

## Supplemental Data

“Phosphotyrosine-based phosphoproteomics for target identification and drug response prediction in AML cell lines”

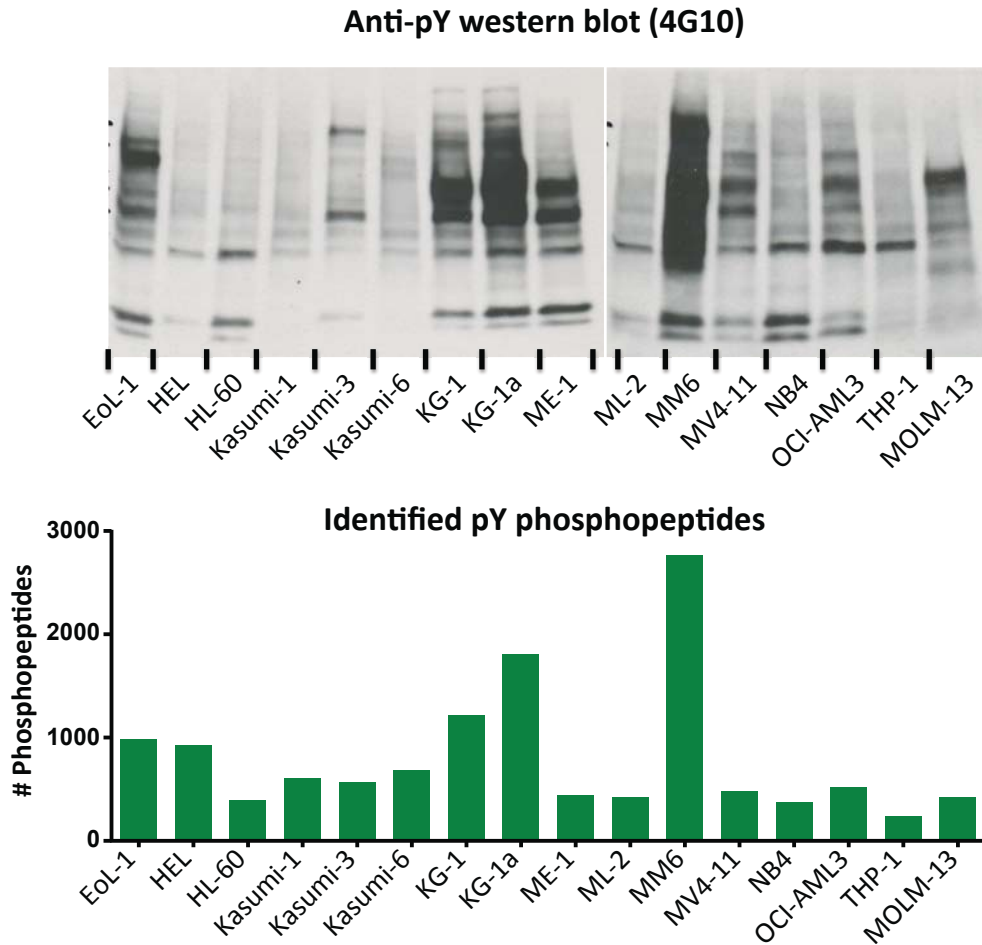
*Carolien van Alphen, Jacqueline Cloos, Robin Beekhof, David G.J. Cucchi, Sander R. Piersma, Jaco C. Knol, Alex Henneman, Thang V. Pham, Johan van Meerloo, Gert J. Ossenkoppele, Henk M.W. Verheul, Jeroen J.W.M. Janssen, and Connie R. Jimenez*

### **Supplemental Figures (pages)**

- Figure S1 (S-2): Overall levels of phosphotyrosine (pY) detected on western blot and numbers of pY phosphopeptides identified by phosphoproteomics show a general correlation.
- Figure S2 (S-3): Clustering and outlier analysis of phosphoproteomic data on protein kinases in an AML cell line panel.
- Figure S3 (S-4-20): Summaries of phosphoproteomic and pharmacological analyses of a panel of AML cell lines.
- Figure S4 (S-21): Drug response curves for quizartinib, ponatinib, and ibrutinib.
- Figure S5 (S-22): Comparison of cell line sensitivity to KIs targeting INKA-identified drivers and those with other targets.
- Figure S6 (S-23-24): Summary of phosphoproteomic analysis of blood sample from AML patients.
- Figure S7 (S-25): Comparison between FLT3-ITD mutant cell line data and FLT3-ITD mutant patient data.
- Figure S8 (S-26-1684): Annotated MS/MS spectra of phosphopeptides identified in AML cell lines.
- Figure S9 (S-1685-1949): Annotated MS/MS spectra of phosphopeptides identified in AML patient samples.

### **Supplemental Tables (separate Excel files)**

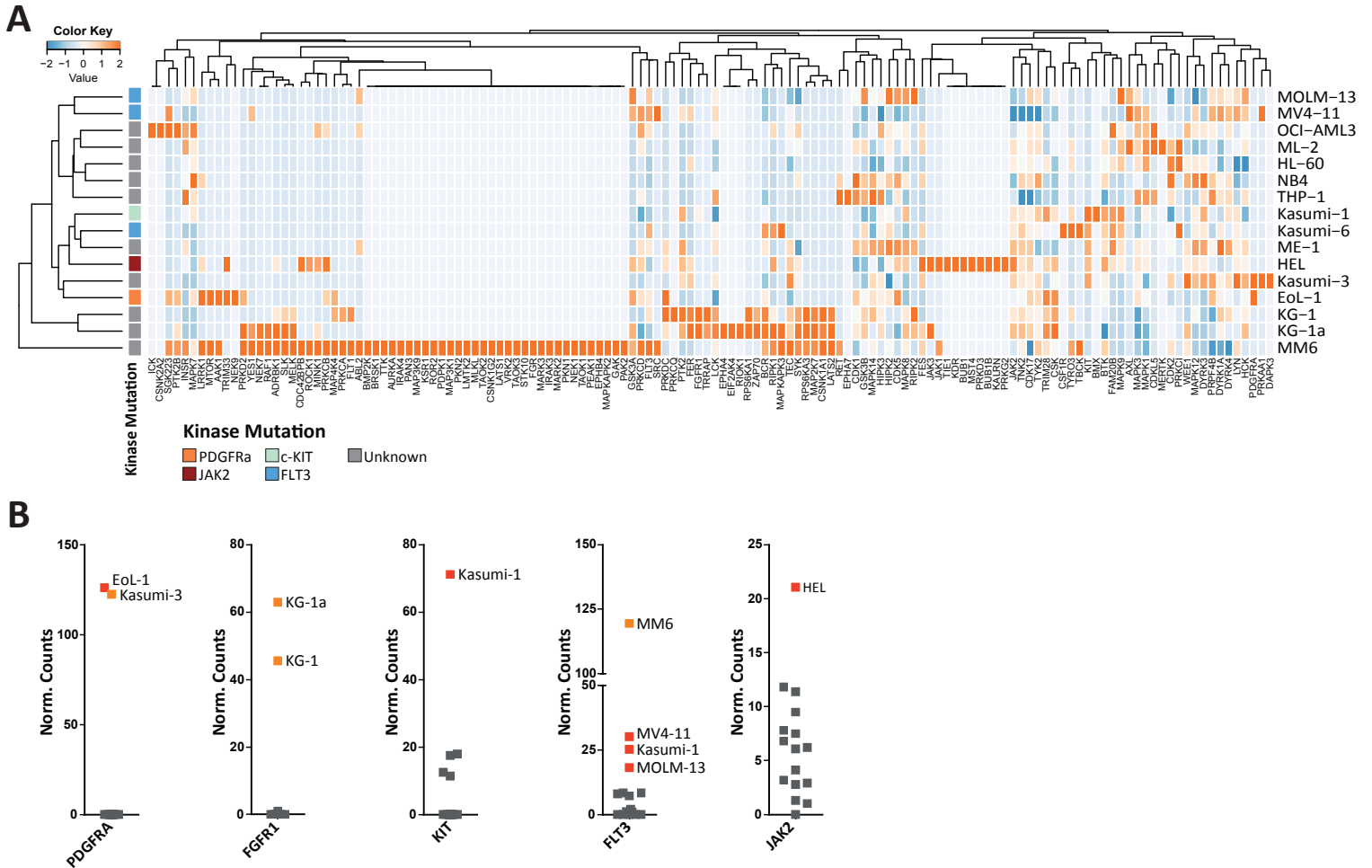
- Supplemental Table S1: (Phospho)peptide and protein identification data.
- Supplemental Table S2: Kinase inhibitor target space analysis.
- Supplemental Table S3: List of AML signaling associated genes.
- Supplemental Table S4: Supporting information for EC50 comparison of KIs targeting identified cell line drivers and those with other targets.
- Supplemental Table S5: Summary of AML cell line analysis results.
- Supplemental Table S6: Overview of AML cell line aberrations.
- Supplemental Table S7: Overview of AML patient sample aberrations.
- Supplemental Table S8: Summary of AML patient analysis results.



**Supplemental Figure S1. Overall levels of phosphotyrosine (pY) detected on western blot and numbers of pY phosphopeptides identified by phosphoproteomics show a general correlation.**

*Upper.* Western blotting for global phosphotyrosine levels (4G10 anti-pY antibody) in the cell line lysates.

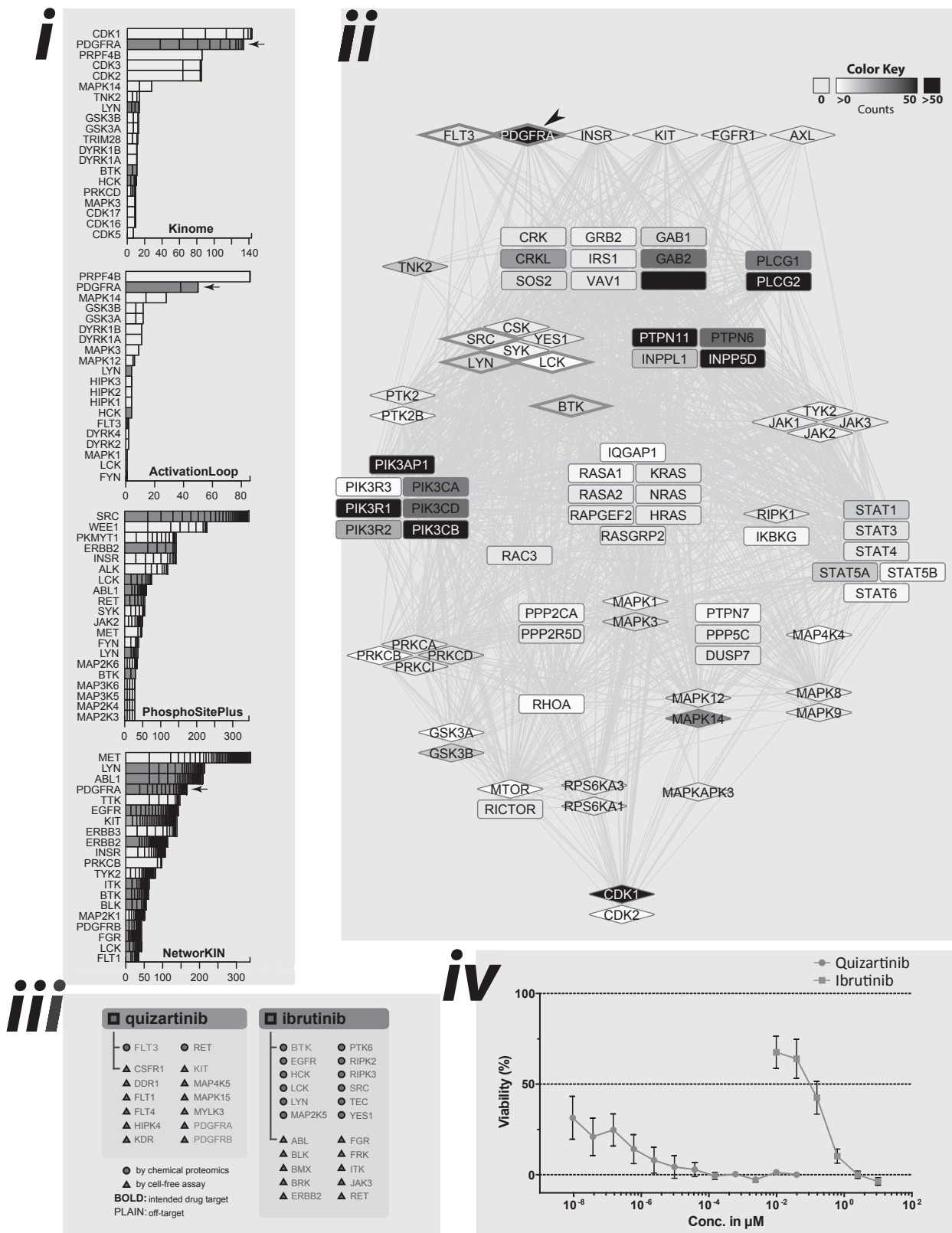
*Lower.* Bar graph depicting the number of pY phosphopeptides immunoprecipitated and identified by mass spectrometry. There is a correlation between global phosphotyrosine levels and the number of identified pY phosphopeptides per cell line. The data confirm signaling heterogeneity in this panel of AML cell lines.



**Supplemental Figure S2. Clustering and outlier analysis of phosphoproteomic data on protein kinases in an AML cell line panel.**

**A.** Hierarchical clustering of the phosphokinome of the 16 AML cell lines. Unsupervised clustering is based on normalized phosphokinase counts. Similar to phosphopeptide based clustering, samples clustered roughly based on high or low levels of phosphorylation. Kinase mutations are indicated as found by standard diagnostic mutational analysis.

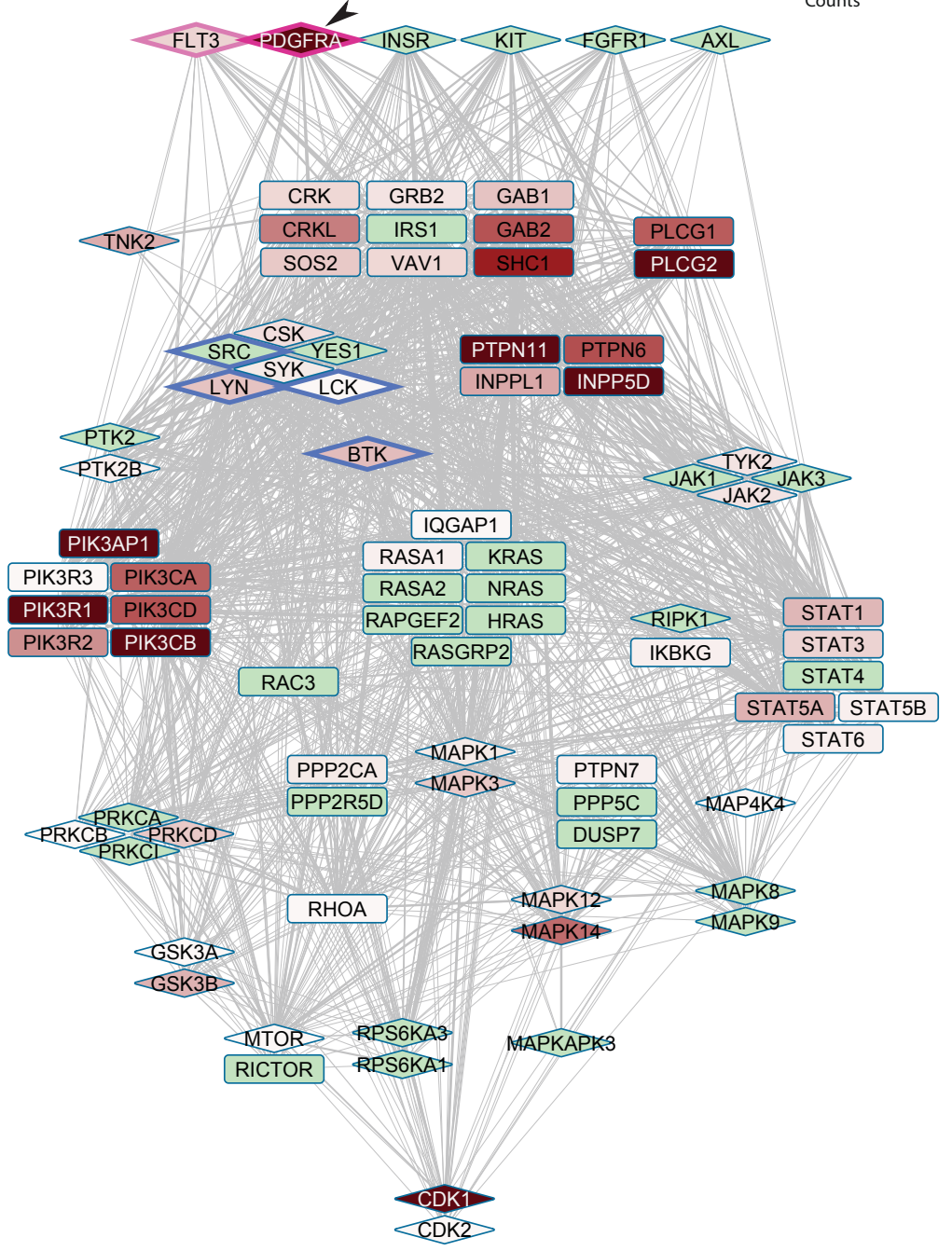
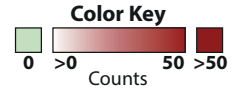
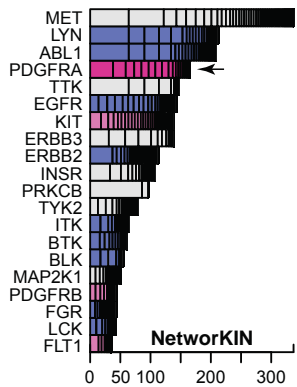
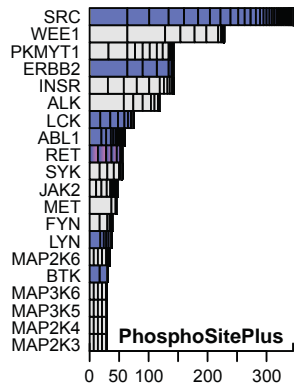
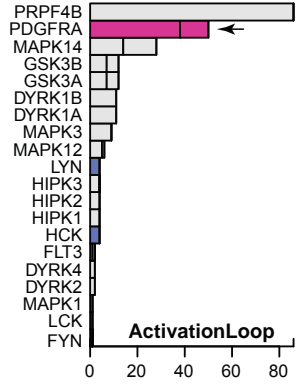
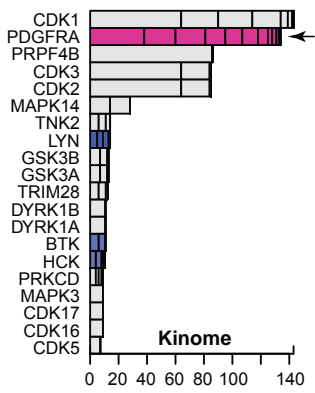
**B.** Outlier analysis of known AML driver kinases shows that kinase phosphorylation levels in TK mutant cell lines (red) are outliers compared to non-mutant cell lines. Additionally, the KG-1, KG-1a, MM6, and Kasumi-3 cell lines (orange) show similar outlier behavior, indicating potential TK driver activity.



### Supplemental Figure S3 General Legend. Summaries of phosphoproteomic and pharmacological analyses of a panel of AML cell lines.

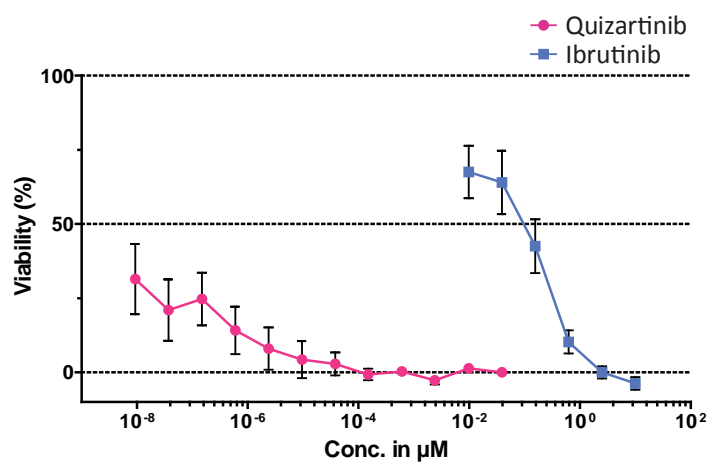
Each sheet contains (i) an overview of bar graphs ranking kinases according to their phosphorylation (*kinome*, phosphorylation anywhere on the protein; *activation loop*, phosphorylation of the activation segment) and according to the phosphorylation of their substrates (*PhosphoSitePlus*, known substrates cataloged in the PhosphoSitePlus database; *NetworkKIN*, predicted substrates assigned by the NetworkKIN algorithm), (ii) a graph of inferred kinase 'activities' mapped onto (primarily) PI3K/RAS-MAPK/JAK-STAT pathways in a signaling network (edges from the STRING database); node fill colors indicate spectral counts as a proxy of activity, (iii) a panel of relevant drugs with intended targets and off-targets (affinity below 500 nM) as determined through chemical proteomics (43) or cell-free assays (42,56,57); and (iv) drug response curves. Kinases from panel iii with a top 20 kinome/activation loop ranking are indicated with corresponding bar fill or node border colors in the different panels. In the following, summary sheets are presented for (A) EoL-1, (B) Kasumi-1, (C) MV4-11, (D) MOLM-13, (E) Kasumi-6, (F) HEL, (G) KG-1, (H) KG-1a, (I) MM6, (J) Kasumi-3, (K) HL-60, (L) ML-2, (M) OCI-AML3, (N) THP-1, (O) NB4, and (P) ME-1.

# EoL-1



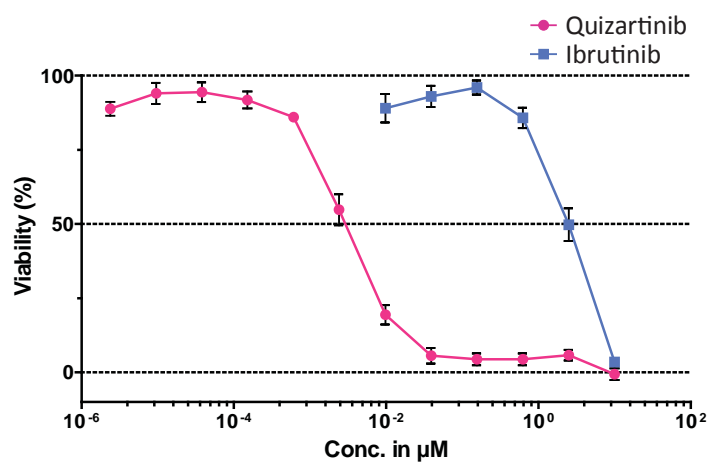
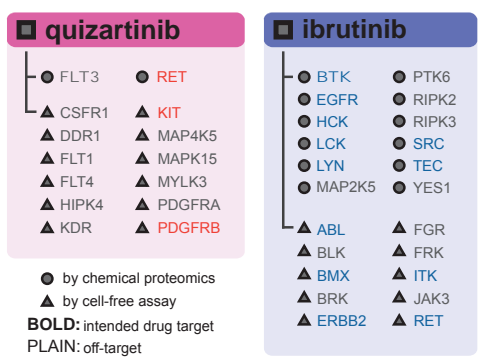
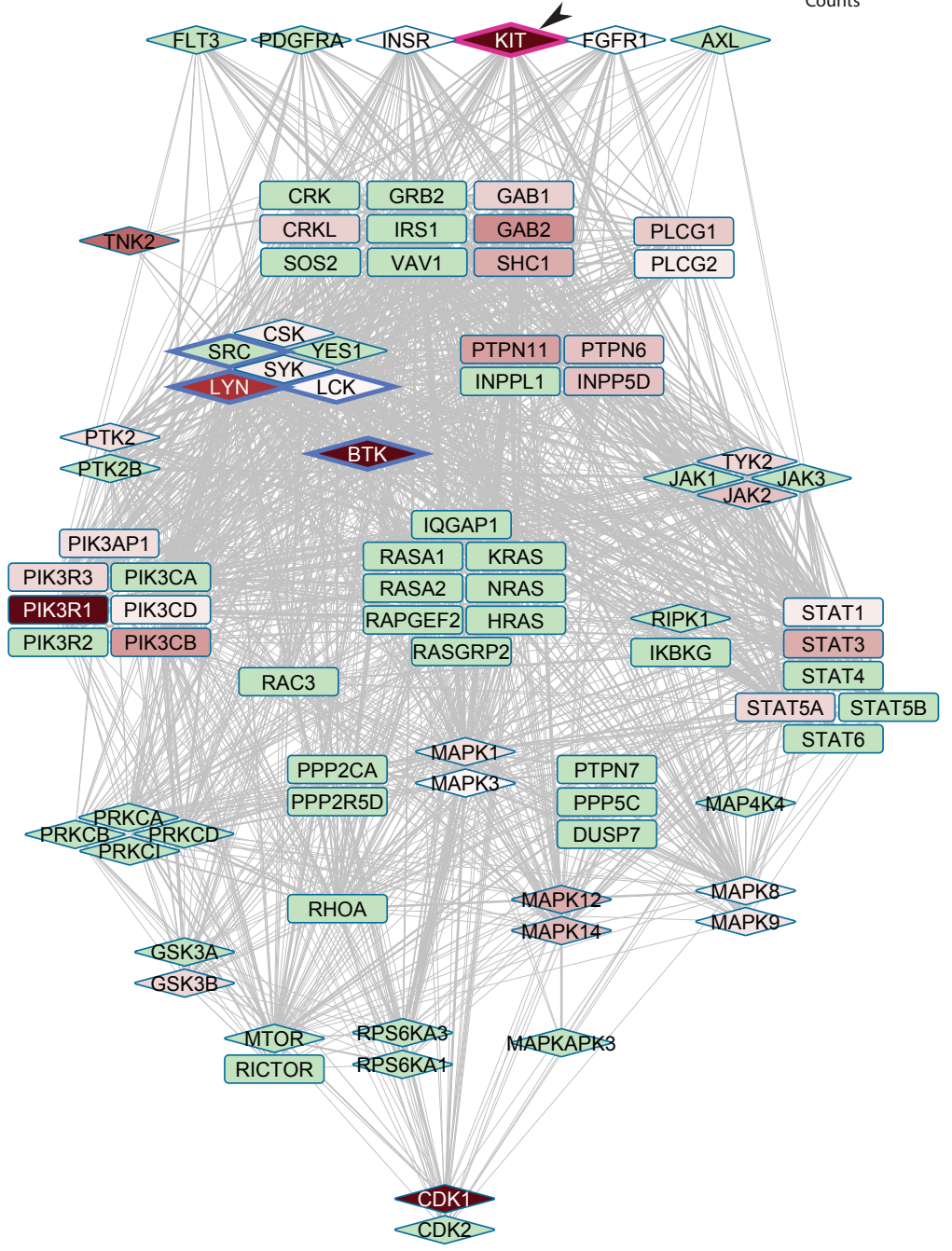
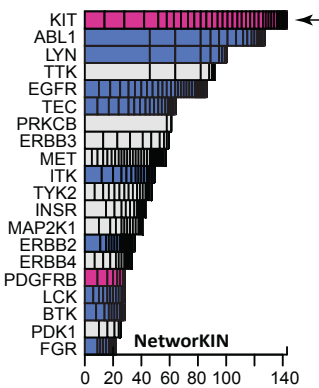
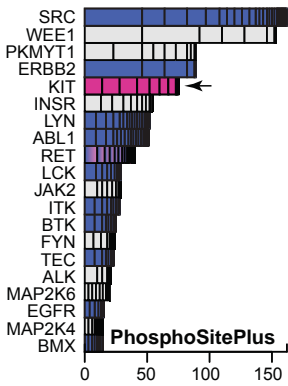
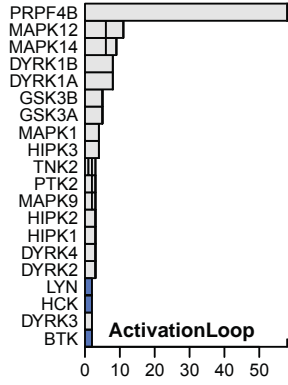
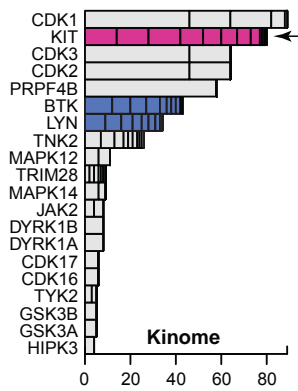
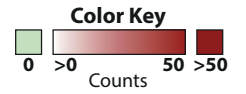
● Quizartinib  
■ Ibrutinib

■ quizartinib		■ ibrutinib	
● FLT3	● RET	● BTK	● PTK6
▲ CSFR1	▲ KIT	● EGFR	● RIPK2
▲ DDR1	▲ MAP4K5	● HCK	● RIPK3
▲ FLT1	▲ MAPK15	● LCK	● SRC
▲ FLT4	▲ MYLK3	● LYN	● TEC
▲ HIPK4	▲ PDGFRA	● MAP2K5	● YES1
▲ KDR	▲ PDGFRB		
● by chemical proteomics		▲ ABL	▲ FGR
▲ by cell-free assay		▲ BLK	▲ FRK
<b>BOLD:</b> intended drug target		▲ BMX	▲ ITK
<b>PLAIN:</b> off-target		▲ BRK	▲ JAK3
		▲ ERBB2	▲ RET



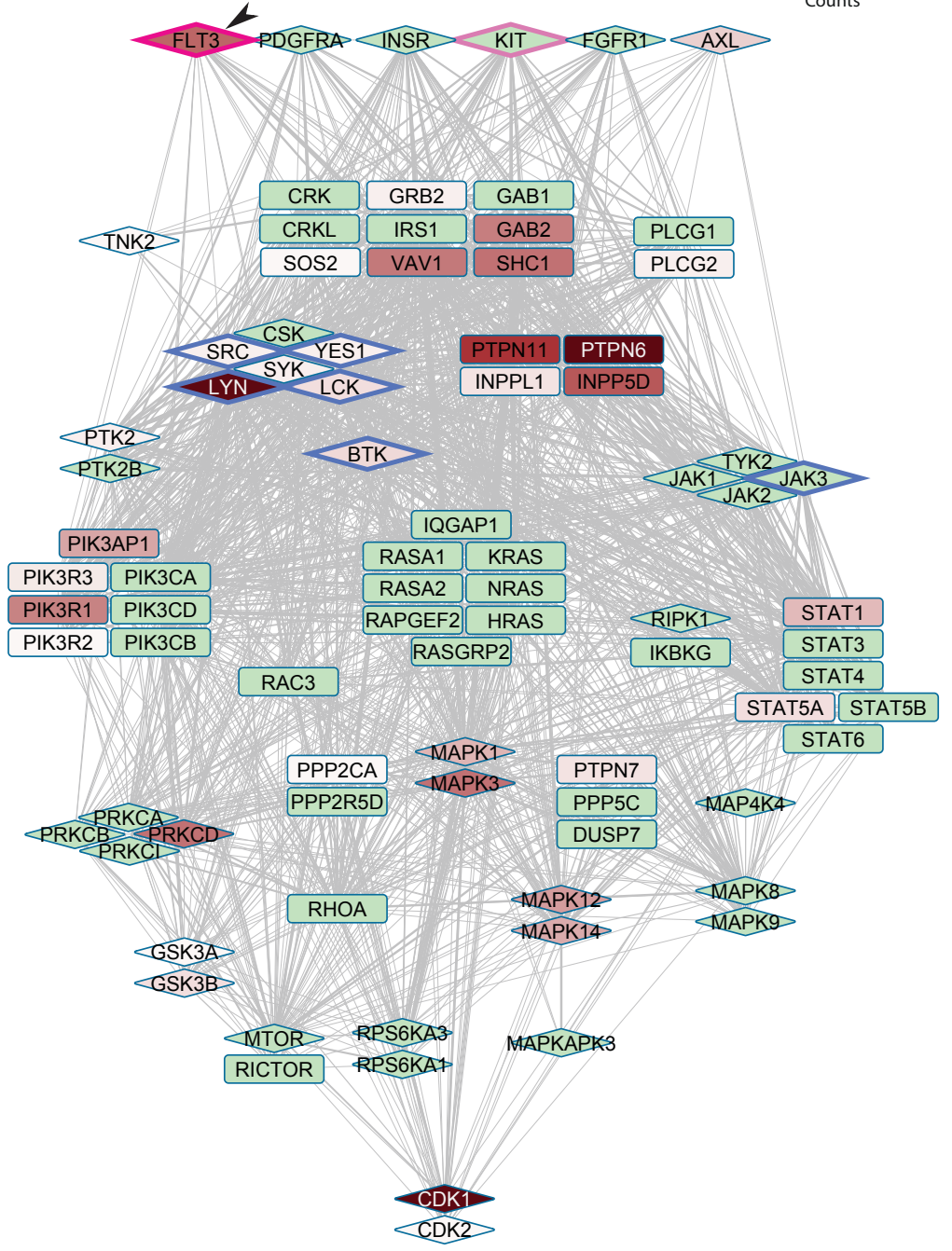
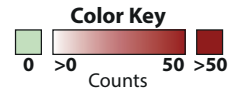
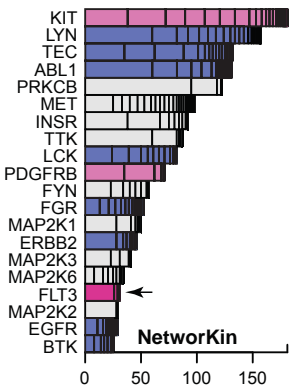
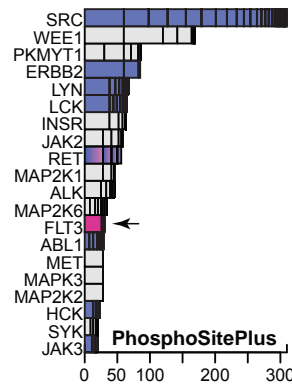
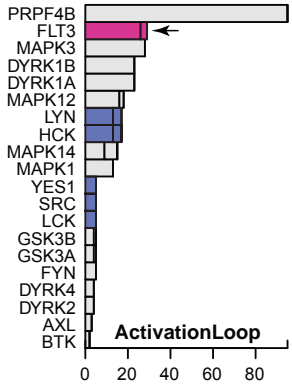
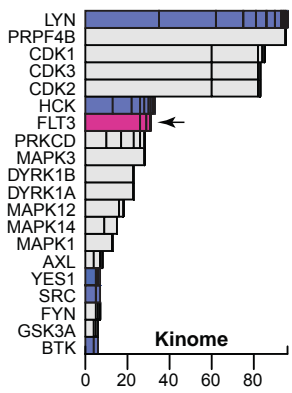
**Supplemental Figure S3A. Analysis of AML cell line EoL-1 carrying a *FIP1L1-PDGFR* fusion.** Arrows (bar graphs) and arrow head (network) indicate the implicated *PDGFRA* kinase.

# Kasumi-1

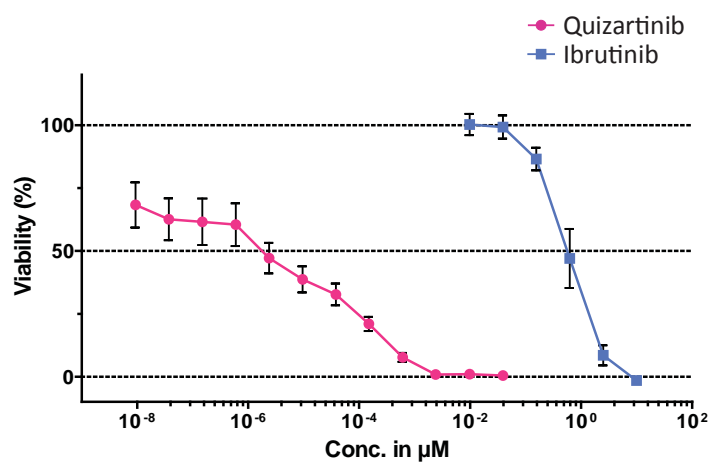


**Supplemental Figure S3B. Analysis of AML cell line Kasumi-1 carrying an N822K KIT point mutation.** Arrows (bar graphs) and arrow head (network) indicate the implicated KIT kinase.

# MV4-11

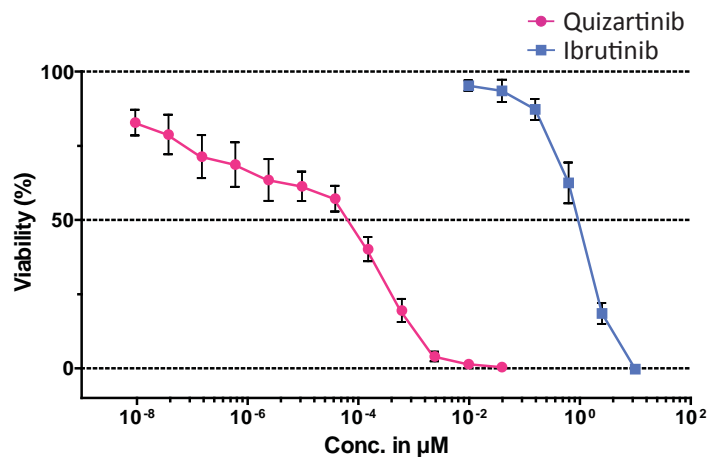
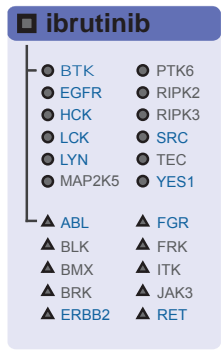
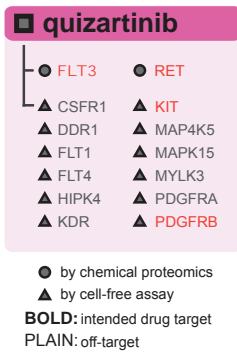
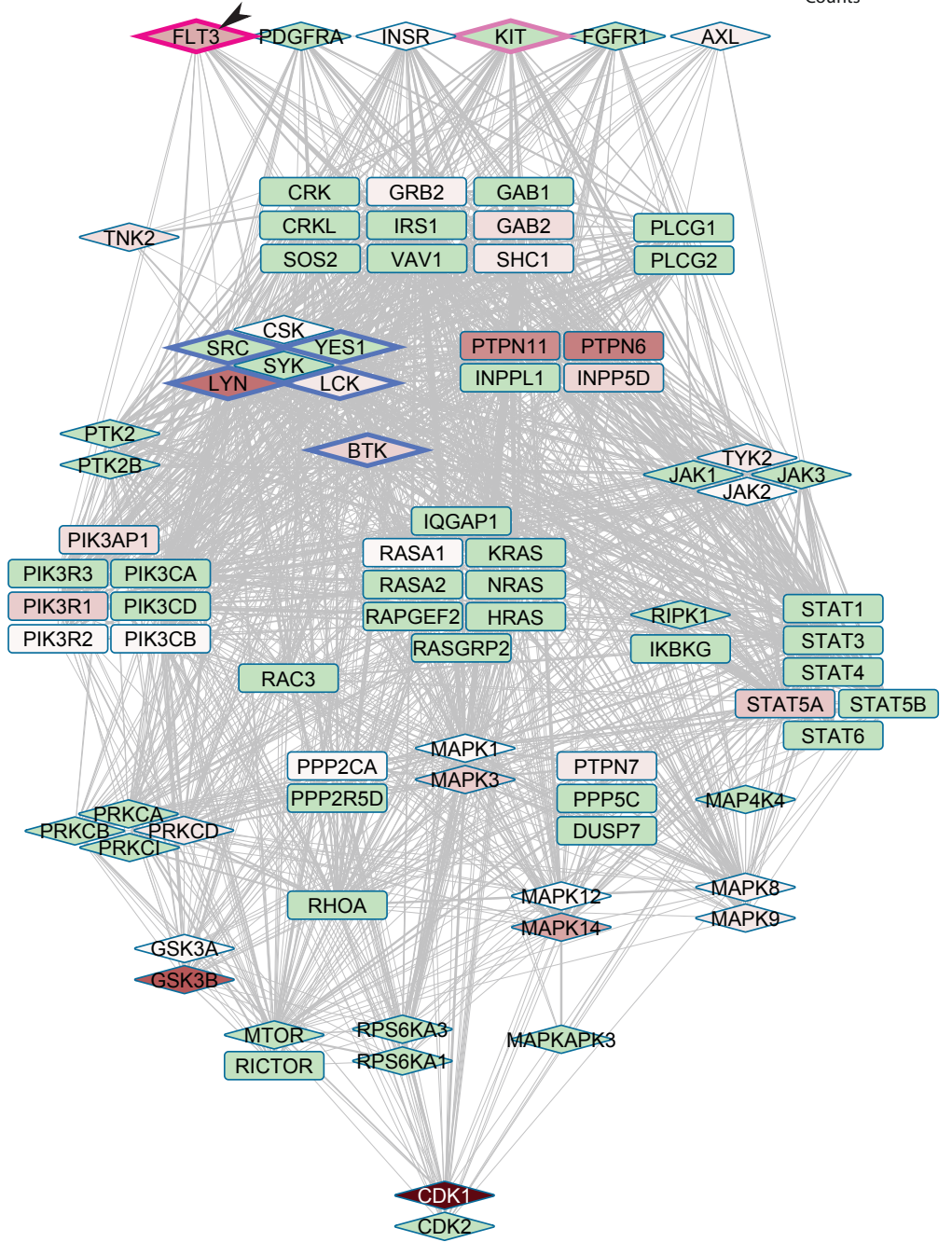
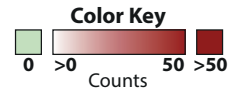
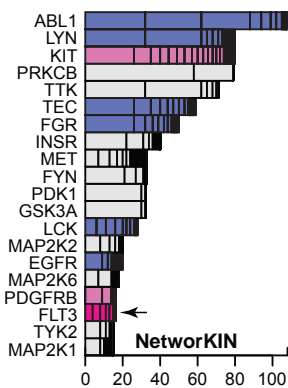
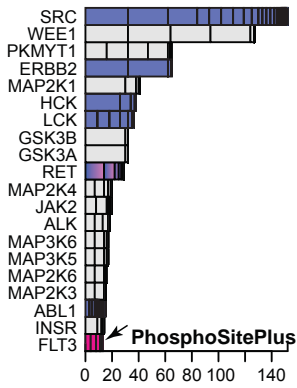
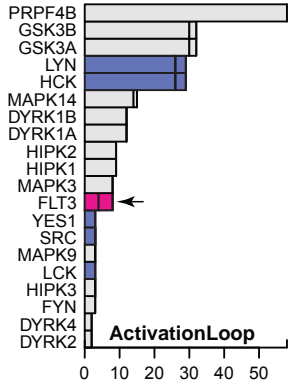
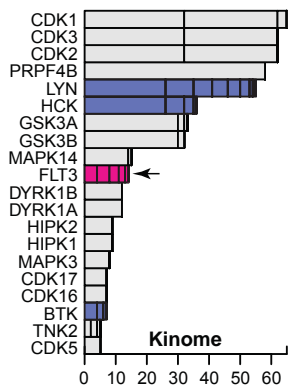


quizartinib		ibrutinib	
● FLT3	● RET	● BTK	● PTK6
▲ CSFR1	▲ KIT	● EGFR	● RIPK2
▲ DDR1	▲ MAP4K5	● HCK	● RIPK3
▲ FLT1	▲ MAPK15	● LCK	● SRC
▲ FLT4	▲ MYLK3	● LYN	● TEC
▲ HIPK4	▲ PDGFRA	● MAP2K5	● YES1
▲ KDR	▲ PDGFRB		
● by chemical proteomics		▲ ABL	▲ FGR
▲ by cell-free assay		▲ BLK	▲ FRK
<b>BOLD:</b> intended drug target		▲ BMX	▲ ITK
<b>PLAIN:</b> off-target		▲ BRK	▲ JAK3
		▲ ERBB2	▲ RET



**Supplemental Figure S3C. Analysis of AML cell line MV4-11 carrying a FLT3-ITD point mutation.**  
 Arrows (bar graphs) and arrow head (network) indicate the implicated FLT3 kinase.

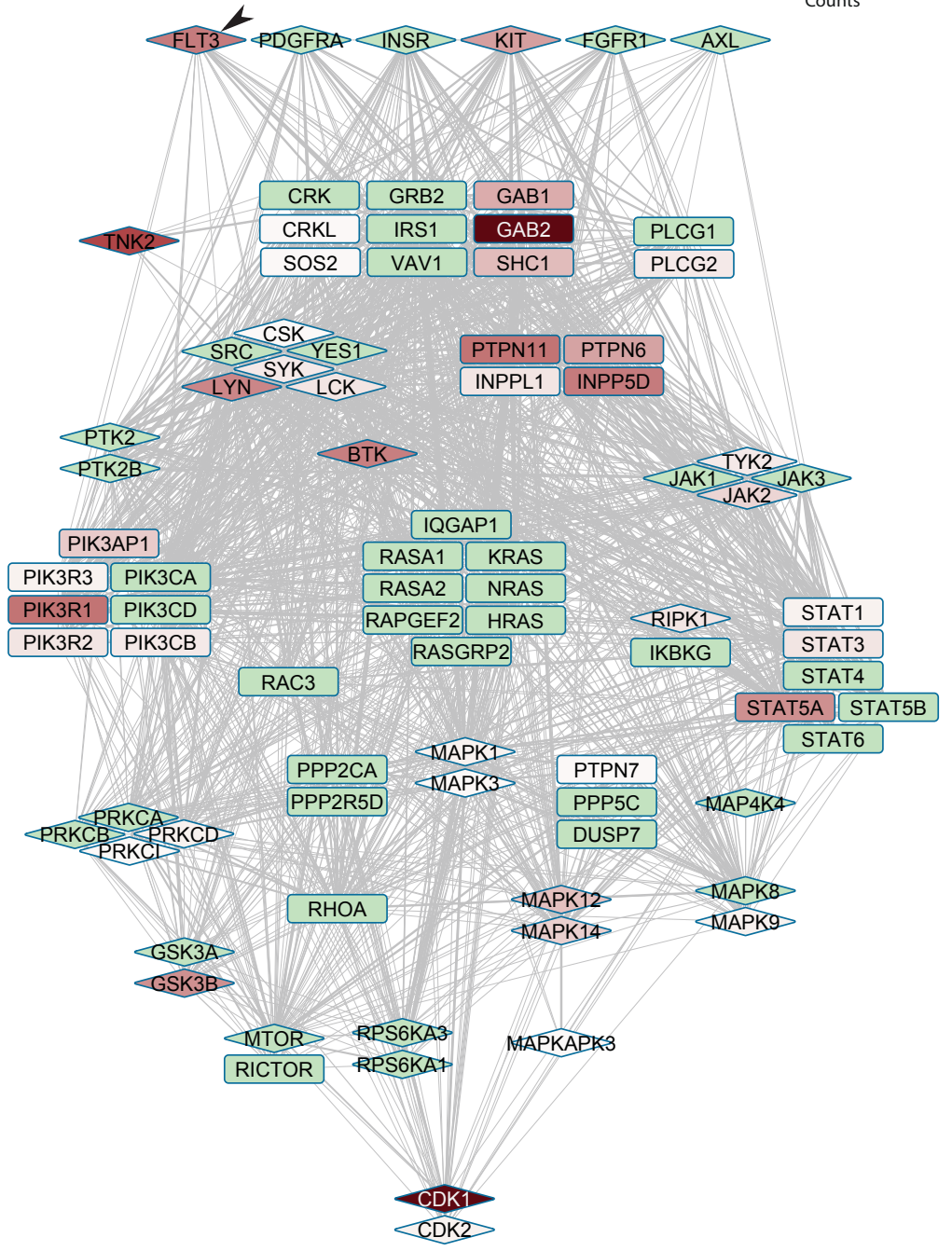
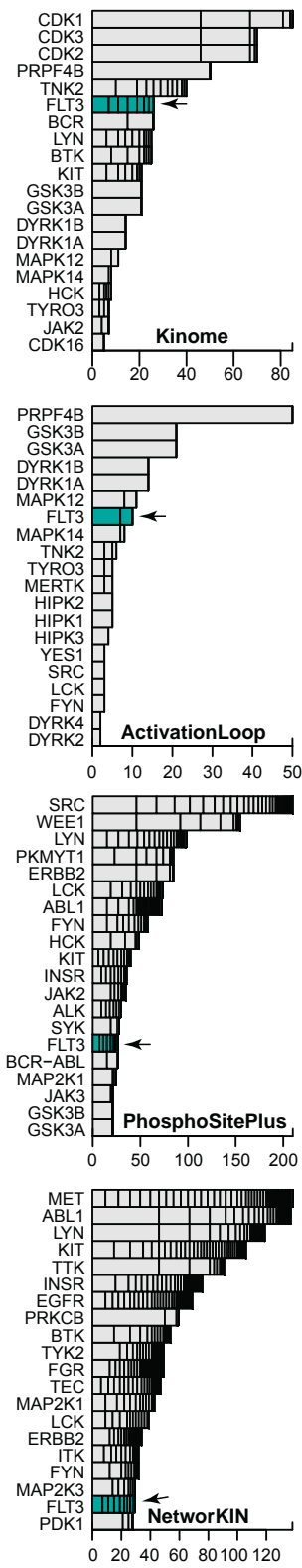
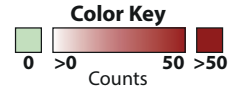
# MOLM-13



**Supplemental Figure S3D. Analysis of AML cell line MOLM-13 carrying a FLT3-ITD point mutation.** Arrows (bar graphs) and arrow head (network) indicate the implicated FLT3 kinase.

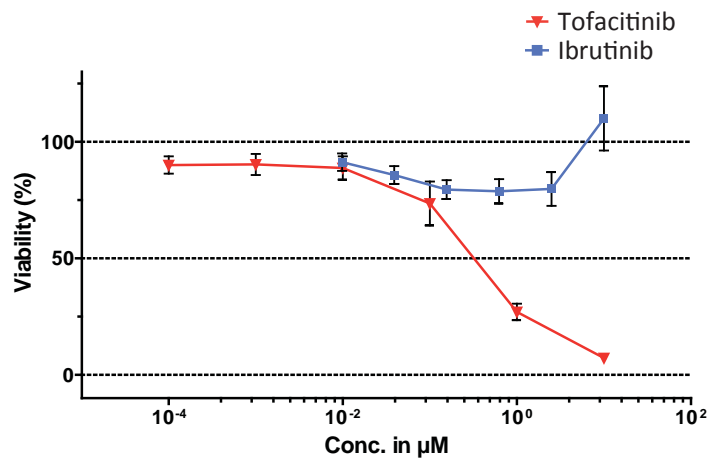
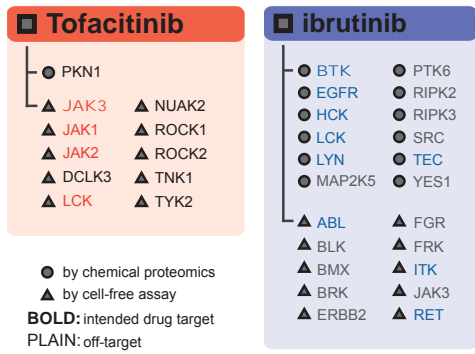
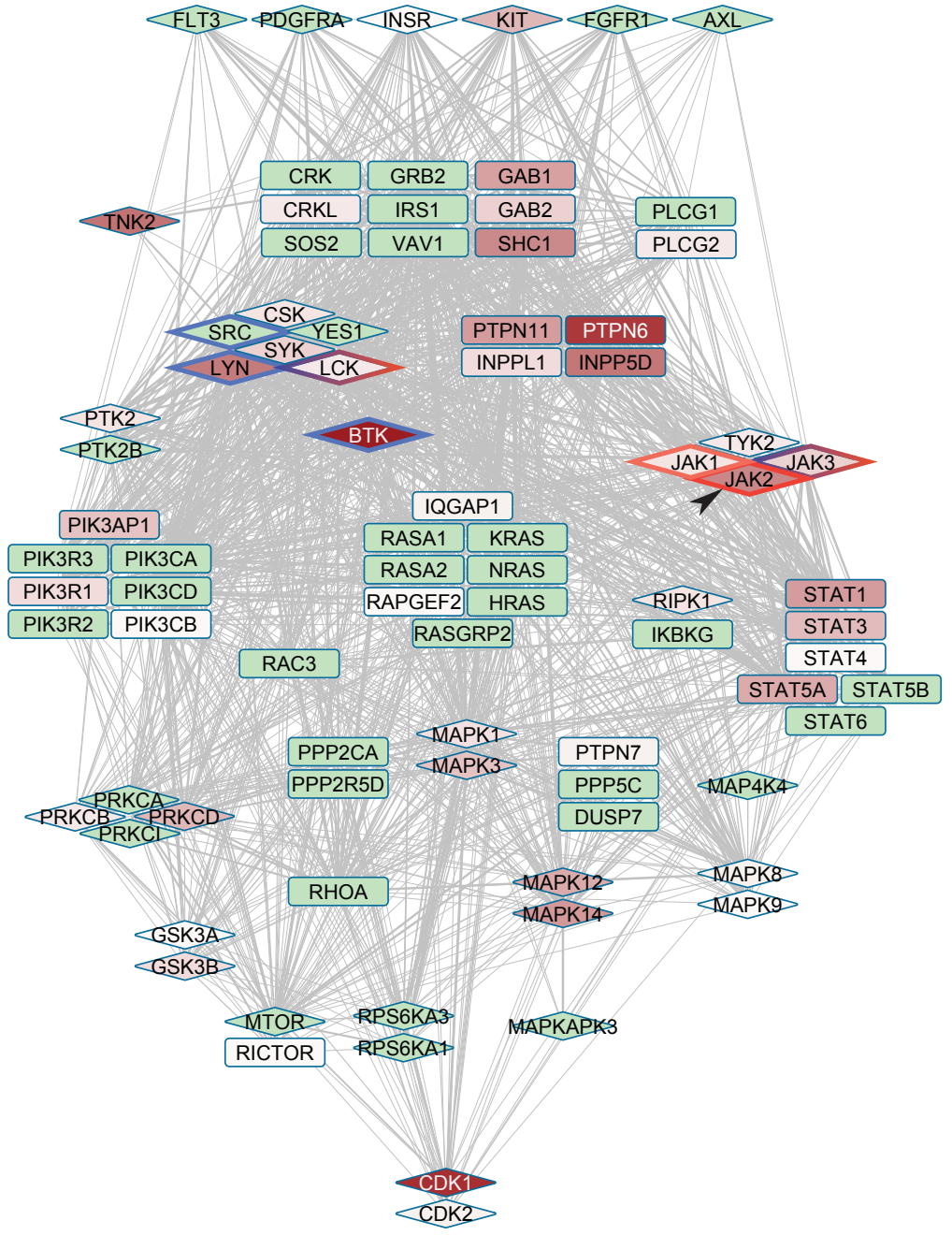
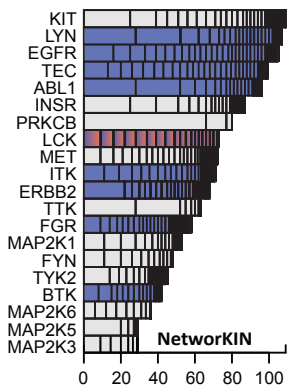
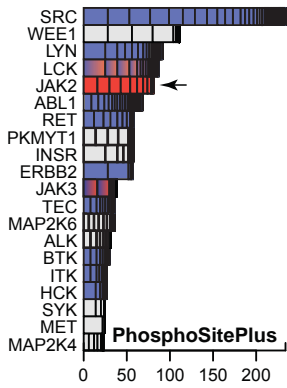
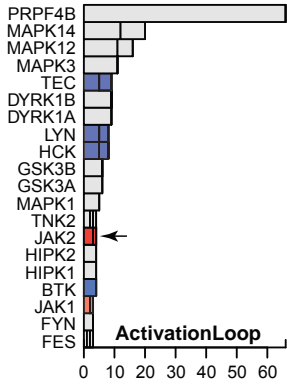
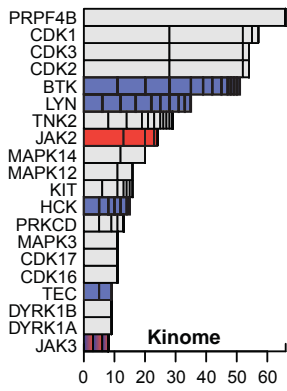
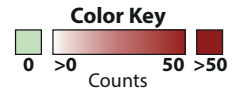


# Kasumi-6



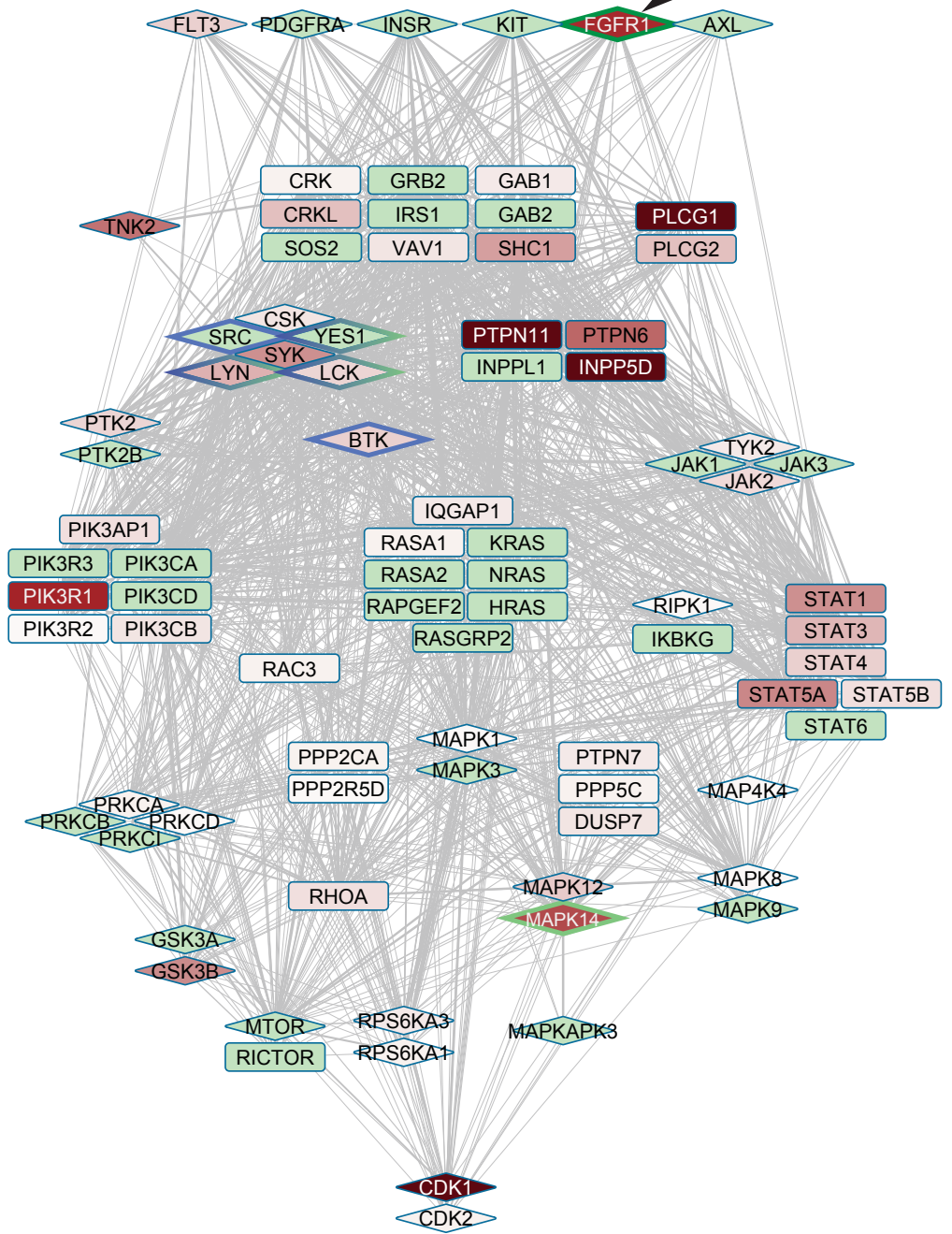
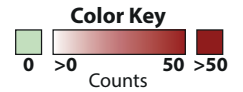
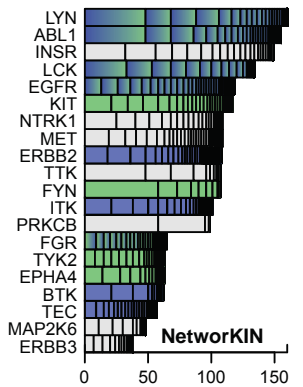
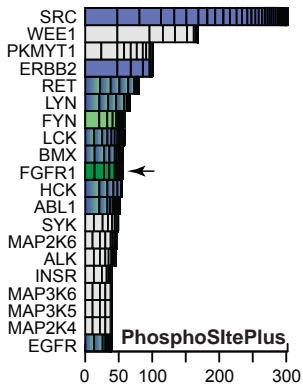
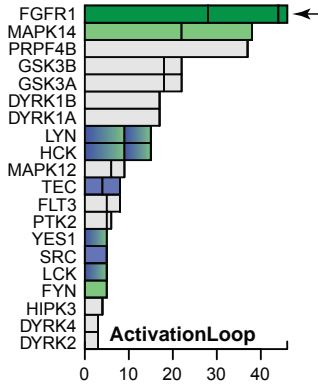
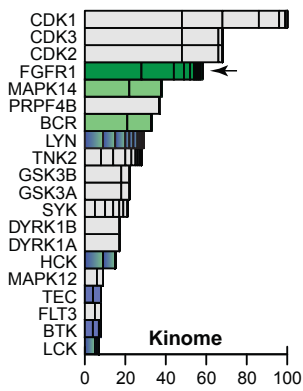
**Supplemental Figure S3E. Analysis of AML cell line Kasumi-6 carrying a *FLT3*-ITD point mutation.** Arrows (bar graphs) and arrow head (network) indicate the implicated *FLT3* kinase.

# HEL



**Supplemental Figure S3F. Analysis of AML cell line HEL carrying a V617F JAK2 mutation.**  
 Arrows (bar graphs) and arrow heads (network) indicate the implicated JAK2 kinase.

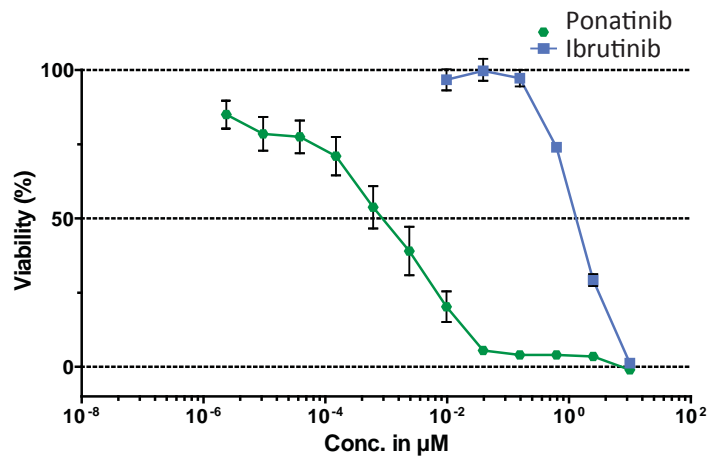
# KG-1



- ponatinib**
- ABL1
  - BCR
  - ABL2
  - CDK17
  - DDR1
  - DDR2
  - EPHA2
  - EPHA5
  - FRK
  - FYN
  - GRB2
  - HCK
  - LCK
  - LYN
  - MAP4K2
  - MAP14
  - MLTK
  - MYLK
  - RET
  - RIPK2
  - TNK1
  - ▲ ARG
  - ▲ ERBB4
  - ▲ LYNB
  - ▲ VEGFR1-3
  - ▲ BLK
  - ▲ FGFR1
  - ▲ MER
  - ▲ YES1
  - ▲ BMX
  - ▲ FGFR2-4
  - ▲ FMS
  - ▲ TIE2
  - ▲ BRK
  - ▲ FGR
  - ▲ PDGFRA
  - ▲ TRKA-C
  - ▲ EGFR
  - ▲ FMS
  - ▲ PDGFRB
  - ▲ EPHA1-8
  - ▲ JAK1-4
  - ▲ PYK2
  - ▲ EPHB1-4
  - ▲ KIT
  - ▲ TYK2

- ibrutinib**
- BTK
  - EGFR
  - HCK
  - LCK
  - LYN
  - MAP2K5
  - PTK6
  - RIPK2
  - RIPK3
  - SRC
  - TEC
  - YES1
  - ▲ ABL
  - ▲ FGR
  - ▲ BLK
  - ▲ FRK
  - ▲ BMX
  - ▲ ITK
  - ▲ BRK
  - ▲ JAK3
  - ▲ ERBB2
  - ▲ RET

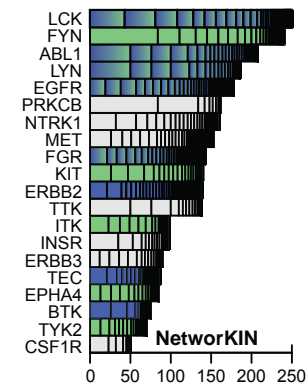
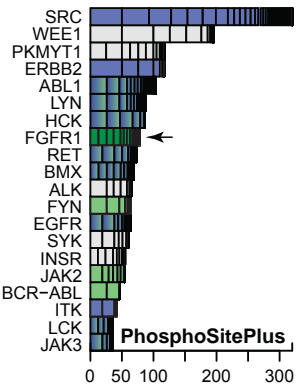
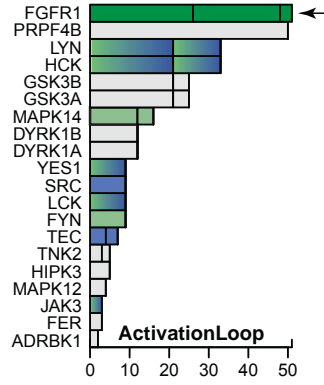
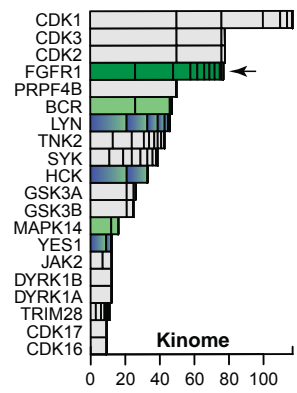
● by chemical proteomics  
▲ by cell-free assay  
BOLD: intended drug target  
PLAIN: off-target



**Supplemental Figure S3G. Analysis of AML cell line KG-1 lacking a known driver mutation.**

Arrows (bar graphs) and arrow head (network) indicate the FGFR1 kinase as a potential driver in this cell line.

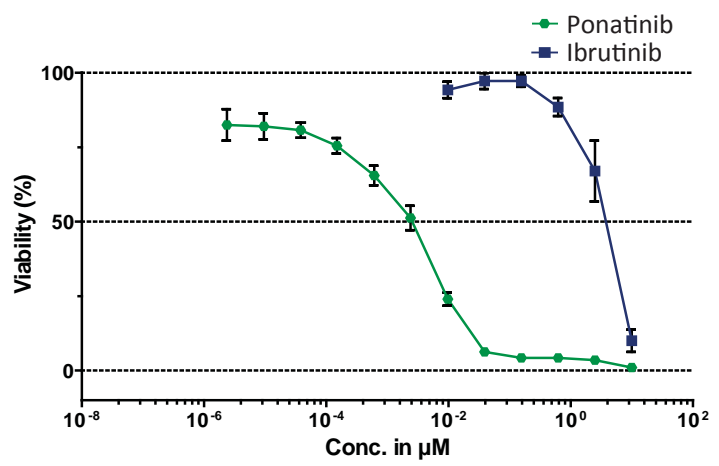
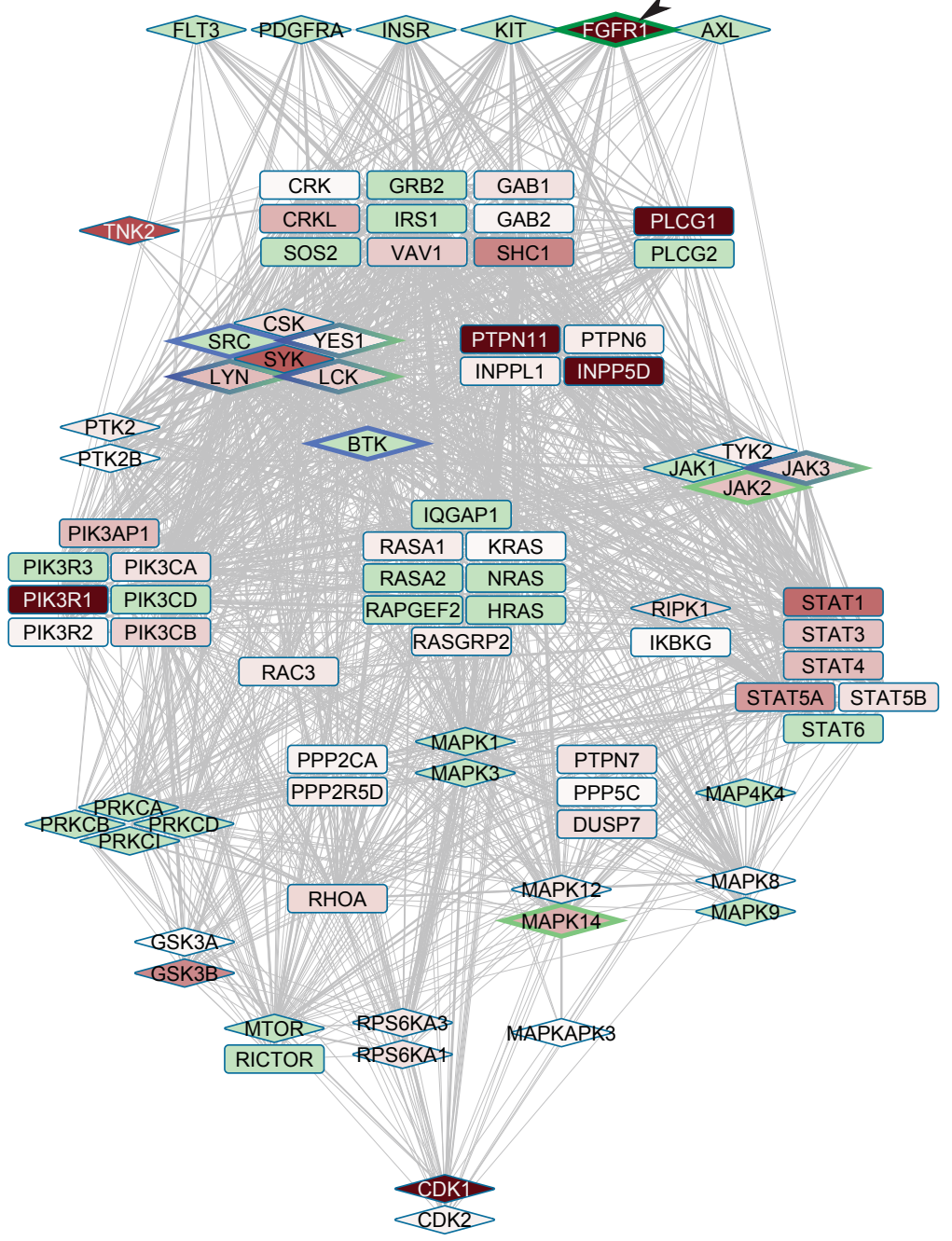
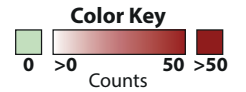
# KG-1a



ponatinib			
● ABL1	● EPHA2	● LCK	● MYLK
● BCR	● EPHA5	● LYN	● RET
● ABL2	● FRK	● MAP4K2	● RIPK2
● CDK17	● FYN	● MAP4K4	● TNK1
● DDR1	● GRB2	● MAPK14	● TNK1
● DDR2	● HCK	● MLTK	
▲ ARG	▲ ERBB4	▲ LYNB	▲ VEGFR1-3
▲ BLK	▲ FGFR1	▲ MER	▲ YES1
▲ BMX	▲ FGFR2-4	▲ FMS	▲ TIE2
▲ BRK	▲ FGR	▲ PDGFRA	▲ TRKA-C
▲ EGFR	▲ FMS	▲ PDGFRB	
▲ EPHA1-8	▲ JAK1-4	▲ PYK2	
▲ EPHB1-4	▲ KIT	▲ TYK2	

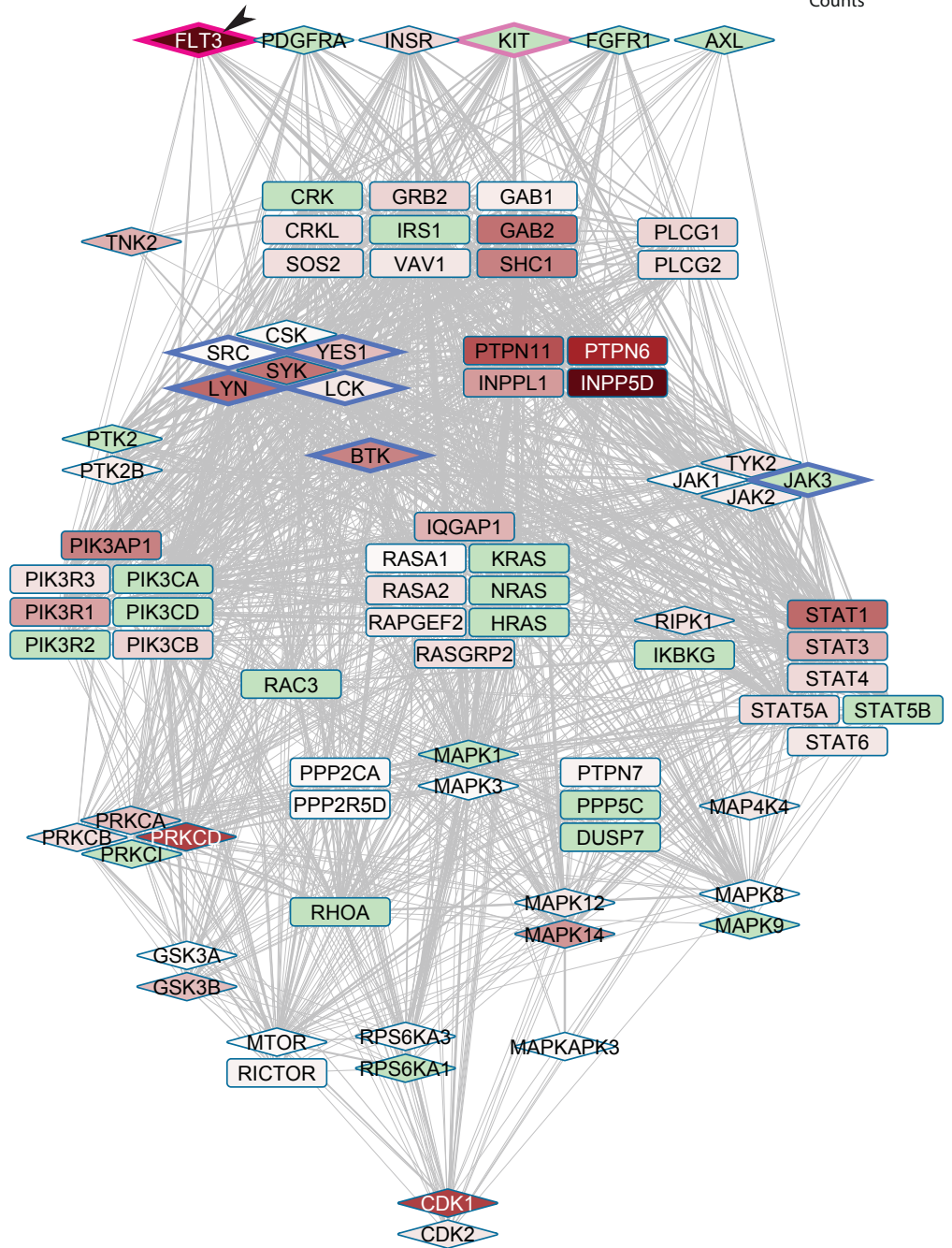
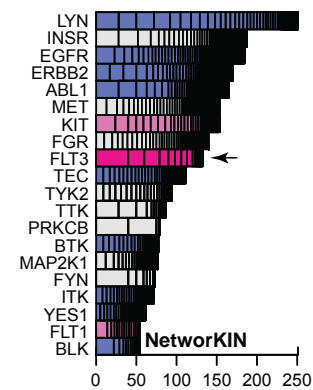
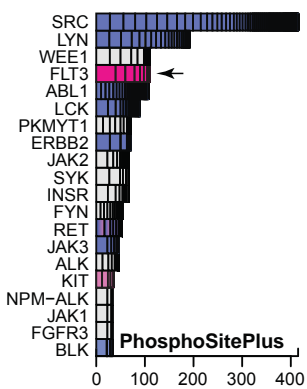
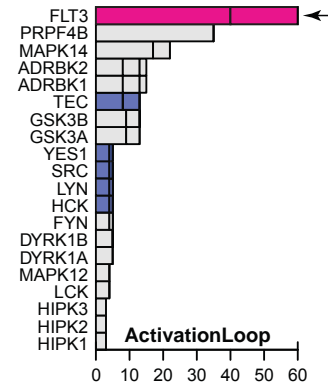
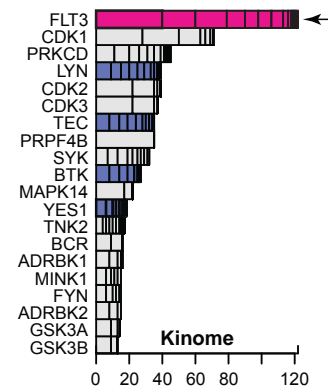
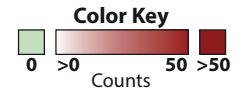
ibrutinib	
● BTK	● PTK6
● EGFR	● RIPK2
● HCK	● RIPK3
● LCK	● SRC
● LYN	● TEC
● MAP2K5	● YES1
▲ ABL	▲ FGR
▲ BLK	▲ FRK
▲ BMX	▲ ITK
▲ BRK	▲ JAK3
▲ ERBB2	▲ RET

● by chemical proteomics  
▲ by cell-free assay  
BOLD: intended drug target  
PLAIN: off-target



**Supplemental Figure S3H. Analysis of AML cell line KG-1a lacking a known driver mutation.**  
Arrows (bar graphs) and arrow head (network) indicate the FGFR1 kinase as a potential driver in this cell line.

## MM6



**quizartinib**



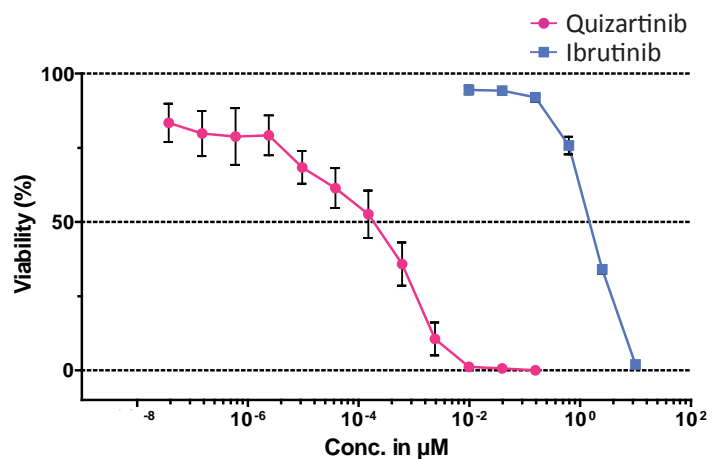
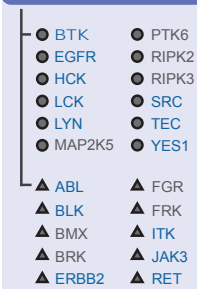
● by chemical proteomics

▲ by cell-free assay

**BOLD:** intended drug target

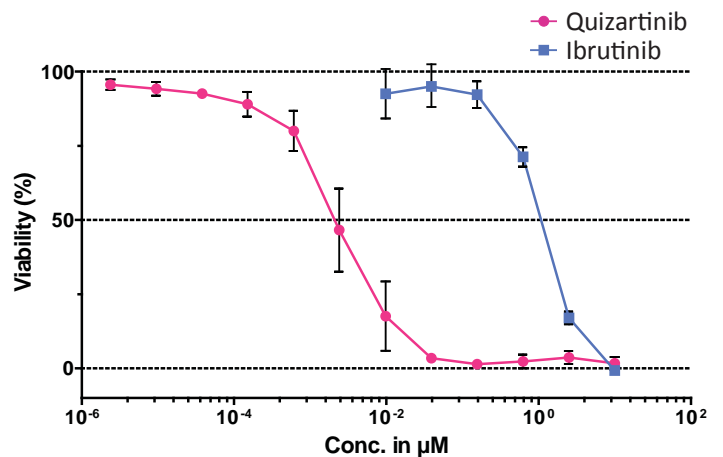
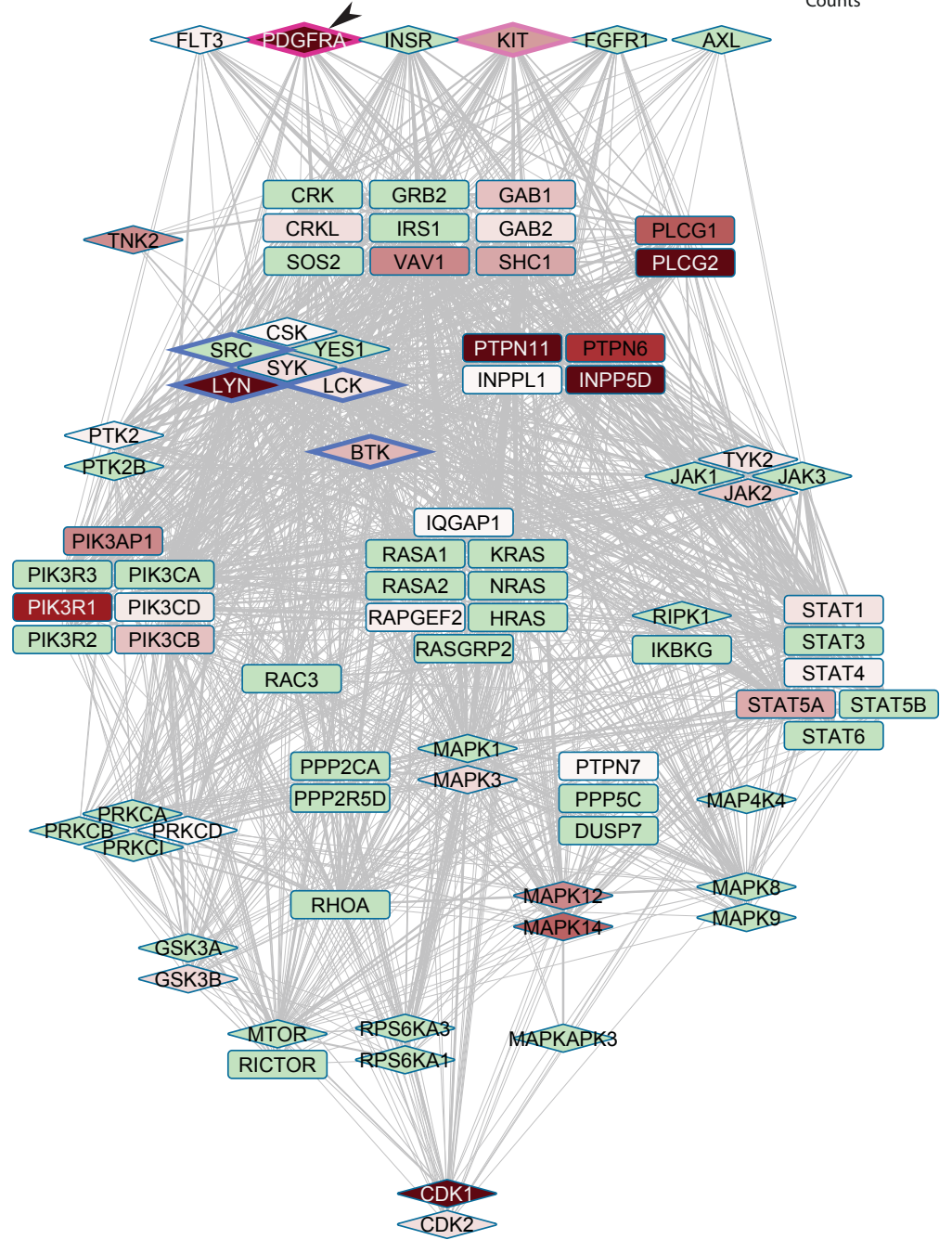
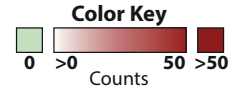
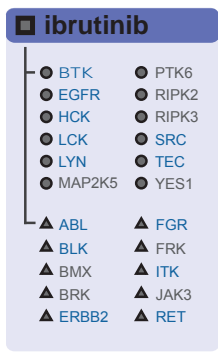
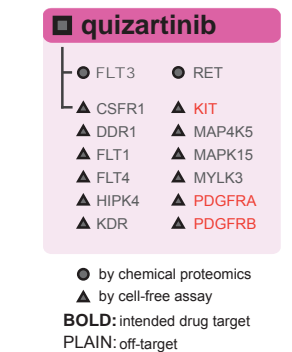
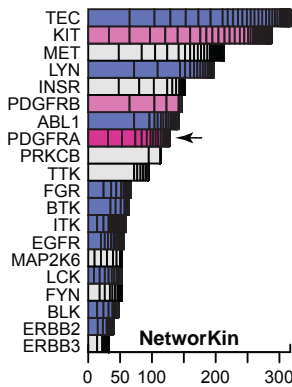
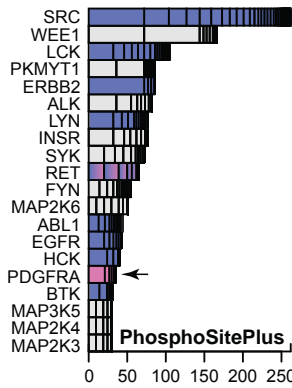
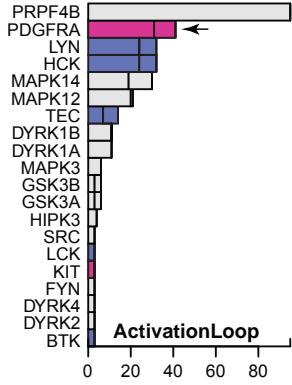
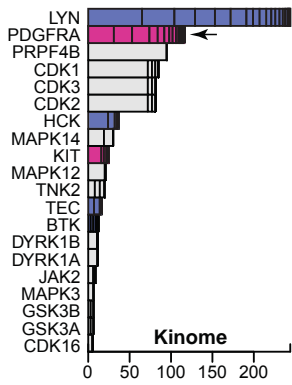
**PLAIN:** off-target

**ibrutinib**



**Supplemental Figure S3I. Analysis of AML cell line MM6 (Mono-Mac-6) lacking a known driver mutation.** Arrows (bar graphs) and arrow head (network) indicate the FLT3 kinase as a potential driver in this cell line.

# Kasumi-3

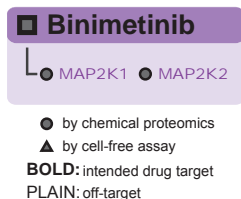
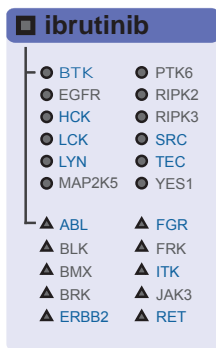
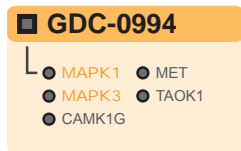
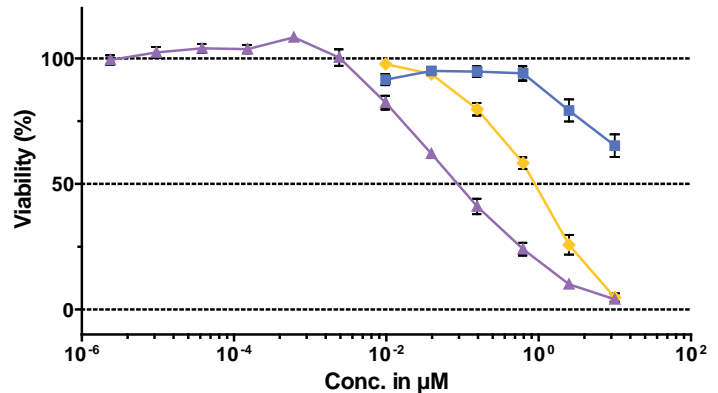
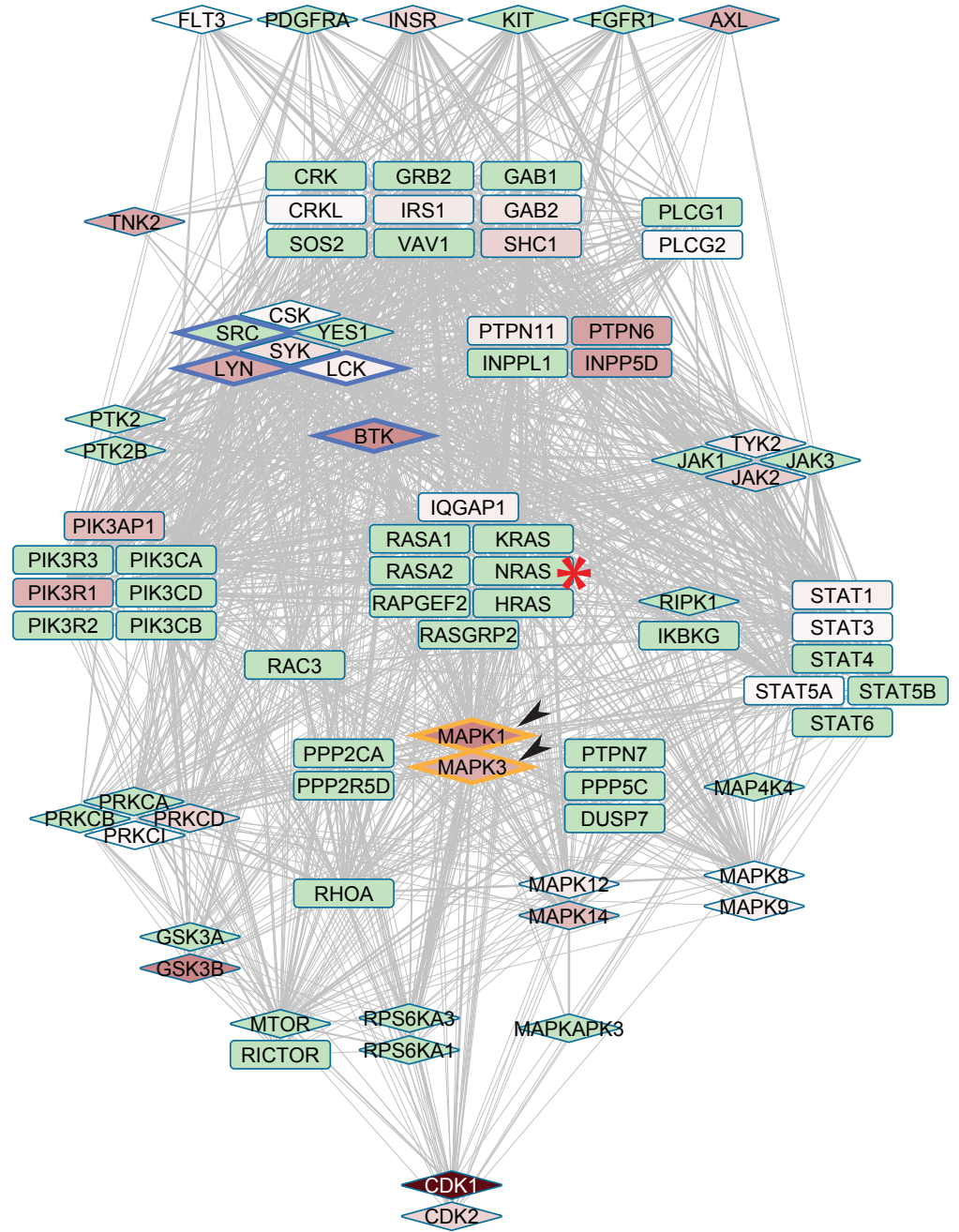
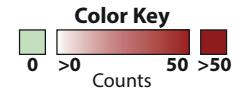
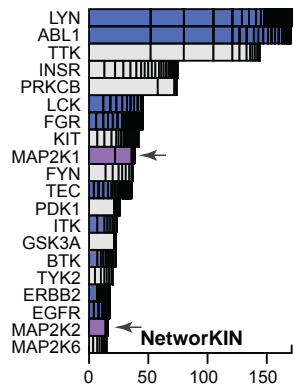
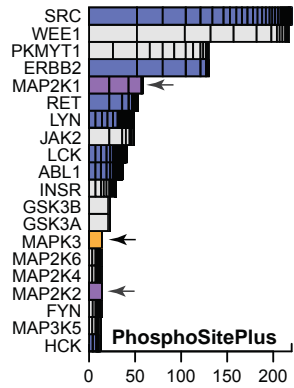
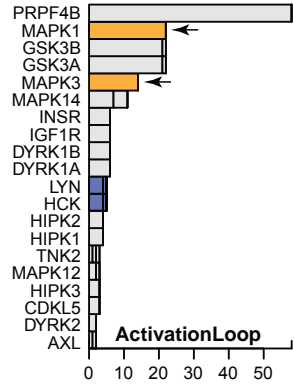
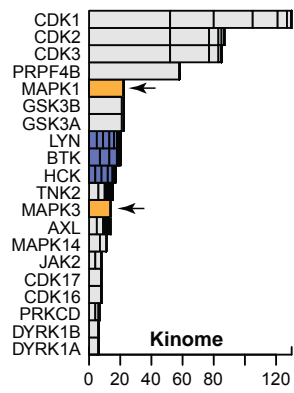


**Supplemental Figure S3J. Analysis of AML cell line Kasumi-3 lacking a known driver mutation.**

Arrows (bar graphs) and arrow head (network) indicate the PDGFRA kinase as a potential driver in this cell line.



# ML-2

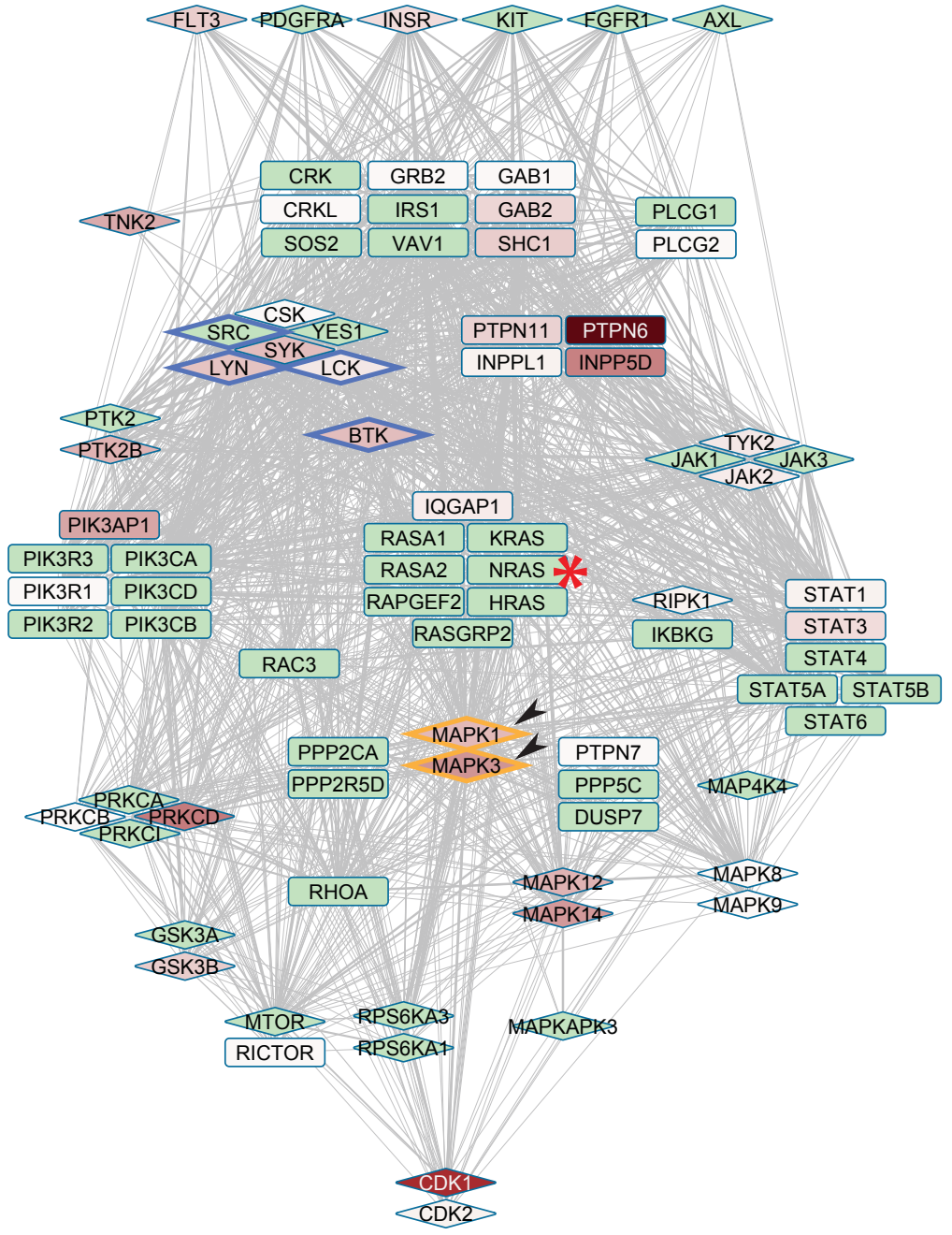
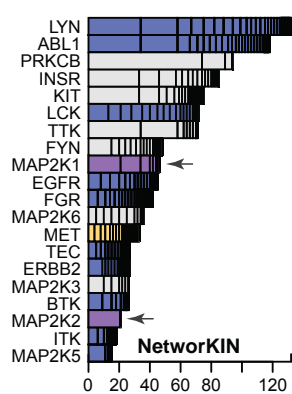
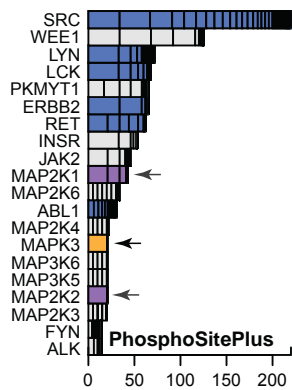
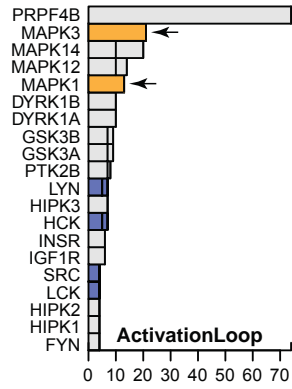
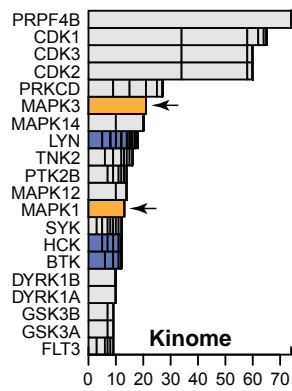
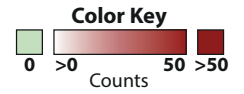


**Supplemental Figure S3L. Analysis of AML cell line ML-2 lacking a known driver mutation.**

Arrows/arrow heads indicate high-ranking kinases of the MAPK1/3 pathway; red asterisk: NRAS mutated.



# OCI-AML3



- ◆ GDC-0994
- ▲ Binimetinib
- Ibrutinib

**GDC-0994**

- MAPK1
- MET
- MAPK3
- TAOK1
- CAMK1G

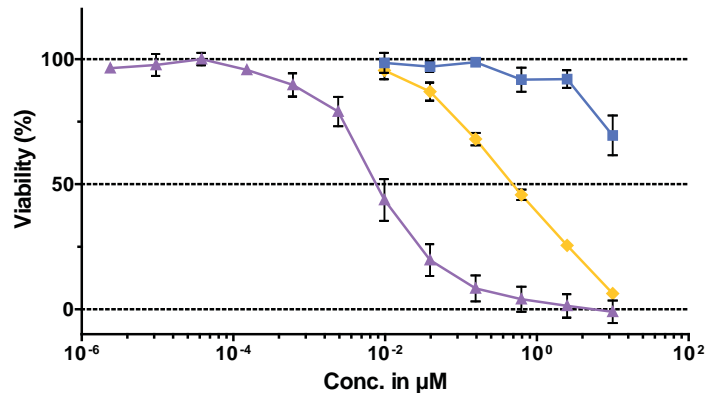
**Binimetinib**

- MAP2K1
- MAP2K2

● by chemical proteomics  
▲ by cell-free assay  
BOLD: intended drug target  
PLAIN: off-target

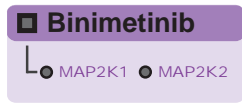
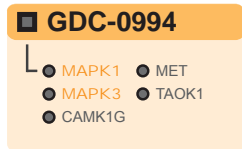
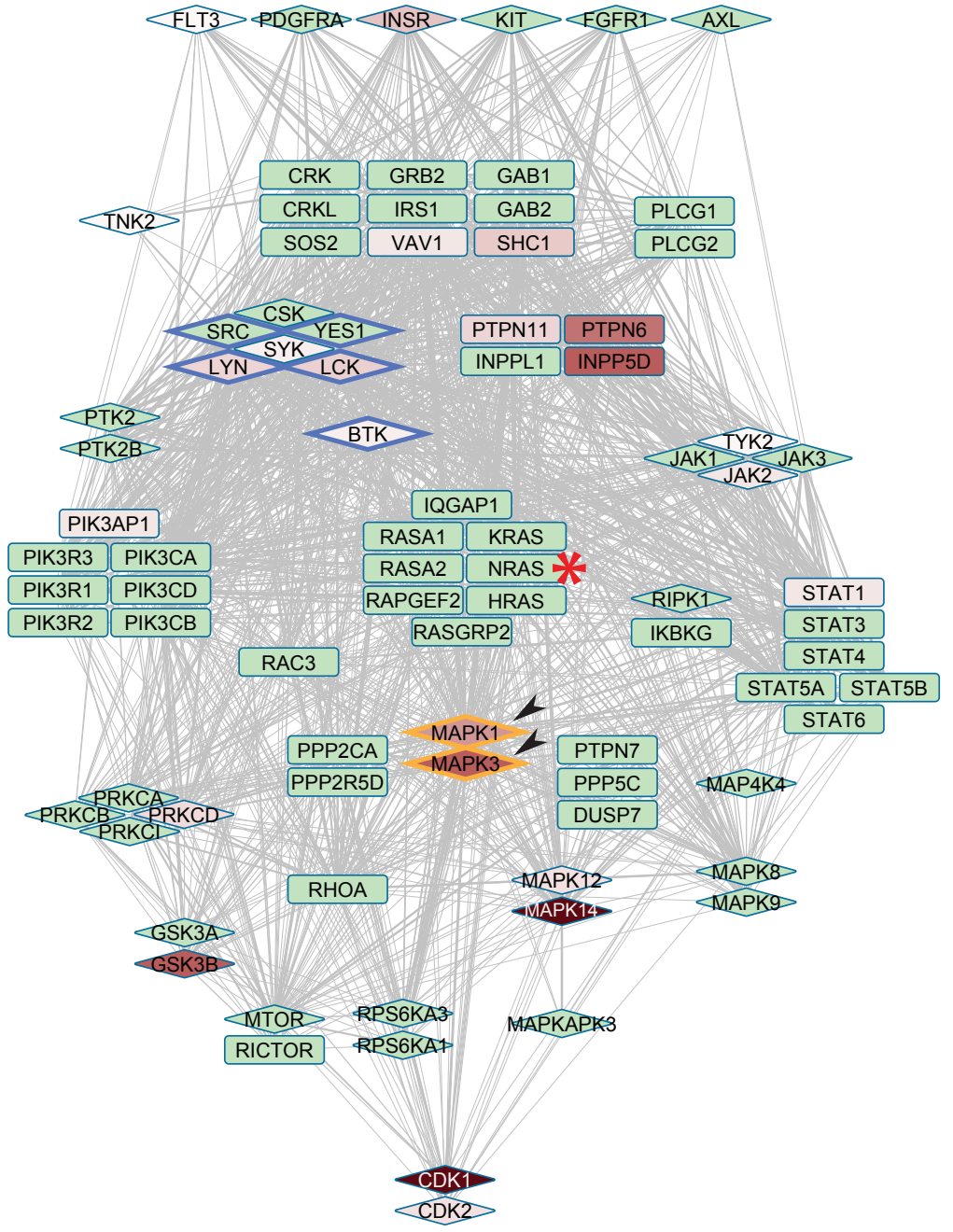
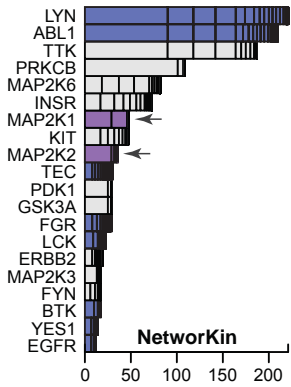
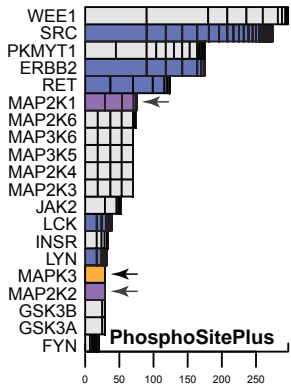
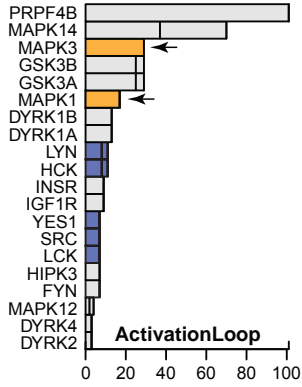
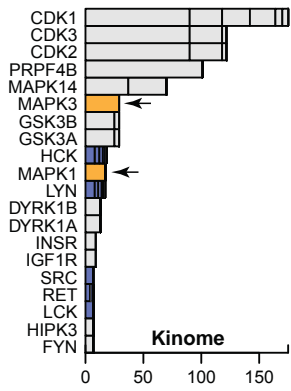
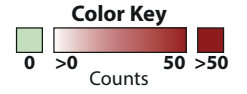
**ibrutinib**

- BTK
- EGFR
- HCK
- LCK
- LYN
- MAP2K5
- PTK6
- RIPK2
- RIPK3
- SRC
- TEC
- YES1
- ▲ ABL
- ▲ BLK
- ▲ BMX
- ▲ BRK
- ▲ ERBB2
- ▲ FGR
- ▲ FRK
- ▲ ITK
- ▲ JAK3
- ▲ RET

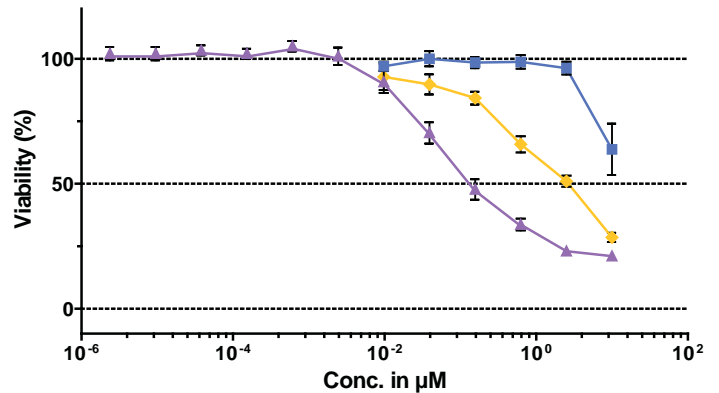
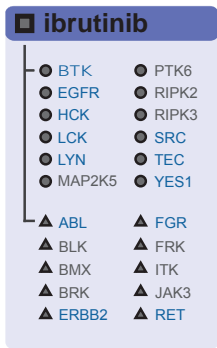


**Supplemental Figure S3M. Analysis of AML cell line OCI-AML3 lacking a known driver mutation.**  
Arrows/arrow heads indicate high-ranking kinases of the MAPK1/3 pathway; red asterisk: NRAS mutated.

# THP-1



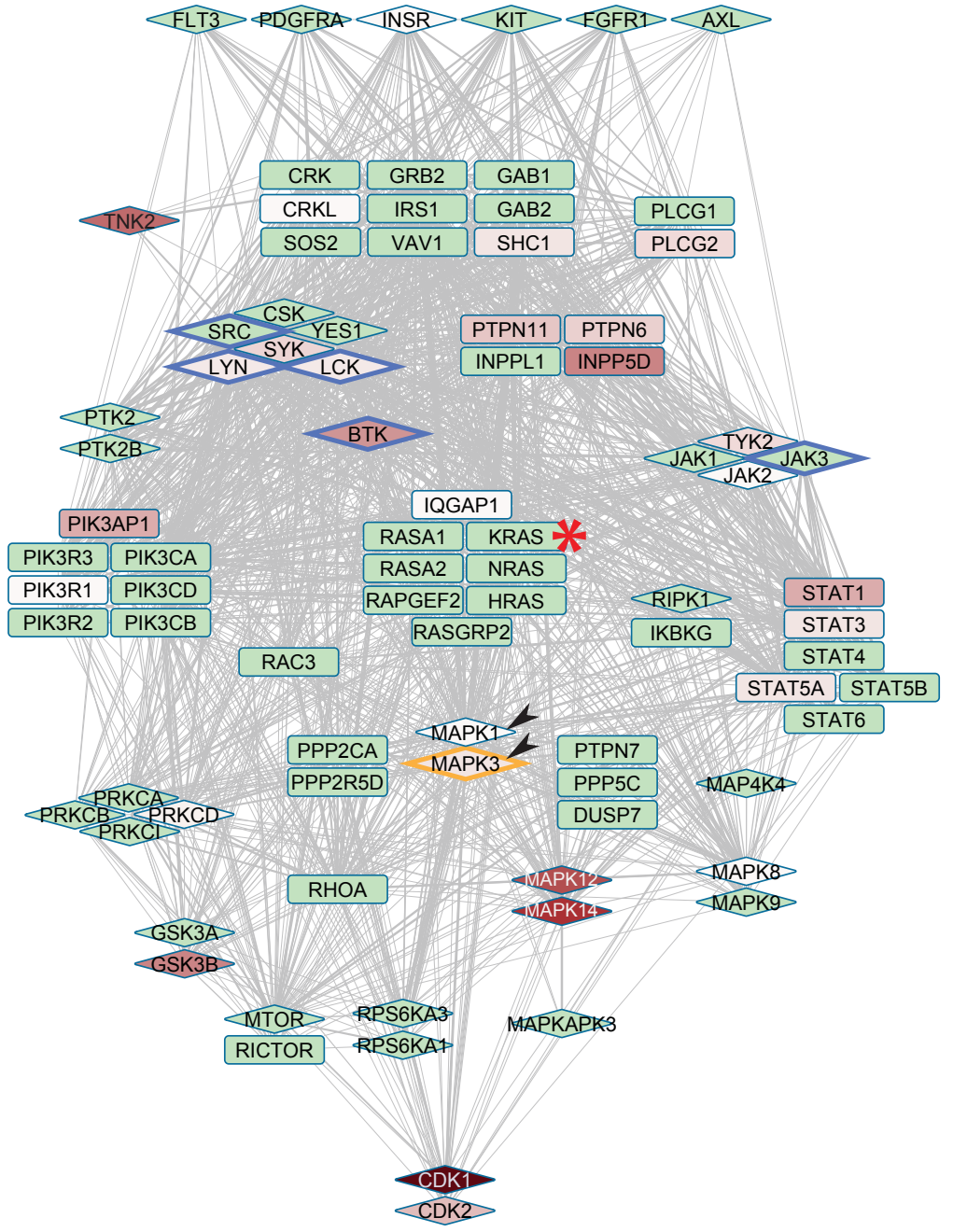
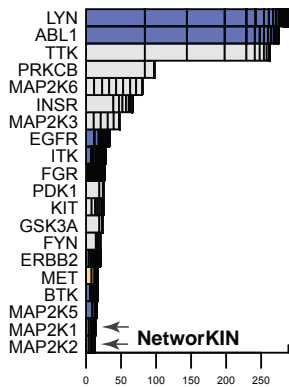
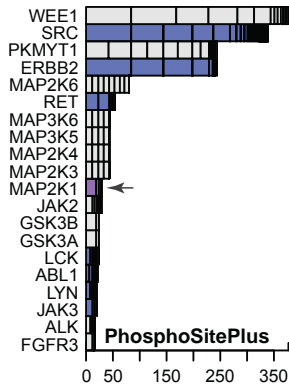
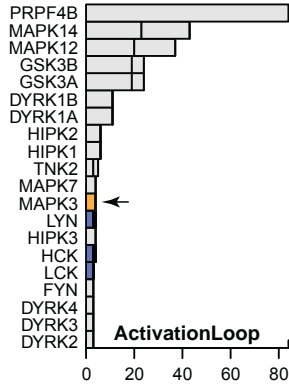
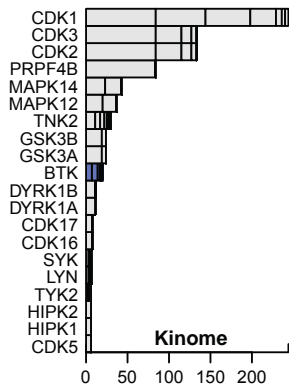
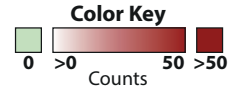
● by chemical proteomics  
▲ by cell-free assay  
**BOLD**: intended drug target  
**PLAIN**: off-target



**Supplemental Figure S3N. Analysis of AML cell line THP-1 lacking a known driver mutation.**

Arrows/arrow heads indicate high-ranking kinases of the MAPK1/3 pathway; red asterisk: NRAS mutated.

# NB4



**GDC-0994**

- MAPK1 ● MET
- MAPK3 ● TAOK1
- CAMK1G

**Binimetinib**

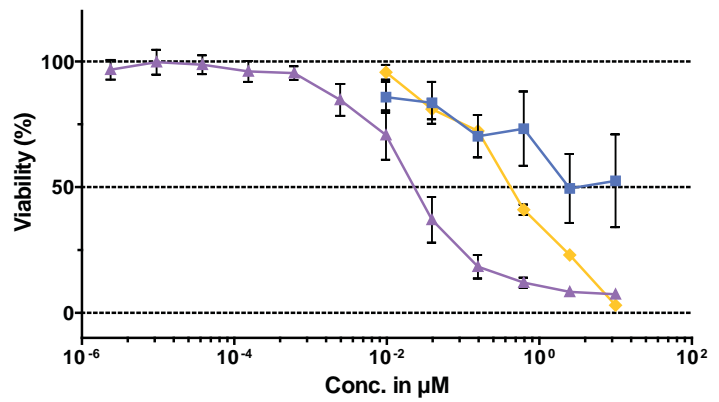
- MAP2K1 ● MAP2K2

● by chemical proteomics  
▲ by cell-free assay  
BOLD: intended drug target  
PLAIN: off-target

**ibrutinib**

- BTK ● PTK6
- EGFR ● RIPK2
- HCK ● RIPK3
- LCK ● SRC
- LYN ● TEC
- MAP2K5 ● YES1
- ▲ ABL ▲ FGR
- ▲ BLK ▲ FRK
- ▲ BMX ▲ ITK
- ▲ BRK ▲ JAK3
- ▲ ERBB2 ▲ RET

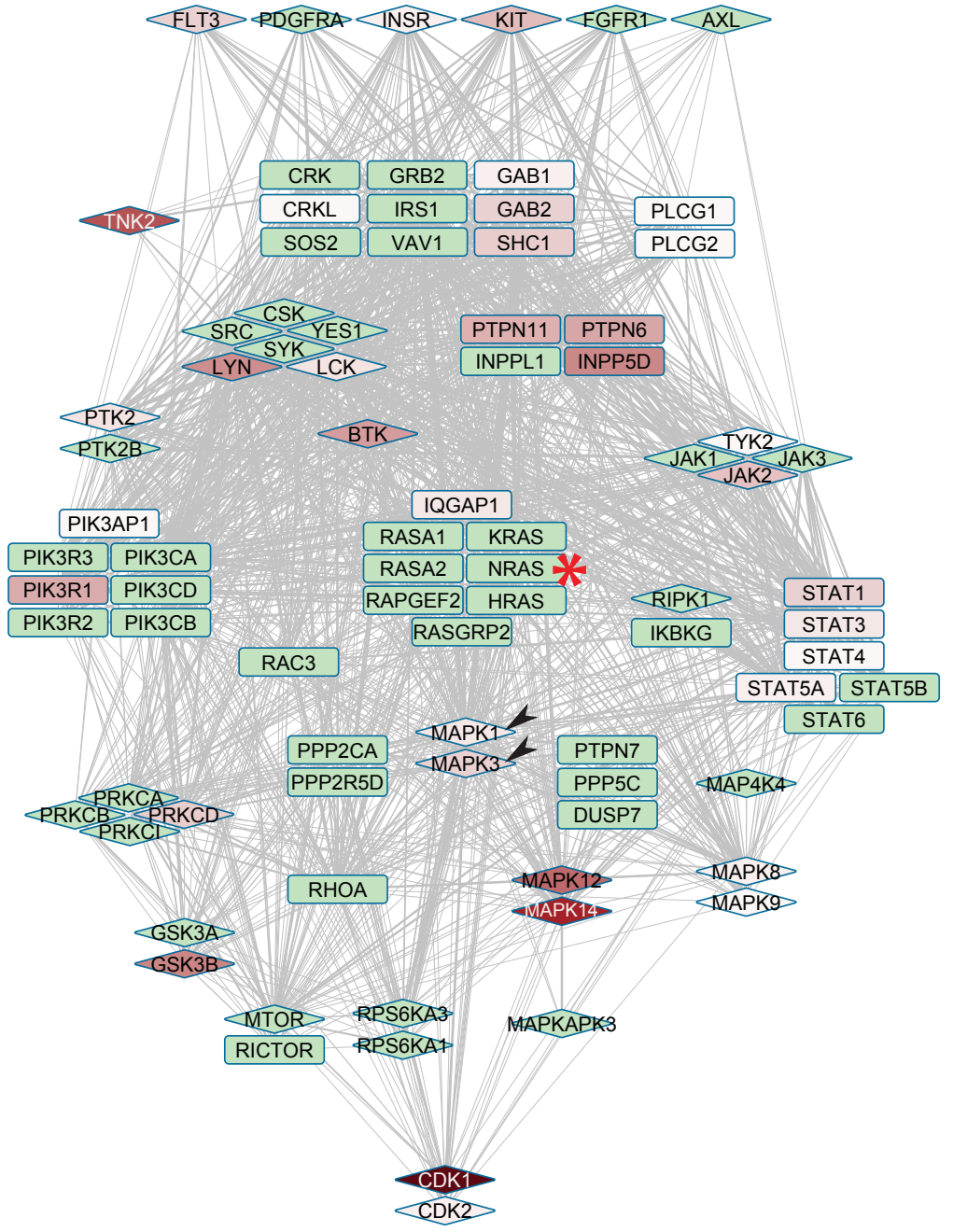
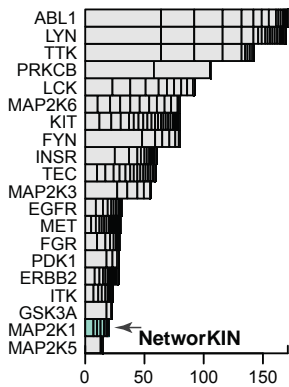
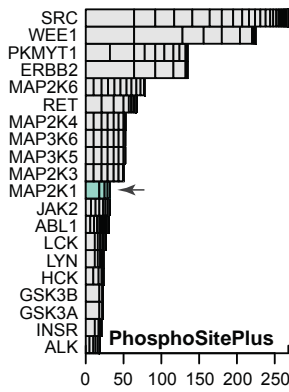
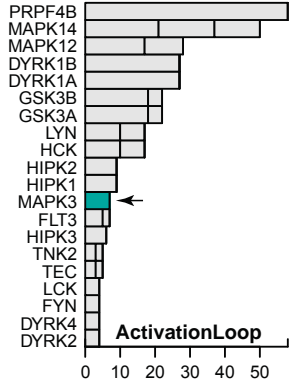
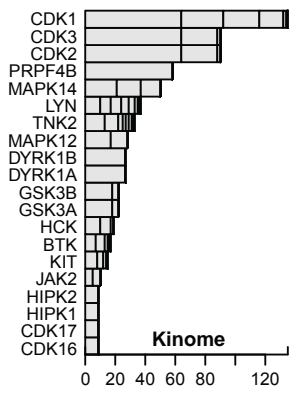
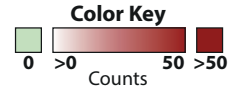
◆ GDC-0994  
▲ Binimetinib  
■ Ibrutinib



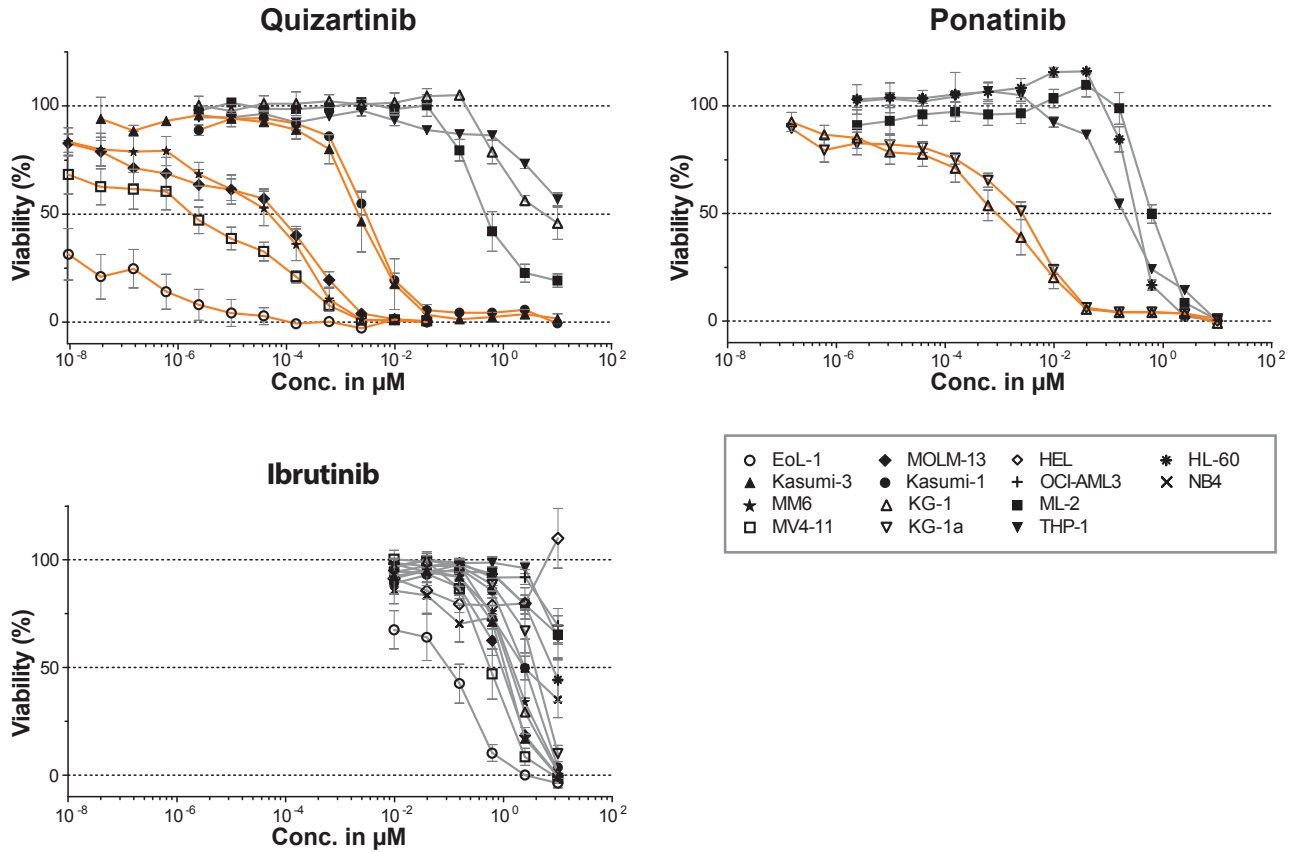
**Supplemental Figure S30. Analysis of AML cell line NB4 lacking a known driver mutation.**

Arrows/arrow heads indicate high-ranking kinases of the MAPK1/3 pathway; red asterisk: KRAS mutated.

# ME-1

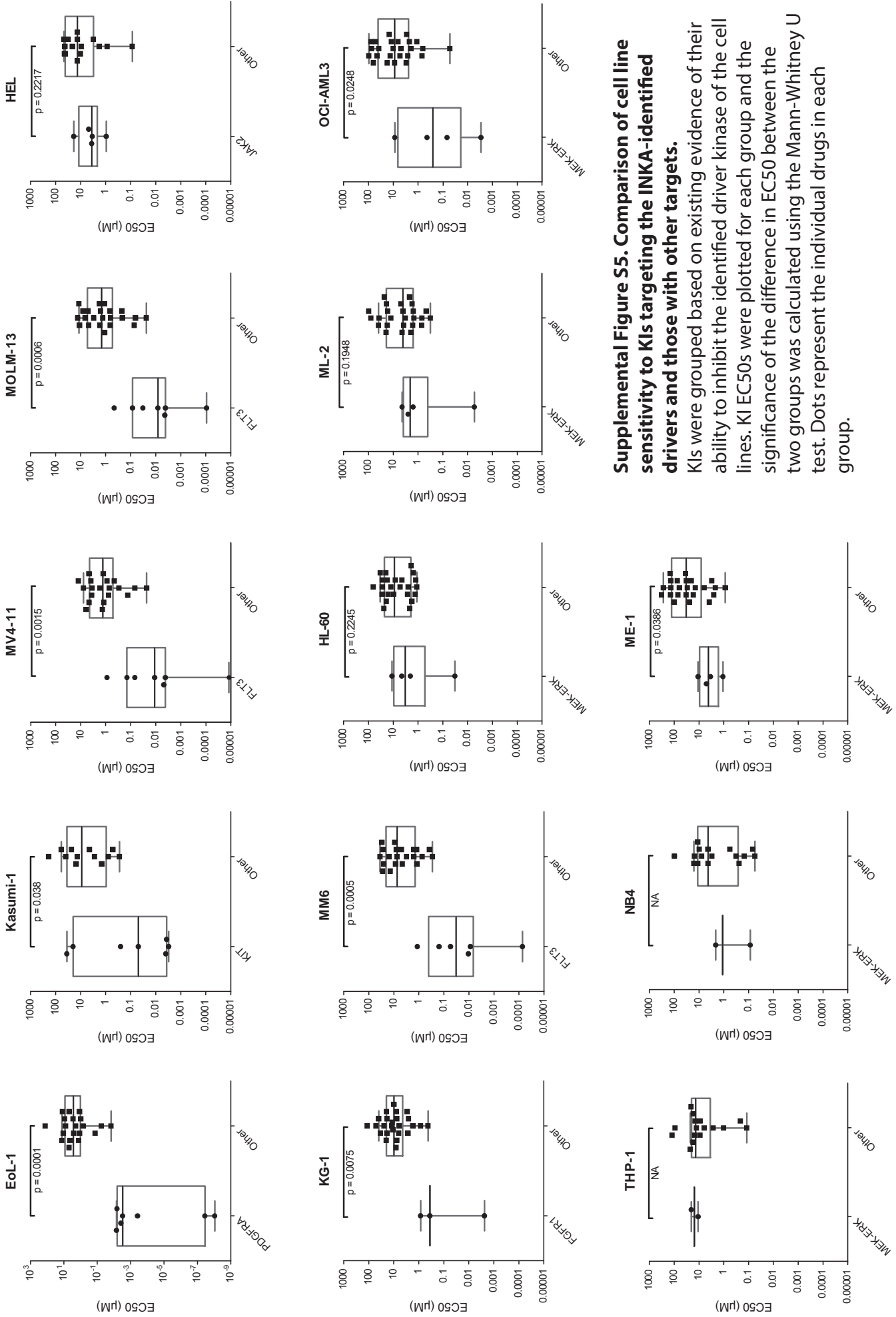


**Supplemental Figure S3P. Analysis of AML cell line ME-1 lacking a known driver mutation.**  
 Arrows/arrow heads indicate high-ranking kinases of the MAPK1/3 pathway; red asterisk: NRAS mutated.



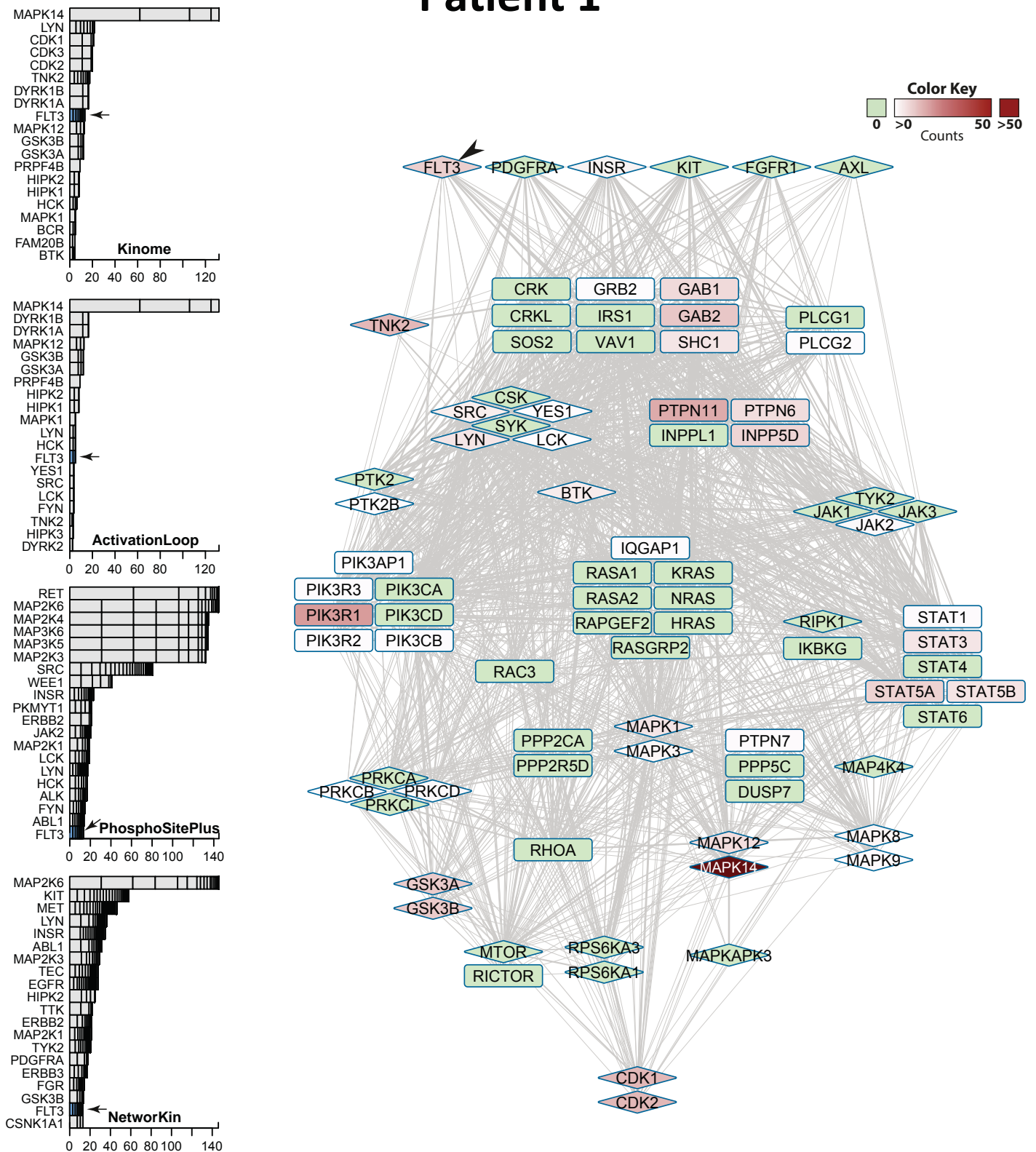
**Supplemental Figure S4. Drug response curves for quizartinib, ponatinib, and ibrutinib.**

Graphs show the drug response curves of cell lines for a drug. Cell lines with an indicated driver kinase targeted by a drug (orange line) show drug responses at lower drug concentrations, compared to control cell lines without a target kinase (grey line). The drug ibrutinib was used as a control, with similar responses for all cell lines.



**Supplemental Figure S5. Comparison of cell line sensitivity to KIs targeting the INKA-identified drivers and those with other targets.** KIs were grouped based on existing evidence of their ability to inhibit the identified driver kinase of the cell lines. KI EC50s were plotted for each group and the significance of the difference in EC50 between the two groups was calculated using the Mann-Whitney U test. Dots represent the individual drugs in each group.

# Patient 1

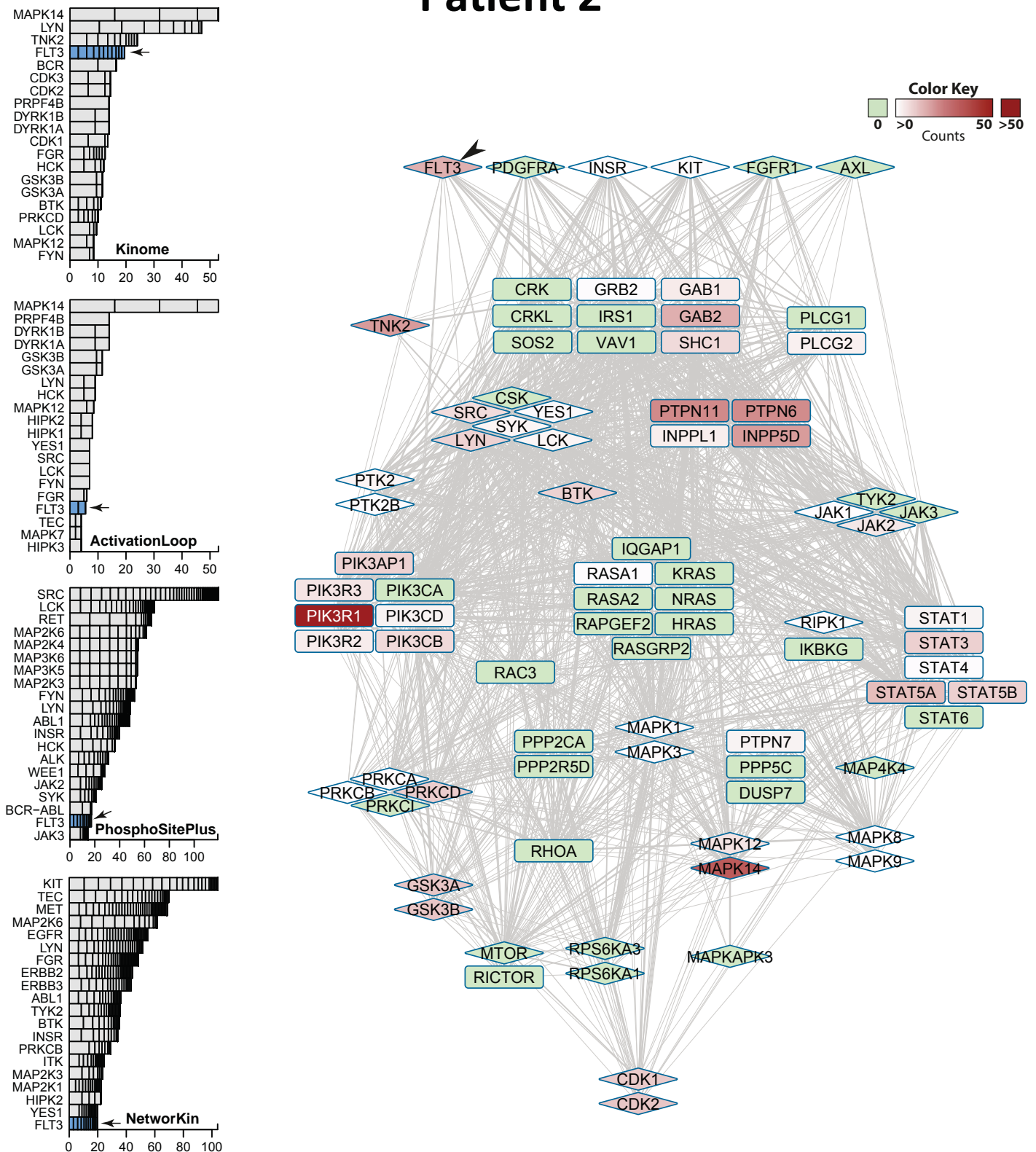


## Supplementary Figure S6A. Summary of phosphoproteomic analysis of blood sample from patient 1.

*Left.* Overview of bar graphs ranking kinases according to their own phosphorylation (*kinome*, phosphorylation anywhere on the protein; *activation loop*, phosphorylation of the activation segment) and according to the phosphorylation of their substrates (*PhosphoSitePlus*, known substrates cataloged in the PhosphoSitePlus database; *NetworKin*, predicted substrates assigned by the NetworKin algorithm). Arrows and blue highlighting indicate the FLT3 kinase.

*Right.* Kinases inferred to be active were mapped in Cytoscape onto pathways in a signaling network (edges from the STRING knowledge base). Node fill colors were rendered according to spectral counts as a proxy of activity. Arrow head indicates the FLT3 kinase.

## Patient 2

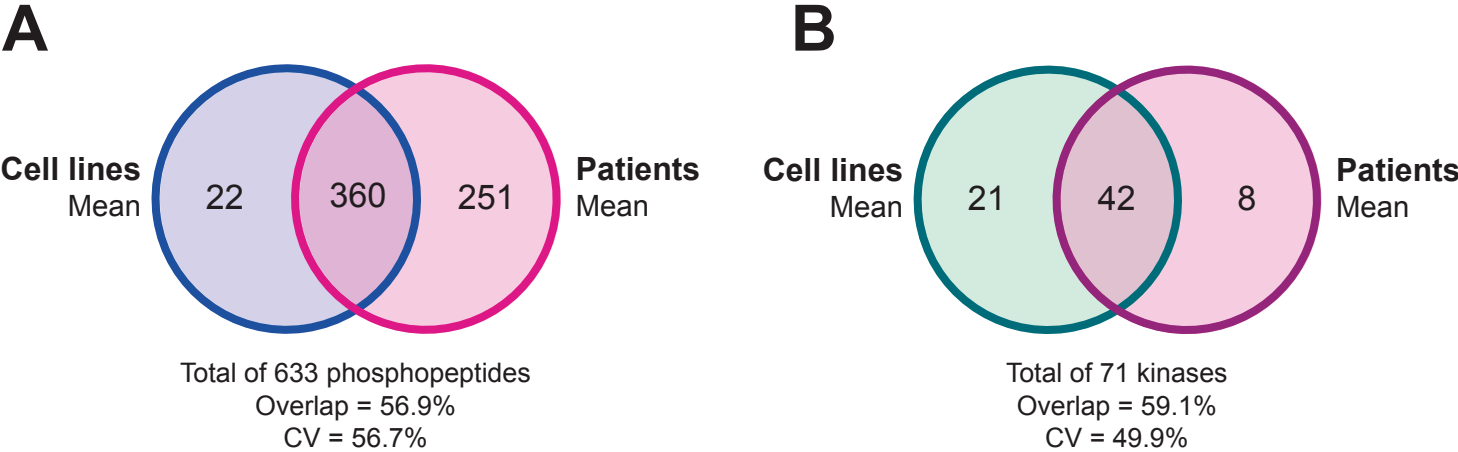


### Supplementary Figure S6B. Summary of phosphoproteomic analysis of blood sample from patient 2.

*Left.* Overview of bar graphs ranking kinases according to their own phosphorylation (*kinome*, phosphorylation anywhere on the protein; *activation loop*, phosphorylation of the activation segment) and according to the phosphorylation of their substrates (*PhosphoSitePlus*, known substrates cataloged in the PhosphoSitePlus database; *NetworkKin*, predicted substrates assigned by the NetworkKin algorithm). Arrows and blue highlighting indicate the FLT3 kinase.

*Right.* Kinases inferred to be active were mapped in Cytoscape onto pathways in a signaling network (edges from the STRING knowledge base). Node fill colors were rendered according to spectral counts as a proxy of activity. Arrow head indicates the FLT3 kinase.





**Supplemental Figure S7. Comparison between FLT3-ITD mutant cell line data and FLT3-ITD mutant patient data.**

A. Overlap of phosphopeptides found in the three FLT3-ITD positive cell lines MV4-11, MOLM-13, and Kasumi-6 (blue) and those found in two FLT3-ITD positive patient samples (red).  
B. Overlap of phosphokinases found in FLT3-ITD positive cell lines and those found in FLT3-ITD positive patient samples.

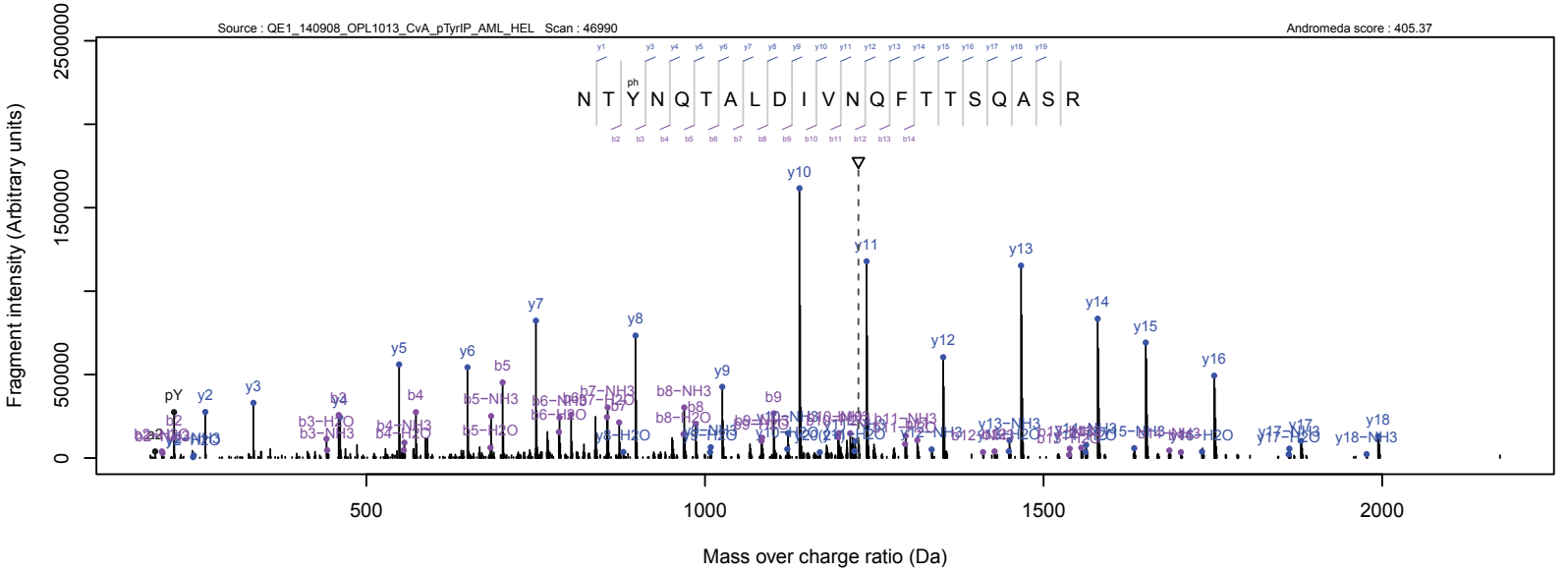
# Supplemental Figure 8

Annotated MS/MS spectra of phosphopeptides  
identified in AML cell lines

NTY(ph)NQTALDIVNQFTTSQASR-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 46990

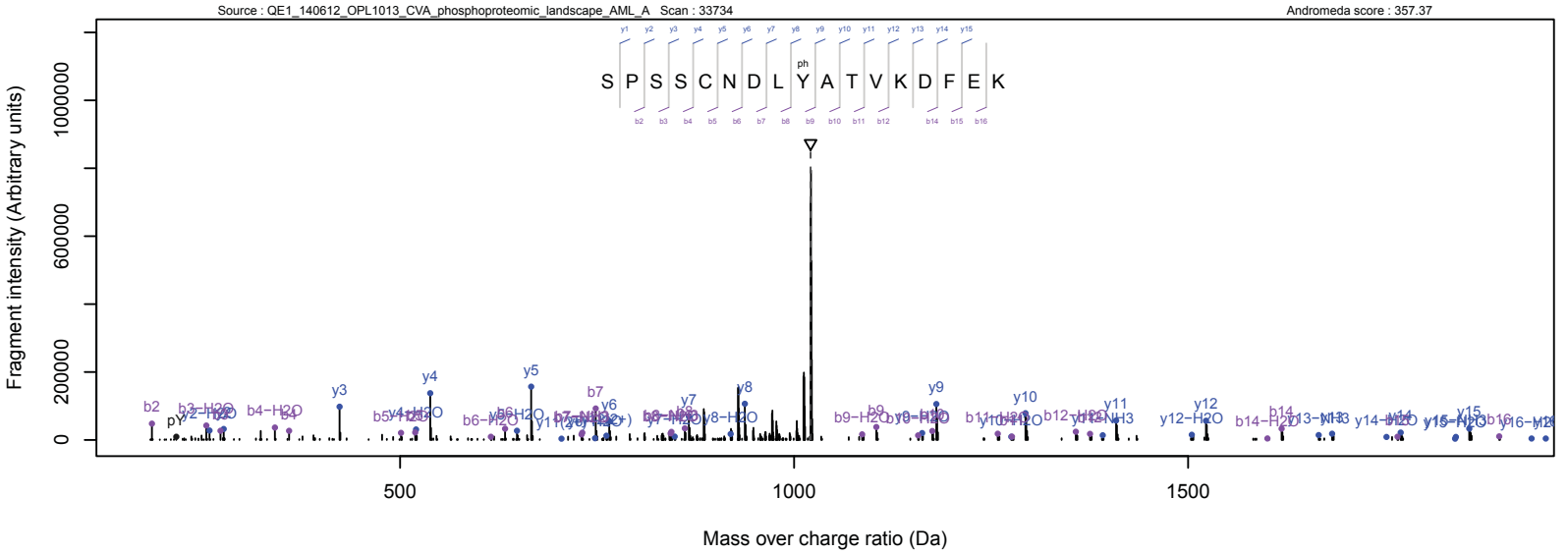
Andromeda score : 405.37



SPSSCNDLY(ph)ATVKDFEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33734

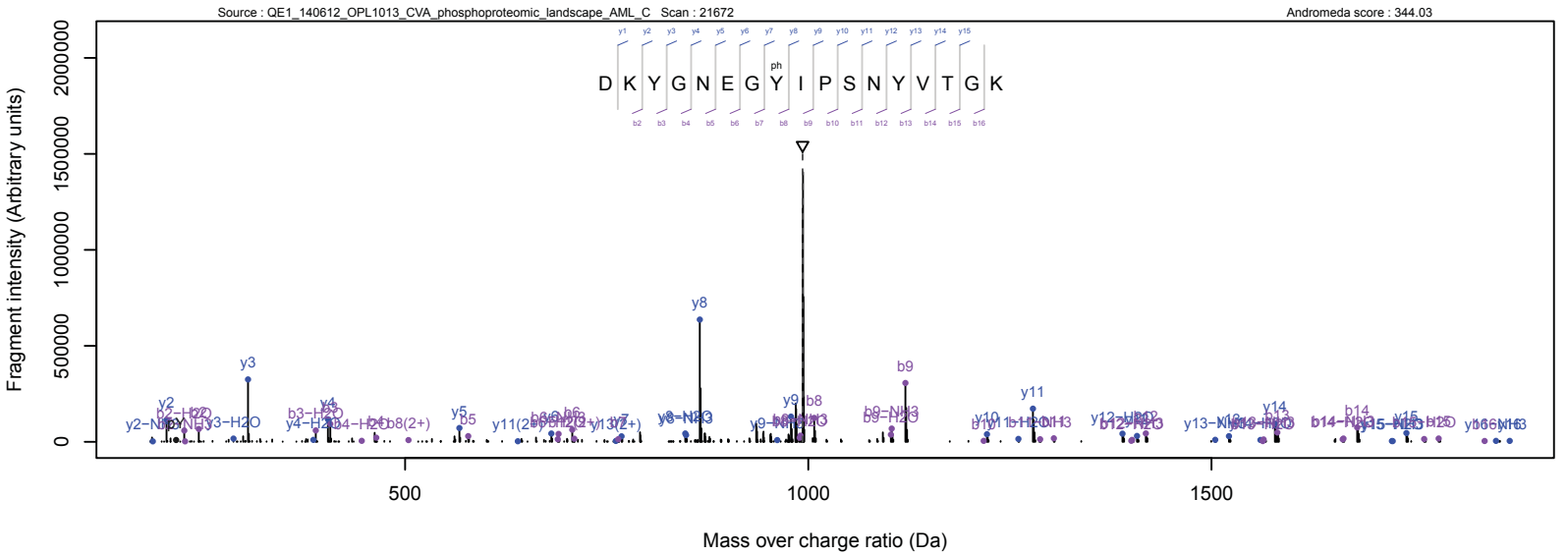
Andromeda score : 357.37



DKYGNEGY(ph)IPSNYVTGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21672

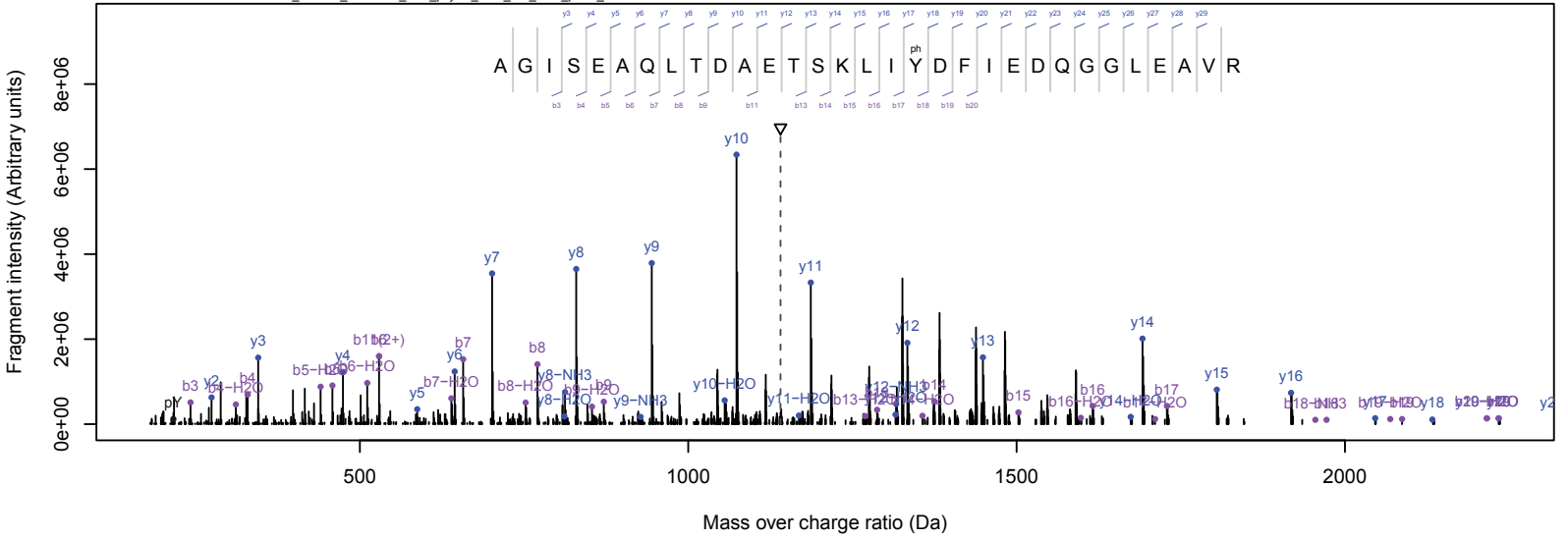
Andromeda score : 344.03



AGISEAQLTDAETSKLIY(ph)DFIEDQGGLEAVR-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 45402

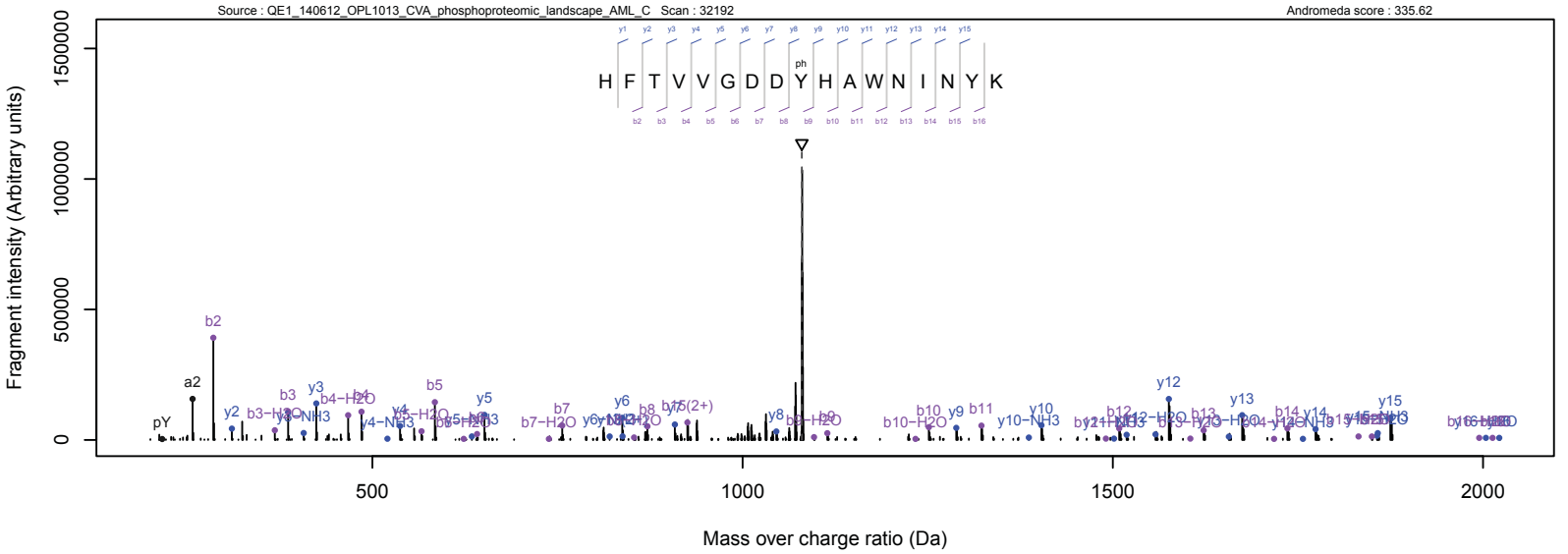
Andromeda score : 340.95



HFTVVGDDY(ph)HAWNINYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32192

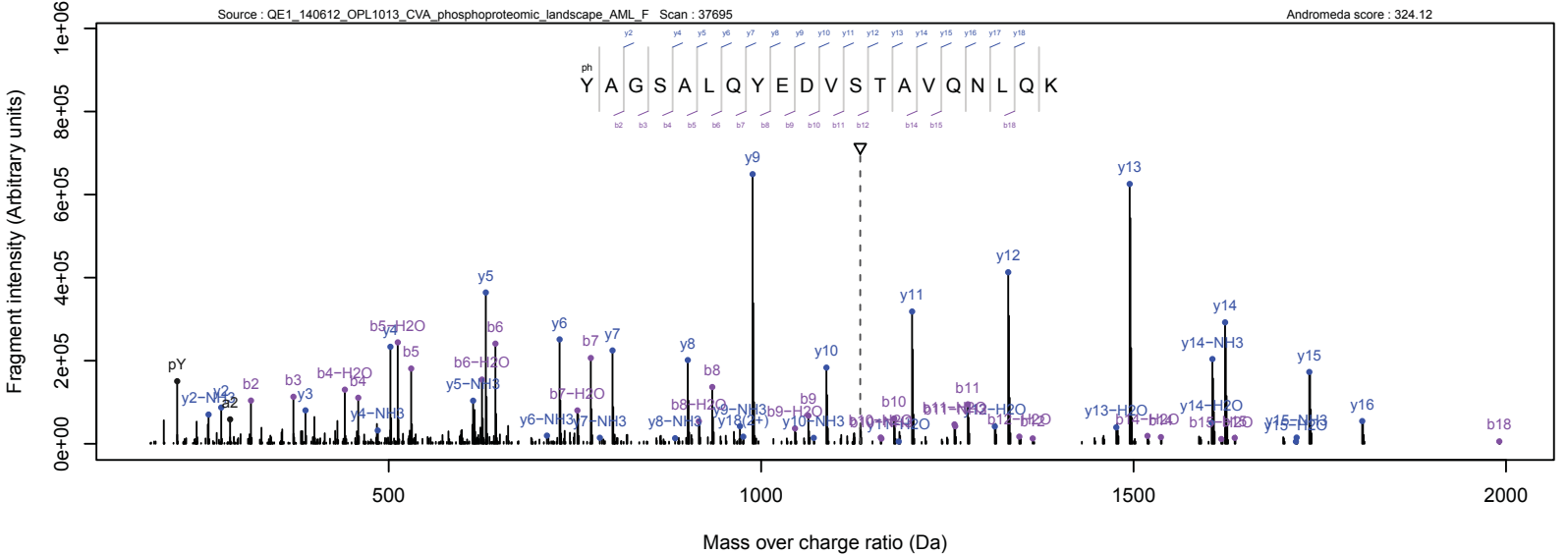
Andromeda score : 335.62



Y(ph)AGSALQYEDVSTAVQNLQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 37695

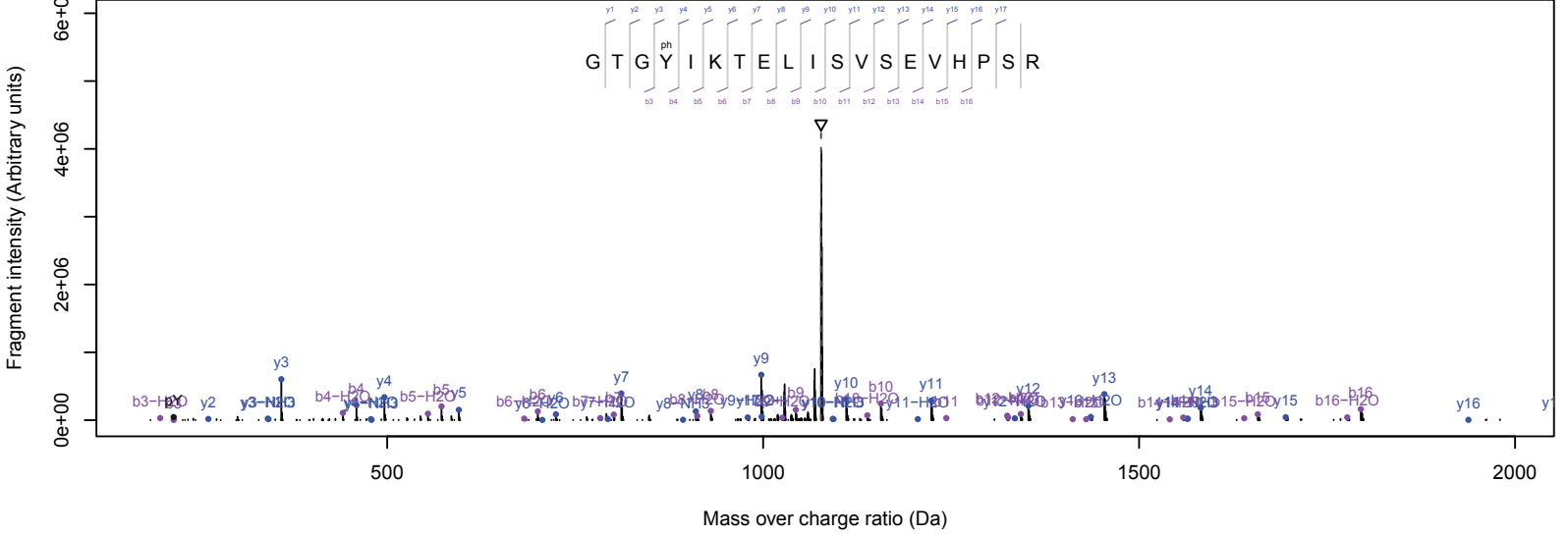
Andromeda score : 324.12



GTGY(ph)IKTELISVSEVHPSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28218

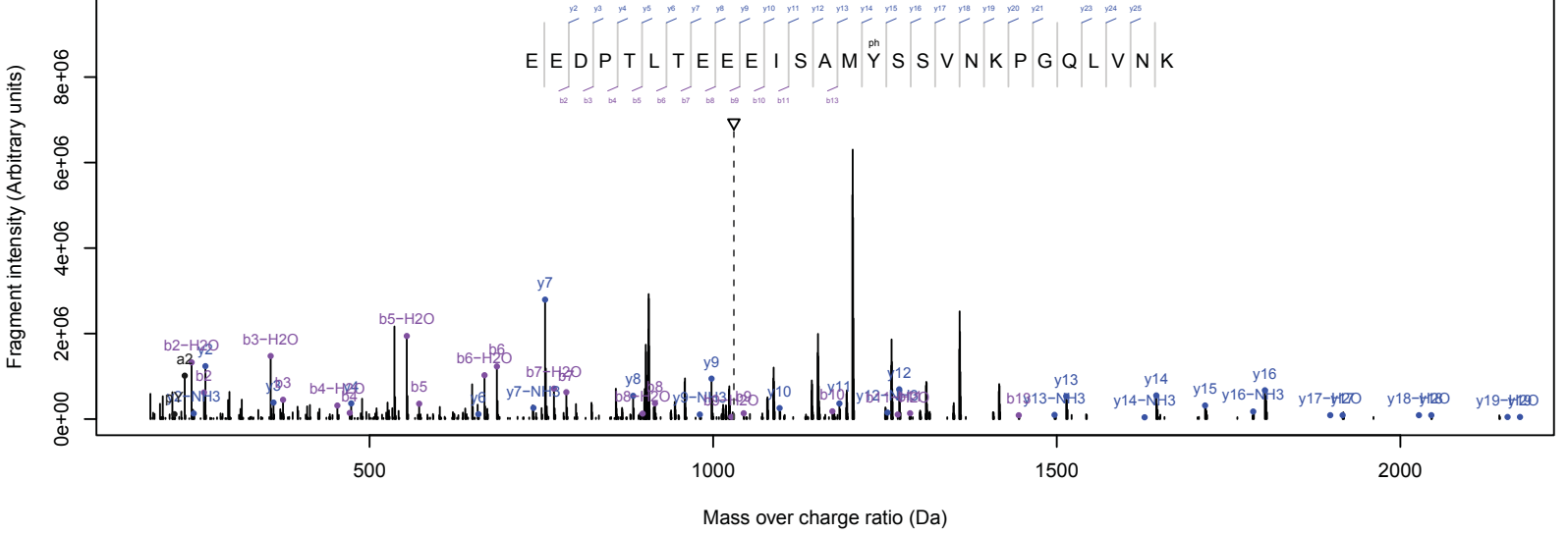
Andromeda score : 323.14



EEDPTLTEEISAMY(ph)SSVKNKPGQLVNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38611

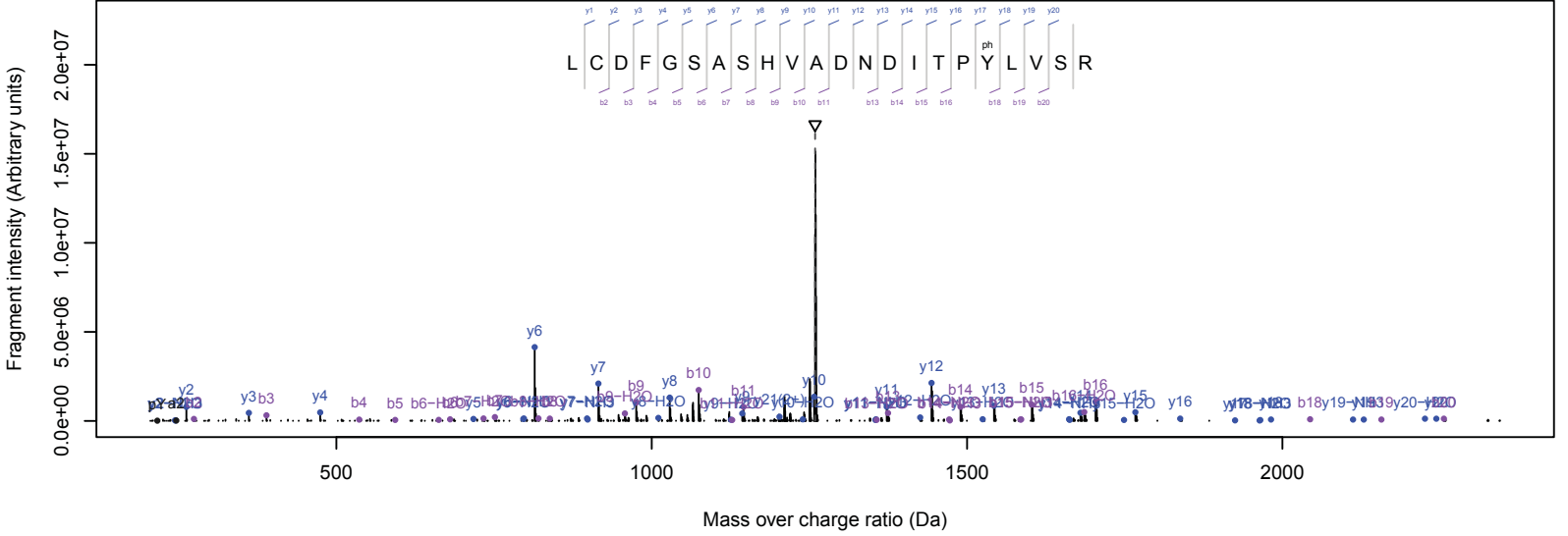
Andromeda score : 323.04



LCDFGSASHVADNDITPYLVS R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 35089

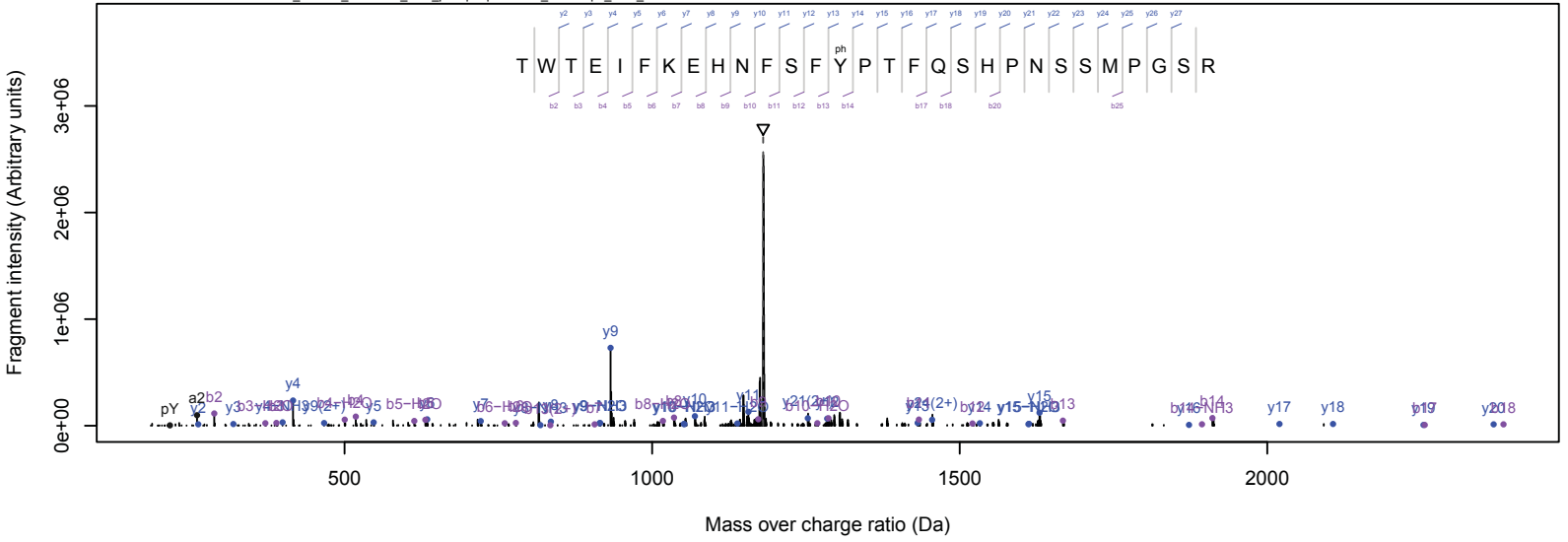
Andromeda score : 320.88



**TWTEIFKEHNFSFY(ph)PTFQSHPNSSMPGSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43699

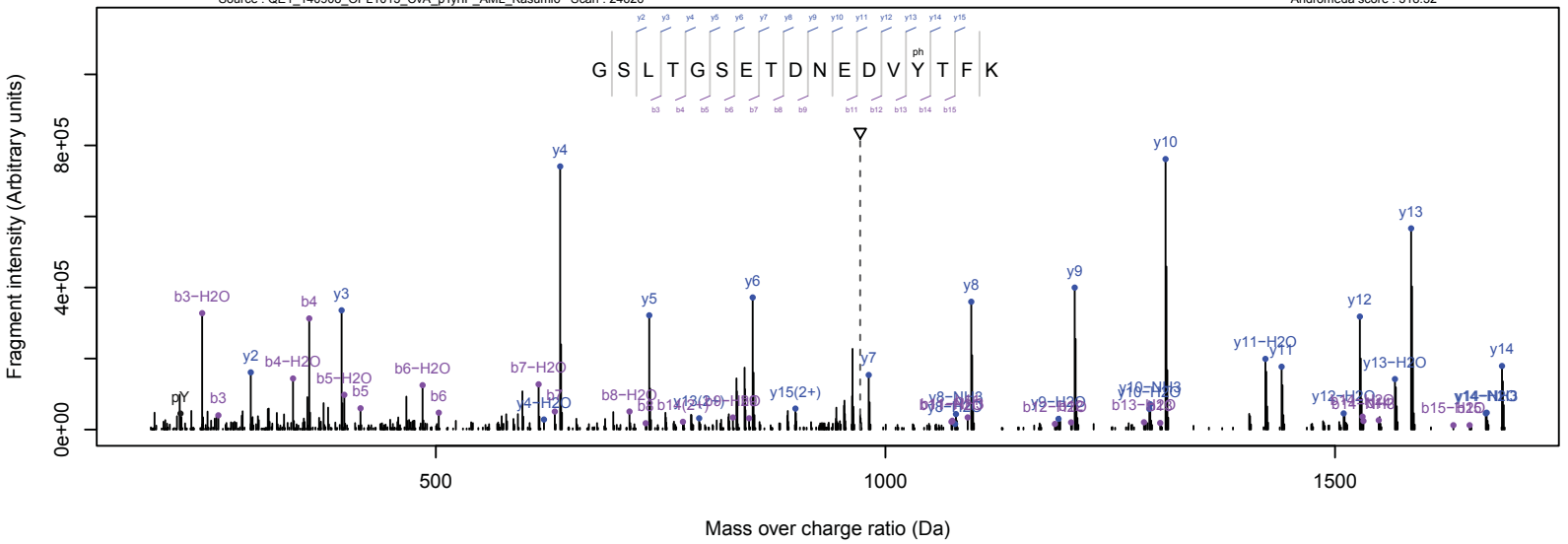
Andromeda score : 318.57



**GSLTGSETDNEDVY(ph)TFK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 24626

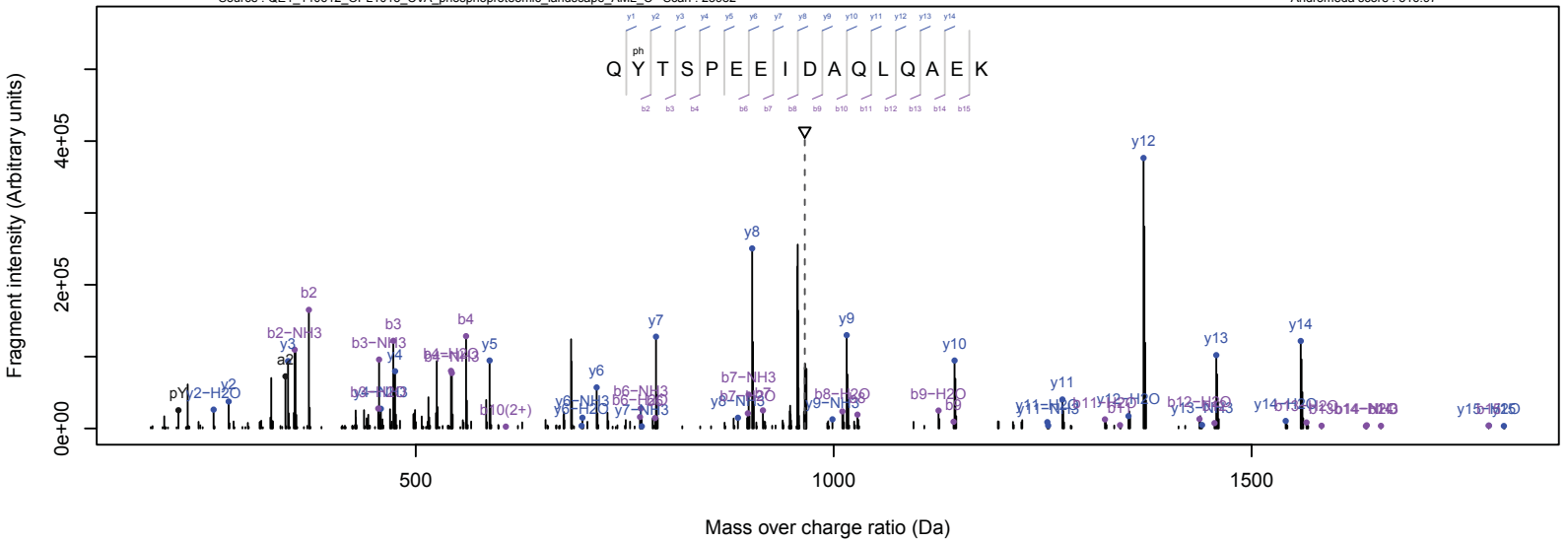
Andromeda score : 318.32



**QY(ph)TSPEEIDAQLQAEK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25982

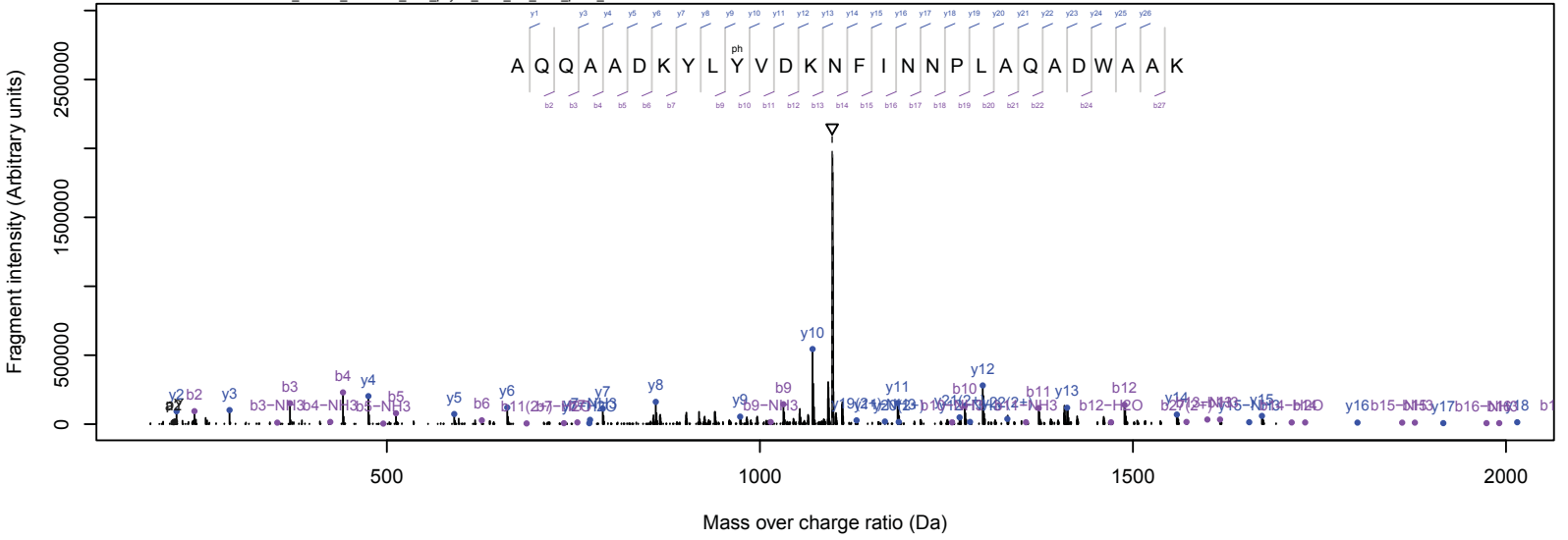
Andromeda score : 316.97



(ac)AQQAADKYL(ph)VDKNFINNPLAQADWAAK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 41777

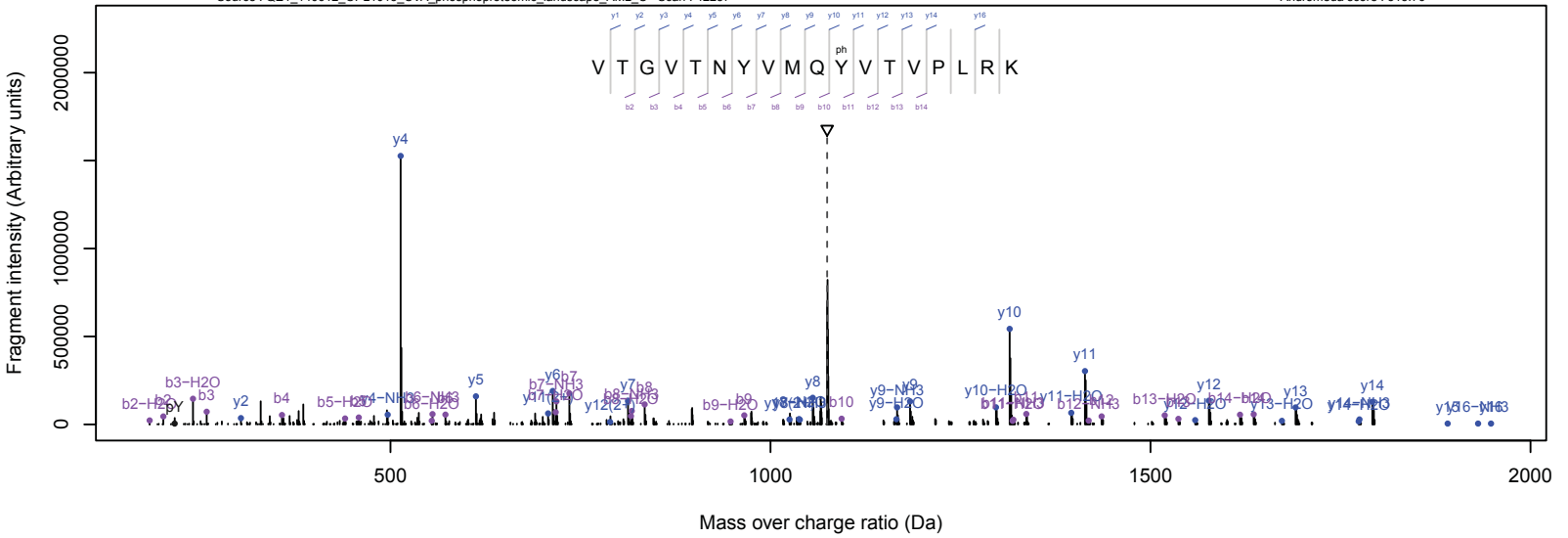
Andromeda score : 314.57



VTGVTNYVMQY(ph)VTVPLRK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42257

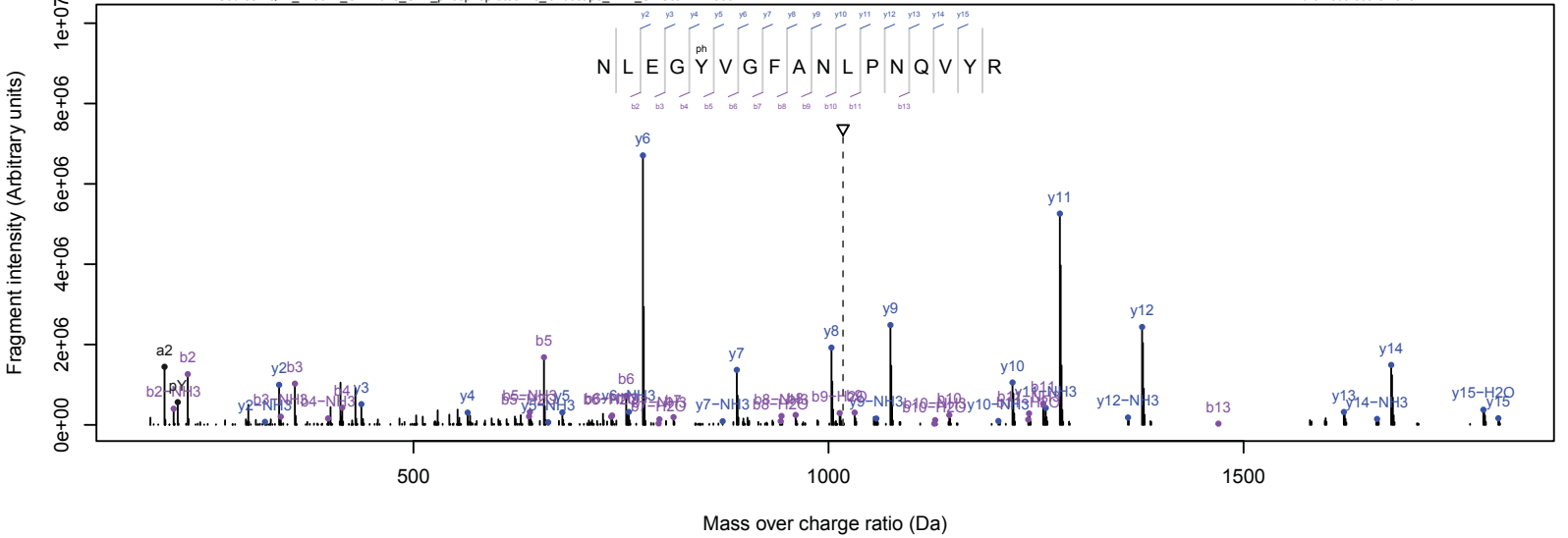
Andromeda score : 313.76



NLEGY(ph)VGfanLPNQVYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42356

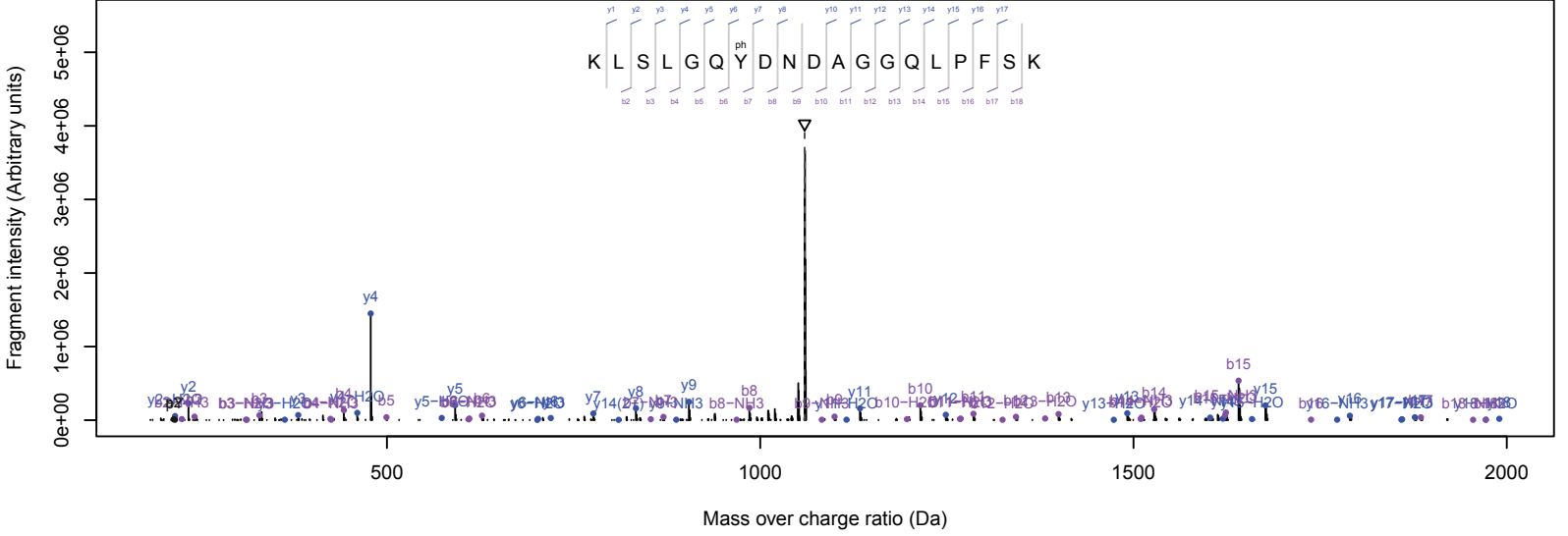
Andromeda score : 310.74



**KLSLGQY(ph)DNDAGGQLPFSK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30150

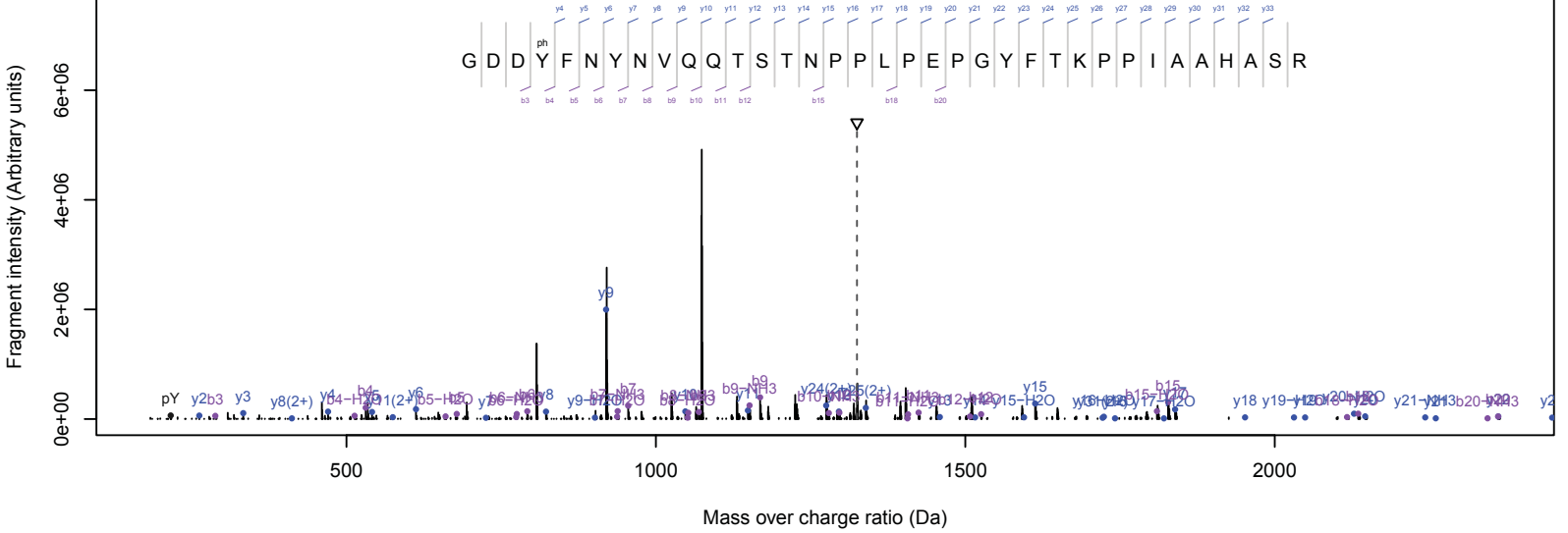
Andromeda score : 309.77



**GDDY(ph)FNYNVQQTSTNPLPEPGYFTKPPIAAHSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39738

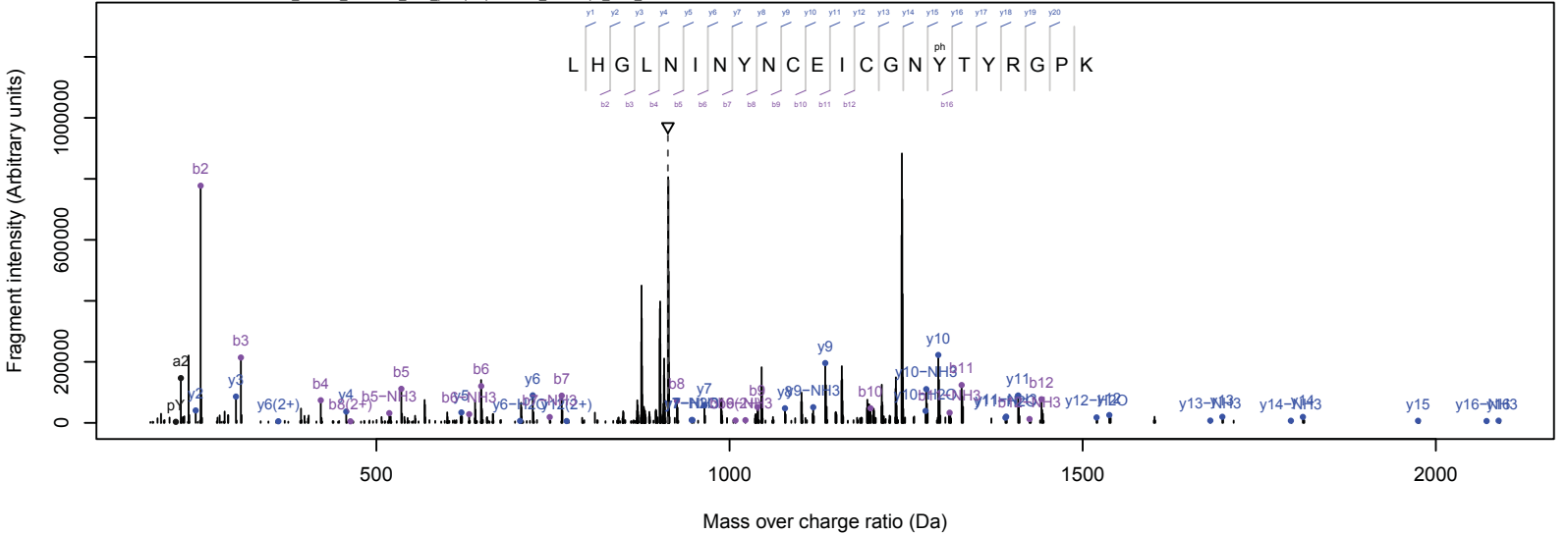
Andromeda score : 307.16



**LHGLNINYNCEICGNYPYRGPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27996

Andromeda score : 306.69



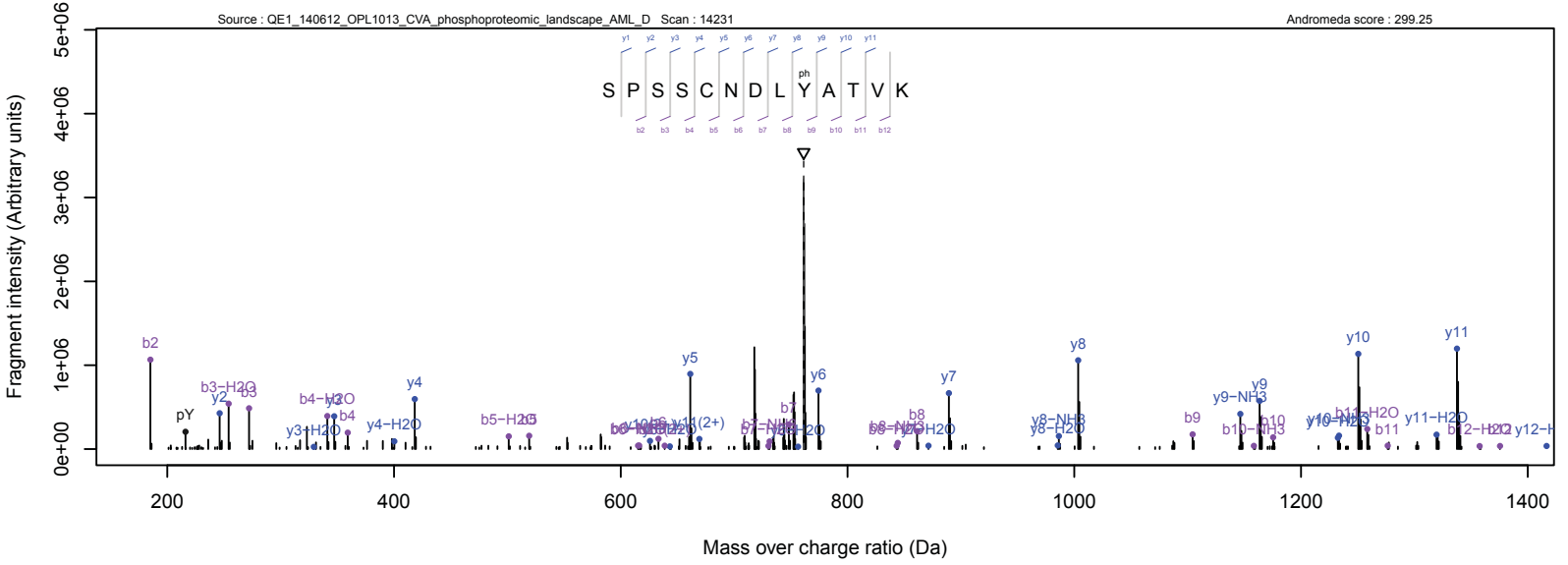




SPSSCNDLY(ph)ATVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14231

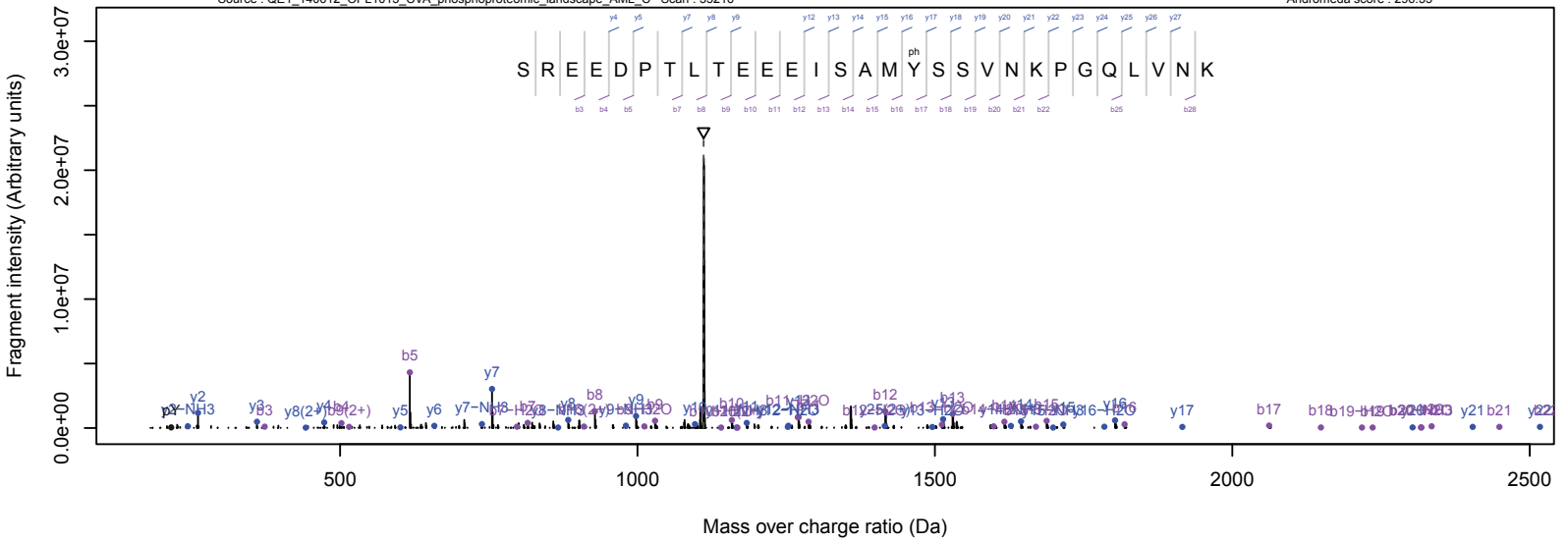
Andromeda score : 299.25



SREEDPTLTEEEISAMY(ph)SSVNKPGQLVNVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35216

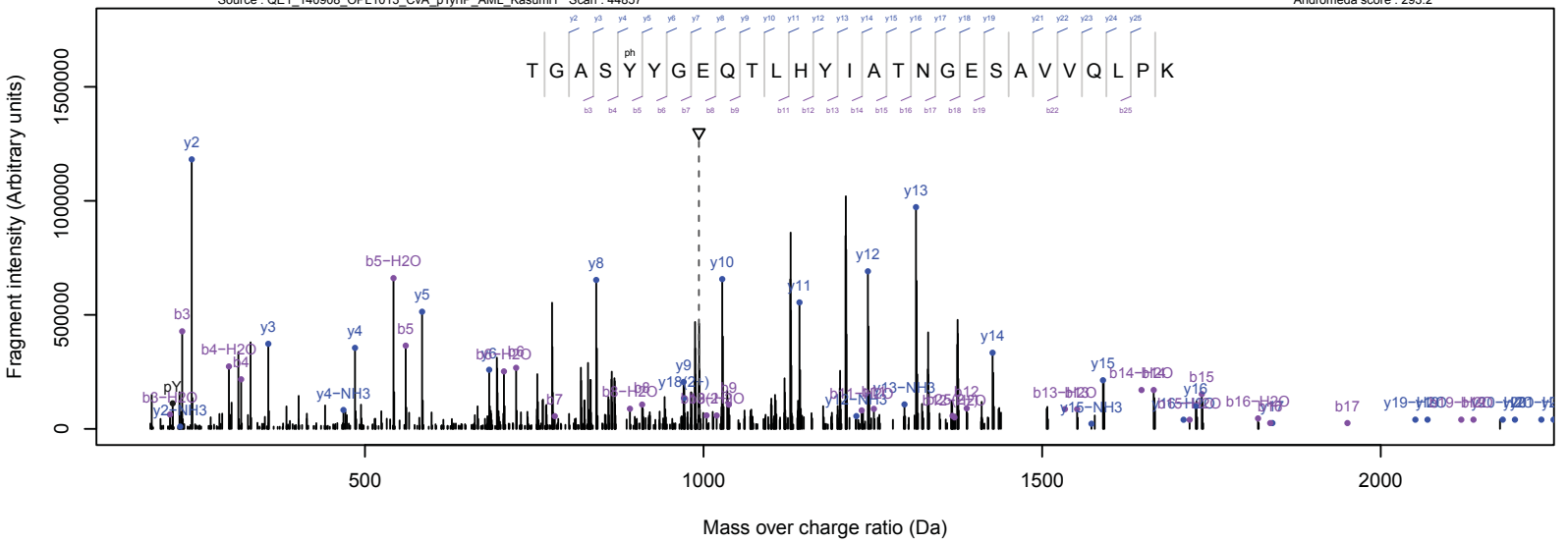
Andromeda score : 296.35



TGASY(ph)YGEQLTHYIATNGESAVVQLPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 44857

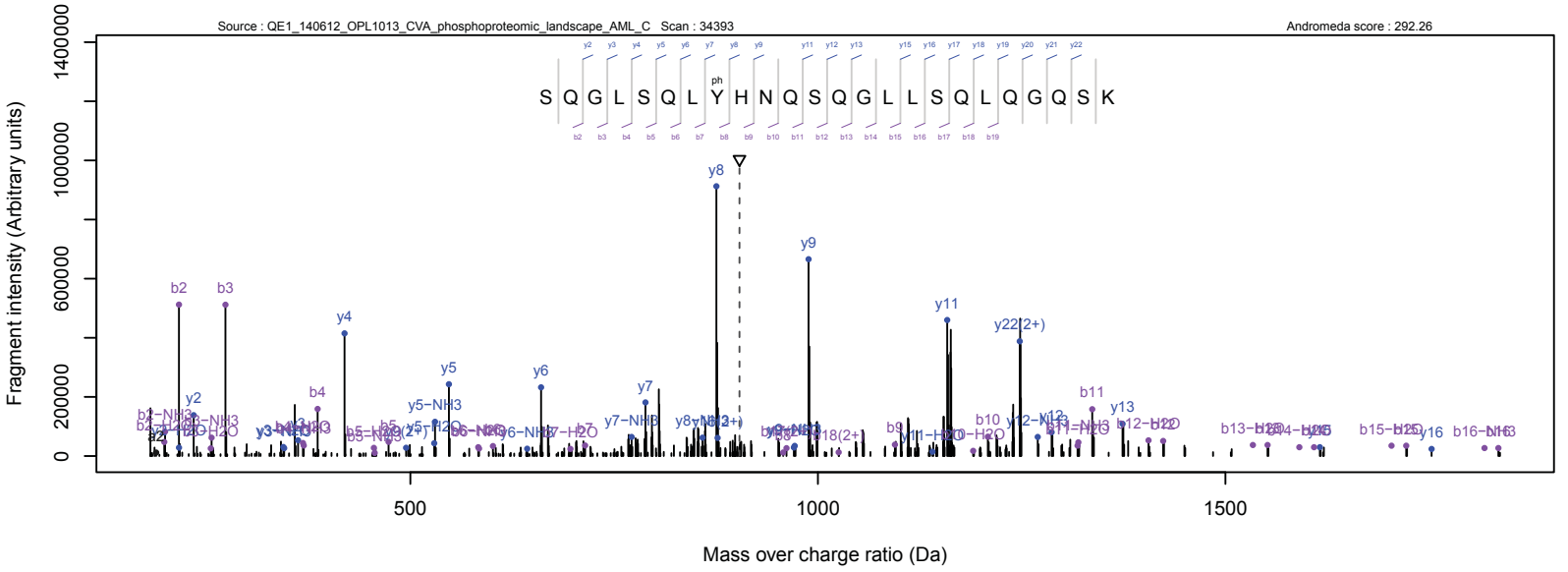
Andromeda score : 293.2



**SQGLS<sup>ph</sup>QLYHNQSQGLLSQLQGQSK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34393

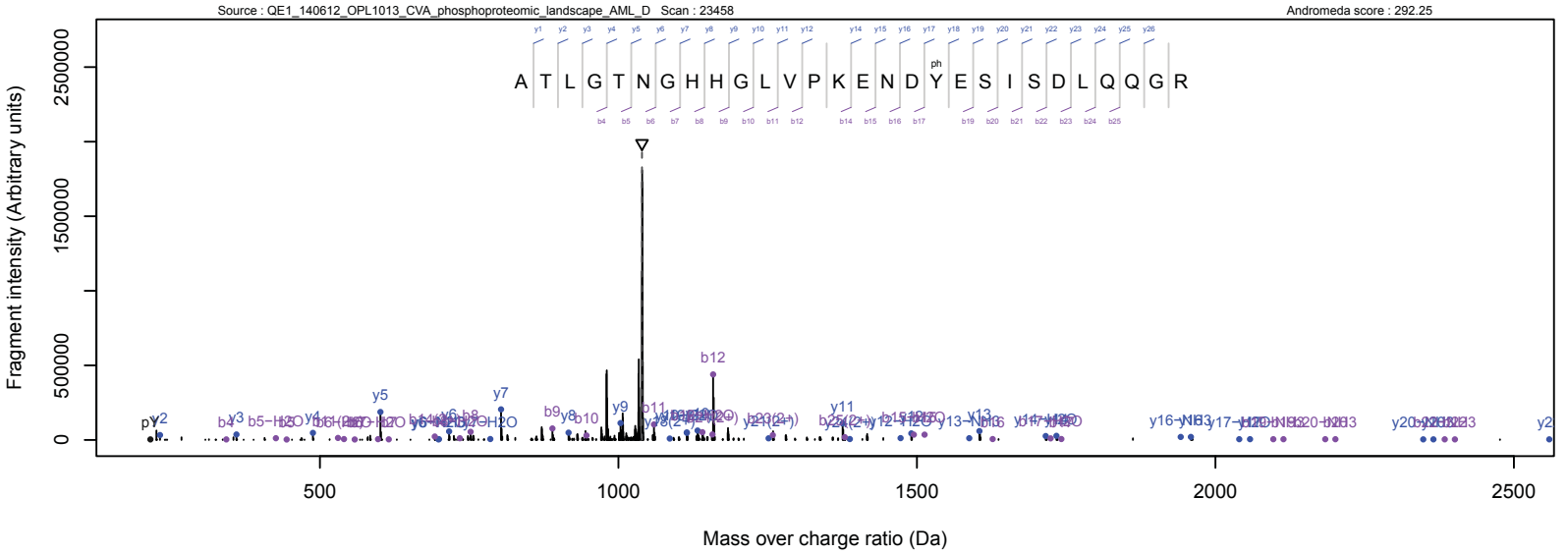
Andromeda score : 292.26



**ATLGTNGHHGLV<sup>ph</sup>PKENDYESISDLQQGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23458

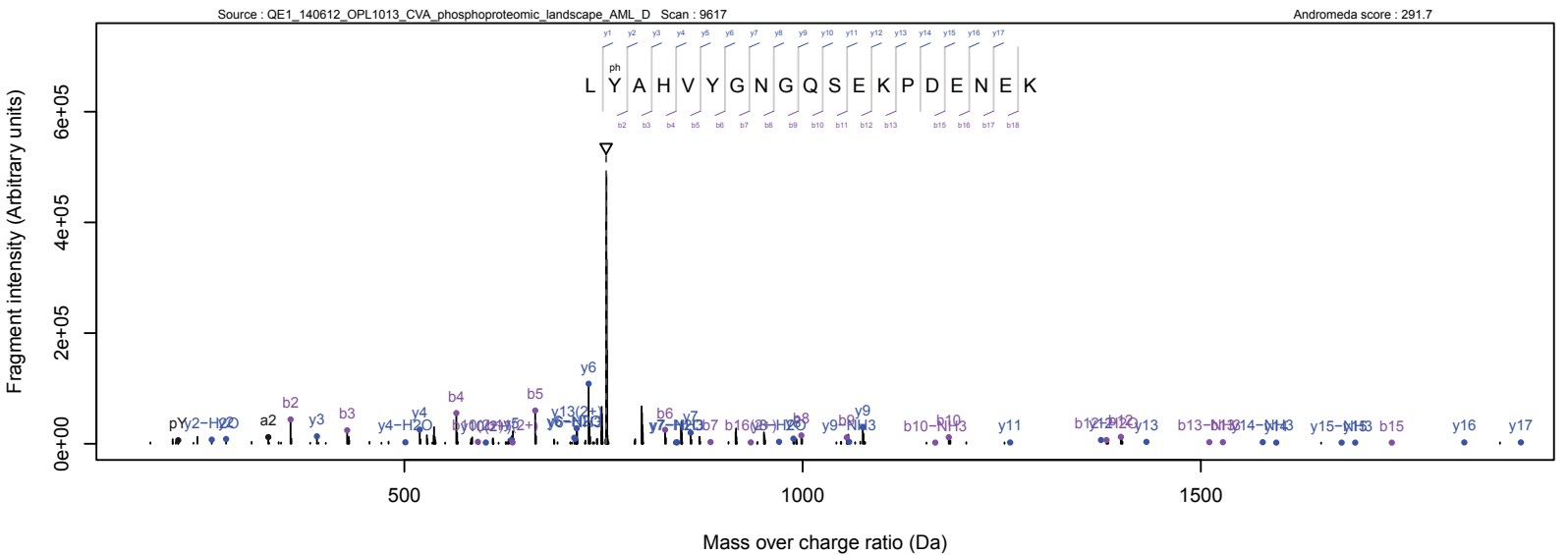
Andromeda score : 292.25



**LY<sup>ph</sup>AHVYNGQSEKPDENEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9617

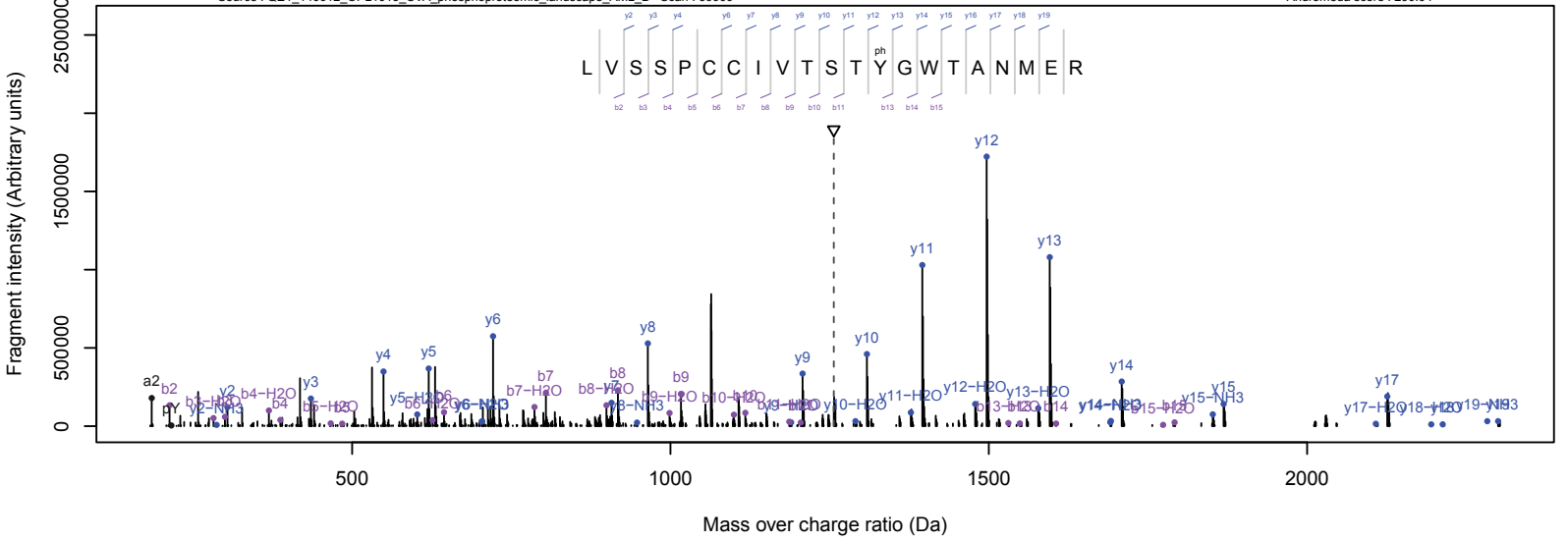
Andromeda score : 291.7



LVSSPCCIVTSTY(ph)GWTANMER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39909

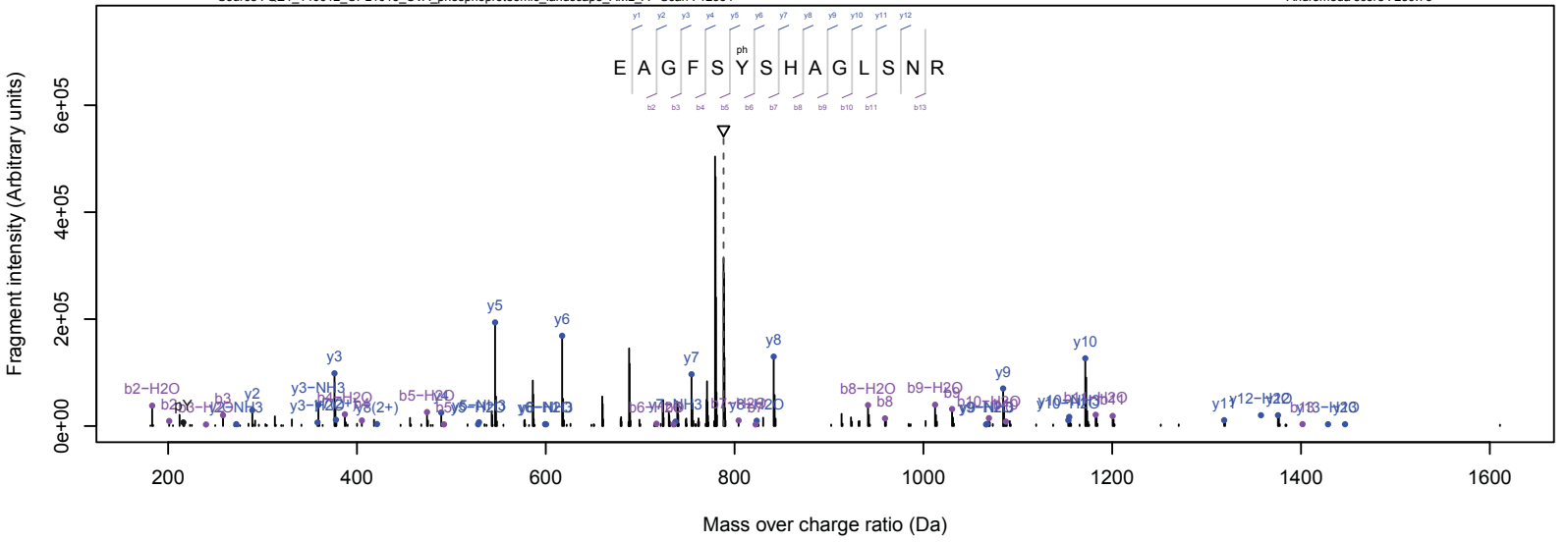
Andromeda score : 290.94



EAGFSY(ph)SHAGLSNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12334

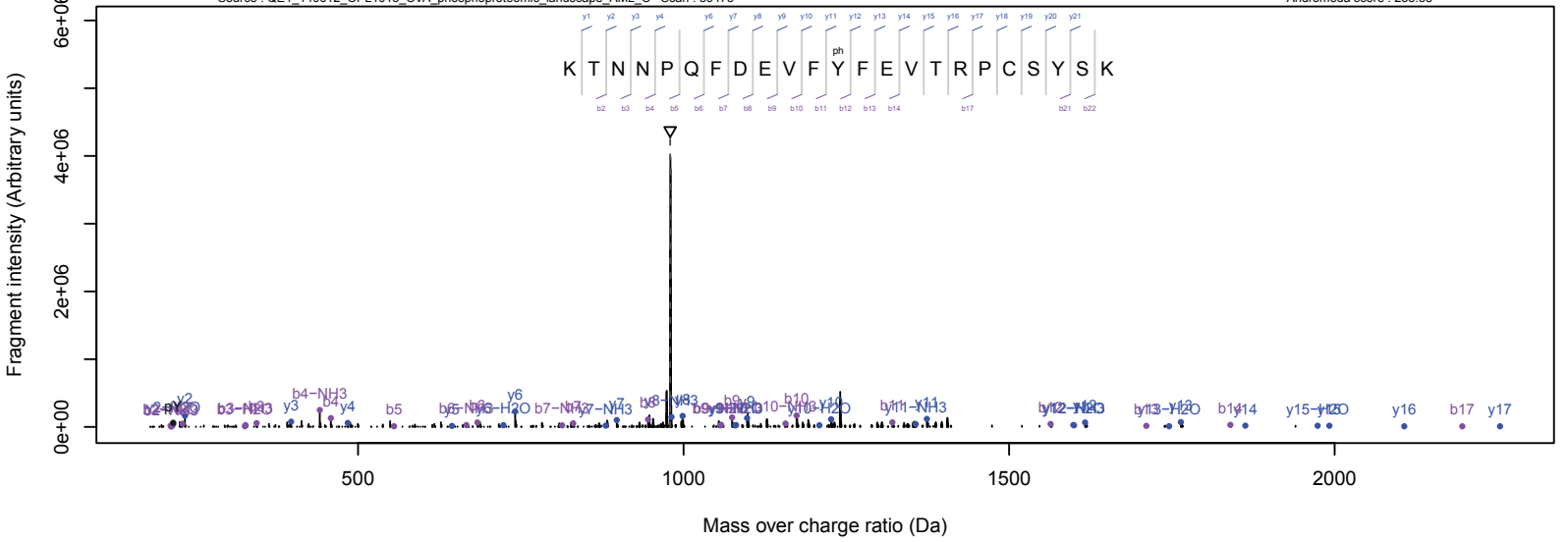
Andromeda score : 289.75



KTNNPQFDEVFY(ph)FEVTRPCSYSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39475

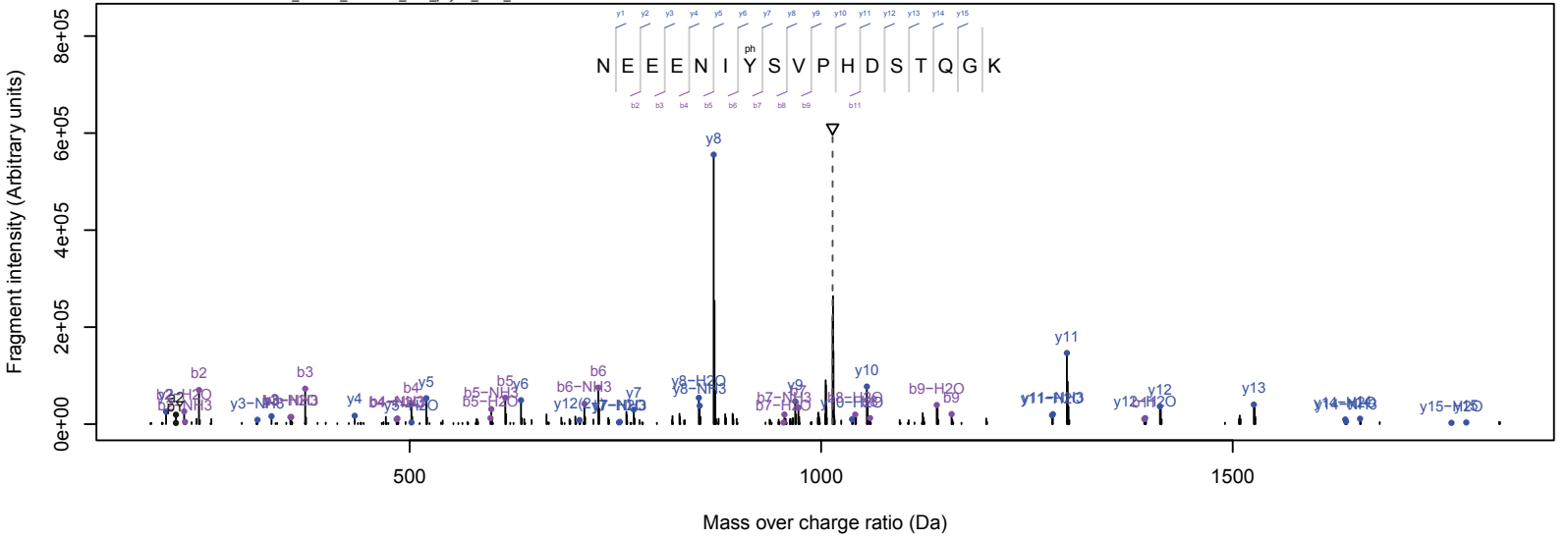
Andromeda score : 288.58



NEEENIY(ph)SVPHDSTQGK-2+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan: 12978

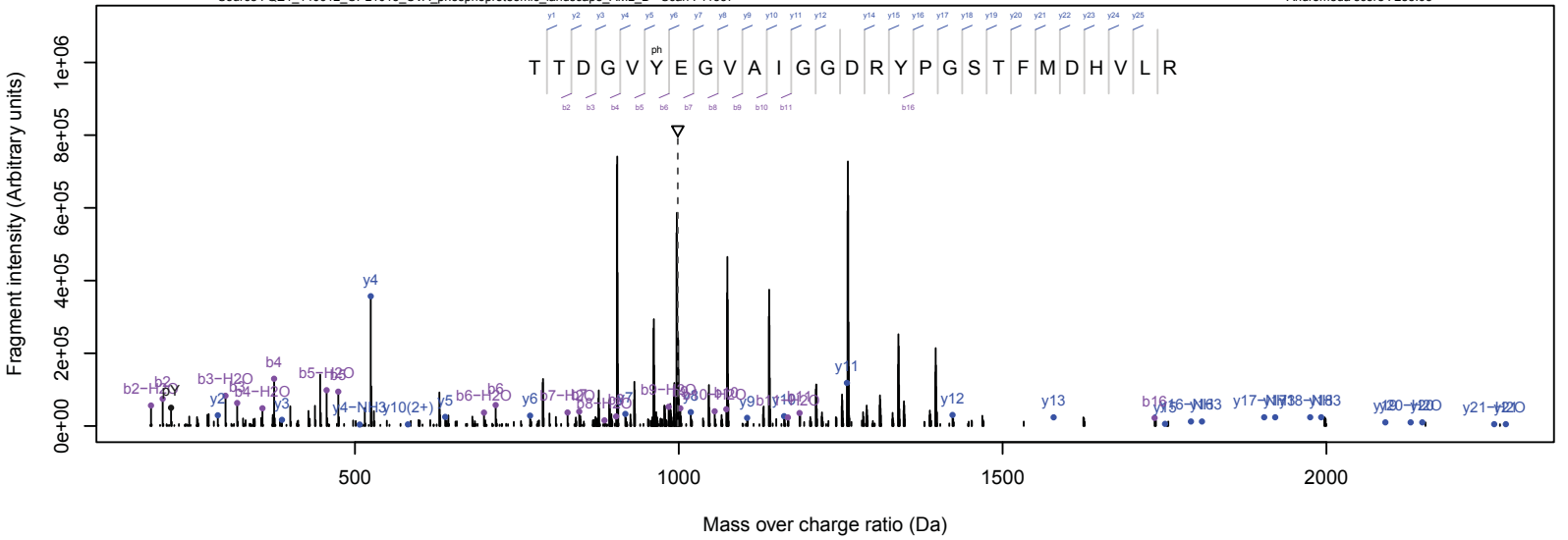
Andromeda score: 286.14



TTDGVY(ph)EGVAIGDRYPGSTFMDHVL R-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 41337

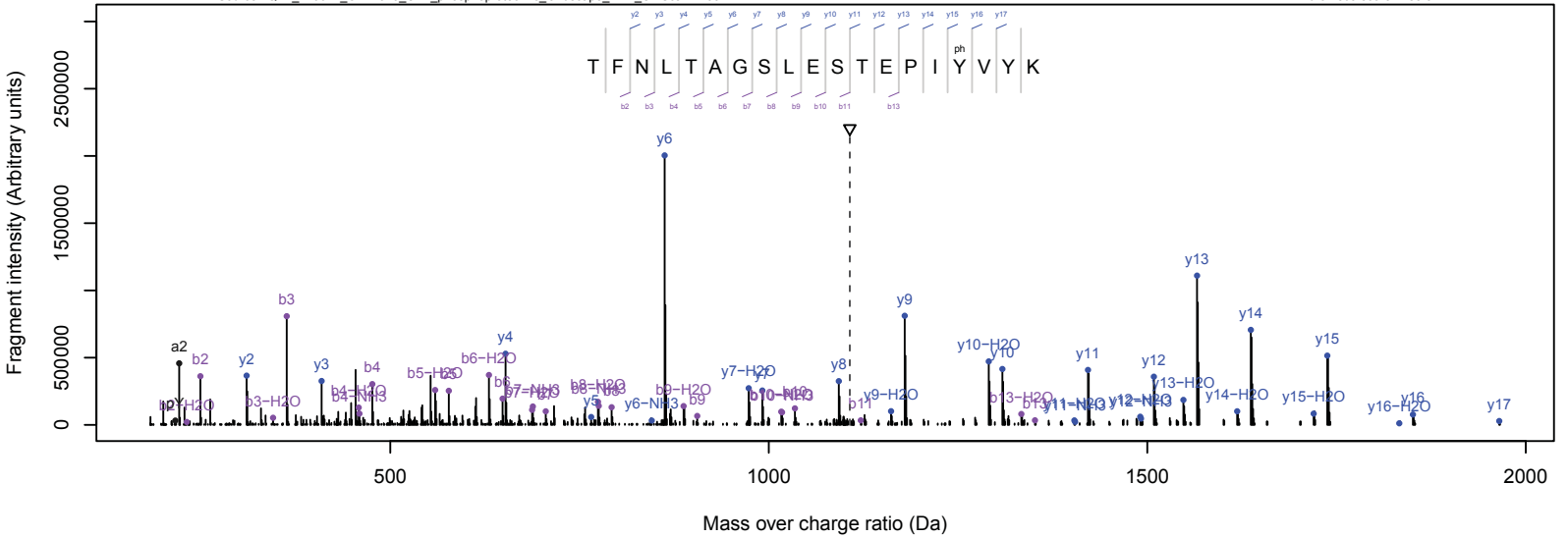
Andromeda score: 285.95



TFNLTAGSLESTEPIYVYK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 43522

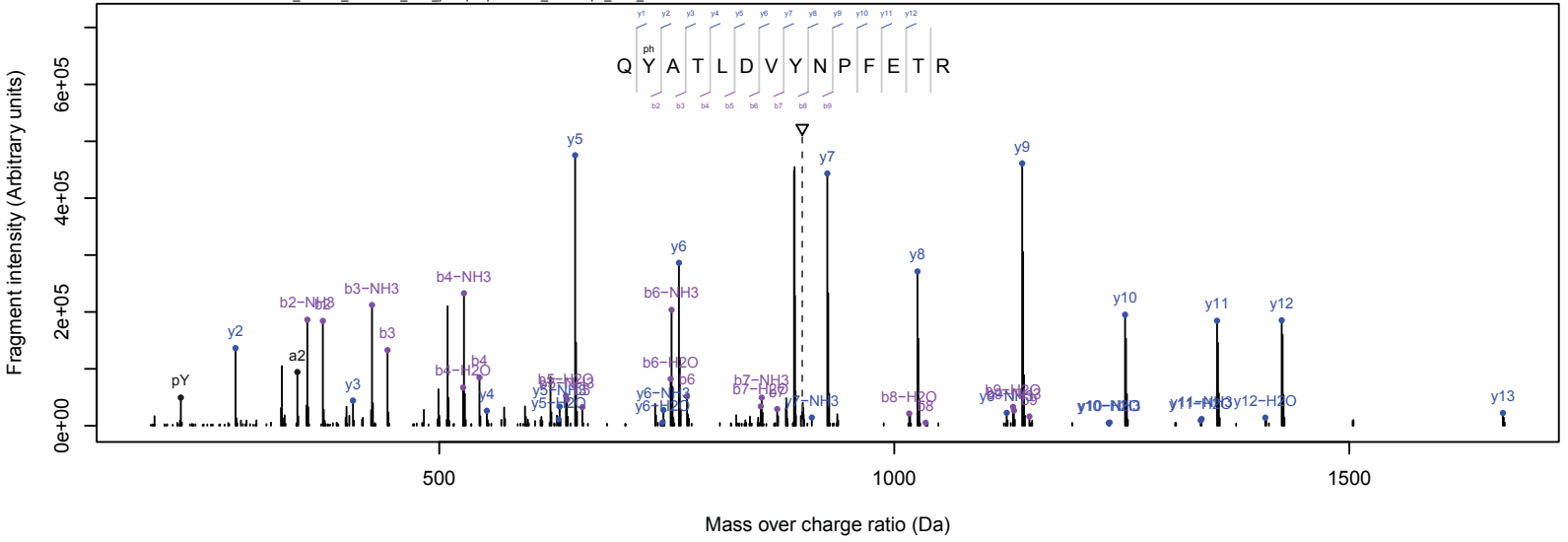
Andromeda score: 285.5



QY(ph)ATLDVYNPFETR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42434

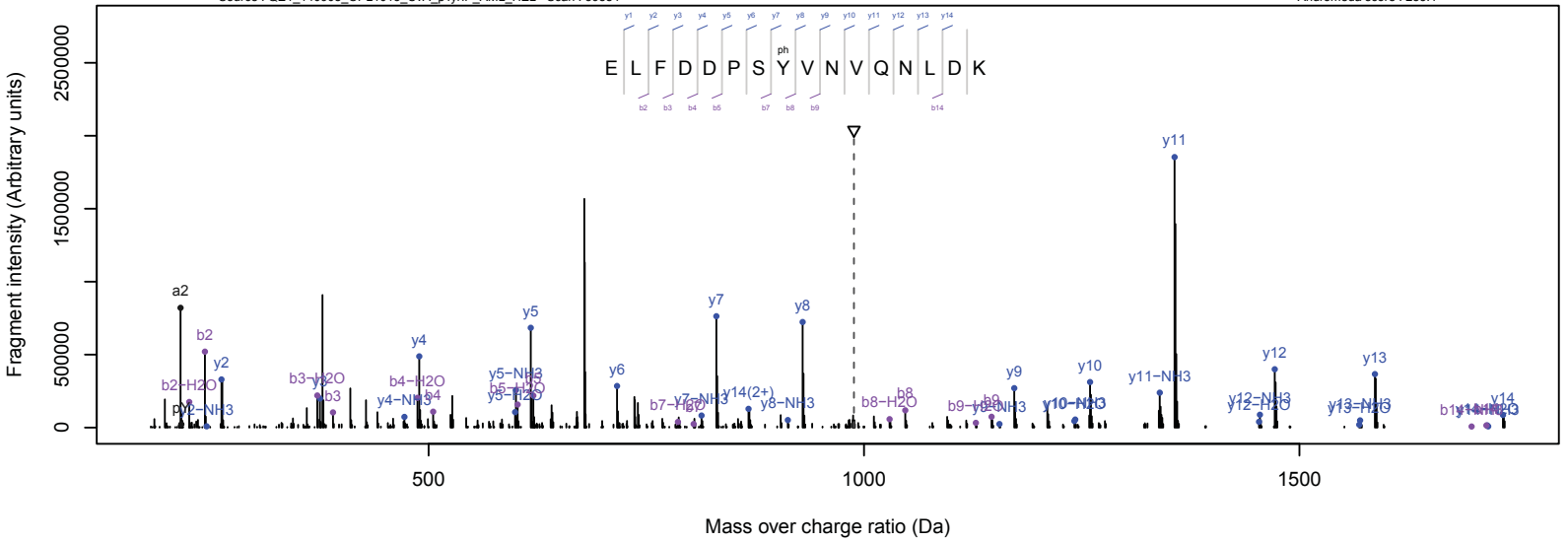
Andromeda score : 283.45



ELFDDPSY(ph)VNVQLDK-2+

Source : QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_HEL Scan : 39884

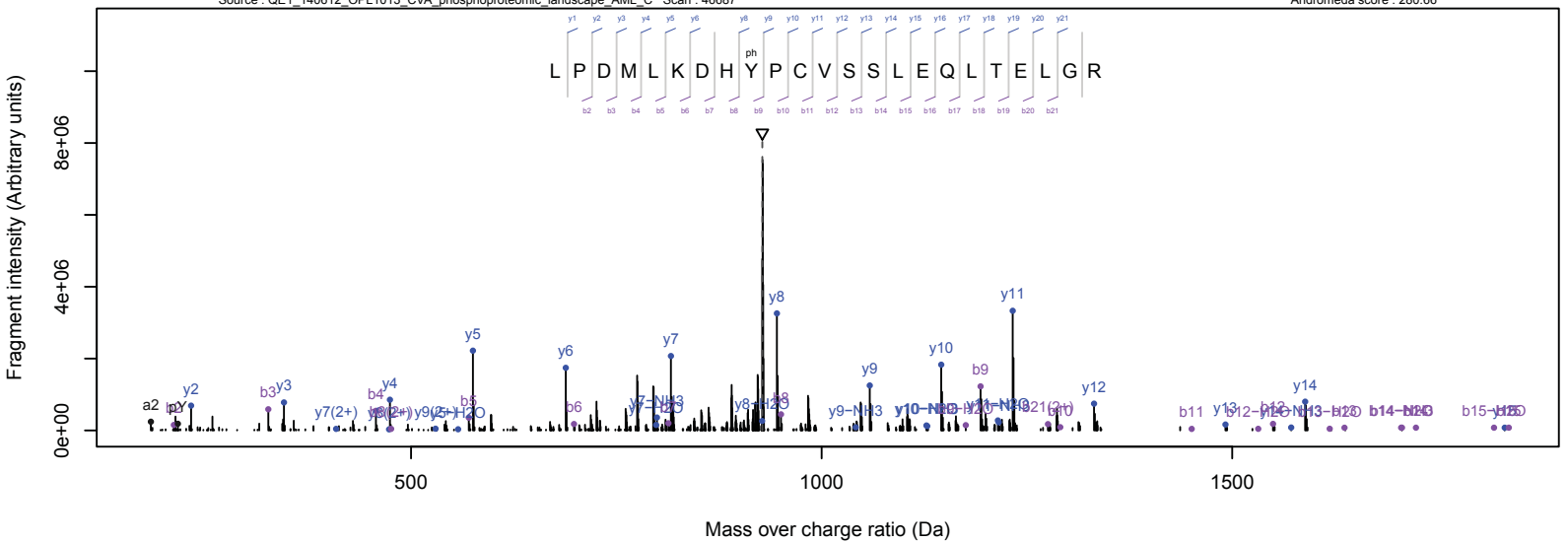
Andromeda score : 283.1



LPDMLKDHY(ph)PCVSSLEQLTELGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46687

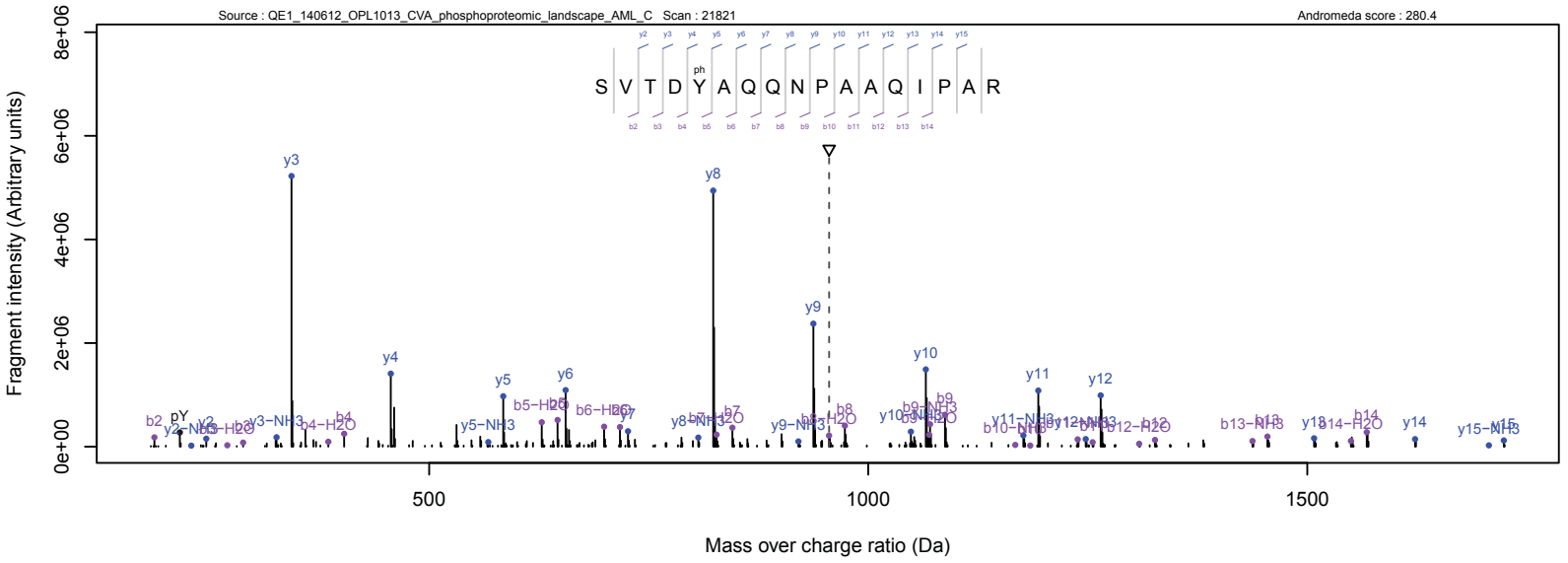
Andromeda score : 280.66



SVTDY(ph)AQQNPAQIPAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21821

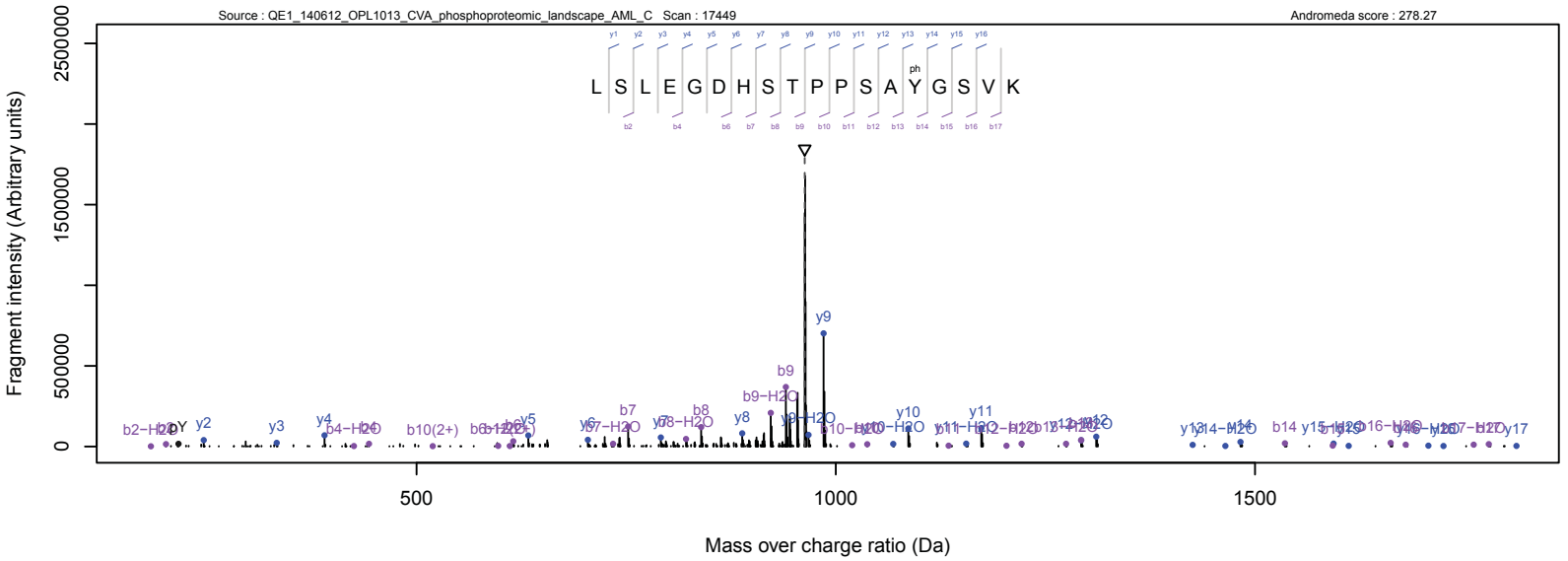
Andromeda score : 280.4



LSLEGDHSTPPSAY(ph)GSVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17449

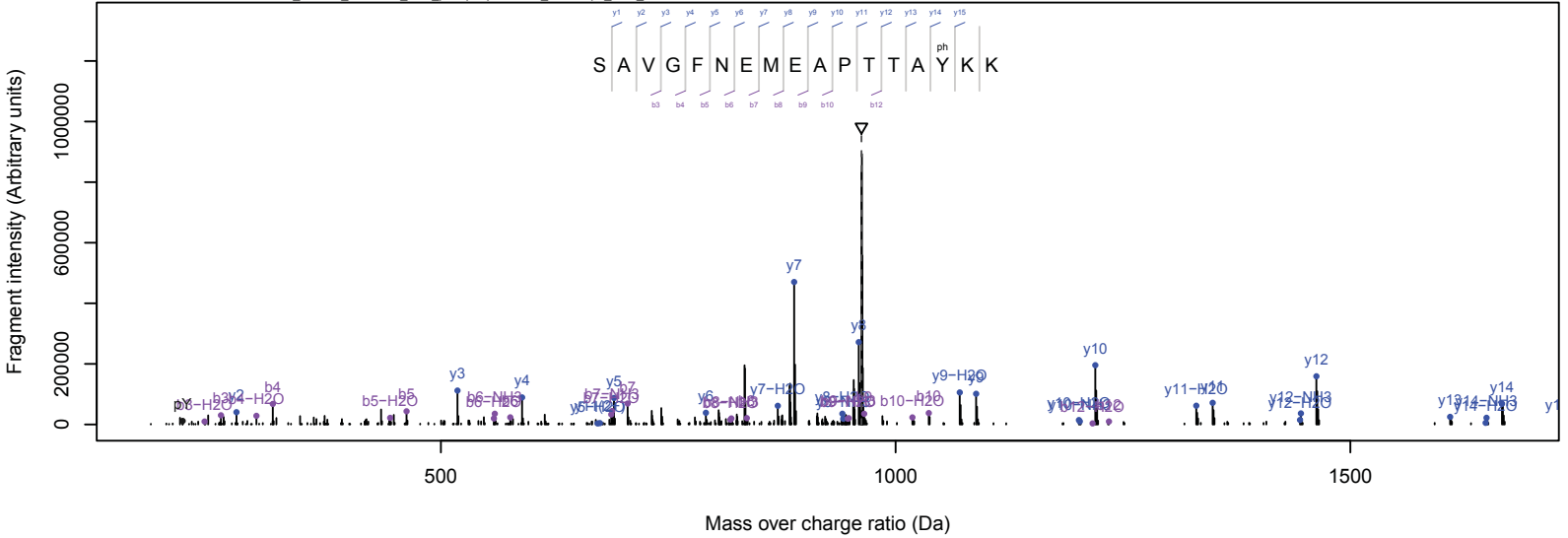
Andromeda score : 278.27



SAVGFNEMEAPTAY(ph)KK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20989

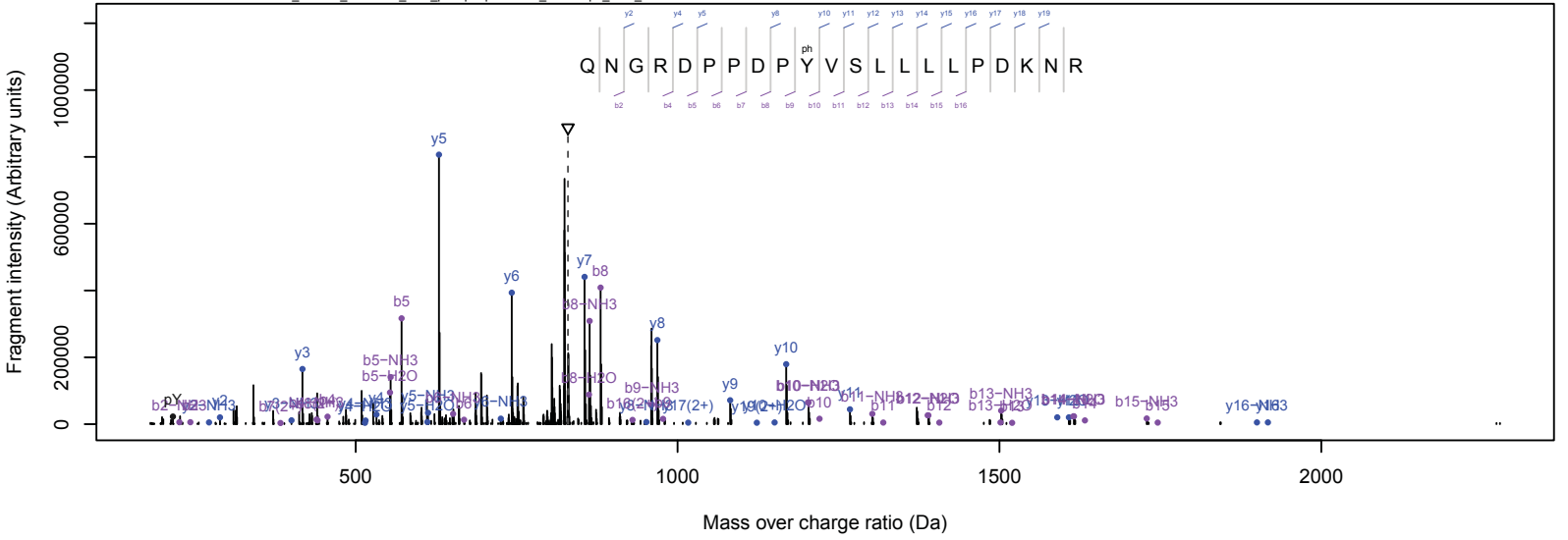
Andromeda score : 277.34



QNGRDPPY(ph)VLLLLPKNR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 36446

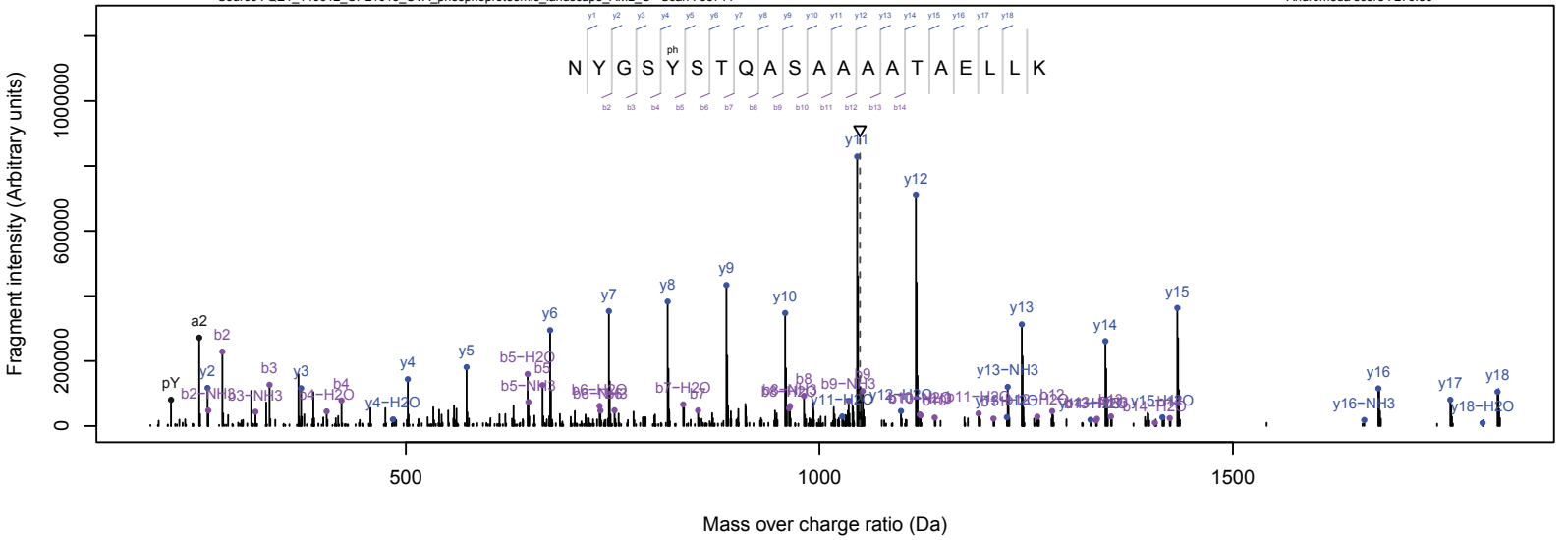
Andromeda score: 277.03



NYGSY(ph)STQASAAAATAELLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 35711

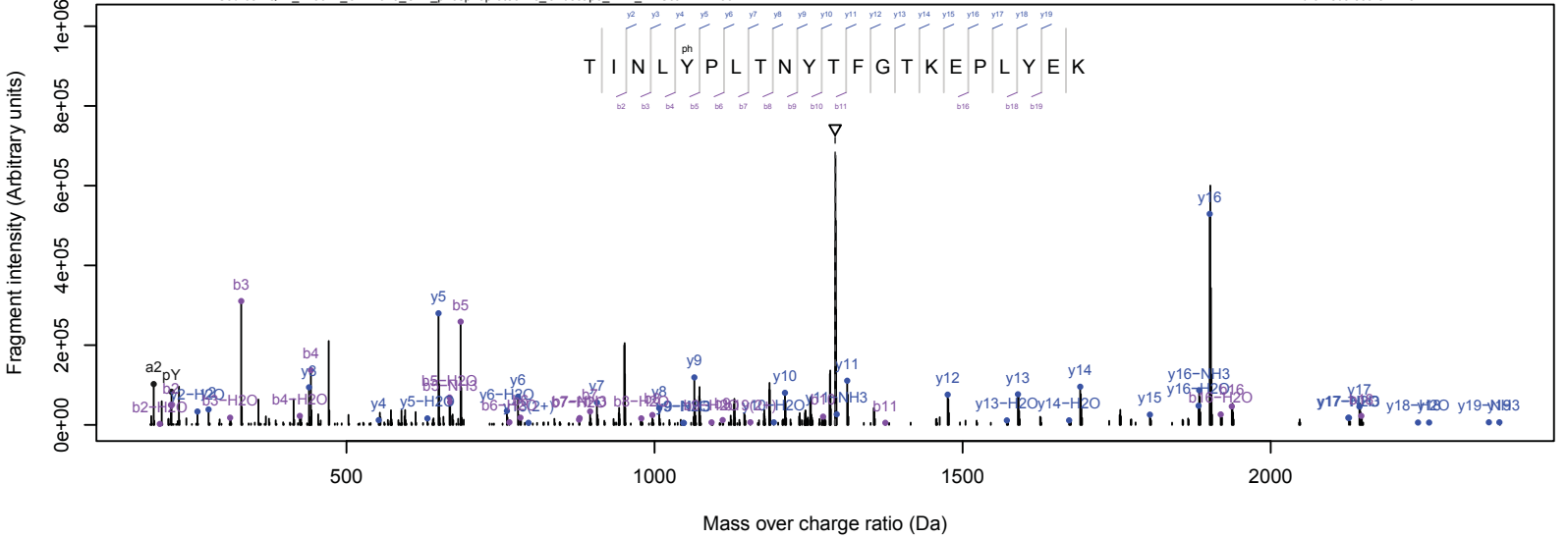
Andromeda score: 276.88



TINLY(ph)PLTNYTFGKELYEK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 44706

Andromeda score: 275.41

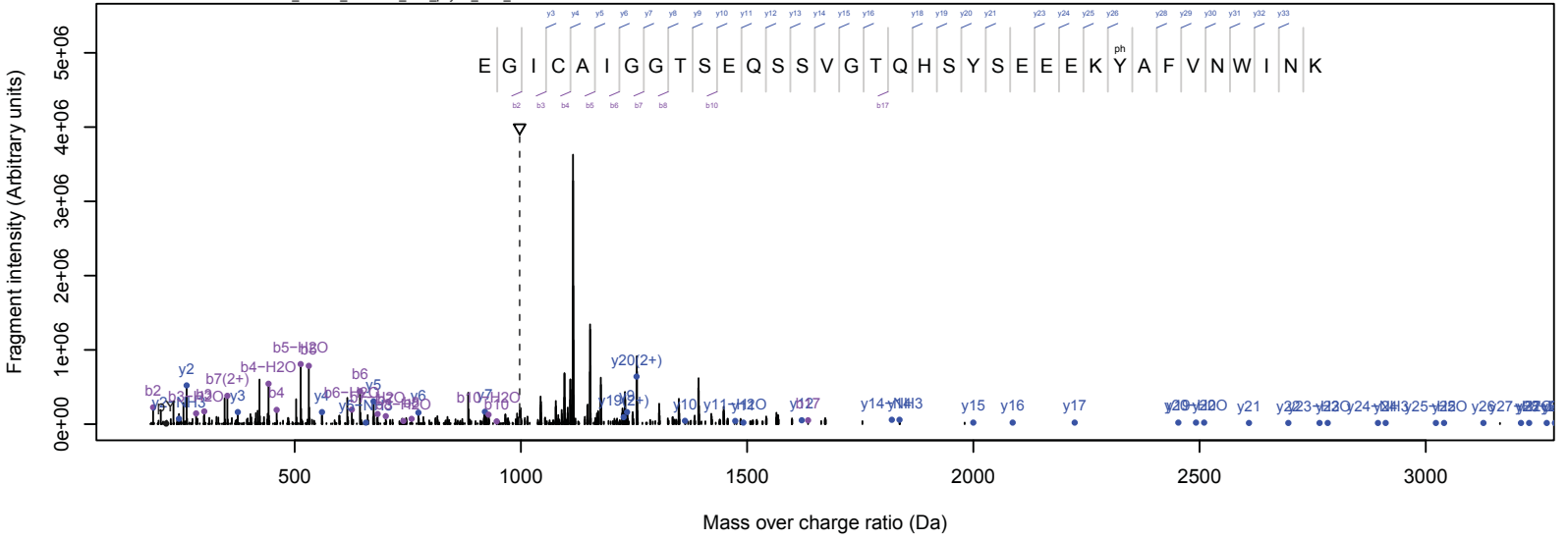




**EGICAIGGTSEQSSVGTQHSYSEEEKY(ph)AFVNWINK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 43737

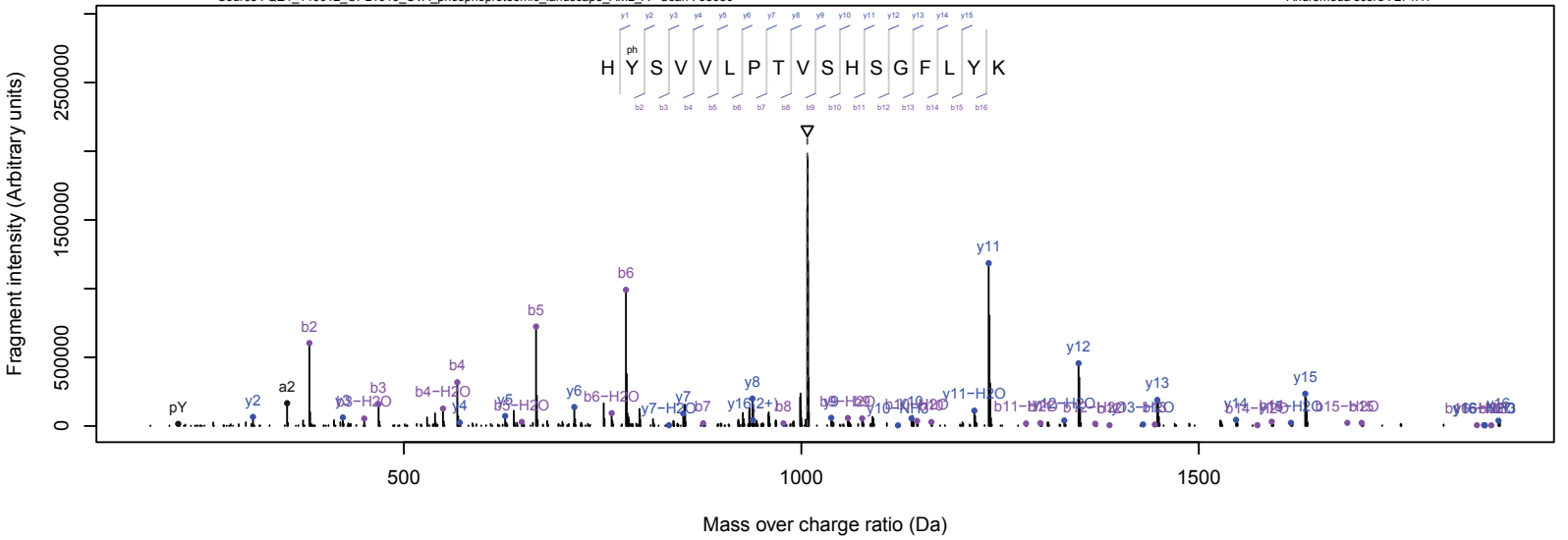
Andromeda score : 274.96



**HY(ph)SVLPTVSHSGFLYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33080

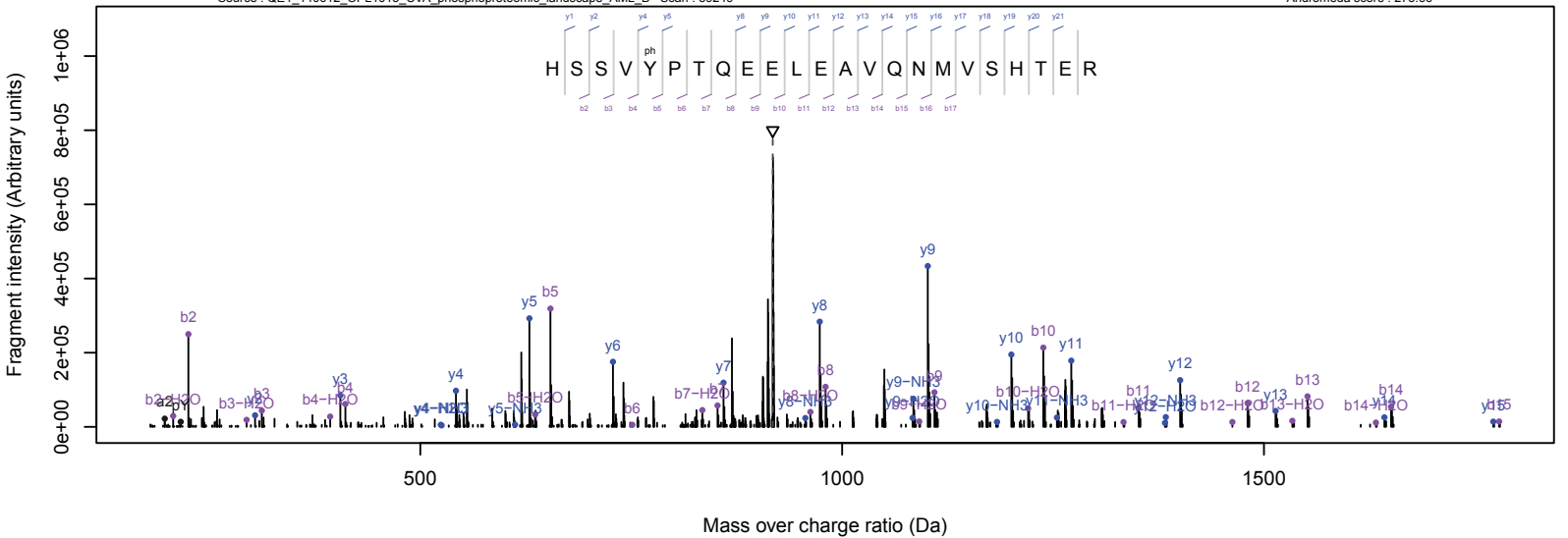
Andromeda score : 274.17



**HSSVY(ph)PTQEELEAVQNMVSHTER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39249

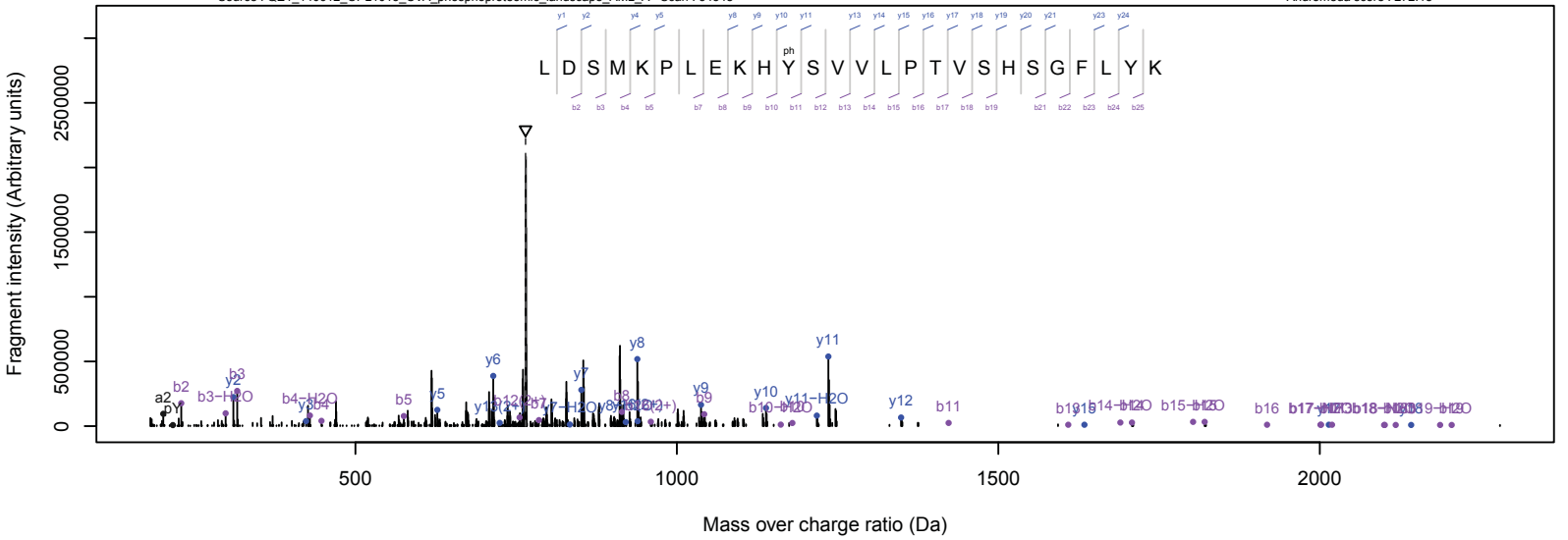
Andromeda score : 273.66



**LDSMKPLEKHY(ph)SVVLP TVSHSGFLYK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34548

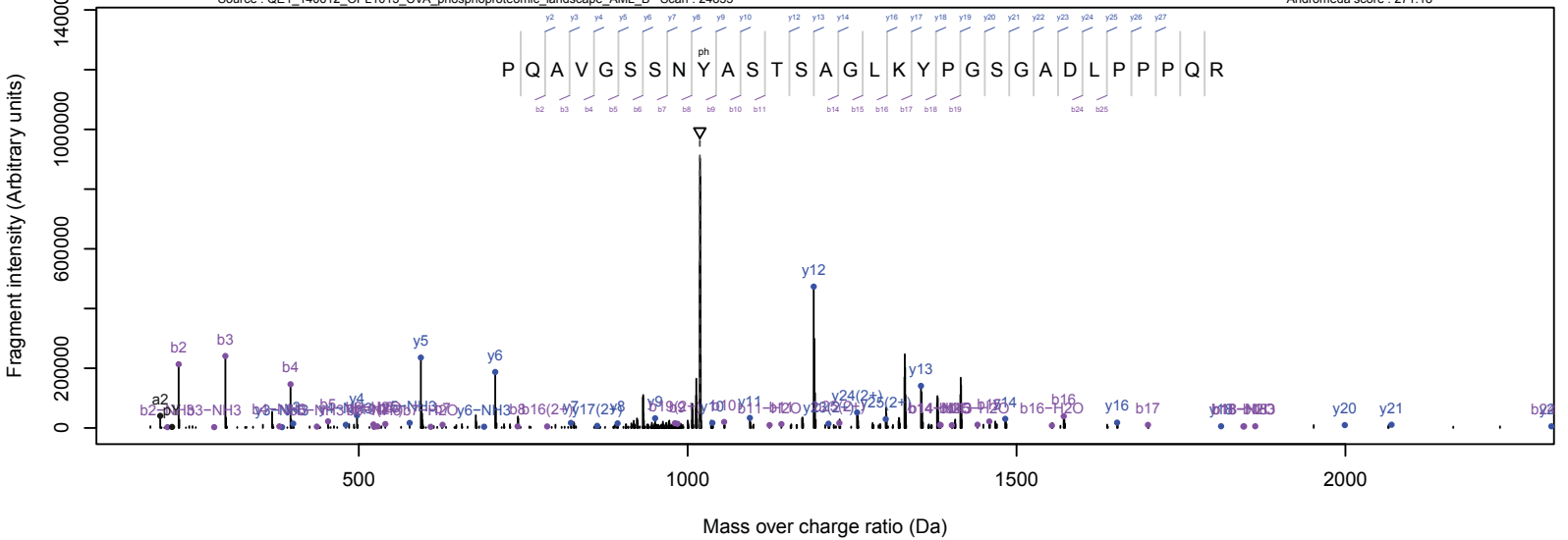
Andromeda score : 272.18



**PQAVGSSNY(ph)ASTSAGLKYPGSGADLPPQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24855

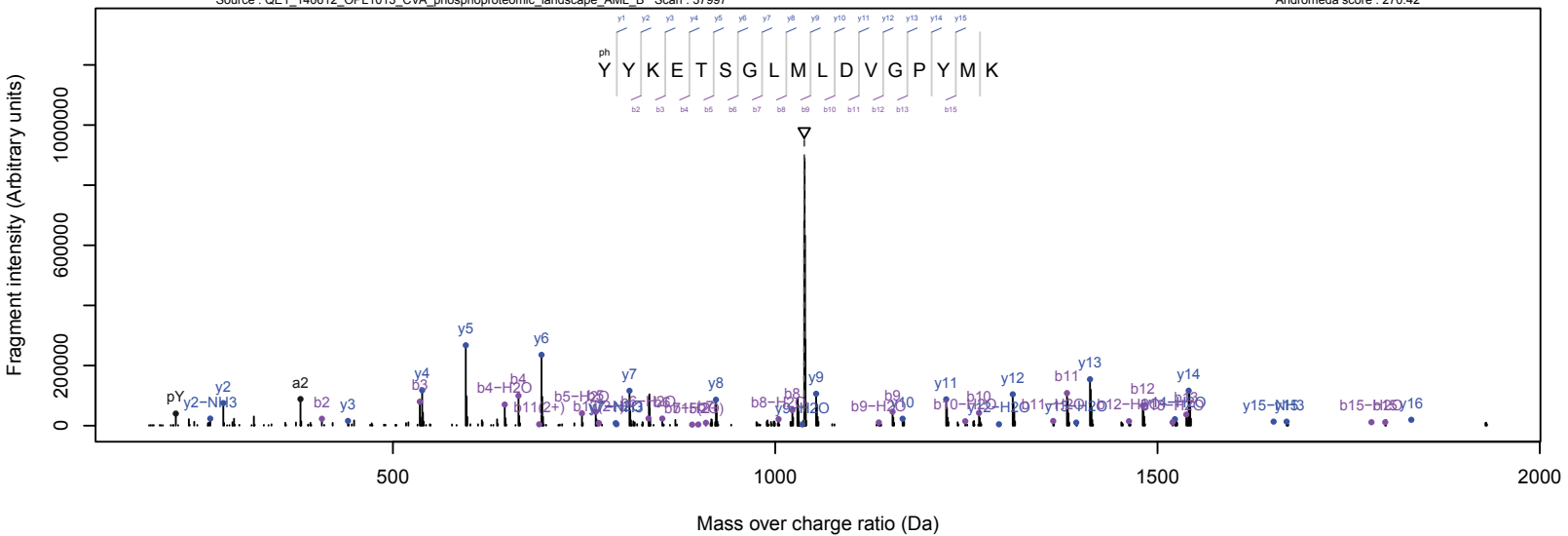
Andromeda score : 271.18



**Y(ph)YKETSGMLLDVGPYMK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37997

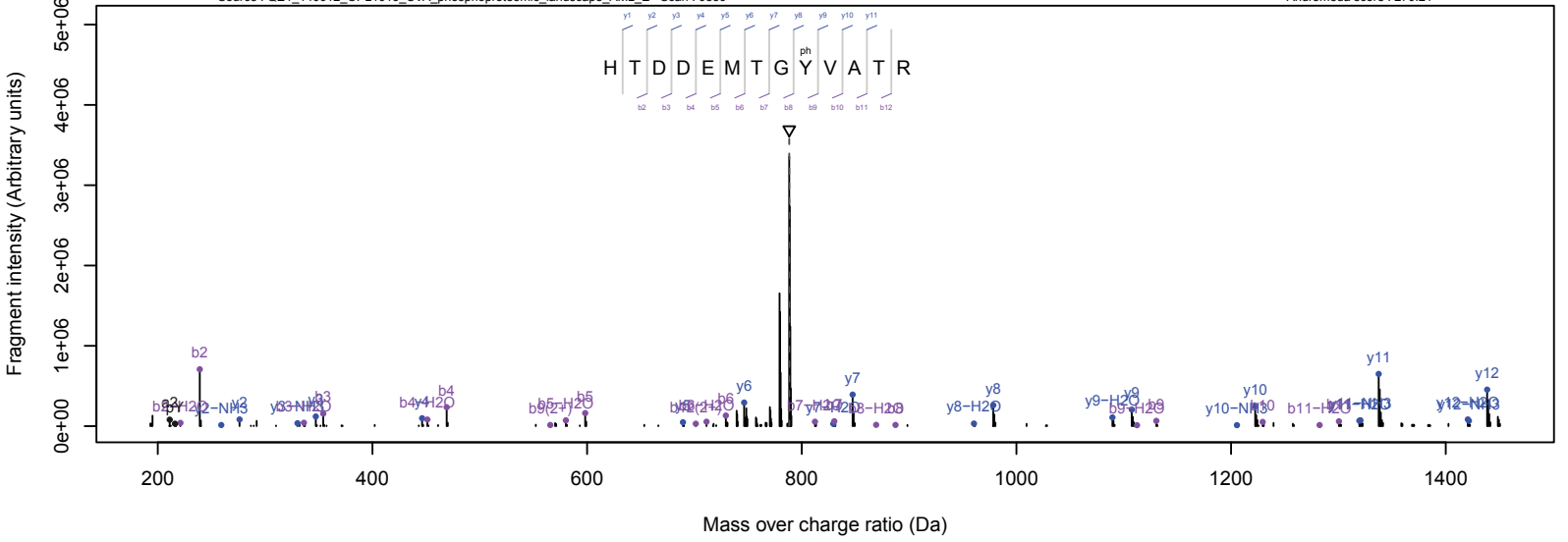
Andromeda score : 270.42



HTDDEMTGY(ph)VATR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 9859

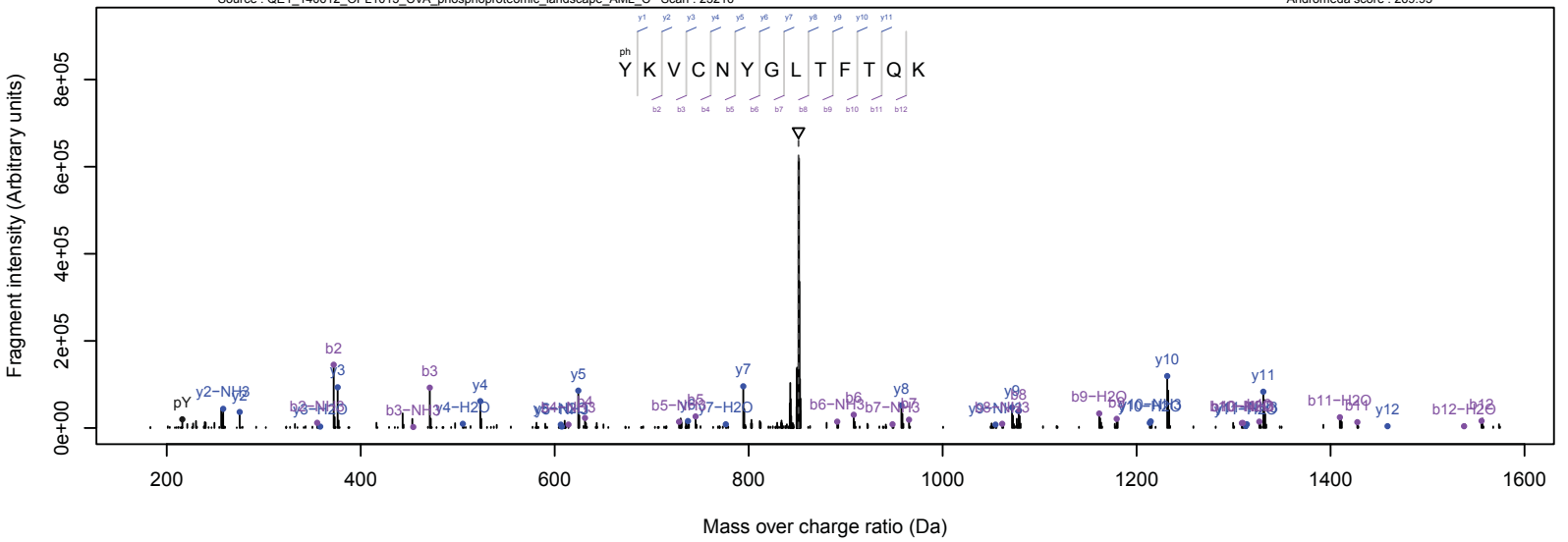
Andromeda score : 270.21



Y(ph)KVCNYGLTFTQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23216

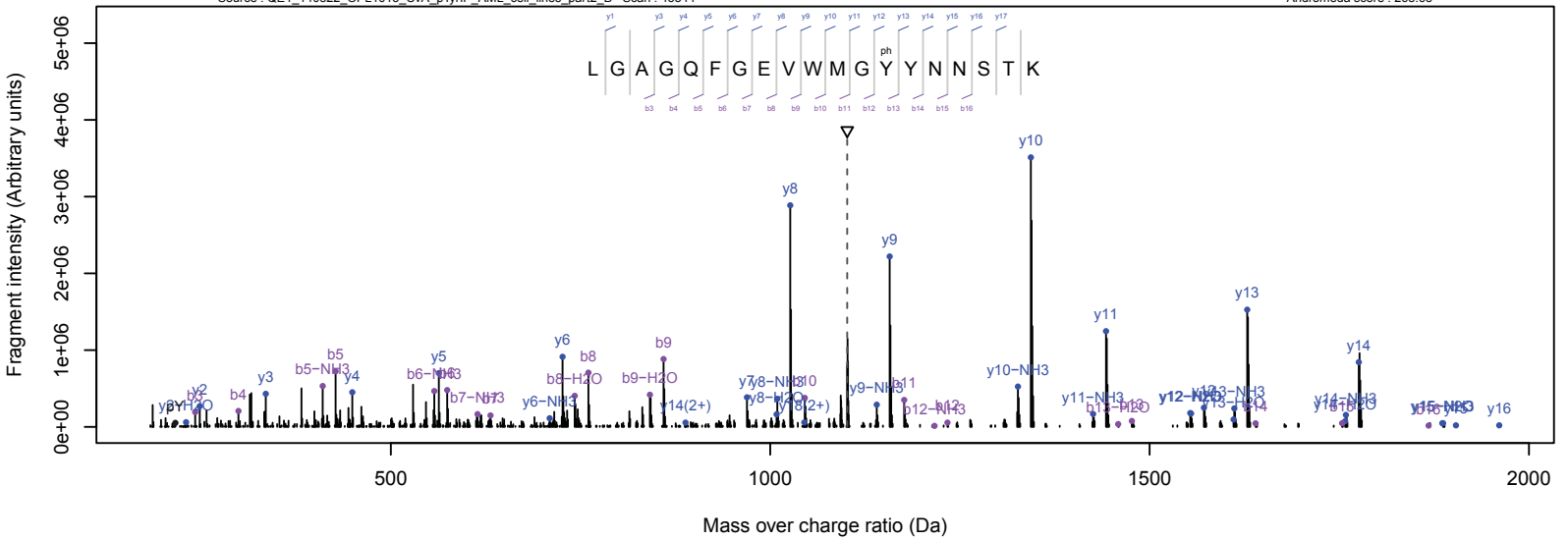
Andromeda score : 269.95



LGAGQFGEVWMGYYNNSTK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 45344

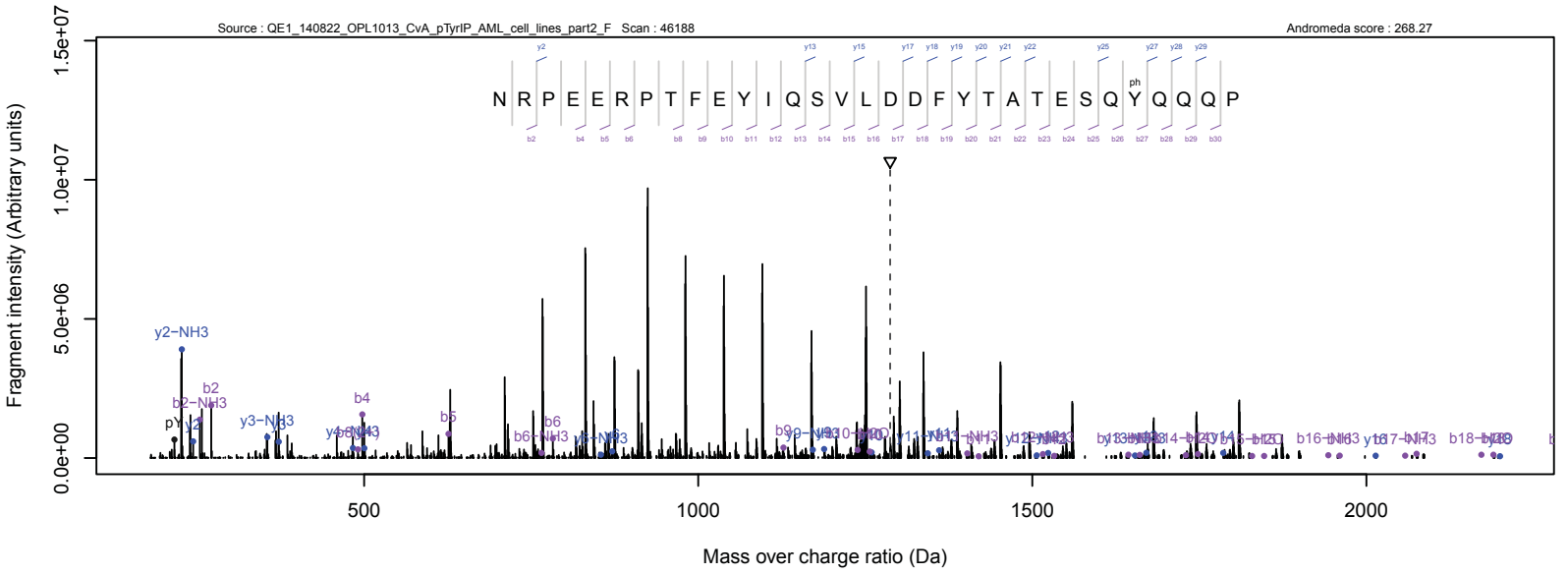
Andromeda score : 268.65



NRPEERPTFEYIQSVLDDDFYTATESQY(ph)QQQP-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 46188

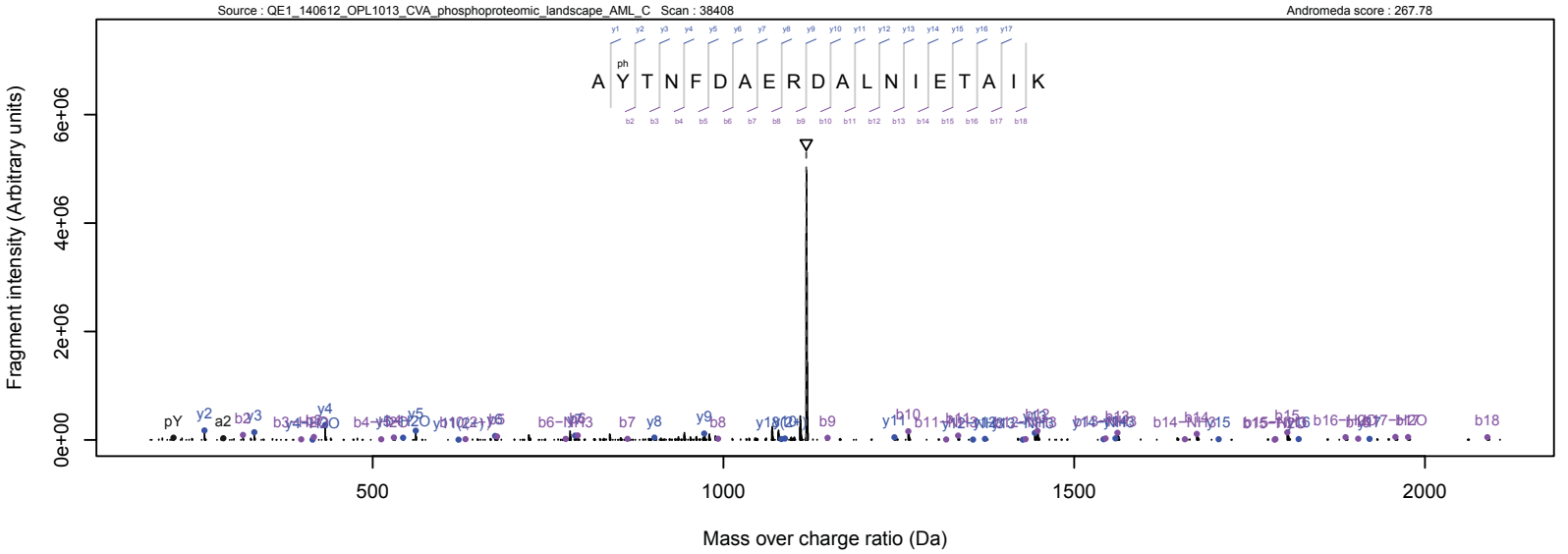
Andromeda score : 268.27



AY(ph)TNFDAERDALNIETAIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38408

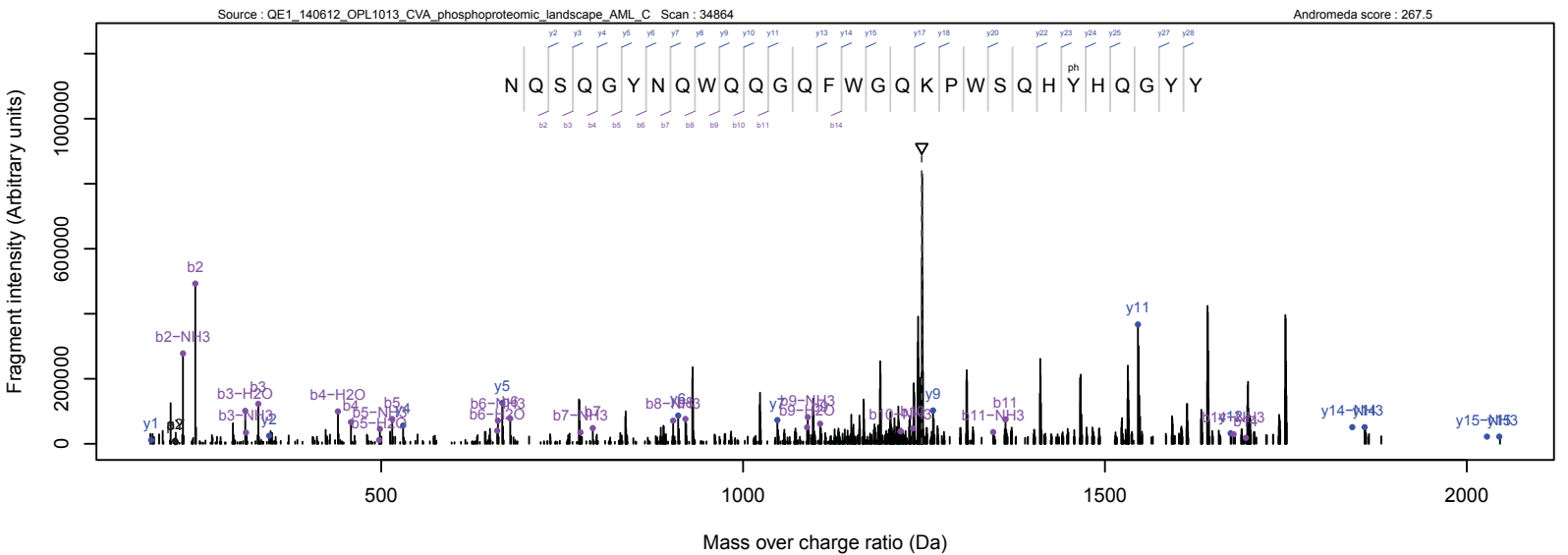
Andromeda score : 267.78



NQSQGYNQWQQGQFWGQK PWSQH Y(ph)HQGY-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34864

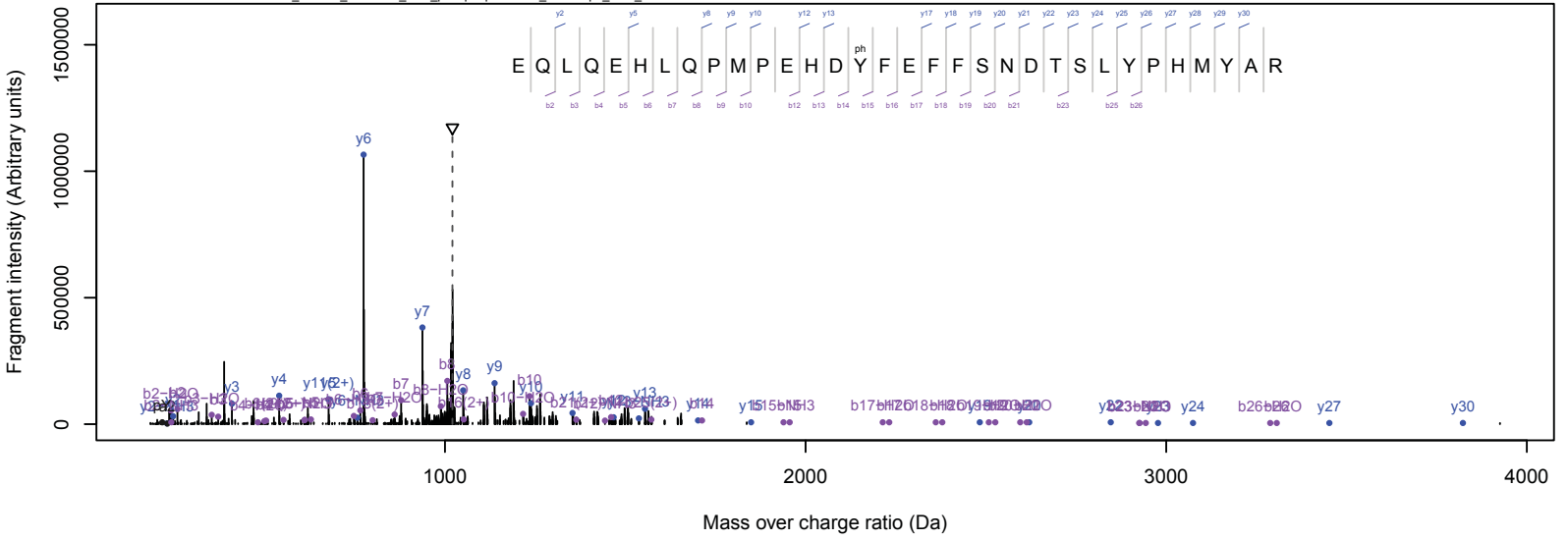
Andromeda score : 267.5



**EQLQEHLQPMPEHDY(ph)FEFFSNDTSLYPHMYAR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45852

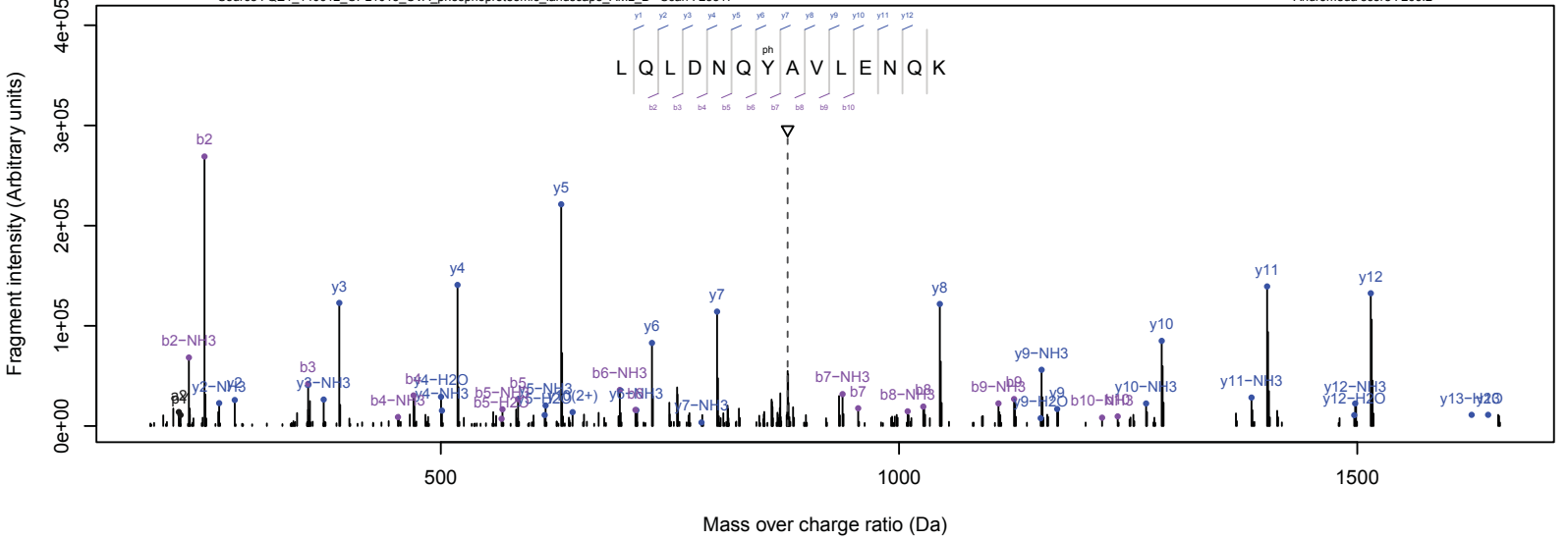
Andromeda score : 266.57



**LQLDNQY(ph)AVLENQK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25317

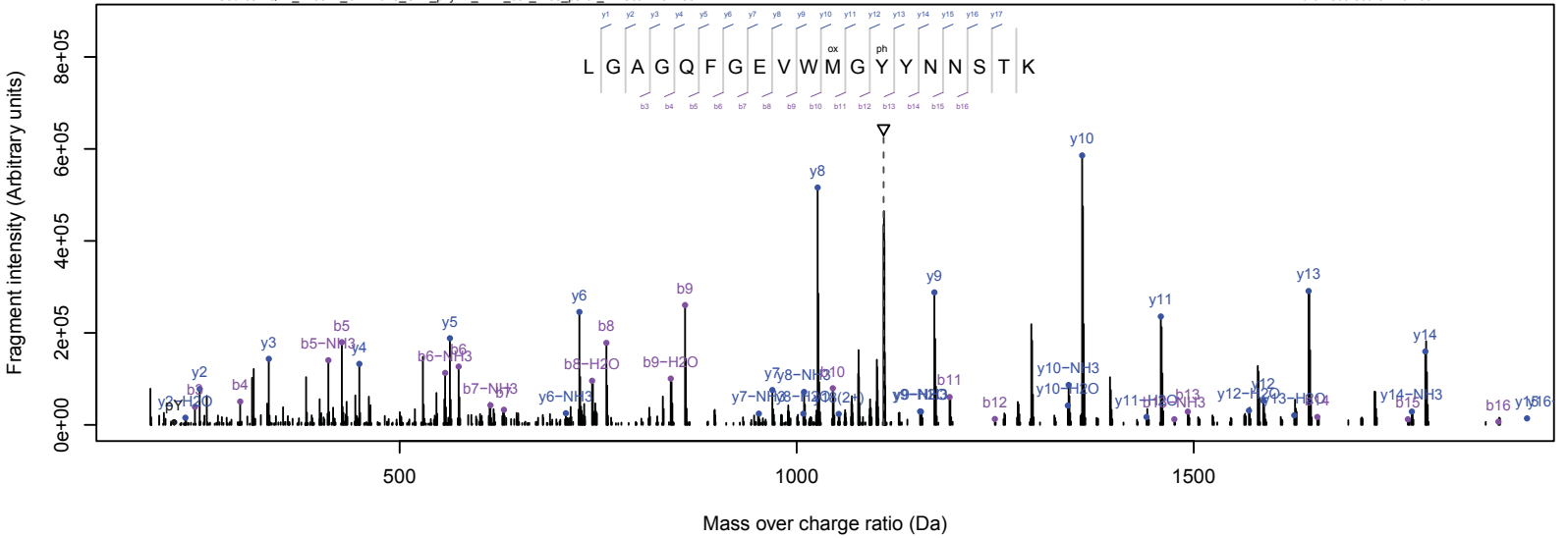
Andromeda score : 266.2



**LGAGQFGEVWMG(ph)YNNSTK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 32153

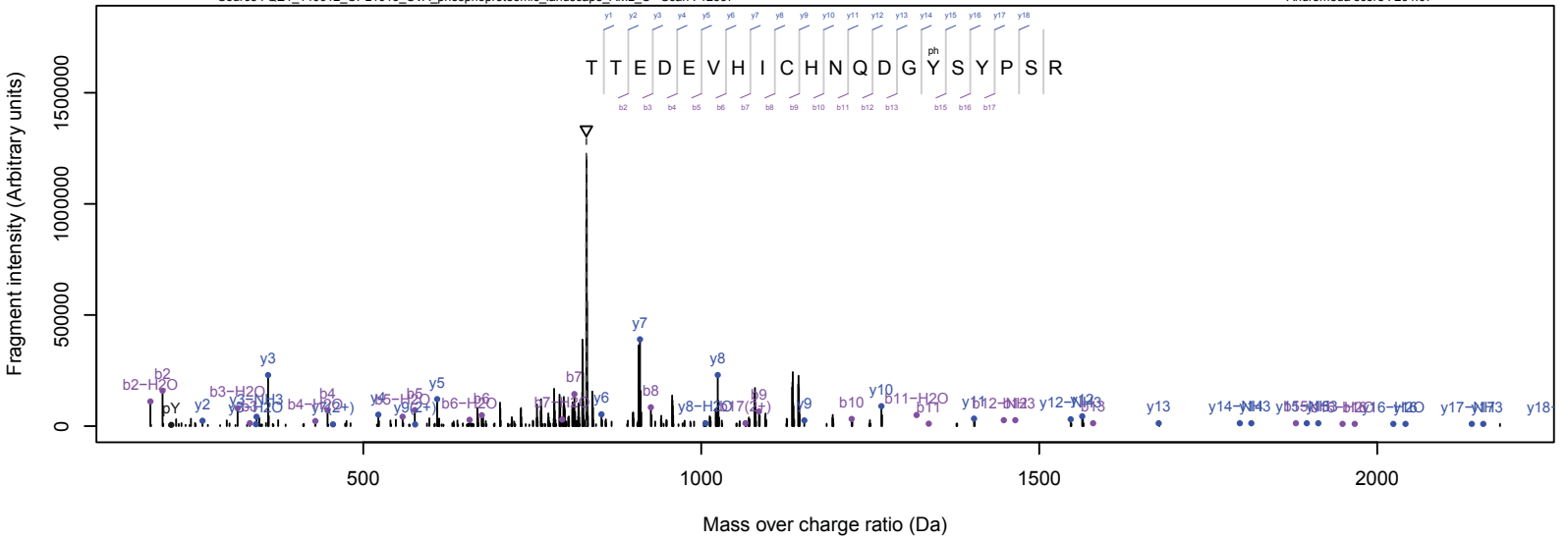
Andromeda score : 264.69



TTEDEVHICHNQDGY(ph)SYPSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12537

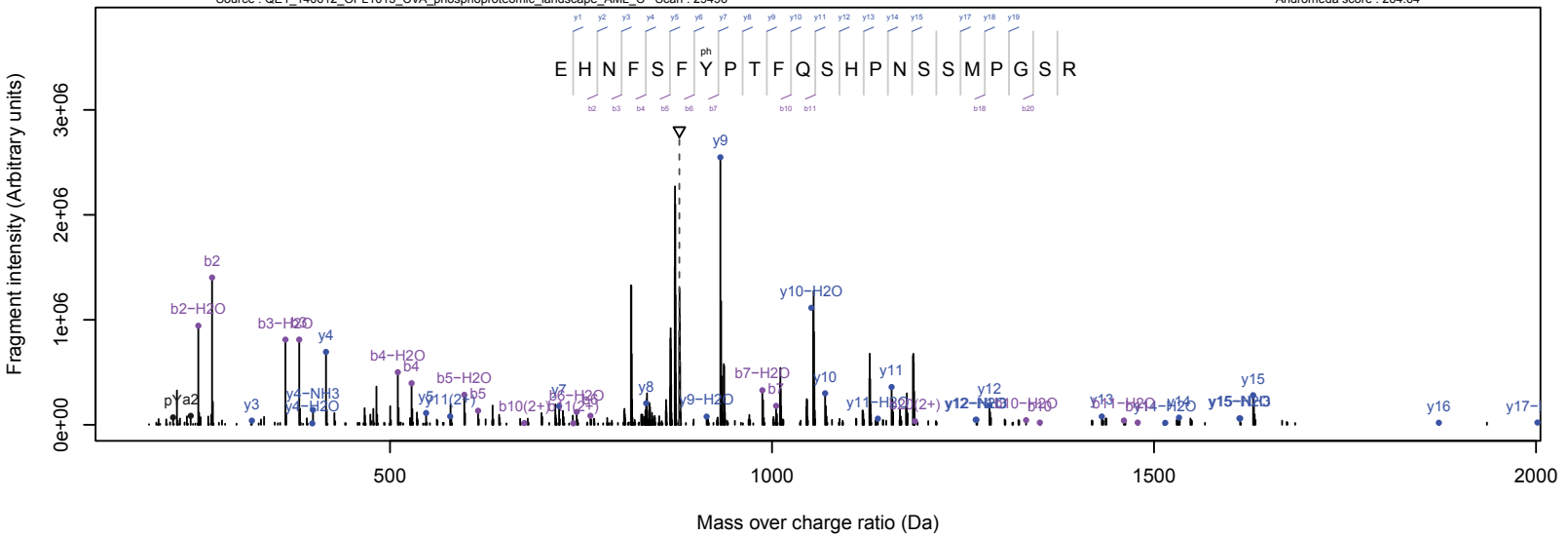
Andromeda score : 264.67



EHNFSFY(ph)PTFQSHPNSSMPGSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29496

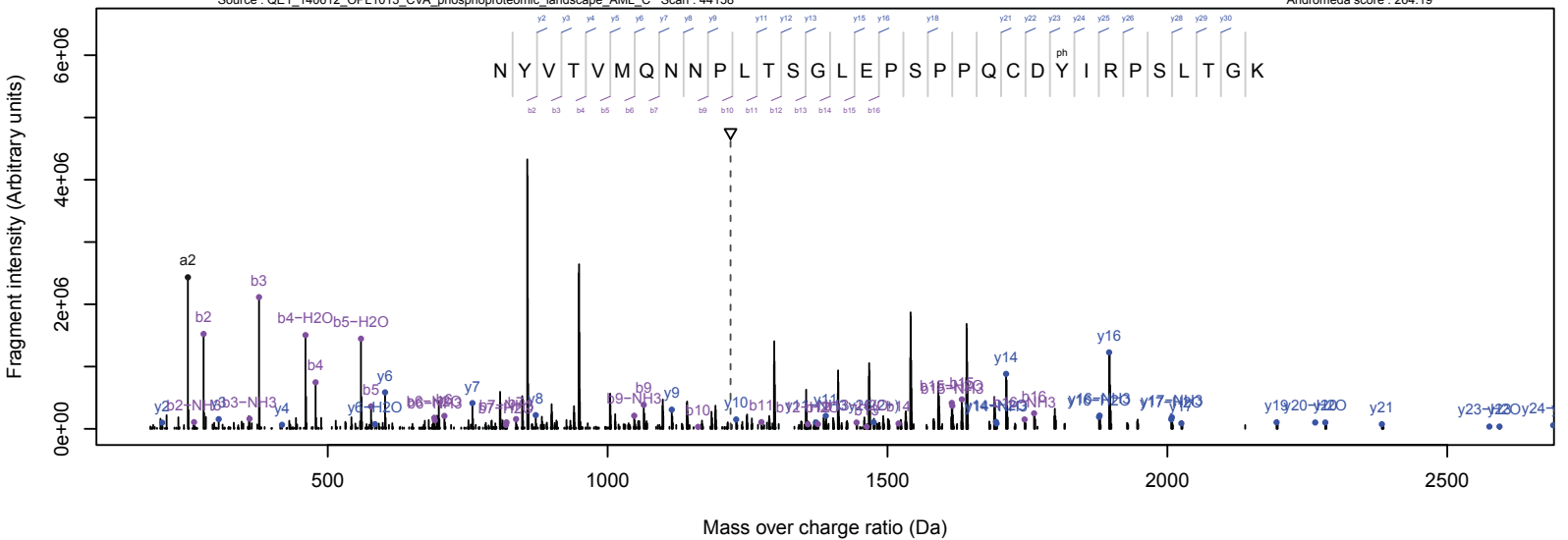
Andromeda score : 264.64



NYVTVMQNNPLTSLGLEPSPQCDY(ph)IRPSLTGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44158

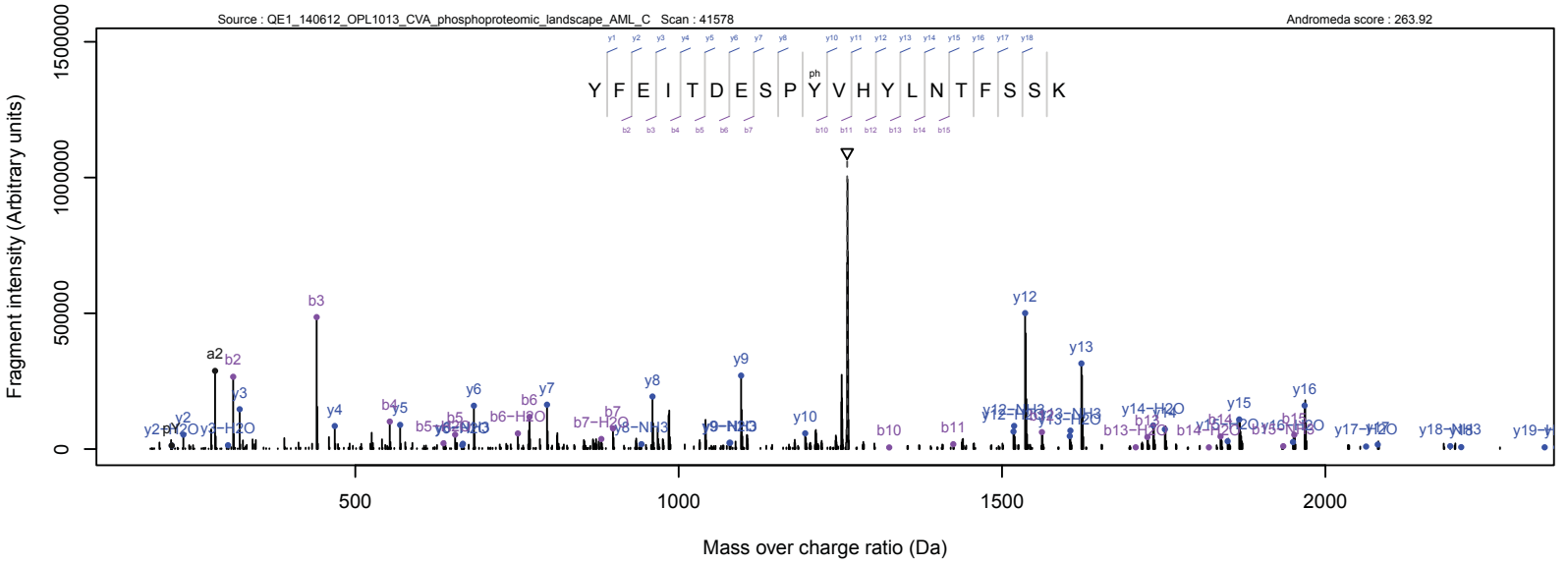
Andromeda score : 264.19



**YFEITDESPY(ph)VHYLNTFSSK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41578

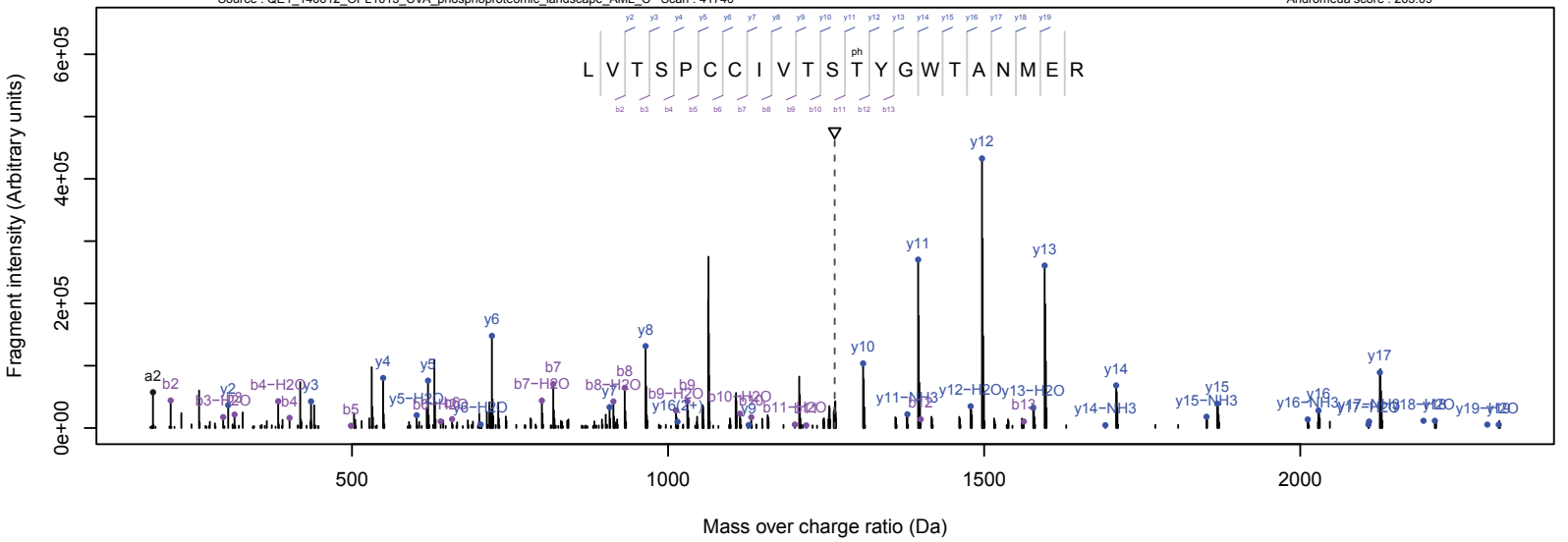
Andromeda score : 263.92



**LVTSPPCCIVTST(ph)YGWTANMER-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41740

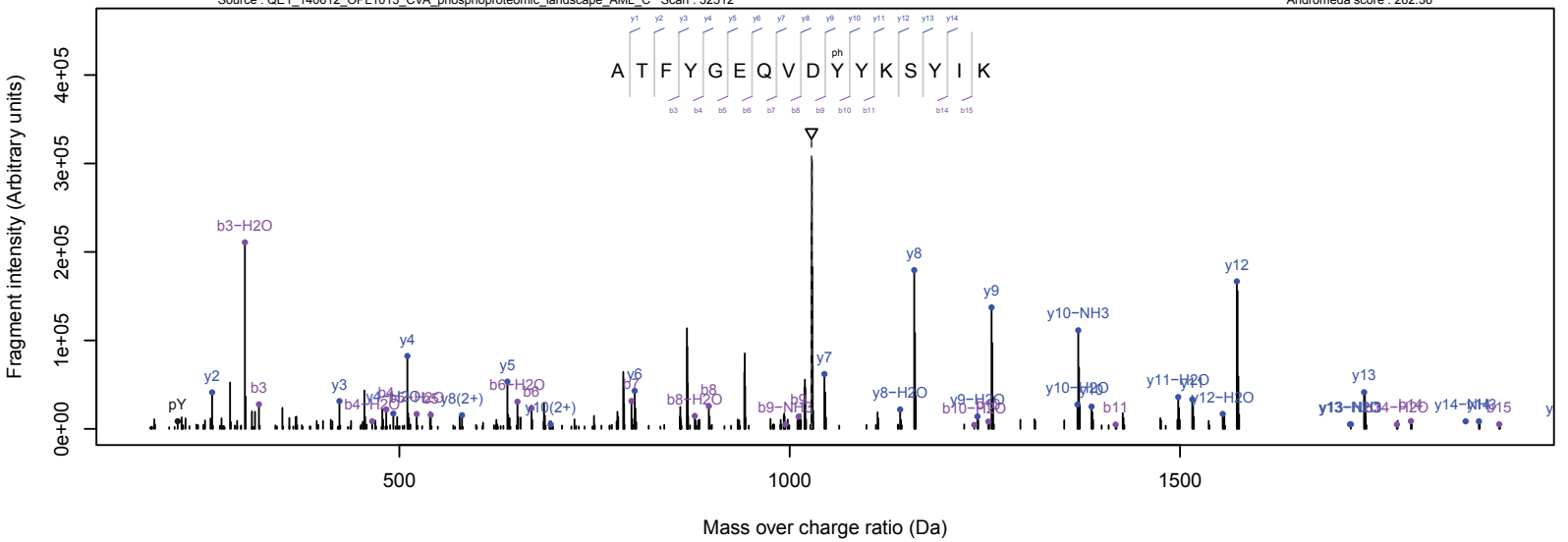
Andromeda score : 263.09



**ATFYGEQVDY(ph)YKSYIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32512

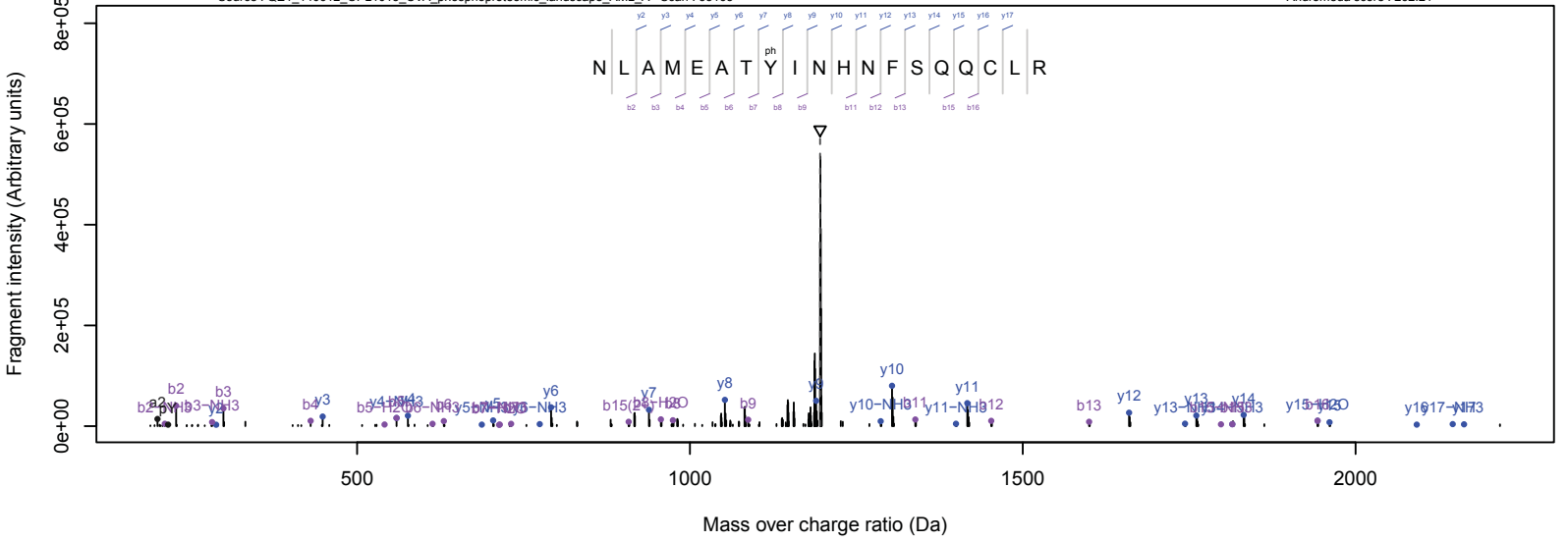
Andromeda score : 262.36



NLAMEATY(ph)INHNSQQCLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33163

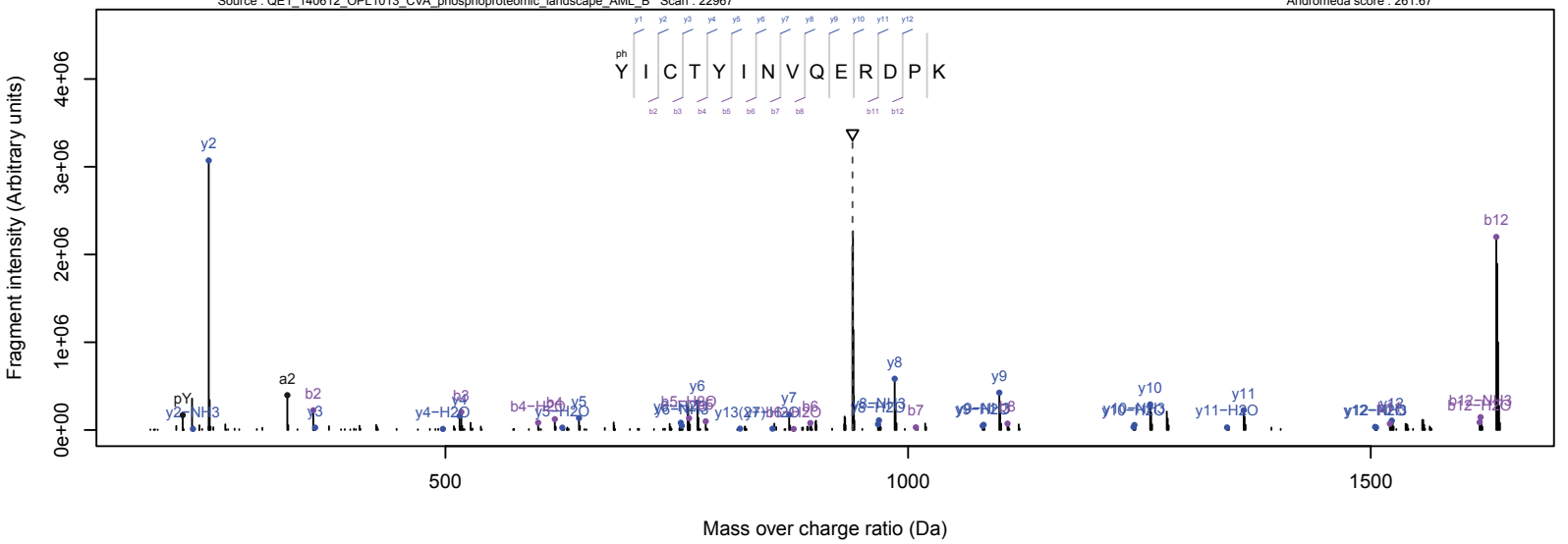
Andromeda score : 262.21



Y(ph)ICTYINVQERDPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22967

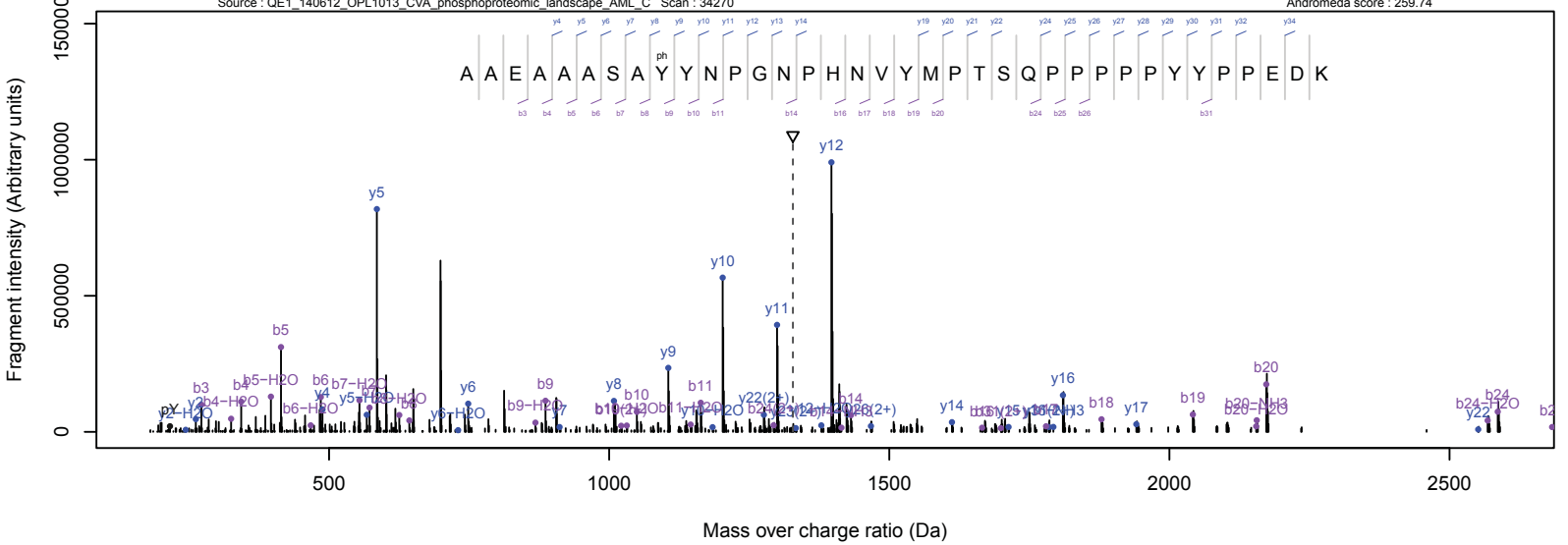
Andromeda score : 261.67



AAEAAASAY(ph)YNPGNPHNVYMPPTSQPPPPYPPEDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34270

Andromeda score : 259.74

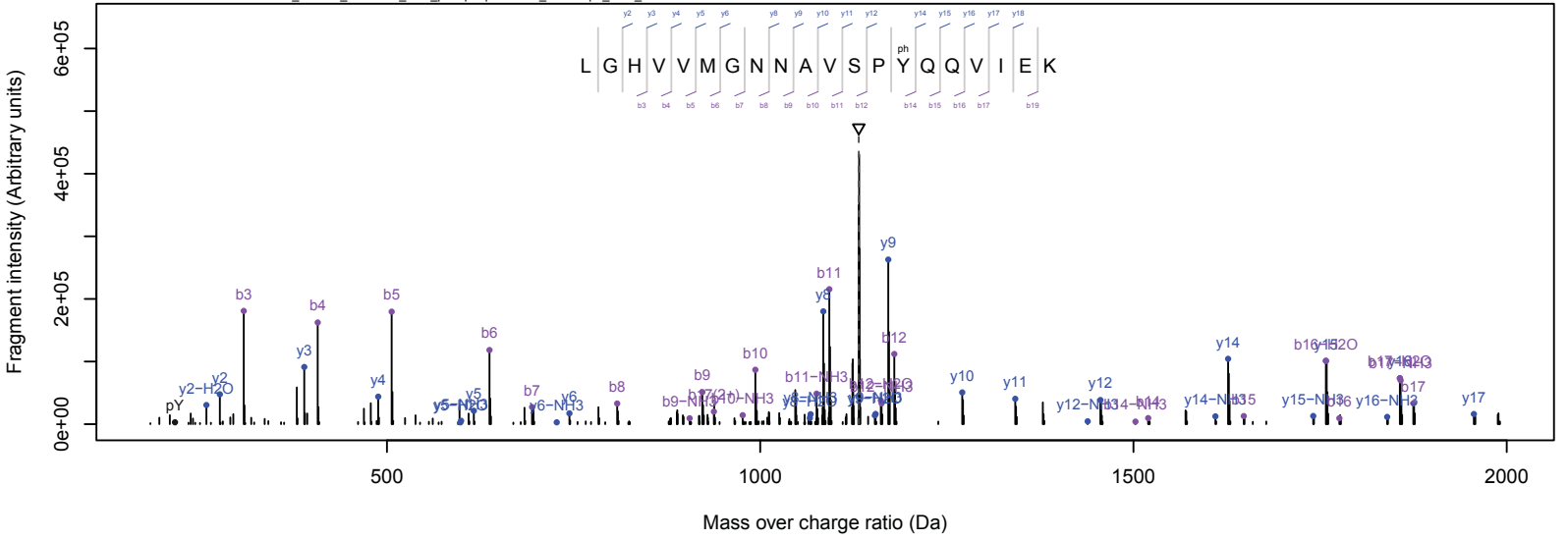




LGHVVMGNNAVSPYQQVIEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24882

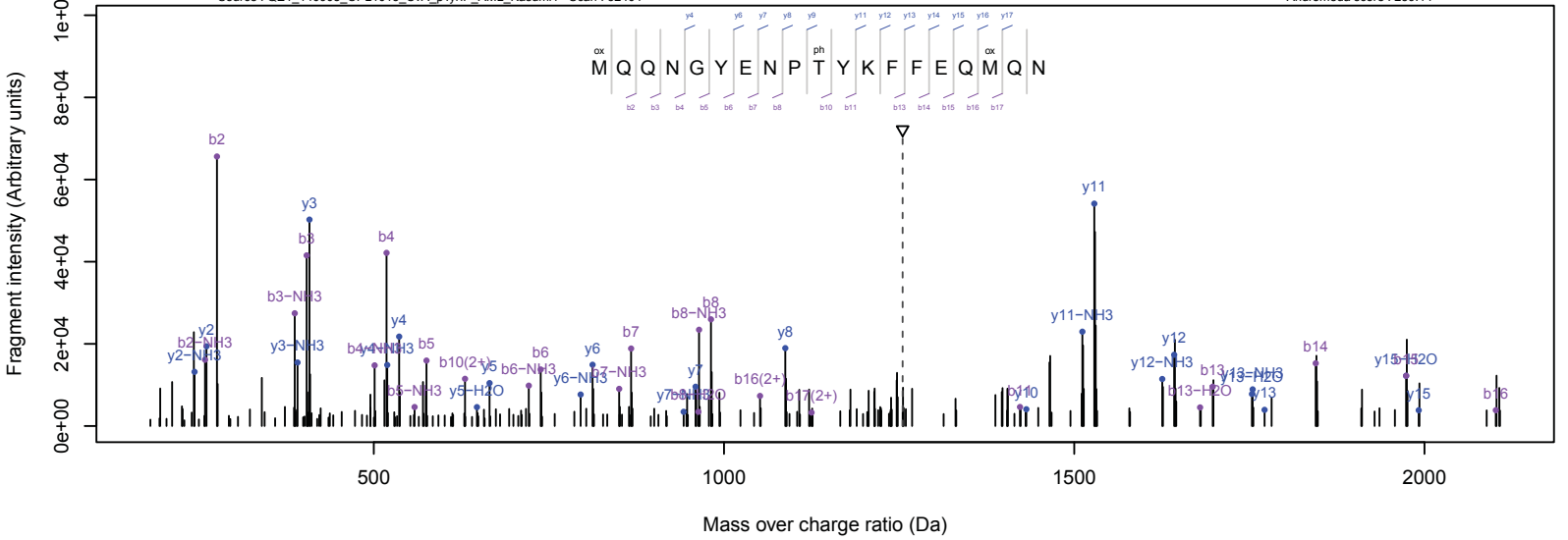
Andromeda score : 259.18



M(ox)QQNGYENPT(ph)YKFFEQM(ox)QN-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 32404

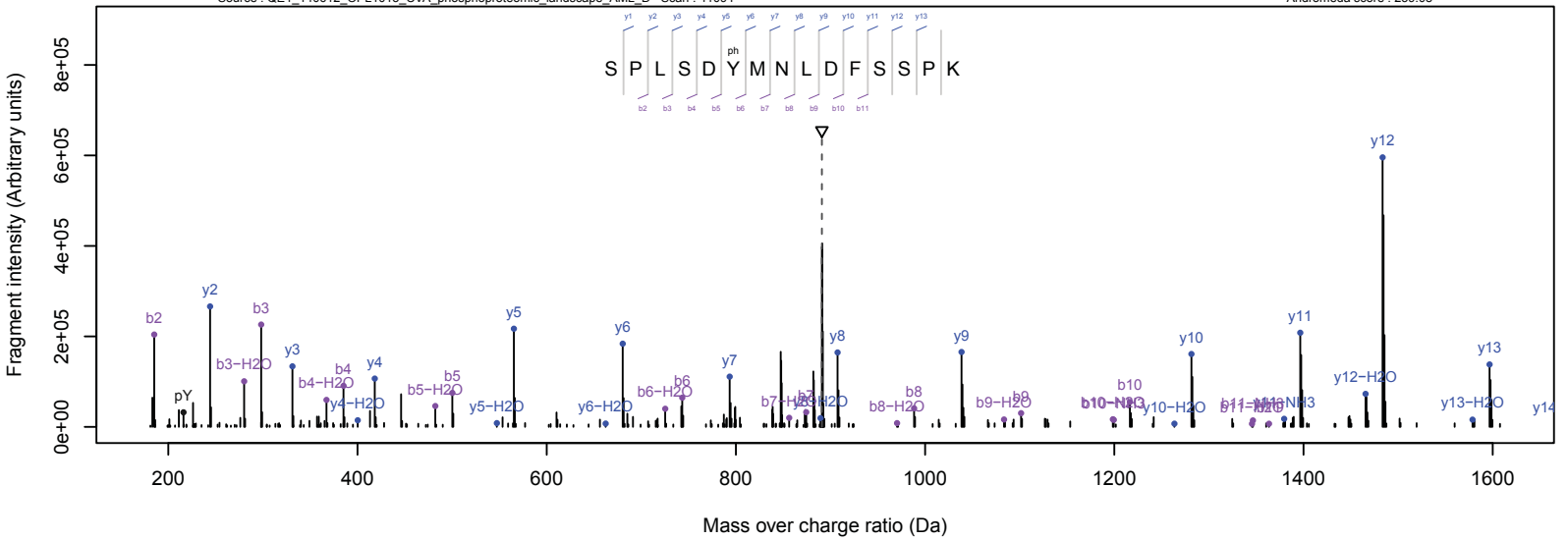
Andromeda score : 259.14



SPLSDY(ph)MNLDFSSPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41094

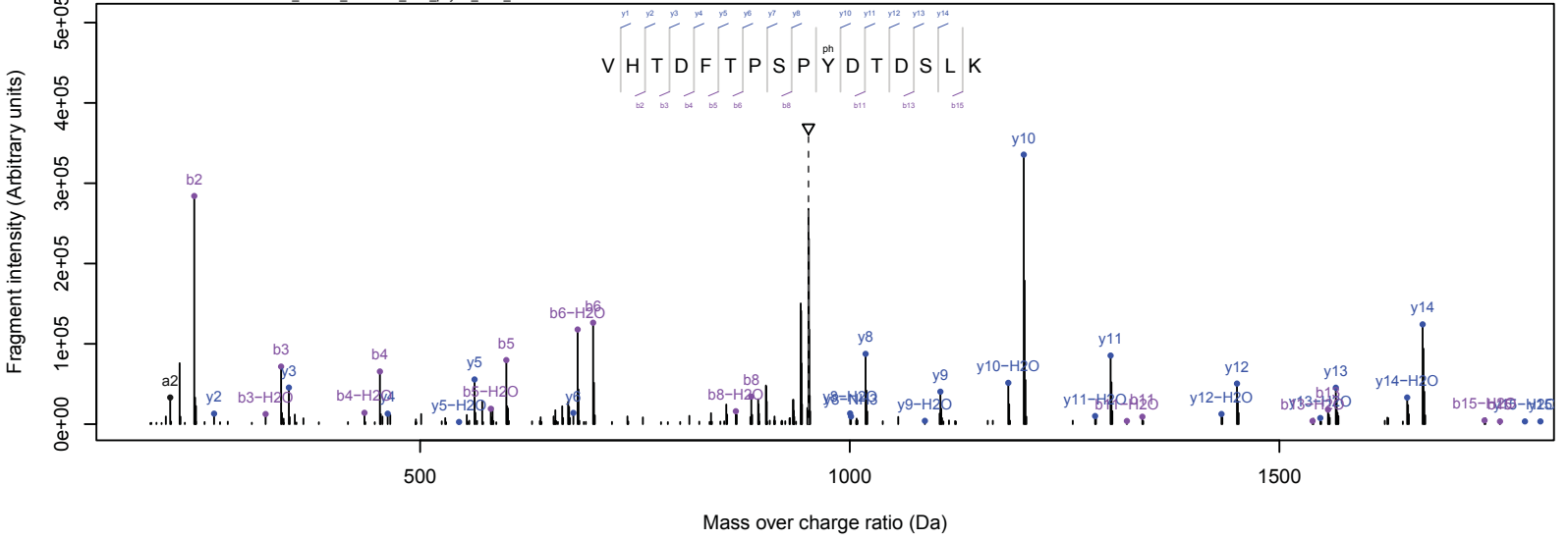
Andromeda score : 259.08



VHTDFTSPY(ph)DTSLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23341

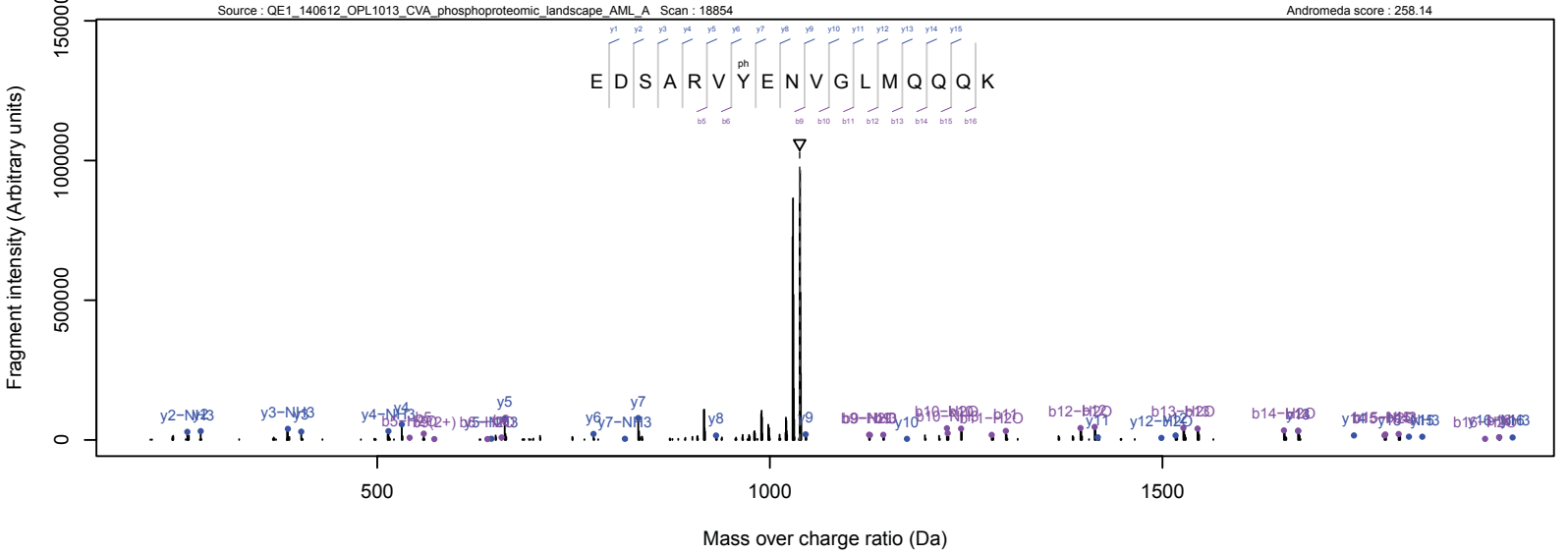
Andromeda score : 258.41



EDSARVY(ph)ENVGLMQQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 18854

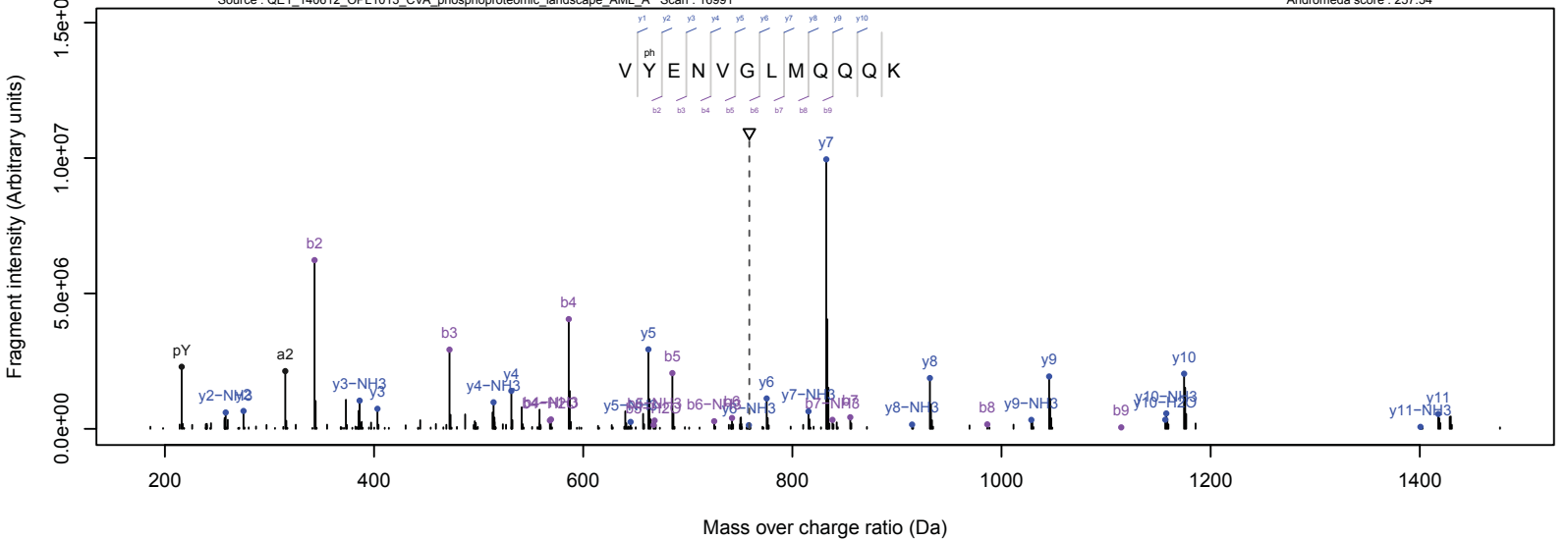
Andromeda score : 258.14



VY(ph)ENVGLMQQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16991

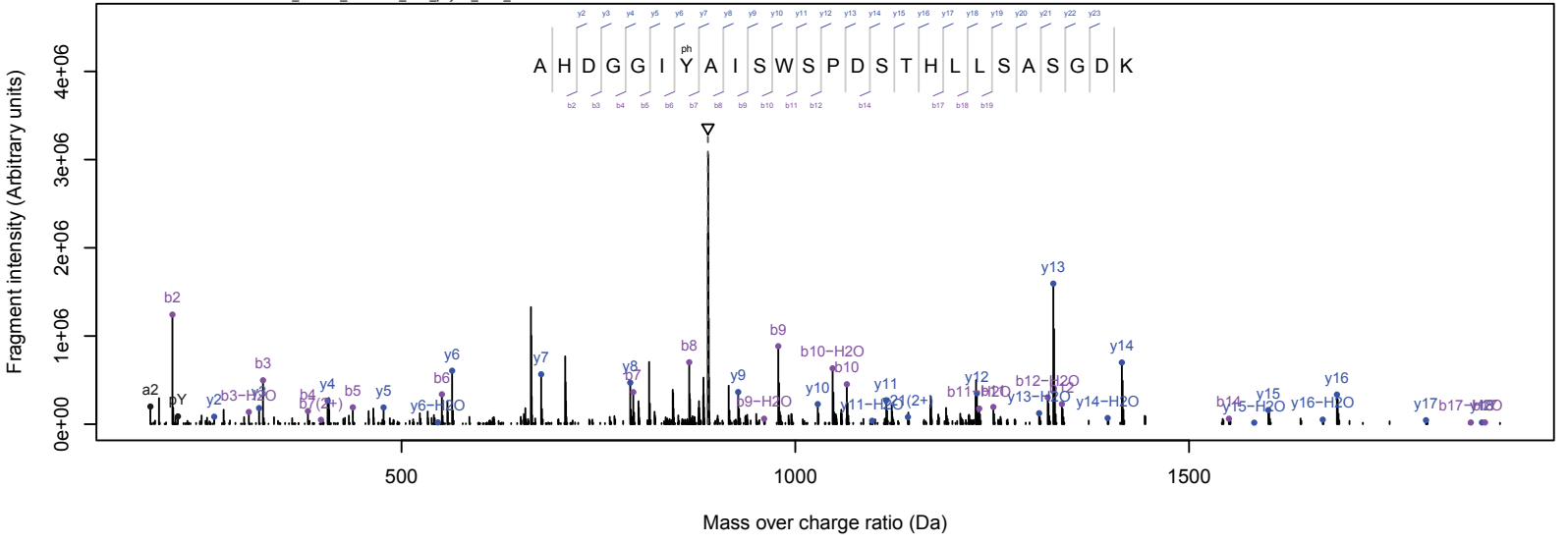
Andromeda score : 257.54



AHDGGIY(ph)AISWSPDSTHLLSASGDK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 38533

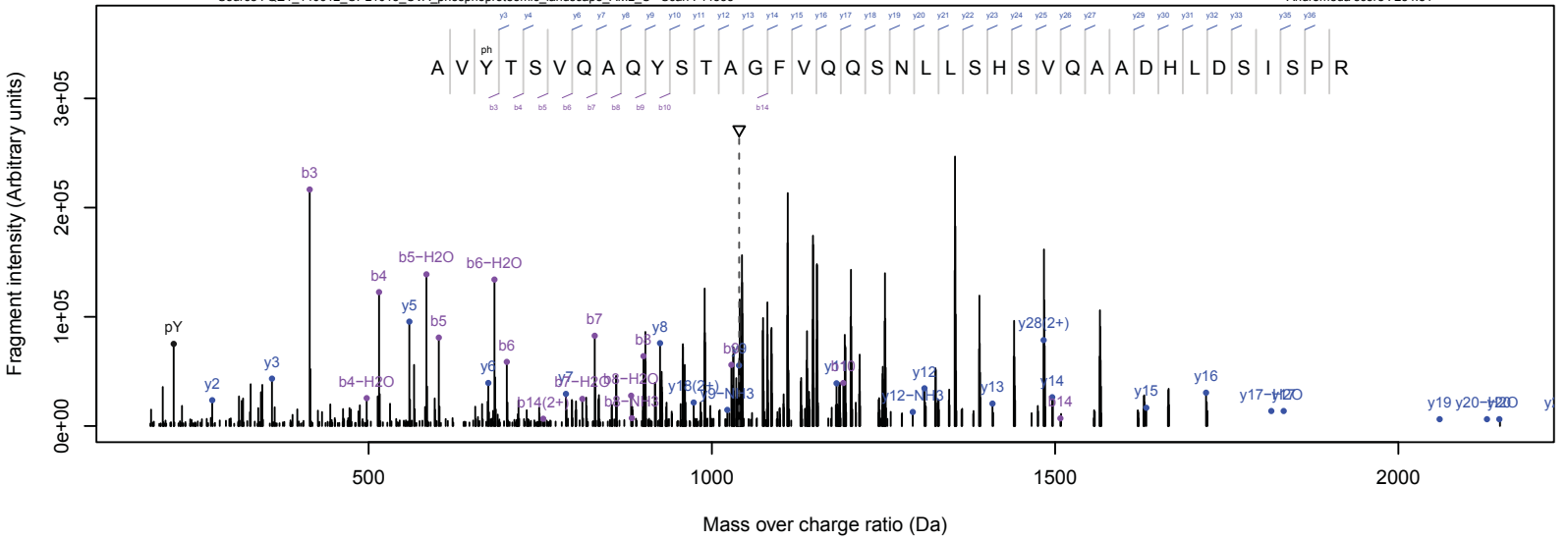
Andromeda score : 257.51



AVY(ph)TSVQAQYSTAGFVQQSNLLSHSVQAADHLDSISPR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44533

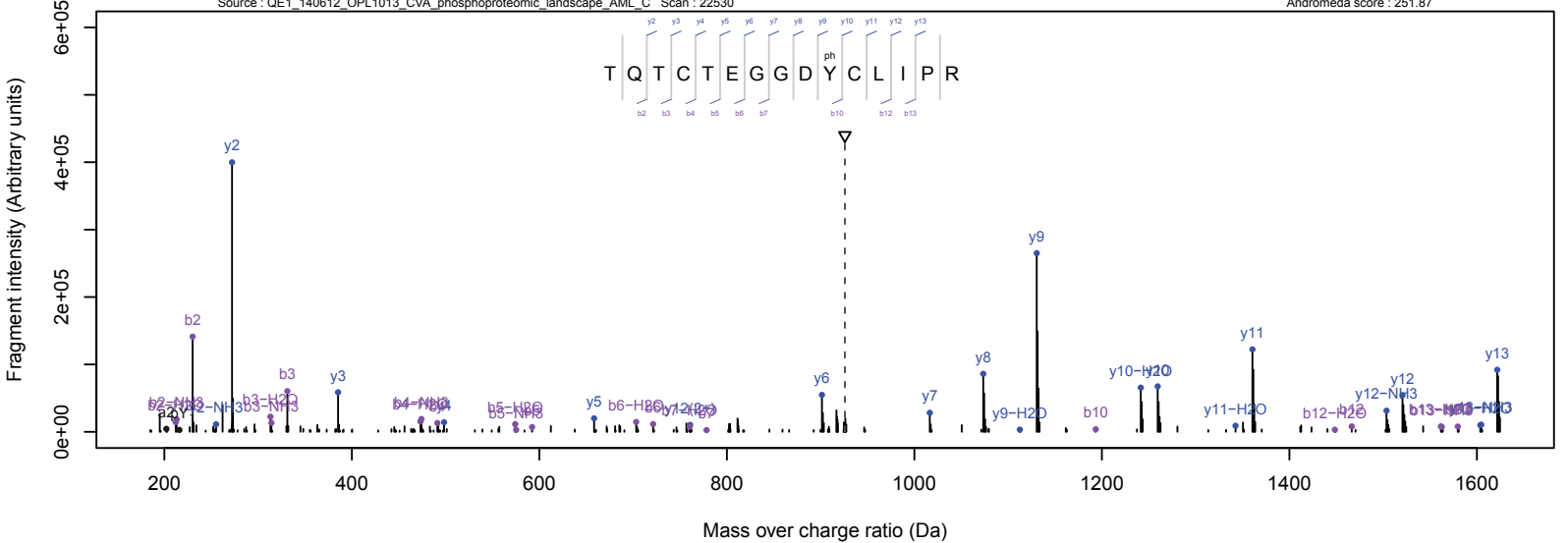
Andromeda score : 254.81



TQTCTEGGDY(ph)CLIPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22530

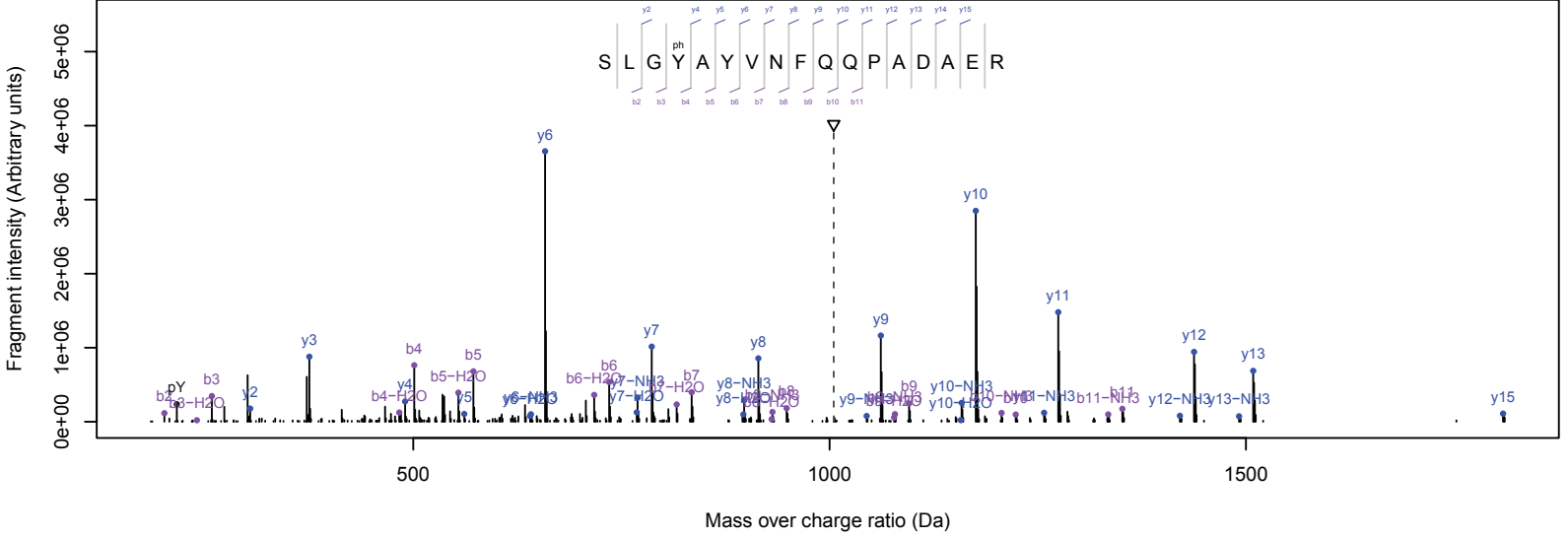
Andromeda score : 251.87



SLGY(ph)AYVNFQPPADAER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31119

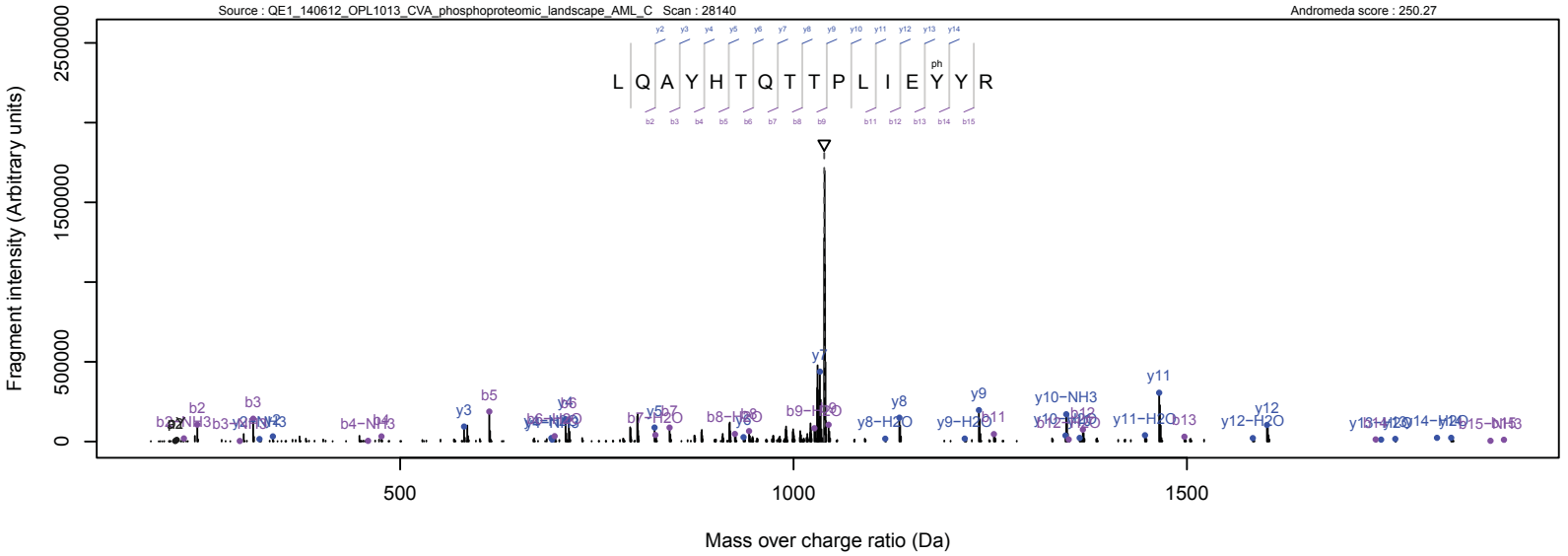
Andromeda score : 251.02



LQAYHTQTTPLEIYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28140

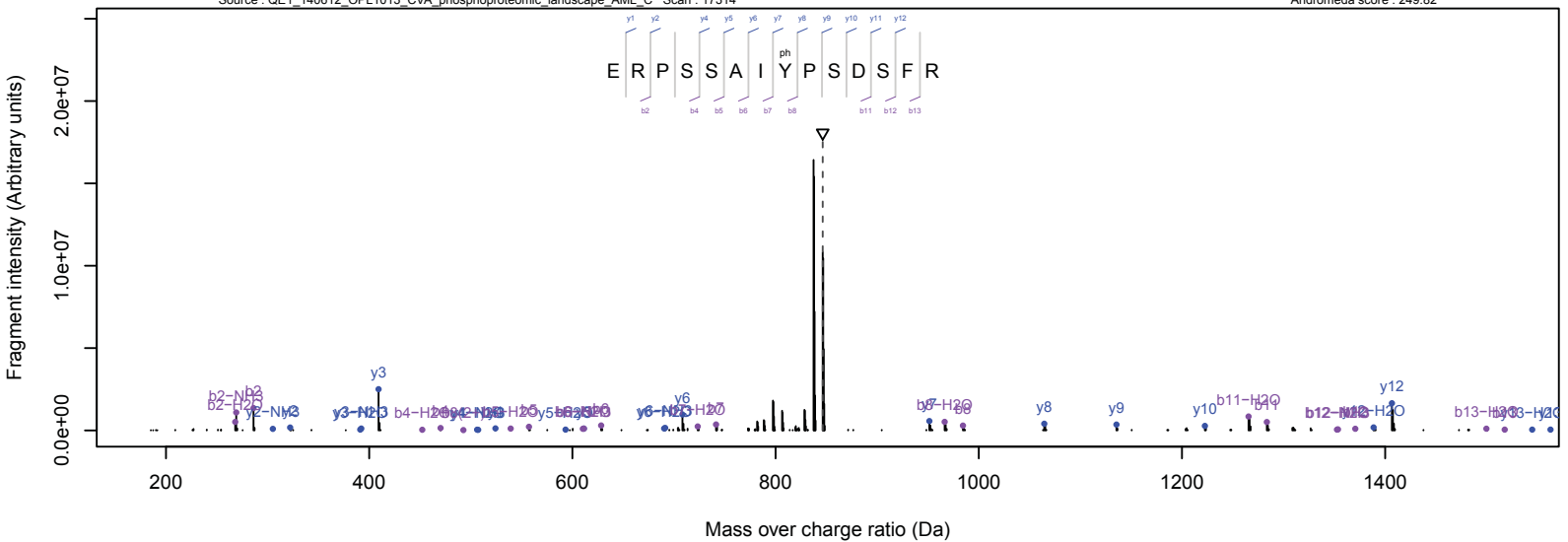
Andromeda score : 250.27



ERPSSAIYPSDSFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17314

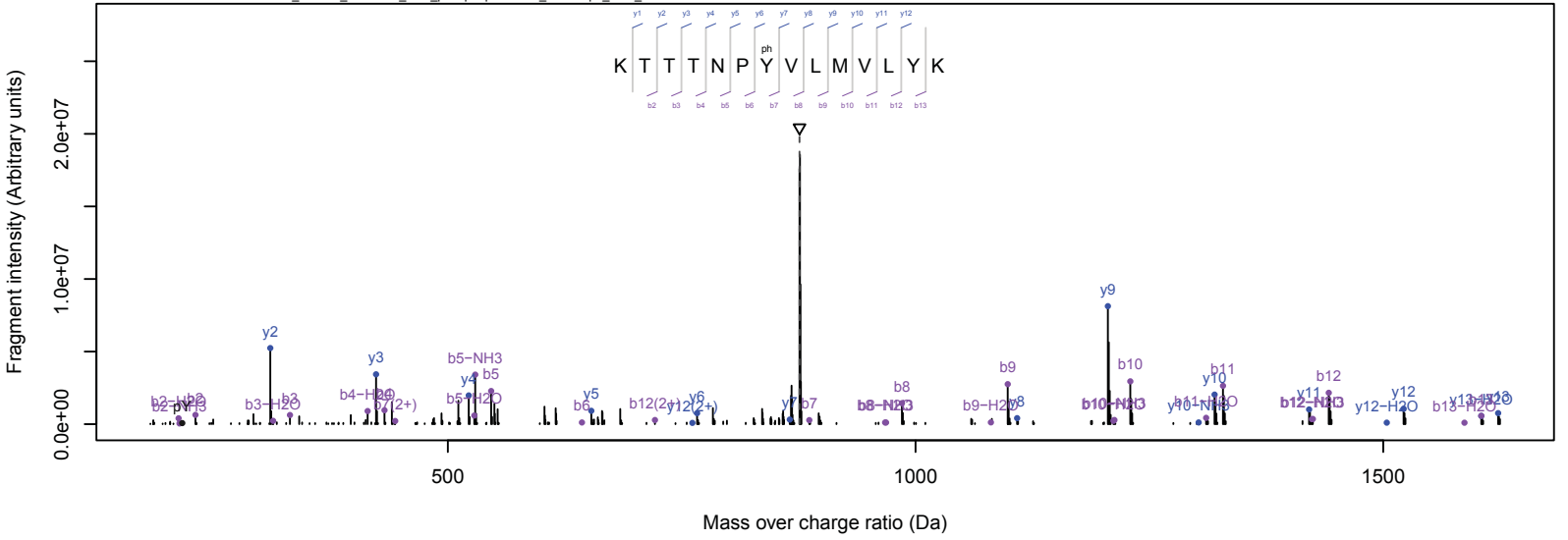
Andromeda score : 249.82



**KTTTNPY(ph)VLMVLYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36139

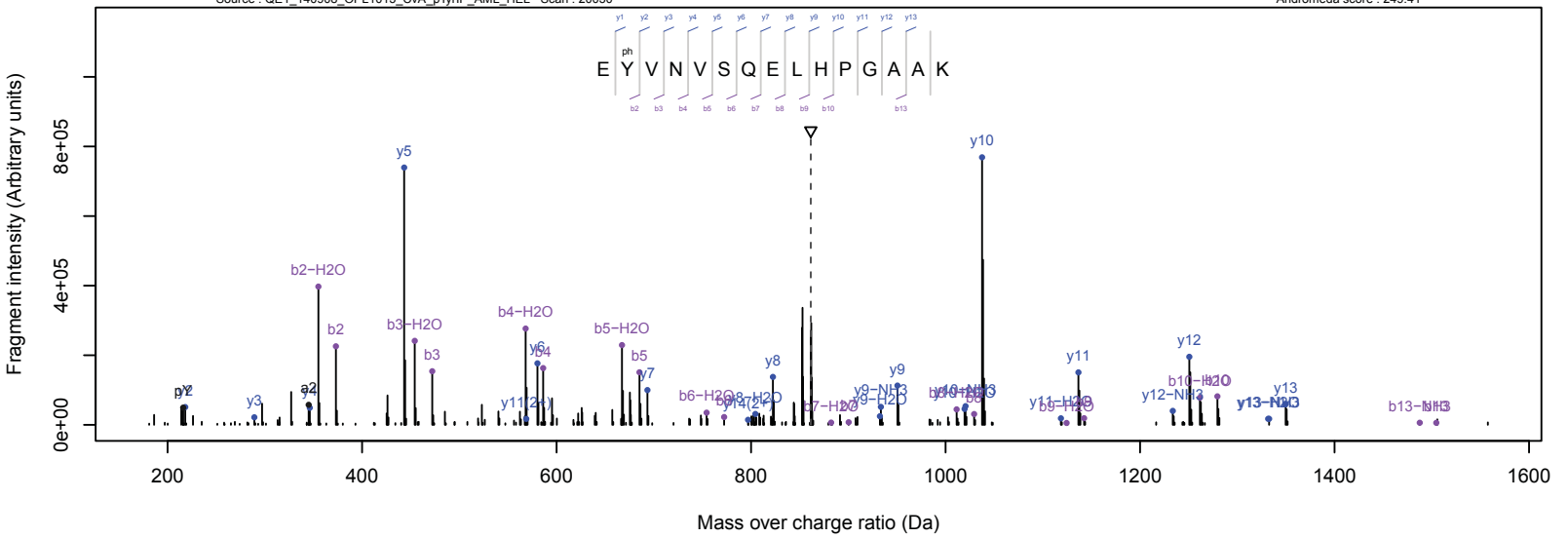
Andromeda score : 249.82



**EY(ph)VNVSQELHPGAAK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 20030

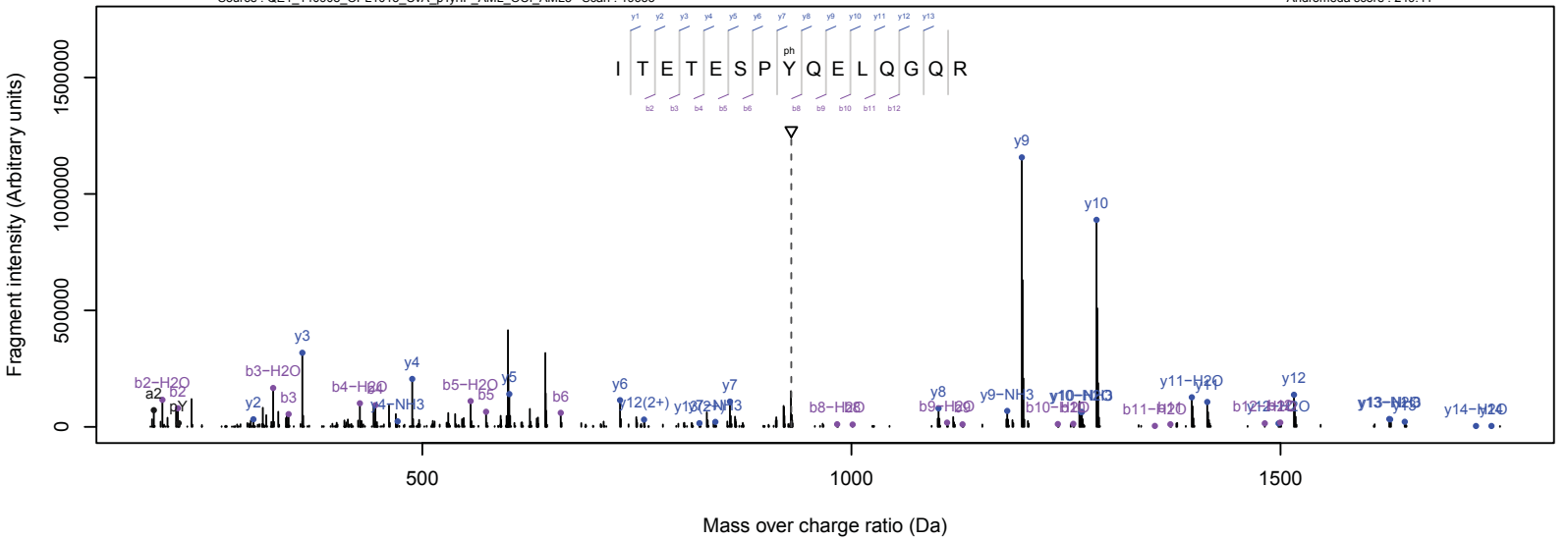
Andromeda score : 249.41



**ITETESPY(ph)QELGQR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 19653

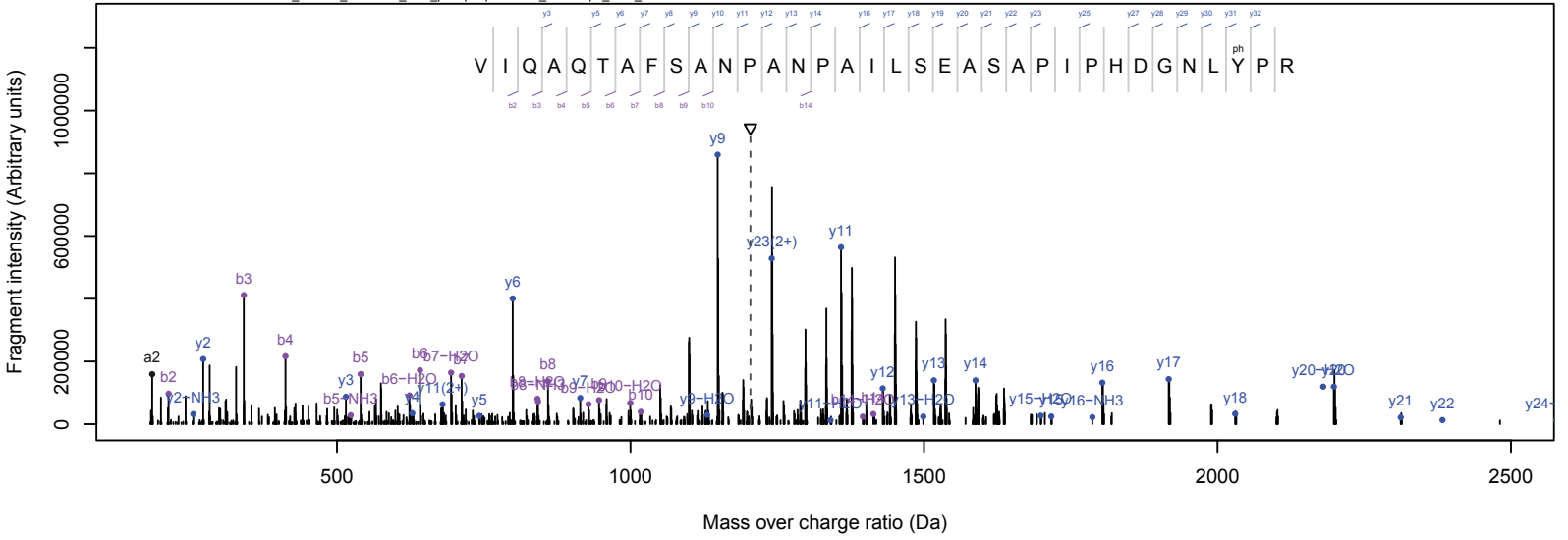
Andromeda score : 249.41



VIQAQTAFSANPANPAILSEASAPIPHDGNLYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 43240

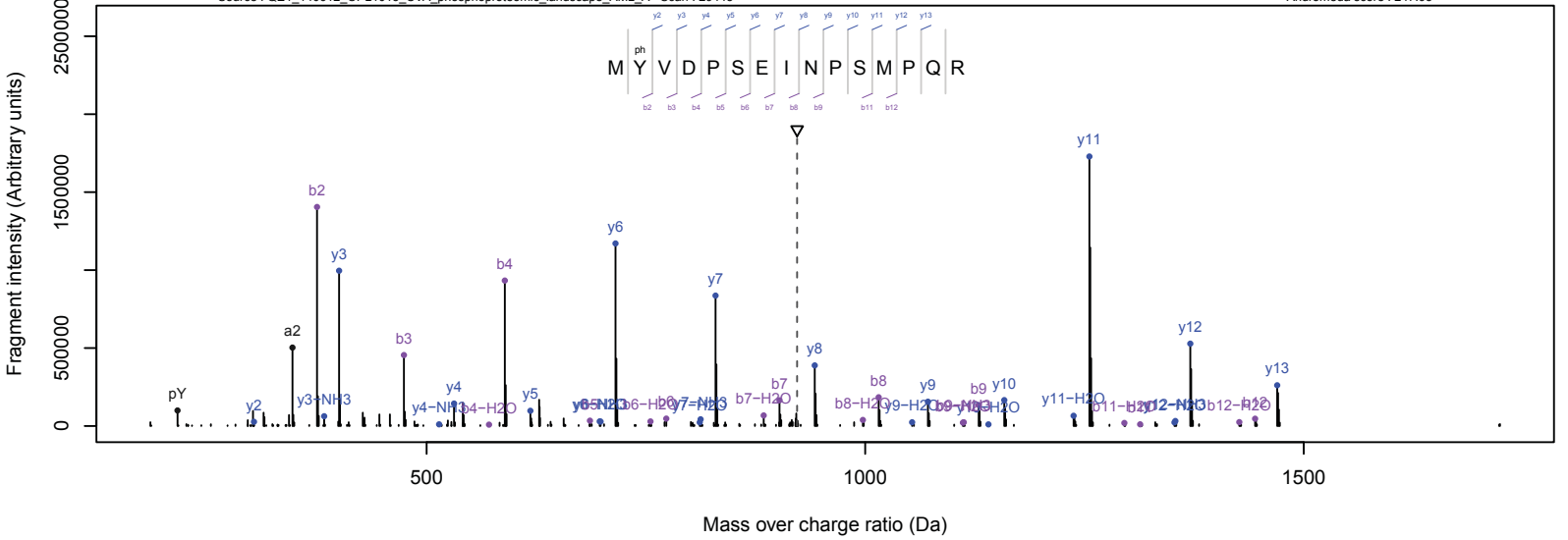
Andromeda score : 248.82



MY(ph)VDPSEINPSPMPQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29443

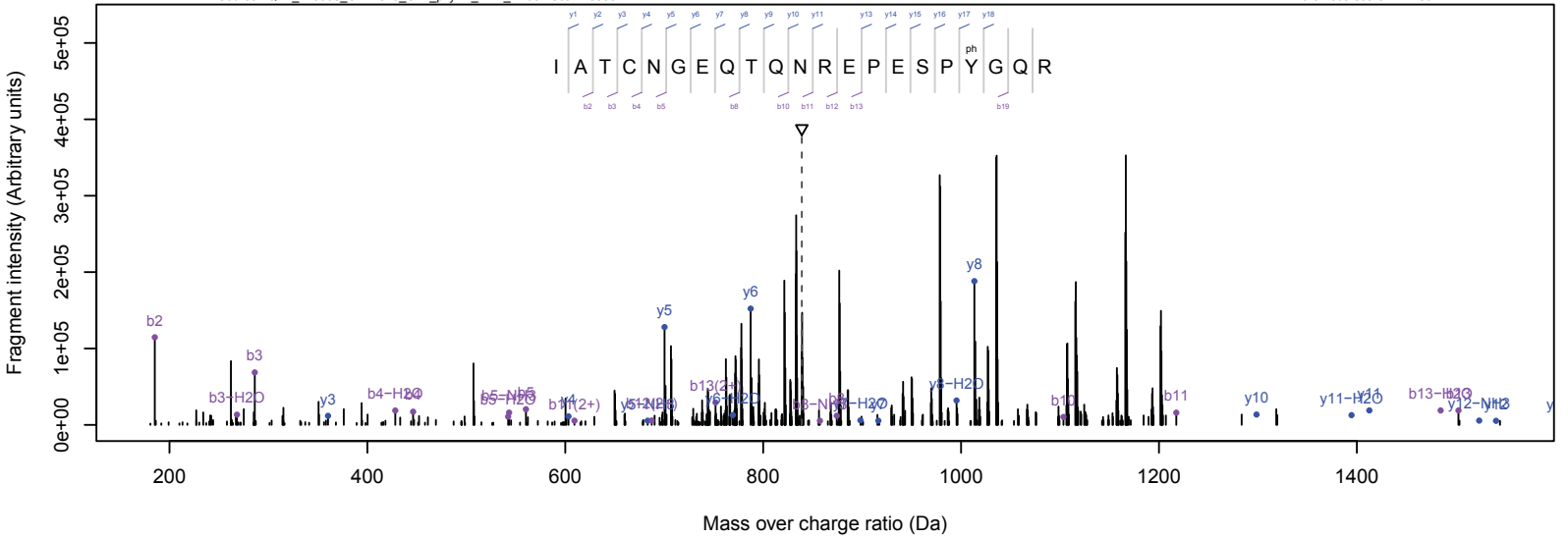
Andromeda score : 247.93

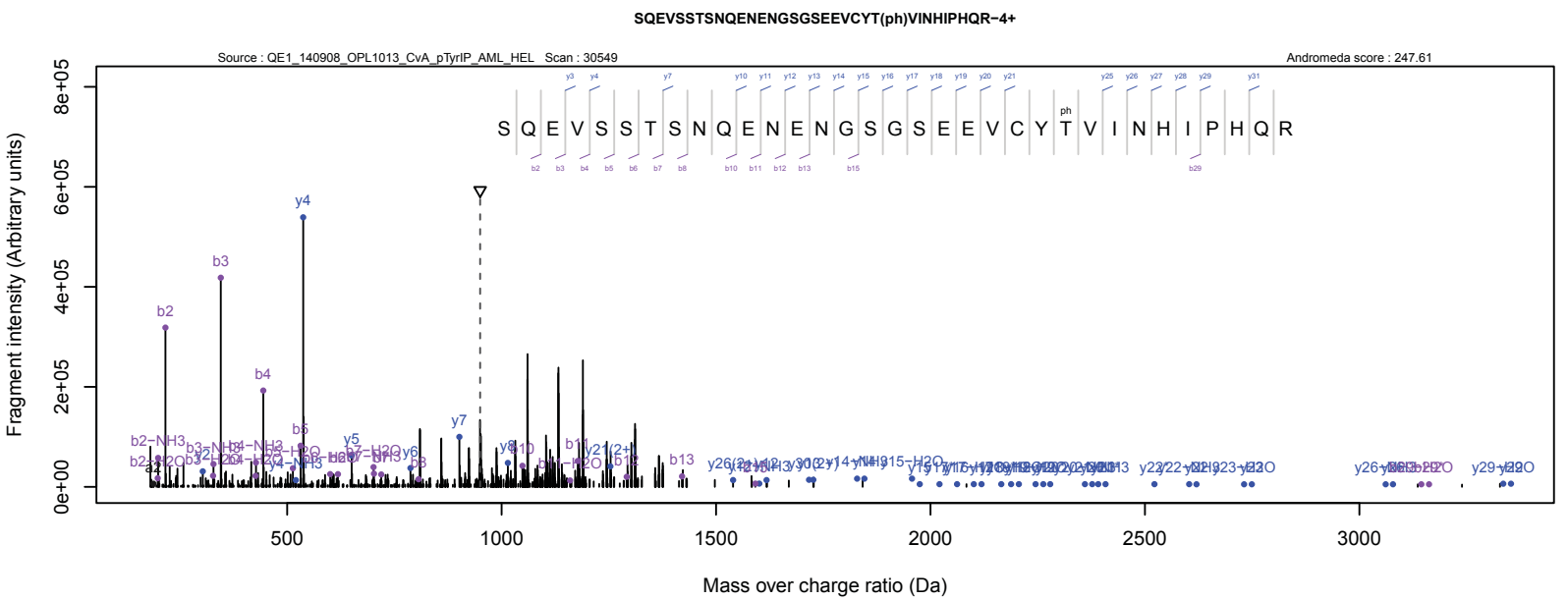
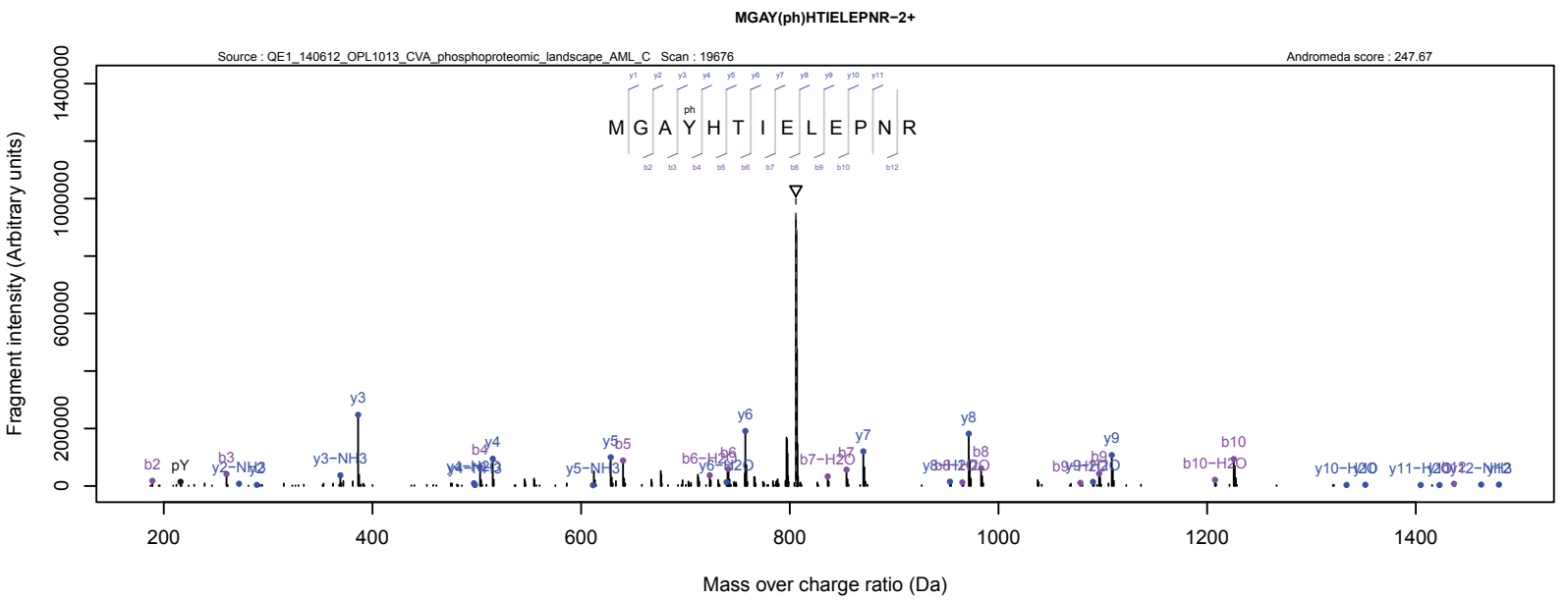
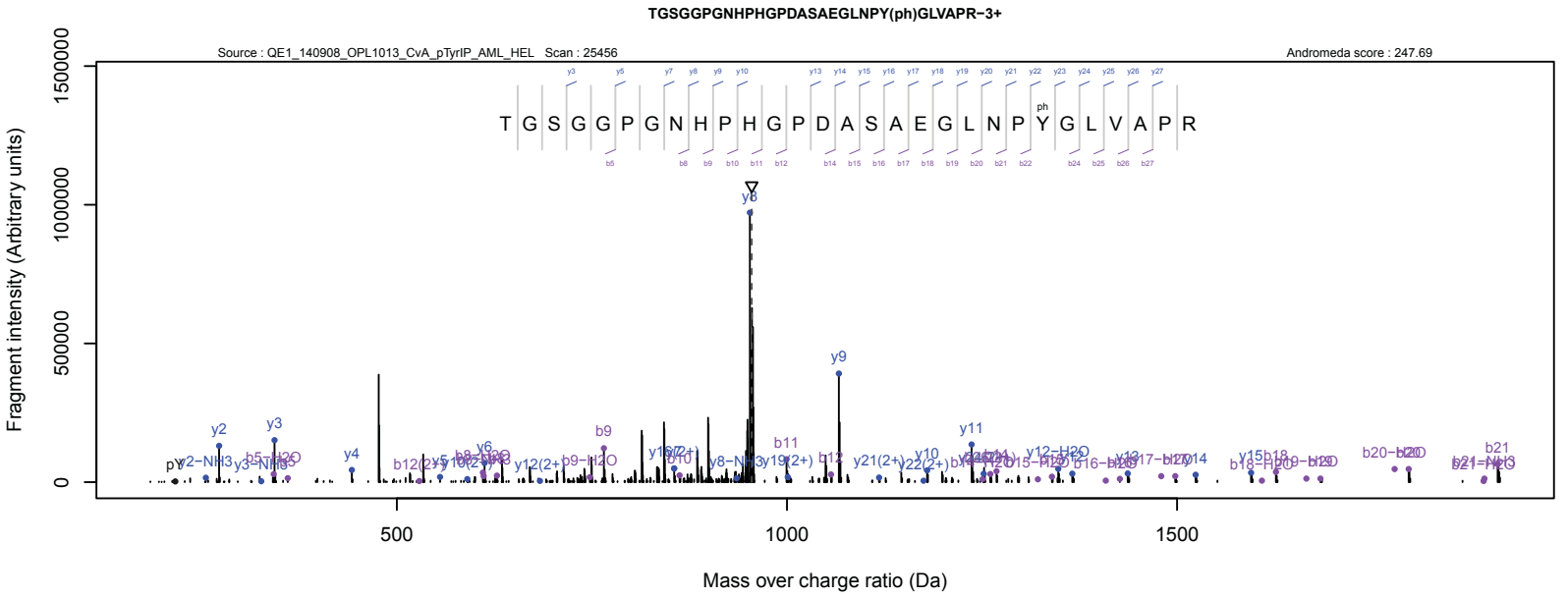


IATCNGETQNRPEESPY(ph)GQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 8553

Andromeda score : 247.89





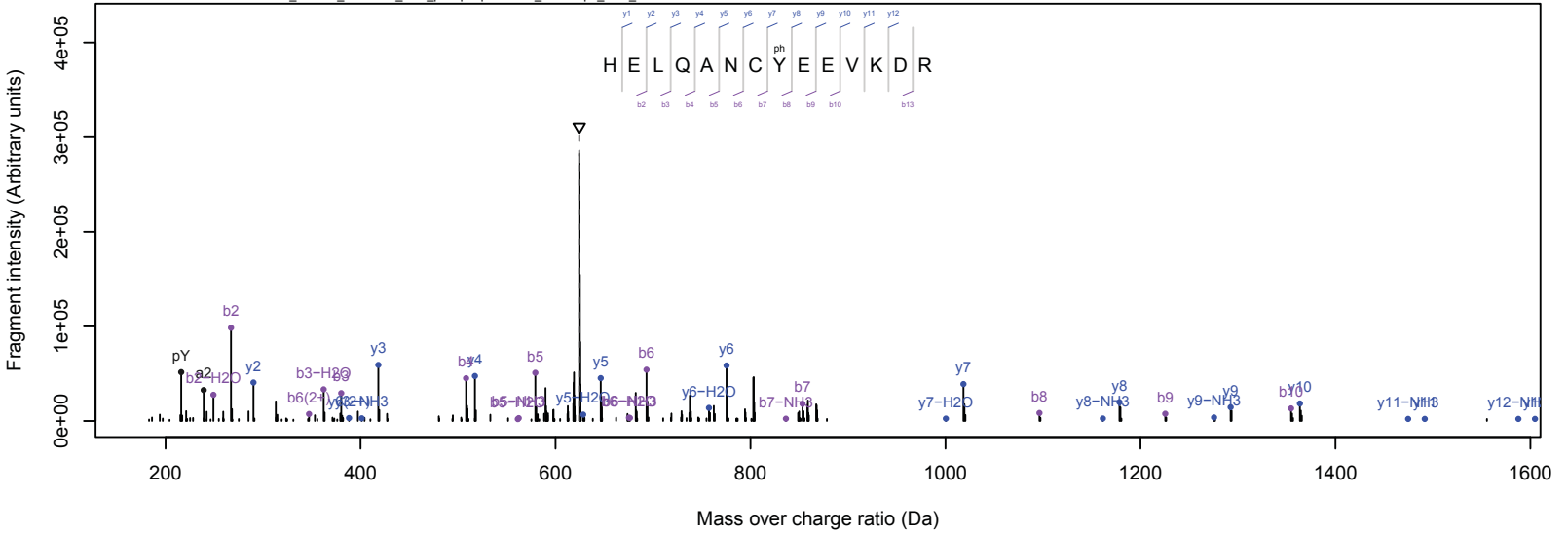




HELQANCY(ph)EEVKDR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 8577

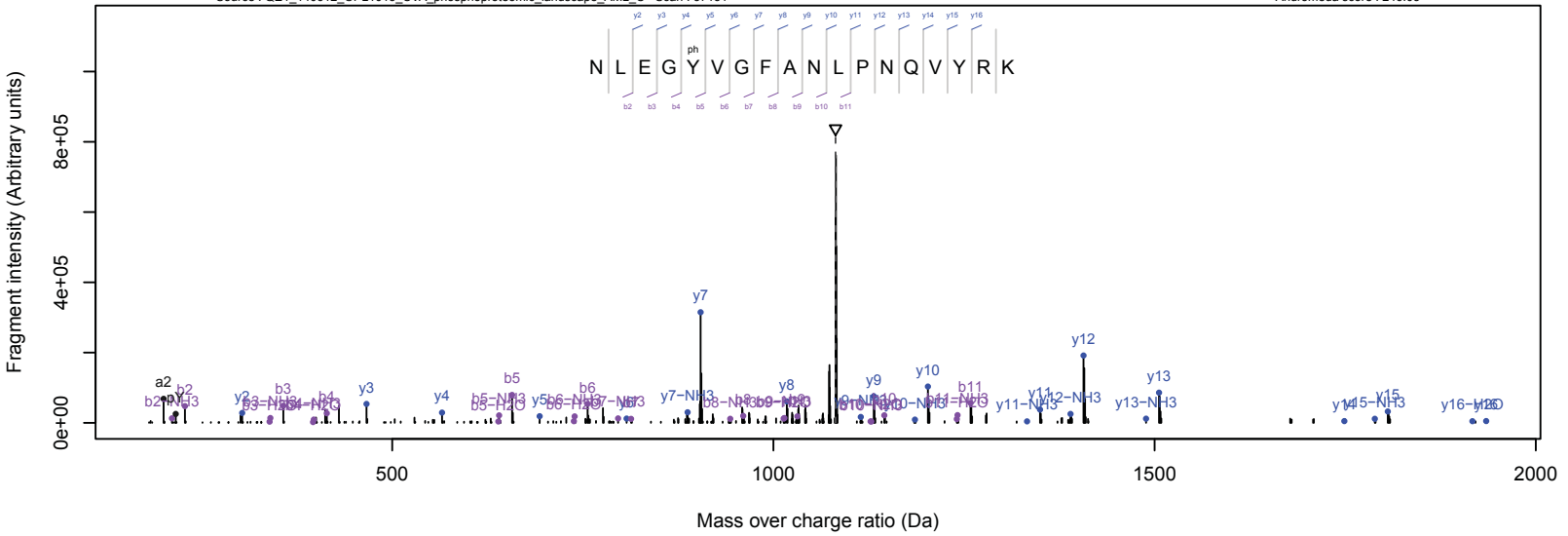
Andromeda score: 246.18



NLEGY(ph)VGFLNPQVYRK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37131

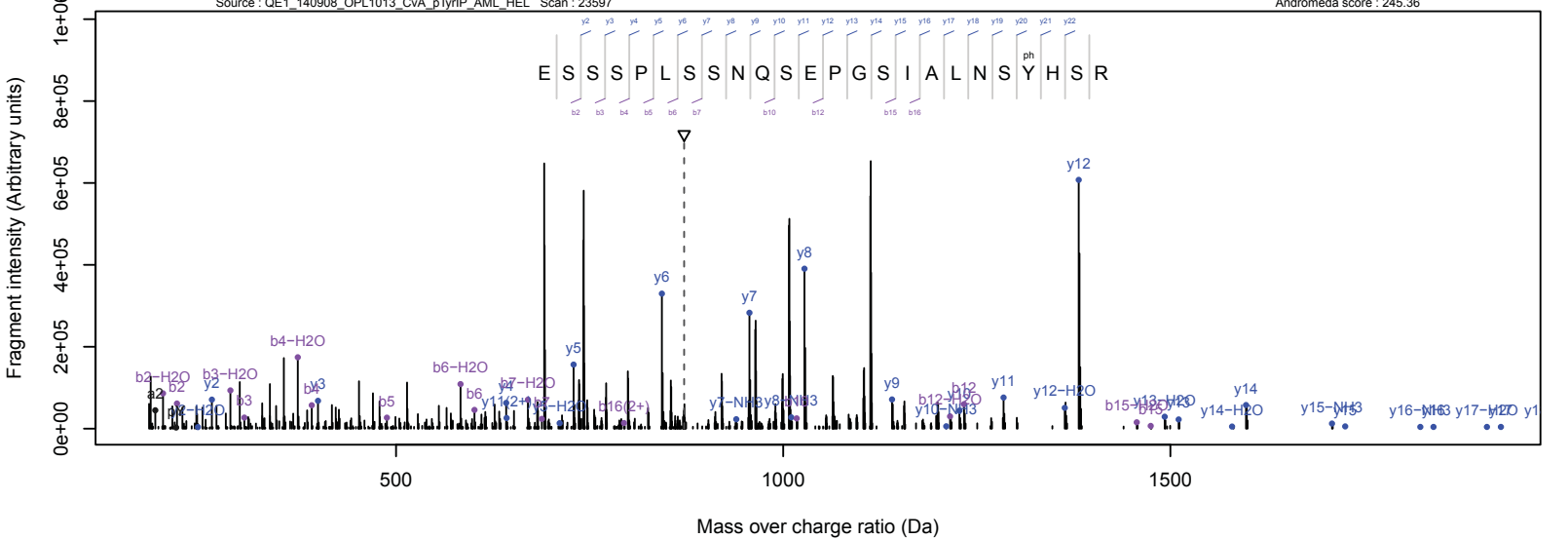
Andromeda score: 246.03



ESSSPLSNQSEPGSIALNSY(ph)HSR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 23597

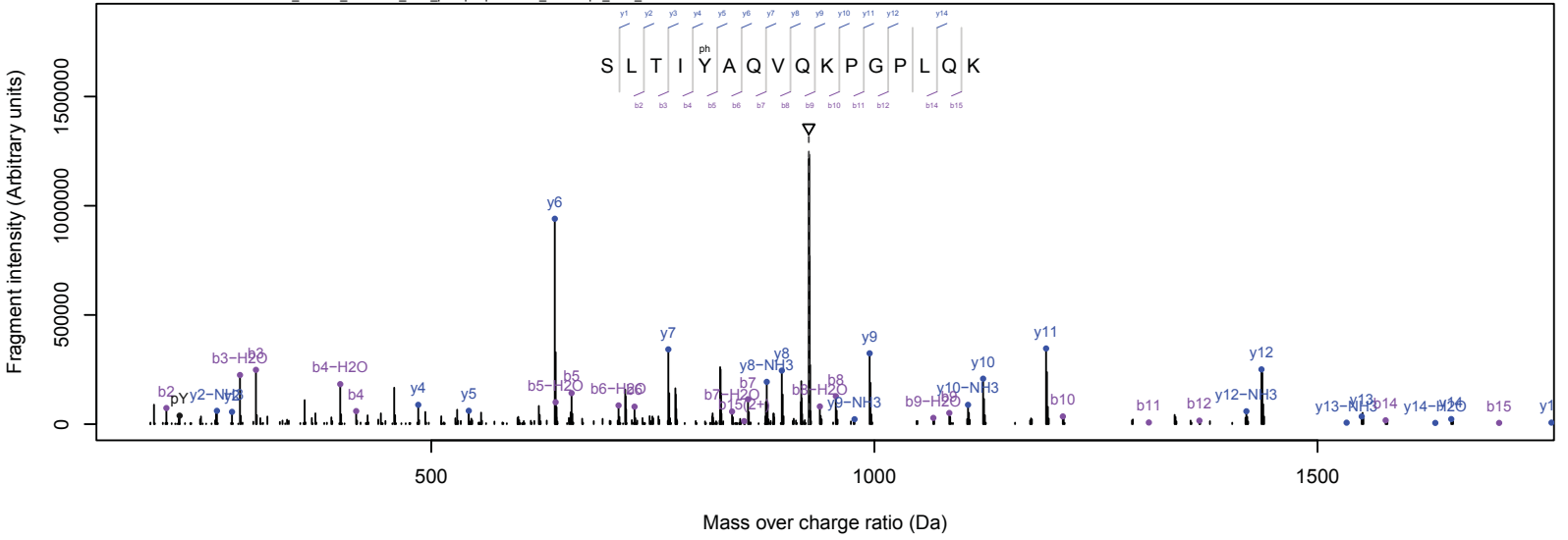
Andromeda score: 245.36



SLTIY(ph)AQVQKPGPLQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25302

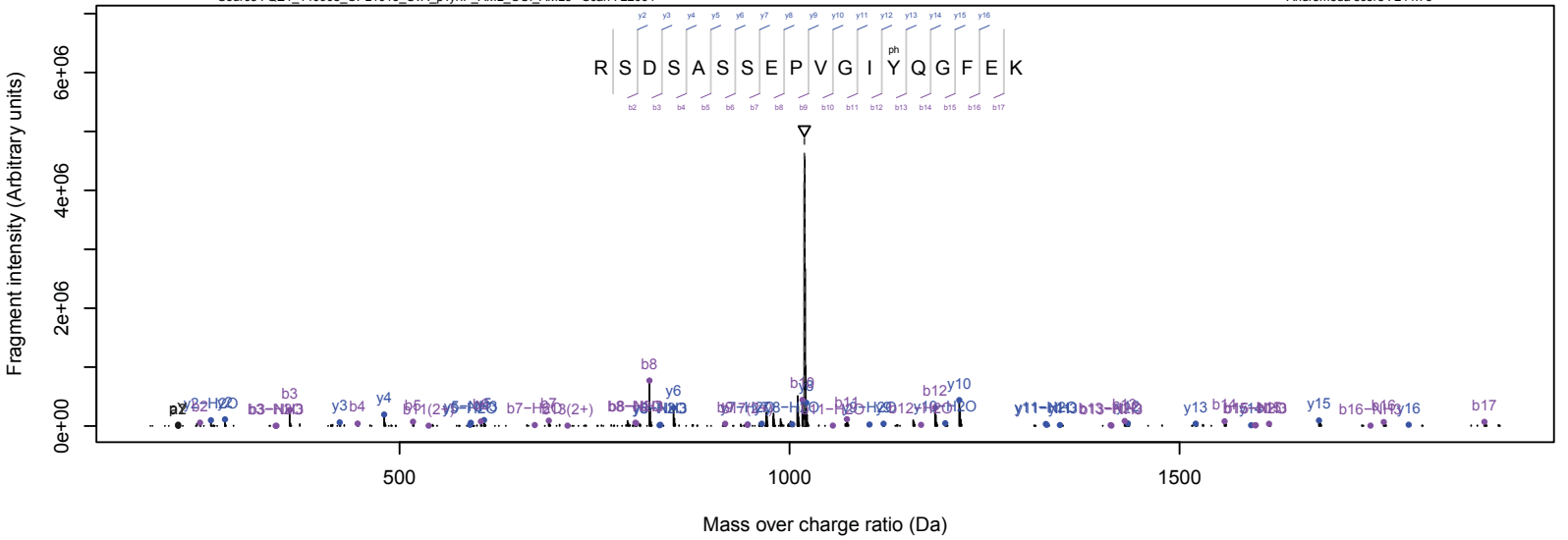
Andromeda score : 245.19



RSDSASSEPVGIYQGFEEK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 22664

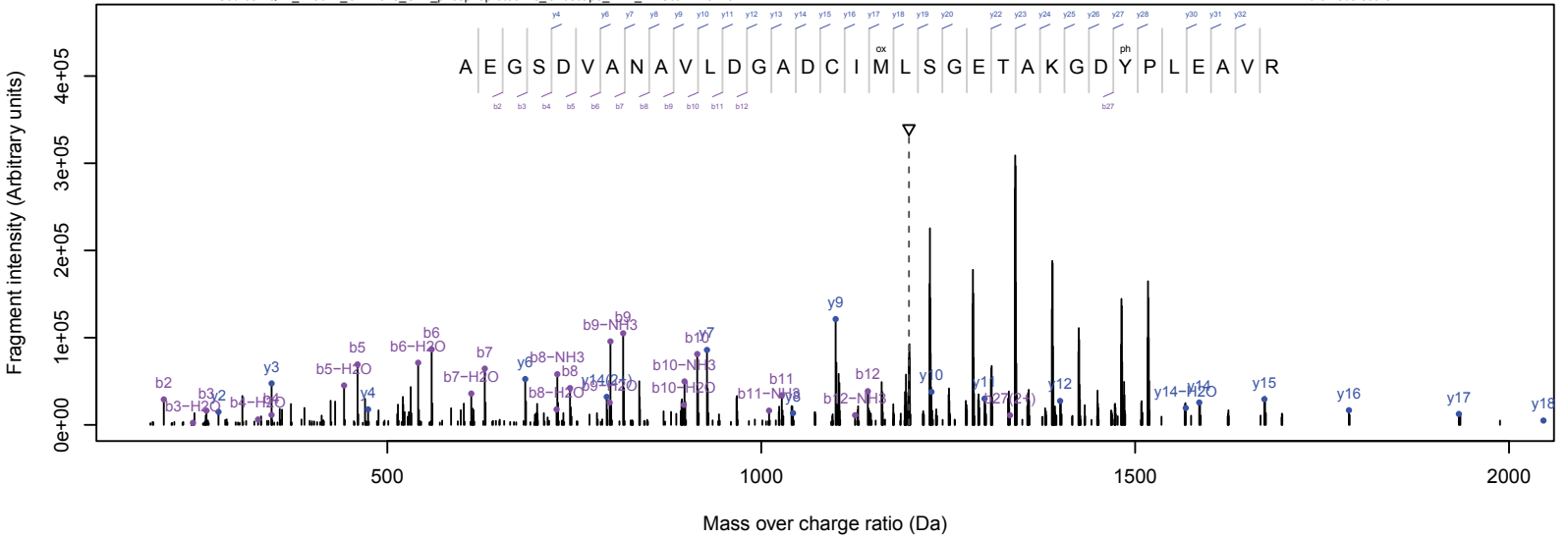
Andromeda score : 244.73



AEGSDVANAVLDGADCIMLSGETAKGDYPLEAVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 45143

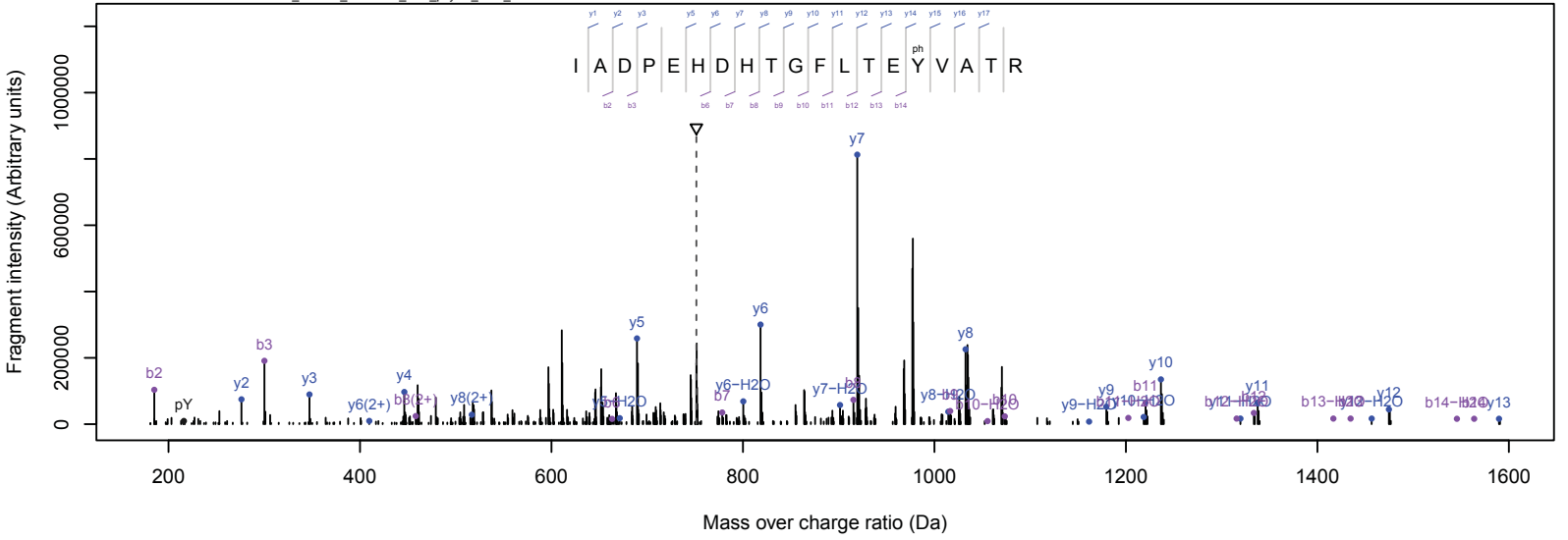
Andromeda score : 244.1



IADPEHDHTGFLTEY(ph)VATR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 28691

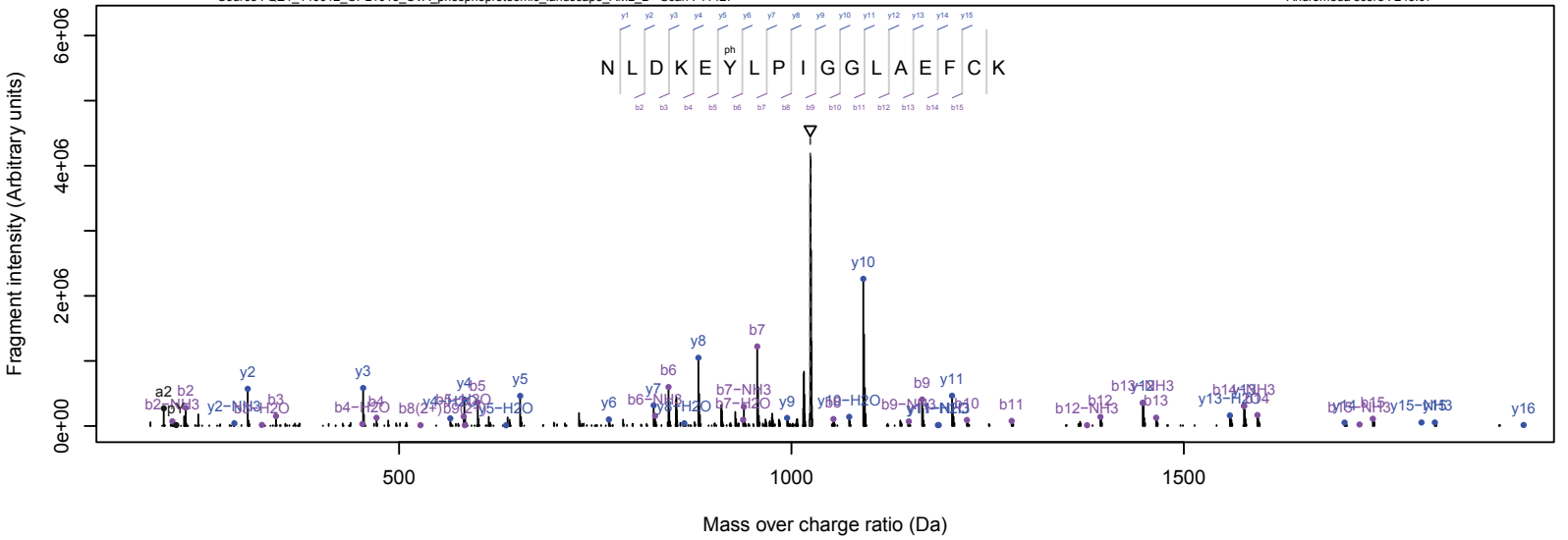
Andromeda score : 244.1



NLDKEY(ph)LPIGGLAEFC-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41427

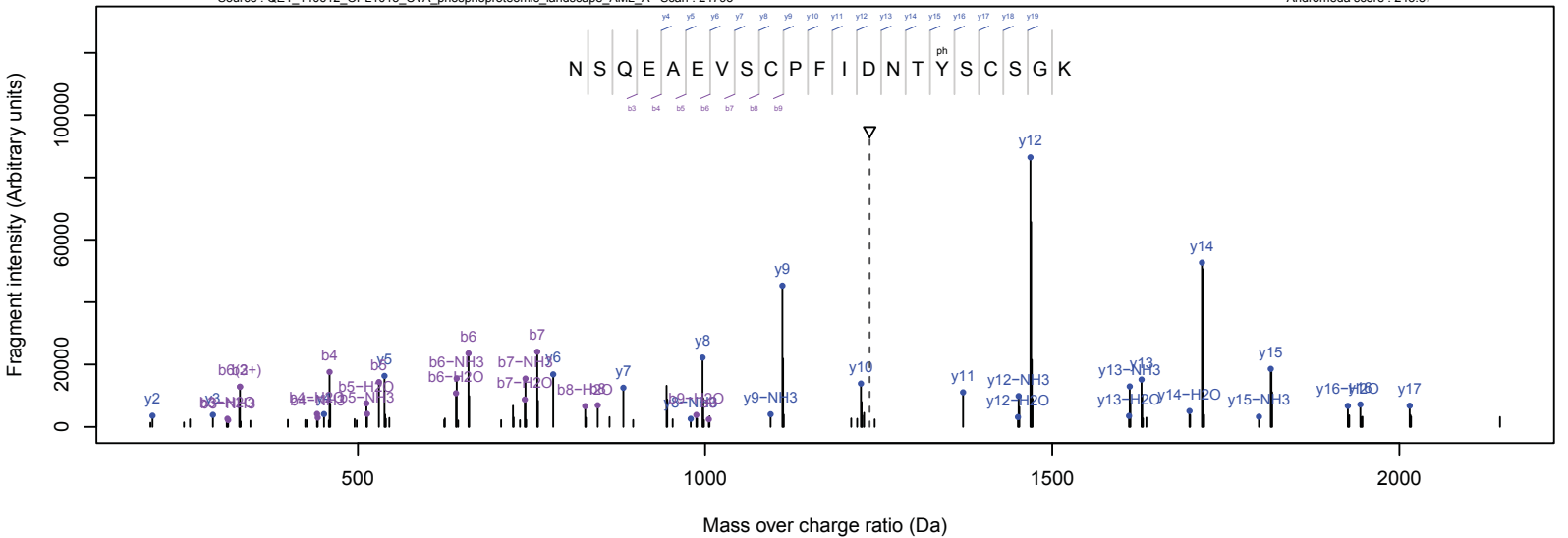
Andromeda score : 243.97



NSQAEVSCPFIDNTY(ph)SCSGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 24793

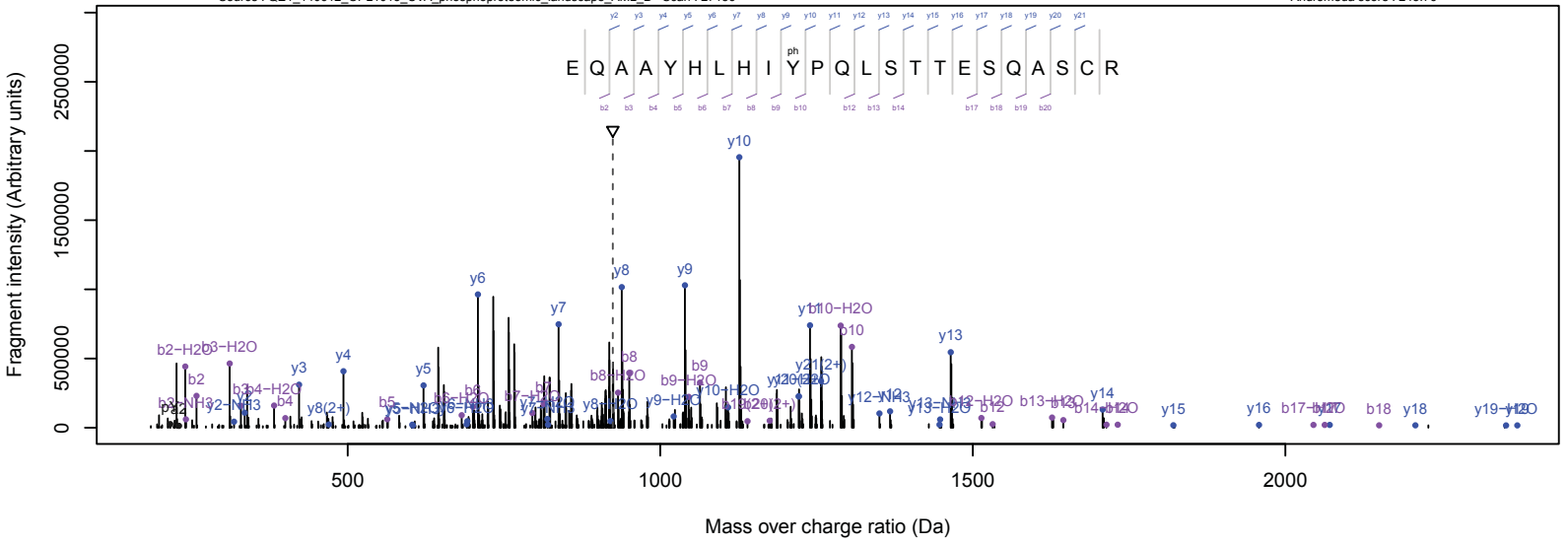
Andromeda score : 243.87



**EQAAYHLHIY(ph)PQLSTTESQASCR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27156

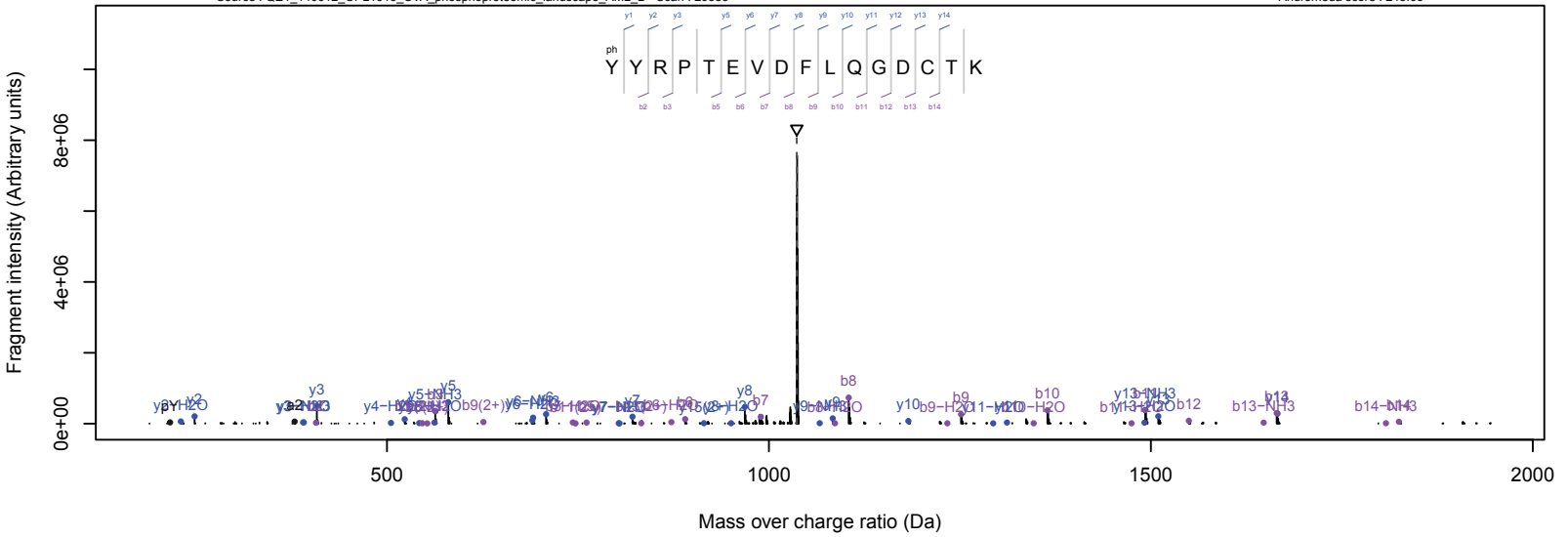
Andromeda score : 243.76



**Y(ph)YRPTEVDFLQGDCTK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29583

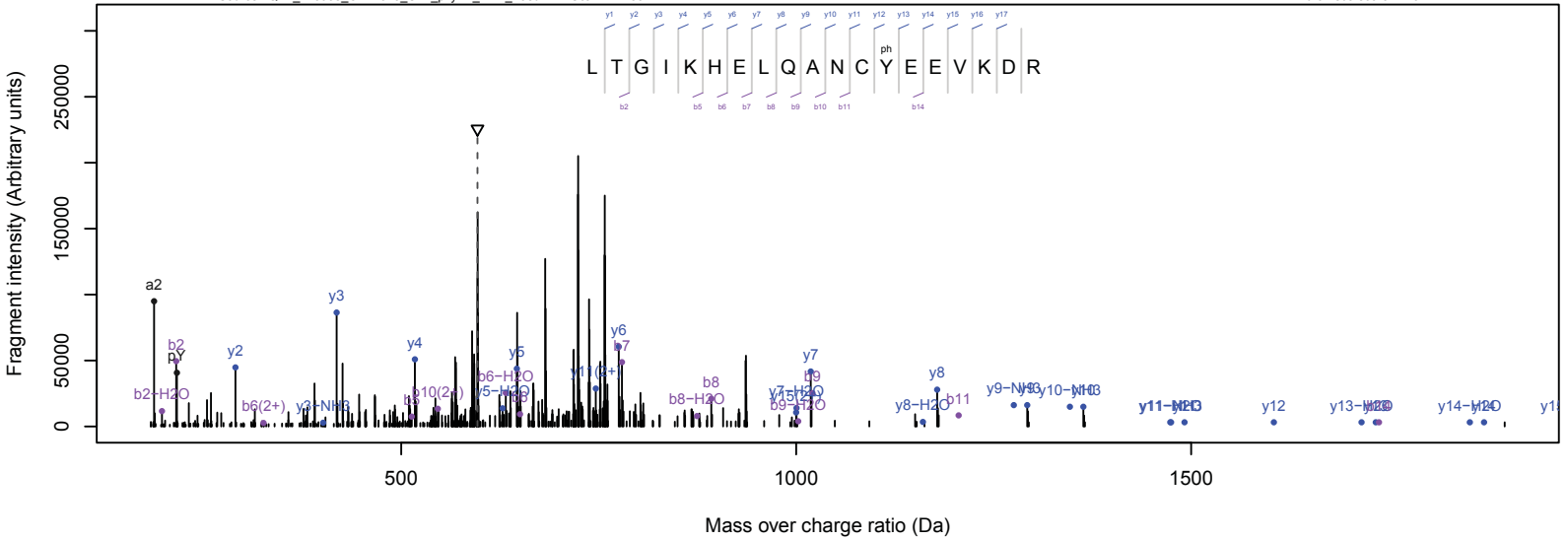
Andromeda score : 243.63



**LTGIKHELQANCY(ph)EEVKDR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 14453

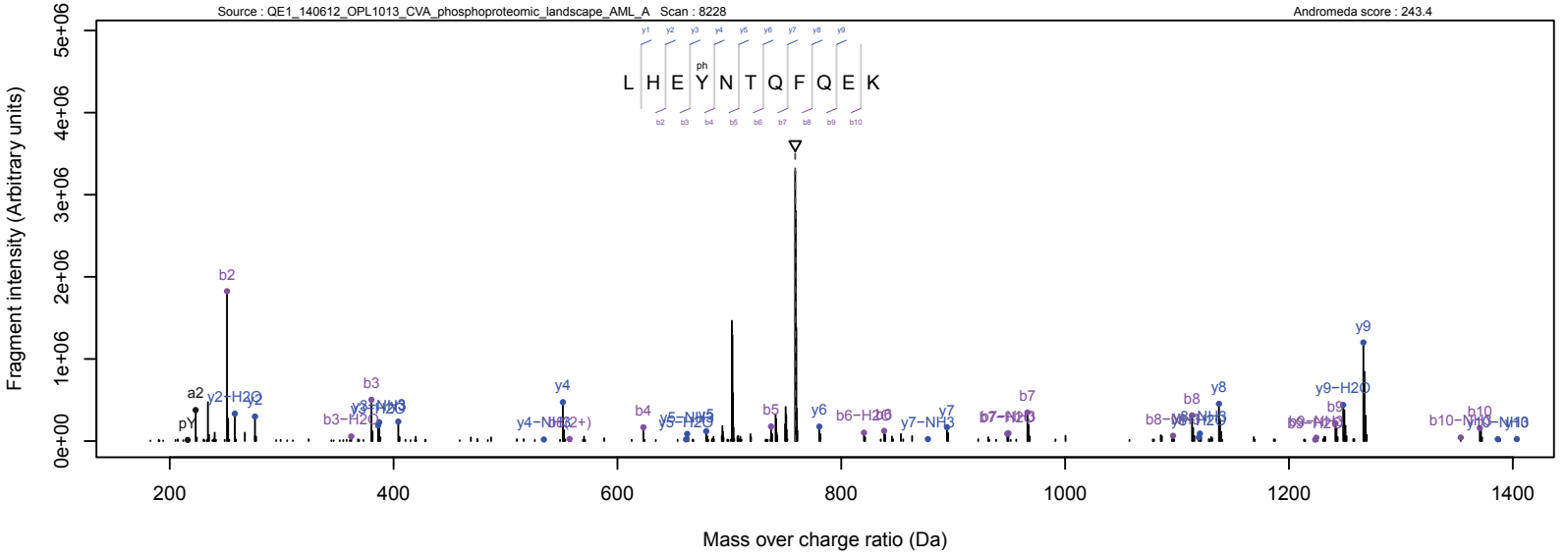
Andromeda score : 243.41



LHEY(ph)NTQFQEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 8228

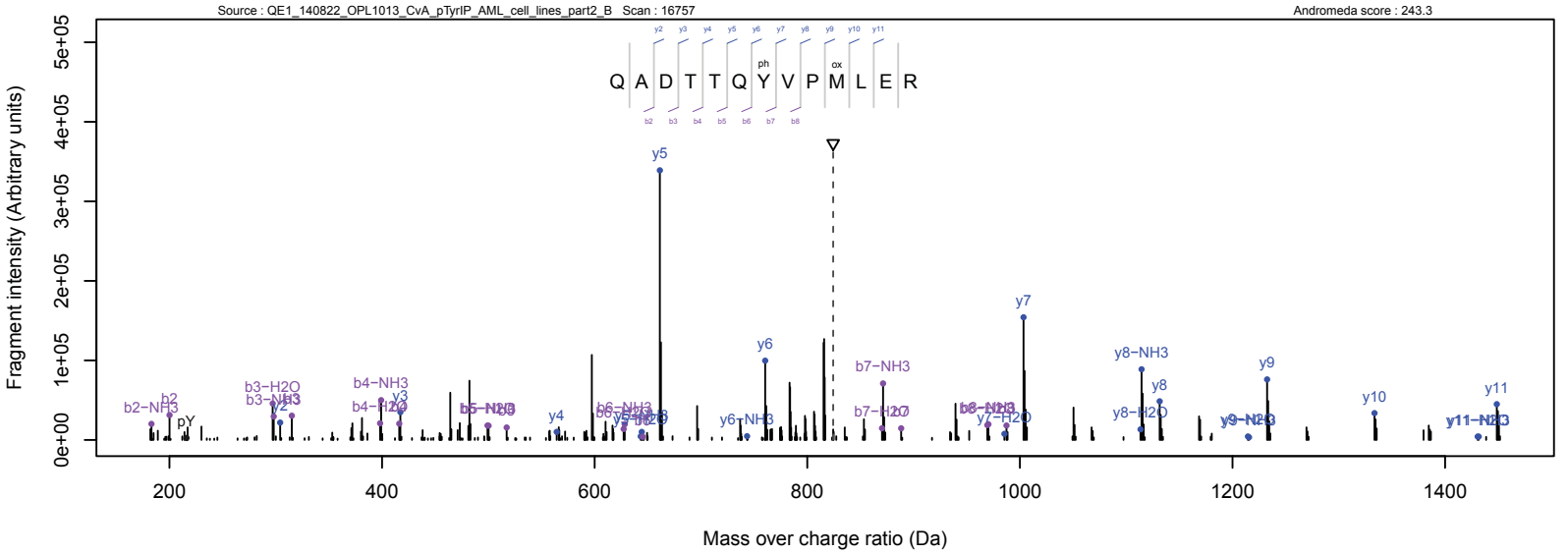
Andromeda score : 243.4



QADTTQY(ph)VPM(ox)LER-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 16757

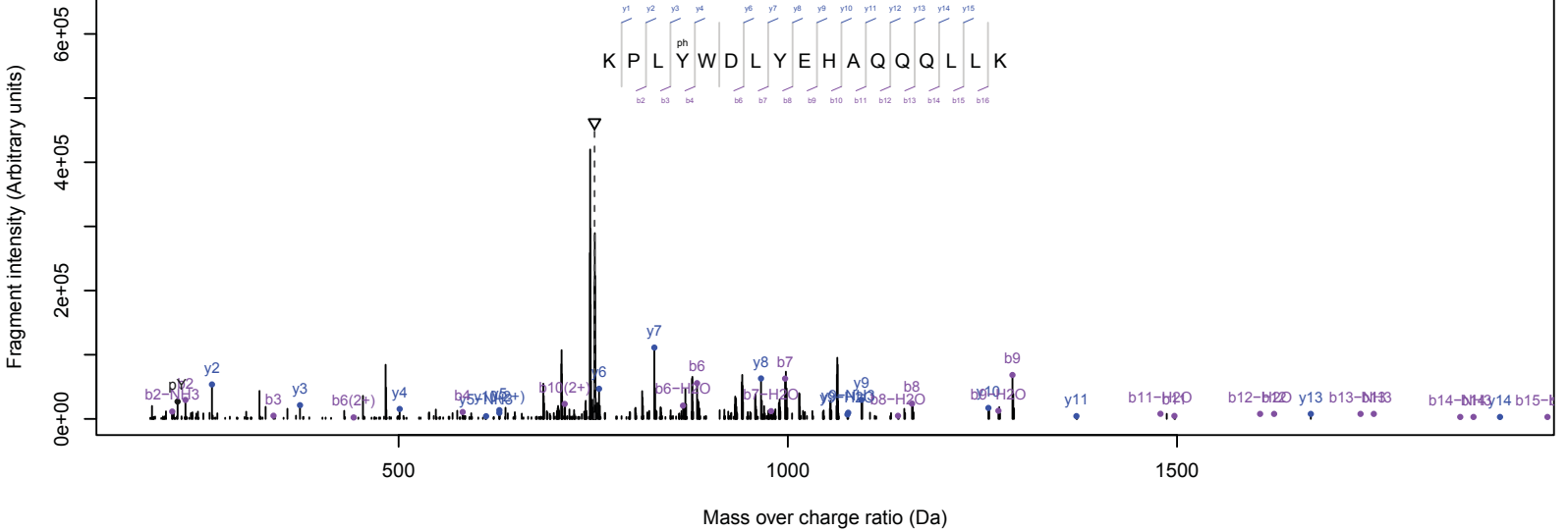
Andromeda score : 243.3



KPLY(ph)WDLYEHAQQQLLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39090

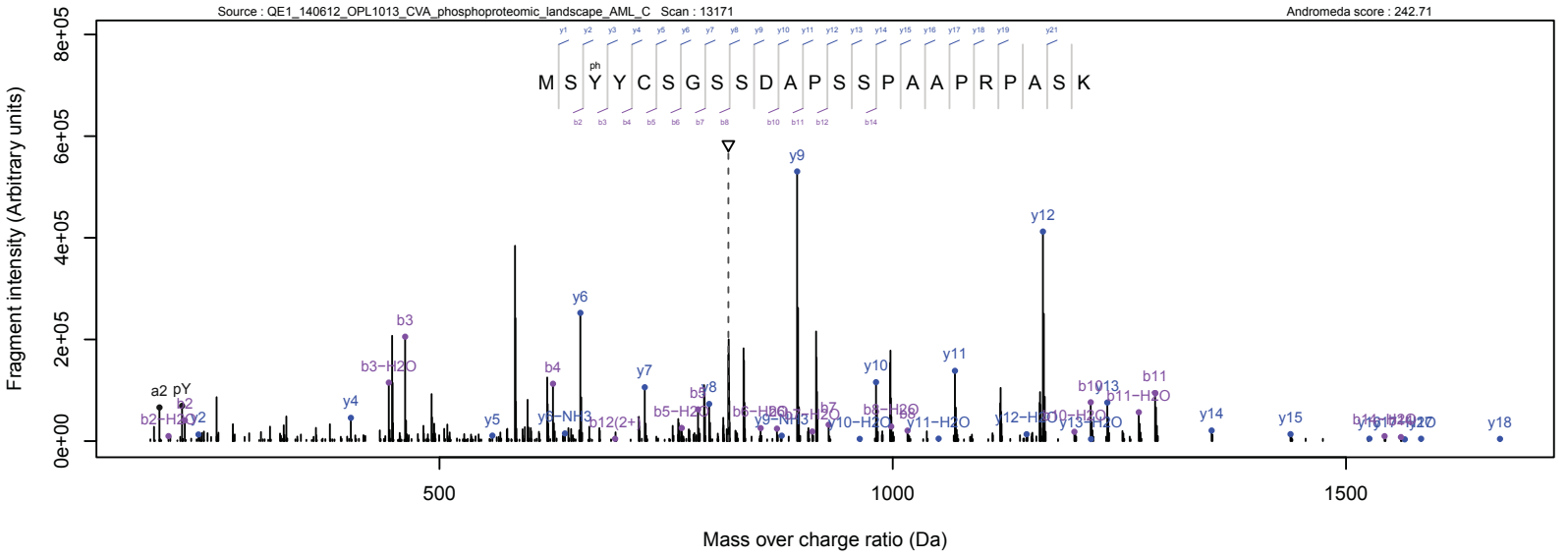
Andromeda score : 243.09



MSY(ph)YCSGSSDAPSSPAAPRPASK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13171

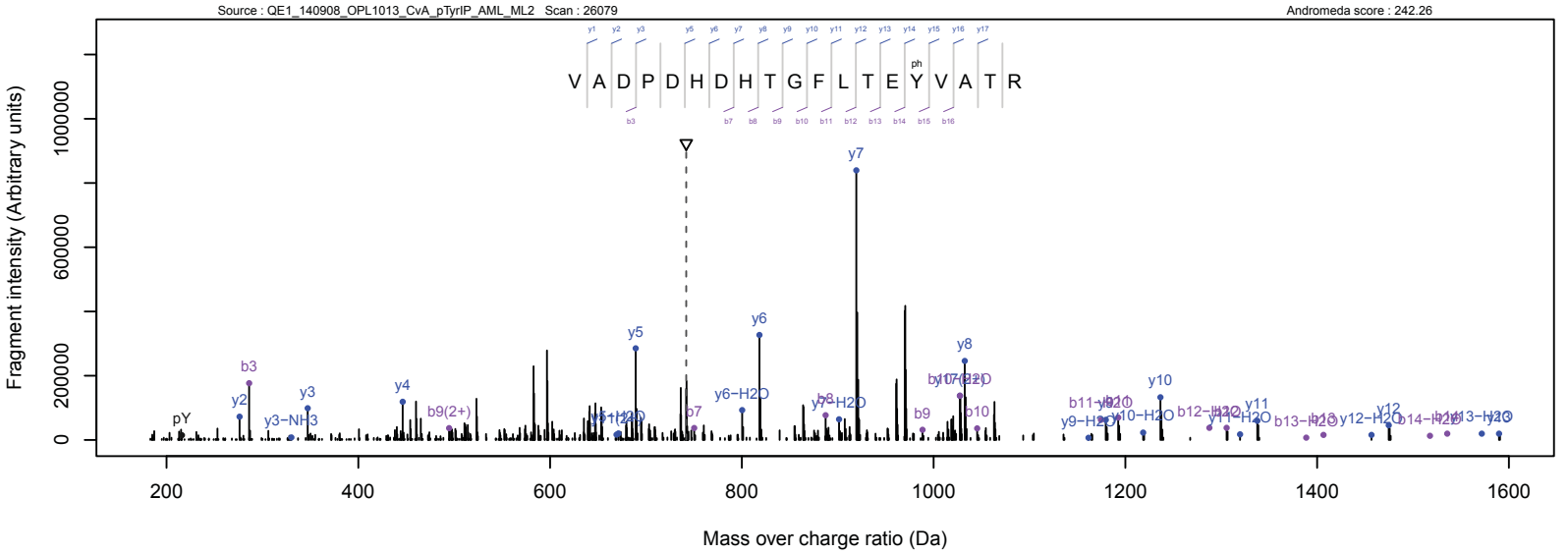
Andromeda score : 242.71



VADPDHDHTGFLTEY(ph)VATR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 26079

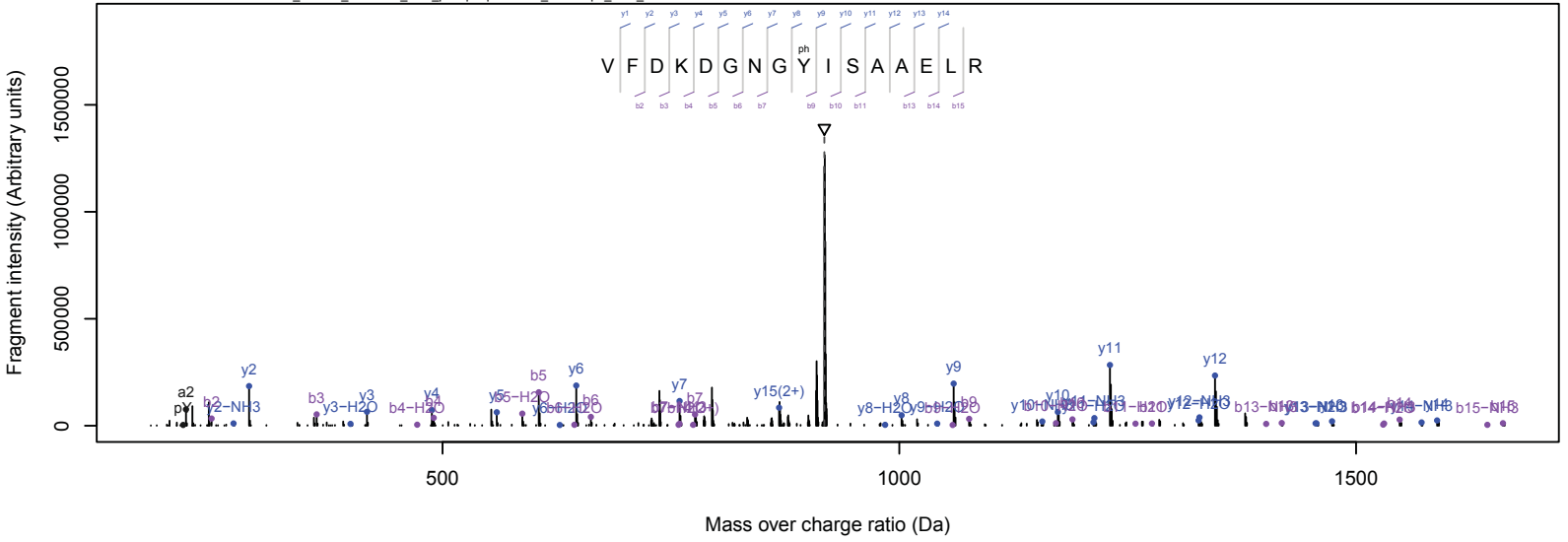
Andromeda score : 242.26



VFDKDGNGY(ph)ISAAELR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25391

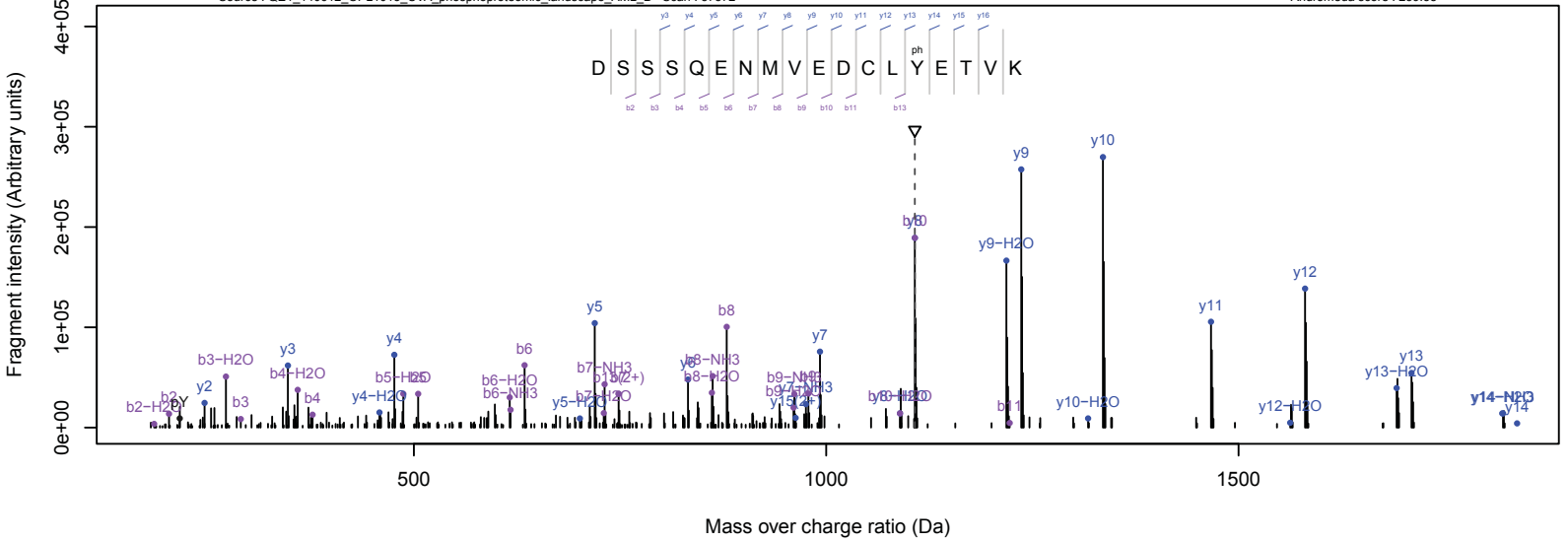
Andromeda score : 240.78



DSSSQENMVEDCLY(ETVK)-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37572

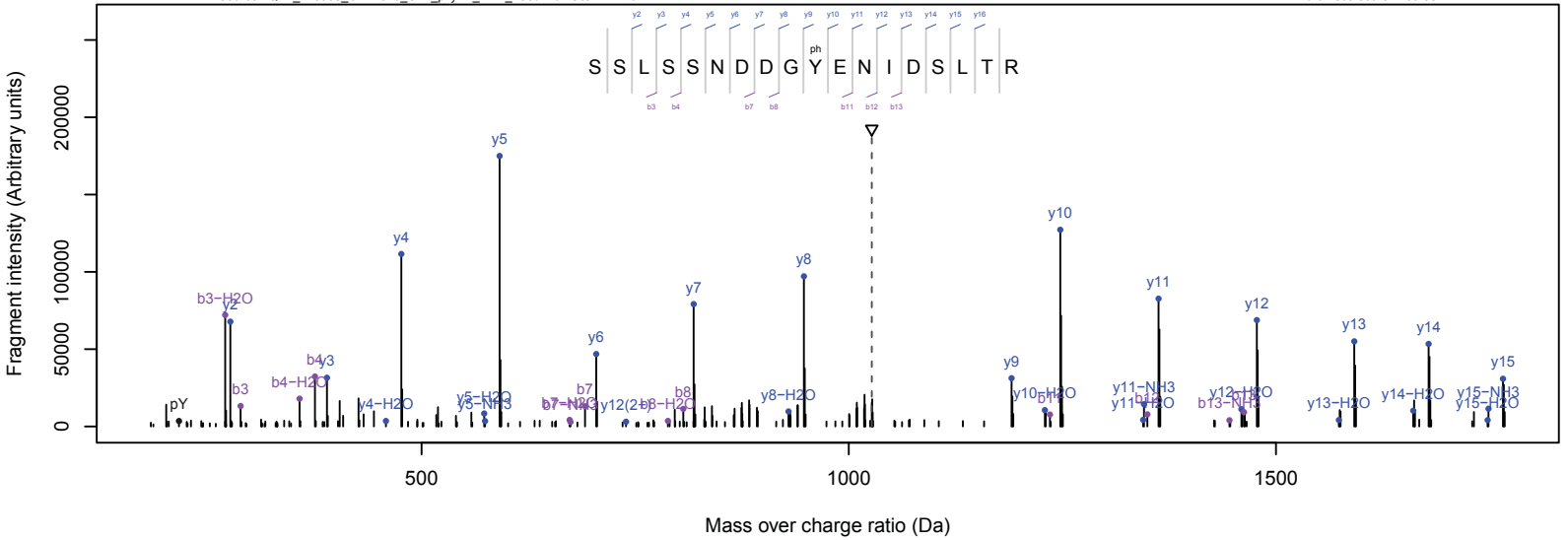
Andromeda score : 239.58

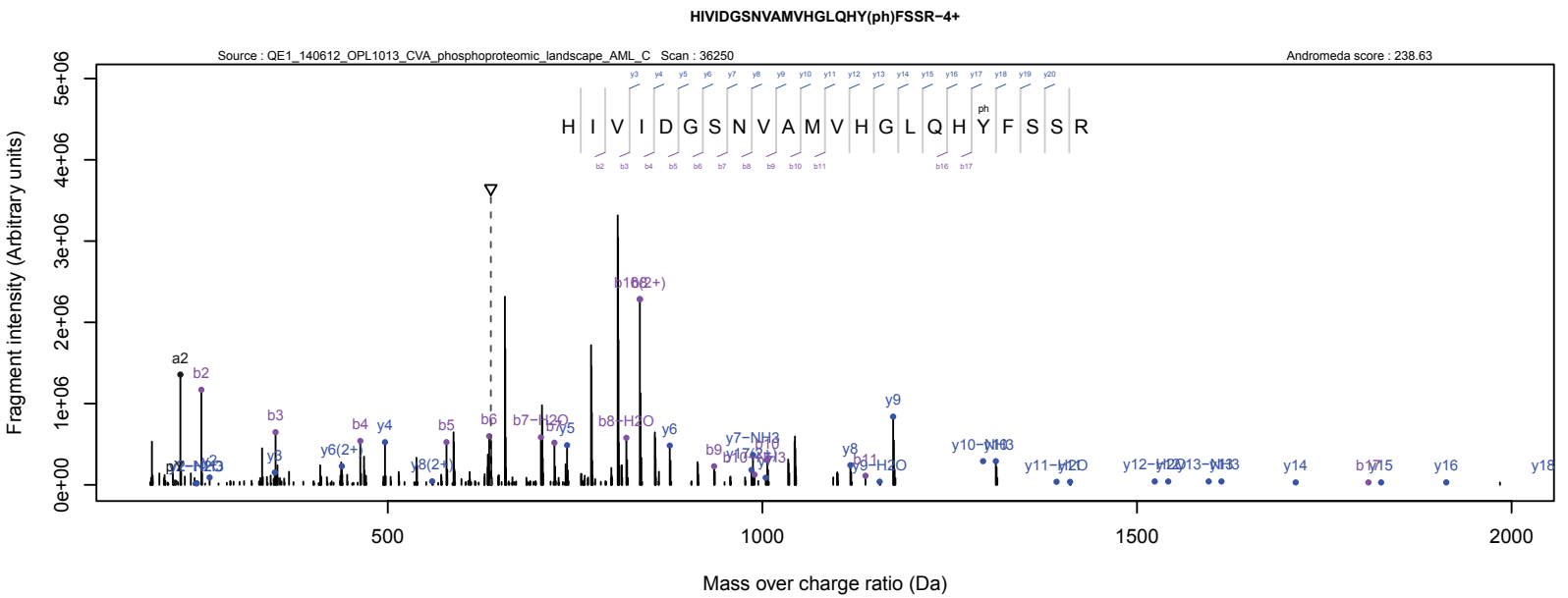
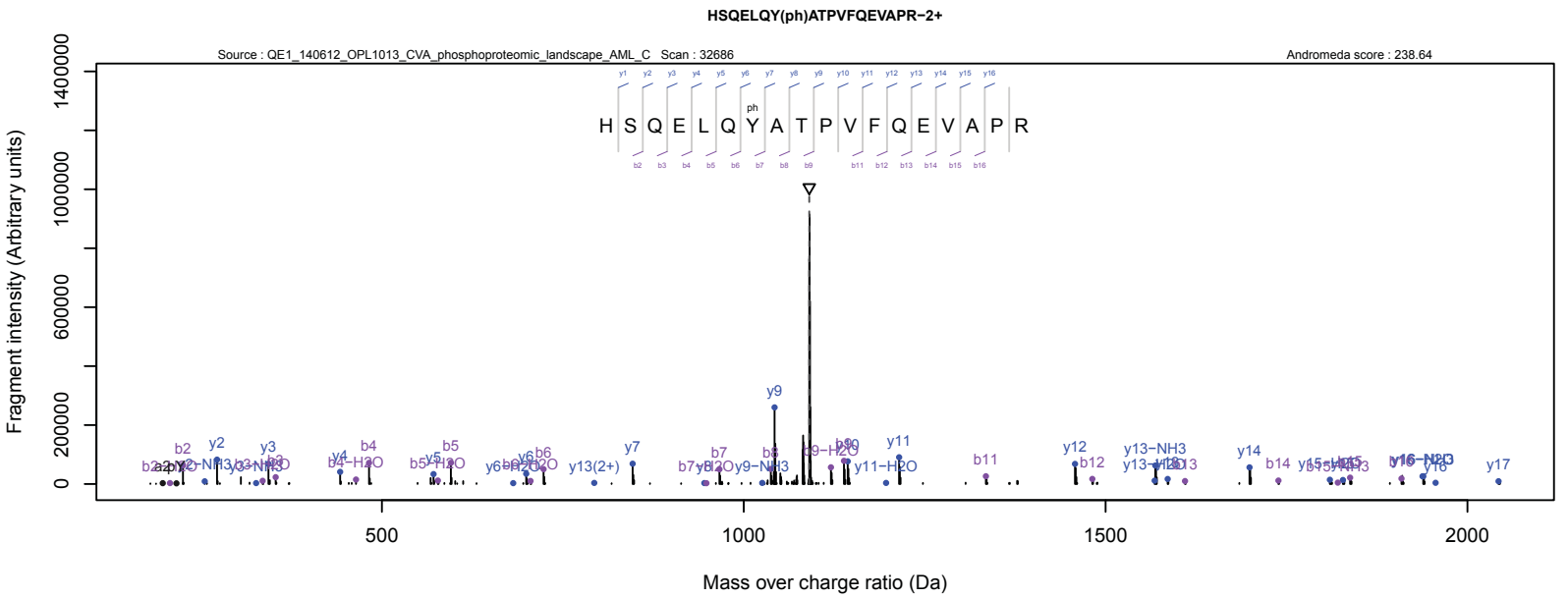
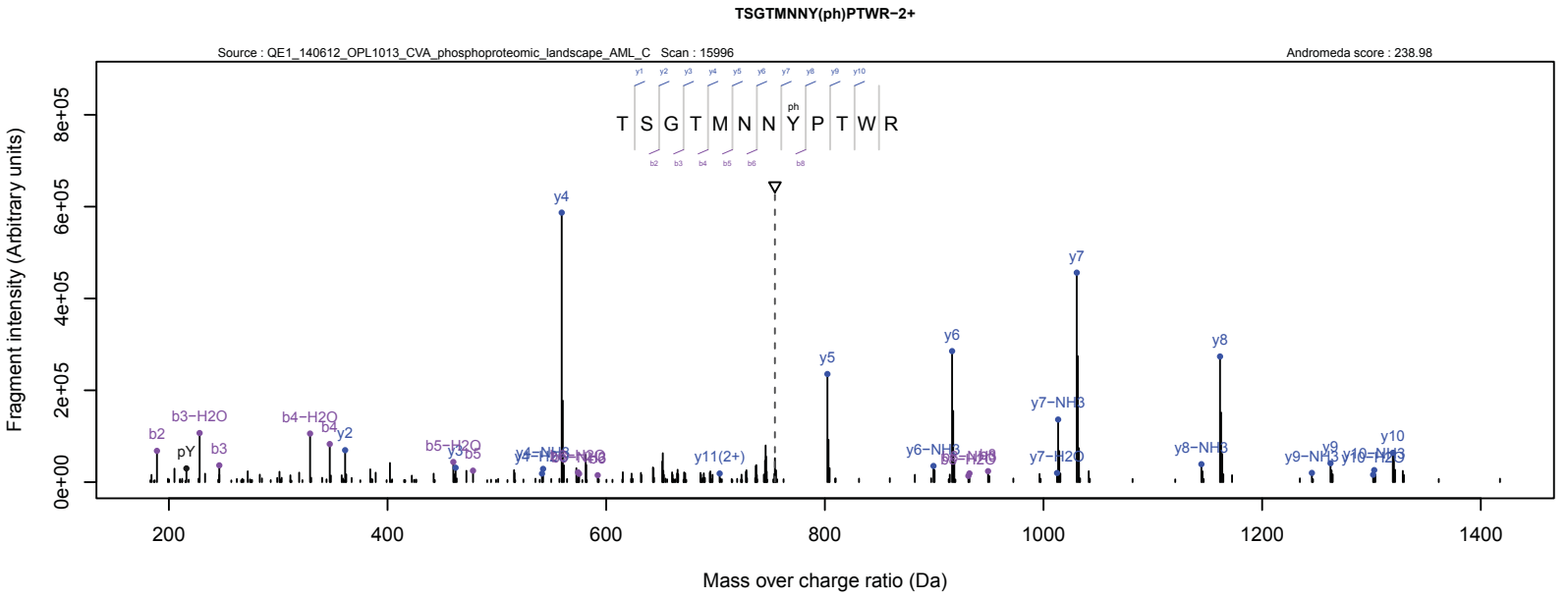


SSLSSNDGYP(h)ENIDSLTR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 27461

Andromeda score : 239.58



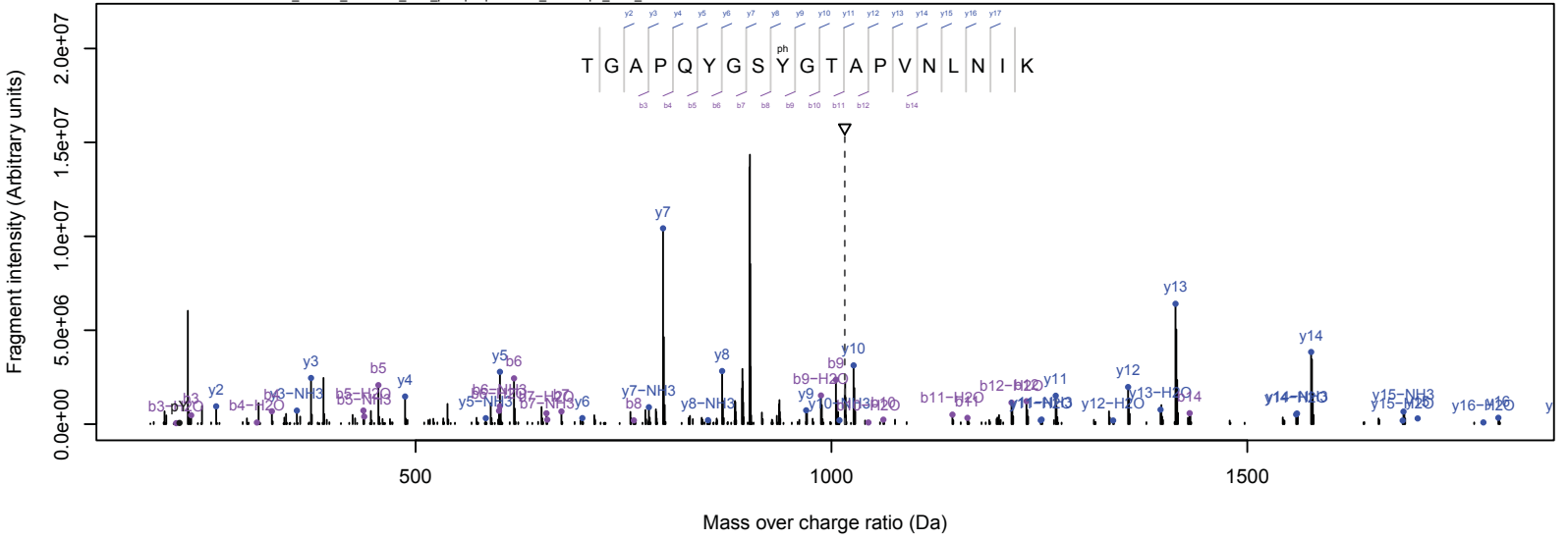




TGAPQYGSY(ph)GTAPVNLNIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29322

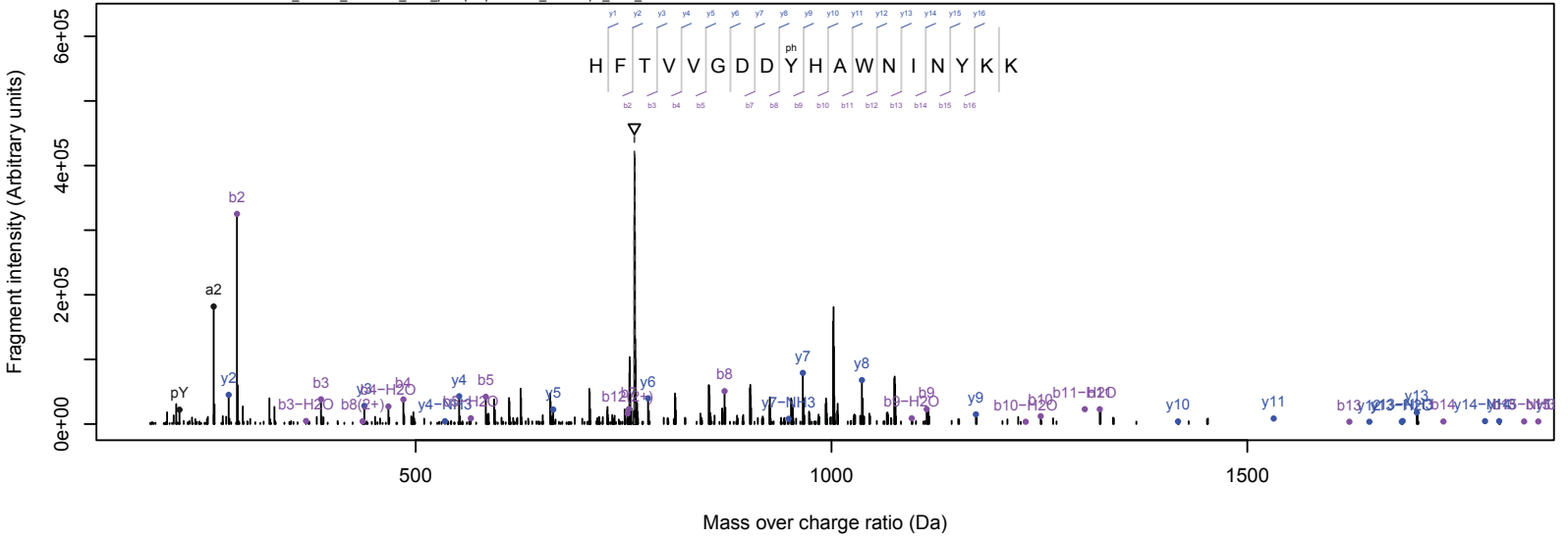
Andromeda score : 238.19



HFTVVGDDY(ph)HAWNINYKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27877

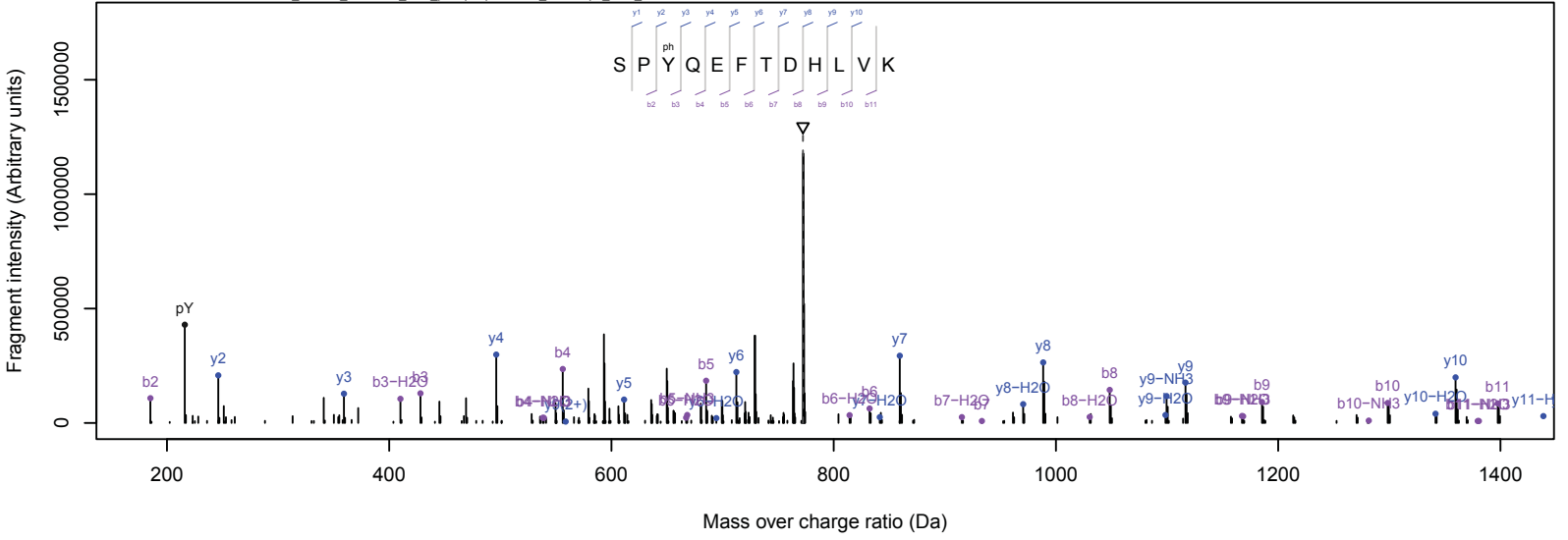
Andromeda score : 237.46

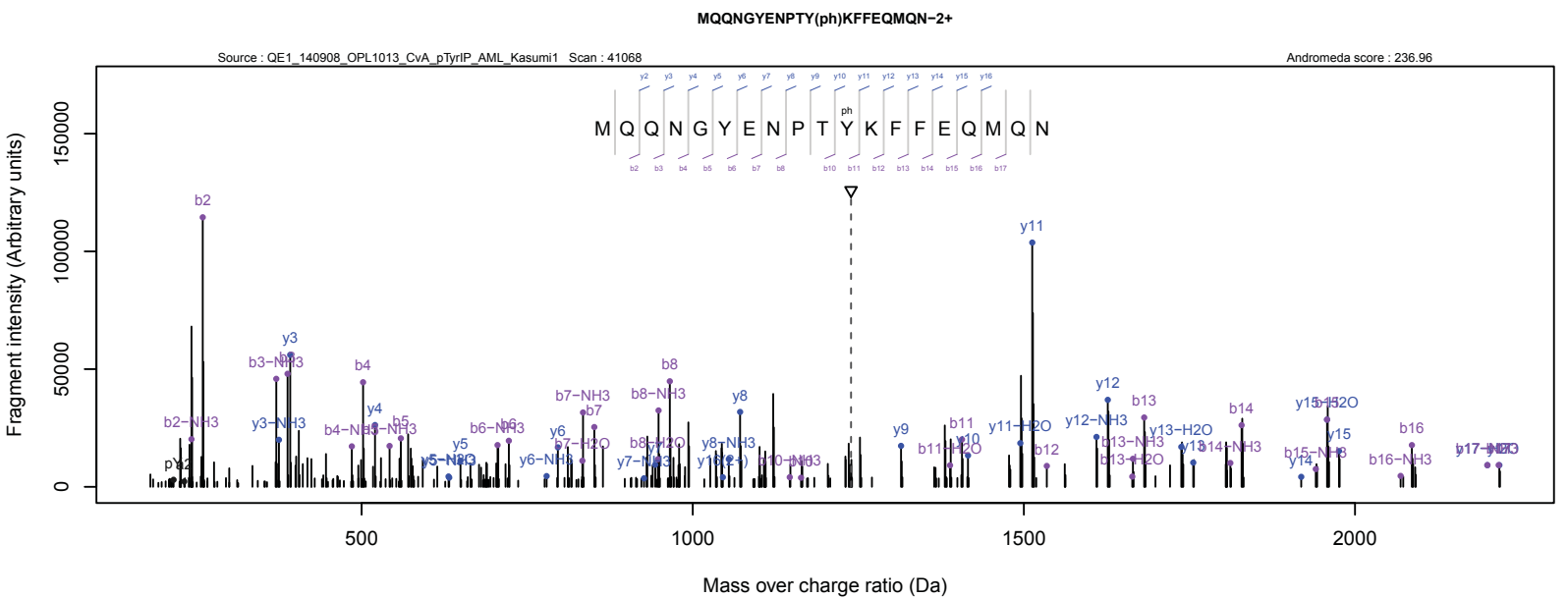
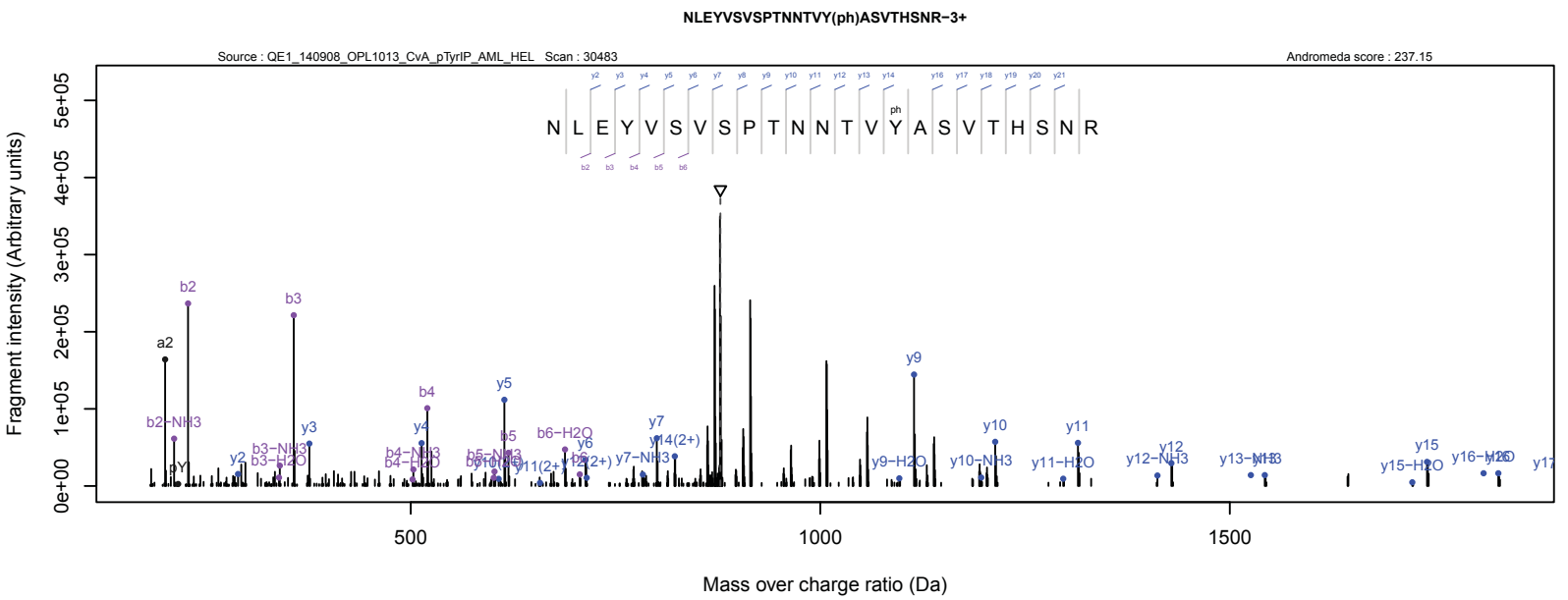
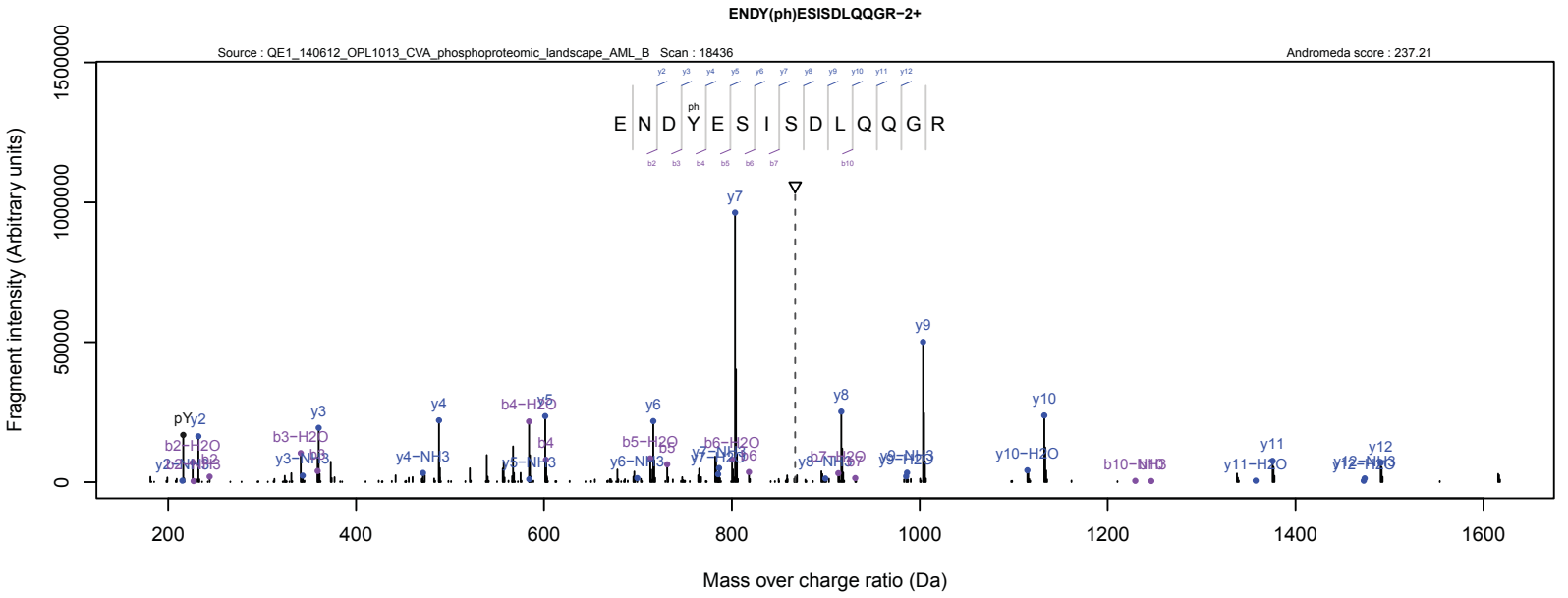


SPY(ph)QEFTDHLVK-2+

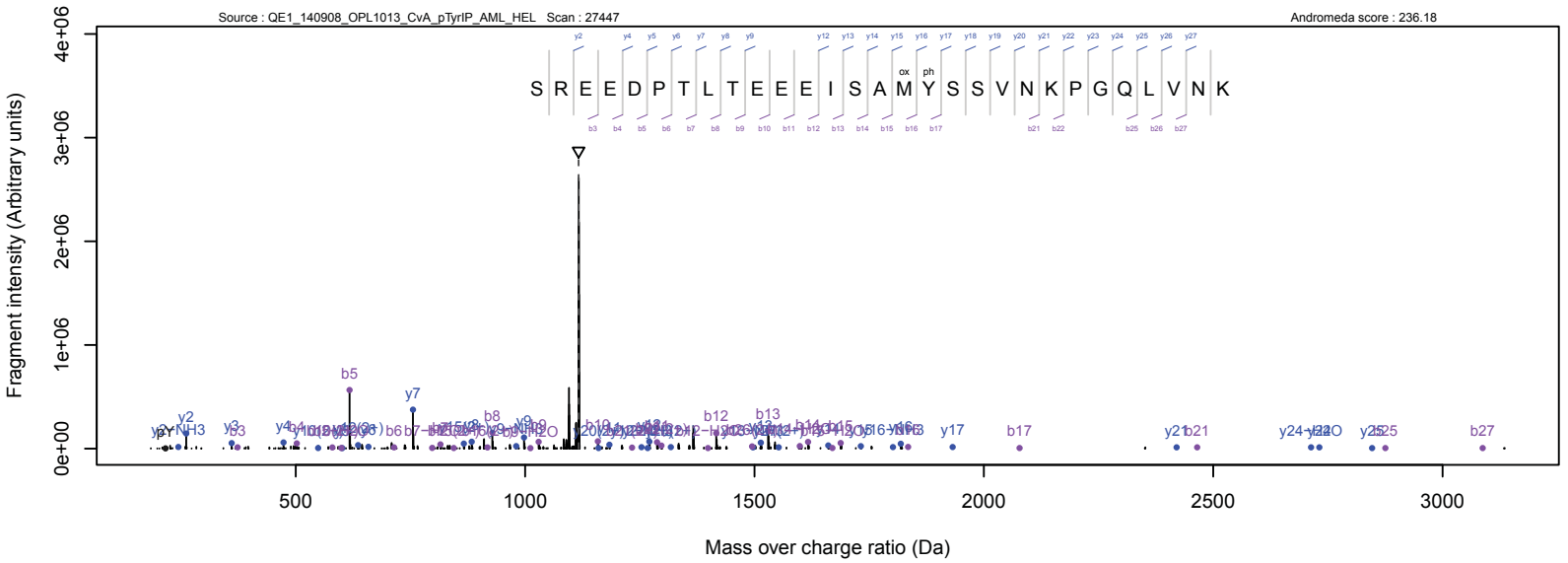
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25368

Andromeda score : 237.41

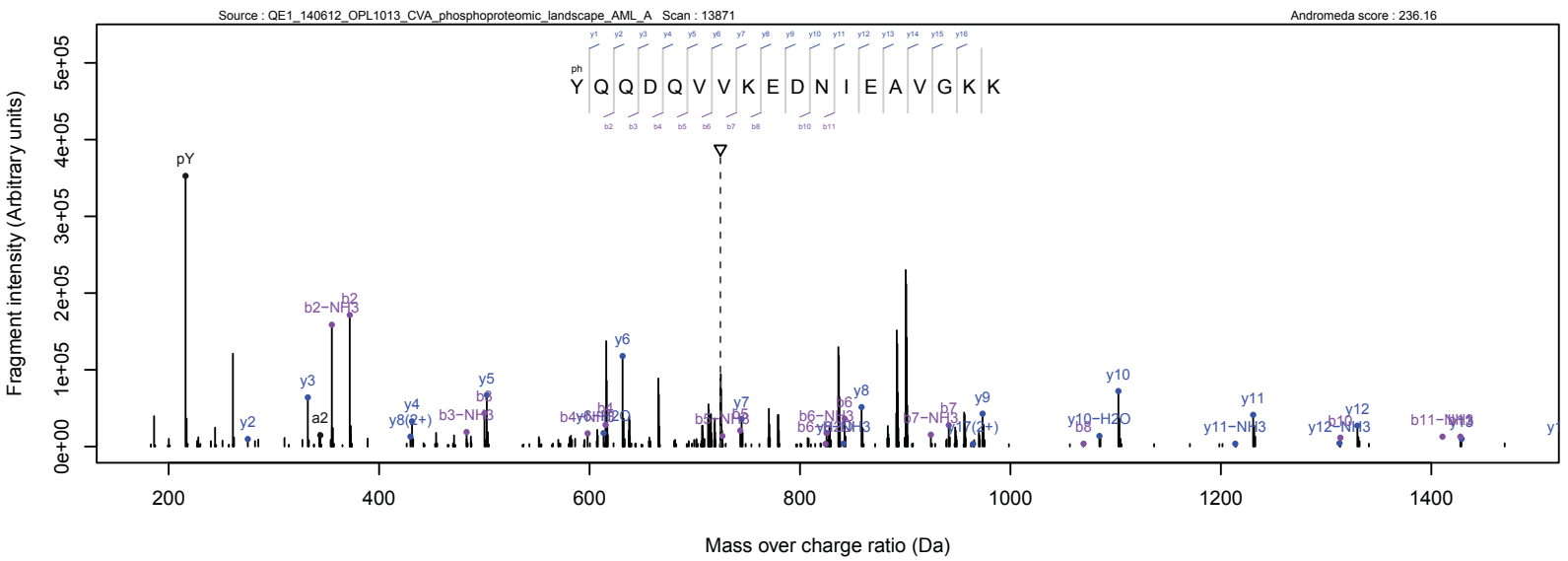




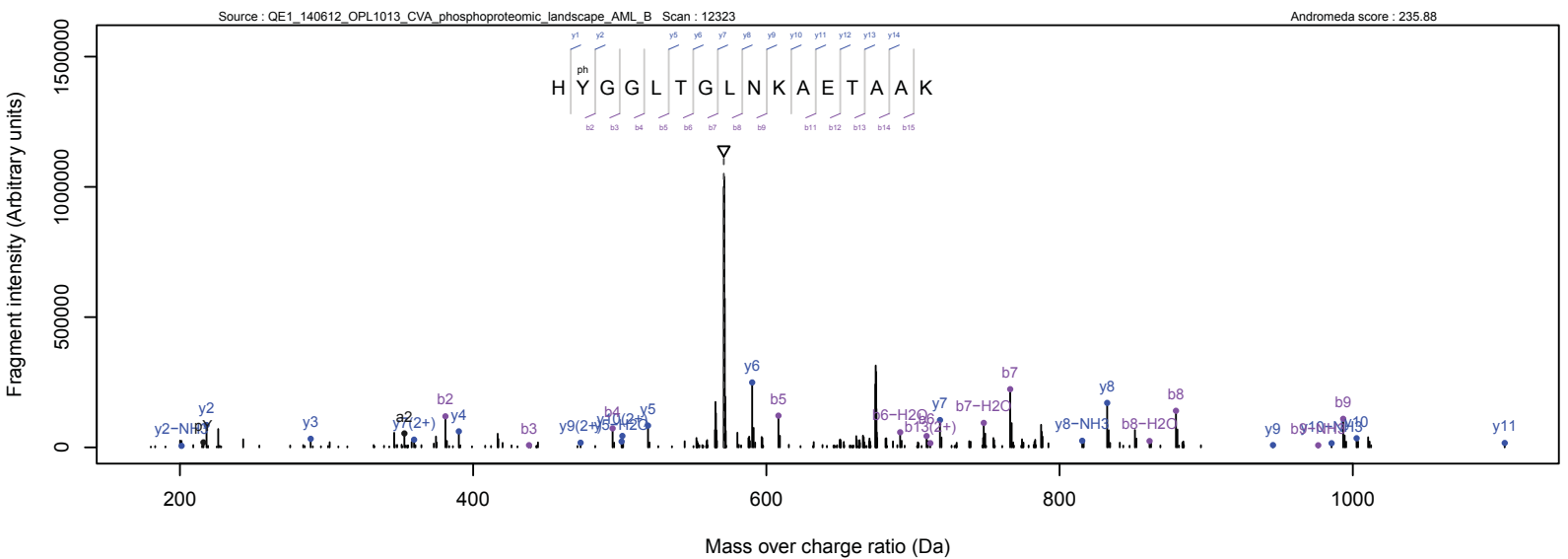
**SREEDPTLTEEEISAM(ox)Y(ph)SSVNKPGQLVNK-3+**



**Y(ph)QQDQVVKEDNIEAVGKK-3+**



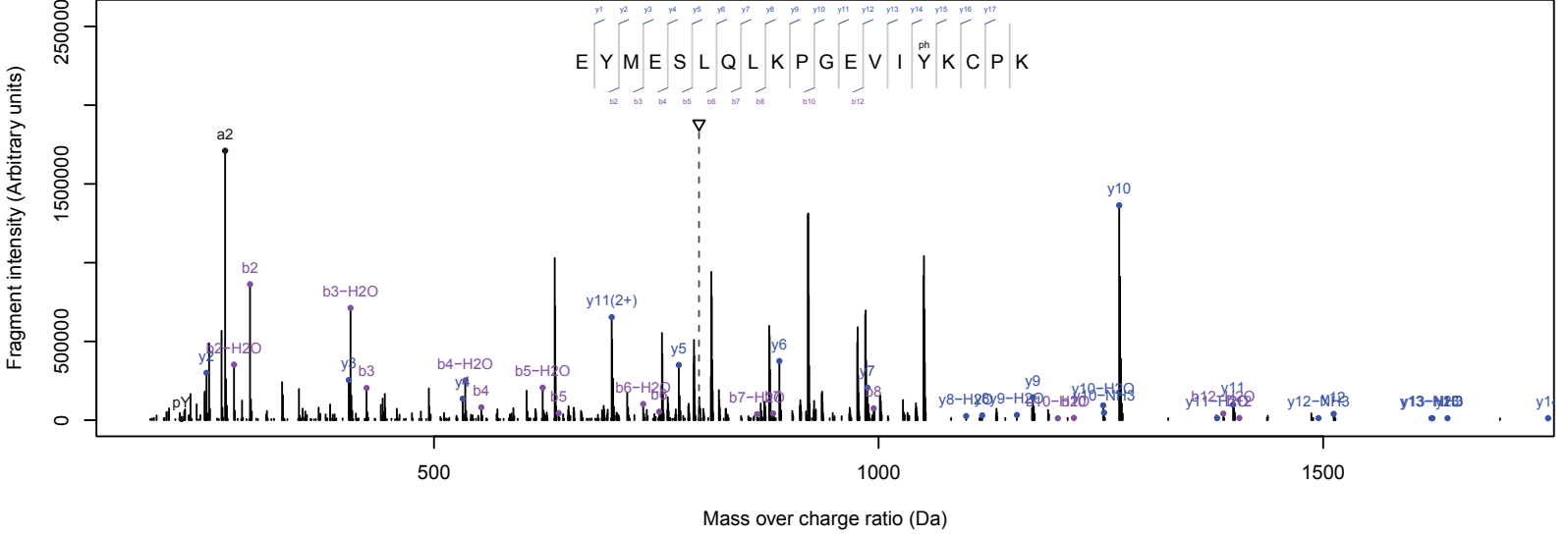
**HY(ph)GGLTGLNKAETAAK-3+**



EYMESLQLKPGEVY(ph)KCPK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 28954

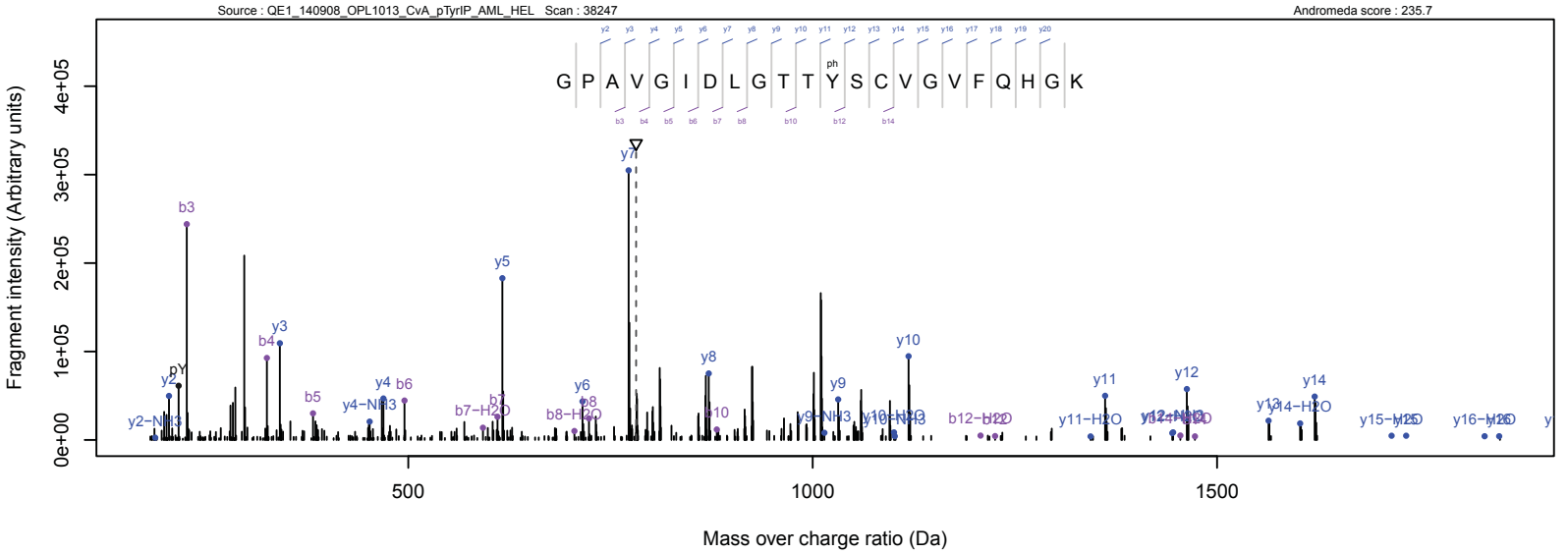
Andromeda score: 235.75



GPAVGIDLTTY(ph)SCVGVFQHGK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 38247

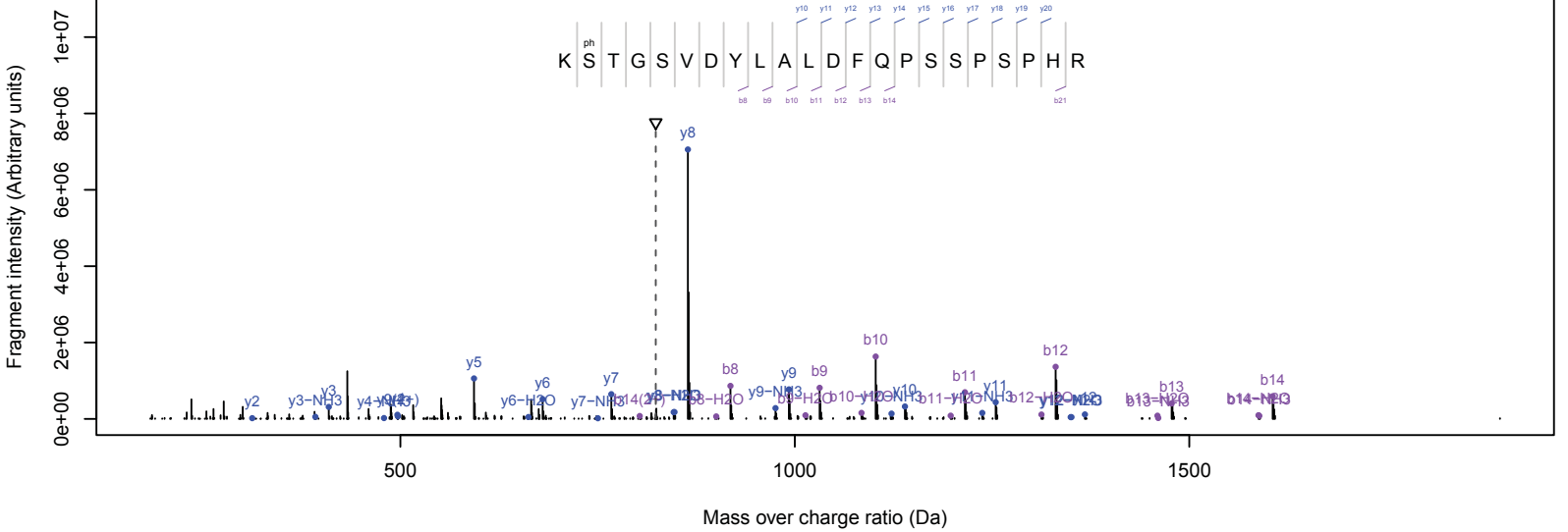
Andromeda score: 235.7



KS(ph)TGSVDYLALDFQPSSPSPHR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 30727

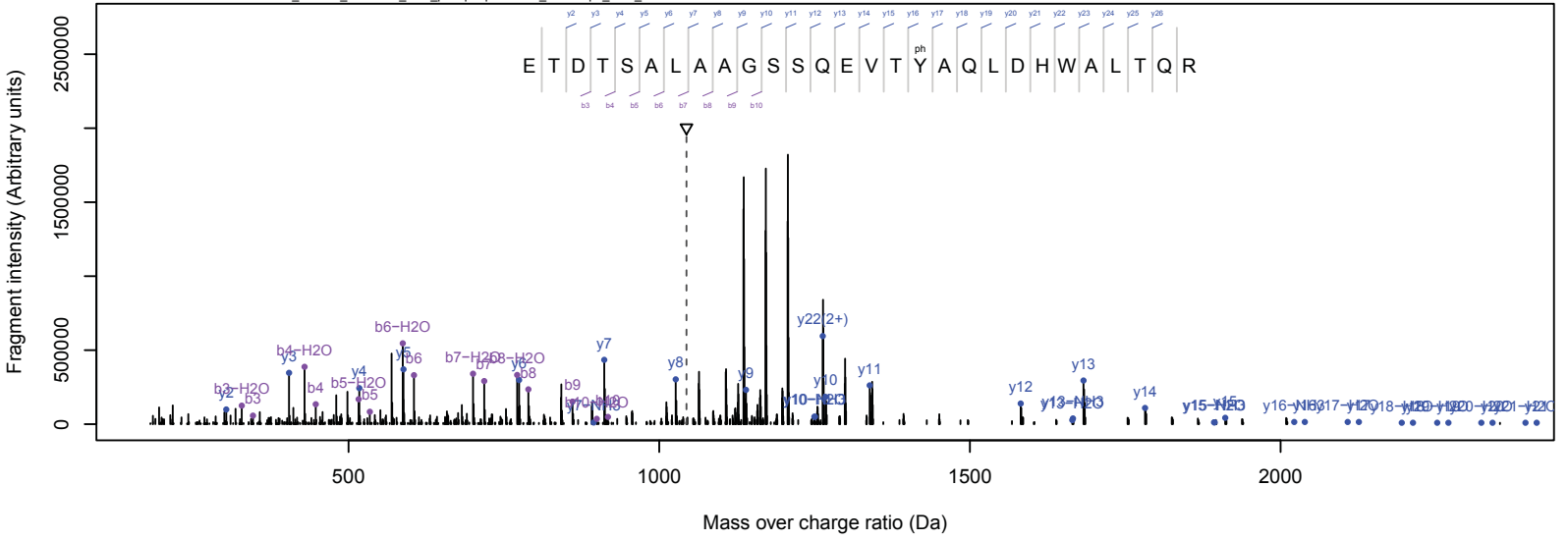
Andromeda score: 235.7



ETDTSALAAGSSQEVTY(ph)AQLDHWALTQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39156

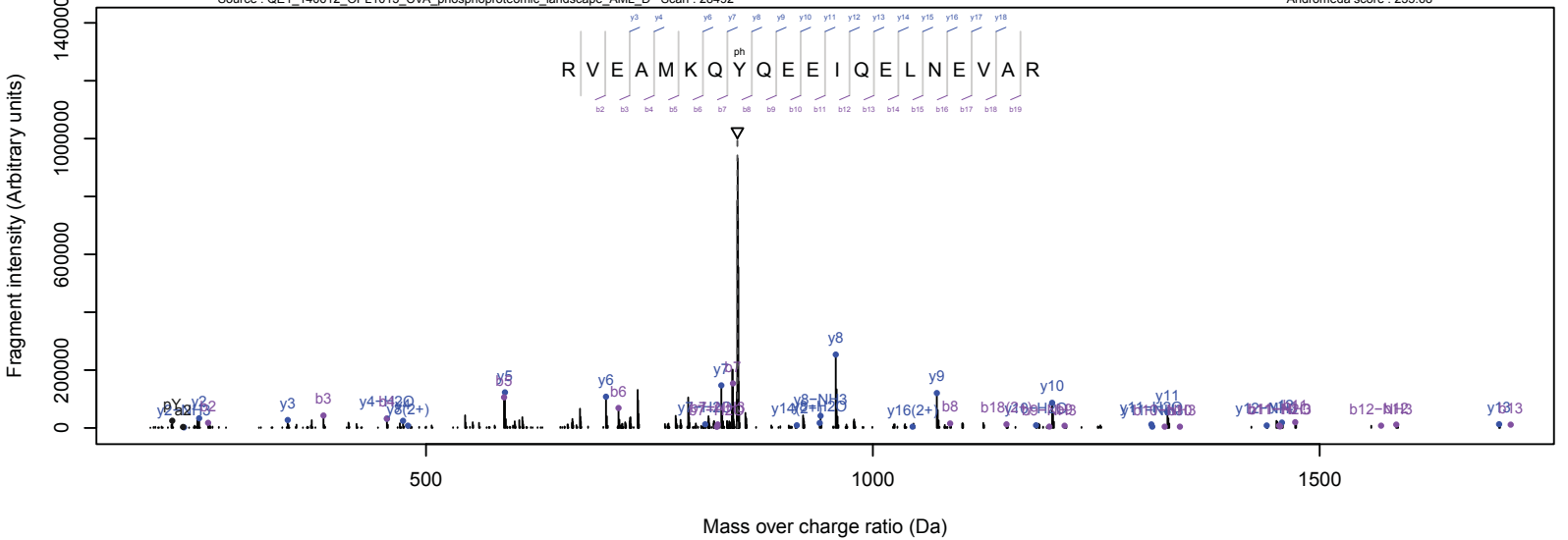
Andromeda score : 235.69



RVEAMKQY(ph)QEEIQELNEVAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28492

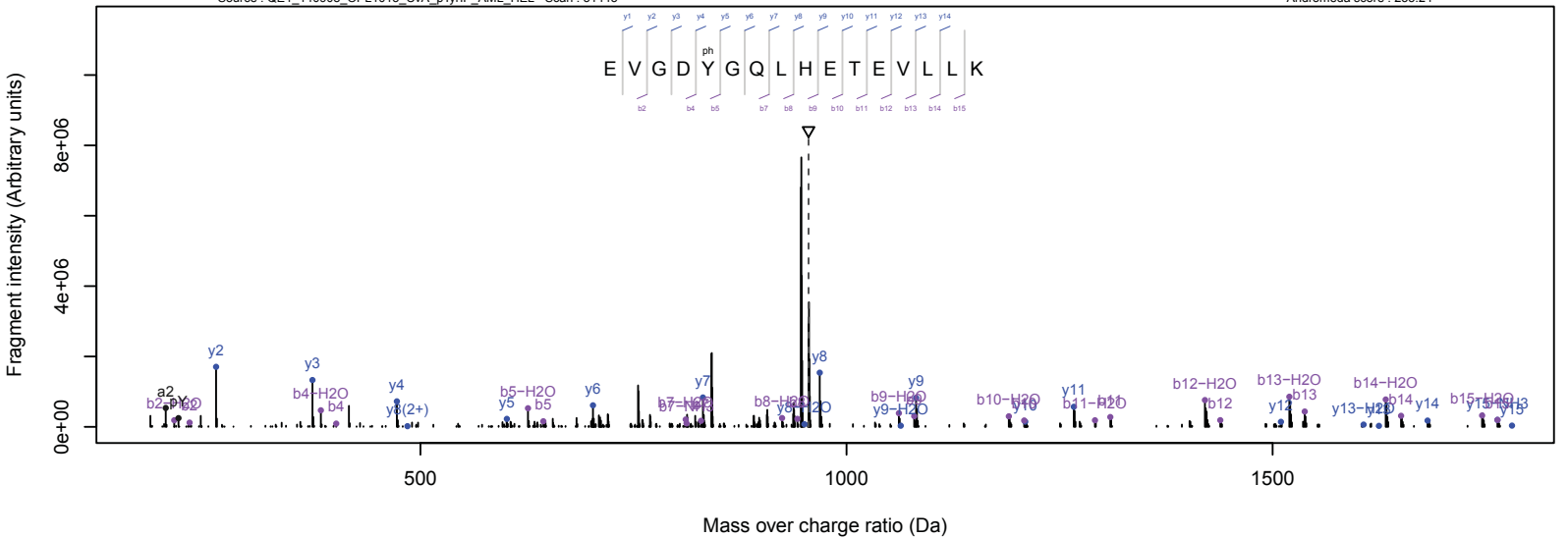
Andromeda score : 235.68



EVGDY(ph)GQLHETEVLLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31443

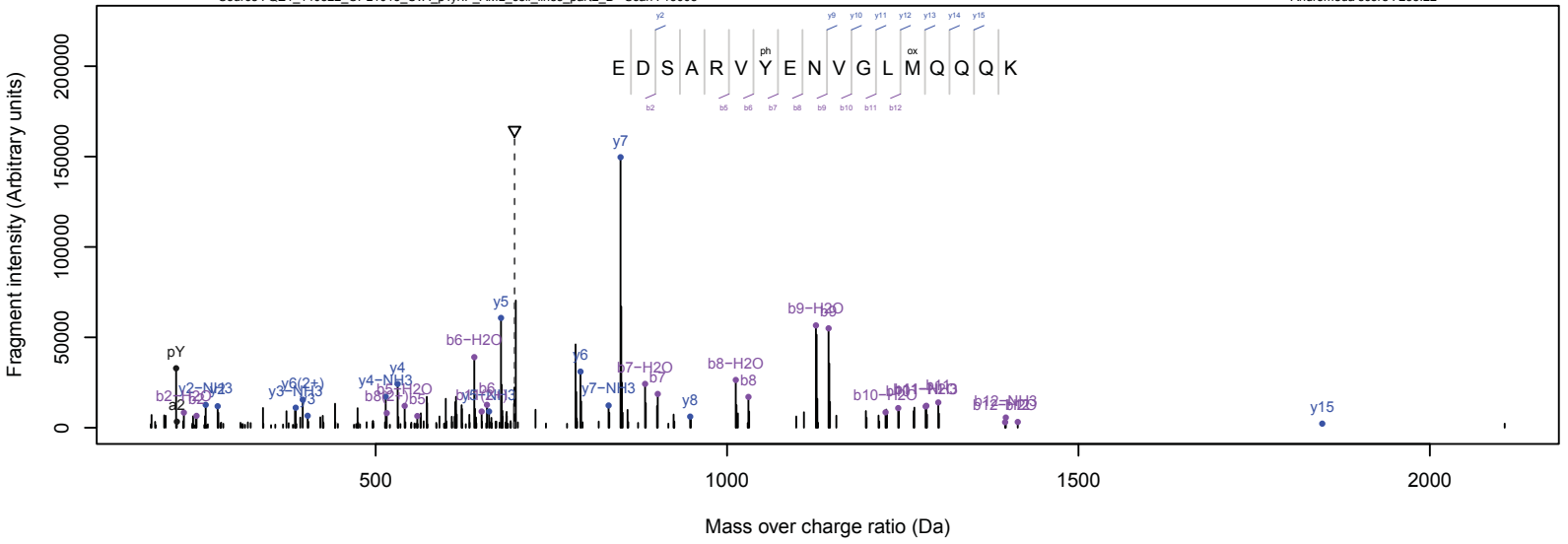
Andromeda score : 235.24



EDSARVY(ph)ENVGLM(ox)QQQK-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 13008

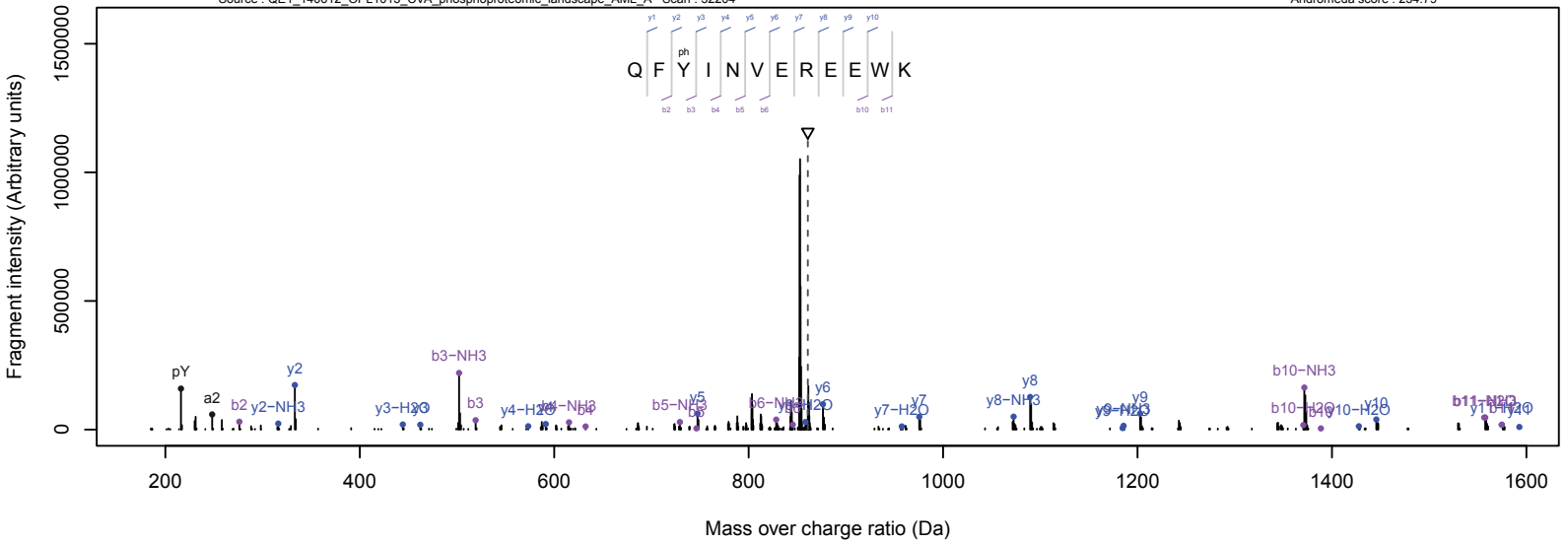
Andromeda score : 235.22



QFY(ph)INVEREEWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32204

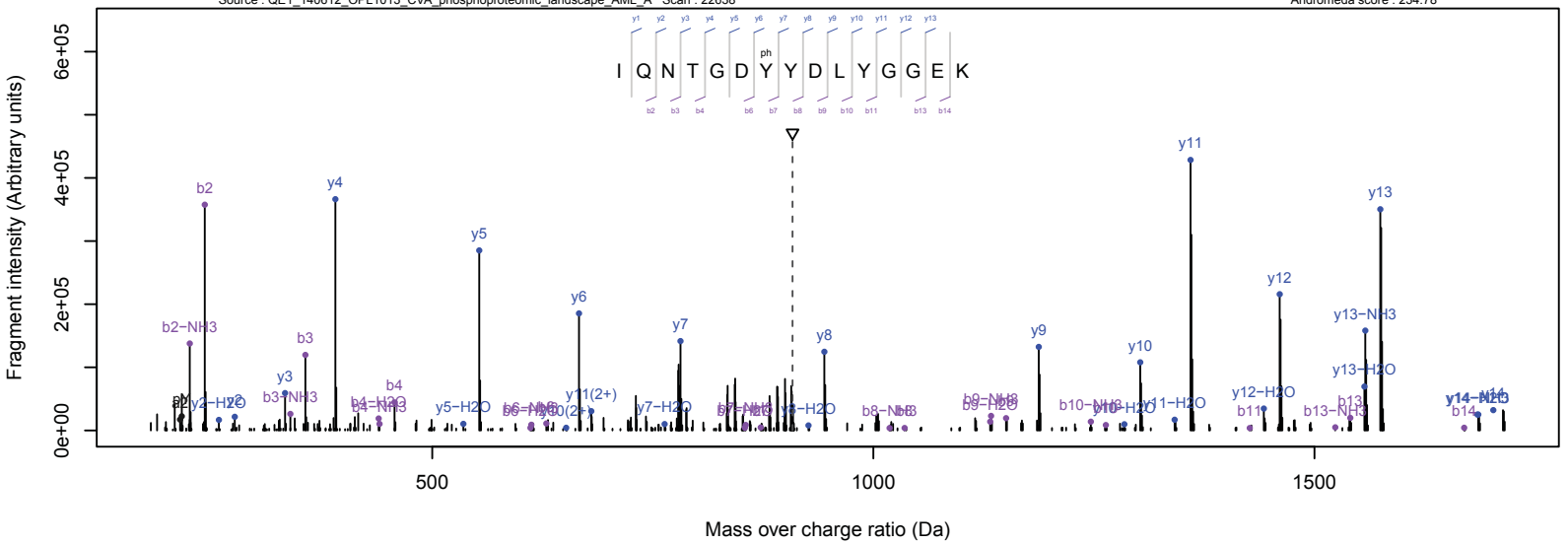
Andromeda score : 234.79



IQNTGDY(ph)YDLYGGEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 22638

Andromeda score : 234.78

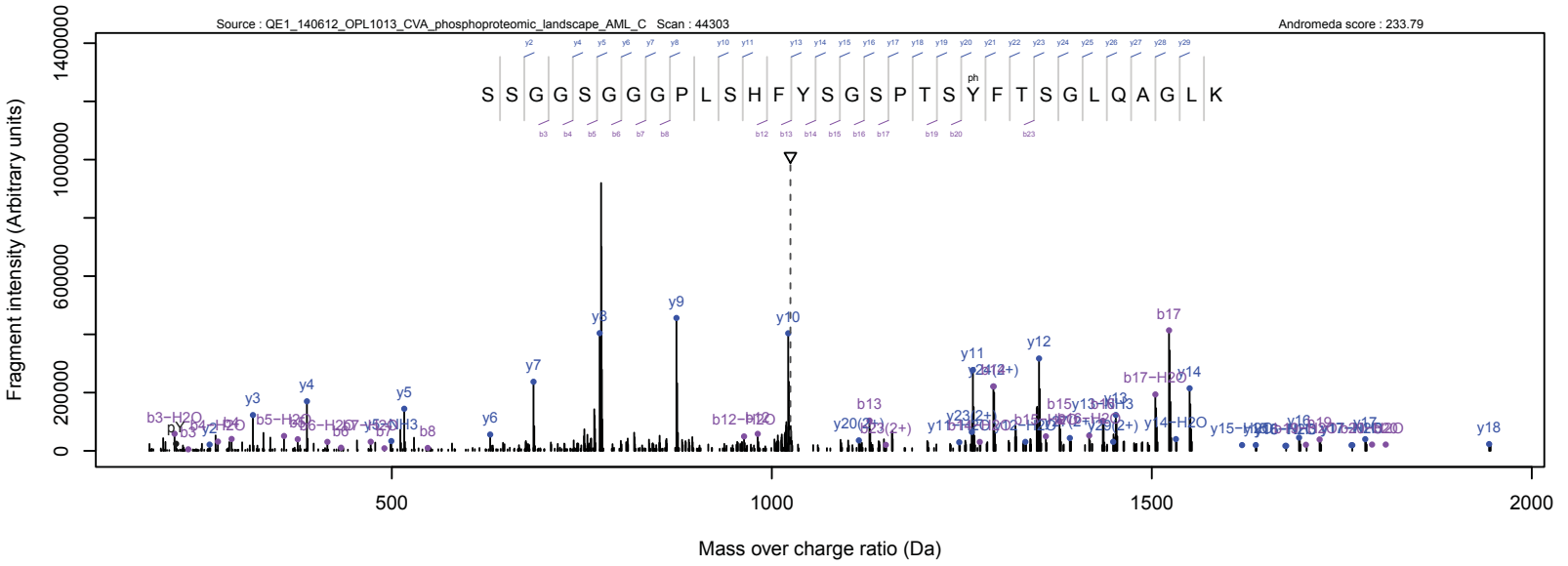




SSGGSGGPLSHFYSGSPTSY(ph)FTSGLQAGLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44303

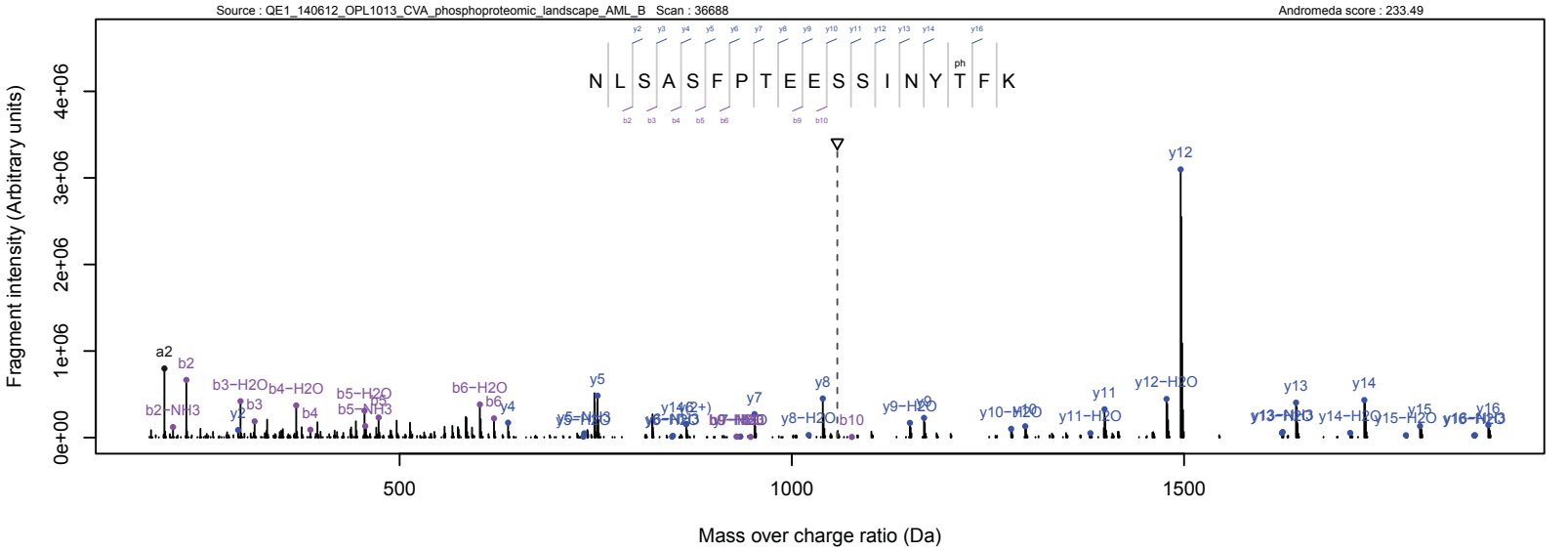
Andromeda score : 233.79



NLSASFTEESSINYT(ph)FK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36688

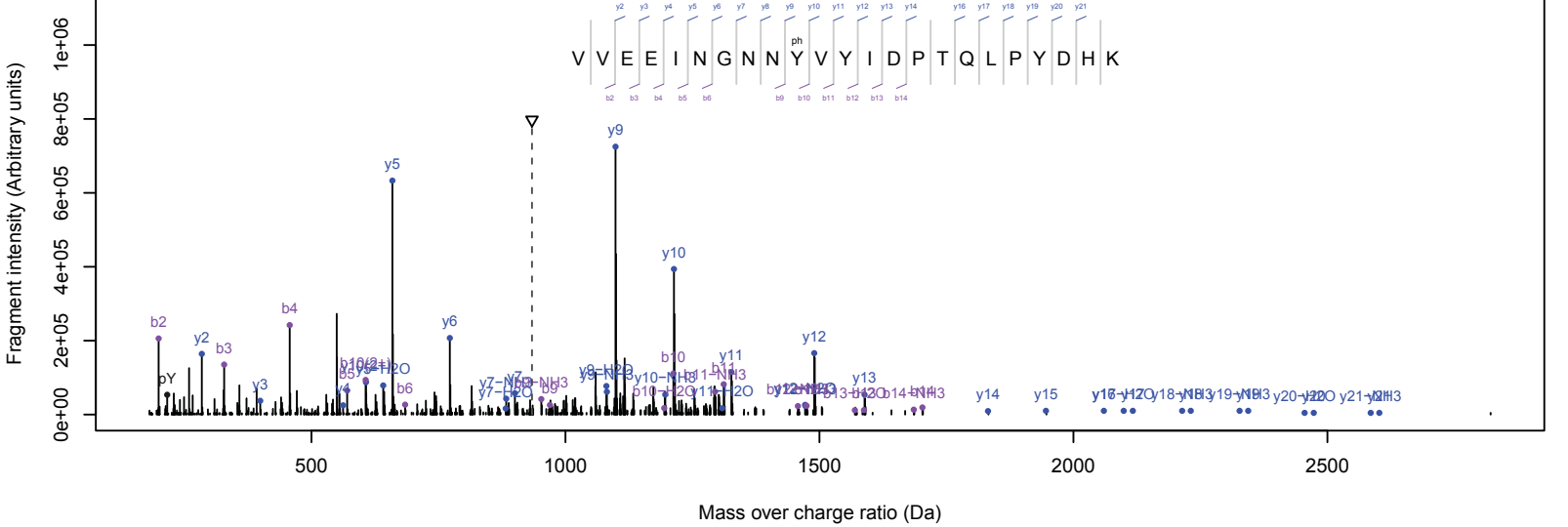
Andromeda score : 233.49



VVEEINGNYYVVIDPTQLPYDHK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 36510

Andromeda score : 233.26



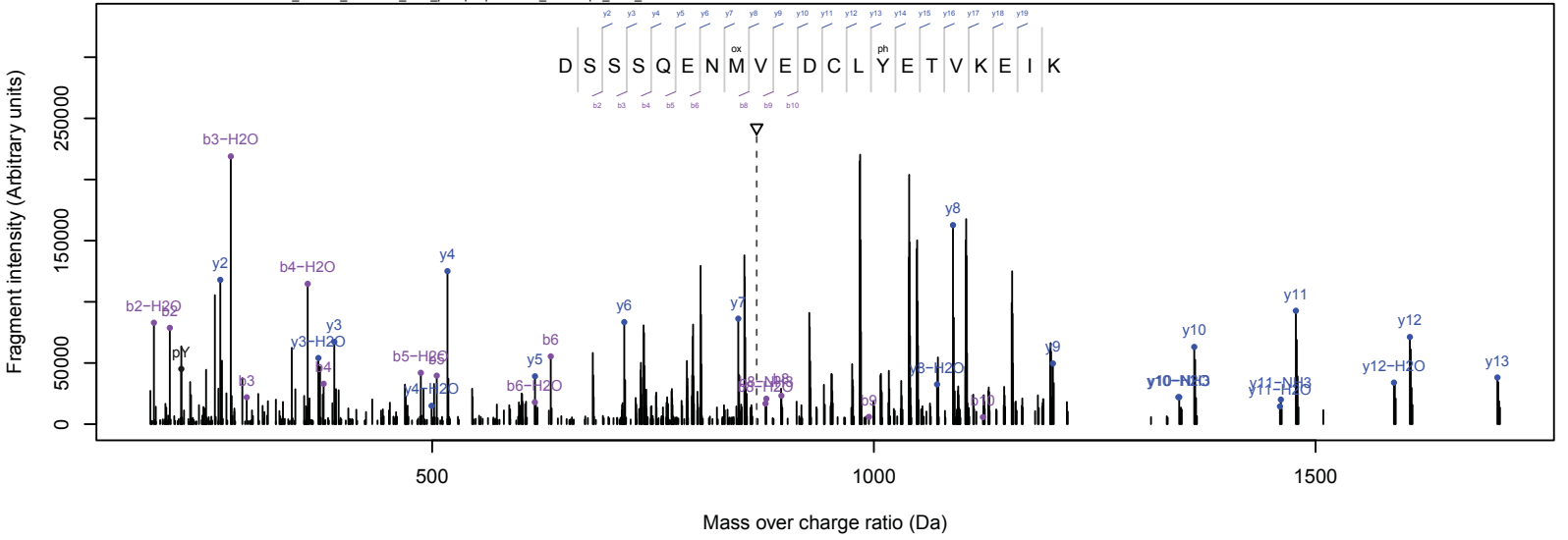




DSSSQENM(ox)VEDCLY(ph)ETVKEIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44914

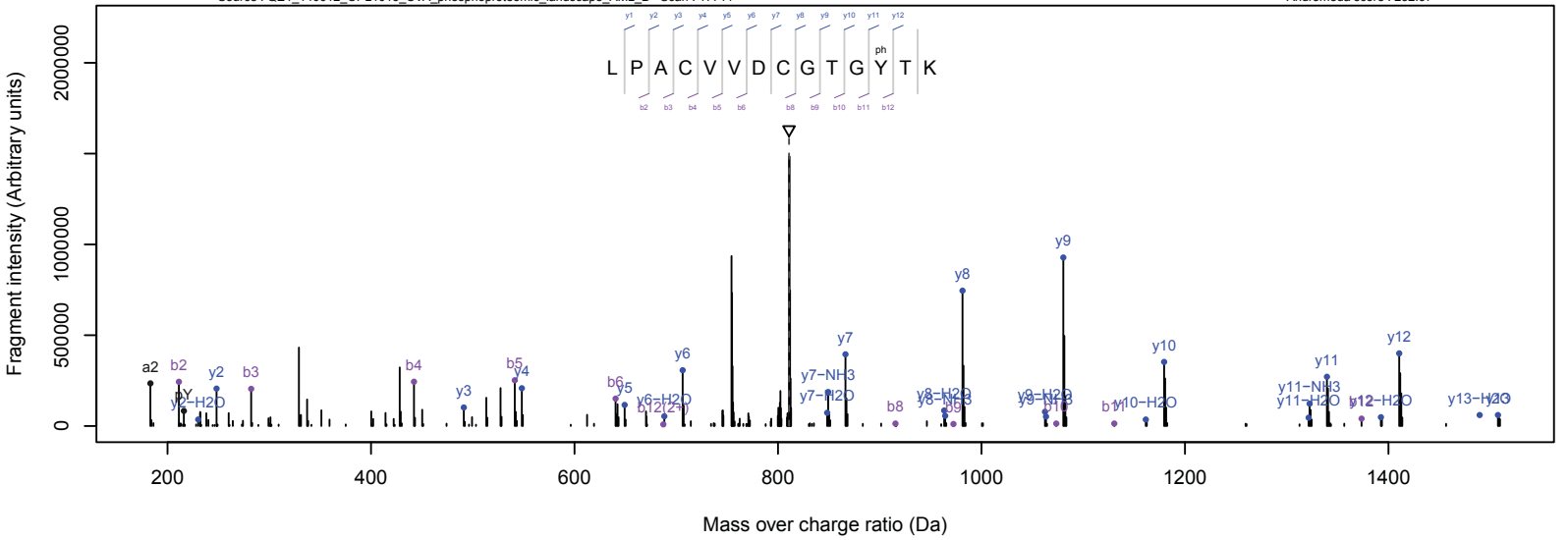
Andromeda score : 232.54



LPACVVDCTG(ph)TK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17711

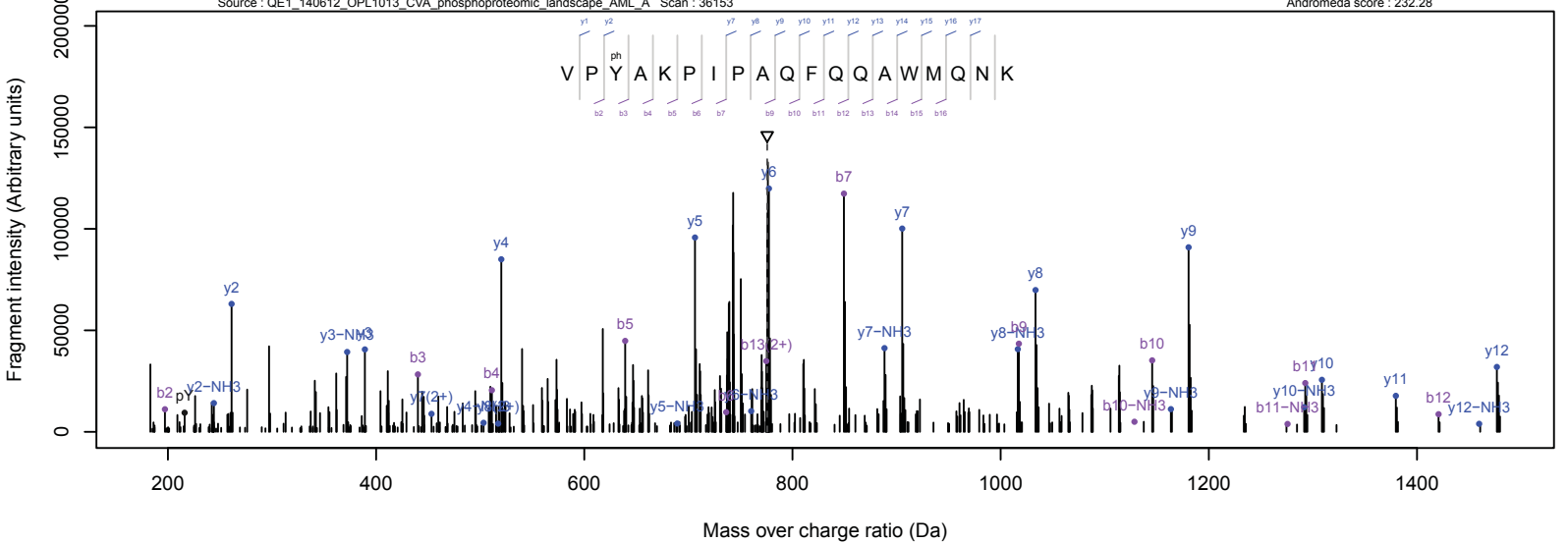
Andromeda score : 232.37



VPY(ph)AKPIPAQFQQAWMQNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36153

Andromeda score : 232.28

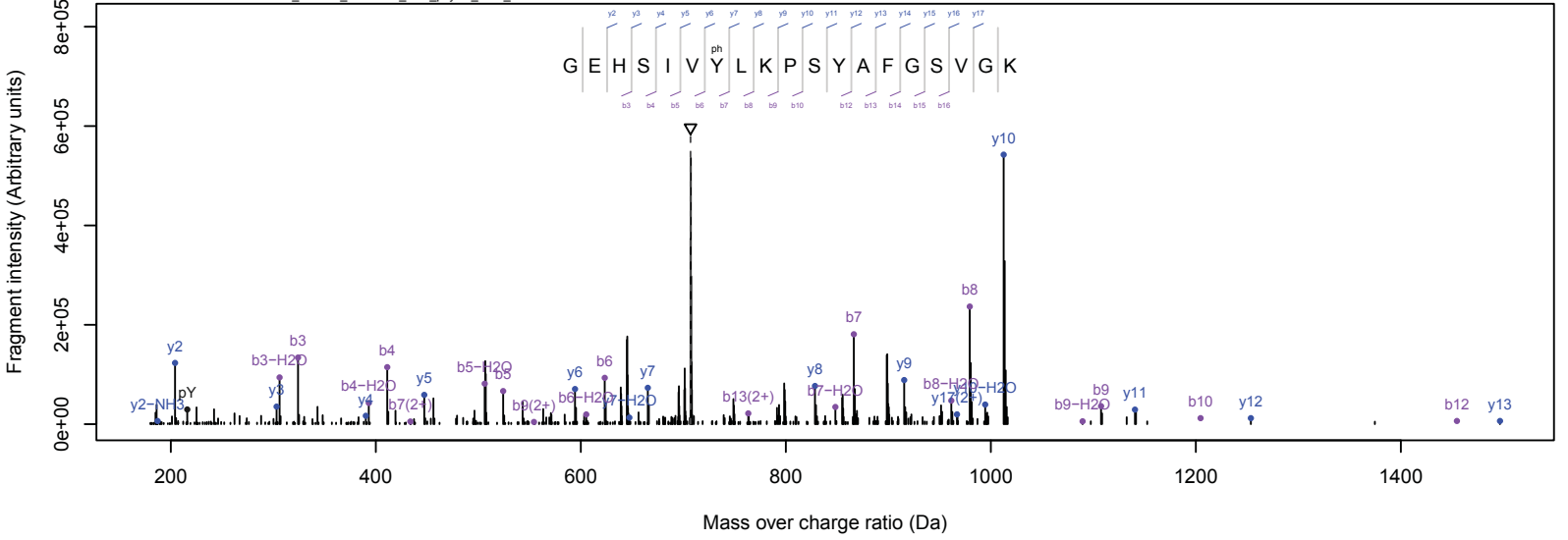




GEHSIVY(ph)LKPSYAFGSVGK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 32311

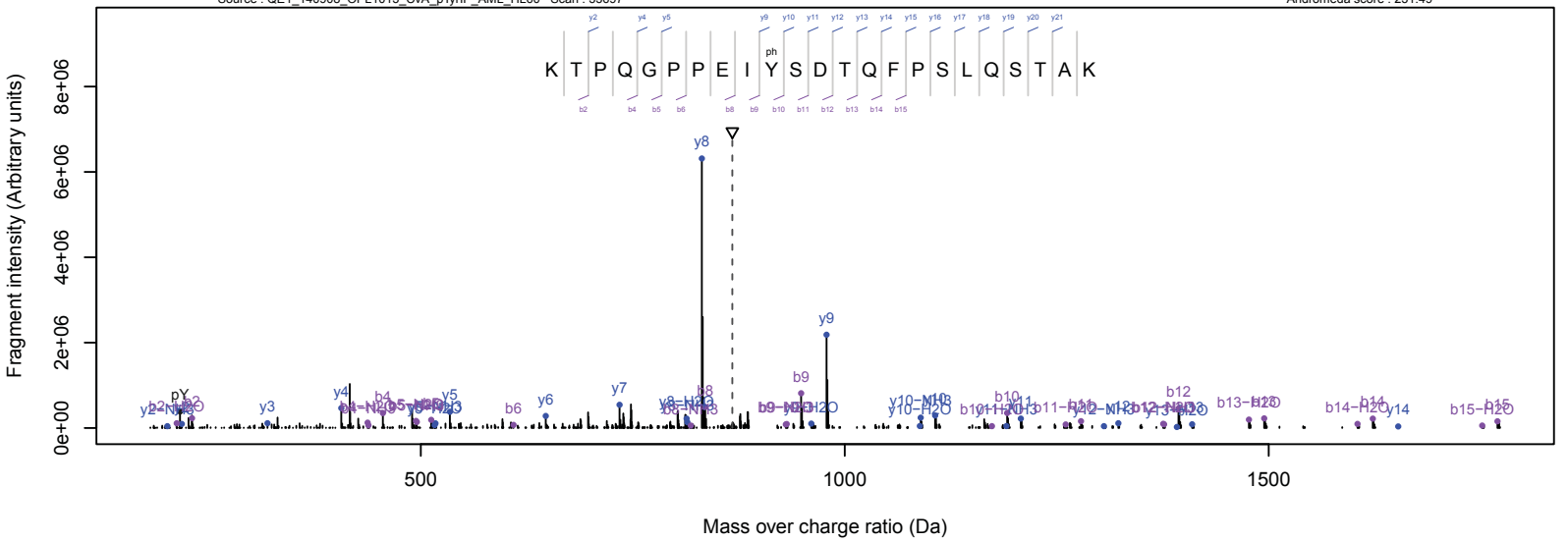
Andromeda score : 231.77



KTPQGPEIY(ph)SDTQFPSLQSTAK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 33697

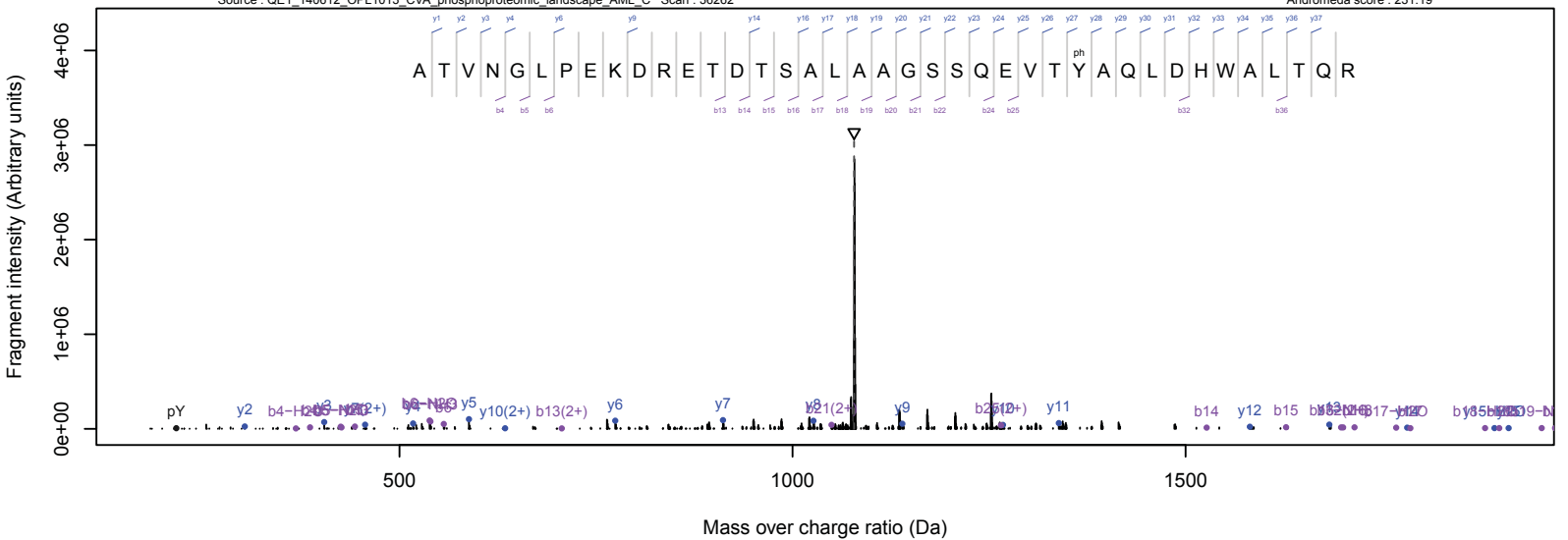
Andromeda score : 231.49



ATVNGLPEKDR(ETD)TSALAAGSSQEVY(ph)AQLDHWALTQR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36262

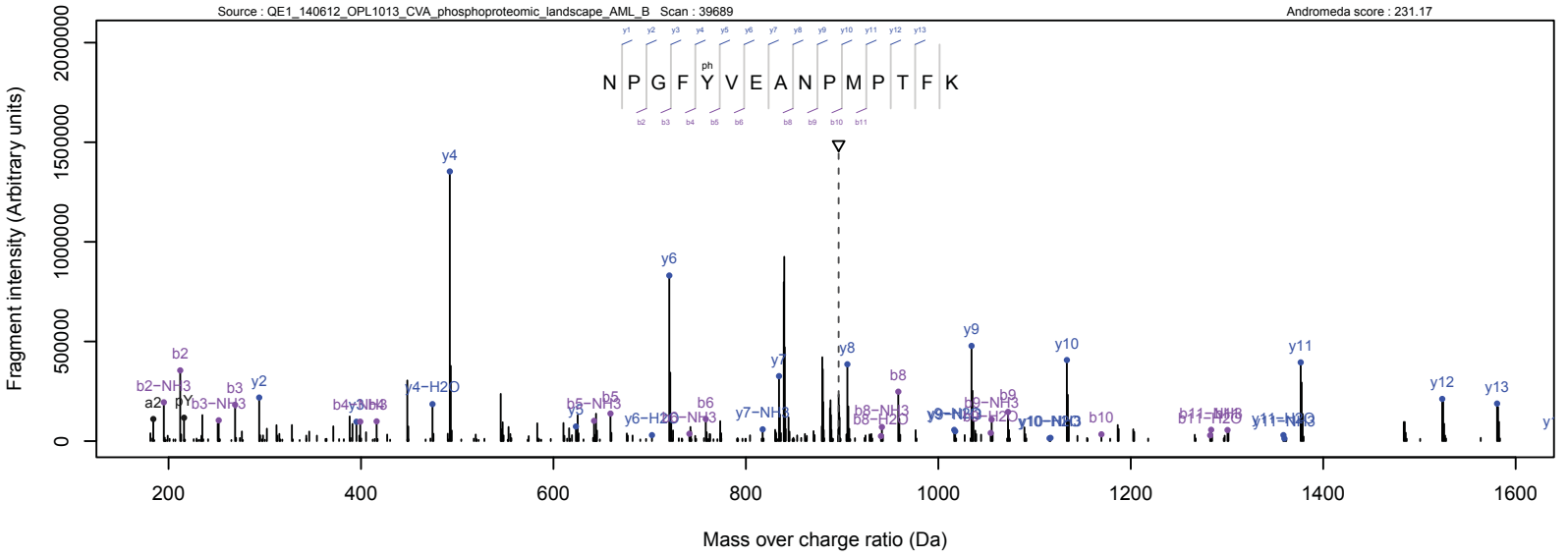
Andromeda score : 231.19



**NPGFY(ph)VEANPMP TFK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39689

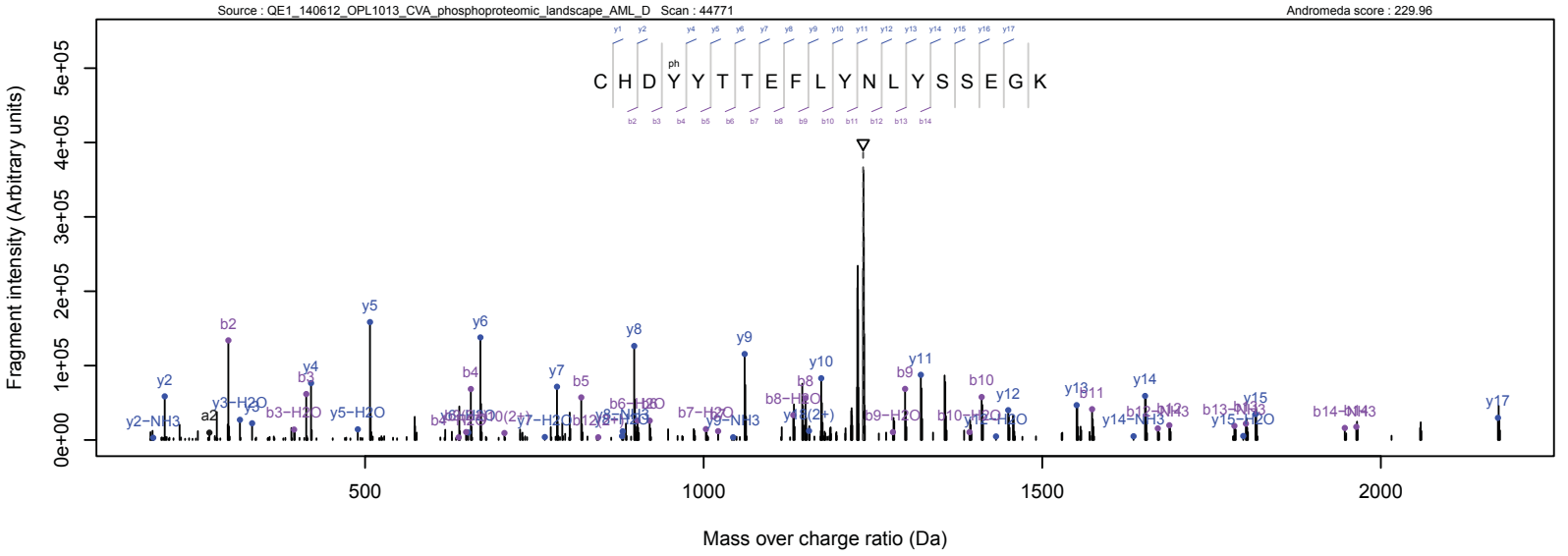
Andromeda score : 231.17



**CHDY(ph)YTTEFLYNLYSSE GK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 44771

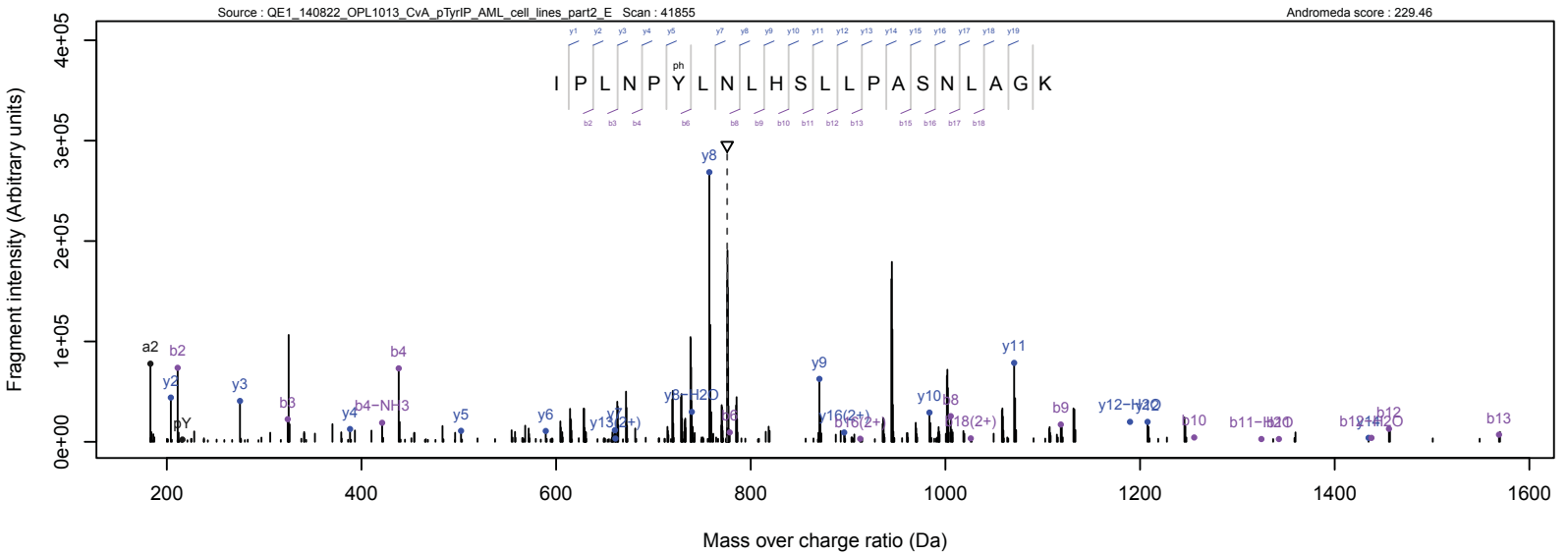
Andromeda score : 229.96



**IPLNPY(ph)LNLHSLLPASNL AGK-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 41855

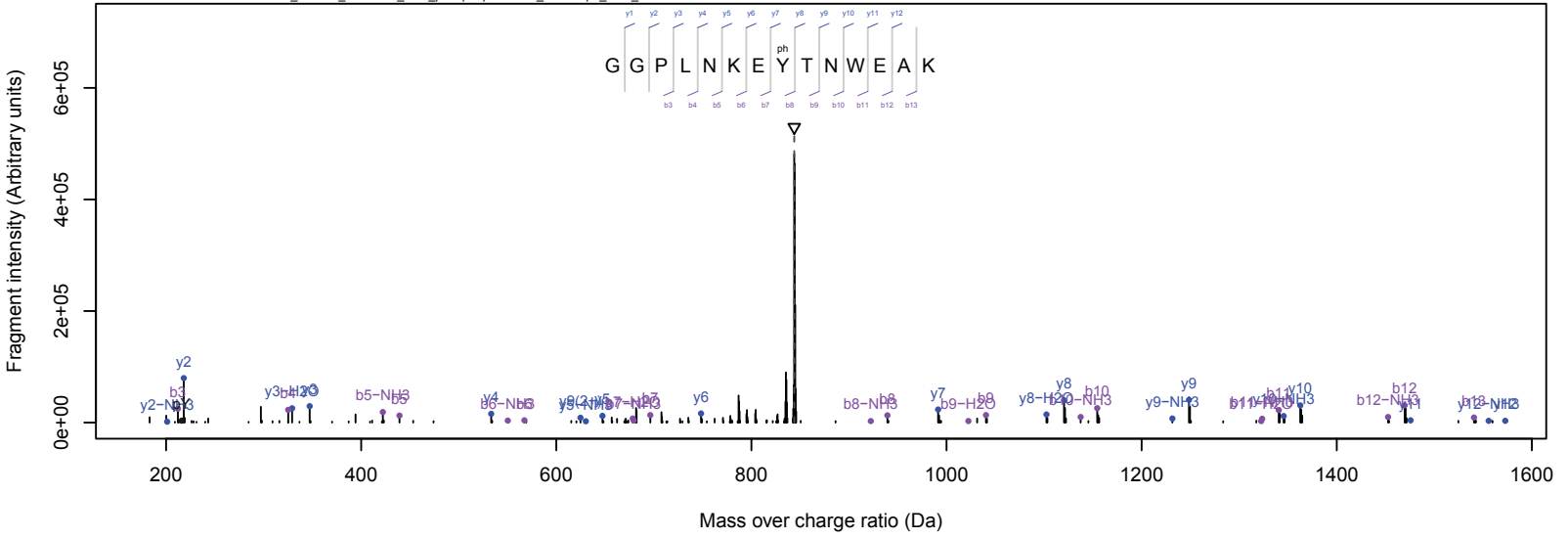
Andromeda score : 229.46



GGPLNKEY(ph)TNWEAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16224

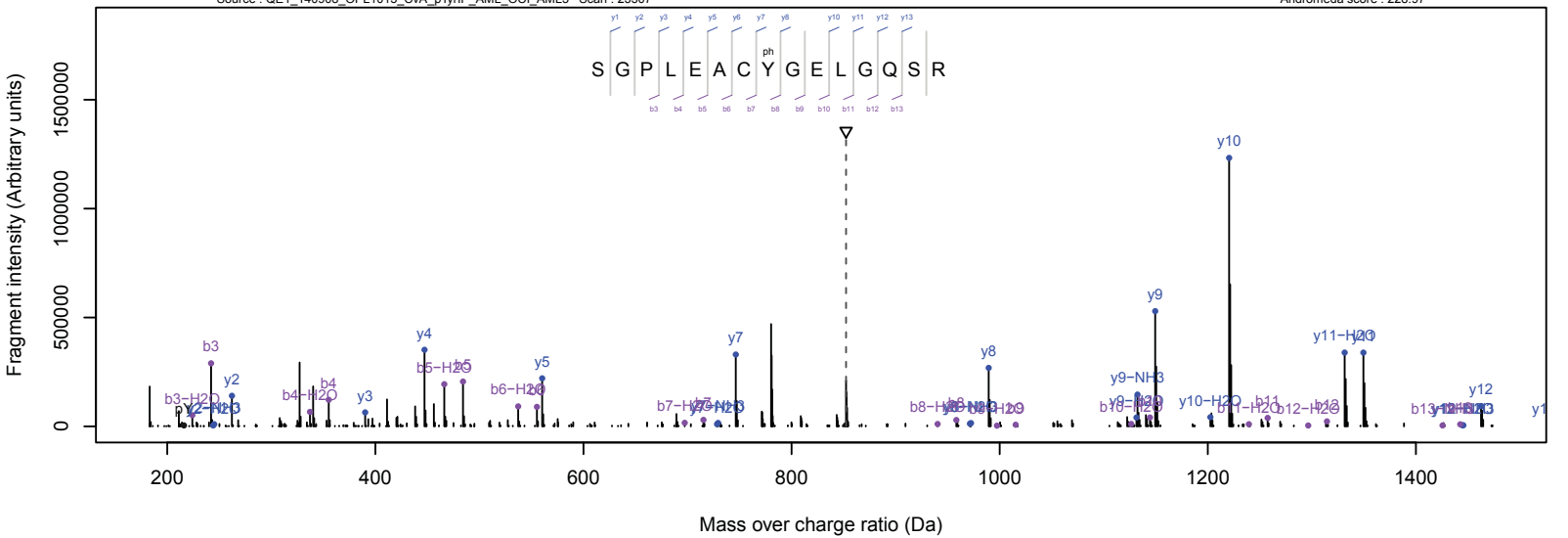
Andromeda score : 229.23



SGPLEACY(ph)GELGQSR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 23307

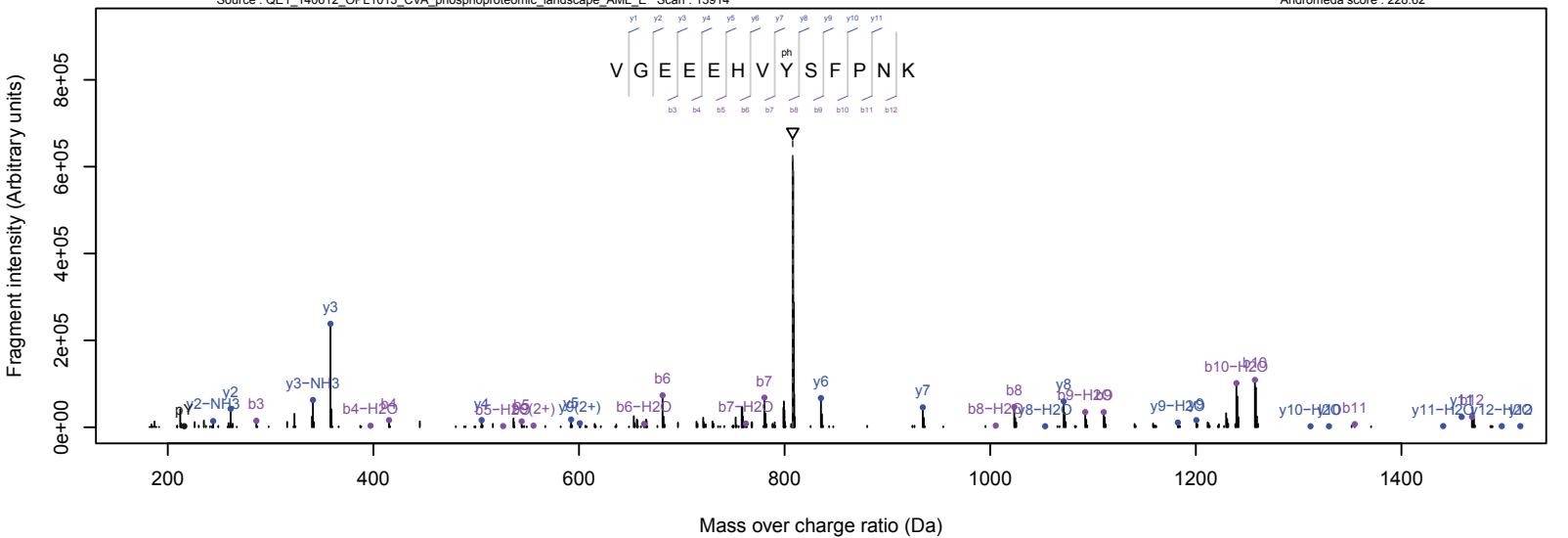
Andromeda score : 228.97



VGEEEHVY(ph)SFPNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 13914

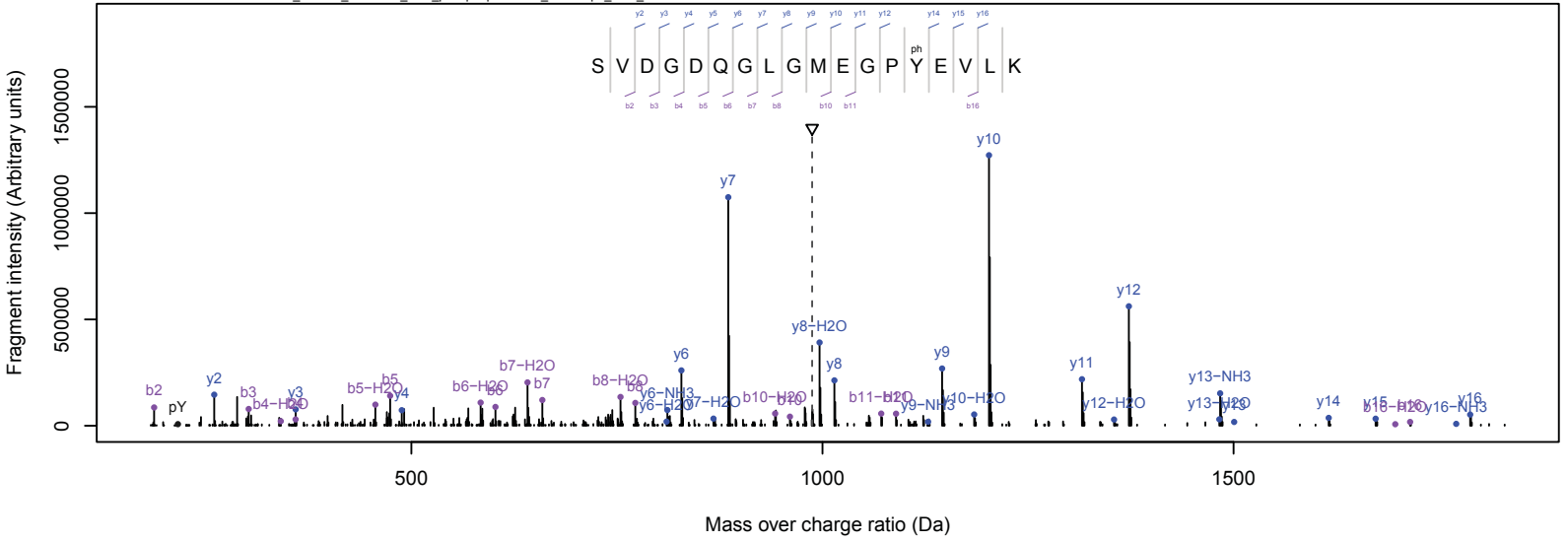
Andromeda score : 228.62



SVDGDQLGMEGPY(ph)EVLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35118

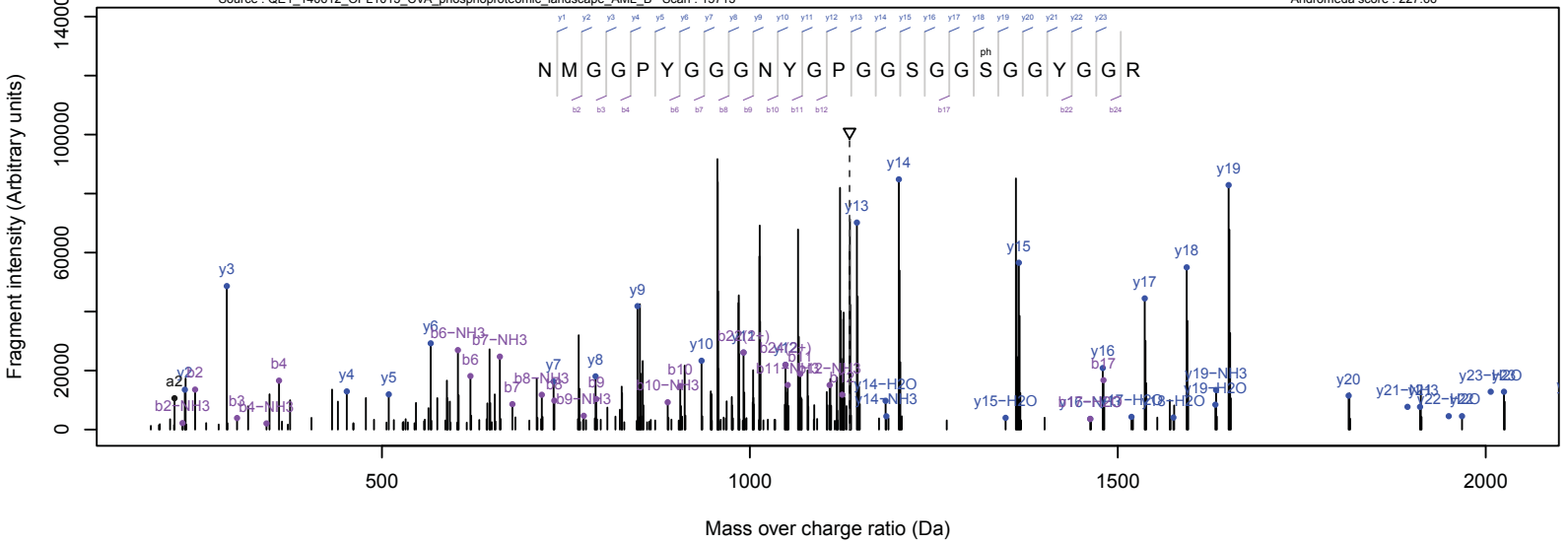
Andromeda score : 228.46



NMGGPYGGNYGPGSSGS(ph)GGYGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15713

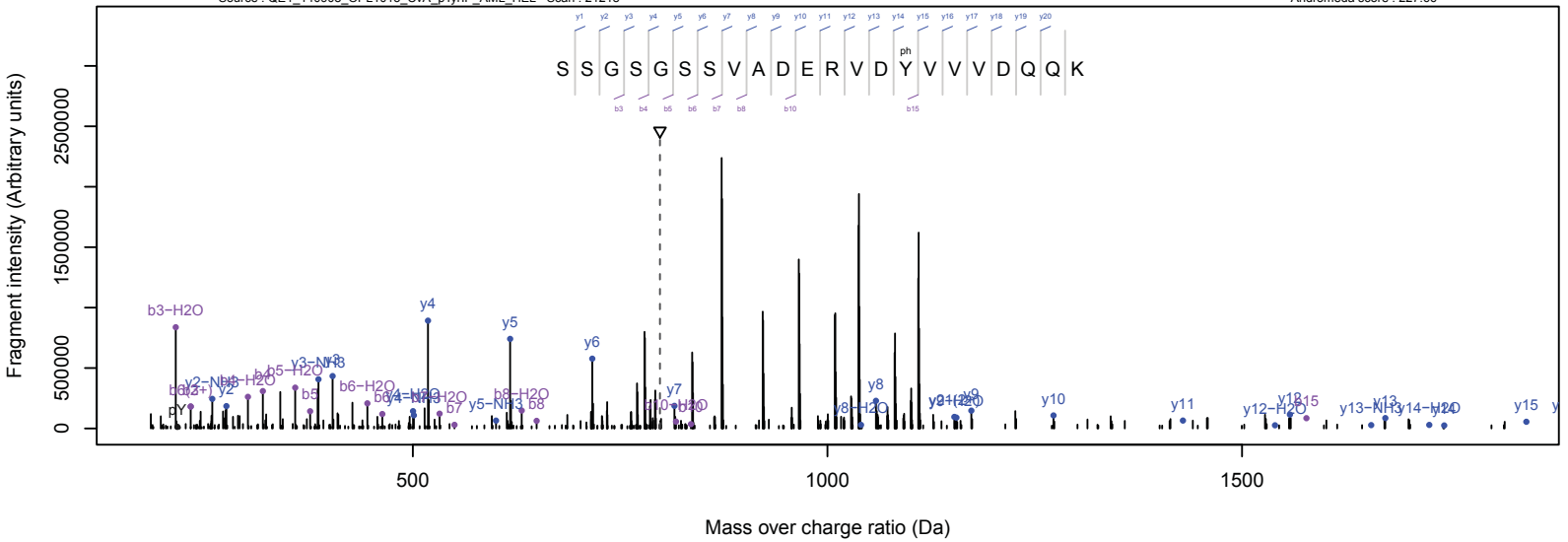
Andromeda score : 227.66



SSGSSVADERVDY(ph)VVDQK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 21218

Andromeda score : 227.66



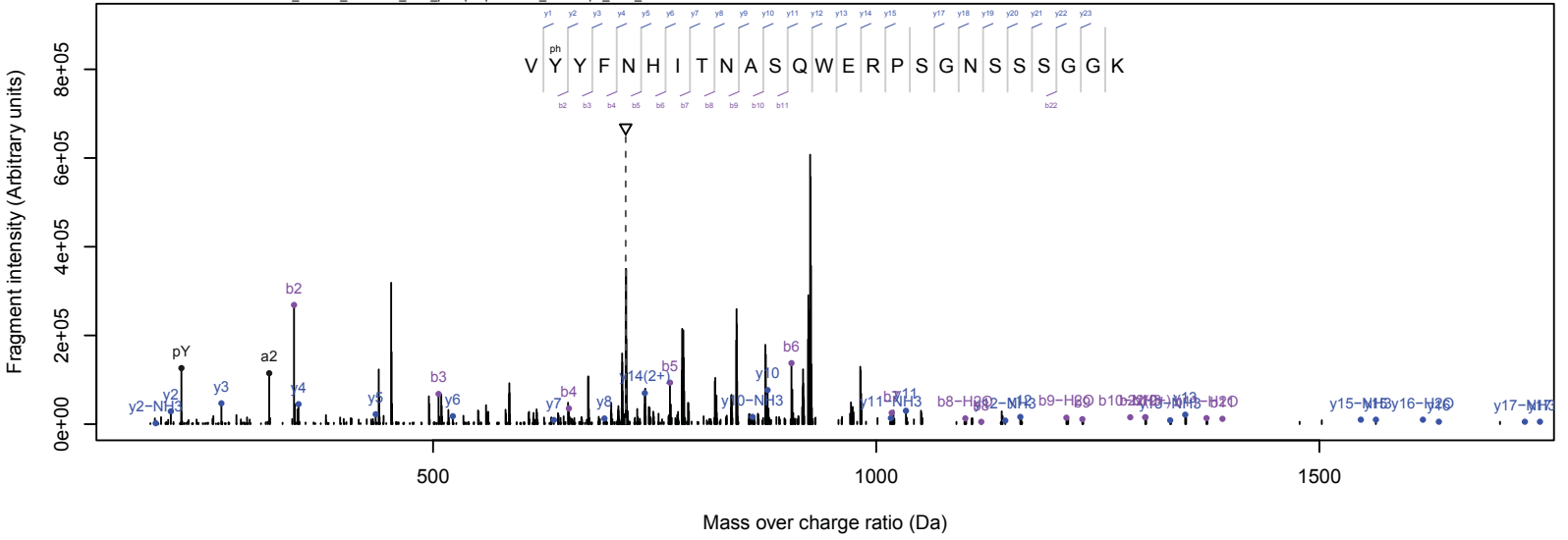




VY(ph)YFNHITNASQWERPSGNSSSGGK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27218

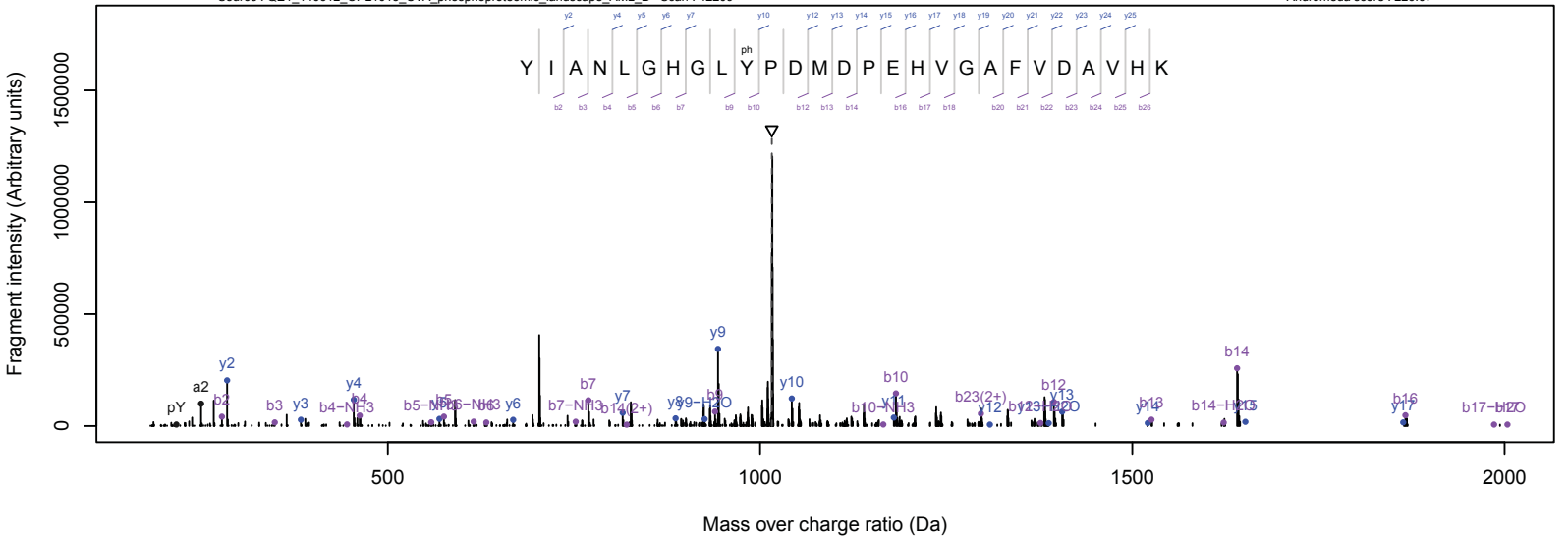
Andromeda score : 226.56



YIANLGHGLY(ph)PDMDPEHVGFVDAVHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42209

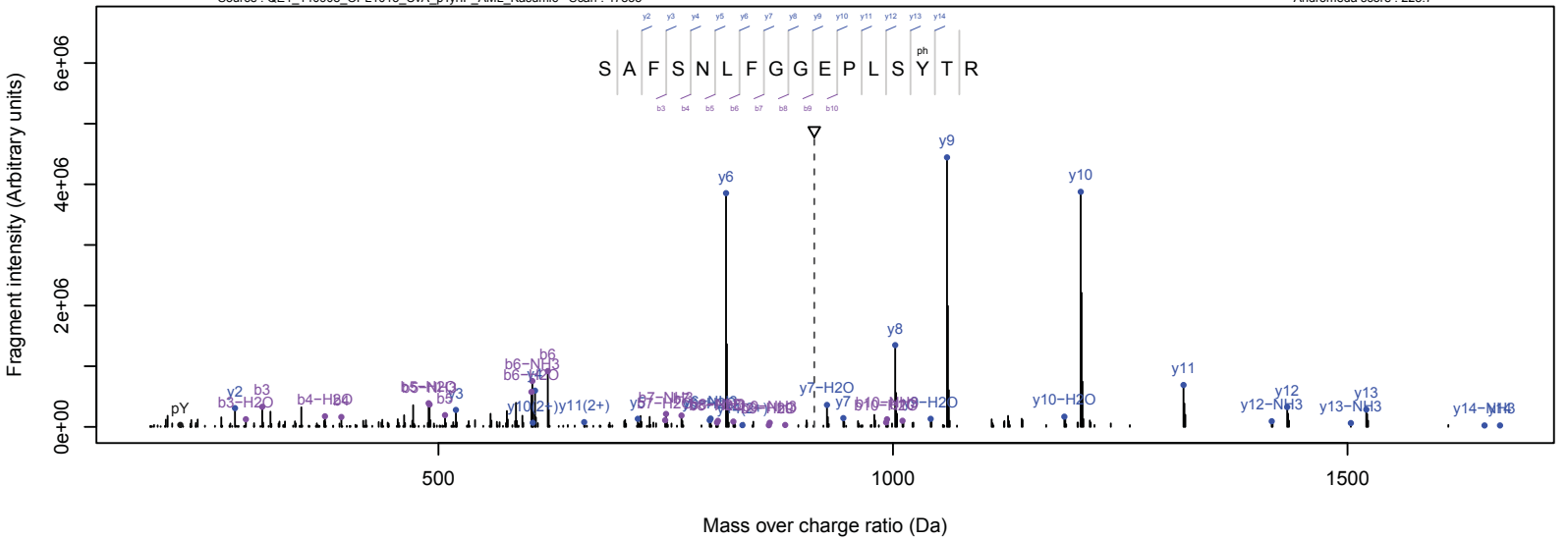
Andromeda score : 226.07



SAFSNLFGGELPSY(ph)TR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 47855

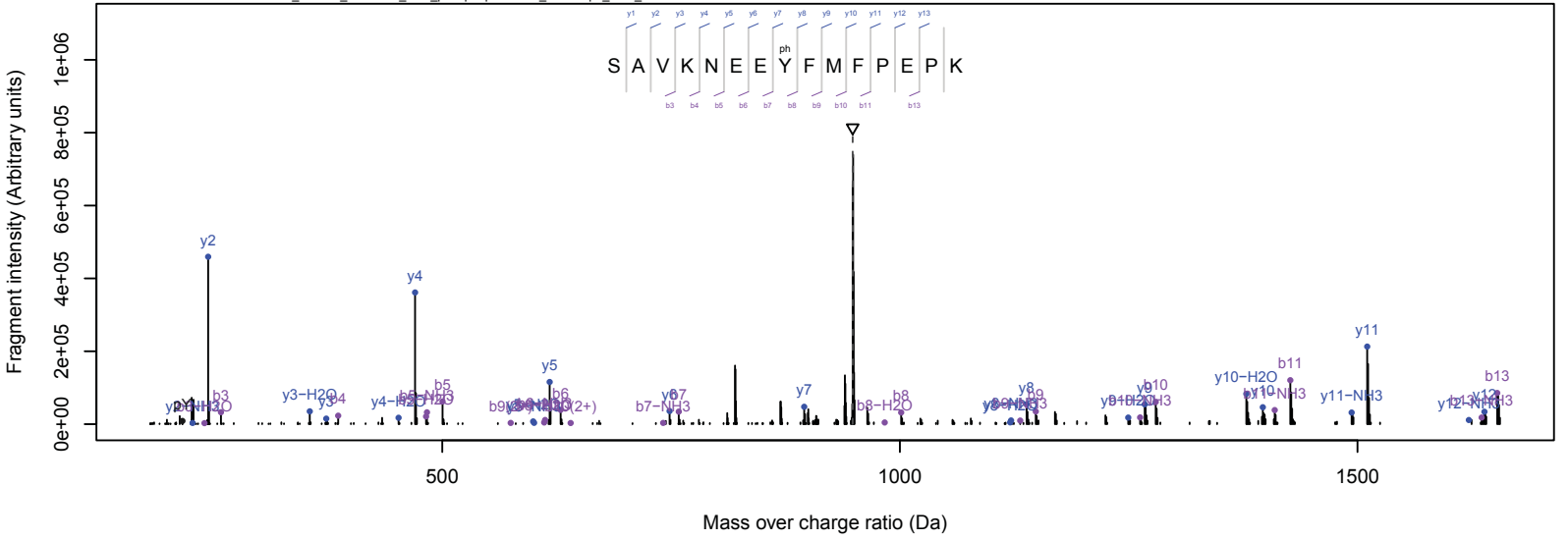
Andromeda score : 225.7



SAVKNEEY(ph)FMFPEPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31265

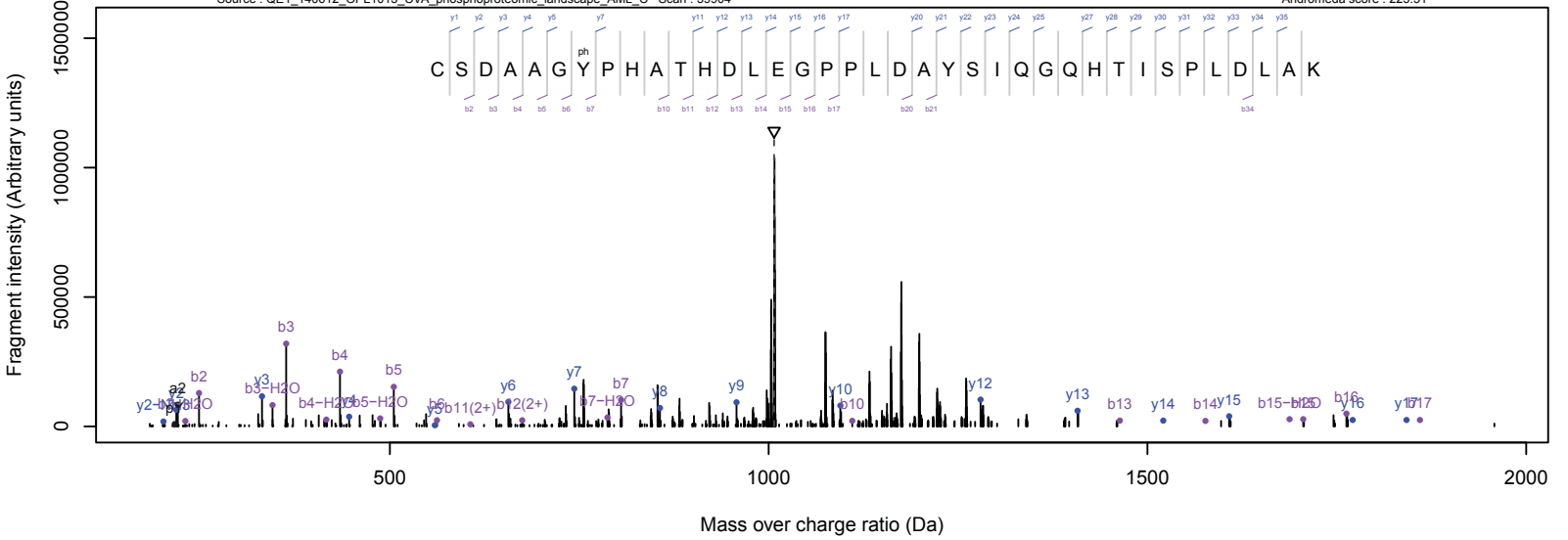
Andromeda score : 225.7



CSDAAGY(ph)PHATHDLEGPPLDAYSIQGQHTISPLDLAK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39904

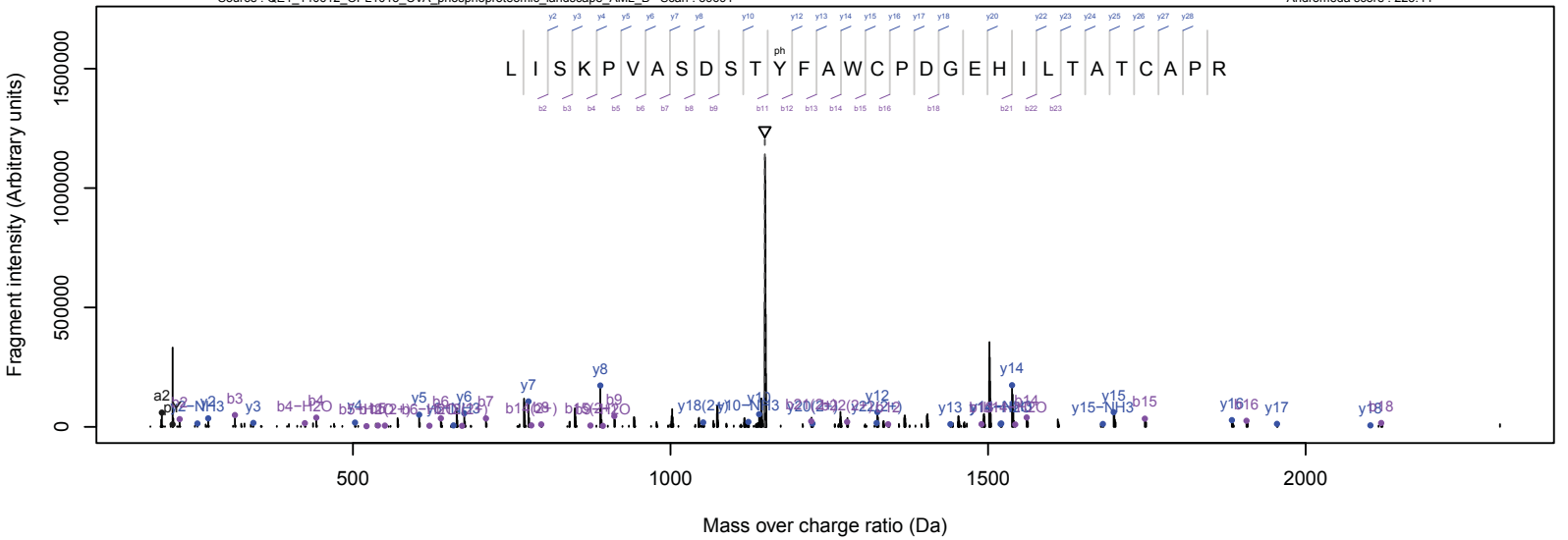
Andromeda score : 225.51



LISKPVASDSTY(ph)FAWC PDGEHILTATCAPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39691

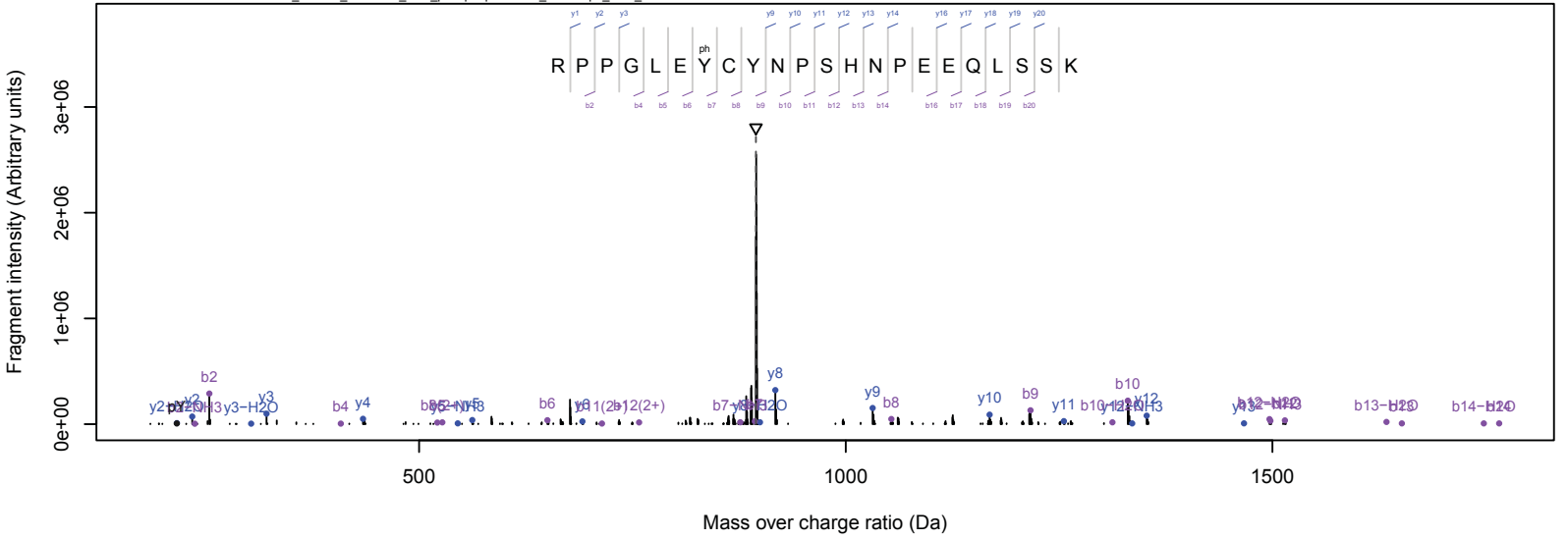
Andromeda score : 225.44



RPPGLELY(ph)CYNP SHNP EEQLSSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 21889

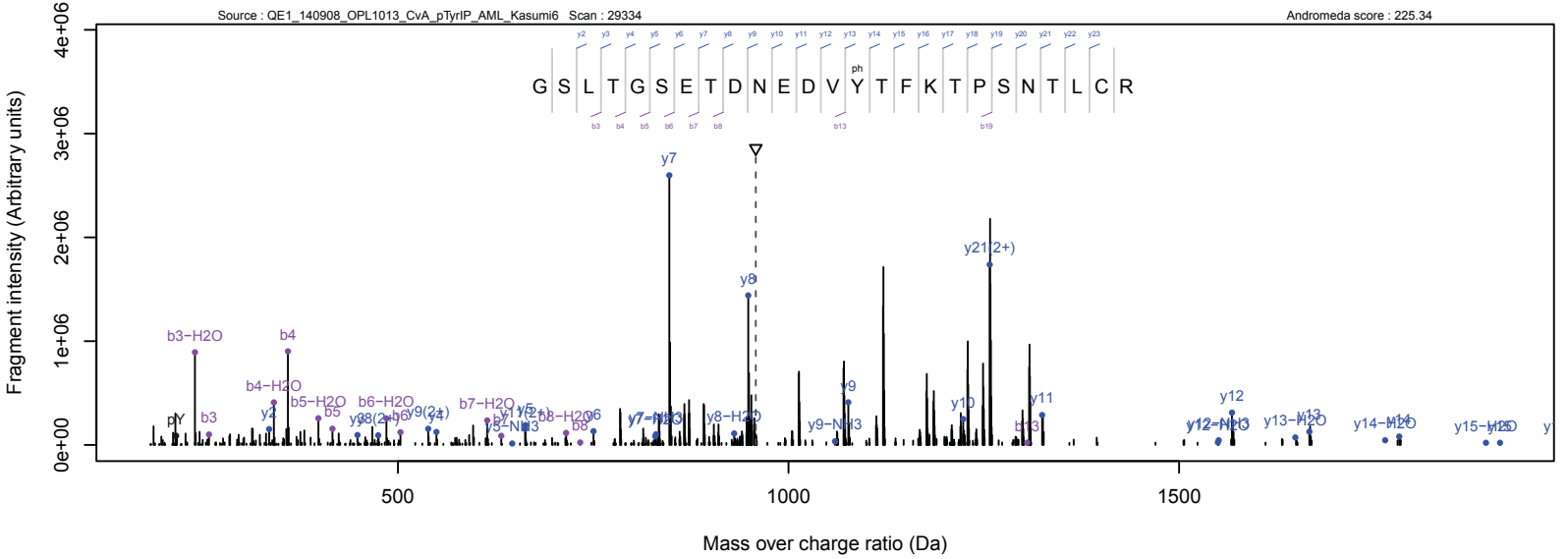
Andromeda score: 225.39



GSLTGSETDNEDVY(ph)TFKTPSNTLCR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 29334

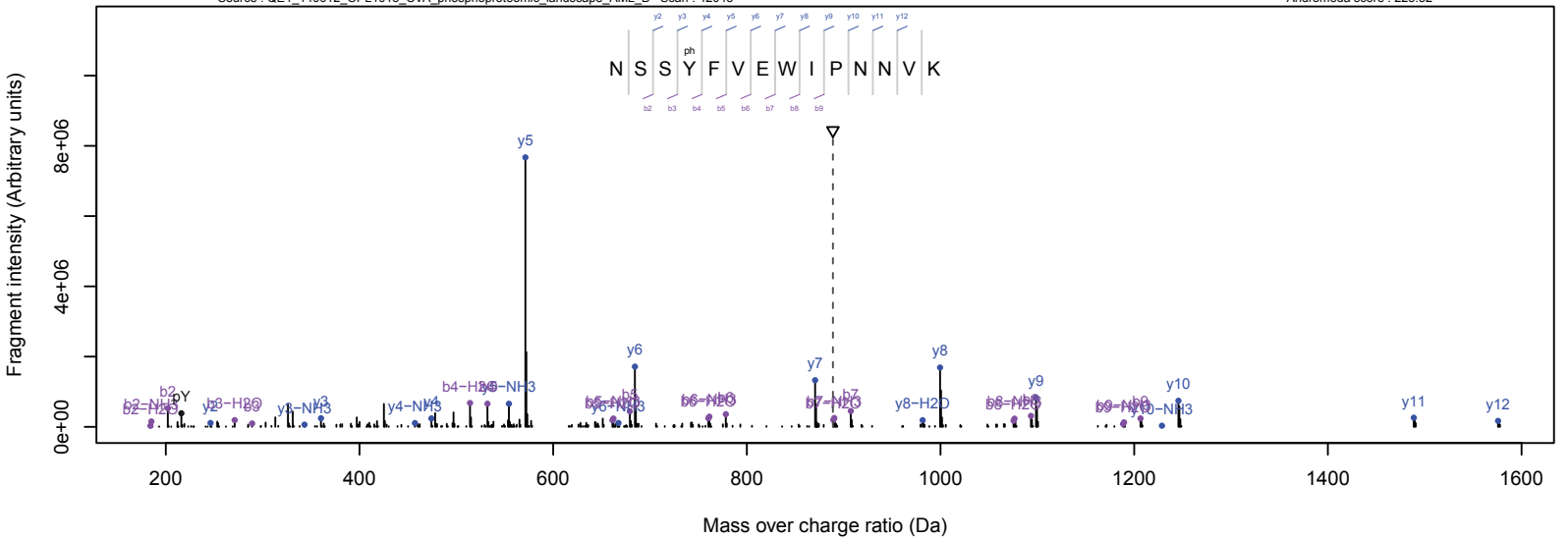
Andromeda score: 225.34



NSSY(ph)FVEWIPNNVK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 42043

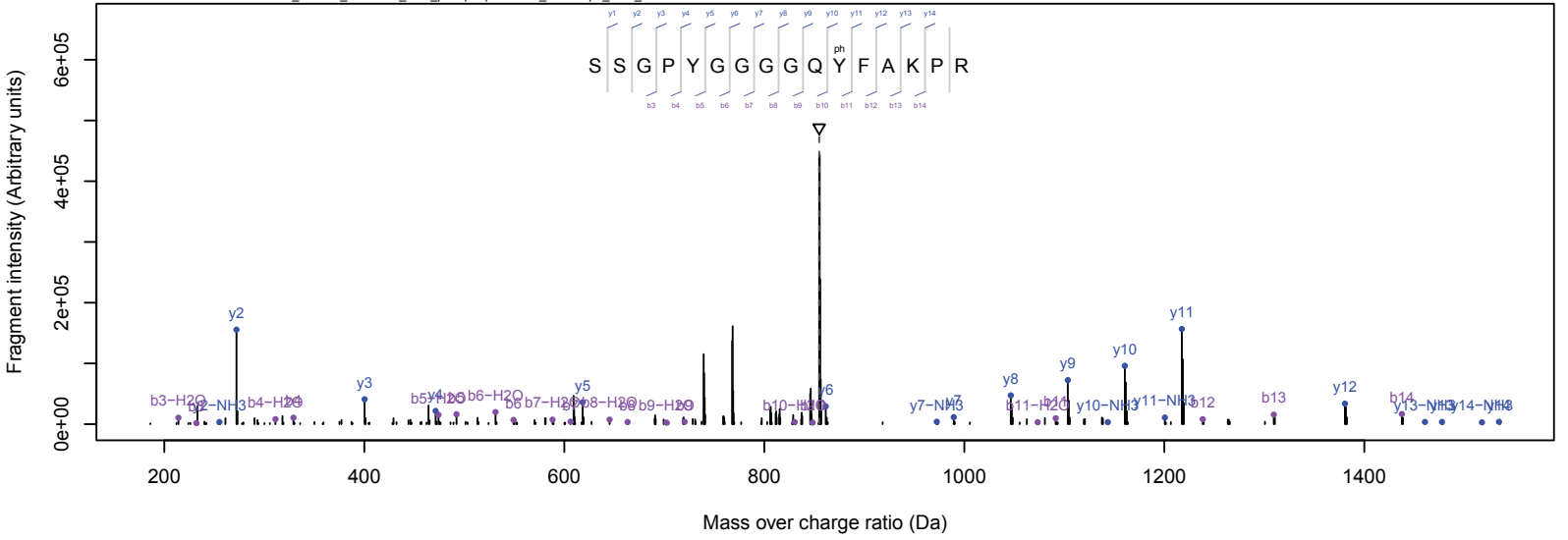
Andromeda score: 225.32



SSGPYGGGGQY(ph)FAKPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 13123

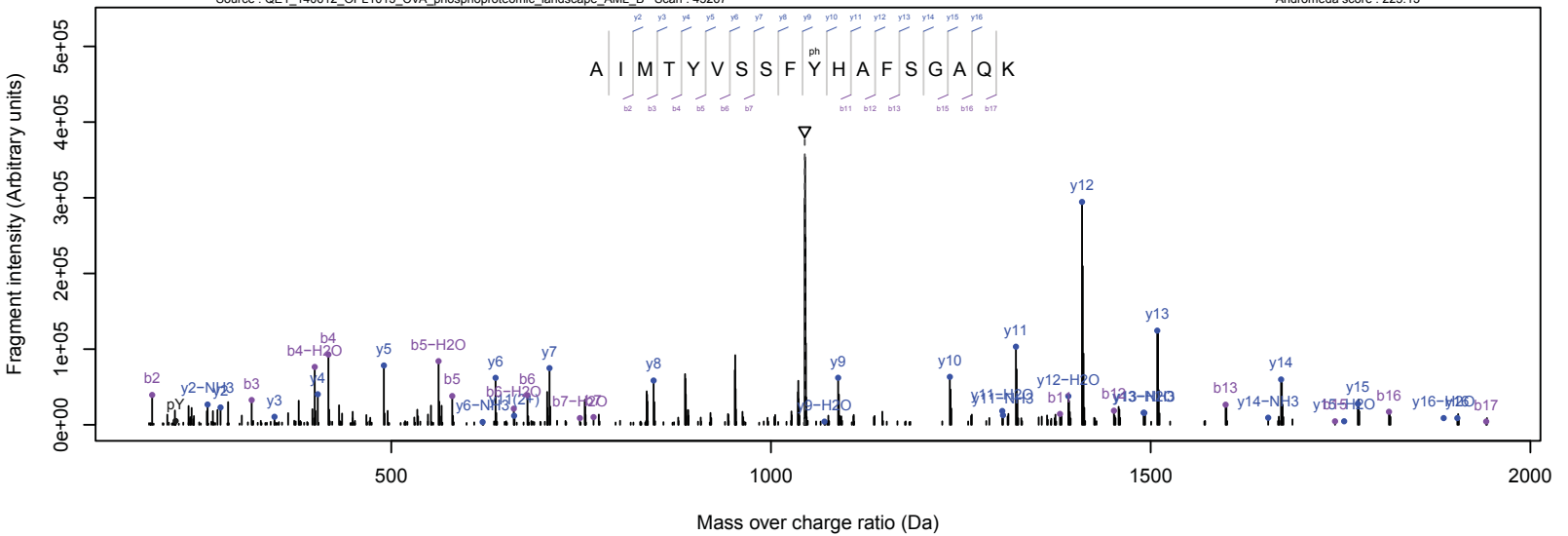
Andromeda score : 225.29



AIMTYVSSFY(ph)HAFSGAQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45267

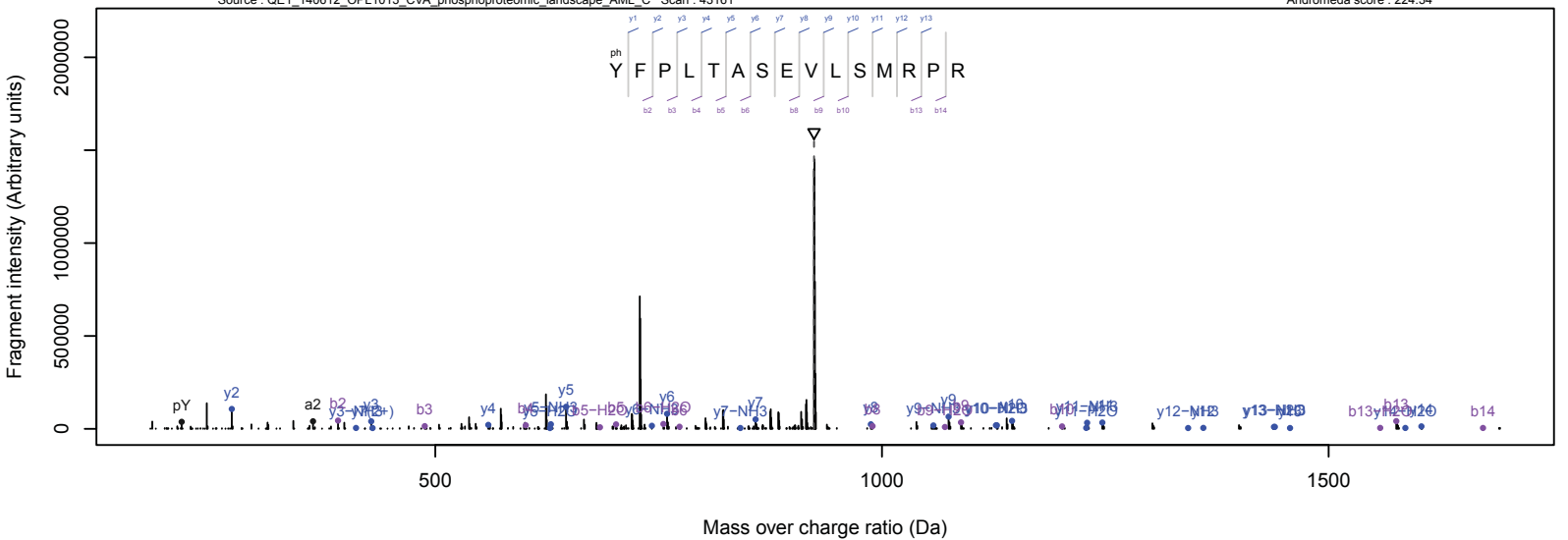
Andromeda score : 225.13



Y(ph)FPLTASEVLSMRPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43161

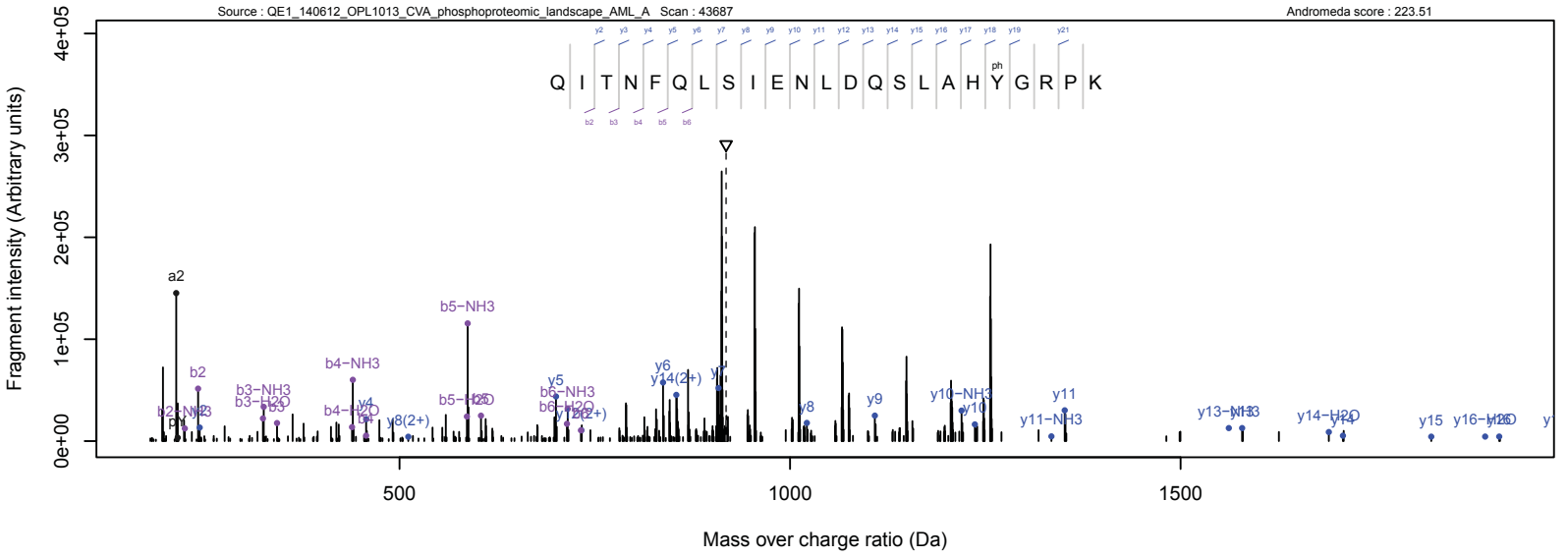
Andromeda score : 224.34



**QITNFQLSIENLDQSLAHY(ph)GRPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 43687

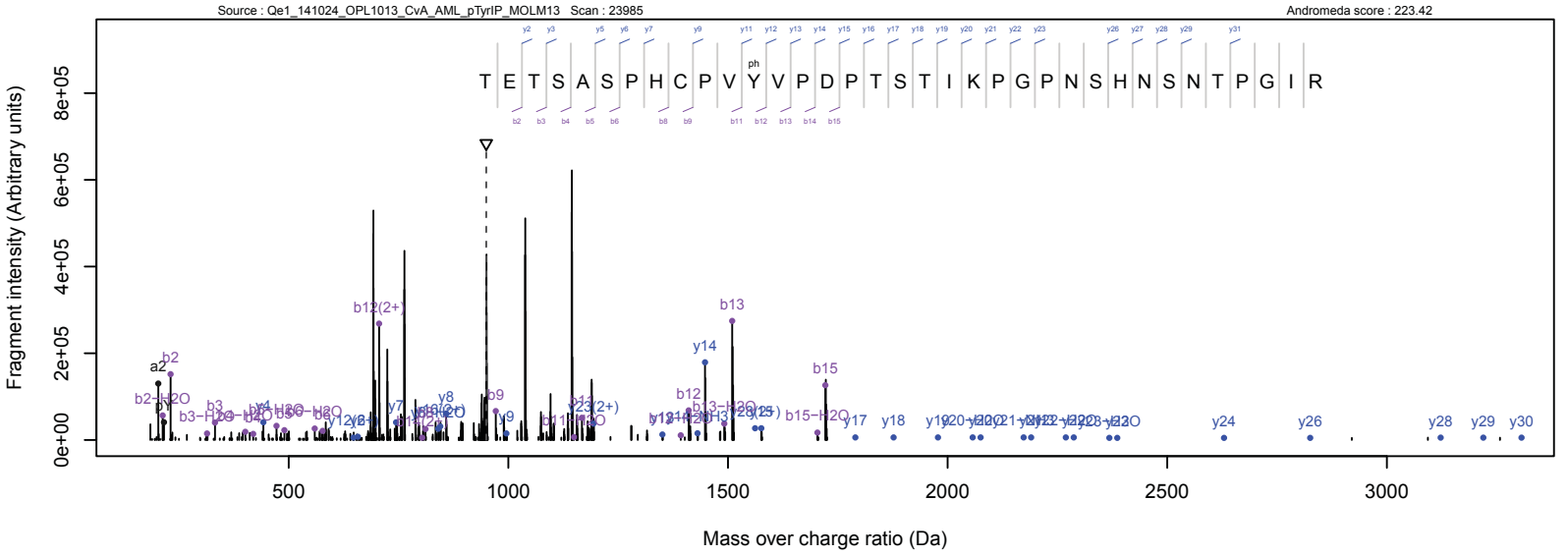
Andromeda score : 223.51



**TETSASPHCPVY(ph)VPDPTSTIKPGPNSHNSNTPGIR-4+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 23985

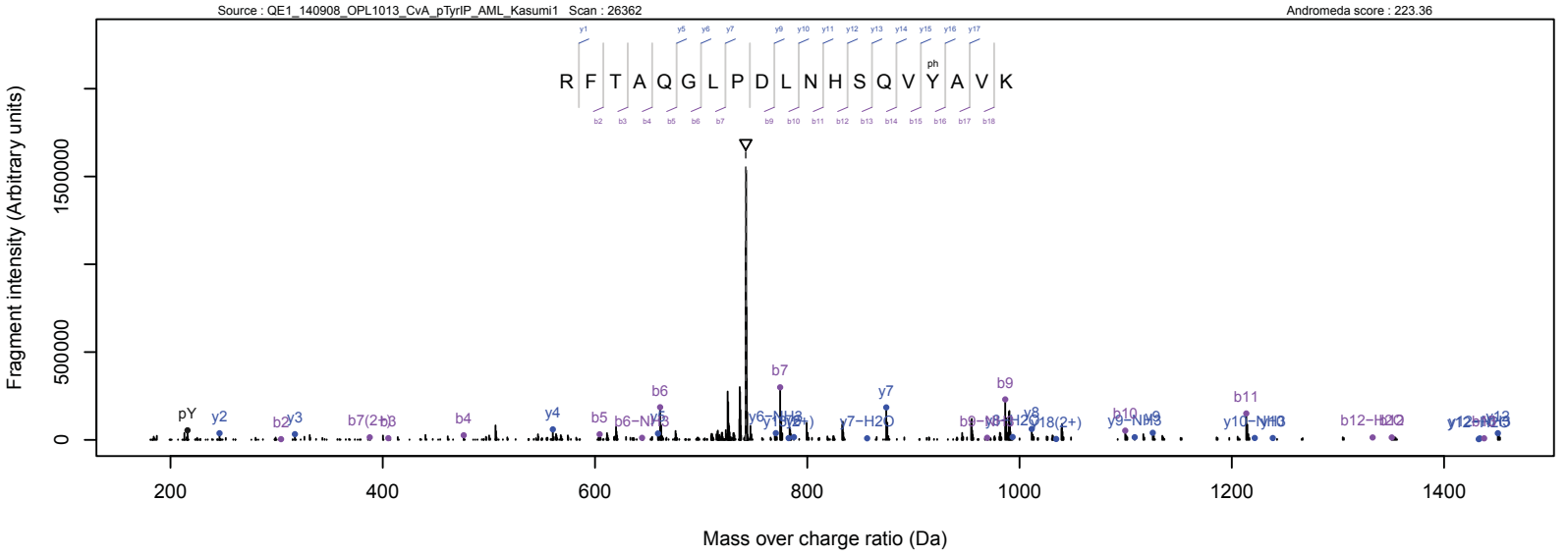
Andromeda score : 223.42



**RFTAQGLPDLNHSQVYA V K-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 26362

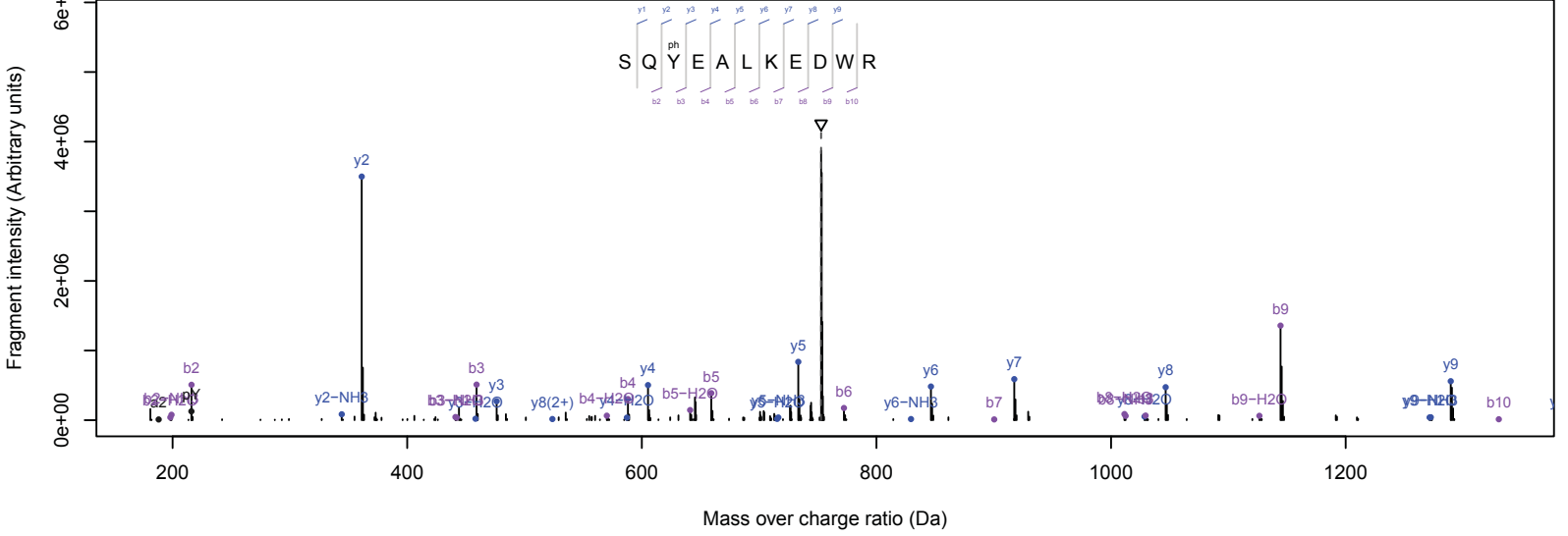
Andromeda score : 223.36



SQY(ph)EALKEDWR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17667

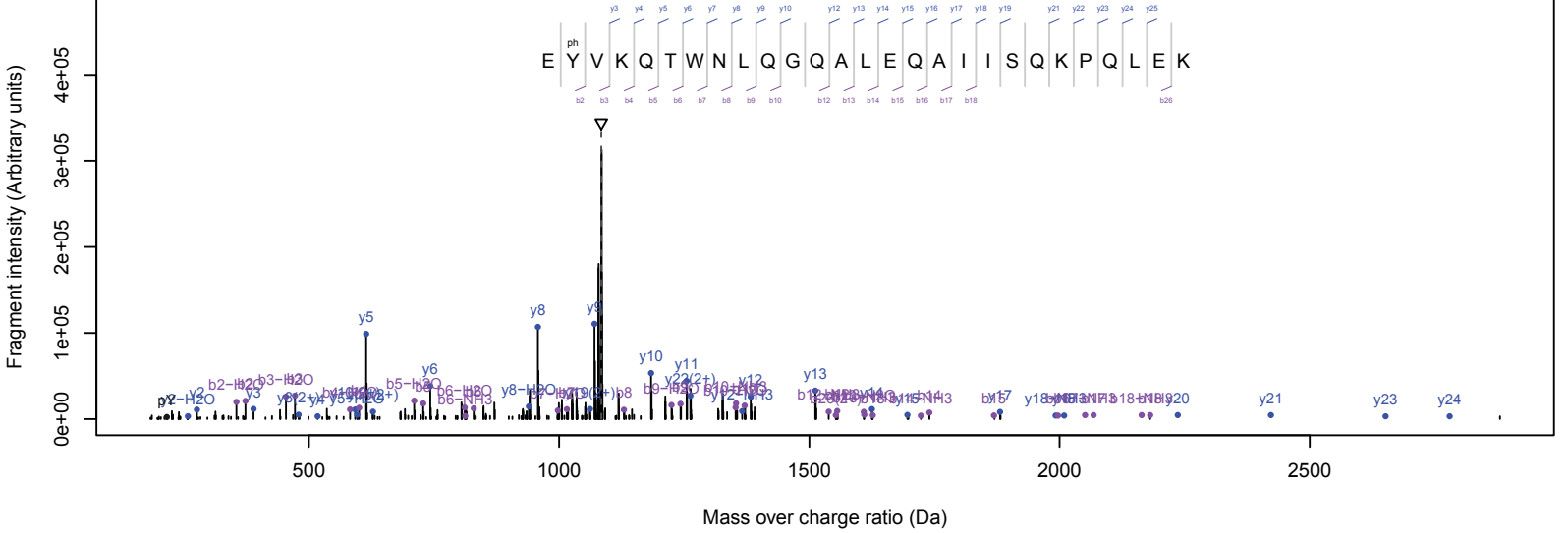
Andromeda score : 223.22



EY(ph)VKQTNLQGGALEQAISQKPQLEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41031

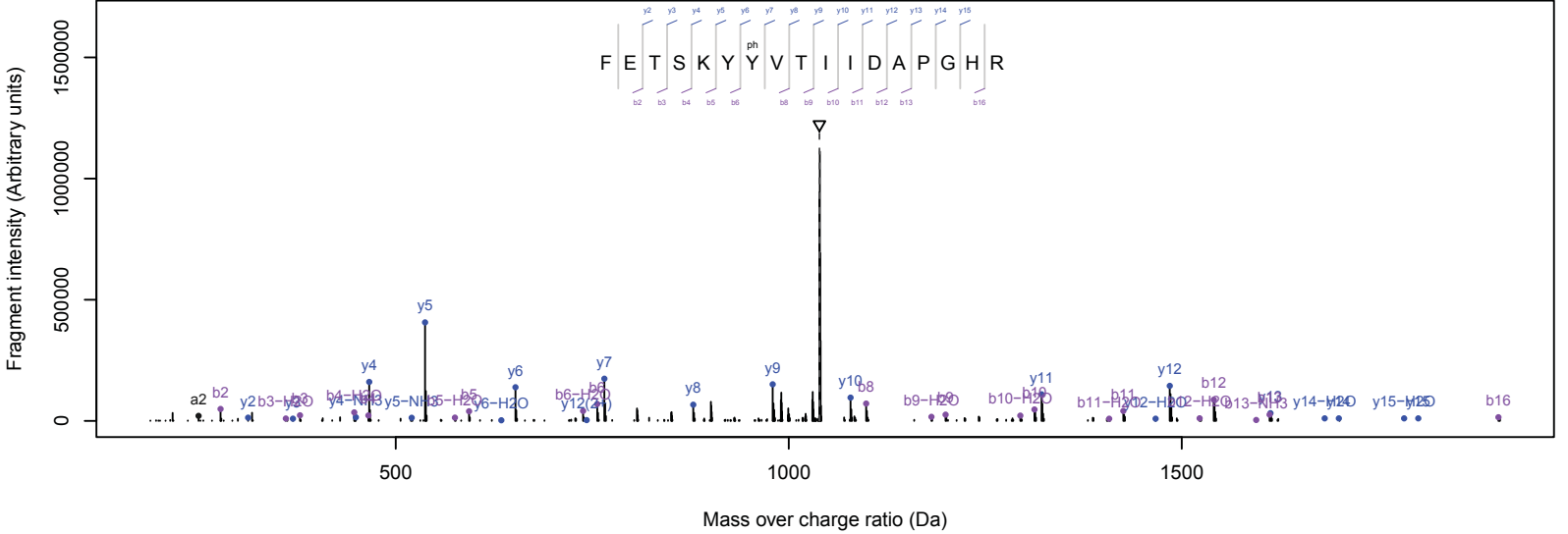
Andromeda score : 223.15



FETSKYY(ph)VTIIDAPGHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28237

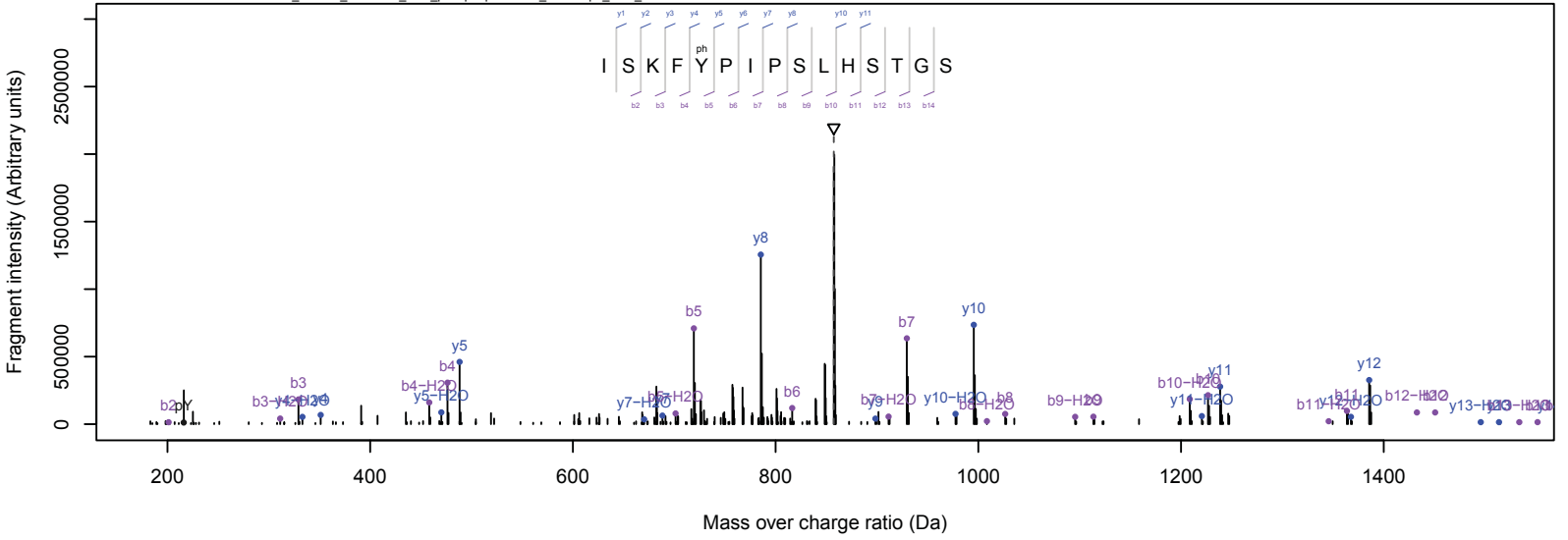
Andromeda score : 223.06



ISKFY(ph)PIPSLHSTGS-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 30475

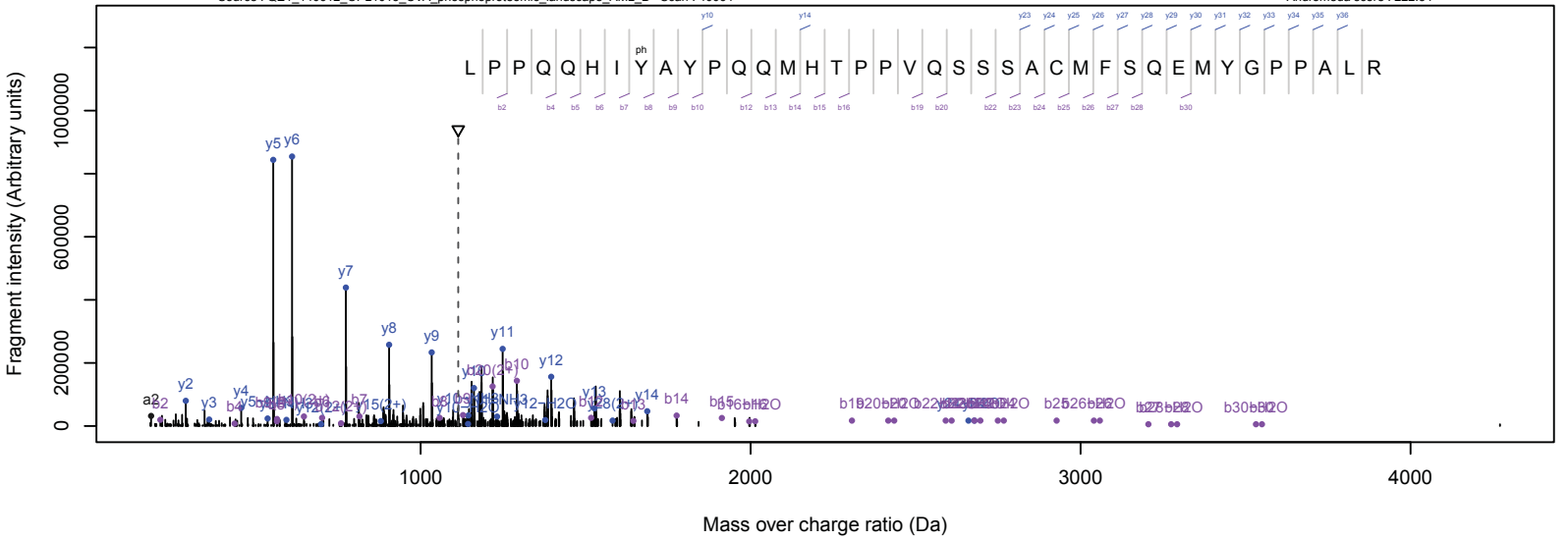
Andromeda score: 222.75



LPPQHHIY(ph)AYPQQMHTPPVQSSSACMFSEQEMYGPPALR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 40064

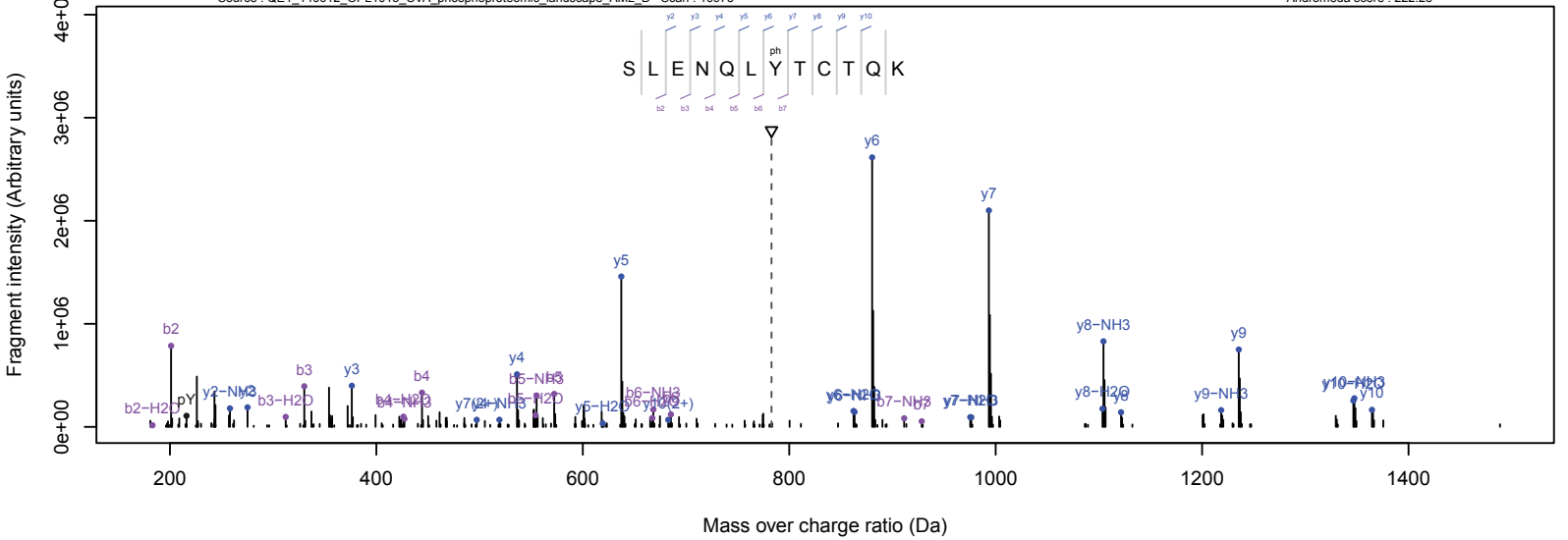
Andromeda score: 222.34



SLENQLY(ph)TCTQK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 16975

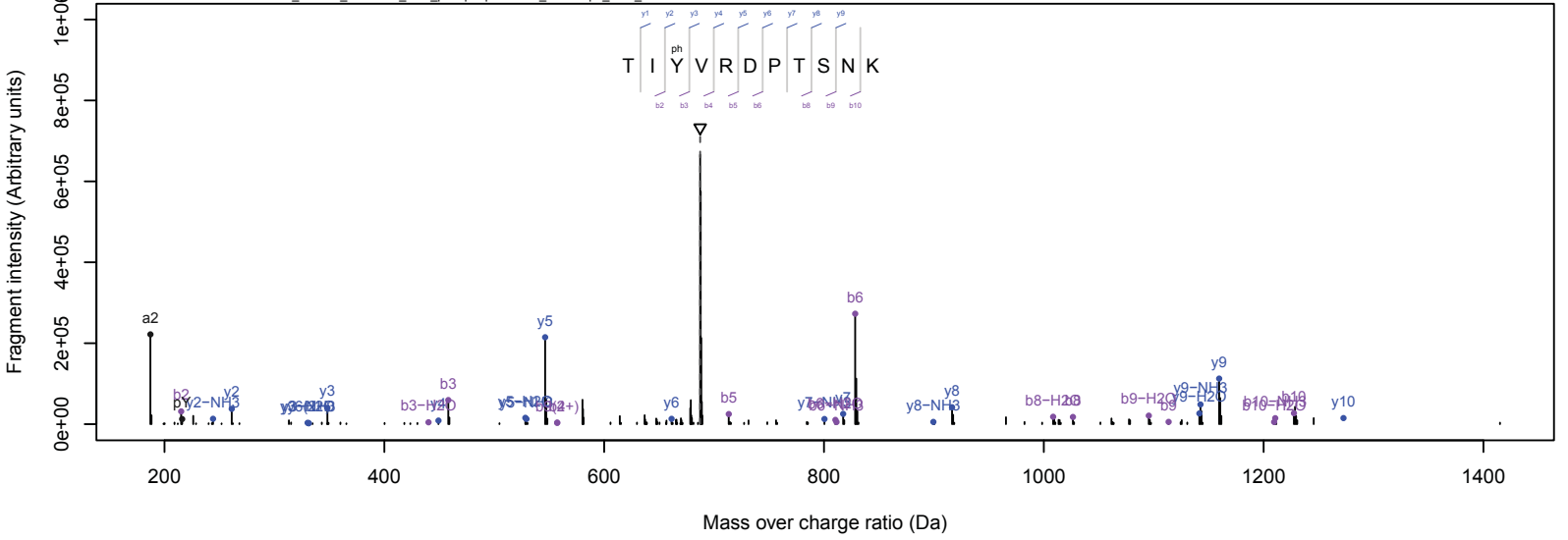
Andromeda score: 222.25



TIY(ph)VRDPTS<sup>2+</sup>

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10585

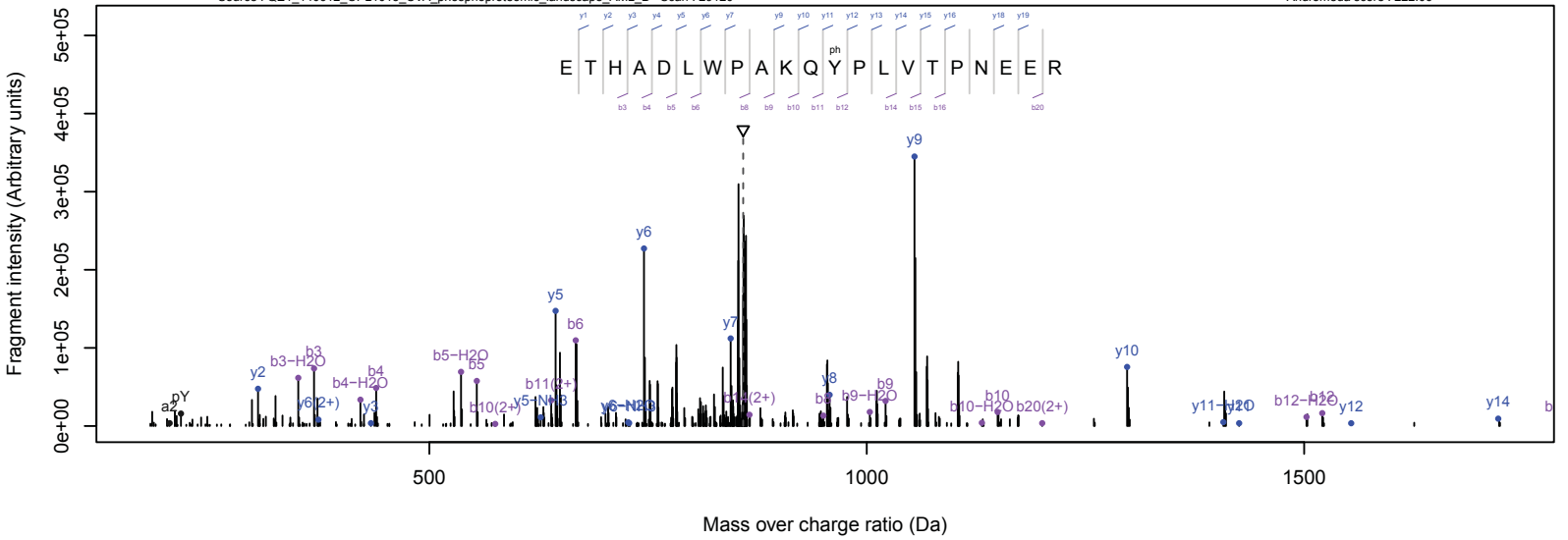
Andromeda score: 222.2



ETHADLWPAKQY(ph)PLVTPNEER-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 29120

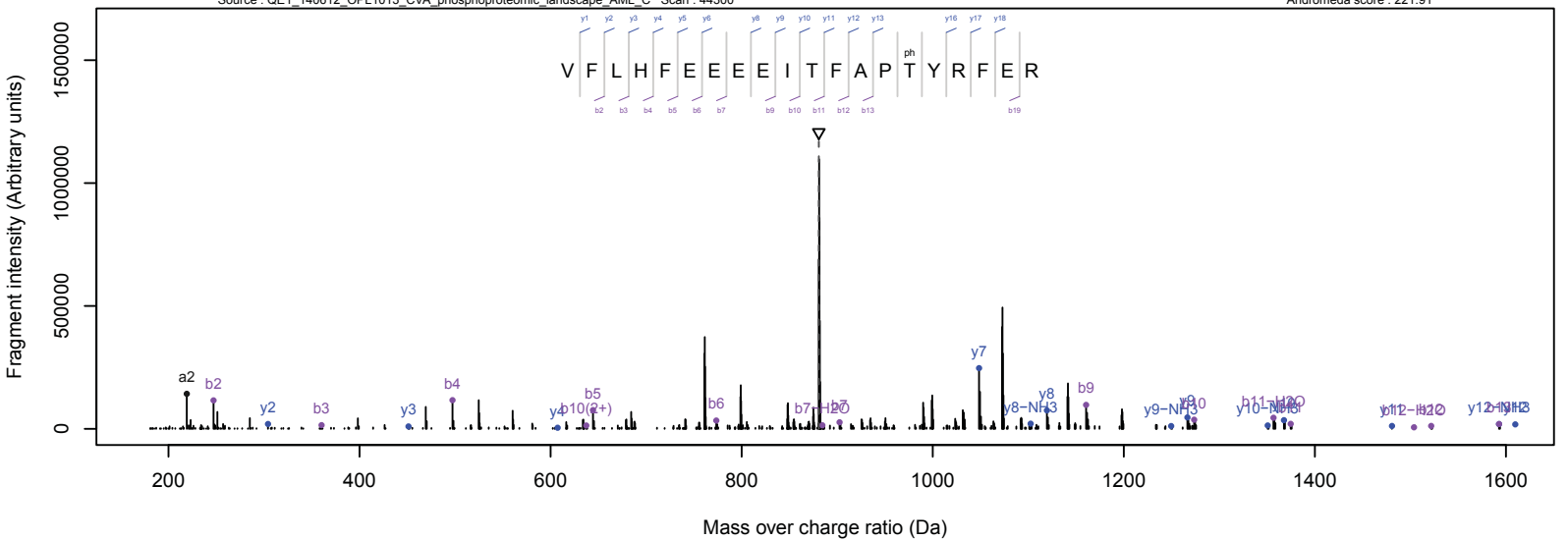
Andromeda score: 222.06



VFLHFEETITFAPTYRFER-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 44300

Andromeda score: 221.91

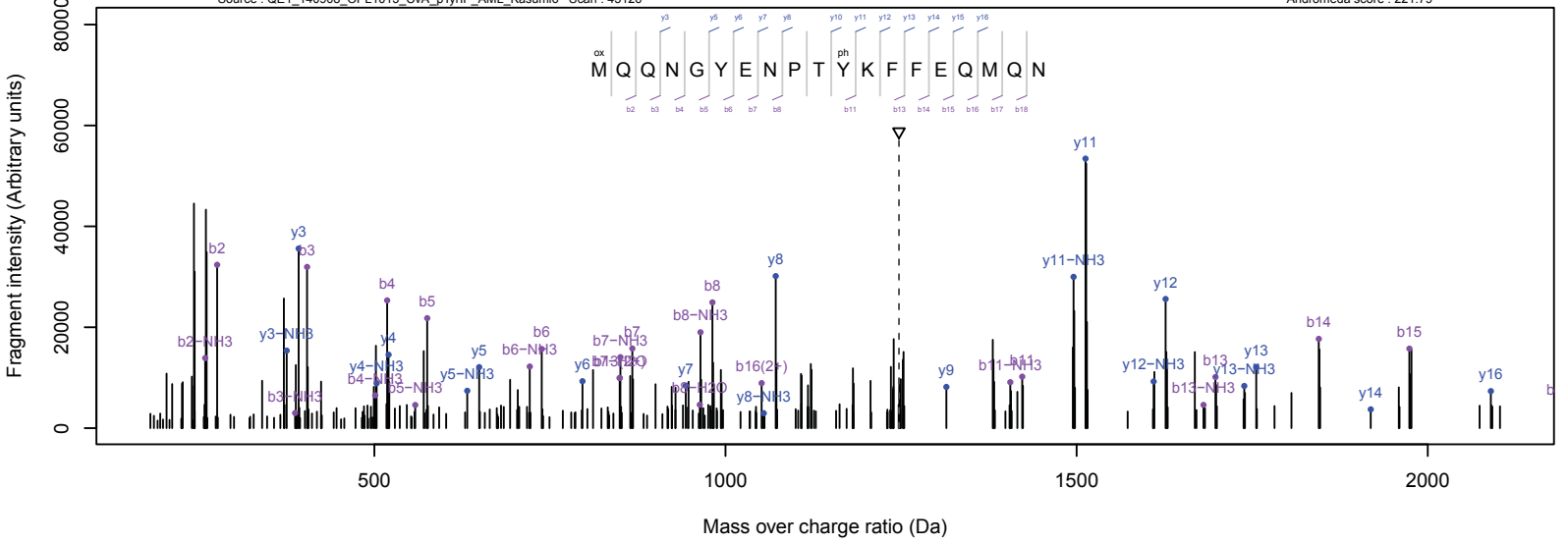




M(ox)QQNGYENPTYKFFEQMQN-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 43126

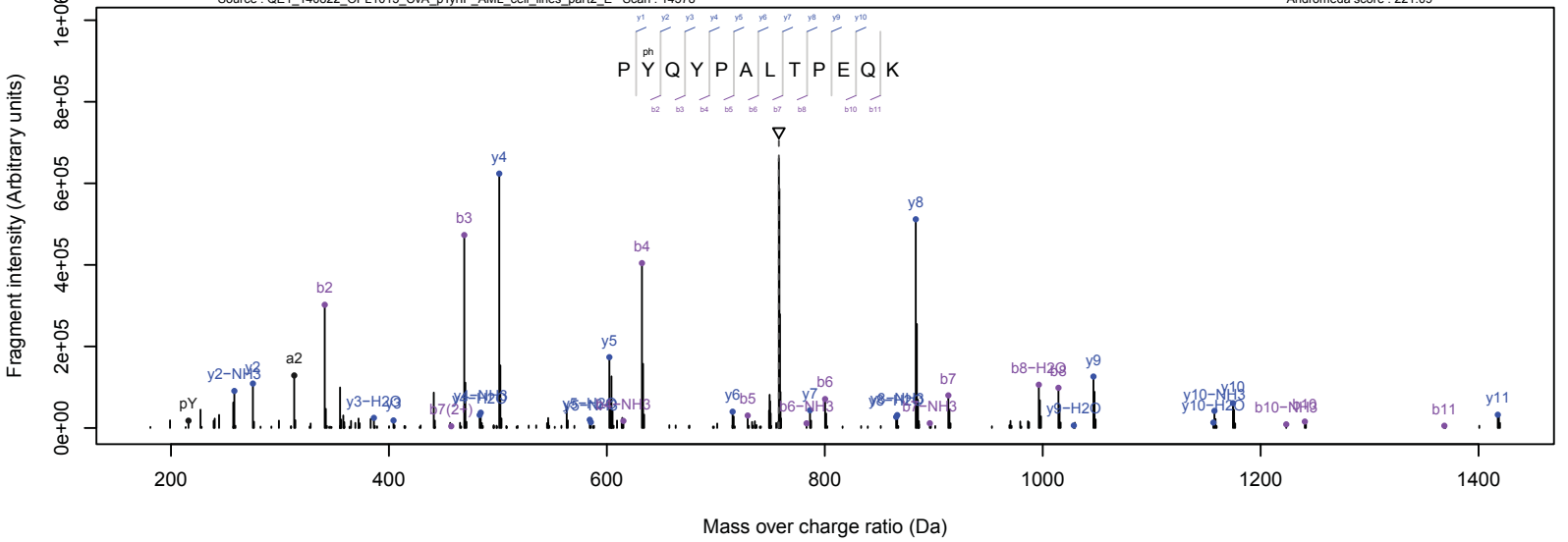
Andromeda score: 221.79



PY(ph)QYPALTPQK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 14578

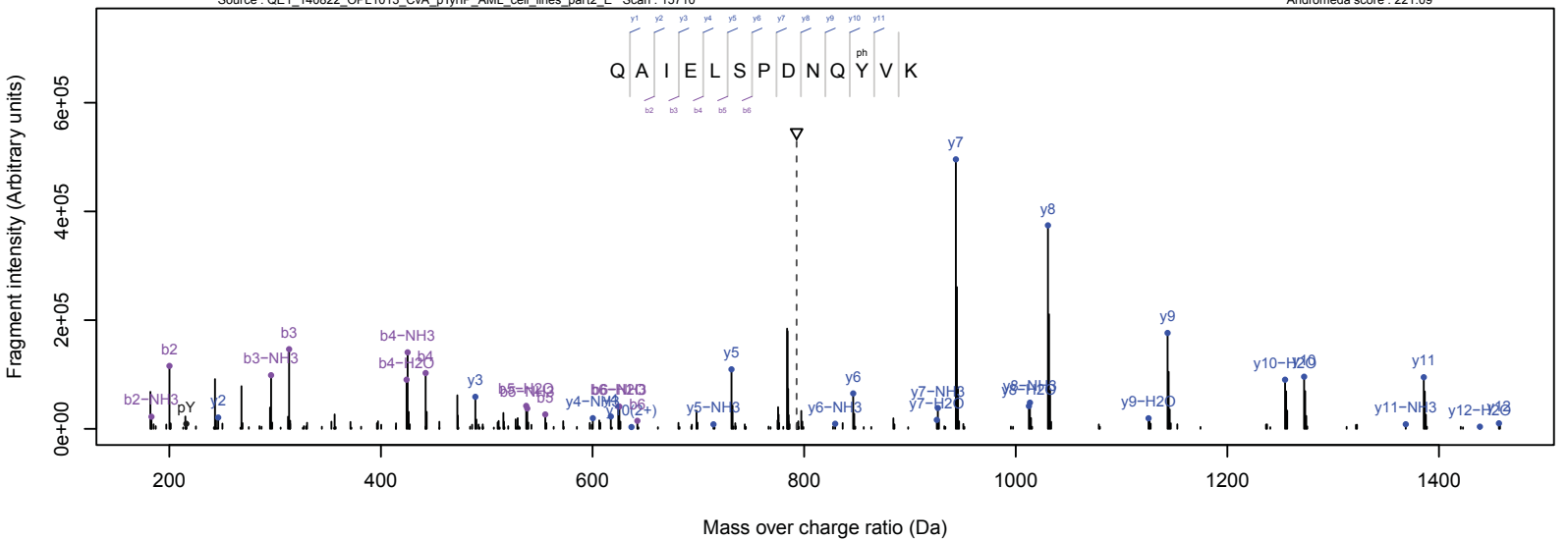
Andromeda score: 221.09



QAIELSPDNQYVK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 15710

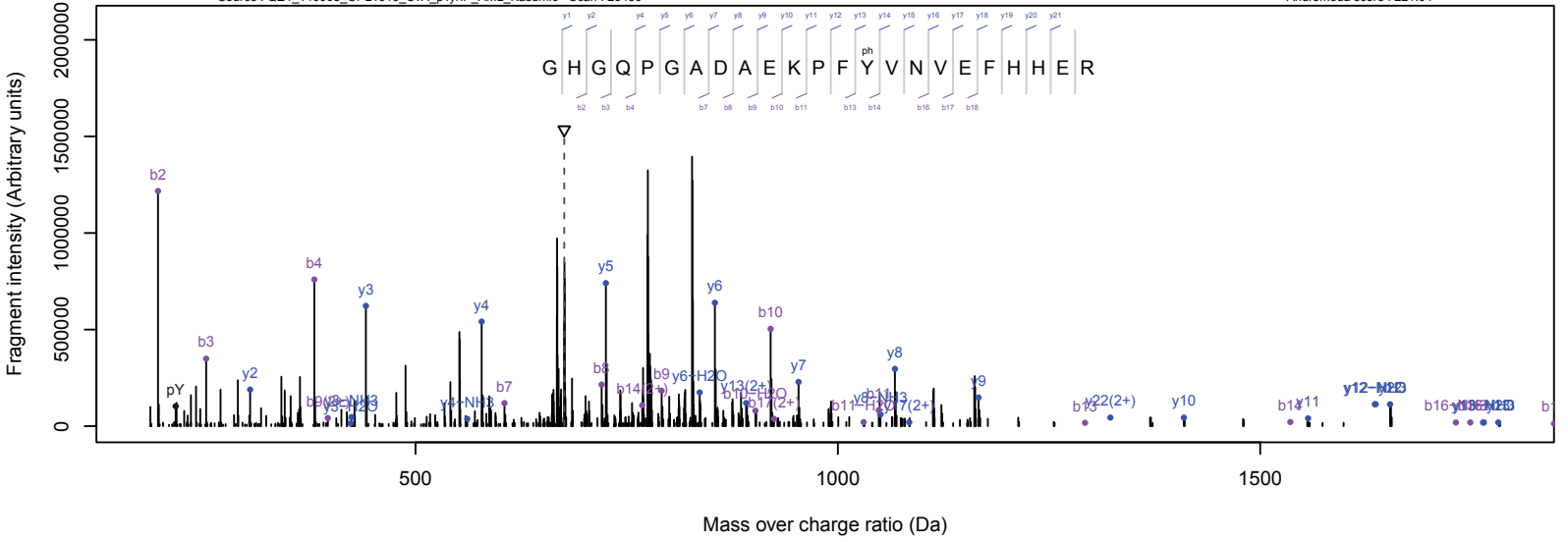
Andromeda score: 221.09



**GHGQPGADAEKPFY(ph)VNVEFHER-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 23450

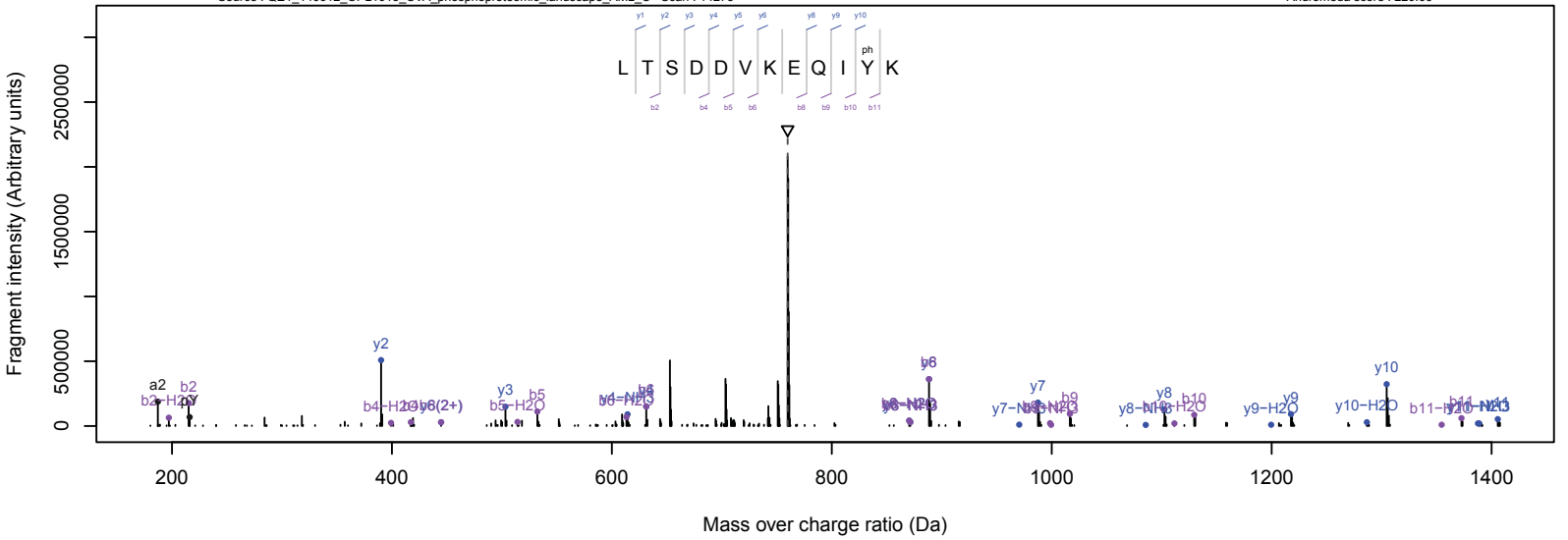
Andromeda score : 221.04



**LTSDDVKEIQY(ph)K-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14273

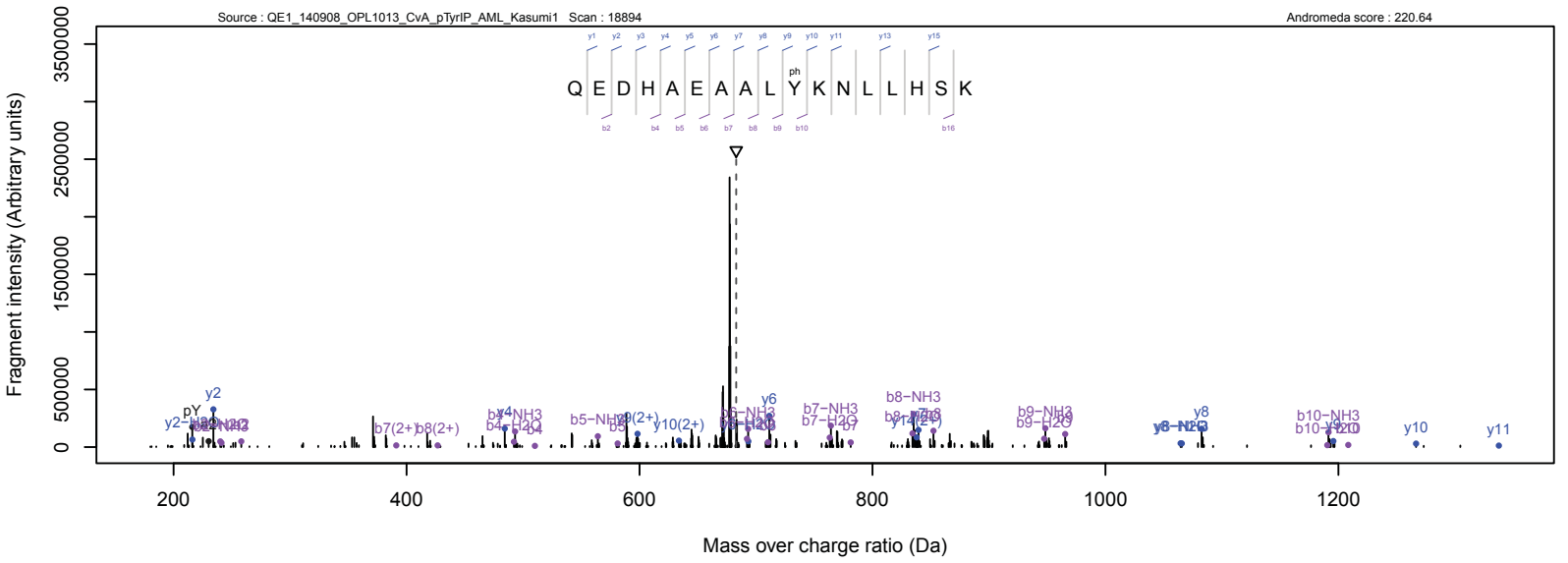
Andromeda score : 220.66



QEDHAEALYK<sup>ph</sup>NLLHSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 18894

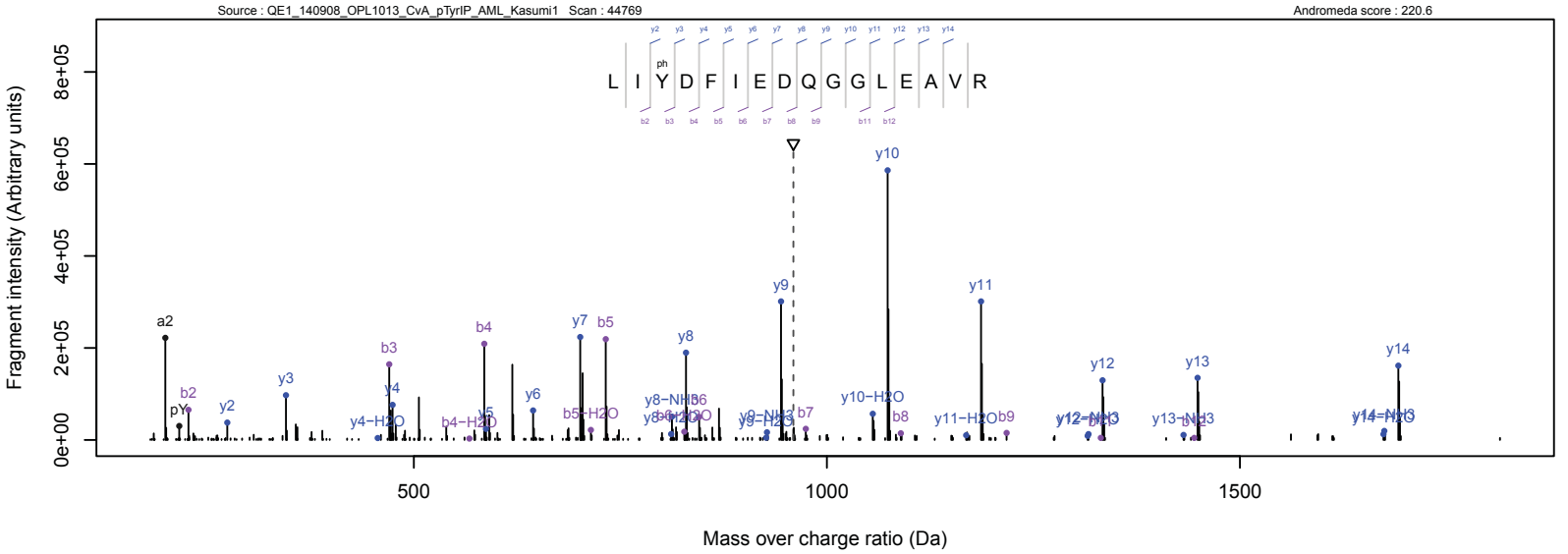
Andromeda score : 220.64



LIY<sup>ph</sup>DFIEDQGGLEAVR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 44769

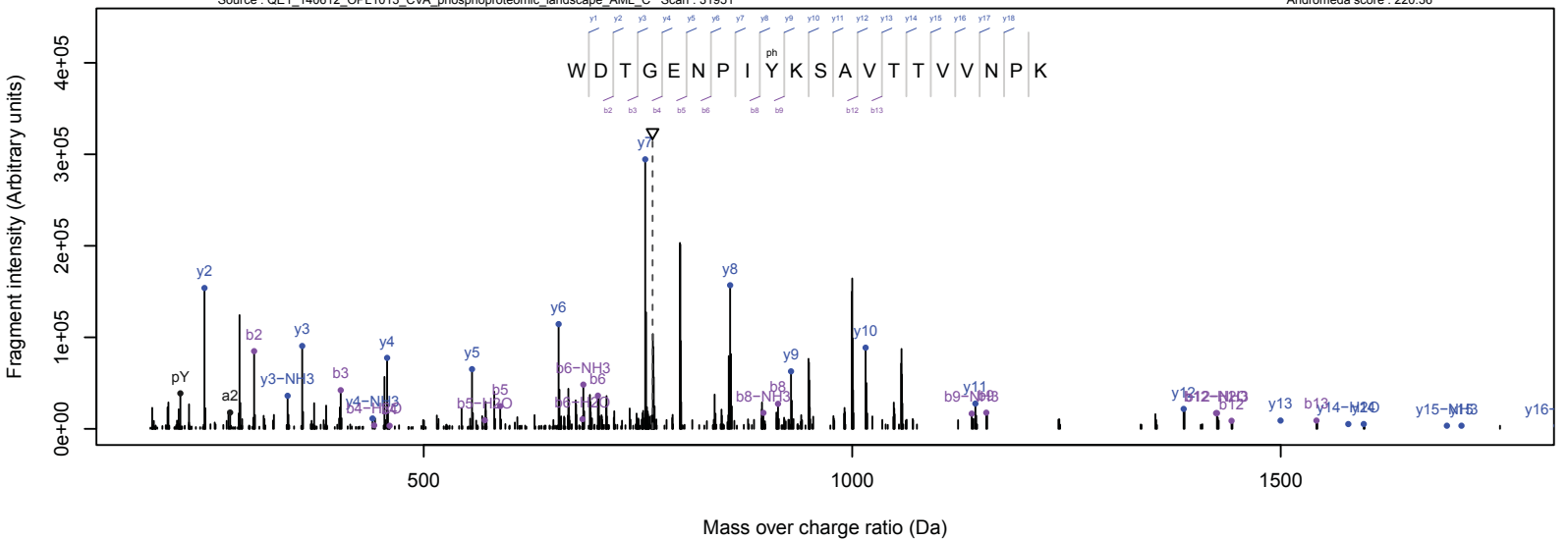
Andromeda score : 220.6



WDTGENPIY<sup>ph</sup>KSAVTTVVNPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31951

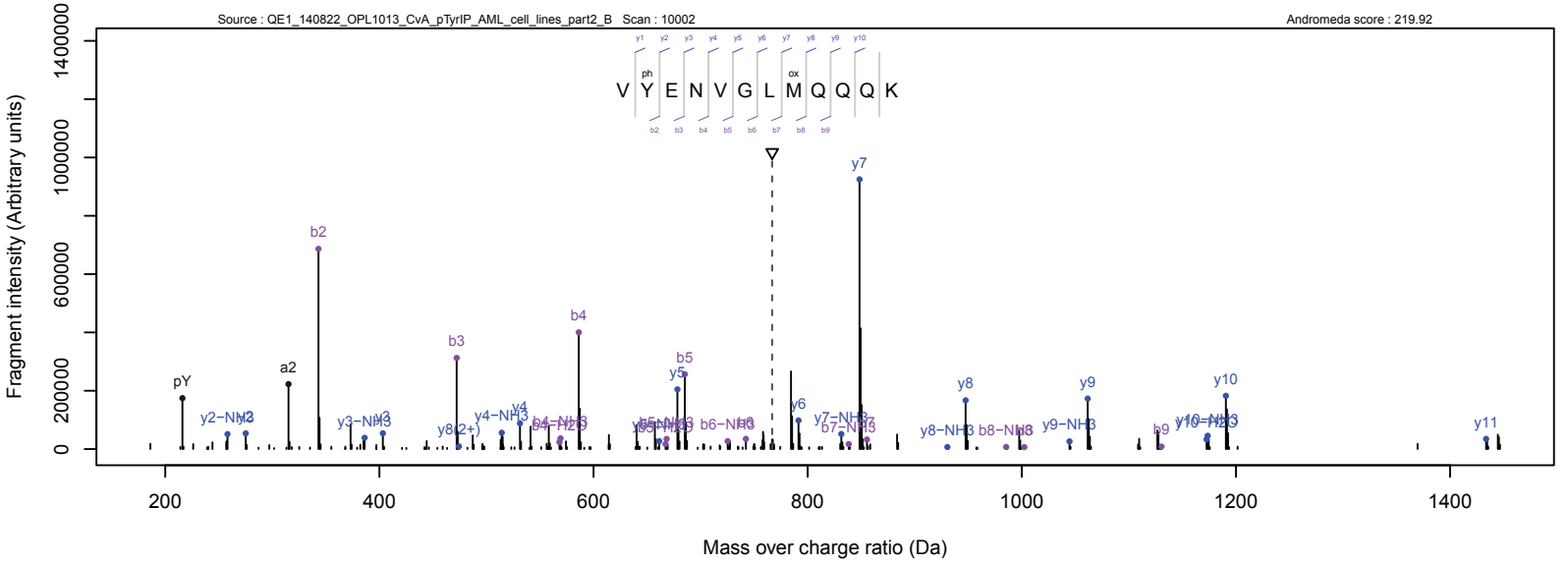
Andromeda score : 220.36



VY(ph)ENGLM(ox)QQK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 10002

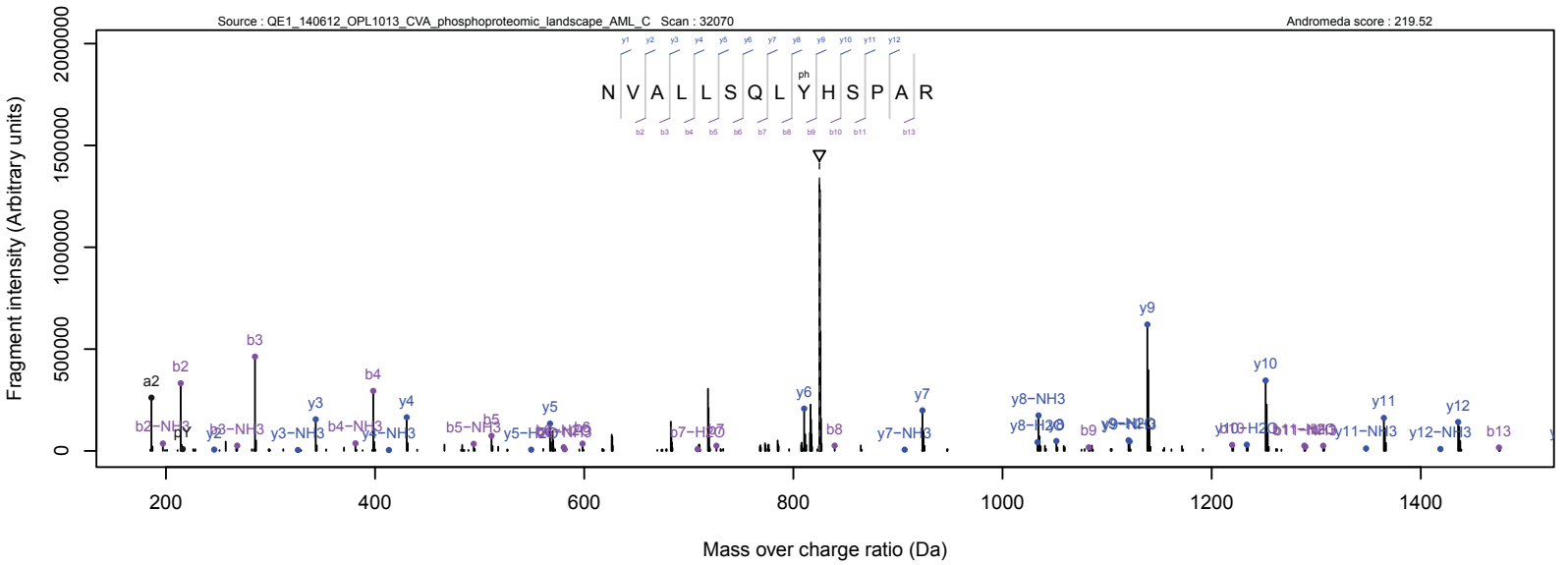
Andromeda score : 219.92



NVALLSQLY(ph)HSPAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32070

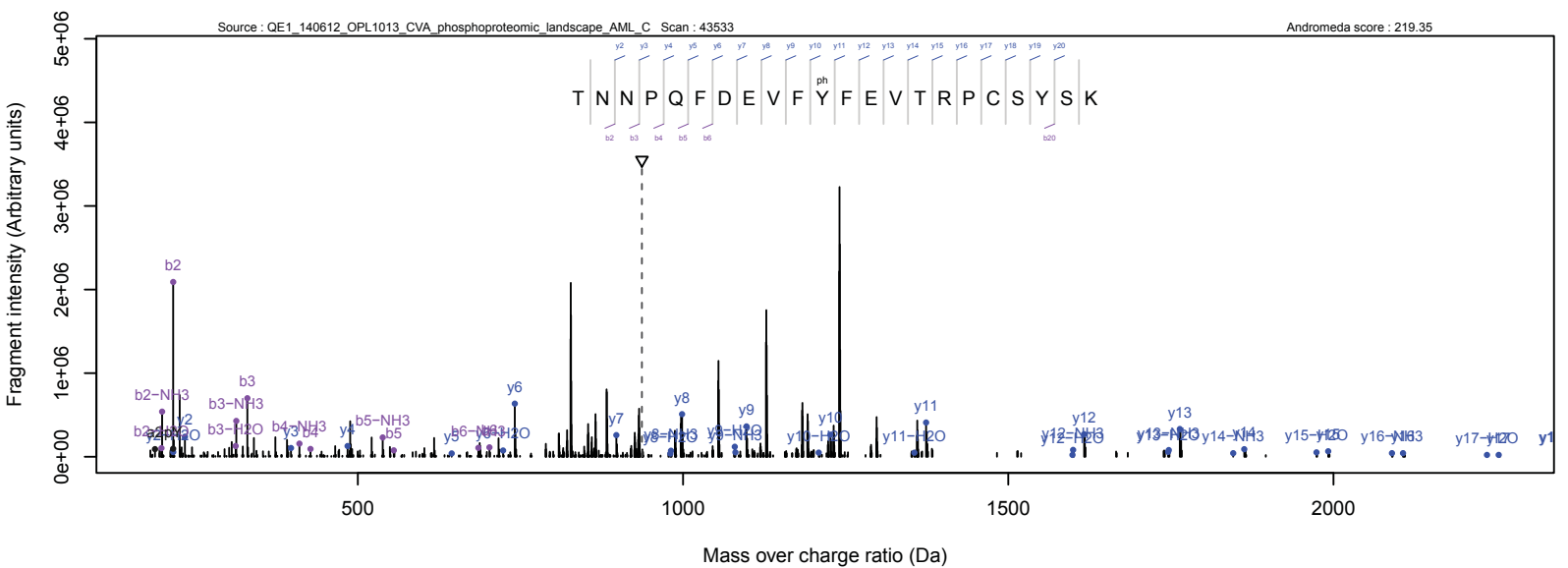
Andromeda score : 219.52



TNNPQFDEVF(ph)FEVTRPCSYSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43533

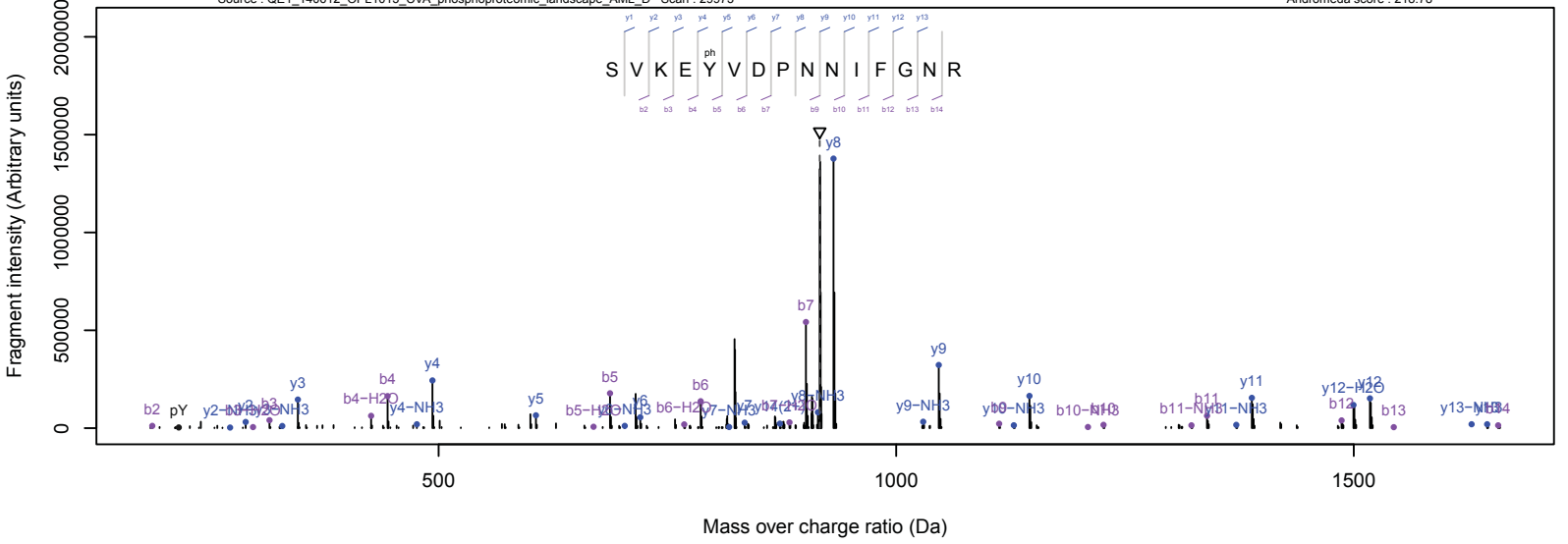
Andromeda score : 219.35



SVKEY(ph)VDPNNIFGNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 29973

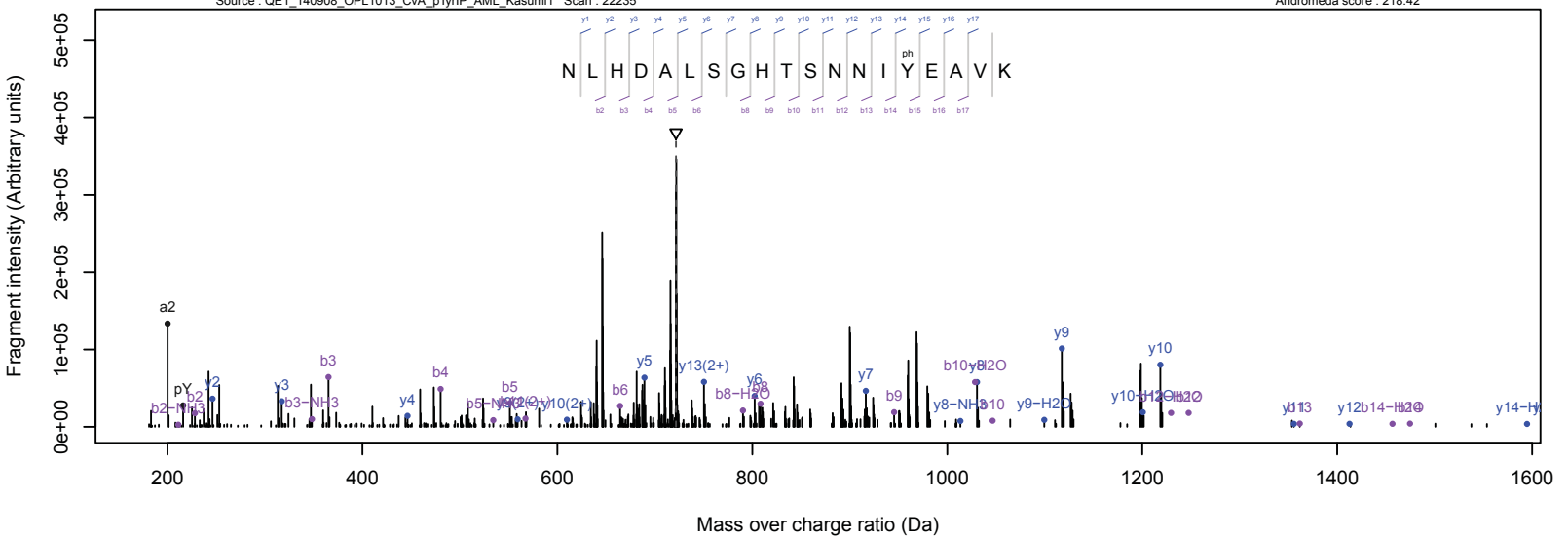
Andromeda score : 218.78



NLHDALSGHTSNNI(ph)EAVK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 22235

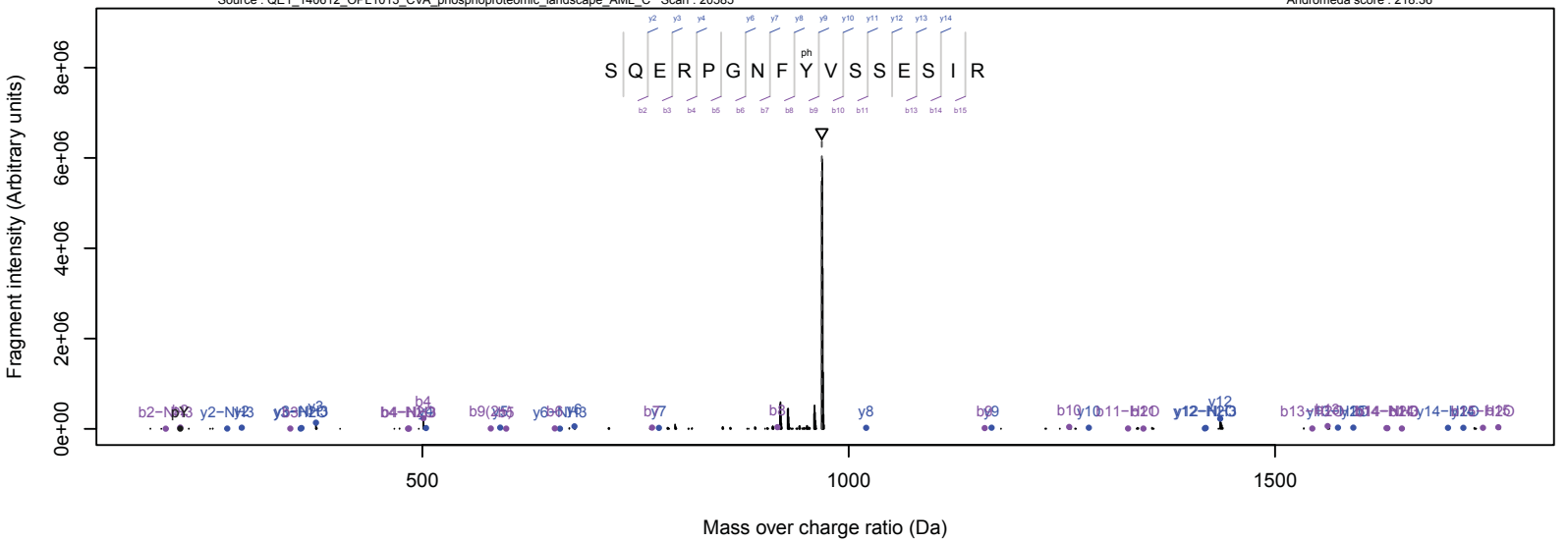
Andromeda score : 218.42



SQERPGNFYVSSESISIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20585

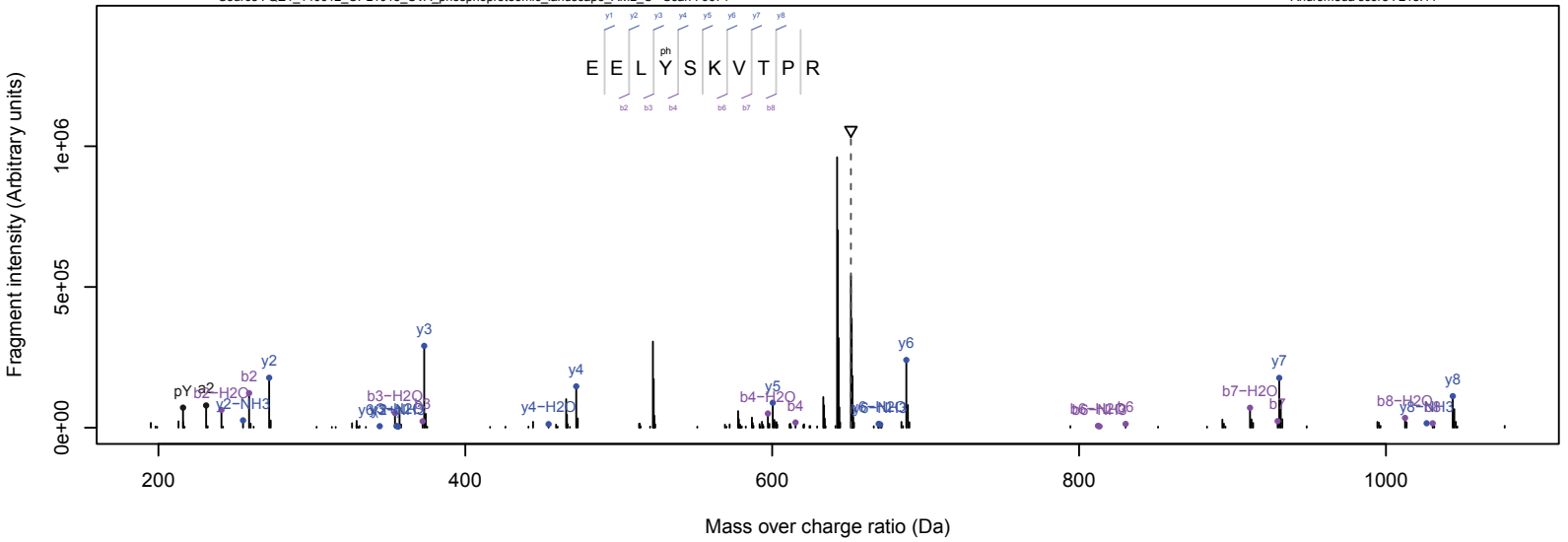
Andromeda score : 218.36



**EELY(ph)SKVTPR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 9671

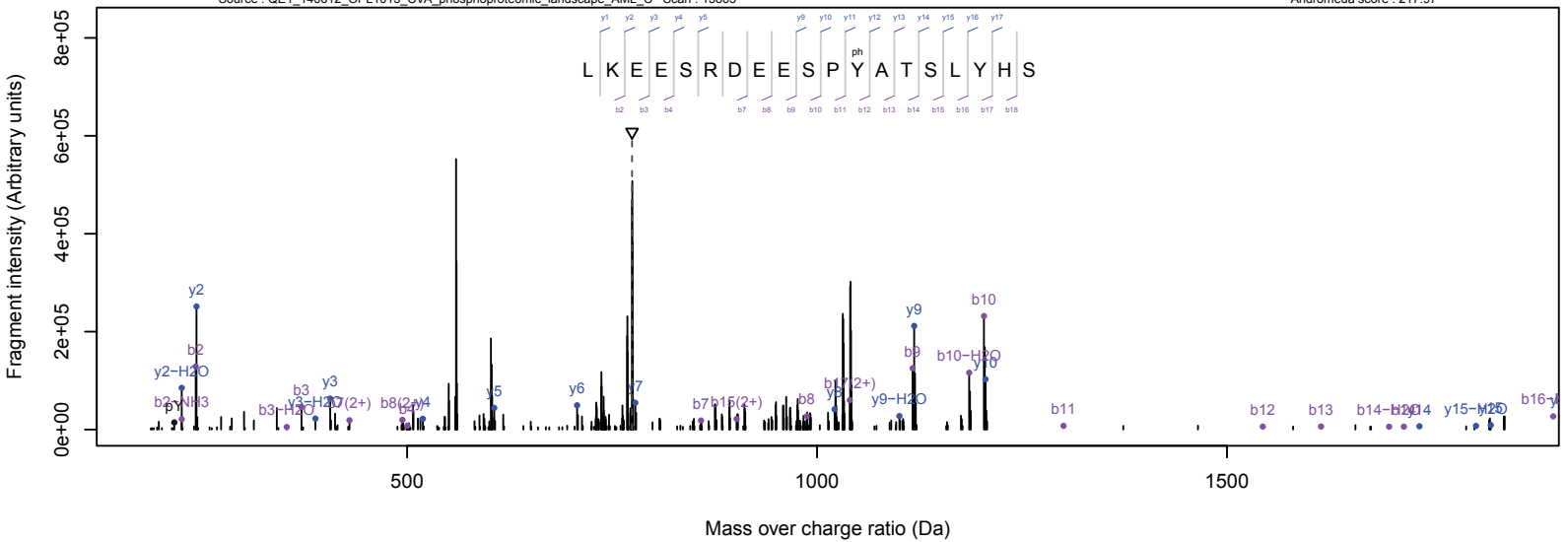
Andromeda score: 218.11



**LKEESRDEESPY(ph)ATSLYHS-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13865

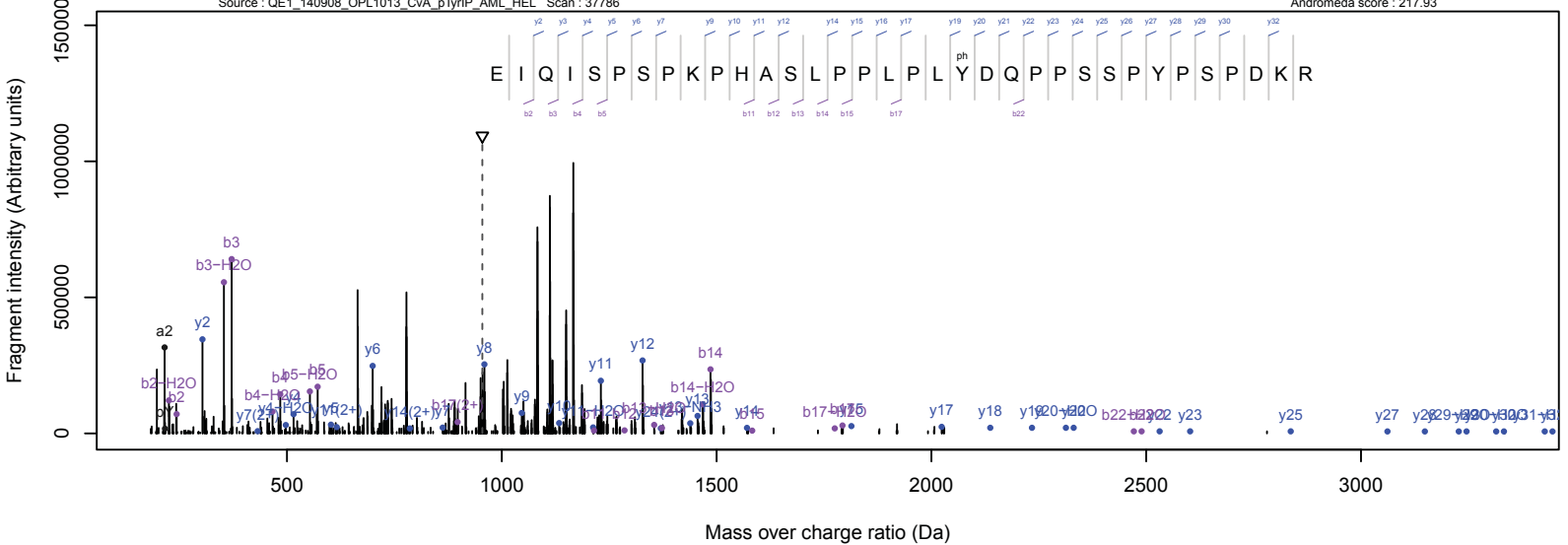
Andromeda score: 217.97



**EIQISPSKPHASLPPLPPLY(ph)DQPPSSPYSPDKR-4+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 37786

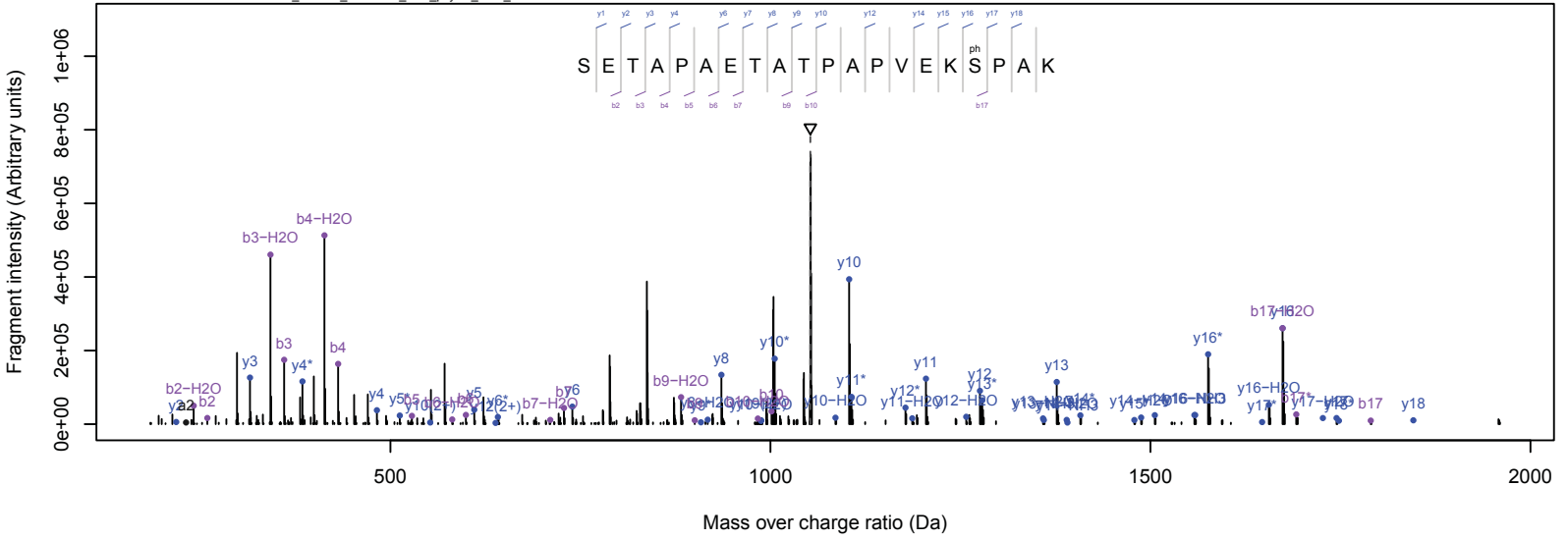
Andromeda score: 217.93



(ac)SETAPAETATPAPVEK(ph)PAK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_ML2 Scan : 16927

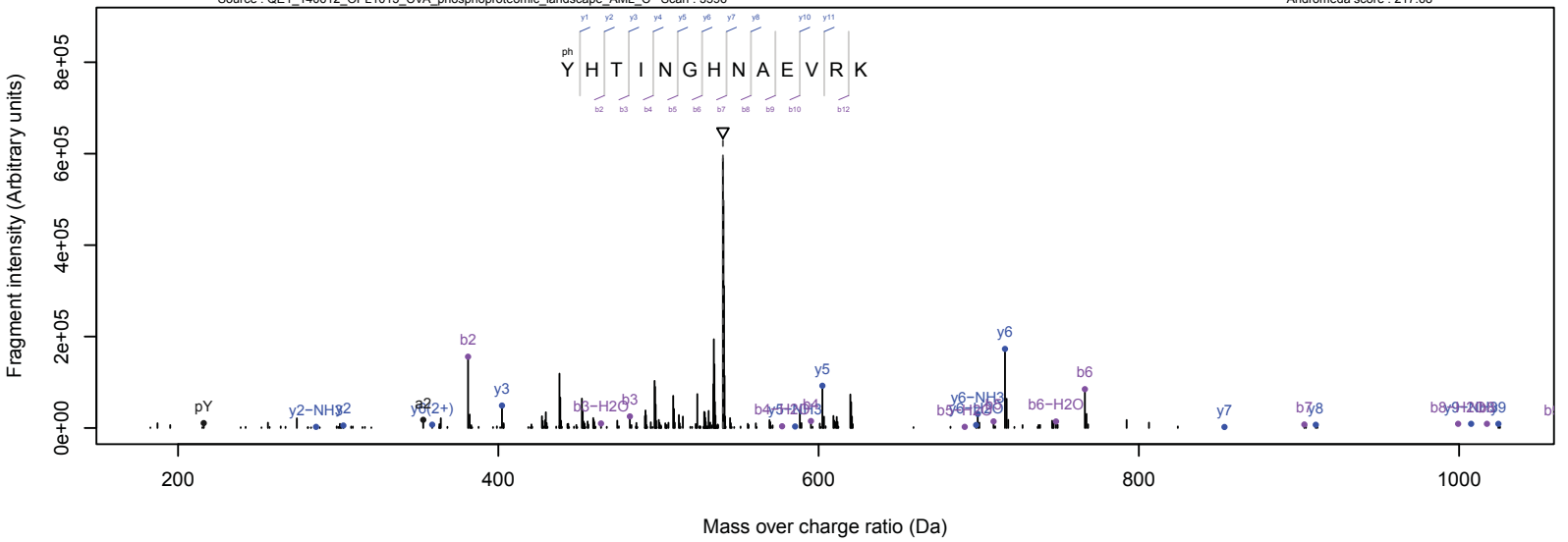
Andromeda score : 217.89



Y(ph)HTINGHNAEVRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3596

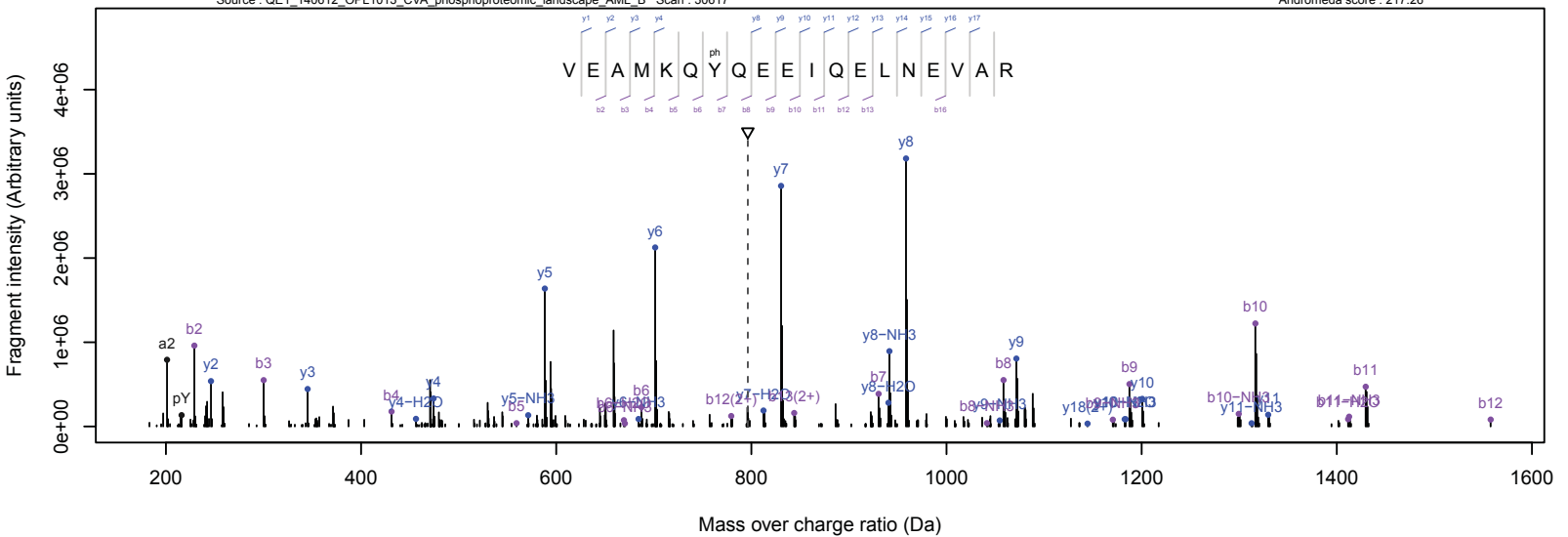
Andromeda score : 217.68



VEAMKQY(ph)QEEIQELNEVAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30617

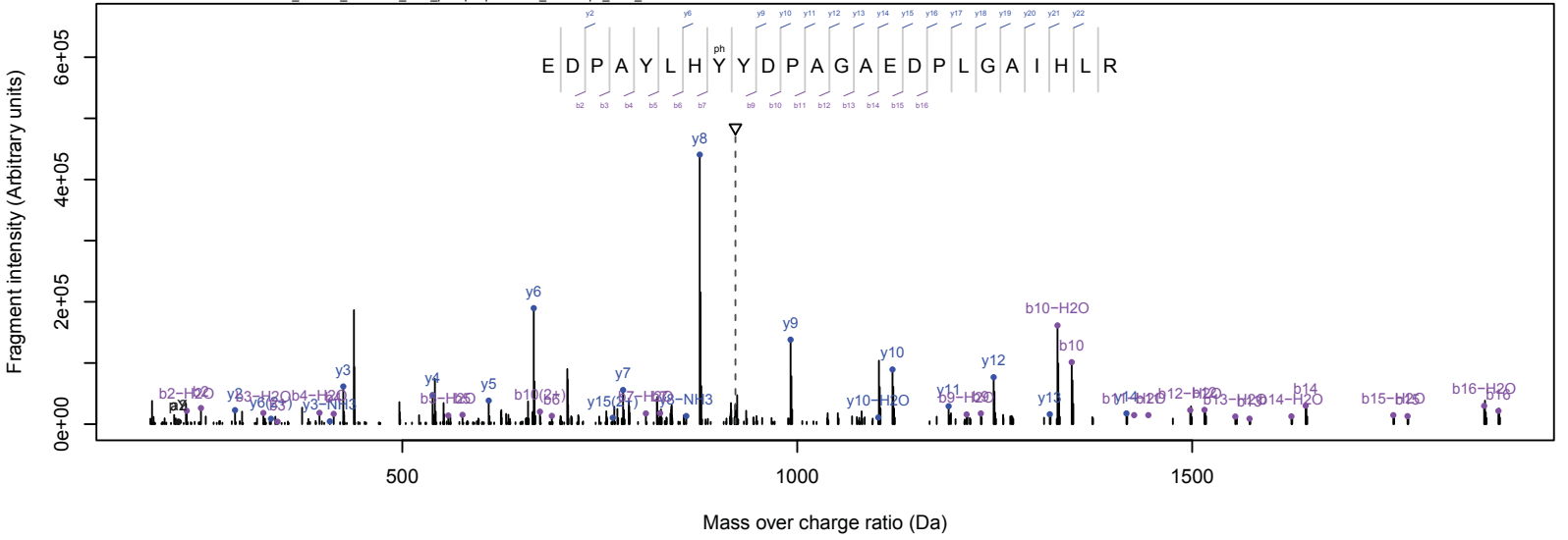
Andromeda score : 217.26



EDPAYLHY(ph)YDPAGAEDPLGAIHLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43838

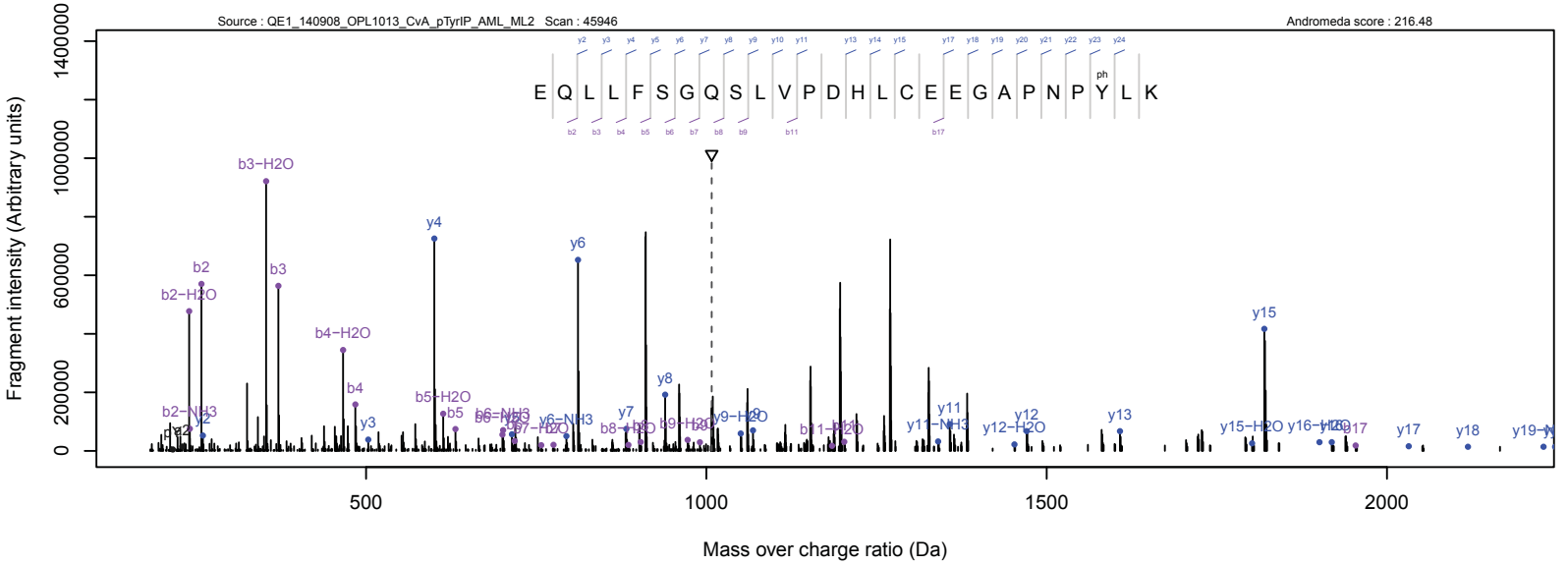
Andromeda score : 216.75



EQLLFSGQSLVPDHLCEEGAPNPY(ph)LK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 45946

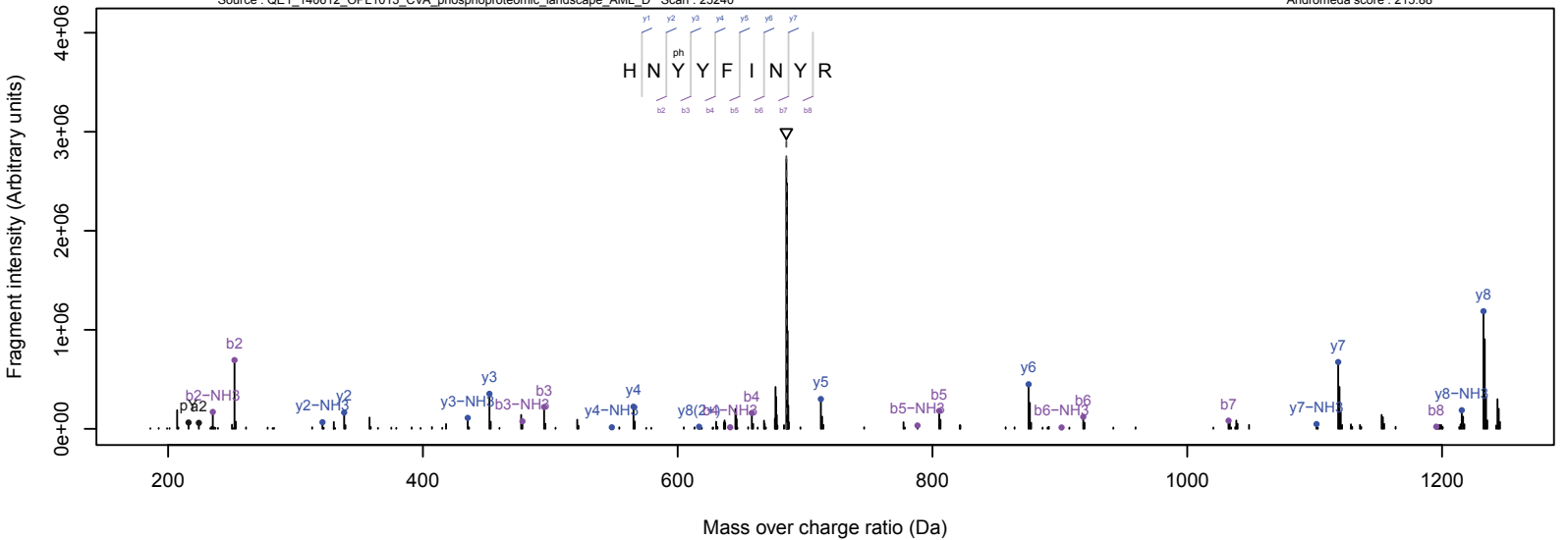
Andromeda score : 216.48



HNY(ph)YFINYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 25240

Andromeda score : 215.88

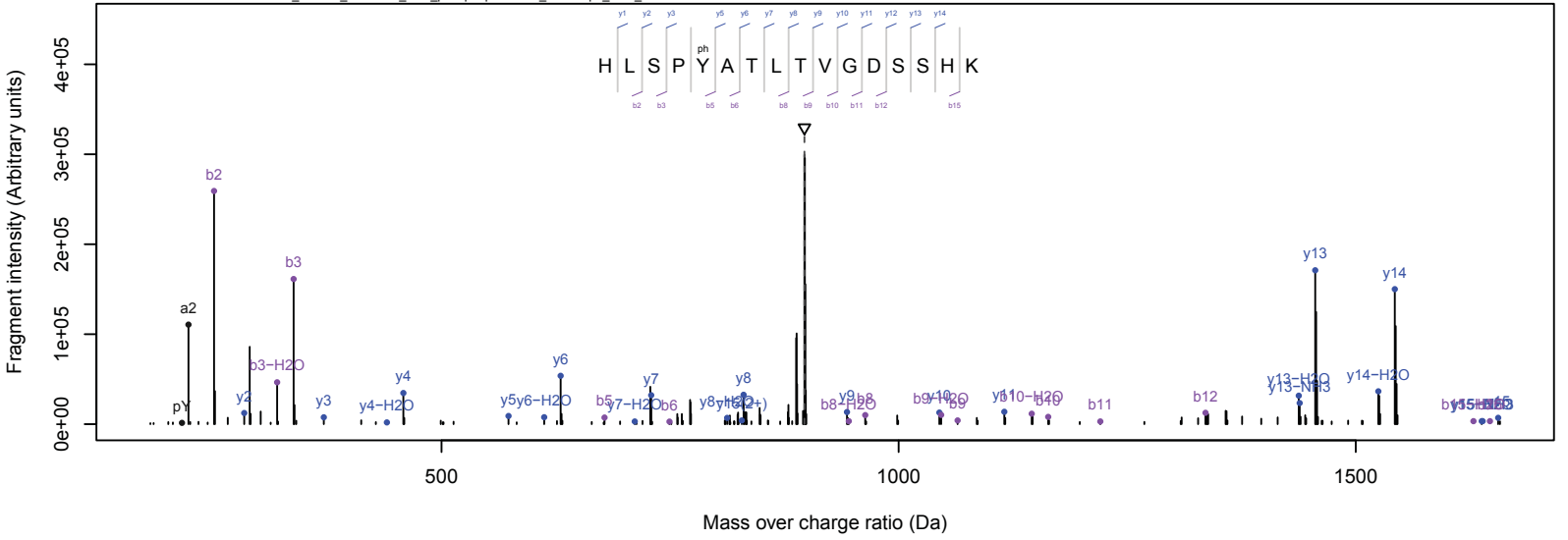




HLSPY(ph)ATLTVGDSSHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17436

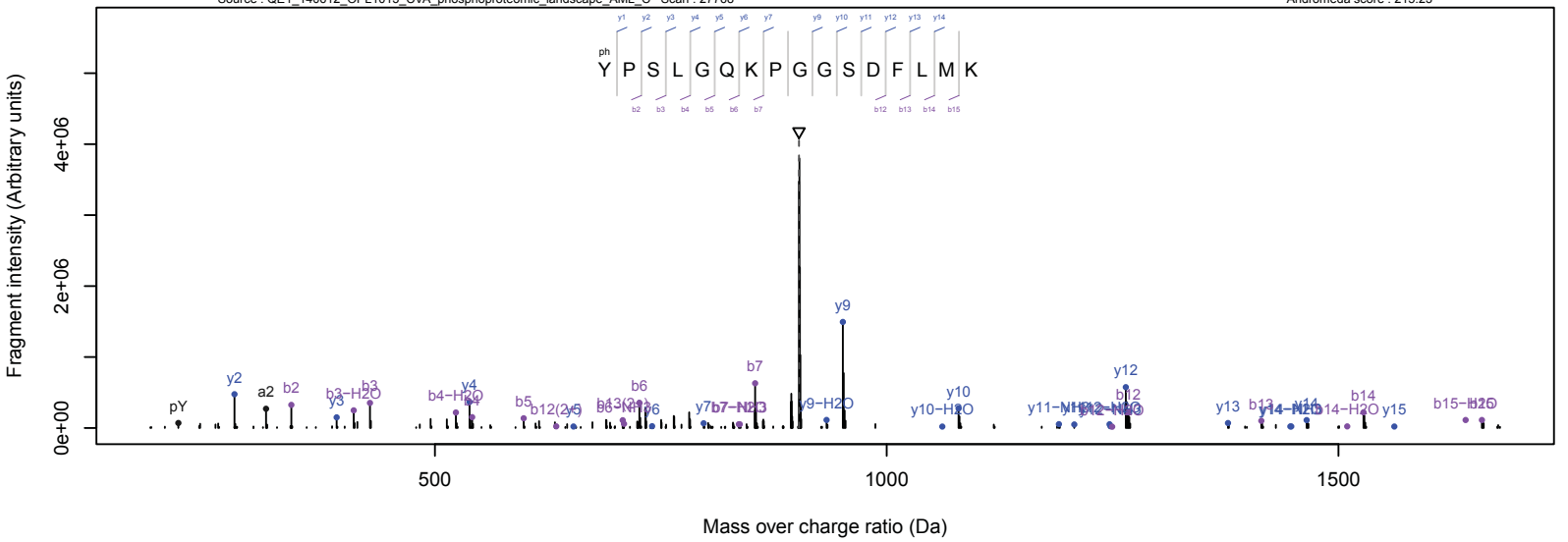
Andromeda score : 215.48



Y(ph)PSLGQKPGSDFLMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27768

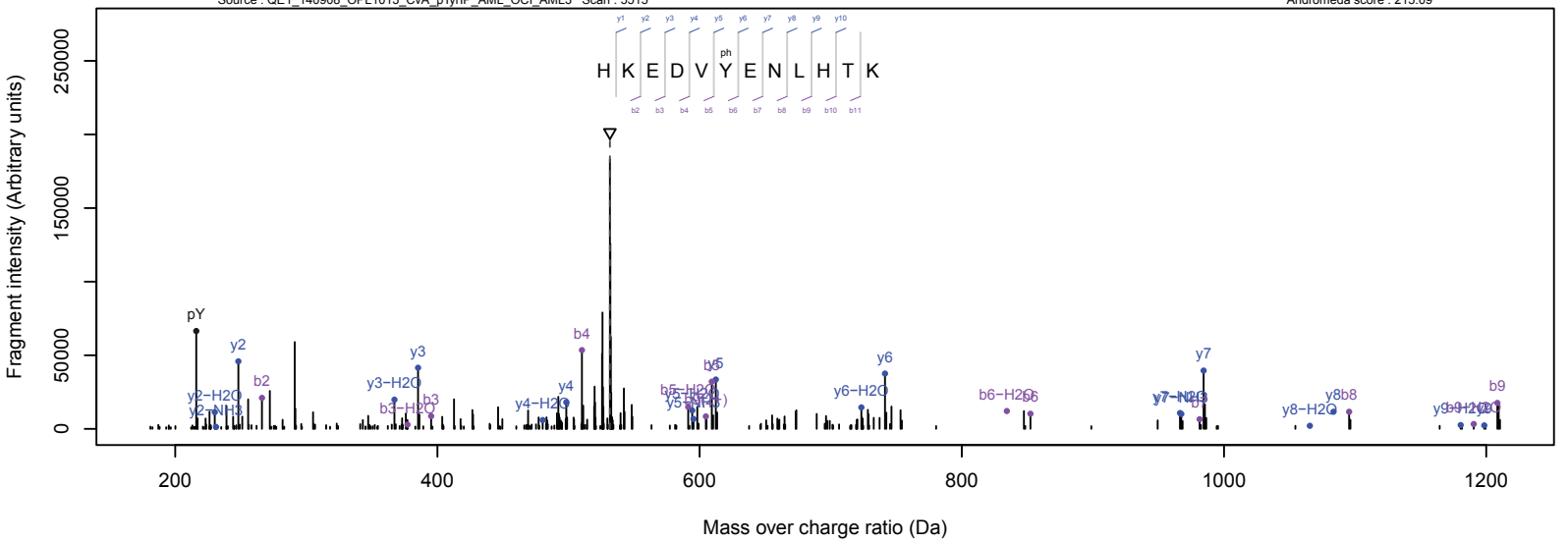
Andromeda score : 215.23



HKEDVY(ph)ENLHTK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 5515

Andromeda score : 215.09

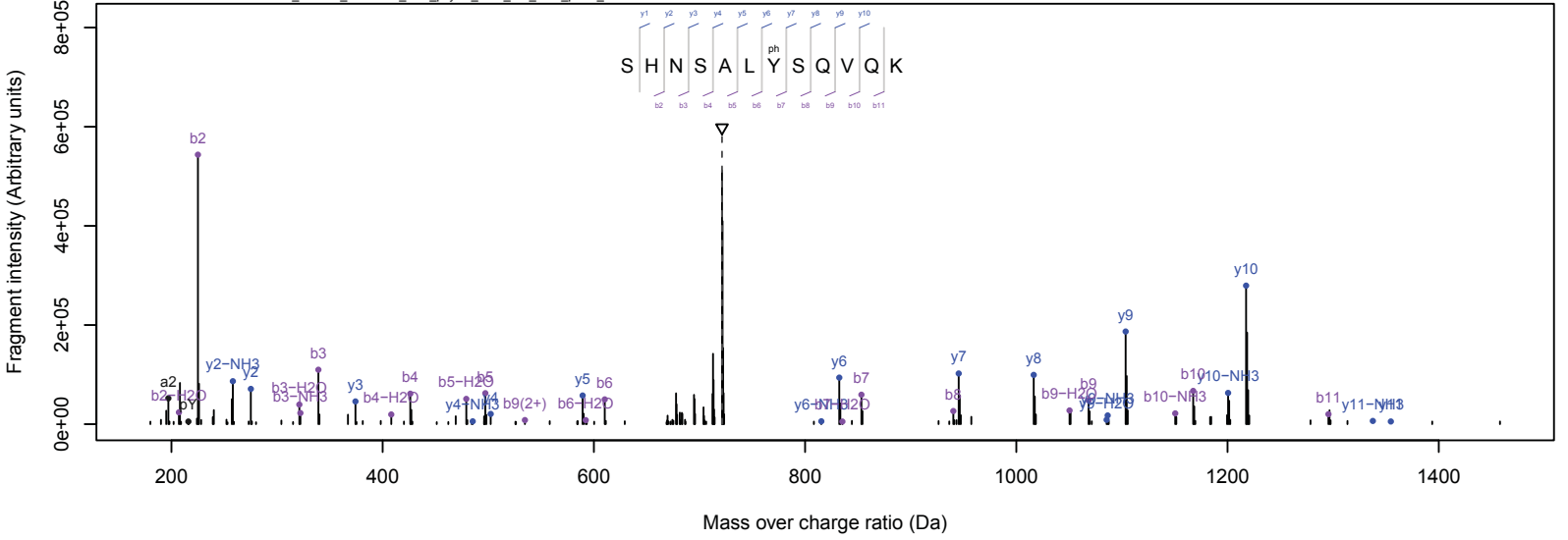




SHNSALY(ph)SQVQK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 4296

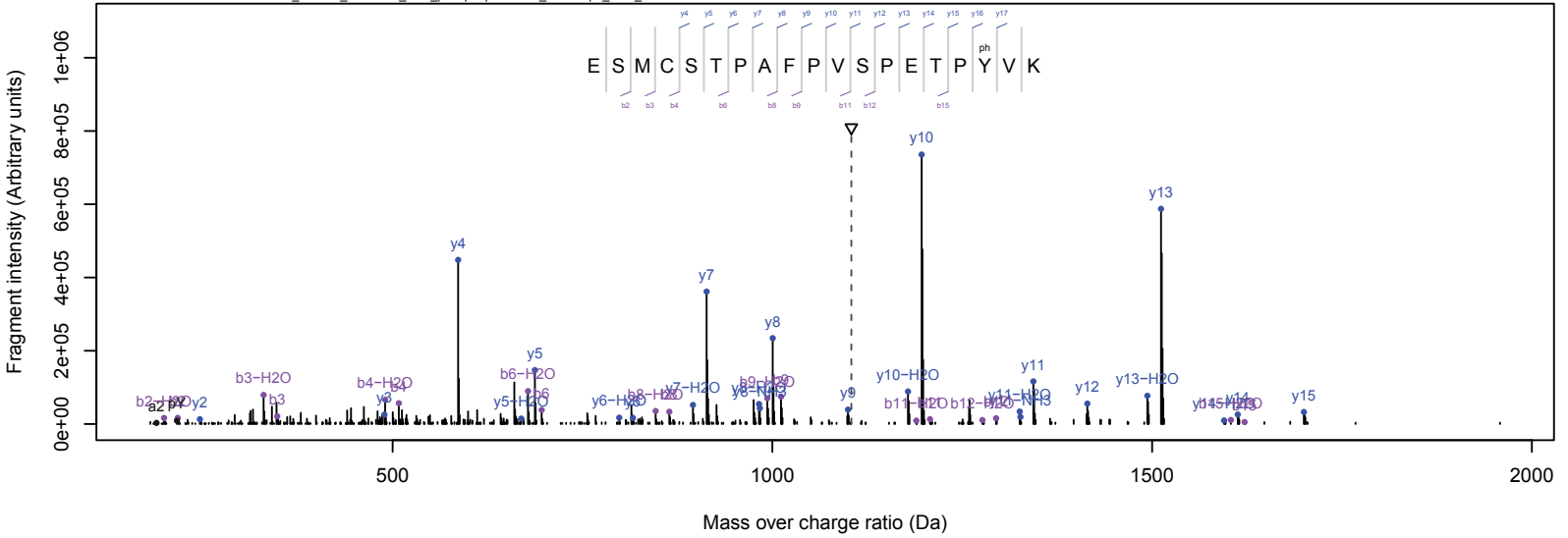
Andromeda score : 214.61



ESMCSTPAFPVSPETPY(ph)VK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34846

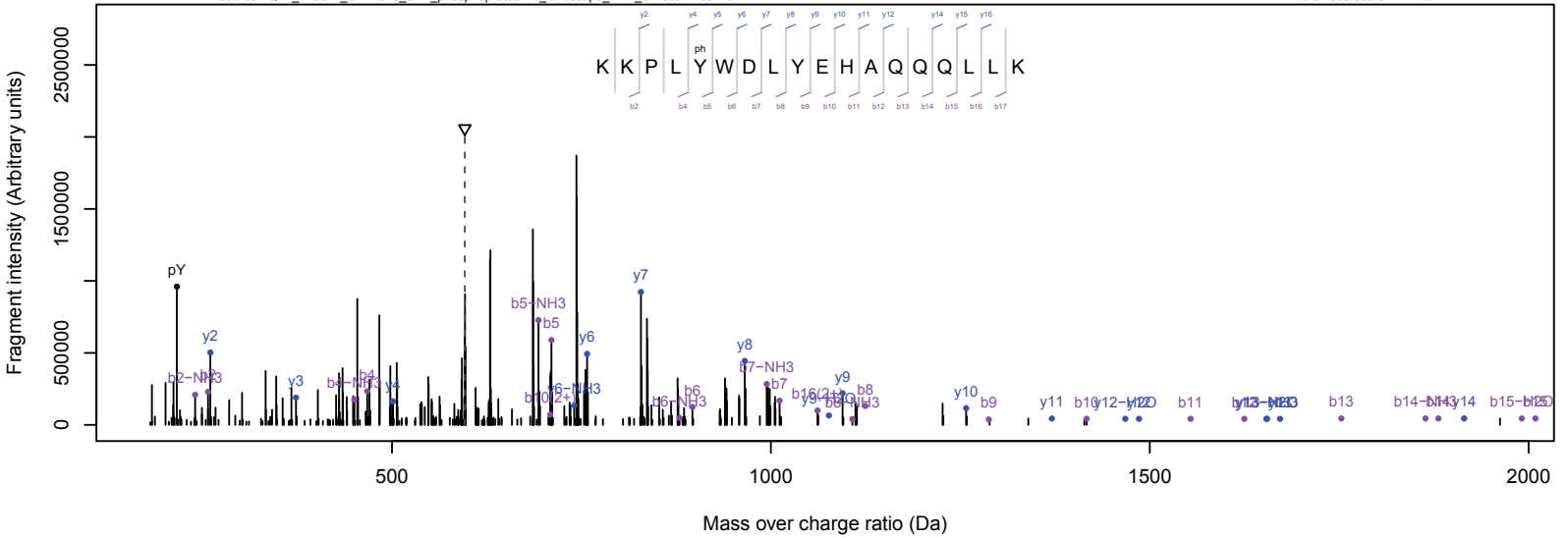
Andromeda score : 214.5



KKPLY(ph)WDLYEHAQQQLLK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35128

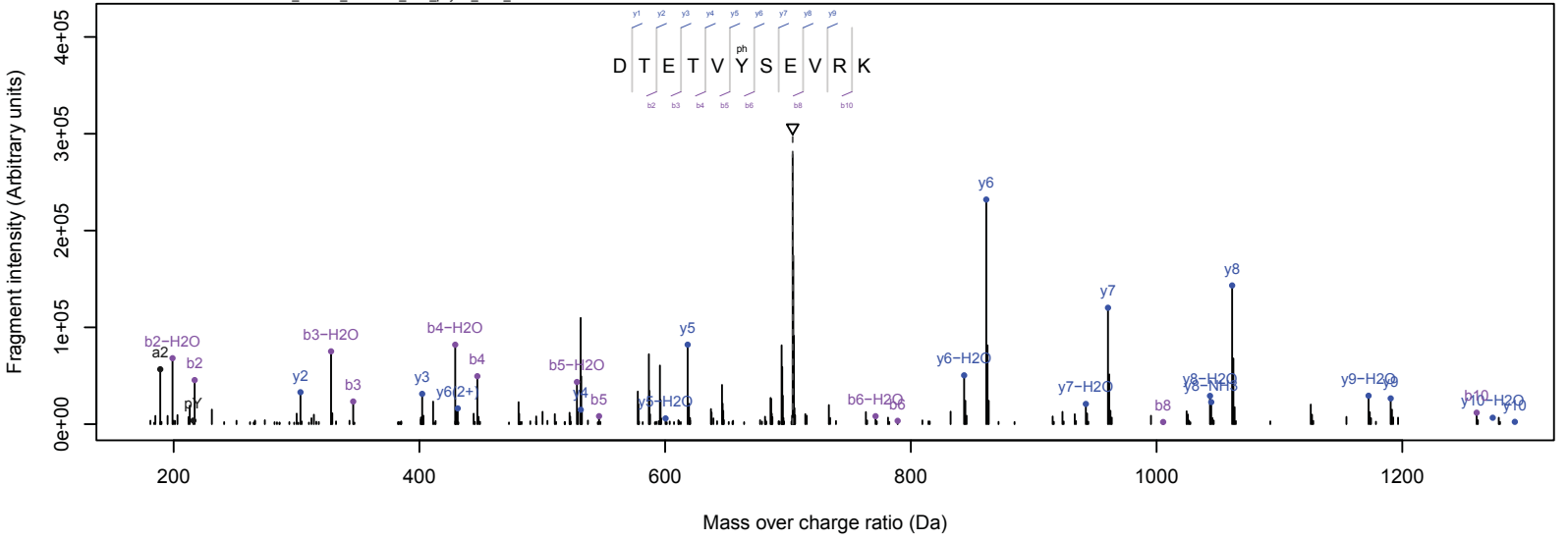
Andromeda score : 214.46



DTETVY(ph)SEVRK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 9741

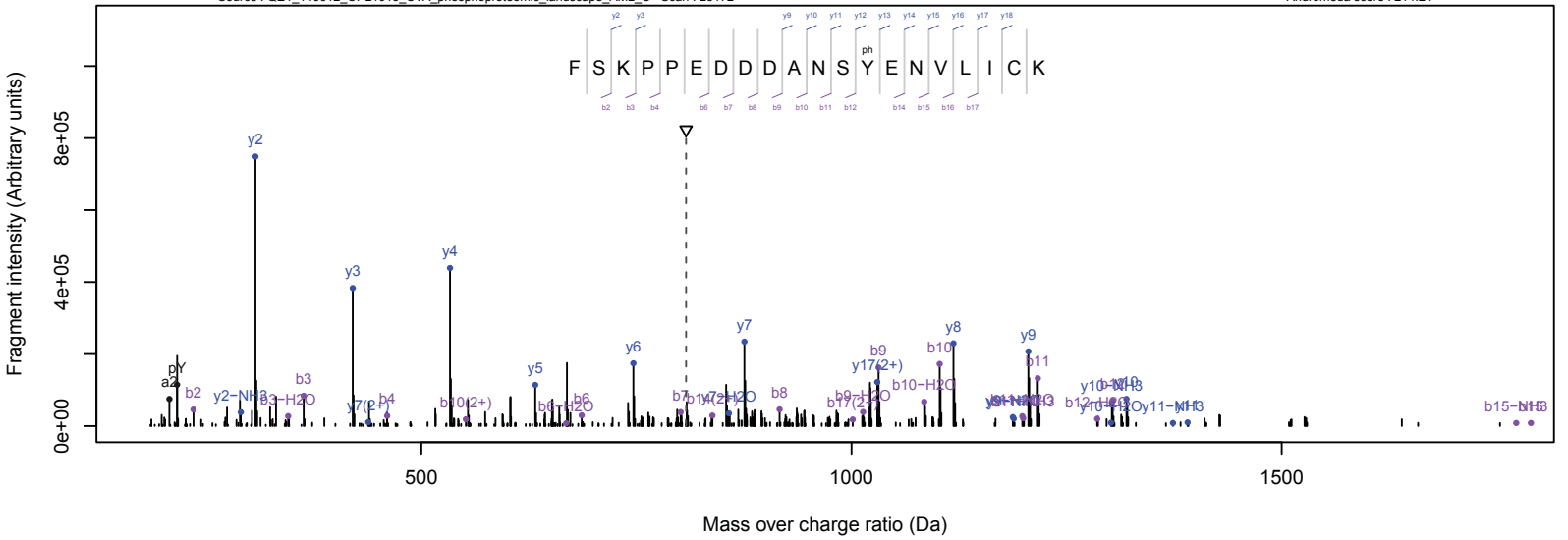
Andromeda score : 214.26



FSKPPEDDANSY(ph)ENVLICK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25172

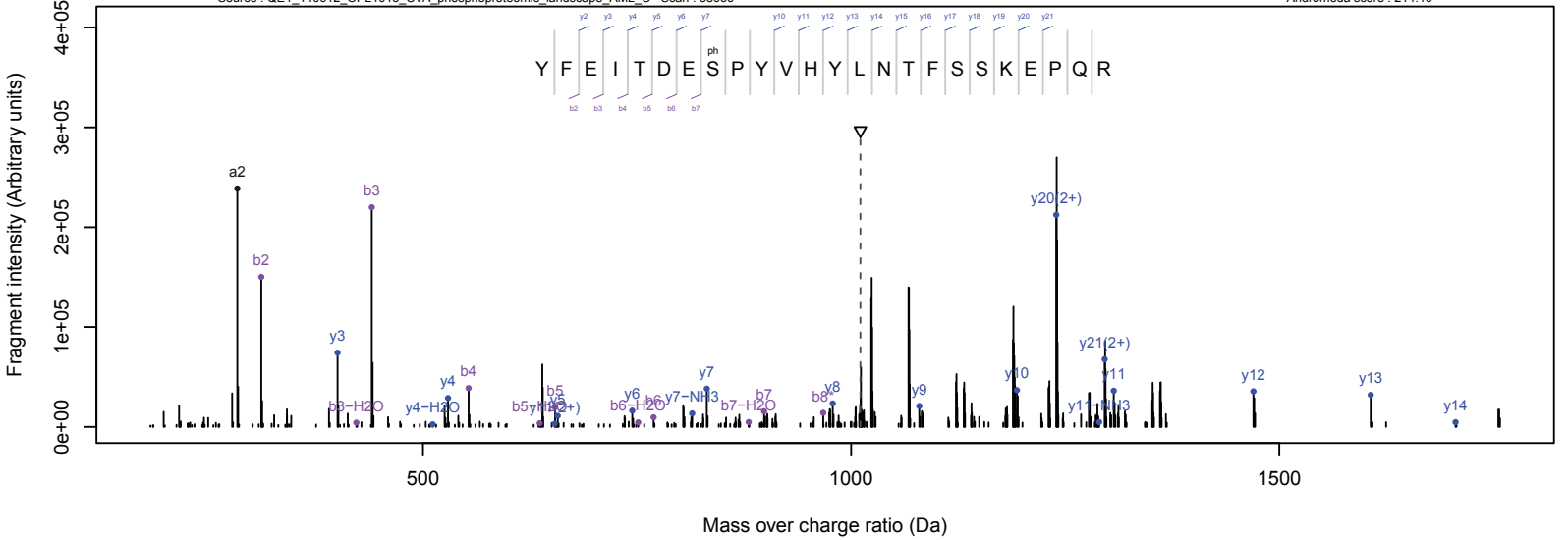
Andromeda score : 214.24



YFEITDES(ph)PYVHYLNTFSSKEPQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38066

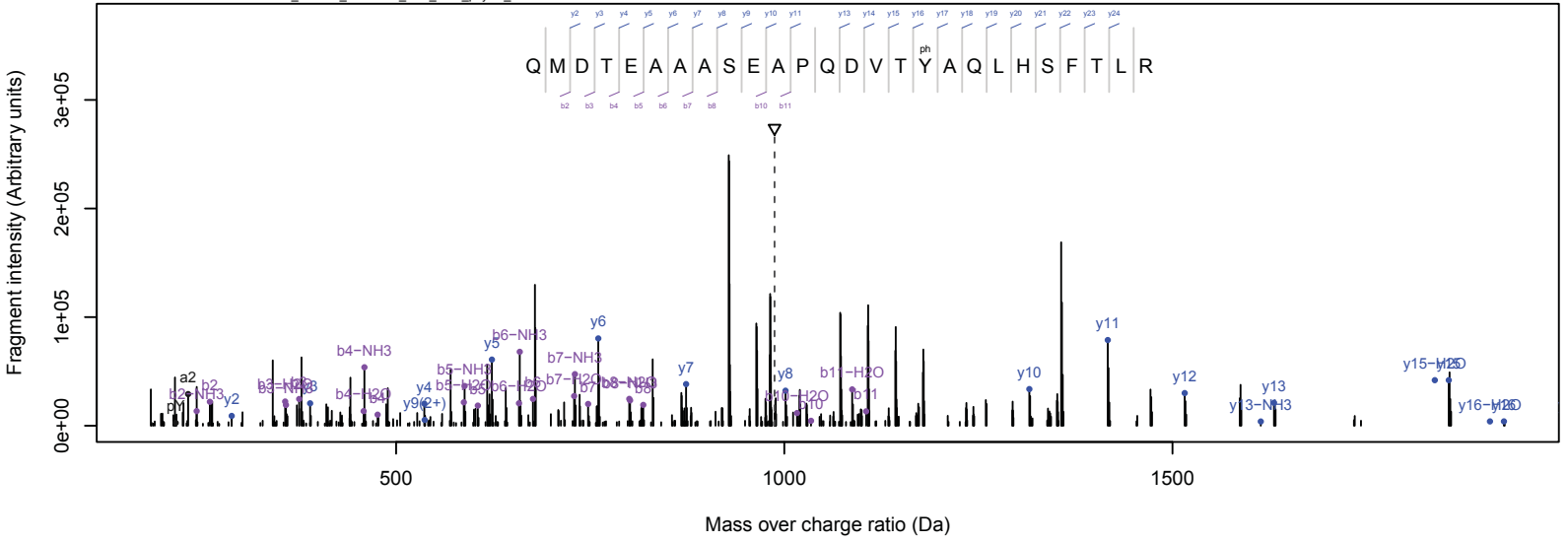
Andromeda score : 214.19



QMDTEAAASEAPQDVTY(ph)AQLHSFTLR-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 37050

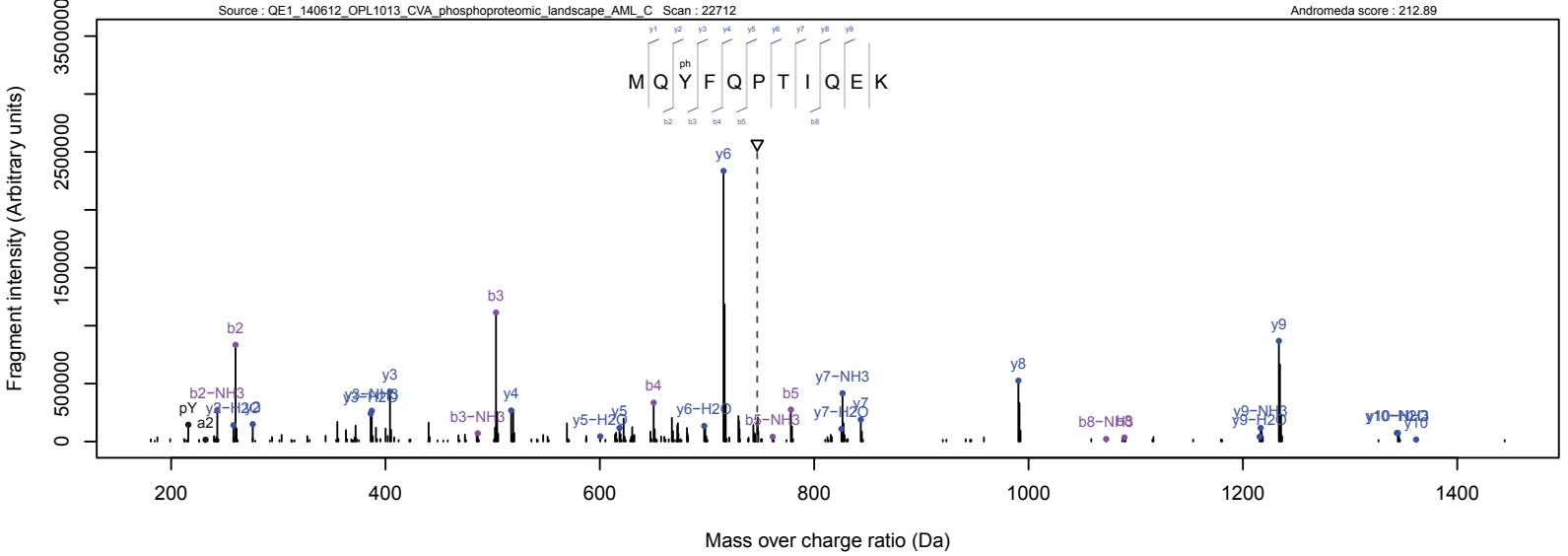
Andromeda score : 214.05



MQY(ph)FQPTIQEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22712

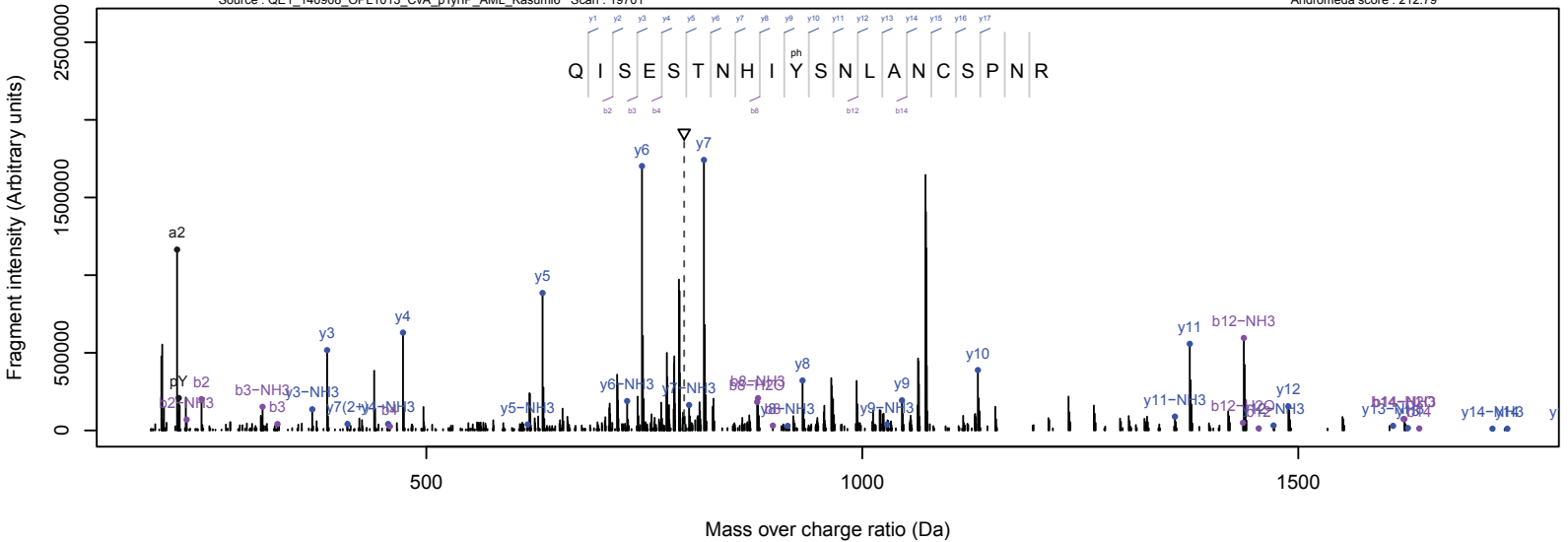
Andromeda score : 212.89



QIESTNHIY(ph)SNLANCSPNR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 19701

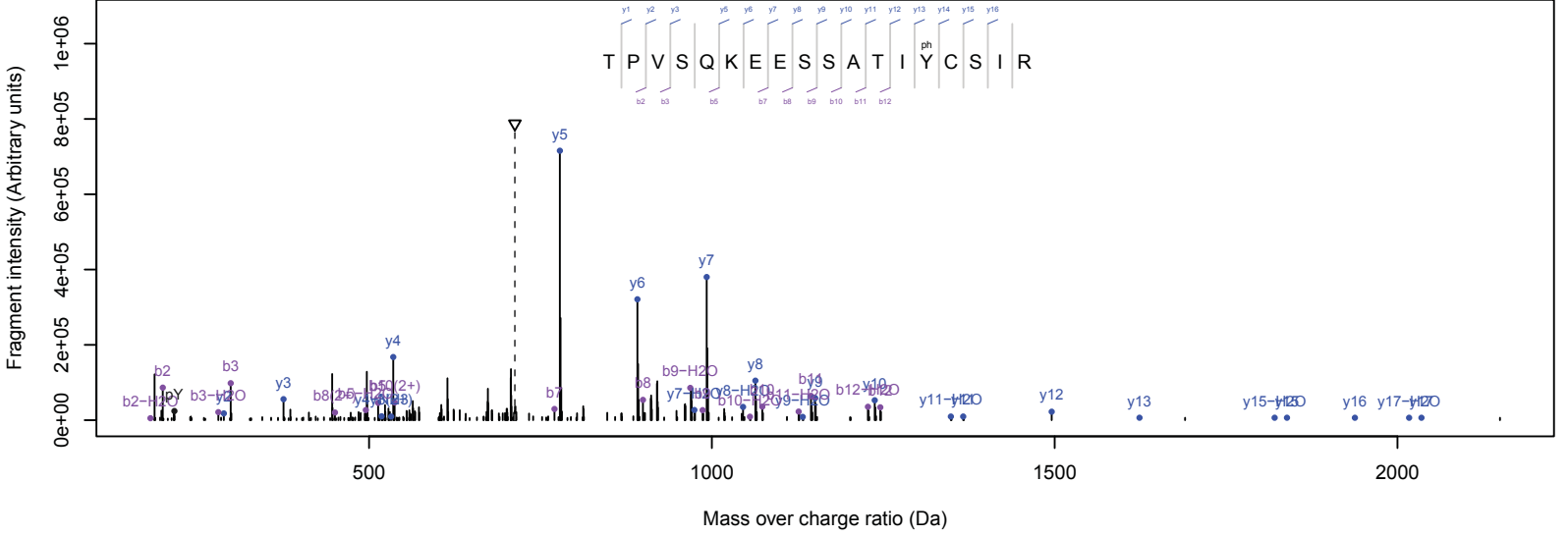
Andromeda score : 212.79



TPVSQKEESSATIY(ph)CSIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16029

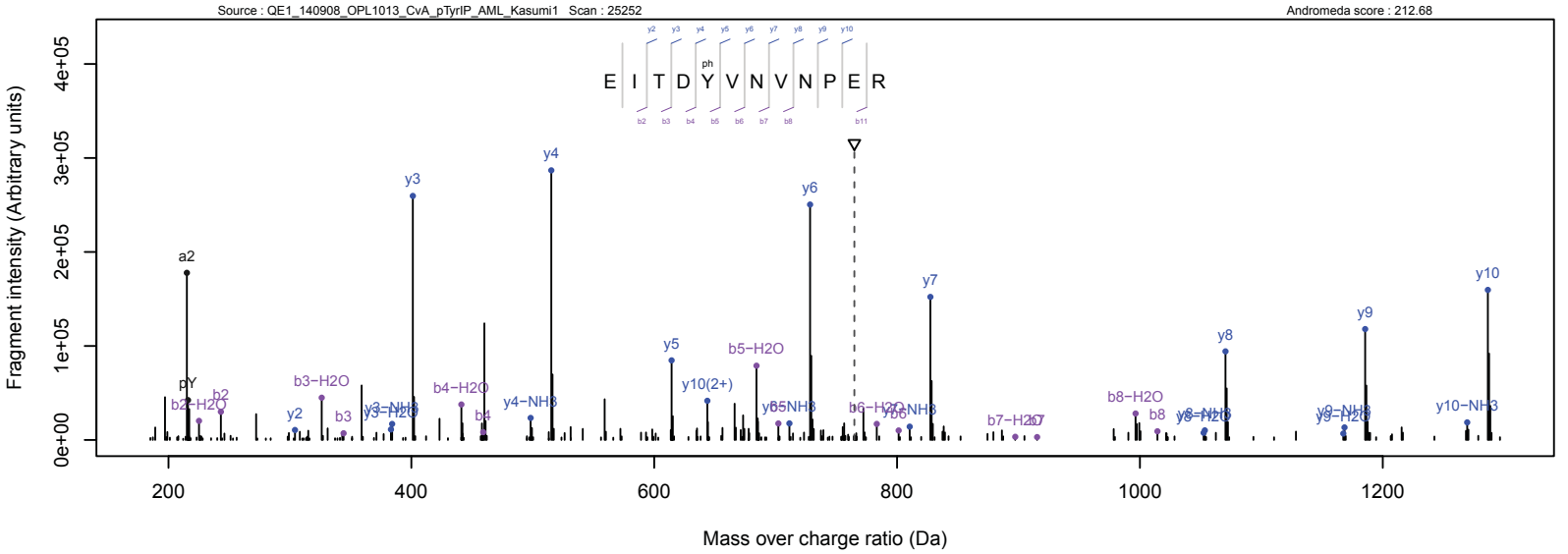
Andromeda score: 212.76



EITDY(ph)VNVNPER-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 25252

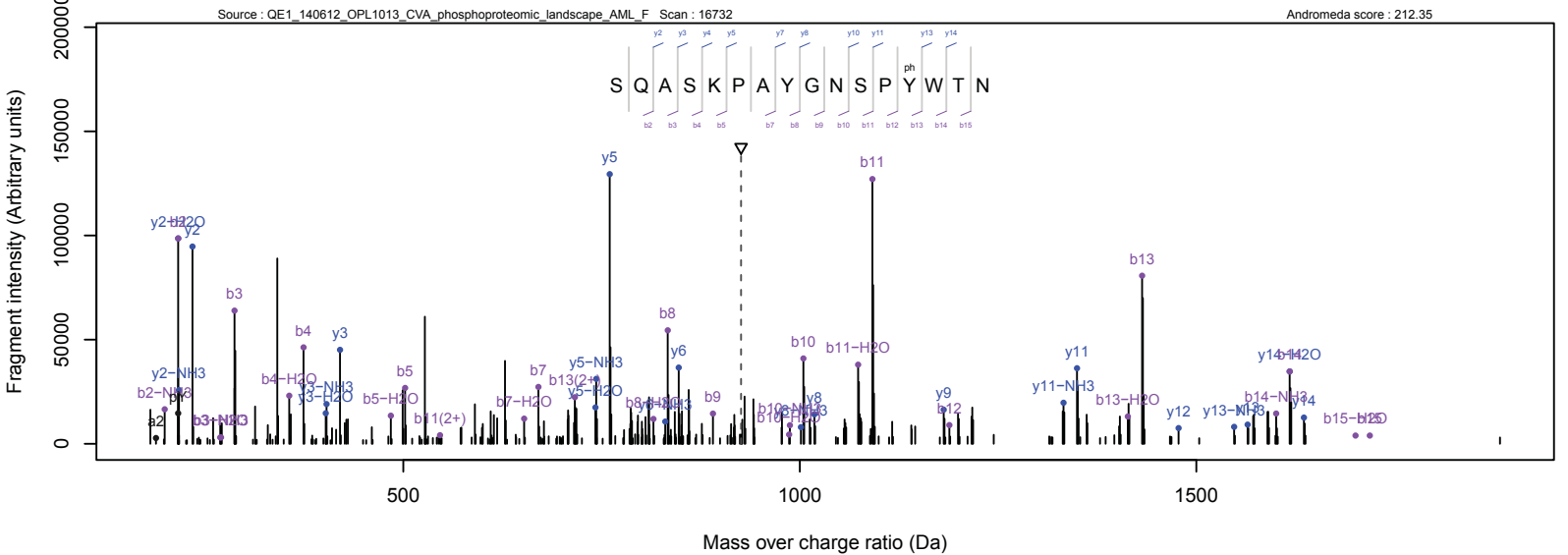
Andromeda score: 212.68



SQASKPAYGNSPY(ph)WTN-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan: 16732

Andromeda score: 212.35

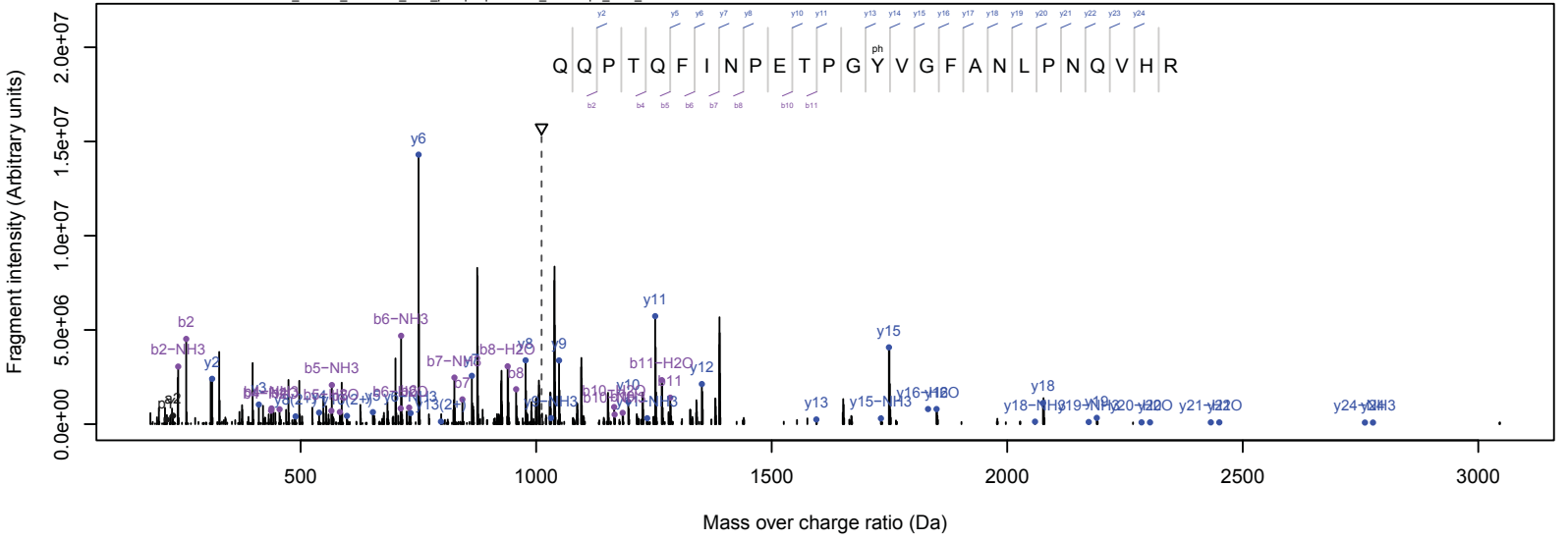




**QQPTQFINPETPGYVGFANLPNQVHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40025

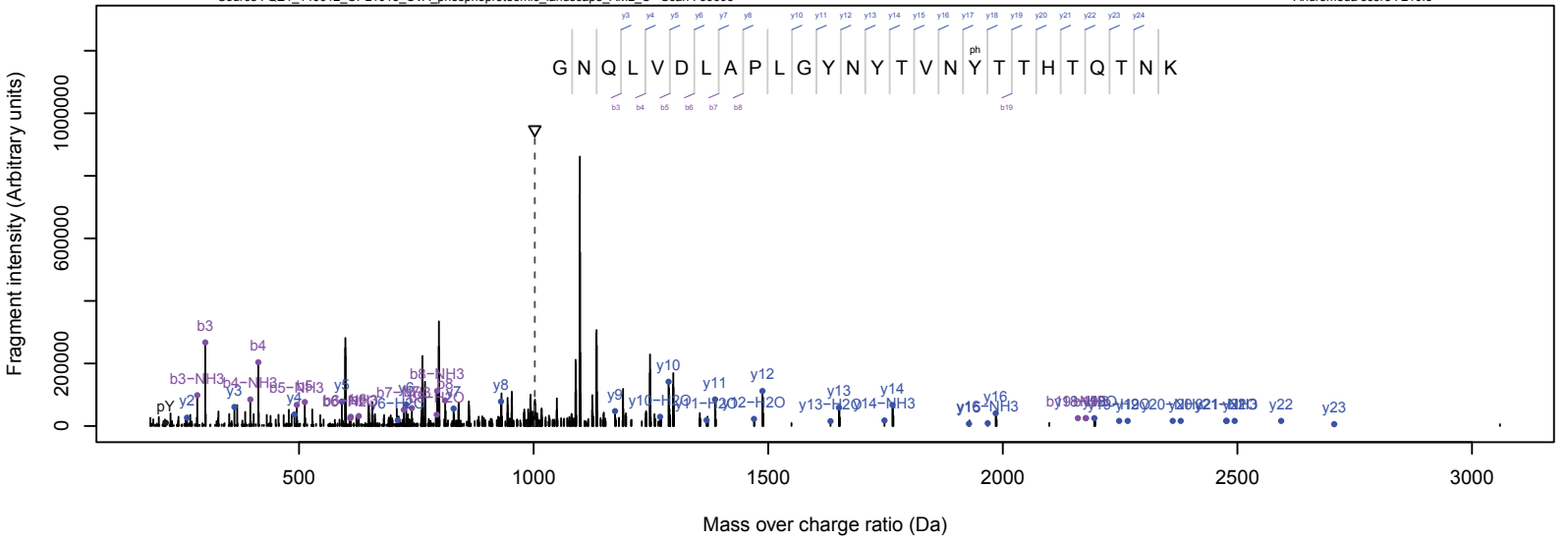
Andromeda score : 210.64



**GNQLVDLAPLGYNYTVNYTTHTQTNK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39695

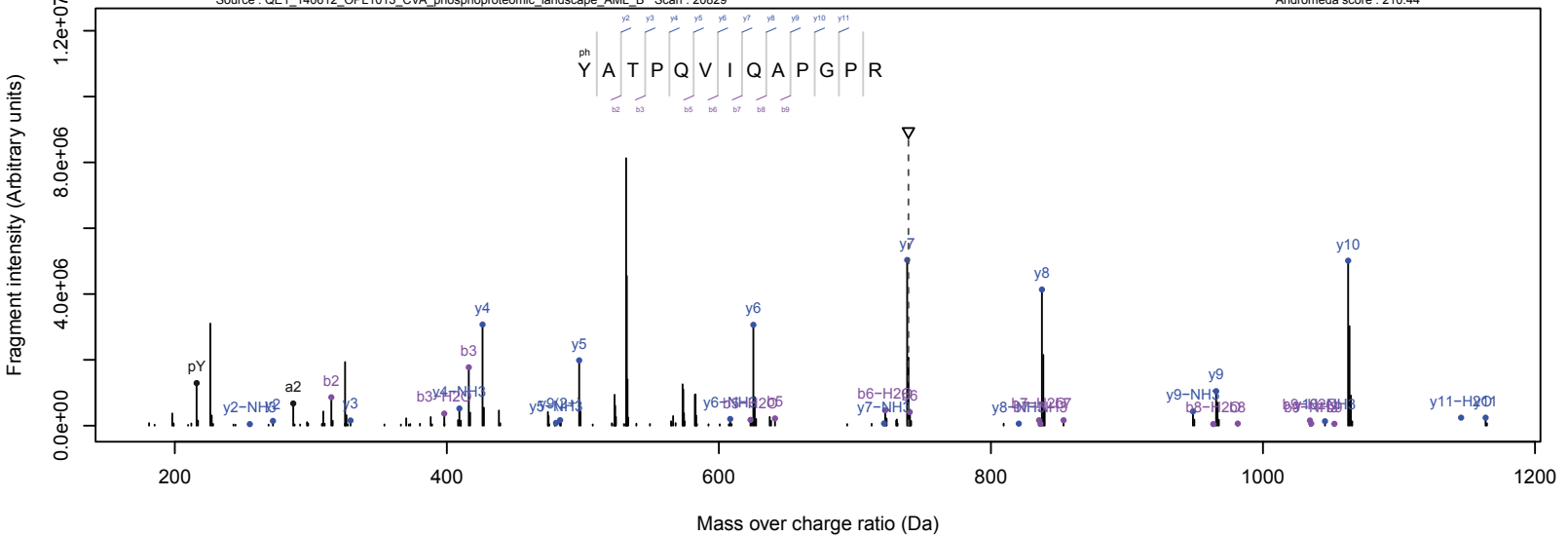
Andromeda score : 210.5



**Y(ph)ATPQVIQAPGPR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20829

Andromeda score : 210.44

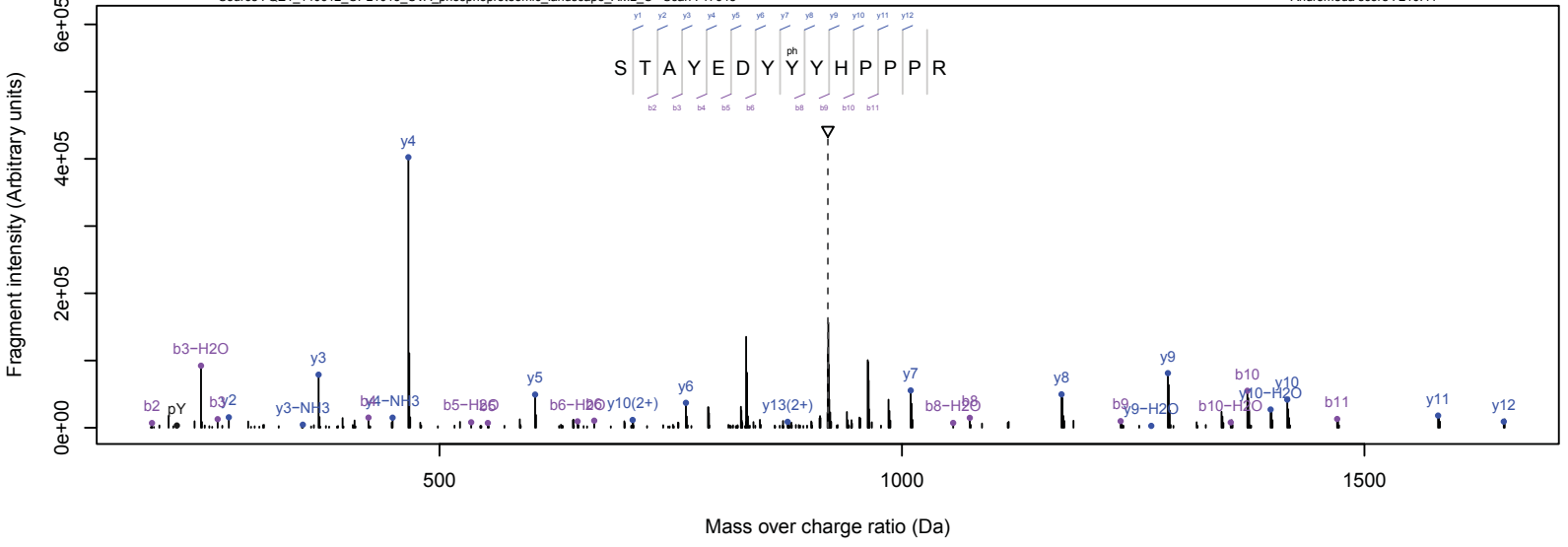




STAYEDYY(ph)YHPPPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17648

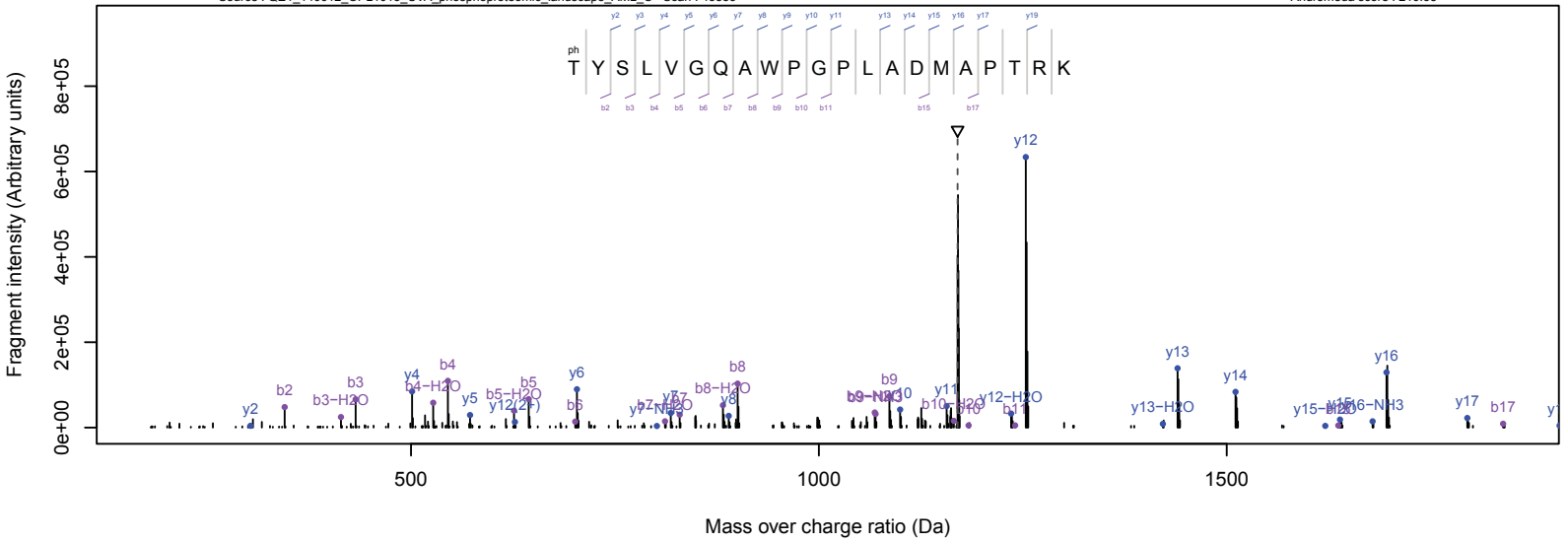
Andromeda score : 210.41



T(ph)YSLVGQAWPGPLADMAPTRK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43883

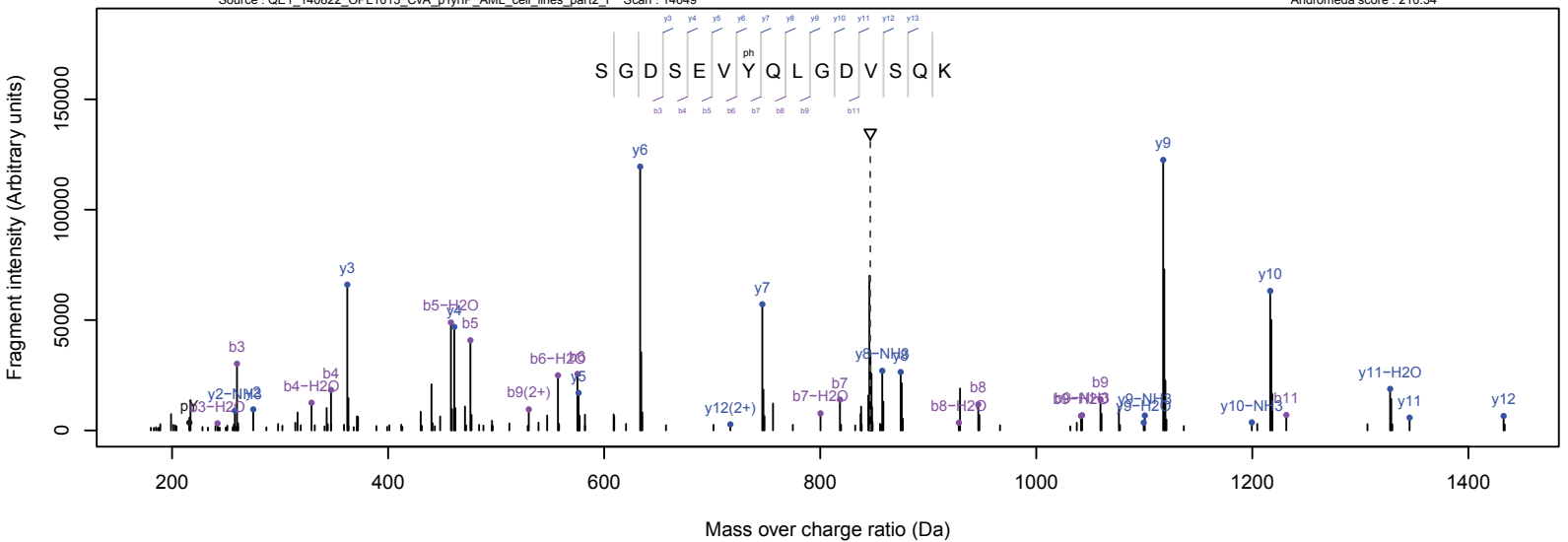
Andromeda score : 210.38



SGDSEVYQLGDVSQK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 14649

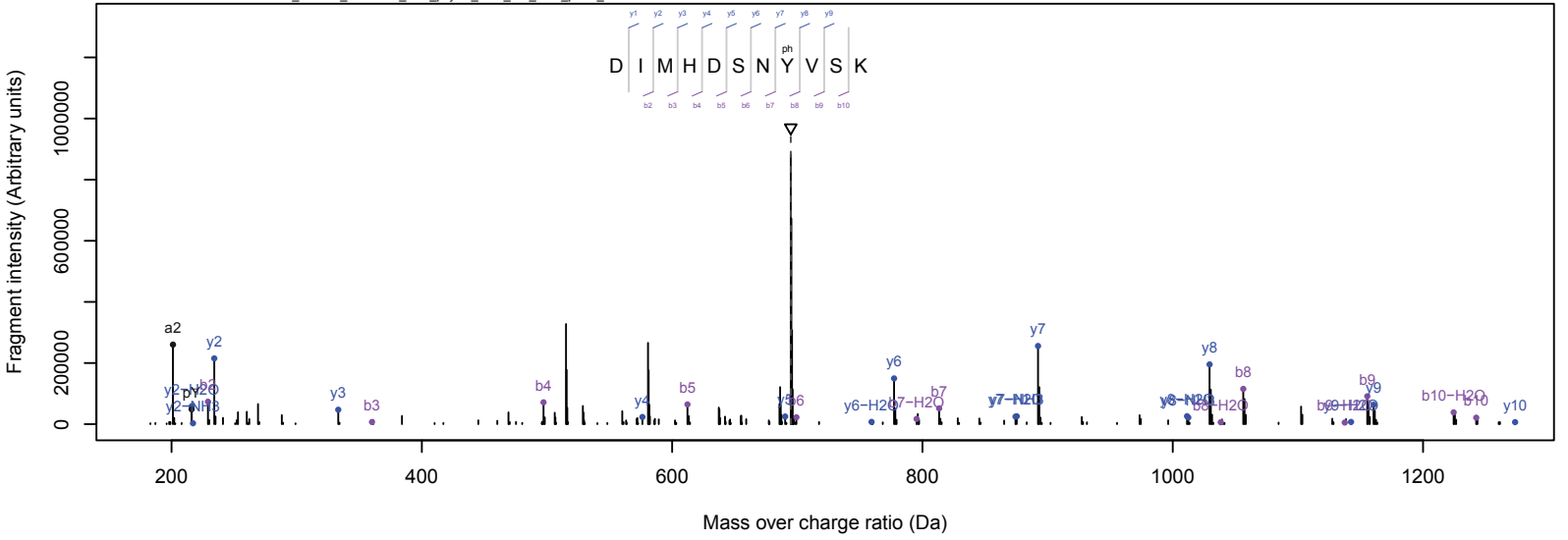
Andromeda score : 210.34



DIMHDSNY(ph)VSK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 8198

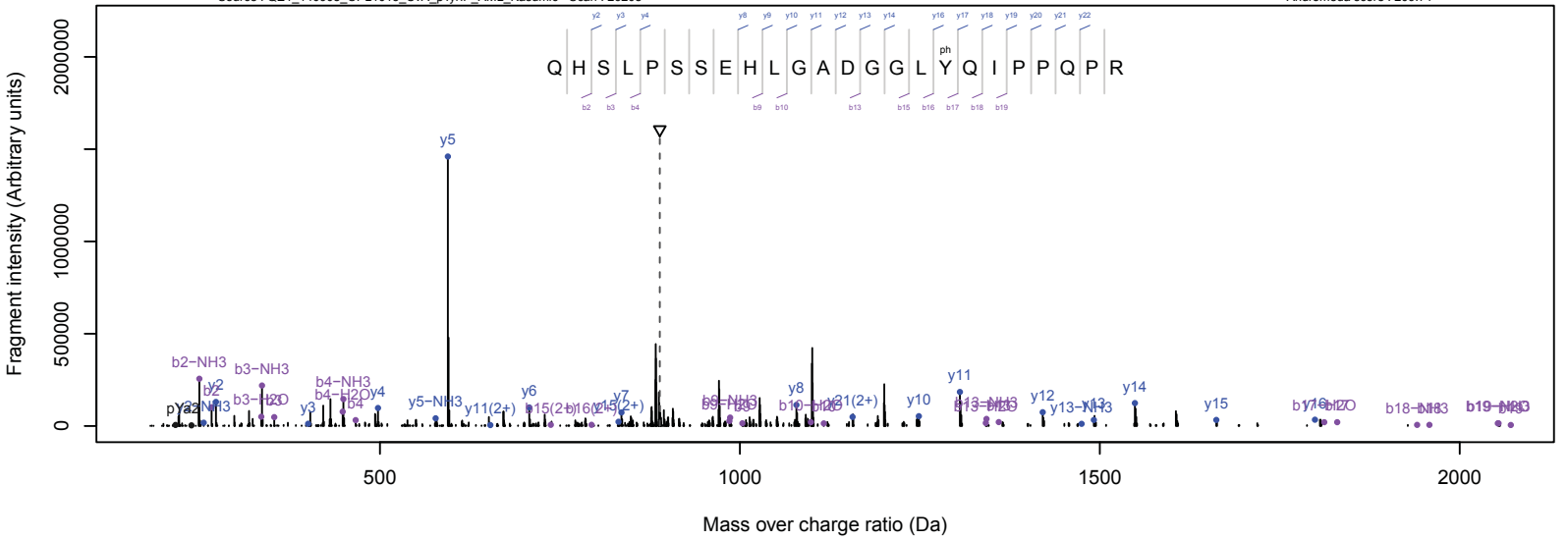
Andromeda score : 210.15



QHSLPSEHLGADGGLYQIIPPQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 26208

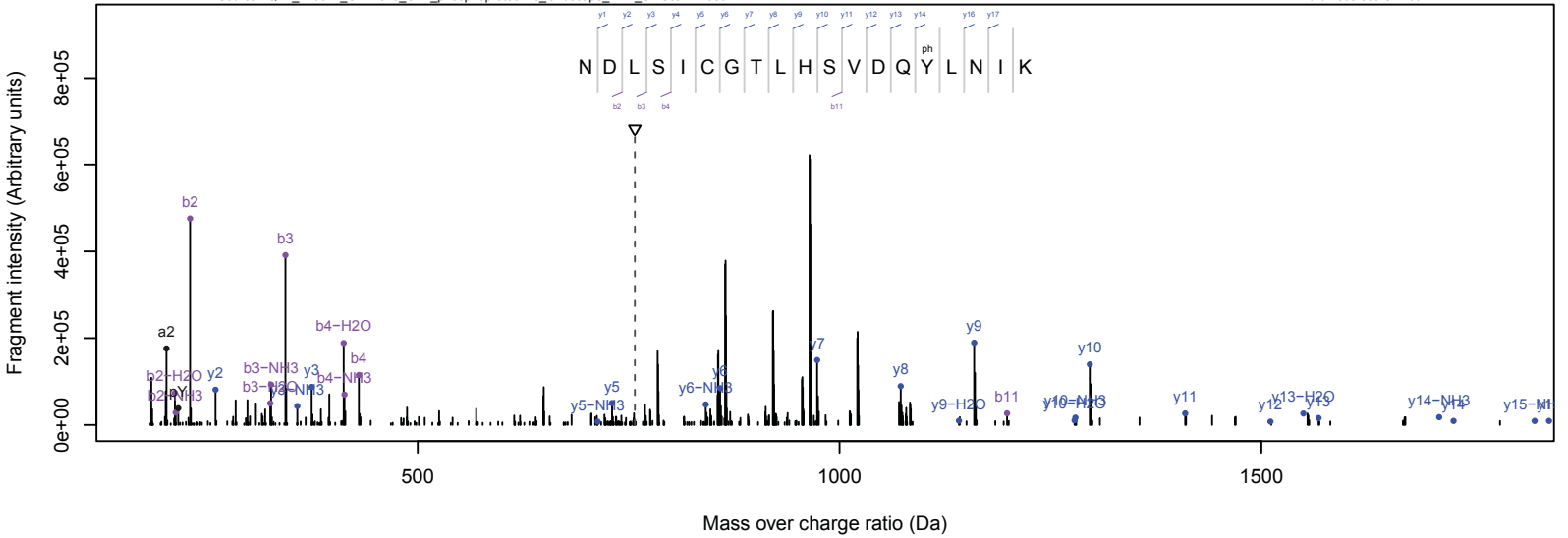
Andromeda score : 209.71



NDSLICGTLHSVDQYLNLIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43894

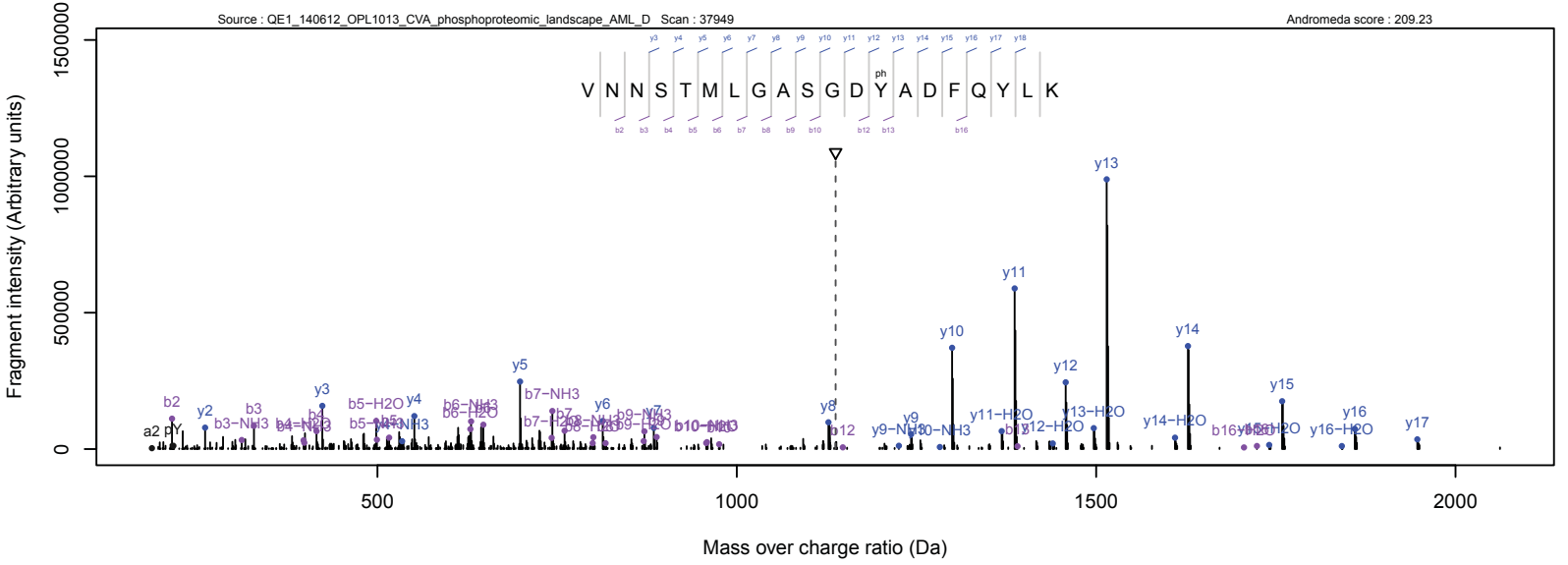
Andromeda score : 209.7



VNNSTMLGASG DY ADFQYLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 37949

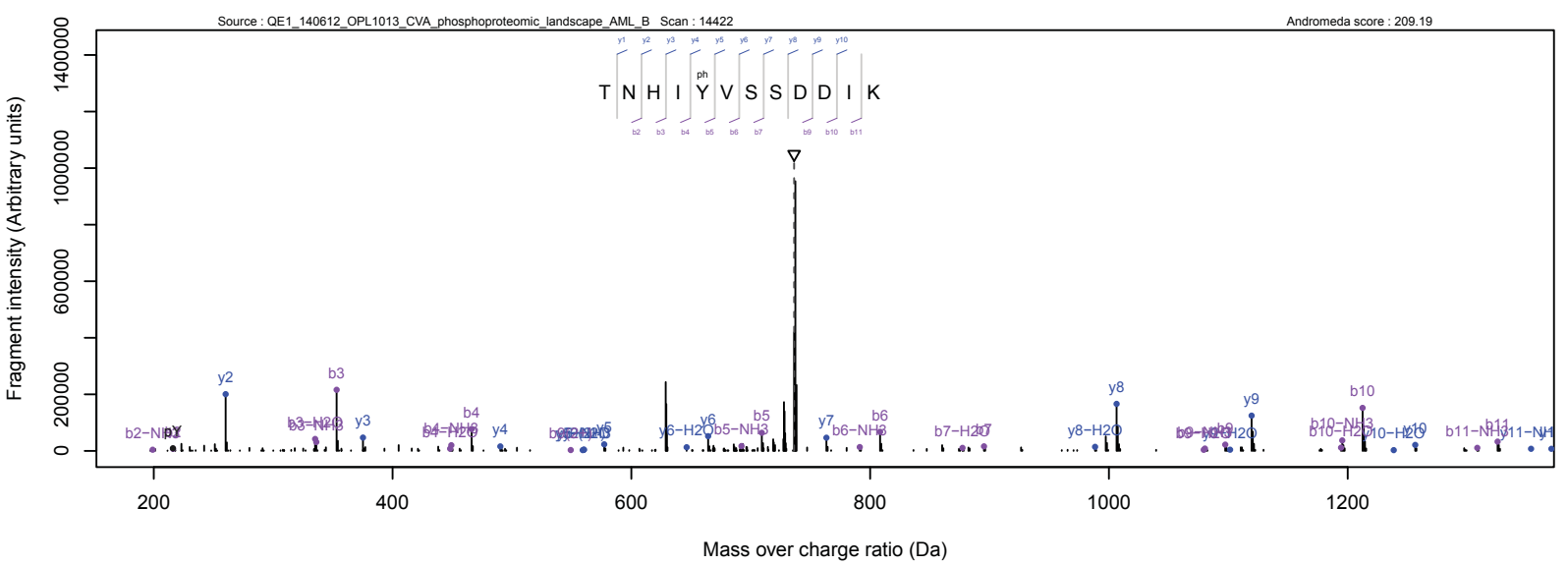
Andromeda score : 209.23



TNHIVSSDDIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14422

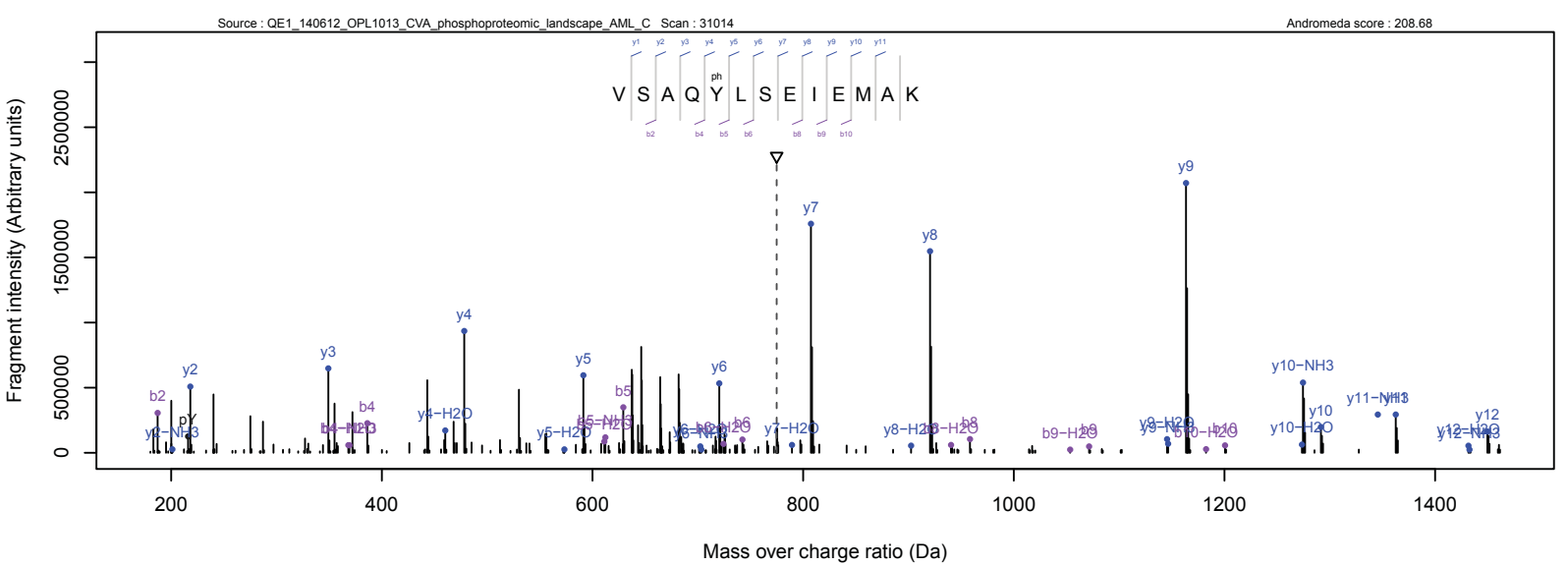
Andromeda score : 209.19



VSAQYLSEIEMAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31014

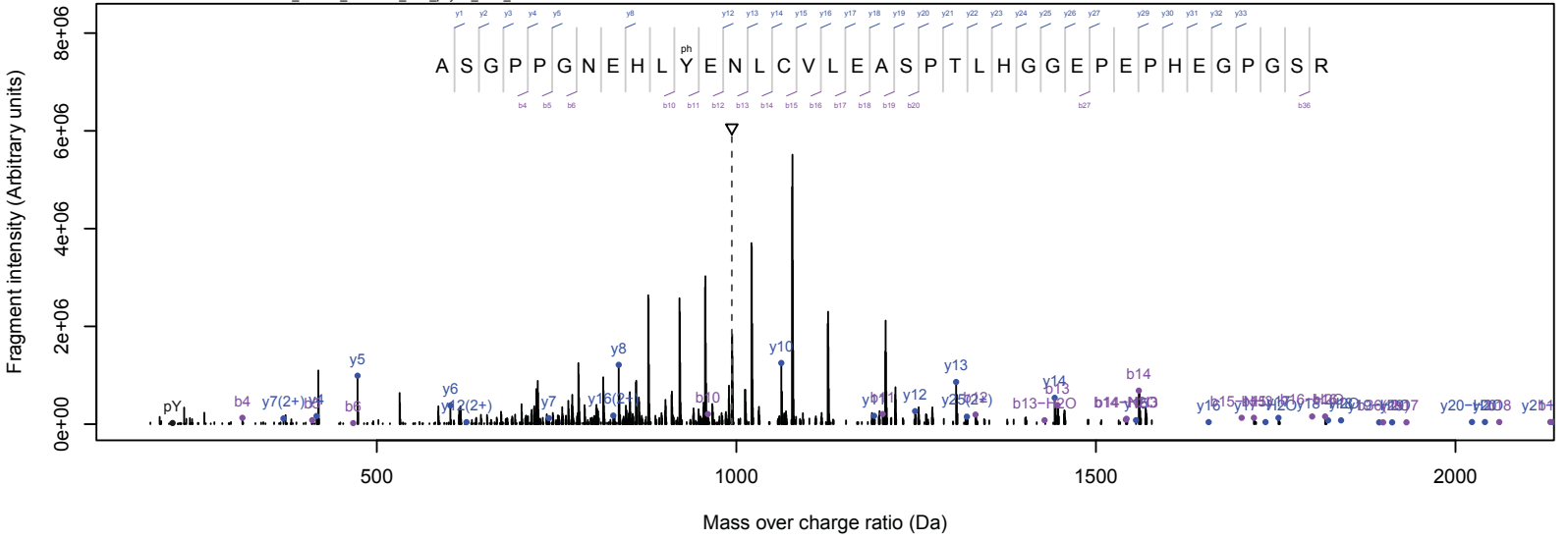
Andromeda score : 208.68



ASGPPGNEHLY(ph)ENLCVLEASPTLHGGEPEPEHEGPGSR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 33993

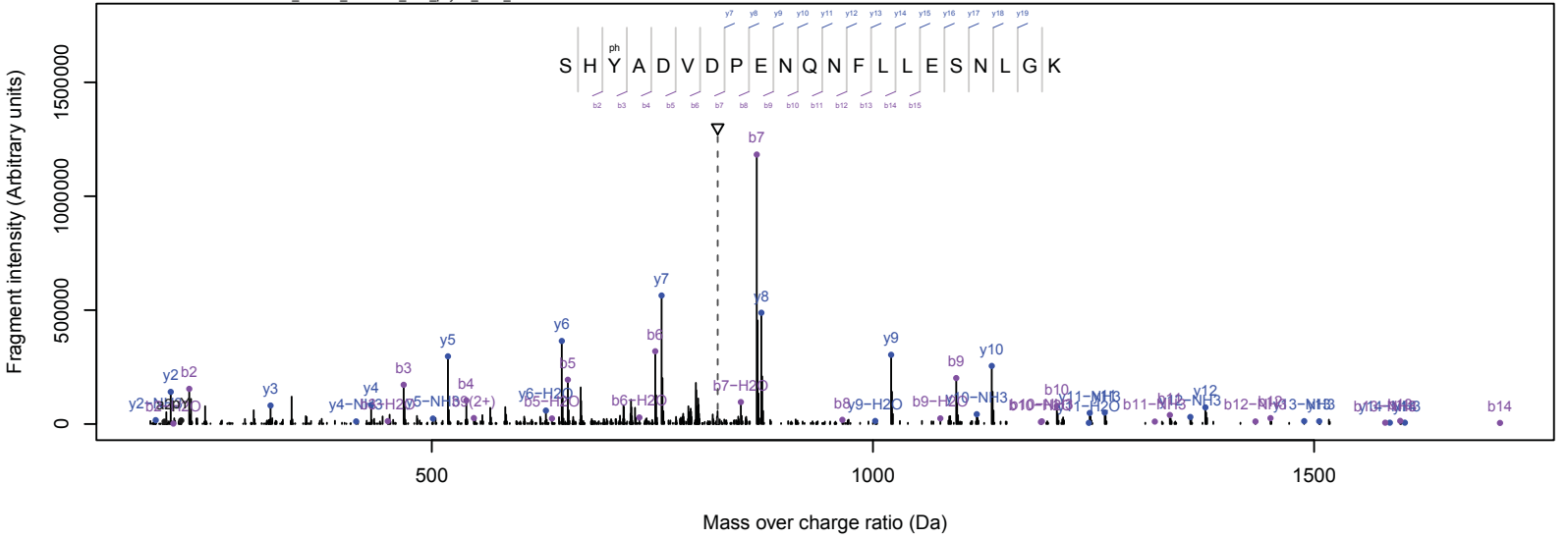
Andromeda score : 208.52



SHY(ph)ADVDPENQNFLLLESNLGK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 34198

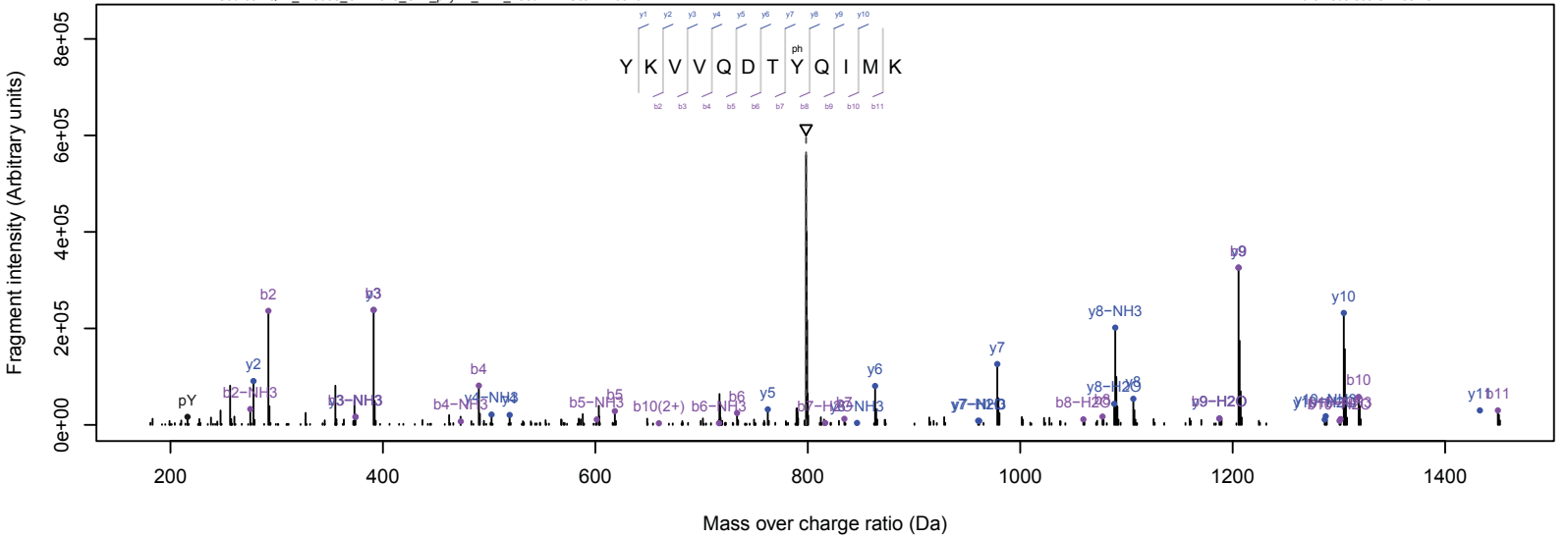
Andromeda score : 208.47



YKVVQDTY(ph)QIMK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 20516

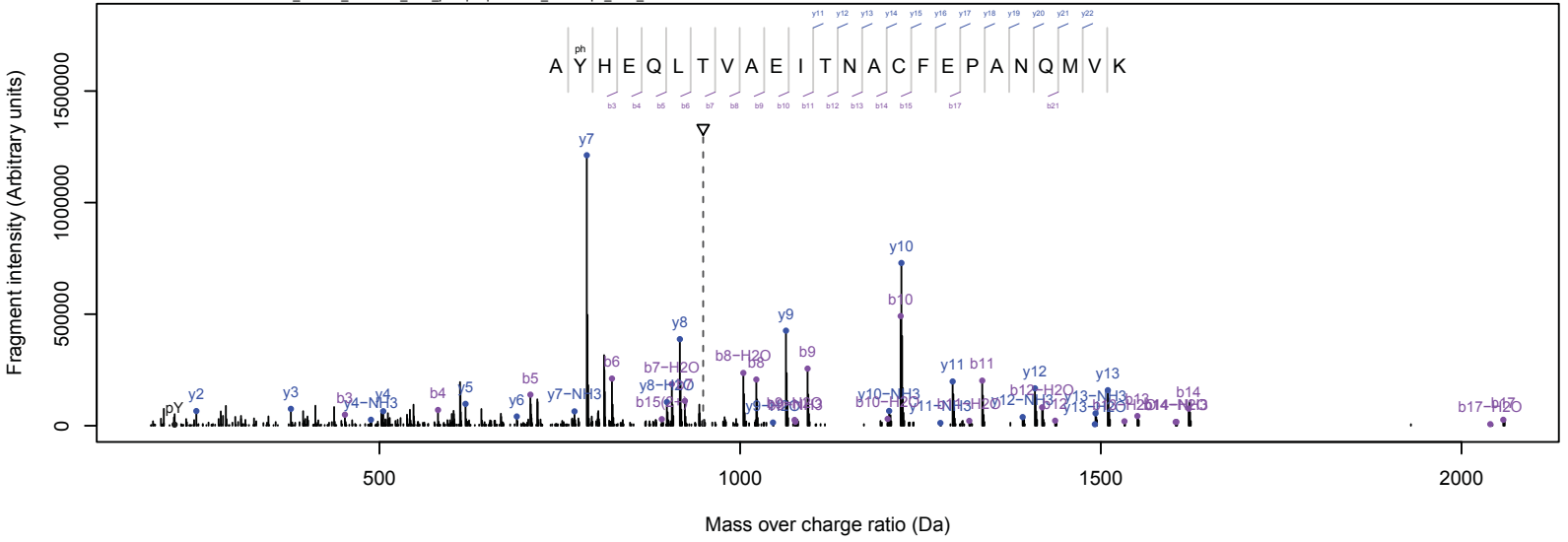
Andromeda score : 208.28



AY(ph)HEQLTVAEITNACFEPANQMVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 39211

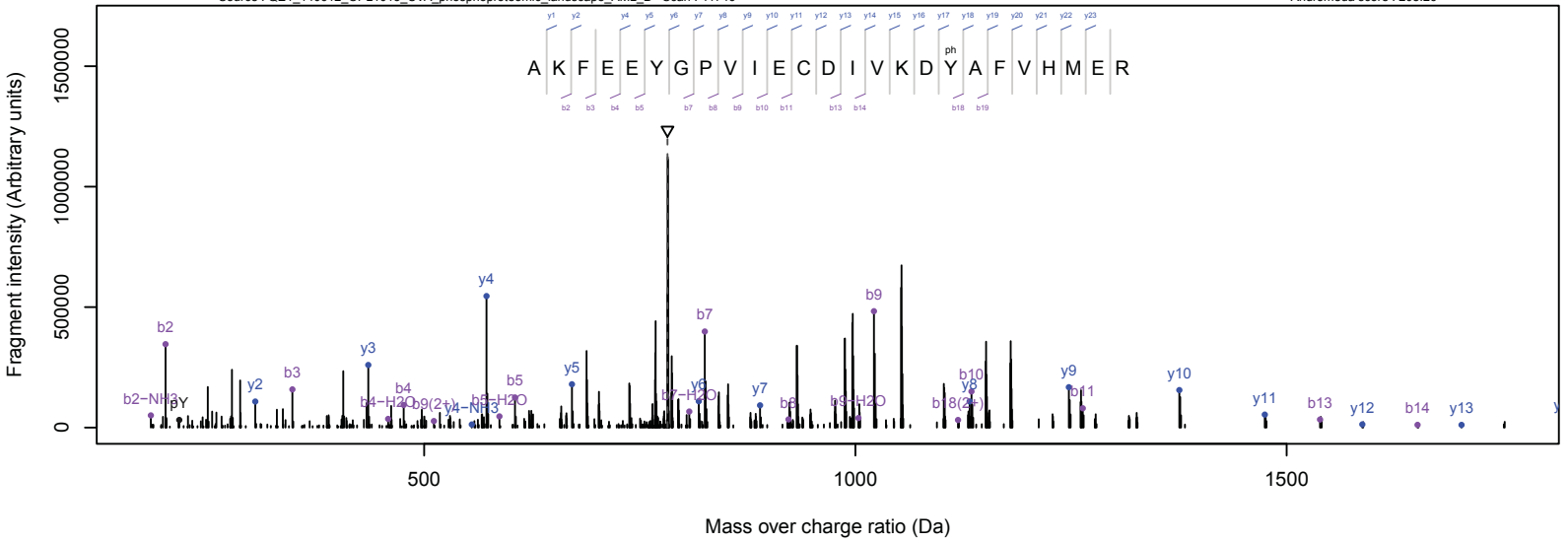
Andromeda score: 208.27



AKFEEYGPVIECDIVKDY(ph)AFVHMER-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 41746

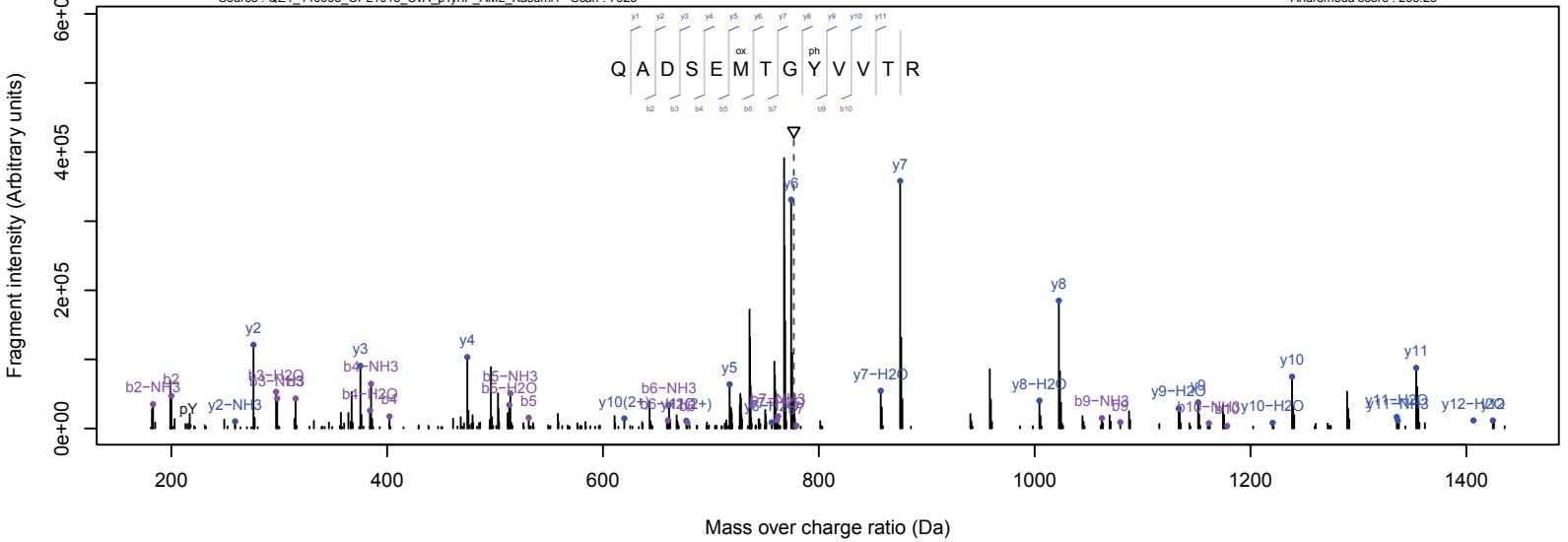
Andromeda score: 208.26



QADSEM(ox)TGY(ph)VVTR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 7925

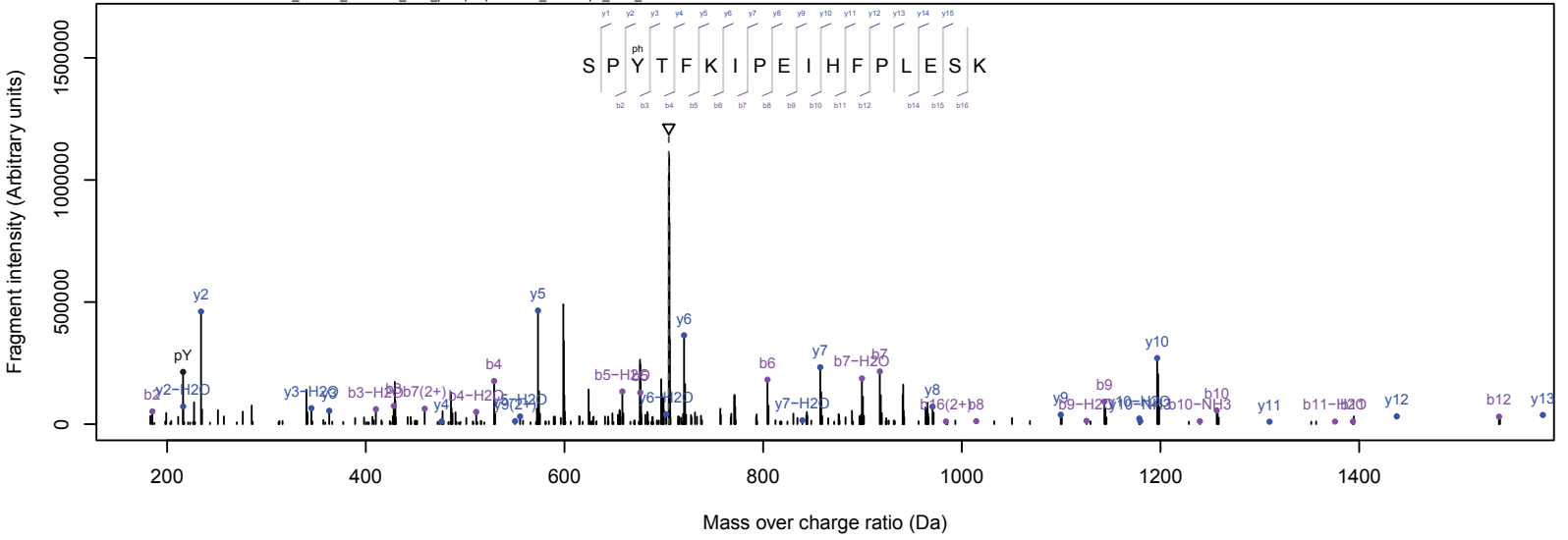
Andromeda score: 208.23



**SPY(ph)TFKIPEIHFPLESK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45378

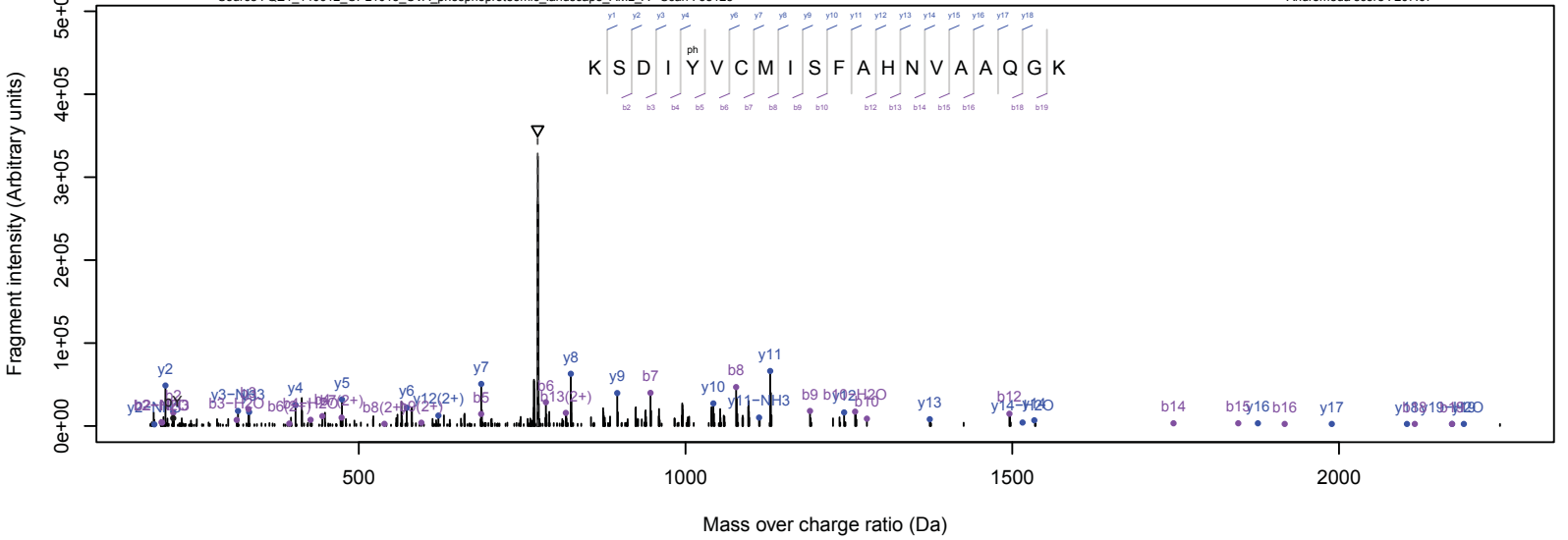
Andromeda score : 208.17



**KSDIY(ph)VCMI SFAHNVAAGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 38125

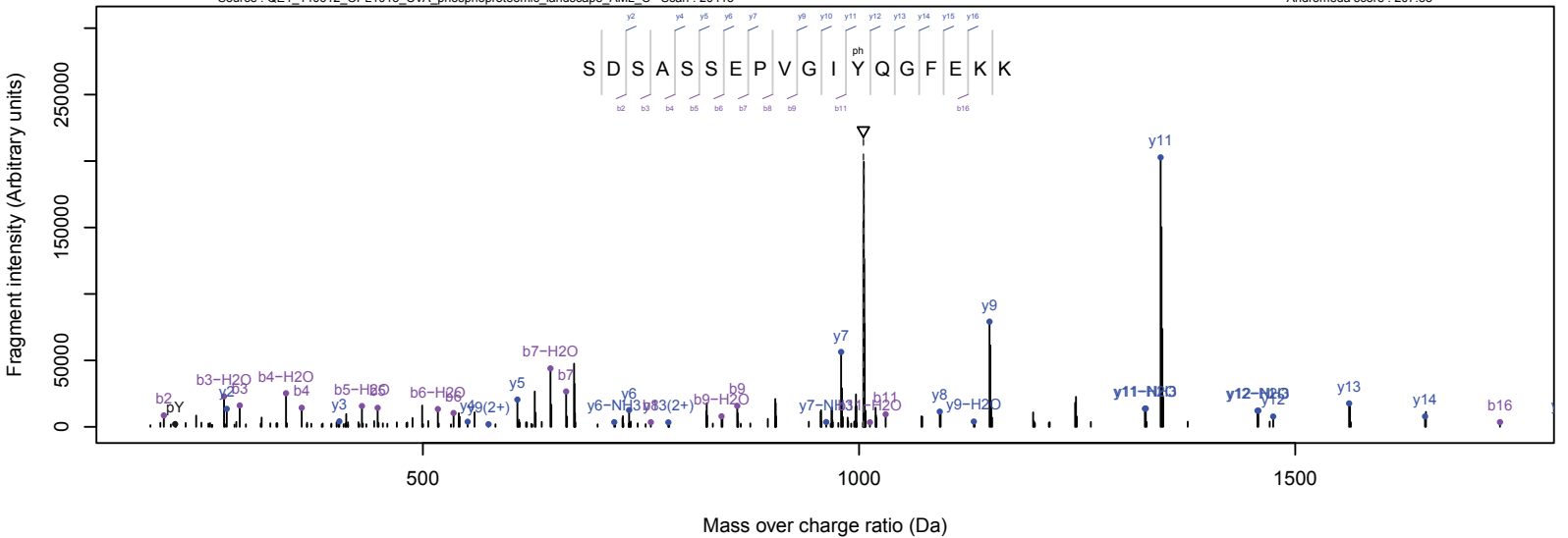
Andromeda score : 207.67



**SDSASSEVGIY(ph)QGFEKK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20415

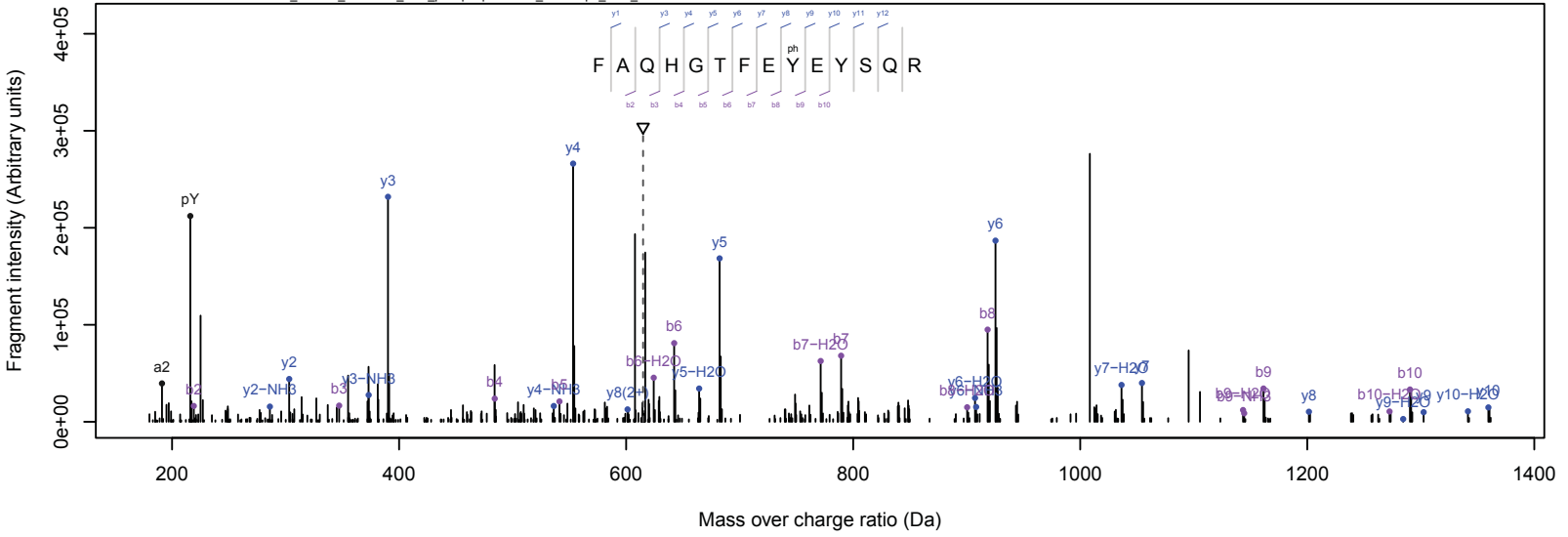
Andromeda score : 207.53



FAQHGTFEY(ph)EYSQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16695

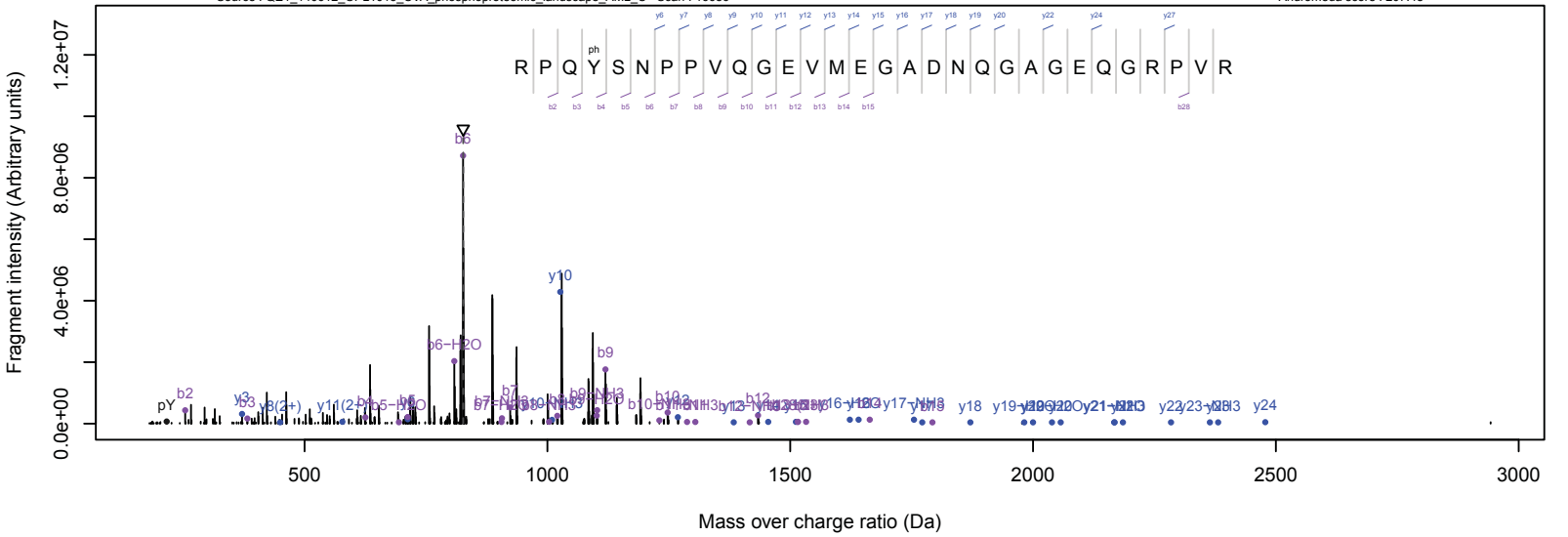
Andromeda score : 207.52



RPQY(ph)SNPPVQGEVMEGADNQGAGEQGRPVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16686

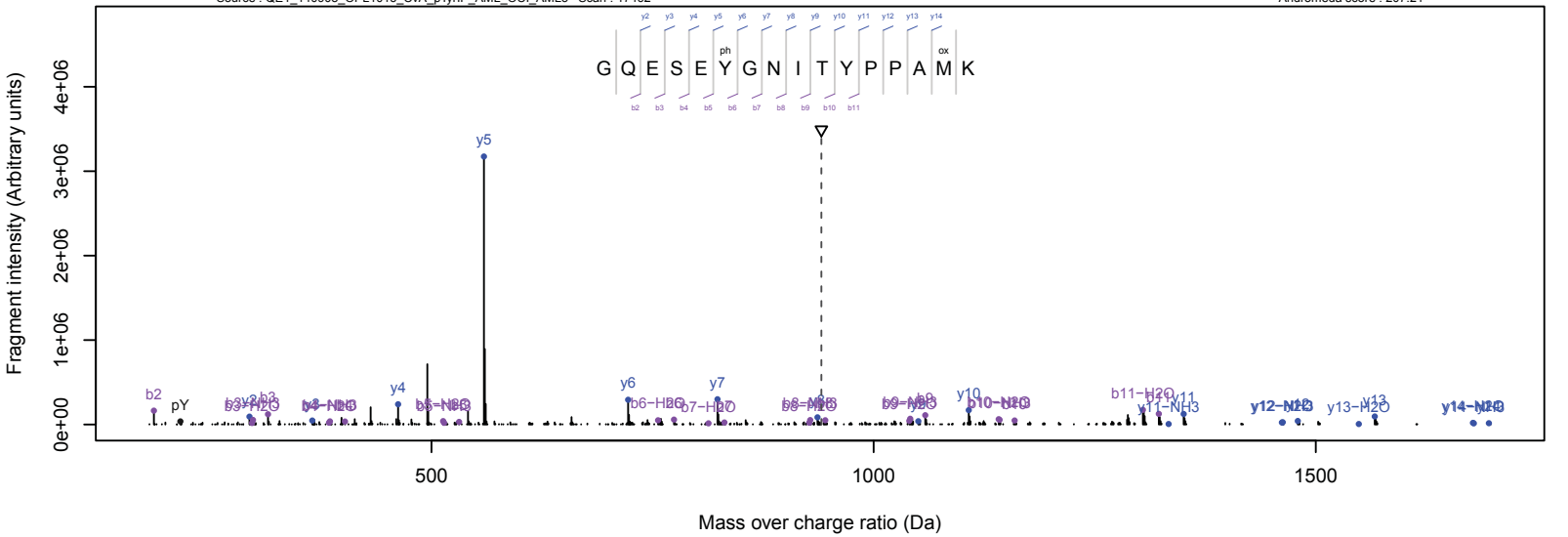
Andromeda score : 207.48



GQESEY(ph)GNITYPPAM(ox)K-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 17462

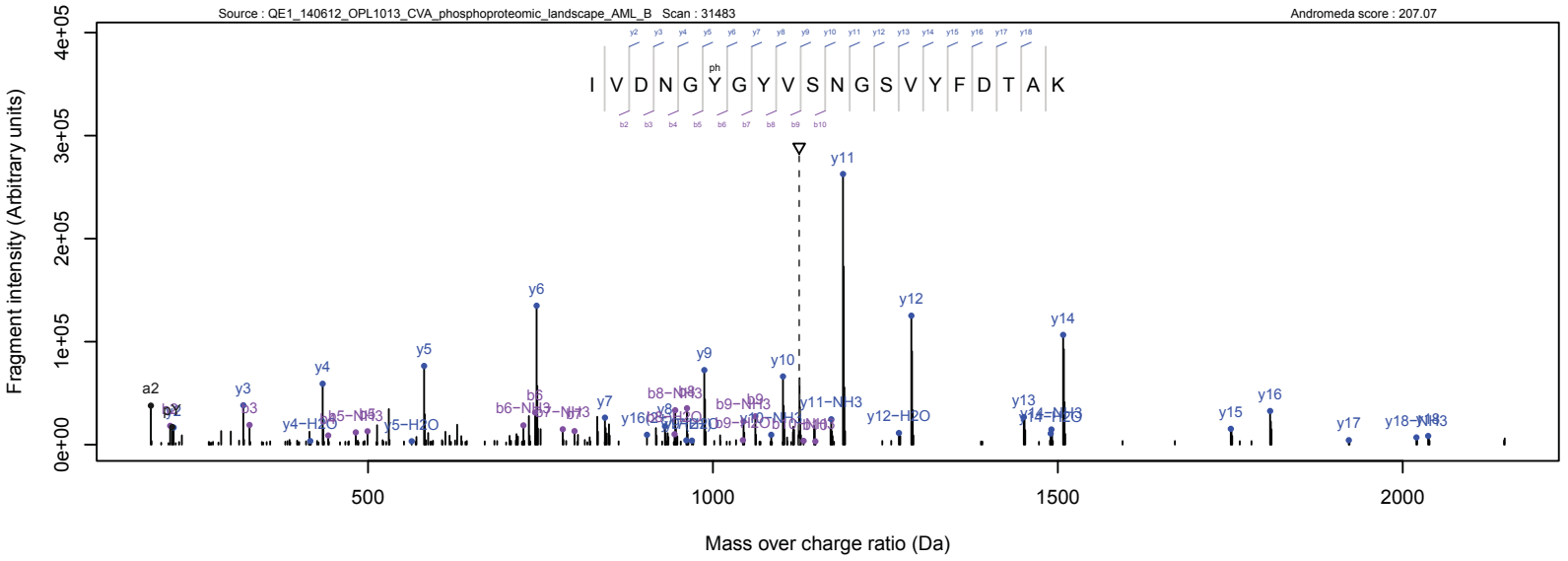
Andromeda score : 207.21



**IVDNGY(ph)GYVSNNGSVYFDTAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31483

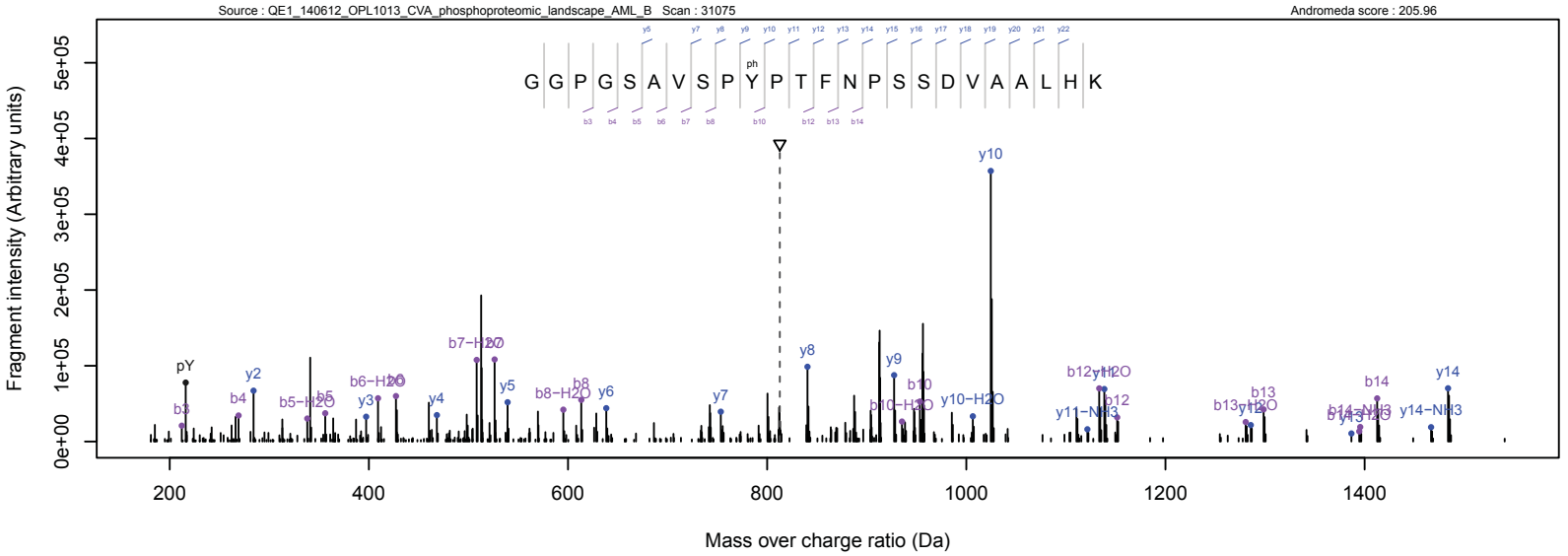
Andromeda score : 207.07



**GGPGSAVSPY(ph)PTFNPSSDVAALHK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31075

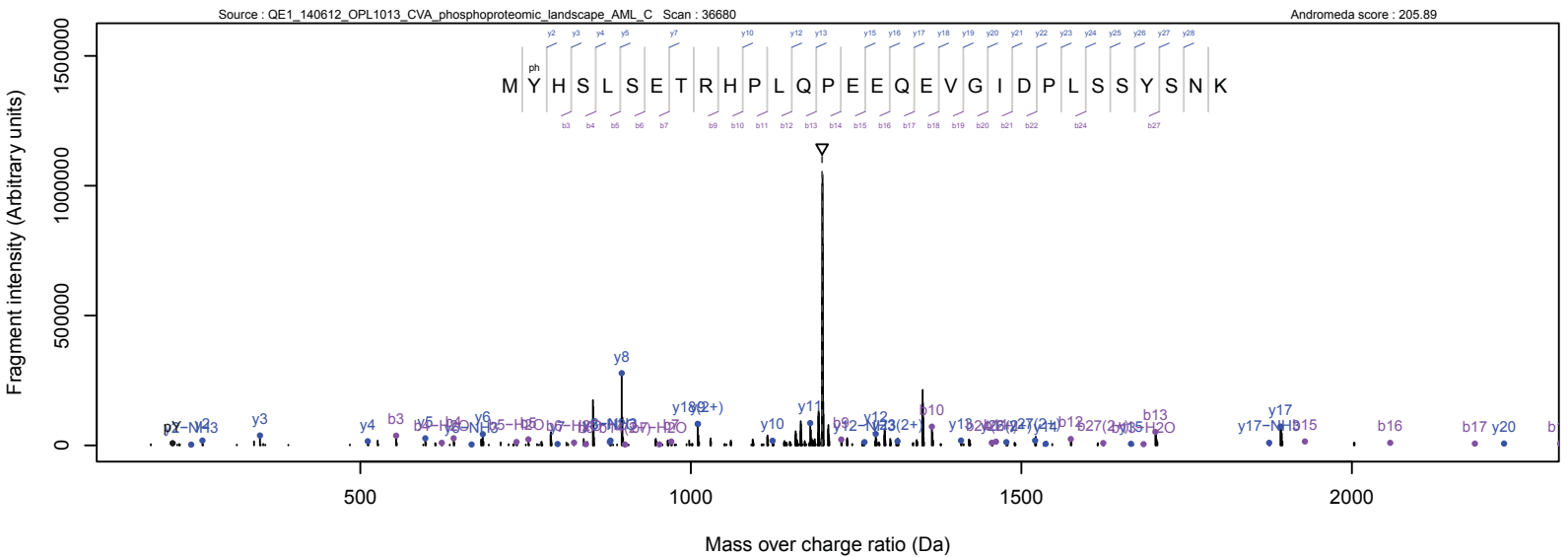
Andromeda score : 205.96



**(ac)MY(ph)HSLSETRHPLQPEEQEVGIDPLSSYSNK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36680

Andromeda score : 205.89

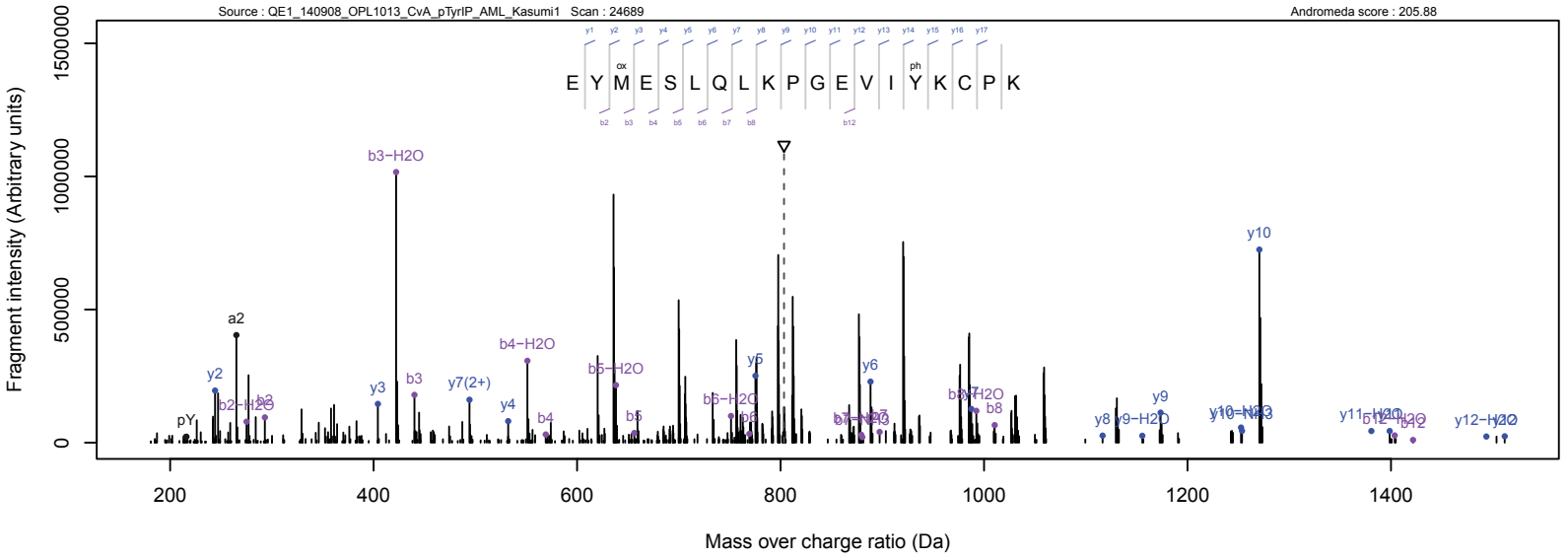




EYM(ox)ESLQLKPGEVY(ph)KCPK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 24689

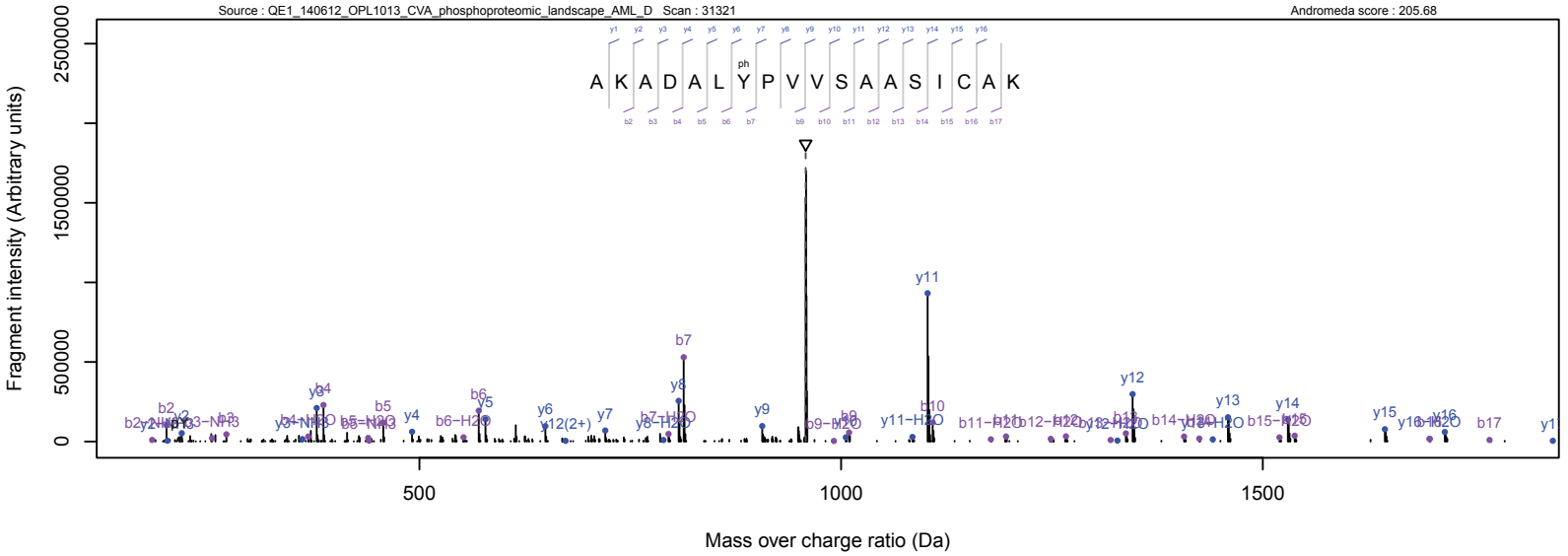
Andromeda score : 205.88



AKADALY(ph)PVVSAASICK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 31321

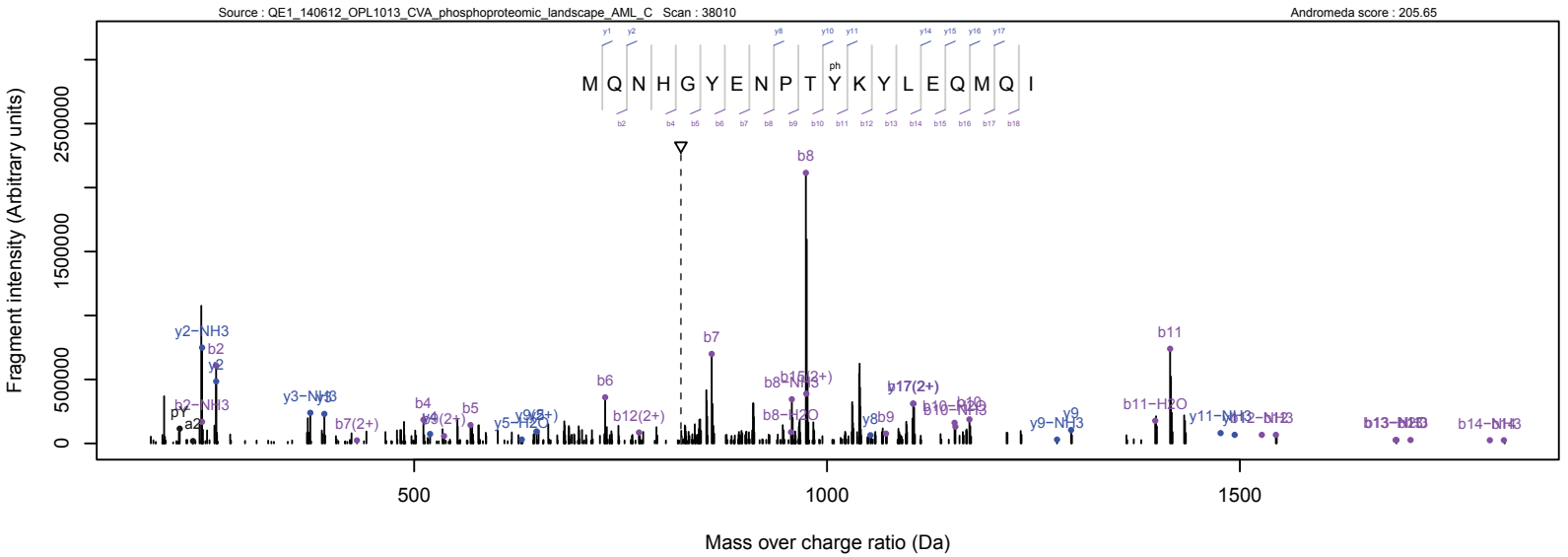
Andromeda score : 205.68



MQNHGYENPTY(ph)KYLEQMQL-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38010

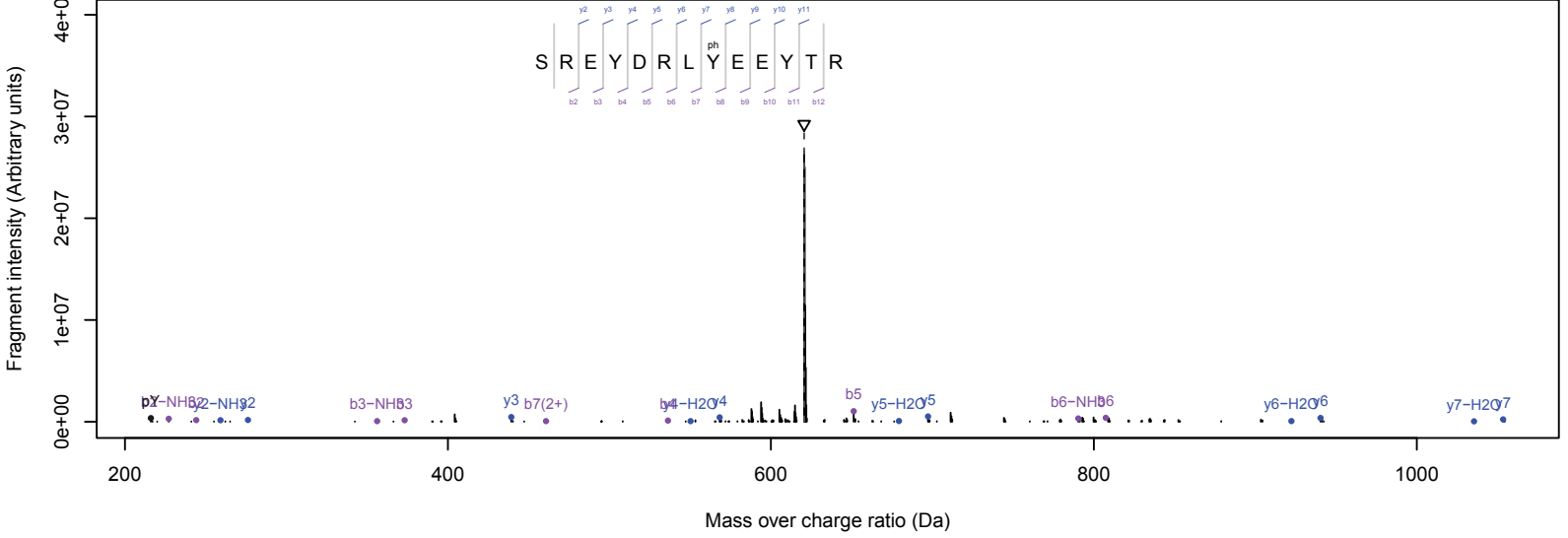
Andromeda score : 205.65



**SREYDRLY(ph)EEYTR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 16654

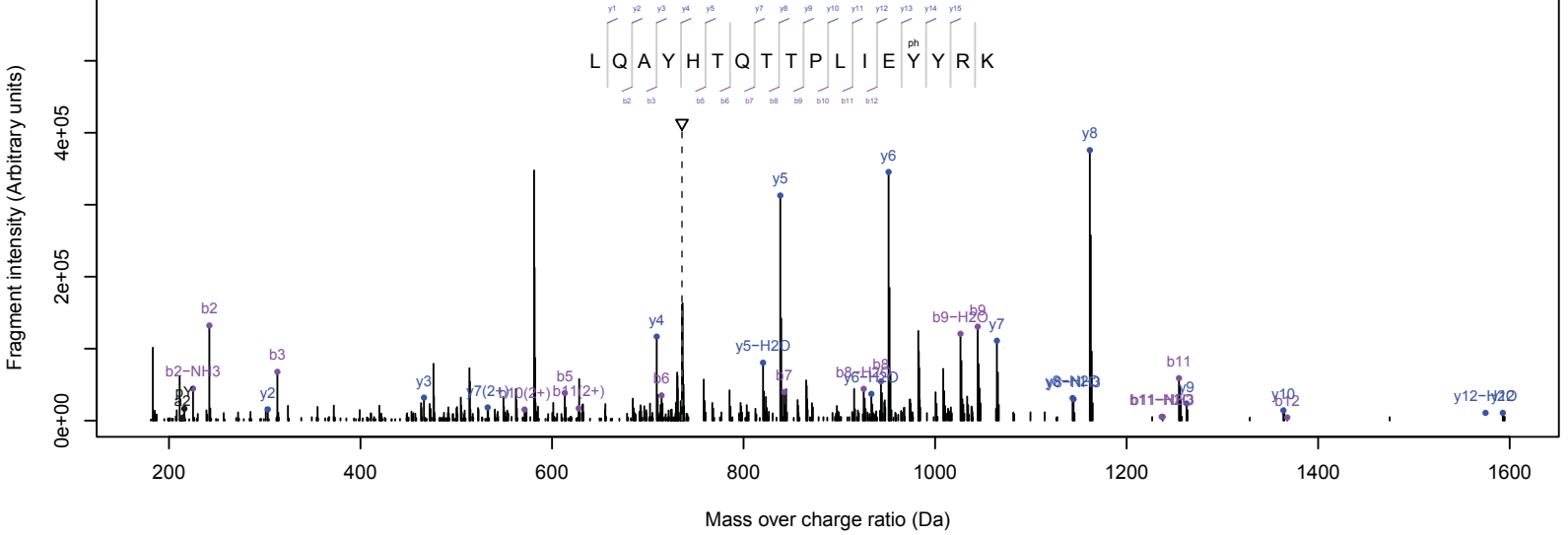
Andromeda score : 205.27



**LQAYHTQTTPLIEY(ph)YRK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23487

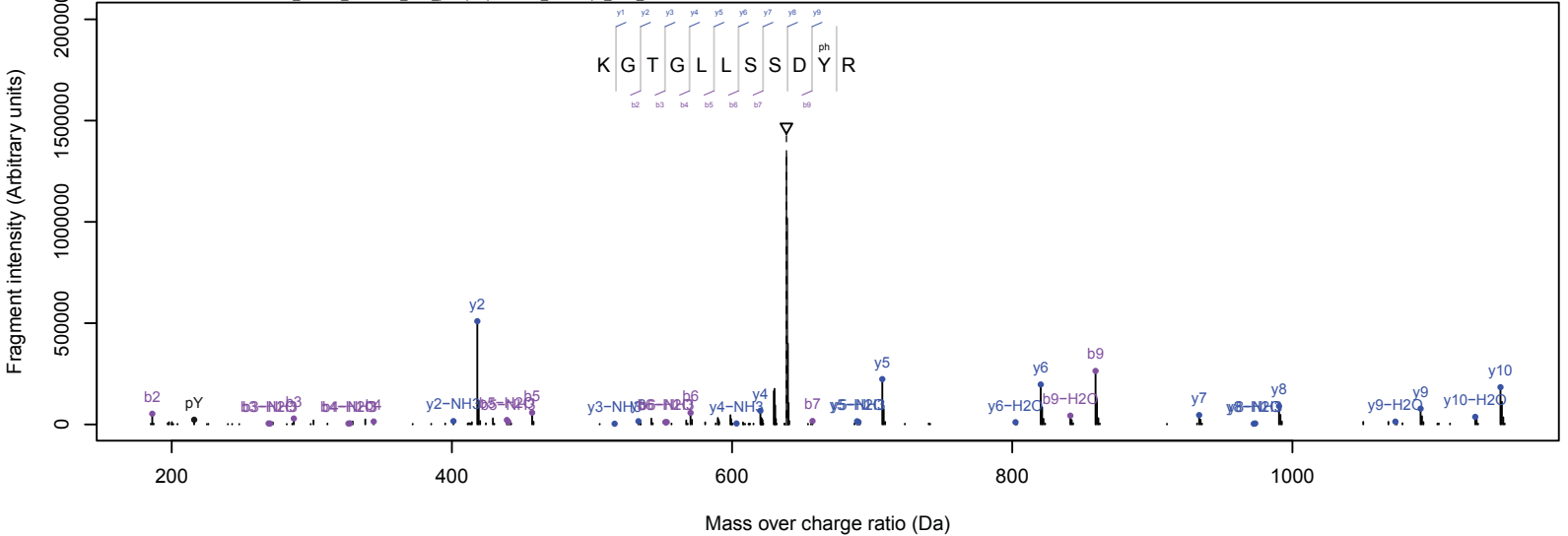
Andromeda score : 205.18



**KGTLSSDY(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11171

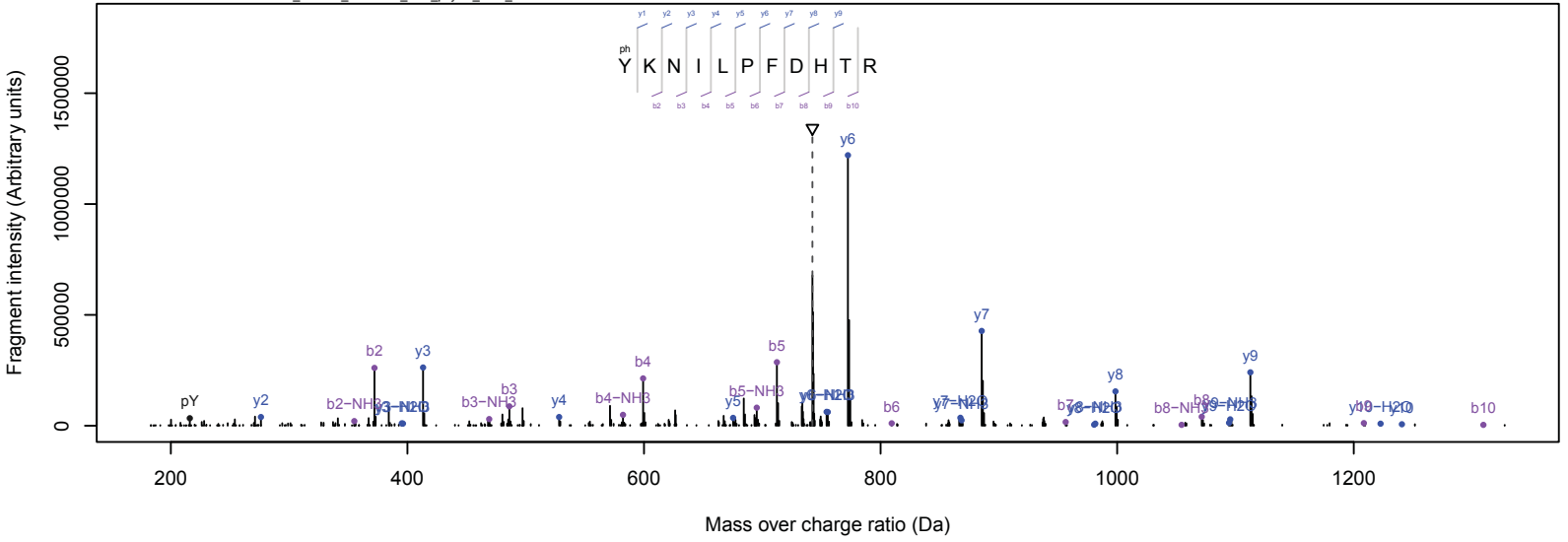
Andromeda score : 205.02



Y(ph)KNILPFDHTR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 22922

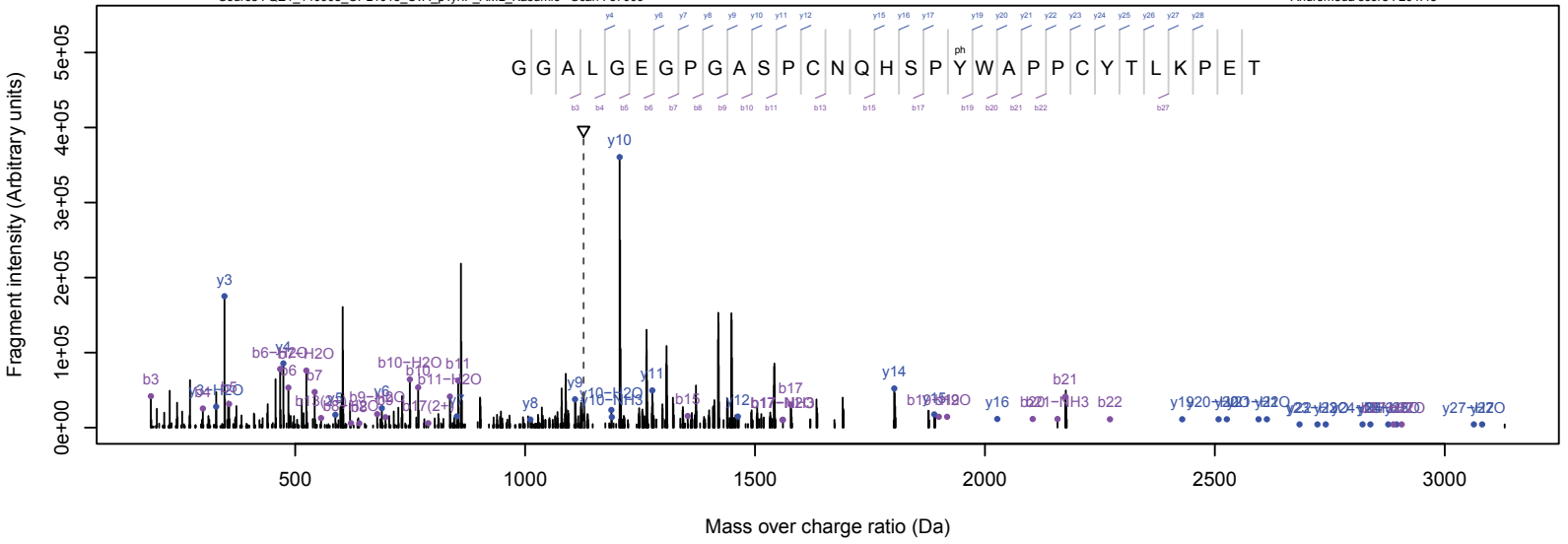
Andromeda score : 205.02



GGALGEGPGASPCNQHSFY(ph)WAPPCYTLKPET-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 37009

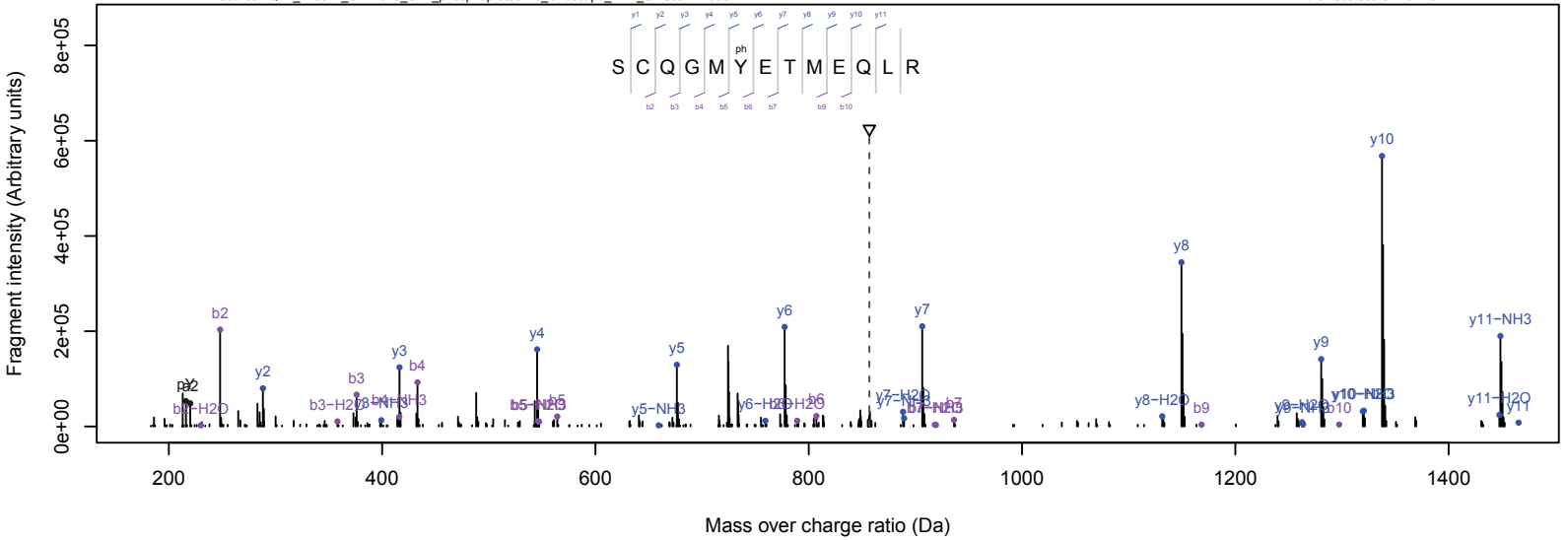
Andromeda score : 204.48



SCQGMYPETMEQLR-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 26004

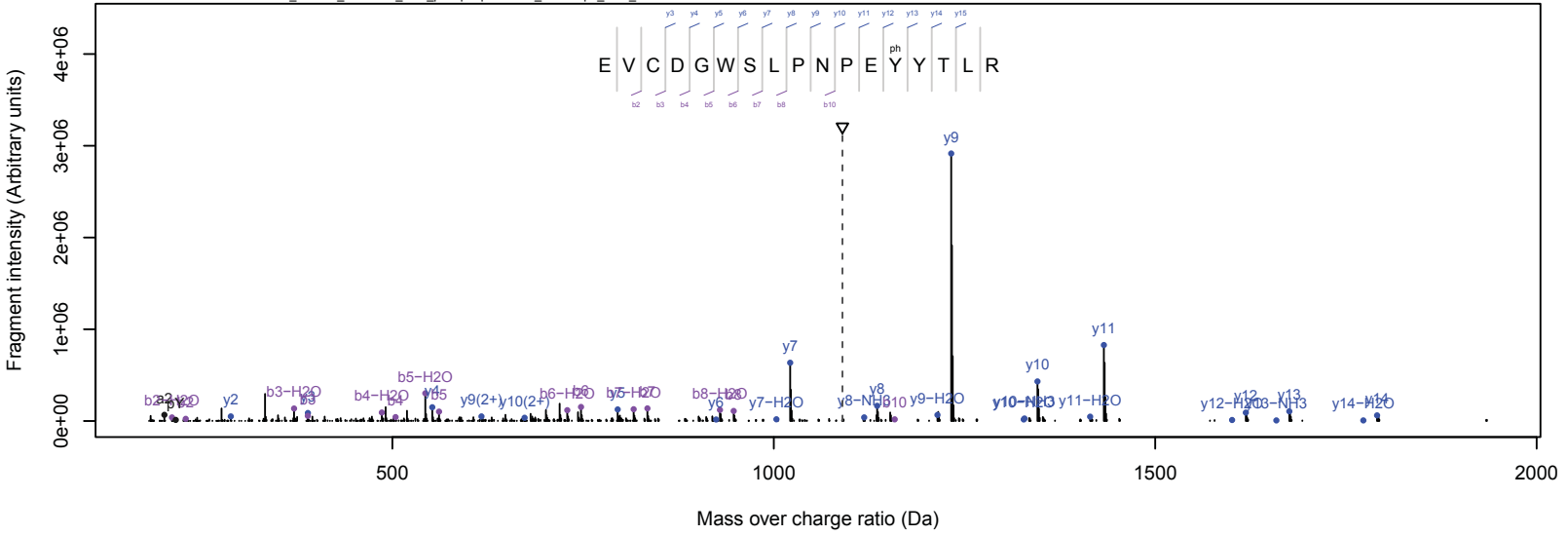
Andromeda score : 204.18



**EVCDGWSLPNPEY(ph)YTLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47754

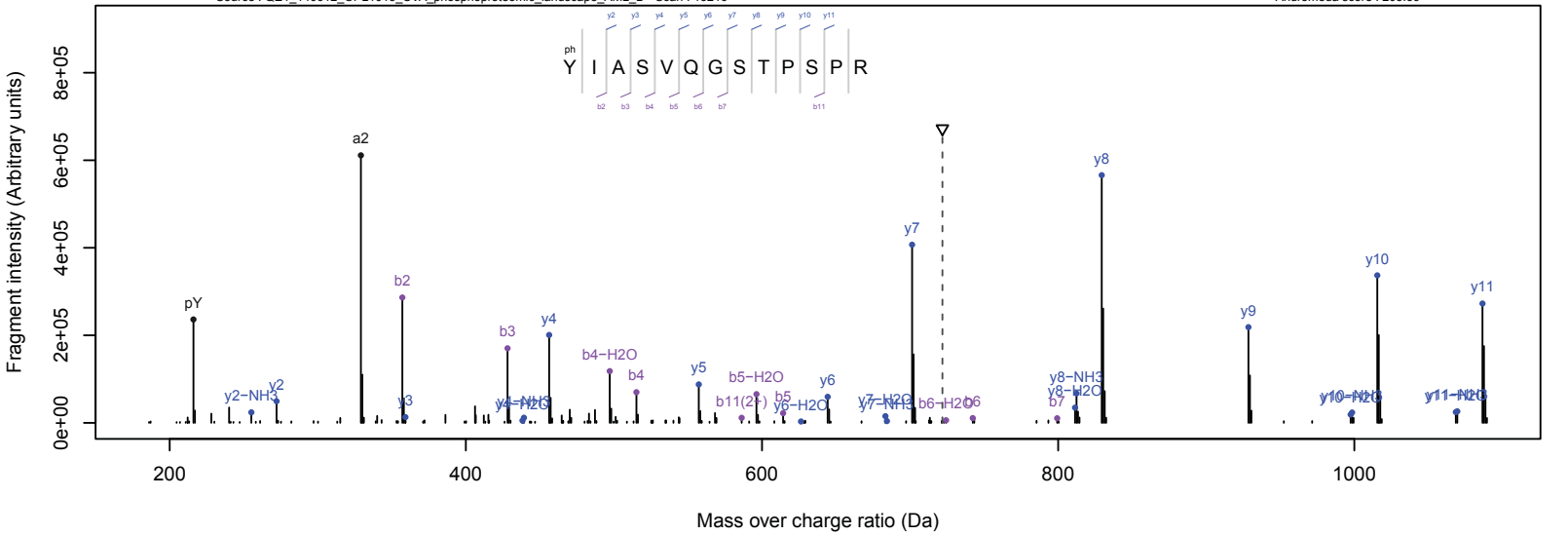
Andromeda score : 203.53



**Y(ph)IASVQGSTPSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 13216

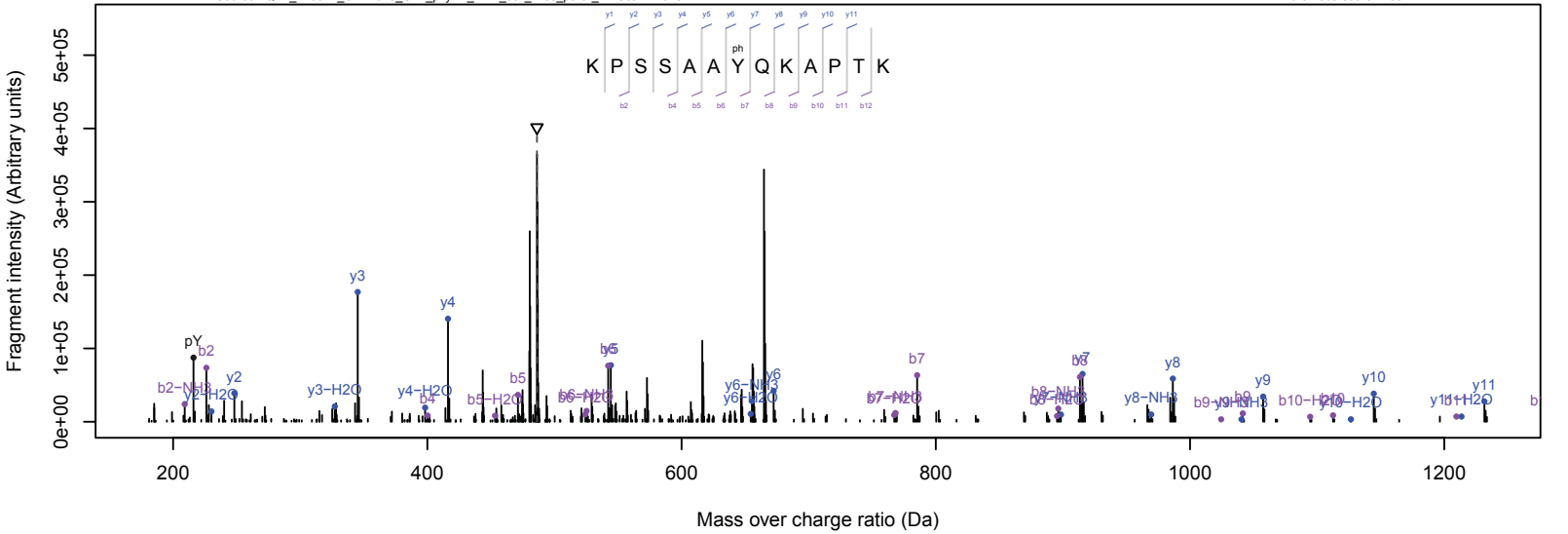
Andromeda score : 203.39



**KPSSAAYQKAPT-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 2626

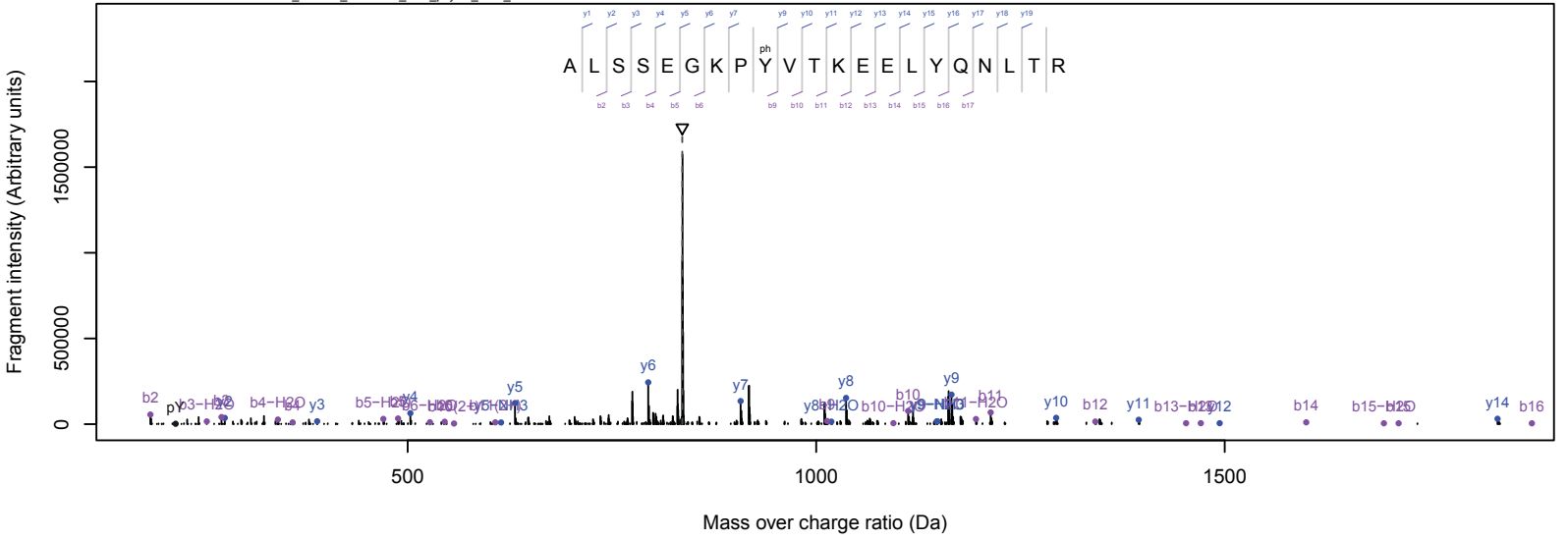
Andromeda score : 203.27



ALSSEGKPY(ph)VTKEELYQLTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 26103

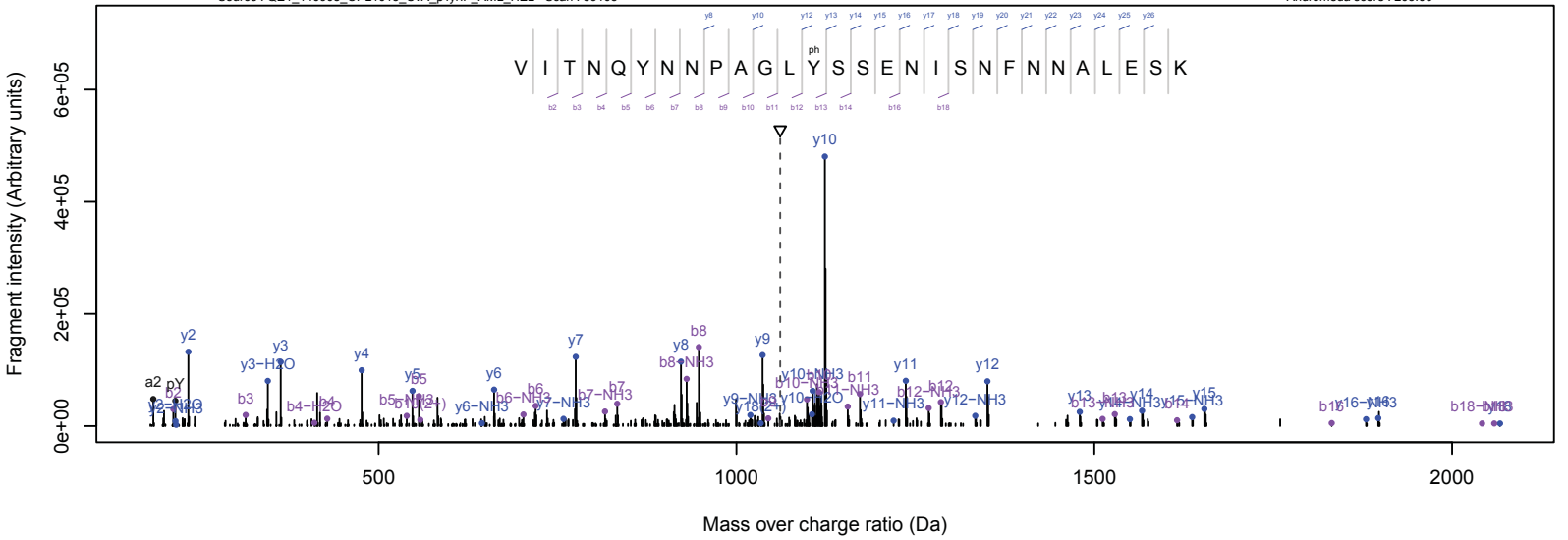
Andromeda score : 203.2



VITNQYNNPAGLY(ph)SSEENISNFNNALESK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 39105

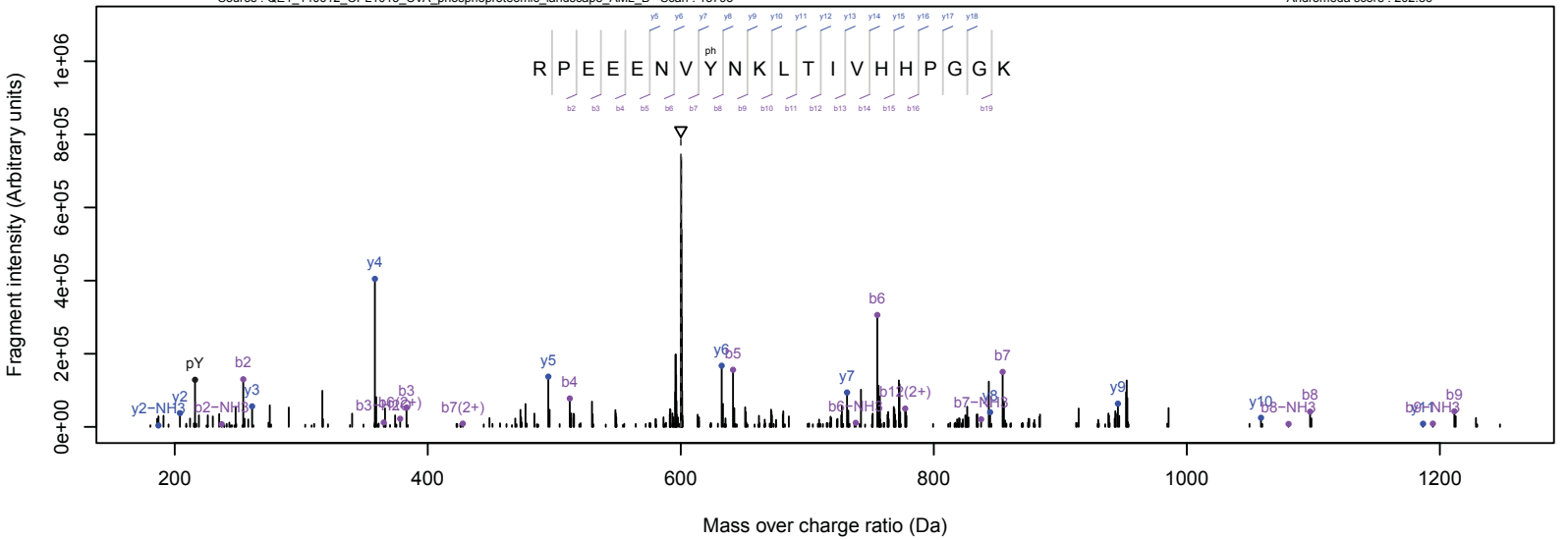
Andromeda score : 203.06



RPEEENVY(ph)NKLTIVHHPGGK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15793

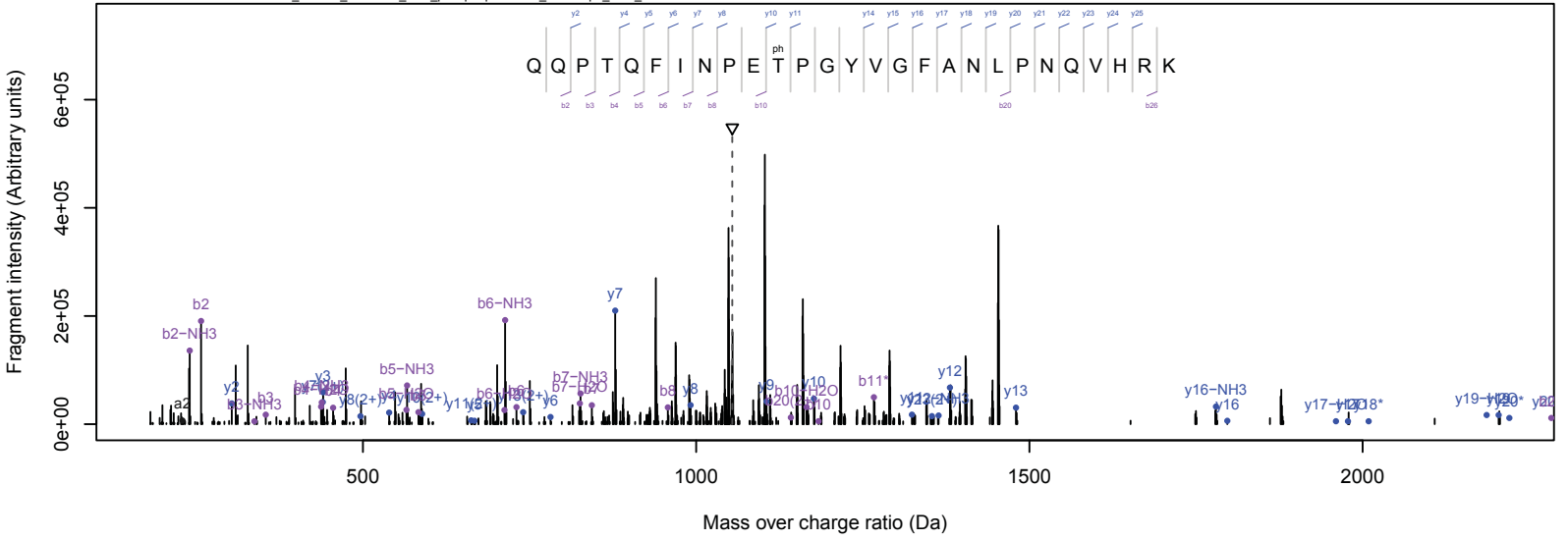
Andromeda score : 202.86



QQPTQFINPETPGYVGFANLPNQVHRK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36261

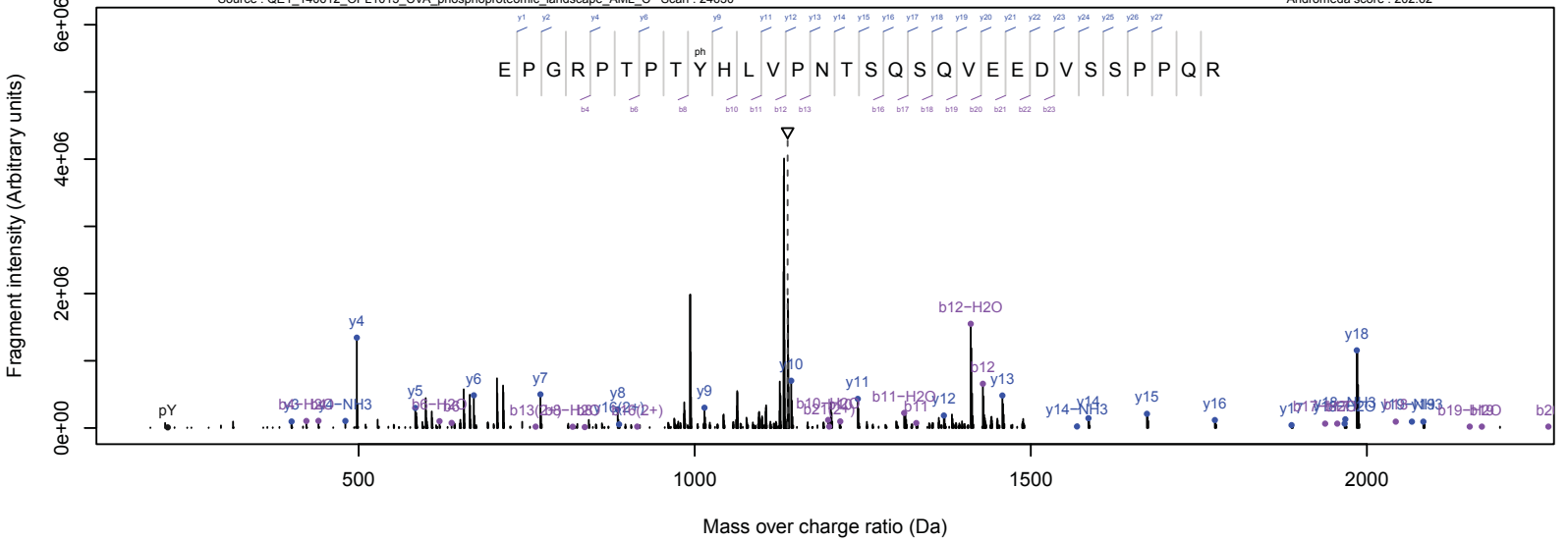
Andromeda score: 202.81



EPGRPTPTYHLVPNTSQQSQVEEDVSSPPQR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24630

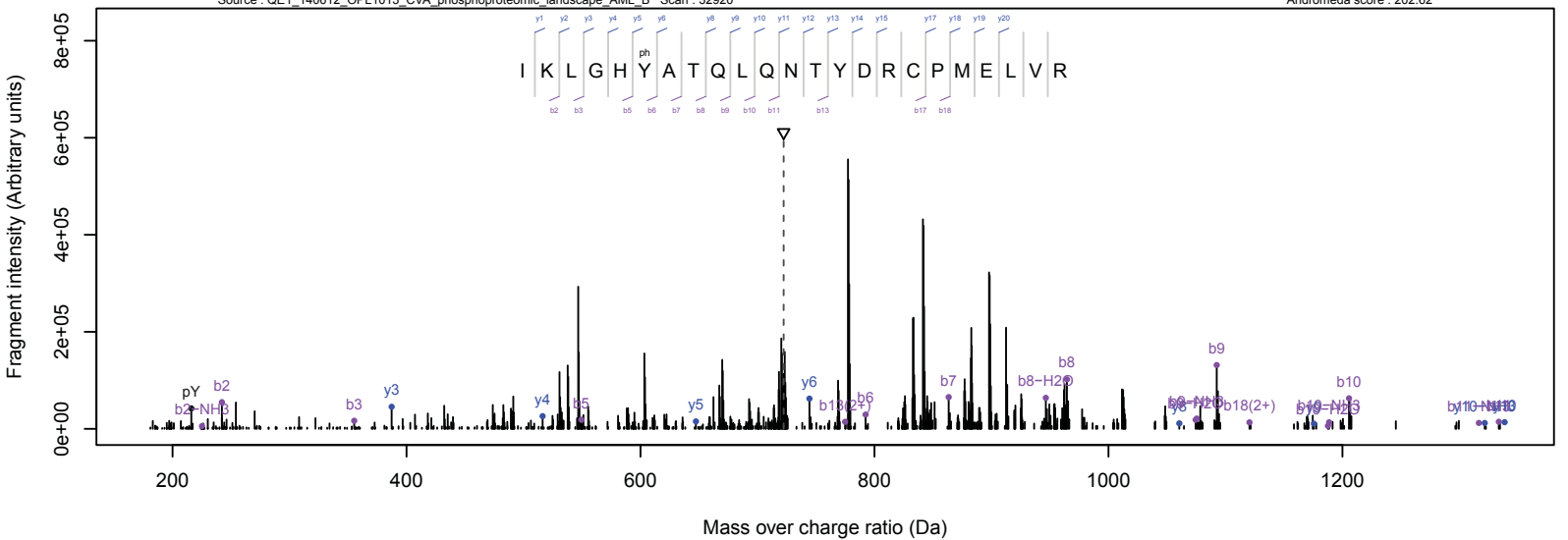
Andromeda score: 202.62



IKLGHYATQLQNTYDRCPMELVR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 32920

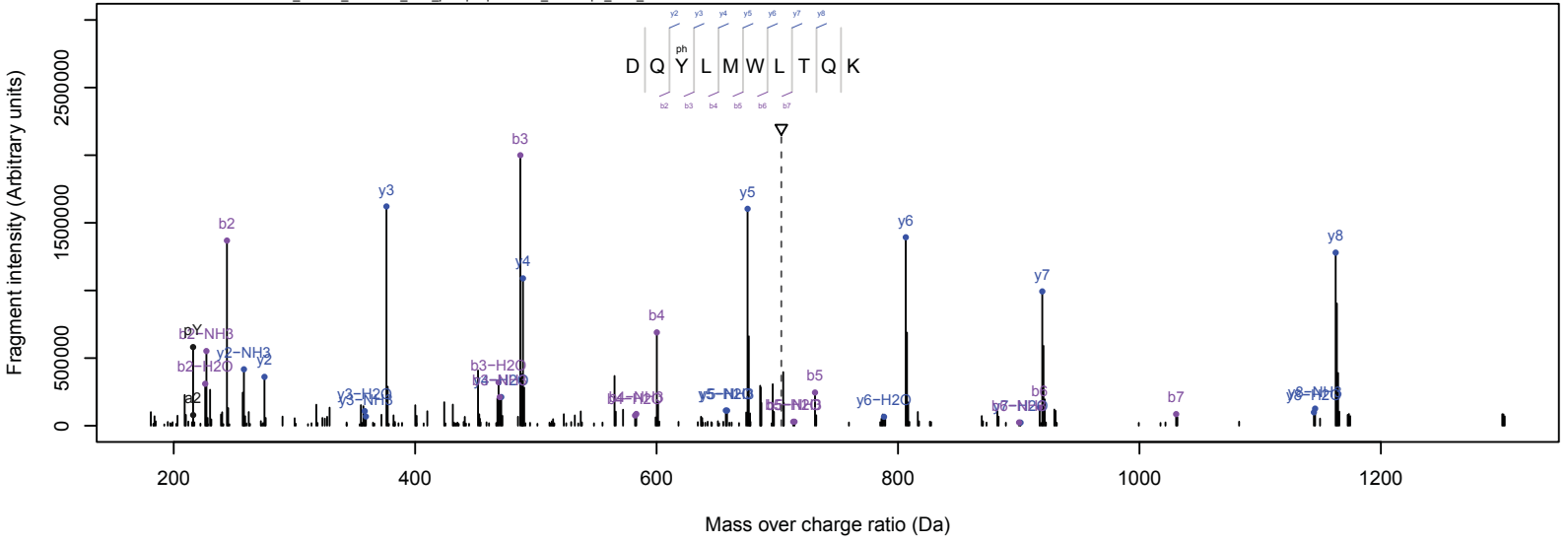
Andromeda score: 202.62



DQY(ph)LMWLTQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 40307

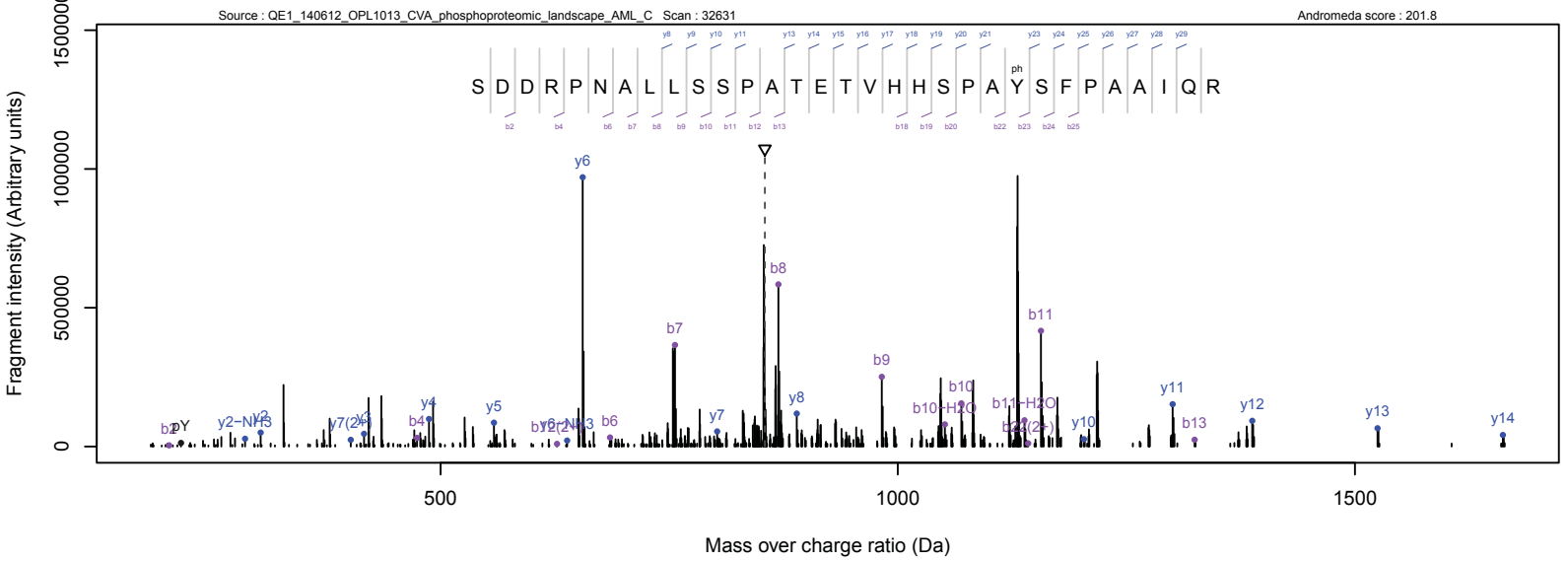
Andromeda score : 202.03



SDDRPNALLSSPATETVHHSPAY(ph)SFPAAIQR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32631

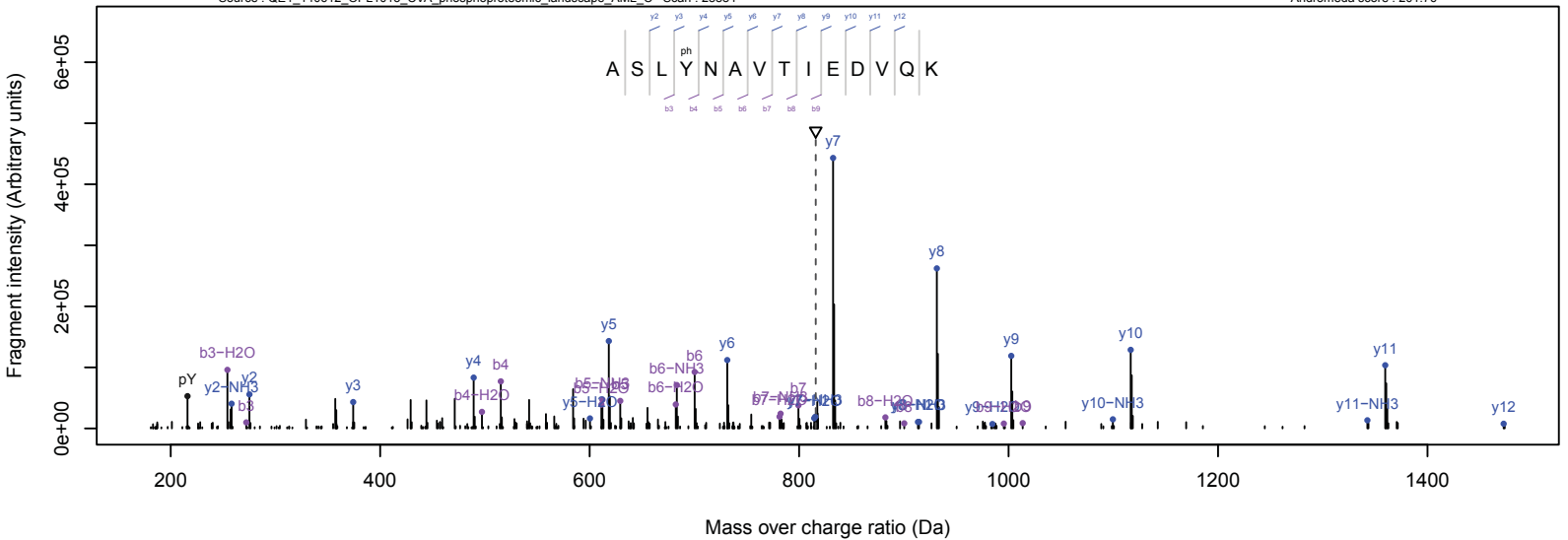
Andromeda score : 201.8



ASLY(ph)NAVTIEDVQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28884

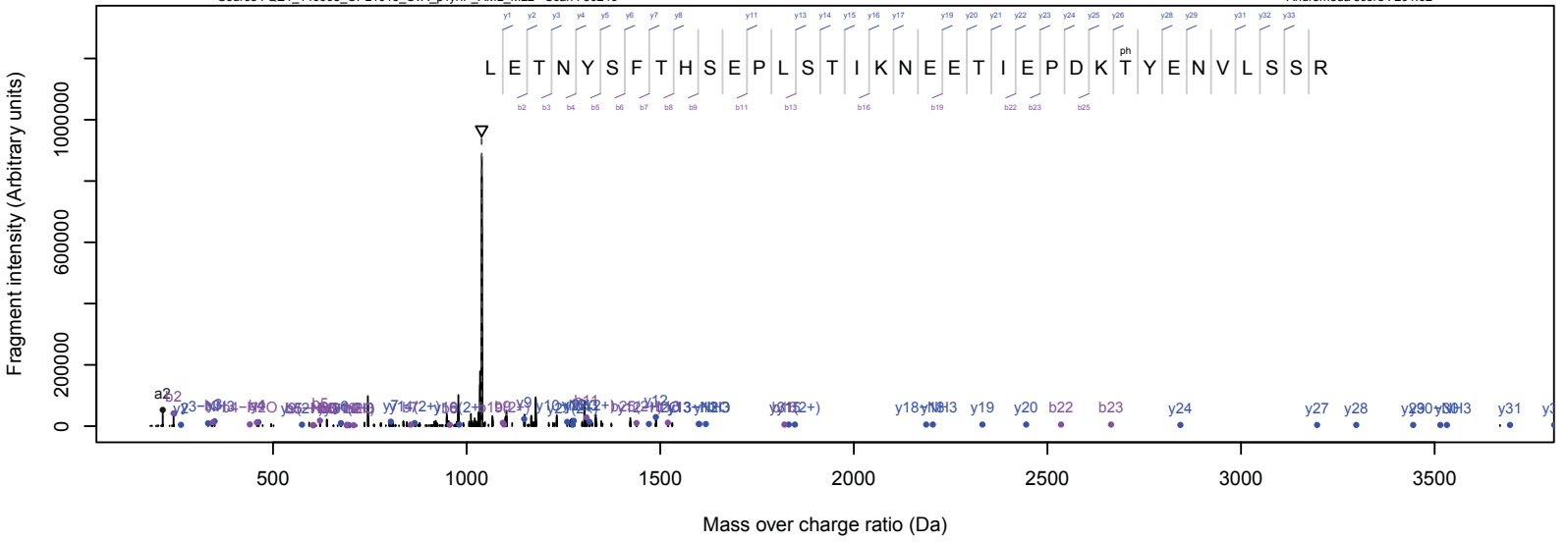
Andromeda score : 201.76



LETNYSFTHSEPLSTIKNEETIEPDKT(ph)YENVLSSR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 36246

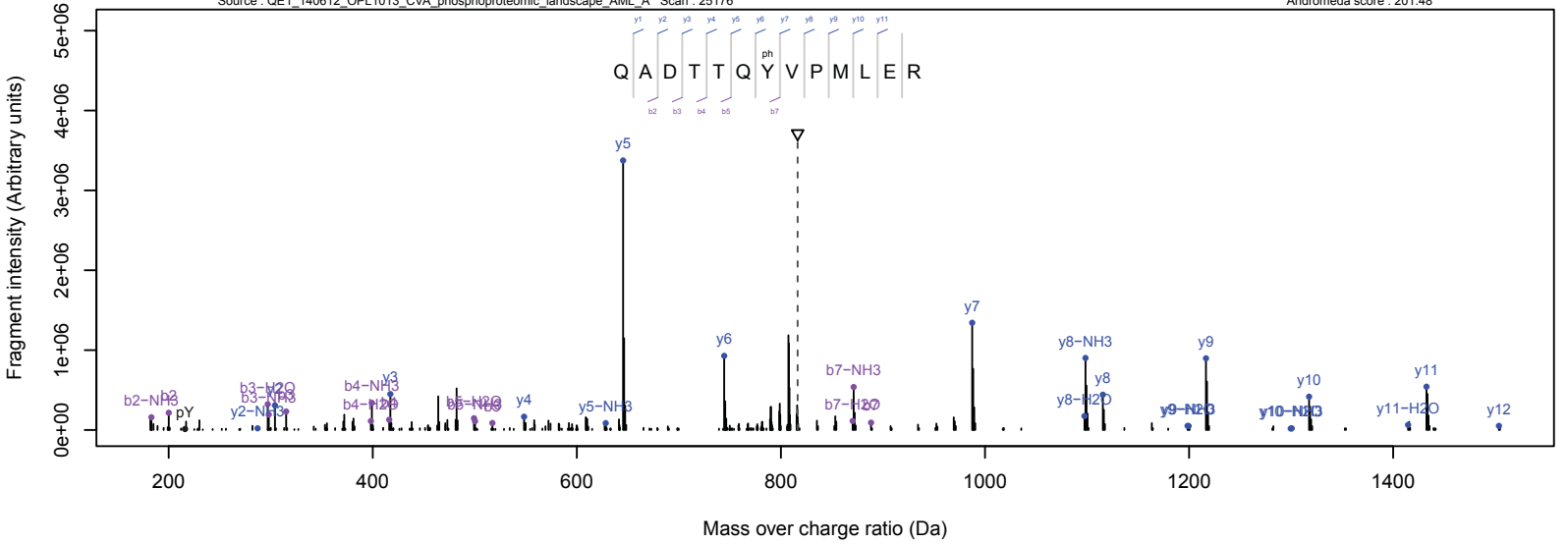
Andromeda score : 201.62



QADTTQ(ph)VPMLER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25176

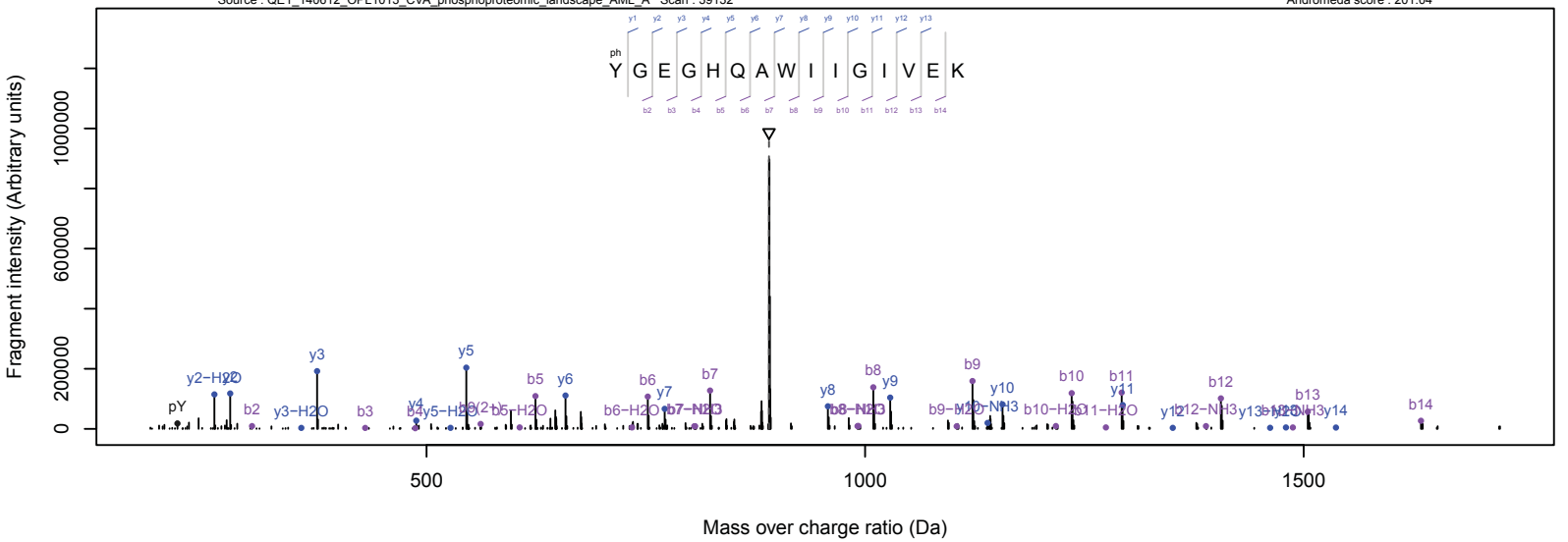
Andromeda score : 201.48



Y(ph)GEGHQAWIIGIVEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 39132

Andromeda score : 201.04

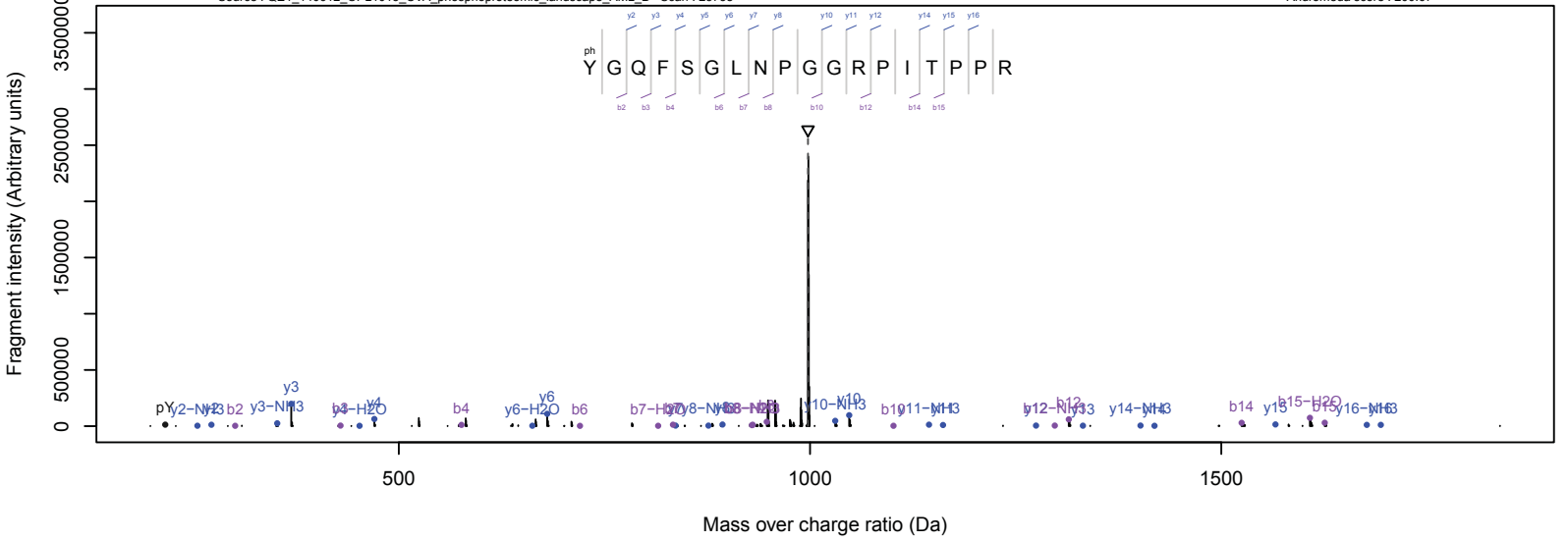




Y(ph)GQFSGLNPGGRIPPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25738

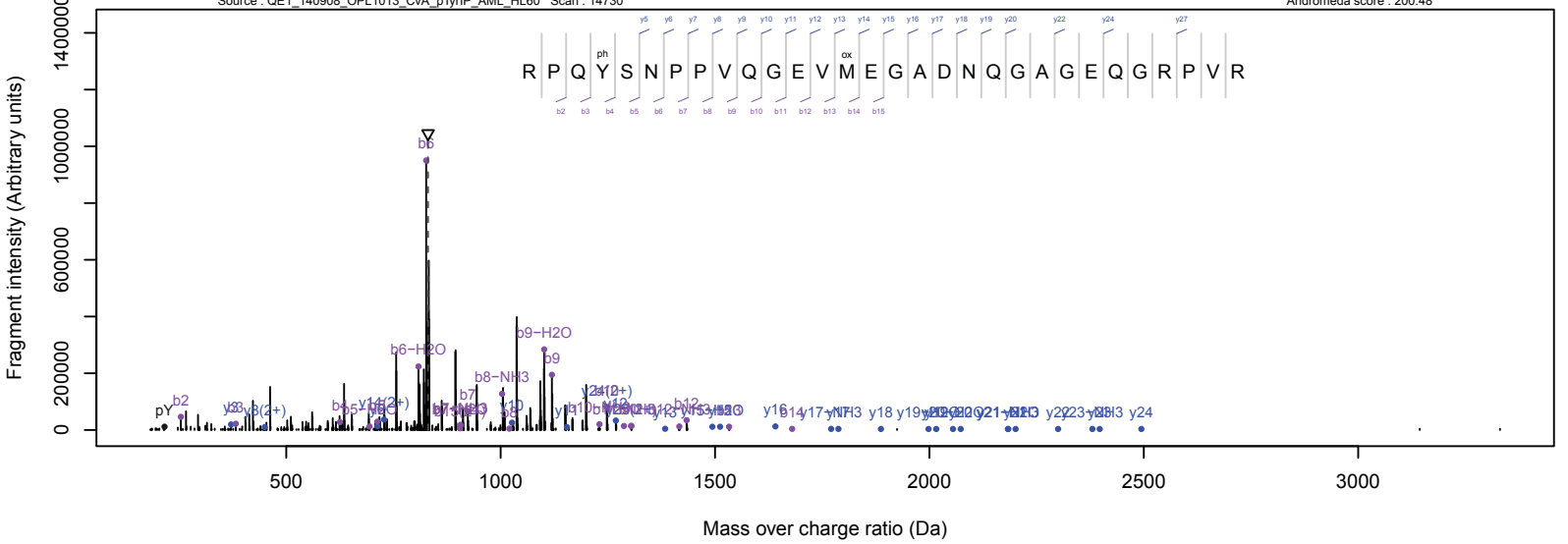
Andromeda score : 200.67



RPQY(ph)SNPPVQGEVM(ox)EGADNQGAGEQGRPVR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 14730

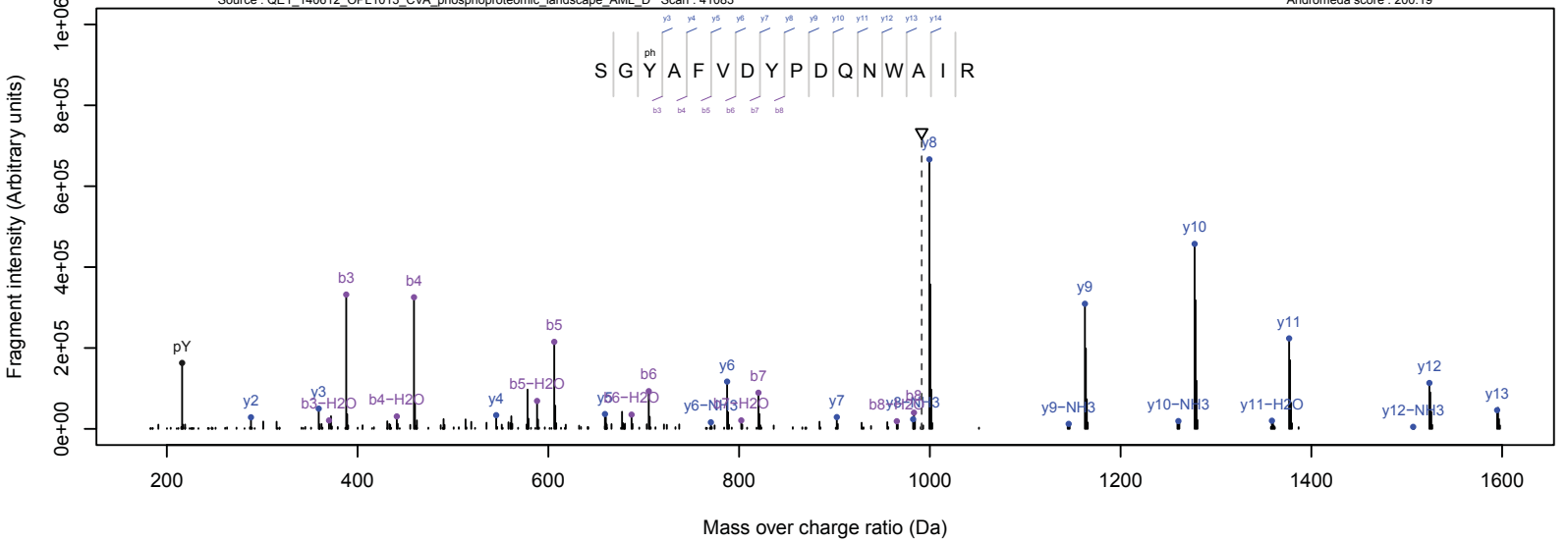
Andromeda score : 200.48



SGY(ph)AFVDYPDQNWAI R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41083

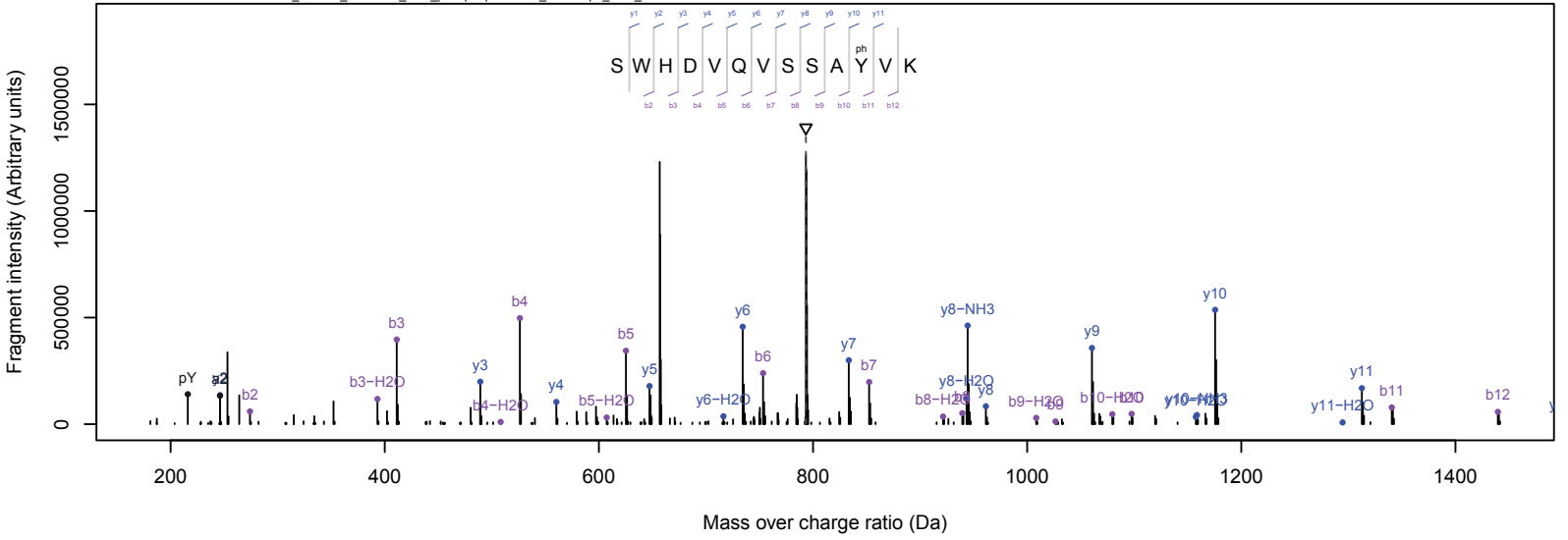
Andromeda score : 200.19



SWHDVQVSSAY(ph)VK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19063

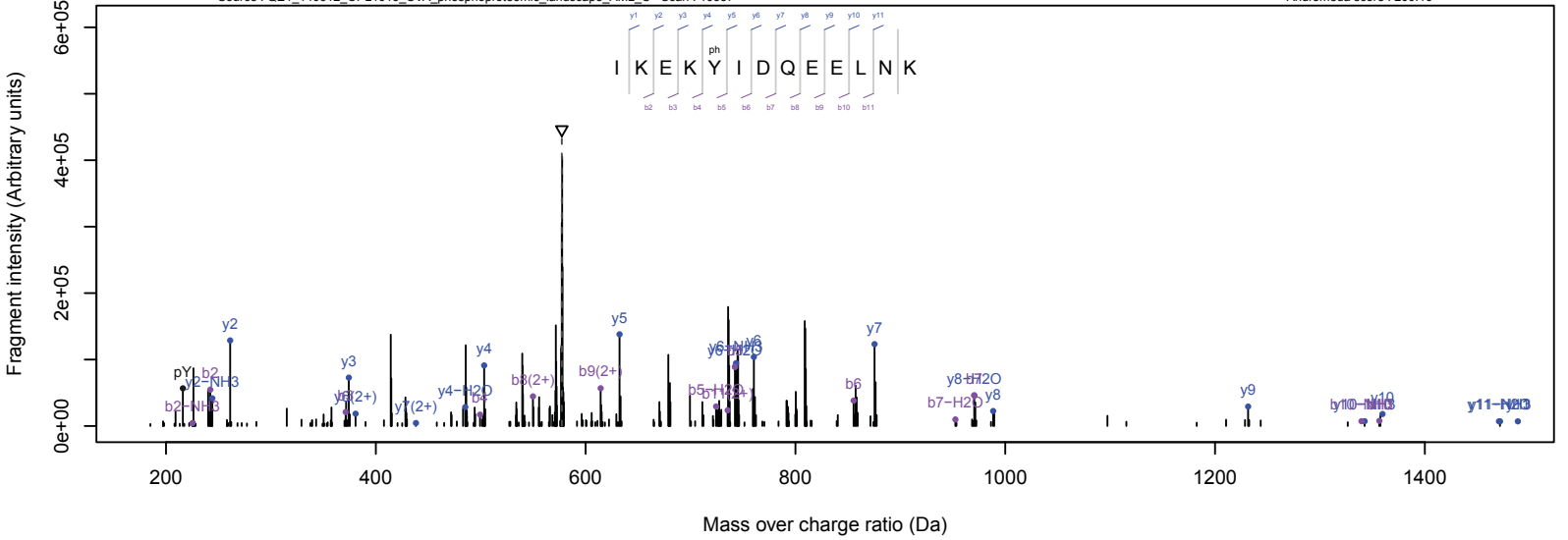
Andromeda score : 200.19



IKEKY(ph)IDQEELNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10557

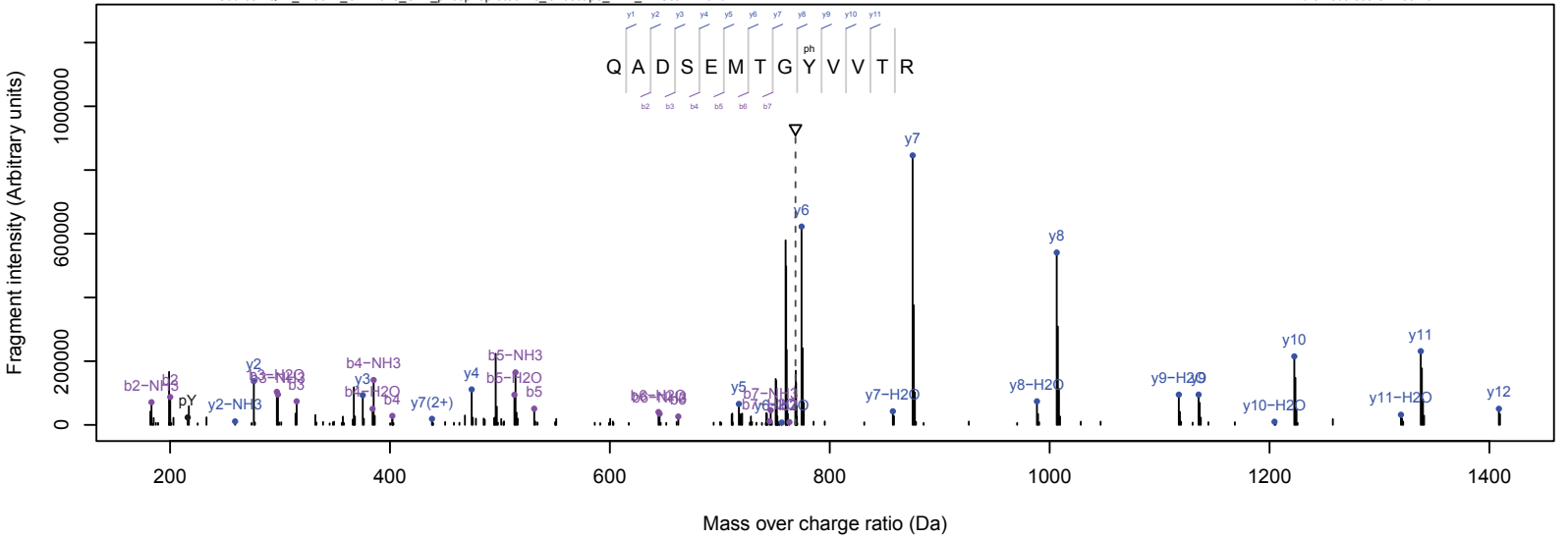
Andromeda score : 200.16



QADSEMTGY(ph)VVTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 15794

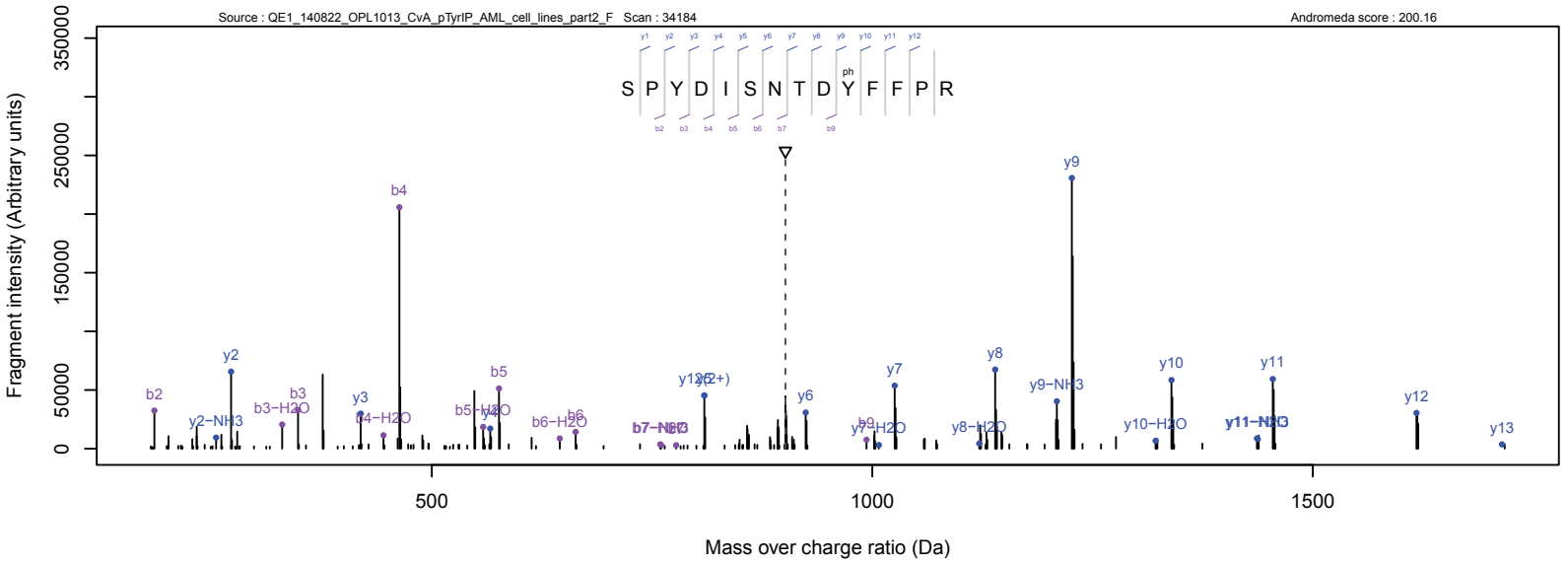
Andromeda score : 200.16



SPYDISNTDY(ph)FFPR-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 34184

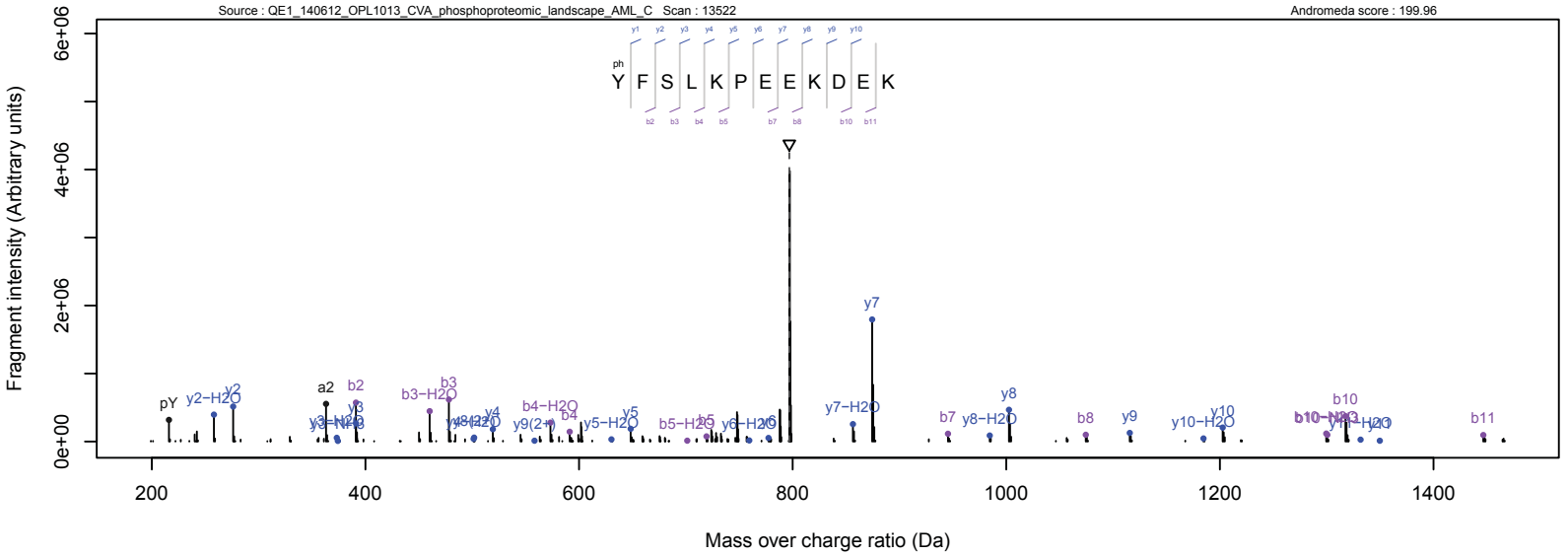
Andromeda score : 200.16



Y(ph)FSLKPEEKDEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13522

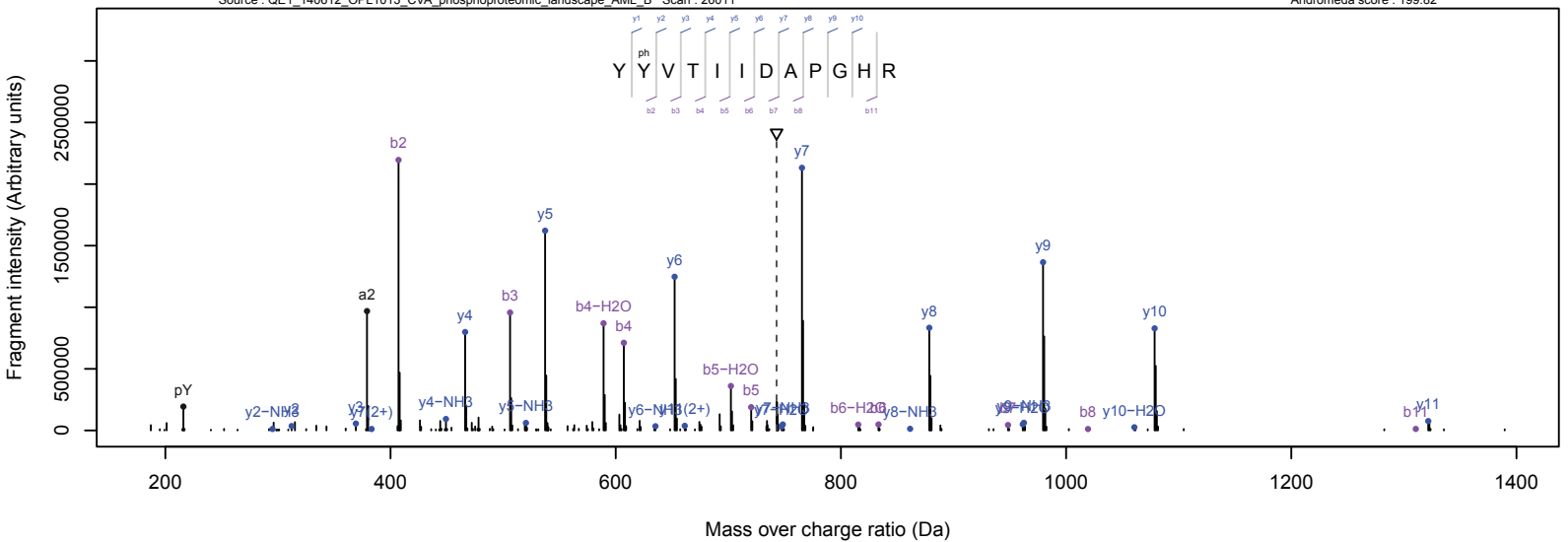
Andromeda score : 199.96



YY(ph)VTIIDAPGHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26011

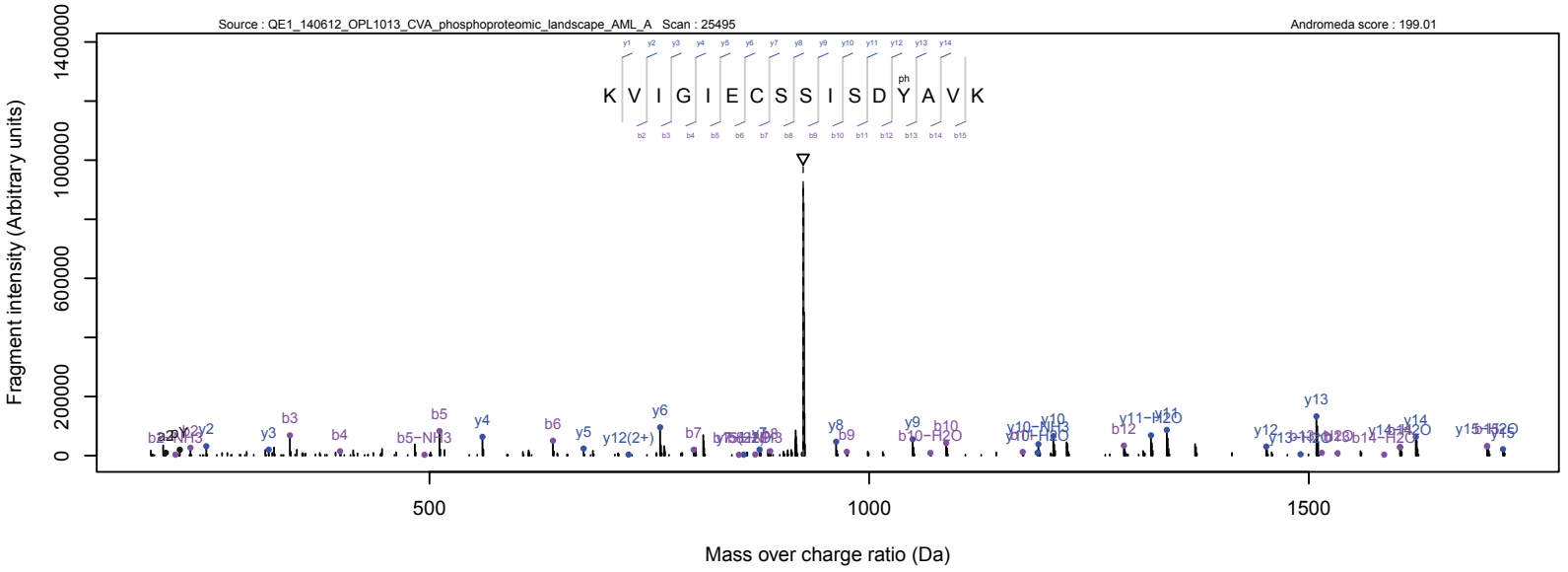
Andromeda score : 199.82



**KVIGIECSSISDY(ph)AVK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25495

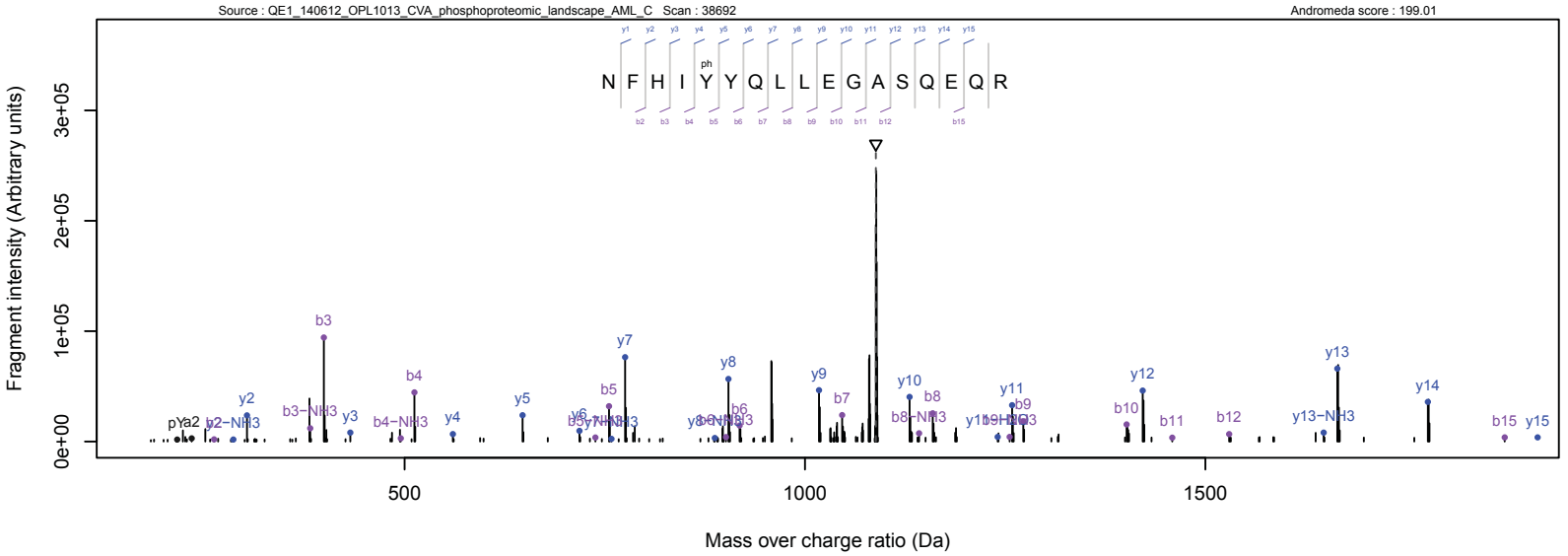
Andromeda score : 199.01



**NFHIIYQLLEGASQEQR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38692

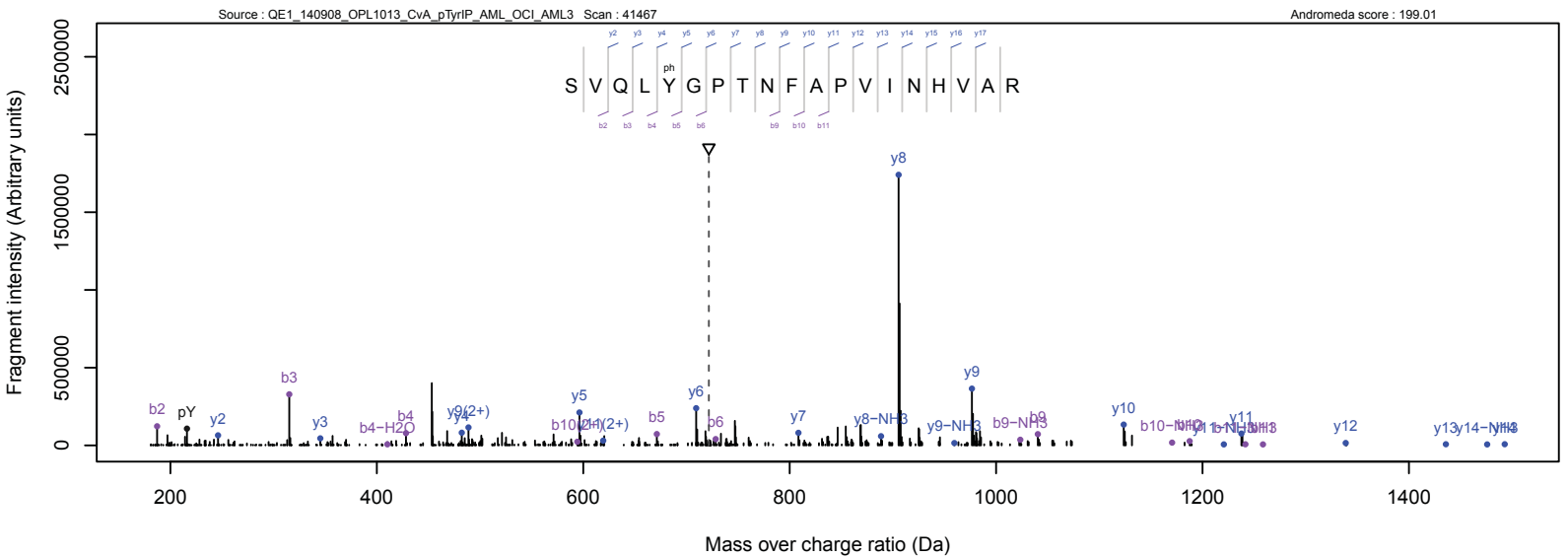
Andromeda score : 199.01



**SVQLY(ph)GPTNFAPVINHVAR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 41467

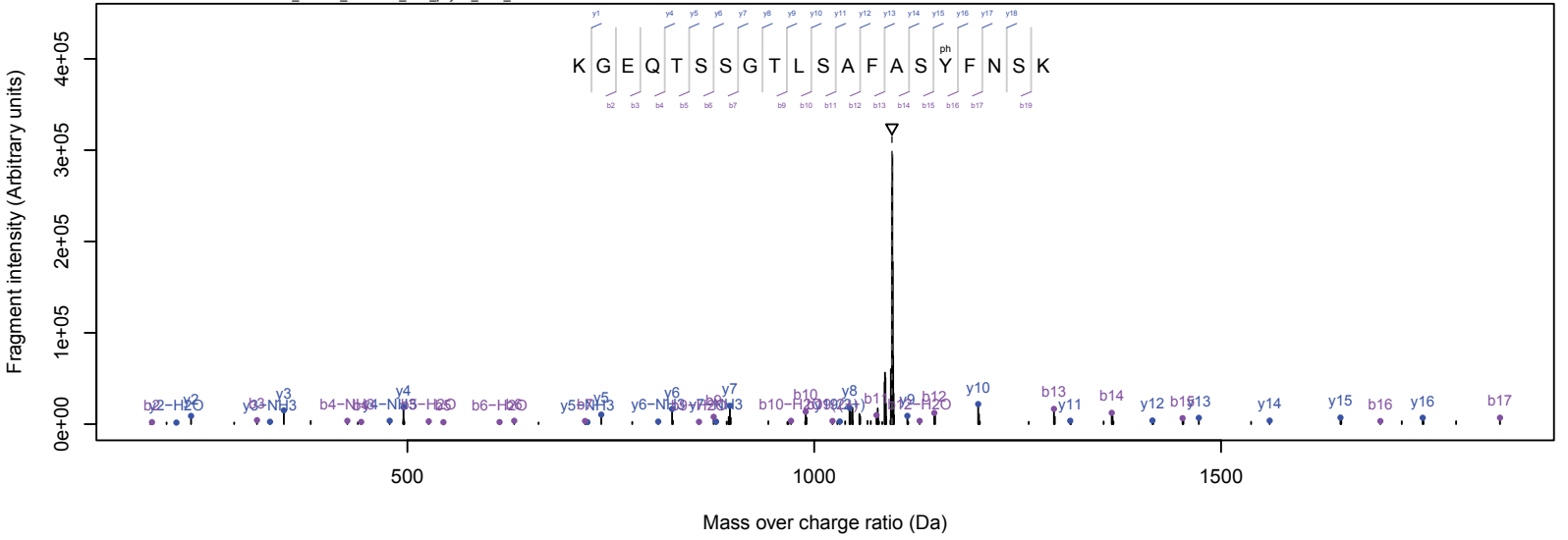
Andromeda score : 199.01



KGEQTSSGTLSAFASY(ph)FNSK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 35594

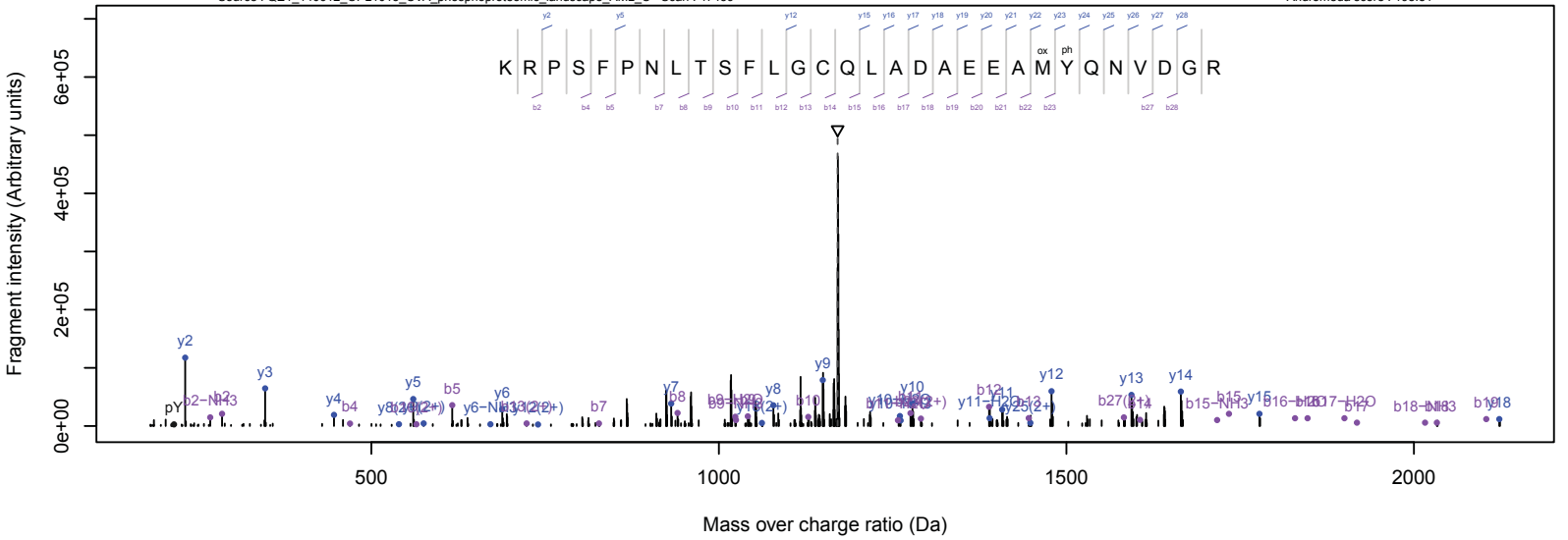
Andromeda score : 198.84



KRPSFPNLTSLGCLADAEEM(ox)Y(ph)QNV DGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47459

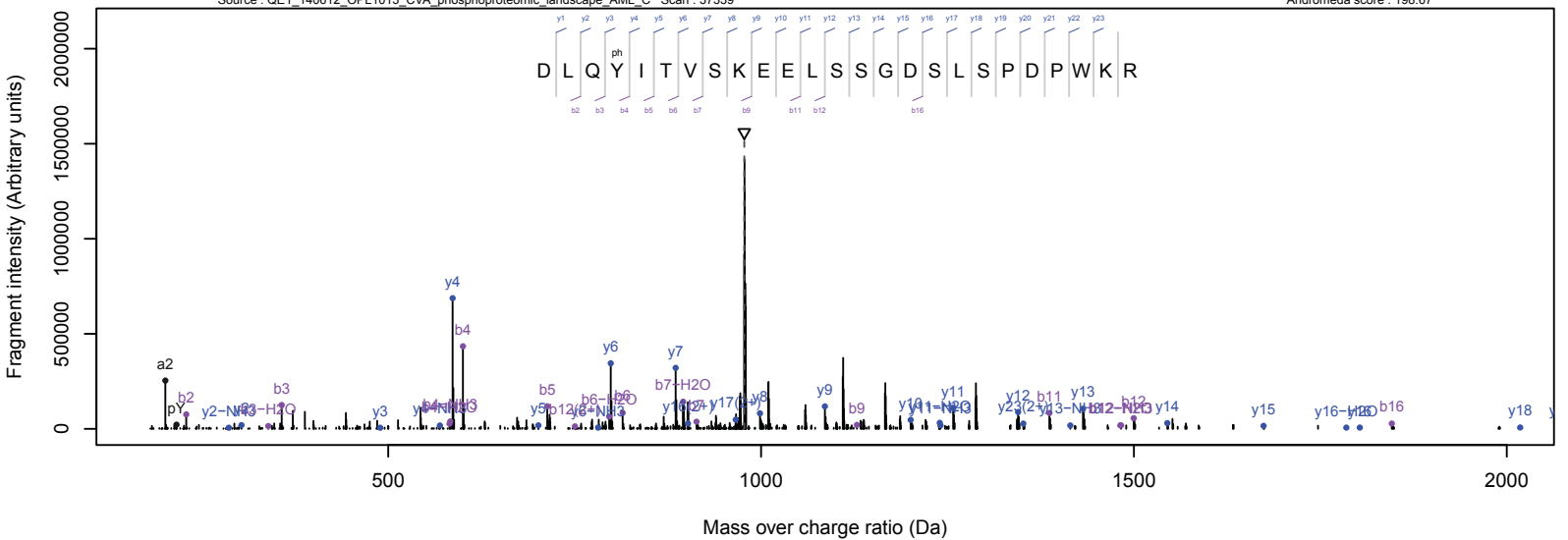
Andromeda score : 198.81



DLQY(ph)ITVSKEELSSGDSLSPDWKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37339

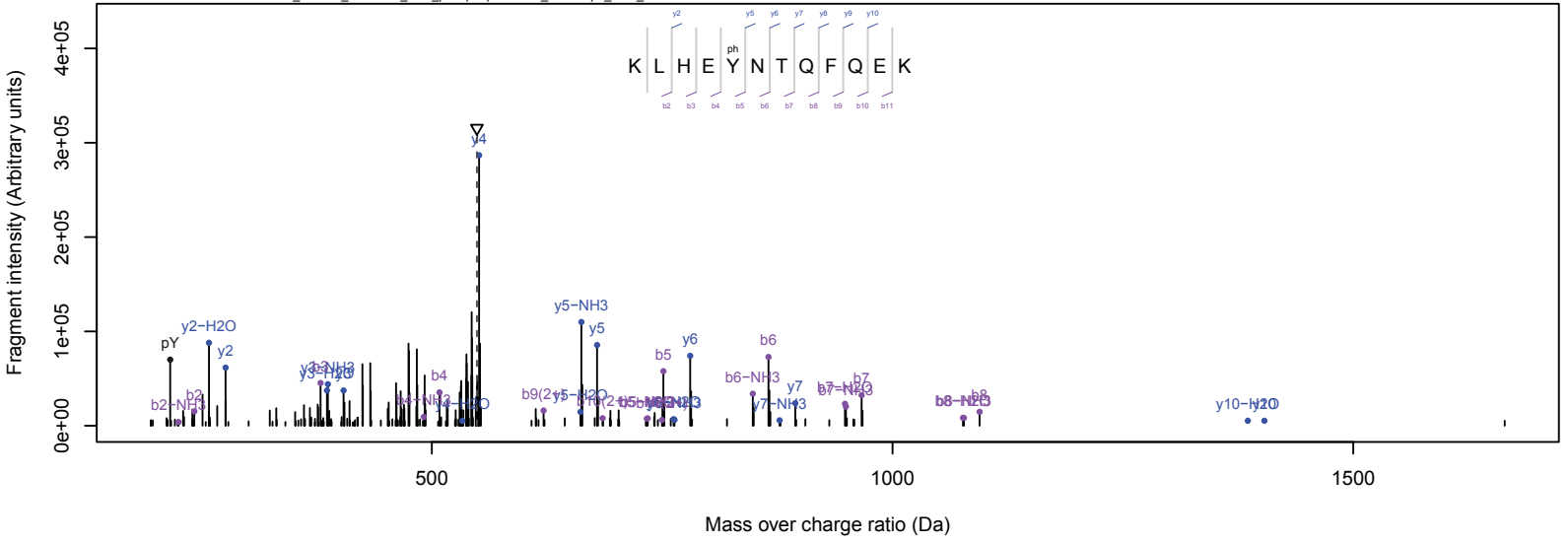
Andromeda score : 198.67



KLHEY(ph)NTQFQEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 7277

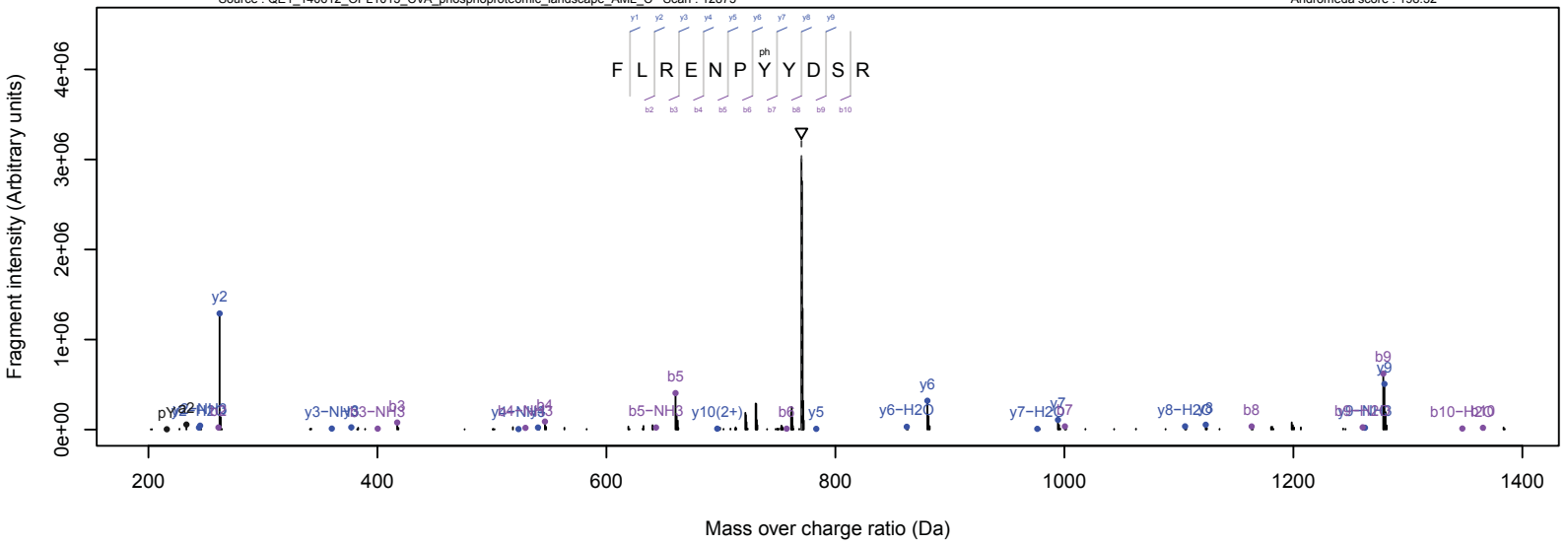
Andromeda score : 198.57



FLRENPY(ph)YDSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12875

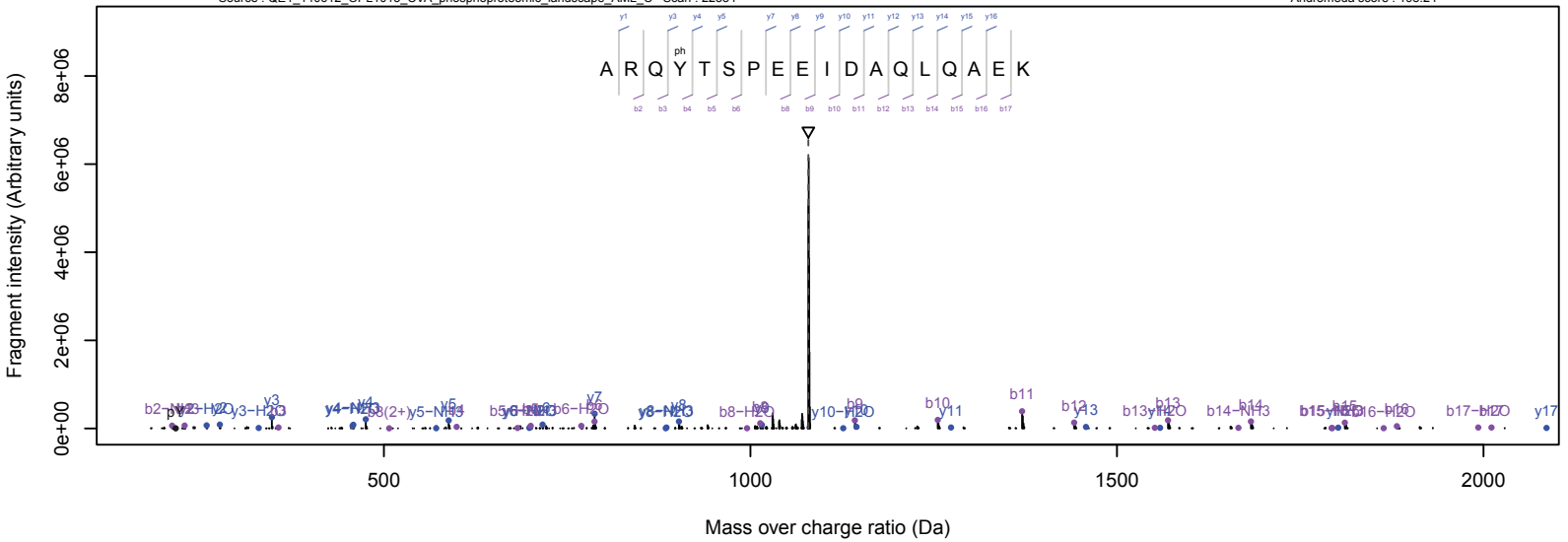
Andromeda score : 198.32



ARQY(ph)TSPEIDAQLQAEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22354

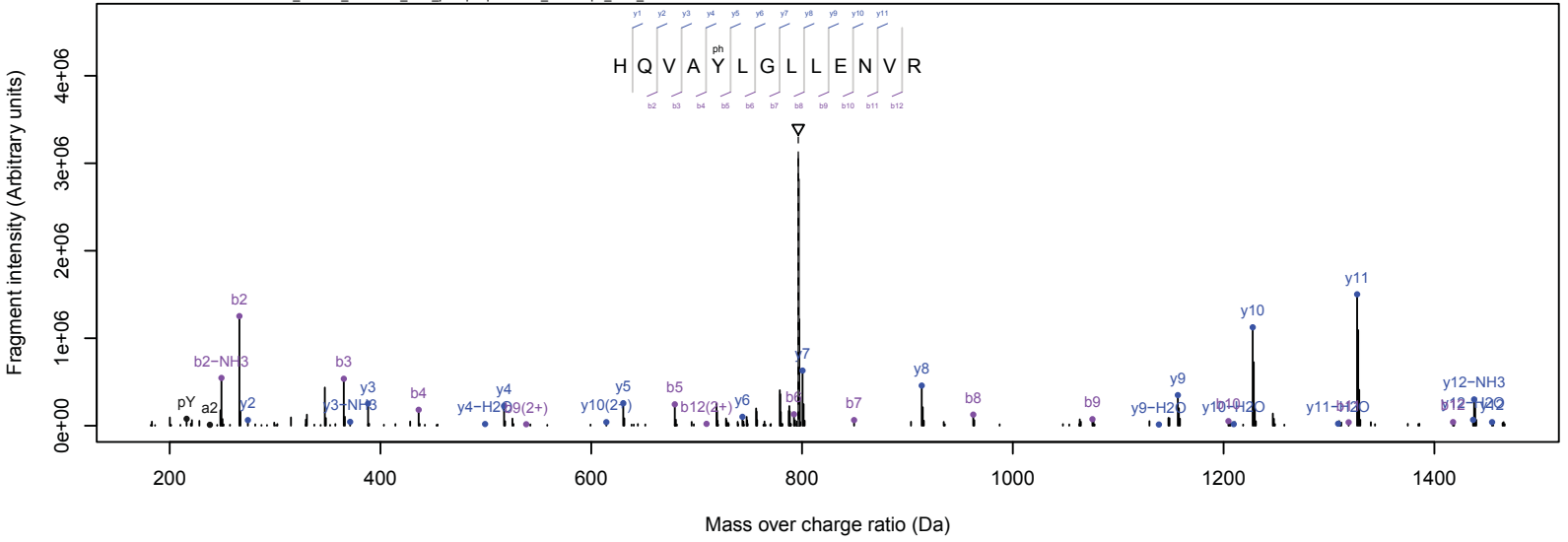
Andromeda score : 198.24



HQVAY(ph)LGLLENVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35744

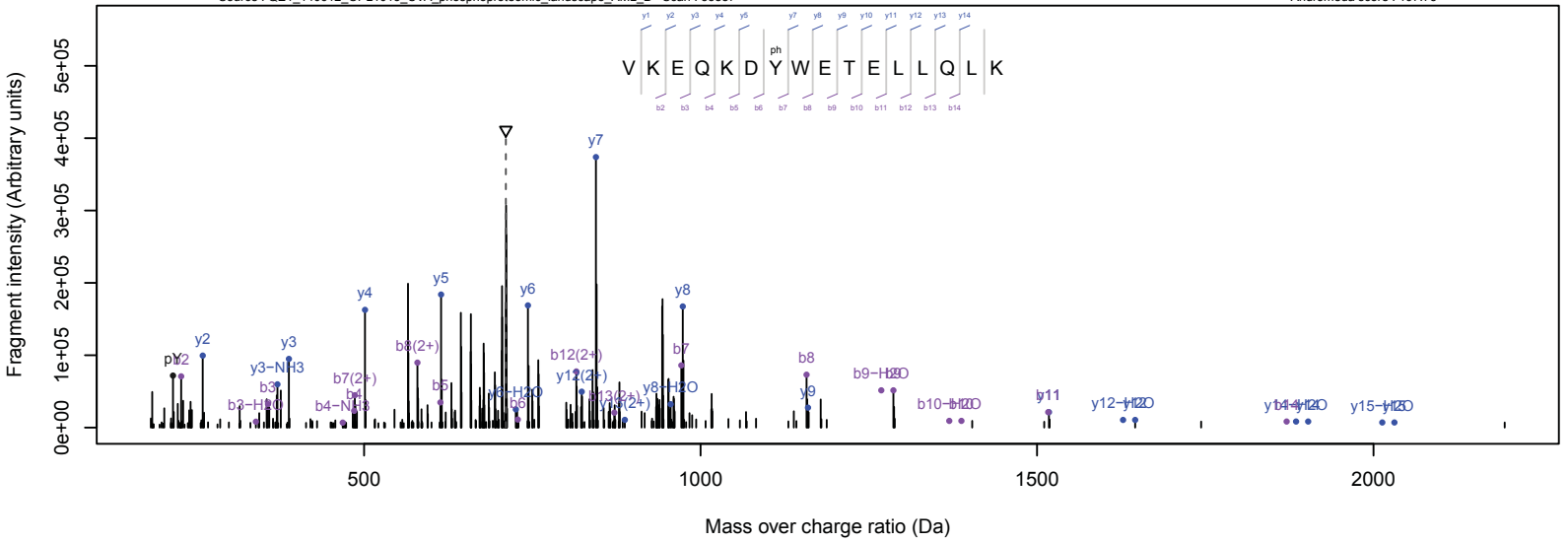
Andromeda score : 198.23



VKEQKDY(ph)WETELLQLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35887

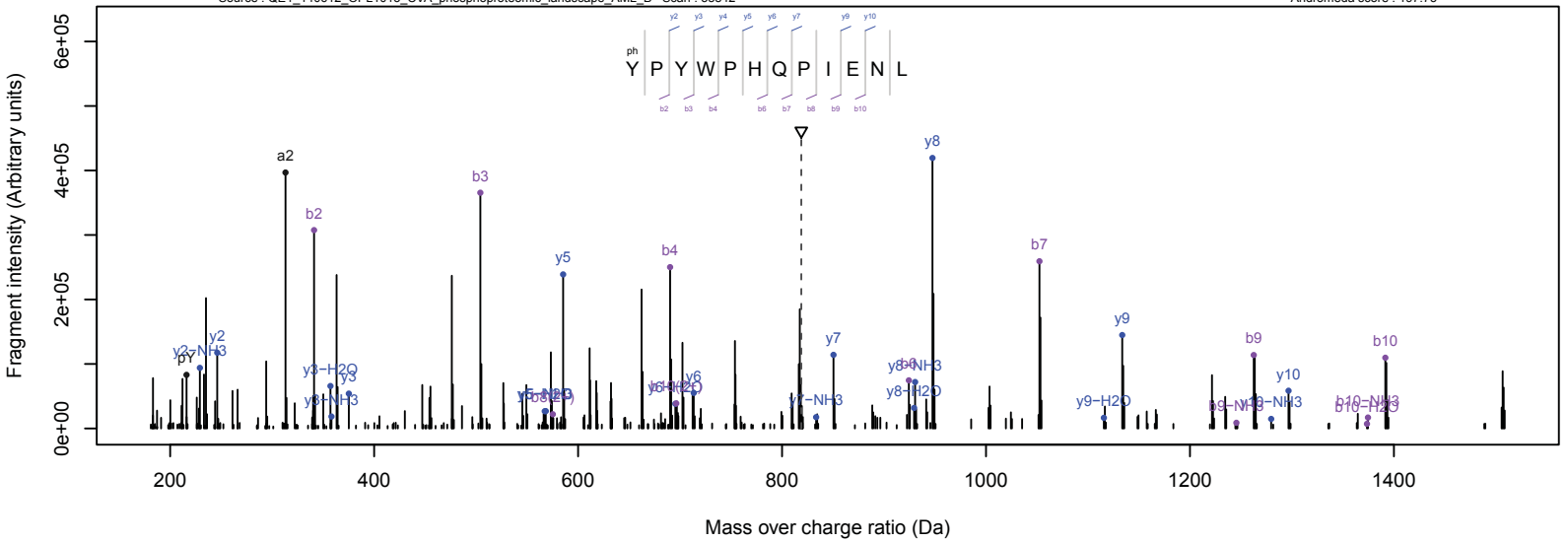
Andromeda score : 197.75



Y(ph)PYWPHQPIENL-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38842

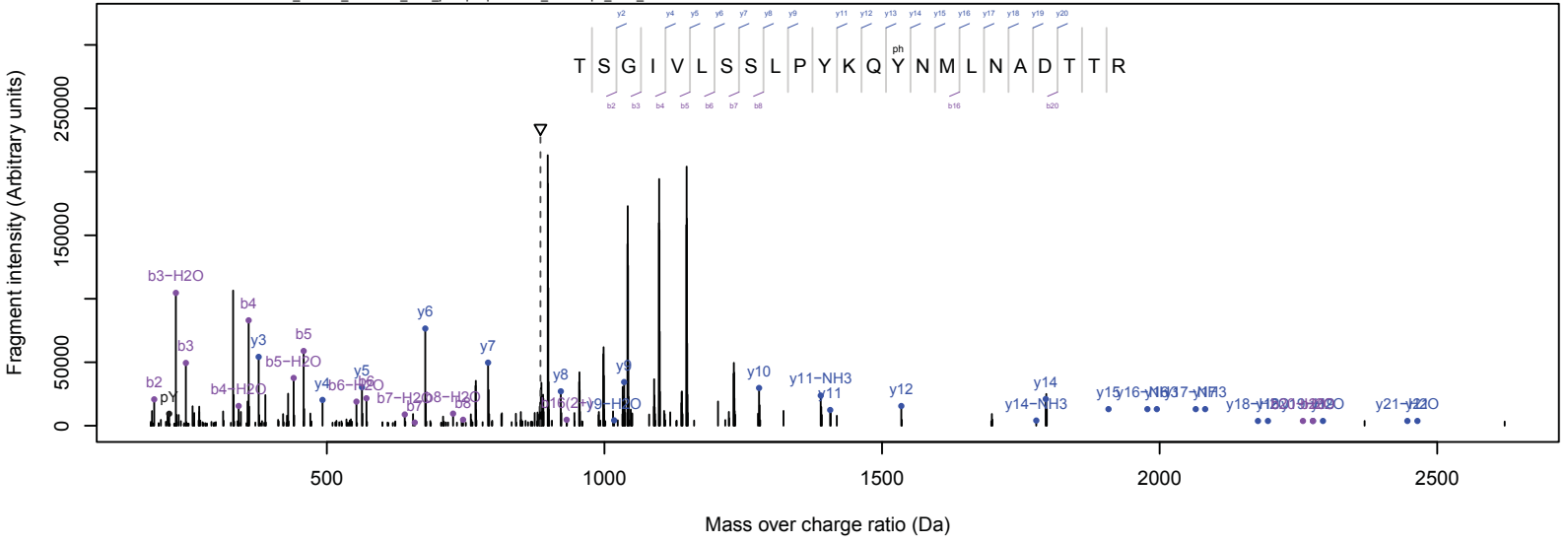
Andromeda score : 197.73



**TSGIVLSSLPYKQ(ph)NMLNADTTR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38232

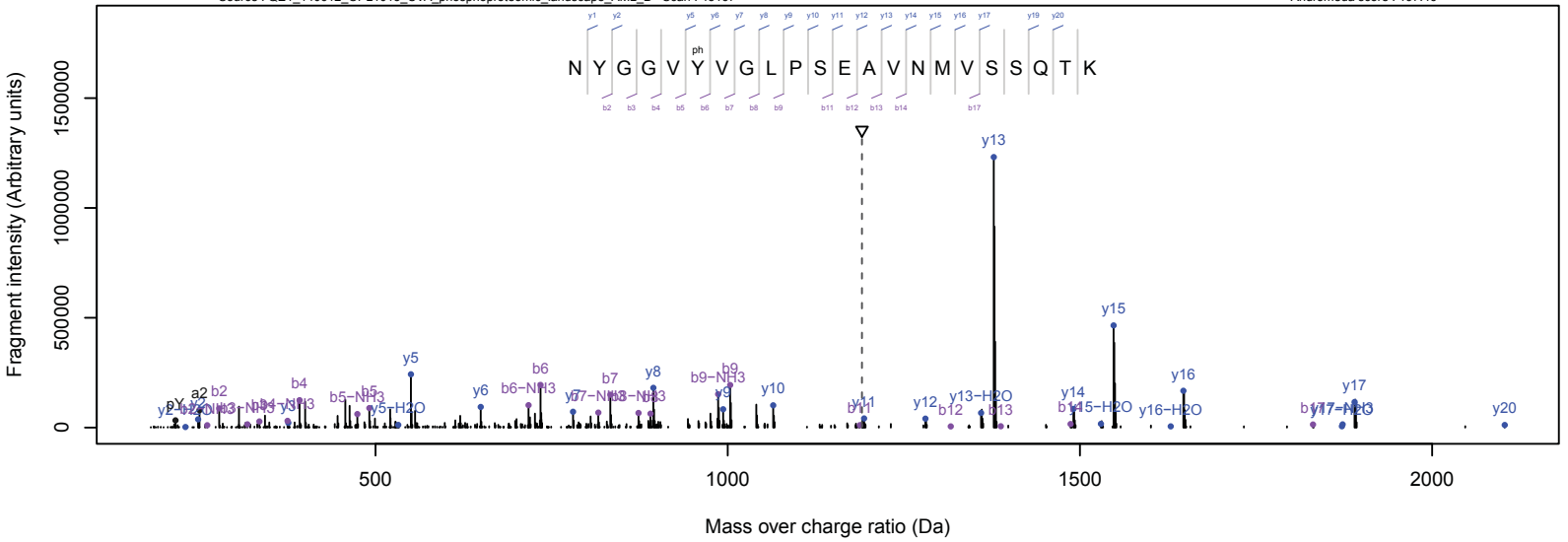
Andromeda score : 197.68



**NYGGVY(ph)VGLPSEAVNMVSSQTK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43167

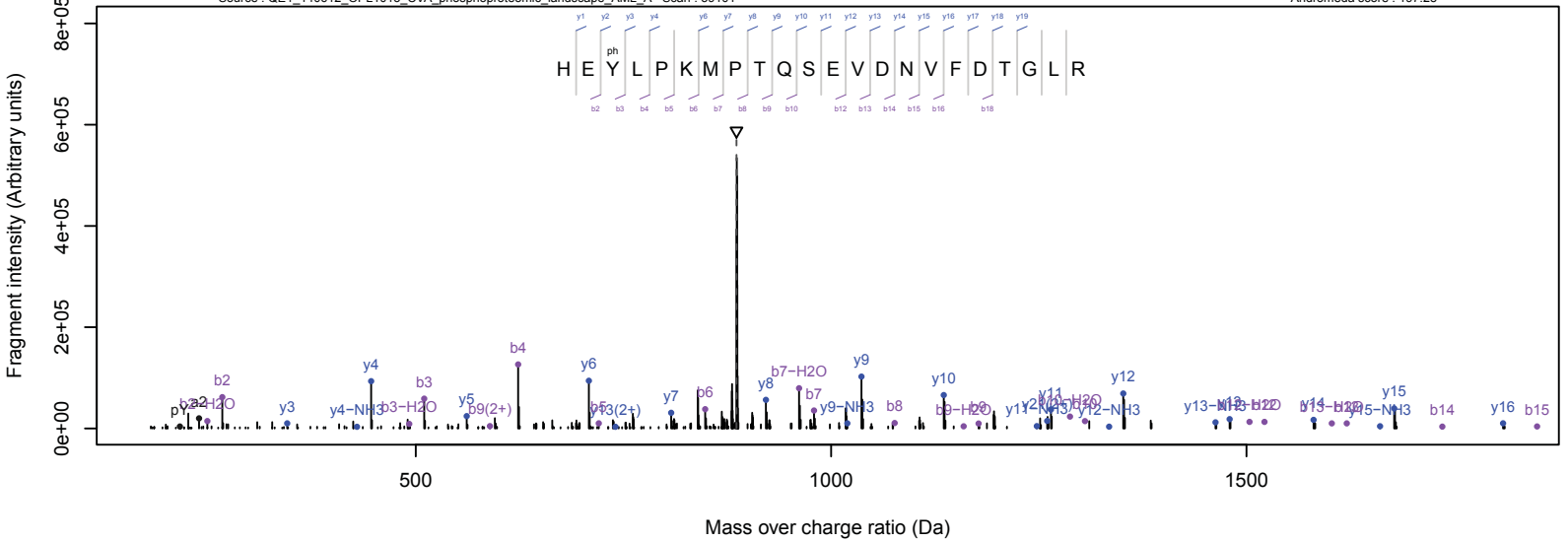
Andromeda score : 197.46



**HEY(ph)LPKMPTQSEVDNVFD TGLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 39101

Andromeda score : 197.28

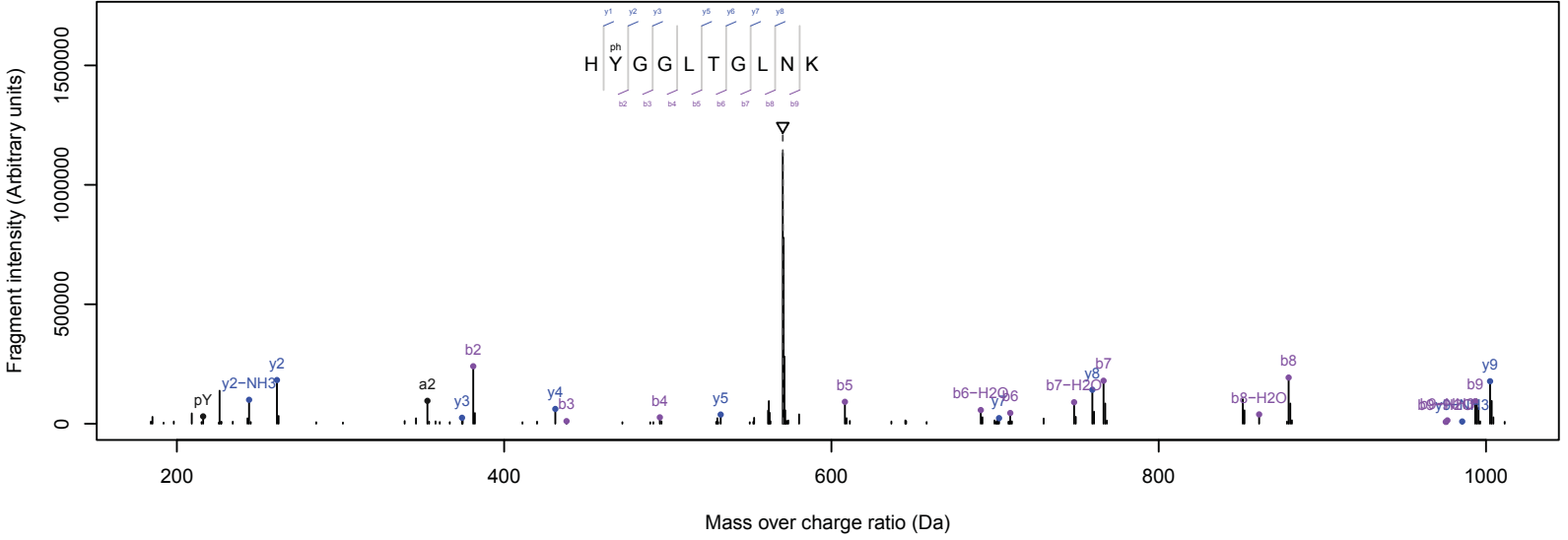




HY(ph)GGLTGLNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9802

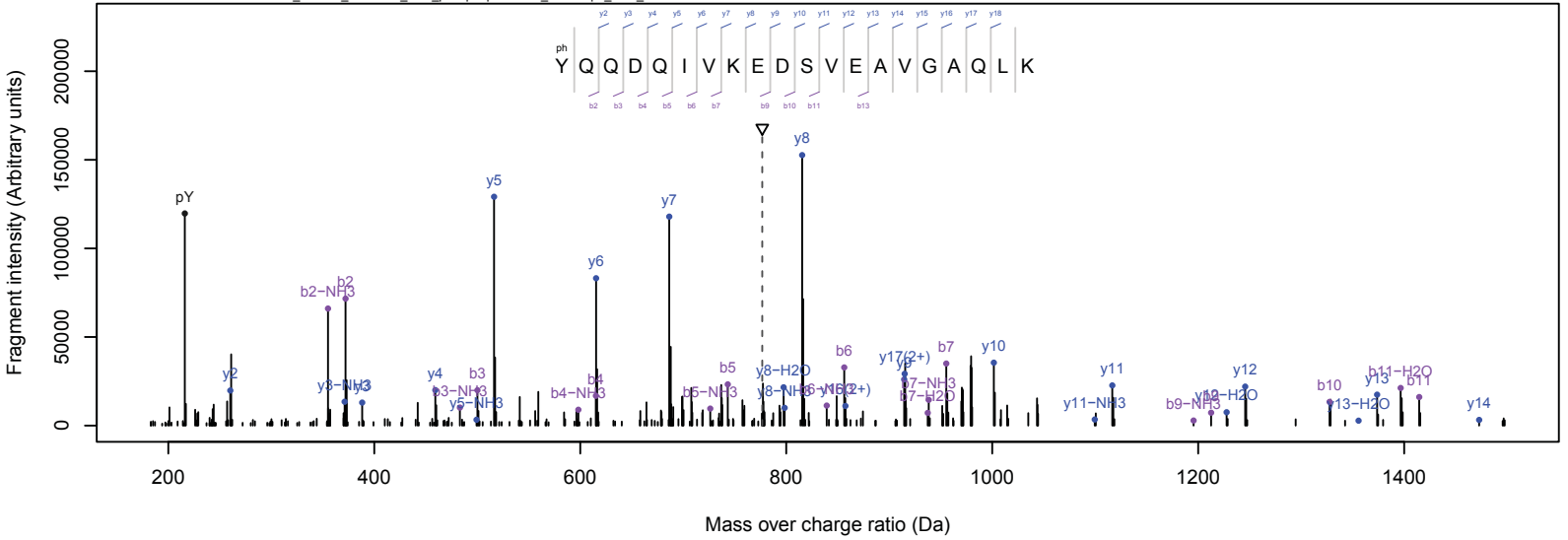
Andromeda score : 197.27



Y(ph)QQDQIVKEDSVEAVGAQLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27809

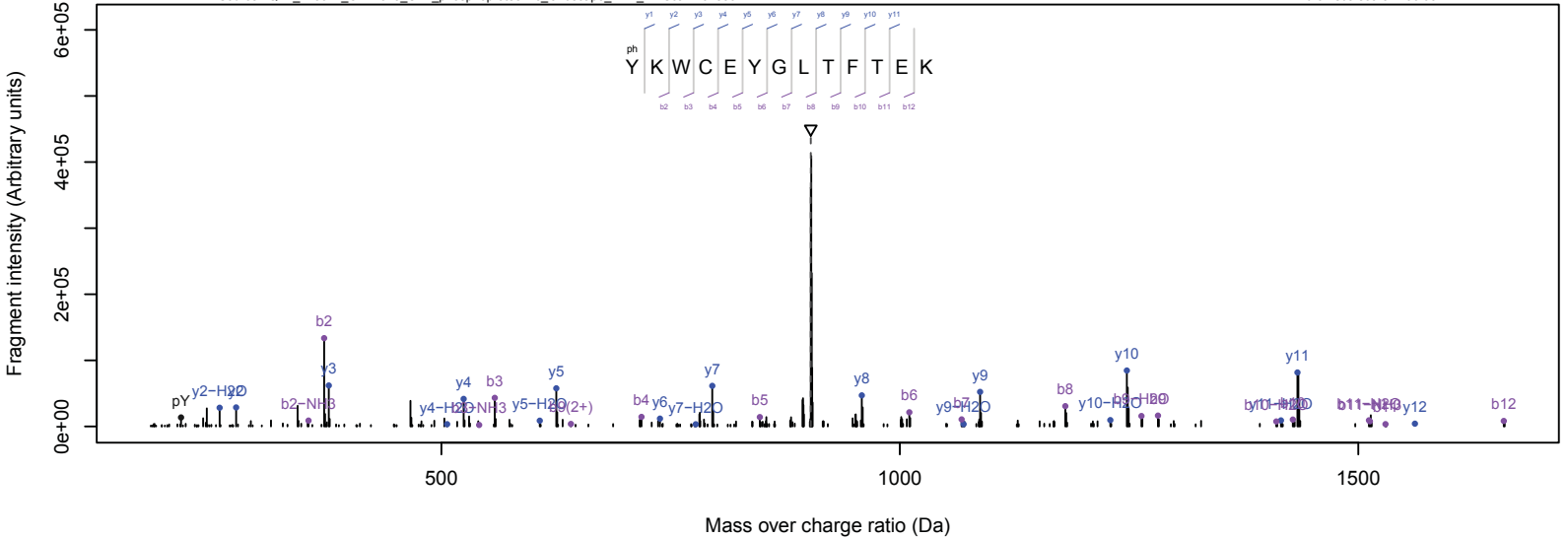
Andromeda score : 197.1



Y(ph)KWCEYGLTFTEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 32850

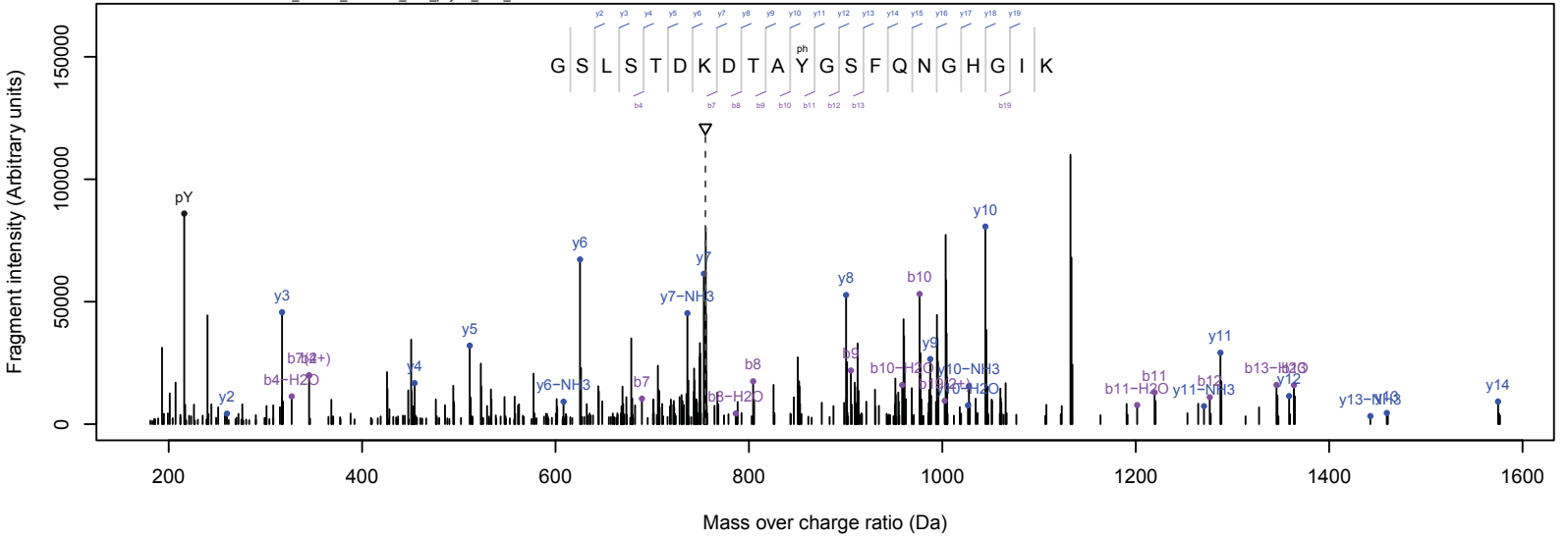
Andromeda score : 196.95



**GSLSTDKTAY(ph)GSFQNGHGIK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 15754

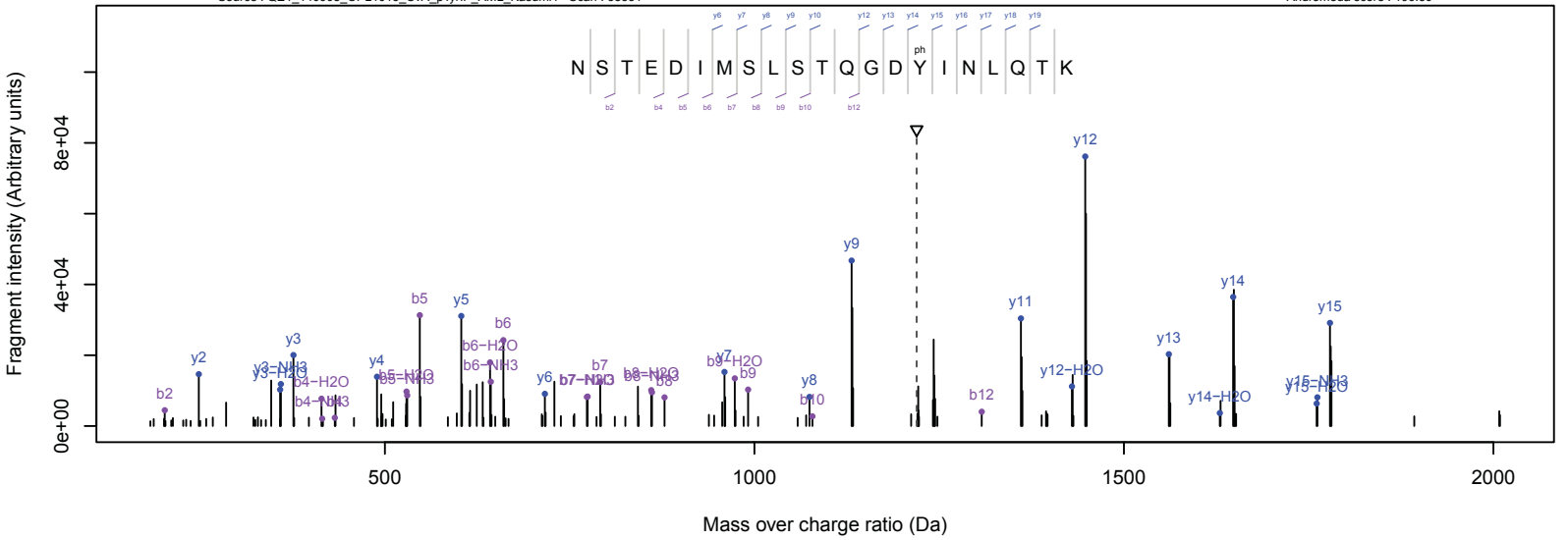
Andromeda score : 196.89



**NSTEDIMSLSTQGDY(ph)INLQTK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 38351

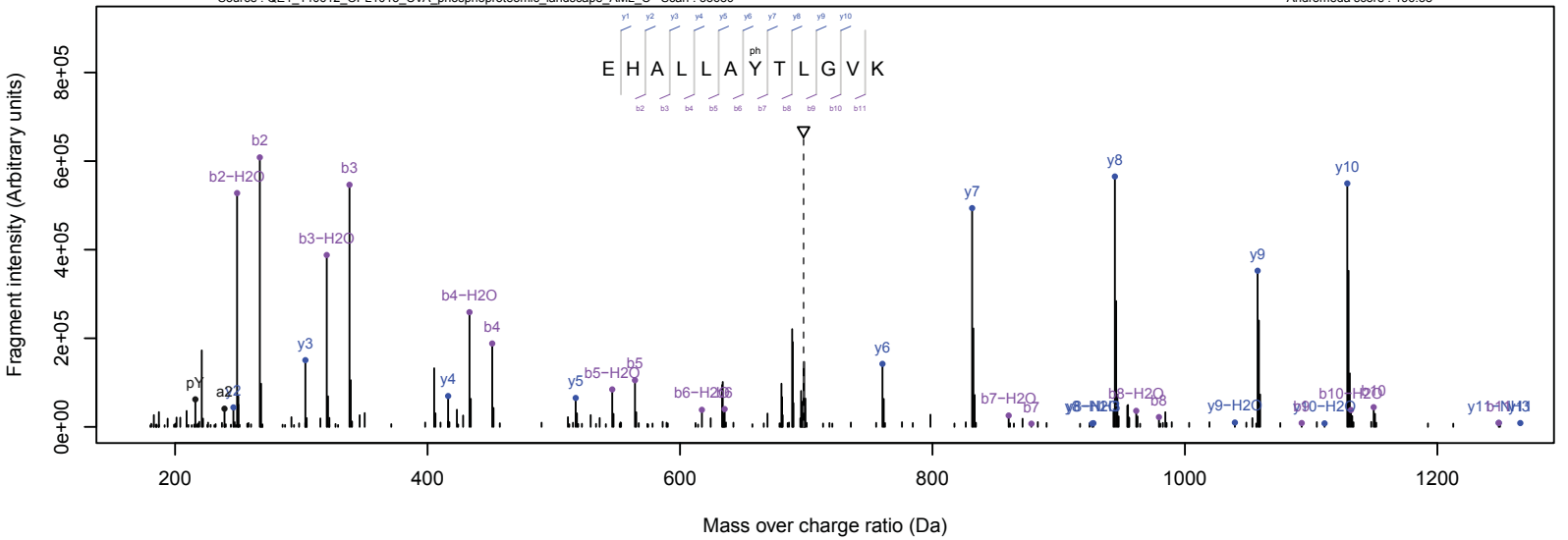
Andromeda score : 196.89



**EHALLAY(ph)TLGVK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33689

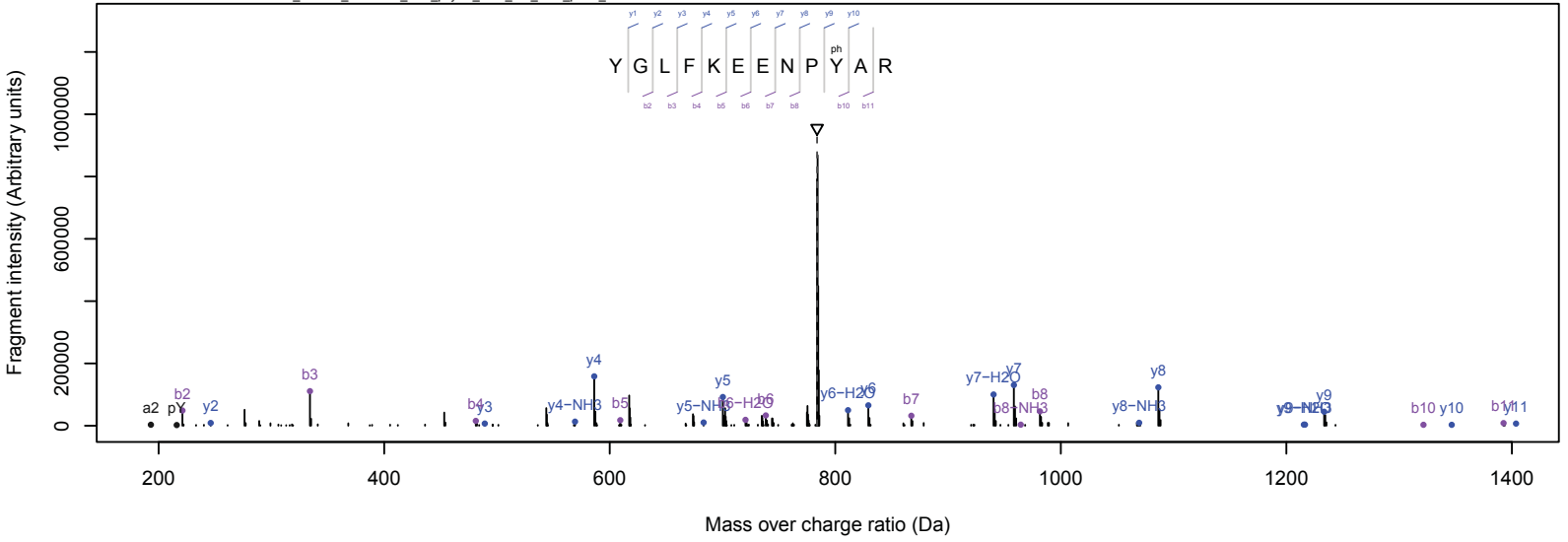
Andromeda score : 196.38



YGLFKEENPY(ph)AR-2+

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan: 14420

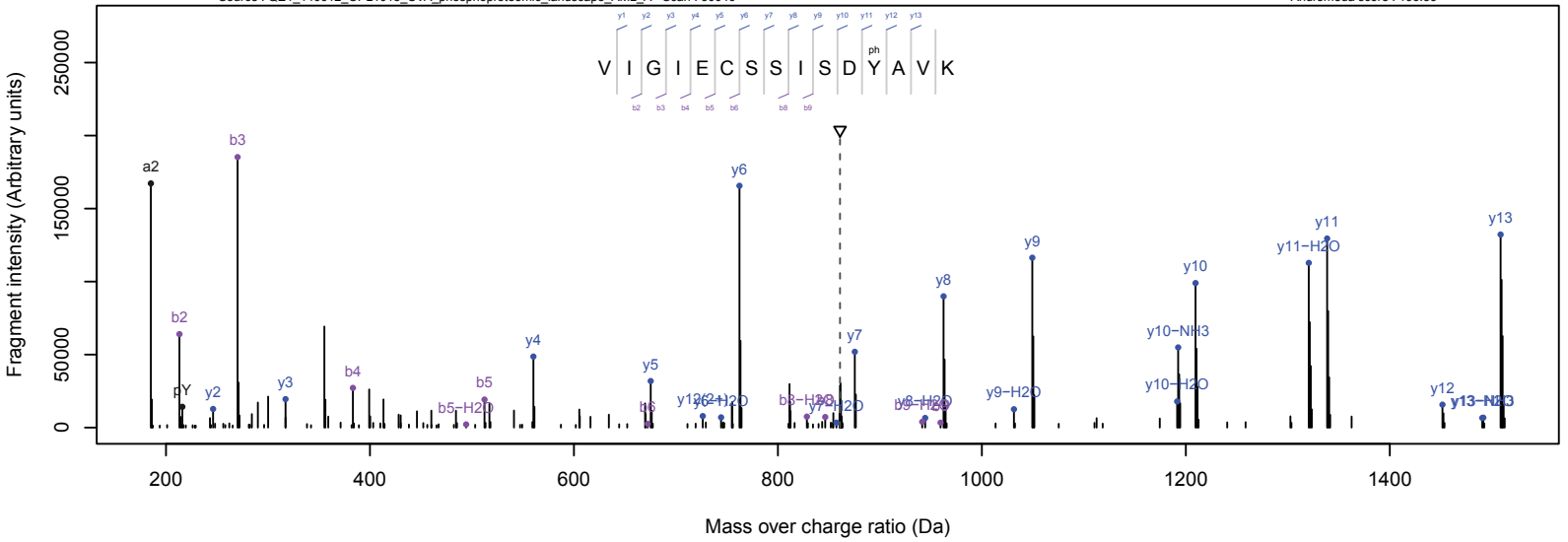
Andromeda score: 196.38



VIGIECSSISD(ph)AVK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 30046

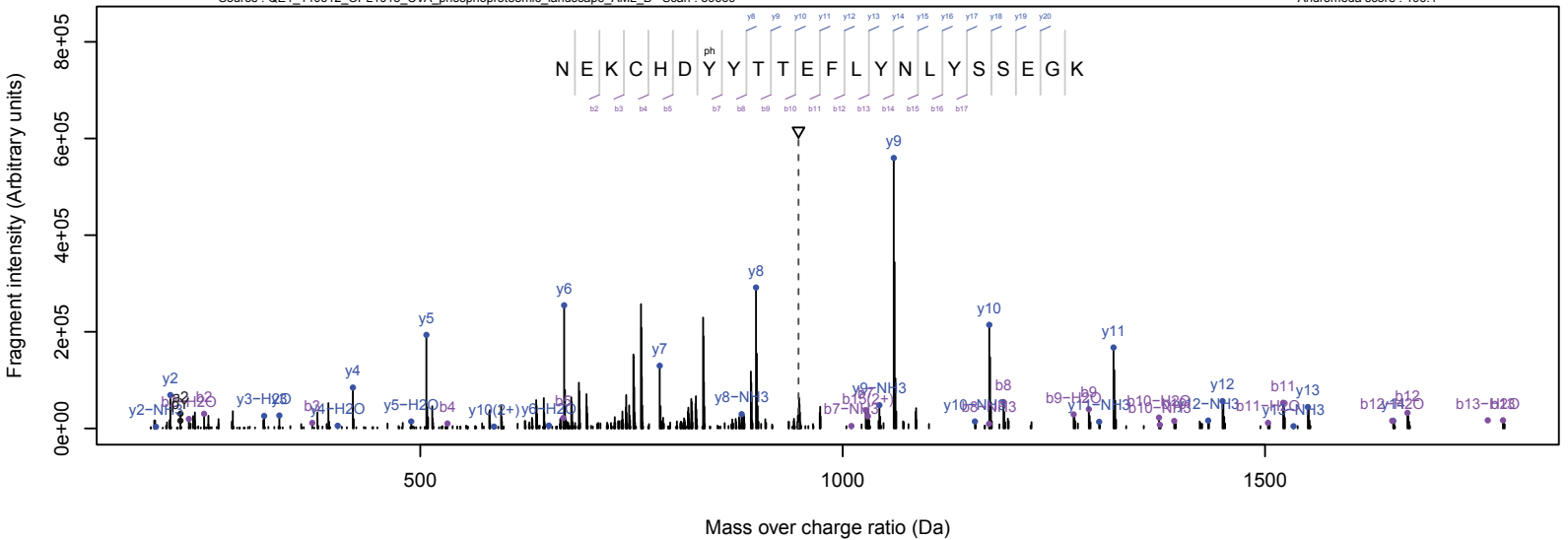
Andromeda score: 196.33



NEKCHDY(ph)YTTEFLYNLYSSEGK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 39669

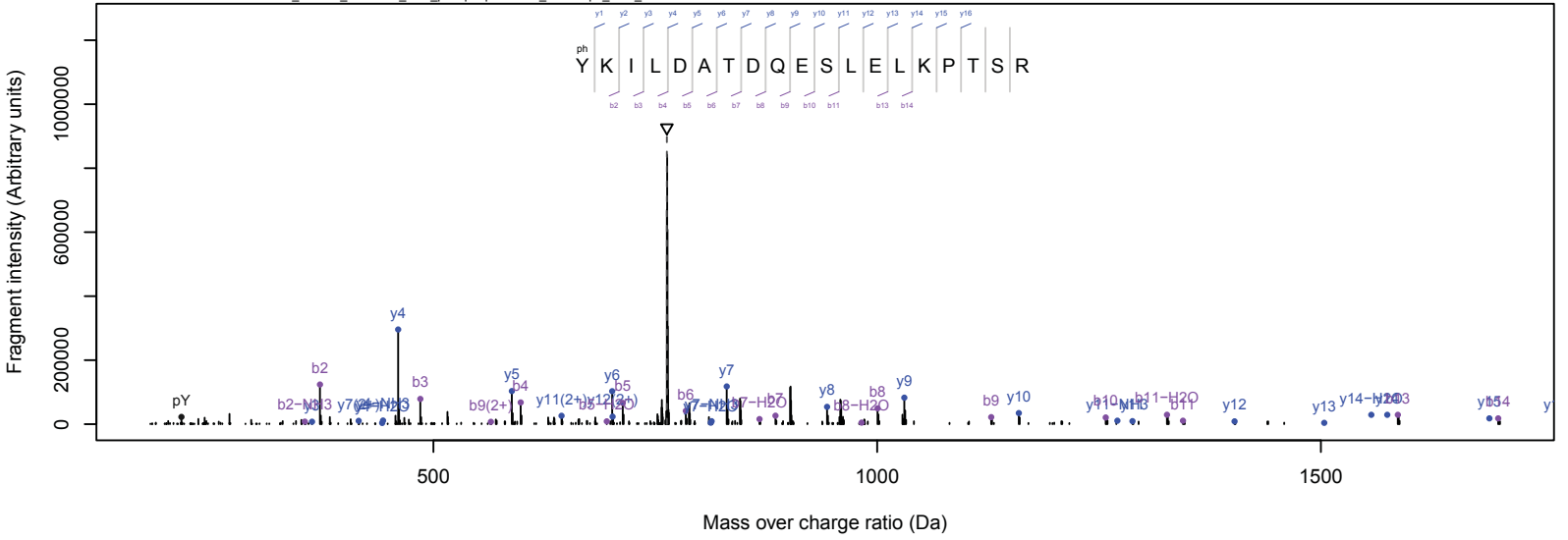
Andromeda score: 196.1



Y(ph)KILDATDQESLELKPTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25560

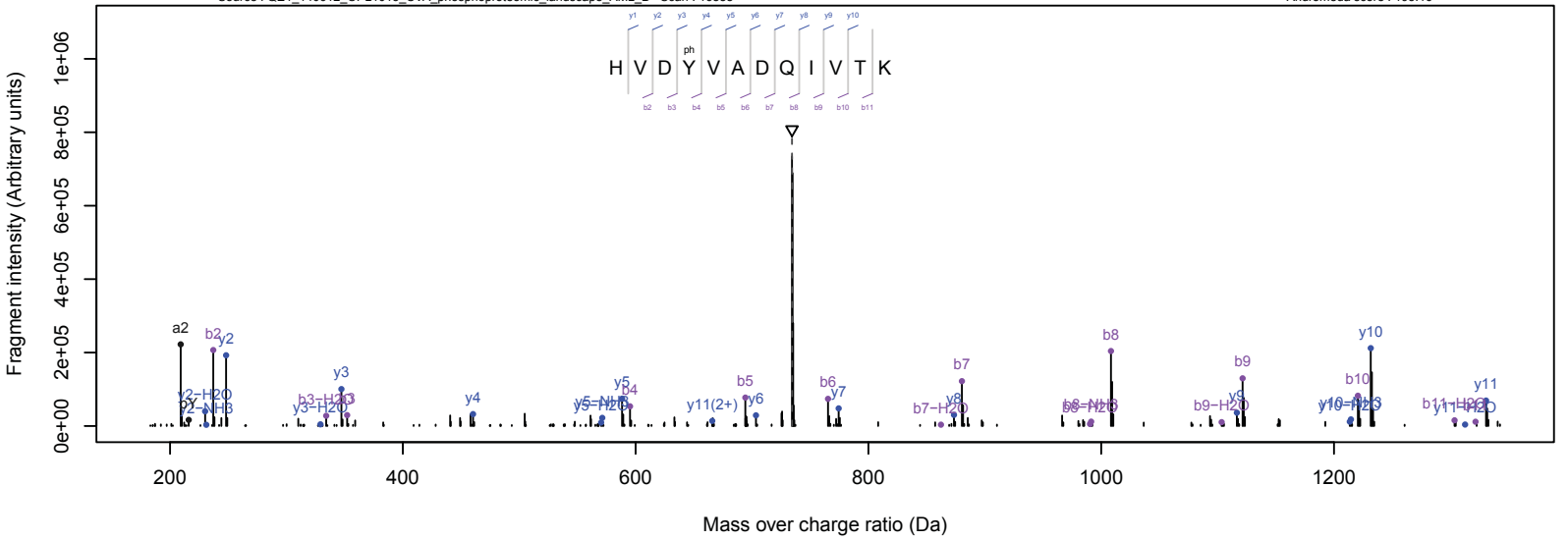
Andromeda score : 195.56



HVDY(ph)VADQIVTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16888

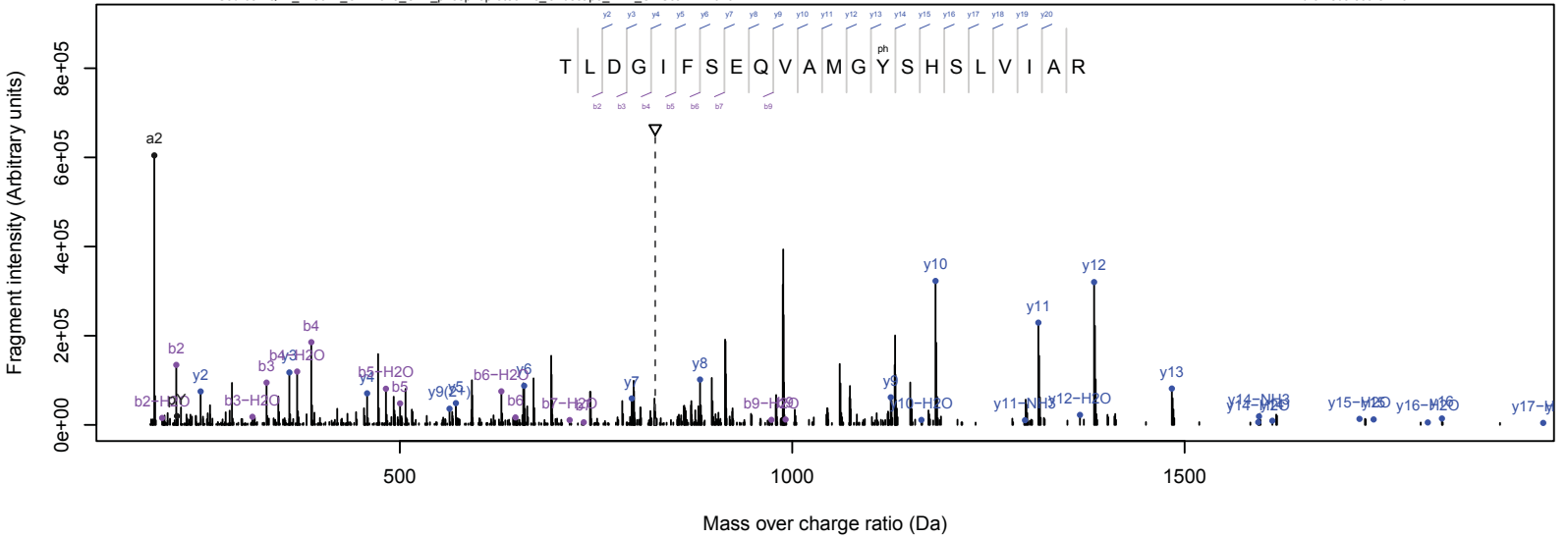
Andromeda score : 195.16



TLDGIFSEQVAMGY(ph)SHSLVIAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44610

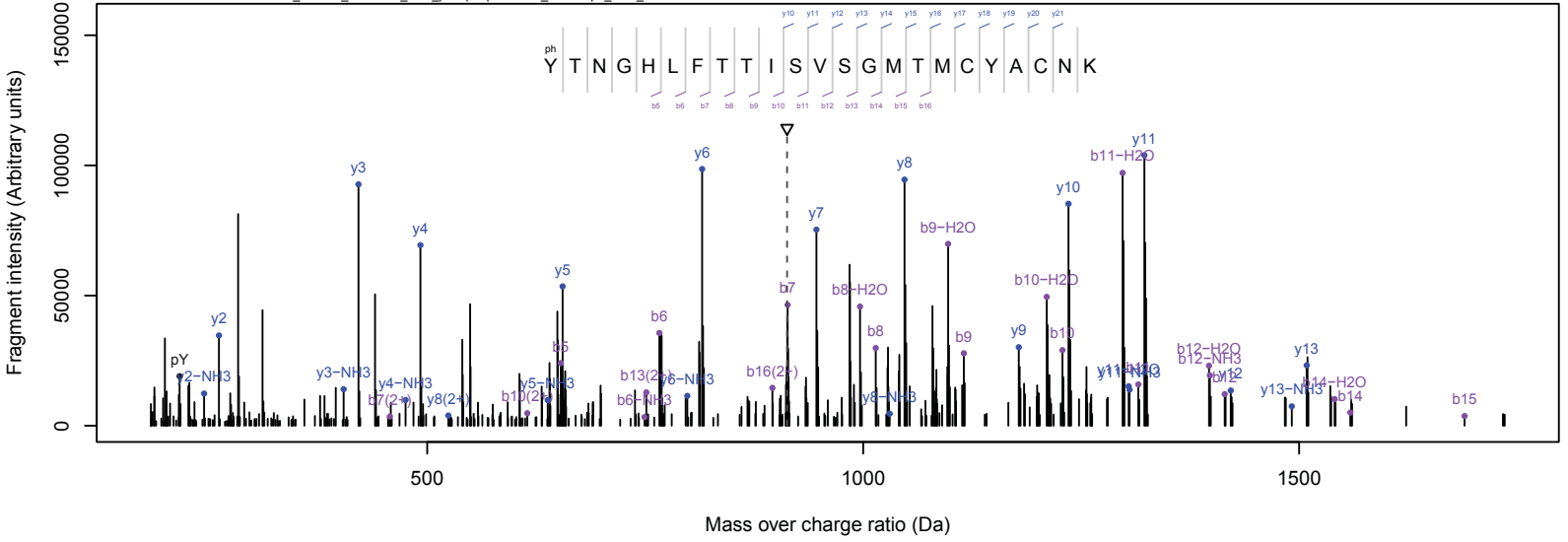
Andromeda score : 194.24



**Y(ph)TNGHLFTTISVSGMTMICYACNK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41542

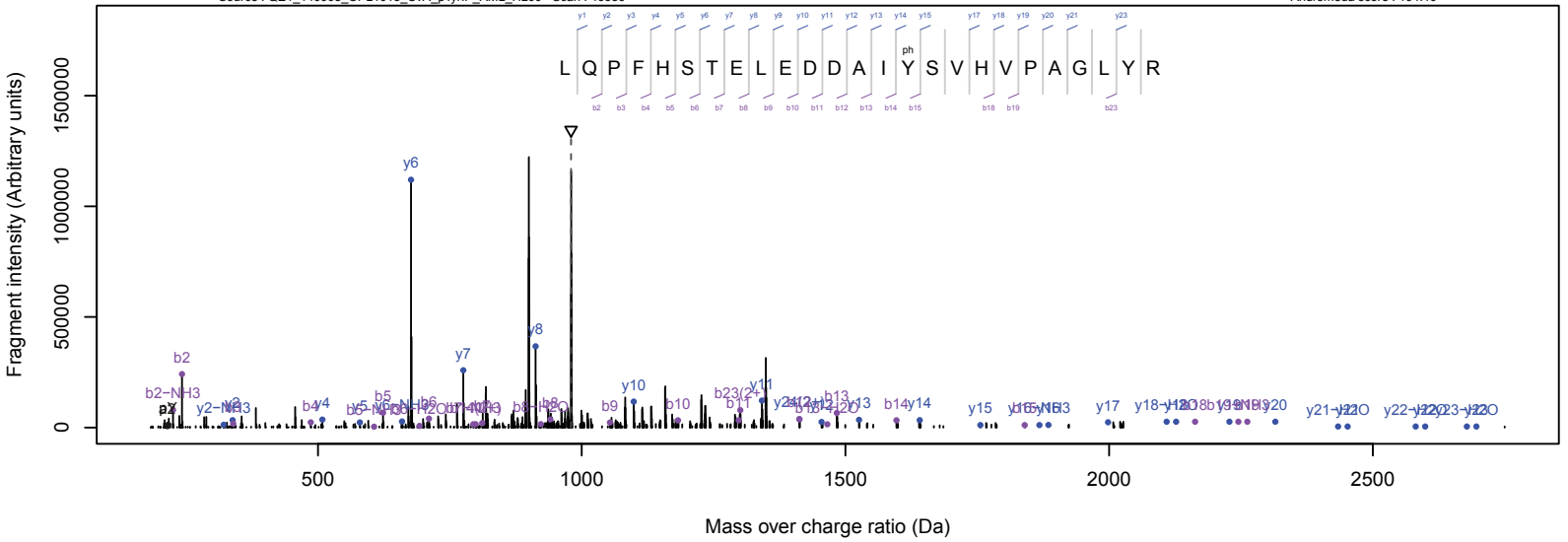
Andromeda score : 194.24



**LQPFHSTELEDDAIY(ph)SVHVPAAGLYR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 46586

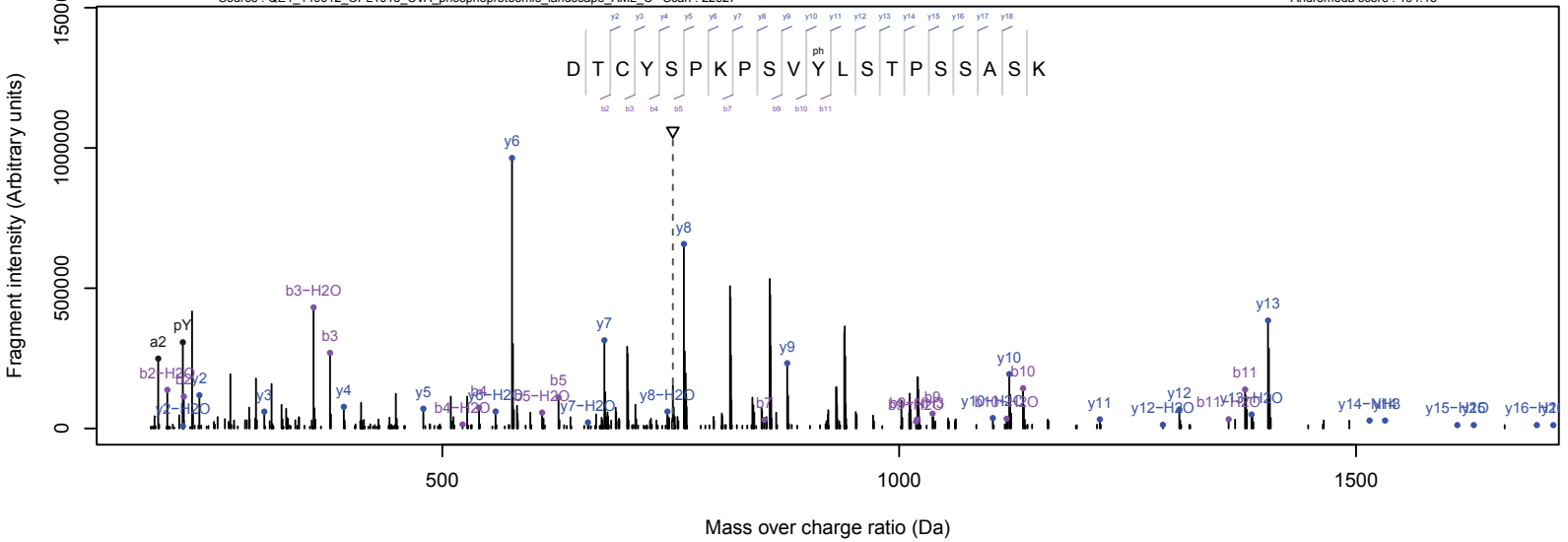
Andromeda score : 194.19



**DTCYSPKSPV(ph)LSTPSSASK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22627

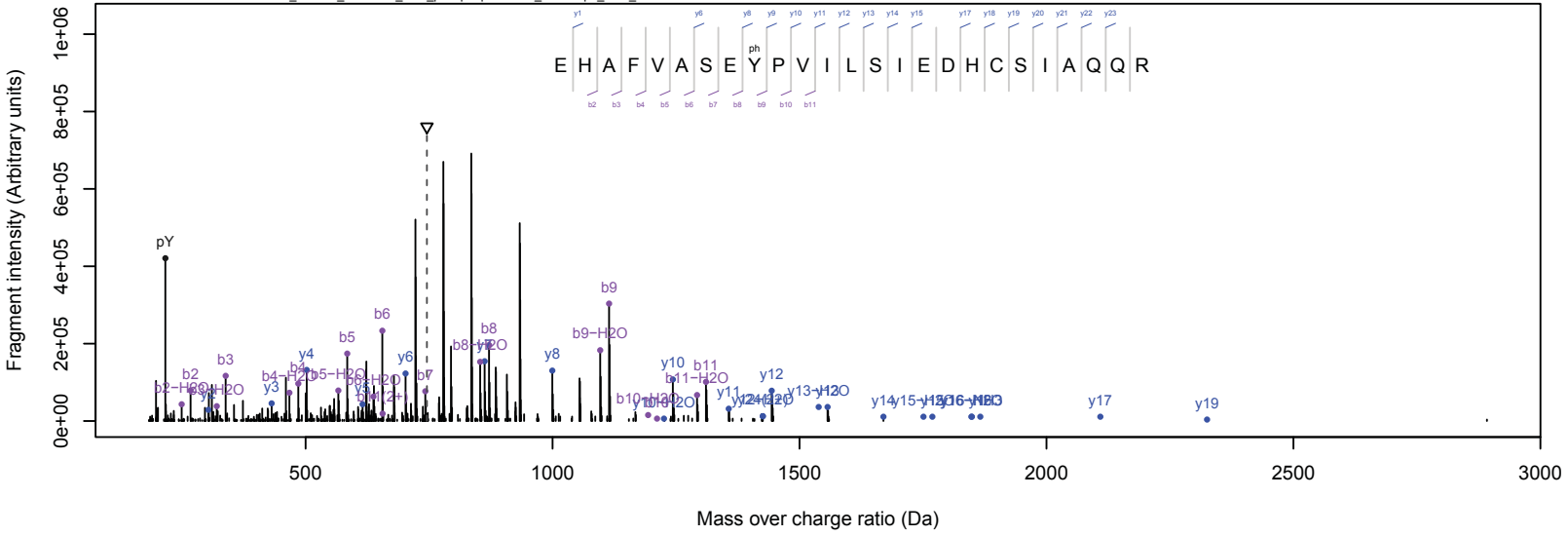
Andromeda score : 194.15



EHAHVASEY(ph)PVILSIEDHCSIAQQR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 40390

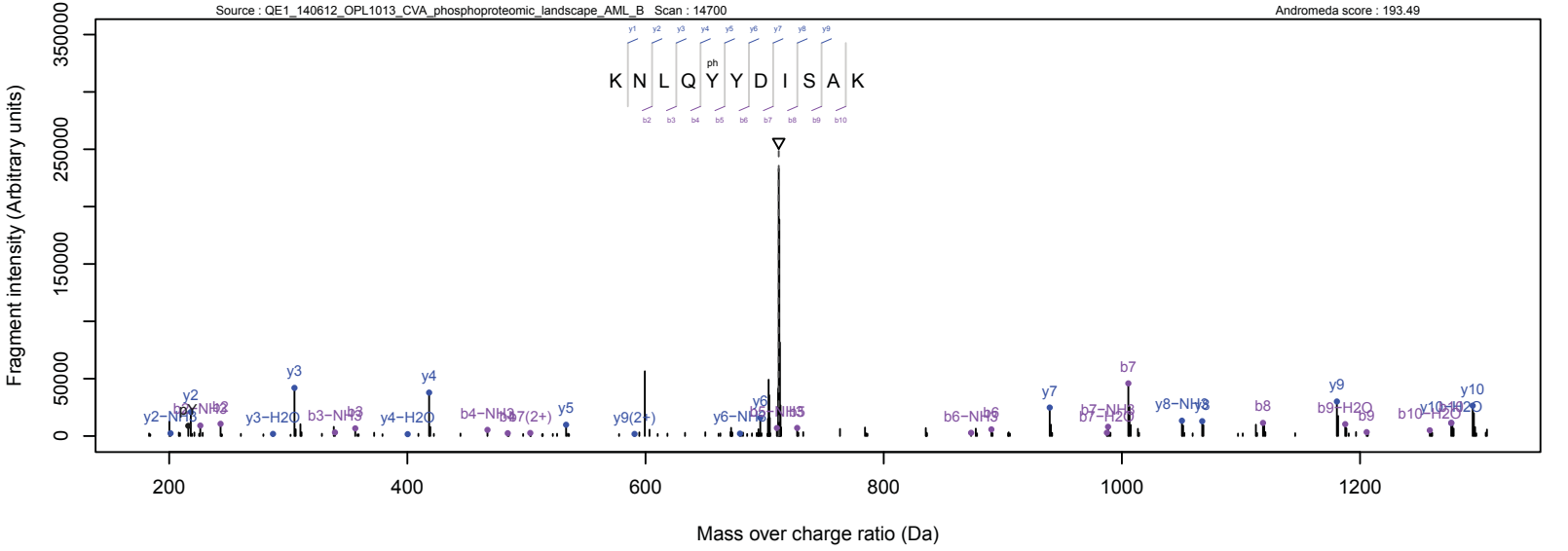
Andromeda score: 193.87



KNLQY(ph)YDISAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14700

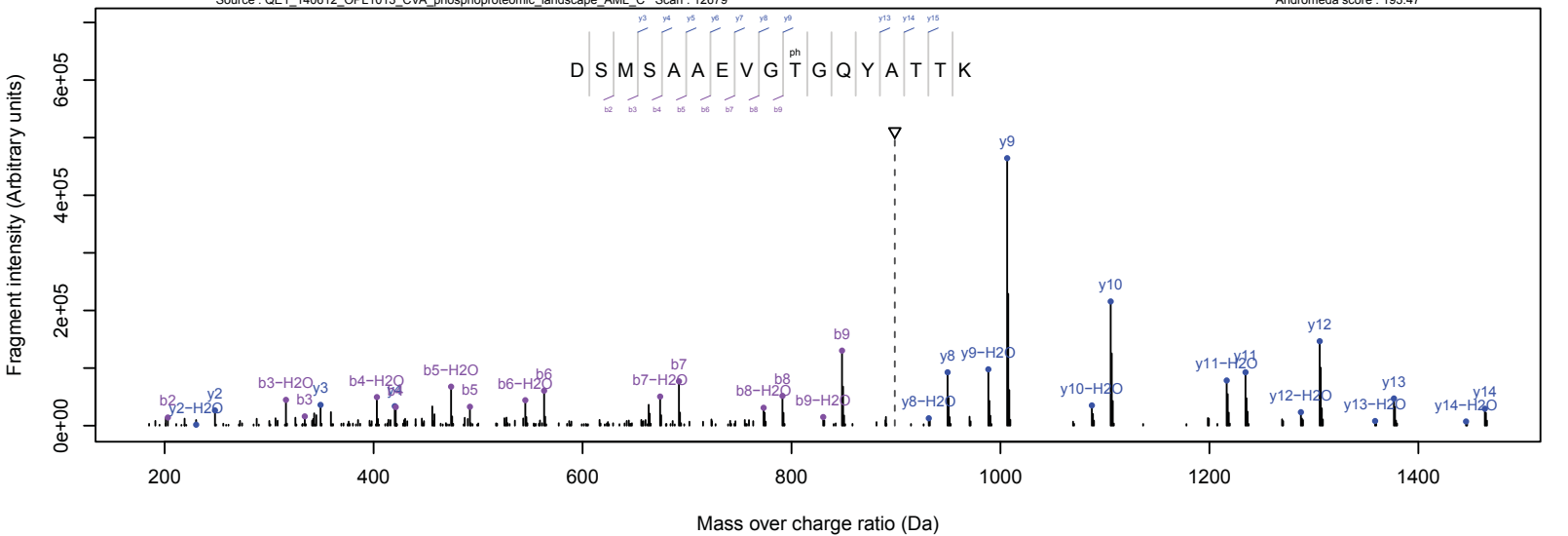
Andromeda score: 193.49



DSMSAAEVGT(ph)GYATTK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12679

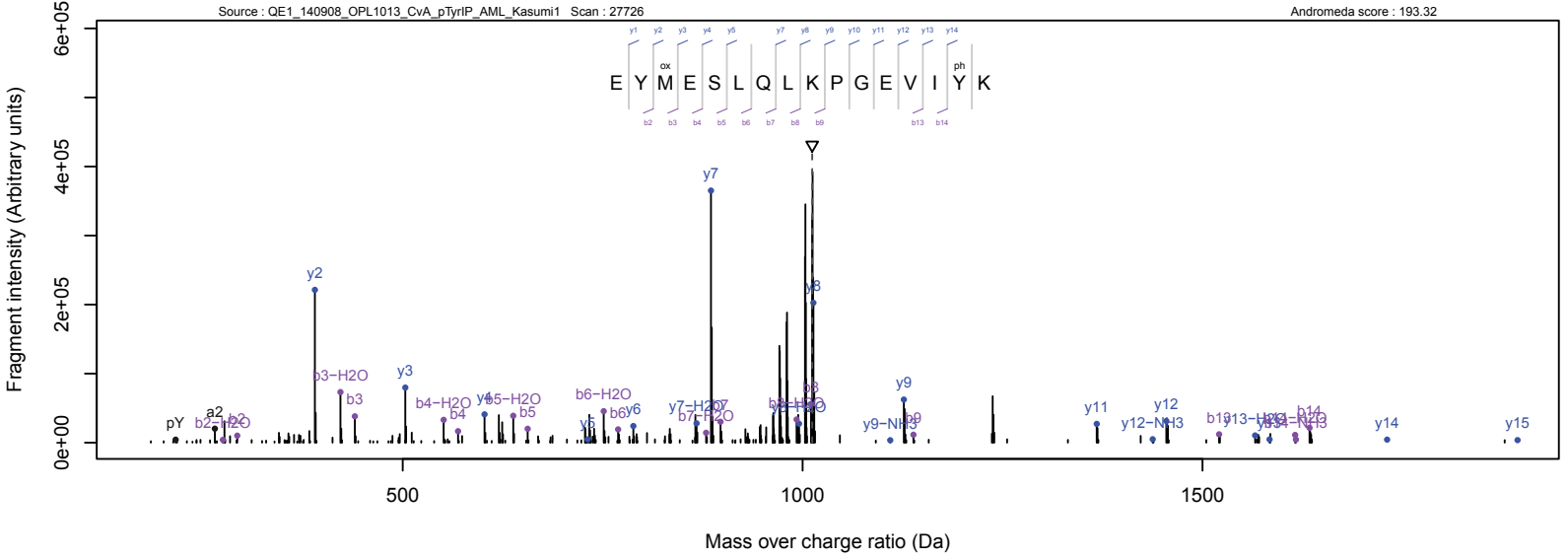
Andromeda score: 193.47



**EYM(ox)ESLQLKPGEVIYK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27726

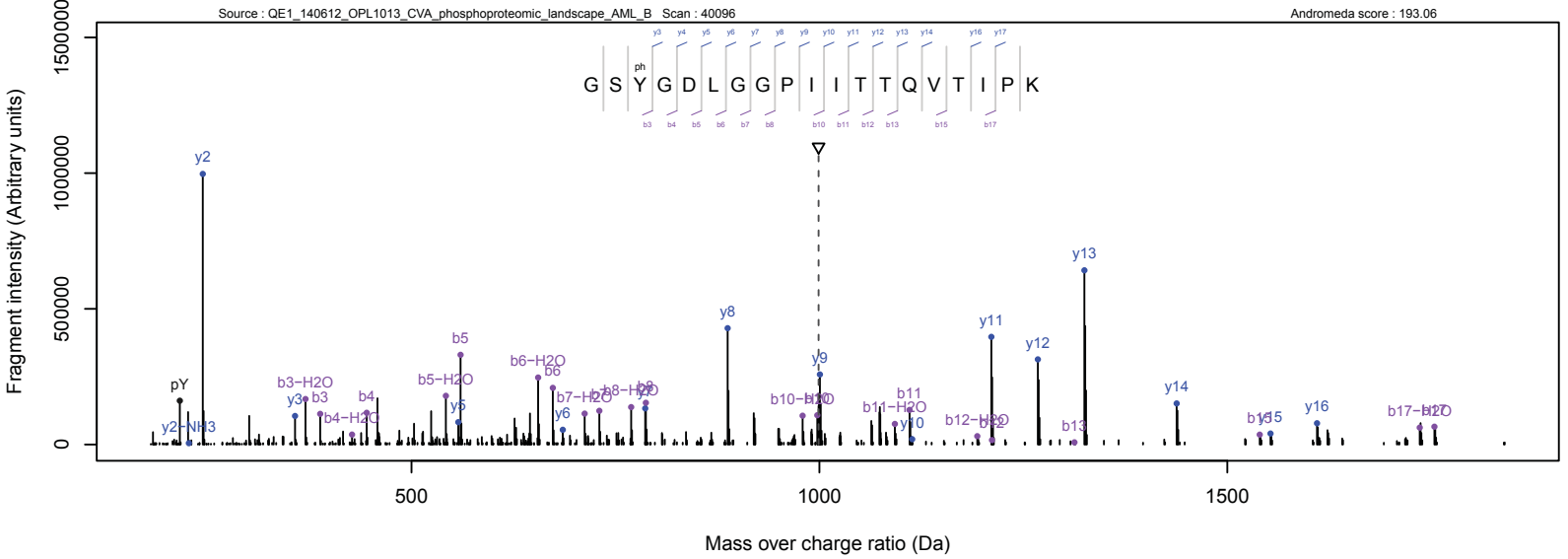
Andromeda score : 193.32



**GSY(ph)GDLGGPIITTQVTIPK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40096

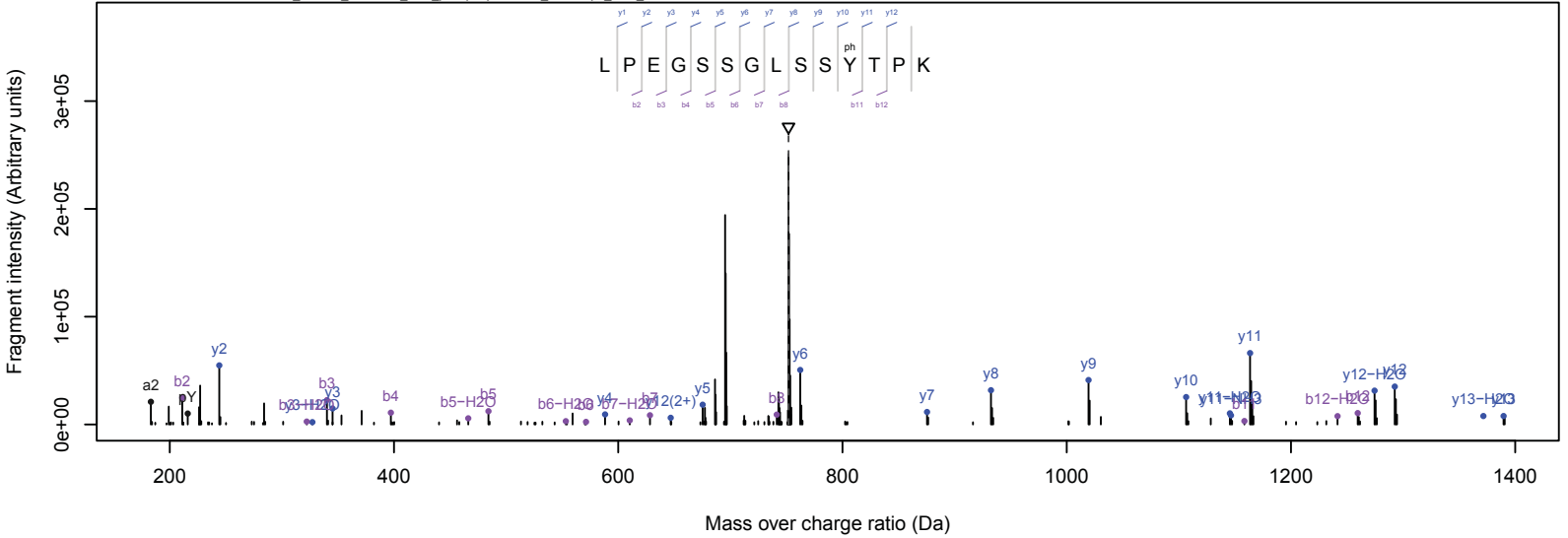
Andromeda score : 193.06



**LPEGSSGLSSYPK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16567

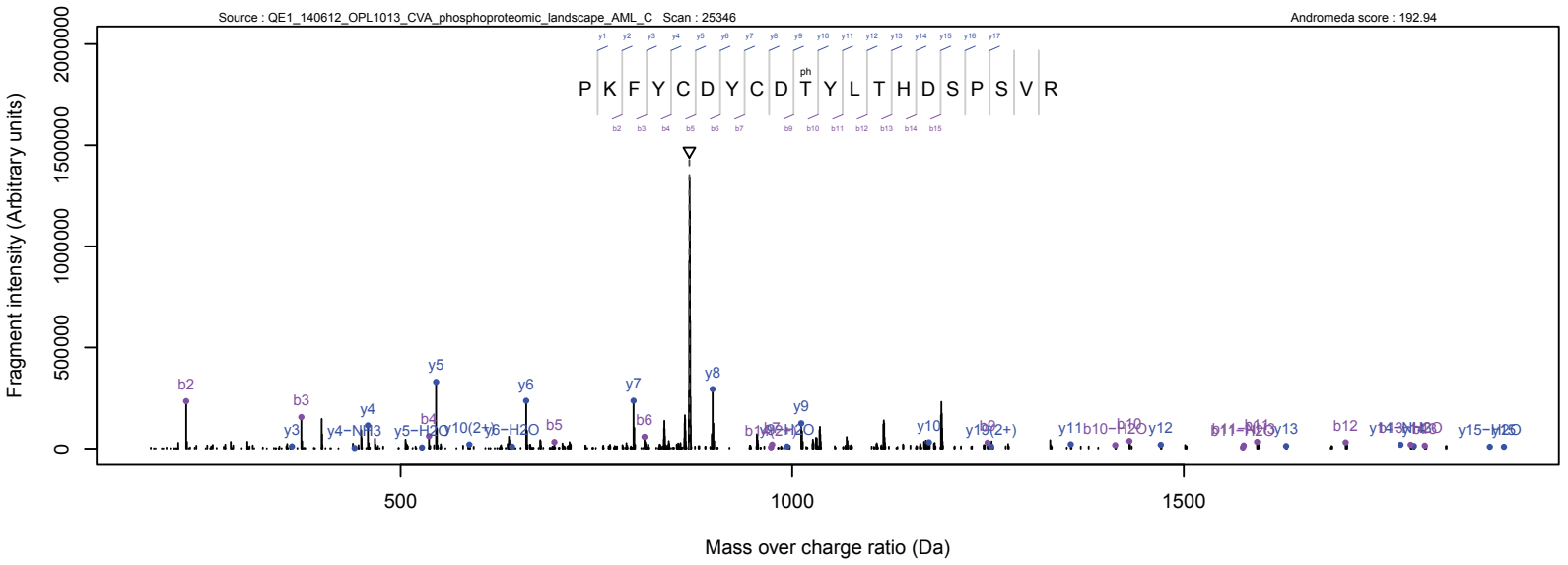
Andromeda score : 193.06



PKFYCDYCDT(ph)YLTHDPSVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25346

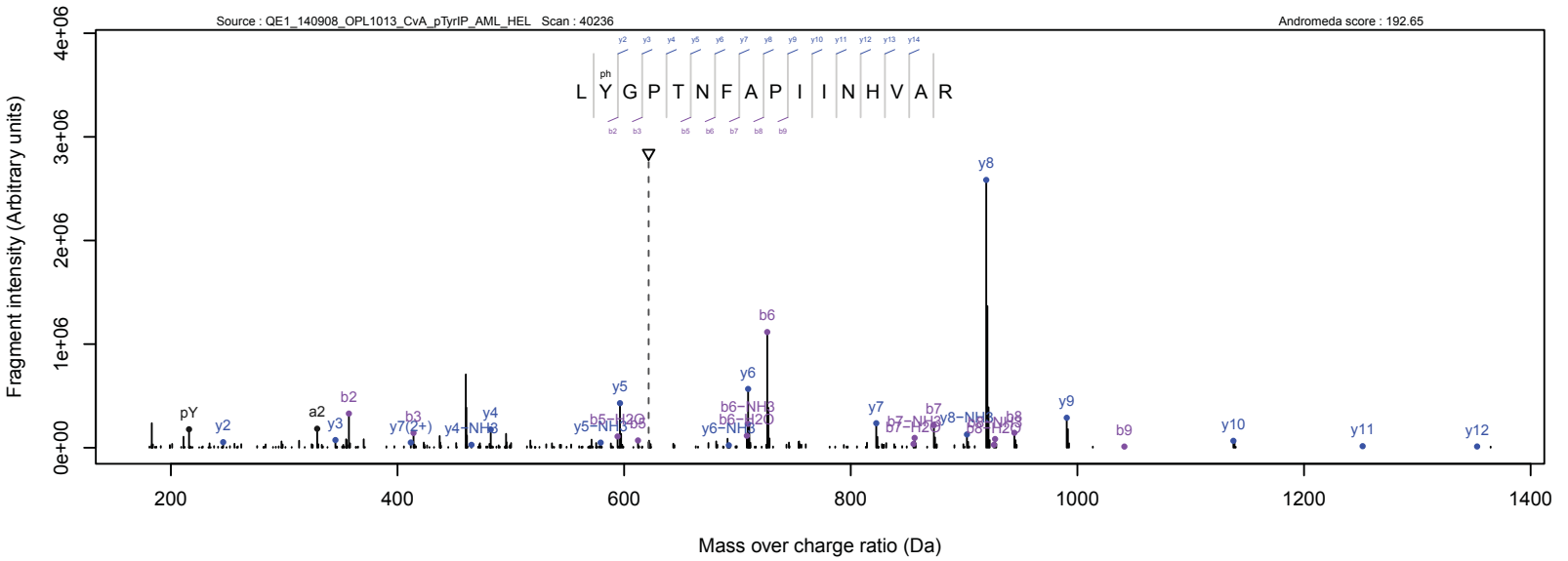
Andromeda score : 192.94



LY(ph)GPTNFAPINHVAR-3+

Source : QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_HEL Scan : 40236

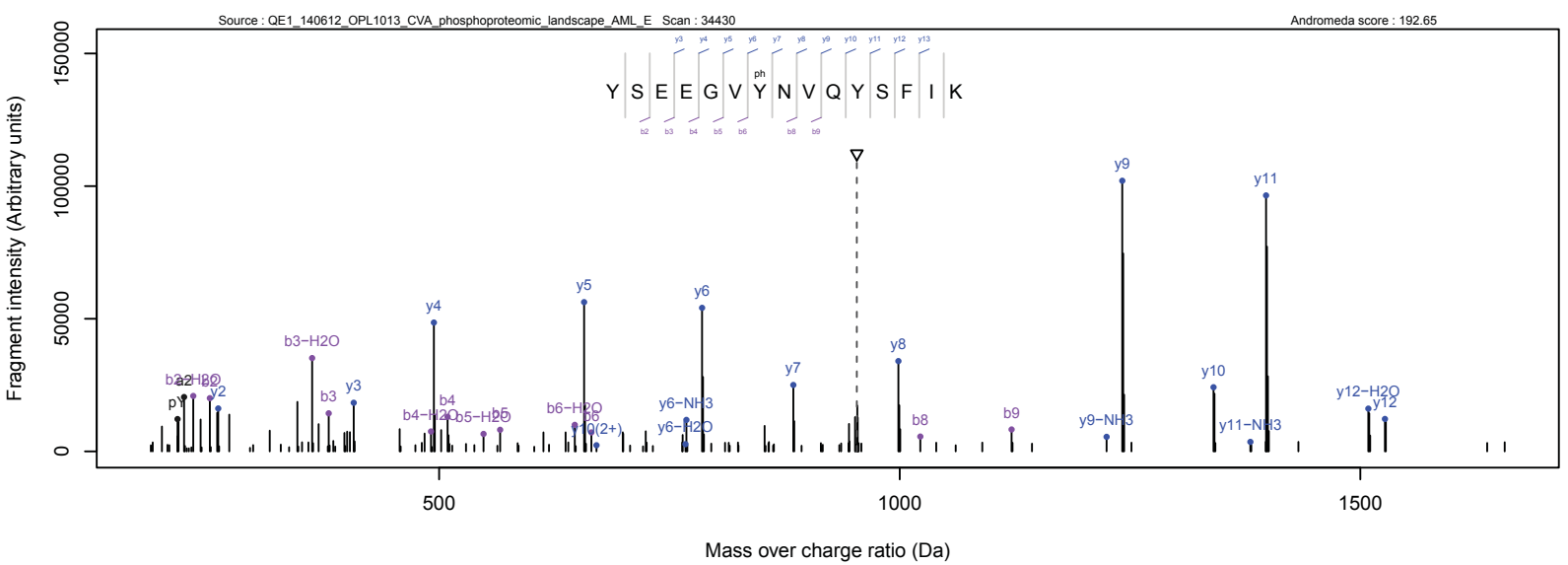
Andromeda score : 192.65



YSEEGVY(ph)NVQYSFIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 34430

Andromeda score : 192.65

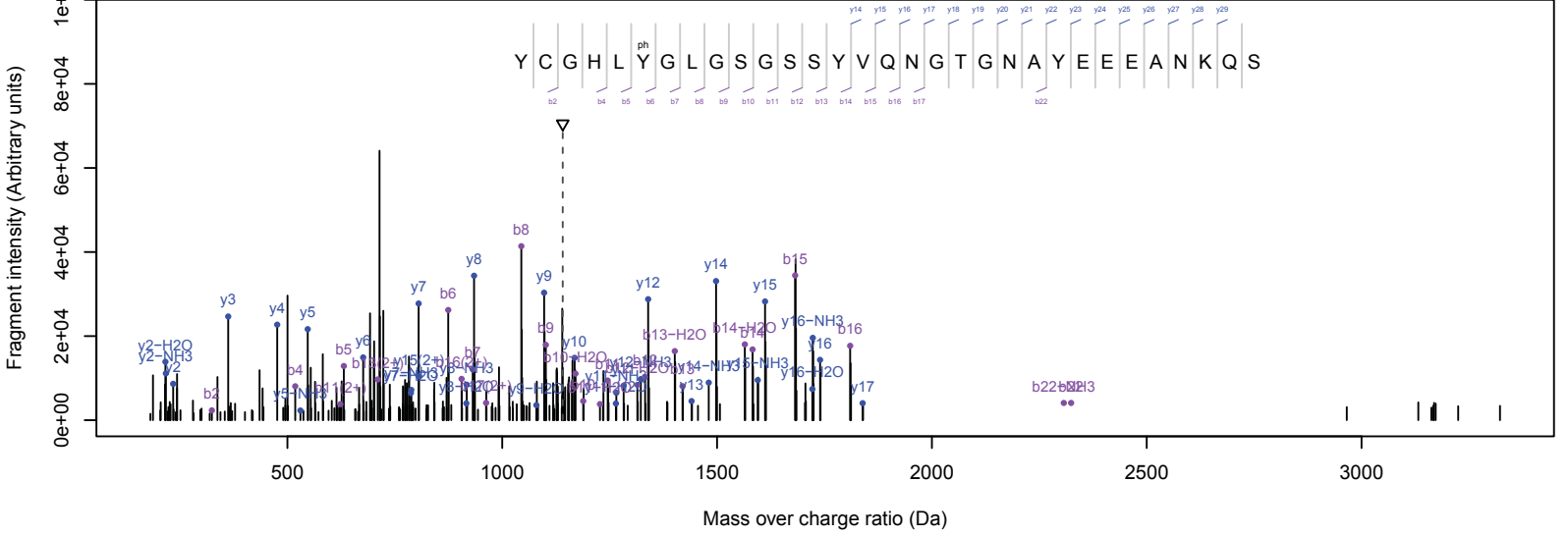




YCGHLY(ph)GLGSGSSVQNGTGNAYEEEANKQS-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 29420

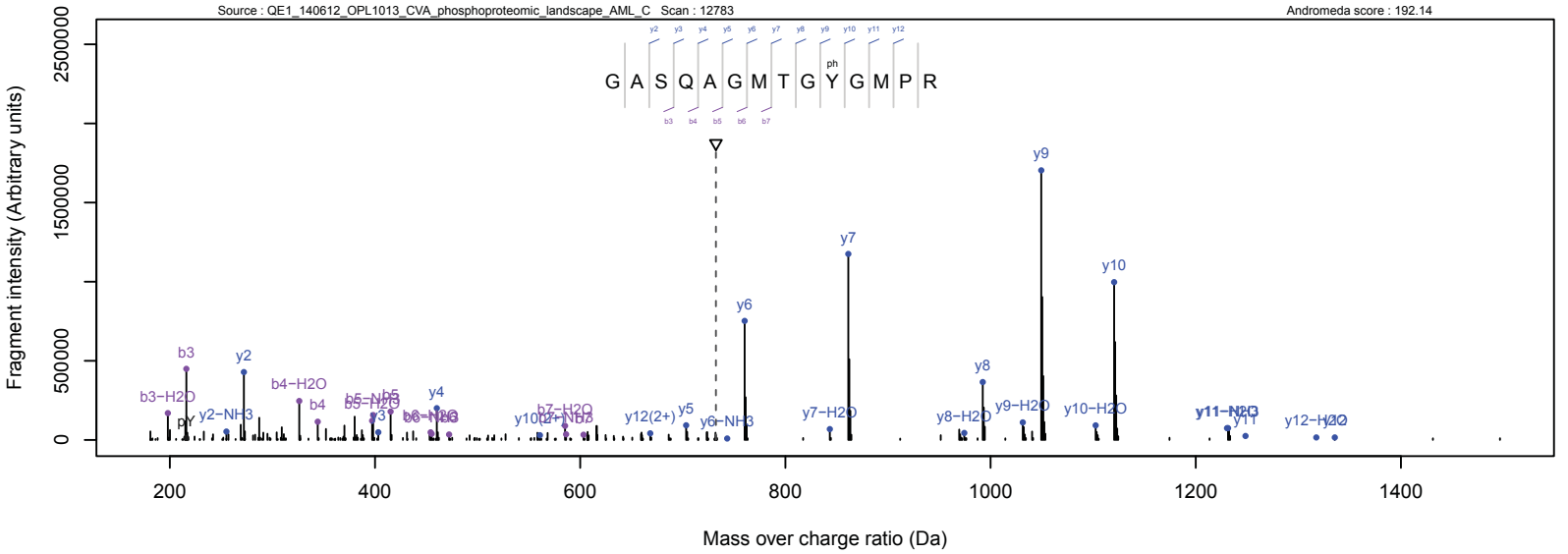
Andromeda score: 192.24



GASQAGMTG(ph)GMPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12783

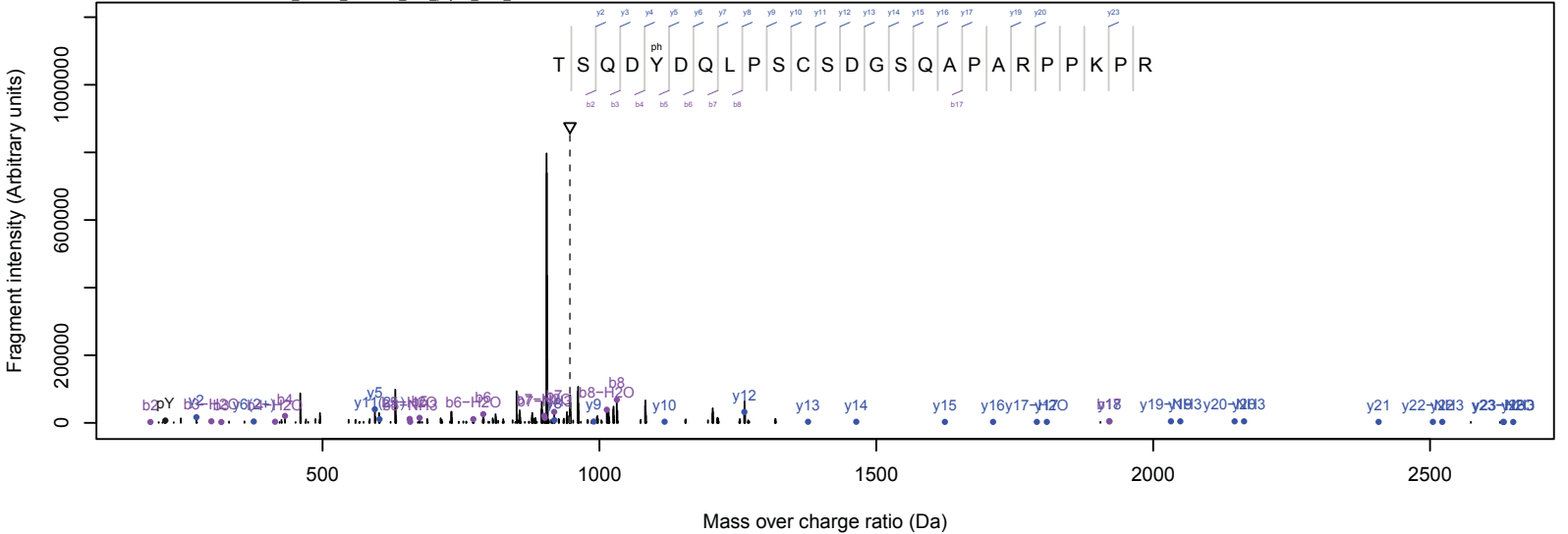
Andromeda score: 192.14



TSQDY(ph)DQLPSCDGSQAPARPPKPR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 15502

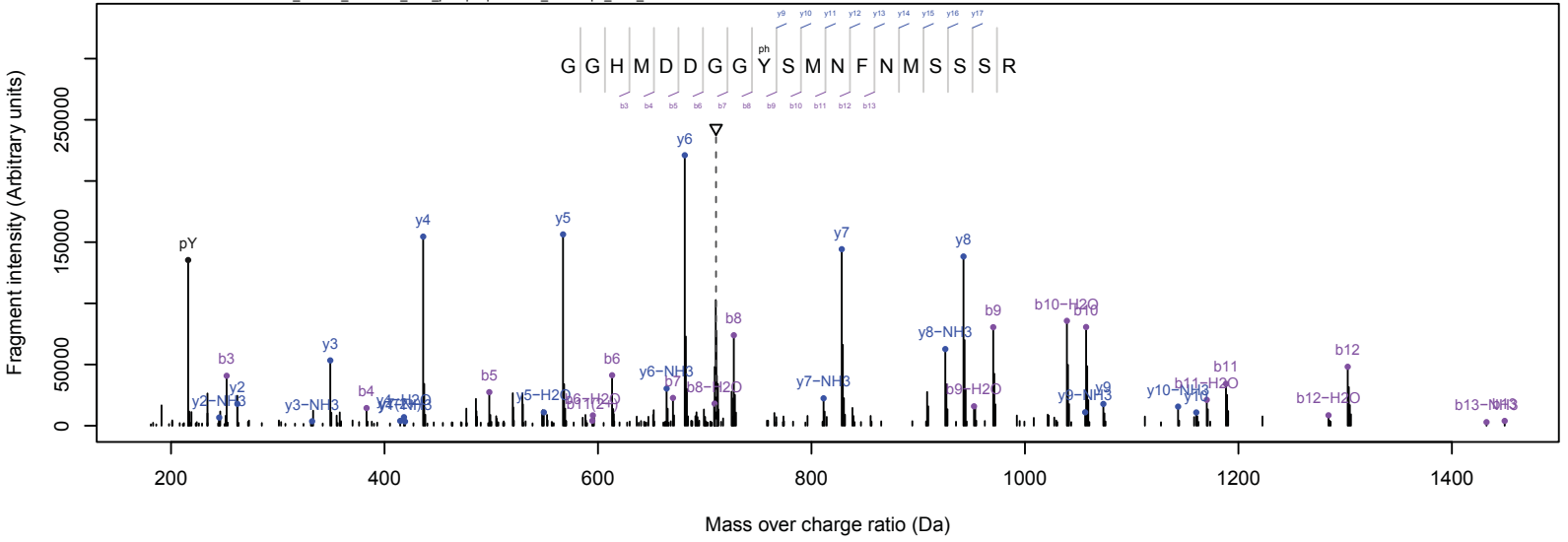
Andromeda score: 192.1



GGHMDGGY(ph)SMNFMSSSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21441

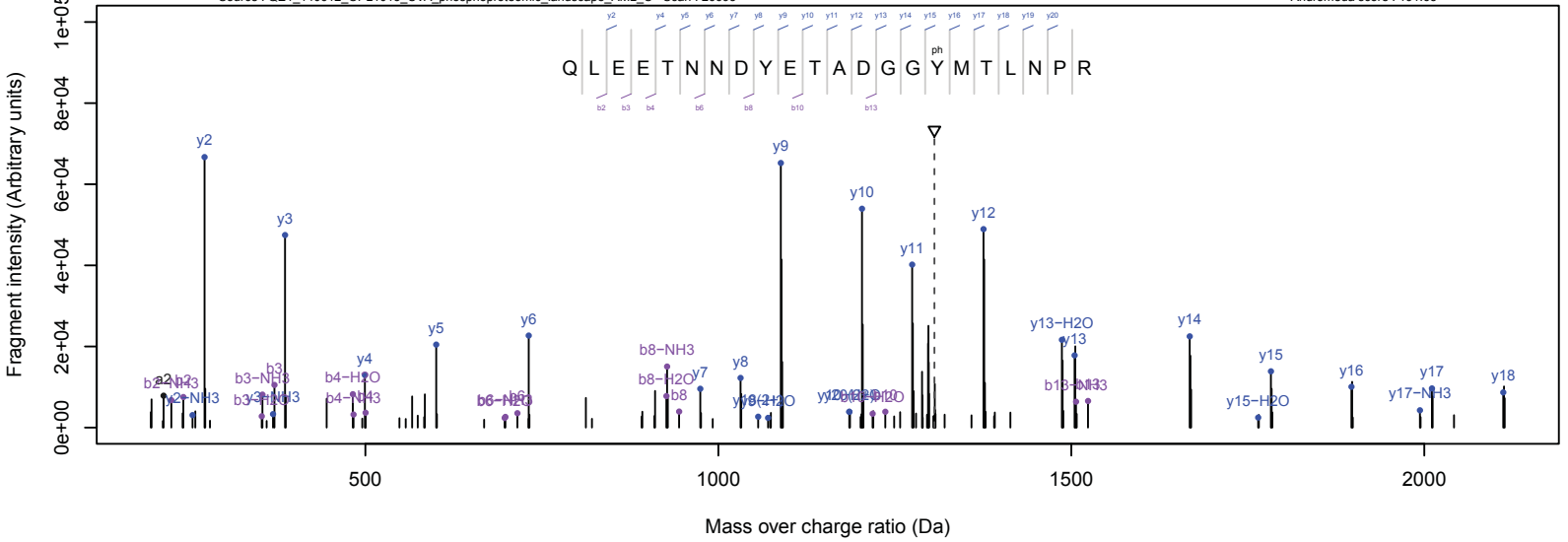
Andromeda score : 191.93



QLEETNNDYETADGGY(ph)MTLNPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26693

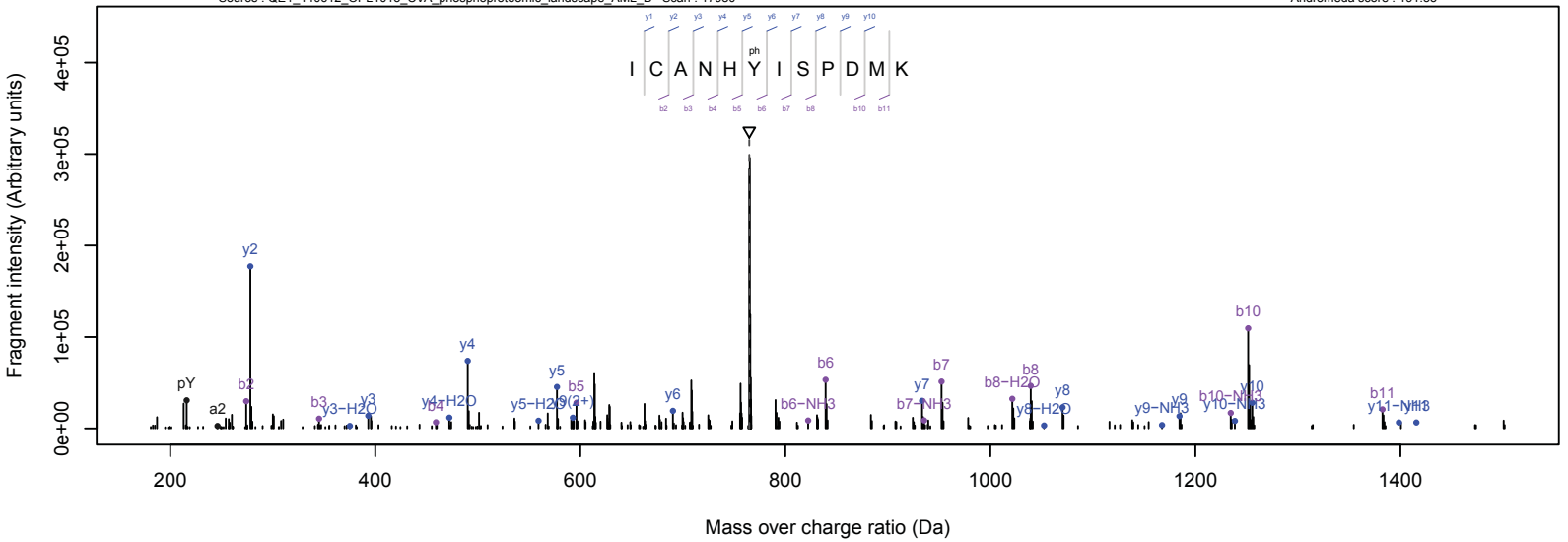
Andromeda score : 191.65



ICANHYP(h)SPDMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17980

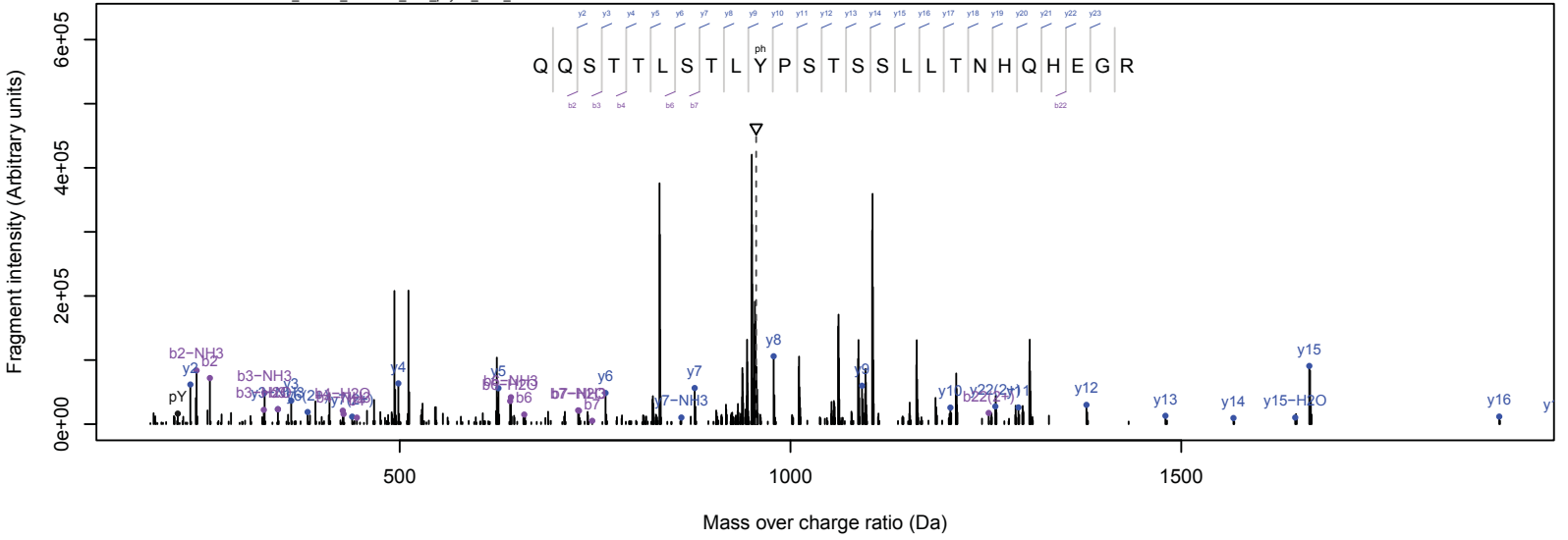
Andromeda score : 191.53



QQSTTLSTLY(ph)PSTSSLLTNHQHEGR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27745

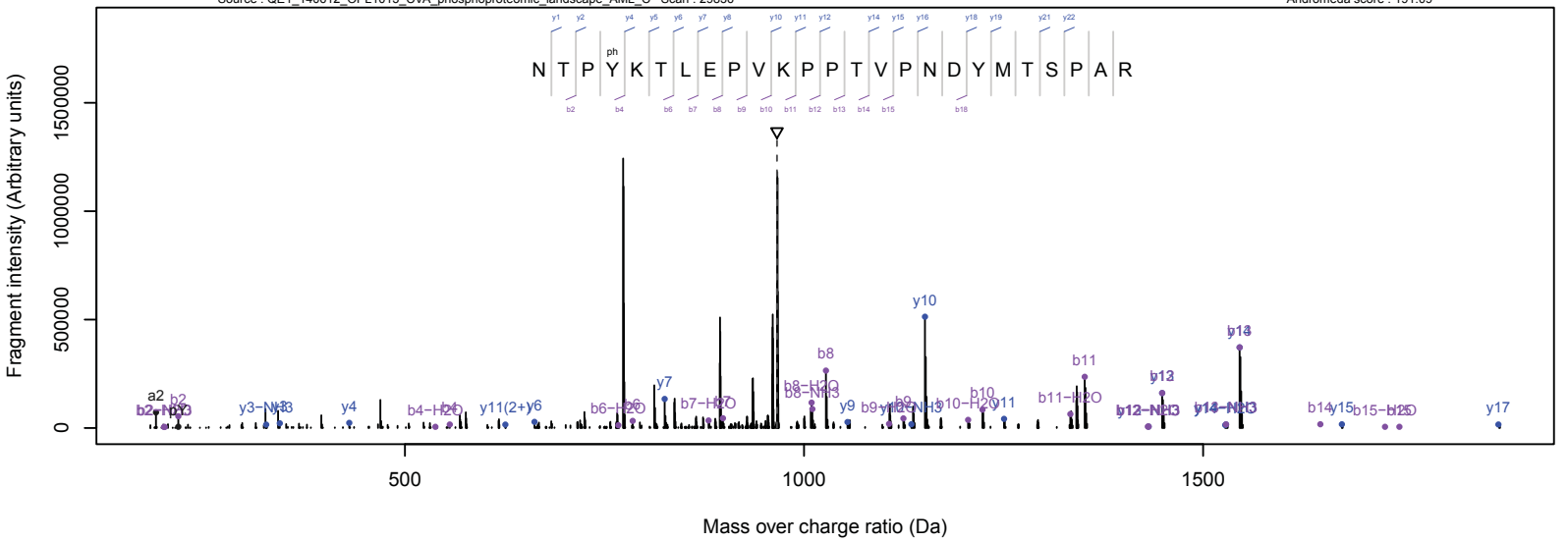
Andromeda score : 191.2



NTPY(ph)KTLEPVKPPPTVPNDYMTSPAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29836

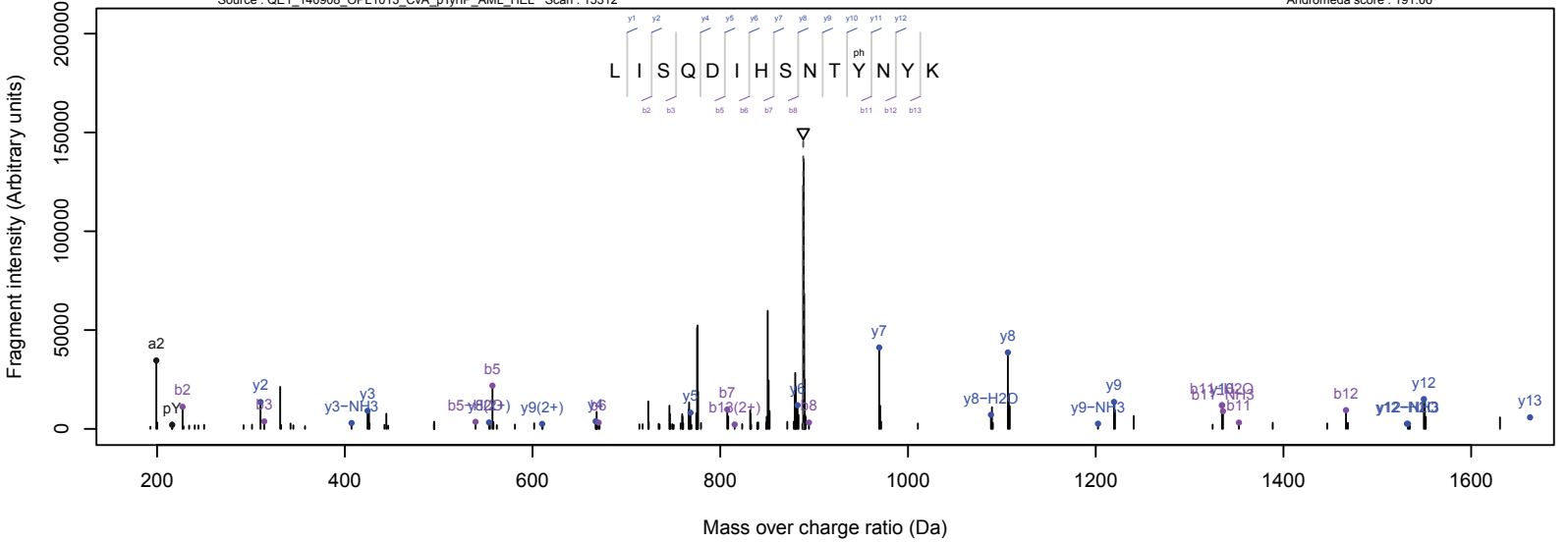
Andromeda score : 191.09



LISQDIHSNTYNYK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 15312

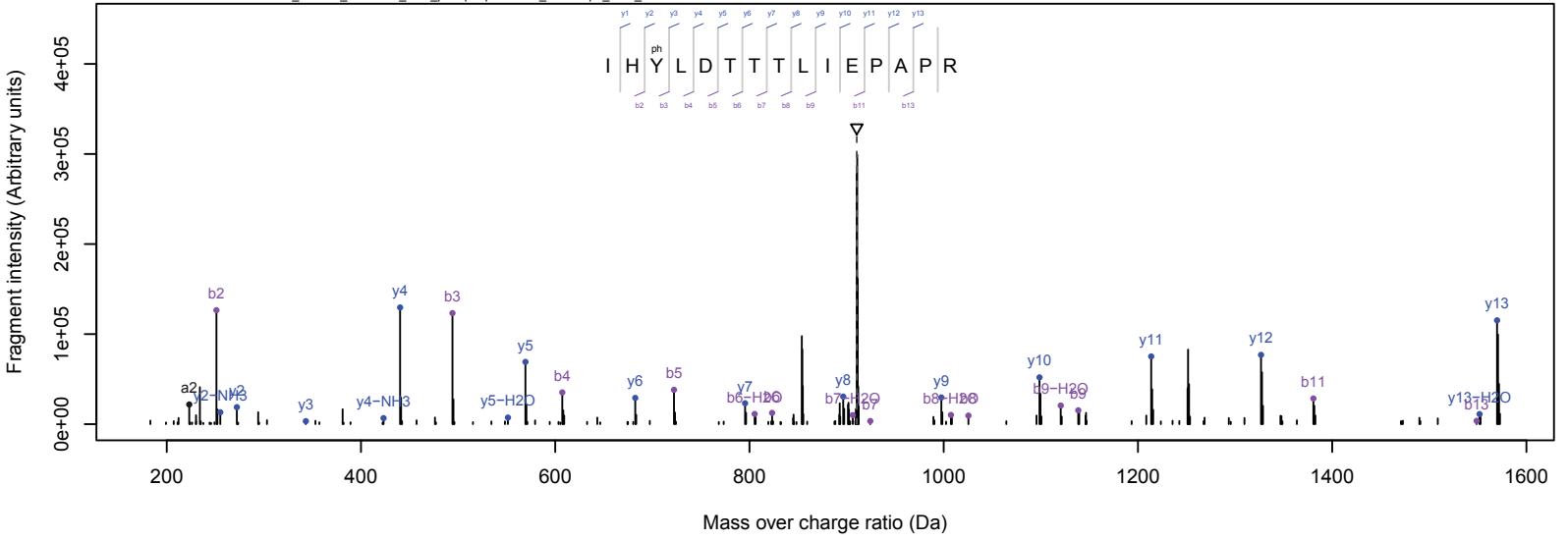
Andromeda score : 191.06



IHY(ph)LDTTTLIEPAPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 32285

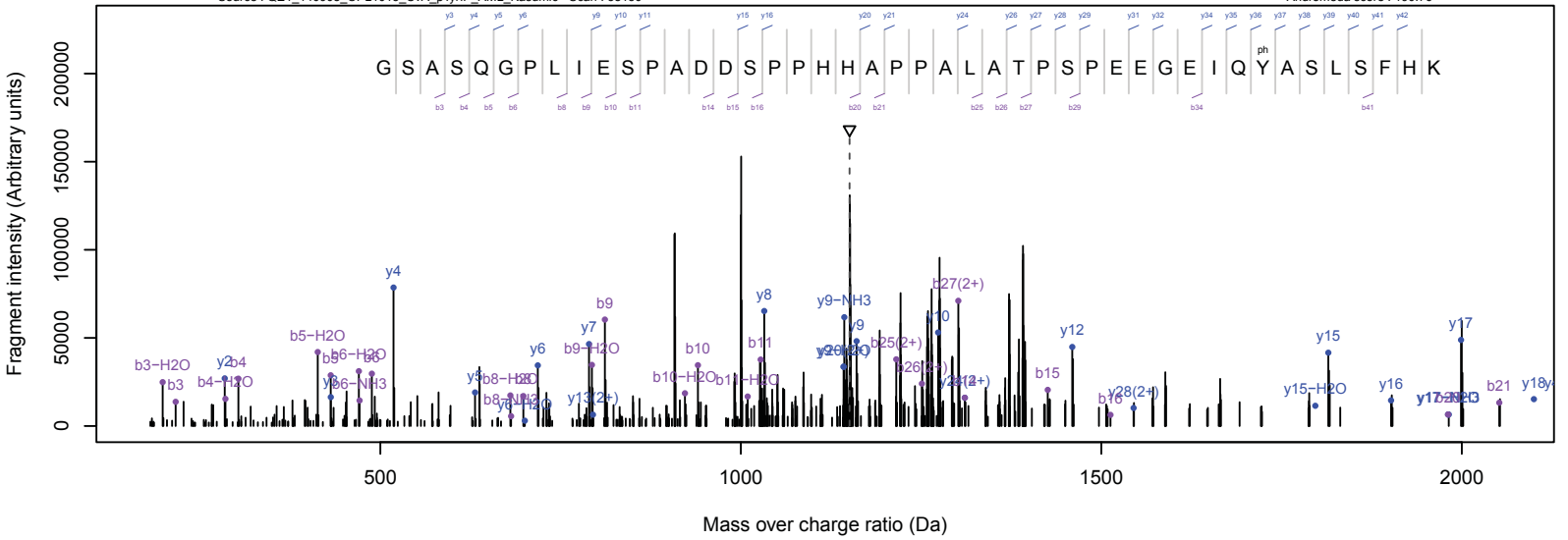
Andromeda score : 190.86



GSASQGLIESPADDSPPHAPPALATPSPEEGEIYQ(ph)ASLSFHK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 38150

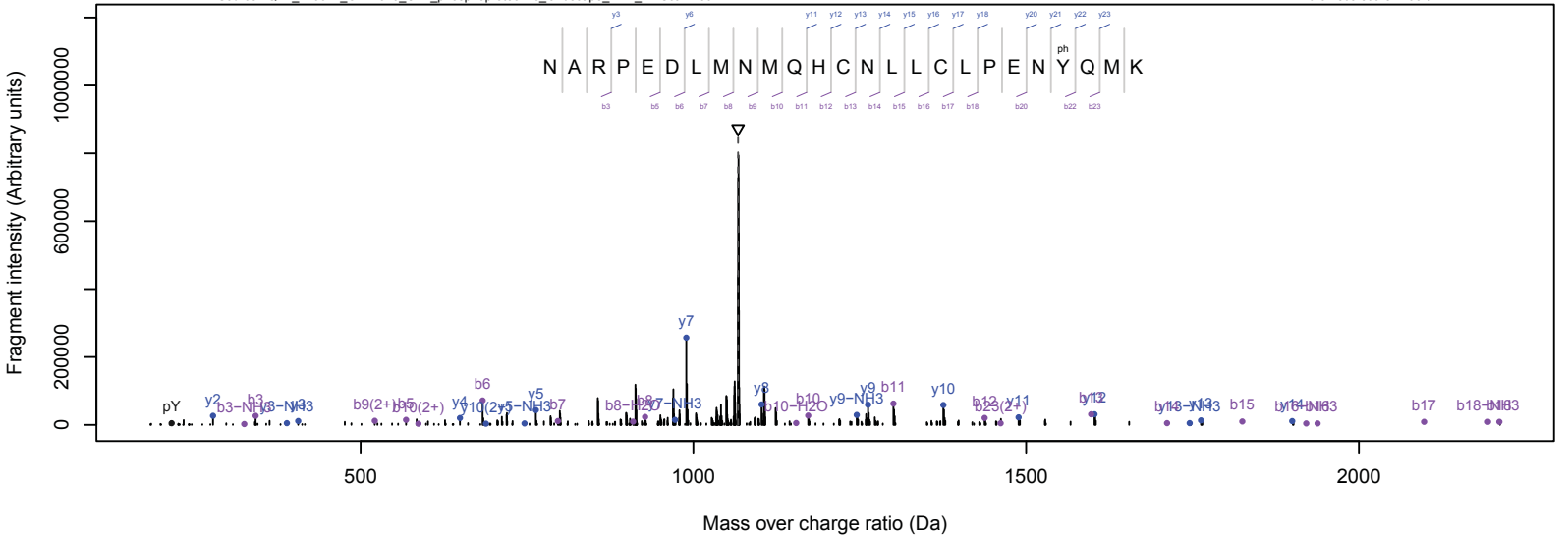
Andromeda score : 190.76



NARPELDMNQHCNLLCLPENYQMK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 39427

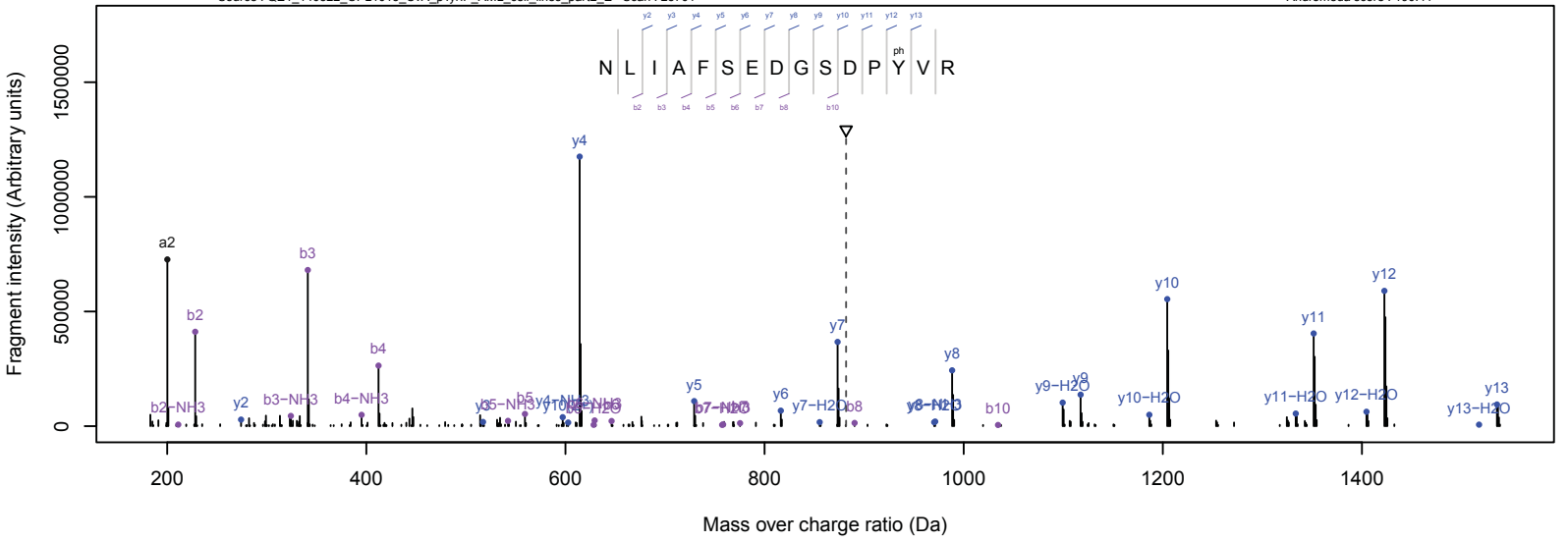
Andromeda score : 190.5



NLIAFSEGDSPY(ph)VR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 25701

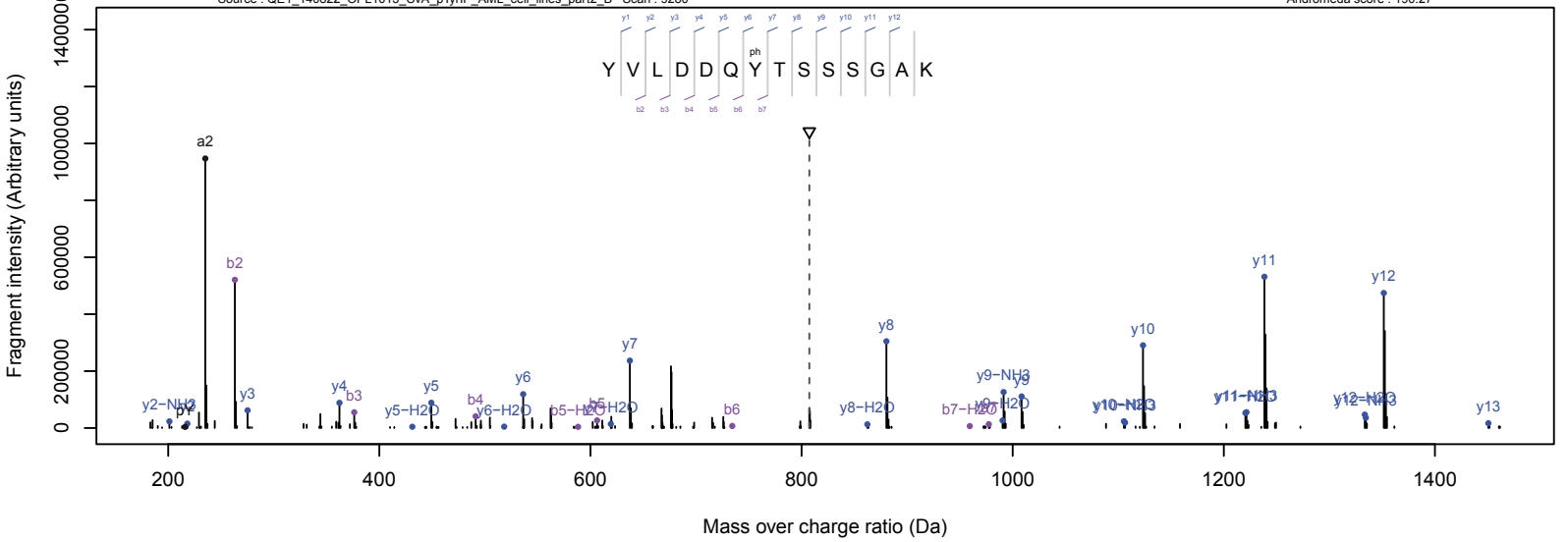
Andromeda score: 190.41



YVLDDQY(ph)TSSSGAK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 9280

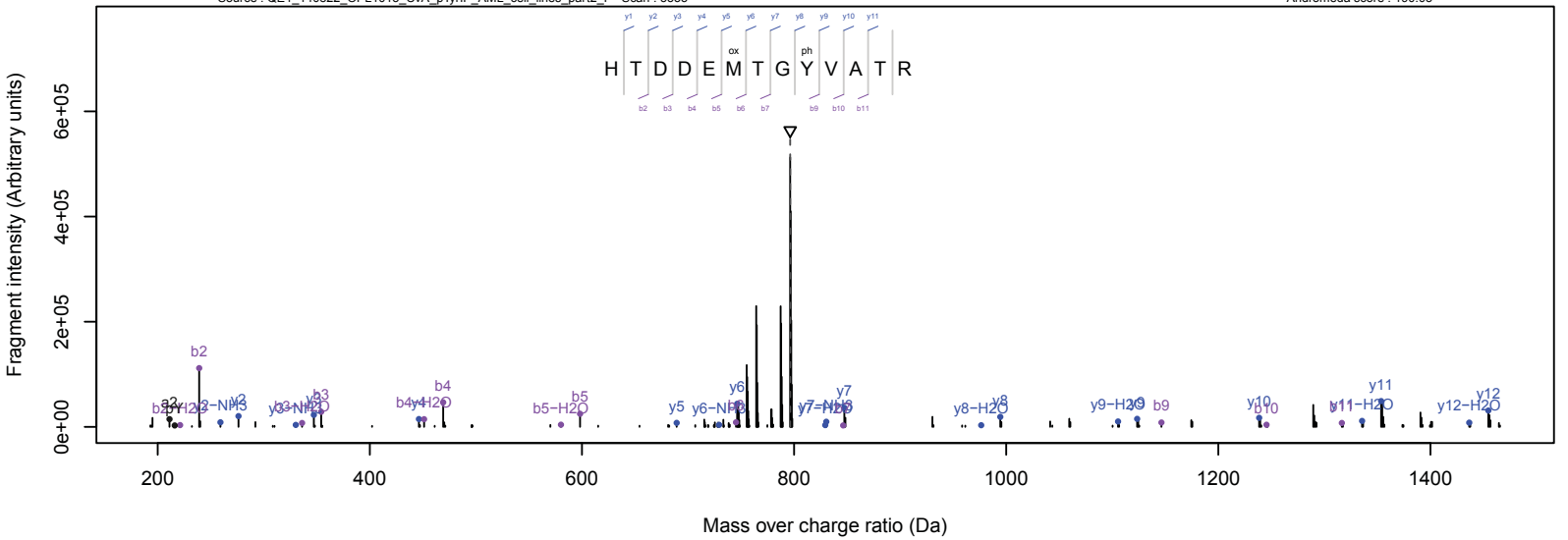
Andromeda score: 190.27



HTDDEM(ox)TGY(ph)VATR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan: 5533

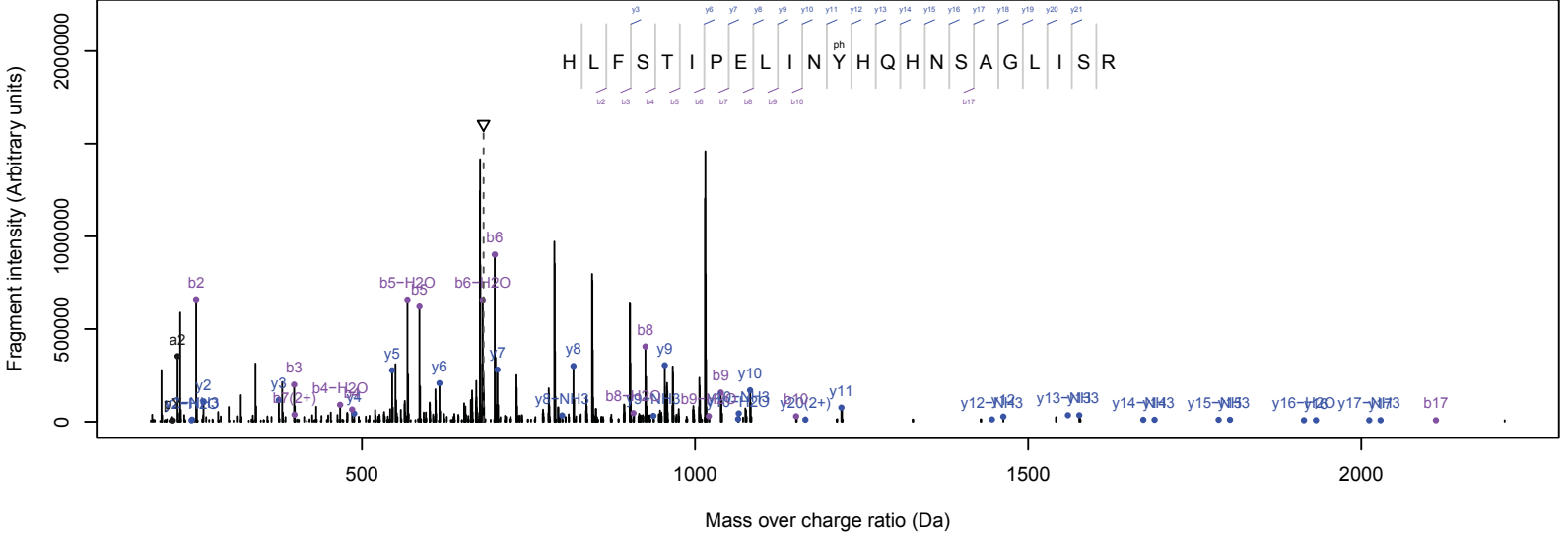
Andromeda score: 190.03



**HLFSTIPELINY(ph)HQHNSAGLISR-4+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 39122

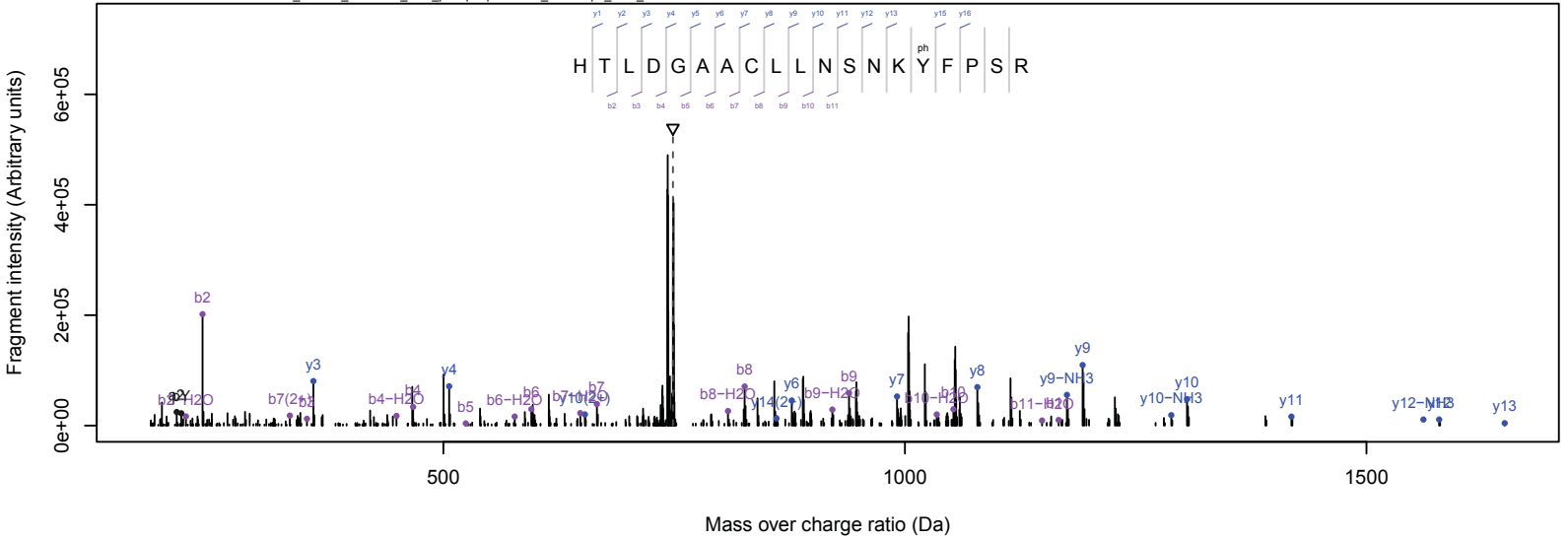
Andromeda score : 189.82



**HTLDGAACLLNSNKY(ph)FPSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30321

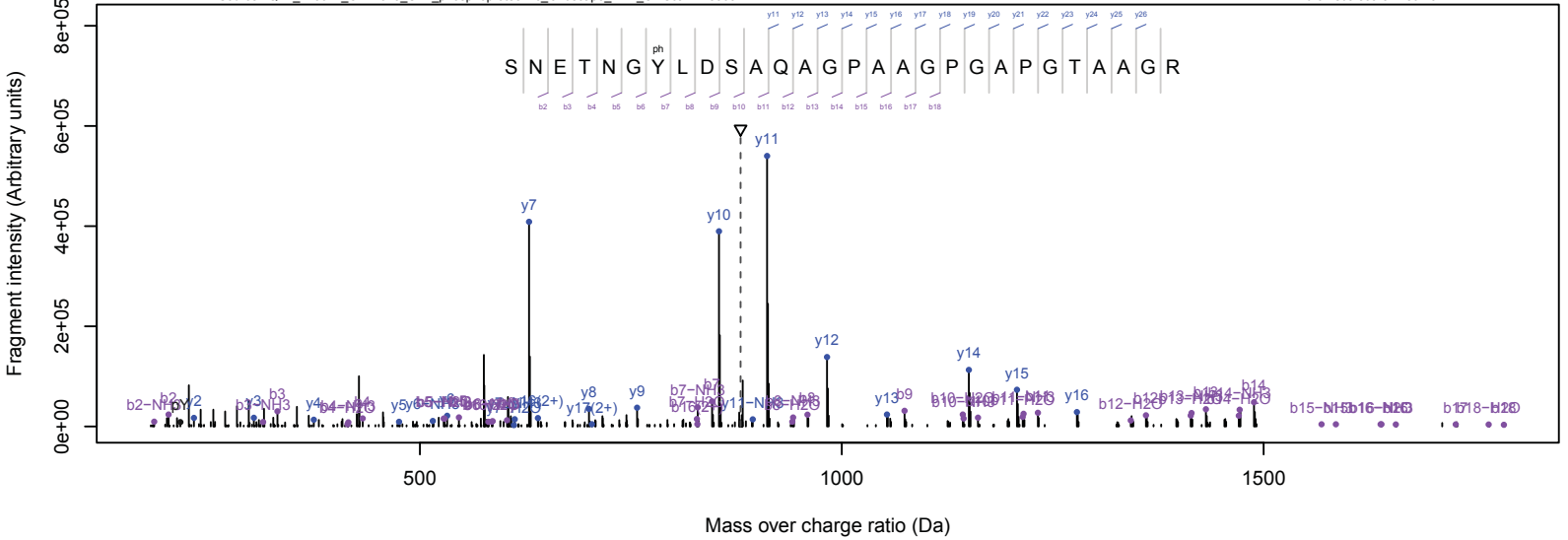
Andromeda score : 189.77



**SNETNGY(ph)LDSAQAGPAAGPGTAAGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18365

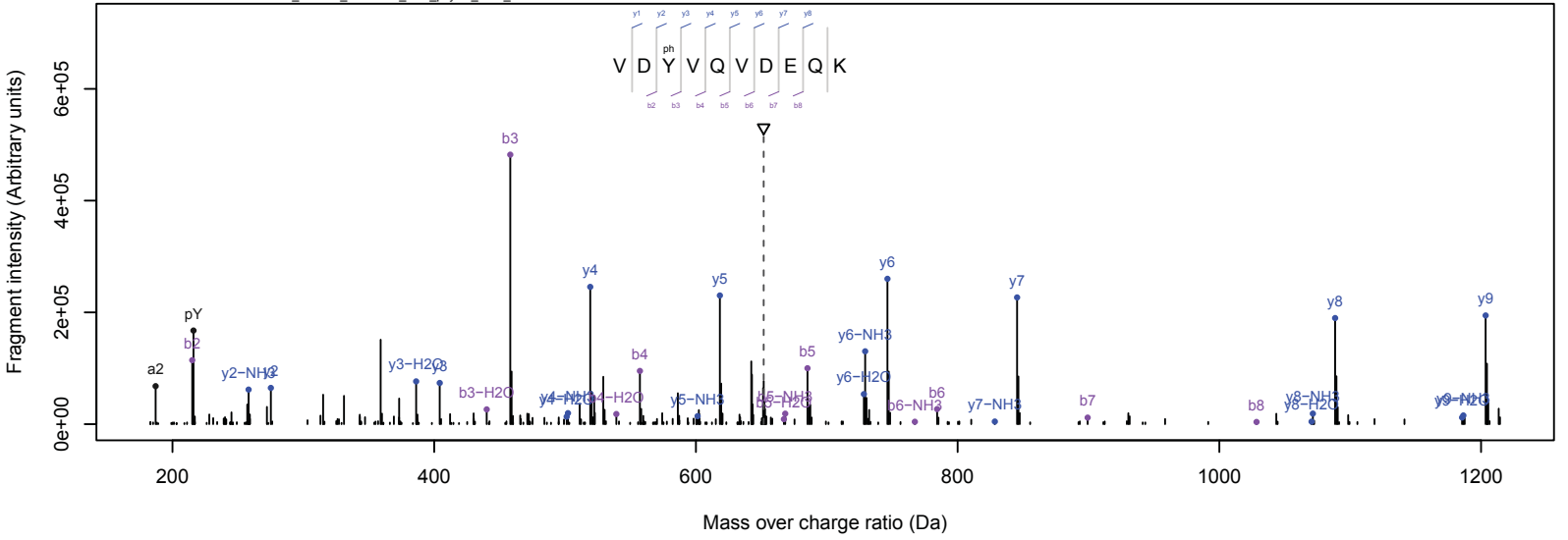
Andromeda score : 189.75



VDY(ph)VQVDEQK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 11639

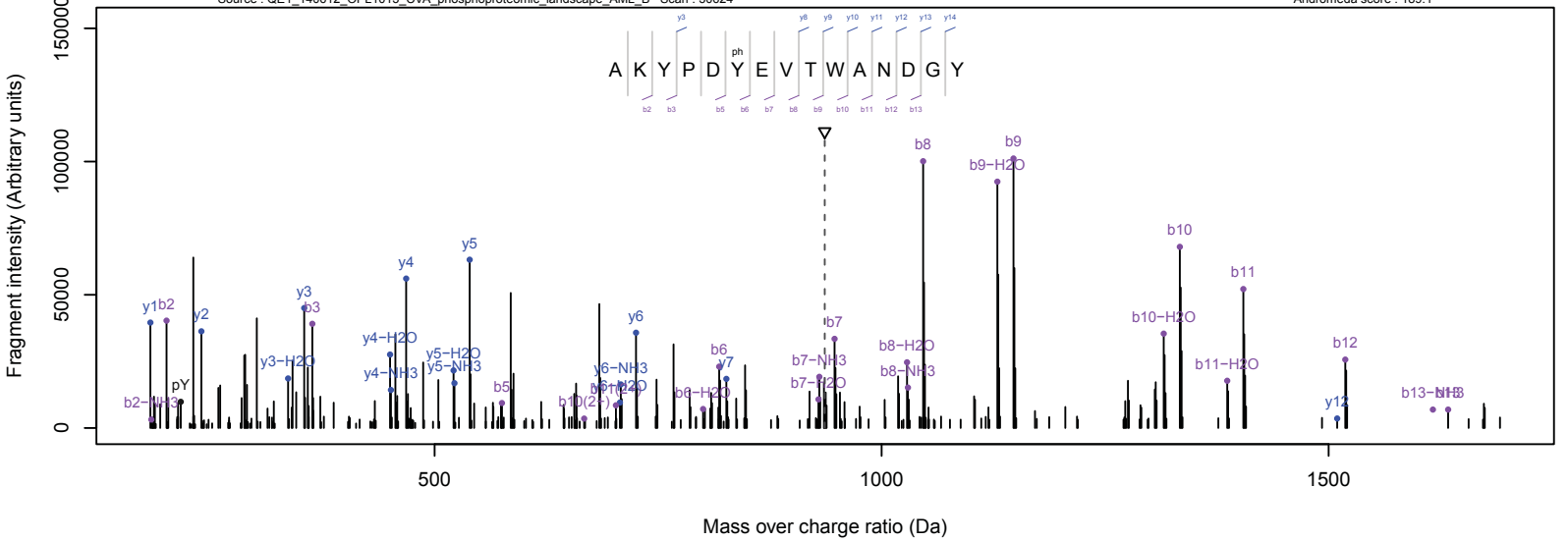
Andromeda score: 189.17



AKYPDY(ph)EVTWANDGY-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 30624

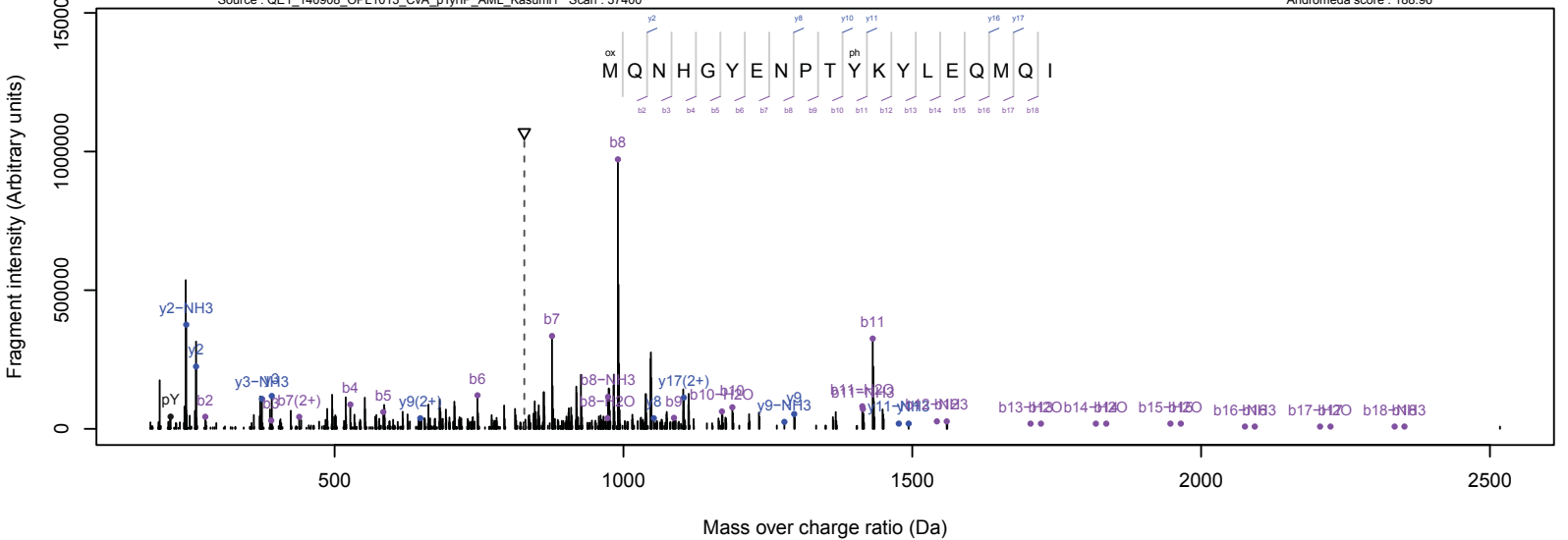
Andromeda score: 189.1



M(ox)QNHGYENPTY(ph)KYLEQMQL-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 37400

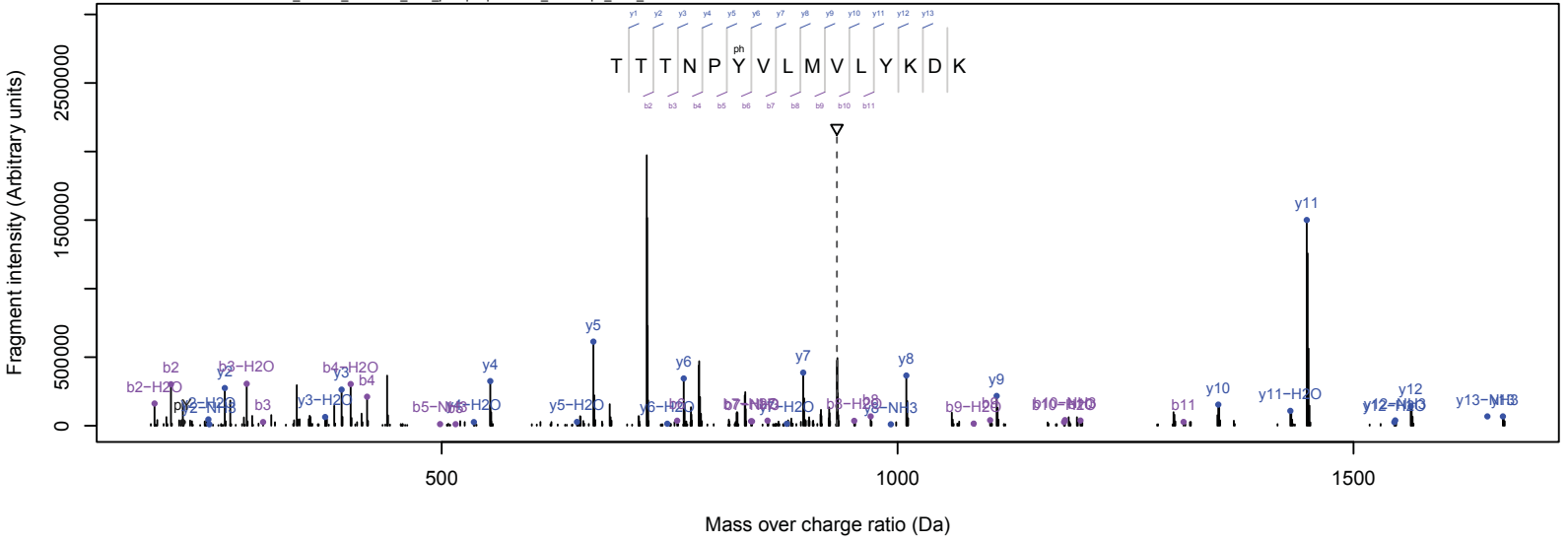
Andromeda score: 188.96



TTTNPY(ph)VLMVLYKDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 38169

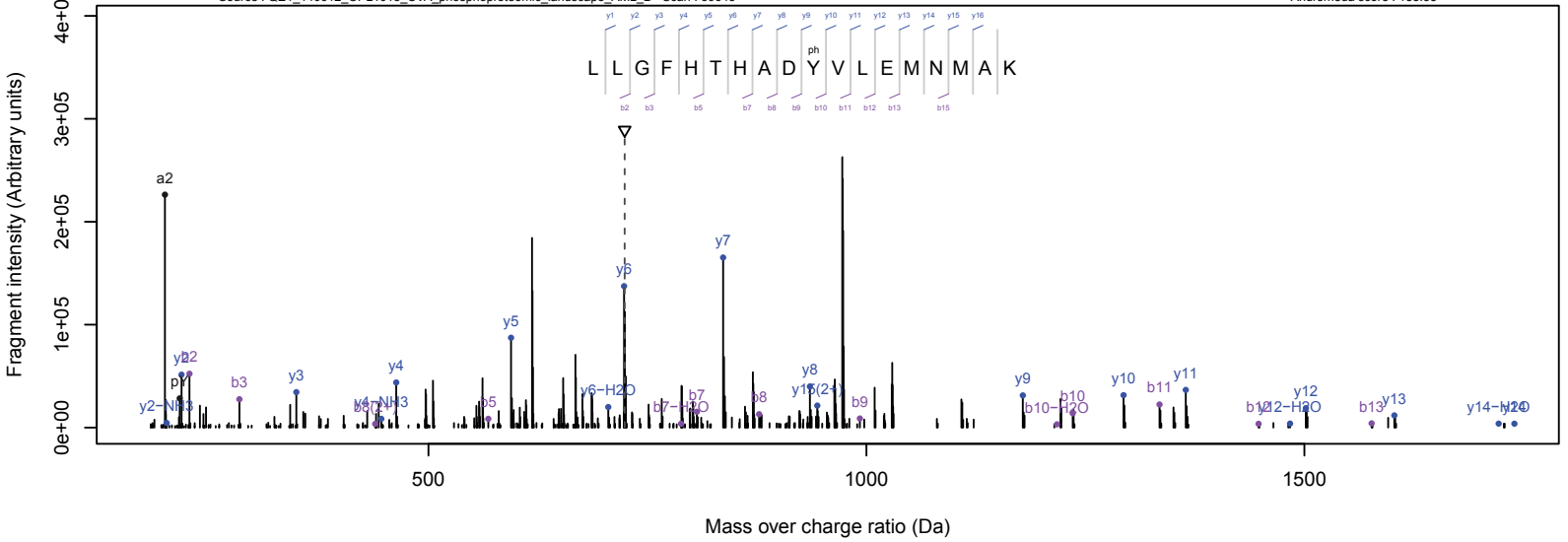
Andromeda score : 188.85



LLGFHTHADY(ph)VLENNMAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39645

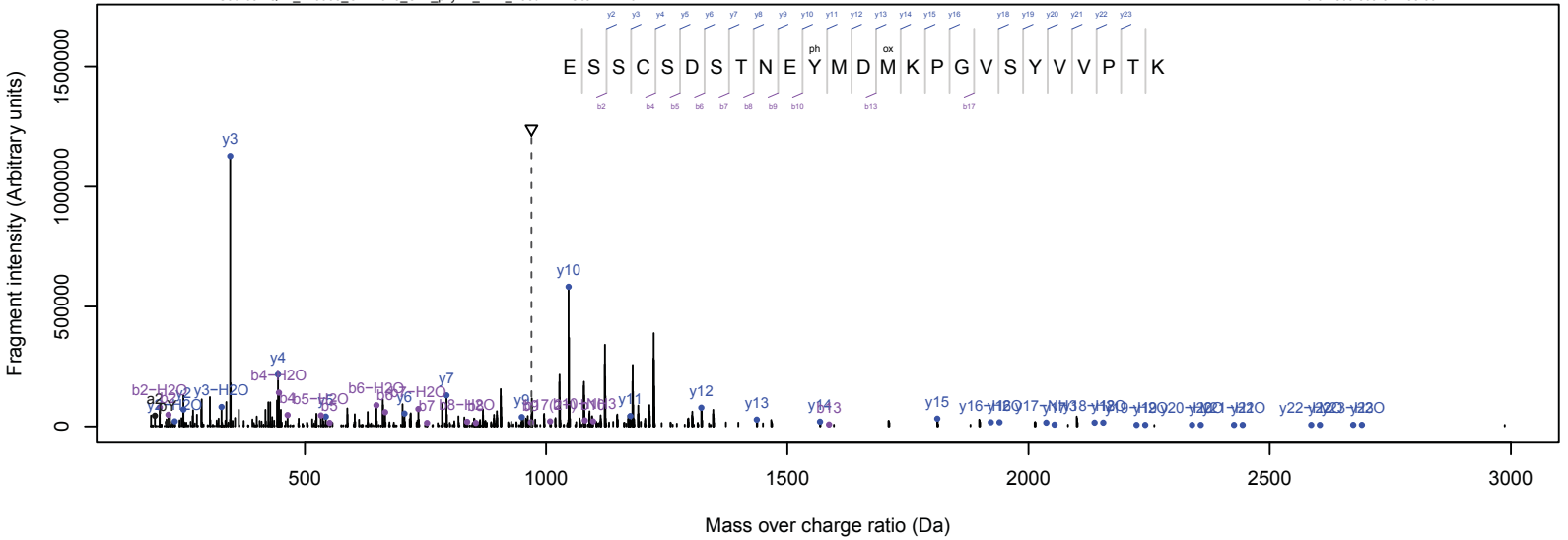
Andromeda score : 188.58



ESSCSDSTNEY(ph)MDM(ox)KPGVSYVPTK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 22494

Andromeda score : 188.39

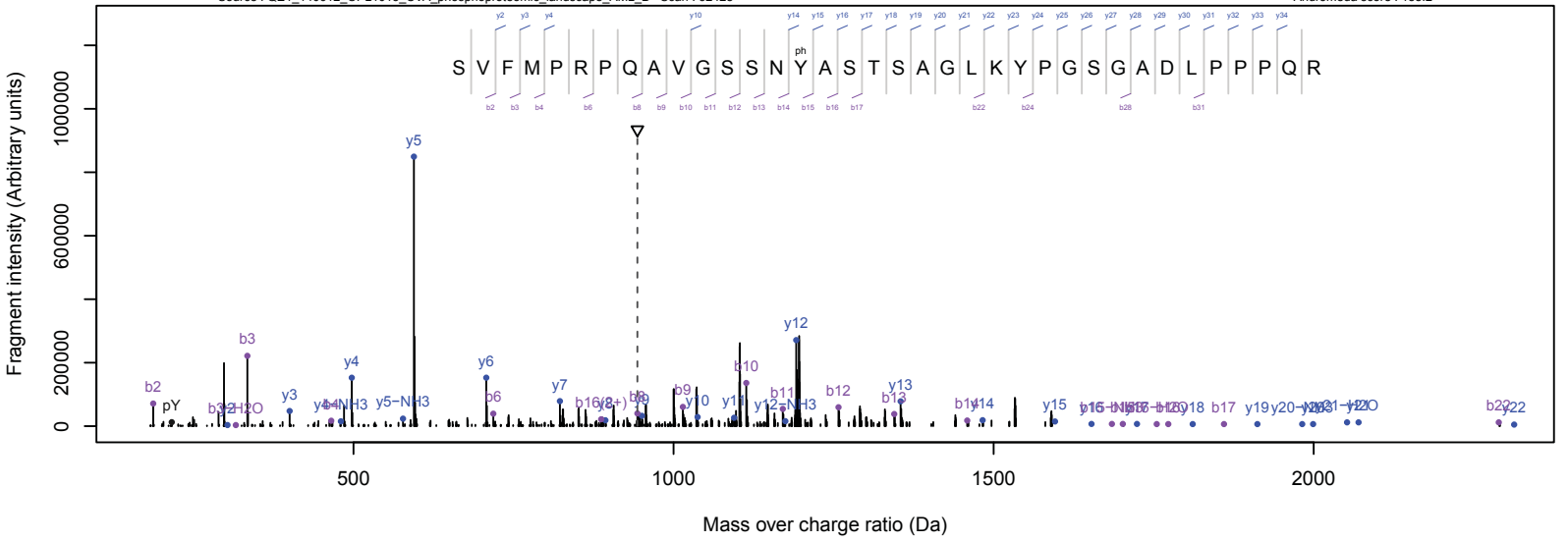




**SVFMPRPQAVGSSNYASTSAGLKYPGSGADLPPQR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32428

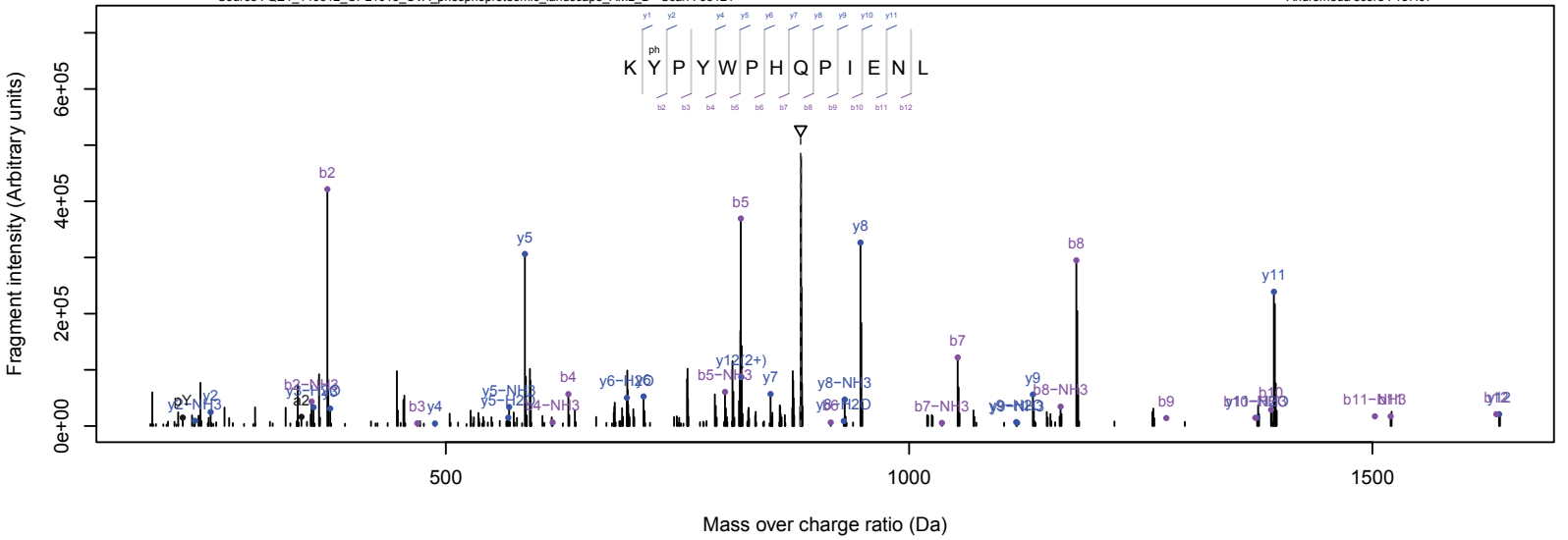
Andromeda score : 188.2



**KY(ph)PYWPHQPIENL-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33121

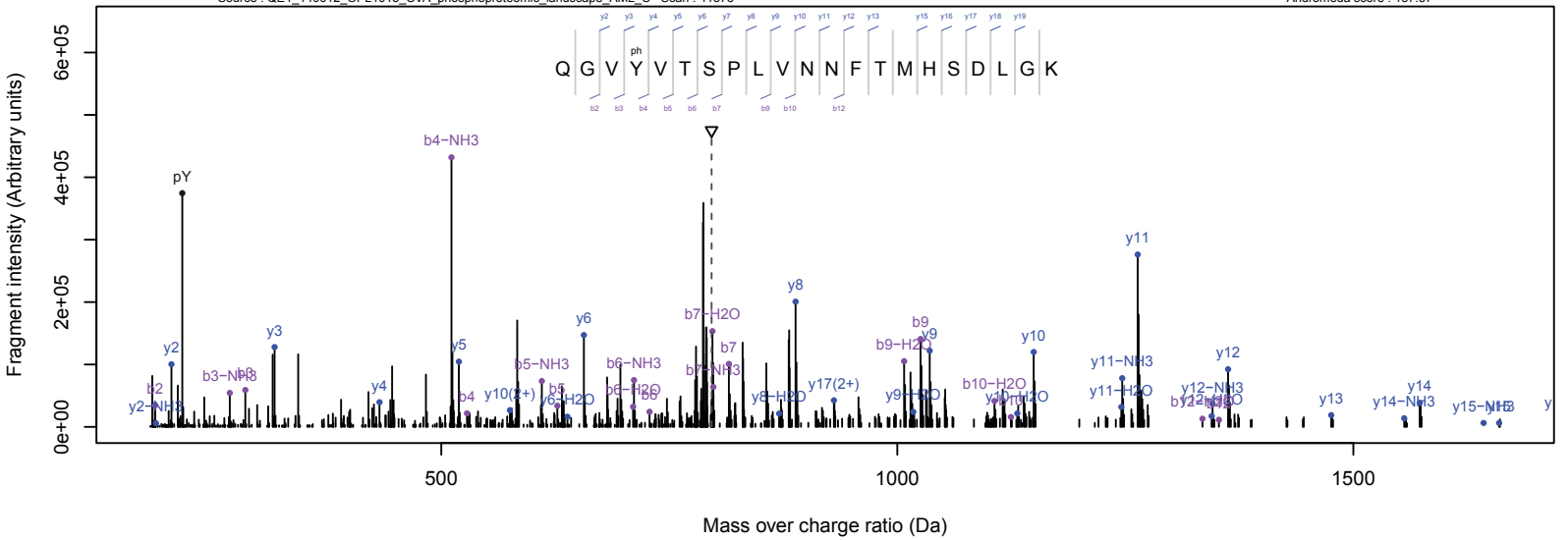
Andromeda score : 187.97



**QGVY(ph)VTSPLVNNTMHSDLGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41378

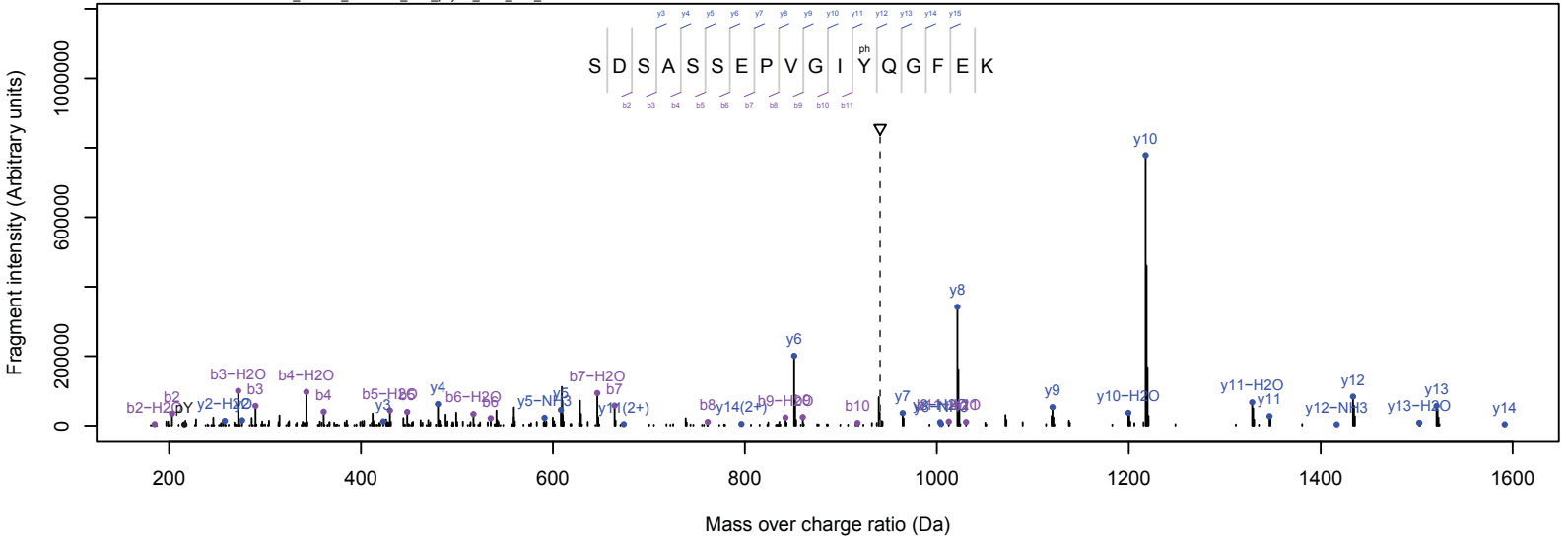
Andromeda score : 187.57



SDSASSEPVGIY(ph)QGFEK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 26645

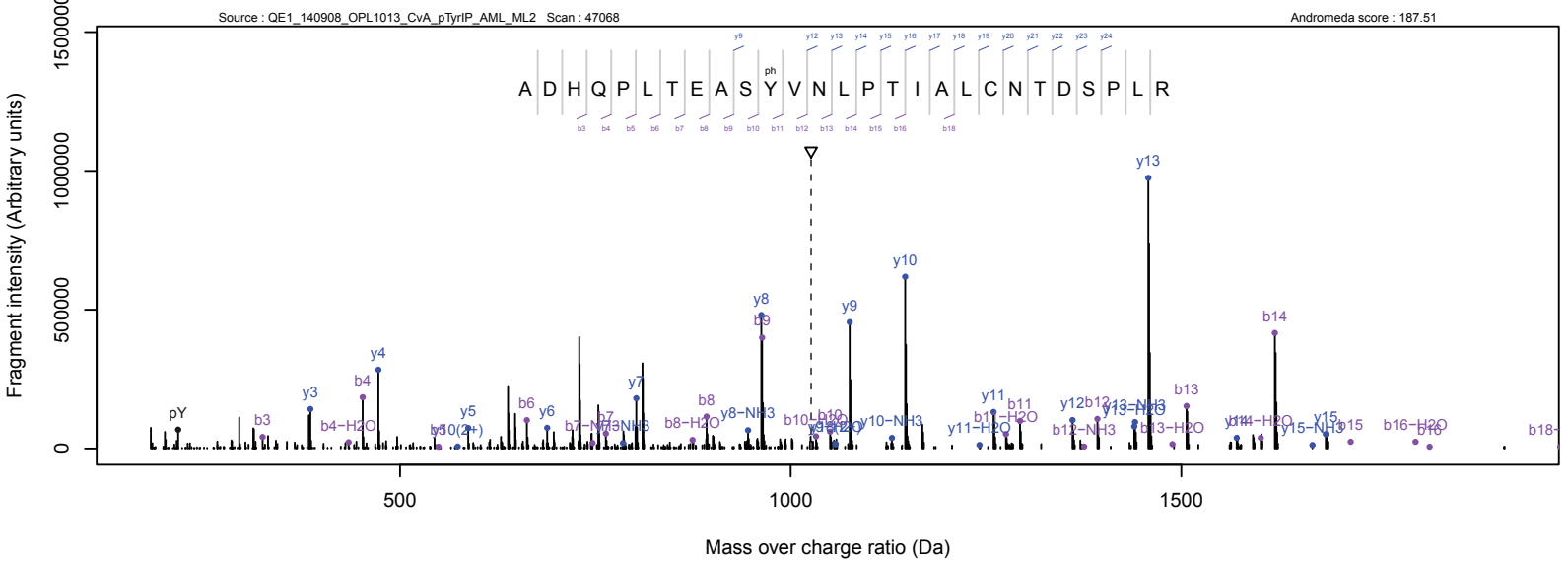
Andromeda score : 187.56



ADHQLTEASY(ph)VNLPTIALCNTDSPLR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 47068

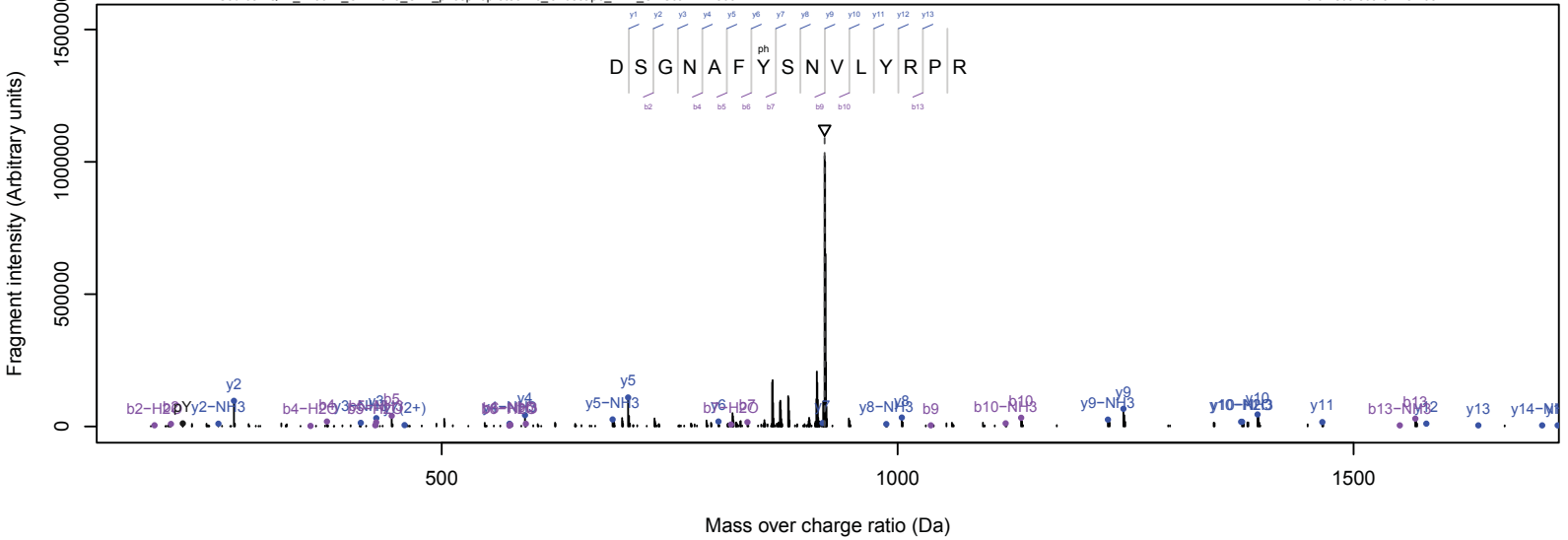
Andromeda score : 187.51

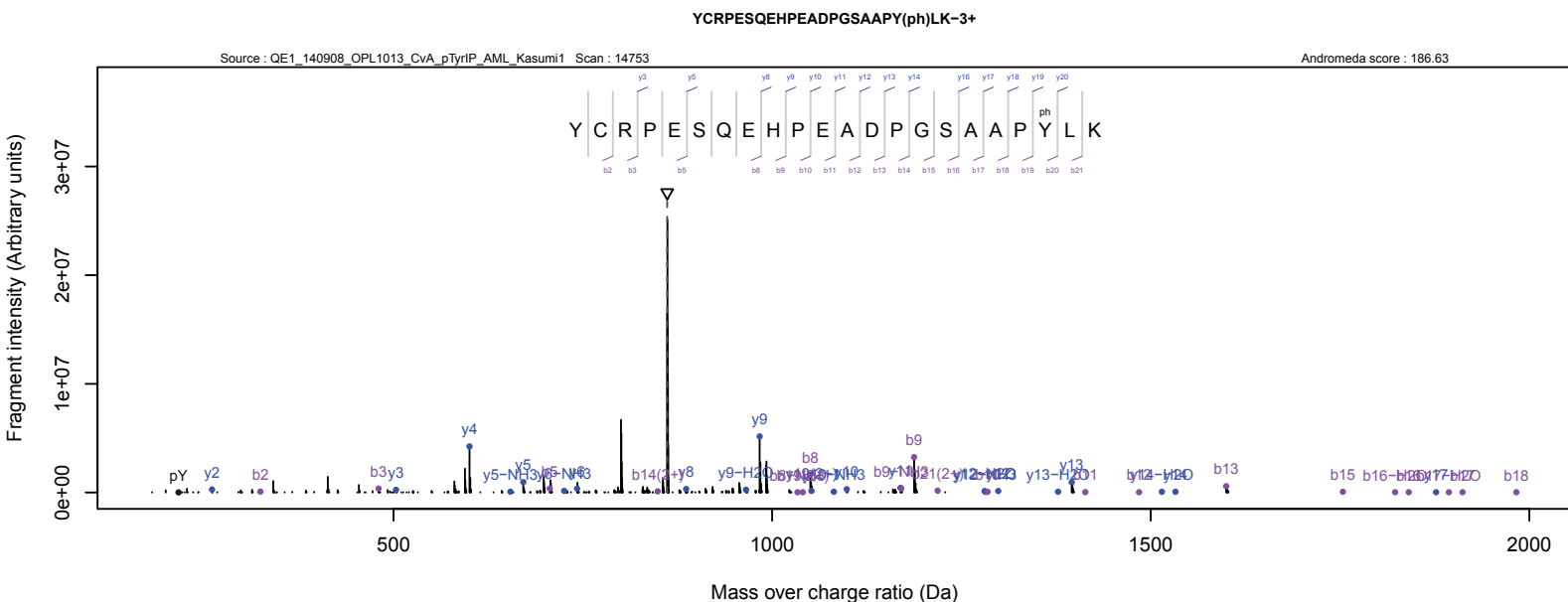
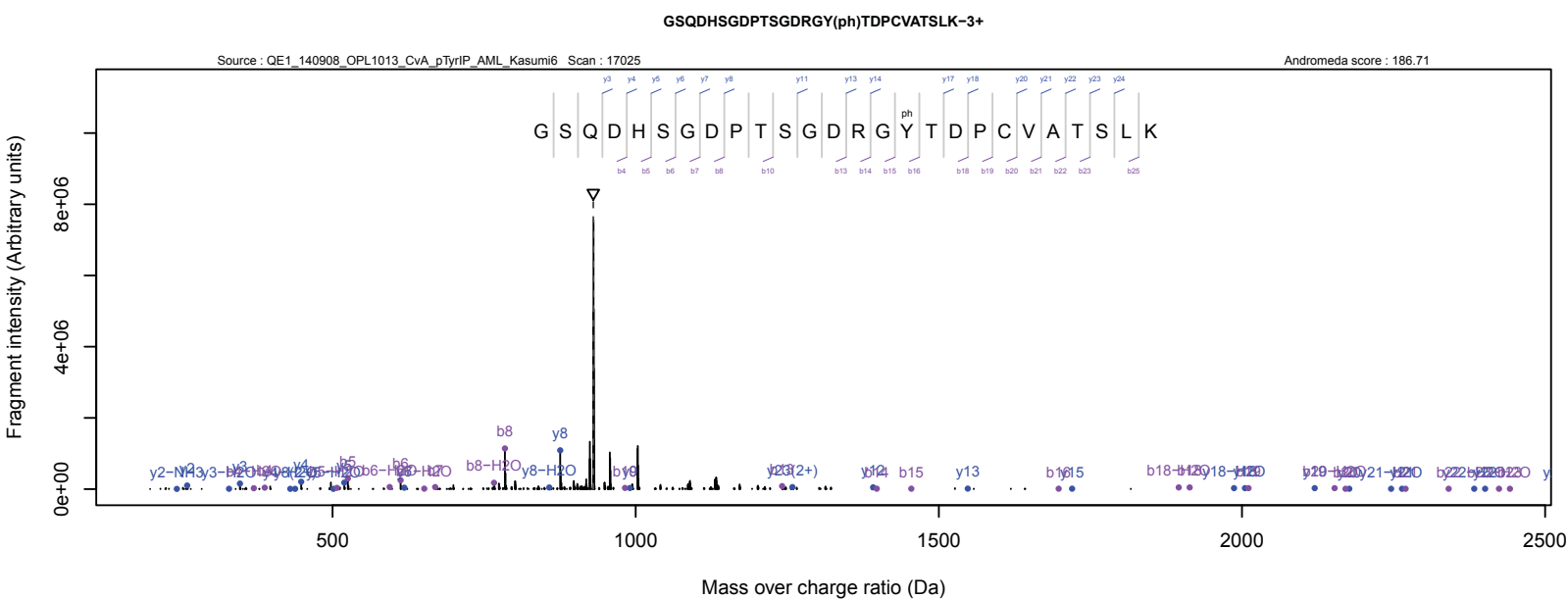
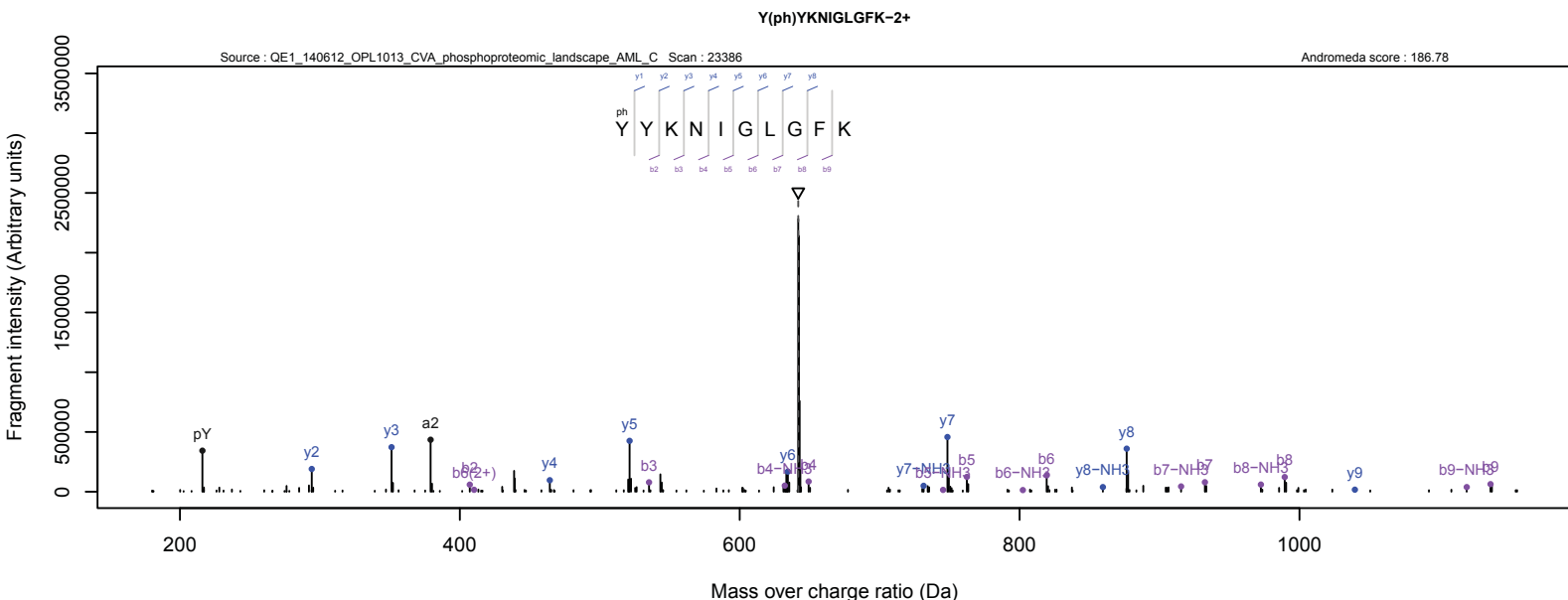


DSGNAFY(ph)SNVLYRPR-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 27865

Andromeda score : 187.08

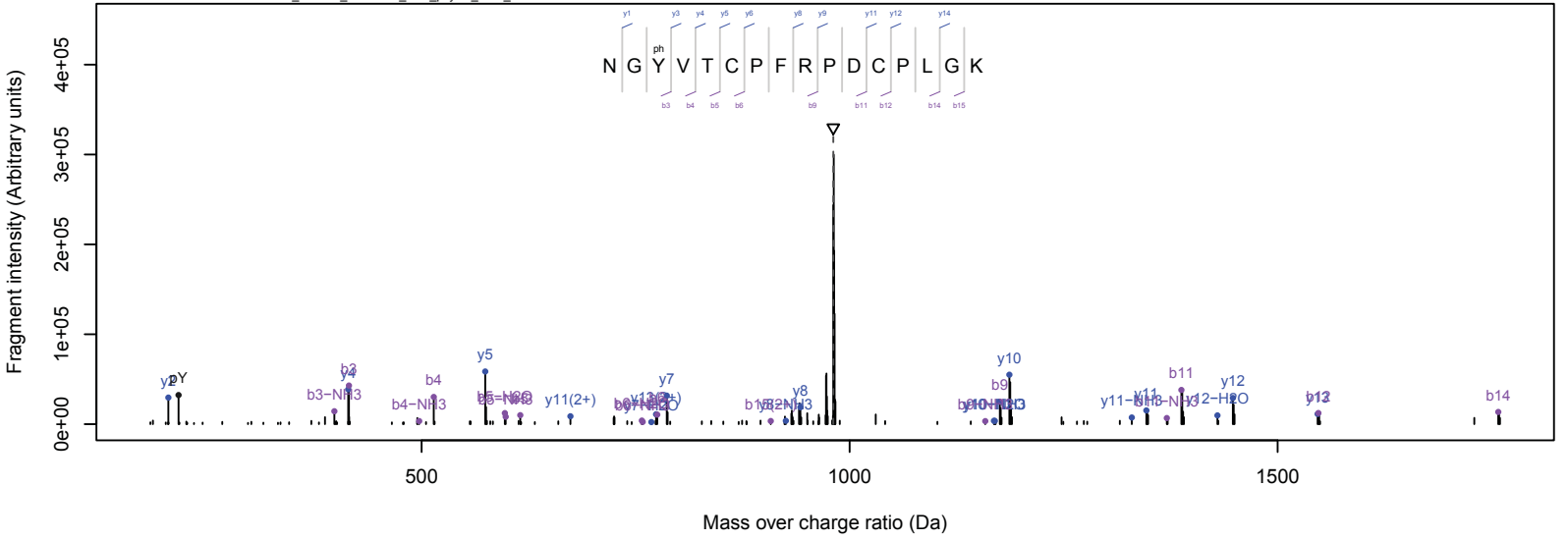




**NGY(ph)VTCPFRPDCPLGK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 27815

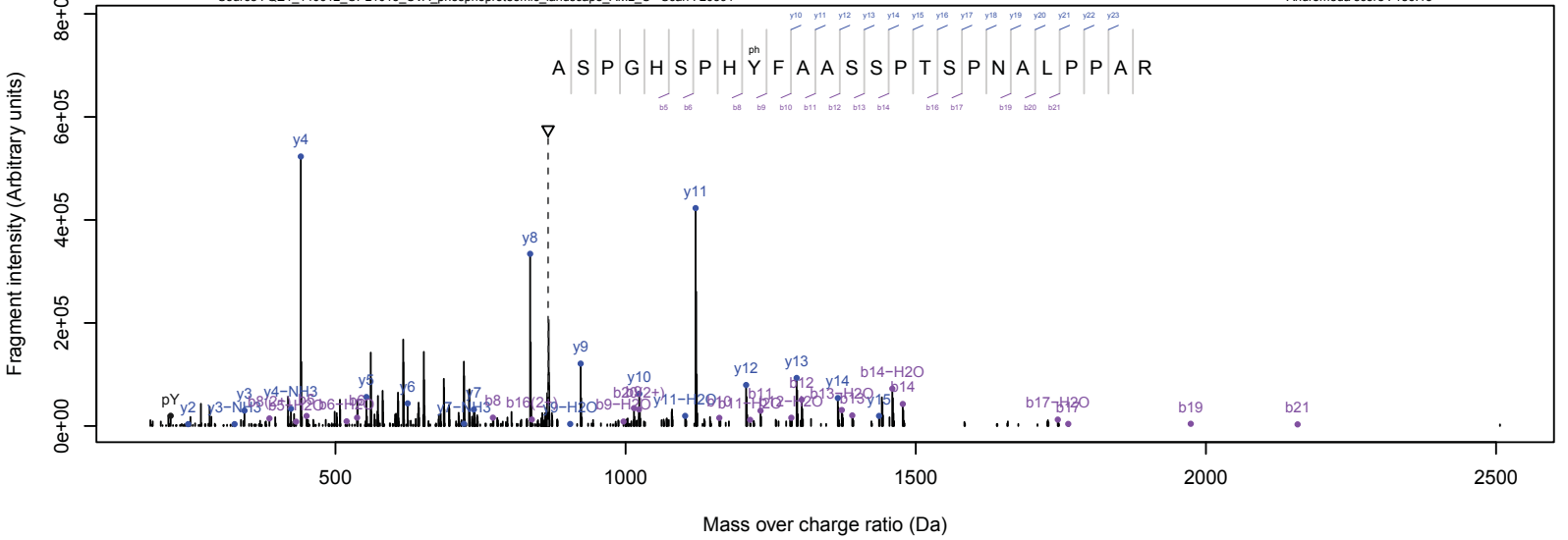
Andromeda score : 186.42



**ASPGHSPHY(ph)FAASSPTSPNALPPAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20391

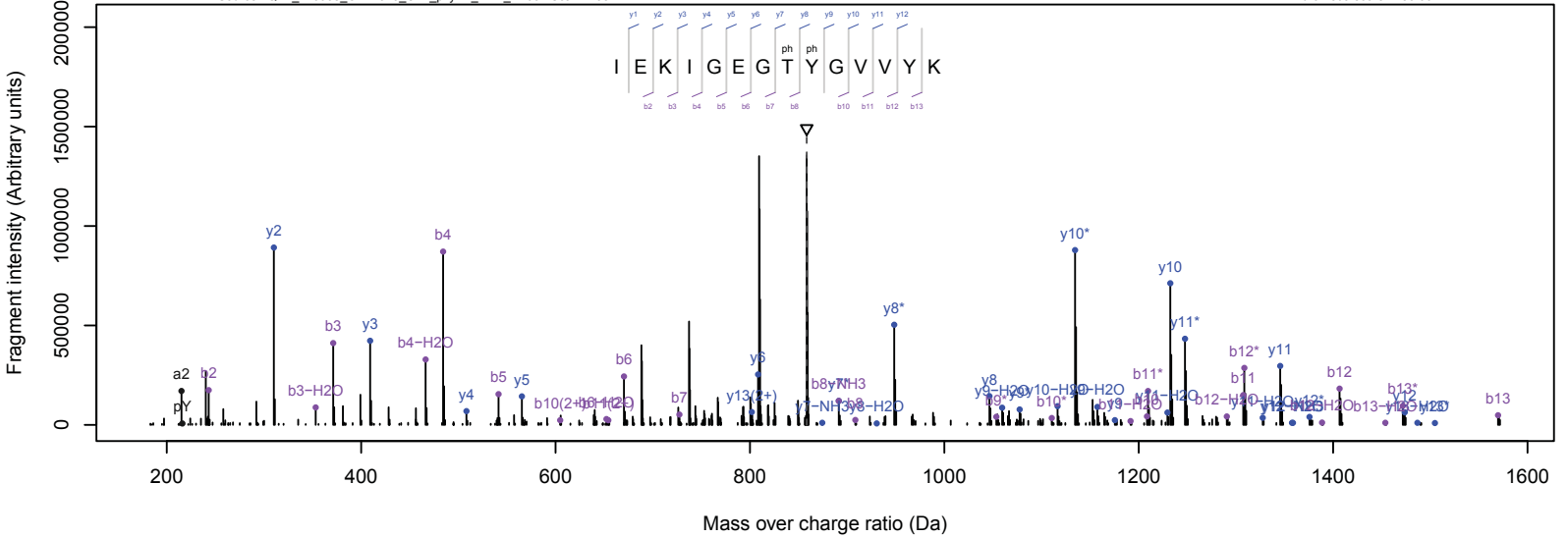
Andromeda score : 186.13



**IEKIGEGT(ph)Y(ph)GVVYK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 23822

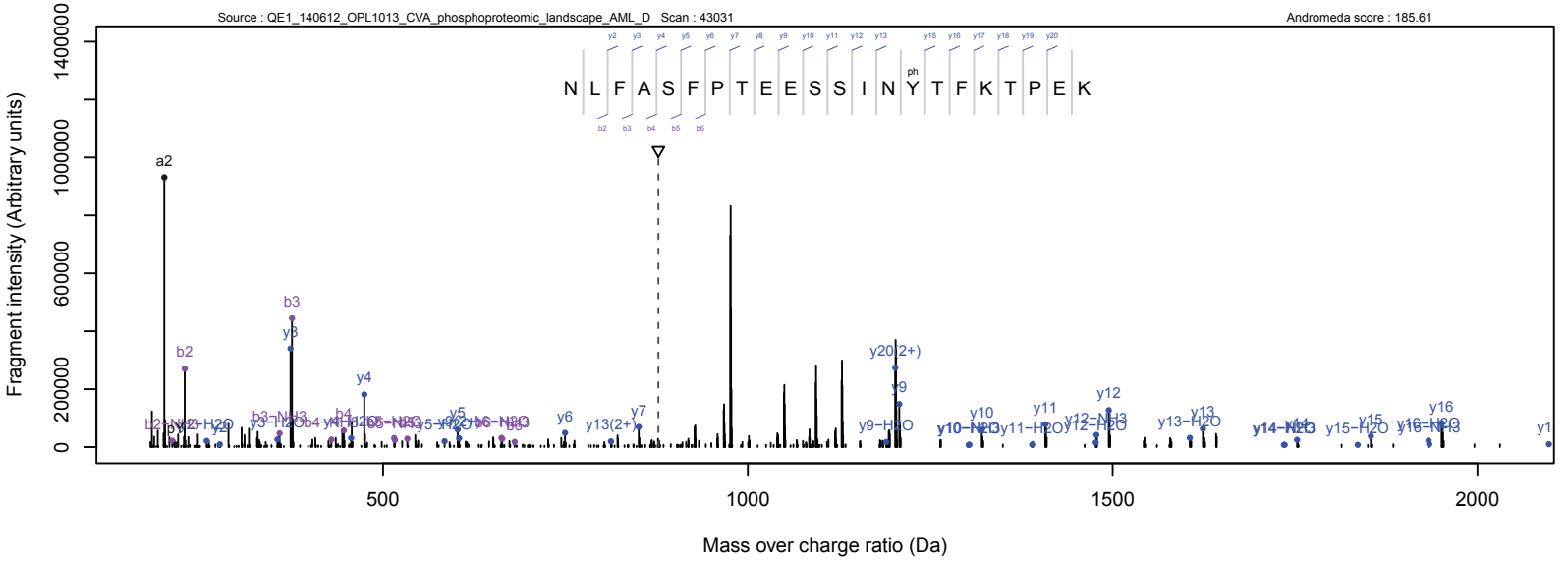
Andromeda score : 186.06



NLFASFPTEESSINY(ph)TFKTPEK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 43031

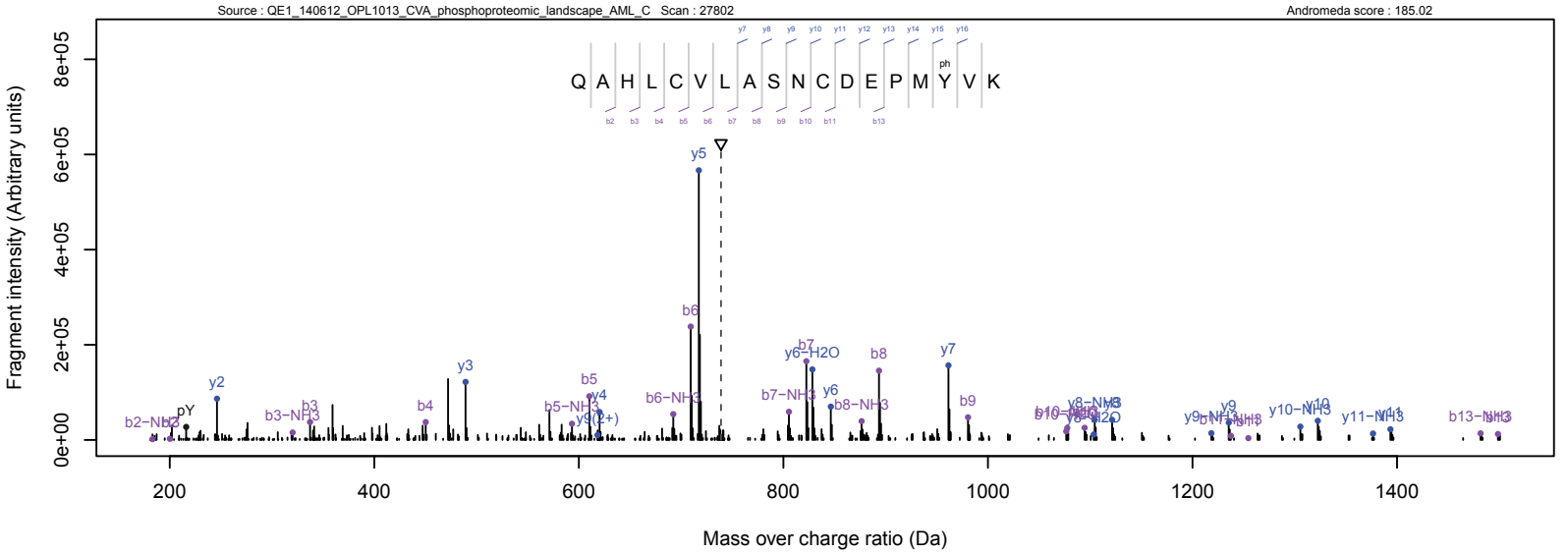
Andromeda score : 185.61



QAHLCLVLANCDEPMY(ph)VK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27802

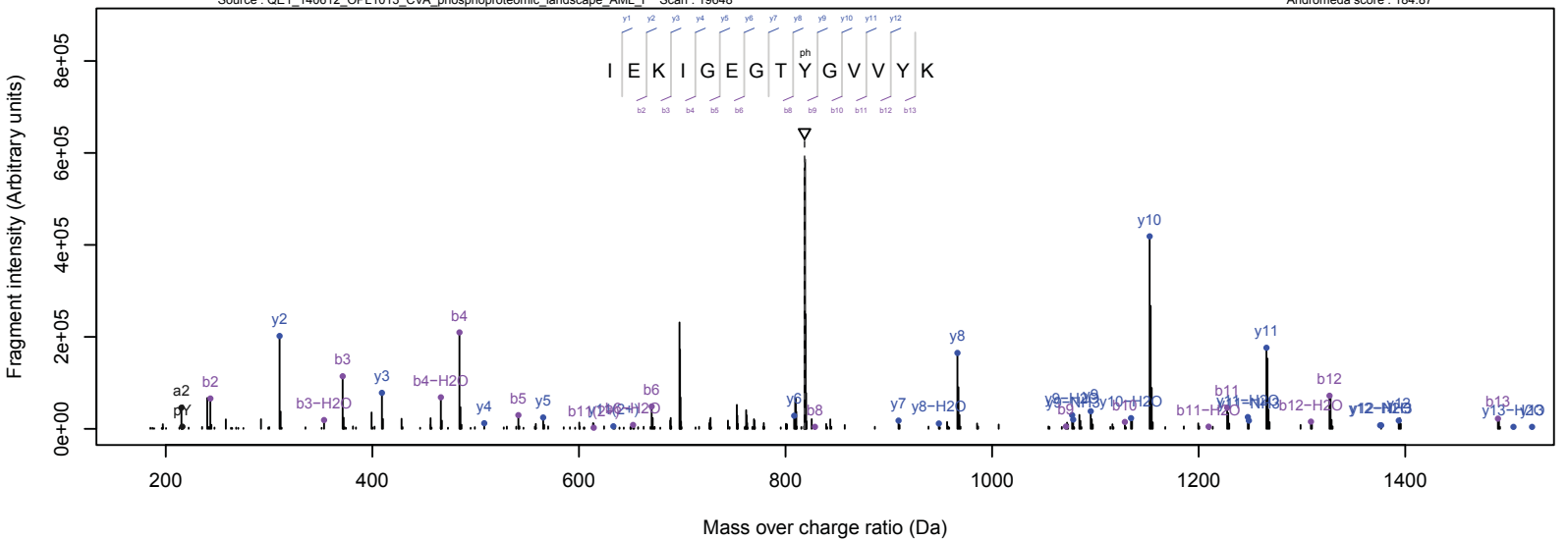
Andromeda score : 185.02



IEKIGEGTY(ph)GVVYK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 19648

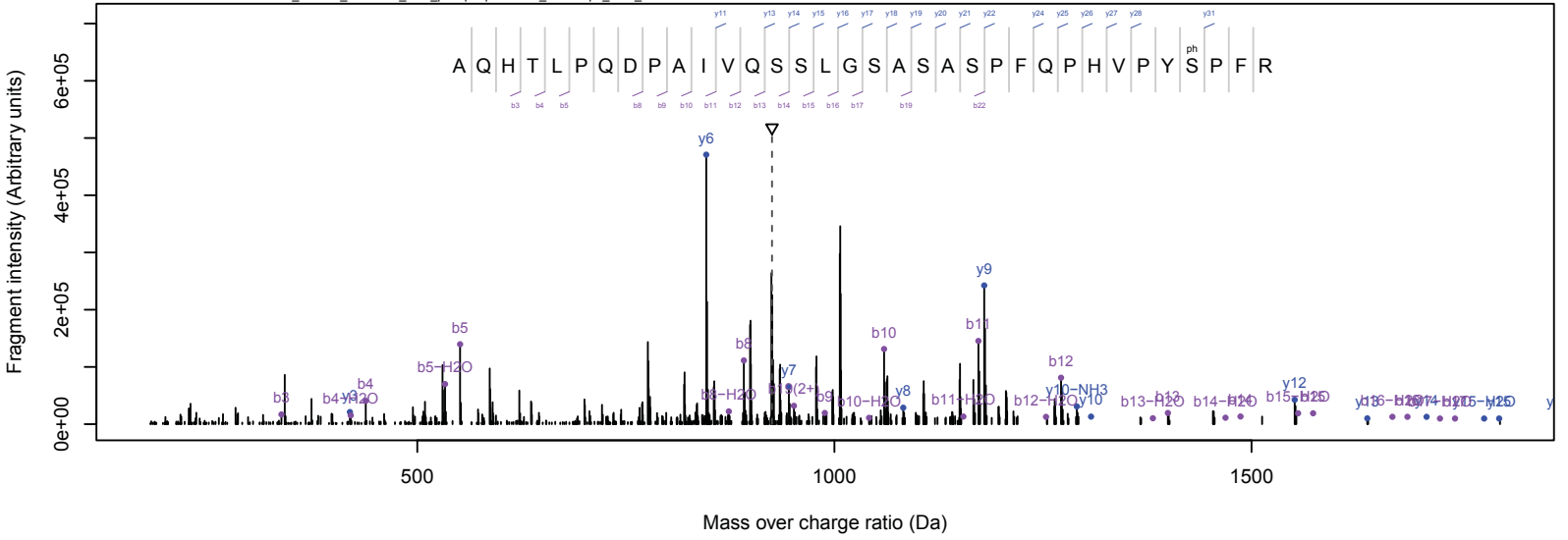
Andromeda score : 184.87



**AQHTLPQDPAIVQSSLSGSASASPFQPHVPYS(ph)PFR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42103

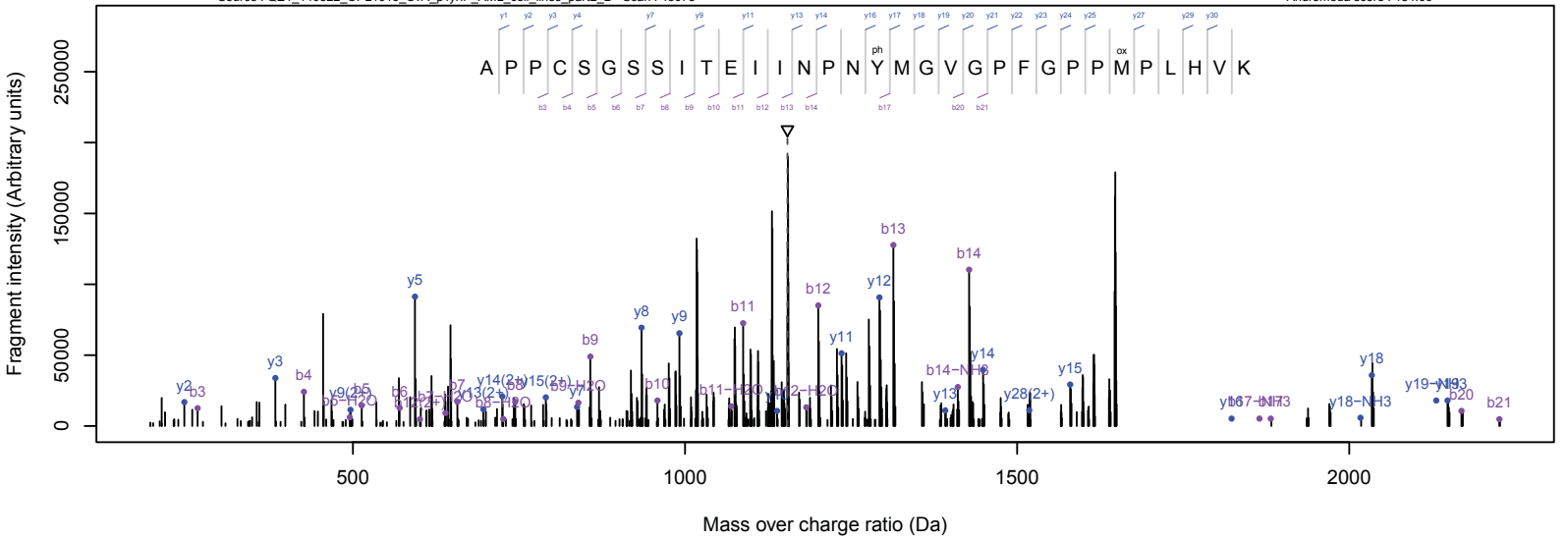
Andromeda score : 184.82



**APPCSGSSITEIINPNYMGVGFPPM(ox)PLHVK-3+**

Source : QE1\_140822\_OPL1013\_Cva\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 45078

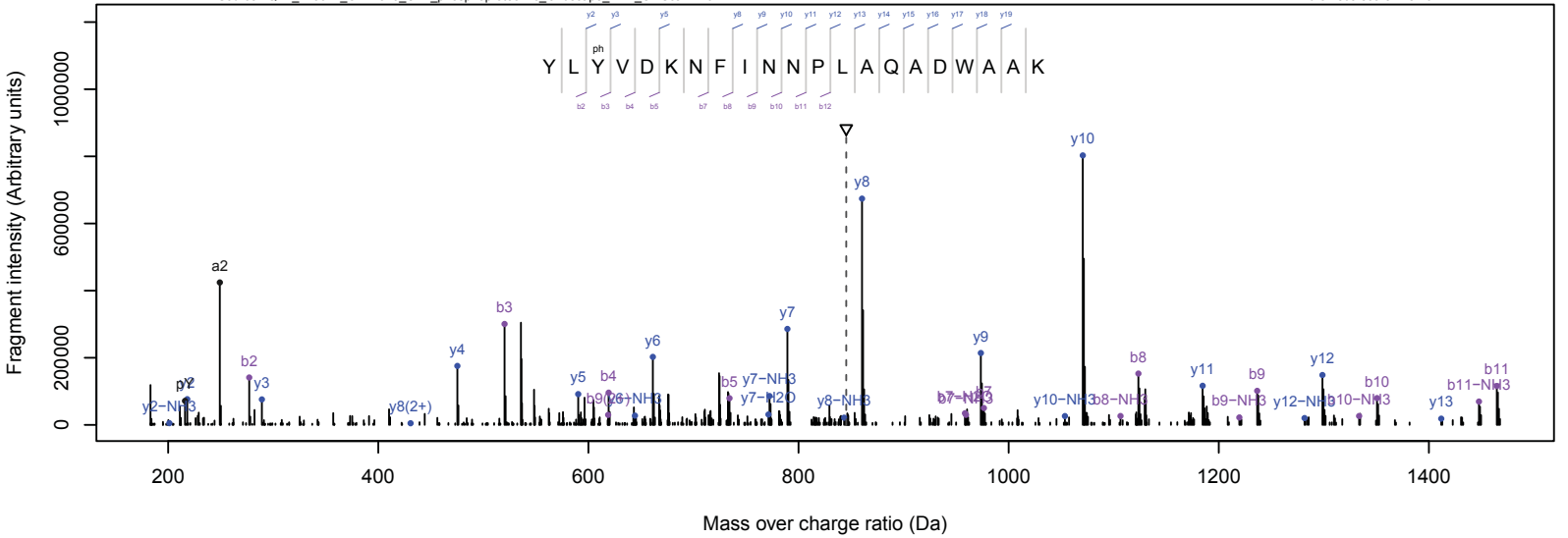
Andromeda score : 184.66



**YLY(ph)VDKFINNPLAQADWAAK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46412

Andromeda score : 184.6

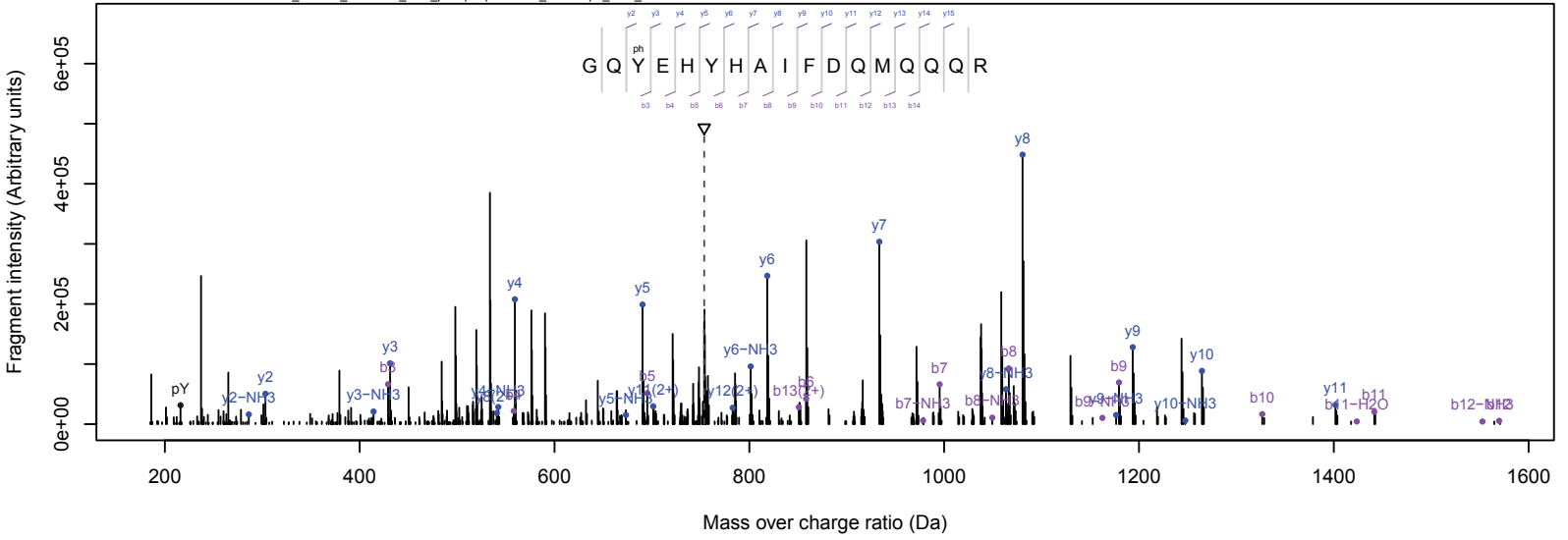




**GQY(ph)EHYHAIFDQMQQQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23845

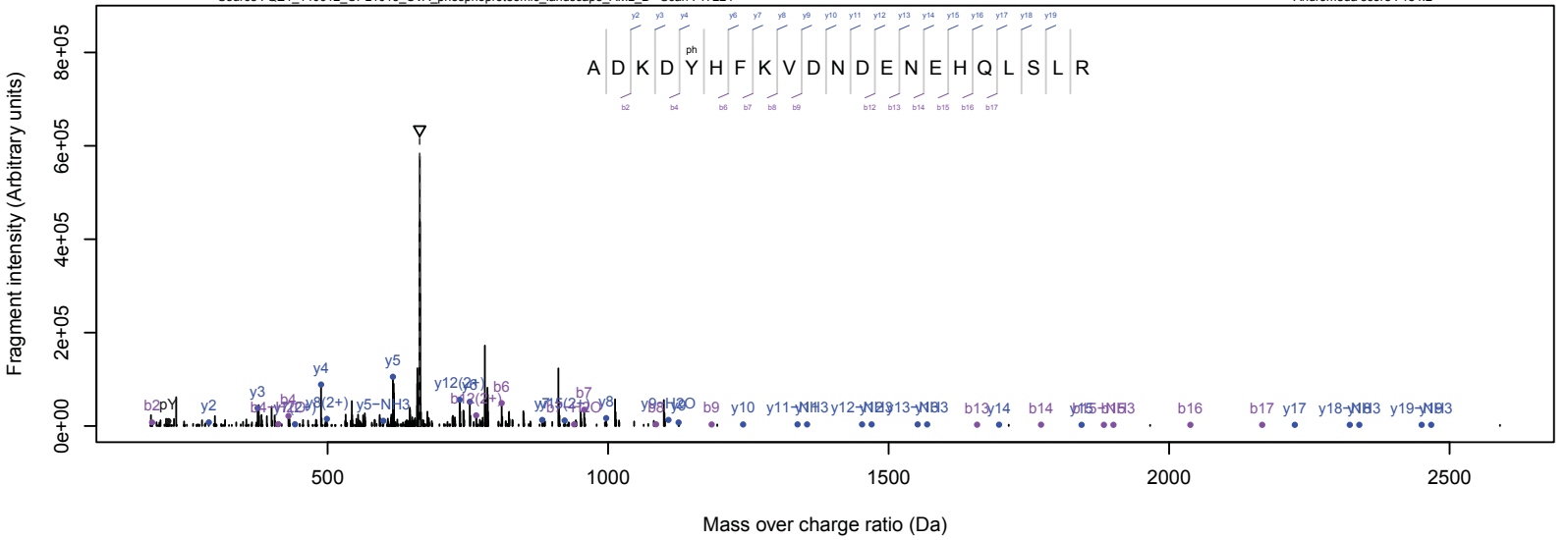
Andromeda score : 184.21



**ADKDY(ph)HFKVDNENEHQLSLR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17224

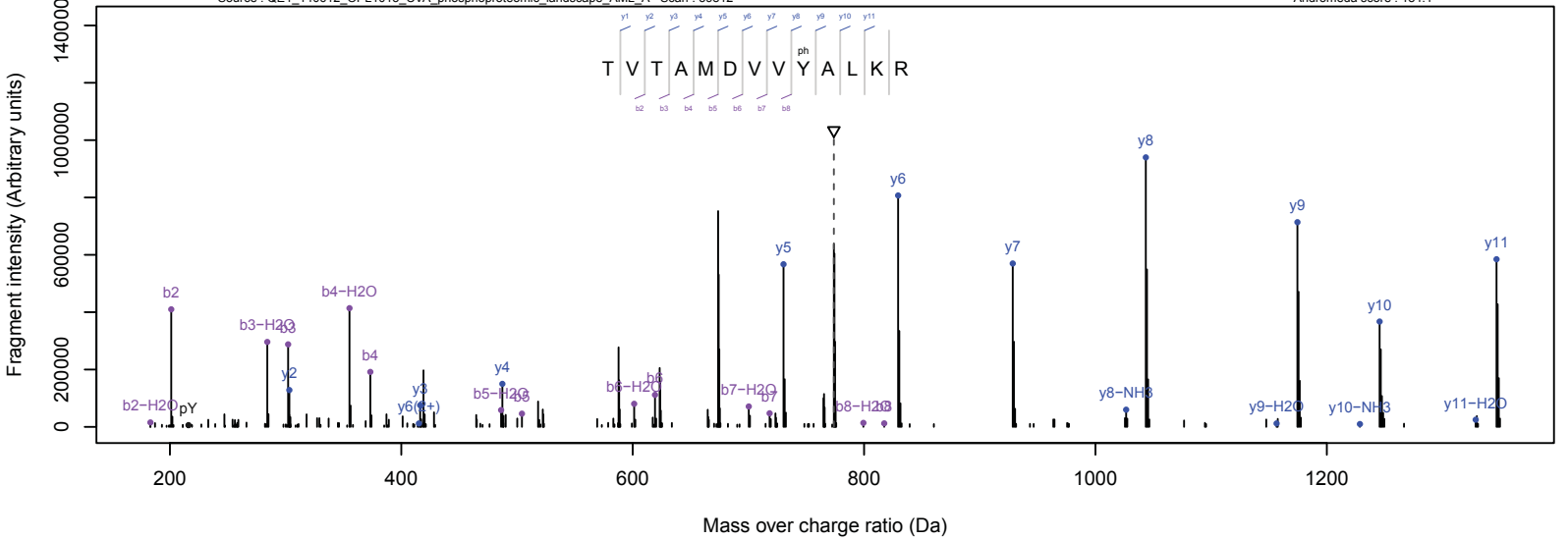
Andromeda score : 184.2



**TVTAMDVVY(ph)ALKR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 39812

Andromeda score : 184.1

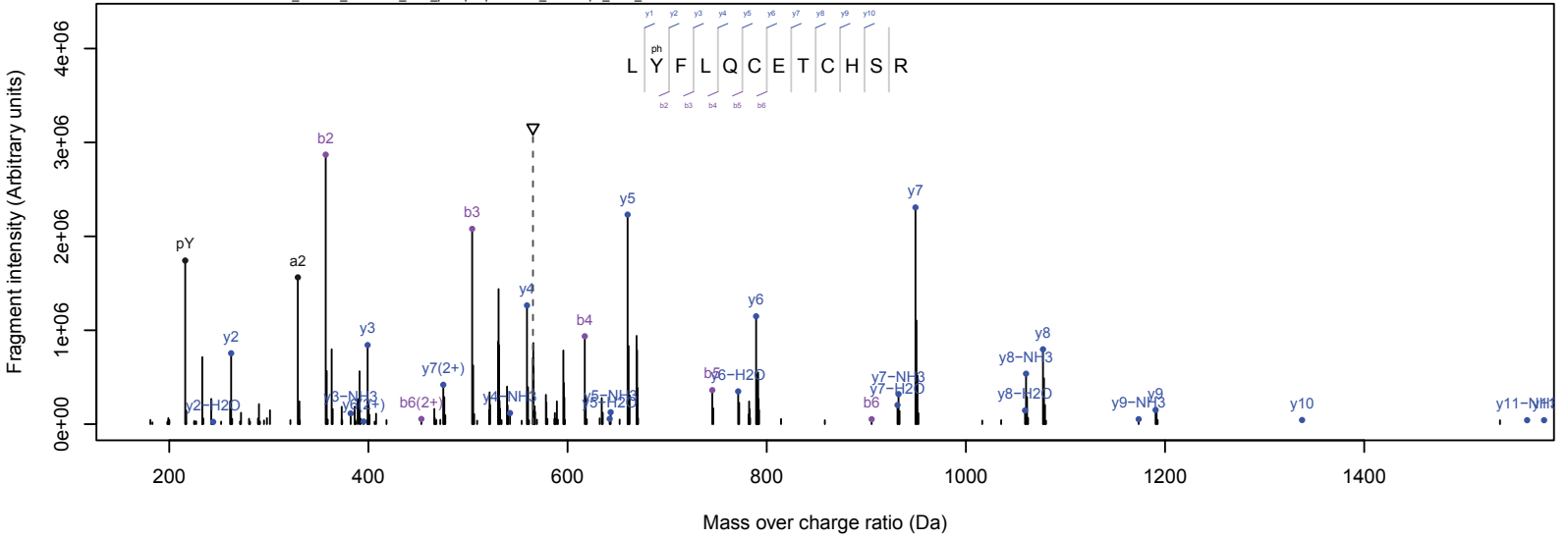




LY(ph)FLQCETCHSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23661

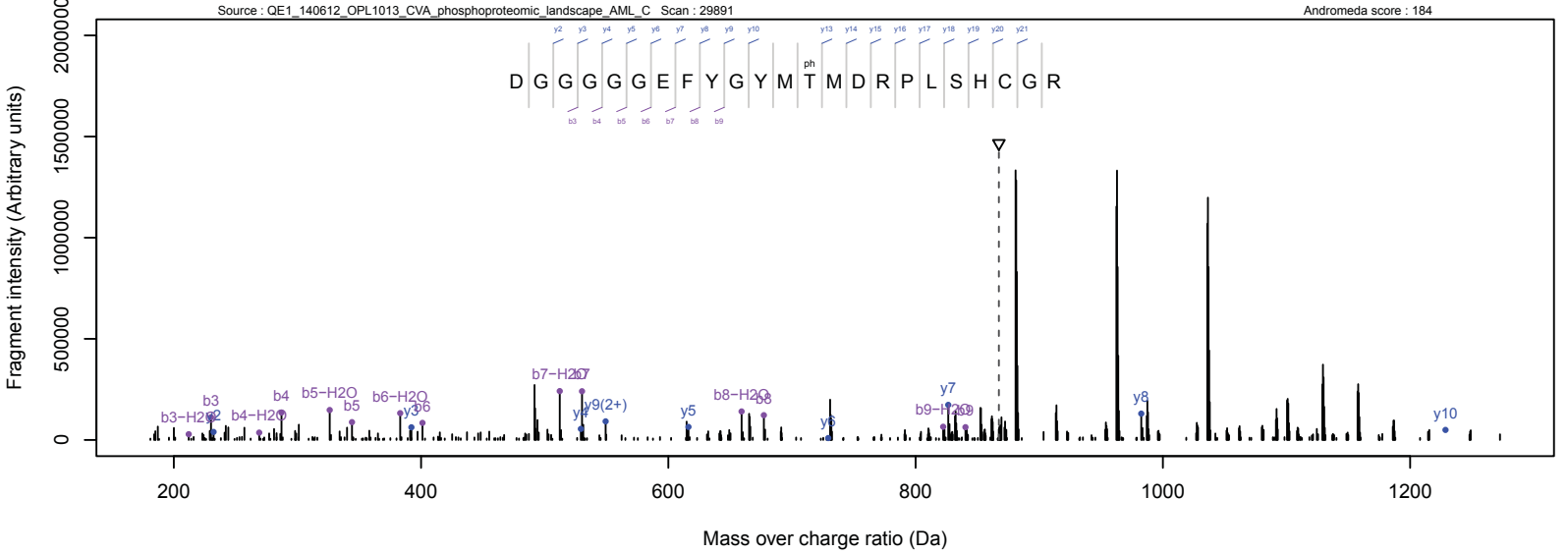
Andromeda score : 184.03



DGGGGGEFYGYM(ph)MDRPLSHCGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29891

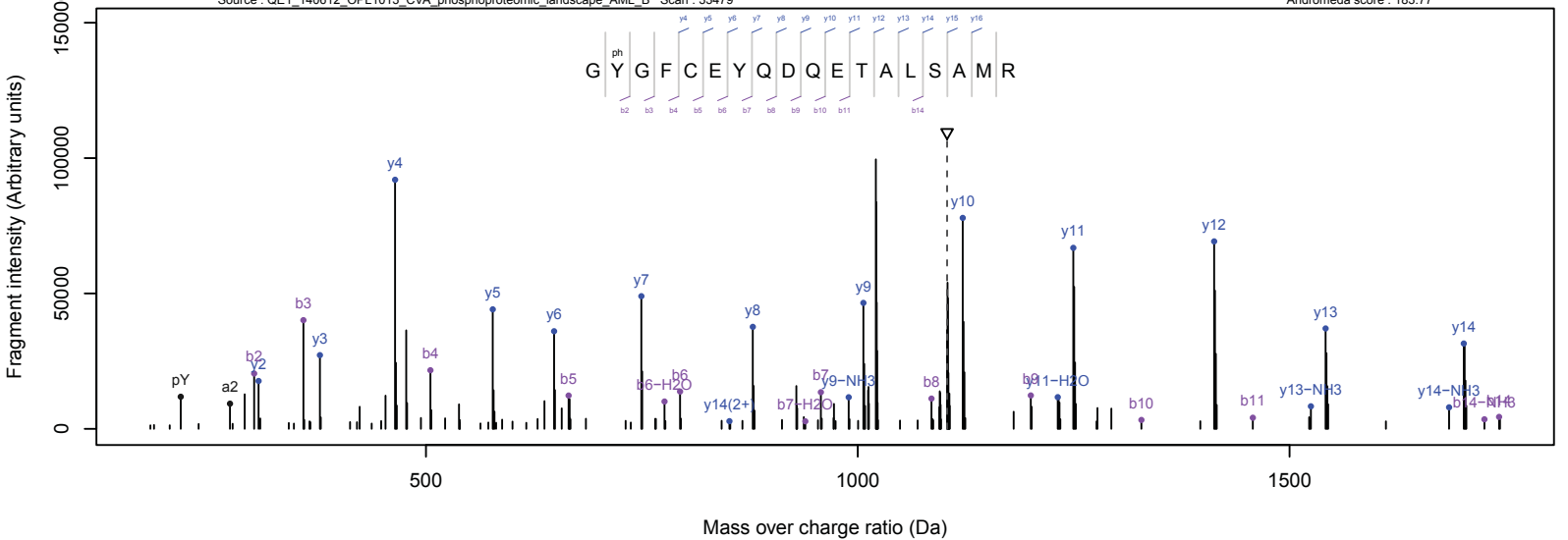
Andromeda score : 184



GY(ph)GFCEYQDQETALSAMR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33479

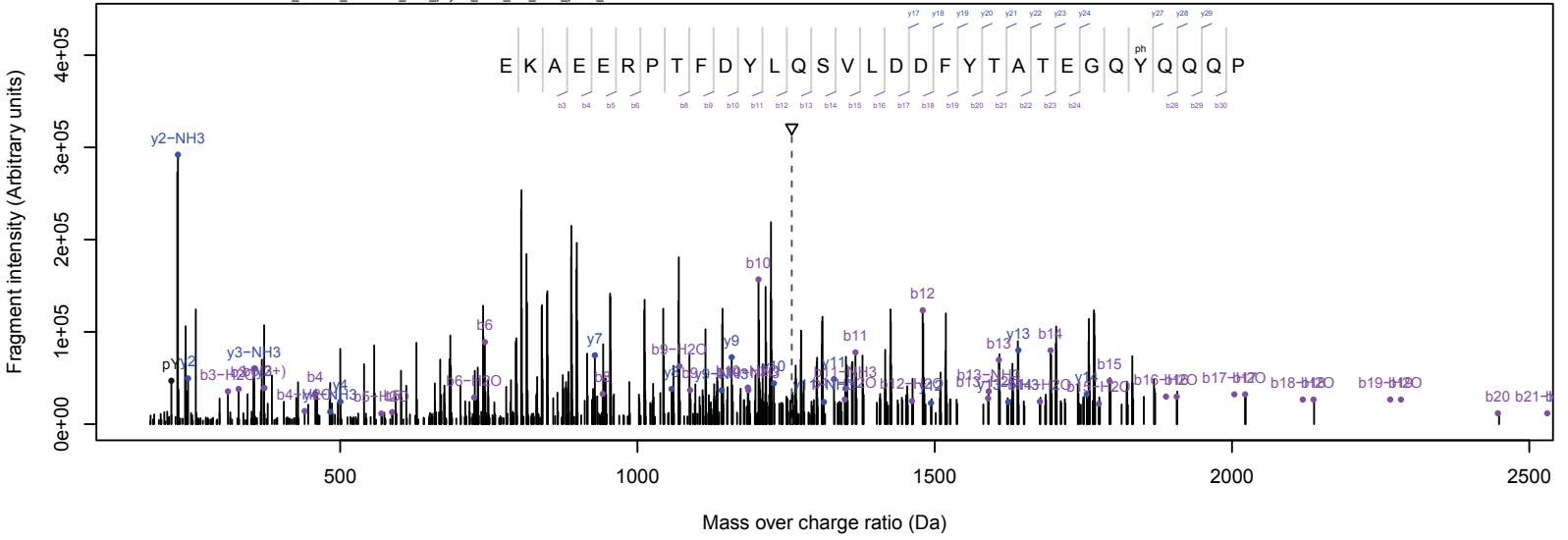
Andromeda score : 183.77



**EKAERPTFDYLSVLDDFYTATEGQYQQP-3+**

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 45436

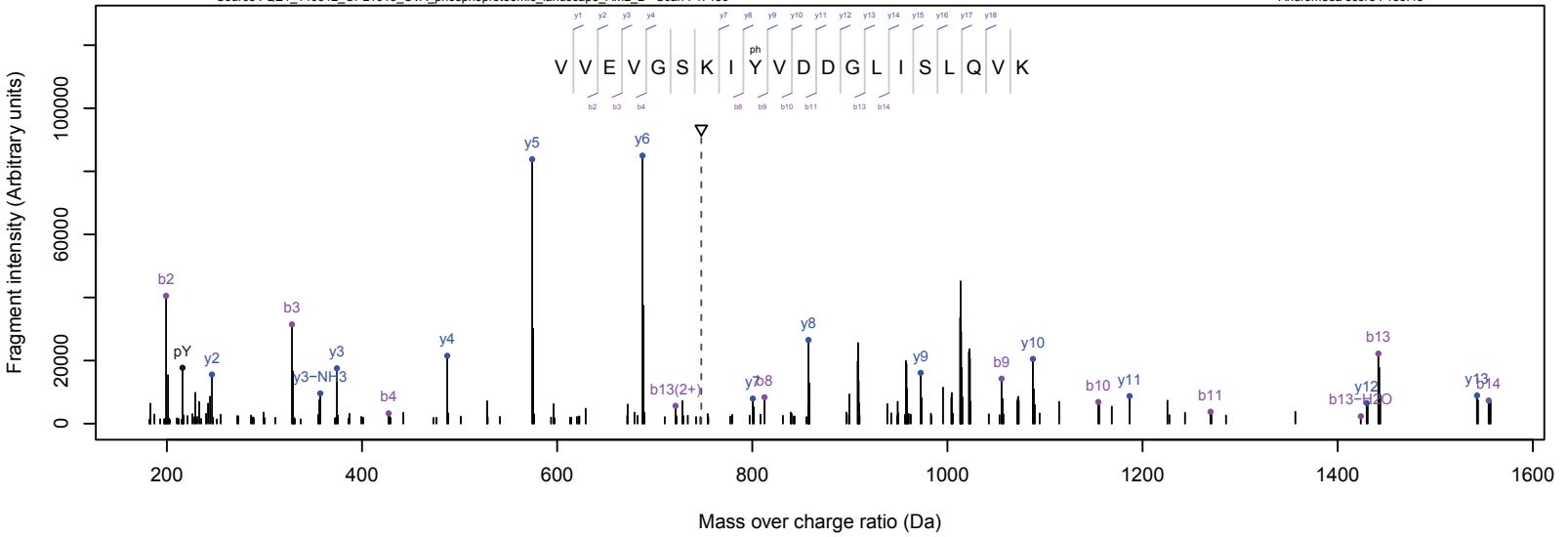
Andromeda score: 183.49



**VVEVGSKIYVDDGLISLQVK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 47486

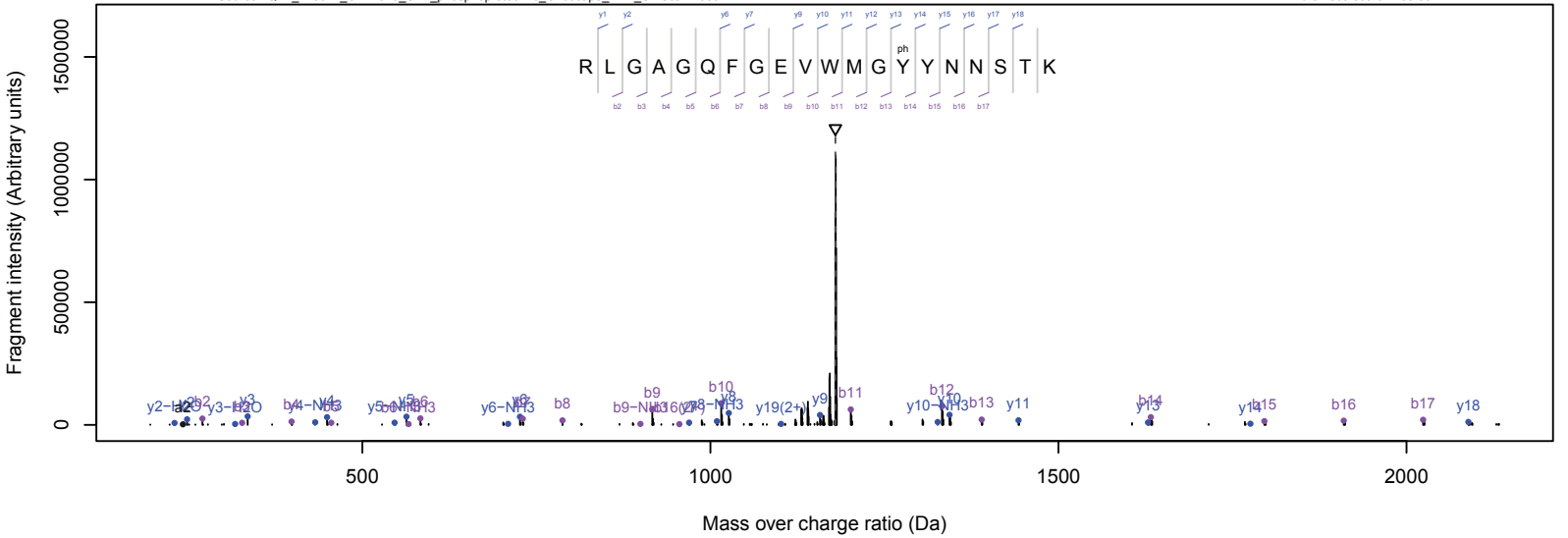
Andromeda score: 183.45



**RLGAGQFGEVWGYNNSTK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36517

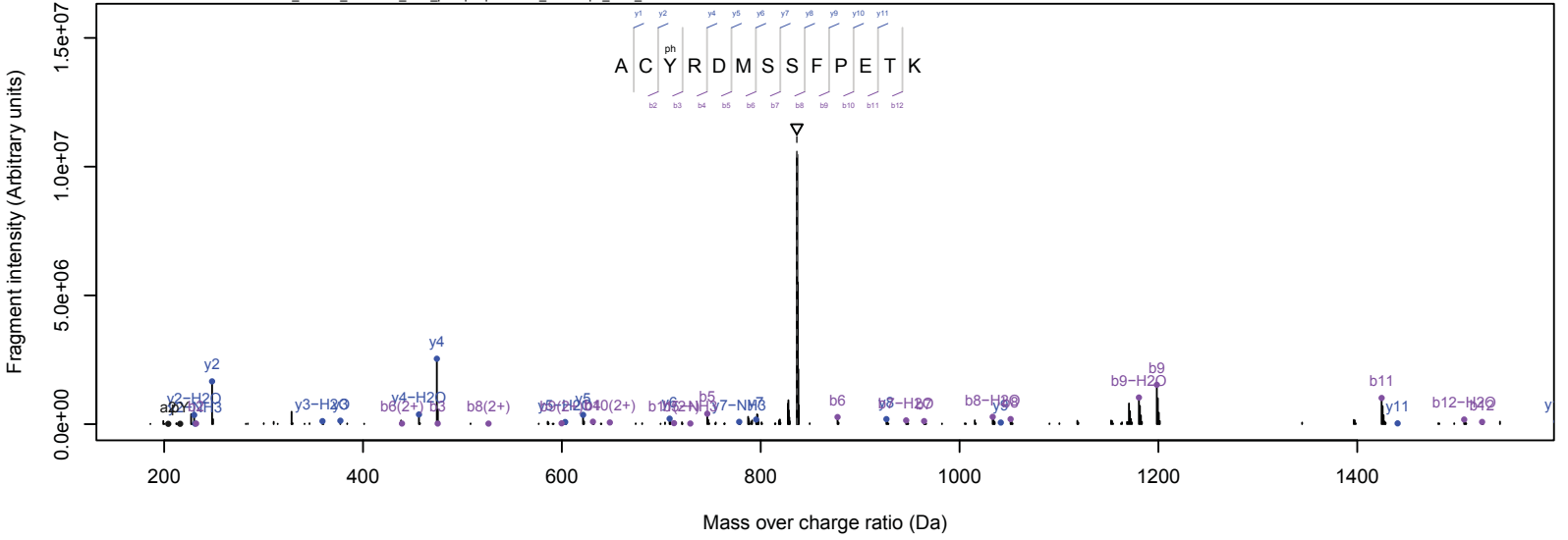
Andromeda score: 183.09



ACY(ph)RDMSSFPETK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 15210

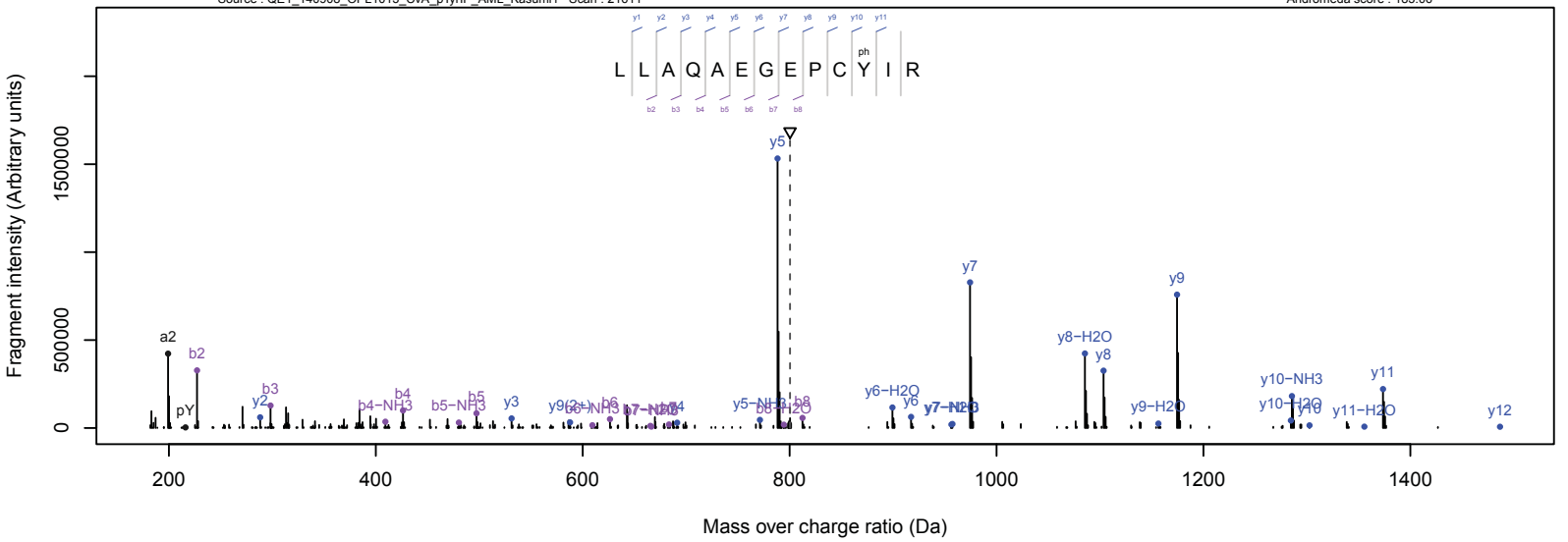
Andromeda score : 183.06



LLAQAEGPCY(ph)IR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 21611

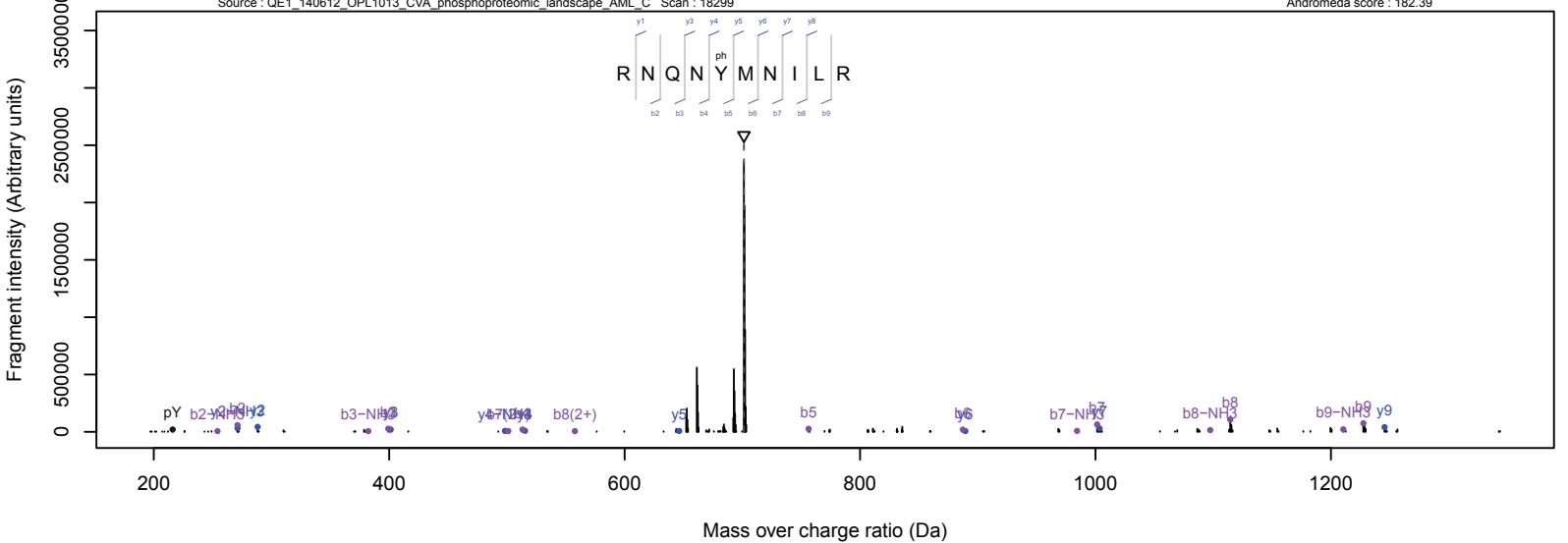
Andromeda score : 183.06



RNQNY(ph)MNILR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18299

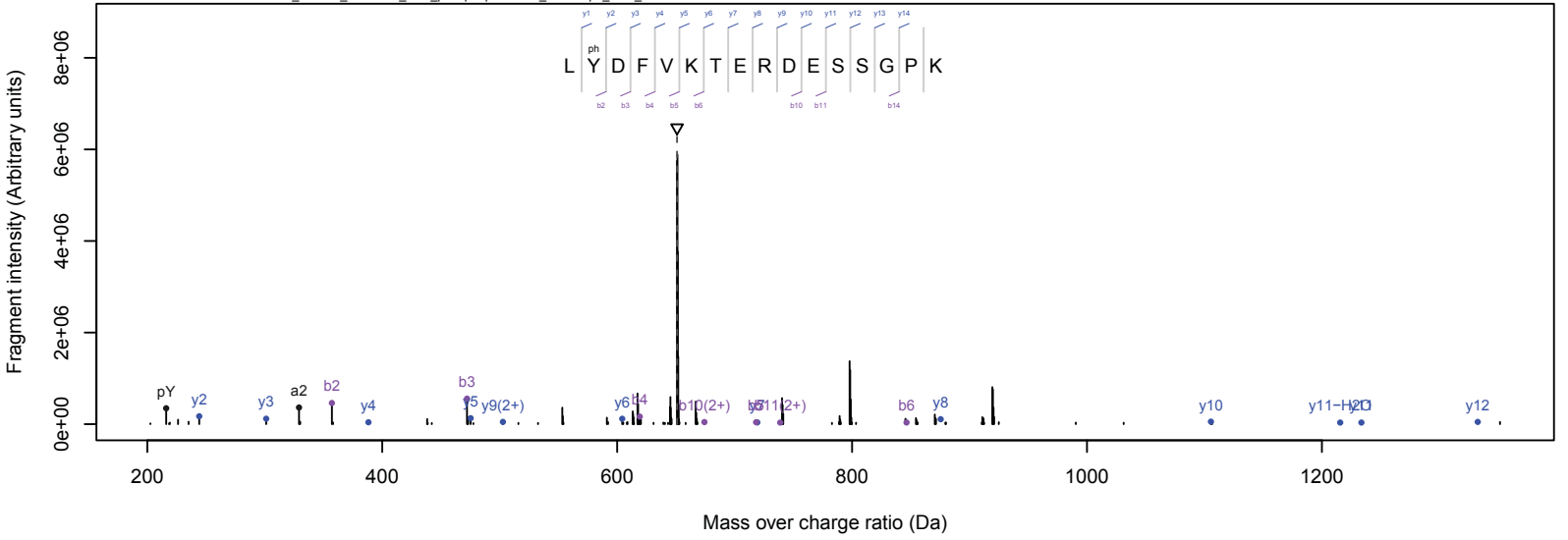
Andromeda score : 182.39



LY(ph)DFVKTERDESSGPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16449

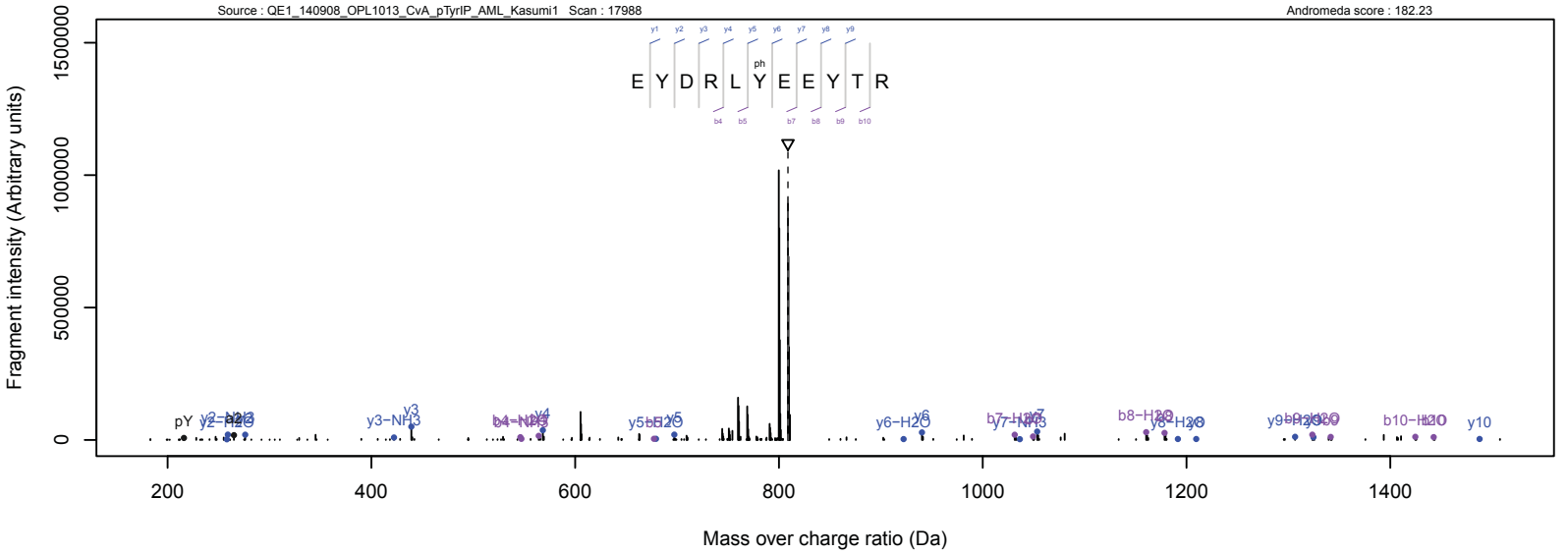
Andromeda score : 182.35



EYDRLY(ph)EEYTR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 17988

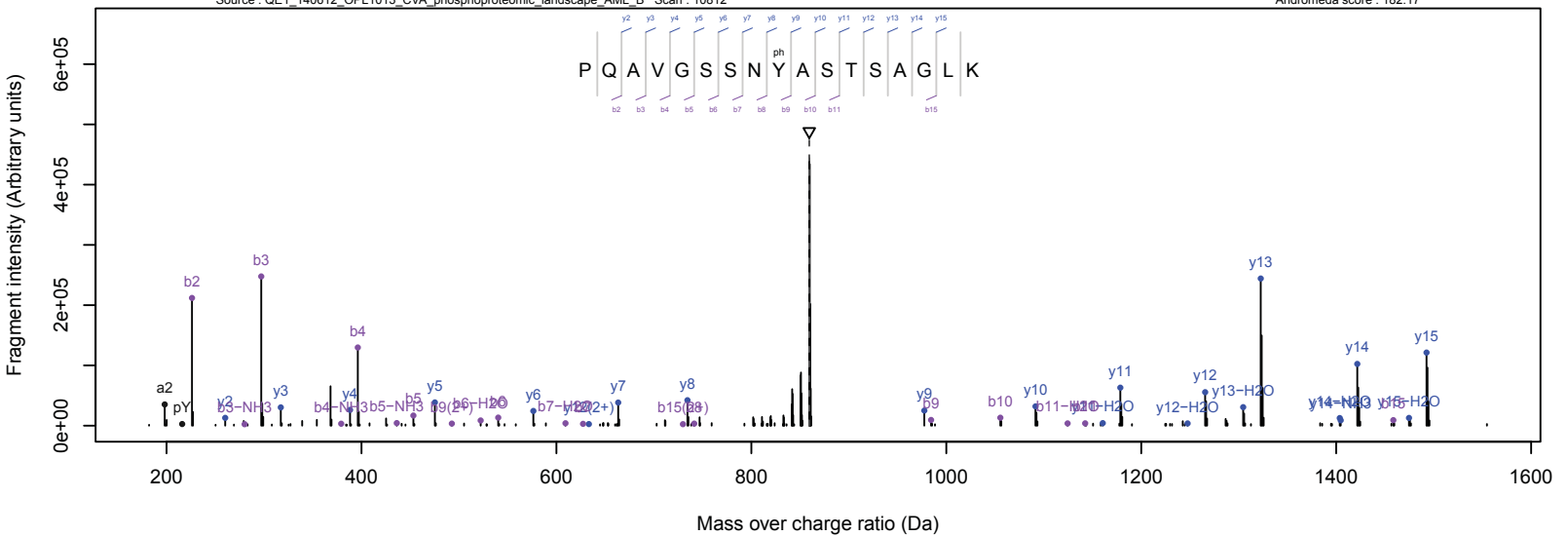
Andromeda score : 182.23



PQAVGSSNY(ph)ASTSAGLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10812

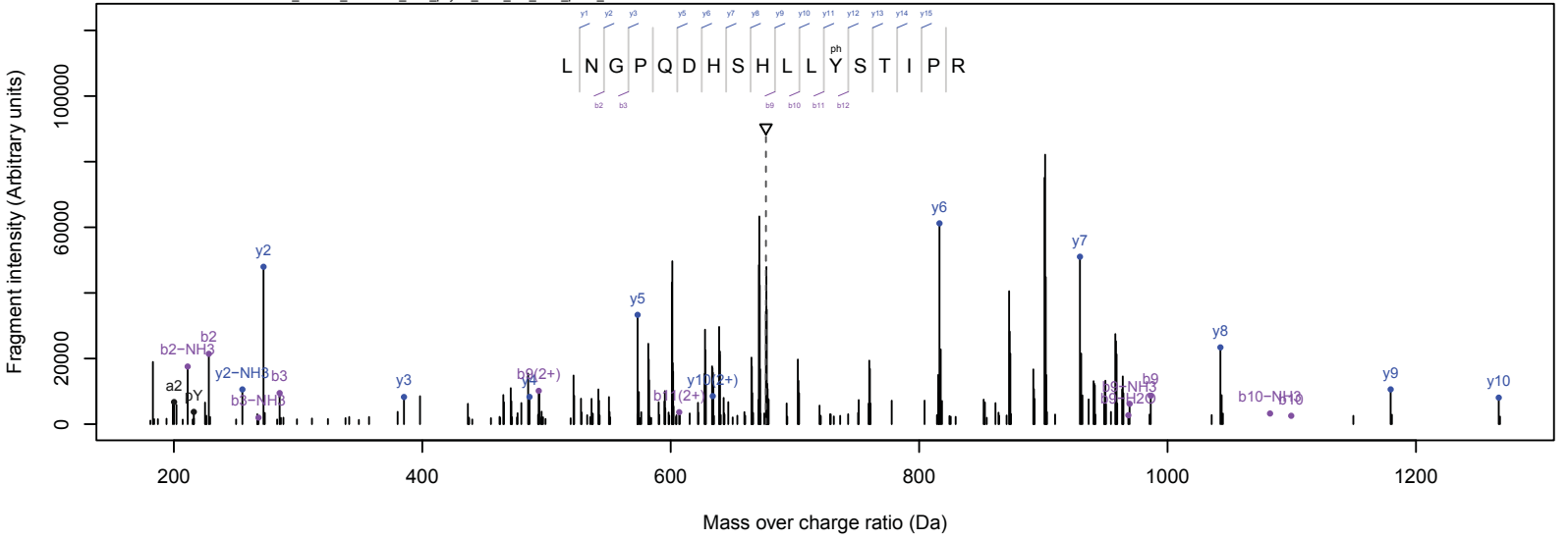
Andromeda score : 182.17



LNGPQDHSLLY(ph)STIPR-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 18729

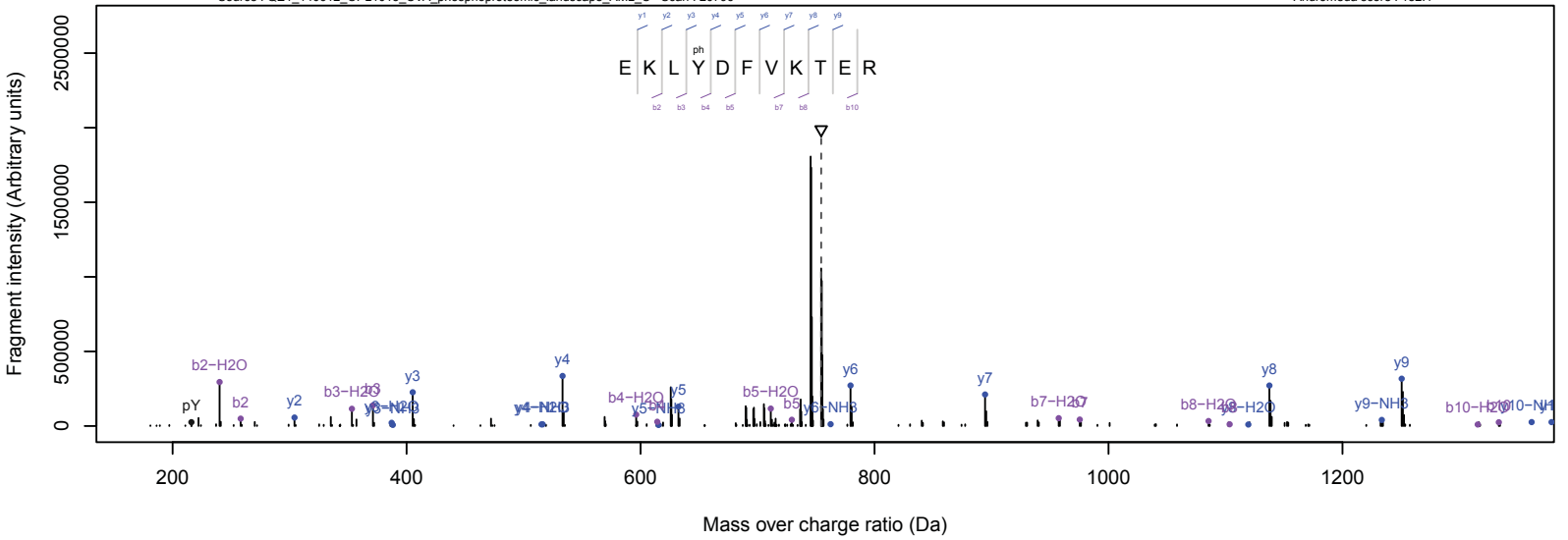
Andromeda score : 182.15



EKLY(ph)DFVKTER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20796

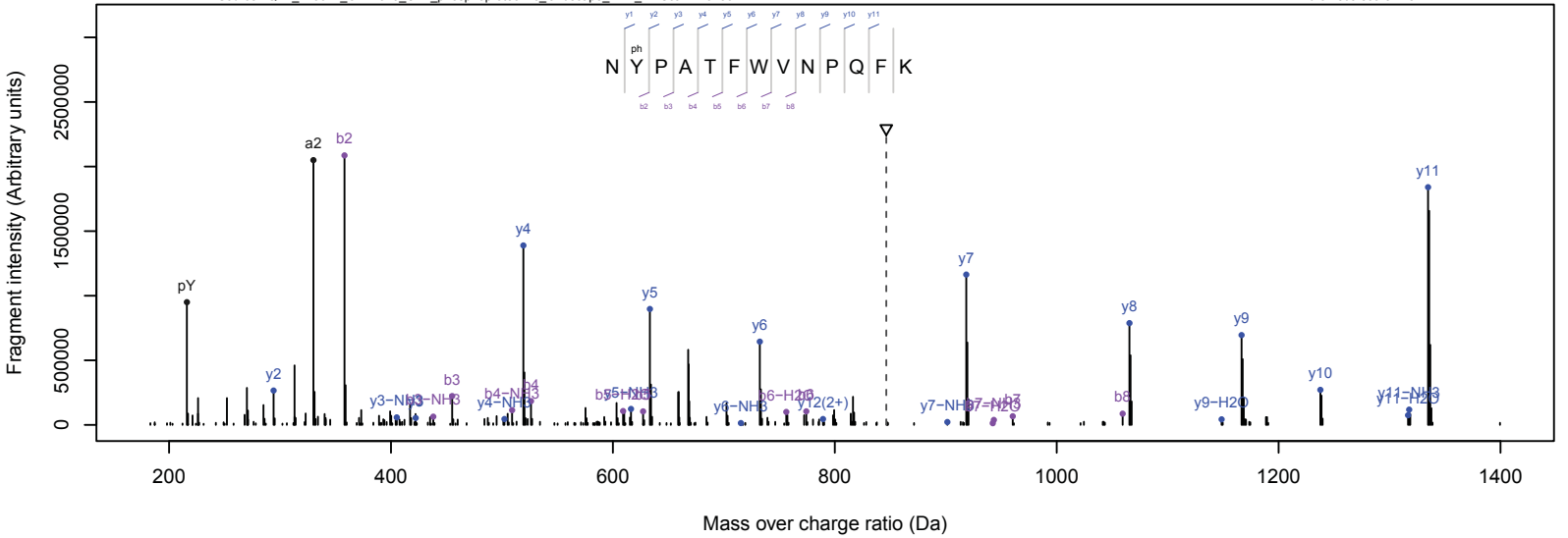
Andromeda score : 182.1



NY(ph)PATFWNPQFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40756

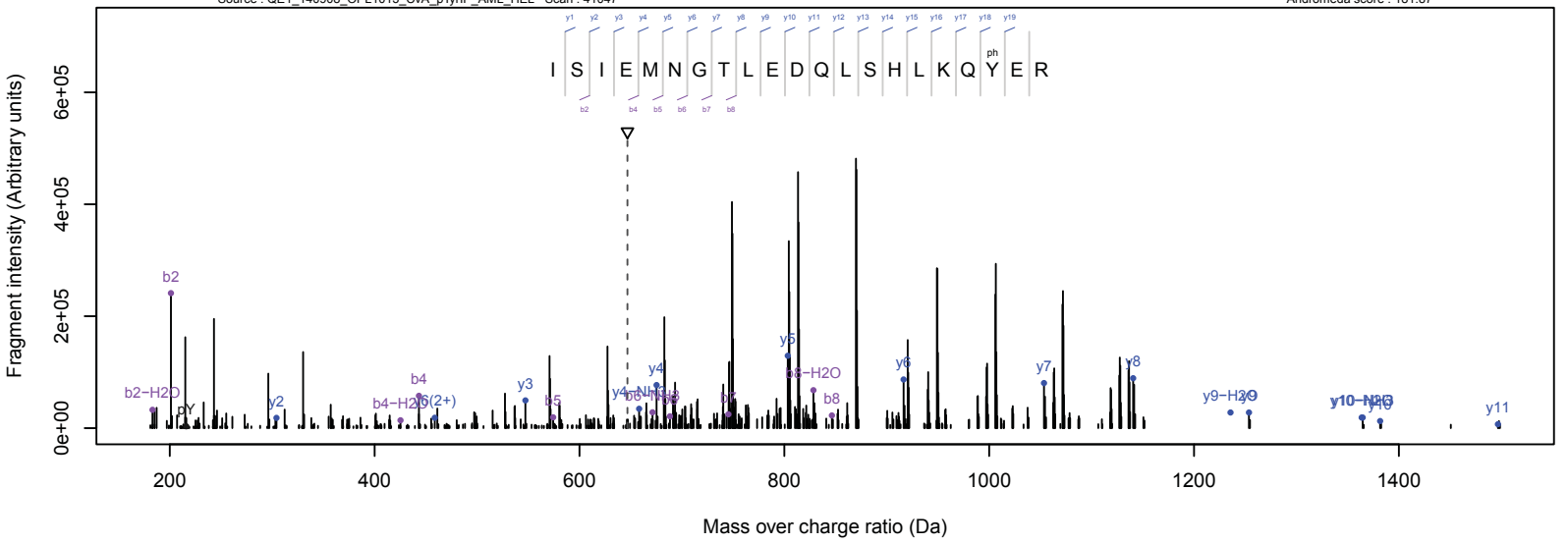
Andromeda score : 182.1



**ISIEMNGTLEDQLSHLKQY(ph)ER-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 41047

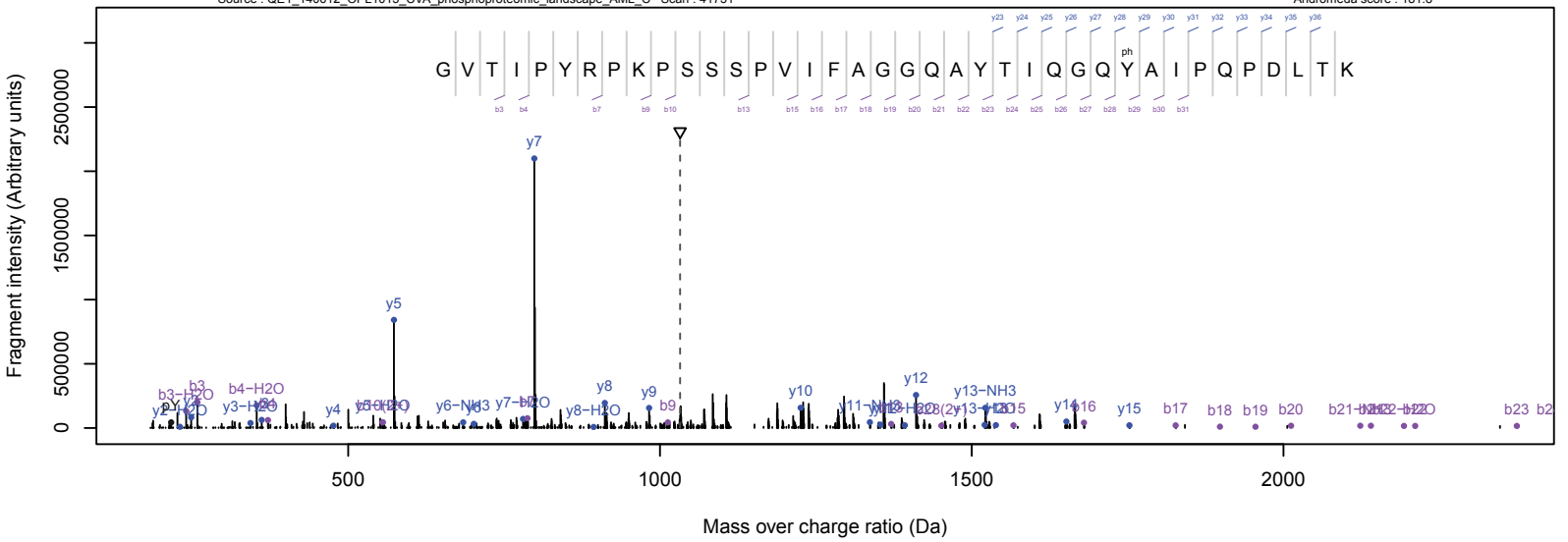
Andromeda score : 181.87



**GVTIPYRKPSSSPVIFAGGQAYTIQQQY(ph)AIPQPDLTk-4+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 41731

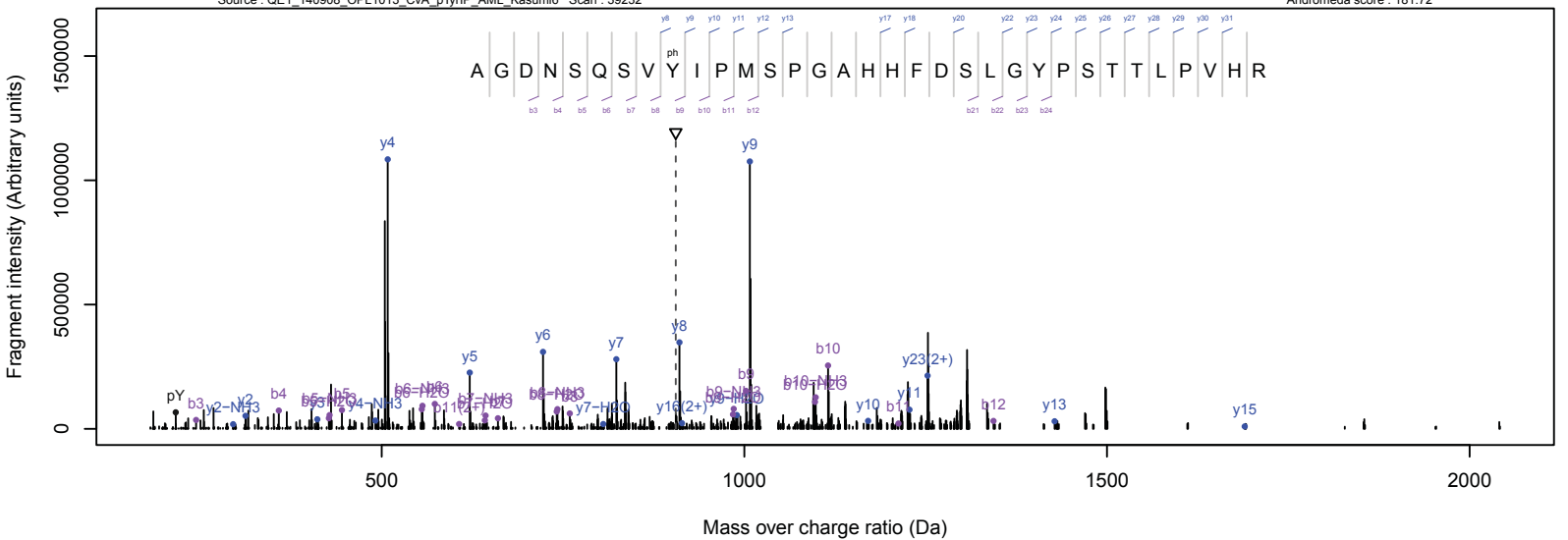
Andromeda score : 181.8



**AGDNSQSVY(ph)IPMSPGAHHFDSLGYPSTTLPVHR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 39232

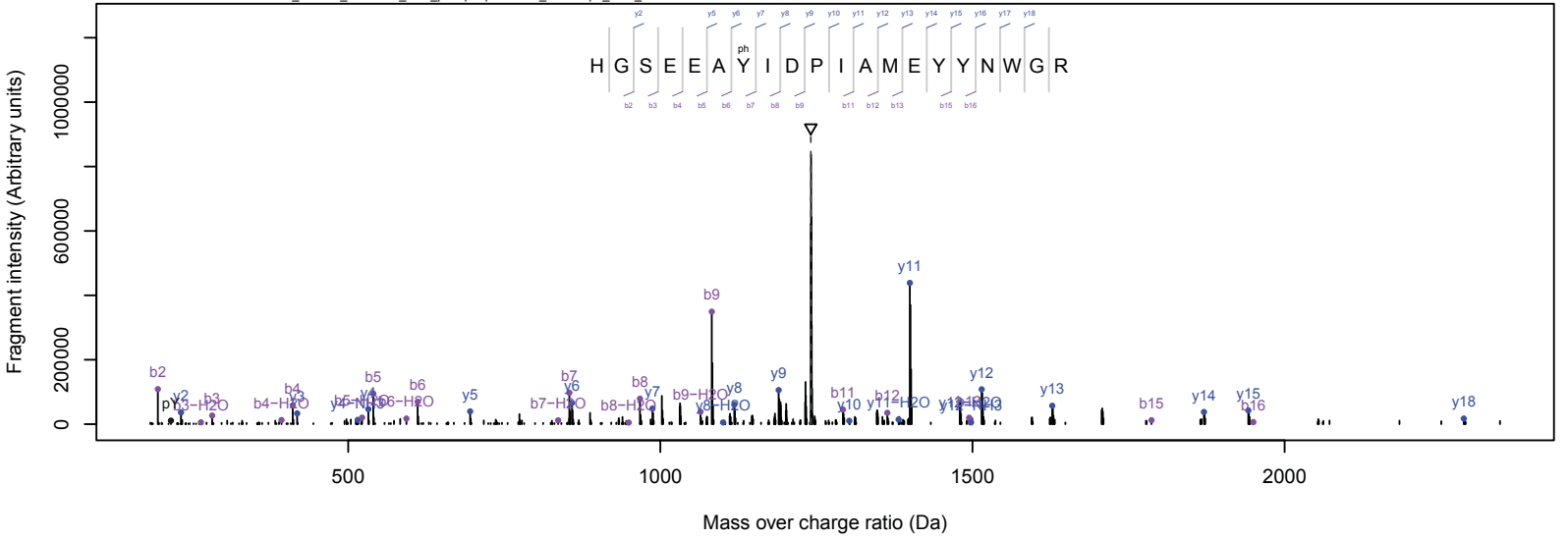
Andromeda score : 181.72



HGSEEA(ph)IDPIAMEYYNWGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44576

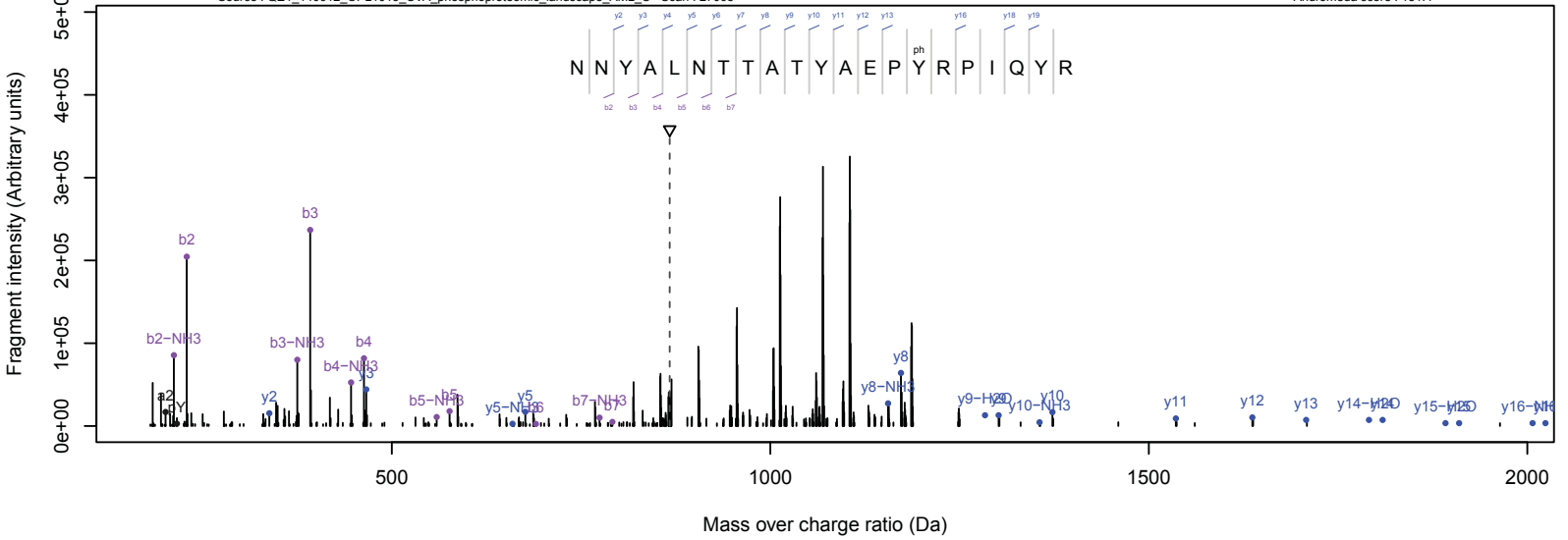
Andromeda score : 181.4



NNYALNTTATYAEPY(ph)RPIQYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27068

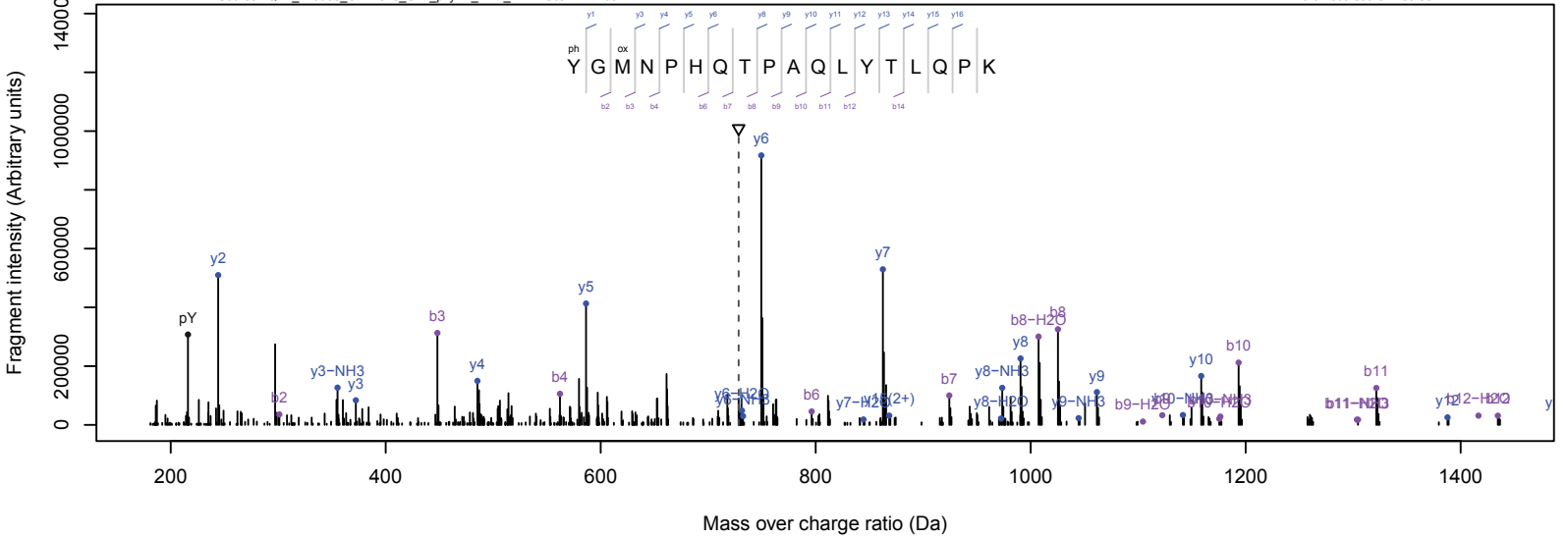
Andromeda score : 181.4



Y(ph)GM(ox)NPHQTPAQLYTLQPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24403

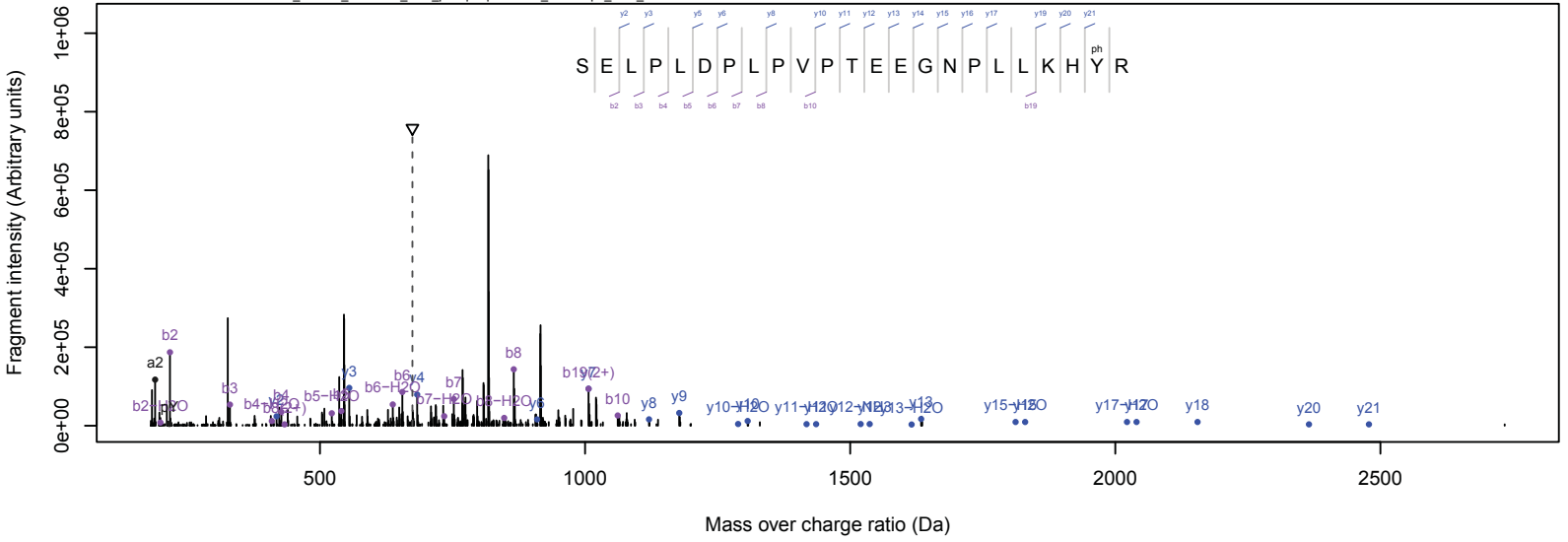
Andromeda score : 180.88



**SELPLDPLVPTEEGNPLLKHYPH-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 43753

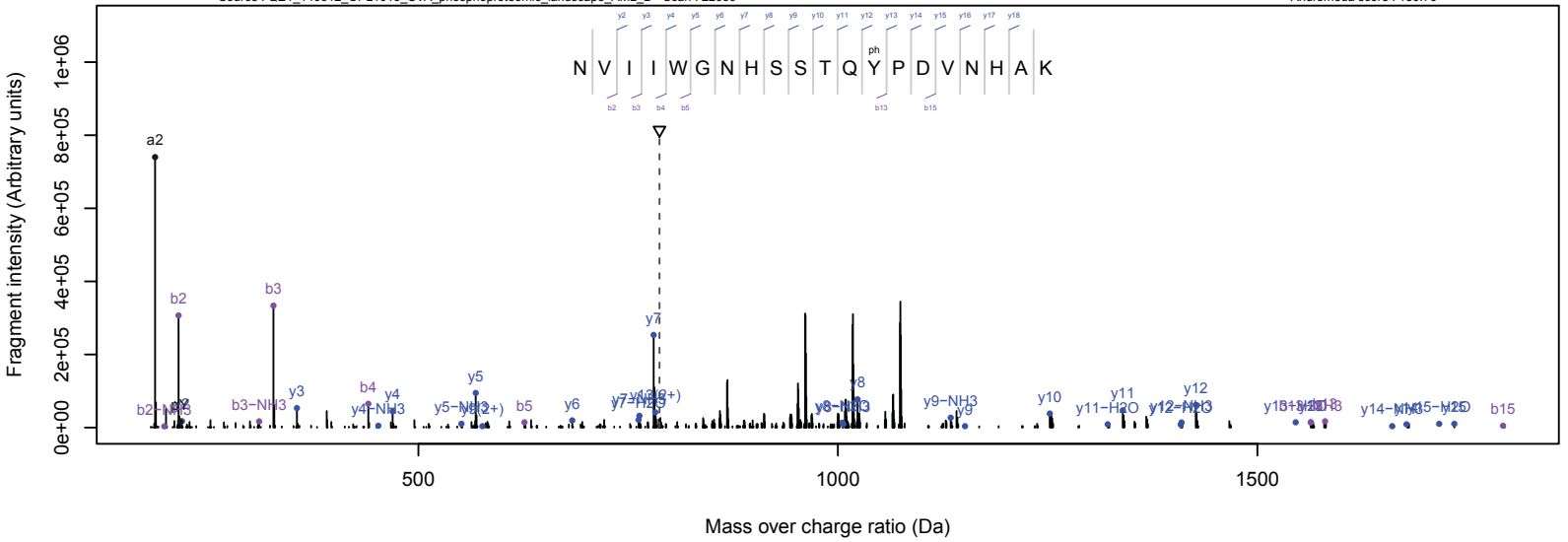
Andromeda score: 180.86



**NVIIWGNHSSSTQYHPD VN HAK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 22659

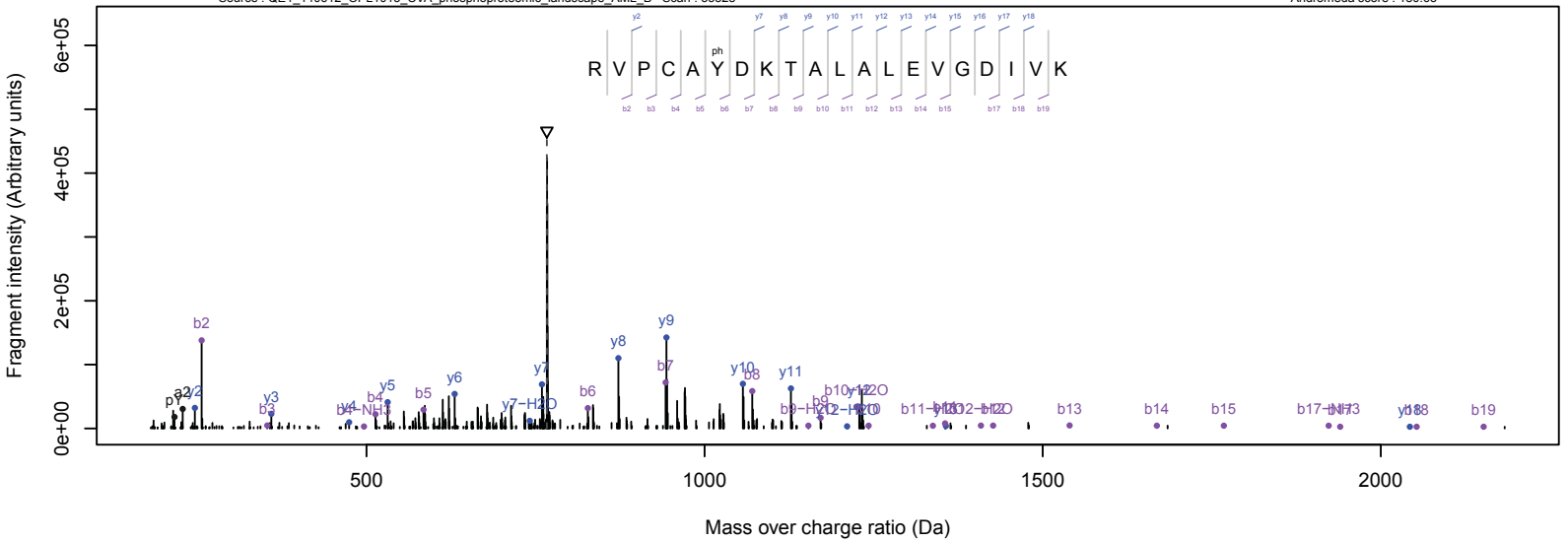
Andromeda score: 180.76



**RVPCAYDKTALALEVGDIVK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 35528

Andromeda score: 180.63

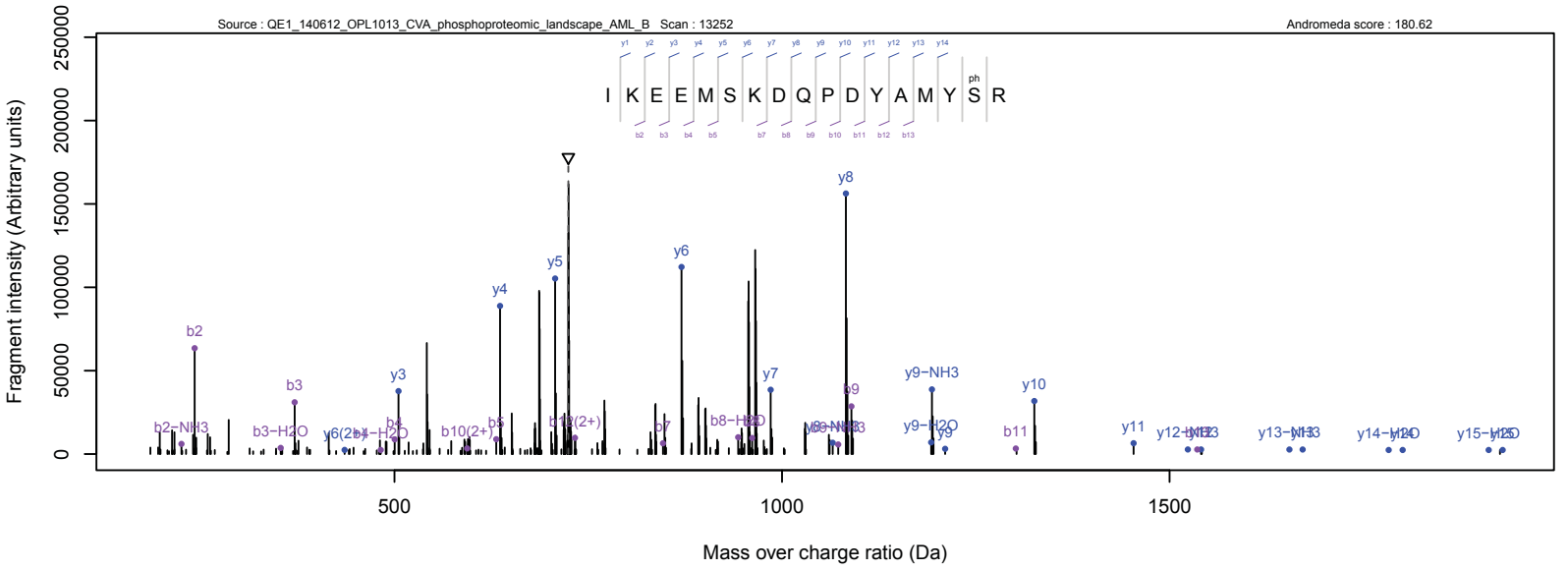




IKEEMSKDPDYAMYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13252

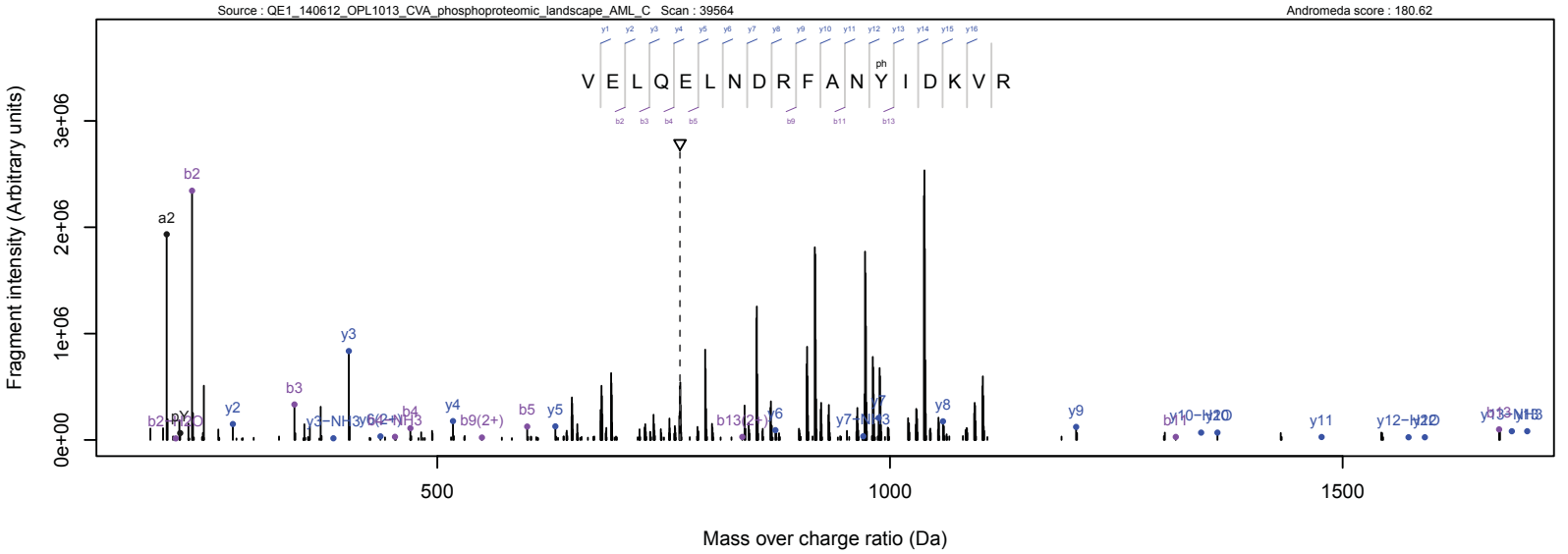
Andromeda score : 180.62



VELQELNDRFANYIDKVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39564

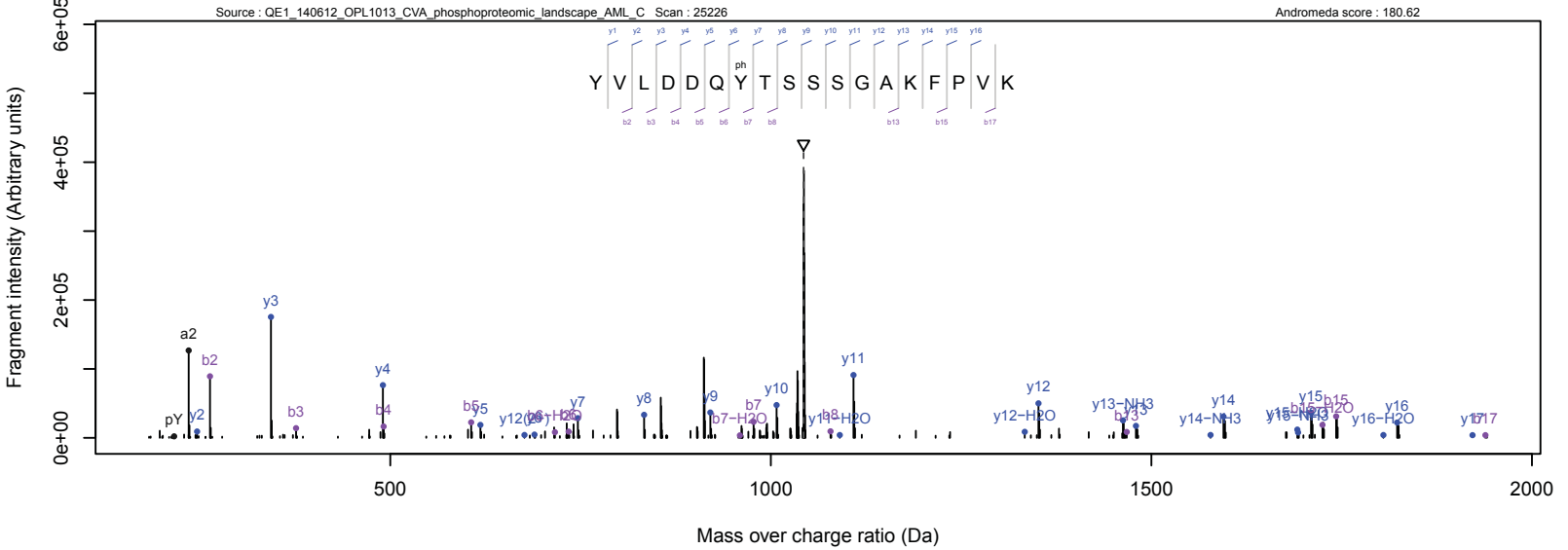
Andromeda score : 180.62



YVLDQY(ph)TSSGAKFPVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25226

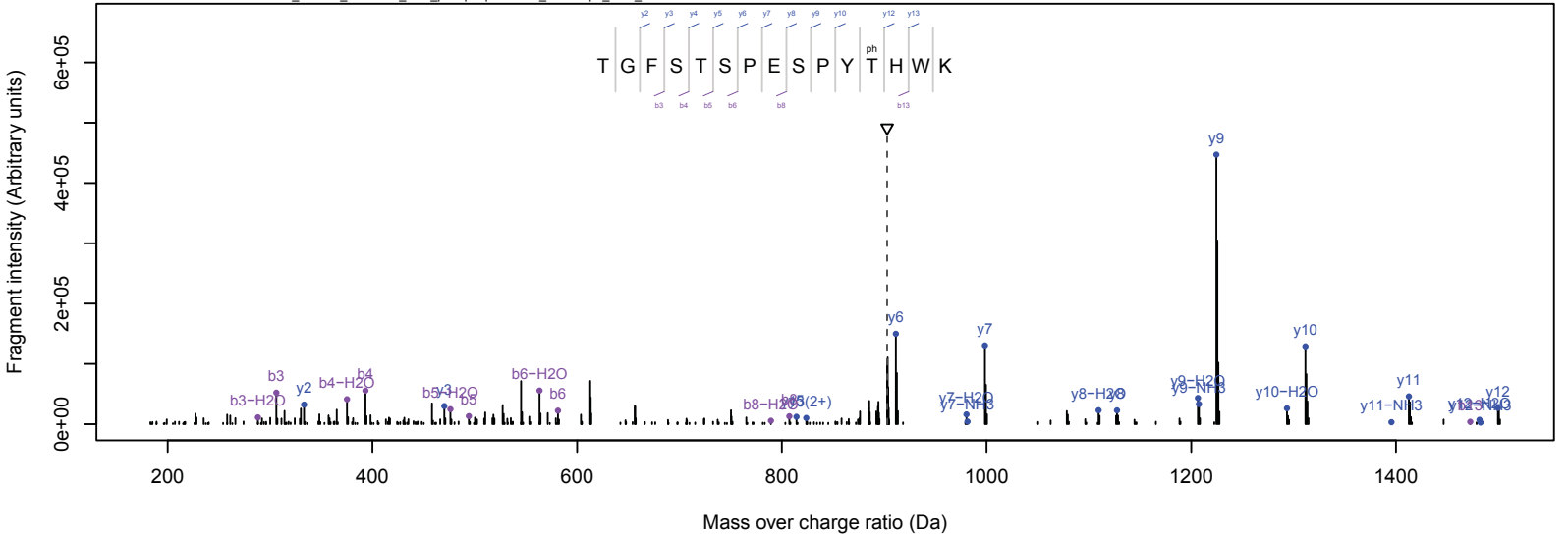
Andromeda score : 180.62



TGFSTSPESPYT<sup>ph</sup>HWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20613

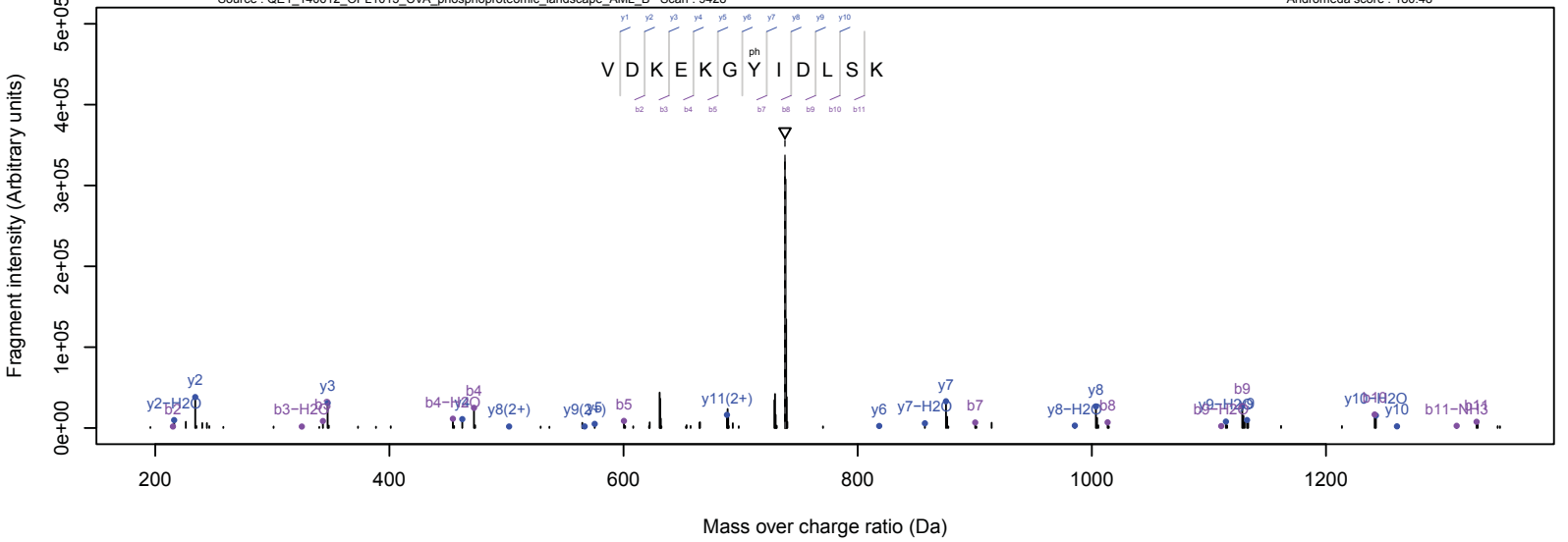
Andromeda score : 180.48



VDKEKGY<sup>ph</sup>IDLSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9428

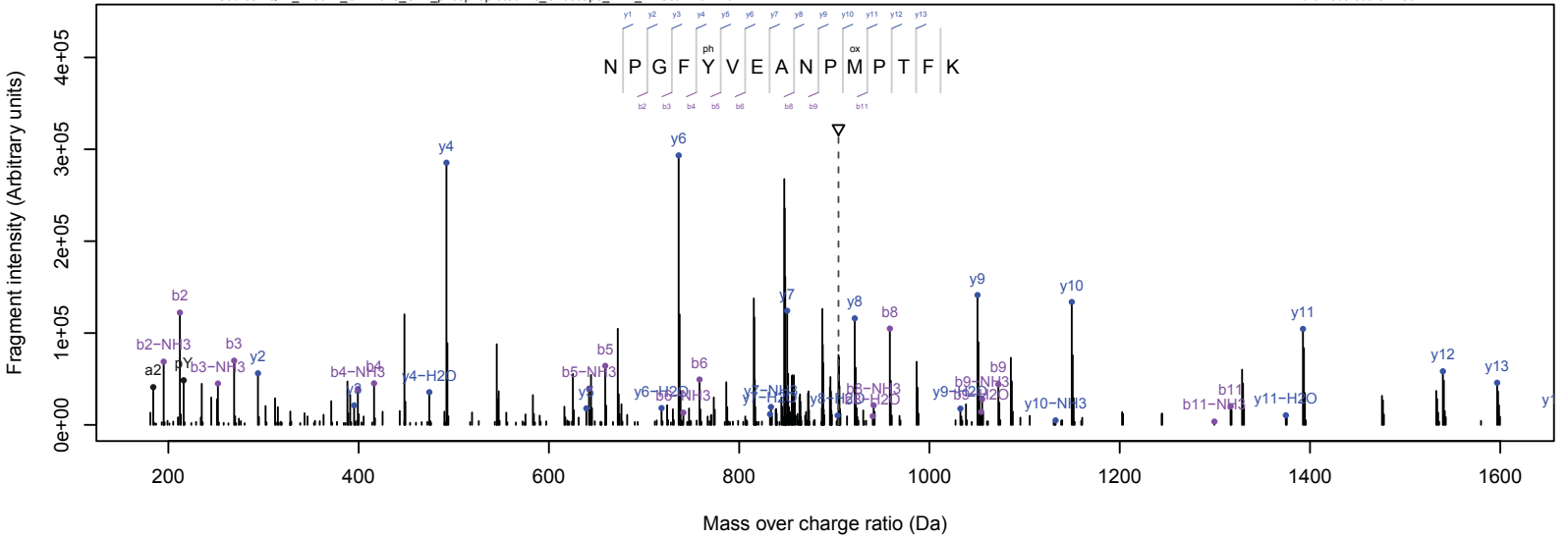
Andromeda score : 180.48



NPGFY<sup>ph</sup>VEANP<sup>ox</sup>MPTFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31176

Andromeda score : 180.11

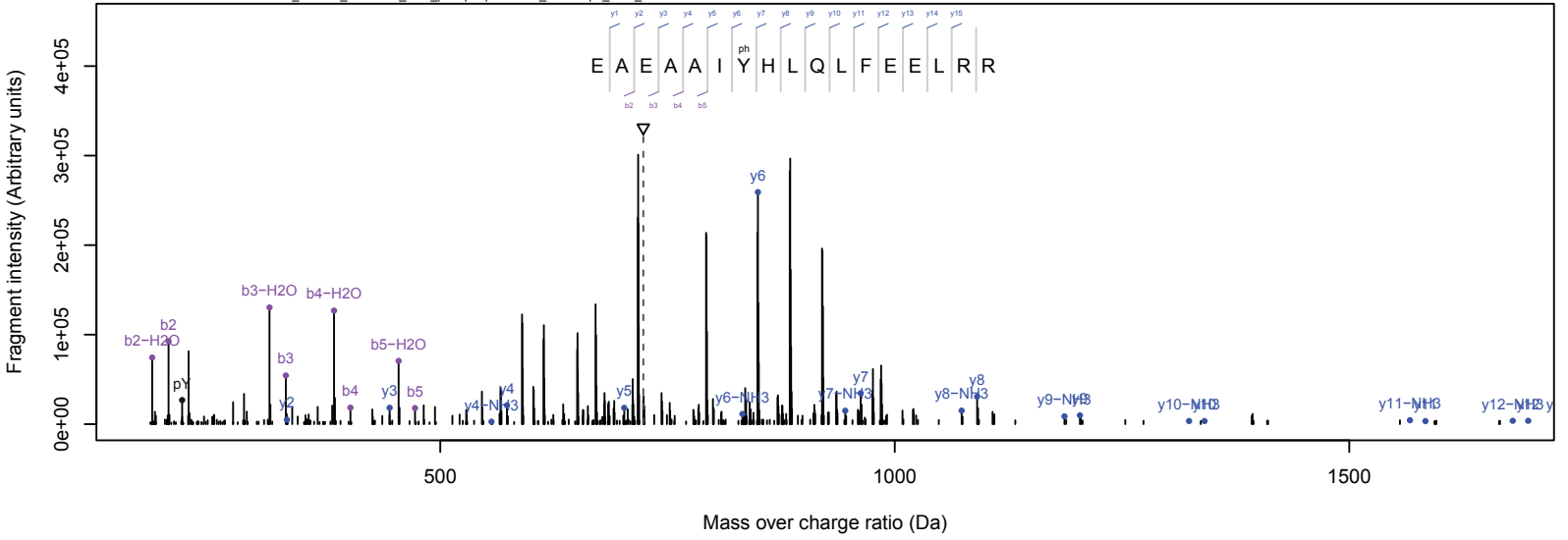




**EAEAAIYHLQFEELRR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41894

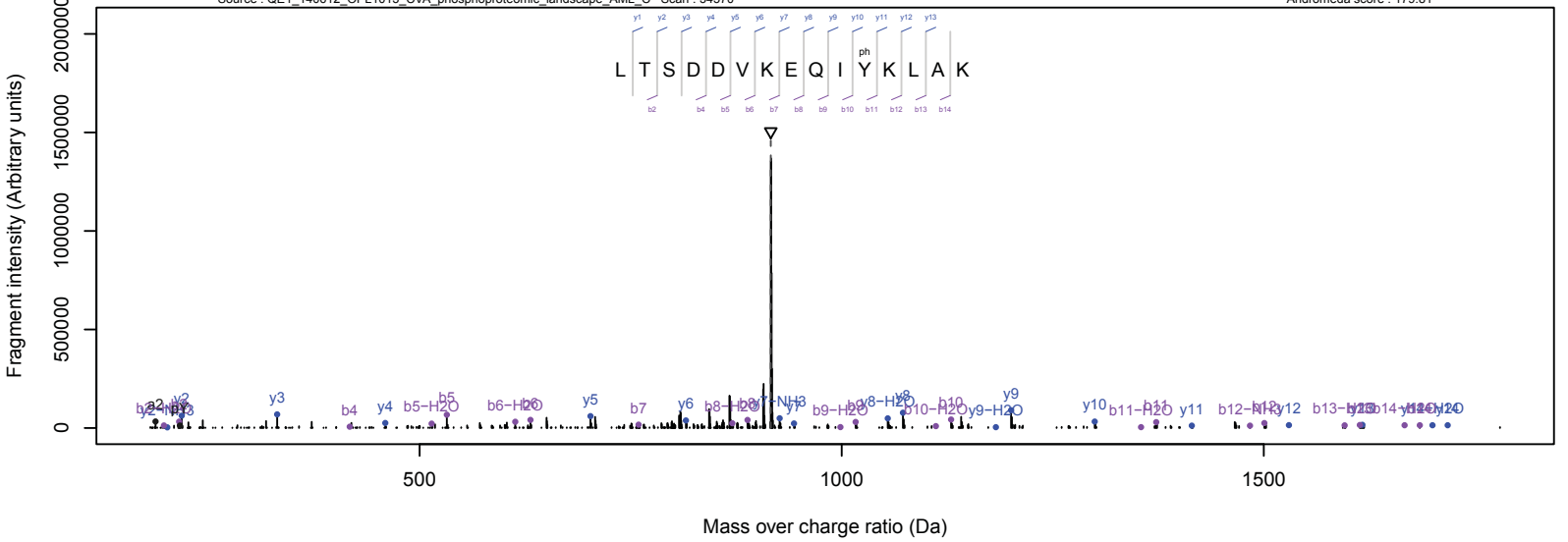
Andromeda score : 179.81



**LTSDDVKEIQIKLAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34570

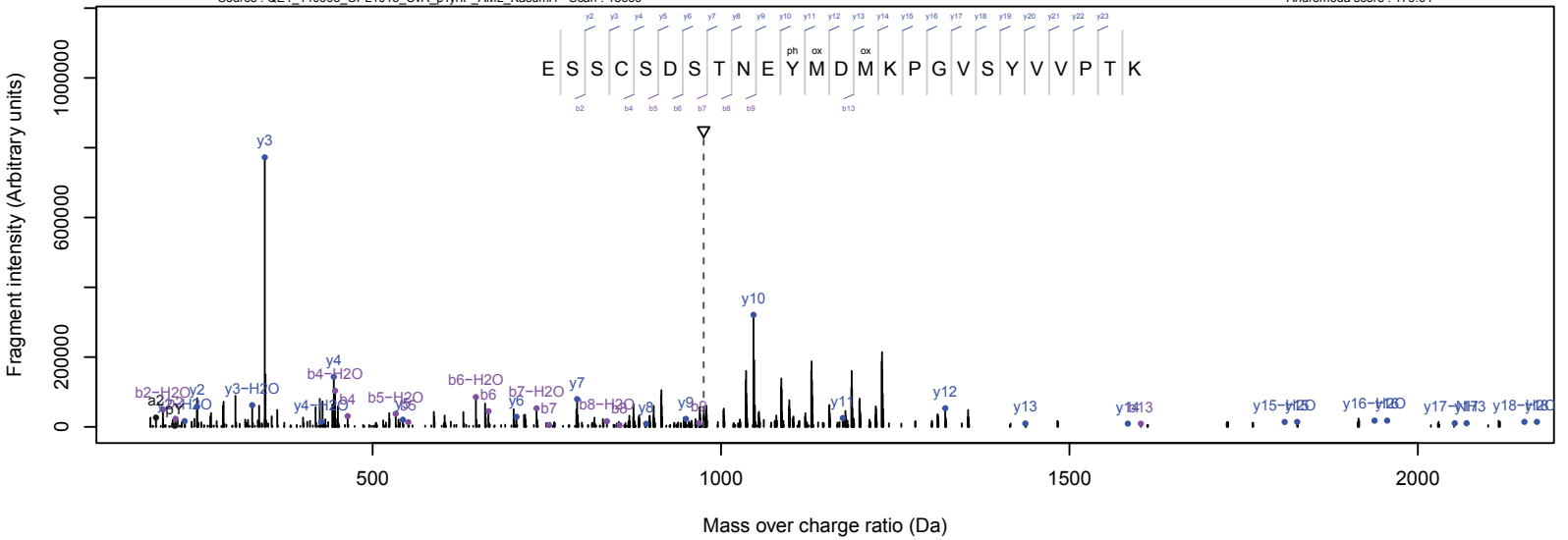
Andromeda score : 179.81



**ESSCSDSTNEYM(ox)DMKPGVSYVVPVK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 18659

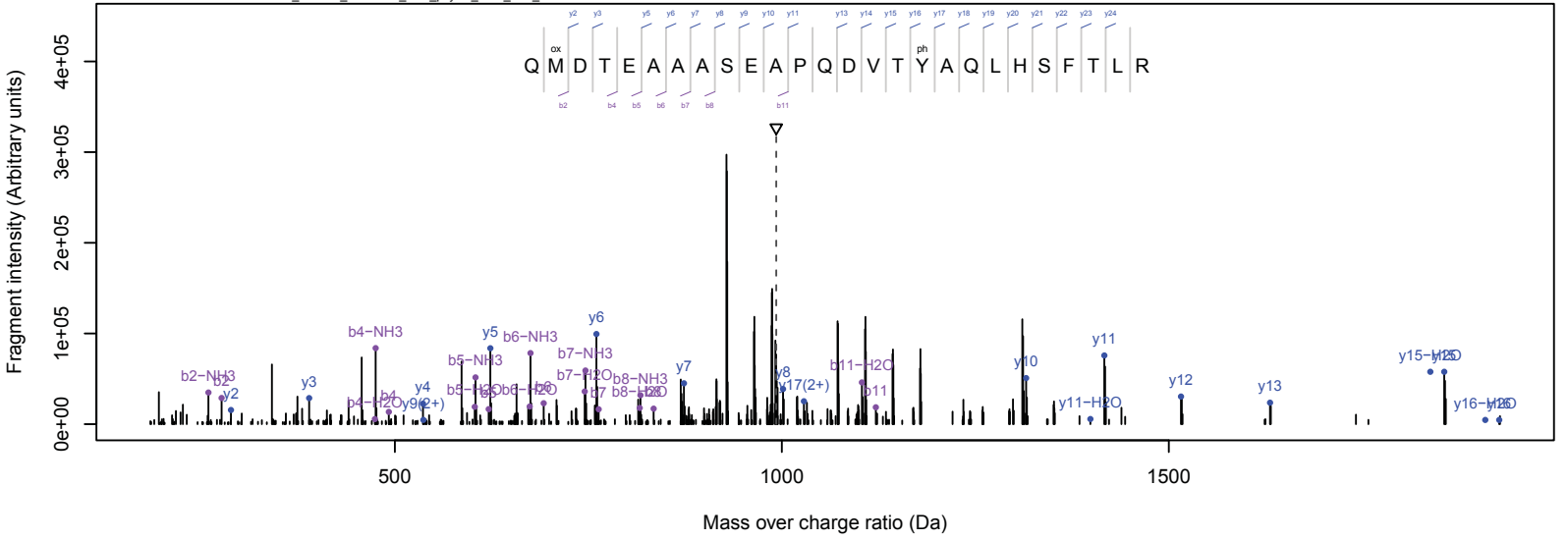
Andromeda score : 179.64



QM(ox)DTEAAASEAPQDVTY(ph)AQLHSFTLR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_OCI\_AML3 Scan : 39133

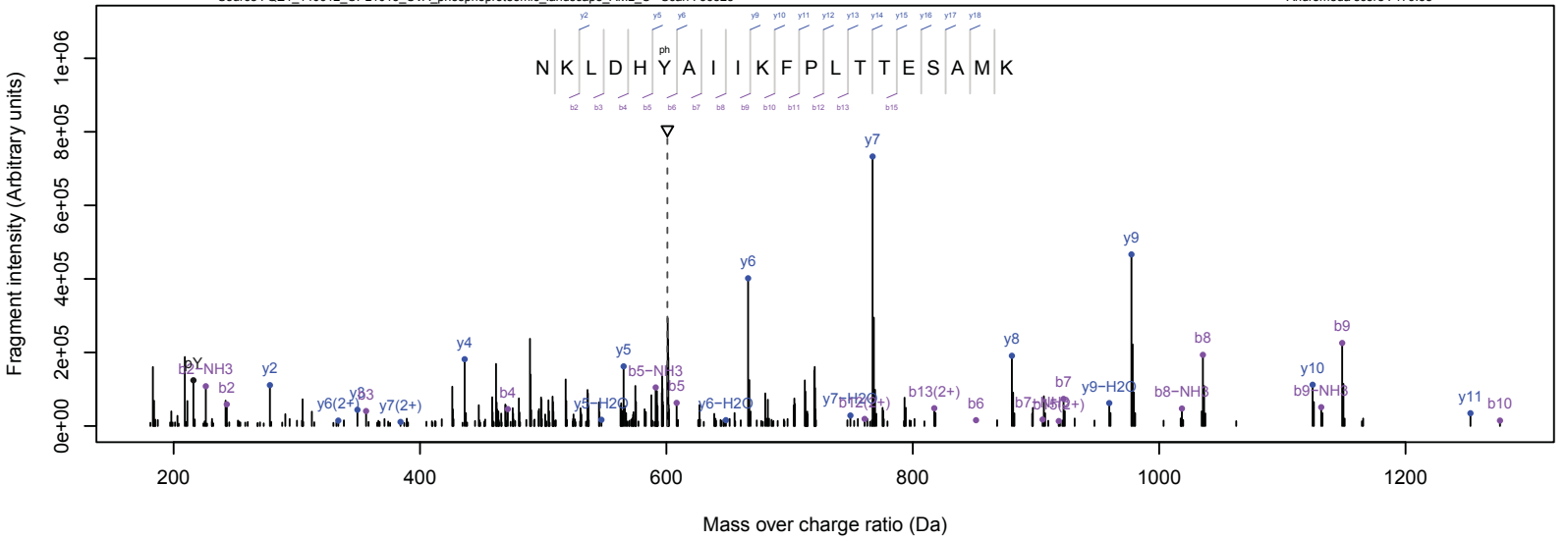
Andromeda score : 179.64



NKLDHY(ph)AIKFLTTESAMK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36525

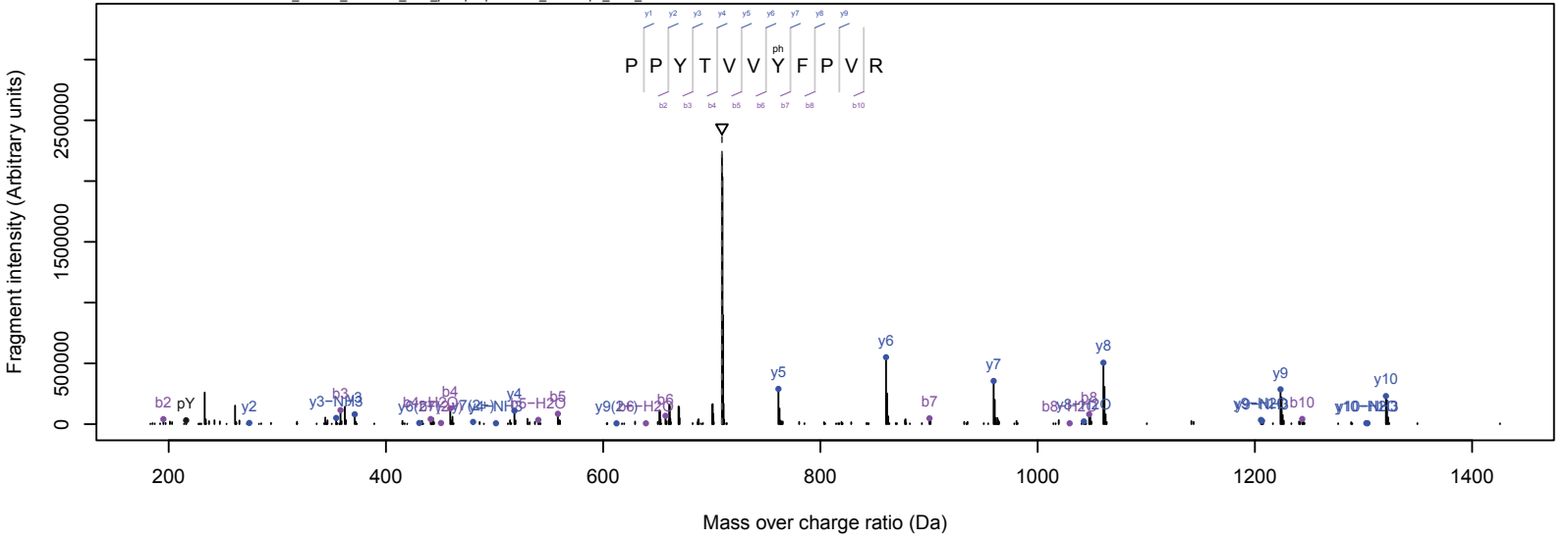
Andromeda score : 179.58



PPYTVVY(ph)FPVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40104

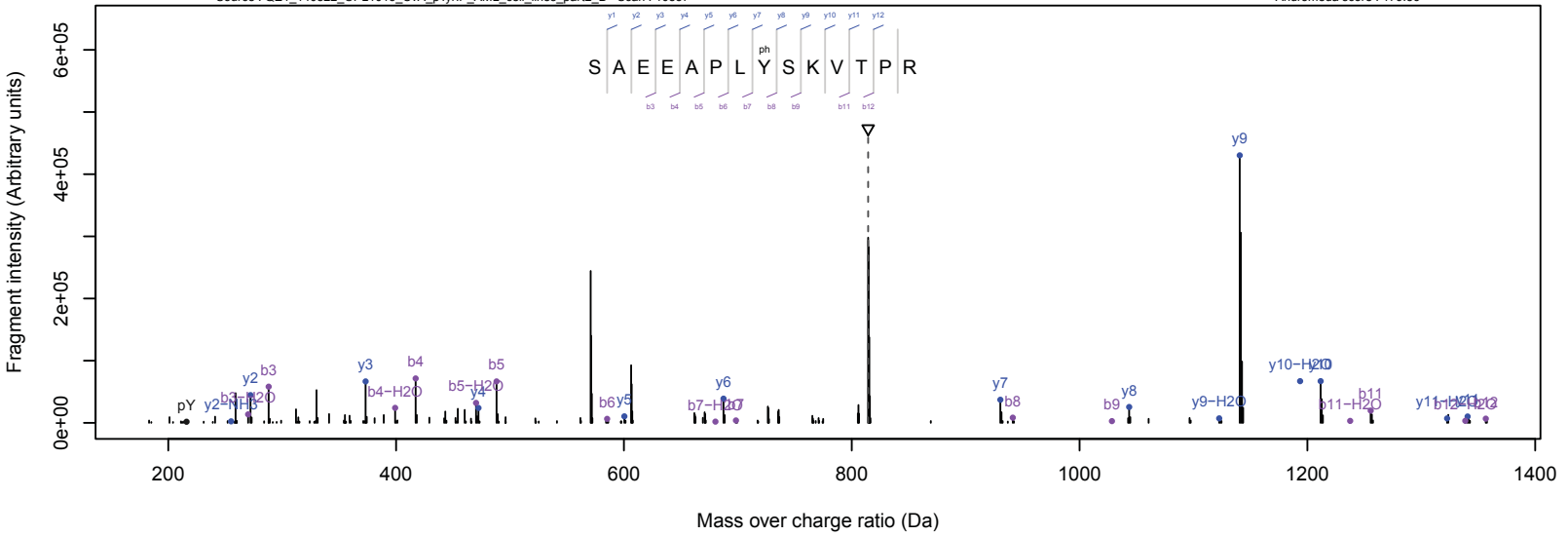
Andromeda score : 179.57



SAEEAPLY(ph)SKVTPR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 10857

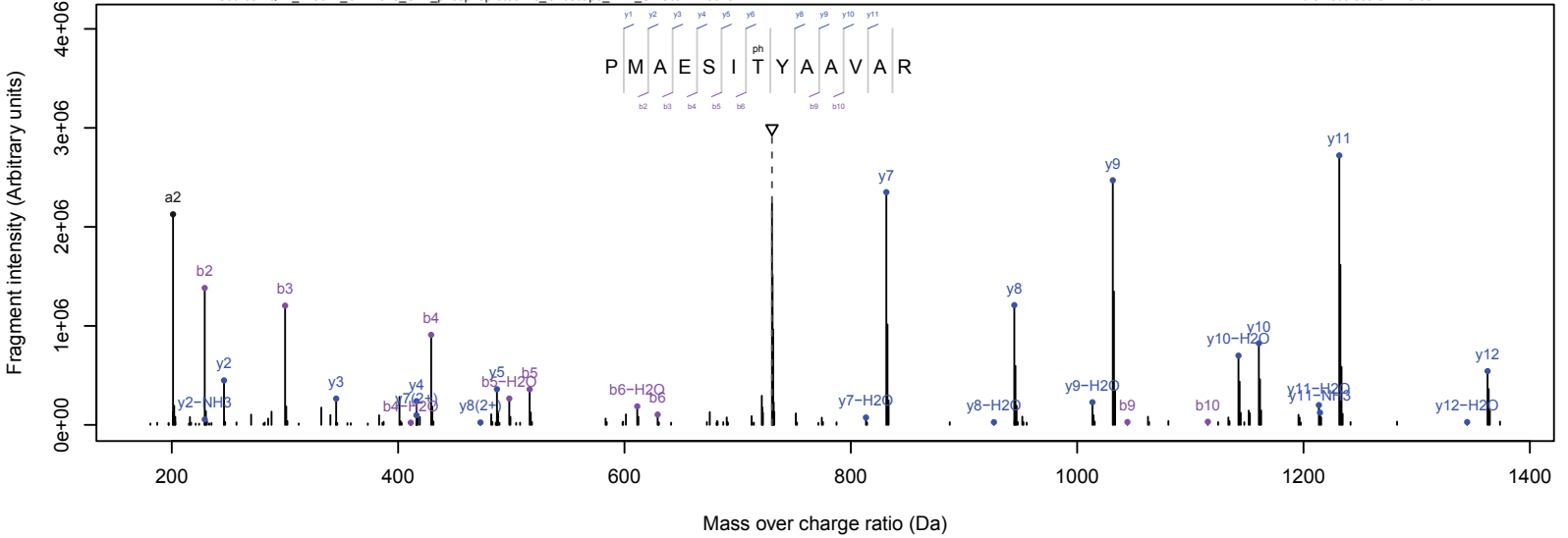
Andromeda score : 179.56



PMAESIT(ph)YAAVAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25028

Andromeda score : 179.35

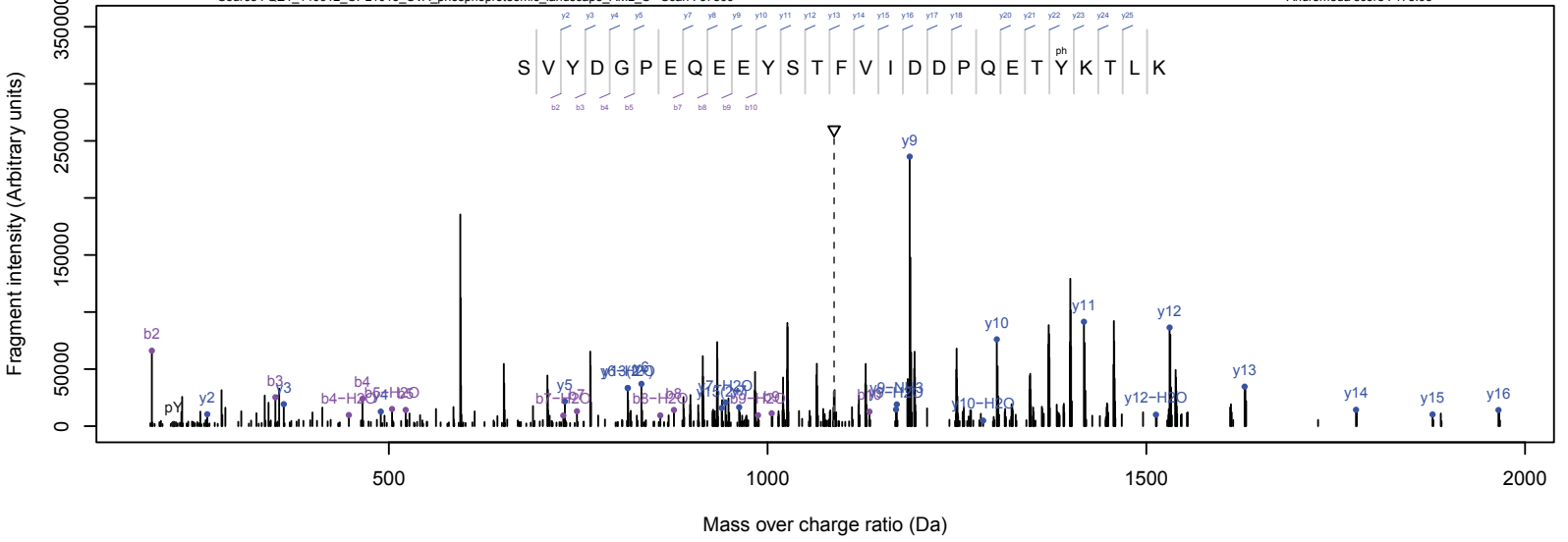




SVYDGP<sup>ph</sup>EQE<sup>ph</sup>EYSTFVIDDPQETYKTLK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37889

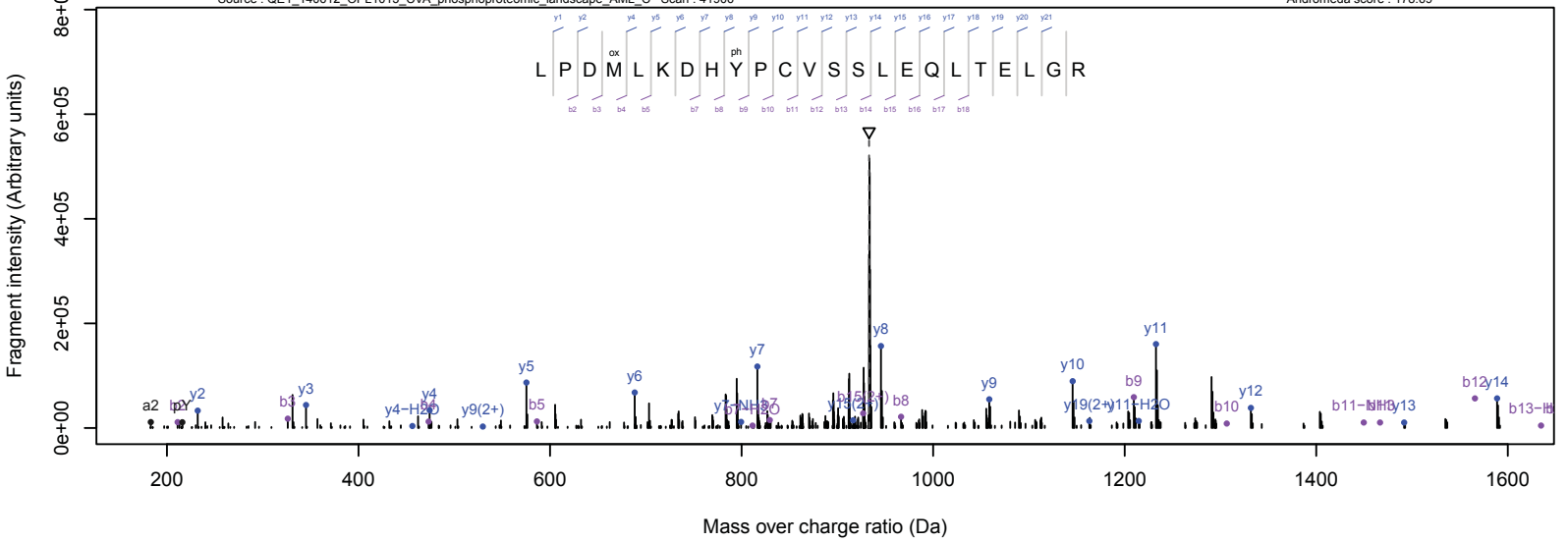
Andromeda score: 178.95



LPDM<sup>ox</sup>LKDHY<sup>ph</sup>PCVSSLEQLTELGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41906

Andromeda score: 178.69

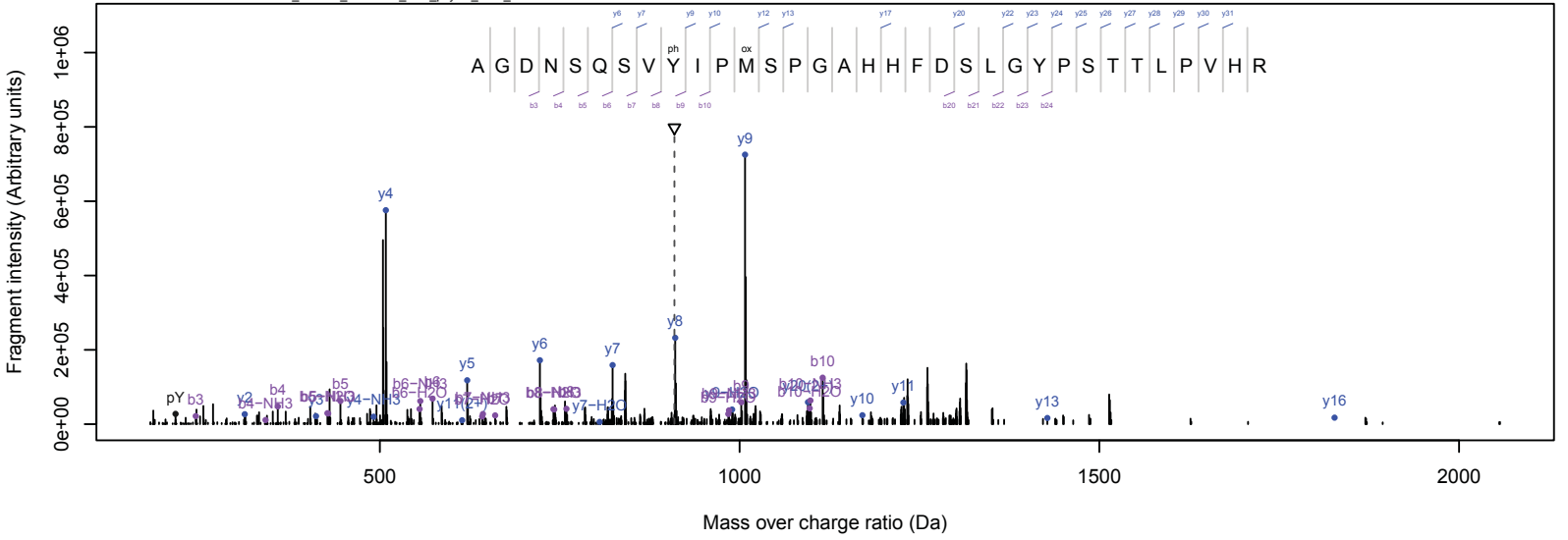




AGDNSQSVY(ph)IPMSPGAHHFDSLGYPSTTLPVHR-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 32114

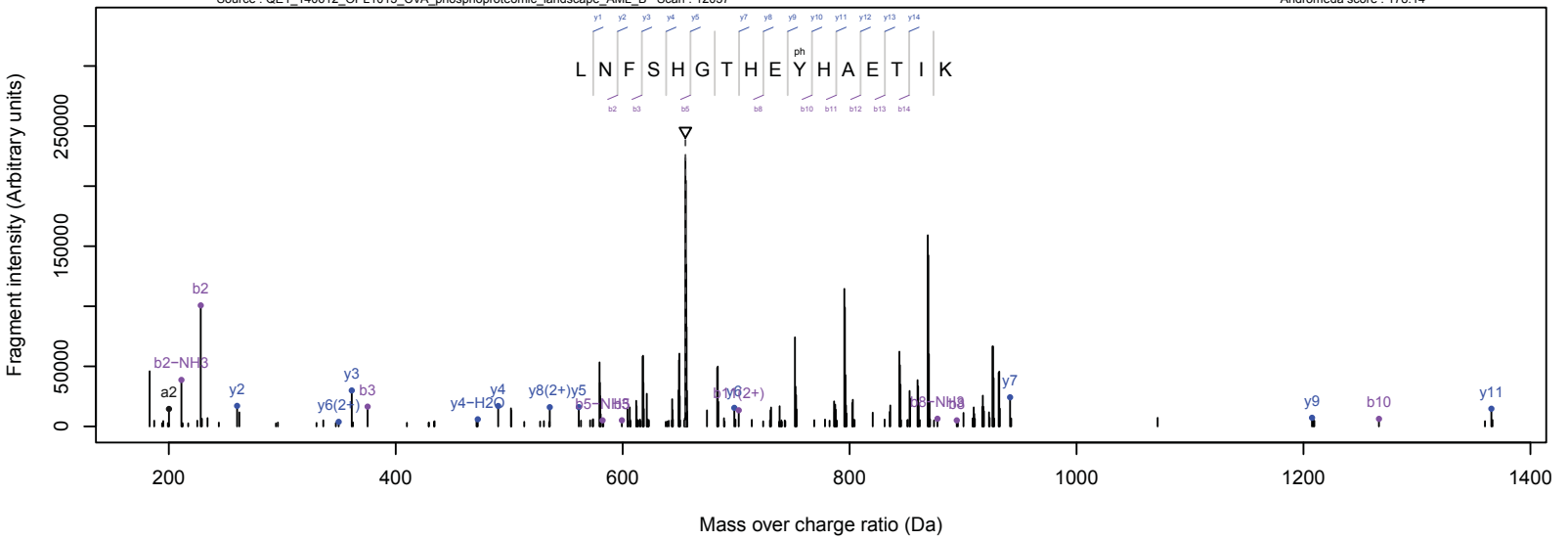
Andromeda score: 178.44



LNFSHGTHEY(ph)HAETIK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 12037

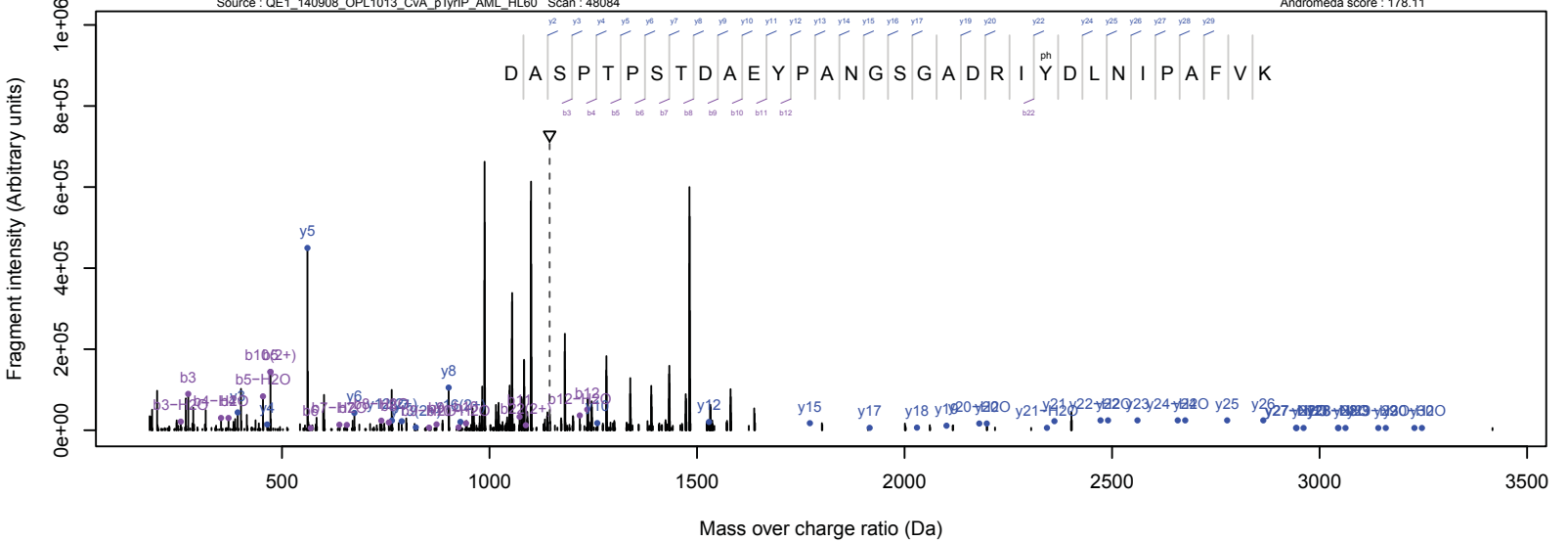
Andromeda score: 178.14



DASPTPSTDAEYPANGSGADRIY(ph)DLNIPAFVK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan: 48084

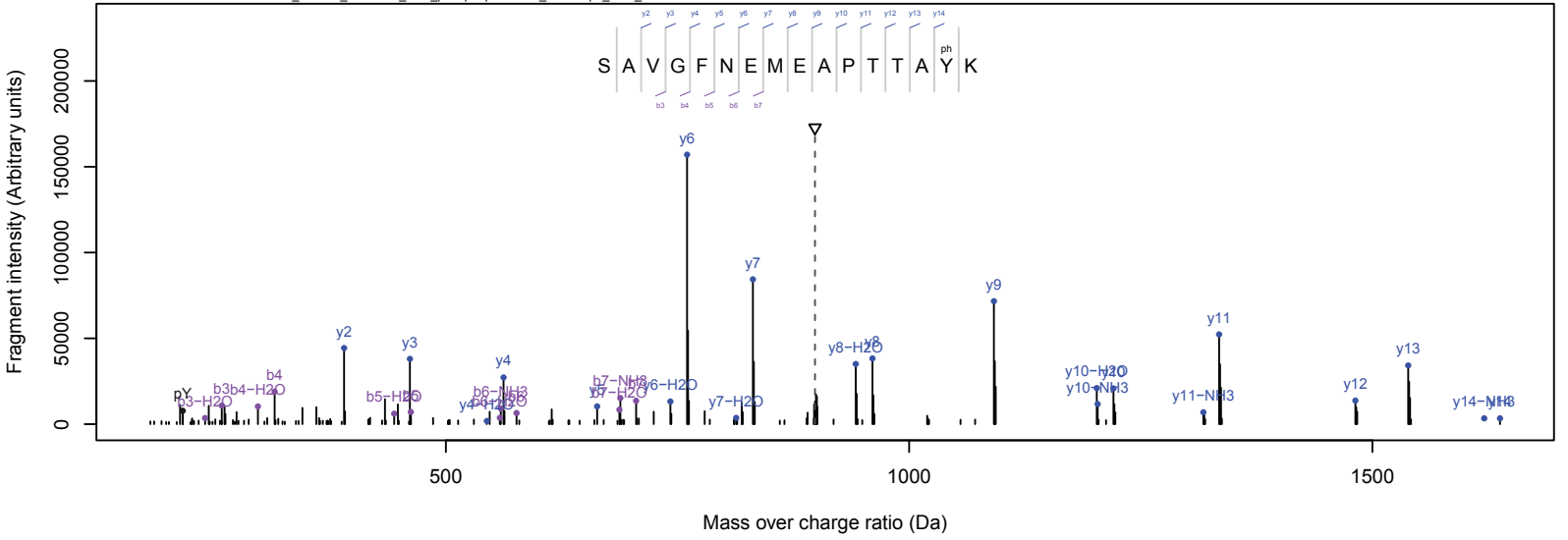
Andromeda score: 178.11



SAVGFNEMEAPTAYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26465

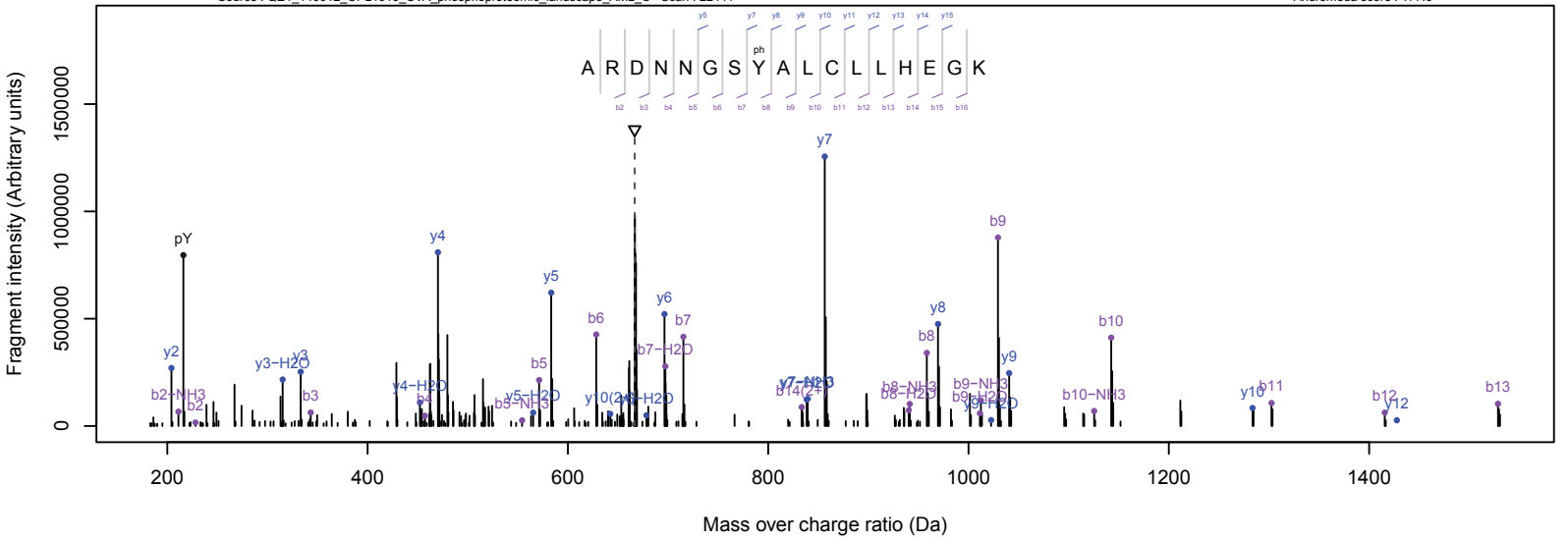
Andromeda score : 178.03



ARDNNGSYALCLLHEGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22441

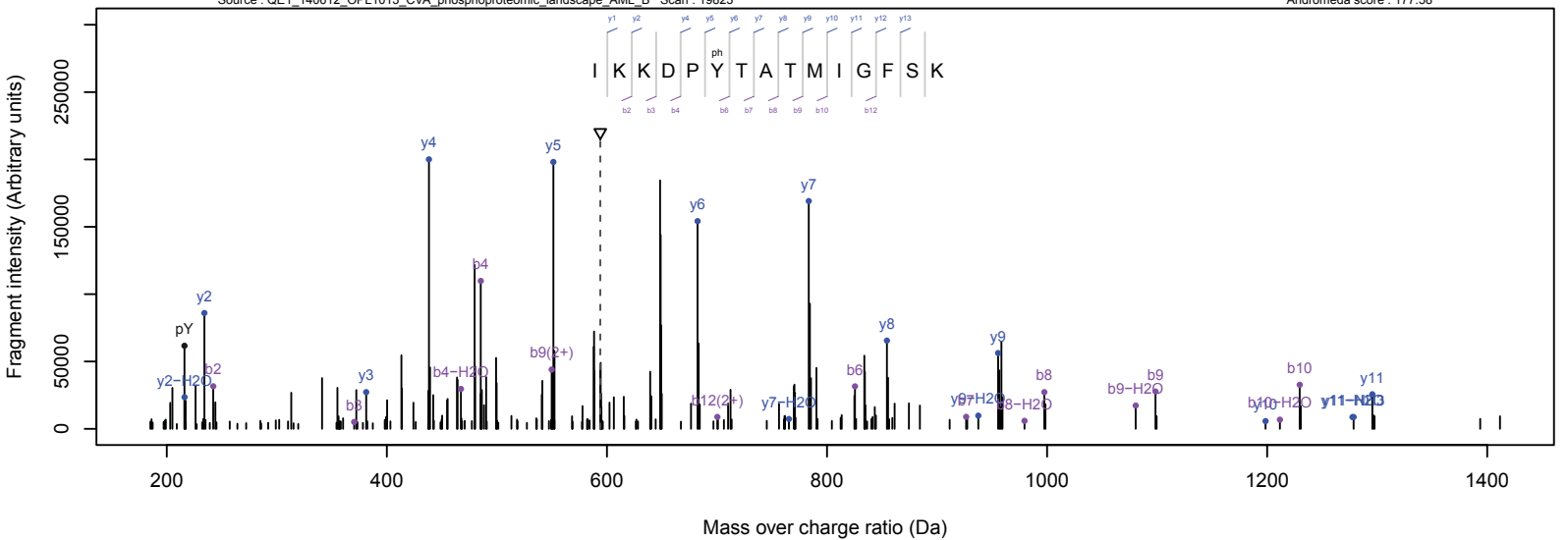
Andromeda score : 177.6



IKKDPYTATMIGFSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19823

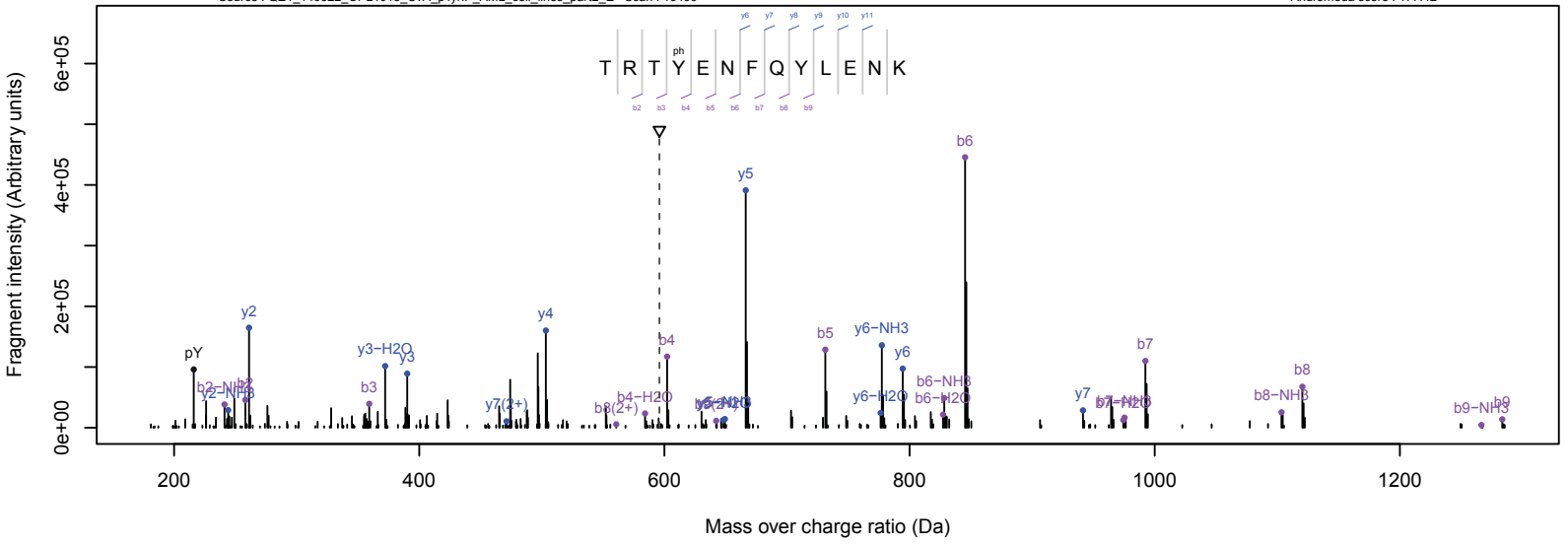
Andromeda score : 177.58



TRTY(ph)ENFQYLENK-3+

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 13499

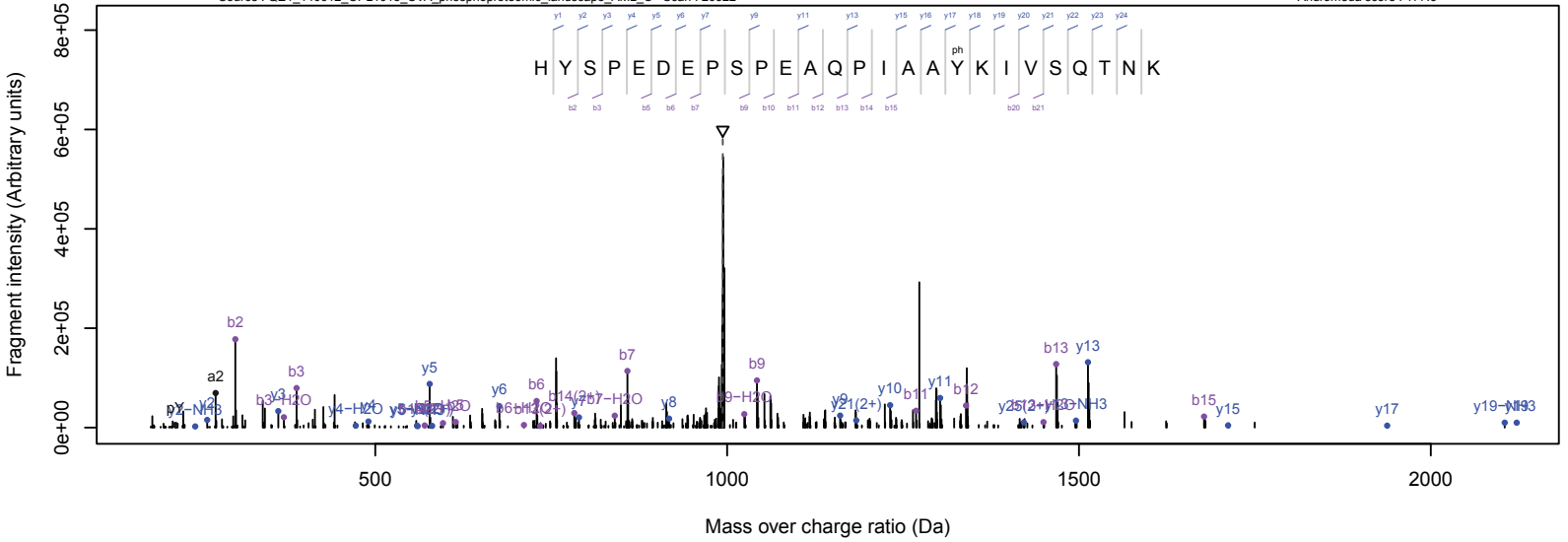
Andromeda score: 177.42



HYSPEDEPSPEAQPIAAYKIVSQTNK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26522

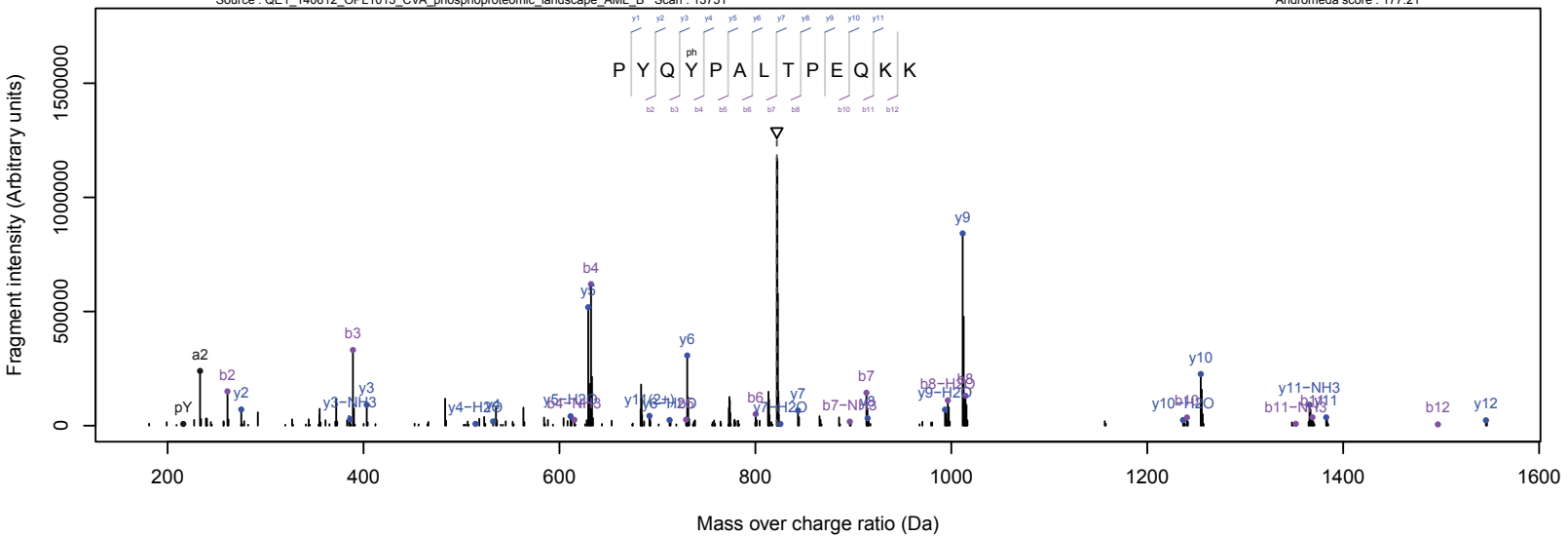
Andromeda score: 177.3



PYQY(ph)PALTPEQKK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 13751

Andromeda score: 177.21

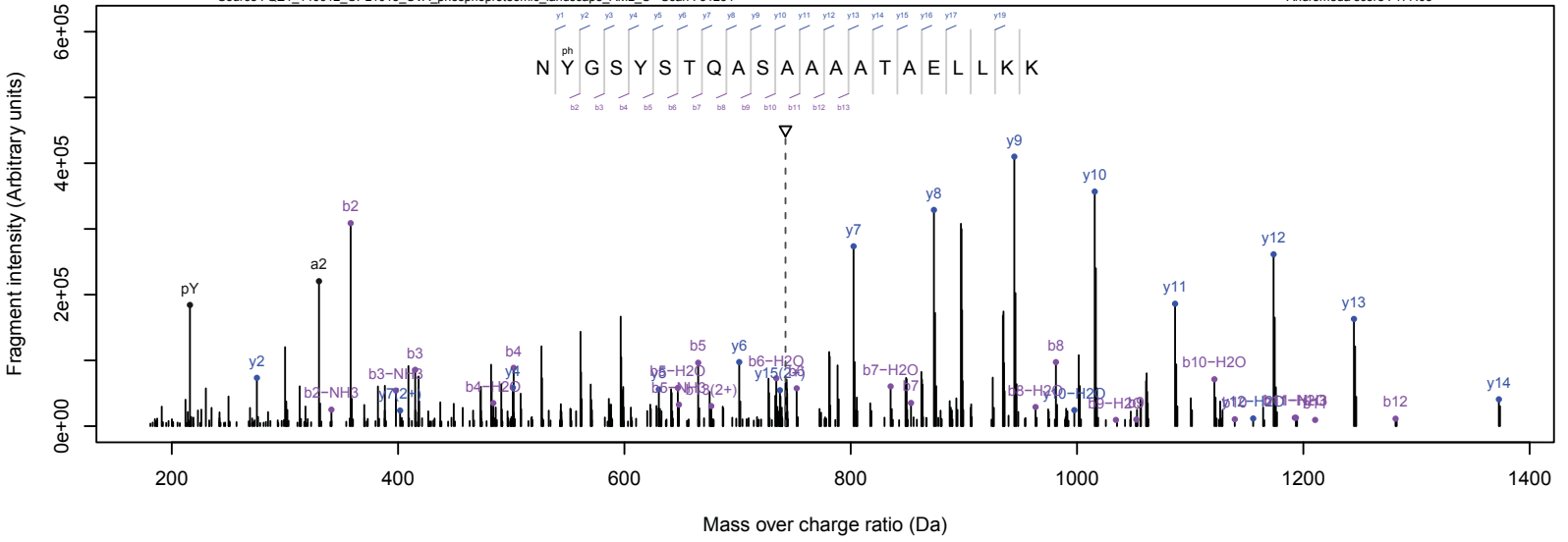




**NY(ph)GSYSTQASAAATAELLKK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31254

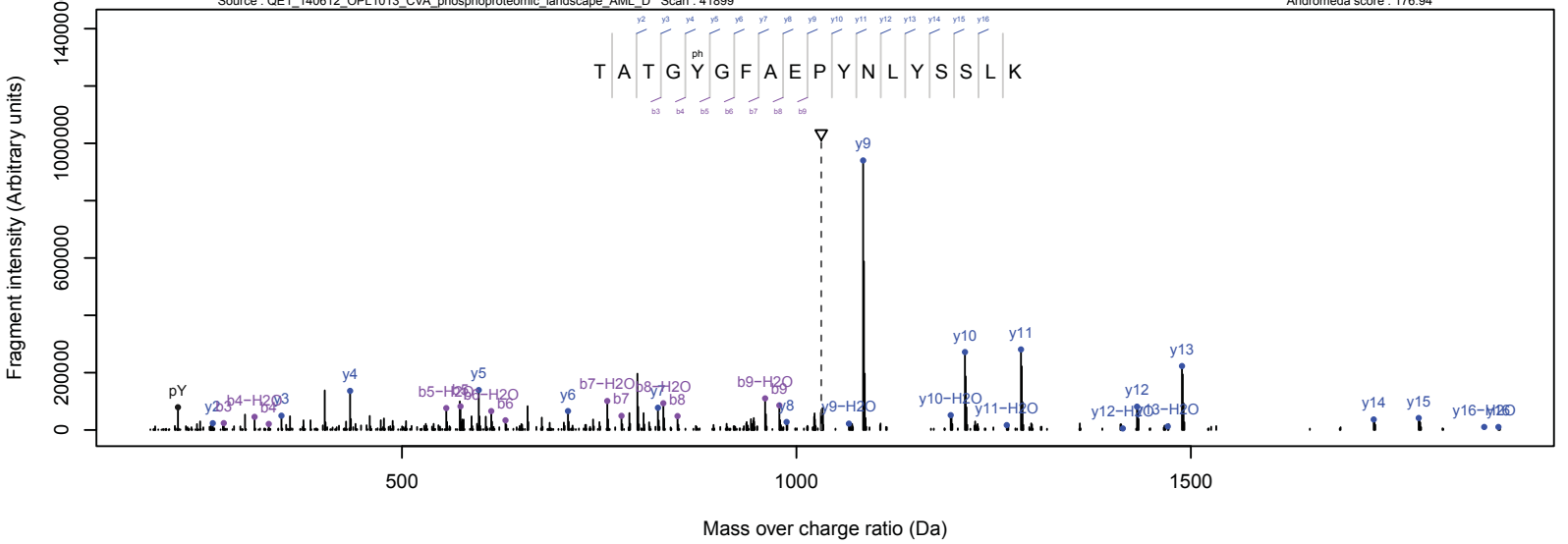
Andromeda score : 177.03



**TATGY(ph)GFAEPYNLYSSLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41899

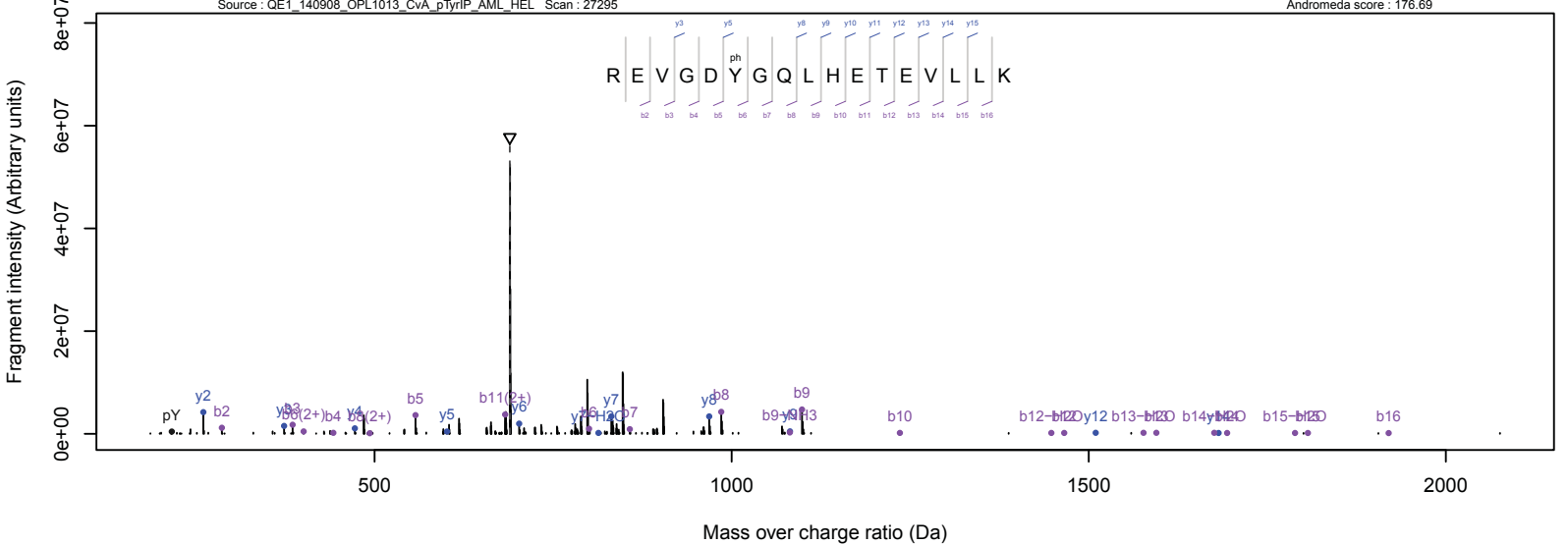
Andromeda score : 176.94



**REVG DY(ph)GQLHETEVLLK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 27295

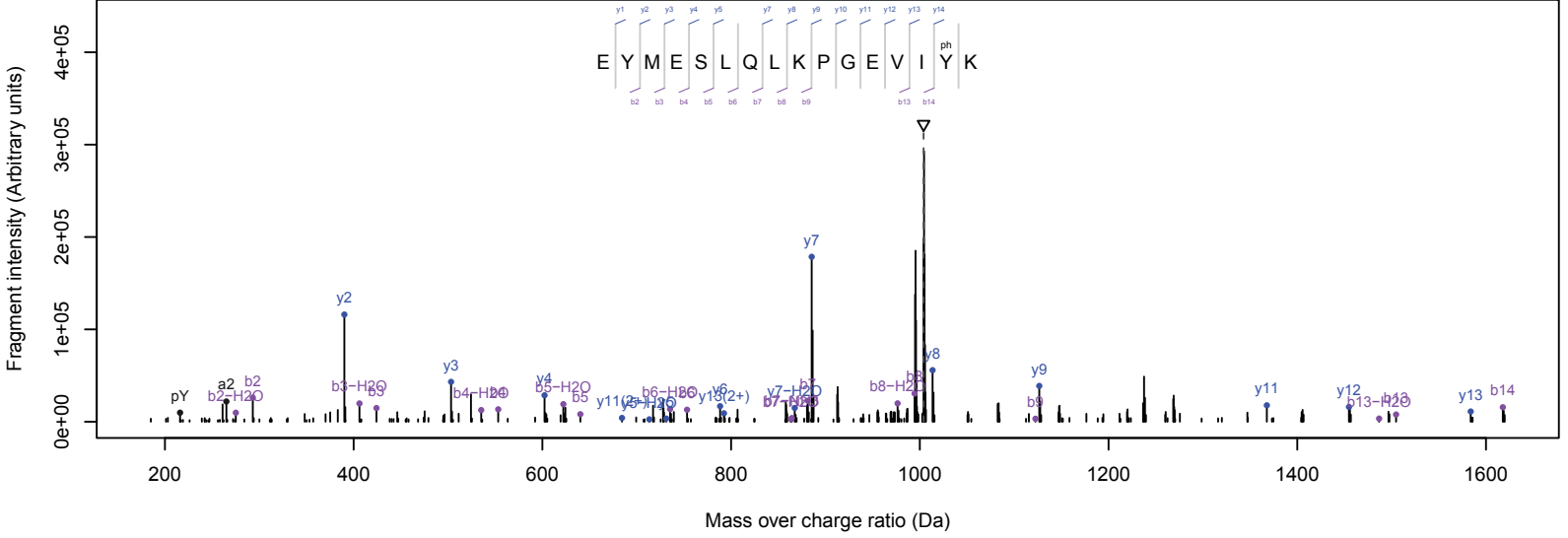
Andromeda score : 176.69



**EYMESLQLKPGEVY(ph)K-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 32479

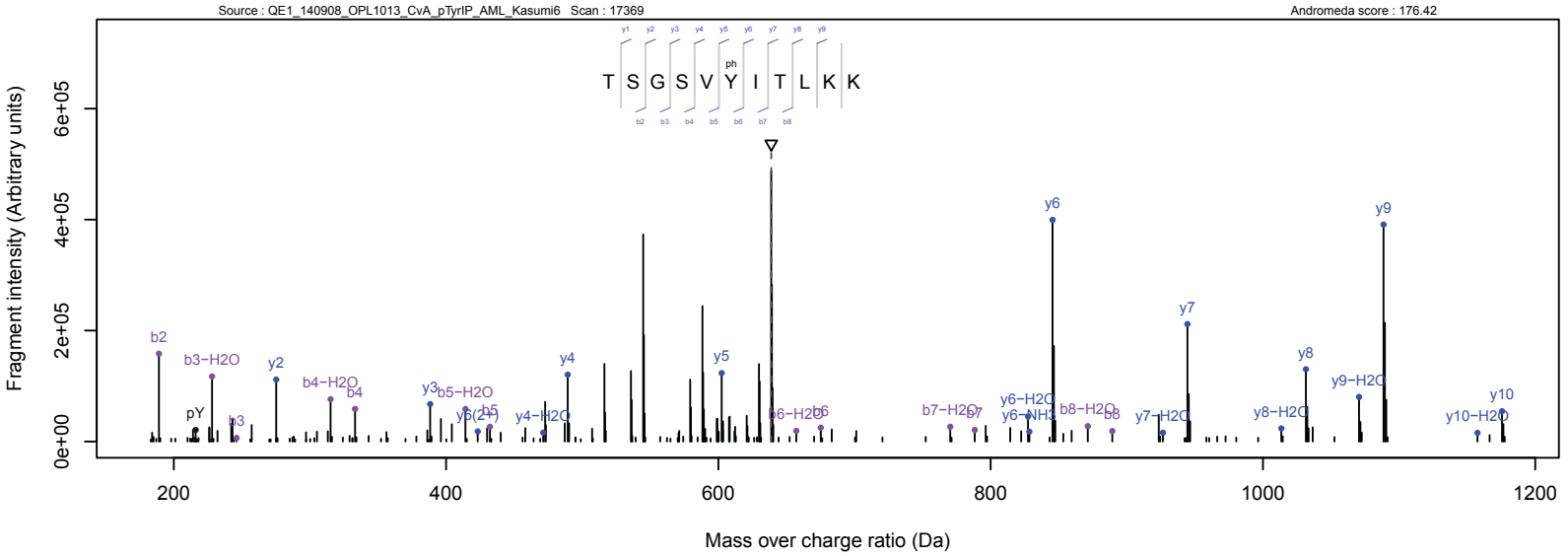
Andromeda score : 176.49



**TSGSVY(ph)ITLKK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 17369

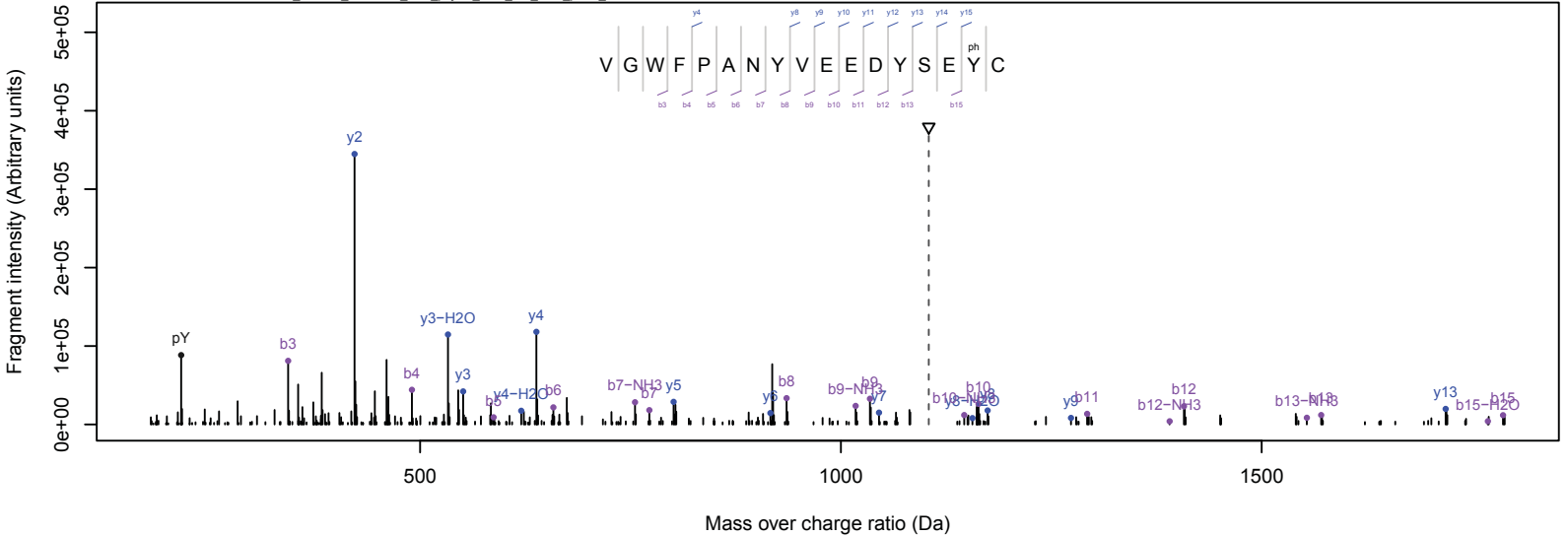
Andromeda score : 176.42



**VGWFpanyveedysey(ph)C-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 42939

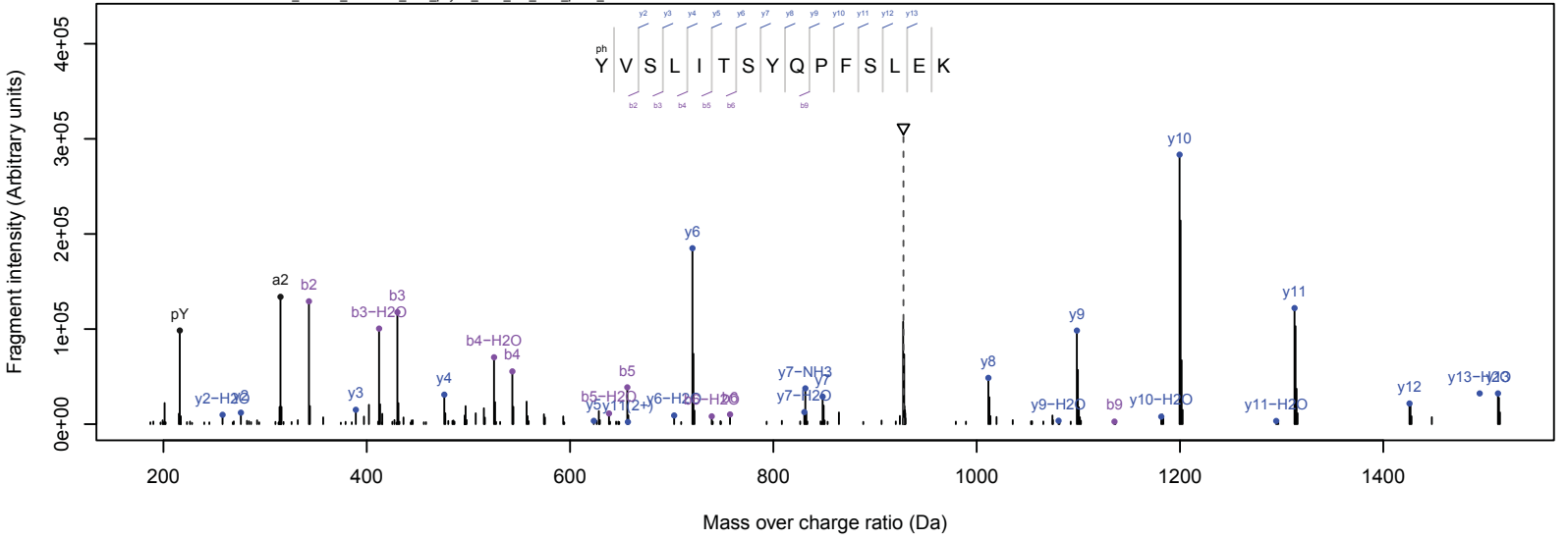
Andromeda score : 176.4



**Y(ph)VSLITSYQPFSLK-2+**

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan: 38995

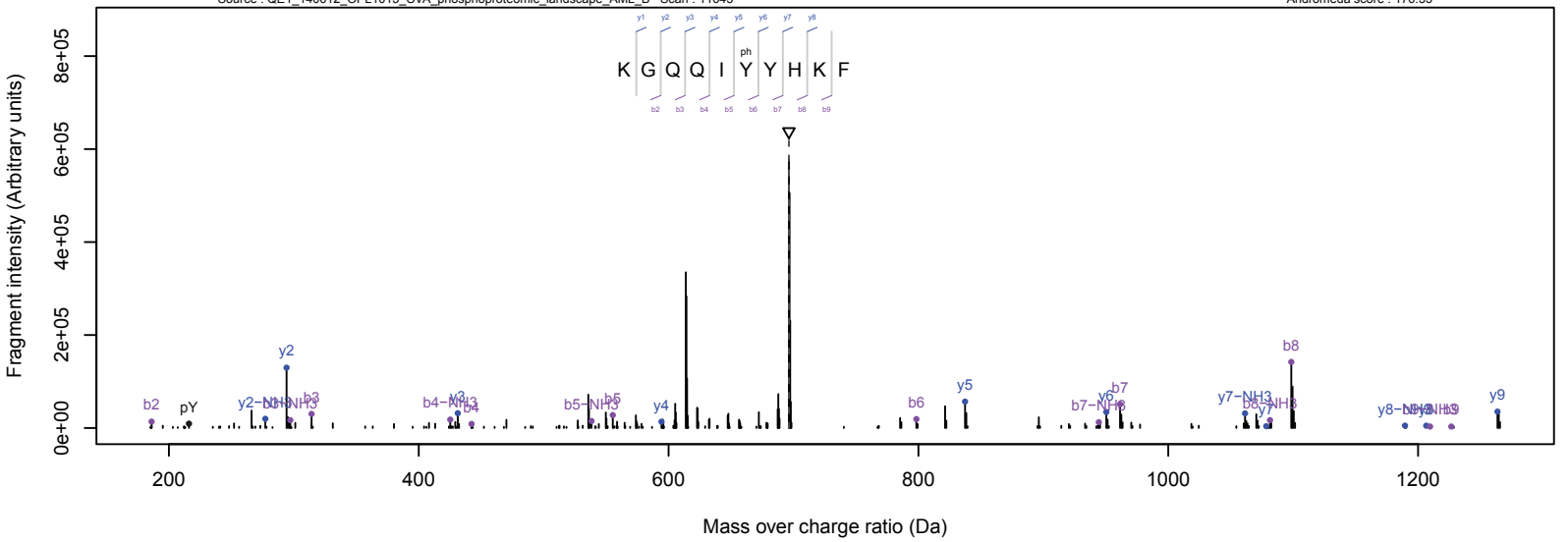
Andromeda score: 176.4



**KGQQIY(ph)YHKF-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 11643

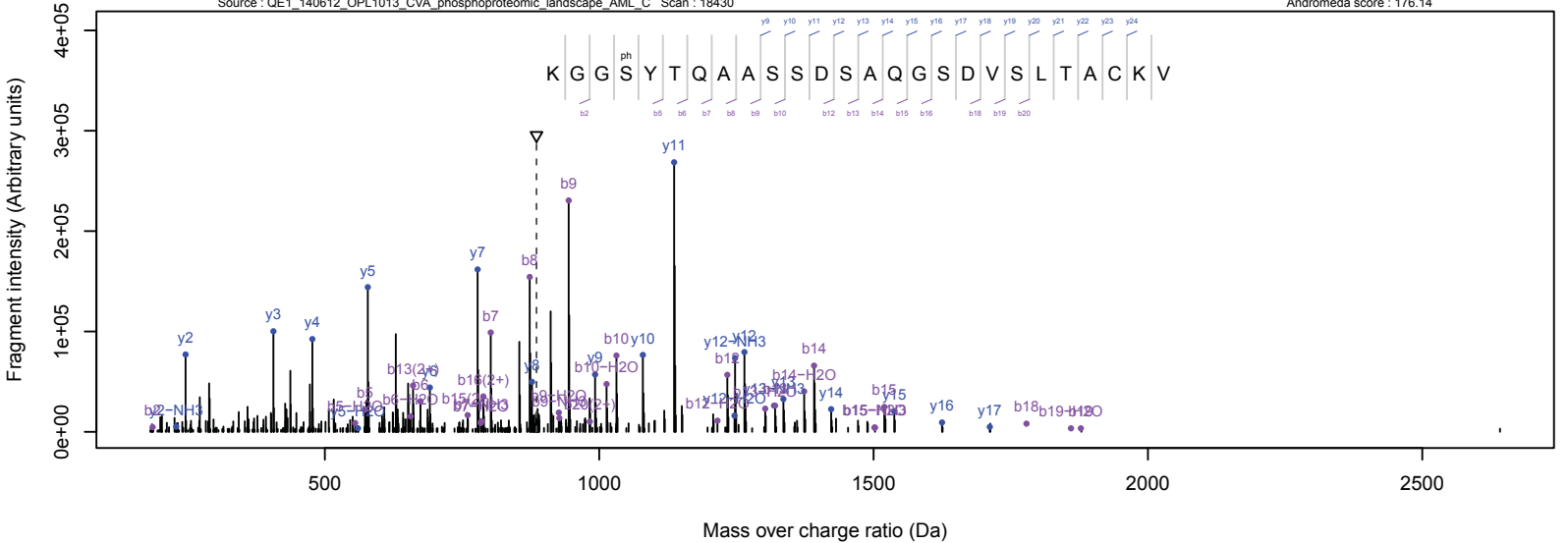
Andromeda score: 176.33



**KGGS(ph)YTQAASSDSAQGS DVSLTACKV-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18430

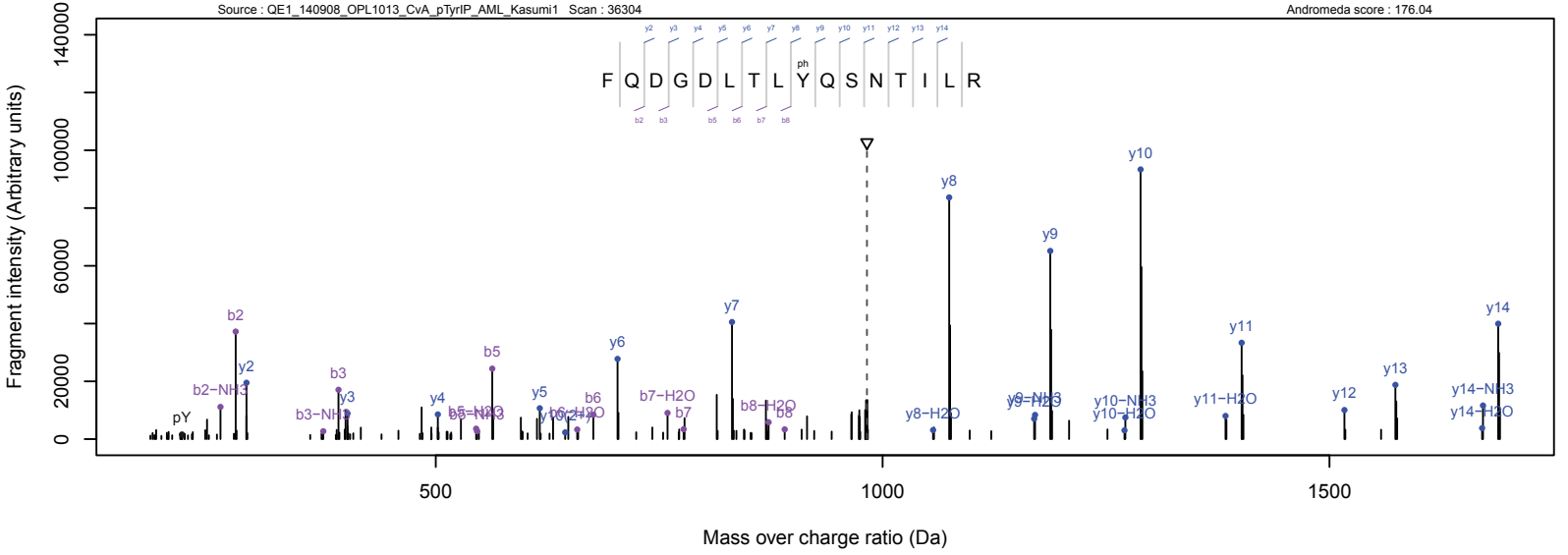
Andromeda score: 176.14



FQDGDLTLY(ph)QSNTILR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 36304

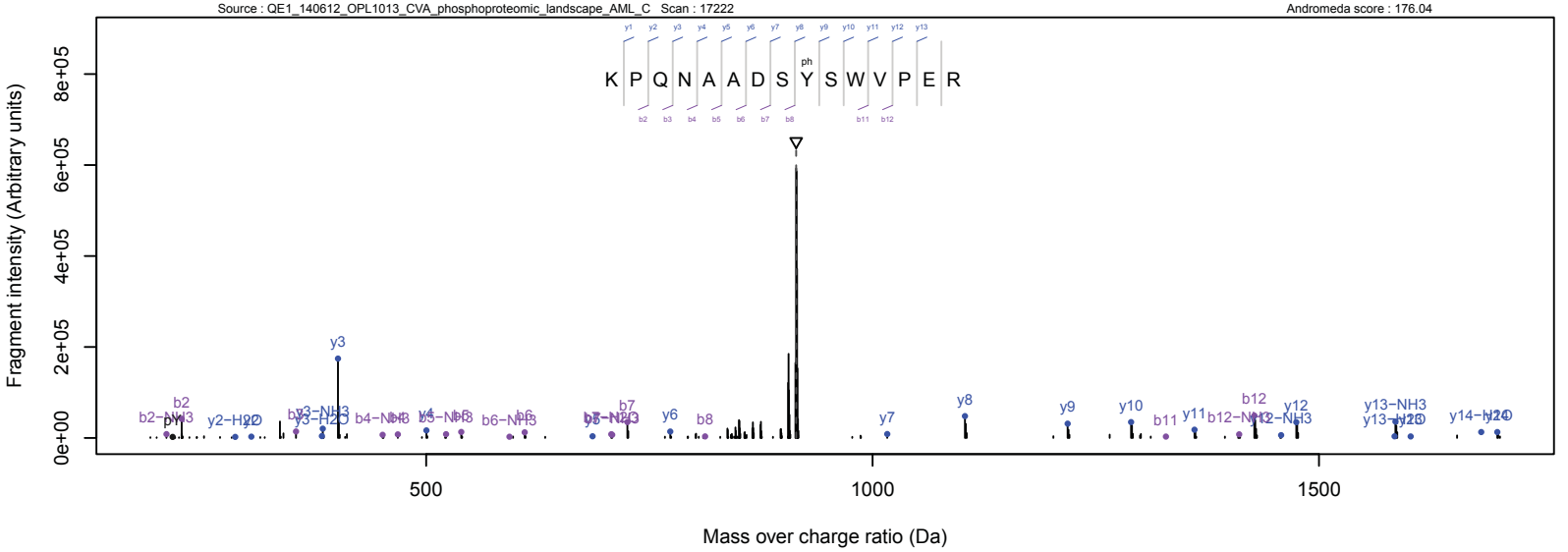
Andromeda score : 176.04



KPQNAADSY(ph)SWVPER-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 17222

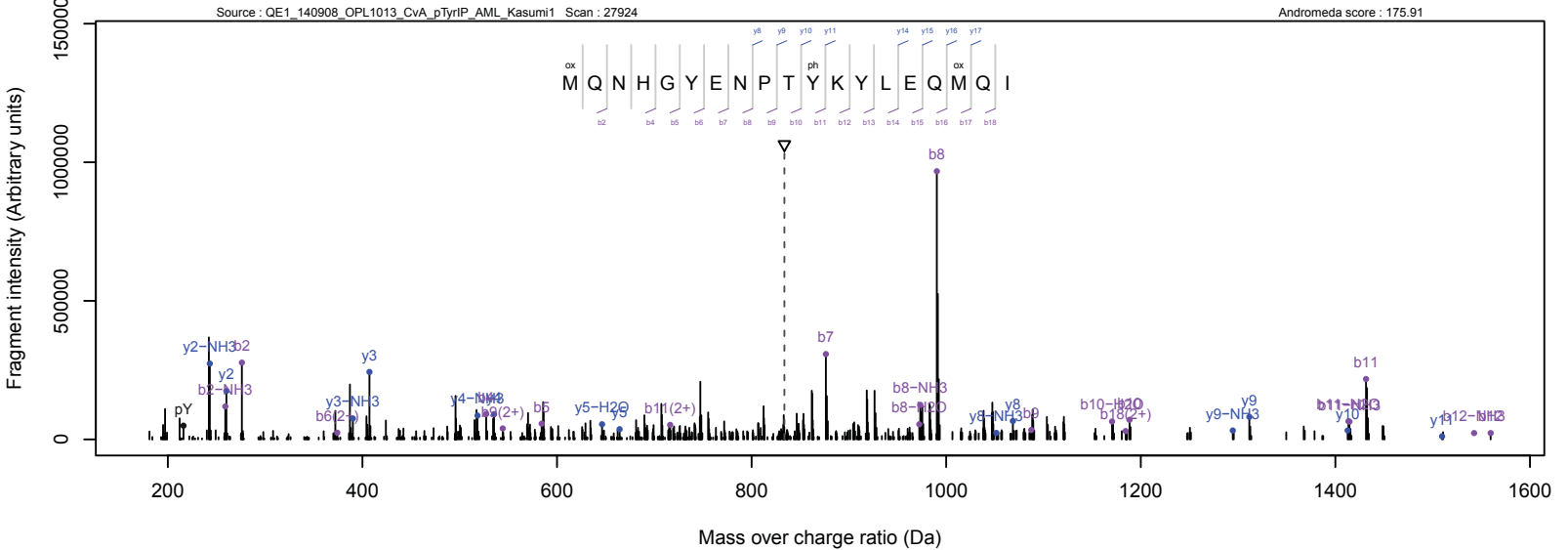
Andromeda score : 176.04



M(ox)QNHGYLENPTY(ph)KYLEQM(ox)QI-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27924

Andromeda score : 175.91

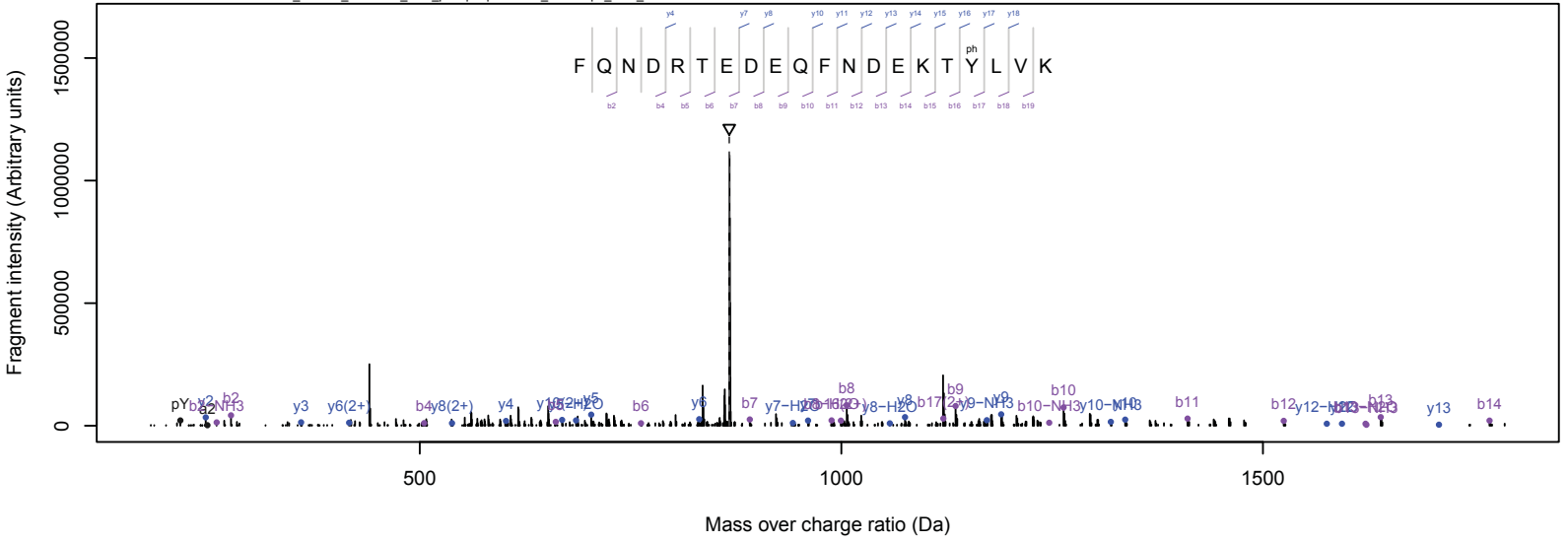




FQNDRTEDQFNDEKTY(ph)LVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19868

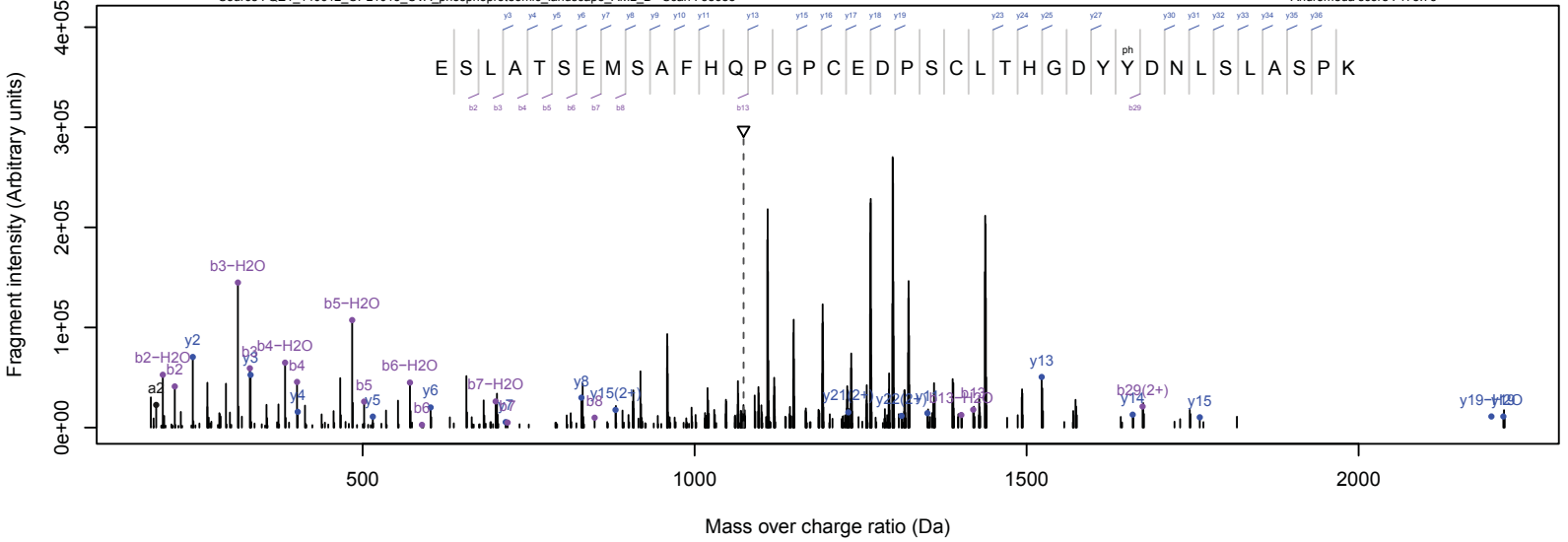
Andromeda score : 175.77



ESLATSEMSAFHQGPCEDPSCSLTHGDYY(ph)DNLSLSPK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38688

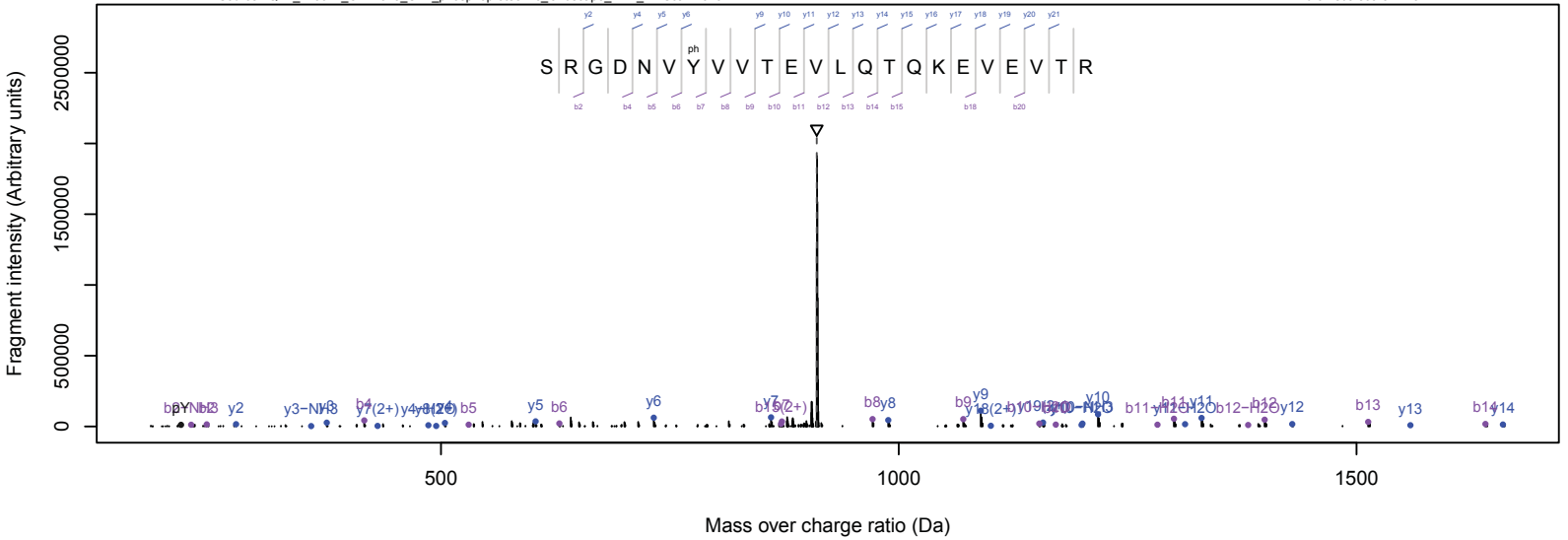
Andromeda score : 175.75



SRGDNVY(ph)VVTEVLQTQKEVETR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 37324

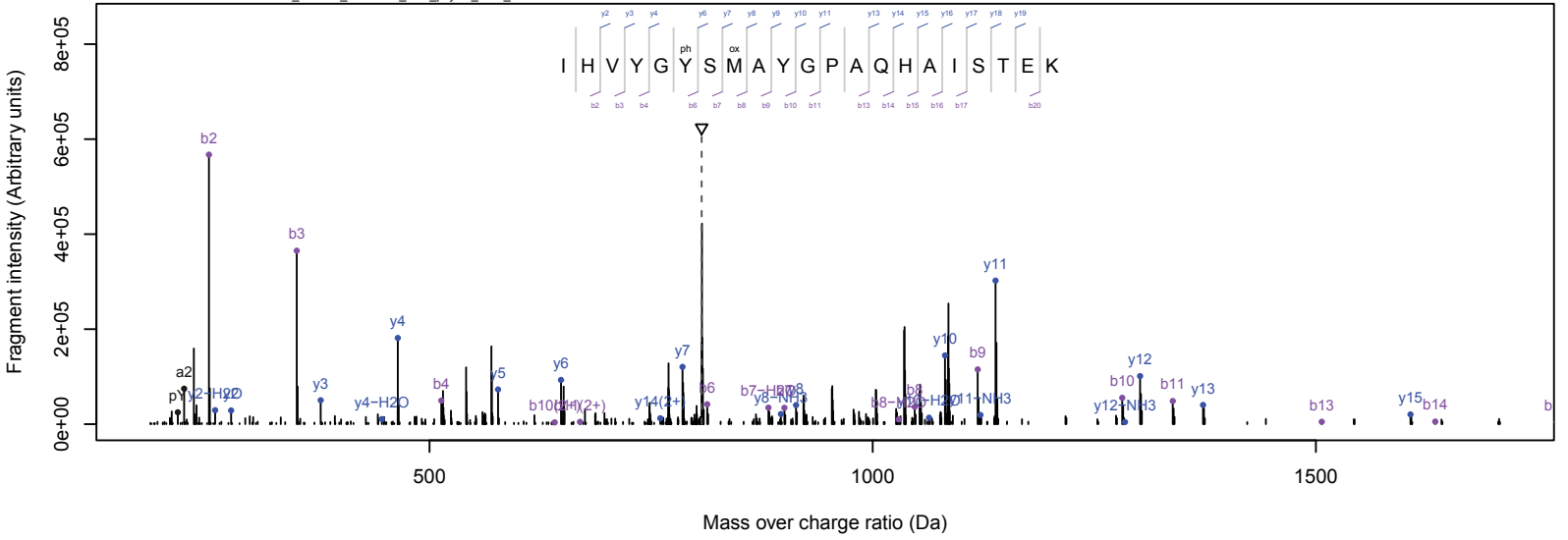
Andromeda score : 175.71



IHVYGY(ph)SM(ox)AYGPAQHAISTEK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23100

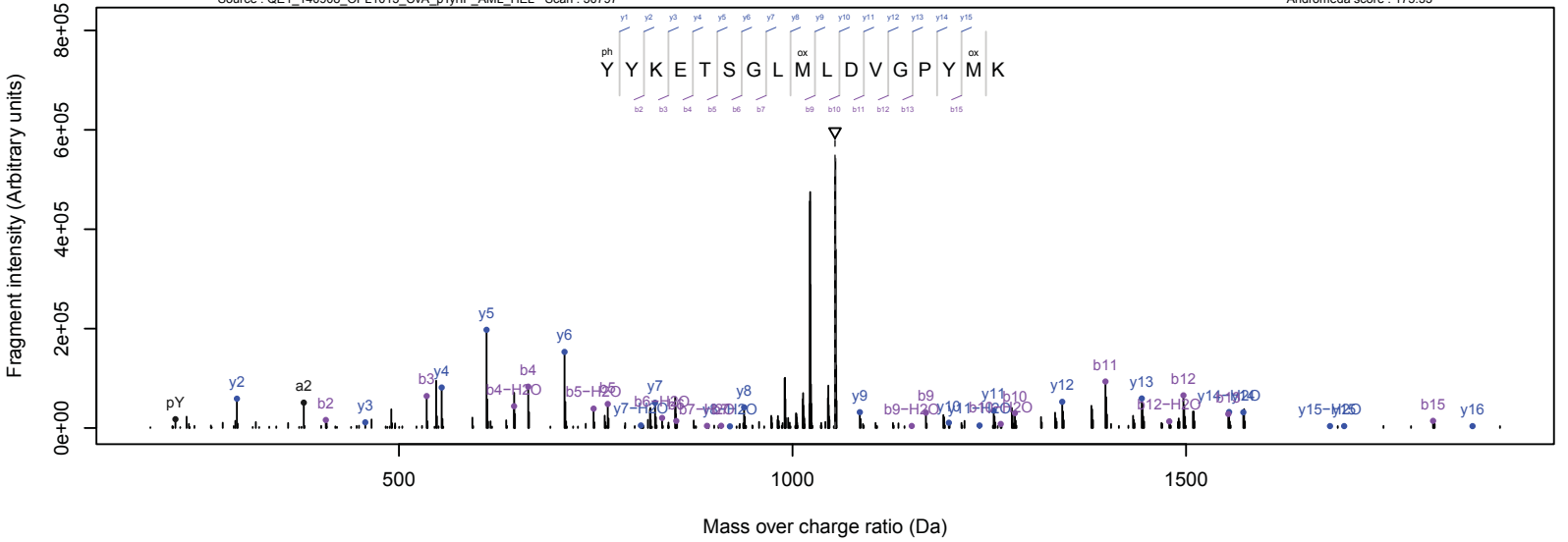
Andromeda score : 175.65



Y(ph)YKETSGLM(ox)LDVGPYM(ox)K-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 30797

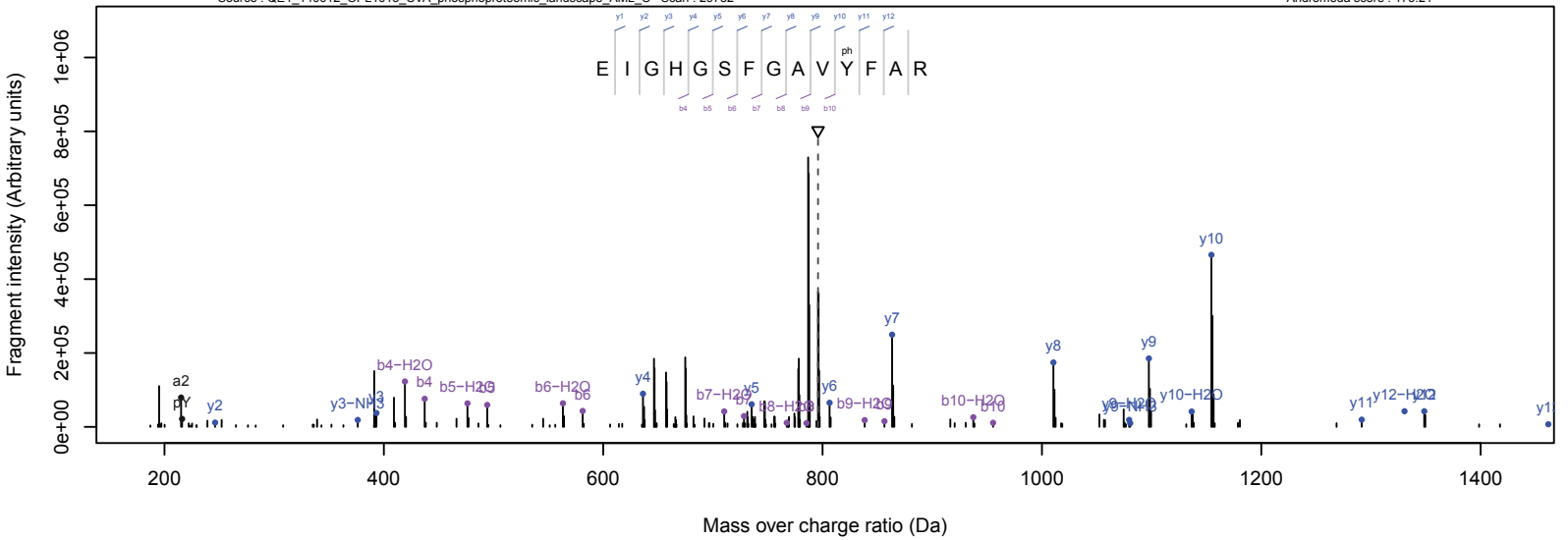
Andromeda score : 175.33



EIGHGSFGAVY(ph)FAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29782

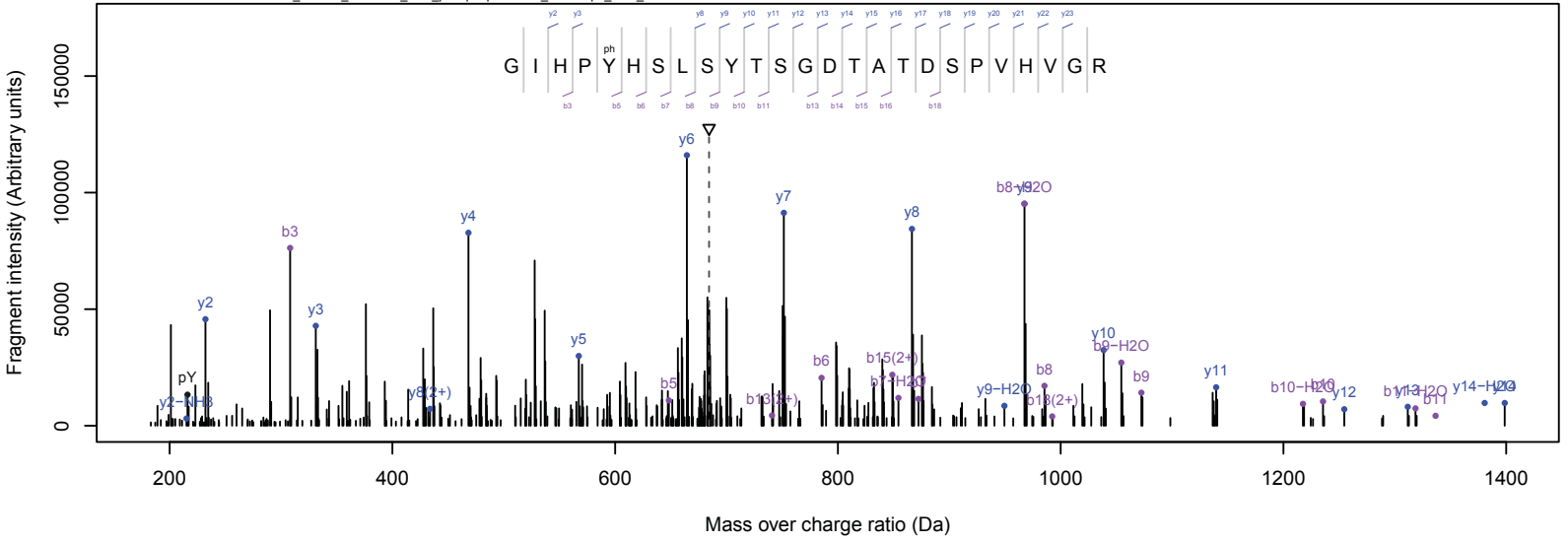
Andromeda score : 175.21



GIHPY(ph)HLSYTSYSGDTATDSPVHVGR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 20607

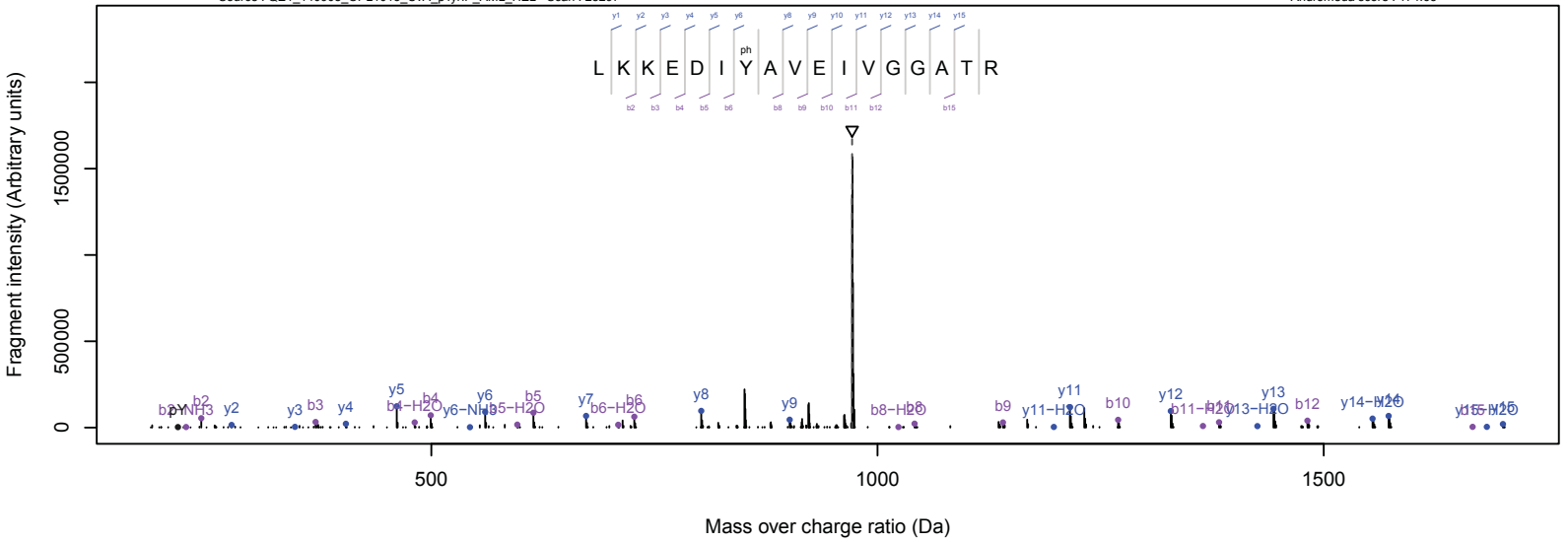
Andromeda score: 174.7



LKKEDIY(ph)AVEIVGGATR-2+

Source: QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_HEL Scan: 28297

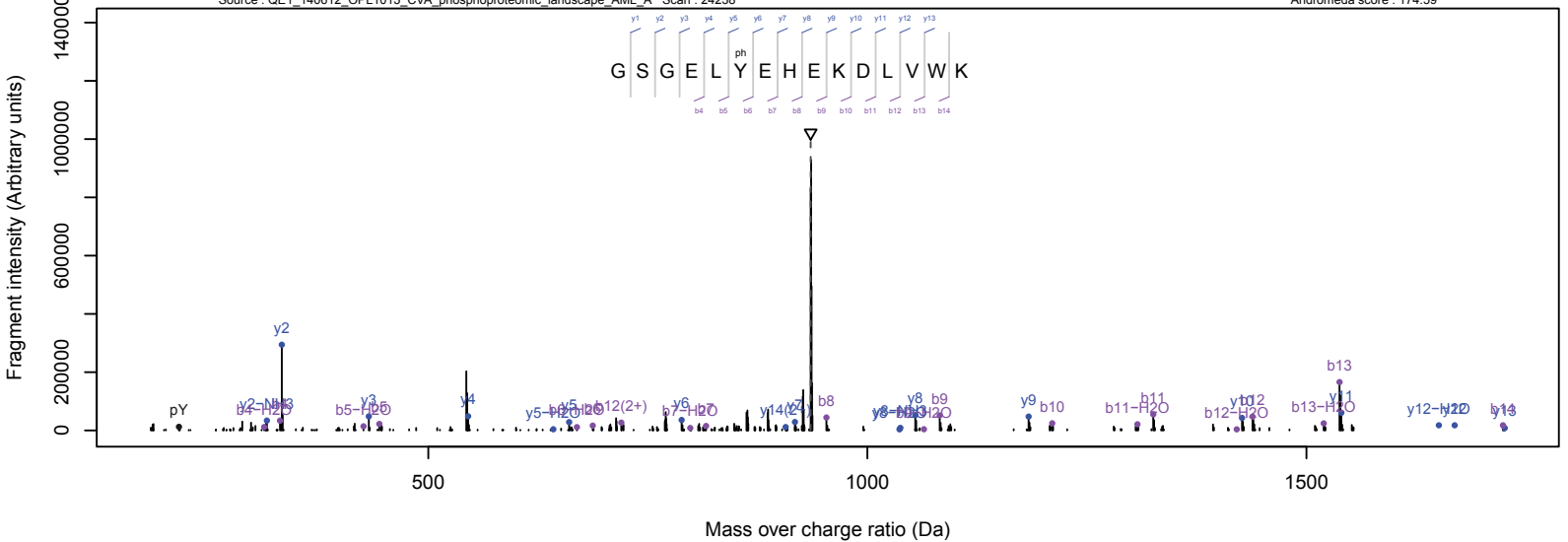
Andromeda score: 174.65



GSGELY(ph)EHEKDLVWK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 24238

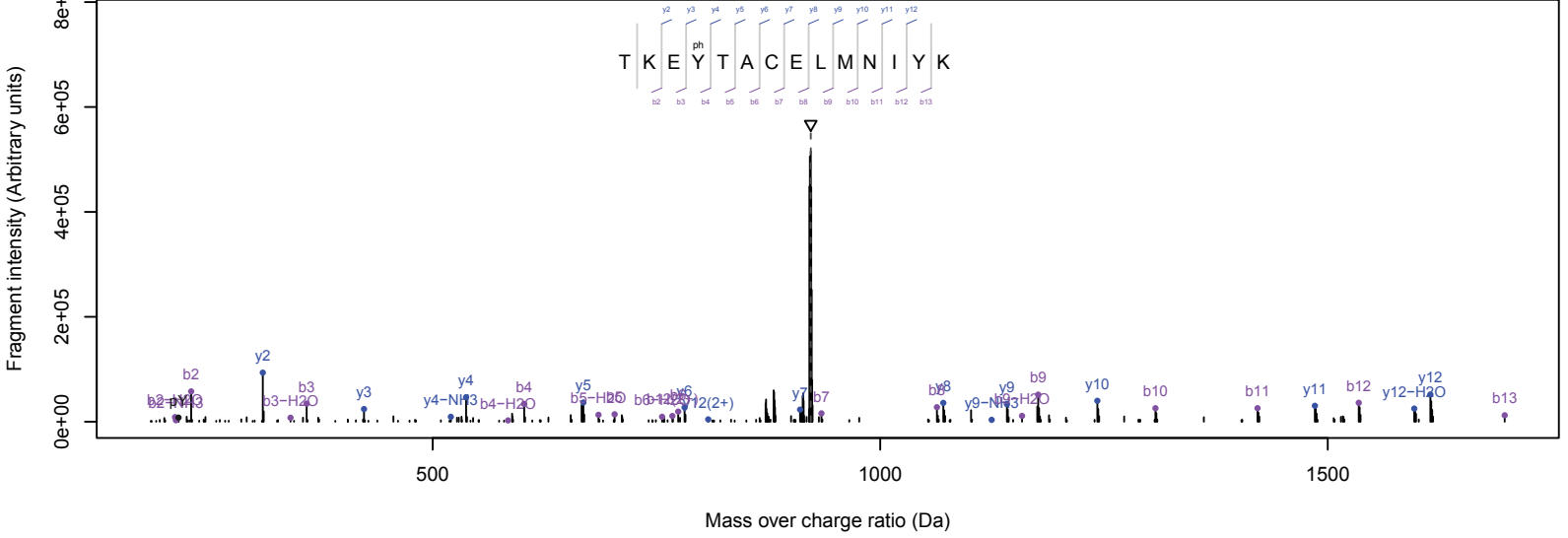
Andromeda score: 174.59



TKEY(ph)TACELMNIYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29859

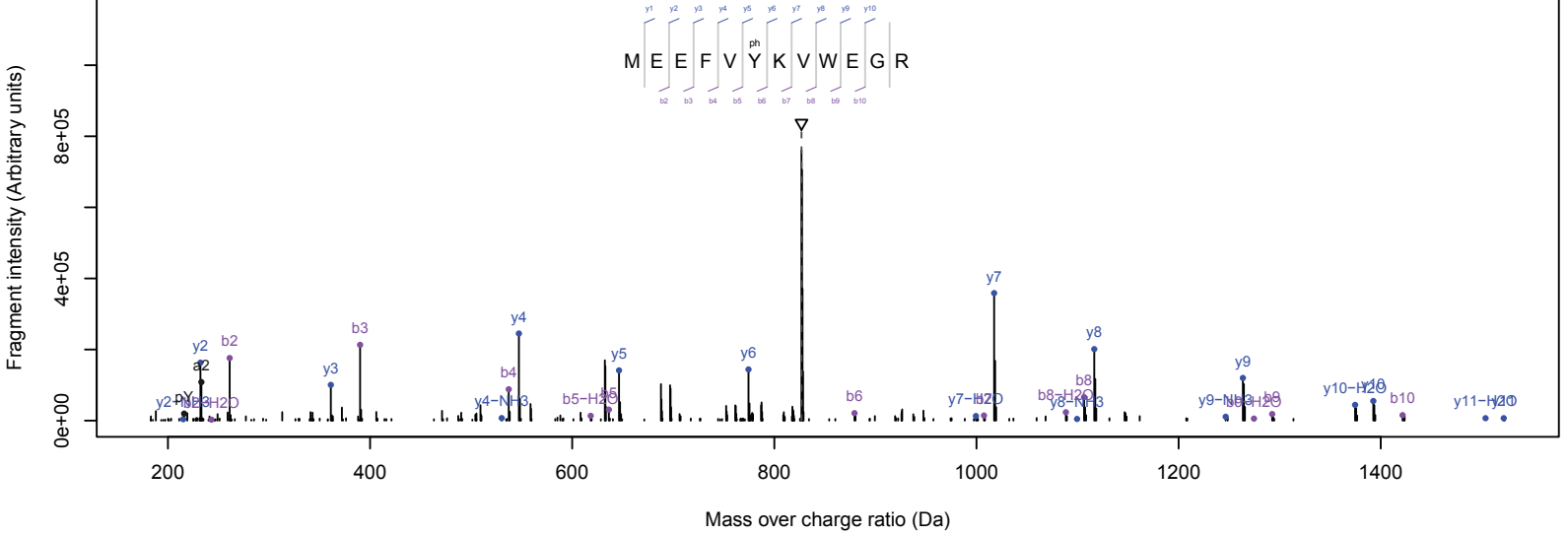
Andromeda score : 174.59



MEEFVY(ph)K VWEGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35745

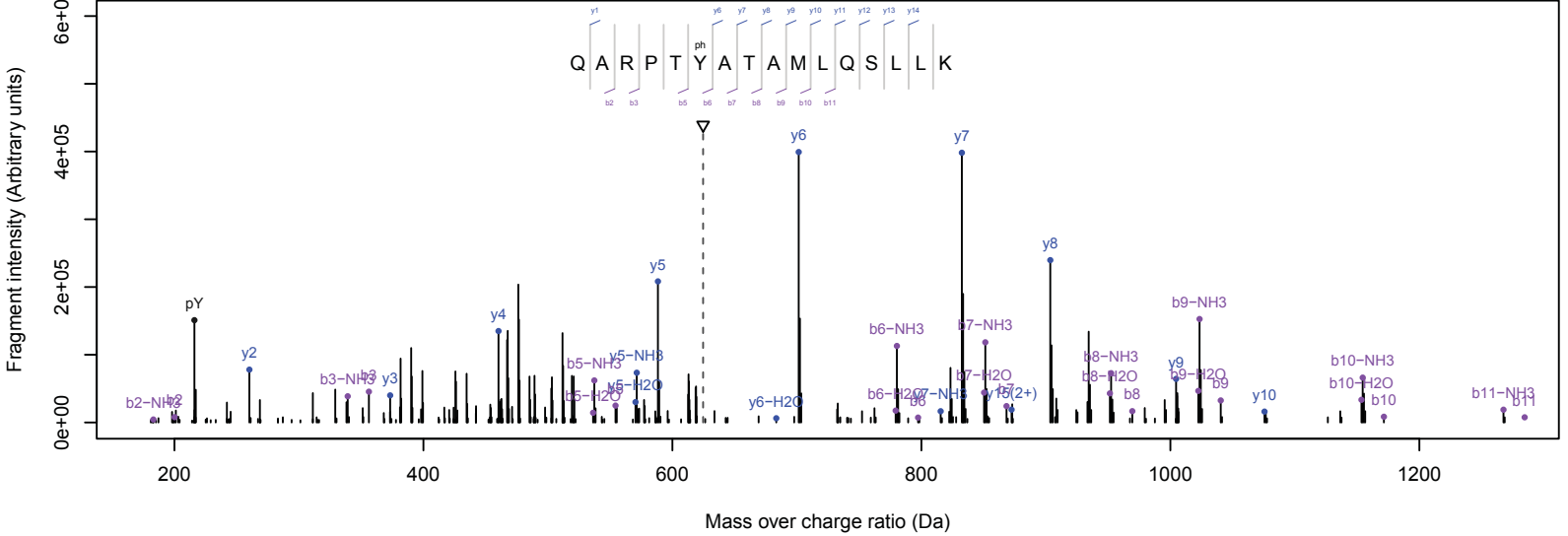
Andromeda score : 174.39



QARPTY(ph)ATAMLSLLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43159

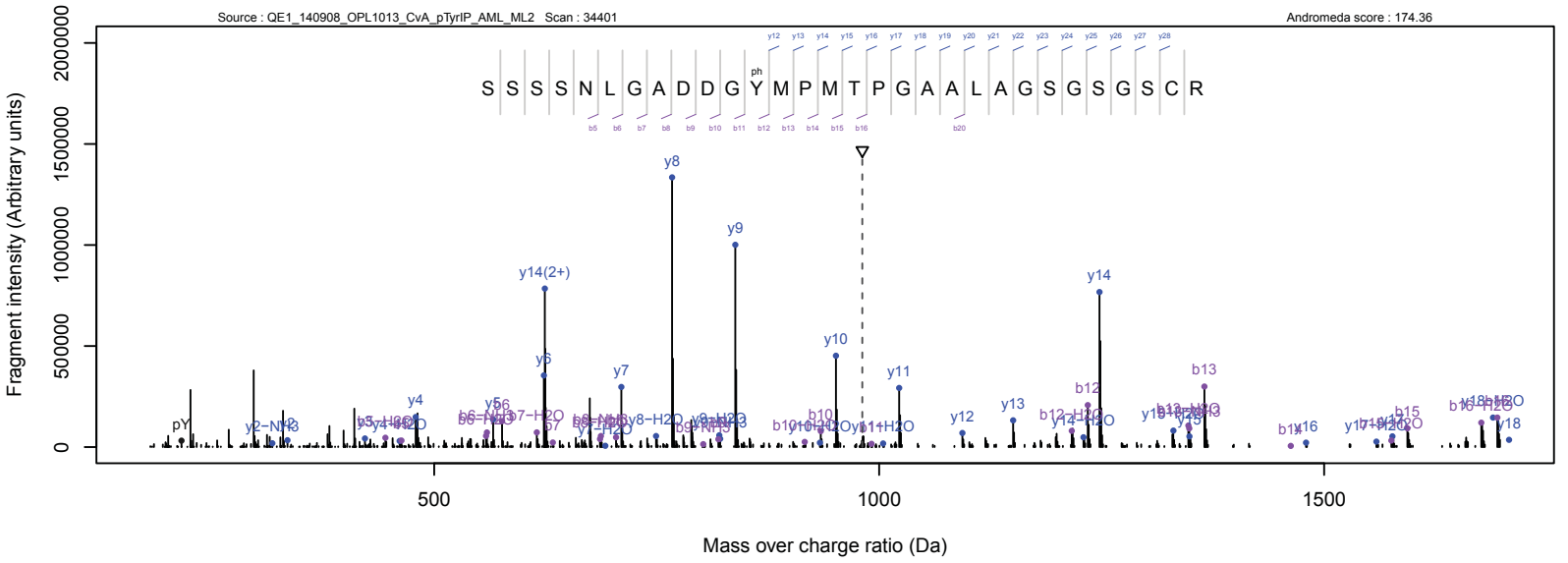
Andromeda score : 174.39



SSSSNLGADDGY(ph)MPMTPGAALAGSGSGSCR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 34401

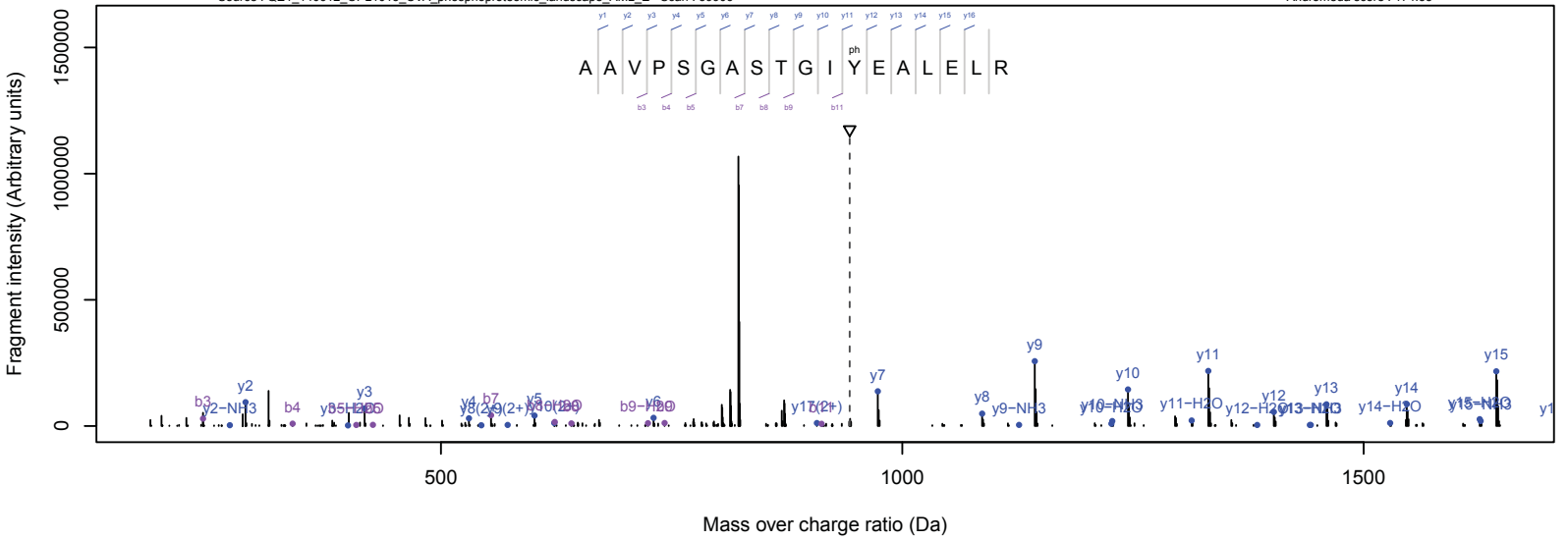
Andromeda score : 174.36



AAVPSGASTGIY(ph)EALELR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 39906

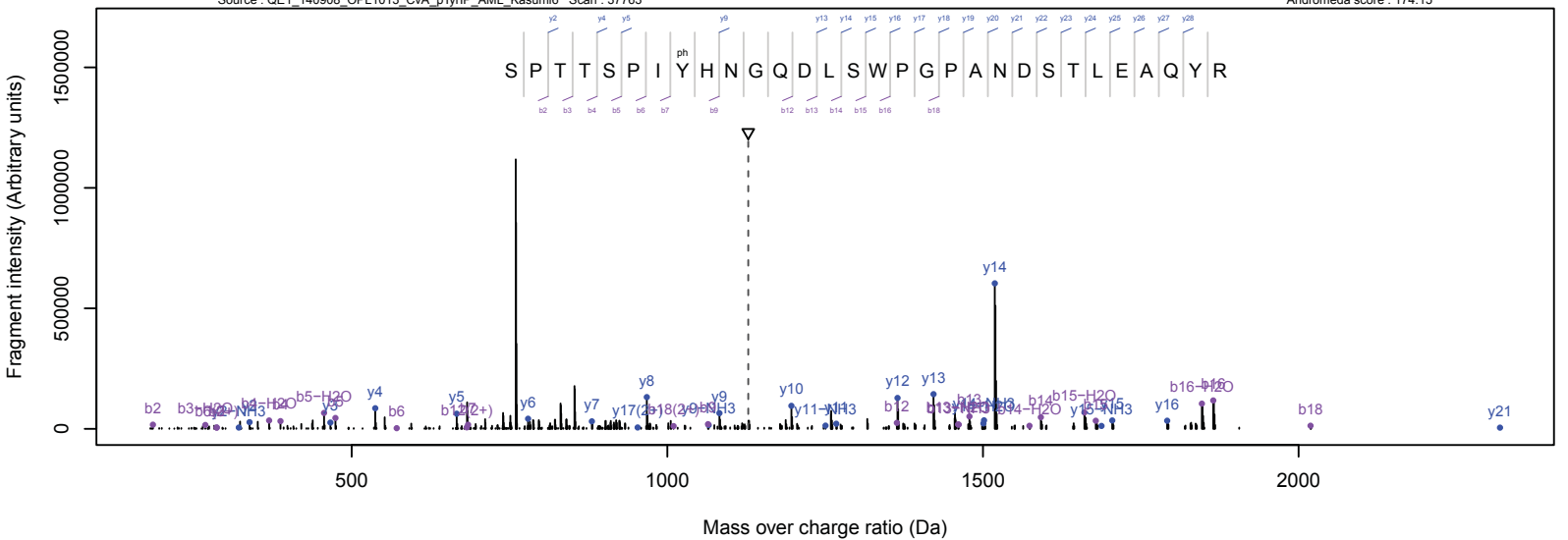
Andromeda score : 174.35



SPTTSPYIYHNGQDLSWPGPANDSTLEAQYR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 37763

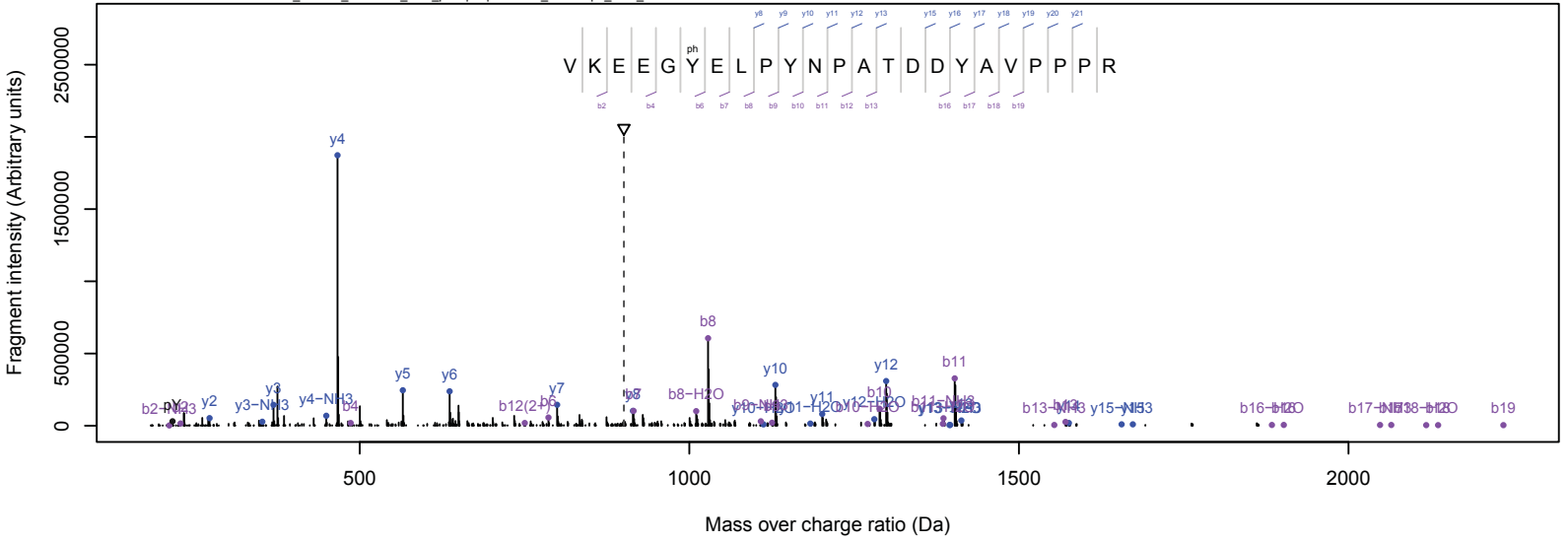
Andromeda score : 174.15



VKEEGY(ph)ELPYNPATDDYAVPPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31210

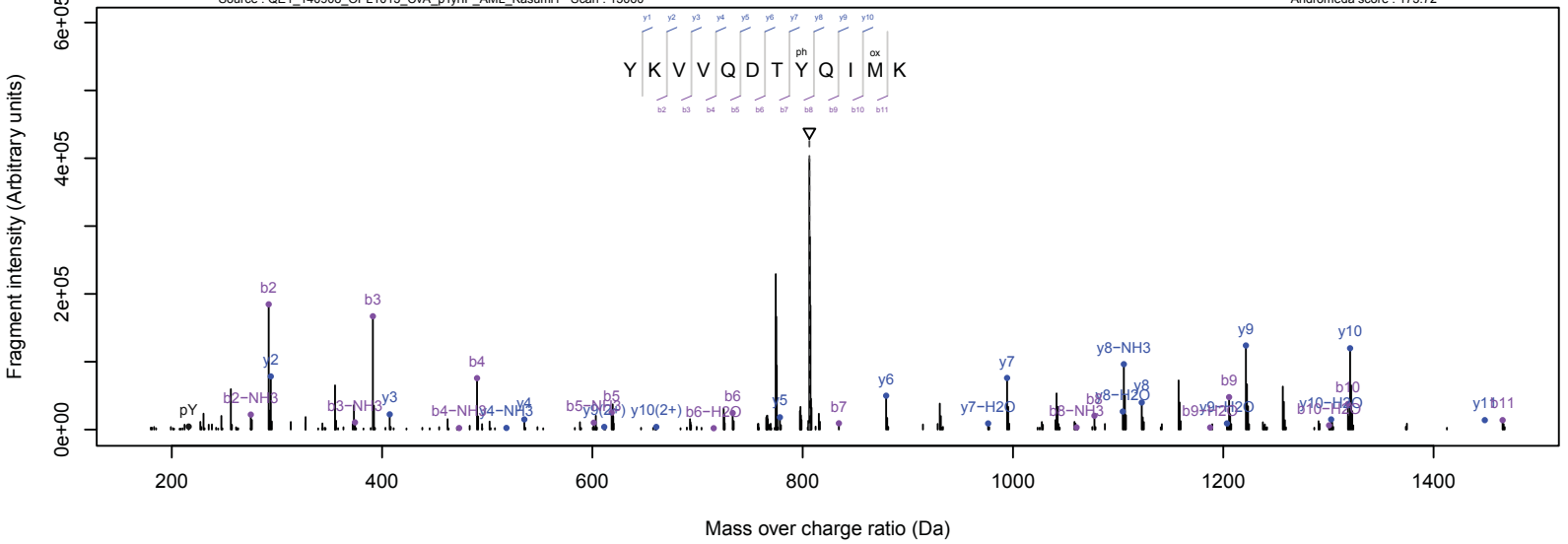
Andromeda score: 174.01



YKVVQDTY(ph)QIM(ox)K-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 13060

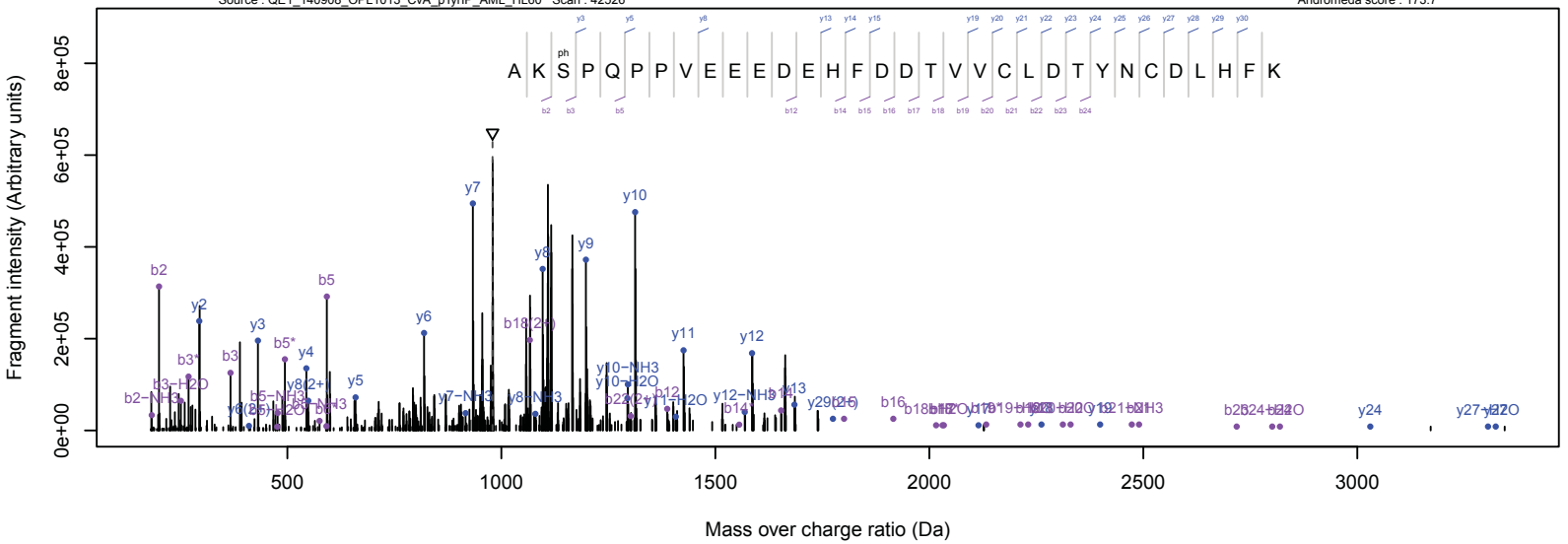
Andromeda score: 173.72



AKS(ph)PQPVEEEDHFD DTVVCLDTYNCDLHFK-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan: 42526

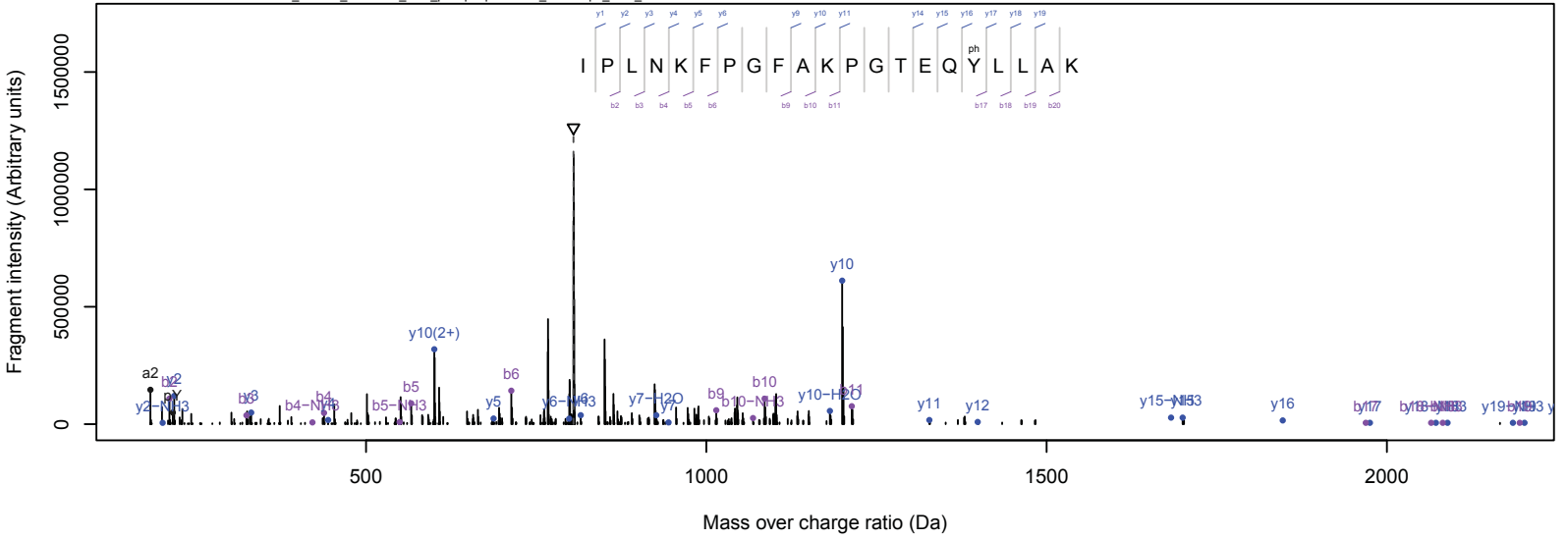
Andromeda score: 173.7



IPLNKFPGFAPGTEQY(ph)LLAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37097

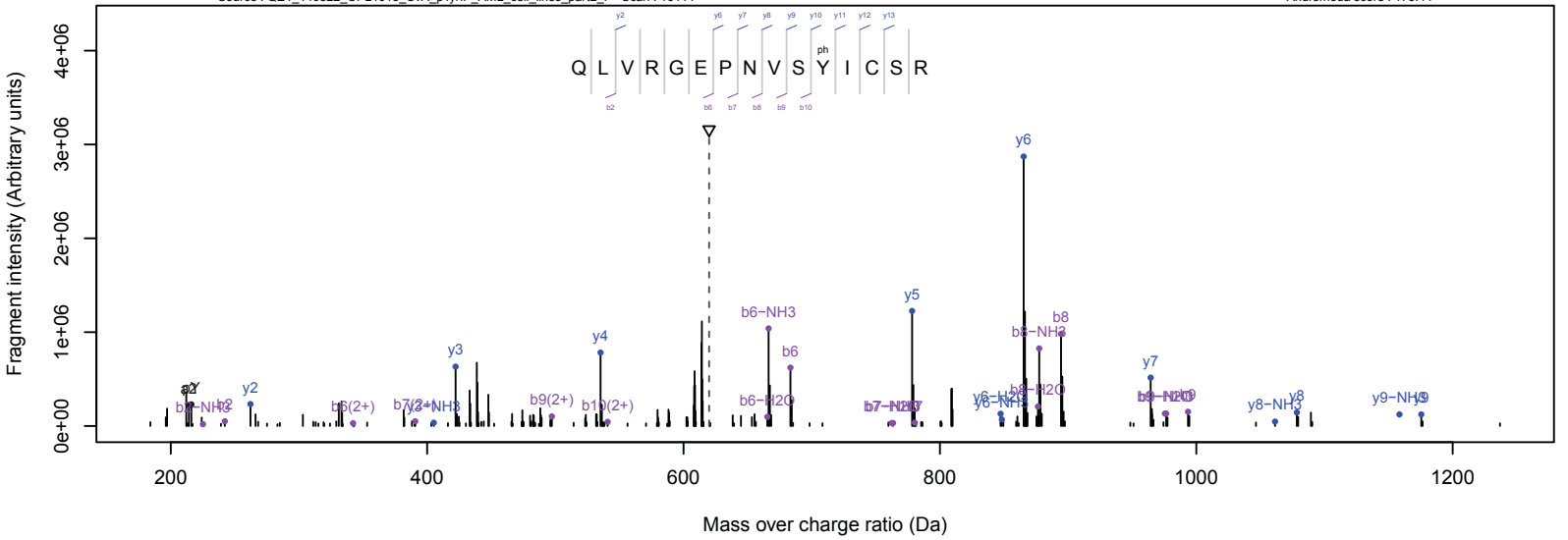
Andromeda score : 173.56



QLVRGEPNVSYP(h)ICSR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 16114

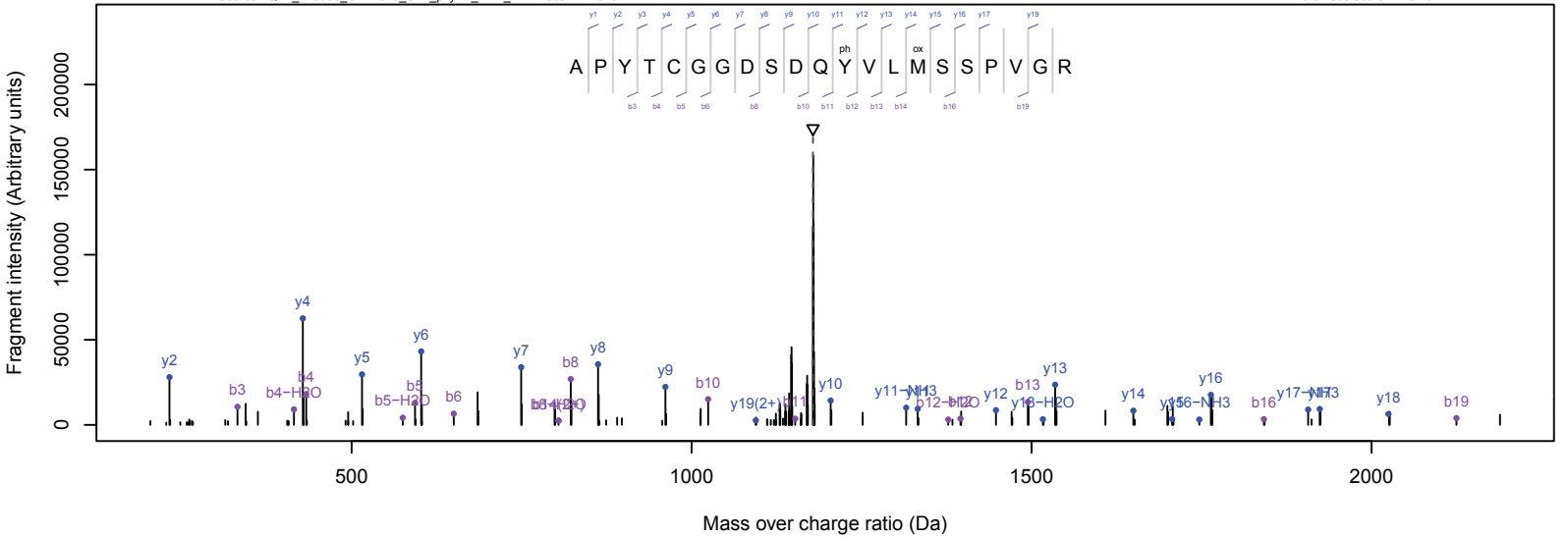
Andromeda score : 173.44



APYTCGGDSQY(ph)VLMSSPVGGR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 27375

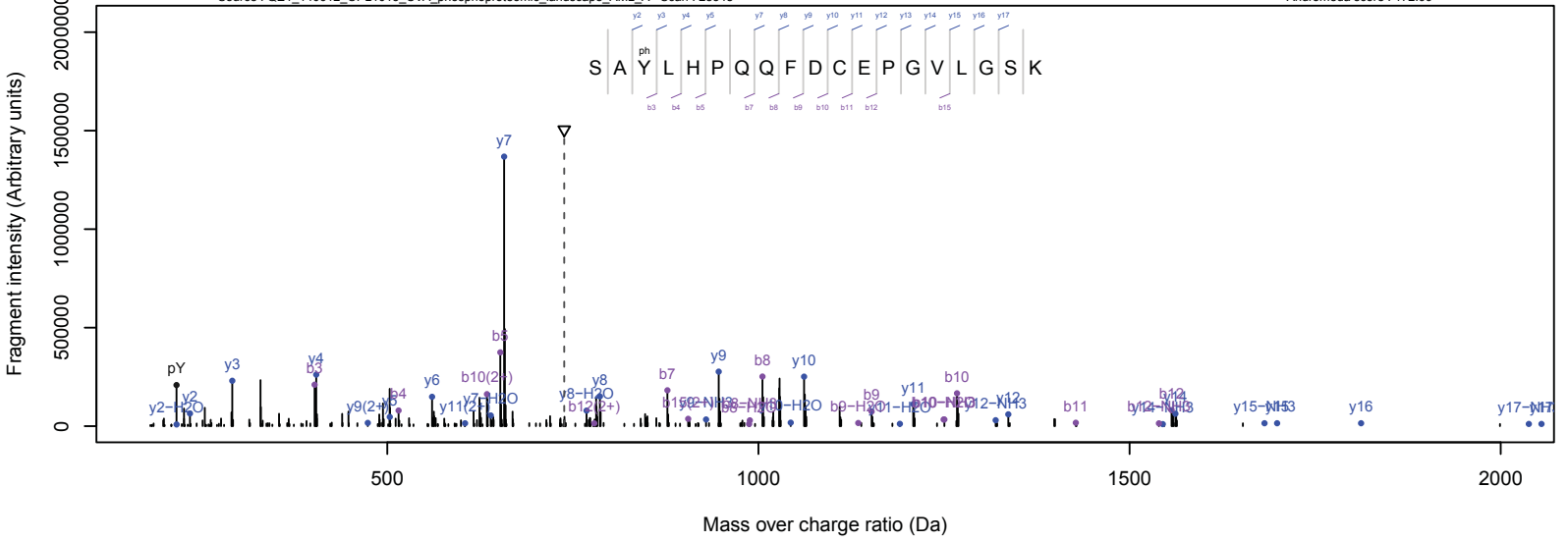
Andromeda score : 173.43



SAY(ph)LHPQQFDCPEPGLGSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 28348

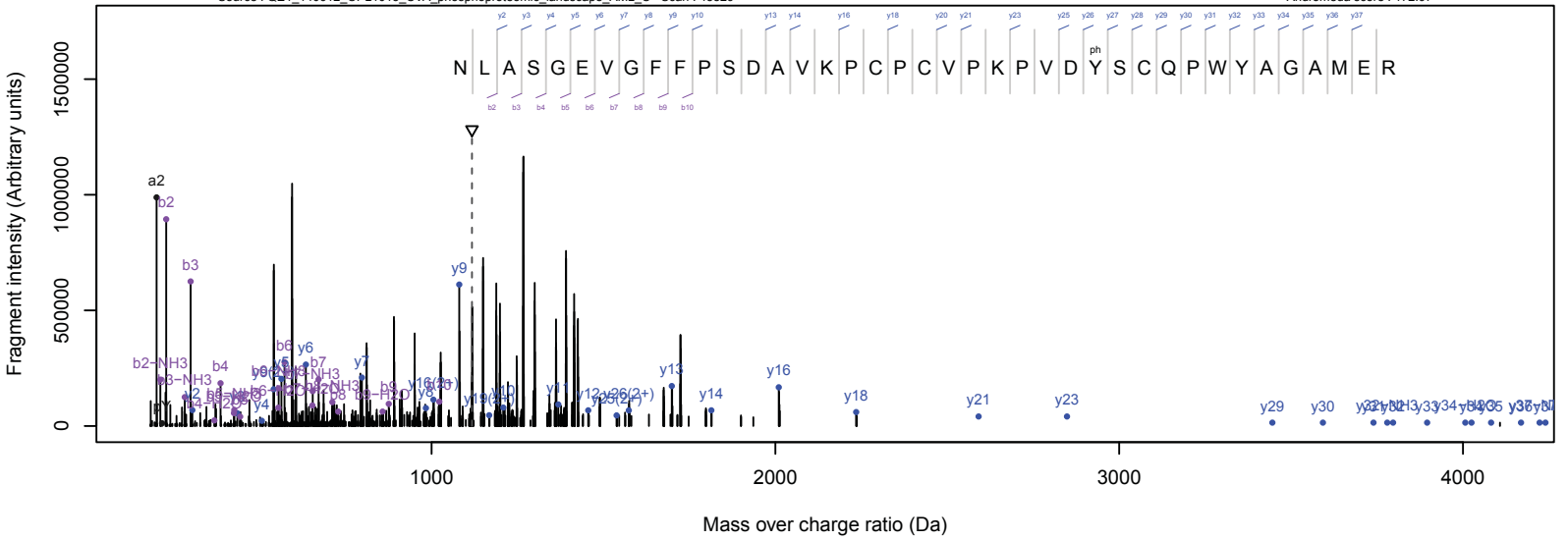
Andromeda score : 172.99



NLASEGVFFPSDAVKPCVCPKVDY(ph)SCQPWYAGAMER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45829

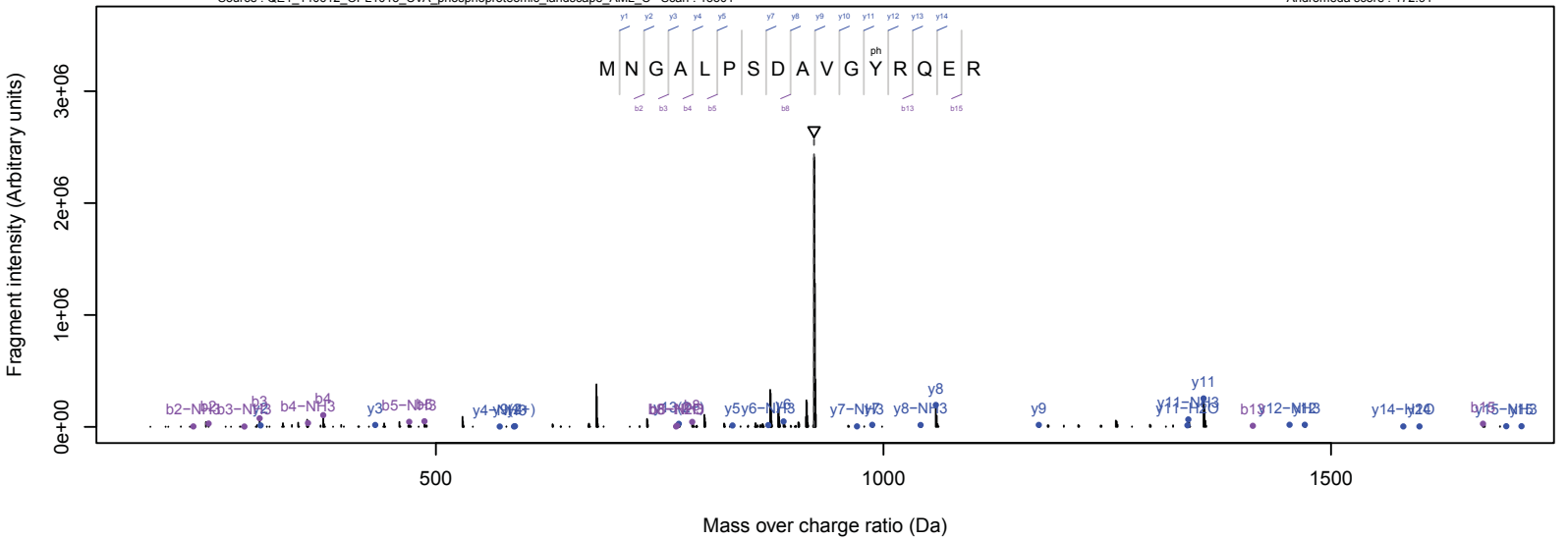
Andromeda score : 172.97



MNGALPSDAVGYRQER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15301

Andromeda score : 172.91

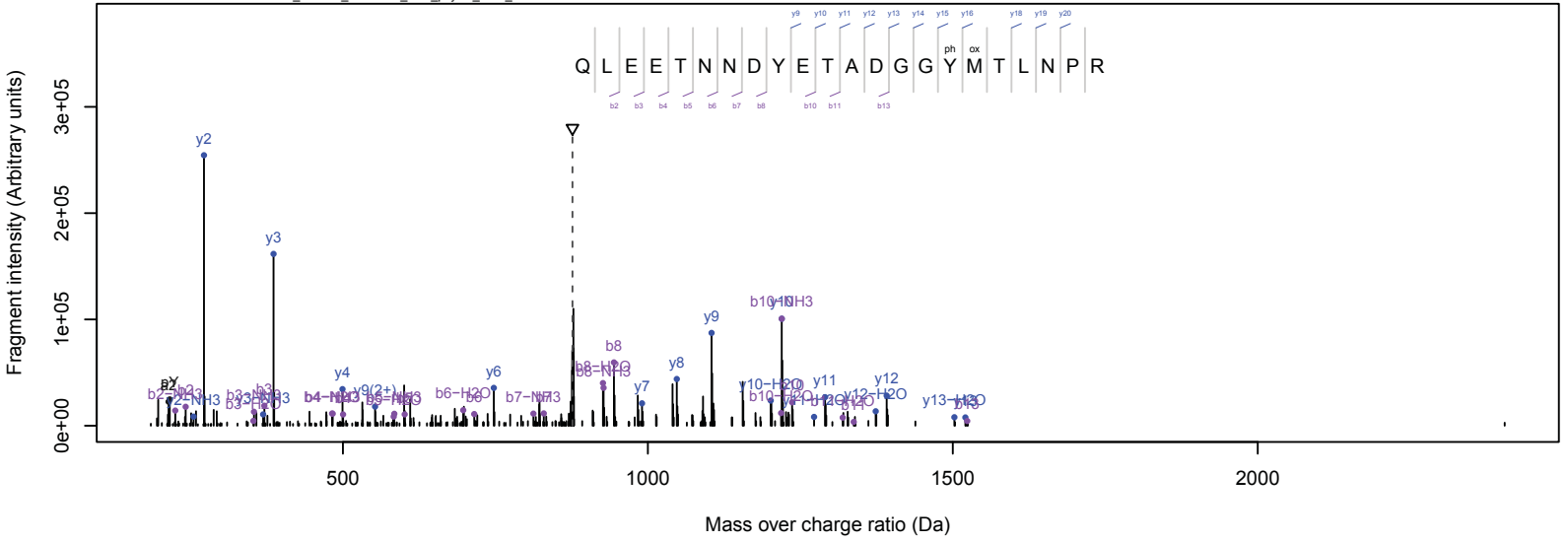




QLEETNNDYETADGGY(ph)M(ox)TLNPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 22042

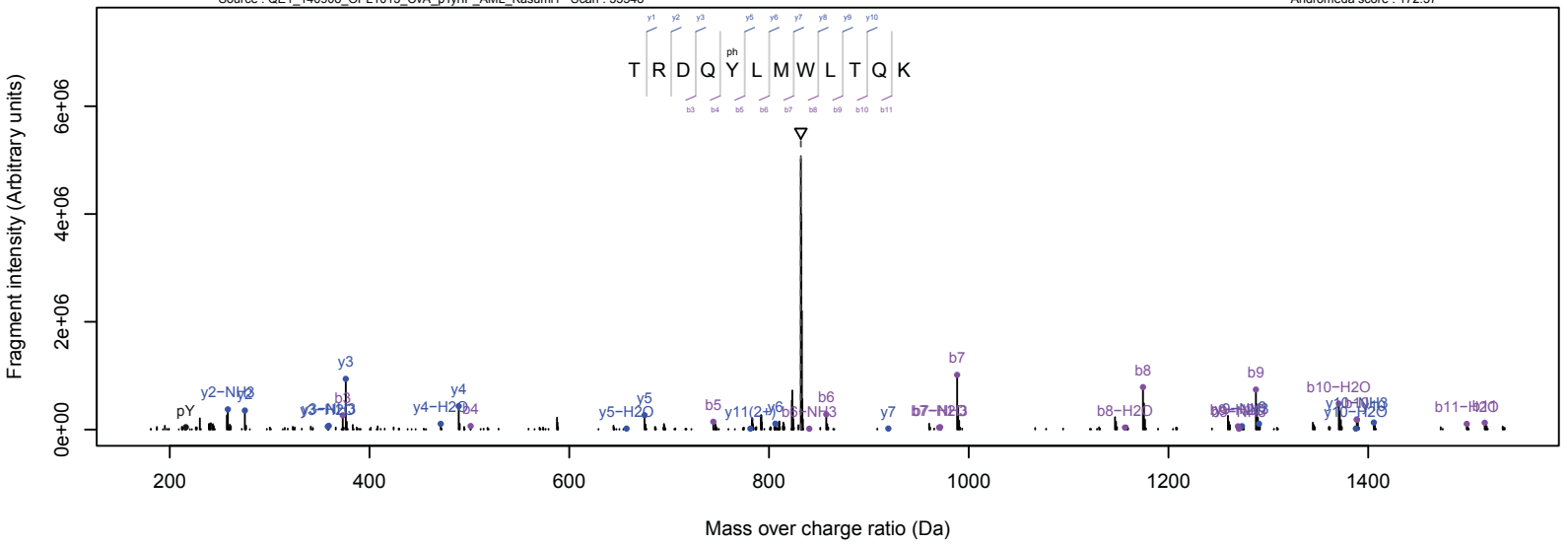
Andromeda score : 172.57



TRDQY(p)LMWLTQK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 33548

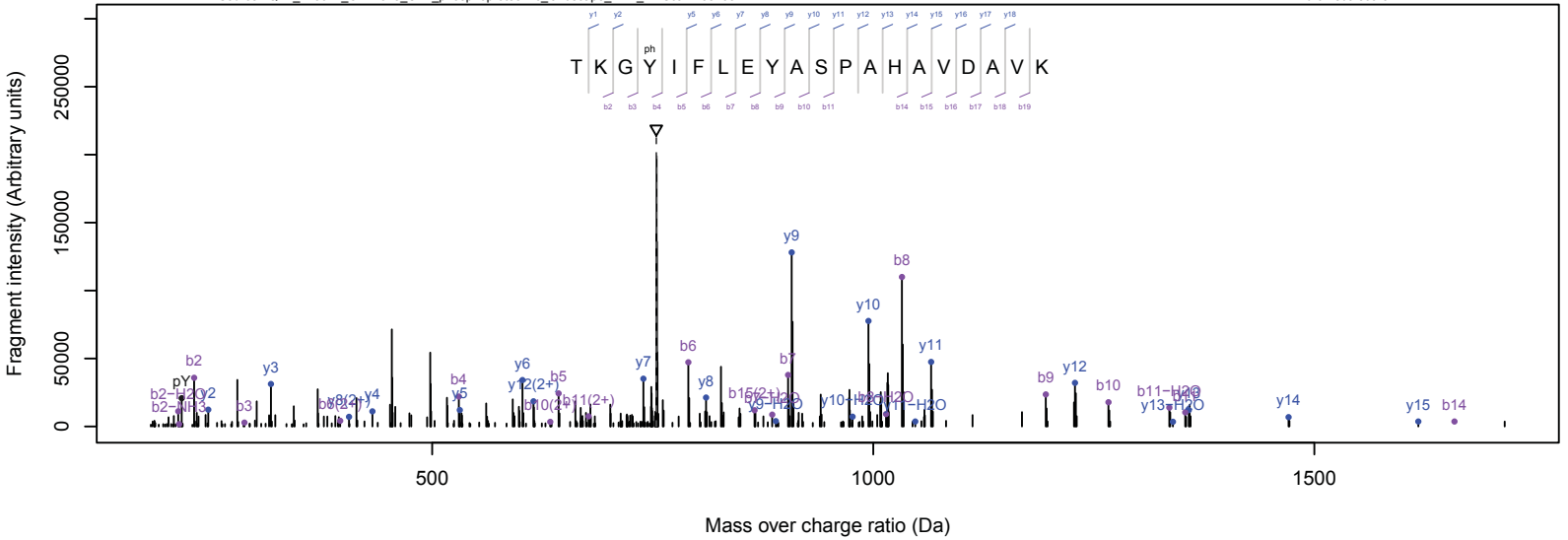
Andromeda score : 172.57



TKGY(p)IFLEYASPAHAVDAVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35435

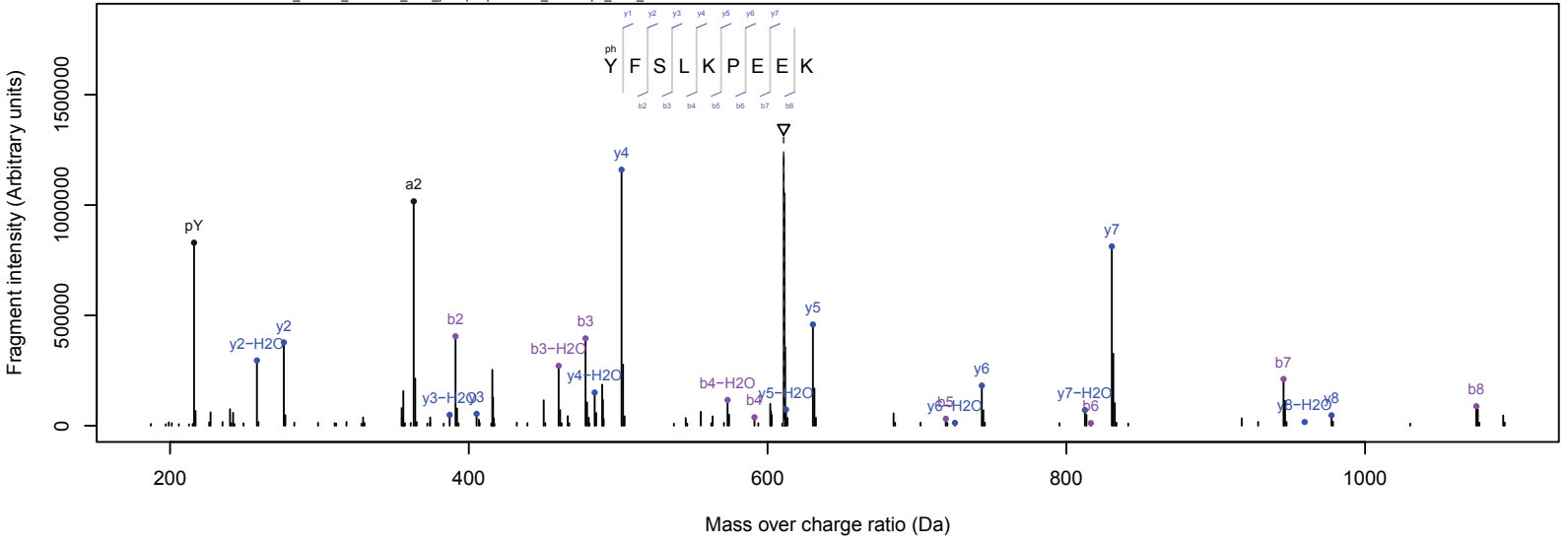
Andromeda score : 172.42



**Y(ph)FSLKPEEK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16543

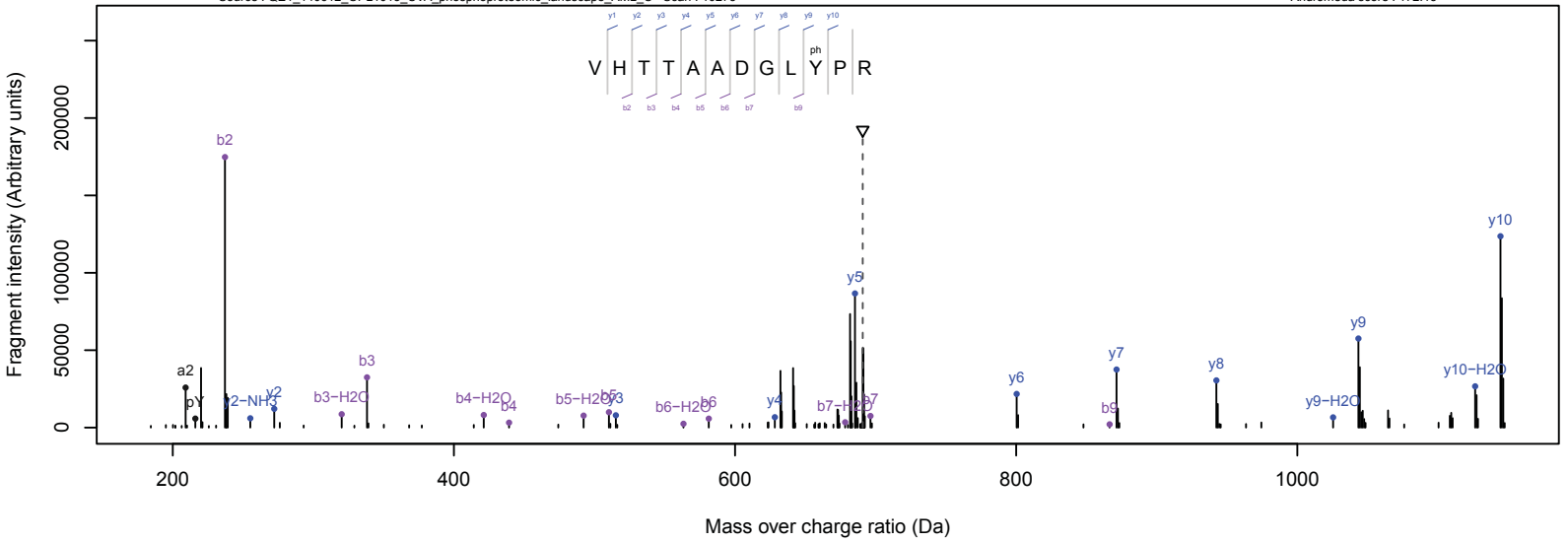
Andromeda score : 172.27



**VHTTAADGLY(ph)PR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10275

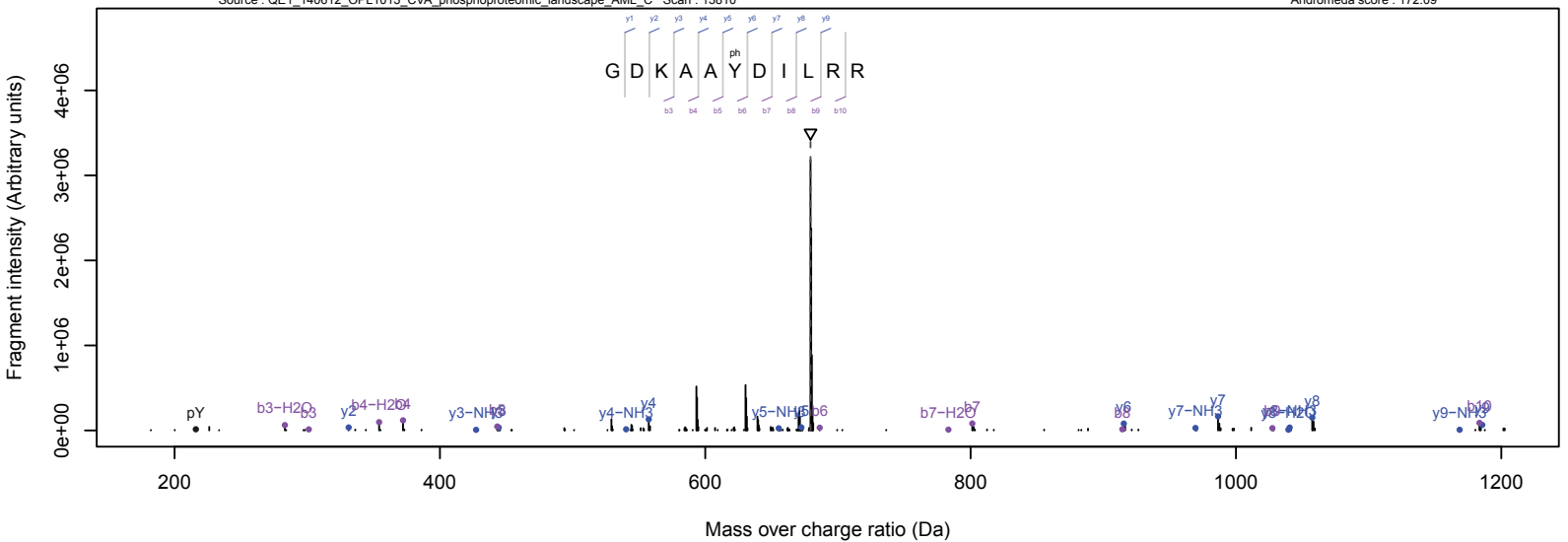
Andromeda score : 172.13



**GDKAAY(ph)DILRR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13810

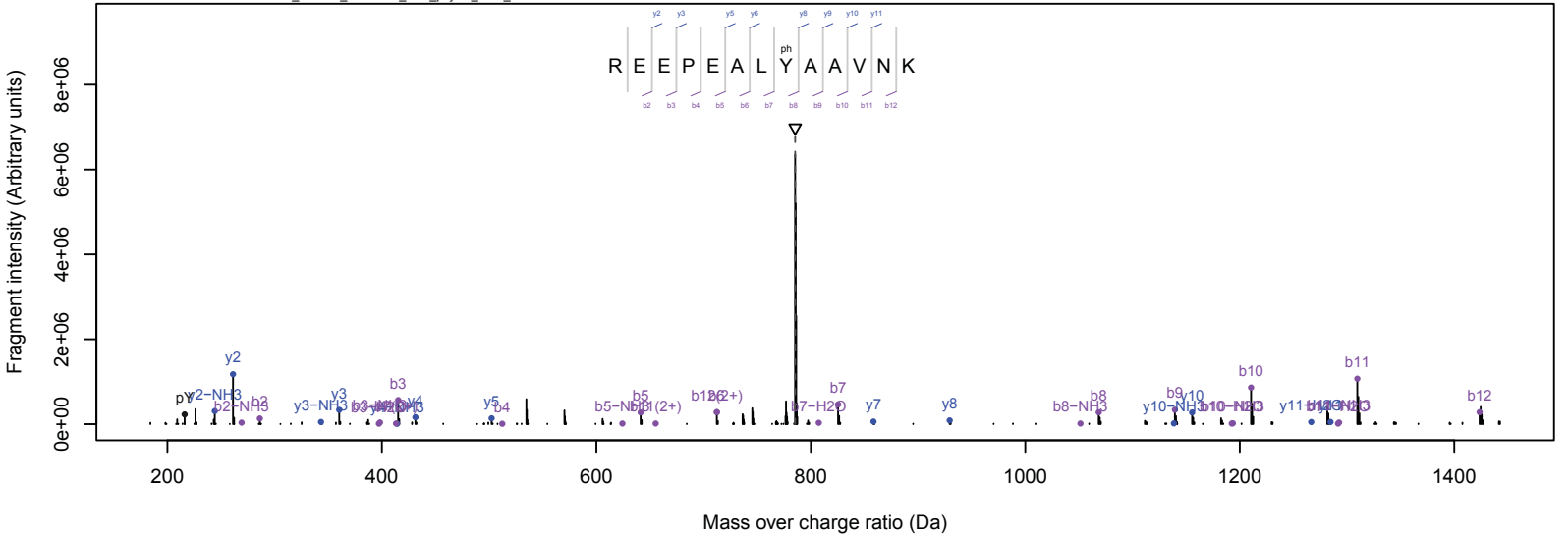
Andromeda score : 172.09



REEPEAL(ph)AAVNK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 19406

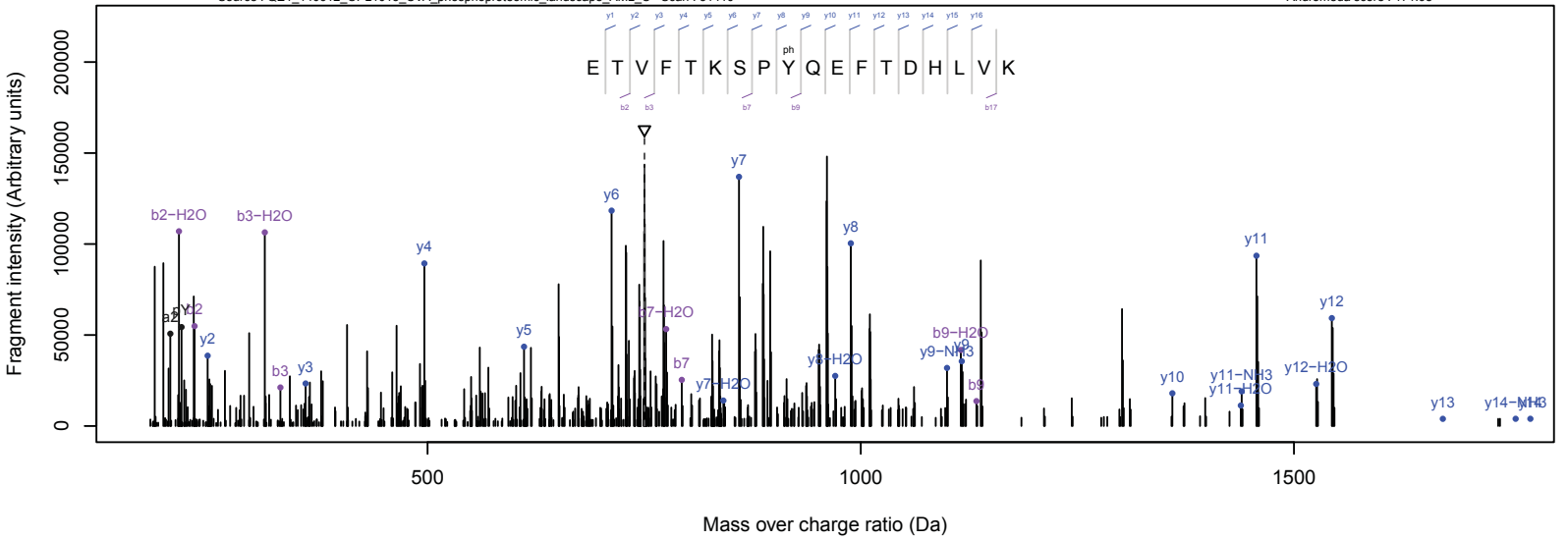
Andromeda score : 172.09



ETVFTKSPY(ph)QEFTDHLVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31410

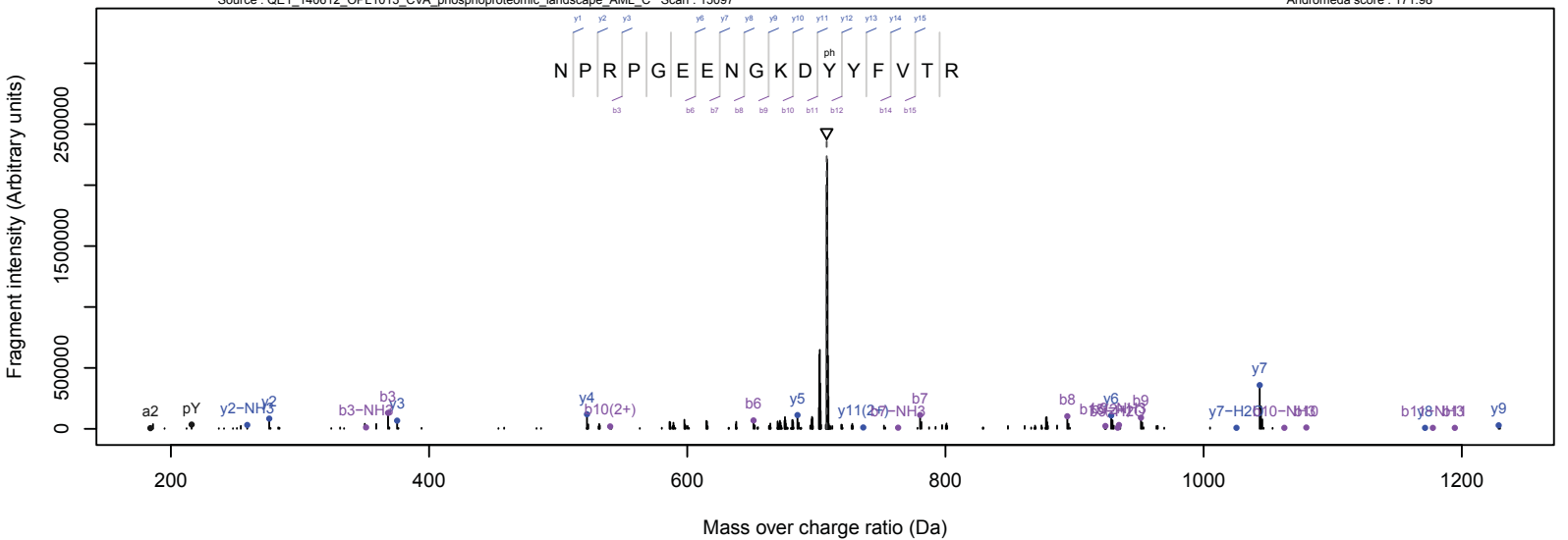
Andromeda score : 171.98



NPRPGEENGKDY(ph)YFVTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15097

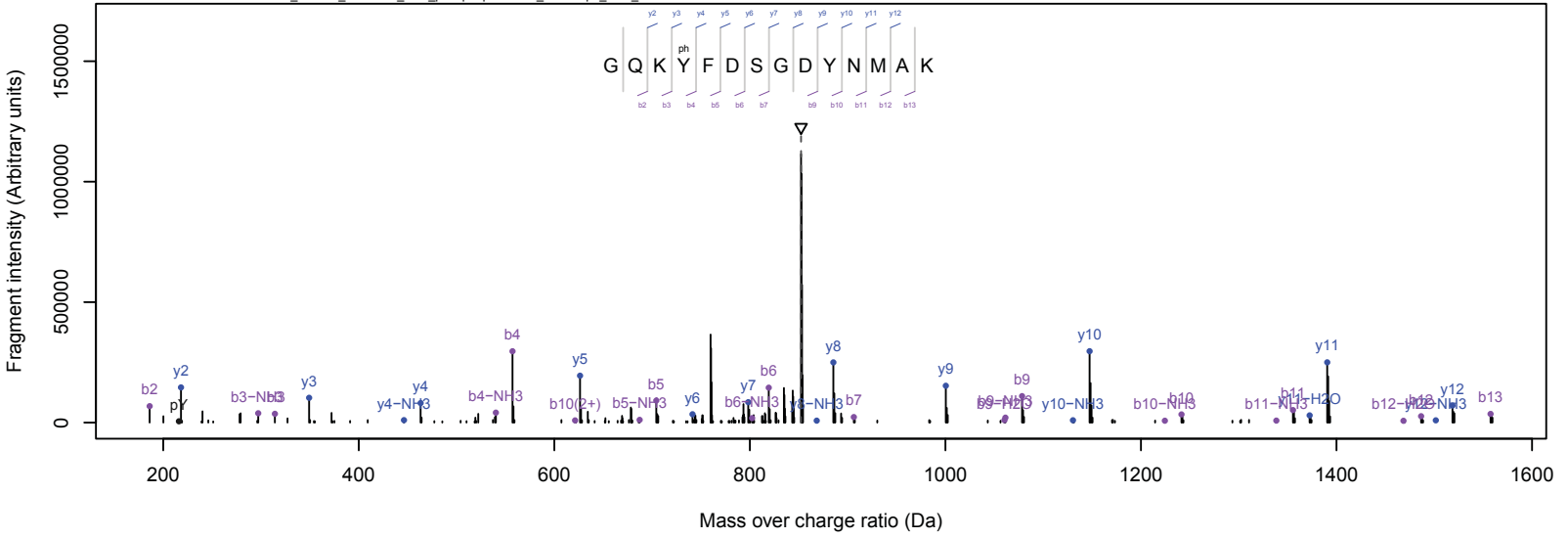
Andromeda score : 171.98



**GQKY(ph)FDSGDYNMAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14890

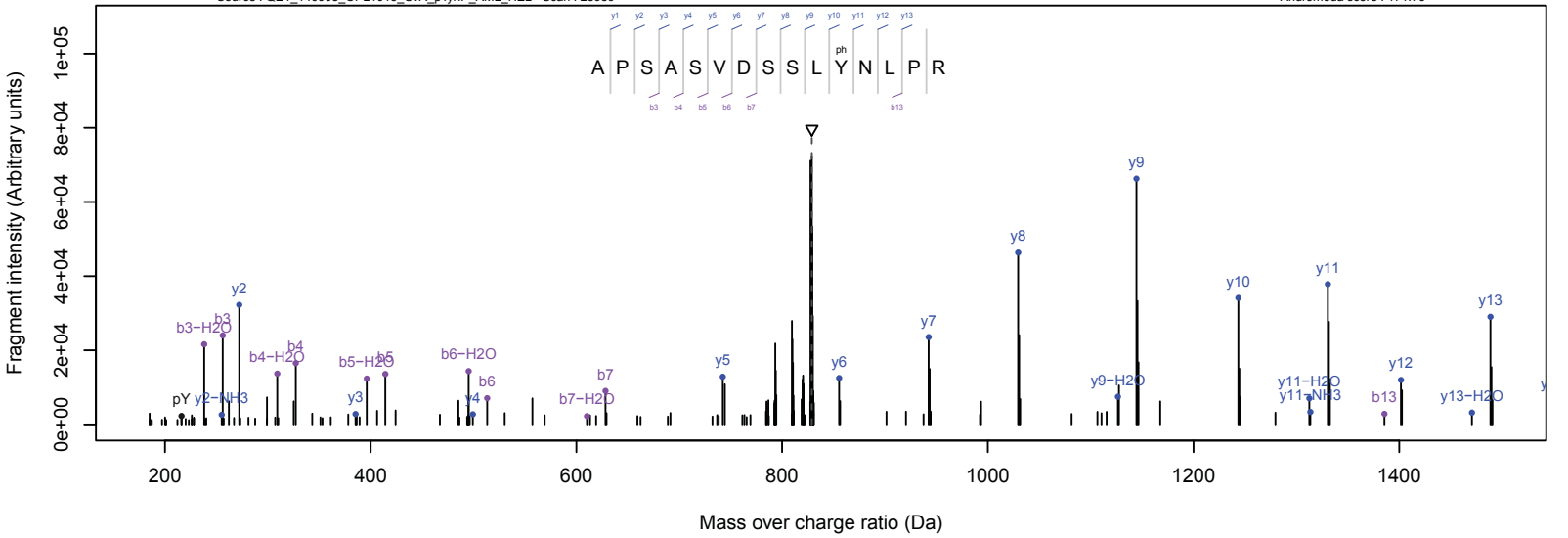
Andromeda score : 171.81



**APSASVDSLLY(ph)NLPR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 28989

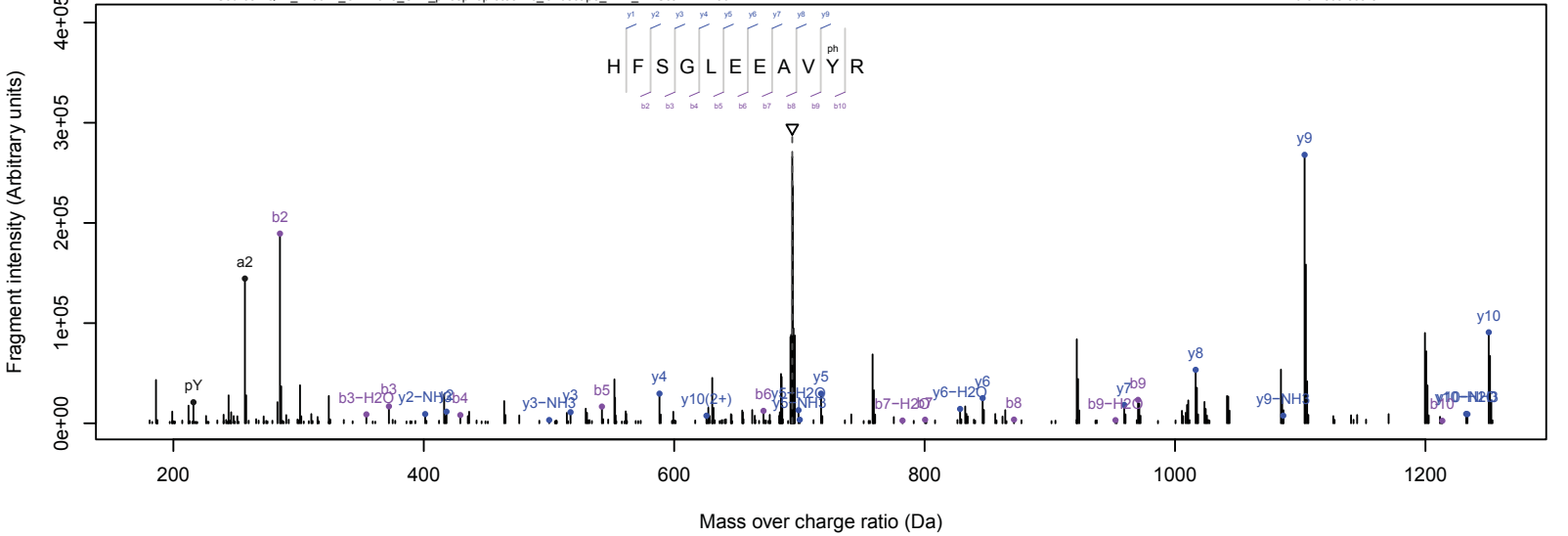
Andromeda score : 171.76



**HFSGLEEAVY(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21136

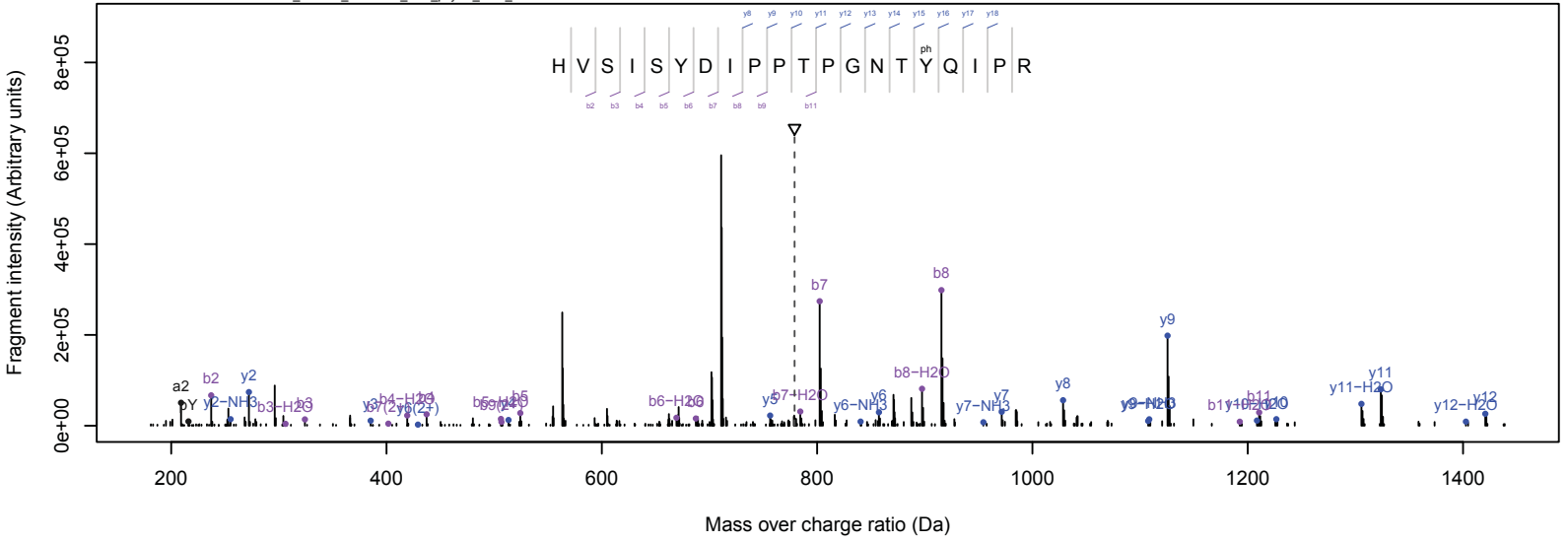
Andromeda score : 171.74



**HVSISYDIPPTPGNTY(ph)QIPR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32705

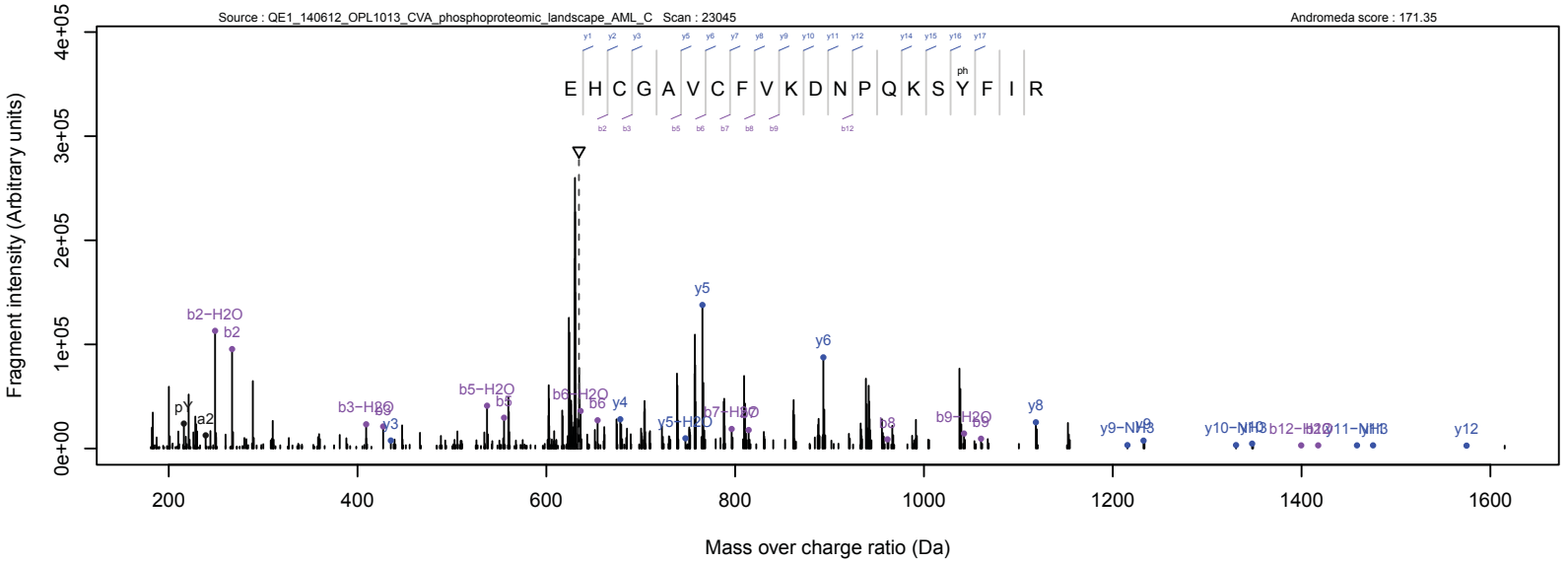
Andromeda score : 171.71



**EHC GAVCFVKDNPQKSY(ph)FIR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23045

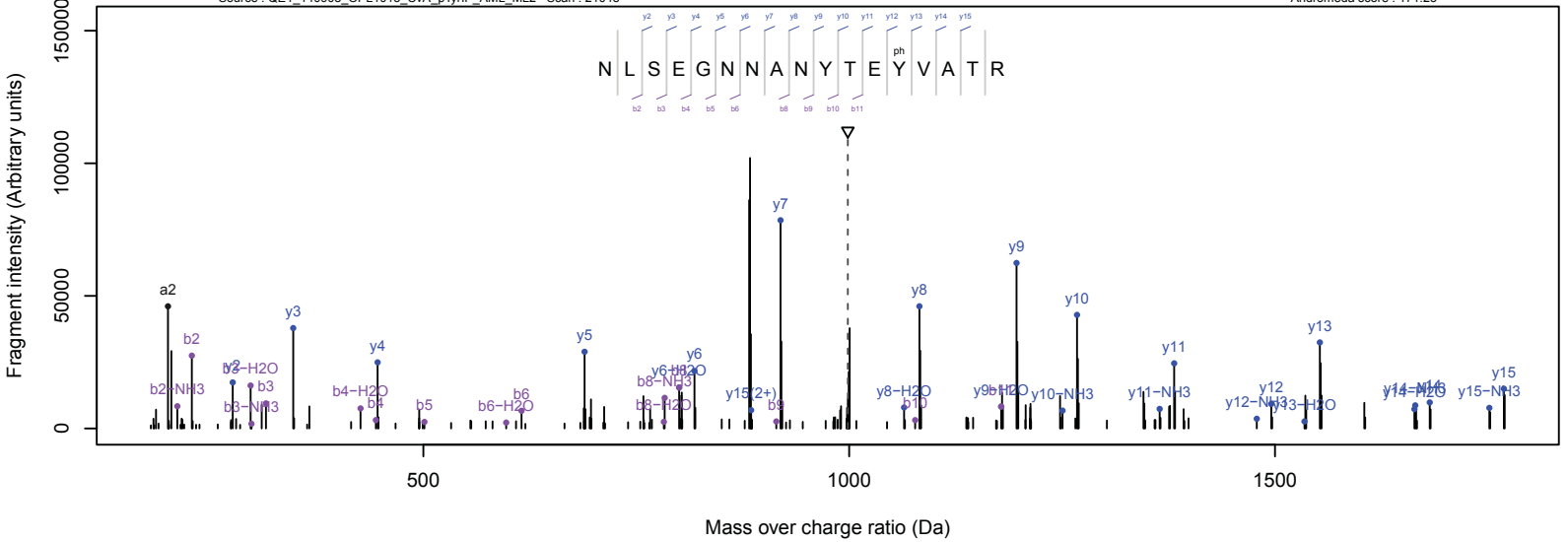
Andromeda score : 171.35



**NLSEGNANYTEY(ph)VATR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 21048

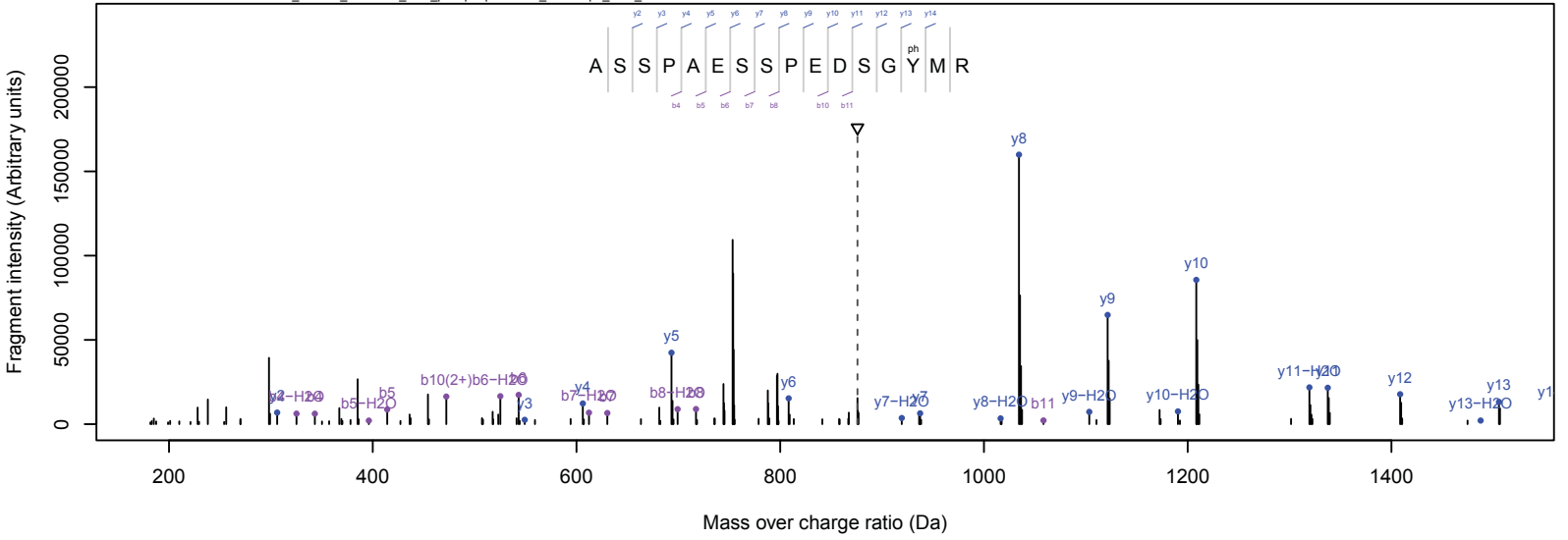
Andromeda score : 171.25



**ASSPAESSPEDSGY<sup>ph</sup>MR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 8965

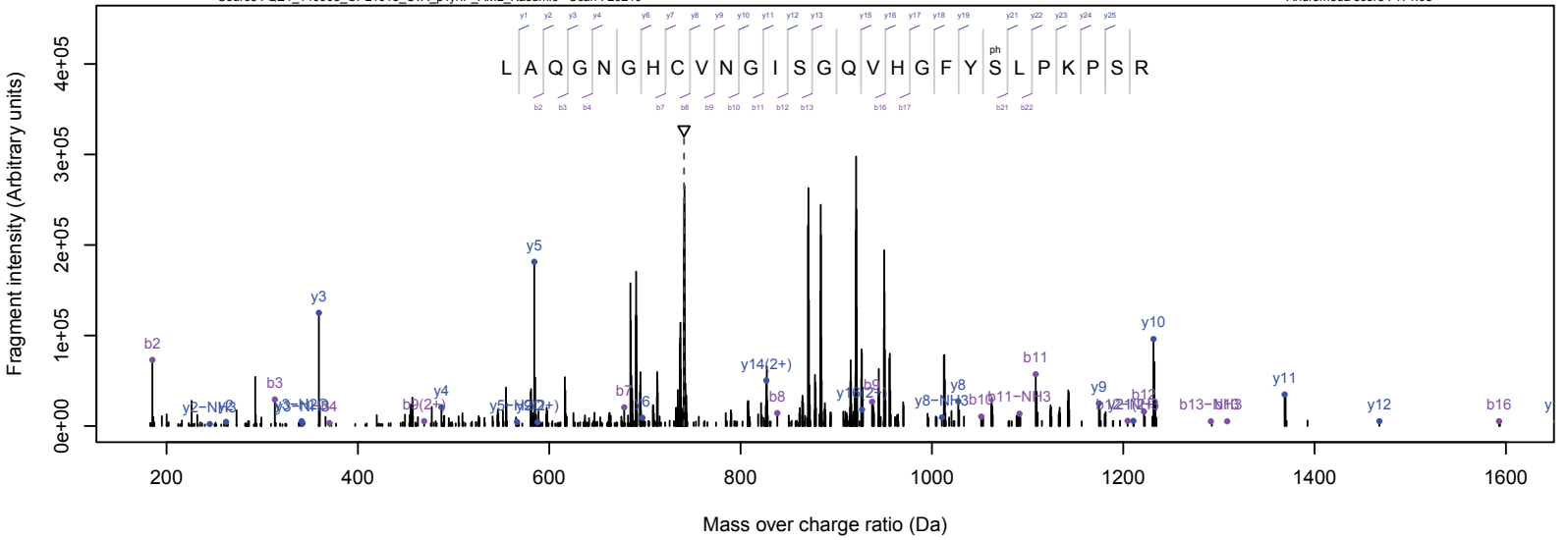
Andromeda score : 171.08



**LAQGNHCVNGISGVHGFYS<sup>ph</sup>LPKPSR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 26219

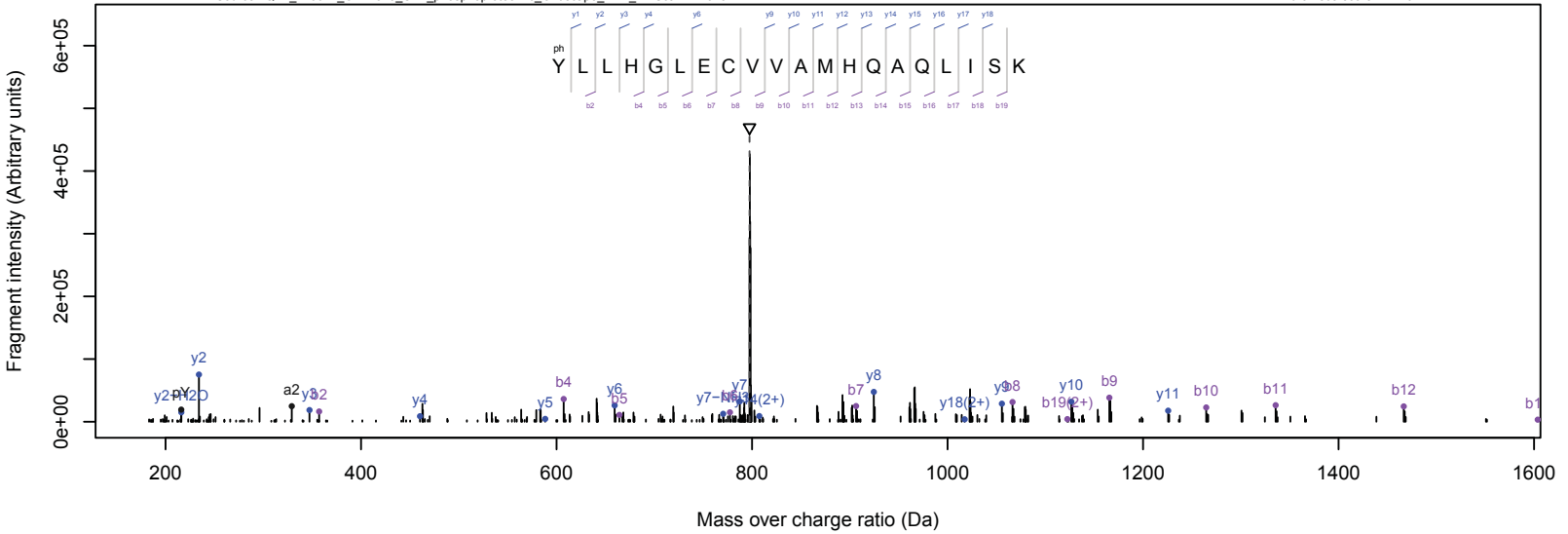
Andromeda score : 171.08



**Y<sup>ph</sup>LLHGLECVVAMHQAQLISK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42025

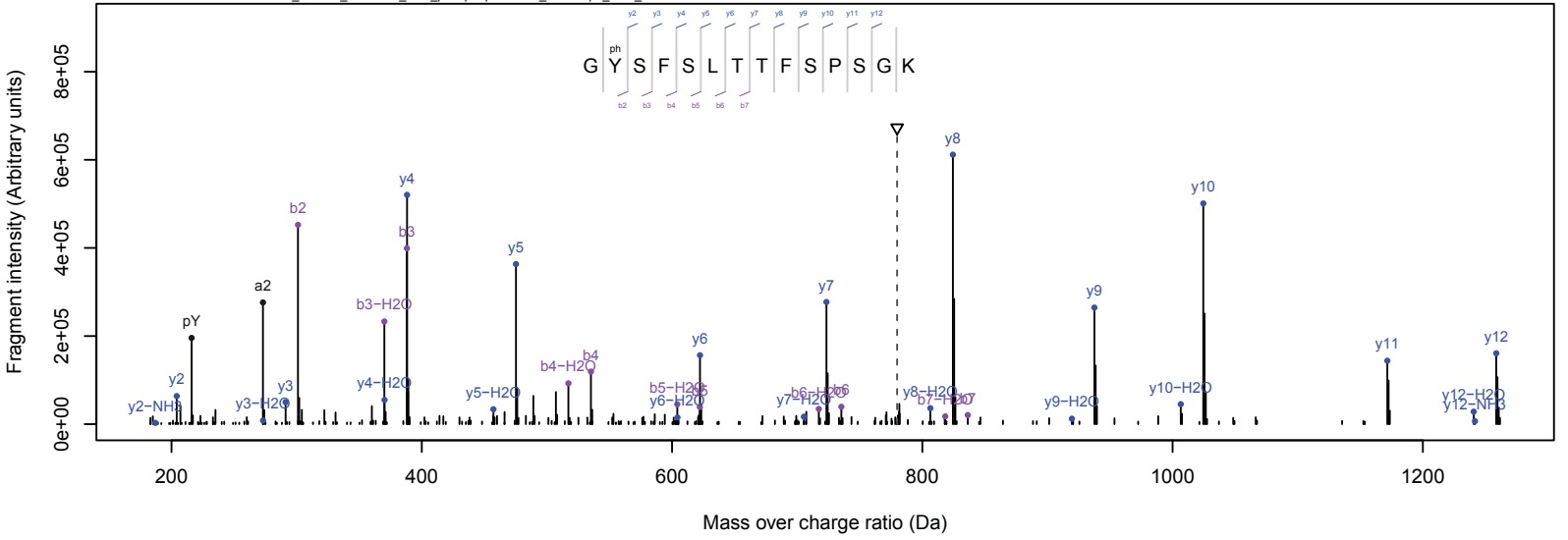
Andromeda score : 171.07



**GY(ph)SFLTTFSPGK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36767

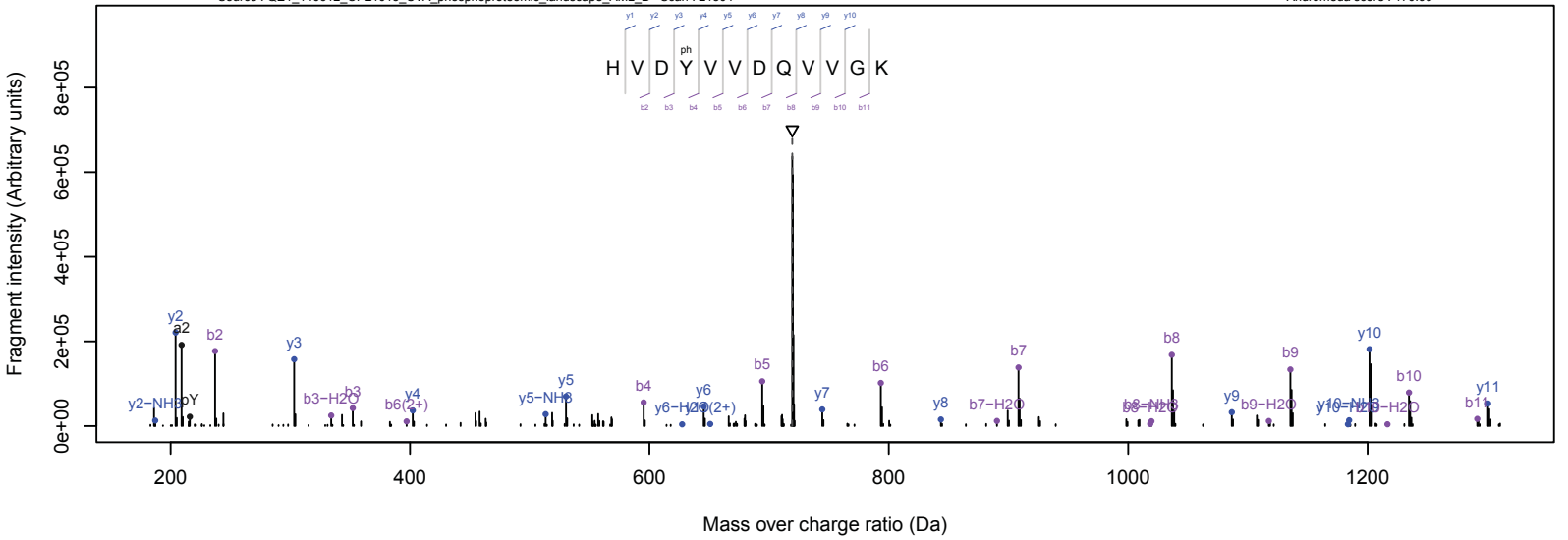
Andromeda score: 170.95



**HVDY(ph)VVDQVVGK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 21304

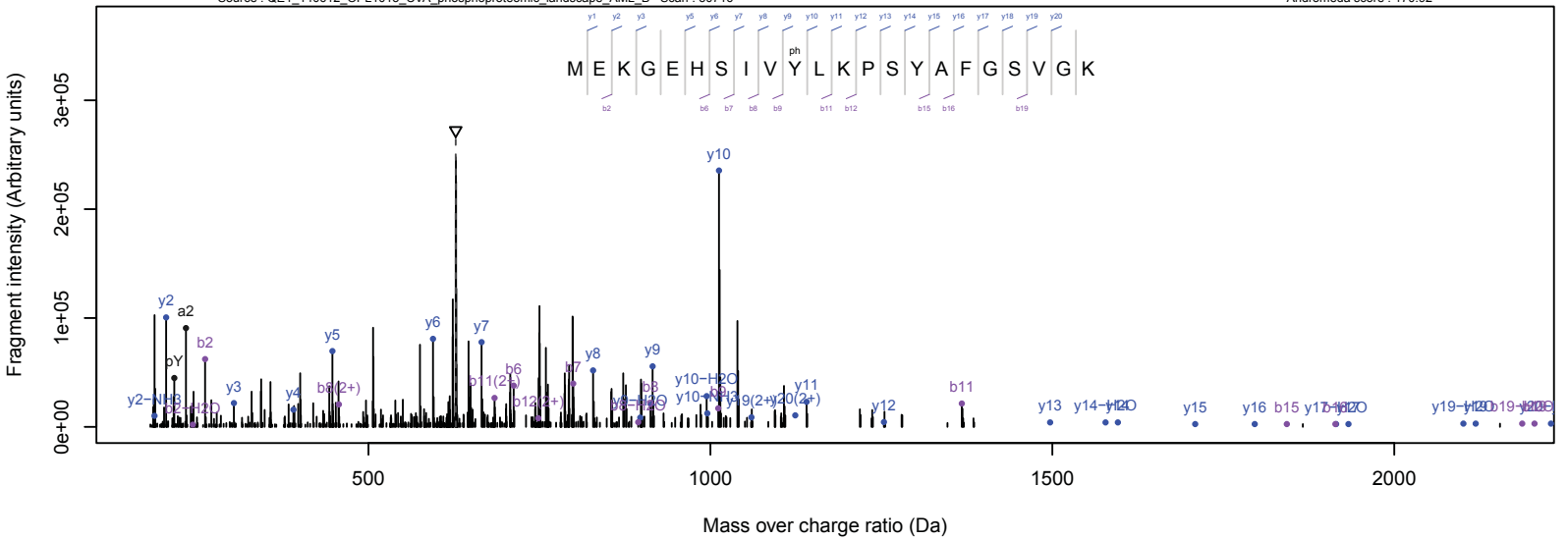
Andromeda score: 170.95



**MEKGEHSIVY(ph)LKPSYAFGSVVGK-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 30716

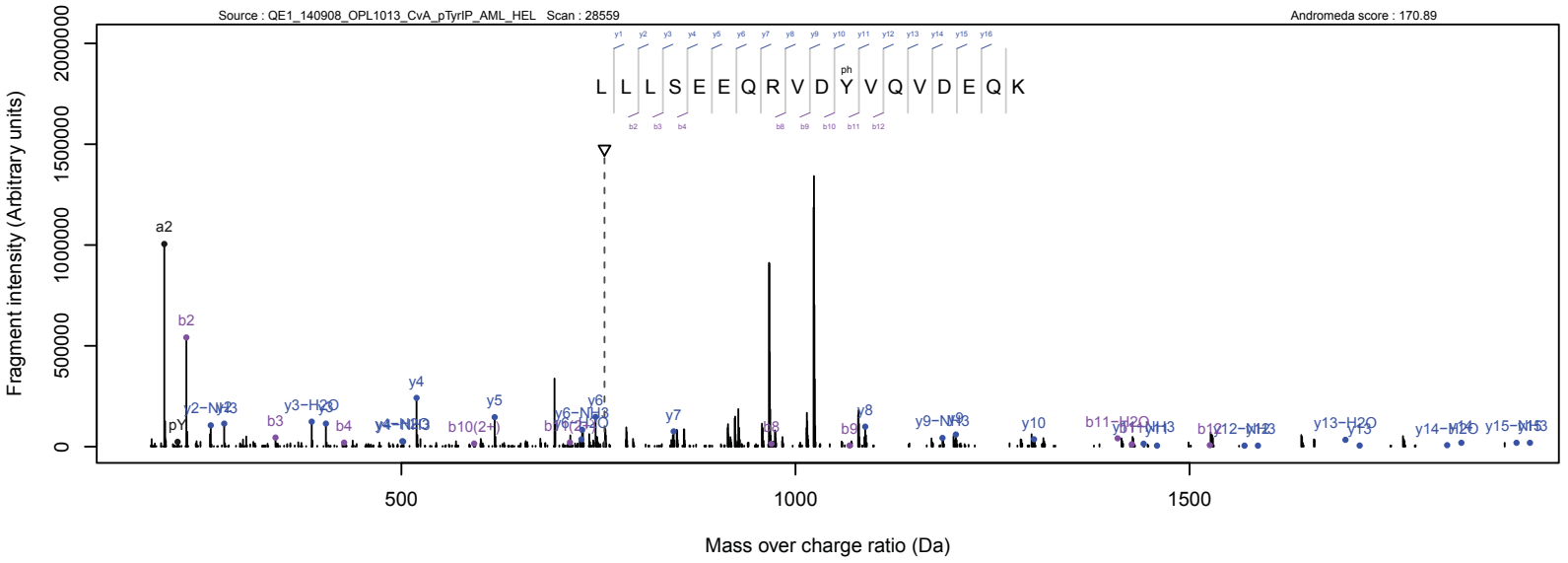
Andromeda score: 170.92



LLLSEEQRDY(ph)VQVDEQK-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan: 28559

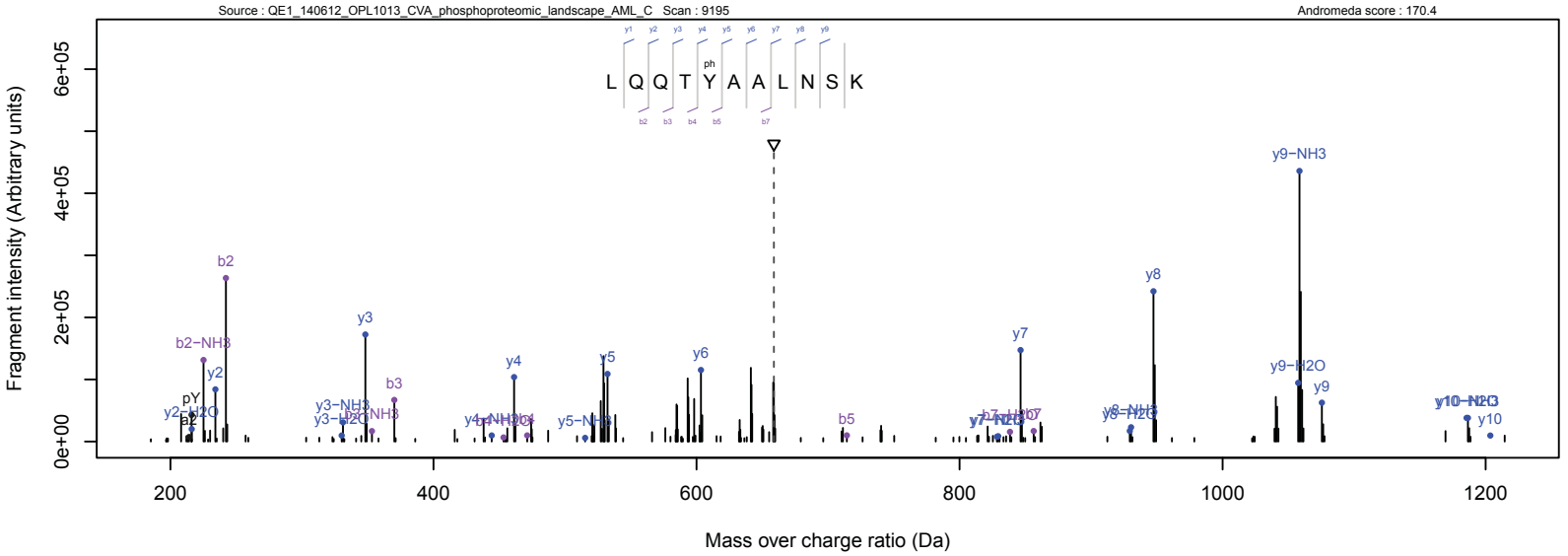
Andromeda score: 170.89



LQTTY(ph)AALNSK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 9195

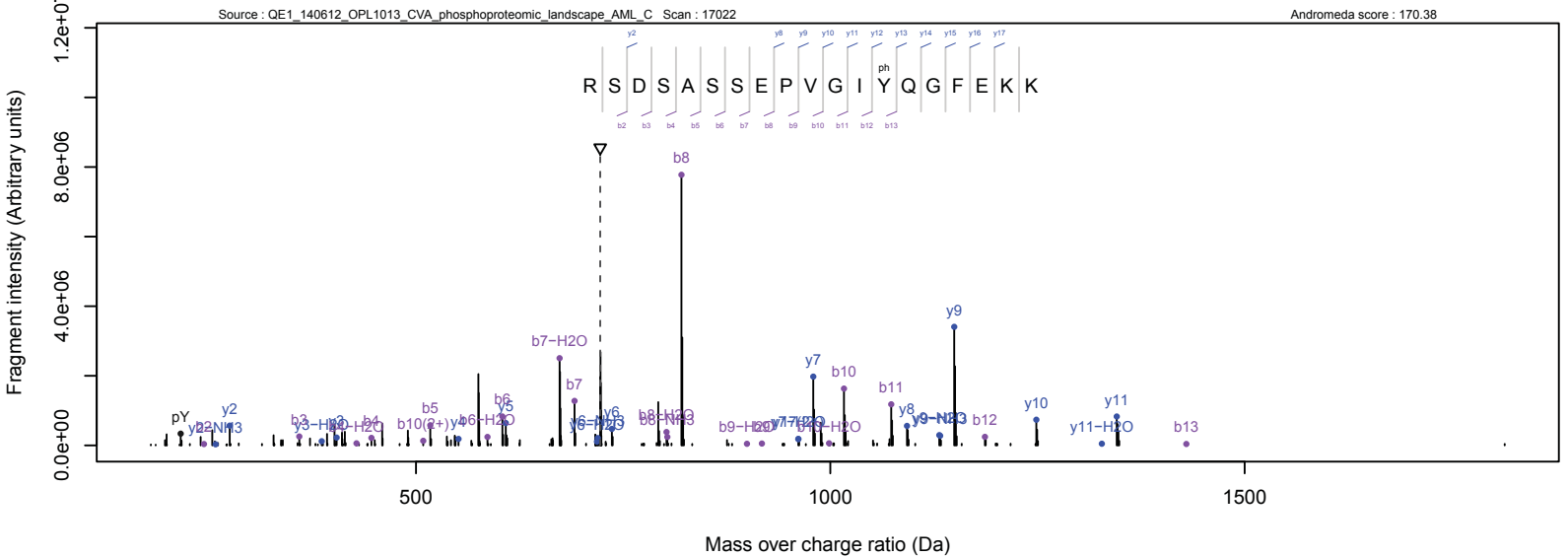
Andromeda score: 170.4



RSDSASSEPVGIYQGFEEK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17022

Andromeda score: 170.38



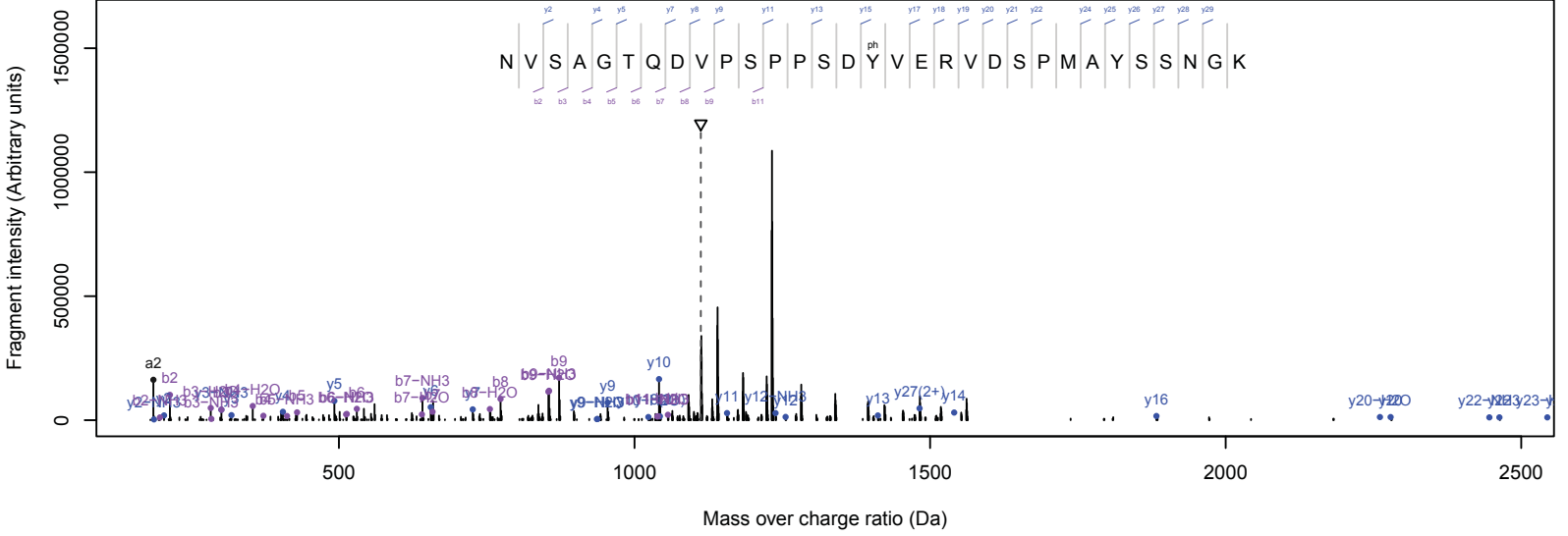




NVSAGTQDVPPSPDY(ph)VERVDSPMAYSSNGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30666

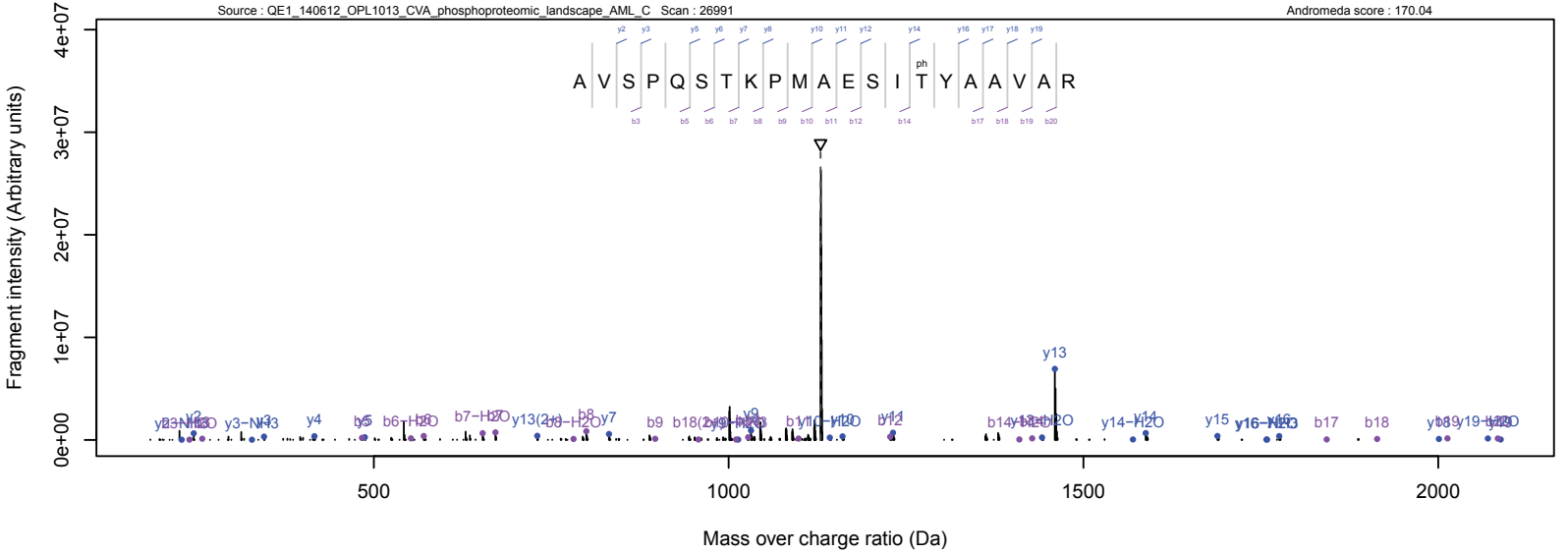
Andromeda score : 170.11



AVSQSTKPMASIT(ph)YAAVAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26991

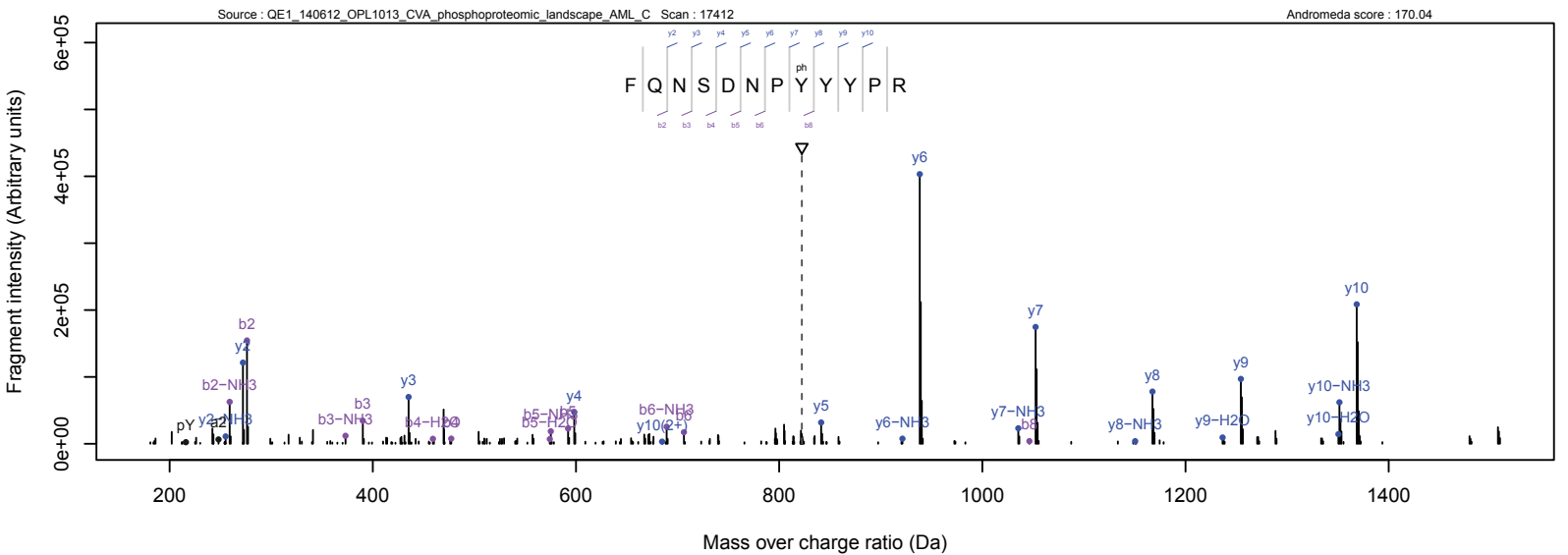
Andromeda score : 170.04



FQNSDNPYYYP(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17412

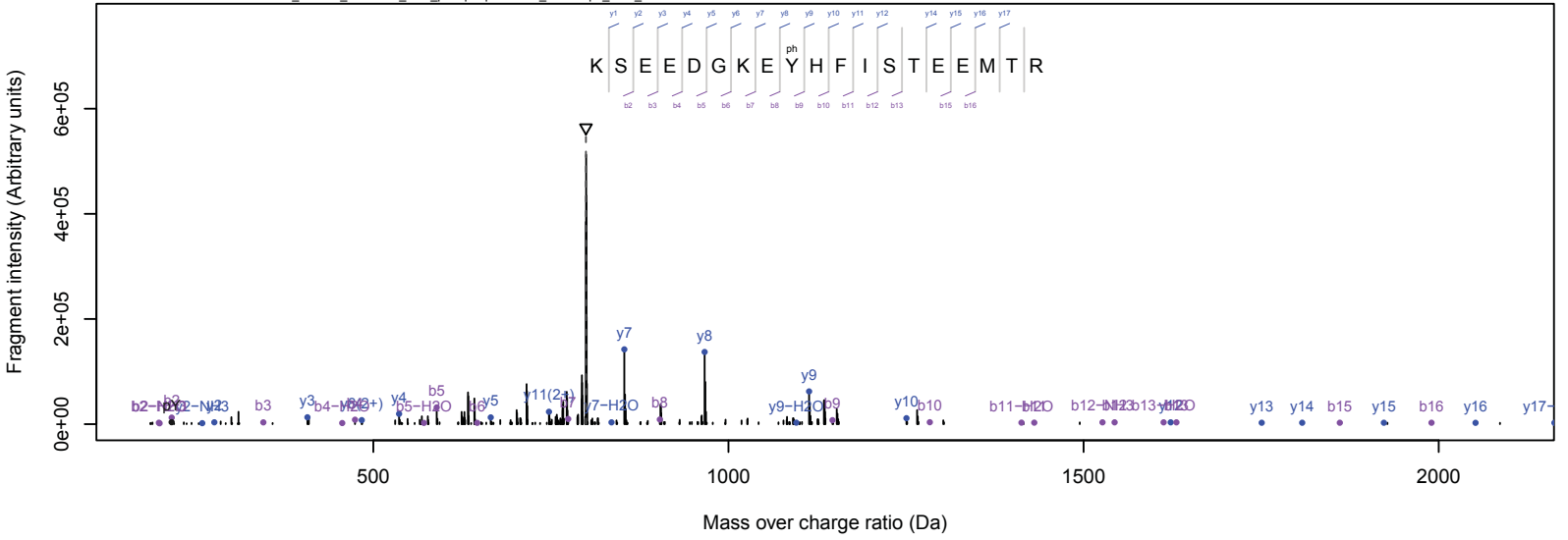
Andromeda score : 170.04



KSEEDGKEY(ph)HFISTEEMTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16570

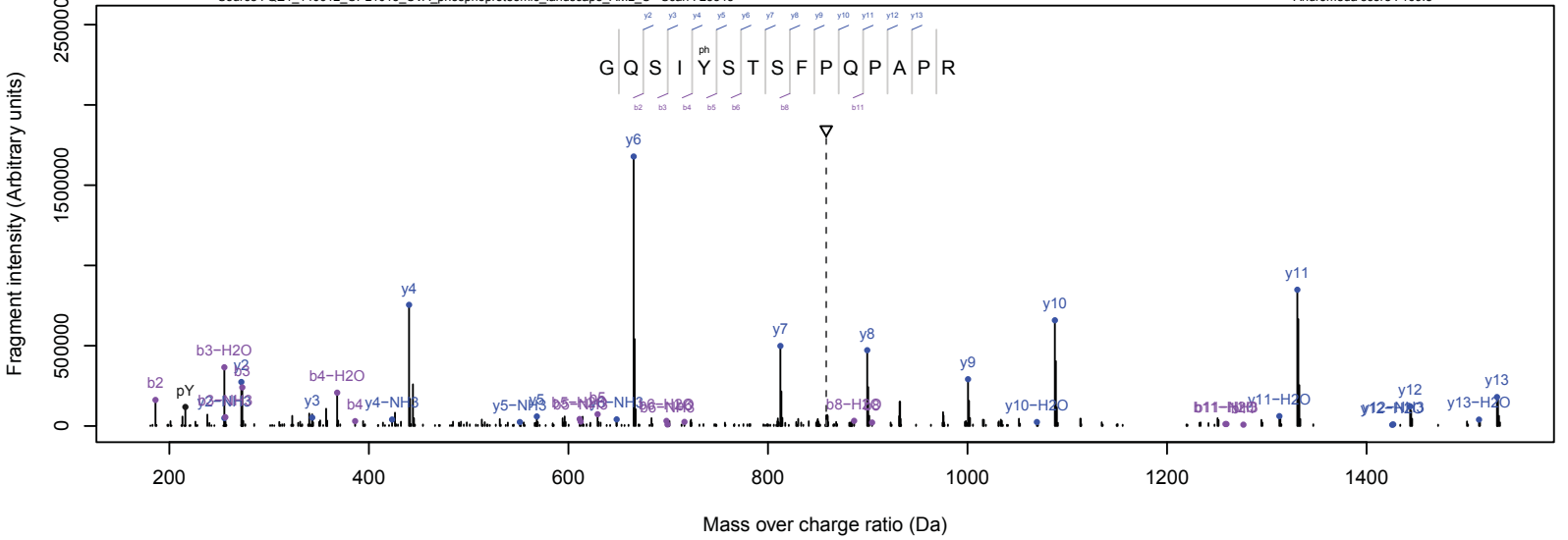
Andromeda score : 169.9



GQSIY(ph)STSFQPAPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23343

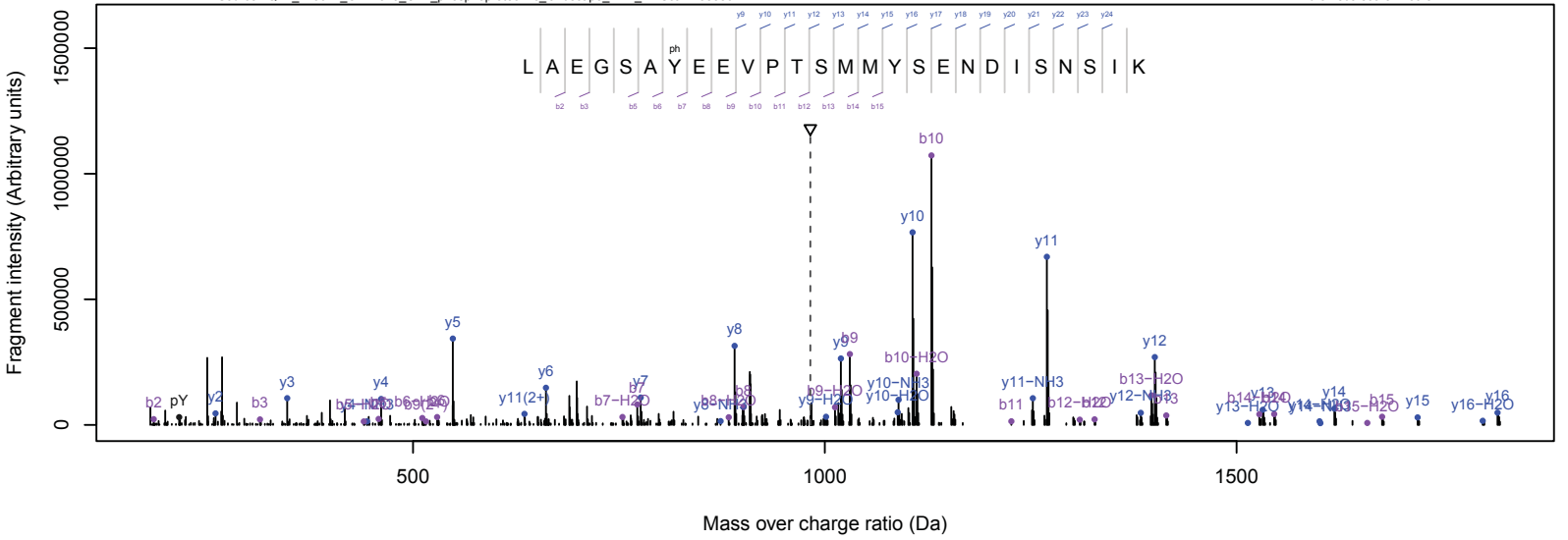
Andromeda score : 169.8



LAEGSAY(ph)EEVPTSMMYSENDISNSIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38886

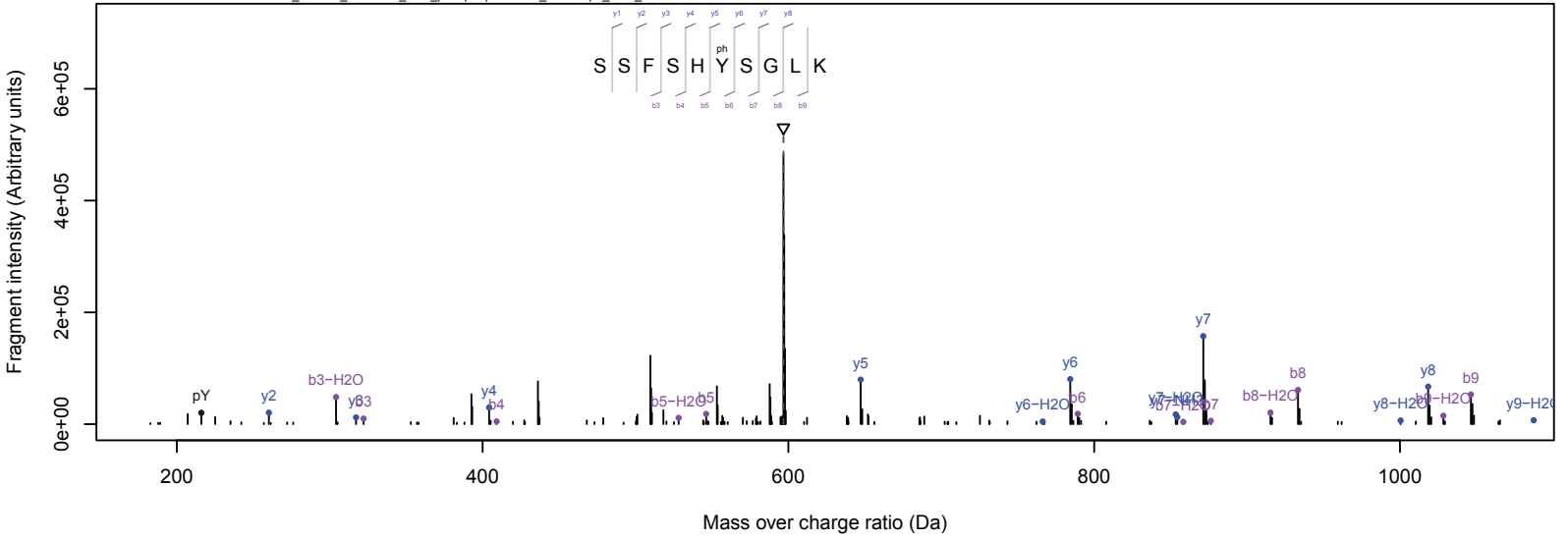
Andromeda score : 169.8



SSFSHY(ph)SGLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10215

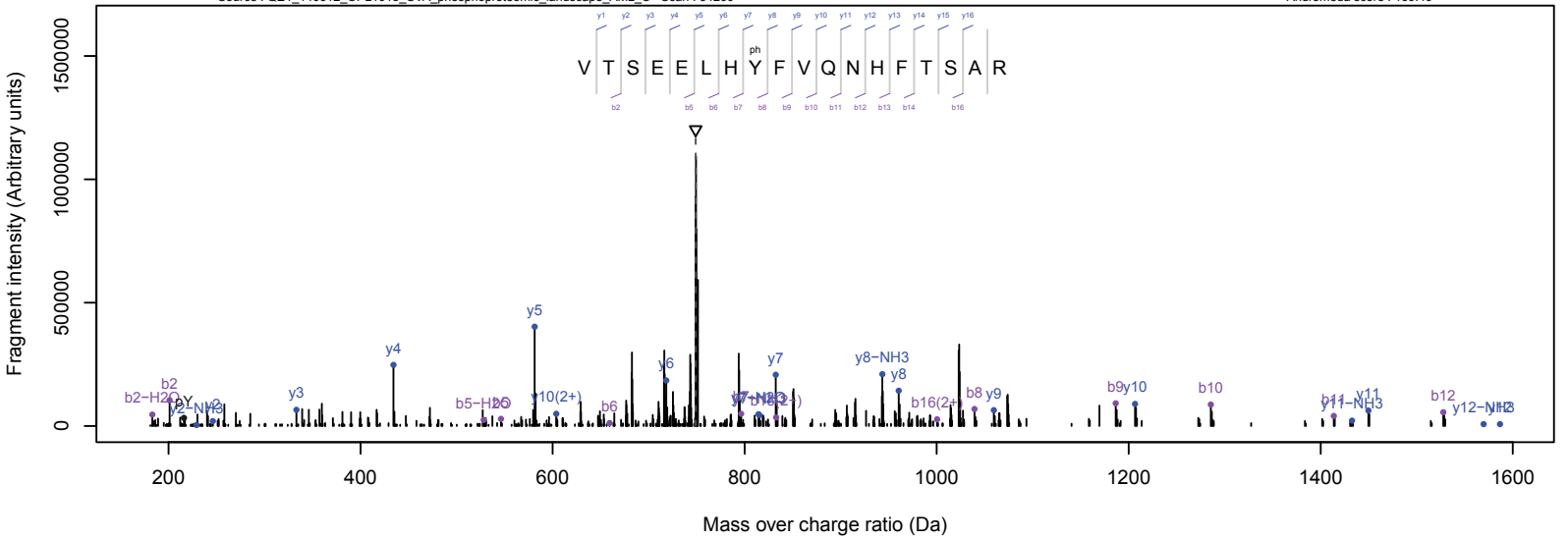
Andromeda score : 169.65



VTSEELHY(ph)FVQNHTSAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34260

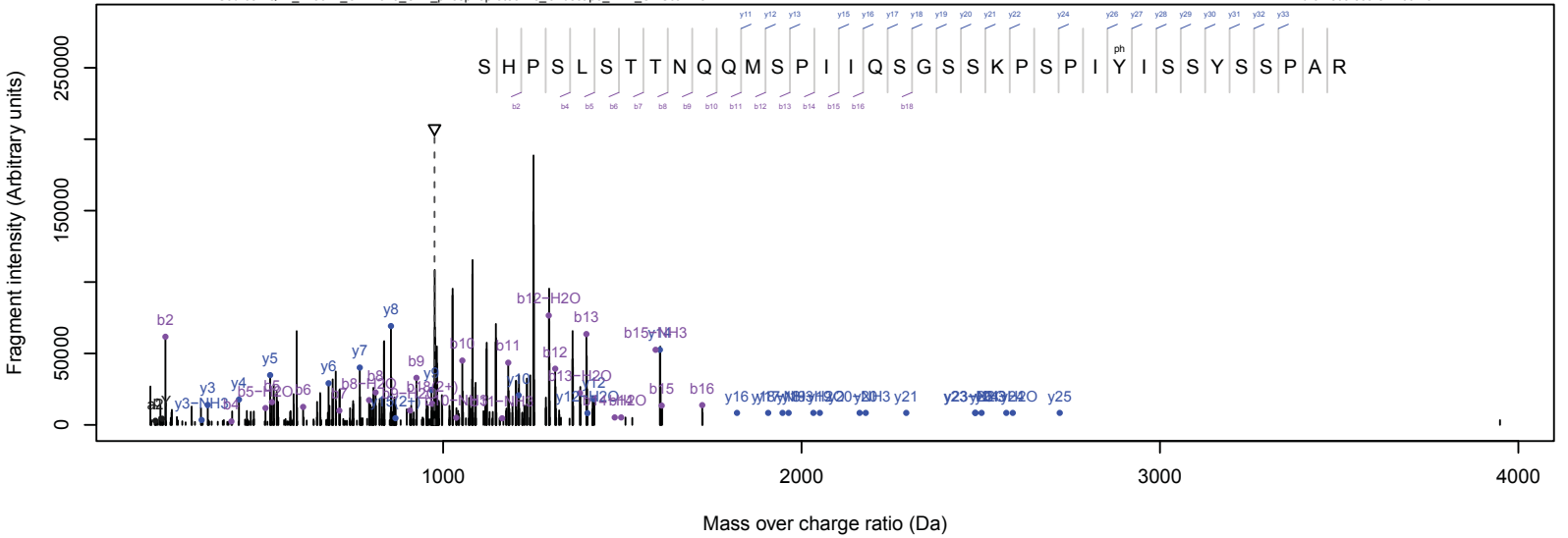
Andromeda score : 169.49



SHPSLSTTNQQMSP(i)IQSGSSKPSPIY(i)ISSYSSPAR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32724

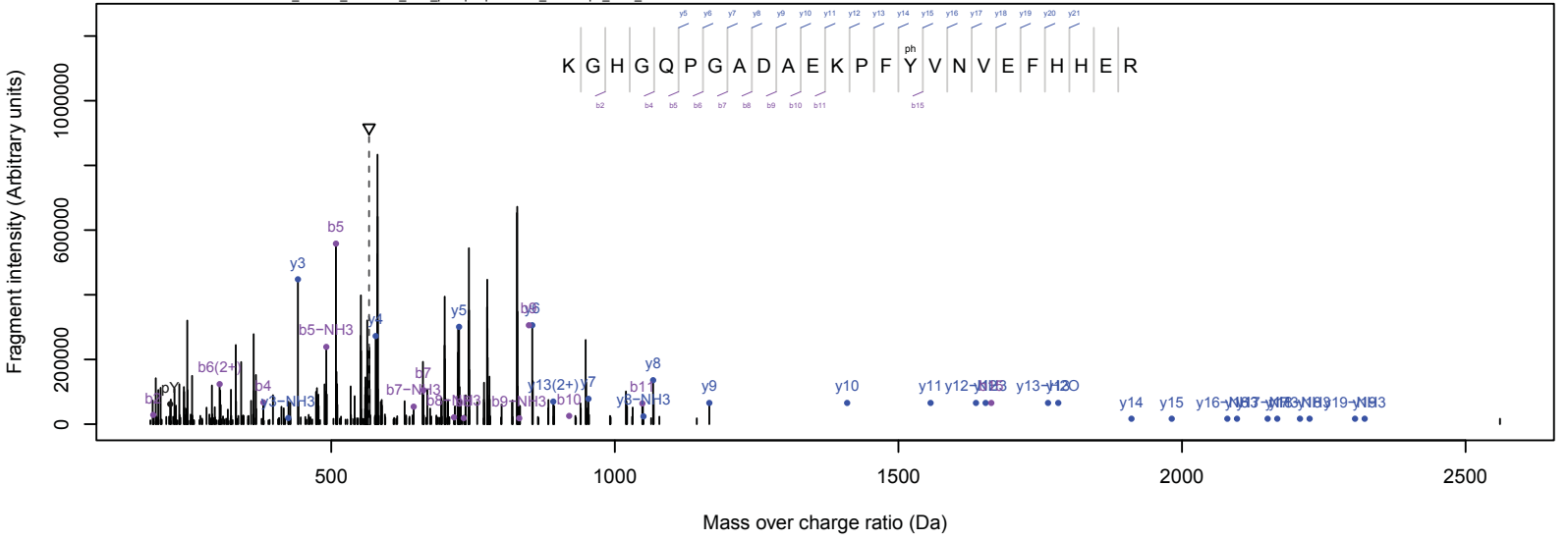
Andromeda score : 169.26



KGHGQPGADA EKPFYVNV E F H H E R

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 20743

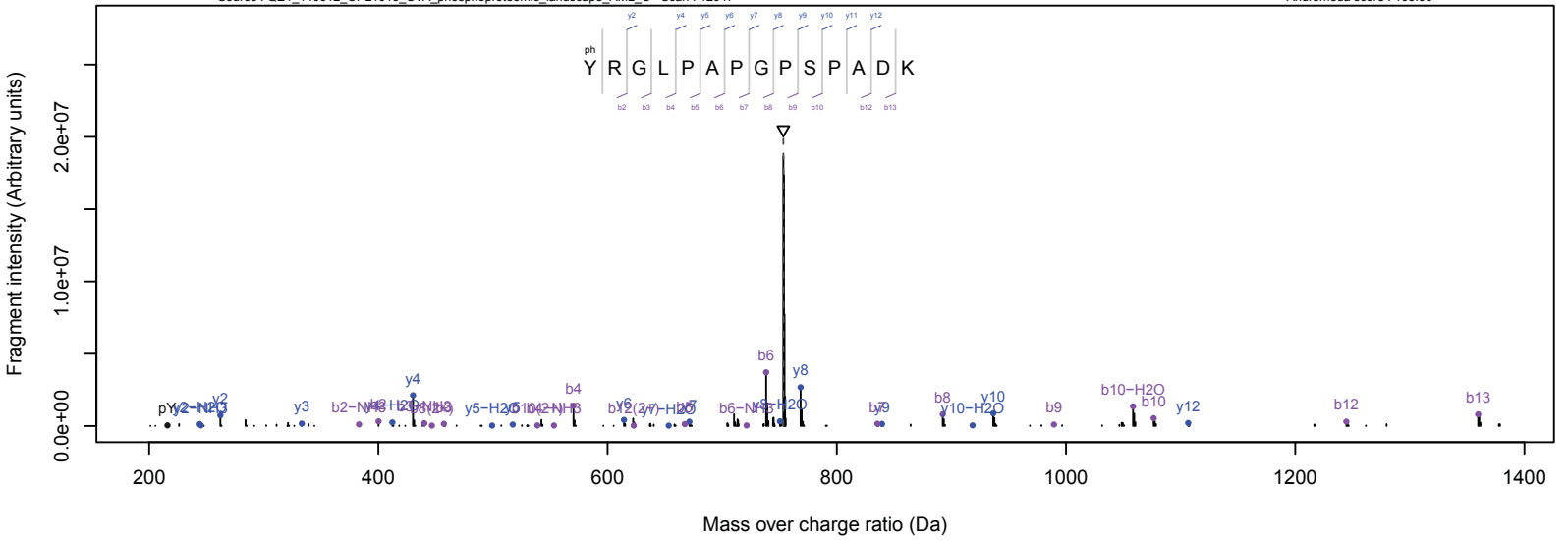
Andromeda score : 169.08



Y(ph)RGLPAPGPS PADK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12947

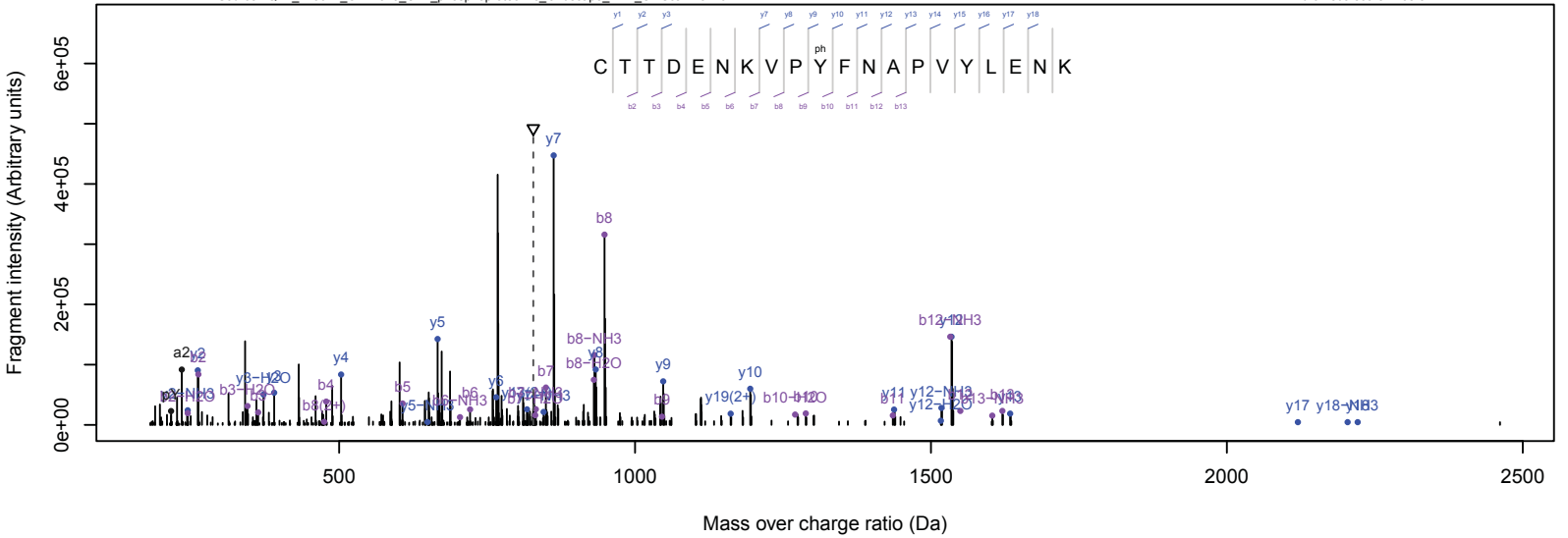
Andromeda score : 168.93



CTTDENKVPYFNAPVYLENK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34292

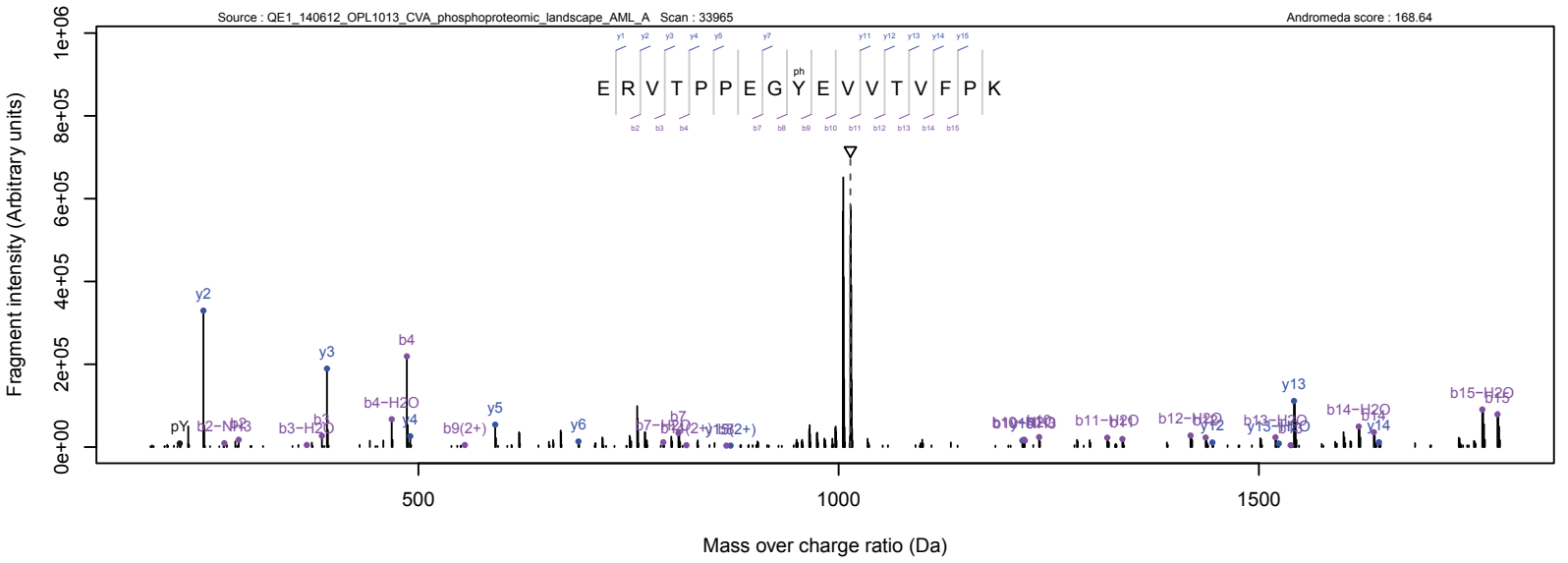
Andromeda score : 168.81



ERVTPPEGY(ph)EVVTVFPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33965

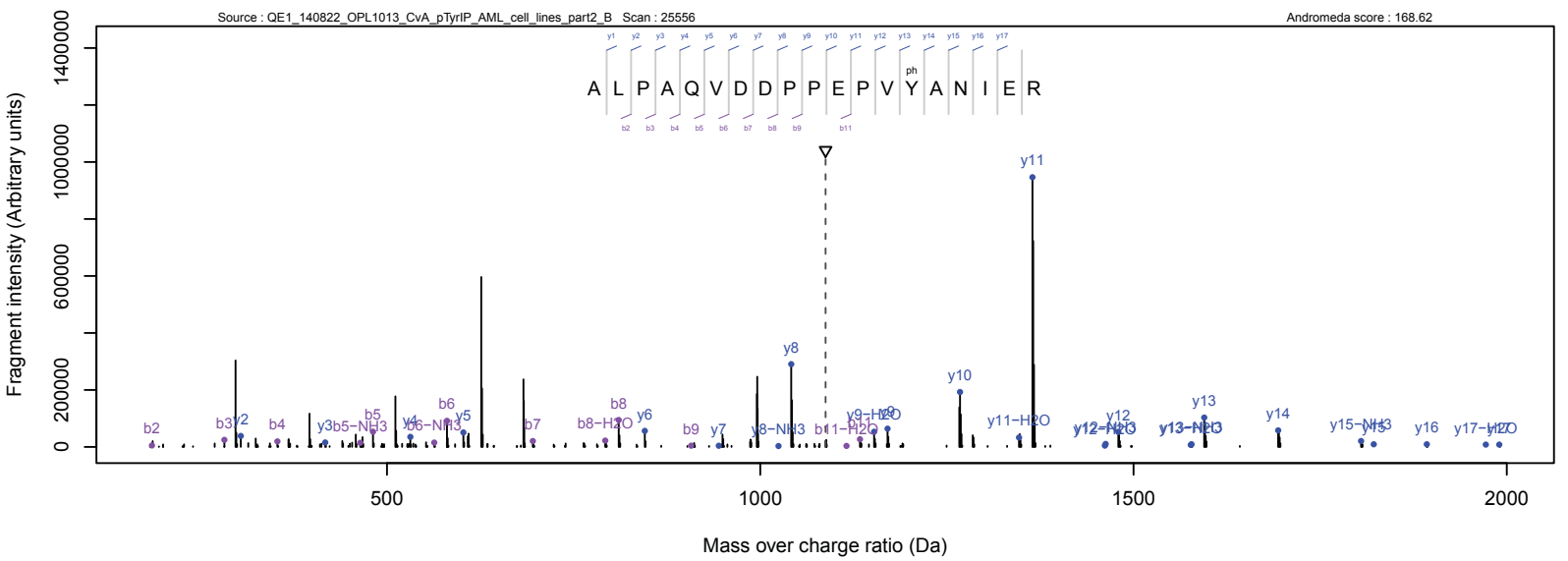
Andromeda score : 168.64



ALPAQVDDPPEV(ph)ANIER-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 25556

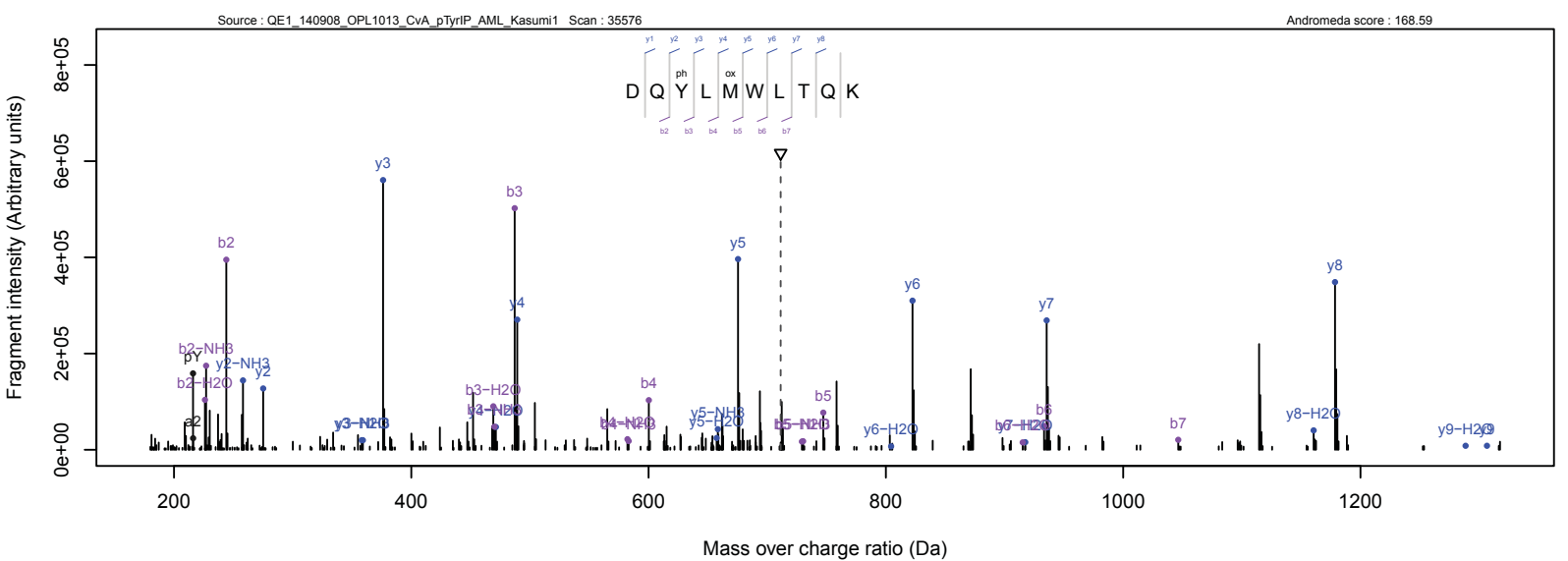
Andromeda score : 168.62



DQY(ph)LMWLTQK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 35576

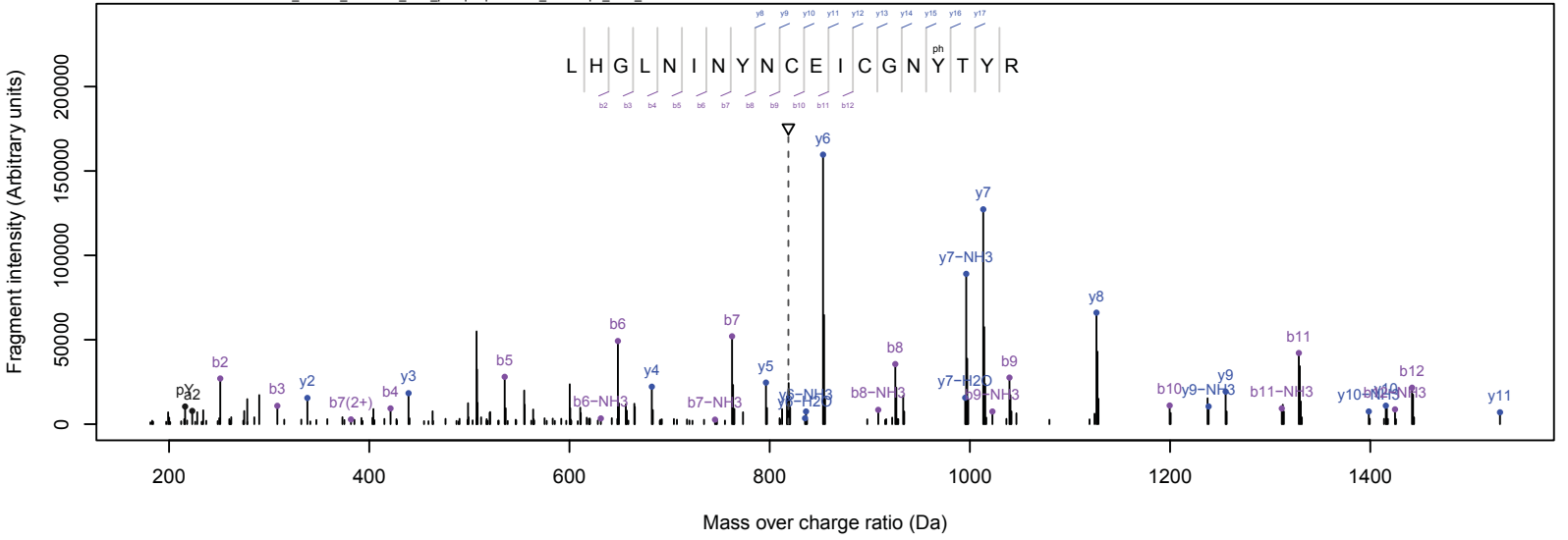
Andromeda score : 168.59



LHGLNINYNCEICGN<sup>ph</sup>TYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 29720

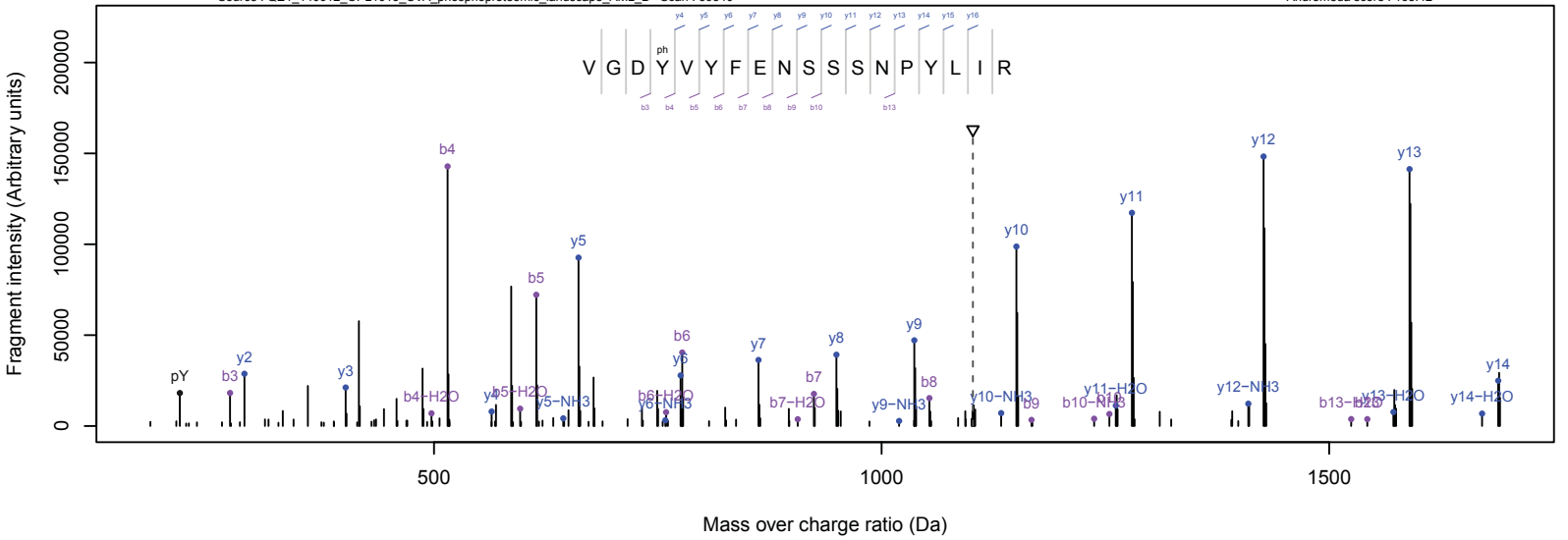
Andromeda score: 168.42



VG DY<sup>ph</sup> V Y F E N S S S N P Y L I R-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 39340

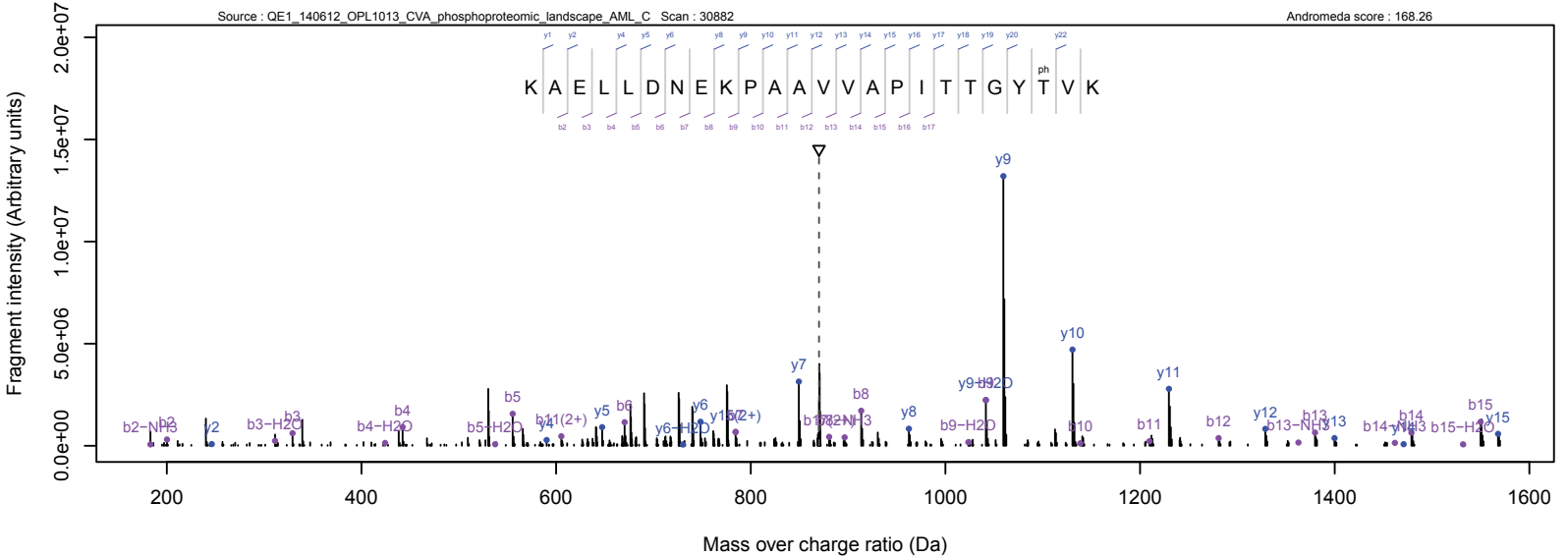
Andromeda score: 168.42



KA E L L D N E K P A A V V A P I T T G Y T V K-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 30882

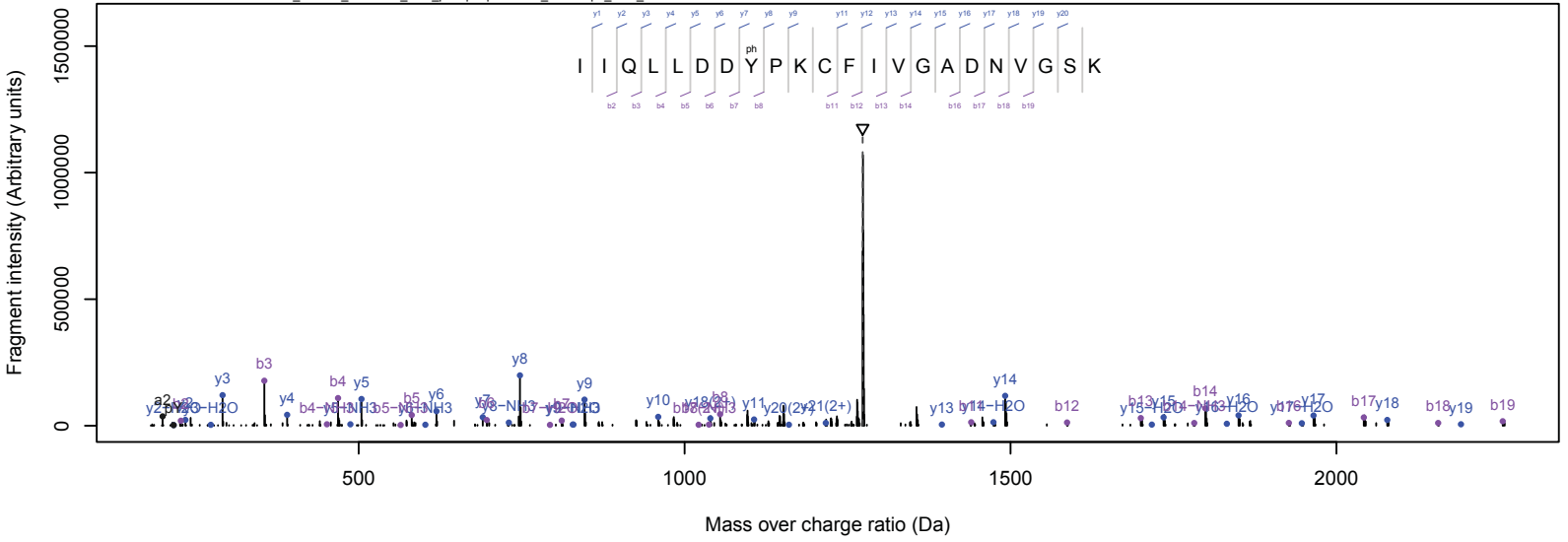
Andromeda score: 168.26



**IIQLDDY(ph)PKCFIVGADNVGSK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44529

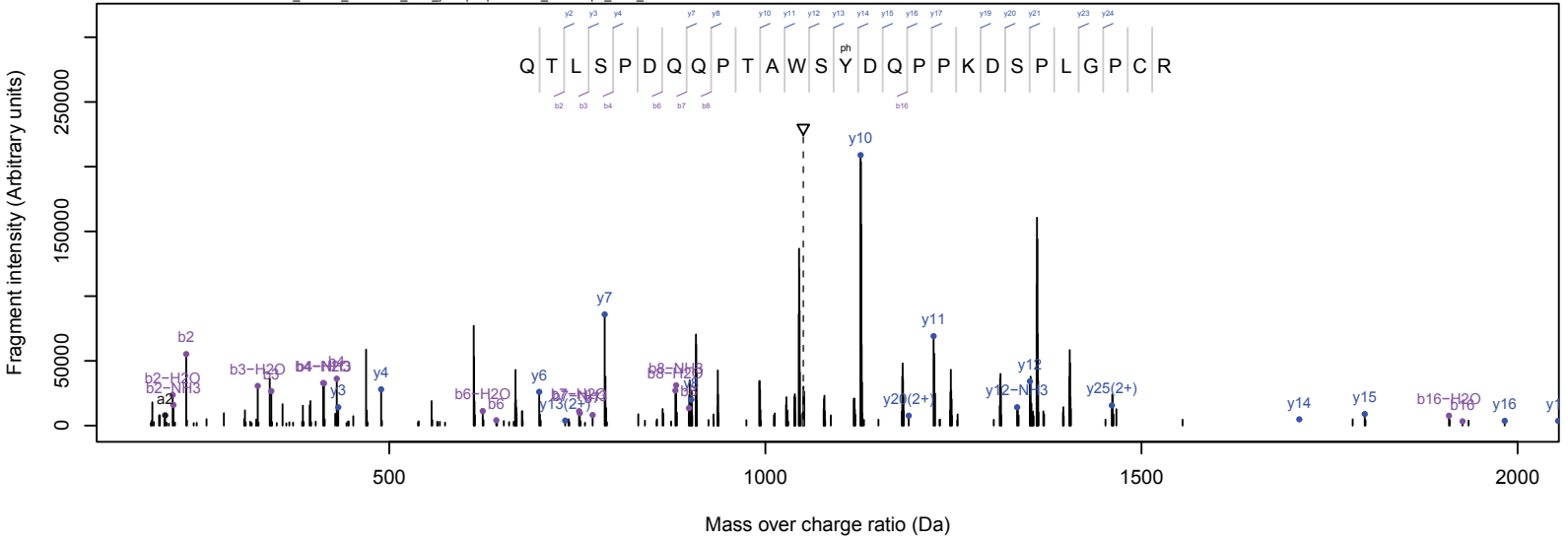
Andromeda score : 168.25



**QTLSPDQQPTAWSY(ph)DQPPKDSPLGPCR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 30037

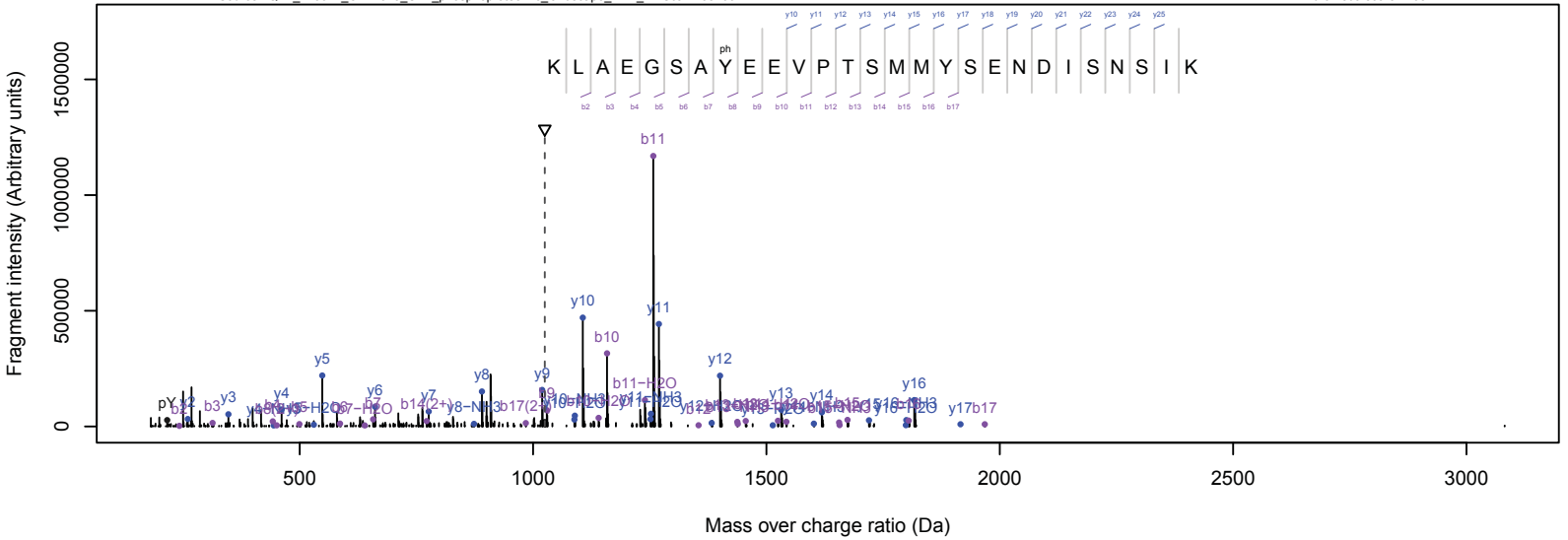
Andromeda score : 168.25



**KLAEGSAY(ph)EEVPTSMMYSENDISNSIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35400

Andromeda score : 168.2

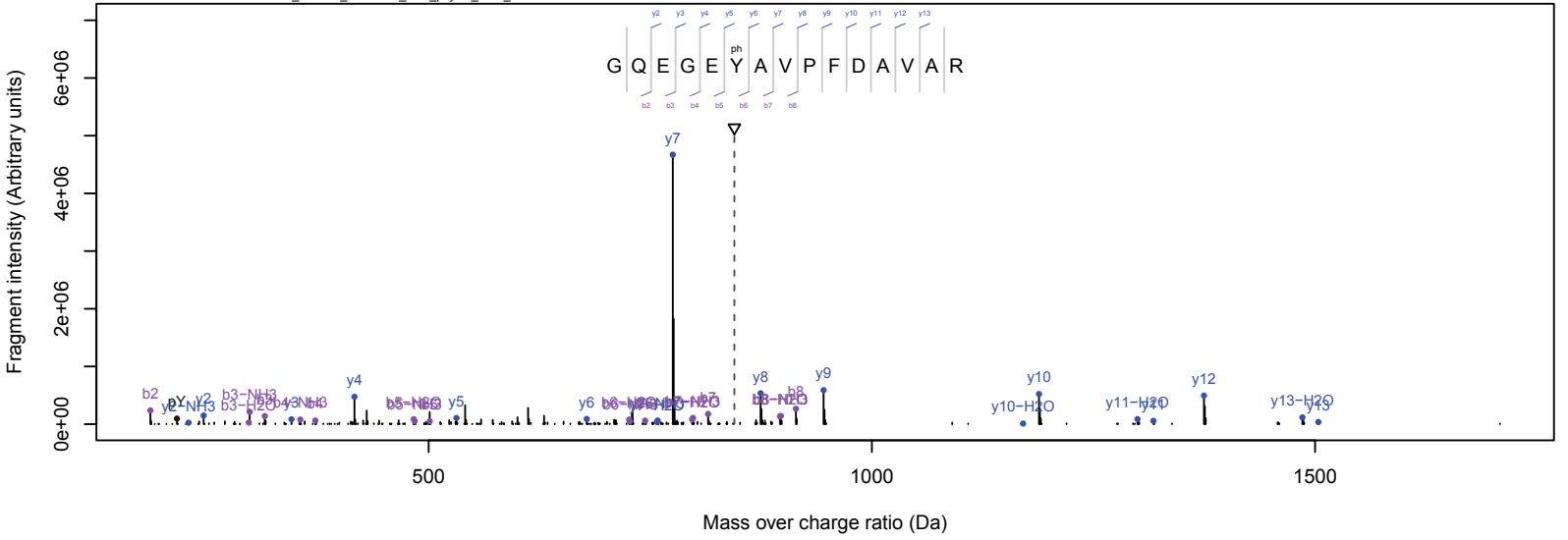




GQEGEY(ph)AVPFDVAR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31267

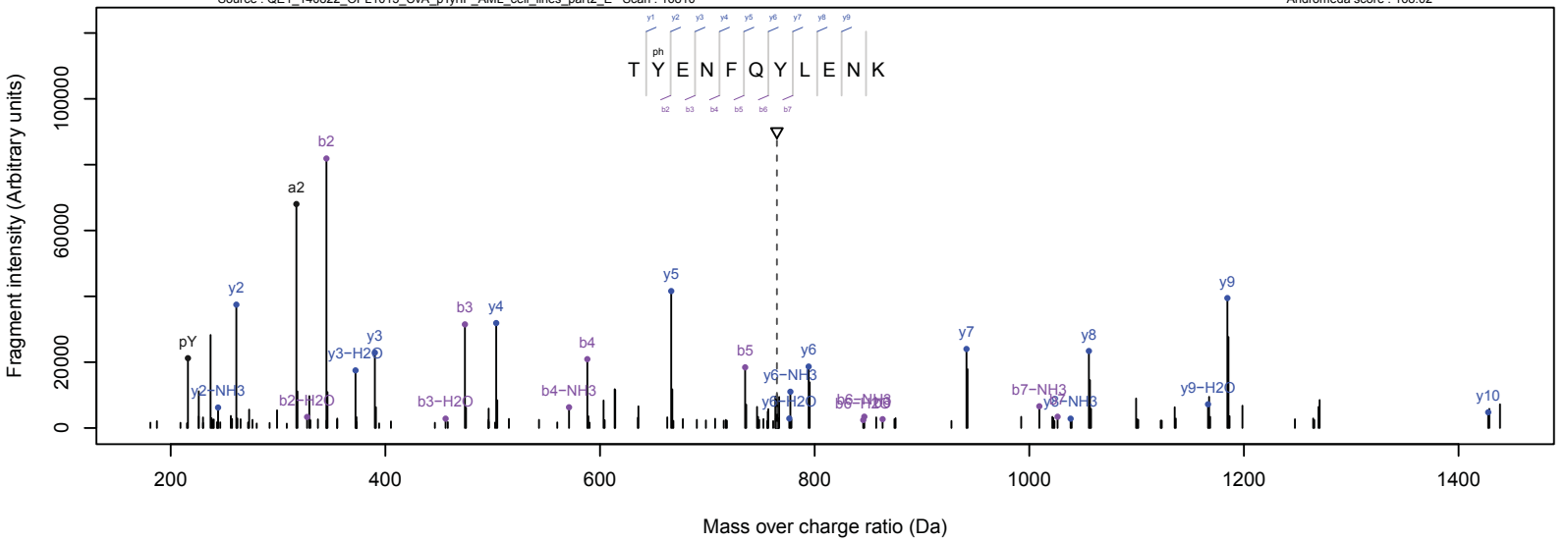
Andromeda score : 168.1



TY(ph)ENFQYLENK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 16810

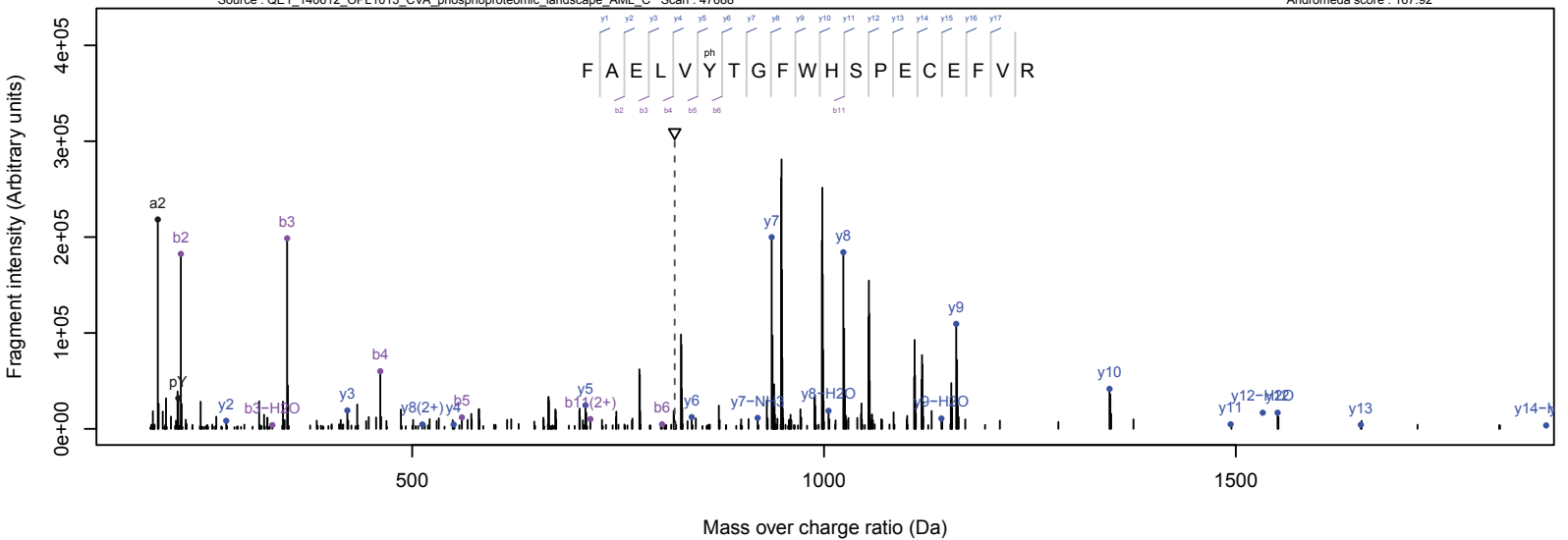
Andromeda score : 168.02



FAELVY(ph)TGFWHSPECFVR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 47688

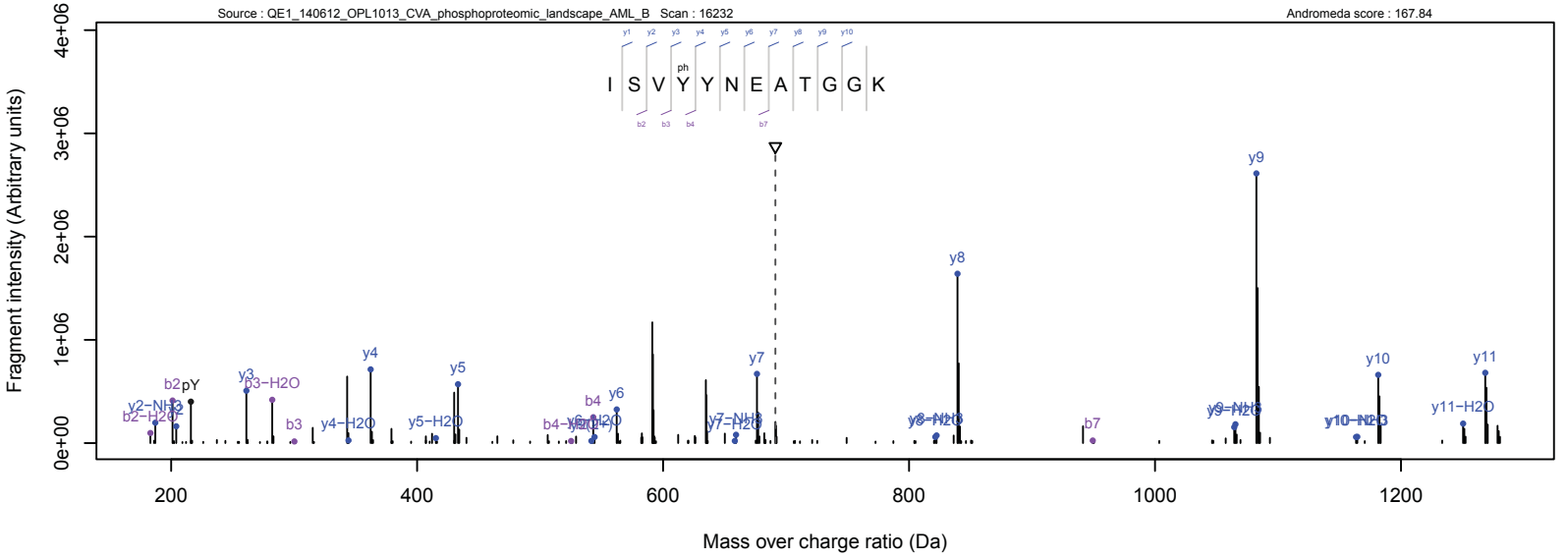
Andromeda score : 167.92



ISVY(ph)YNEATGGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16232

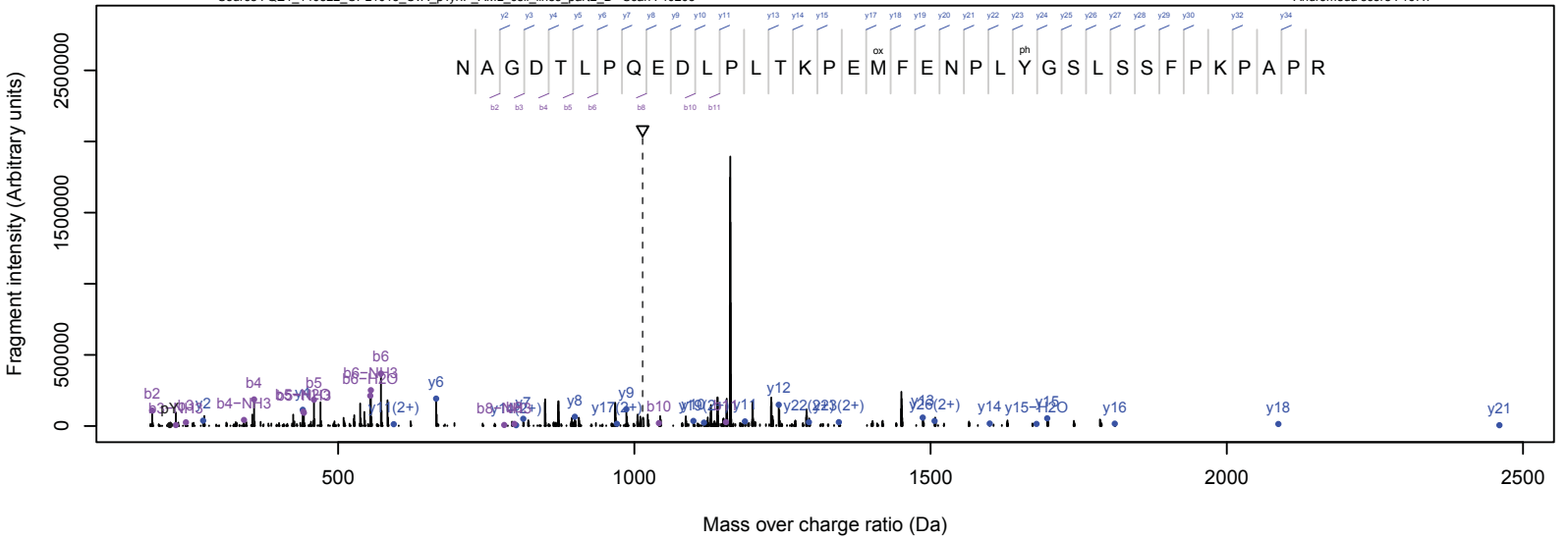
Andromeda score : 167.84



NAGDTLPQEDLPLTKPEM(ox)FENPLY(ph)GSLSSFPKPAPR-4+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 40205

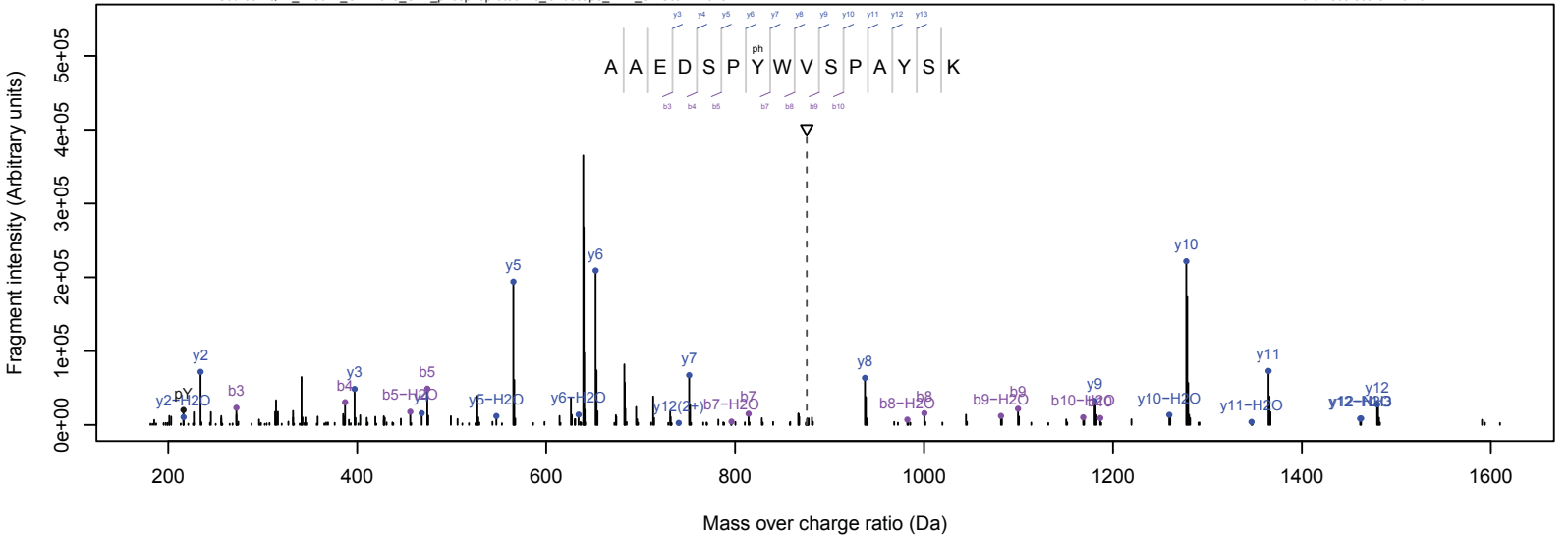
Andromeda score : 167.7



AAEDSPY(ph)WVSPAYSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25487

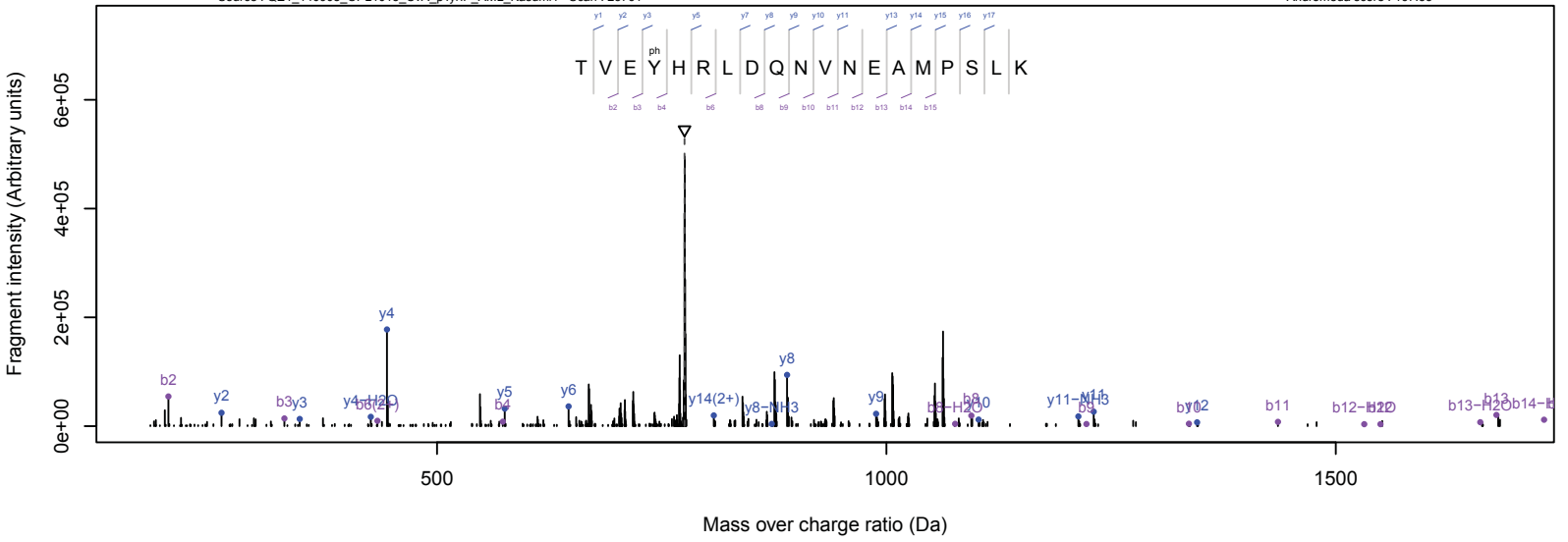
Andromeda score : 167.61



TVEY(ph)HRLDQNVNEAMP<sup>SLK</sup>-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 23761

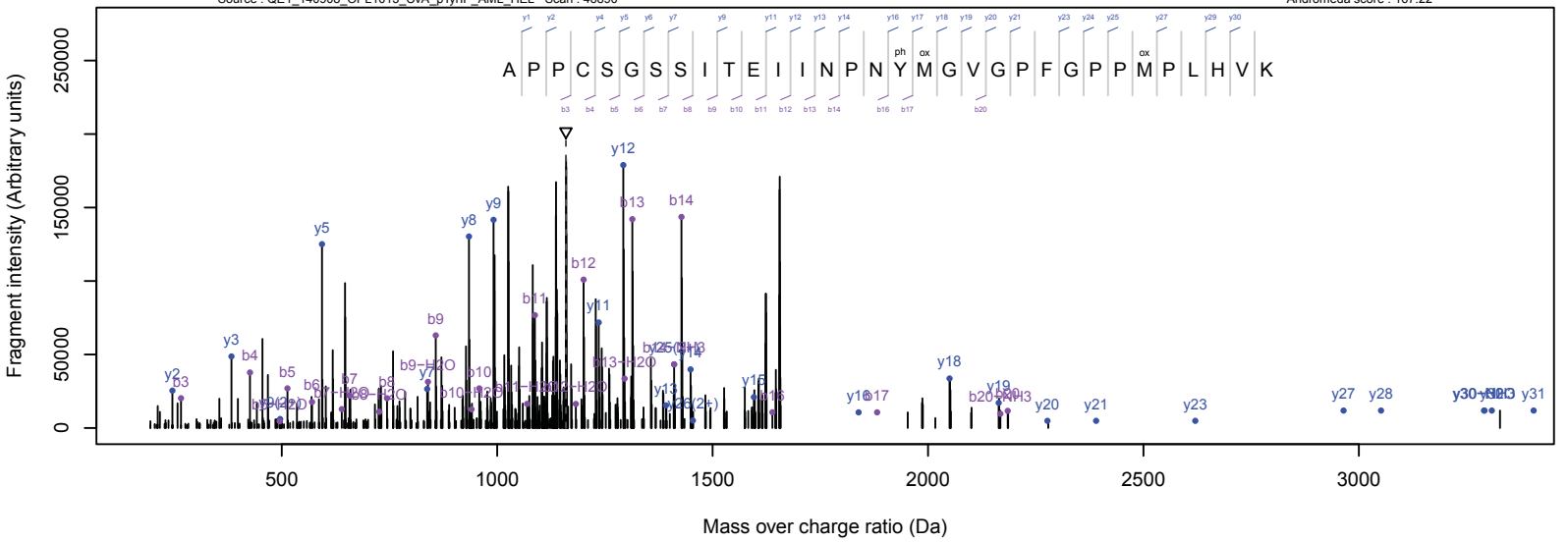
Andromeda score : 167.53



APPCSGSSITEIINP<sup>NYM</sup>(ox)GVGVPFGPP<sup>ox</sup>MPLHVK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 46890

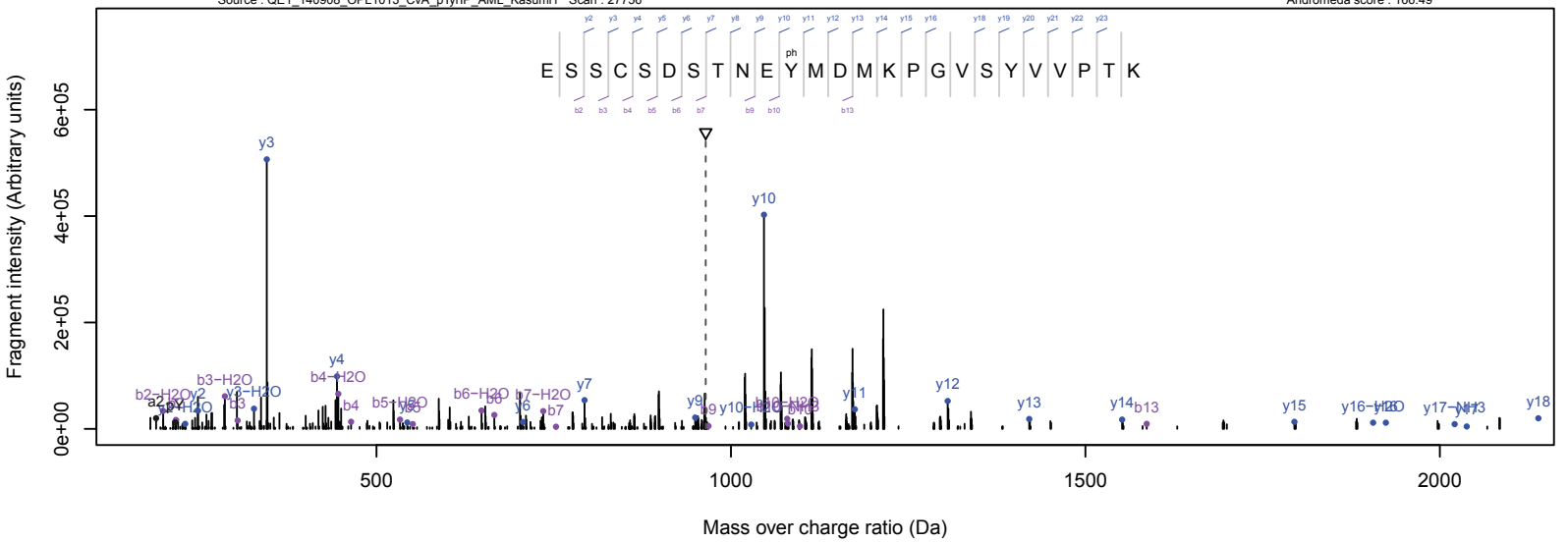
Andromeda score : 167.22



ESSCSDSTNEY(ph)MDMKPGVSYVPTK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27756

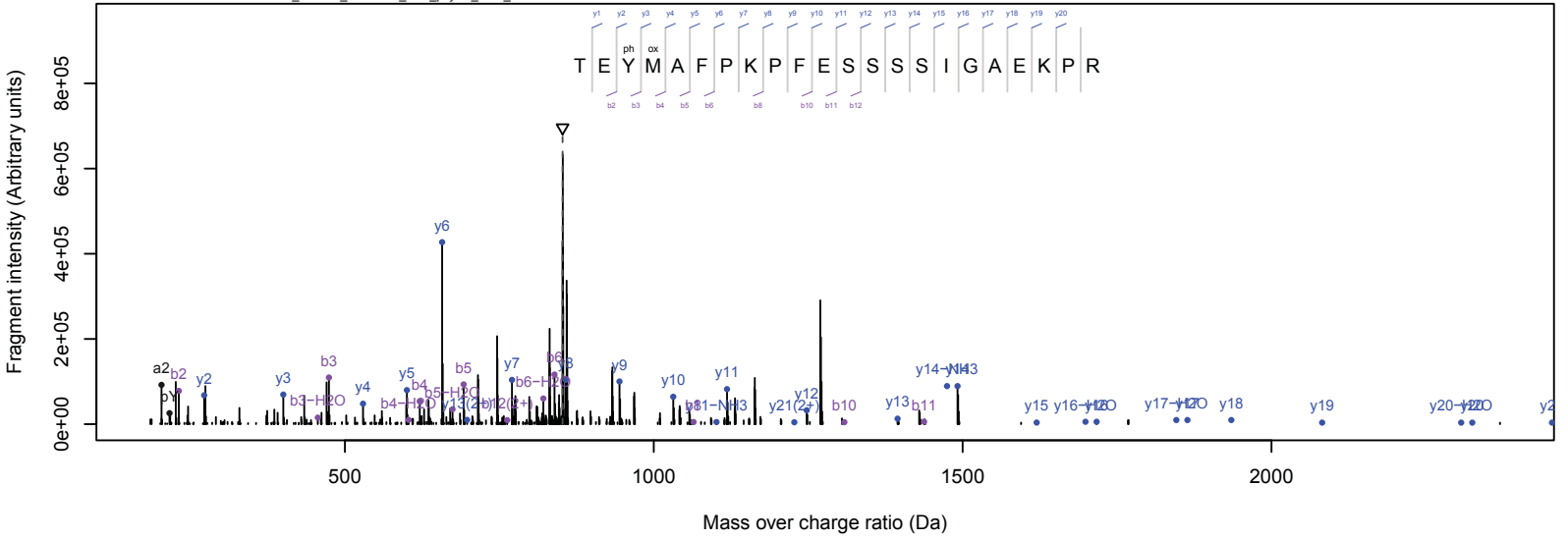
Andromeda score : 166.49



TEY(ph)M(ox)AFPKPFESSSSIGAEKPR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 26978

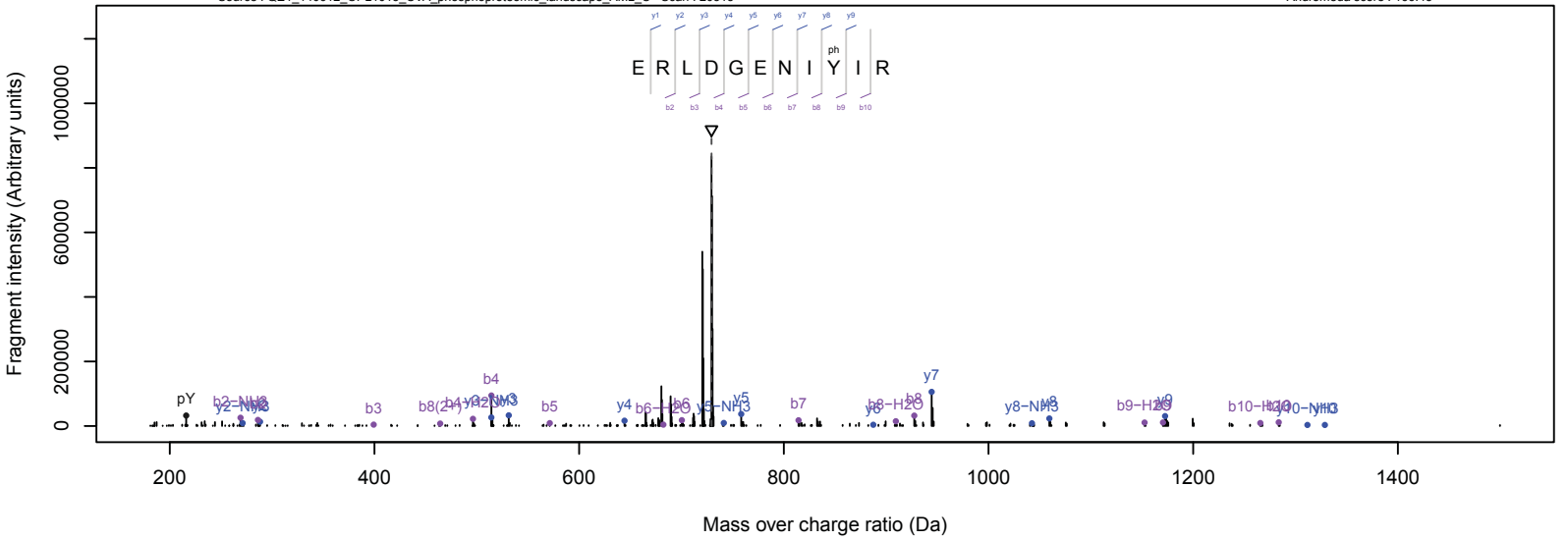
Andromeda score : 166.49



ERLDGENIY(ph)IR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20513

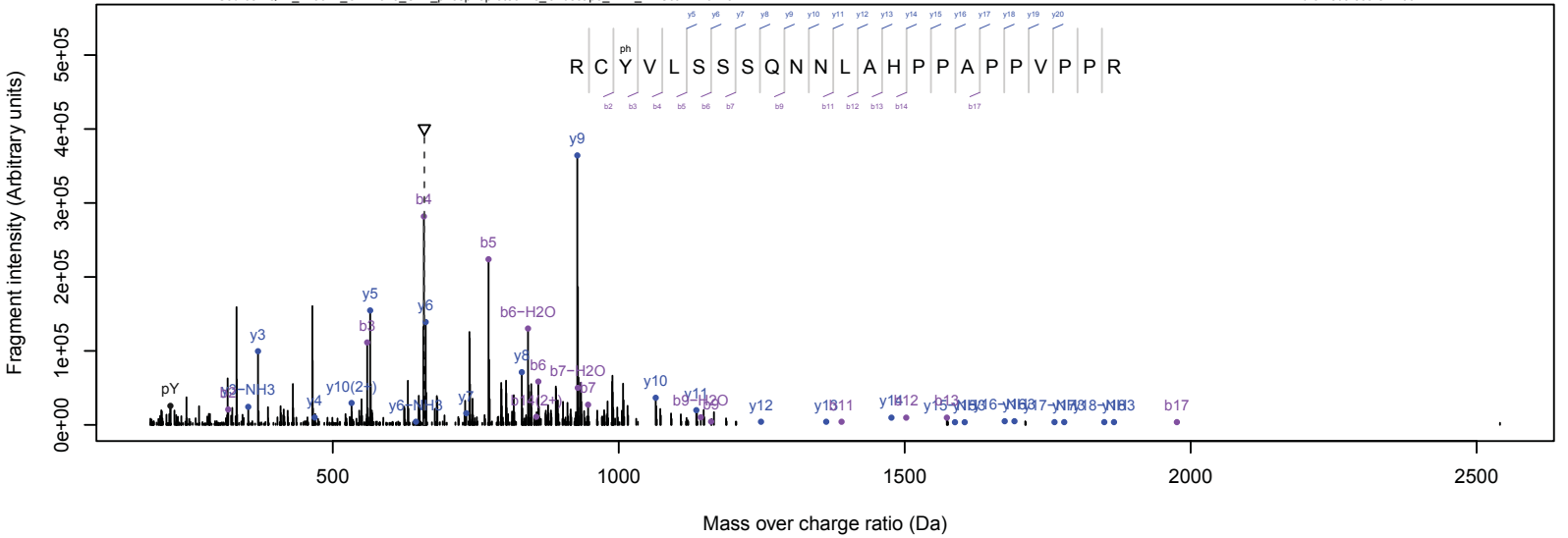
Andromeda score : 166.45



RCY(ph)VLSSSQNNLAHPPAPPVPPR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26175

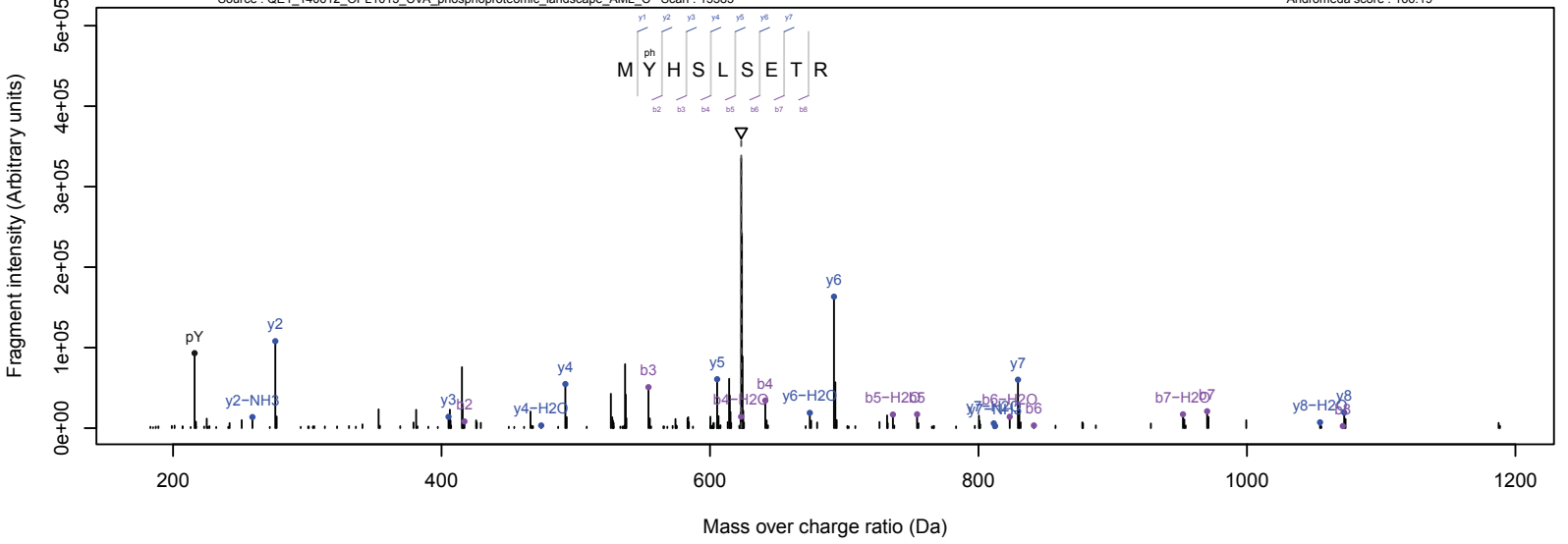
Andromeda score : 166.27



(ac)MY(ph)HSLSETR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15585

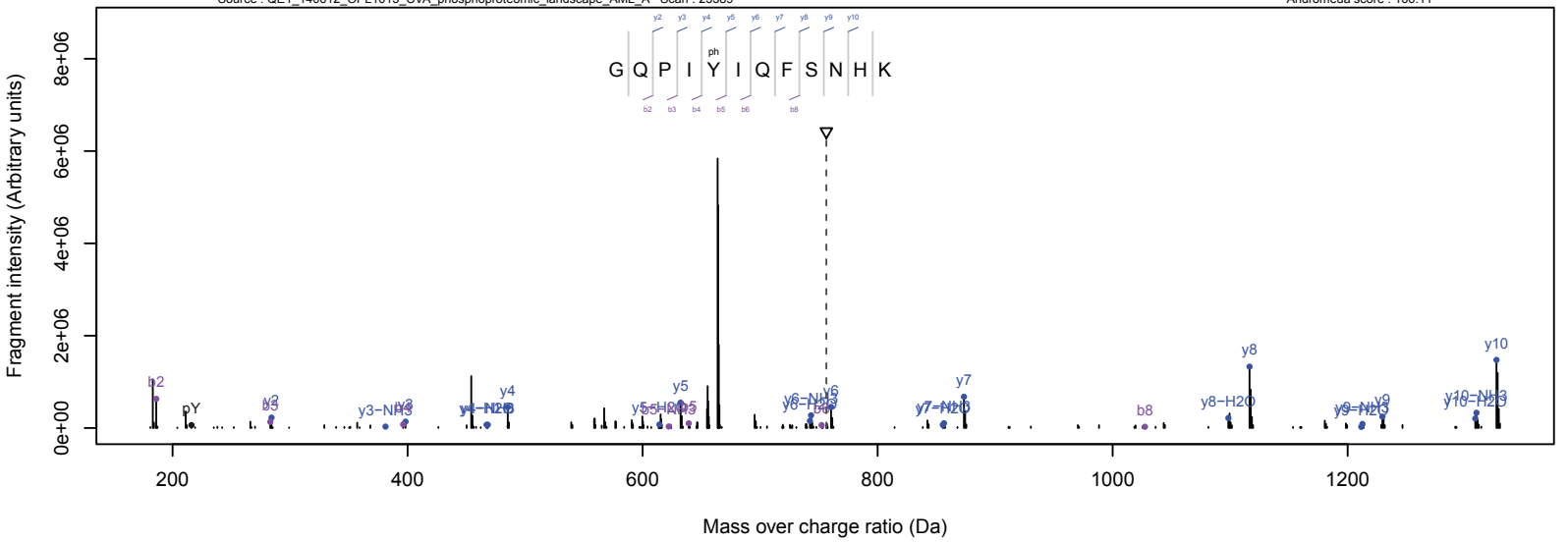
Andromeda score: 166.19



GQPIY(ph)IQFSNHK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 23389

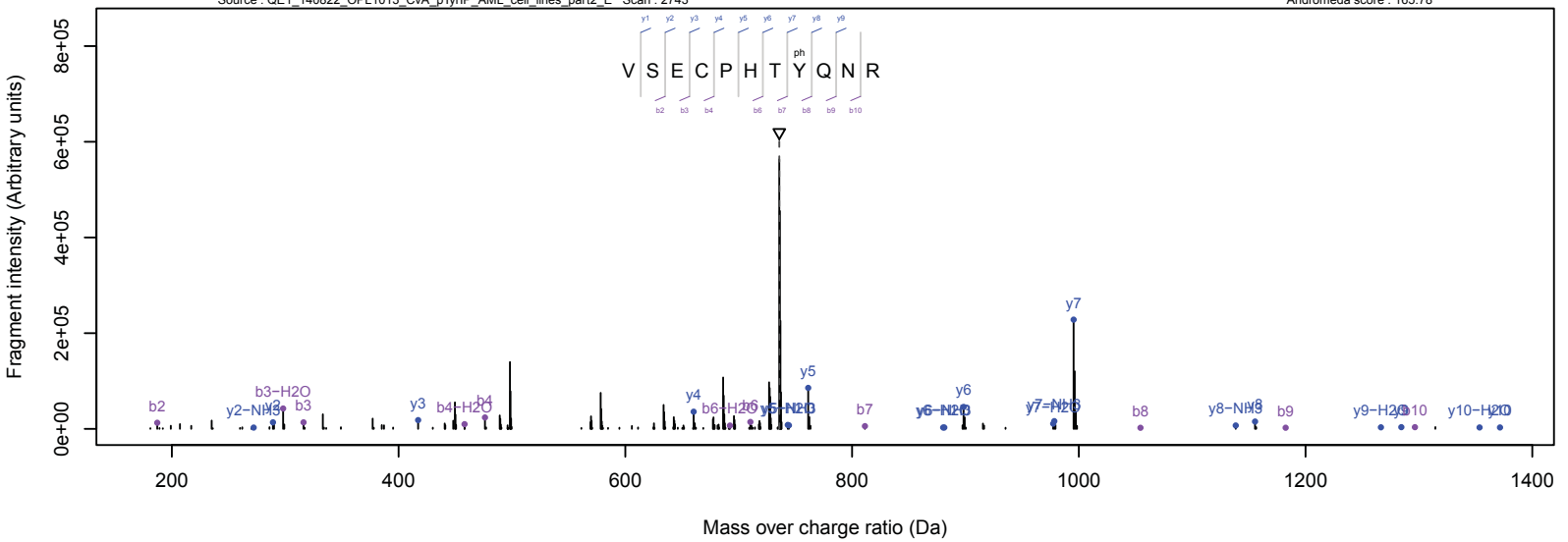
Andromeda score: 166.11



VSECPHTY(ph)QNR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 2743

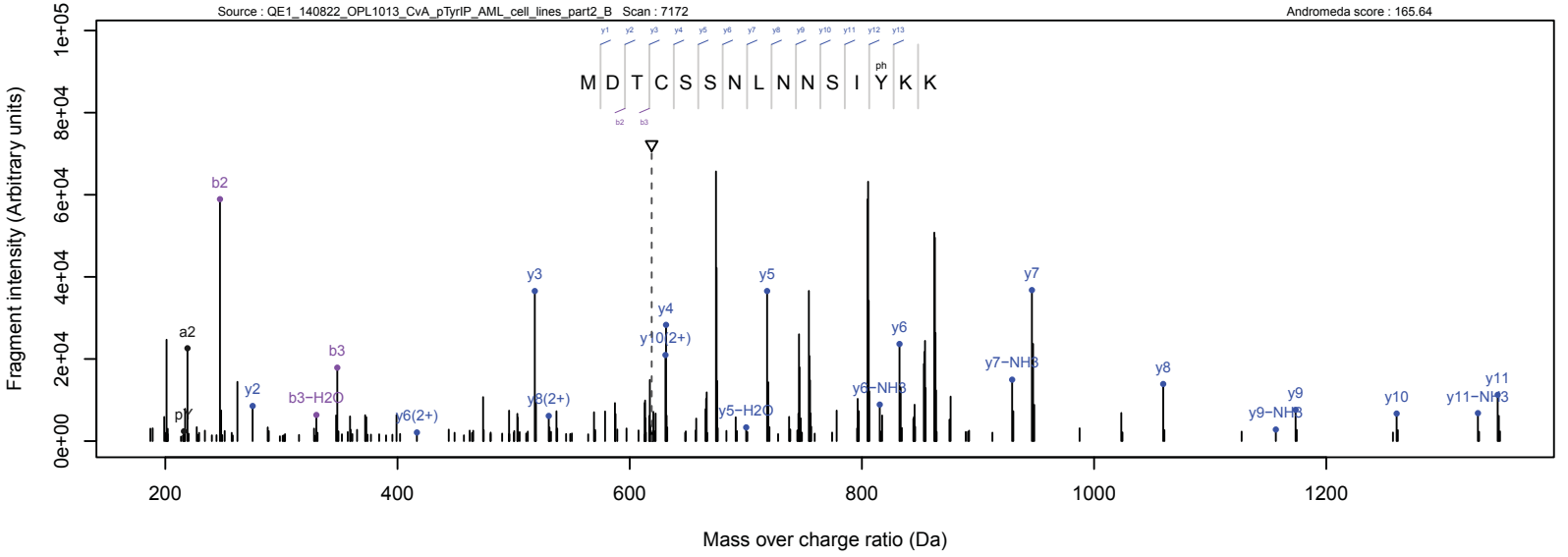
Andromeda score: 165.78



**MDTCSSNLNSIY(ph)KK-3+**

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 7172

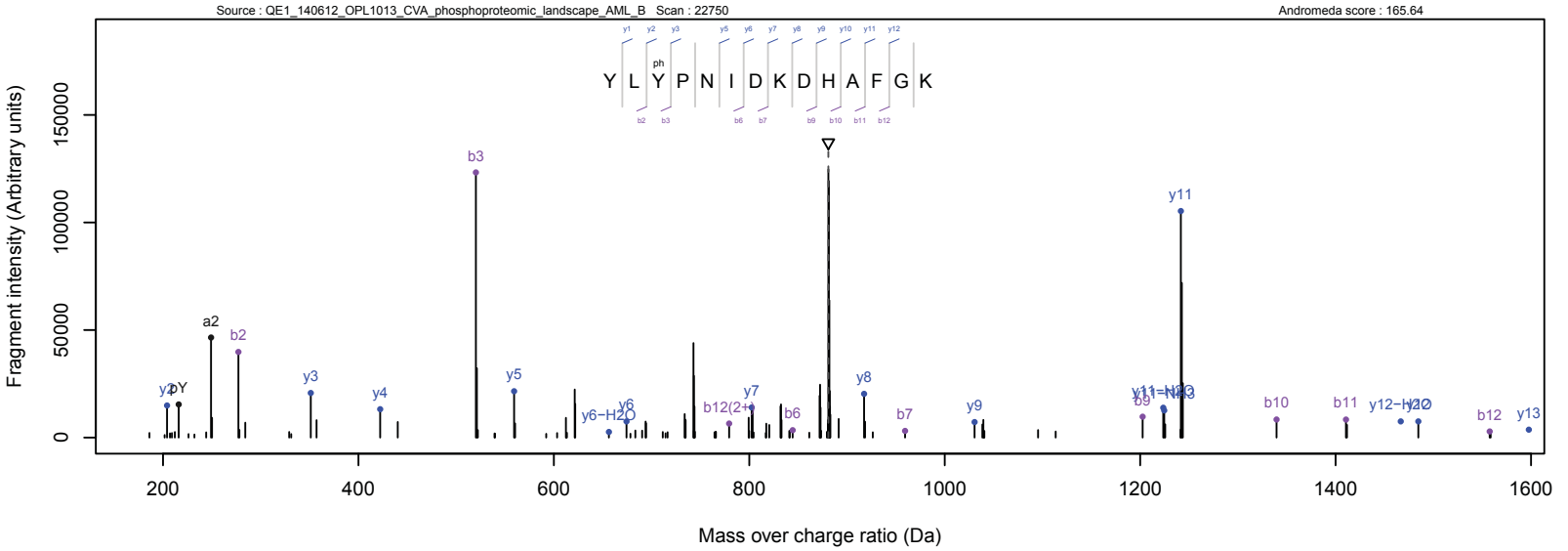
Andromeda score : 165.64



**YLY(ph)PNIDKDHAFGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22750

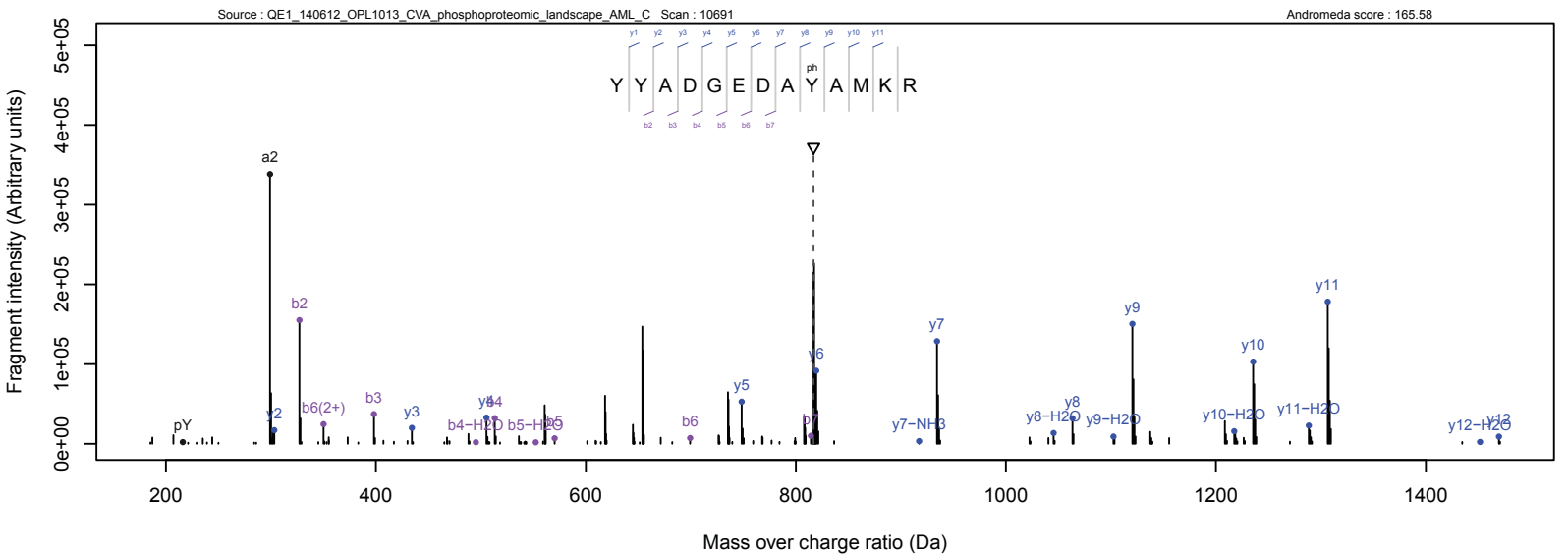
Andromeda score : 165.64



**YYADGEDAY(ph)AMKR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10691

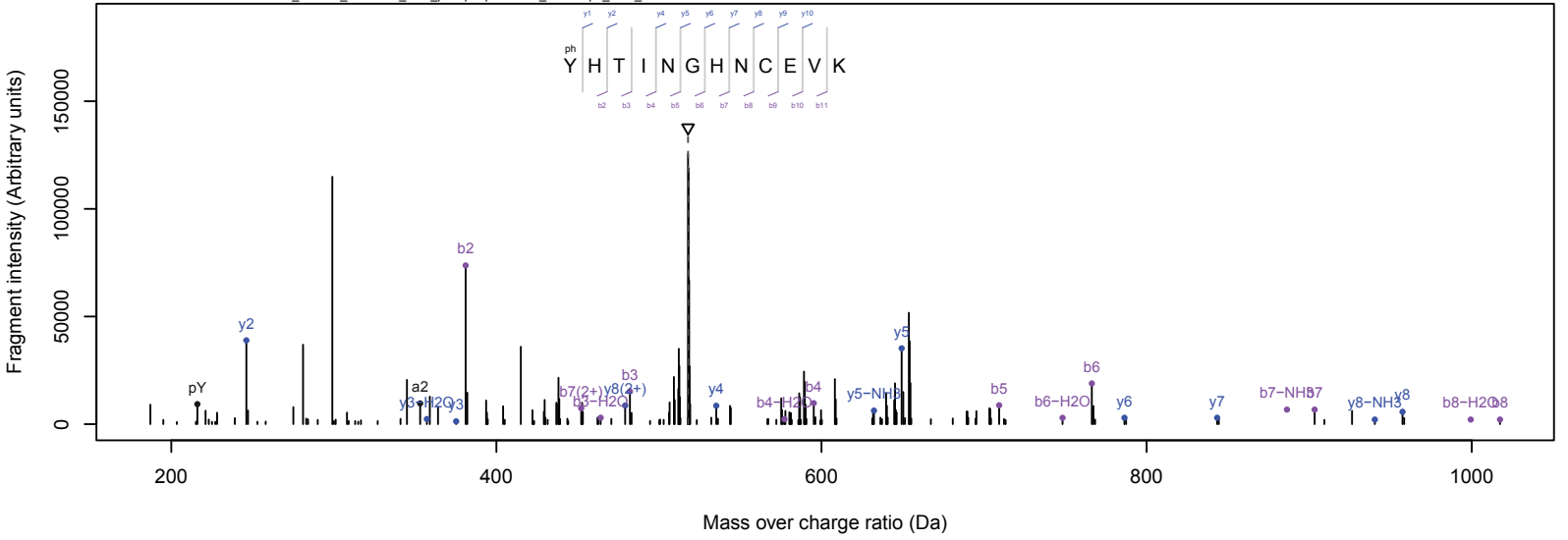
Andromeda score : 165.58



Y(ph)HTINGHNCEVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 3897

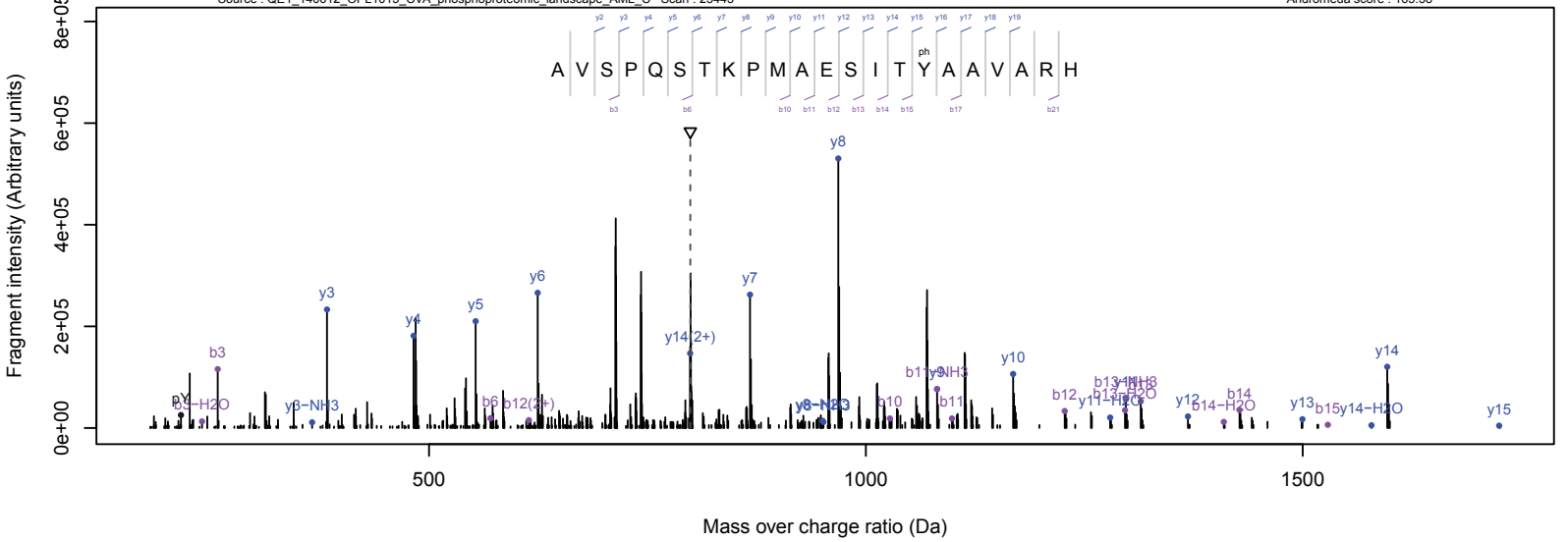
Andromeda score: 165.51



AVSPQSTKPMAESITY(ph)AAVARH-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25443

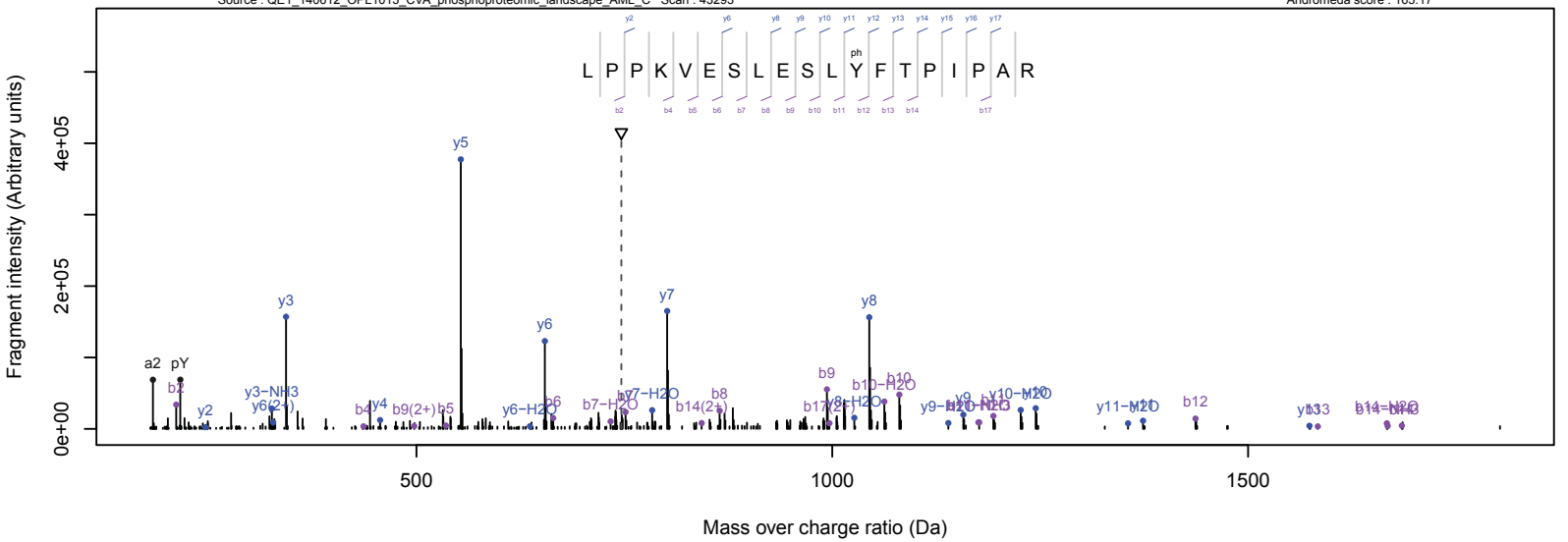
Andromeda score: 165.38



LPPKVESLESLY(ph)FTPIPAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 43293

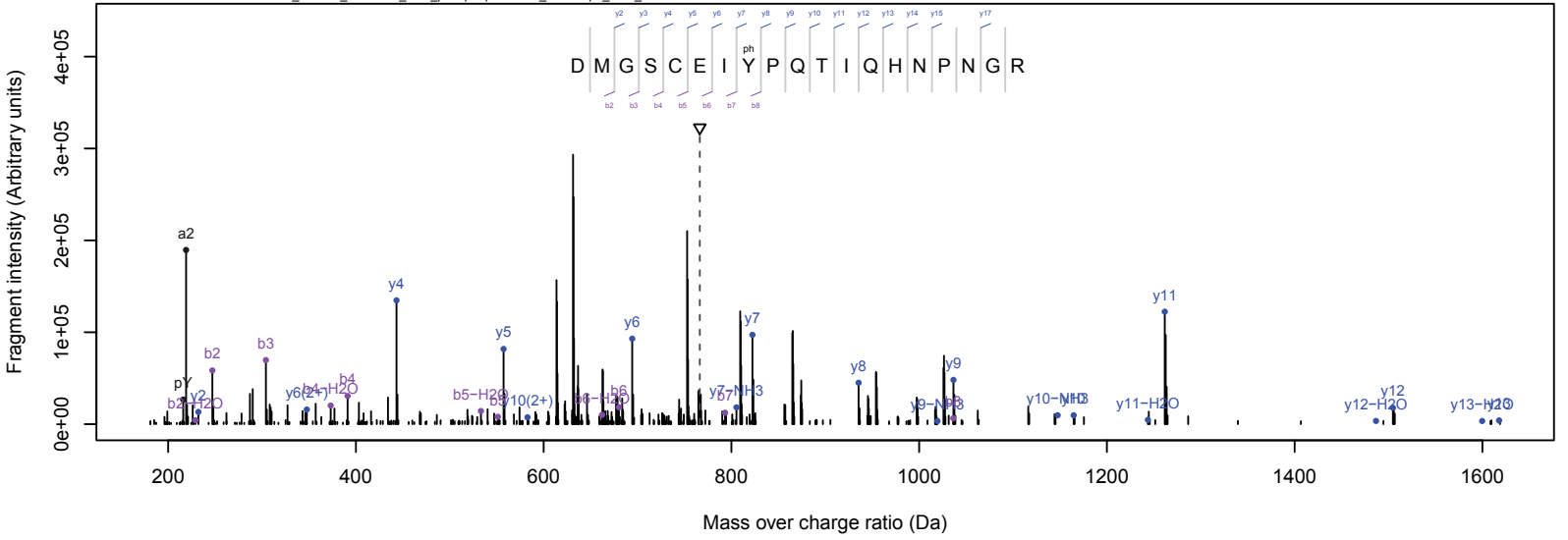
Andromeda score: 165.17



**DMGSCEIY(ph)PQTIQHNPNGR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18488

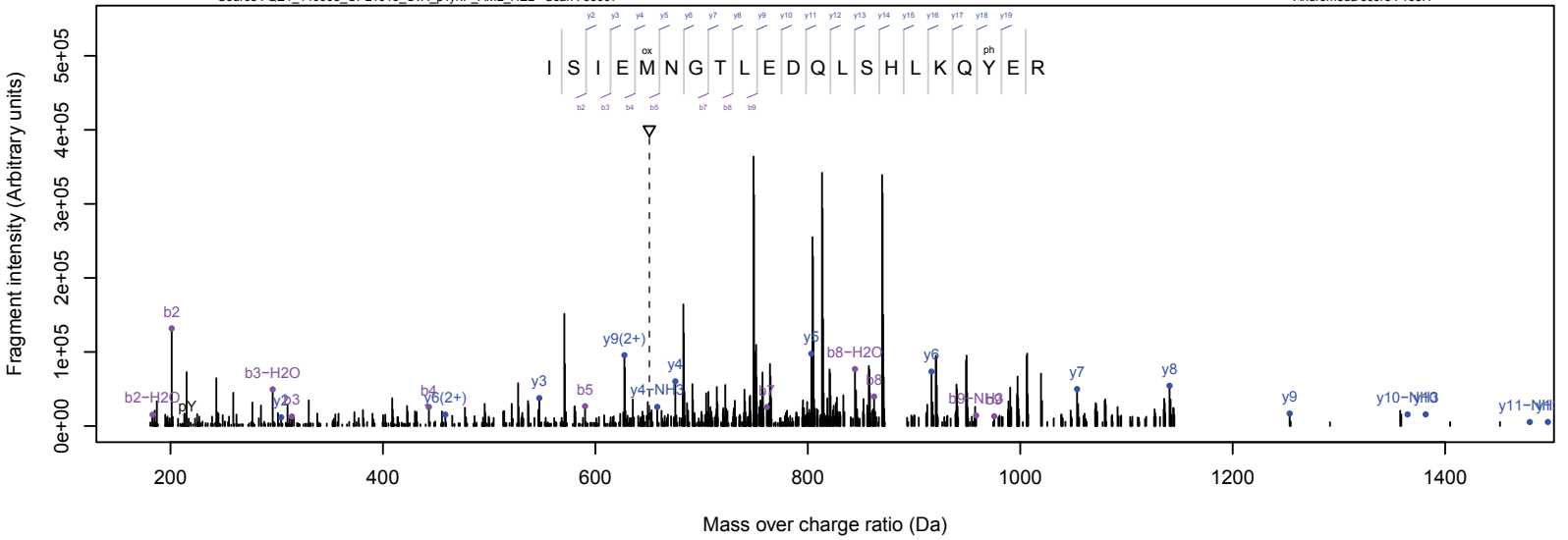
Andromeda score: 165.16



**ISIEM(ox)NGTLEDQLSHLKQY(ph)ER-4+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 35997

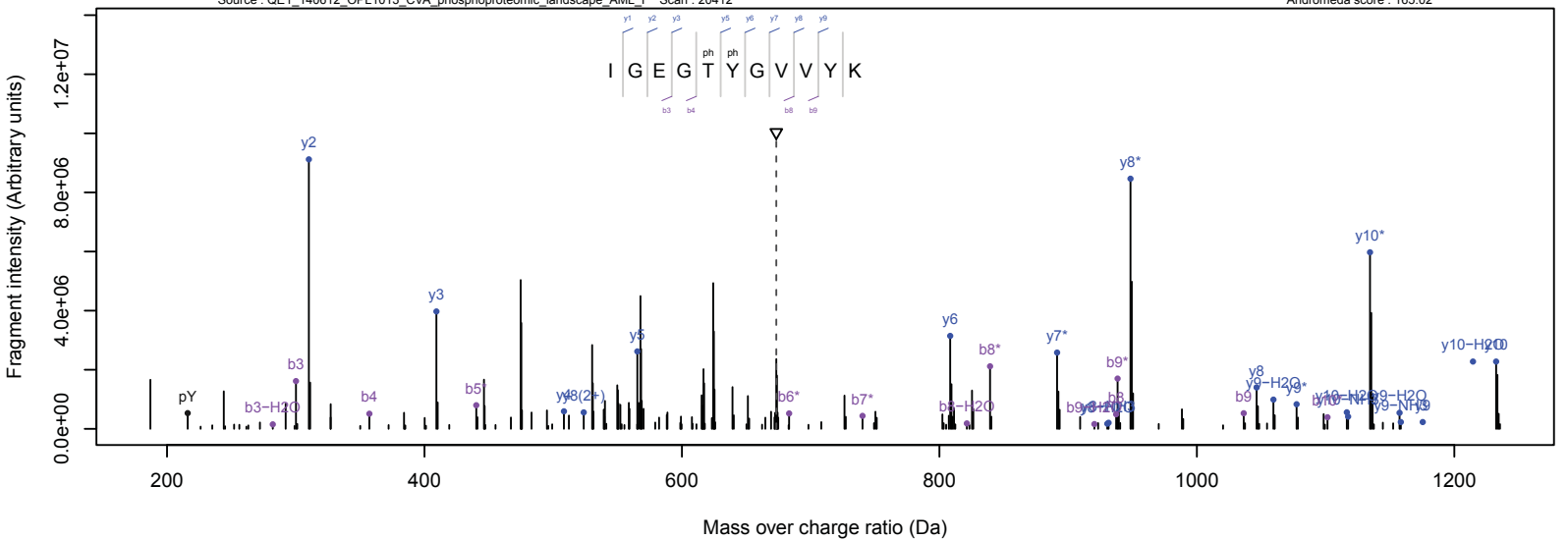
Andromeda score: 165.1



**IGEGT(ph)Y(ph)GVVYK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan: 20412

Andromeda score: 165.02

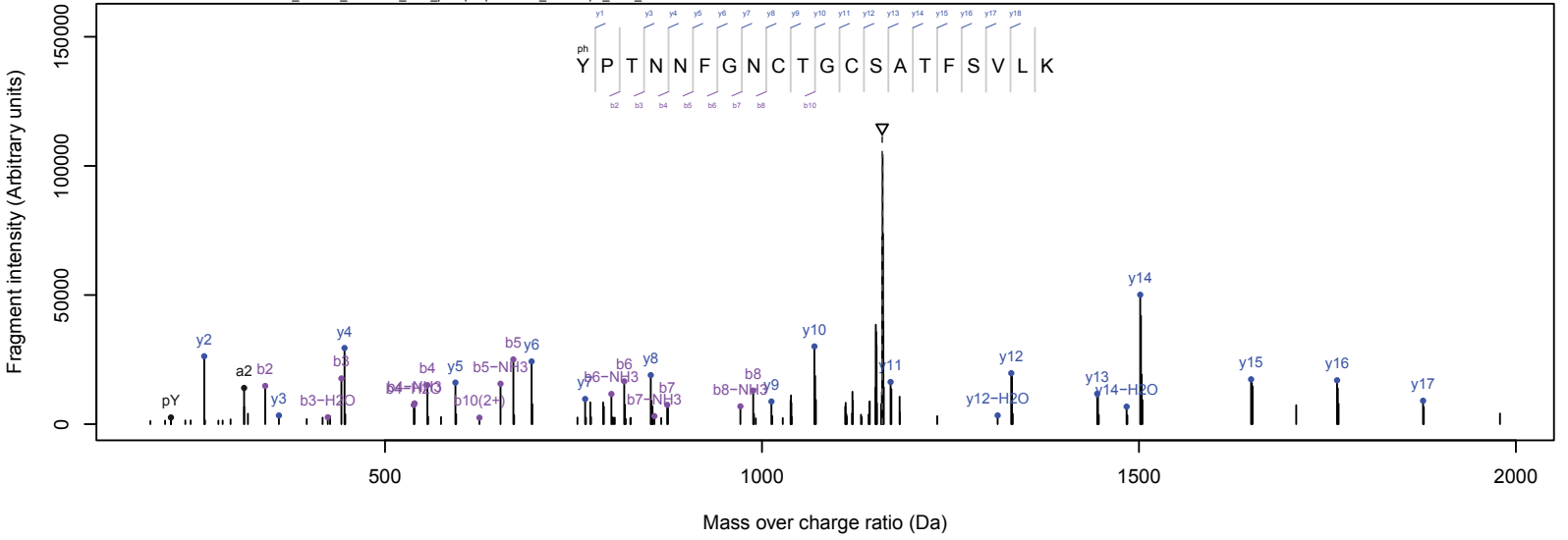




Y(ph)PTNFGNCTGCSATFSVLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34076

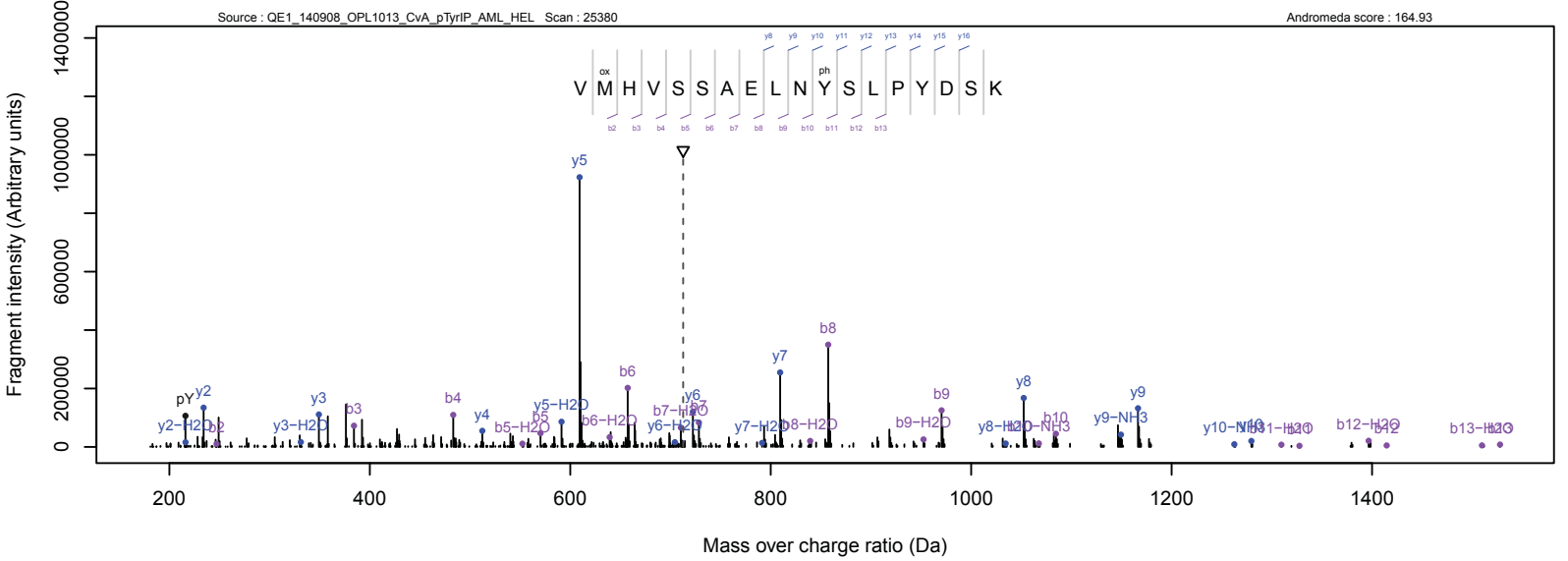
Andromeda score : 164.94



VM(ox)HVSSAELNY(ph)SLPYDSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 25380

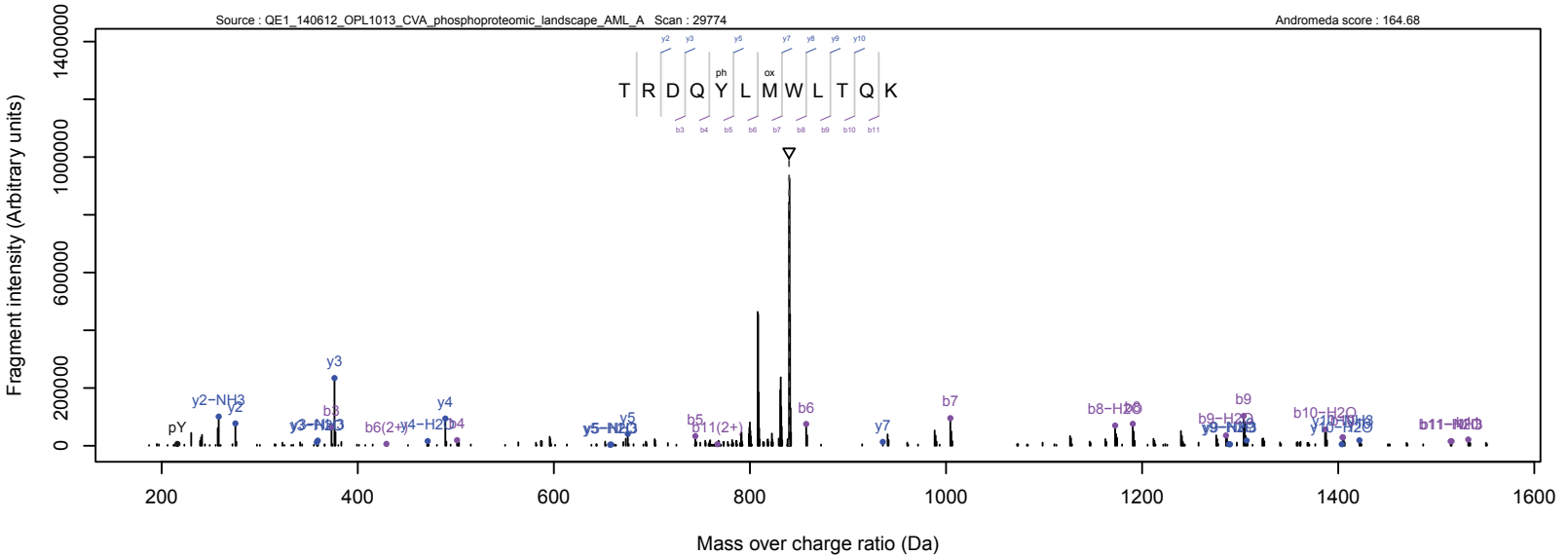
Andromeda score : 164.93



TRDQY(ph)LM(ox)WLTQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29774

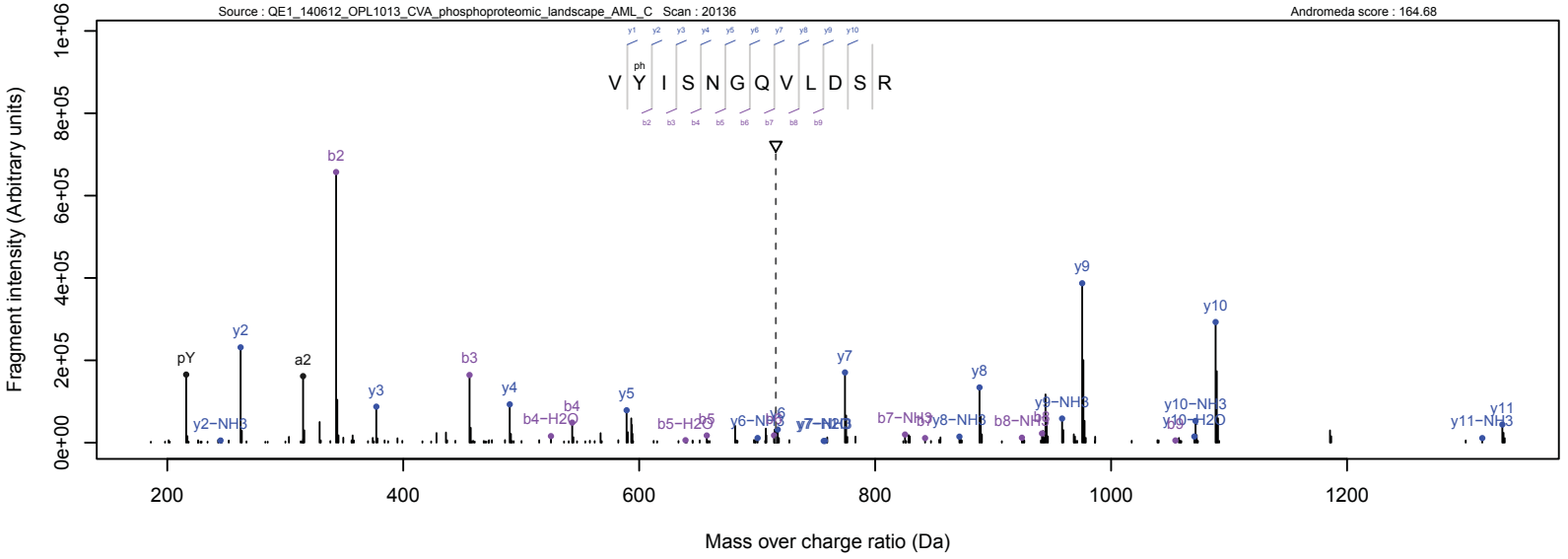
Andromeda score : 164.68



VY(ph)ISNGQLDSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20136

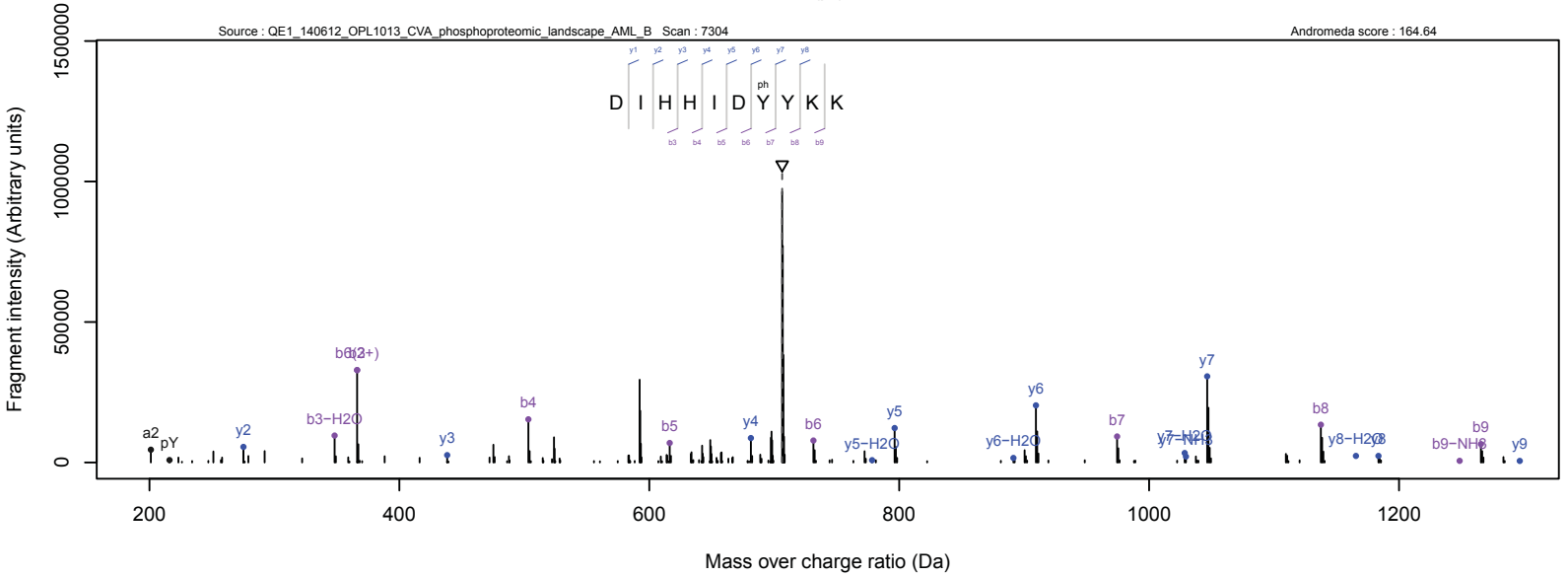
Andromeda score : 164.68



DIHHIDY(ph)YKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7304

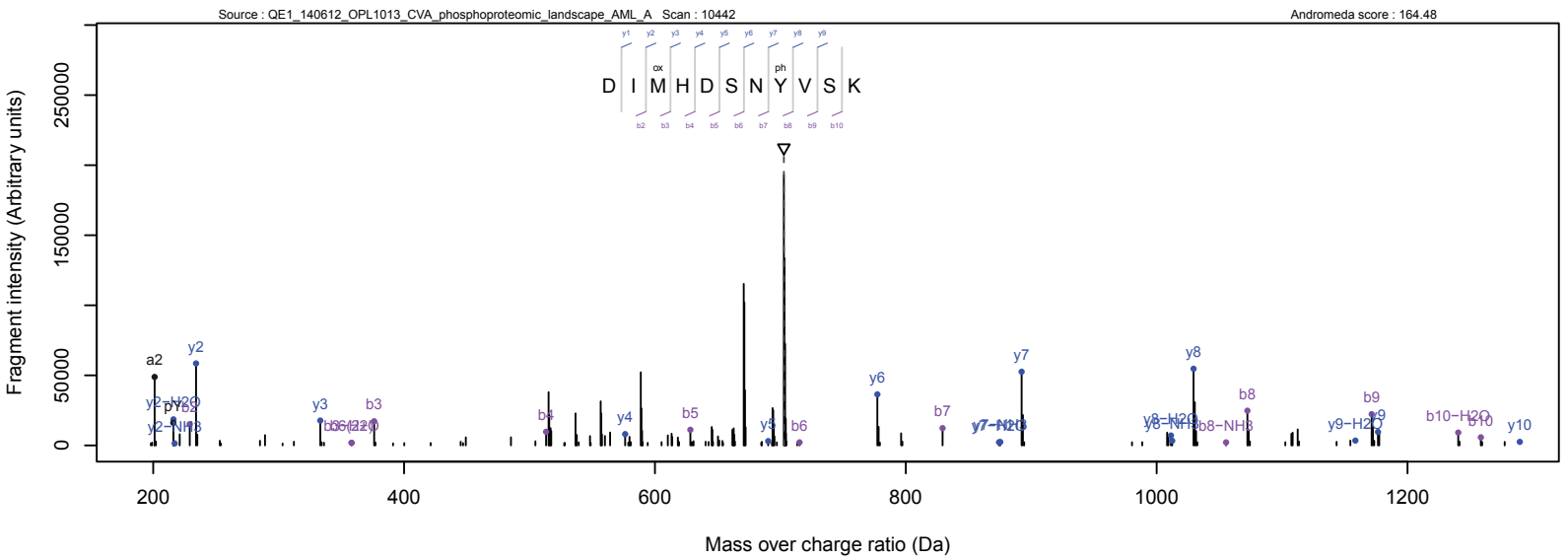
Andromeda score : 164.64



DIM(ox)HDSNY(ph)VSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 10442

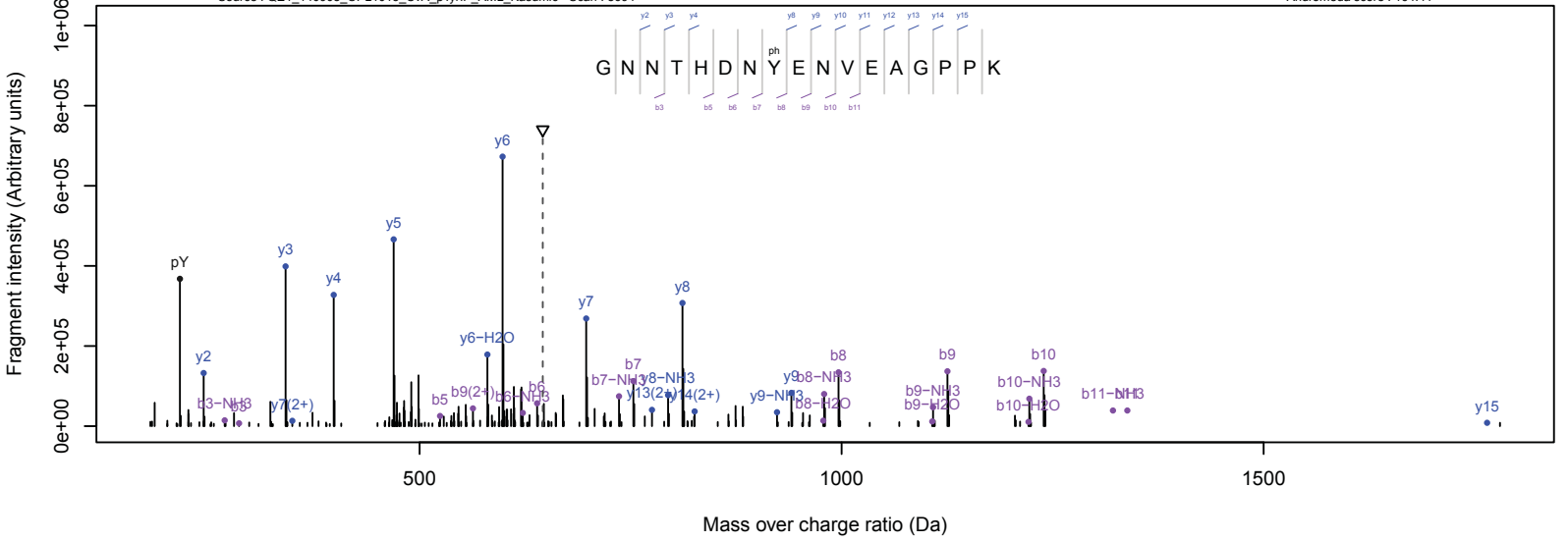
Andromeda score : 164.48



**GNNTHDNY(ph)ENVEAGPPK-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 8694

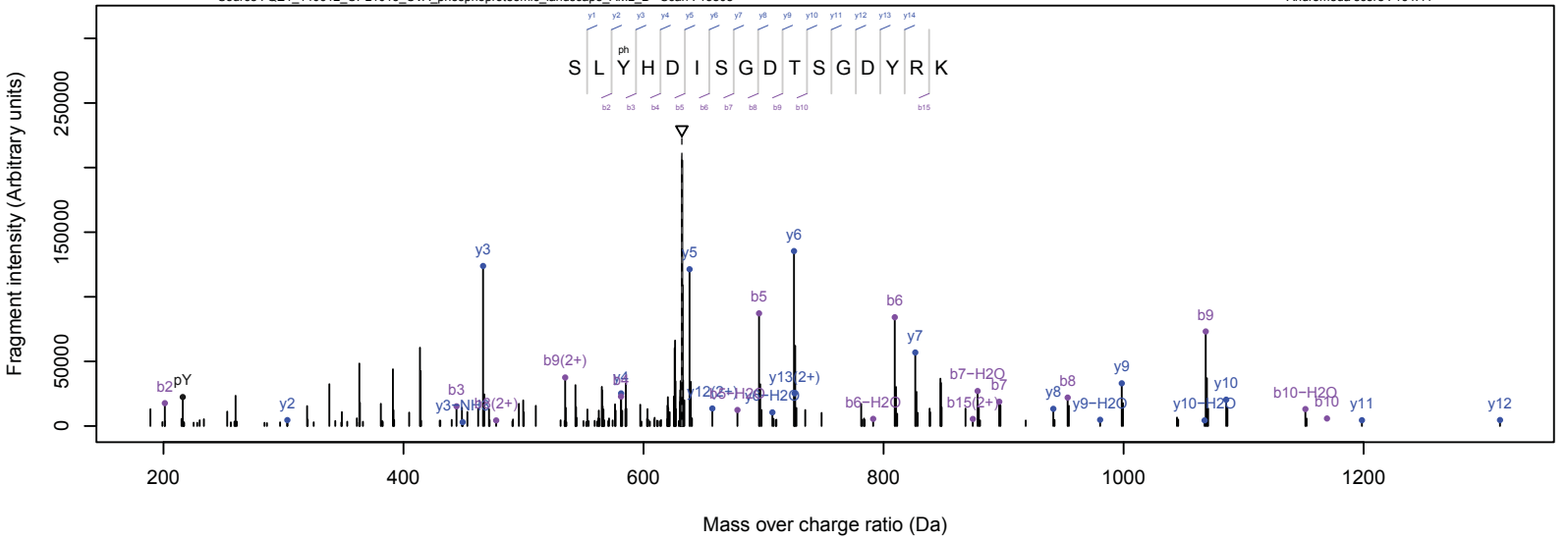
Andromeda score : 164.41



**SLY(ph)HDISGDTSGDYRK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13808

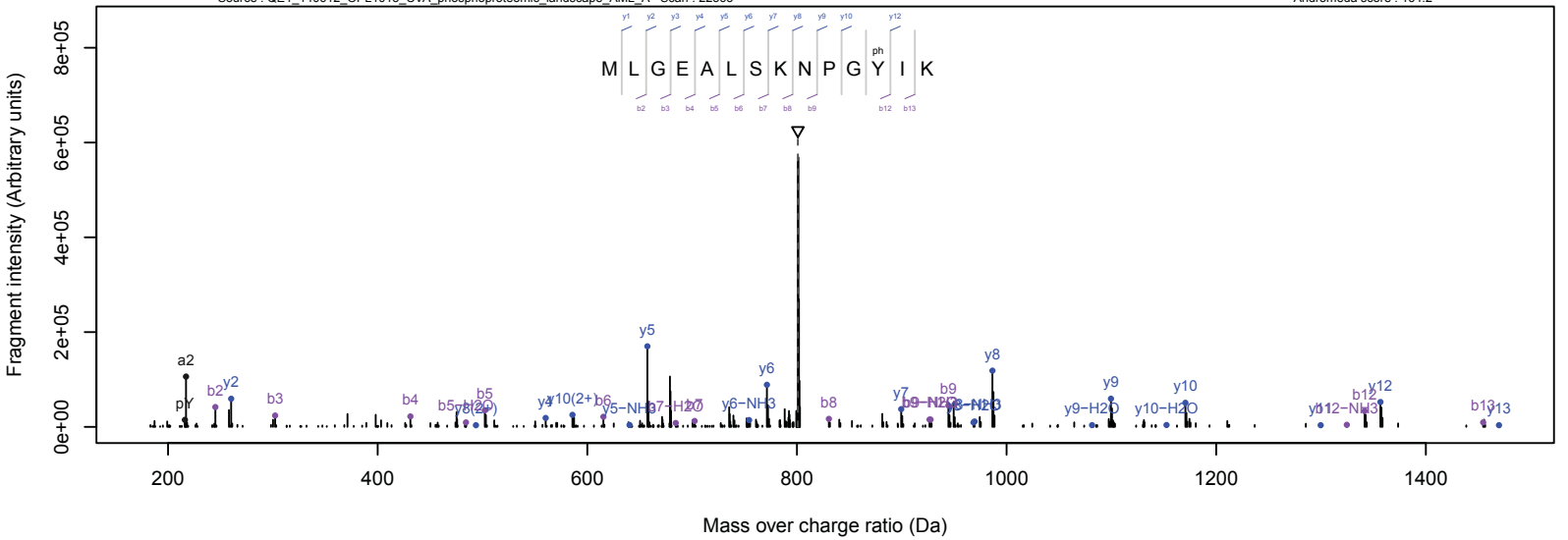
Andromeda score : 164.41



**MLGEALSKNP GYIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 22365

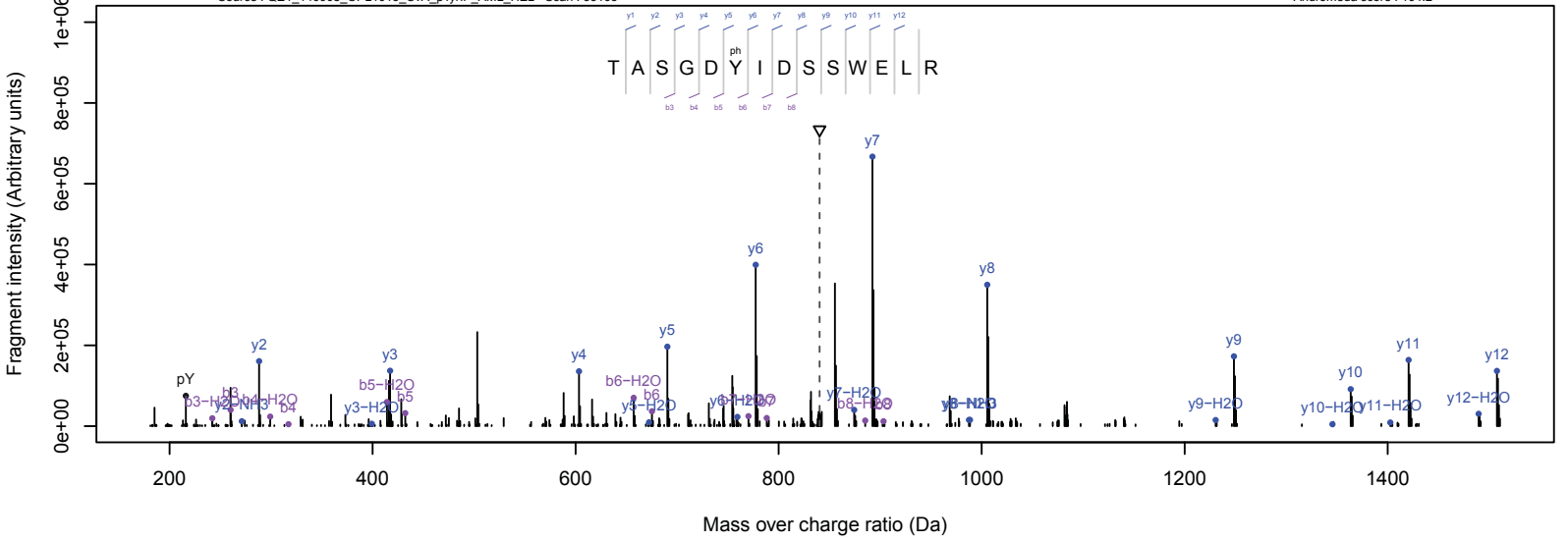
Andromeda score : 164.2



TASGDY(ph)IDSSWELR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 33163

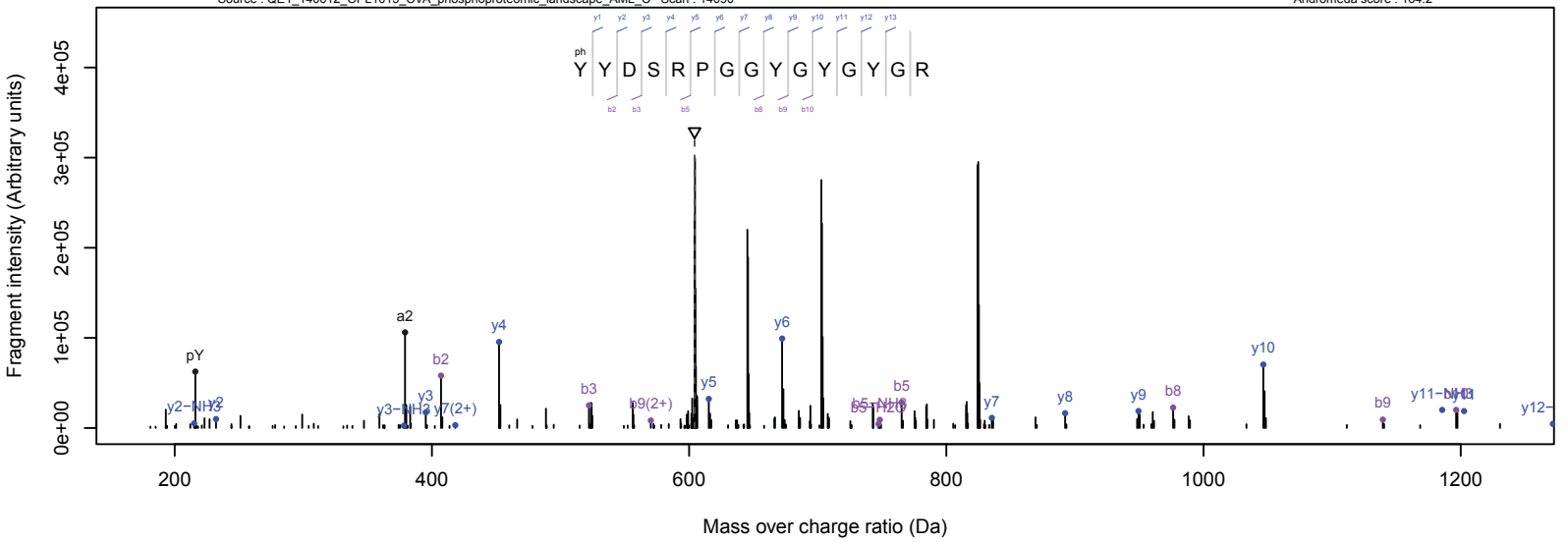
Andromeda score : 164.2



Y(ph)YDSRPGGYGYGR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 14090

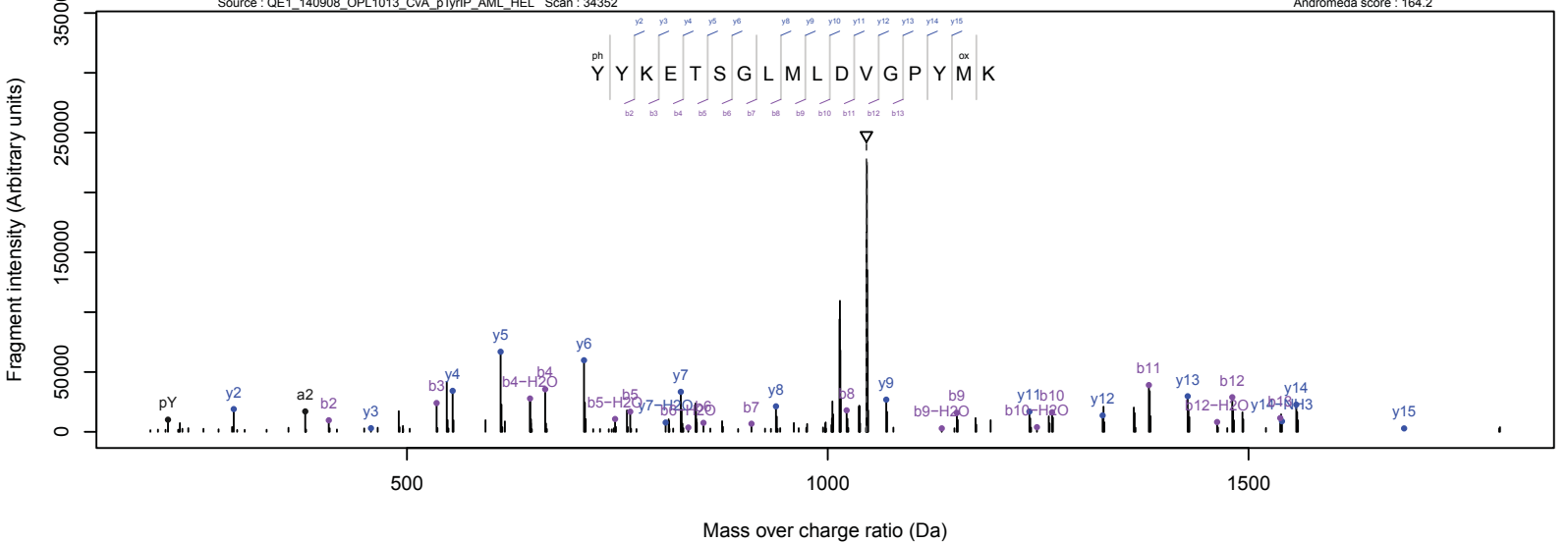
Andromeda score : 164.2



Y(ph)YKETSGLMLDVGPYMK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 34352

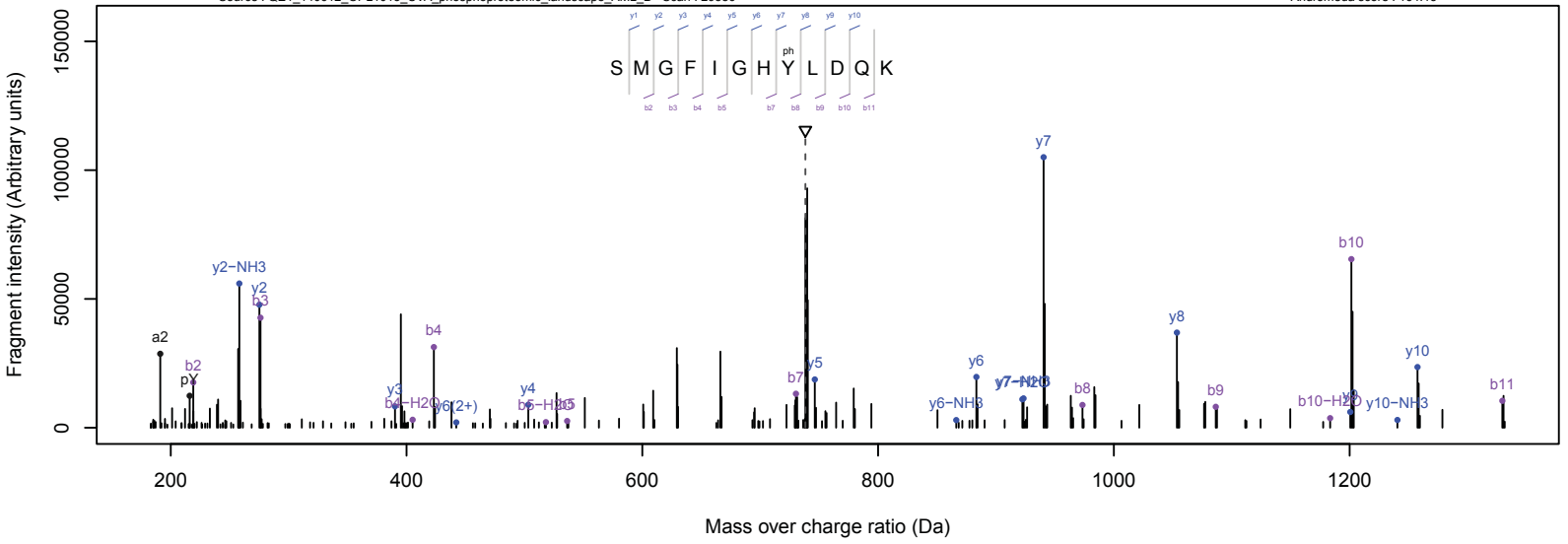
Andromeda score : 164.2



SMGFIGHY(ph)LDQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29586

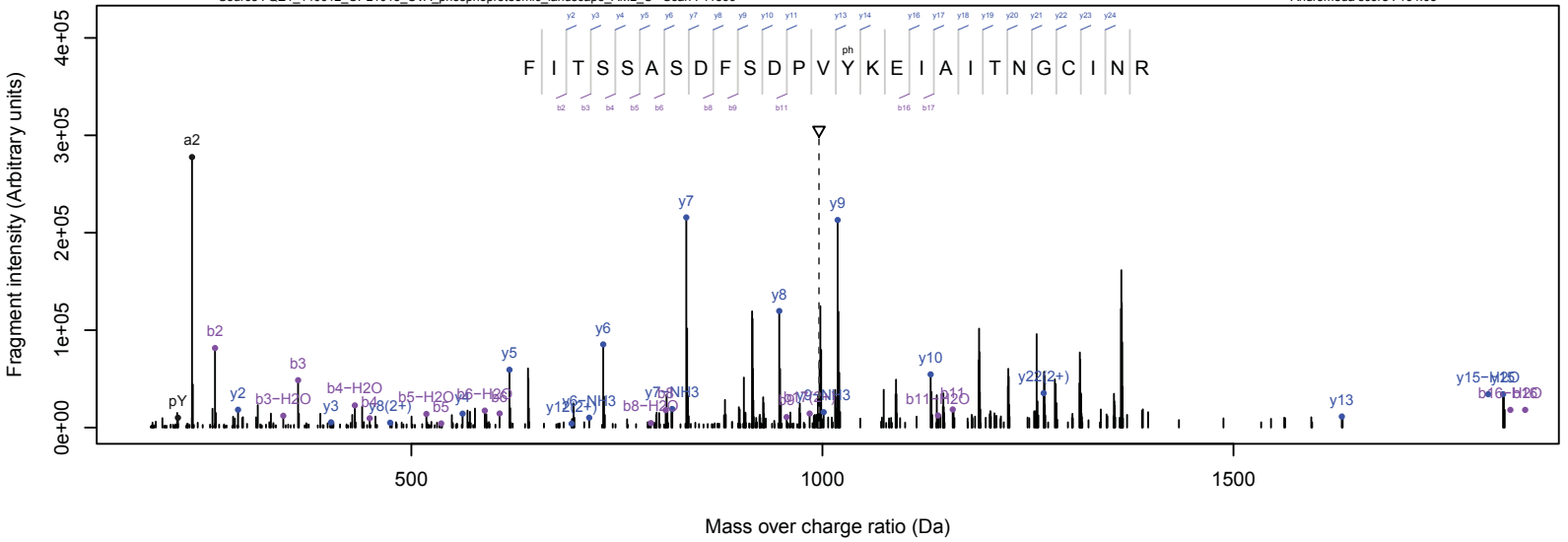
Andromeda score : 164.13



FITSSASDFSDPVY(ph)KEIAITNGCINR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41399

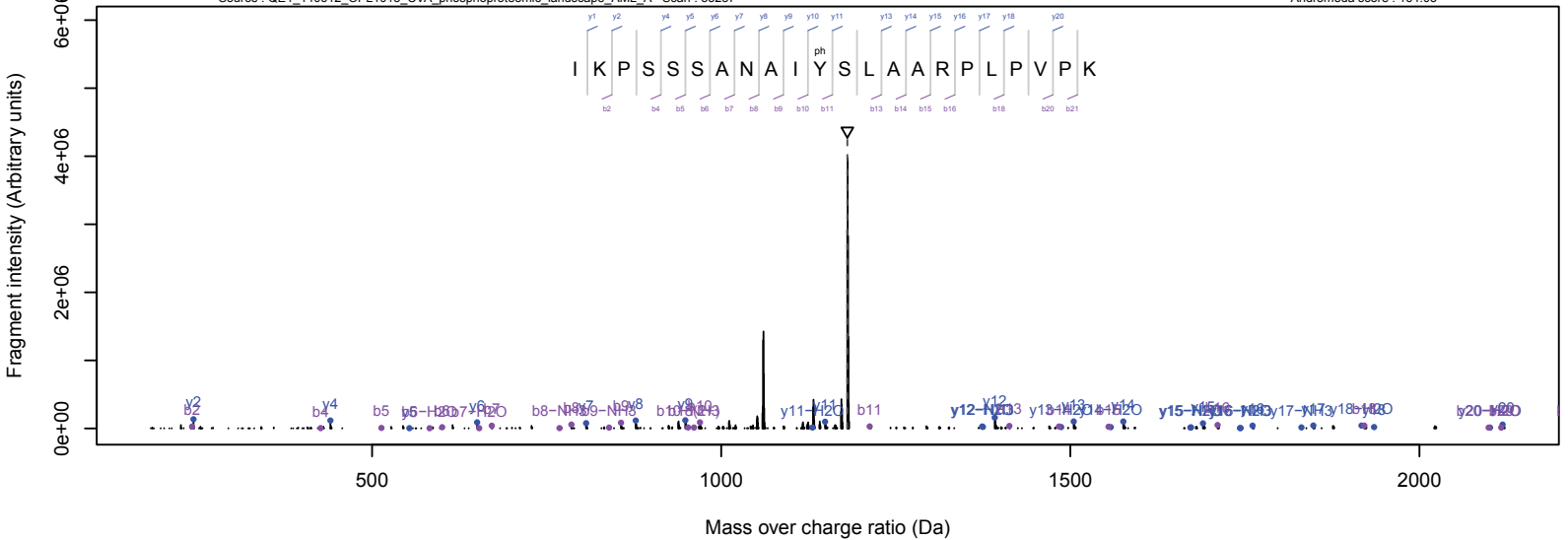
Andromeda score : 164.05



IKPSSANAIY(ph)SLAARPLVPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33257

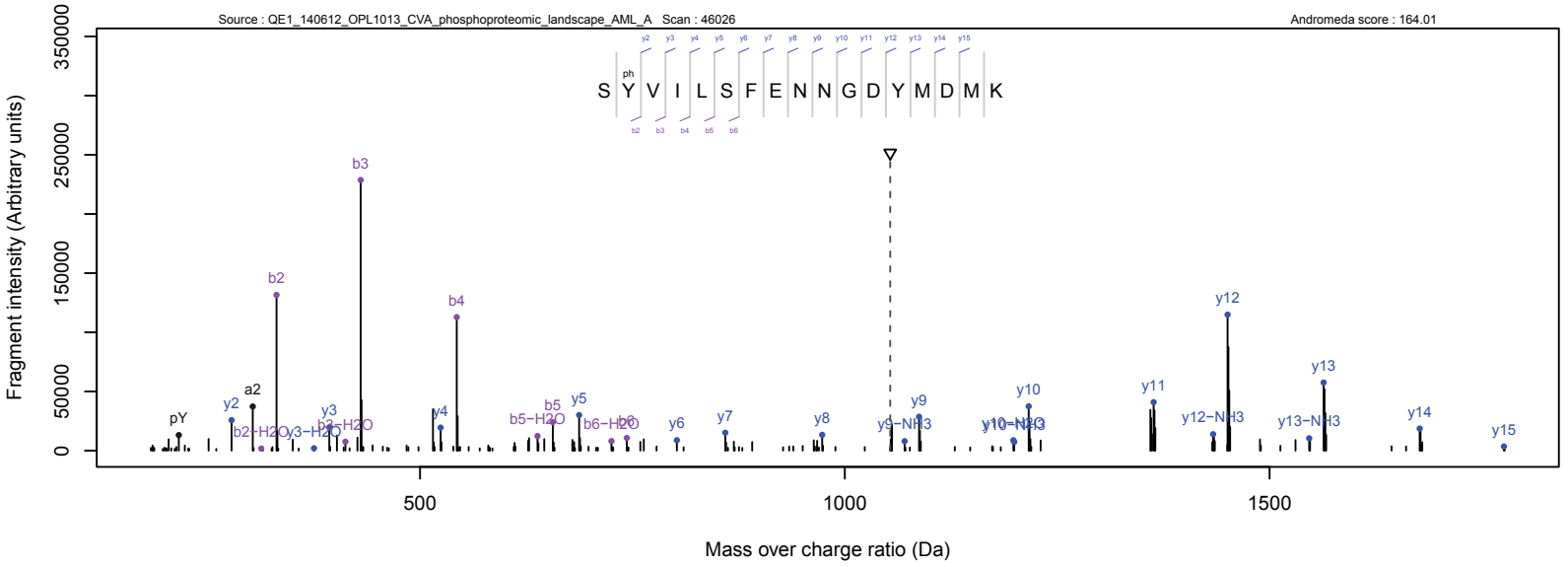
Andromeda score : 164.05



**SY(ph)VILSFENNGDYMDMK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 46026

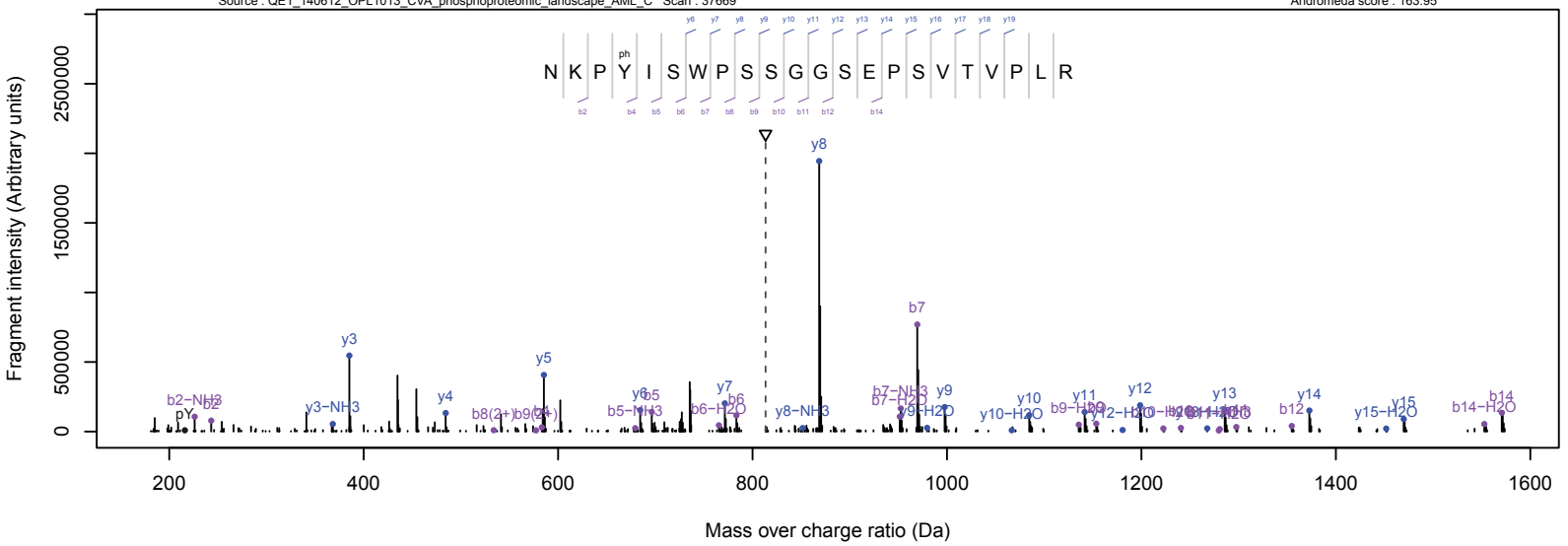
Andromeda score : 164.01



**NKPY(ph)ISWPSSGGSEPSVTVPLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37669

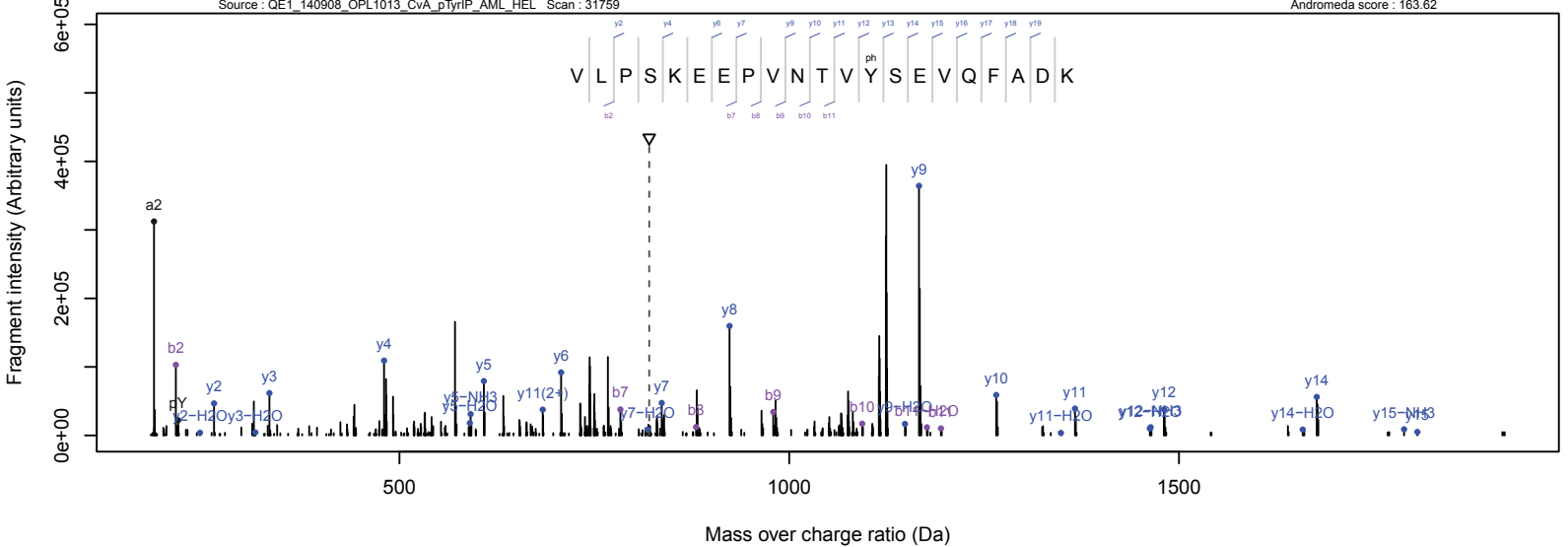
Andromeda score : 163.95



**VLPKEEPVNTVY(ph)SEVQFADK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31759

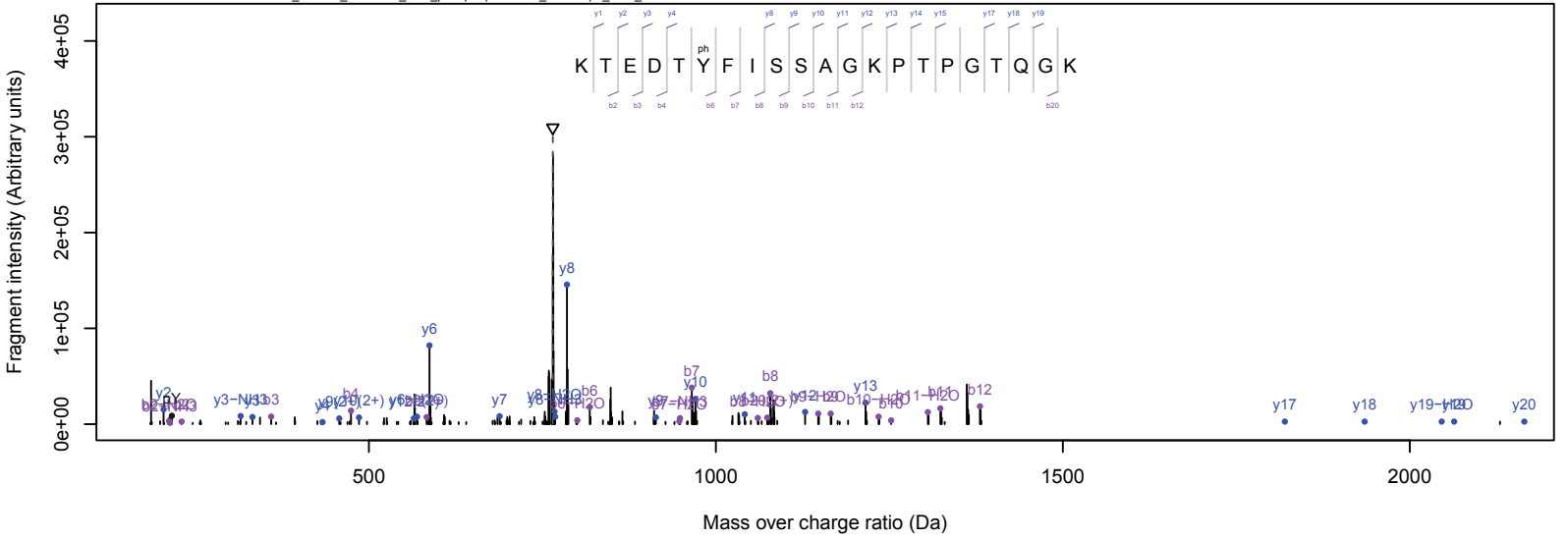
Andromeda score : 163.62



**KTEDTY(ph)FISSAGKPTPGTGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15489

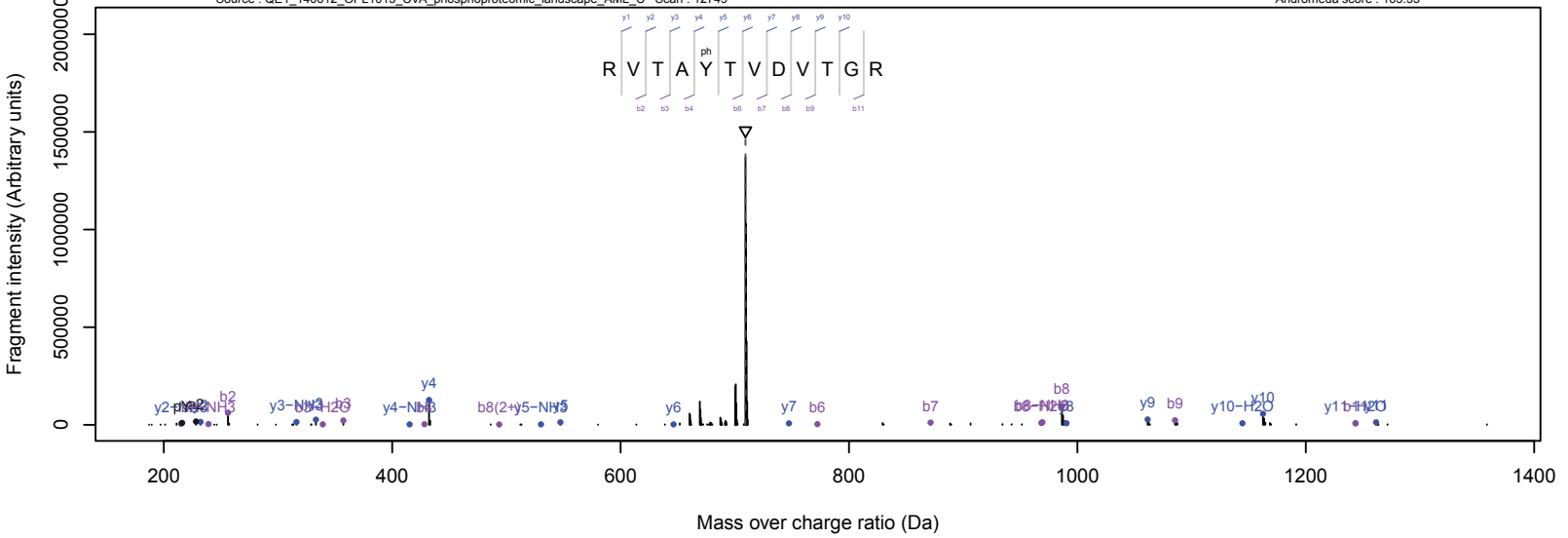
Andromeda score : 163.49



**RVTAY(ph)TVDTVTR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12749

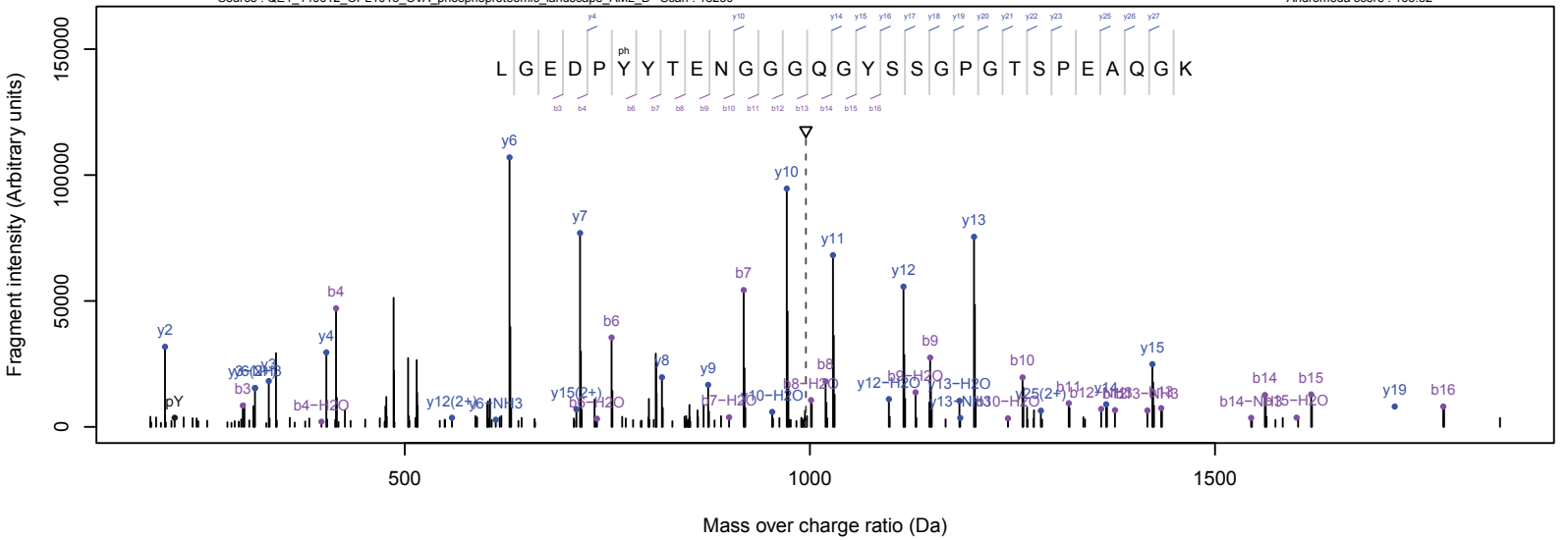
Andromeda score : 163.33



**LGEDPY(ph)YTENGGQGYS SGP GTSPEA QGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18250

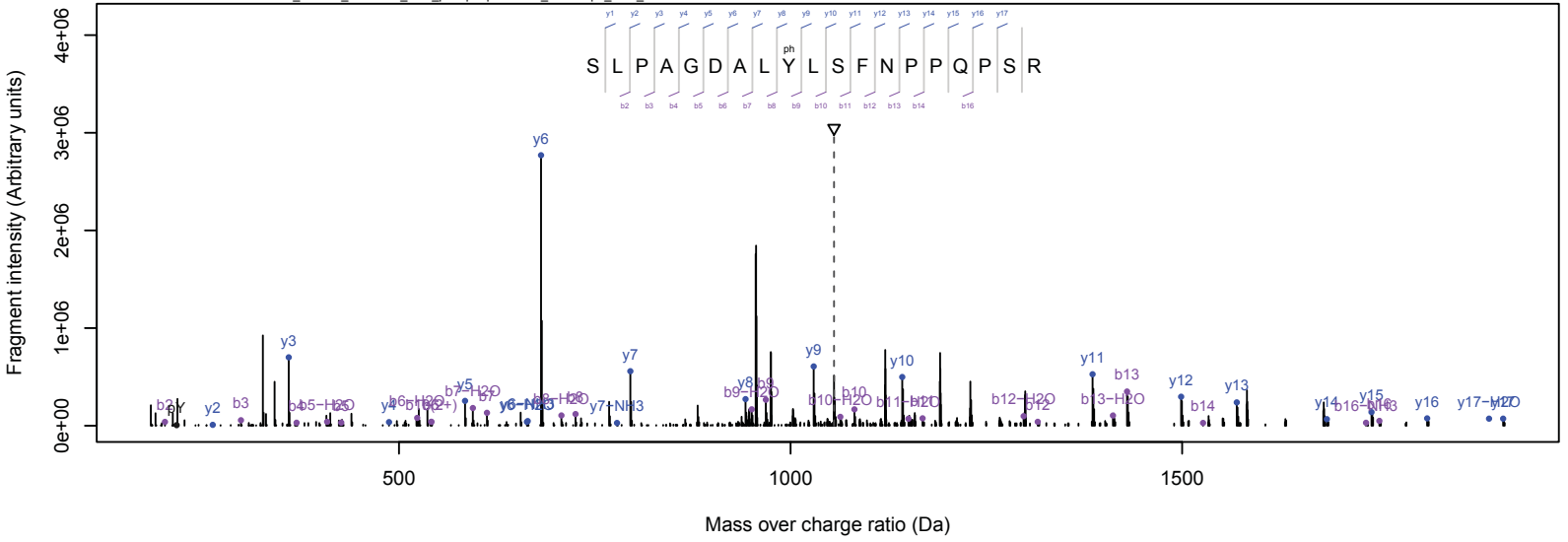
Andromeda score : 163.32



SLPAGDALY(ph)LSFNPPQPSR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 43724

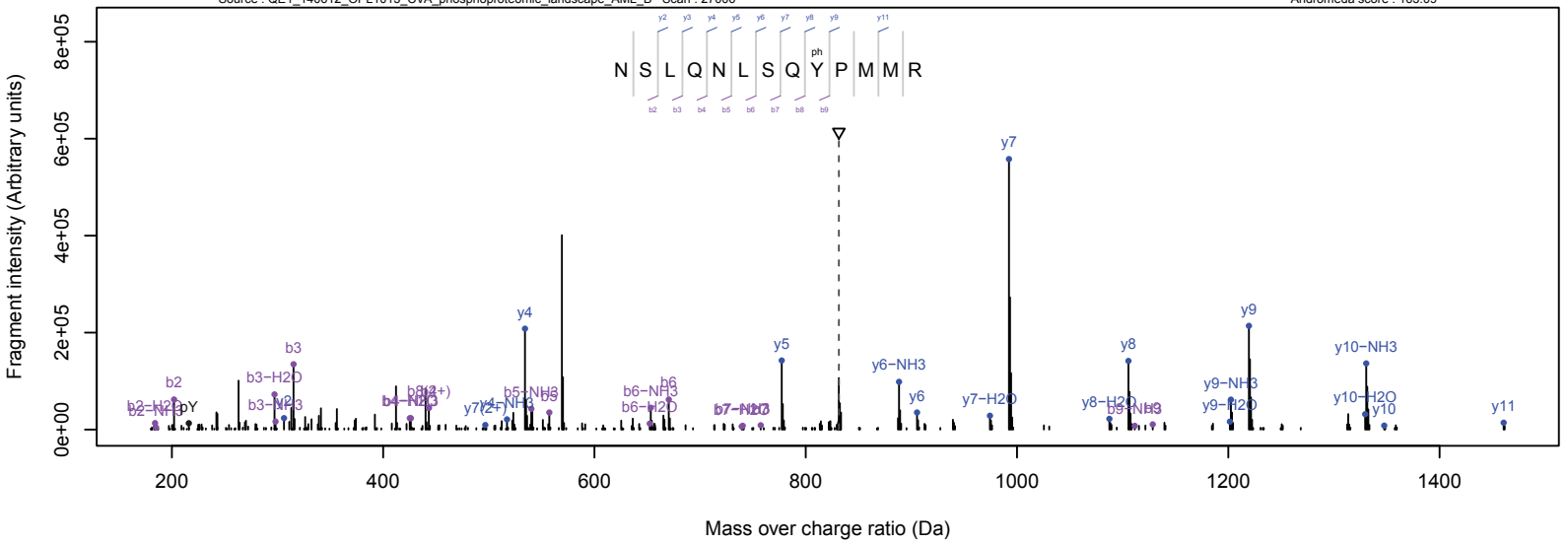
Andromeda score: 163.15



NSLQNLQY(ph)PMMR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27006

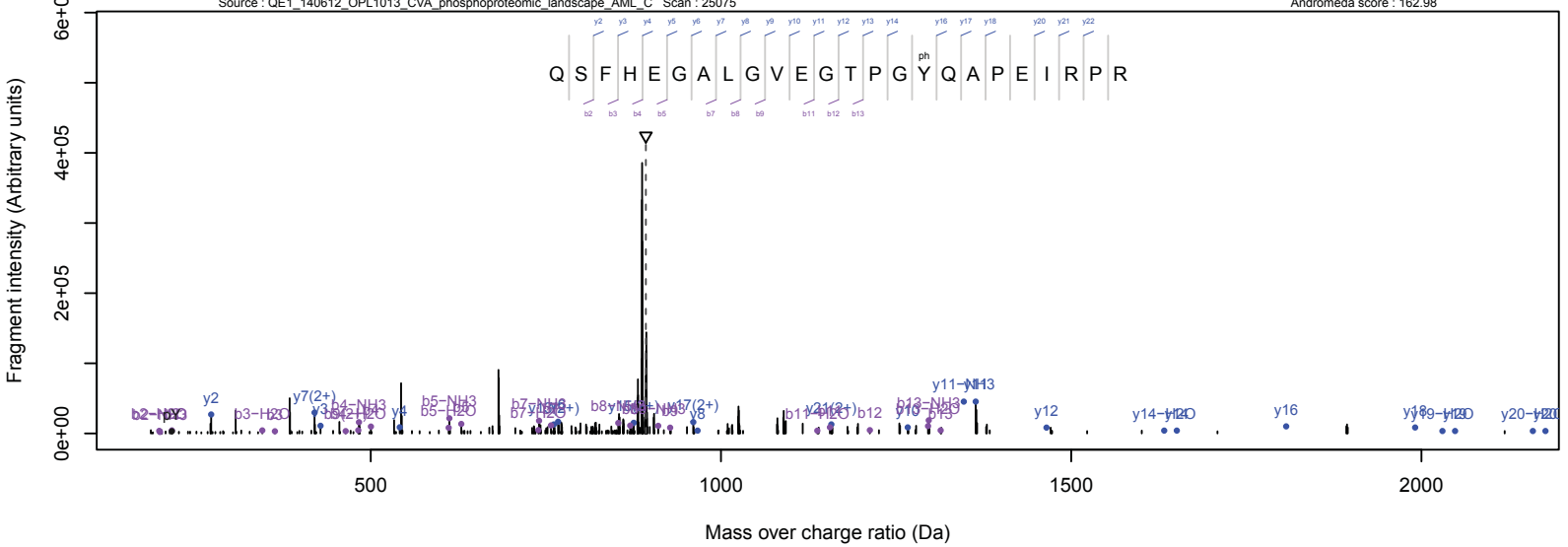
Andromeda score: 163.09



QSFHEGALGVETPGYQAPEIRP-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25075

Andromeda score: 162.98

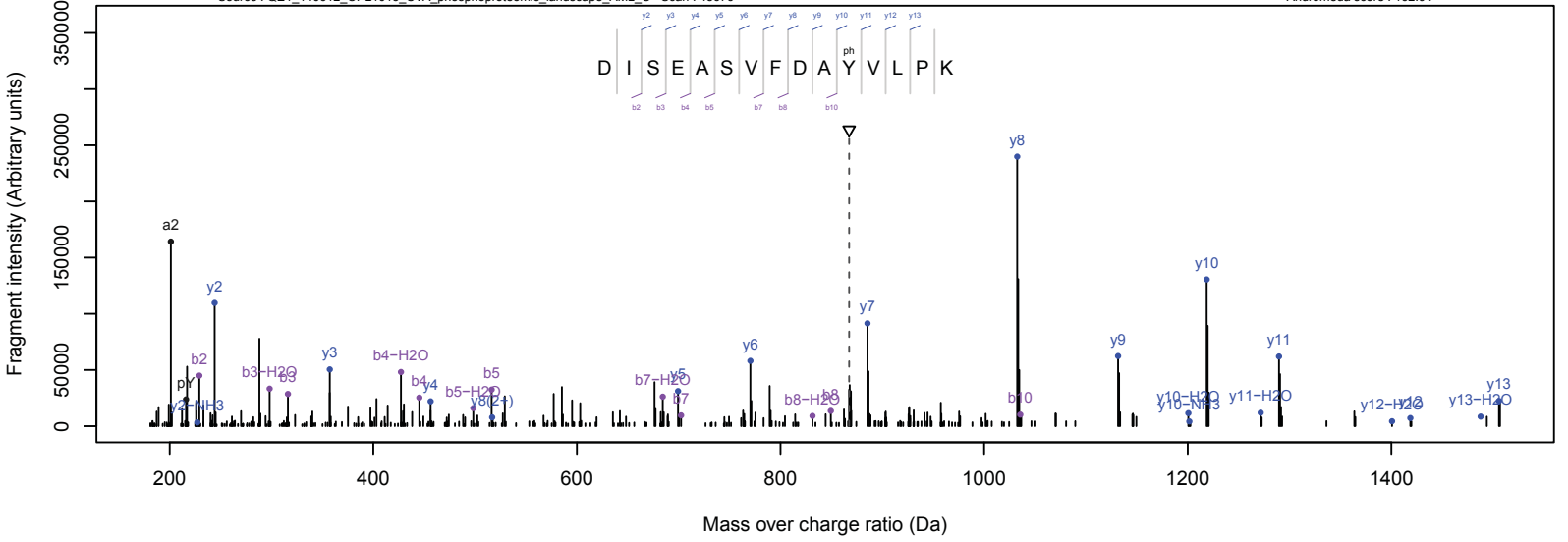




DISEASVFDAYVLPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 45676

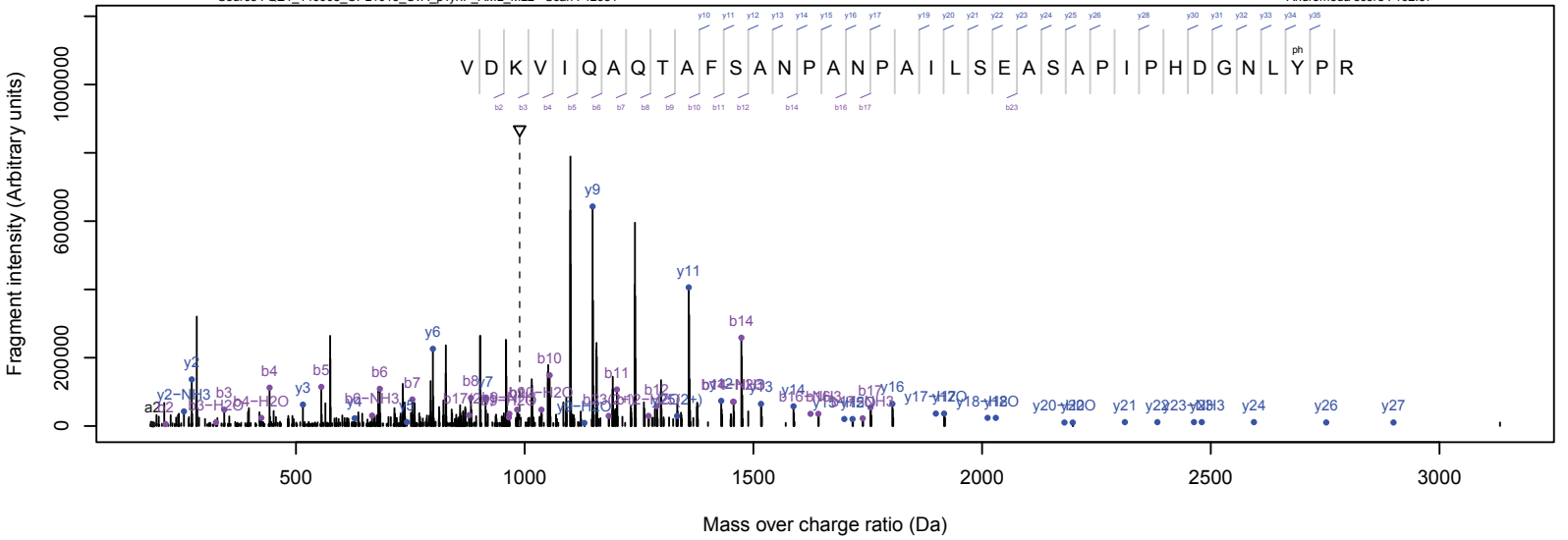
Andromeda score: 162.94



VDKVIQAQTAFSANPANPAILSEASAPIPHDGNLYPR-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan: 42381

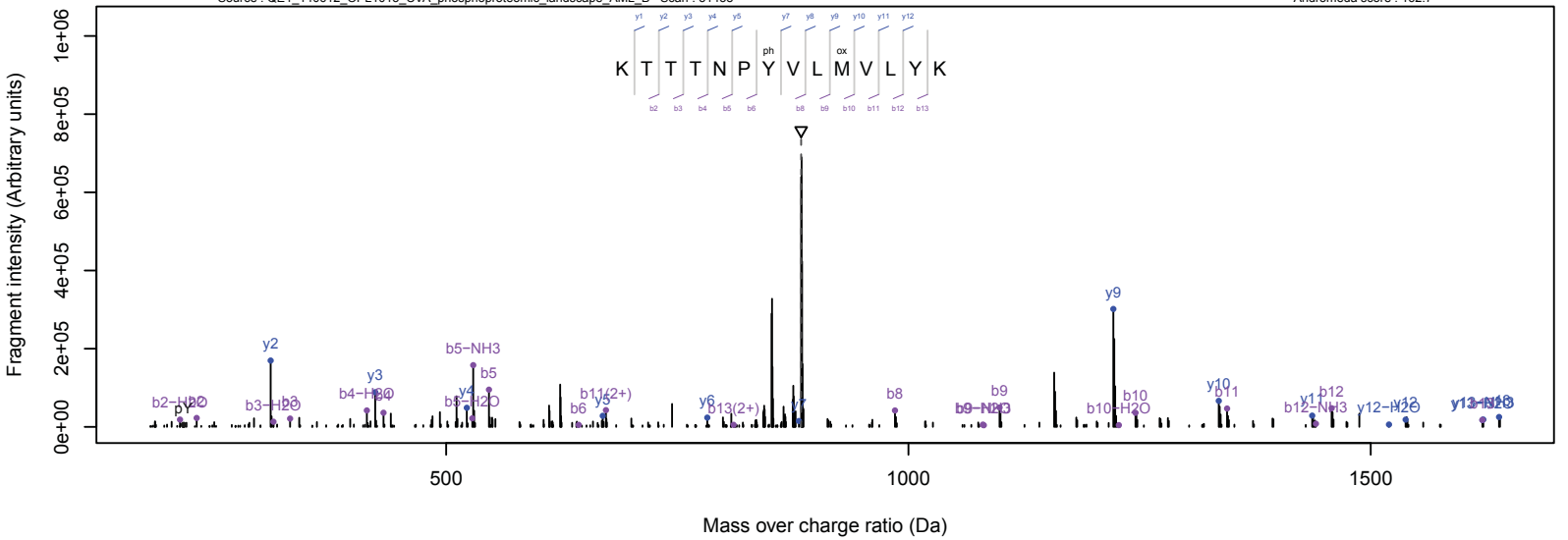
Andromeda score: 162.87



KTTTNPYVLMVLYK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 31438

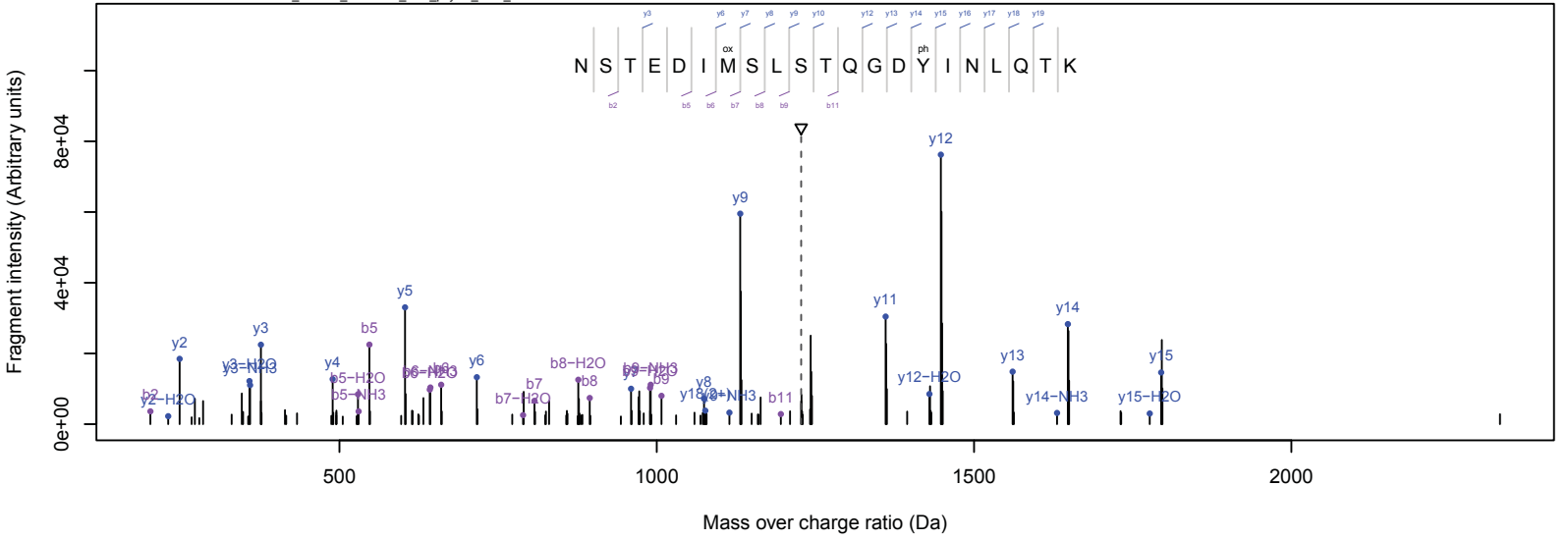
Andromeda score: 162.7



**NSTEDIM(ox)SLSTQGDY(ph)INLQTK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 30682

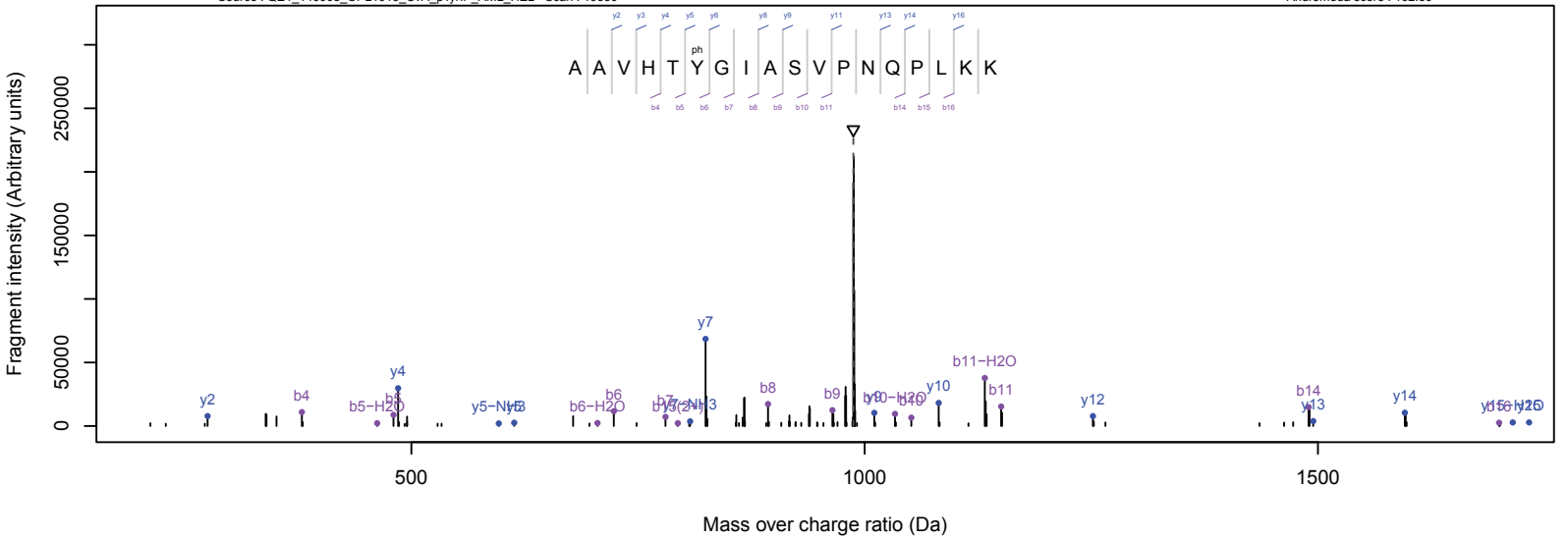
Andromeda score : 162.65



**AAVHTY(ph)GIASVNPQLKK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 19830

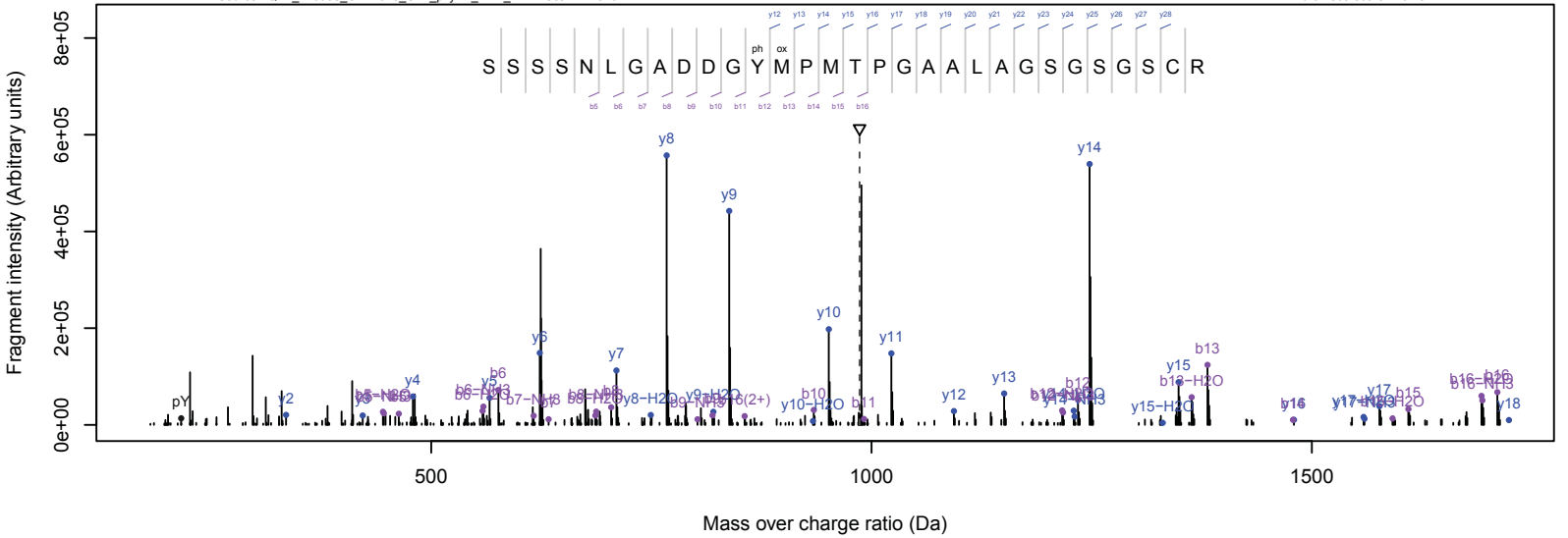
Andromeda score : 162.59



**SSSNLGADDGY(ph)M(ox)PMTPGAALAGSGSGSCR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 27576

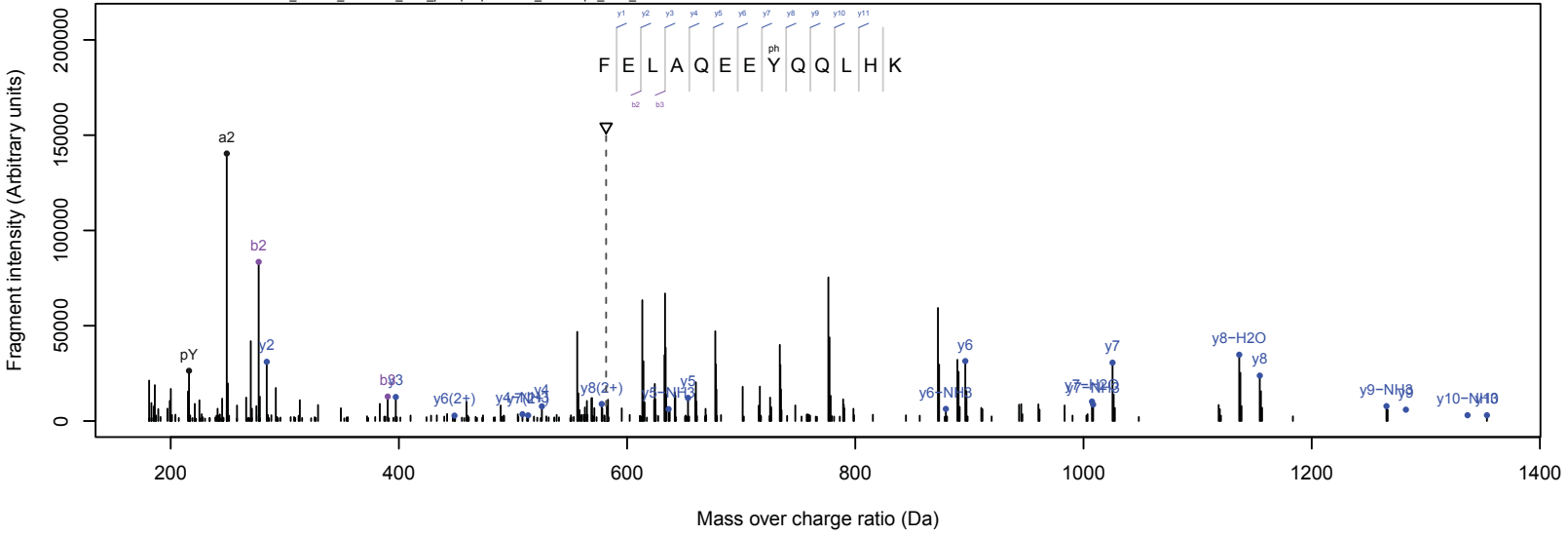
Andromeda score : 162.52



FELAQEEY(ph)QLHK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 23336

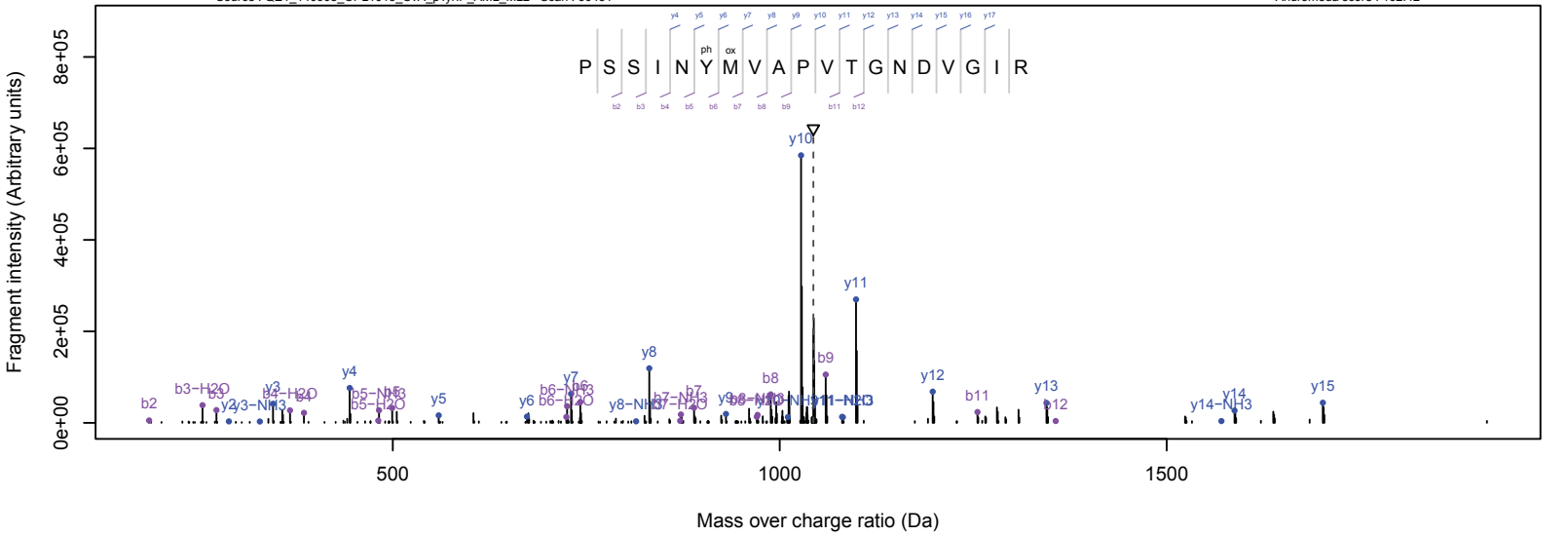
Andromeda score: 162.49



PSSINY(ph)M(ox)VAPVTGNDVGIR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan: 30431

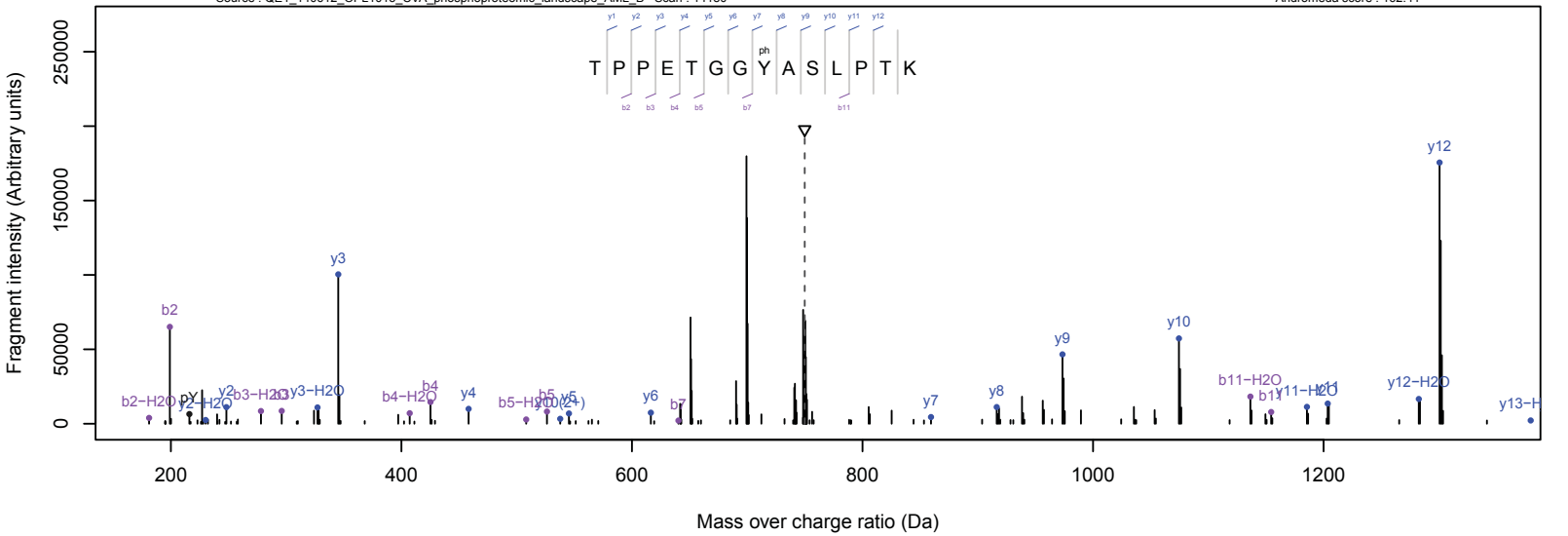
Andromeda score: 162.42



TPPETGGY(ph)ASLPTK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14180

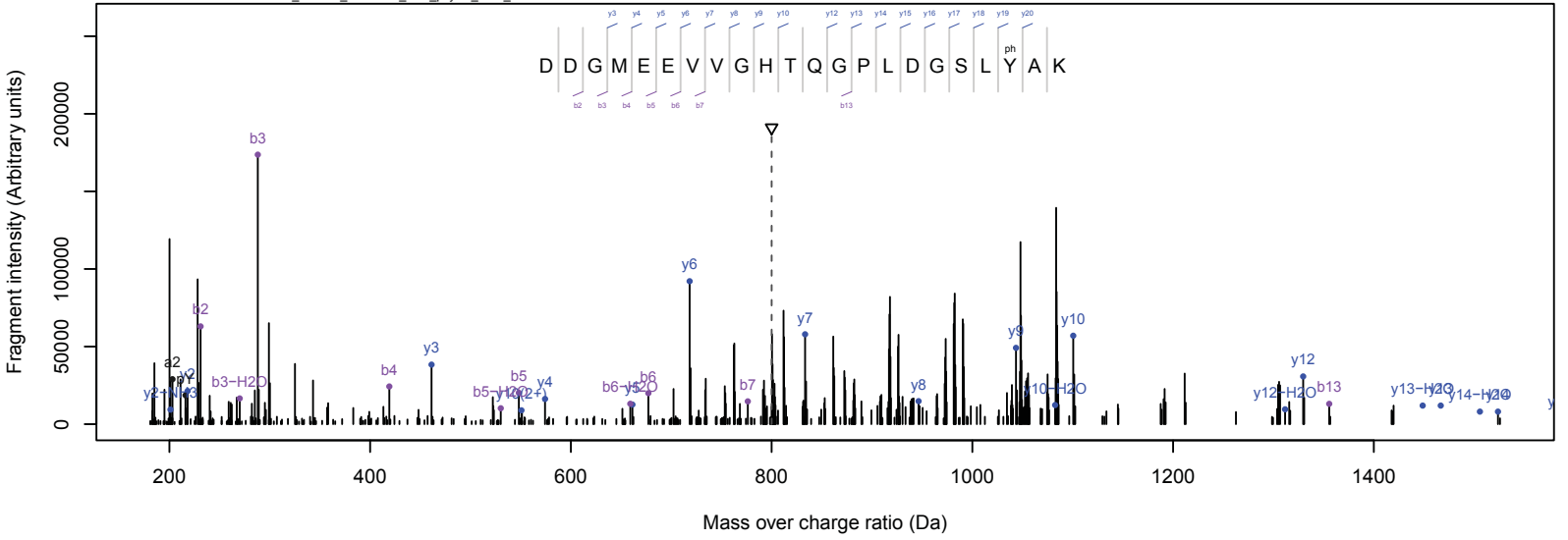
Andromeda score: 162.41



DDGMEEVVGHTQGPLDGSLYAK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35179

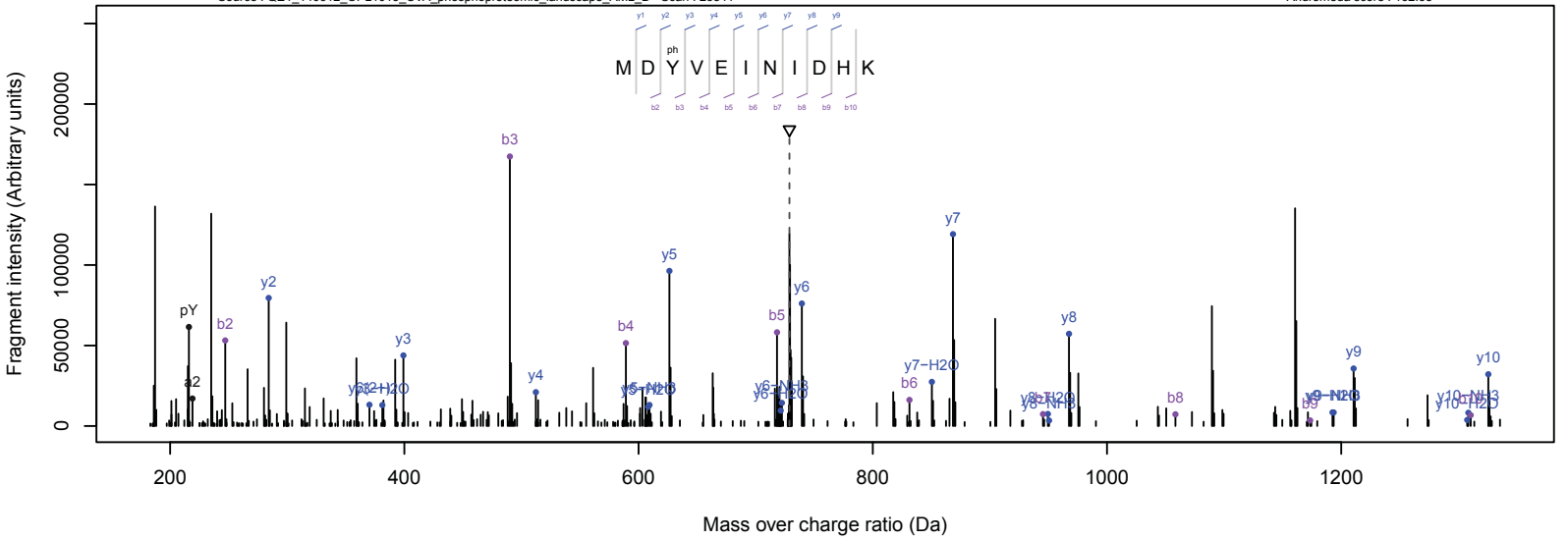
Andromeda score : 162.39



MDY(ph)VEINIDHK-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan : 23841

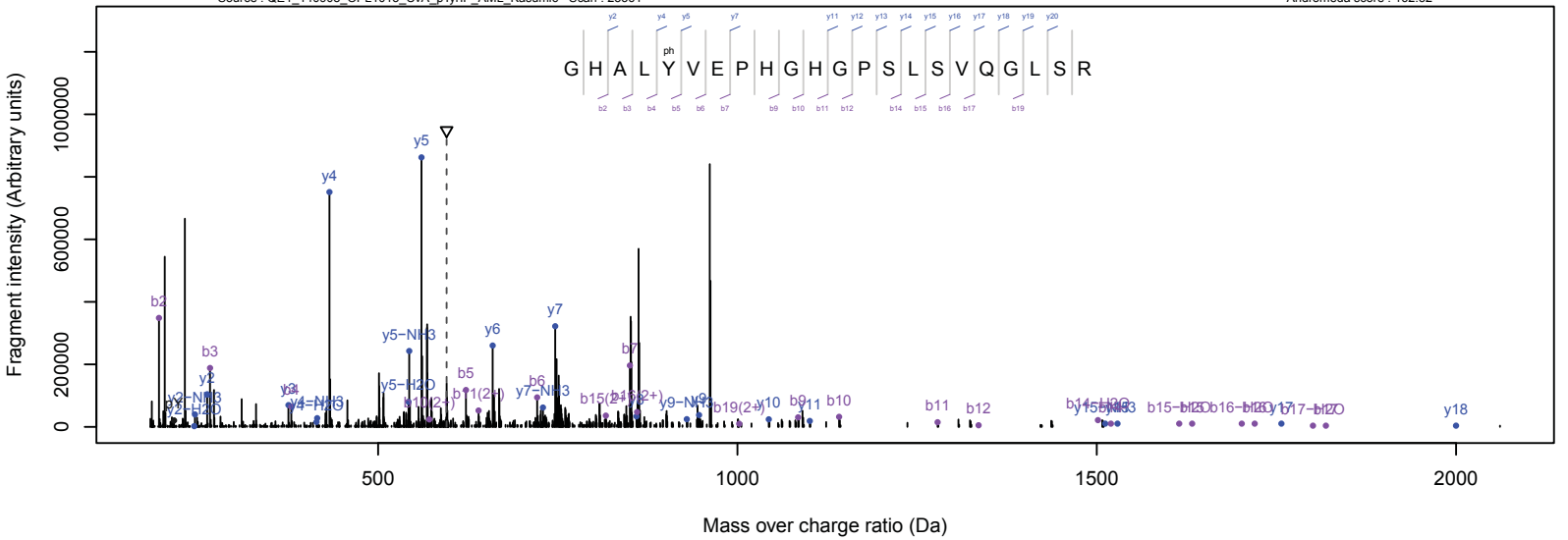
Andromeda score : 162.38



GHALY(ph)VEPHGHGPSLSVQGLSR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 25361

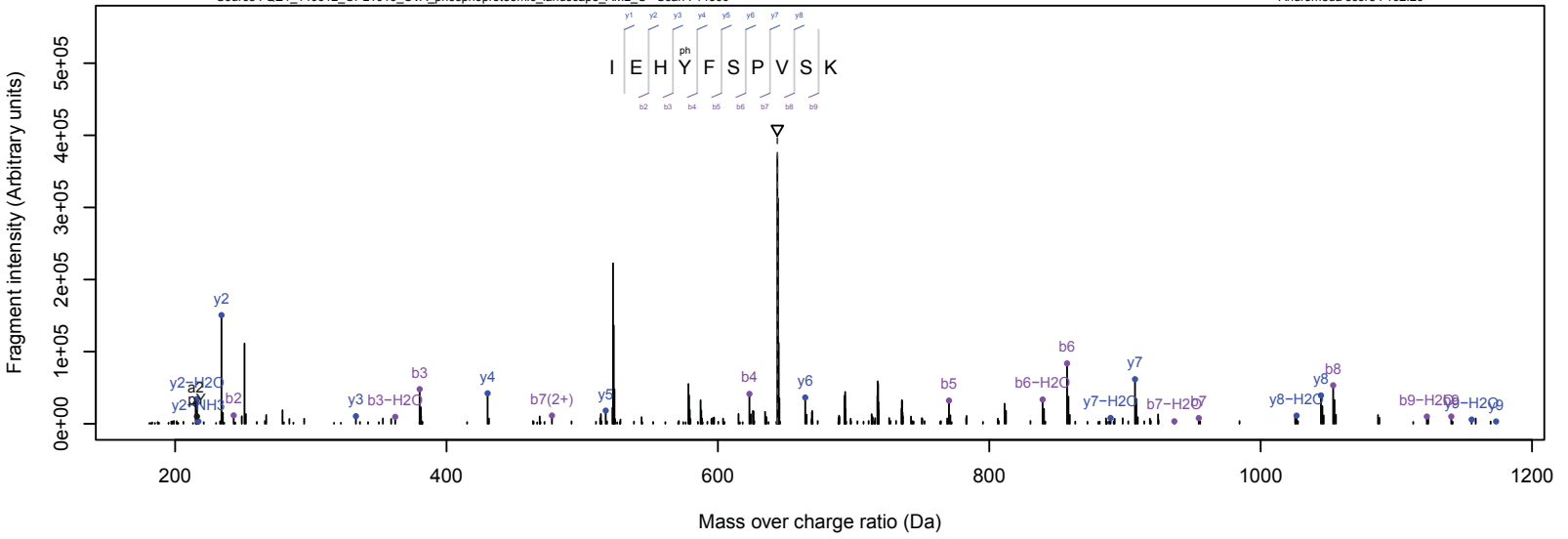
Andromeda score : 162.32



IEHY(ph)FSPVSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14509

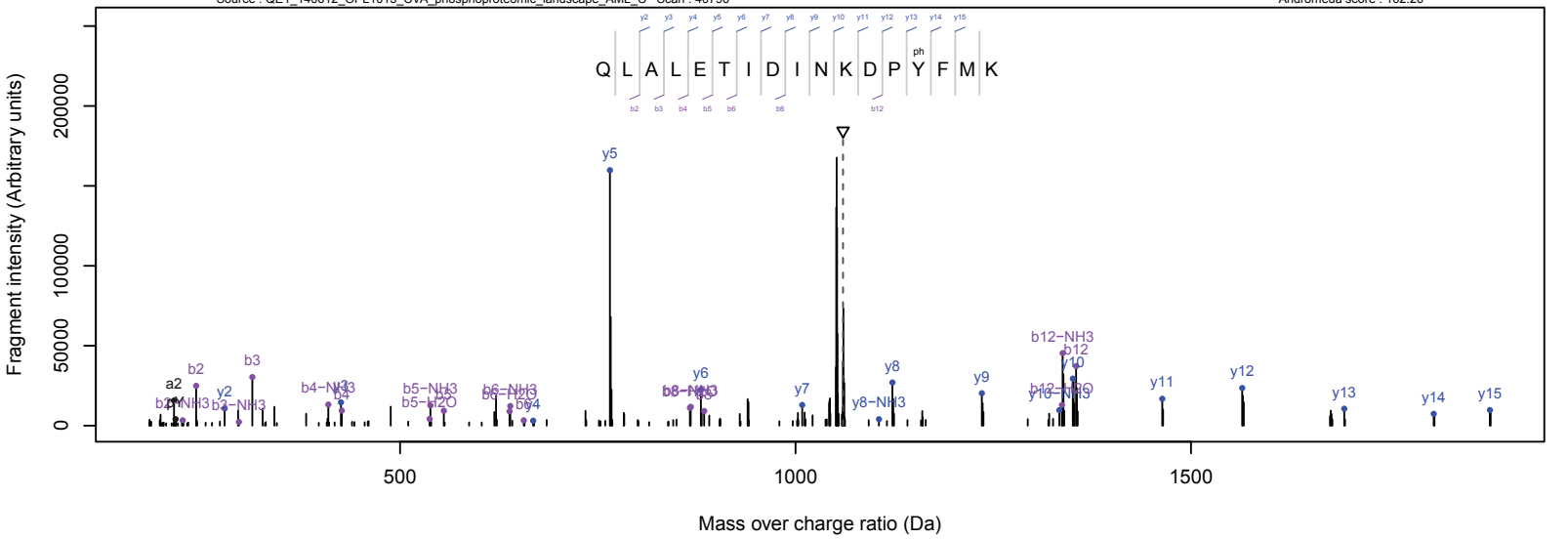
Andromeda score : 162.26



QLALETIDINKDPY(ph)FMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40796

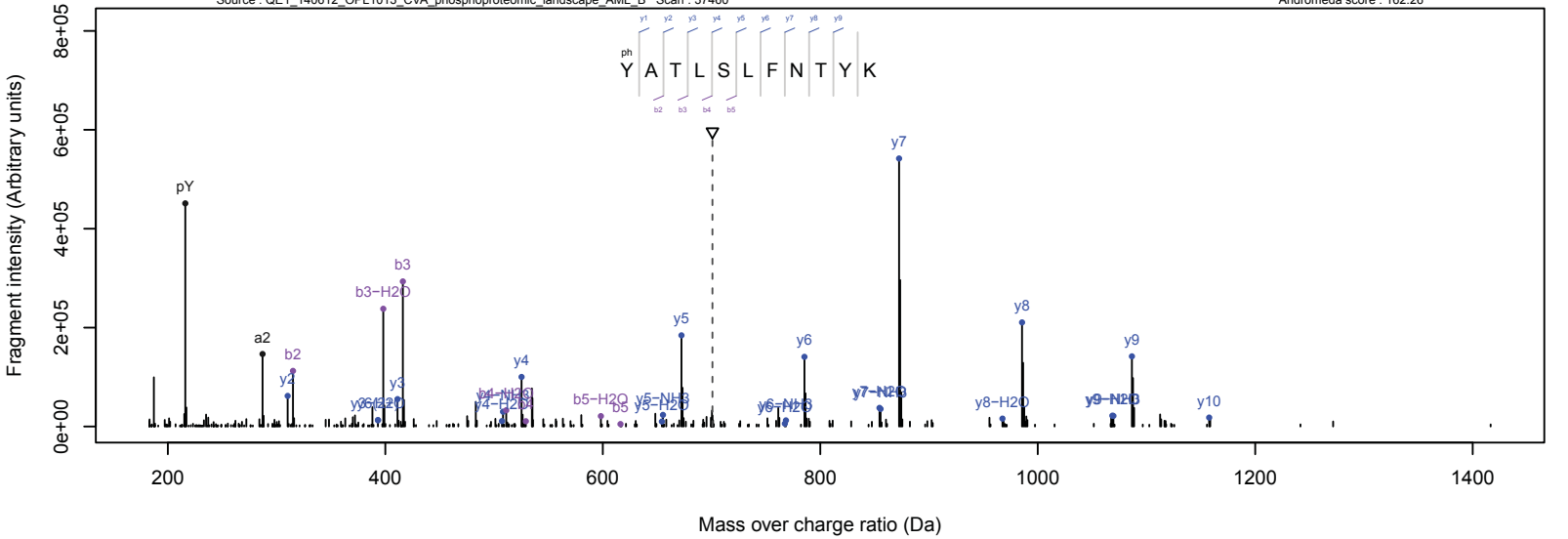
Andromeda score : 162.26



Y(ph)ATLSLFNTYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37460

Andromeda score : 162.26

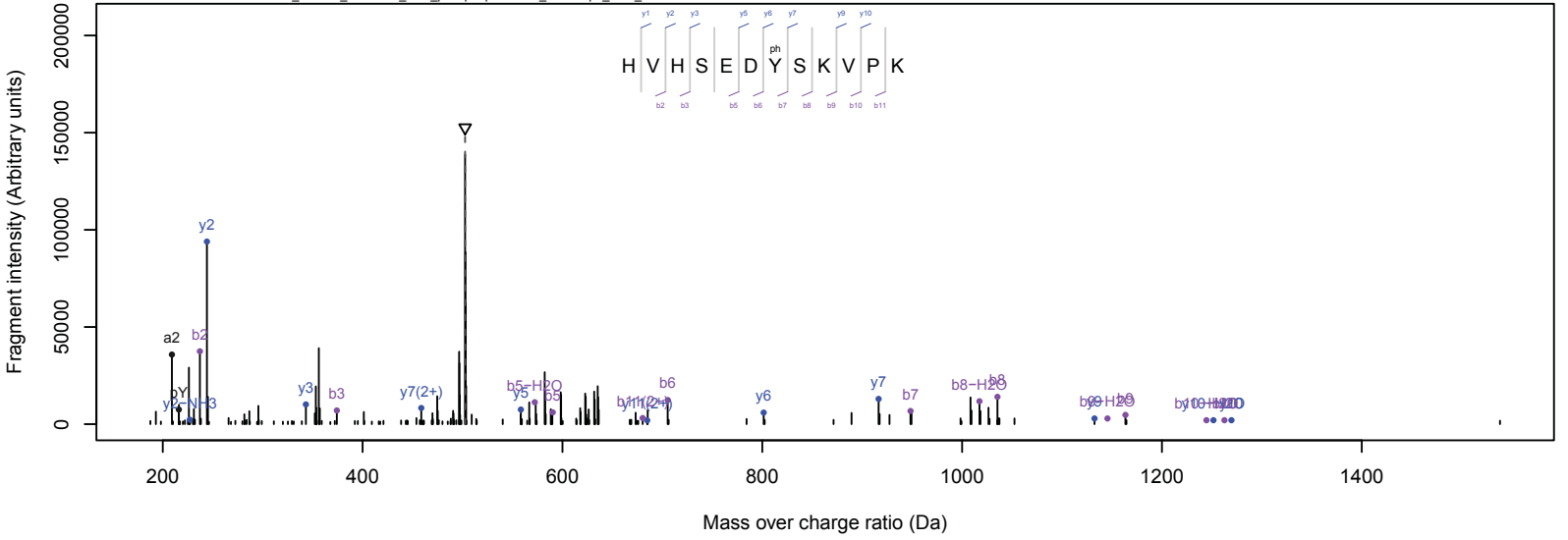




HVHSEDY(ph)SKVPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 3627

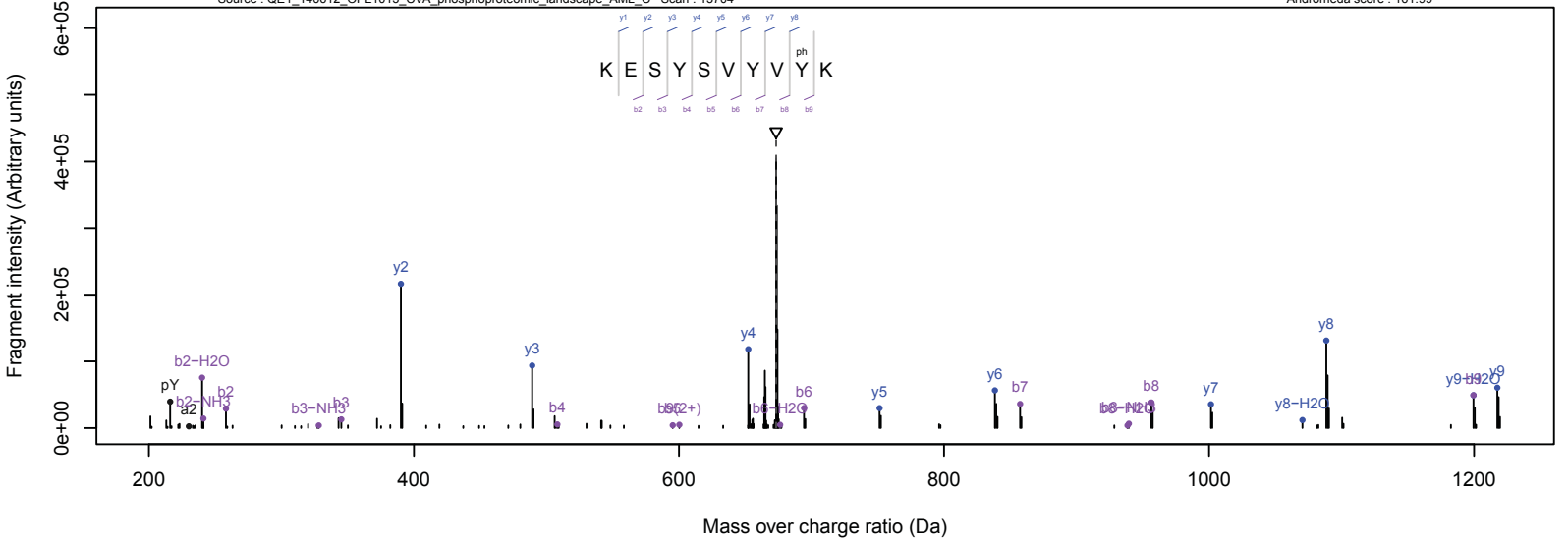
Andromeda score : 161.99



KESYSVYVY(ph)K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13704

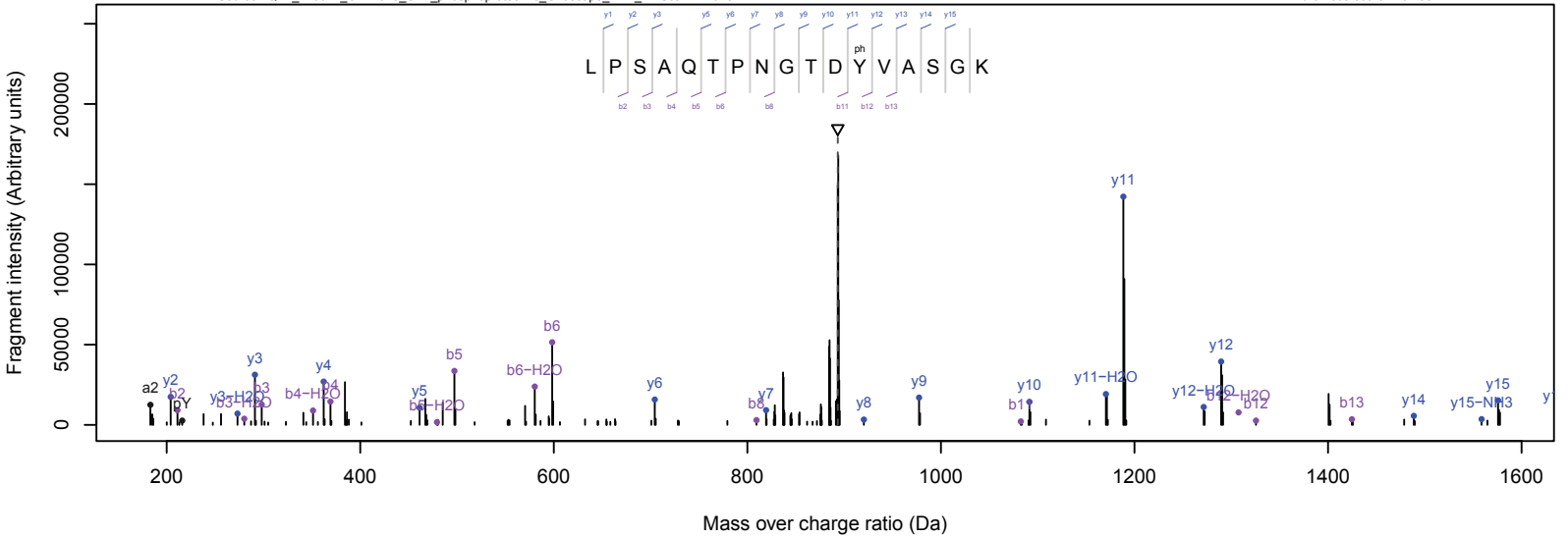
Andromeda score : 161.99



LPSAQTPTNGTDY(ph)VASGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14915

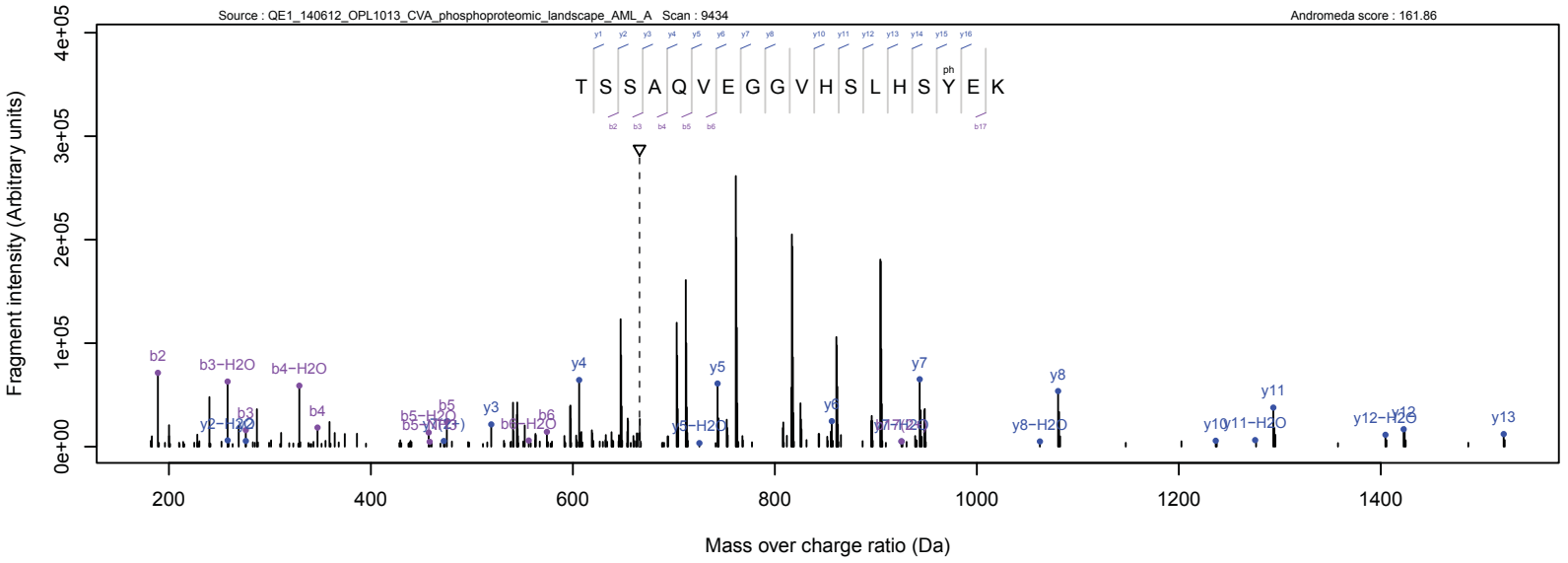
Andromeda score : 161.86



**TSSAQVEGGVHSLHSY(ph)EK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 9434

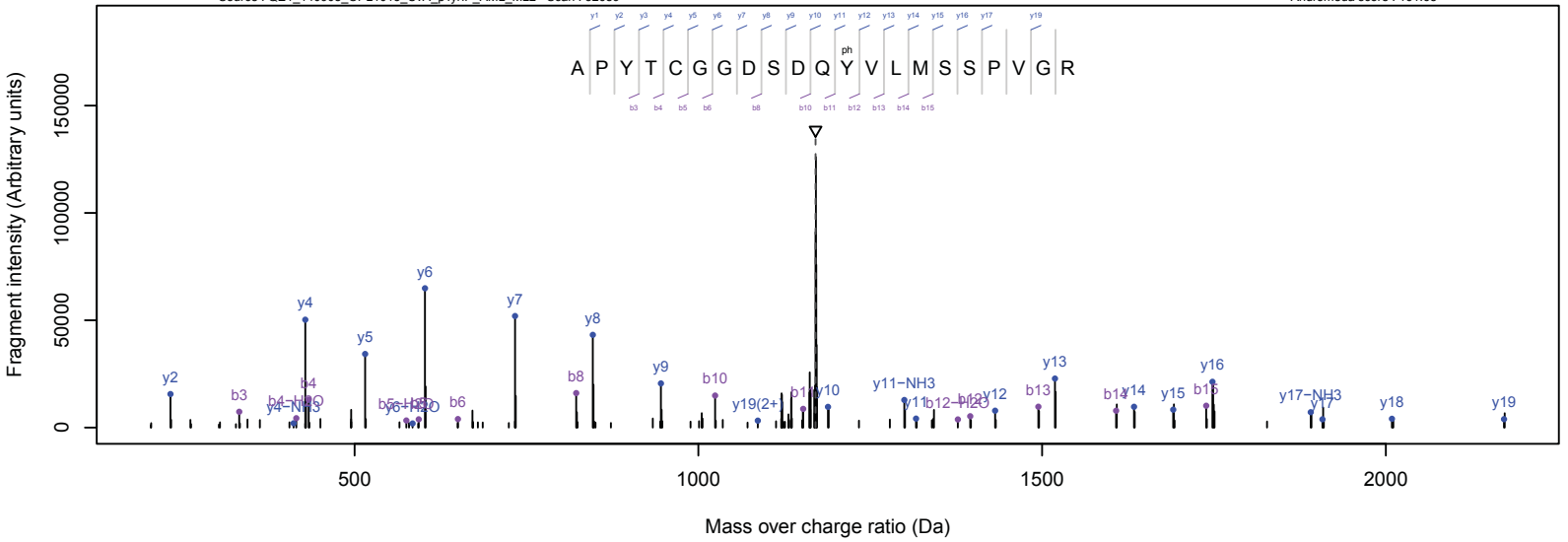
Andromeda score : 161.86



**APYTCGGDSDQY(ph)VLMSSPVG R-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 32689

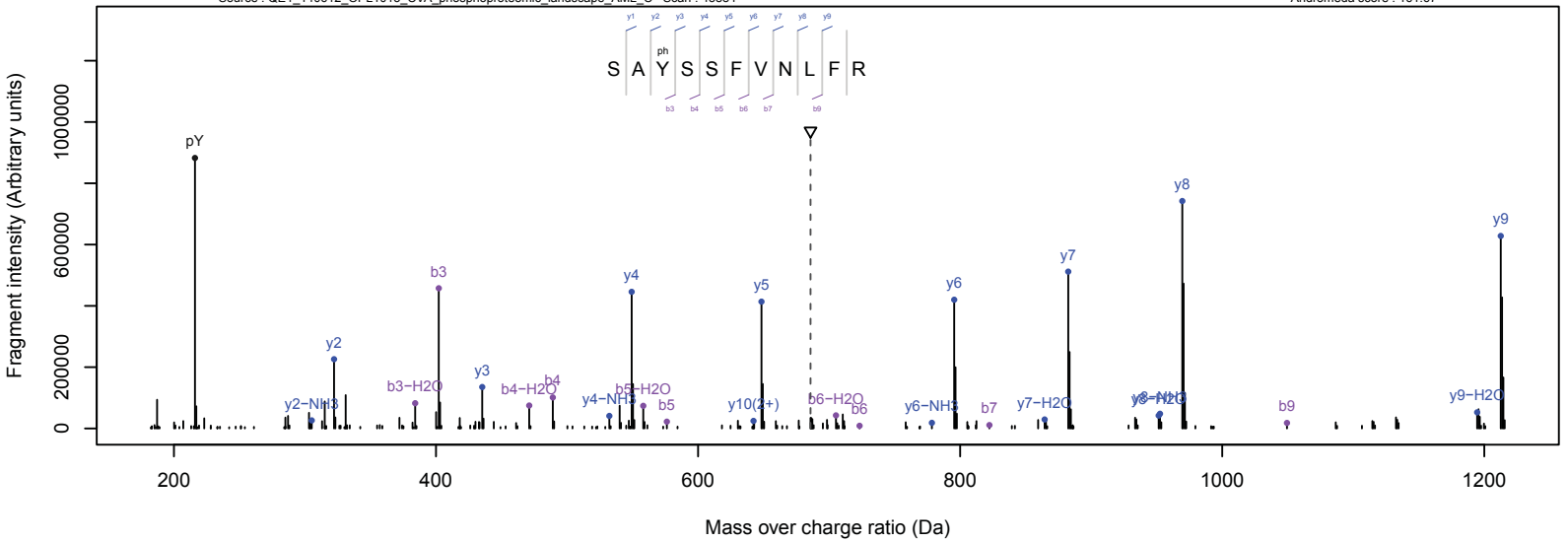
Andromeda score : 161.68



**SAY(ph)SSFVNLFR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45884

Andromeda score : 161.67

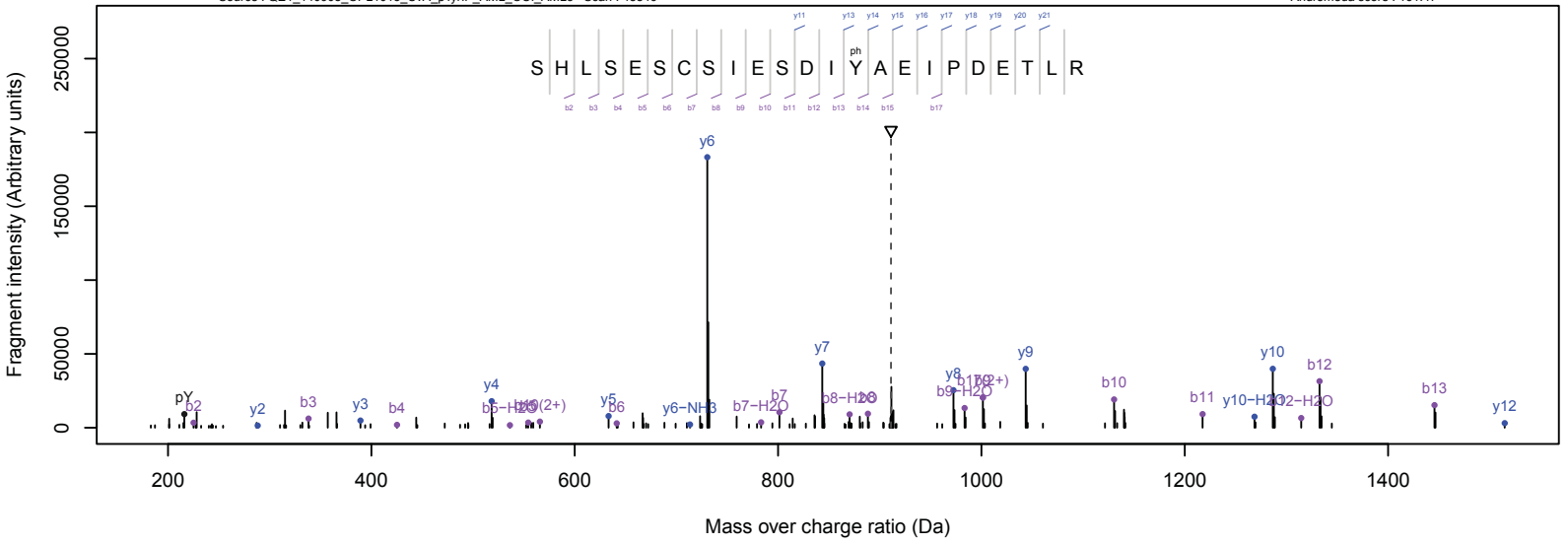




SHLSESCSI ESDIY A E I P D E T L R-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_OCI\_AML3 Scan : 48846

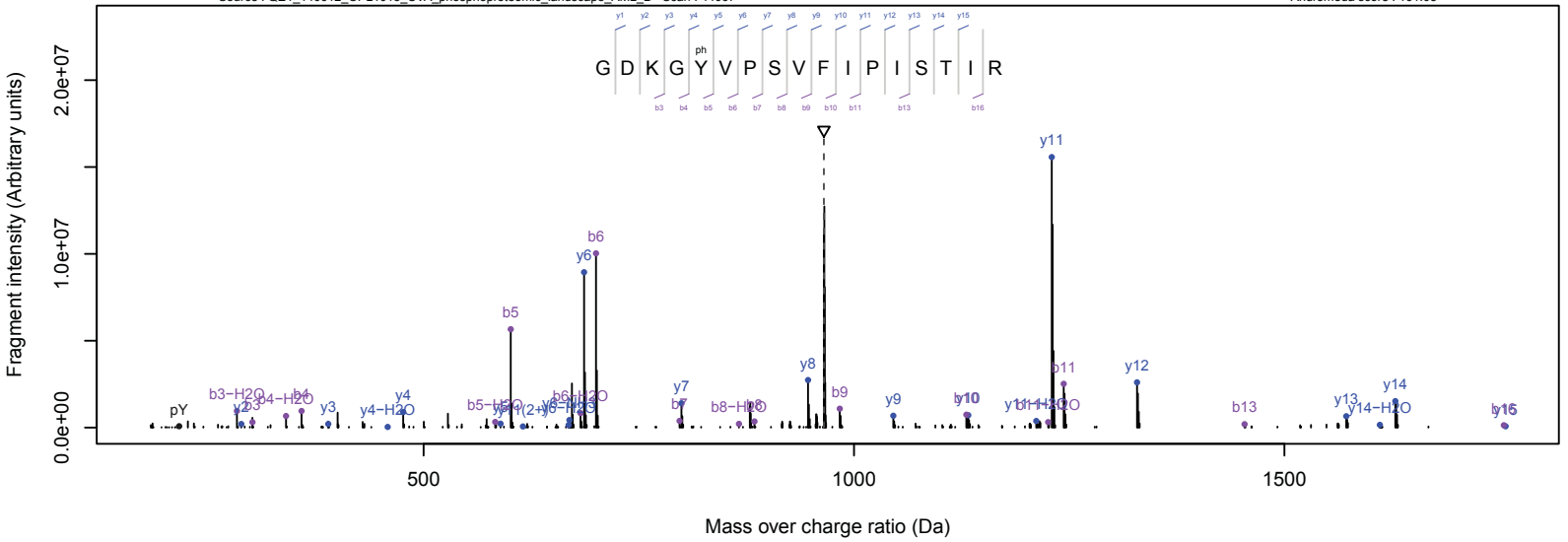
Andromeda score : 161.47



GDKGY(ph) VPSVFIPISTIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44067

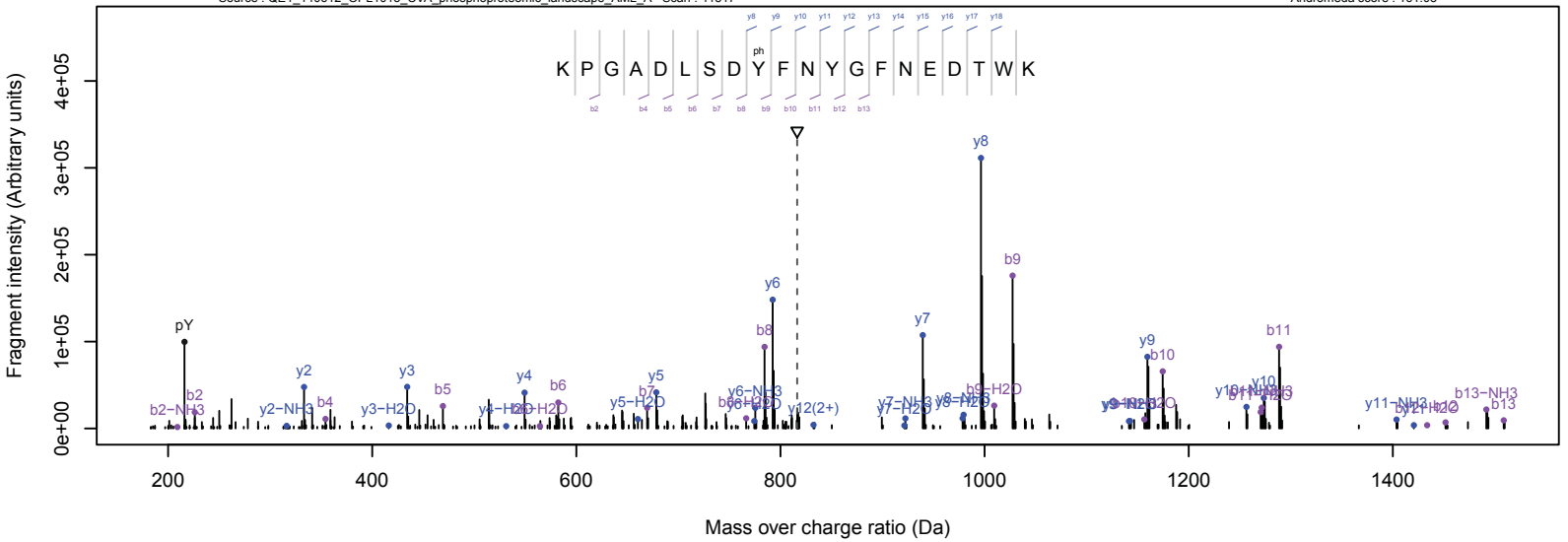
Andromeda score : 161.08



KPGADLSDY(ph) FNYGFNEDTWK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 41517

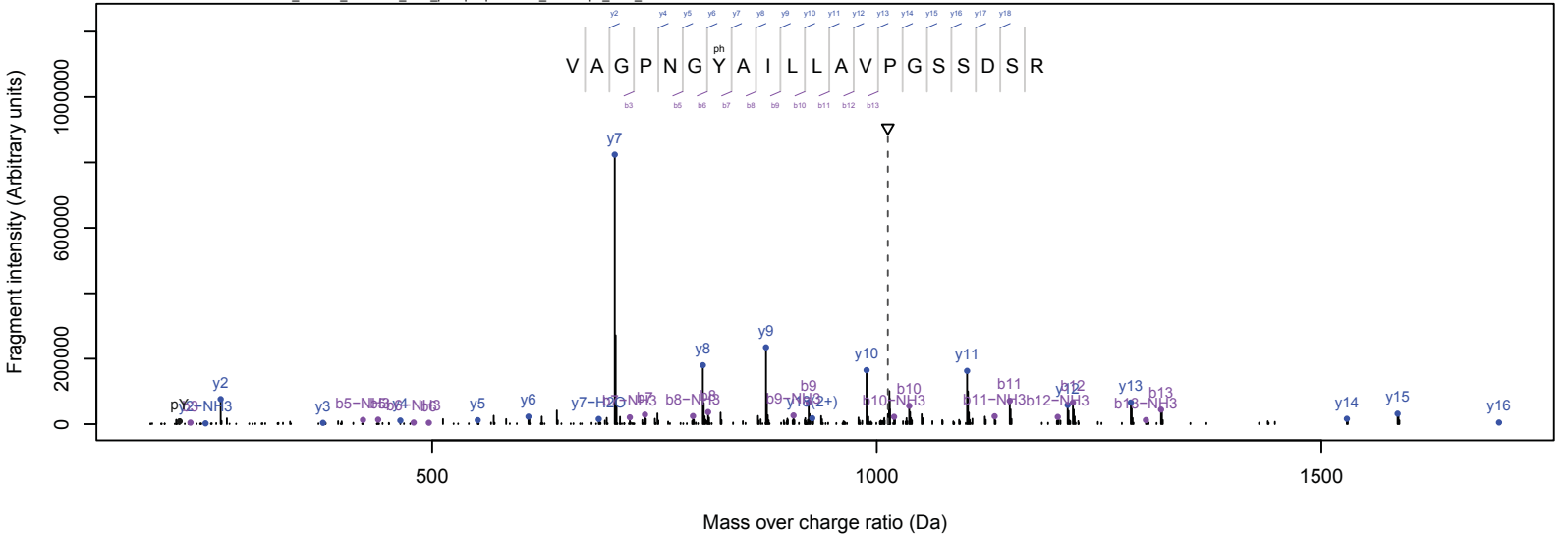
Andromeda score : 161.08



VAGPNGY(ph)AILLAVPGSSDSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40250

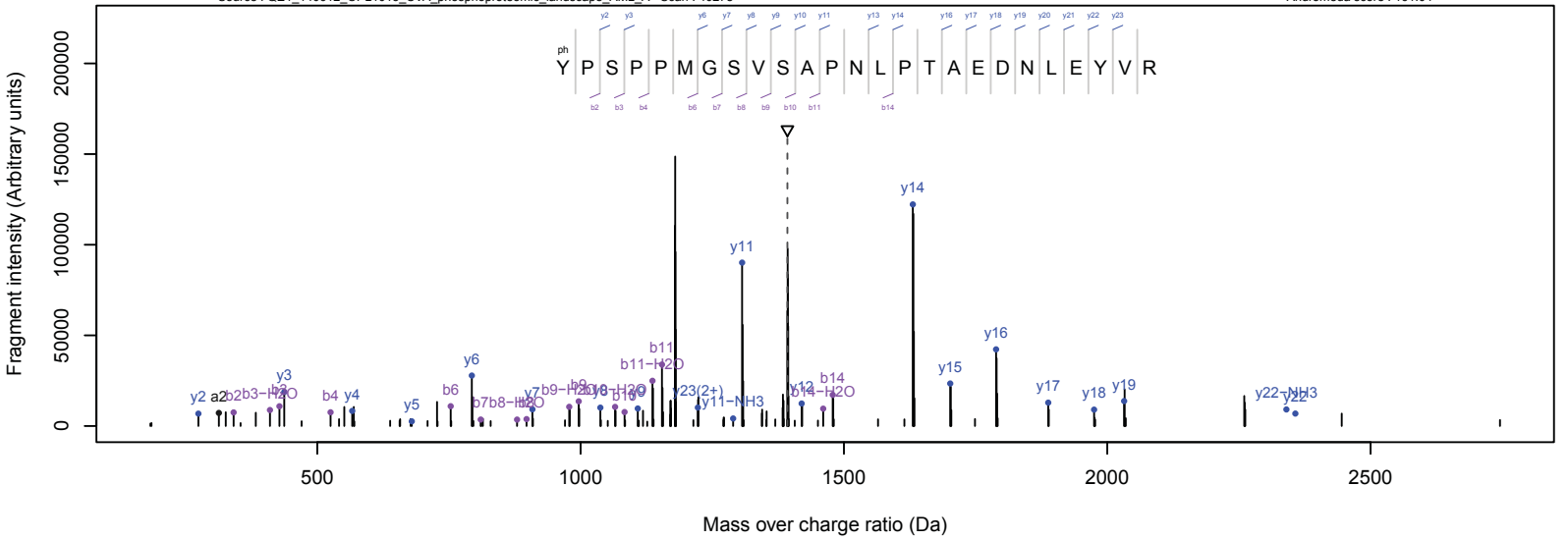
Andromeda score : 161.08



Y(ph)PSPPMGVSAPNLPPTAEDNLEYVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 40278

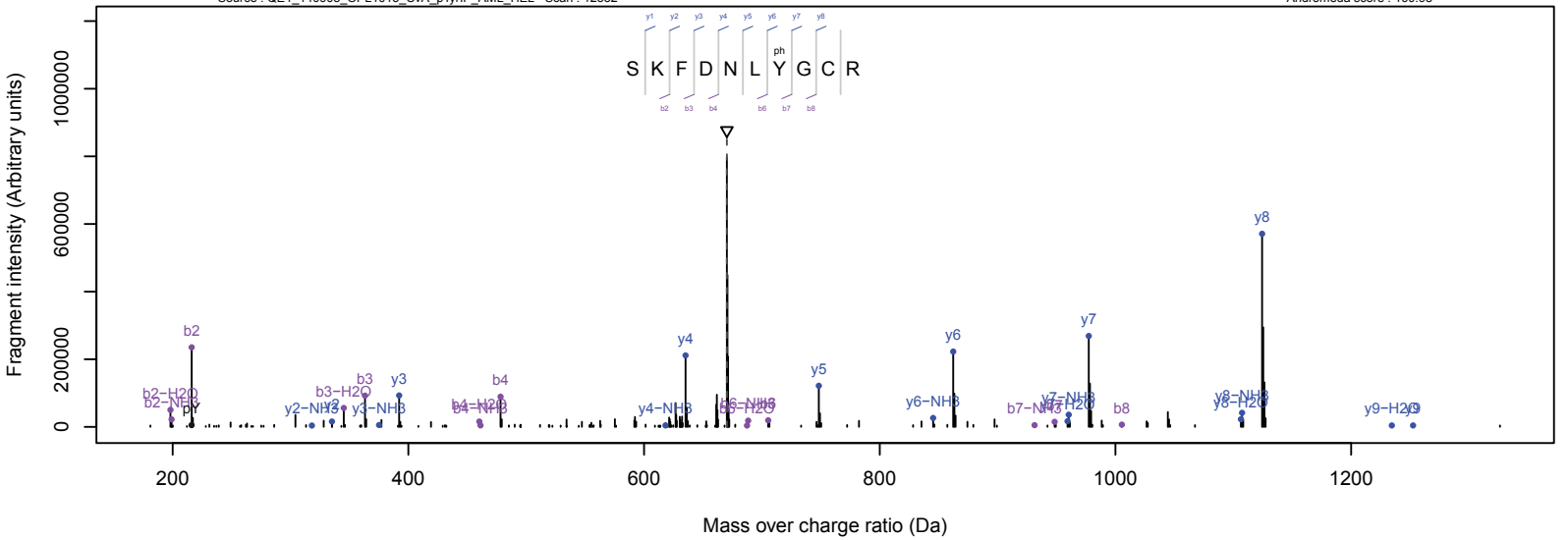
Andromeda score : 161.04



SKFDNLY(ph)GCR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 12382

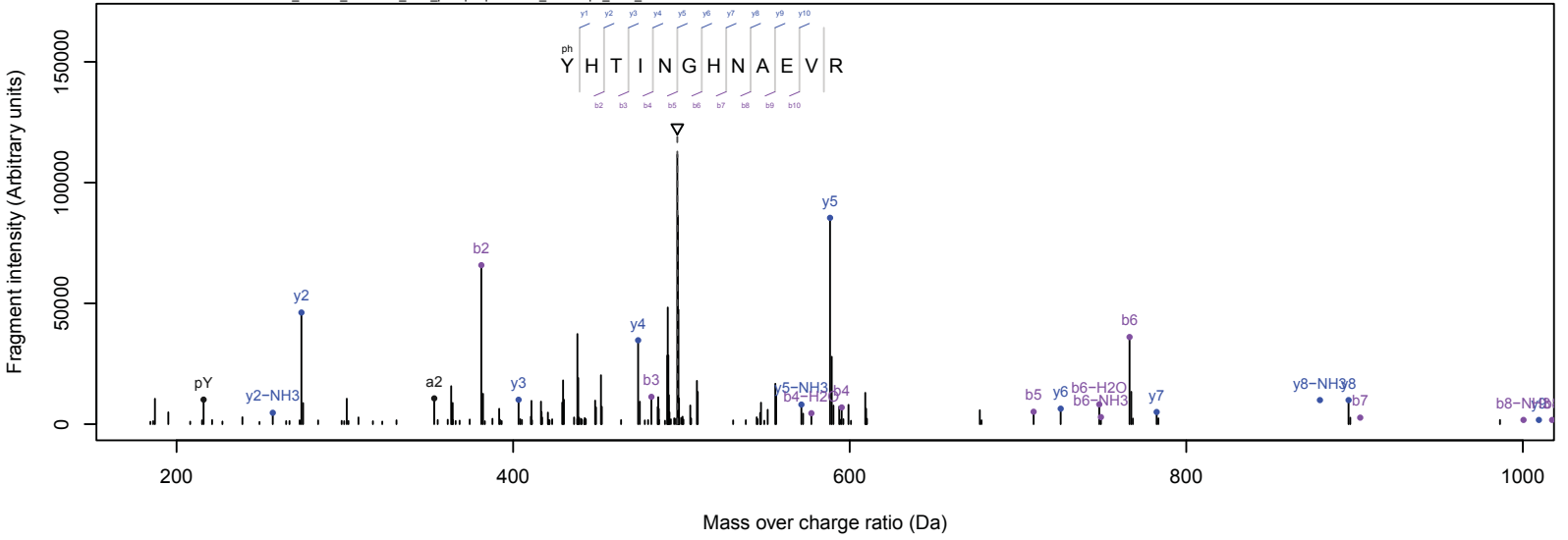
Andromeda score : 160.98



Y(ph)HTINGHNAEVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 4654

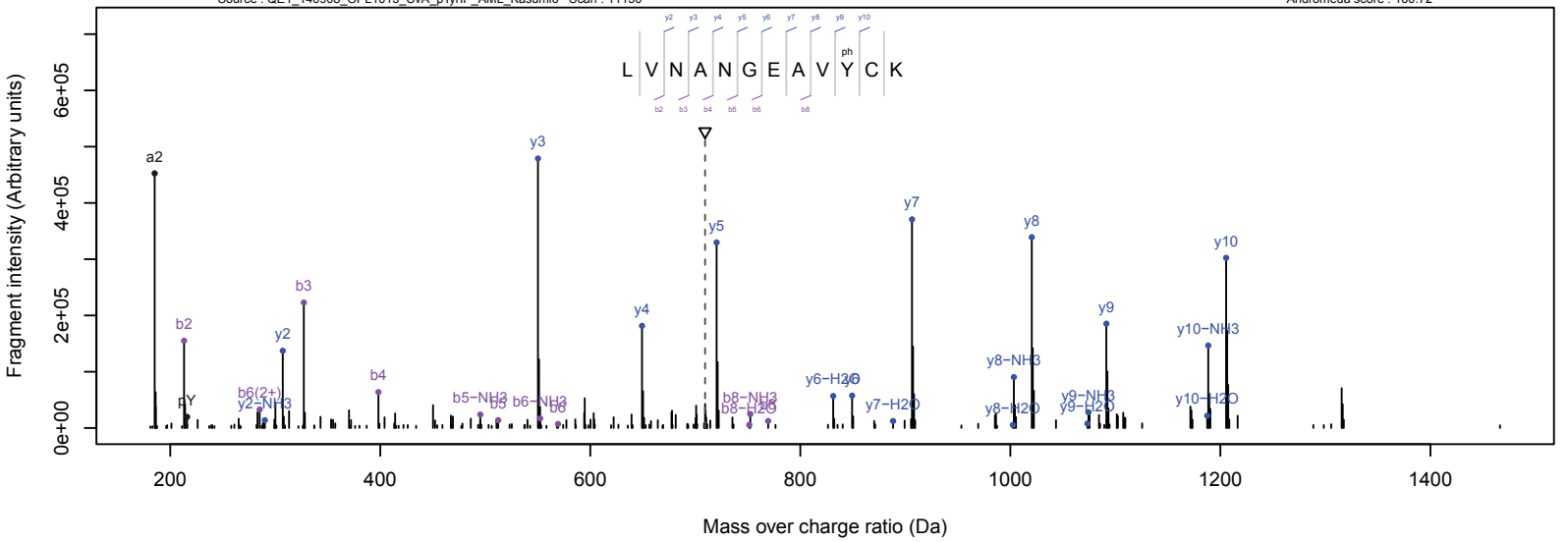
Andromeda score: 160.84



LVNANGEAVY(ph)CK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 11130

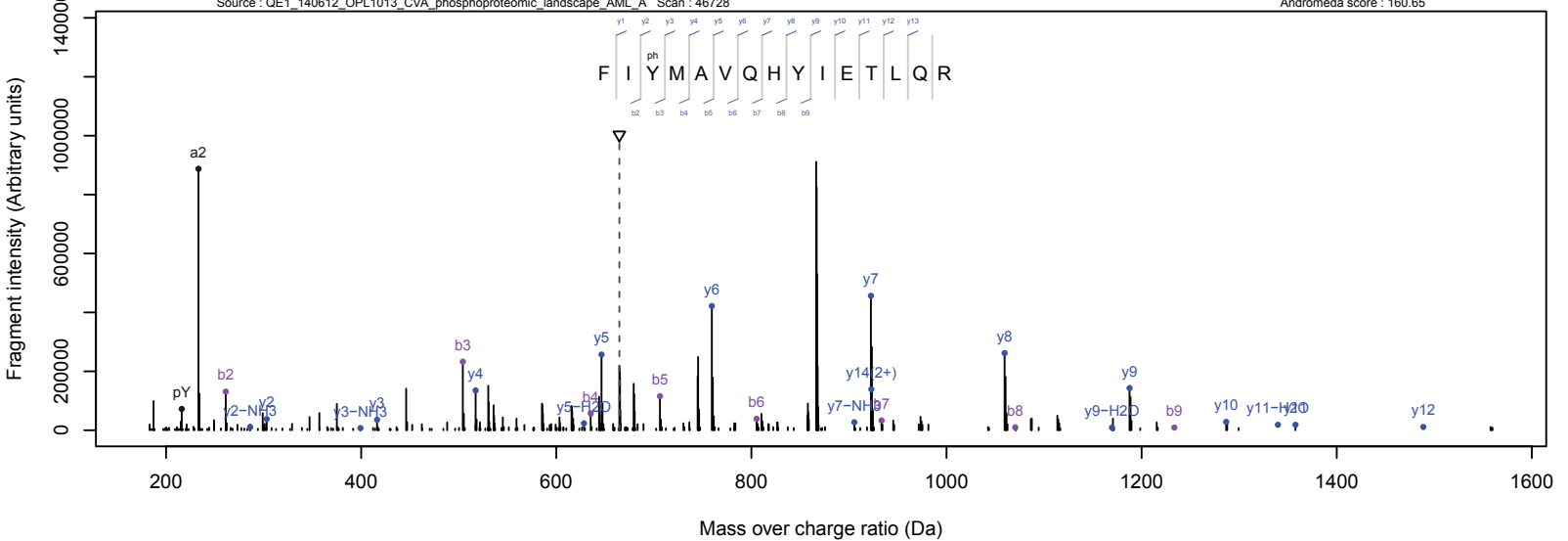
Andromeda score: 160.72



FIY(ph)MAVQHYIETLQR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 46728

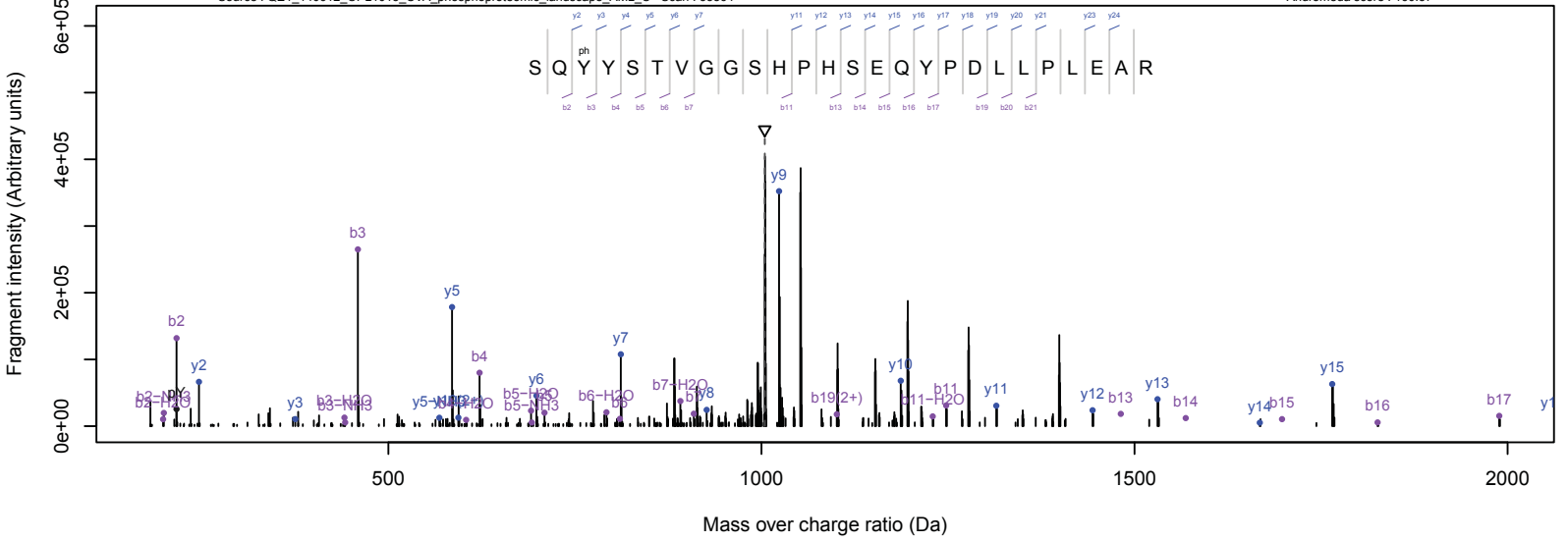
Andromeda score: 160.65



**SQY(ph)YSTVGGSHPHSEQYPDLLPLEAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33864

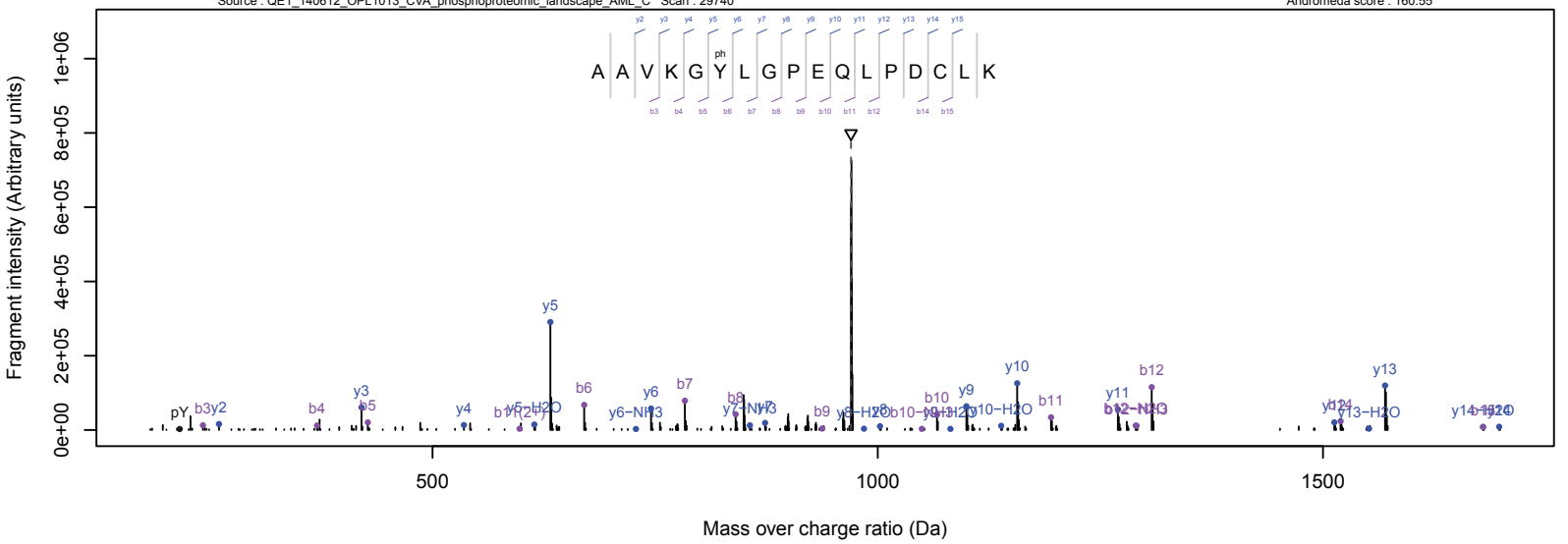
Andromeda score : 160.57



**AAVKY(ph)LGPEQLPDCLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29740

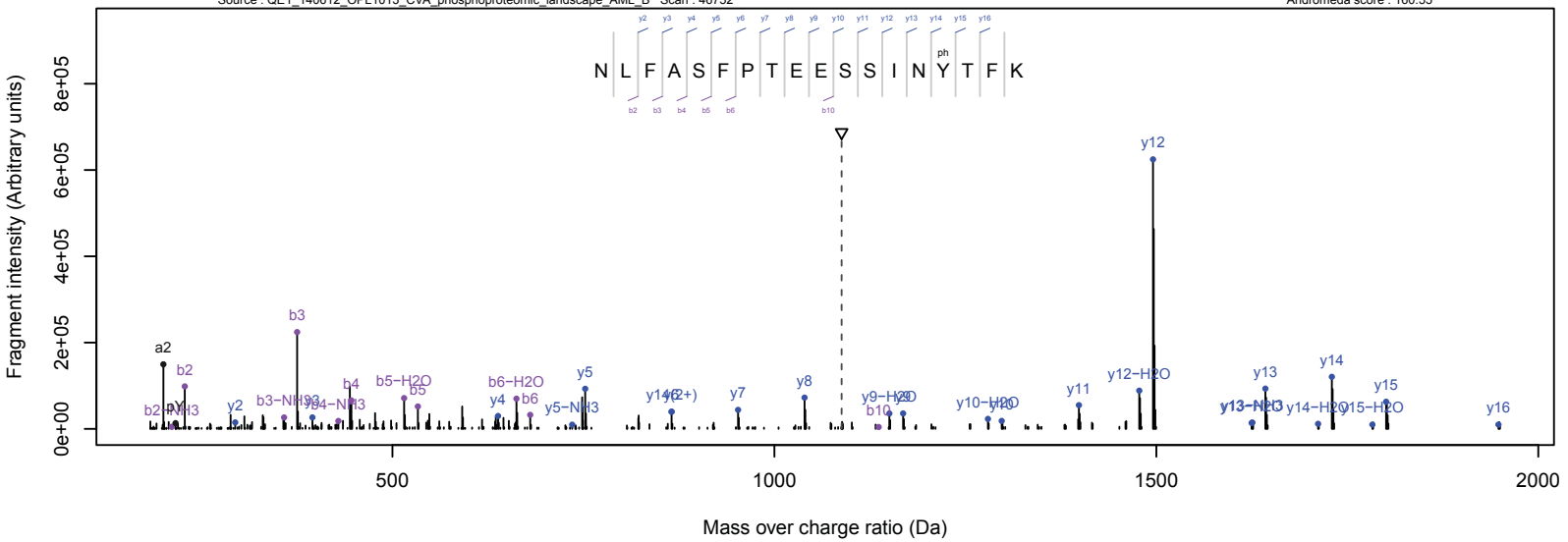
Andromeda score : 160.55



**NLFASFPTTEESSIN(ph)YTFK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46752

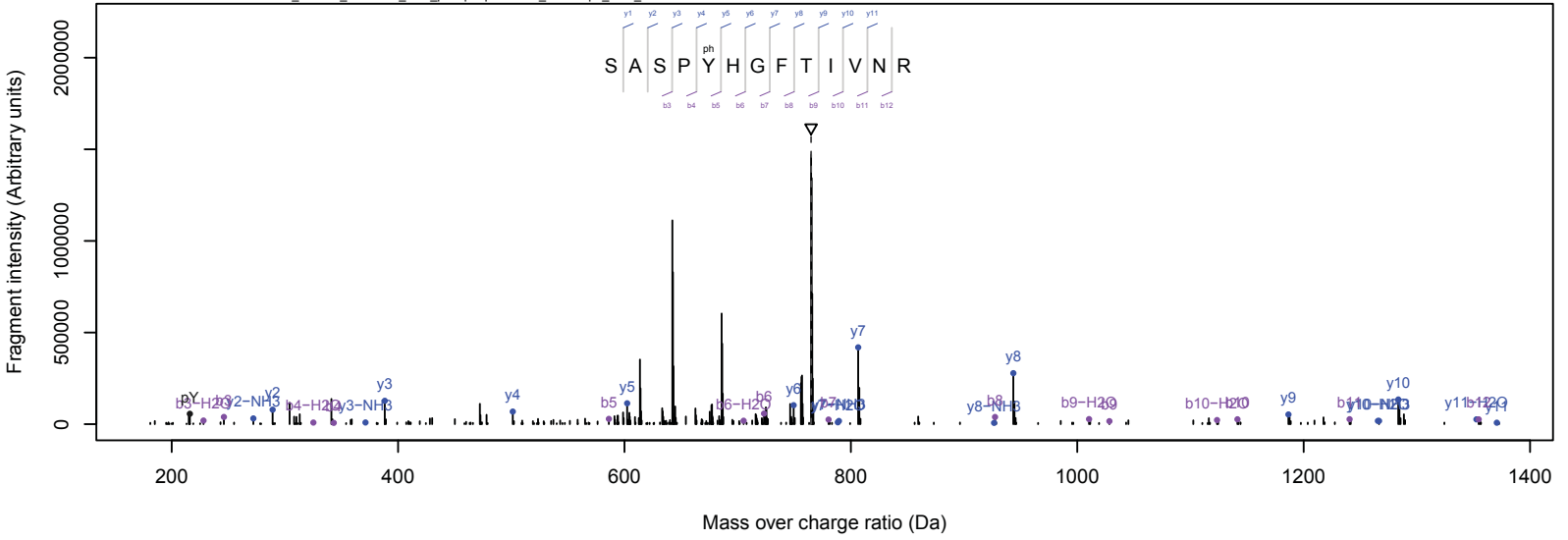
Andromeda score : 160.55



SASPY(ph)HGFTIVNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19758

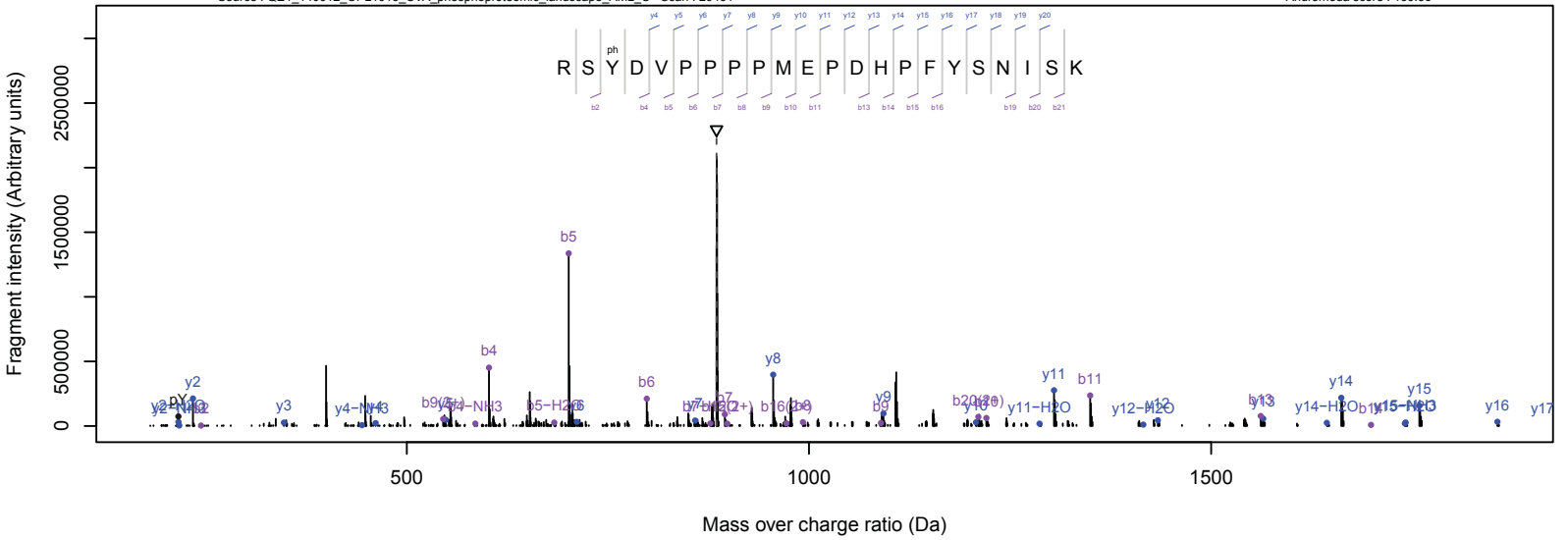
Andromeda score : 160.48



RSY(ph)DVPPPPMEPDHPFYSNISK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29431

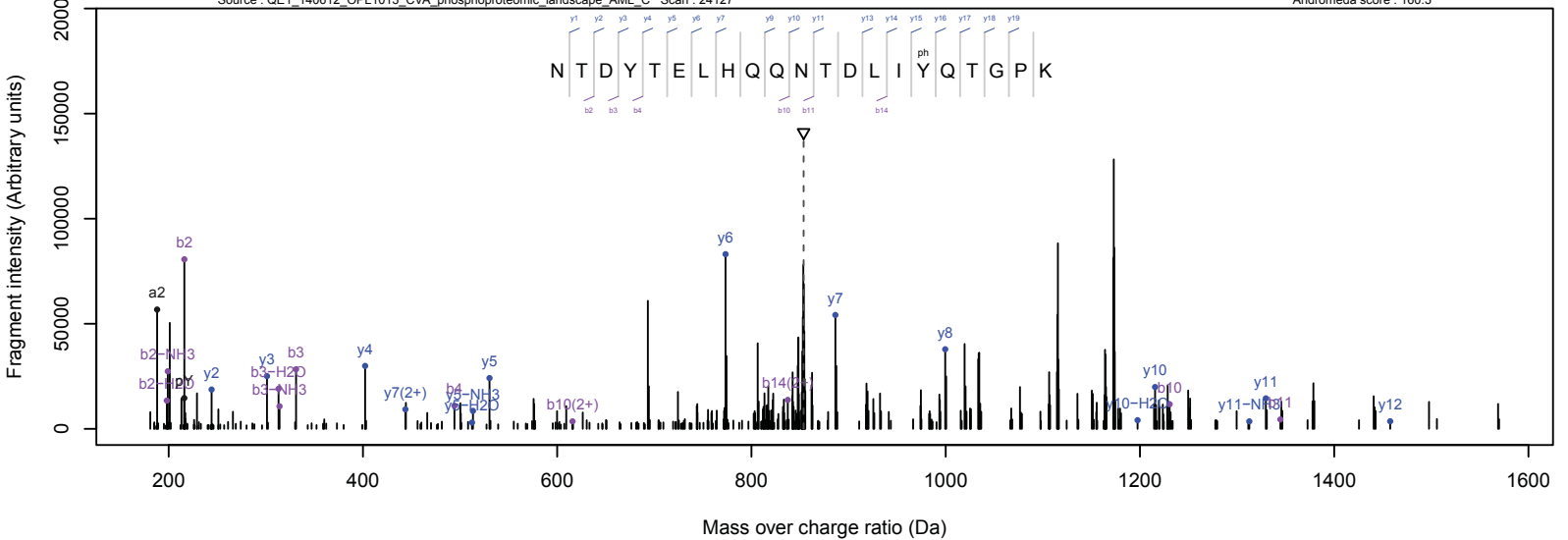
Andromeda score : 160.36



NTDYTELHQNTDLIY(ph)QTGPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24127

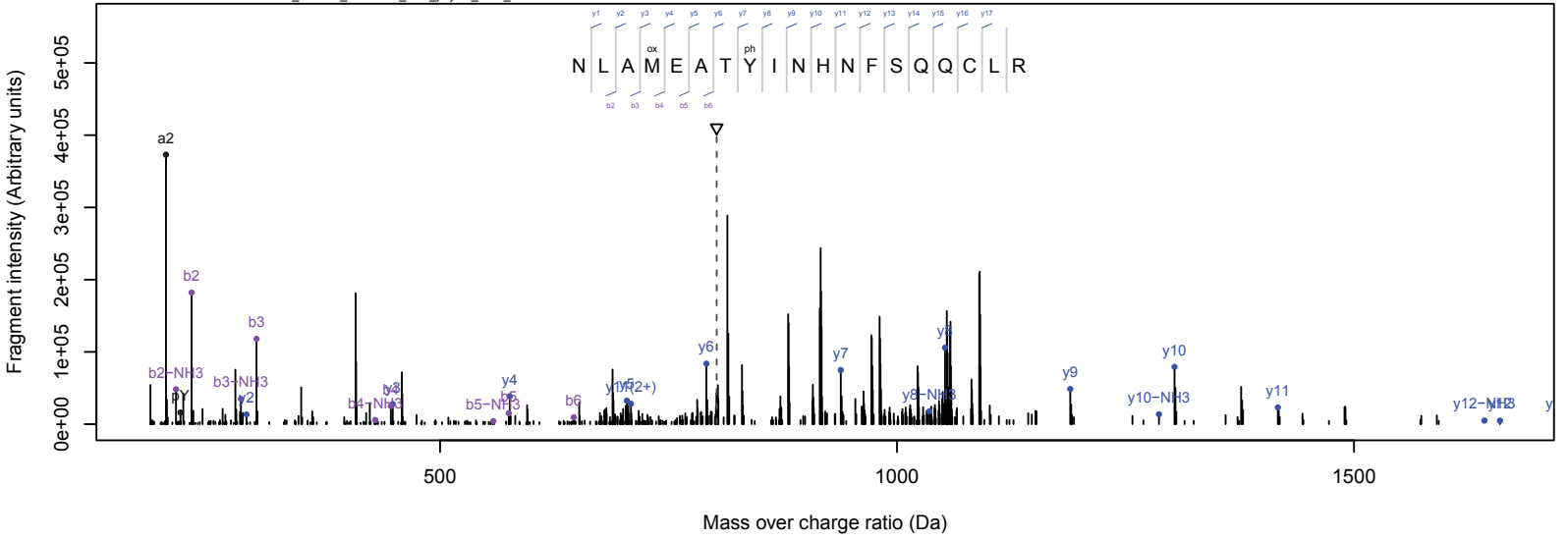
Andromeda score : 160.3



NLAM(ox)EATY(ph)INHNSQCLR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 33991

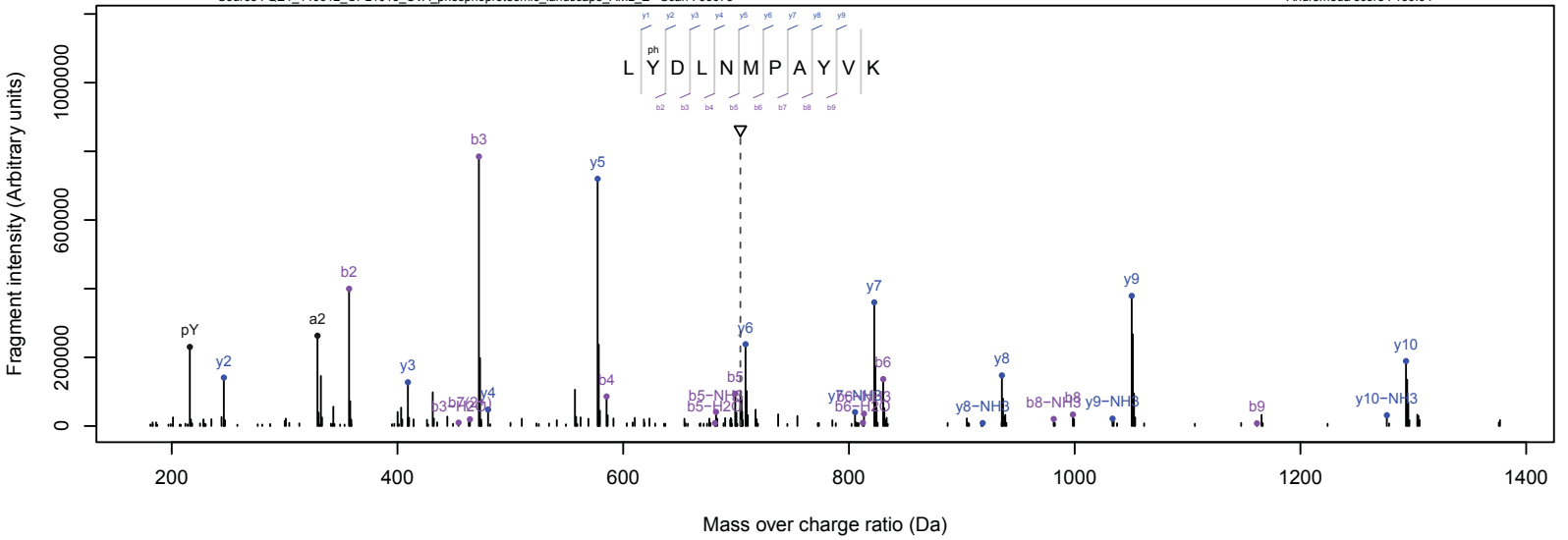
Andromeda score : 159.95



LY(ph)DLNMPAYVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 33073

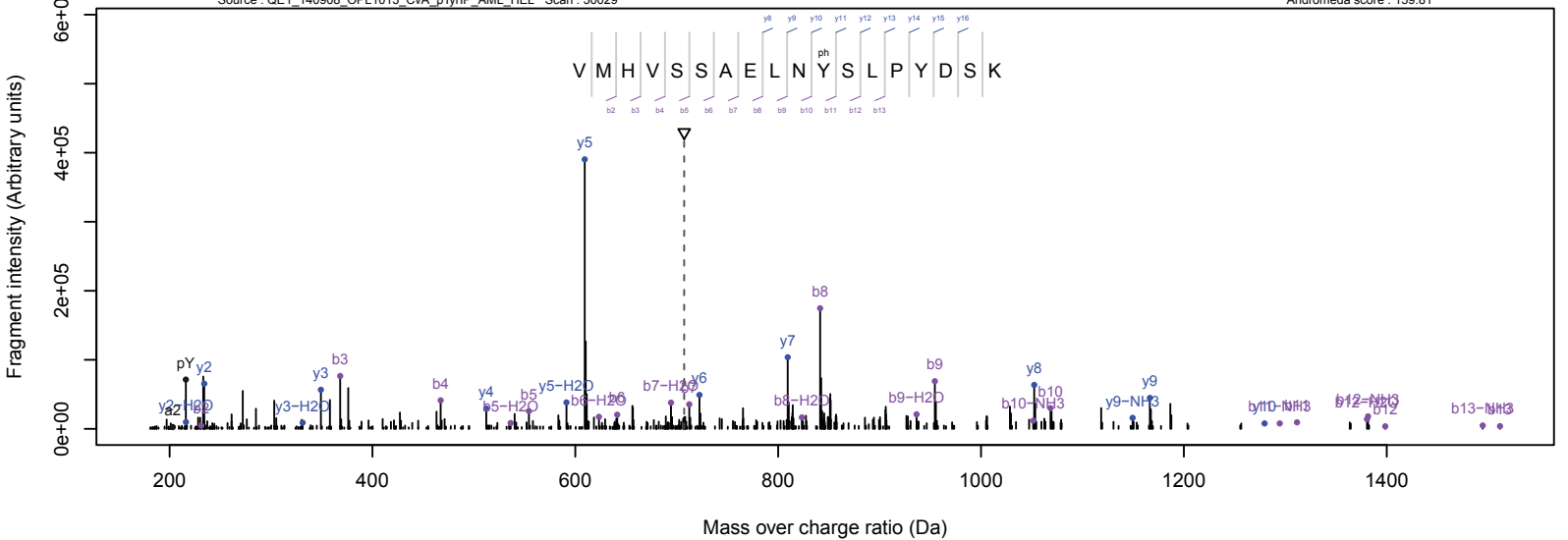
Andromeda score : 159.94



VMHVSSAELNYSLPYDSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 30029

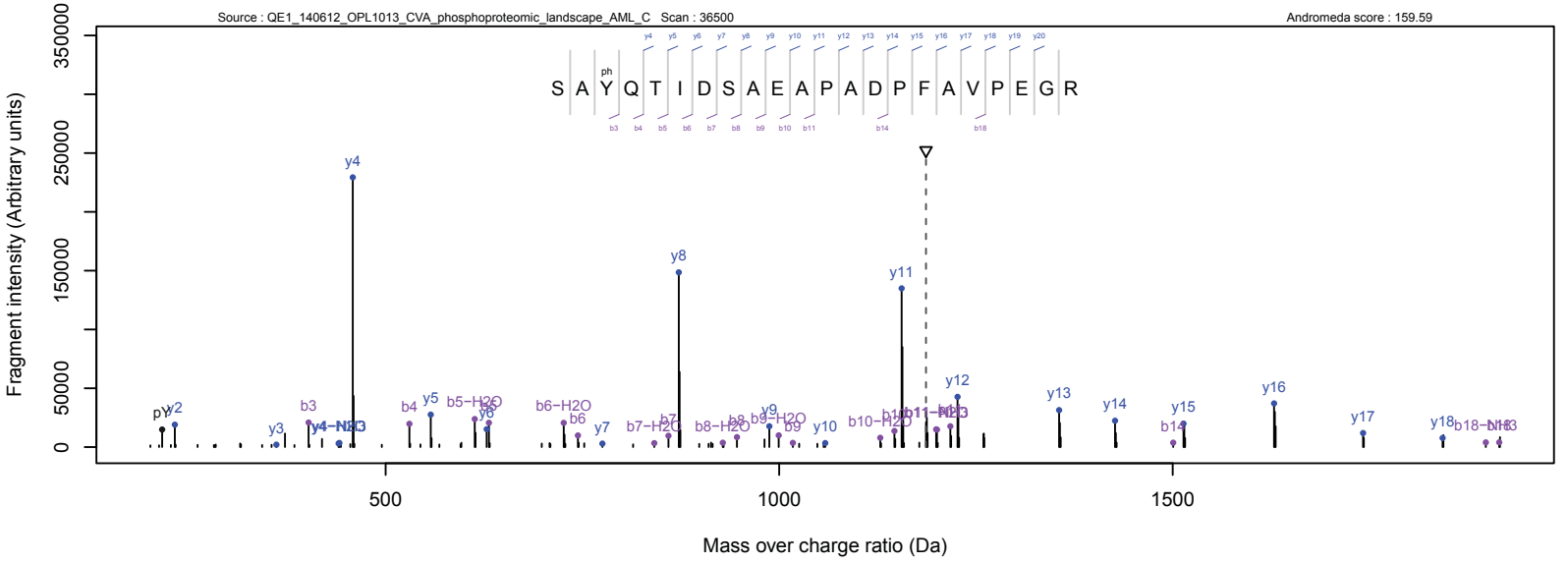
Andromeda score : 159.81



SAY(ph)QTIDSAEAPADPFVPEGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36500

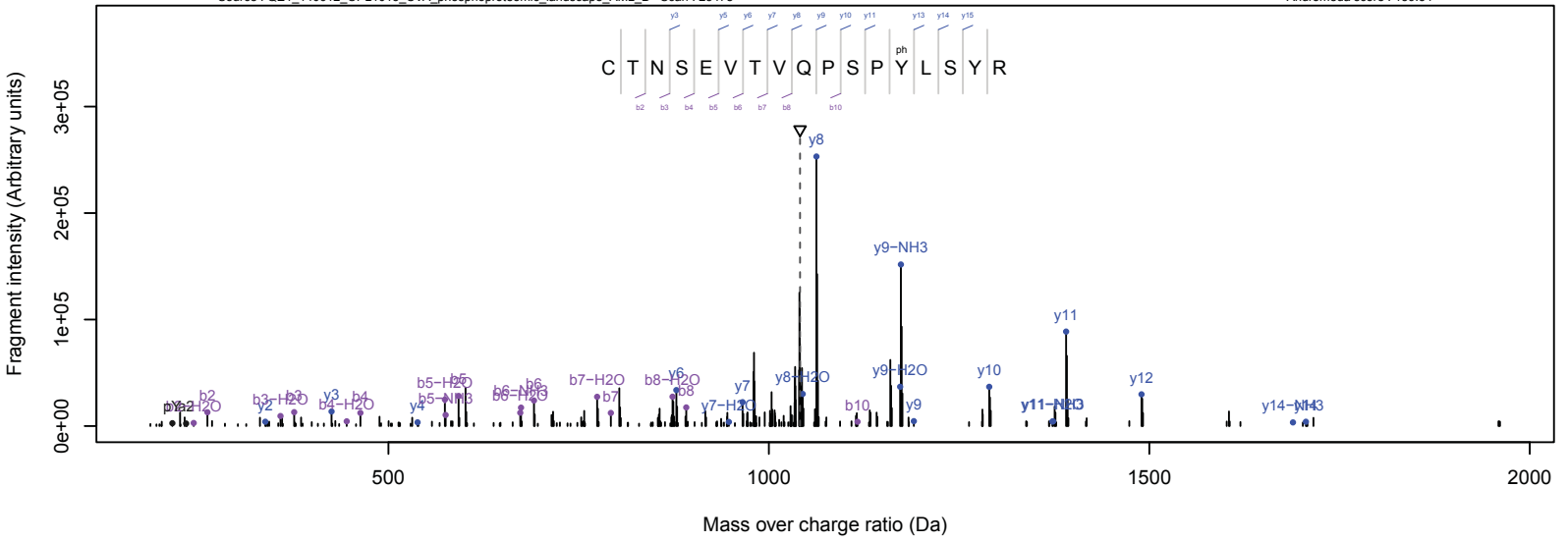
Andromeda score : 159.59



CTNSEVTVQSPY(ph)LSYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25478

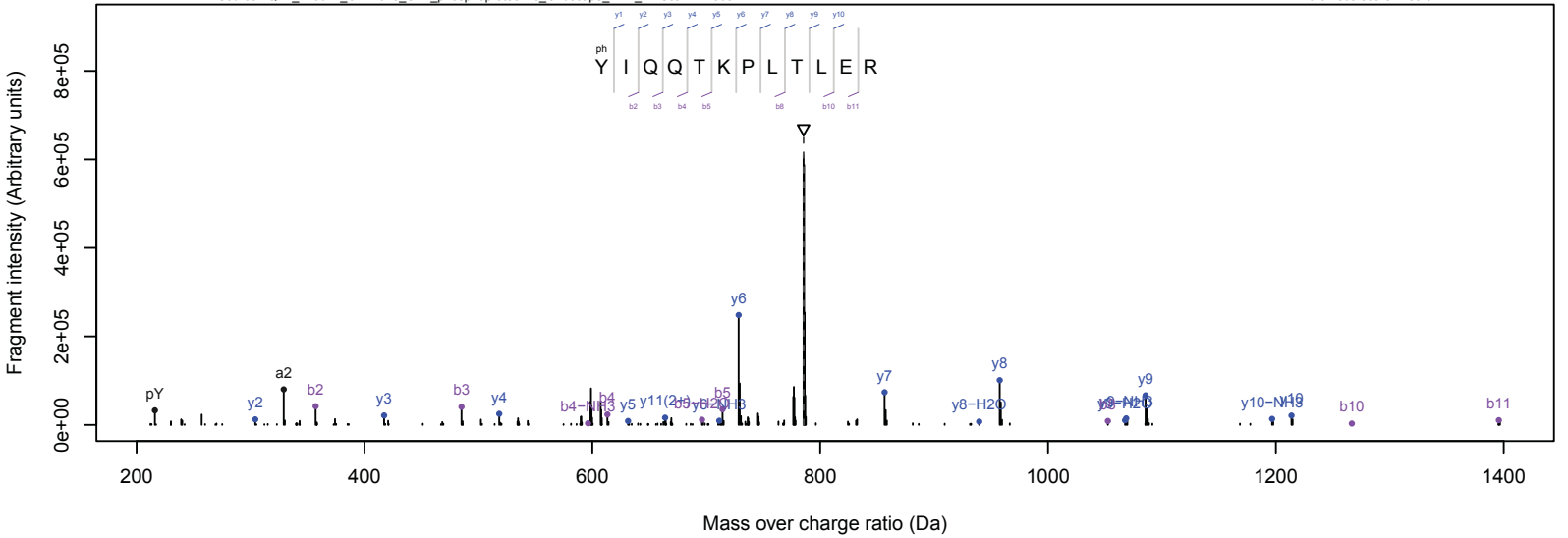
Andromeda score : 159.54



Y(ph)IQQTKPLTLER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17365

Andromeda score : 159.5



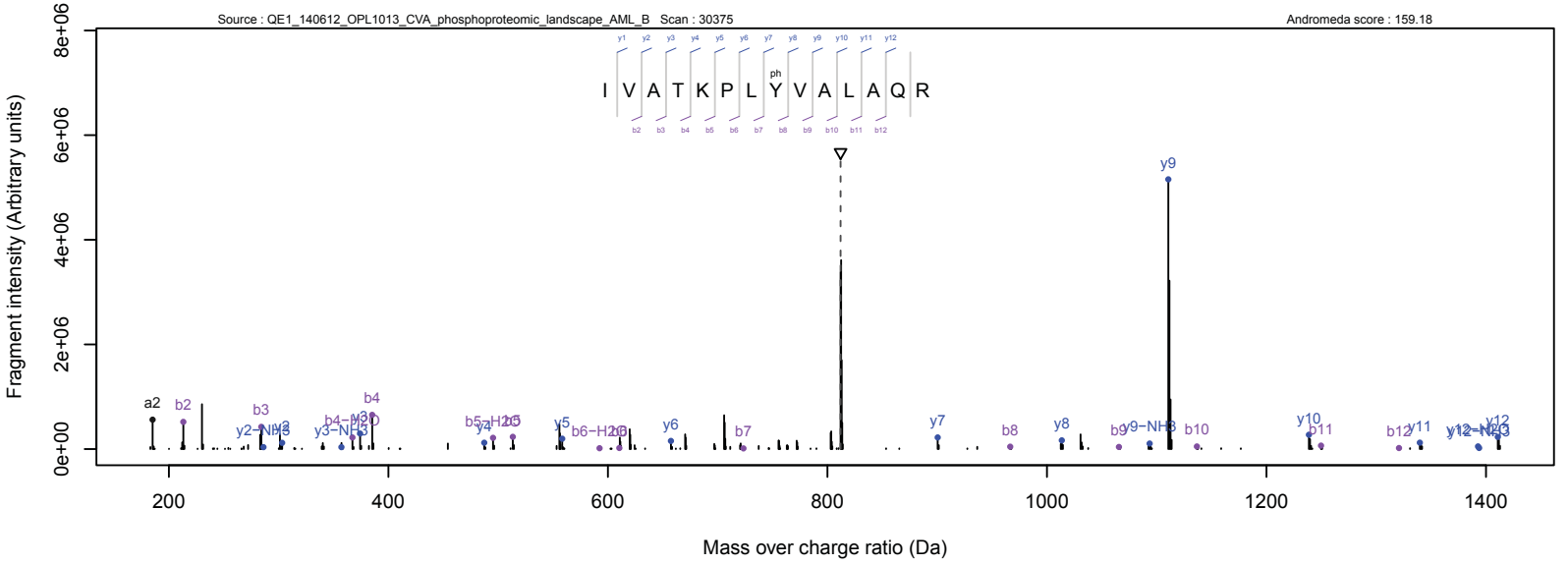




IVATKPLY(ph)VALAQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30375

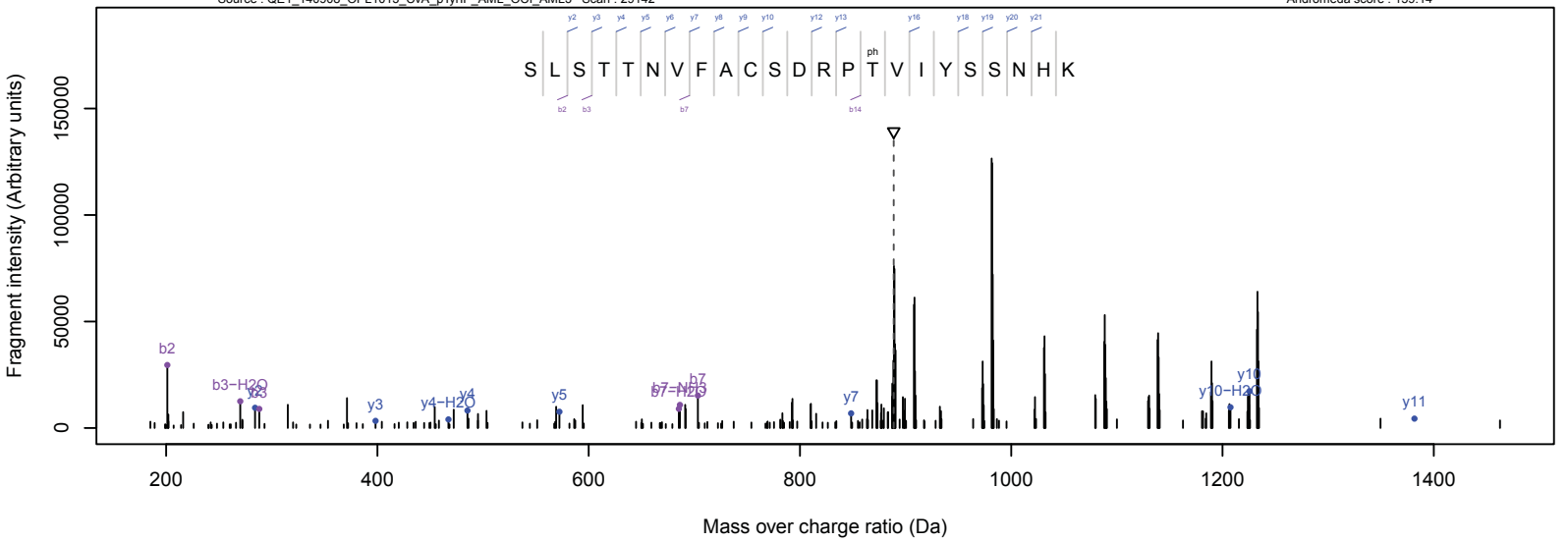
Andromeda score : 159.18



SLSTTNVFACSDRPT(ph)VIYSSNHK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 29142

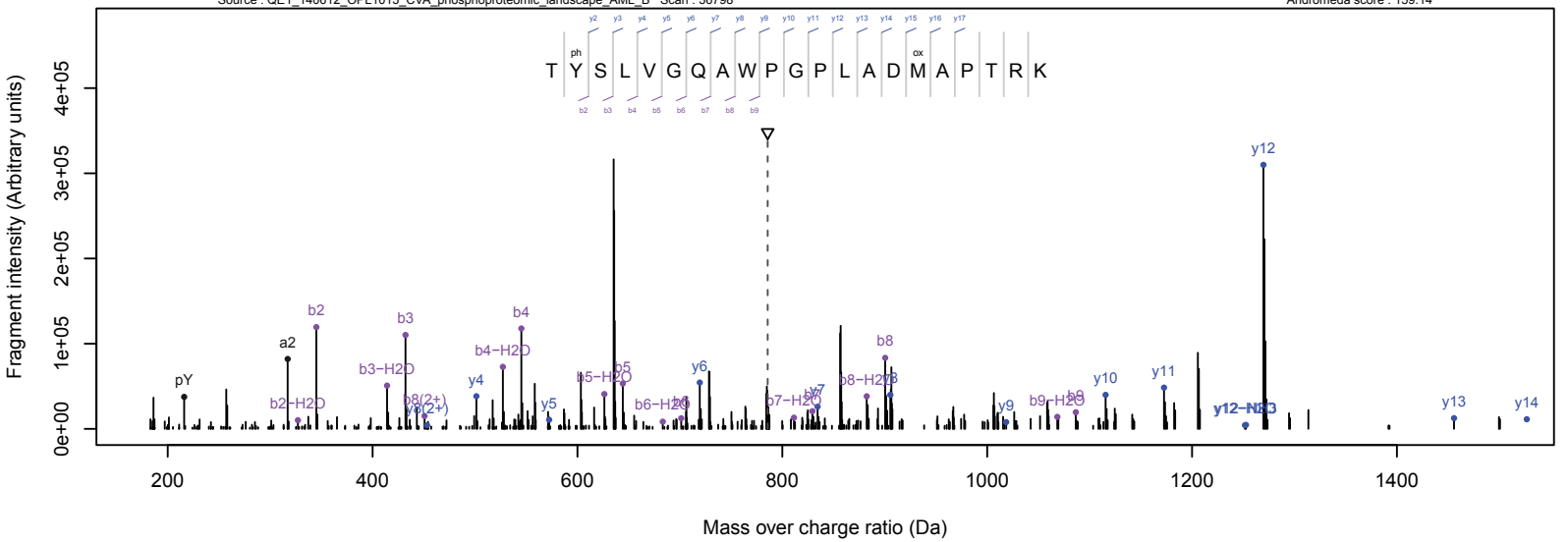
Andromeda score : 159.14



TY(ph)SLVGQAWPGLADM(ox)APTRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36798

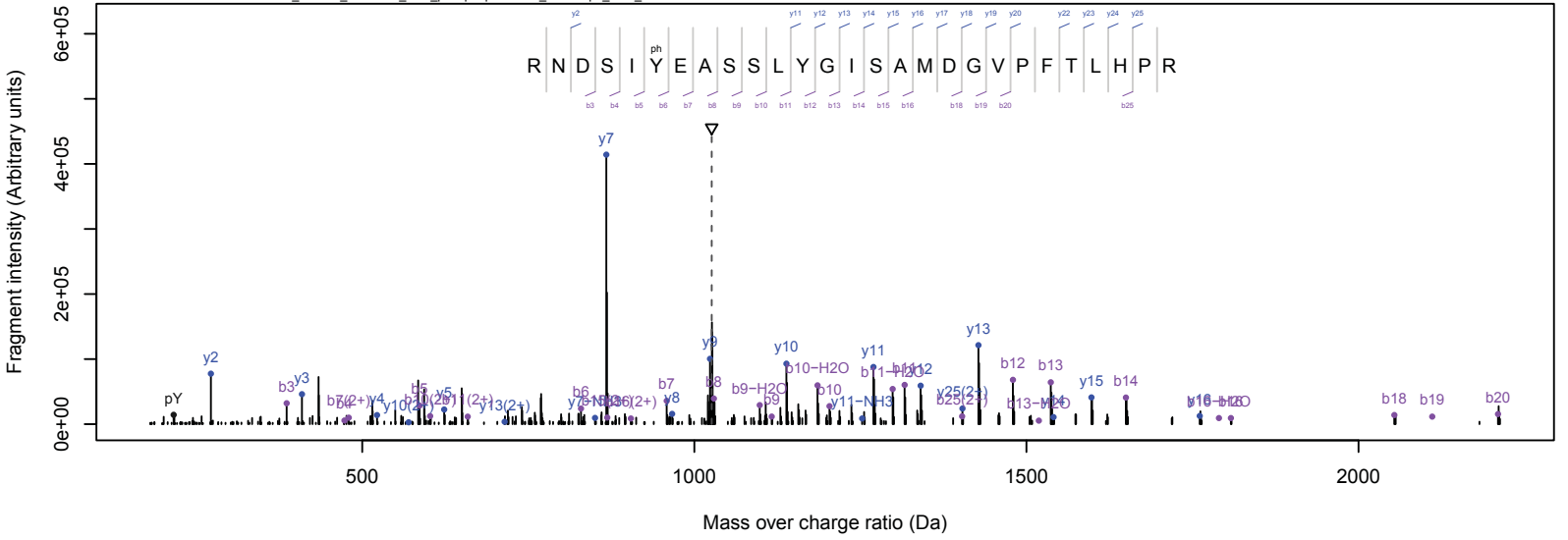
Andromeda score : 159.14



RNDSIY(ph)EASSLYGISAMDGVPTFLHPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47071

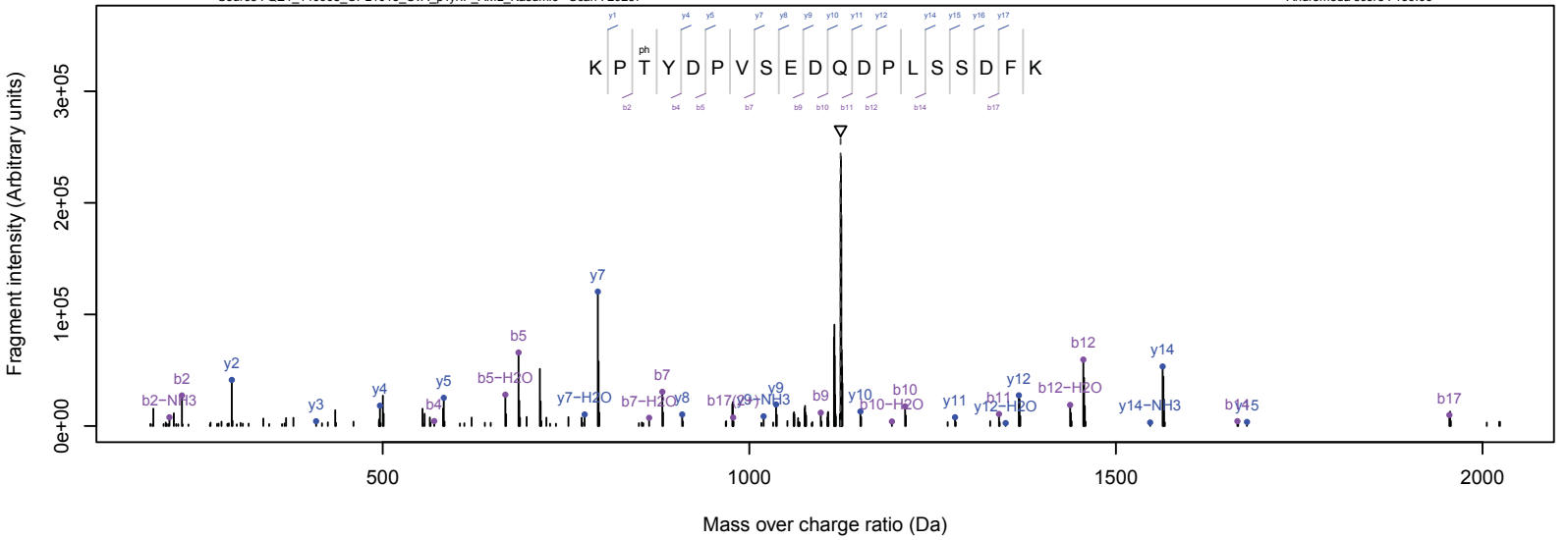
Andromeda score : 159.13



KPT(ph)YDPVSEDDQLSPSSDFK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 29257

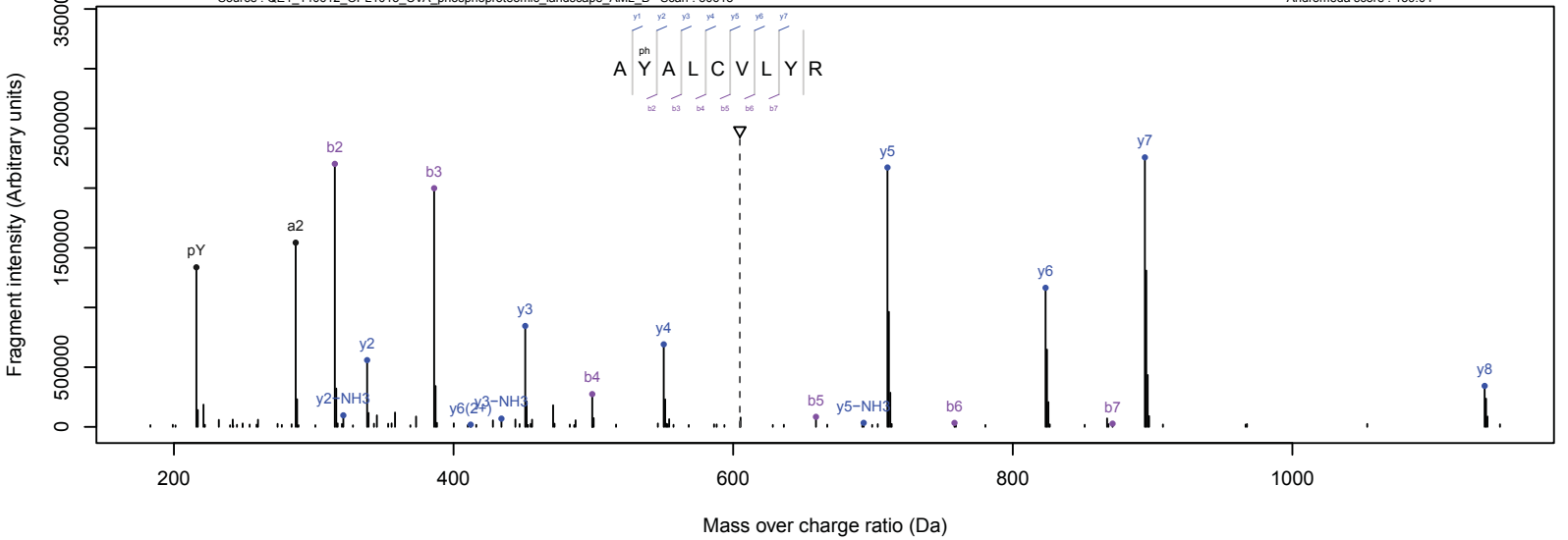
Andromeda score : 159.06



AY(ph)ALCVLYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30618

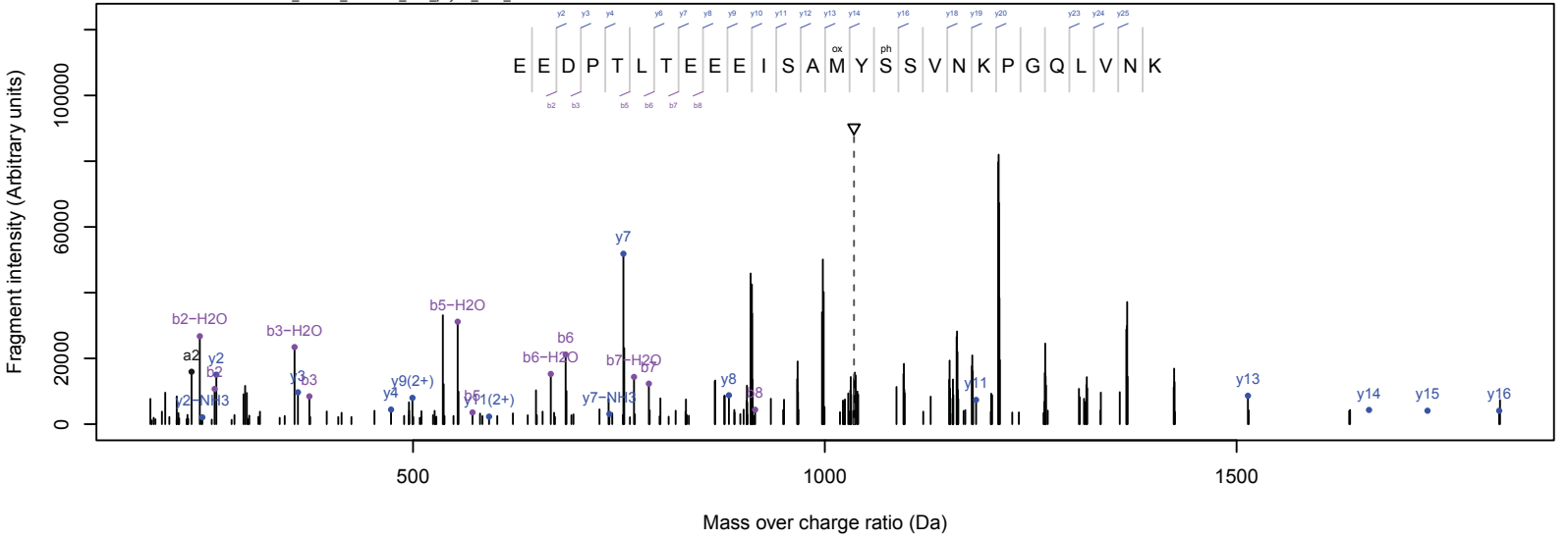
Andromeda score : 159.04



EEDPTLLEEISAM(ox)YS(ph)SVNKPQLVNK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_ML2 Scan : 30781

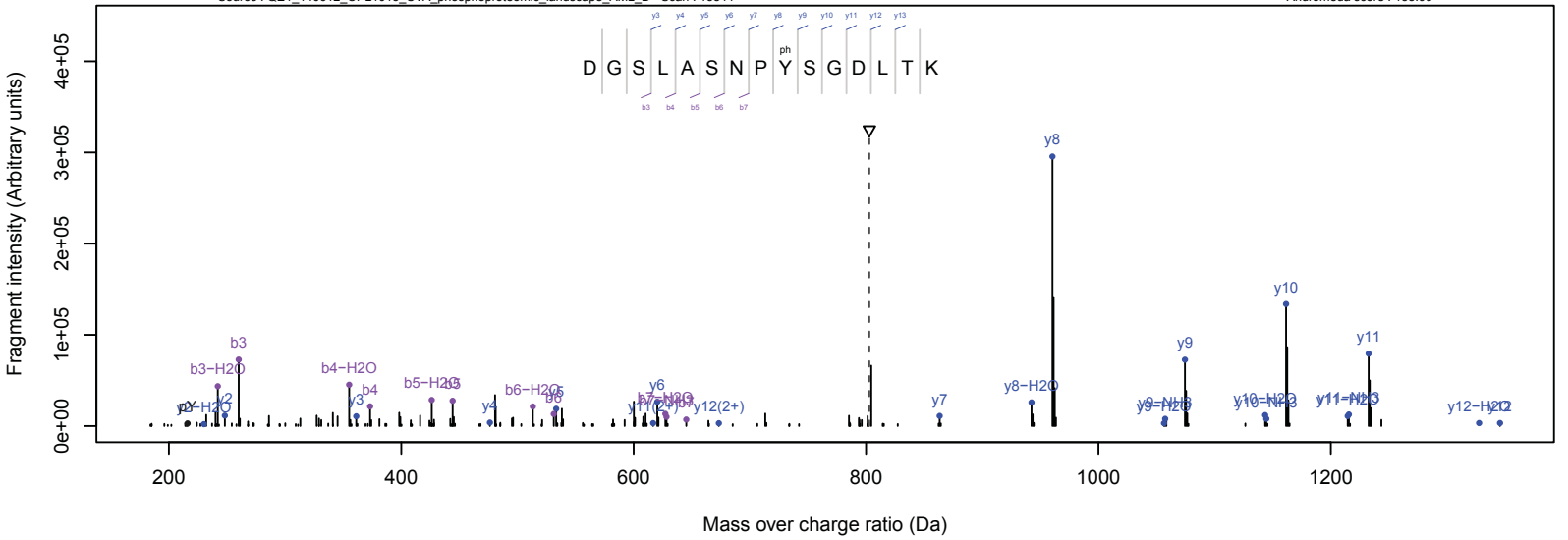
Andromeda score : 158.94



DGSLASNPY(ph)SGDLTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 15844

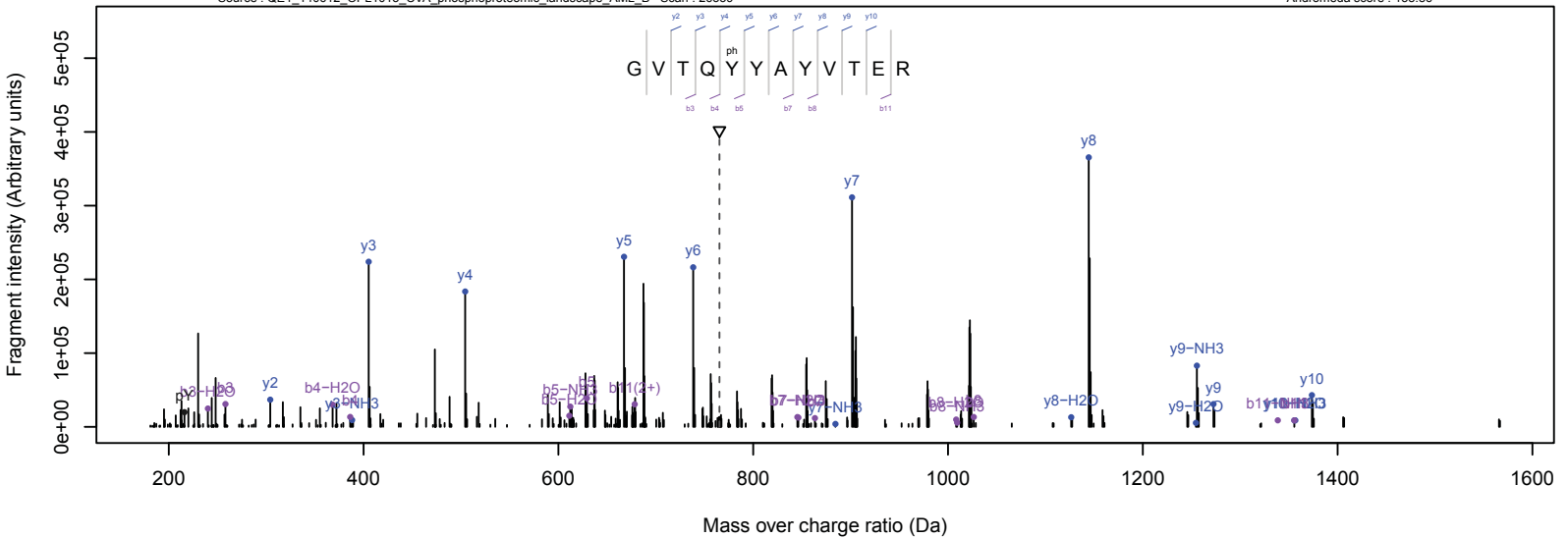
Andromeda score : 158.93



GVTQY(ph)YAYVTER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26339

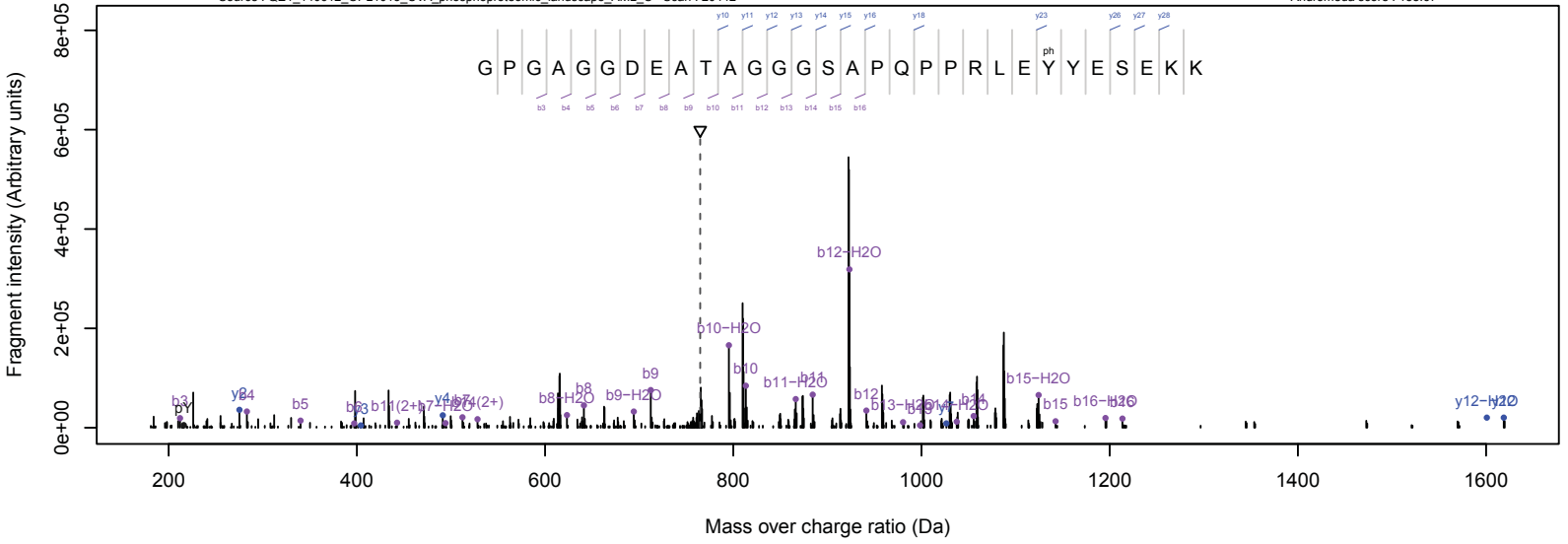
Andromeda score : 158.86



GPGAGGDEATAGGGSAPQPPRLEYESEKK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 20442

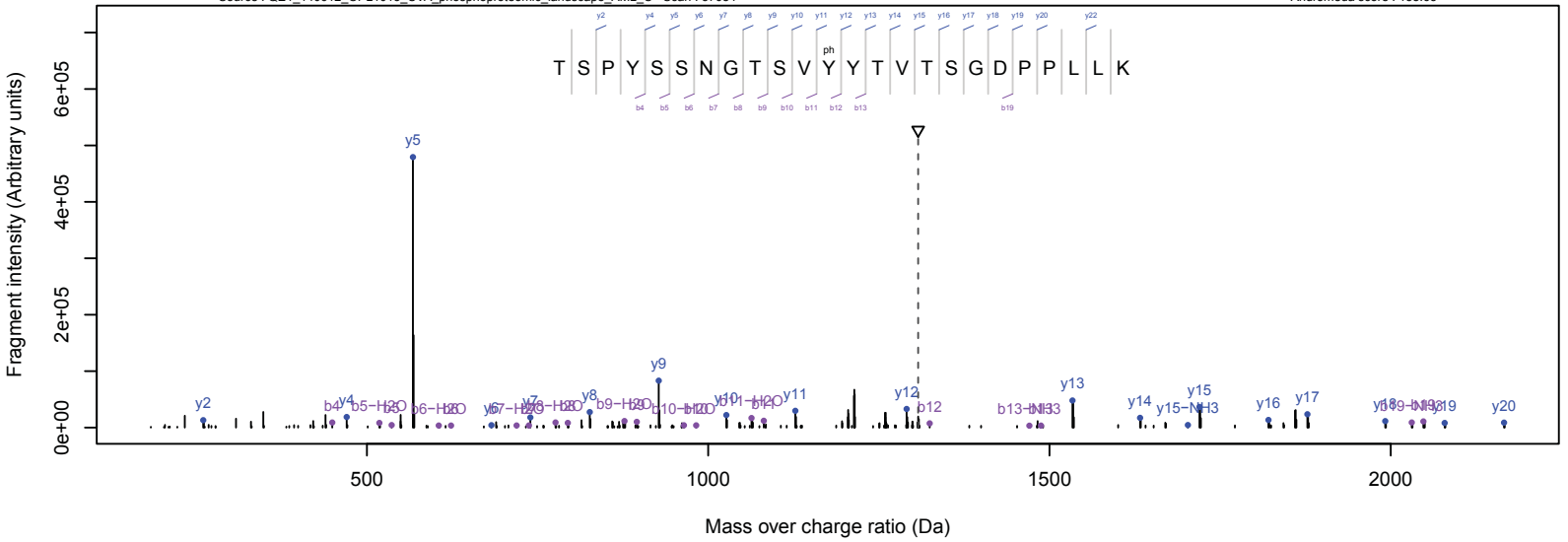
Andromeda score: 158.67



TSPYSSNGTSVY(Y)YTVTSGDPPLLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37084

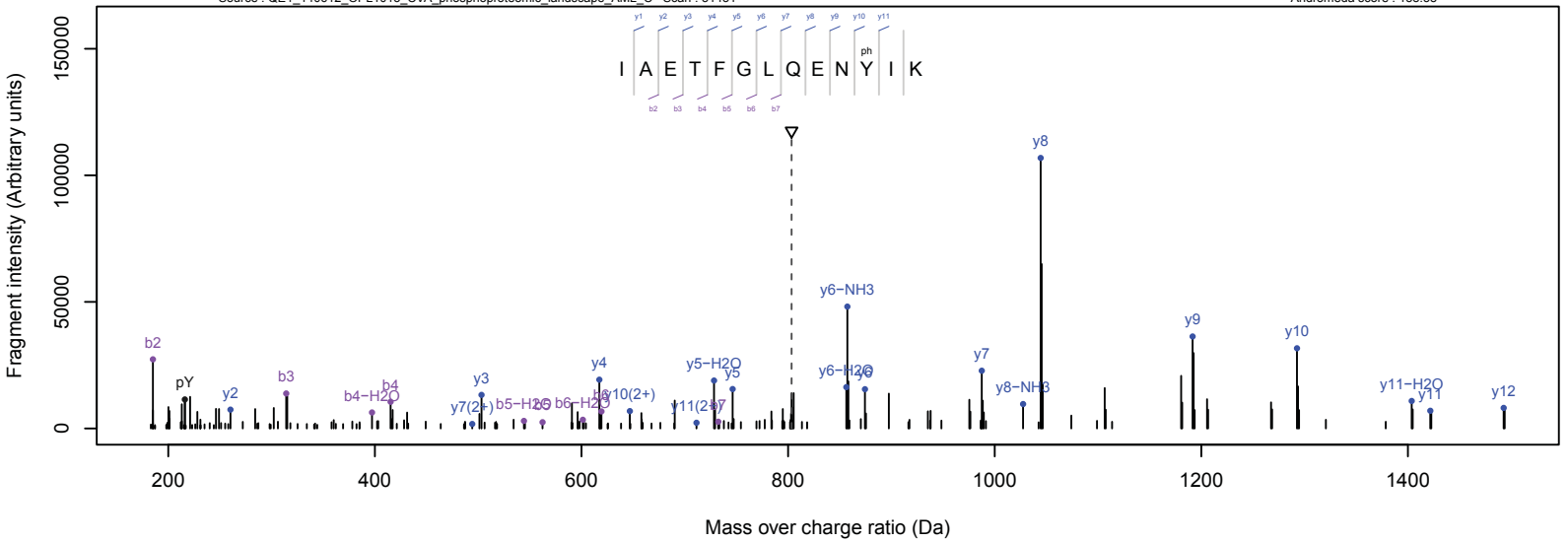
Andromeda score: 158.65



IAETFGLQENY(I)K-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31431

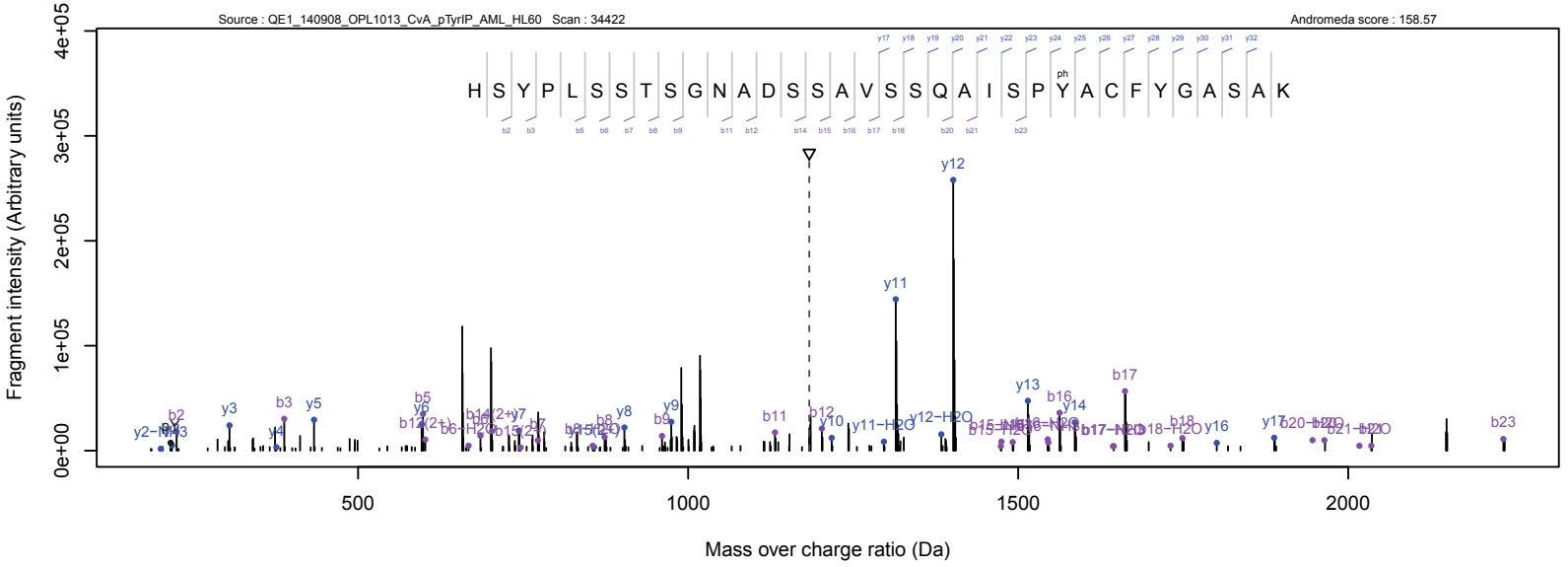
Andromeda score: 158.58



HSYPLSSTSGNADSSAVSSQAISPY(ph)ACFYGASAK-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan: 34422

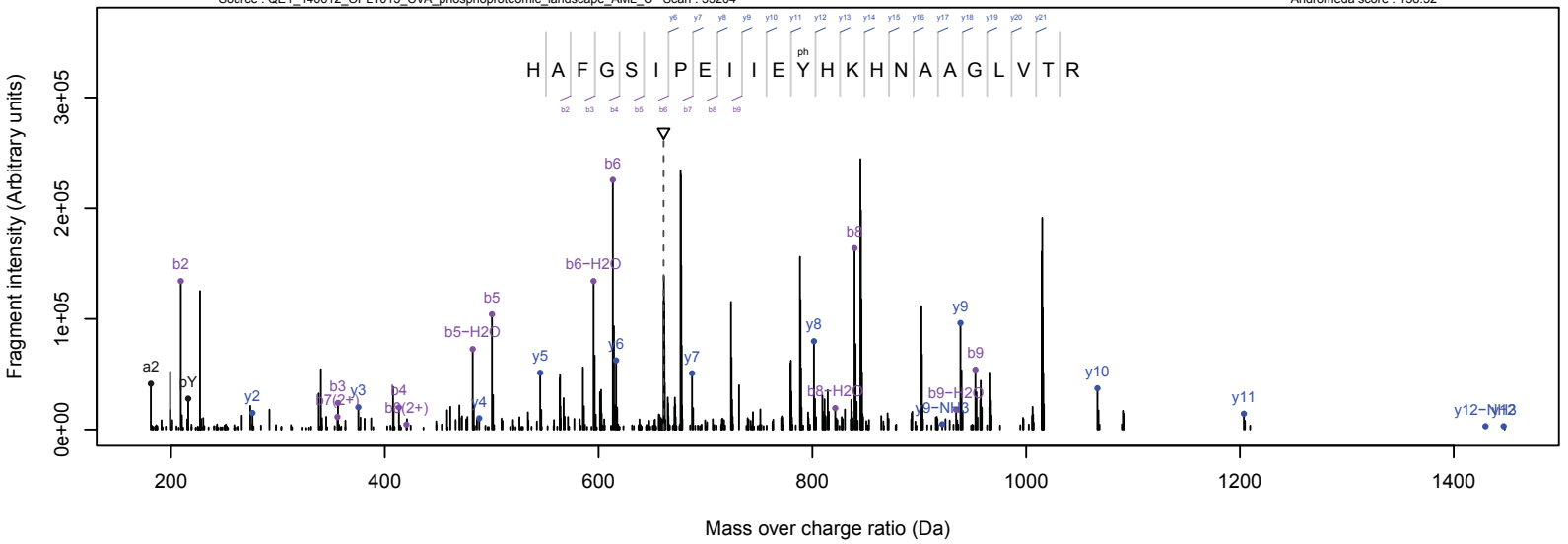
Andromeda score: 158.57



HAFGSIPEIIEY(ph)HKHNAAGLVTR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 33204

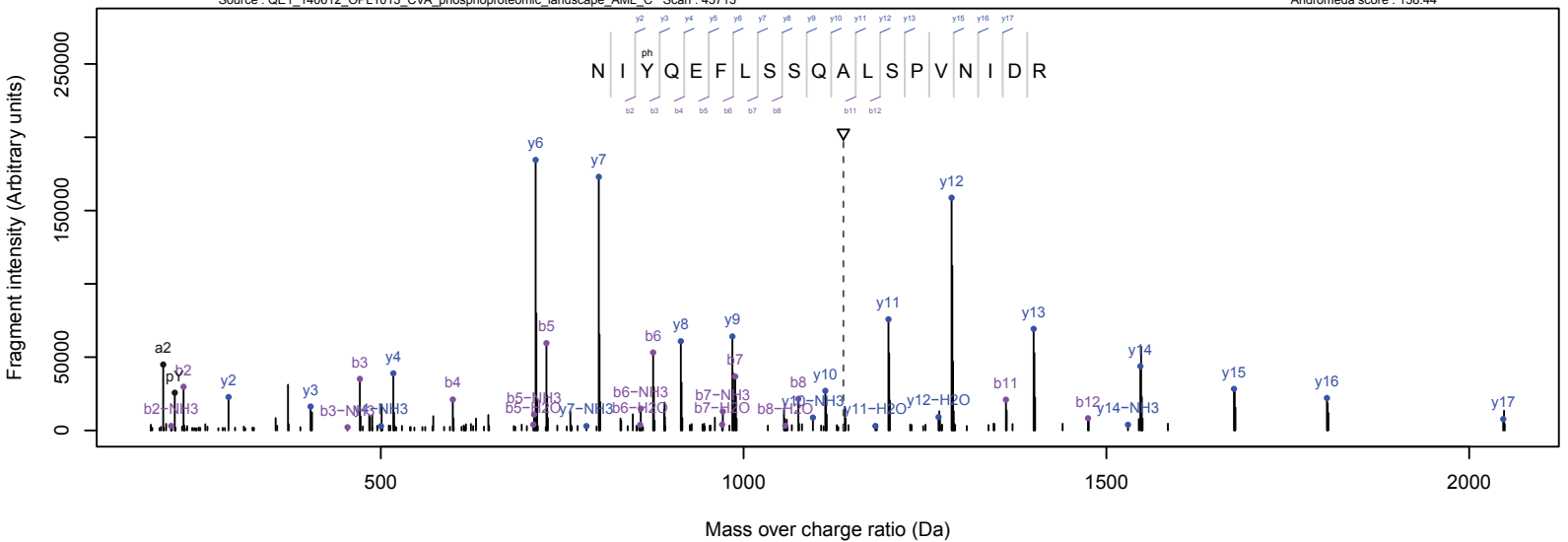
Andromeda score: 158.52



NIY(ph)QEFLSSQALSPVNIDR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 45713

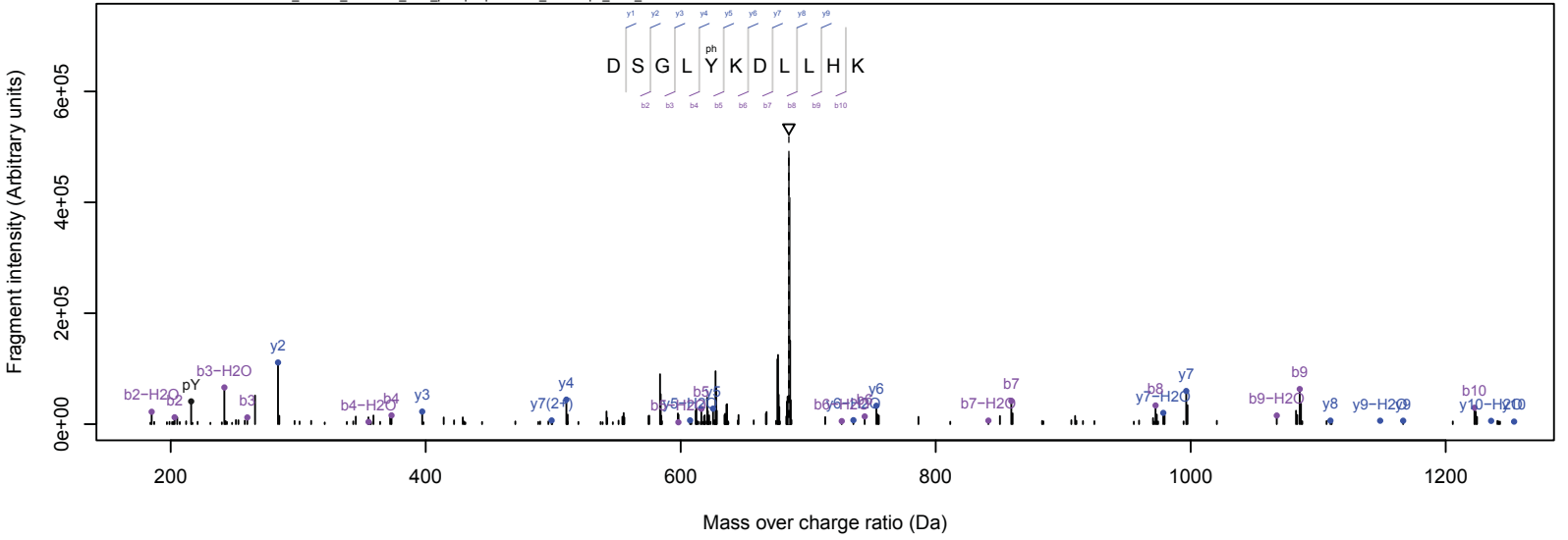
Andromeda score: 158.44



DSGLY(ph)KDLLHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19289

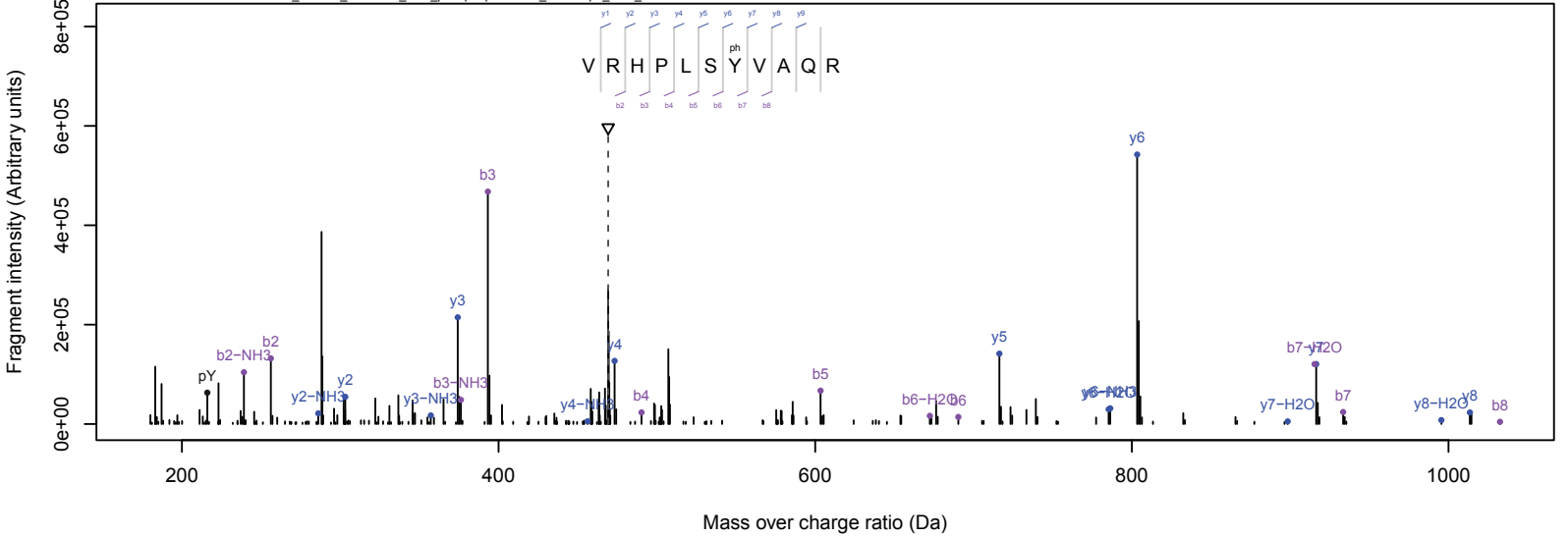
Andromeda score : 158.34



VRHPLSY(ph)VAQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12267

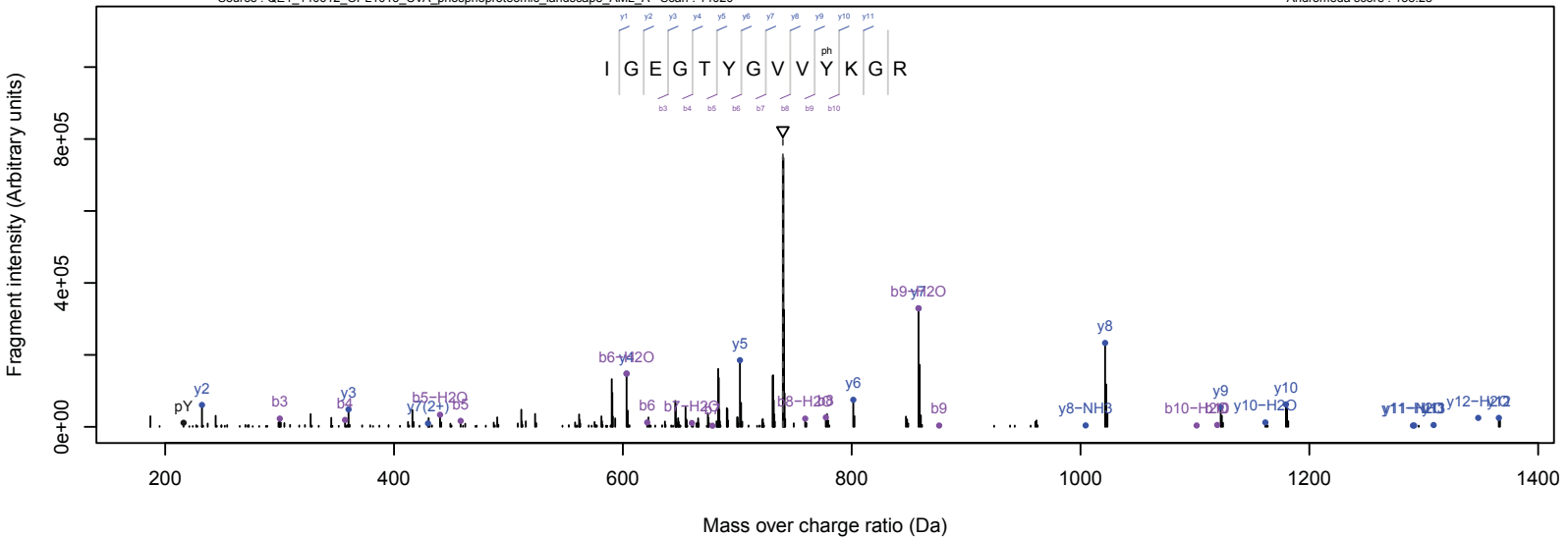
Andromeda score : 158.3



IGEGTYGVVYKGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 14020

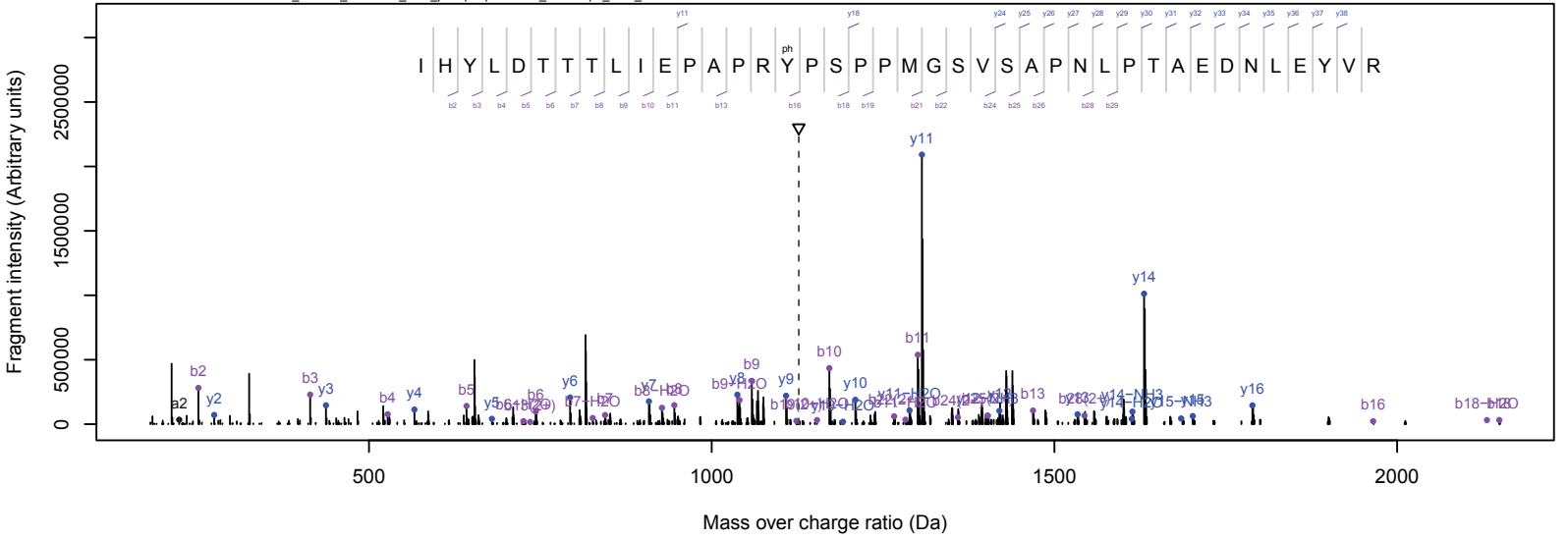
Andromeda score : 158.25



IHYLDTTTLIEPAPRY(ph)PSPPMGVSAPSAPNLPTAEDNLEYVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 45827

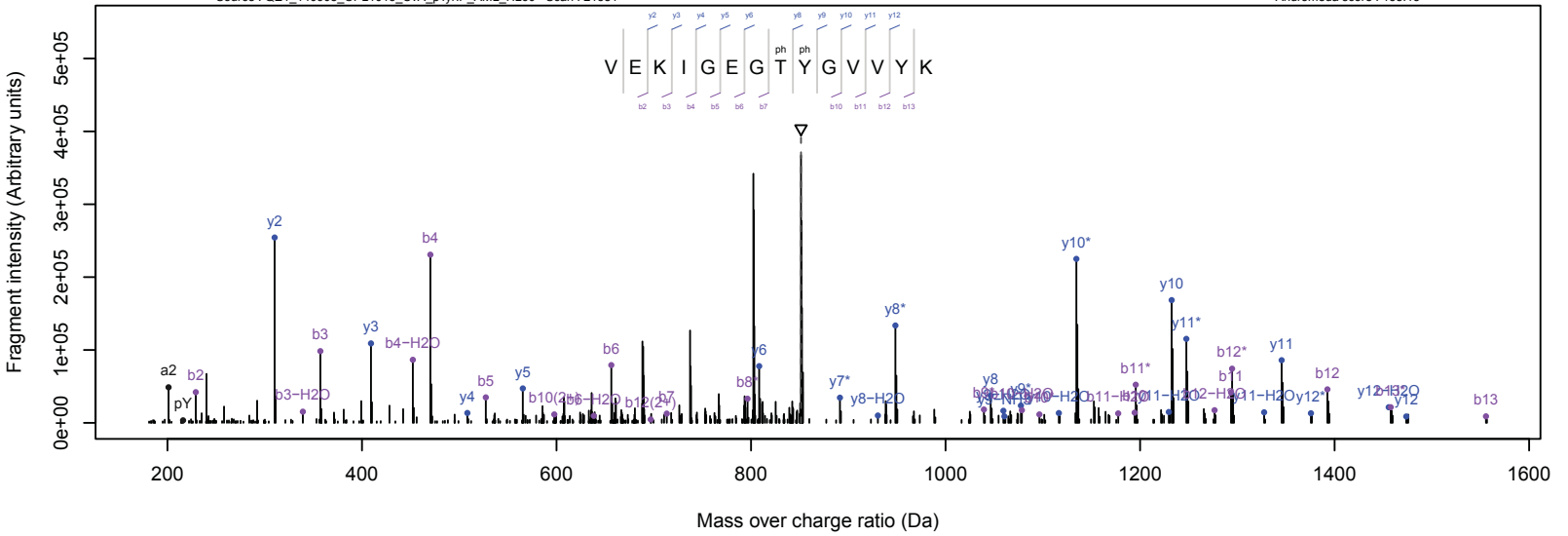
Andromeda score : 158.2



VEKIGEGT(ph)Y(ph)GVVYK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 21581

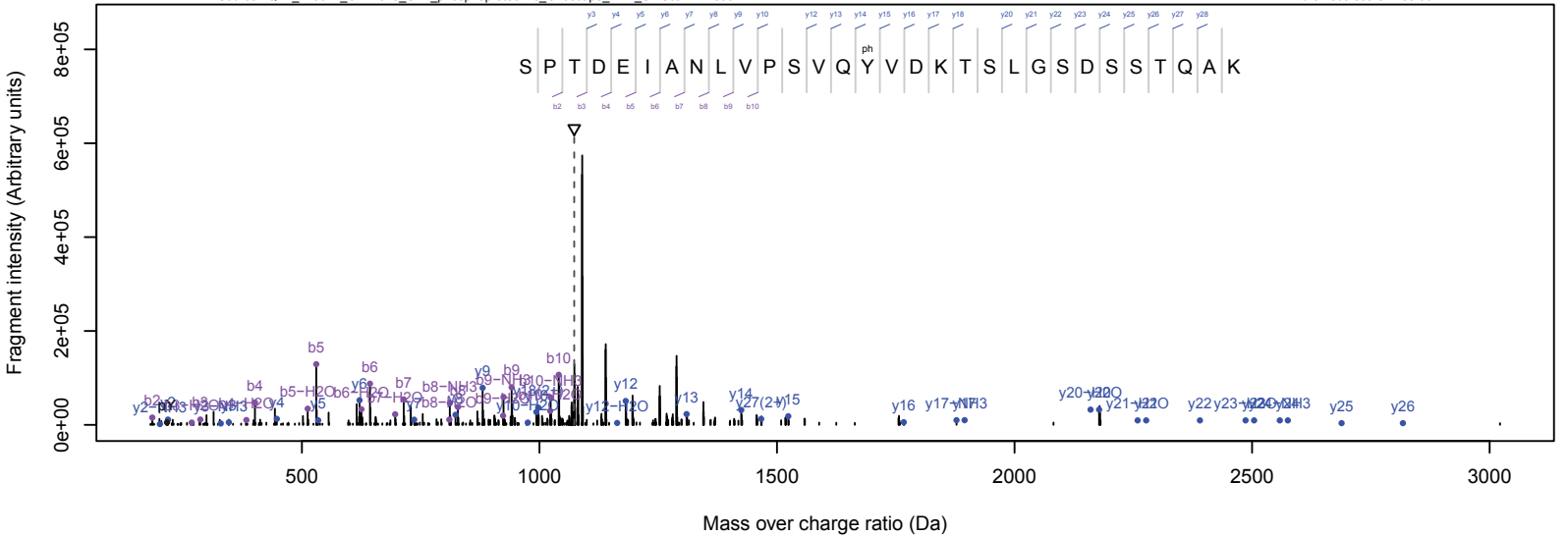
Andromeda score : 158.15



SPTDEIANLVPSVQY(ph)VDKTSLGSDSSSTQAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42830

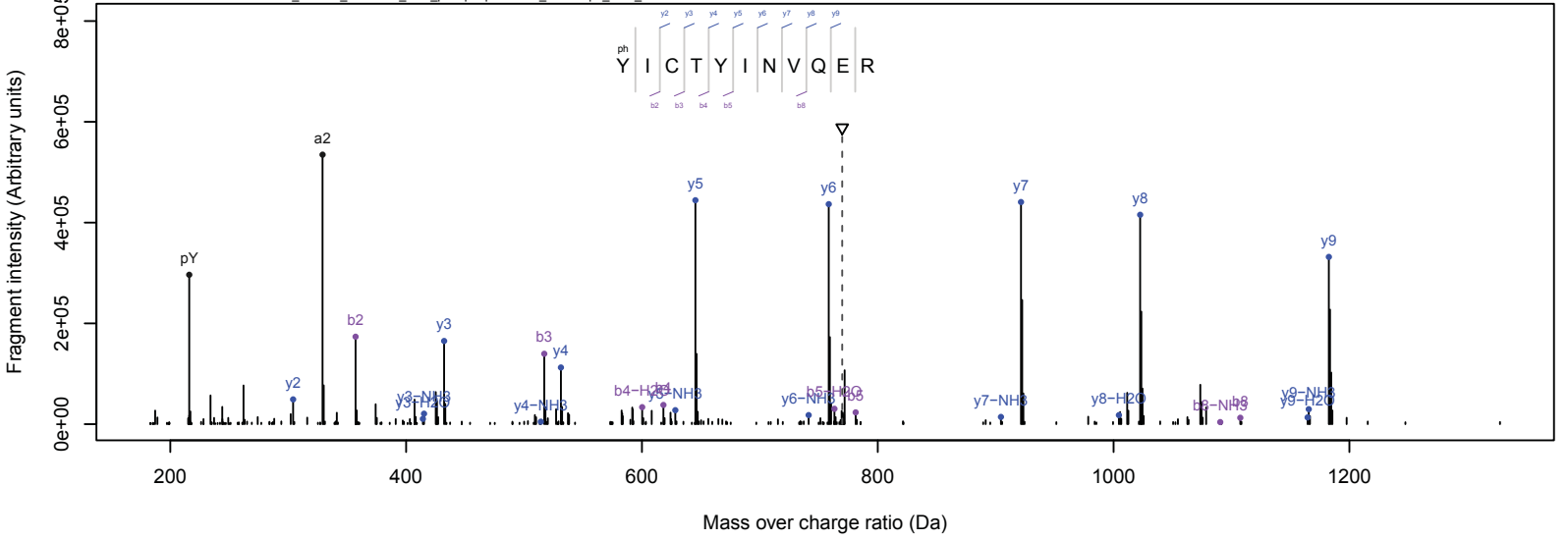
Andromeda score : 158.06



**Y(ph)ICTYINVQER-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26981

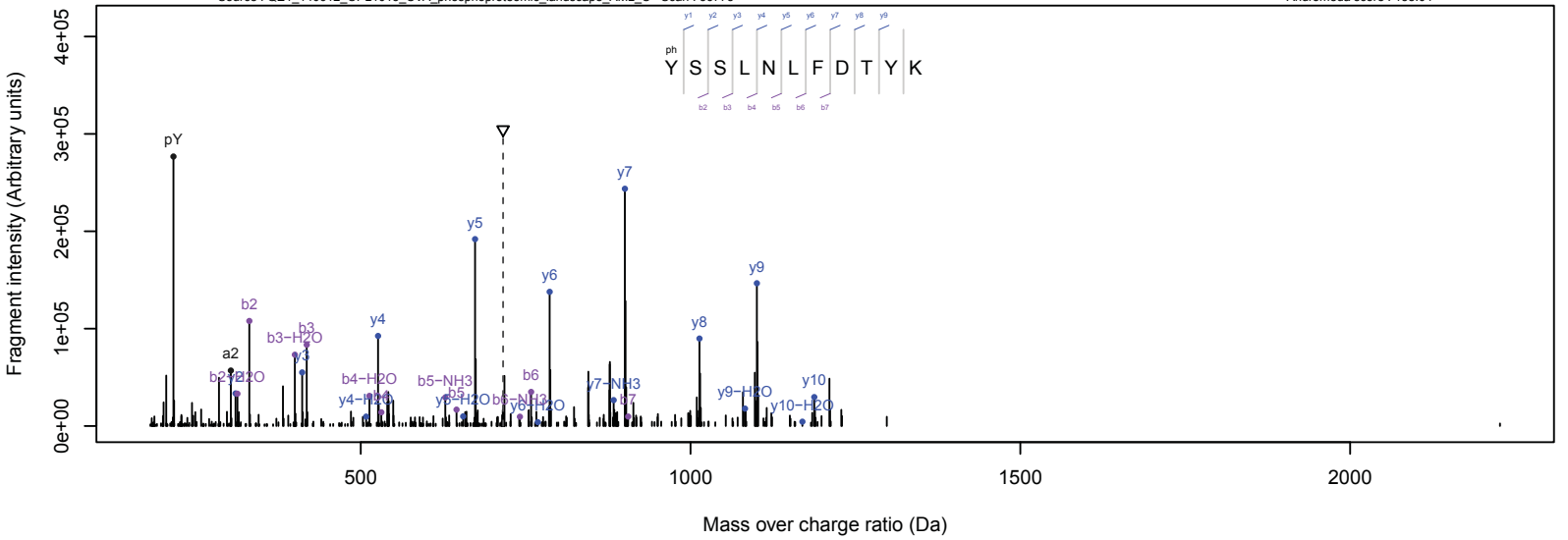
Andromeda score : 158.05



**Y(ph)SSLNLFDTYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36773

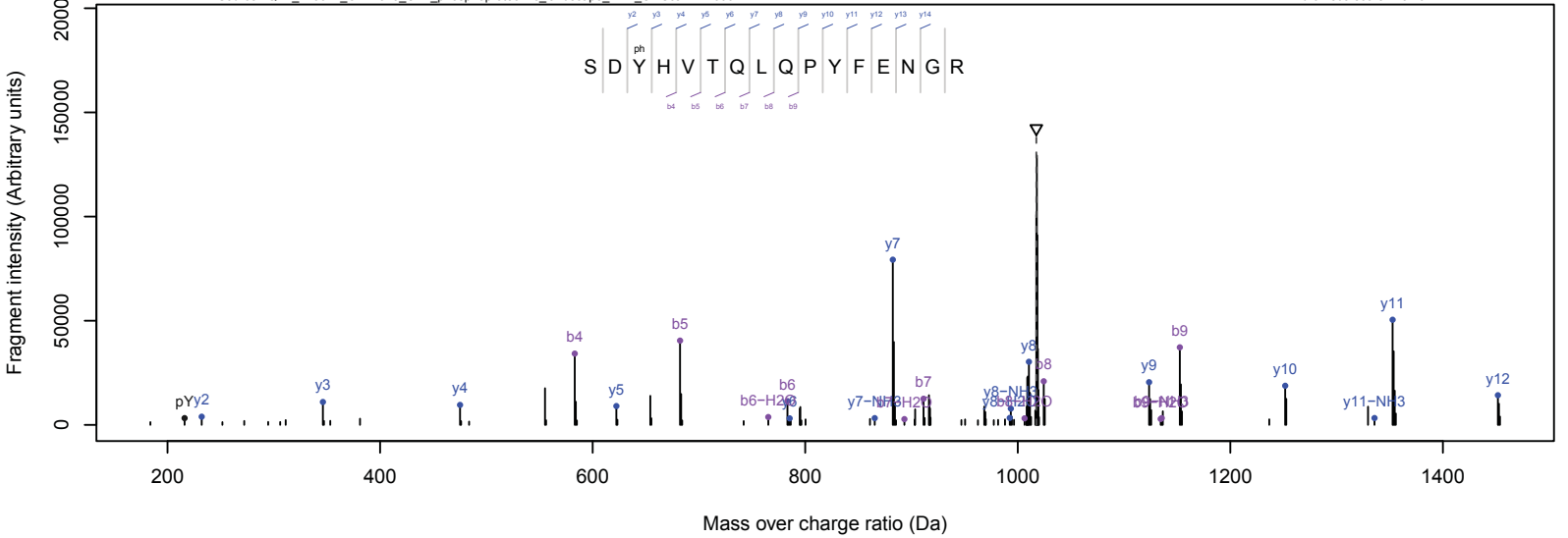
Andromeda score : 158.04



**SDY(ph)HVTQLQPYFENGR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27606

Andromeda score : 157.91

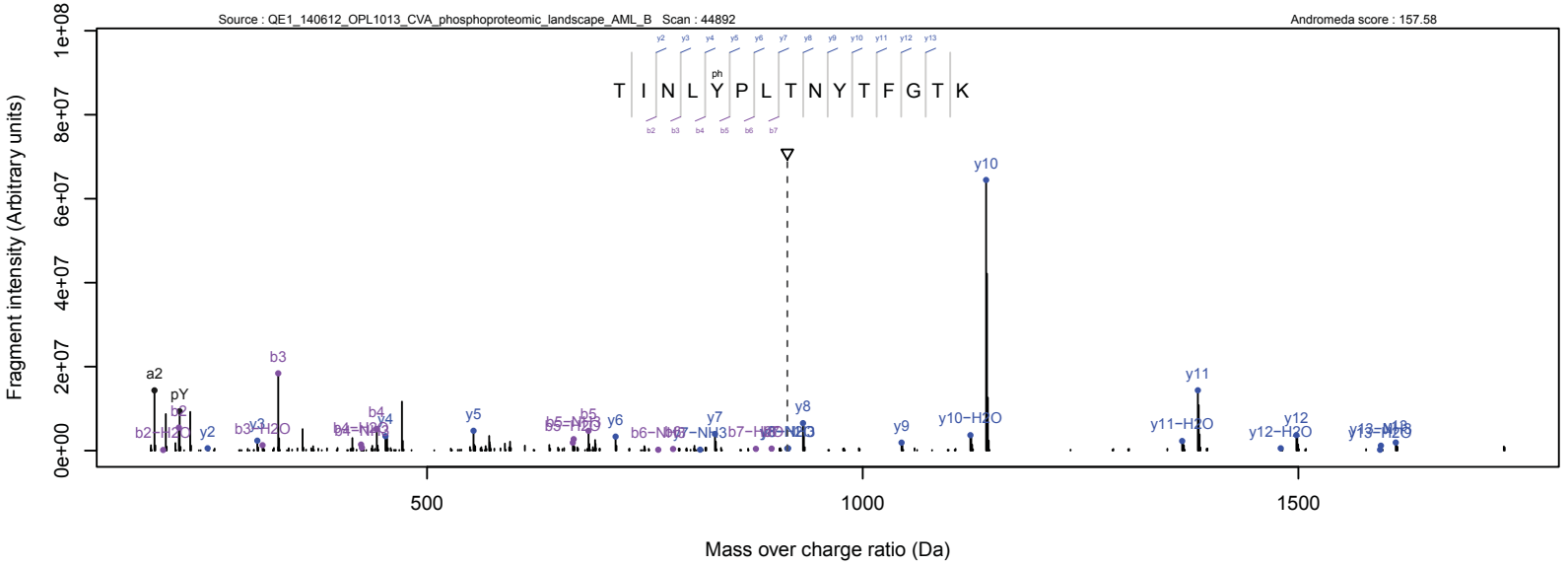




TINLY(ph)PLTNYTFGTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44892

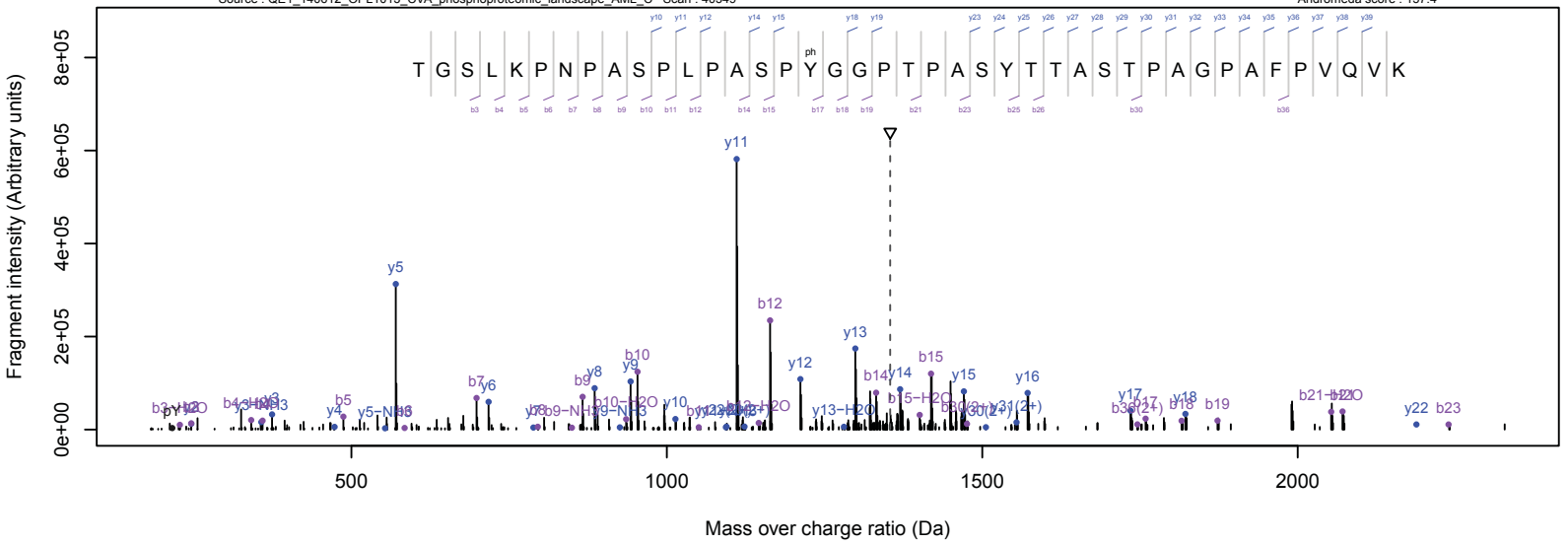
Andromeda score : 157.58



TGSLKPNPASPLPASPY(ph)GGPTASYYTASTPAGPAFPVQVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40349

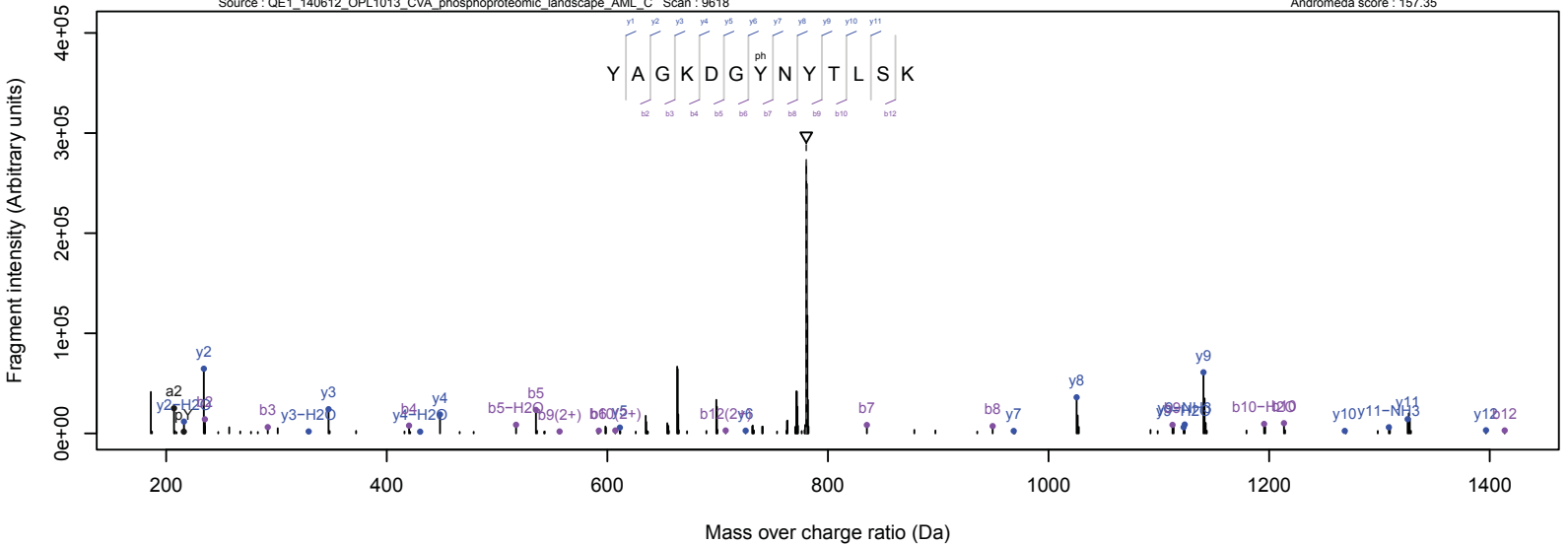
Andromeda score : 157.4



YAGKDGYNYSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9618

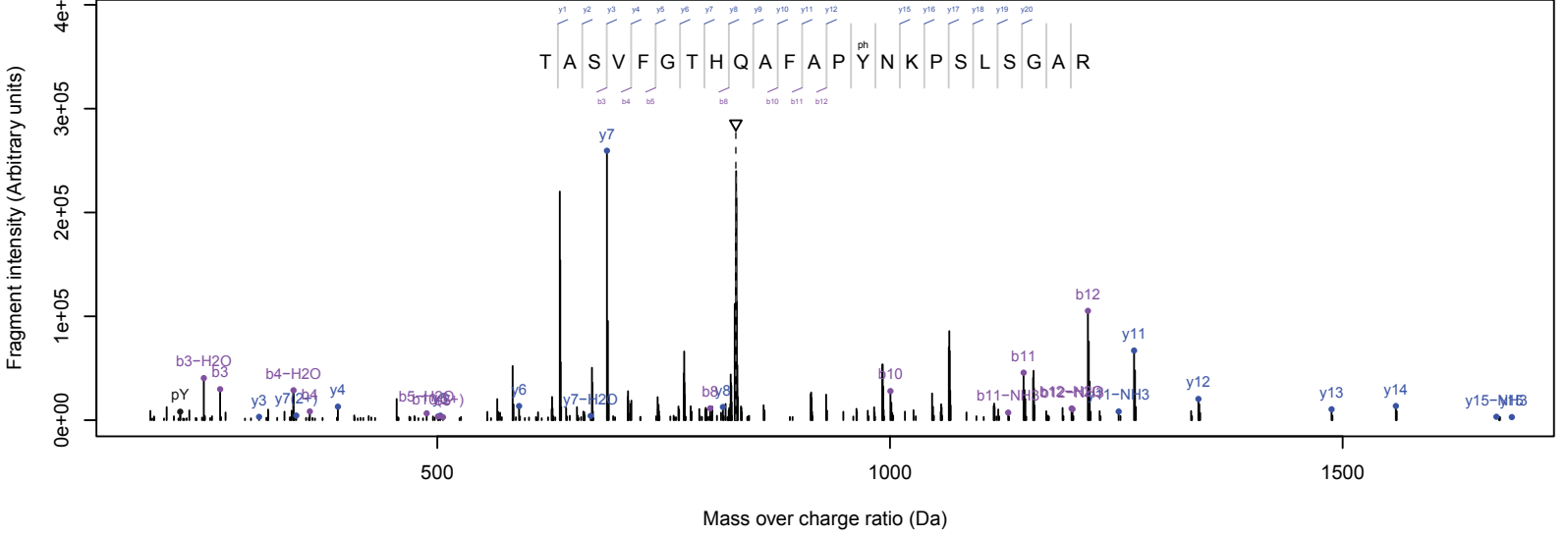
Andromeda score : 157.35



TASVFGTHQAFAPY<sup>ph</sup>NKPSLSGAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25635

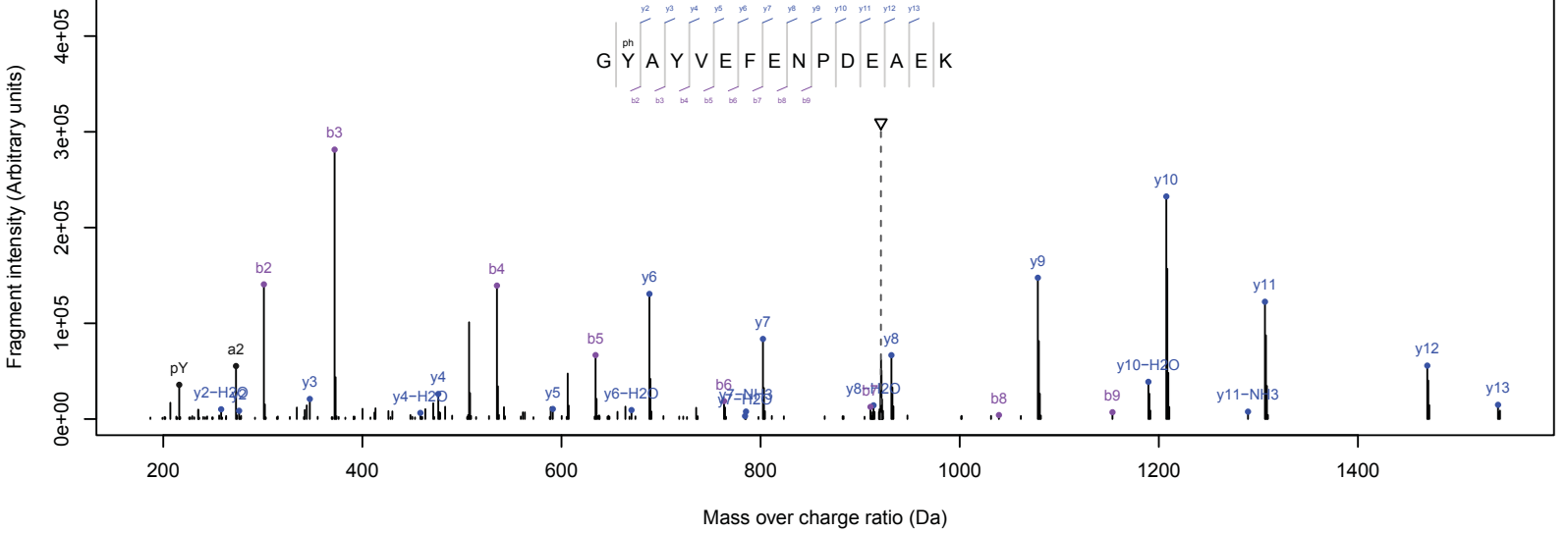
Andromeda score : 157.3



GY<sup>ph</sup>AYVEFENPDEAEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26805

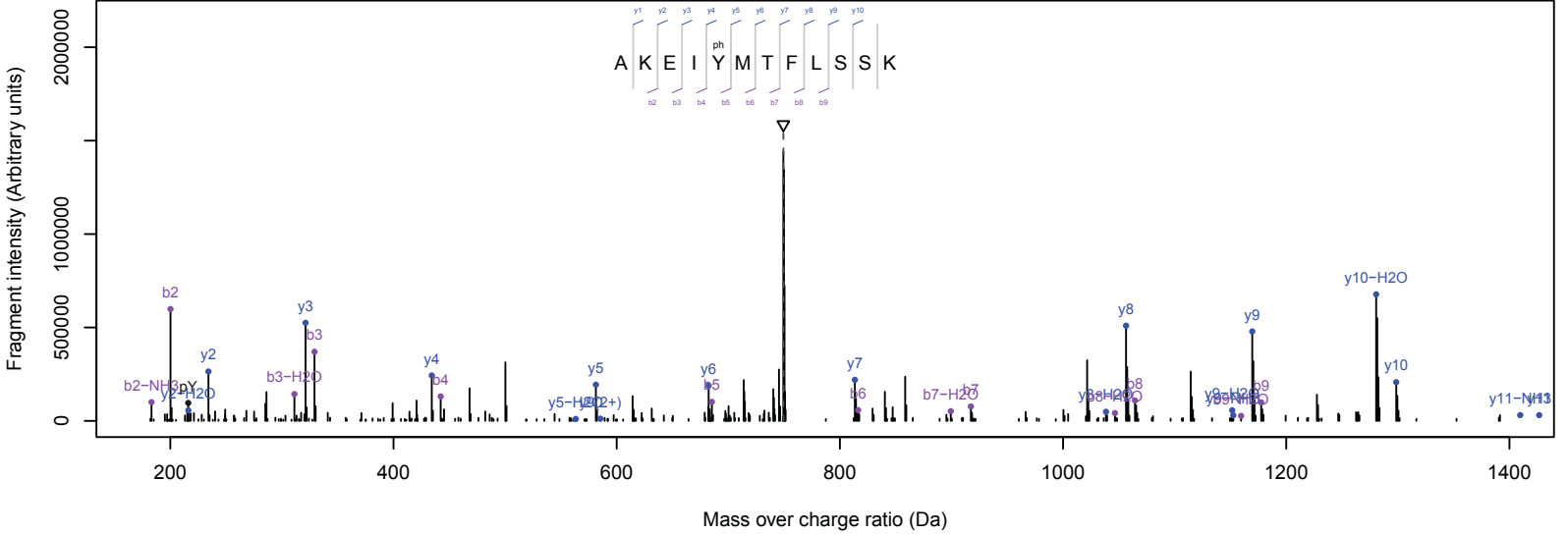
Andromeda score : 157.23



AKEIY<sup>ph</sup>MTFLSSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30289

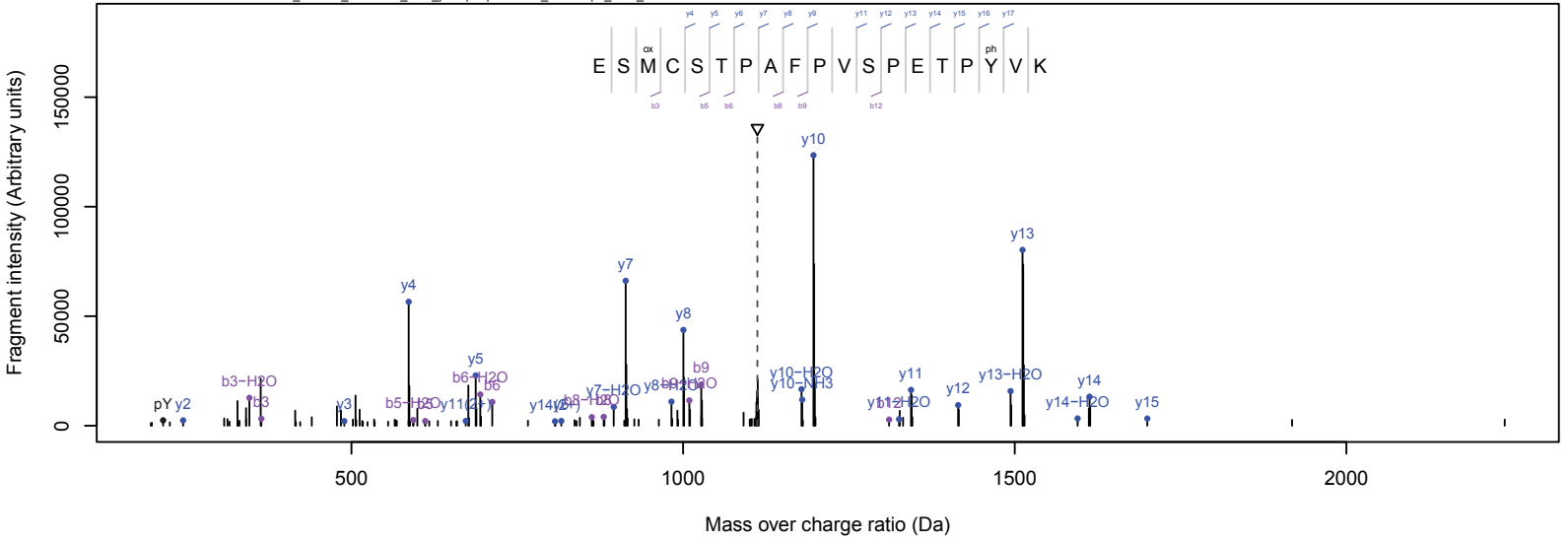
Andromeda score : 157.2



ESM(ox)CSTPAFPVSPETPYVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33475

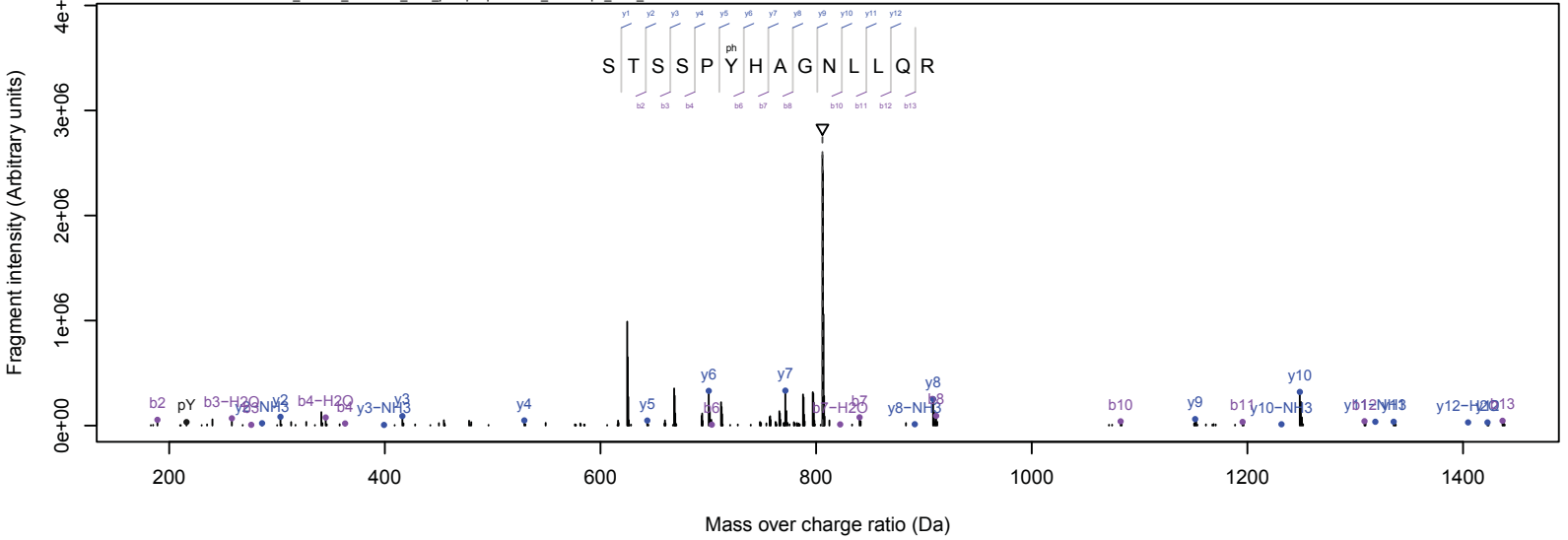
Andromeda score : 157.11



STSSPY(H)HAGNLLQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13906

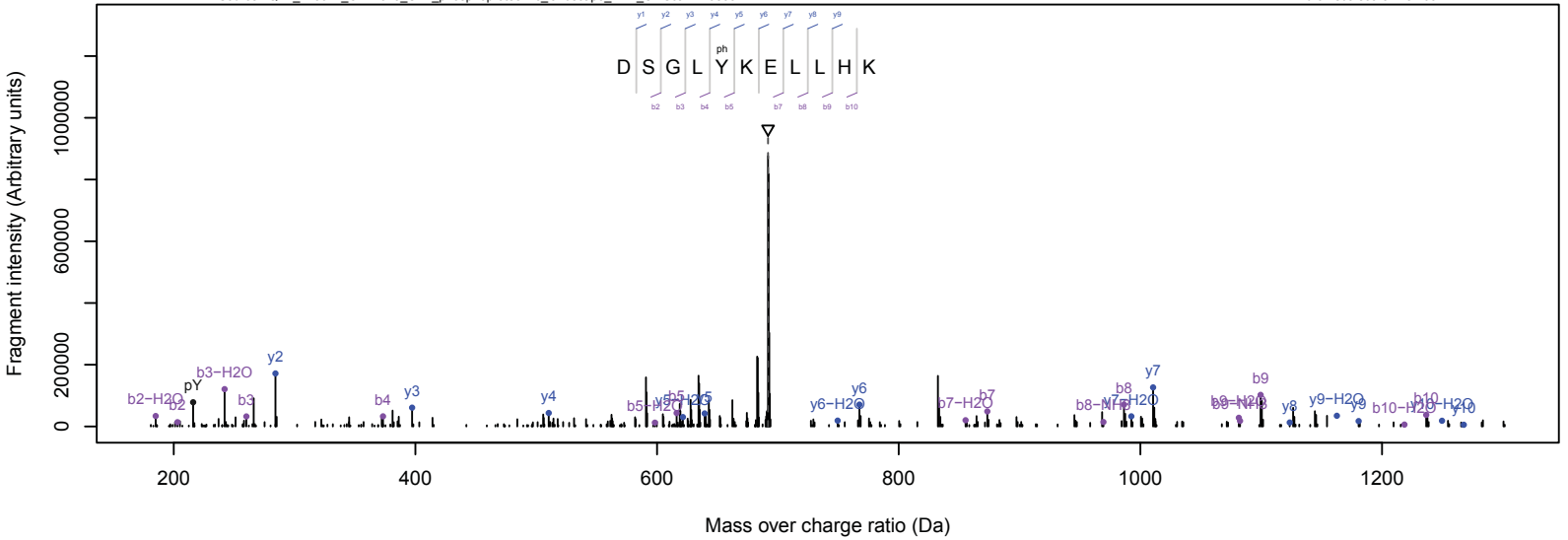
Andromeda score : 157.09



DSGLY(ph)KELLHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19833

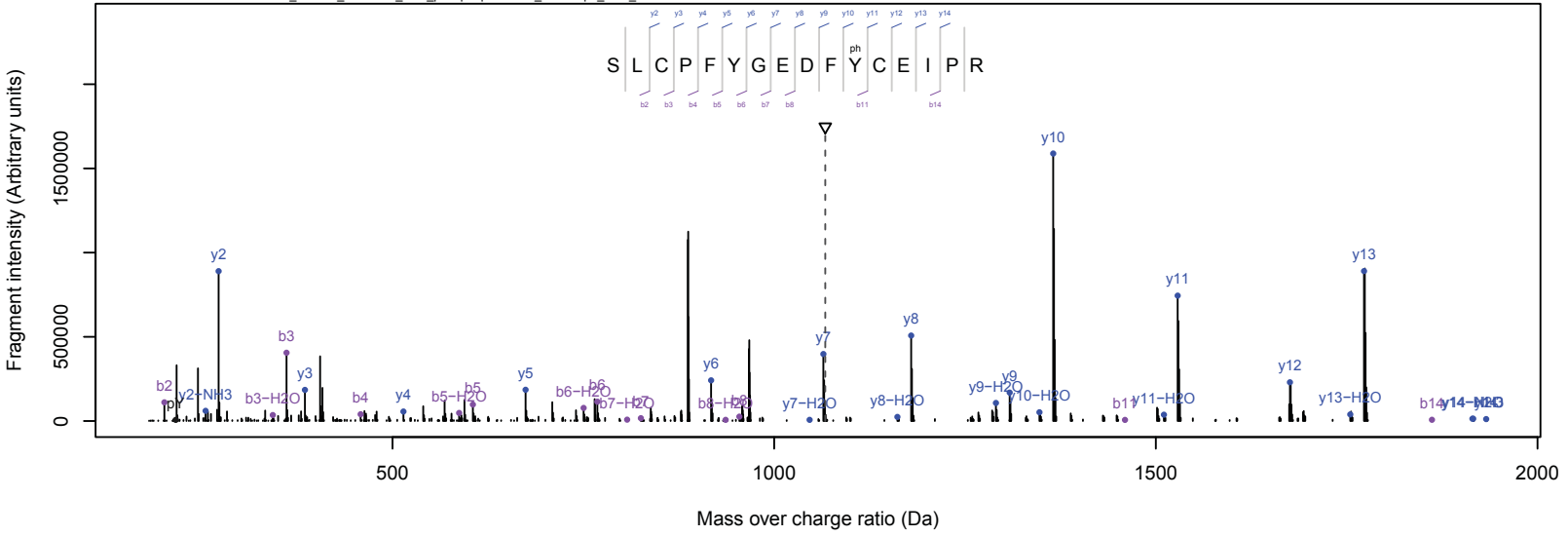
Andromeda score : 157.03



**SLCPFYGEDFY(ph)CEIPR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43960

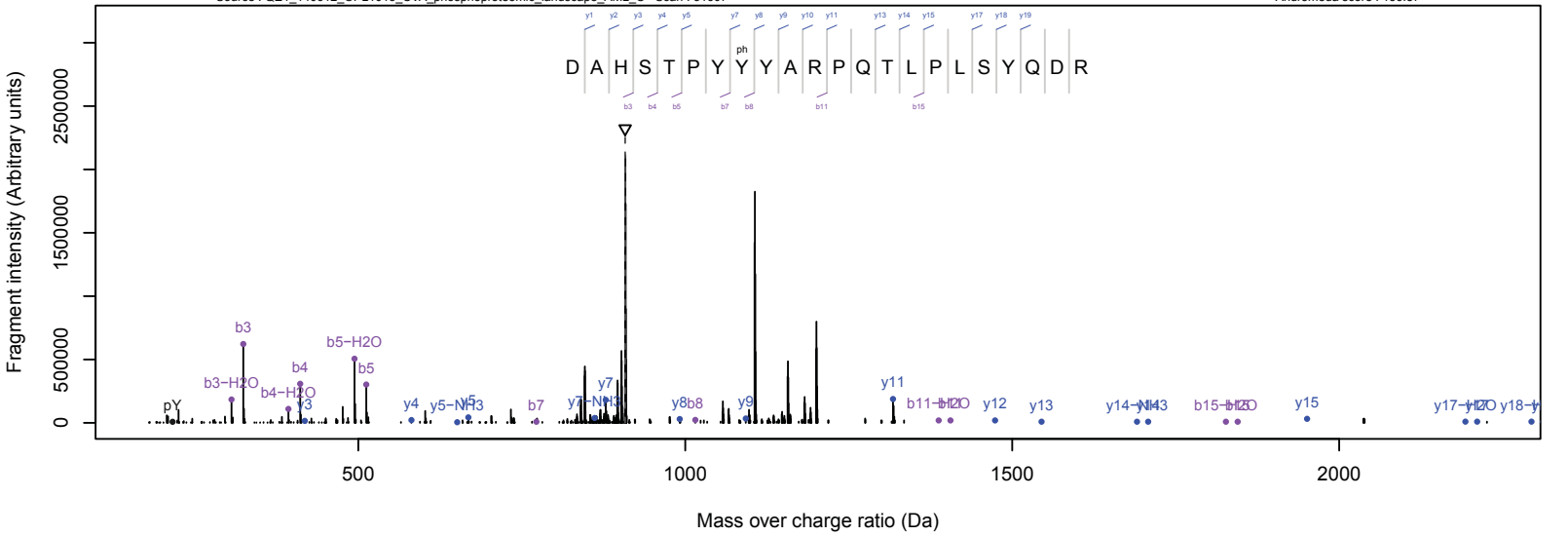
Andromeda score : 156.98



**DAHSTPYY(ph)YARPQTLPLSYQDR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31607

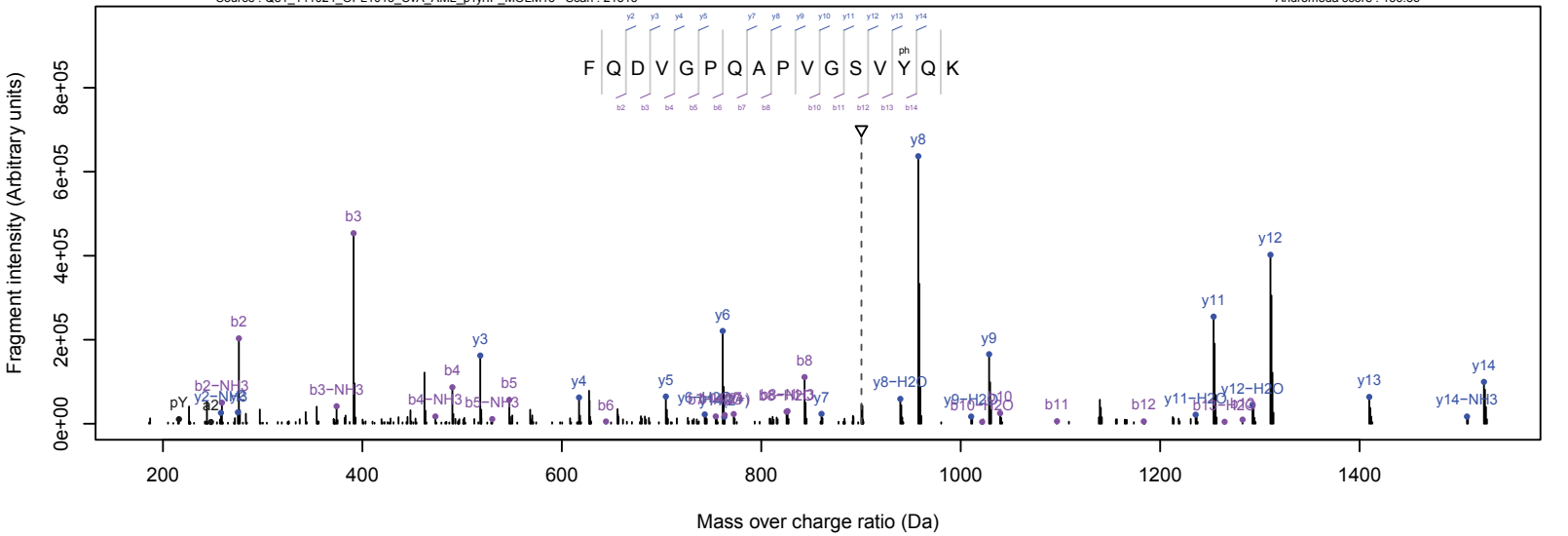
Andromeda score : 156.87



**FQDVGPQAPVGSVY(ph)QK-2+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 21813

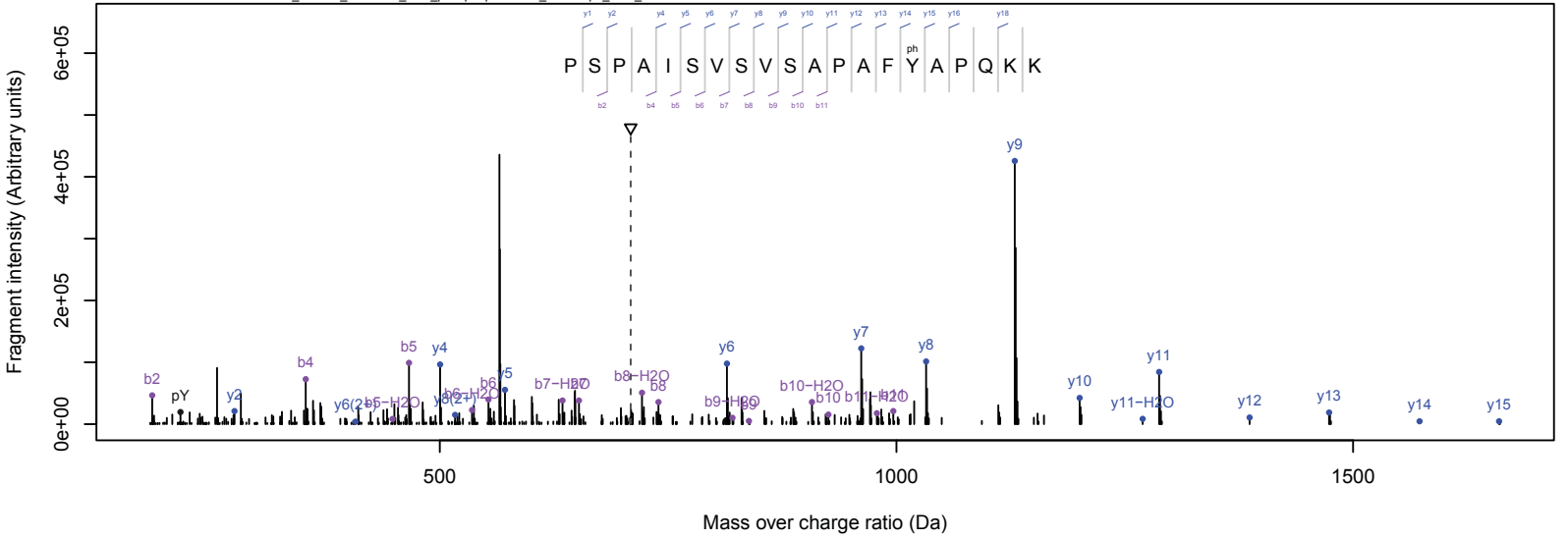
Andromeda score : 156.86



PSPAISVSVSAPAFY(ph)APQKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32675

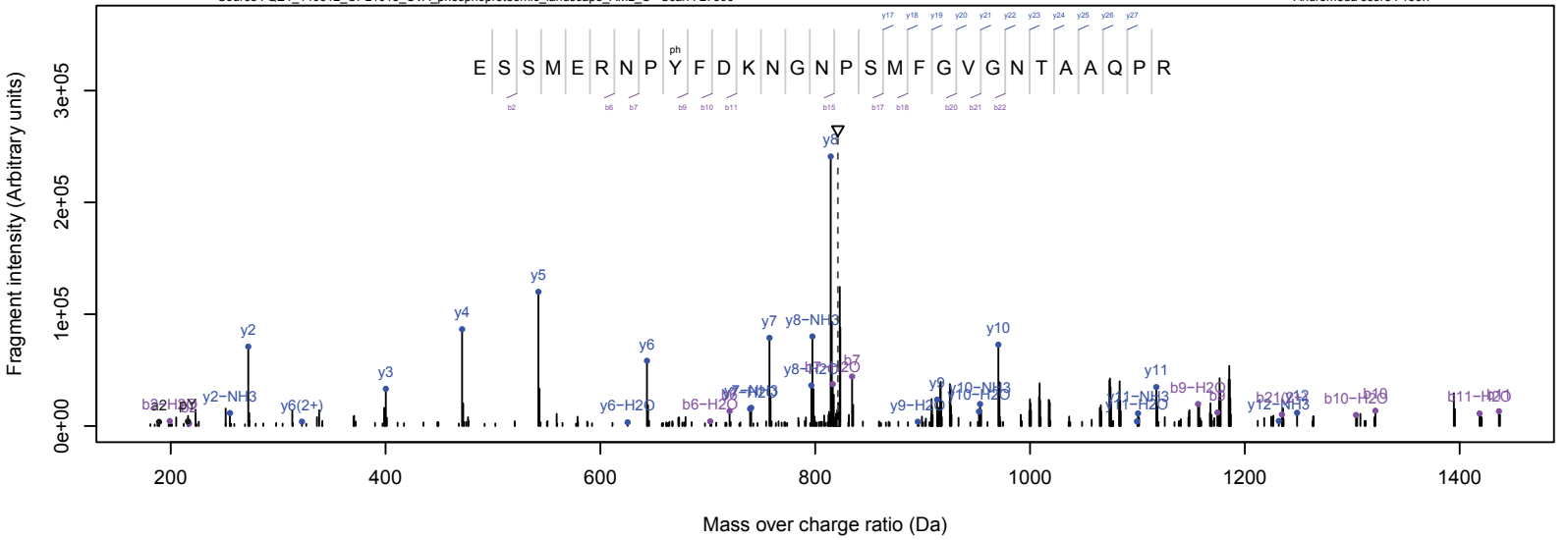
Andromeda score : 156.73



ESSMERNPY(ph)FDKNGNPSMFGVGNNTAAQPR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27396

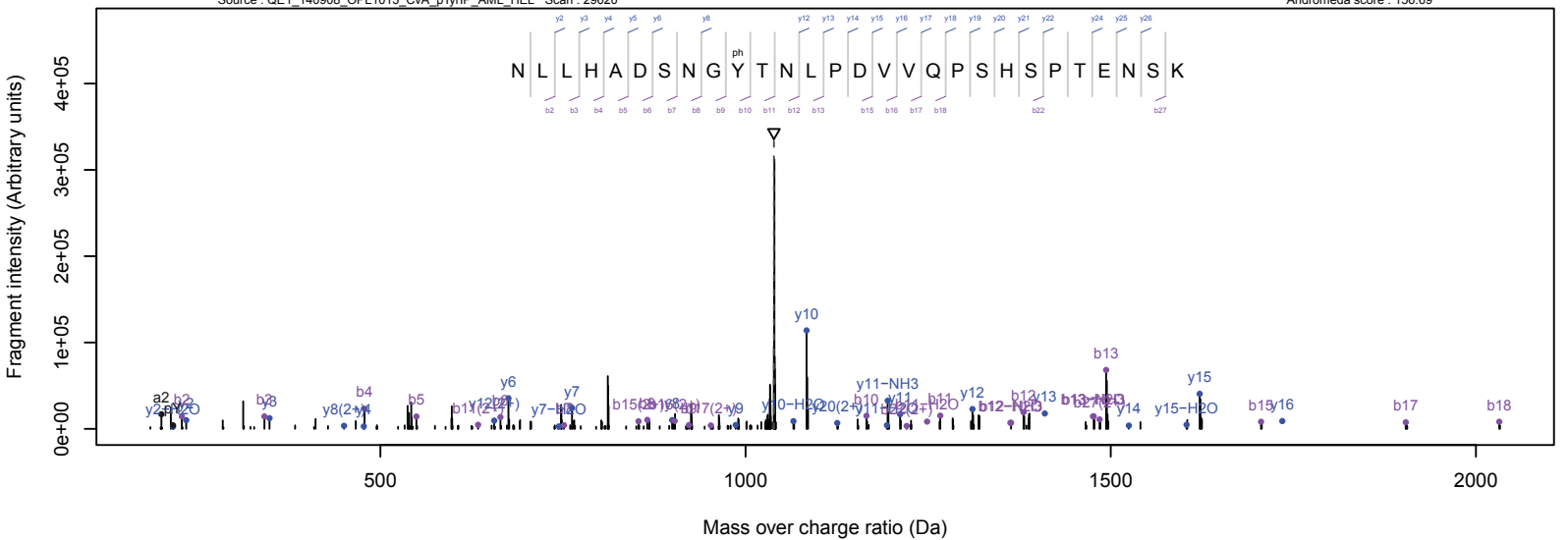
Andromeda score : 156.7



NLLHADSNGY(ph)TNLPDVVQPSHSPTENSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29620

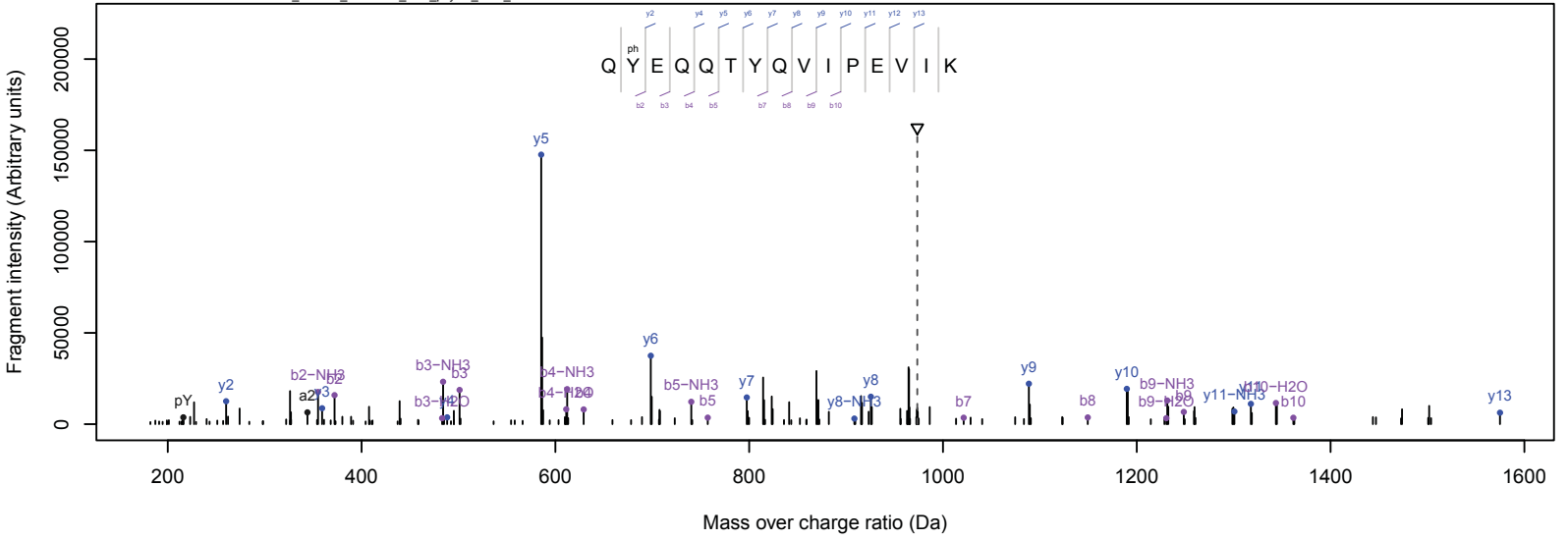
Andromeda score : 156.69



QY(ph)EQQTYQVIPEVIK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37723

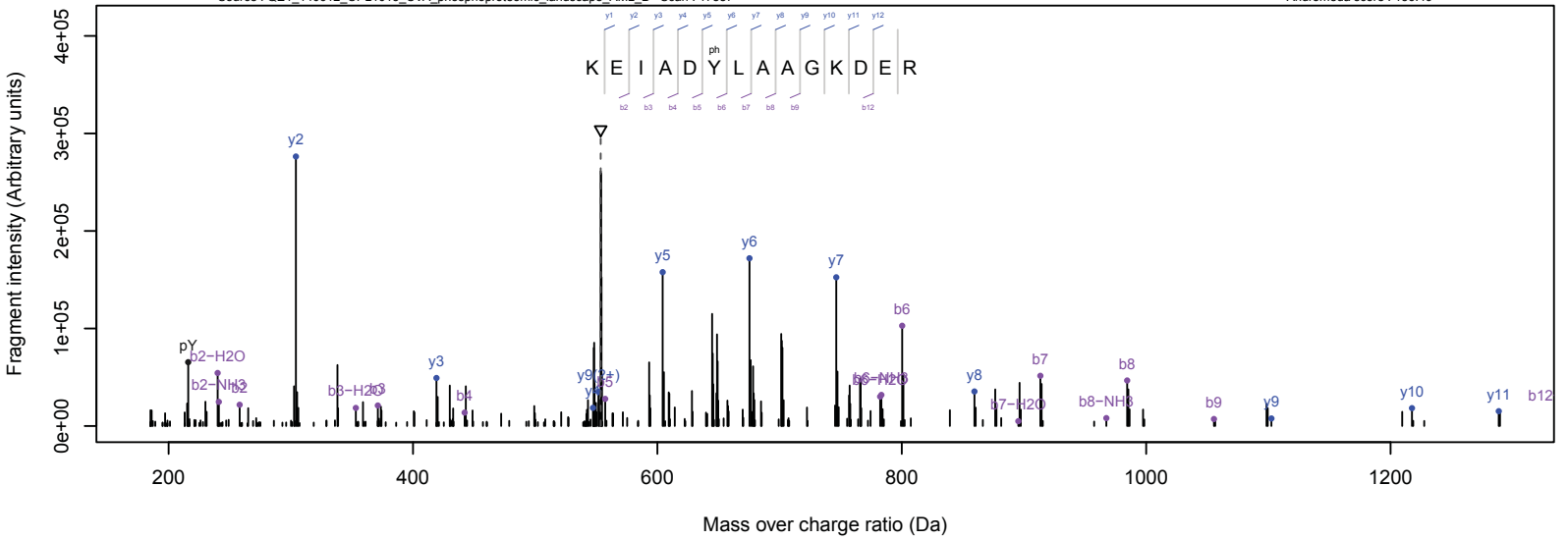
Andromeda score : 156.57



KEIADY(ph)LAAGKDER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17887

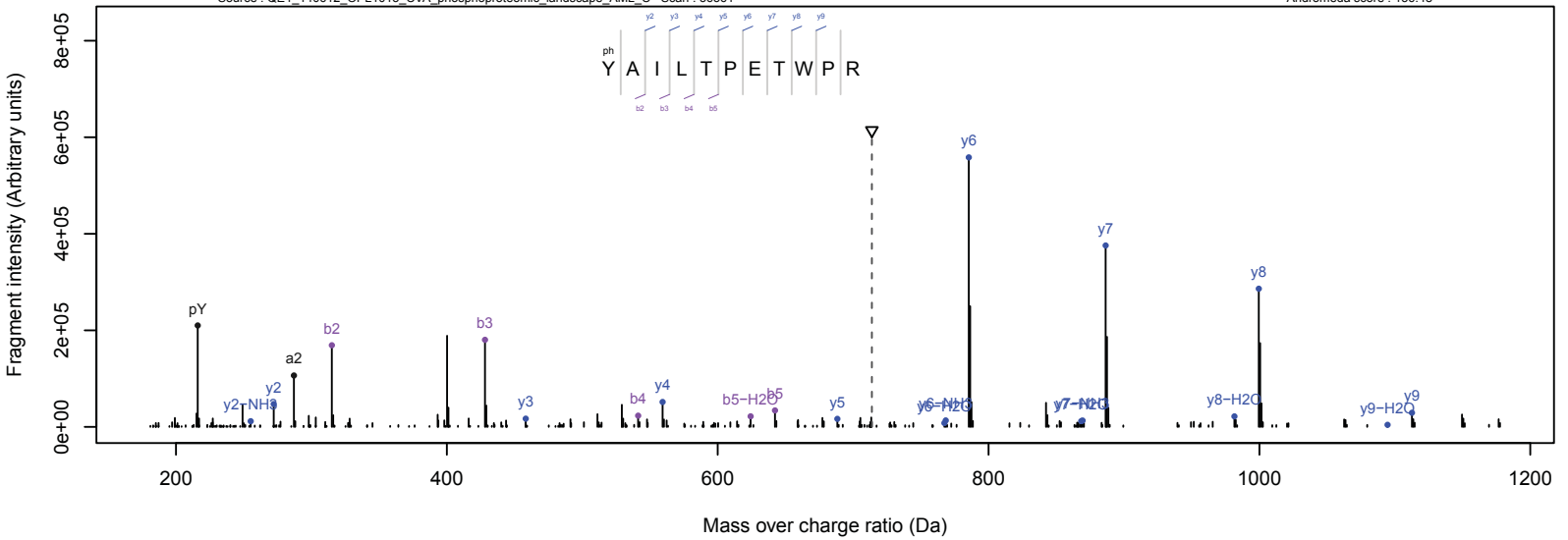
Andromeda score : 156.49



Y(ph)AILTPEWPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36361

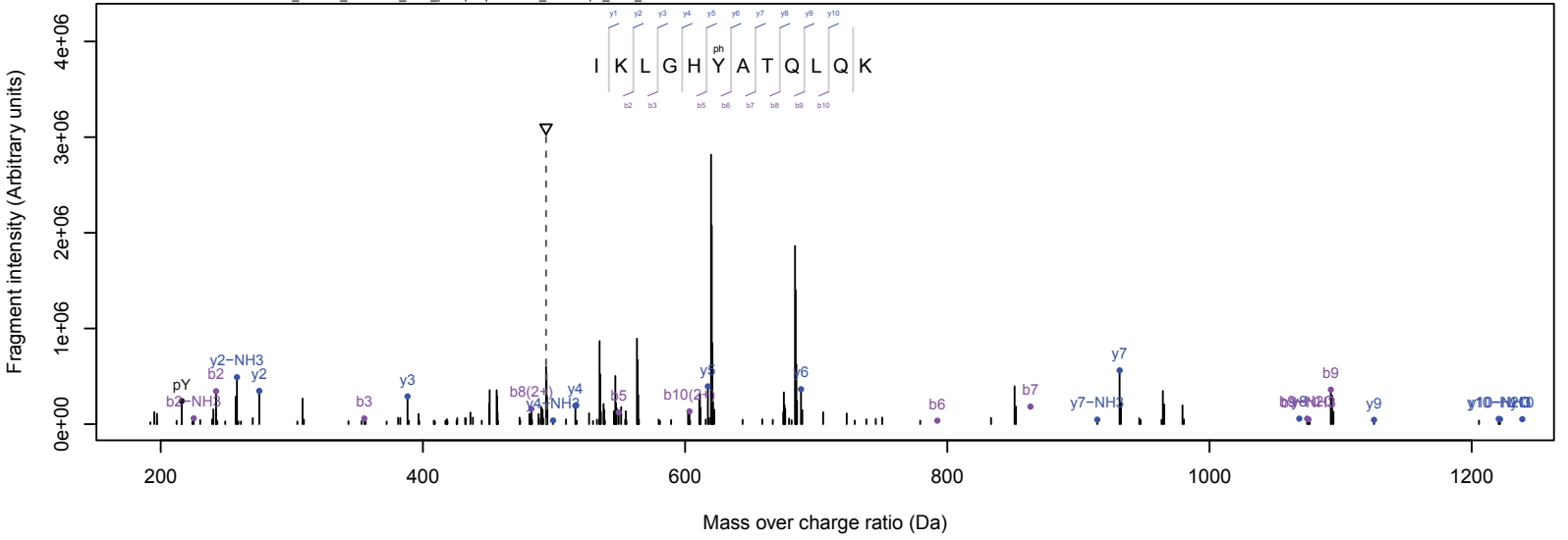
Andromeda score : 156.48



**IKLGHY(ph)ATQLQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15156

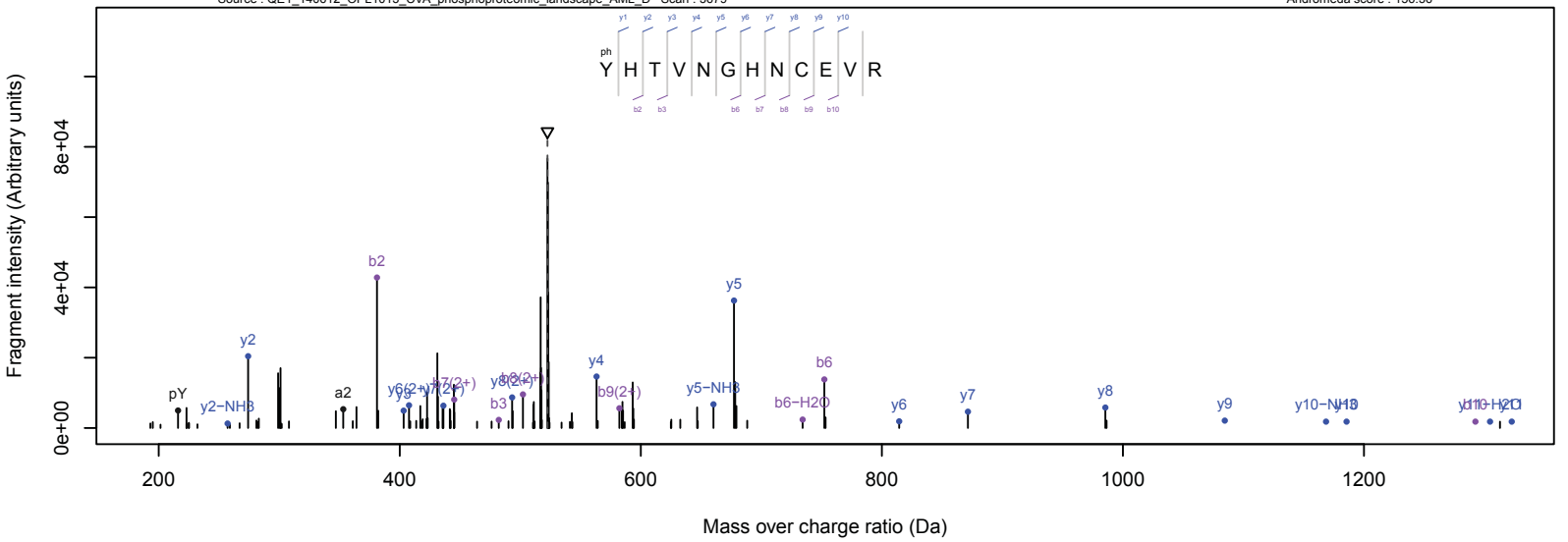
Andromeda score : 156.36



**Y(ph)HTVNGHNCEVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 3679

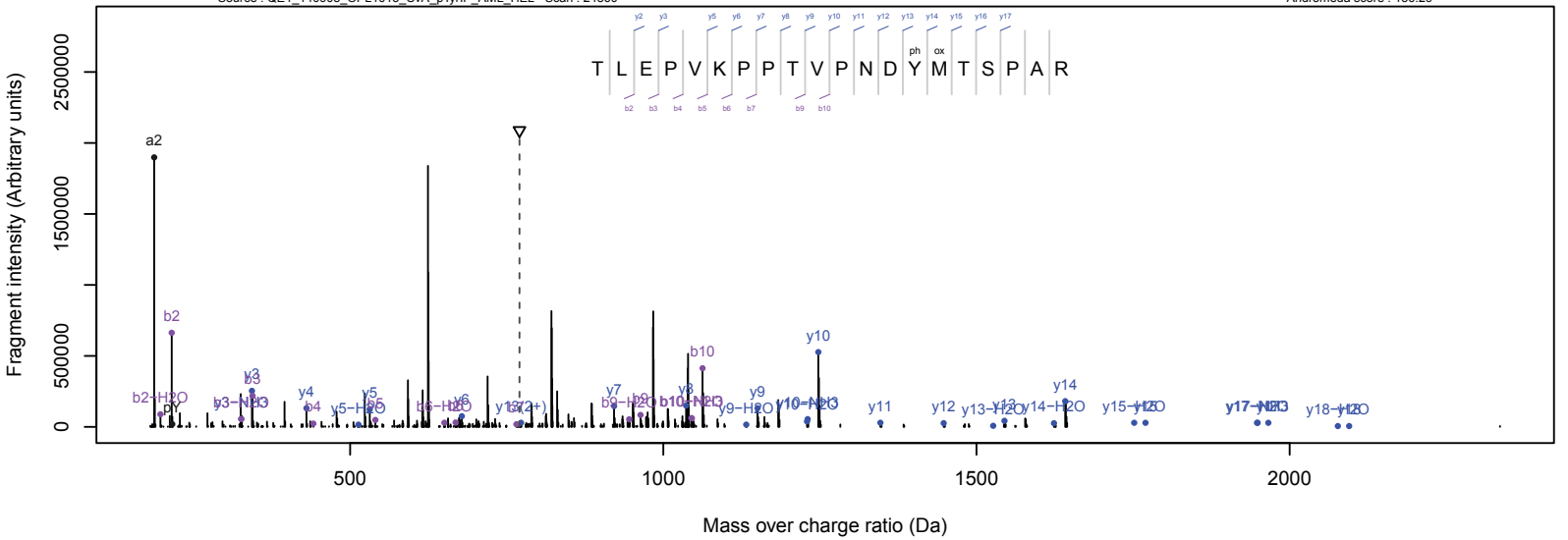
Andromeda score : 156.36



**TLEPVKPTVPNDY(ph)M(ox)TSPAR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24860

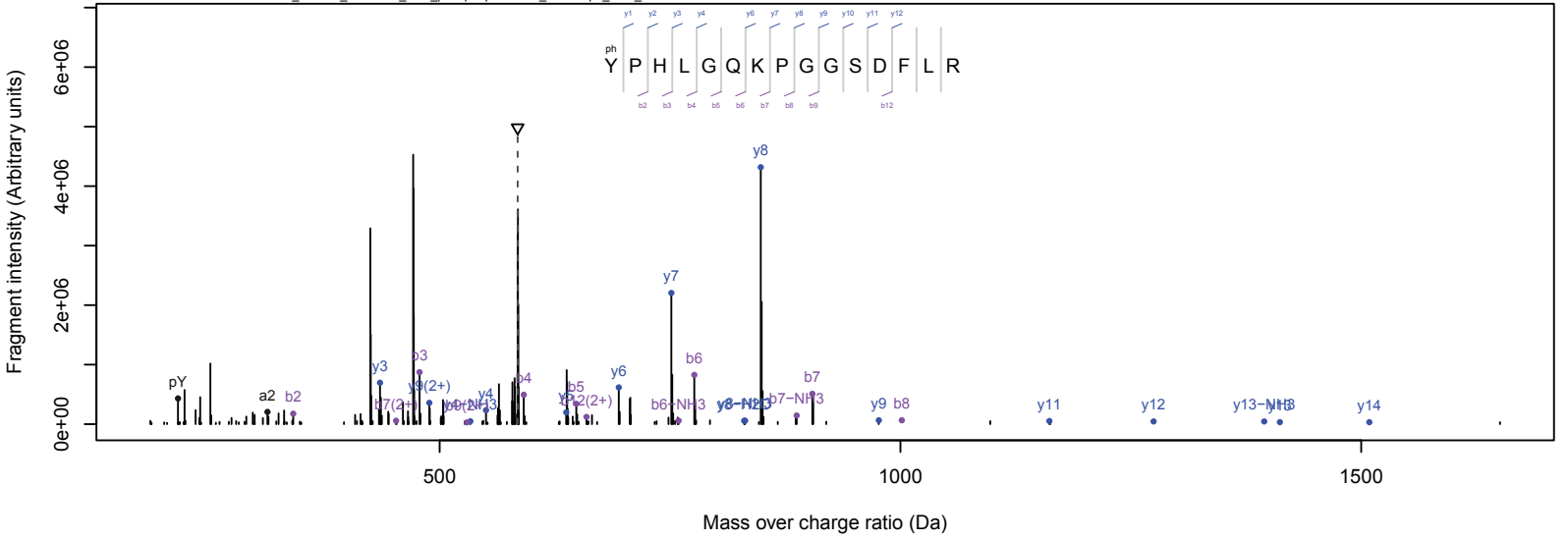
Andromeda score : 156.29



**Y(ph)PHLGQKPGGSDFLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21576

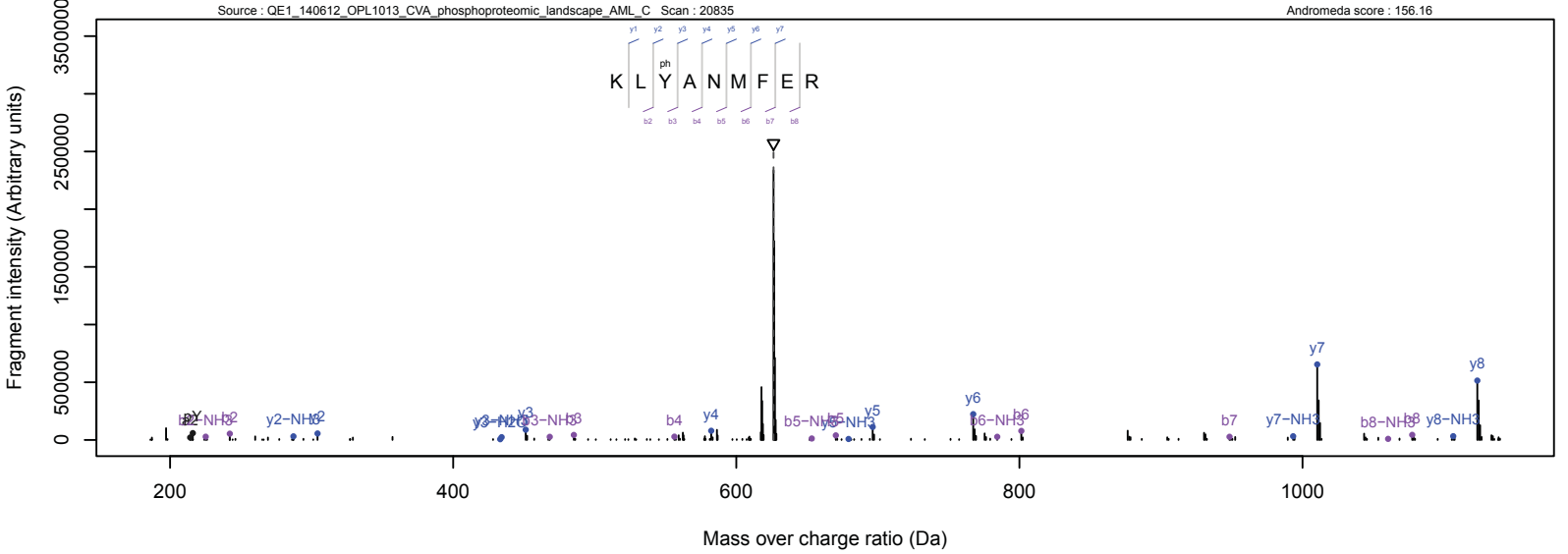
Andromeda score : 156.24



**KLY(ph)ANMFER-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20835

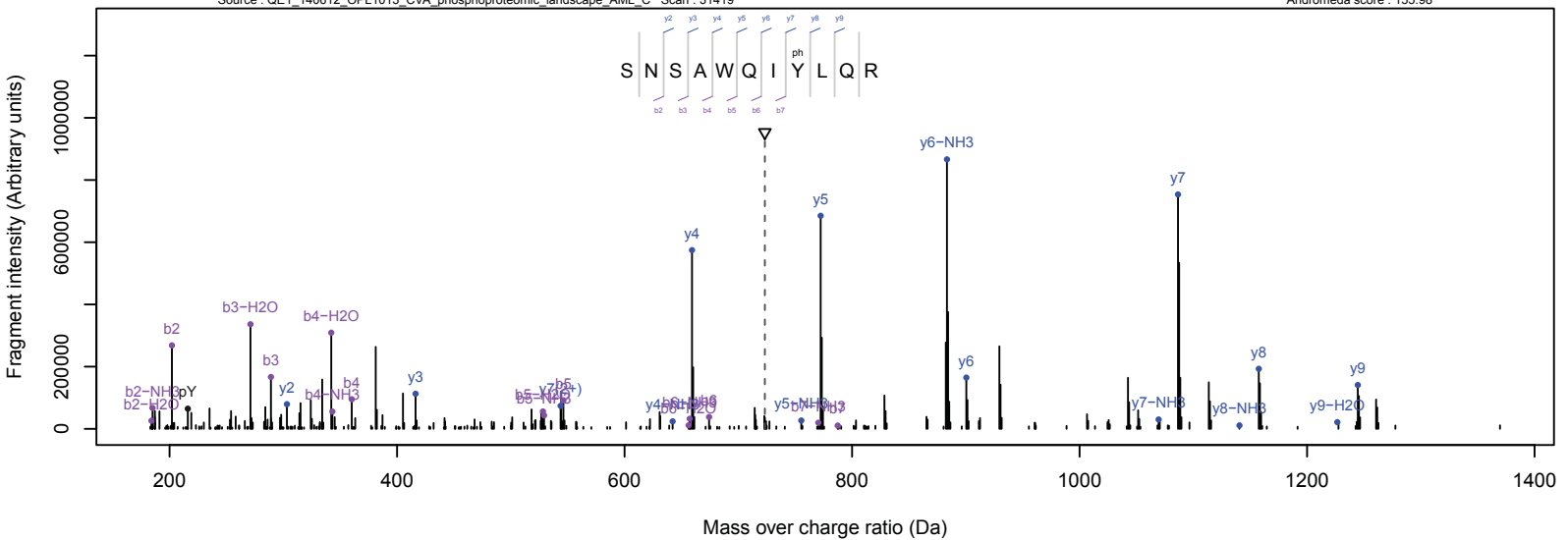
Andromeda score : 156.16



**SNSAWQIY(ph)LQR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31419

Andromeda score : 155.98

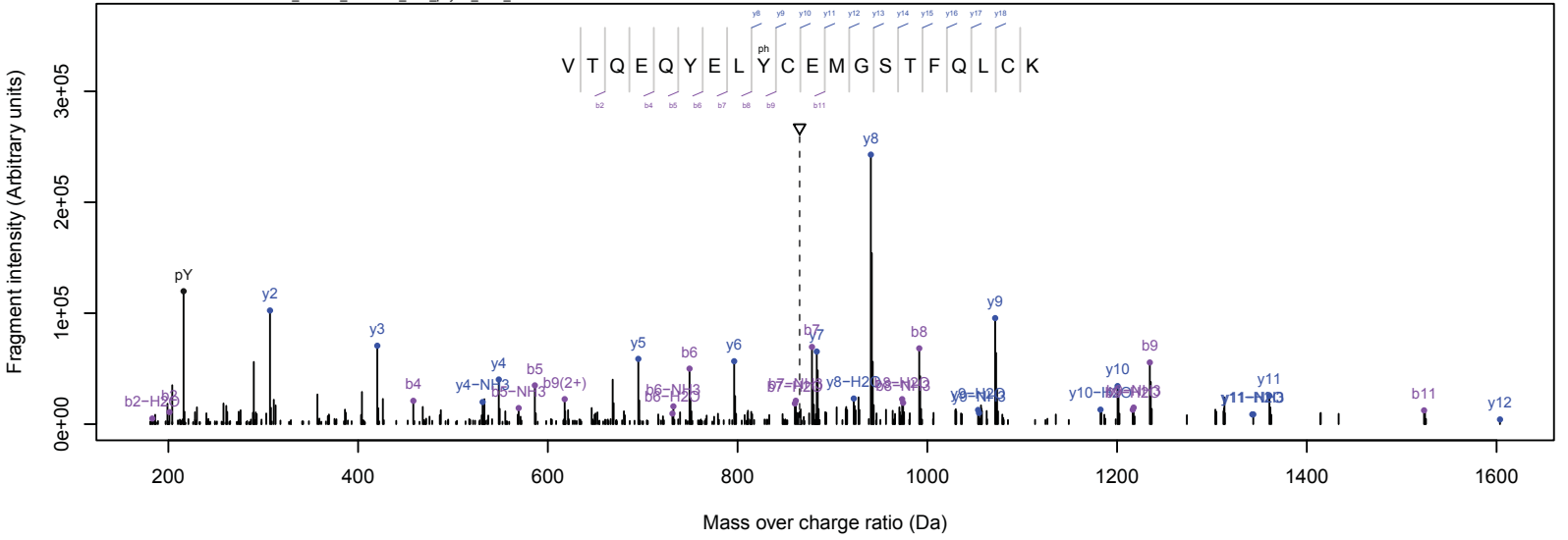




VTQEYELY(ph)CEMGSTFQLCK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 43720

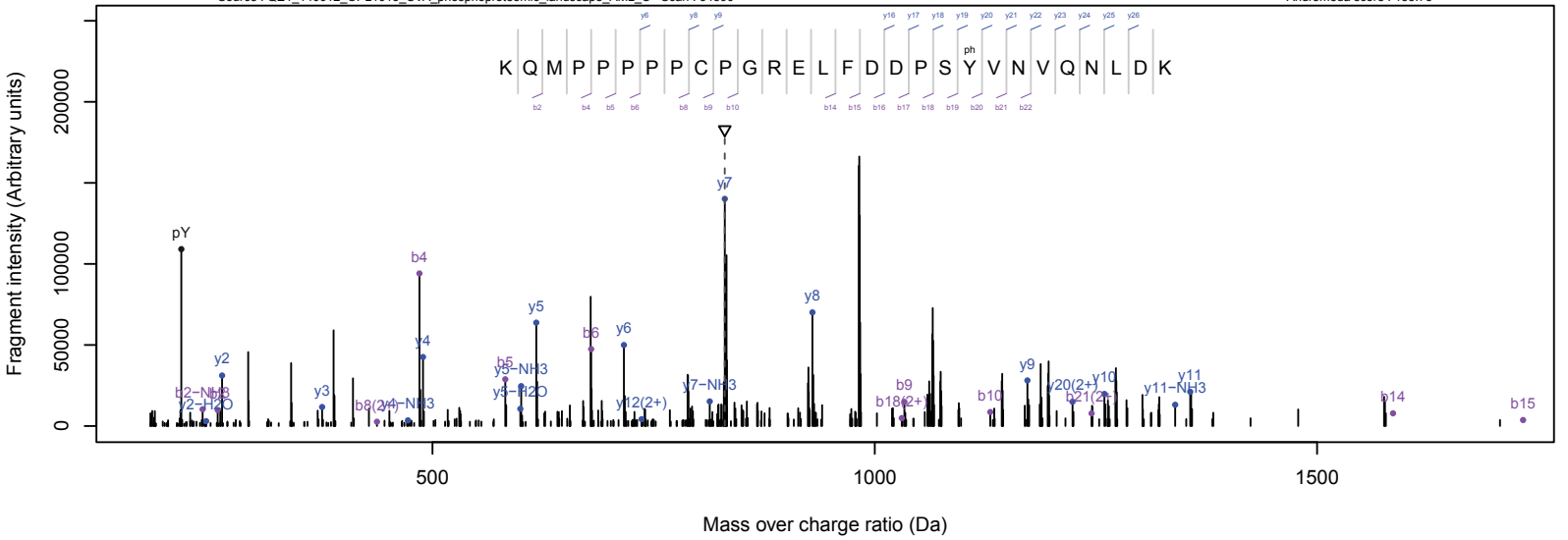
Andromeda score : 155.94



KQMPPPPPCGRELFDDPSY(ph)VNVQLDK-4+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 34590

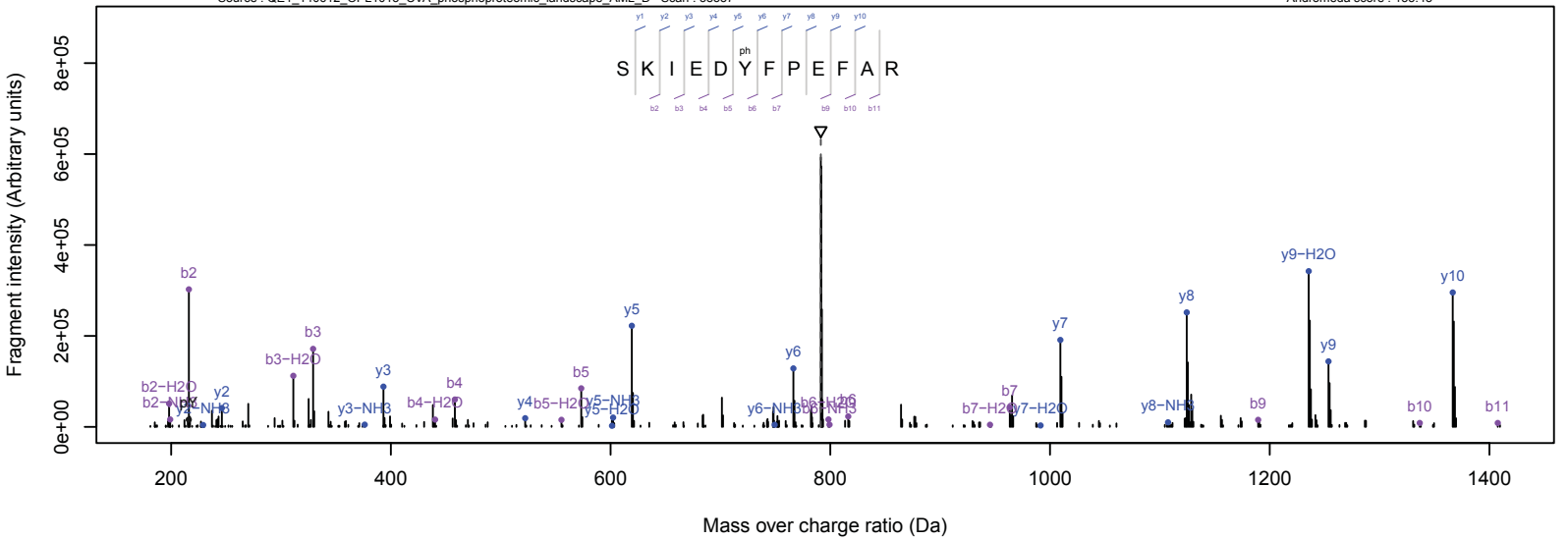
Andromeda score : 155.73



SKIEDY(ph)FPEFAR-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan : 38357

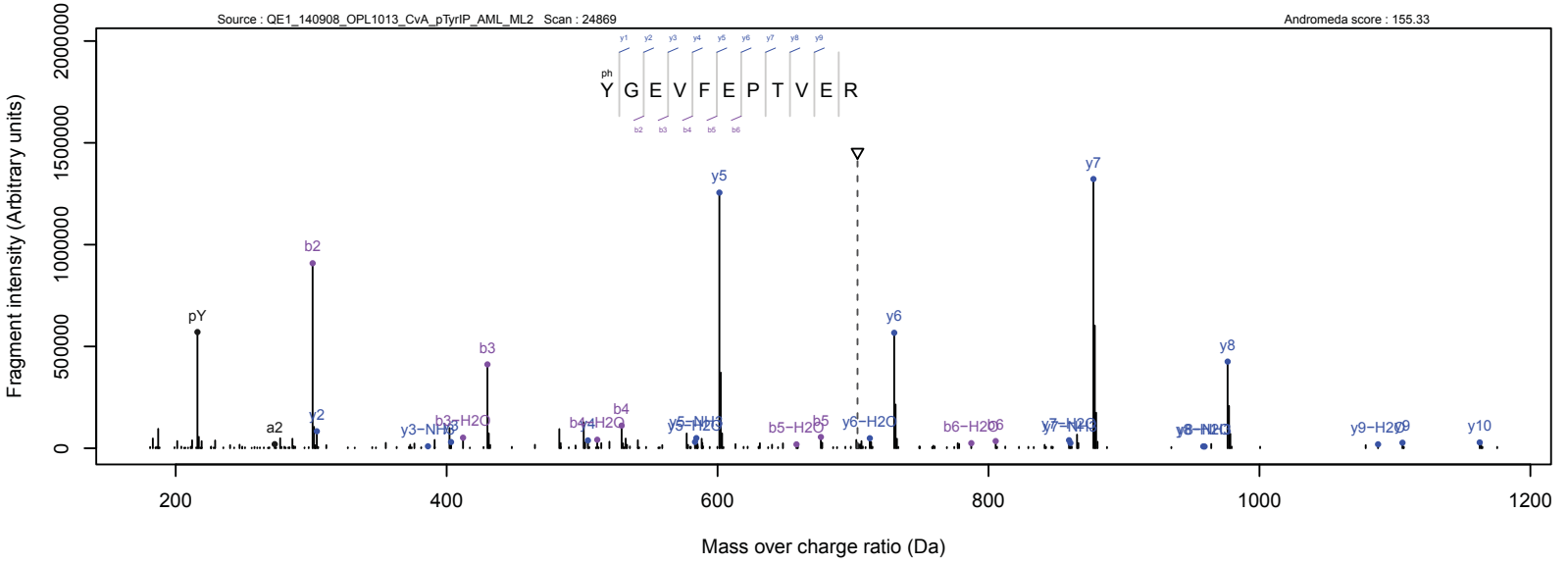
Andromeda score : 155.43



Y(ph)GEVFPTVER-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan: 24869

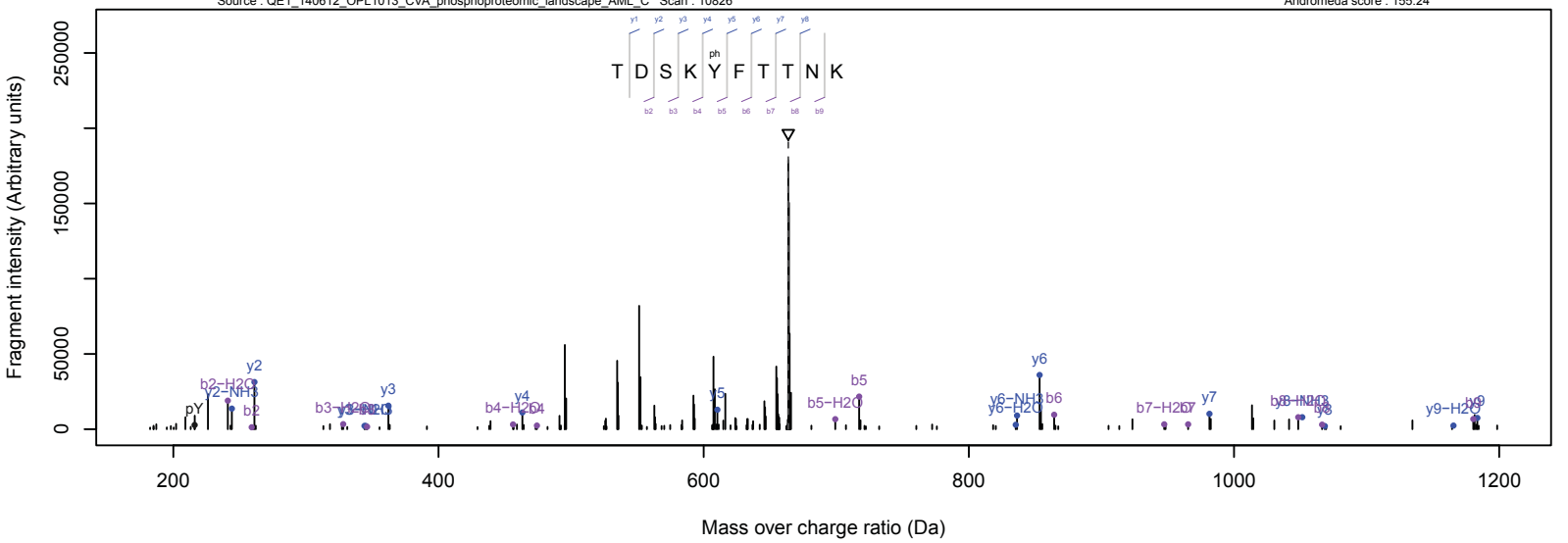
Andromeda score: 155.33



(ac)TDSKY(ph)FTTNK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10826

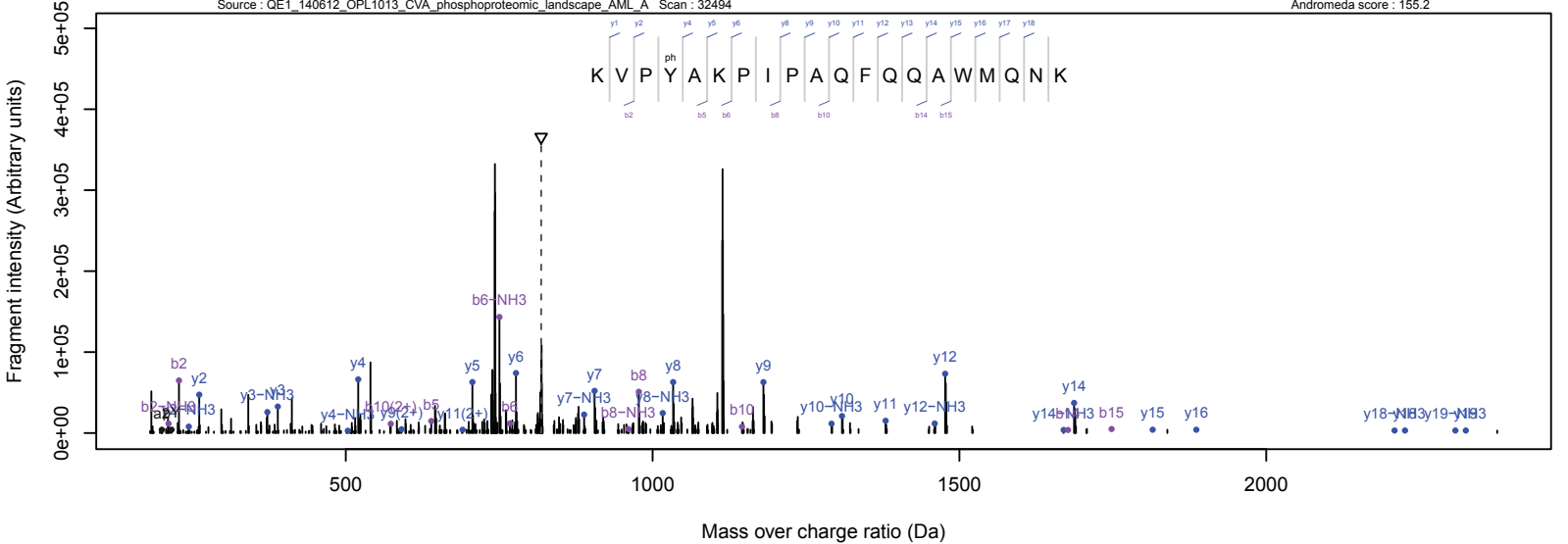
Andromeda score: 155.24



KVPY(ph)AKPIPAQFQQAWMQNK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 32494

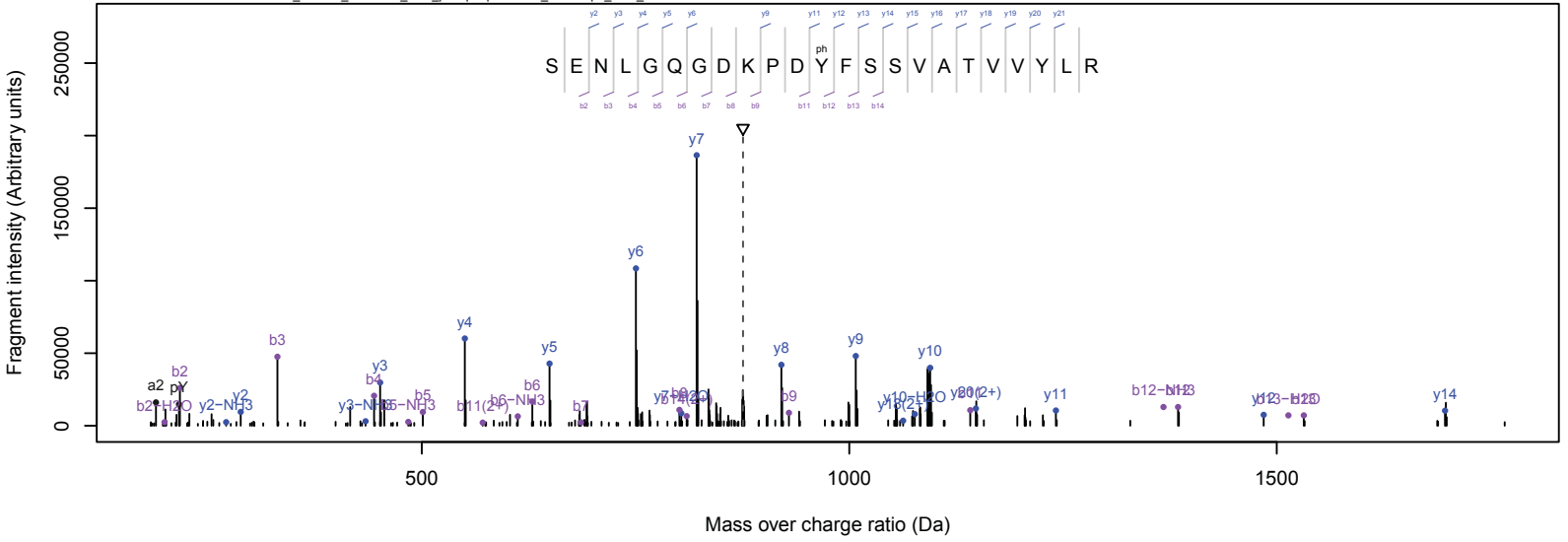
Andromeda score: 155.2



SENLGQGDKPDY(ph)FSSVATVVYLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41116

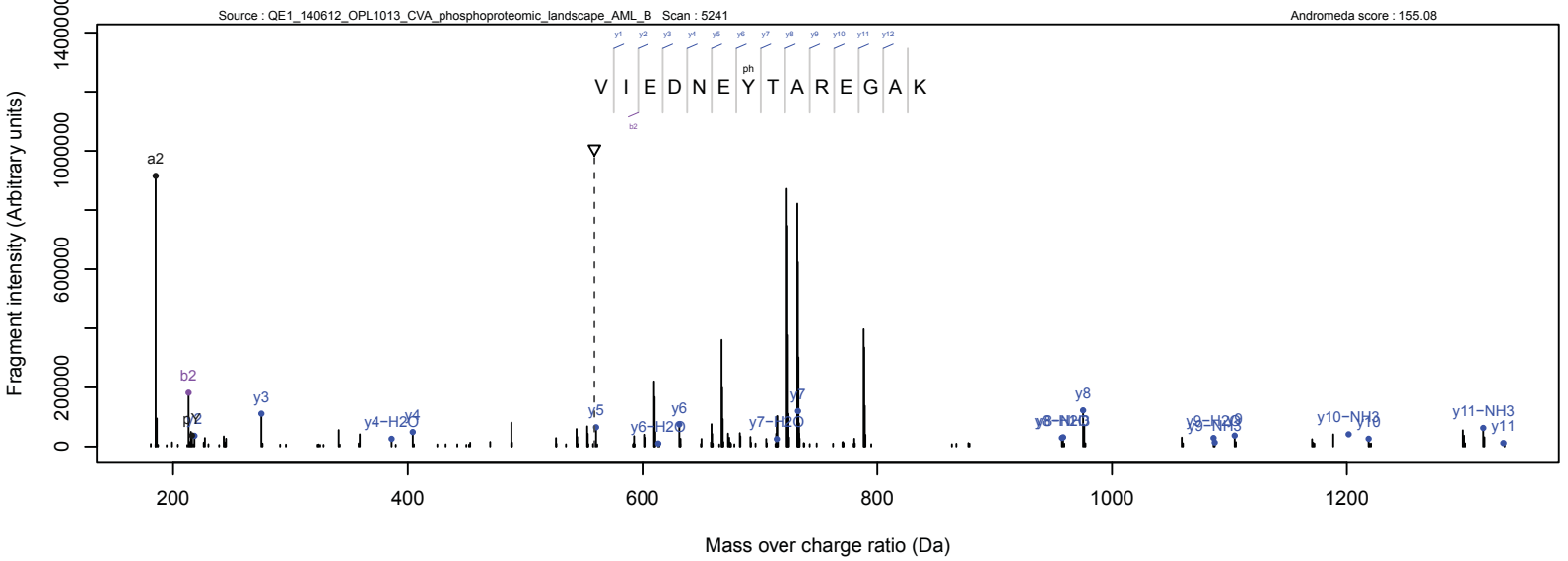
Andromeda score : 155.17



VIDNEY(ph)TAREGAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 5241

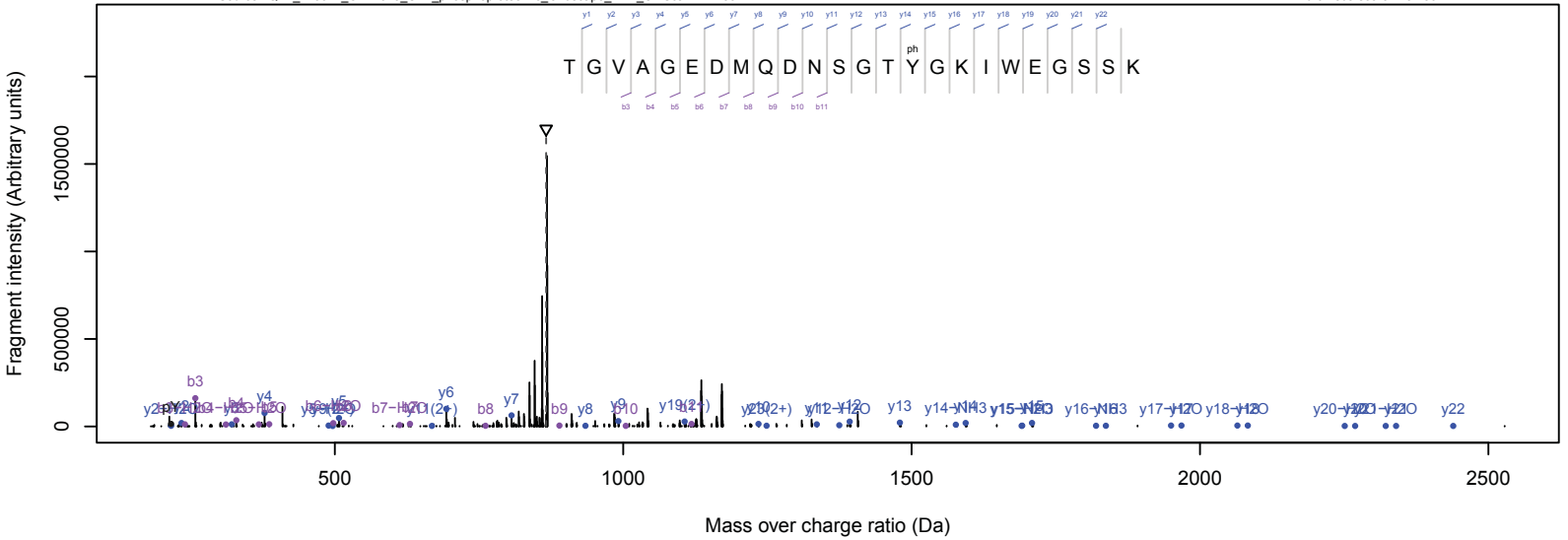
Andromeda score : 155.08



TGVAGEDMQDMSGTY(ph)GIWEGSSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22183

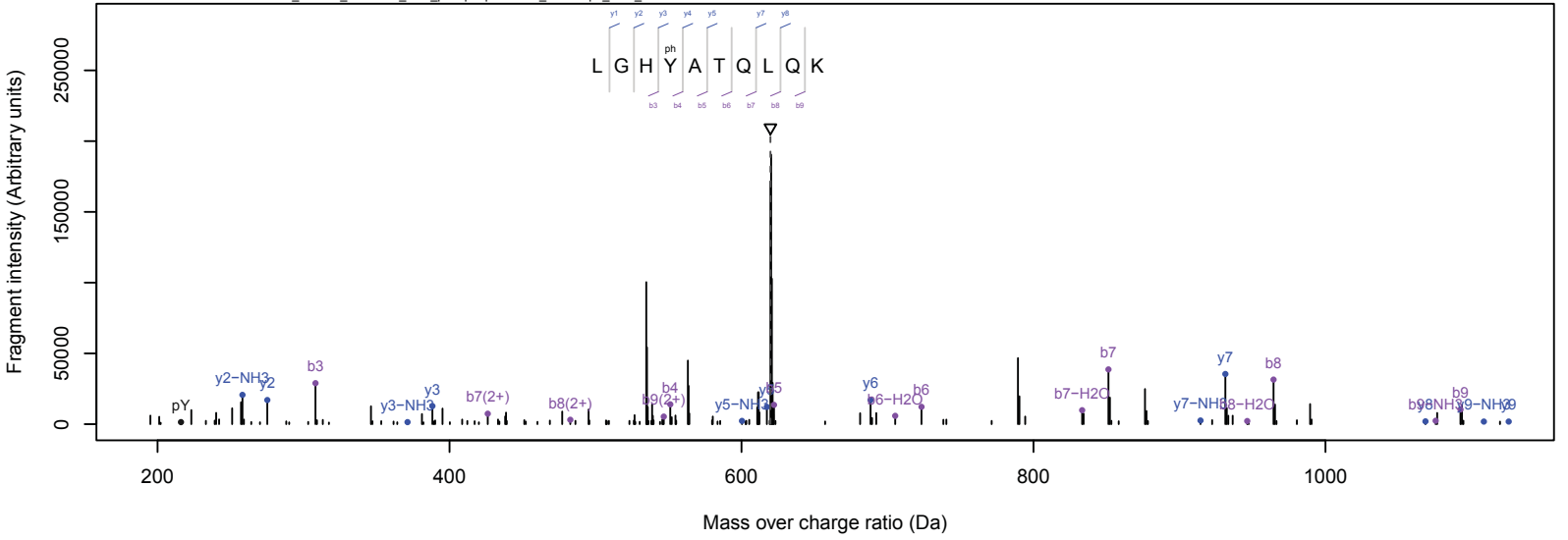
Andromeda score : 154.86



LGHY(ph)ATQLQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7378

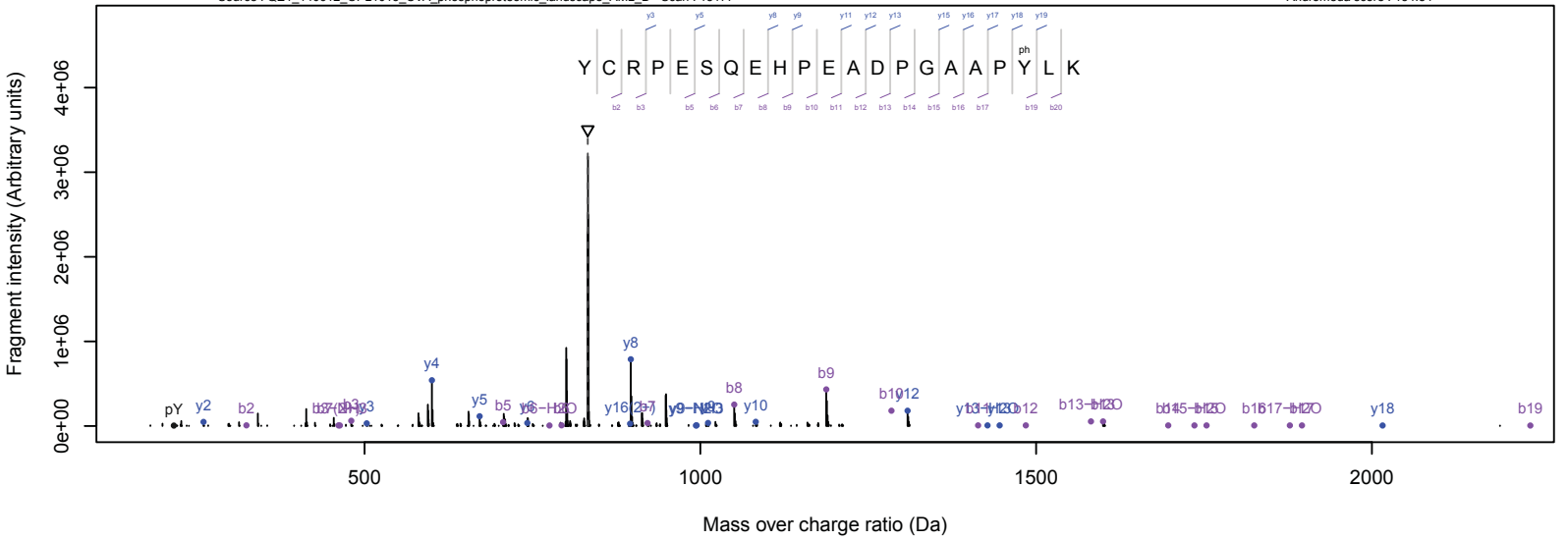
Andromeda score : 154.84



YCRPESQEHPEADPGAAPY(ph)LK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 15177

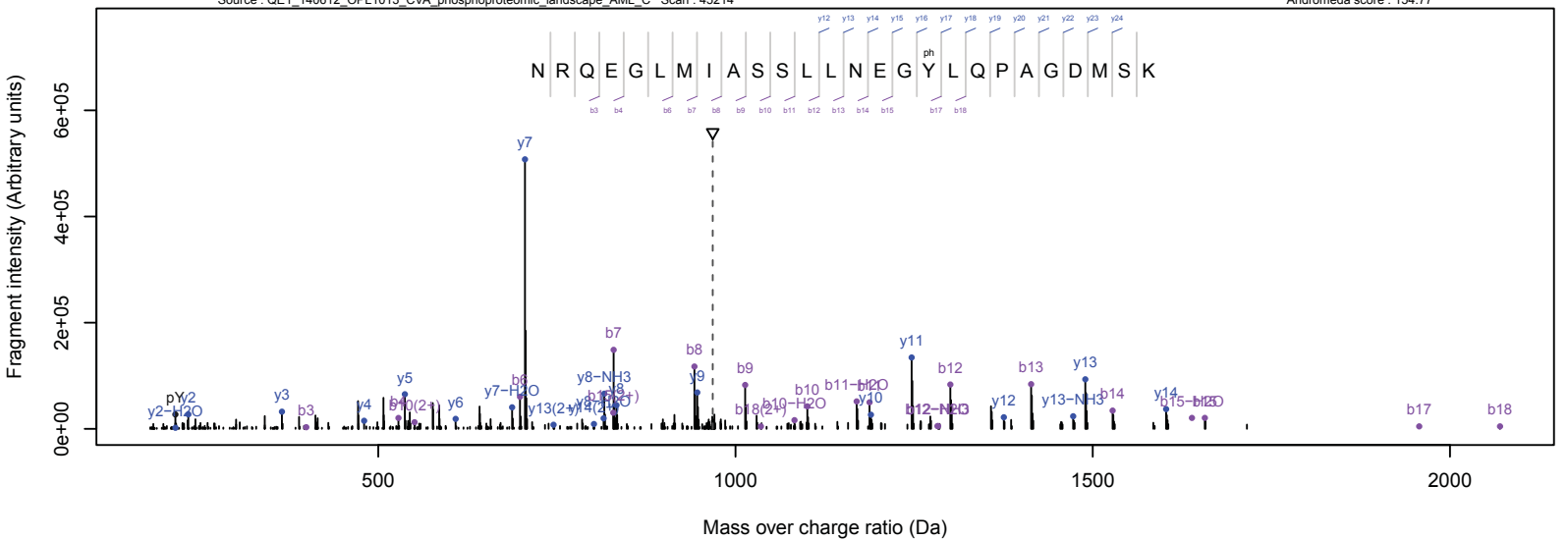
Andromeda score : 154.84



NRQEGLMIASSLLNEGY(ph)LQPAGDMSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45214

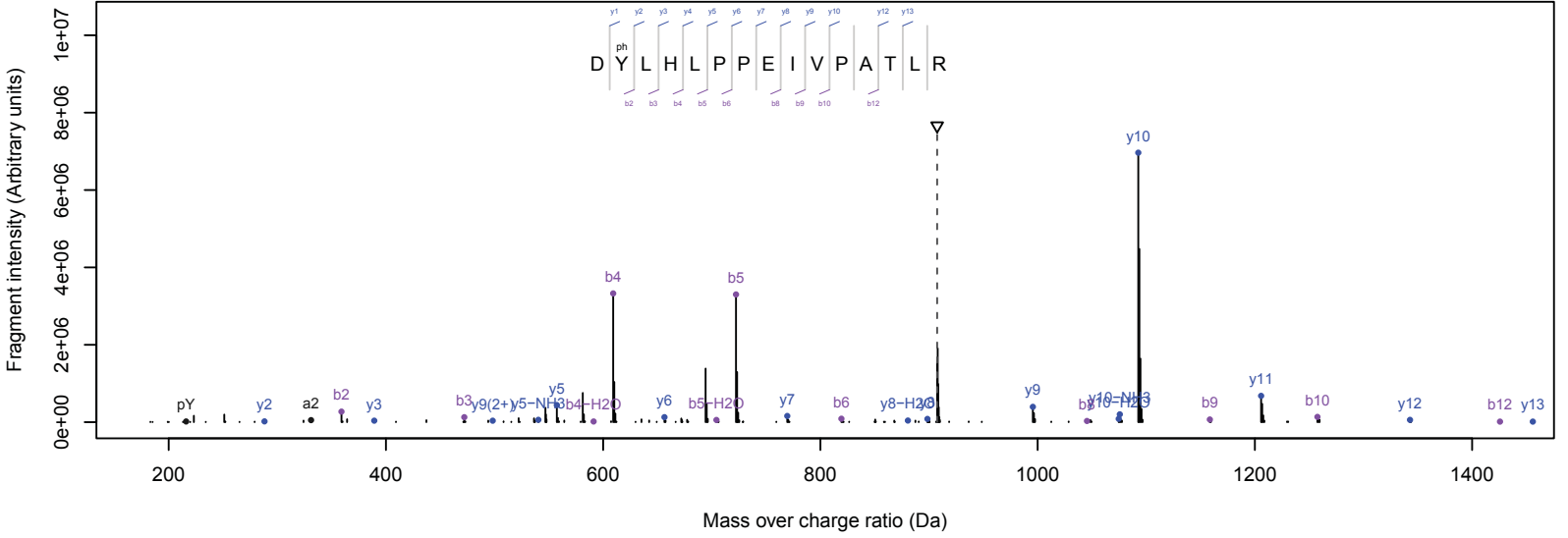
Andromeda score : 154.77



DY(ph)LHLPEIVPATLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45530

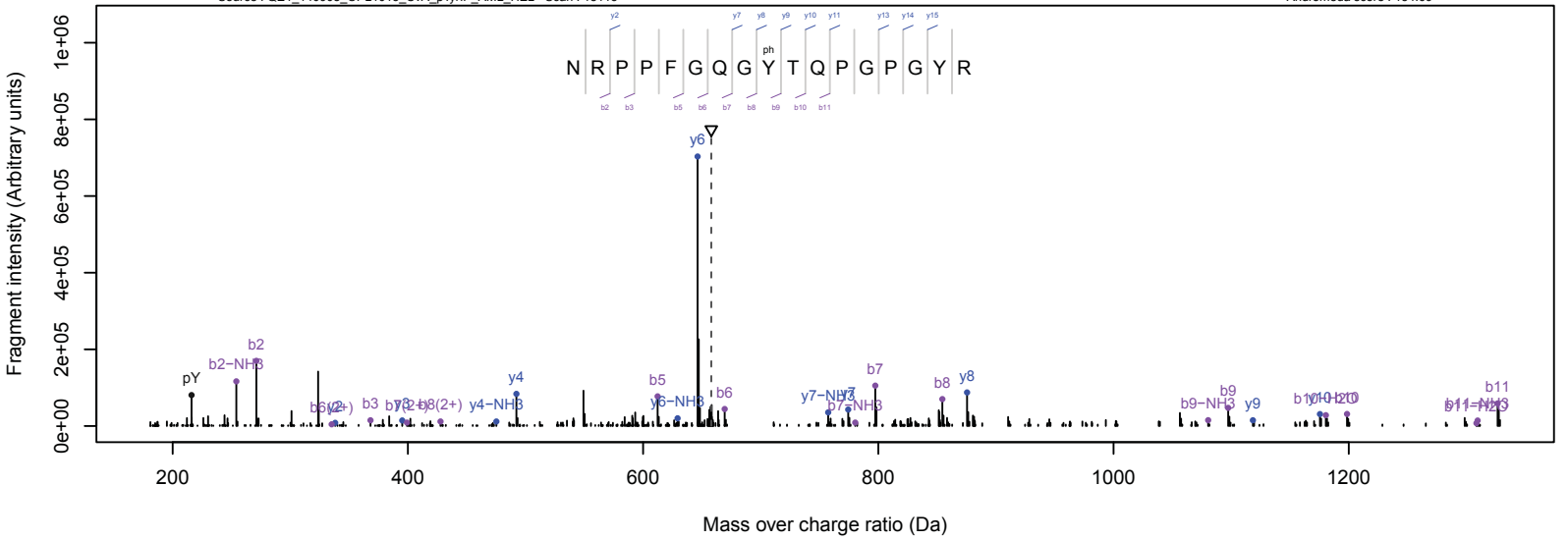
Andromeda score : 154.69



NRPPFGQGY(ph)TQPGPGYR-3+

Source : QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_HEL Scan : 18113

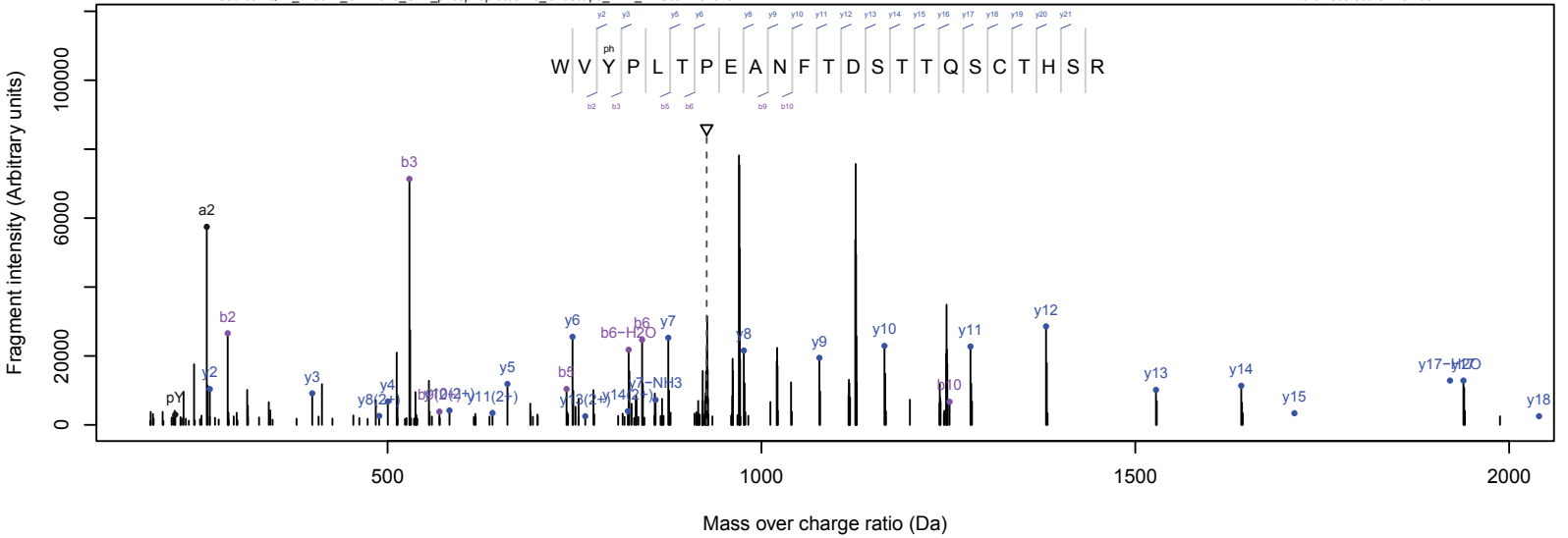
Andromeda score : 154.69



WVY(ph)PLTPEANFTDSTTQSCYHSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34328

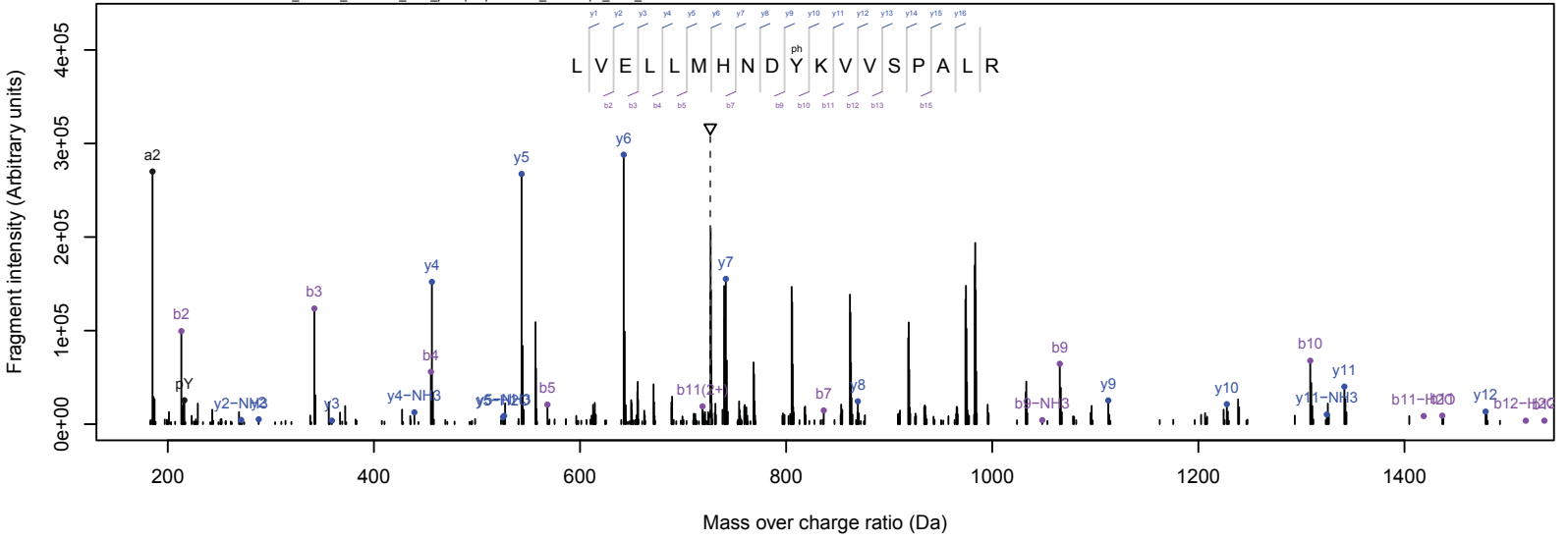
Andromeda score : 154.65



LVELLMHNDY(ph)KVVSPALR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36282

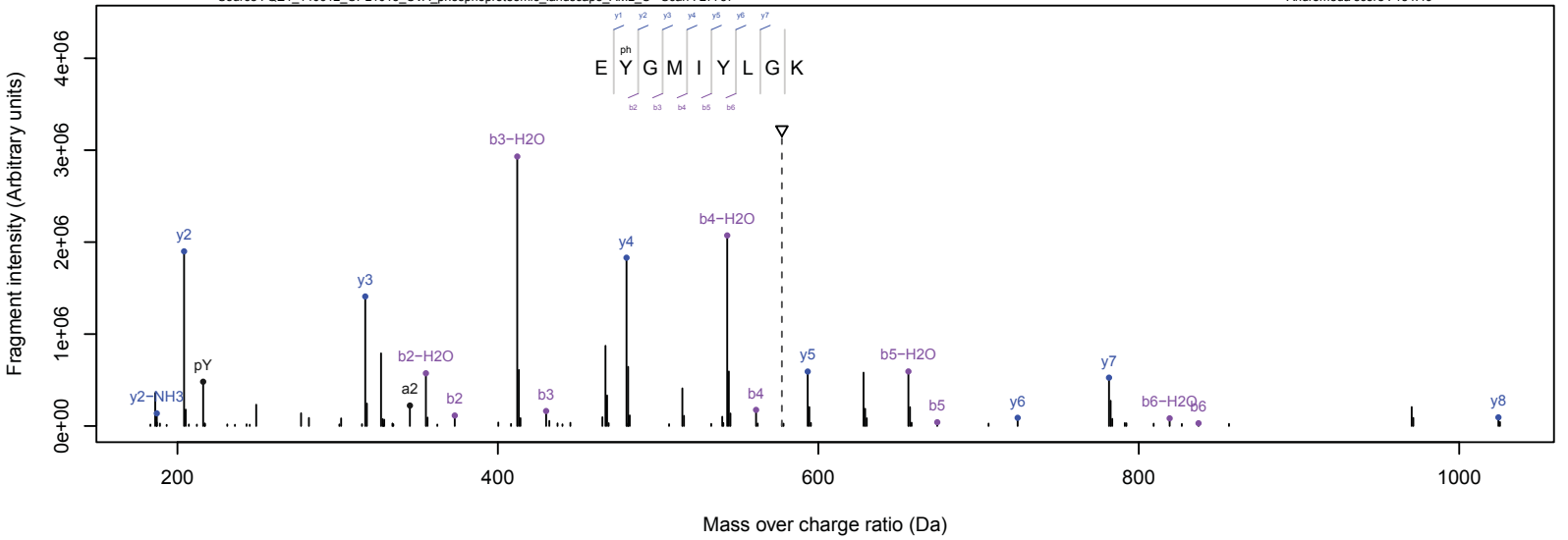
Andromeda score : 154.55



EY(ph)GMIYLGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27757

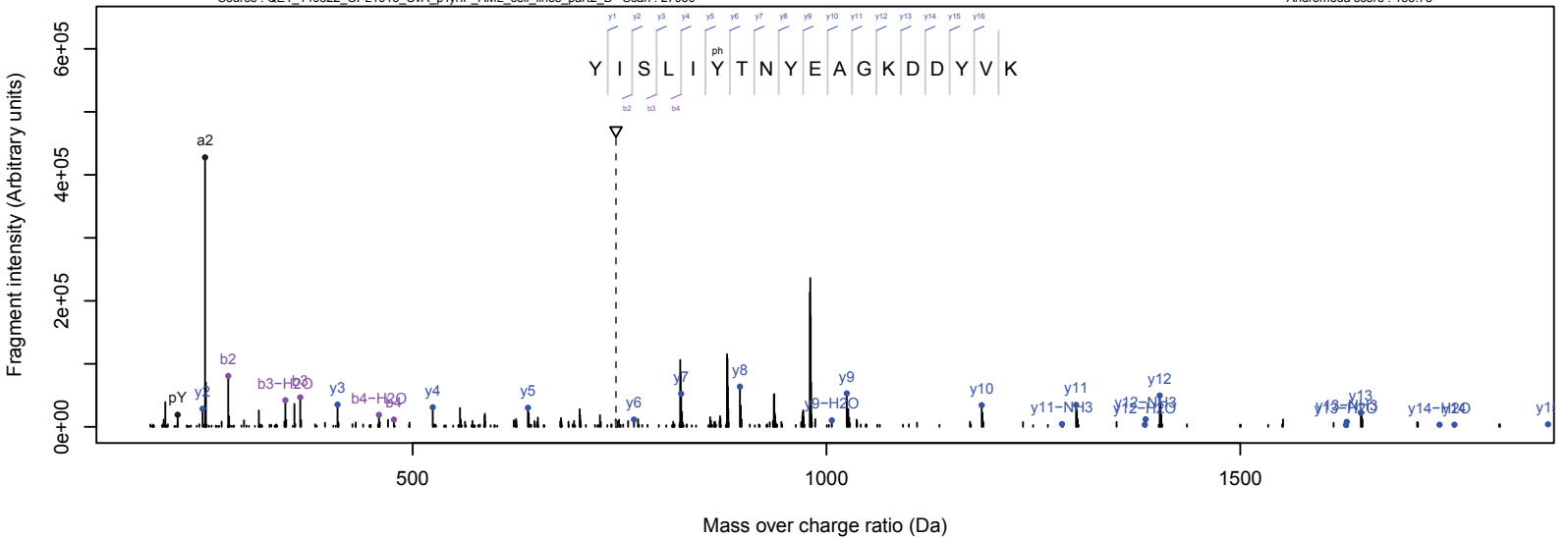
Andromeda score : 154.49



YISLIY(ph)TNYEAGKDDYVK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 27050

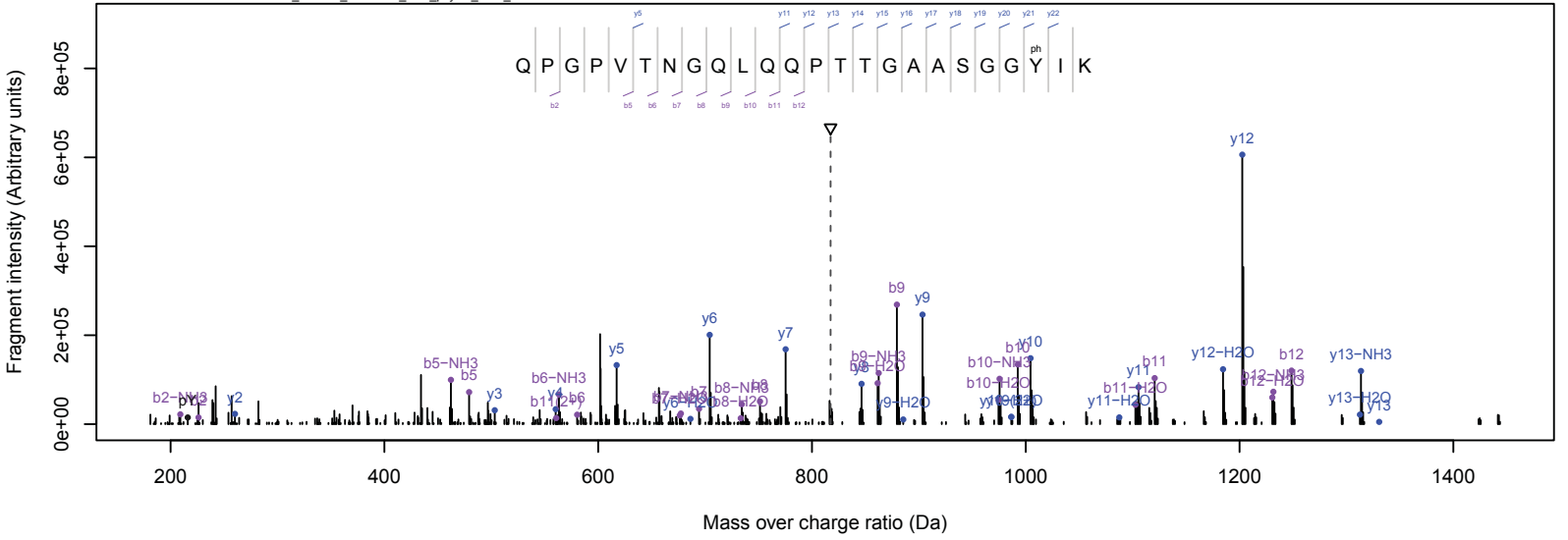
Andromeda score : 153.75



**QPGPVTNGQLQQPTTGAASGGY(ph)K-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 22526

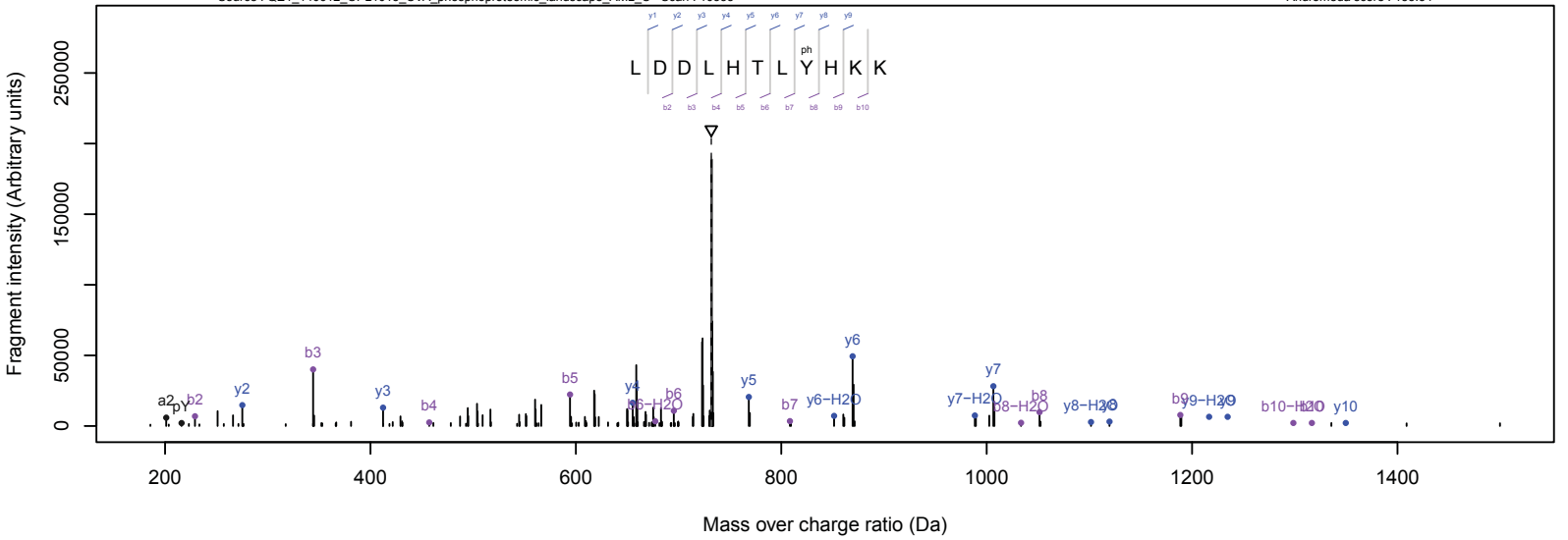
Andromeda score : 153.65



**LDDLHTLY(ph)HKK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10333

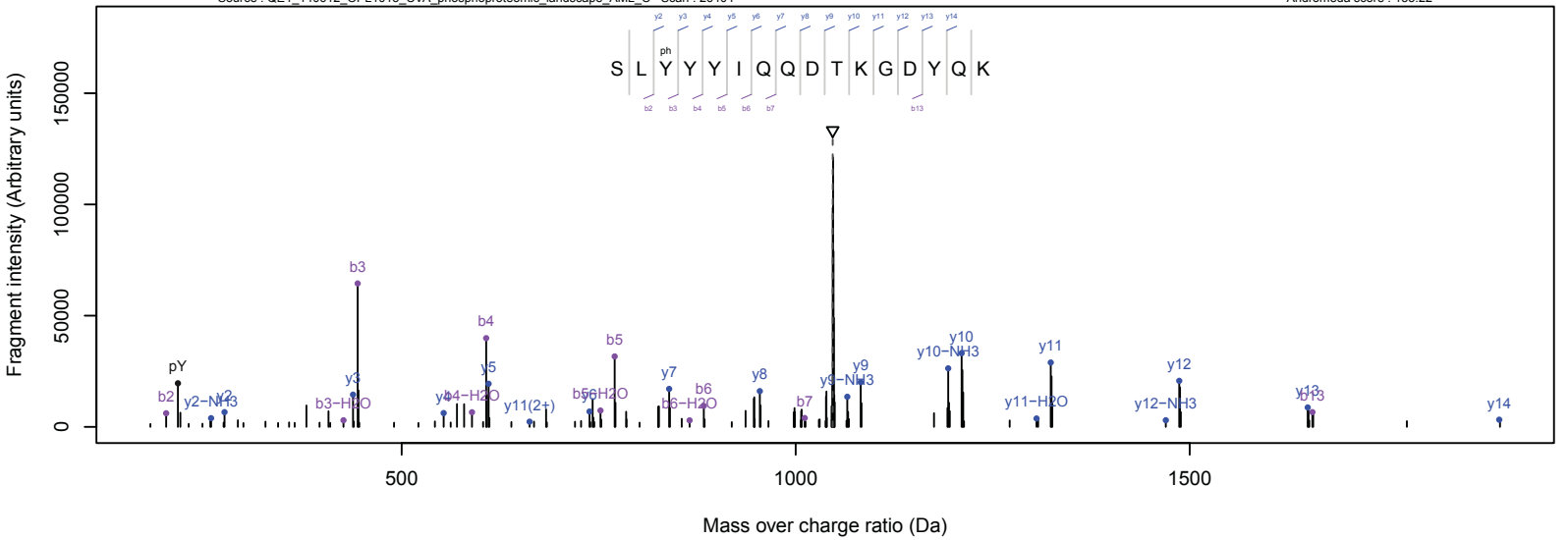
Andromeda score : 153.34



**SLY(ph)YYIQDQTKGDYQK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26404

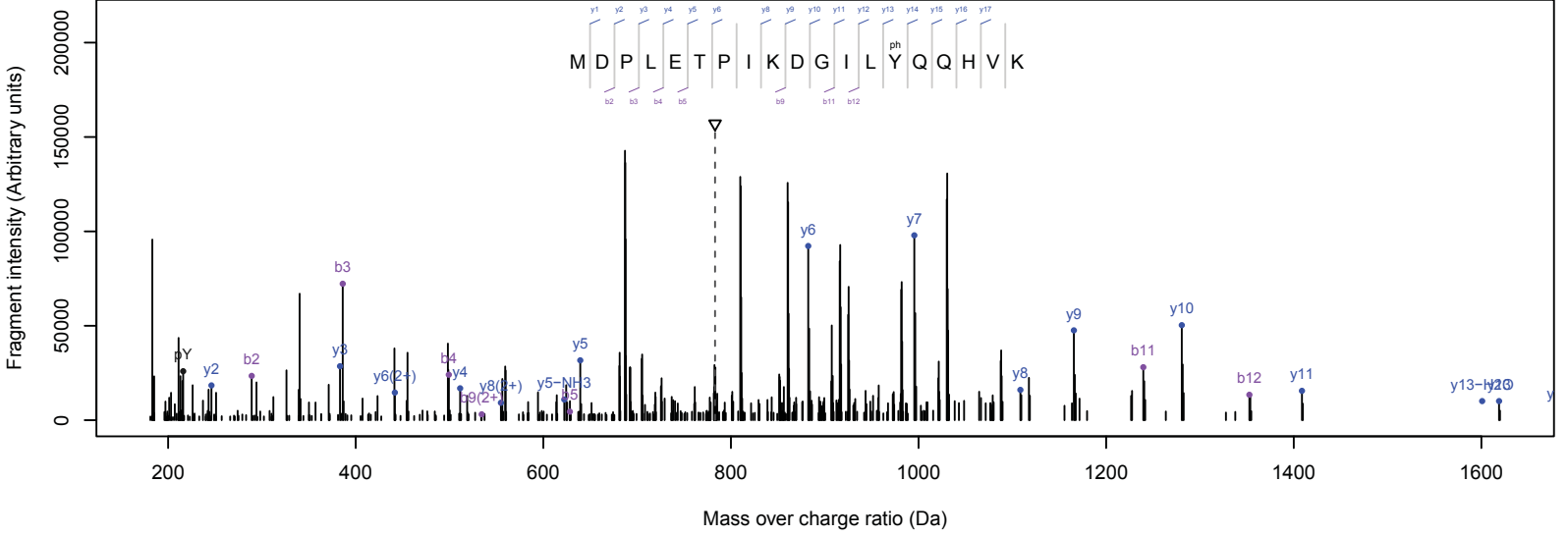
Andromeda score : 153.22



(ac)MDPLETPIKDGILLYQGHVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42873

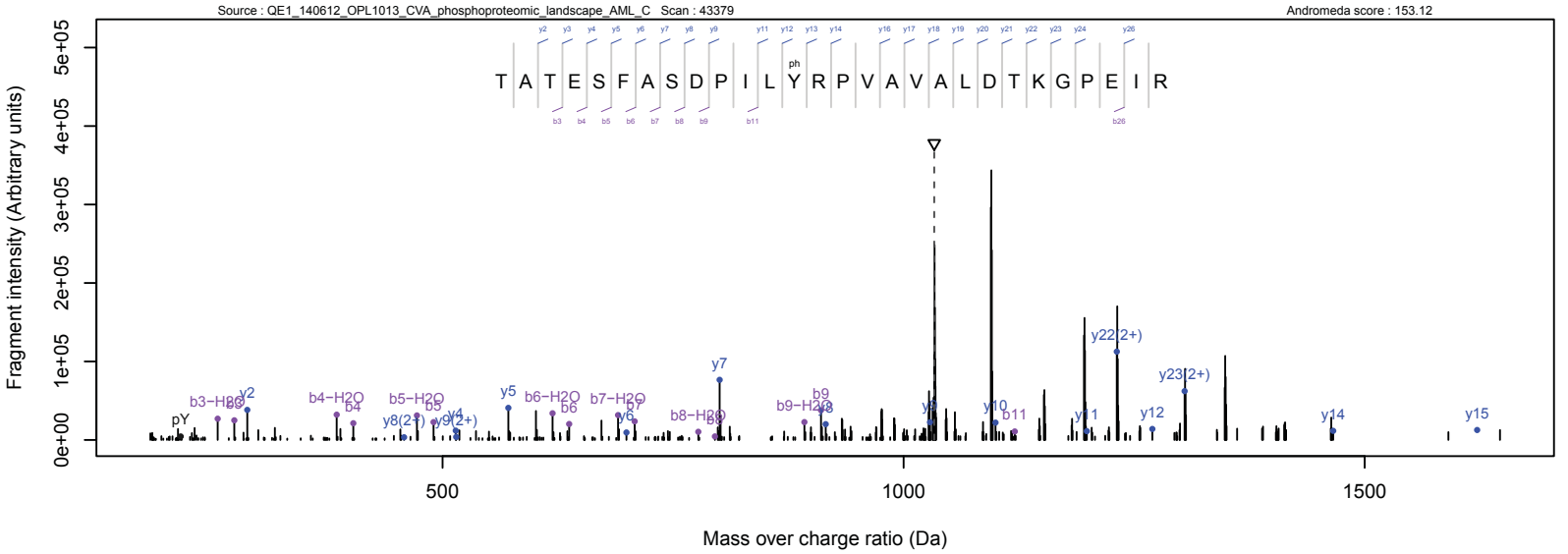
Andromeda score : 153.21



TATESFASDPILYRVPVAVALDTKGPEIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43379

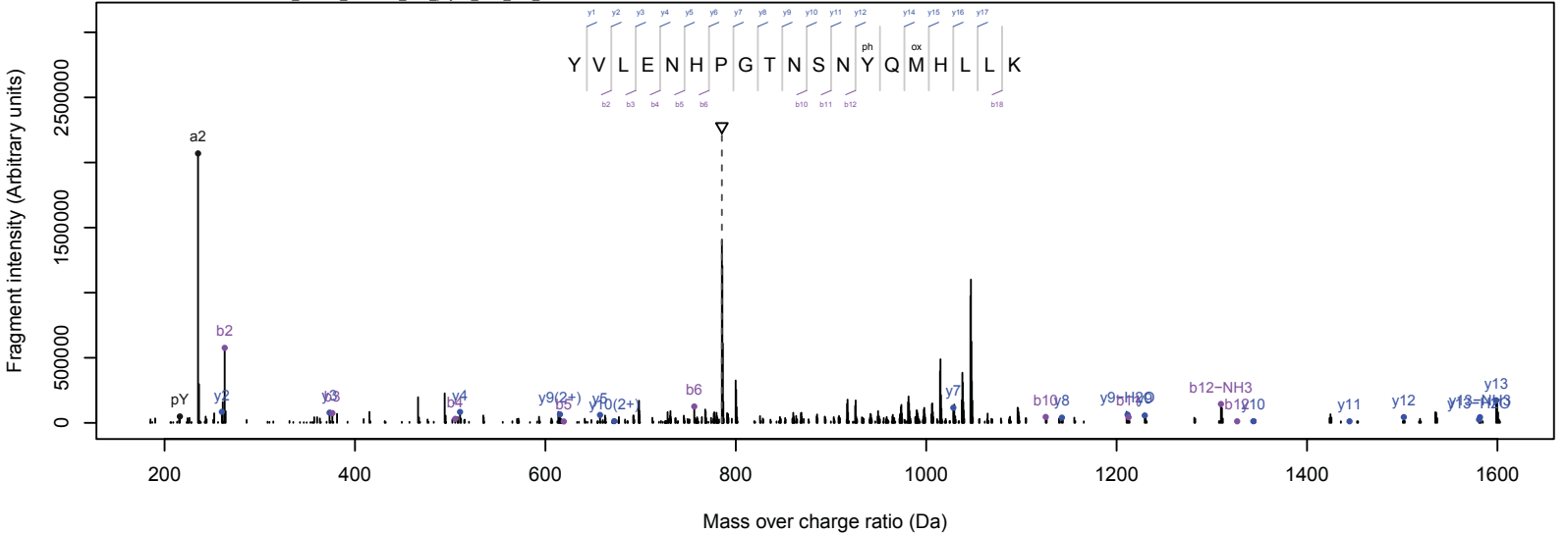
Andromeda score : 153.12



YVLEHHPGTNSNYQM(ox)HLLK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 20424

Andromeda score : 153.04

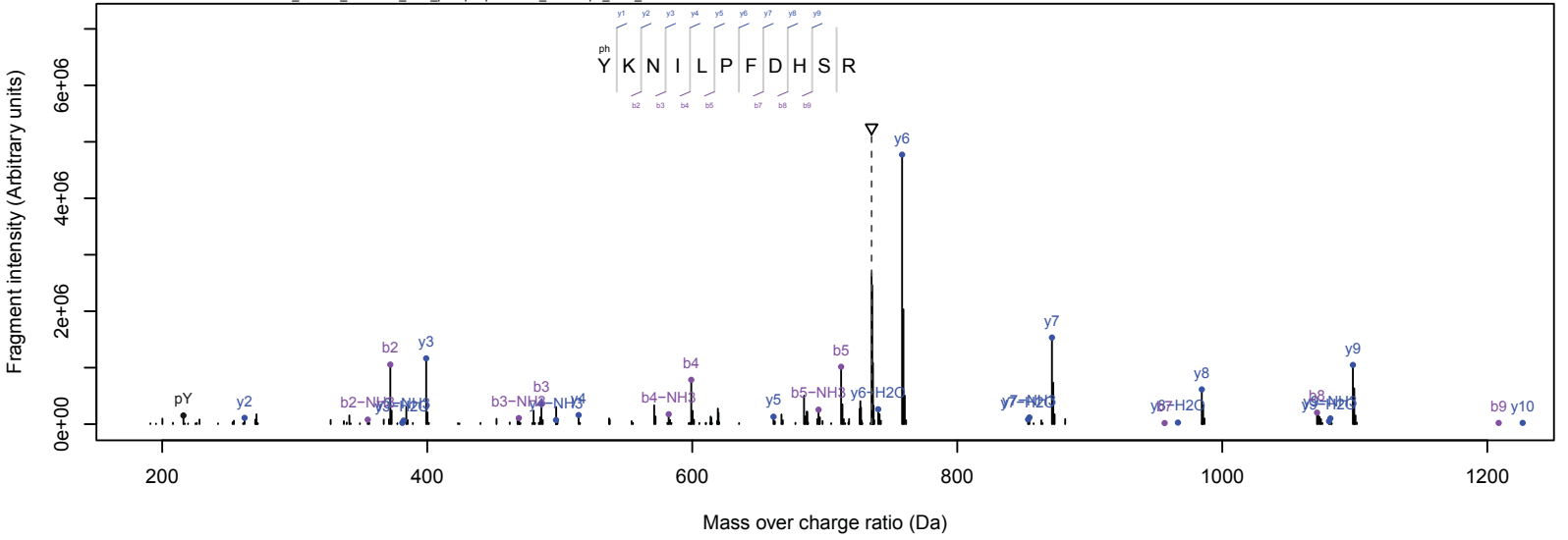




Y(ph)KNILPFDHSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20552

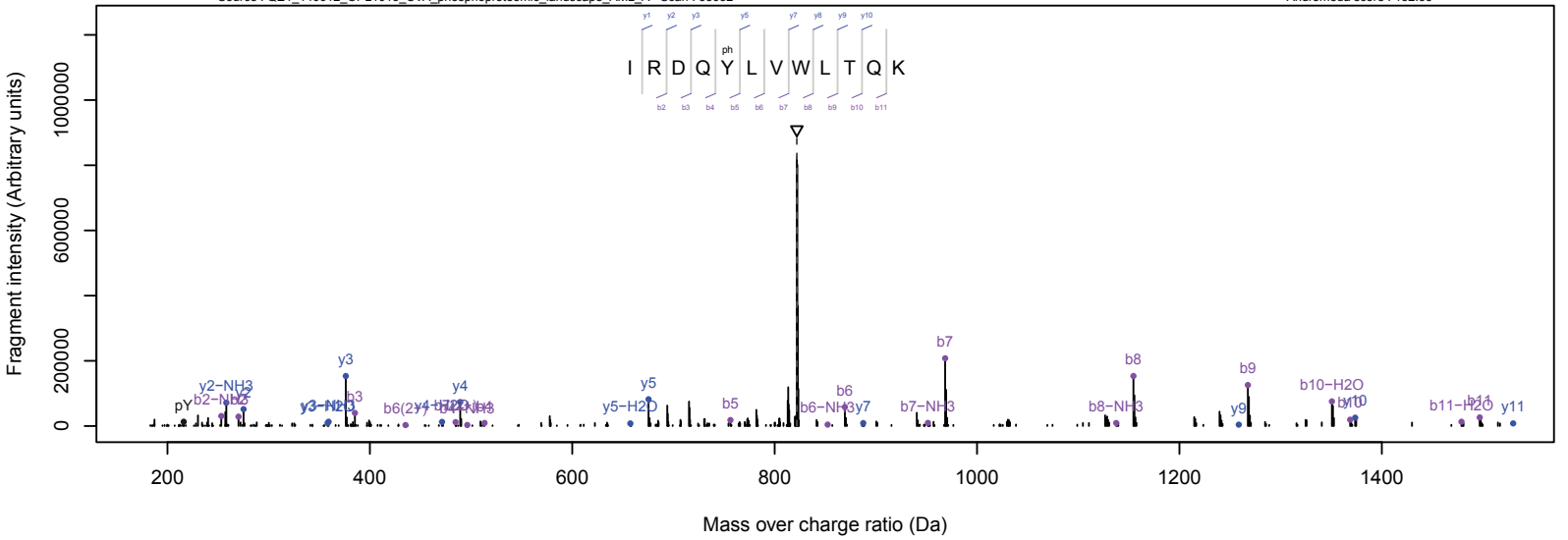
Andromeda score : 152.96



IRDQY(ph)LVWLTQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 35032

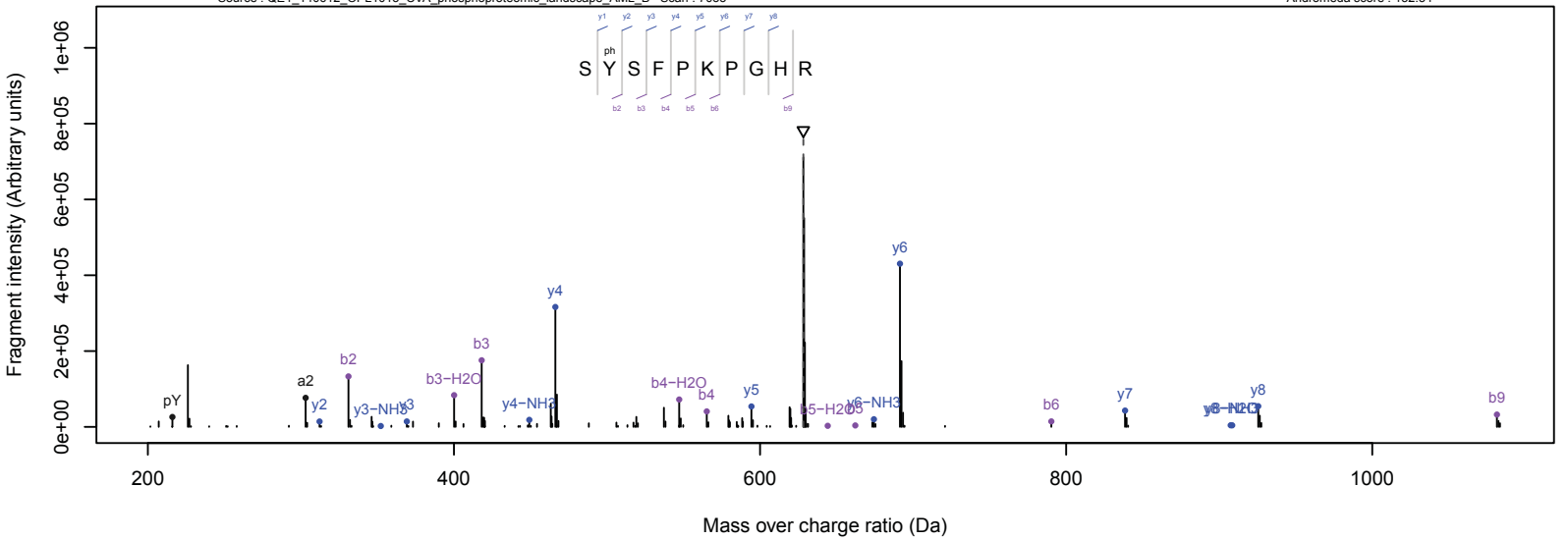
Andromeda score : 152.55



SY(ph)SFPKPGHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7055

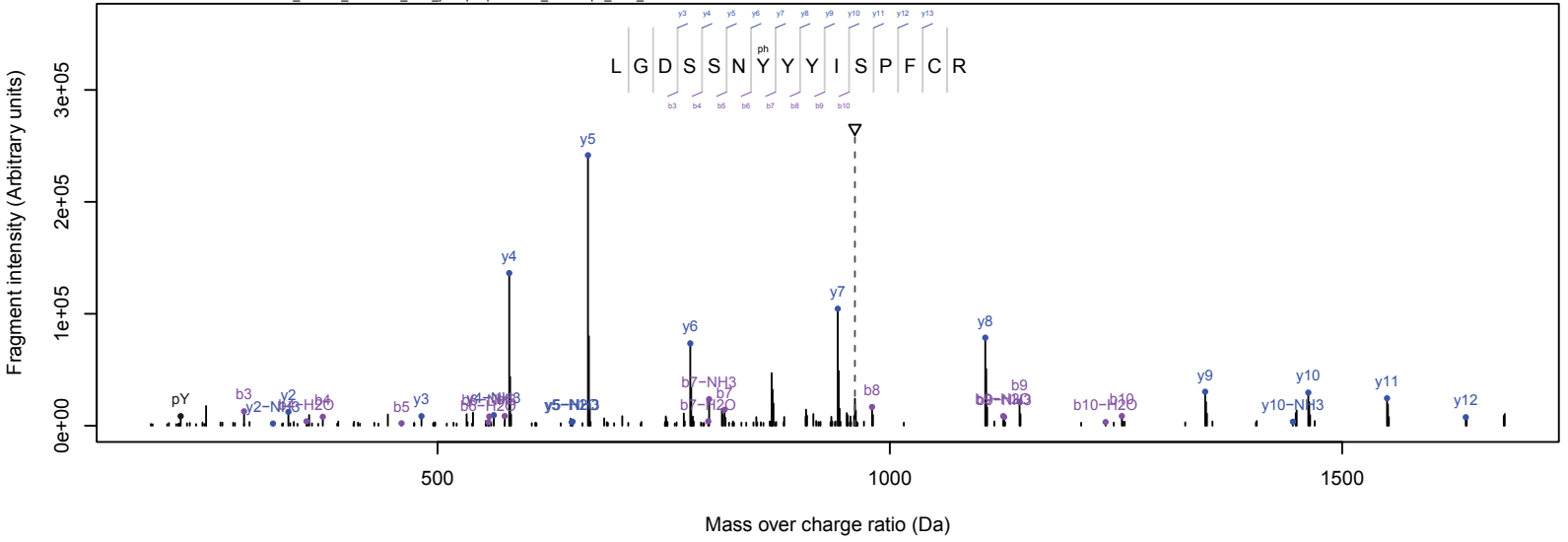
Andromeda score : 152.54



LGSSNY(ph)YYISPF<sup>CR</sup>-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37535

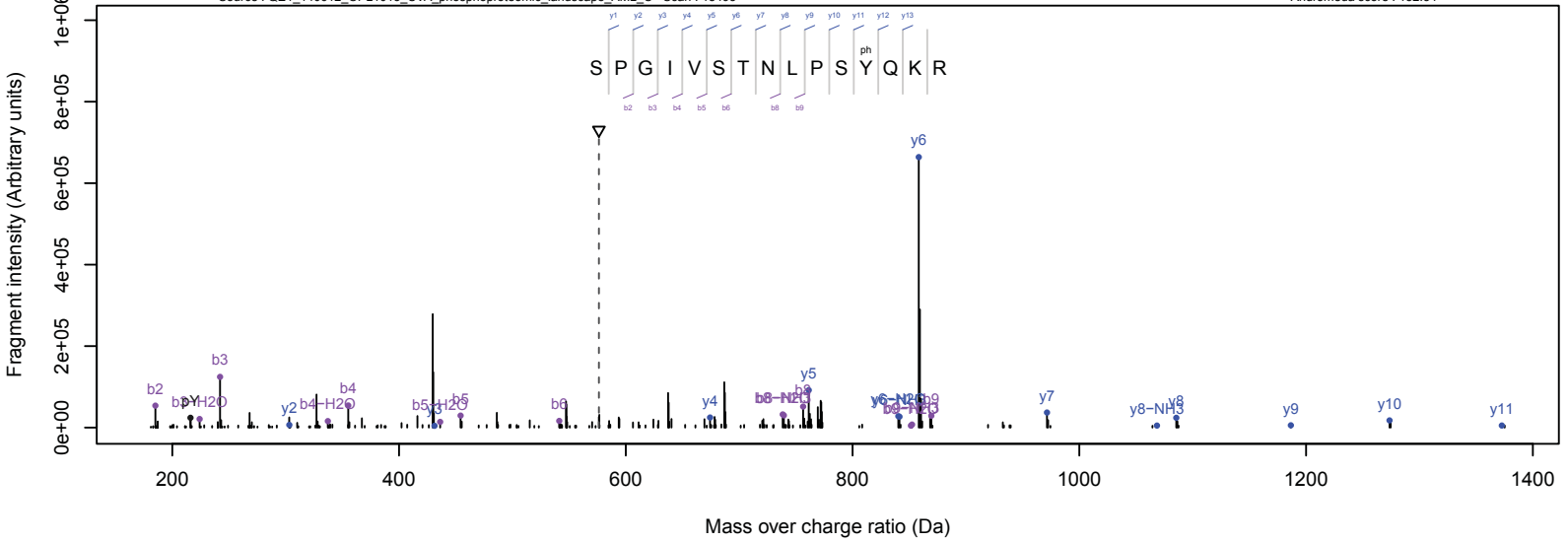
Andromeda score : 152.34



SPGIVSTNLPSY(ph)QKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18493

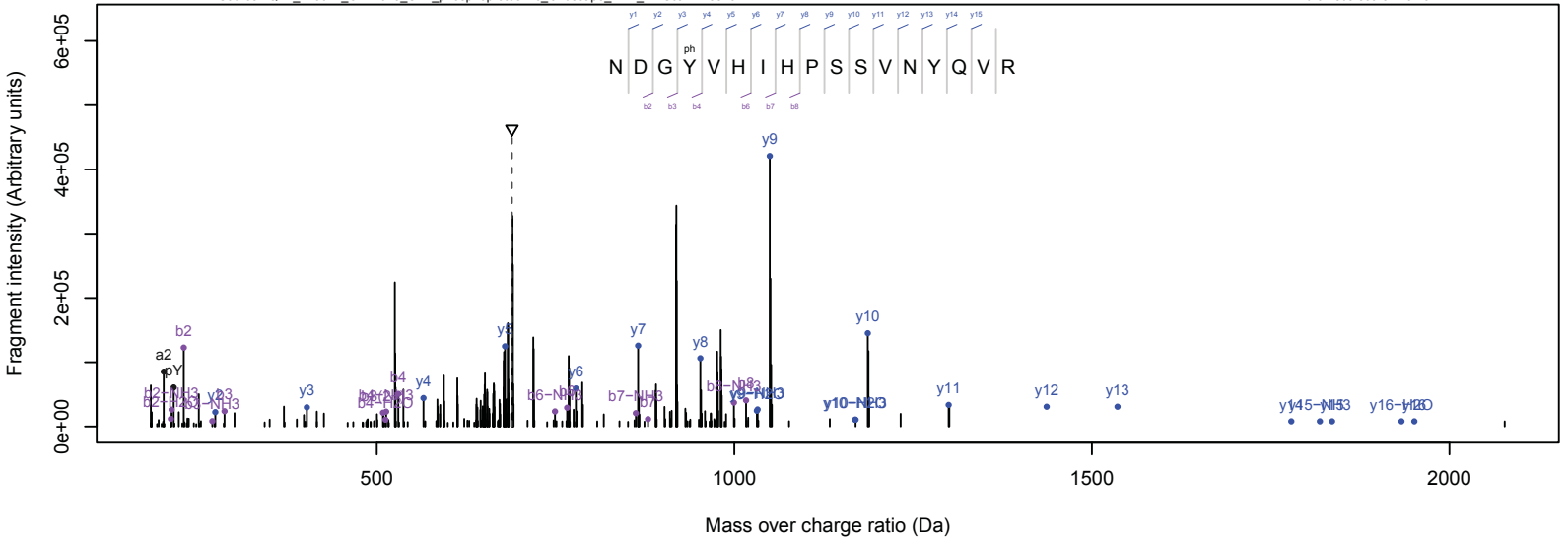
Andromeda score : 152.34



NDGY(ph)VHIHPSSVNYQVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 18813

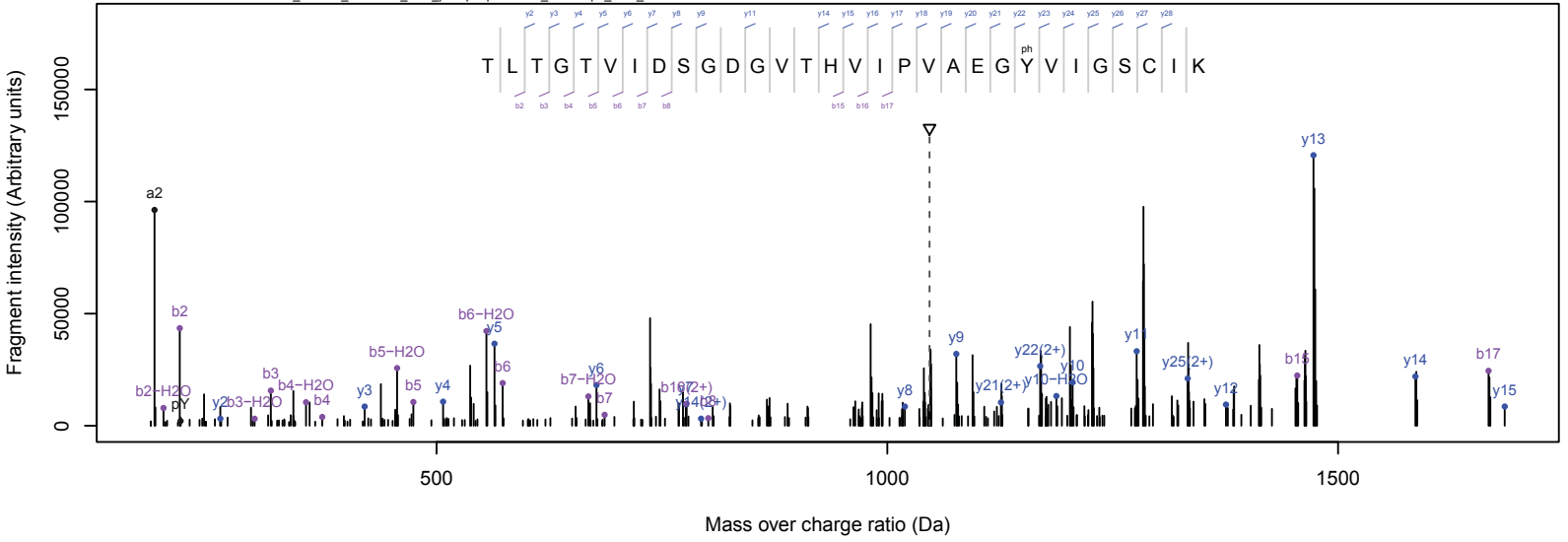
Andromeda score : 152.32



TLTGTVIDSGDGVTHVIPVAEGY(ph)VIGSCIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 46476

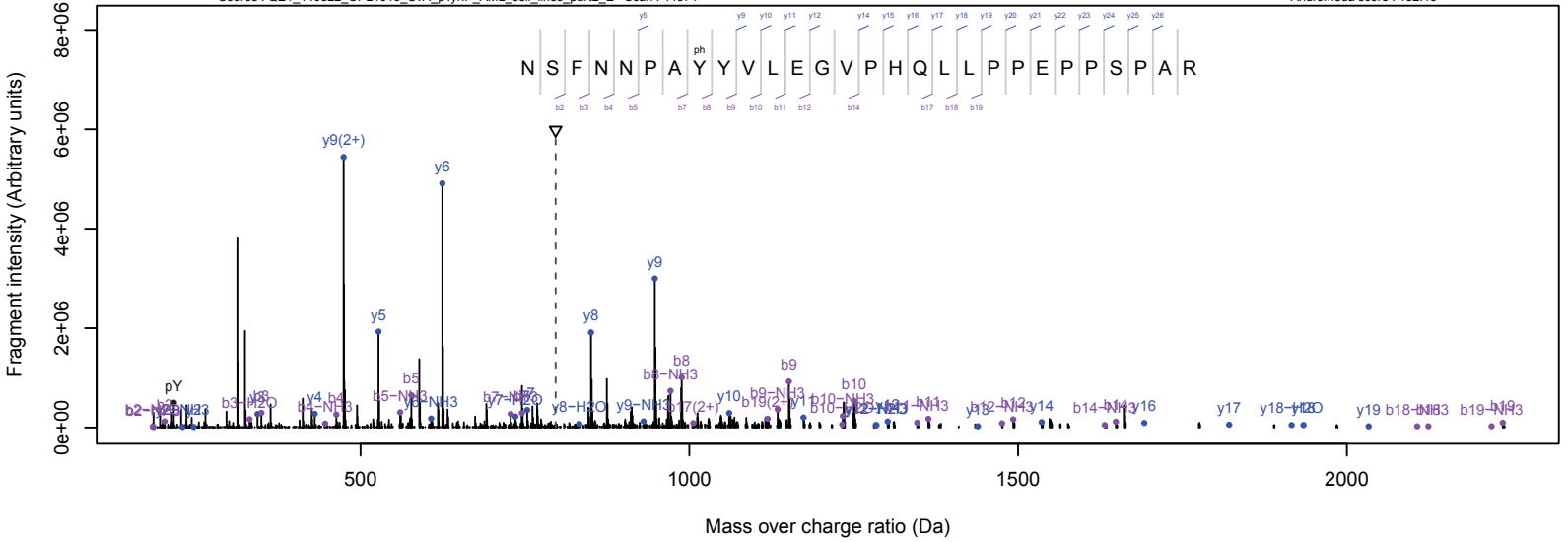
Andromeda score : 152.16



NSFNNPAY(ph)YVLEGVPHQLLPPEPPSPAR-4+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 44374

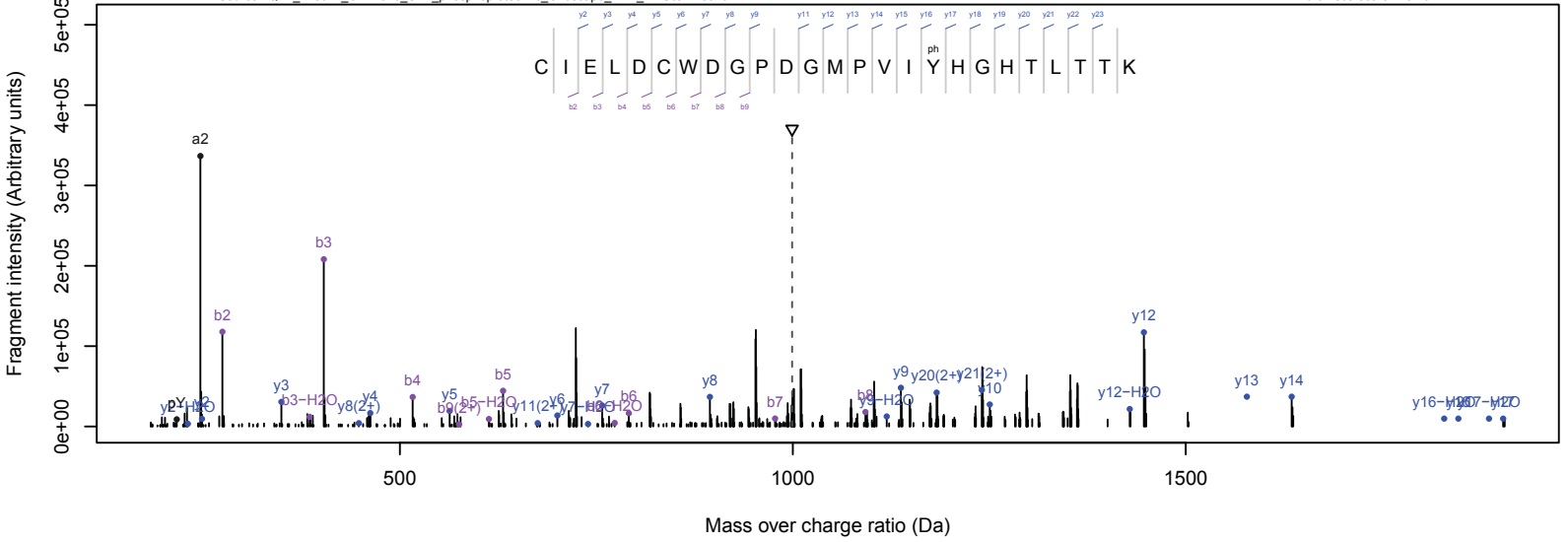
Andromeda score : 152.15



CIELDCWDGPDGMPVIYHGHTLTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39294

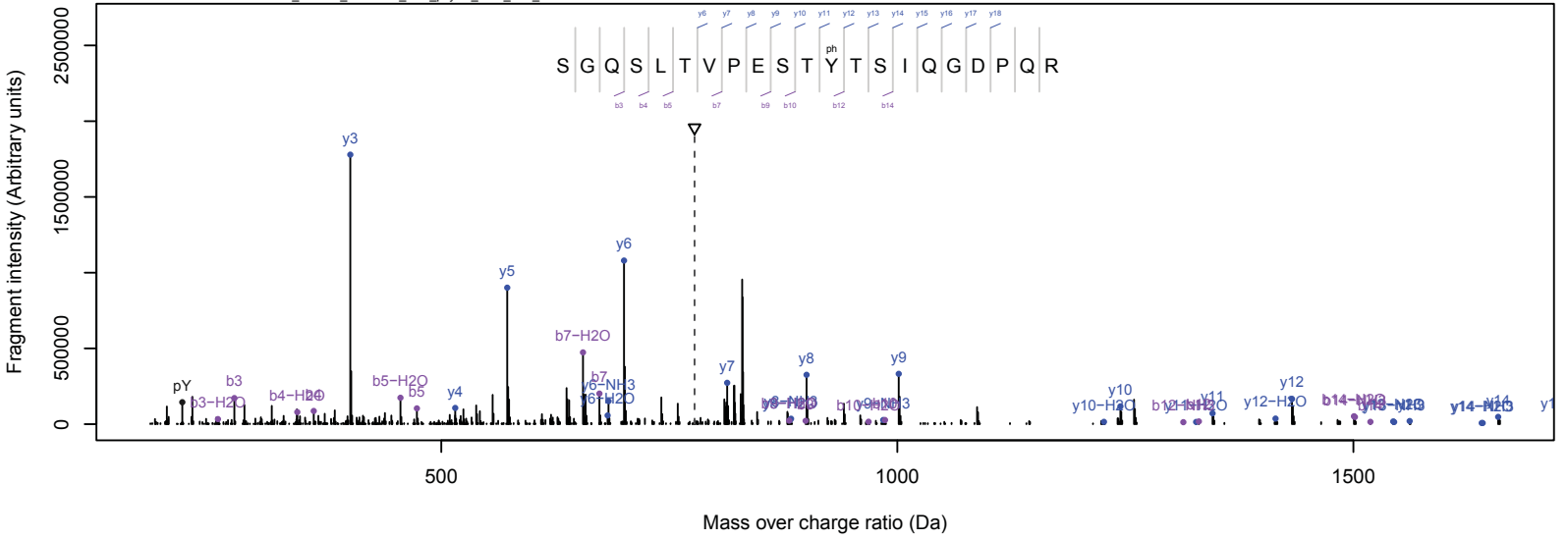
Andromeda score : 152.07



**SGQSLTVPES<sup>ph</sup>TYTSIQGDPQR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 25788

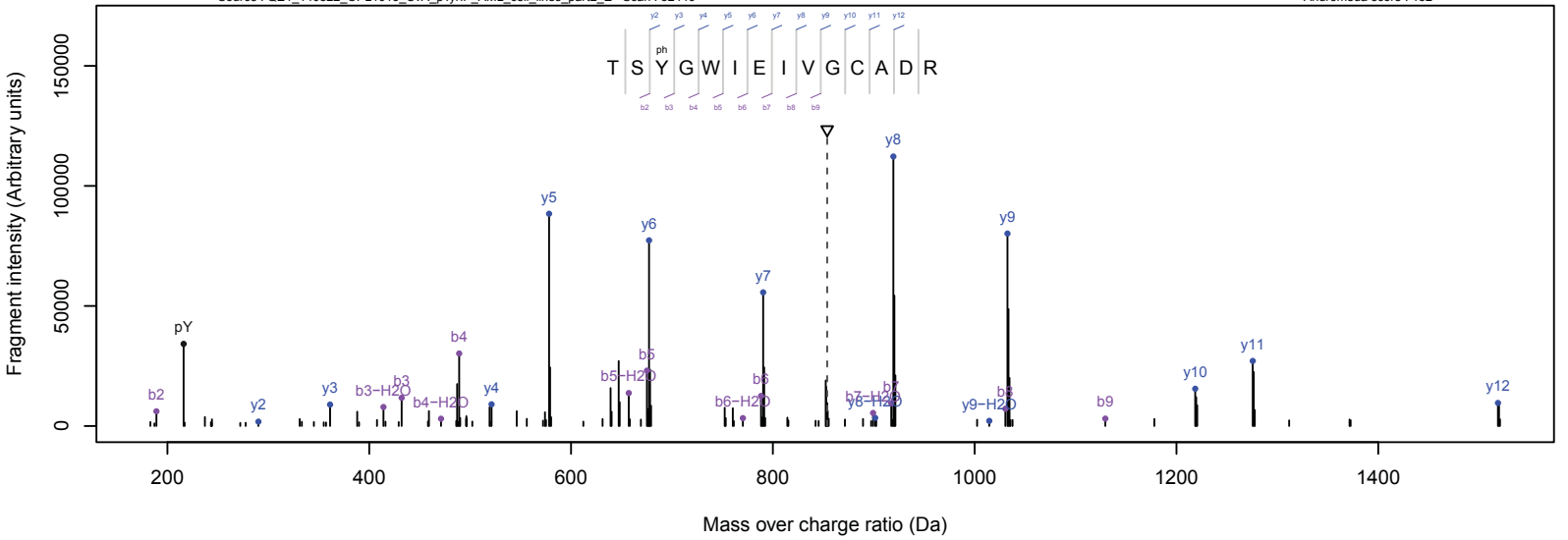
Andromeda score : 152.06



**TSY<sup>ph</sup>GWIEIVGCADR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 32446

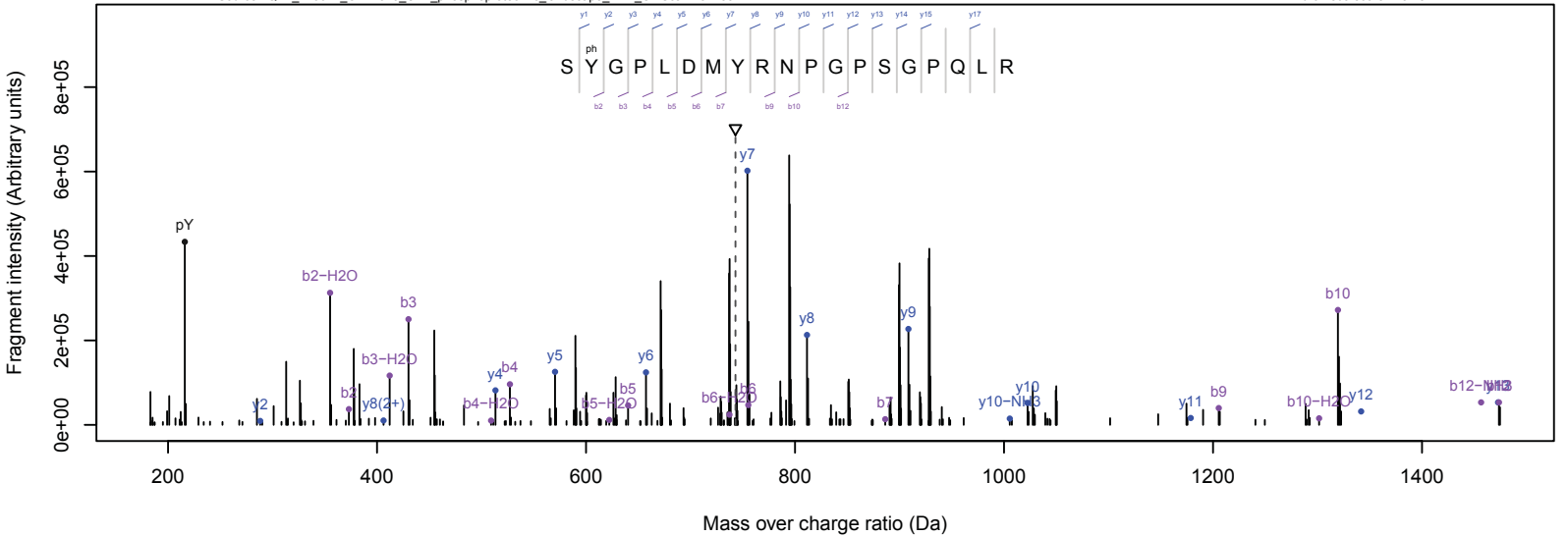
Andromeda score : 152



**(ac)SY<sup>ph</sup>GPLDMYRNPGPSGLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37185

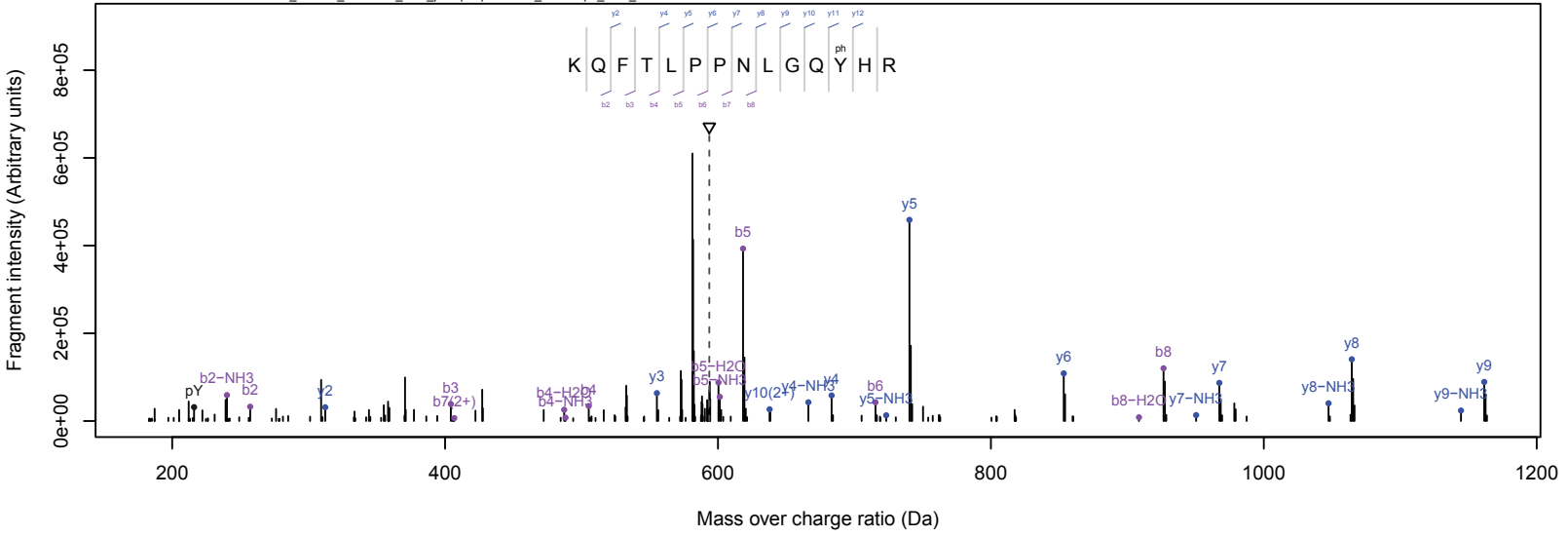
Andromeda score : 151.87



**KQFTLPPN LGQYHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23073

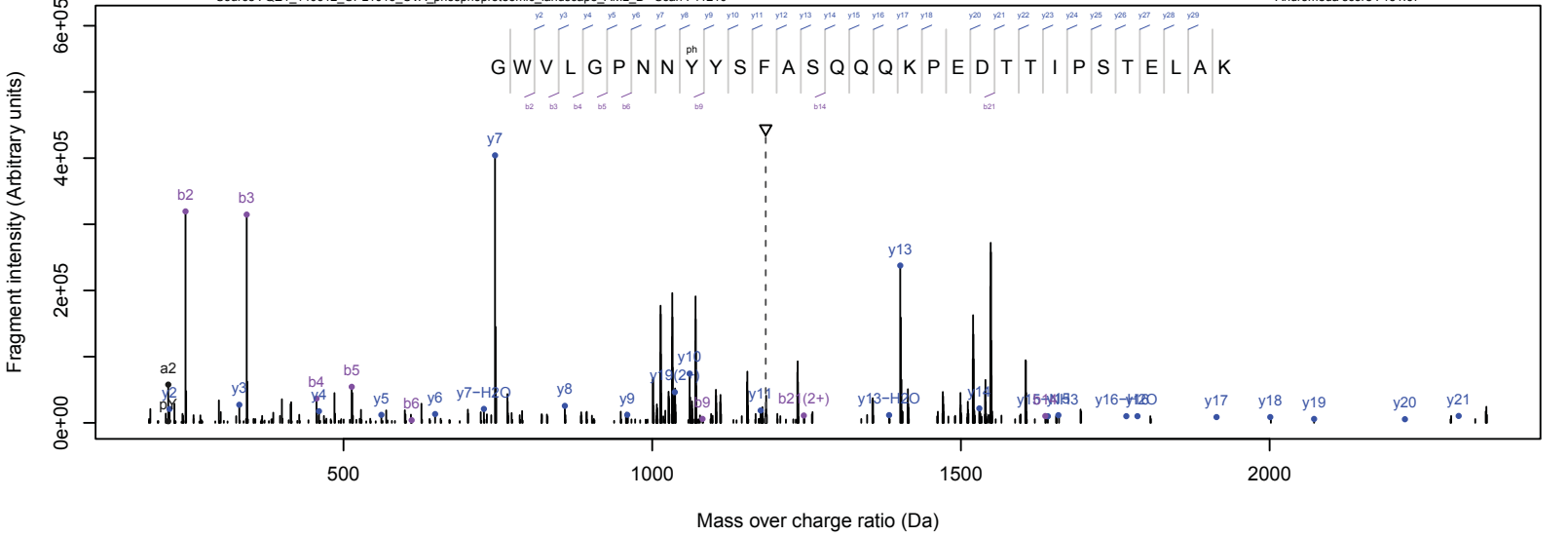
Andromeda score : 151.7



**GWV LGPN NY YSFASQQQKPEDTTIPSTELAK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41210

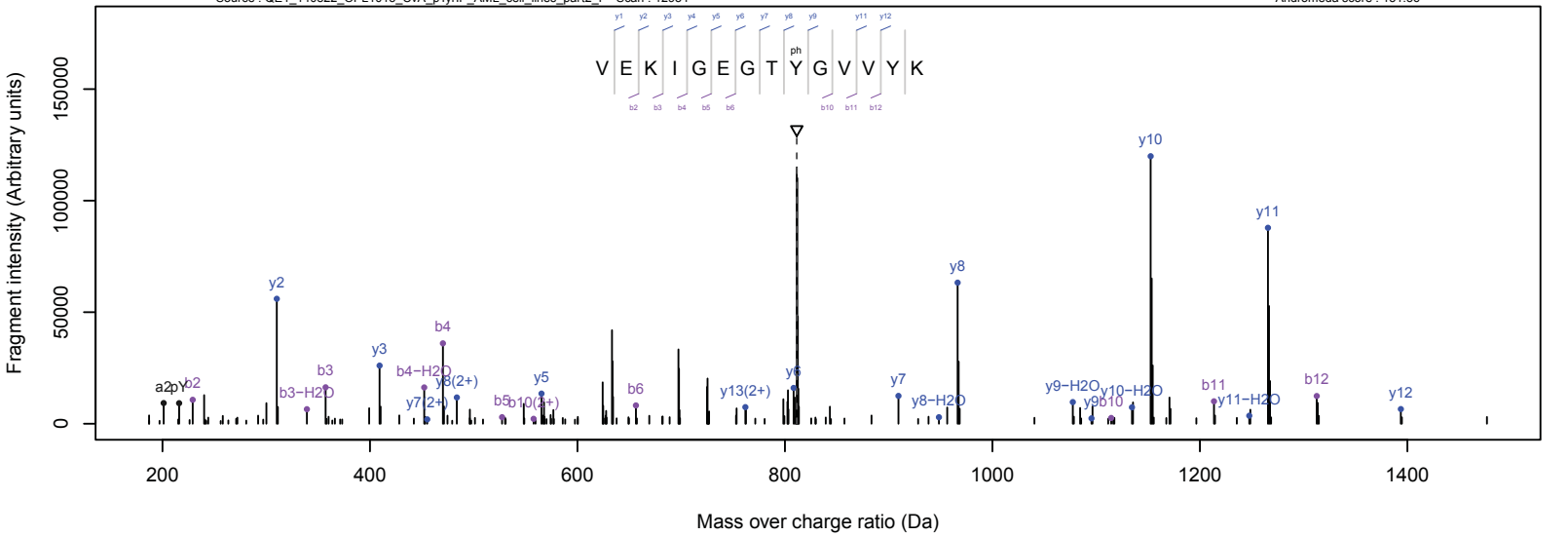
Andromeda score : 151.67



**VEKIGEGTYGVVYK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 12934

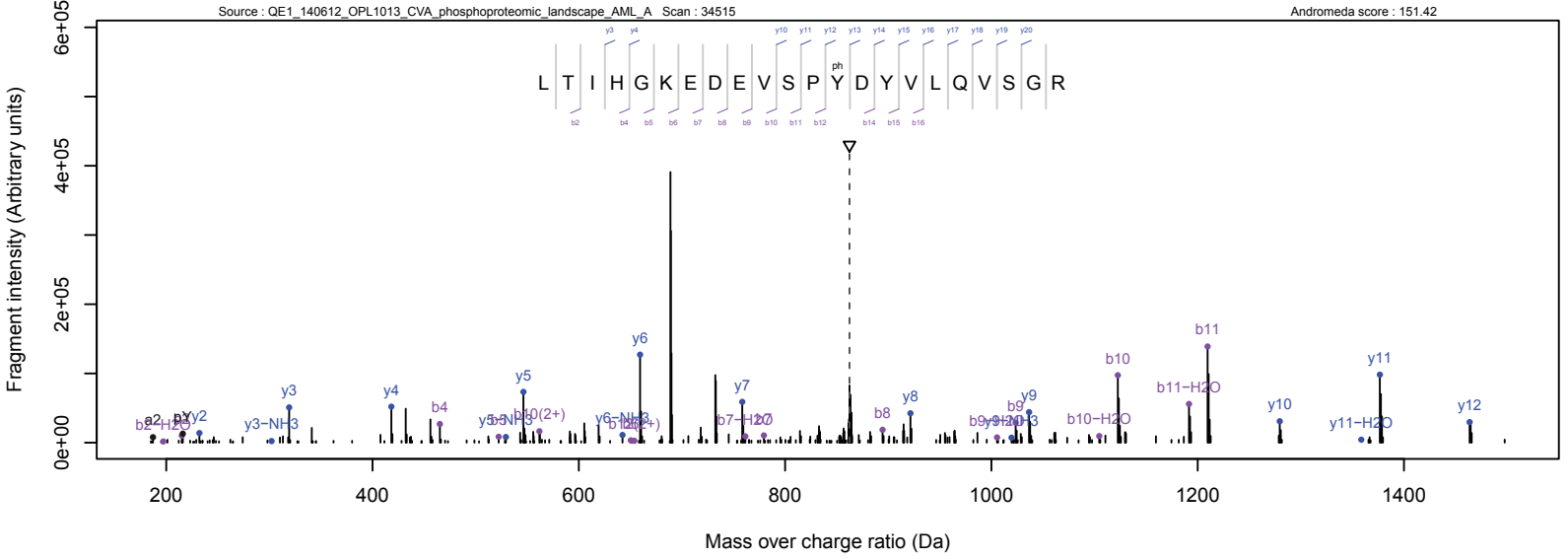
Andromeda score : 151.56



**LTIHGKEDEVSPY(ph)DYVLQVSGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34515

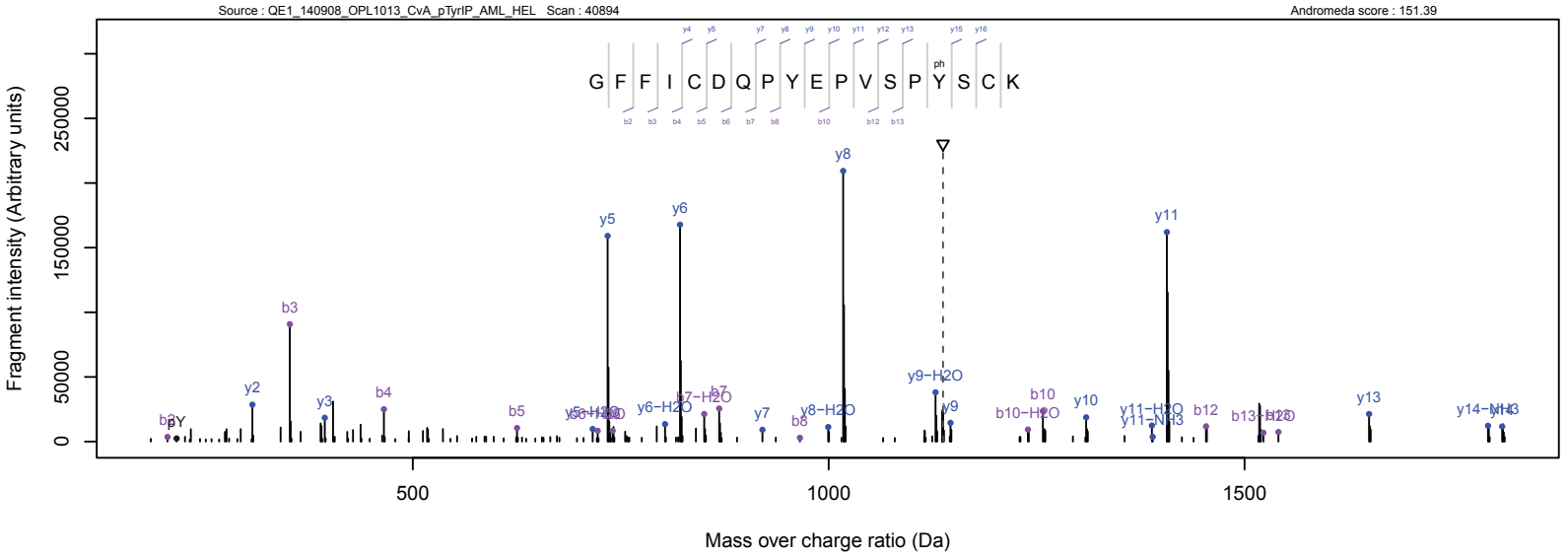
Andromeda score : 151.42



**GFFICDQPYEPVSPY(ph)SCK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 40894

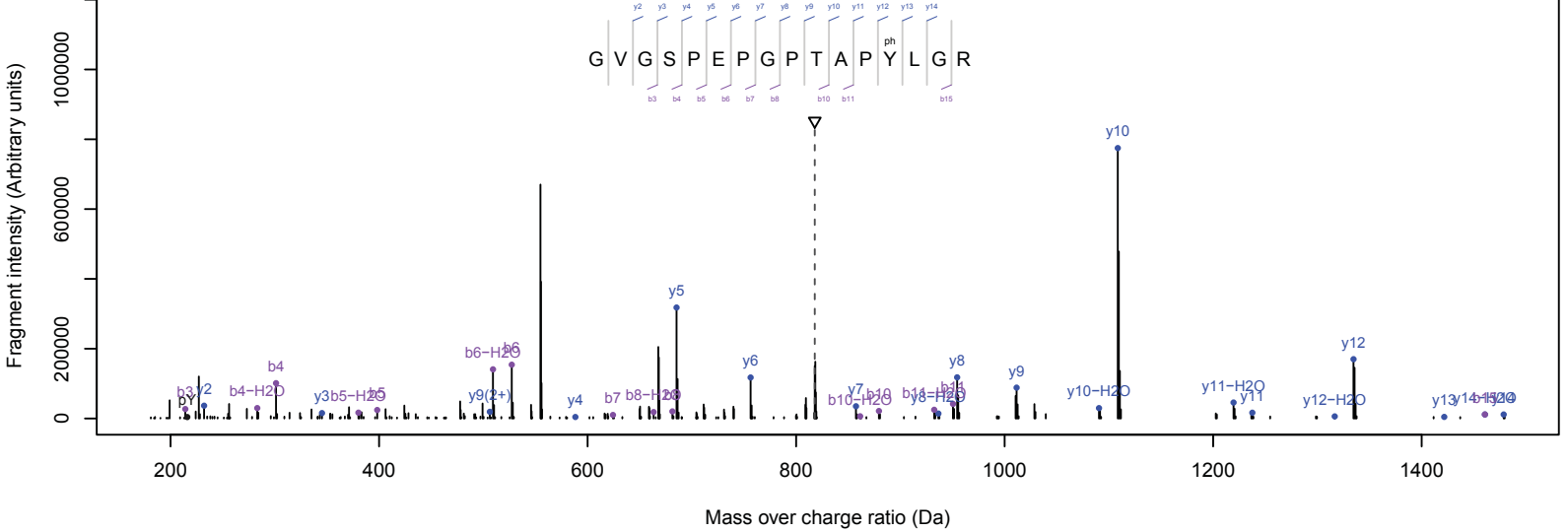
Andromeda score : 151.39



**GVGSPGPATPY(ph)LGR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21937

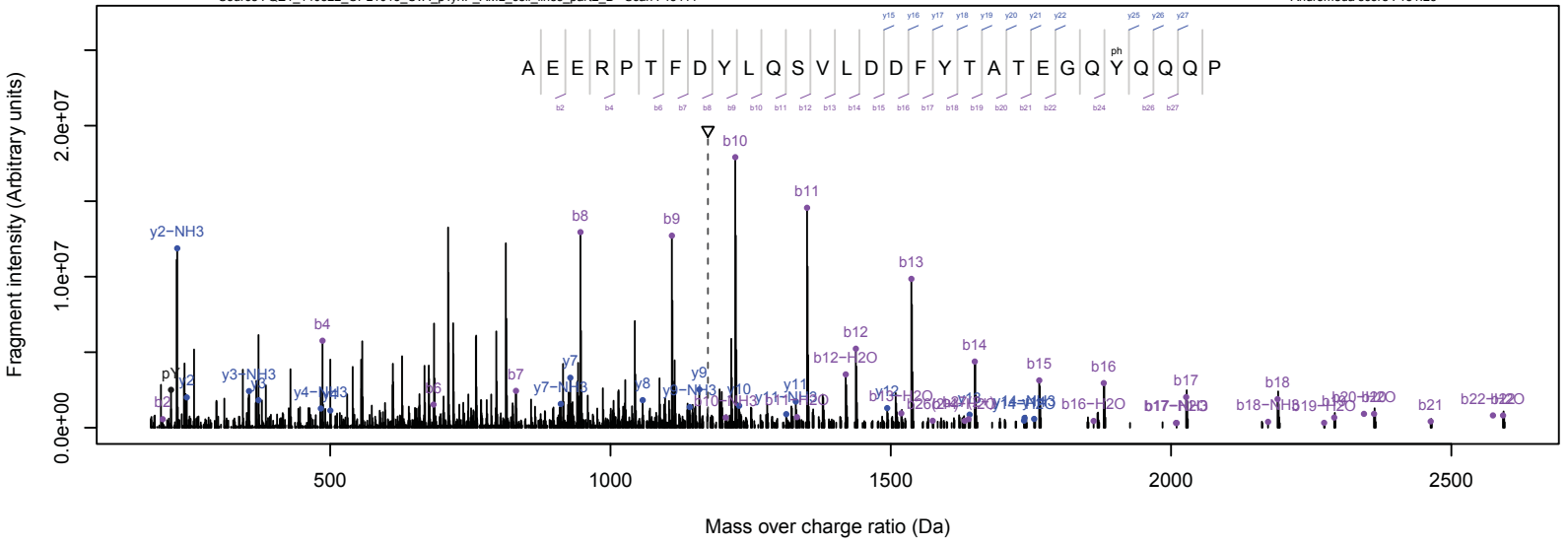
Andromeda score : 151.38



**AEERPTFDYLSVLDDFYTATEGQY(ph)QQQ-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 45411

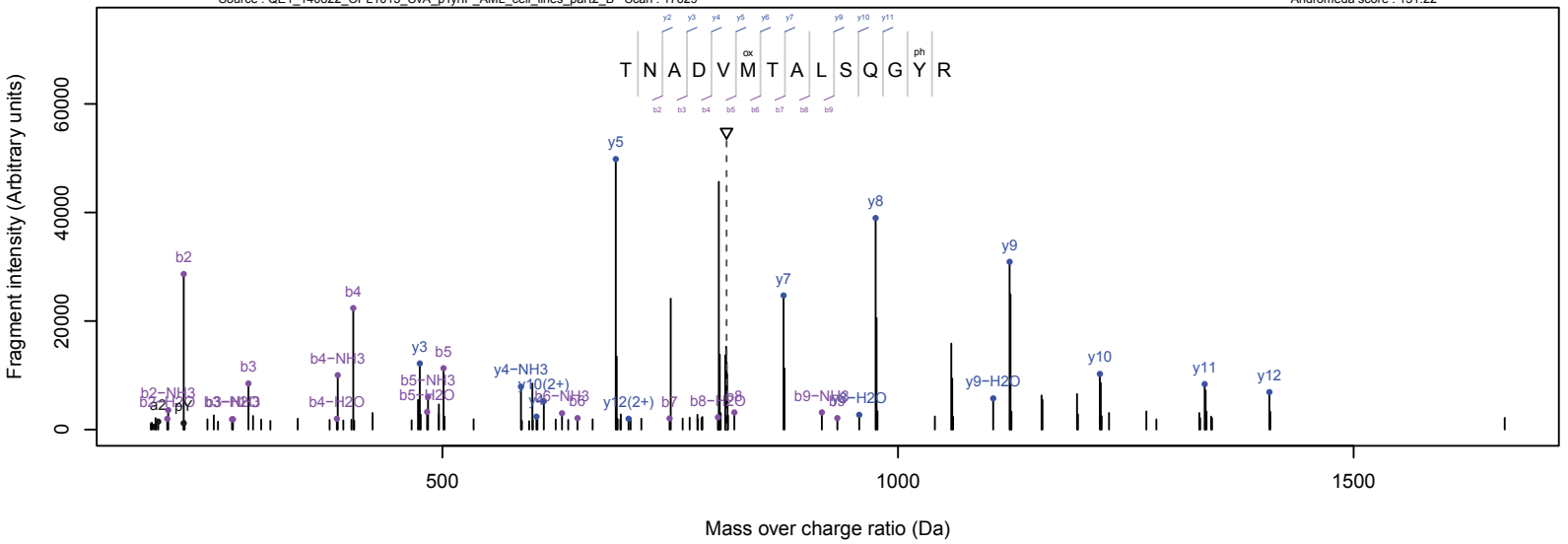
Andromeda score : 151.25



**TNADVM(ox)TALSQGY(ph)R-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 17629

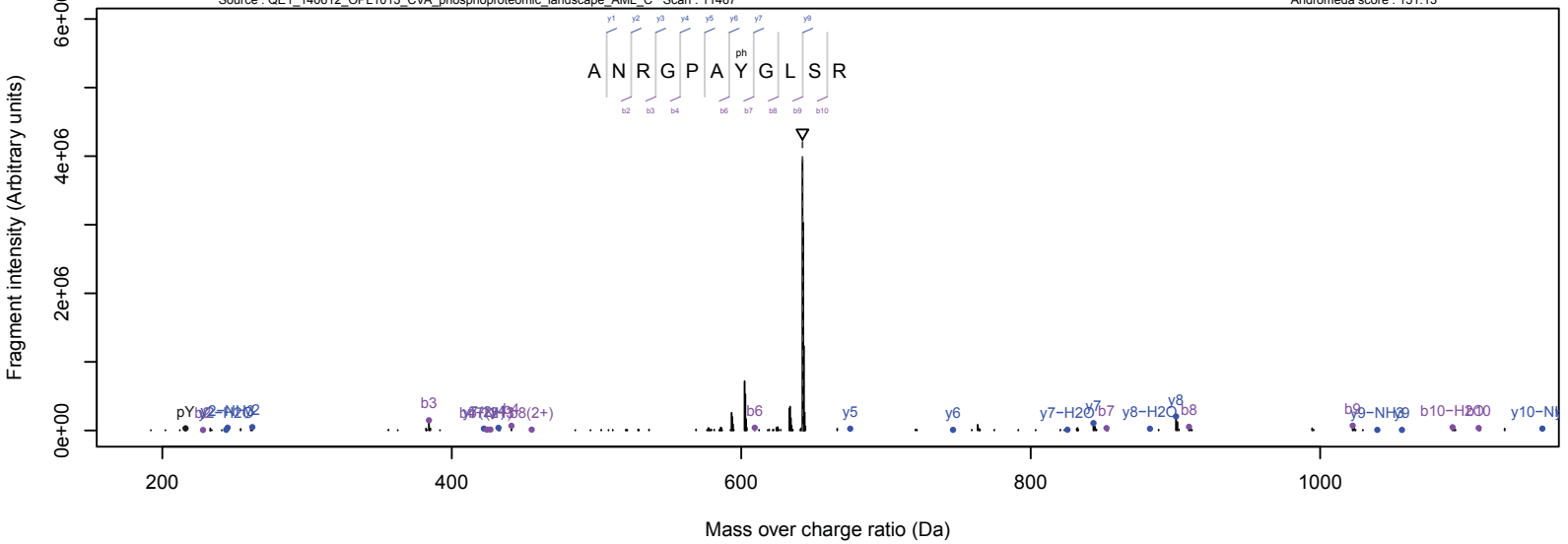
Andromeda score : 151.22



**(ac)ANRGPAY(ph)GLSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11467

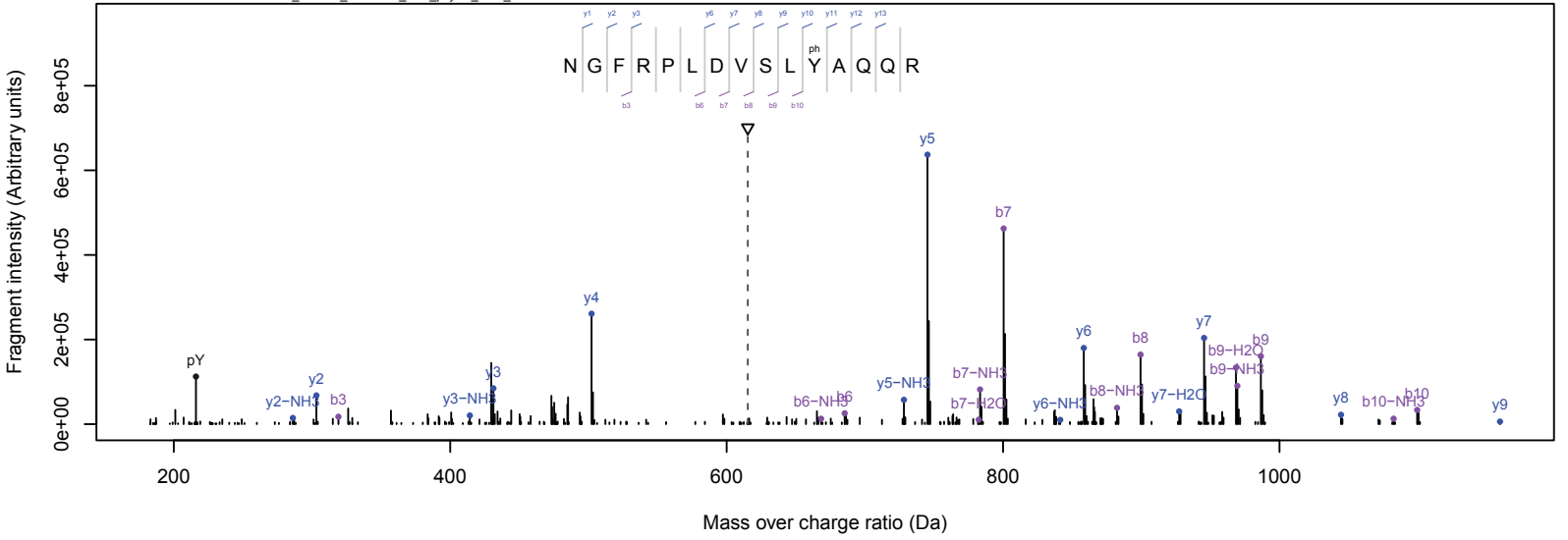
Andromeda score : 151.13



NGFRPLDVSLY(ph)AQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 33009

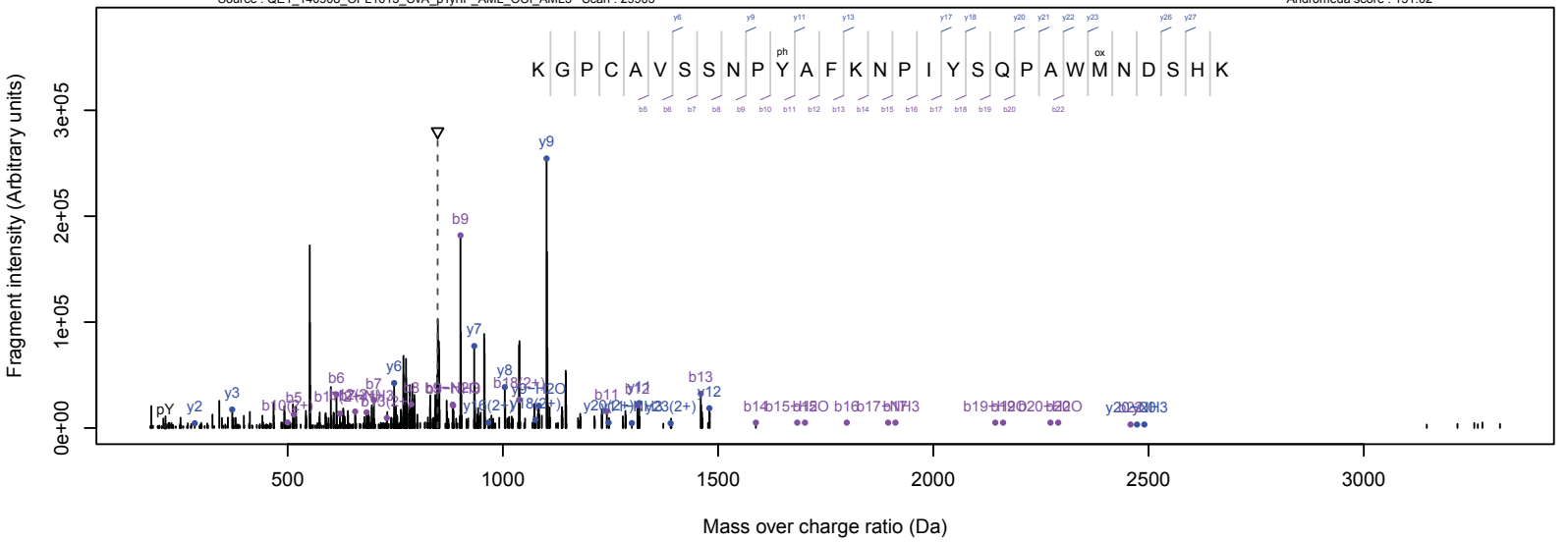
Andromeda score : 151.07



KGPCAVSSNPY(ph)AFKNPIYSQPAWM(ox)NDSHK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 29903

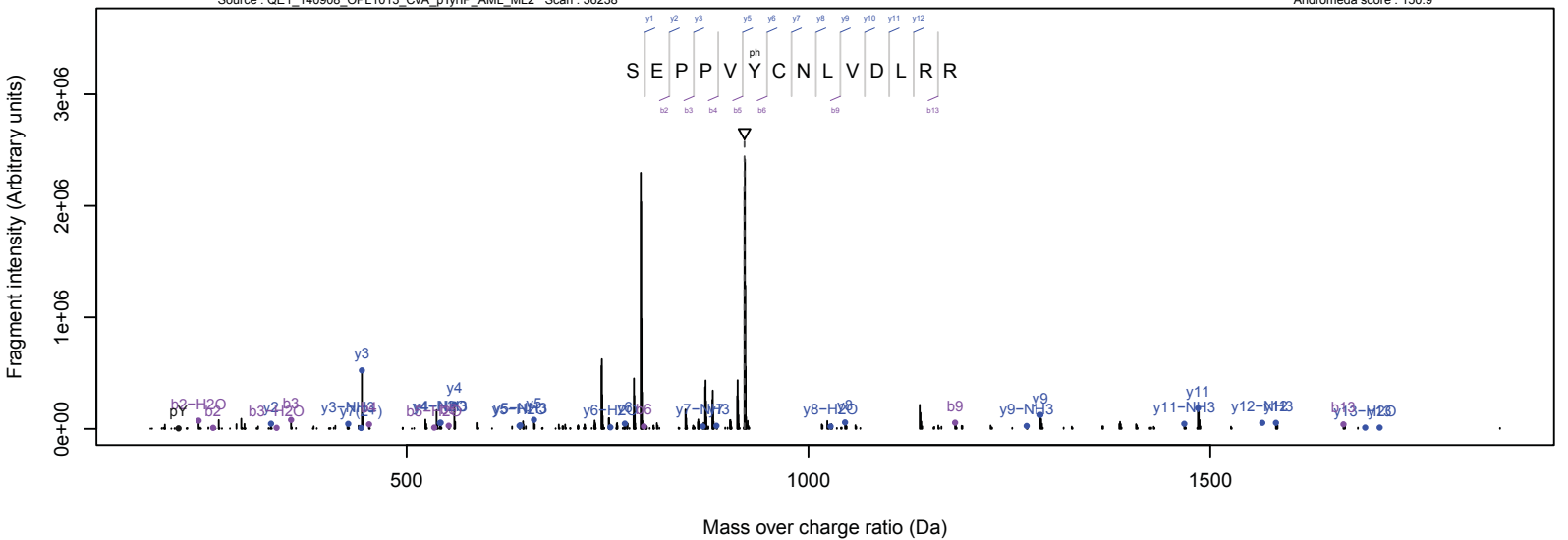
Andromeda score : 151.02



(ac)SEPPVY(ph)CNLVDLRR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 36238

Andromeda score : 150.9

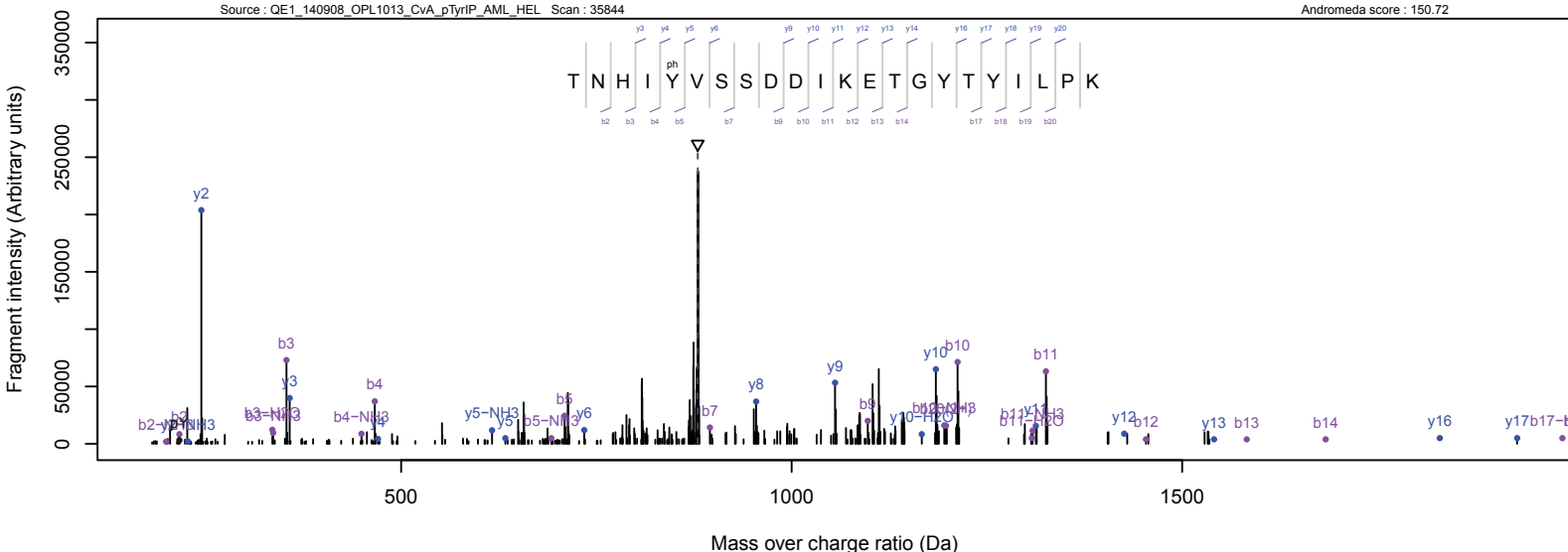




**TNHIIYVSSDDIKETGYTYILPK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35844

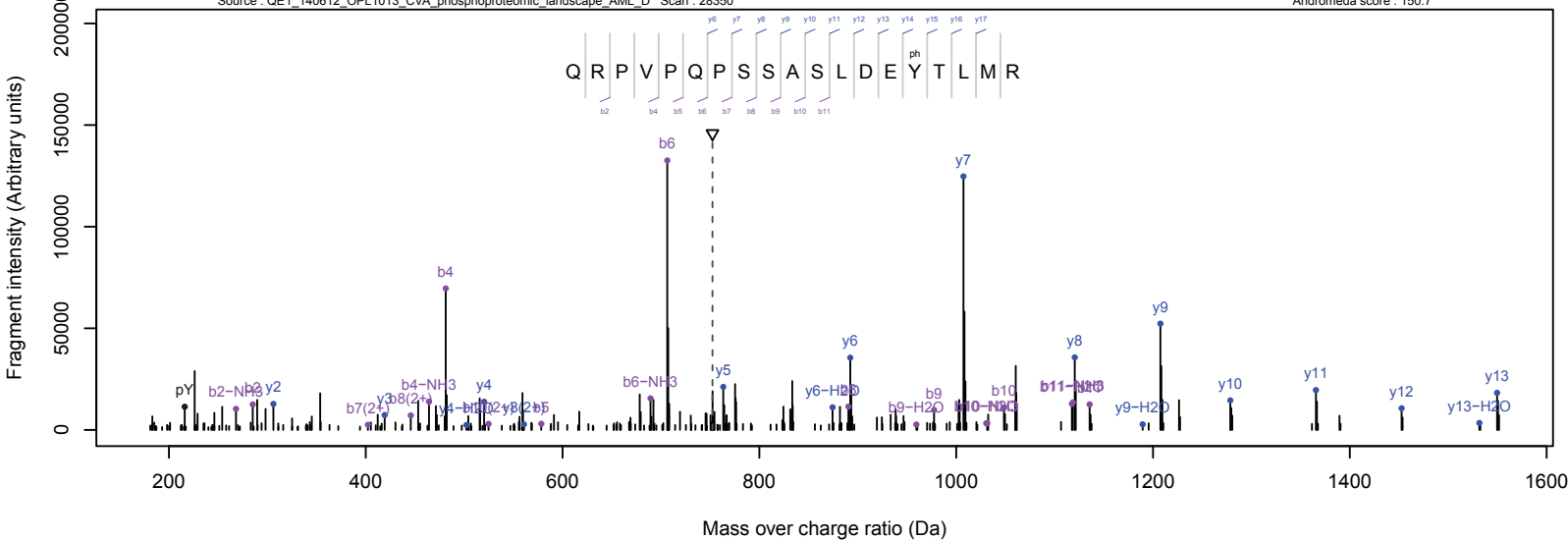
Andromeda score : 150.72



**QRPVPQPSSASLDEYTLMR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28350

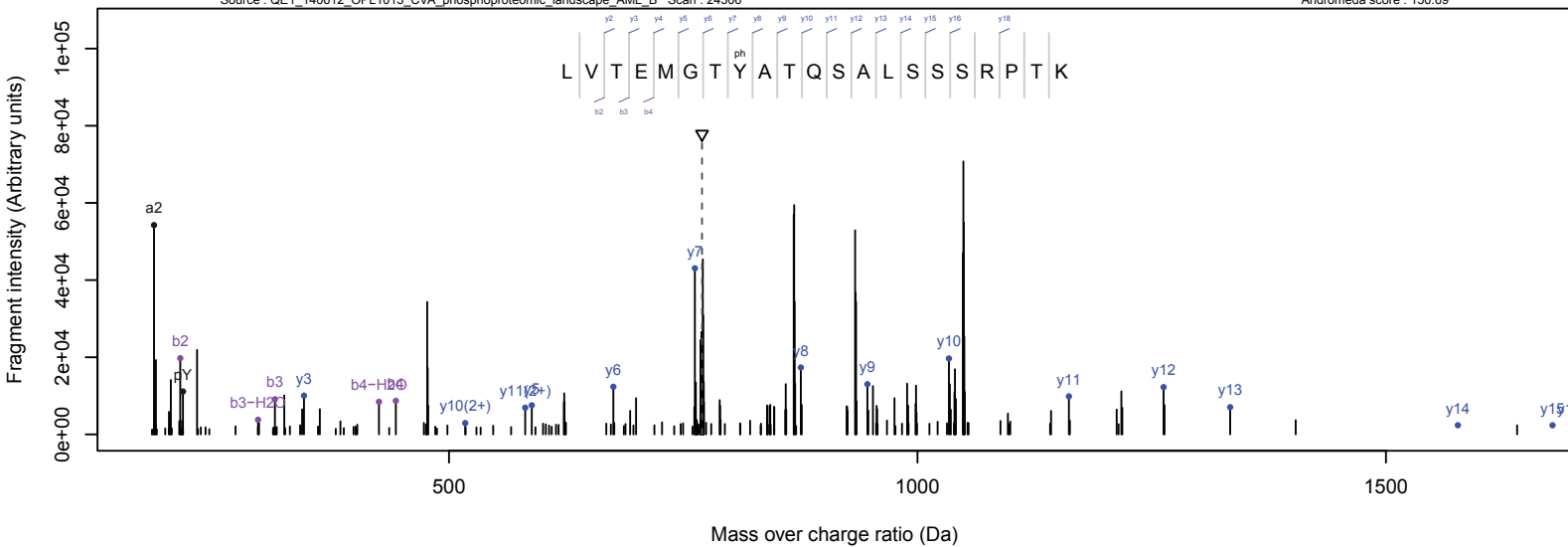
Andromeda score : 150.7



**LVTEMGTYATQSALSSSRPTK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24306

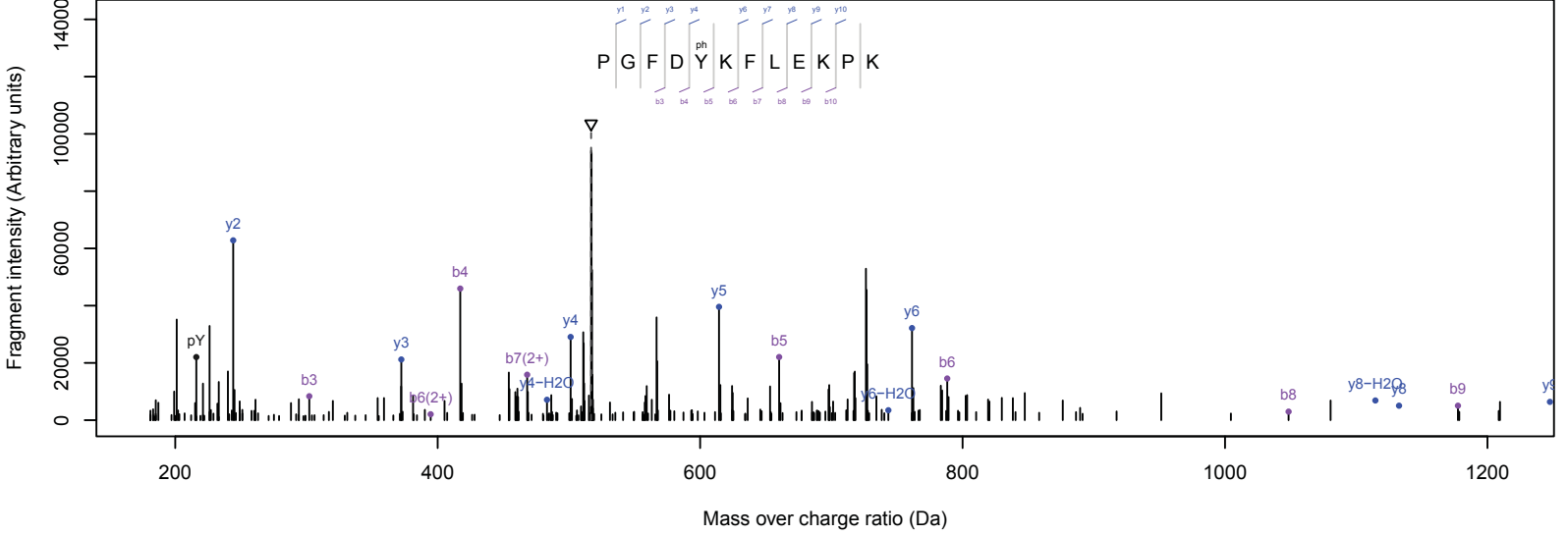
Andromeda score : 150.69



**PGFDY(ph)KFLEKPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 25696

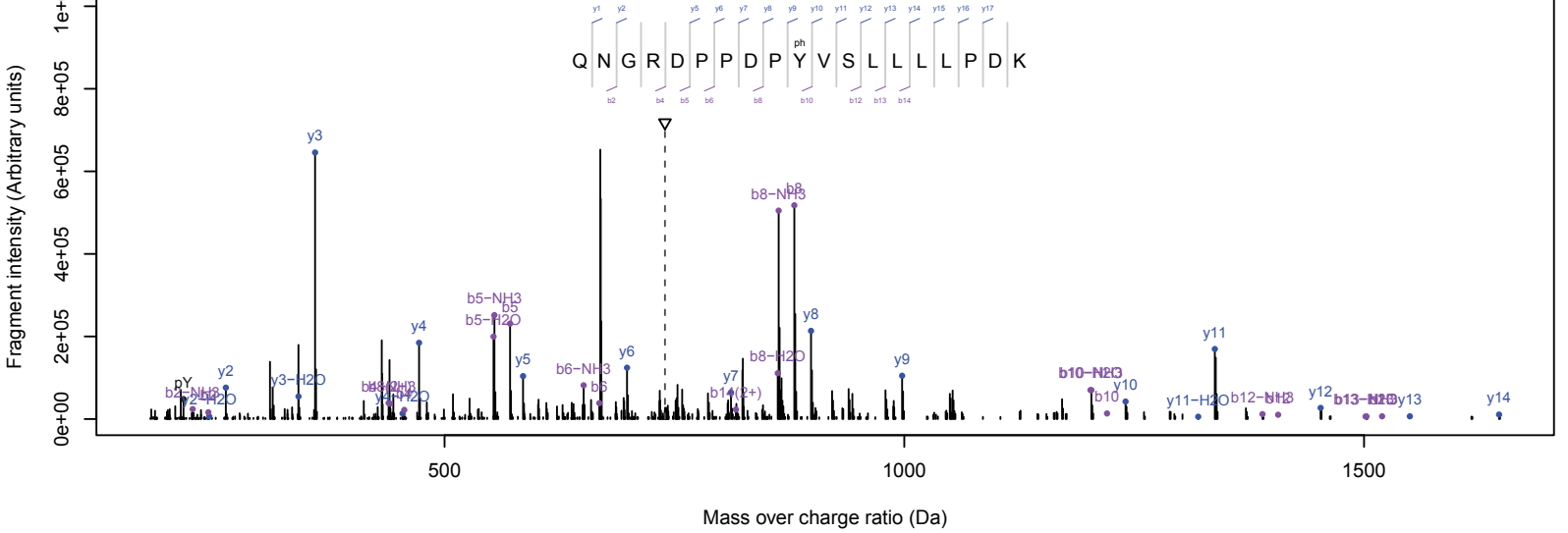
Andromeda score : 150.69



**QNGRDPPDYPVSLLLLPDK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41872

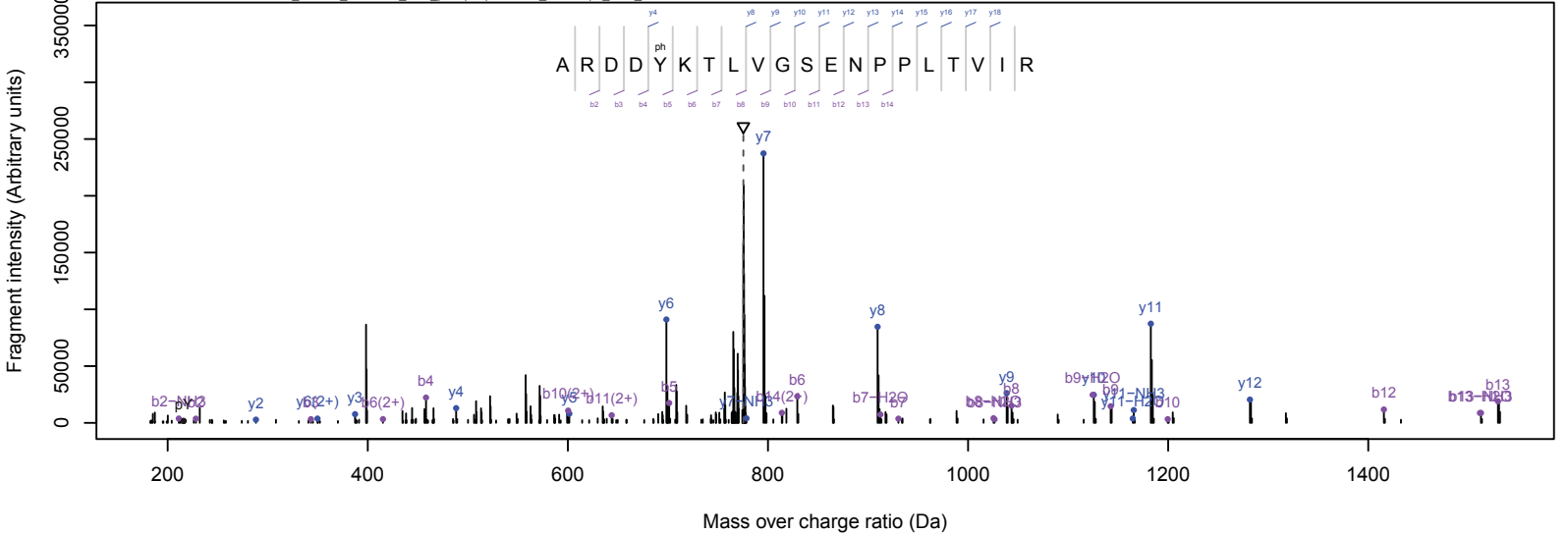
Andromeda score : 150.69



**ARDDY(ph)KTLVGSENPPLTVIR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 29957

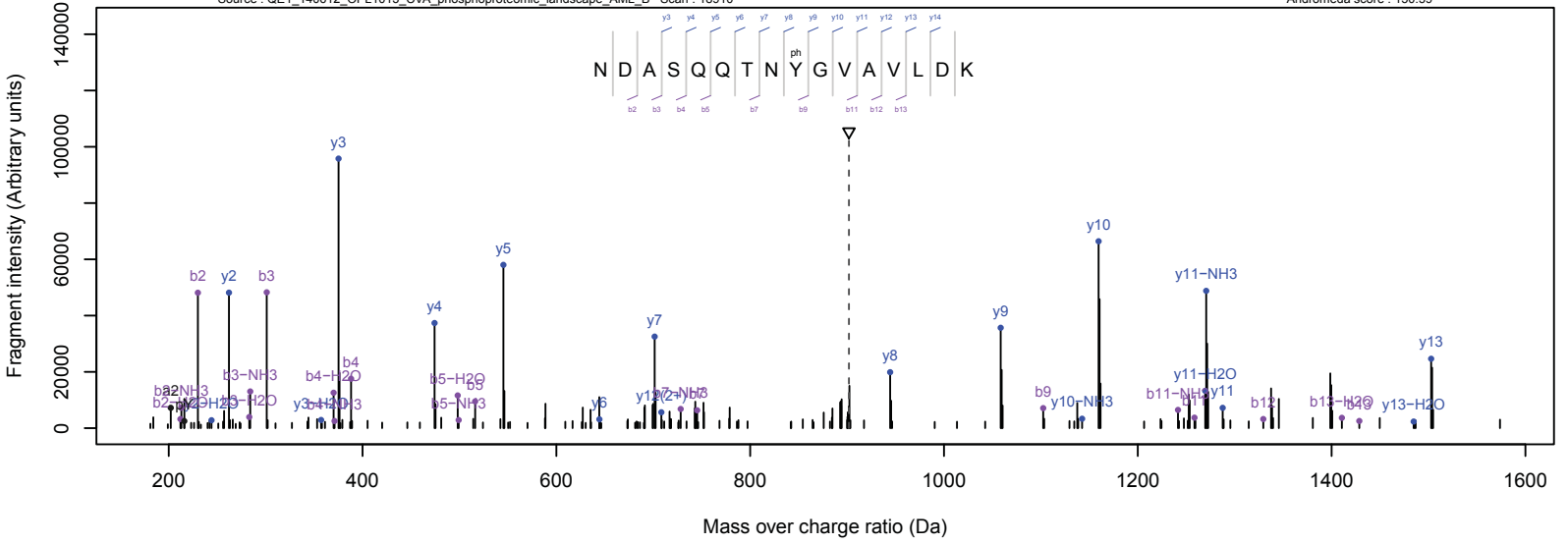
Andromeda score : 150.65



**NDASQQTNY(ph)GVAVLDK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18910

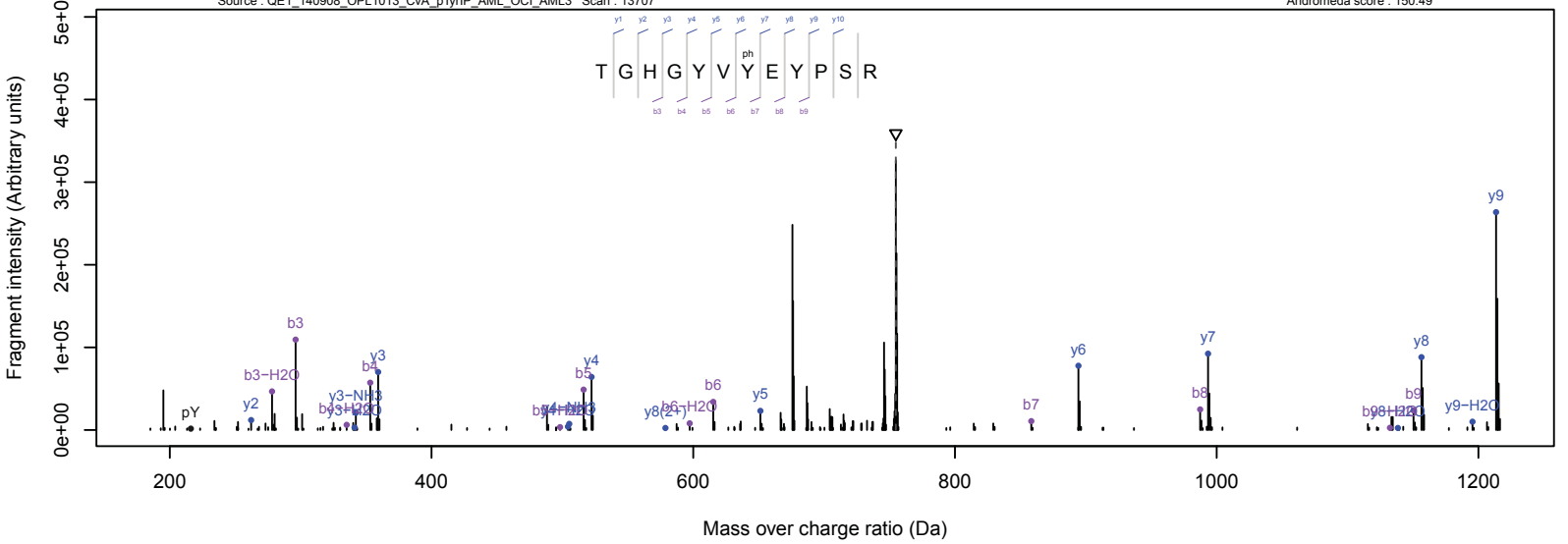
Andromeda score : 150.59



**TGHGYVY(ph)EYPSR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 13707

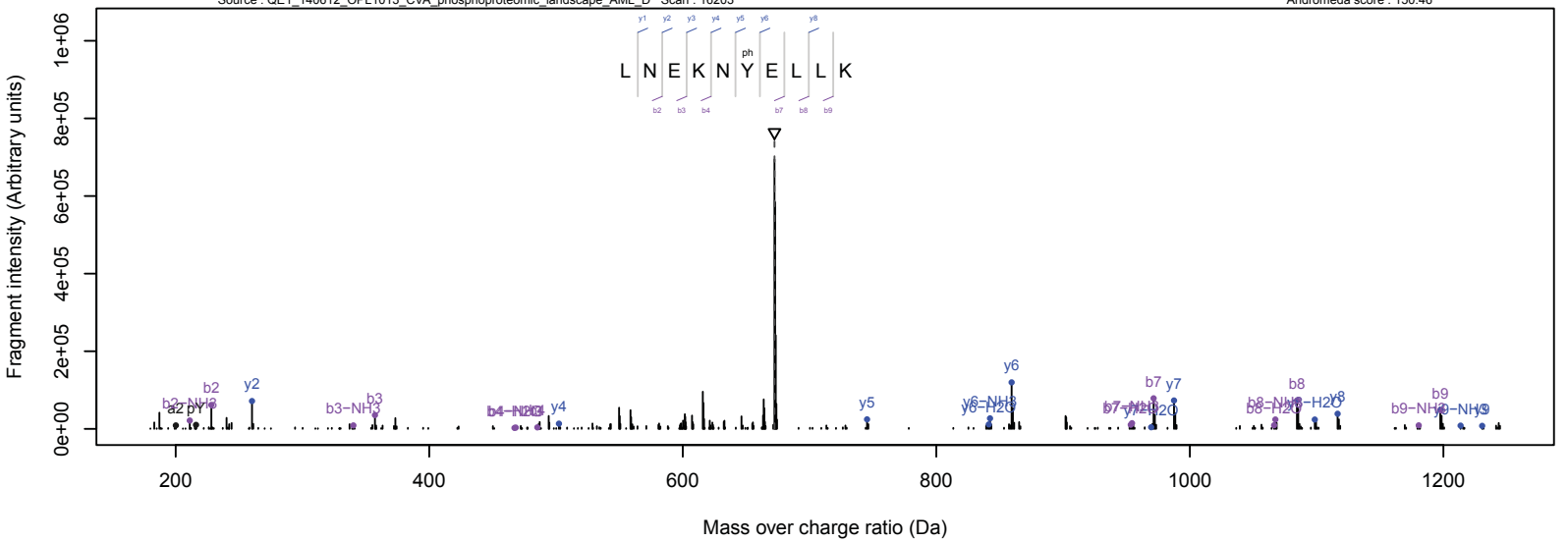
Andromeda score : 150.49



**LNEKNY(ph)ELLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 16203

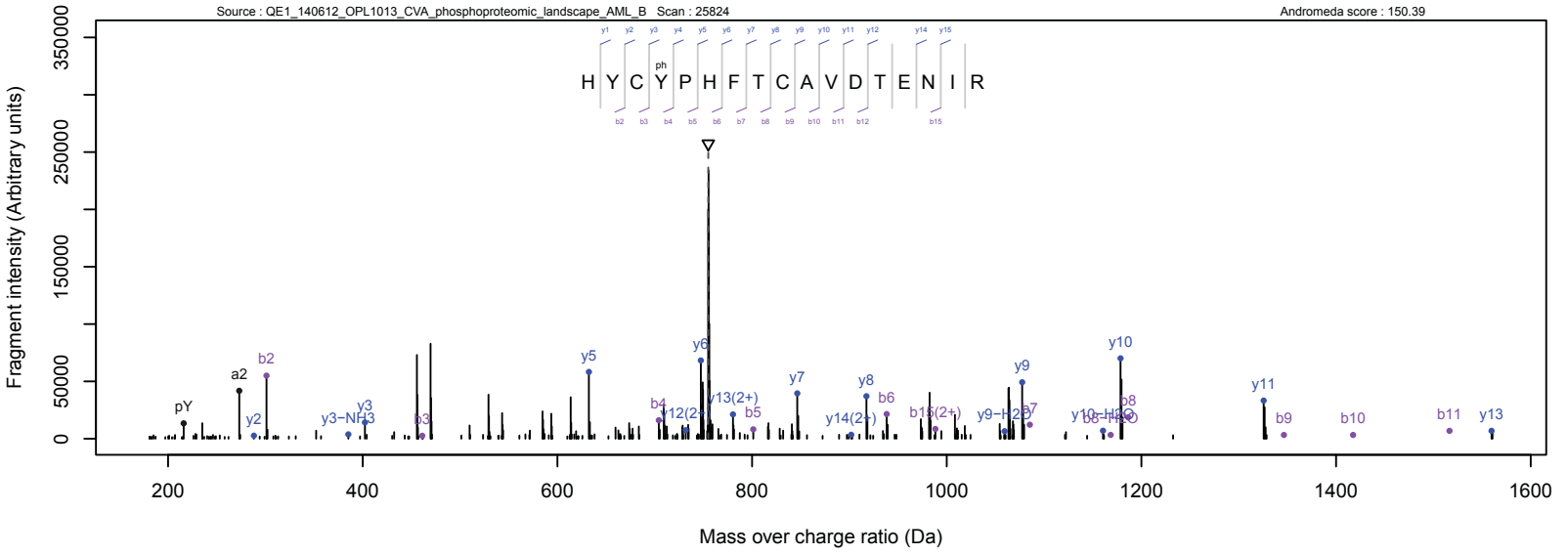
Andromeda score : 150.46



**HYCY(ph)PHFTCAVDTEIR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 25824

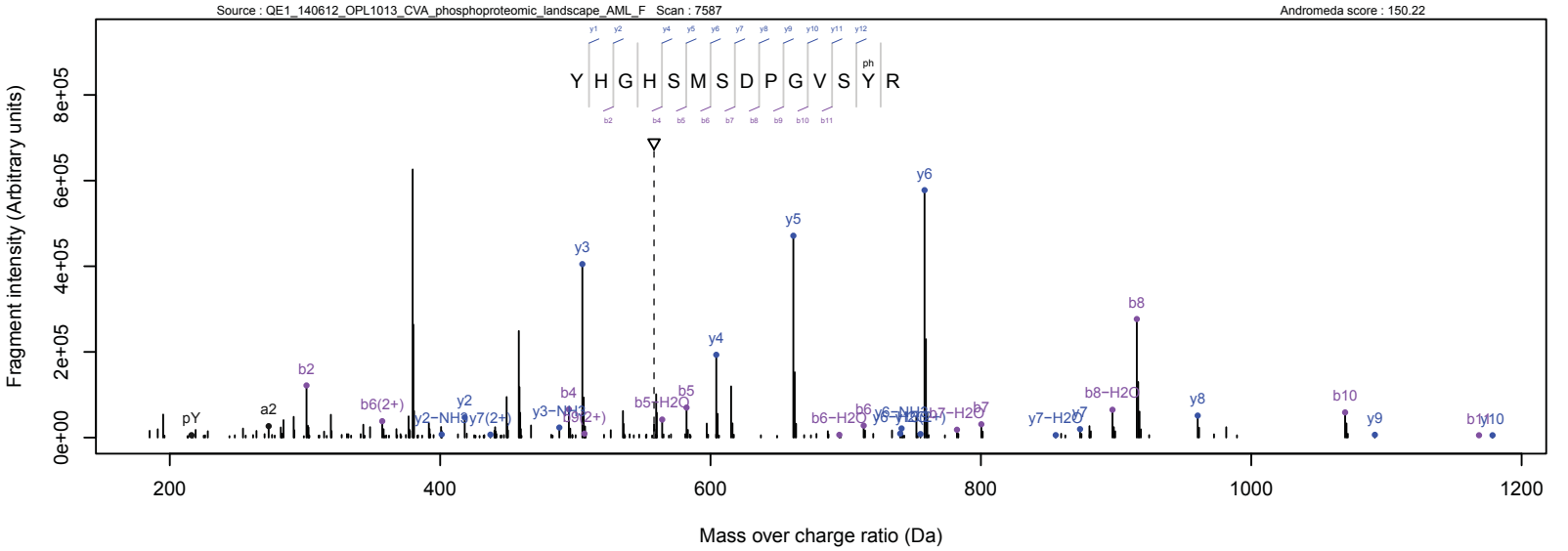
Andromeda score: 150.39



**YHGHSMDPGVSY(ph)R-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan: 7587

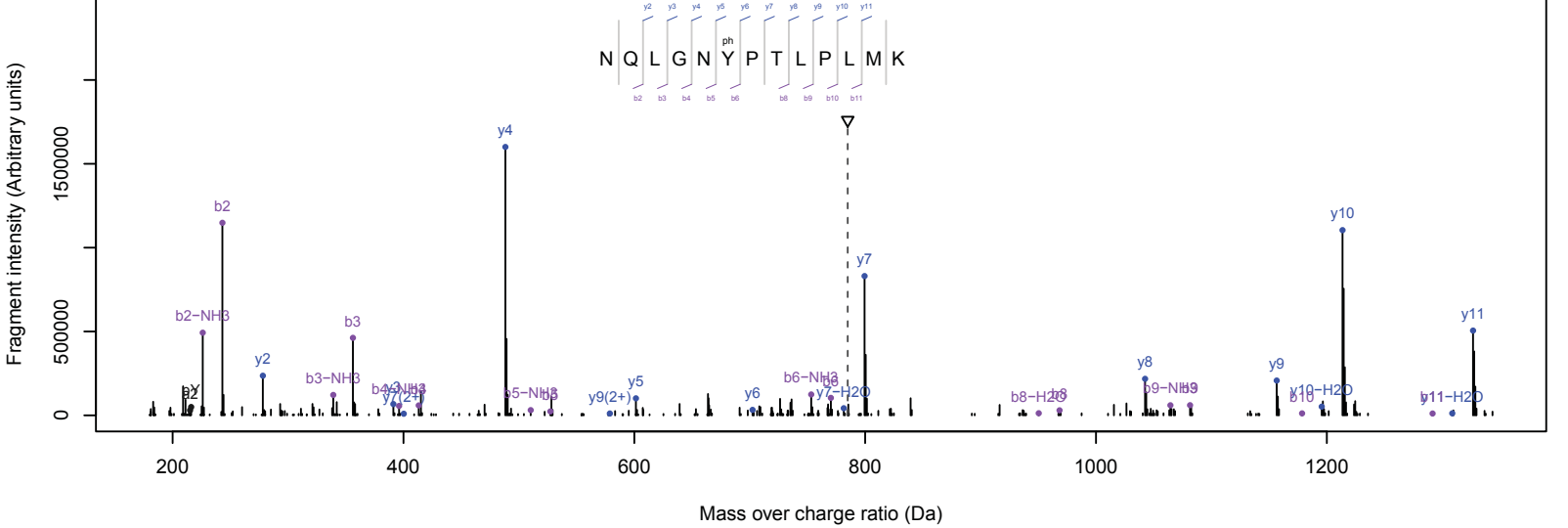
Andromeda score: 150.22



**NQLGNY(ph)PTLPLMK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36338

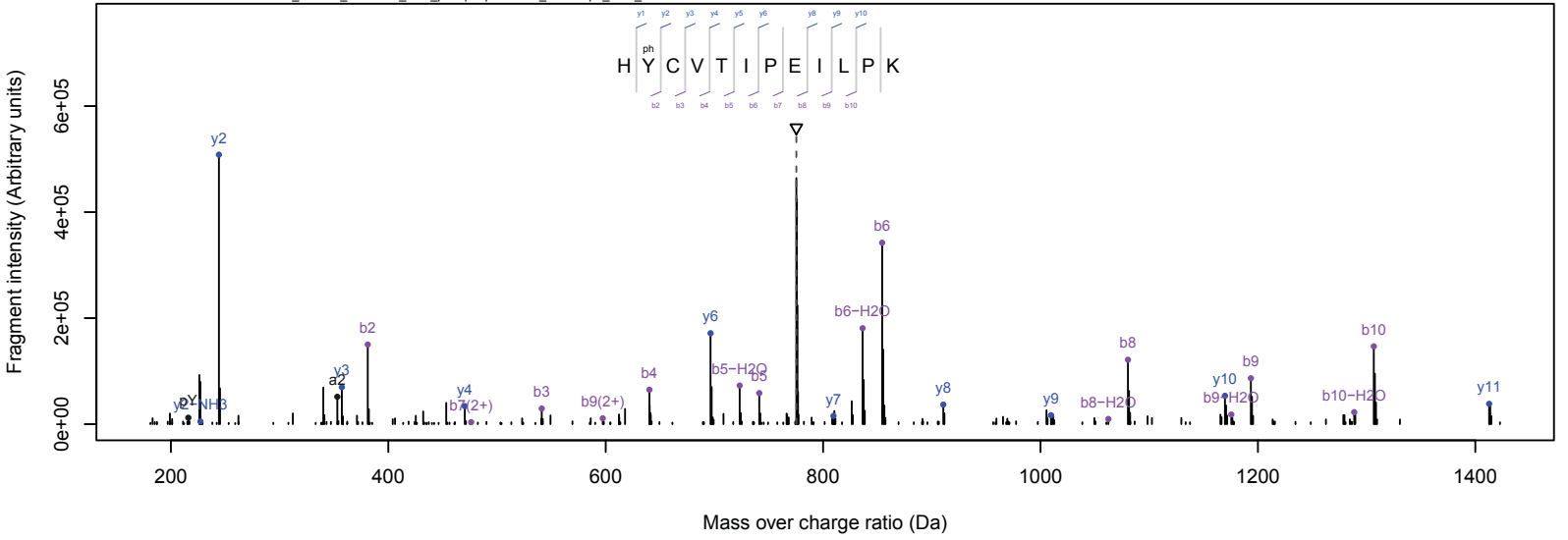
Andromeda score: 150.11



HY(ph)CVTIPEILPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 35131

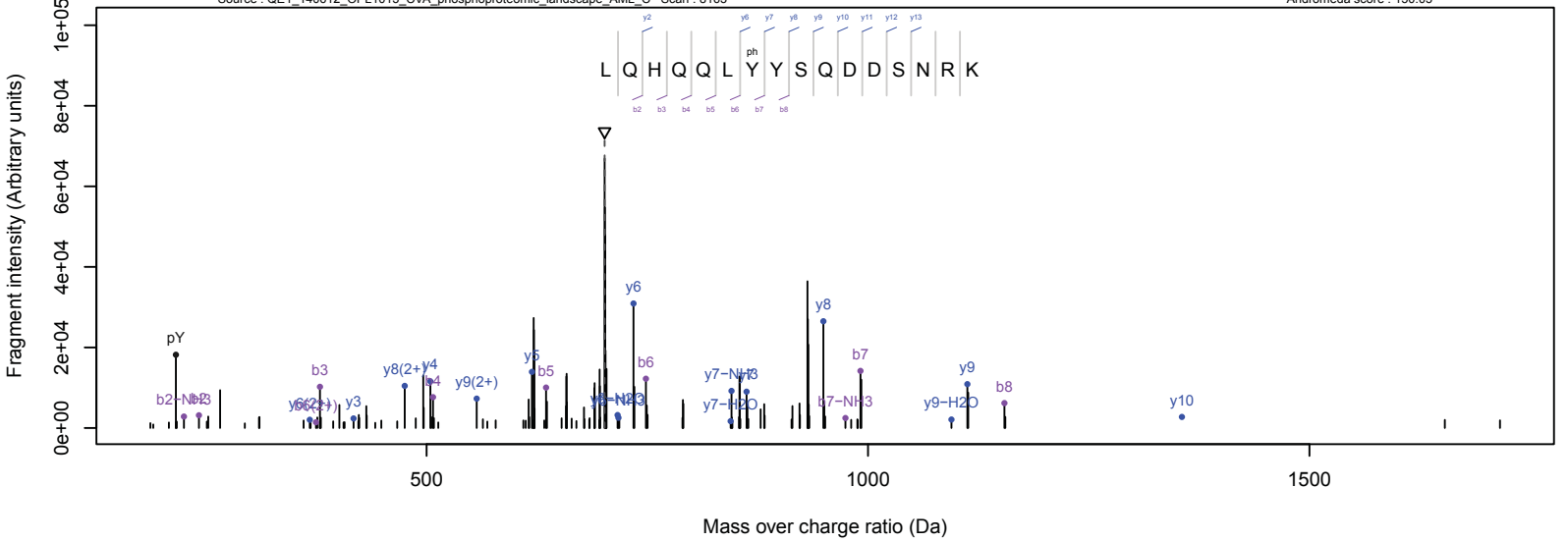
Andromeda score : 150.06



LQHQQLY(ph)YSQDDSNRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8163

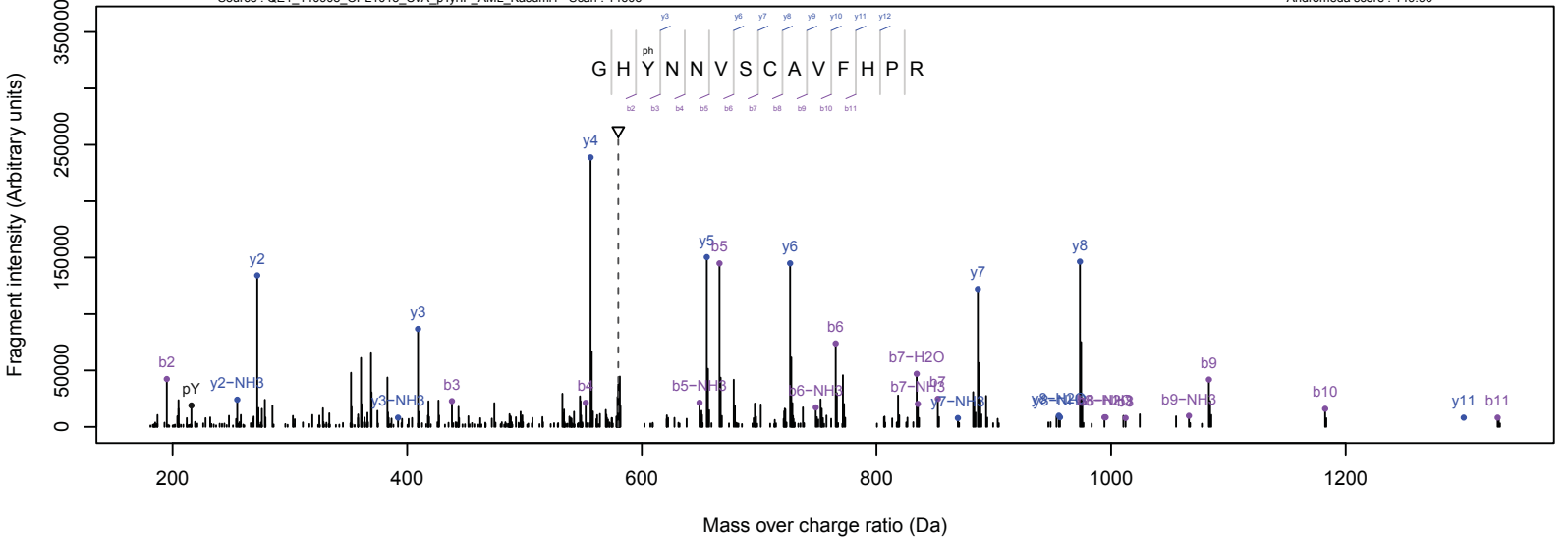
Andromeda score : 150.05

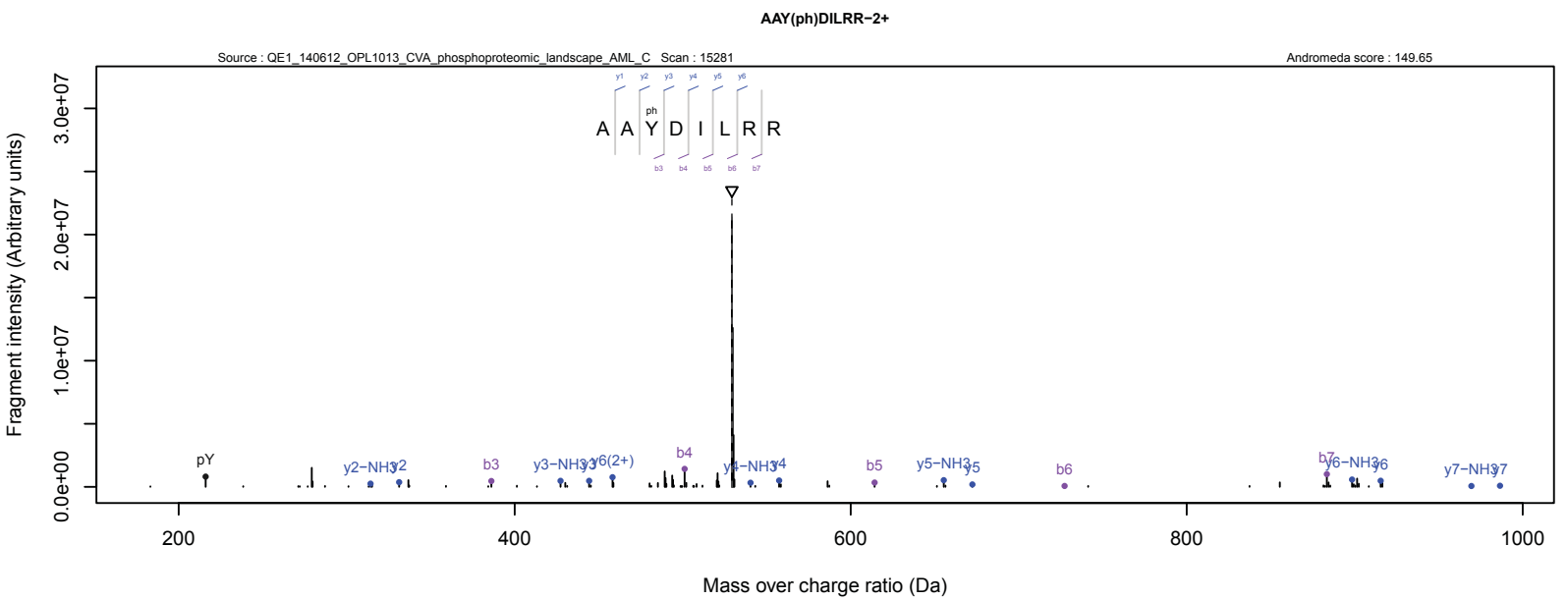
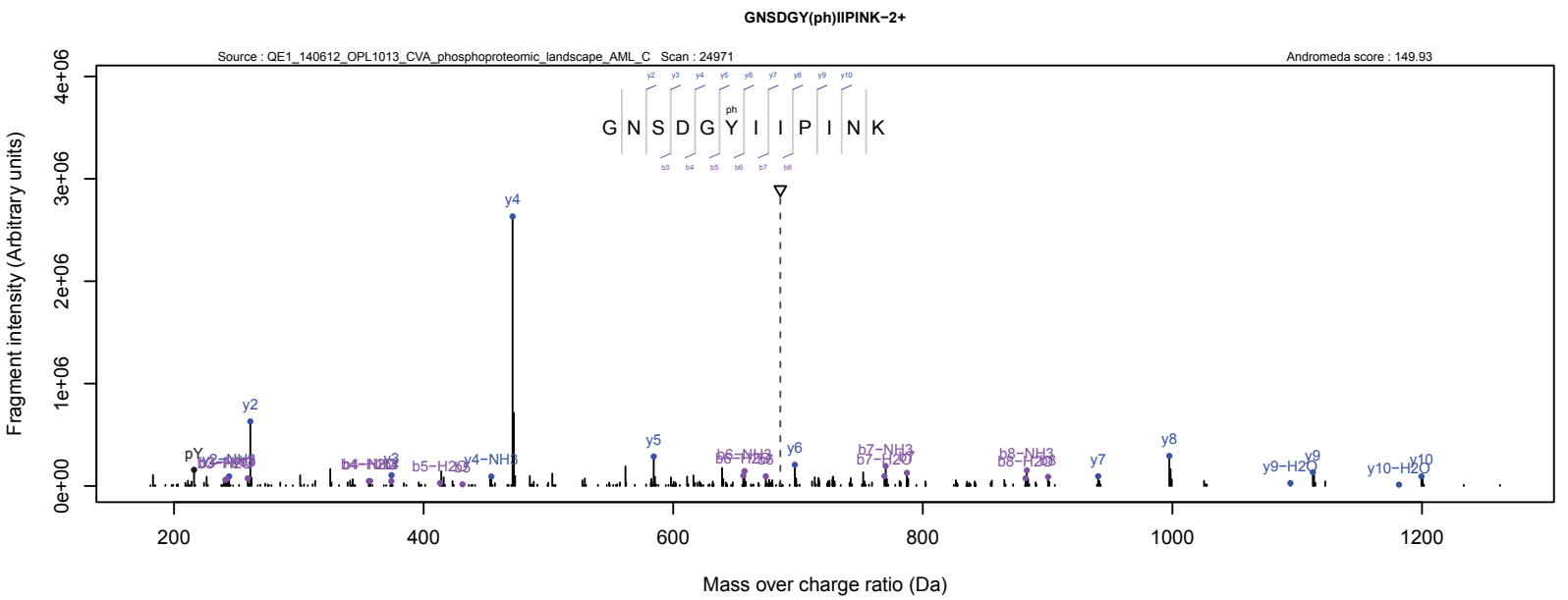
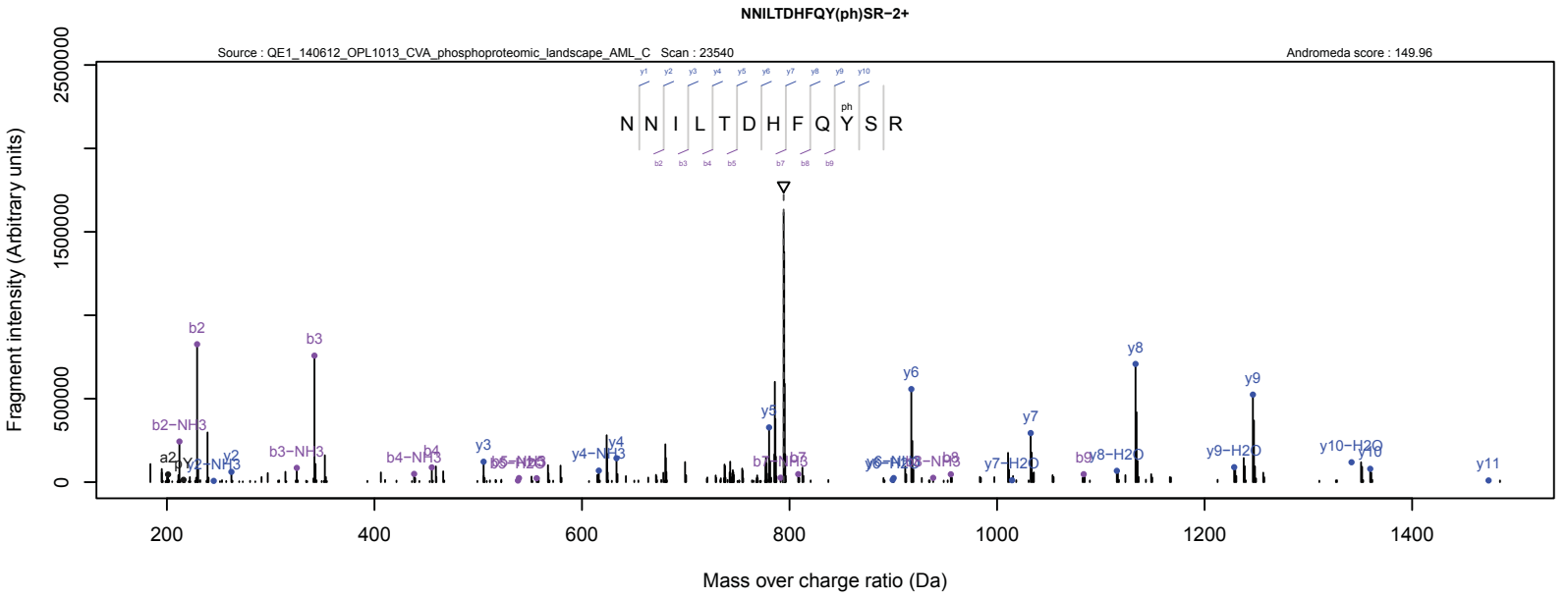


GHY(ph)NNVSCAVFHPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 14606

Andromeda score : 149.96

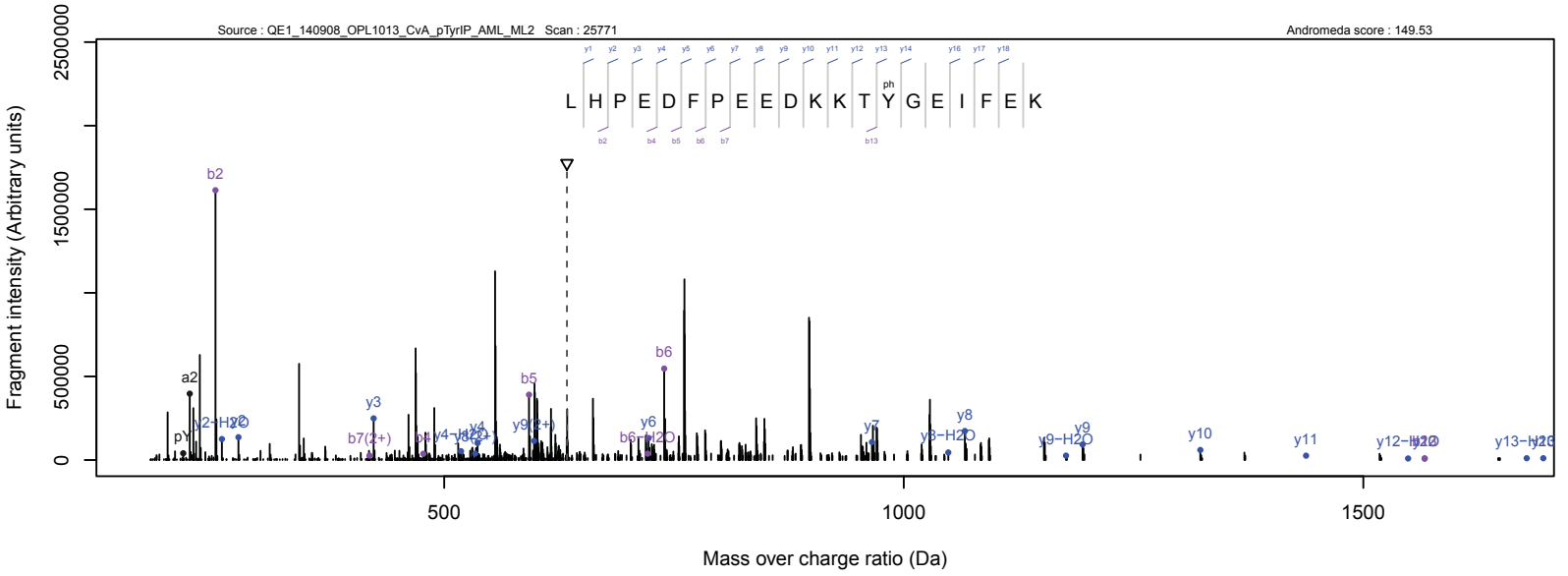




LHPEDFPEEDKKTYPHGEIFEK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 25771

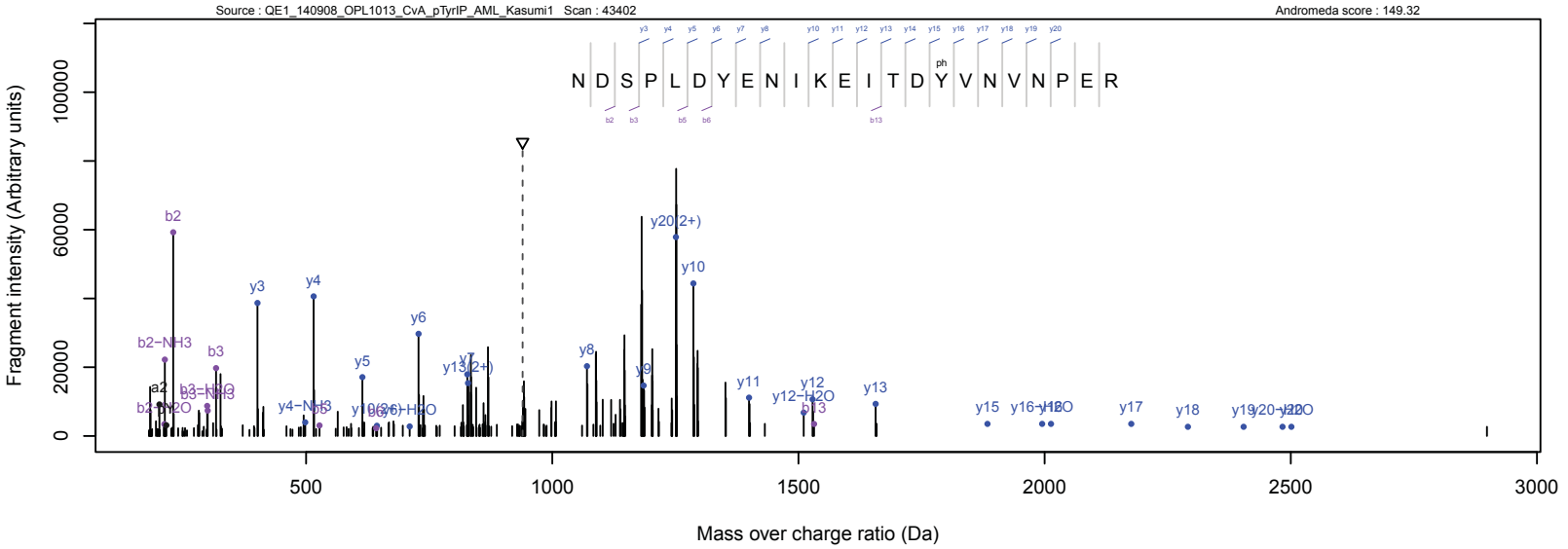
Andromeda score : 149.53



NDSPLDYENIKEITDYVNVNPER-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 43402

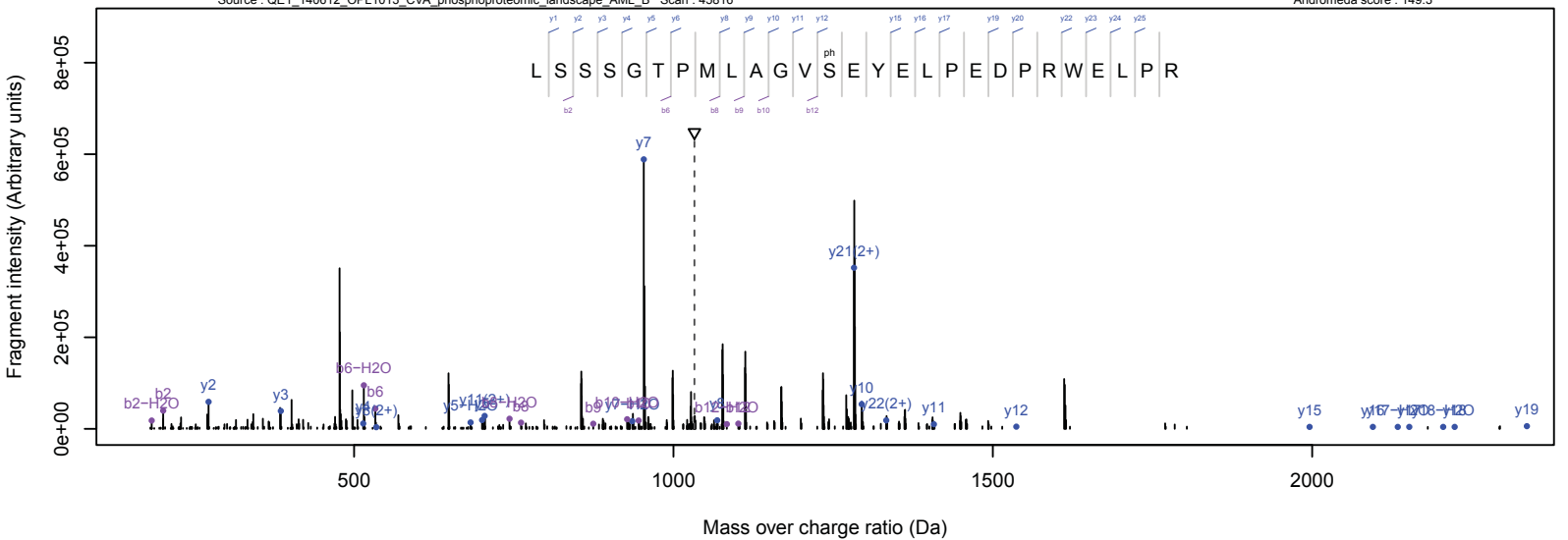
Andromeda score : 149.32

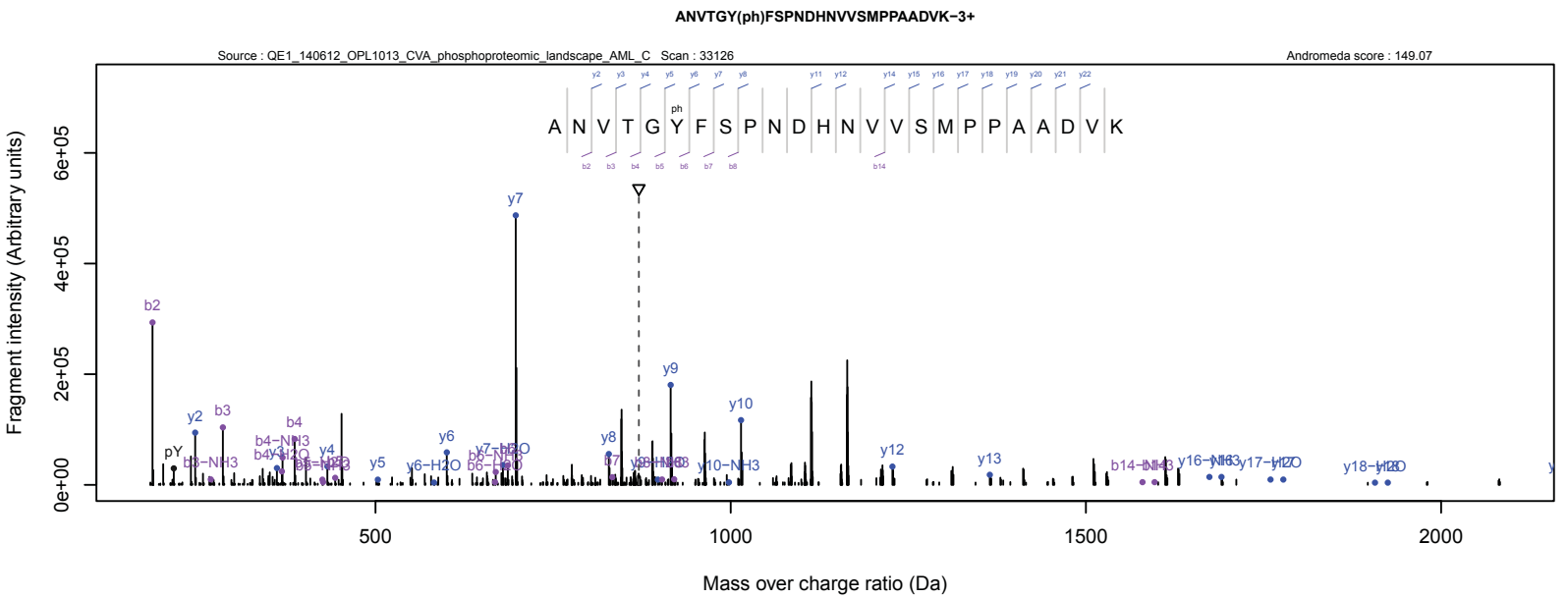
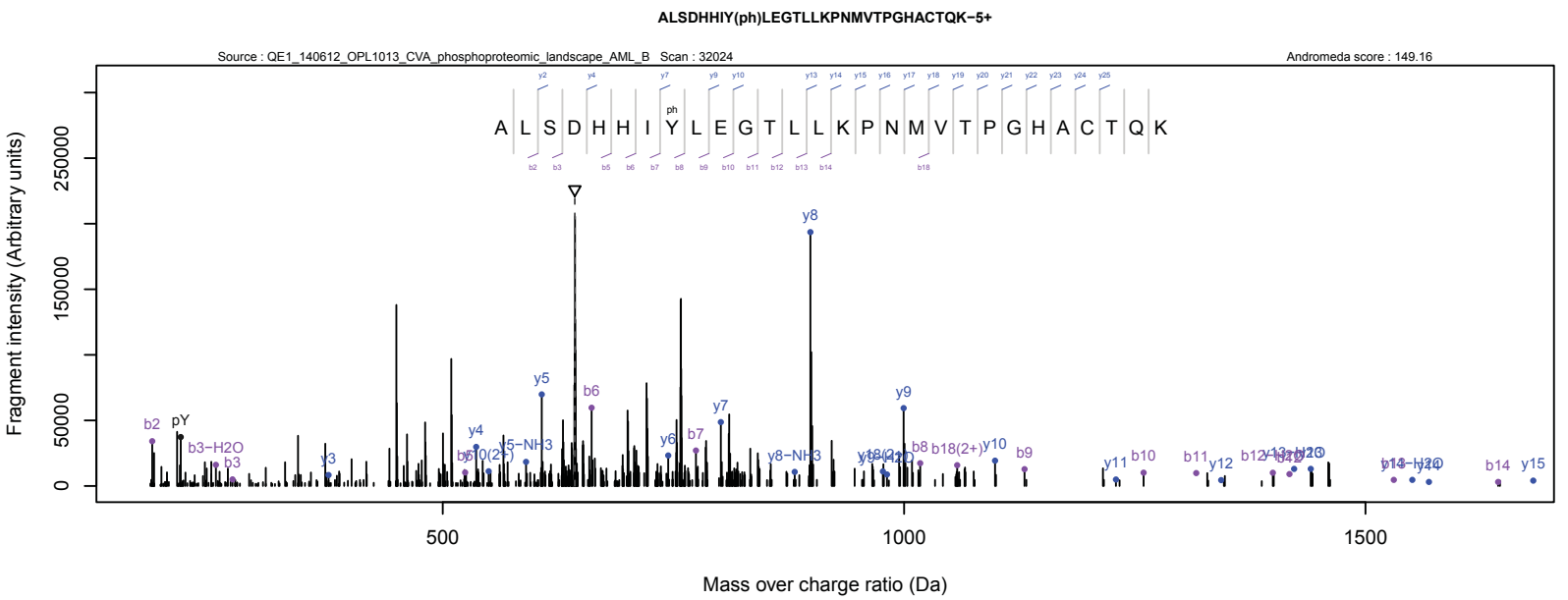
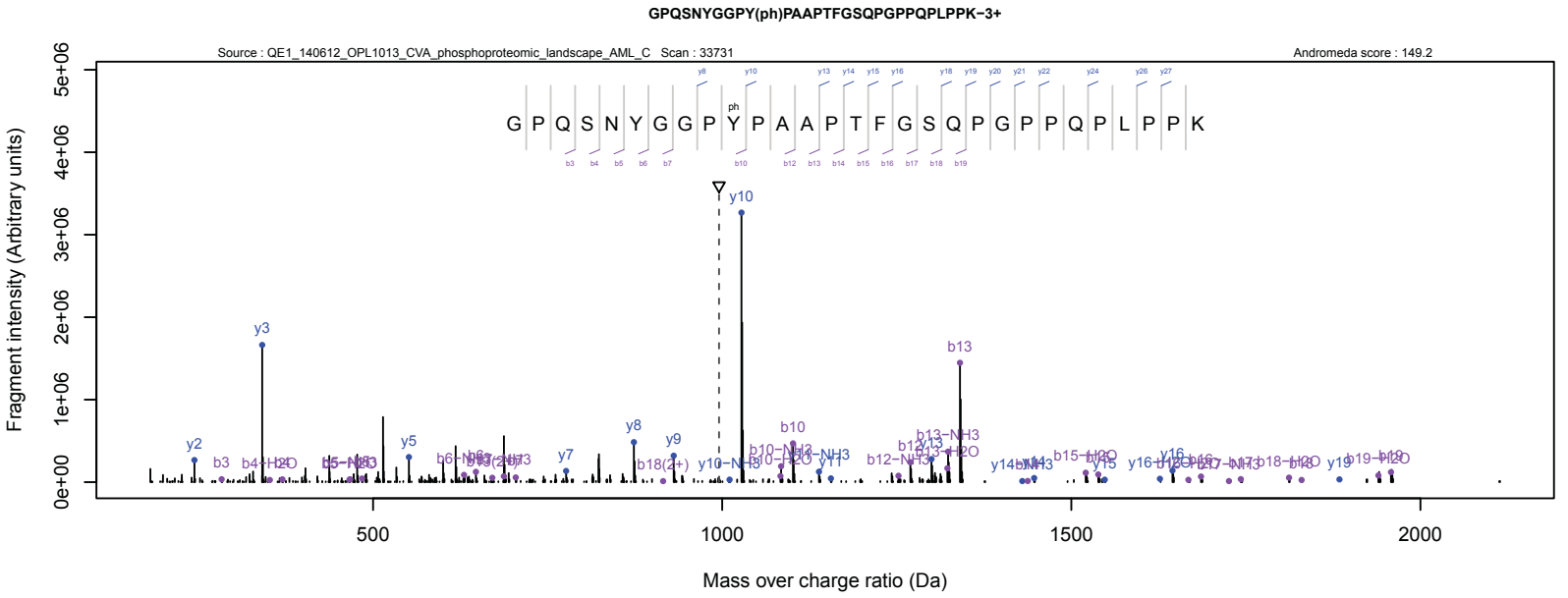


LSSSGTPMLAGVSEYELPEDPRWELPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45816

Andromeda score : 149.3



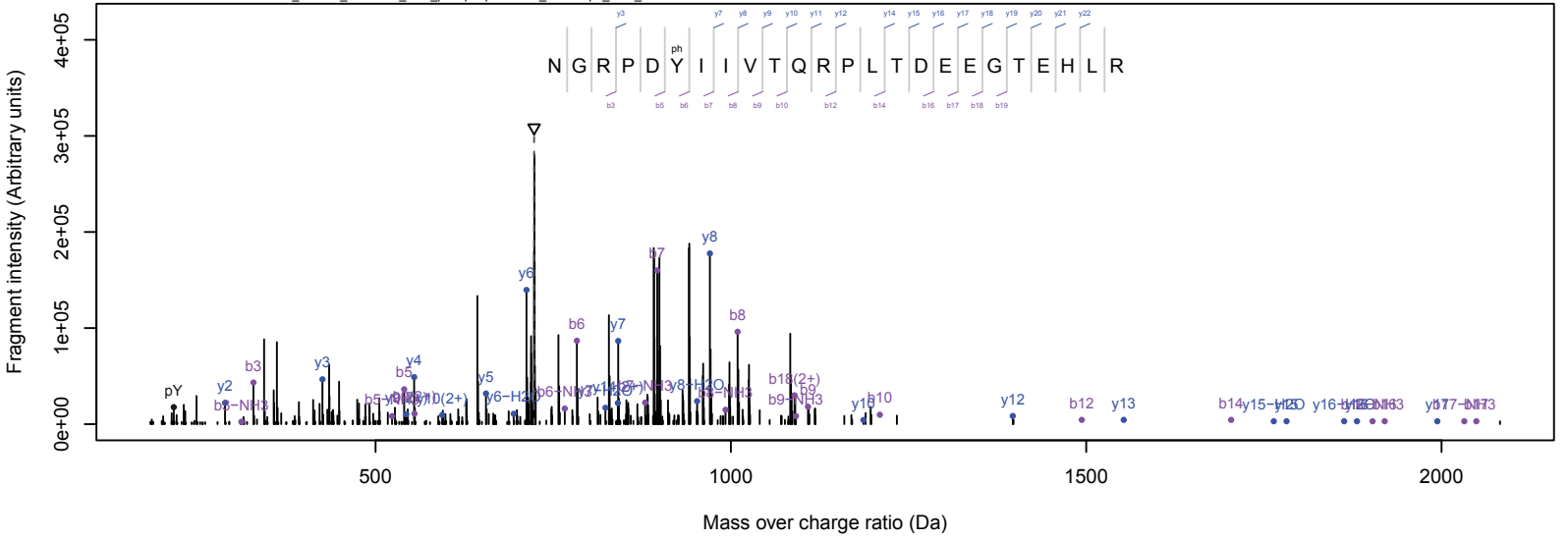




NGRPDY(ph)IIVTQRPLTDEEGTEHLR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25427

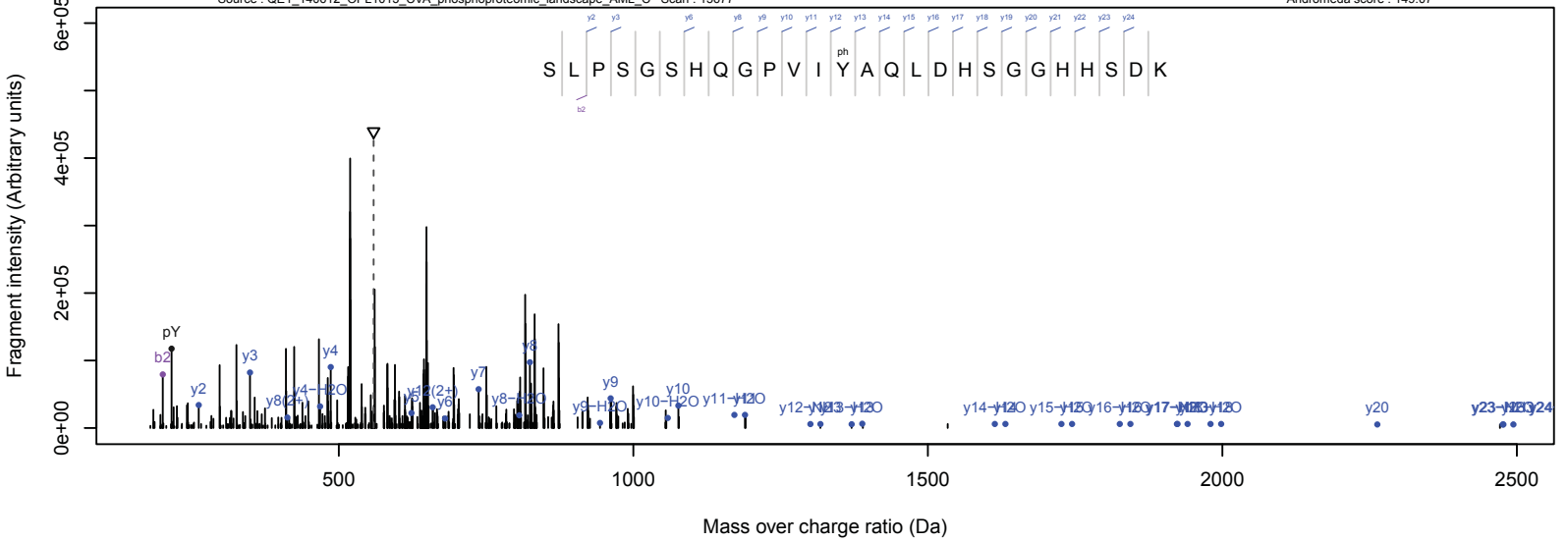
Andromeda score : 149.07



SLPSGSHQGPVIY(ph)AQLDHSGGHHSK-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15677

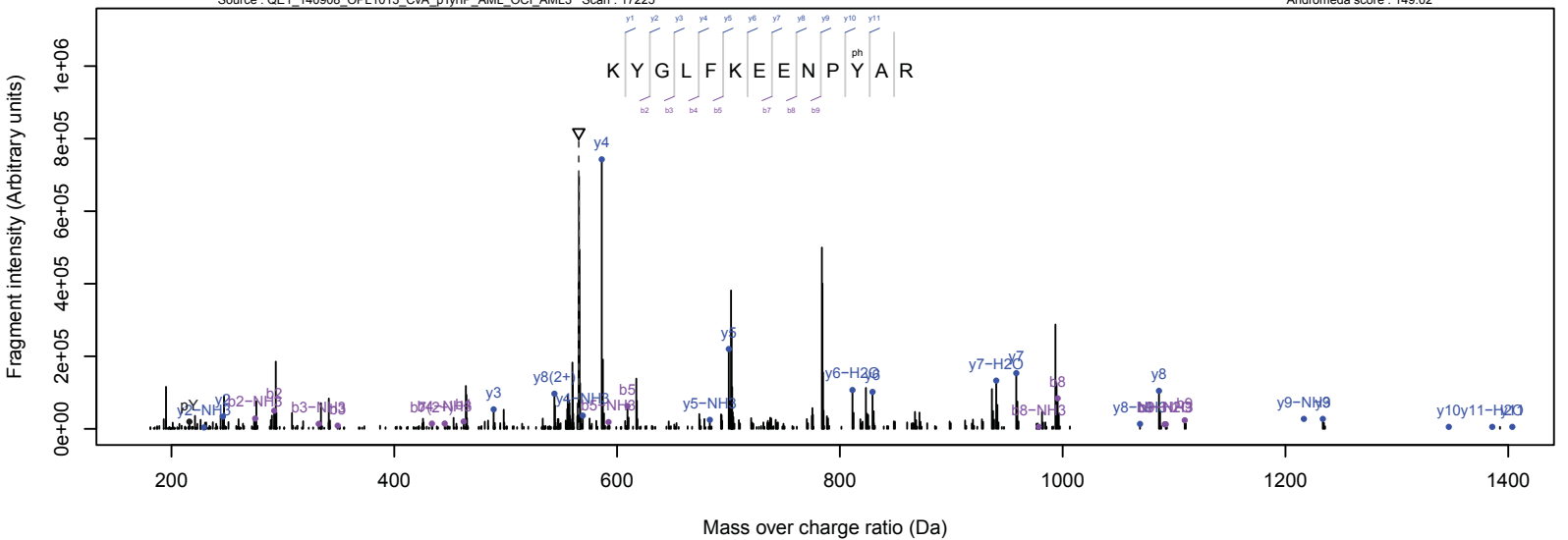
Andromeda score : 149.07



KYGLFKEENPYAR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 17225

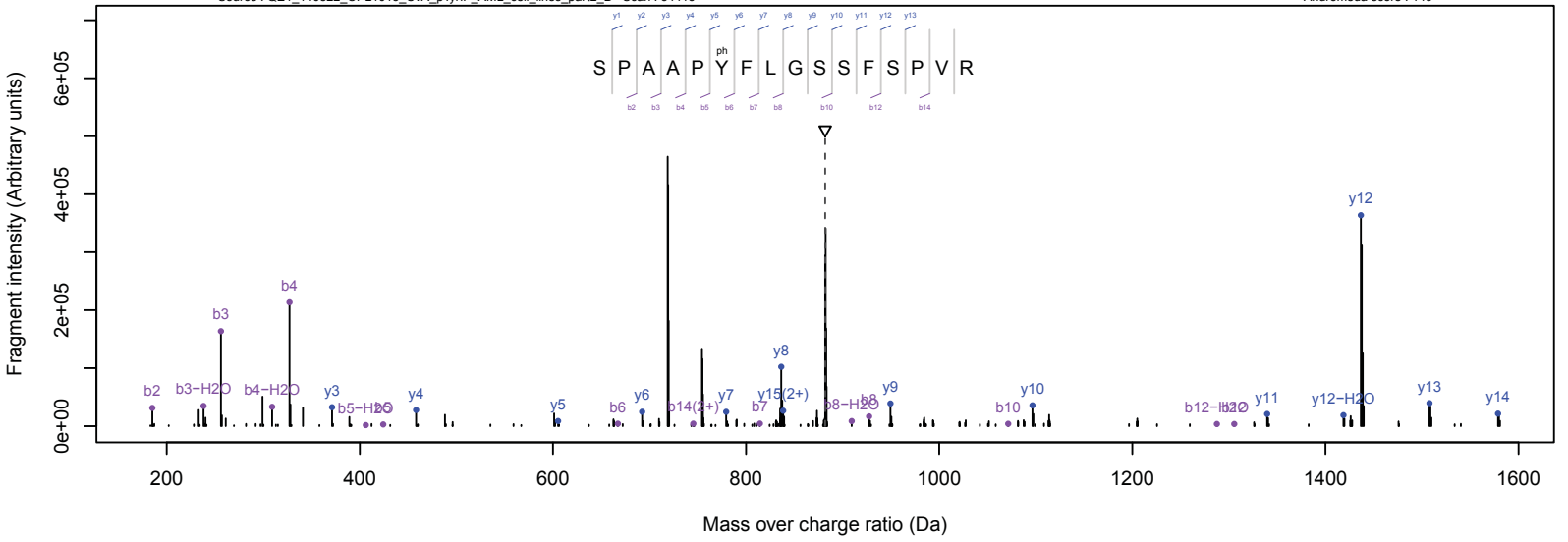
Andromeda score : 149.02



SPAAPY(ph)FLGSSFSFSPVR-2+

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 34415

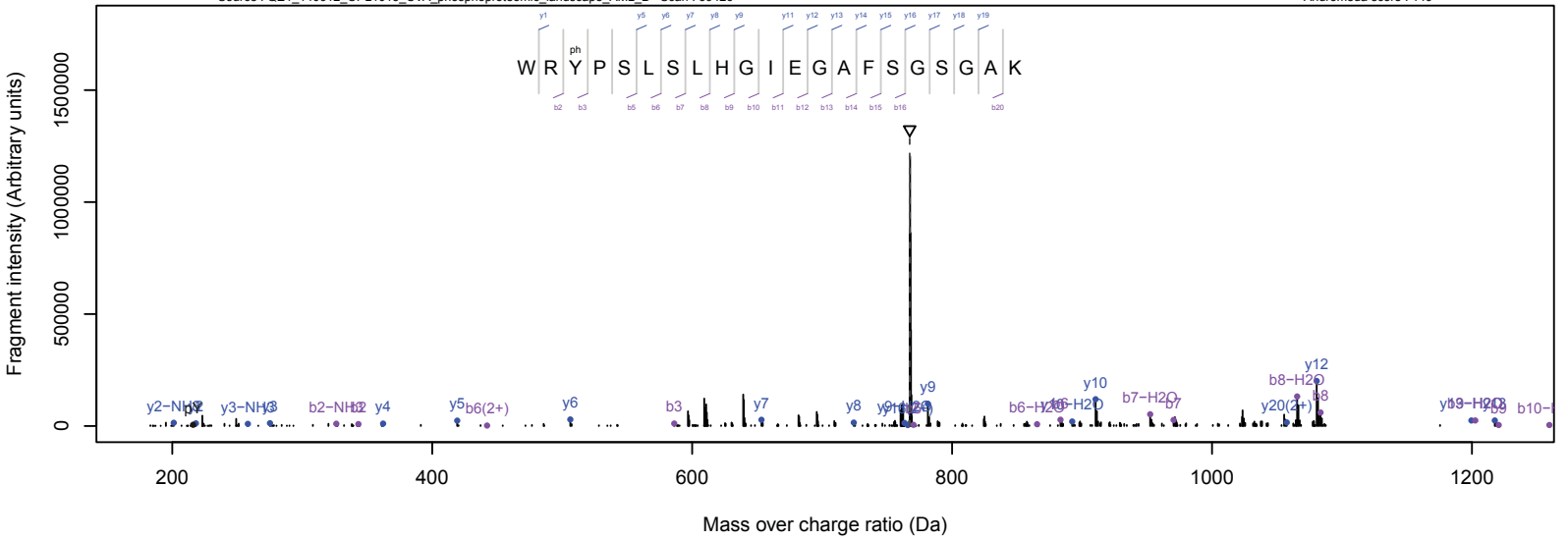
Andromeda score: 149



WRY(ph)PSLSLHGIEGAFSGSGAK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 39426

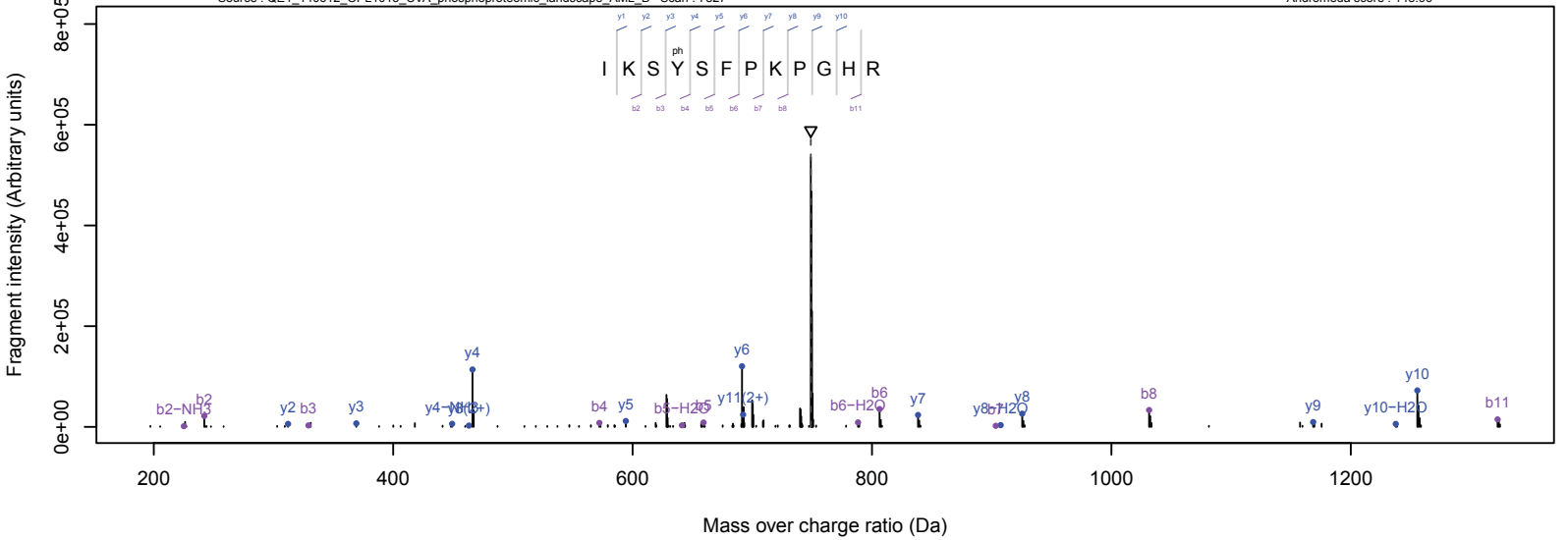
Andromeda score: 149



IKSY(ph)SFPKPGHR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 7827

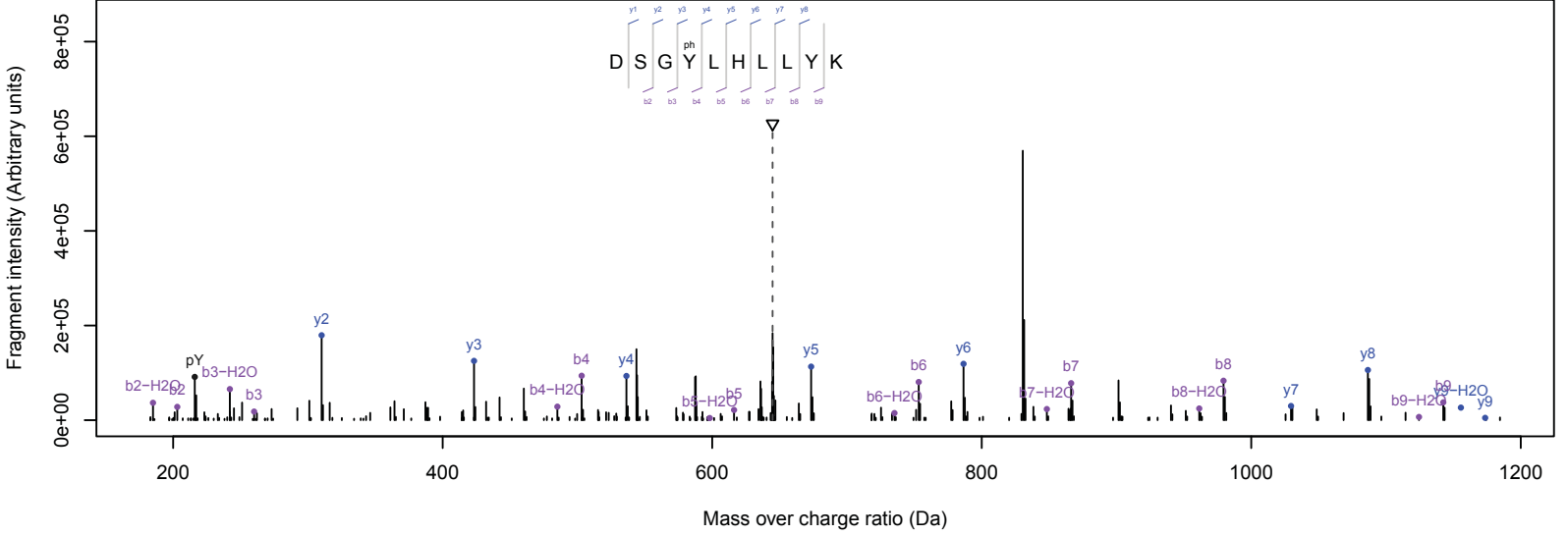
Andromeda score: 148.96



DSGY(ph)LHLLYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 29235

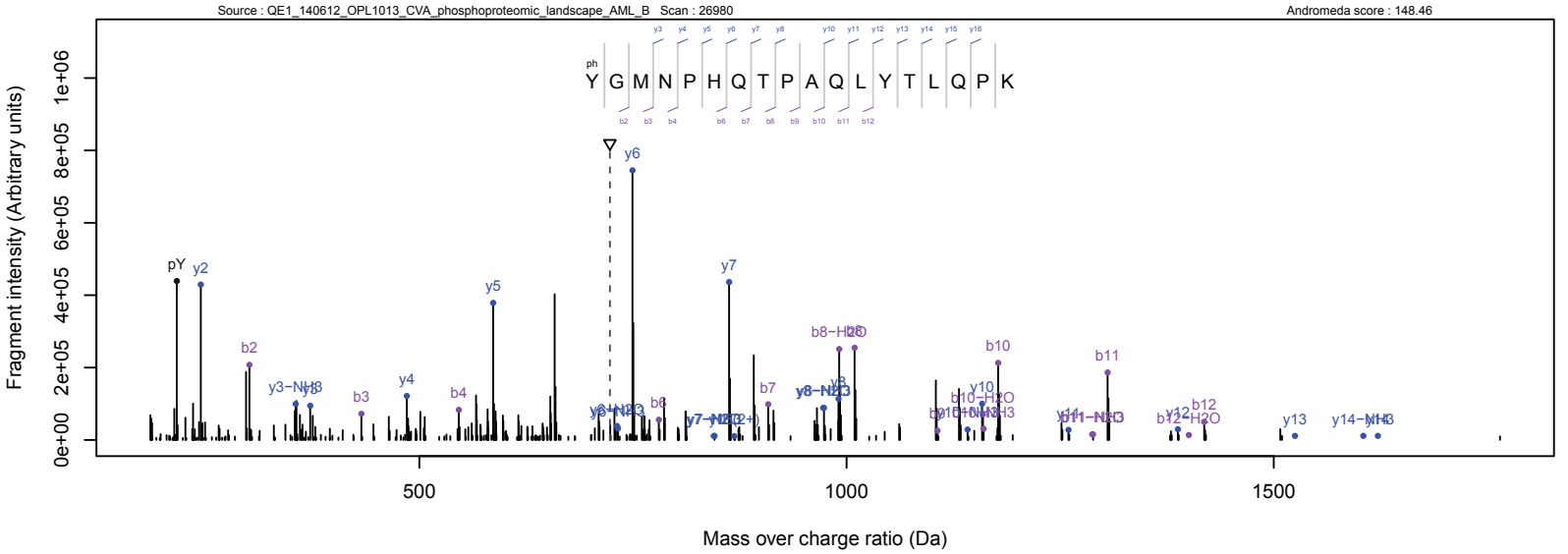
Andromeda score : 148.91



Y(ph)GMNPHQTPAQLYTLQPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26980

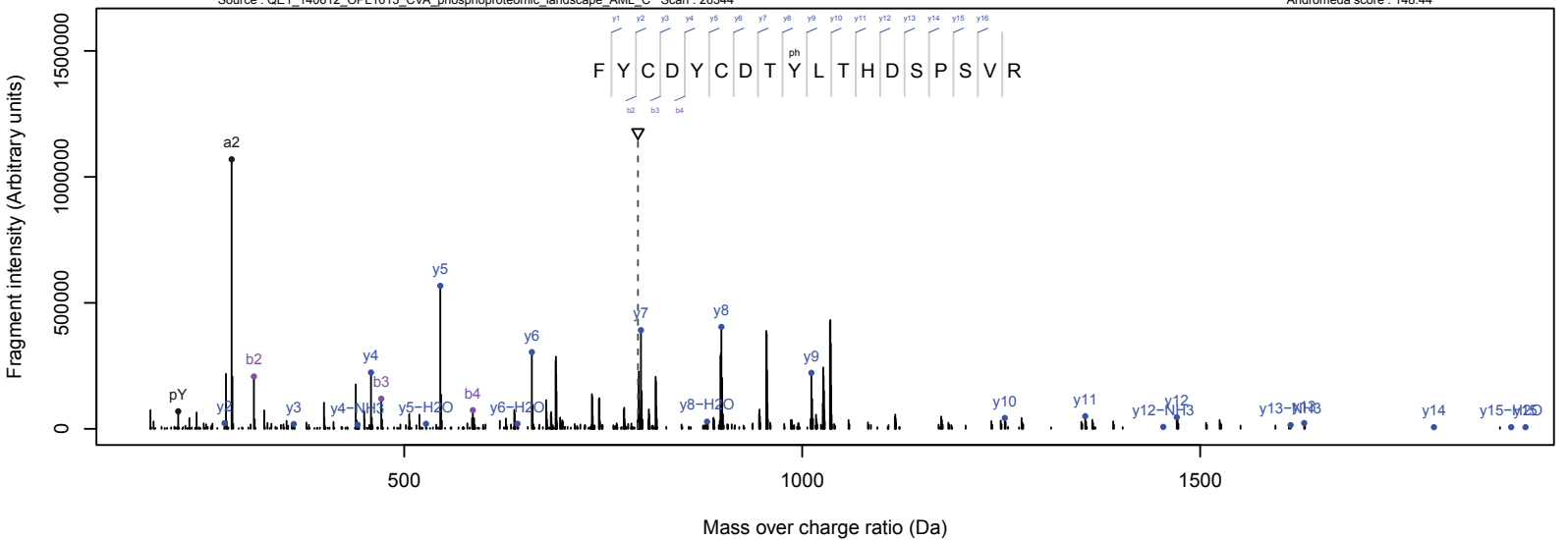
Andromeda score : 148.46



FYCDYCDTY(ph)LTHDPSVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26344

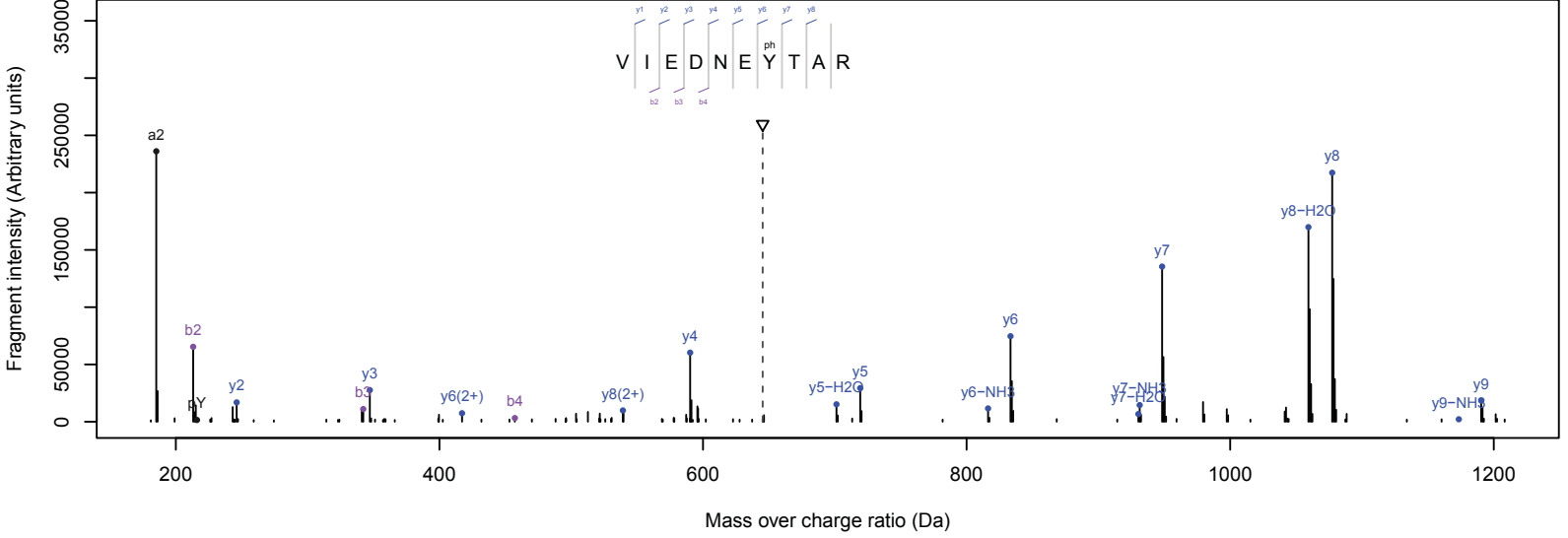
Andromeda score : 148.44



**VIEDNEY(ph)TAR-2+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 7741

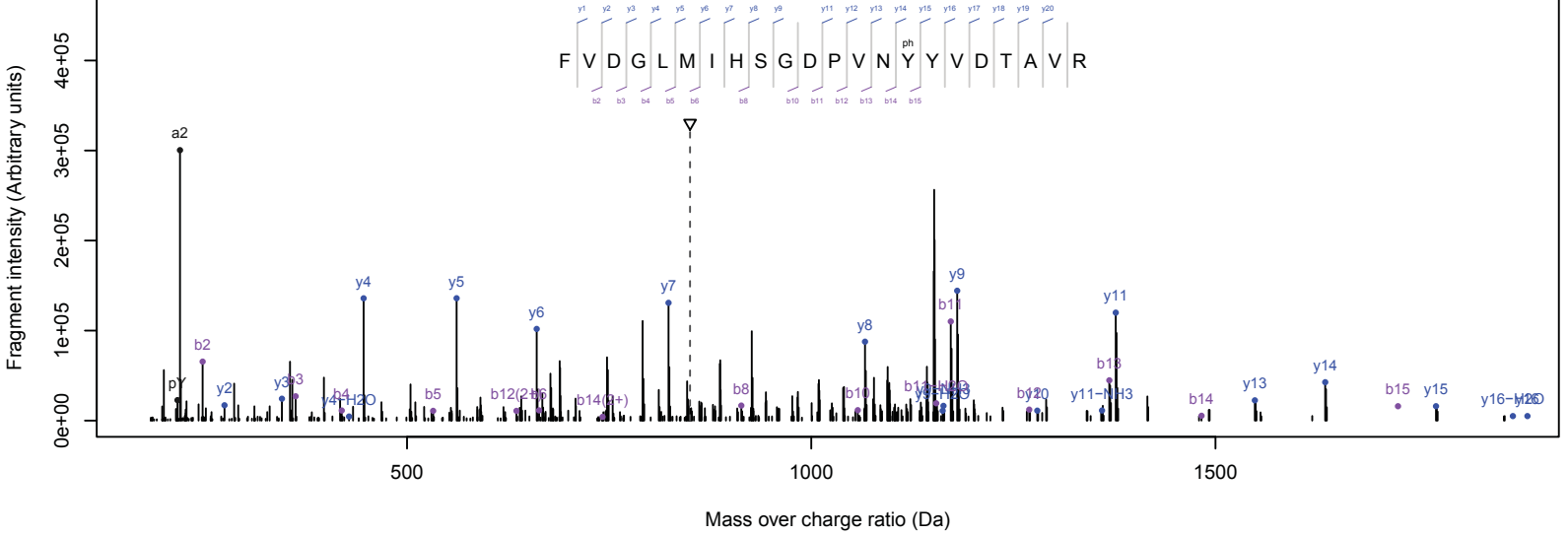
Andromeda score : 148.32



**FVDGLMIHSGDPVNY(ph)YVDTAVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45026

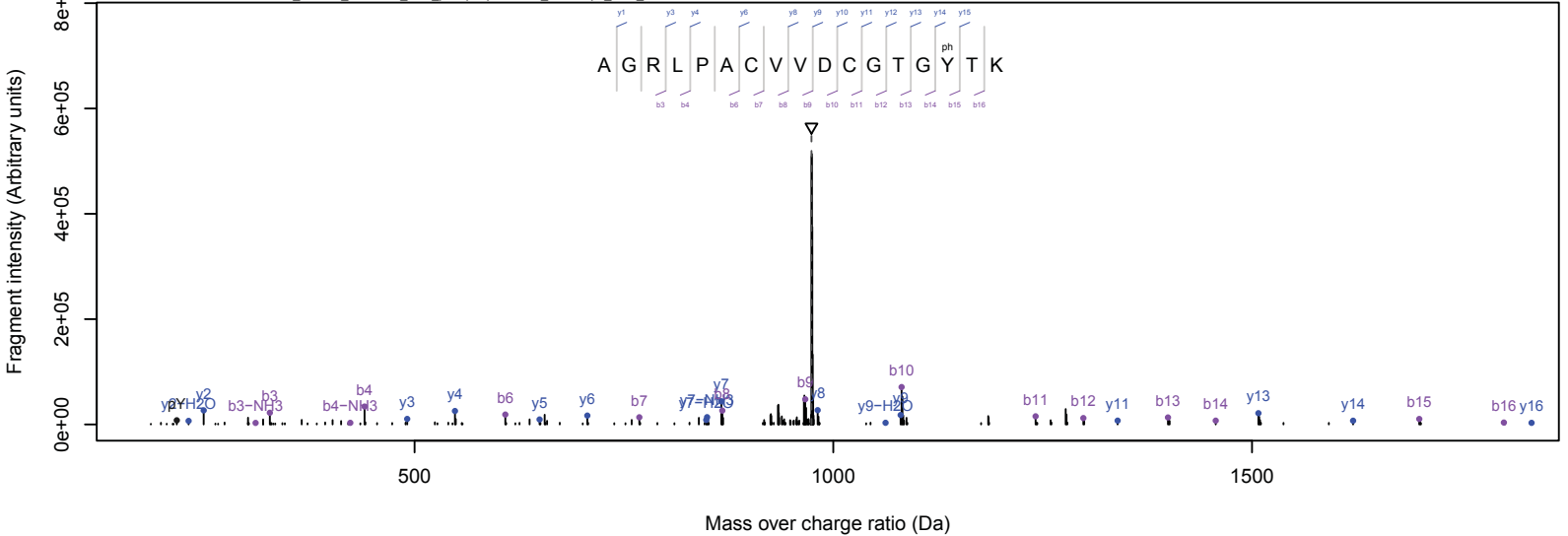
Andromeda score : 148.29



**(ac)AGRLPACVVDCGTGY(ph)TK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27552

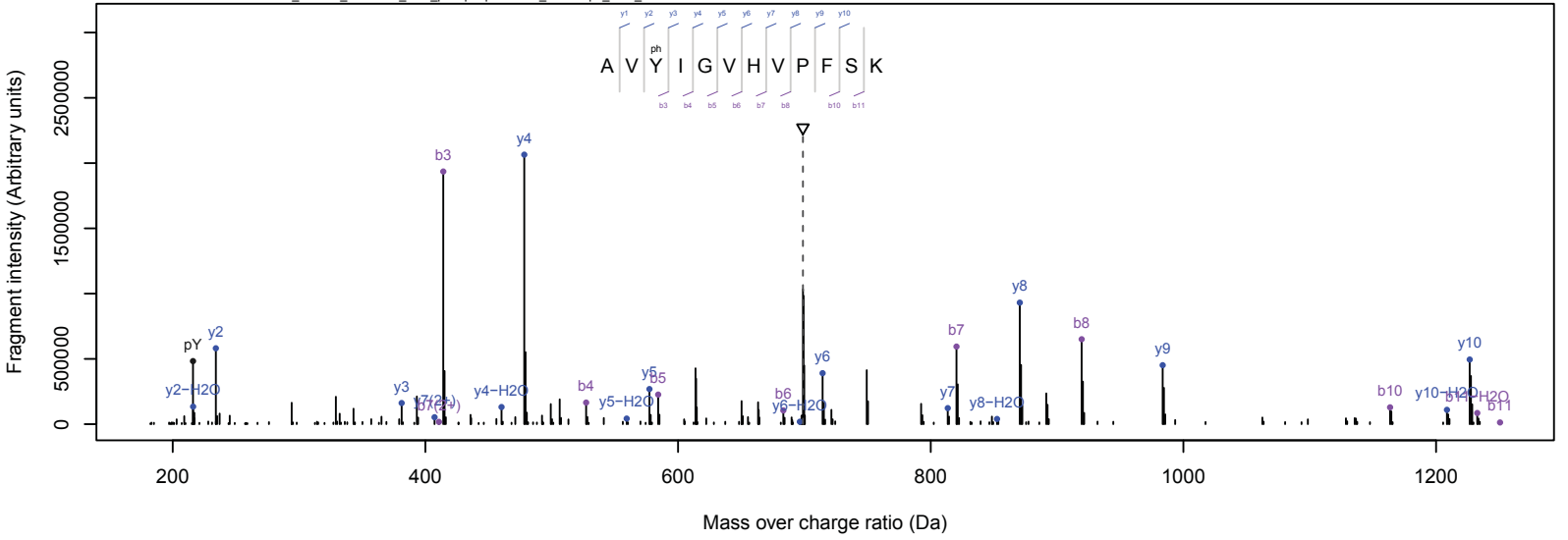
Andromeda score : 148.26



AVY(ph)IGVHVPF<sup>ph</sup>SK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34985

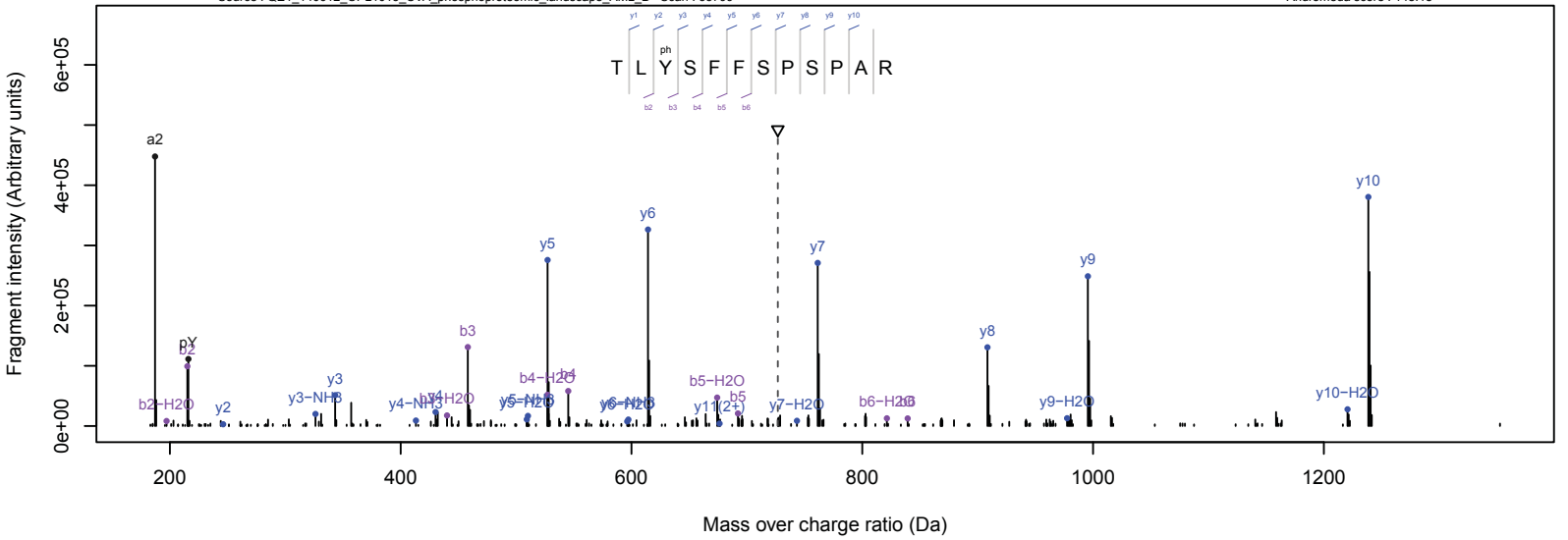
Andromeda score : 148.18



TLY(ph)SFFSPSPAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35799

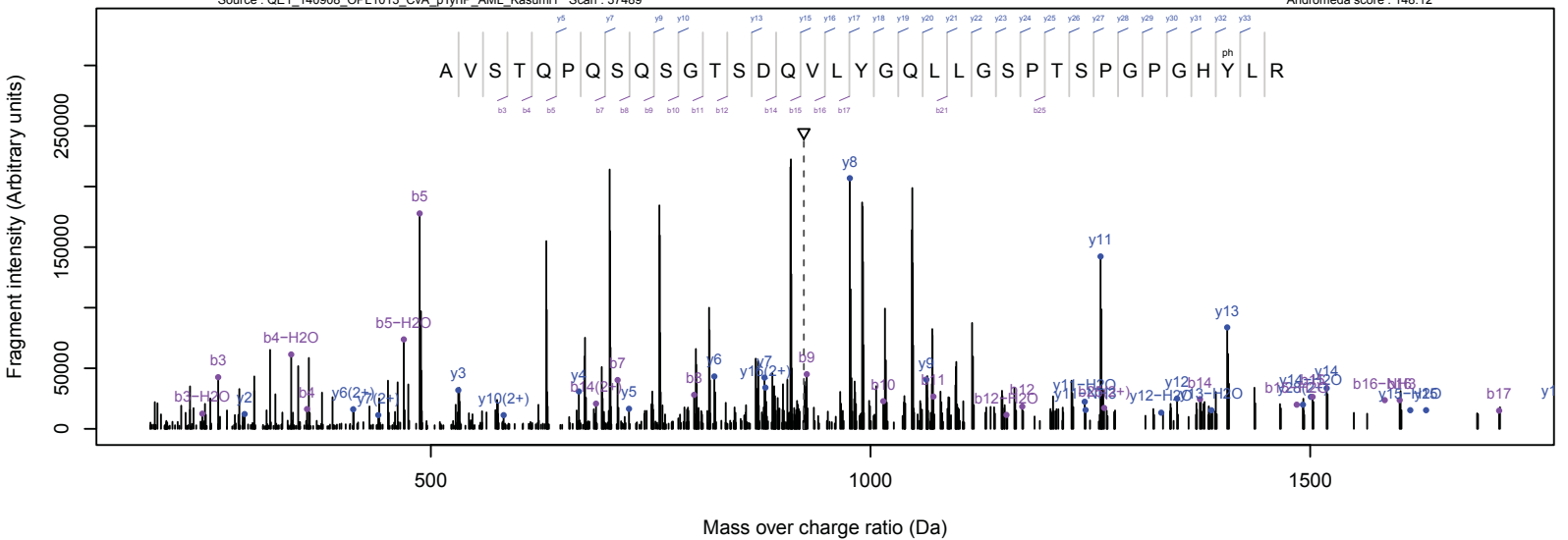
Andromeda score : 148.18



AVSTQPSQSGTSDQVLYGQLLGSPTSPGPGHY(ph)LR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 37489

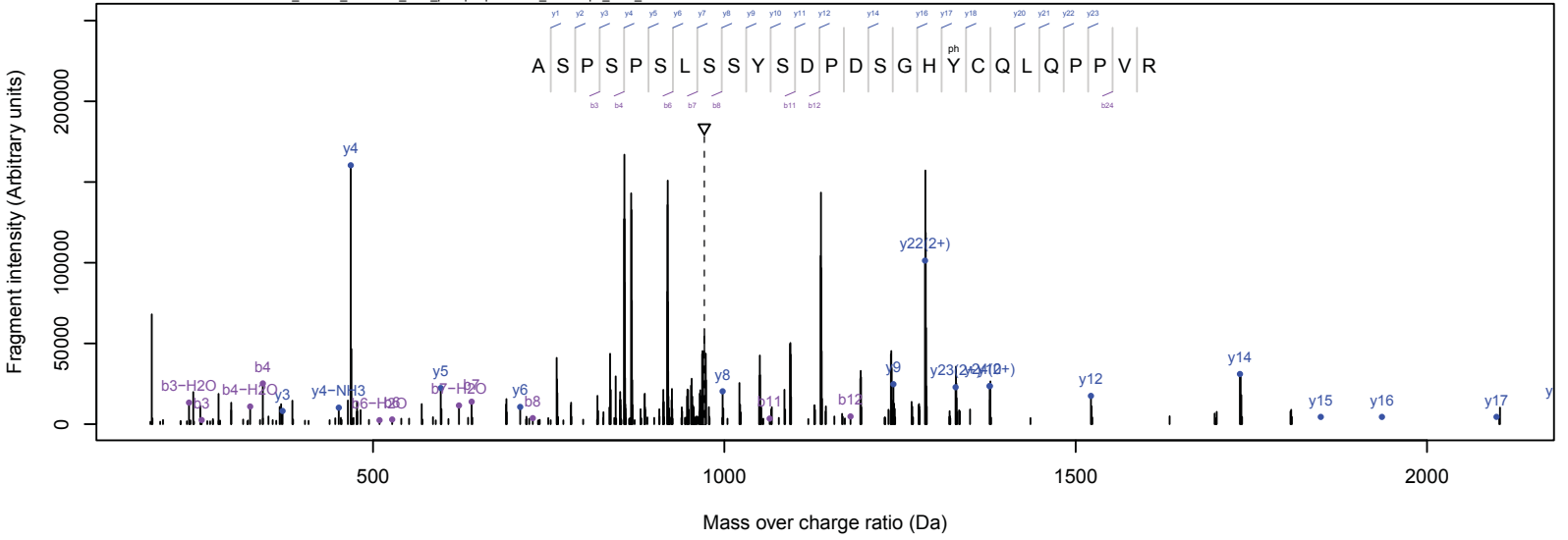
Andromeda score : 148.12



ASPSLSYSDPSGHY(ph)CQLQPPV R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27774

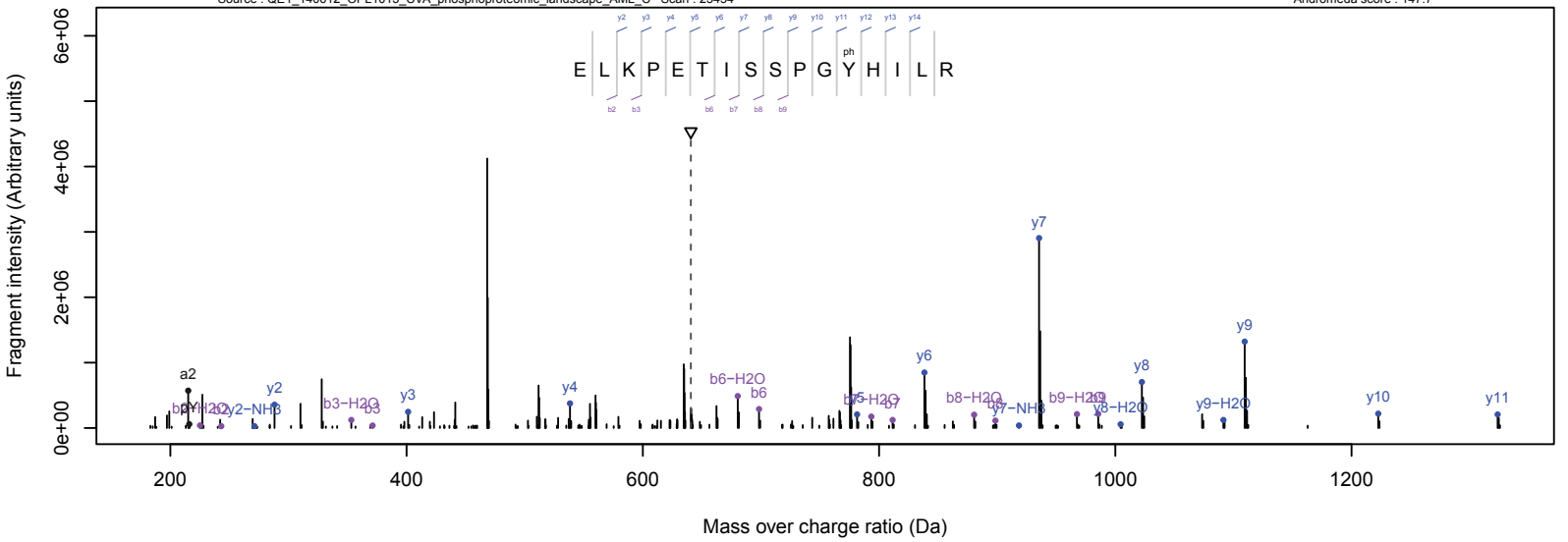
Andromeda score : 148.09



ELKPETISSPGY(ph)HILR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25454

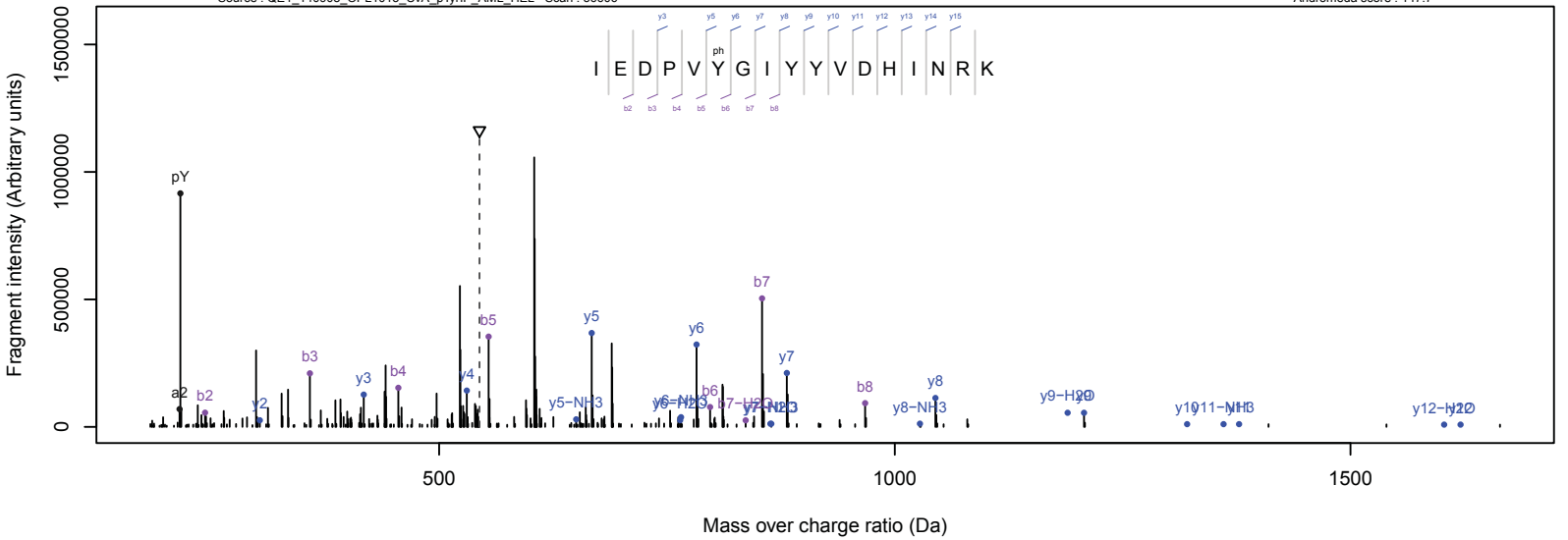
Andromeda score : 147.7



IEDPVYGIYYVDHINR K-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 36606

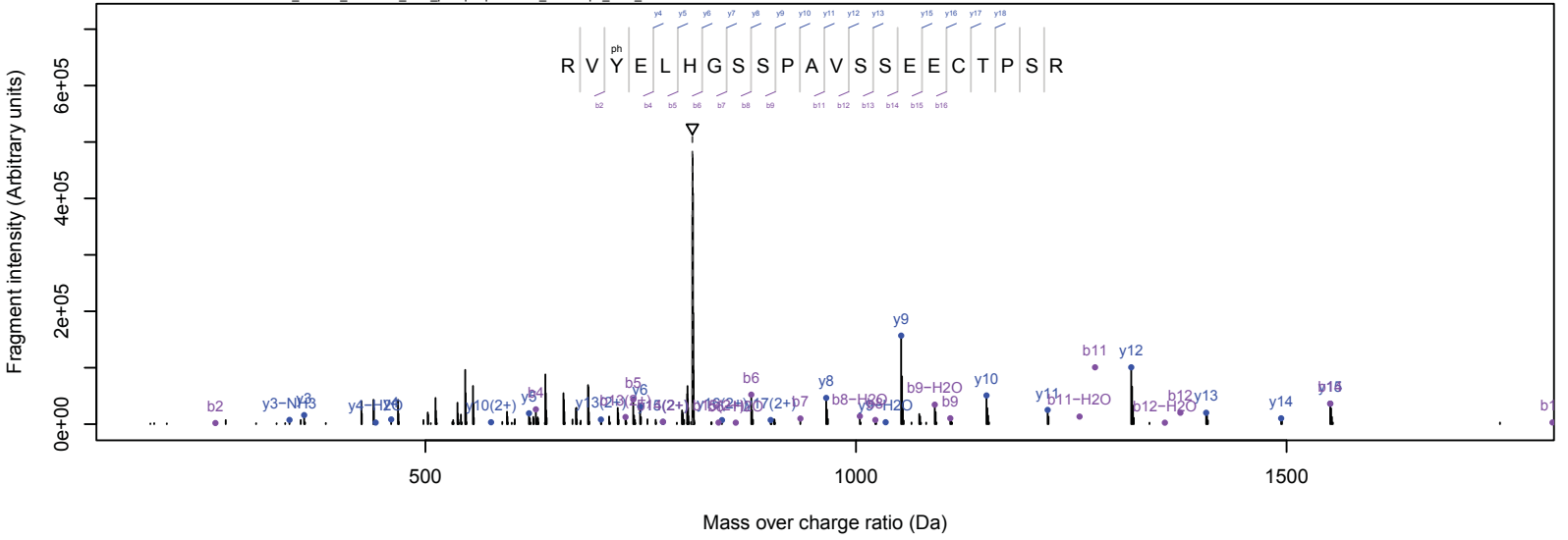
Andromeda score : 147.7



**RVY(ph)ELHGSSPAVSSSEECTPSR**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12026

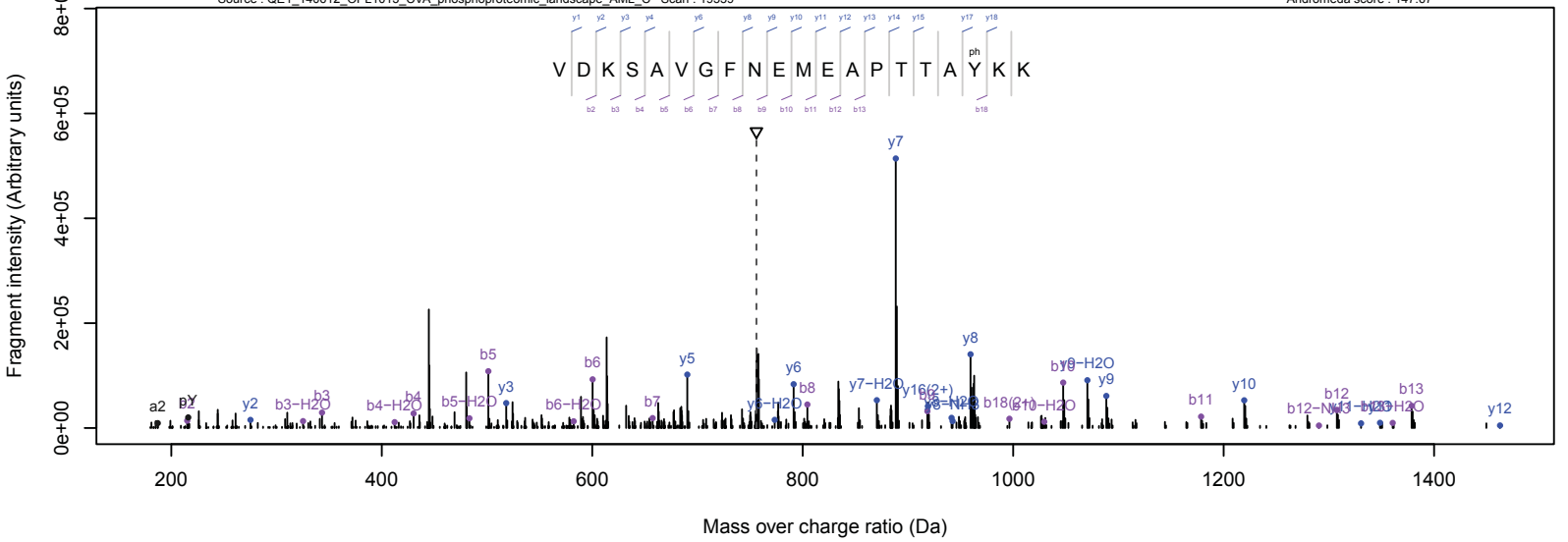
Andromeda score : 147.67



**VDKSAVGFNEMEAPTTAYKK**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19559

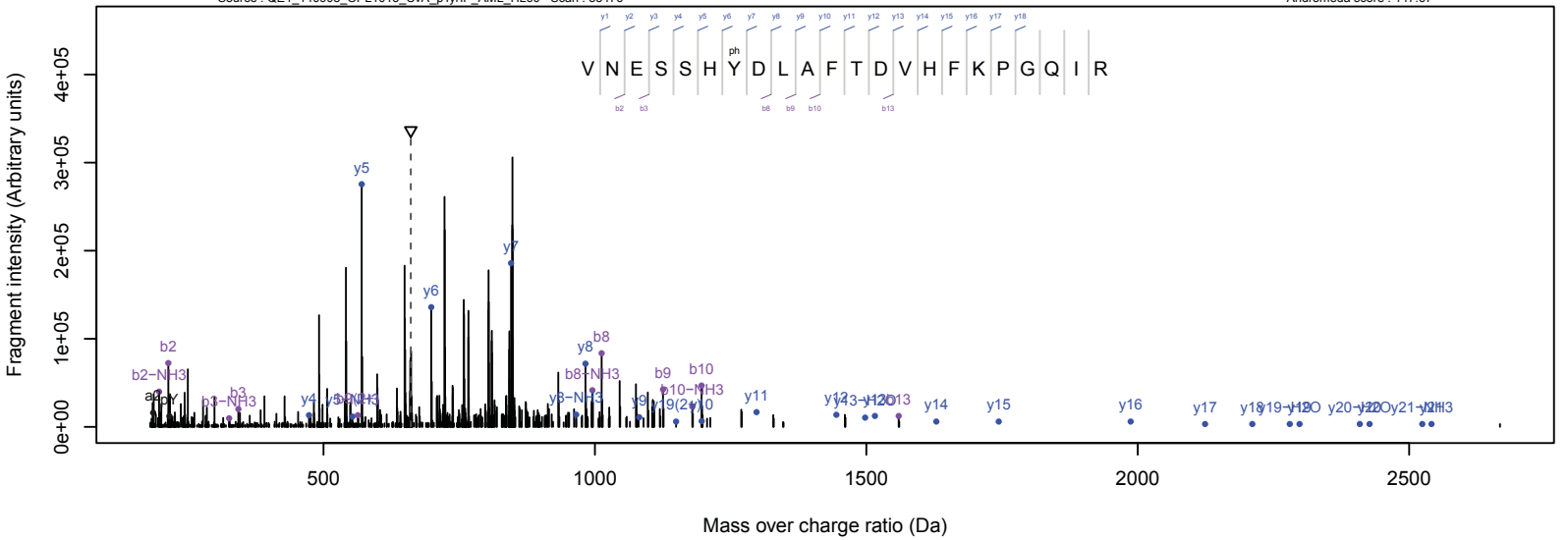
Andromeda score : 147.67



**VNESSHY(ph)DLAFTDVHFKPGQIR**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 33476

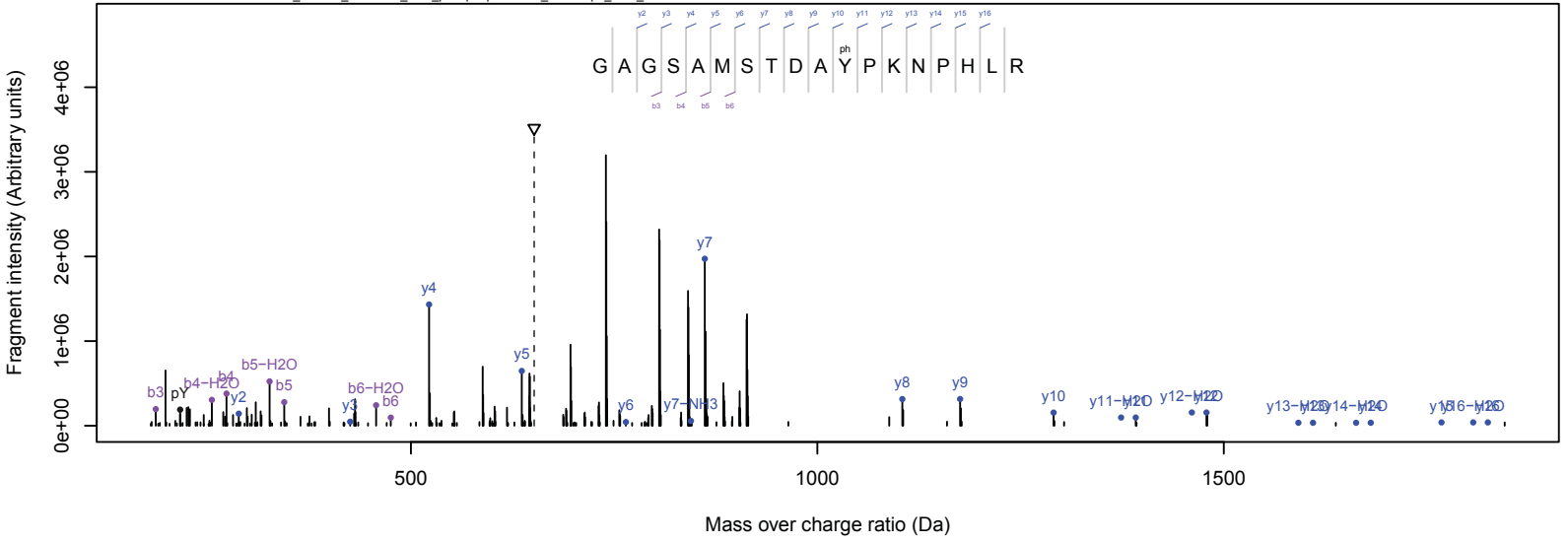
Andromeda score : 147.67



GAGSAMSTDAY(ph)PKNPHLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10971

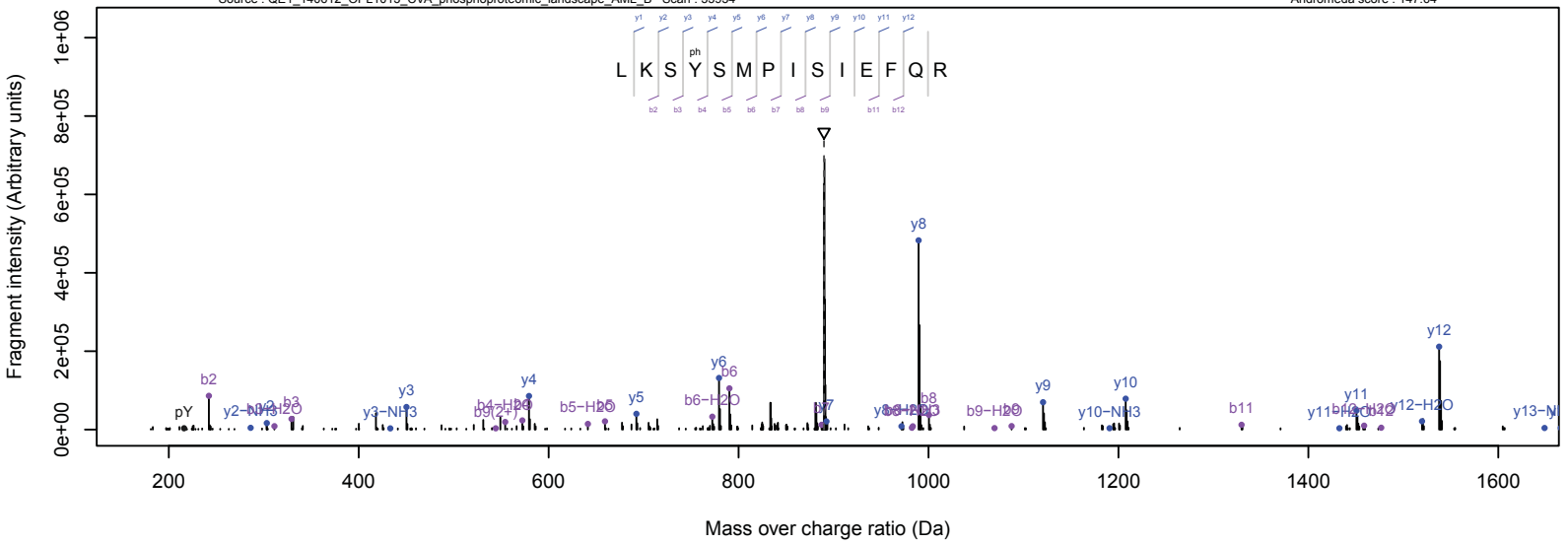
Andromeda score: 147.64



LKSY(ph)SMPISIEFQR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 33934

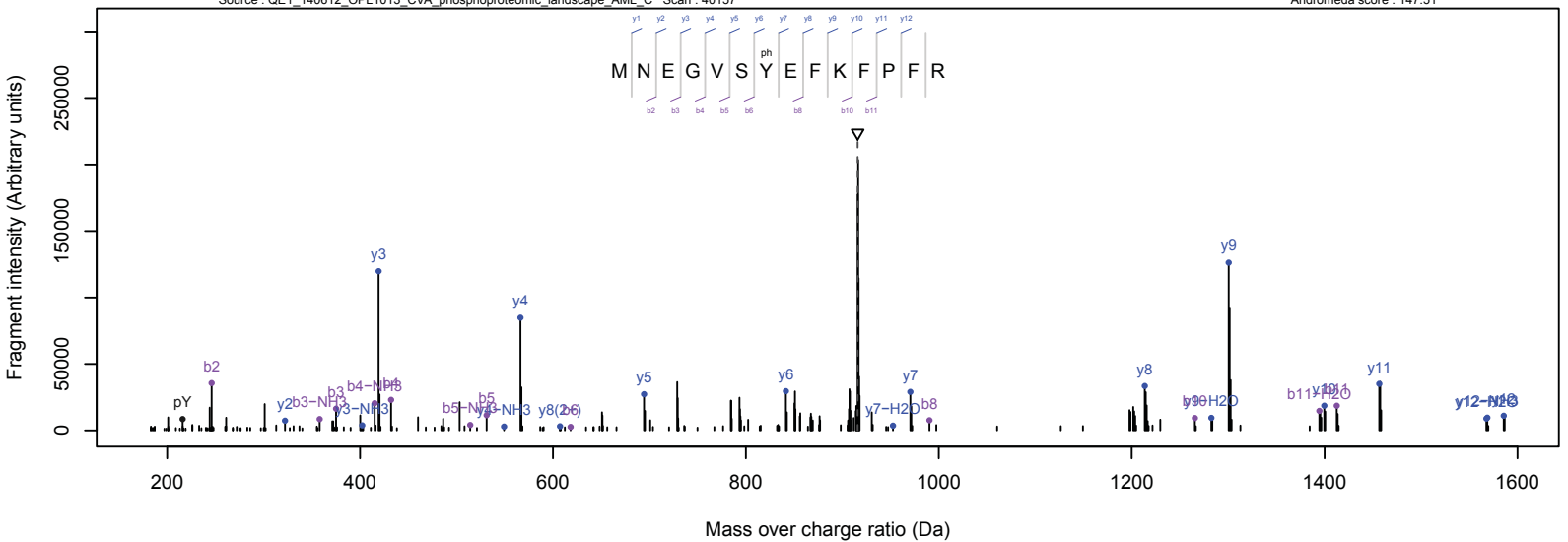
Andromeda score: 147.64



MNEGVSY(ph)EFKFFR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 40157

Andromeda score: 147.51

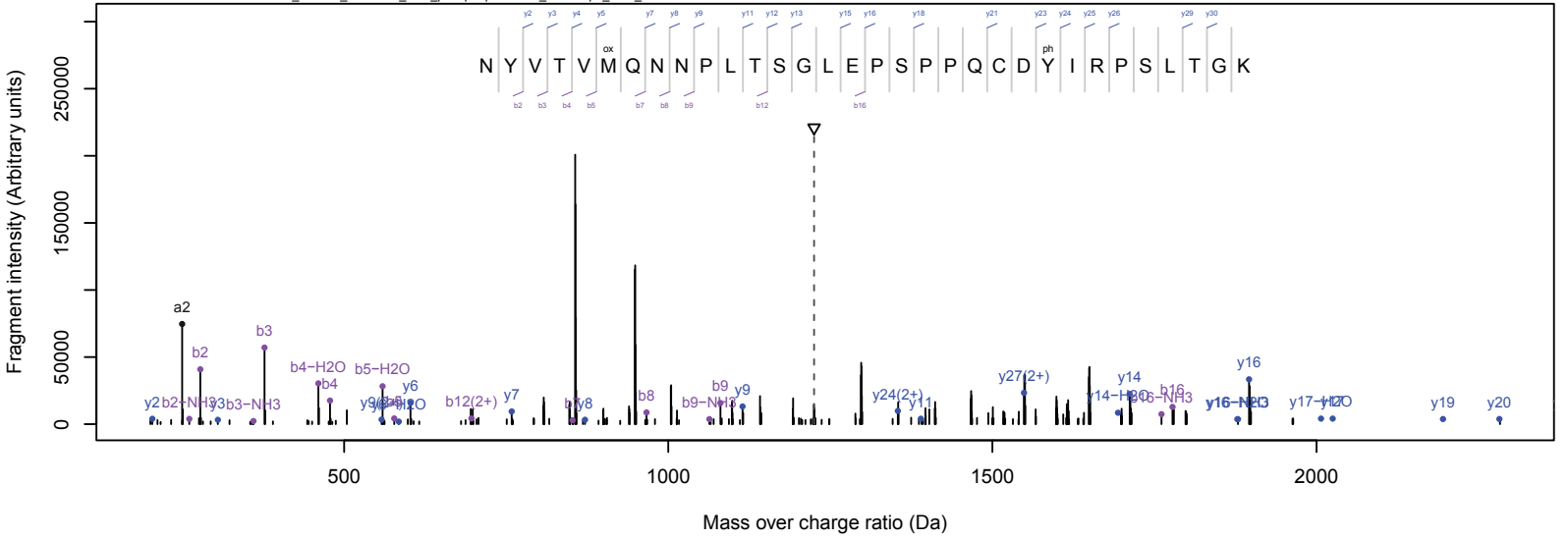




**NYVTVM(ox)QNNPLTSGLEPSPQCDY(ph)IRPSLTGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39046

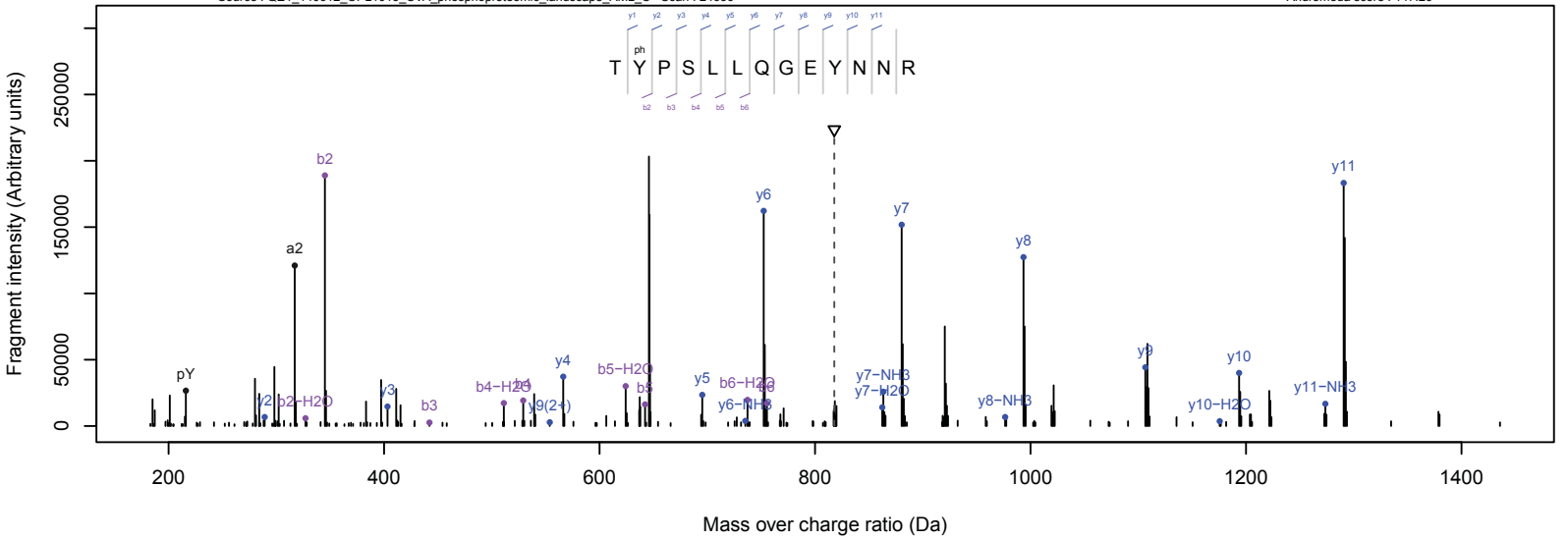
Andromeda score : 147.42



**TY(ph)PSLLQGEYNNR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24656

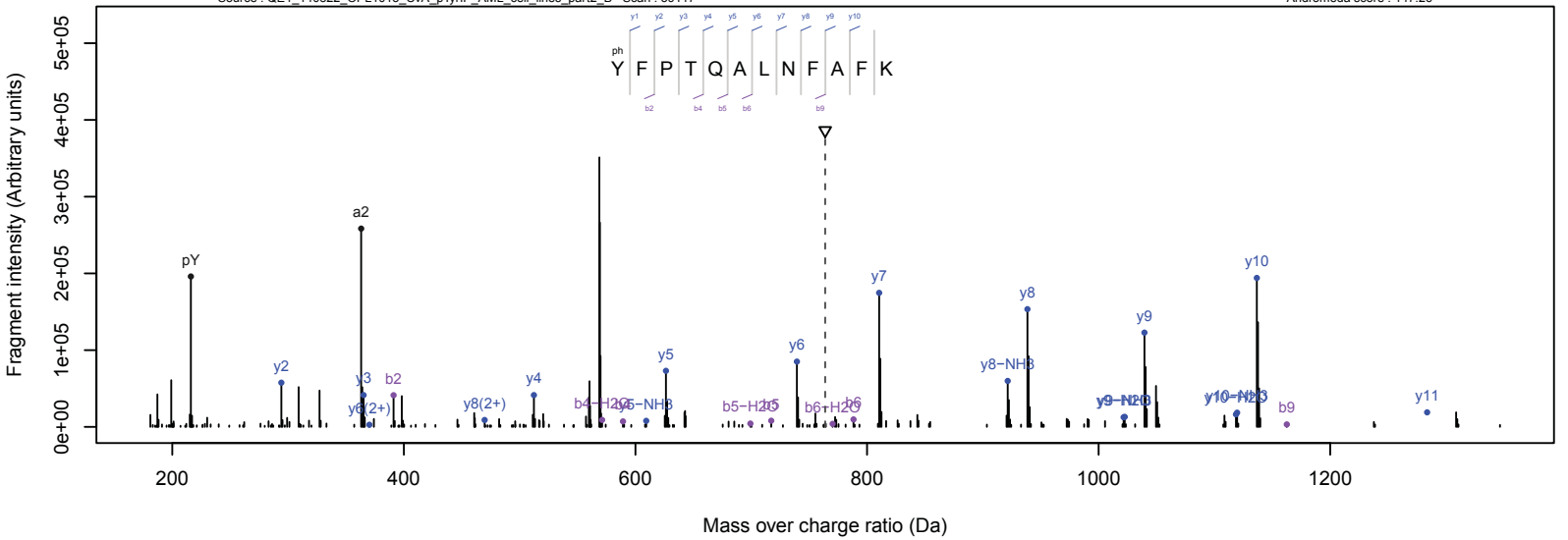
Andromeda score : 147.26



**Y(ph)FPTQALNFAFK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 39117

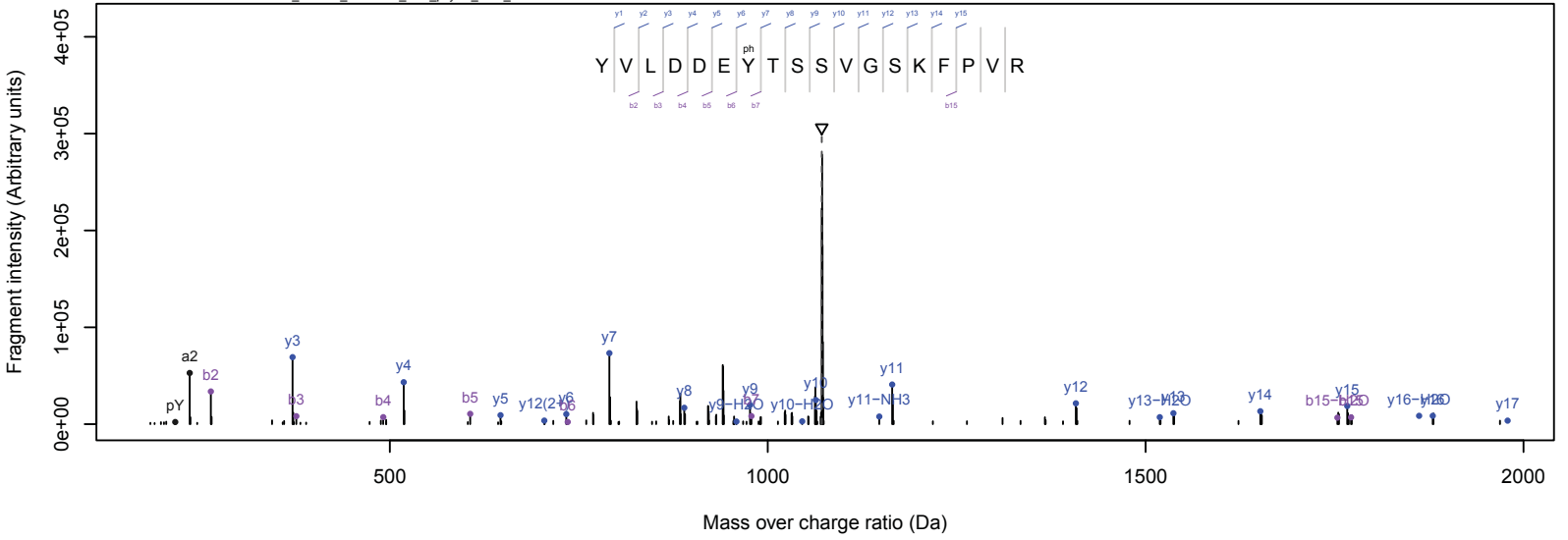
Andromeda score : 147.26



**YVLDDDEY(ph)TSSVGSKFPVR-2+**

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan: 33302

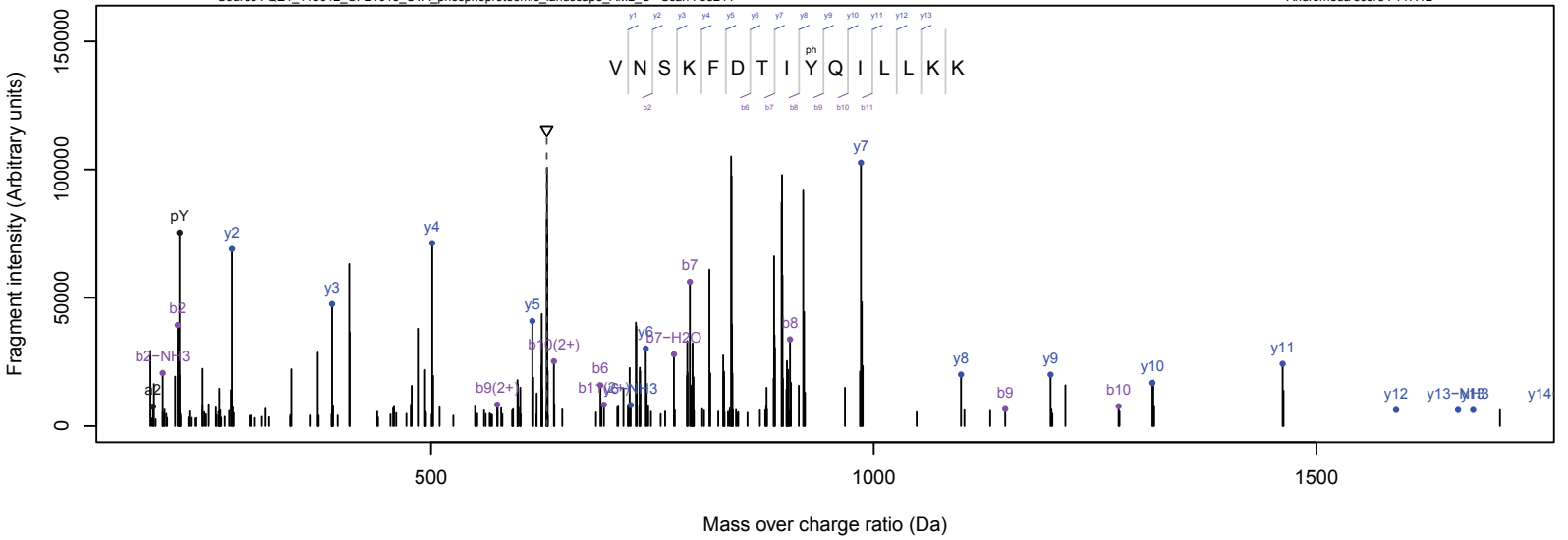
Andromeda score: 147.17



**VNSKFDTIY(ph)QLLKK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38244

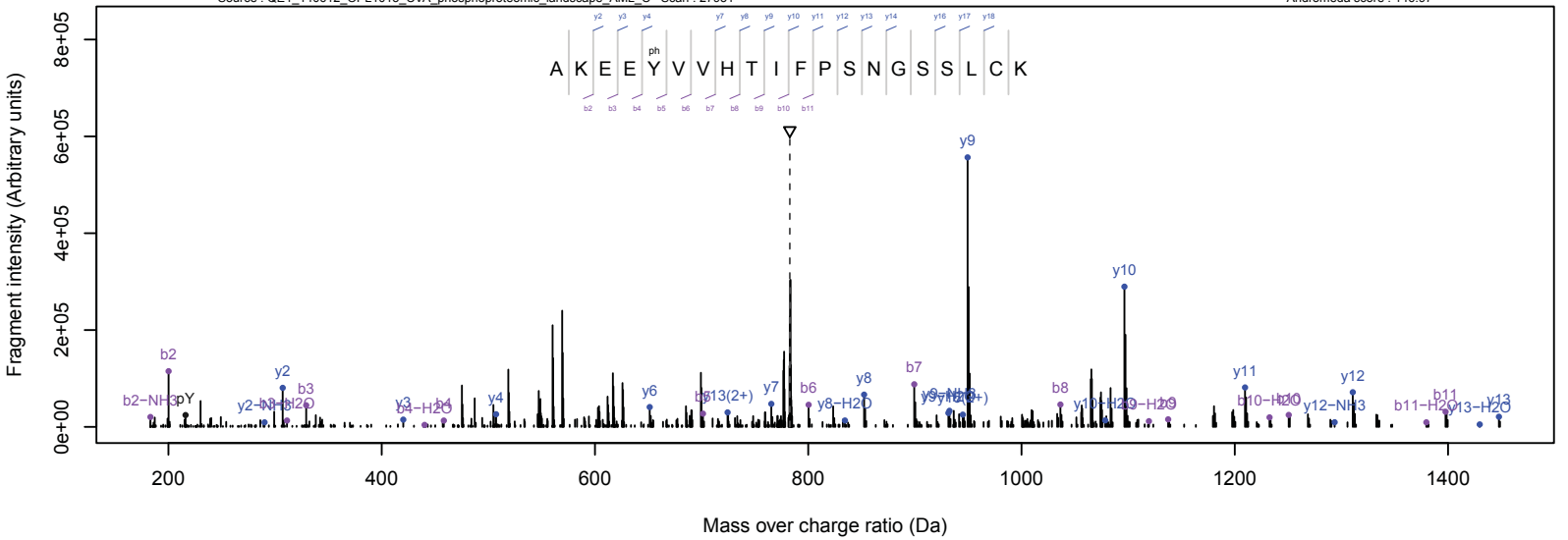
Andromeda score: 147.12



**AKEEY(ph)VVHTIFPSNGSSLCK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 27931

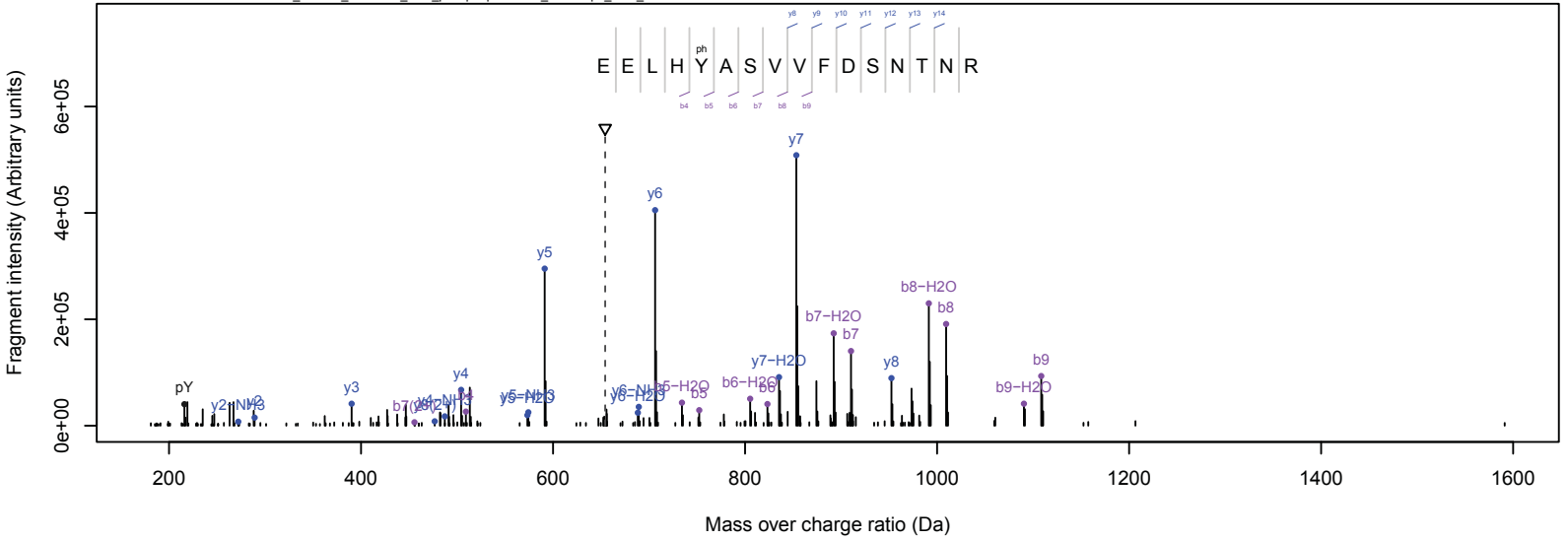
Andromeda score: 146.97



**EELHY(ph)ASVVFDSTNR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27788

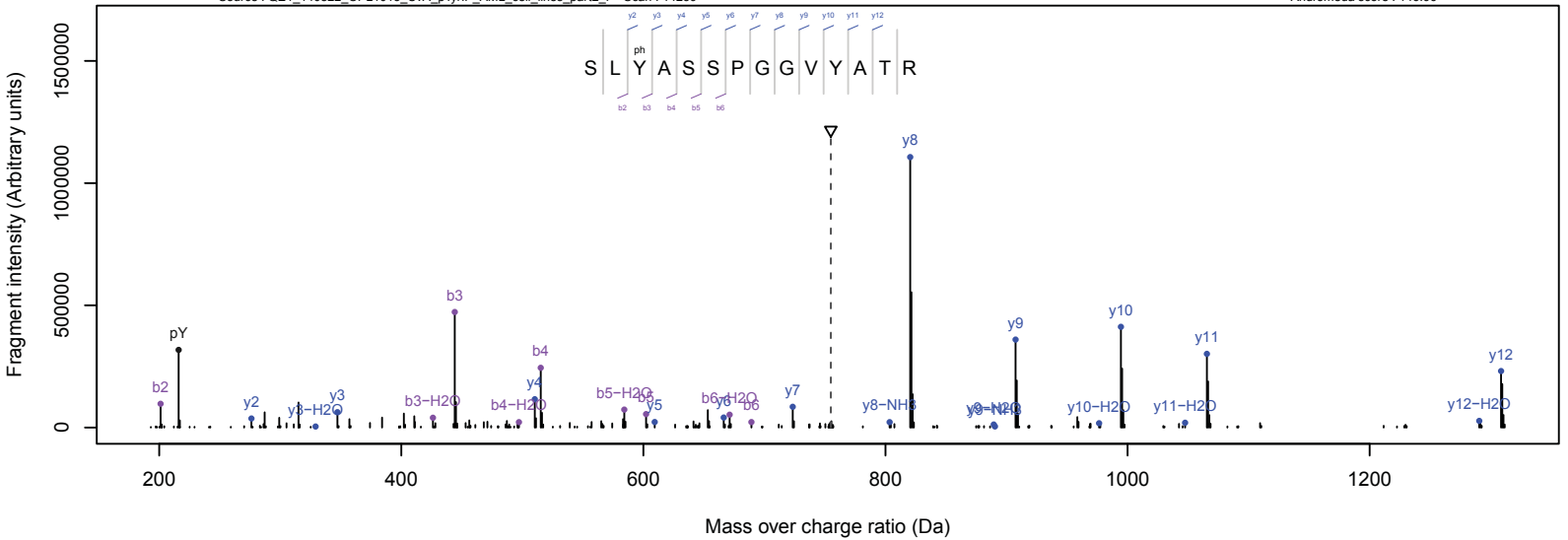
Andromeda score : 146.96



**SLY(ph)ASSPGGVYATR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 14239

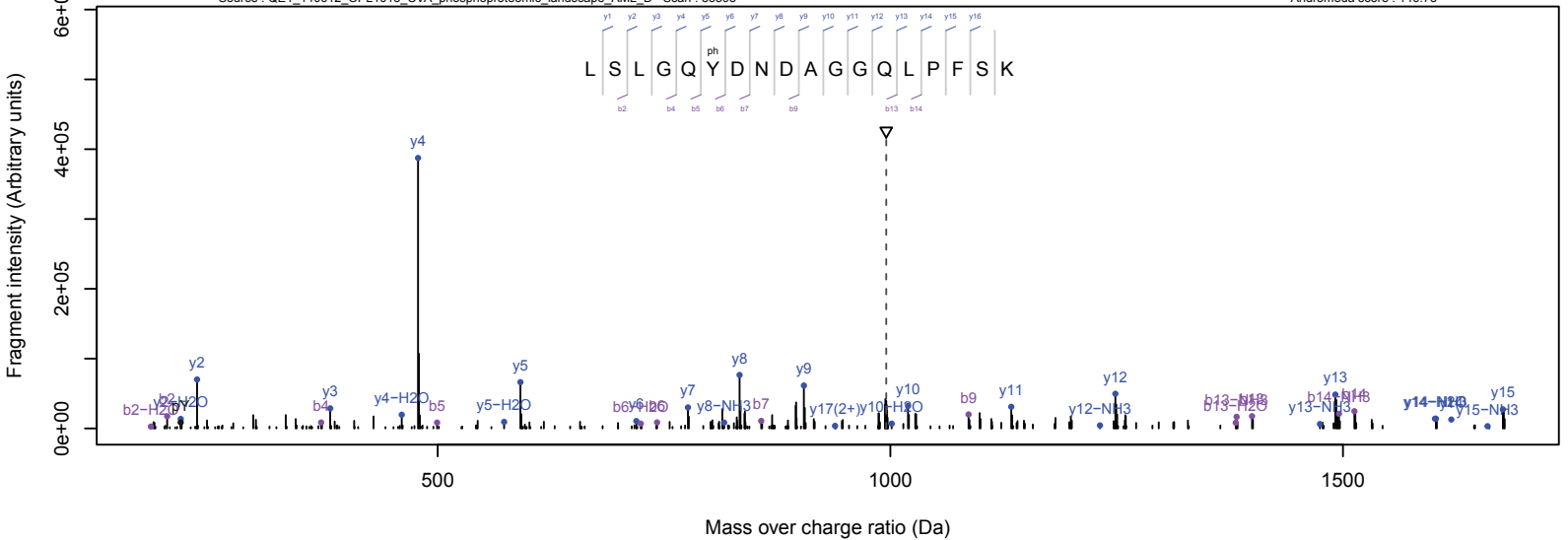
Andromeda score : 146.96



**LSLGQY(ph)DNDAGGQLPFSK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 35593

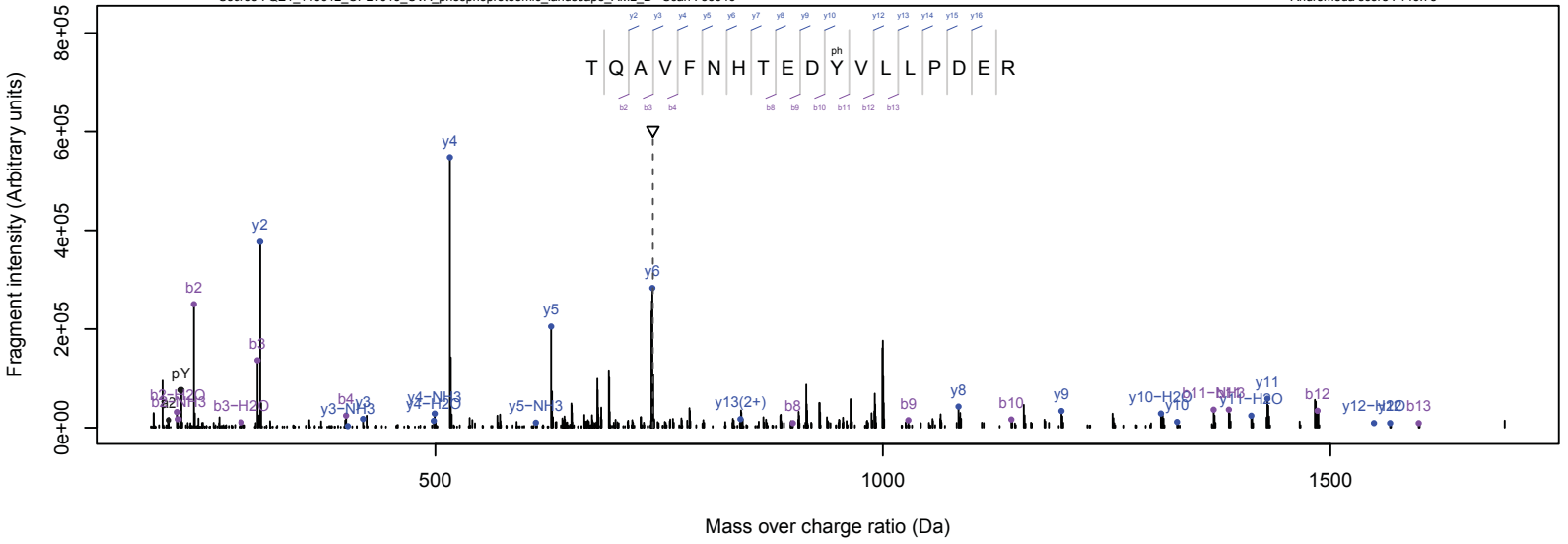
Andromeda score : 146.78



TQAVFNHTEDY(ph)VLLPDER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33046

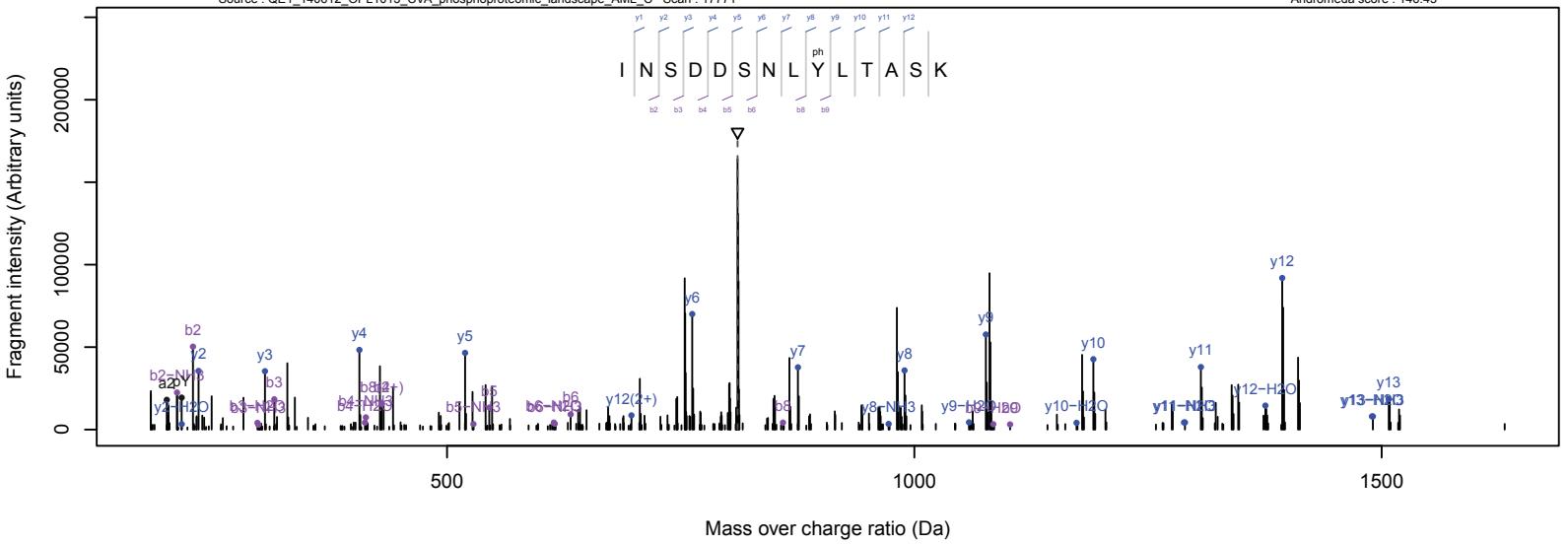
Andromeda score : 146.78



INSDDSNLY(ph)LTASK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17771

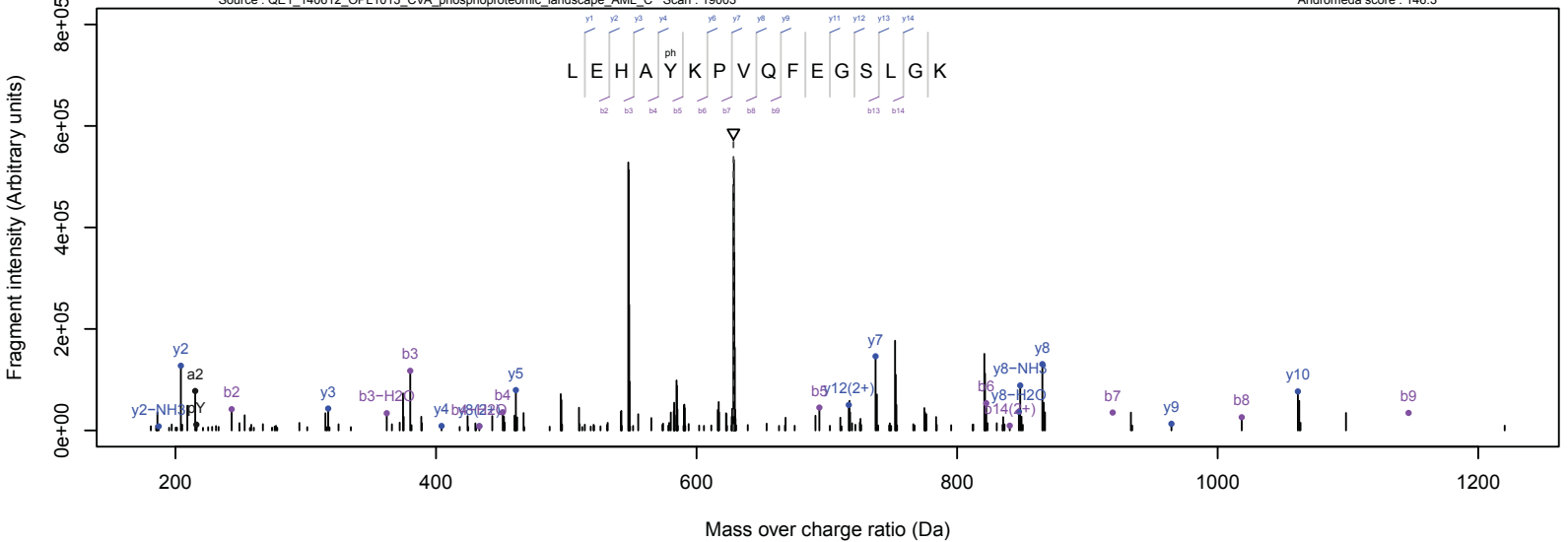
Andromeda score : 146.45



LEHAY(ph)KPVQFEGSLGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19003

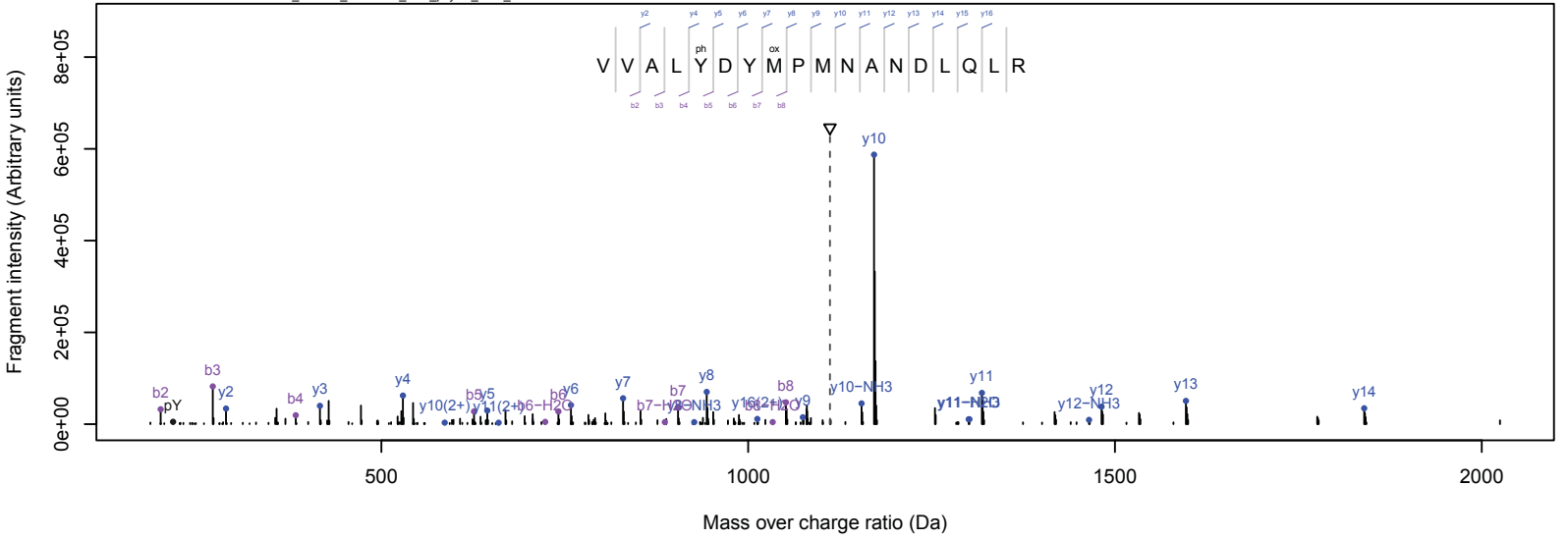
Andromeda score : 146.3



VVALY(ph)DYM(ox)PMNANDLQLR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 43016

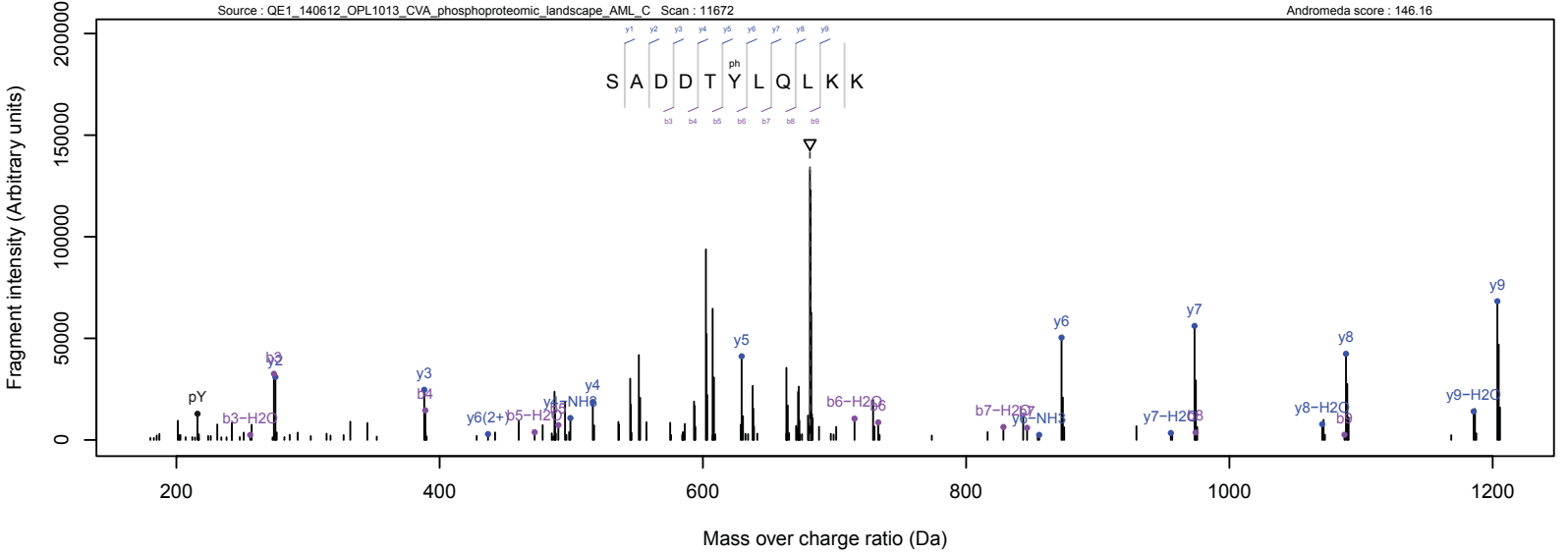
Andromeda score : 146.3



SADDTY(ph)LQLKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11672

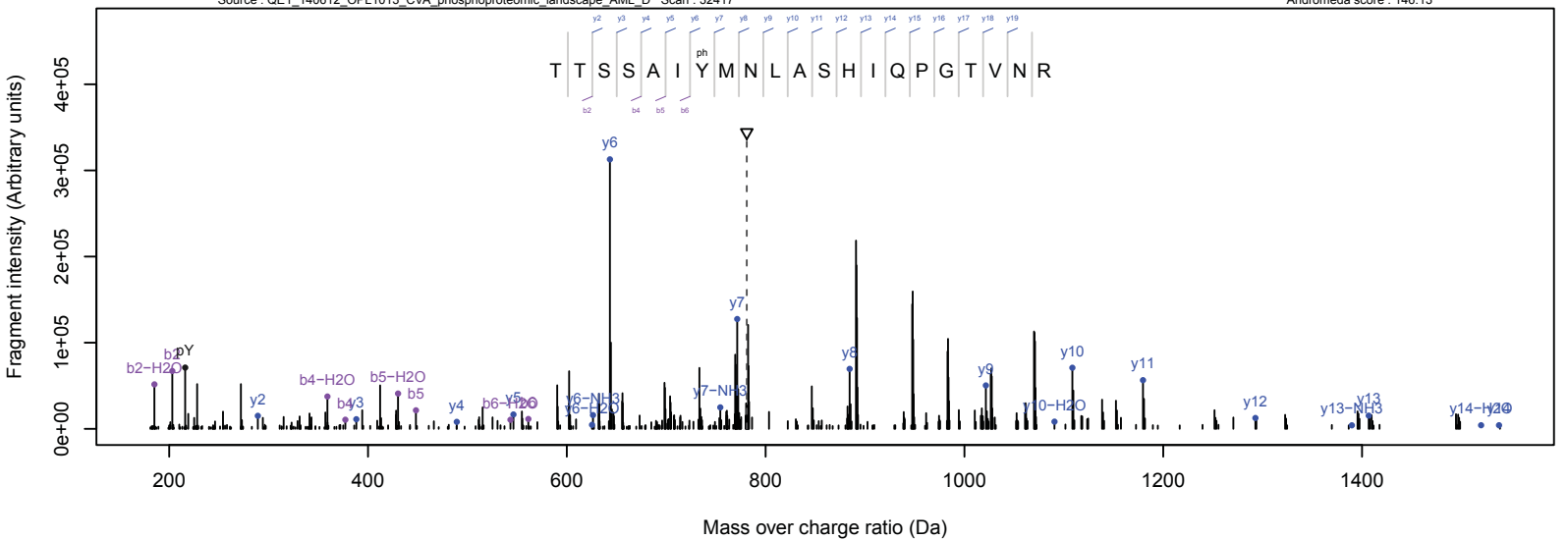
Andromeda score : 146.16



TTSSAIY(ph)MNLASHIQPGTVNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 32417

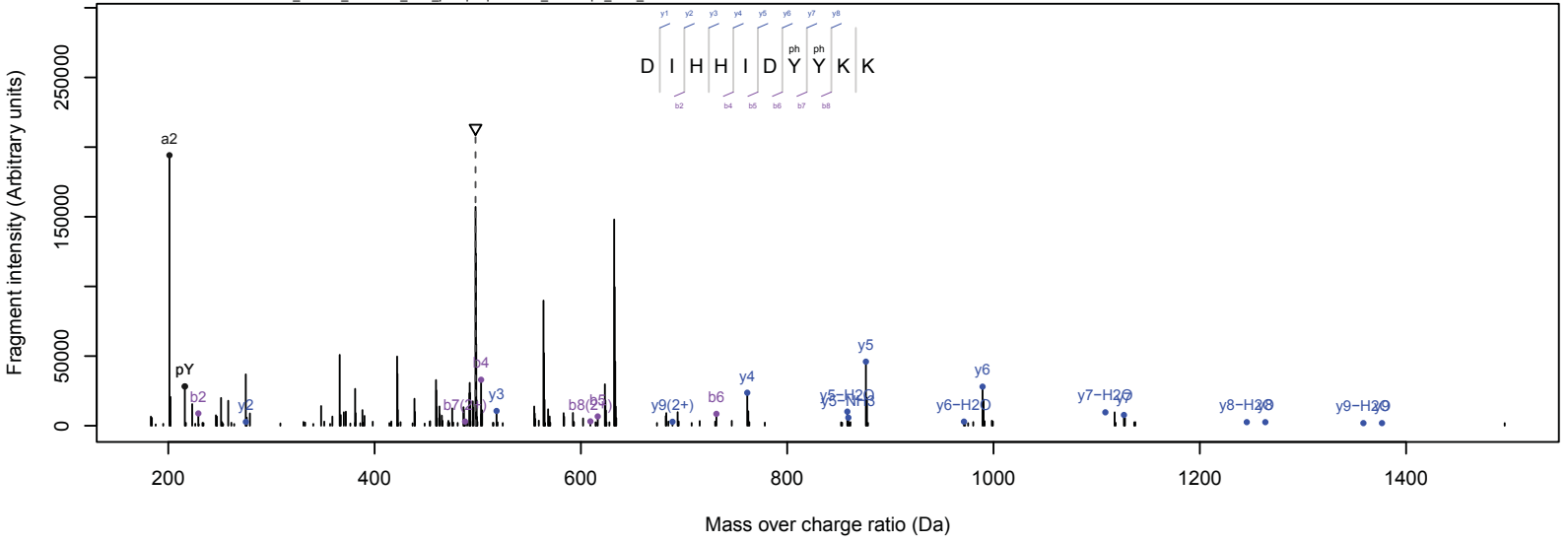
Andromeda score : 146.13



**DIHHIDY(ph)Y(ph)KK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 4906

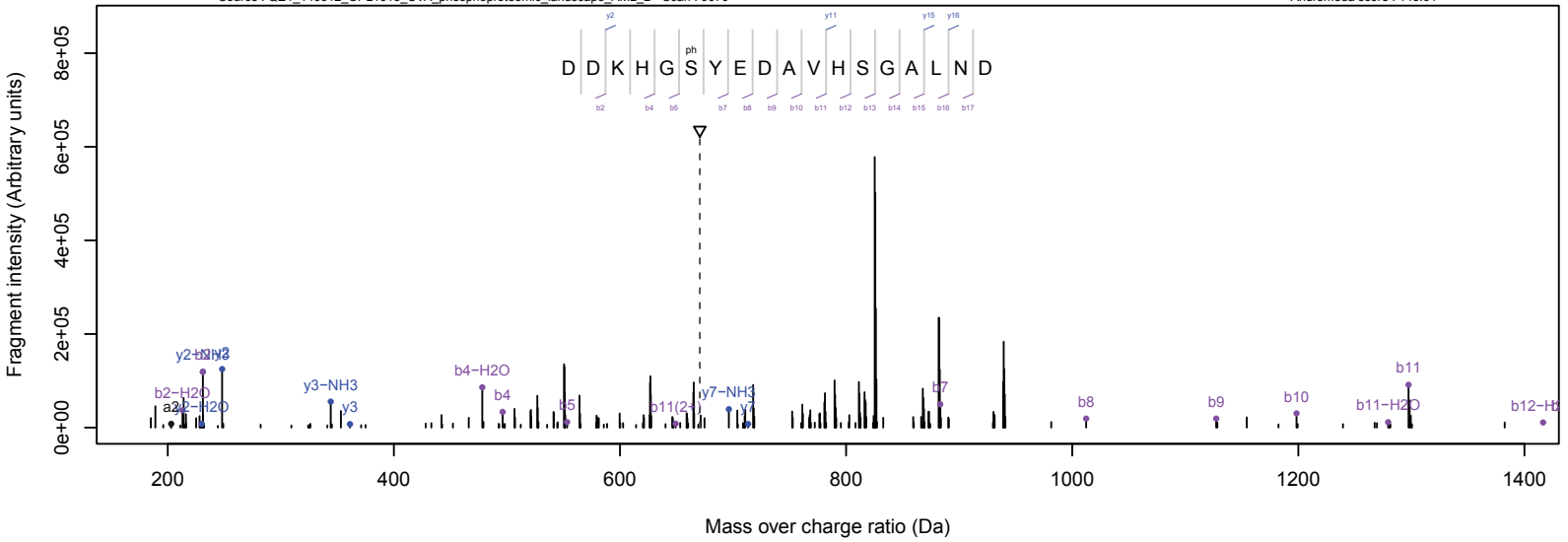
Andromeda score : 146.11



**DDKHGS(ph)YEDAVHSGALND-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9670

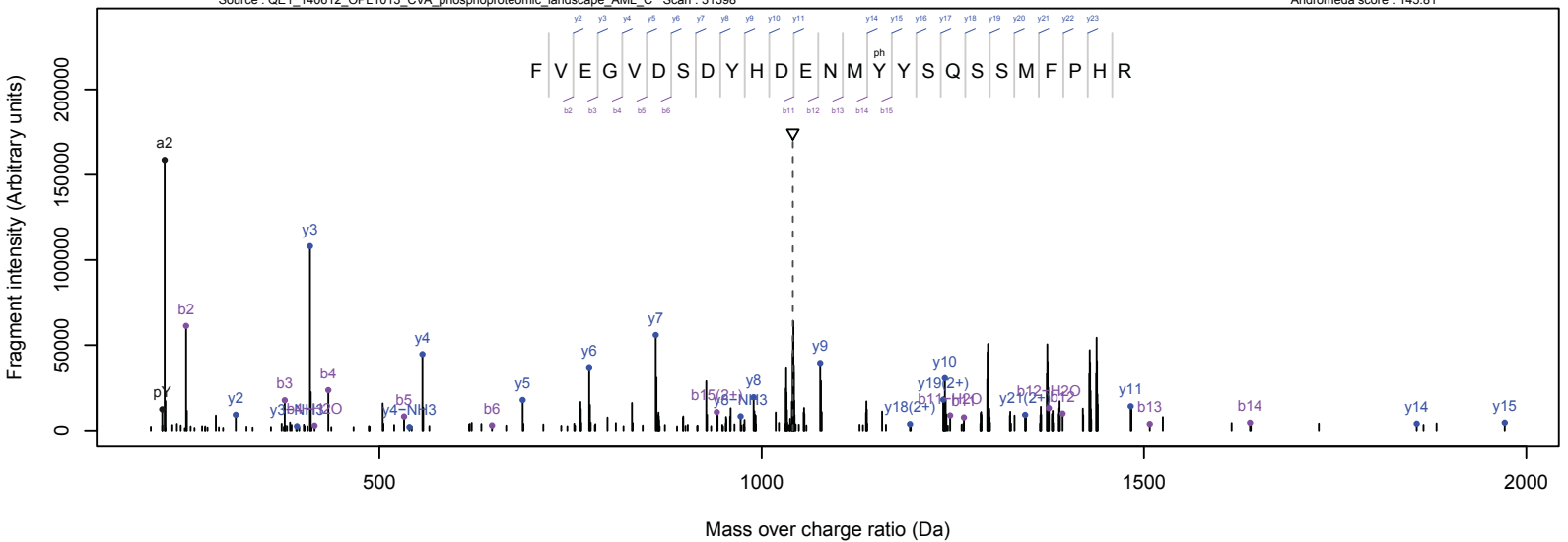
Andromeda score : 145.91



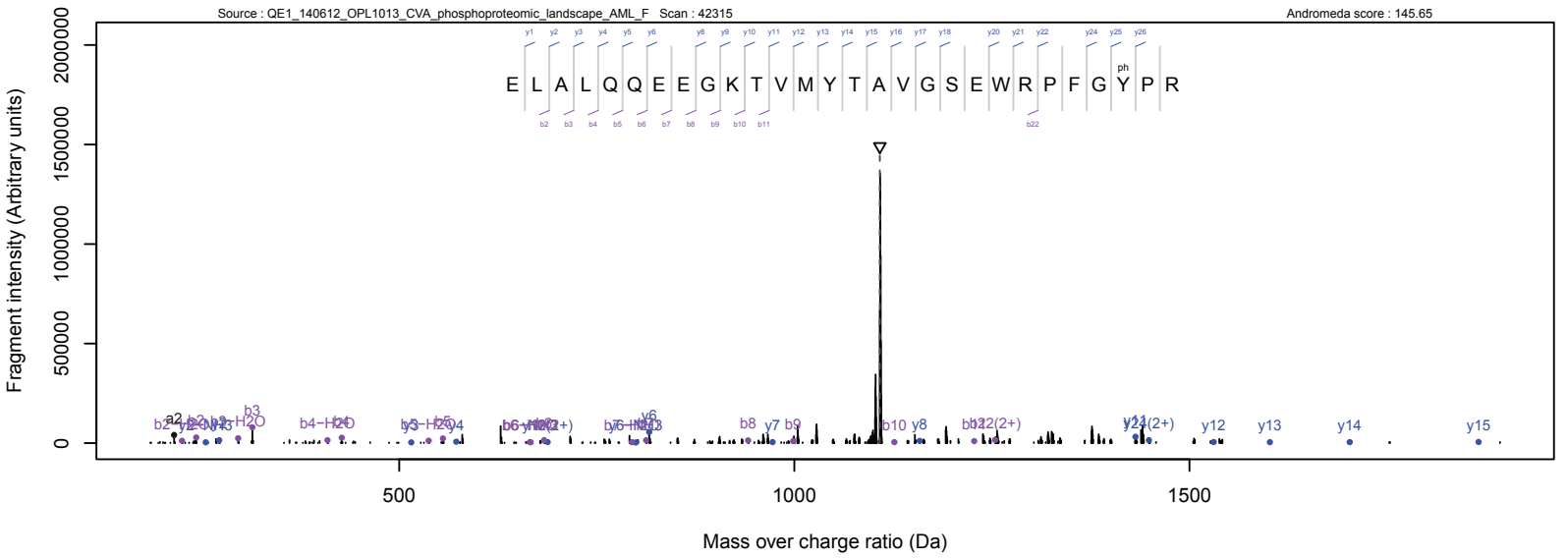
**FVEGVDSYHDENMY(ph)YSQSSMFPHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31398

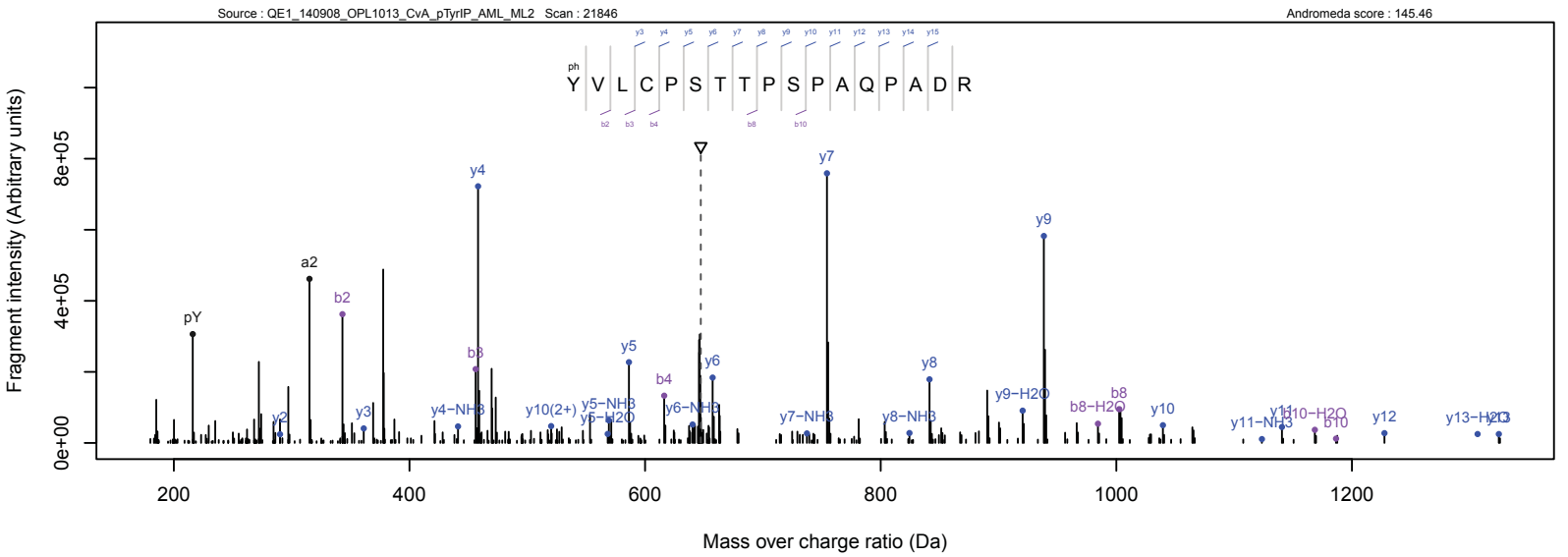
Andromeda score : 145.81



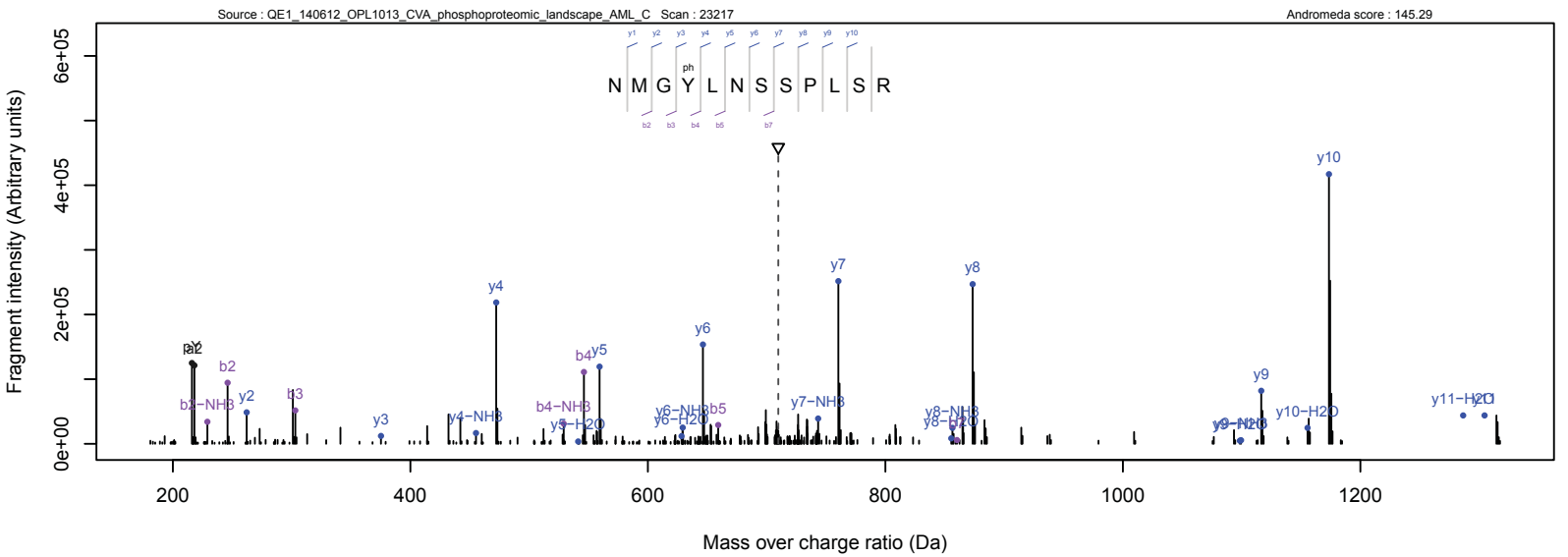
ELALQQEEGKTVMYTAVGSEWRPFGYPR-3+



Y(ph)VLCPSTTSPAQPADR-3+



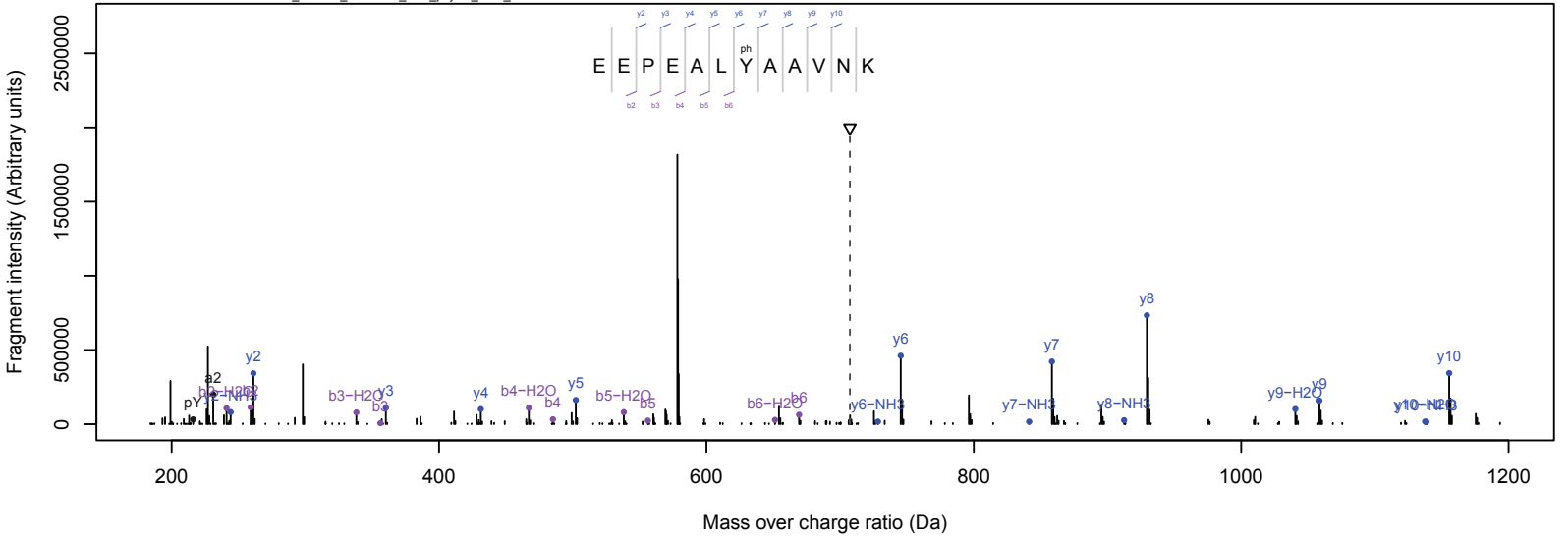
NMGY(ph)LNSSPLSR-2+



EEPEALY(ph)AAVNK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23232

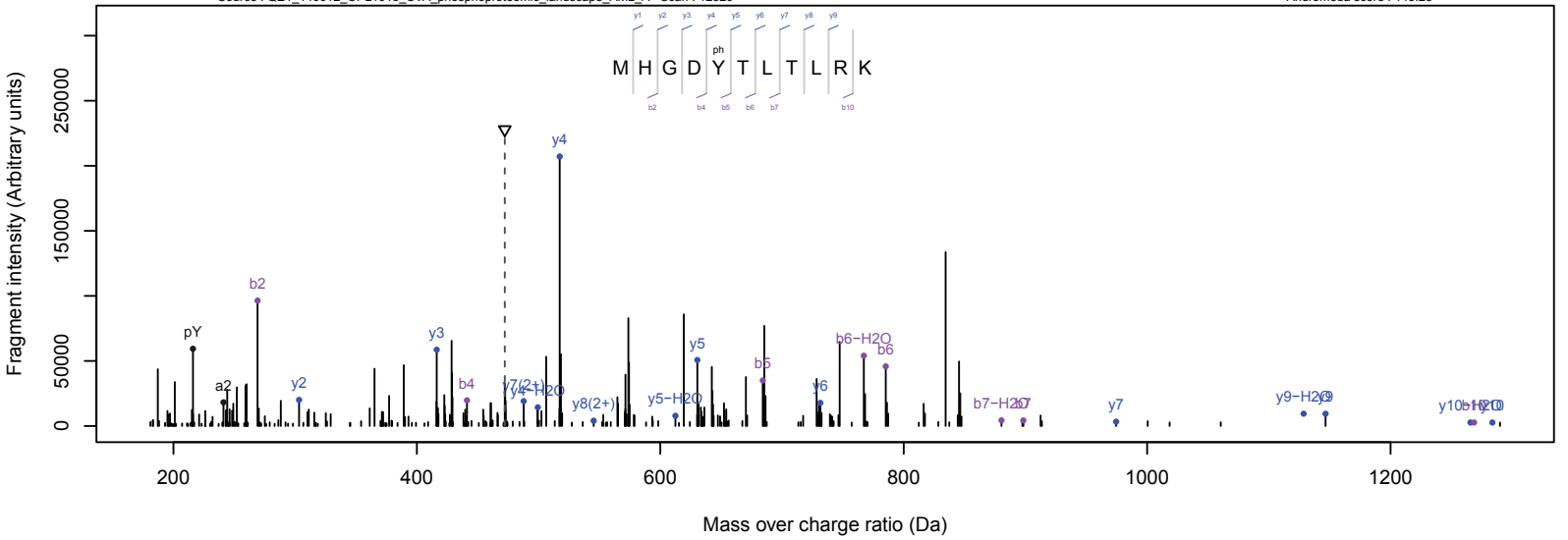
Andromeda score : 145.28



MHGDY(ph)TLTLRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12529

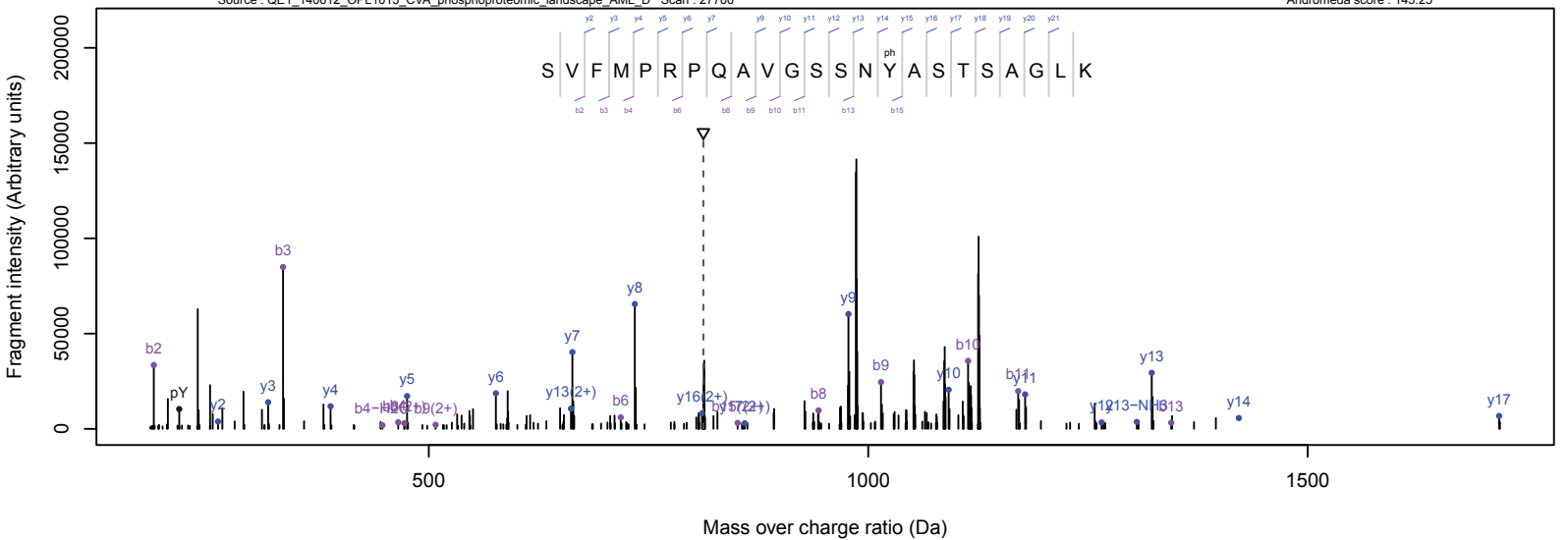
Andromeda score : 145.28



SVFMRPQAVGSSNY(ph)ASTSAGLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27700

Andromeda score : 145.25

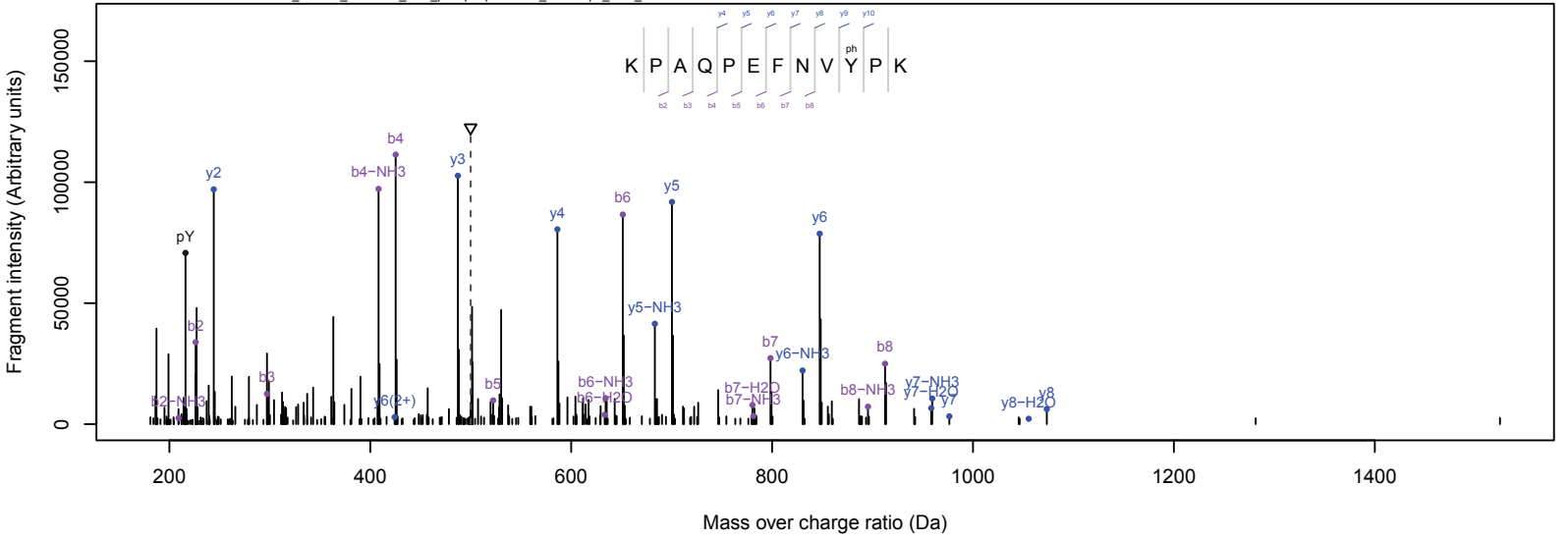




**KPAQPEFNVYPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16276

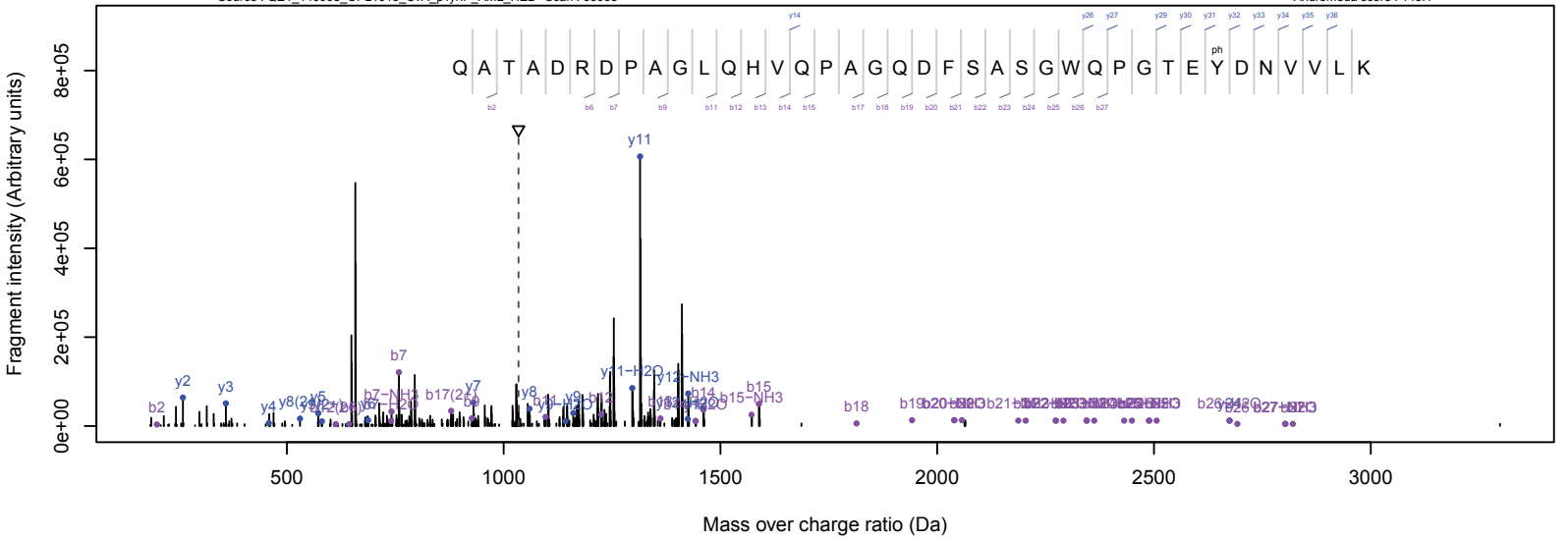
Andromeda score : 145.23



**QATADRDPAQLQHVQPAGQDFASASGWQPGTEY(ph)DNVVLK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35938

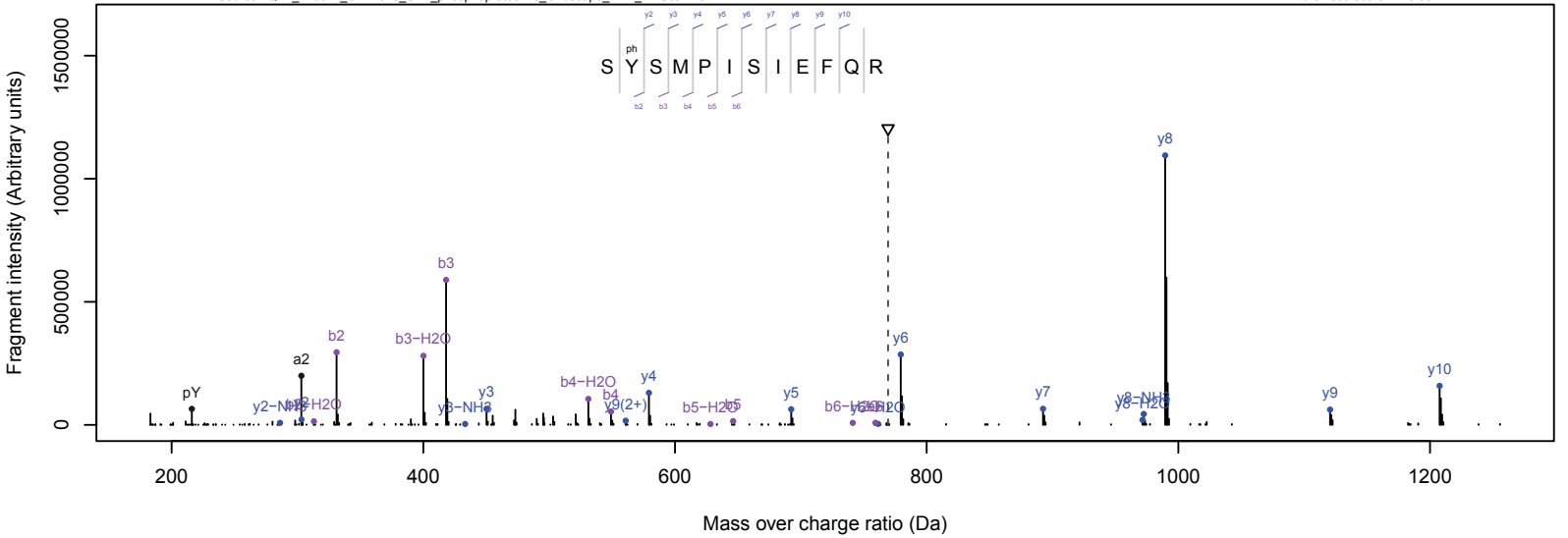
Andromeda score : 145.1



**SY(ph)SMPISIEFQR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37414

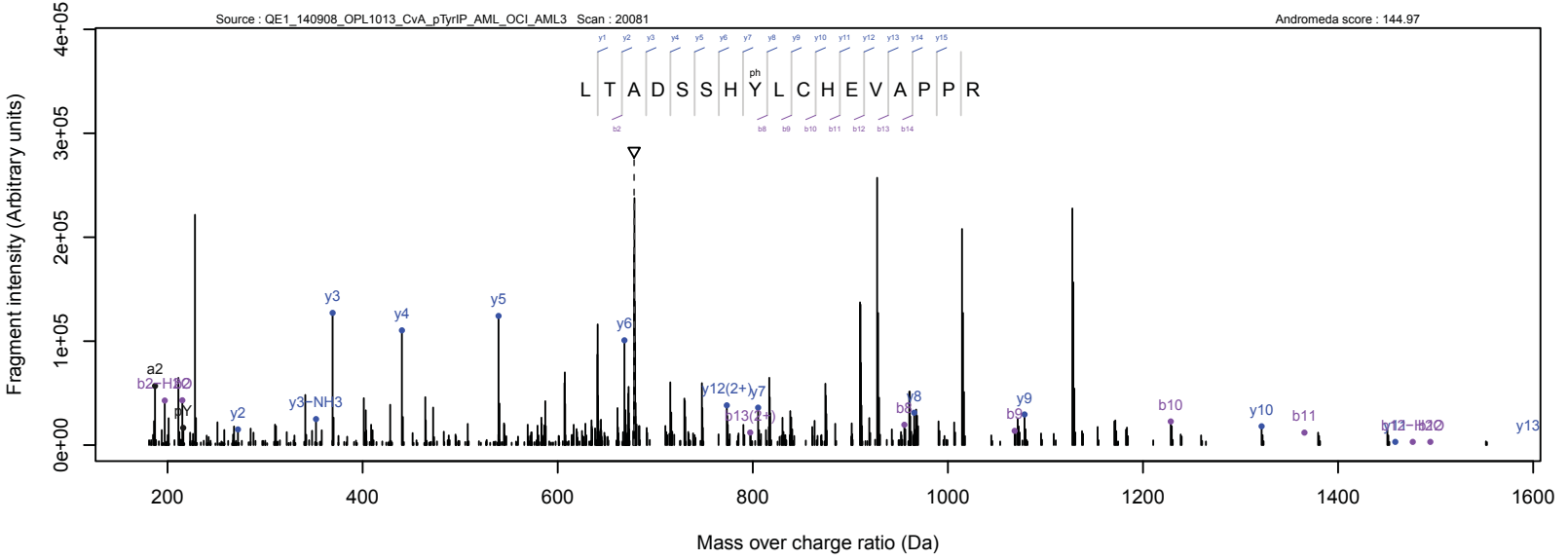
Andromeda score : 145.09



LTADSSHY(ph)LCHEVAPPR-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_OCI\_AML3 Scan: 20081

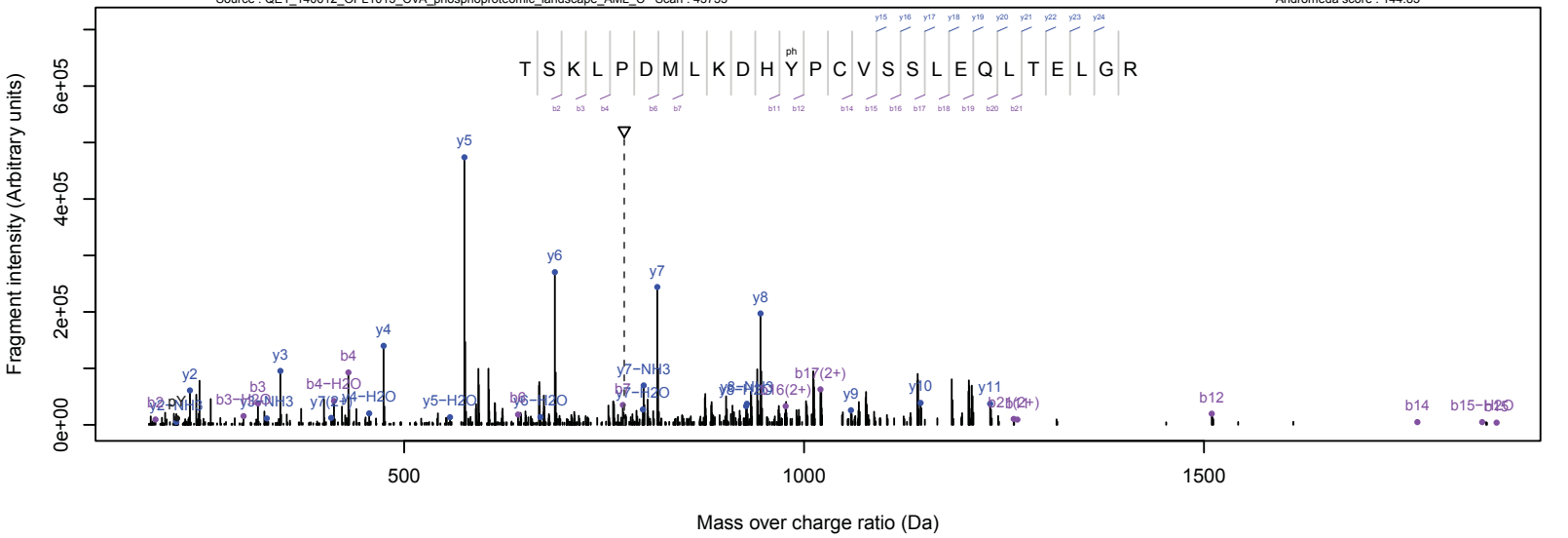
Andromeda score: 144.97



TSKLPDMLKDHY(ph)PCVSSLEQLTELGR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 43755

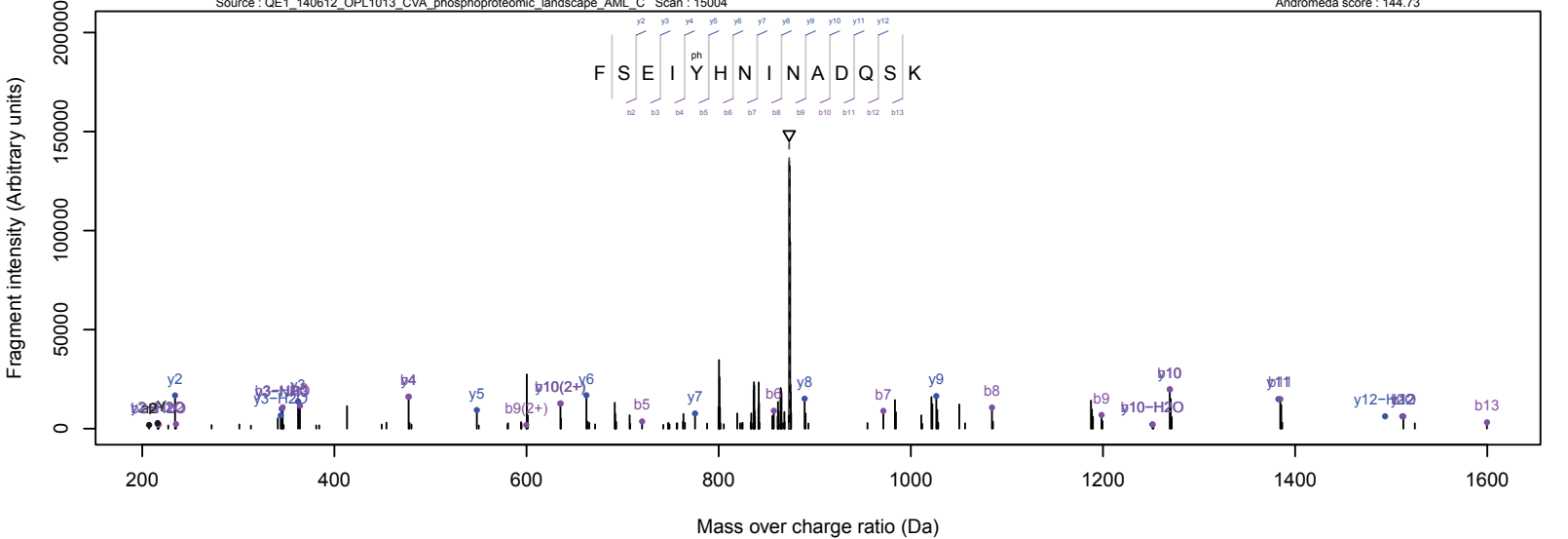
Andromeda score: 144.83



FSEIY(ph)HNINADQSK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15004

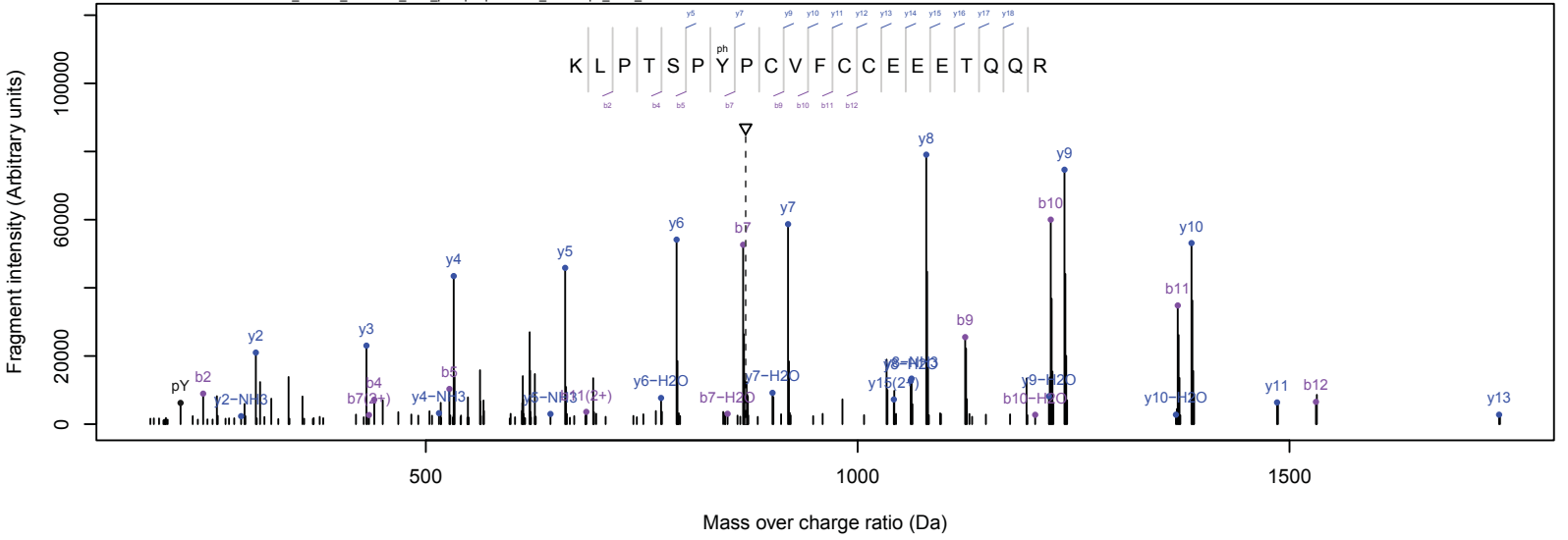
Andromeda score: 144.73



**KLPTSPY(ph)PCVFCCEETQQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27908

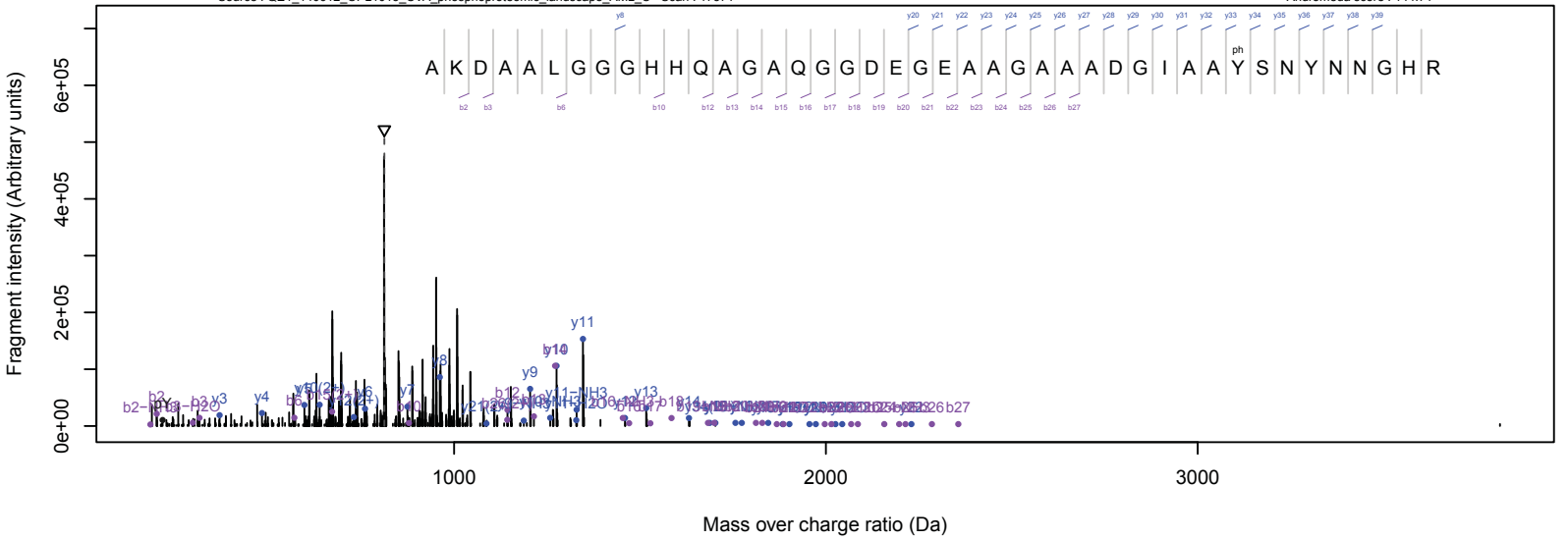
Andromeda score : 144.72



**AKDAALGGHHQAGQGDEGEAAGAADGIAAY(ph)SNYNNNGHR-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17571

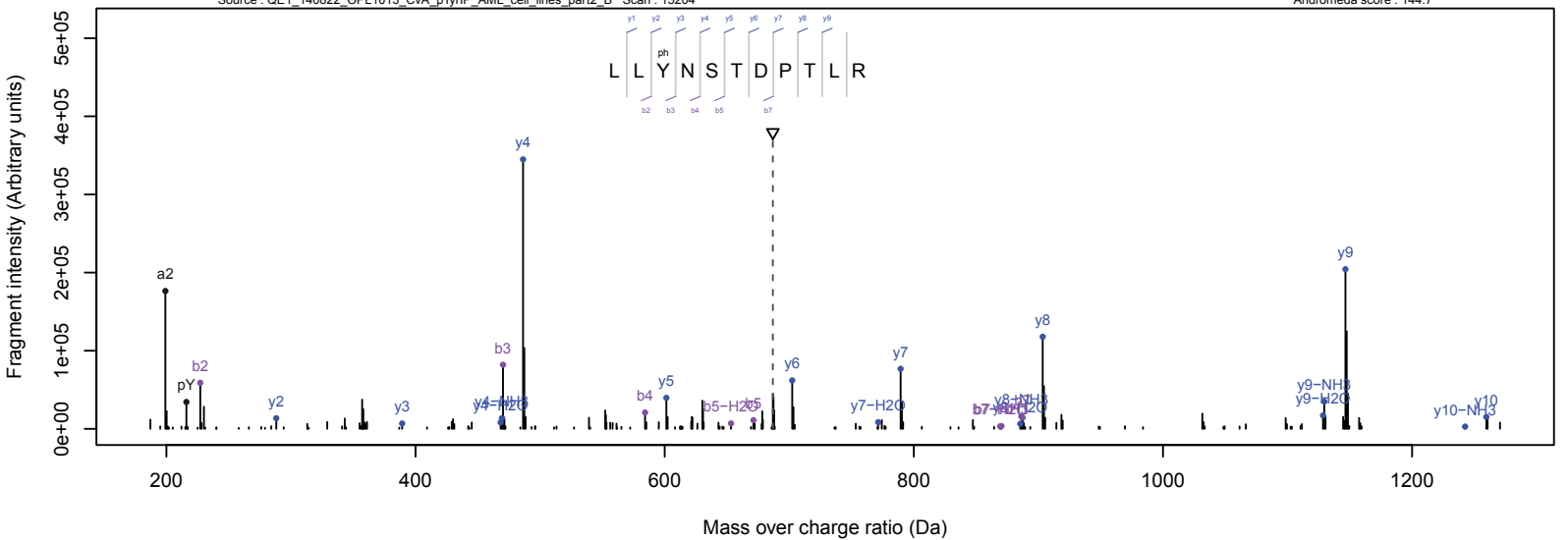
Andromeda score : 144.71



**LLY(ph)NSTDPTLR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 15204

Andromeda score : 144.7

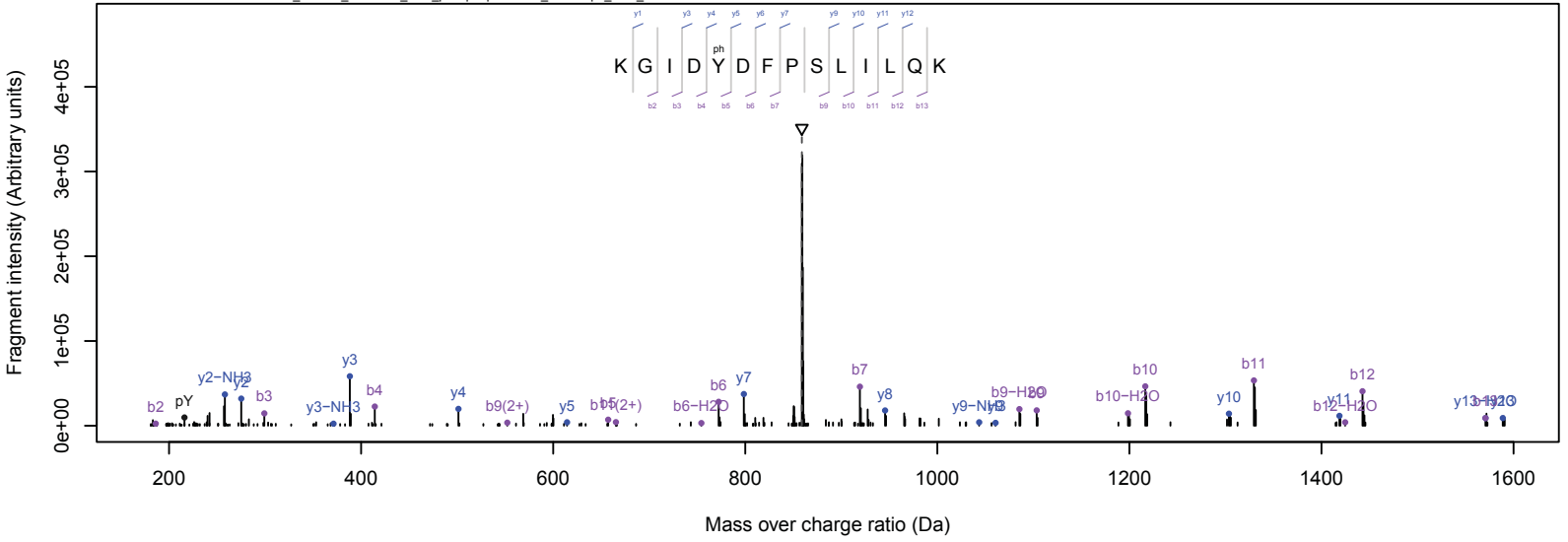




**KGIDY(ph)DFPSLILQK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42048

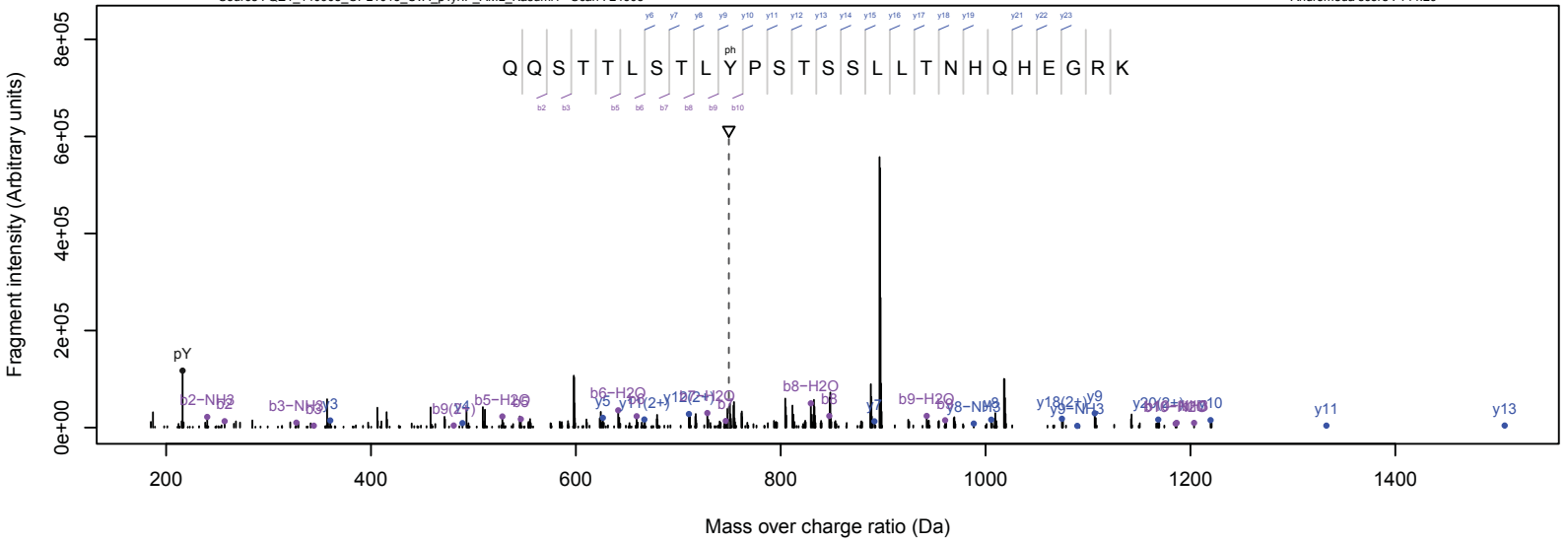
Andromeda score : 144.29



**QQSTTLSTLY(ph)PSTSSLLTNHQHEGRK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 24603

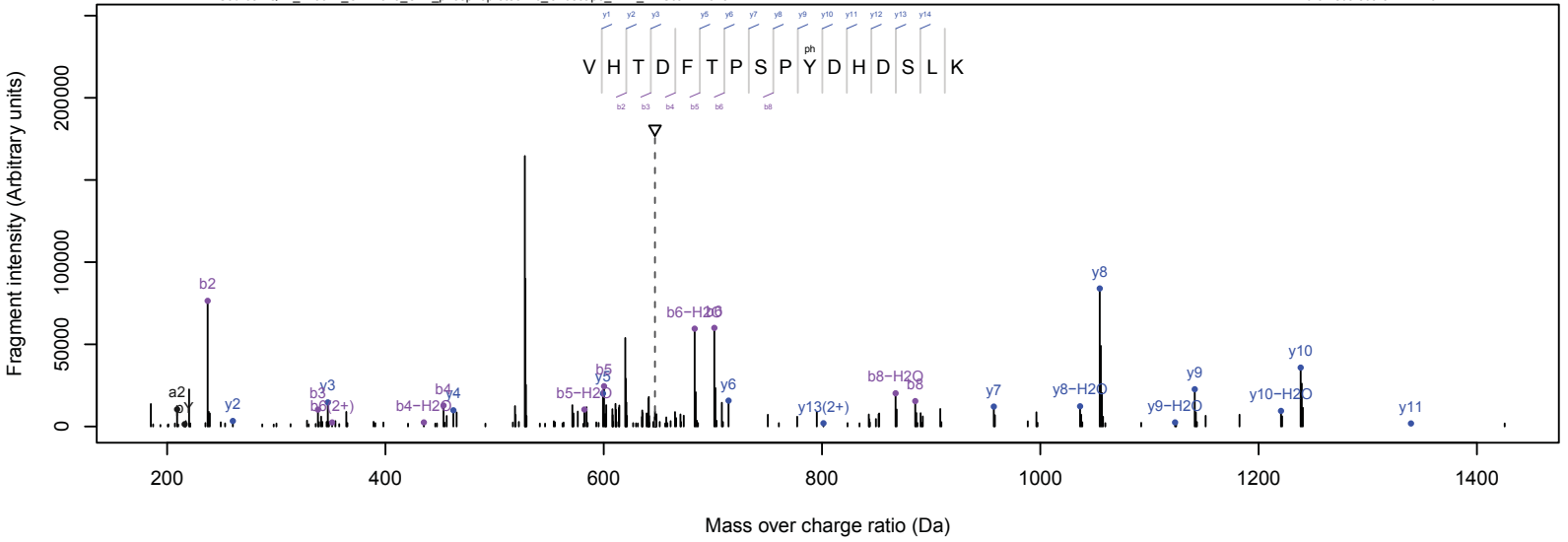
Andromeda score : 144.29

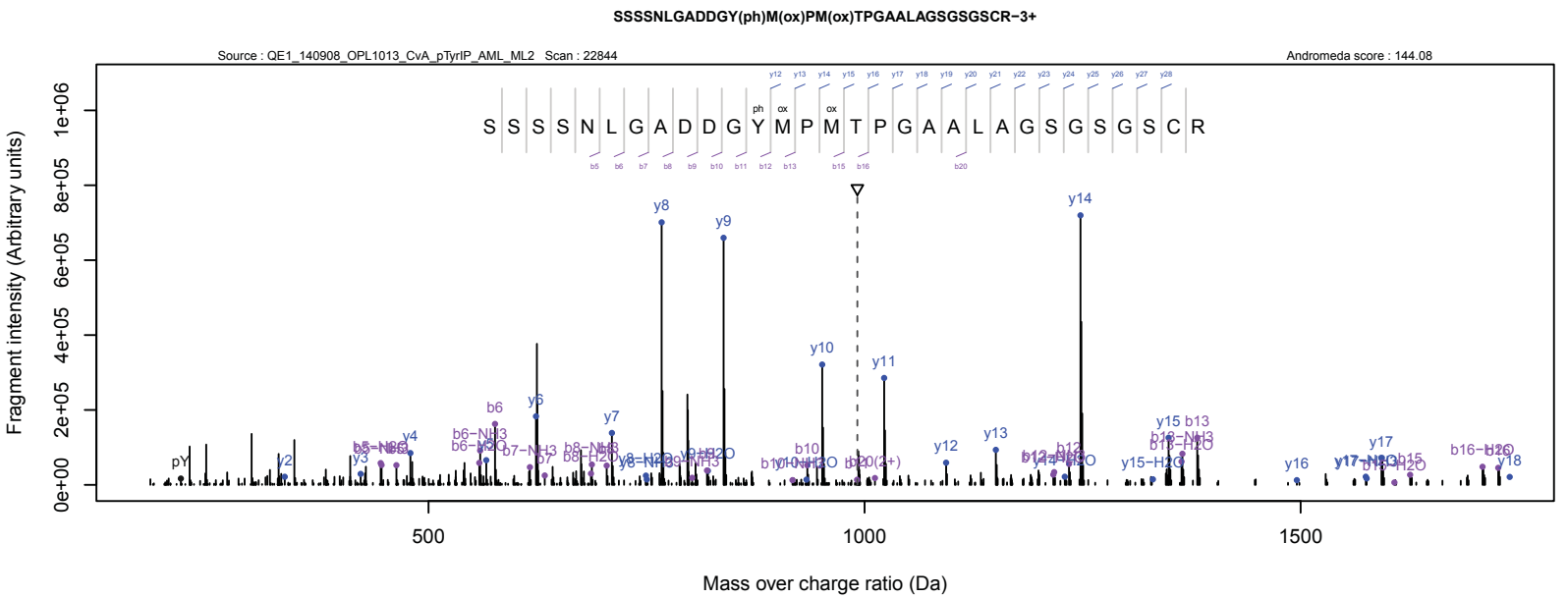
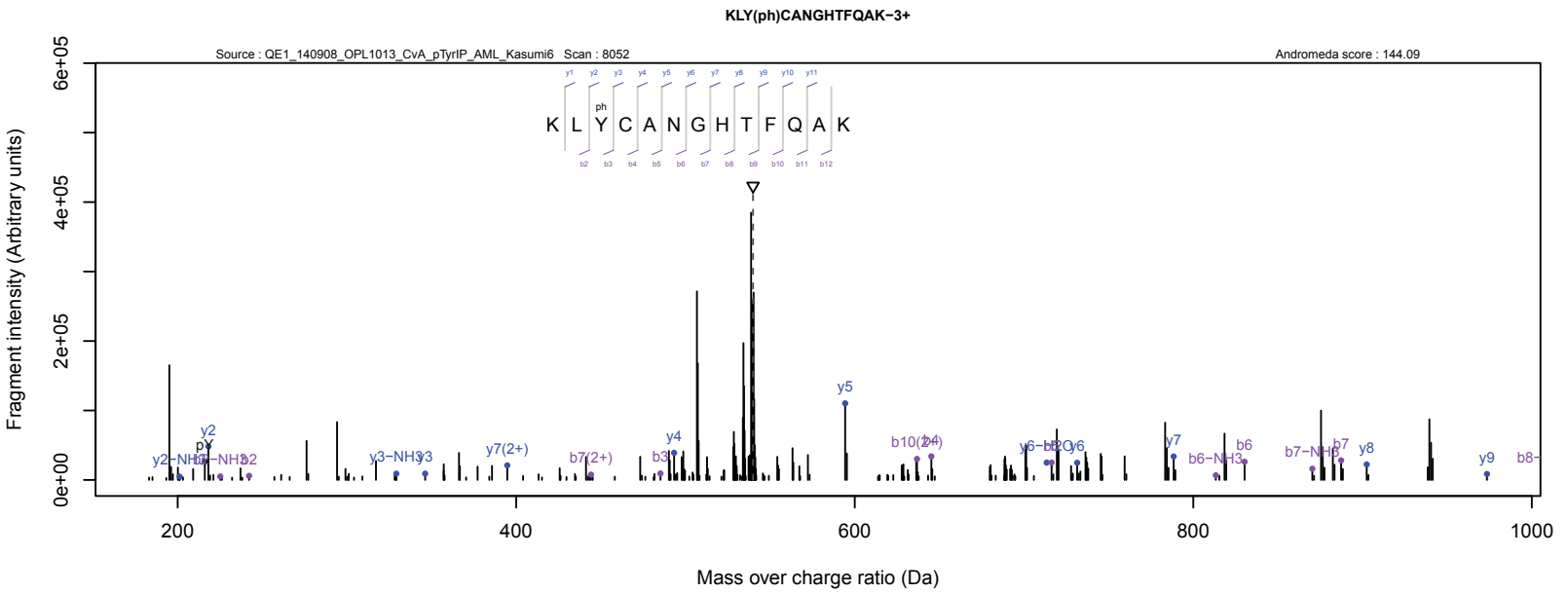
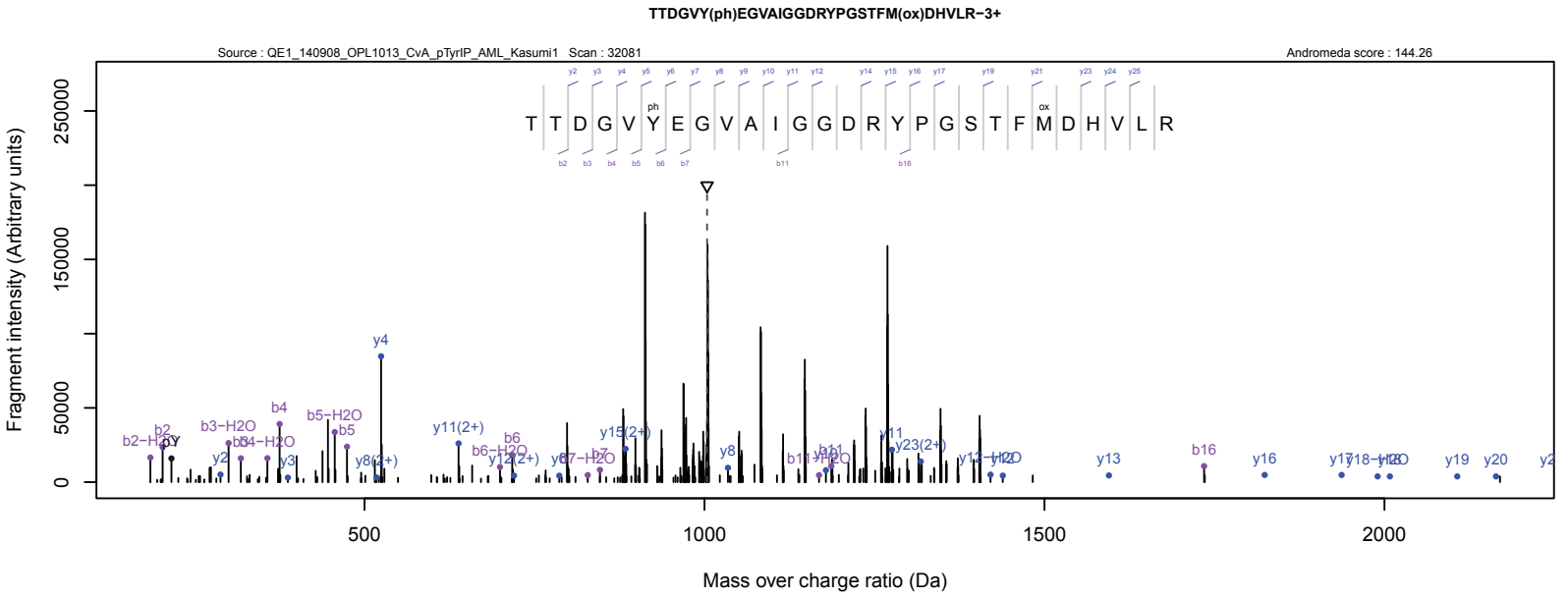


**VHTDFTSPY(ph)DHDSLK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 16781

Andromeda score : 144.29

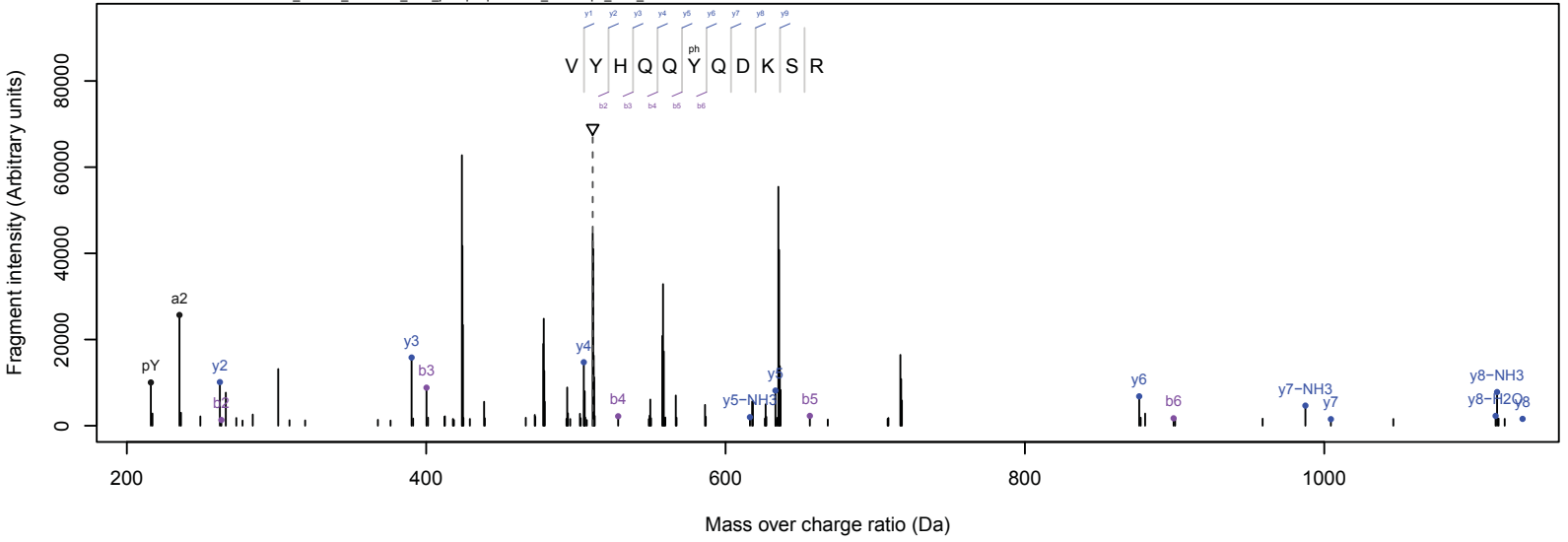




VYHQY(ph)QDKSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 3161

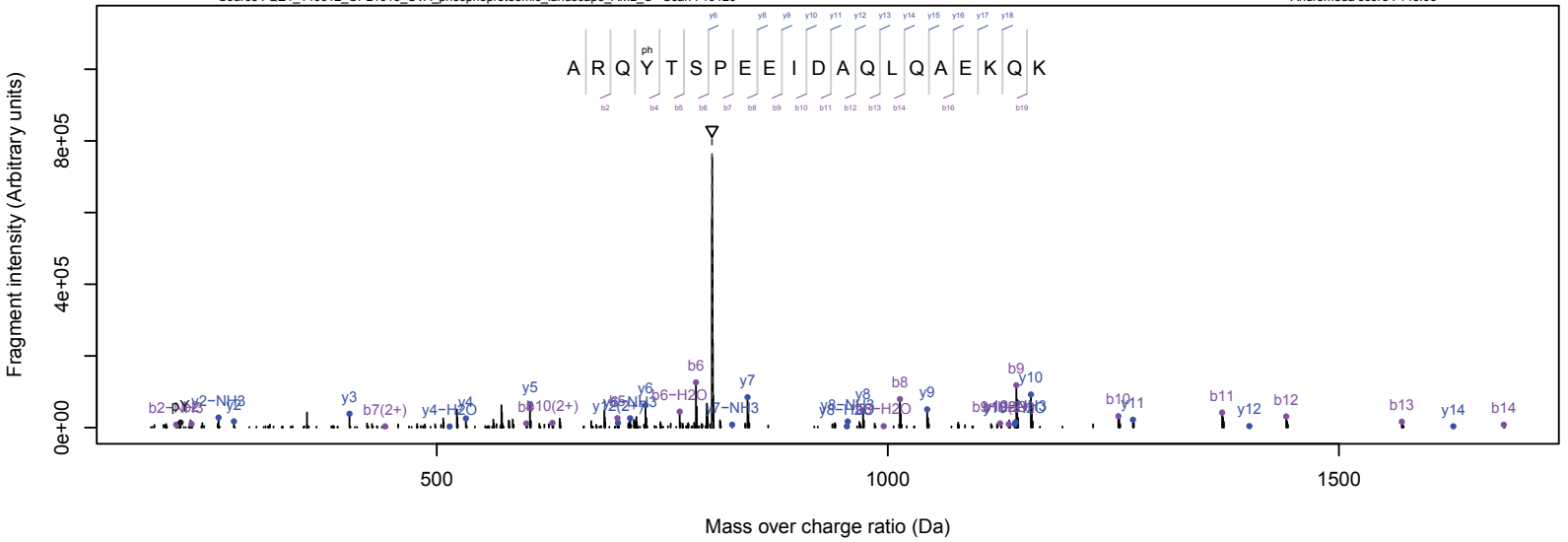
Andromeda score : 143.97



ARQY(ph)TSPEEIDAQLQAEKQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19120

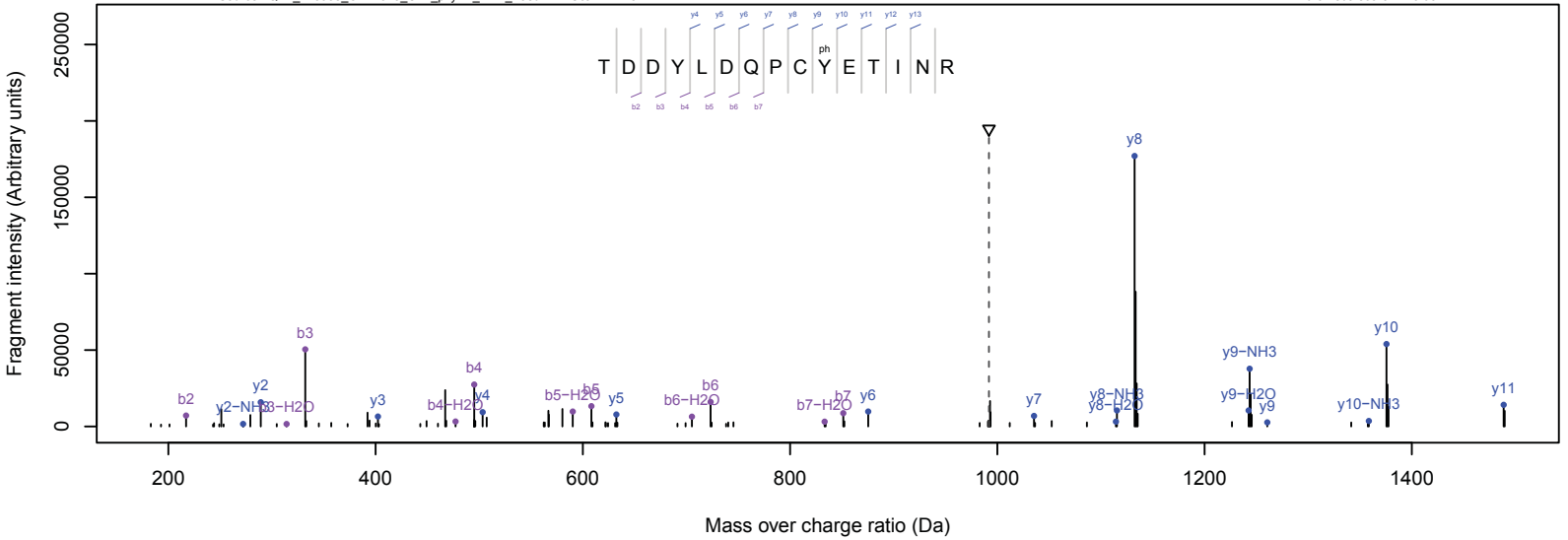
Andromeda score : 143.93



TDDYLDQPCY(ph)ETINR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 24281

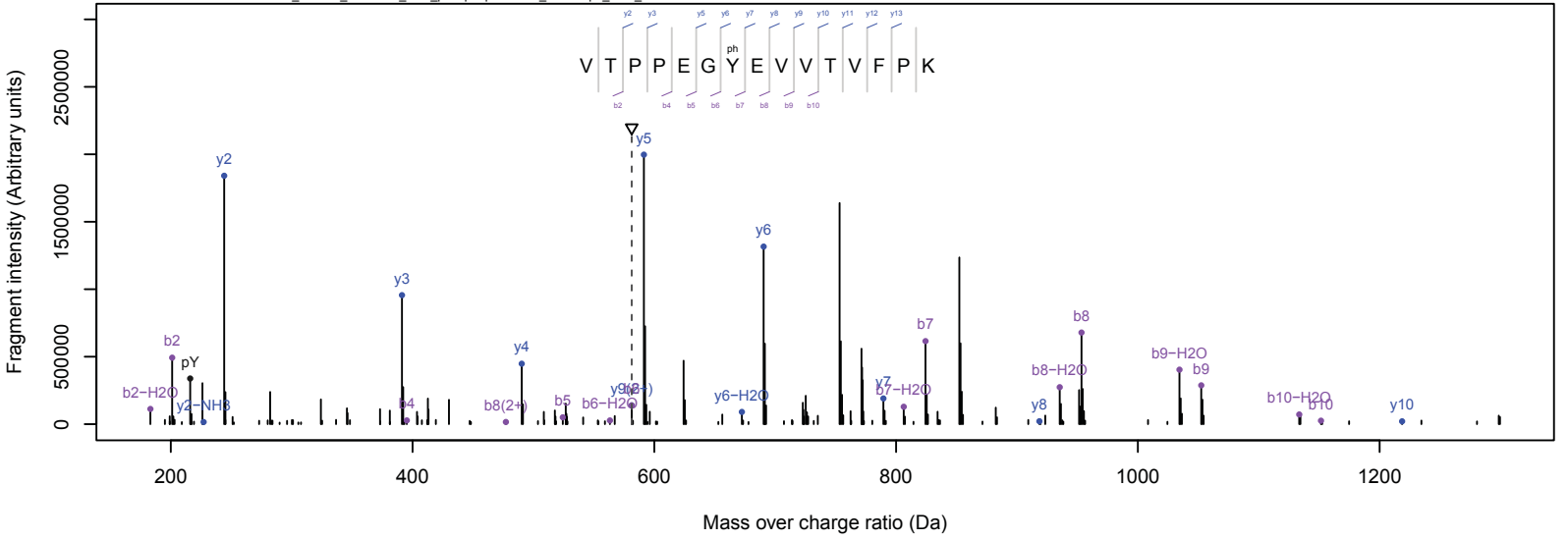
Andromeda score : 143.93



VTPPEGY(ph)EVVTVFPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36461

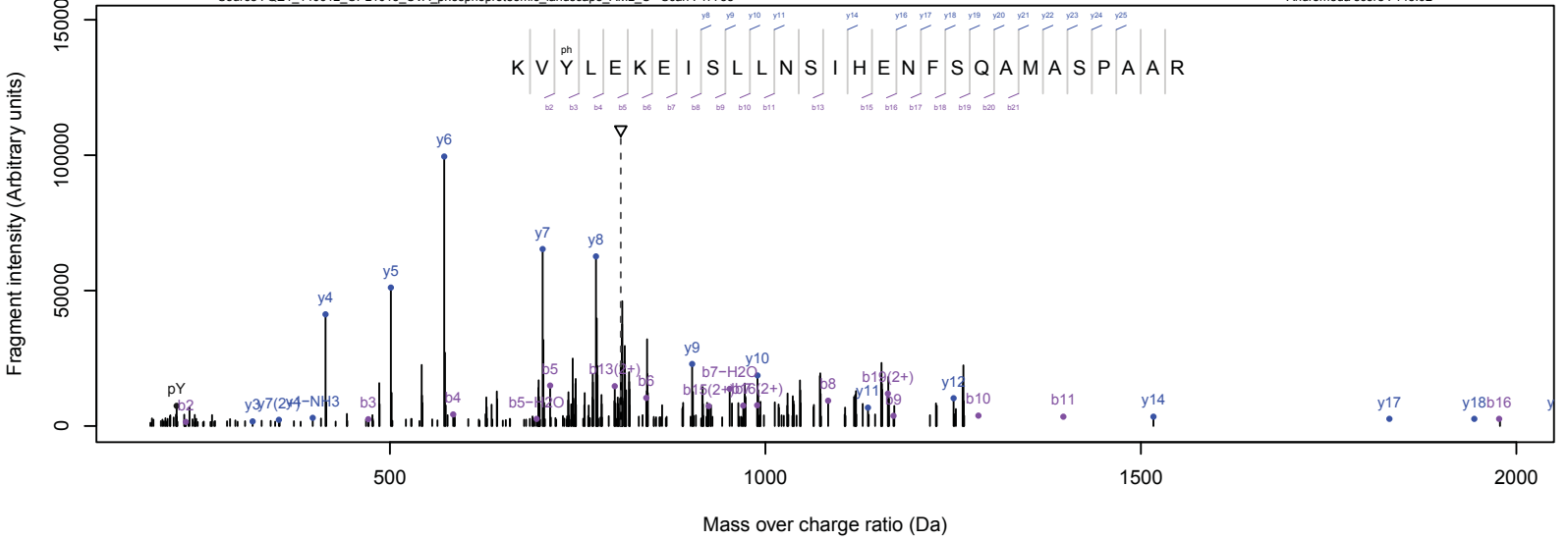
Andromeda score : 143.93



KVY(ph)LEKEISLLNSIHENFSQAMASPAAR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47788

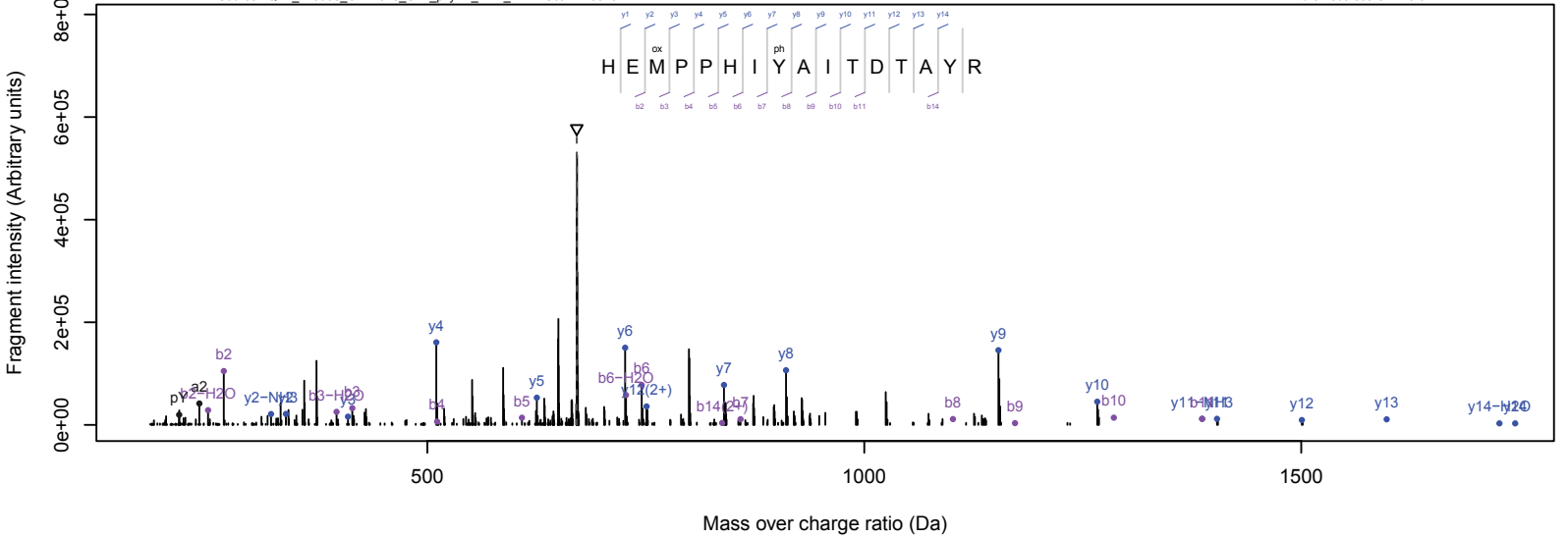
Andromeda score : 143.92



HEM(ox)PPHIY(ph)AITDTAYR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23515

Andromeda score : 143.67

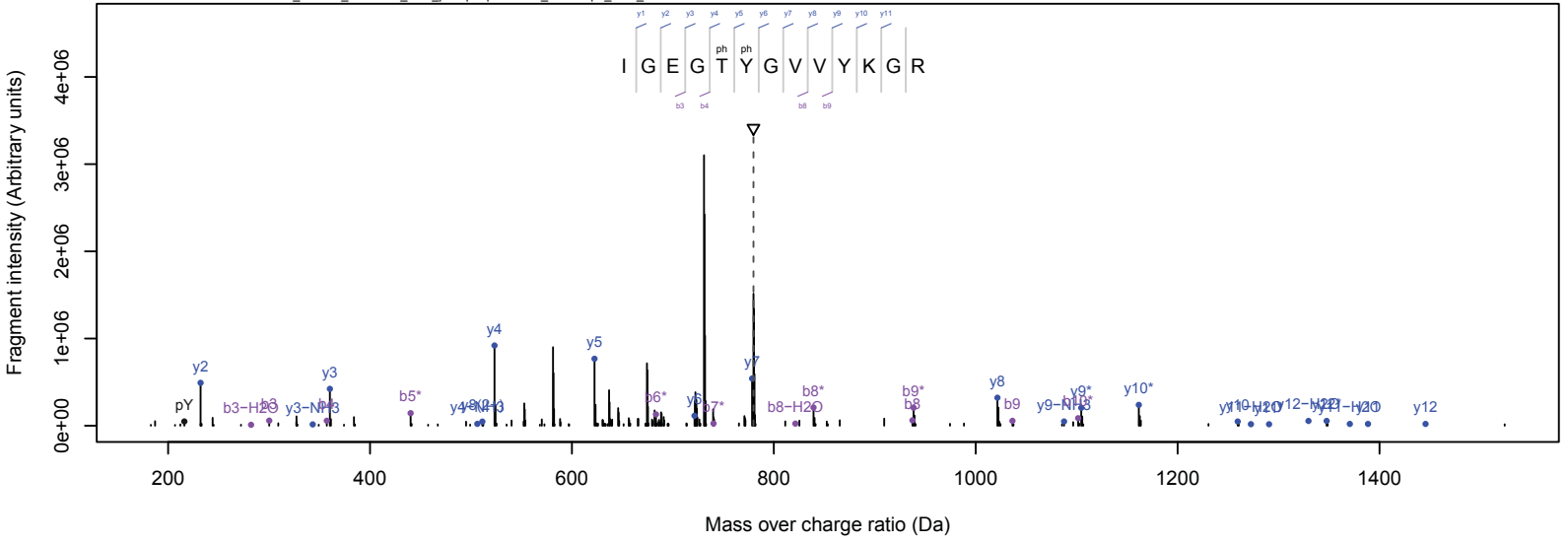




**IGEGT(ph)Y(ph)GVVYKGR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan: 16025

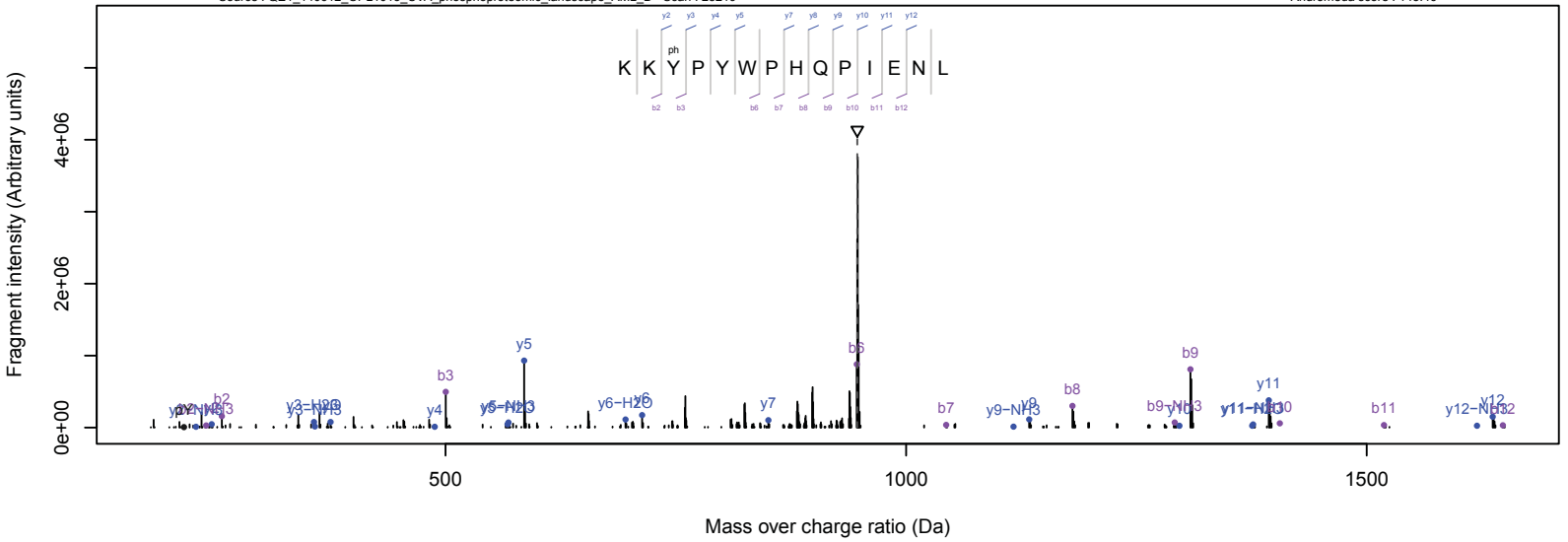
Andromeda score: 143.46



**KKY(ph)PYWPHQPIENL-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 28219

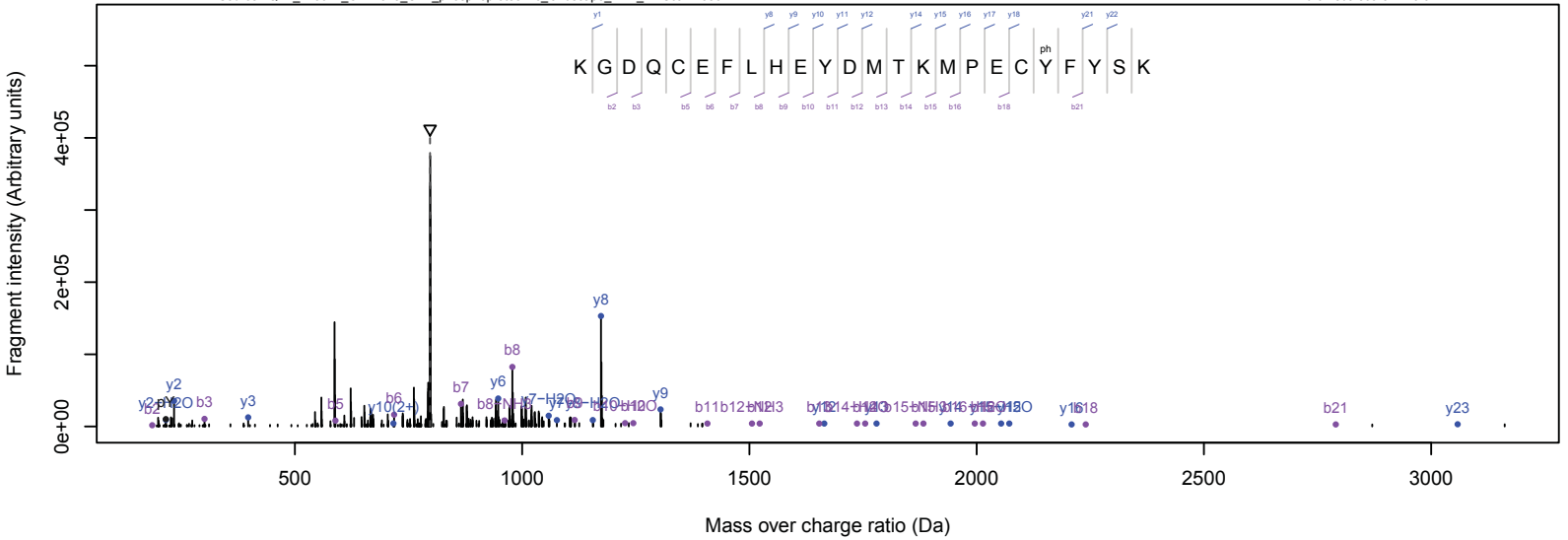
Andromeda score: 143.46



**KGDQCEFLHEYDMTKMPECY(ph)FYSK-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 35612

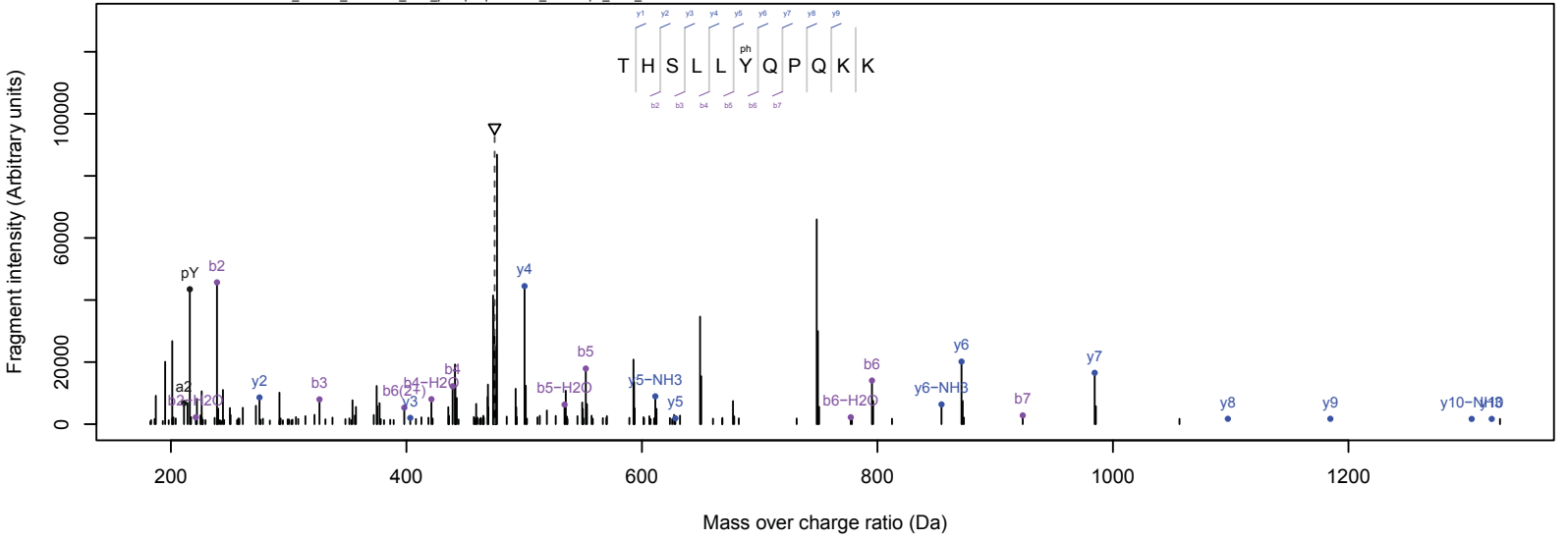
Andromeda score: 143.32



THSLLY(ph)QPQKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8461

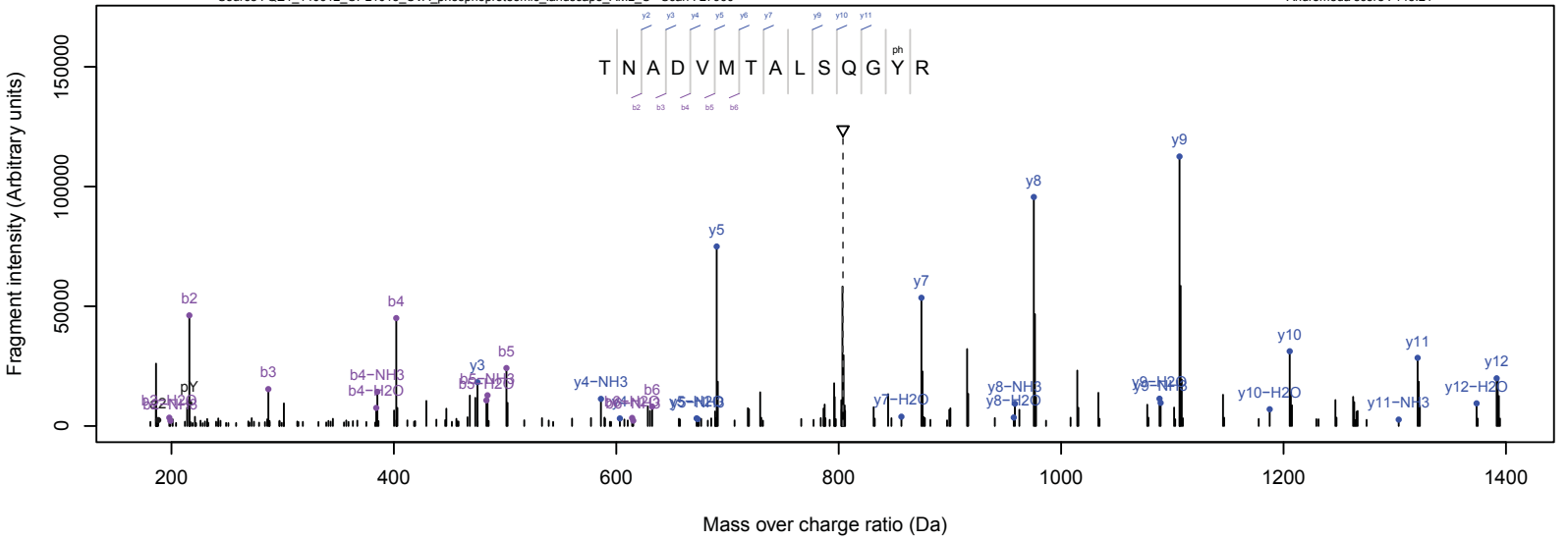
Andromeda score : 143.24



TNADVMTALSQGY(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27960

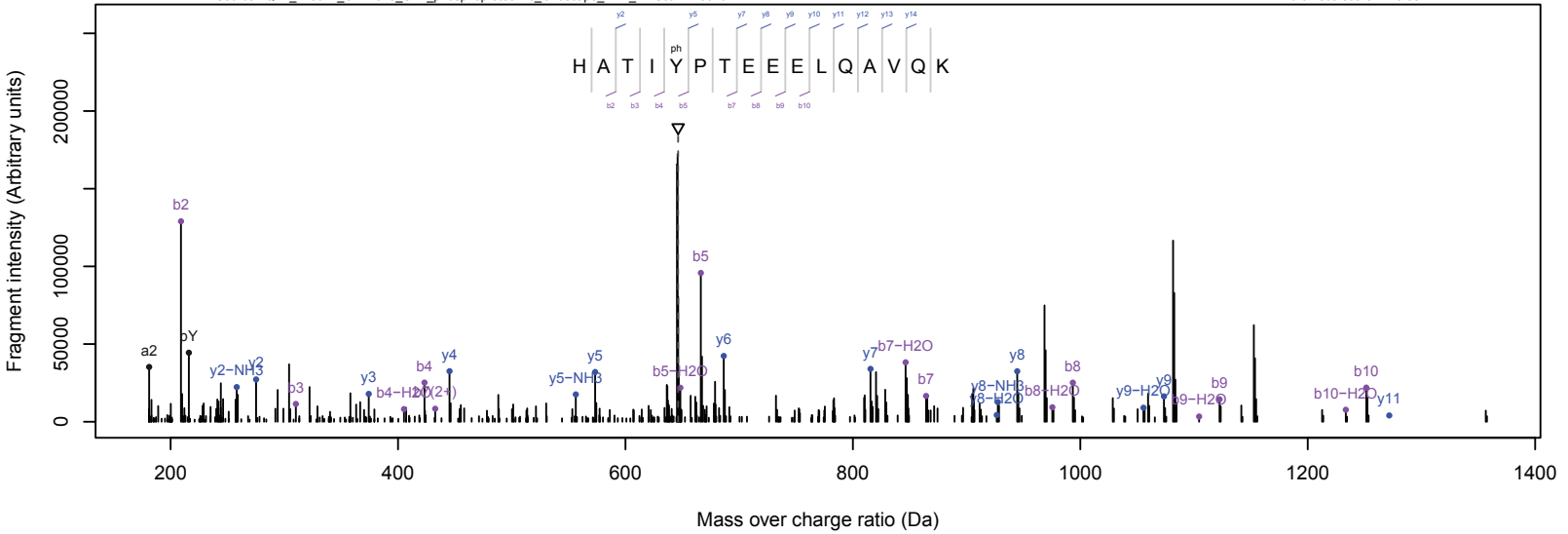
Andromeda score : 143.21



HATIY(ph)PTEELQAVQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23326

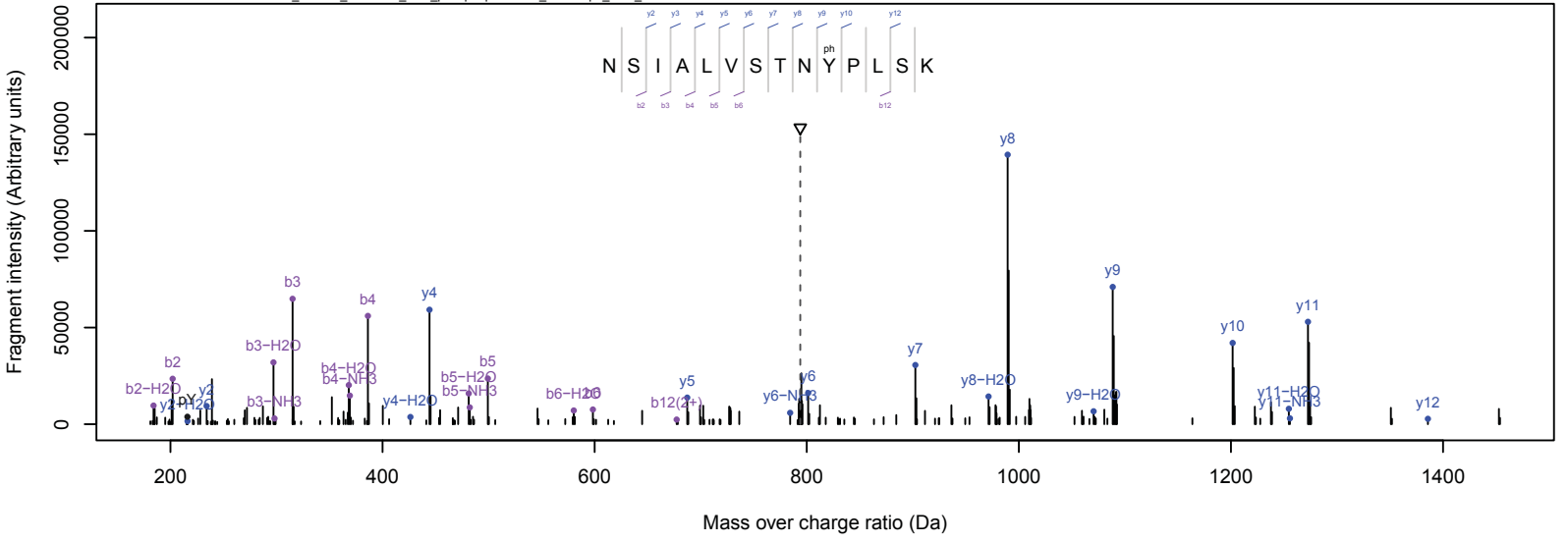
Andromeda score : 143.05



NSIALVSTNY(ph)PLSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28729

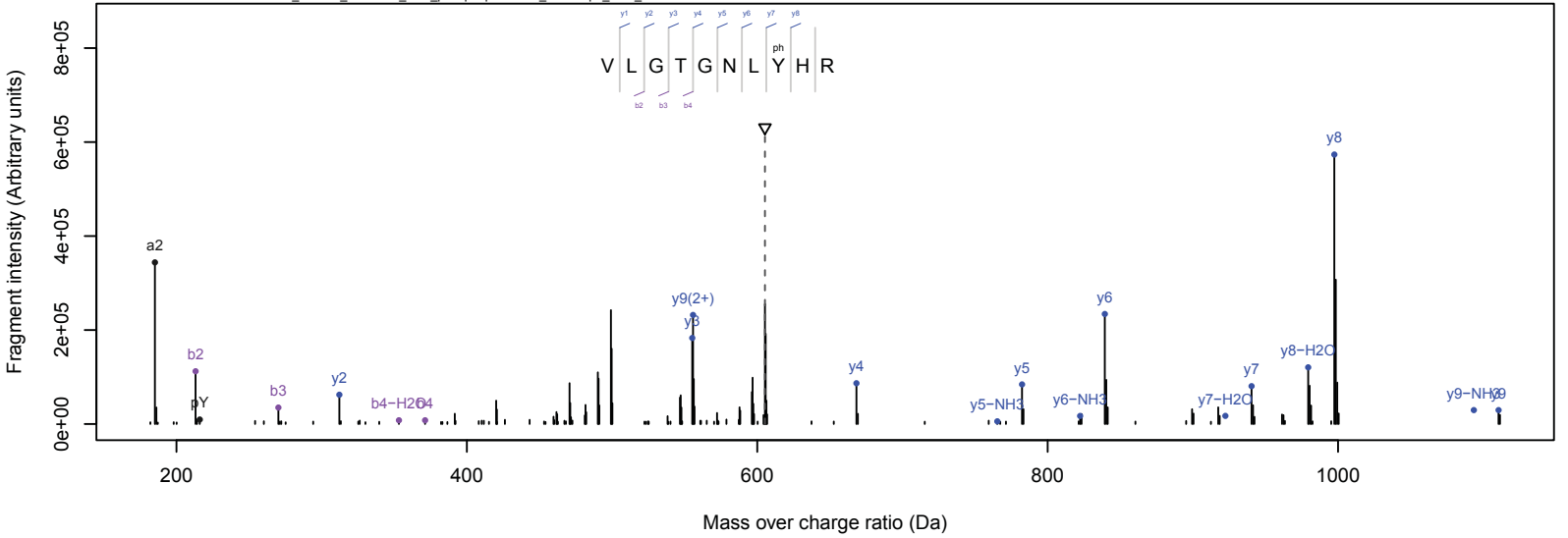
Andromeda score : 143.05



VLGTGNLY(ph)HR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11401

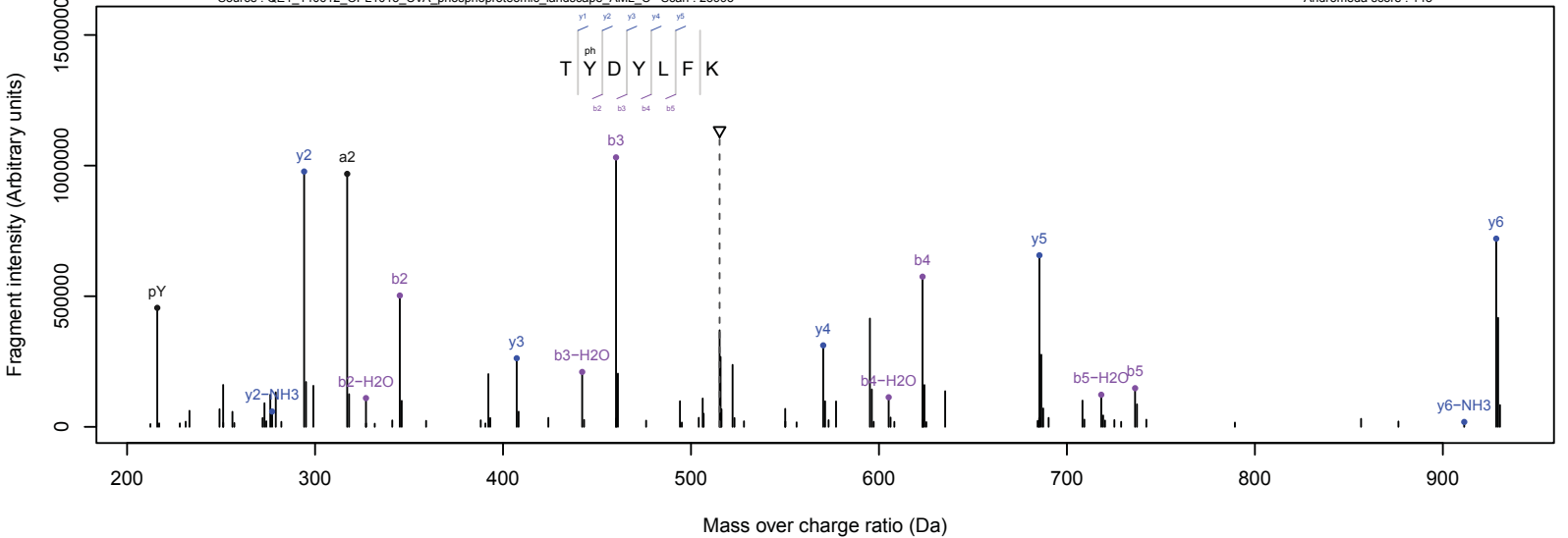
Andromeda score : 143.03



TY(ph)DYLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25993

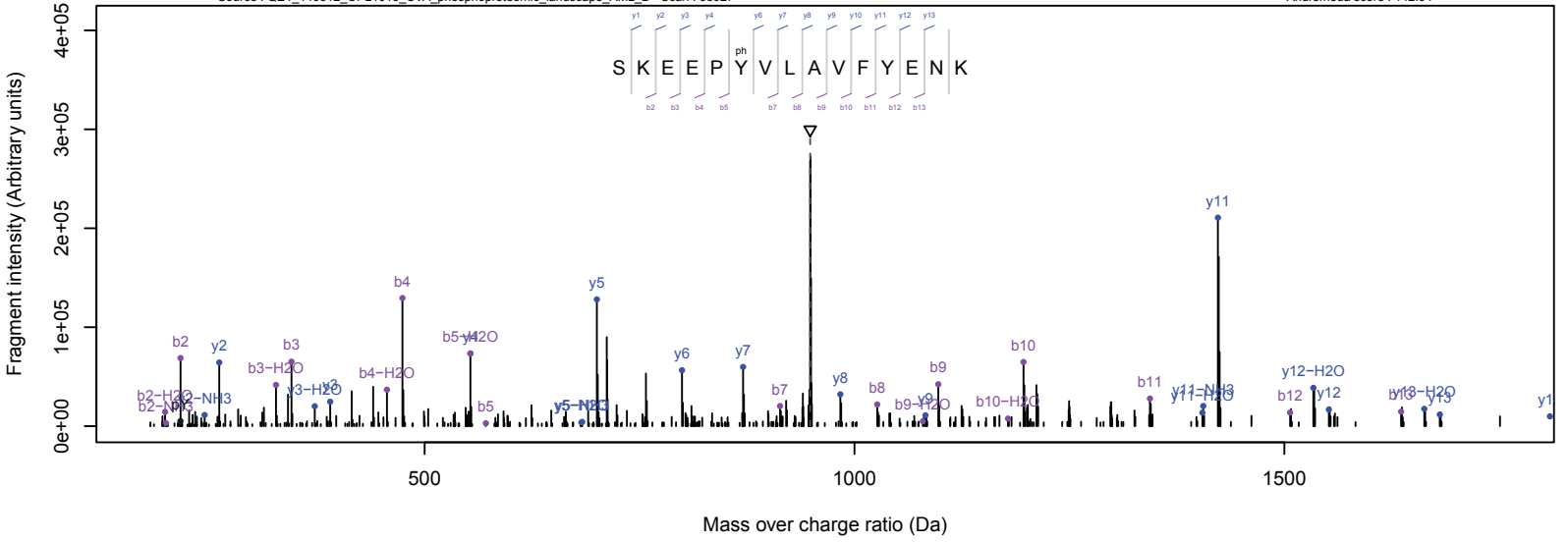
Andromeda score : 143



**SKEEPY(ph)VLAVFYENK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 38927

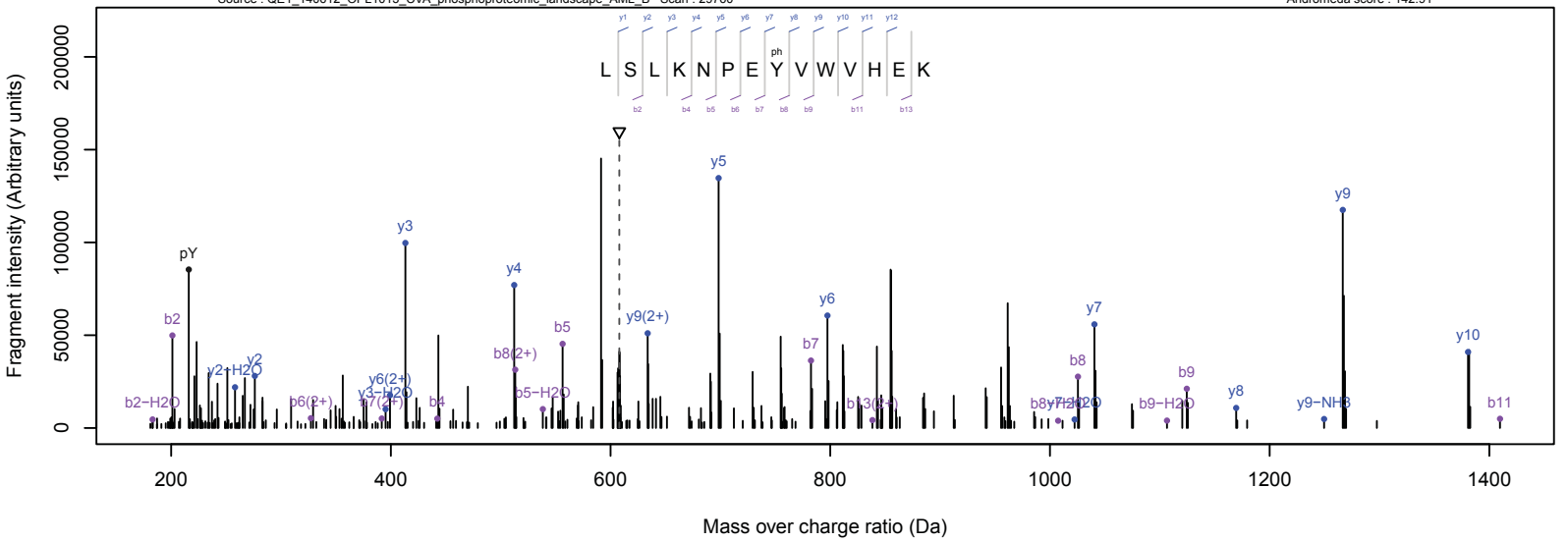
Andromeda score: 142.94



**LSLKNPEY(ph)VWVHEK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 29760

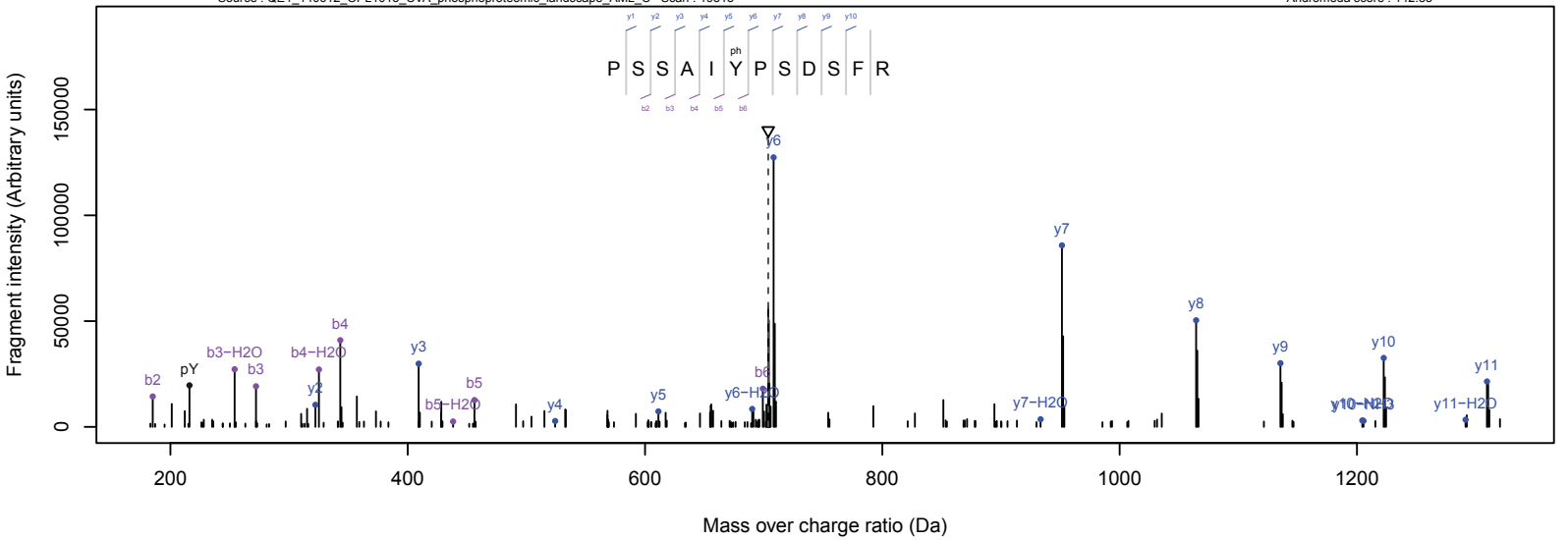
Andromeda score: 142.91



**PSSAIY(ph)PSDSFR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19313

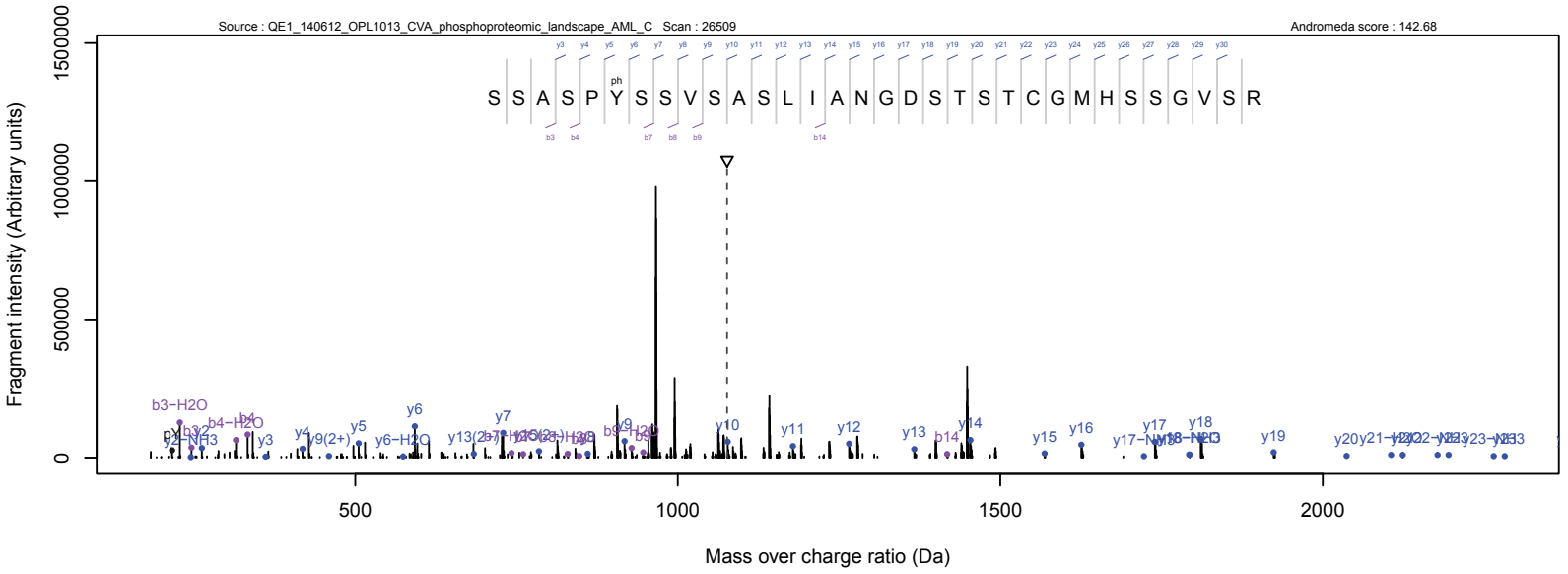
Andromeda score: 142.83



SSASPY(ph)SSVSASLIANGDSTSTCGMHSSGVS R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26509

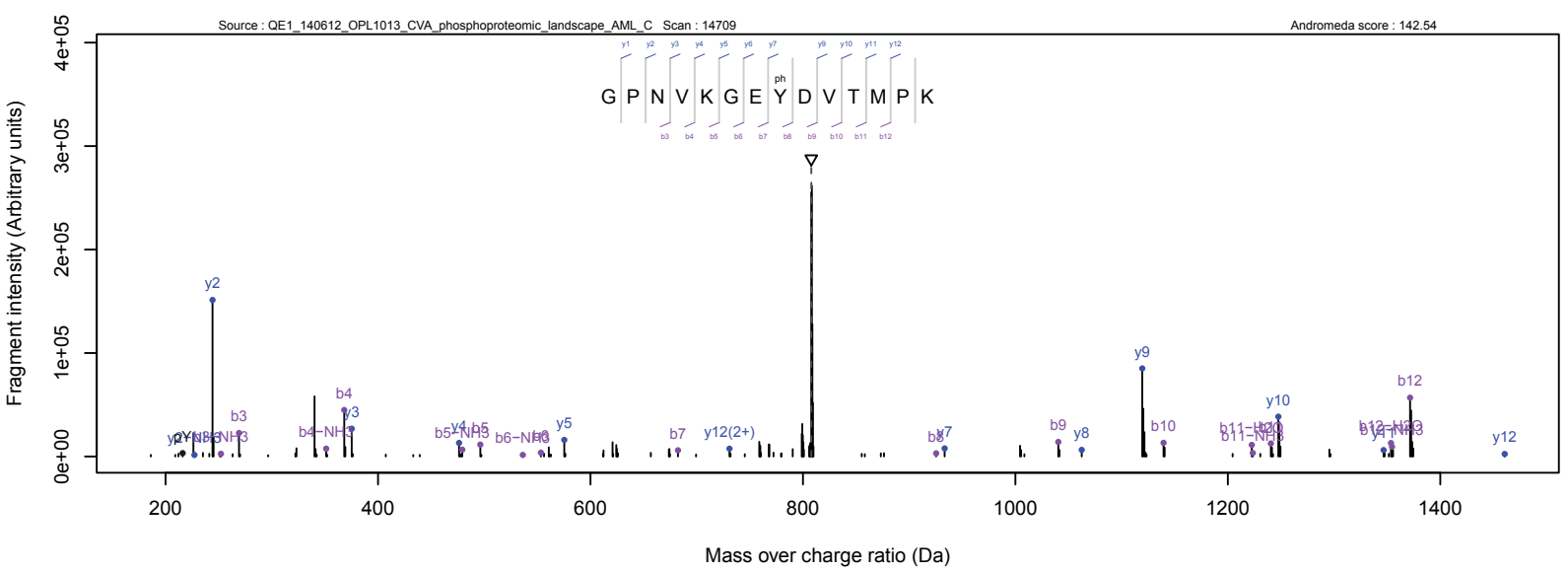
Andromeda score : 142.68



GPNVKGEY(ph)DVTMPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14709

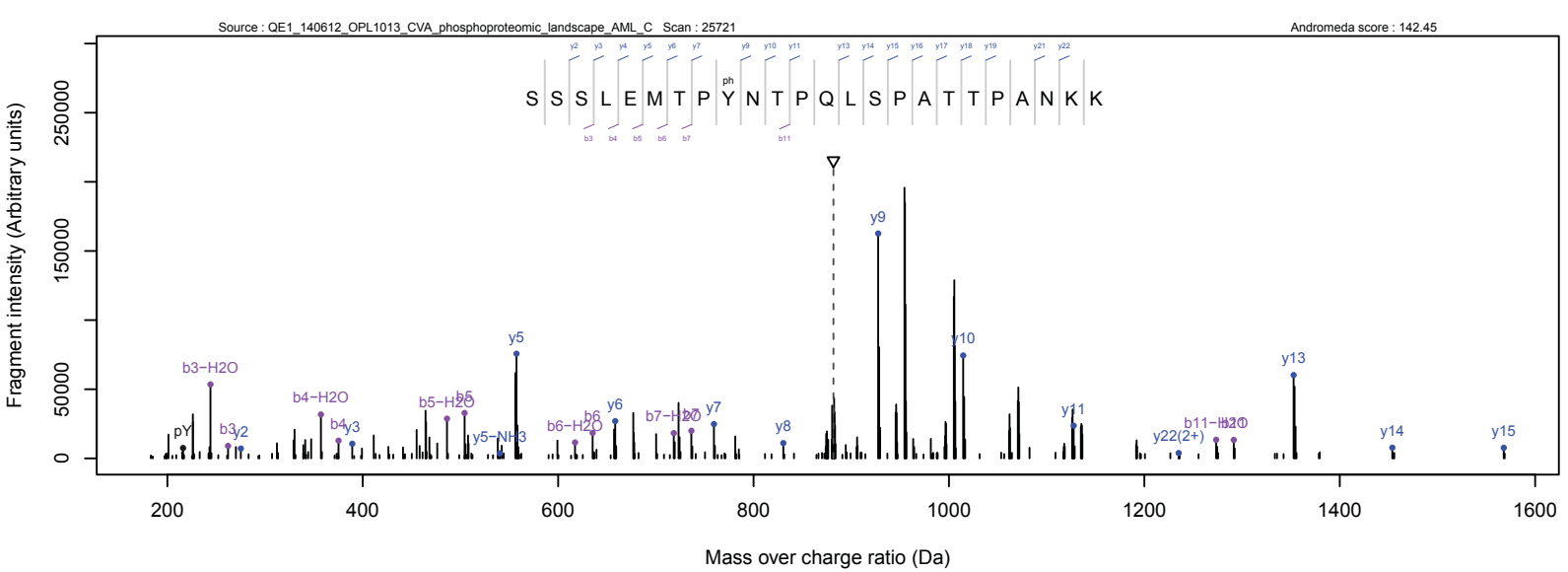
Andromeda score : 142.54



SSSLEMTPY(ph)NTPQLSPATTPANKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25721

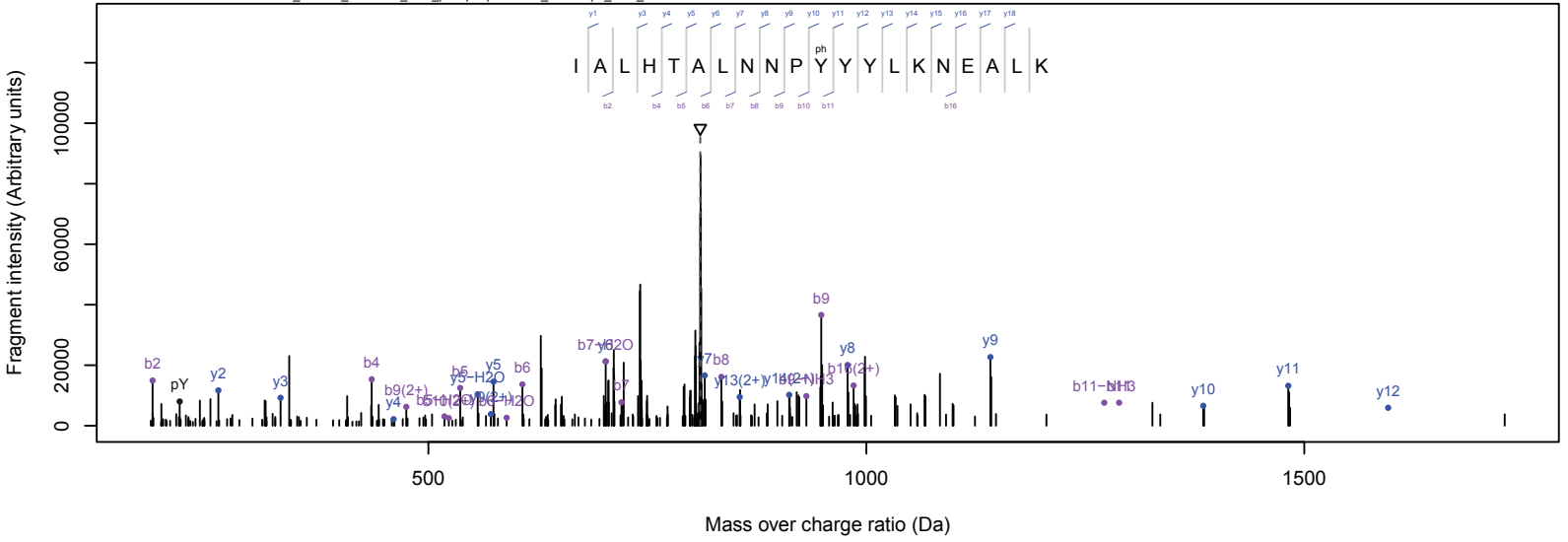
Andromeda score : 142.45



**IALHTALNNPYYYLKNEALK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 36089

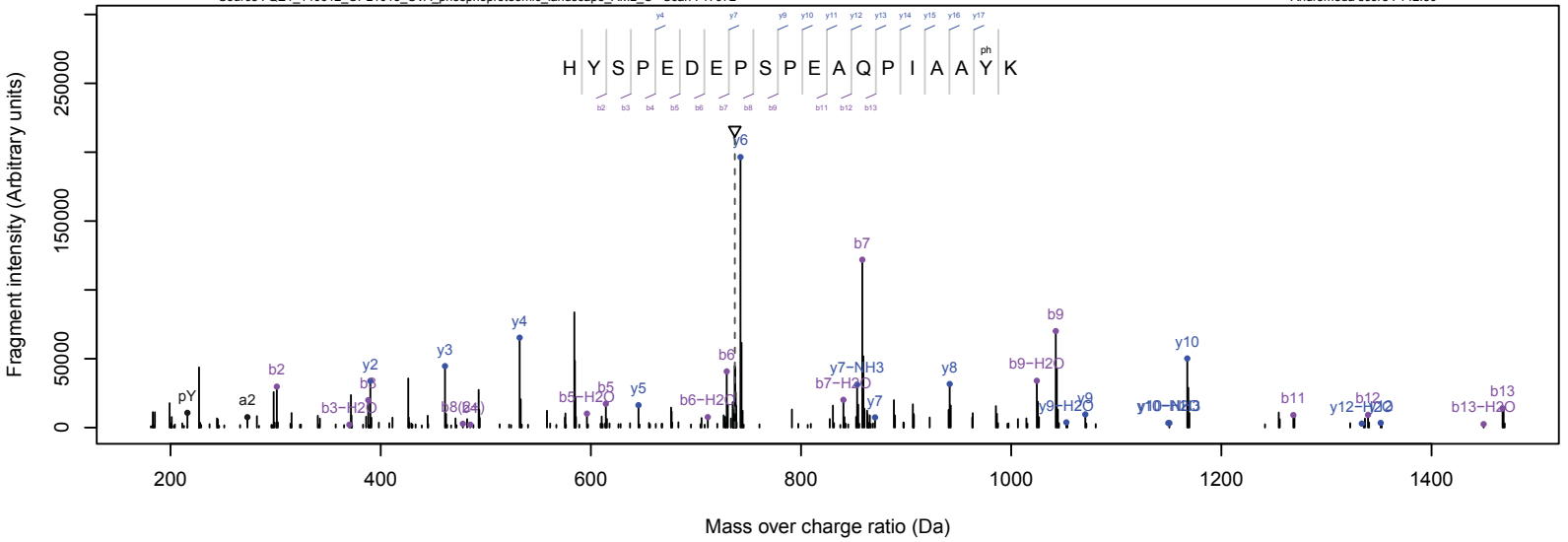
Andromeda score: 142.43



**HYSPEDEPSPEAQPIAAYK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17972

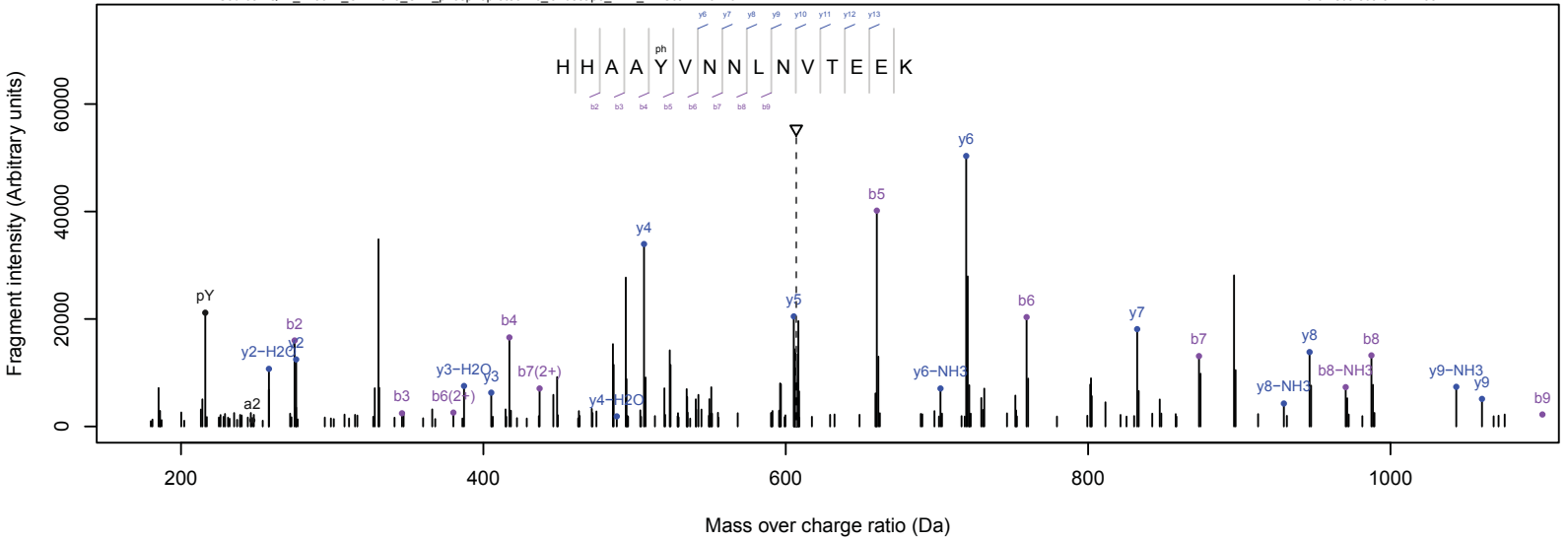
Andromeda score: 142.39



**HHAAY(V)NNLNVTEEK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 13146

Andromeda score: 142.38

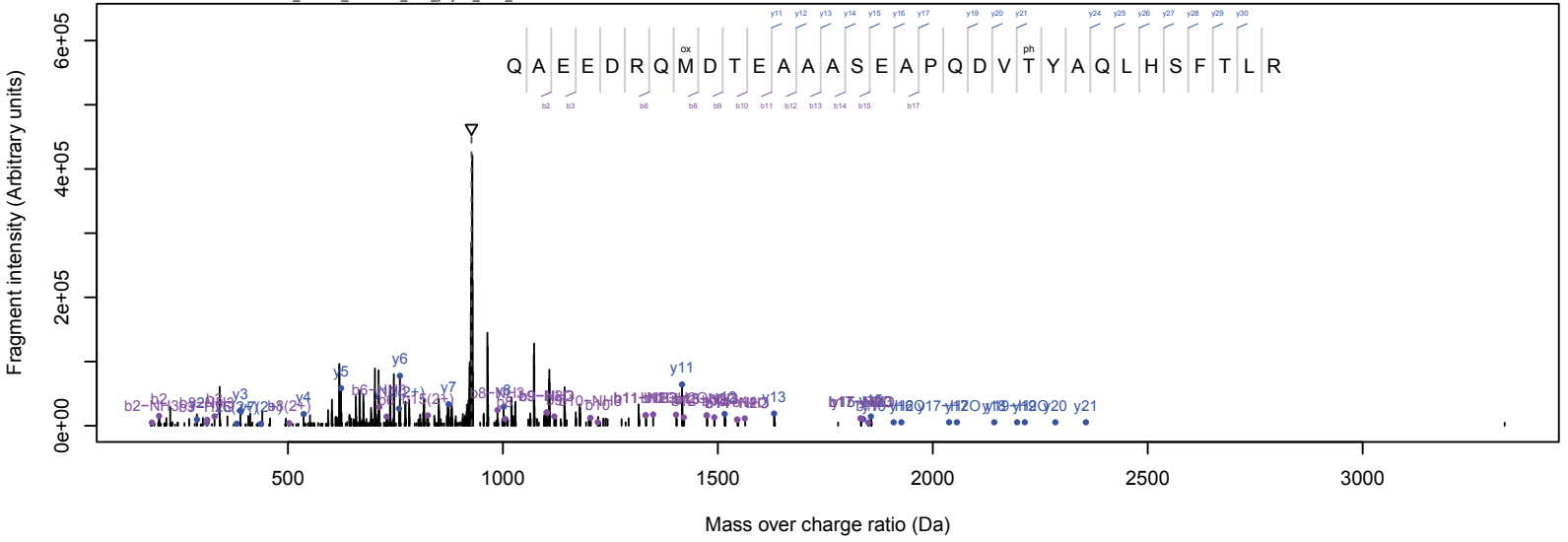




QAEDDRM(ox)DTEAAASEAPQDVT(ph)YAQLHSFTLR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 35660

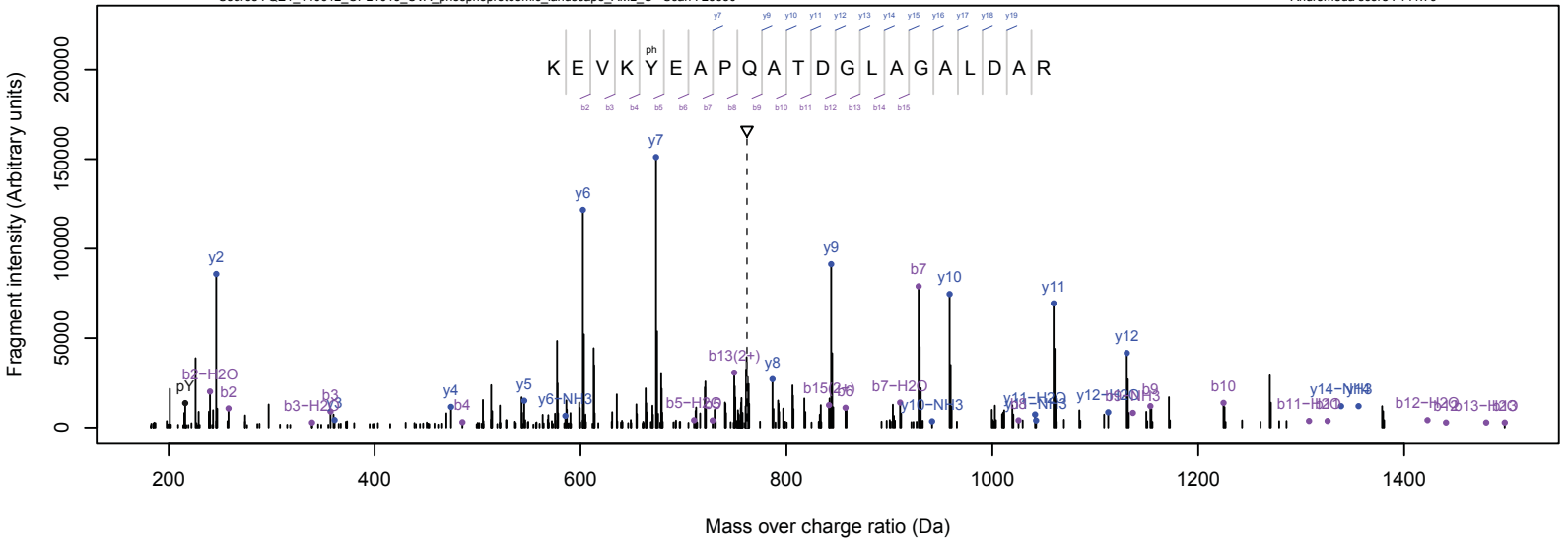
Andromeda score : 141.98



KEVKY(ph)EAPQATDGLAGALDAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23650

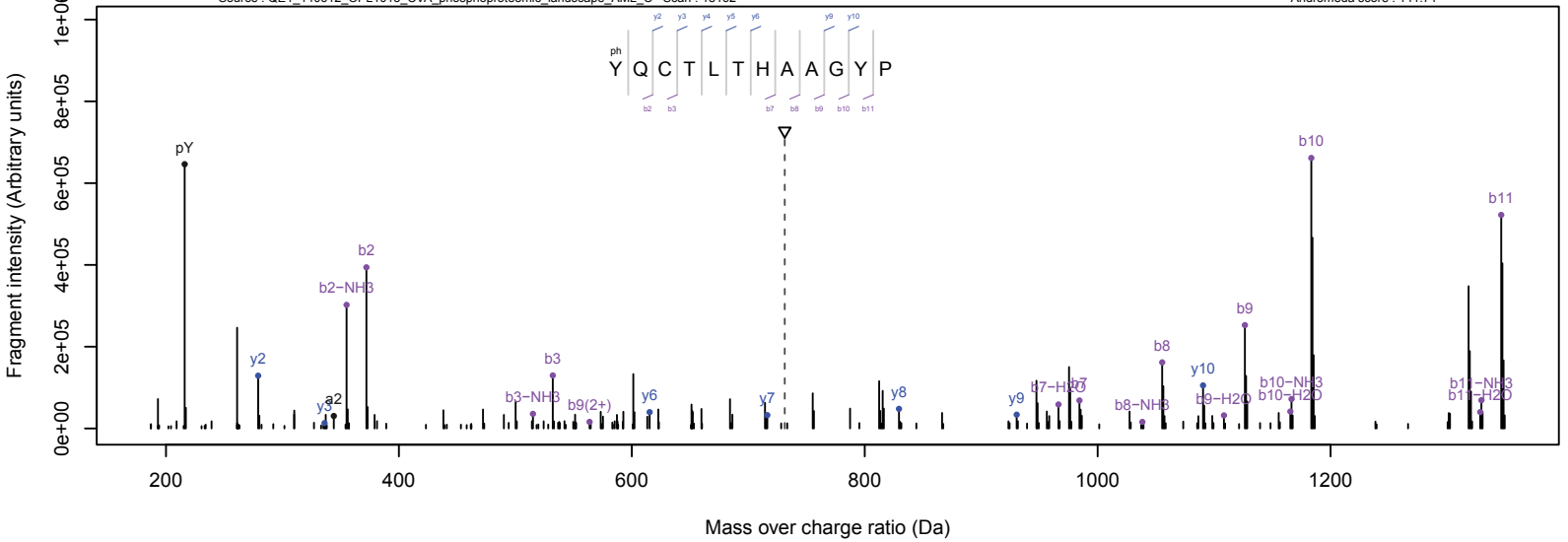
Andromeda score : 141.79



Y(ph)QCTLTHAAGYP-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18162

Andromeda score : 141.71

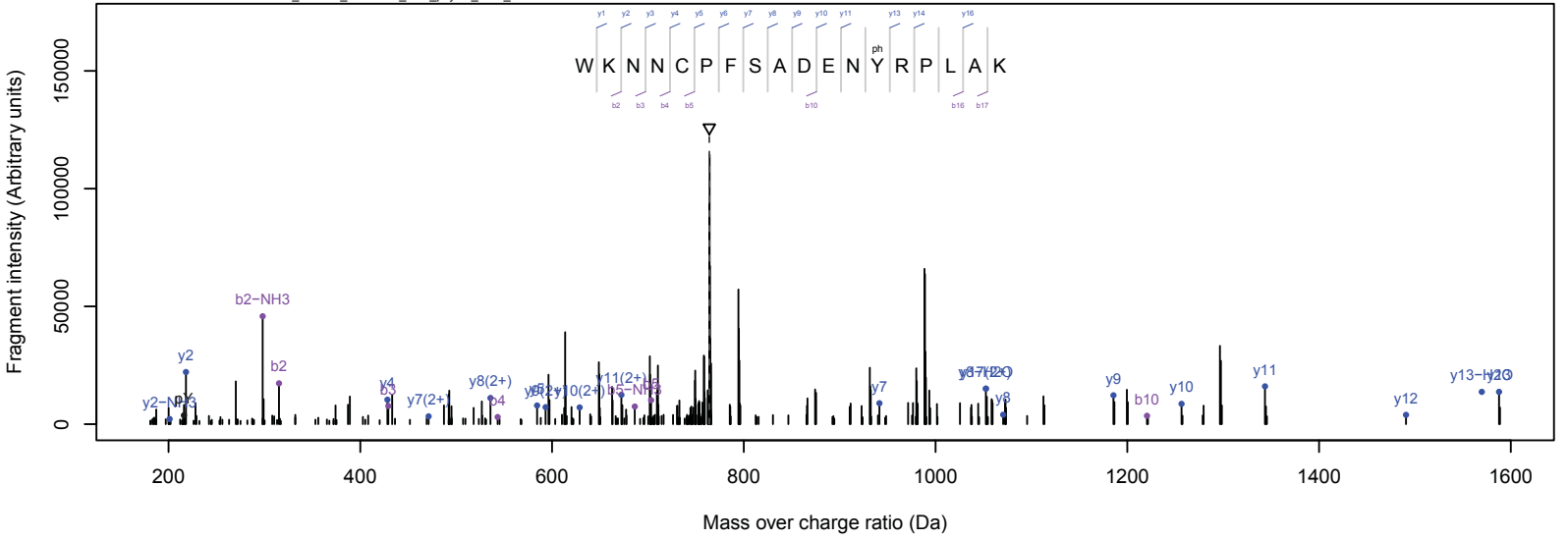




WKNNCPFSADENY<sup>ph</sup>RPLAK-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan: 21804

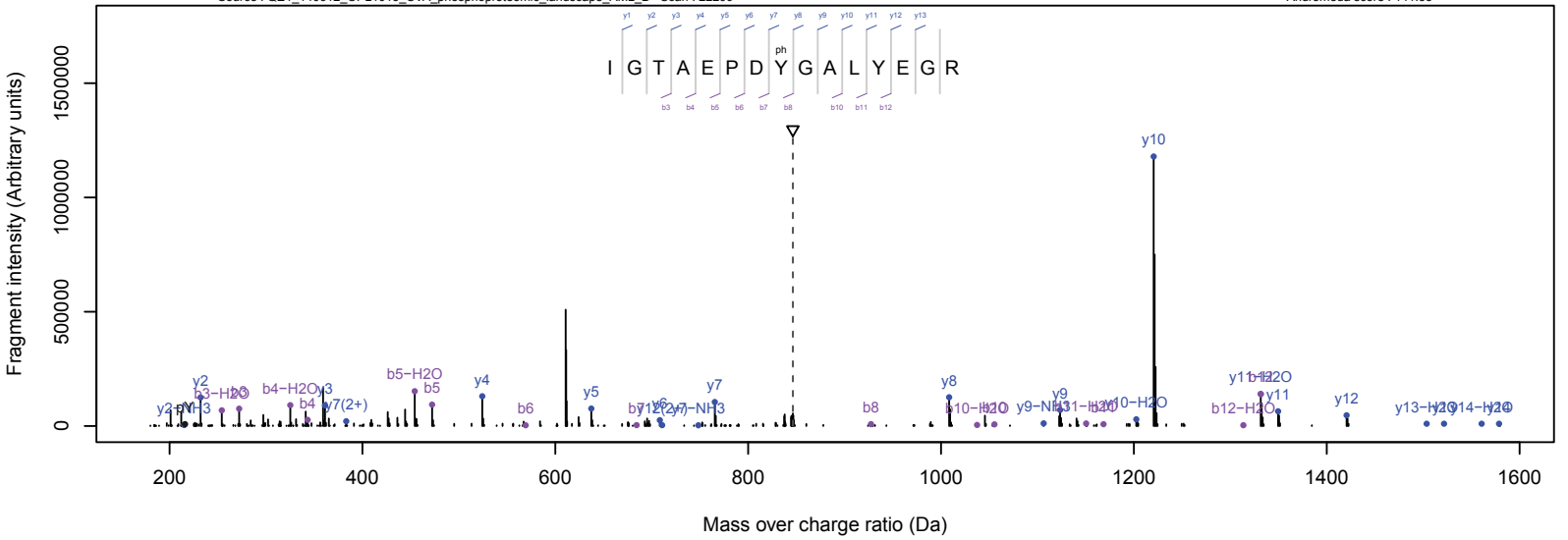
Andromeda score: 141.6



IGTAEPDY<sup>ph</sup>GALYEGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 22299

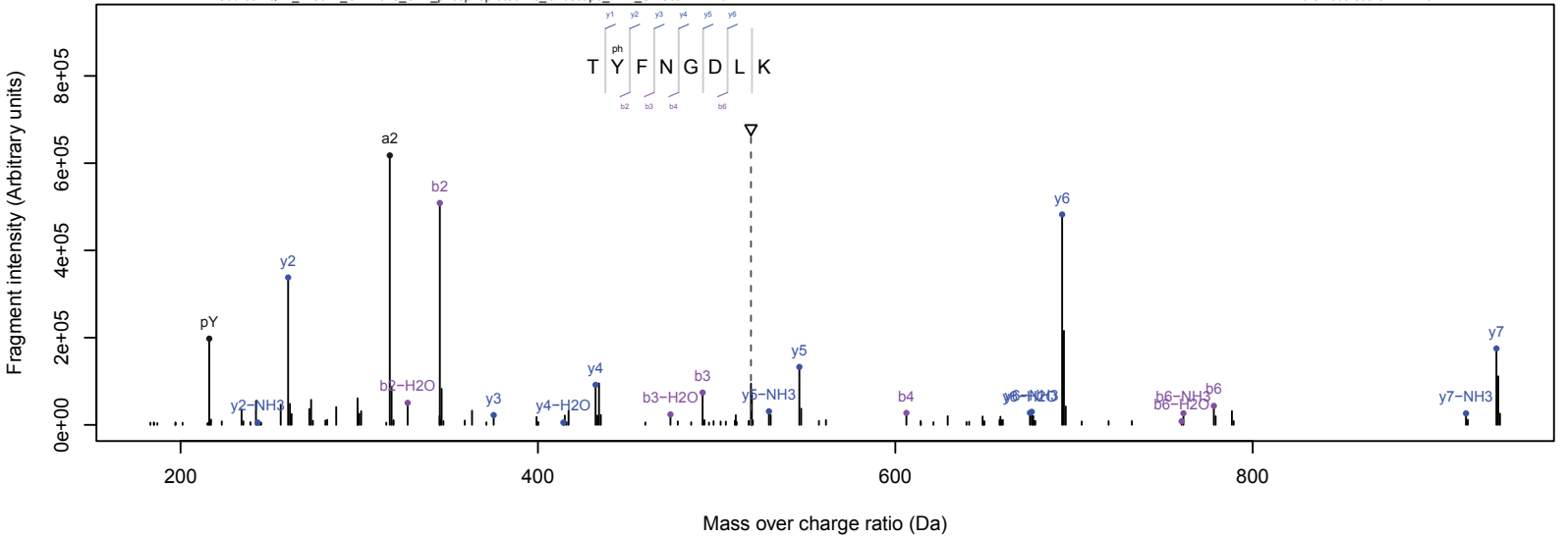
Andromeda score: 141.55



TY<sup>ph</sup>FNGDLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14101

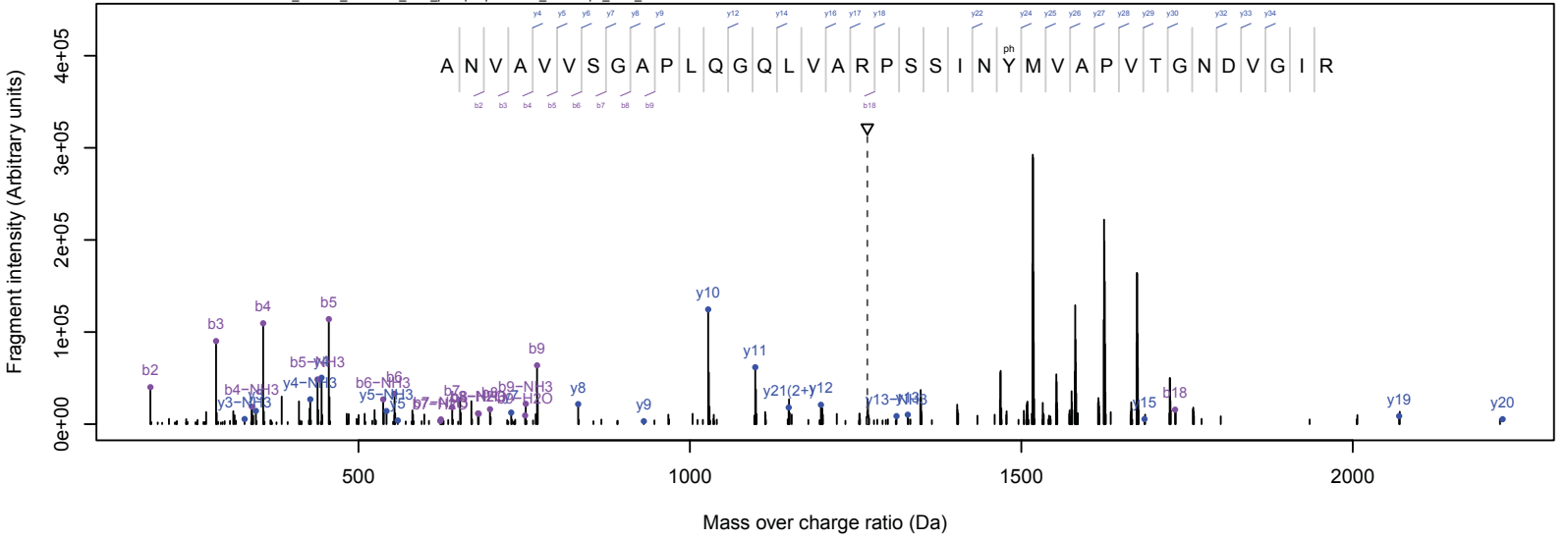
Andromeda score: 141.48



ANVAVVSGAPLQGQLVARPSSINYPH MVAPVTGNDVGIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 47017

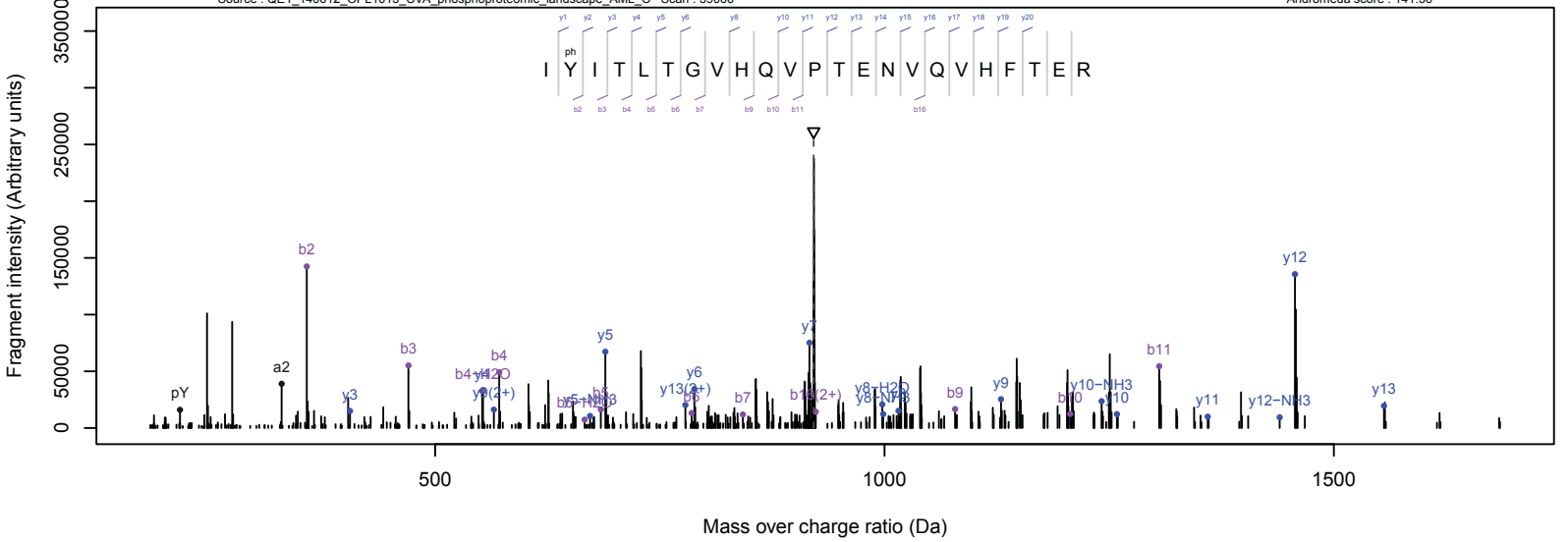
Andromeda score: 141.36



IY(pH)ILTGTVHQVPTENVQVHFTEK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 39006

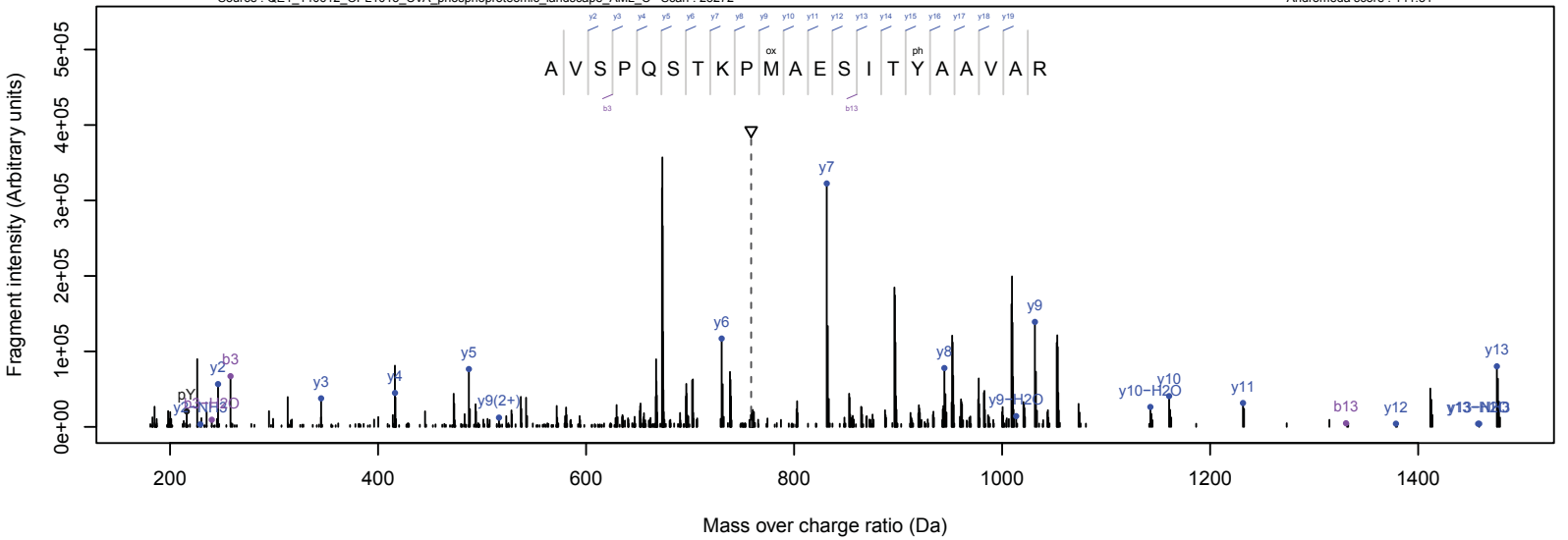
Andromeda score: 141.36



AVSPQSTKPM(ox)AESITYA AAVAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25272

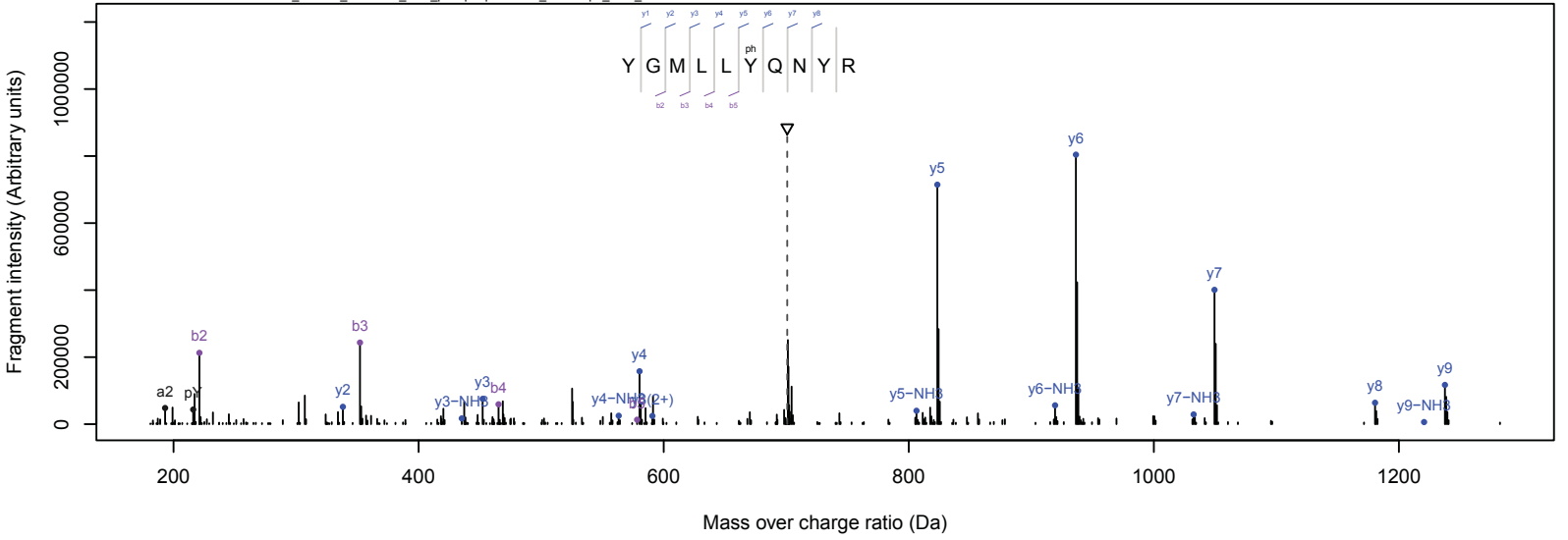
Andromeda score: 141.31



YGMLLY(ph)QNYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29738

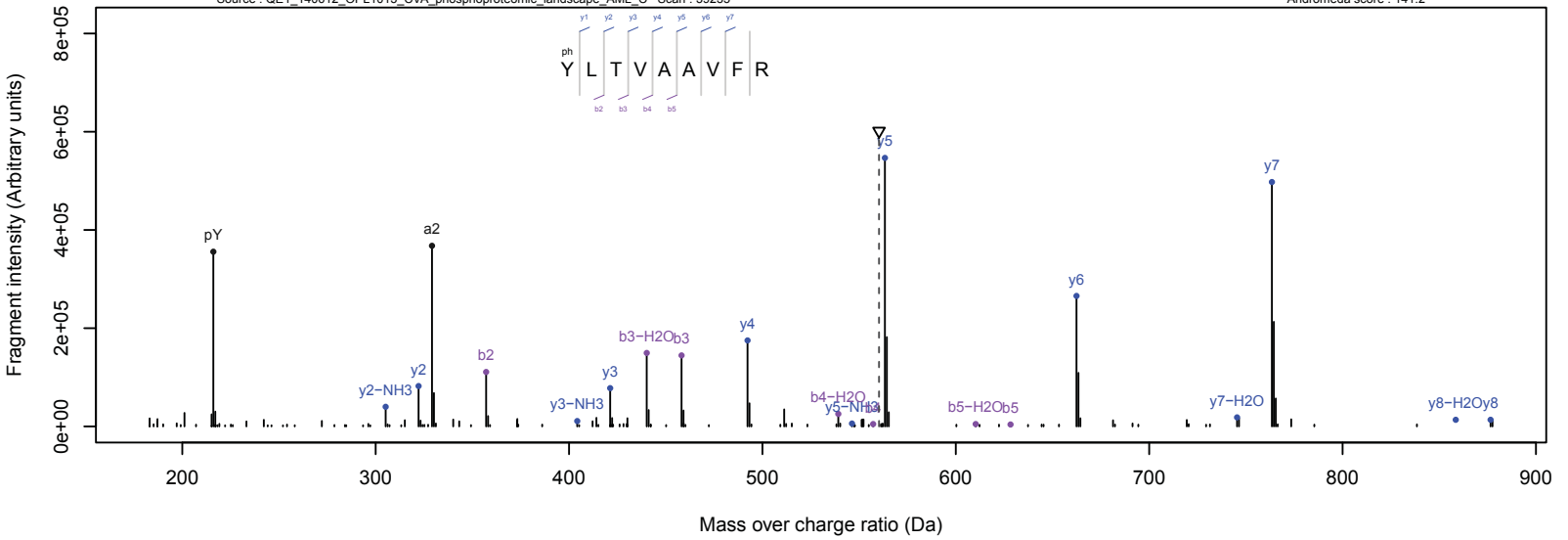
Andromeda score : 141.2



Y(ph)LTVAAVFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39235

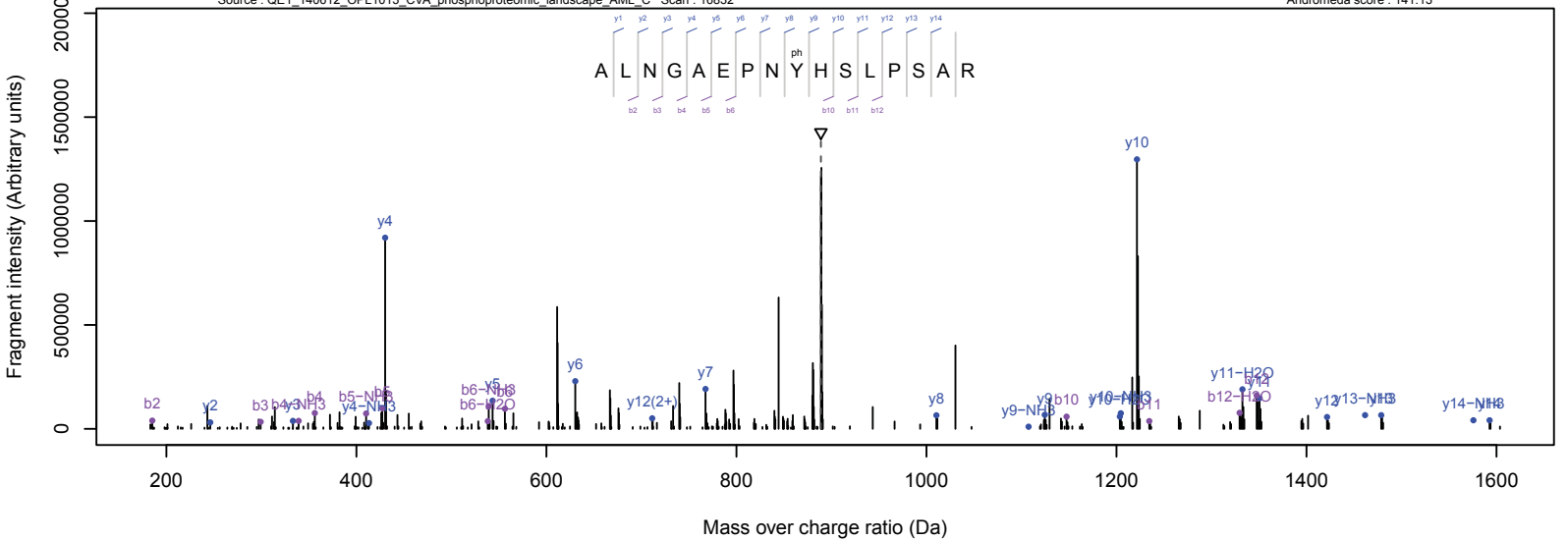
Andromeda score : 141.2



ALNGAEPNY(ph)HSLPSAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16832

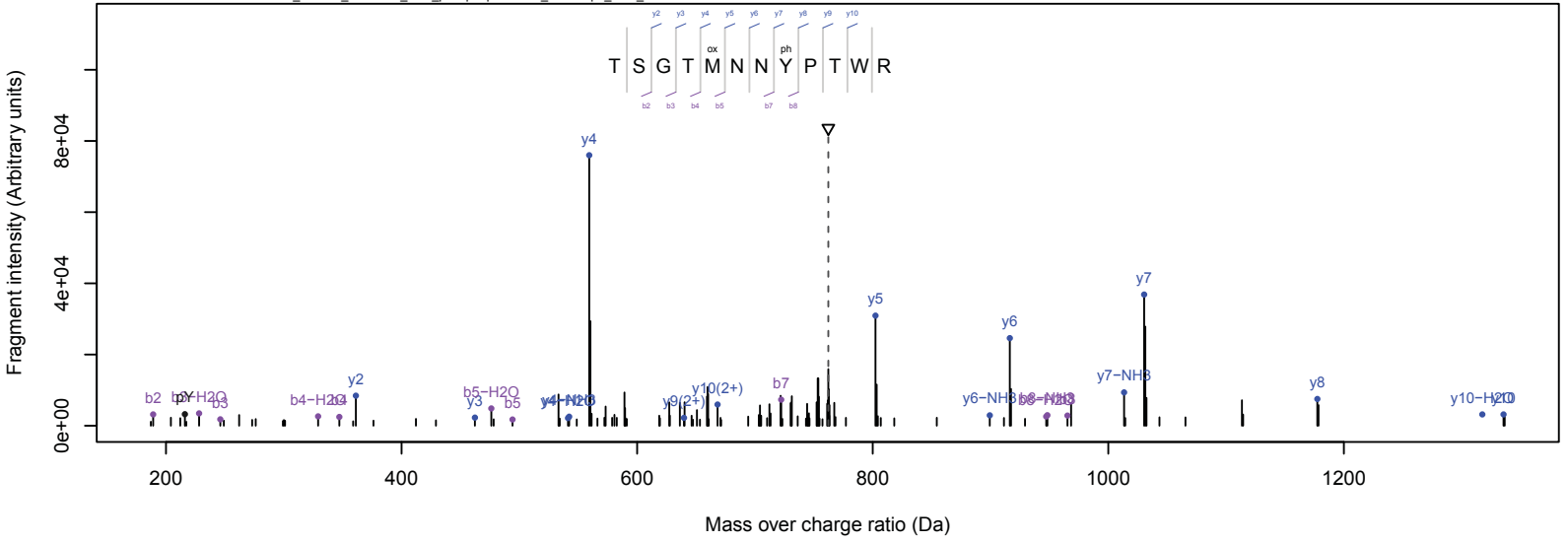
Andromeda score : 141.13



TSGTM(ox)NNY(ph)PTWR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11562

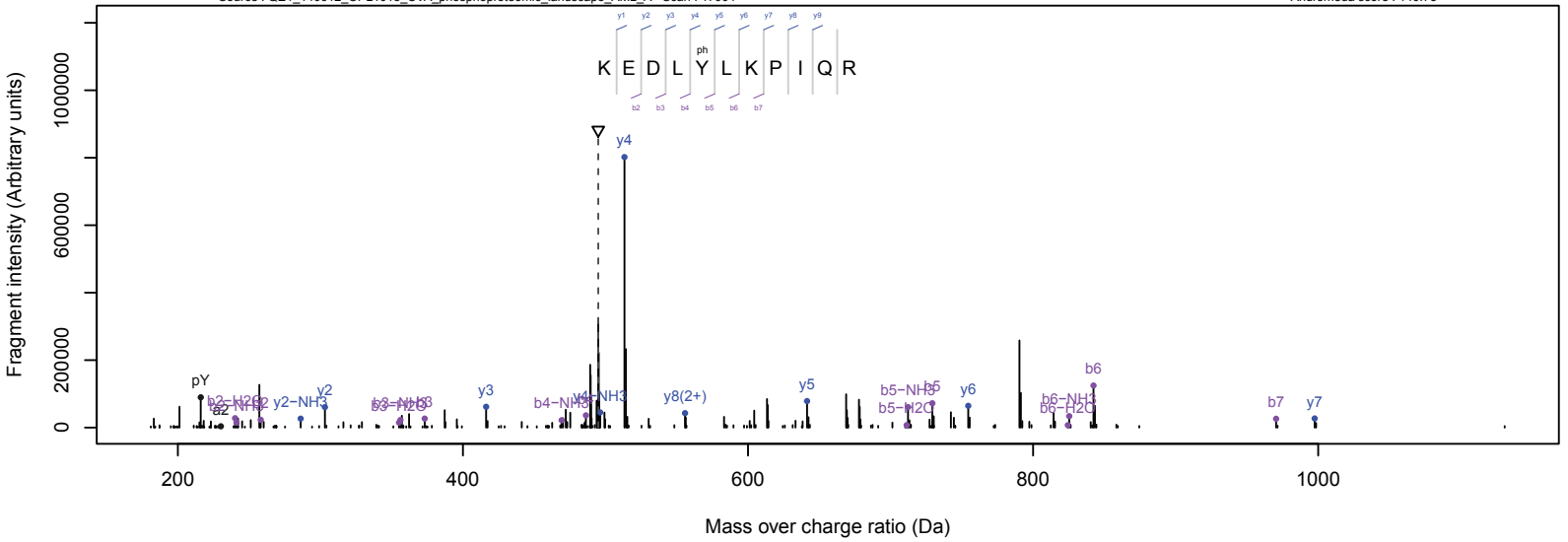
Andromeda score : 140.91



KEDLY(ph)LKPIQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17904

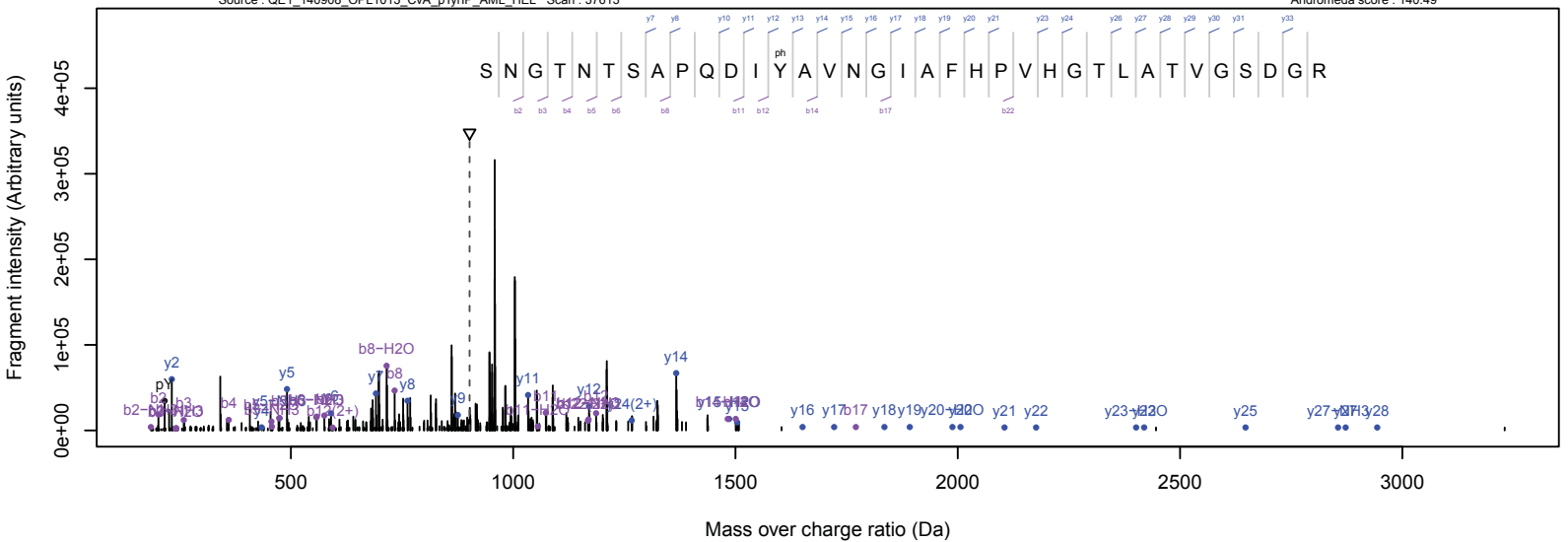
Andromeda score : 140.78



SNGTNTSAPQDIYAVNGIAFHPVHGTLATVGS DGR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37613

Andromeda score : 140.49

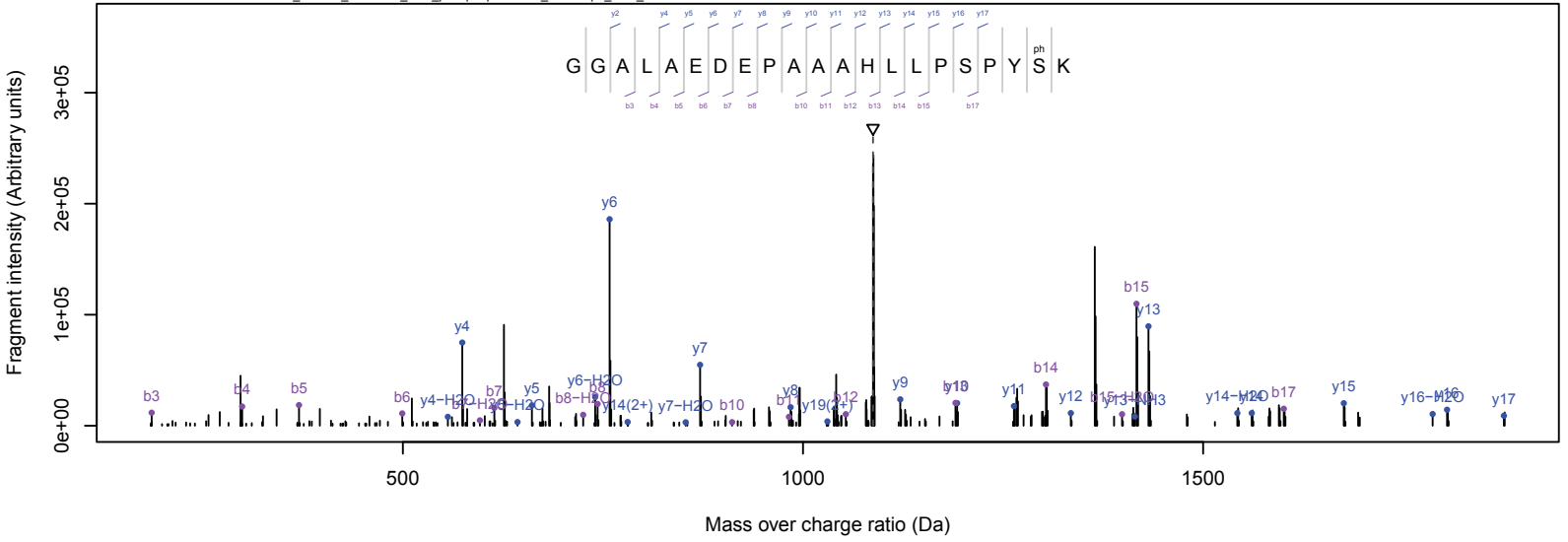




GGALAEDEPAAHLLPSPYS(ph)K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32005

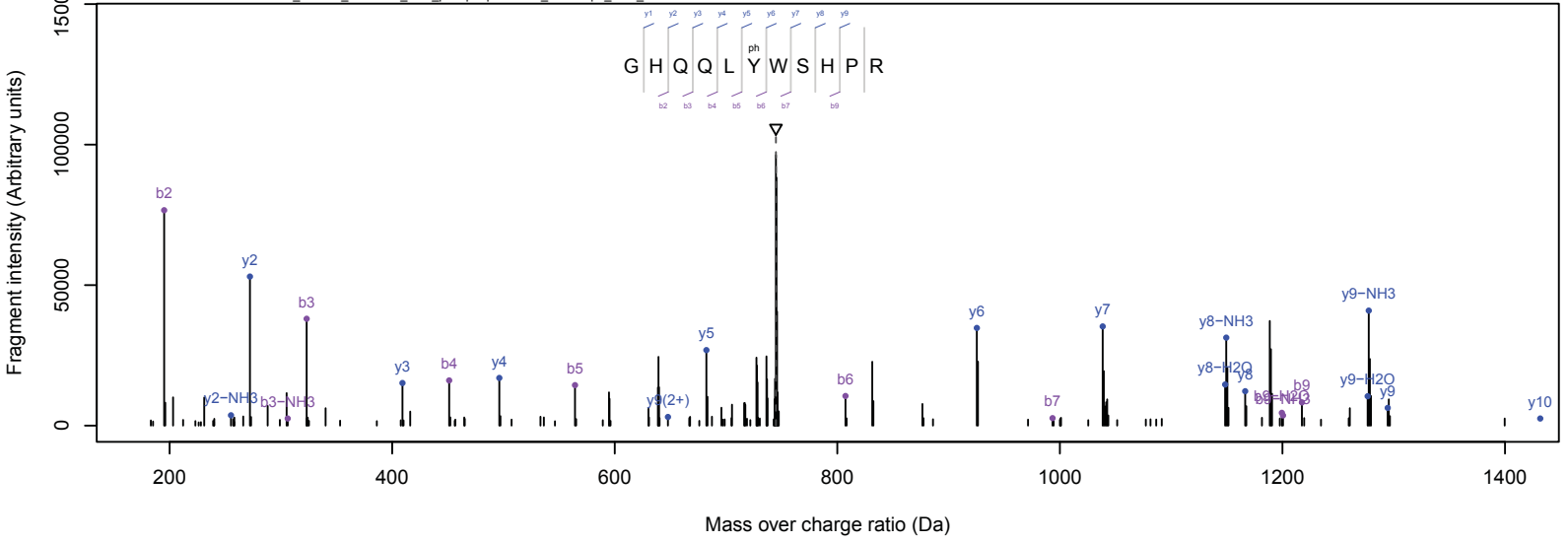
Andromeda score : 140.27



GHQQLY(ph)WSHPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12054

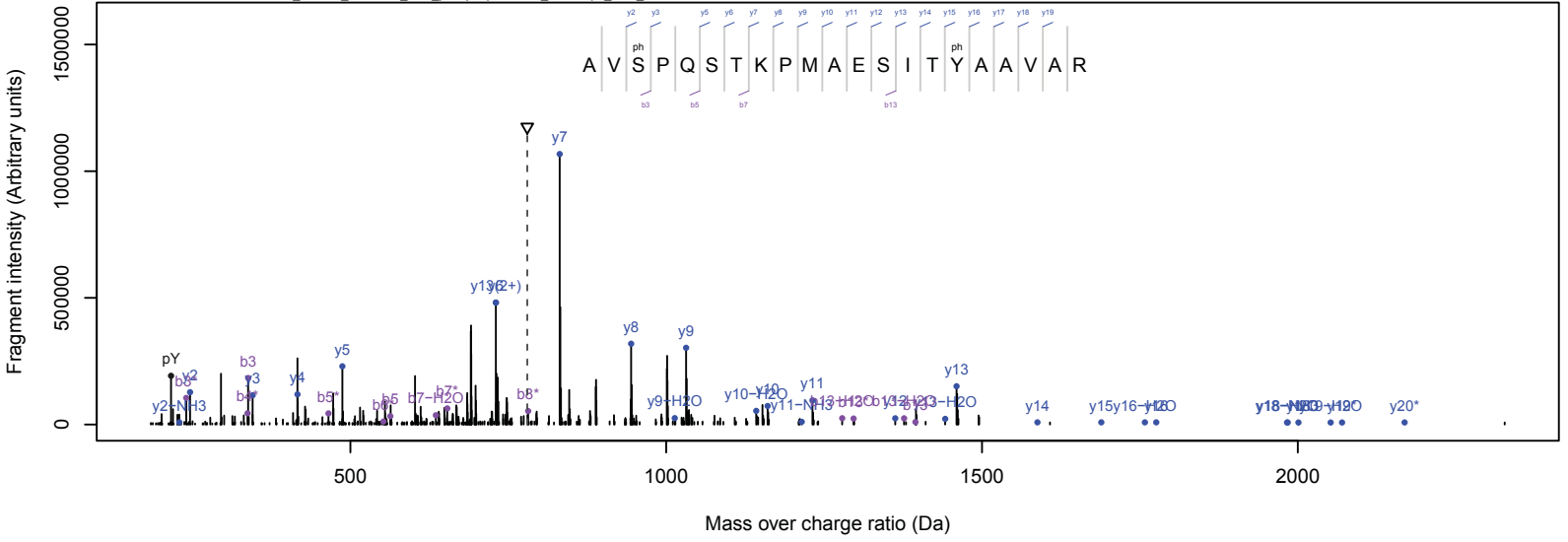
Andromeda score : 140.24



AVS(ph)PQSTKPMAESITYA(ph)AVAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31070

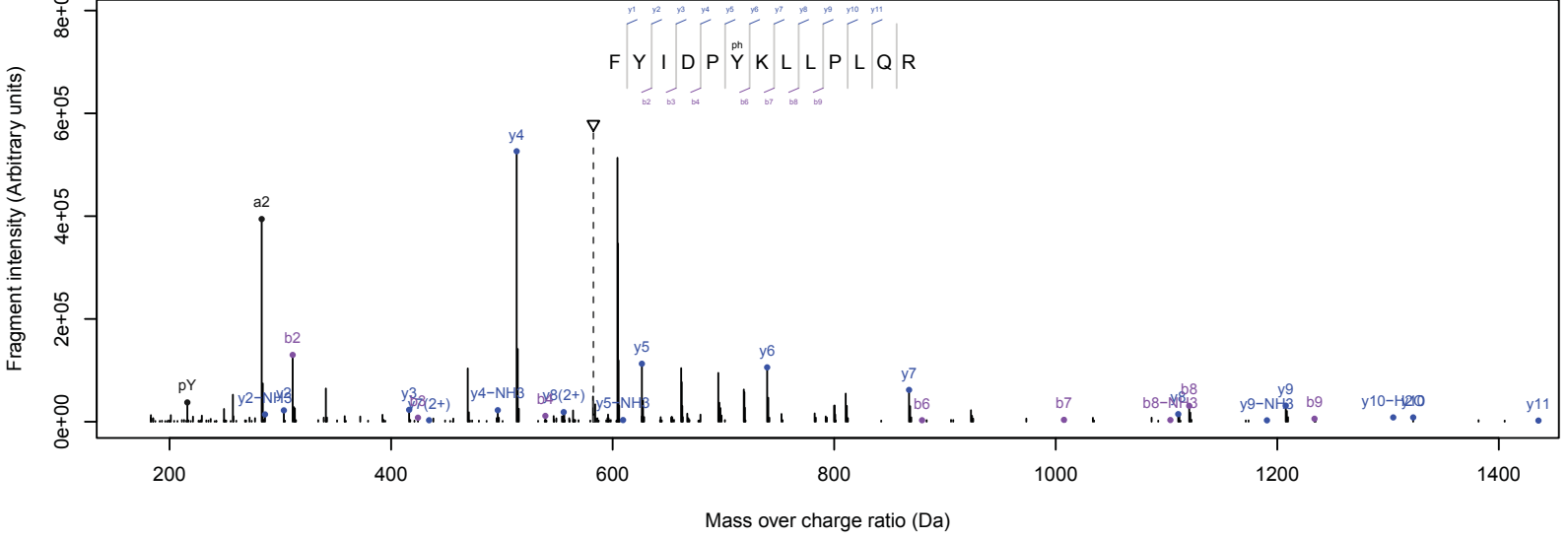
Andromeda score : 140.21



**FYIDPY(ph)KLLPLQR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 48195

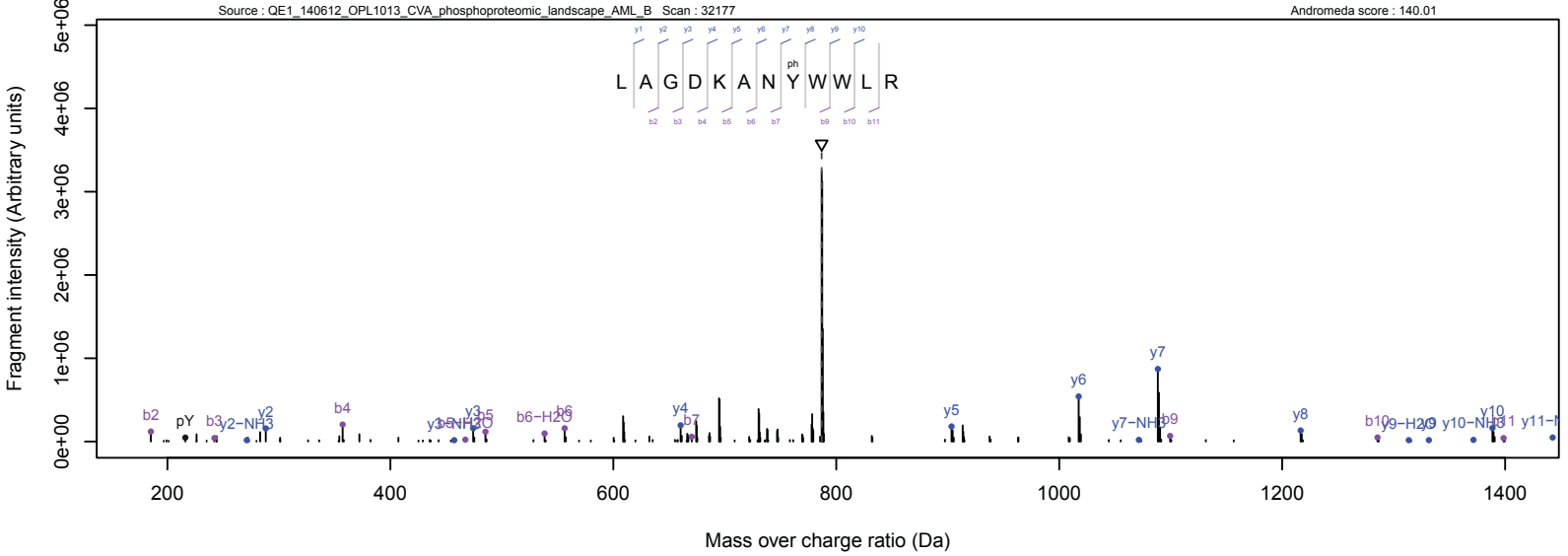
Andromeda score: 140.01



**LAGDKANY(ph)WWLR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 32177

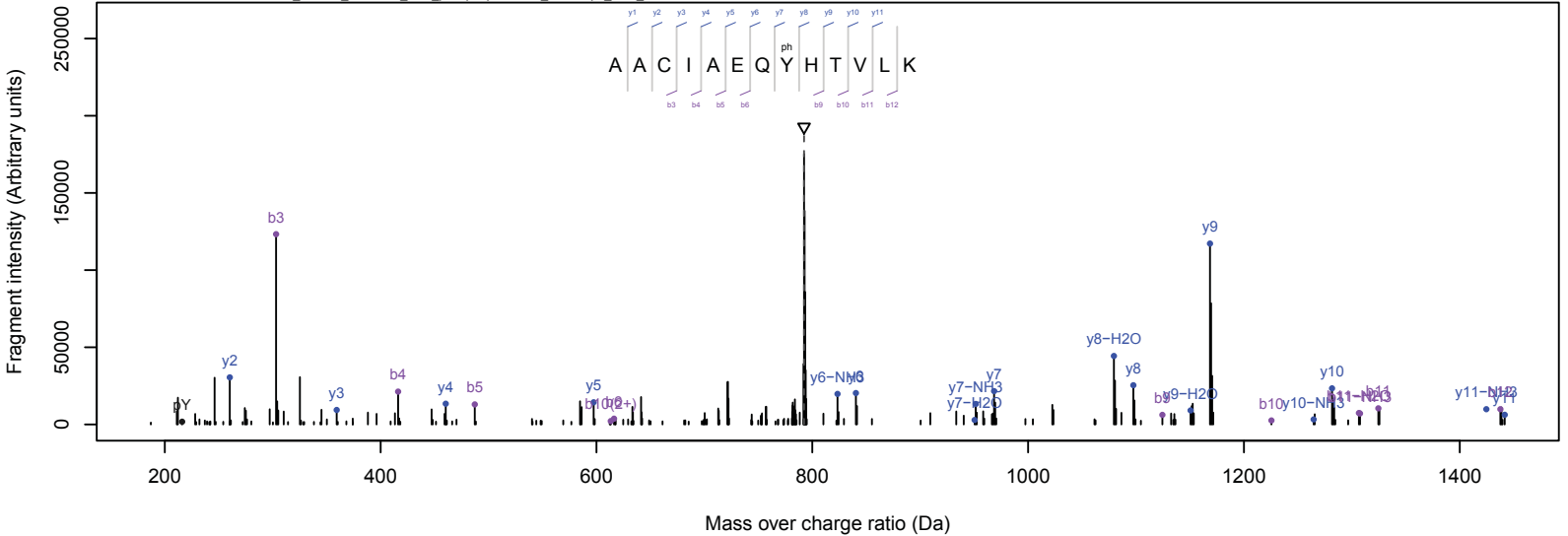
Andromeda score: 140.01



**AACIAEQY(ph)HTVLK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16700

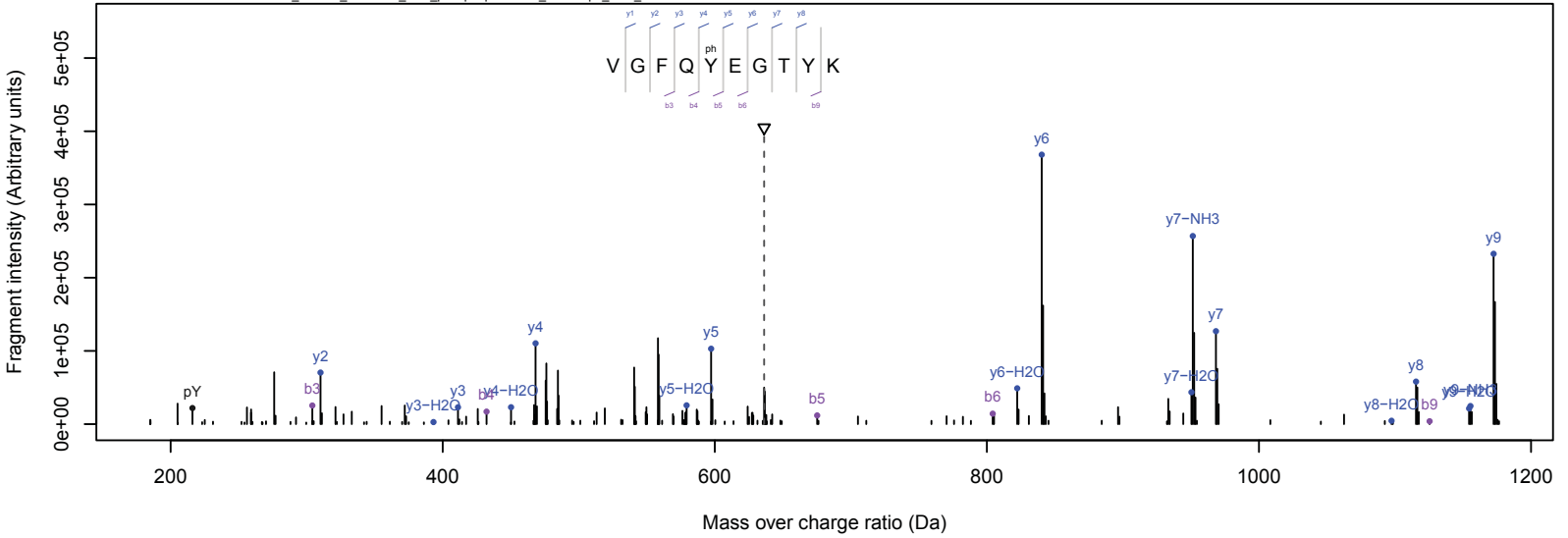
Andromeda score: 139.86



**VG FQ Y(ph) E G T Y K**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 15775

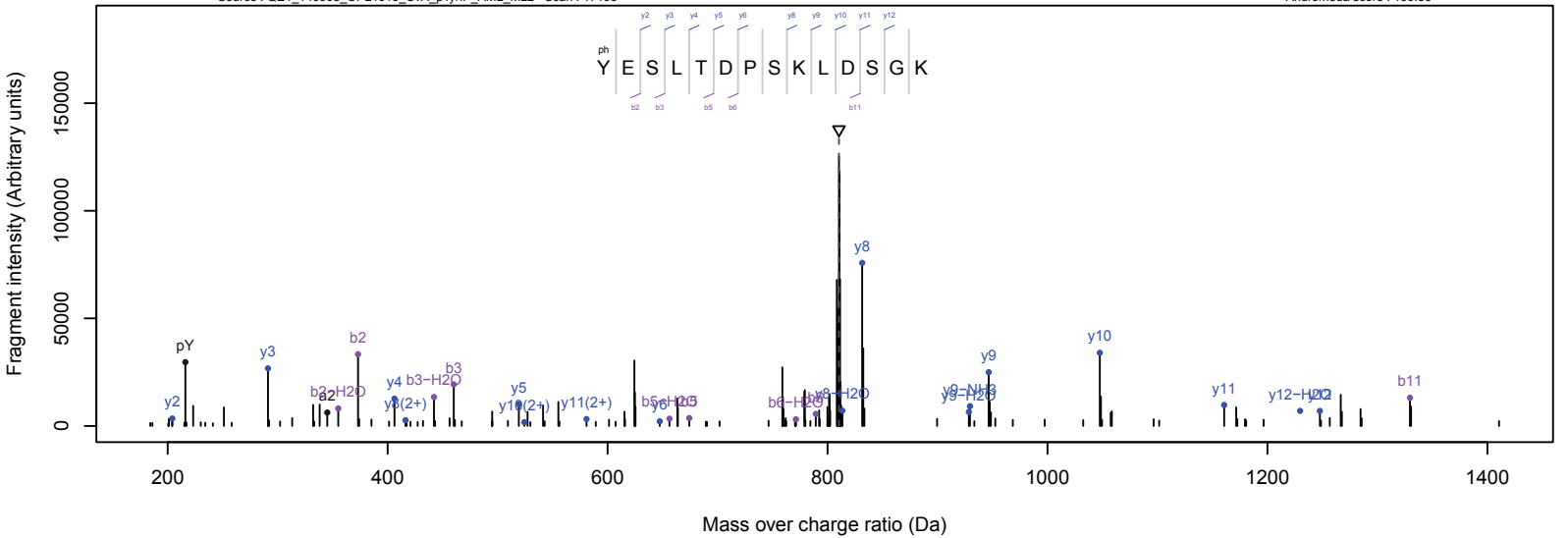
Andromeda score : 139.86



**Y(ph) E S L T D P S K L D S G K**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 17198

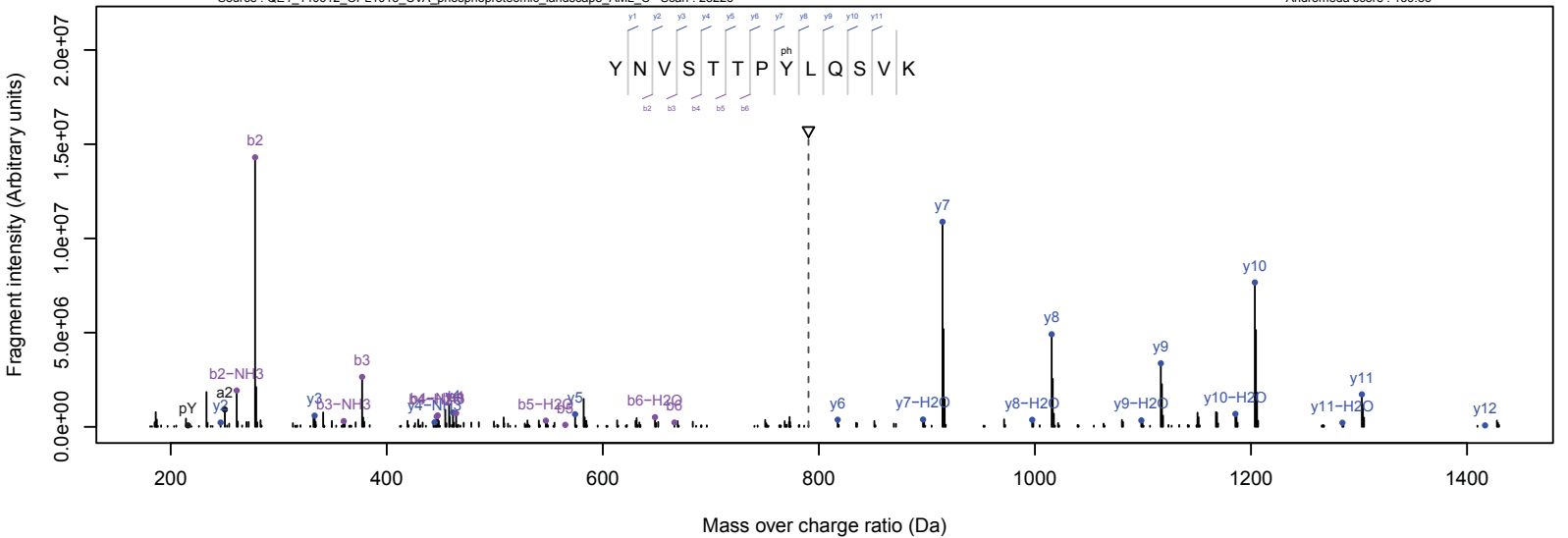
Andromeda score : 139.86



**Y N V S T T P Y L Q S V K**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23226

Andromeda score : 139.86

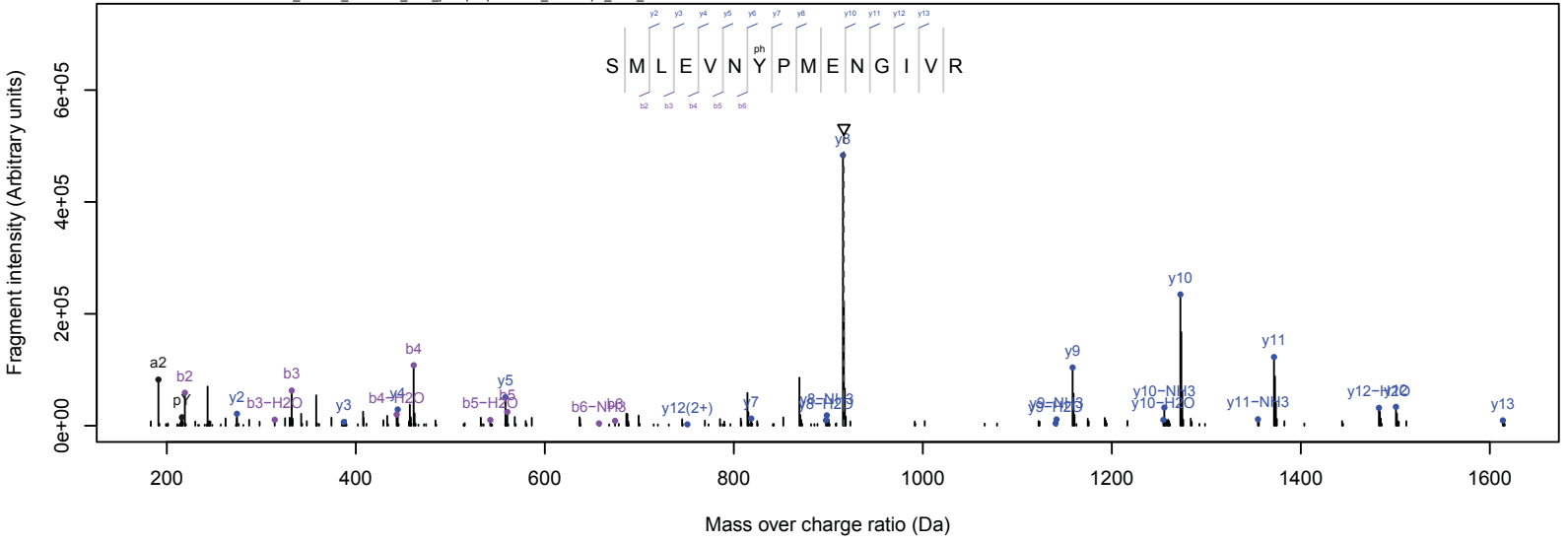




SMLEVNY(ph)PMENGIVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 35390

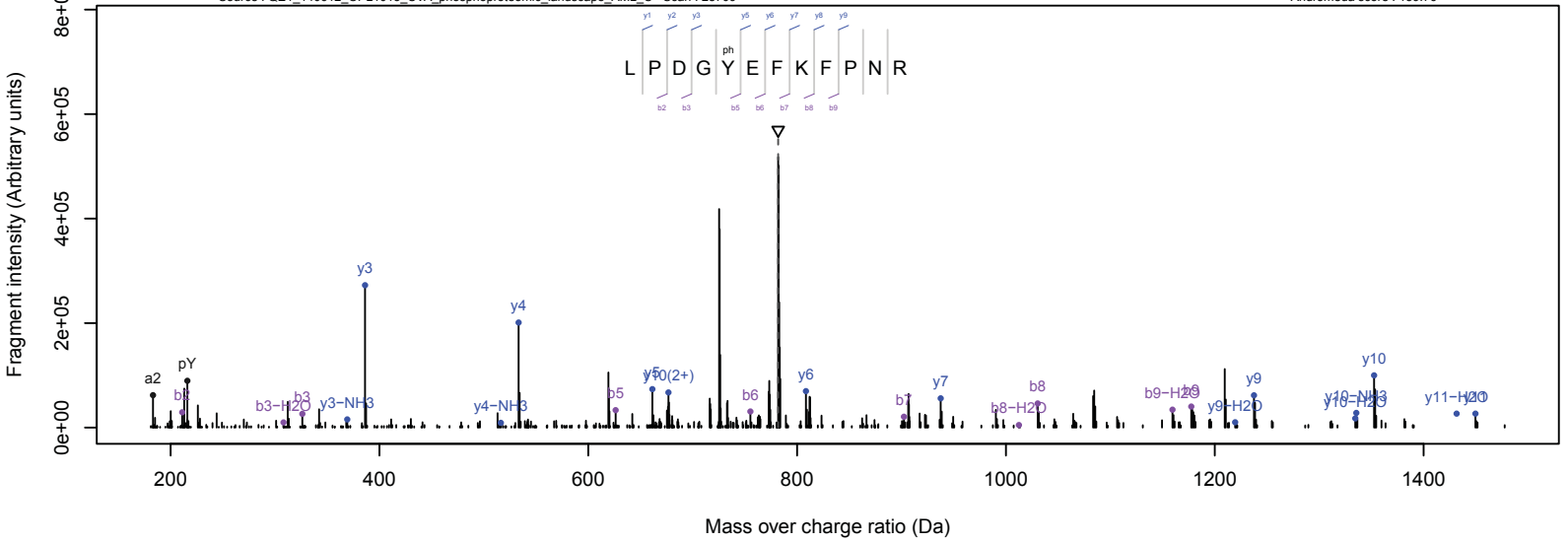
Andromeda score: 139.77



LPDGY(ph)EFKFPNR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 28709

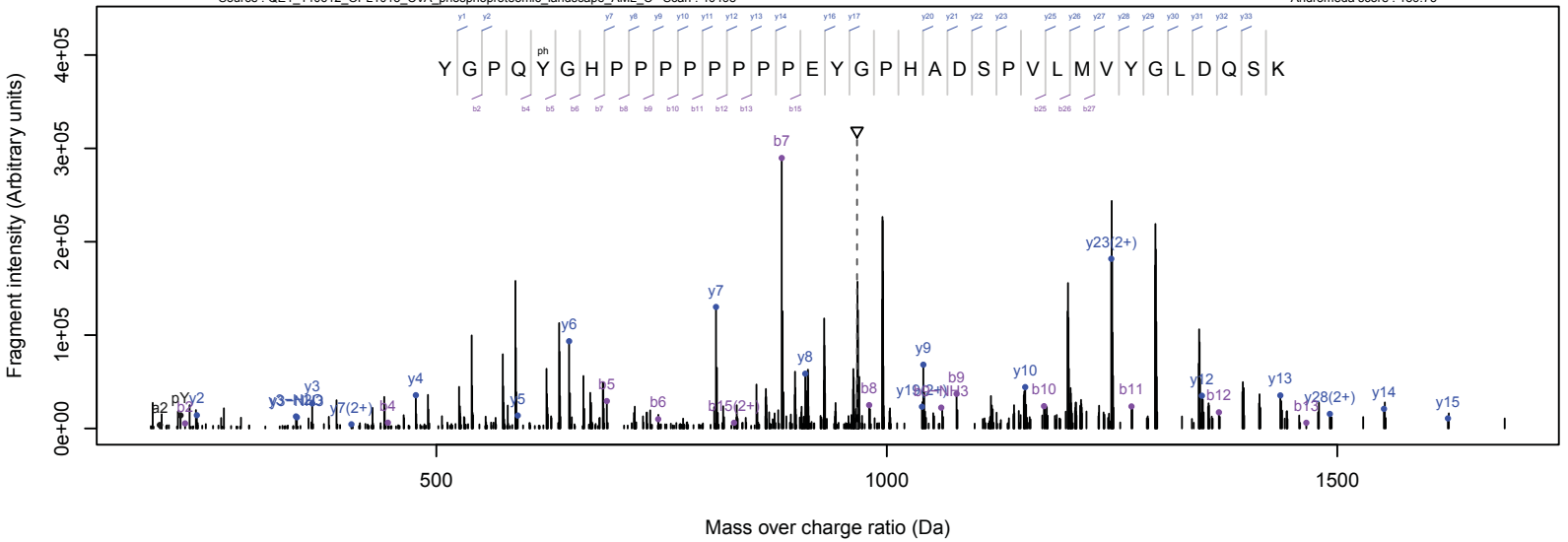
Andromeda score: 139.76



YGPQY(ph)GHPPPPPPPEYGHADSPVLMVYGLDQSK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 40498

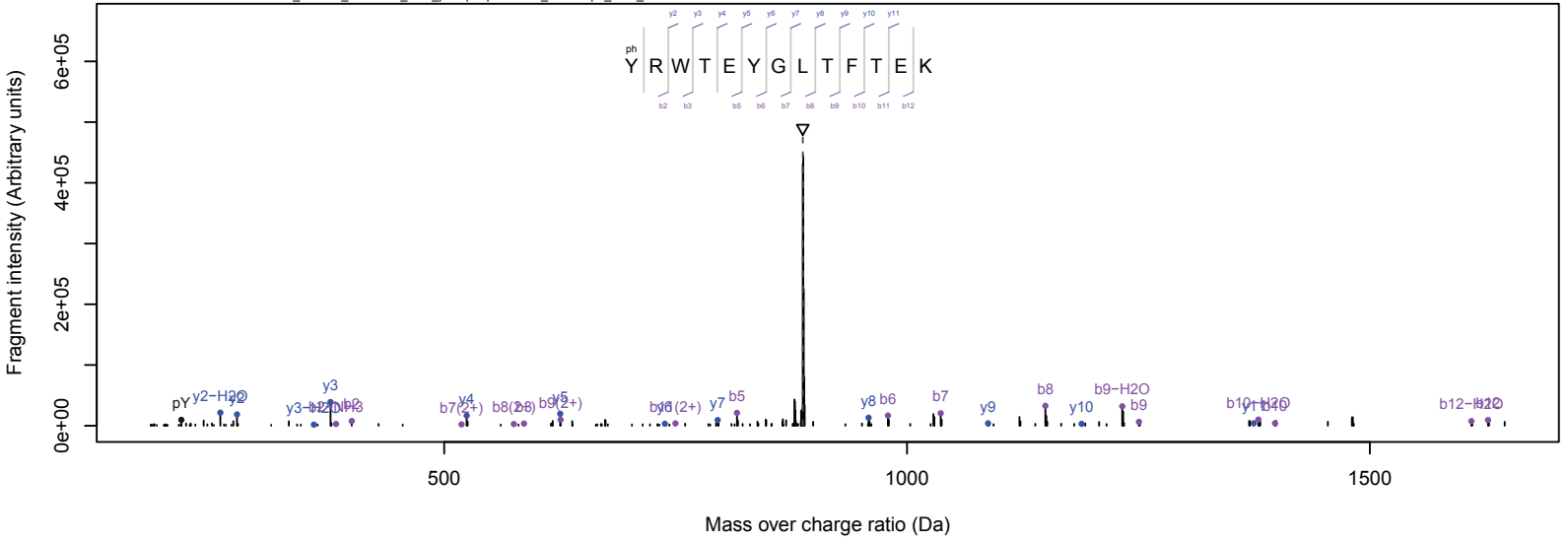
Andromeda score: 139.75



**Y(ph)RWTEYGLTFTEK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32811

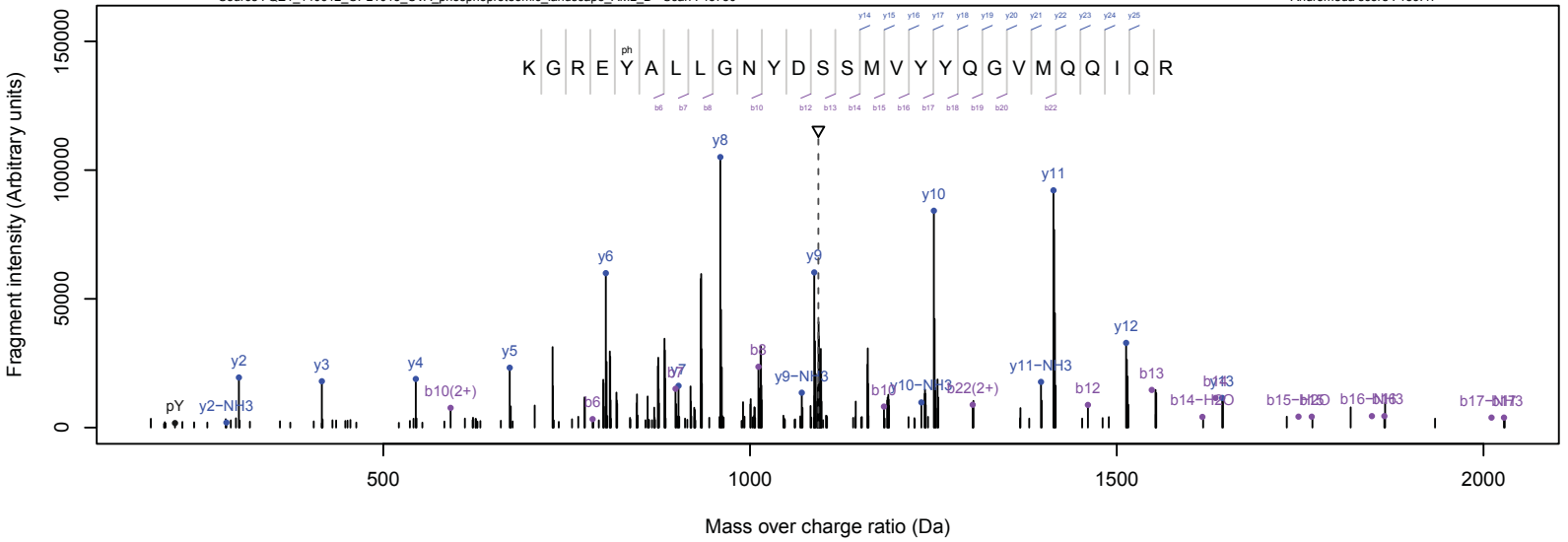
Andromeda score : 139.66



**KGREY(ph)ALLGNYSMYYQGVMQIQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 43750

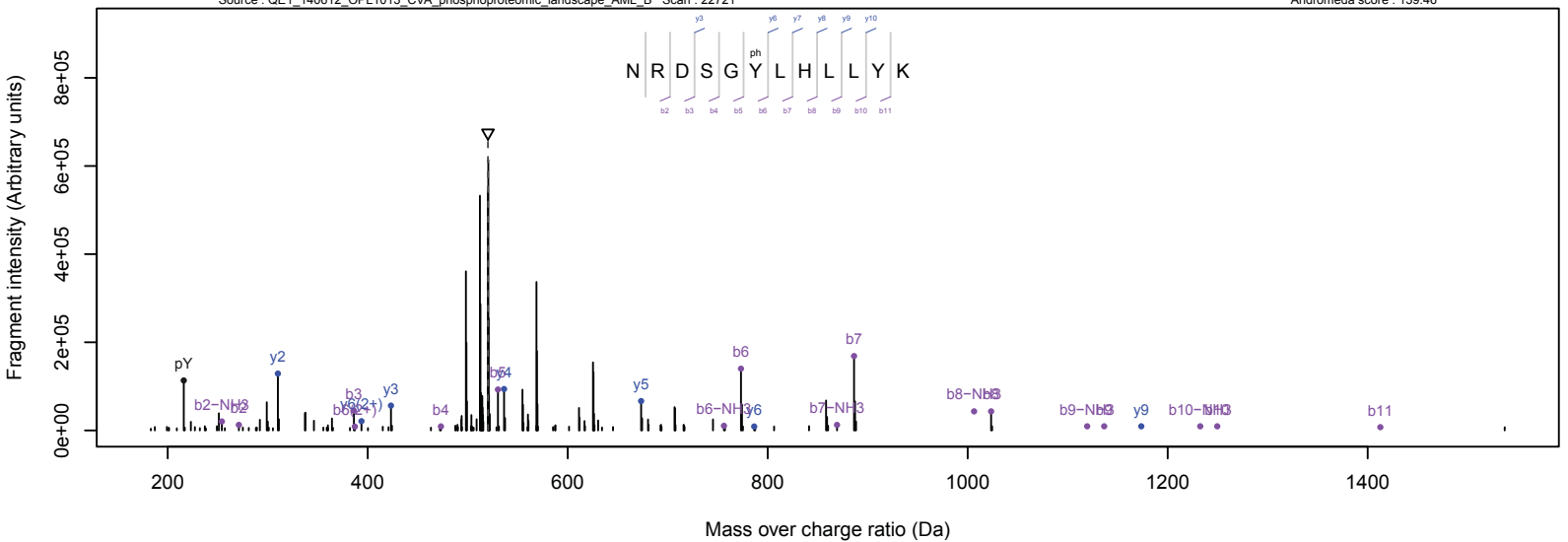
Andromeda score : 139.47



**NRDSGY(ph)LHLLYK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22721

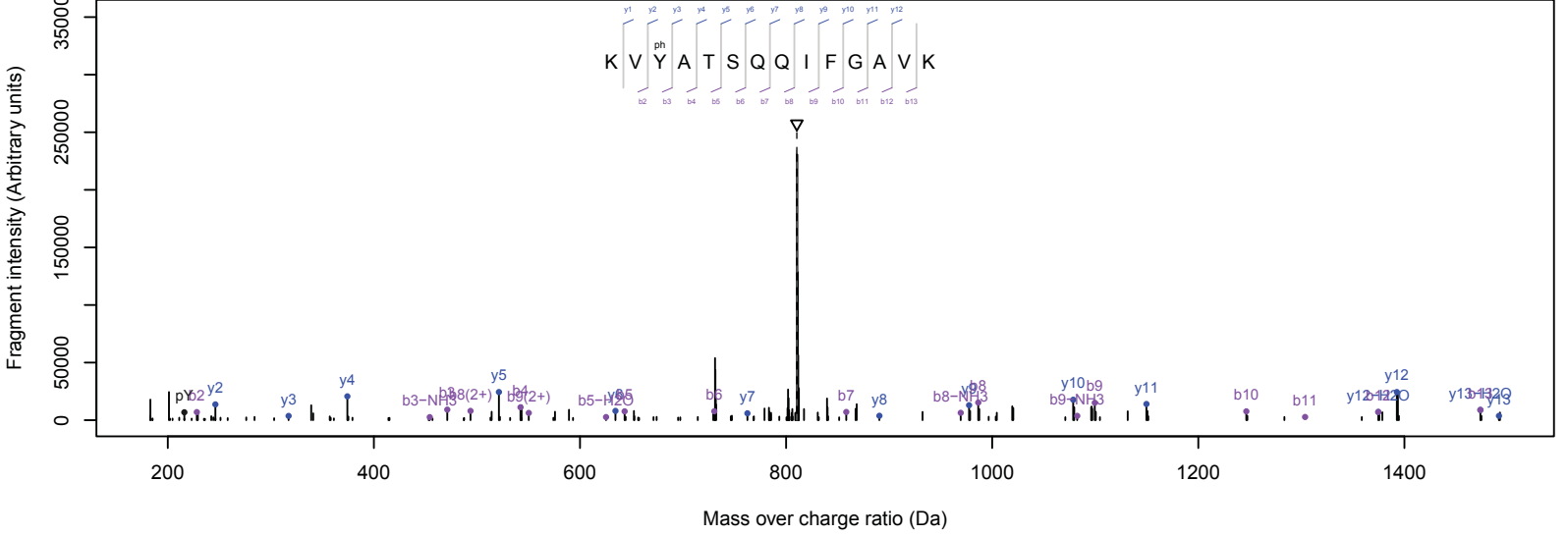
Andromeda score : 139.46



KVY(ph)ATSQQIFGAVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23492

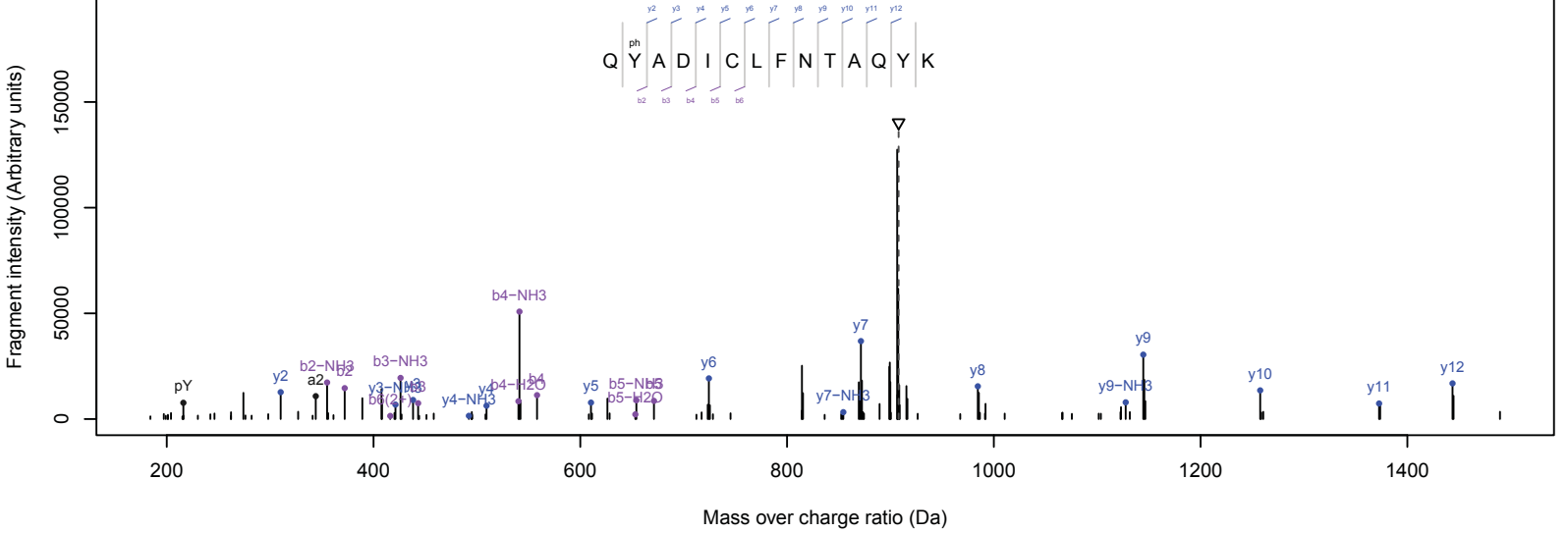
Andromeda score : 139.34



QY(ph)ADICLFNTAQYK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37969

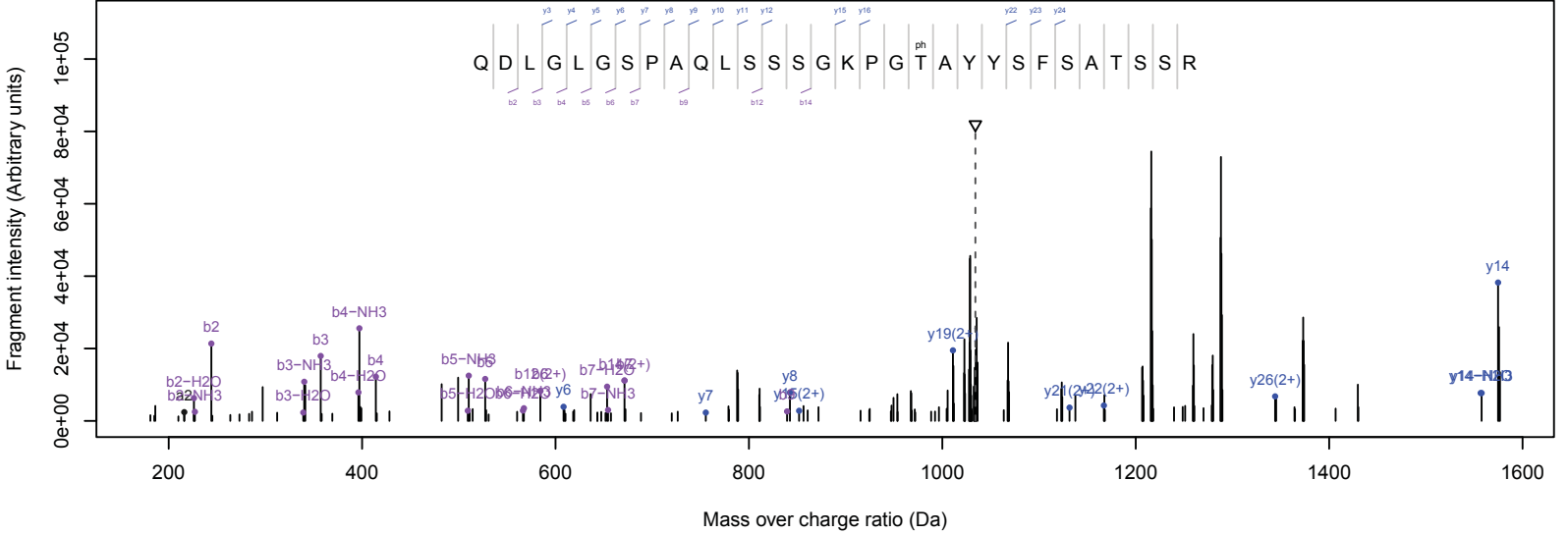
Andromeda score : 139.3



QDLGLGSPAQLSSSGKPGT(ph)AYYSFATSSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33740

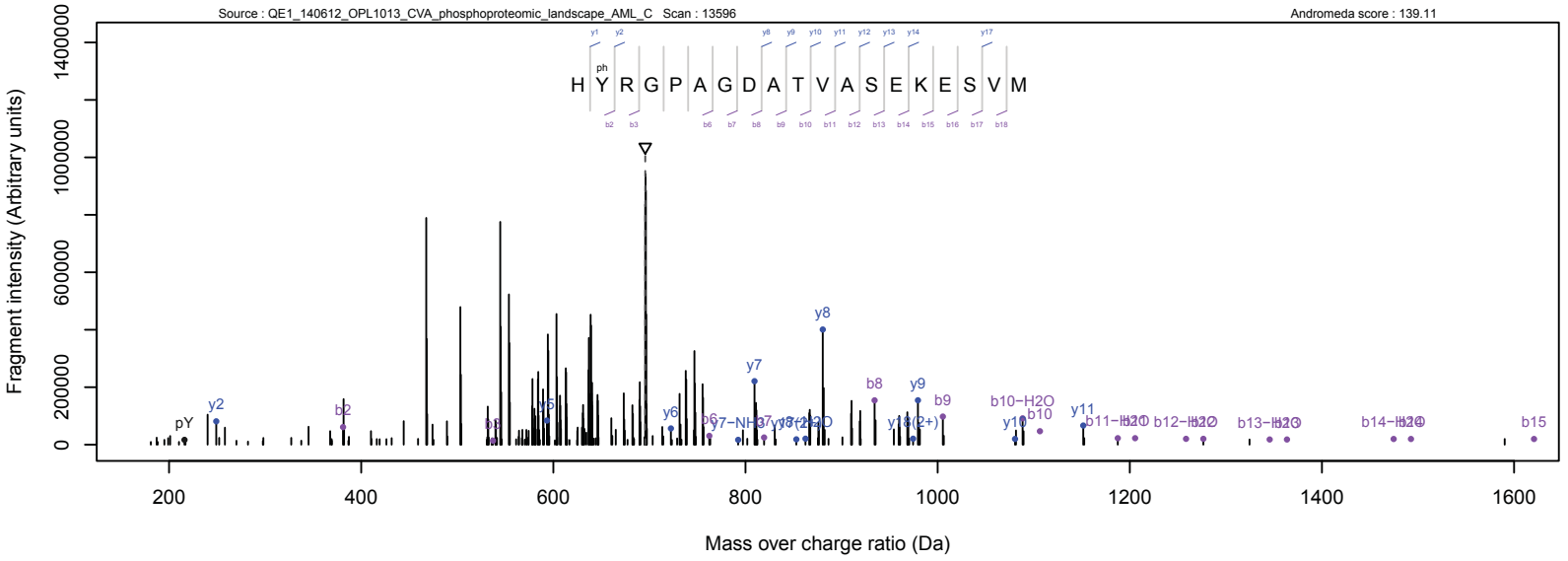
Andromeda score : 139.22



**HY(ph)RGPAGDATVASEKESVM-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13596

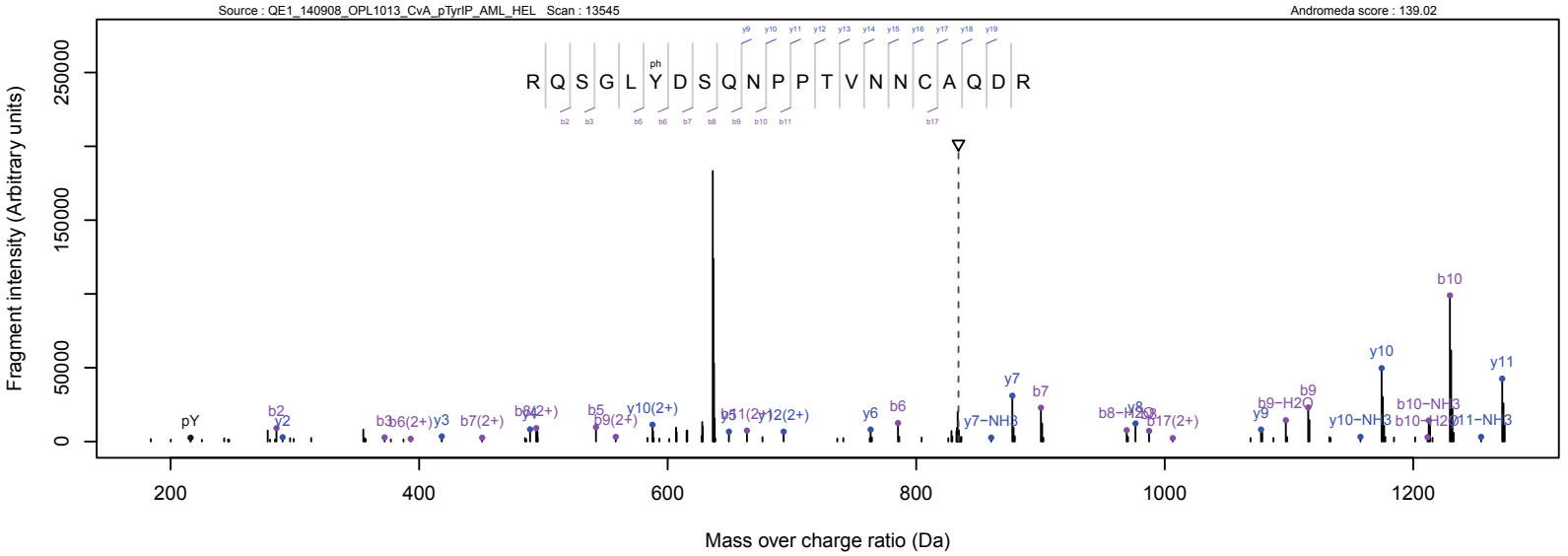
Andromeda score : 139.11



**RQSGLY(ph)DSQNPPVTNNCAQDR-3+**

Source : QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_HEL Scan : 13545

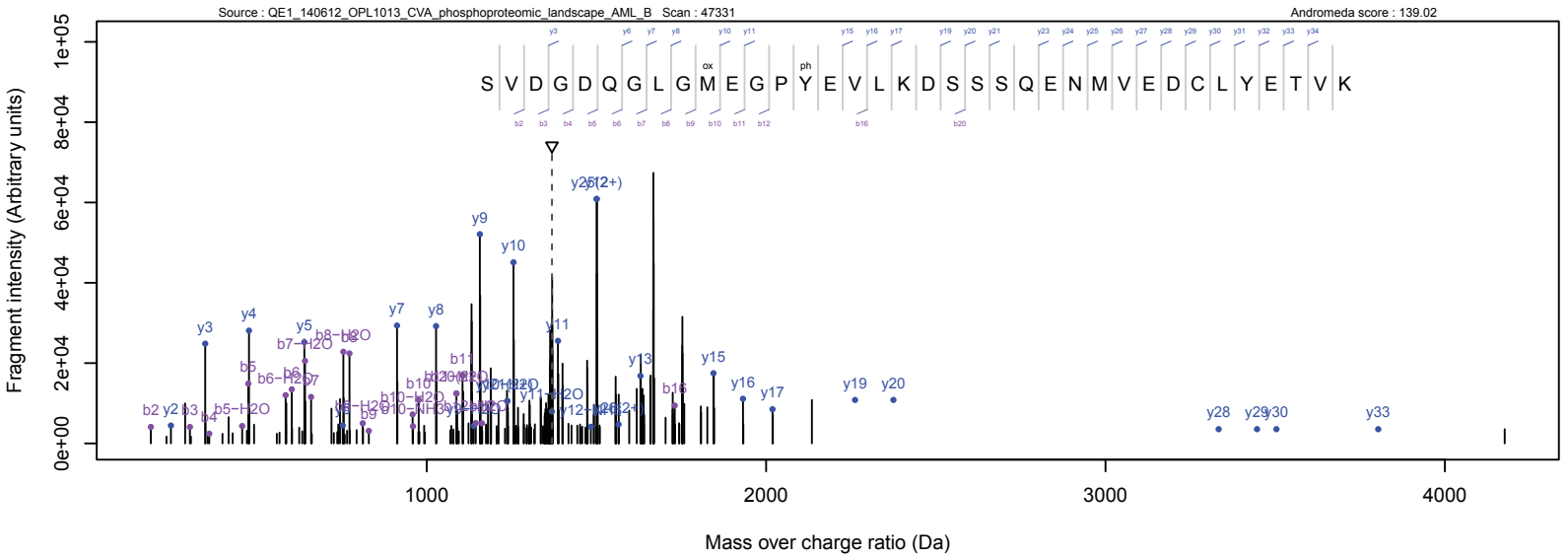
Andromeda score : 139.02



**SVDGQGLGM(ox)EGPY(ph)EVLKDSSENMVEDCLYETVK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47331

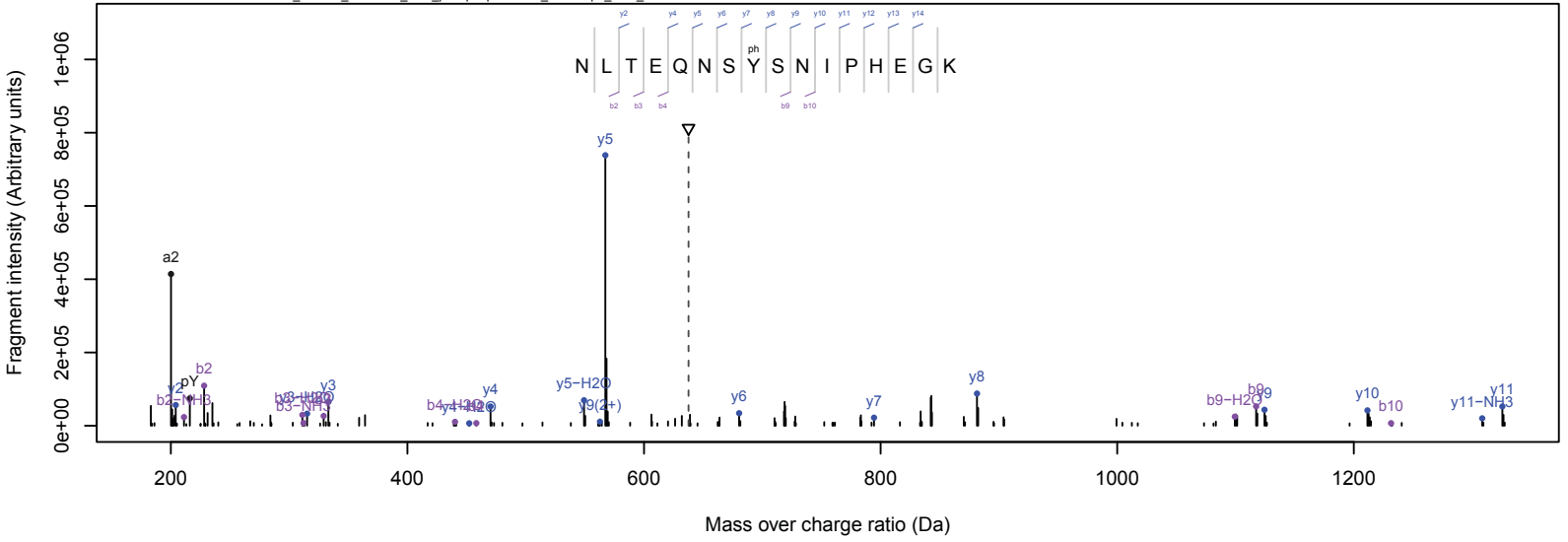
Andromeda score : 139.02



NLTEQNSY(ph)SNIPHEGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10910

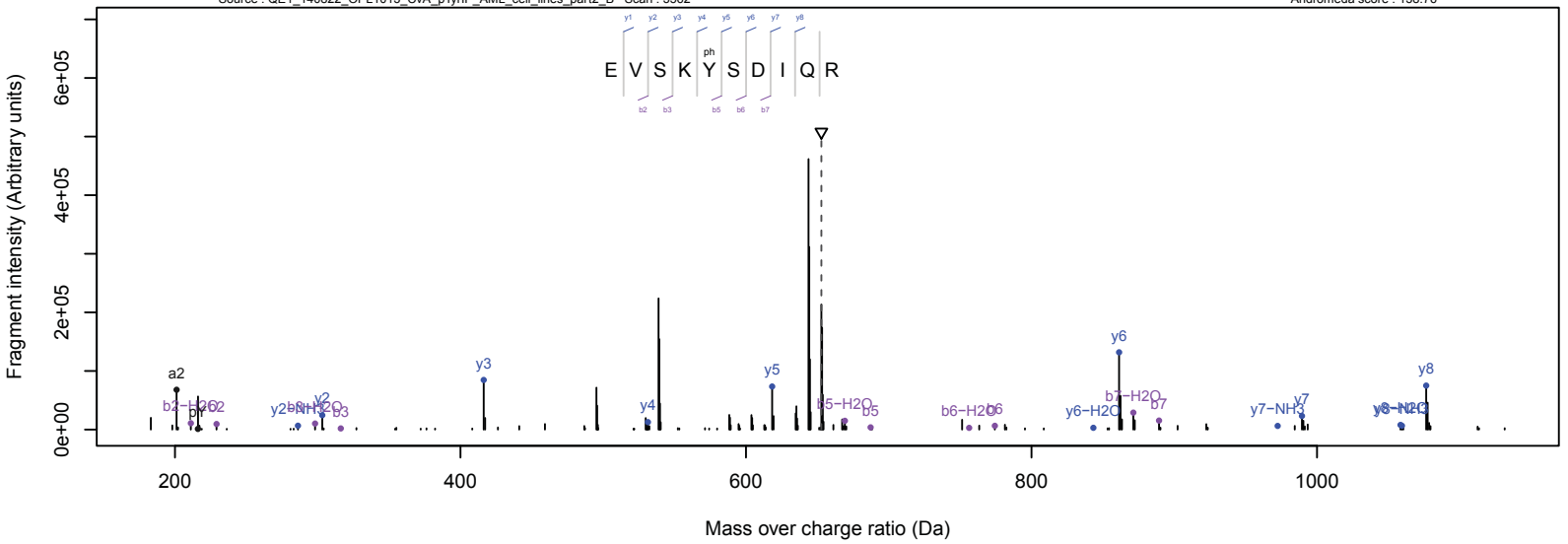
Andromeda score : 138.91



EVSKY(ph)SDIQR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3562

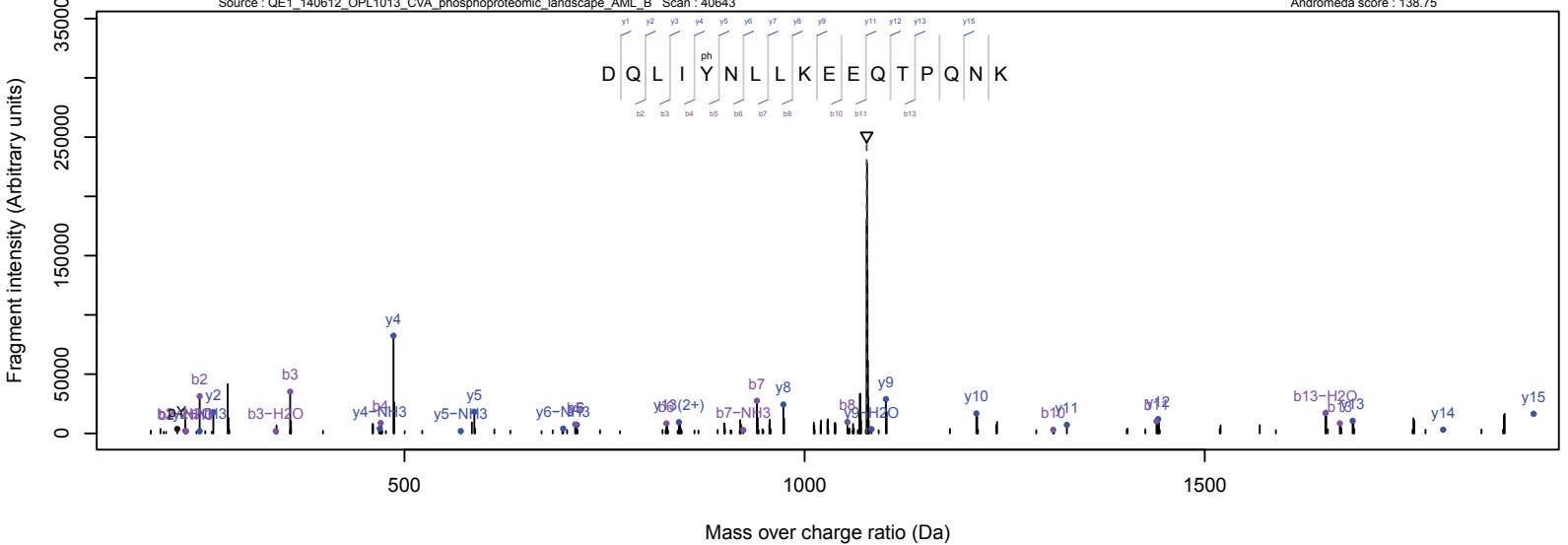
Andromeda score : 138.76

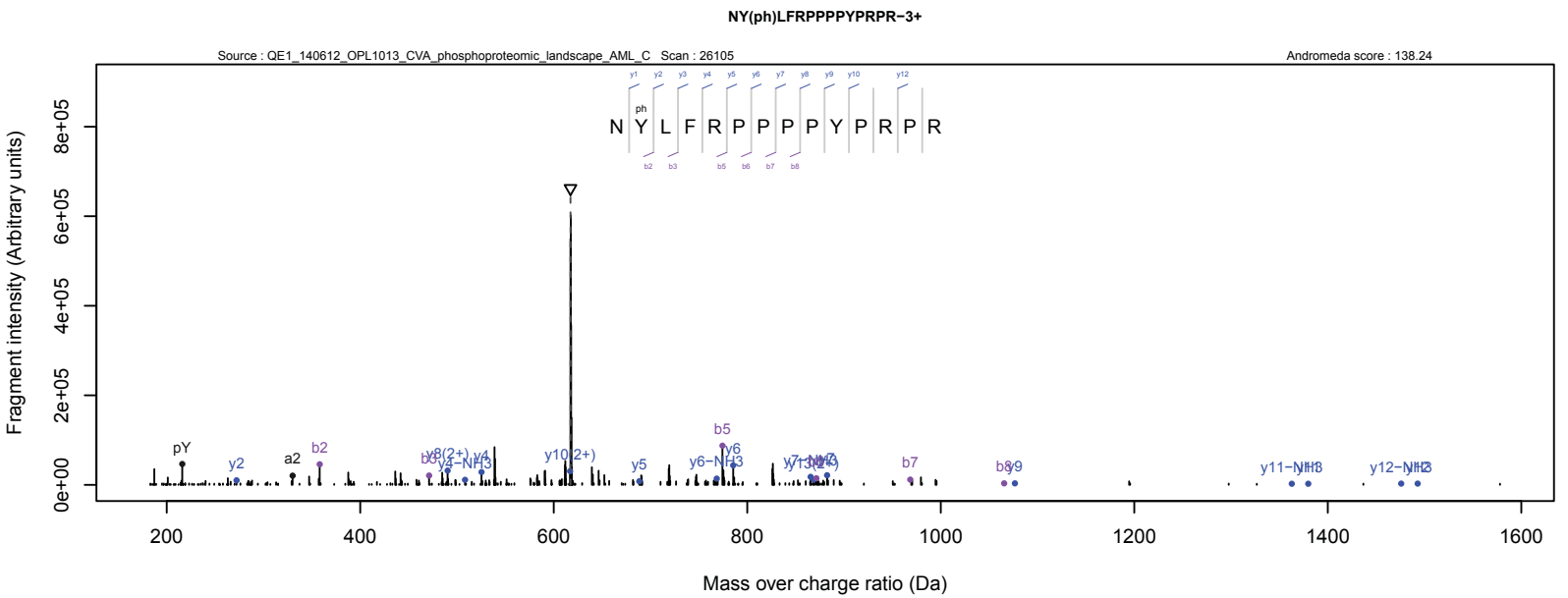
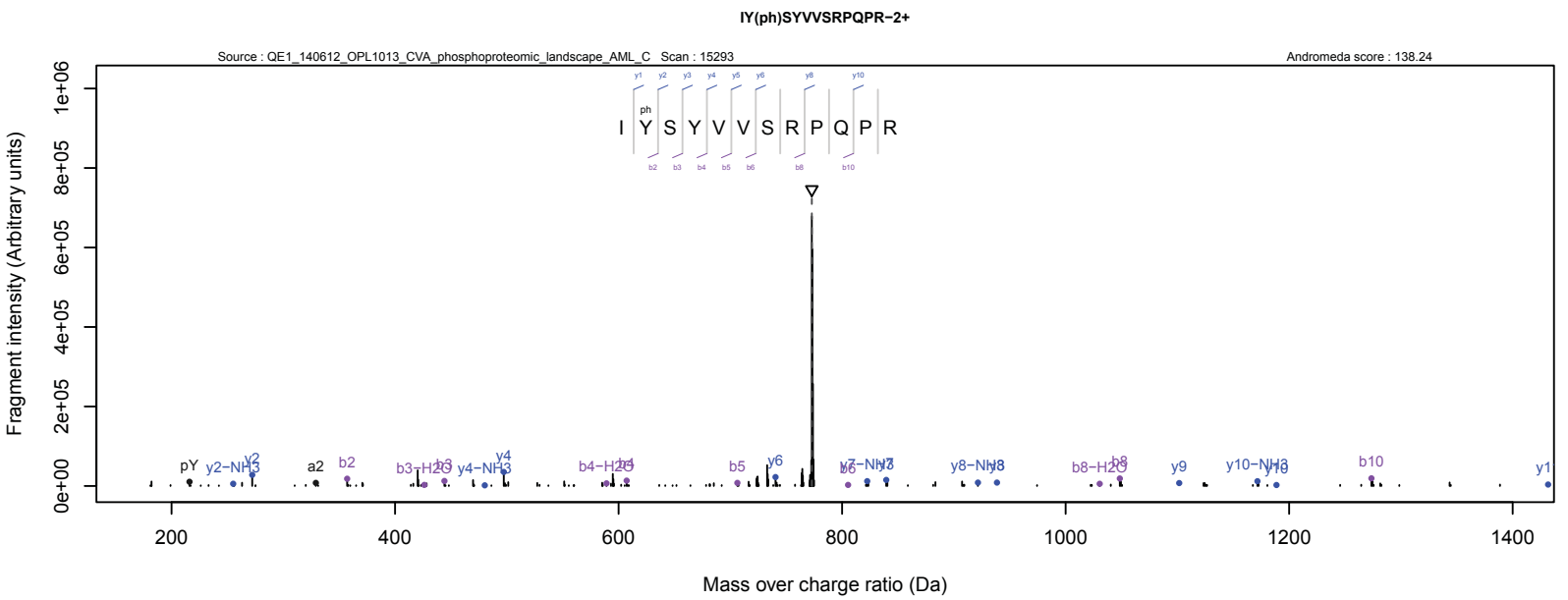
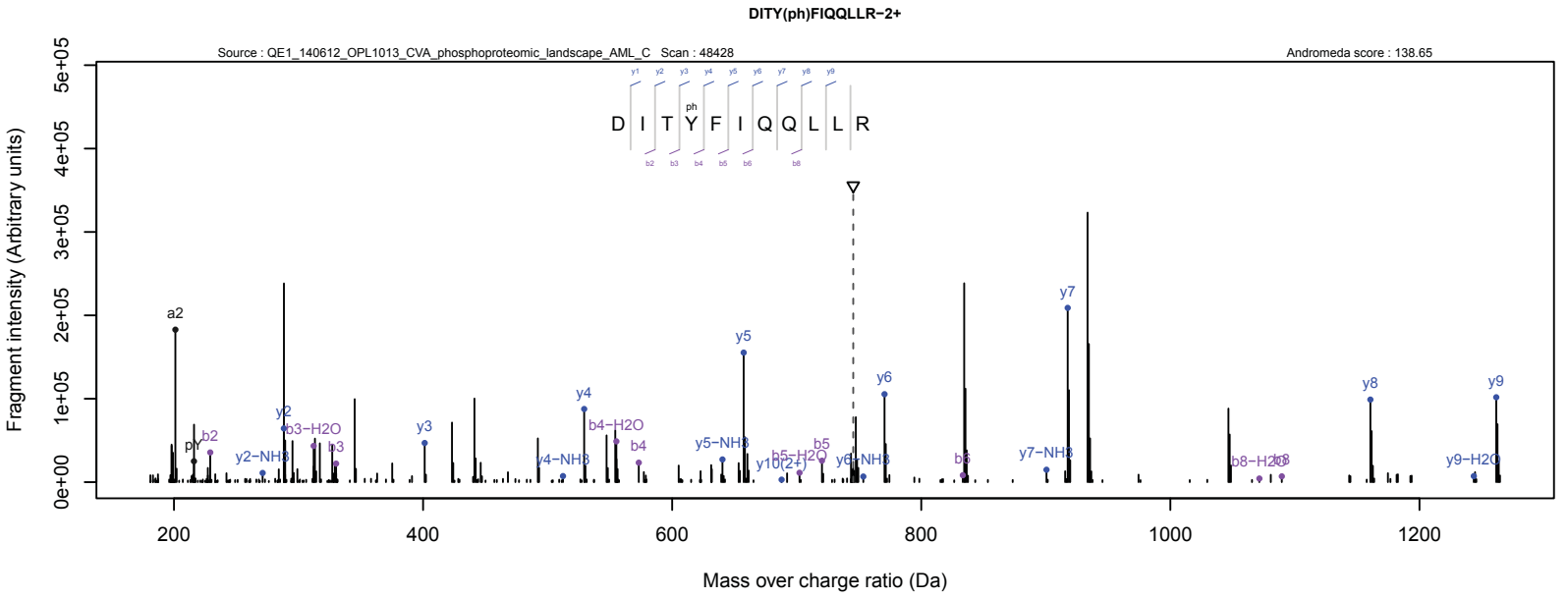


DQLIY(ph)NLLKEEQTPQNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40643

Andromeda score : 138.75

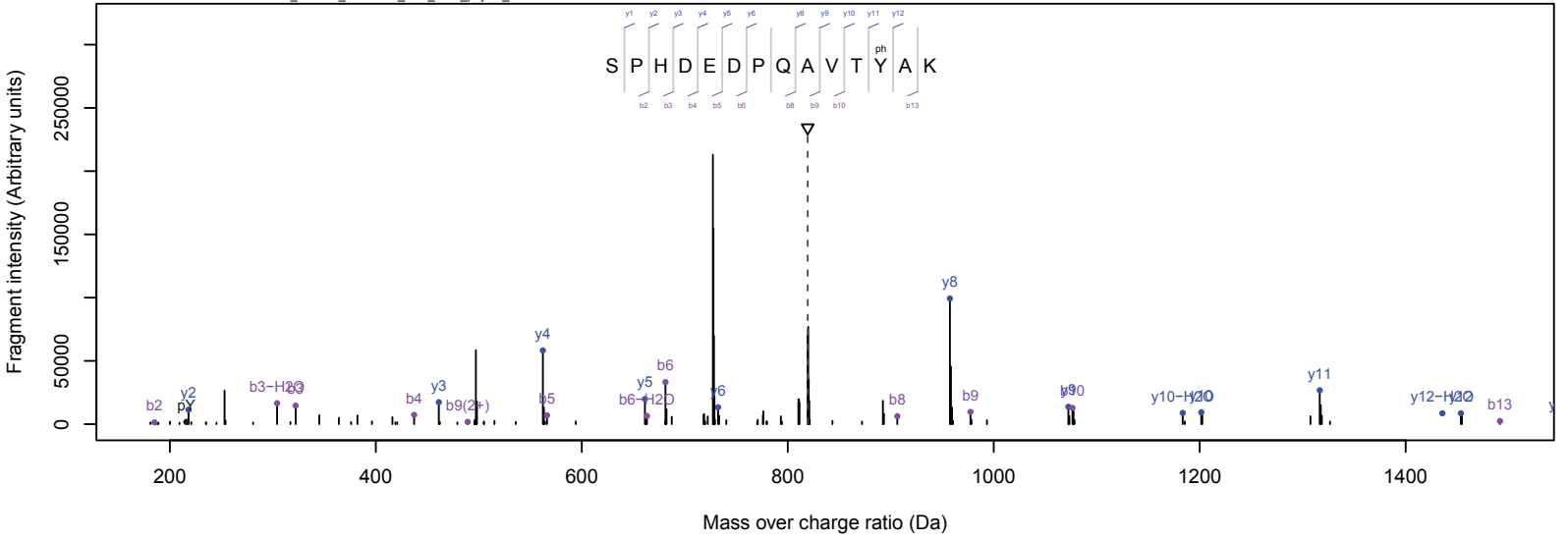




SPHDEDPAVTY(ph)AK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 8314

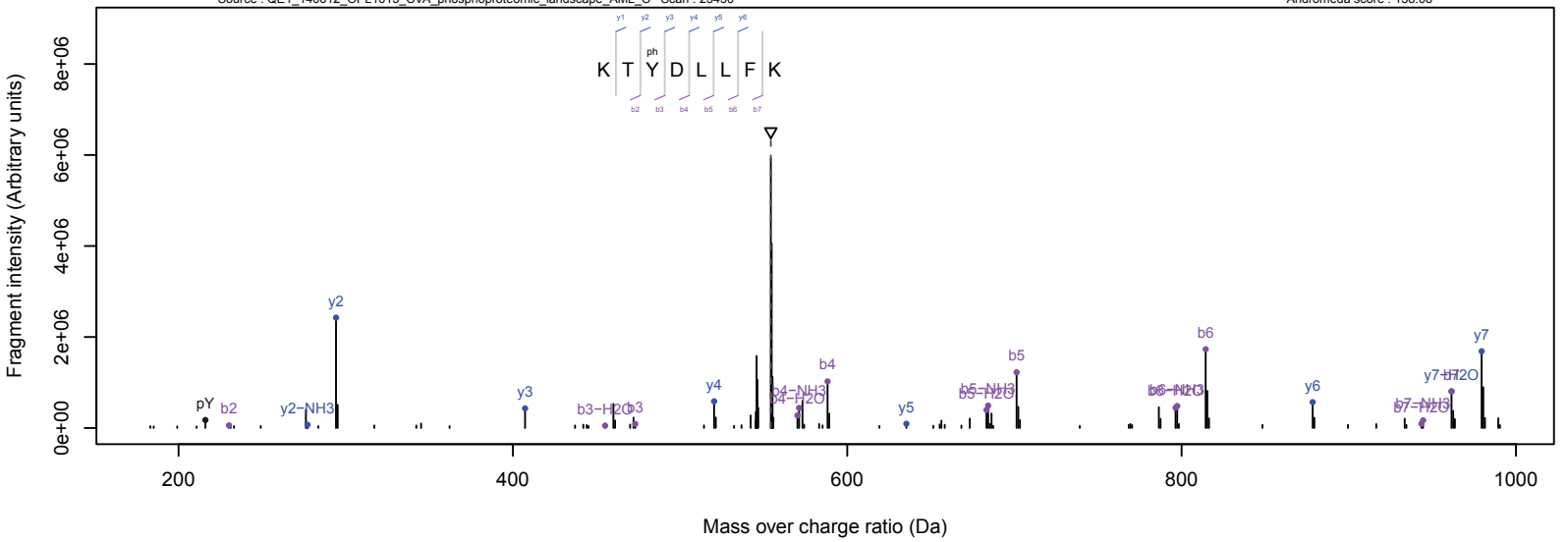
Andromeda score : 138.19



KTY(ph)DLLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23430

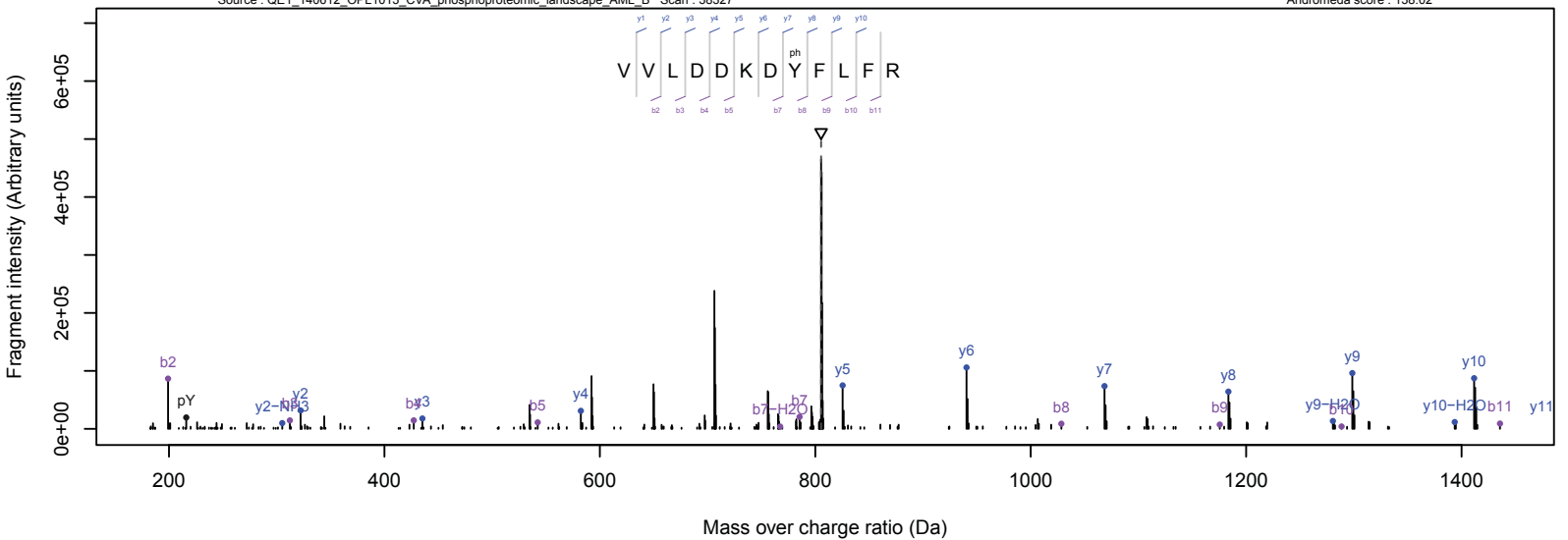
Andromeda score : 138.08



VLLDDKY(ph)FLFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38327

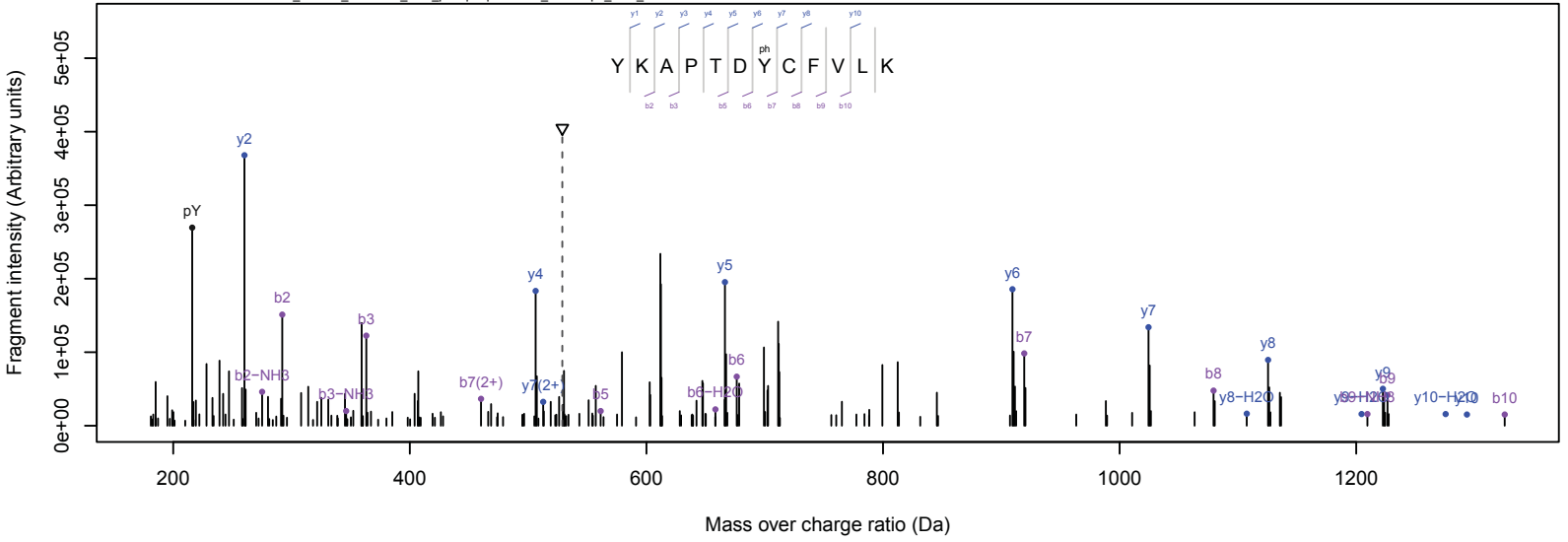
Andromeda score : 138.02



YKAPTDY(ph)CFVLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27980

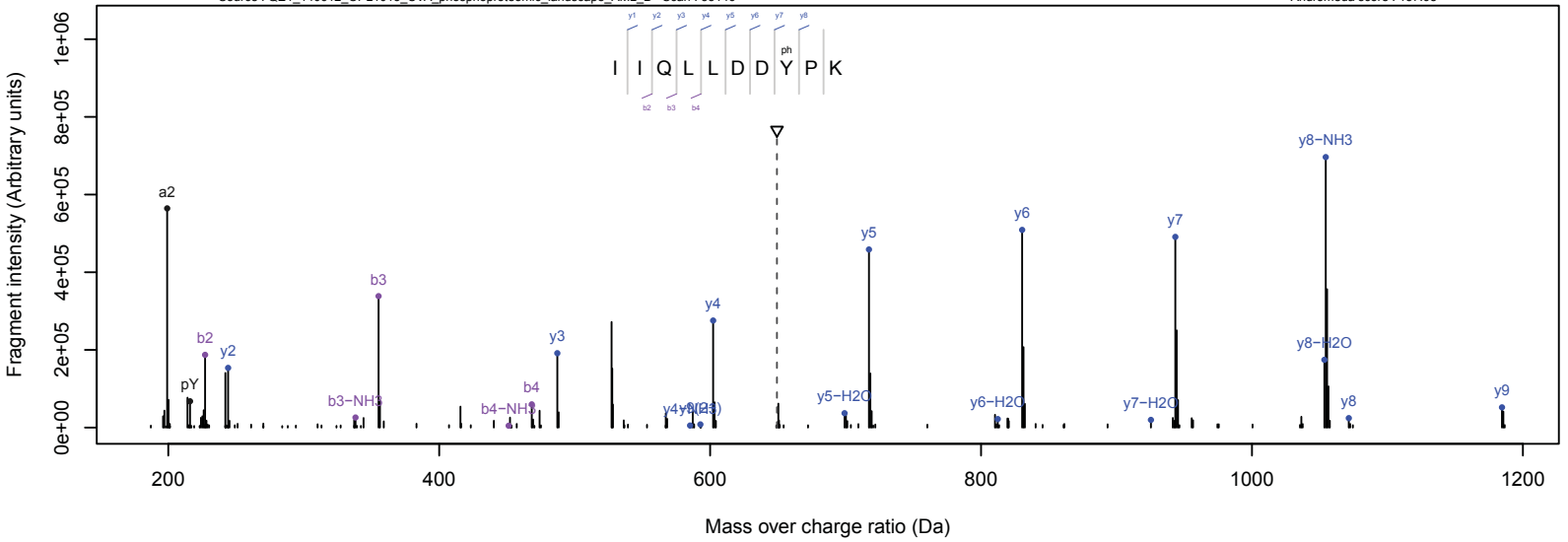
Andromeda score : 138.02



IIQLDDY(ph)PK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33146

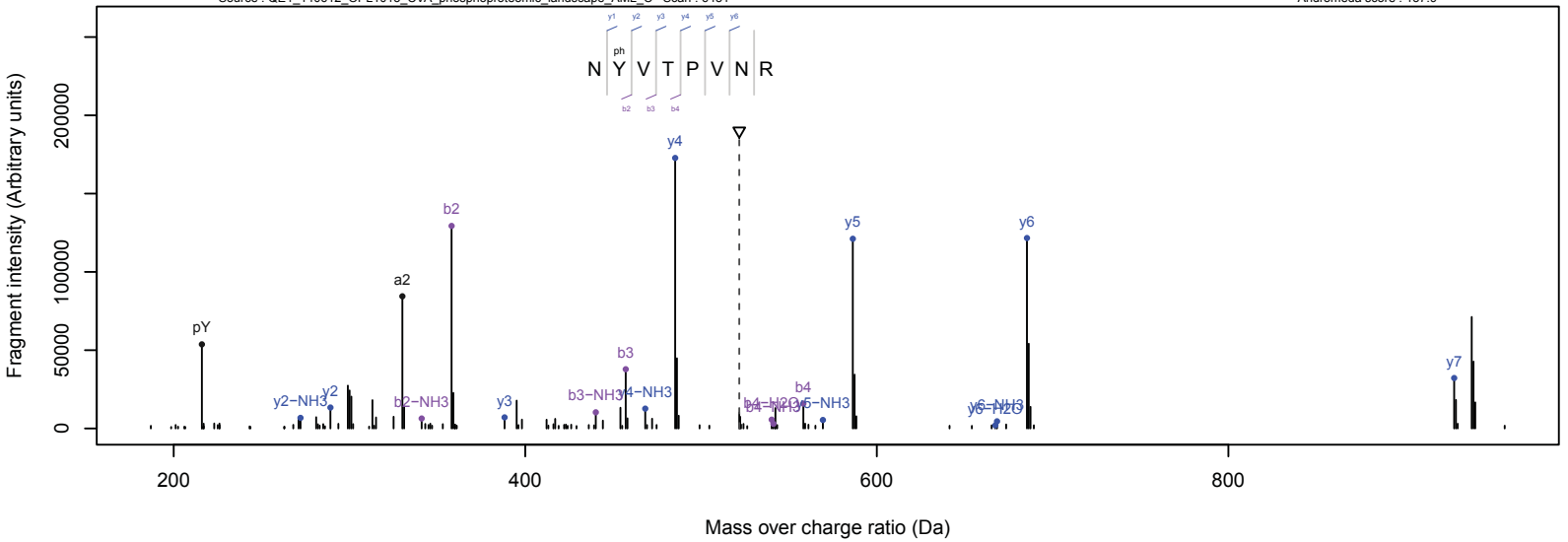
Andromeda score : 137.98



NY(ph)VTPVNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6451

Andromeda score : 137.9



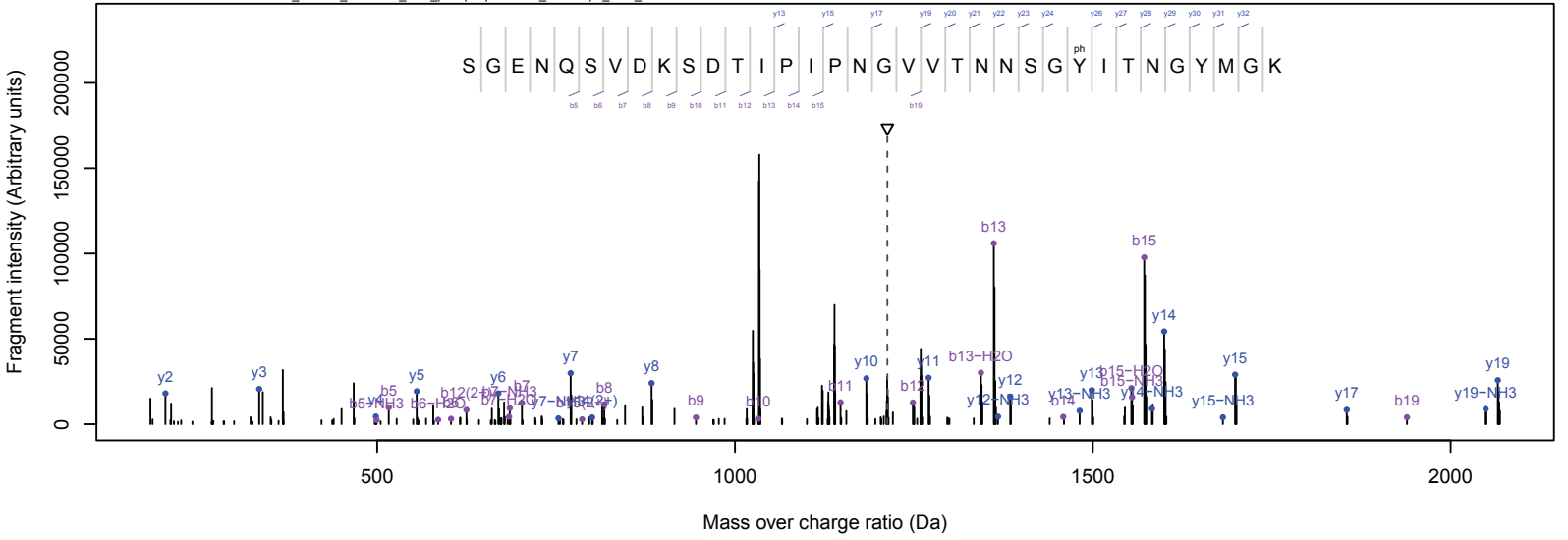




SGENQSVDKSDTIPIPIPNNGVVTTNNSGY(ph)ITNGYMGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 36704

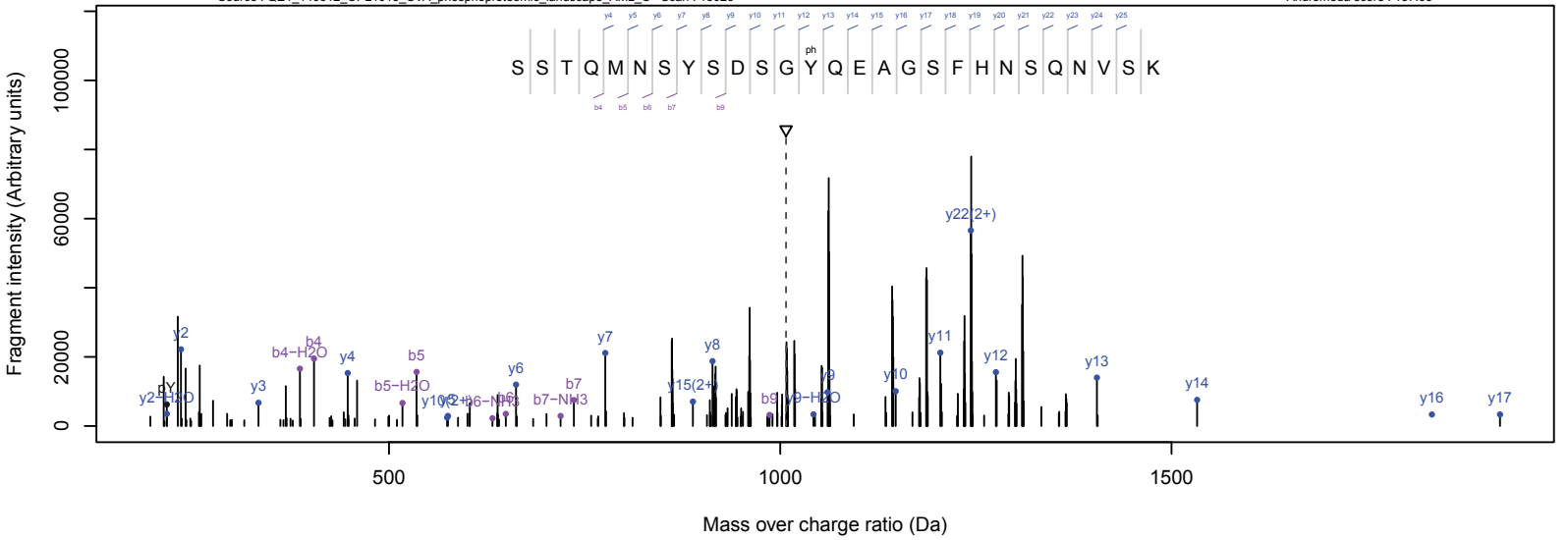
Andromeda score : 137.25



SSTQMNSYSDSGY(ph)QEAGSFHNSQNVSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15026

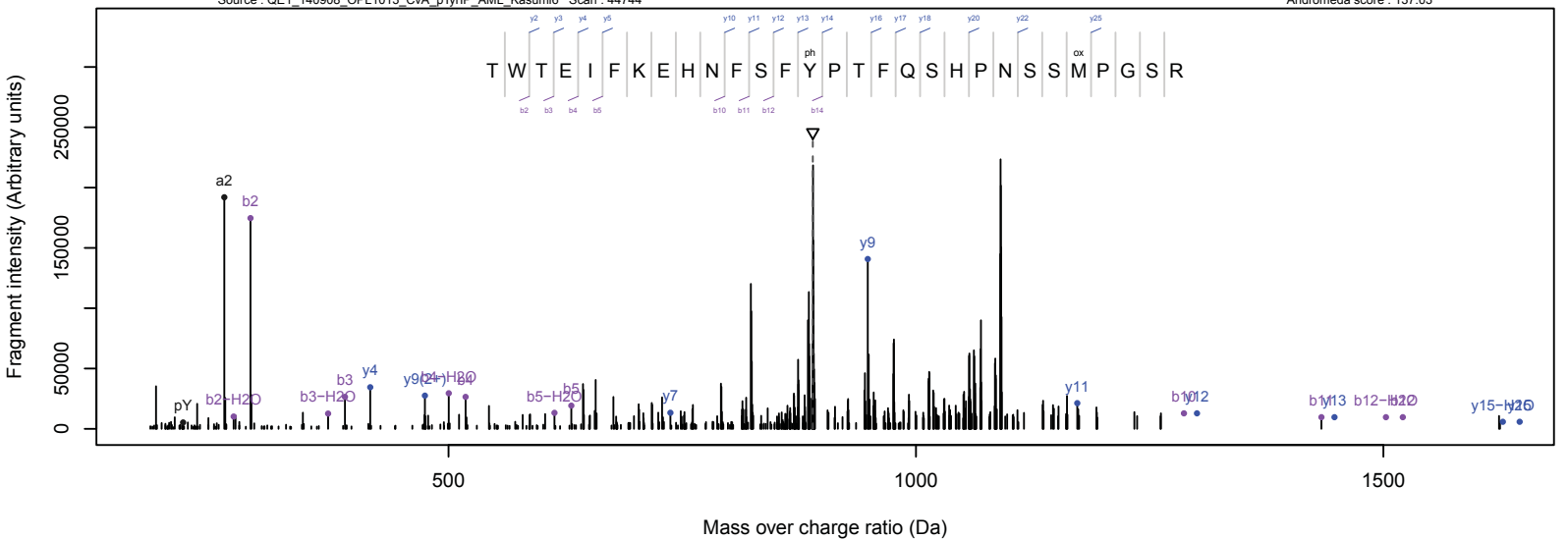
Andromeda score : 137.03



TWTEIFKEHNFSFY(ph)PTFQSHPNSSM(ox)PGSR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 44744

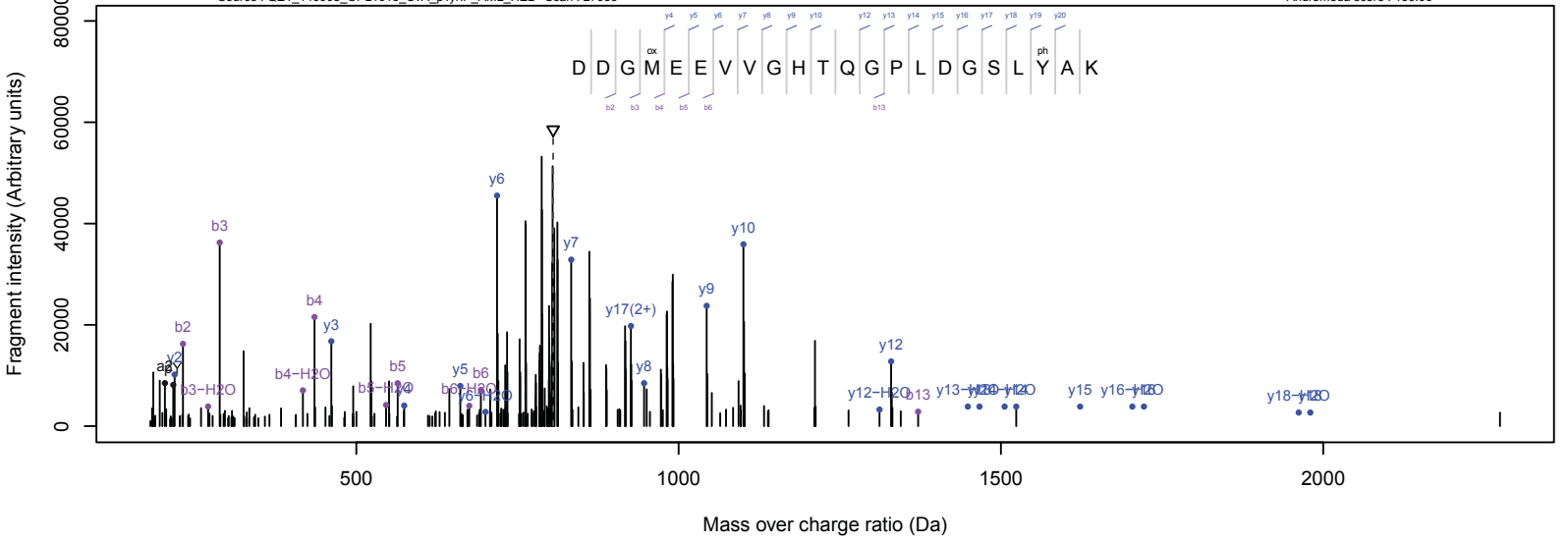
Andromeda score : 137.03



DDGM(ox)EEVVGHTQGPLDGSLY(ph)AK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 27953

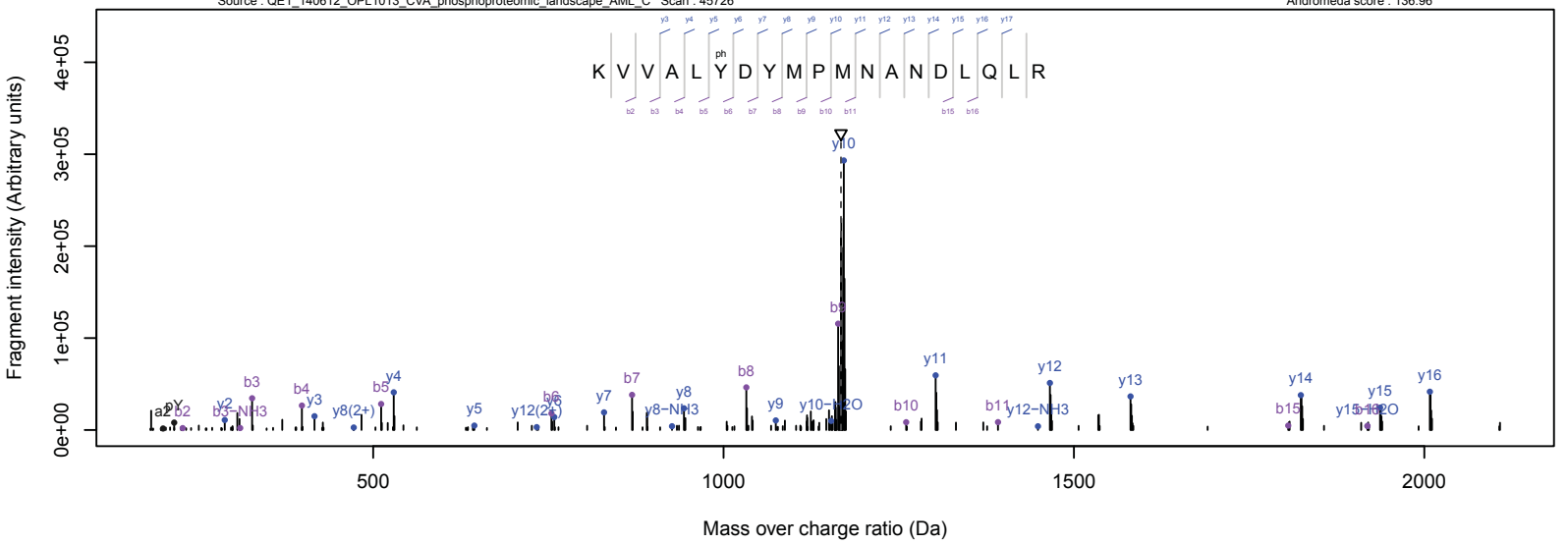
Andromeda score : 136.96



KVVALY(ph)DYMPMNANDLQLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45726

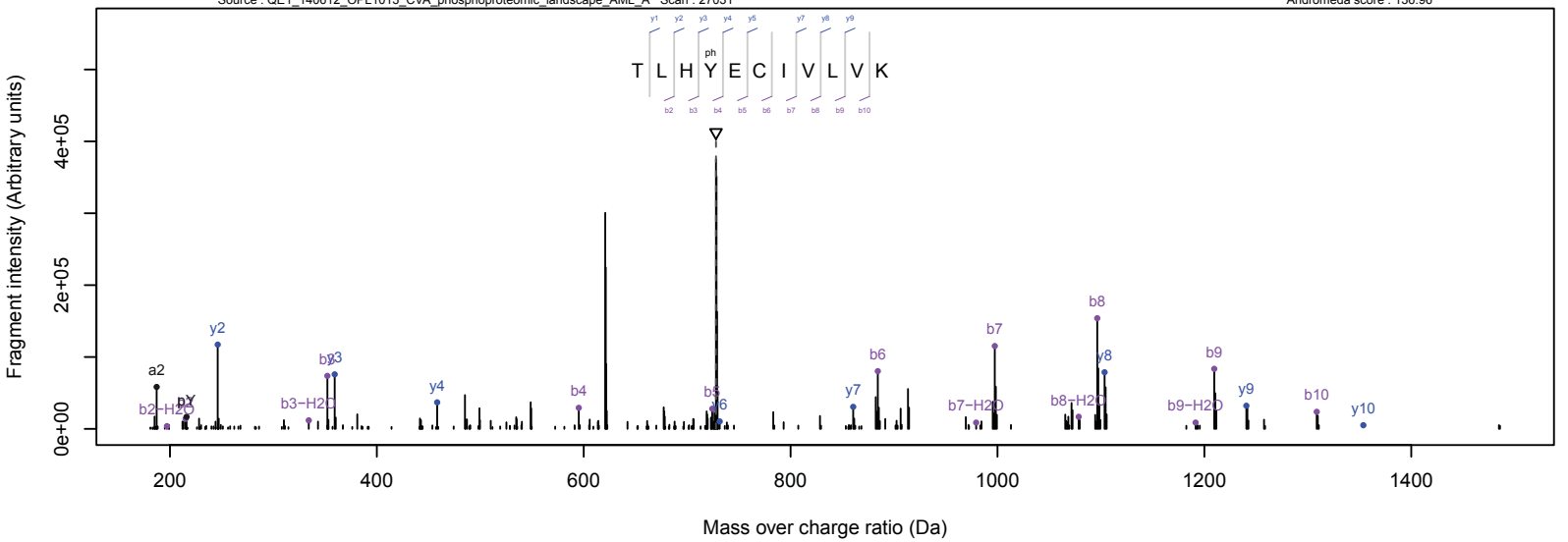
Andromeda score : 136.96



TLHY(ph)ECIVLVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27031

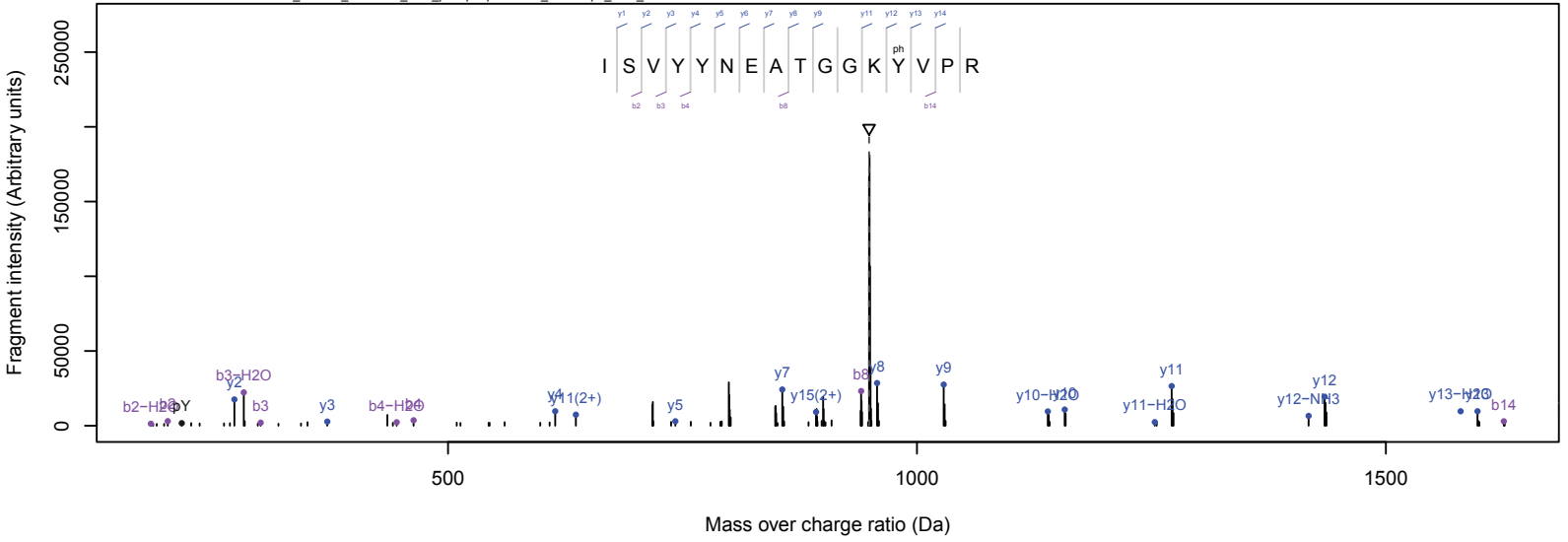
Andromeda score : 136.96



ISVYYNEATGGKY(ph)VPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21715

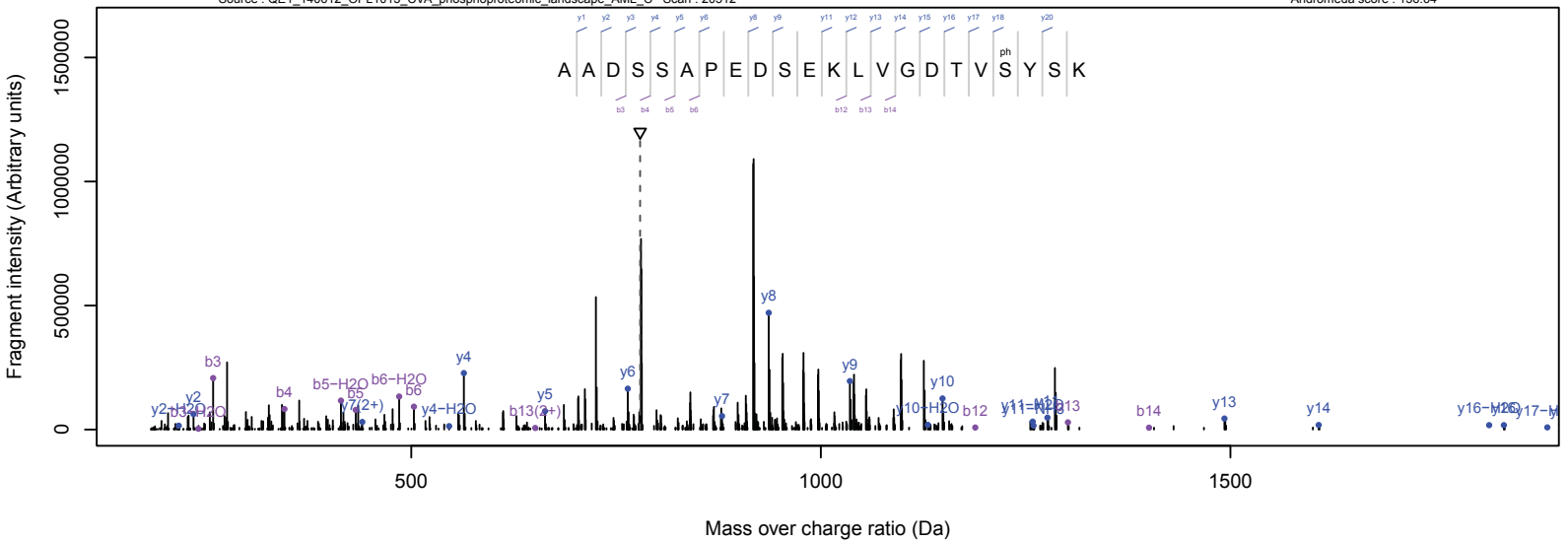
Andromeda score : 136.81



AADSSAPEDSEKLVGDTV(ph)YSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20512

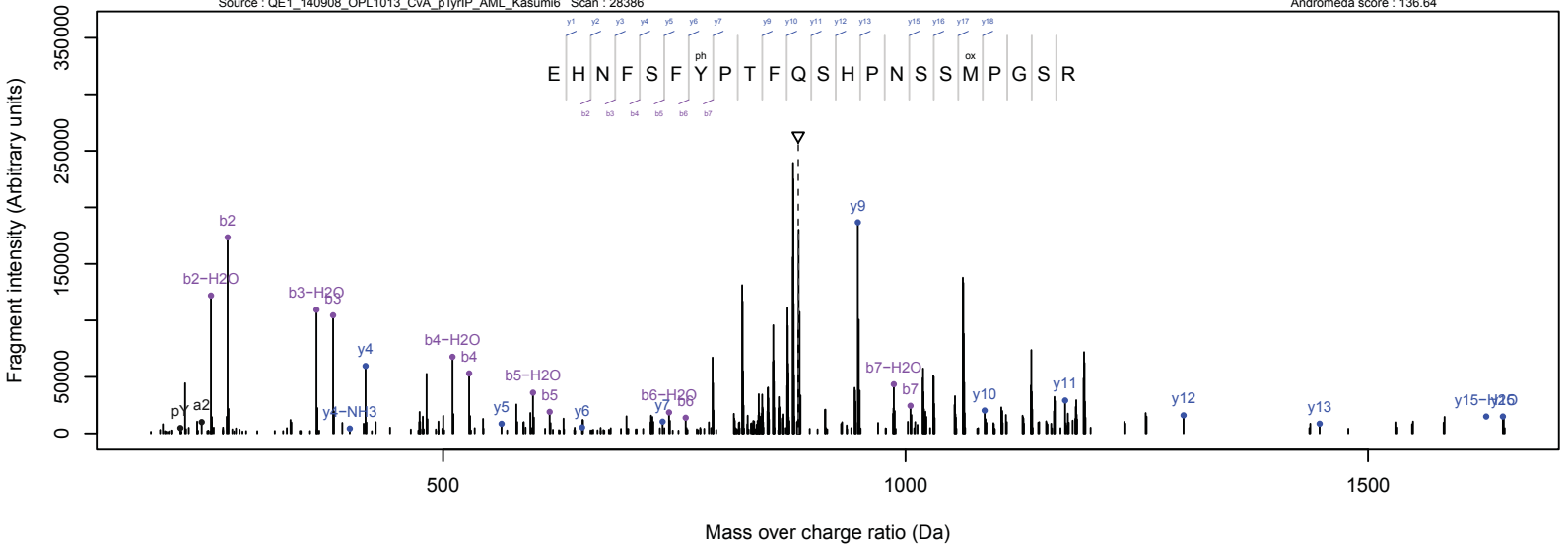
Andromeda score : 136.64



EHNFSFY(ph)PTFQSHPNSSMPGSR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 28386

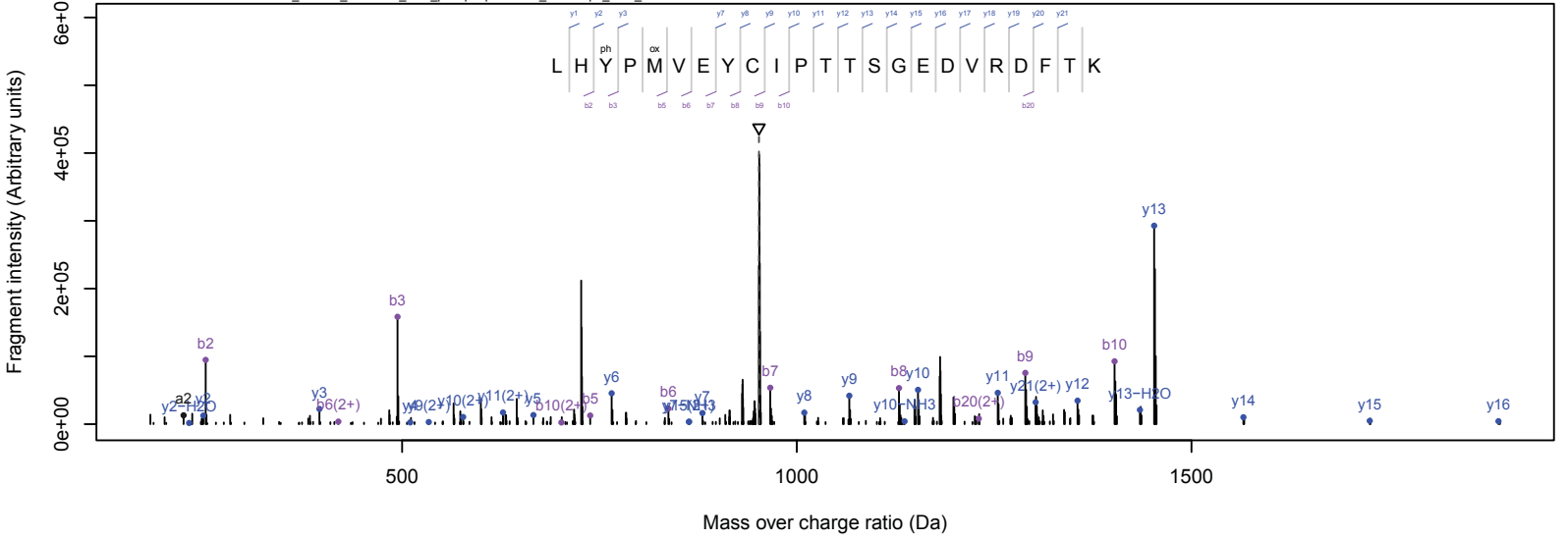
Andromeda score : 136.64



LHY(ph)PM(ox)VEYCIPTTSGEDVRDFTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33790

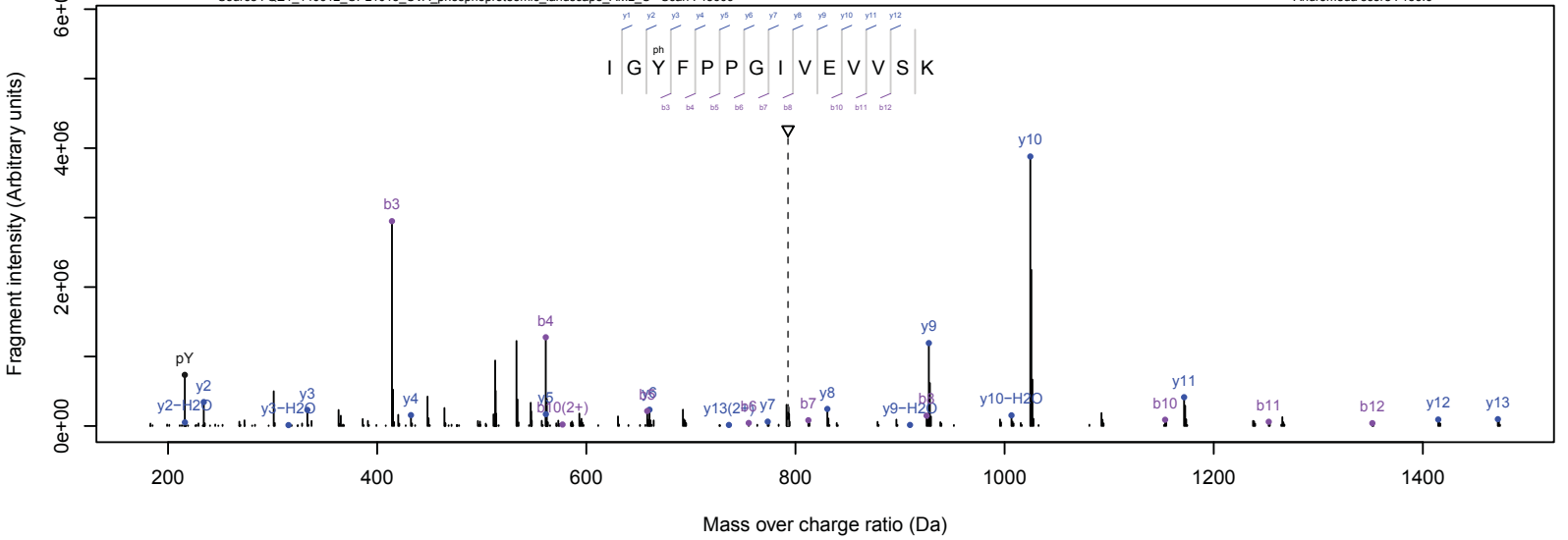
Andromeda score : 136.62



IGY(ph)FPPGIVEVSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45609

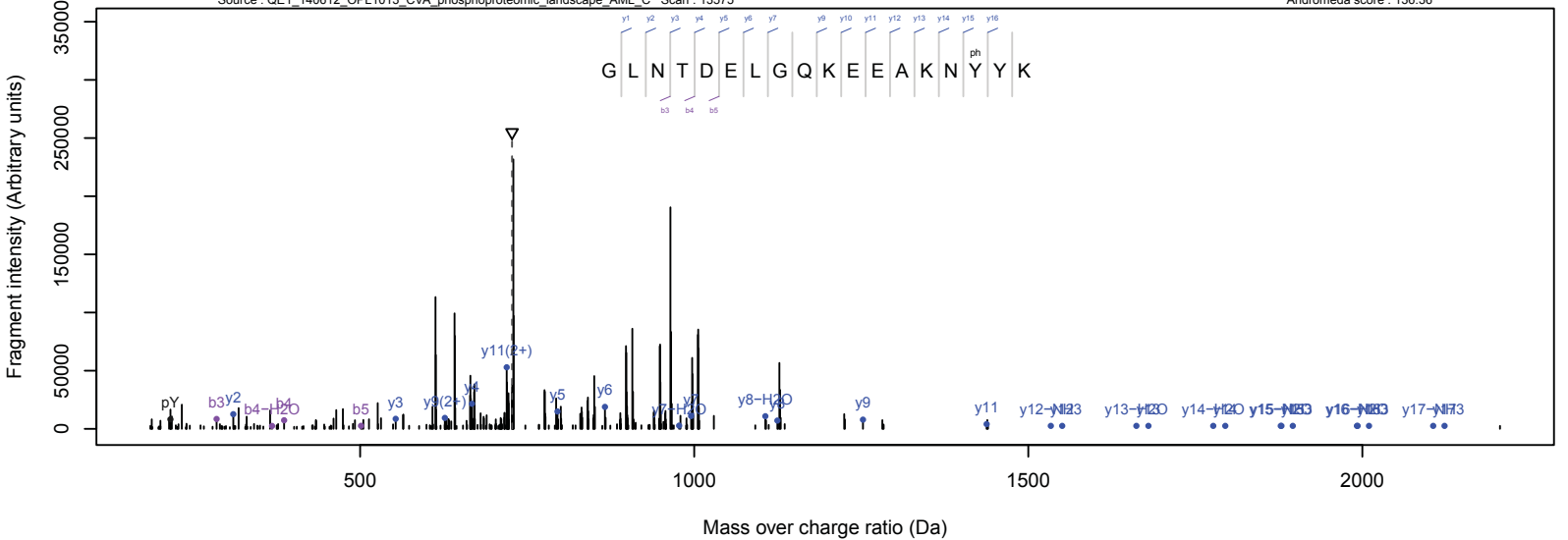
Andromeda score : 136.6



GLNTDELGQKEEAKNY(ph)YK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13575

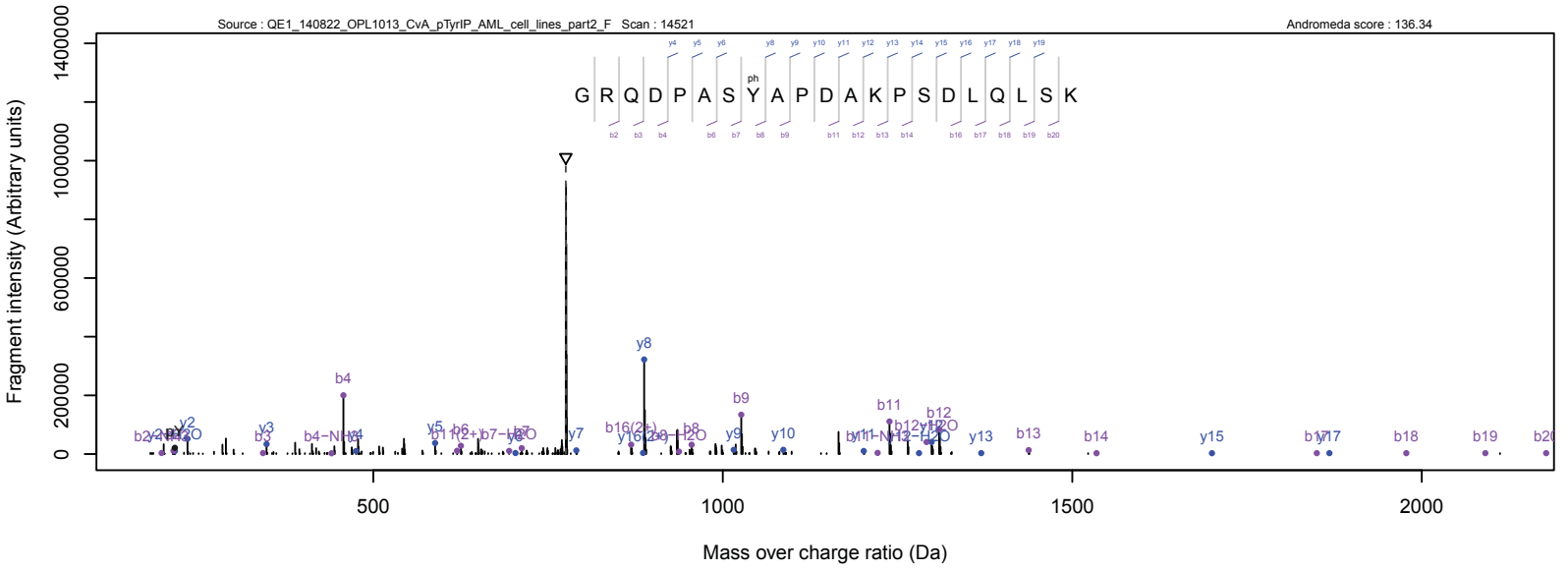
Andromeda score : 136.36



GRQDPASY(ph)APDAKPSDLQLSK-3+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan: 14521

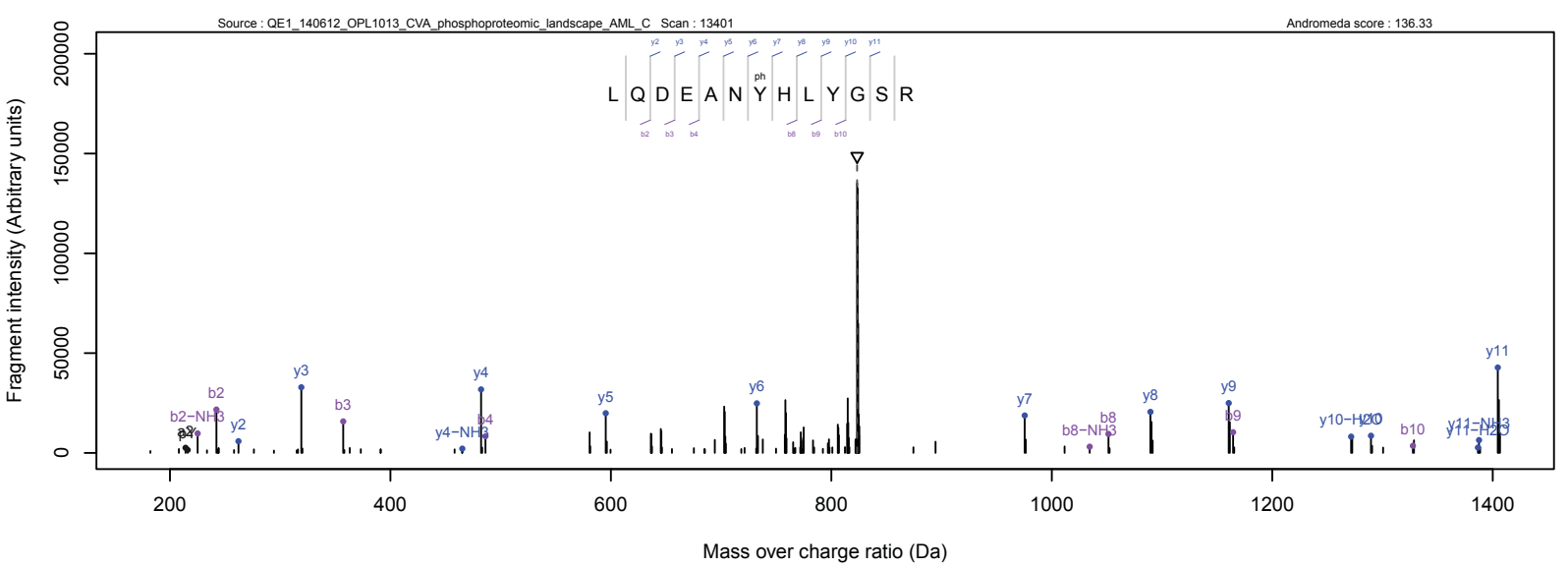
Andromeda score: 136.34



LQDEANY(ph)HLYGSR-2+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 13401

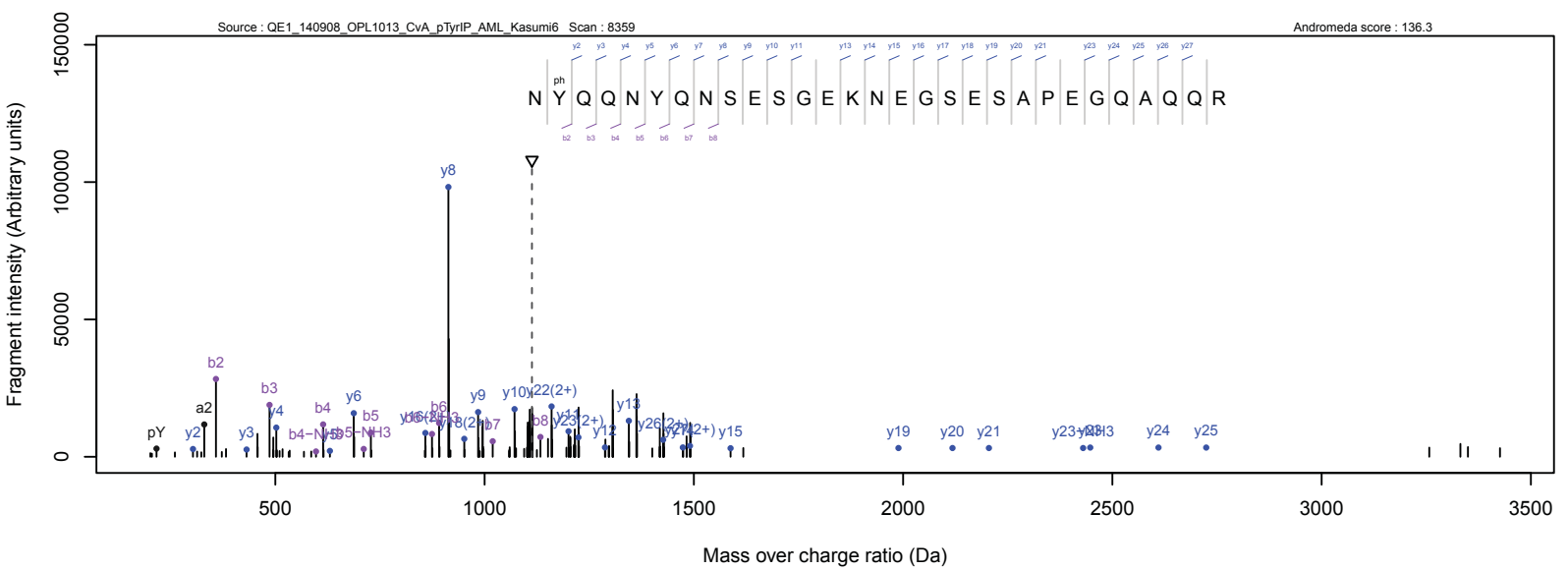
Andromeda score: 136.33



NY(ph)QQNYQNSESGEKNEGSESAPEGQAQR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 8359

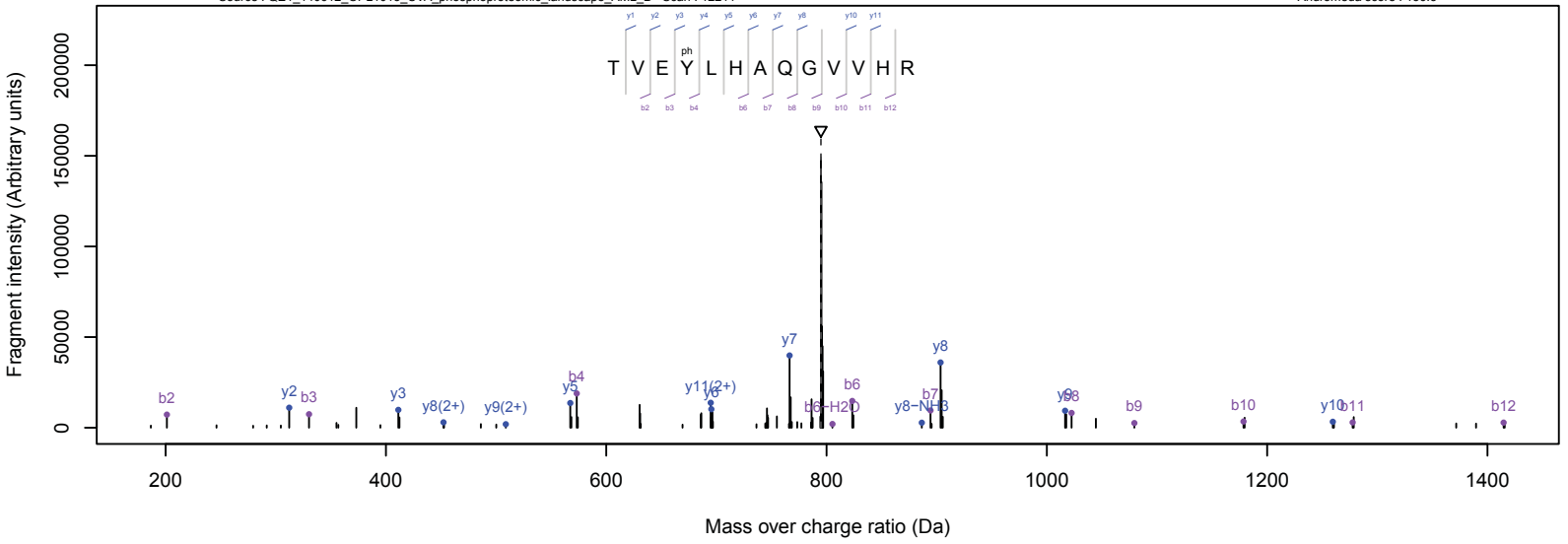
Andromeda score: 136.3



**TVEY(ph)LHAQGVVHR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12211

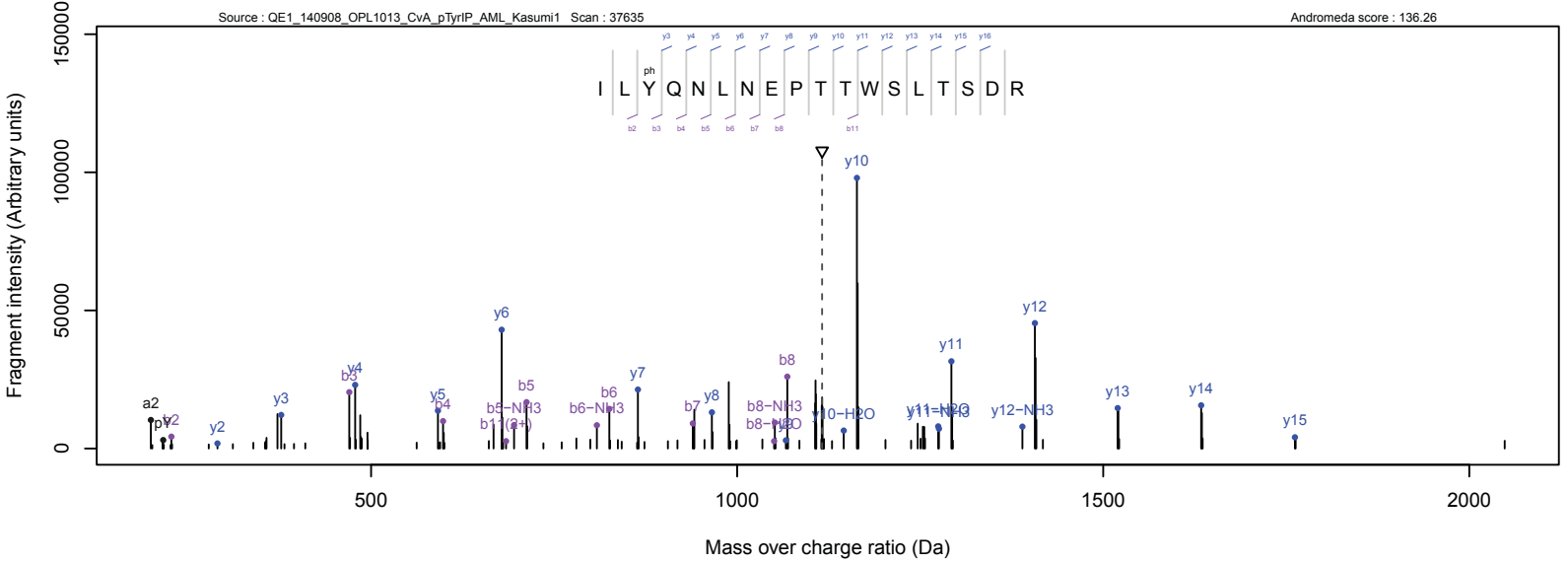
Andromeda score : 136.3



**ILY(ph)QNLNEPTTWSLTSR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 37635

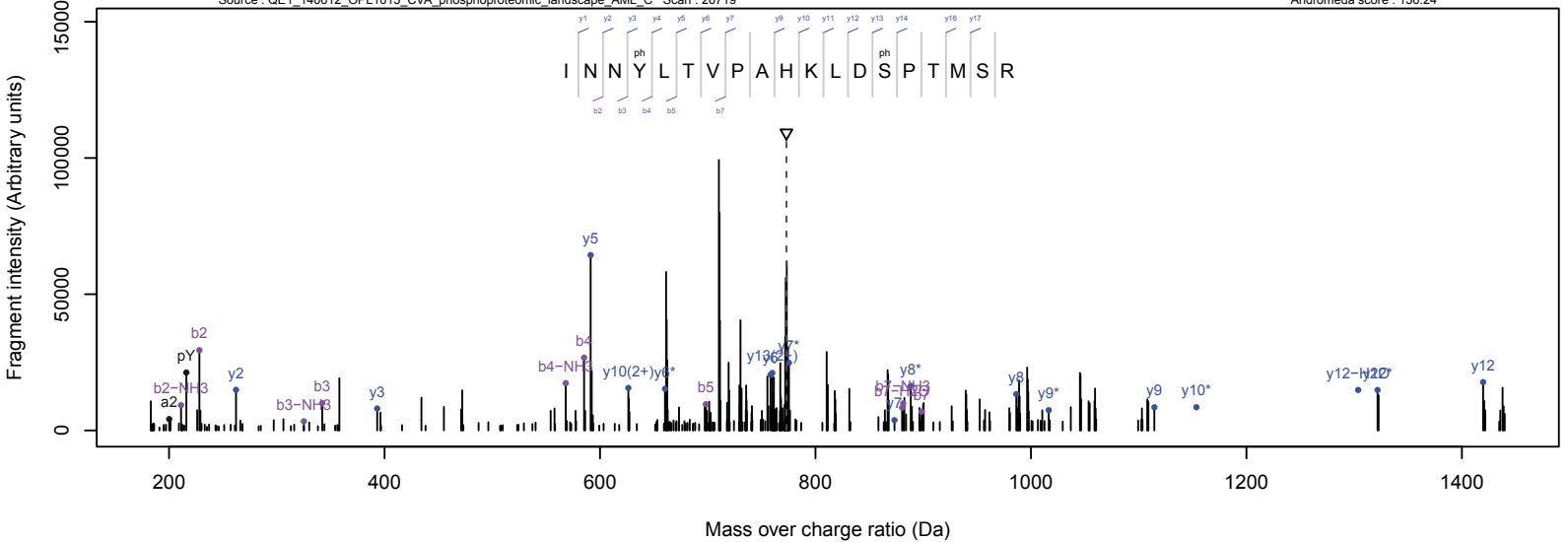
Andromeda score : 136.26



**INNY(ph)LTPAHKLD(ph)PTMSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26719

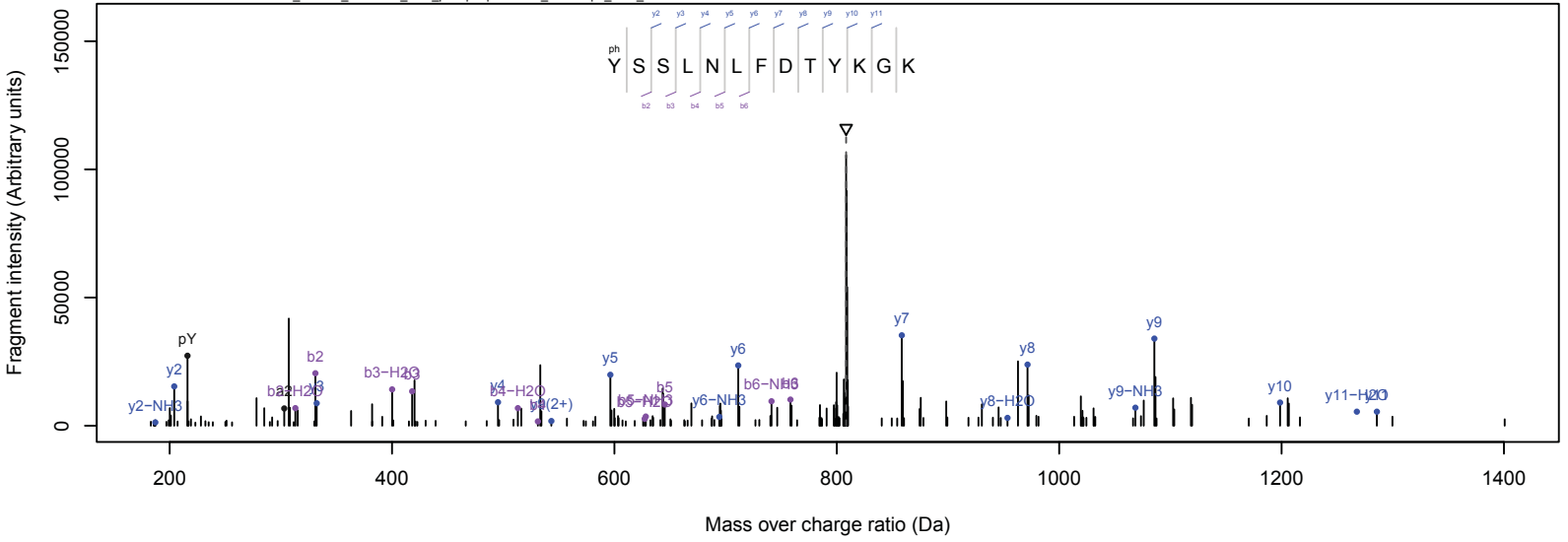
Andromeda score : 136.24



Y(ph)SSLNLFDTYK GK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30561

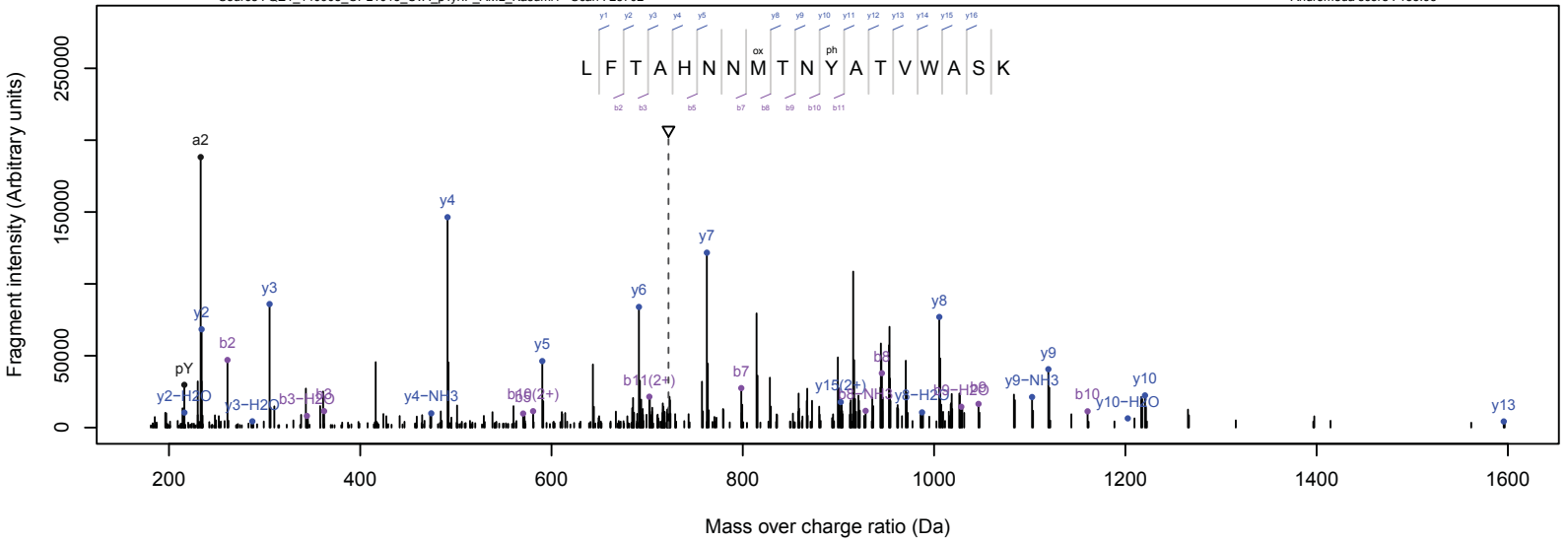
Andromeda score : 136.24



LFTAHHNNM(ox)TNY(ph)ATVWASK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 25702

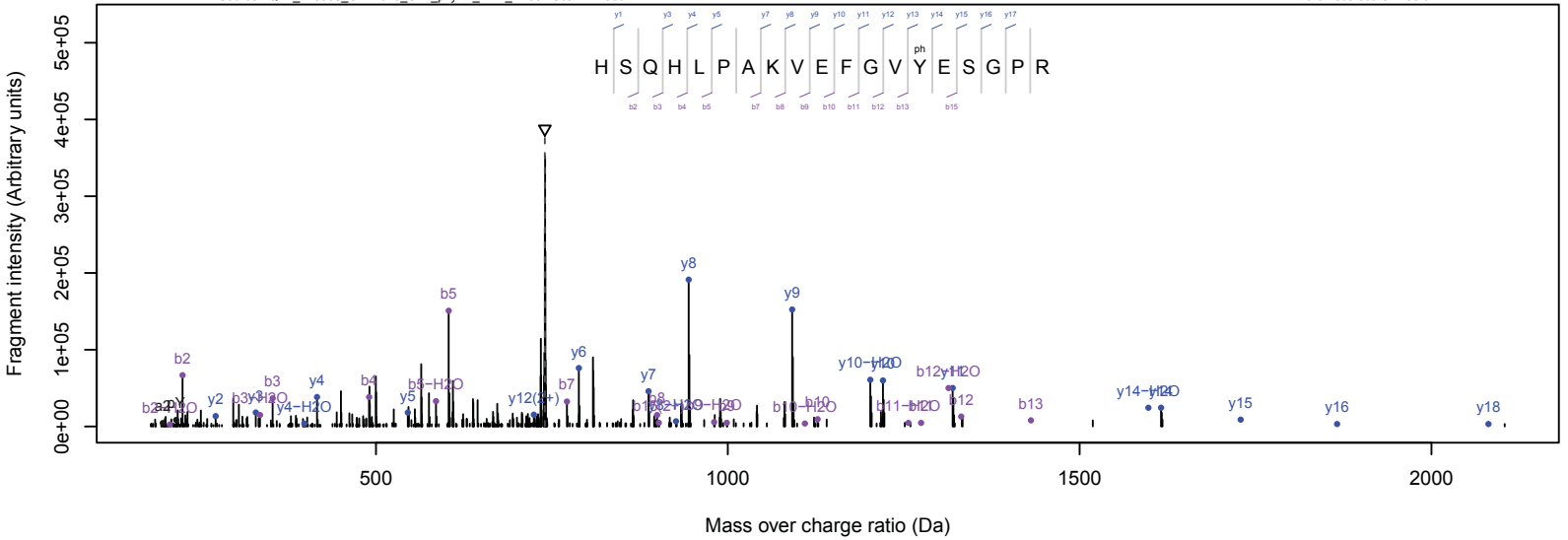
Andromeda score : 135.95



HSQHLPKVEFGVY(ph)ESGPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 21506

Andromeda score : 135.91

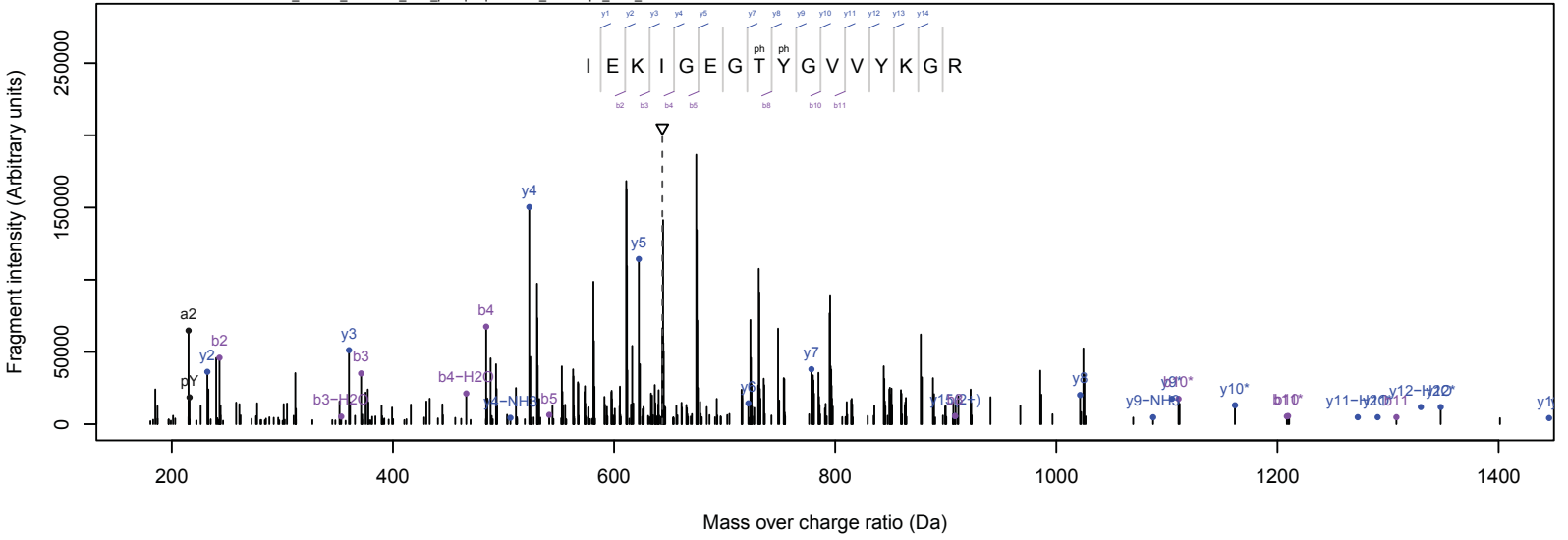




IEKIGEGT(ph)Y(ph)GVVYKGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 18499

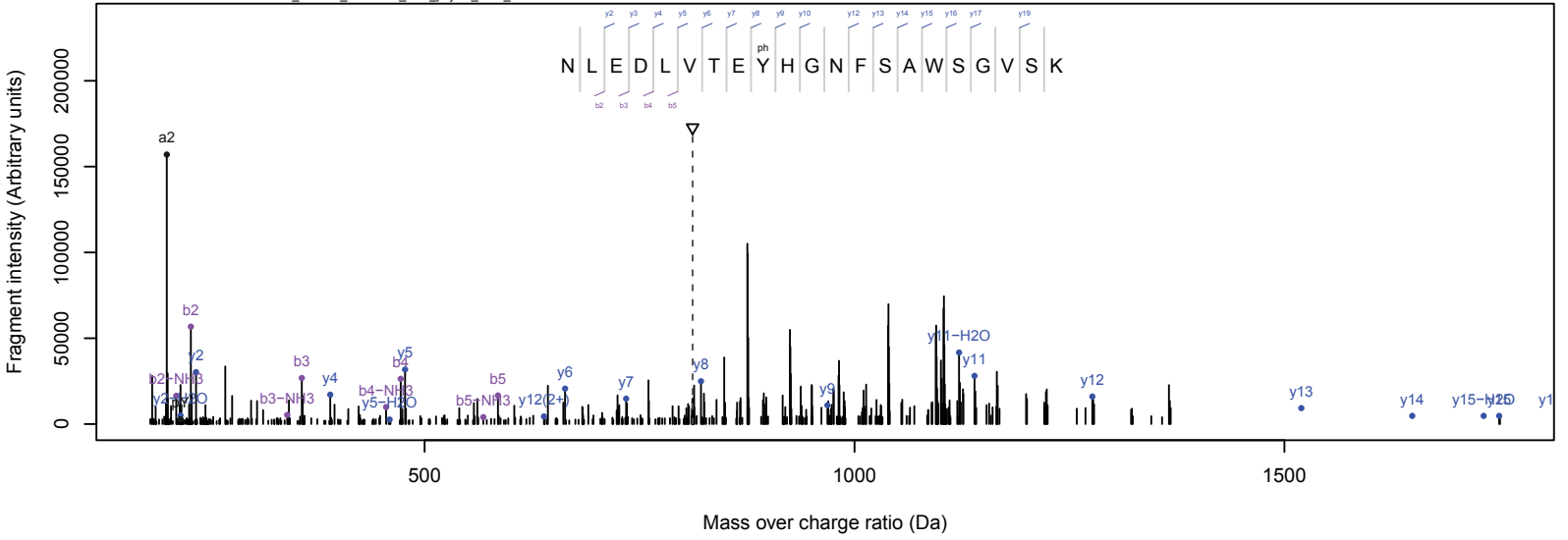
Andromeda score : 135.91



NLEDLVTEY(ph)HGNSAWSGVSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 46591

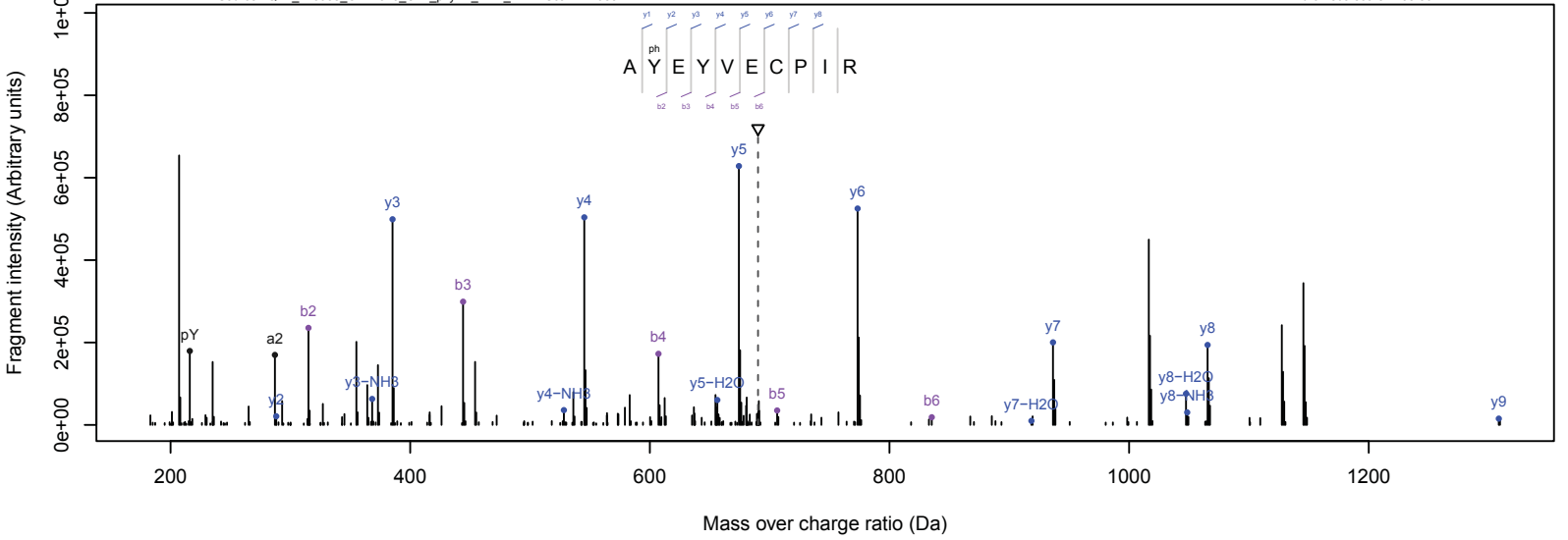
Andromeda score : 135.91

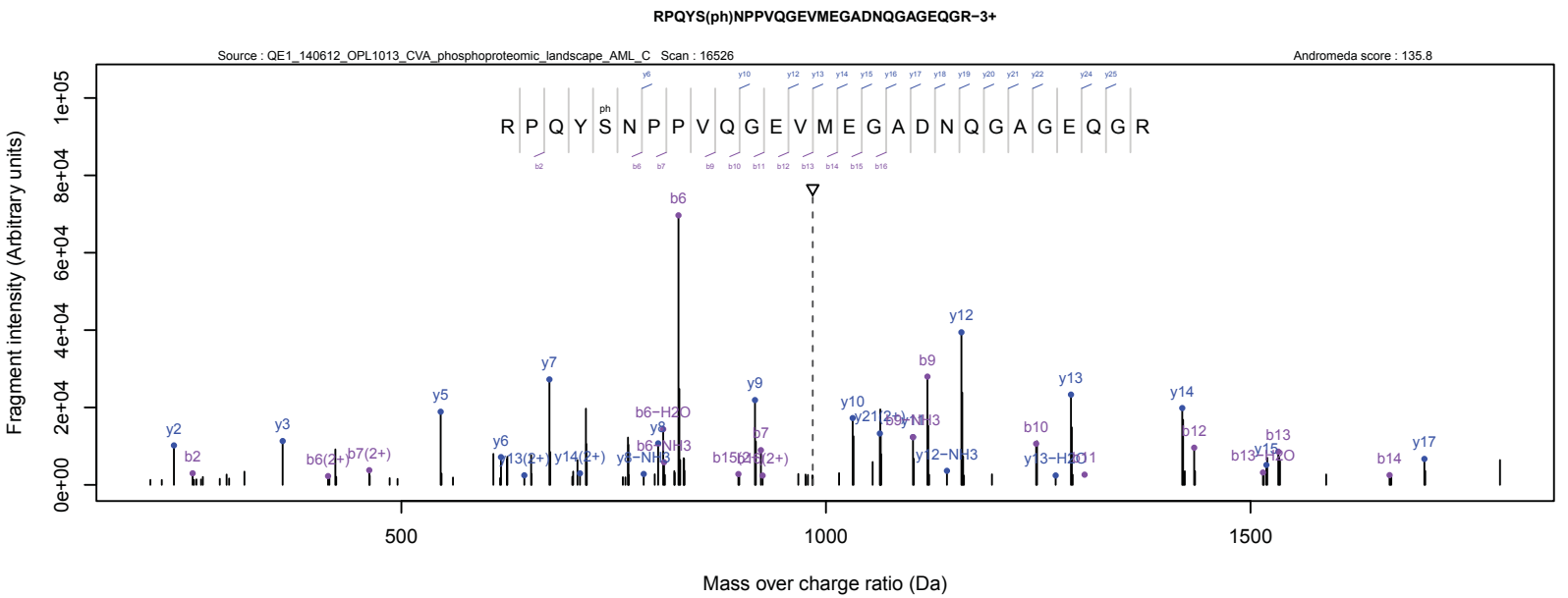
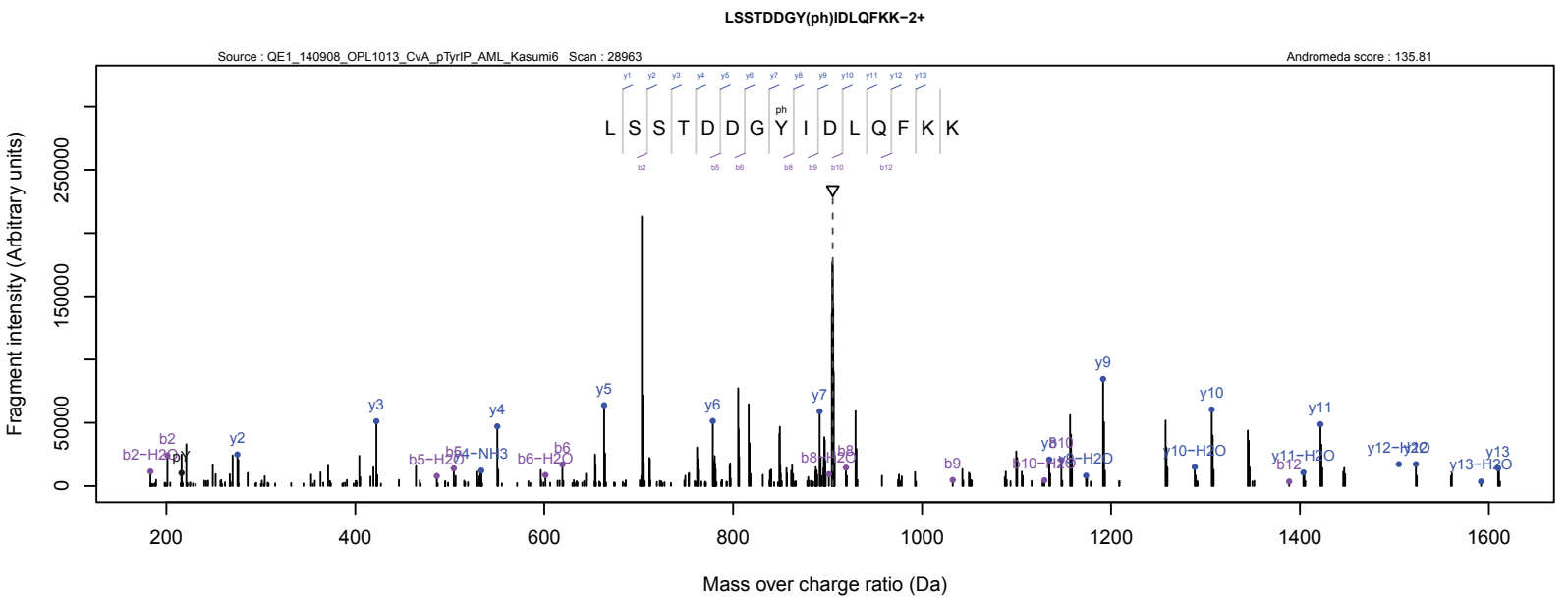
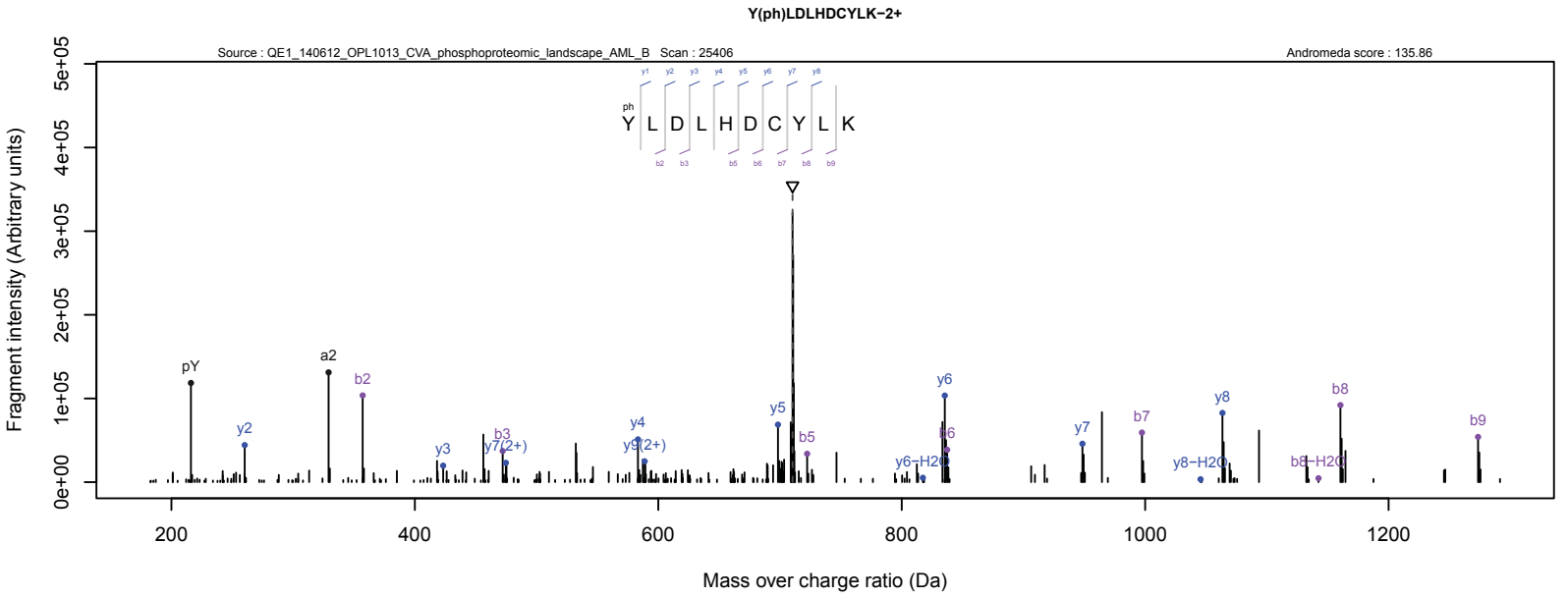


AY(ph)EYVECPPIR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 21985

Andromeda score : 135.86

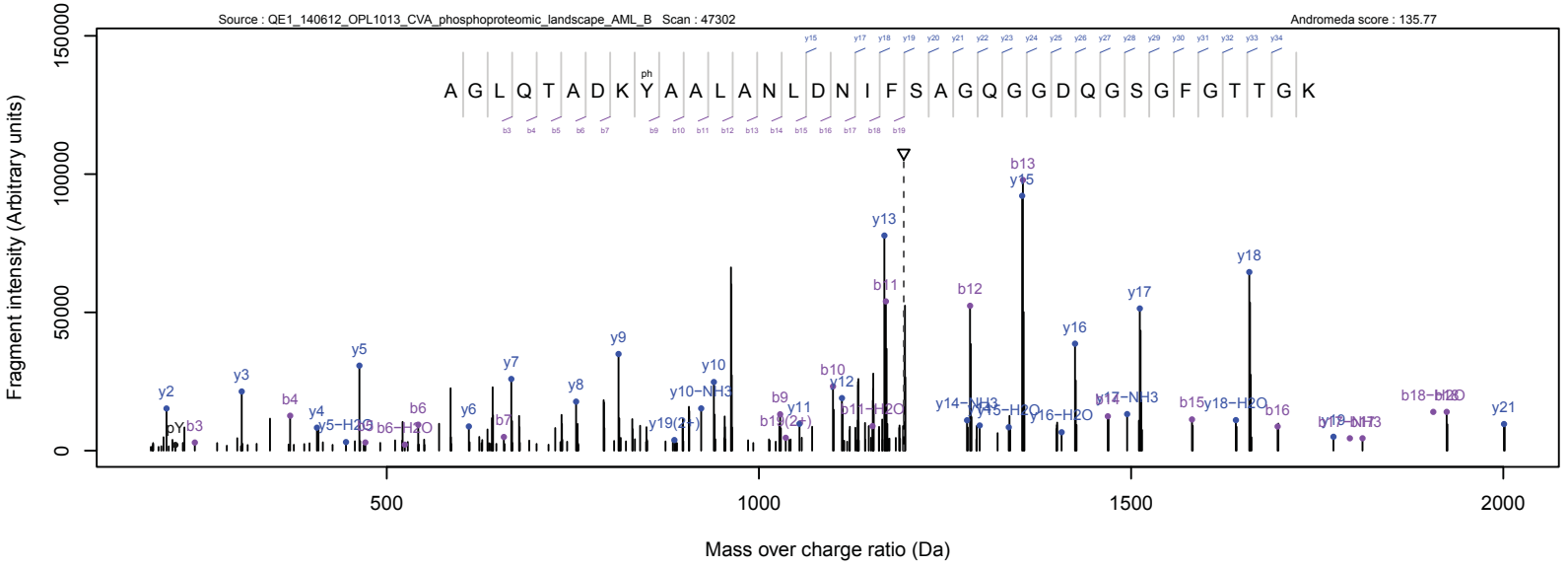




**AGLQTADKY(ph)AALANLDNIFSAAGQGQGGDQGSFGTTGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47302

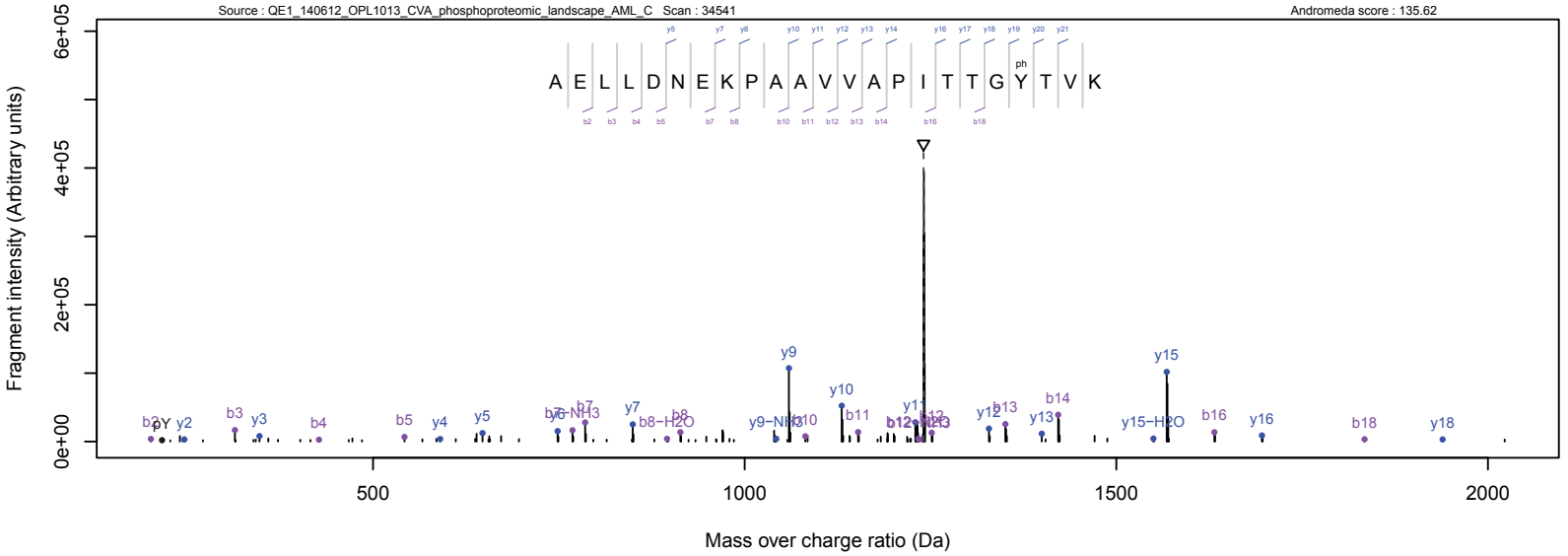
Andromeda score : 135.77



**AELLDNEKPAAVVAPITTTG(ph)TVK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34541

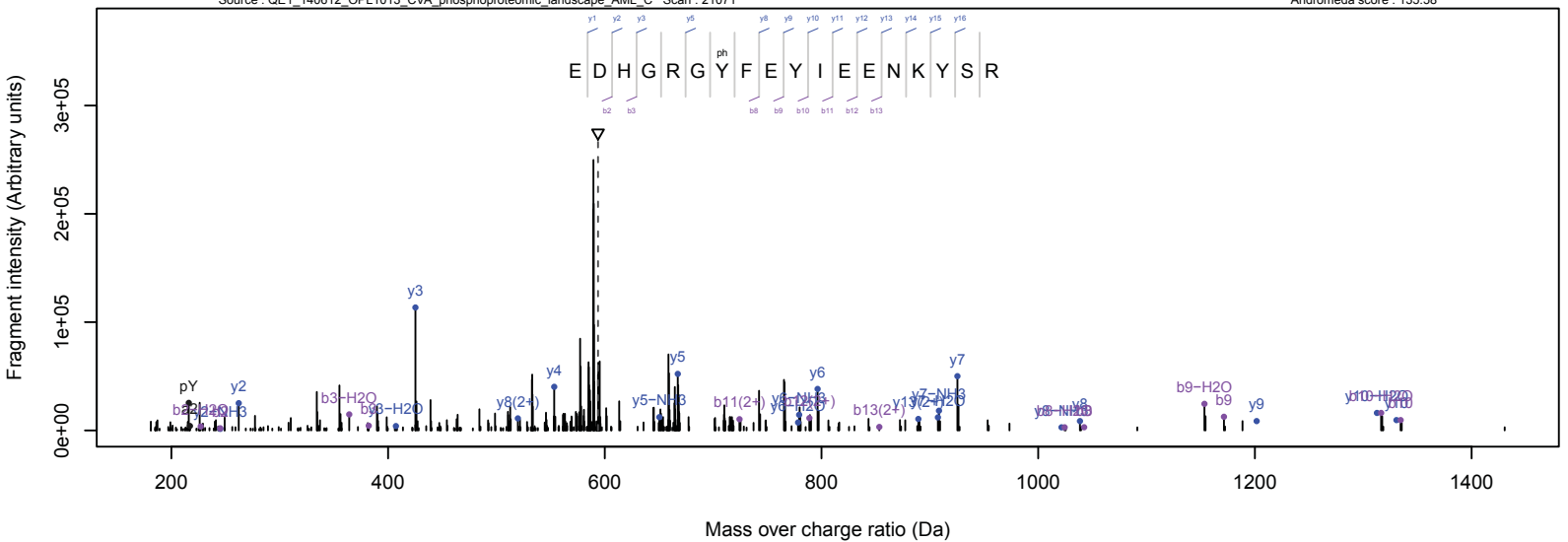
Andromeda score : 135.62



**EDHGRGYP(h)FEYIEENKYSR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21071

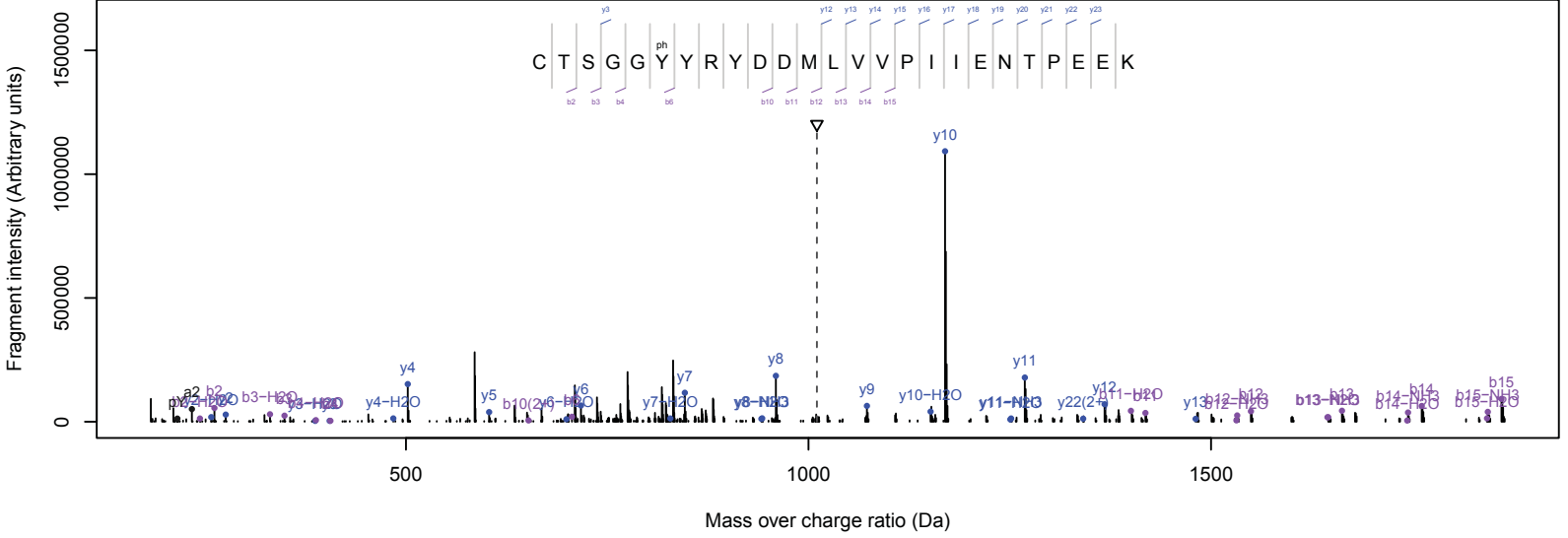
Andromeda score : 135.58



CTSGGY(ph)YRDDMLVVPIIENTPEEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46005

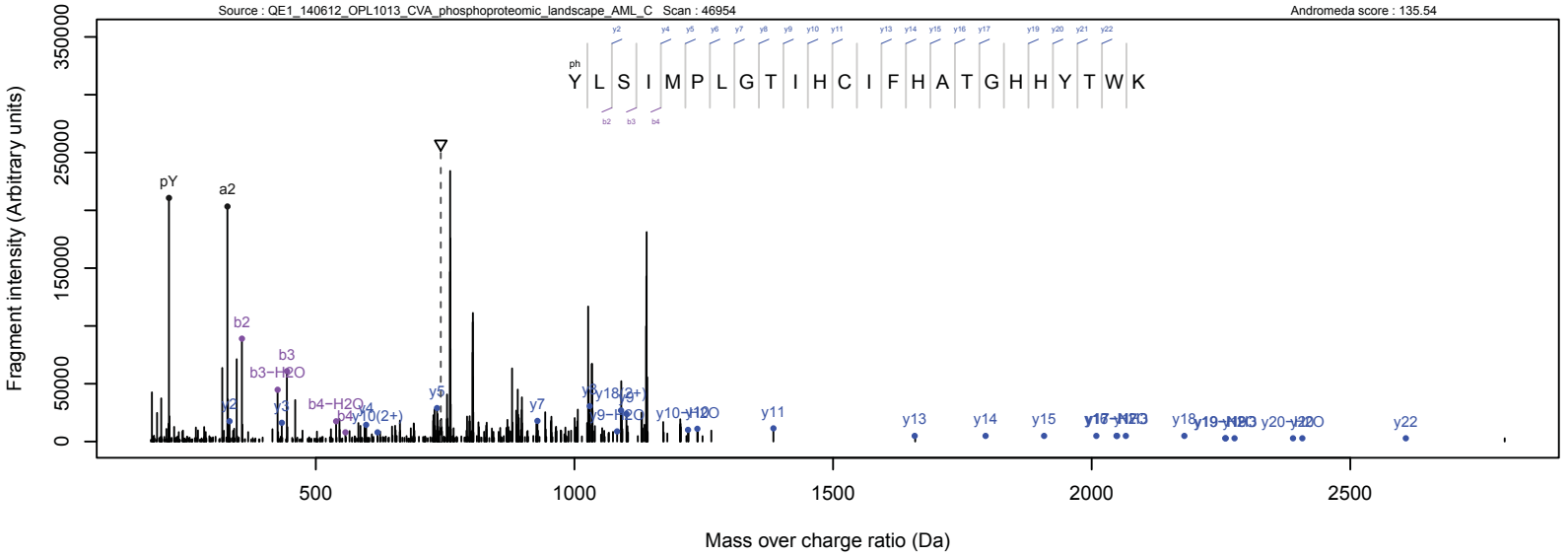
Andromeda score : 135.55



Y(ph)LSIMPLGTIHCIFHATGHHYTWK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46954

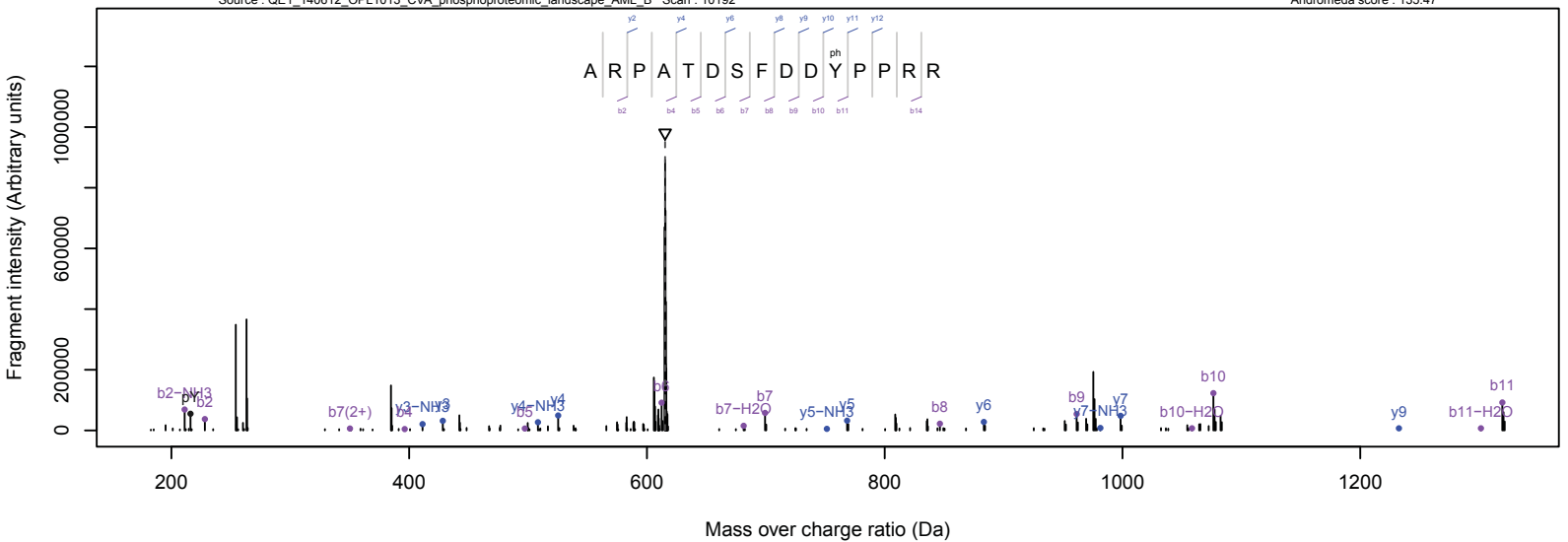
Andromeda score : 135.54



ARPATDSFDDY(ph)PPRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10192

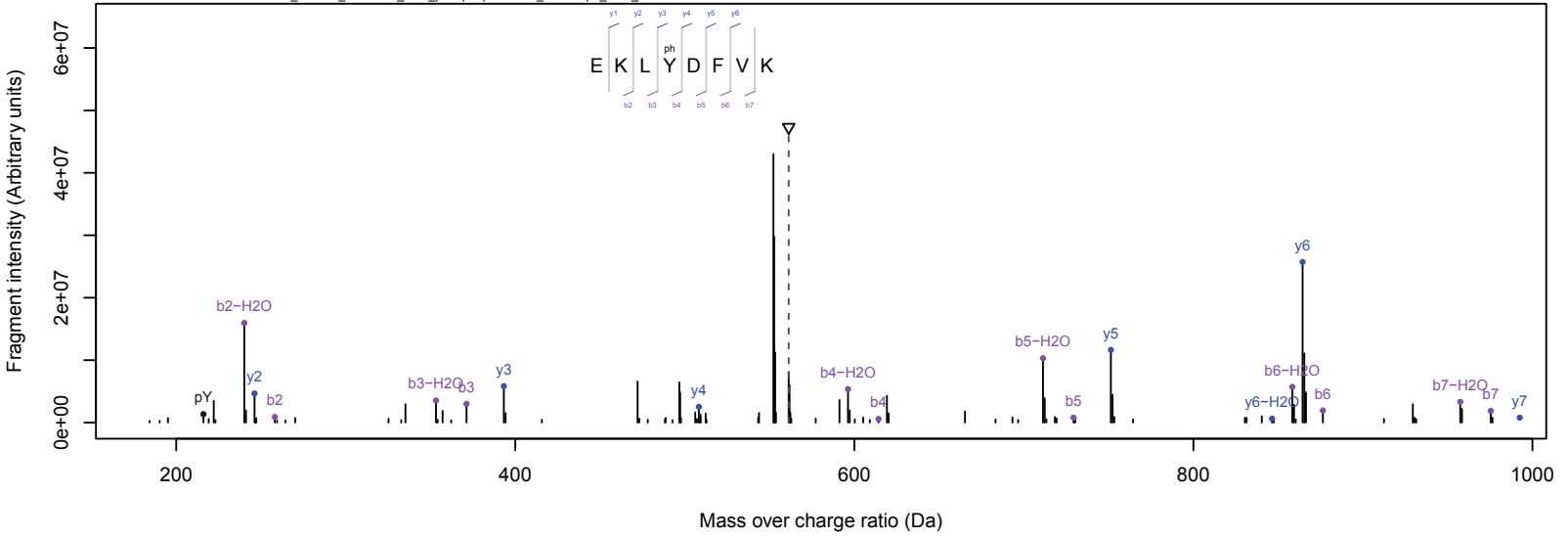
Andromeda score : 135.47



**EKLY(ph)DFVK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 21625

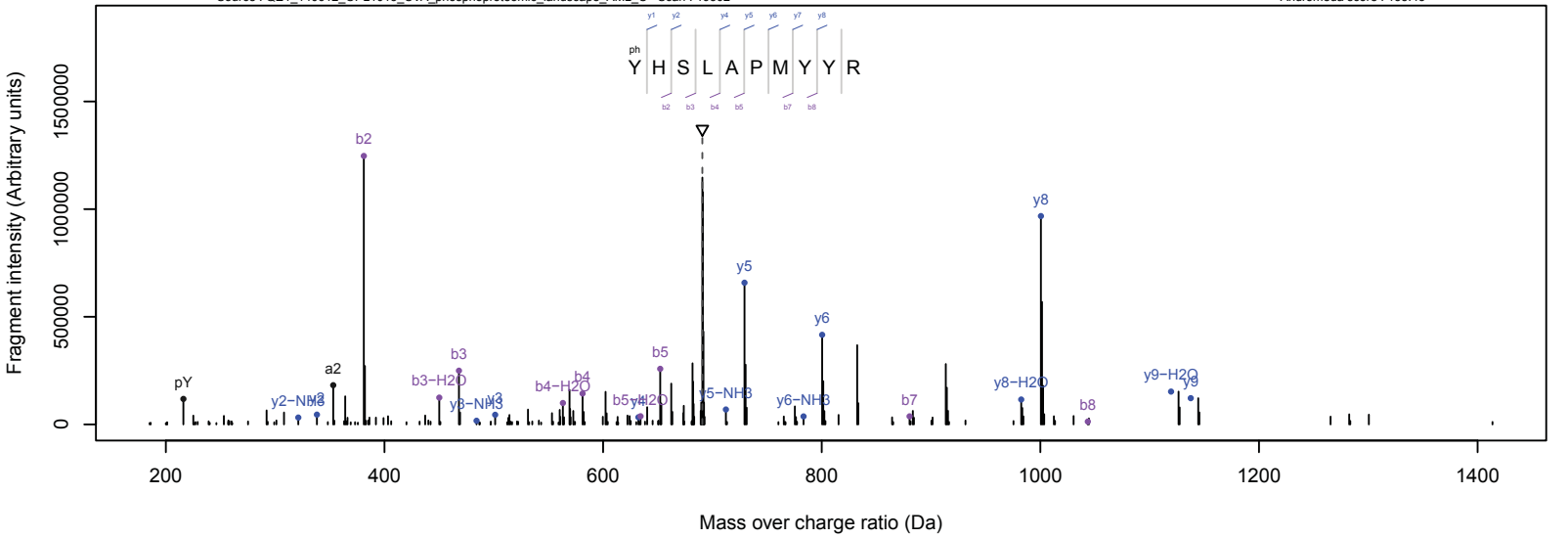
Andromeda score: 135.43



**Y(ph)HSLAPMYR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19302

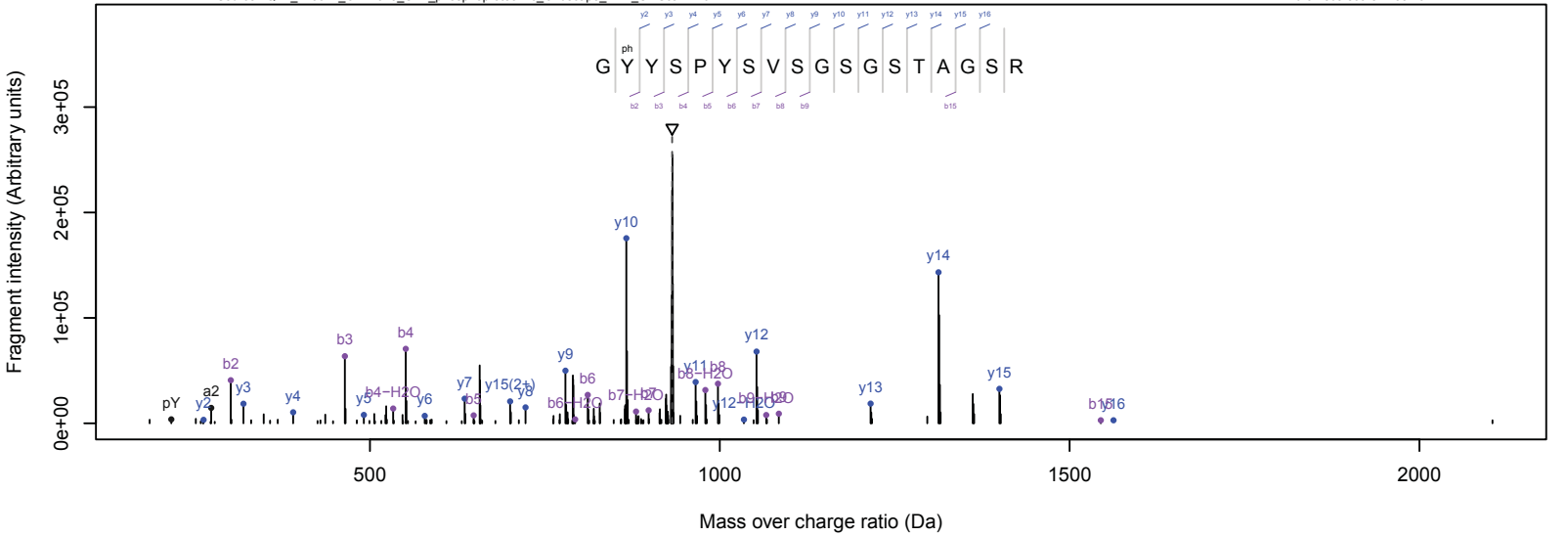
Andromeda score: 135.43



**GY(ph)YSPYVS GSGSTAGSR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16222

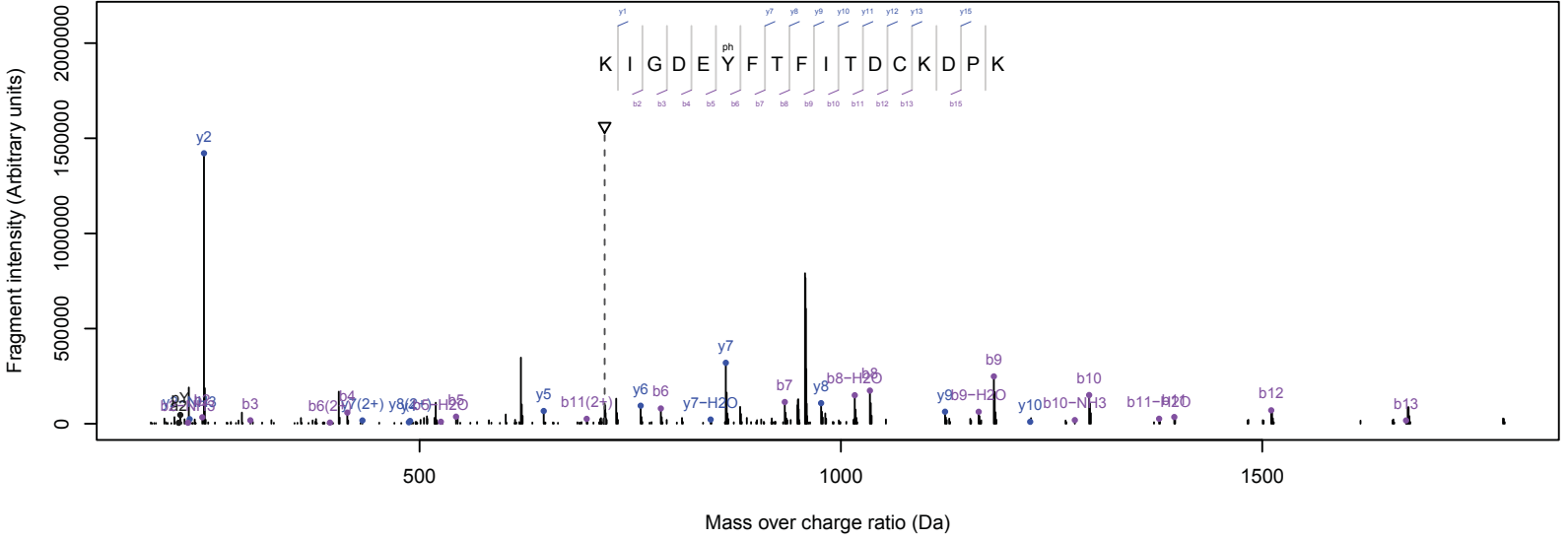
Andromeda score: 135.29



KIGDEY(ph)FTTDCDKPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31944

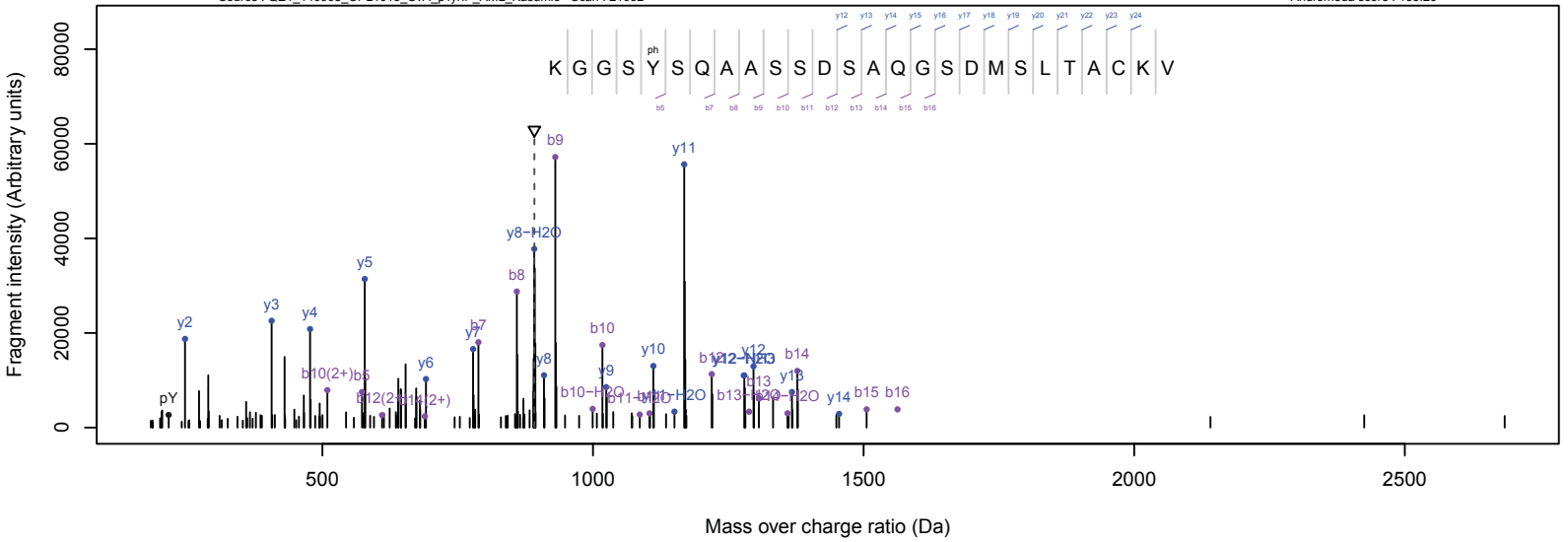
Andromeda score : 135.29



KGGSY(ph)SQAASSDSAQGS DMSLTACKV-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 21052

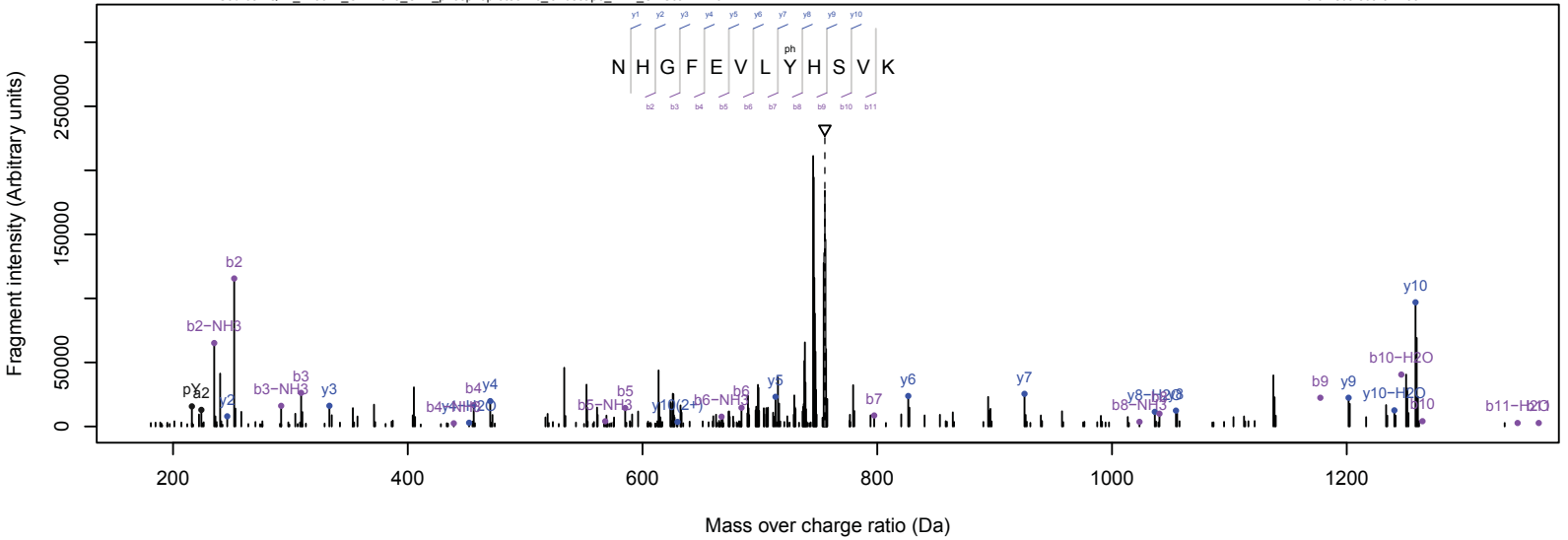
Andromeda score : 135.25



NHGFVLY(ph)HSVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21074

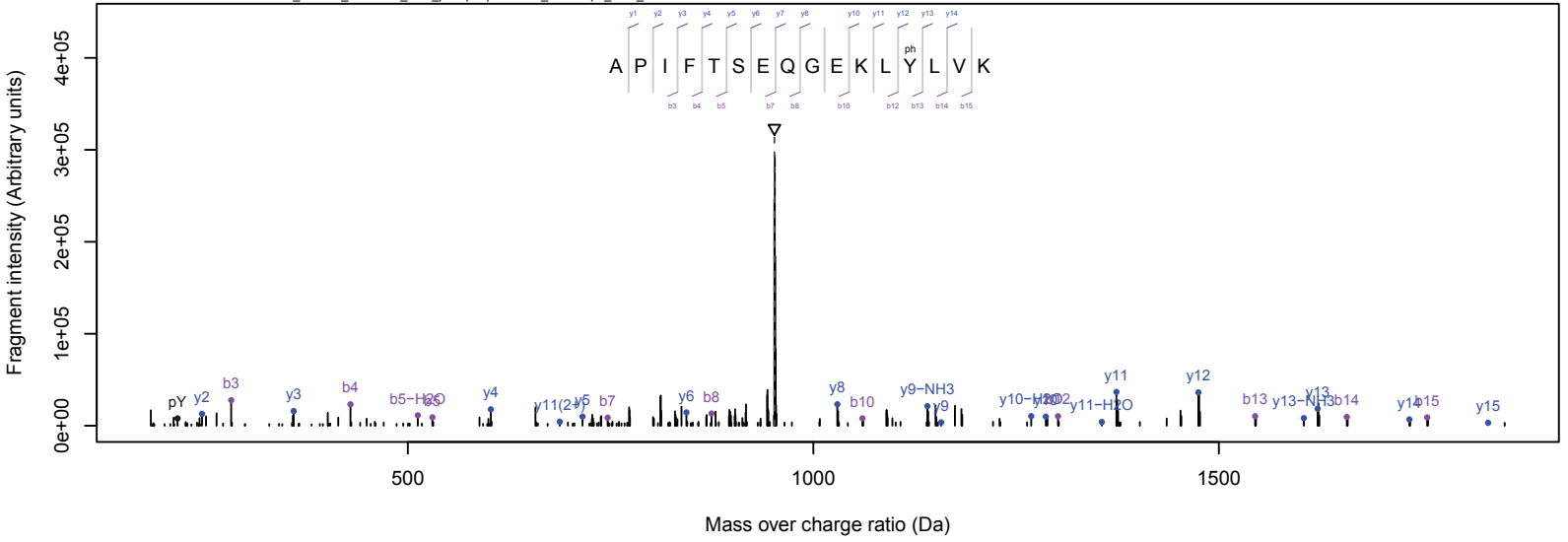
Andromeda score : 135.24



APIFTSEQGEKLY(ph)LVK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 33155

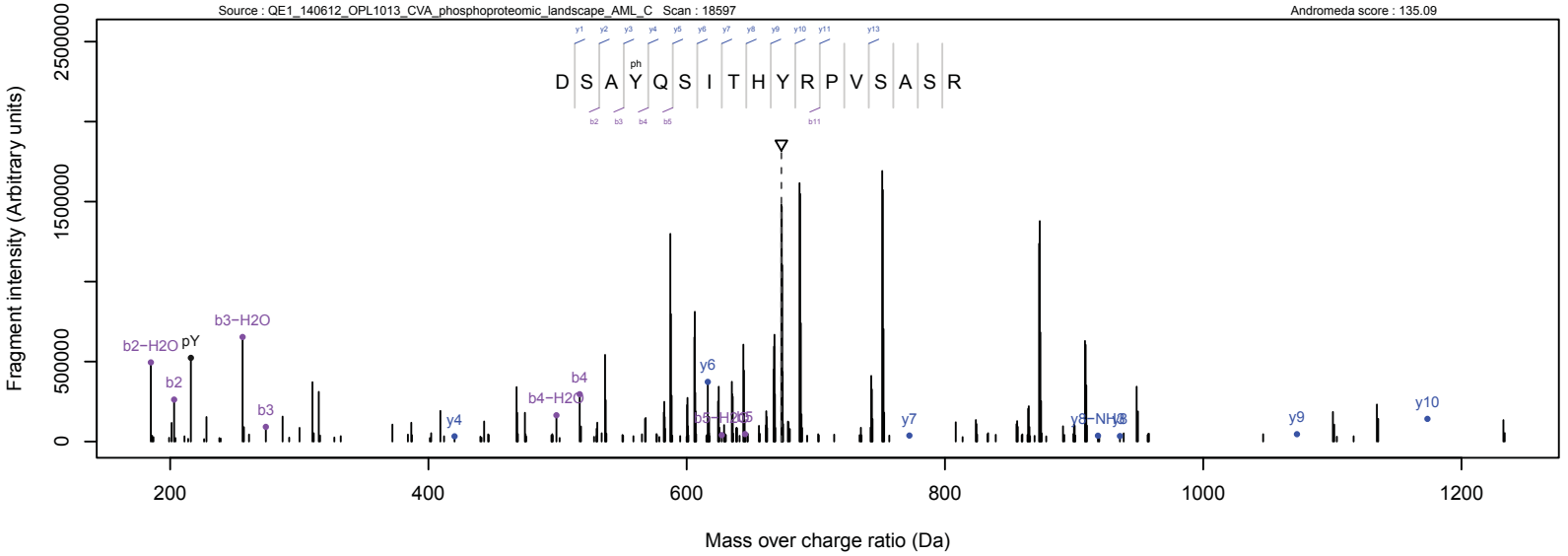
Andromeda score: 135.09



DSAY(ph)QSITHYRVPVSASR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18597

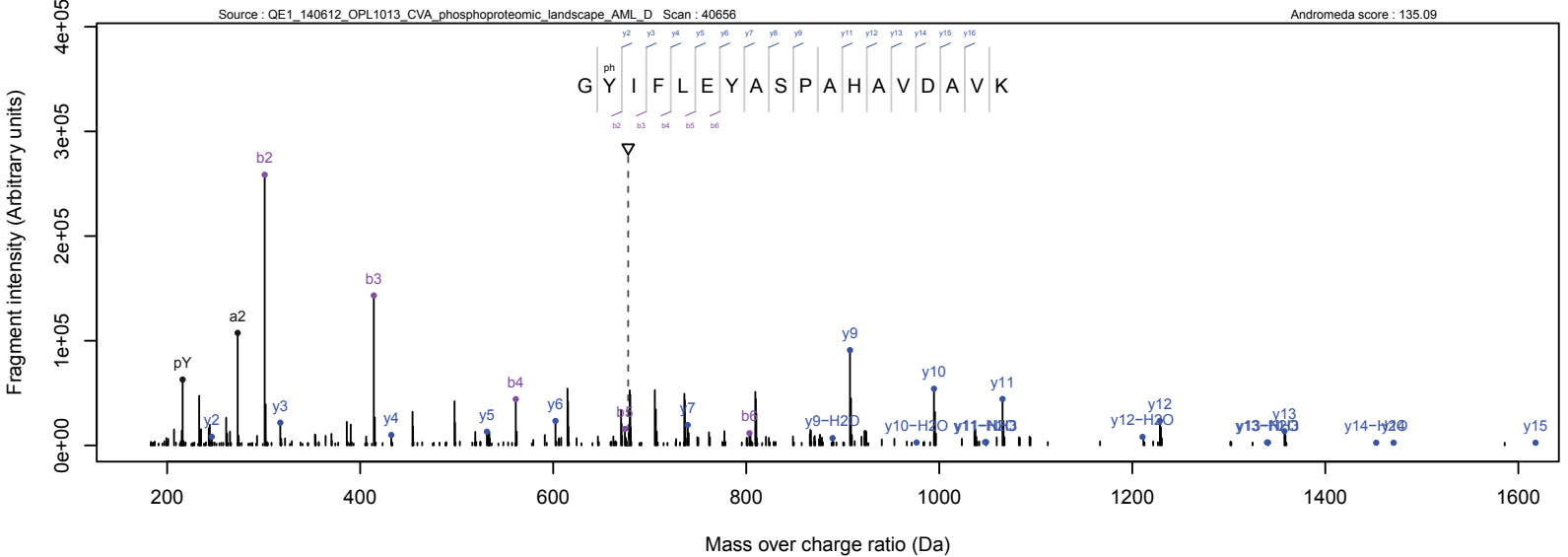
Andromeda score: 135.09



GY(ph)FLEYASPAHVAVDVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 40656

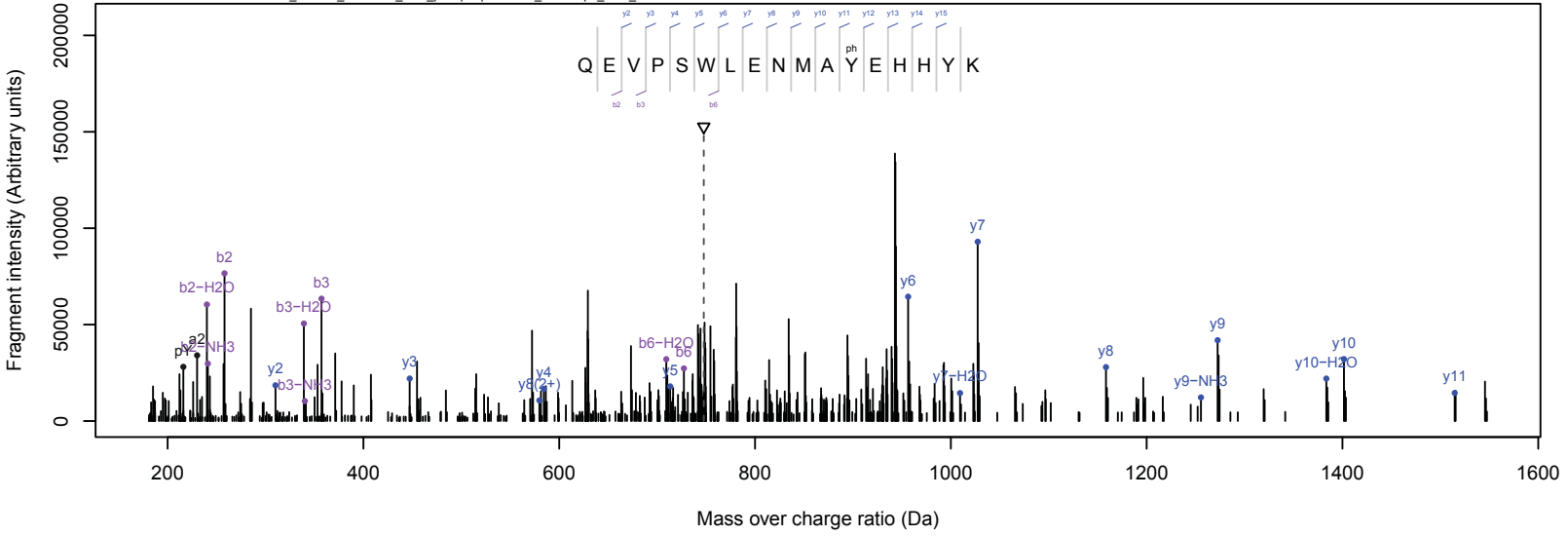
Andromeda score: 135.09



QEVPSWLENMAY(ph)EHYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34700

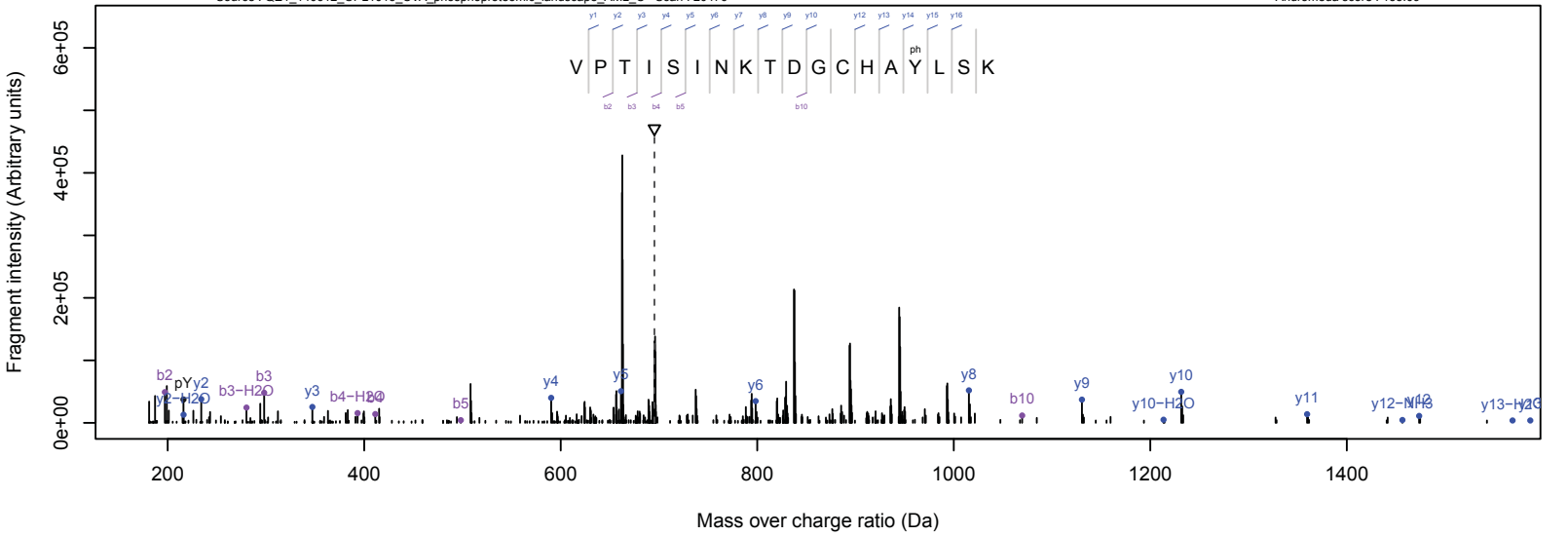
Andromeda score : 135.09



VPTISINKTDGCHAY(ph)LSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20476

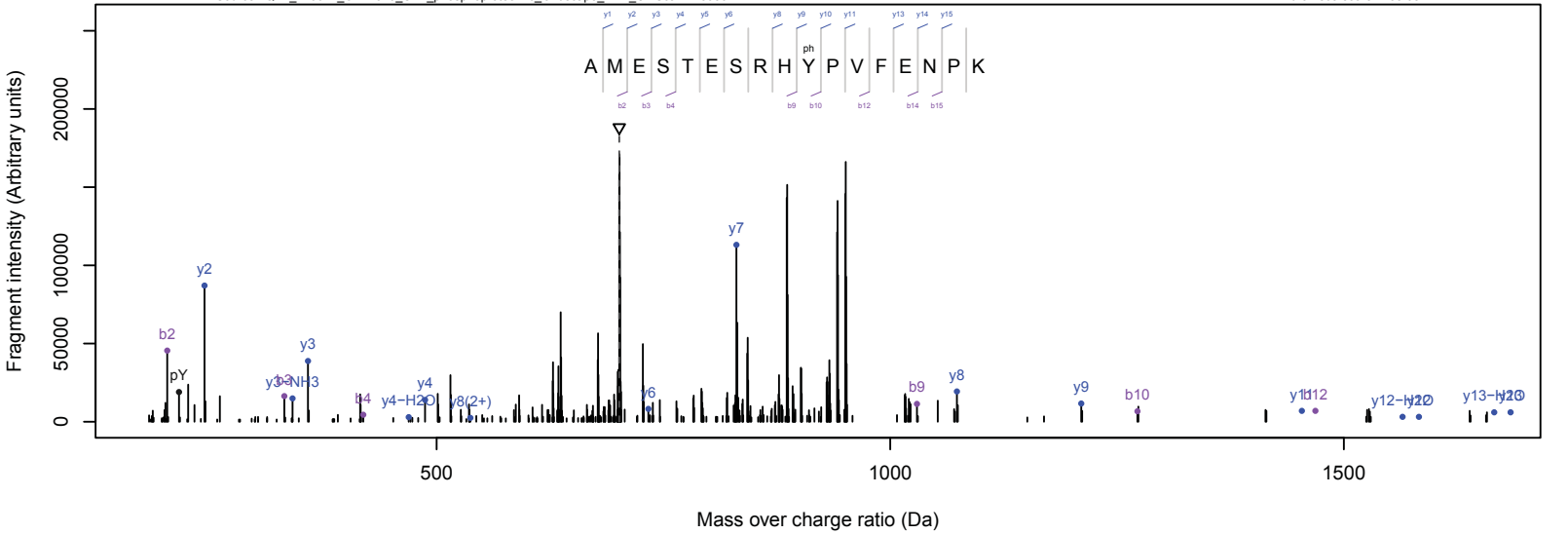
Andromeda score : 135.09



AMESTESRHY(ph)PVFENPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13950

Andromeda score : 135.08

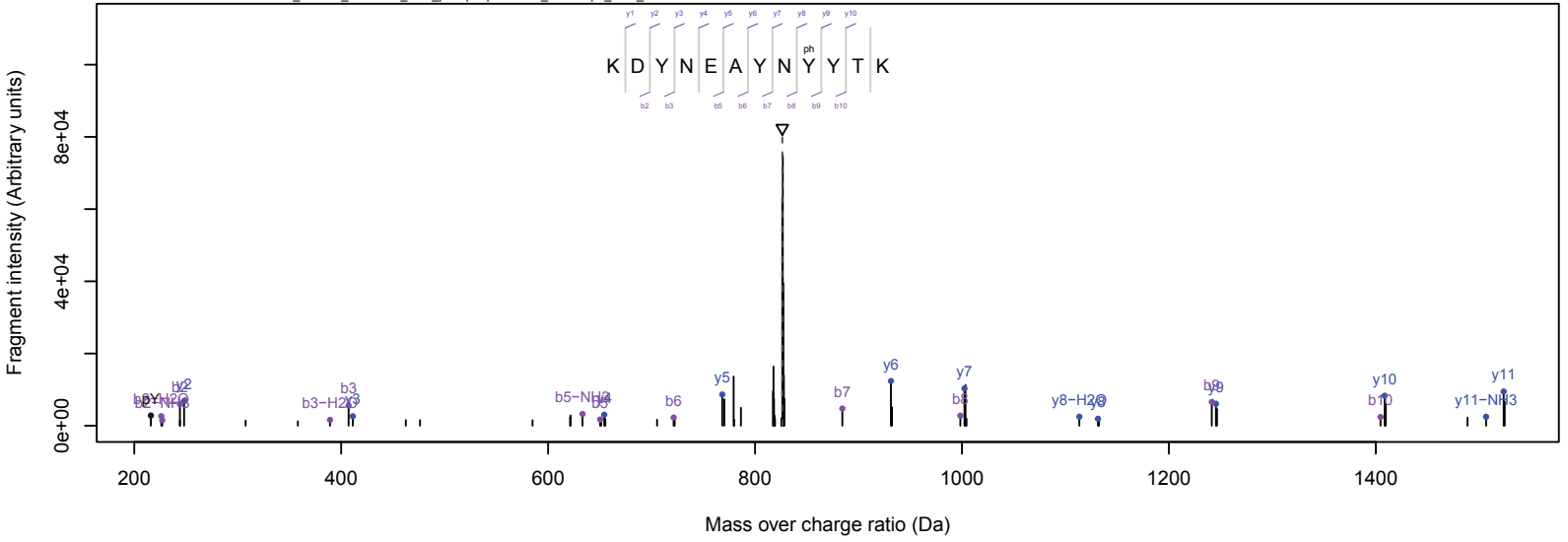




**KDYNEAYNYTK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14246

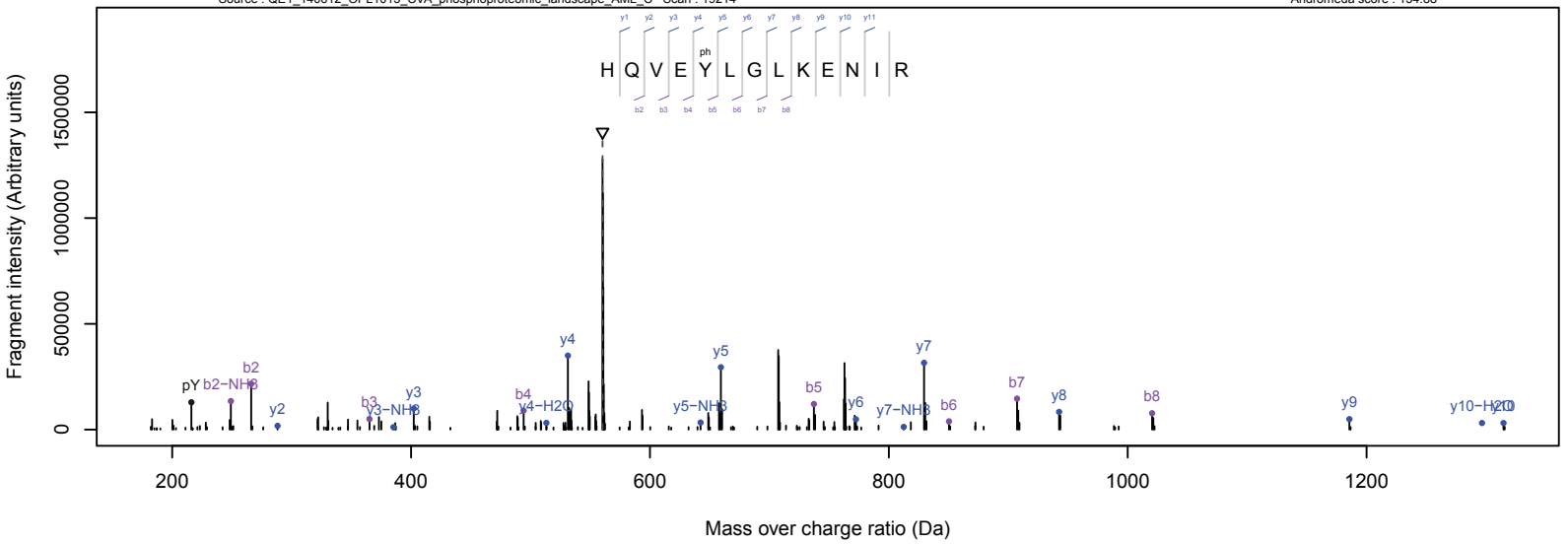
Andromeda score : 135.02



**HQVEY(p)LGLKENIR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19214

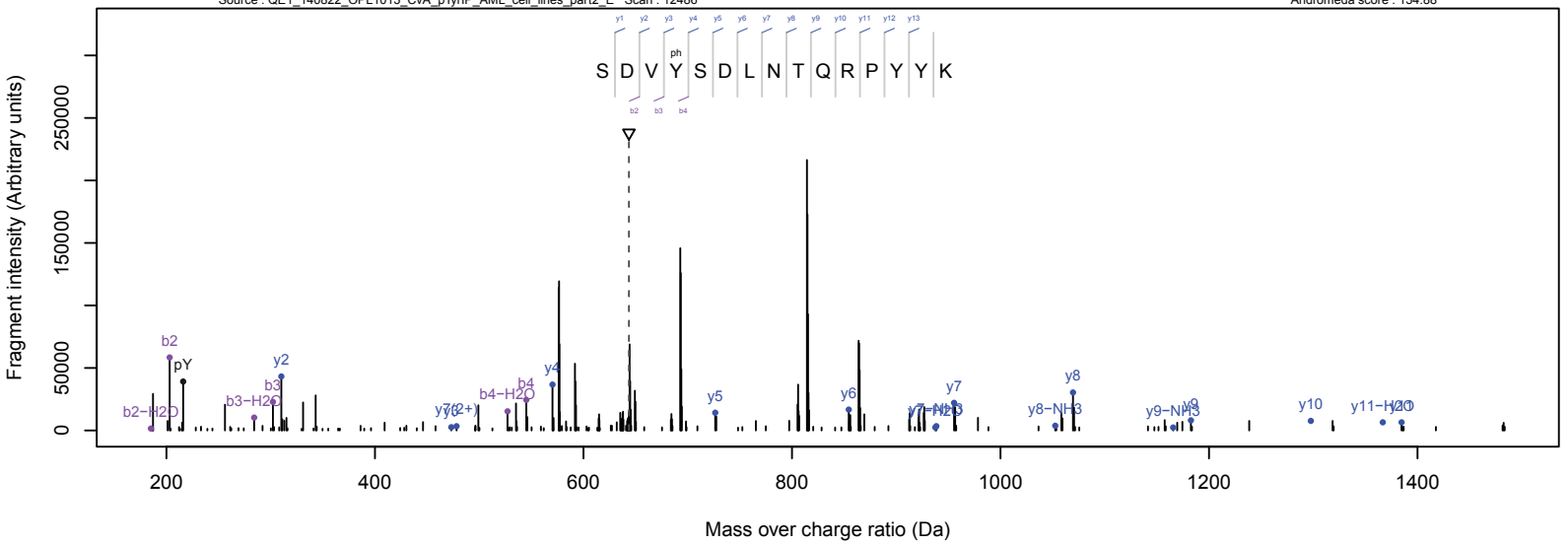
Andromeda score : 134.88



**SDVY(p)SDLNTQRPYYK-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 12486

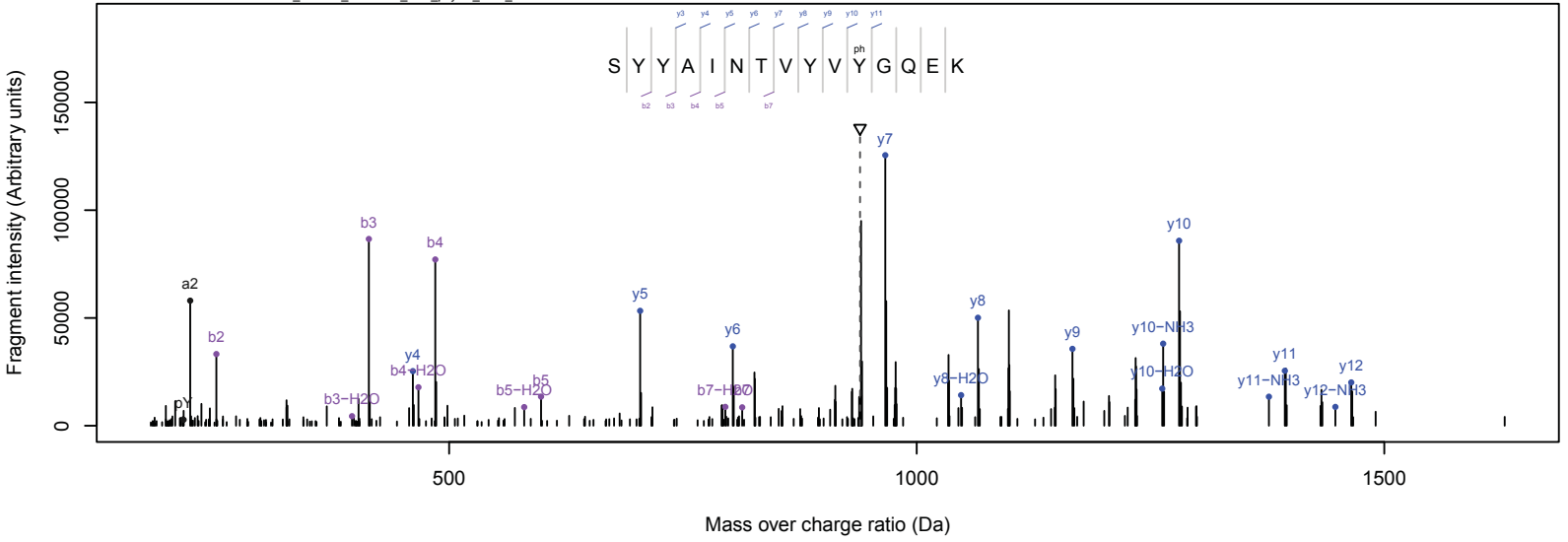
Andromeda score : 134.88



SYIAINTVYVY(ph)GQEK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan : 33726

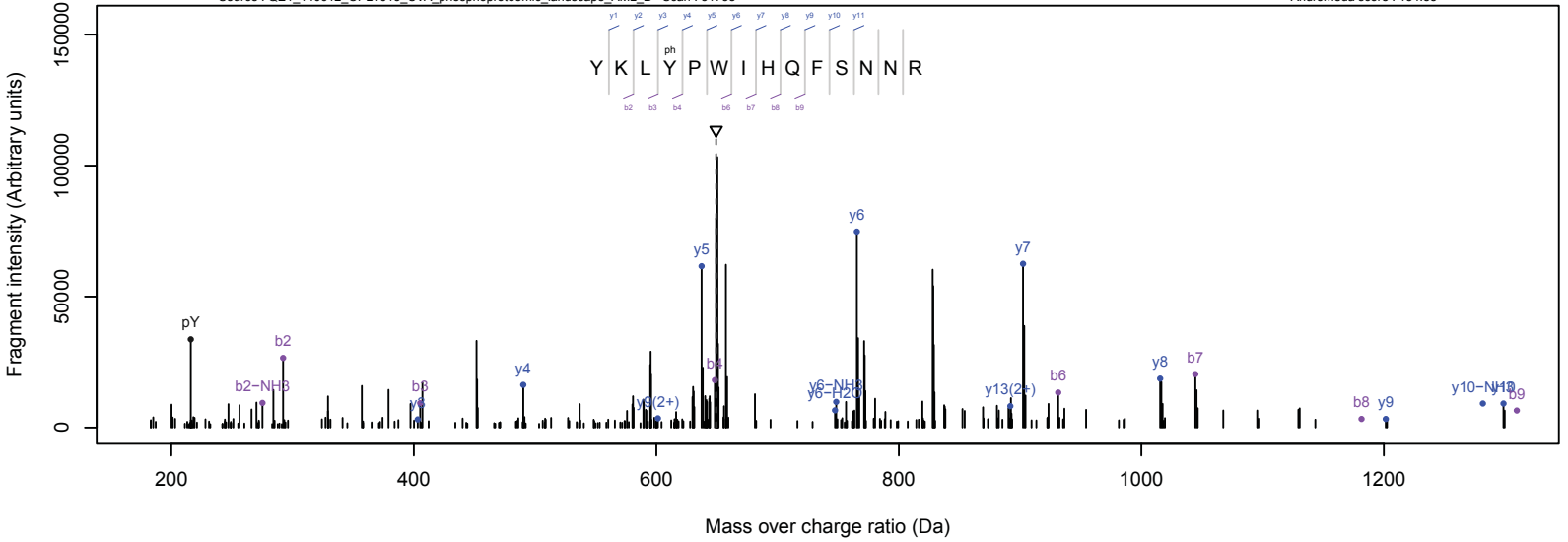
Andromeda score : 134.88



YKLY(ph)PWIHQFSNNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31738

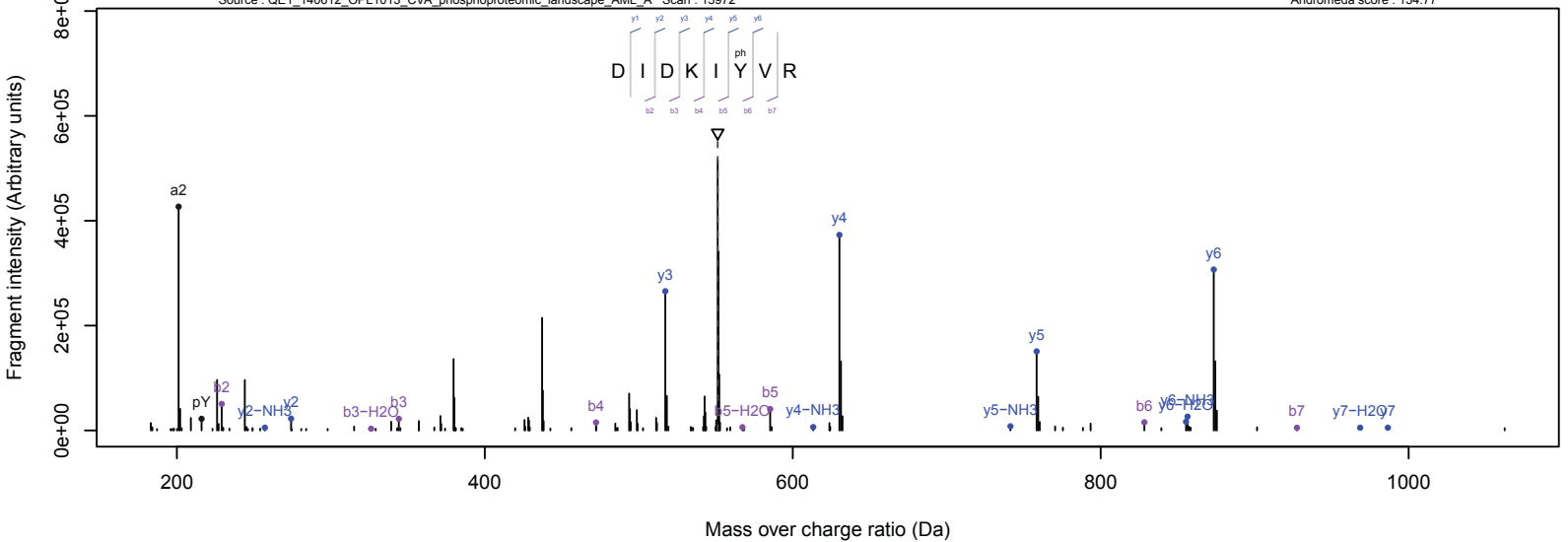
Andromeda score : 134.88



DIDKIYVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 15972

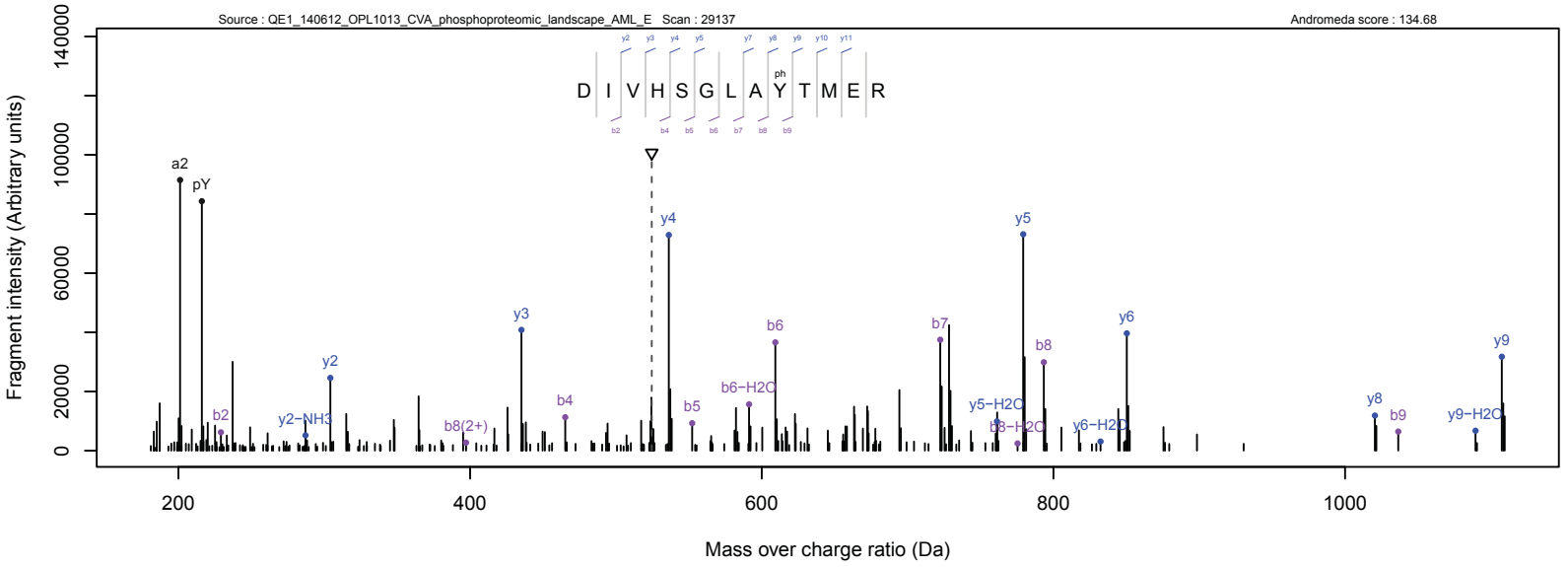
Andromeda score : 134.77



DIVHSGLAY(ph)TMER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 29137

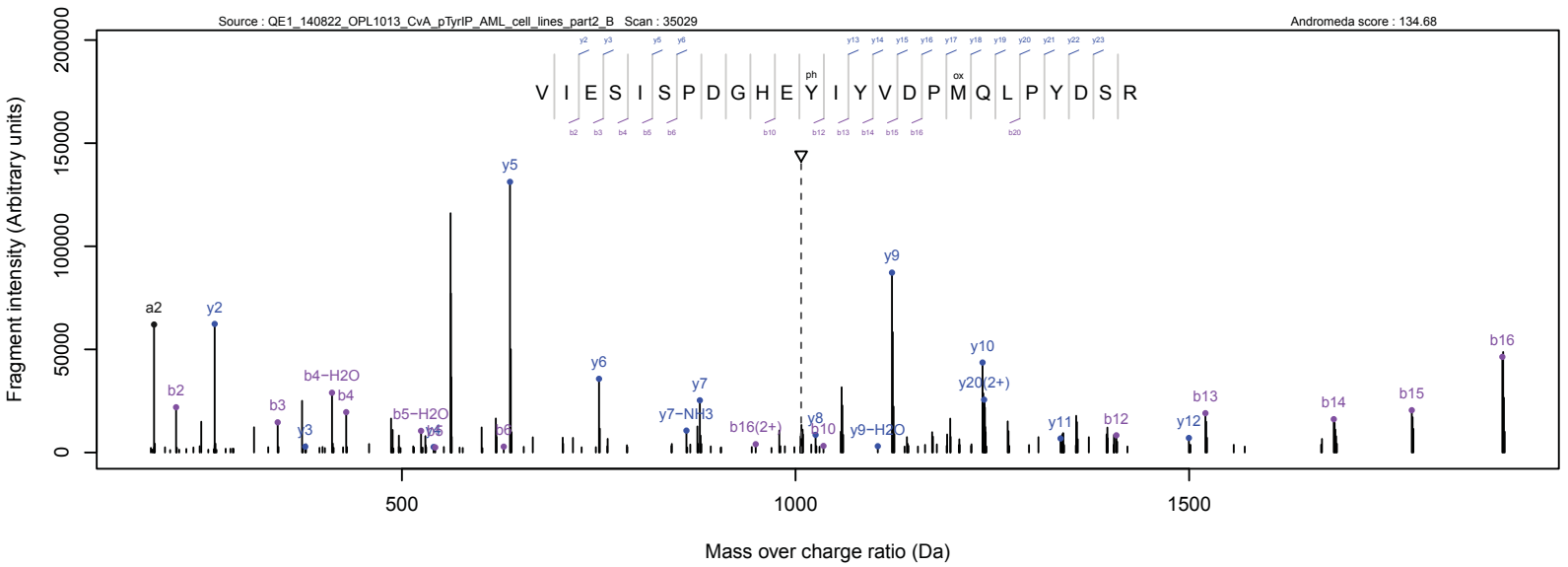
Andromeda score : 134.68



VIESISPDGHEY(ph)IYVDPM(ox)QLPYDSR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 35029

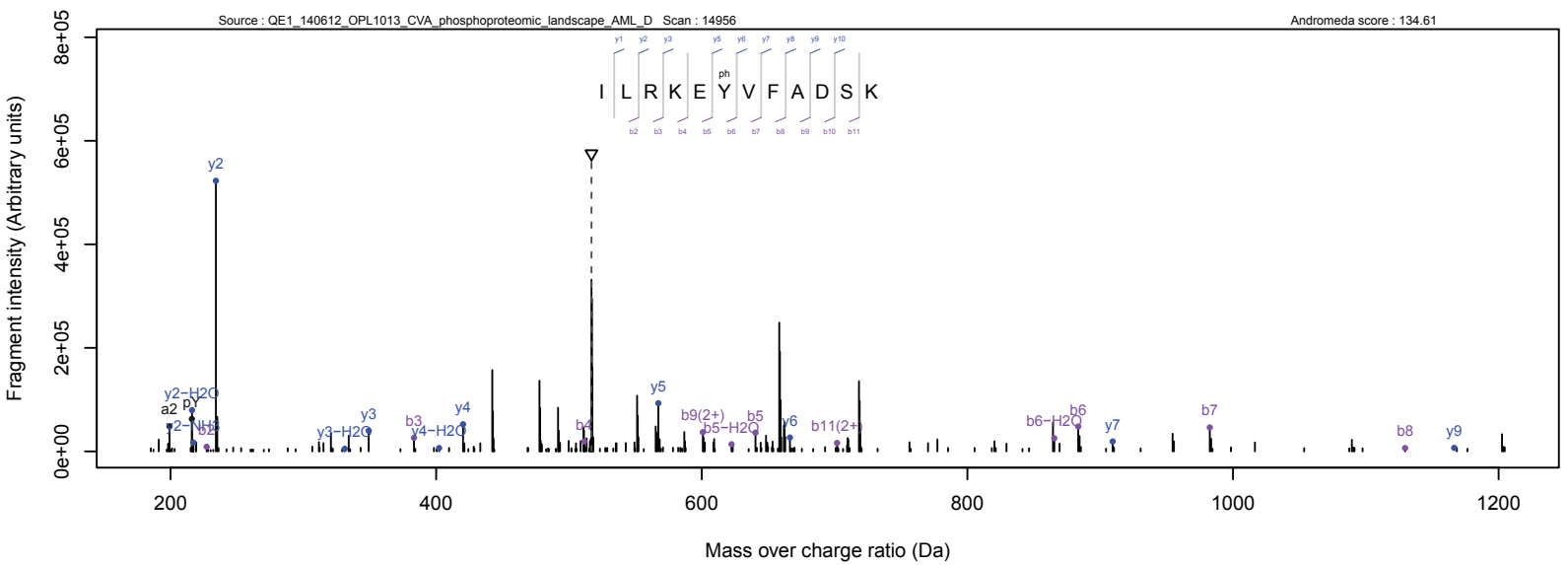
Andromeda score : 134.68



ILRKEY(ph)VFADSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14956

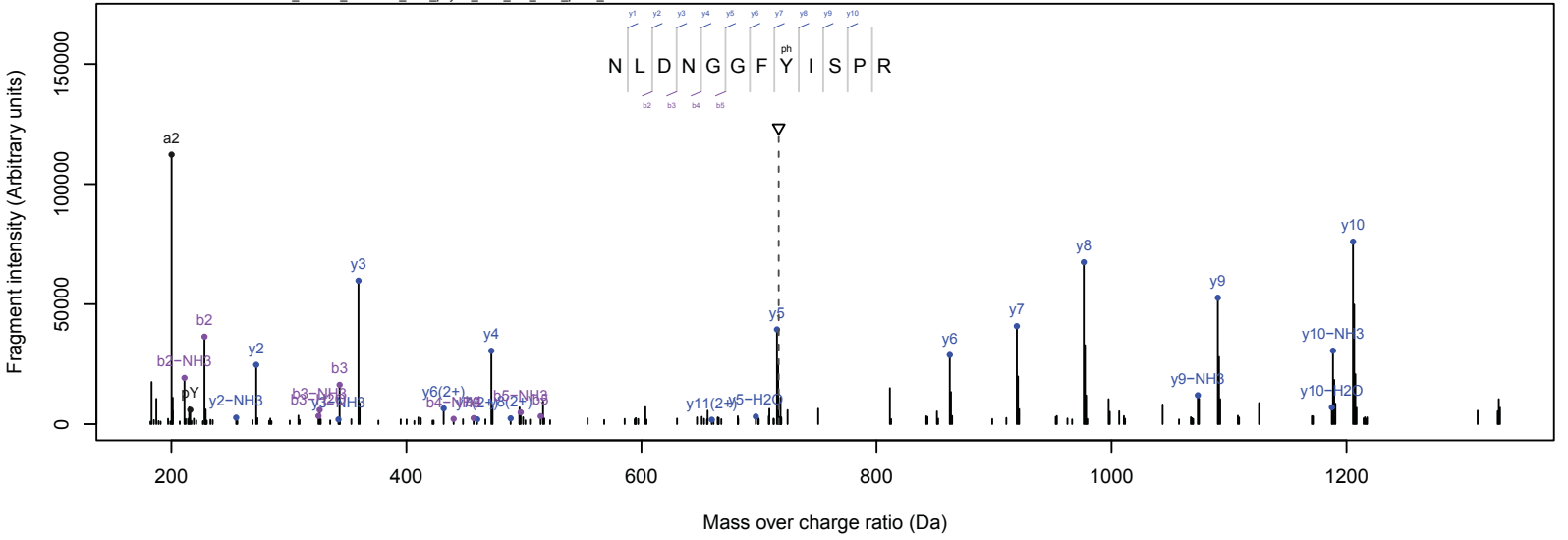
Andromeda score : 134.61



NLDNGGFY(ph)ISPR-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 21982

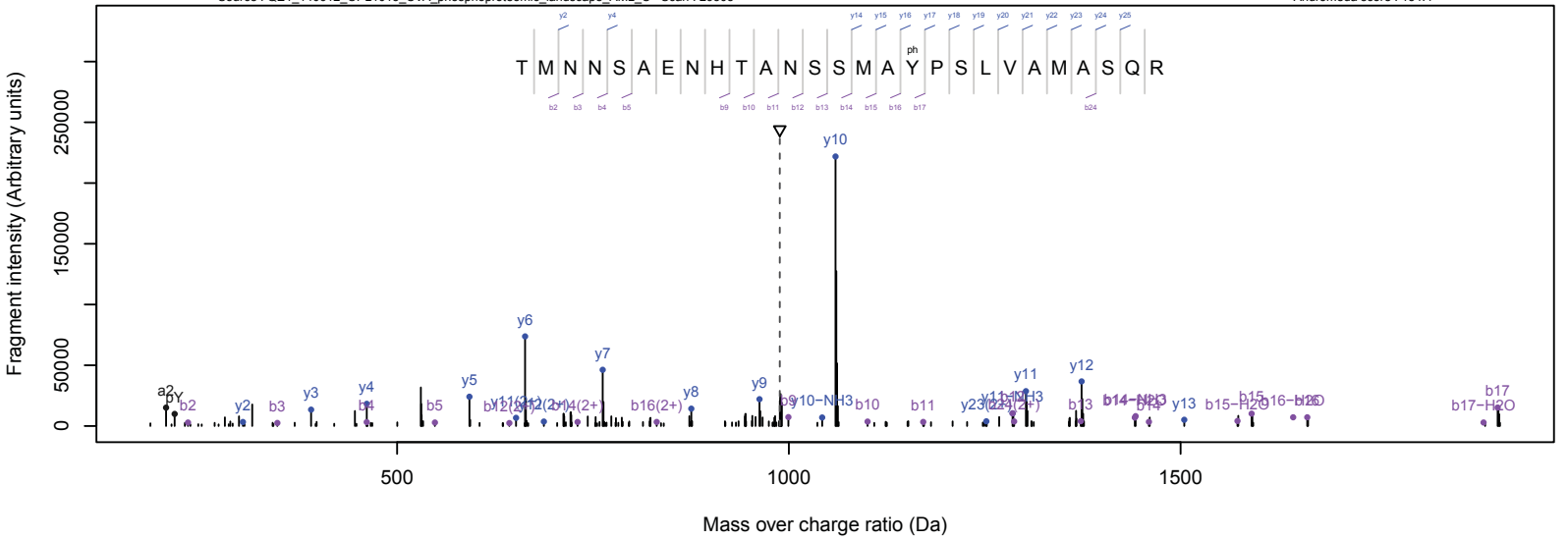
Andromeda score : 134.56



TMNNSAENHTANSSMAY(ph)PSLVAMASQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29503

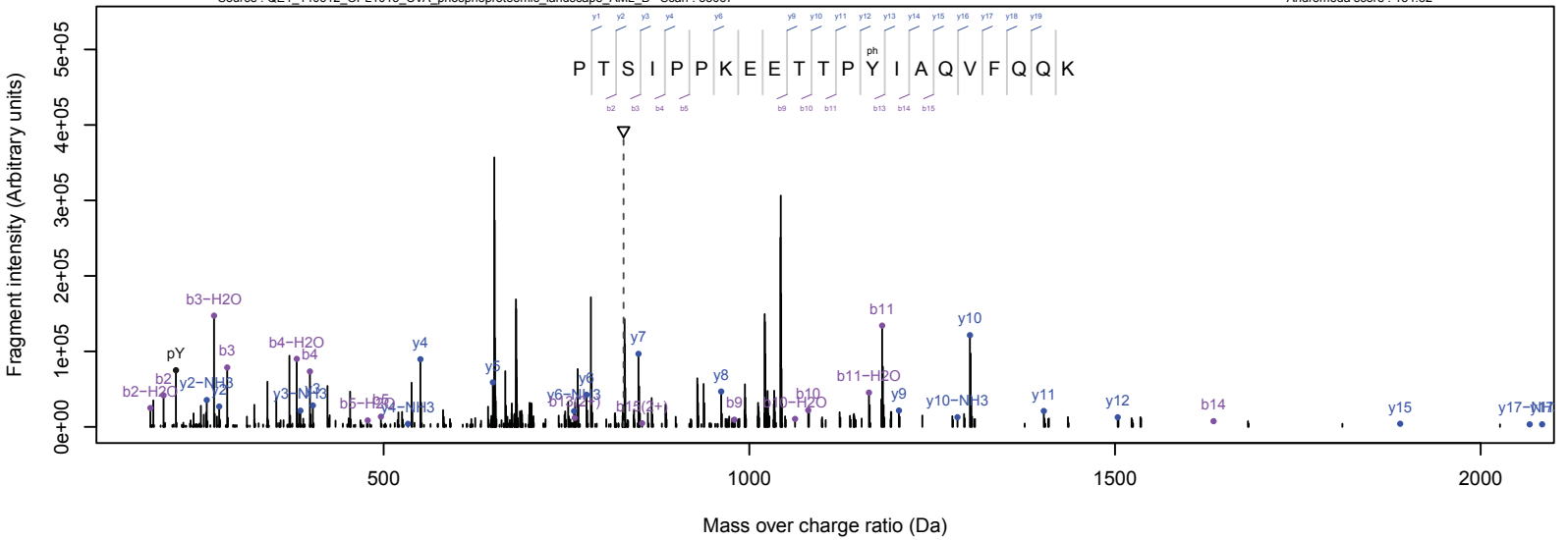
Andromeda score : 134.4



PTSIPPKEETTPY(ph)IAQVFQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33067

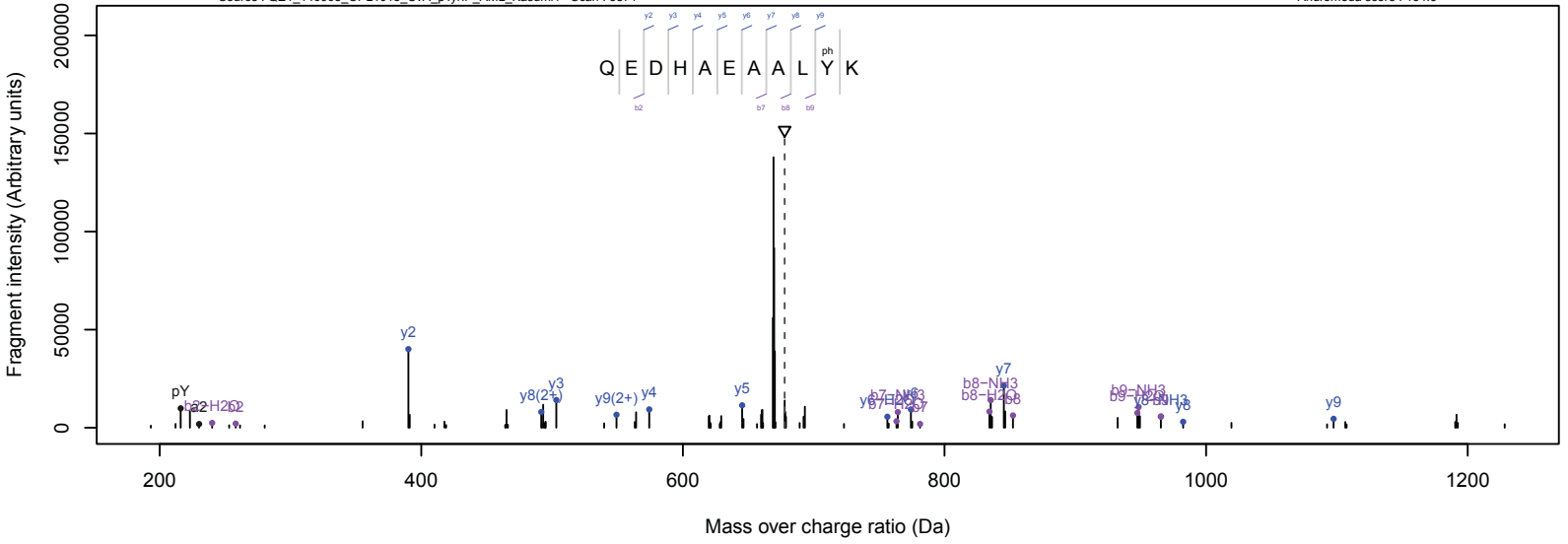
Andromeda score : 134.32



**QEDHAEALYK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 5871

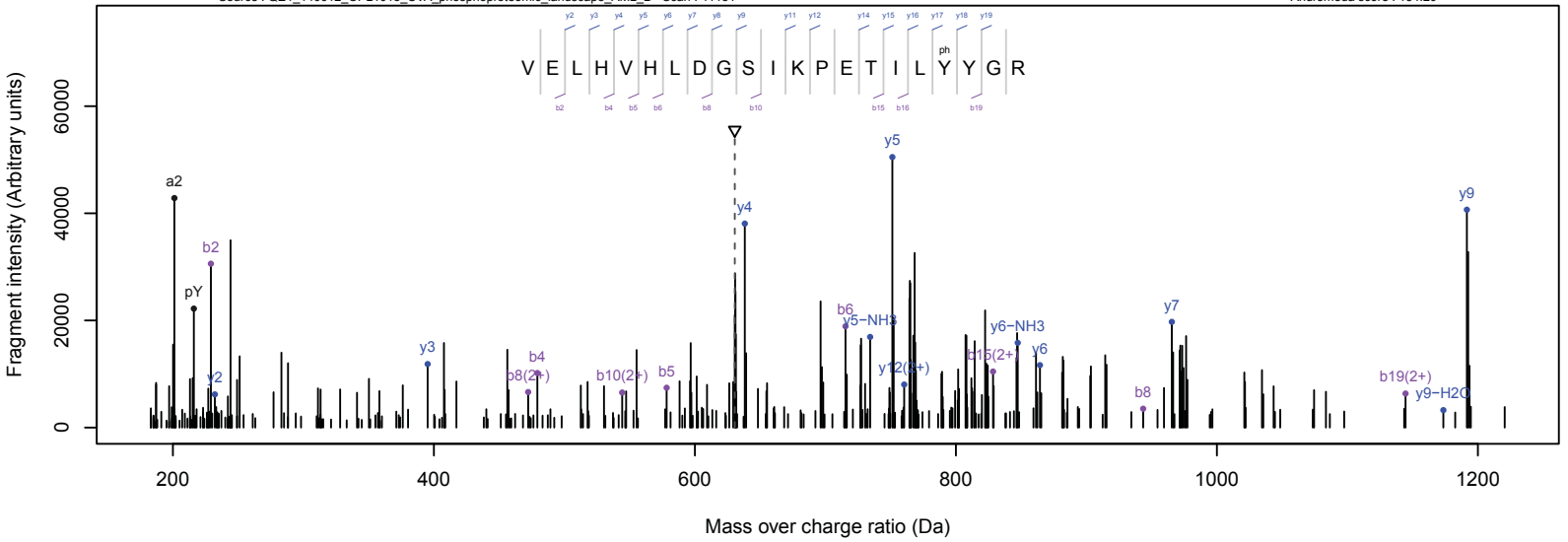
Andromeda score : 134.3



**VELHVHLDGSIKPEITILYGR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41131

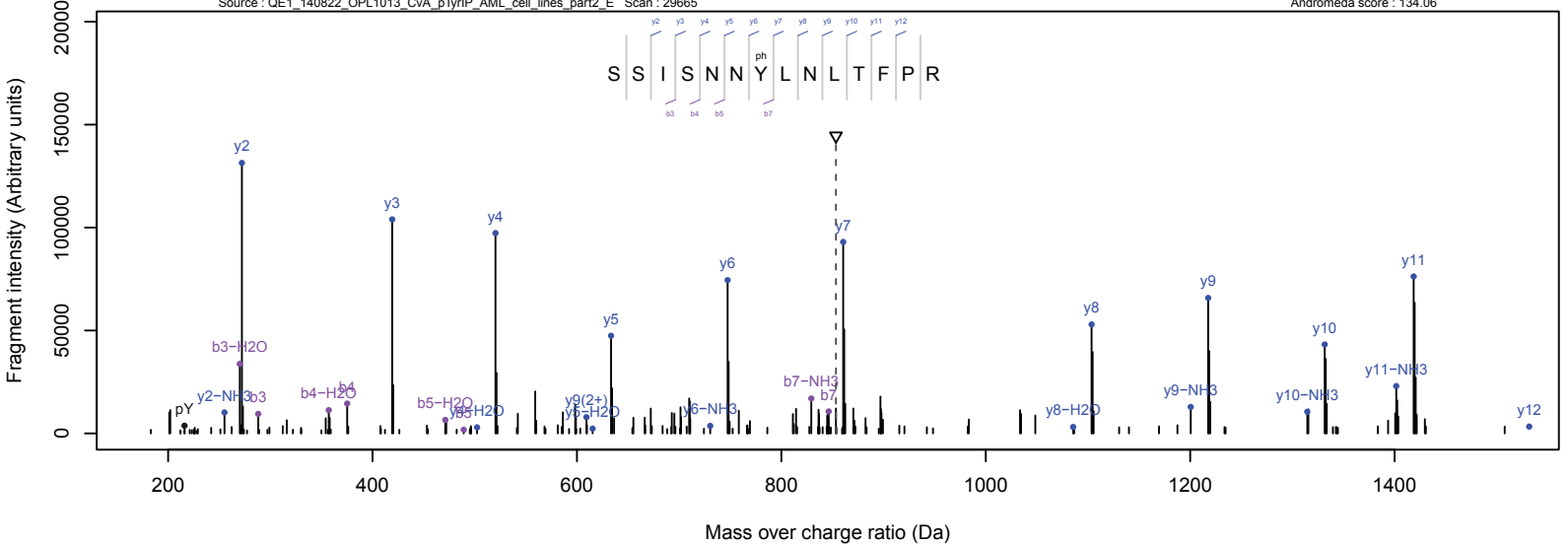
Andromeda score : 134.29



**SSISNNYLNLTFFPR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 29665

Andromeda score : 134.06

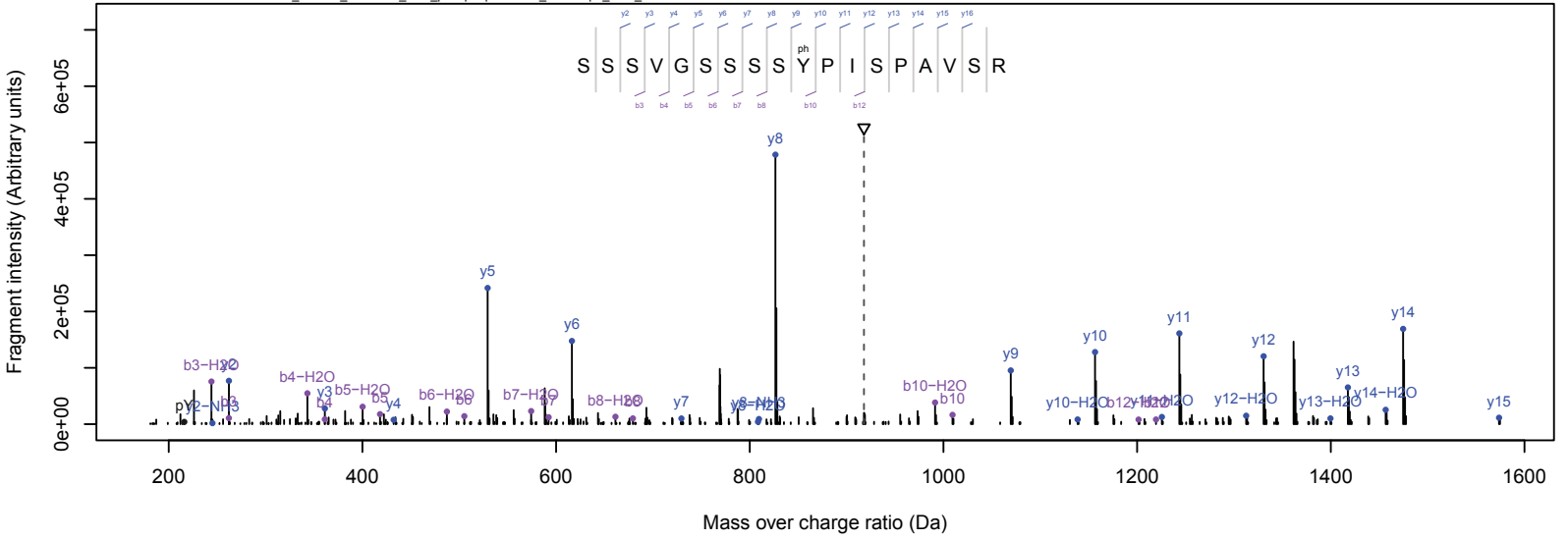




SSSVGSSSSYPISPAVSR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15590

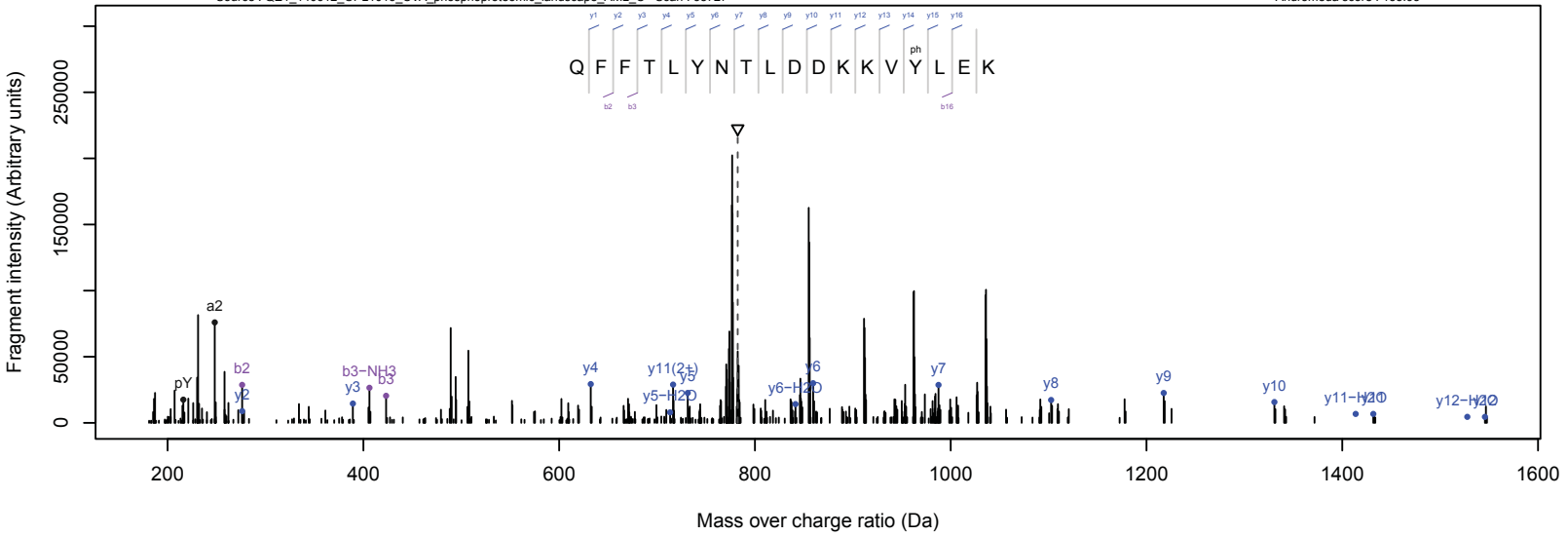
Andromeda score: 133.94



QFFTLYNTLDDKKVYLEK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38727

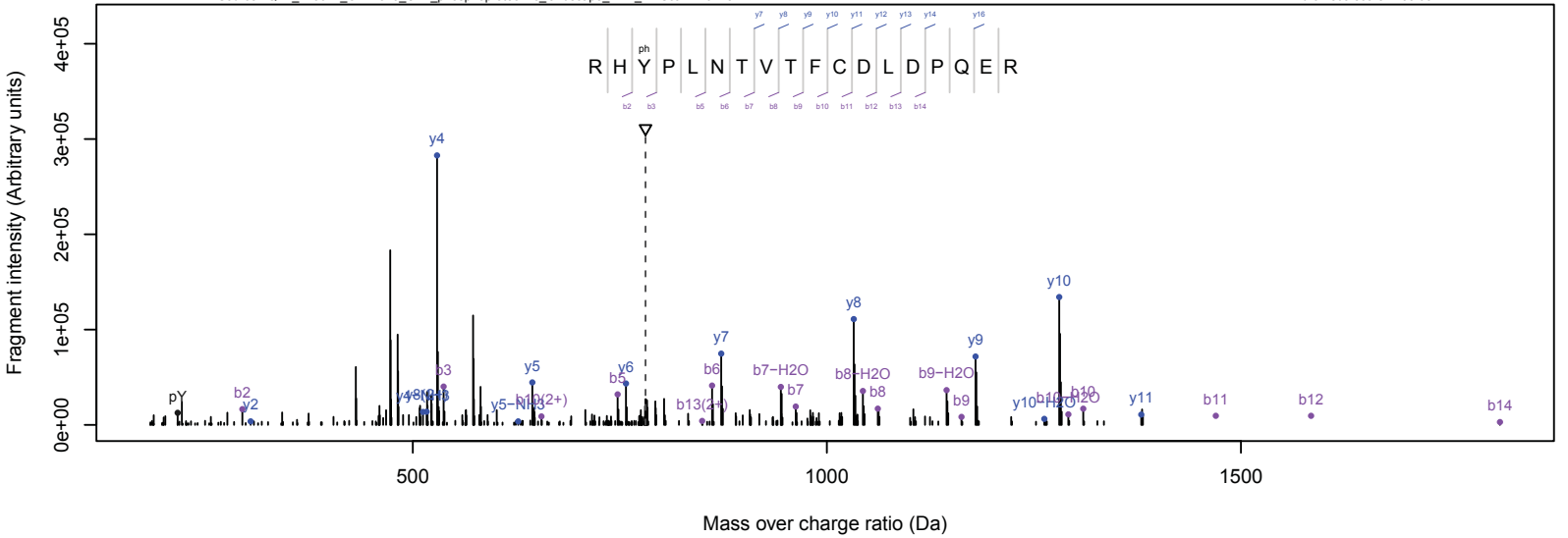
Andromeda score: 133.93



RHY(ph)PLNTVTFCDLPQER-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 29448

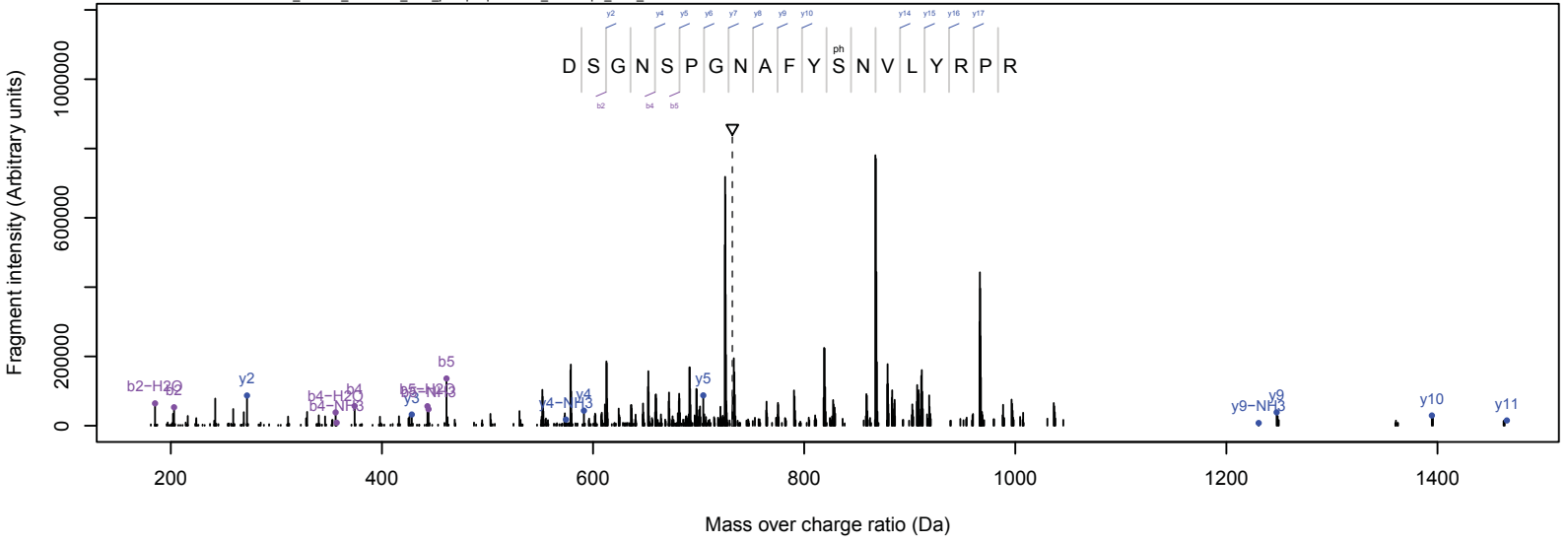
Andromeda score: 133.93



DSGNSPGNAFY(ph)NVLYRPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29540

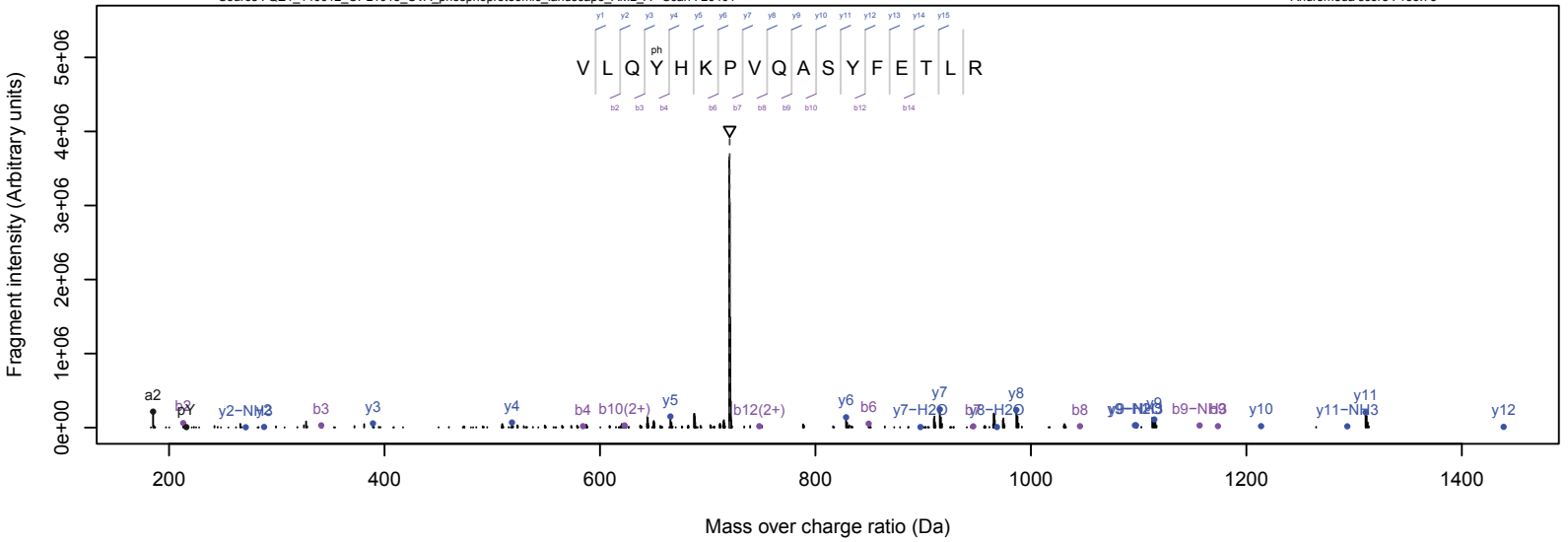
Andromeda score : 133.86



VLQY(ph)HKPVQASYFETLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26401

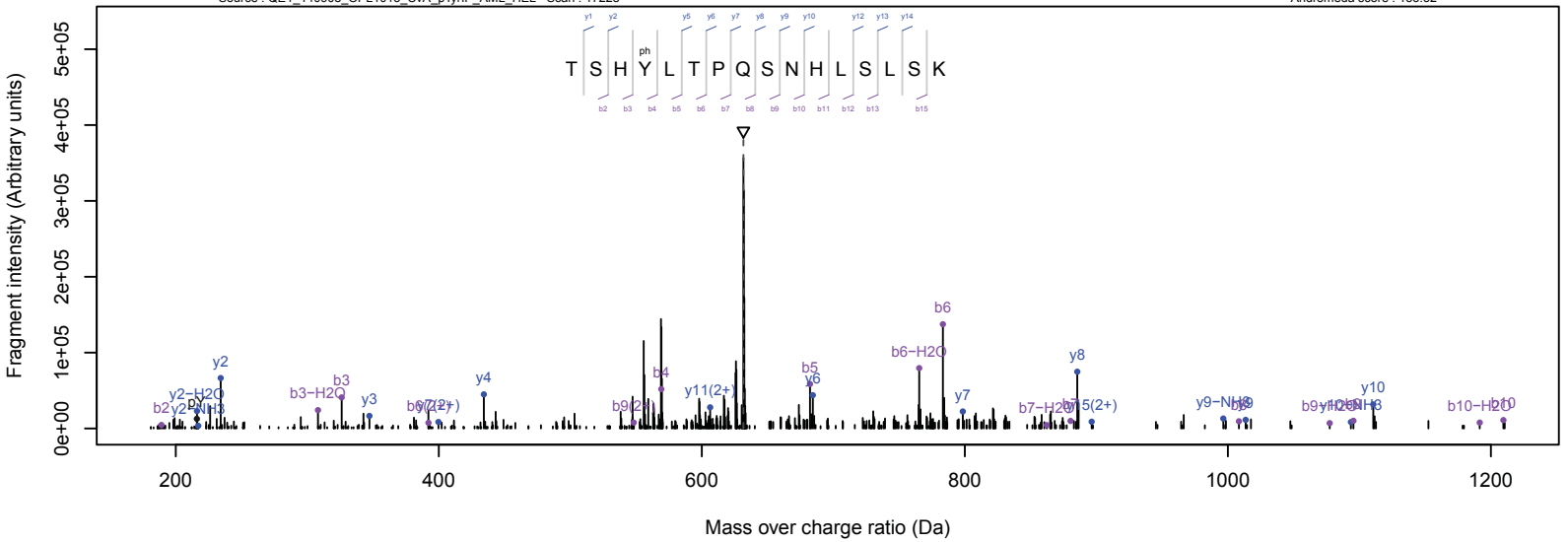
Andromeda score : 133.78



TSHY(ph)LTPQSNHLSLSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 17228

Andromeda score : 133.52

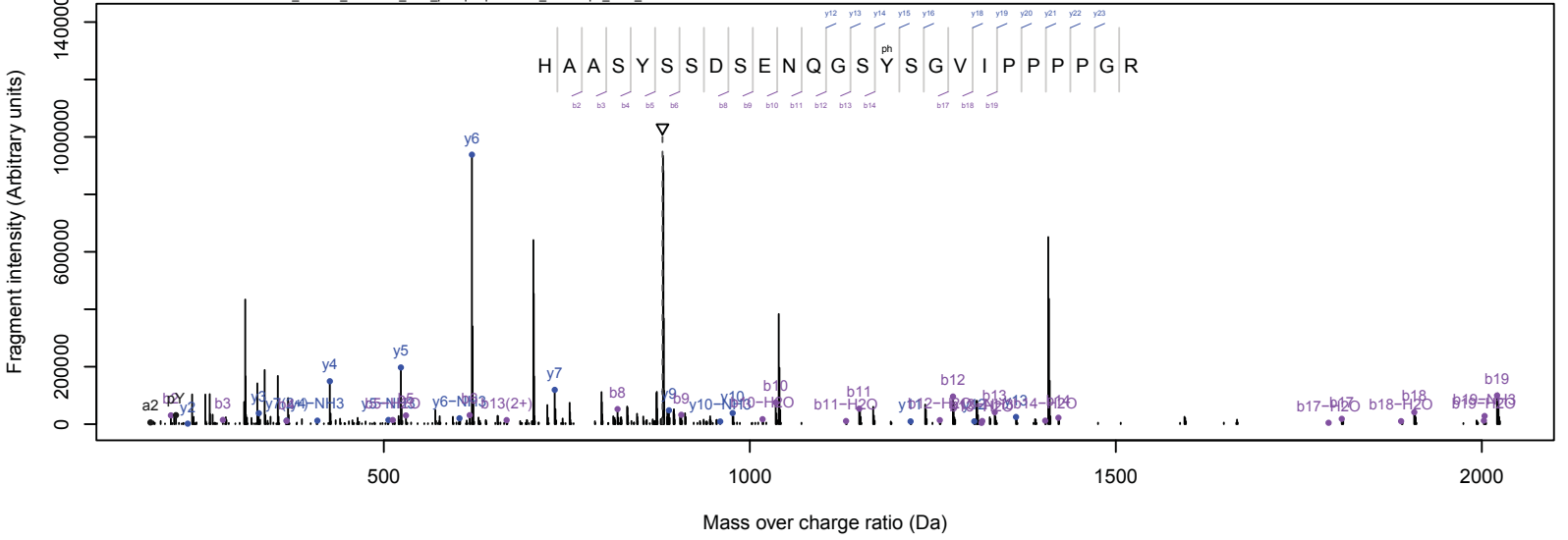




**HAASYSSDSENQGSYPHSGVIPPPGR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18872

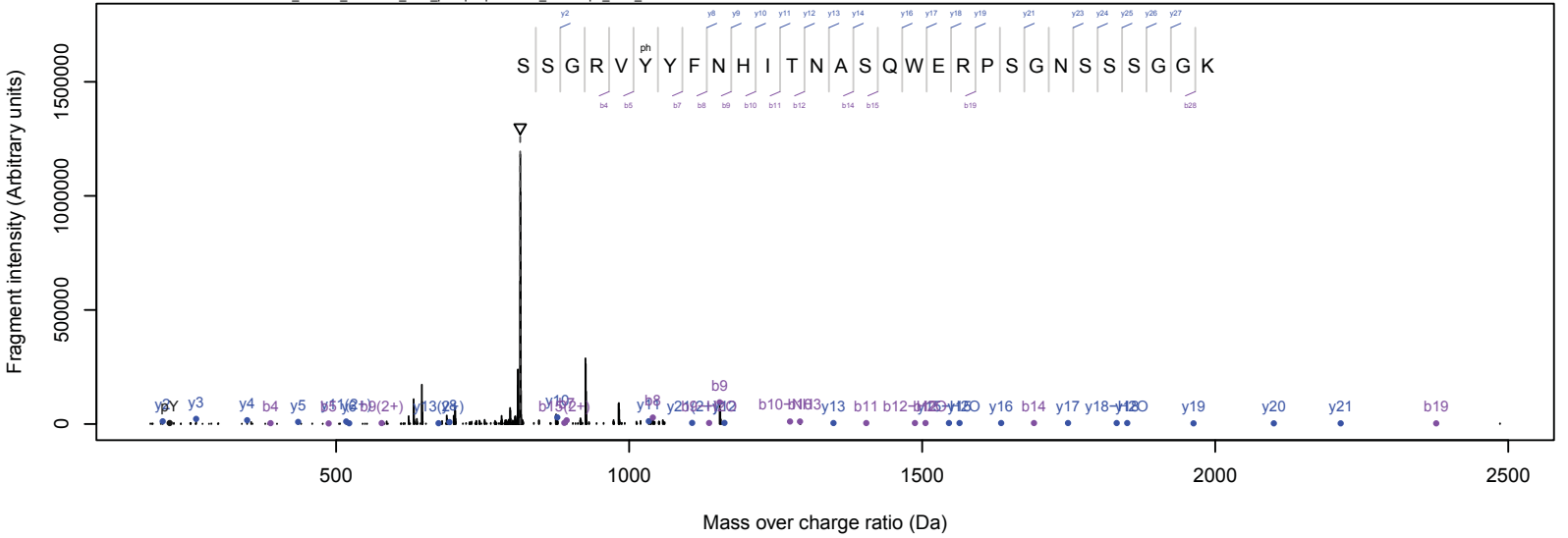
Andromeda score: 133.41



**SSGRVYPHFNHITNASQWERPSGNSSSSGGK-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25107

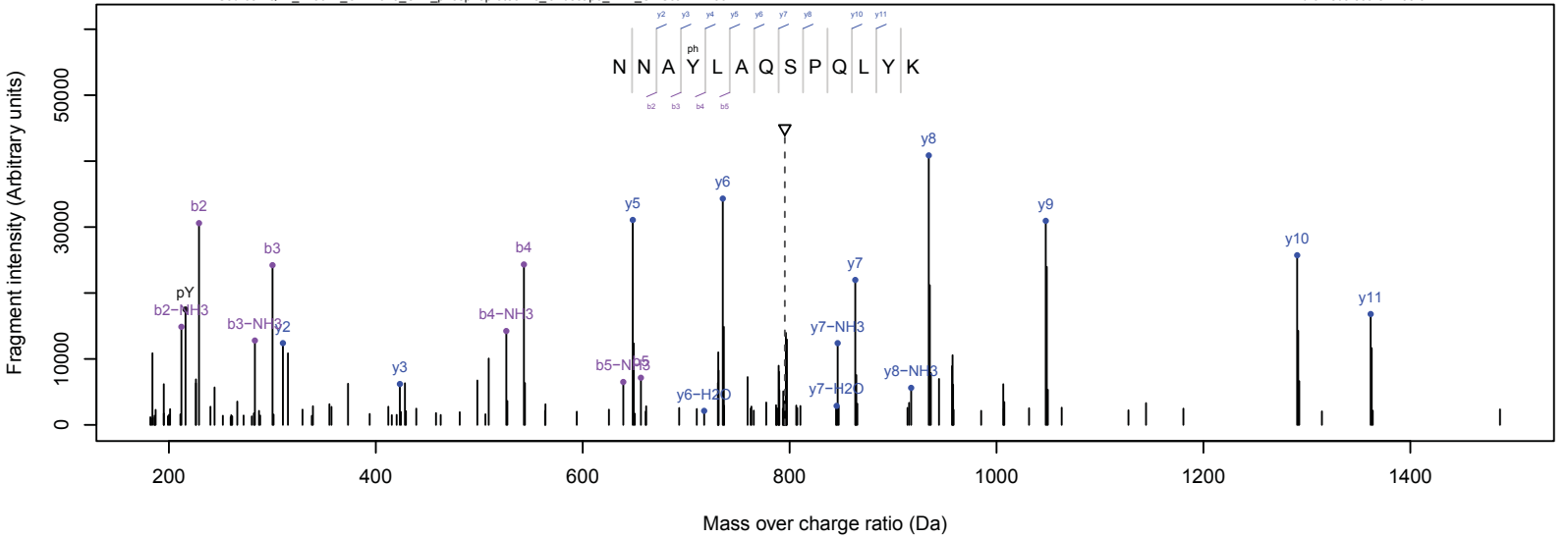
Andromeda score: 133.4



**NNAYLDAQSPQLYK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21564

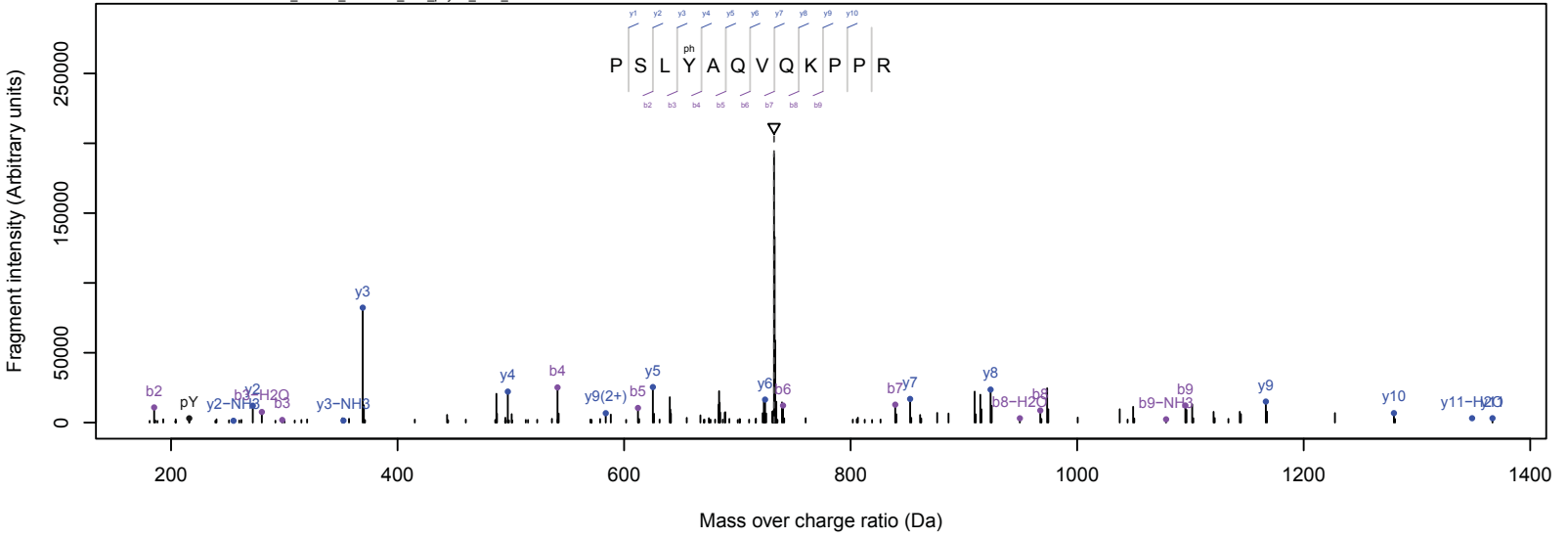
Andromeda score: 133.32



PSLY(ph)AQVQKPPR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 15065

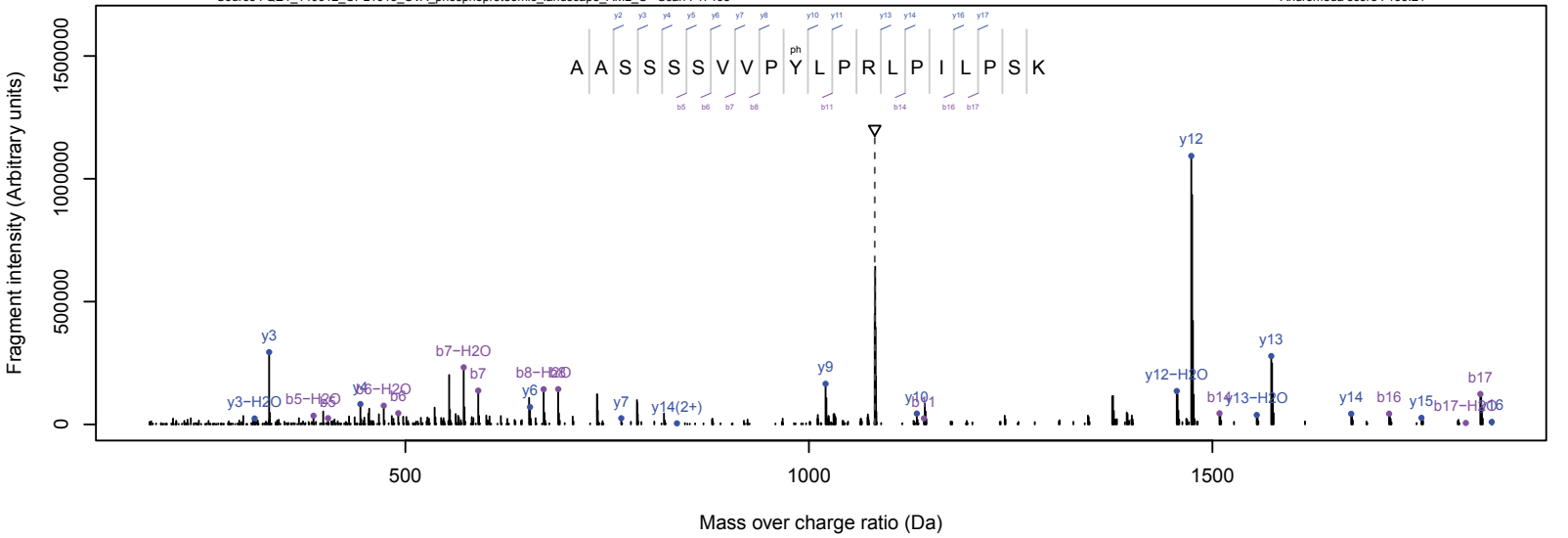
Andromeda score : 133.32



AASSSVVPY(ph)LPRLPILPSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47468

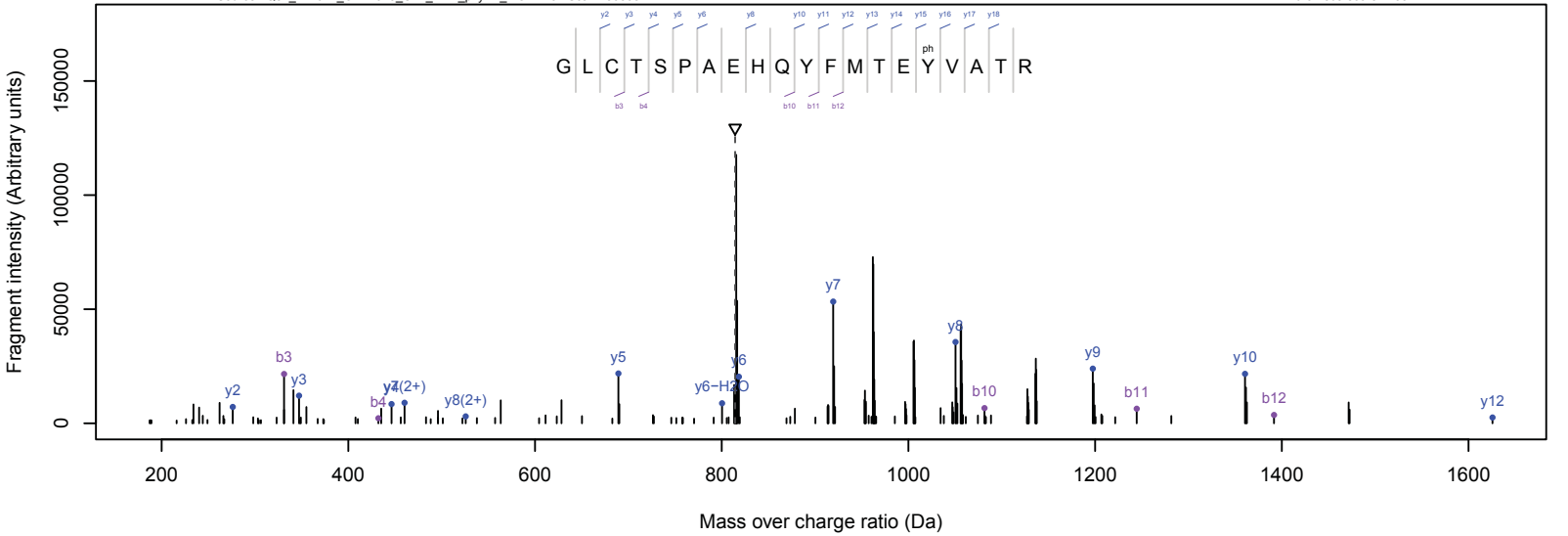
Andromeda score : 133.21



GLCTSPAEHQYFMTEY(ph)VATR-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 30808

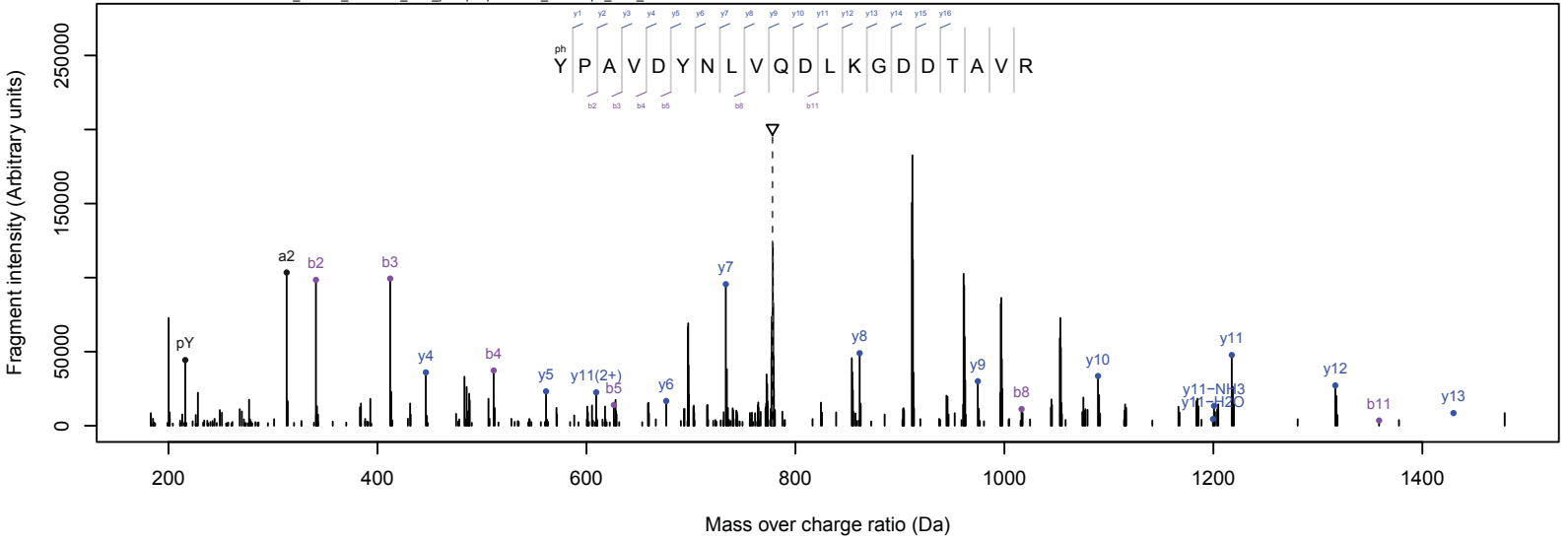
Andromeda score : 133.21



**Y(ph)PAVDYNLVQDLKGGDDTAVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37295

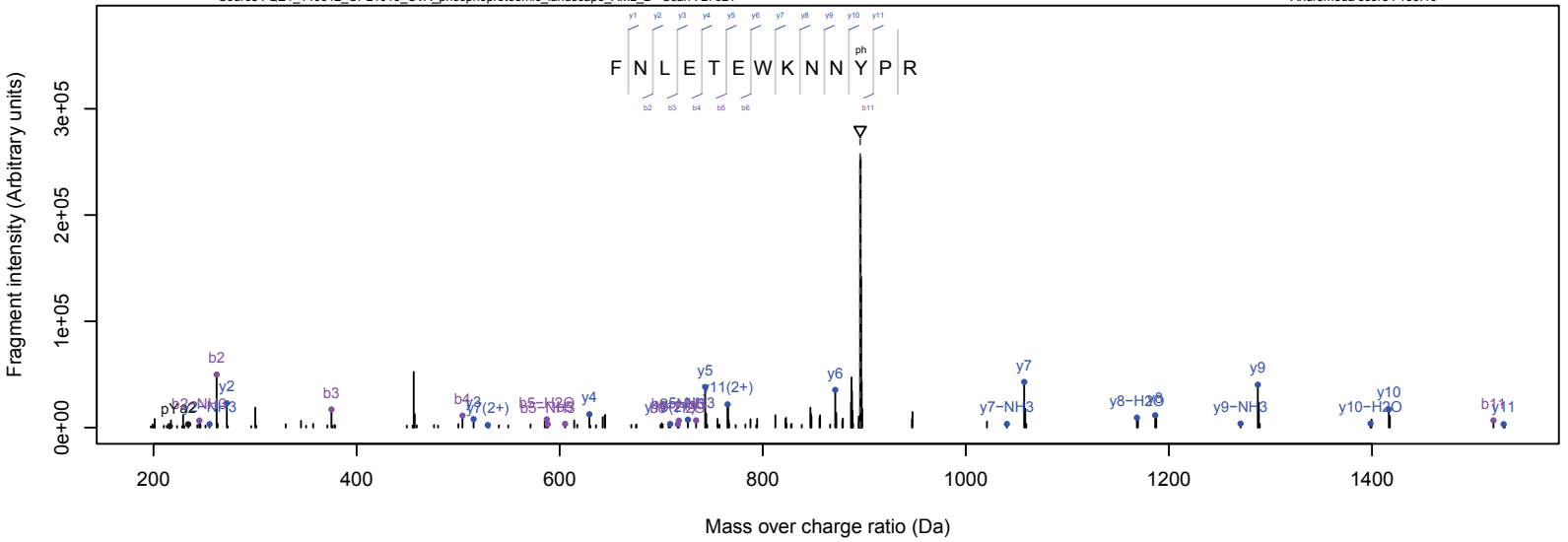
Andromeda score : 133.21



**FNLETEWKNNY(ph)PR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27621

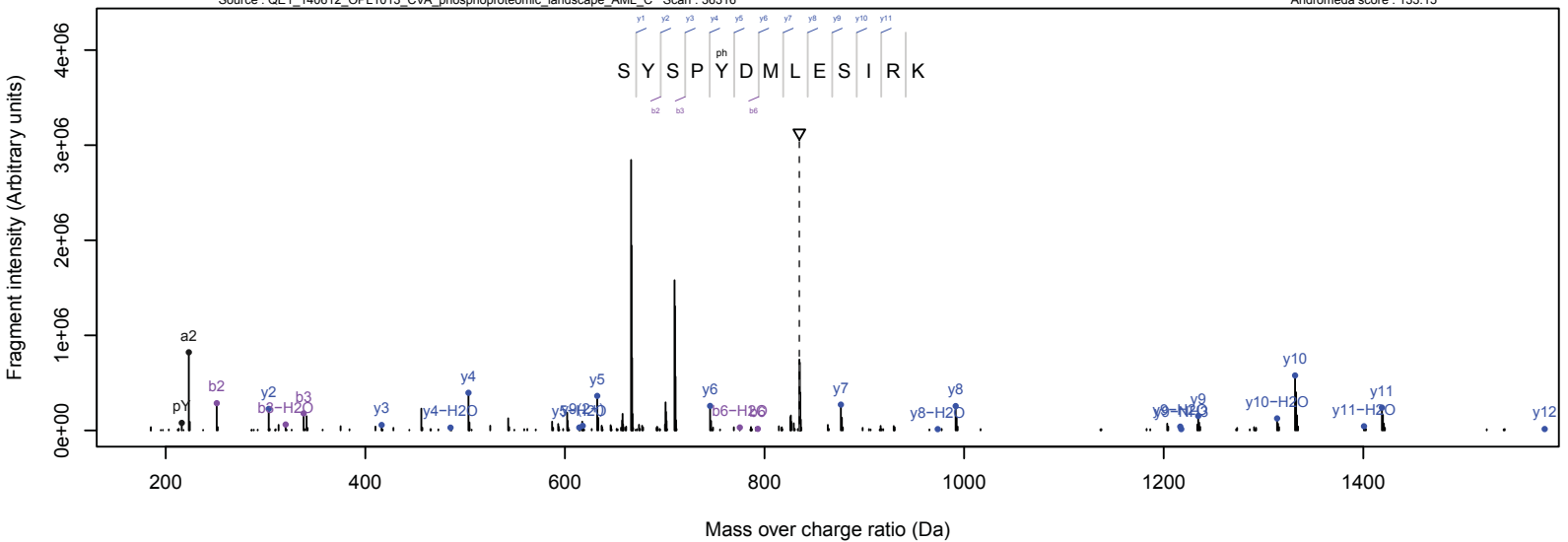
Andromeda score : 133.16



**SYSPY(ph)DMLESIRK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36316

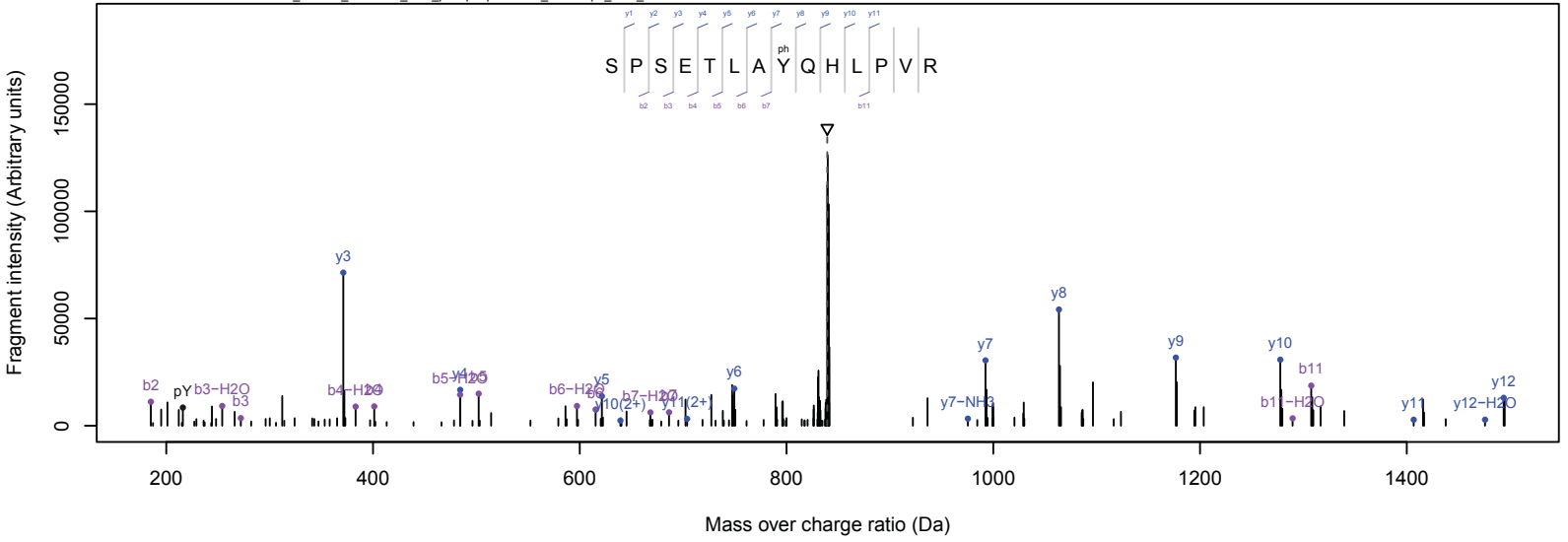
Andromeda score : 133.15



SPSETLAY(ph)QHL PVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22023

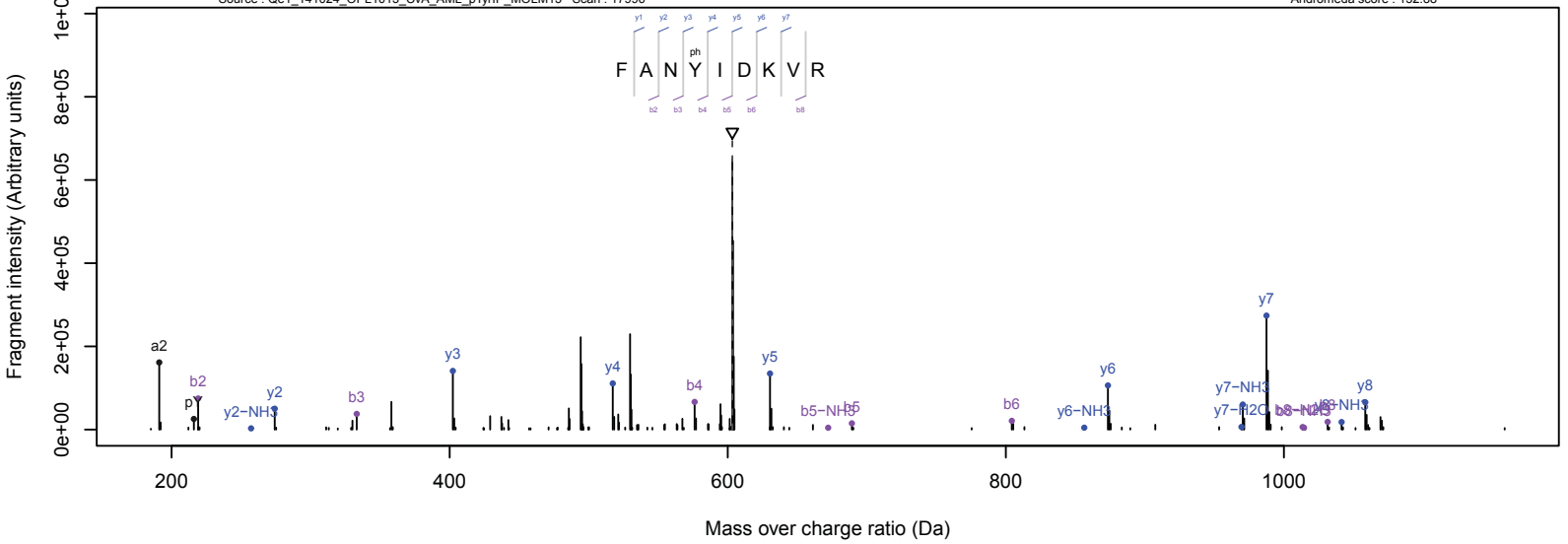
Andromeda score : 132.97



FANY(ph)IDKVR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 17996

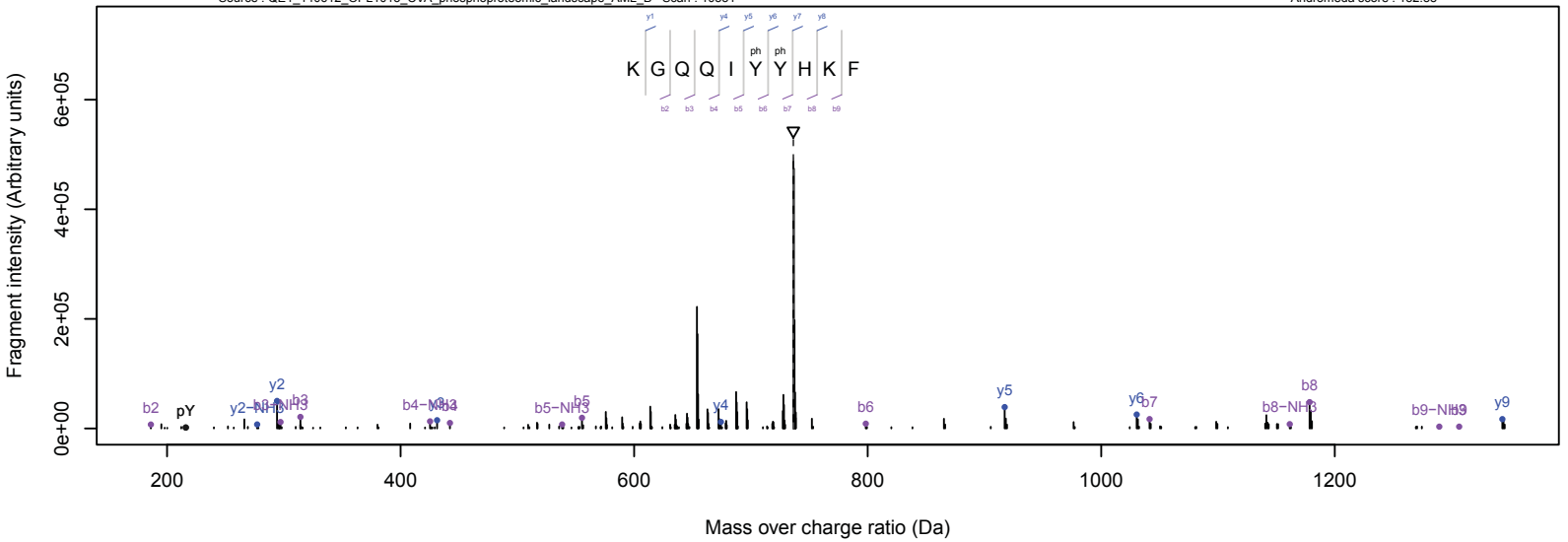
Andromeda score : 132.88



KGQQIY(ph)Y(ph)HKF-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10531

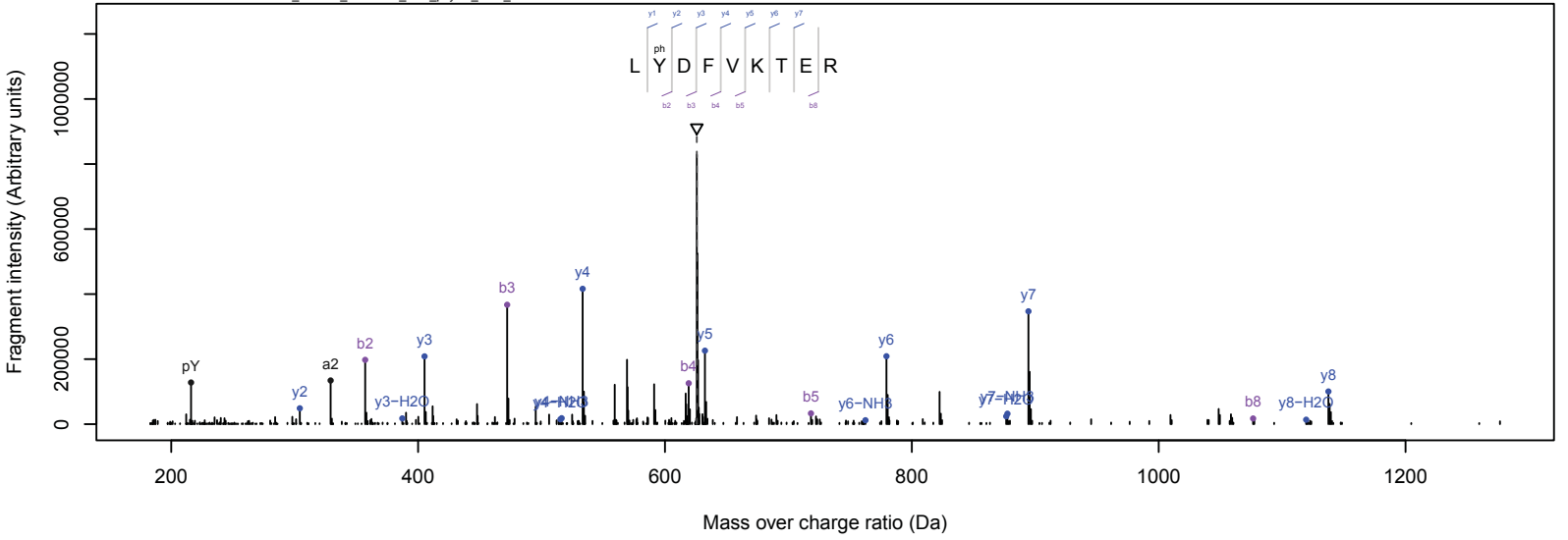
Andromeda score : 132.88



LY(ph)DFVKTER-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 18330

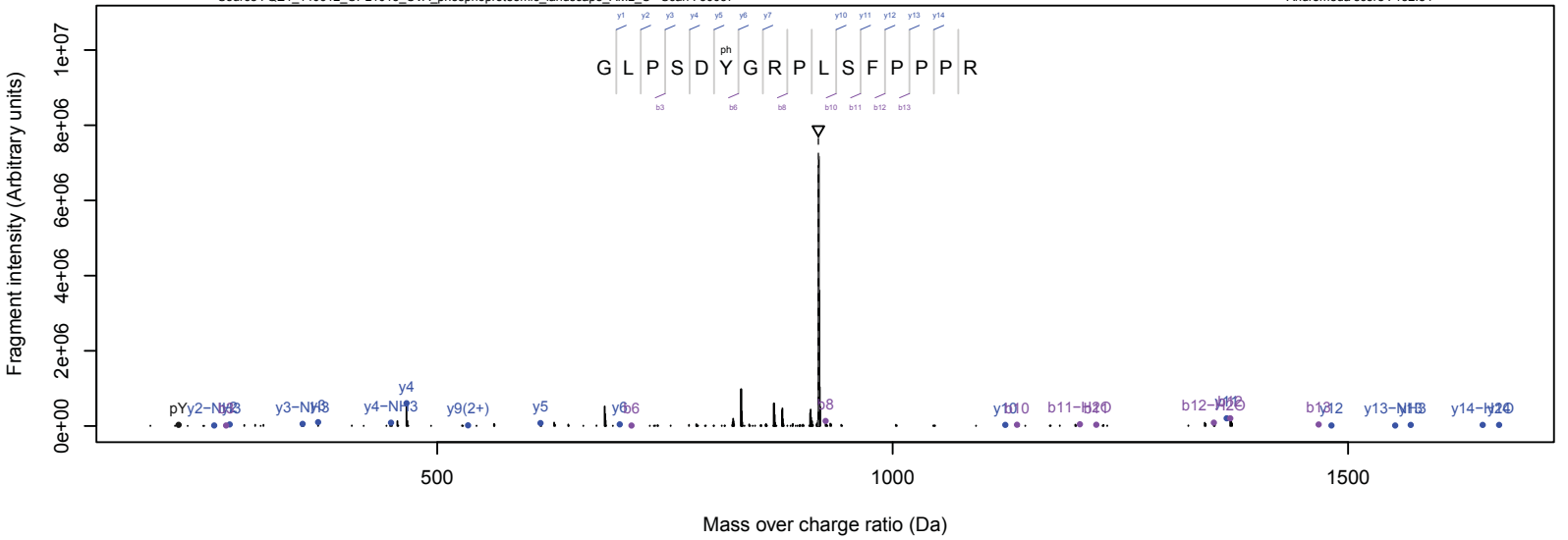
Andromeda score : 132.88



GLPSDY(ph)GRPLSFPPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30037

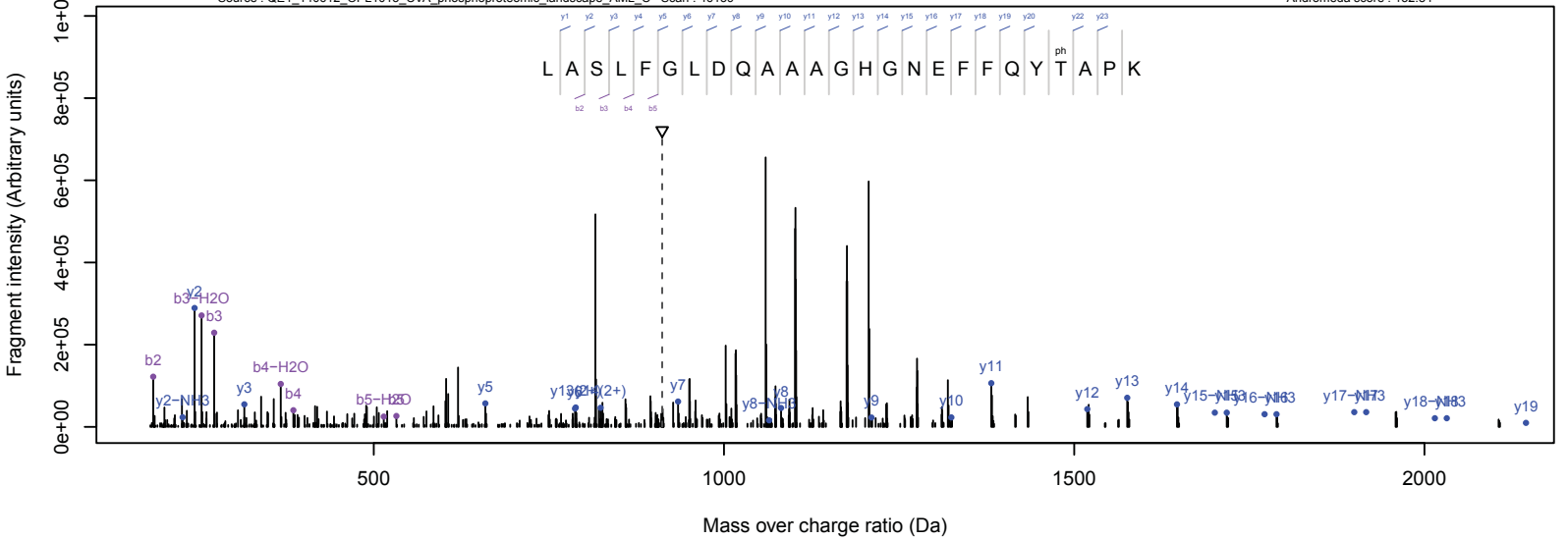
Andromeda score : 132.84



LASLFGLDQAAAGHGNEFFQY(ph)APK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46139

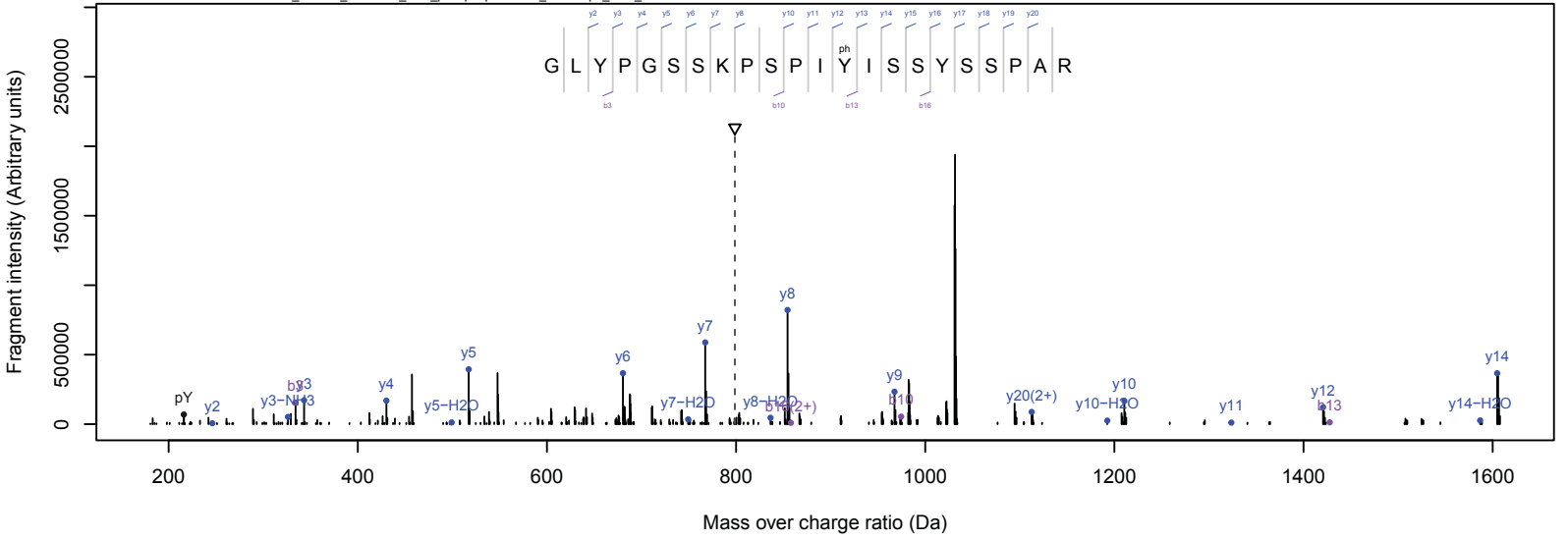
Andromeda score : 132.84



GLYPGSSKPSPIYISSYSSPAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31606

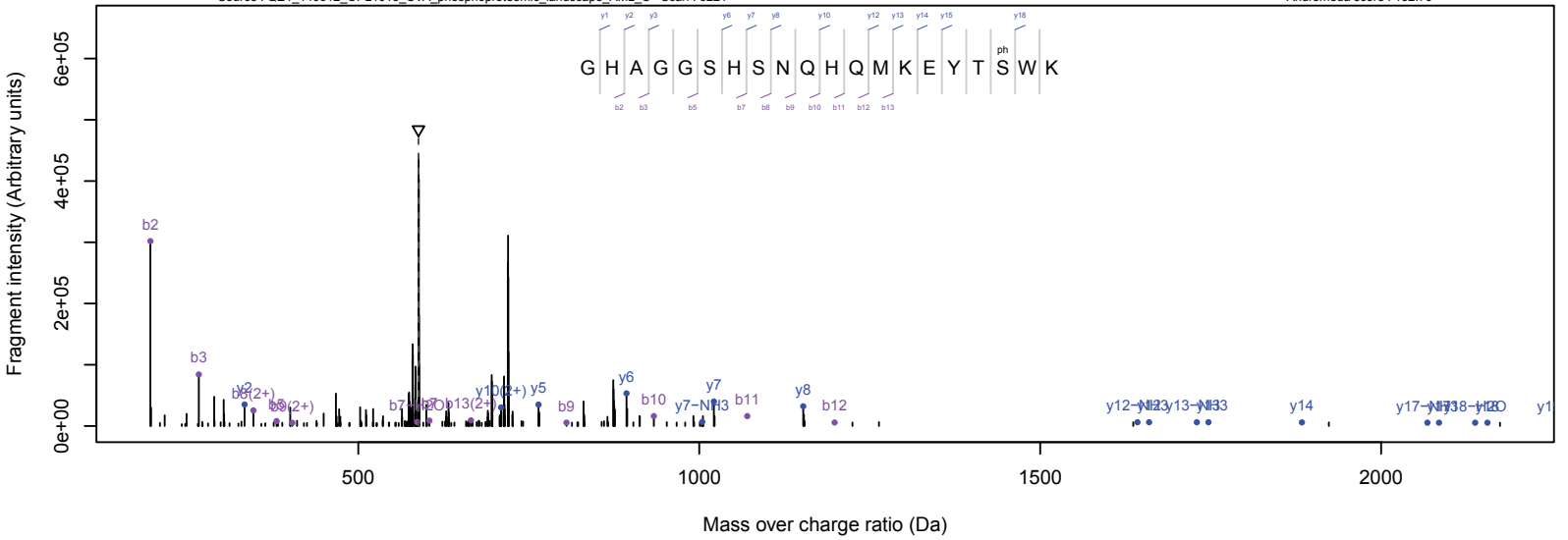
Andromeda score: 132.83



GHAGGSHSNQHMQKEYTSW K-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 6221

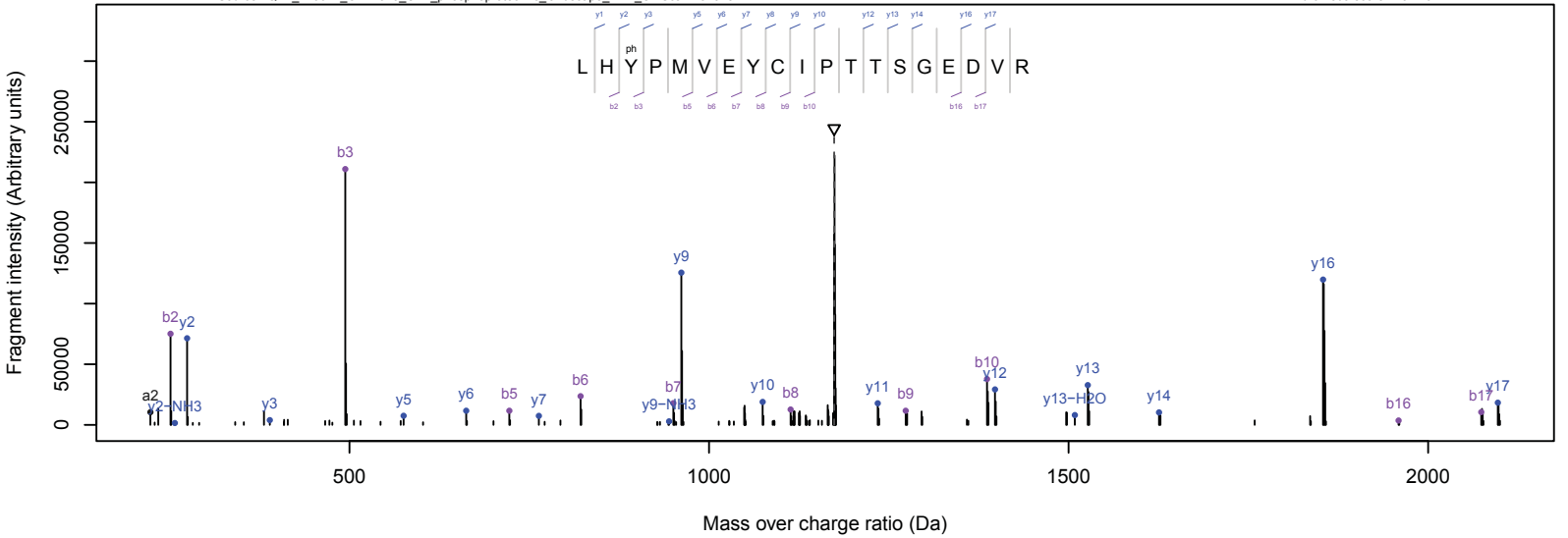
Andromeda score: 132.79



LHY(ph)PMVEYCIPTTSGEDVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 32548

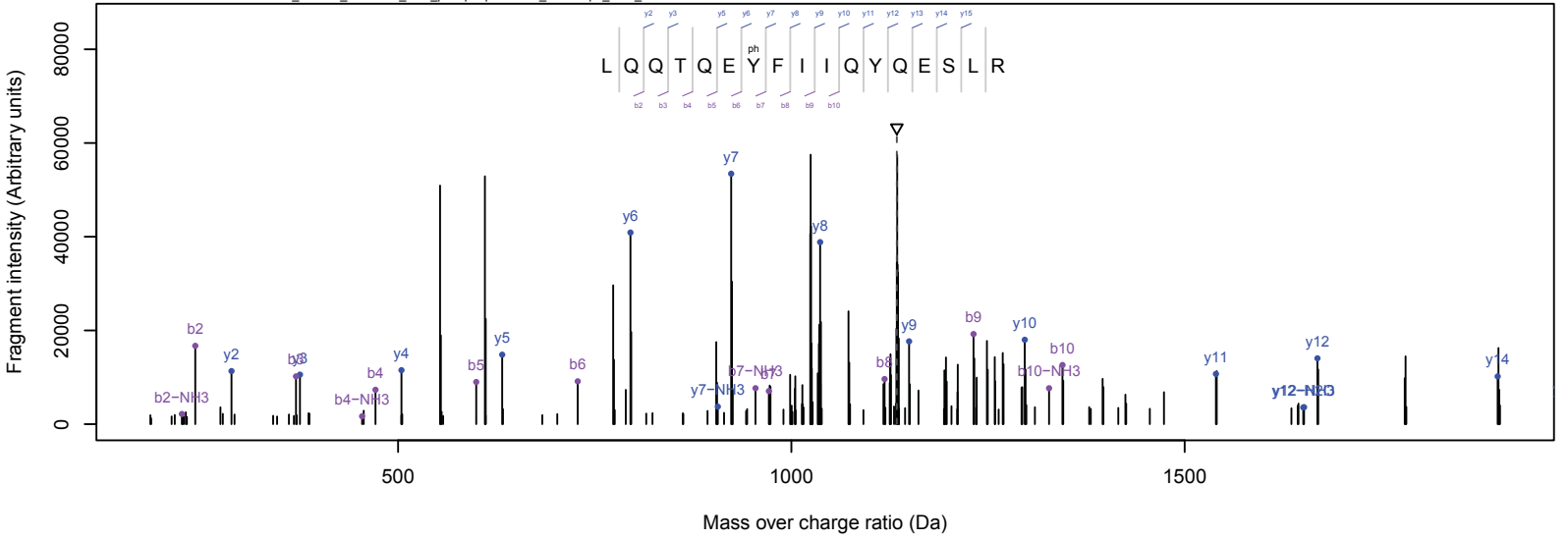
Andromeda score: 132.79



LQQTQEY(ph)FIQYQESLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 42066

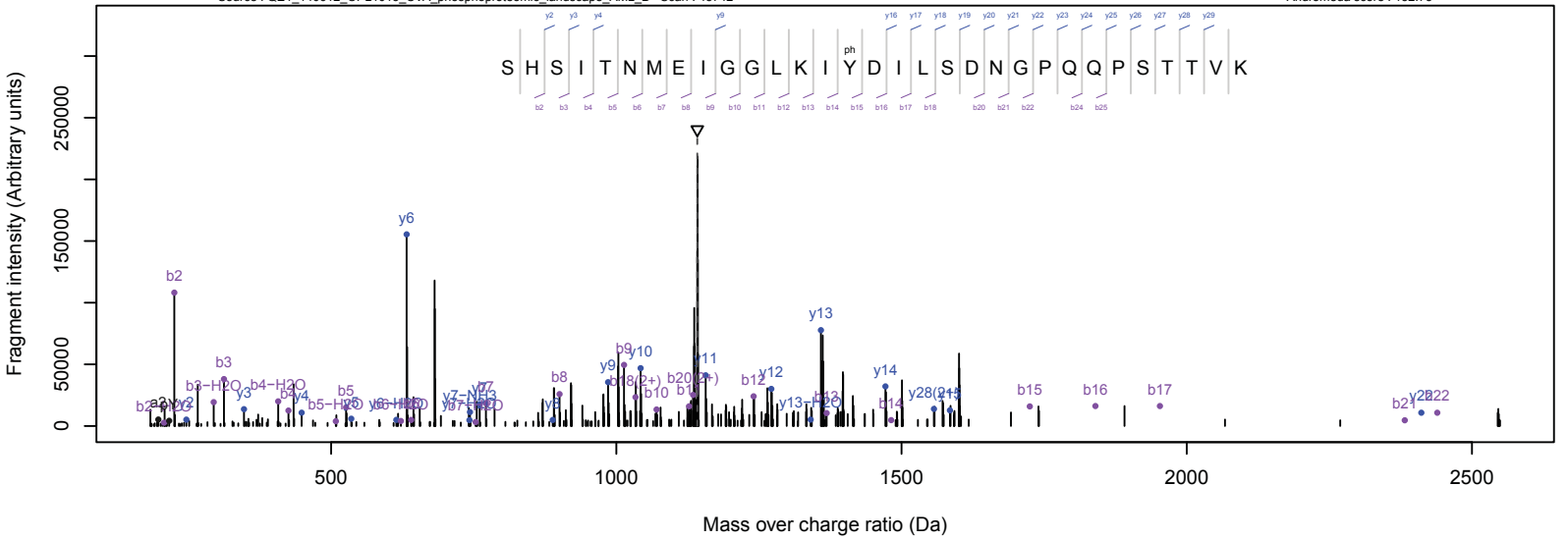
Andromeda score : 132.79



SHSITNMEIGGLKIY(ph)DILSDNGPQQPSTTVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45742

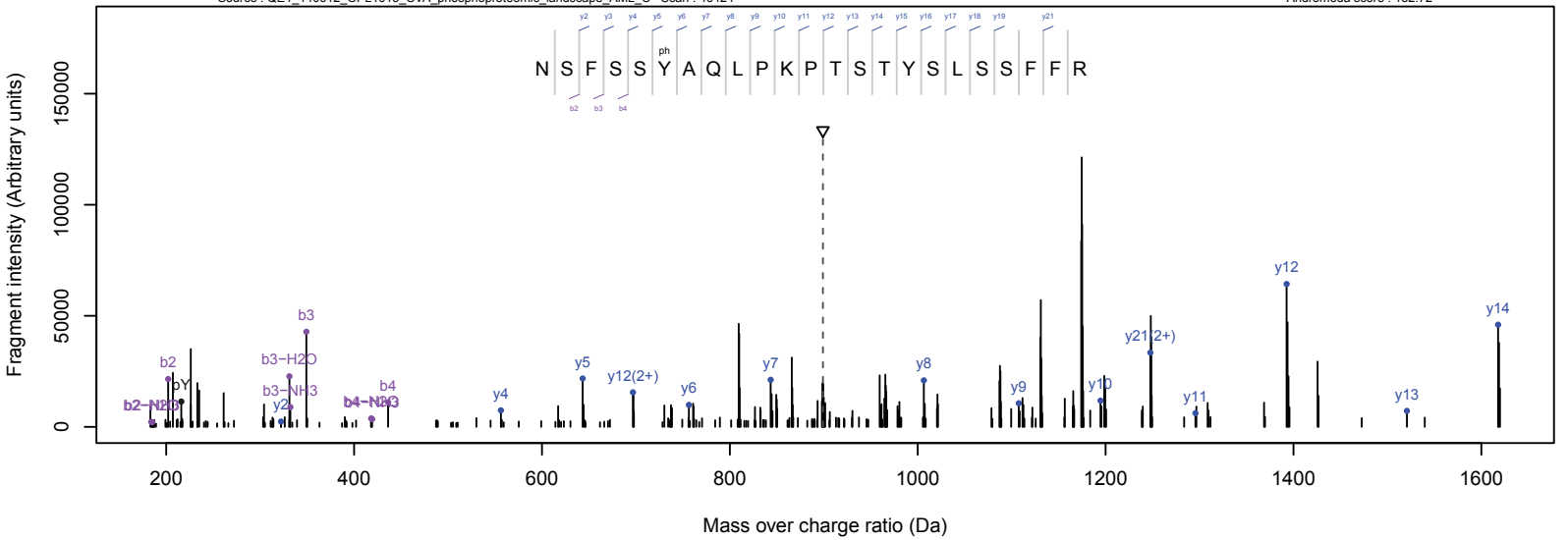
Andromeda score : 132.73



NSFSSY(ph)AQLPKPTSTYSLSSFFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46424

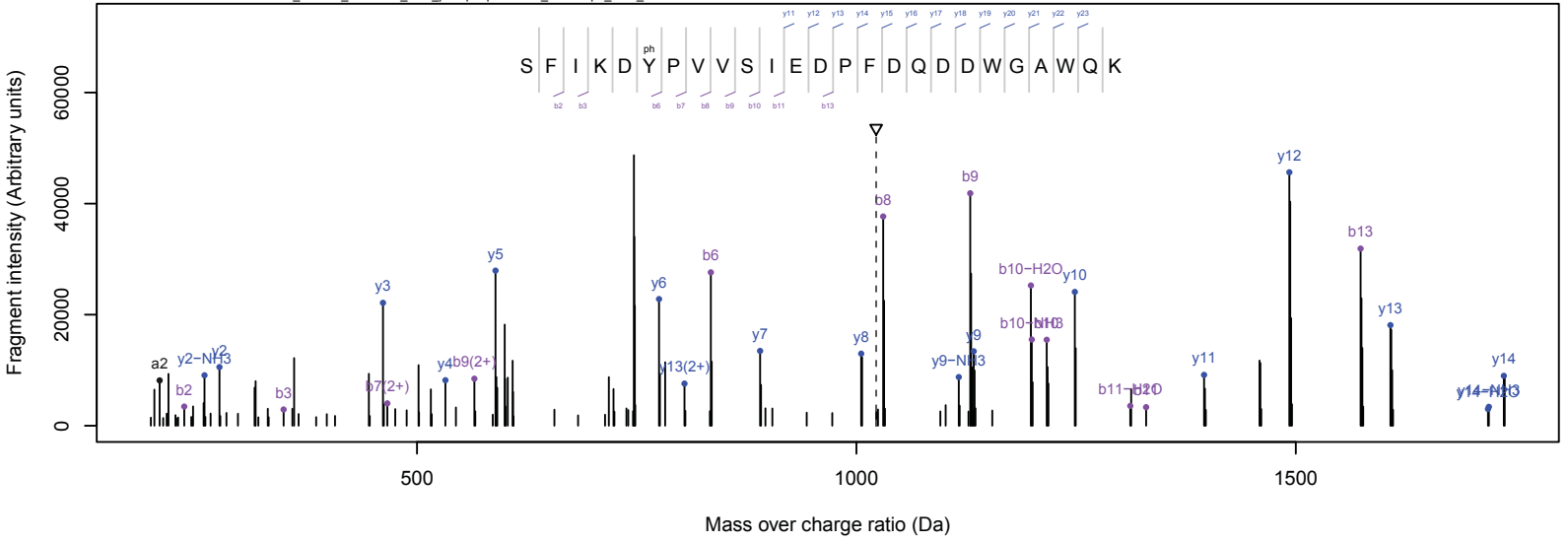
Andromeda score : 132.72



SFIKDY(ph)PVVSIEDPFDQDDWGAWQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 48052

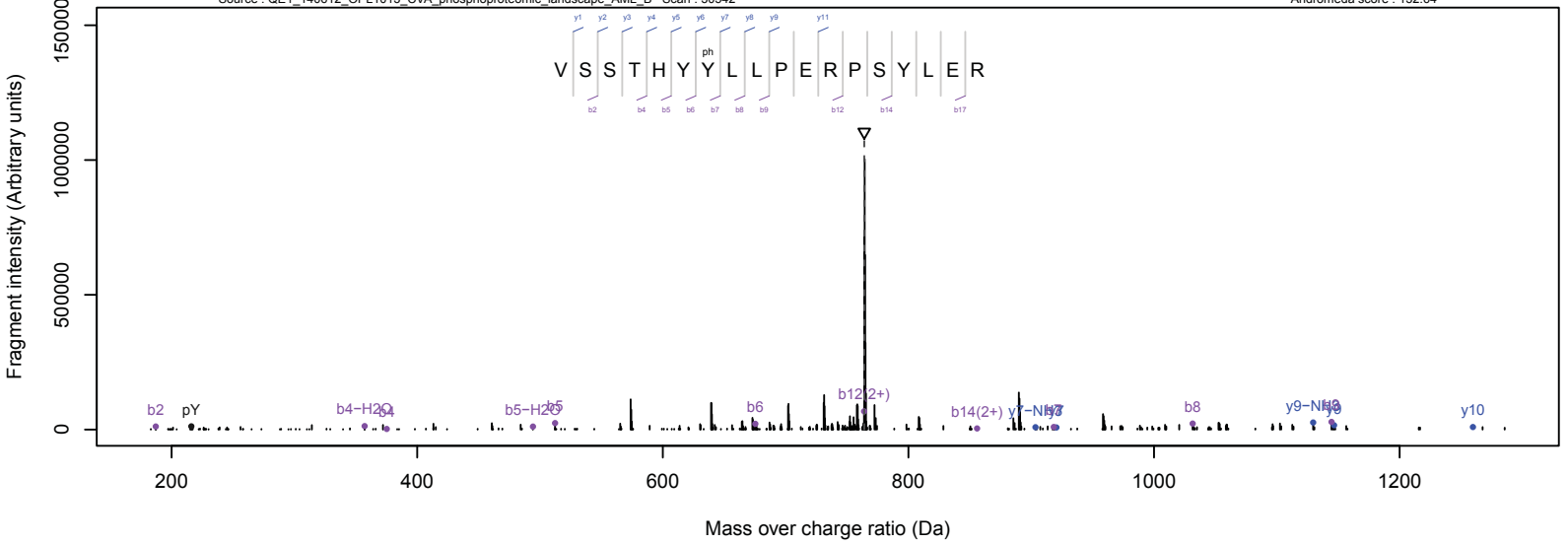
Andromeda score : 132.72



VSSTHY(ph)LLPERPSYLER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30542

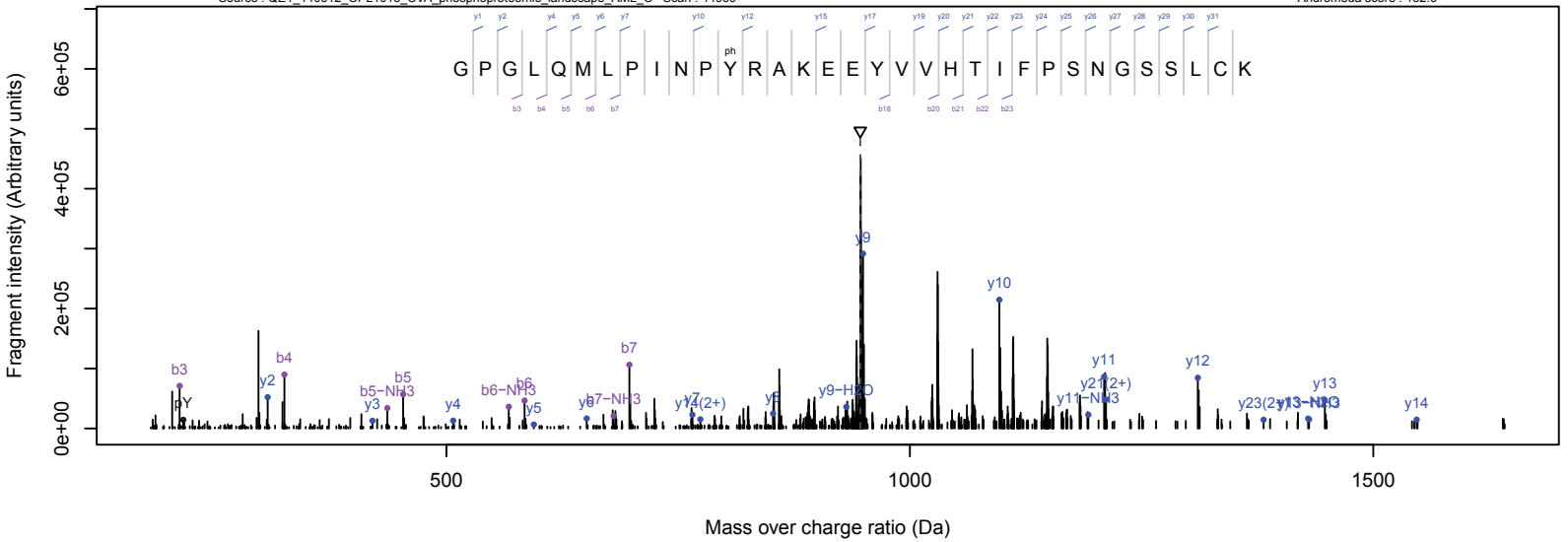
Andromeda score : 132.64



GPGLQMLPINPY(ph)RAKEEYVHTIFPSNGSSLCK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44939

Andromeda score : 132.6

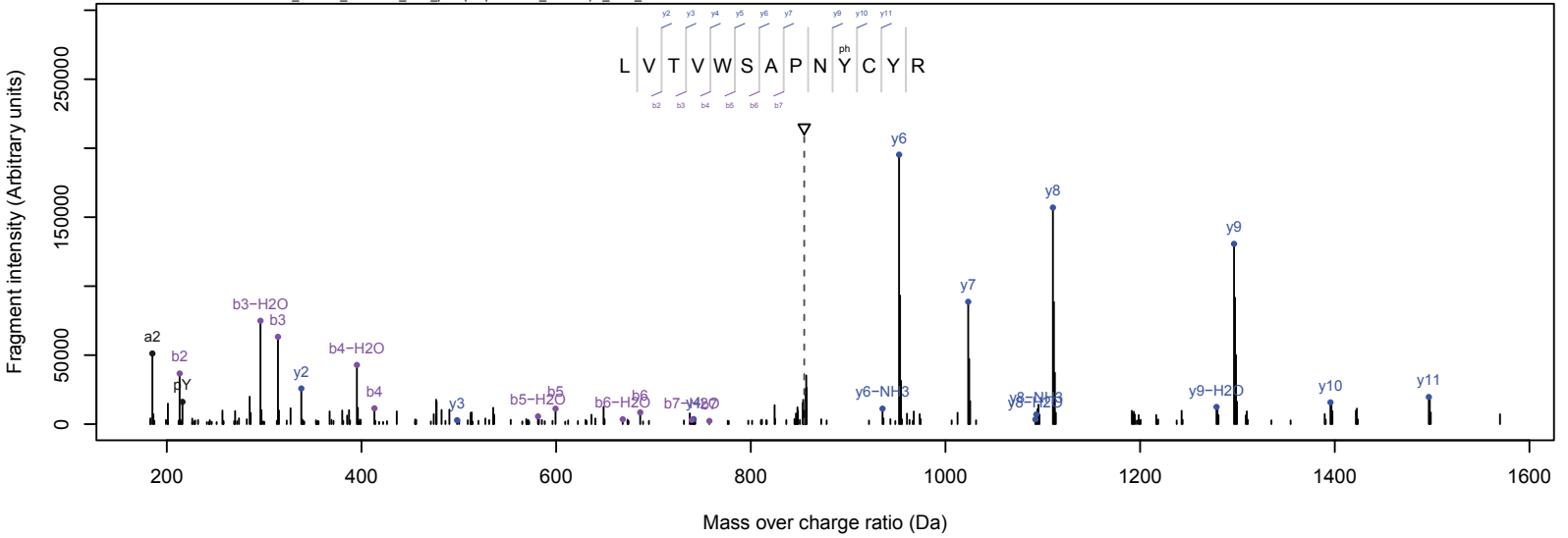




LVTVWSAPNY(ph)CYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32929

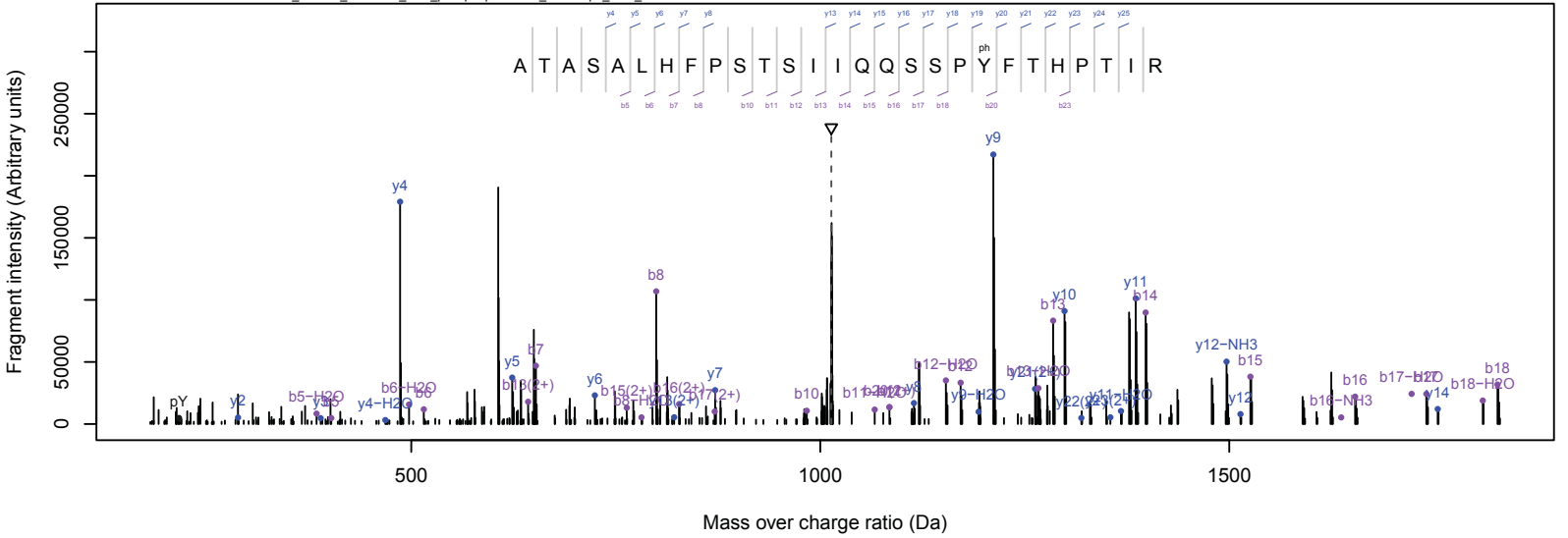
Andromeda score : 132.5



ATASALHFPSTSIQQSSPY(ph)FTHPTIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39360

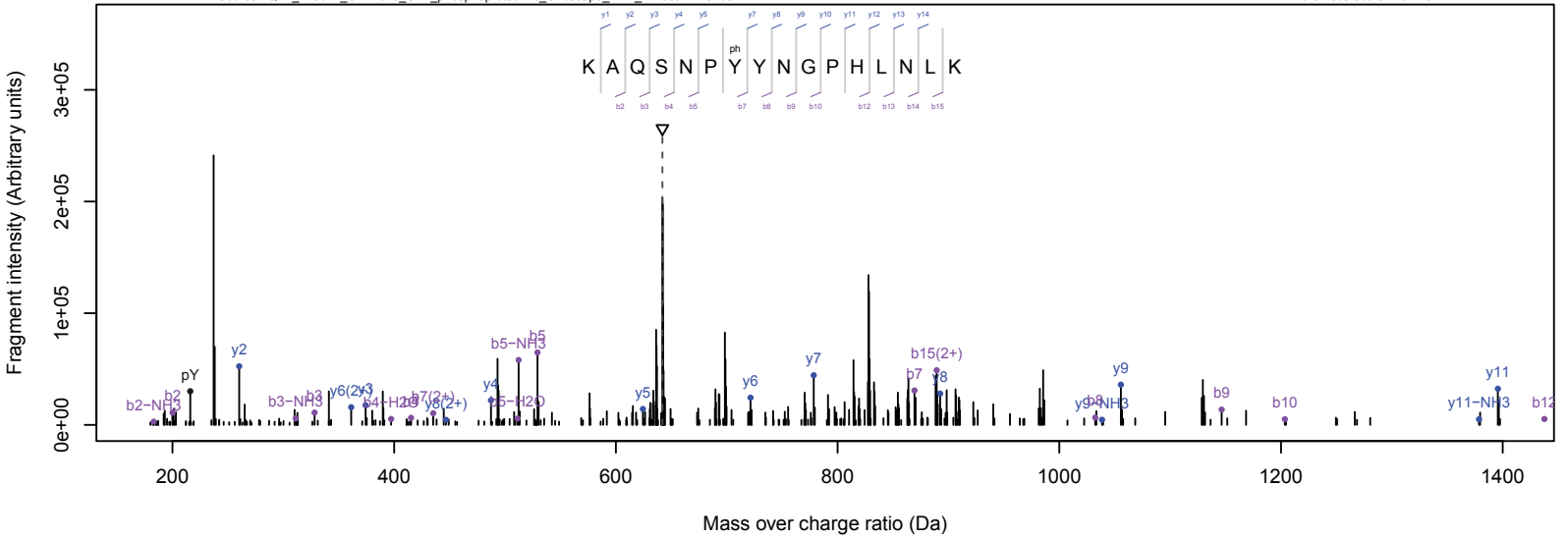
Andromeda score : 132.28



KAQSNPY(ph)YNGPHLNLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 18265

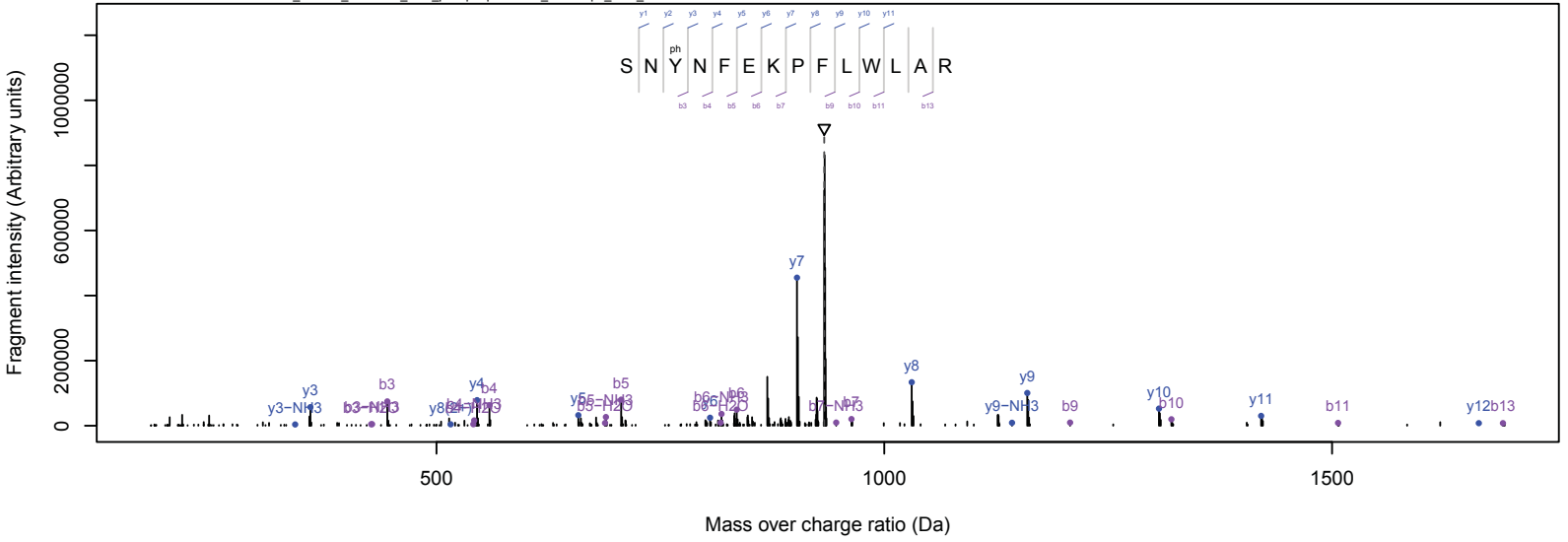
Andromeda score : 132.25



**SNY(ph)NFEKPLWLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45323

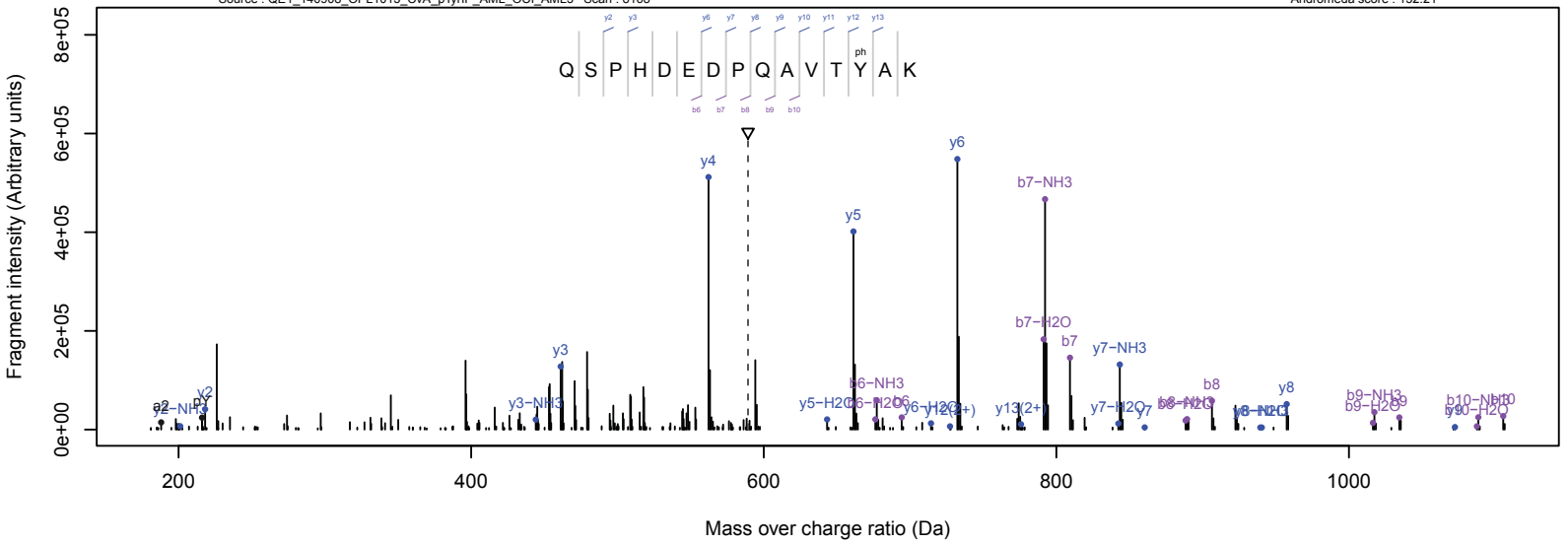
Andromeda score : 132.23



**QSPHDEDPAVTY(ph)AK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 8168

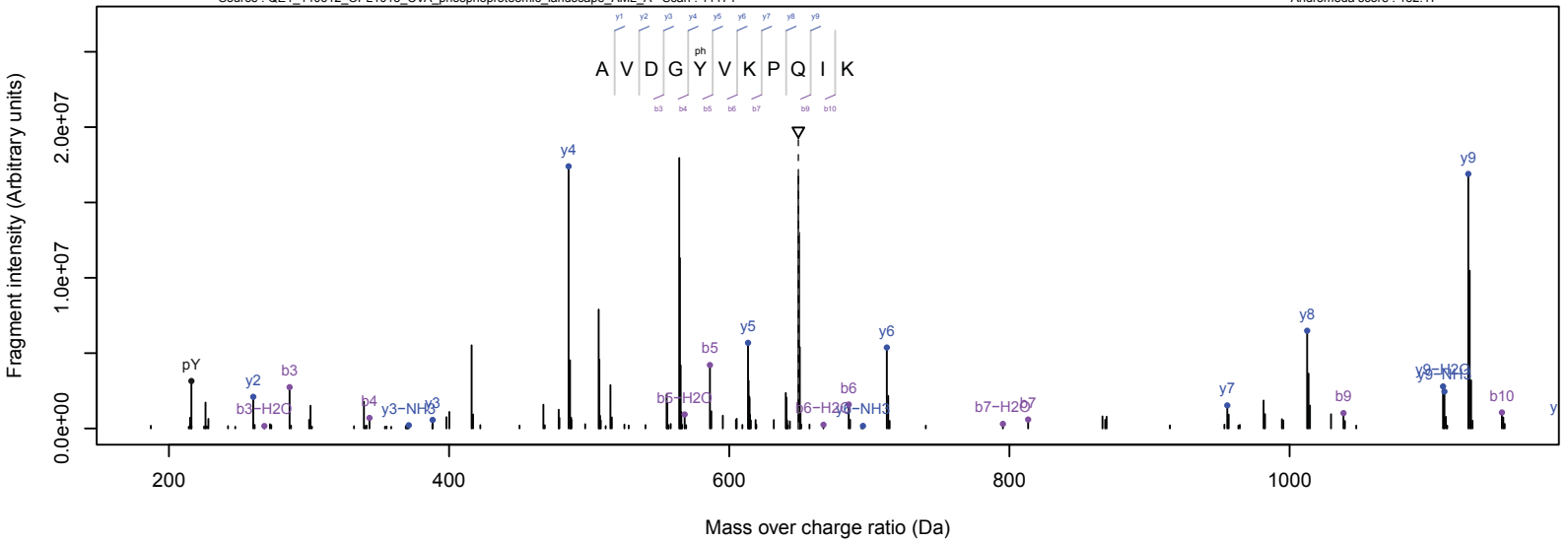
Andromeda score : 132.21



**AVDGY(ph)VKPQIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 11174

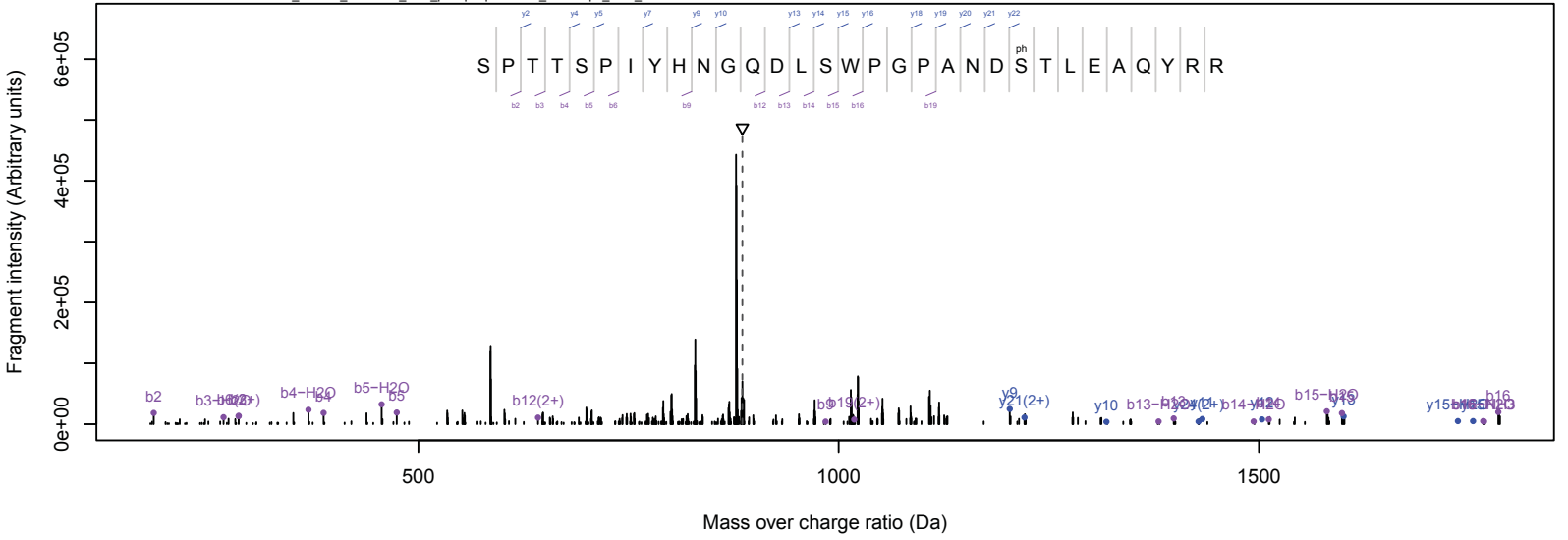
Andromeda score : 132.17



SPTTSPHYHNGQLSWPGPANDS(ph)TLEAQYRR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31244

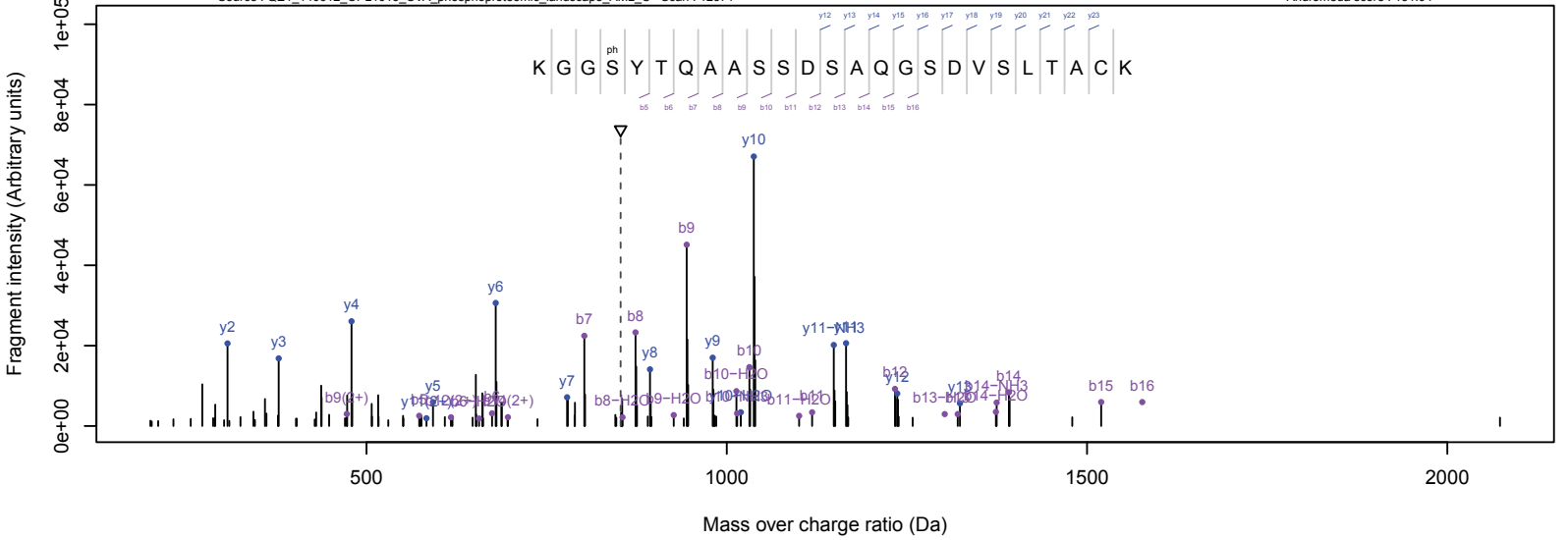
Andromeda score: 131.95



KGGS(ph)YTQAASSDSAQGS DVS LTACK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12371

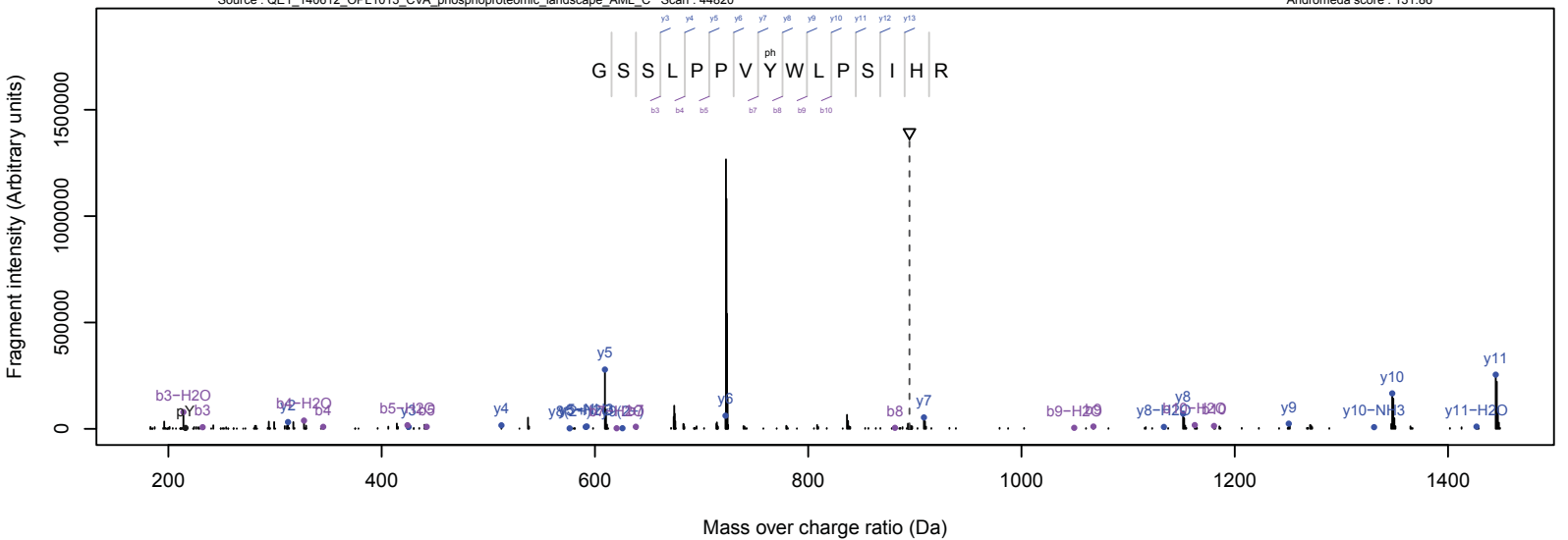
Andromeda score: 131.94



GSSLPPVY(ph)WLPSIHR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 44820

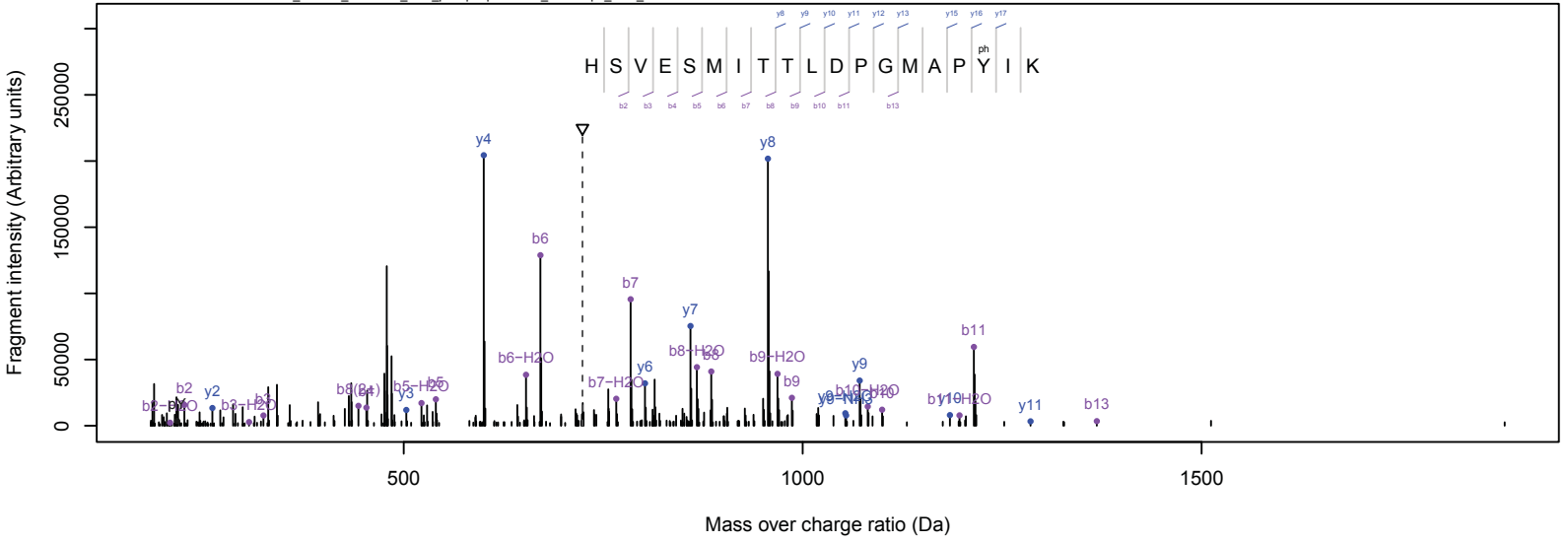
Andromeda score: 131.86



HSVESMITTLDPGMAPY(ph)K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 42246

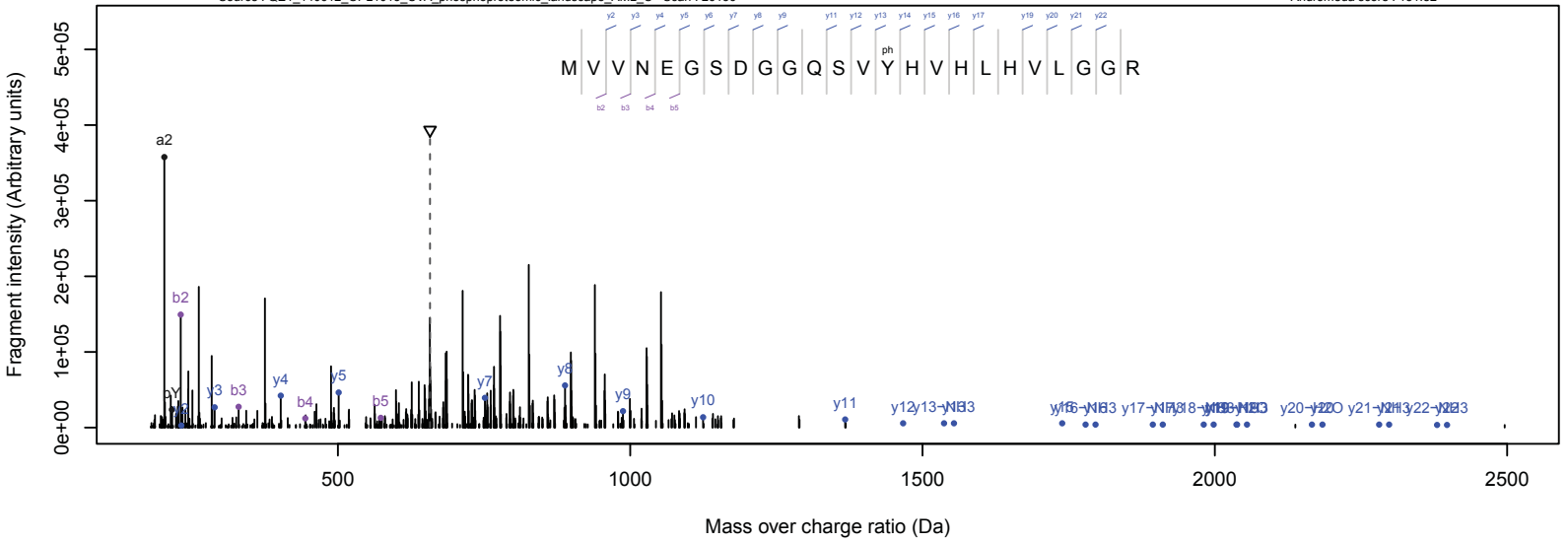
Andromeda score : 131.82



MVNEGSDGGQSVY(ph)HVLHVLGGR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26136

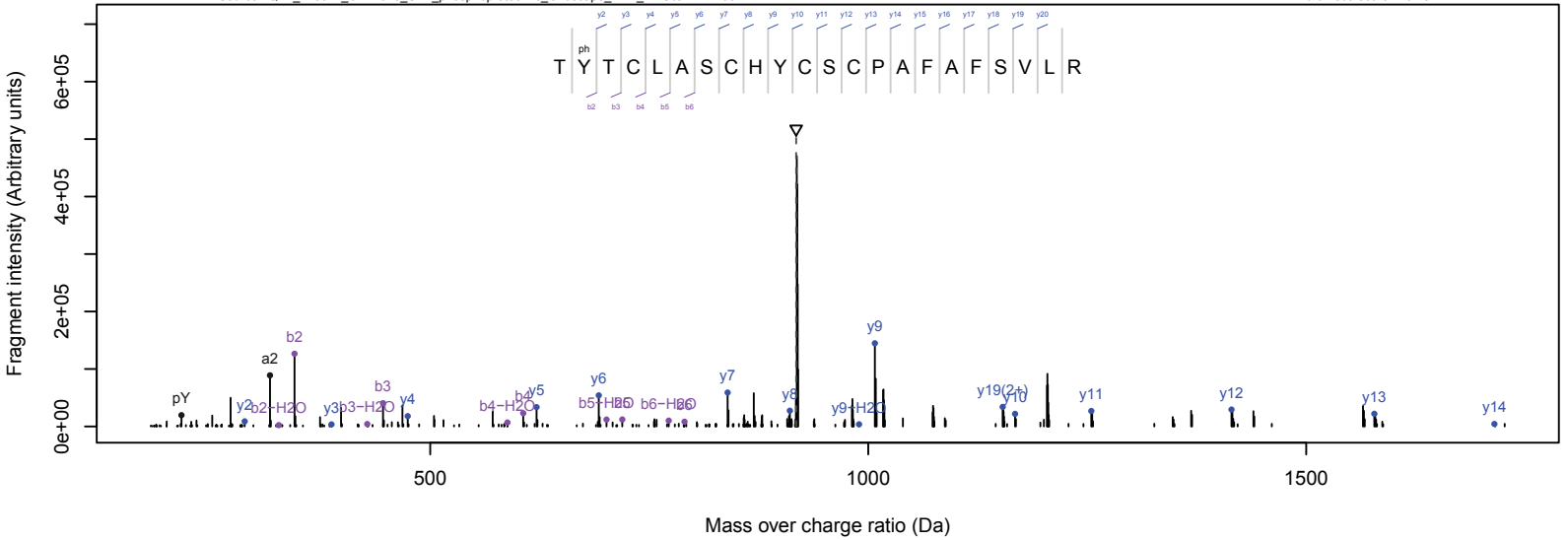
Andromeda score : 131.82



TY(ph)TCLASCHYCSCPFAFASVLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42100

Andromeda score : 131.82

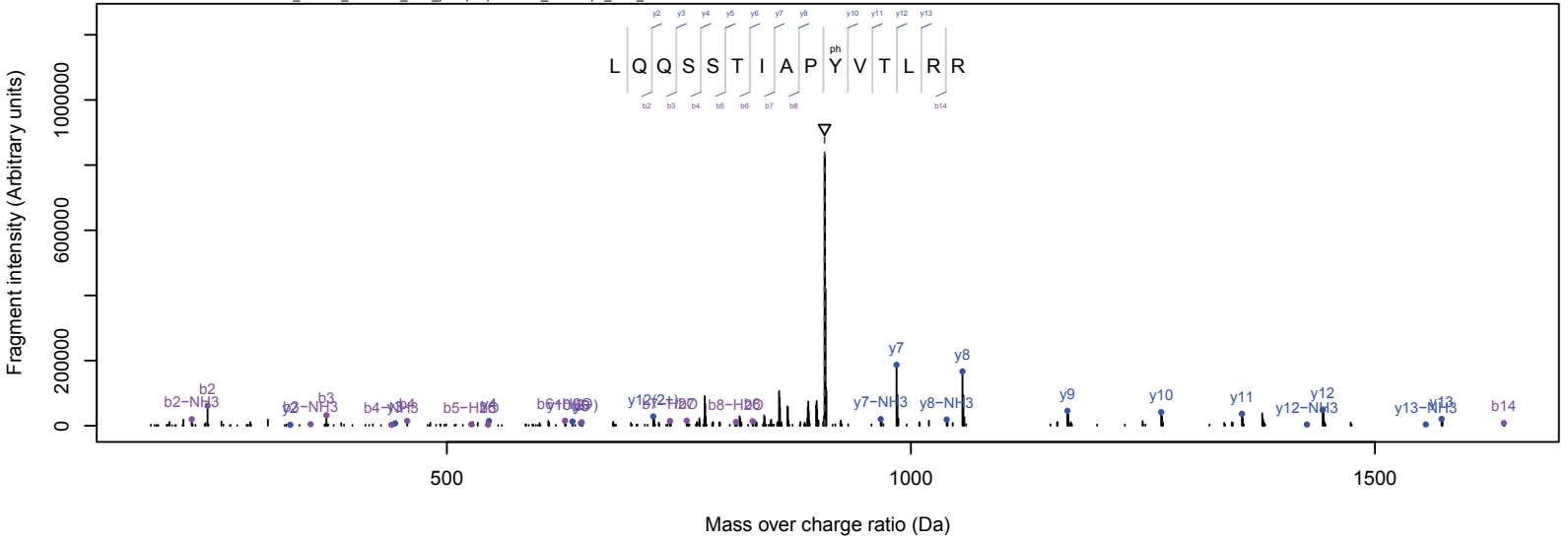




LQQSSTIAPY(ph)VTLRR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24319

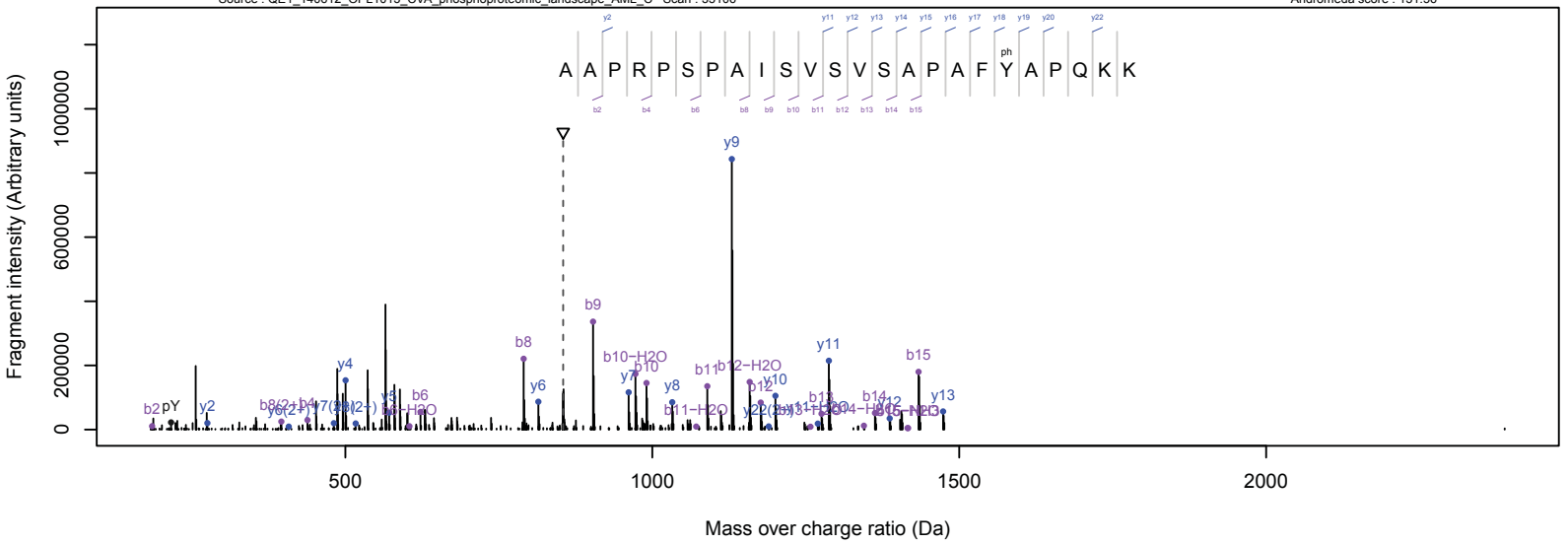
Andromeda score : 131.61



(ac)AAPRSPAISVSVSAPAFY(ph)APQKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35106

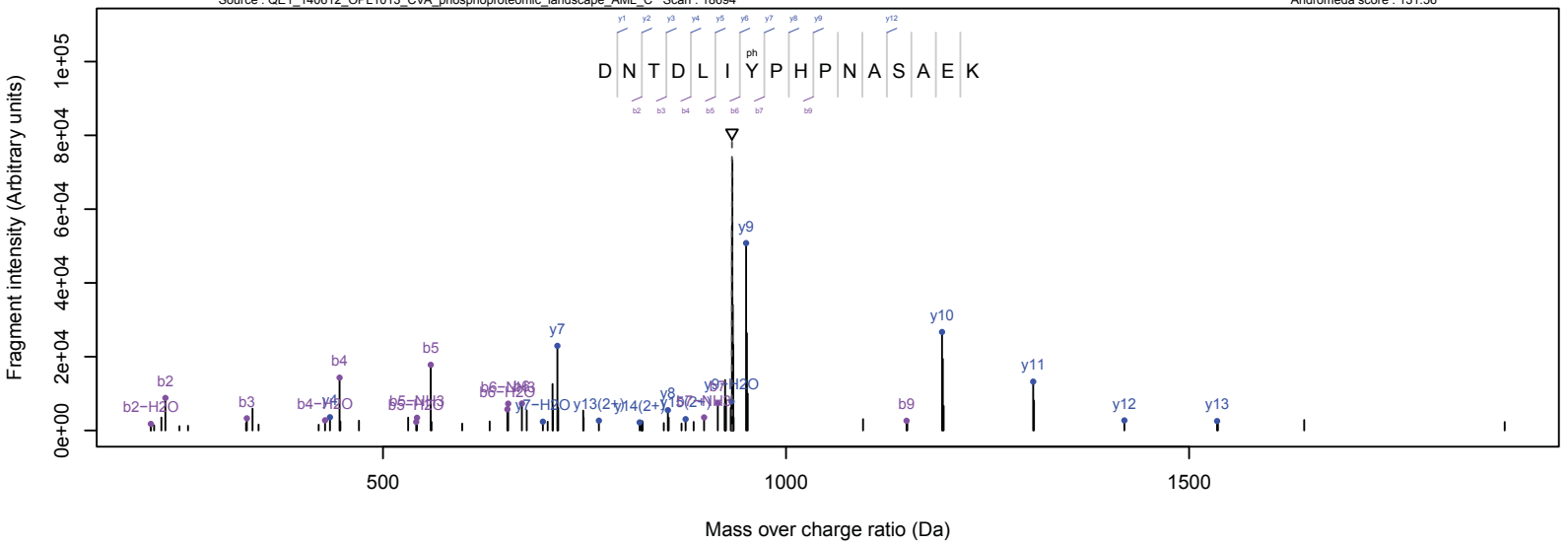
Andromeda score : 131.56



DNTDLIY(PHPN)ASAEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18694

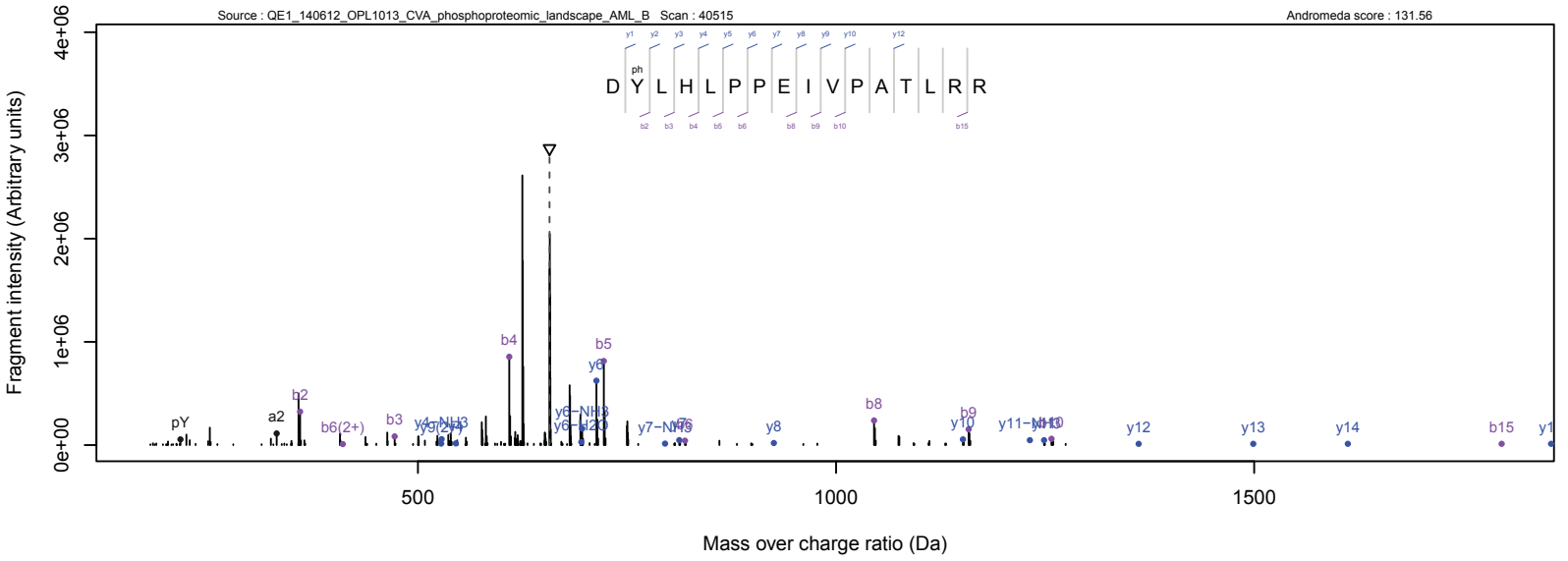
Andromeda score : 131.56



DY(ph)LHLPPEIVPATLRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40515

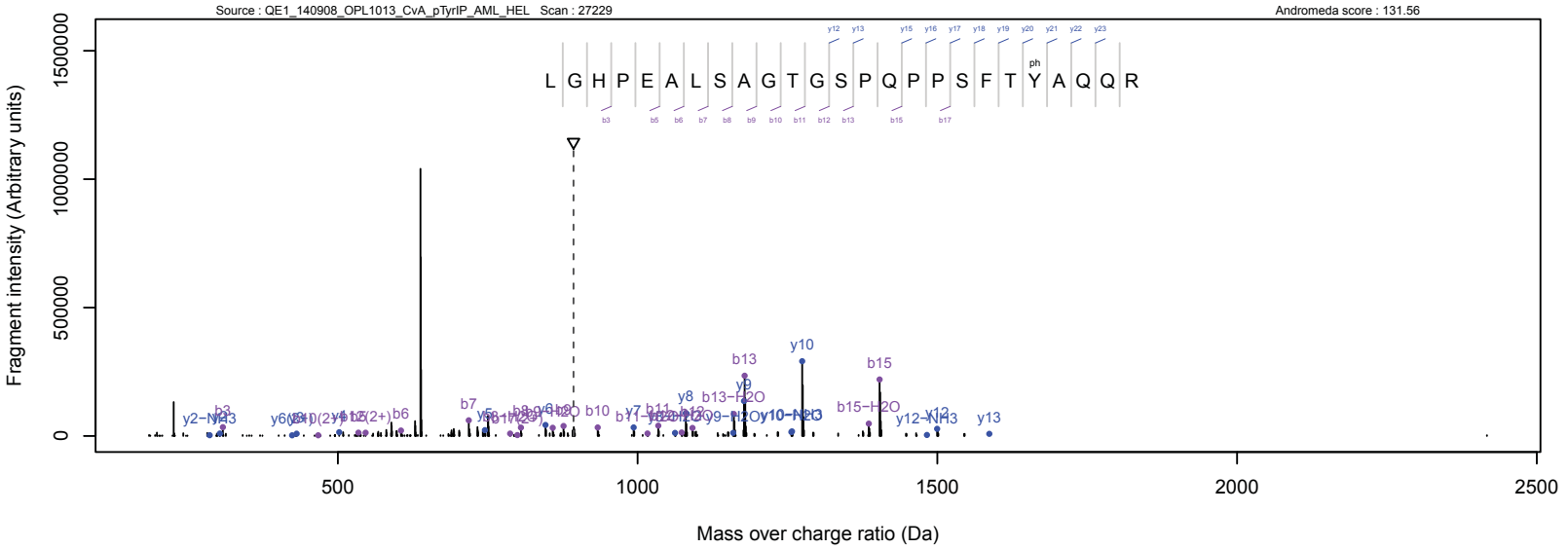
Andromeda score : 131.56



LGHPEALSAGTGSPPSFTY(ph)AQQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 27229

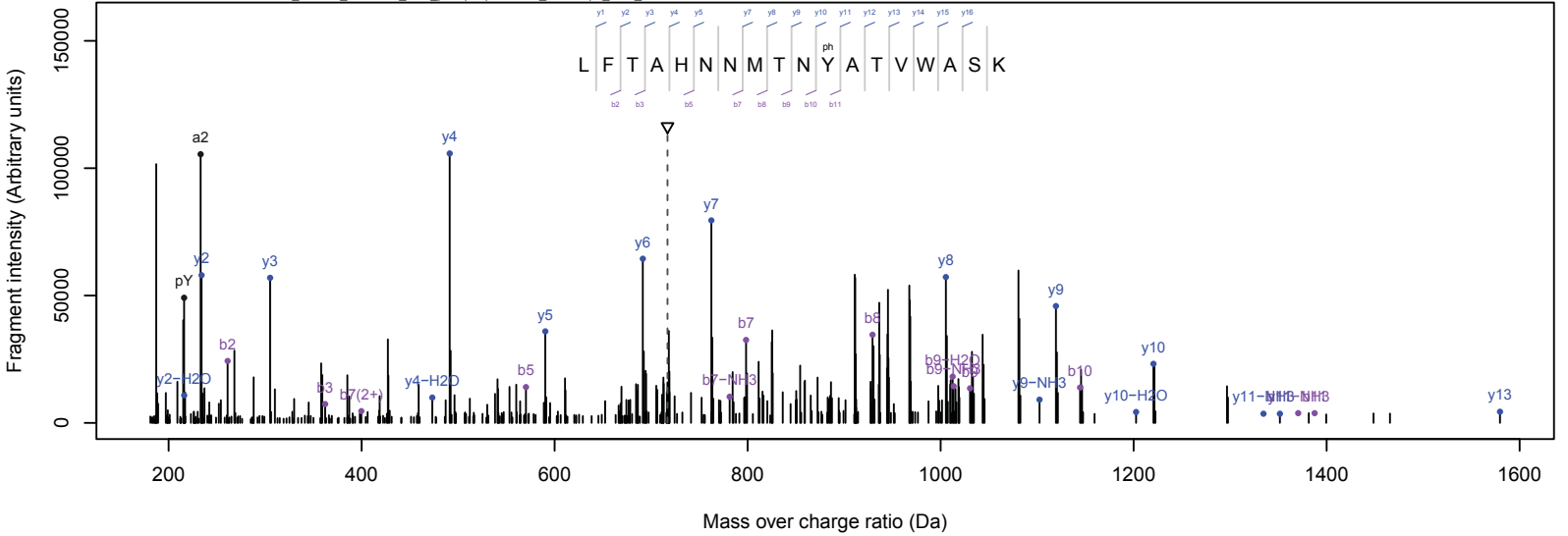
Andromeda score : 131.56



LFTAHHNMTNY(ph)ATVWASK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32038

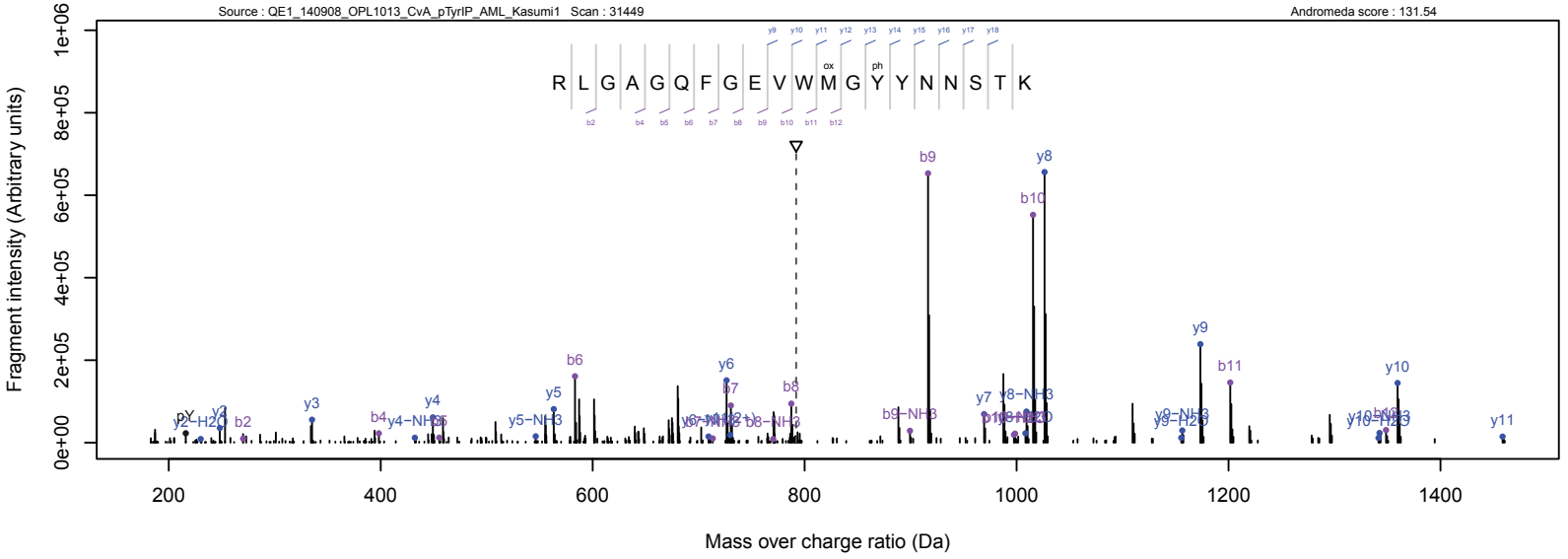
Andromeda score : 131.54



RLGAGQFGEVWVG<sup>ox</sup>GY<sup>ph</sup>YNNSTK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 31449

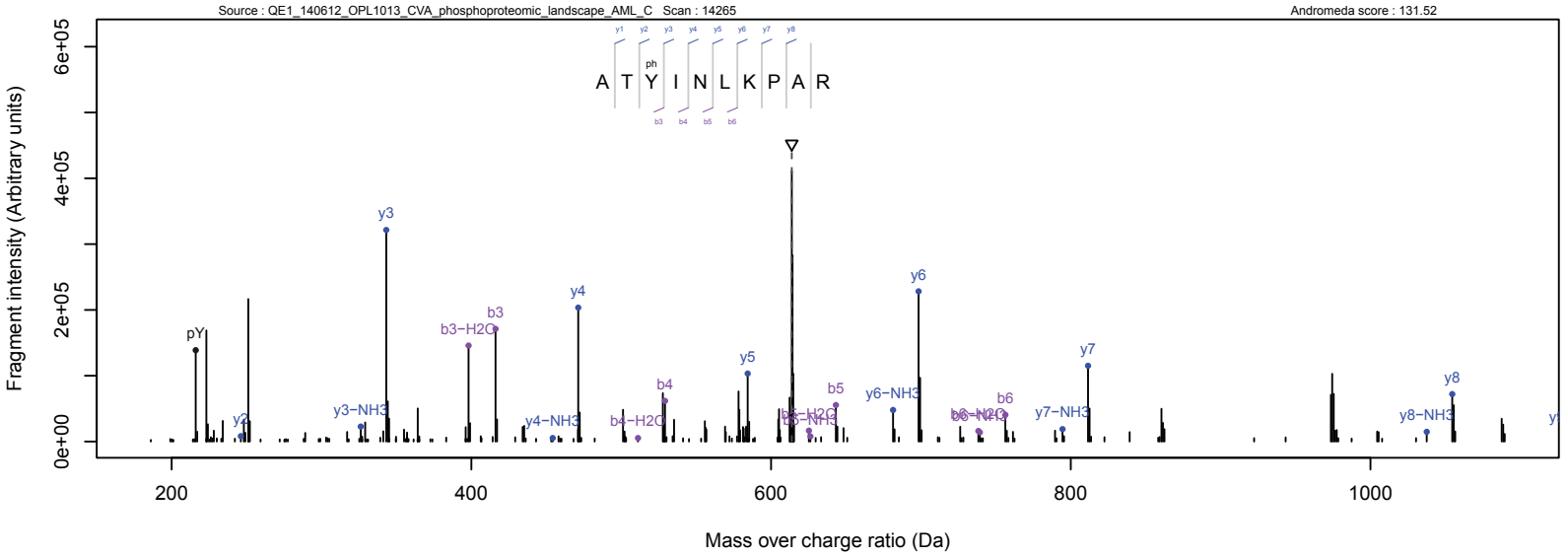
Andromeda score: 131.54



ATY<sup>ph</sup>INLKPAR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14265

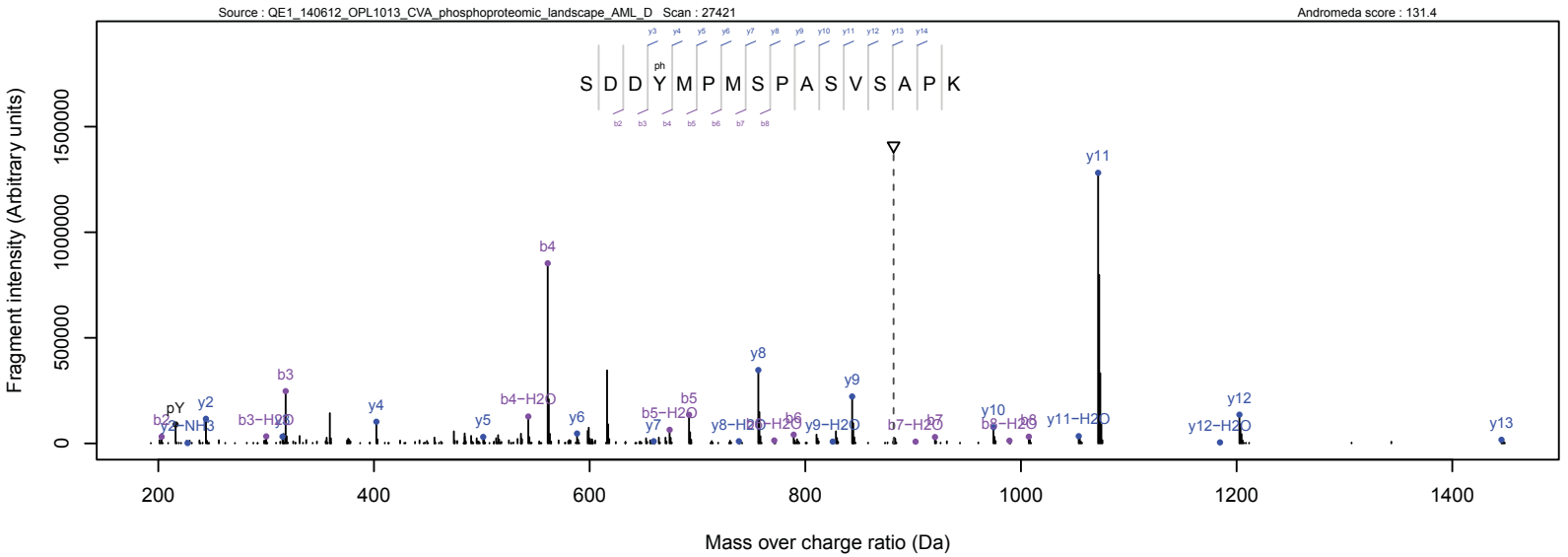
Andromeda score: 131.52



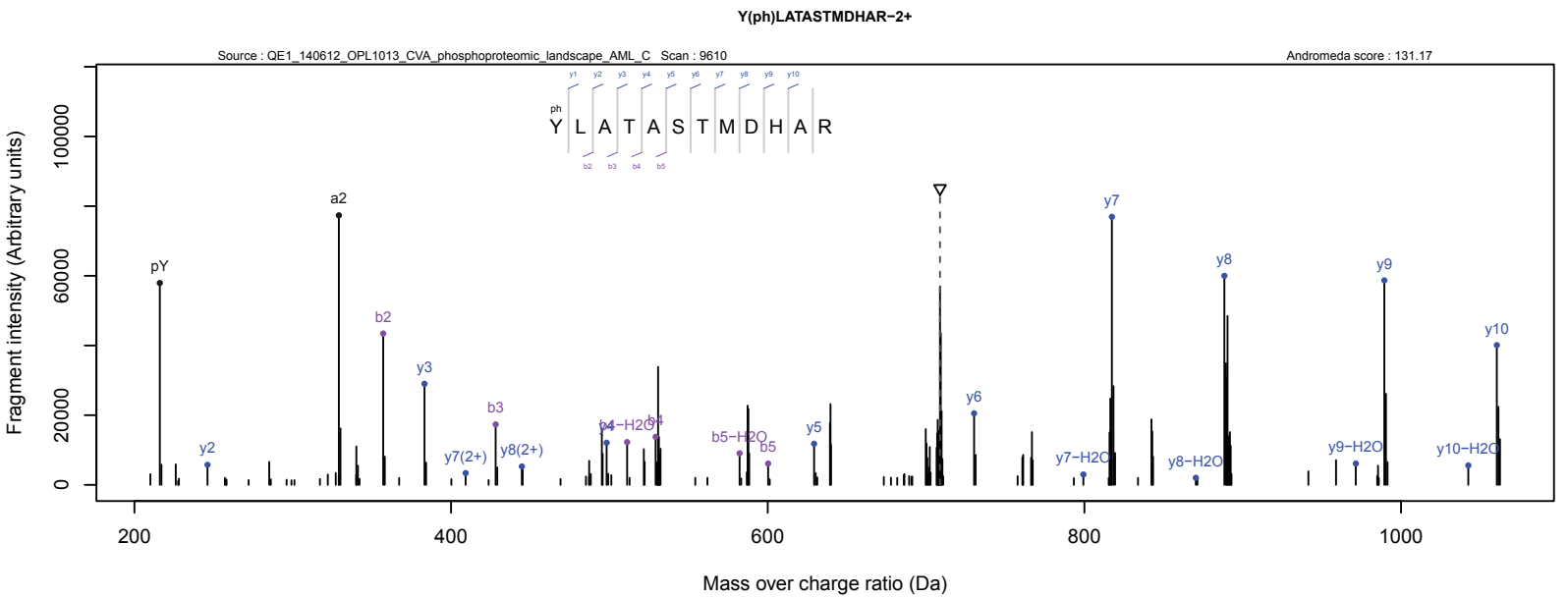
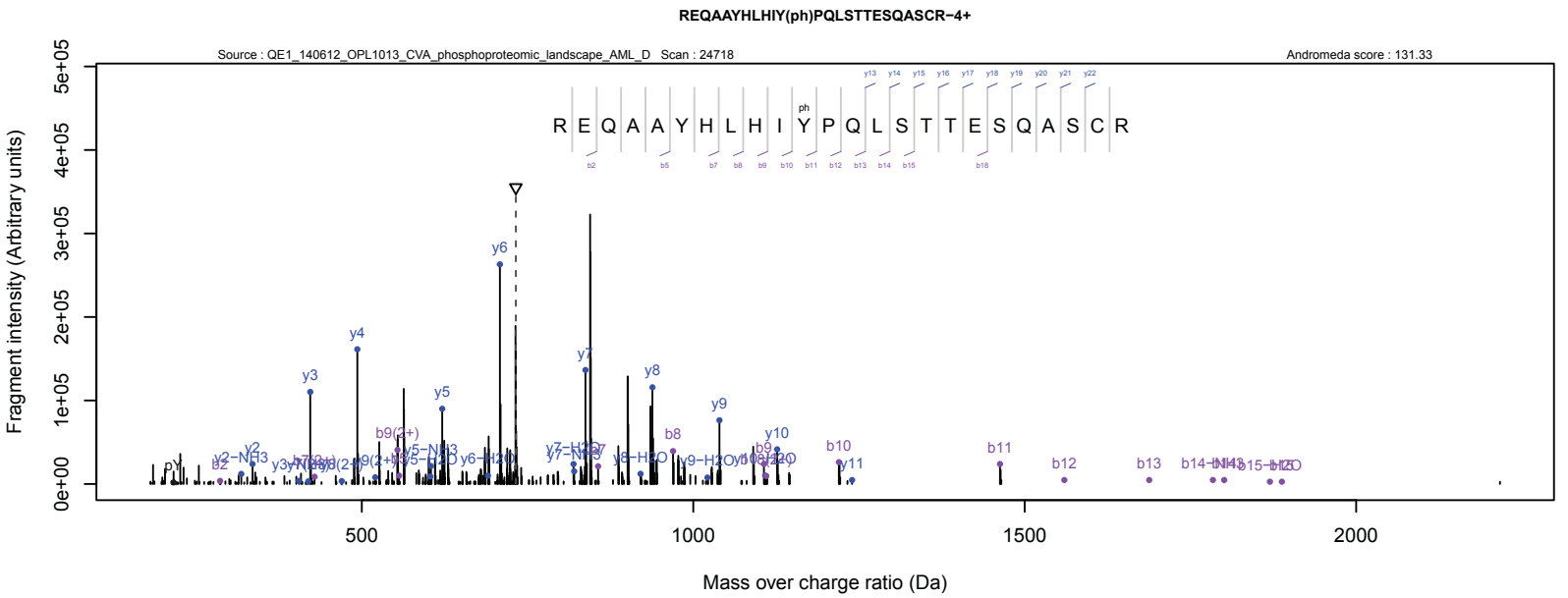
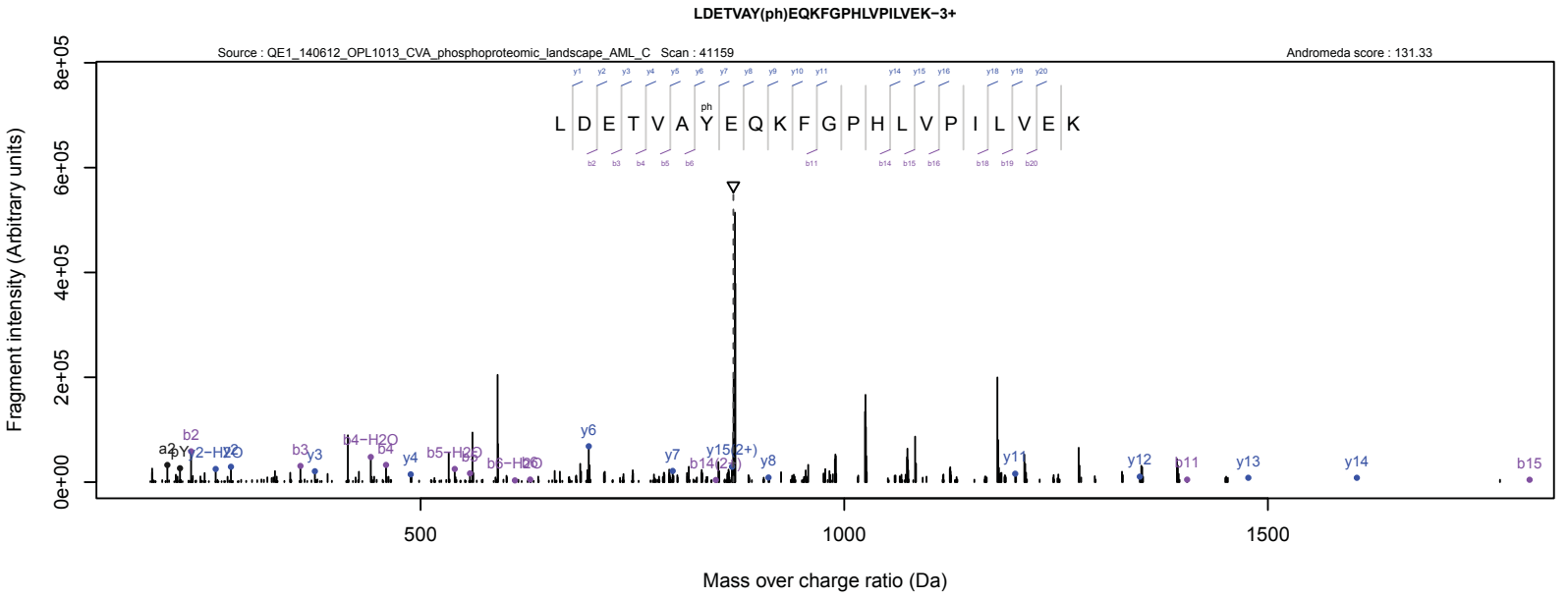
SDDY<sup>ph</sup>MPMSPASVSAPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 27421

Andromeda score: 131.4



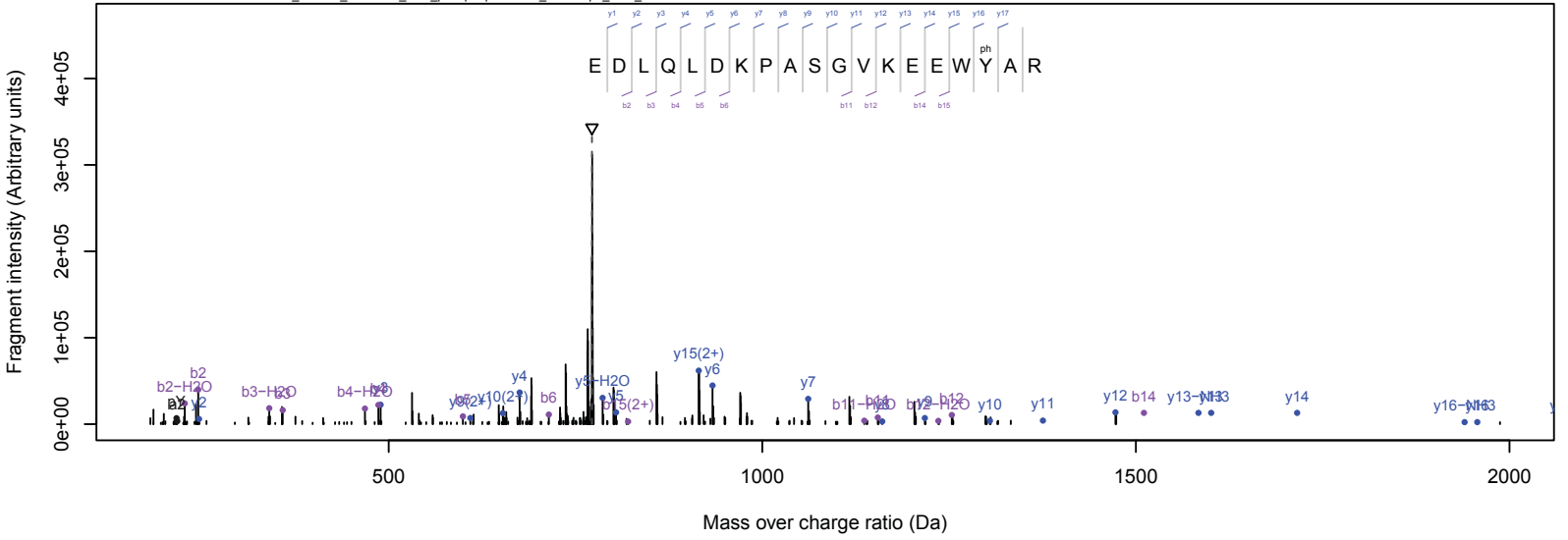




EDLQDKPASGVKEEWY(ph)AR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 25697

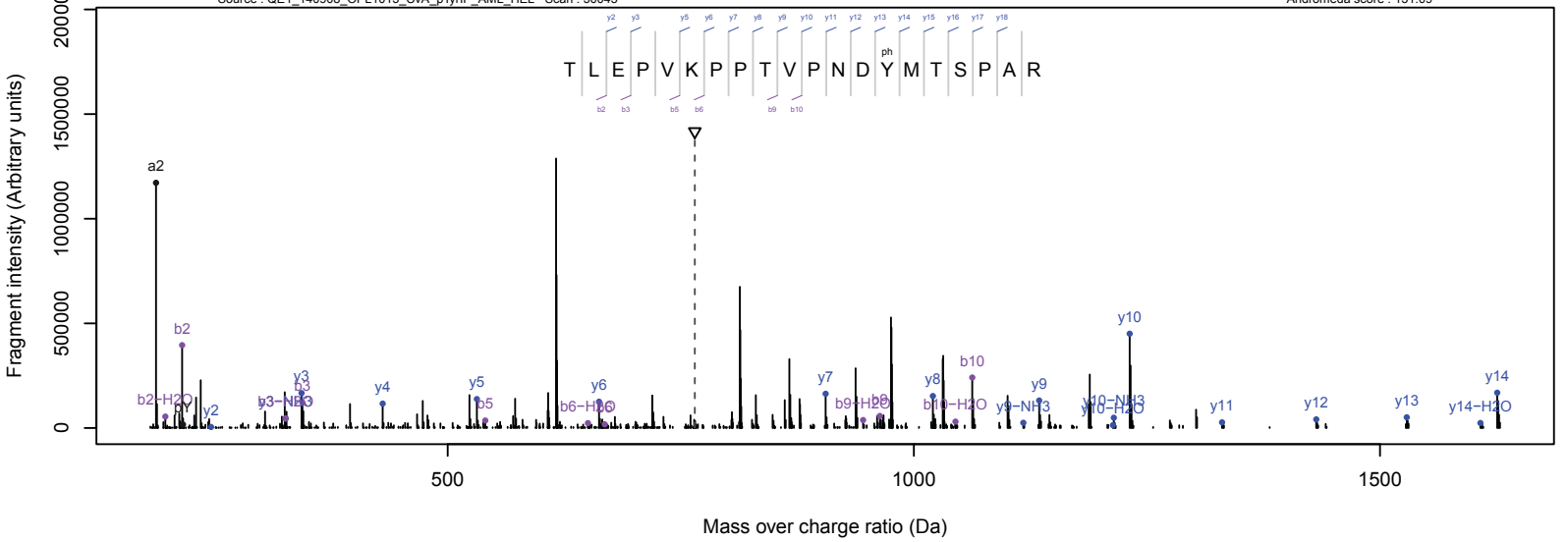
Andromeda score: 131.09



TLEPVKPTVPNDY(ph)MTSPAR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 30045

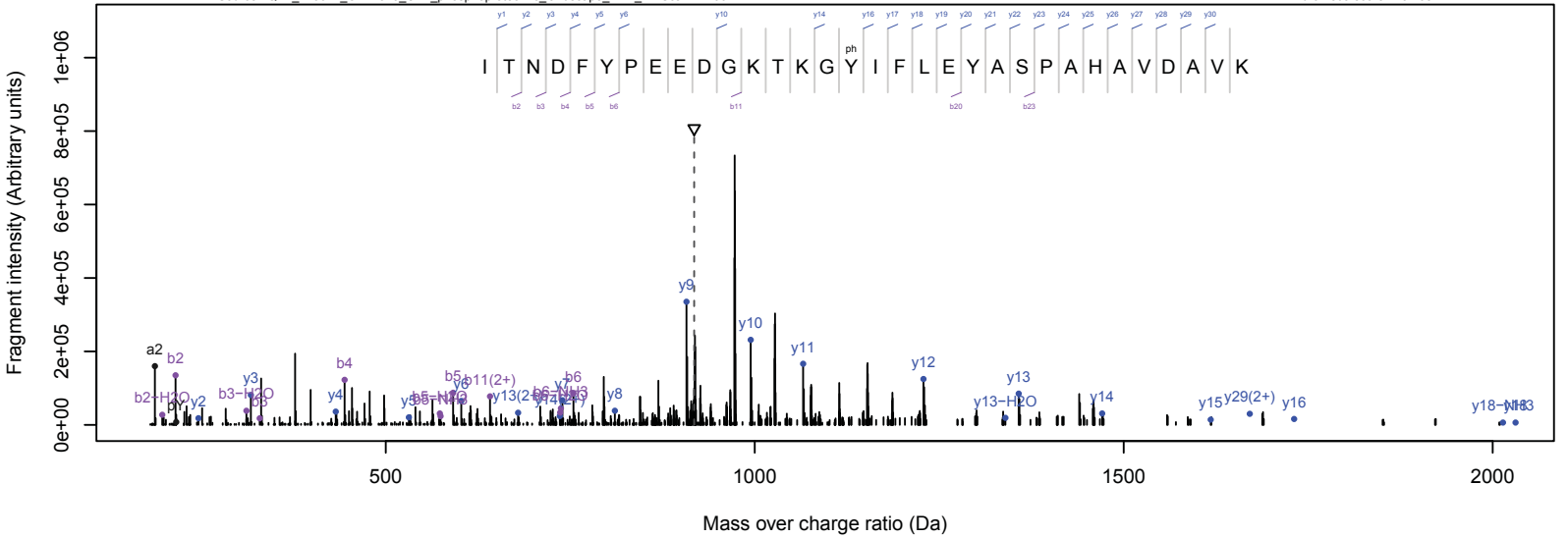
Andromeda score: 131.09



ITNDFYPEEDGKTKGY(ph)IFLEYASPAHA VDAVK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 41061

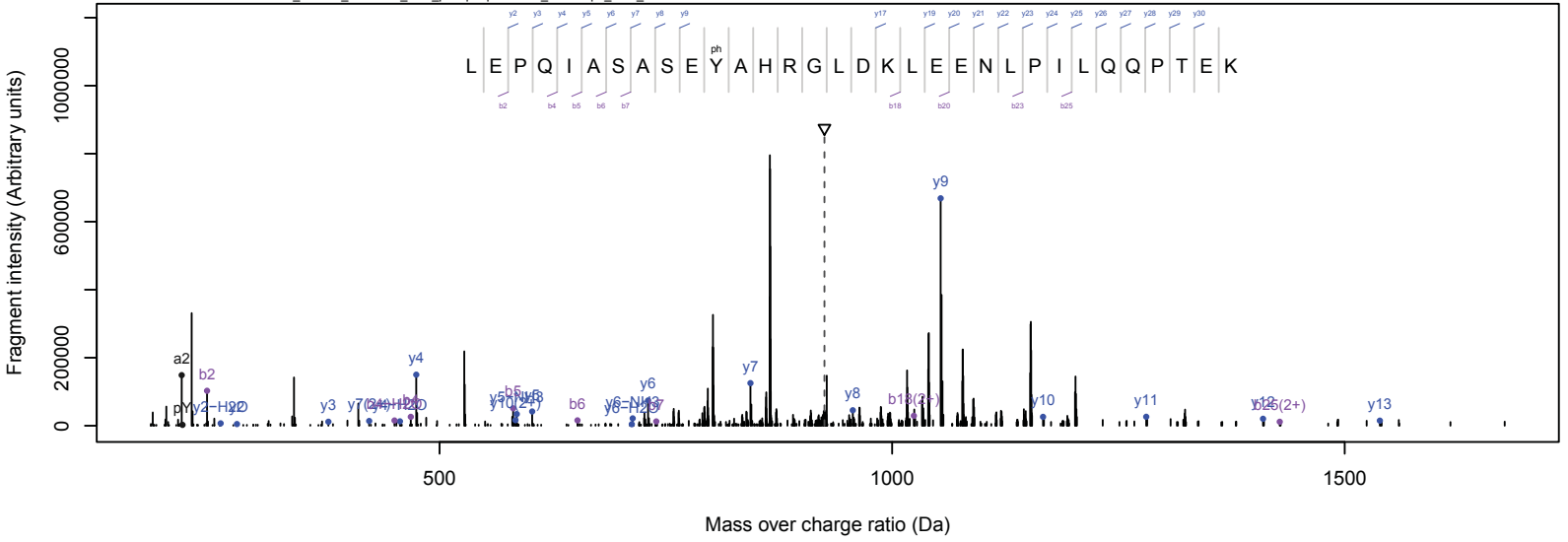
Andromeda score: 131.08



LEPQIASASEY(ph)AHRGLDKLEENLPILQQPTEK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46516

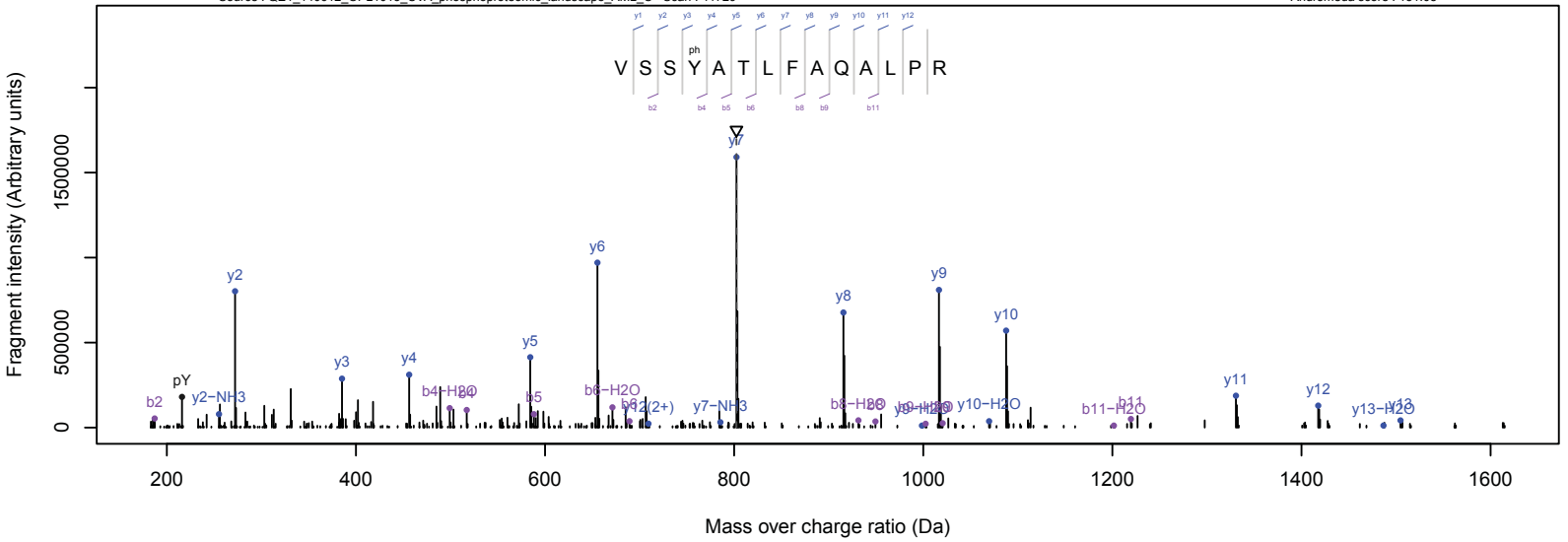
Andromeda score : 131.08



VSSY(ph)ATLFAQALPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41729

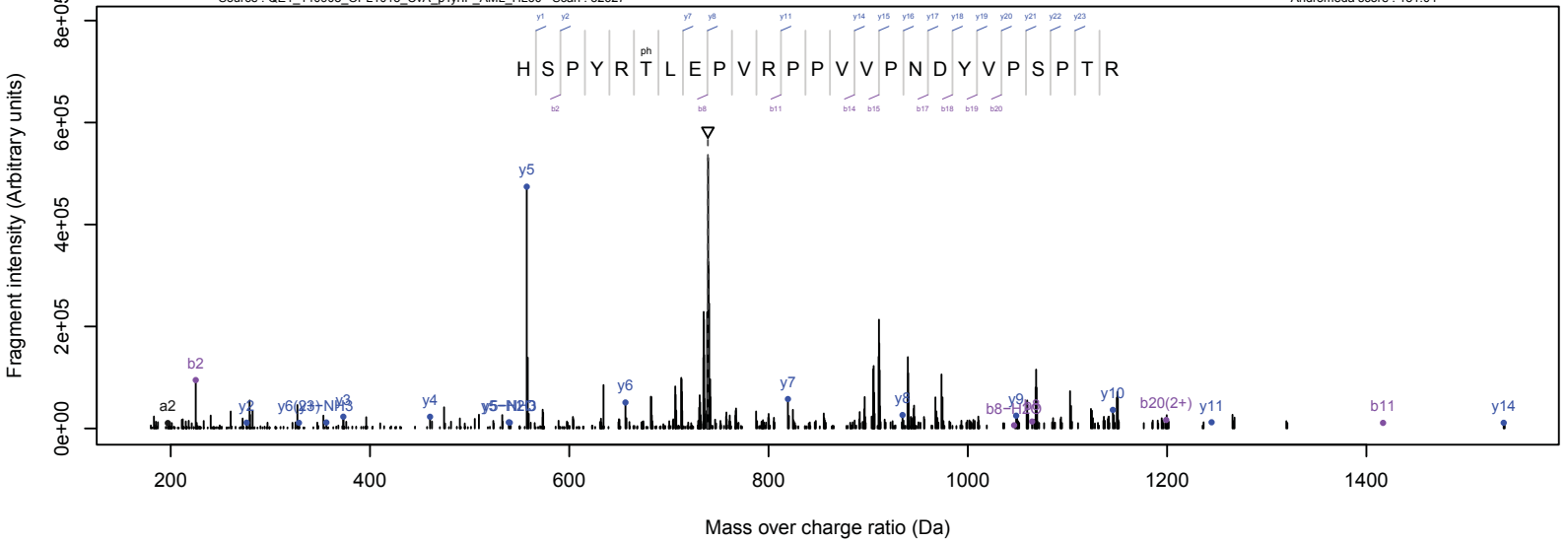
Andromeda score : 131.08



HSPYRT(ph)LEPVRPPVVPNDYVPSPTR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 32627

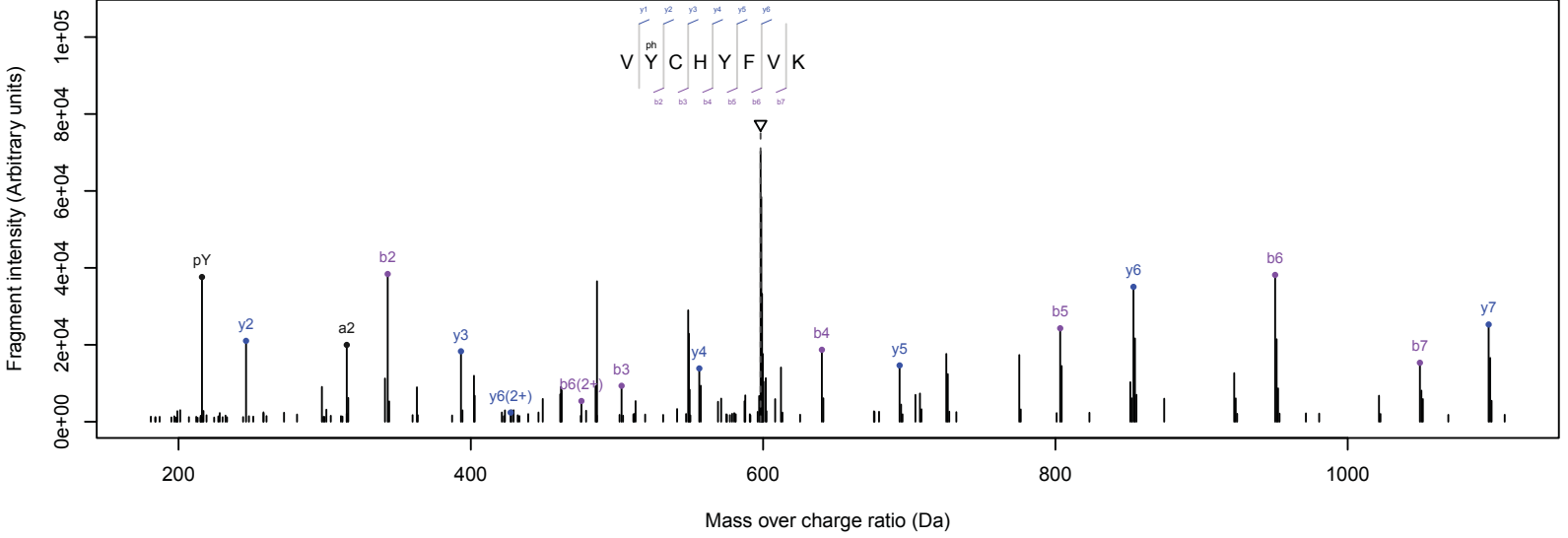
Andromeda score : 131.04



VY(ph)CHYFVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13451

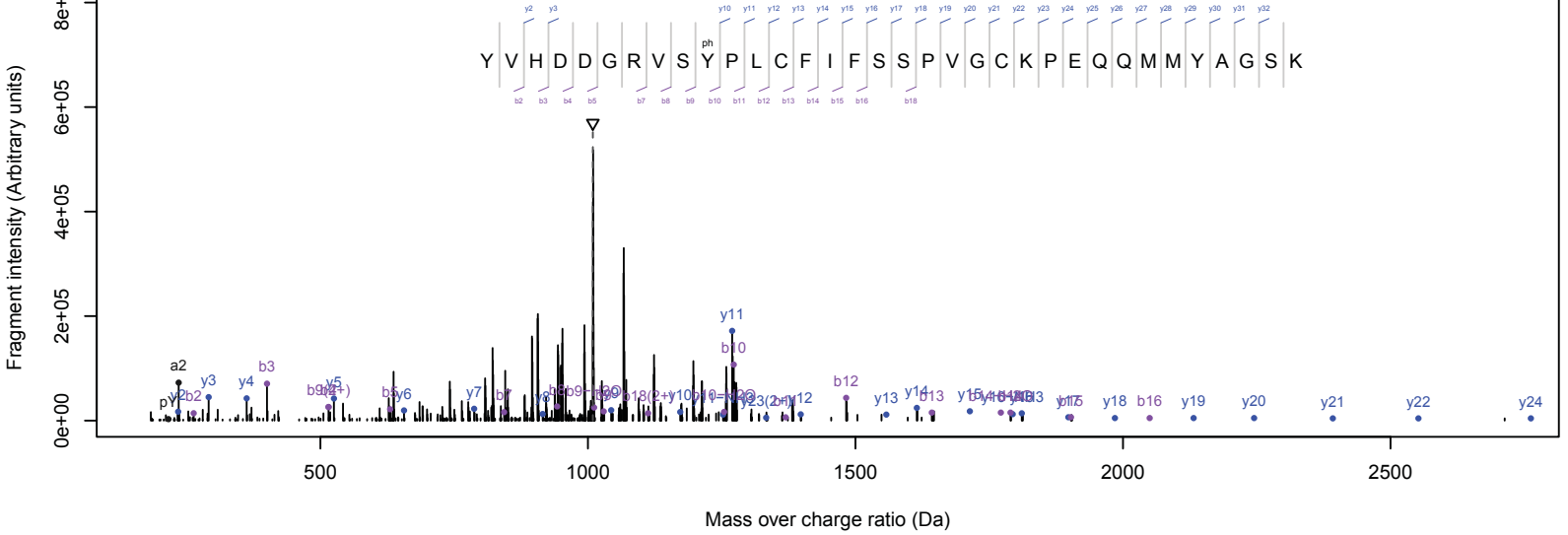
Andromeda score : 131.04



YVHDDGRVSY(ph)PLCFIFSSPVGCKPEQMMYAGSK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44441

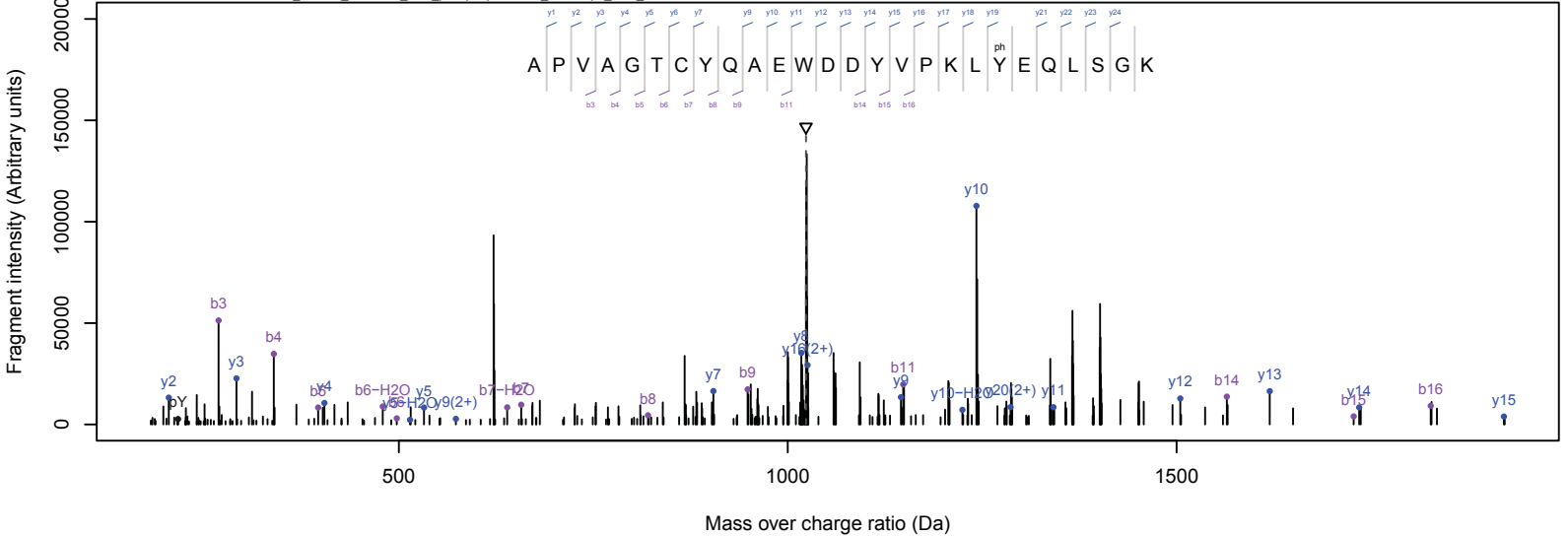
Andromeda score : 130.88



APVAGTCYQAEWDDYVPKLYEQLSGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46223

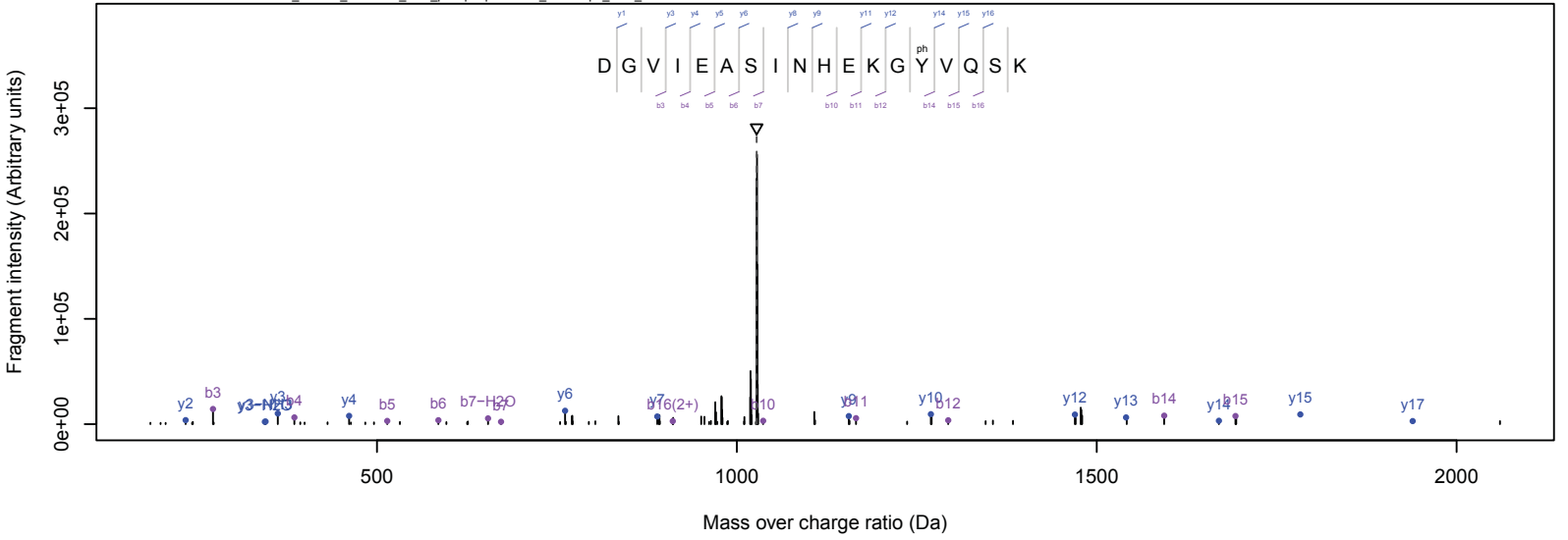
Andromeda score : 130.57



DGVIEASINHEKGYVQSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20739

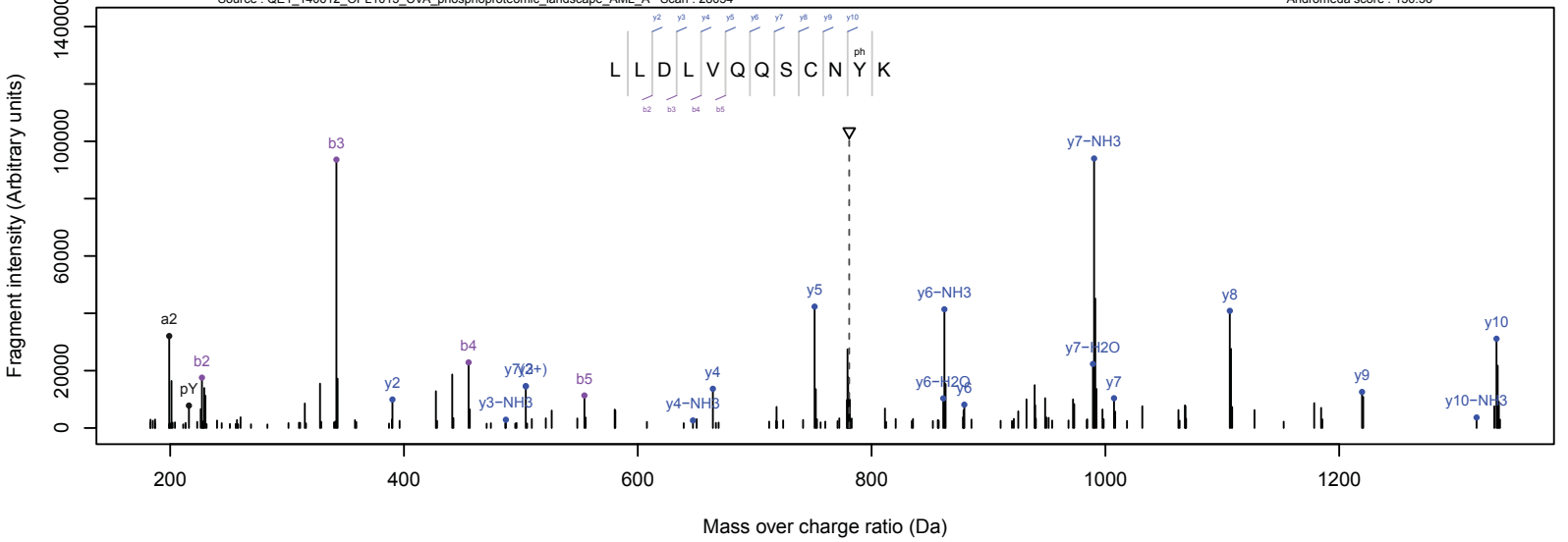
Andromeda score : 130.56



LLDLVQSCNYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 28054

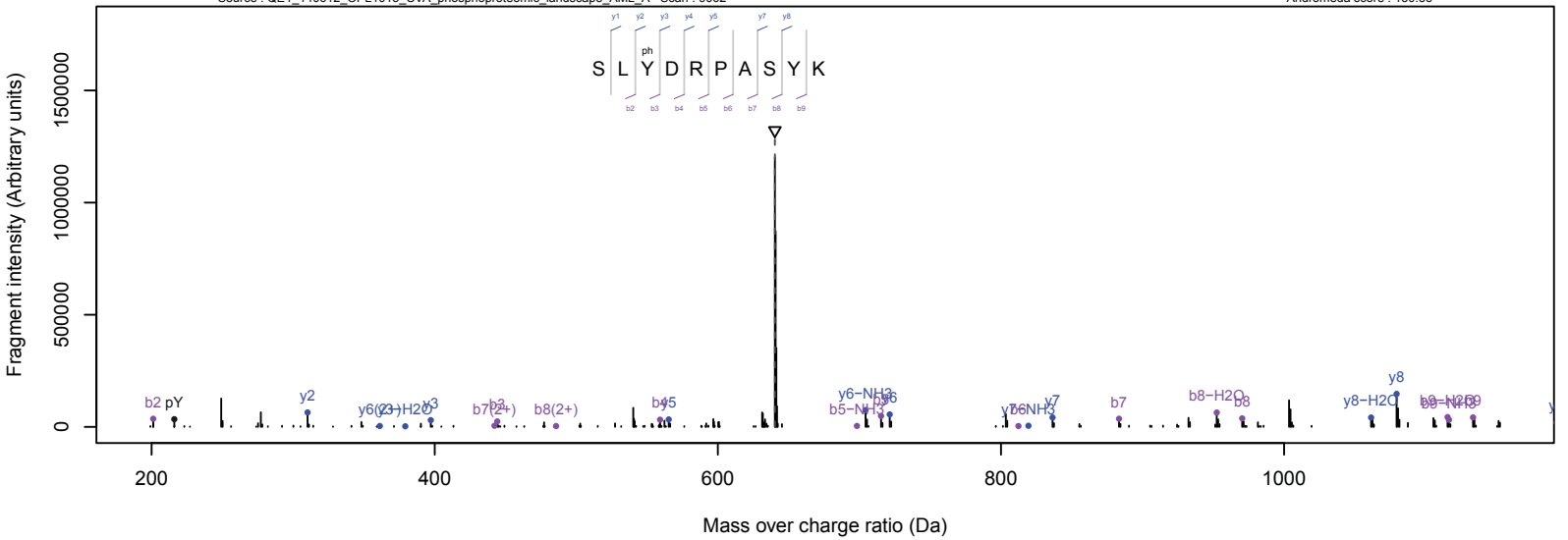
Andromeda score : 130.56



SLY(ph)DRPASYSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 9062

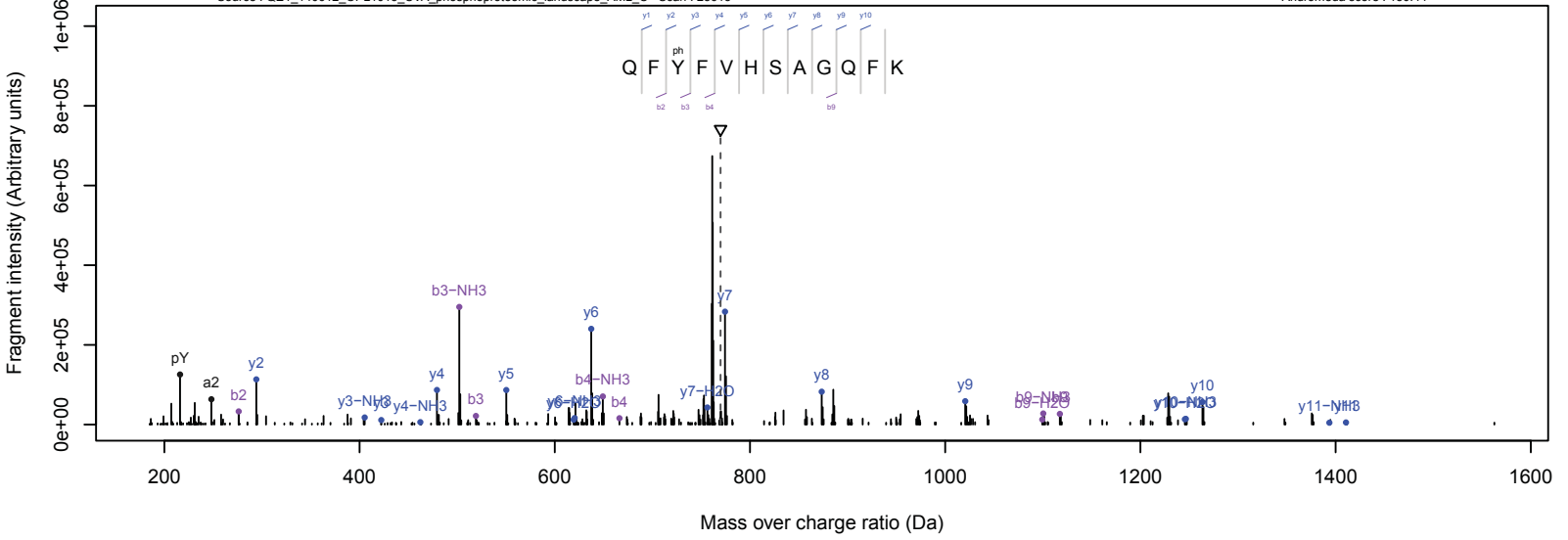
Andromeda score : 130.56



QFY(ph)FVHSAGQFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28615

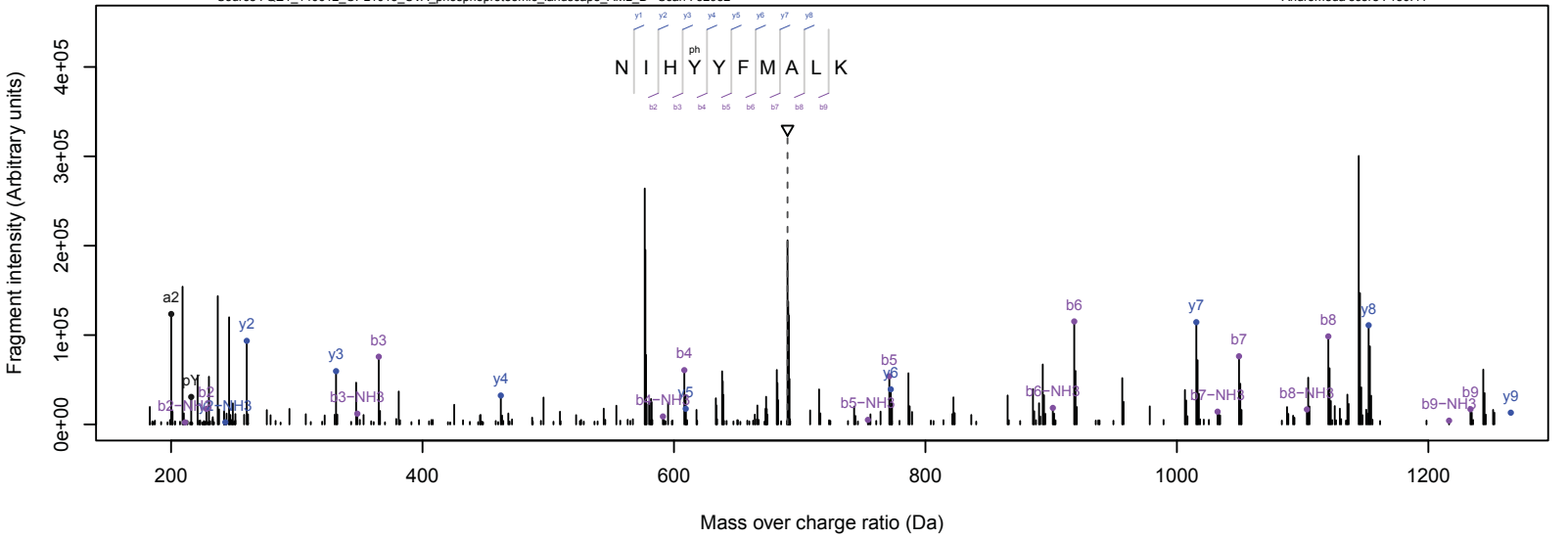
Andromeda score : 130.44



NIHY(ph)YFMALK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32982

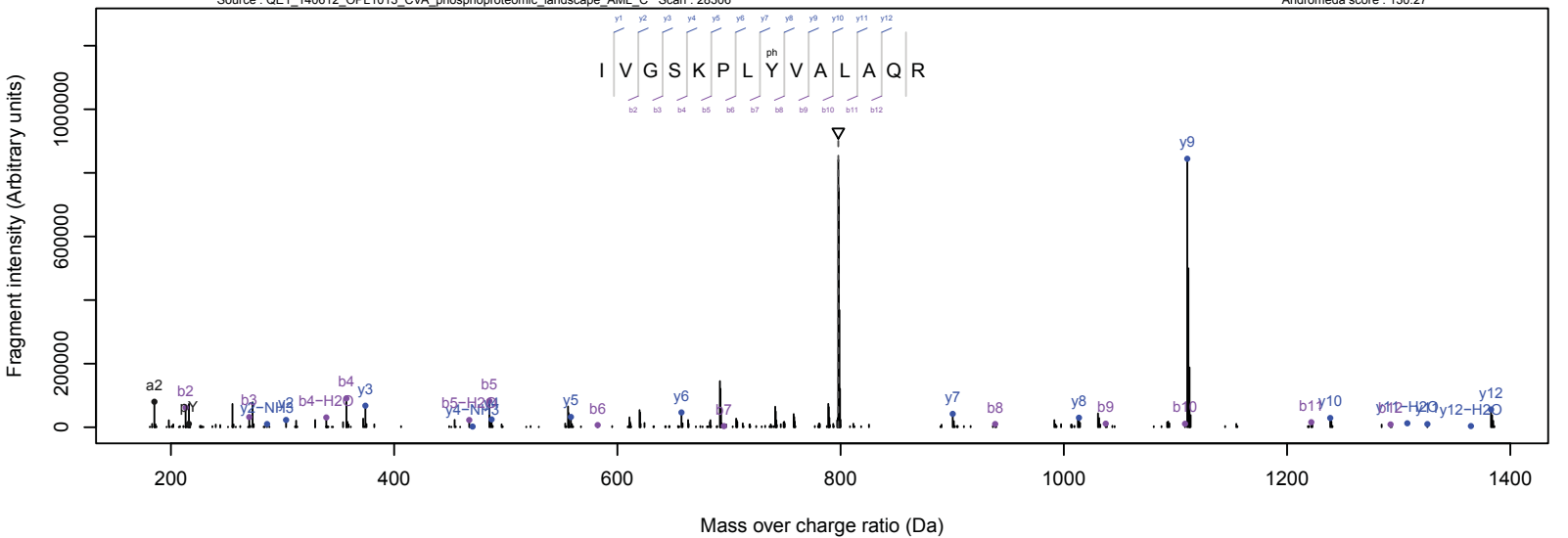
Andromeda score : 130.41



IVGSKPLY(ph)VALAQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28306

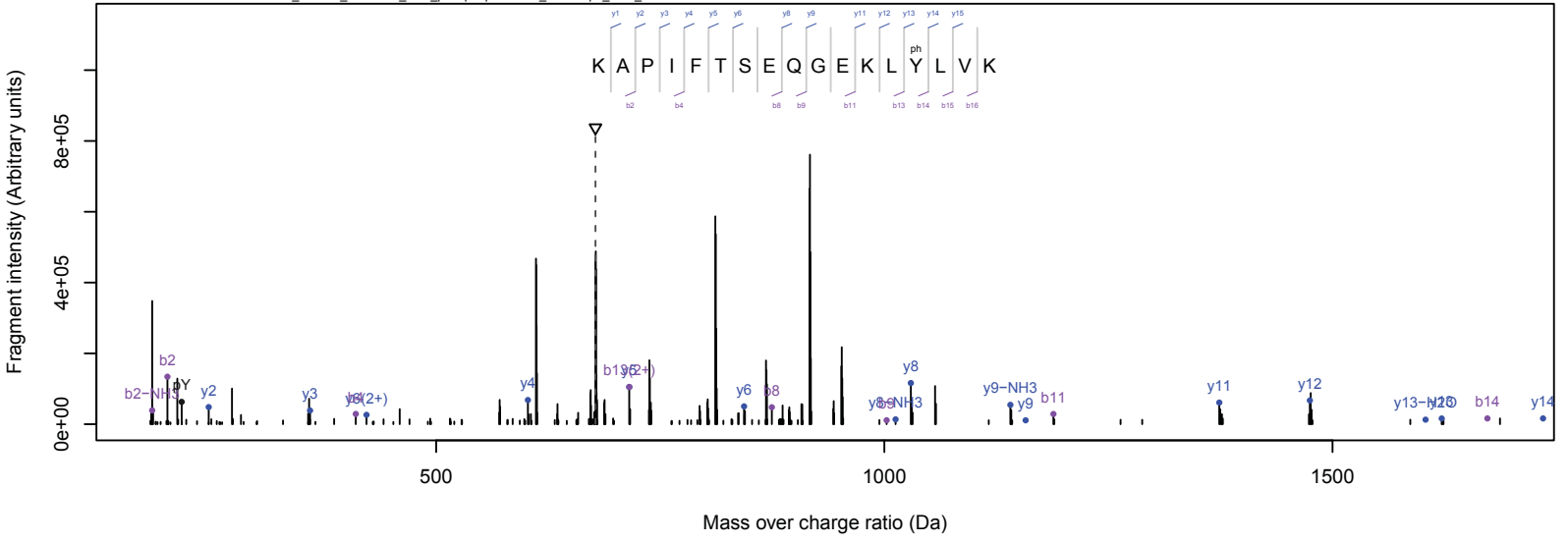
Andromeda score : 130.27



**KAPIFTSEQEKLYLVK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28393

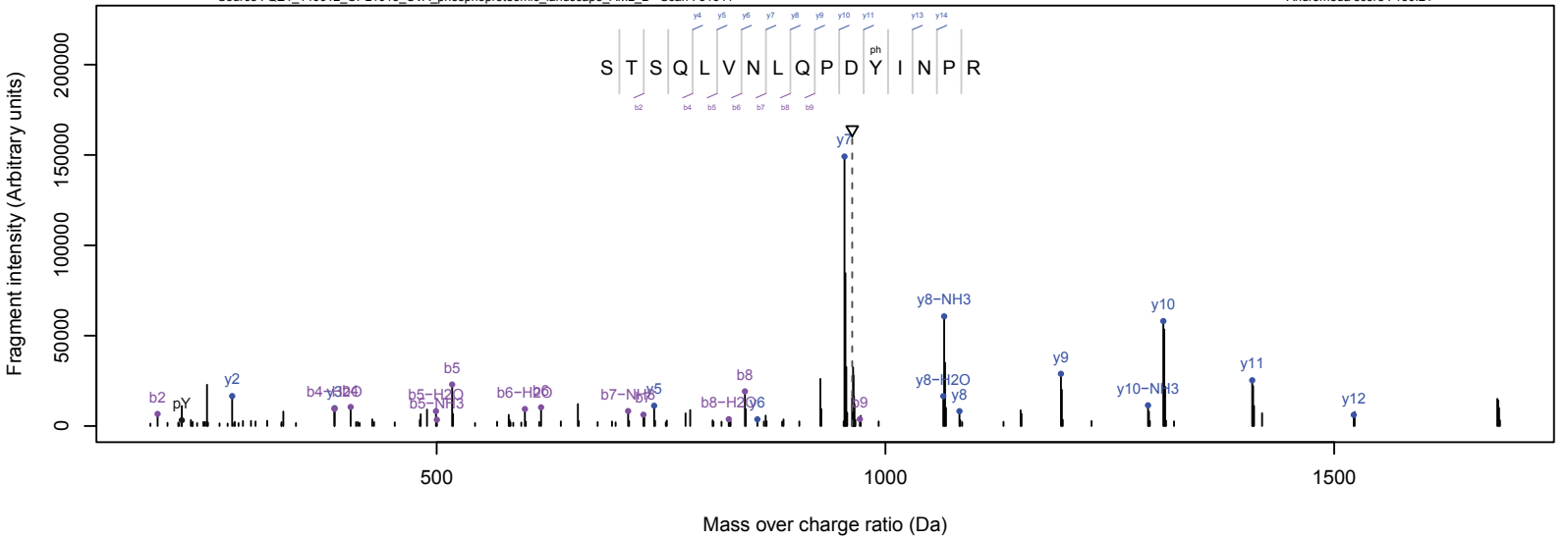
Andromeda score : 130.27



**STSQLVNLQPDYINPR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31641

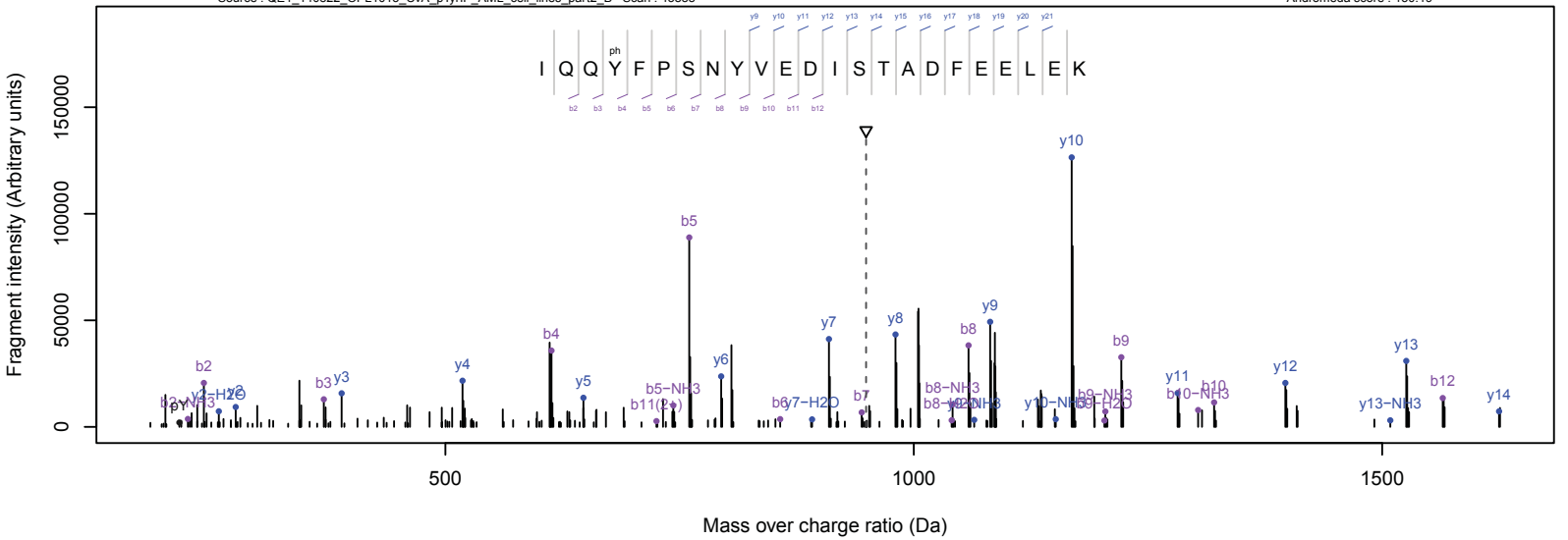
Andromeda score : 130.21



**IQQY(ph)FPSNYVEDISTADFEELK-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 43888

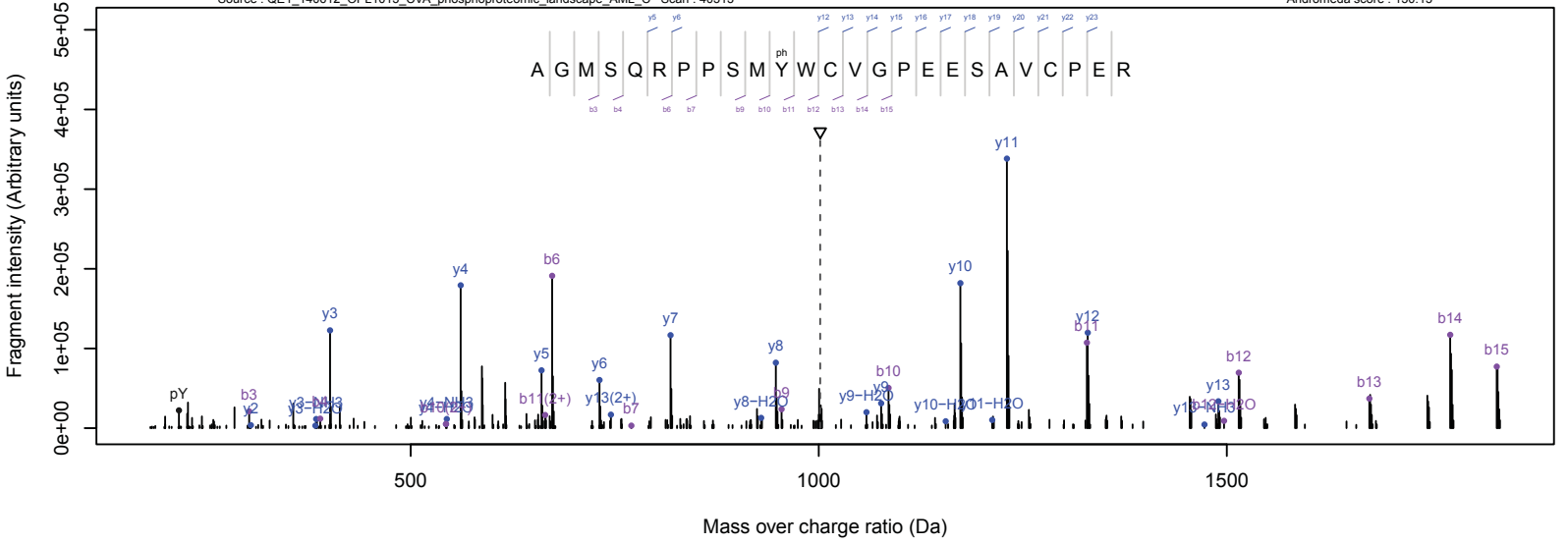
Andromeda score : 130.19



(ac)AGMSQRPPSMY(W)WCVGPEESAVCPER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40313

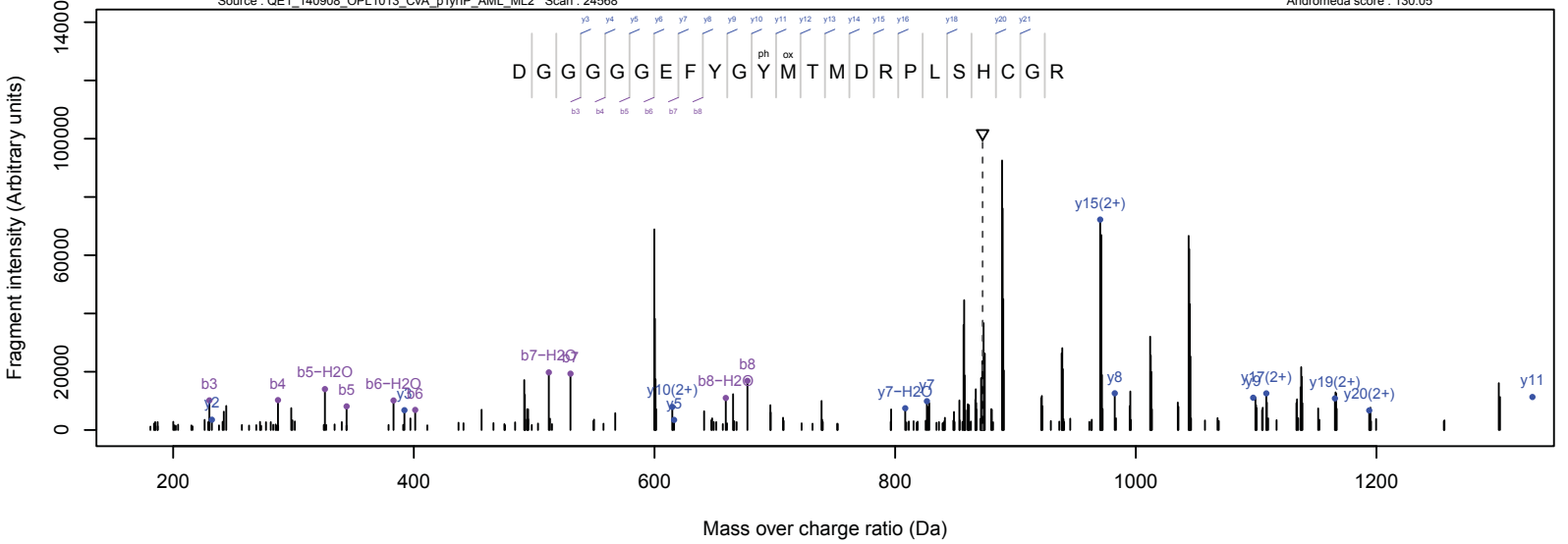
Andromeda score : 130.15



DGGGGGFYGY(M)(ox)TMDRPLSHCGR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 24568

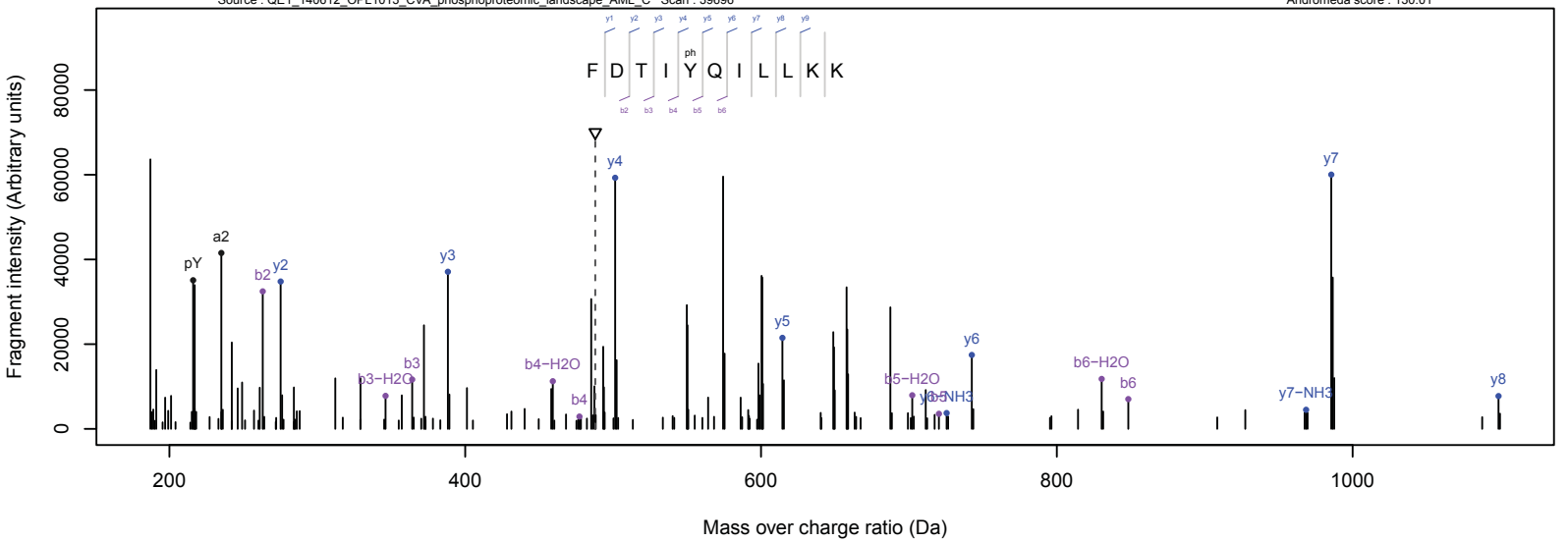
Andromeda score : 130.05



FDTIY(Q)ILLKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39696

Andromeda score : 130.01

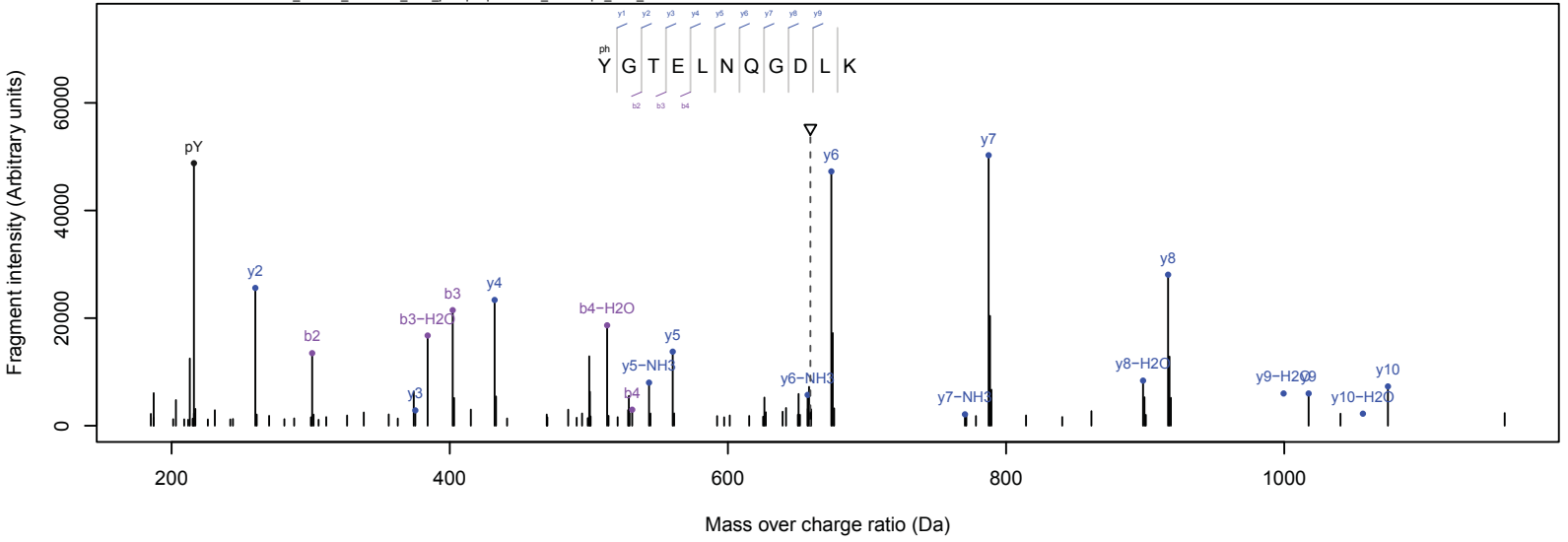




**Y(ph)GTELNQDLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12050

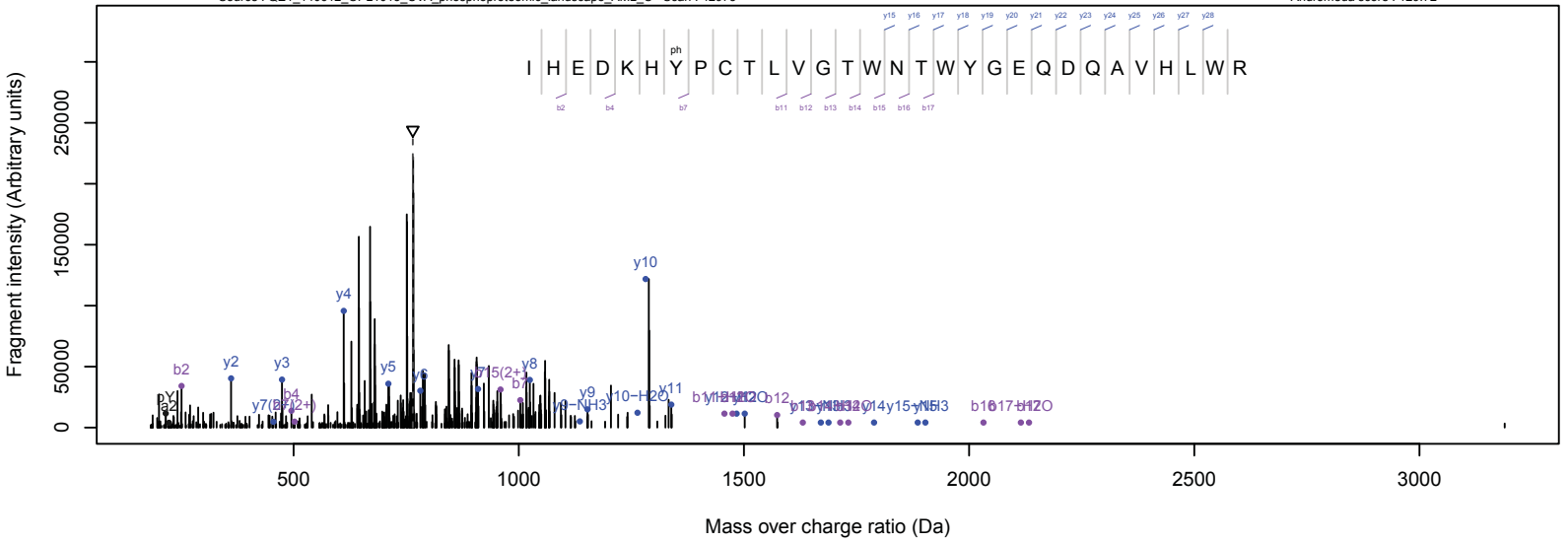
Andromeda score : 129.82



**IHEDKHY(ph)PCTLVGTWNTWYGEQDQAVHLWR-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42973

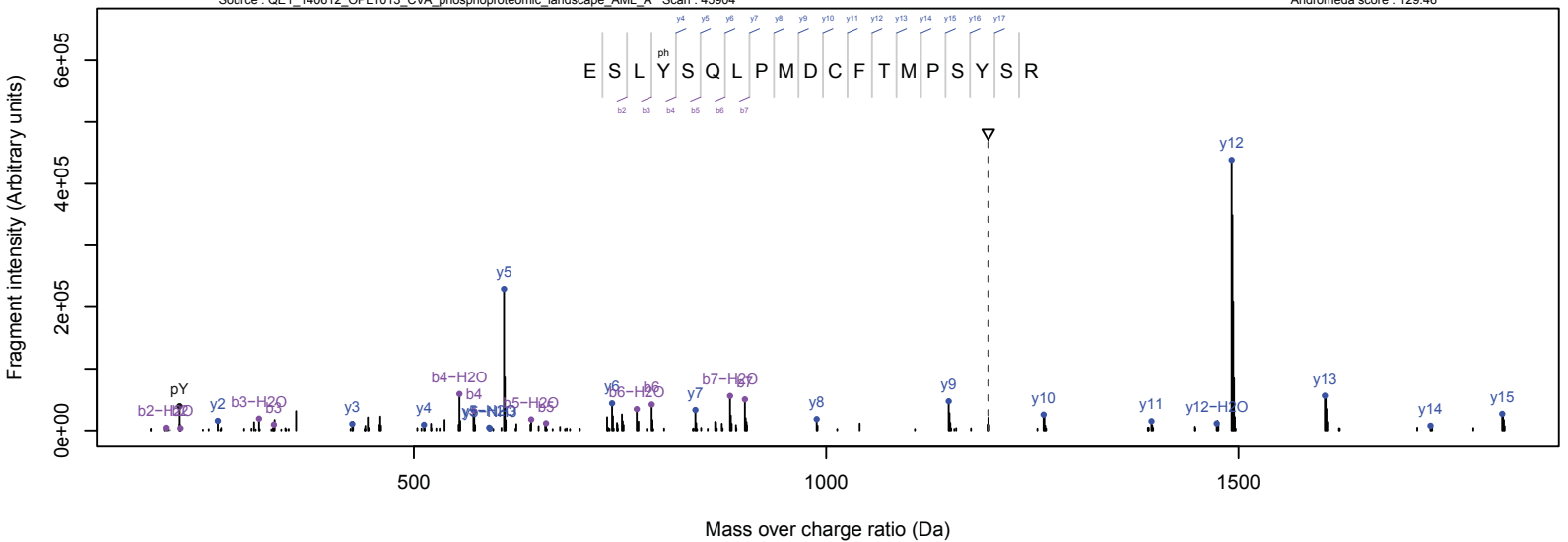
Andromeda score : 129.72



**ESLY(ph)SQLPMDCFMPSYSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 45904

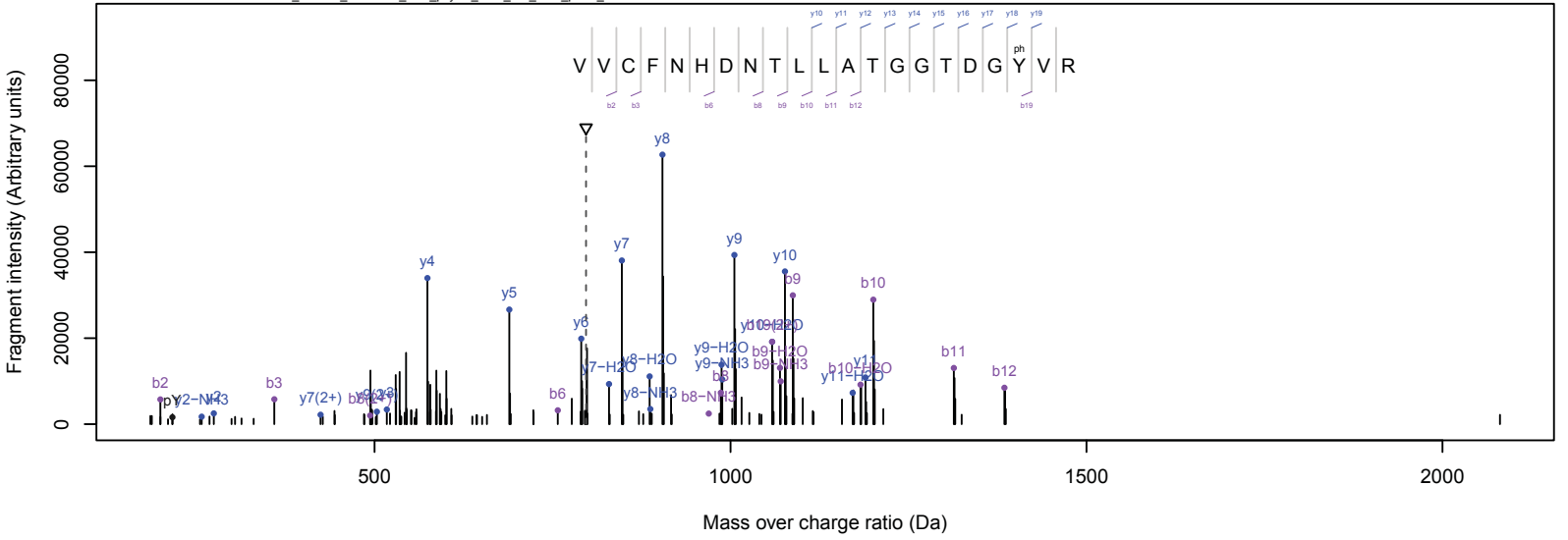
Andromeda score : 129.46



VVCFNHDNTLLATGGTDGYVR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 22064

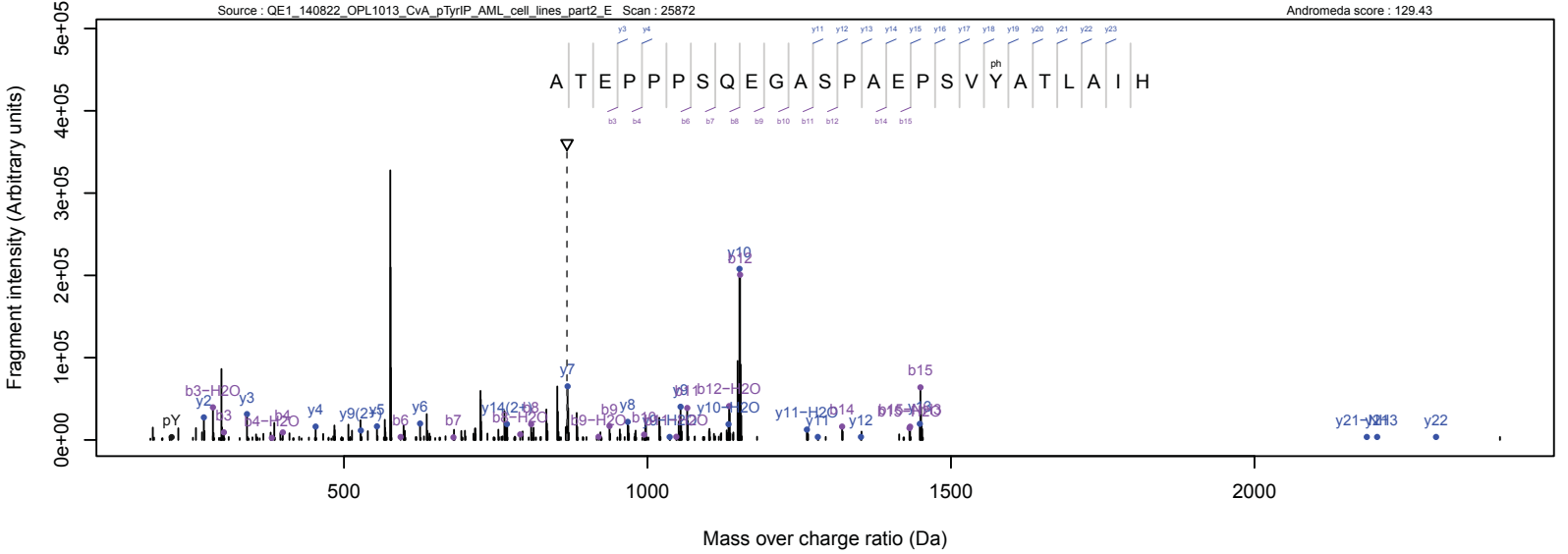
Andromeda score : 129.46



ATEPPSQEGASPAEPSVYATLAIH-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 25872

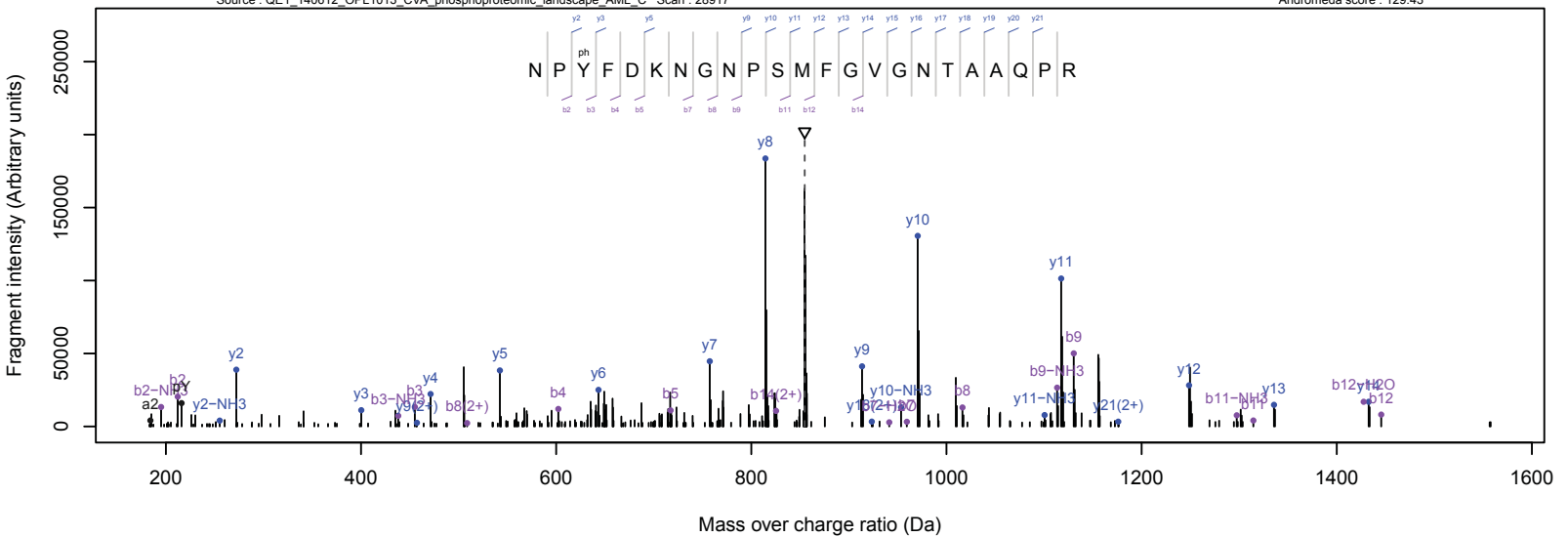
Andromeda score : 129.43



NPY(ph)FDKNGNPSMFGVGNATAAQP R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28917

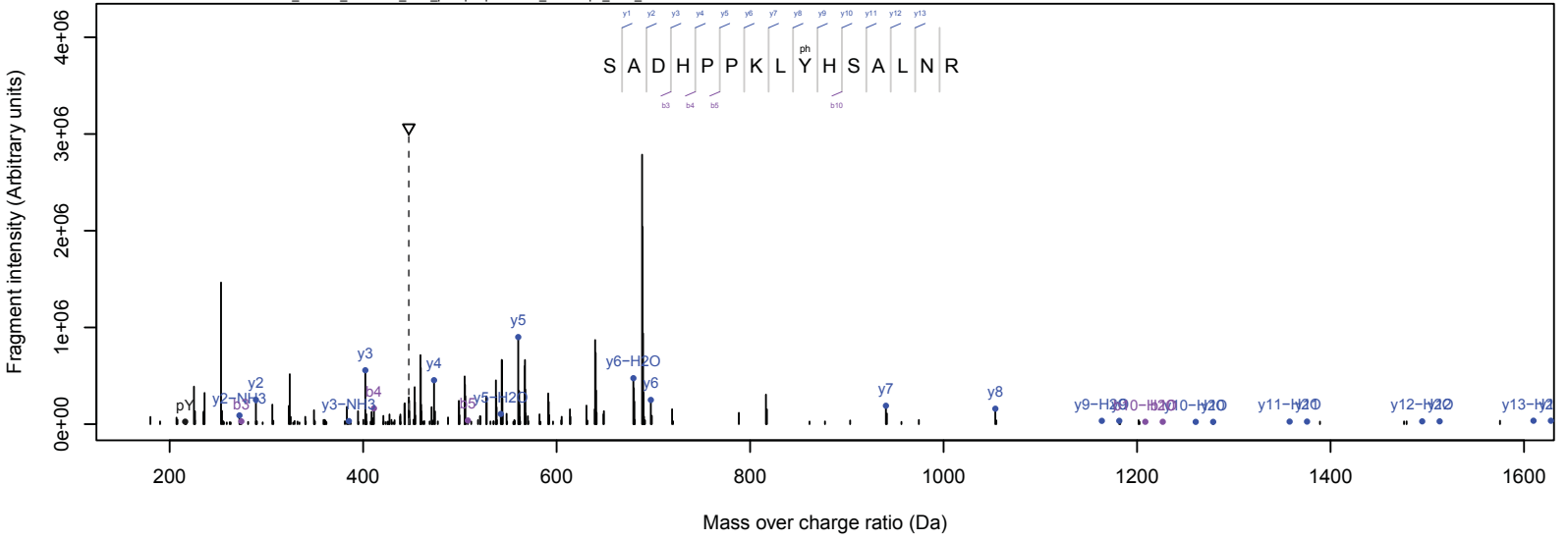
Andromeda score : 129.43



SADHPPKLY(ph)HSALNR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9663

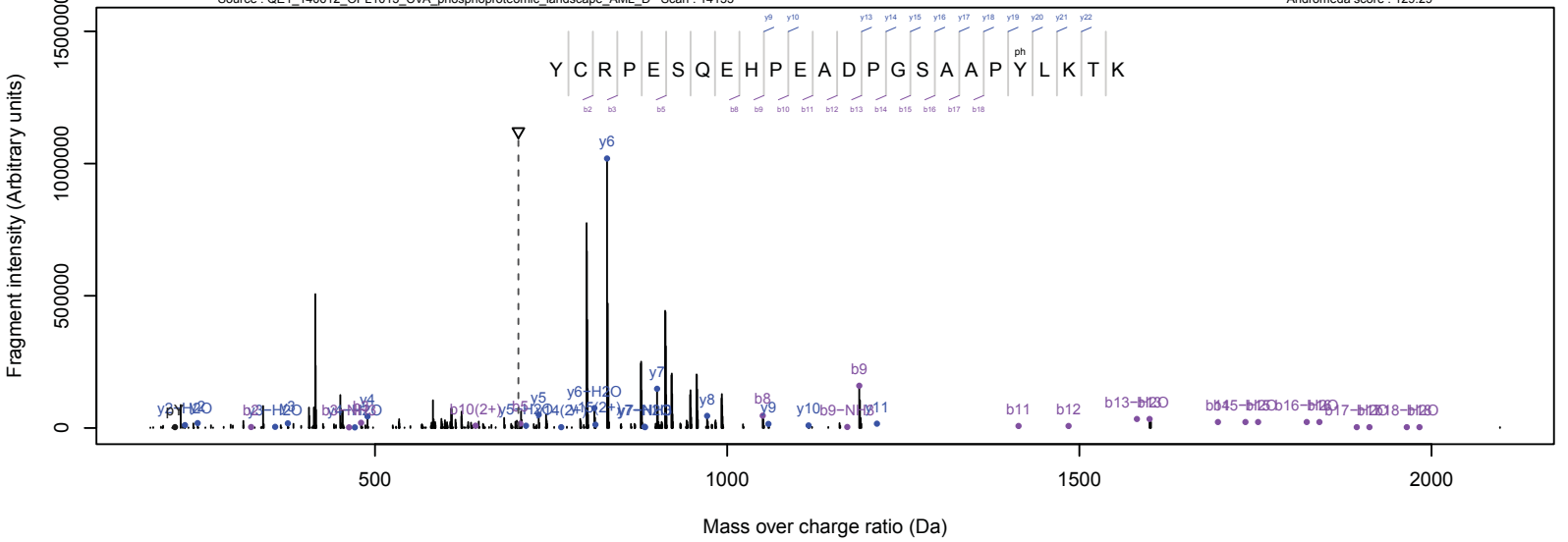
Andromeda score : 129.38



YCRPESQEHPGSAAPY(ph)LKTK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14155

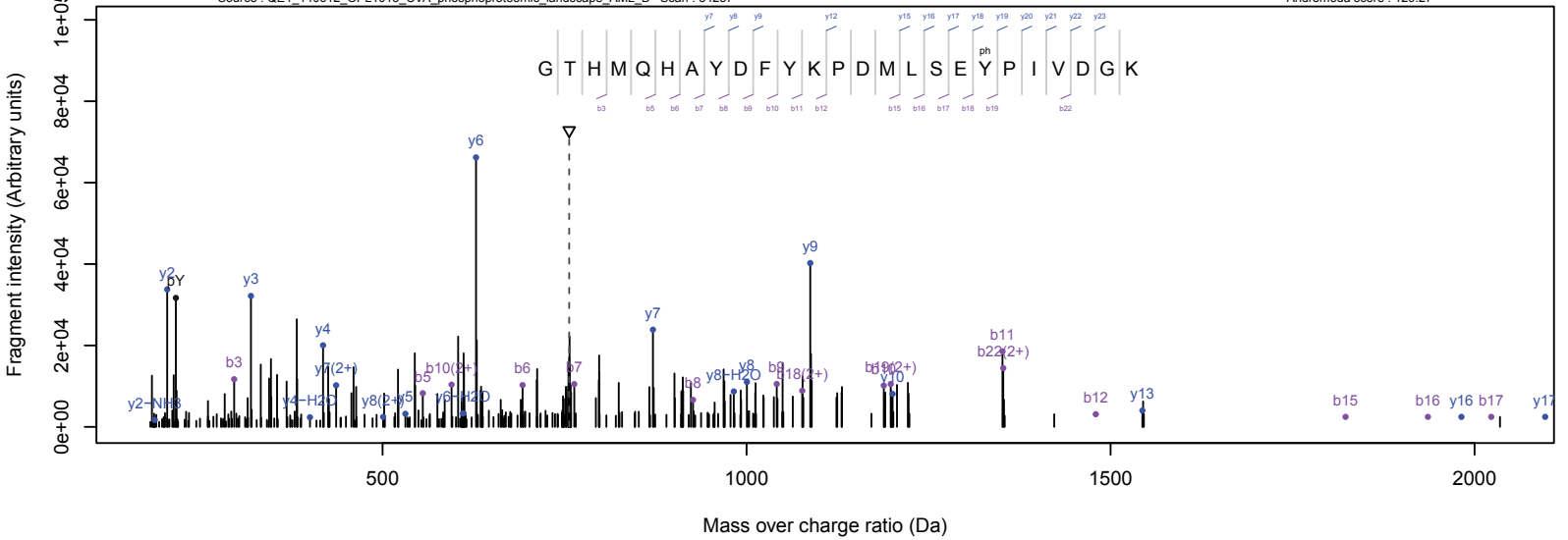
Andromeda score : 129.29



GTHMQHAYDFYKPDMLSEYP IVDGK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34257

Andromeda score : 129.27

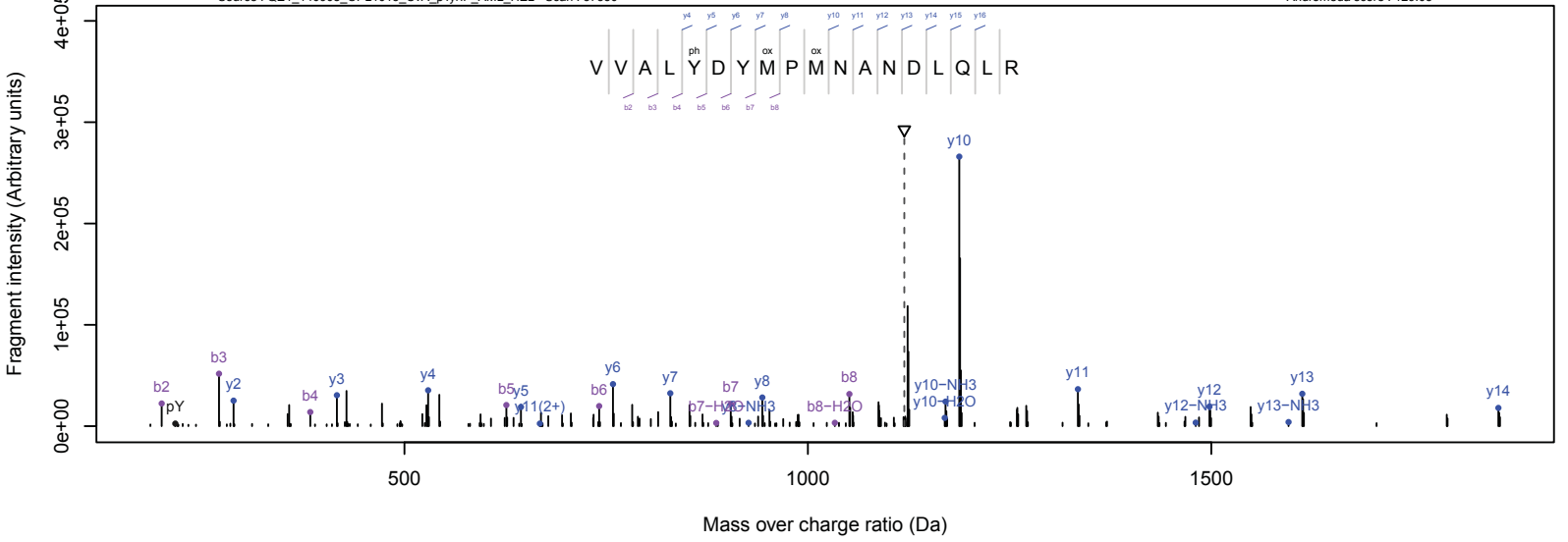




VVALY(ph)DYM(ox)PM(ox)NANDLQLR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37350

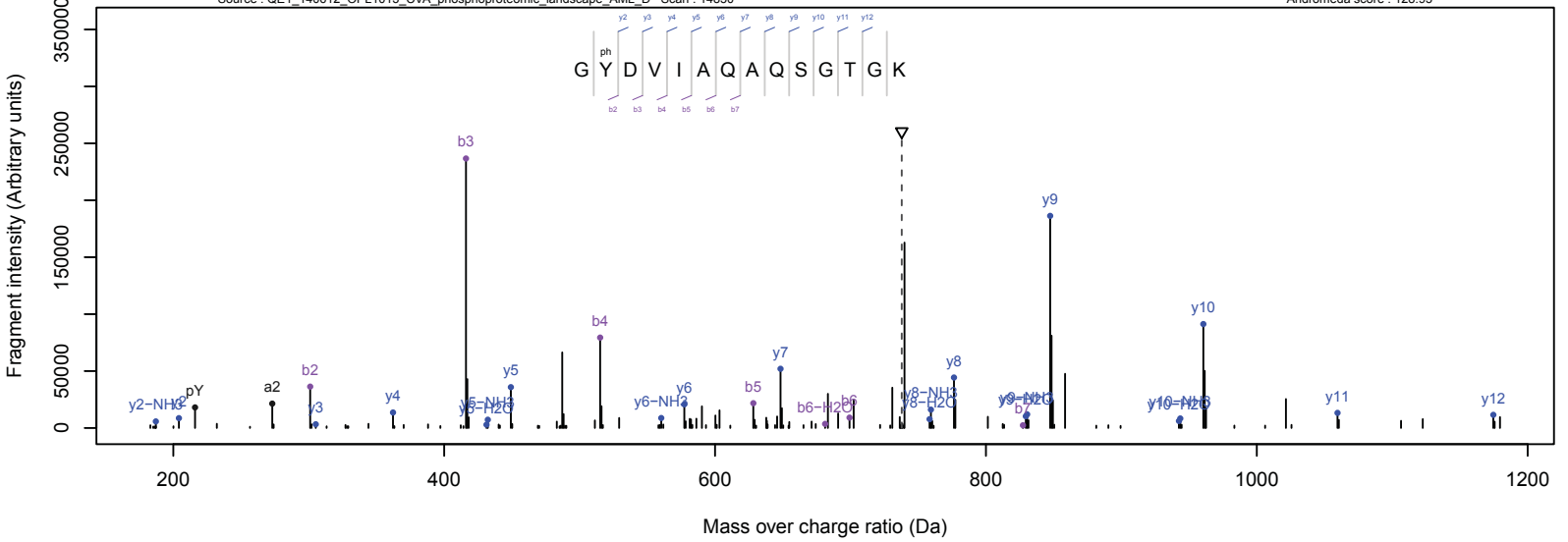
Andromeda score : 129.05



GY(ph)DVIAQAQSGTGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14850

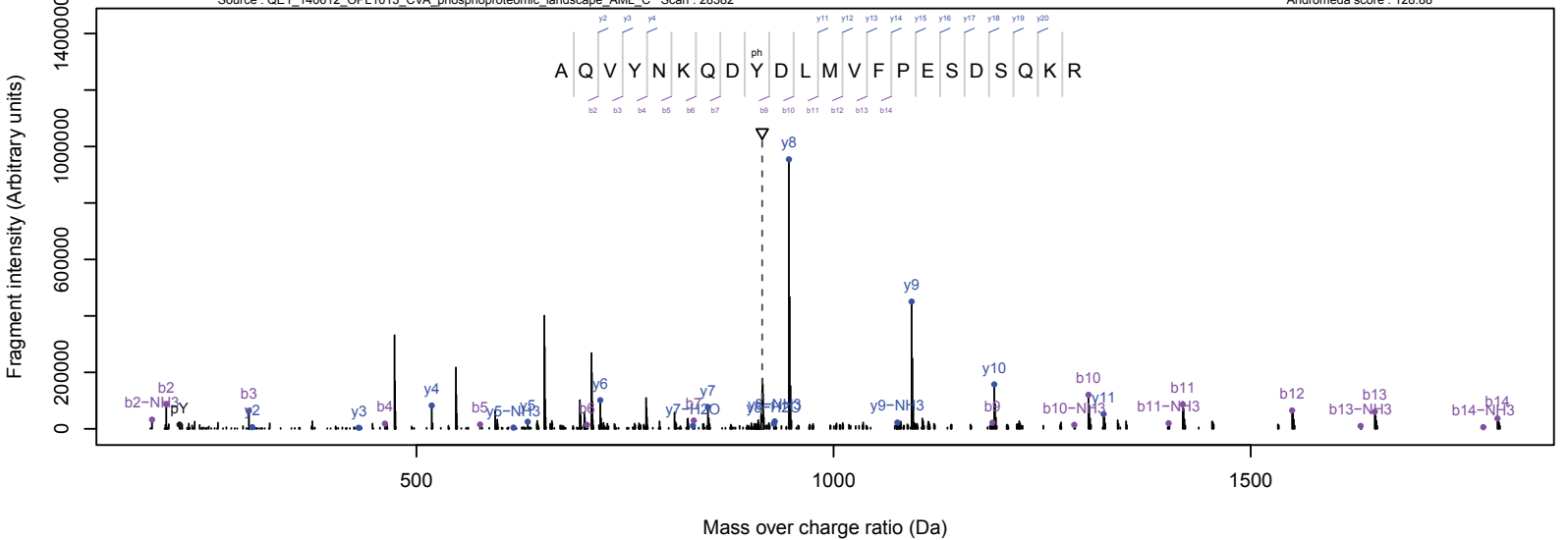
Andromeda score : 128.95



AQVYNKQDY(ph)DLMVFPESDSQKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28382

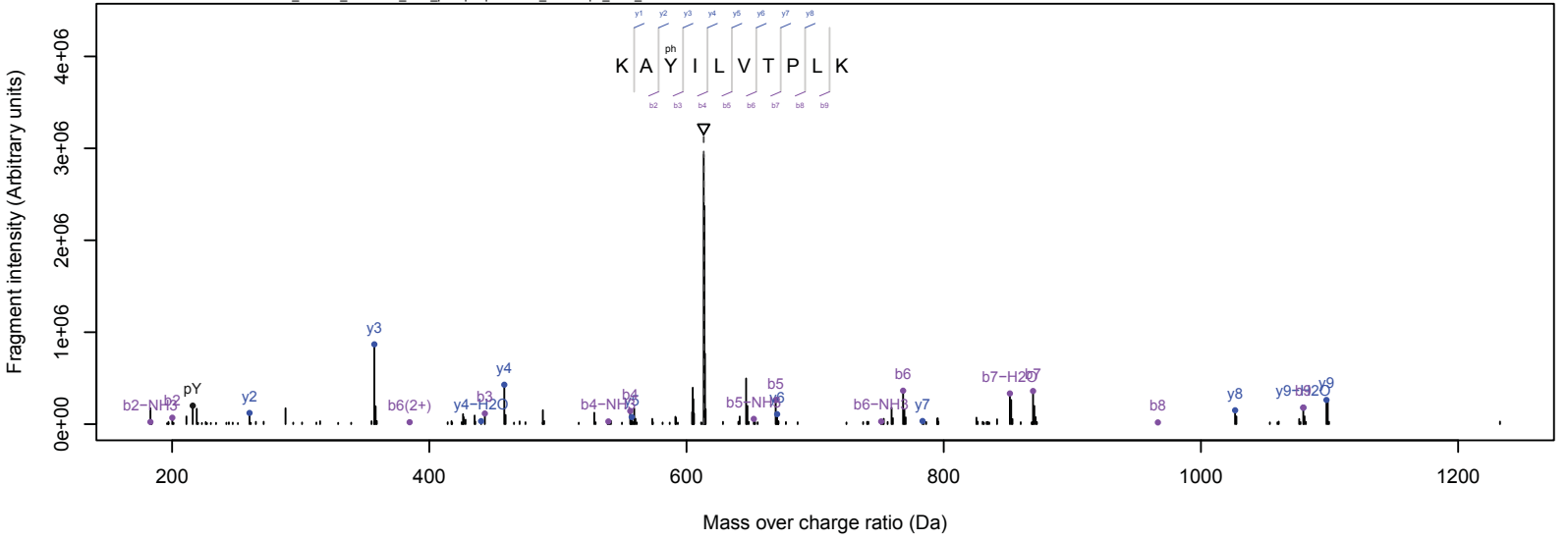
Andromeda score : 128.88



**KAY(ph)ILVTPLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27220

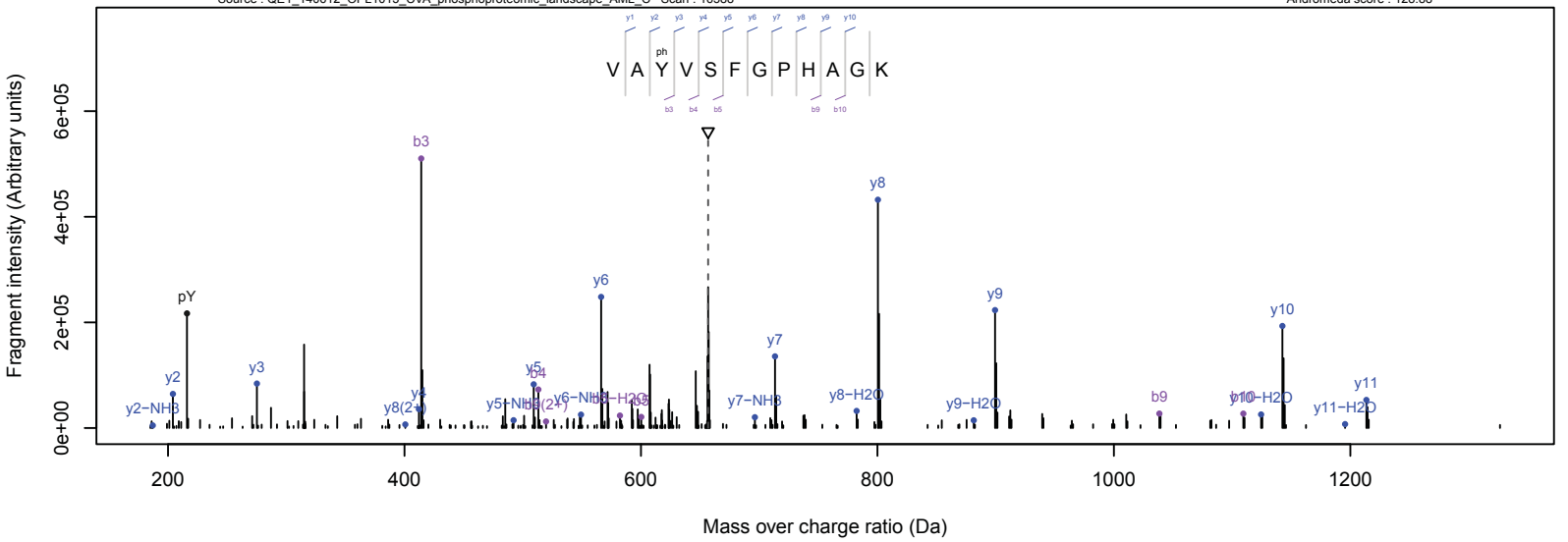
Andromeda score : 128.88



**VAY(ph)VSFGPHAGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16588

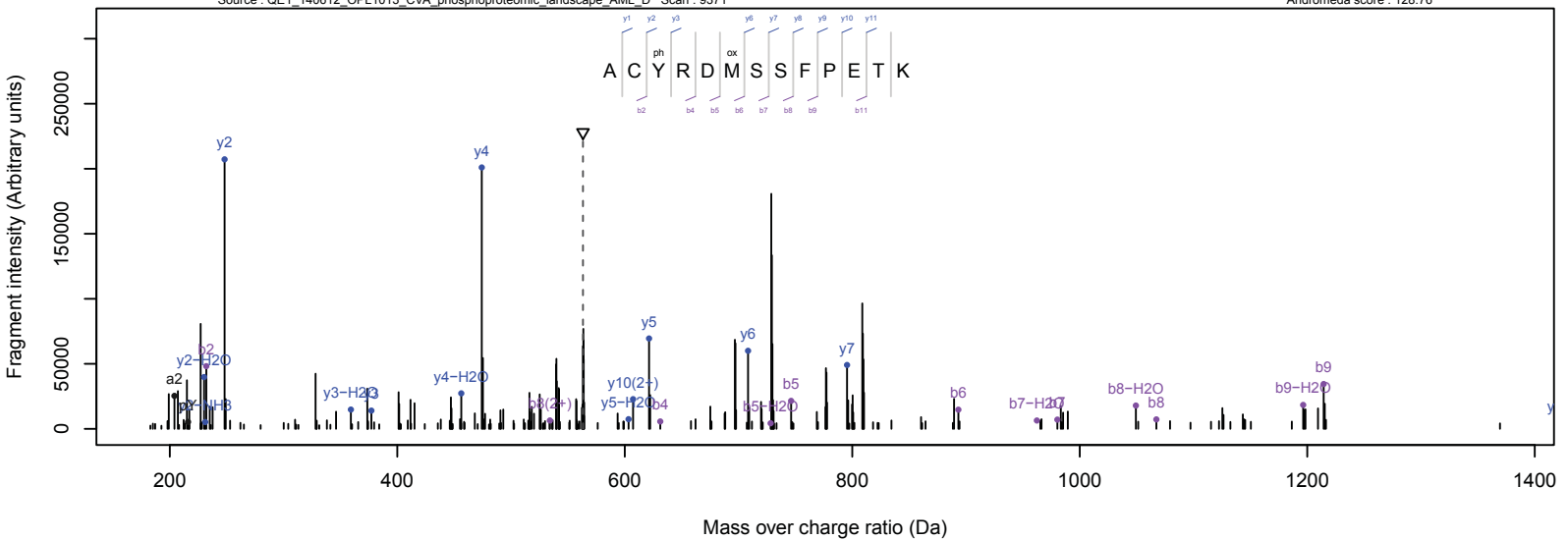
Andromeda score : 128.88



**ACY(ph)RDM(ox)SSFPETK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9371

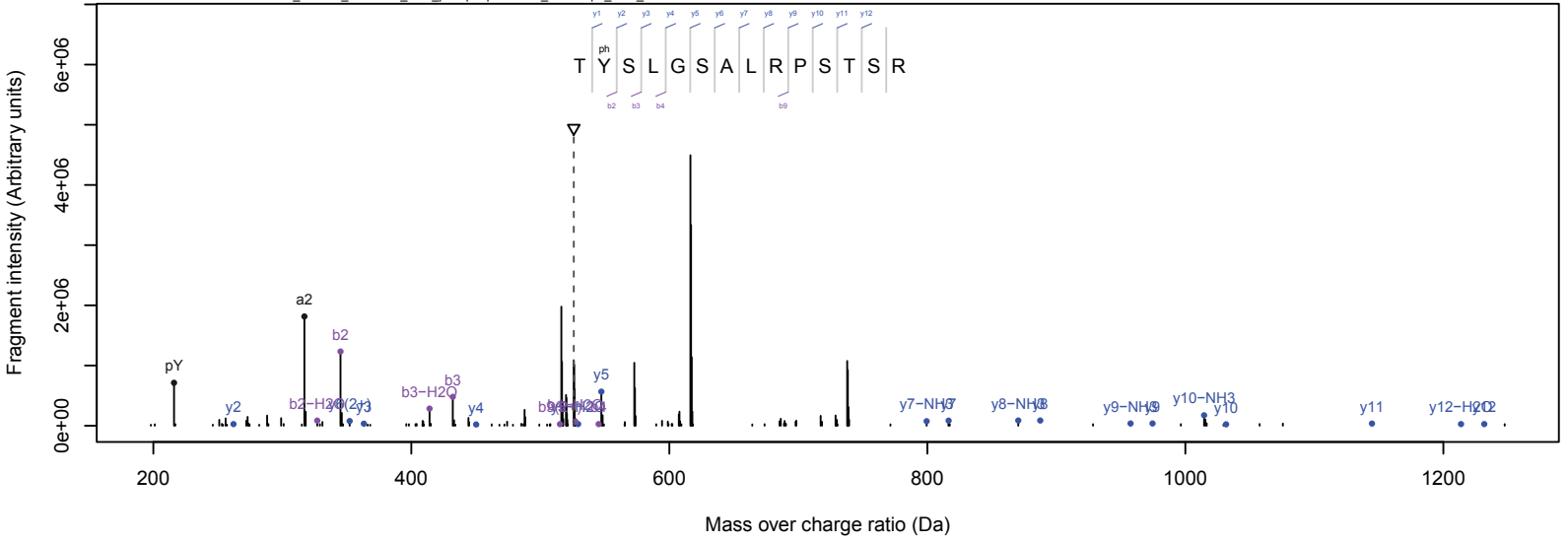
Andromeda score : 128.76



TY(ph)SLGSALRPSTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17445

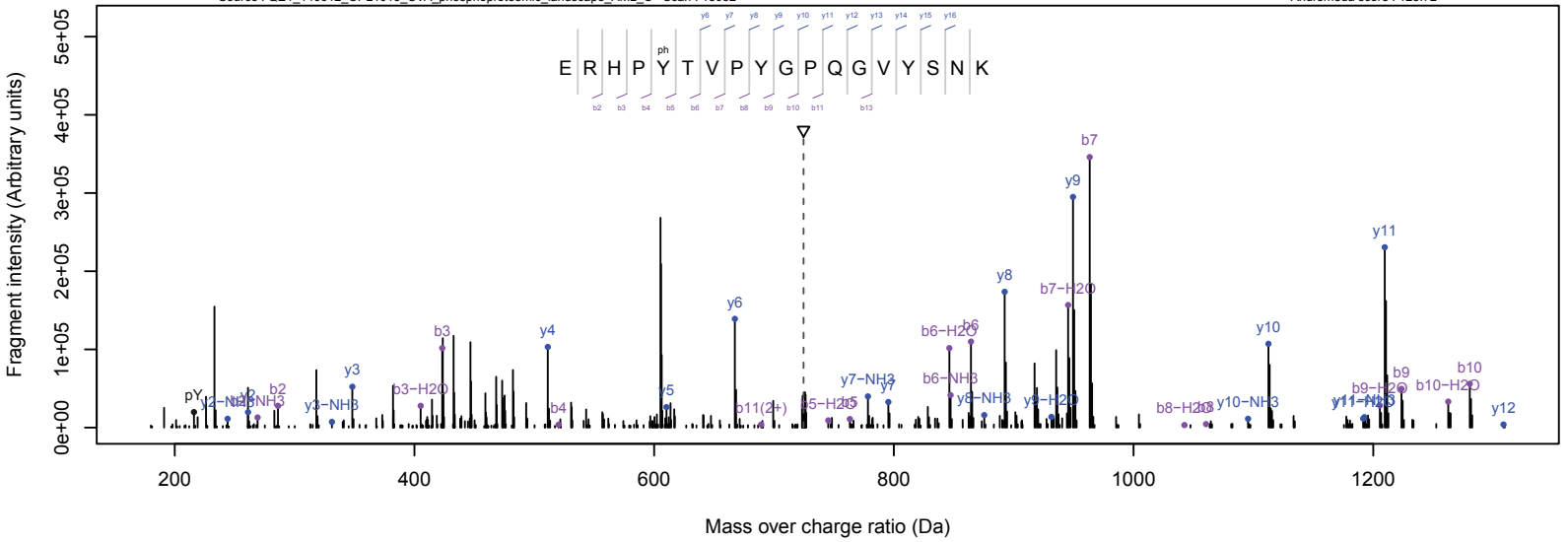
Andromeda score : 128.76



ERHPY(ph)TVPYGPGVYSNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18052

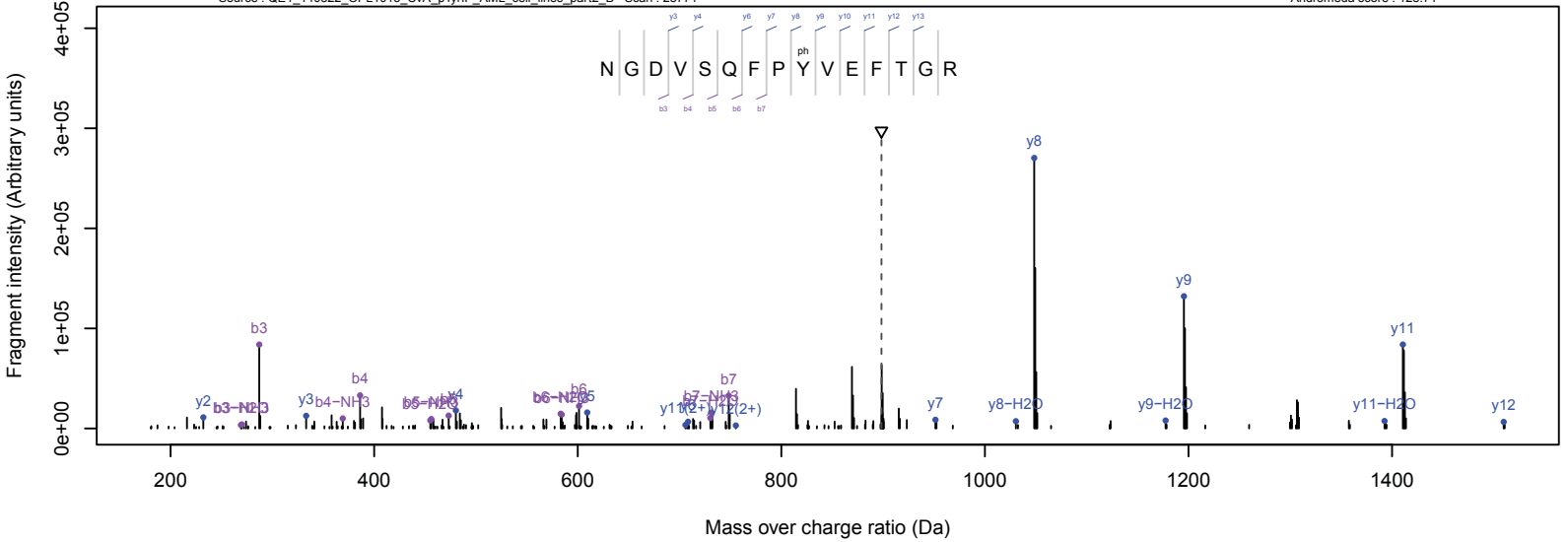
Andromeda score : 128.72



NGDVSQFPY(ph)VEFTGR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 28771

Andromeda score : 128.71



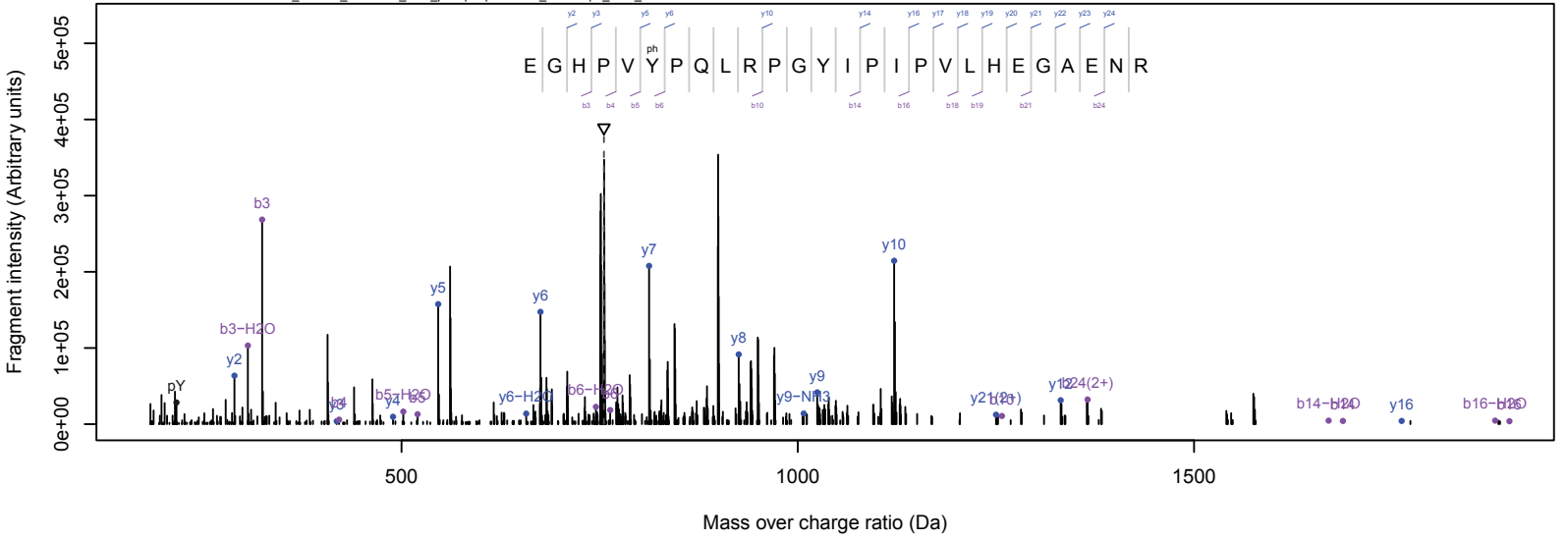




EGHPVY(ph)PQLRPGYIPIPLHEGAENR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38728

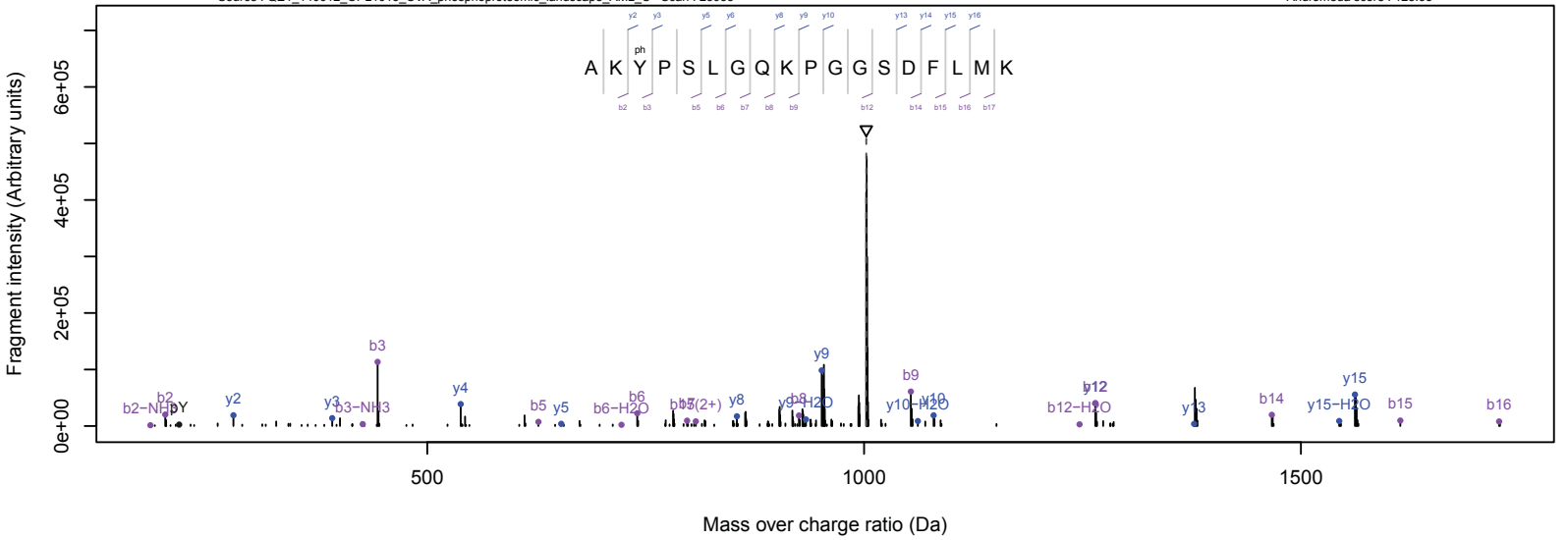
Andromeda score : 128.4



AKY(ph)PSLGQKPGGSDFLMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23935

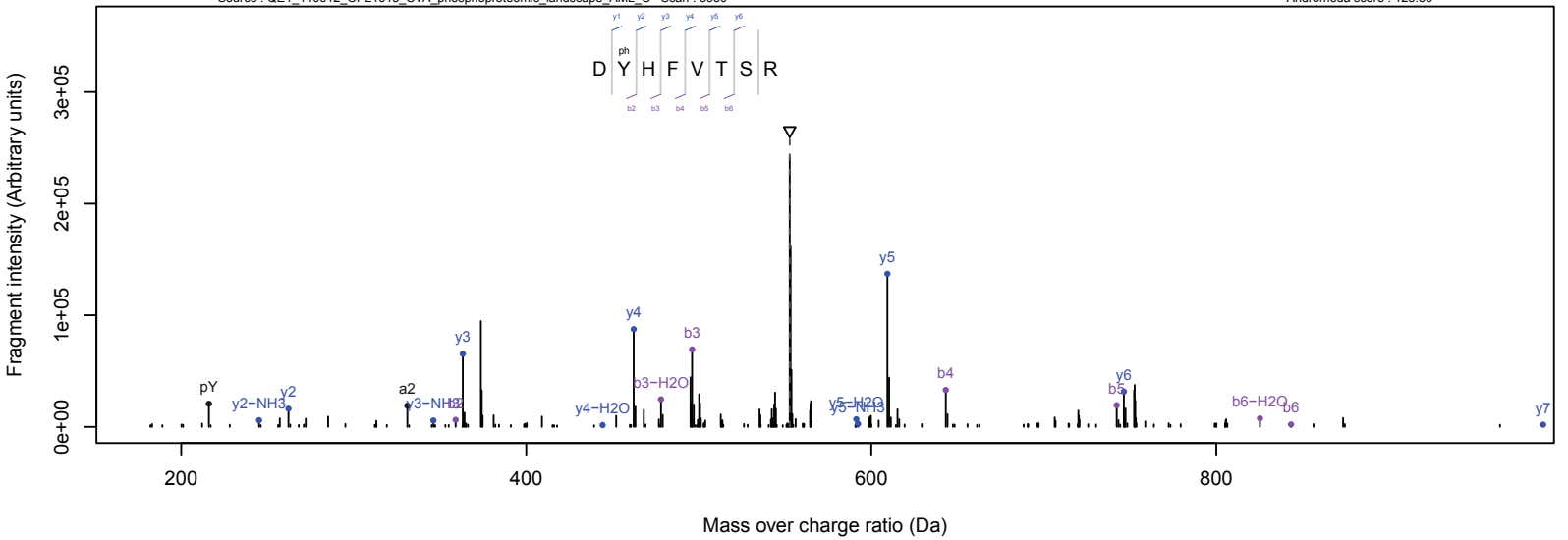
Andromeda score : 128.38



DY(ph)HFVTSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8930

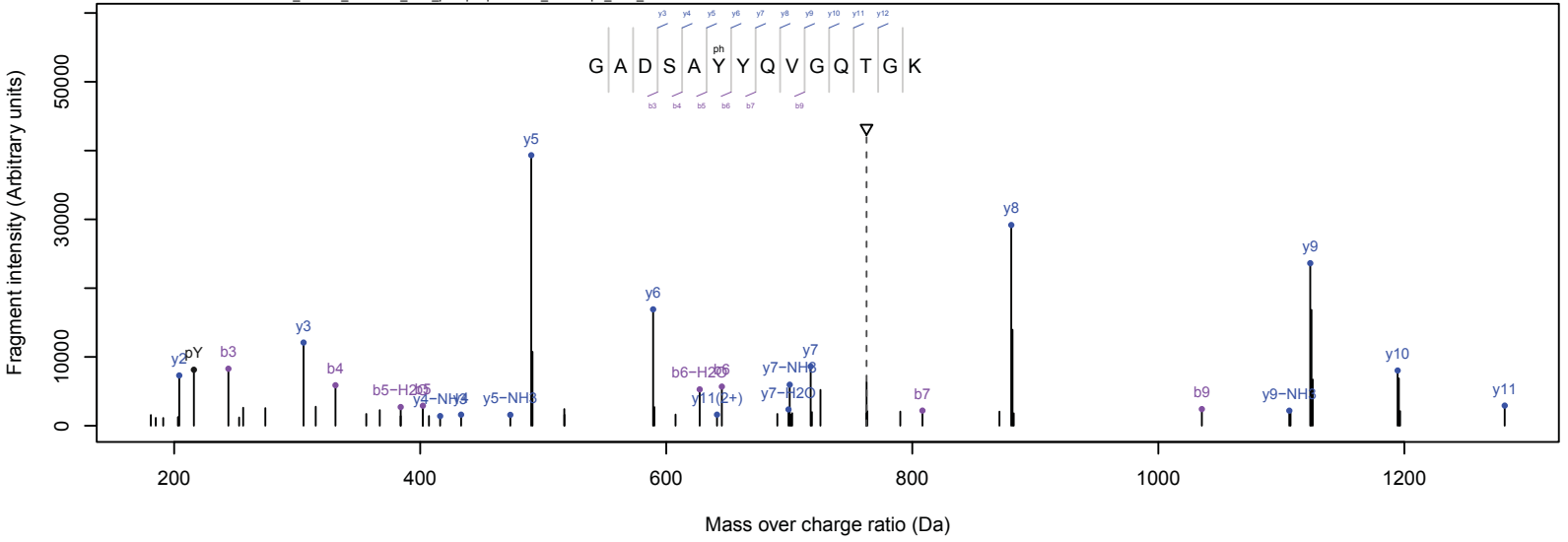
Andromeda score : 128.36



GADSA<sup>ph</sup>YQVGQTGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9241

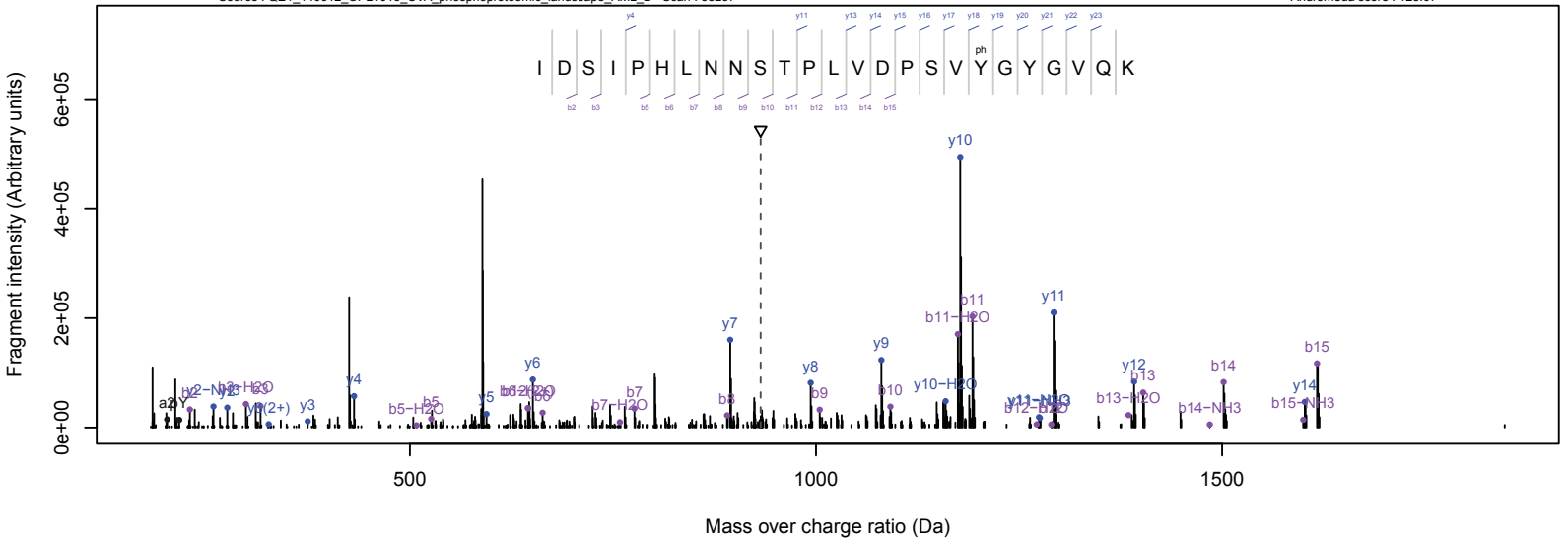
Andromeda score : 128.22



IDSIPLN<sup>ph</sup>NSTPLVDPSVY<sup>ph</sup>GYGVQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38237

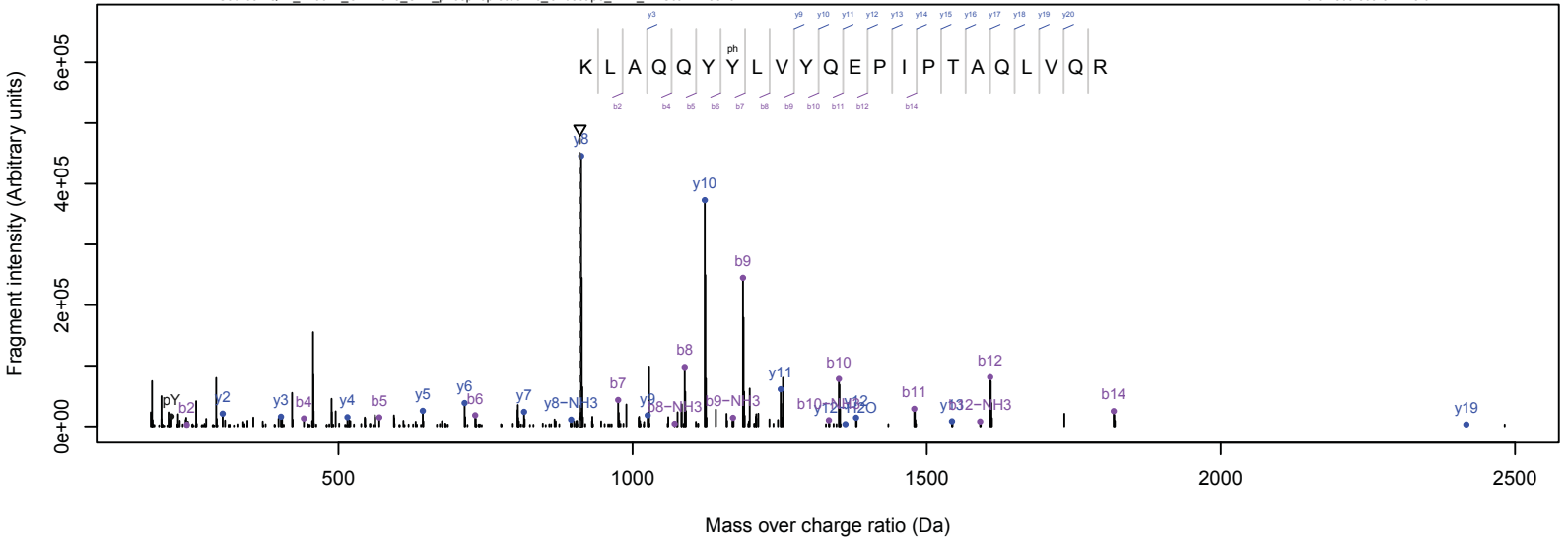
Andromeda score : 128.07



KLAQQY<sup>ph</sup>LVYQEPIPTAQLVQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40329

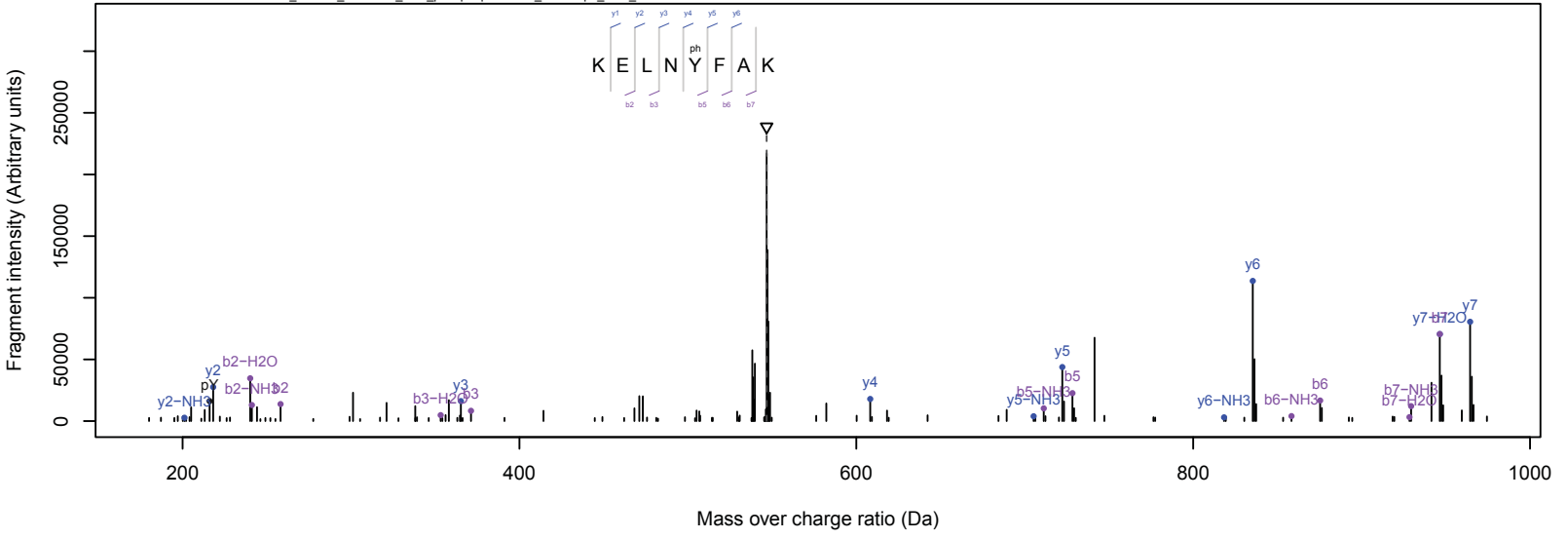
Andromeda score : 128.01



**KELNY(ph)FAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 10846

Andromeda score : 128

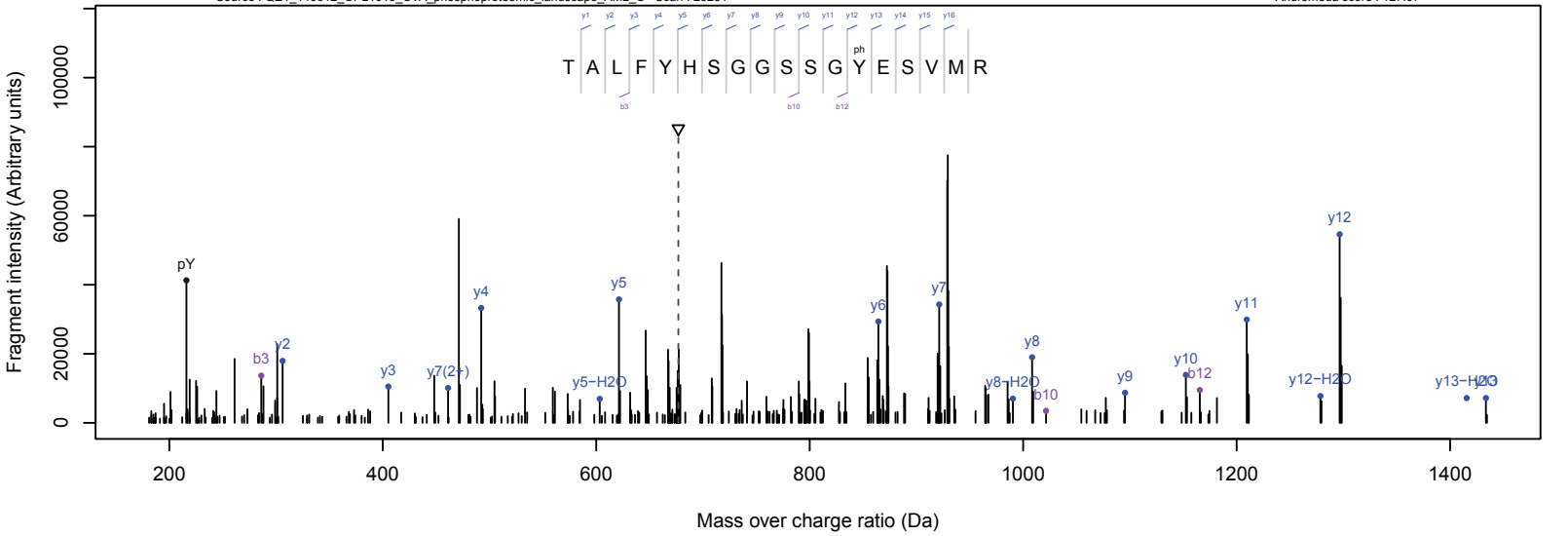


Mass over charge ratio (Da)

**TALFYHSGSSGY(ph)ESVMR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25281

Andromeda score : 127.97

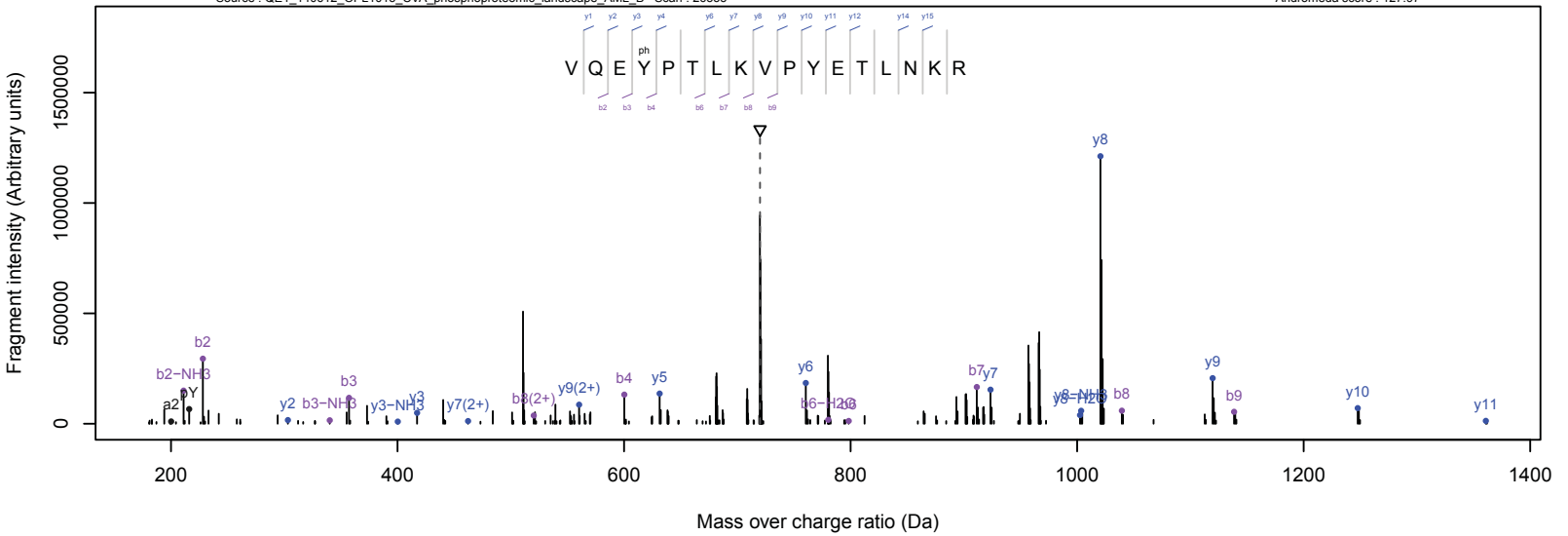


Mass over charge ratio (Da)

**VQEY(ph)PTLKVPYETLNKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26535

Andromeda score : 127.97

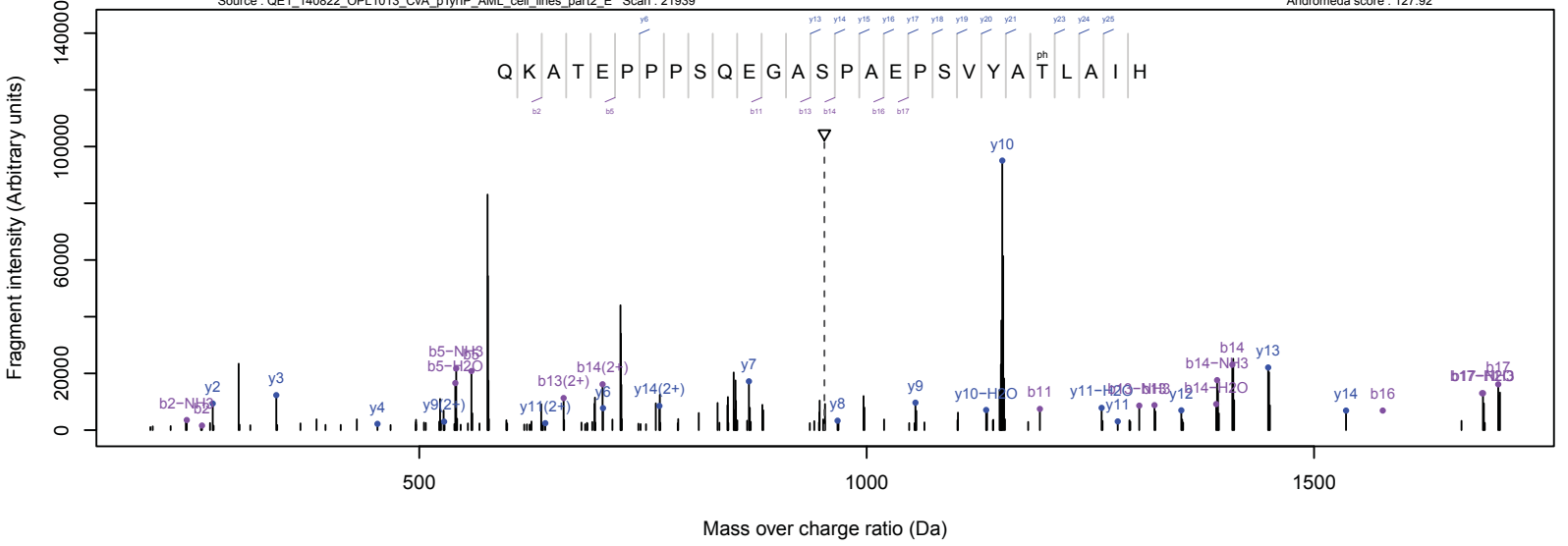


Mass over charge ratio (Da)

**QKATEPPPSQEGASPAEPSVYATLAIH-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 21939

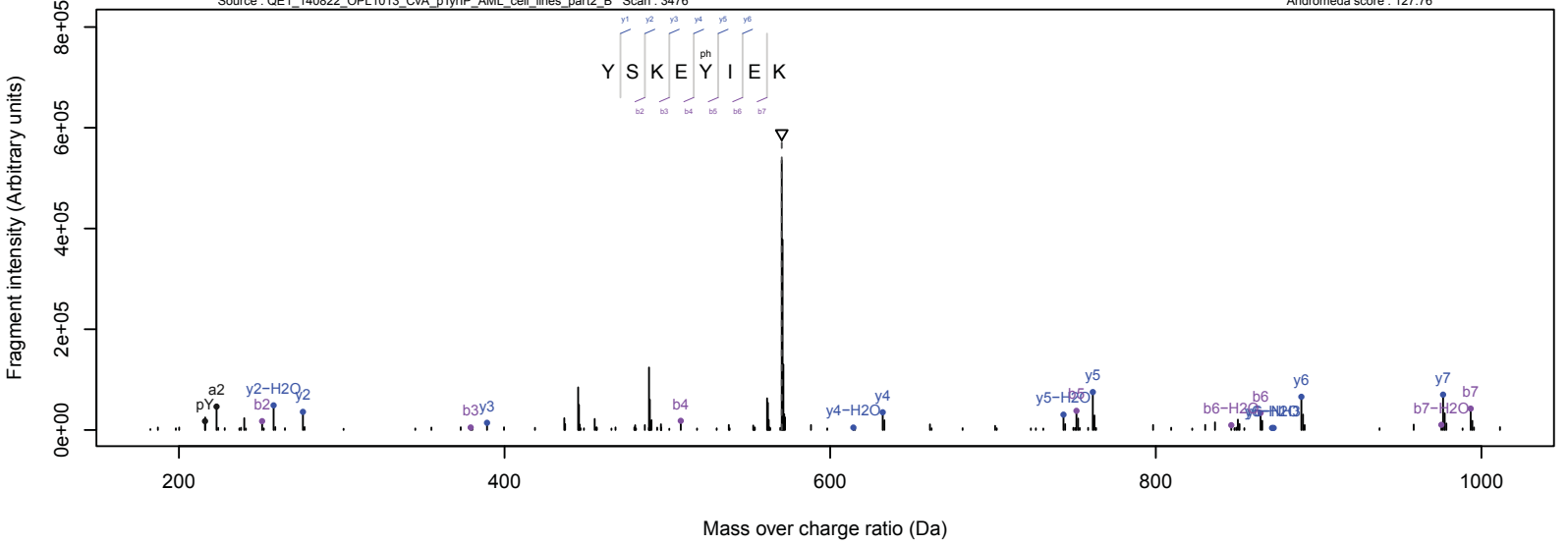
Andromeda score : 127.92



**YSKEY(ph)IEK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3476

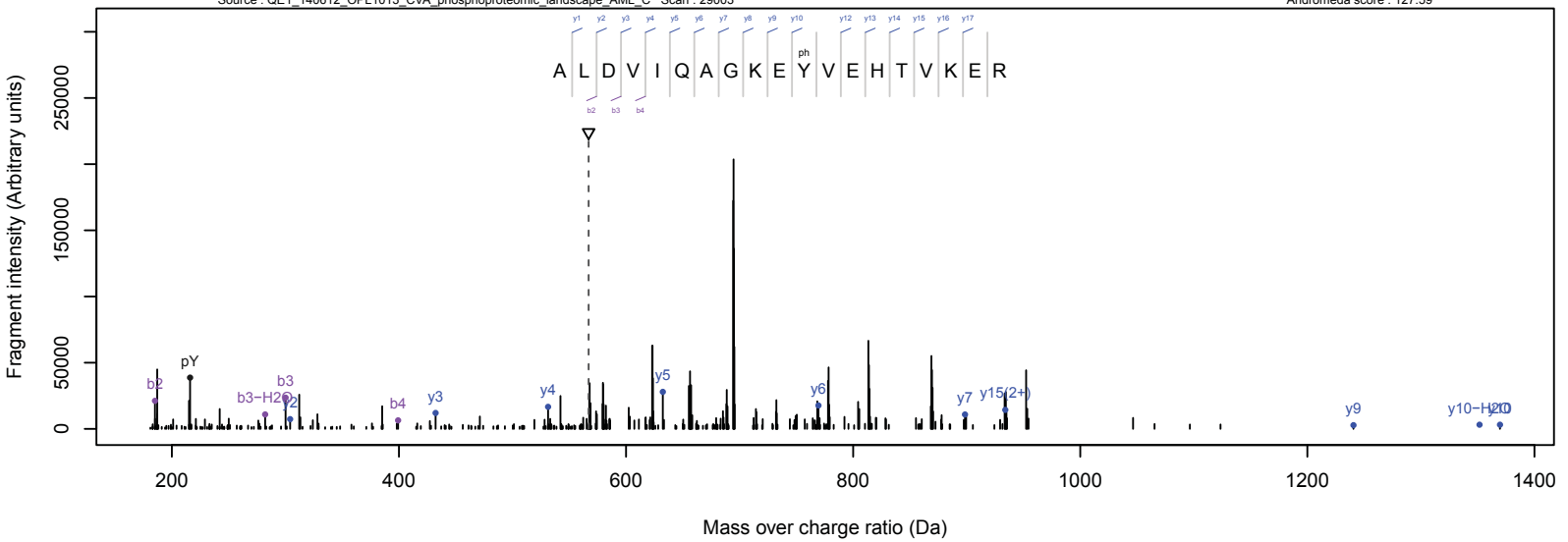
Andromeda score : 127.76



**ALDVIQAGKEY(ph)VEHTVKER-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29003

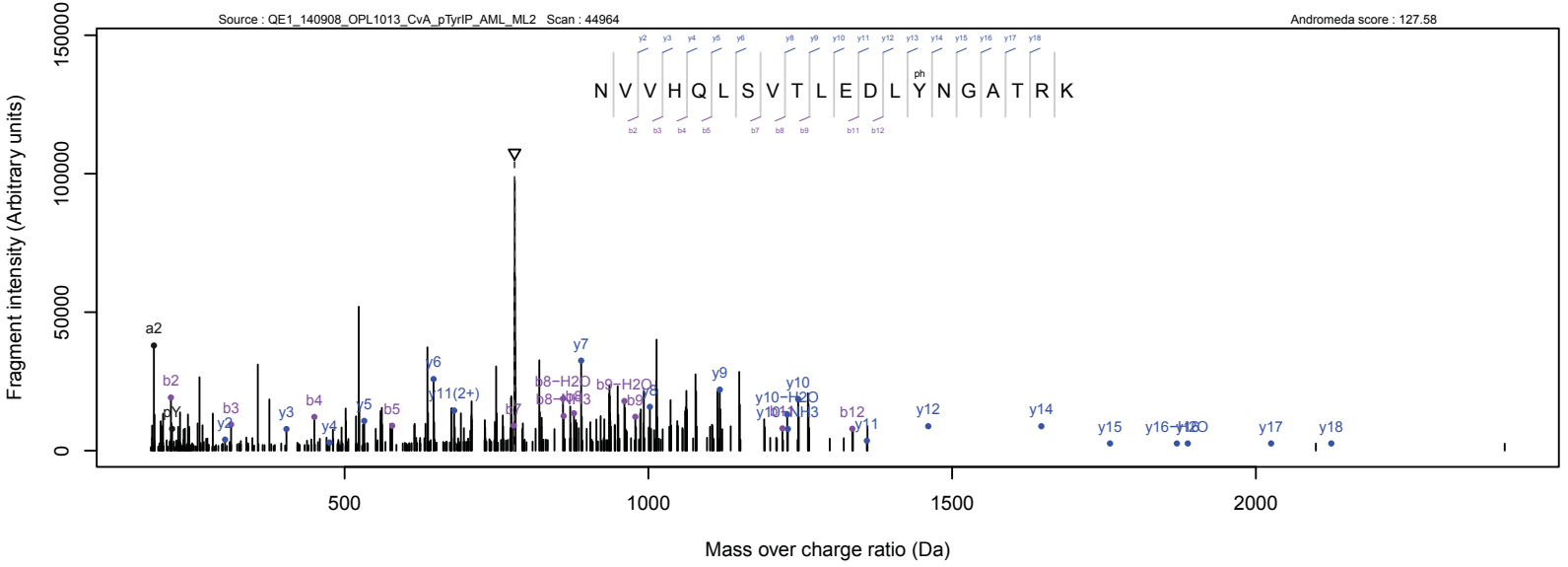
Andromeda score : 127.59



NVVHQLSVTLLEDLY(ph)NGATRK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 44964

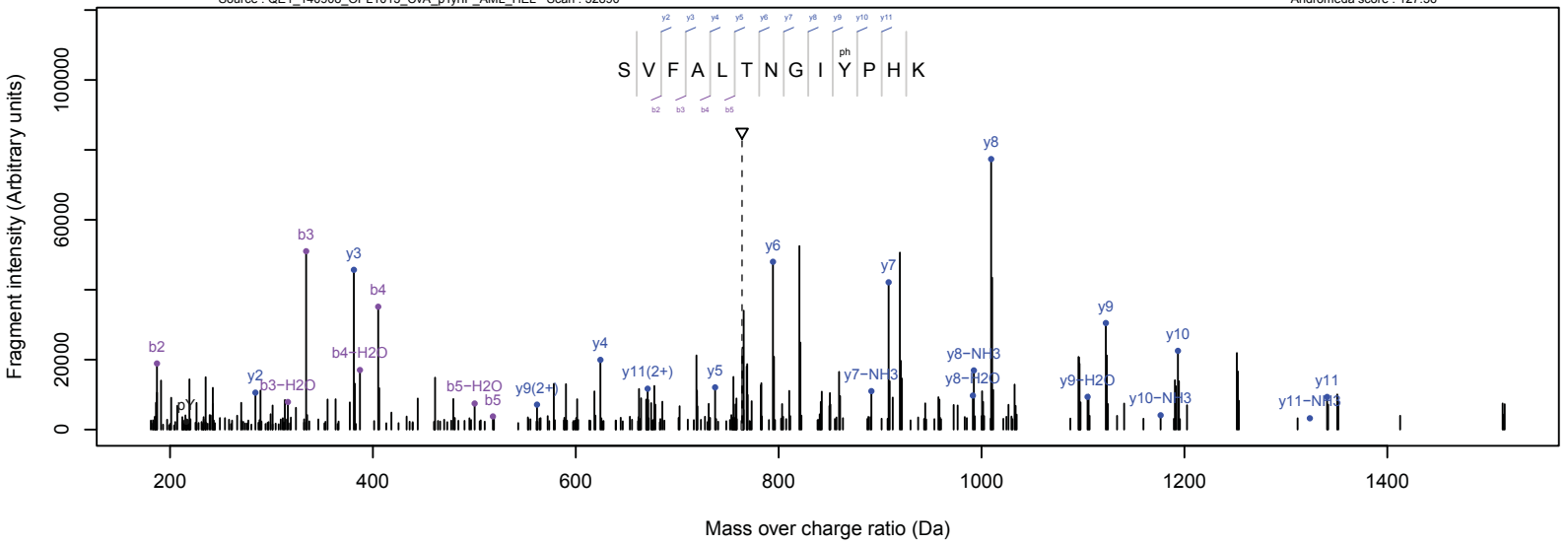
Andromeda score : 127.58



SVFALTNGIY(ph)PHK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32890

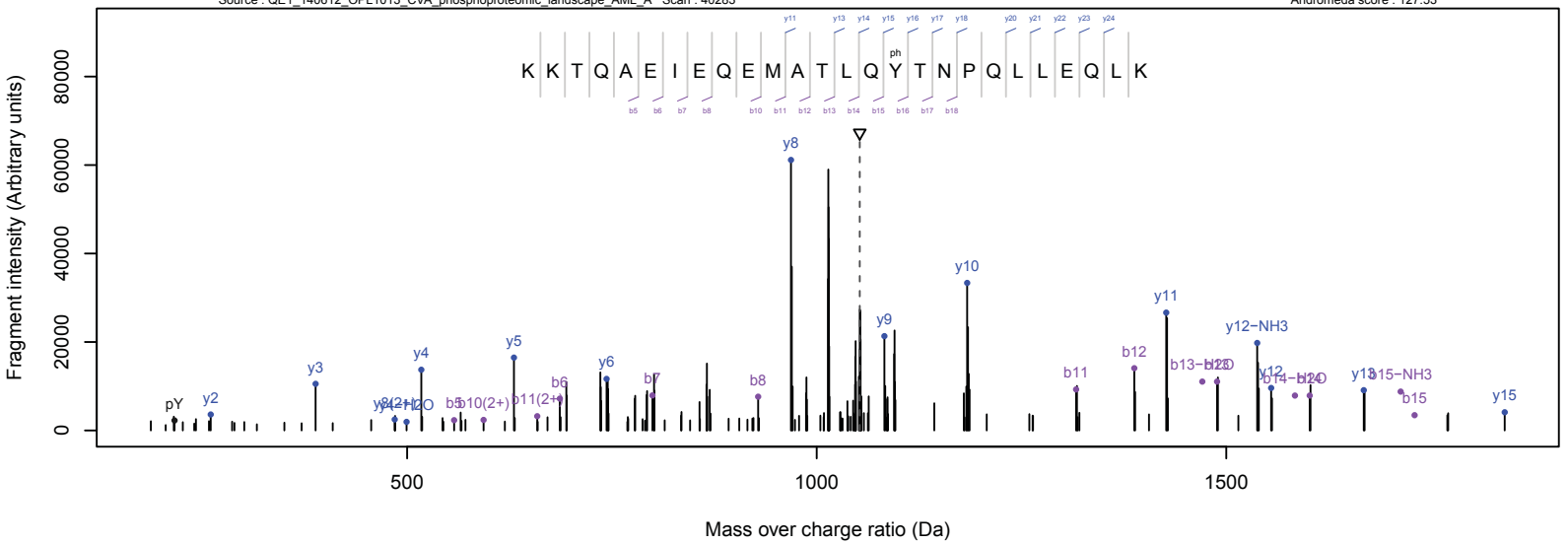
Andromeda score : 127.56



KKTQAEIEQEMATLQY(ph)TNPQLLEQLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 40283

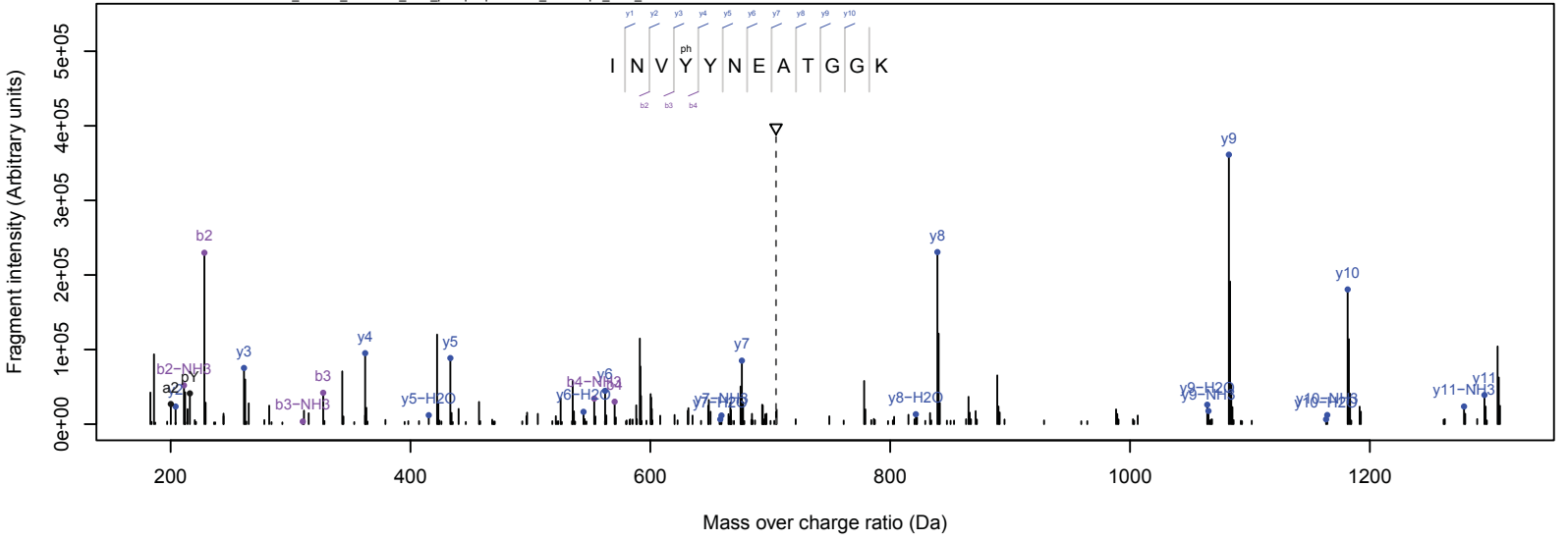
Andromeda score : 127.53



INVY(ph)YNEATGGK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15448

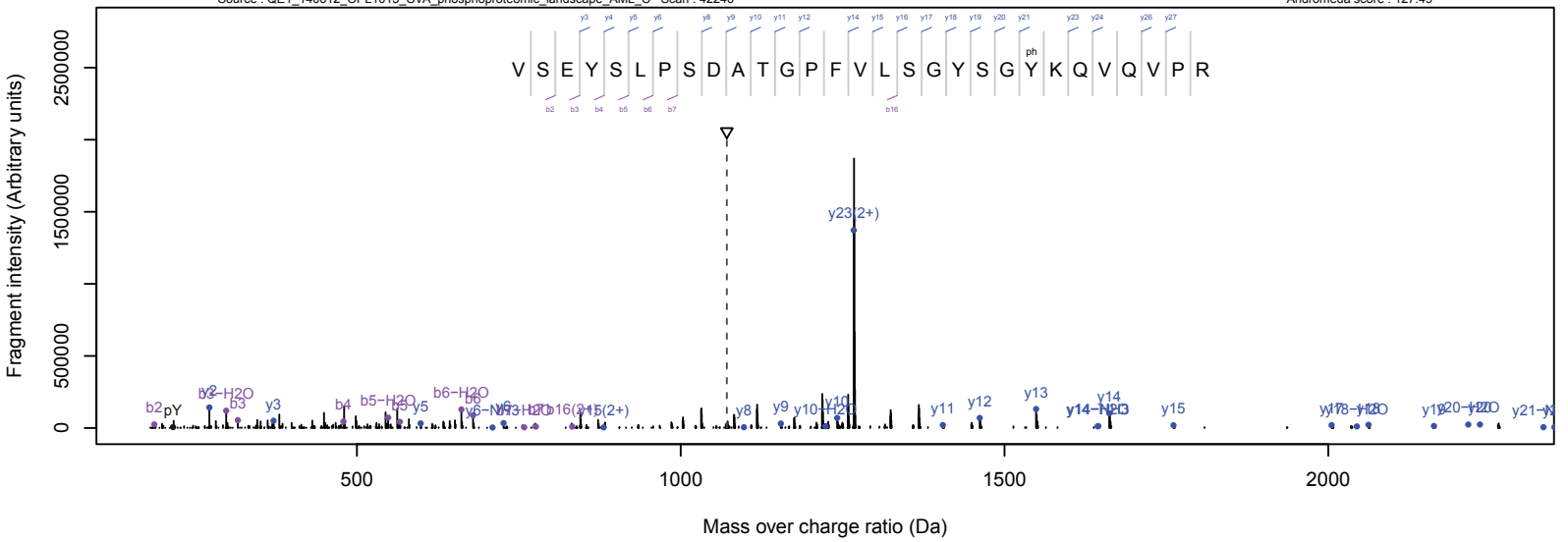
Andromeda score: 127.52



VSEYSLPSDATGPFVLSGYSYKQVQVPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 42246

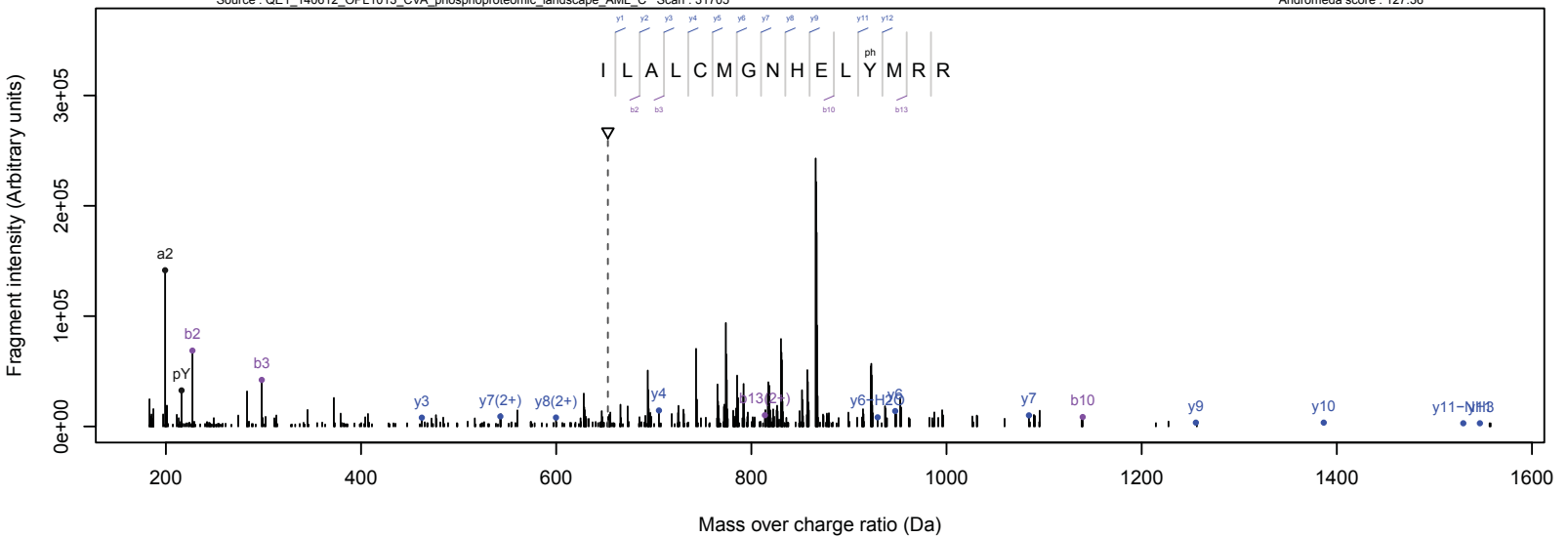
Andromeda score: 127.49



ILALCMGNHELYMRR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31705

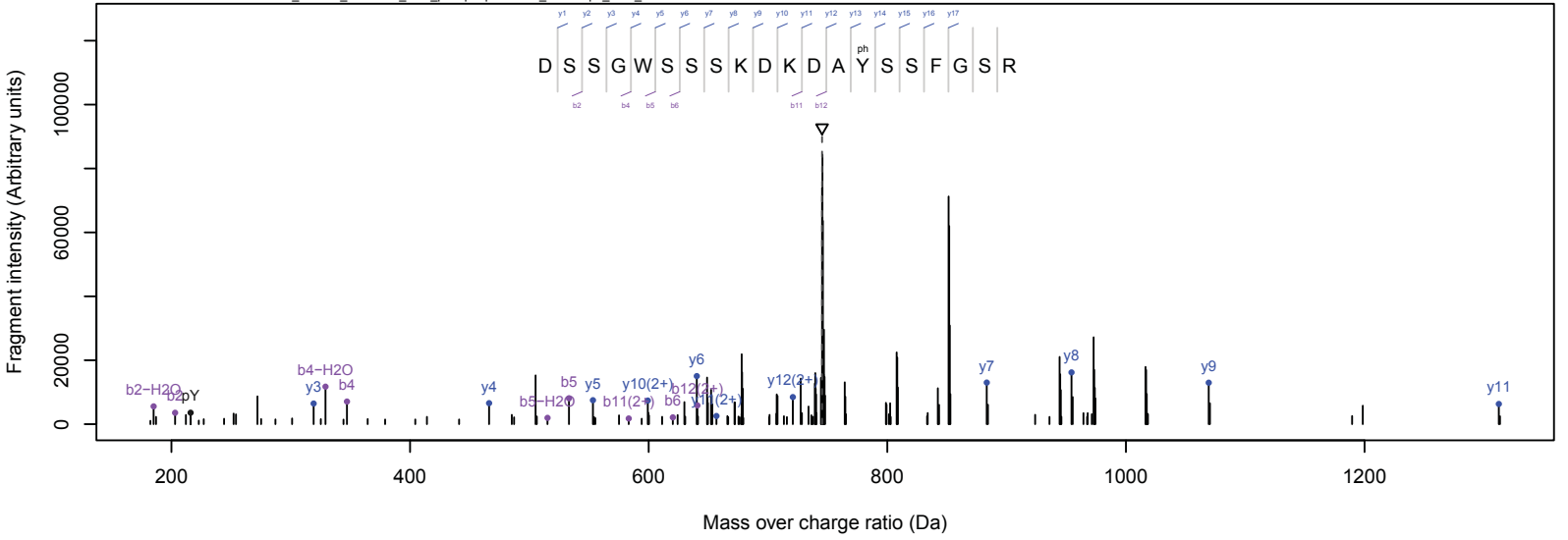
Andromeda score: 127.36



DSSGWSSSKDKDAY(ph)SSFGSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12527

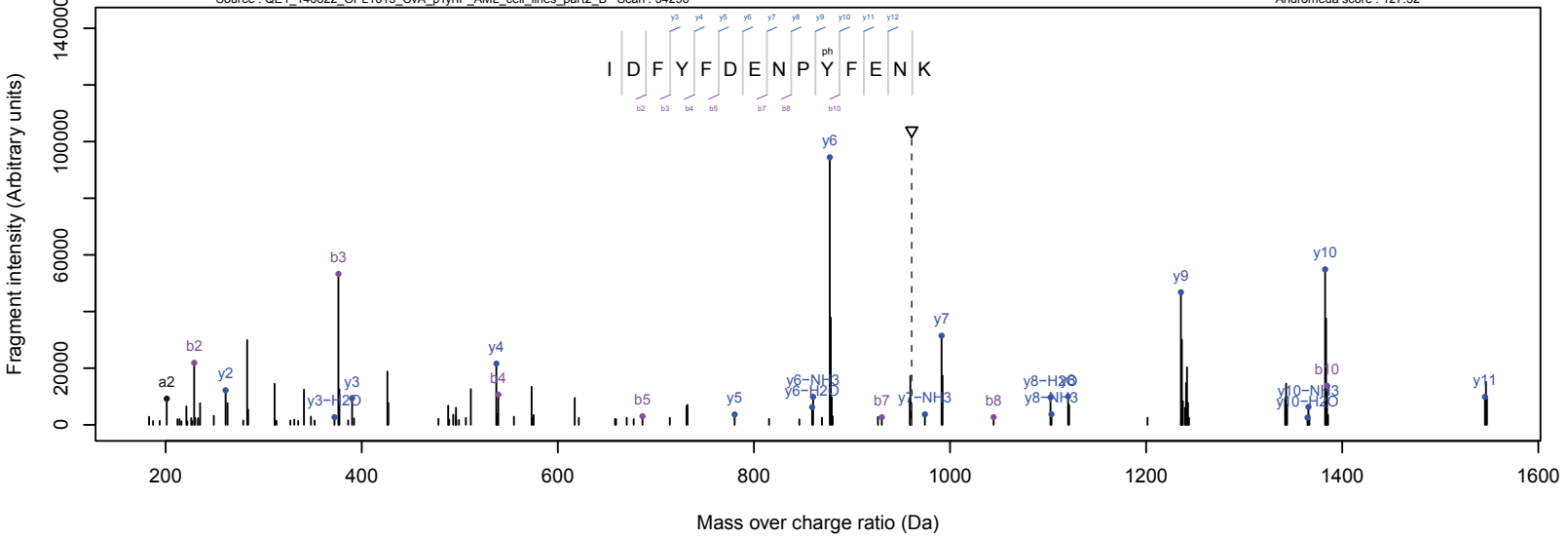
Andromeda score : 127.32



IDFYFDENPY(ph)FENK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 34296

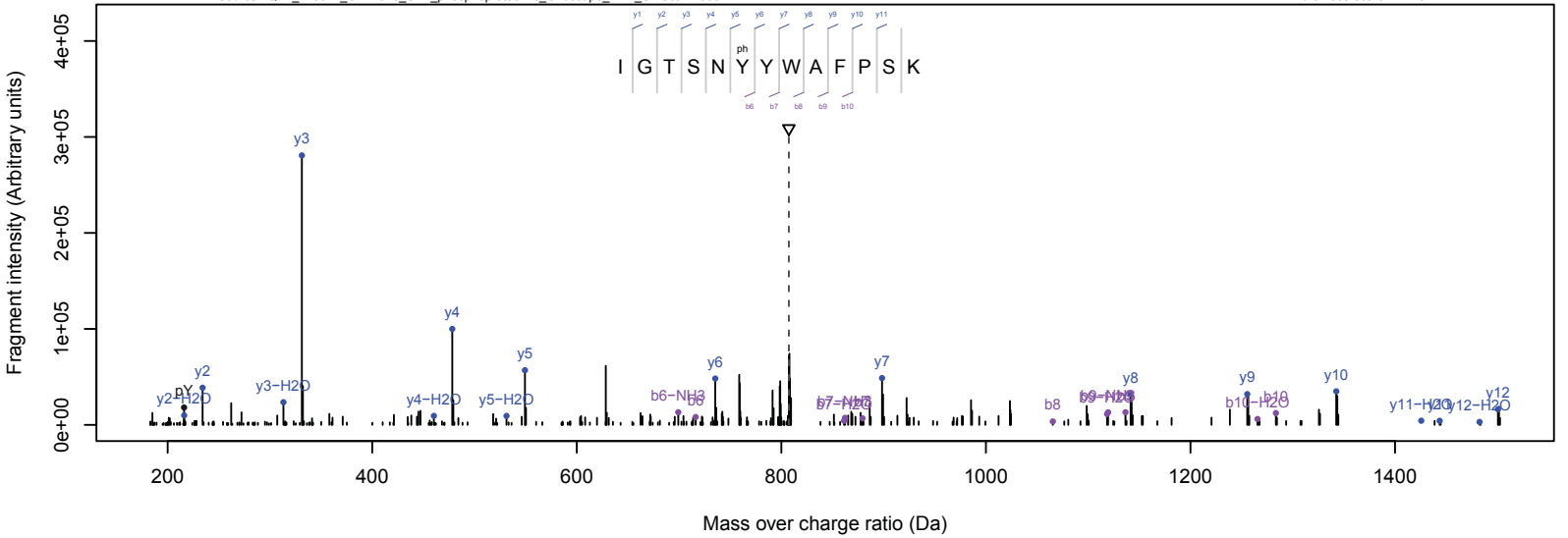
Andromeda score : 127.32



IGTSNY(ph)YWAFPSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36314

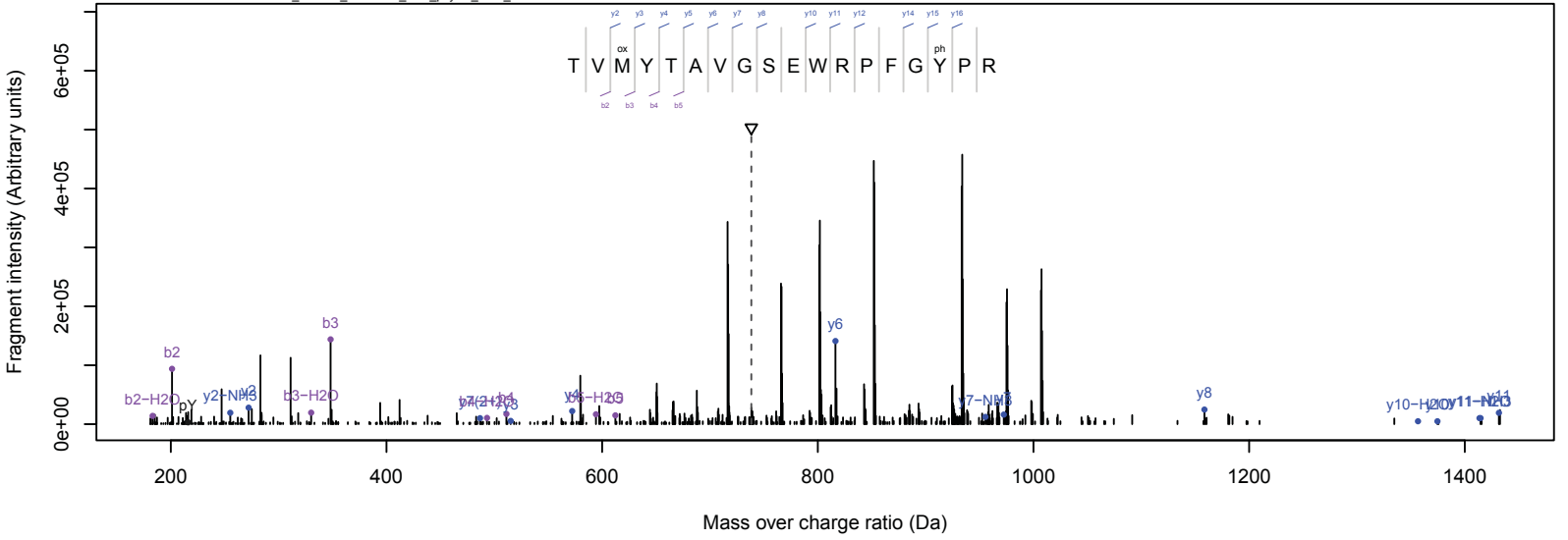
Andromeda score : 127.32



TVM(ox)YTAGSEWRPFY(ph)PR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan : 39015

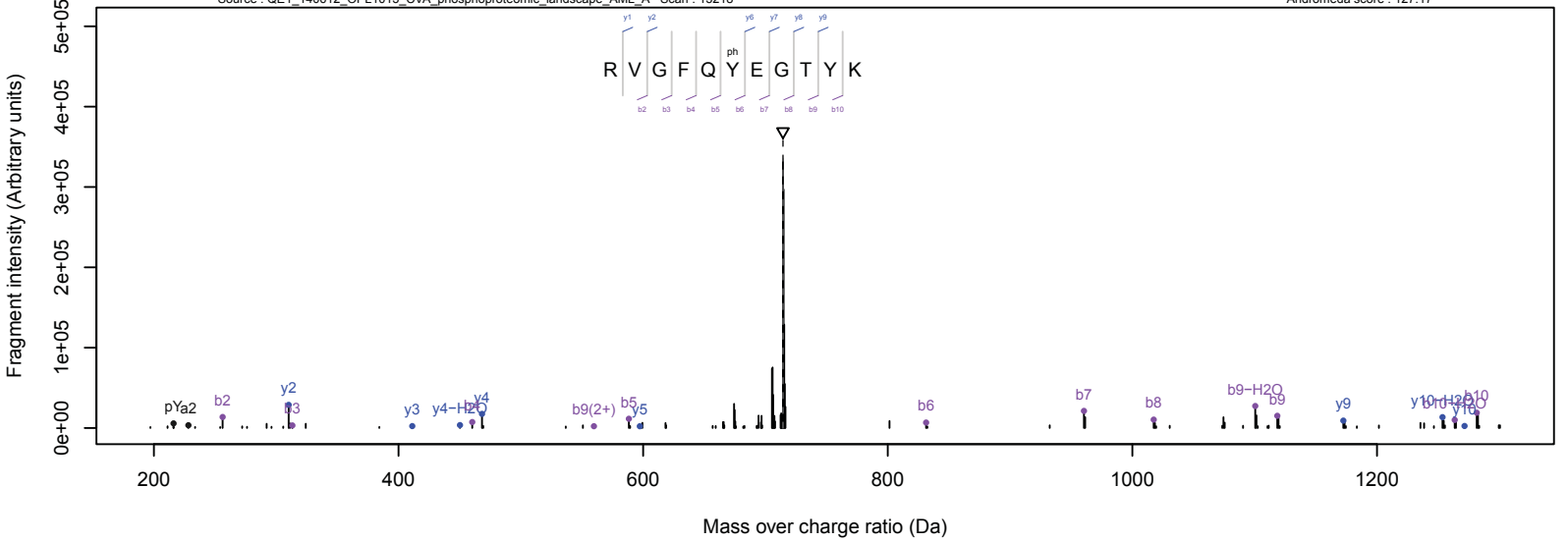
Andromeda score : 127.3



RVGFQY(ph)EGTYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 13218

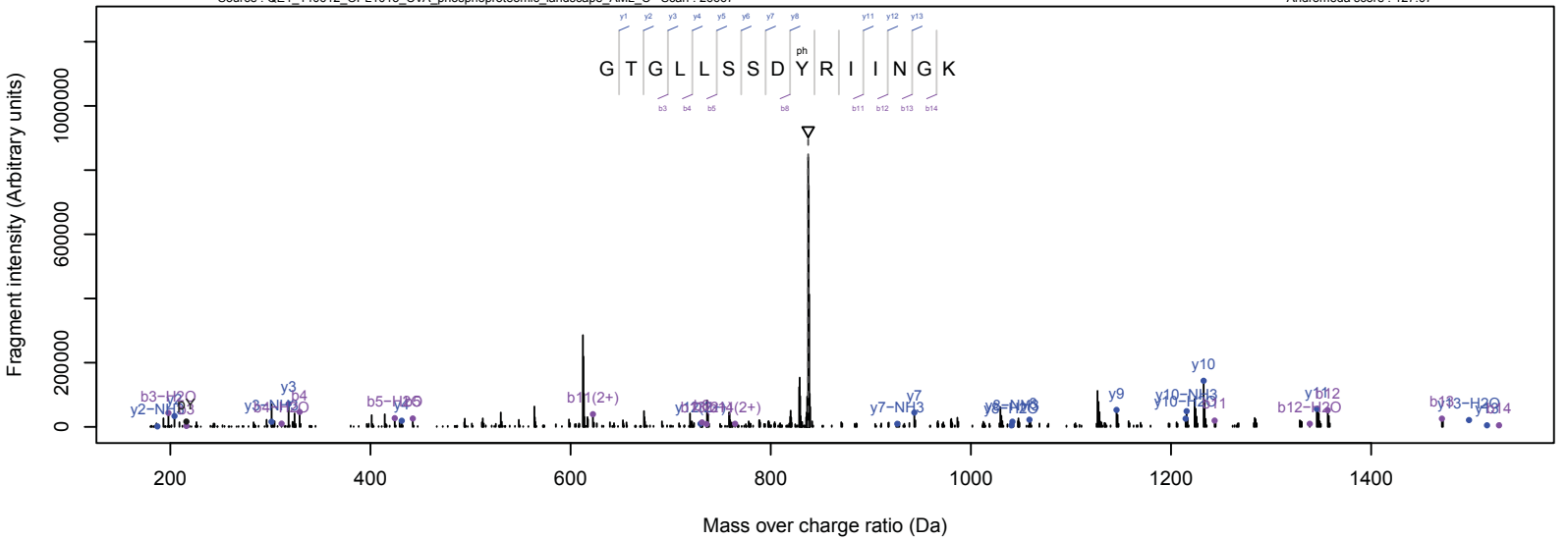
Andromeda score : 127.17



GTGLSSDY(ph)RIINGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26667

Andromeda score : 127.07

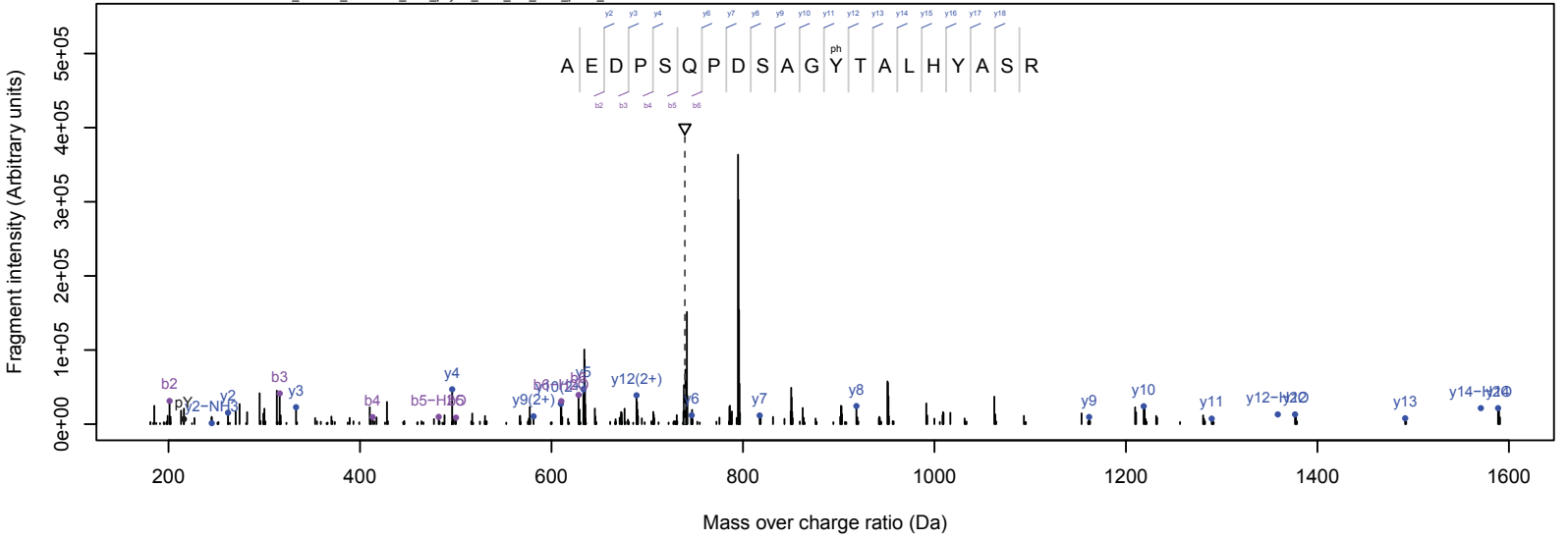




**AEDPSQPSAGY(ph)TALHYASR-3+**

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 15694

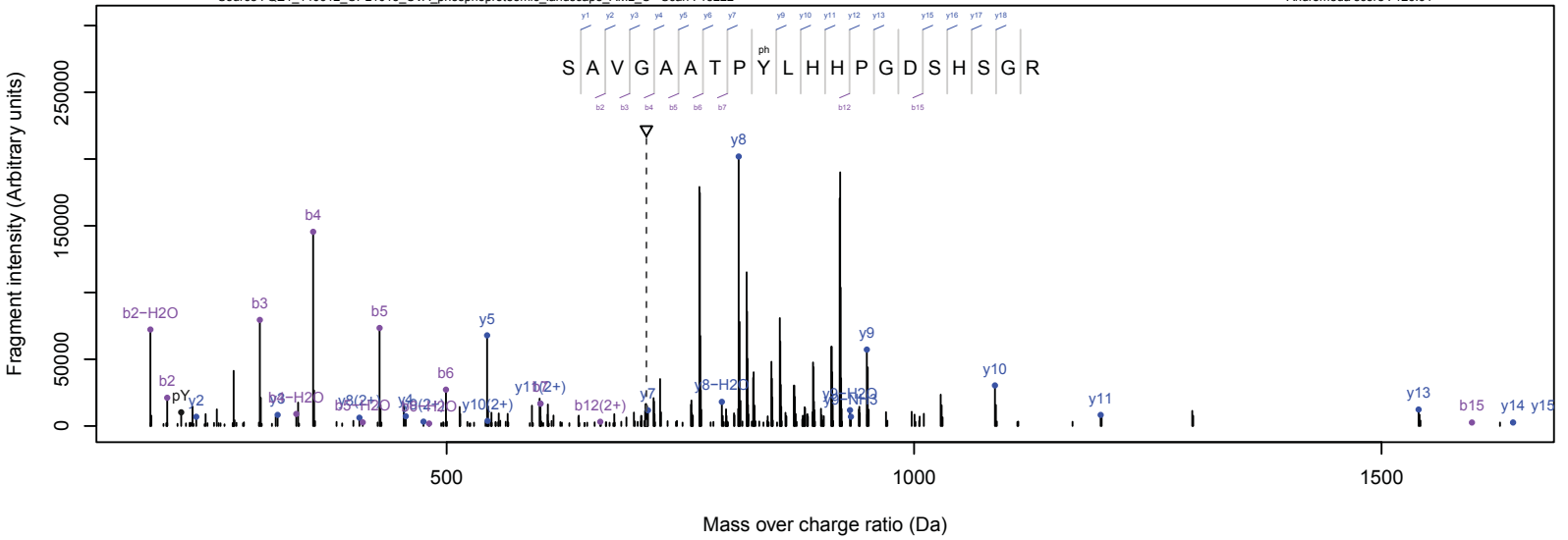
Andromeda score: 126.91



**(ac)SAVGAATPY(ph)LHHPGDSHSGR-3+**

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 15222

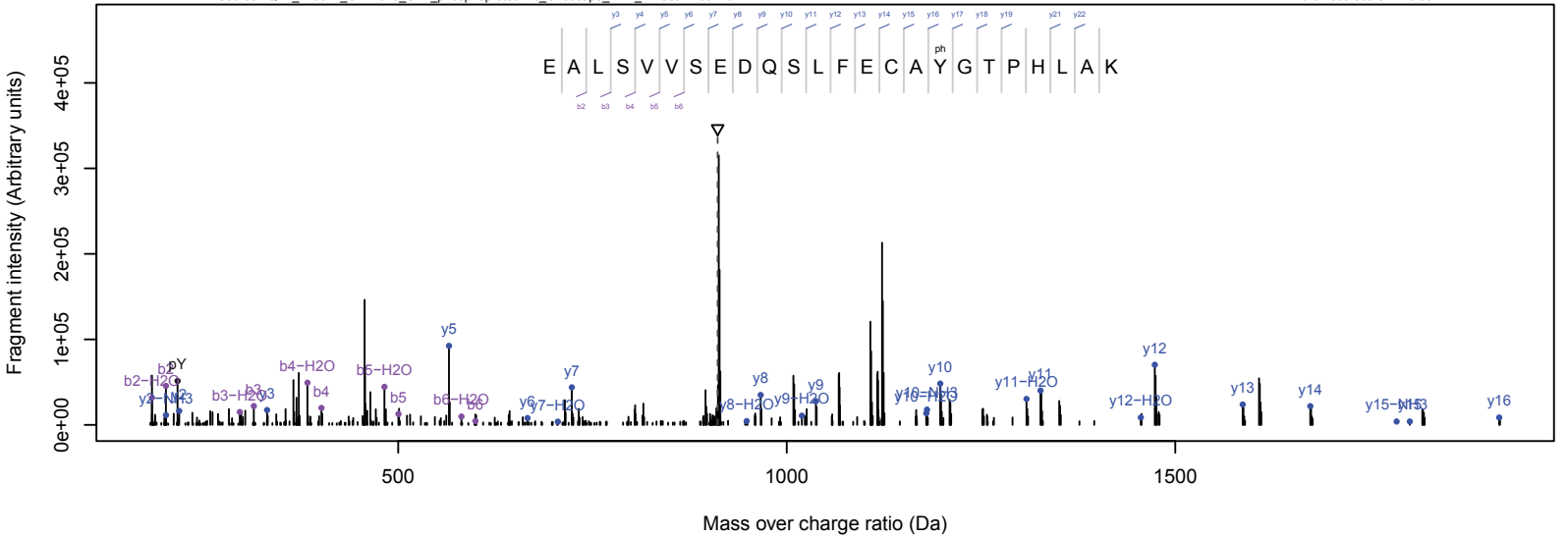
Andromeda score: 126.91



**EALSVVSEDQSLFECAY(ph)GTPHLAK-3+**

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan: 39416

Andromeda score: 126.86

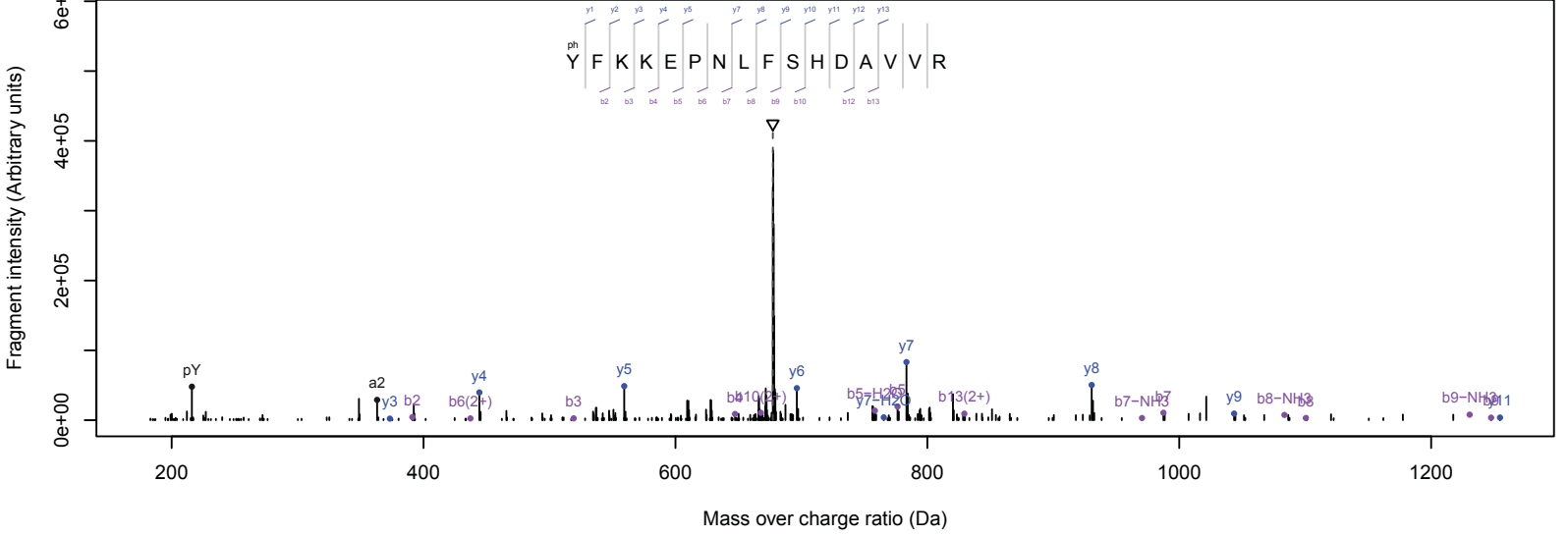




Y(ph)FKKEPNLFSHDAVVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21779

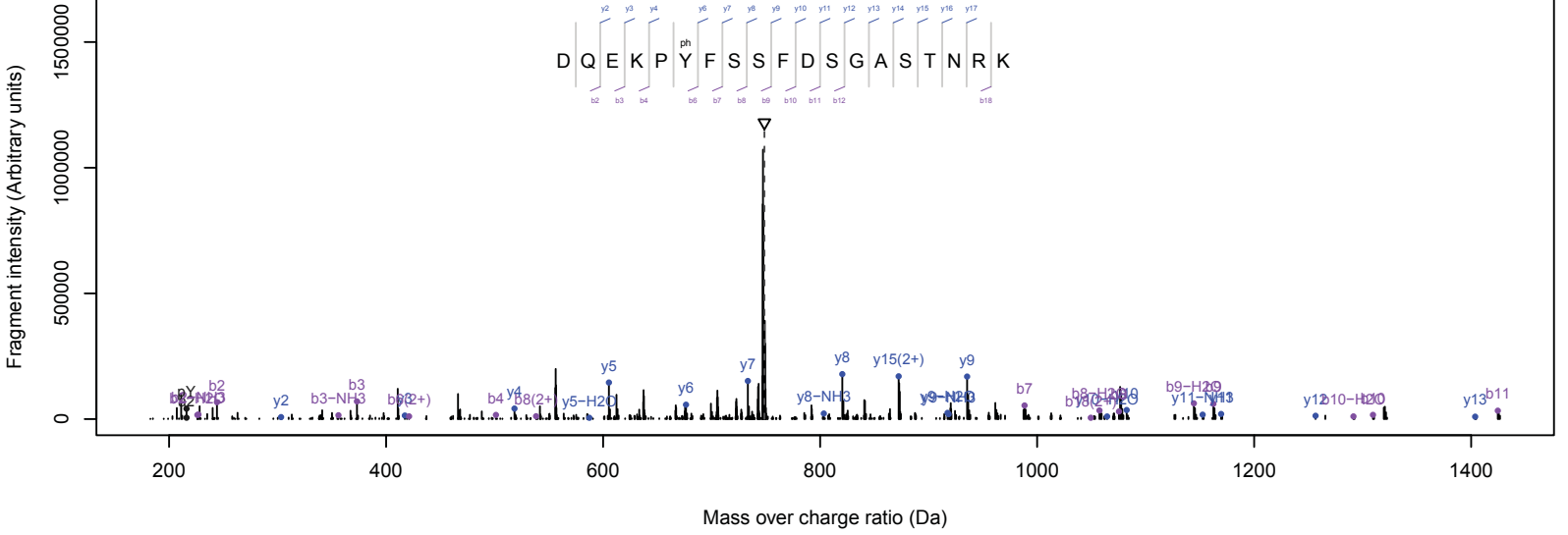
Andromeda score : 126.63



DQEKPY(ph)FSSFDSGASTNRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16983

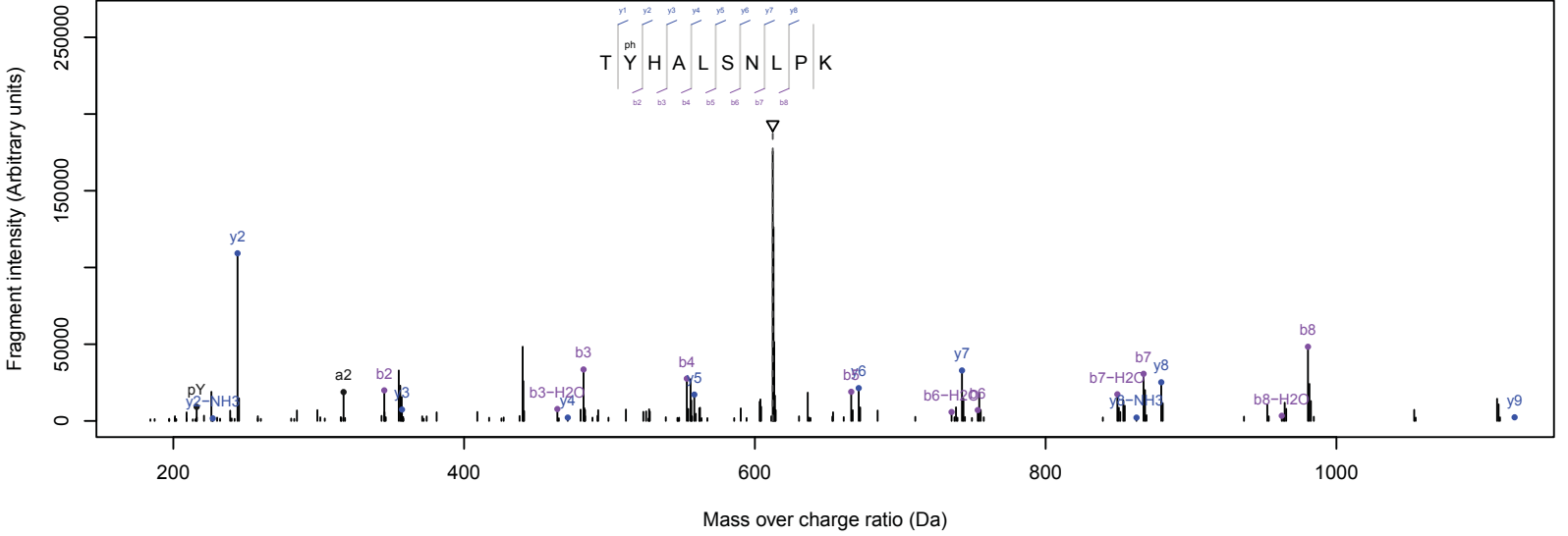
Andromeda score : 126.37



TY(ph)HALSNLPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12236

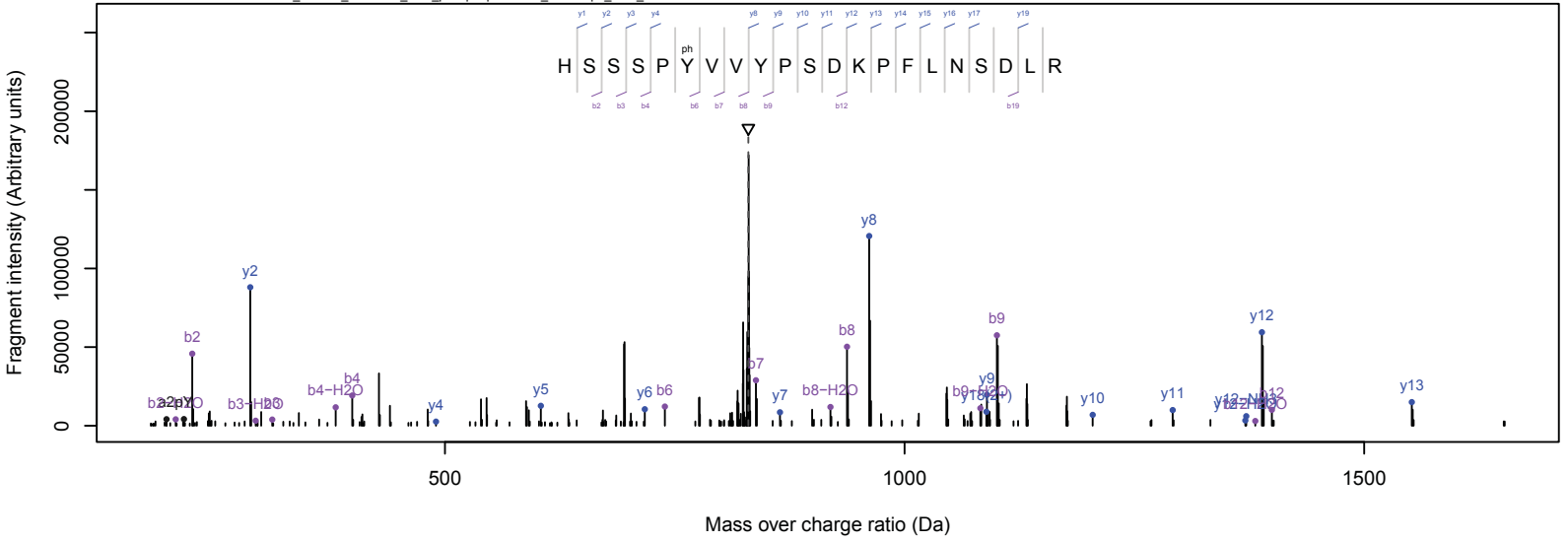
Andromeda score : 126.19



HSSSPY(ph)VVYPSDKPFLNSDLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30834

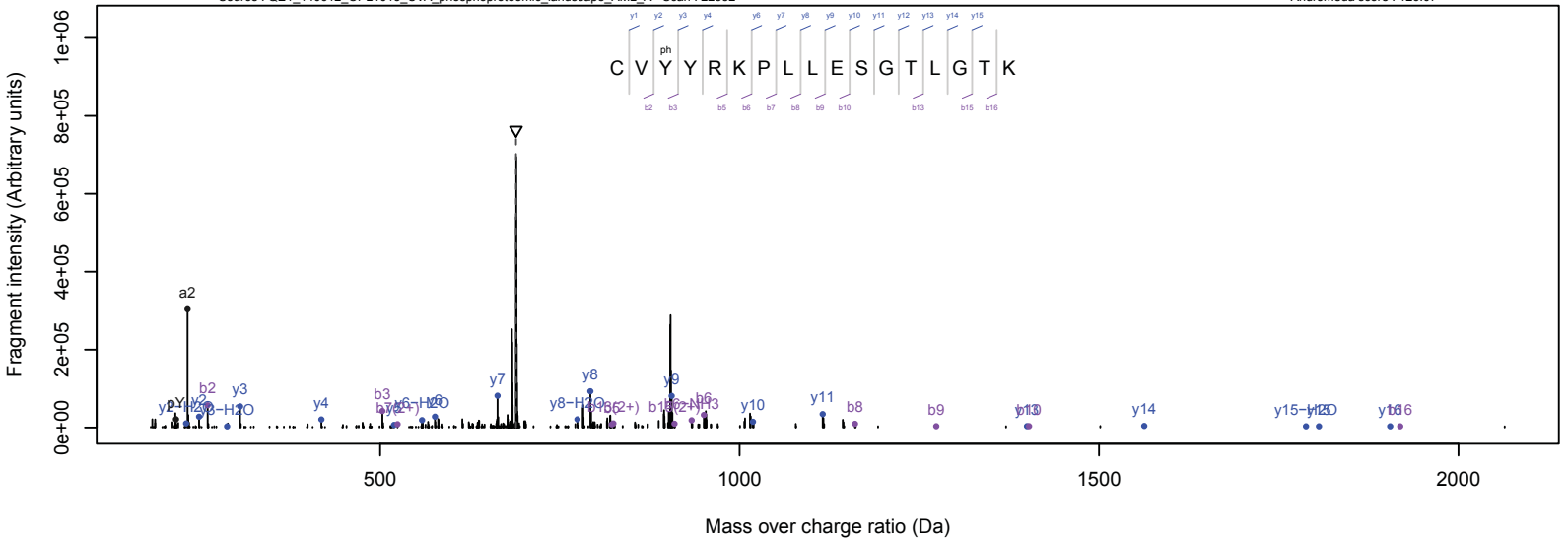
Andromeda score : 126.09



CVY(ph)YRKPLLESGLGTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 22352

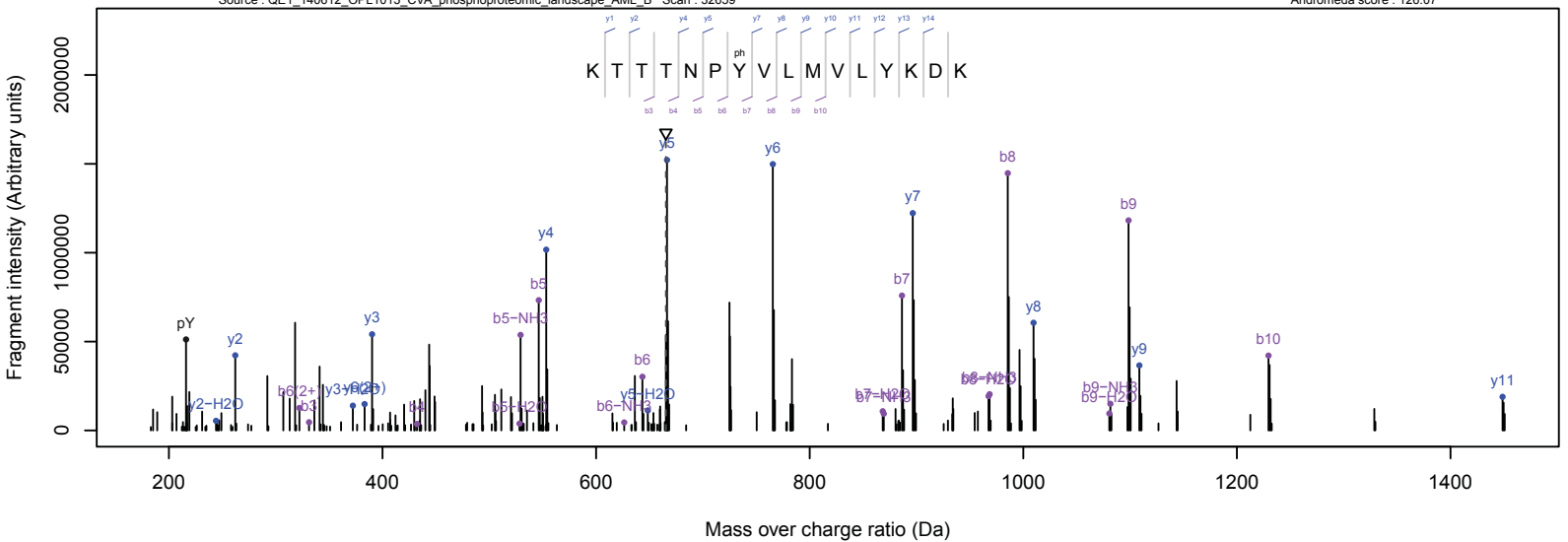
Andromeda score : 126.07



KTTTNPY(ph)VLMVLYKDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32659

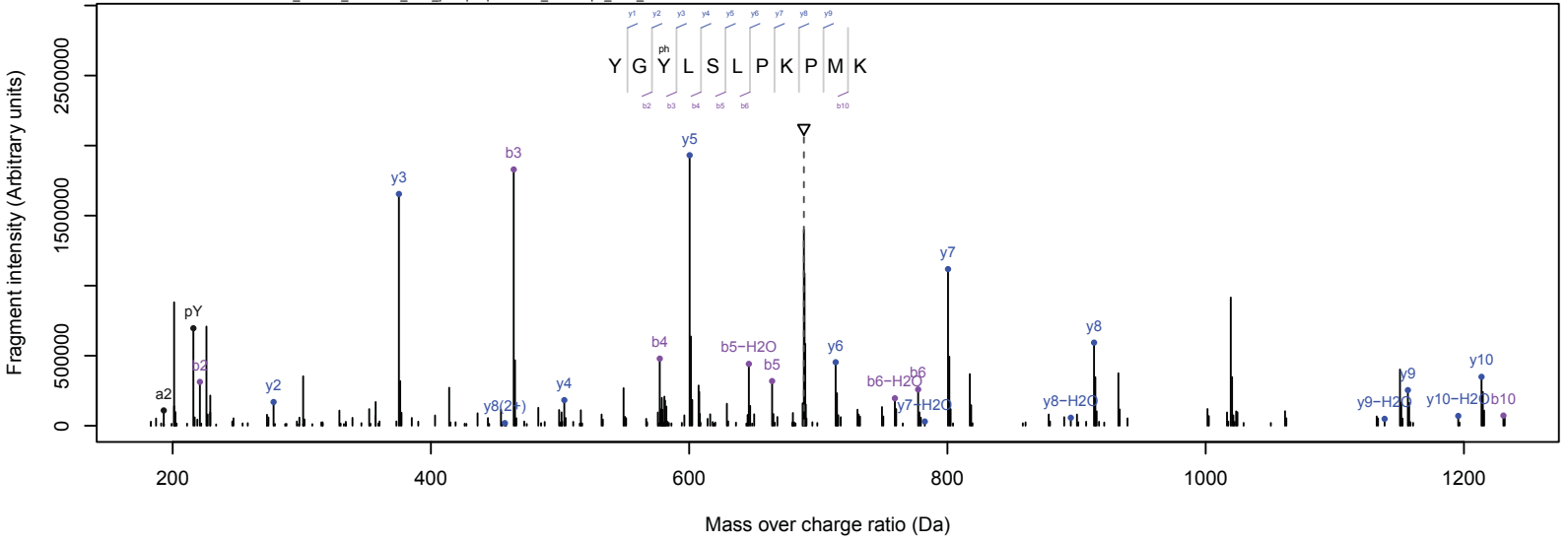
Andromeda score : 126.07



YGY(ph)LSLPKMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28052

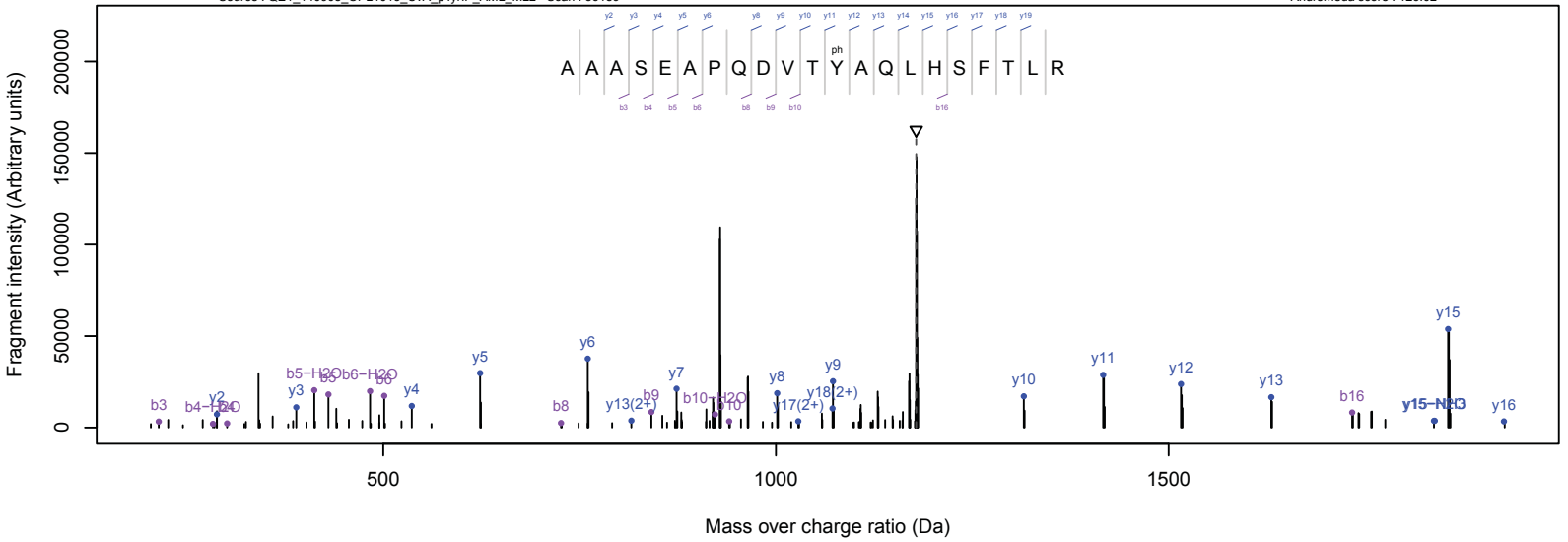
Andromeda score : 126.03



AAASEAPQDVTY(ph)AQLHSFTLR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 36189

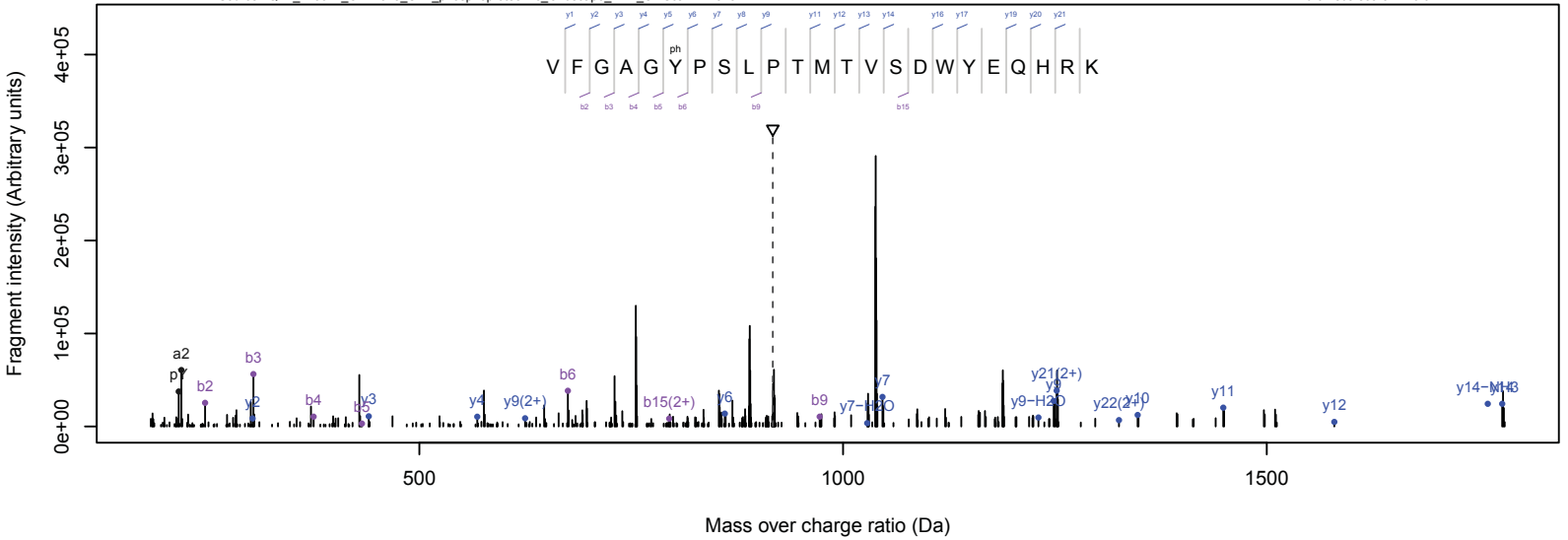
Andromeda score : 126.02



VFGAGY(ph)PSLPTMTVSDWYEQHRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41543

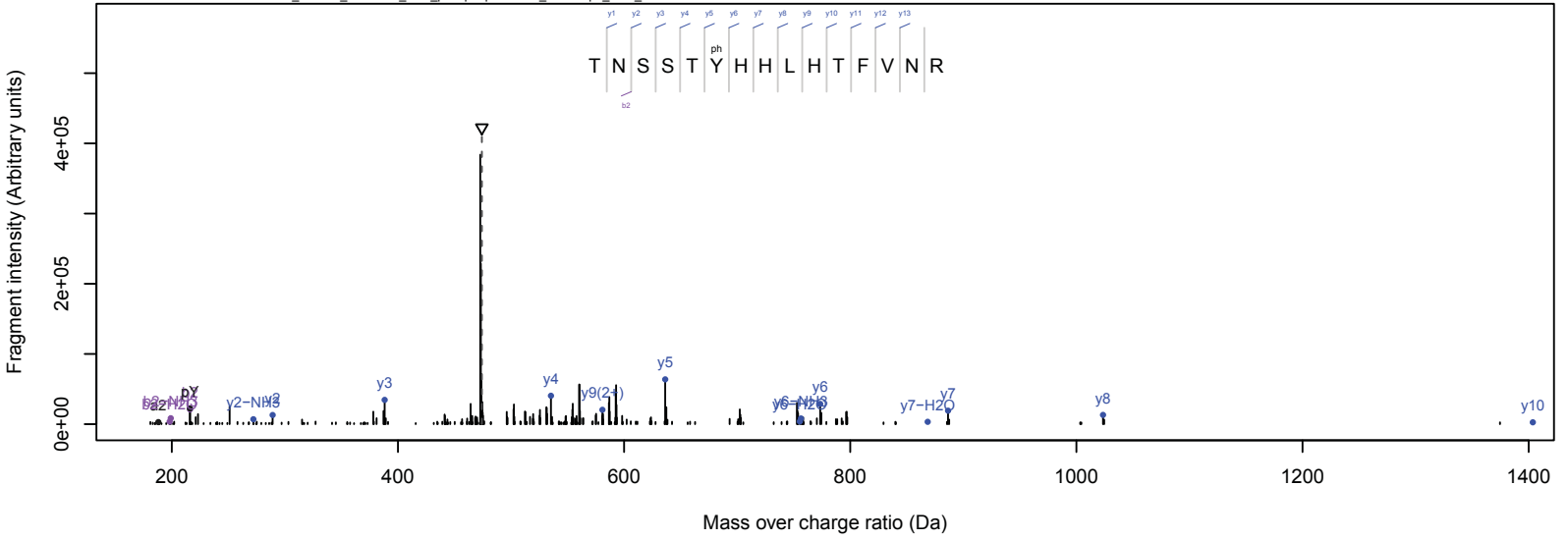
Andromeda score : 126.02



TNSSTY(ph)HLLHTFVNR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10343

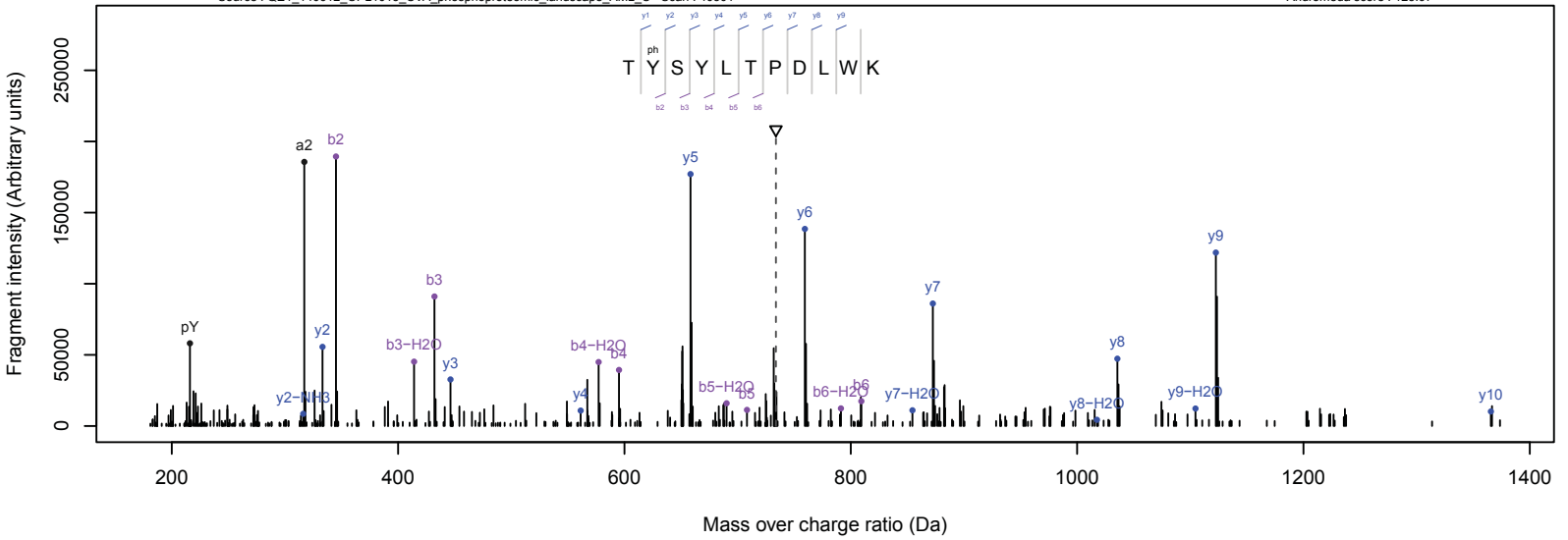
Andromeda score : 125.97



TY(ph)SYLTPDLWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40501

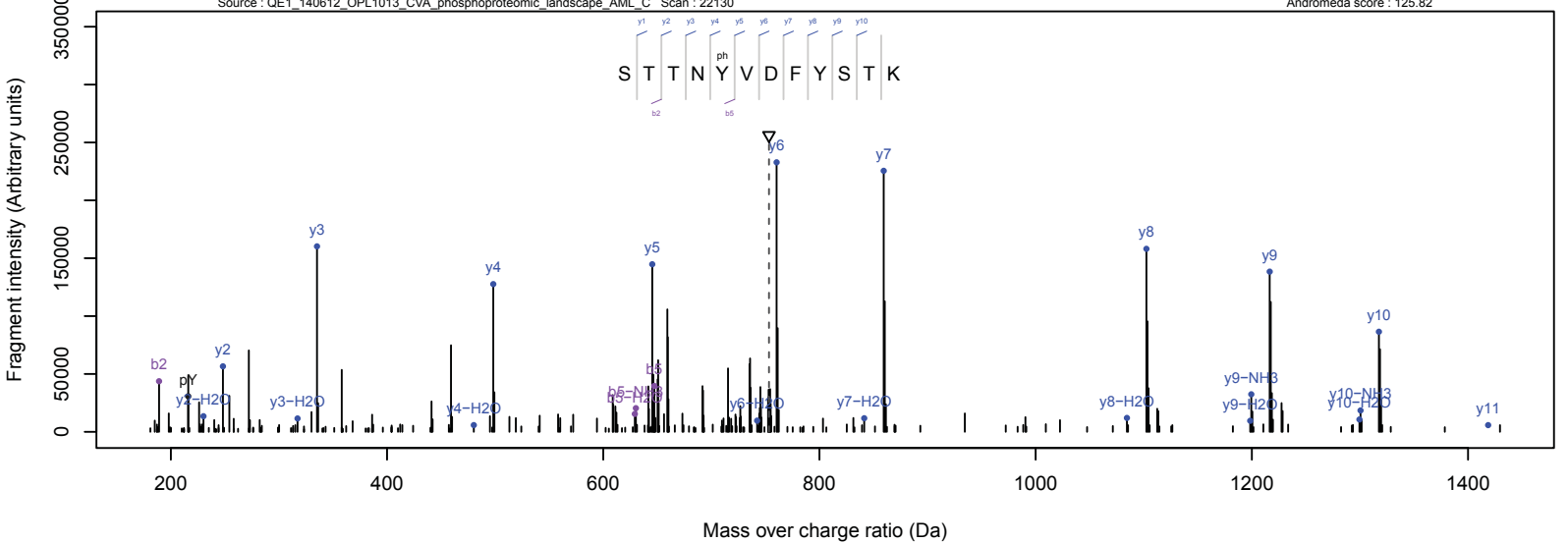
Andromeda score : 125.97

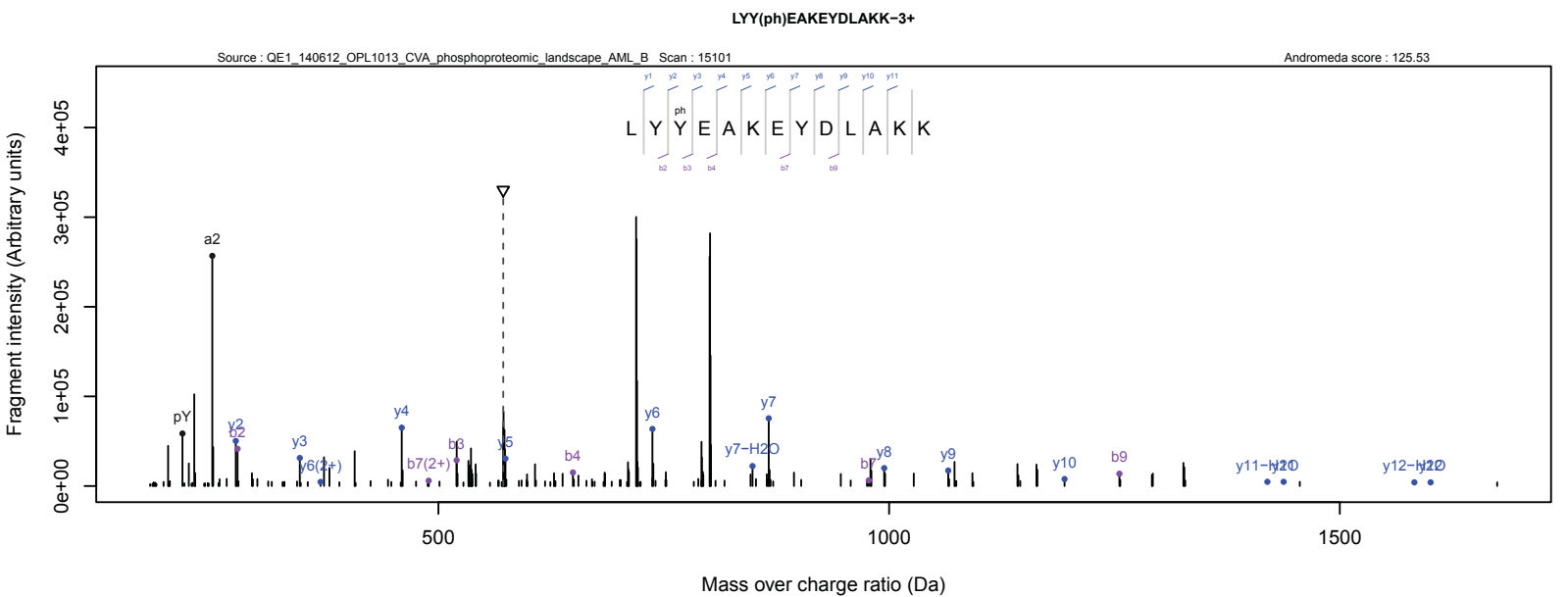
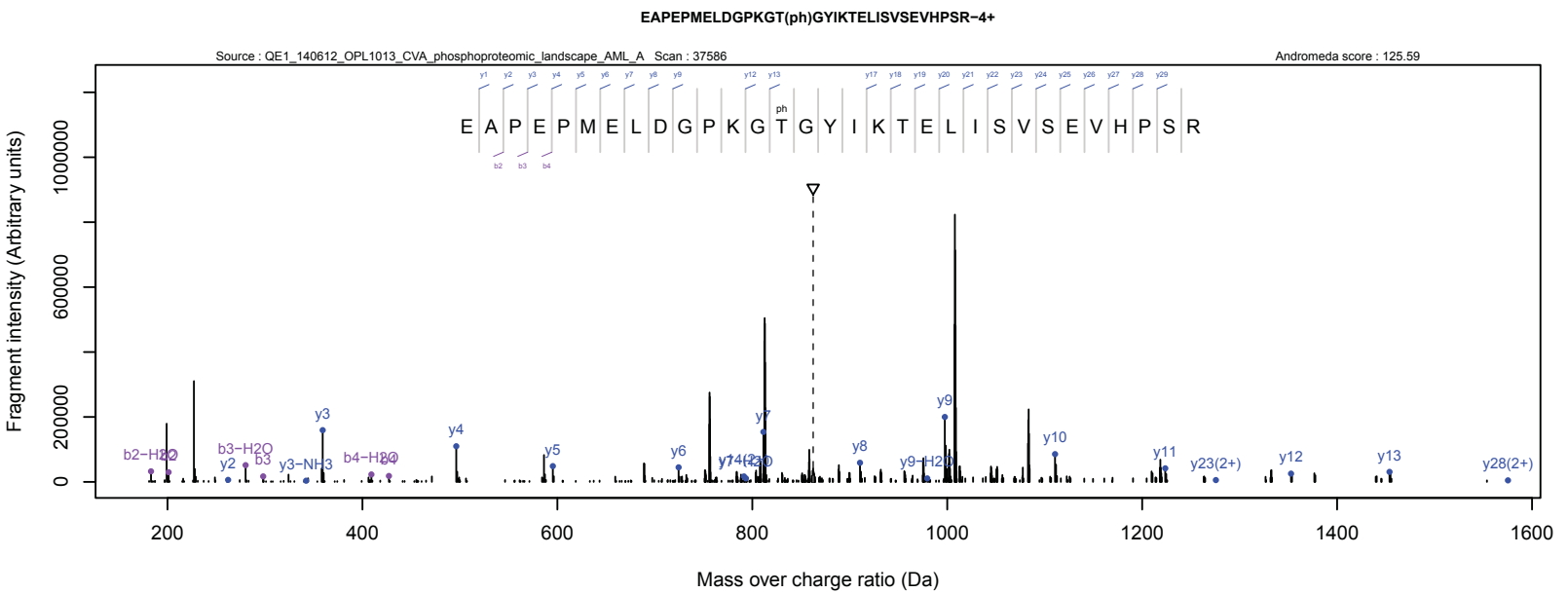
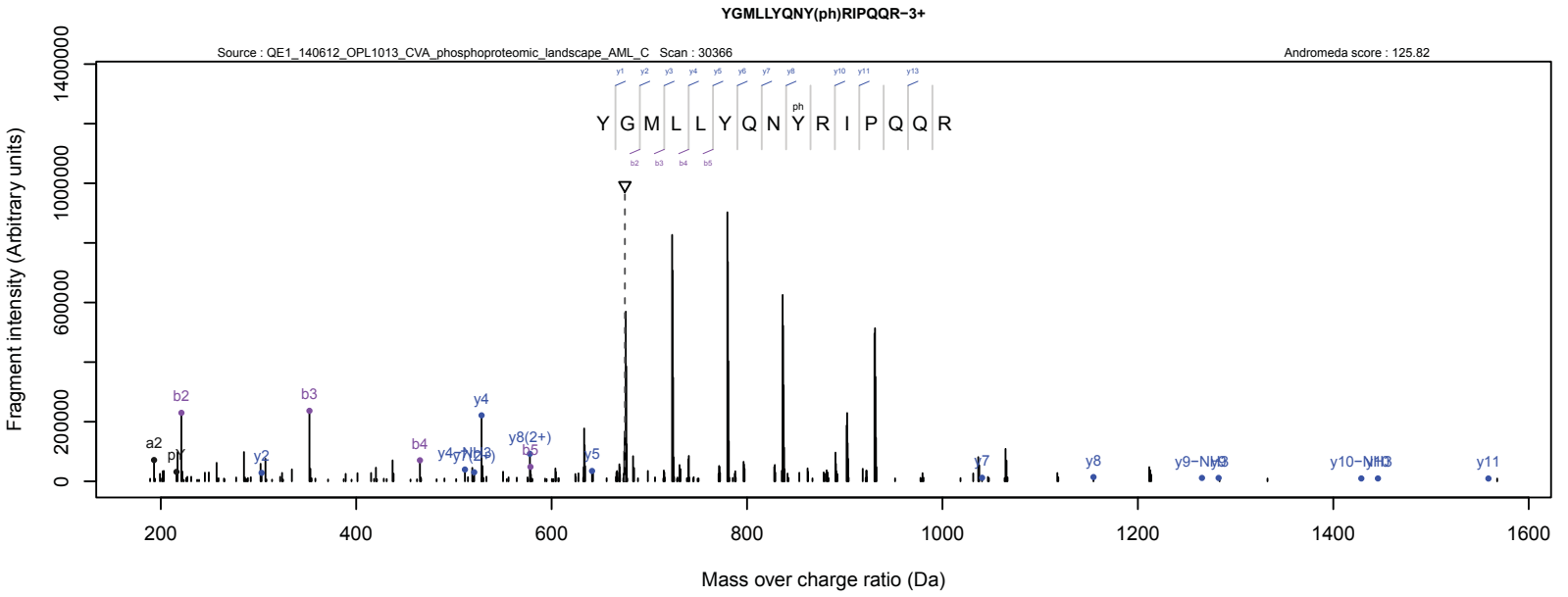


STTNY(ph)VDFYSTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22130

Andromeda score : 125.82

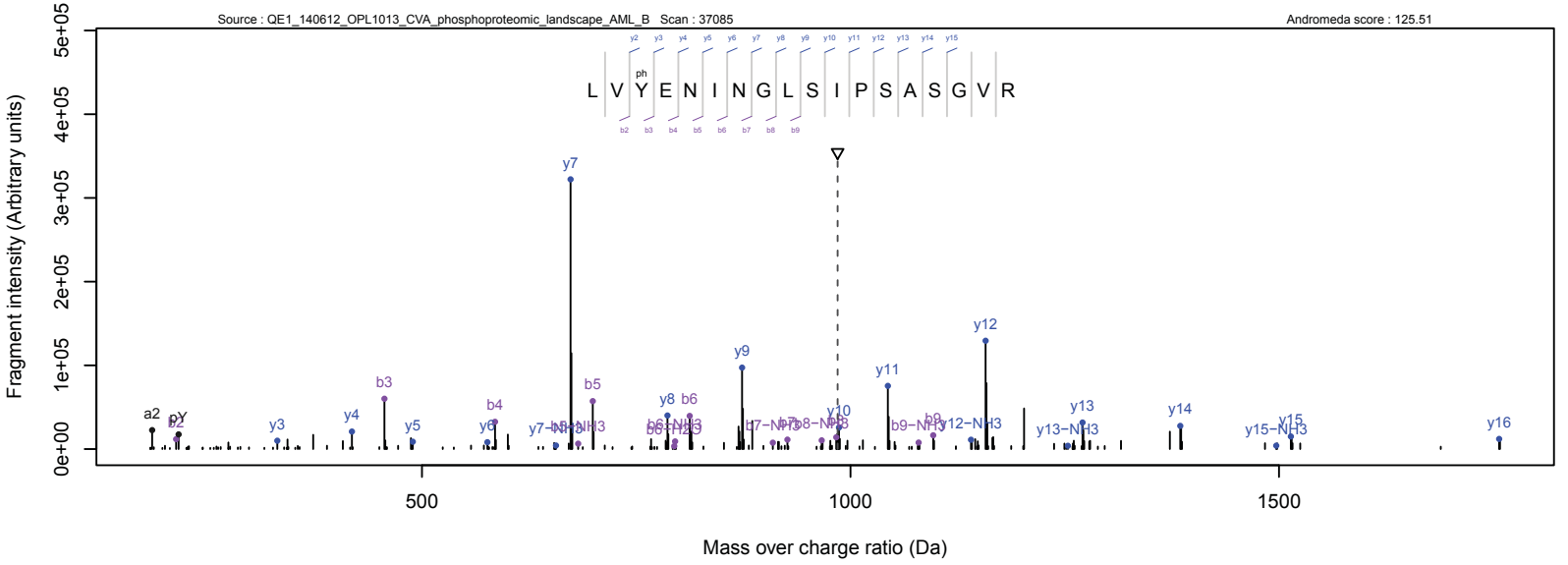




LVY(ph)ENINGLSIPSASGVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37085

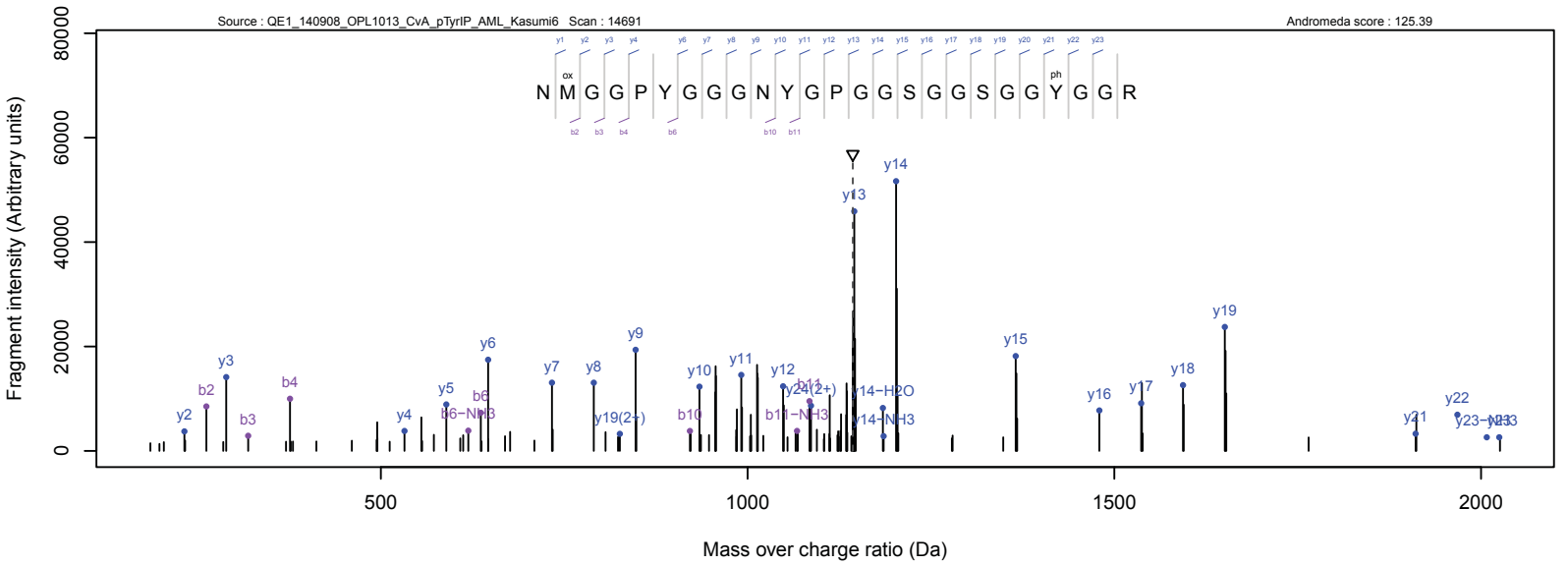
Andromeda score : 125.51



NM(ox)GGPYGGNGYGGSGGGY(ph)GGR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 14691

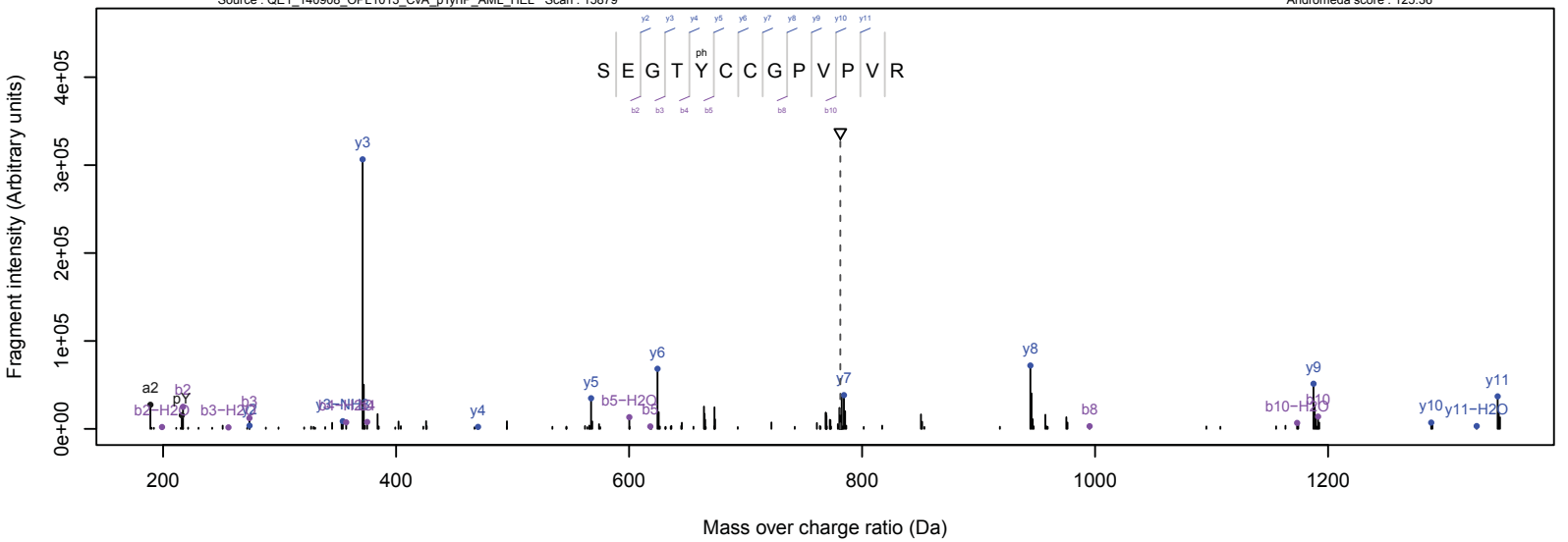
Andromeda score : 125.39



SEGTY(ph)CCGPVVR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 15879

Andromeda score : 125.36

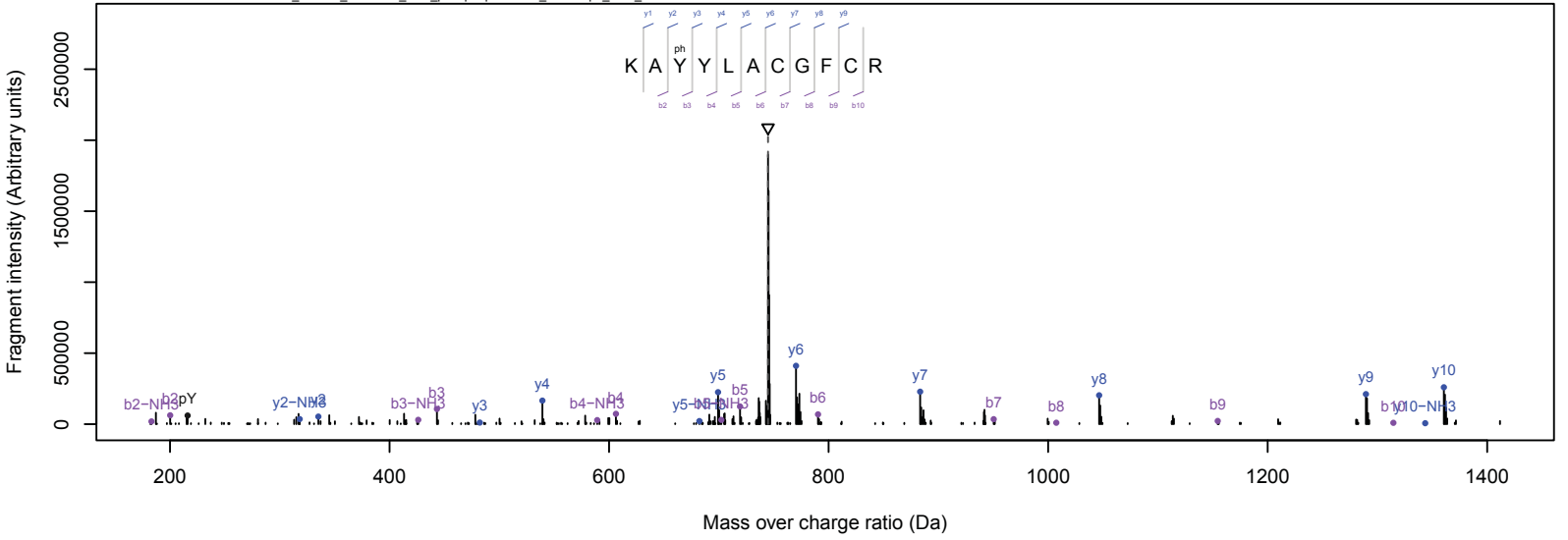




**KAY(ph)YLACGFCR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21374

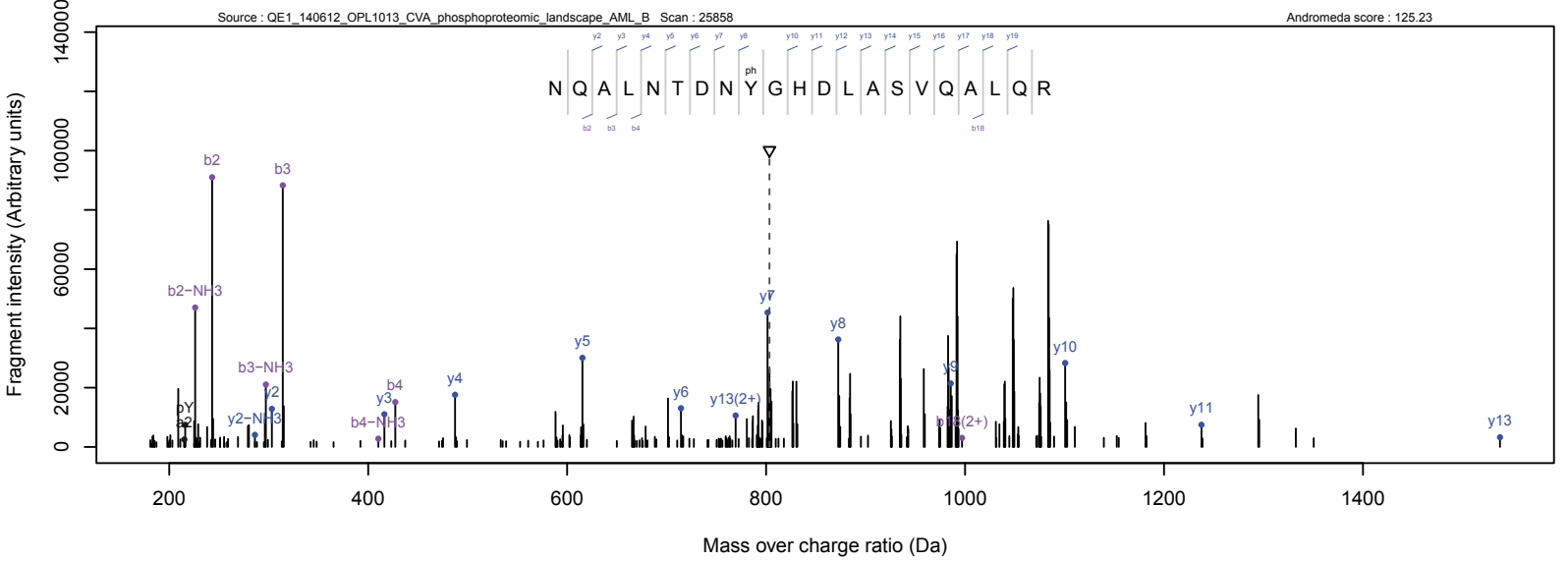
Andromeda score: 125.23



**NQALNTDNY(ph)GHDLASVQALQR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 25858

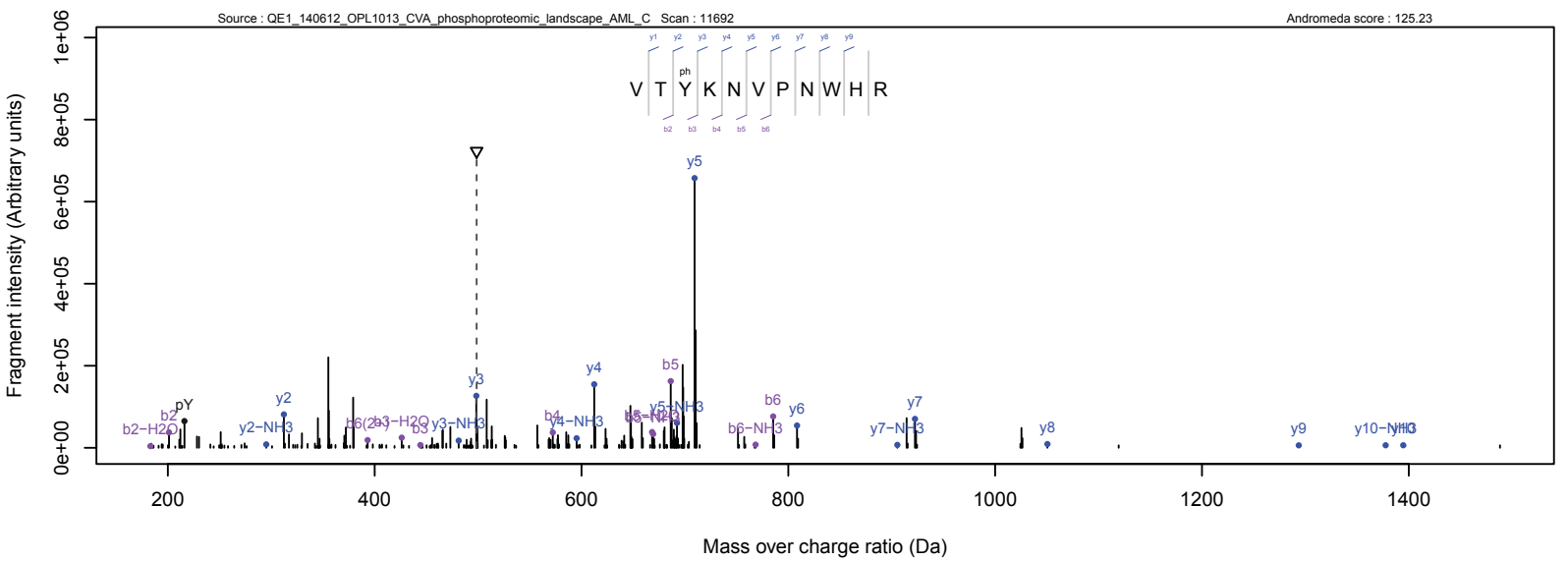
Andromeda score: 125.23



**VTY(ph)KNVPNWHR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11692

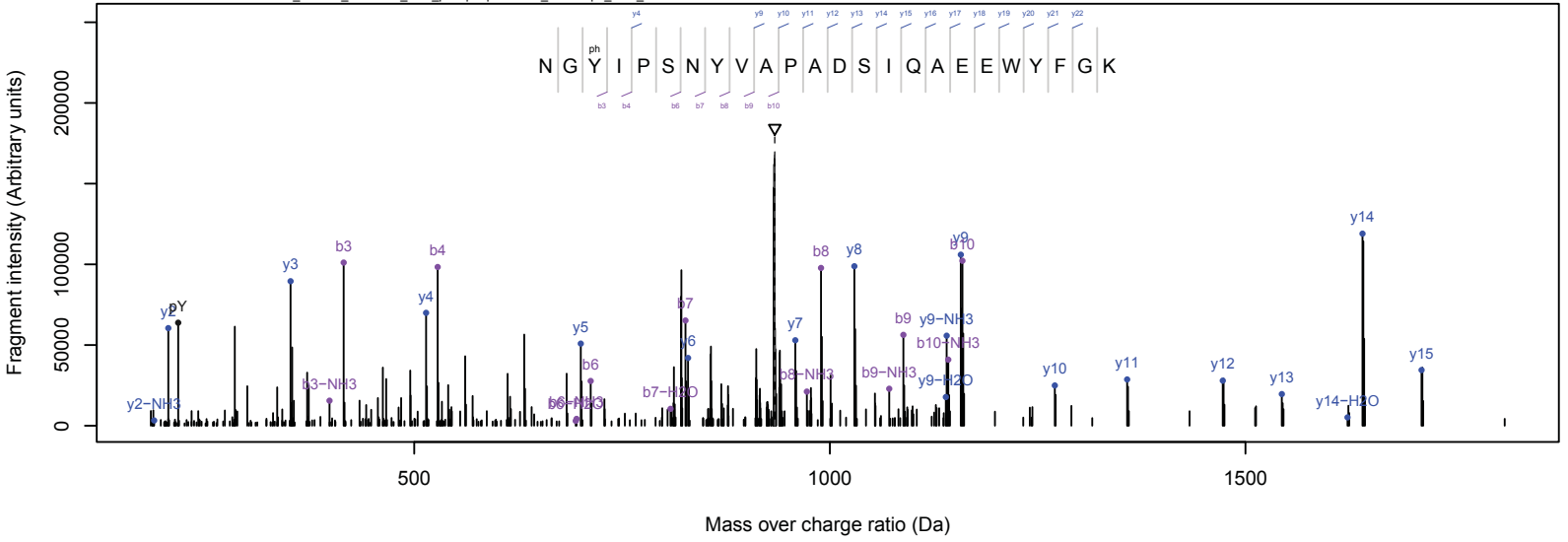
Andromeda score: 125.23



NGY(ph)IPSNYVAPADSIQAEWYFGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47163

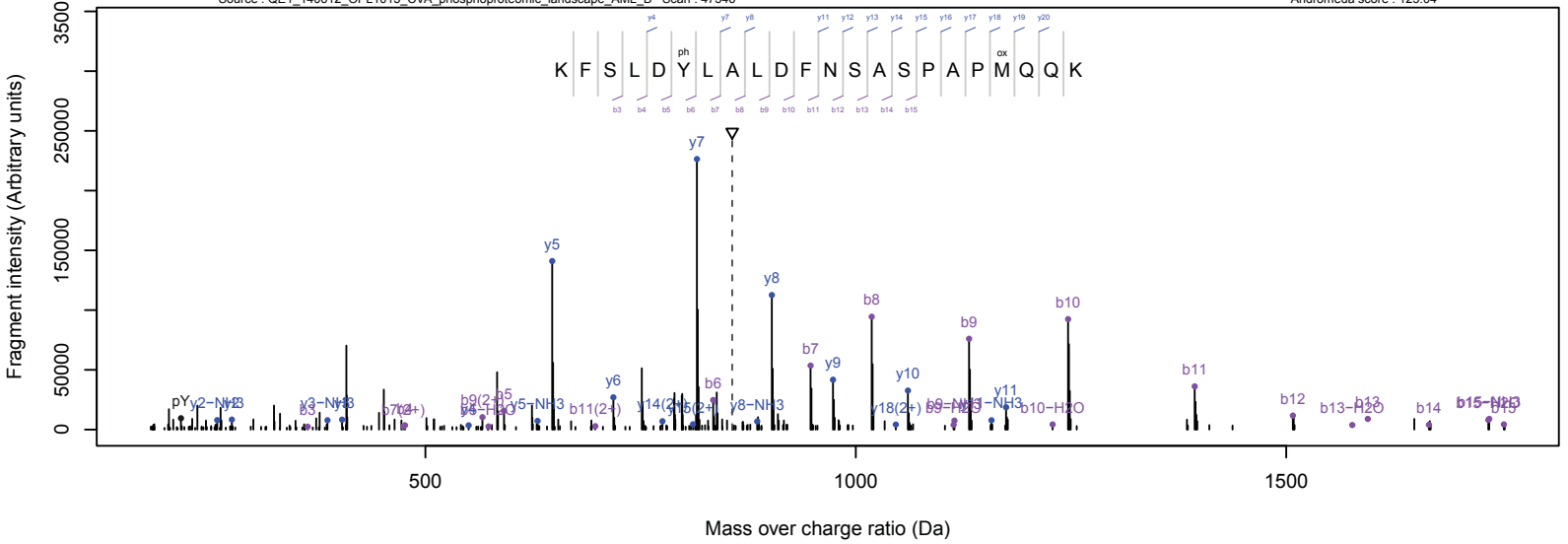
Andromeda score : 125.11



KFSLDY(ph)LALDFNSASPAPMox)QQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47346

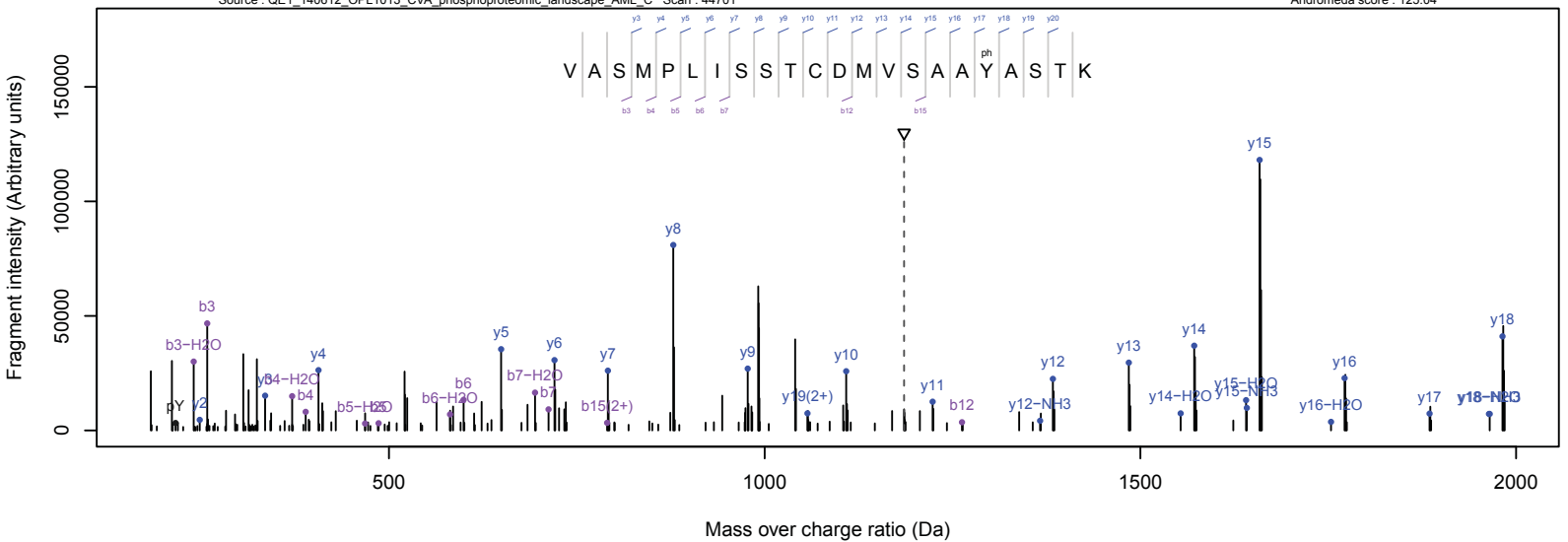
Andromeda score : 125.04



VASMP LISSTCDMVSAAAY(ph)ASTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44701

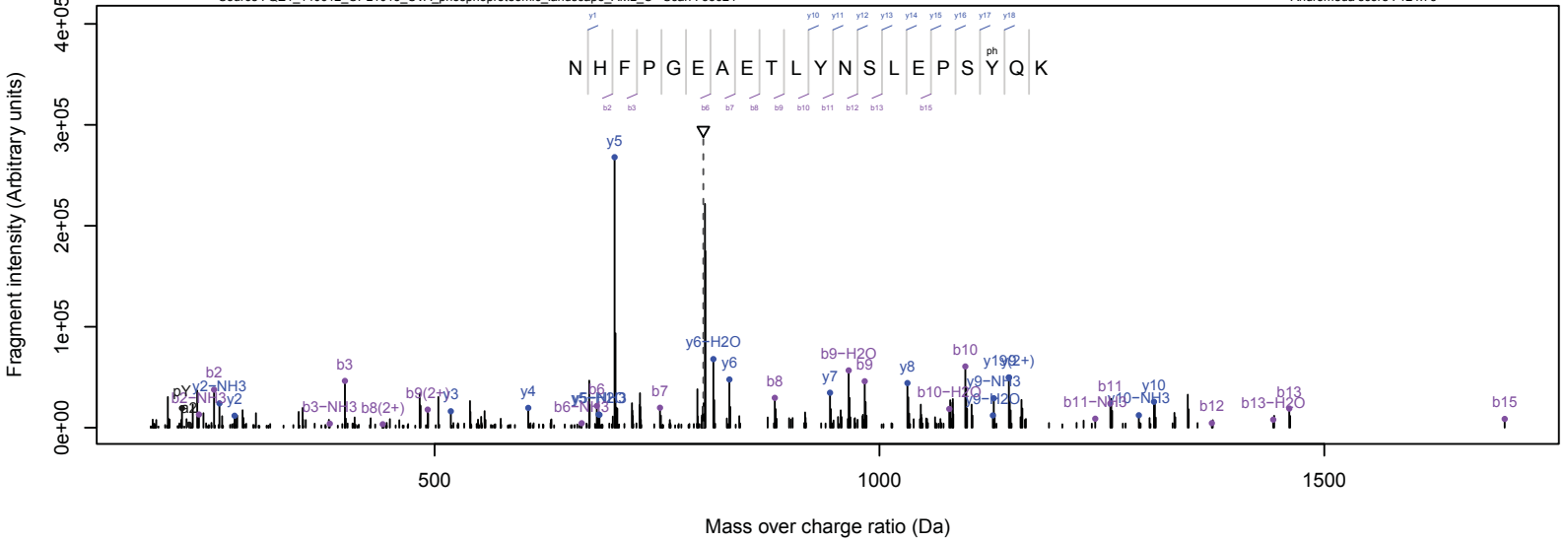
Andromeda score : 125.04



**NHFPGEAETLYNSLEPSYQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33624

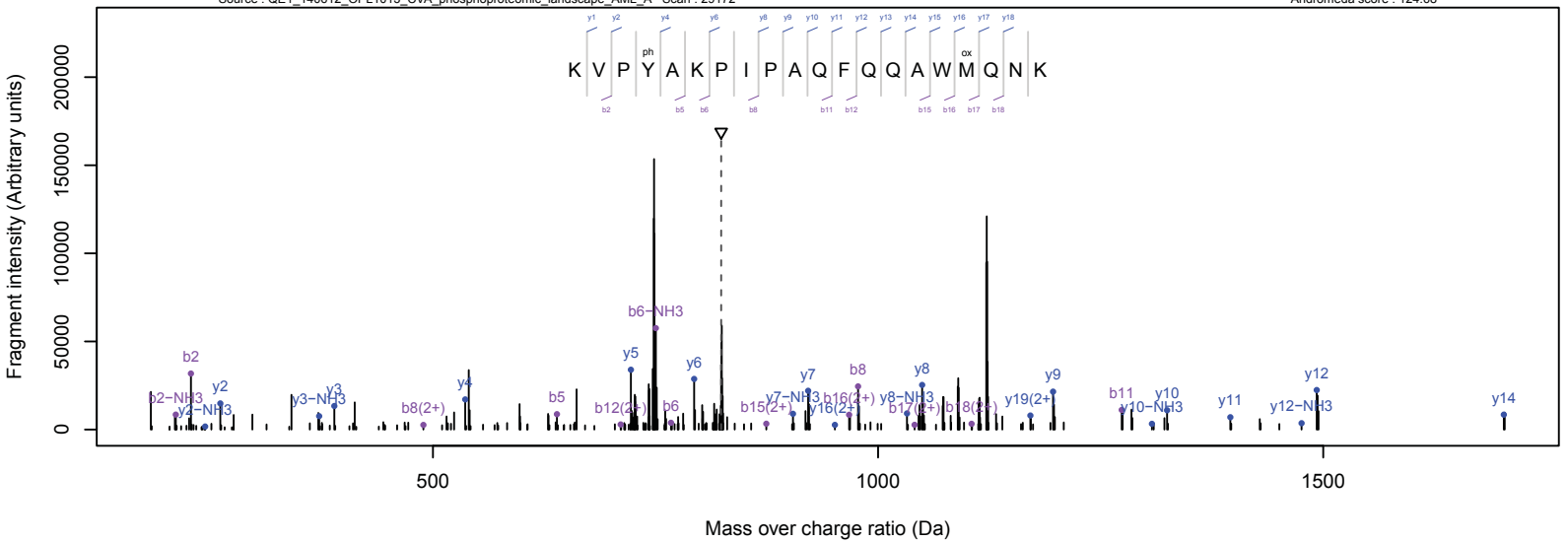
Andromeda score : 124.73



**KVPY(ph)AKPIPAQFQQAWM(ox)QNK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29172

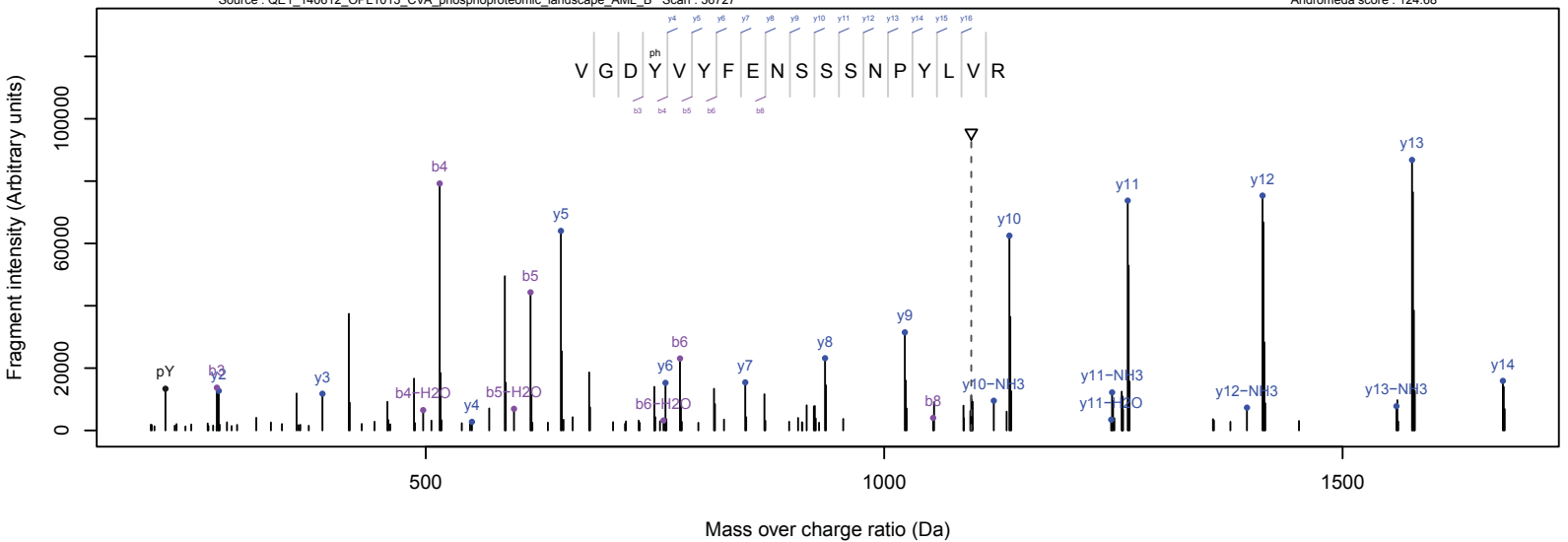
Andromeda score : 124.68



**VGDY(ph)VYFENSSNPYLVR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36727

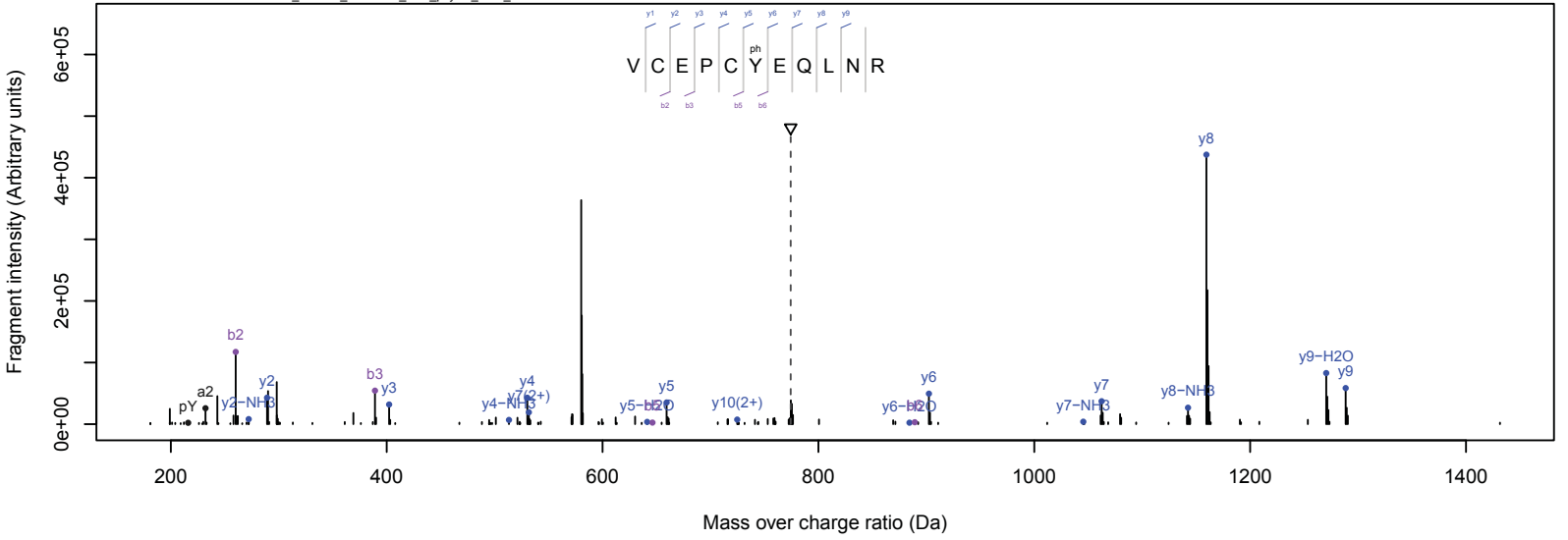
Andromeda score : 124.68



VCEPCY(ph)EQLNR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 15214

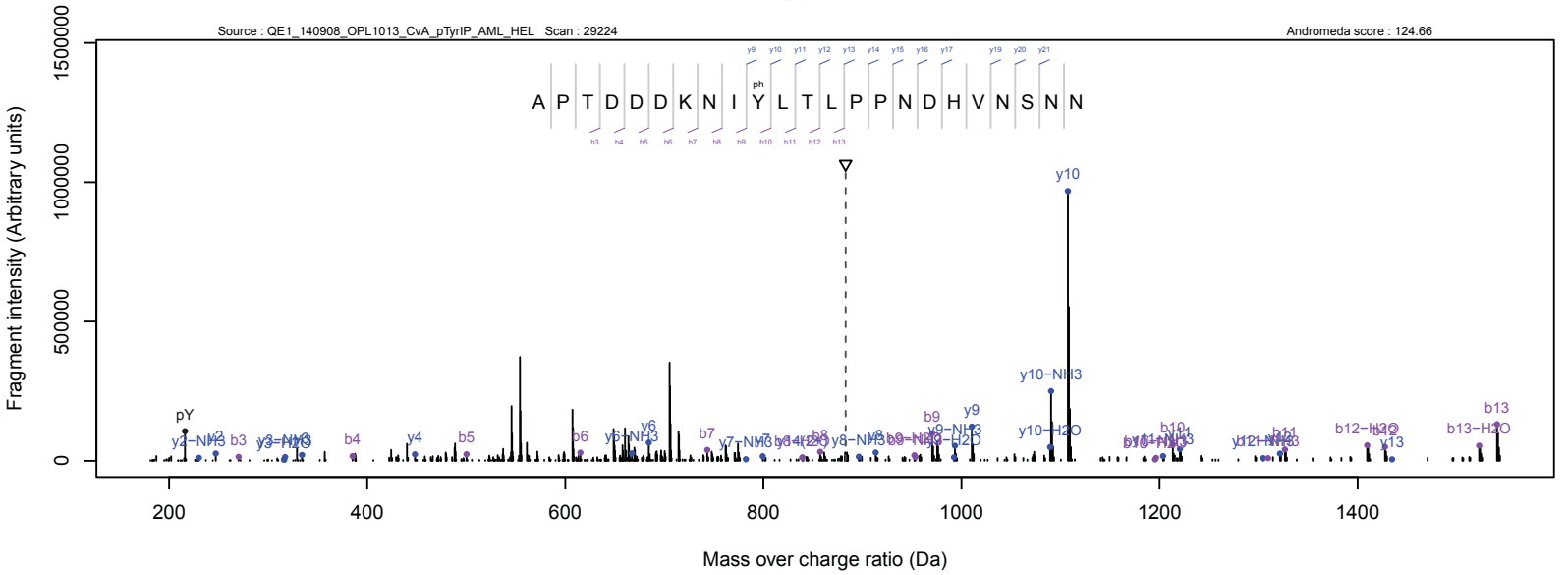
Andromeda score : 124.67



APTDDDKNIY(ph)LTLPNDHVSNN-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29224

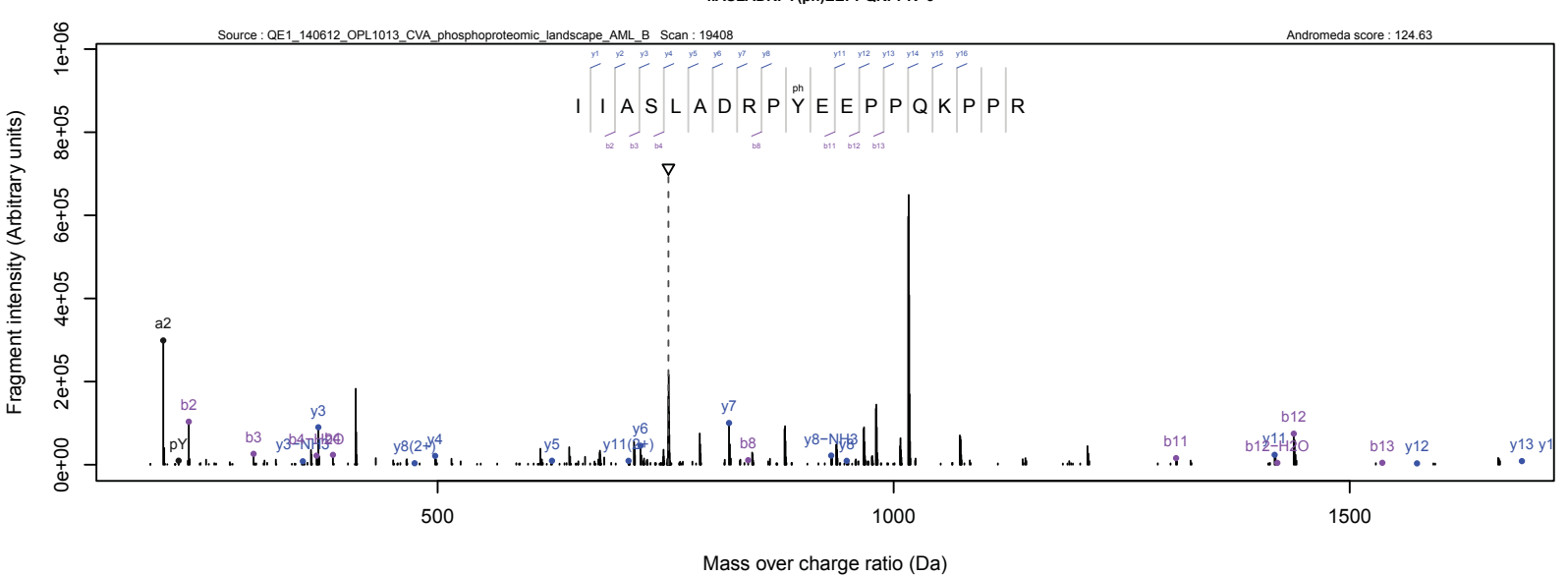
Andromeda score : 124.66



IIASLDRPY(ph)EPPQKPPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19408

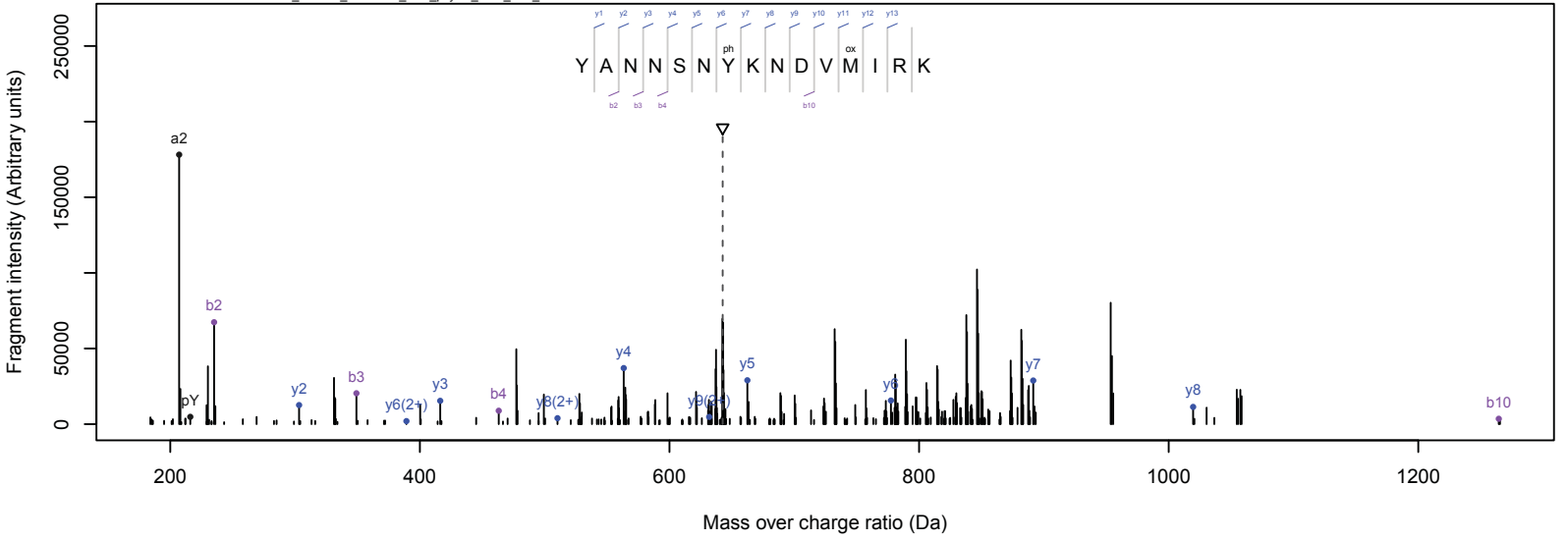
Andromeda score : 124.63



YANNSNY(ph)KNDVMOIRK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 7157

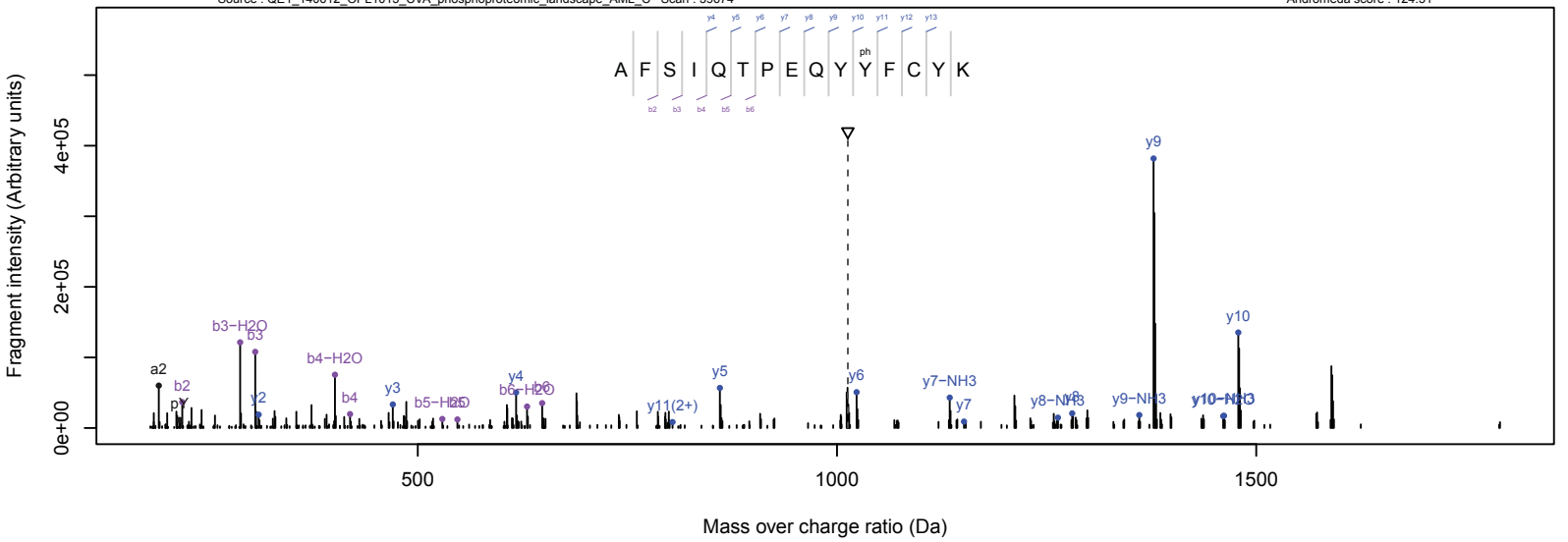
Andromeda score : 124.62



AFSIQTPEQY(ph)FCYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39674

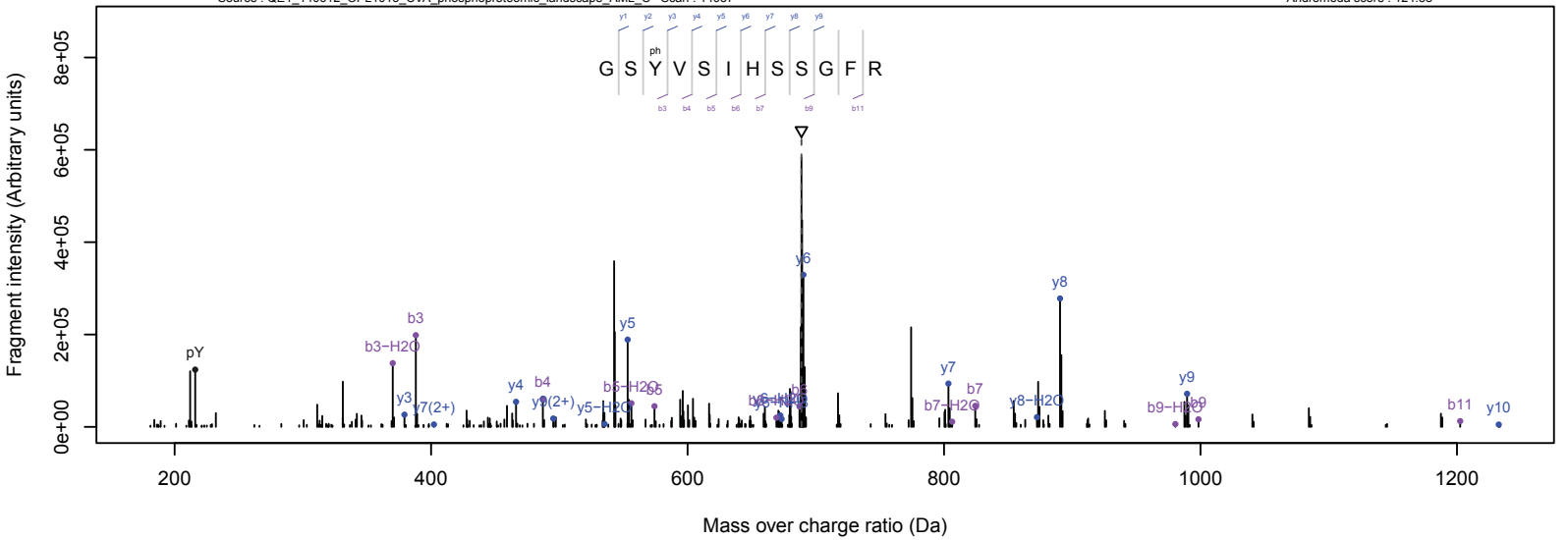
Andromeda score : 124.51



GSY(ph)VSIISSGFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14937

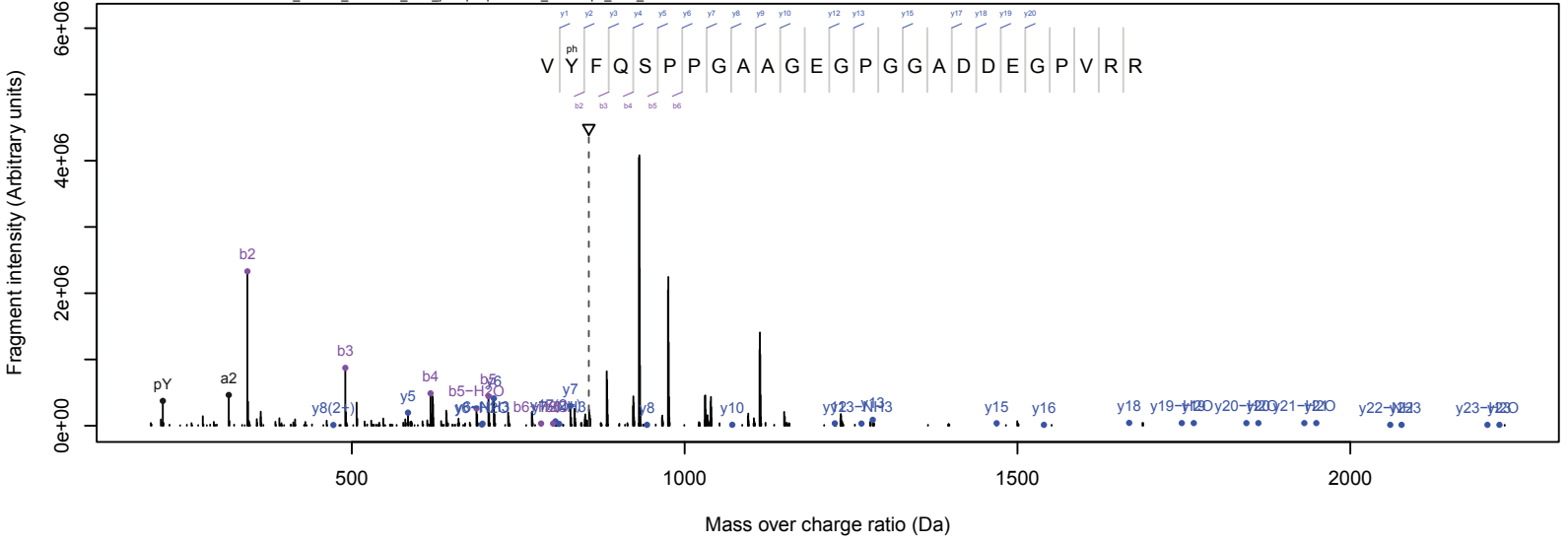
Andromeda score : 124.38



VY(ph)FQSPPGAAGEGPGADDEGPVRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22935

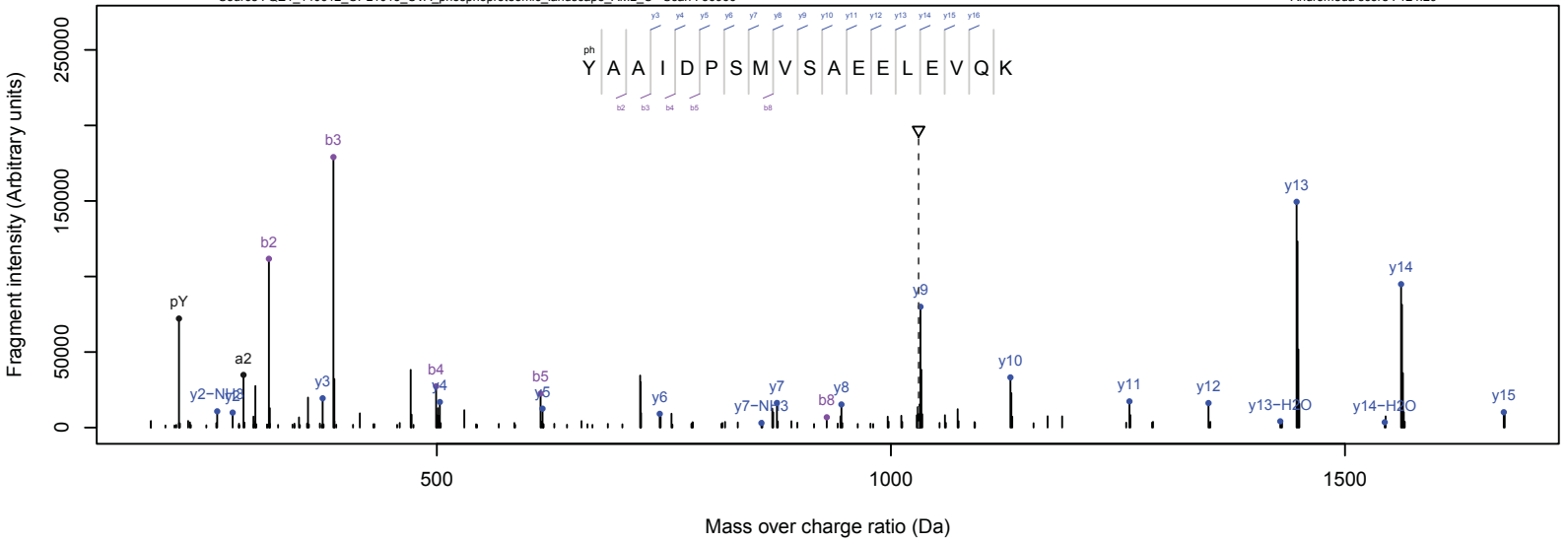
Andromeda score : 124.31



Y(ph)AAIDPSMVAEELEVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36966

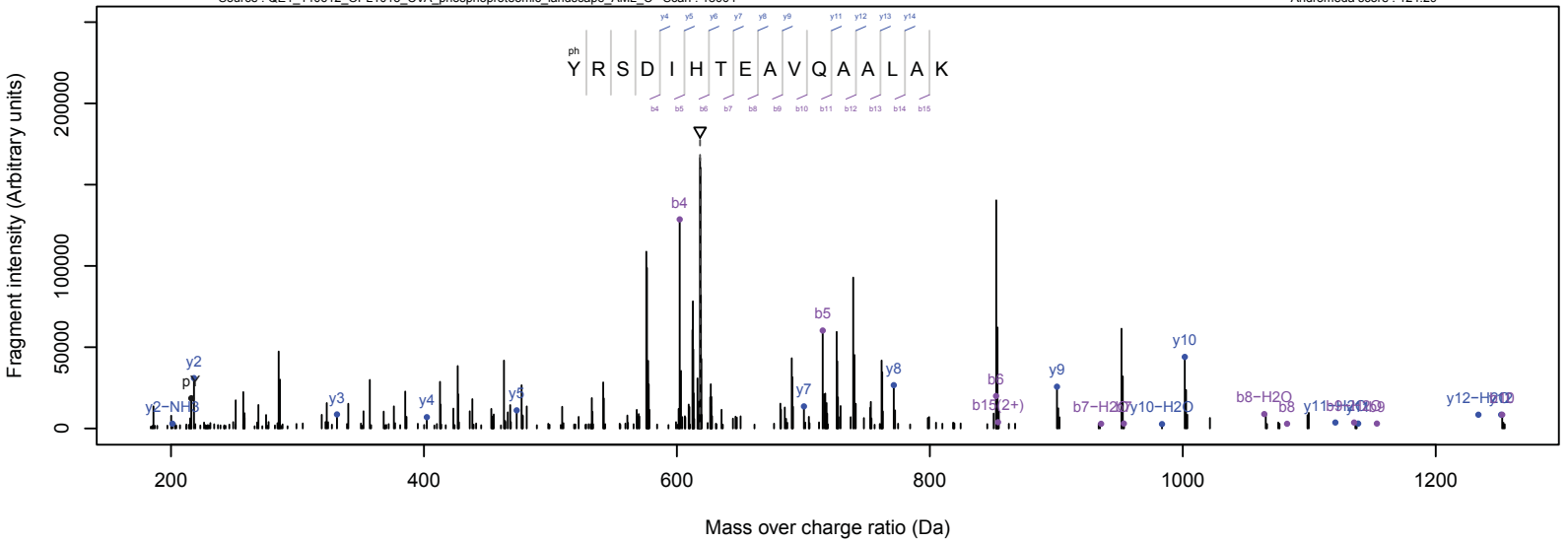
Andromeda score : 124.29



Y(ph)RSDIHTEAVQAALAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18004

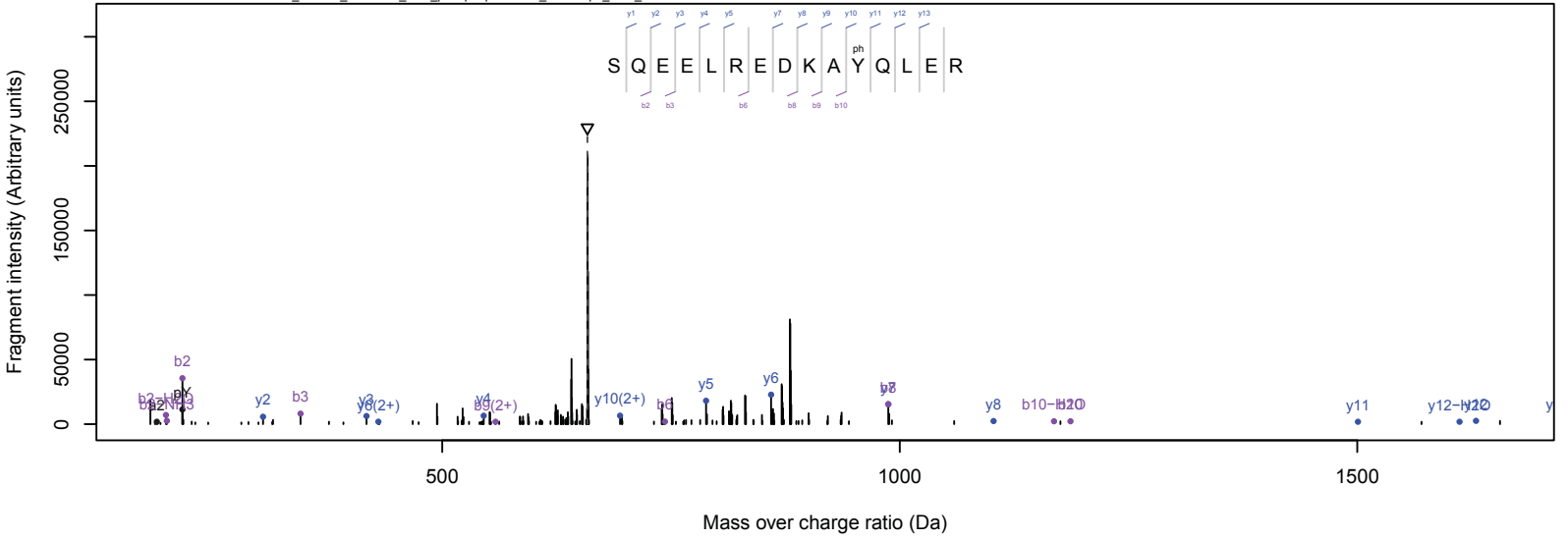
Andromeda score : 124.29



SQEELREDKAYQLER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10442

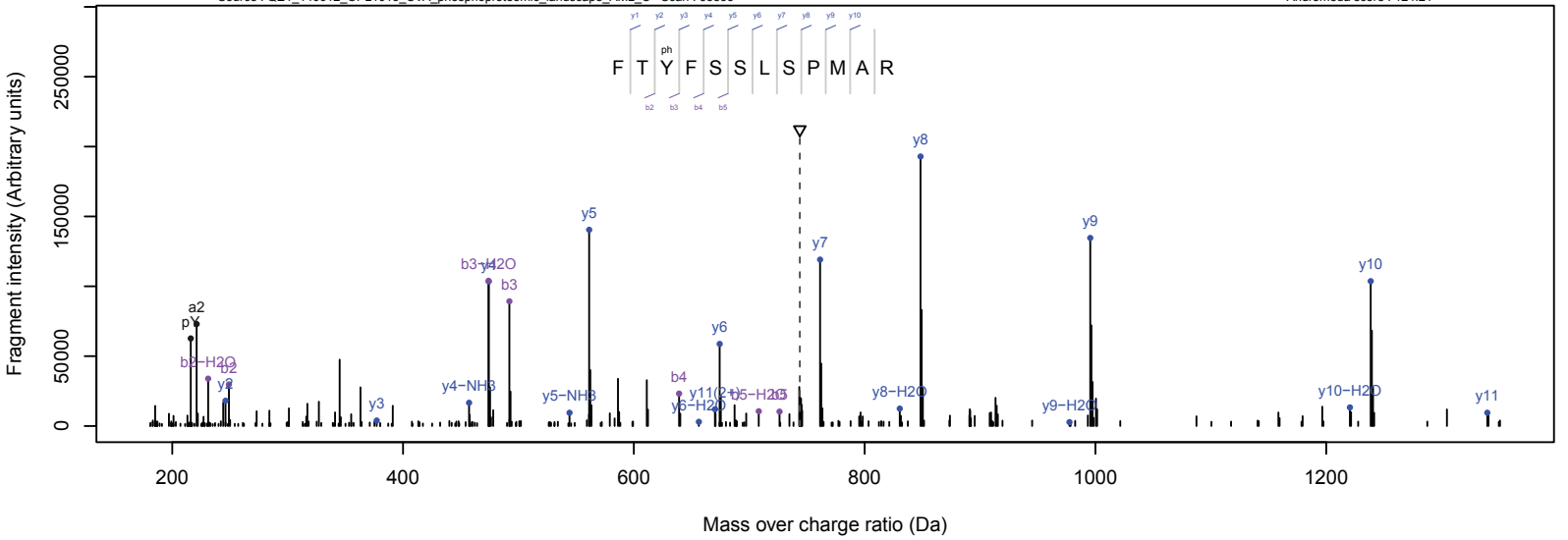
Andromeda score : 124.25



FTY(ph)FSSSPMAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35833

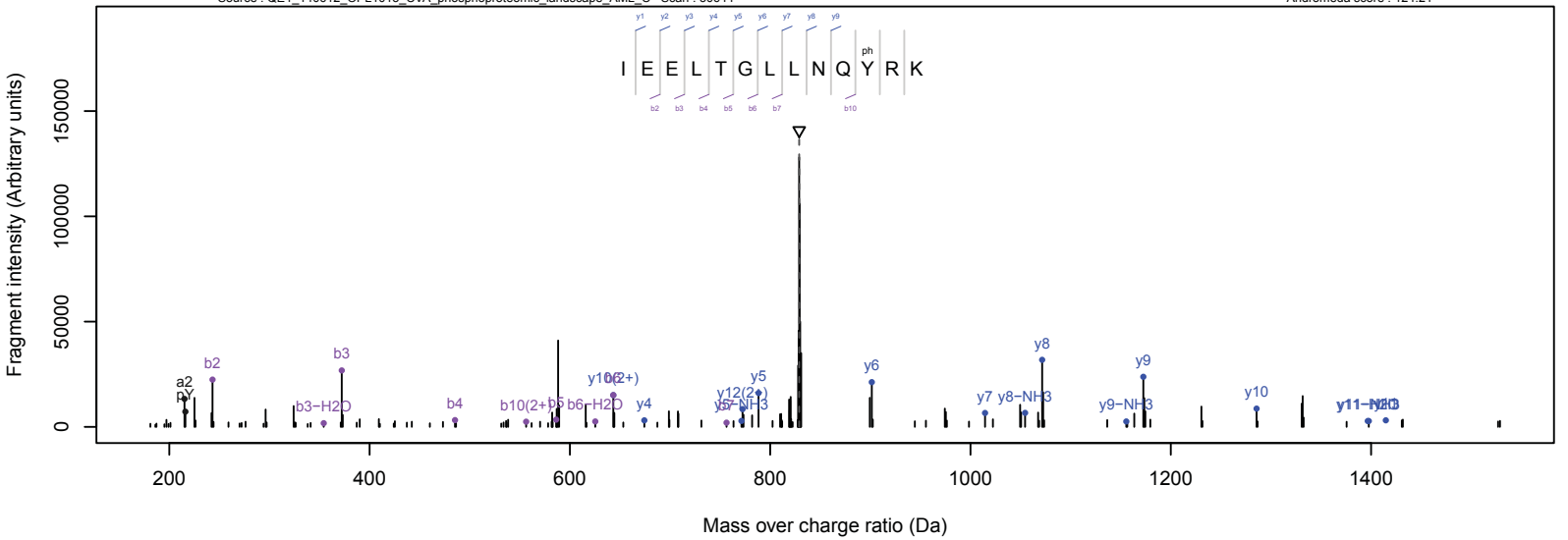
Andromeda score : 124.21



IEELTGLLNQYRK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30644

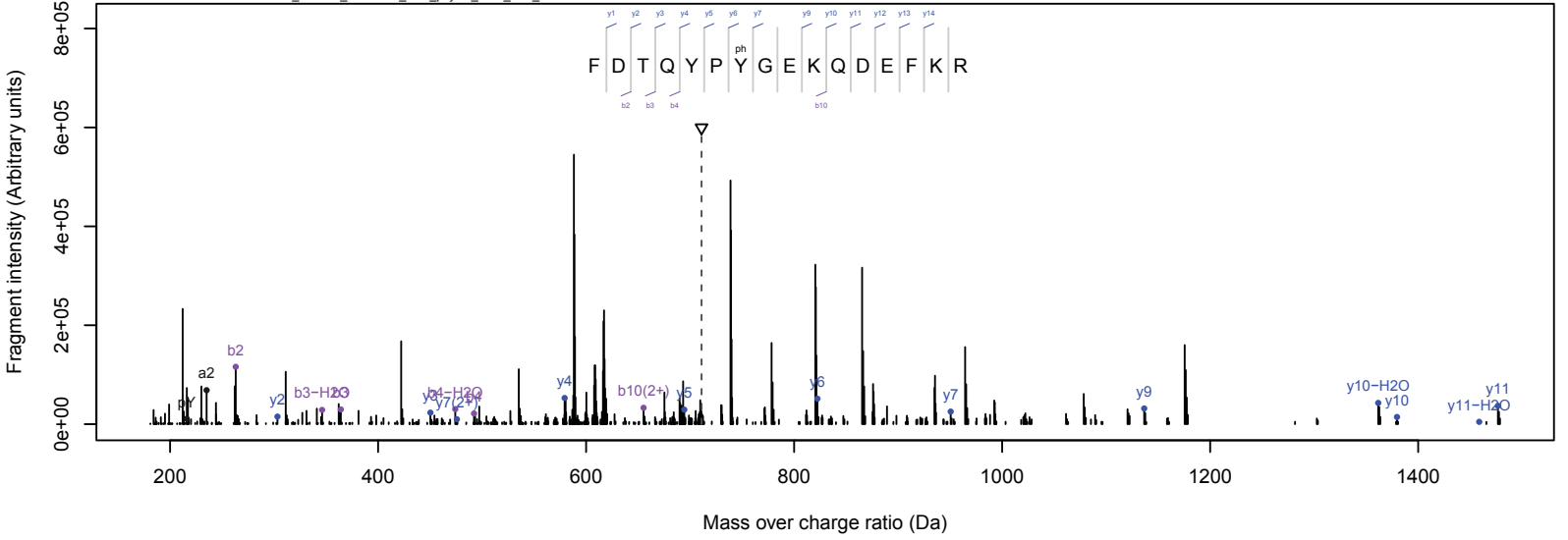
Andromeda score : 124.21



FDTQYPY(ph)GEKQDEFKR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 17926

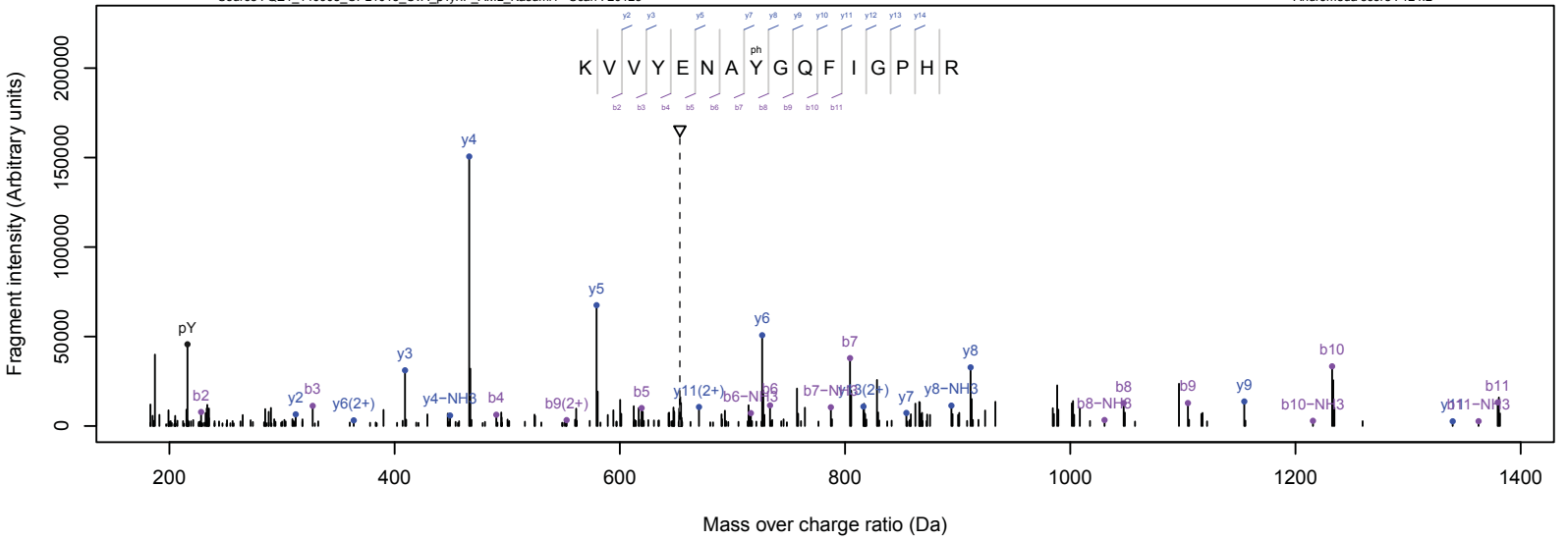
Andromeda score: 124.2



KVYENAY(ph)GQFIGPHR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 20123

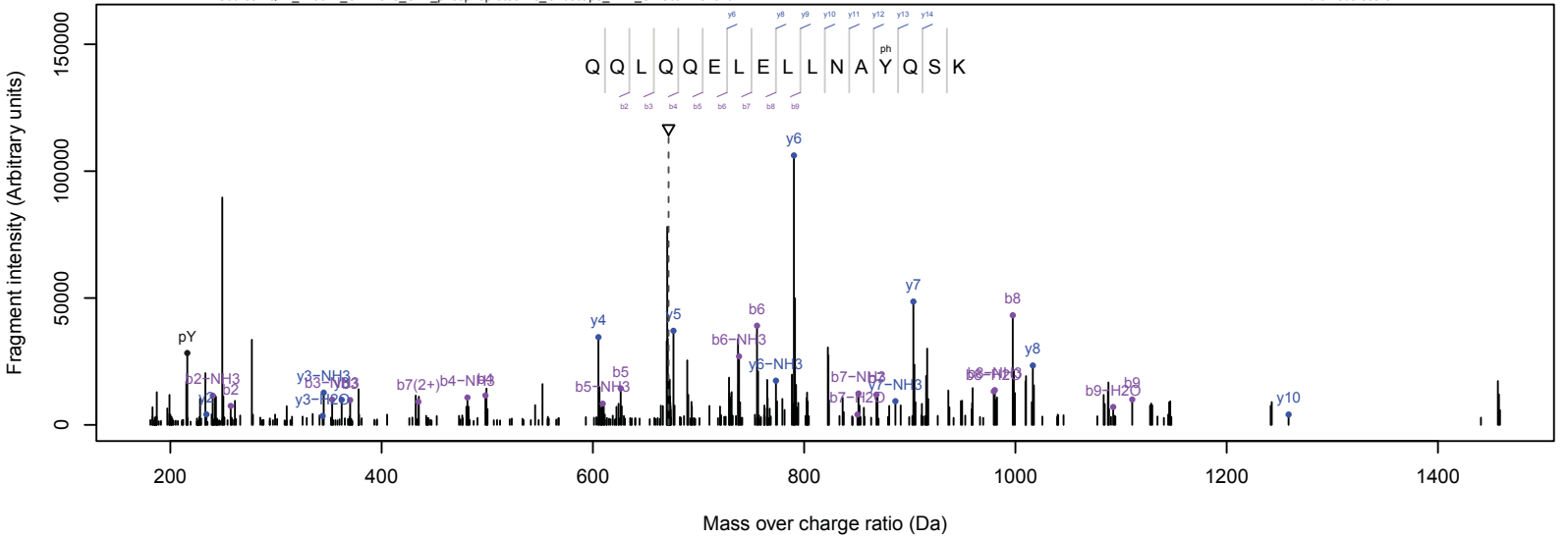
Andromeda score: 124.2



QLLQLELLNAY(ph)QSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37079

Andromeda score: 124.2

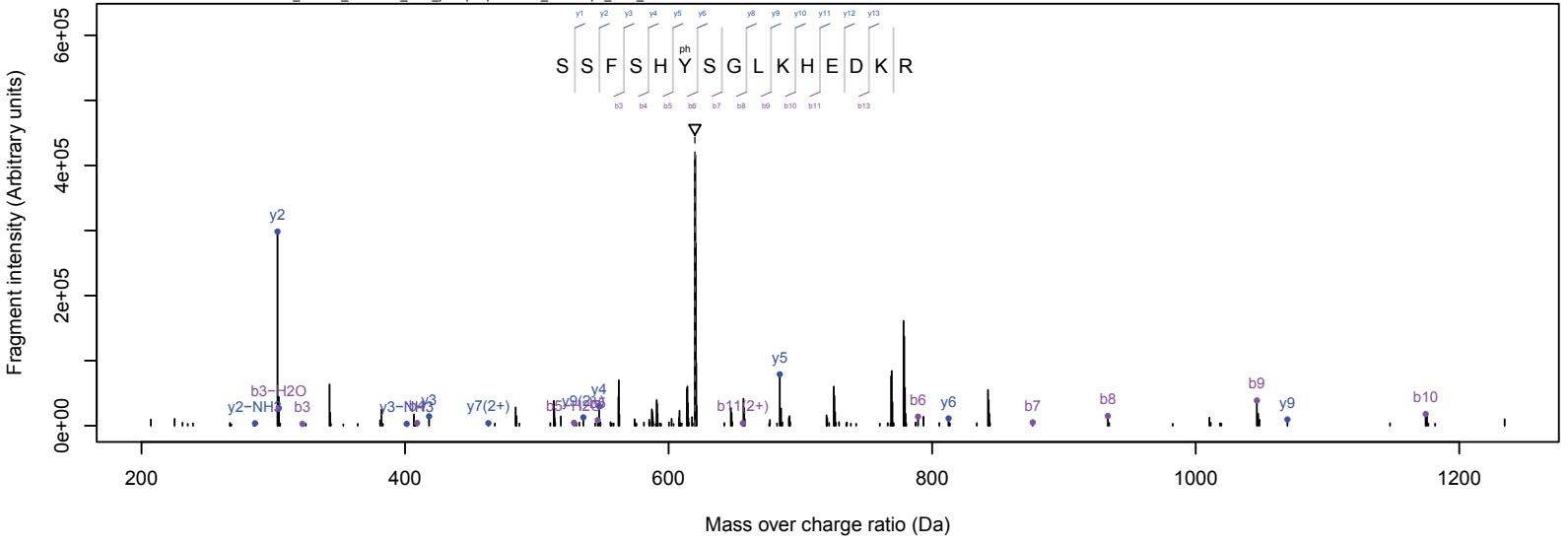




**SSFSHY(ph)SGLKHEDKR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 5360

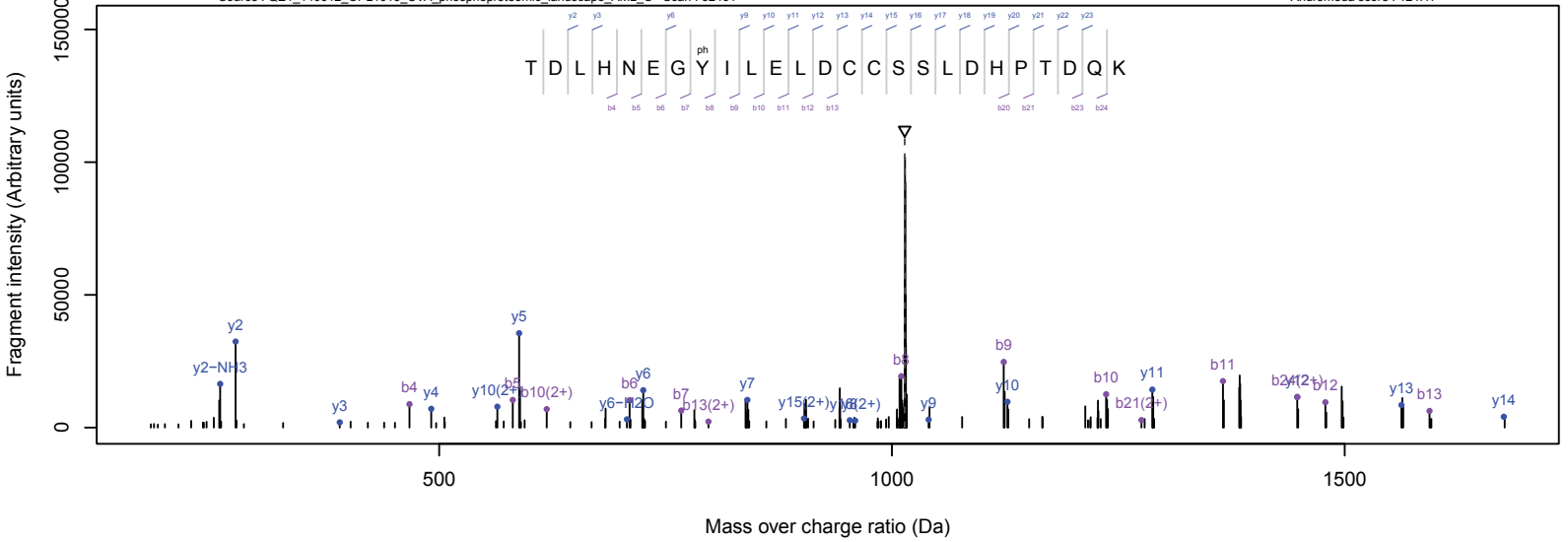
Andromeda score: 124.2



**TDLHNEGY(ph)ILELDCCSLDHPDQK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 32461

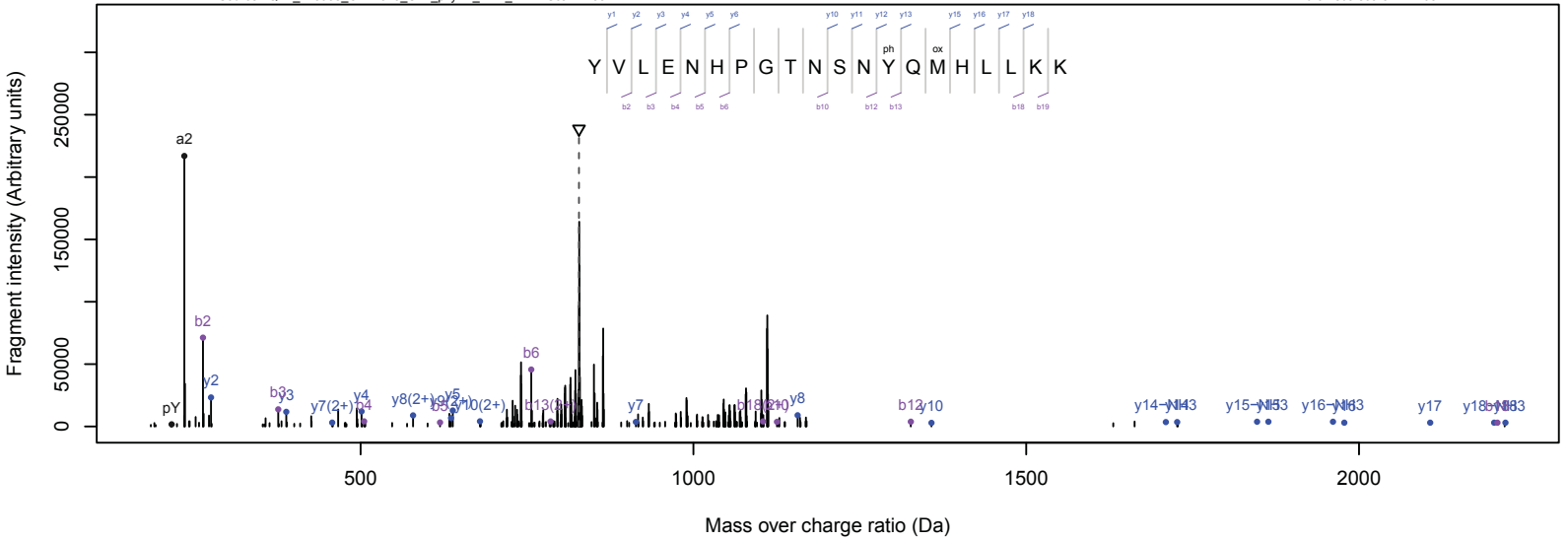
Andromeda score: 124.17



**YVLENHPG(TNSNY(ph)QM(ox))HLLKK-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 15571

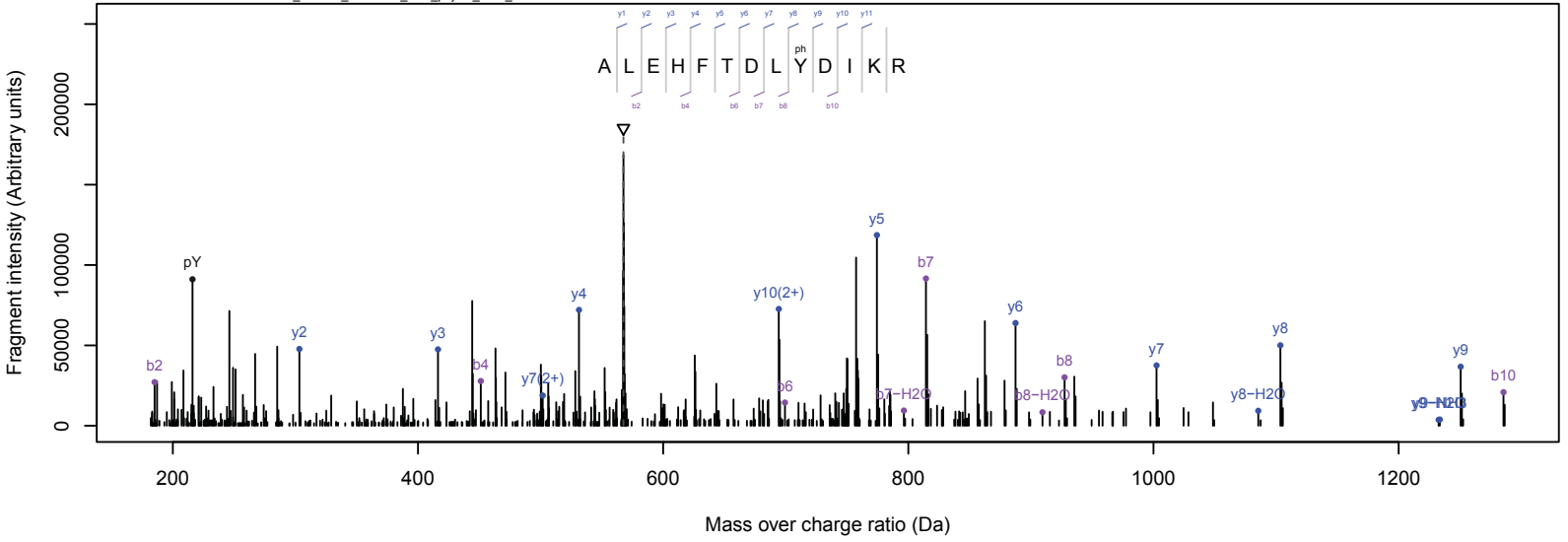
Andromeda score: 124.08



**ALEHFTDLYD<sup>ph</sup>IKR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 28572

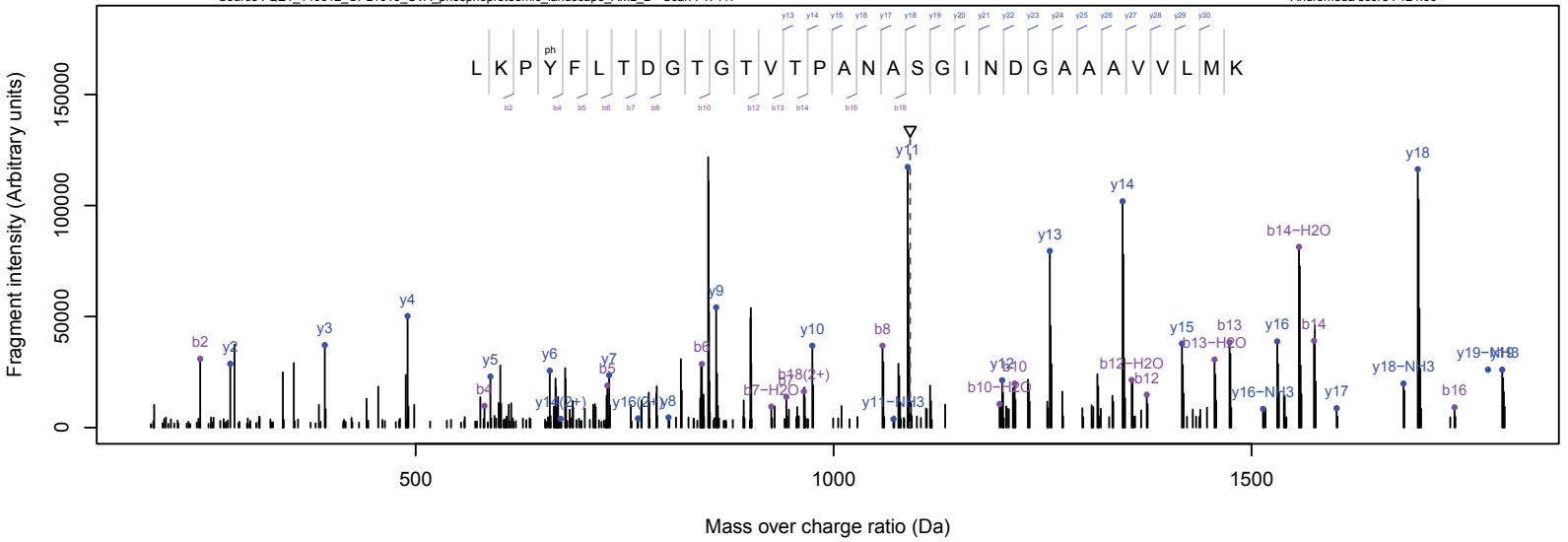
Andromeda score : 124.07



**LKPY<sup>ph</sup>FLTDGTGTVTPANASGINDGAAAVVLMK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 47417

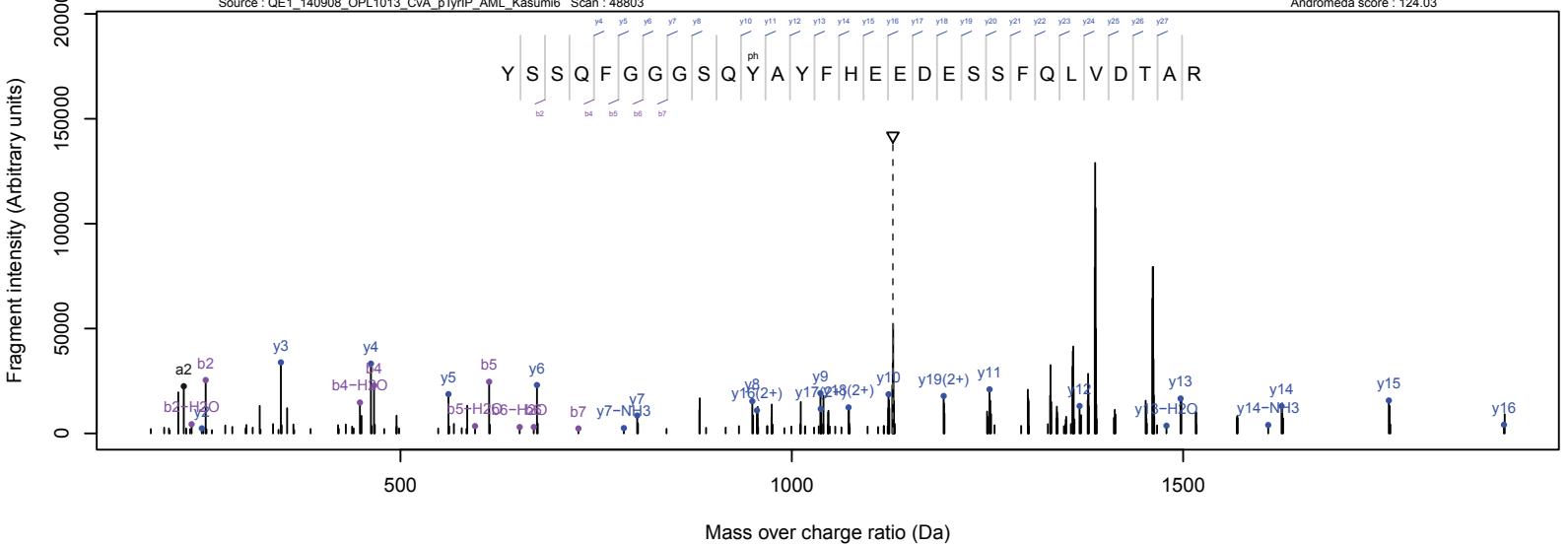
Andromeda score : 124.06



**YSSQFGGGSQY<sup>ph</sup>AYFHEEDESFLVD<sup>ph</sup>TAR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 48803

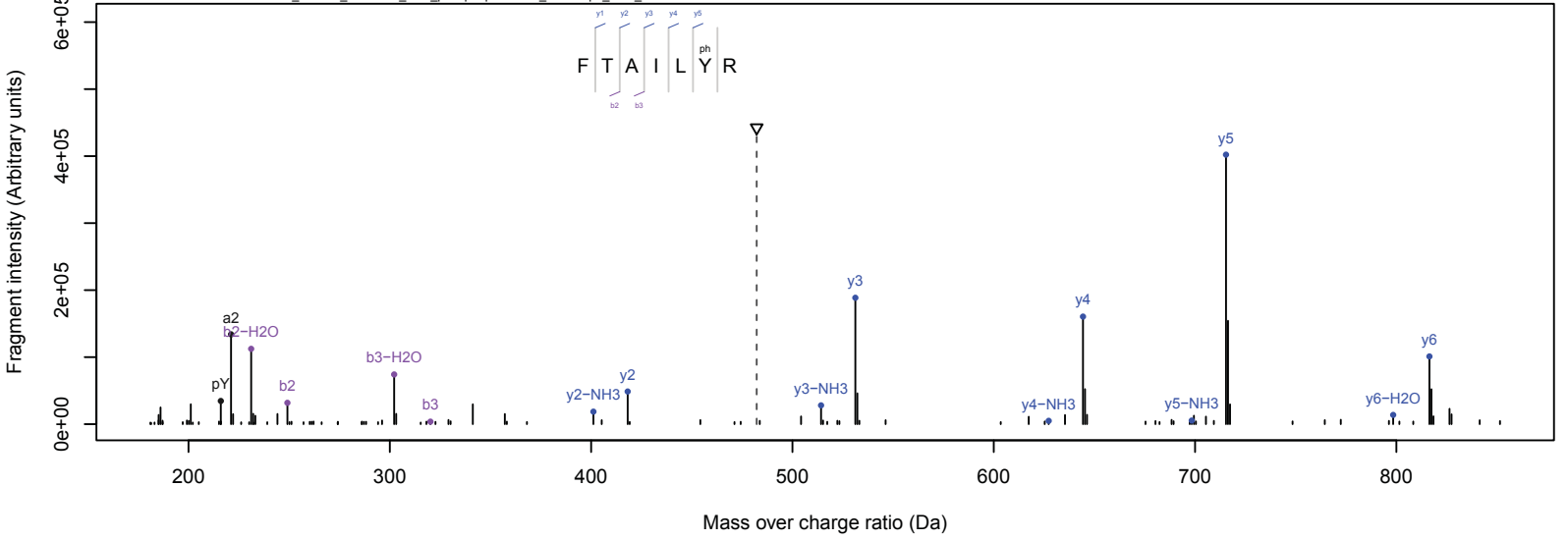
Andromeda score : 124.03



FTAILY(ph)R-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25050

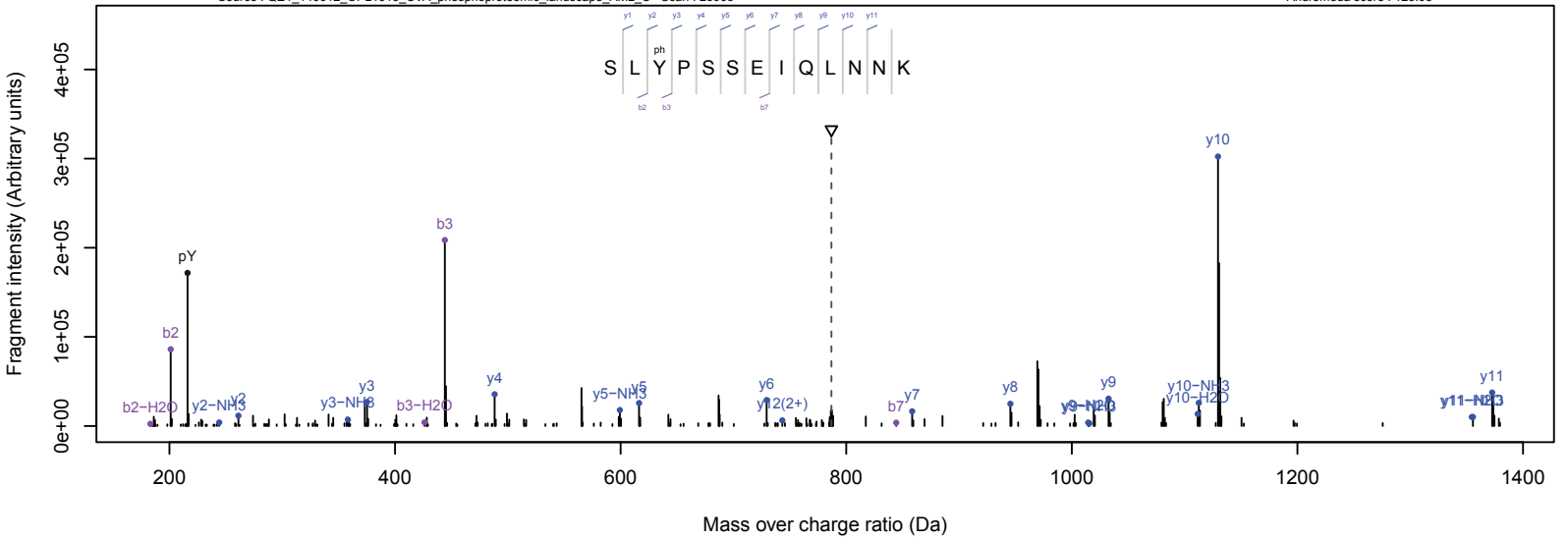
Andromeda score: 123.99



SLY(ph)PSSEIQLNKK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23963

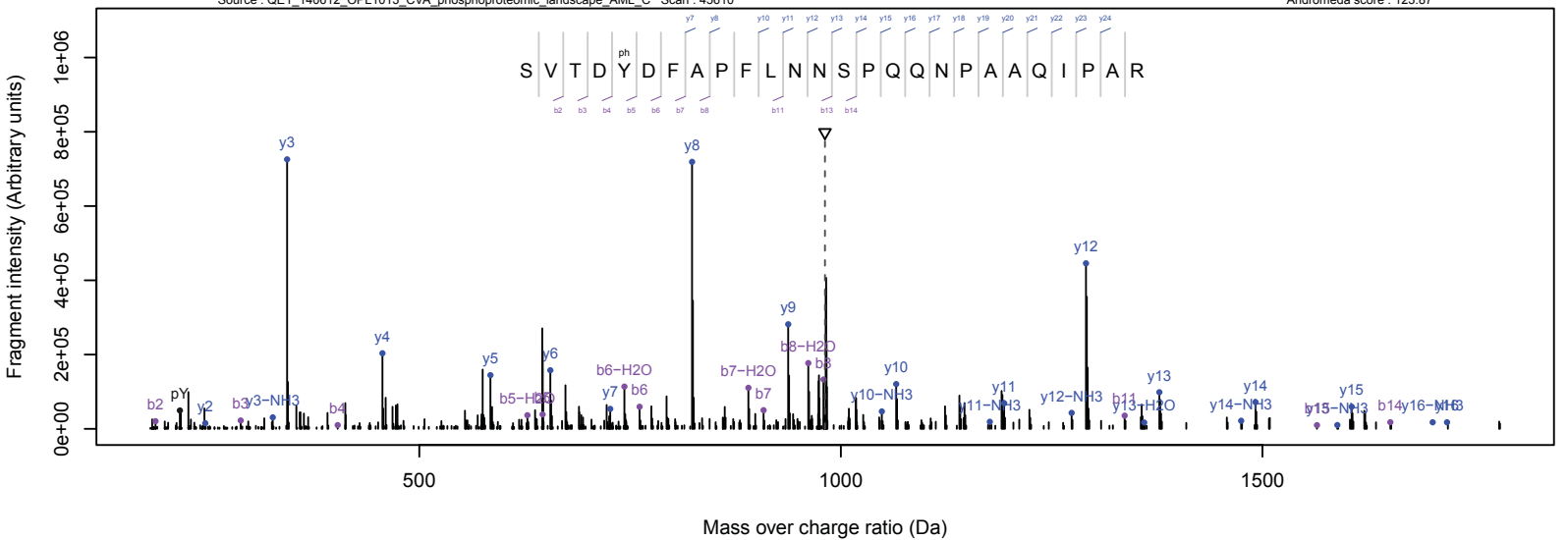
Andromeda score: 123.95



SVTDY(ph)DFAPFLNNSPQQNPAAQIPAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 45610

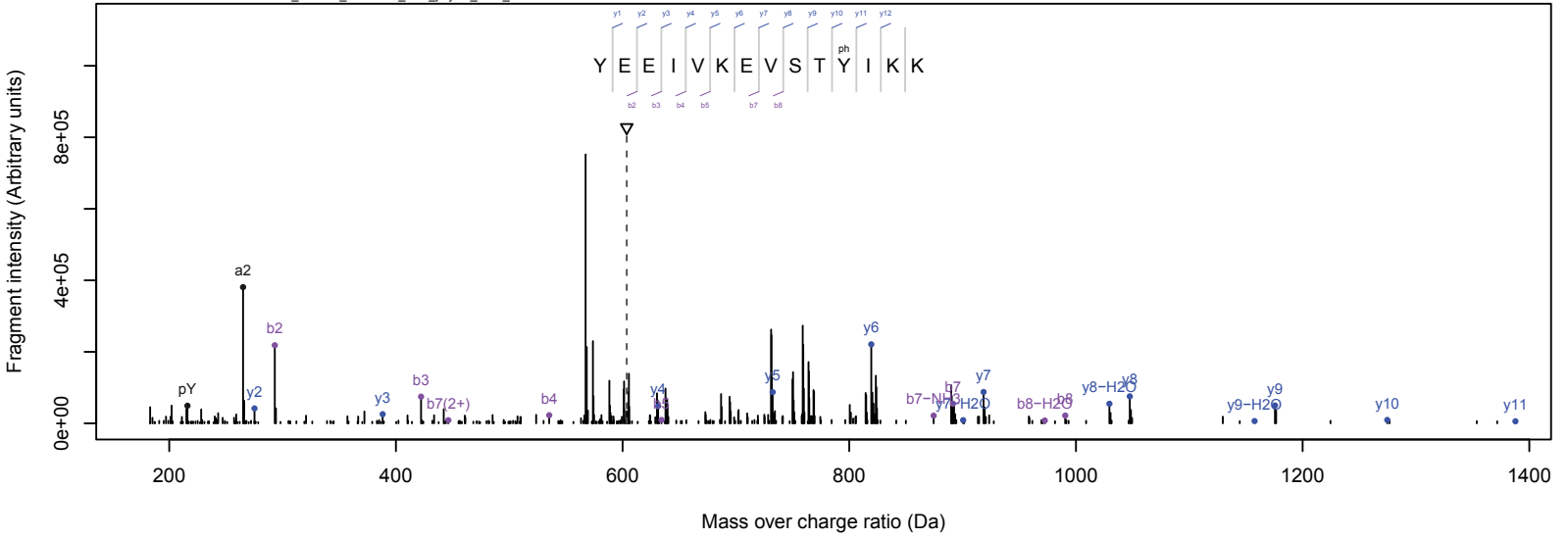
Andromeda score: 123.87



**YEEIVKEVSTY(ph)IKK-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 27547

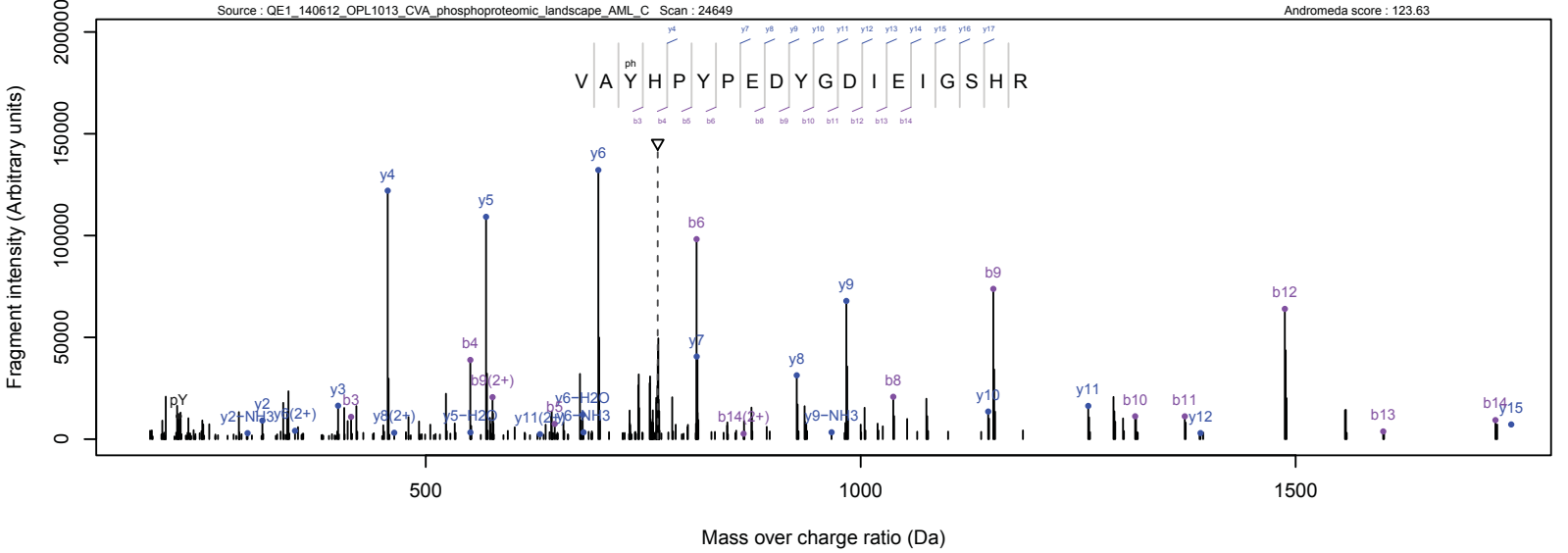
Andromeda score : 123.76



**VAY(ph)HPYPEDYGDIEIGSHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24649

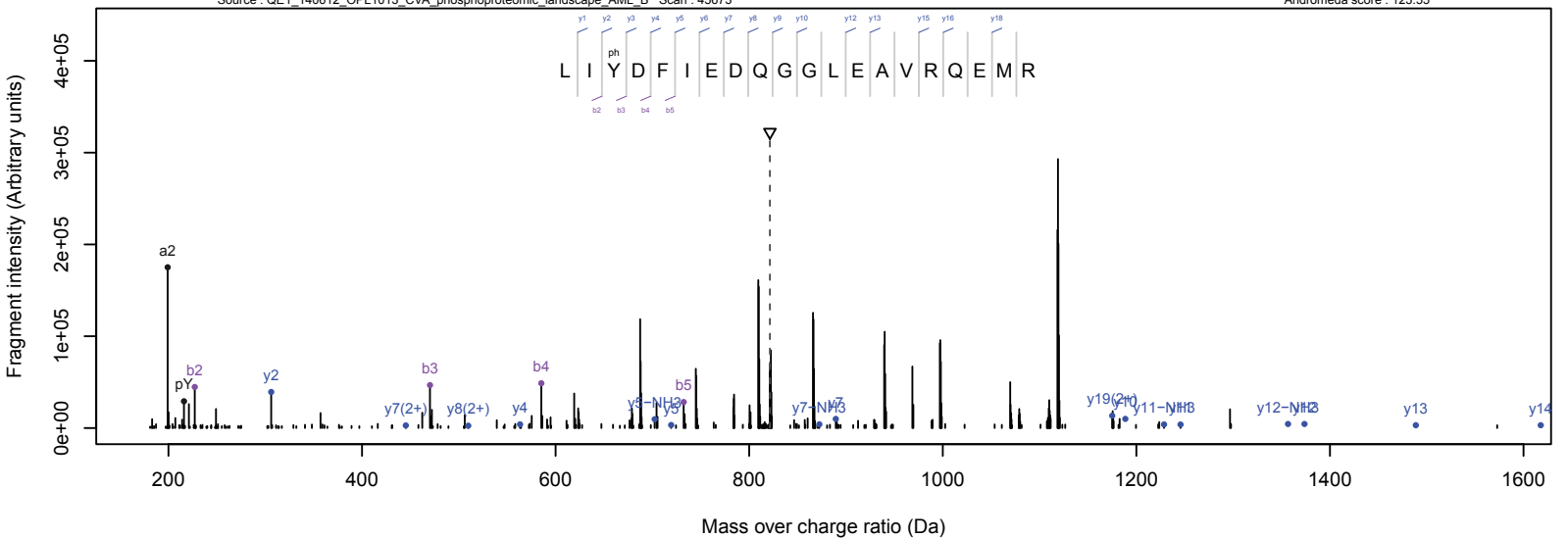
Andromeda score : 123.63

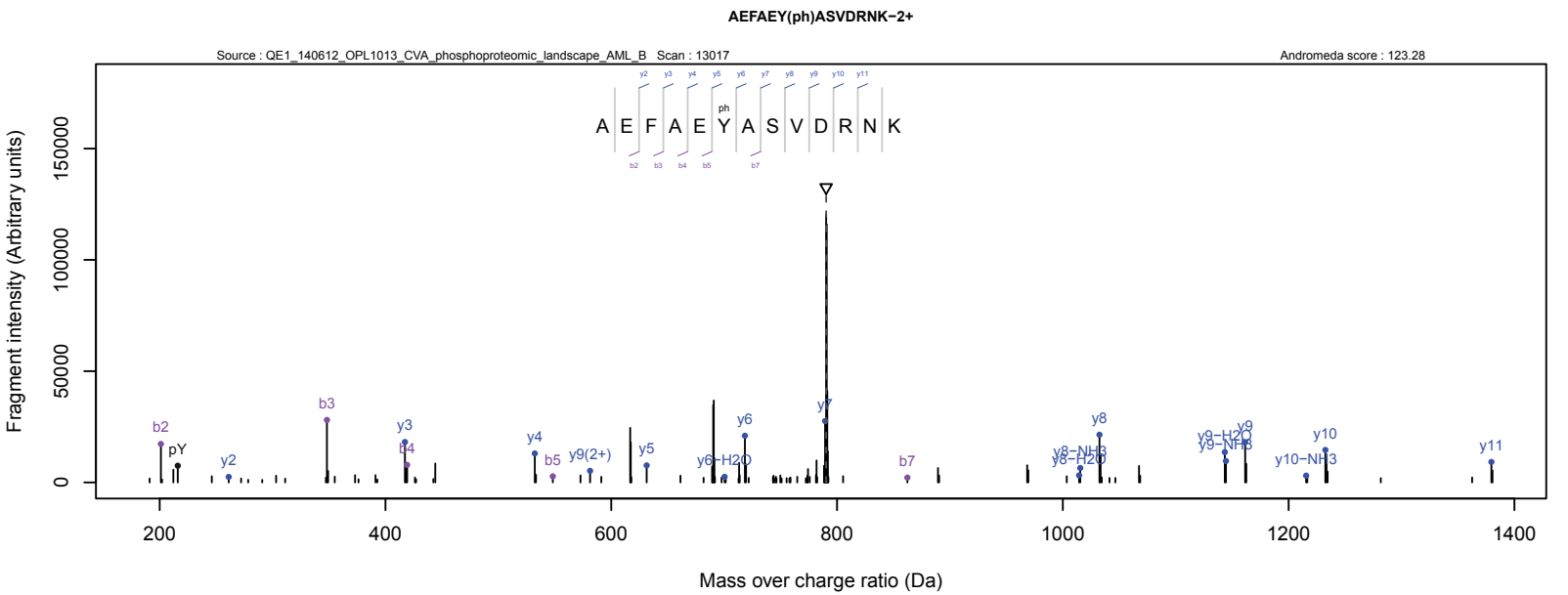
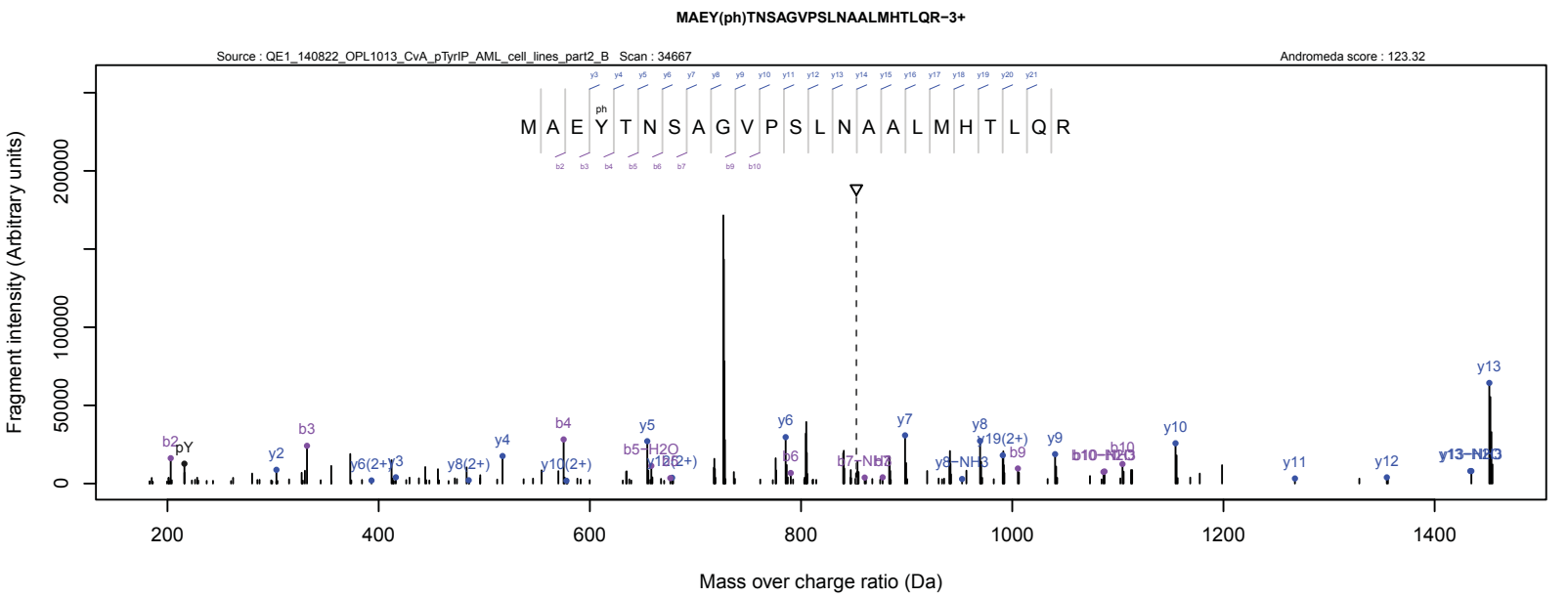
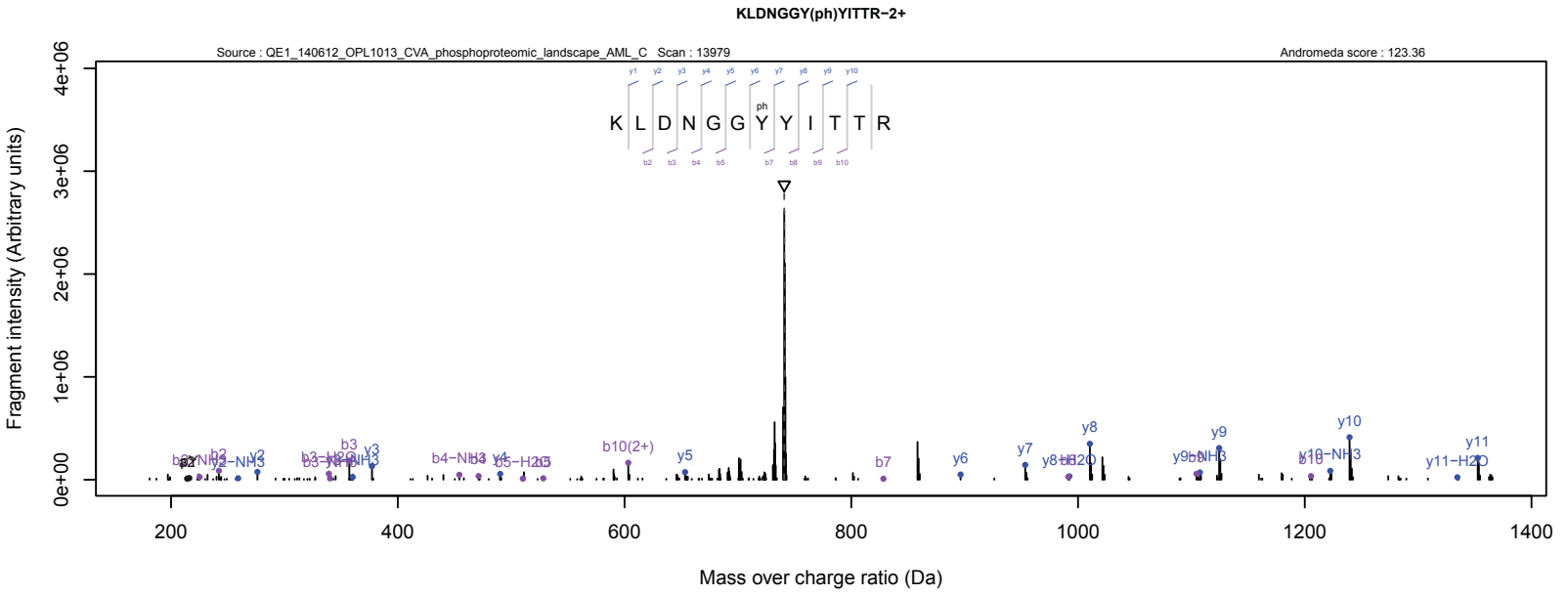


**LIY(ph)DFIEDQGGLEAVRQEMR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45673

Andromeda score : 123.55

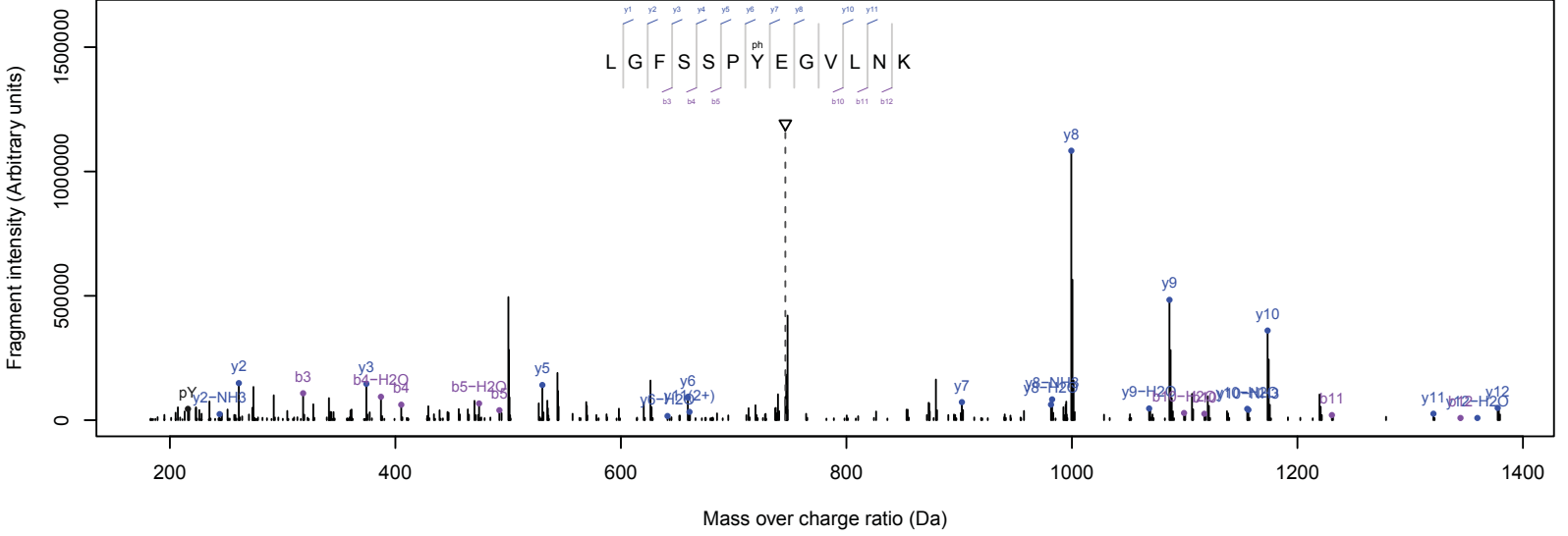




**LGFSSPY(ph)EGLNLK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 29714

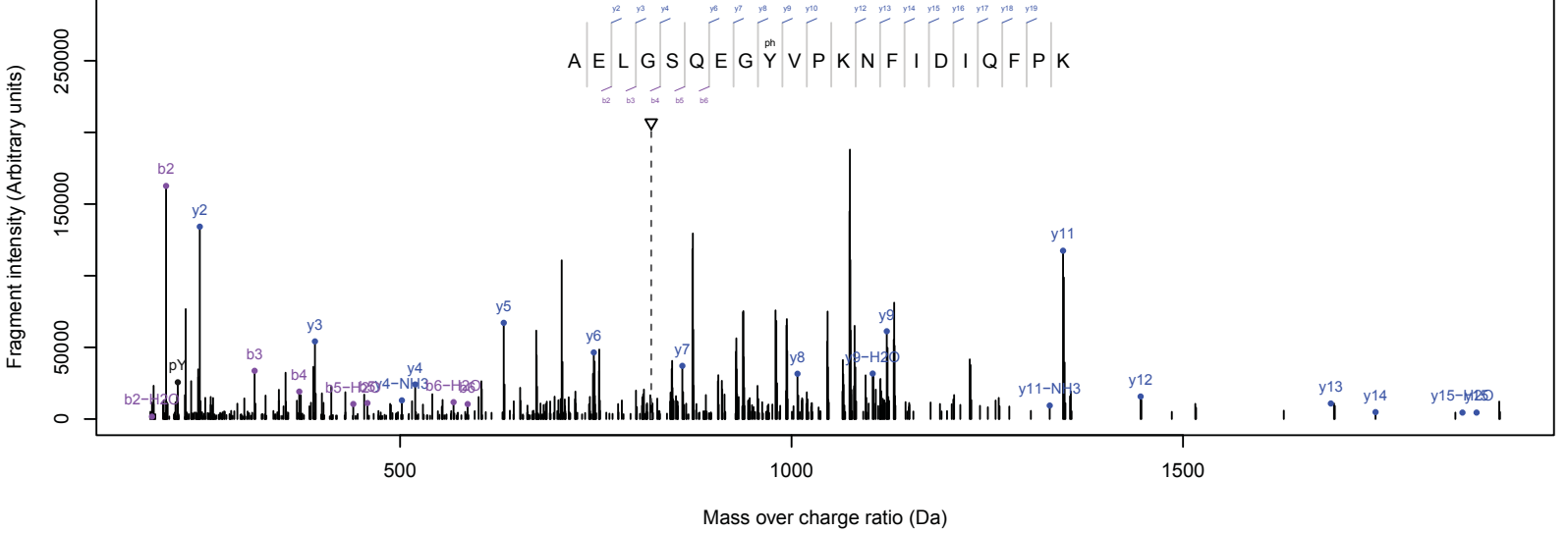
Andromeda score: 123.28



**AELGSQEGY(ph)VPKNFIDIQFPK-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 44899

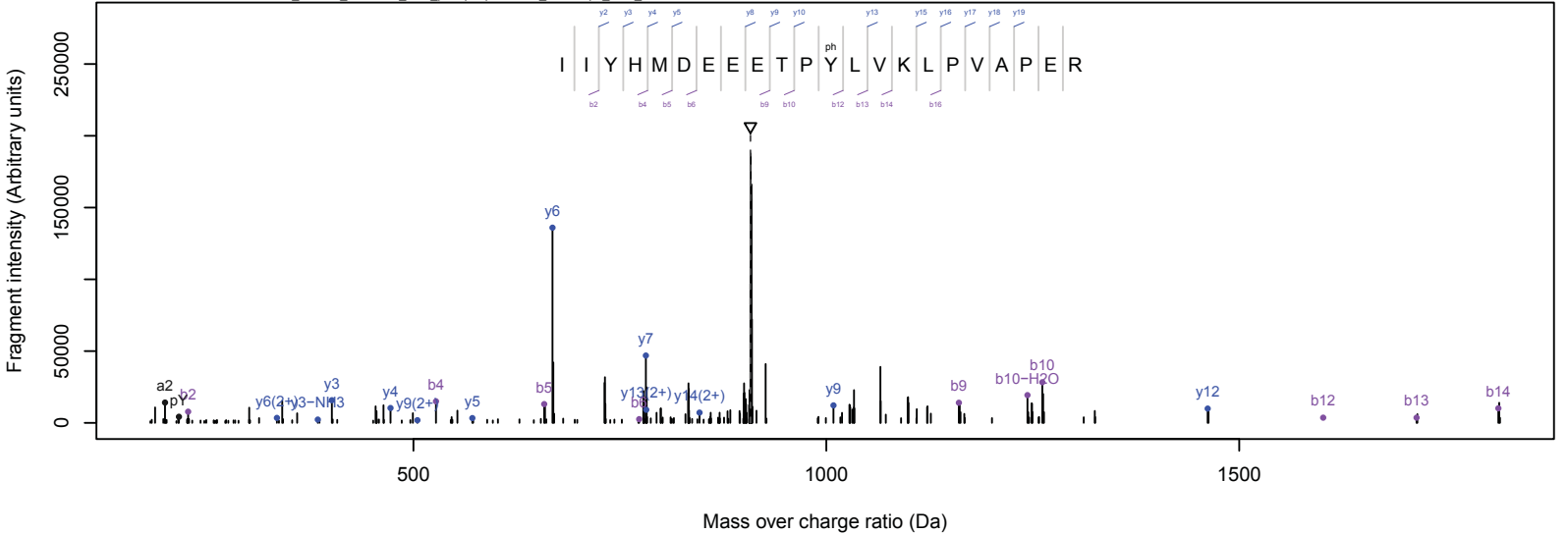
Andromeda score: 123.19



**IYHMDEETPY(ph)LVKLPVAPER-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 39481

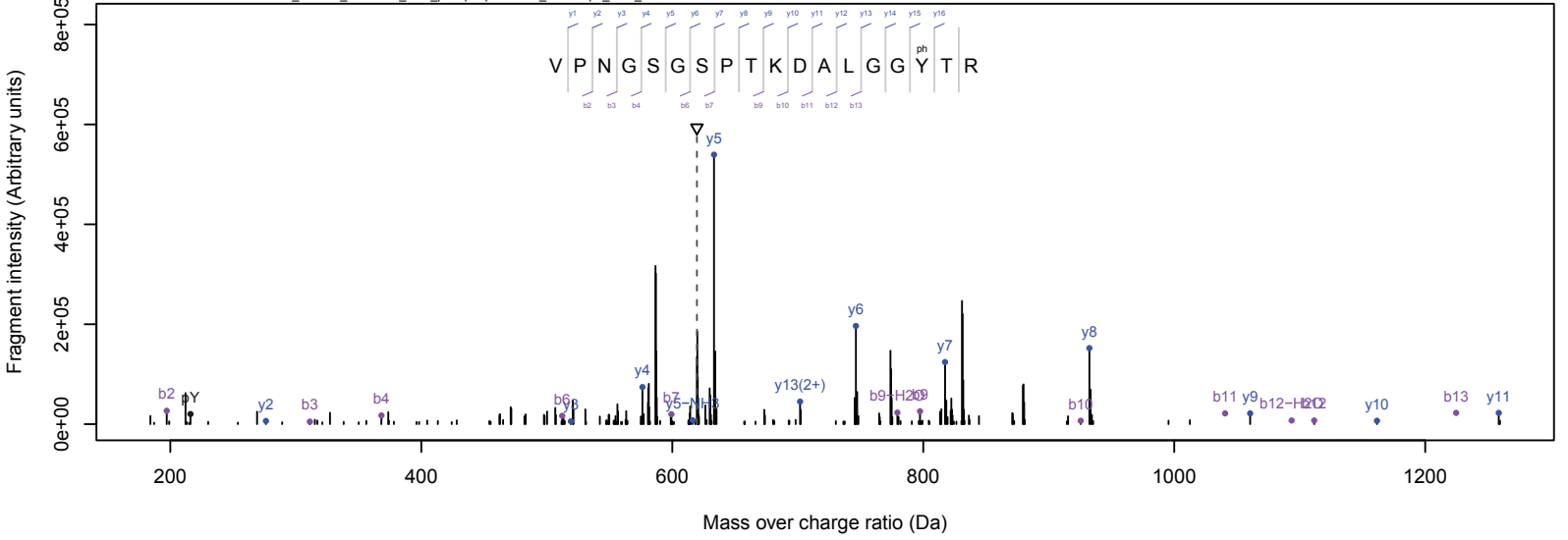
Andromeda score: 123.19



VPNGSGSPDKALGGYTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10824

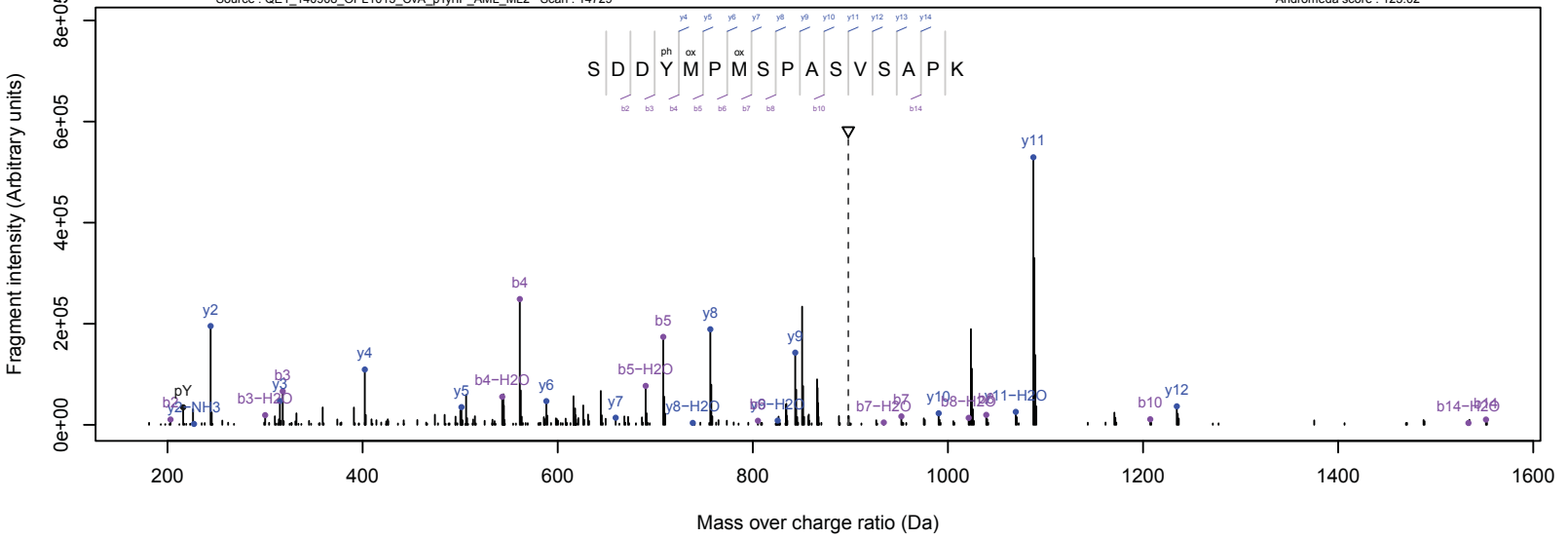
Andromeda score : 123.19



SDDY(ph)M(ox)PM(ox)SPASVSAPK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 14729

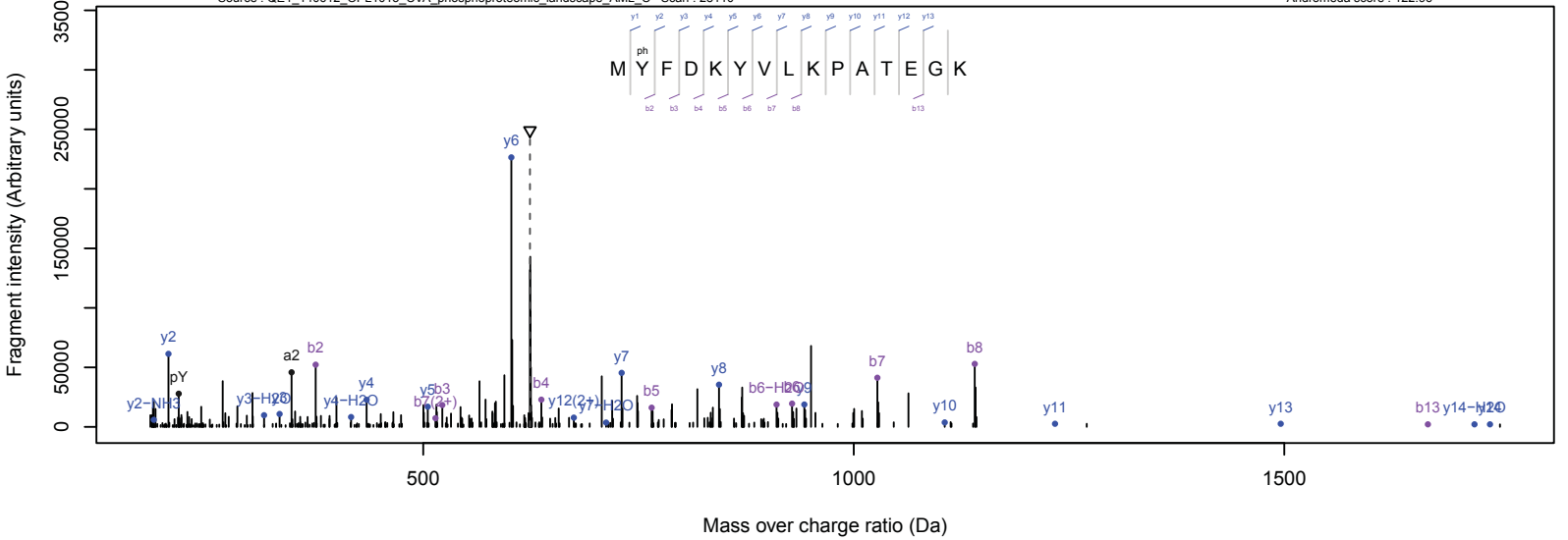
Andromeda score : 123.02



MY(ph)FDKYVLKPALEGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28110

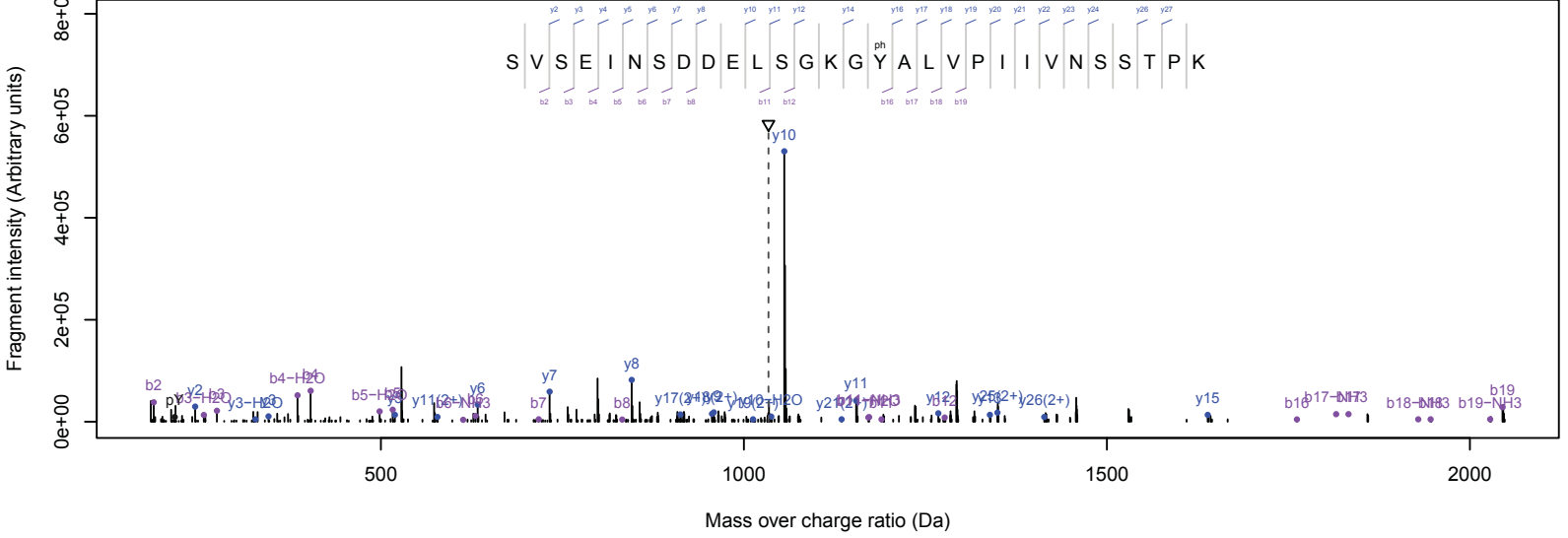
Andromeda score : 122.96



SVSEINSDDELSGKGY(ph)ALVPIIVNSSTPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42456

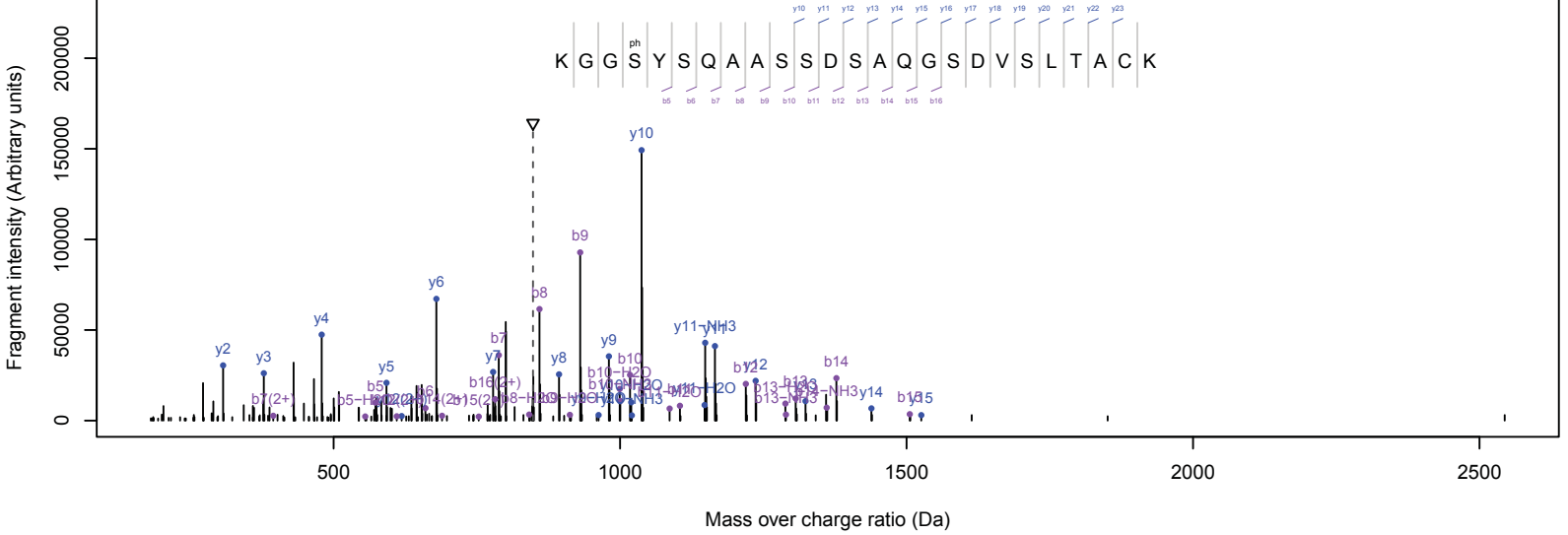
Andromeda score : 122.89



KGGs(ph)YSQAASSDSAQGSVDVSLTACK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 13419

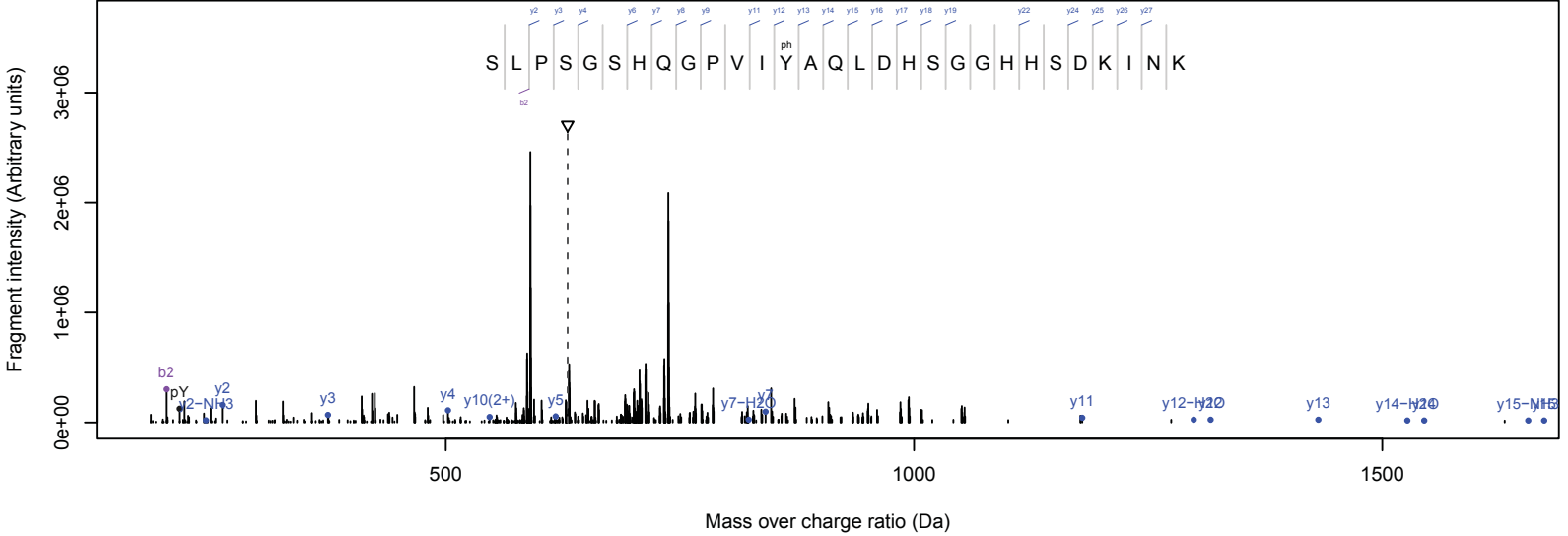
Andromeda score : 122.84



SLPSGSHQGPVIY(ph)AQLDHSGGHSDKINK-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15488

Andromeda score : 122.8

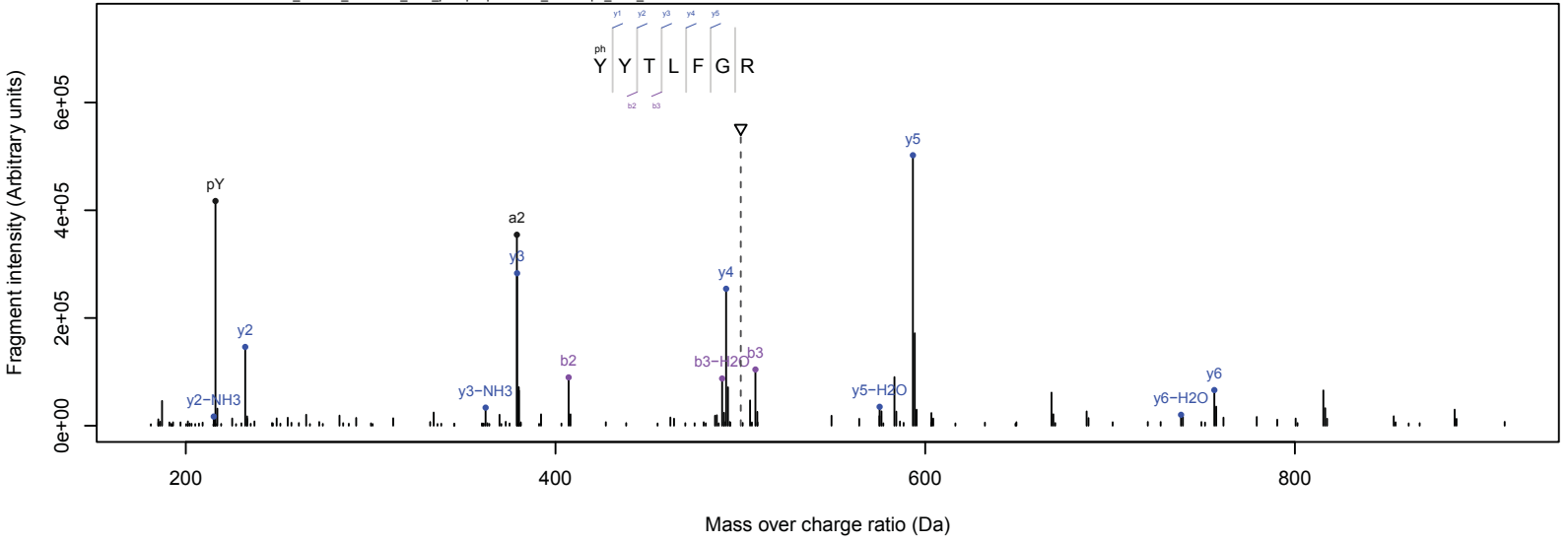




Y(ph)YTLFGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28393

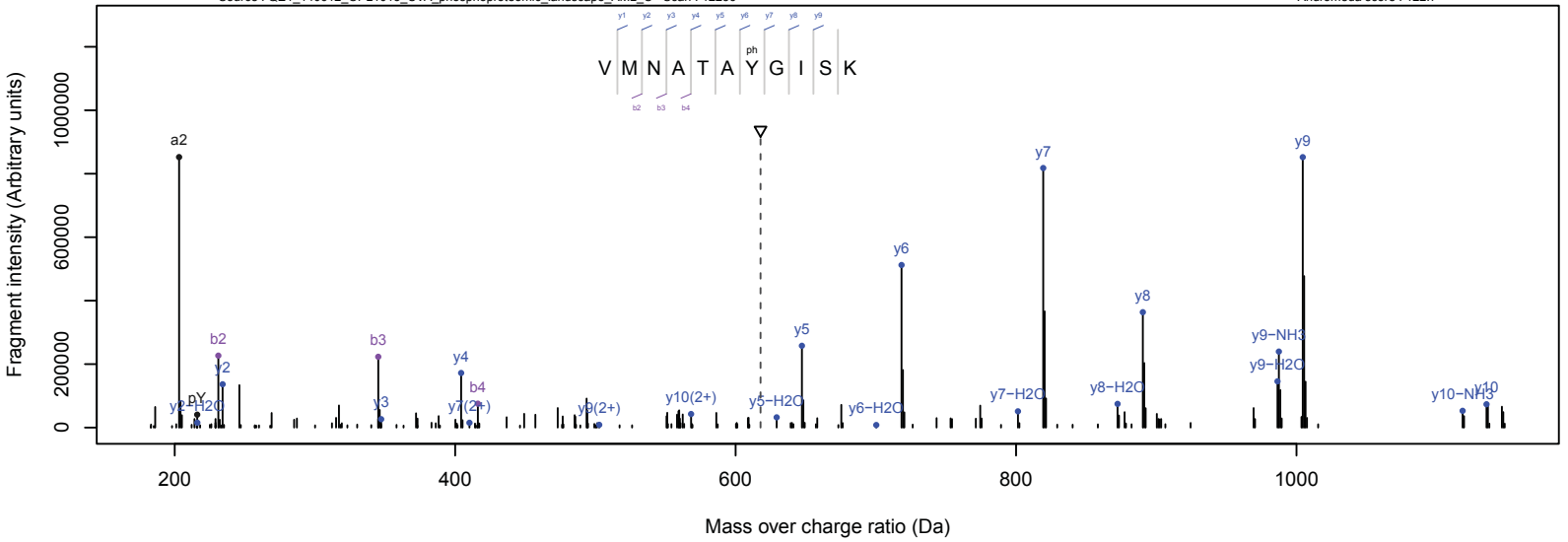
Andromeda score : 122.79



VMNATAY(ph)GISK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12256

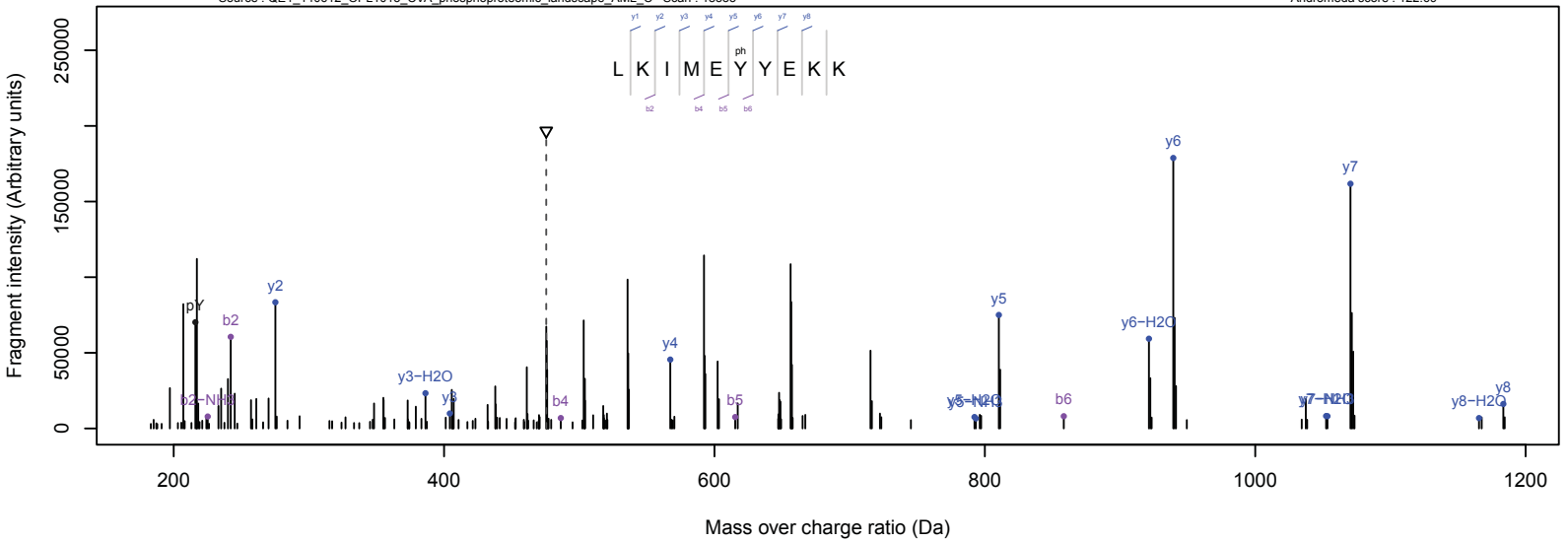
Andromeda score : 122.7



LKIMEY(ph)YEKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18353

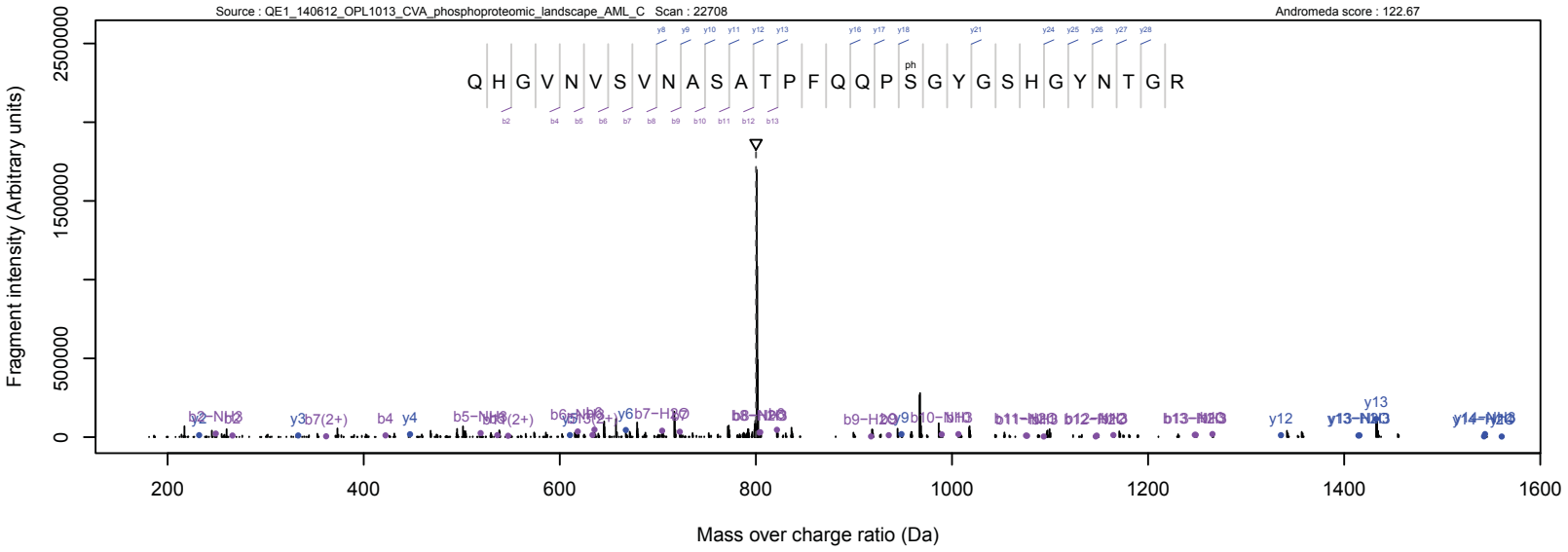
Andromeda score : 122.69



**QHGVNVSINASATPFQQPS(ph)YGSHGYNTGR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22708

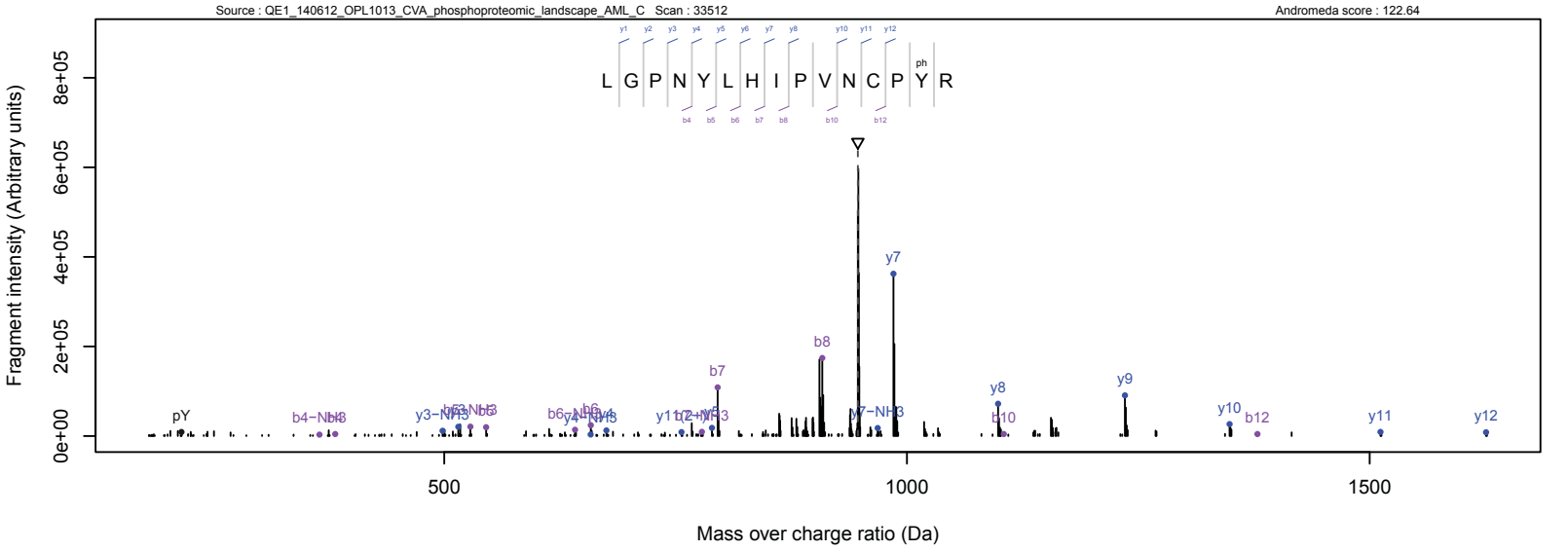
Andromeda score : 122.67



**LGPNYLHIPVNCYP(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33512

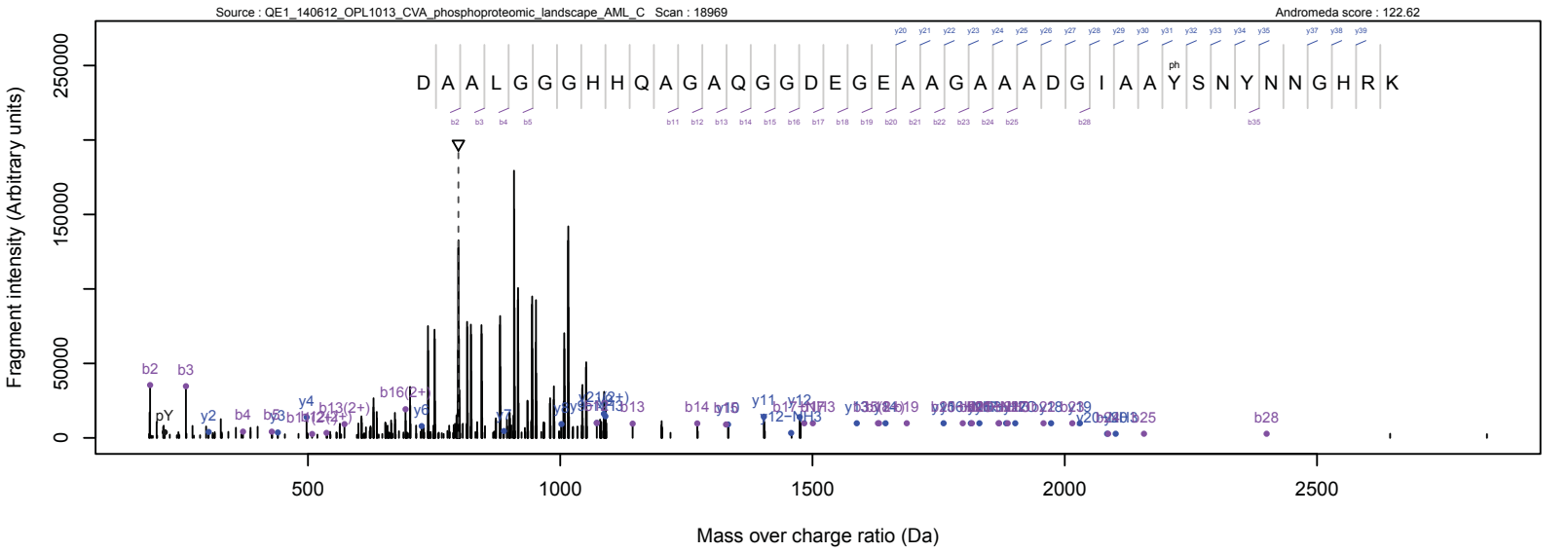
Andromeda score : 122.64



**DAALGGHHQAGQGDEGEAAGAADGIAAY(ph)SNYNGHRK-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18969

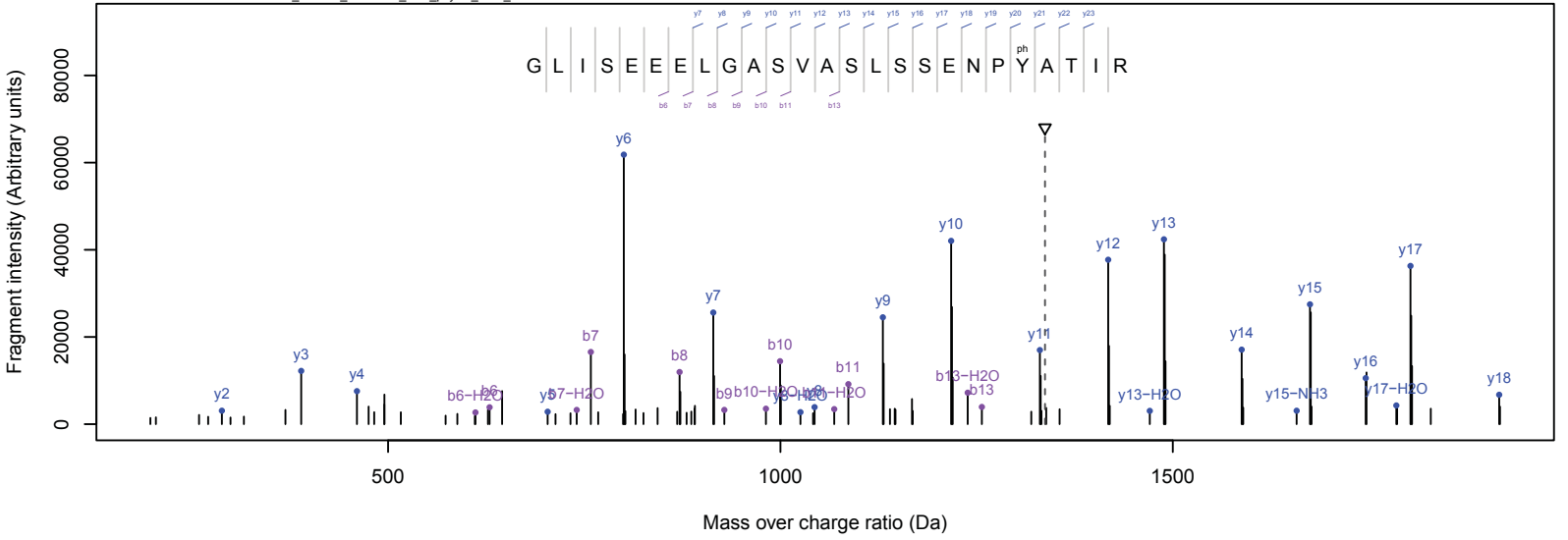
Andromeda score : 122.62



**GLISEEELGASVASLSENPY(ph)ATIR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 47105

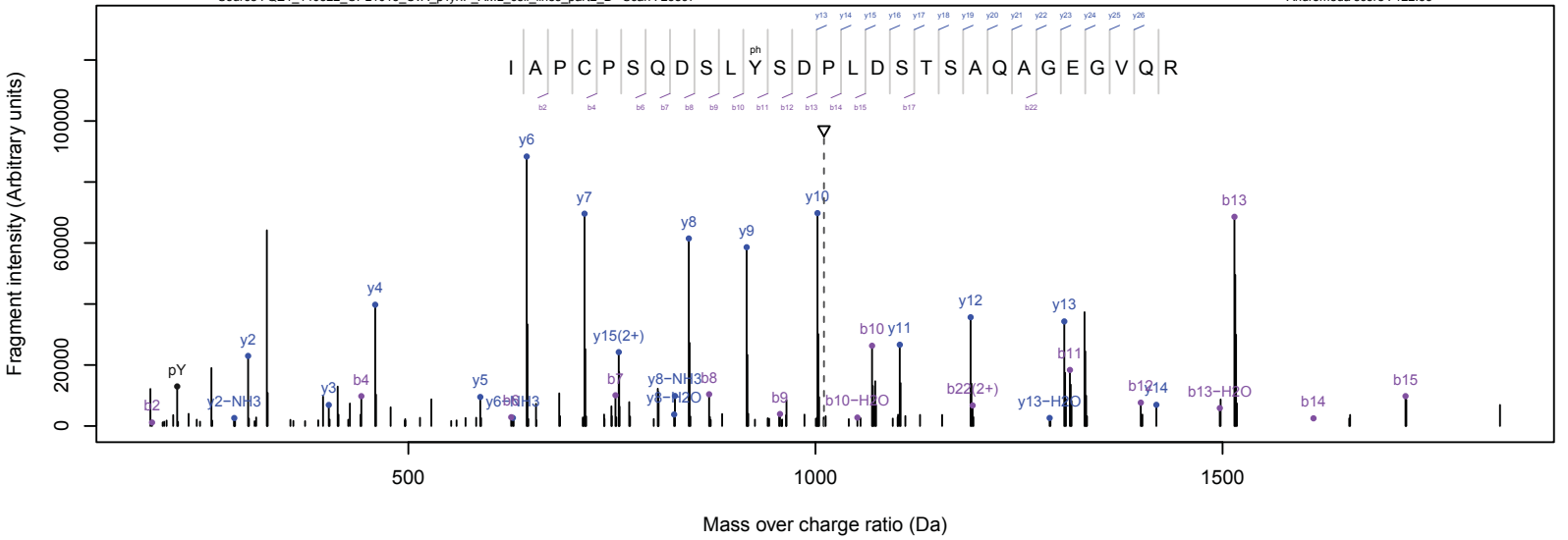
Andromeda score : 122.6



**IAPCPSQDSL(ph)SDPLDSTSAQAGEGVQR-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 23897

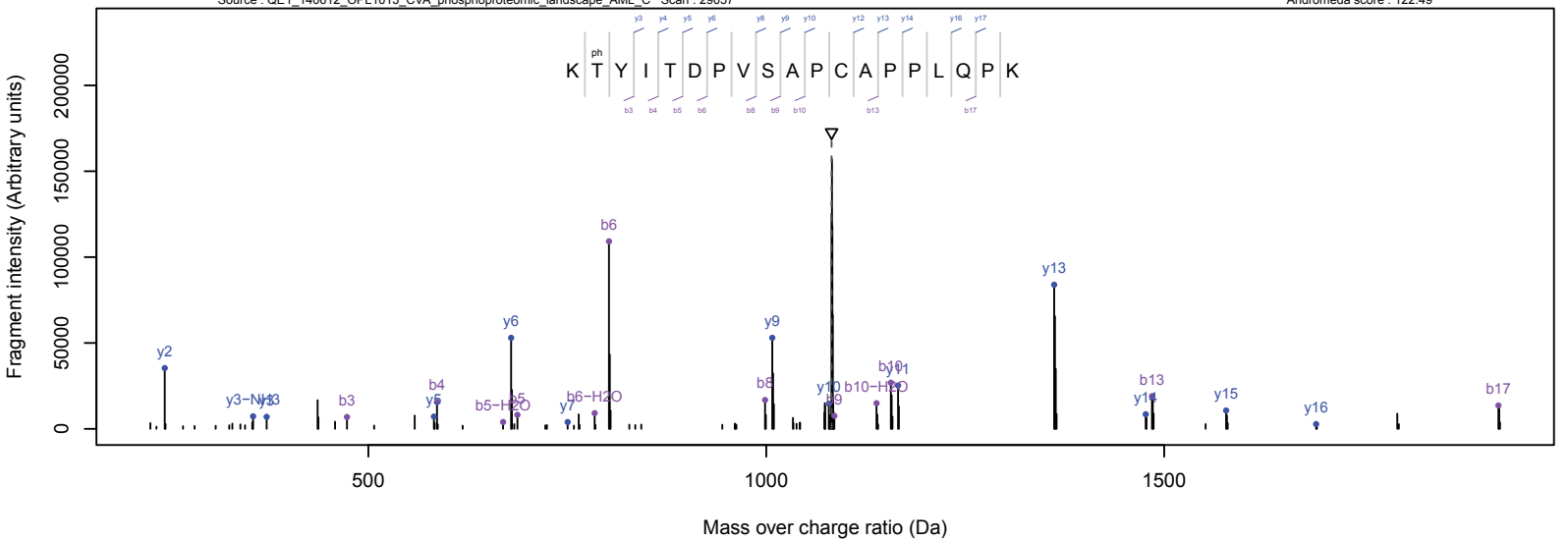
Andromeda score : 122.53



**KT(ph)YITDPVSAPCAPPLQPK-2+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 29057

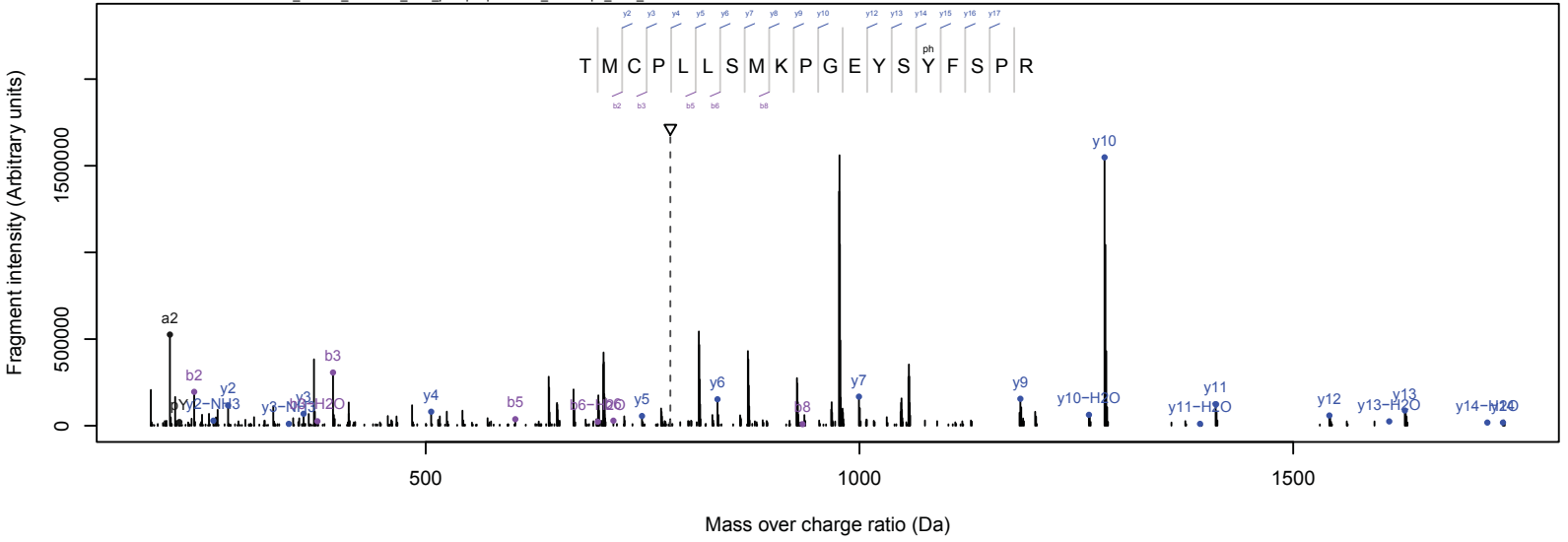
Andromeda score : 122.49



**TMCPLLSMKPGEYS(ph)FSPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40872

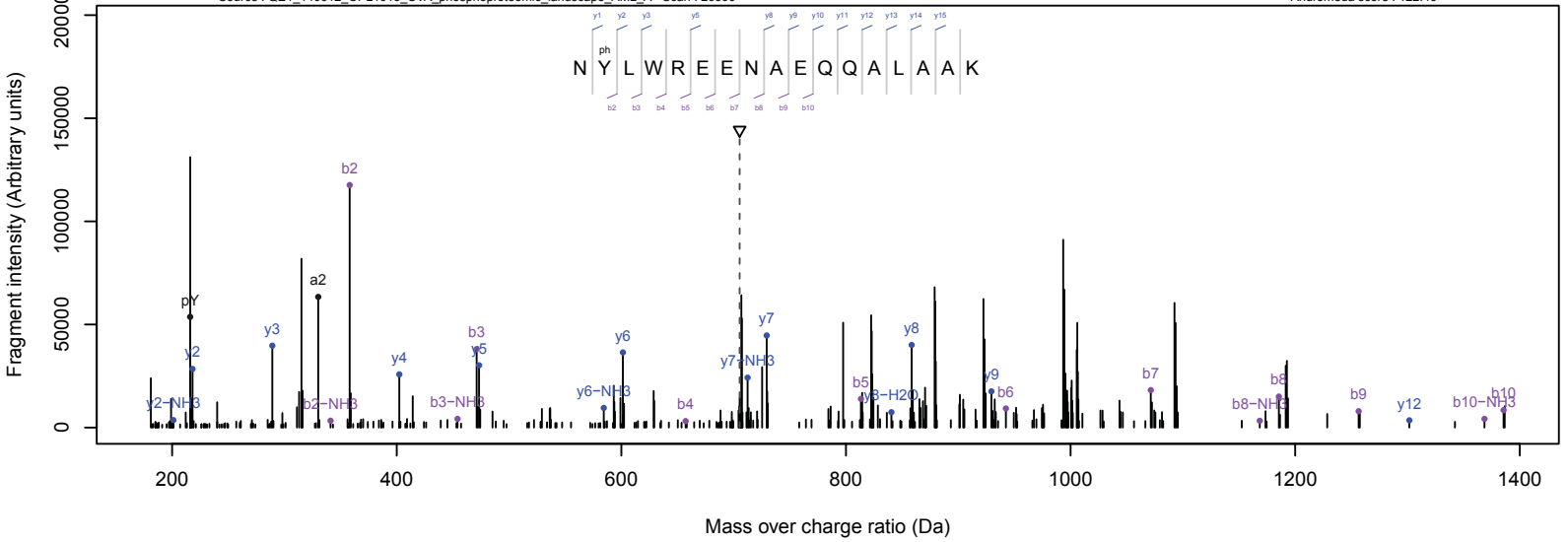
Andromeda score : 122.49



**NY(ph)LWREENAQQALAAK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26390

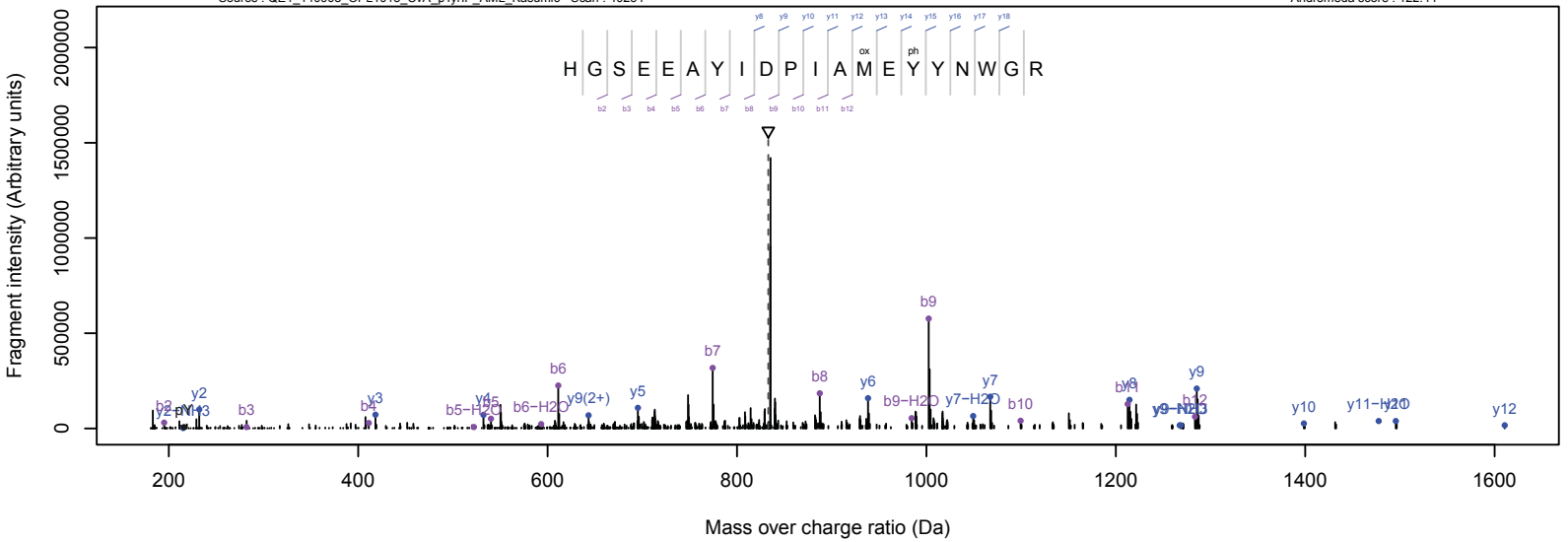
Andromeda score : 122.46



**HGSEEAYIDPIAEMEY(ph)YNWGR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 40254

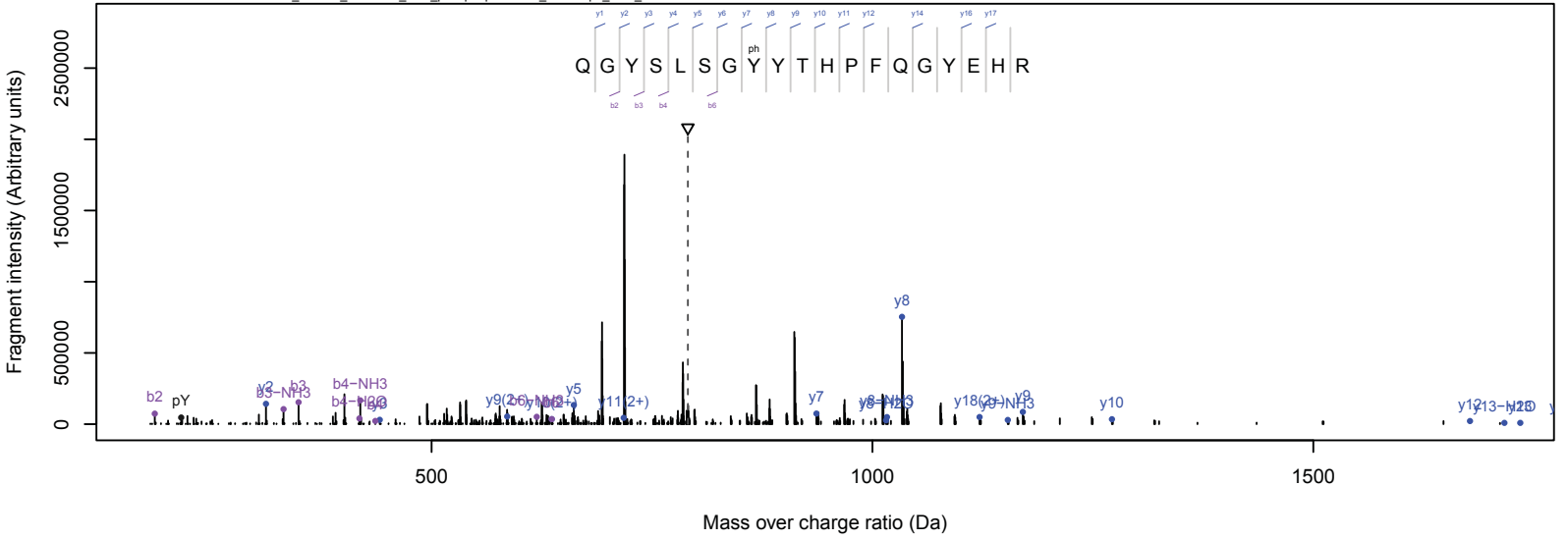
Andromeda score : 122.44



**QGYSLSGY(ph)YTHPFQGYEHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28085

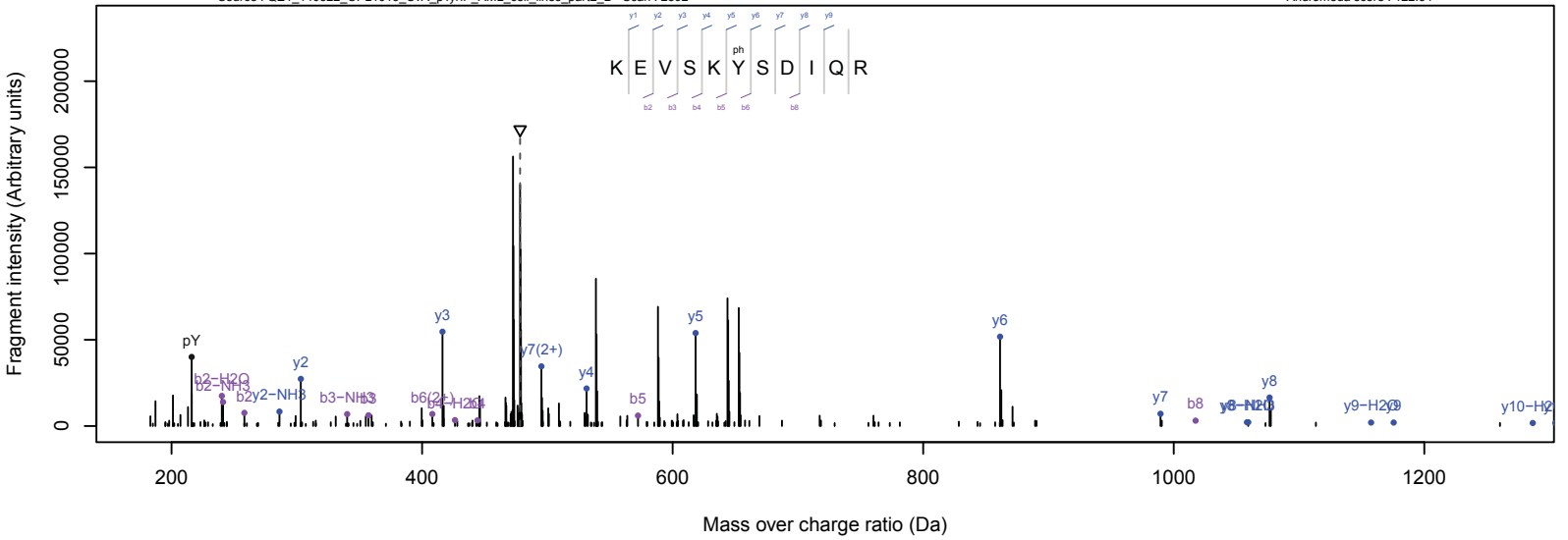
Andromeda score : 122.44



**KEVSKY(ph)SDIQR-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 2892

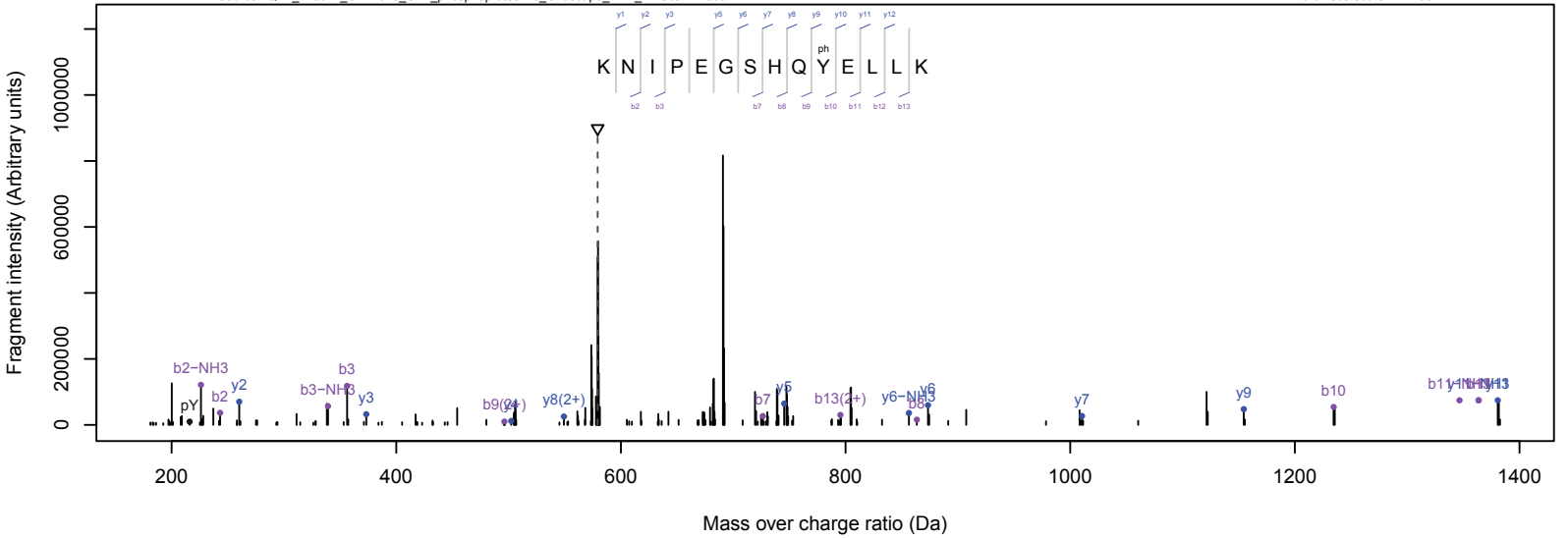
Andromeda score : 122.34



**KNIEGSHQY(ph)ELLK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13331

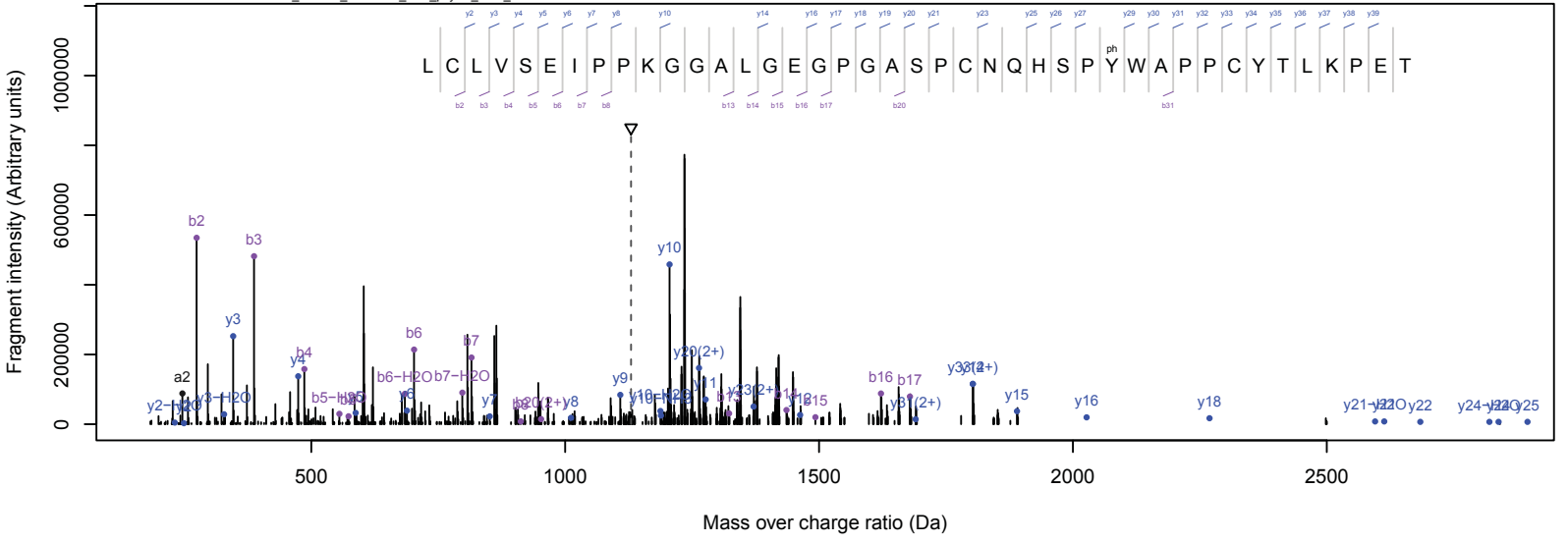
Andromeda score : 122.33



LCLVSEIPPKGALGEGPGASPCNQHSPLYWAPPCTLKPET-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 43455

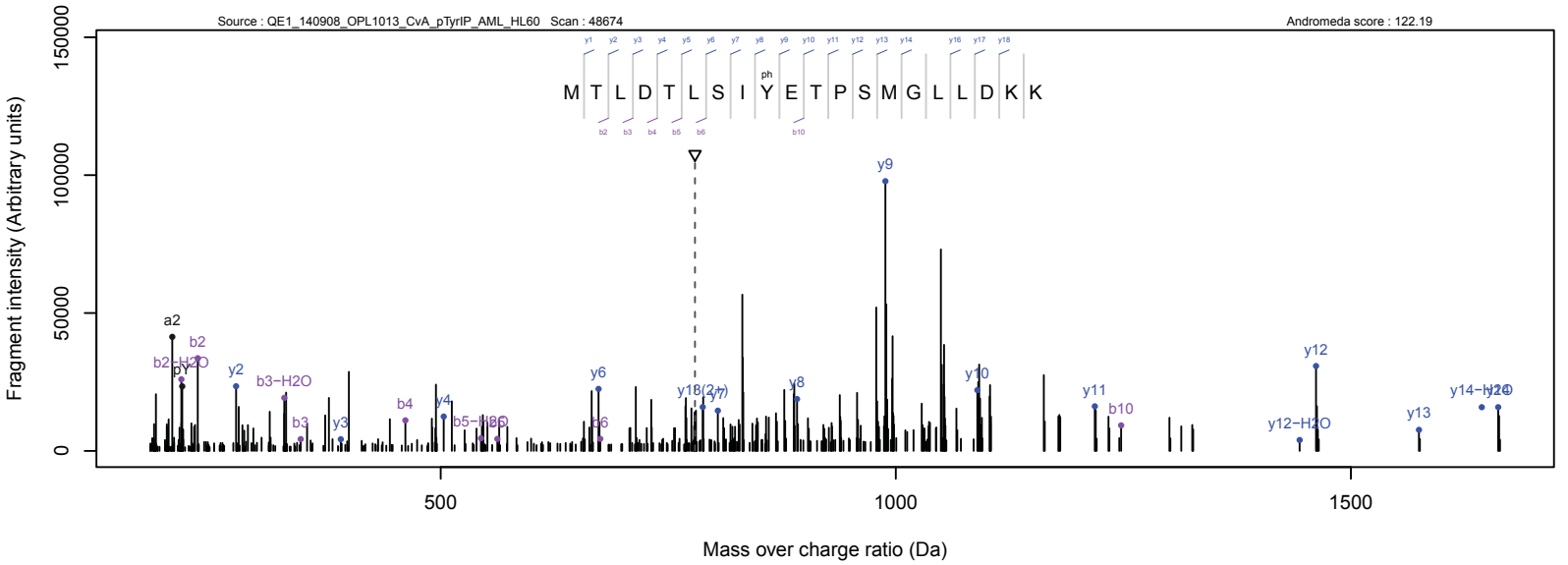
Andromeda score : 122.3



MTLDTLSIY(ph)ETPSMGLLDKK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 48674

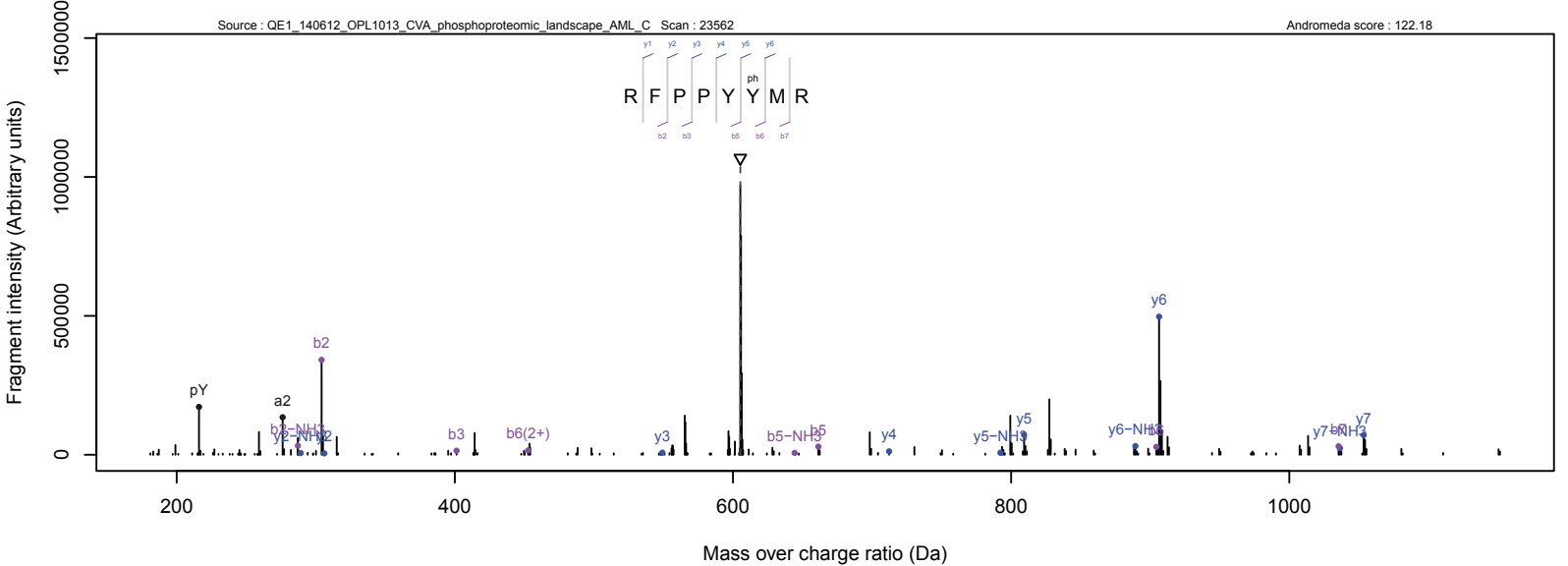
Andromeda score : 122.19



RFPPYY(ph)MR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23562

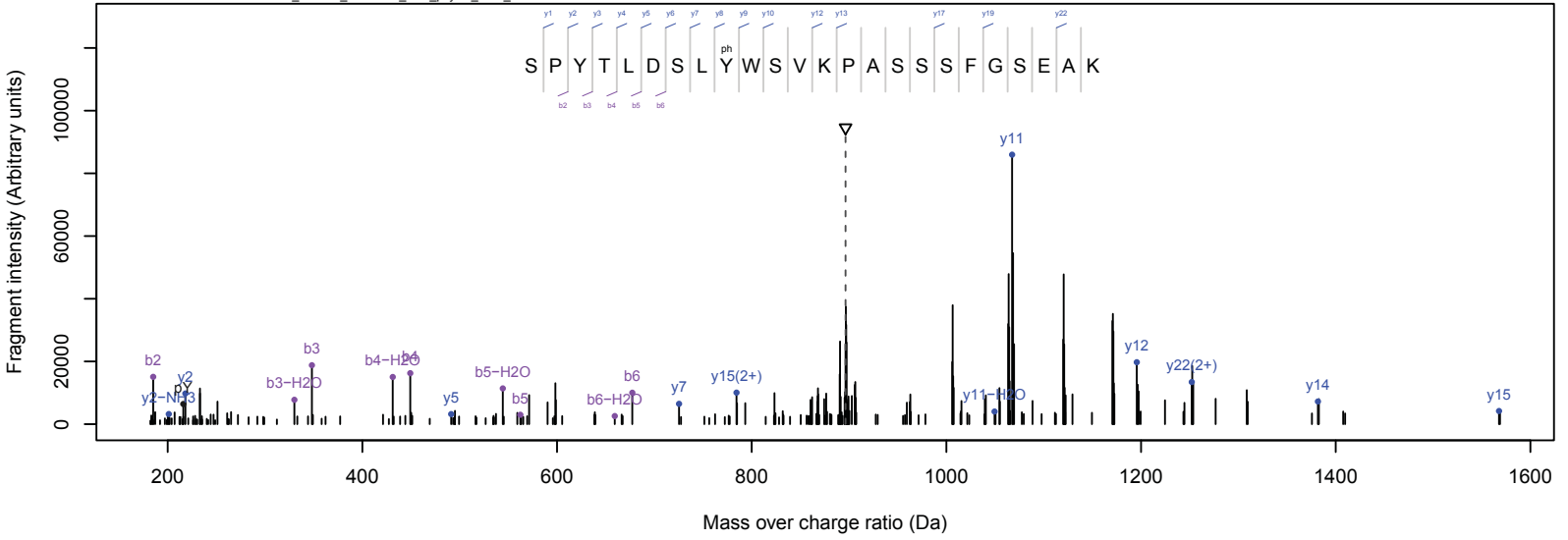
Andromeda score : 122.18



SPYTLDSL(ph)WSVKPASSFGSEAK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 43604

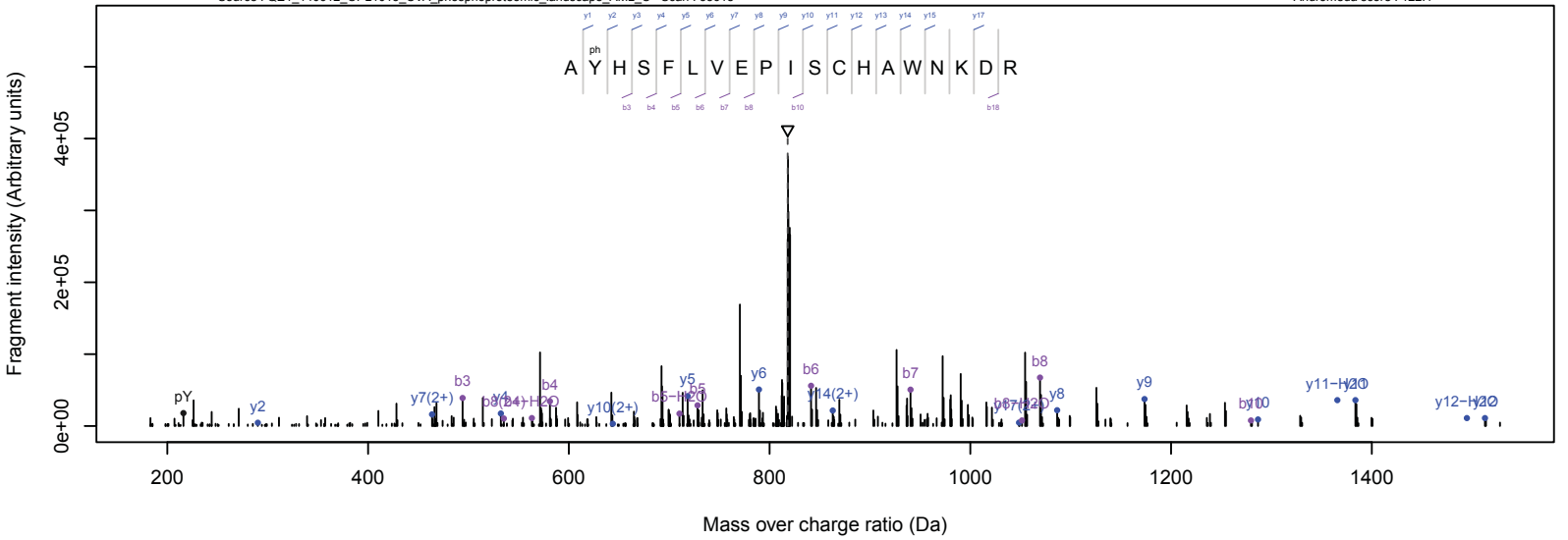
Andromeda score : 122.18



(ac)AY(ph)HSFLVEPISCHAWNKDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35618

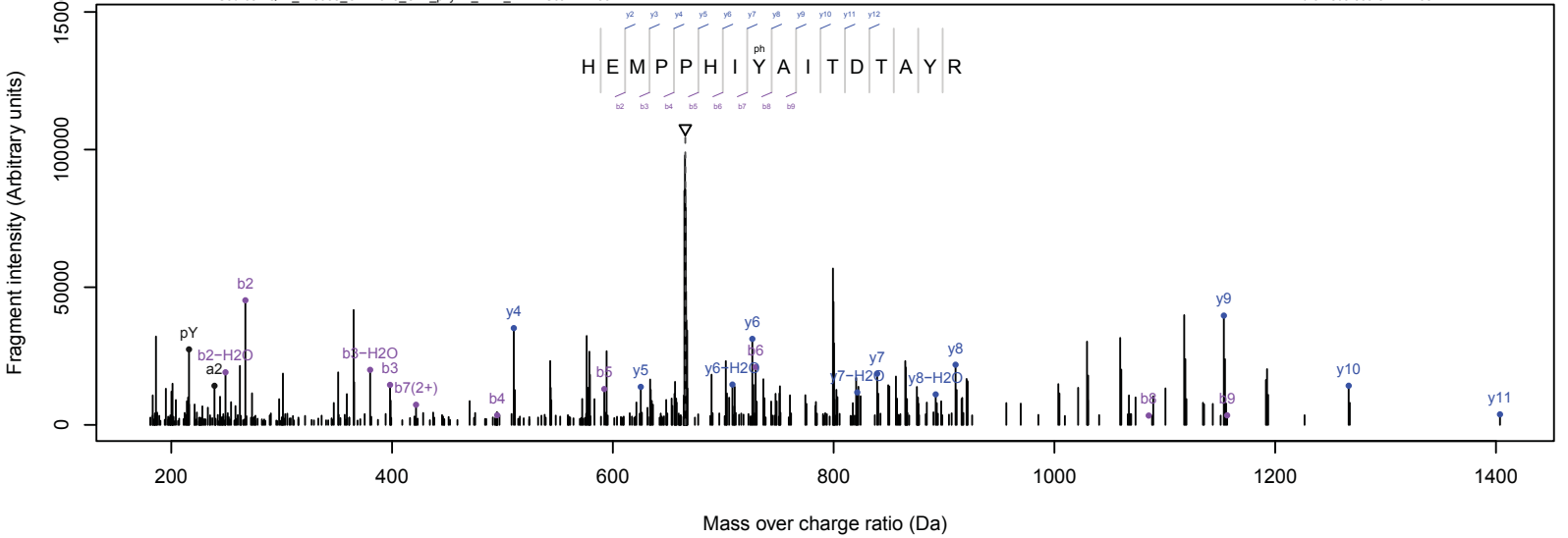
Andromeda score : 122.1



HEMPPHIYAITDTAYR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 27587

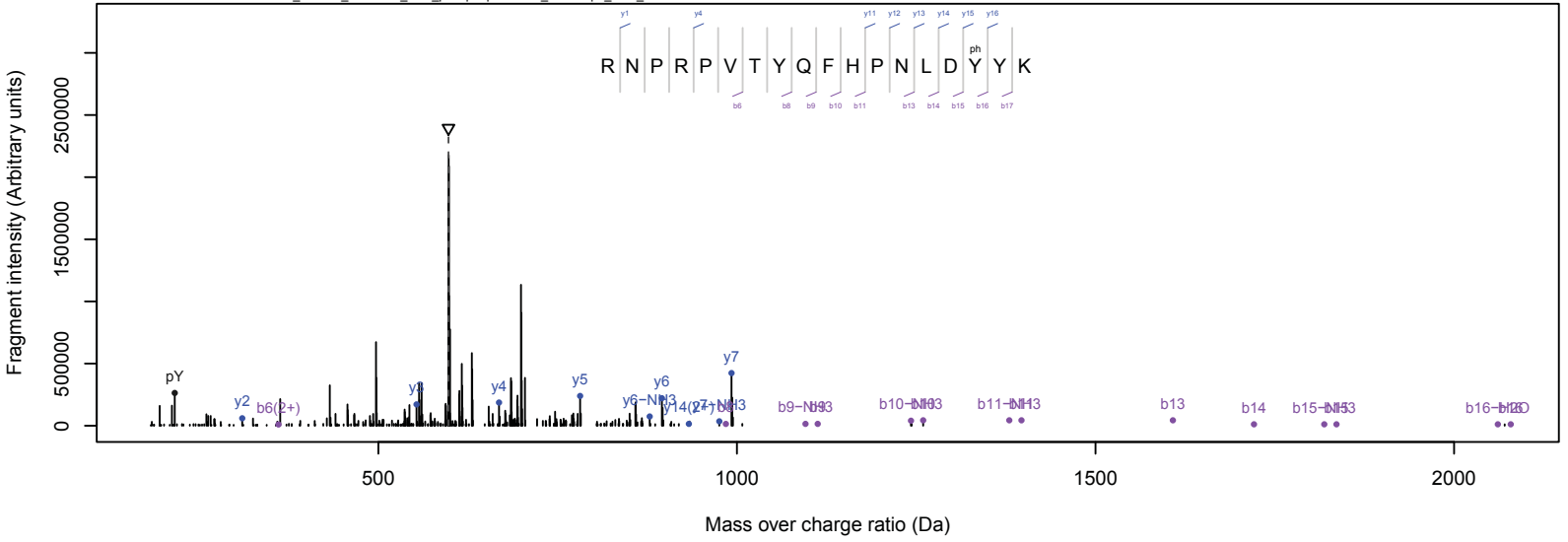
Andromeda score : 122.08



RNRPVTYQFHPNLDYYK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24761

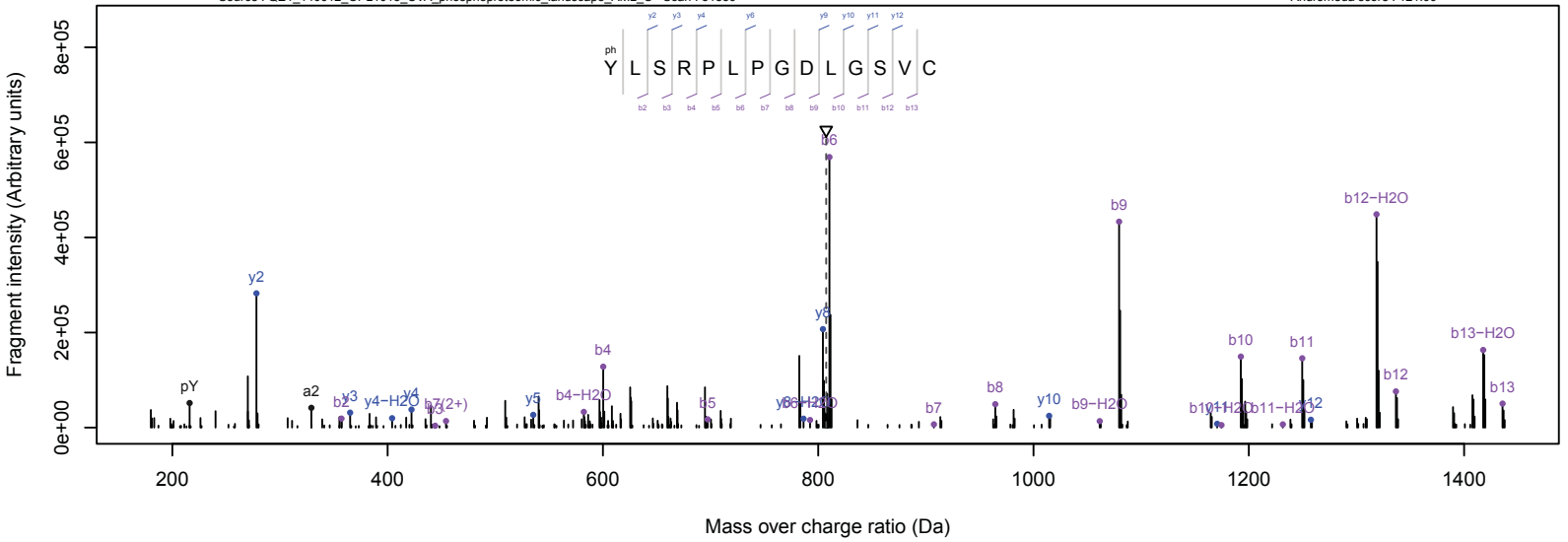
Andromeda score : 122.01



Y(ph)LSRPLPGDLGVC-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31859

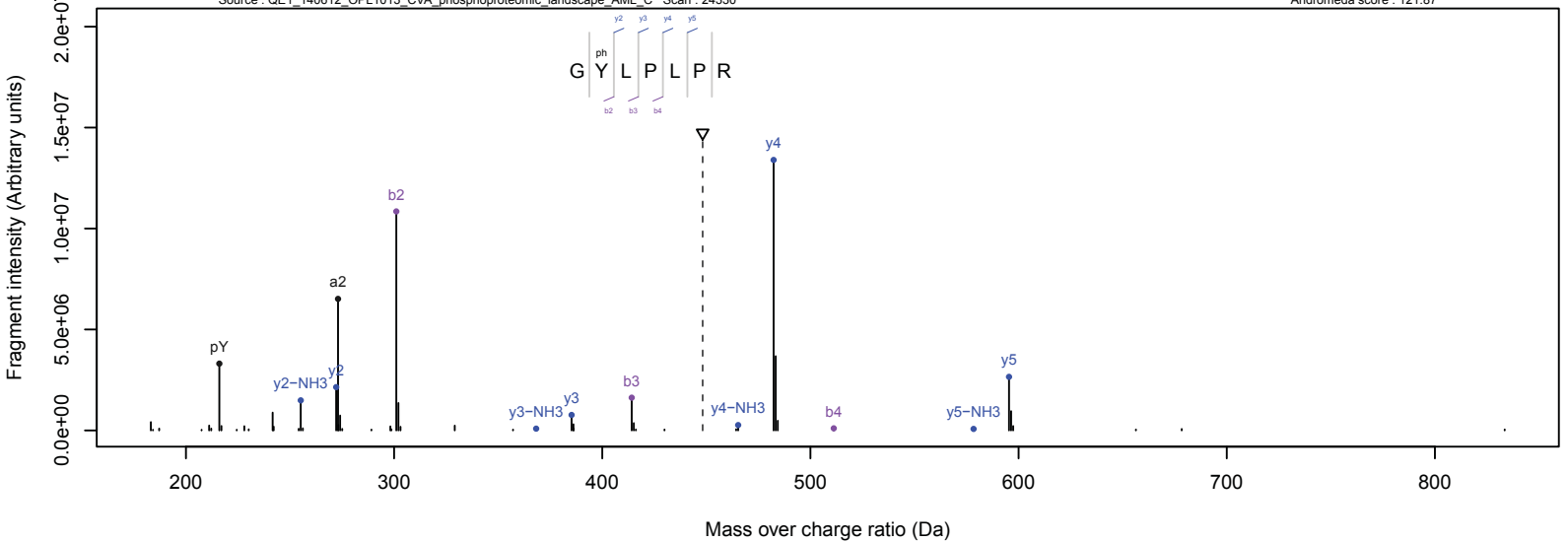
Andromeda score : 121.99



GY(ph)LPLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24330

Andromeda score : 121.87

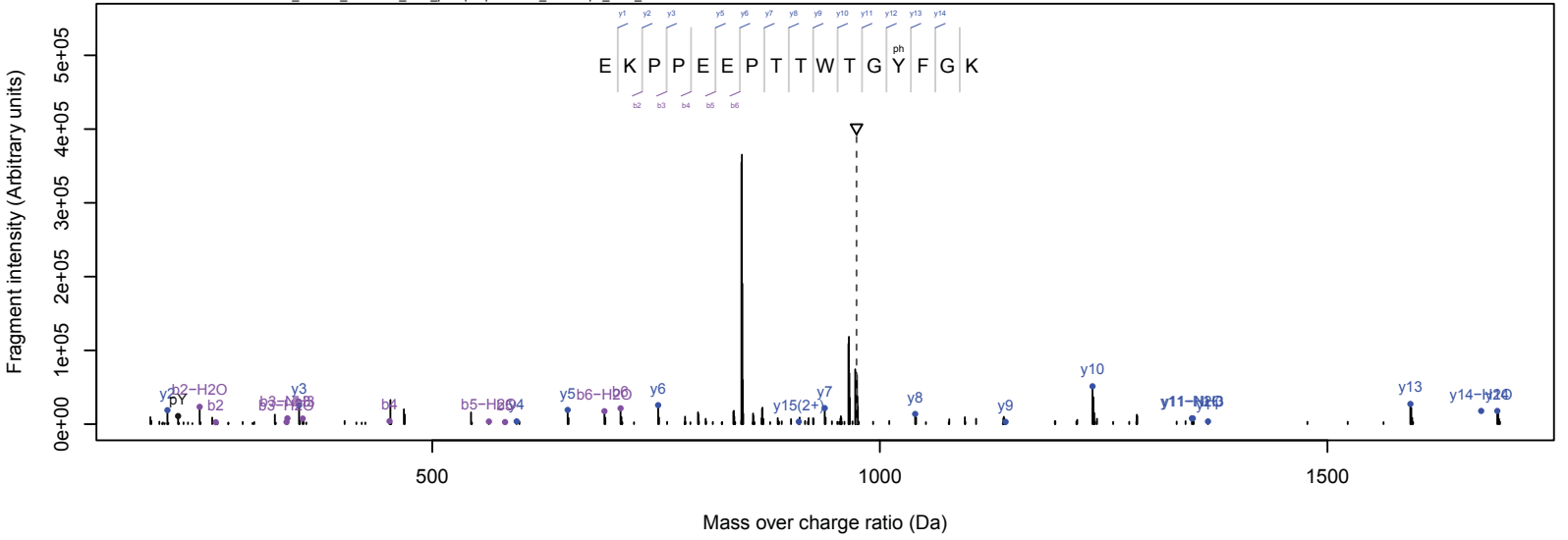




**EKPPEPTWTGY(ph)FGK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 28262

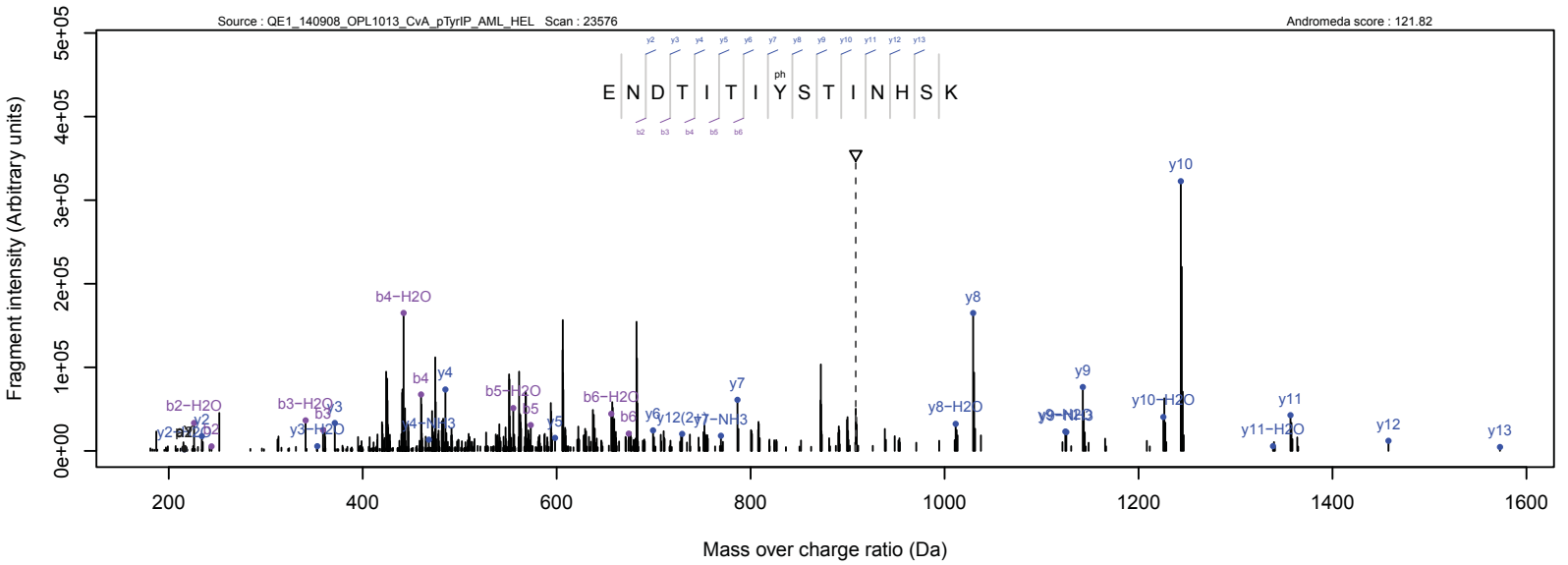
Andromeda score: 121.82



**ENDTITY(ph)STINHSK-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 23576

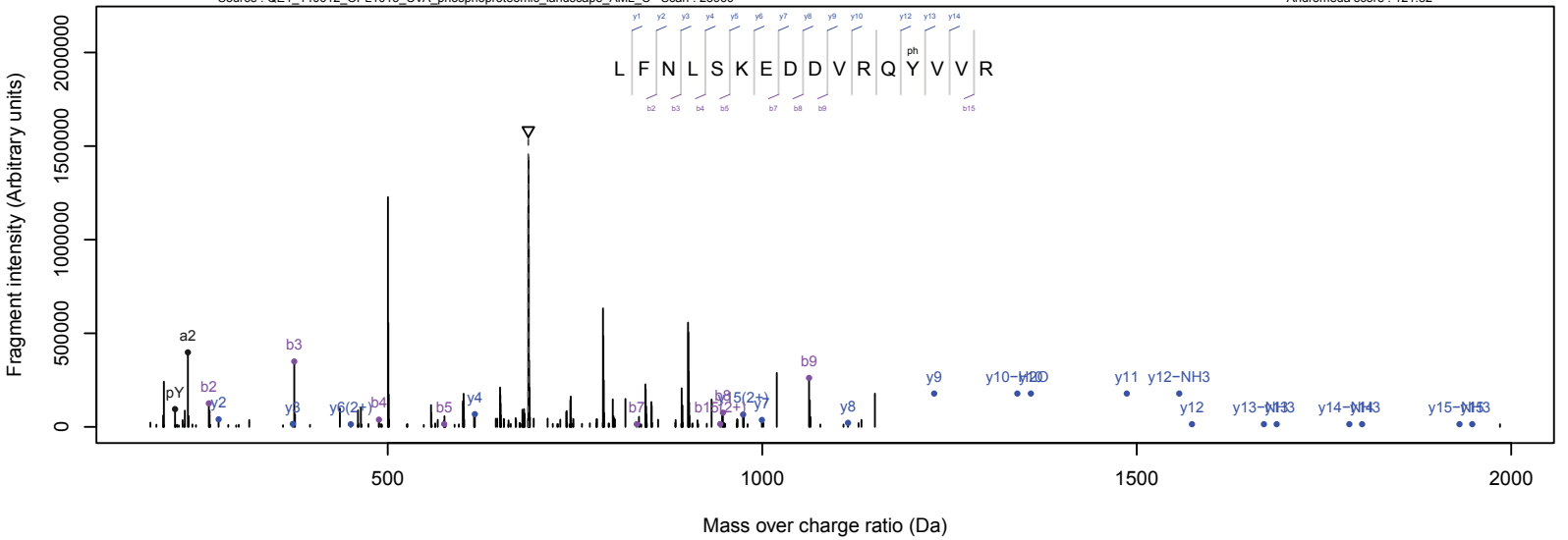
Andromeda score: 121.82



**LFNLSKEDDVRQY(ph)VVR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 28969

Andromeda score: 121.82

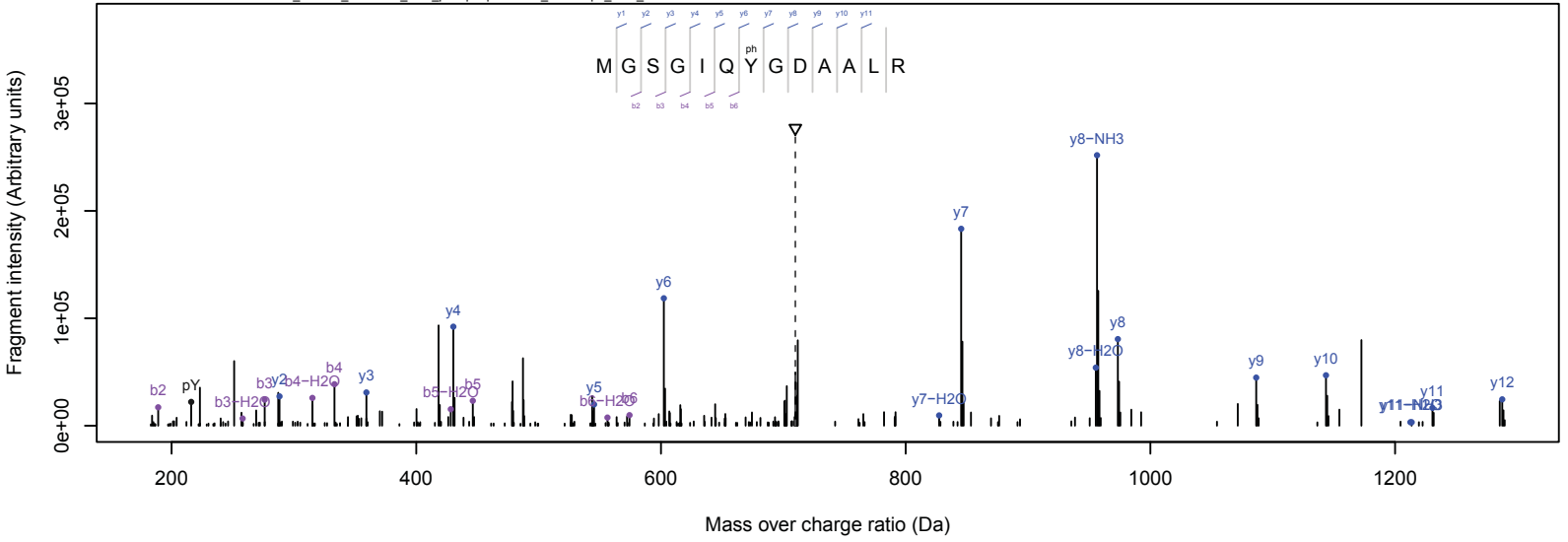




**MGSGIQ(ph)GDAALR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19505

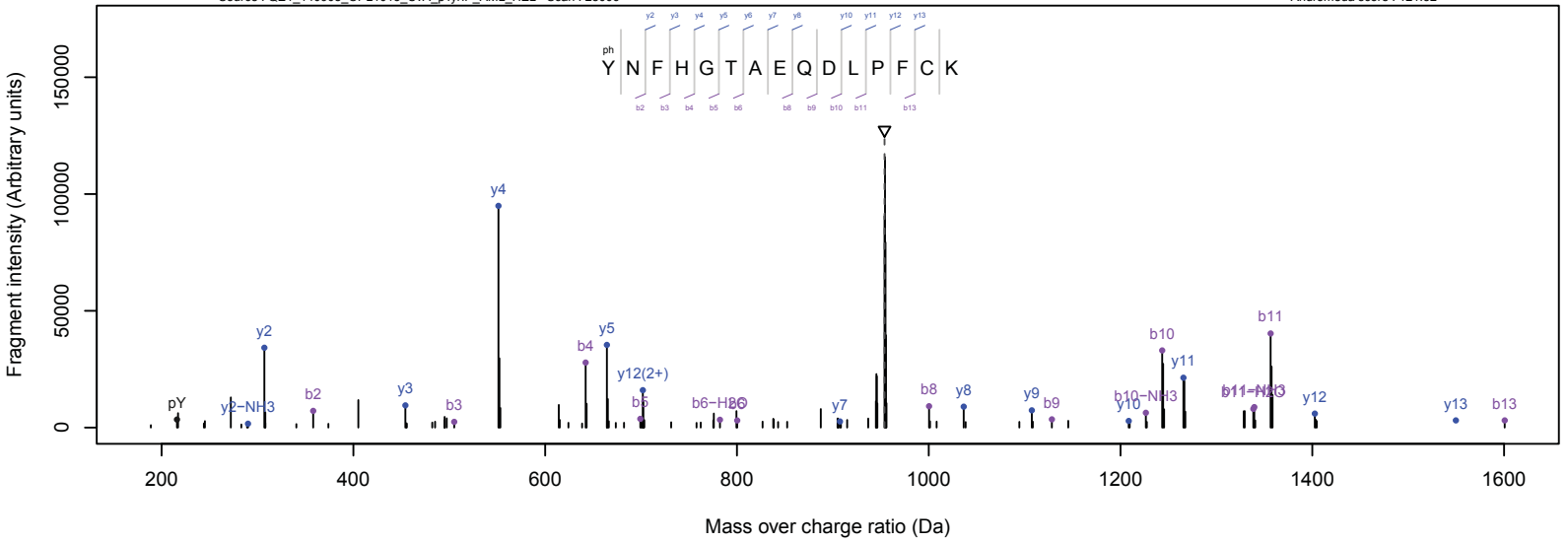
Andromeda score: 121.56



**Y(ph)NFHGTAEQDLPFCK-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 28006

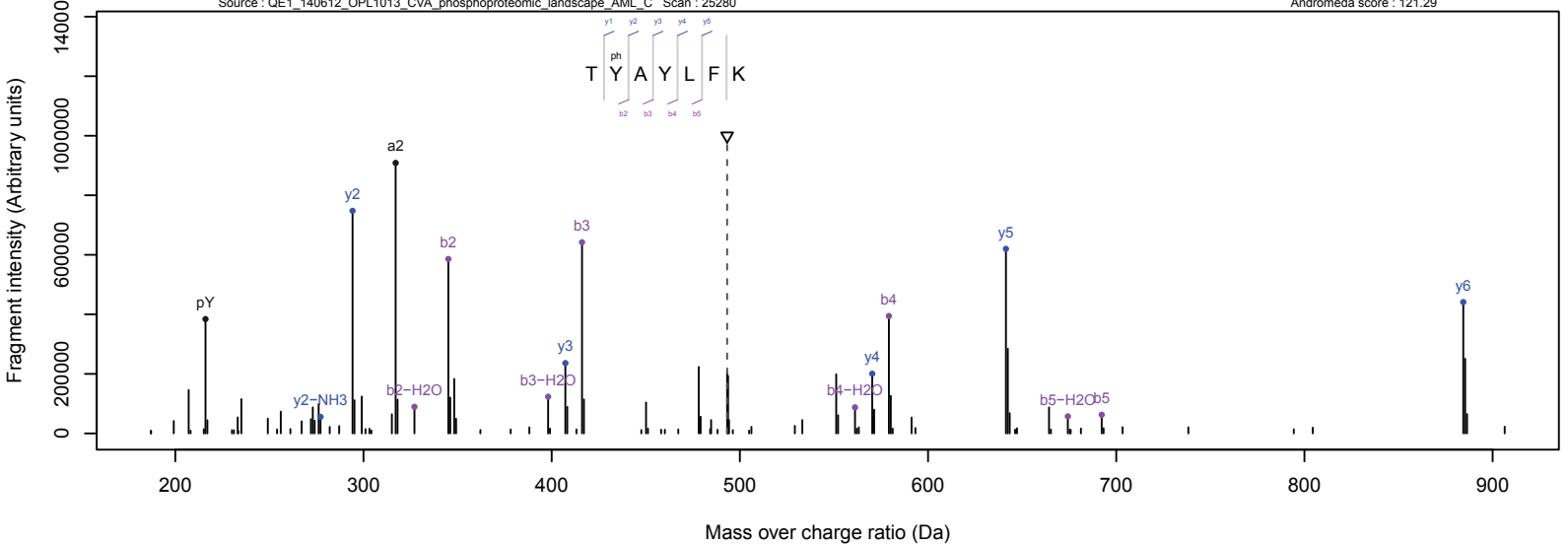
Andromeda score: 121.32



**TY(ph)AYLFK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25280

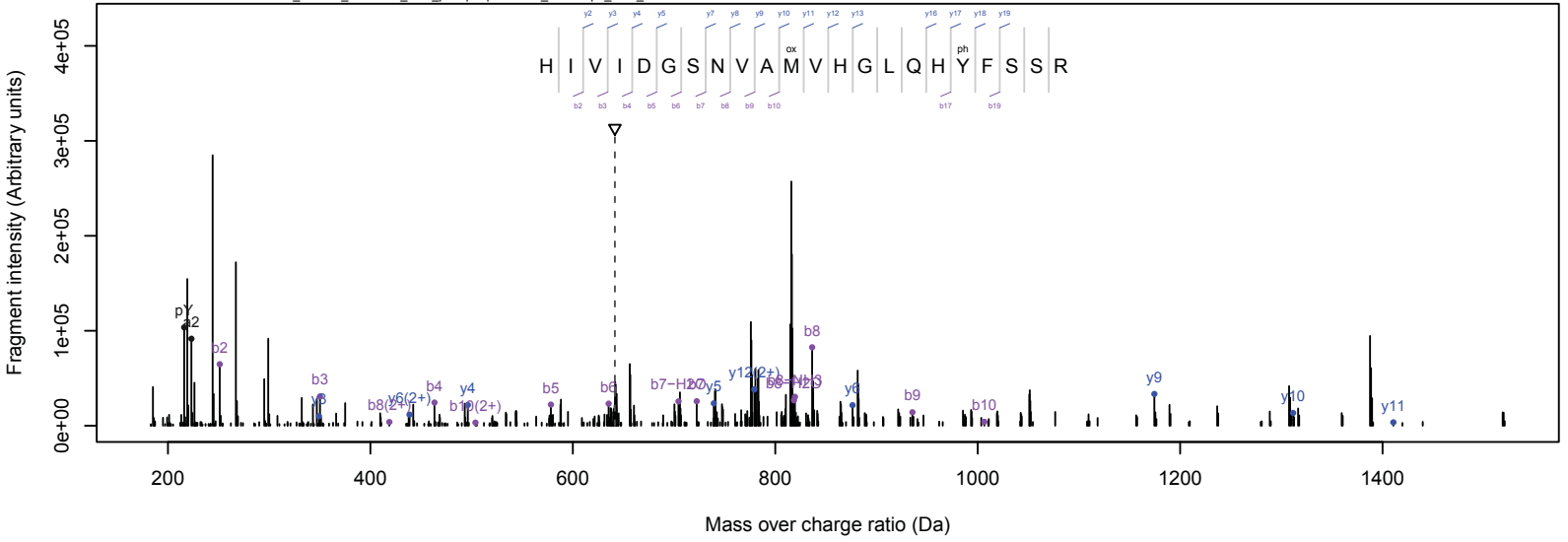
Andromeda score: 121.29



HIVIDGSNVAMVHGLQHY(ph)FSSR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31733

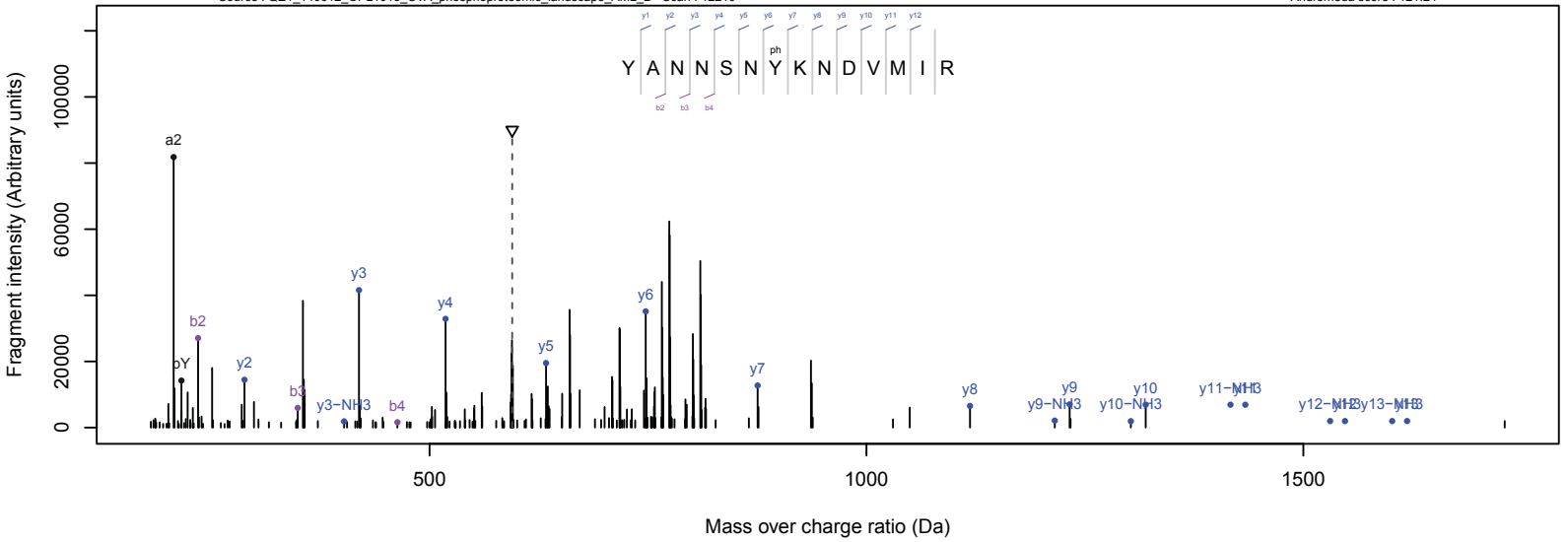
Andromeda score : 121.25



YANNSNY(ph)KNDVMIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12210

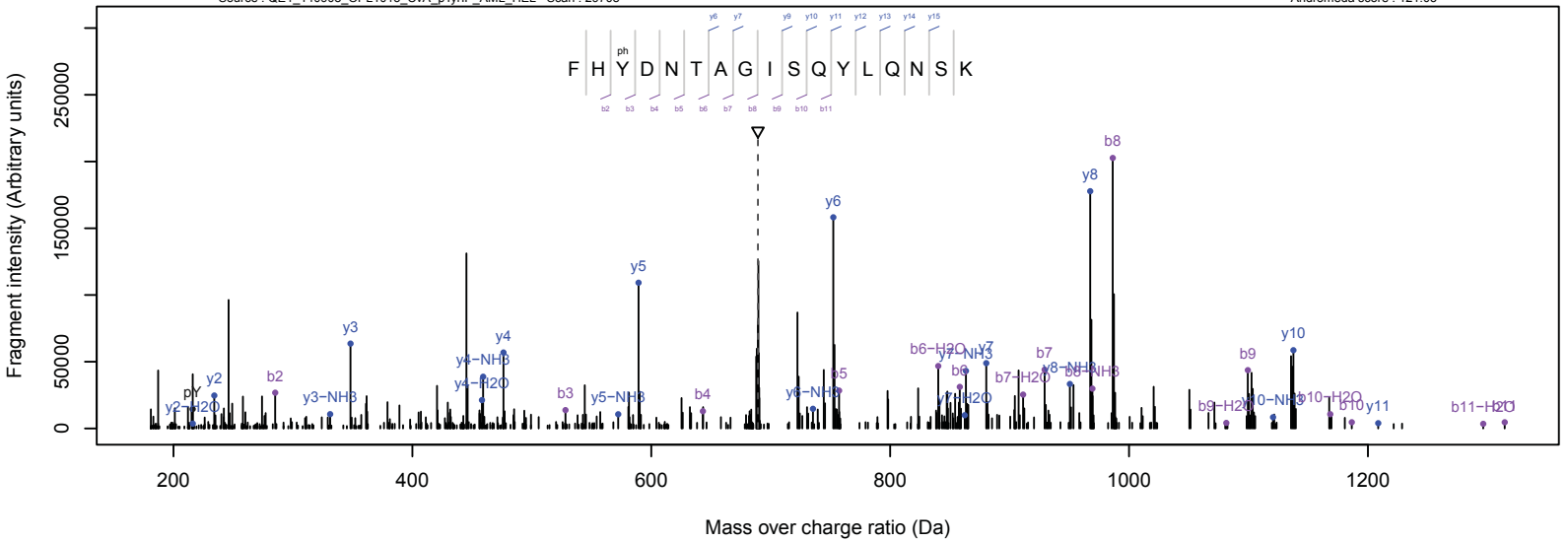
Andromeda score : 121.24



FHY(ph)DNTAGISQYLQNSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29708

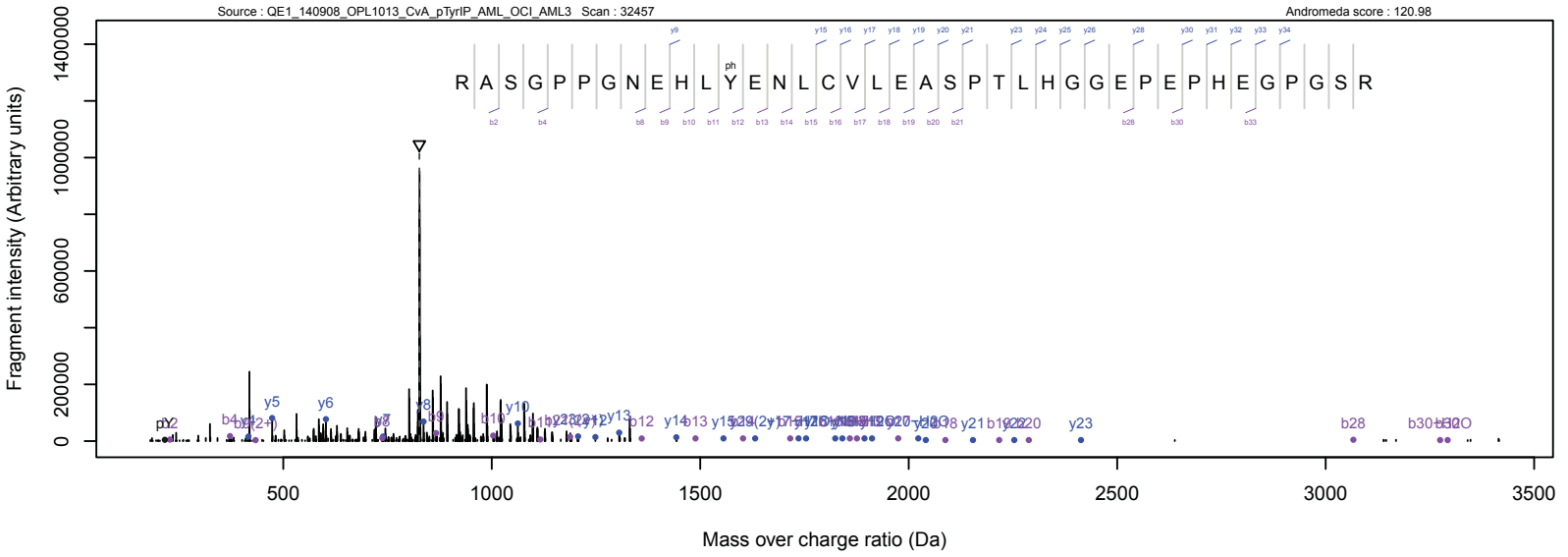
Andromeda score : 121.03



RASGPPGNEHLY(ph)ENLCVLEASPTLHGGEPEPEHGGPSR-5+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_OCI\_AML3 Scan: 32457

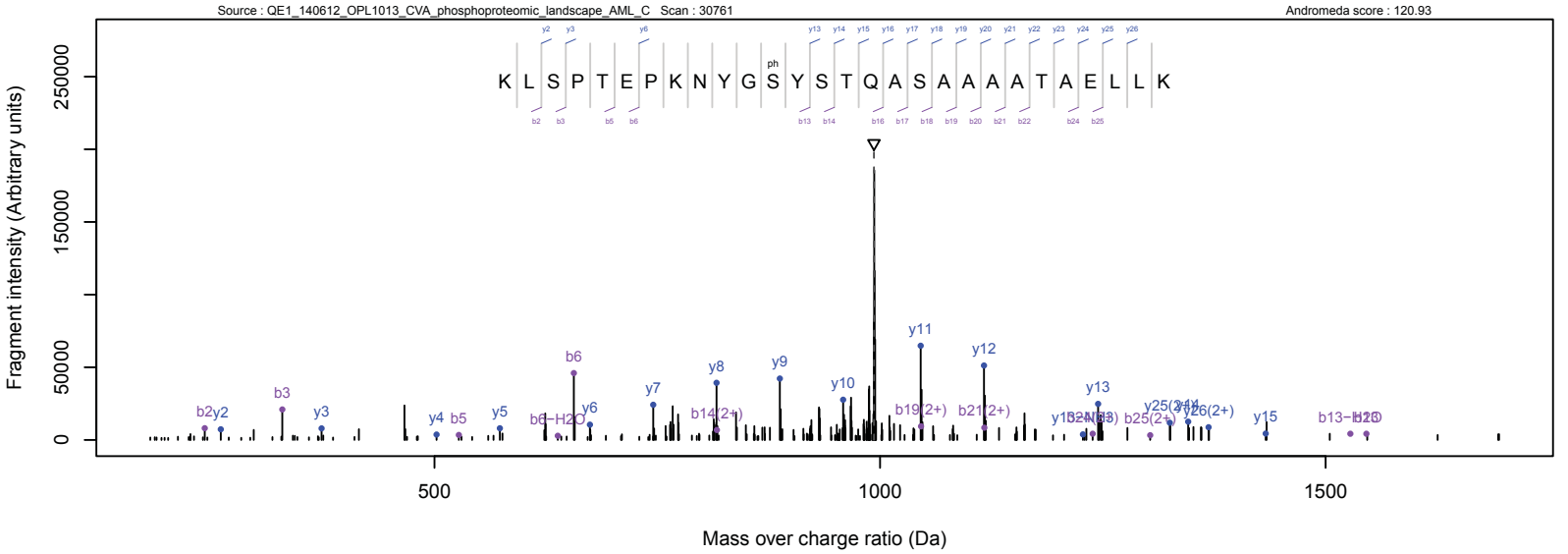
Andromeda score: 120.98



KLSPTPEKNYGS(ph)YSTQASAAAATAELLK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 30761

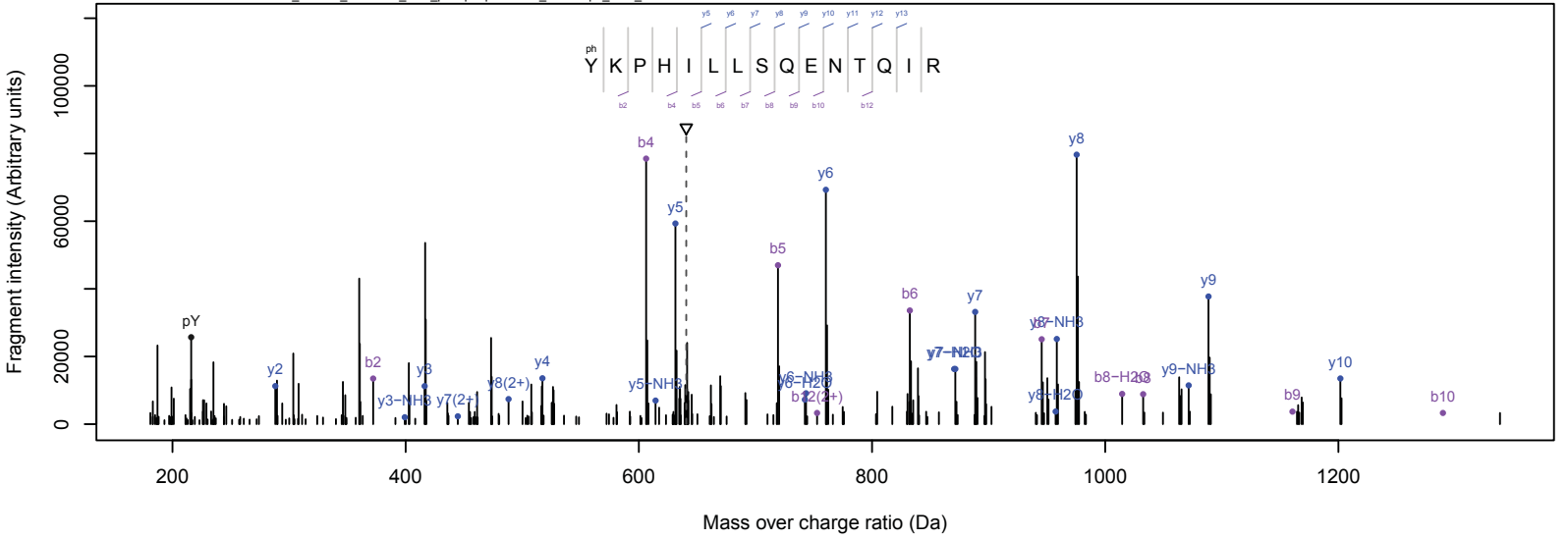
Andromeda score: 120.93



Y(ph)KPHILLSQENTQIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19228

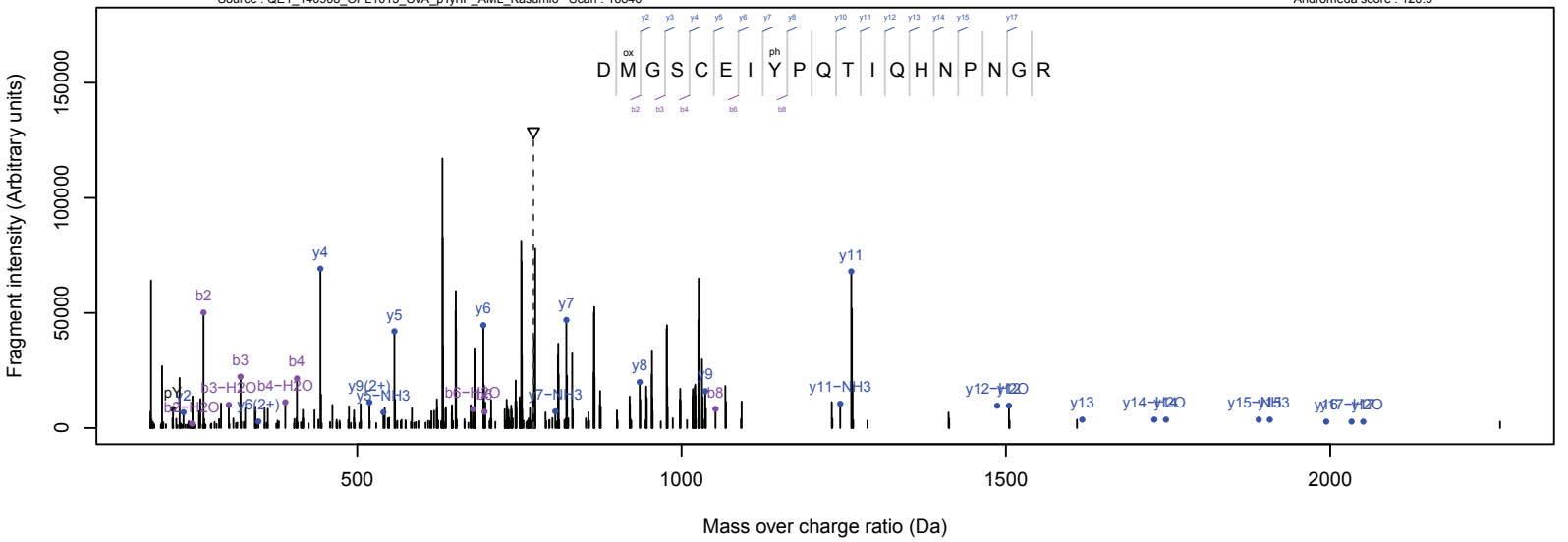
Andromeda score : 120.92



DM(ox)GSCEIY(ph)PQTIQHNPNGR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 16840

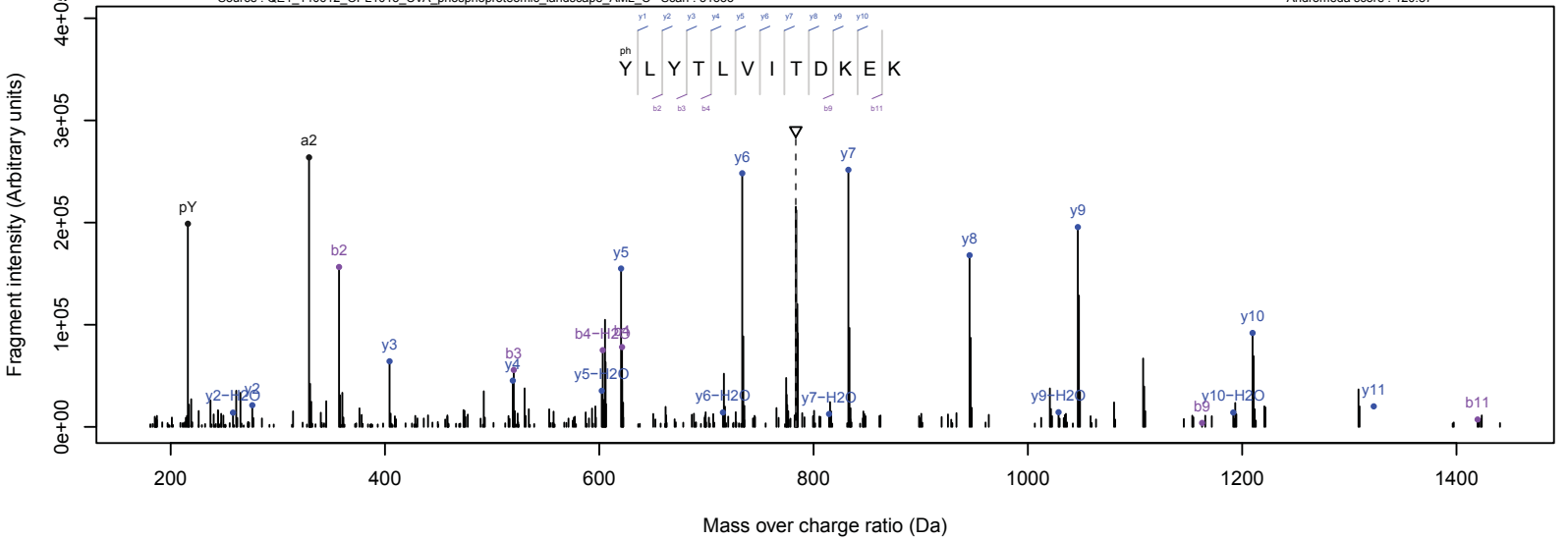
Andromeda score : 120.9



Y(ph)LYTLVITDKEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31553

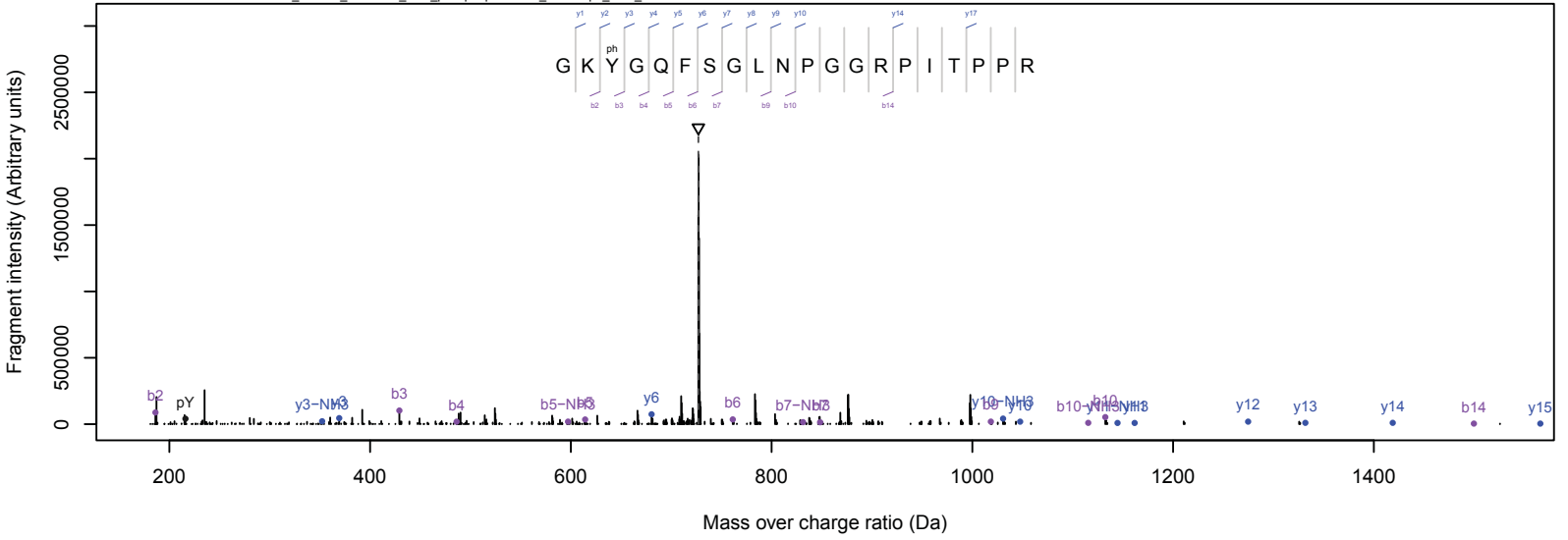
Andromeda score : 120.87



**GKY(ph)GQFSLNPGGRPITPPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22356

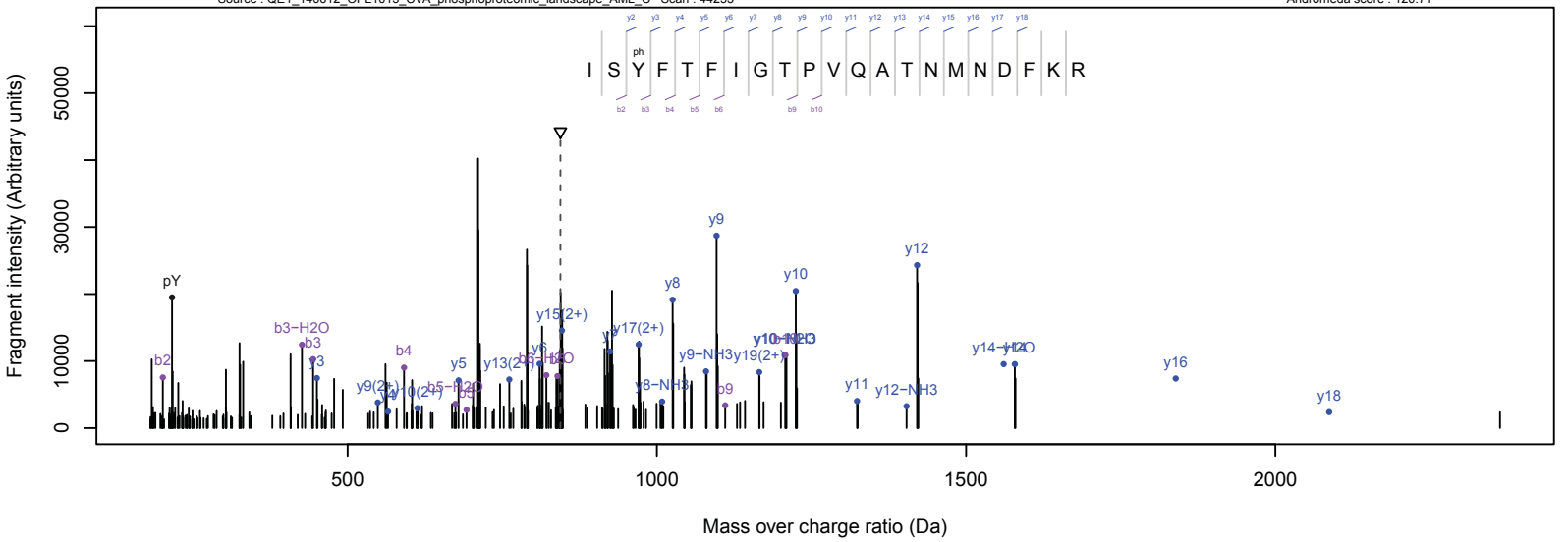
Andromeda score : 120.71



**ISY(ph)FTFIGTPVQATNMNDFKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44253

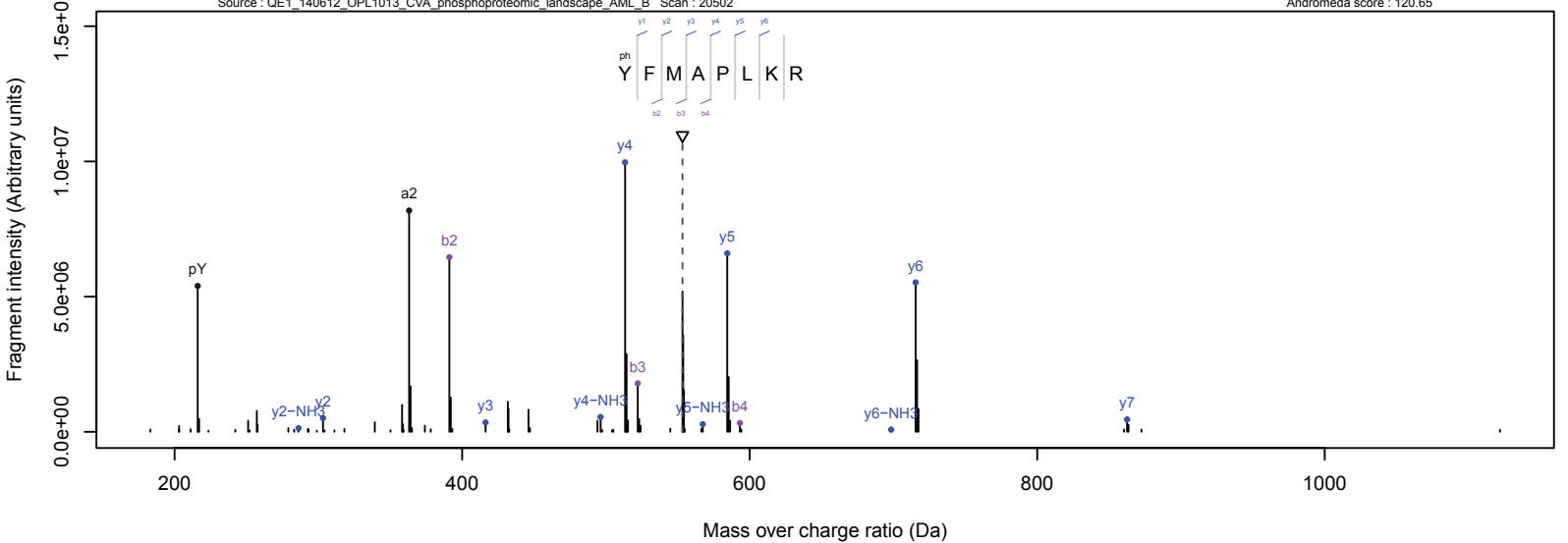
Andromeda score : 120.71



**Y(ph)FMAPLKR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20502

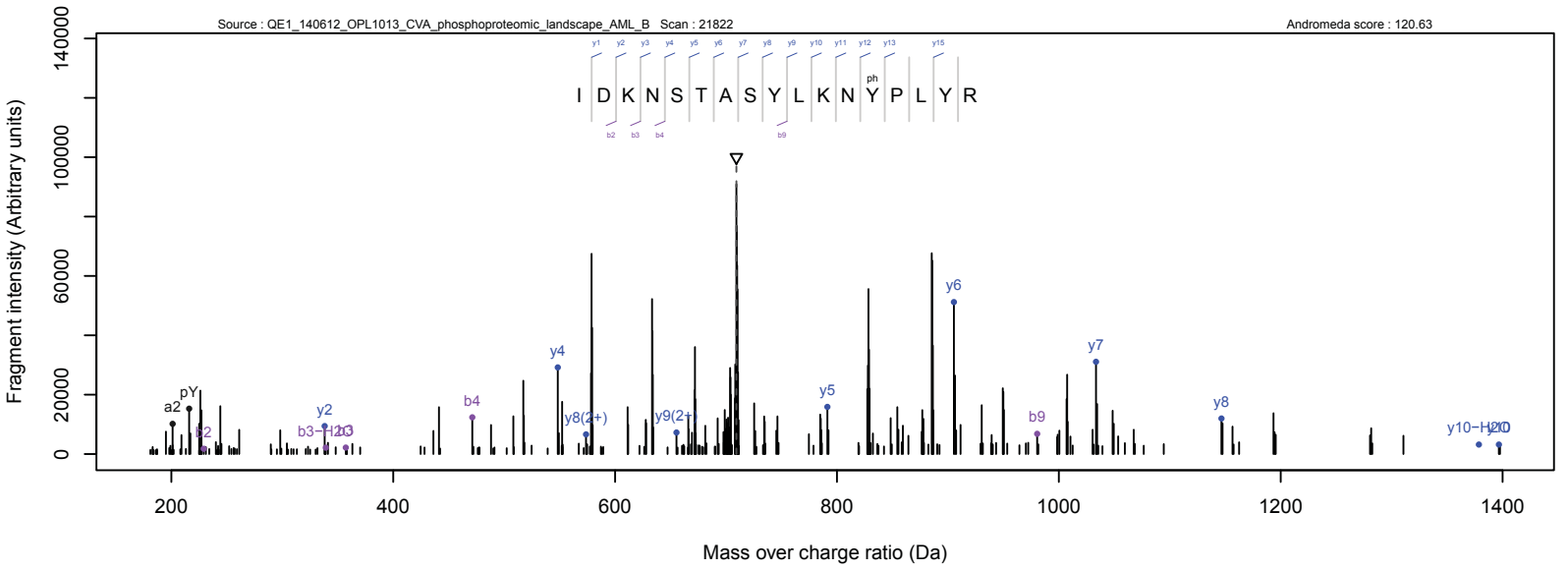
Andromeda score : 120.65



IDKNSTASYLKNY(ph)PLYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21822

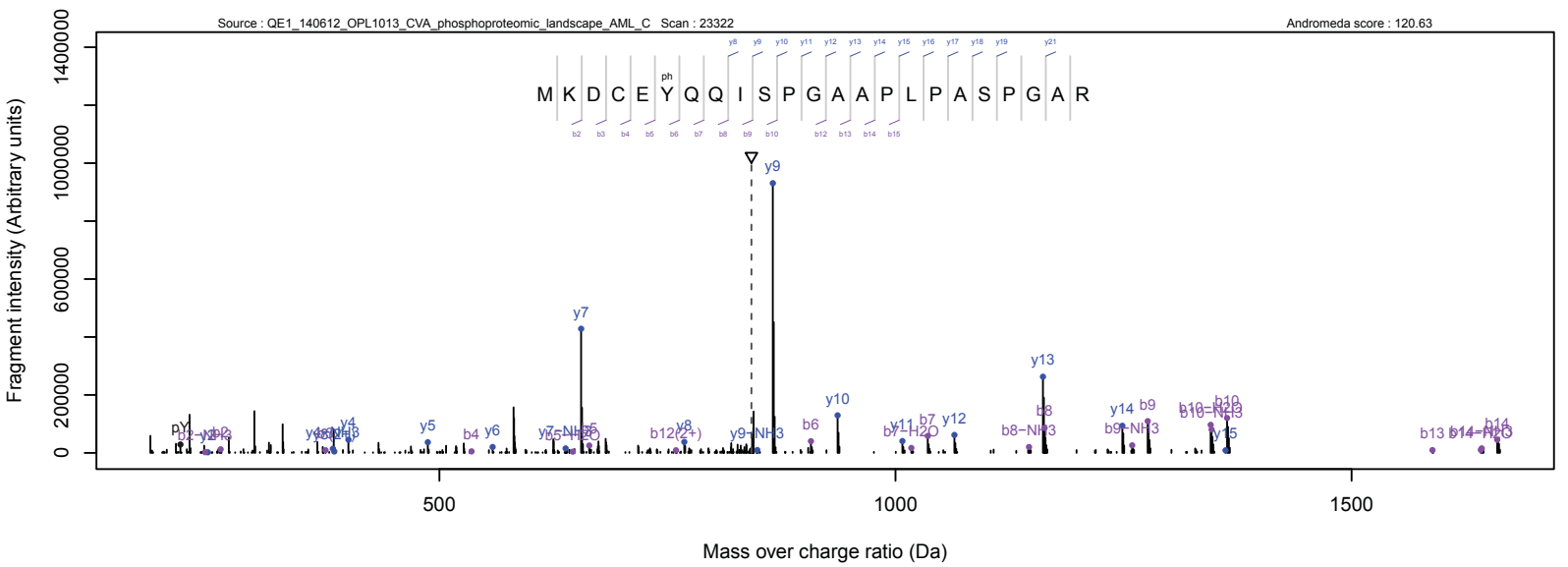
Andromeda score : 120.63



MKDCEY(ph)QQISPGAAPLPASPGAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23322

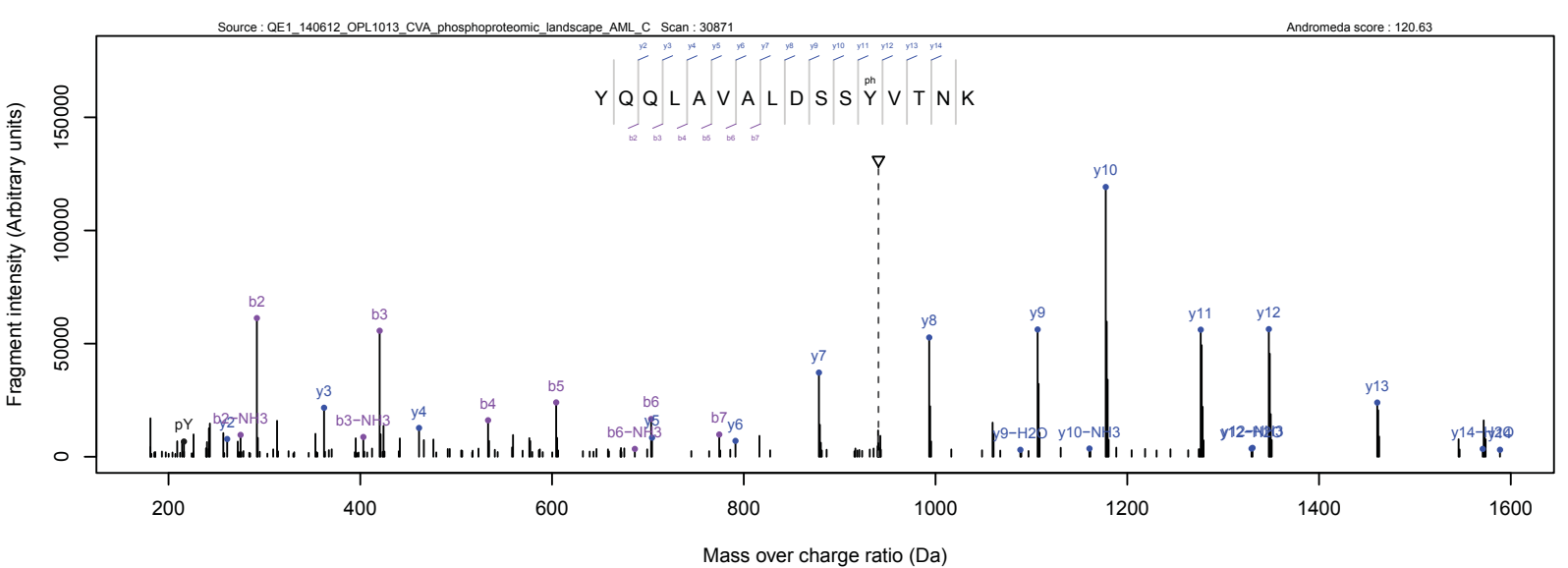
Andromeda score : 120.63



YQQLAVALDSSY(ph)VTNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30871

Andromeda score : 120.63

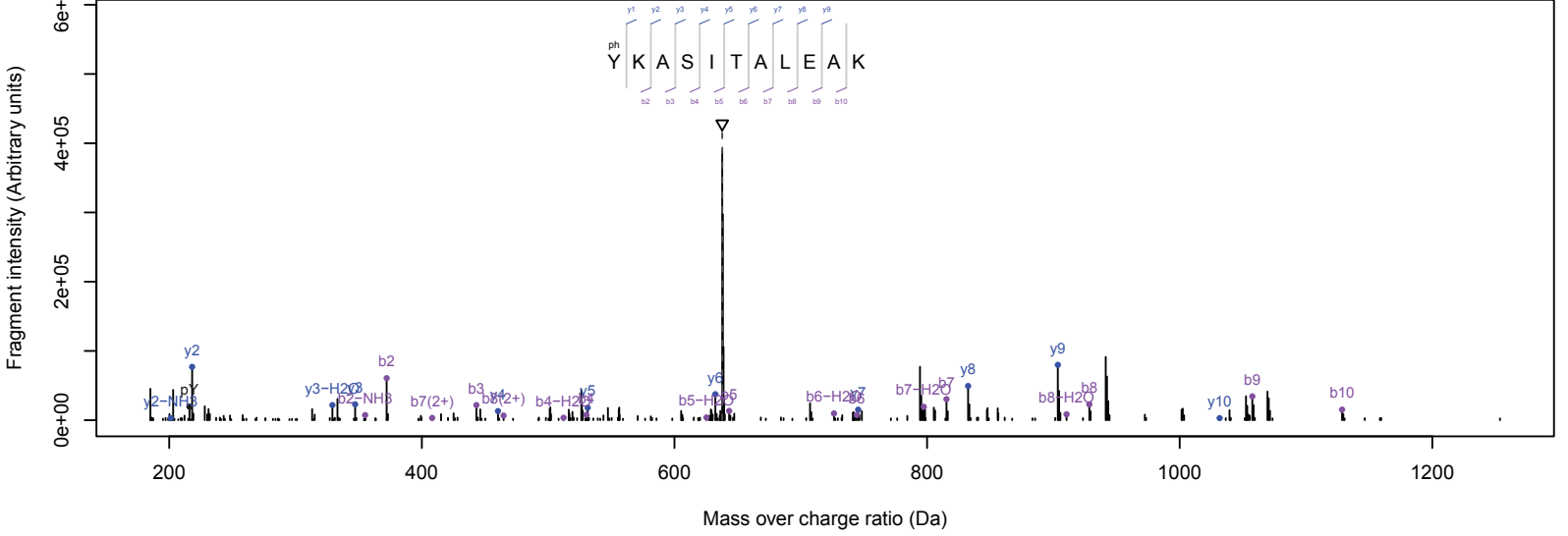




Y(ph)KASITALEAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 13873

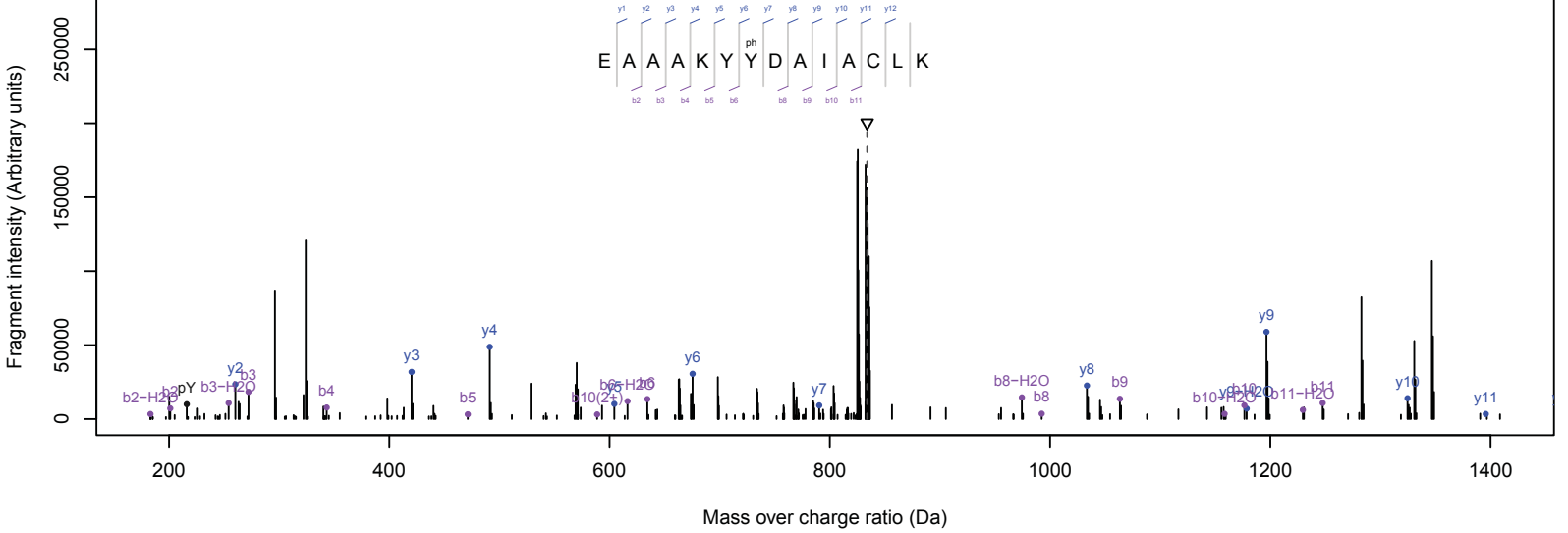
Andromeda score : 120.6



EAAAKYY(ph)DAIACK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26558

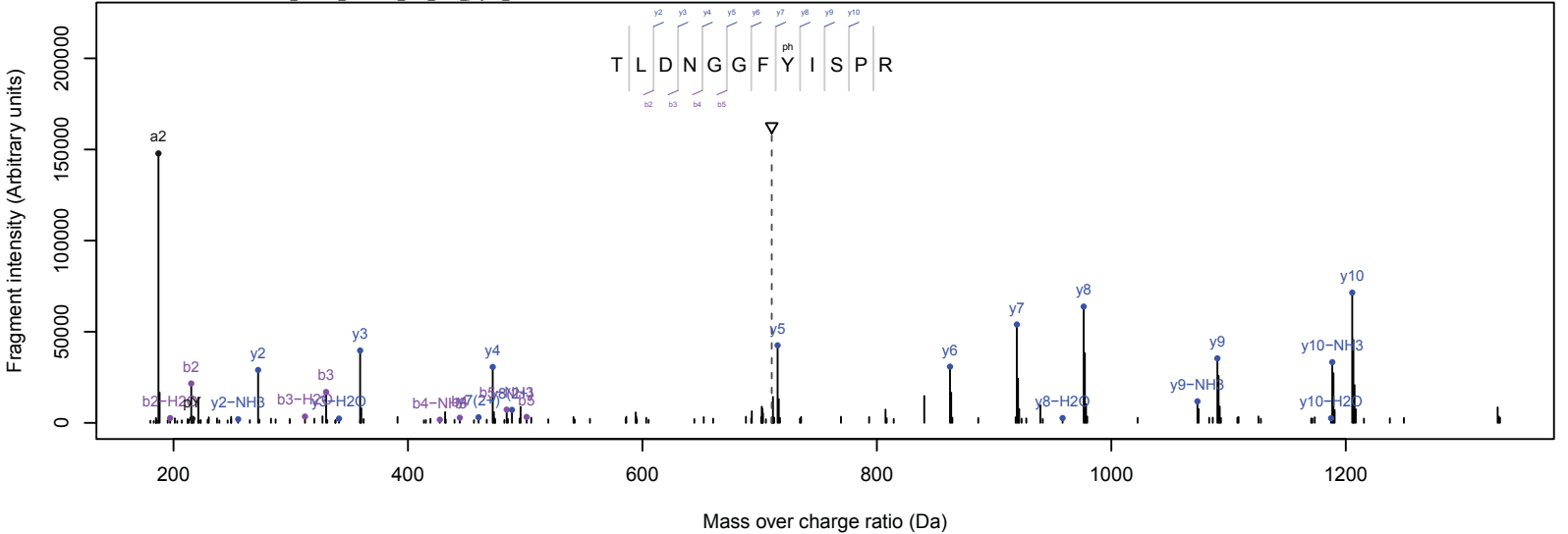
Andromeda score : 120.54



TLDNGGFY(ph)ISPR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 29086

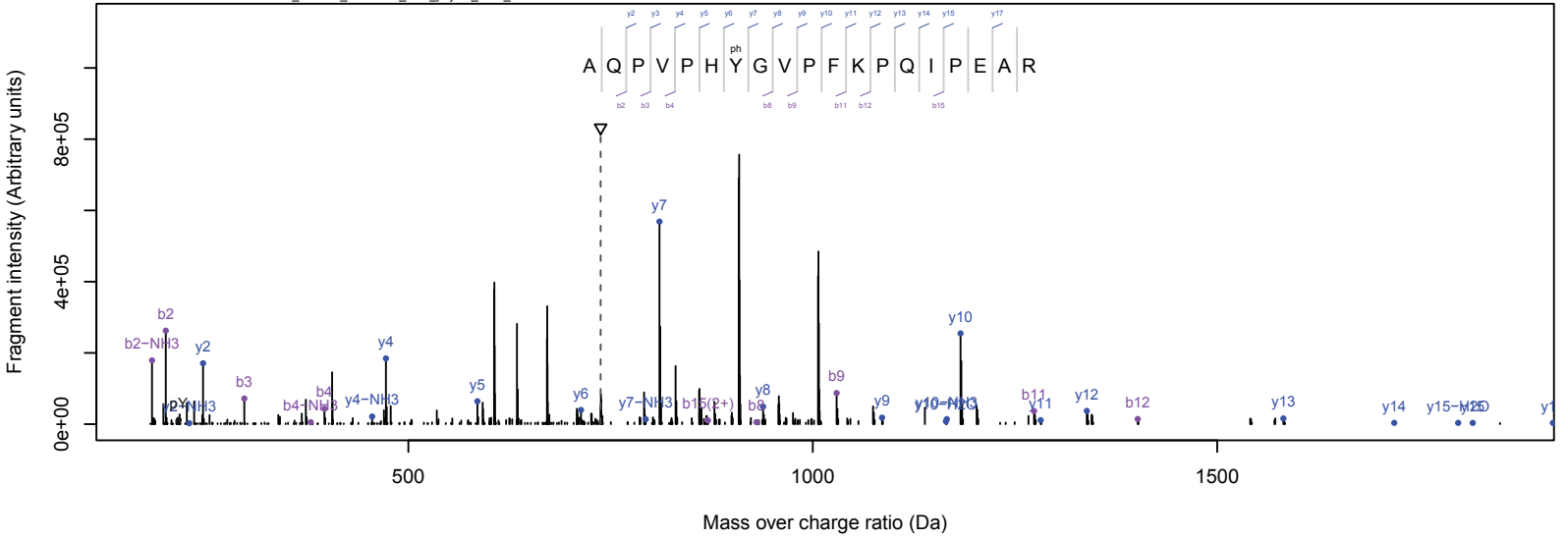
Andromeda score : 120.53



AQPVPHY(ph)GVFPFKPIPEAR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29648

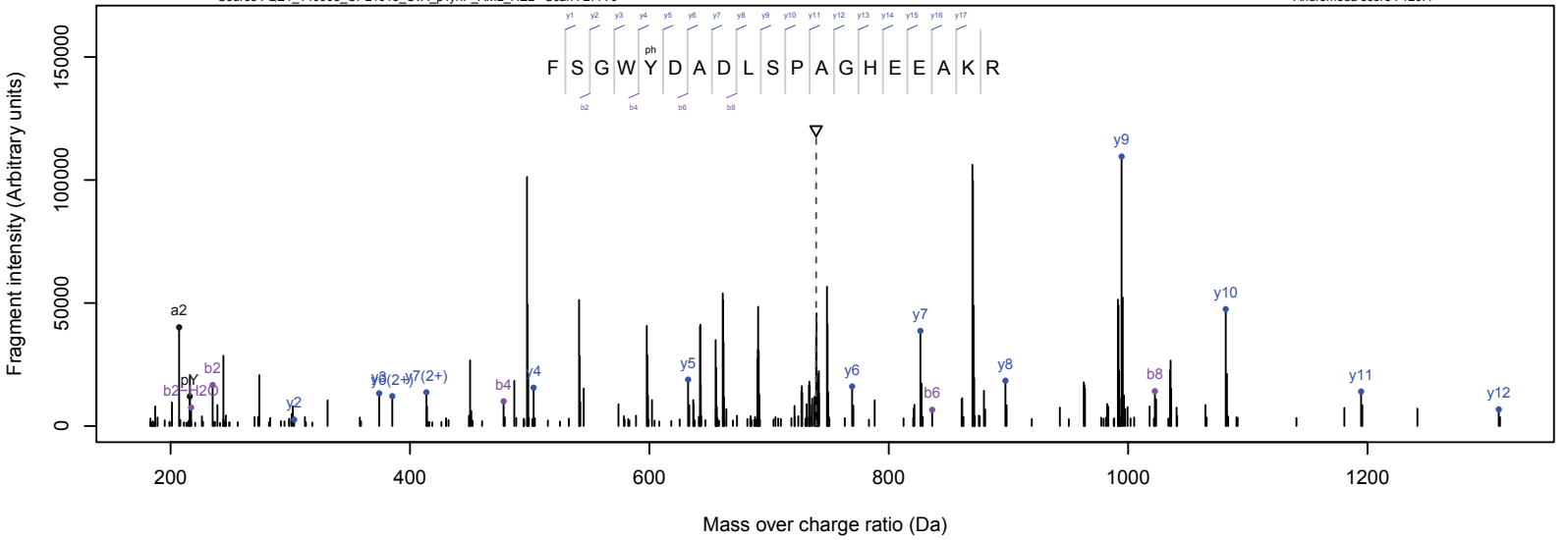
Andromeda score : 120.4



FSGWY(ph)DADLSPAGHEEAKR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 27770

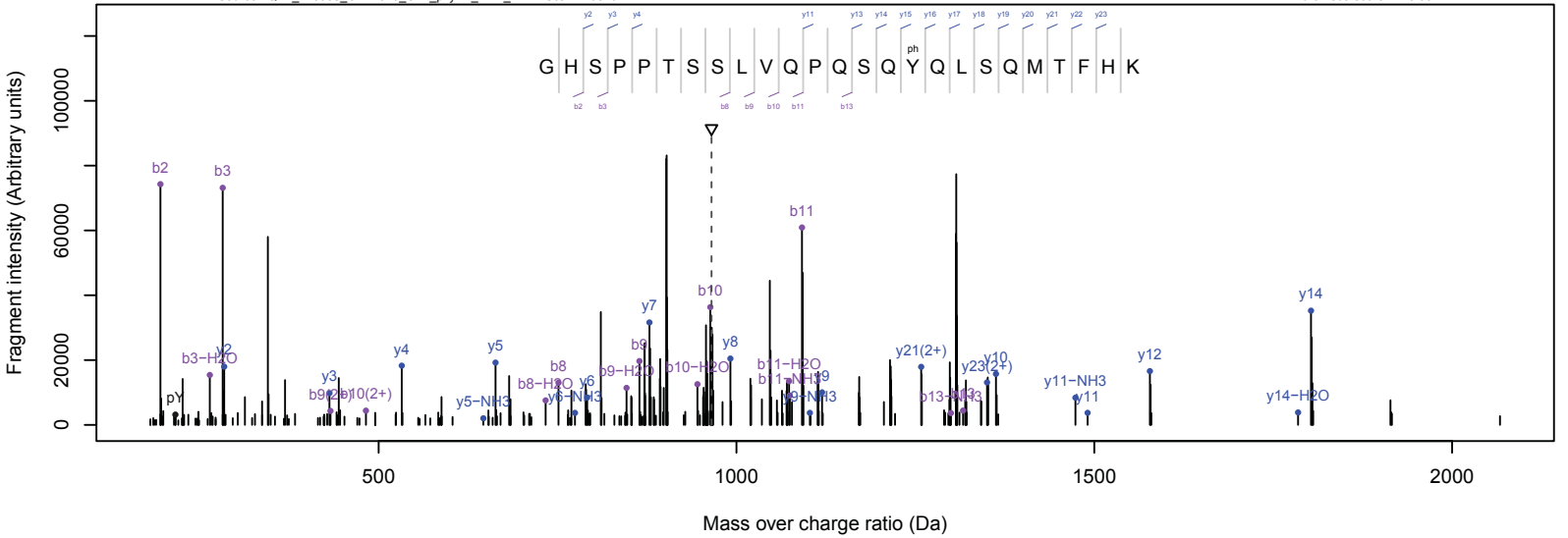
Andromeda score : 120.4



GHSPTSSLVQPQSQYQLSQMTFHK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29549

Andromeda score : 120.33

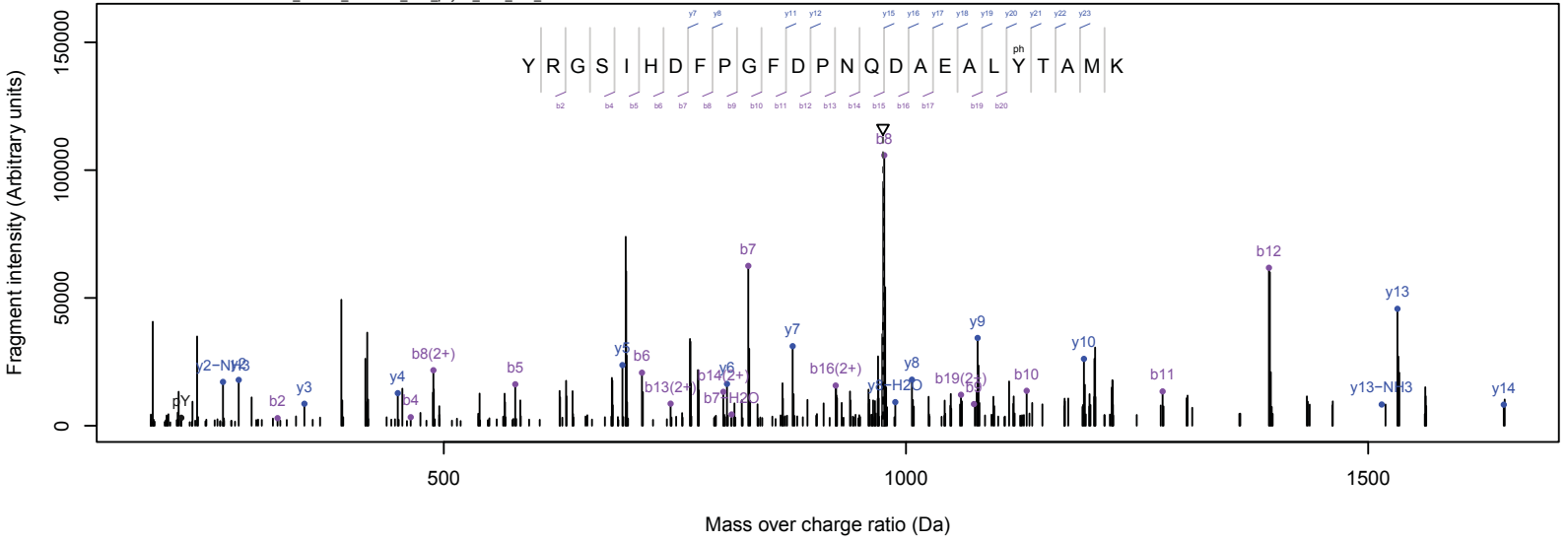




YRGSIHDFPFGDPNQDAEALY<sup>ph</sup>TAMK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 46411

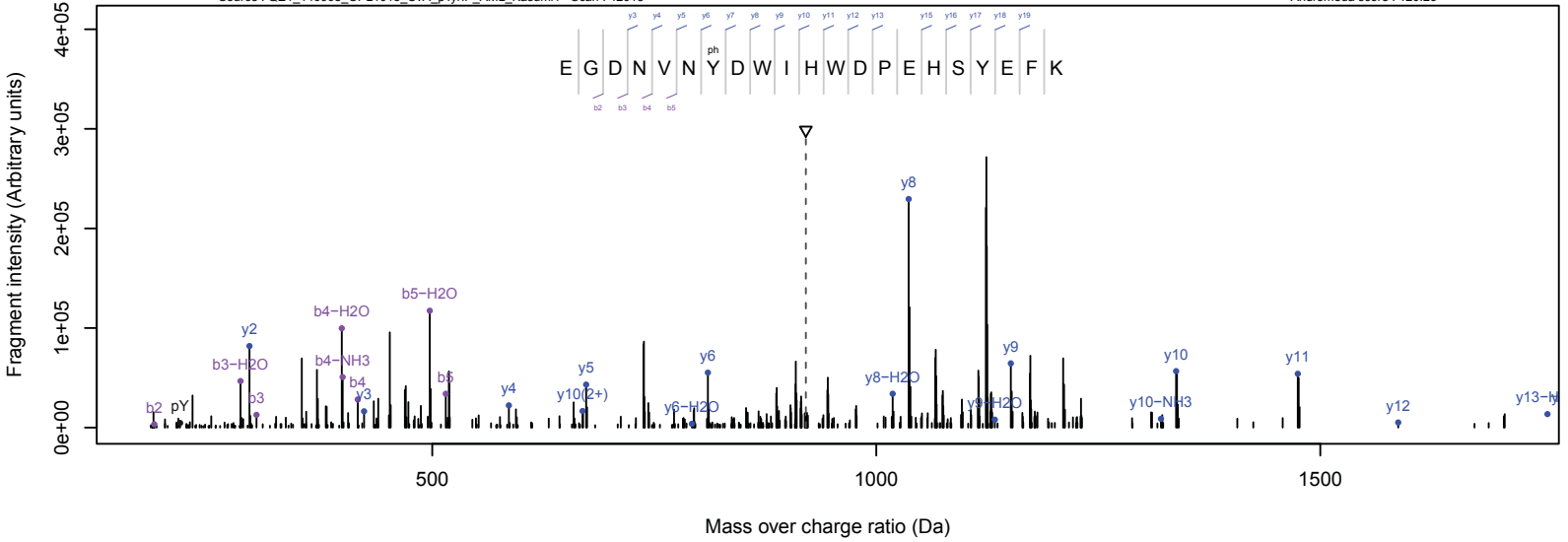
Andromeda score: 120.27



EGD<sup>ph</sup>NVNYD<sup>ph</sup>WIHWDPESHSEYEFK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 42018

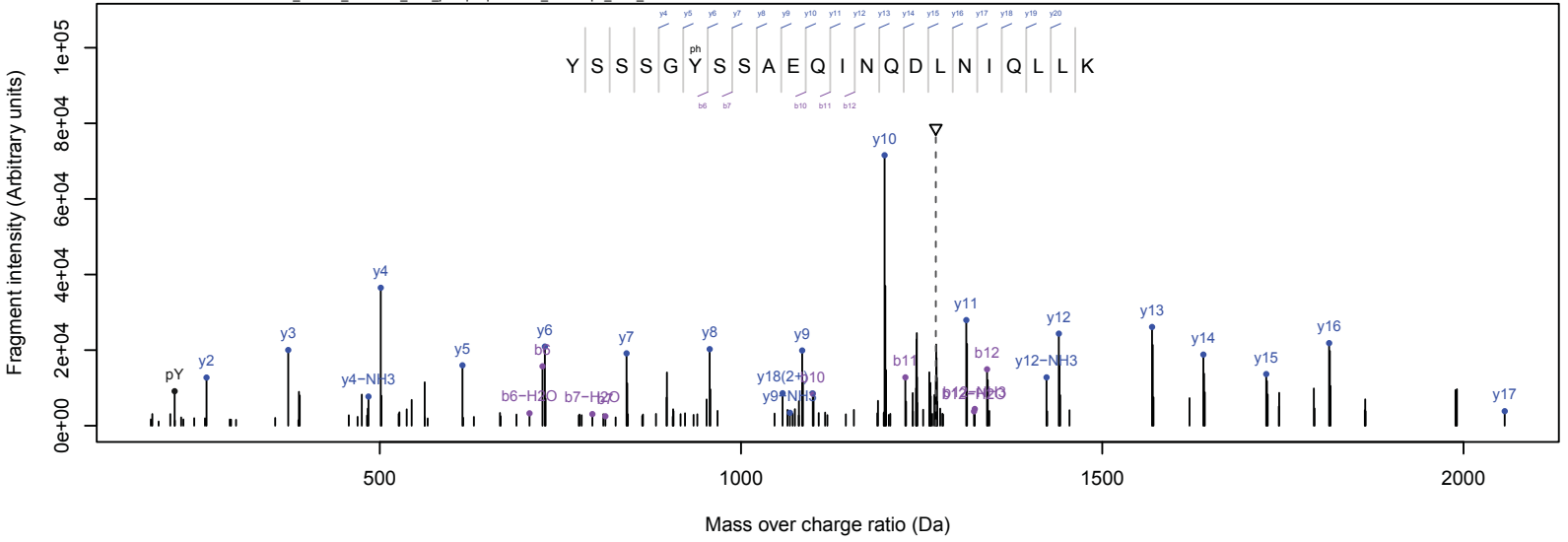
Andromeda score: 120.23



**YSSSGY(ph)SSAEQINQLNQLLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42232

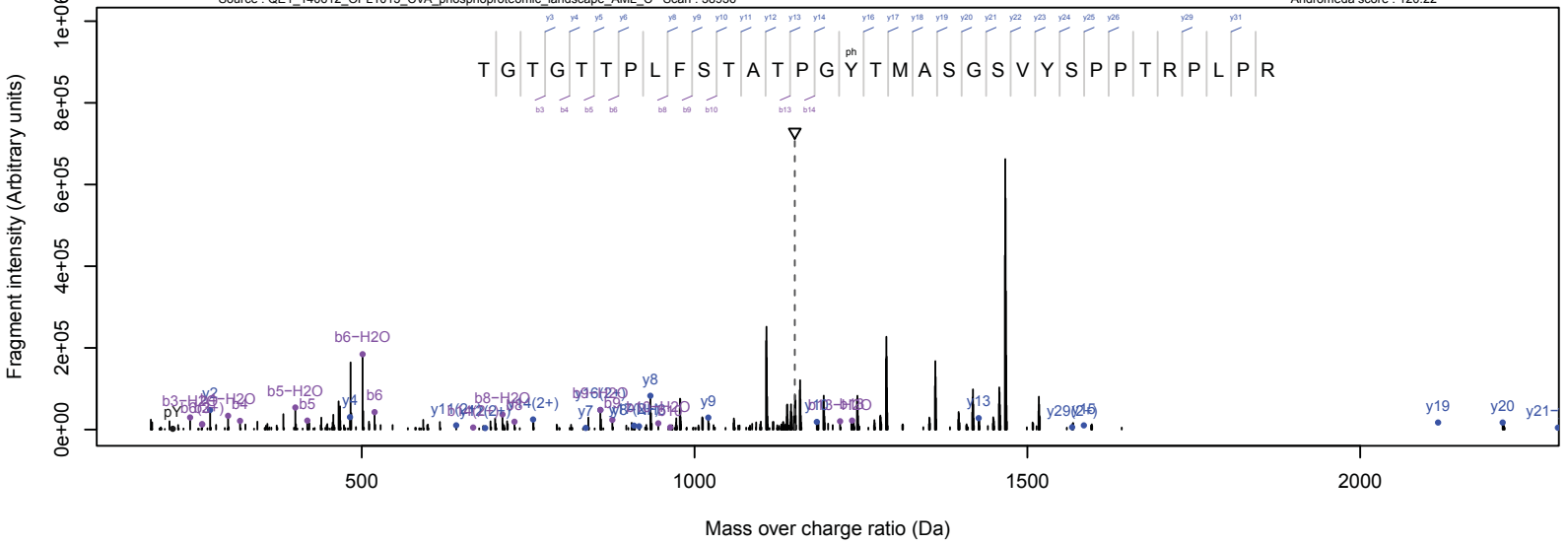
Andromeda score : 120.23



**TGTGTTPLFSTATPGY(ph)TMASGSVYSPPTRPLPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38936

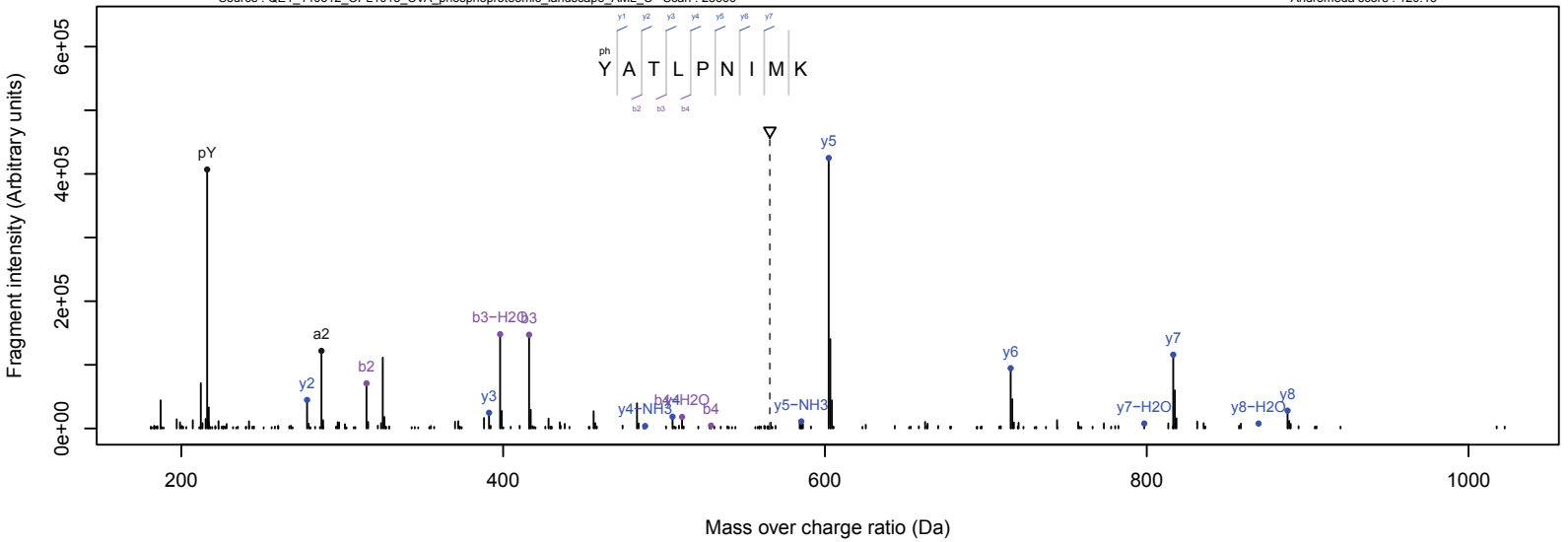
Andromeda score : 120.22



**Y(ph)ATLPNIMK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28666

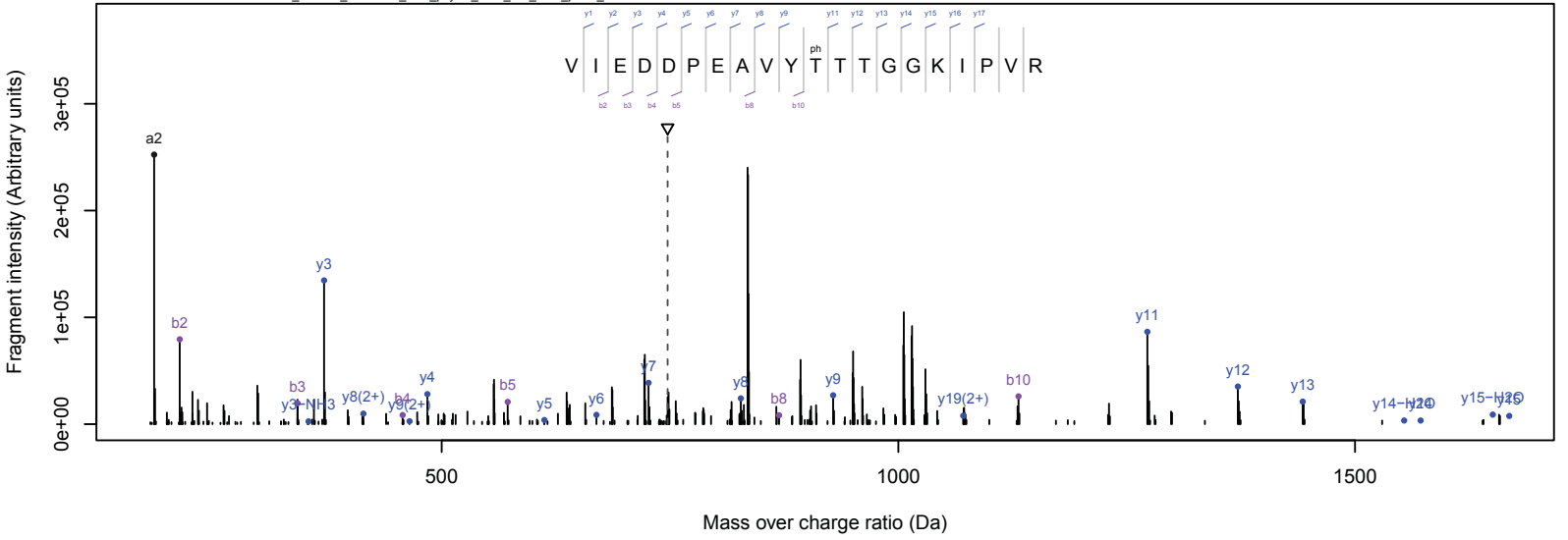
Andromeda score : 120.15



VIEDDPEAVY(ph)TTGGKIPVR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 21418

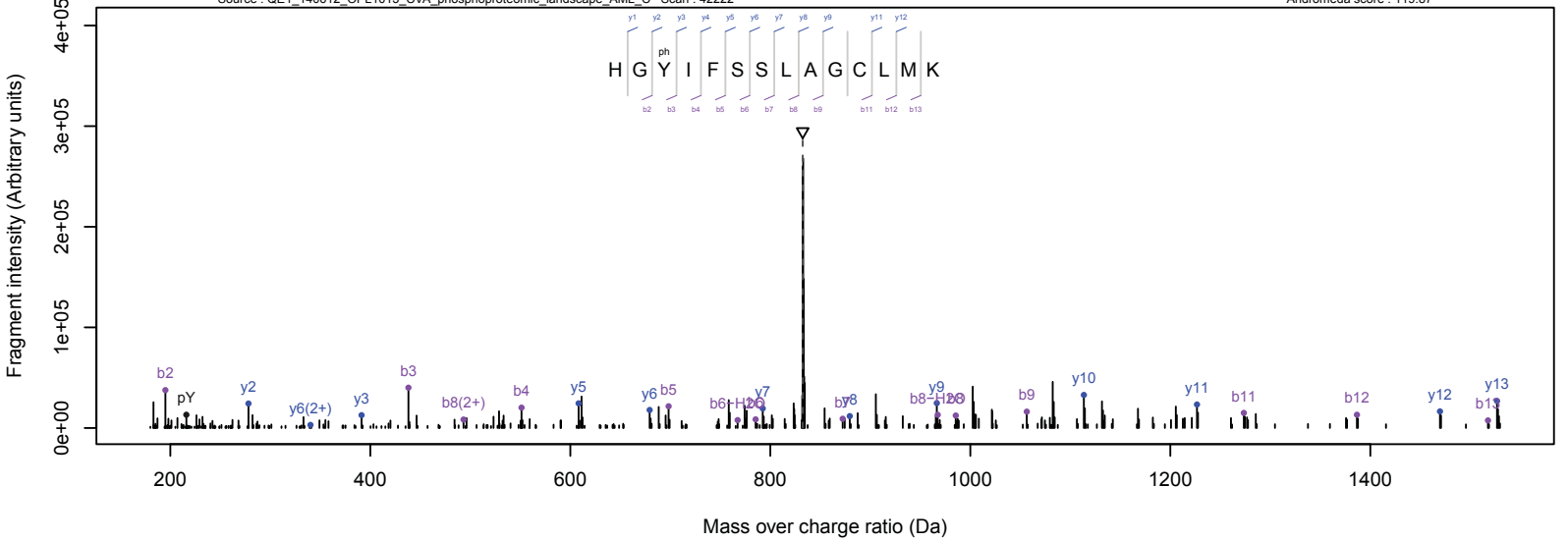
Andromeda score : 120.11



HGY(ph)FSSLAGCLMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42222

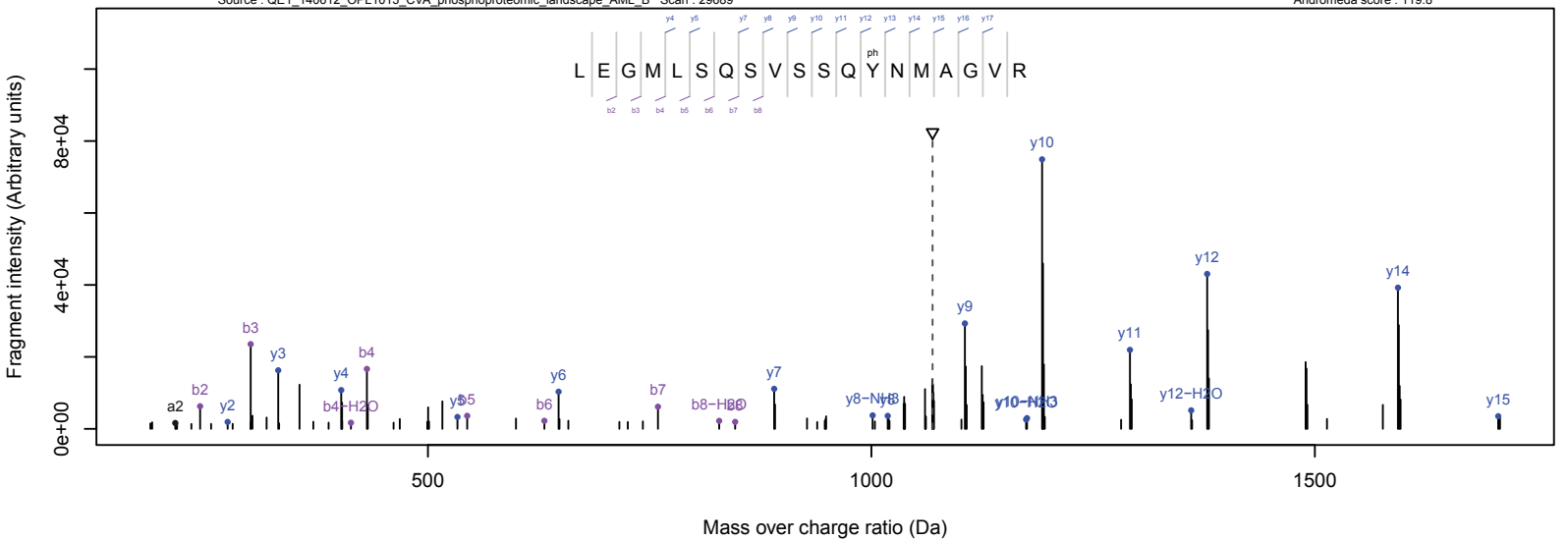
Andromeda score : 119.87



LEGMLSQSVSSQY(ph)NMAGVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29689

Andromeda score : 119.8

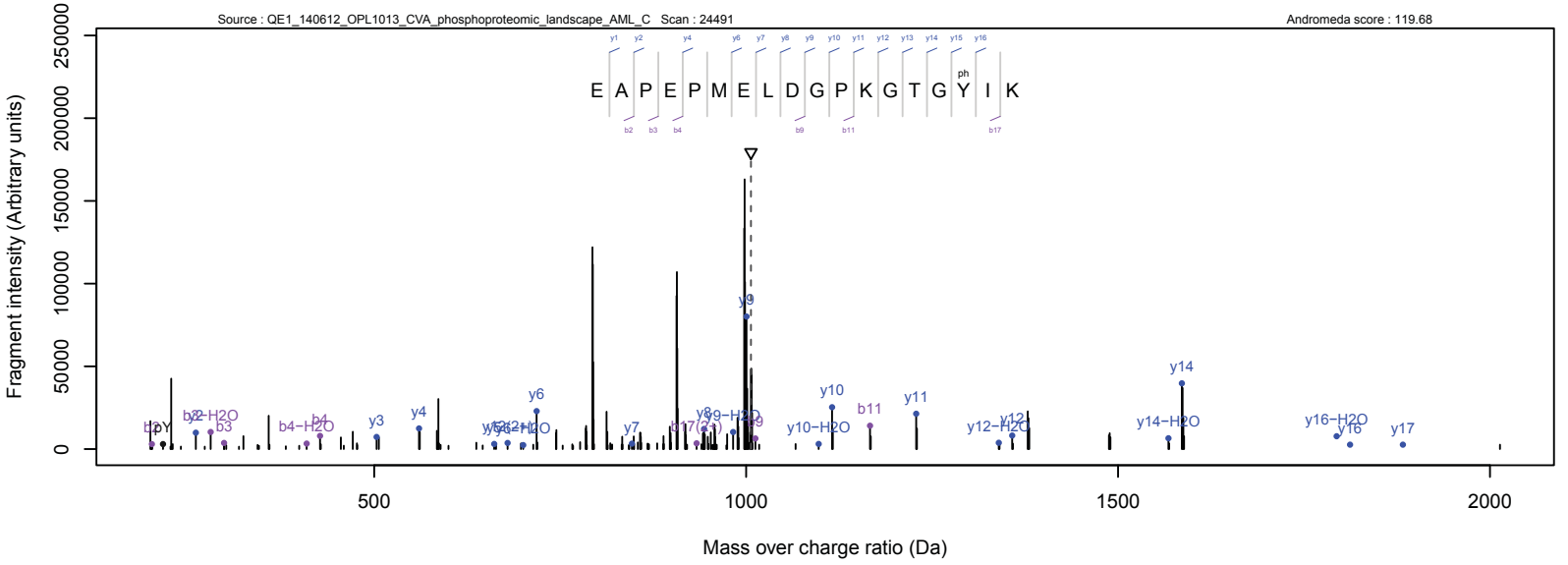




**EAPEPMELDGPKGTGYI<sup>ph</sup>K-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24491

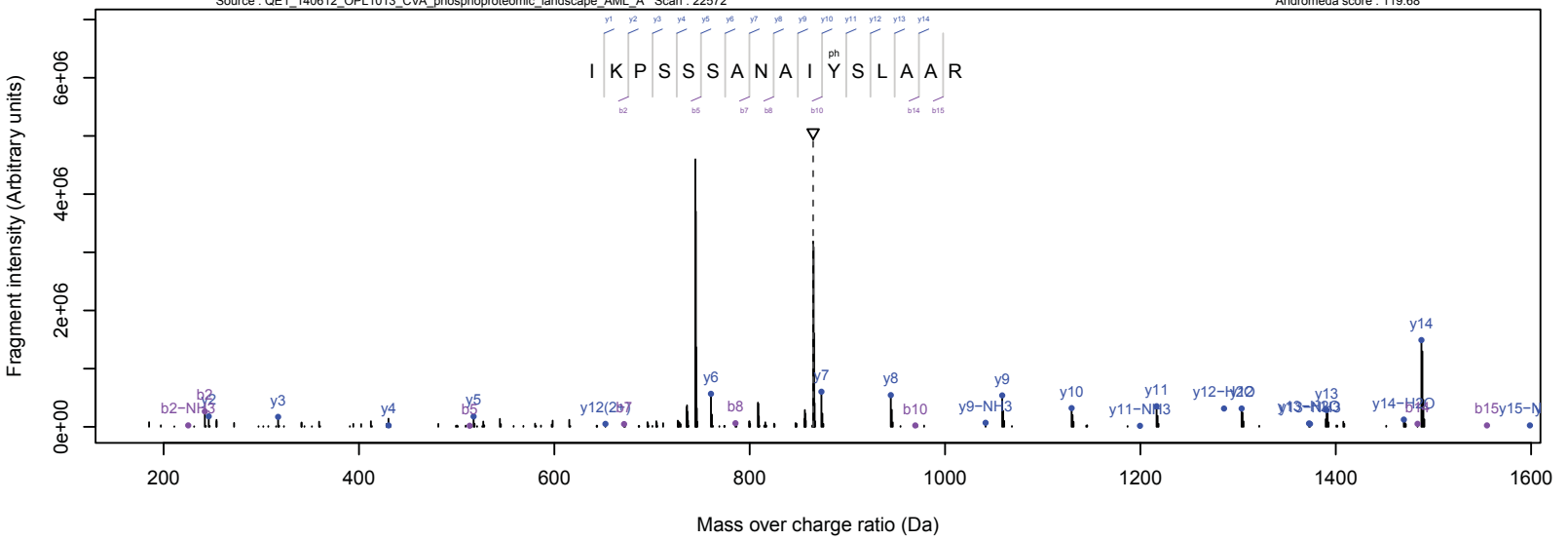
Andromeda score: 119.68



**IKPSSSANA<sup>ph</sup>YSLAAR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 22572

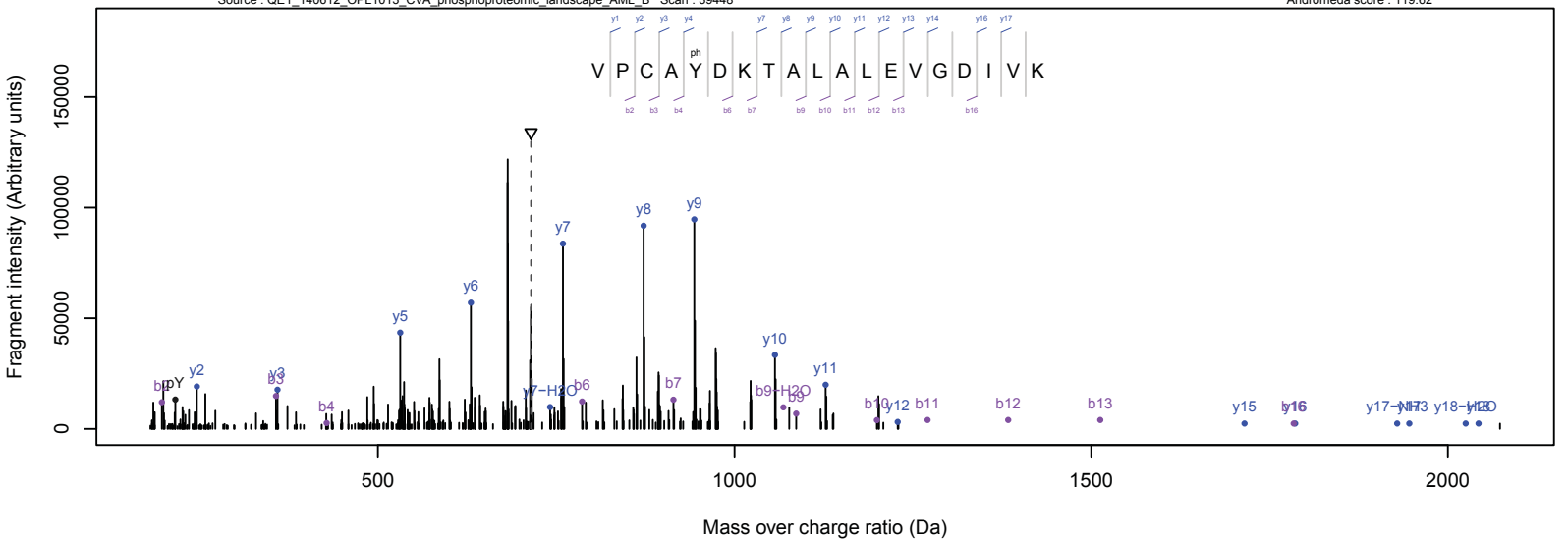
Andromeda score: 119.68



**VPCAY<sup>ph</sup>DKTALALEVGDIVK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 39448

Andromeda score: 119.62

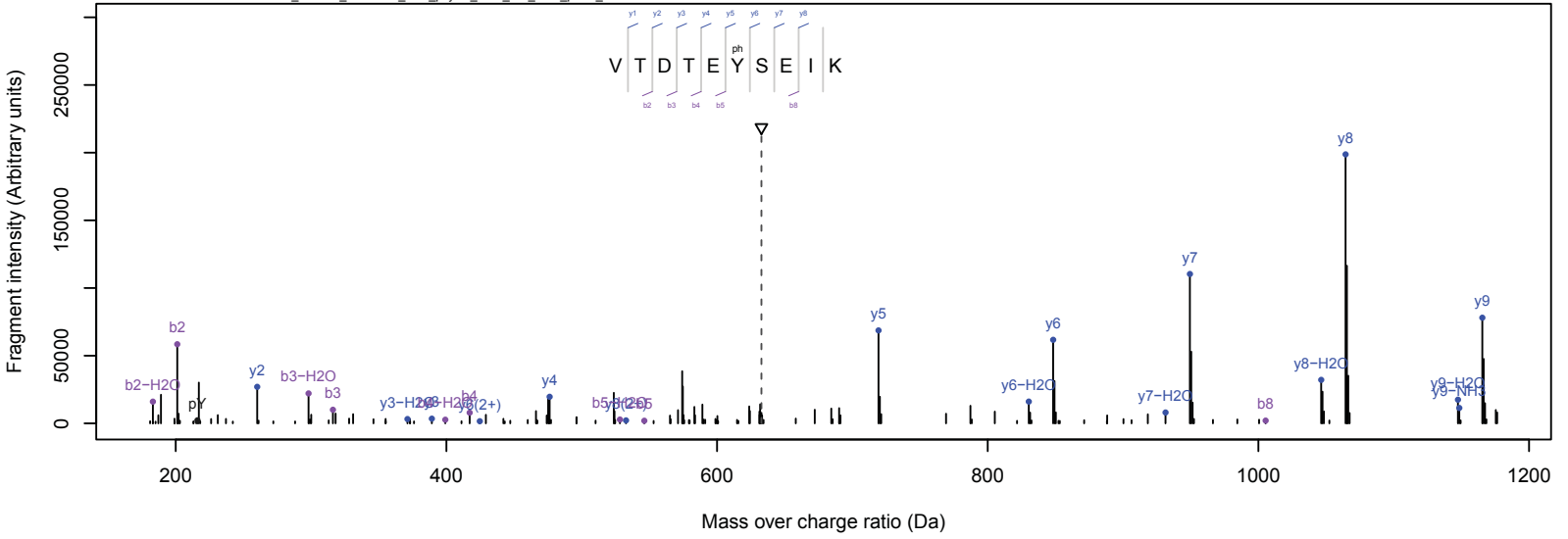




**VTDEY(ph)SEIK-2+**

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 5291

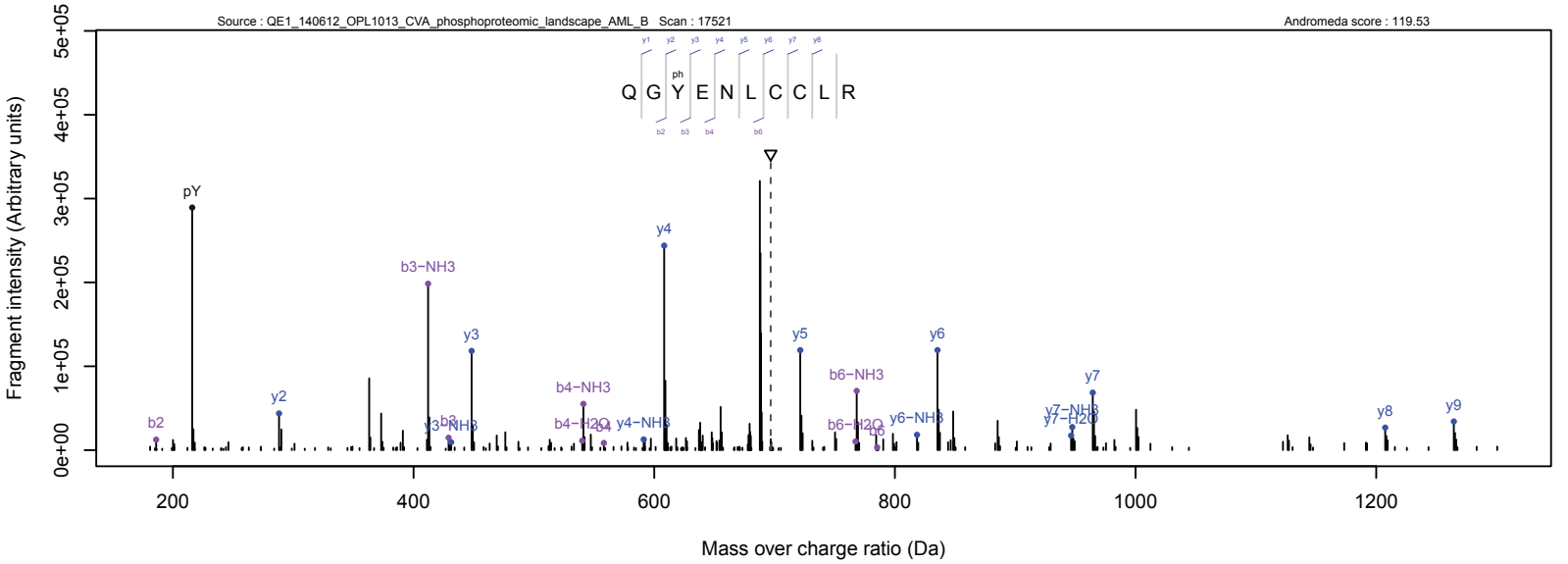
Andromeda score : 119.6



**QGY(ph)ENLCCLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17521

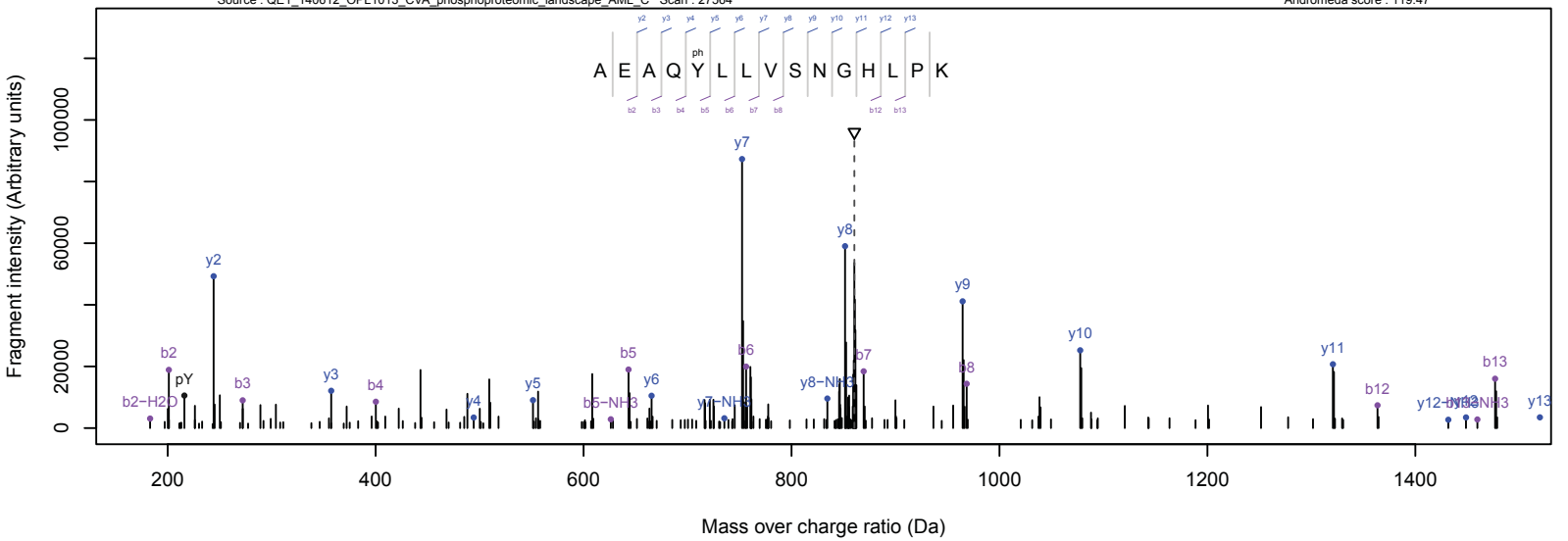
Andromeda score : 119.53



**AEAQY(ph)LLVSNGLHPK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27564

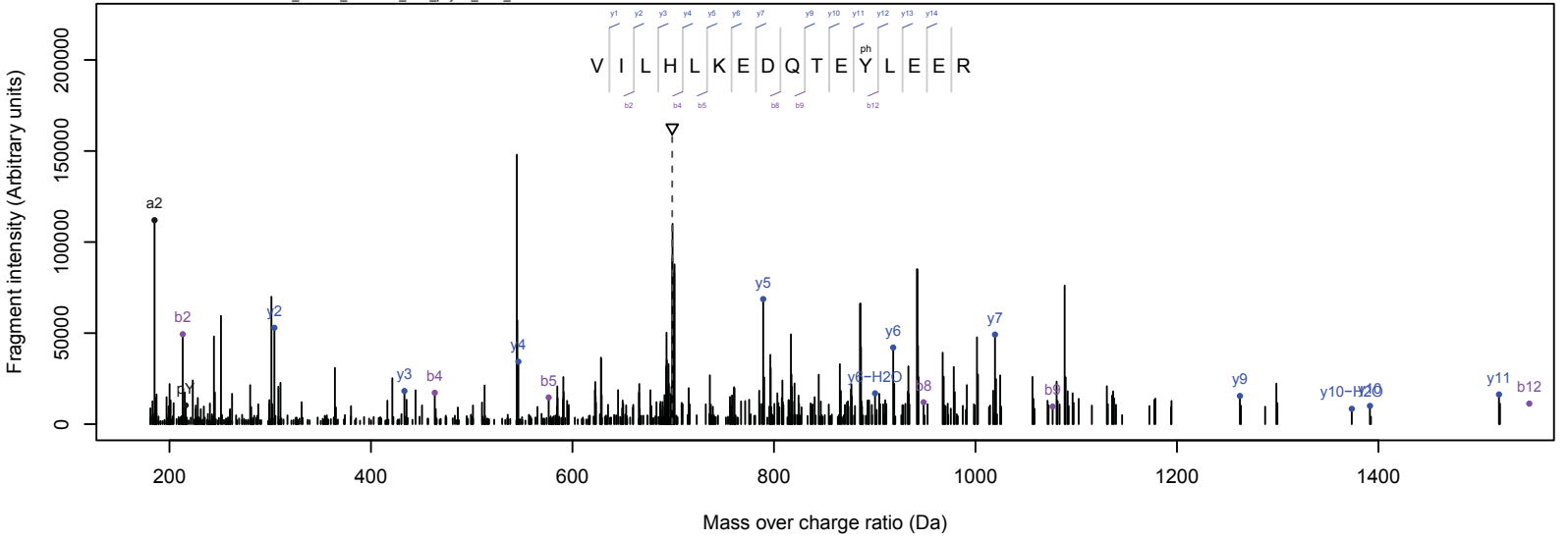
Andromeda score : 119.47



VILHLKEDQTEY(ph)LEER-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 26550

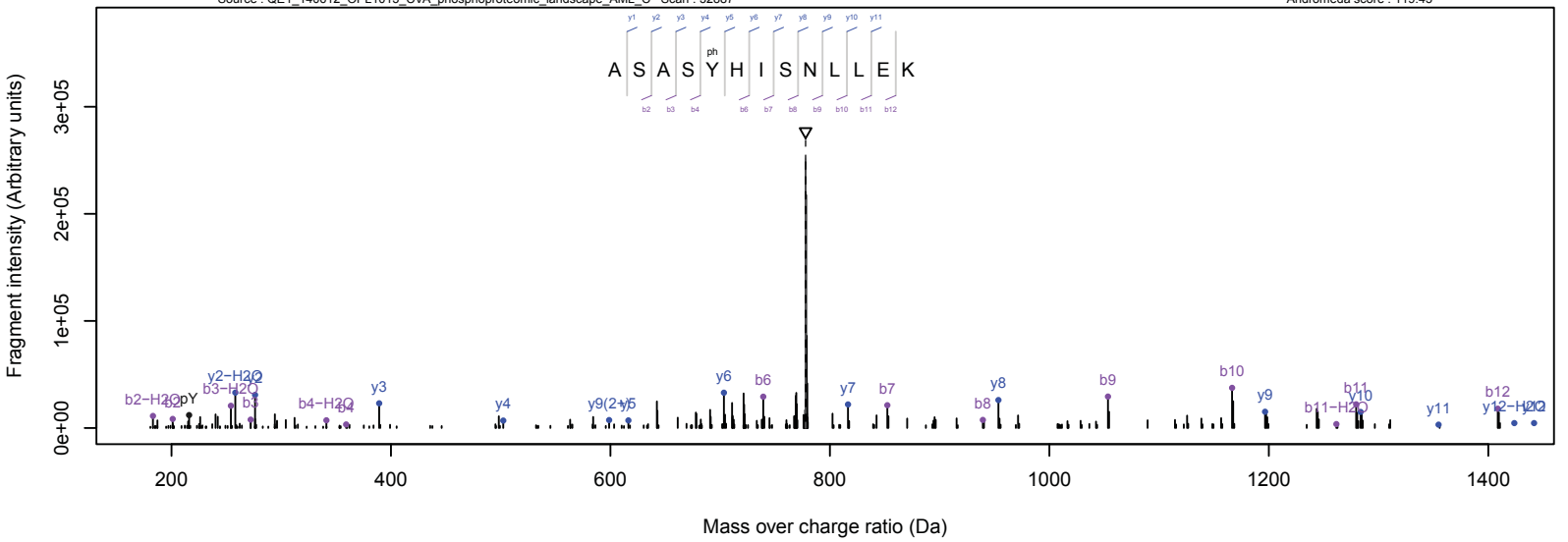
Andromeda score : 119.46



(ac)ASASY(ph)HISNLEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32887

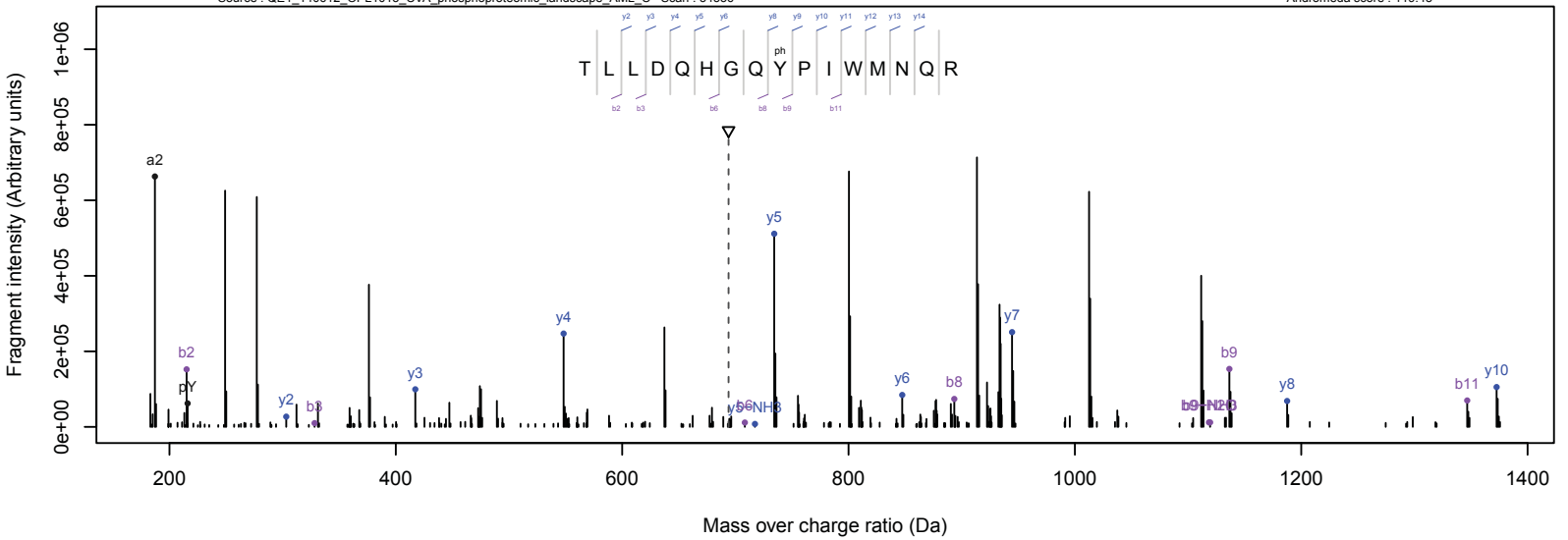
Andromeda score : 119.45



TLLDQHGQY(ph)PIWMNQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34536

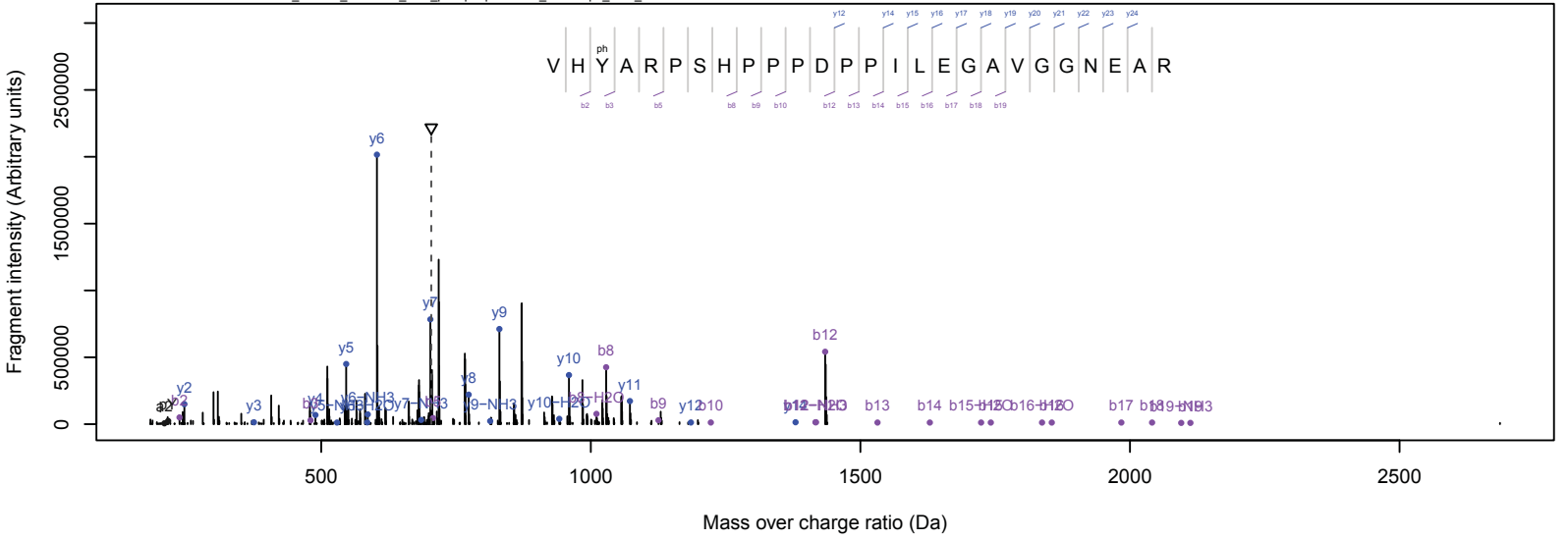
Andromeda score : 119.45



VHY(ph)ARPSHPPDPPILEGAVGGNEAR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22936

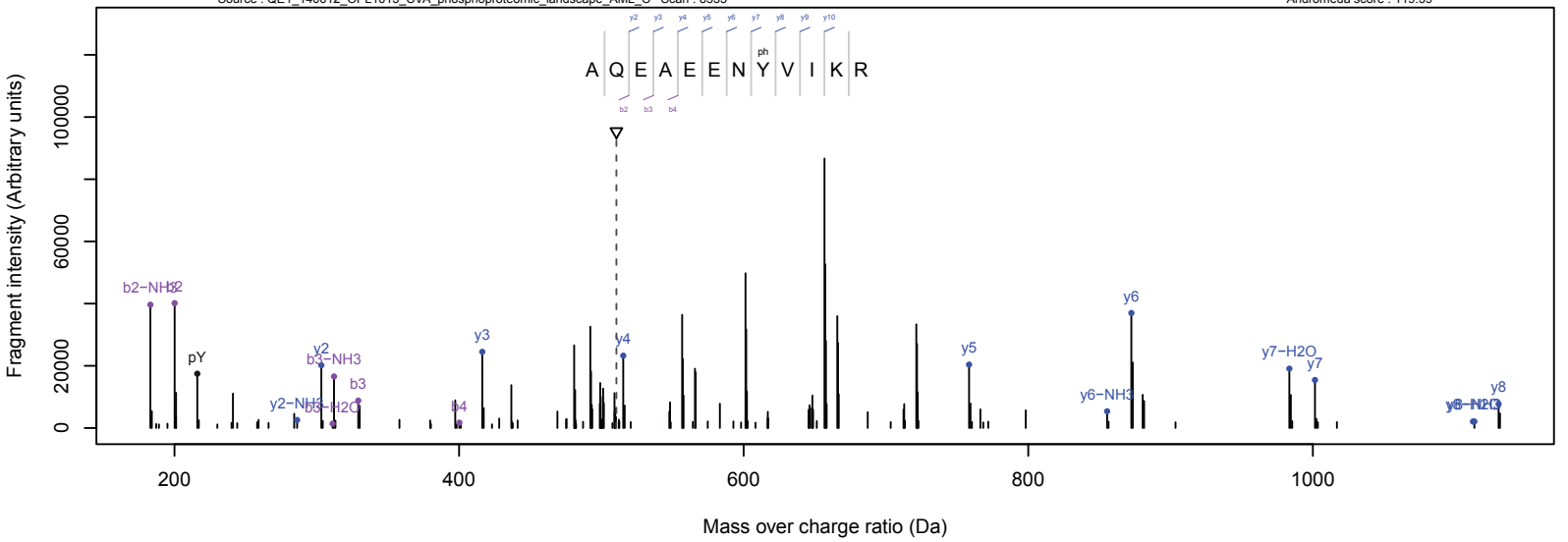
Andromeda score : 119.45



AQEAENY(ph)VIKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8535

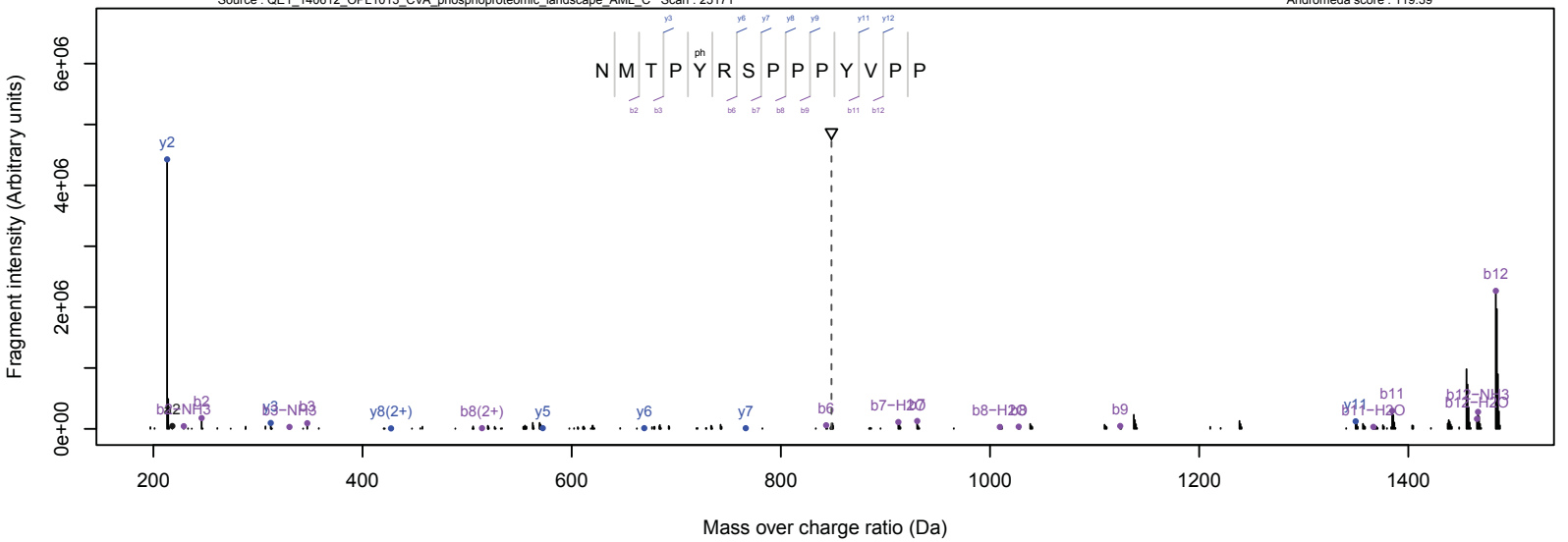
Andromeda score : 119.39



NMTPY(ph)RSPPYVPP-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25171

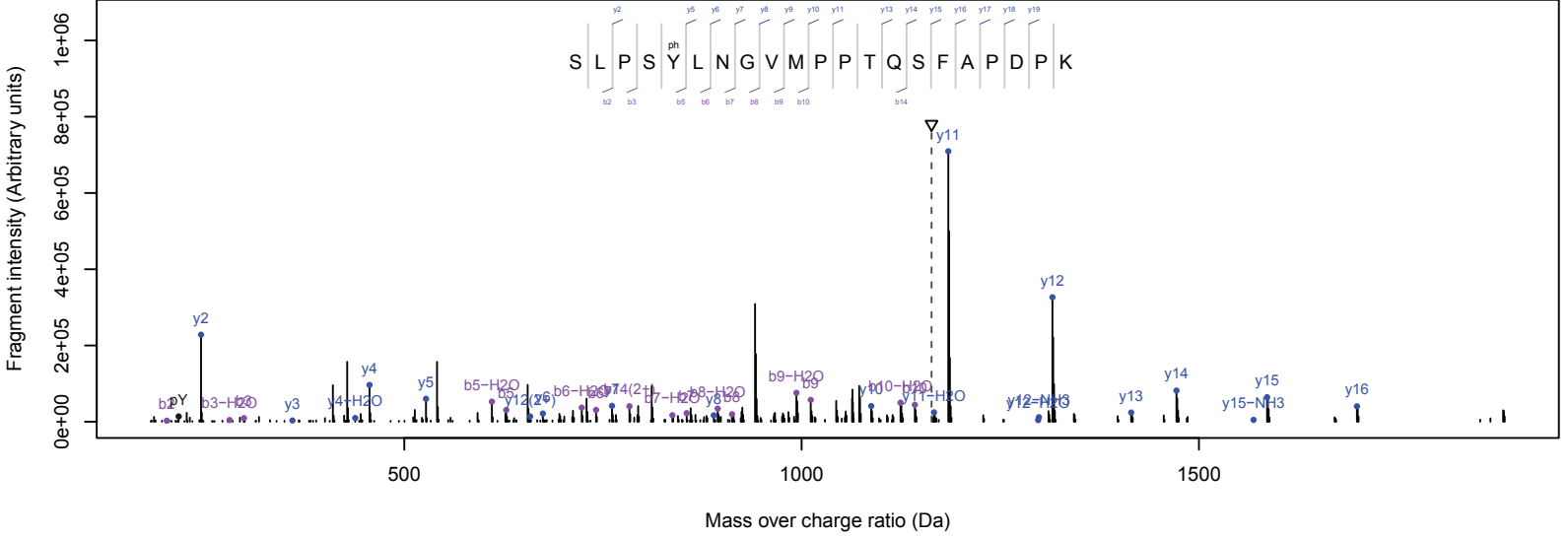
Andromeda score : 119.39



SLPSY(ph)LNGVMPPTQSFAPDPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 44059

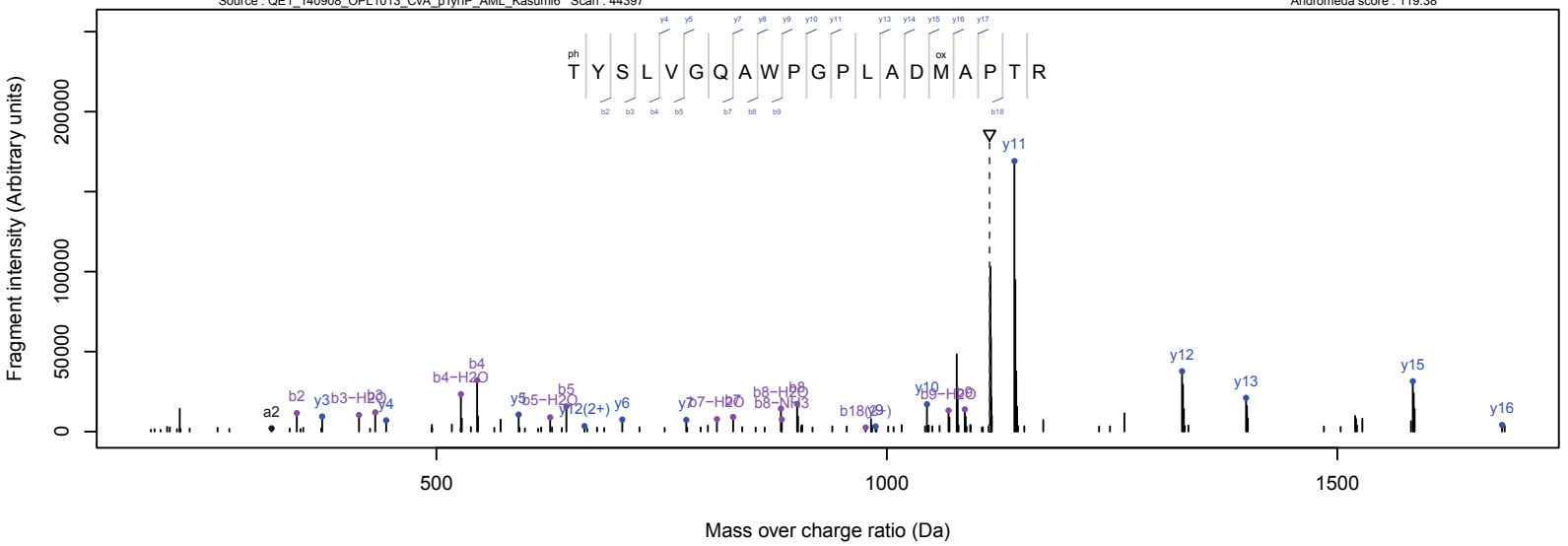
Andromeda score: 119.38



T(ph)YSLVGQAWPGPLADM(ox)APTR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 44397

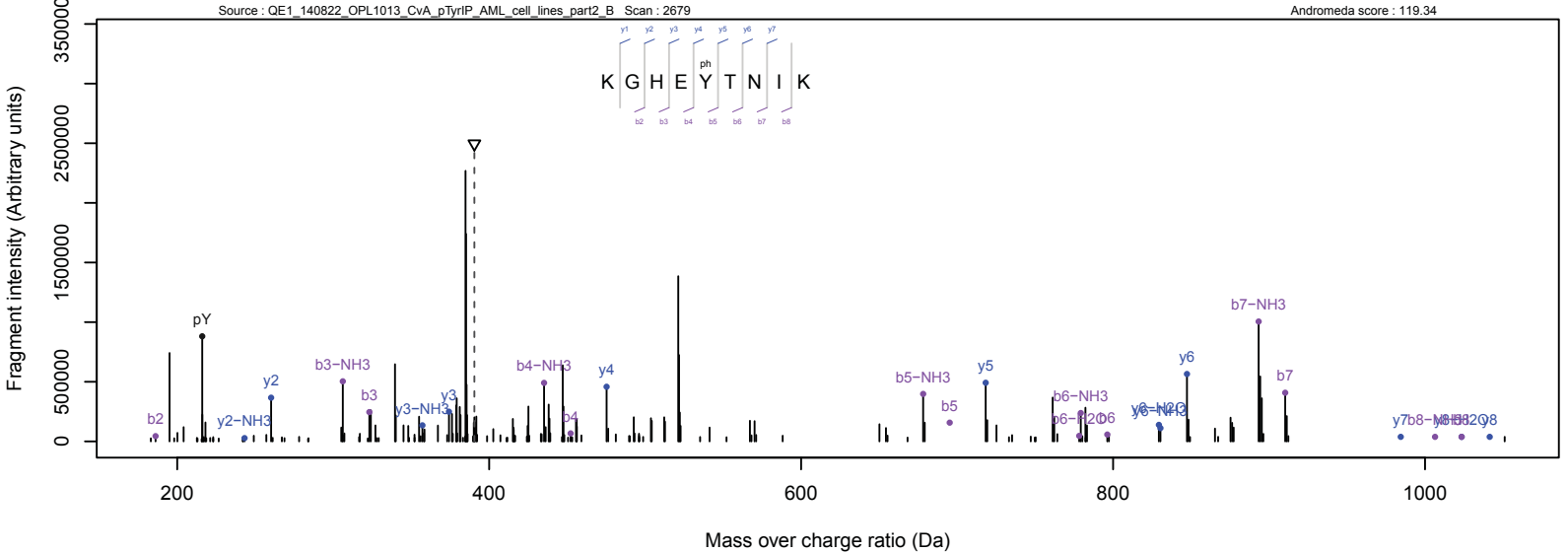
Andromeda score: 119.38



KGHEY(ph)TNIK-3+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 2679

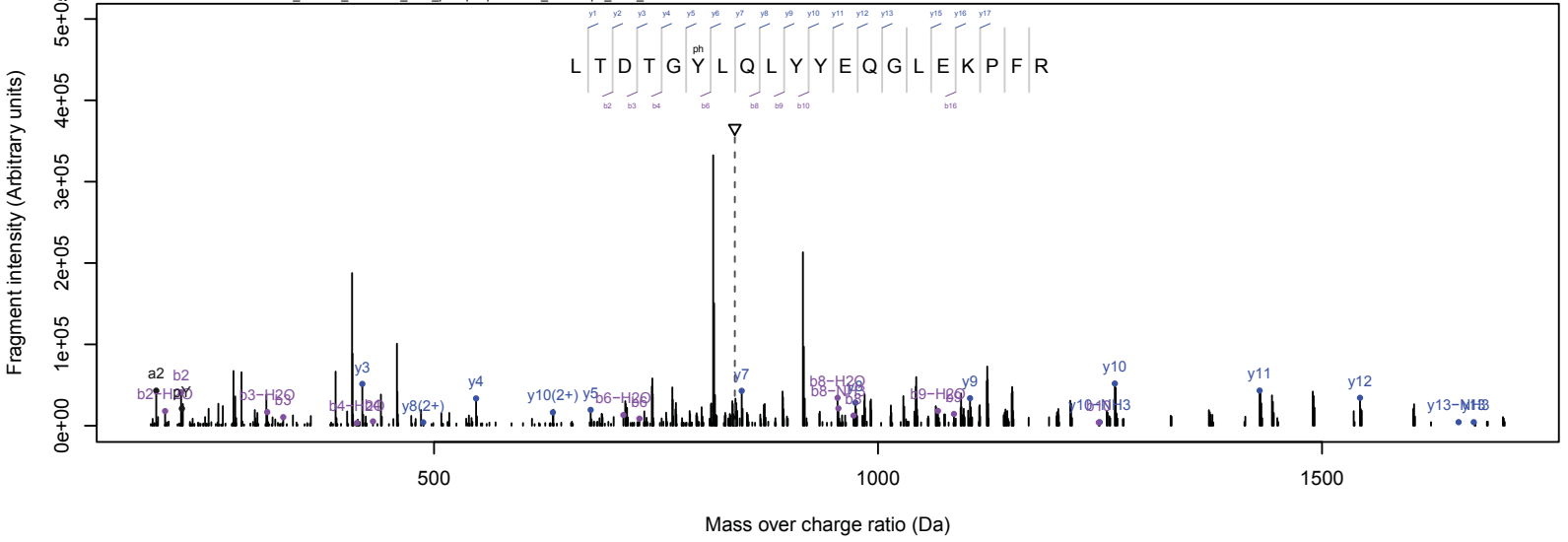
Andromeda score: 119.34



LTDGTG(ph)LQLYYEQGLEKPF<sup>3+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 42514

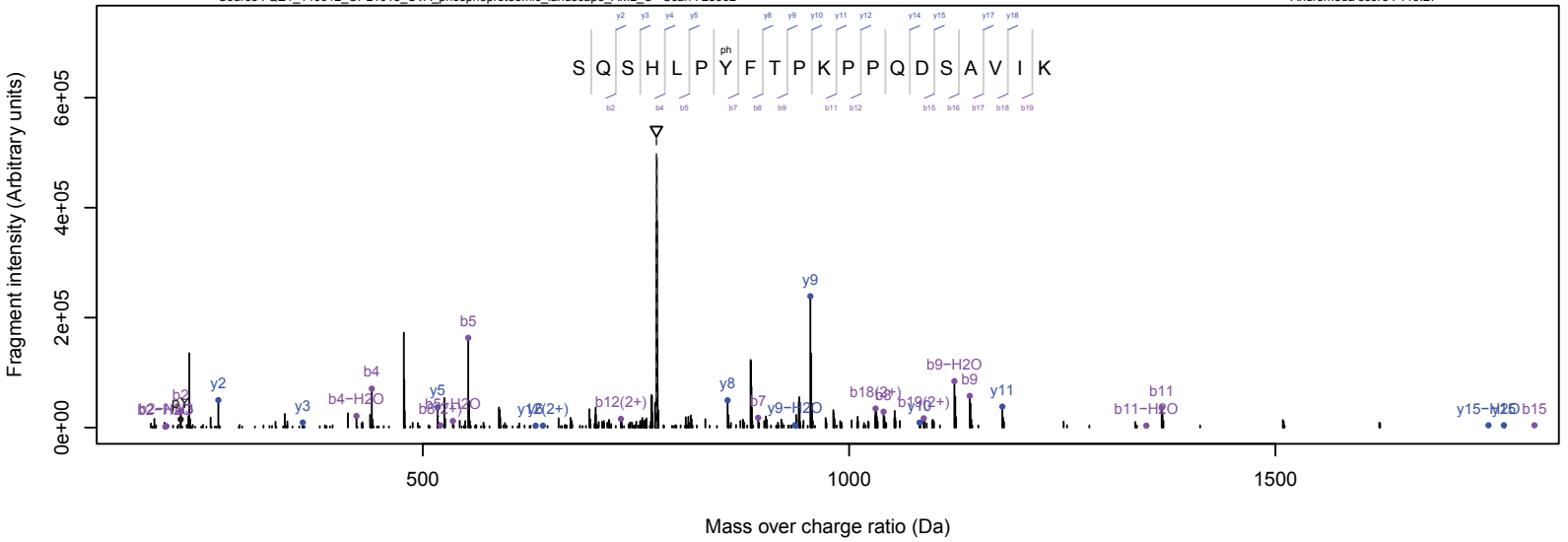
Andromeda score : 119.27



SQSHLPY(ph)FTP<sup>3+</sup>KPPQDSAVIK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28362

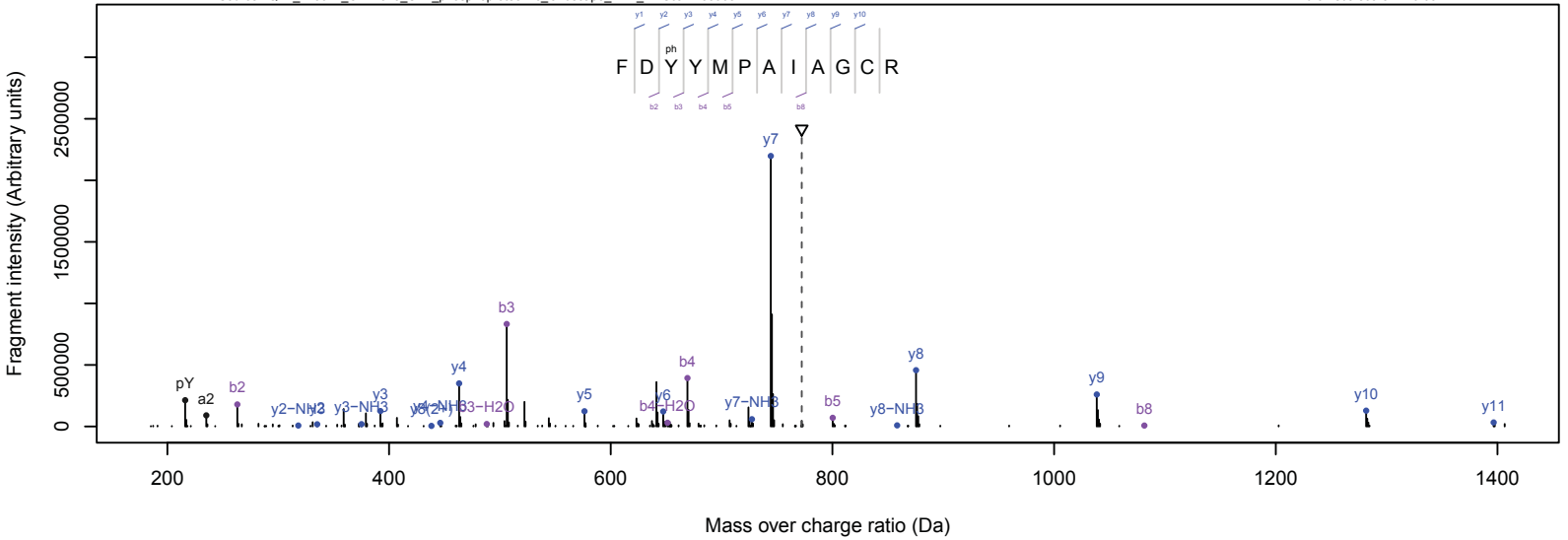
Andromeda score : 119.27



FDY(ph)YMPAIAIGCR<sup>2+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33965

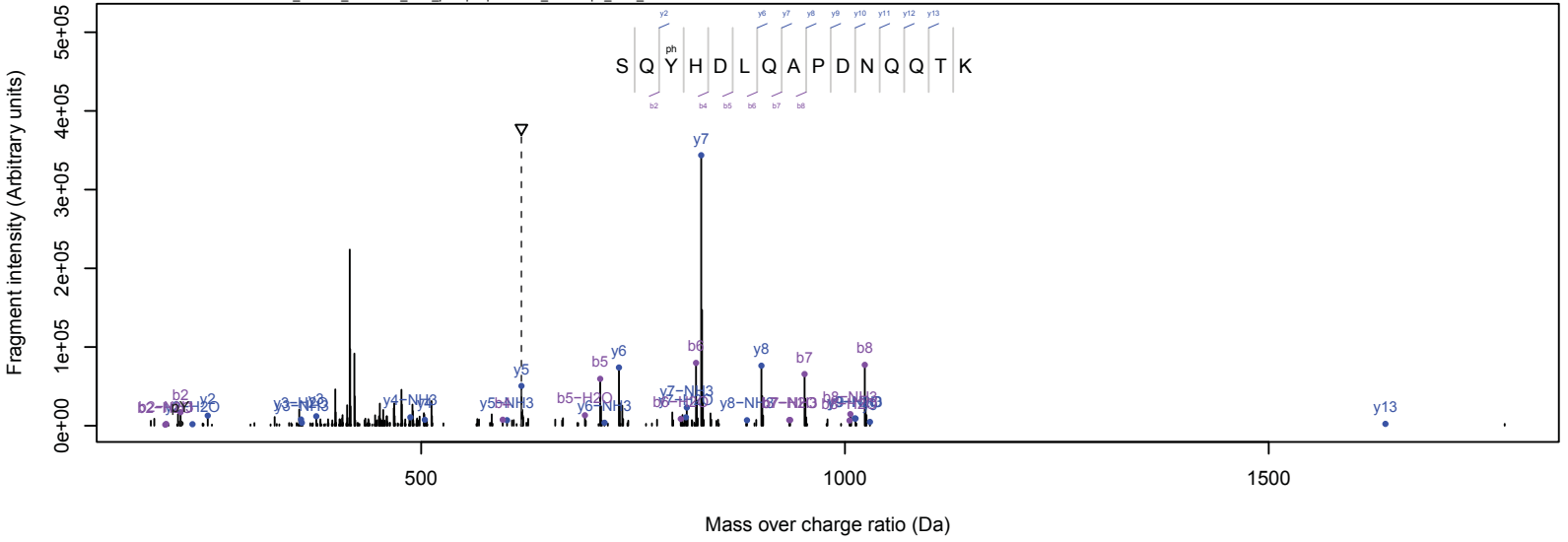
Andromeda score : 119.03



**SQY(ph)HDLQAPDNQQTK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 6467

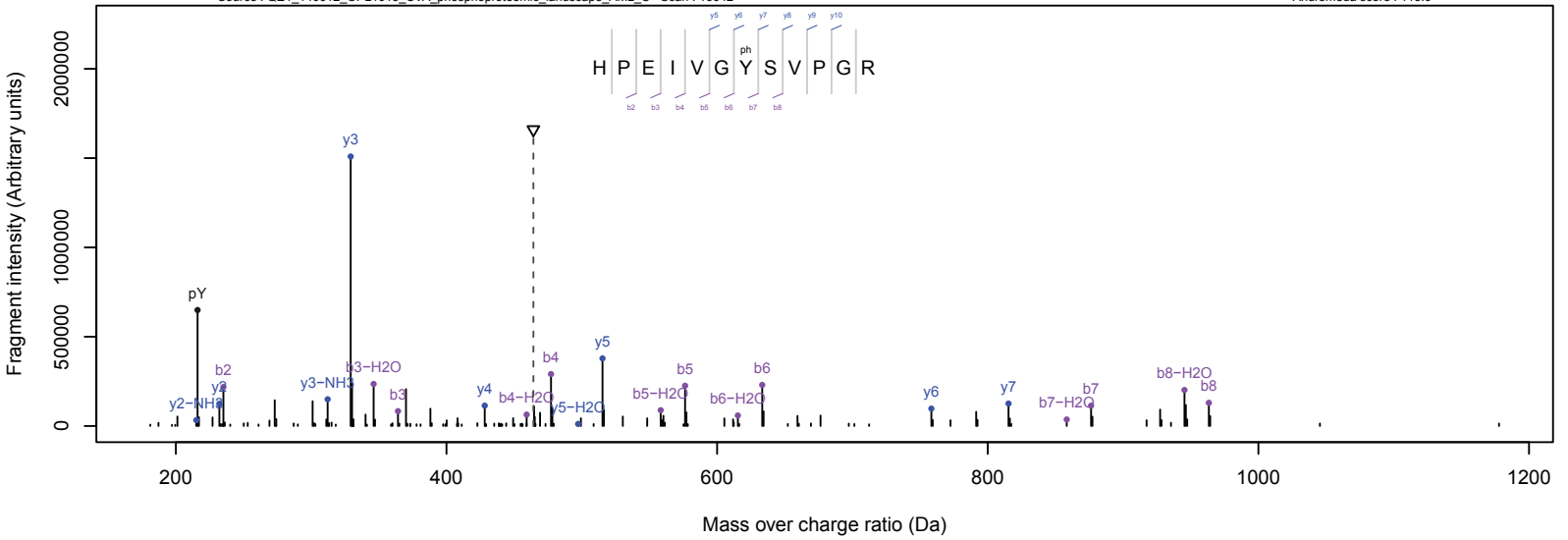
Andromeda score: 119.02



**HPEIVG(ph)SVPGR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16642

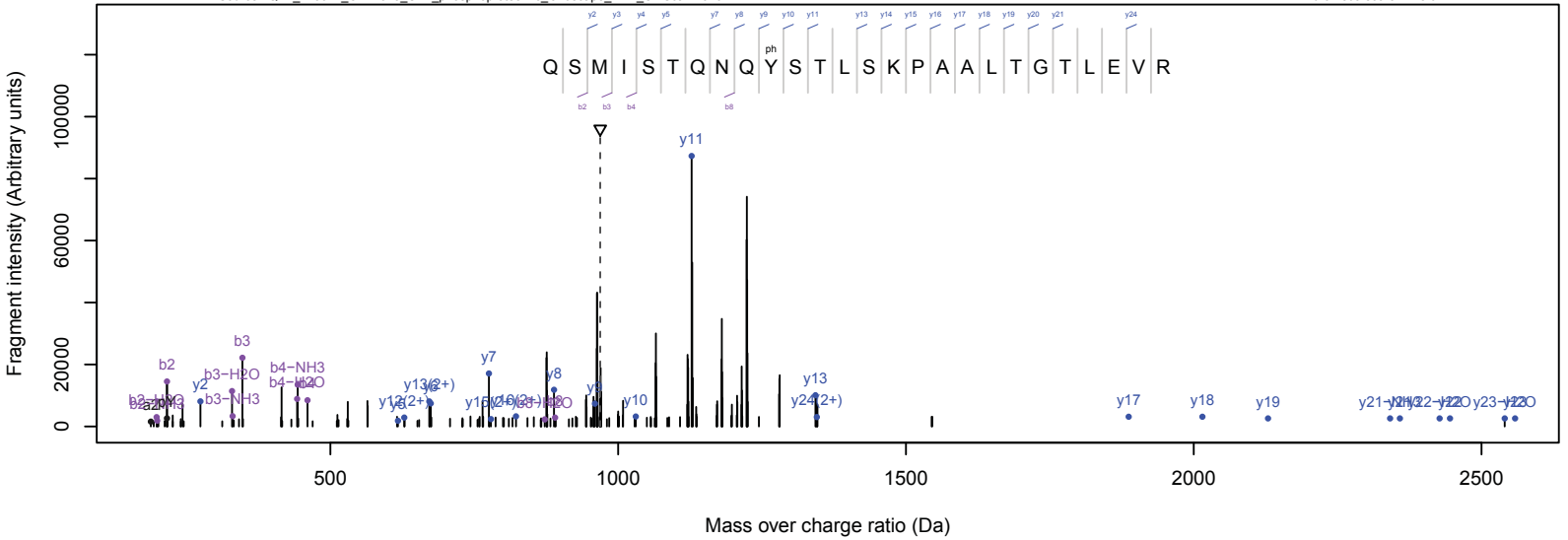
Andromeda score: 118.9



**QSMISTQNQY(ph)STLSKPAALTGTLEVR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34674

Andromeda score: 118.9

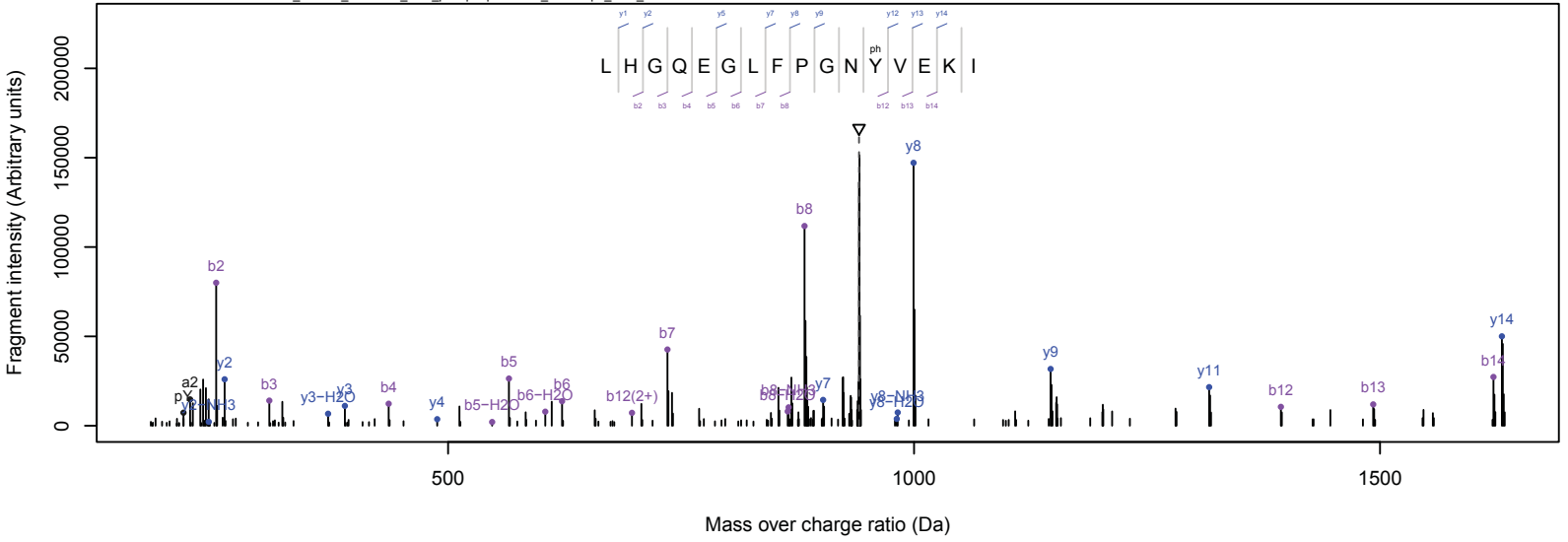




LHGQEGLFPGNY(ph)VEKI-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33846

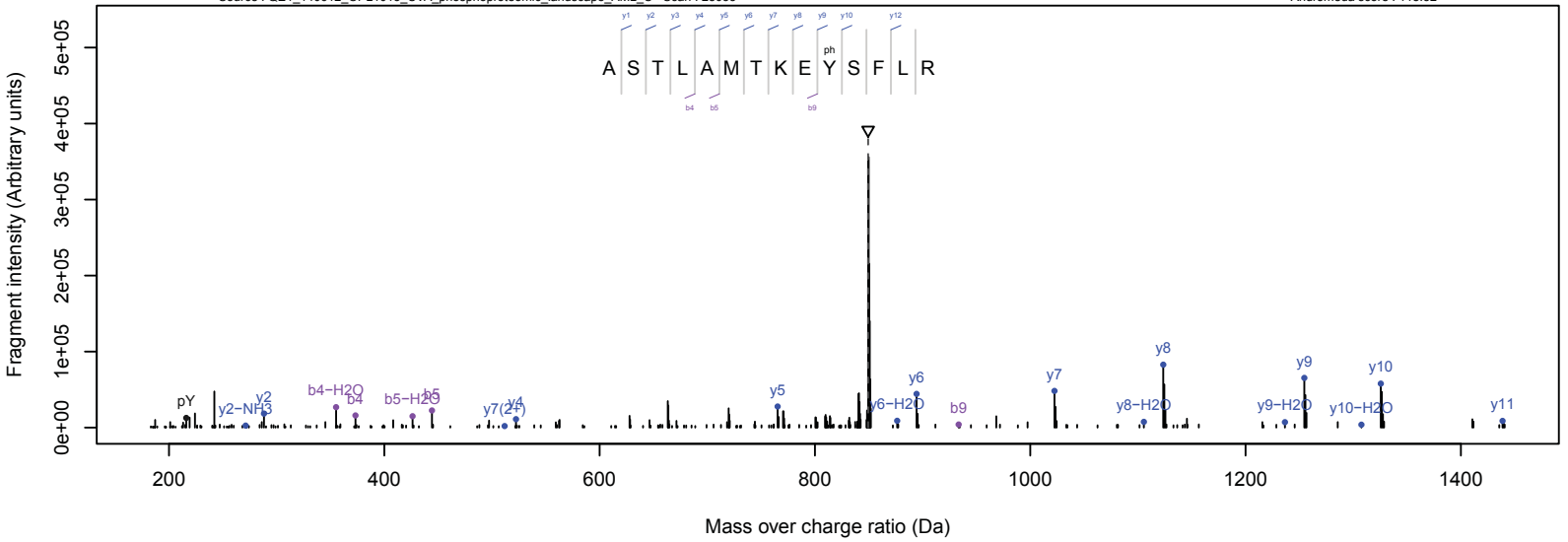
Andromeda score : 118.55



ASTLAMTKEY(ph)SFLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28035

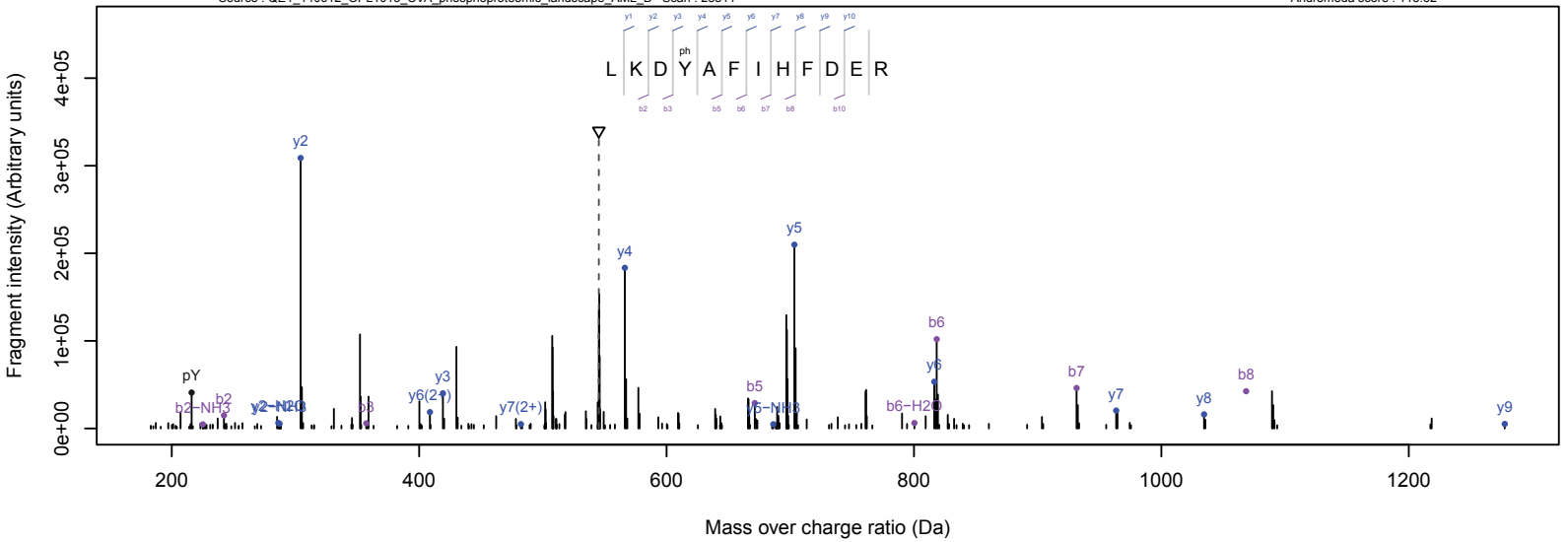
Andromeda score : 118.52



LKDY(ph)AFIHFDER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28511

Andromeda score : 118.52

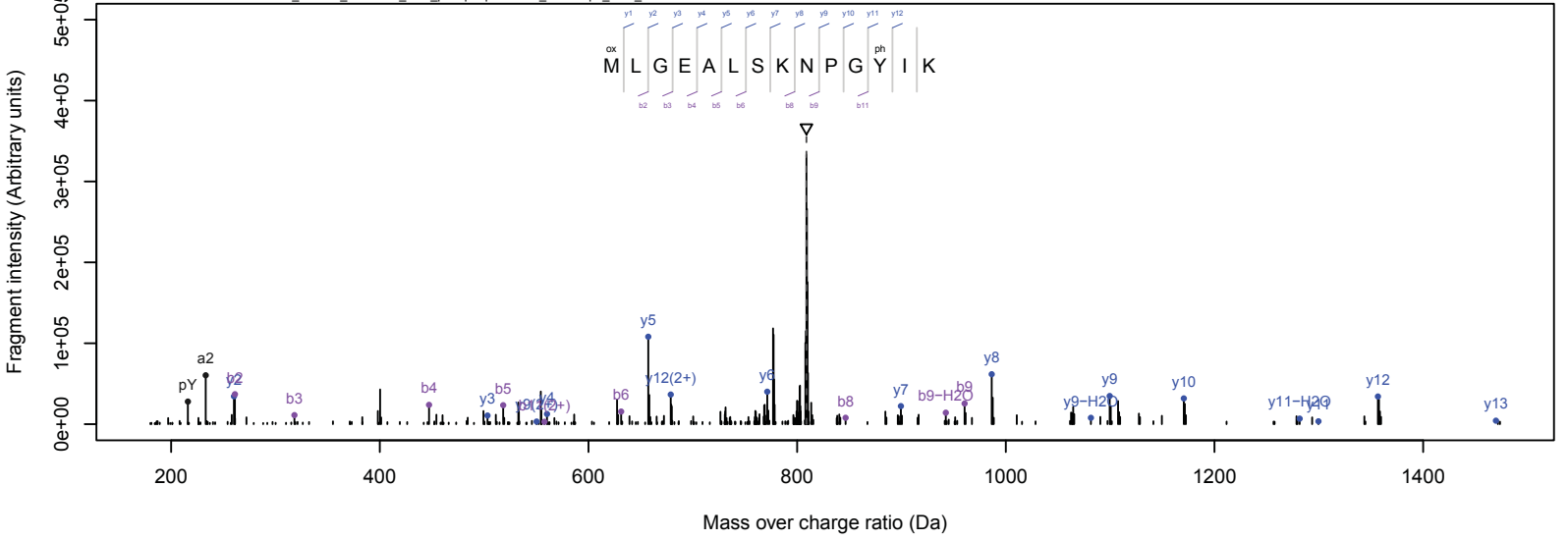




**M(ox)LGEALSKNPGYIK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21624

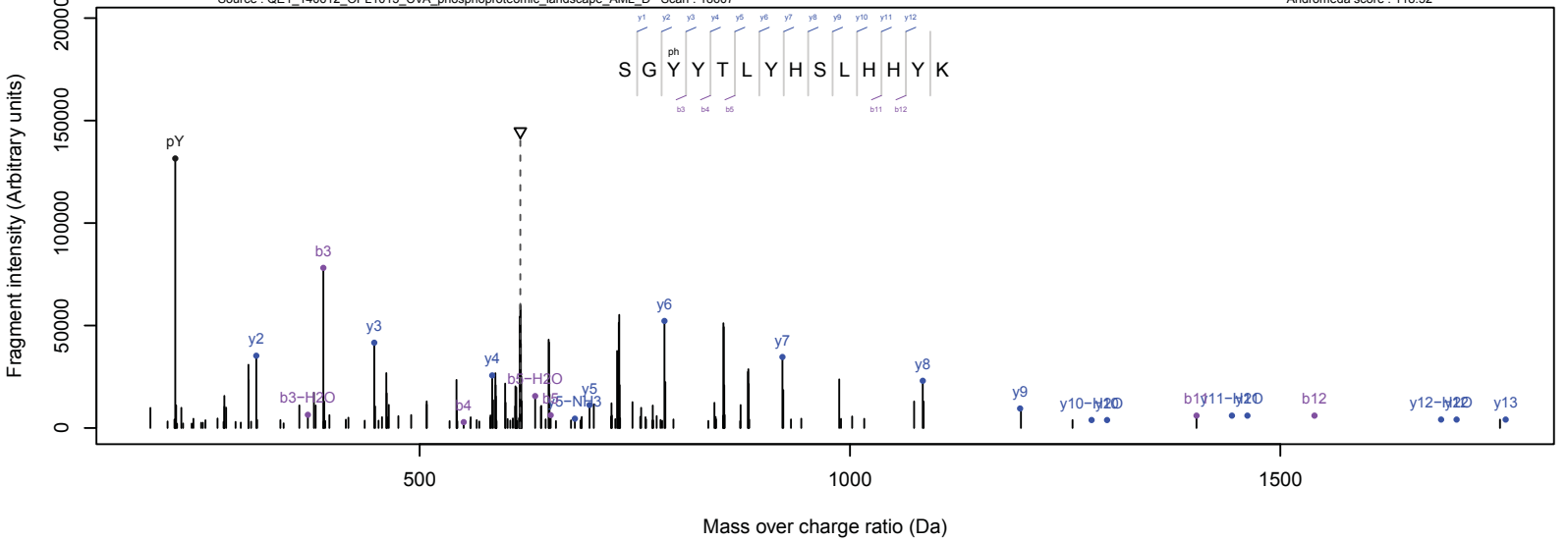
Andromeda score: 118.52



**SGY(ph)YTLYHSLHHYK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 18607

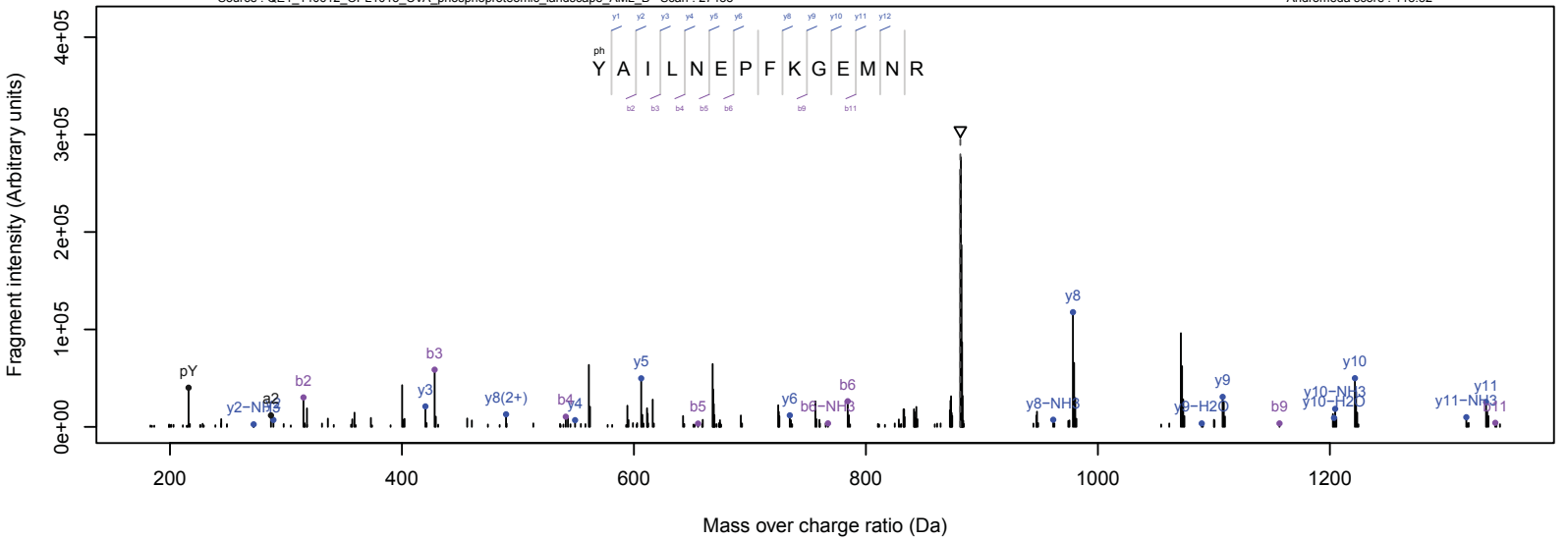
Andromeda score: 118.52



**Y(ph)AILNEPFKGE MN R-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27488

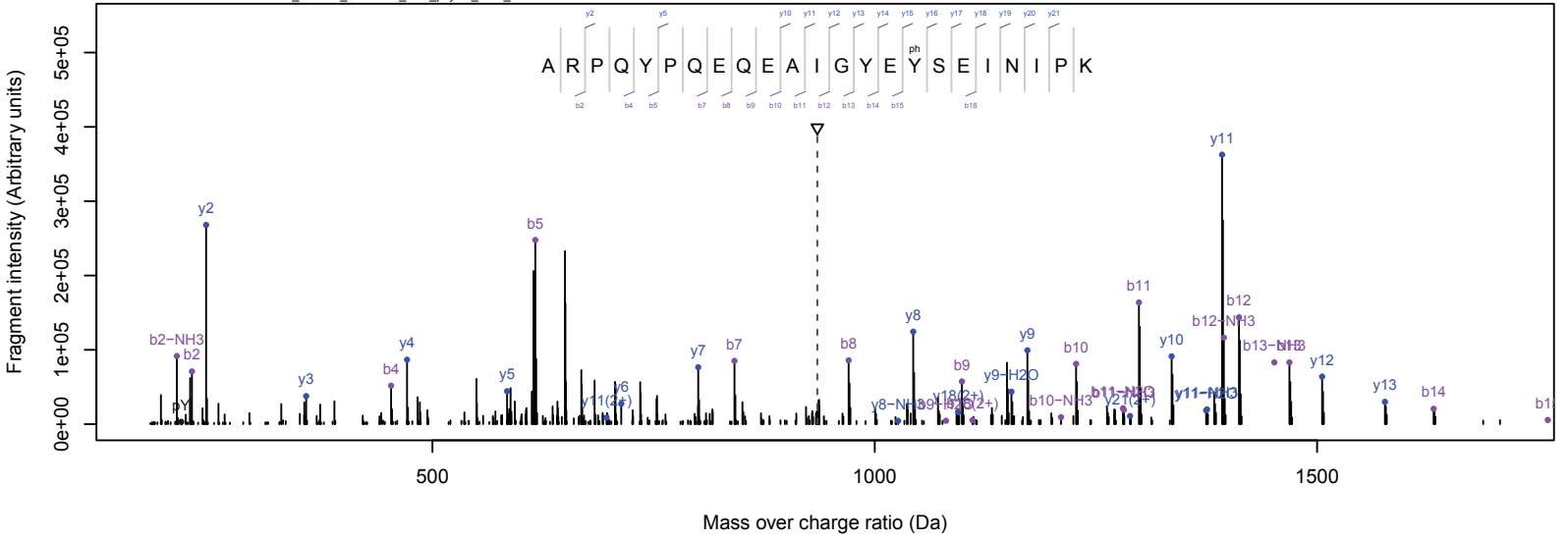
Andromeda score: 118.52



ARPQYPQEAEIGYEY(ph)SEINIPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 33480

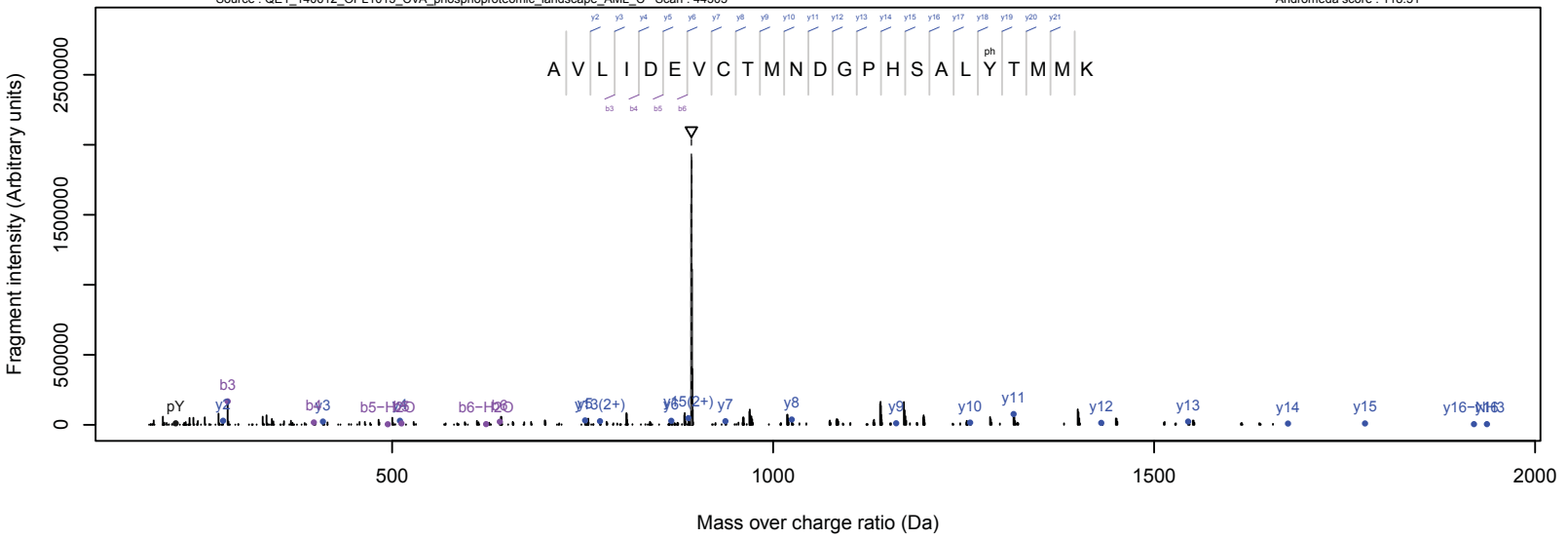
Andromeda score : 118.51



AVLIDEVCTMNDGPHSALY(ph)TMMK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 44305

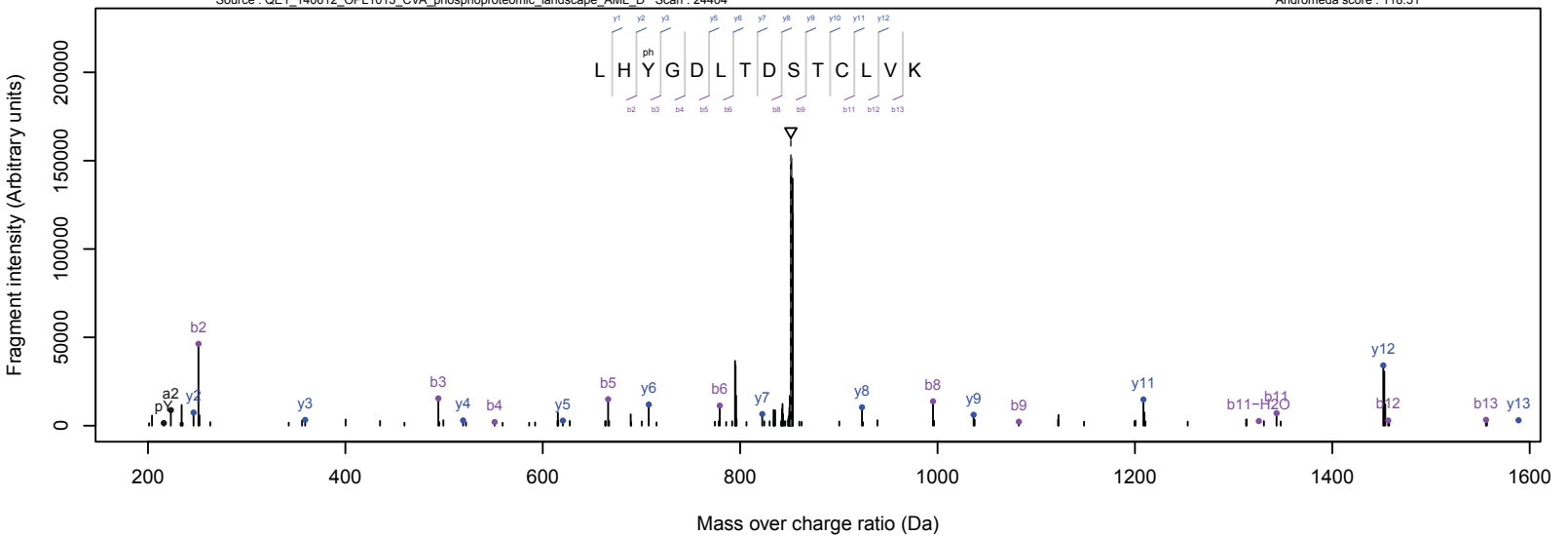
Andromeda score : 118.51



LHY(ph)GDLTDSLTK-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan : 24404

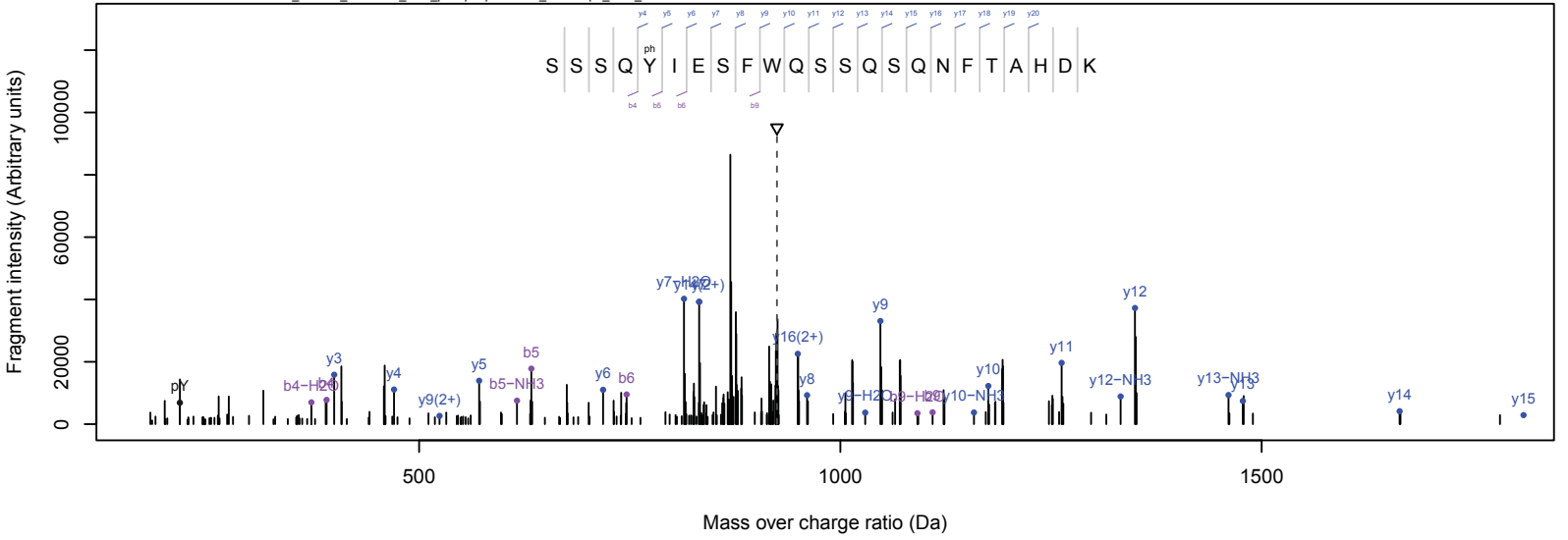
Andromeda score : 118.51



SSSQ(ph)ESFWQSSQSNFTAHDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35971

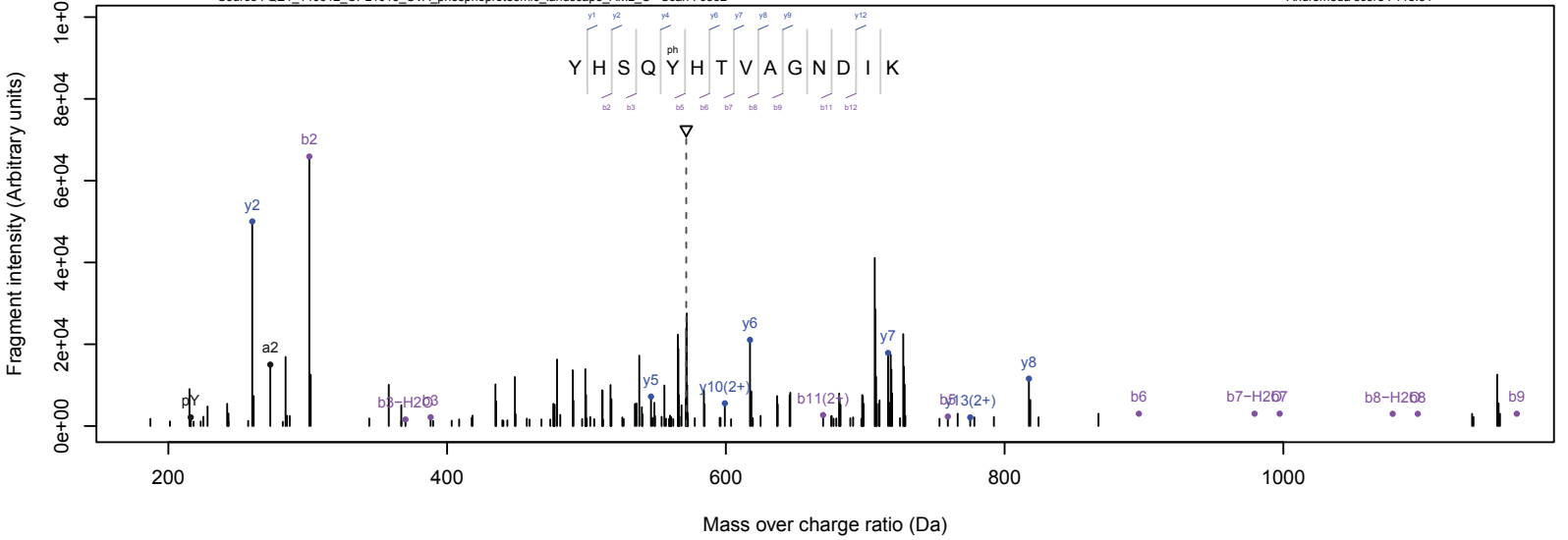
Andromeda score : 118.51



YHSQ(ph)HTVAGNDIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6352

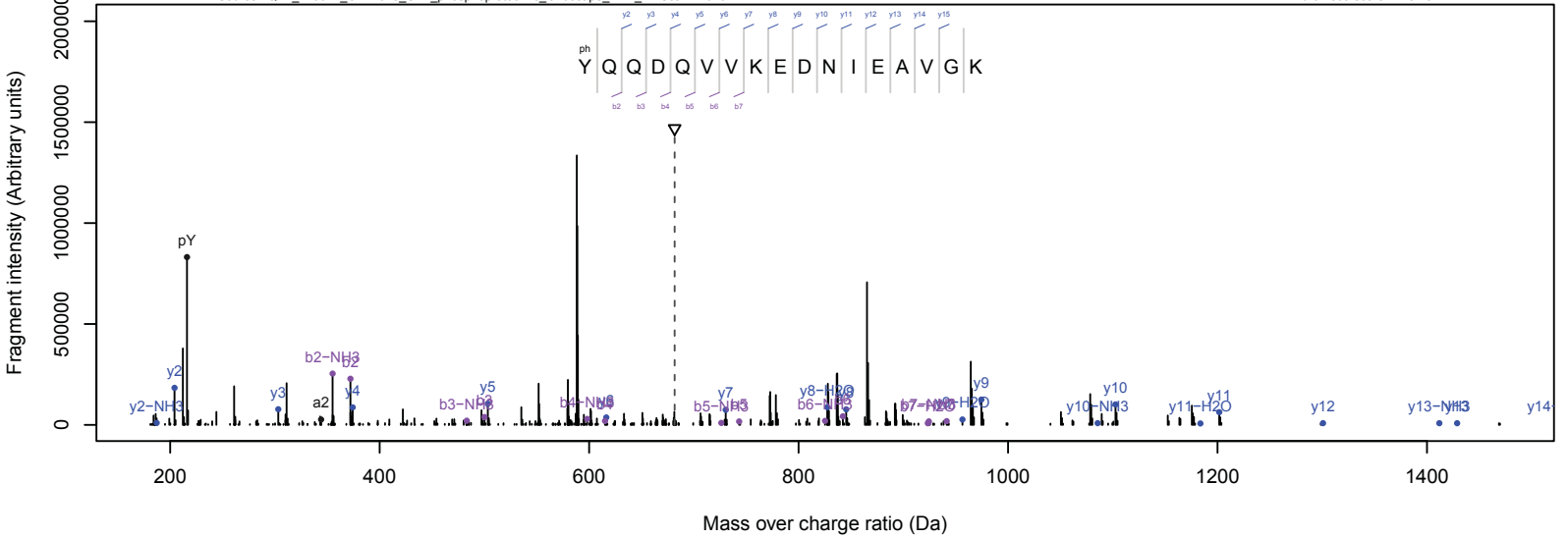
Andromeda score : 118.51



Y(ph)QQDQVVKEDNIEAVGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16734

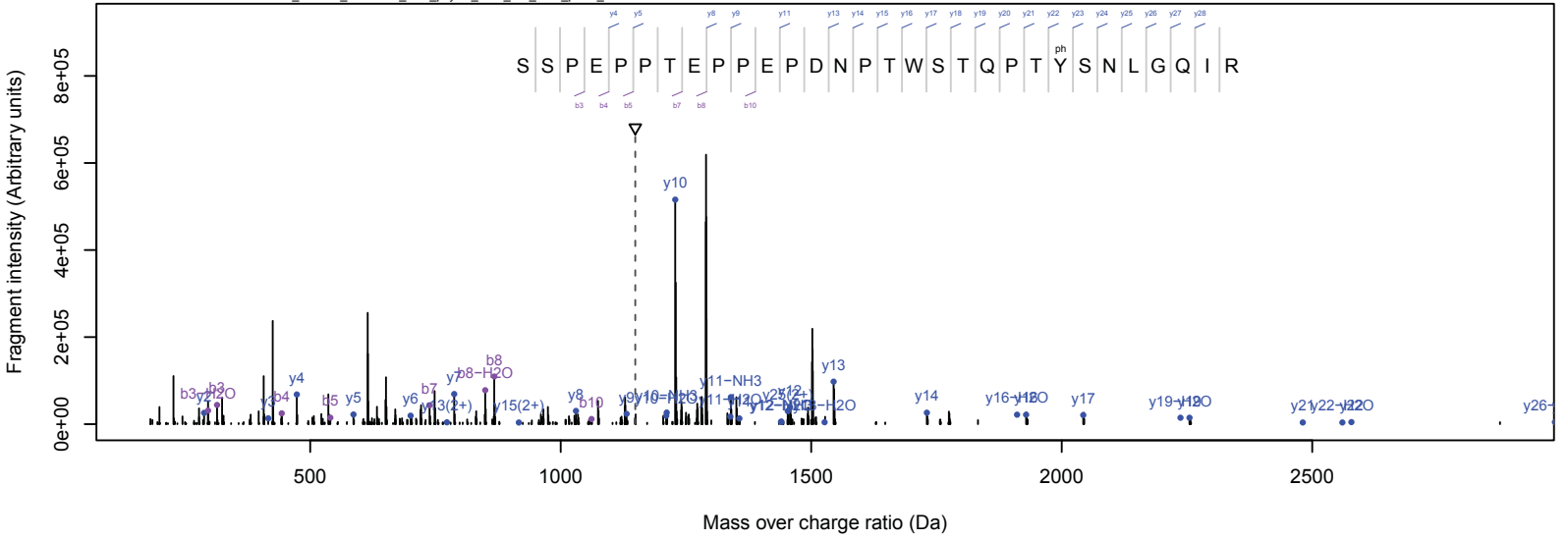
Andromeda score : 118.48



(ac)SSPEPTPEPEPDNPTWSTQPTY(ph)SNLGQIR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 31963

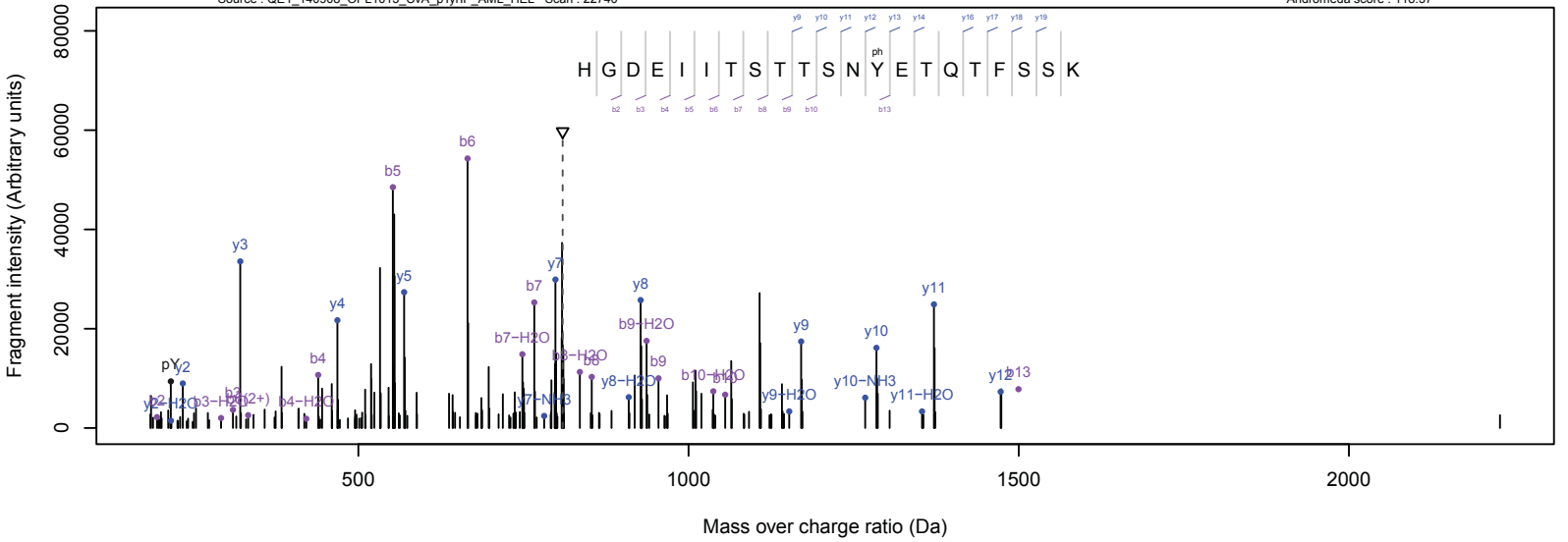
Andromeda score : 118.42



HGDEIITSTTSNY(ph)ETQTFSSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 22740

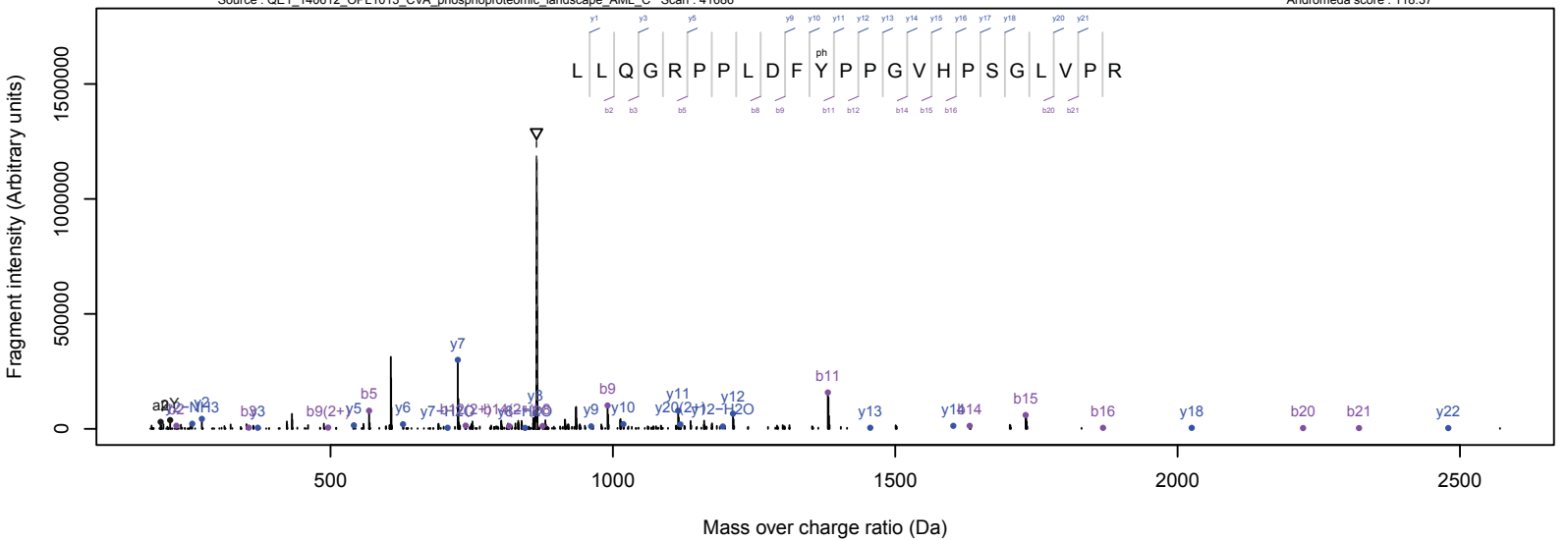
Andromeda score : 118.37



LLQGRPLDFY(ph)PPGVHPSGLVPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41686

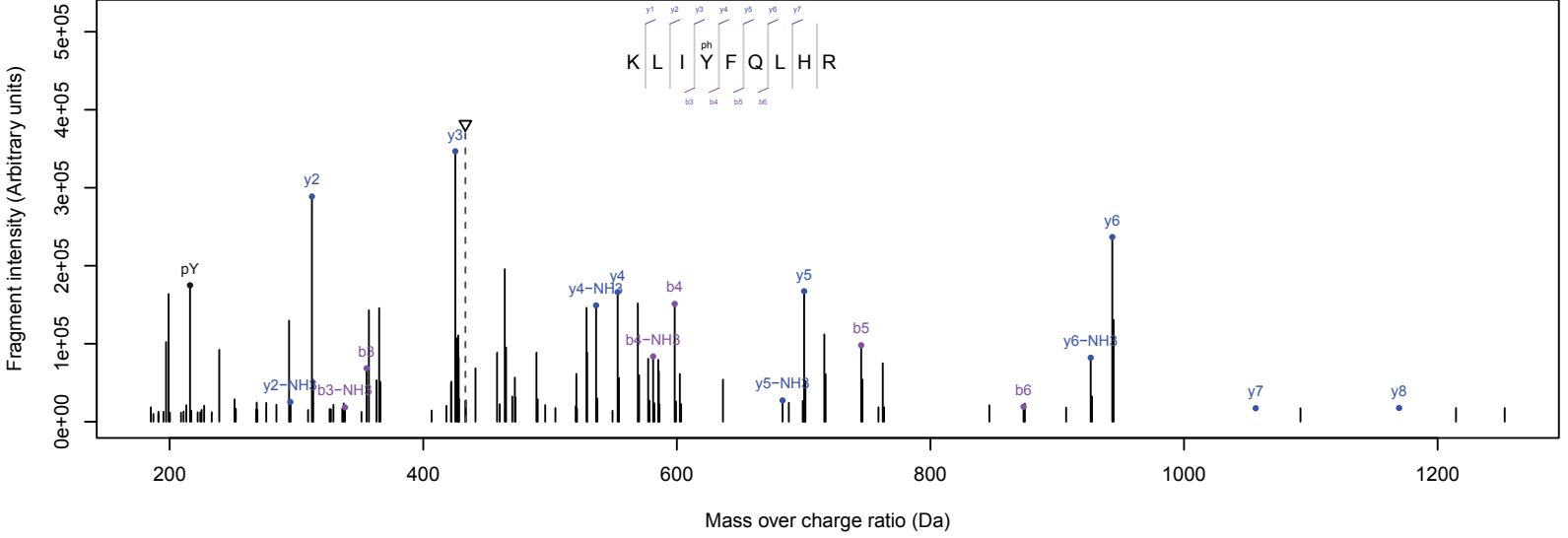
Andromeda score : 118.37



**KLIY(ph)FQLHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27432

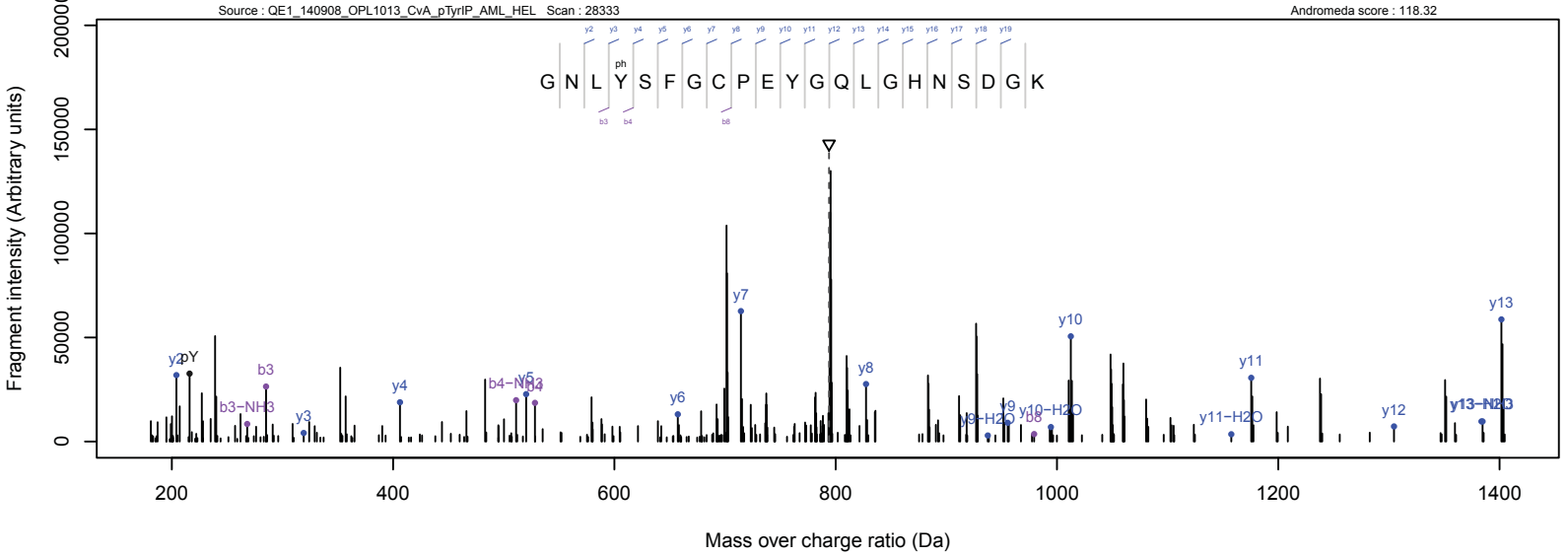
Andromeda score : 118.33



**GNLY(ph)SFGCPYGLGHNSDGK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 28333

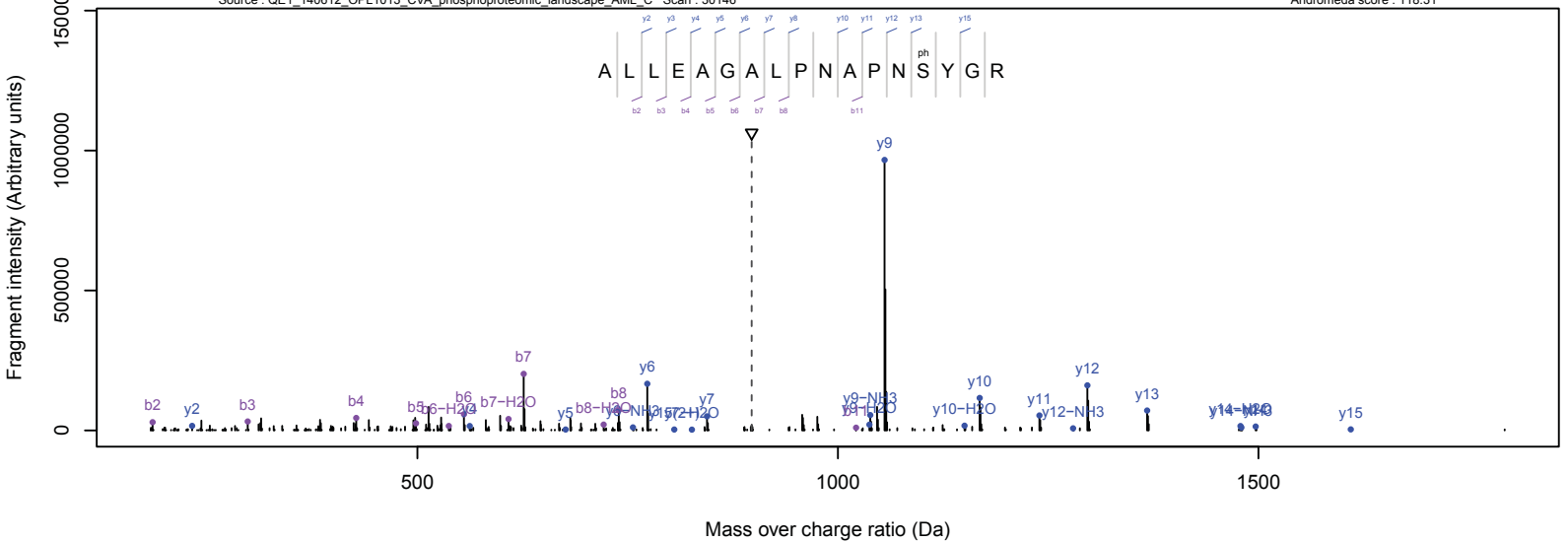
Andromeda score : 118.32



**ALLEAGALPNAPNS(ph)YGR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30146

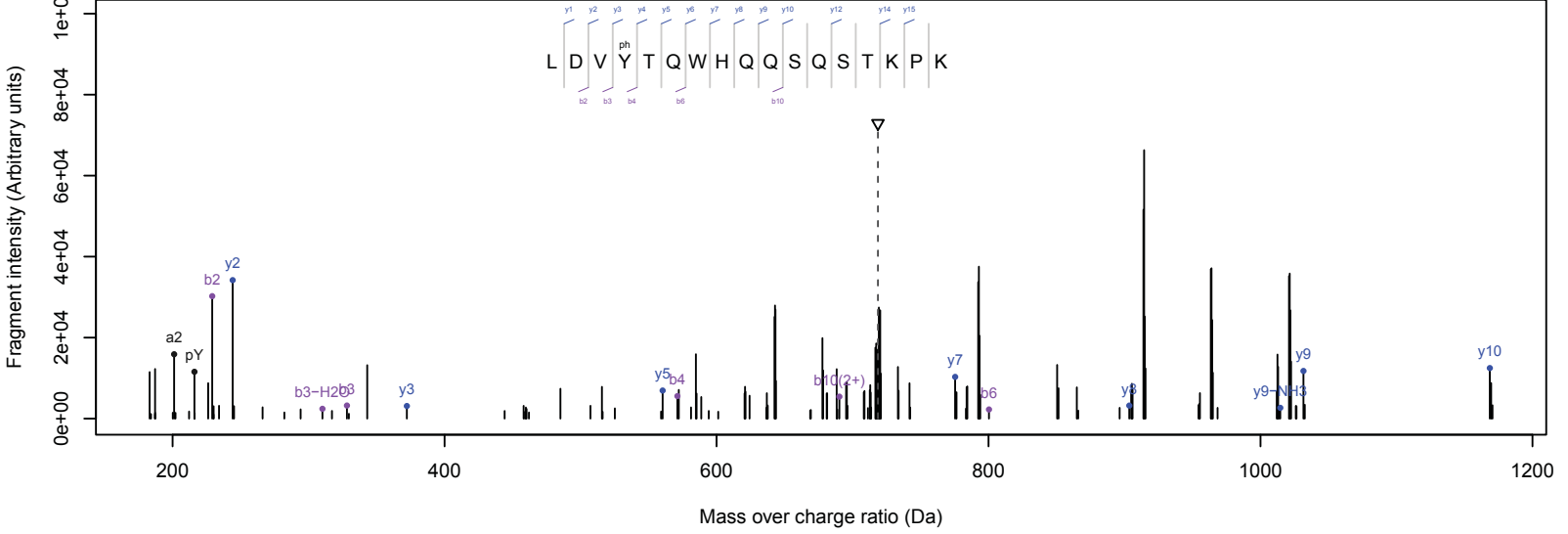
Andromeda score : 118.31



LDVY(ph)TQWHQQSQSTKPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16327

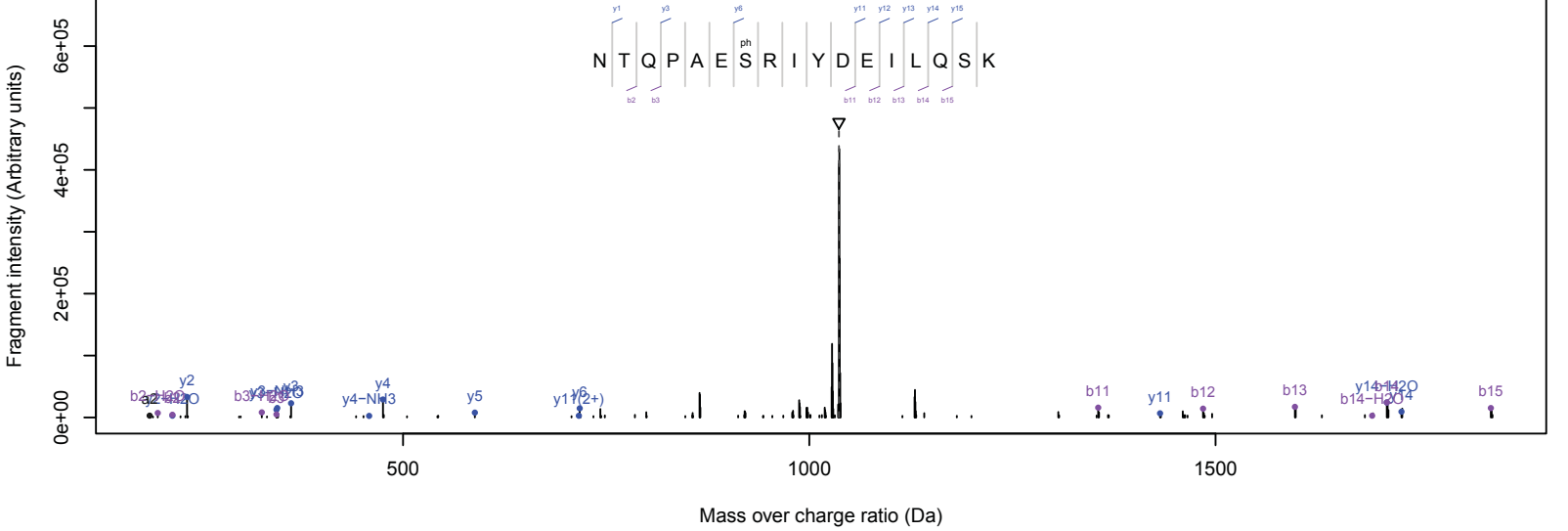
Andromeda score : 118.31



NTQPAES(ph)RIYDEILQSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26804

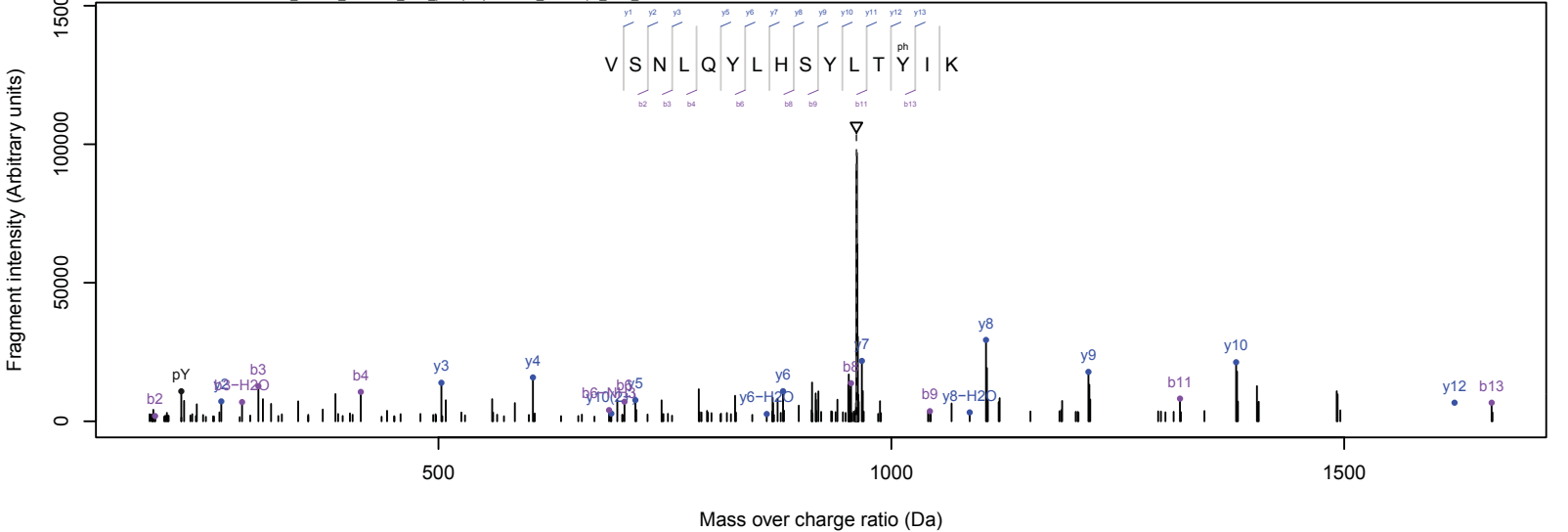
Andromeda score : 118.31



VSNLQYLHSYLT(ph)IK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39409

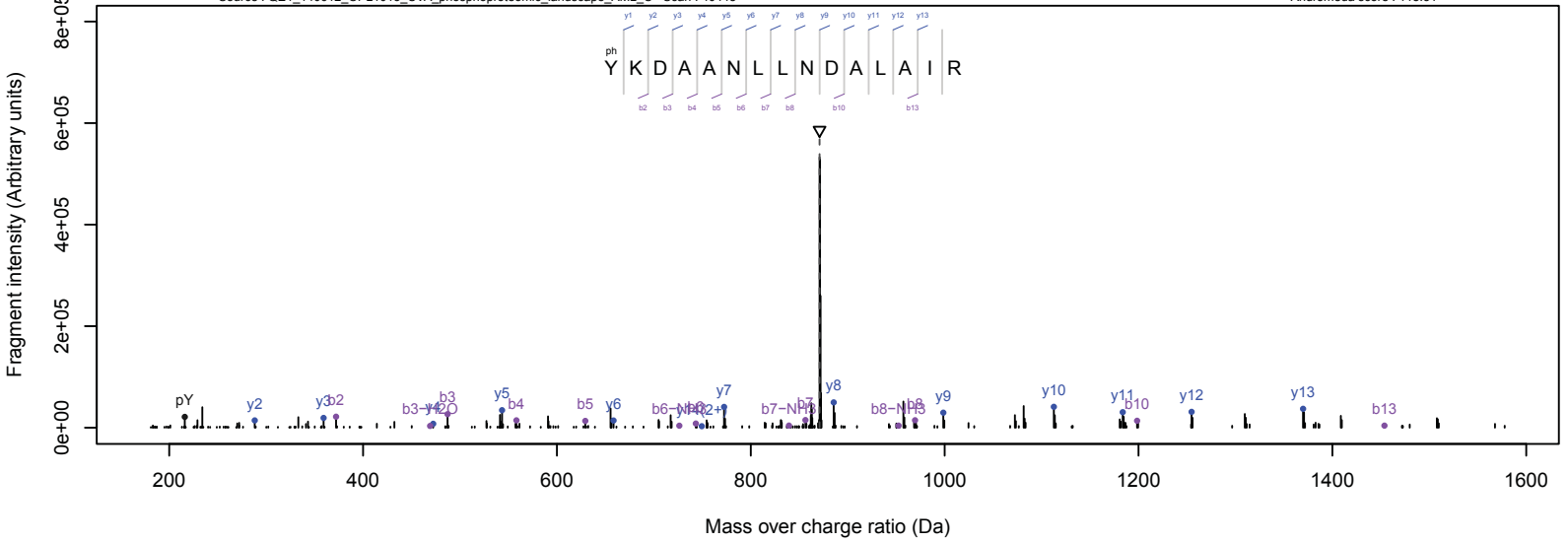
Andromeda score : 118.31



Y(ph)KDAANLLNDALAIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40448

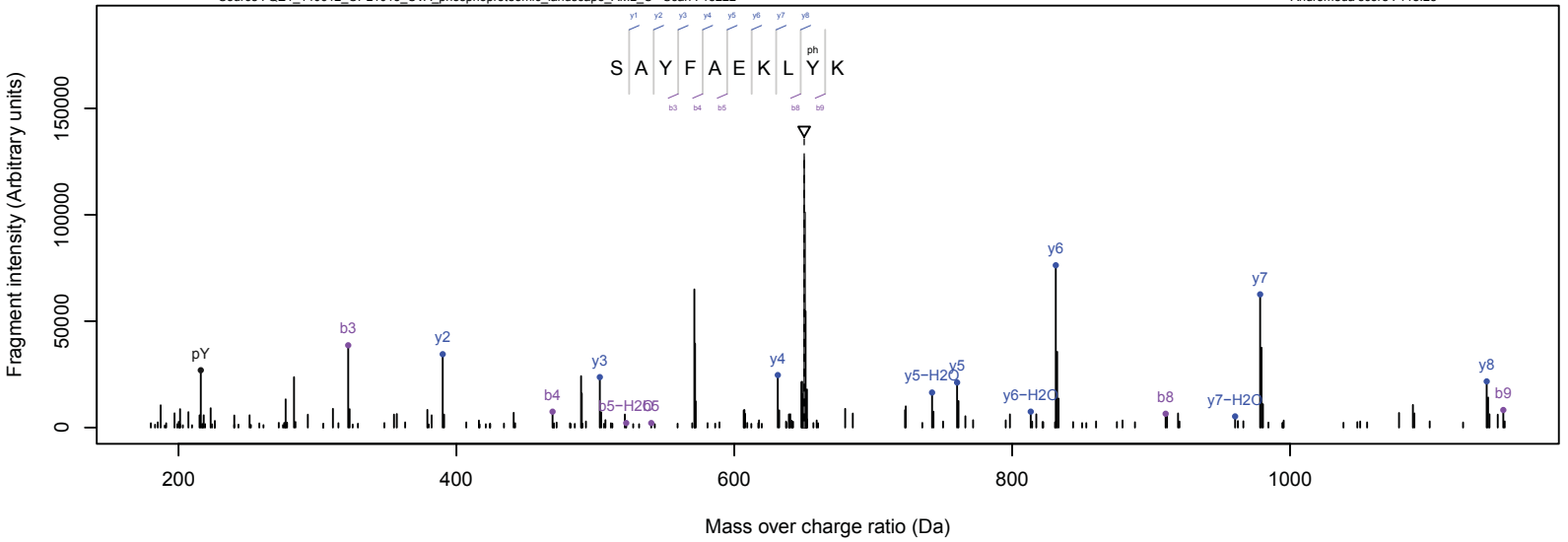
Andromeda score : 118.31



SAYFAEKLY(ph)K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18222

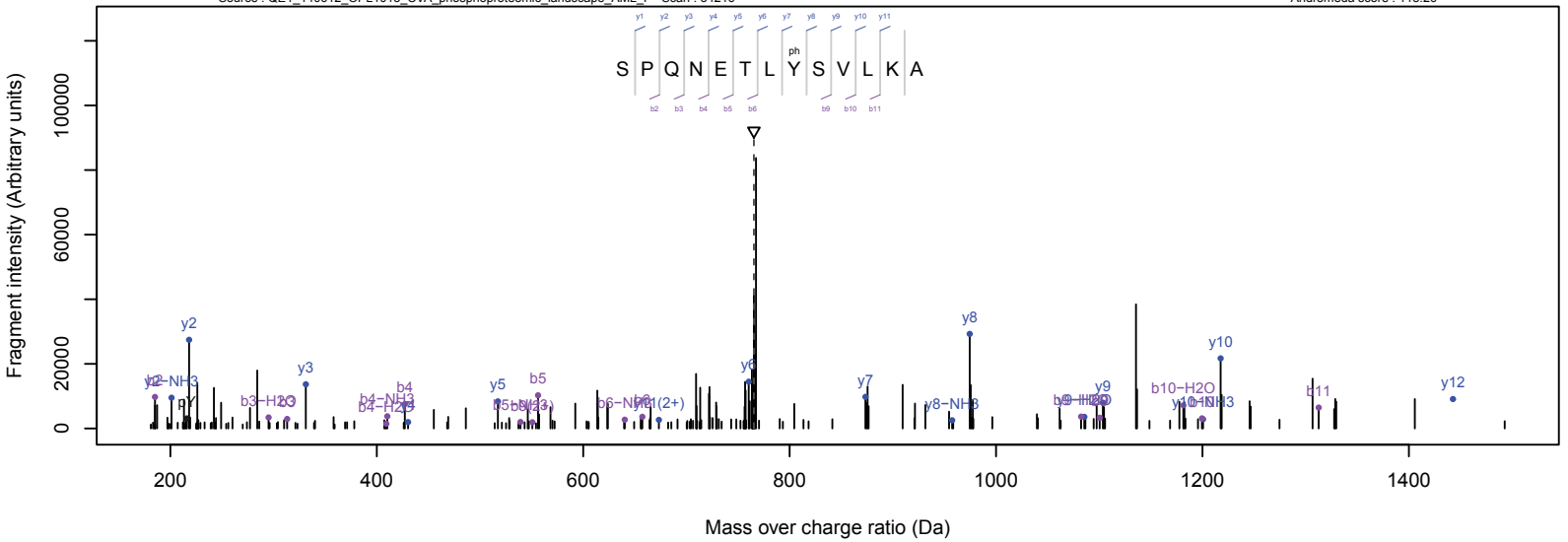
Andromeda score : 118.28



SPQNETLY(ph)SVLKA-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 34216

Andromeda score : 118.26





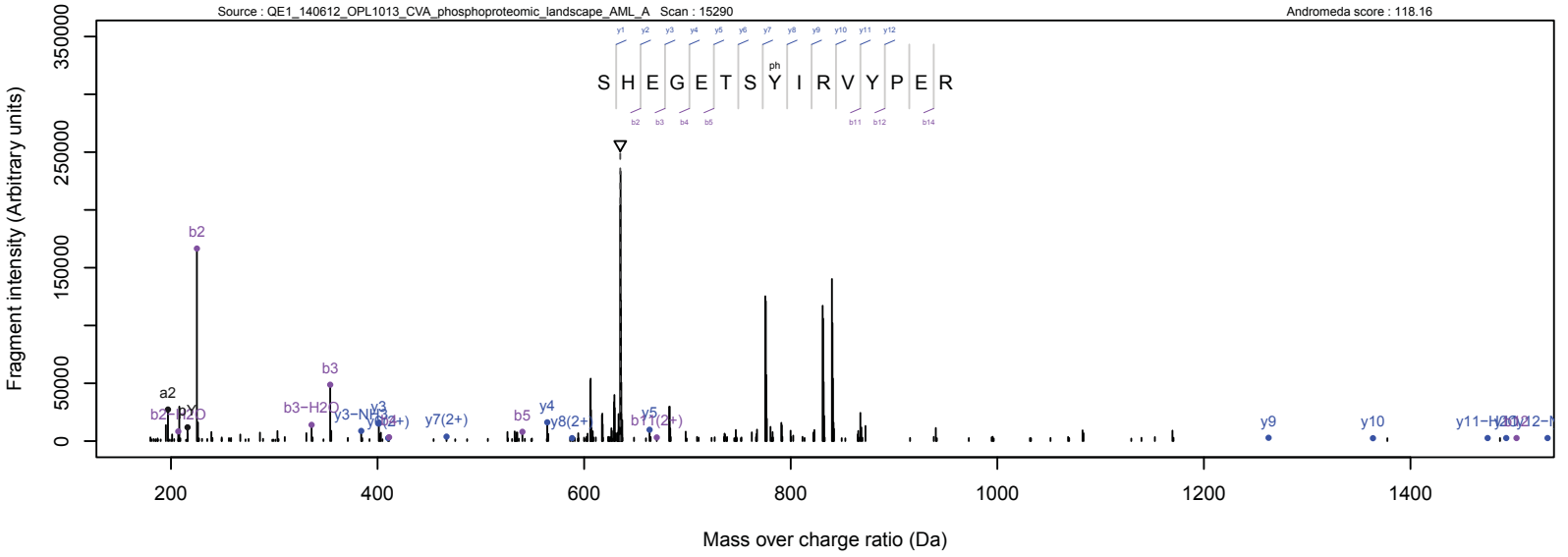




**SHEGETSY(ph)IRVYPER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 15290

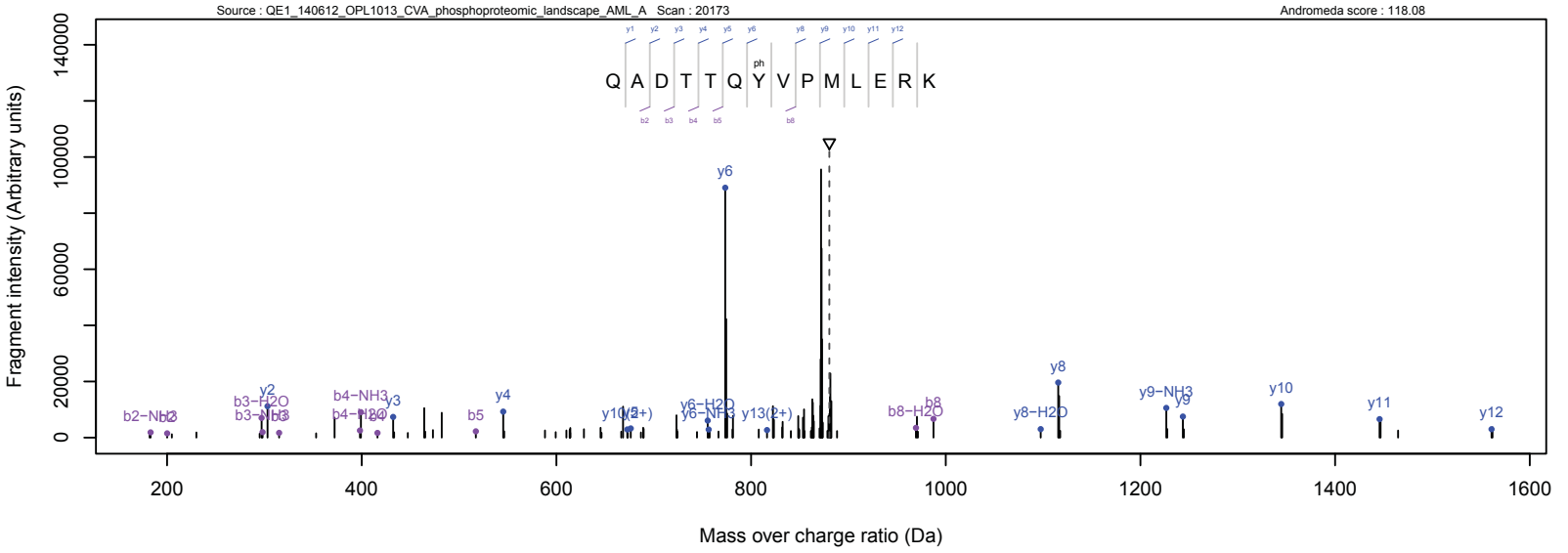
Andromeda score : 118.16



**QADTTQY(ph)VPMLERK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 20173

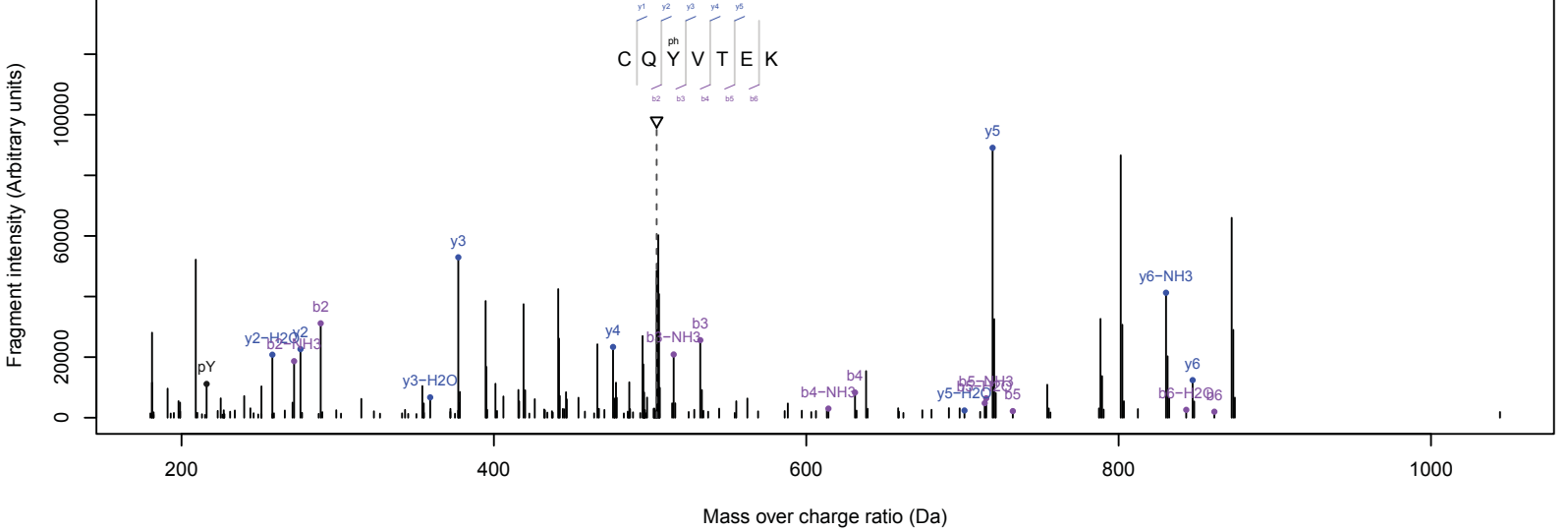
Andromeda score : 118.08



**CQY(ph)VTEK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 2899

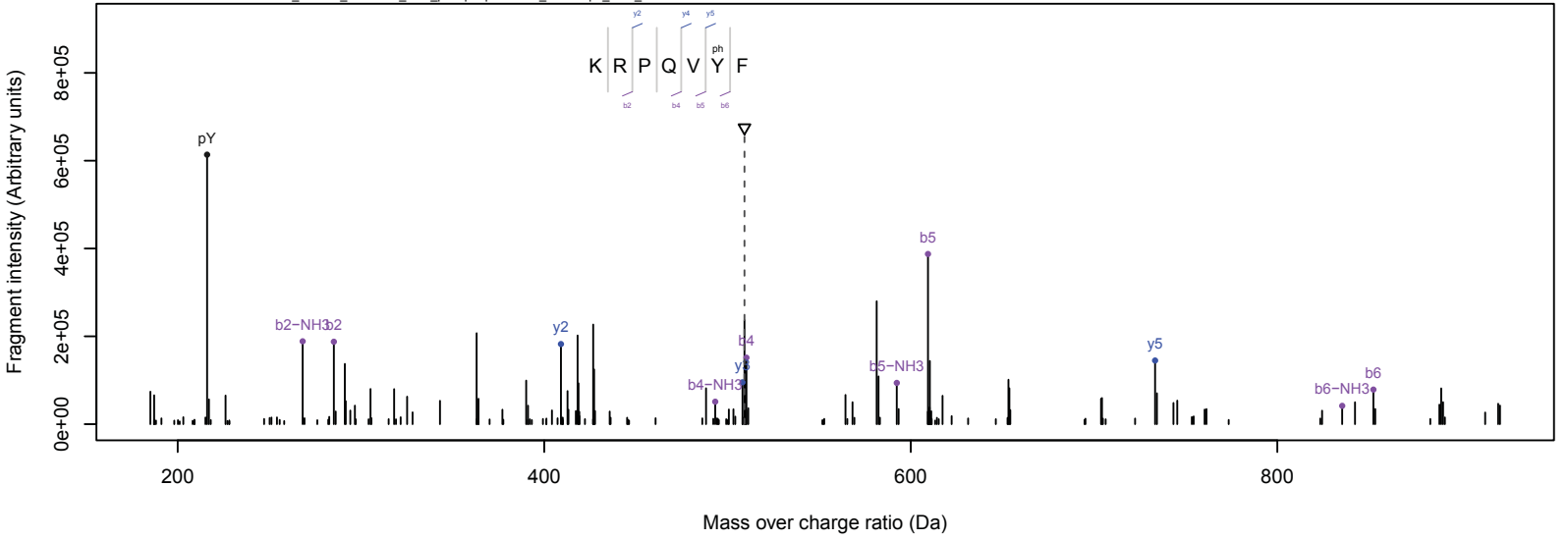
Andromeda score : 118.07



KRPQVY(ph)F-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14480

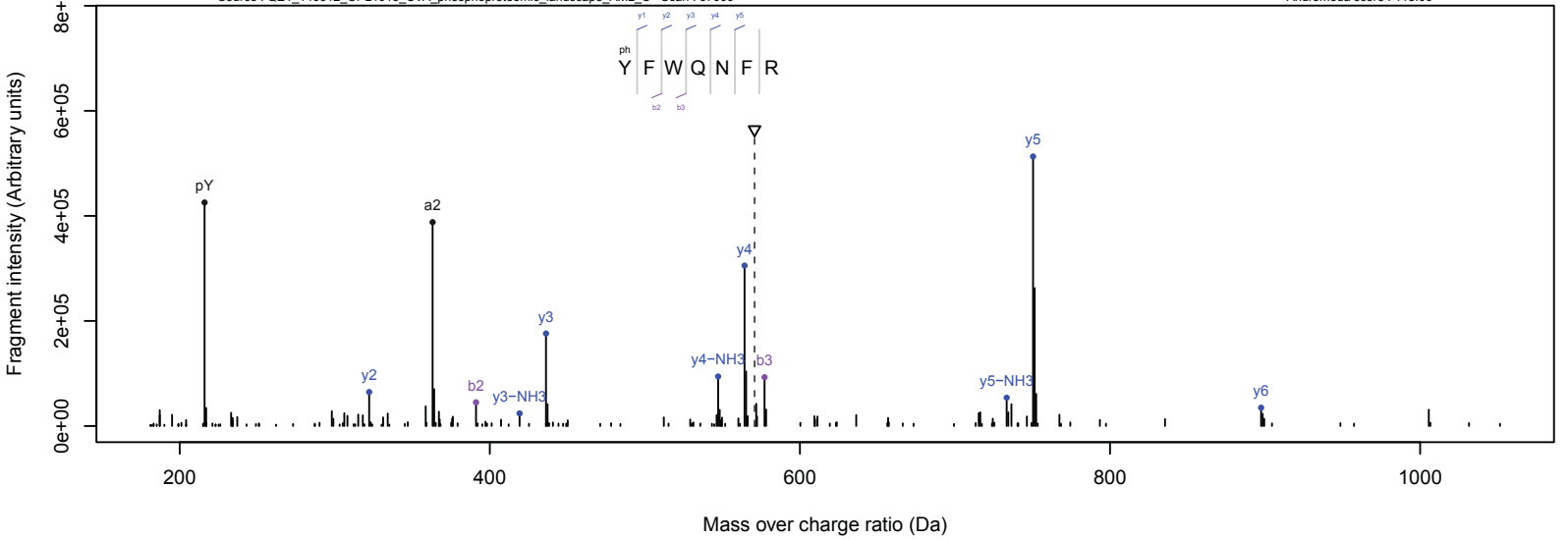
Andromeda score: 118.06



Y(ph)FWQNFR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37009

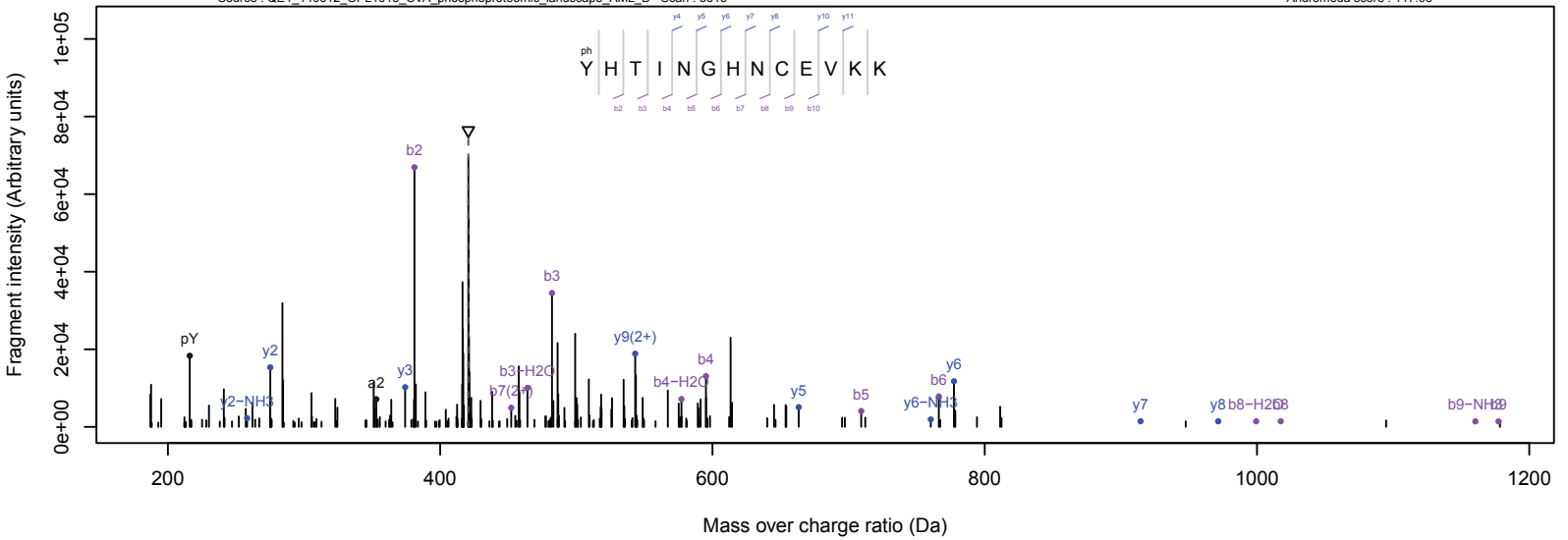
Andromeda score: 118.06



Y(ph)HTINGHNCEVKK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 3616

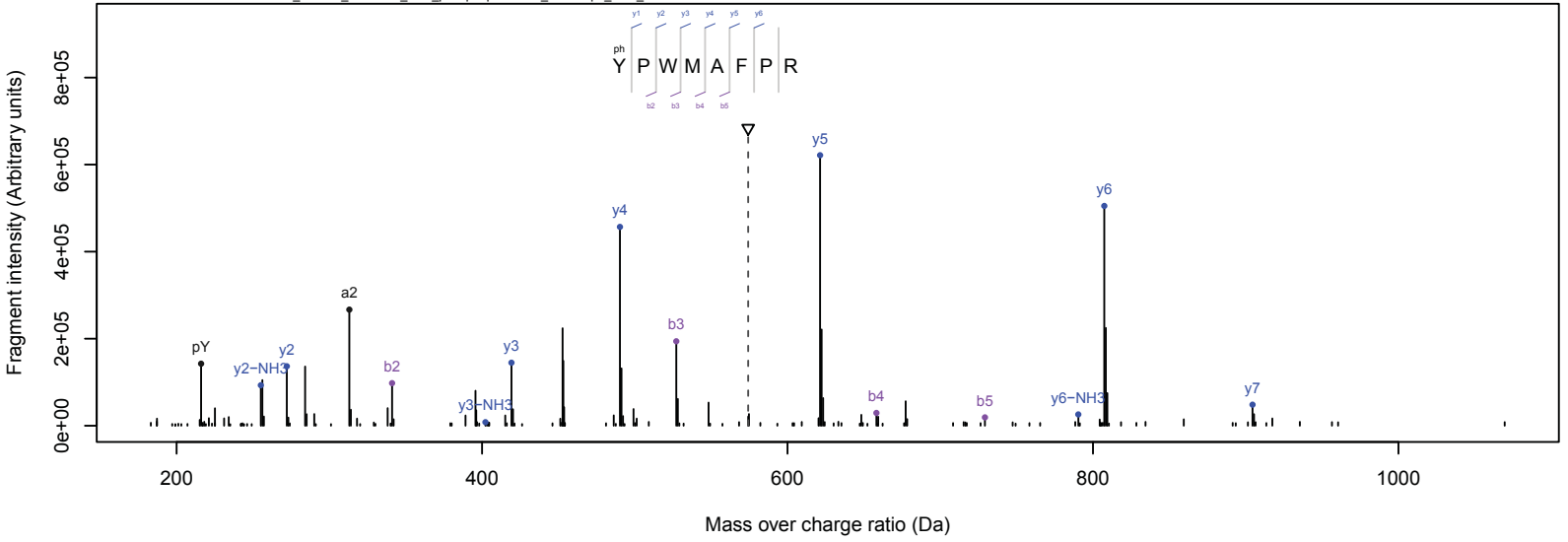
Andromeda score: 117.93



Y(ph)PWMAFPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37395

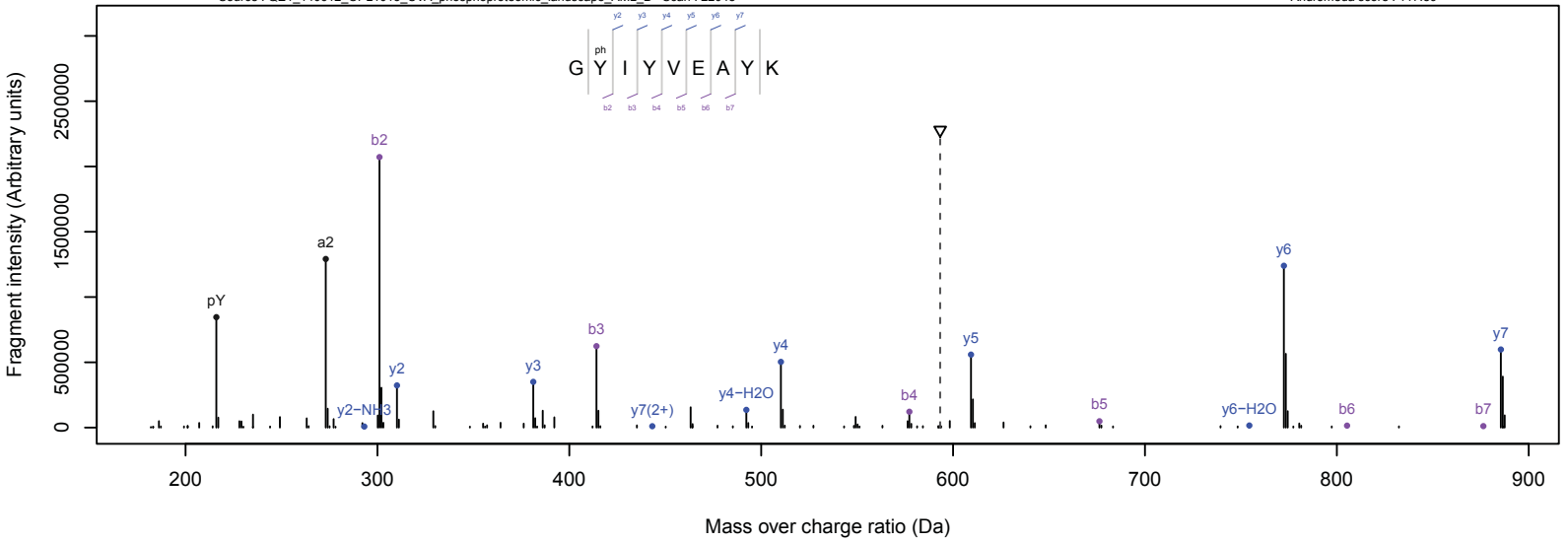
Andromeda score : 117.93



GY(ph)IYVEAYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22048

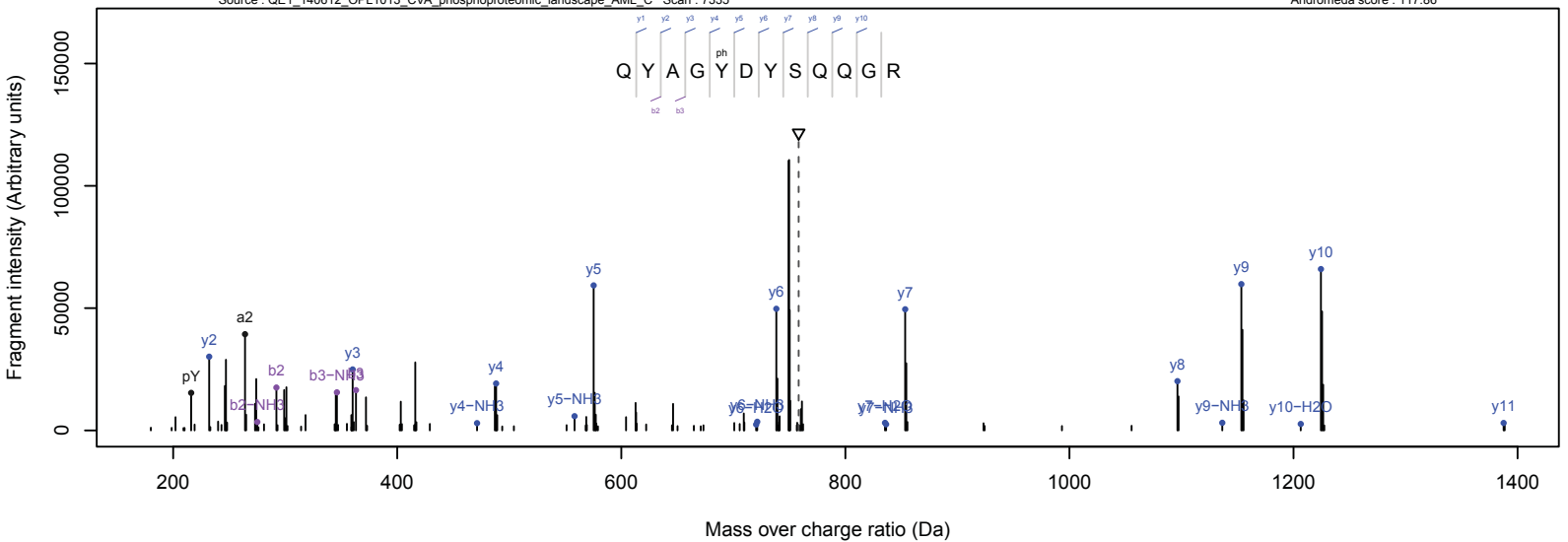
Andromeda score : 117.89



QYAGYDYSQQGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7335

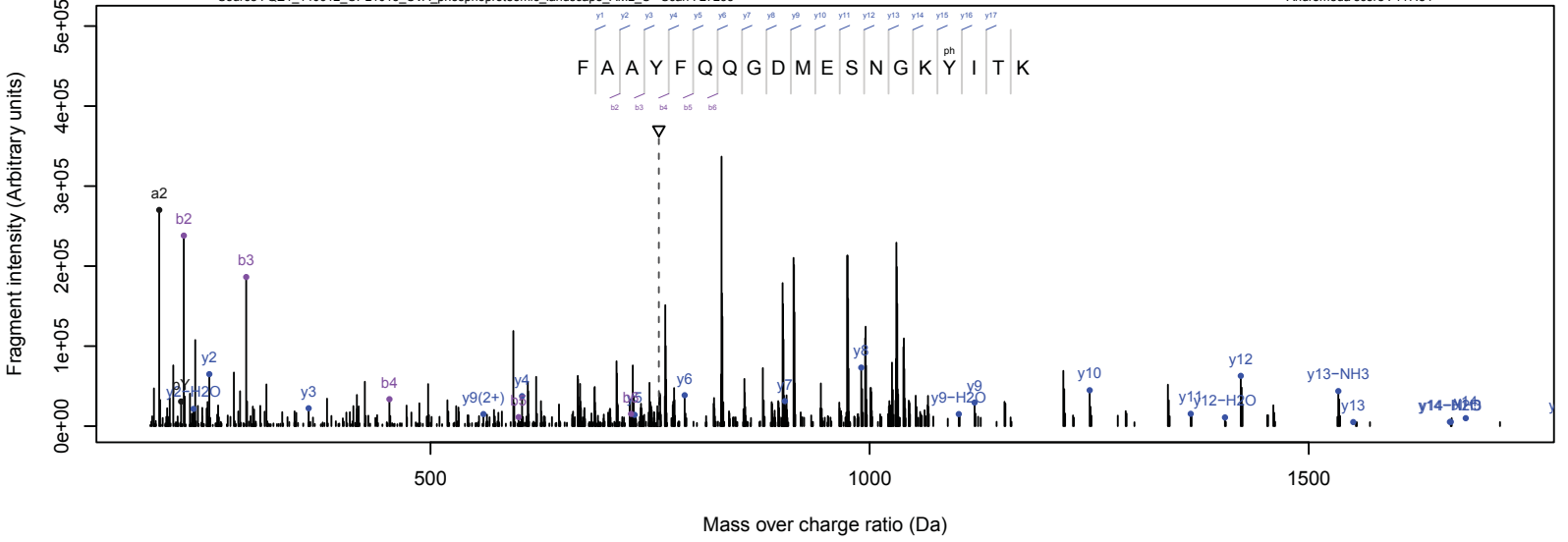
Andromeda score : 117.86



FAAYFQQGDMESNGKY(ph)ITK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27233

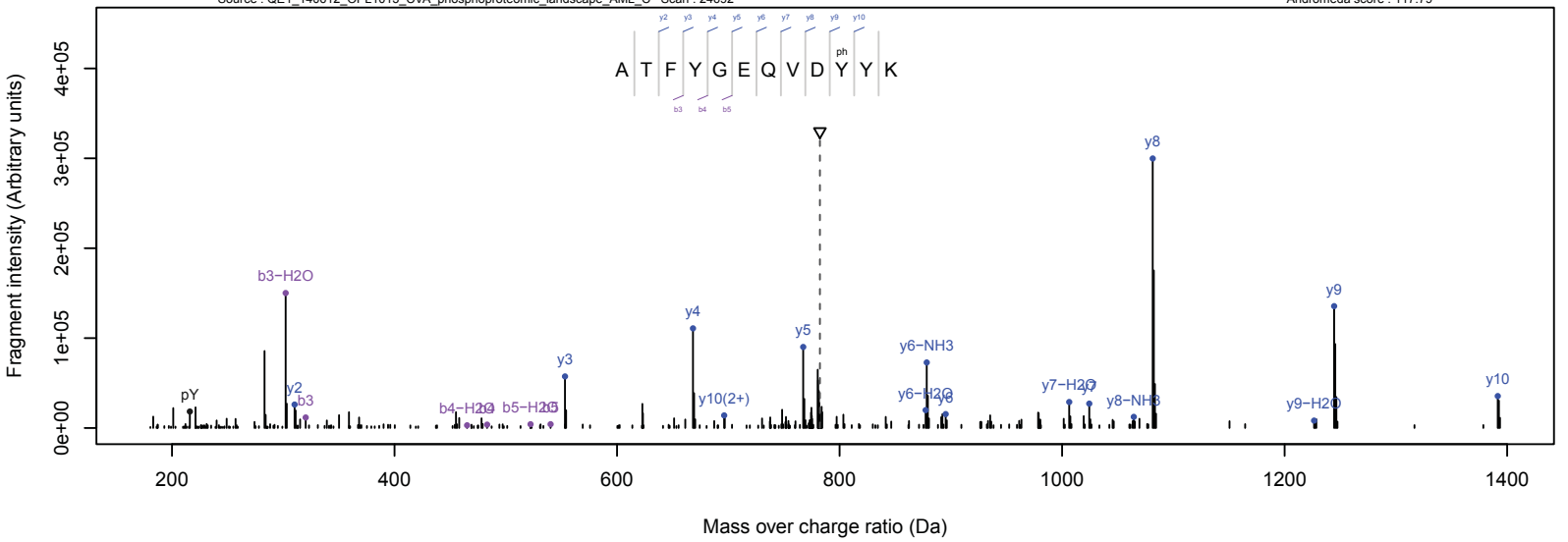
Andromeda score : 117.84



ATFYGEQVDY(ph)YK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24692

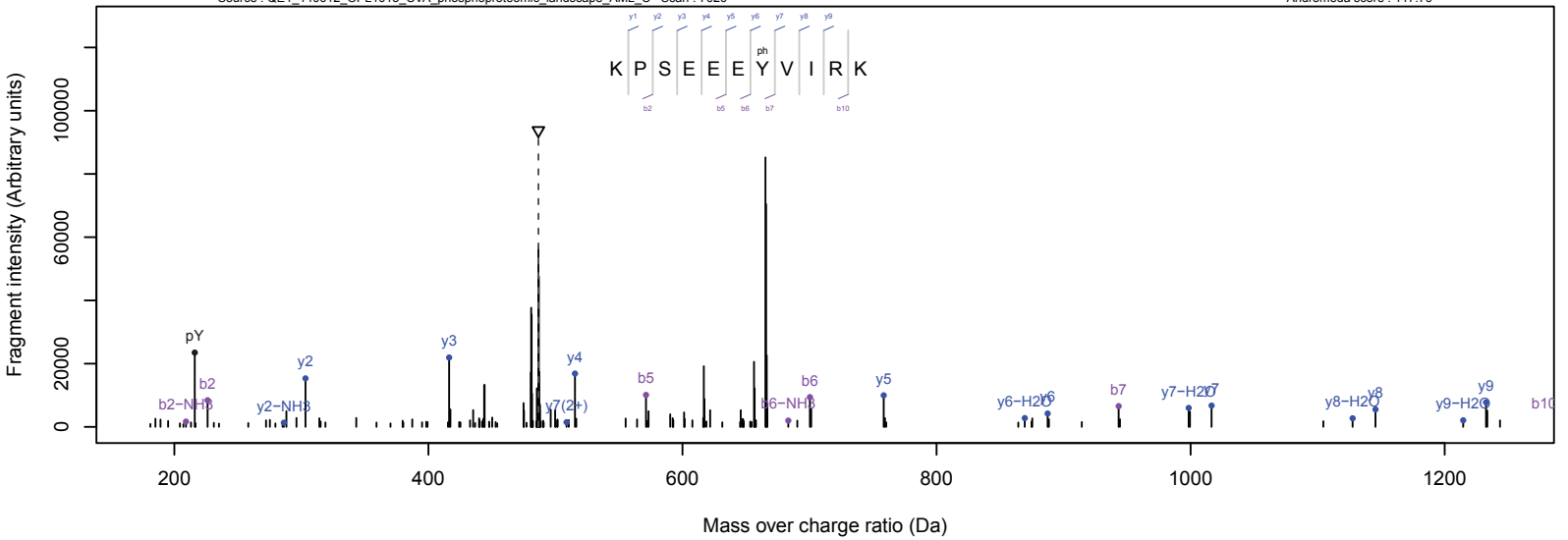
Andromeda score : 117.79



KPSEEEYVIRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7026

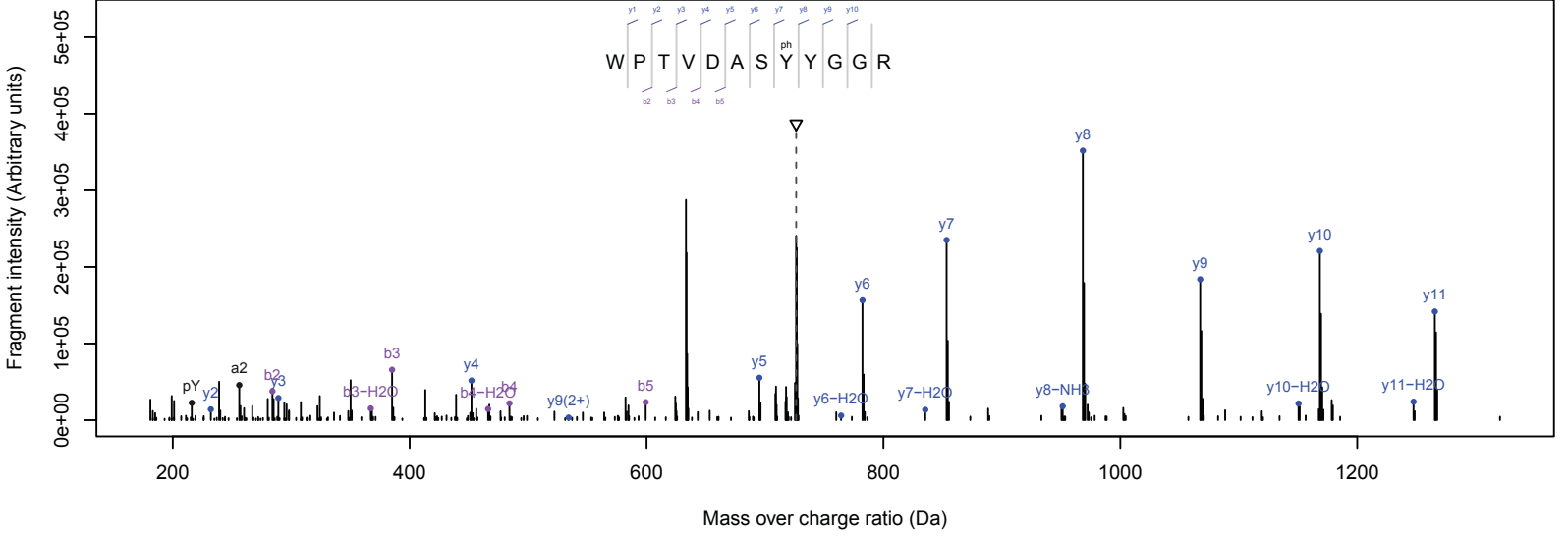
Andromeda score : 117.79



WPTVDASY(ph)YGGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28272

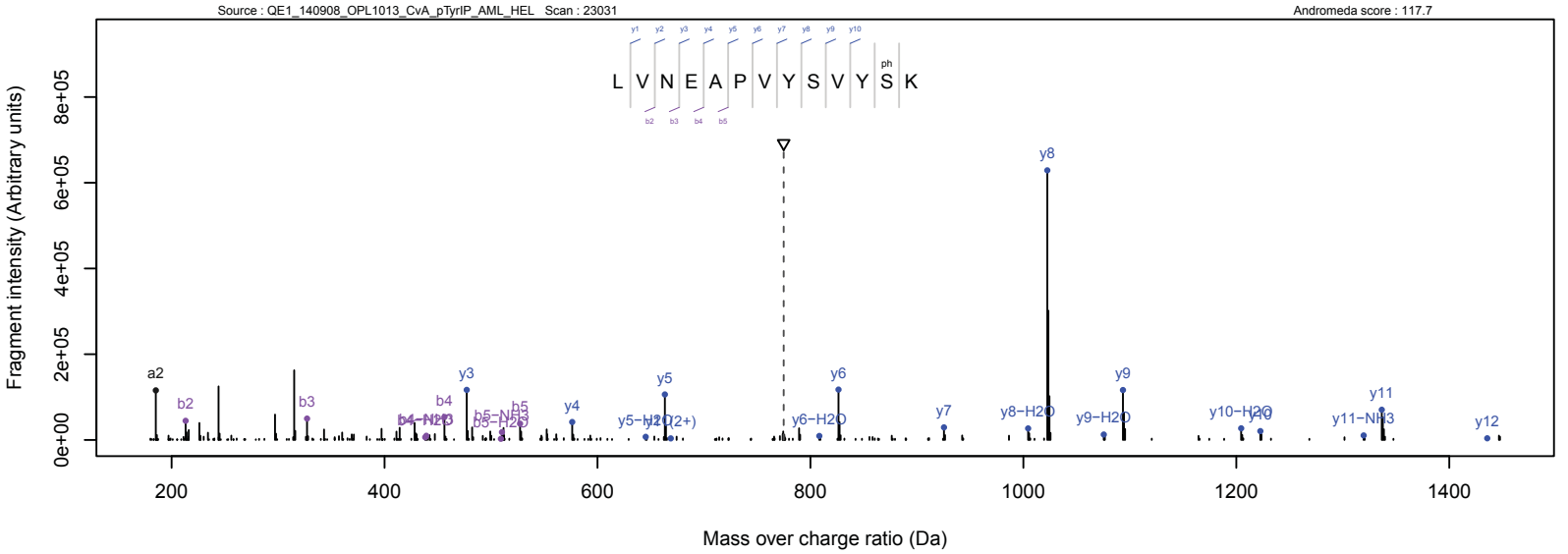
Andromeda score : 117.79



LVNEAPVYSVYS(ph)K-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23031

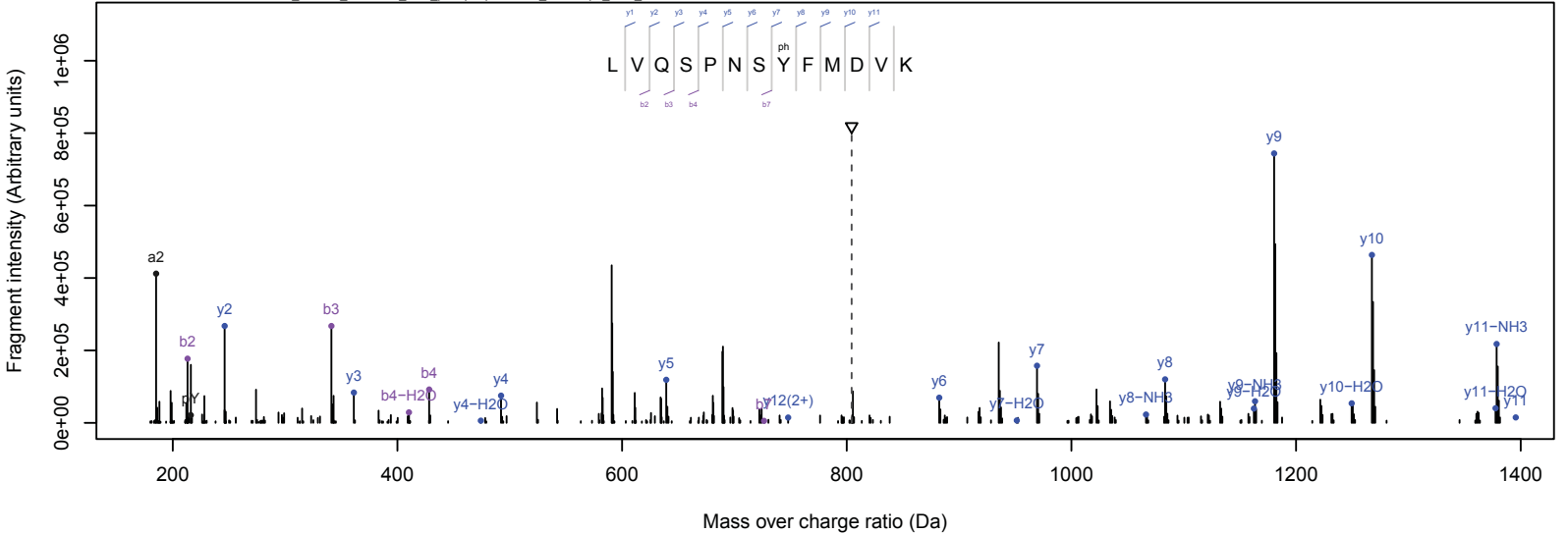
Andromeda score : 117.7



LVQSPNSY(ph)FMDVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29377

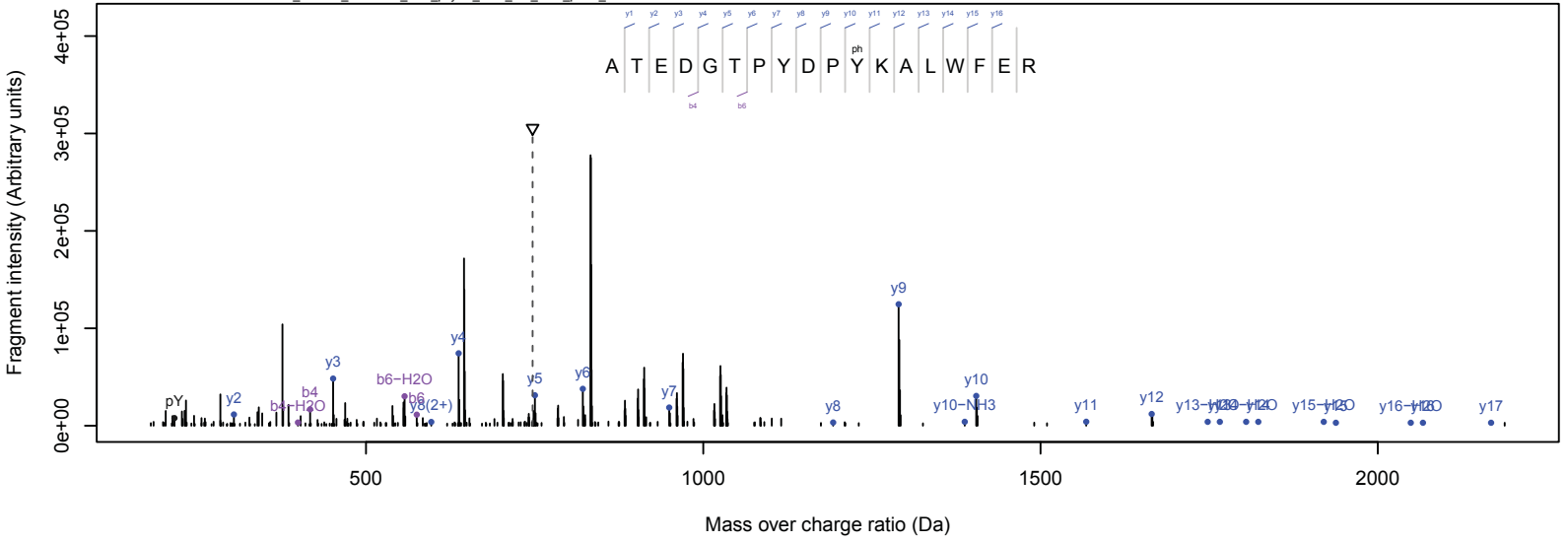
Andromeda score : 117.7



ATEDGTPYDPY(ph)KALWFER-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 29046

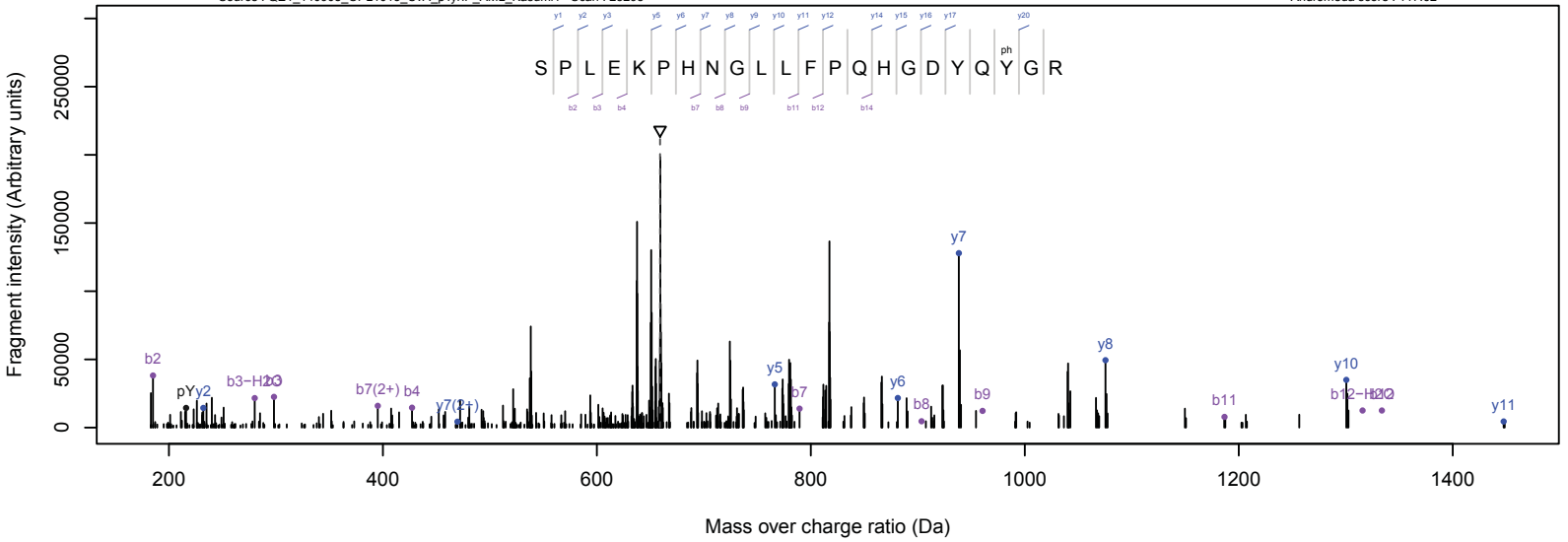
Andromeda score : 117.65



SPLEKPHNGLLPQHG DYQY(ph)GR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 23295

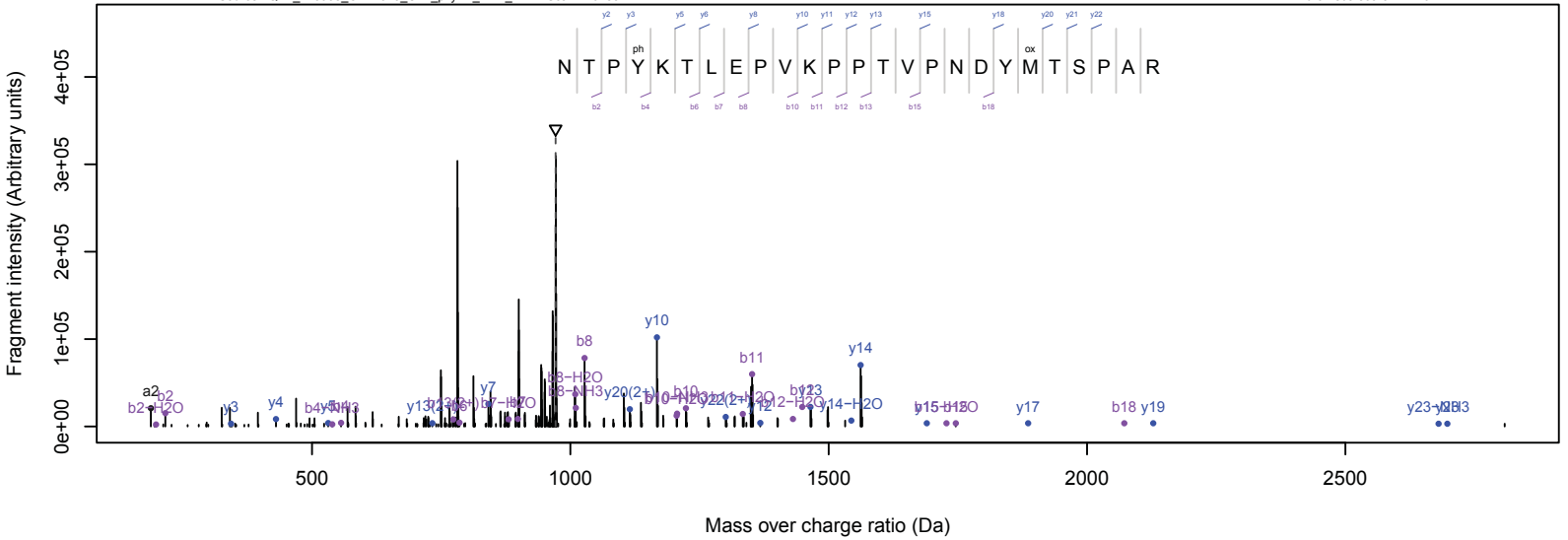
Andromeda score : 117.62



NTPY(ph)KTLPEVKPPTVPNDYMTSPAR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 26285

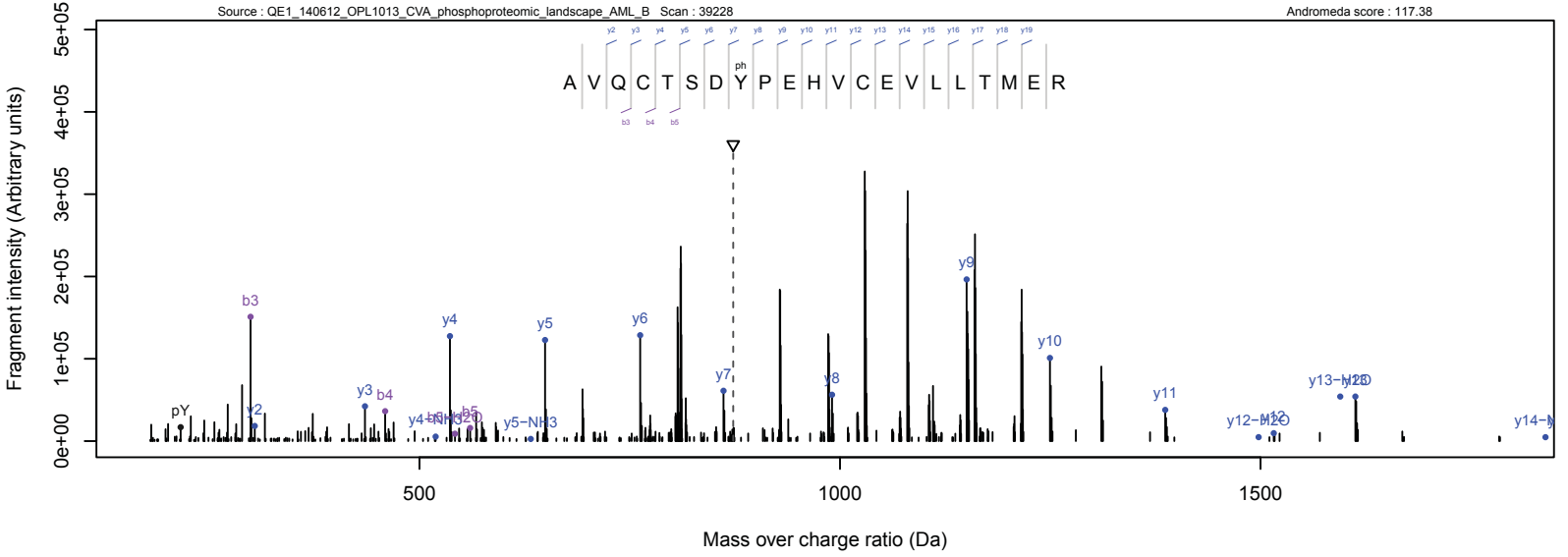
Andromeda score : 117.61



AVQCTSDY(ph)PEHVCEVLLTMER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39228

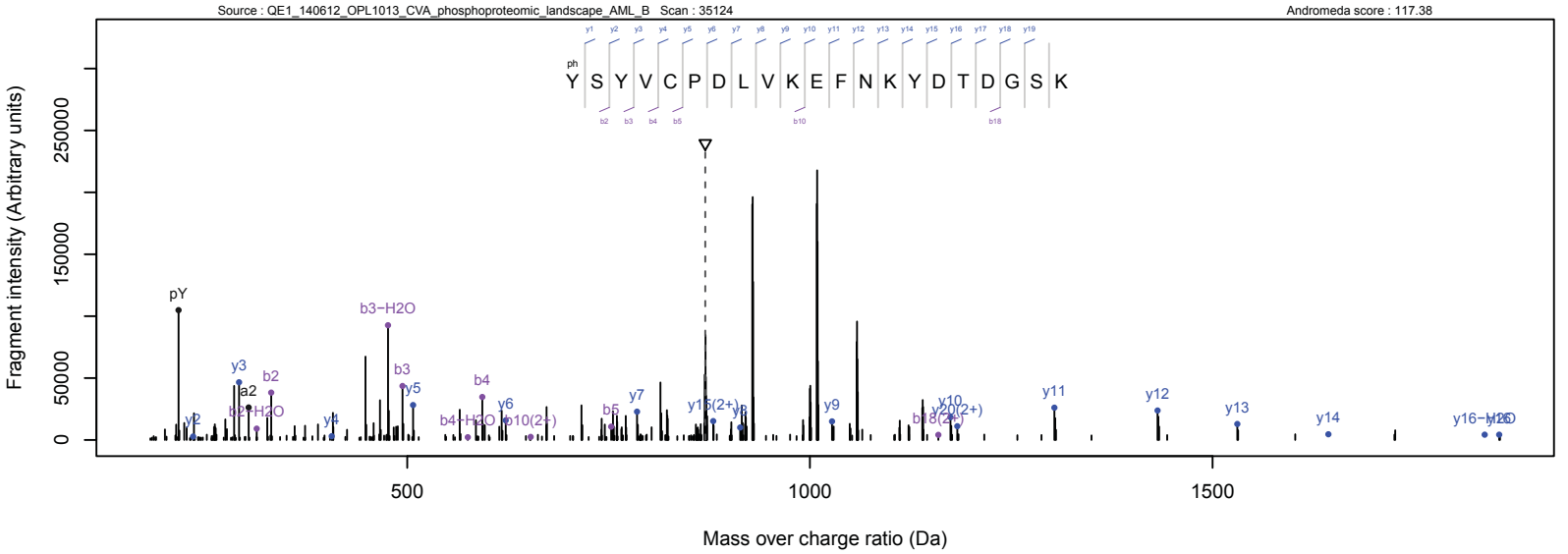
Andromeda score : 117.38



Y(ph)SYVCPDLVKEFNKYDTDGSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35124

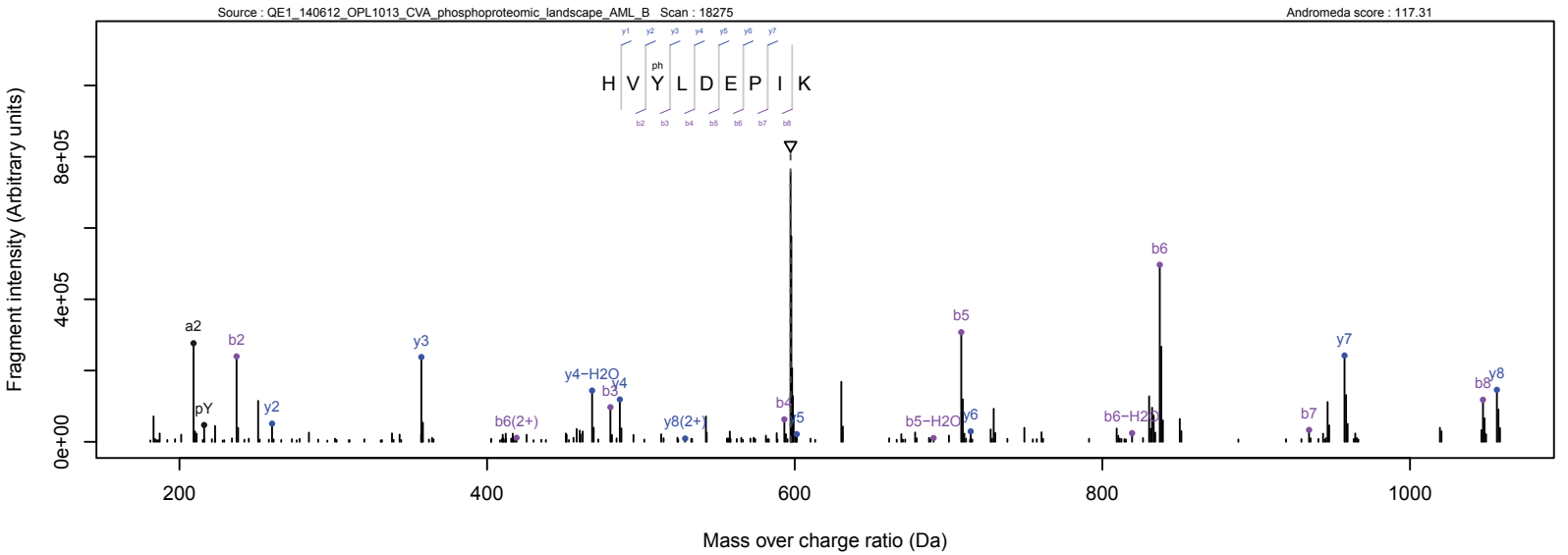
Andromeda score : 117.38



HVY(ph)LDEPIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18275

Andromeda score : 117.31

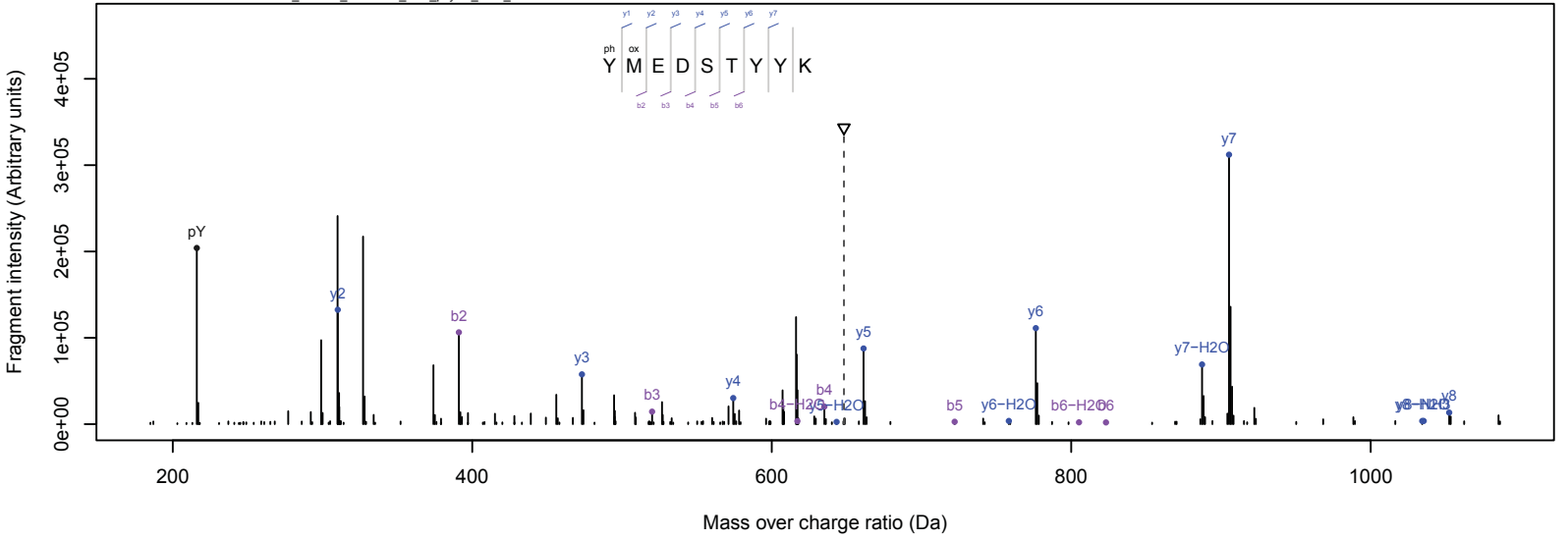




Y(ph)M(ox)EDSTYYK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 7304

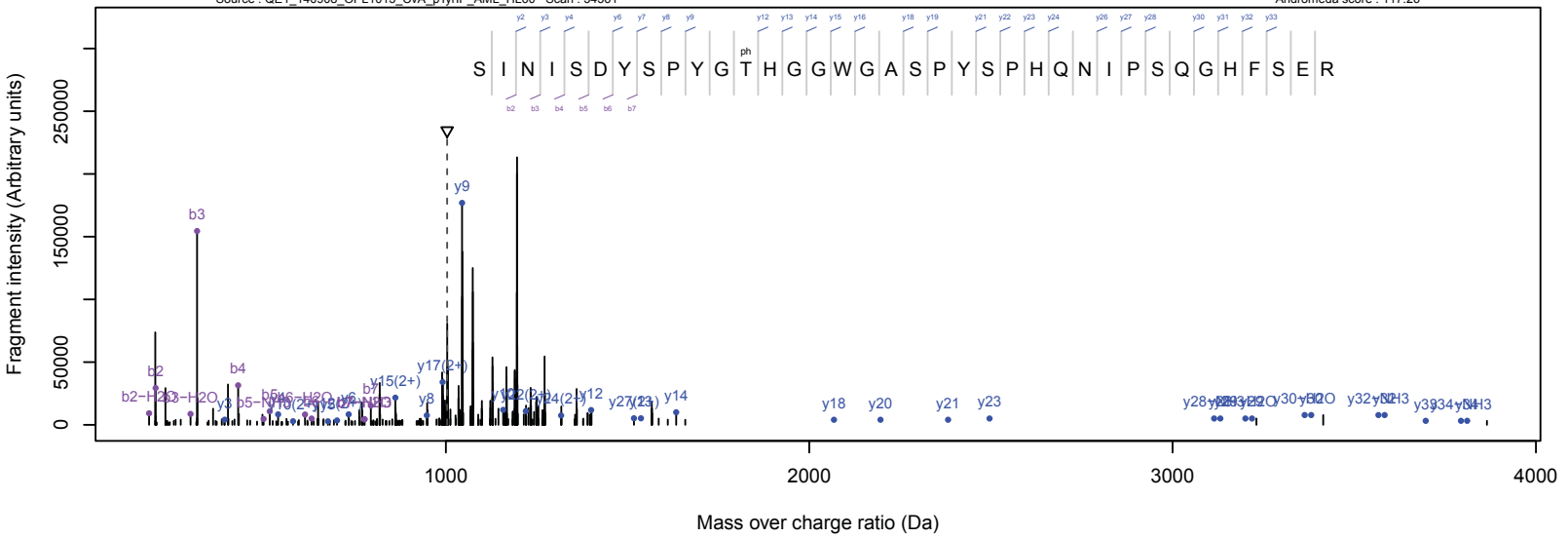
Andromeda score : 117.31



SINISDYSPYGT(ph)HGGWGWASPYSPHQNIP SQGHFSEK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 34361

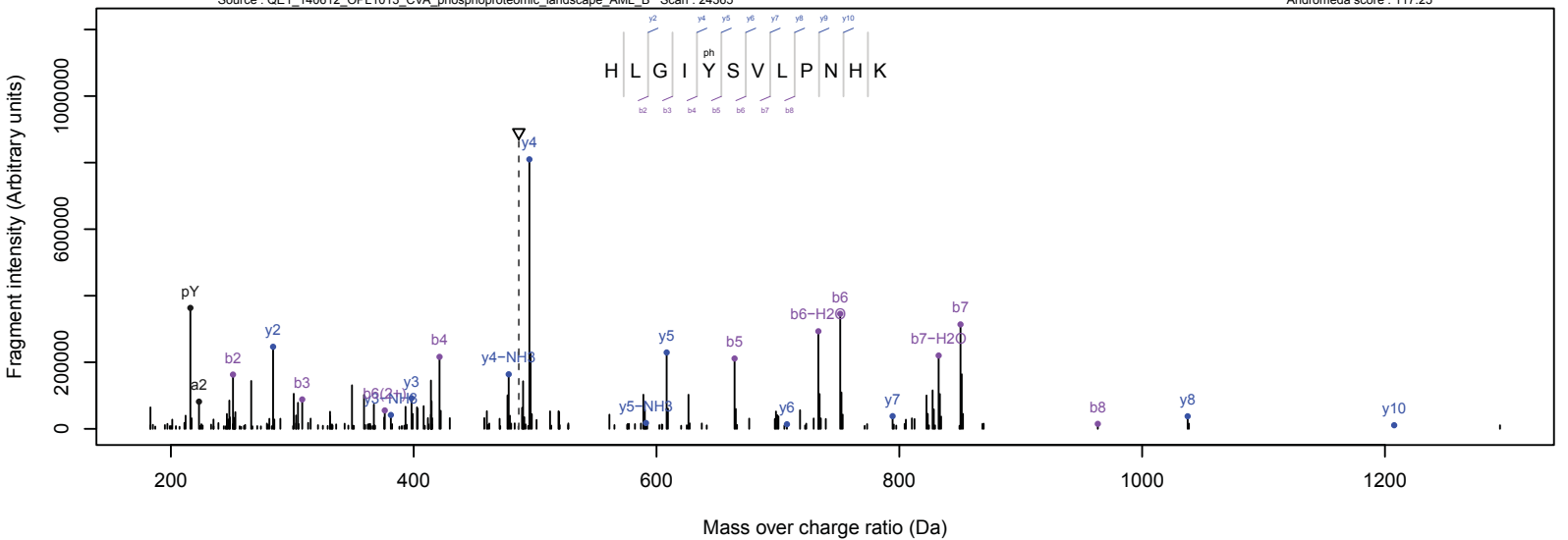
Andromeda score : 117.26



HLGIY(ph)SVLPNHK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan : 24365

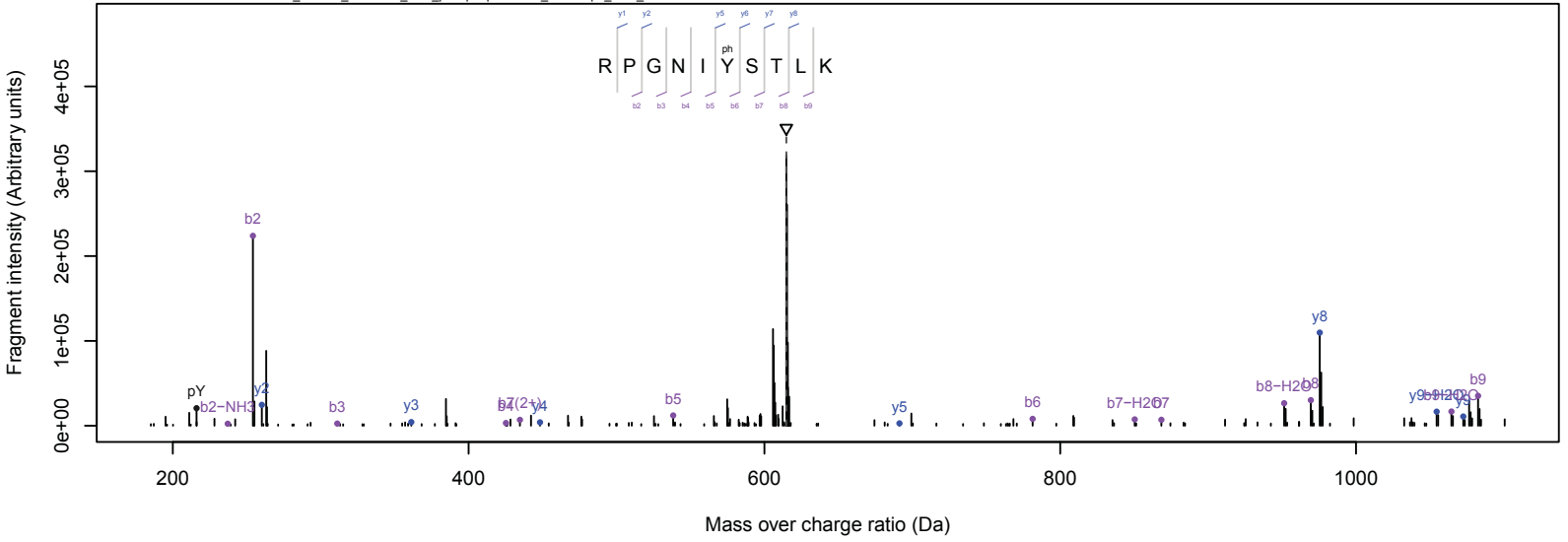
Andromeda score : 117.25



RPGNIY(ph)STLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10078

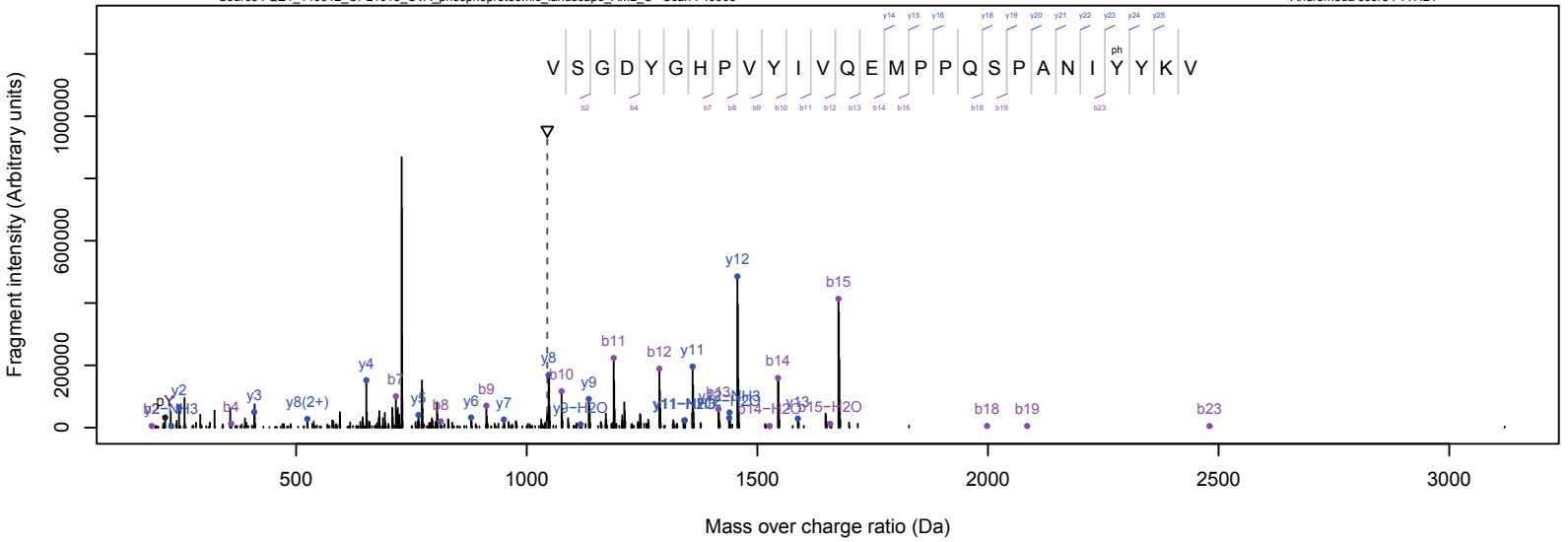
Andromeda score : 117.25



VSGDYGHPVYIQEMPQPSPANIY(ph)YKV-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40905

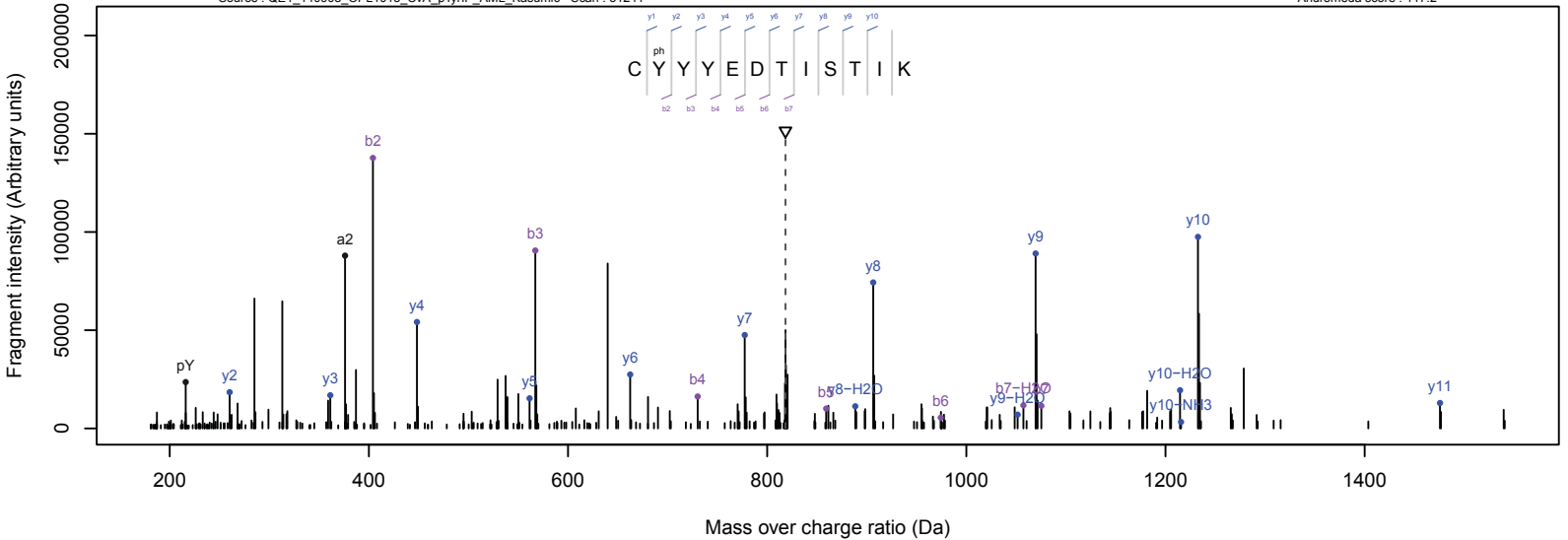
Andromeda score : 117.21



CY(ph)YYEDTISTIK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 31241

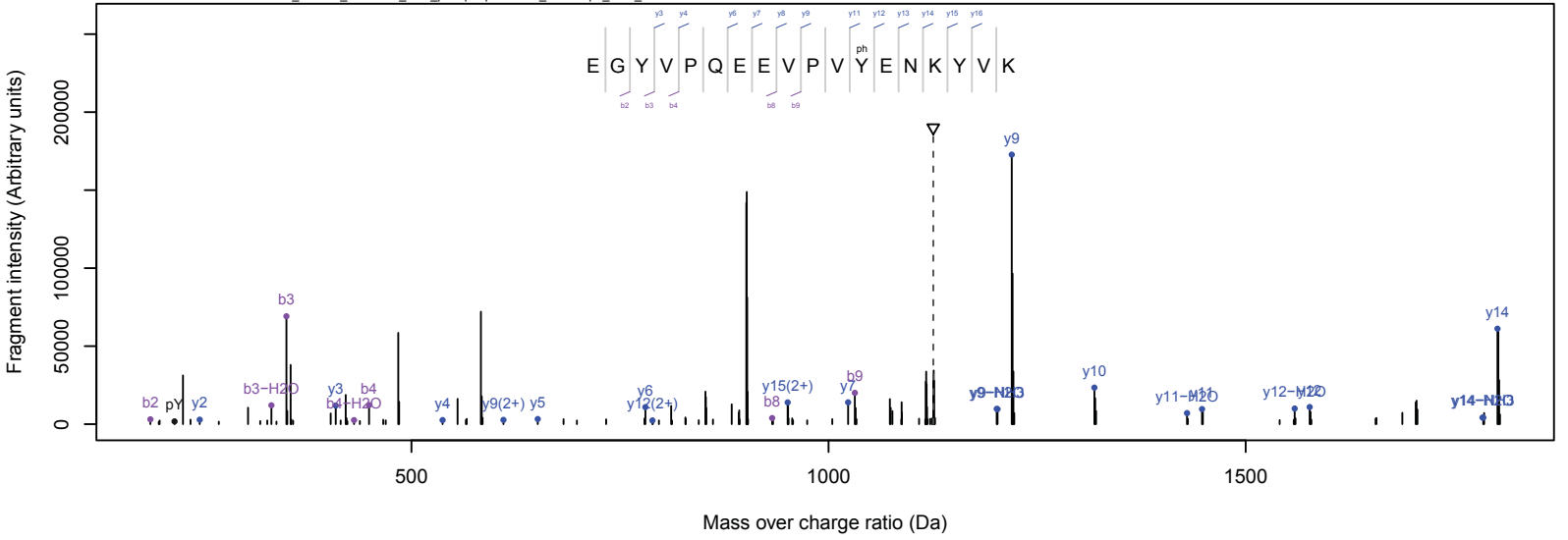
Andromeda score : 117.2



EGYVPQEEVPV(ph)ENKYVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26406

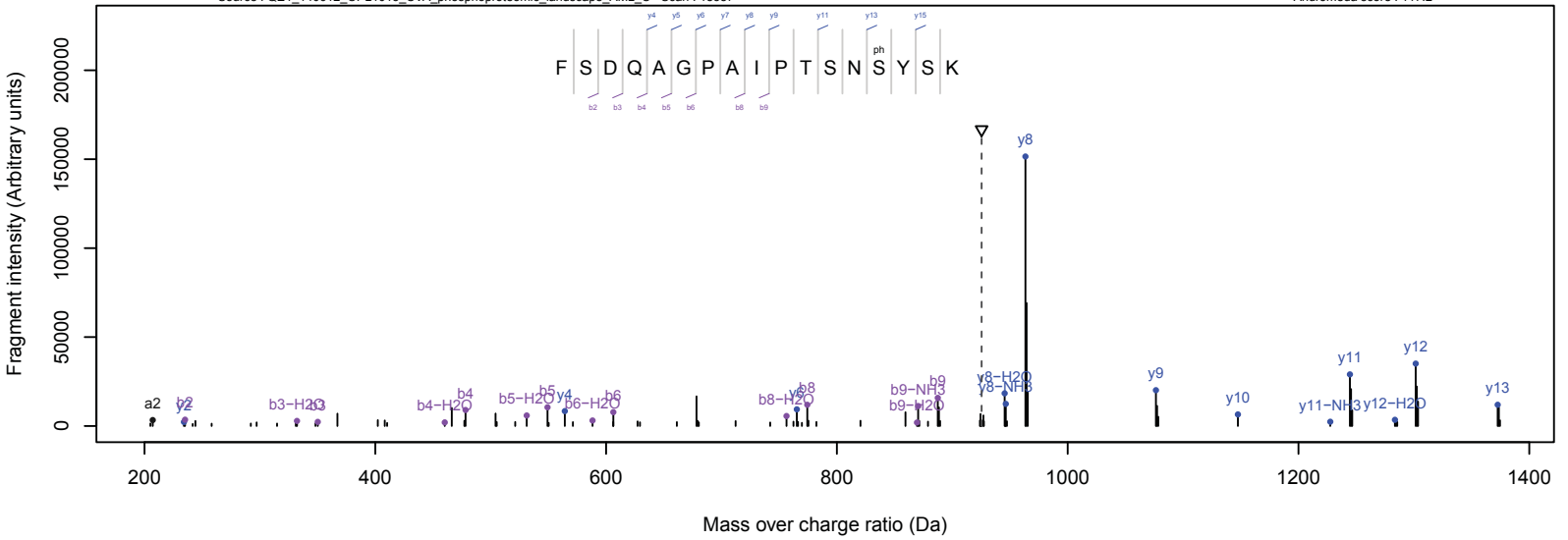
Andromeda score : 117.2



FSDQAGPAITNS(ph)YSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18587

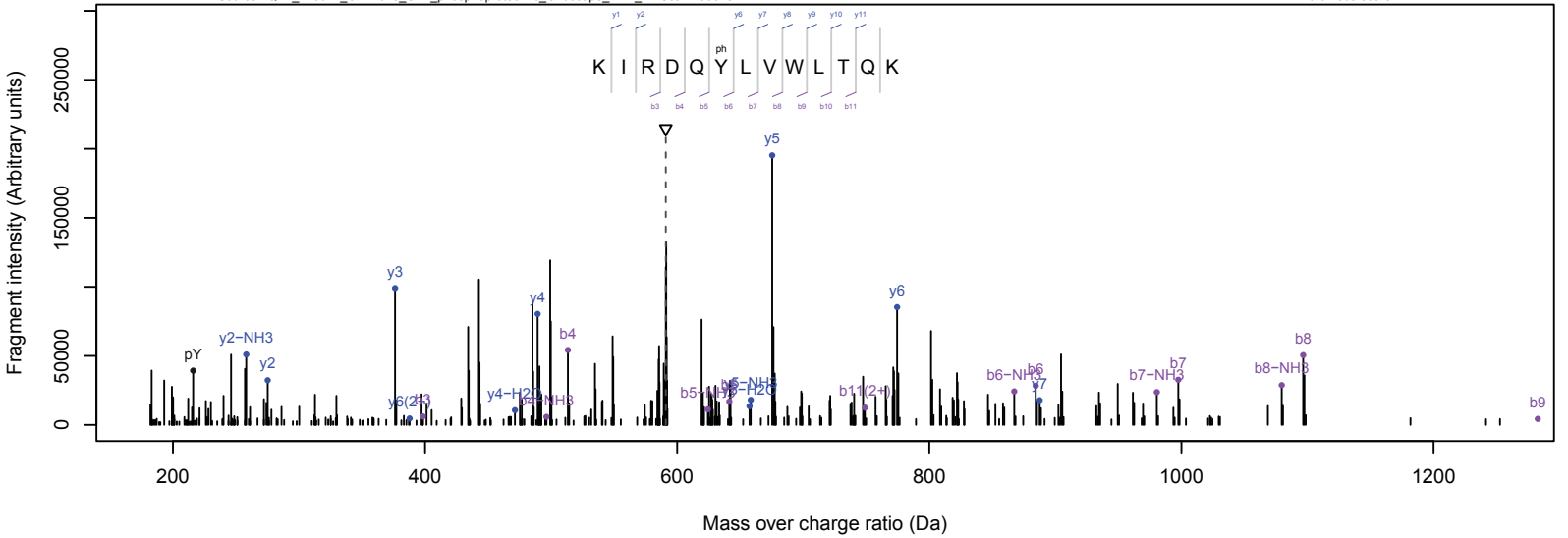
Andromeda score : 117.2



KIRDQY(ph)LVWLTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 30628

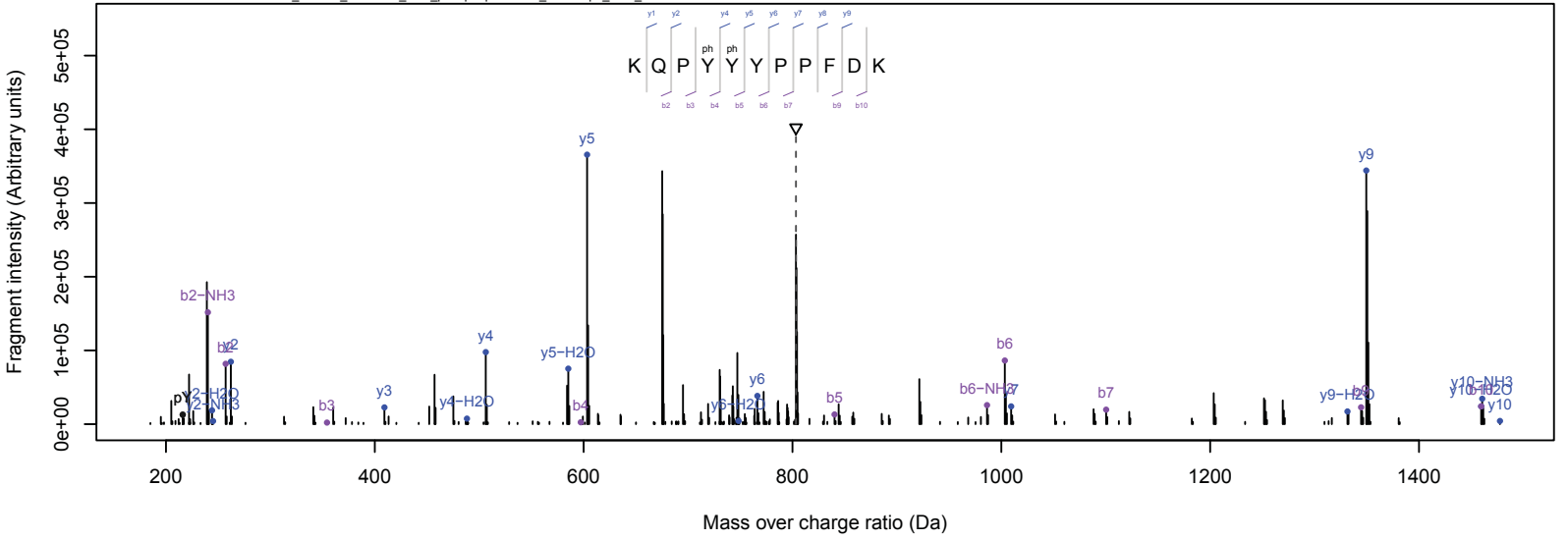
Andromeda score : 117.2



**KQPY(ph)YYPFDK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 20816

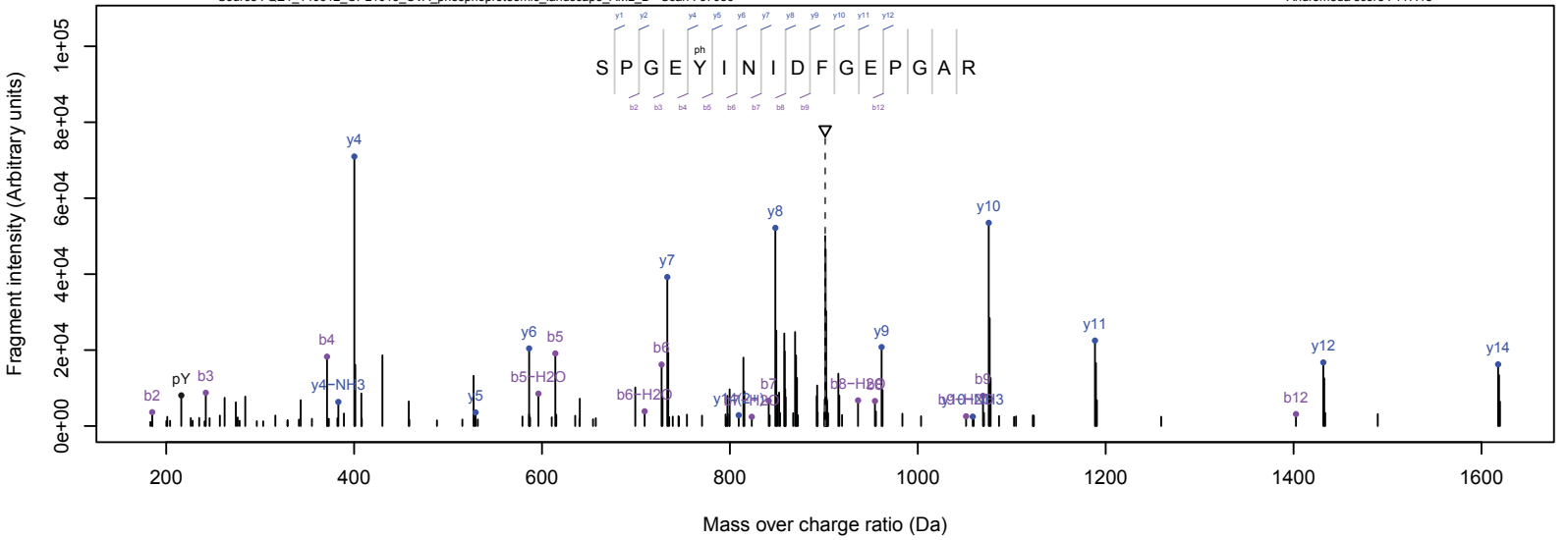
Andromeda score : 117.2



**SPGEY(ph)INIDFGEPGAR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 37086

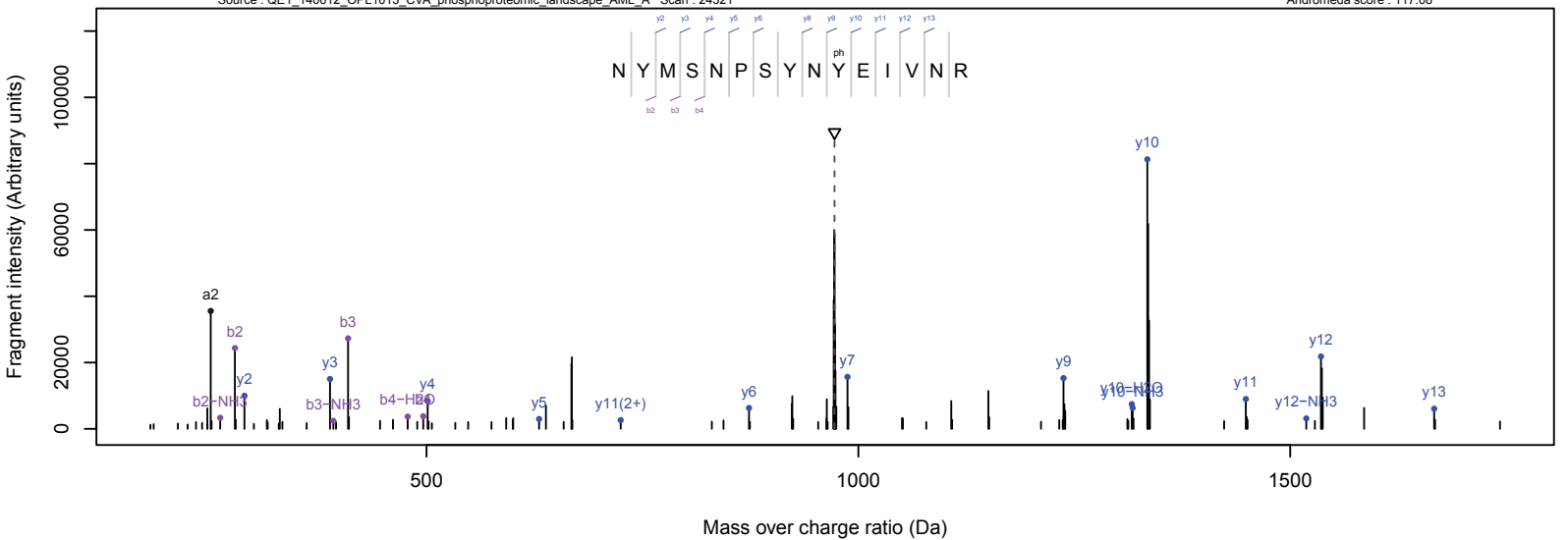
Andromeda score : 117.18



**NYMSNPSYNY(ph)EIVNR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 24321

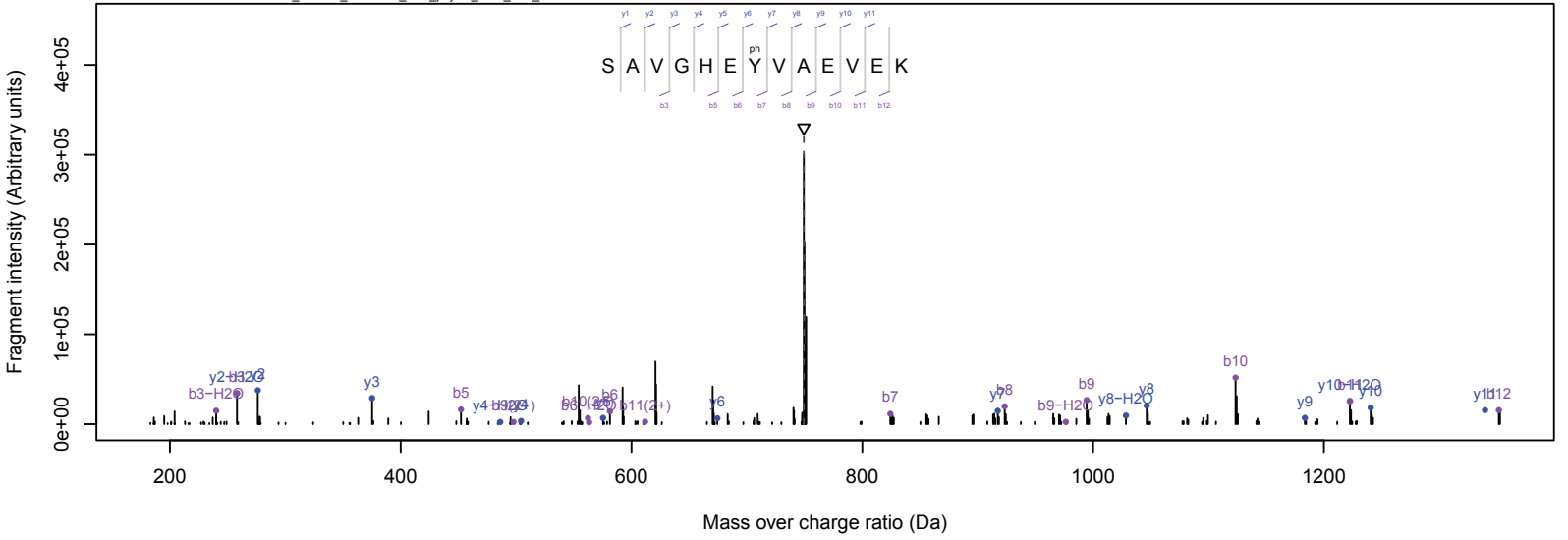
Andromeda score : 117.08



SAVGHEY(ph)VAEVEK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 12436

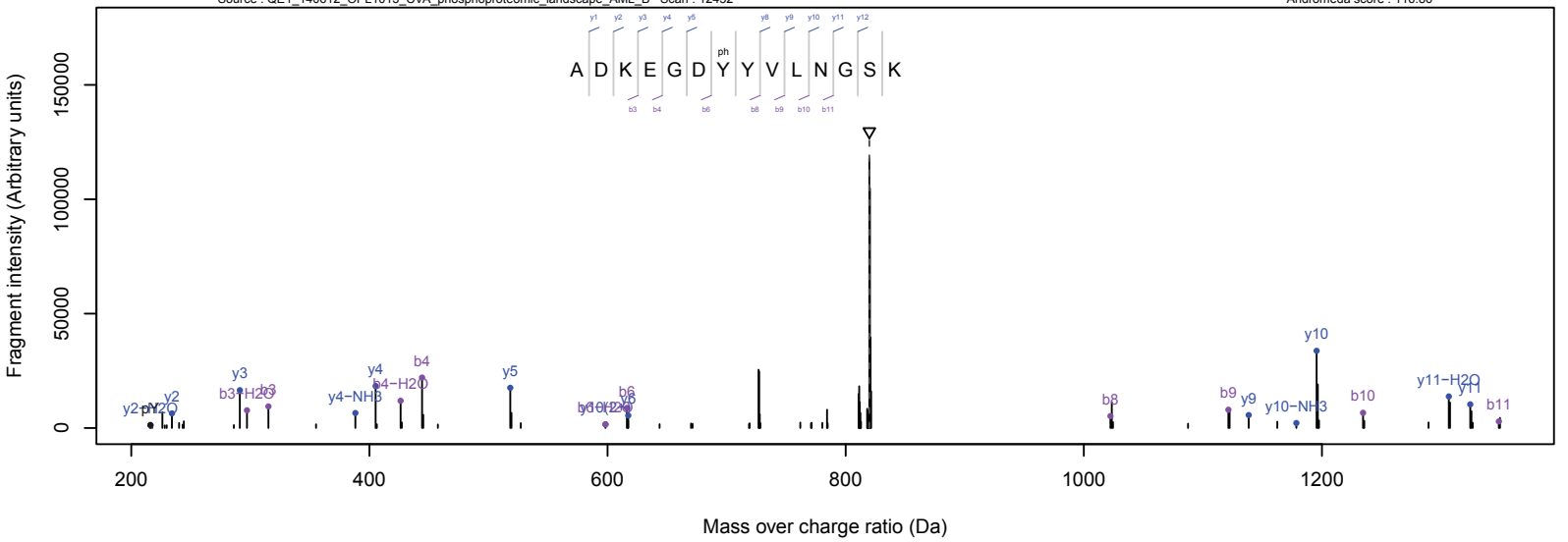
Andromeda score : 117.08



ADKEGDY(ph)YVLNGSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12452

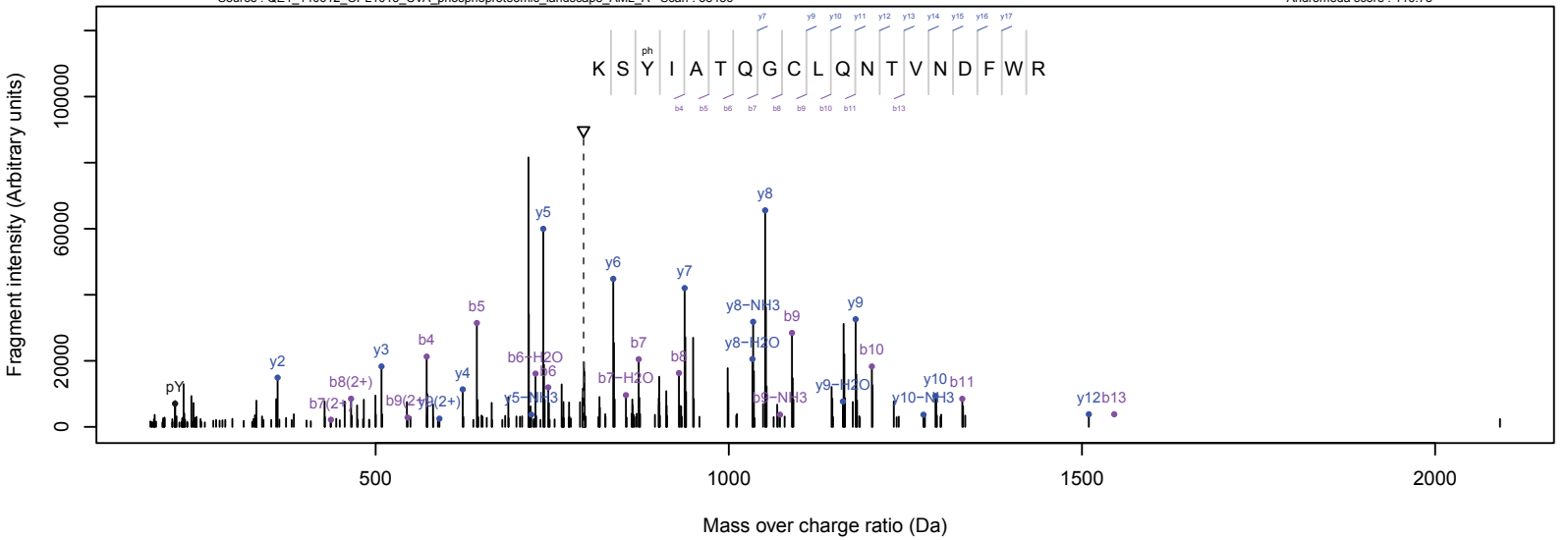
Andromeda score : 116.86



KSY(ph)IATQGCLQNTVDFWR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 38486

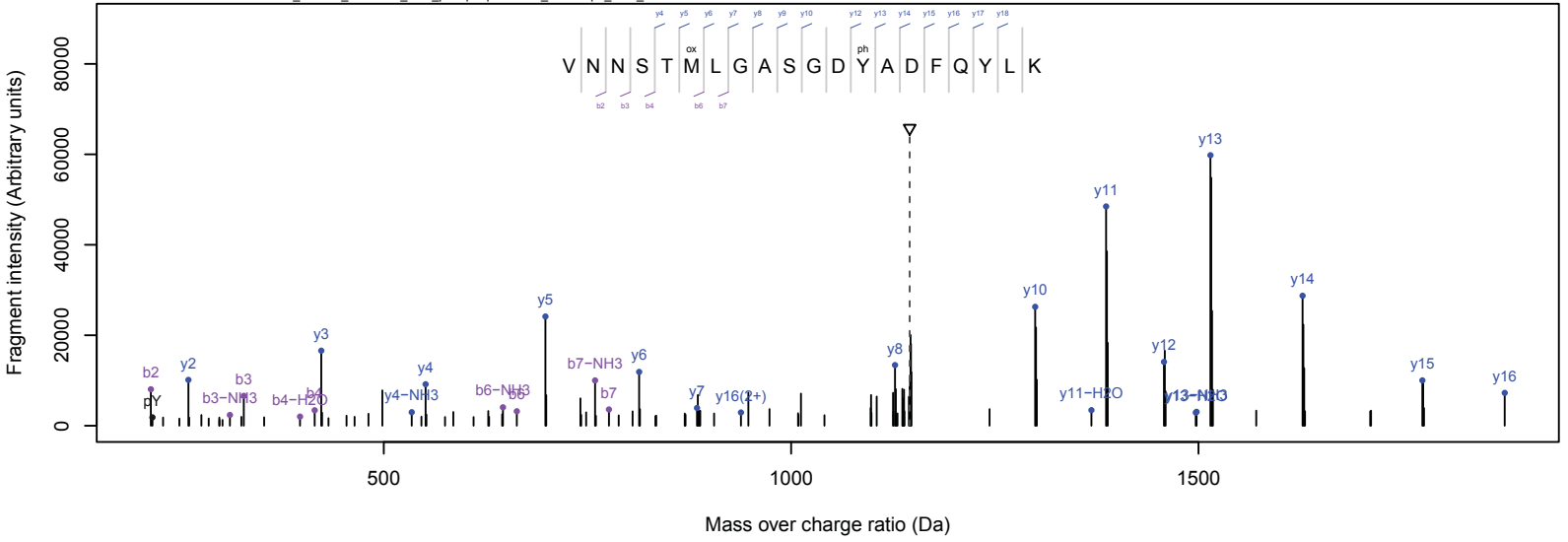
Andromeda score : 116.73



VNNSTM(ox)LGASGDY(ph)ADFQYLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33160

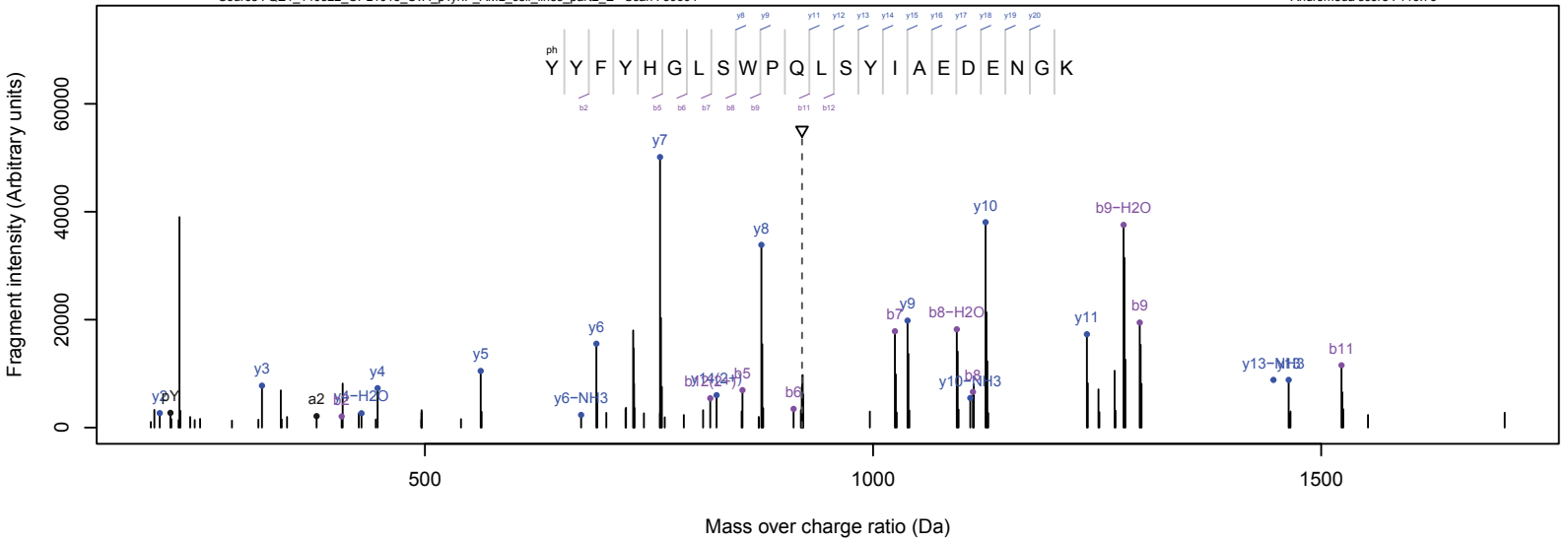
Andromeda score : 116.73



Y(ph)YFYHGLSWPQLSYIAEDENGK-3+

Source : QE1\_140822\_OPL1013\_Cva\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 39604

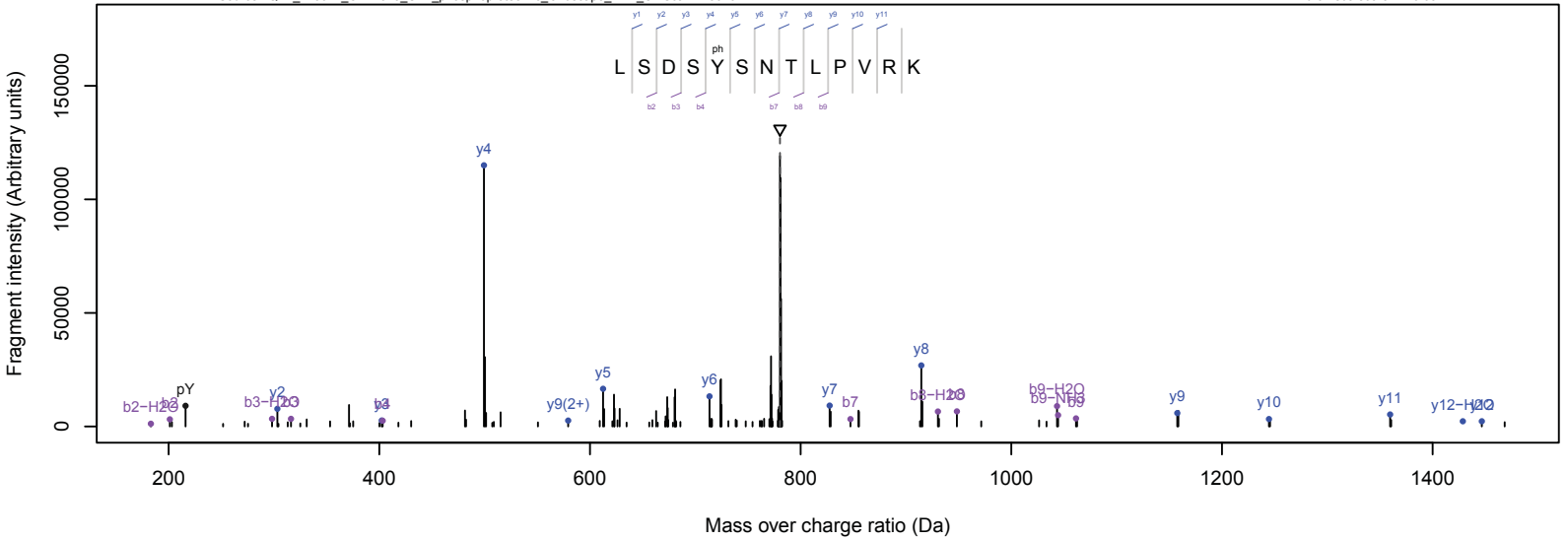
Andromeda score : 116.73



LSDSY(ph)SNTLPVRK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13010

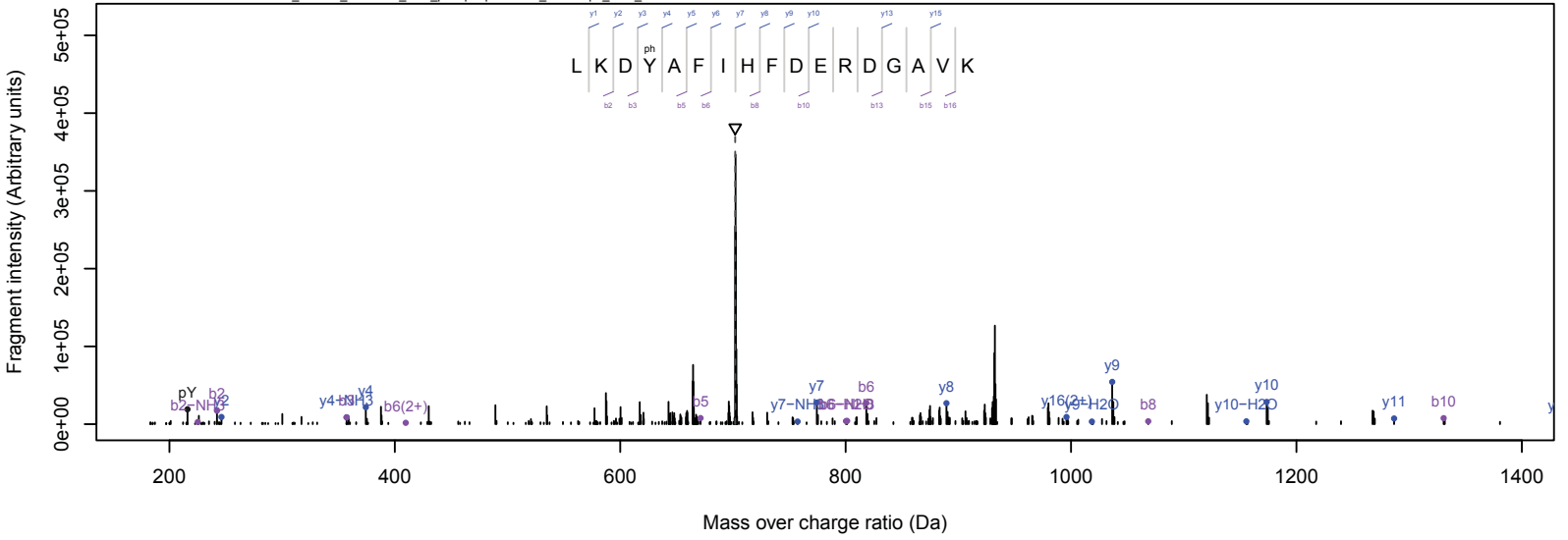
Andromeda score : 116.63



LKDY(ph)AFIHFDERDGAVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26798

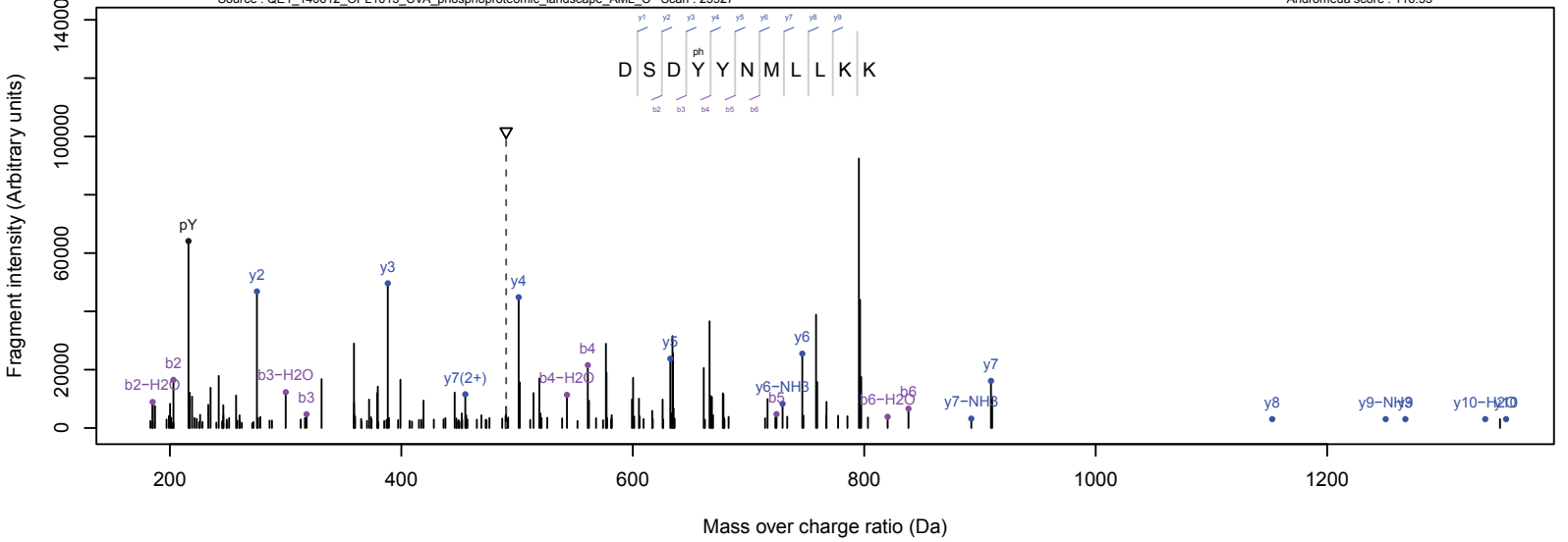
Andromeda score : 116.6



DSDY(ph)YNMLLKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25927

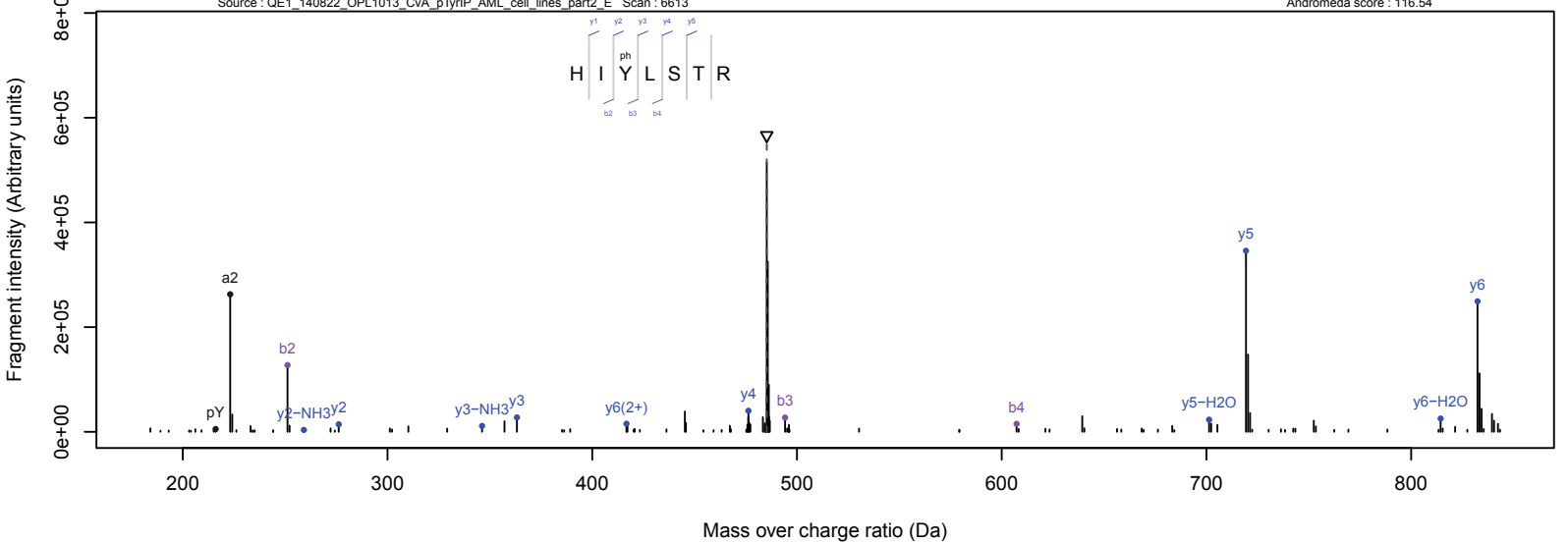
Andromeda score : 116.55



HIY(ph)LSTR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 6613

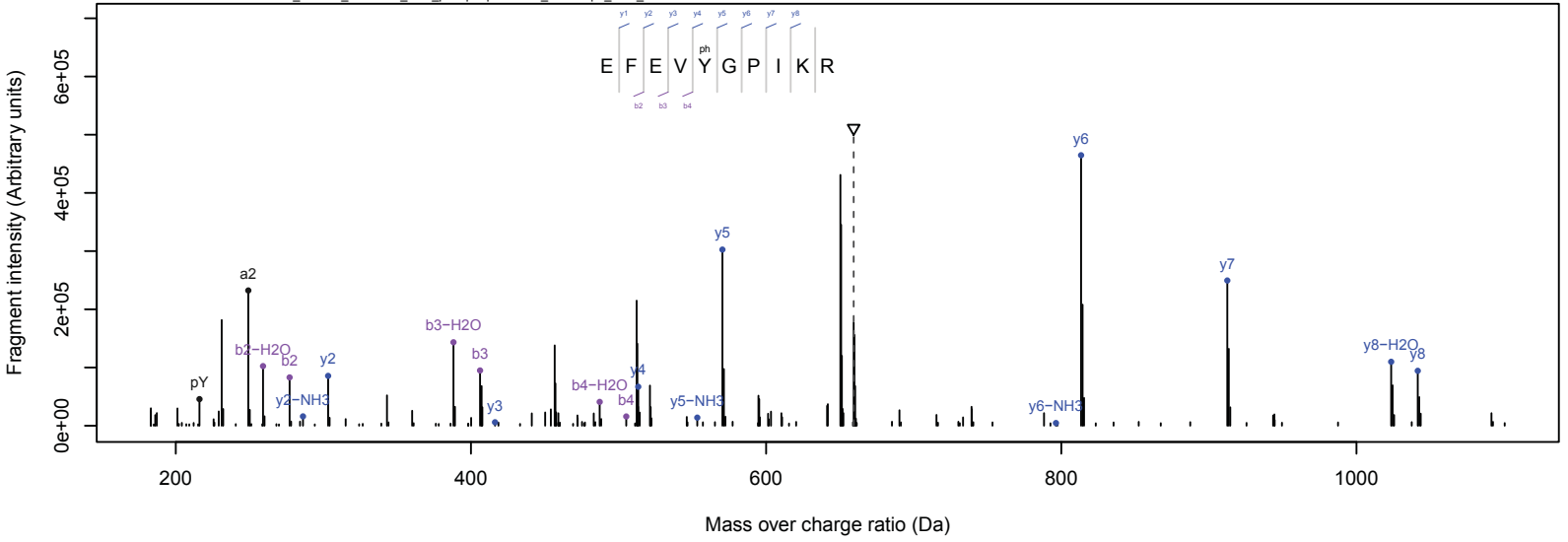
Andromeda score : 116.54



EFEVY(ph)GPIKR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 22196

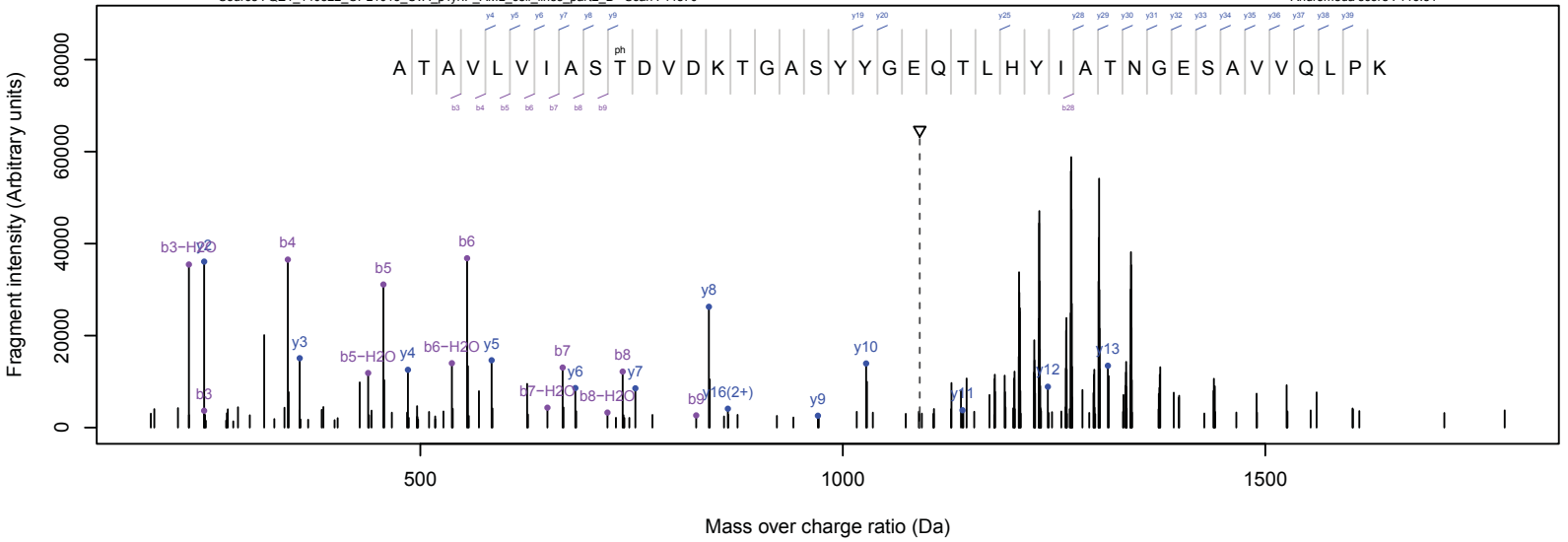
Andromeda score: 116.52



ATAVLVIAS(ph)DVKDTGASYYGEQTLHYIATNGESAVVQLPK-4+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 44870

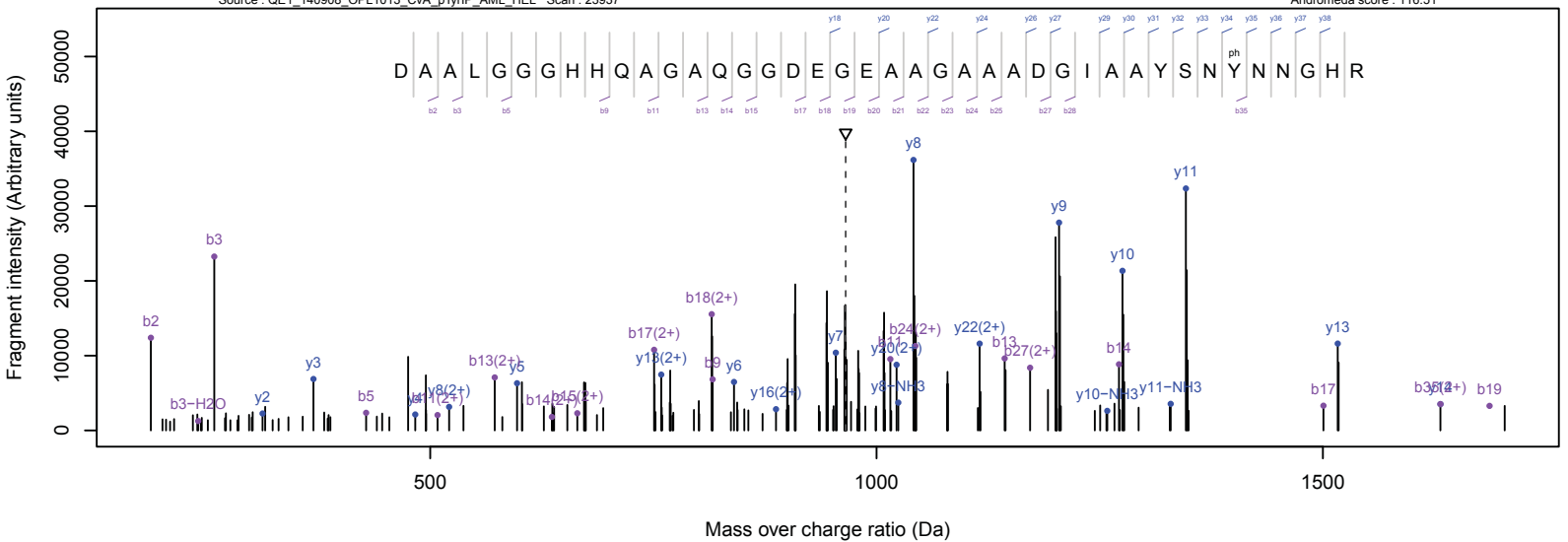
Andromeda score: 116.51



DAALGGHHQAGAQQGDEGEAAGAAADGIAAYSNY(ph)NNGHR-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 23937

Andromeda score: 116.51

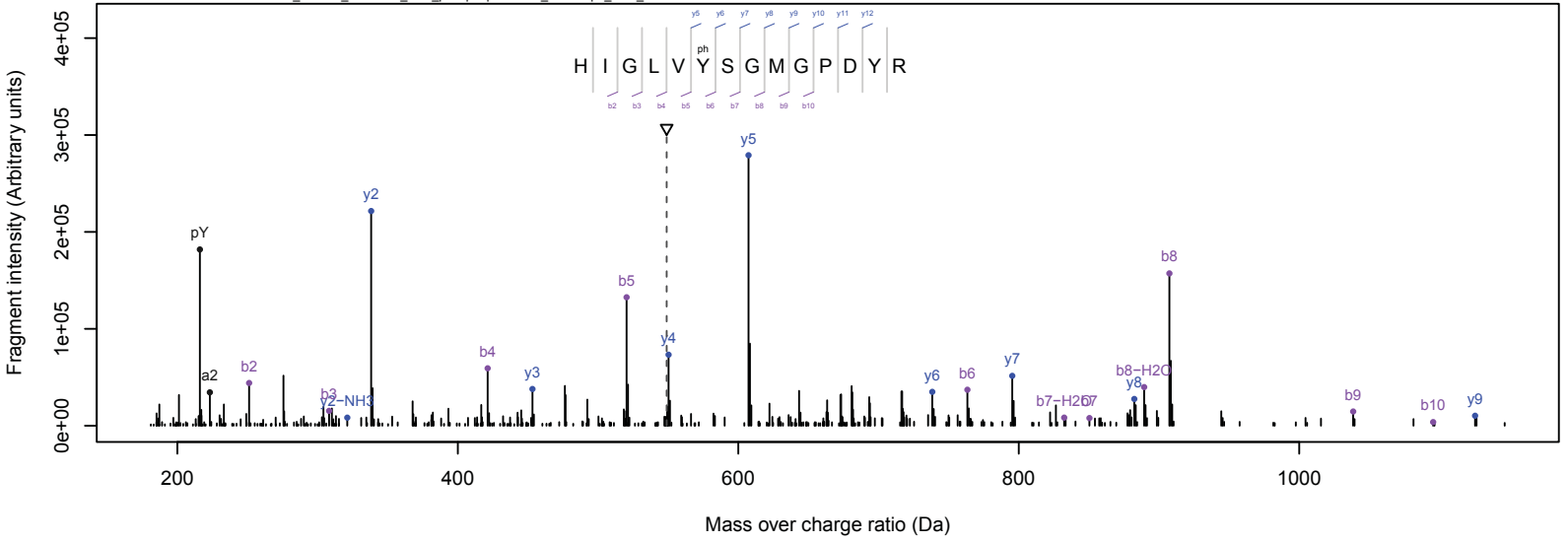




HIGLVY(ph)SGMGPDYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 27044

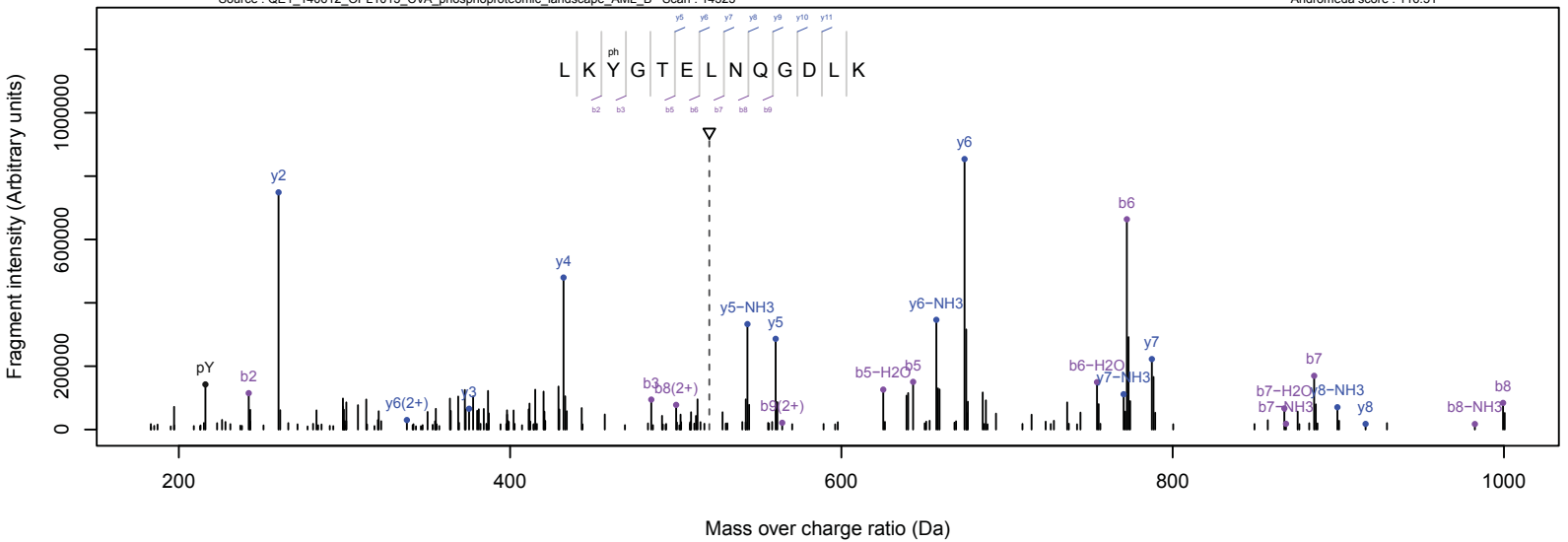
Andromeda score: 116.51



LKY(ph)GTELNQGLK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14323

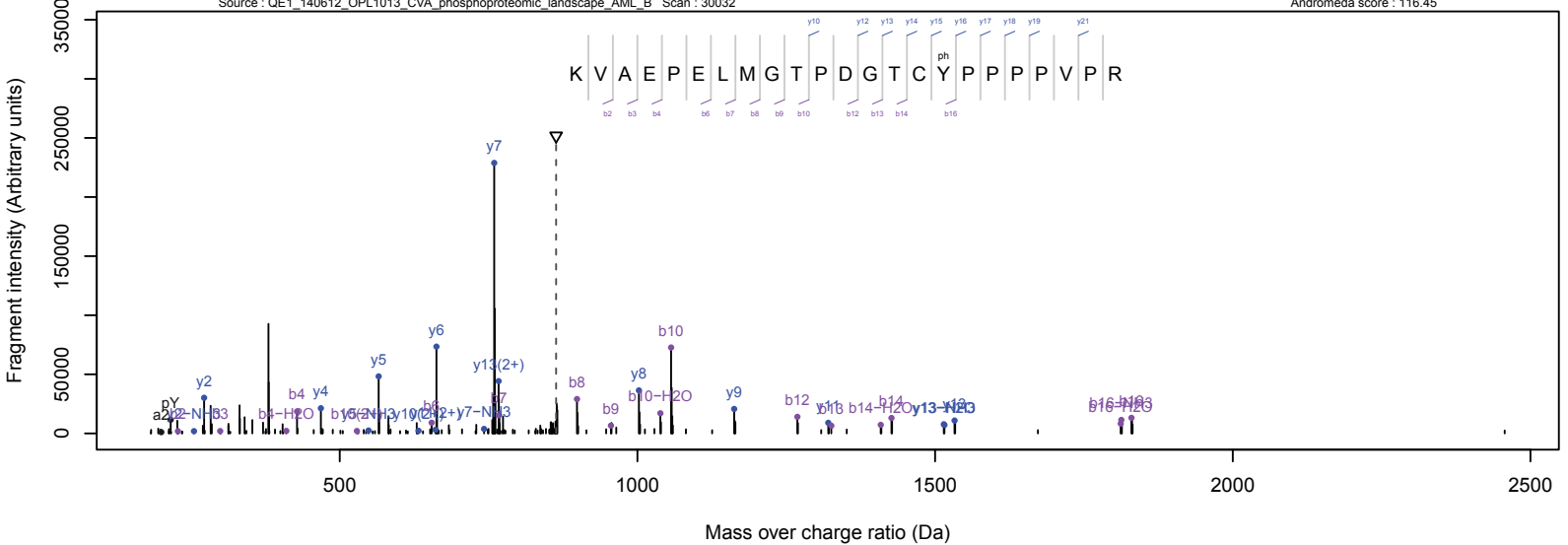
Andromeda score: 116.51



KVAEPELMGTPDGTCTCYPPPPVPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 30032

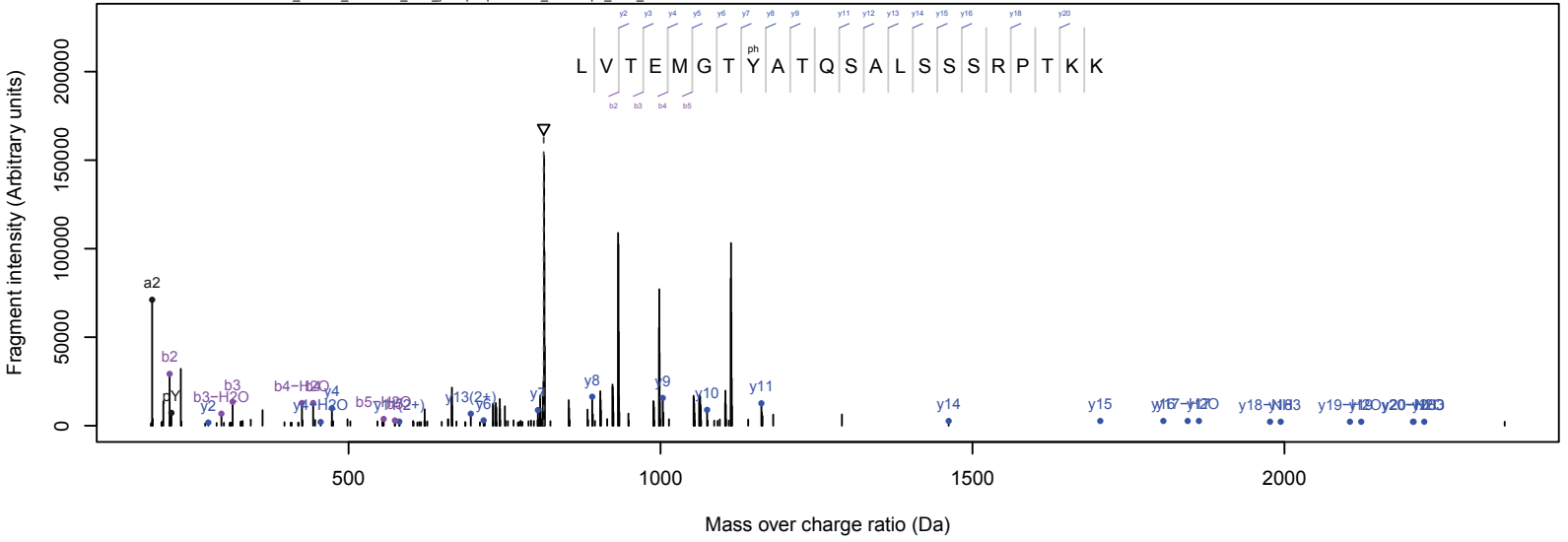
Andromeda score: 116.45



LVTEMGT<sup>(ph)</sup>ATQSALSSSRPTKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21845

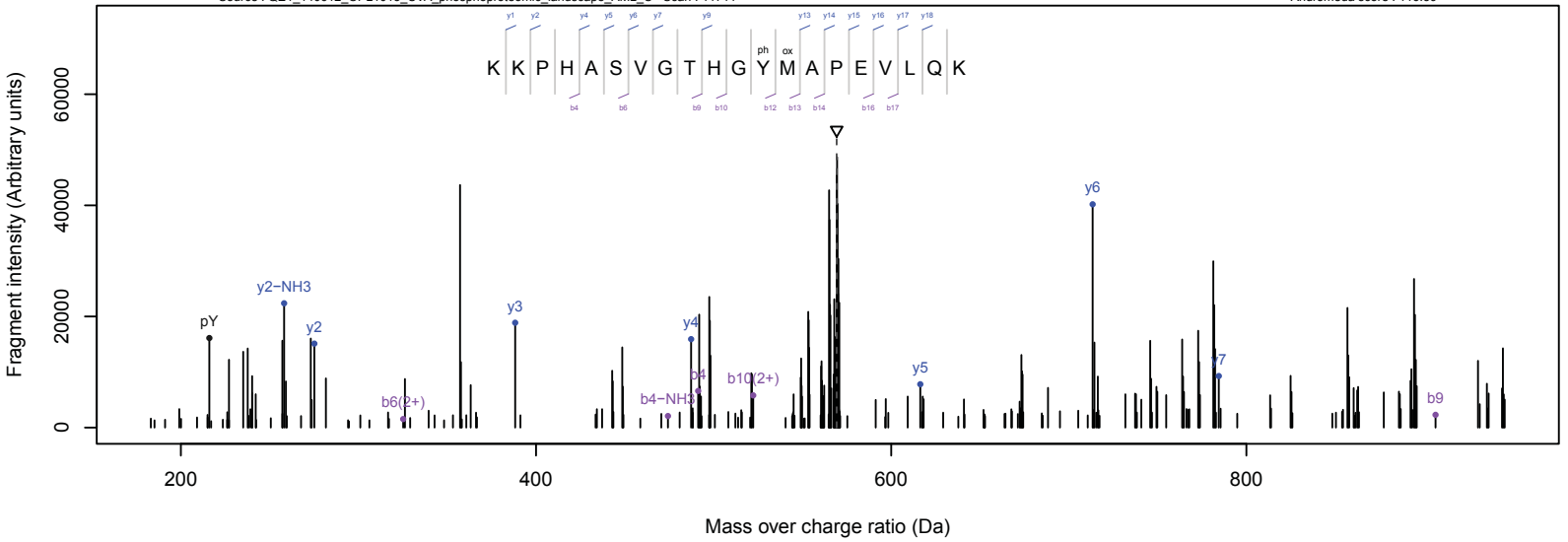
Andromeda score : 116.43



KKPHASVGT<sup>(ph)</sup>H<sup>(ox)</sup>MAPEVLQK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11711

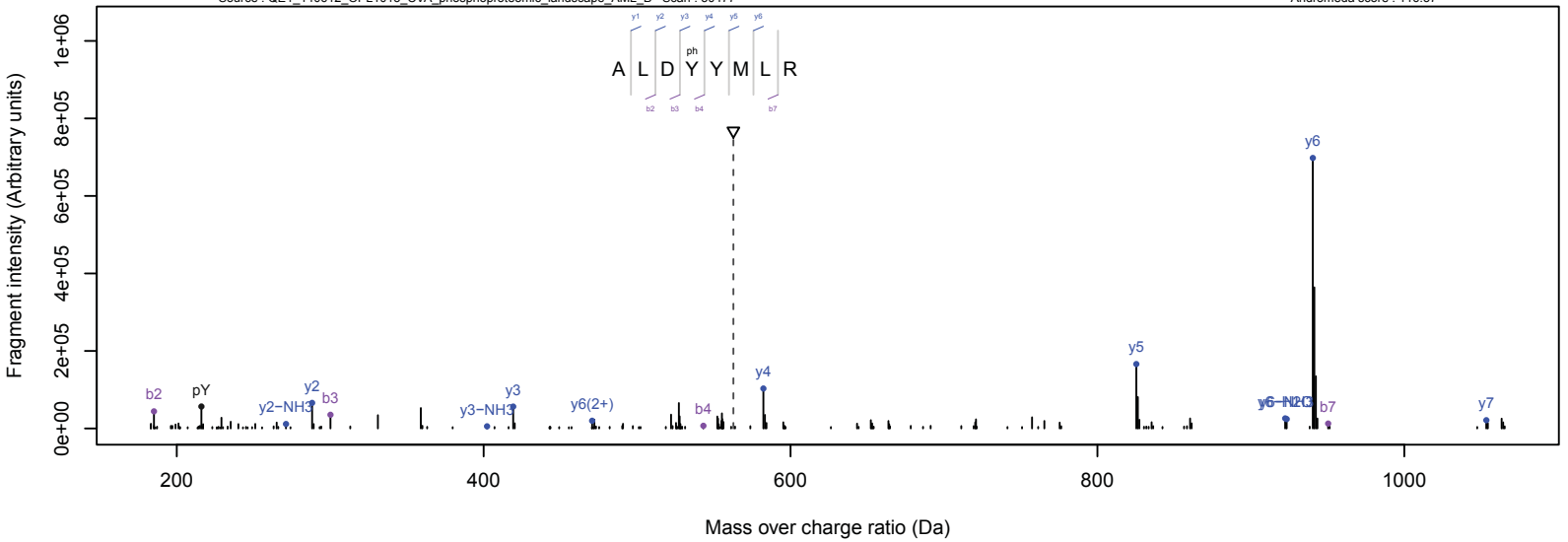
Andromeda score : 116.39



ALDY<sup>(ph)</sup>YMLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30477

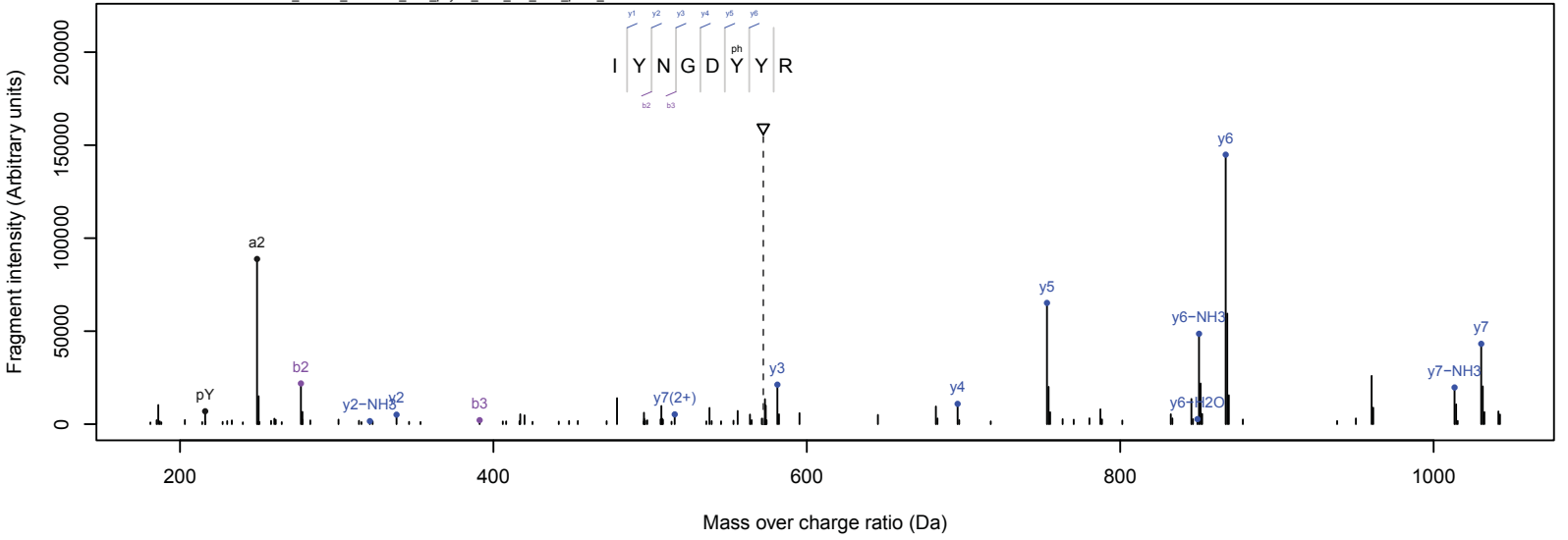
Andromeda score : 116.37



IYNGDY(ph)YR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 6258

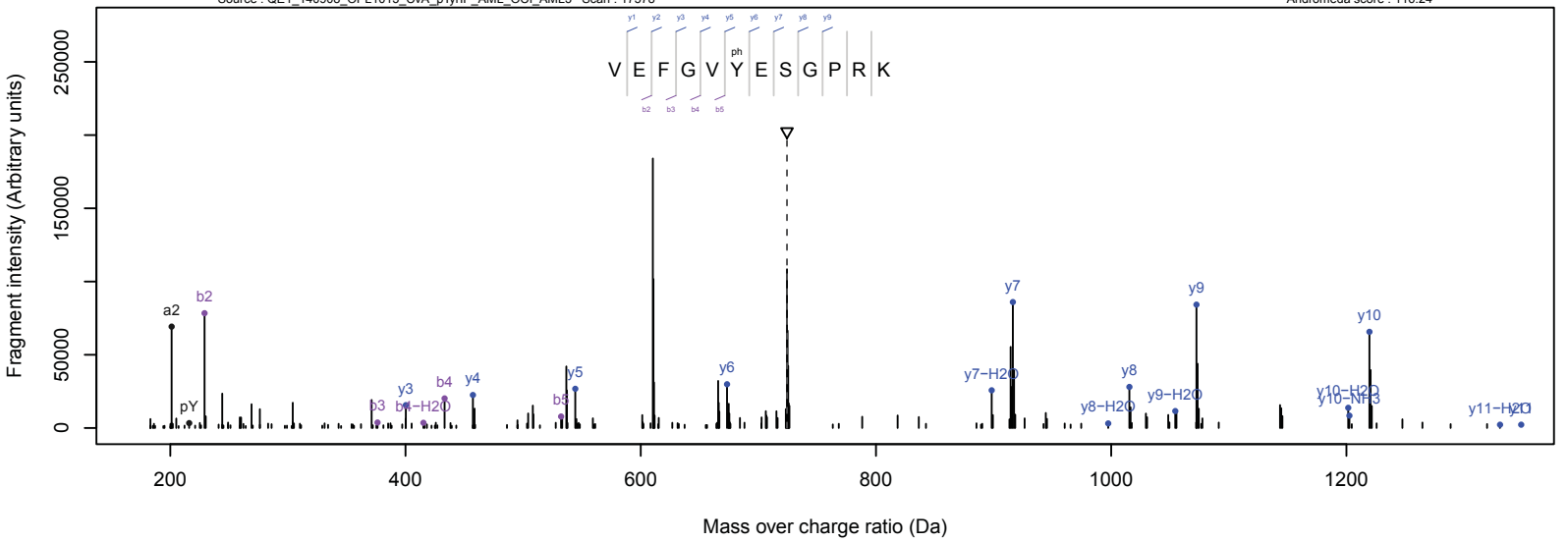
Andromeda score : 116.37



VEFGVY(ph)ESGPRK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 17578

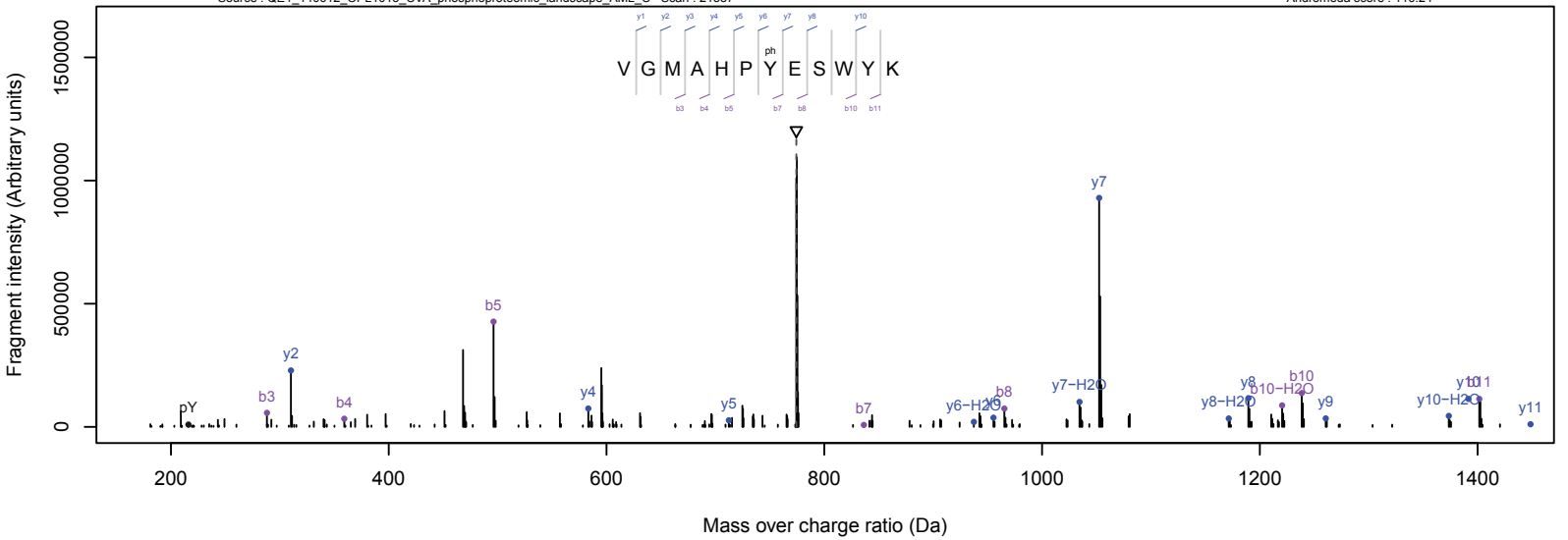
Andromeda score : 116.24



VGMAHPY(ph)ESWYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21387

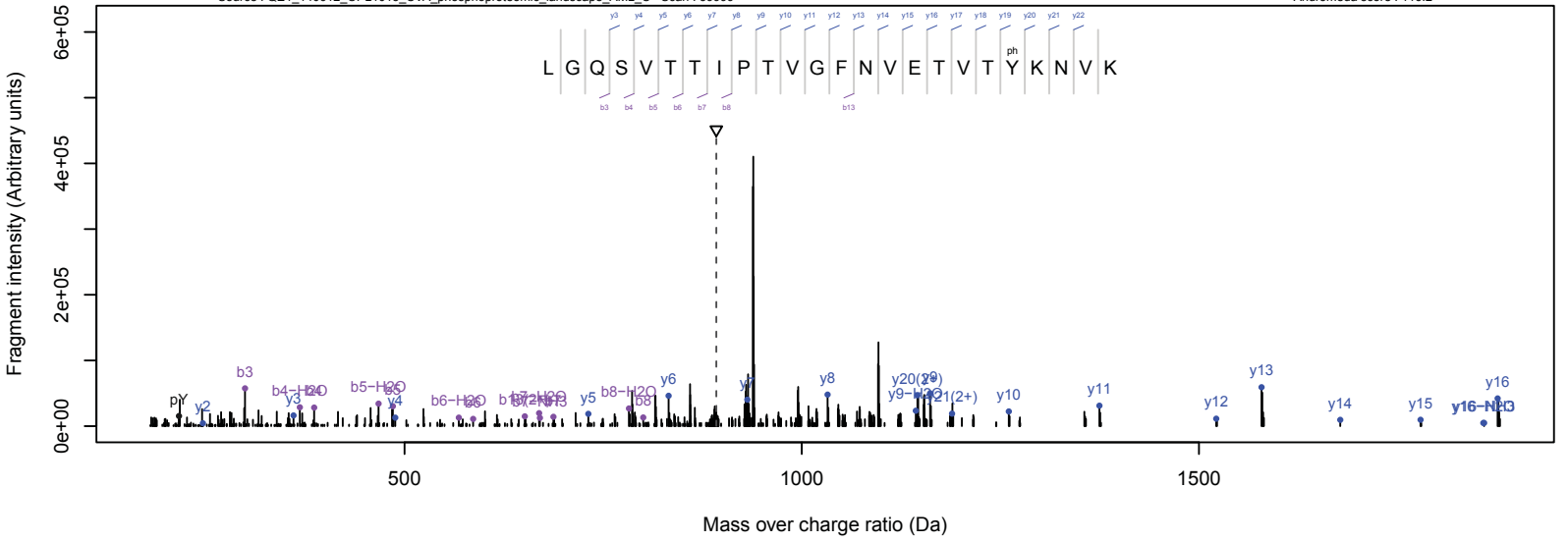
Andromeda score : 116.24



LGQSVTTIPTVGFNVETVY(ph)KNVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39999

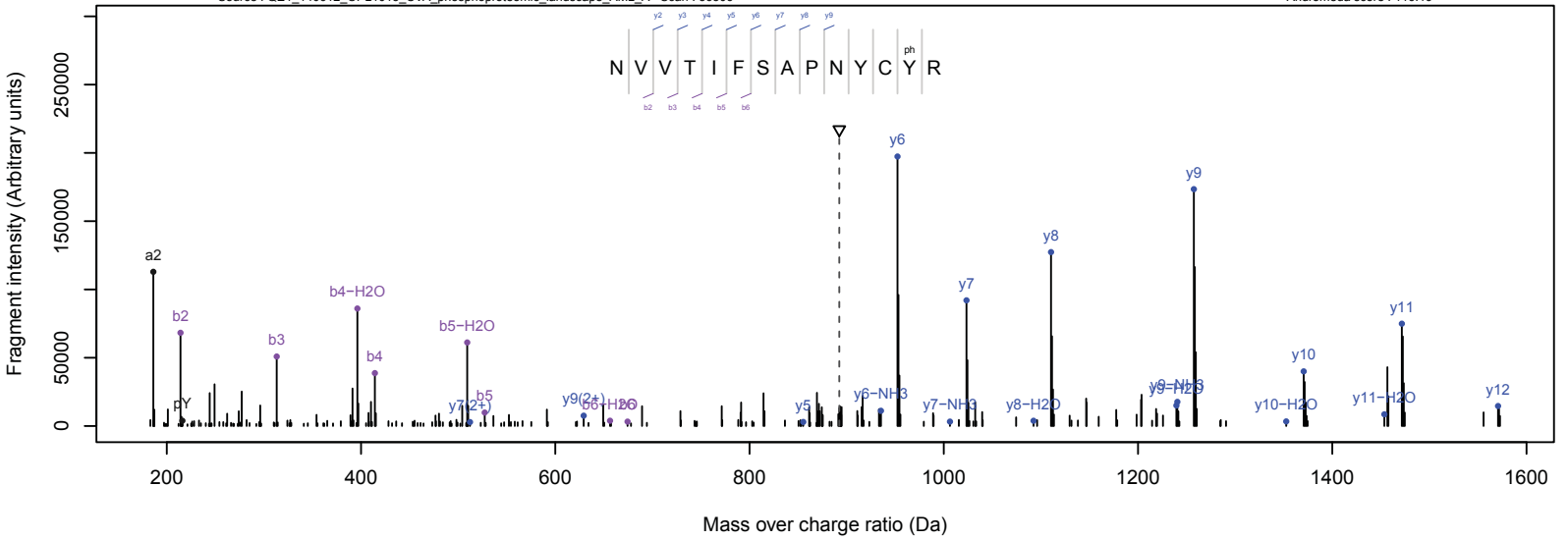
Andromeda score : 116.2



NVVTIFSAPNYC(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36506

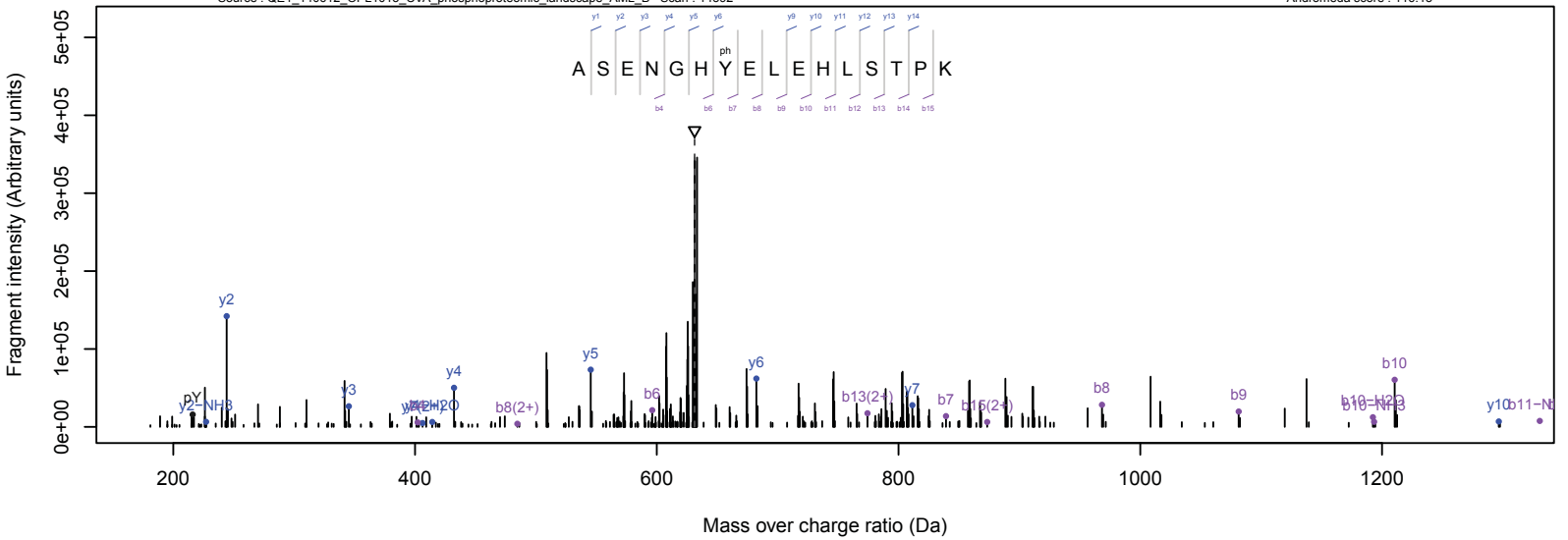
Andromeda score : 116.19



ASENGHY(ph)ELEHLSTPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14592

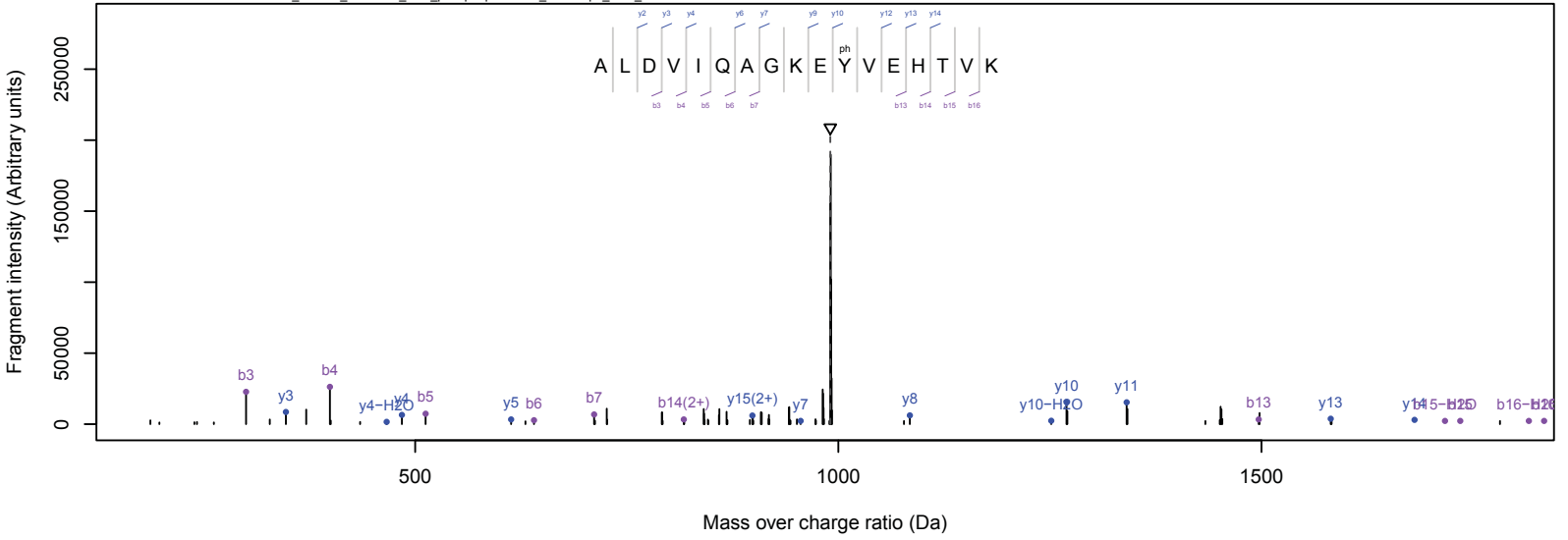
Andromeda score : 116.13



**ALDVIQAGKEY(ph)VEHTVK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28053

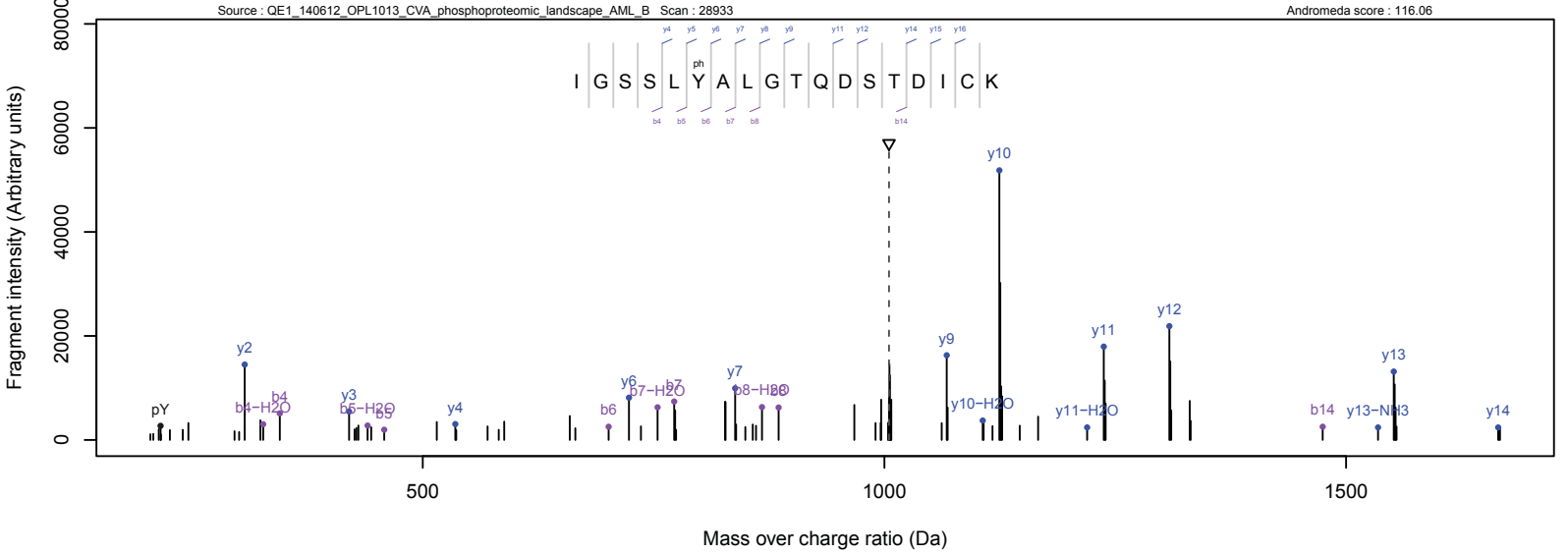
Andromeda score : 116.06



**IGSSLY(ph)ALGTQDSTDICK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28933

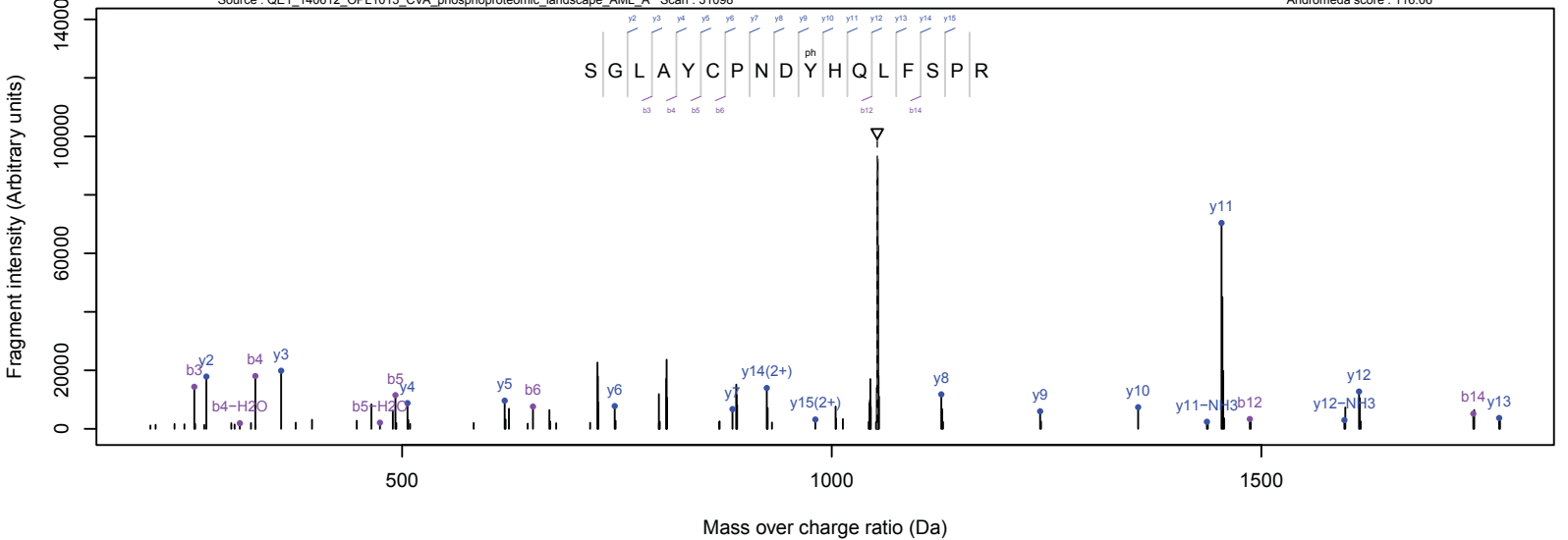
Andromeda score : 116.06



**SGLAYCPNDY(ph)HQLFSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 31098

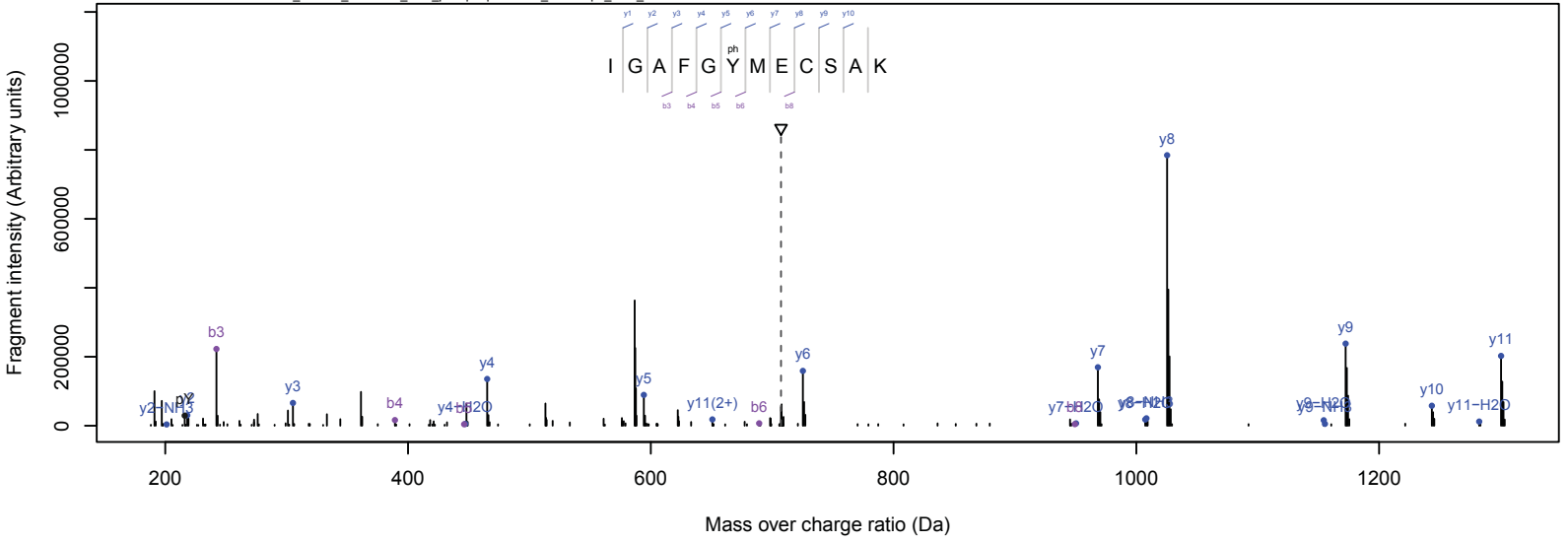
Andromeda score : 116.06



IGAFGY(ph)MECSAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 24892

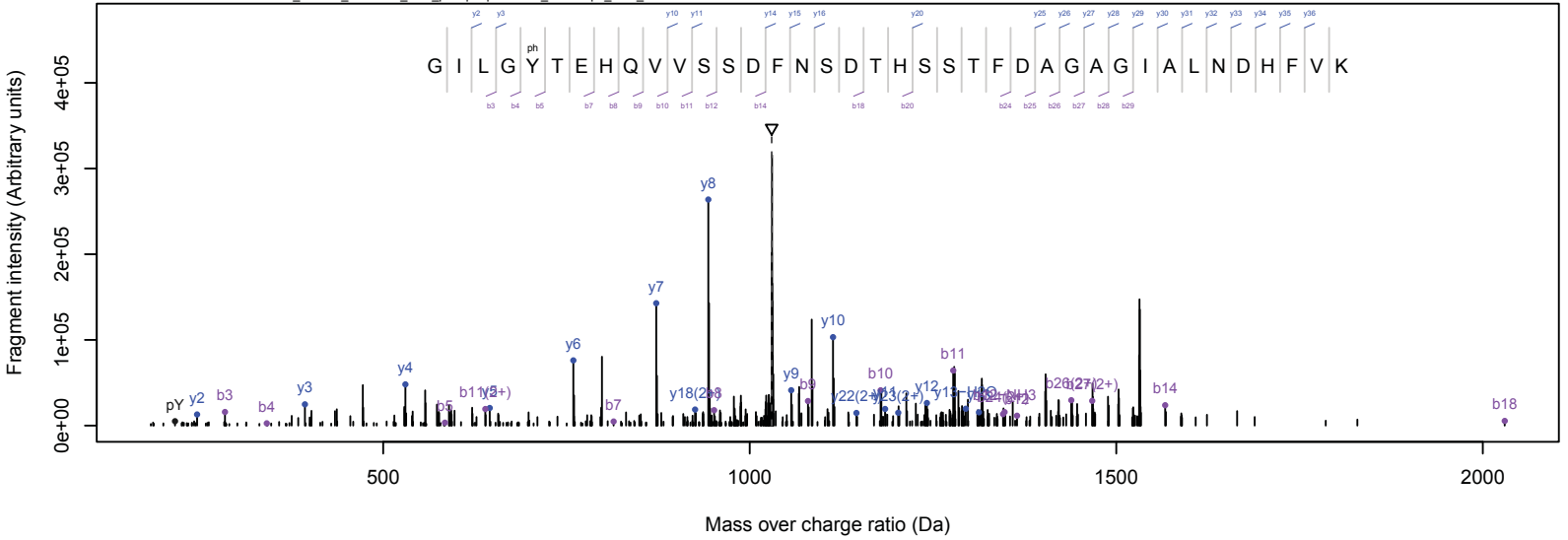
Andromeda score : 116.03



GILGY(ph)TEHQVVSSDFNSDTHSSTFDAGAGIALNDHFVK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41541

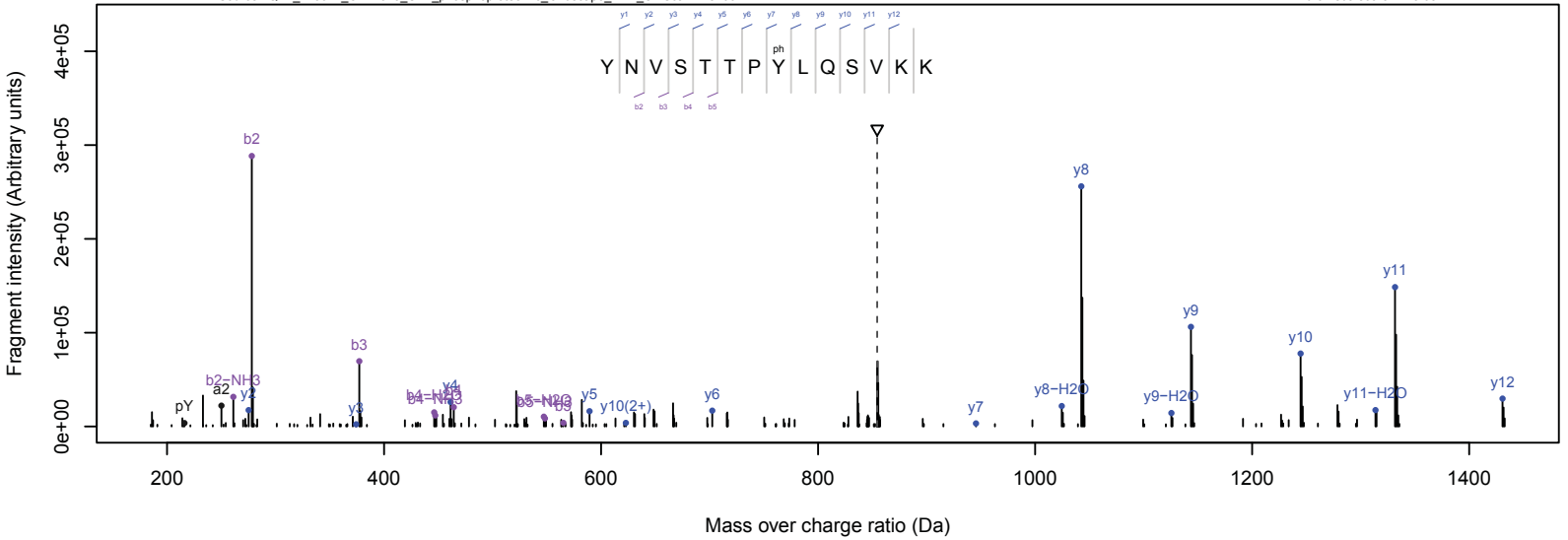
Andromeda score : 115.87



YNVSTTPY(ph)LQSVKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18200

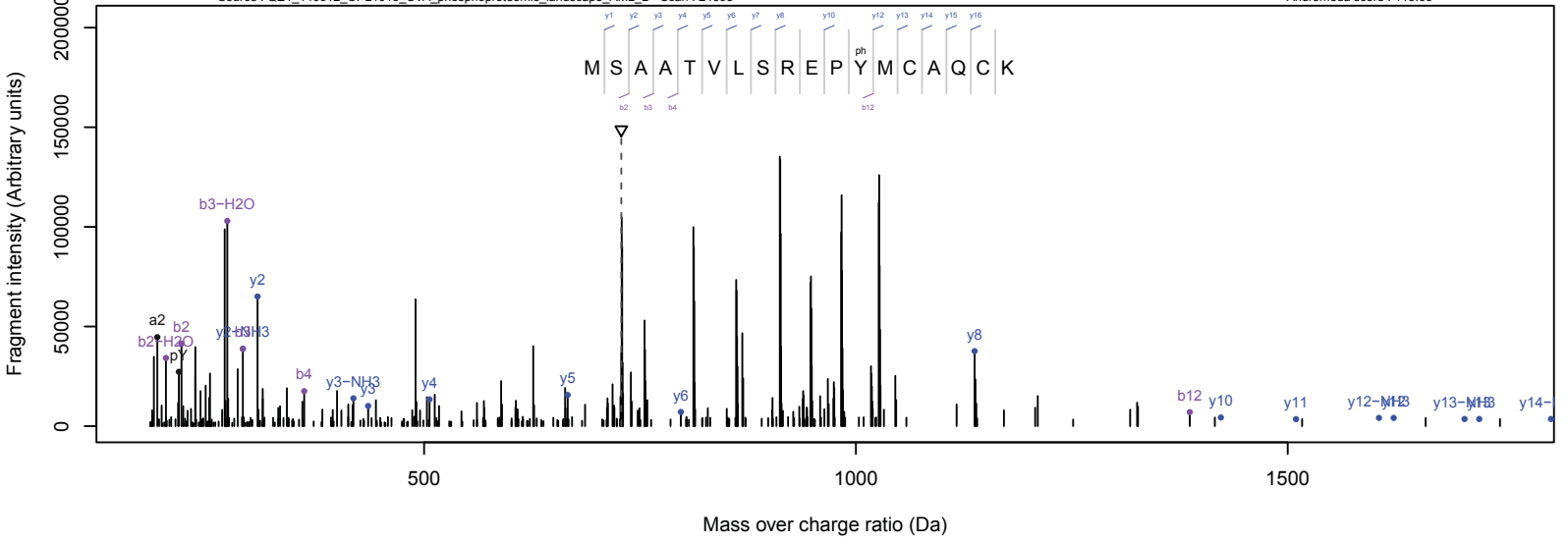
Andromeda score : 115.68



**MSAATVLSREPY(ph)MCAQCK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24698

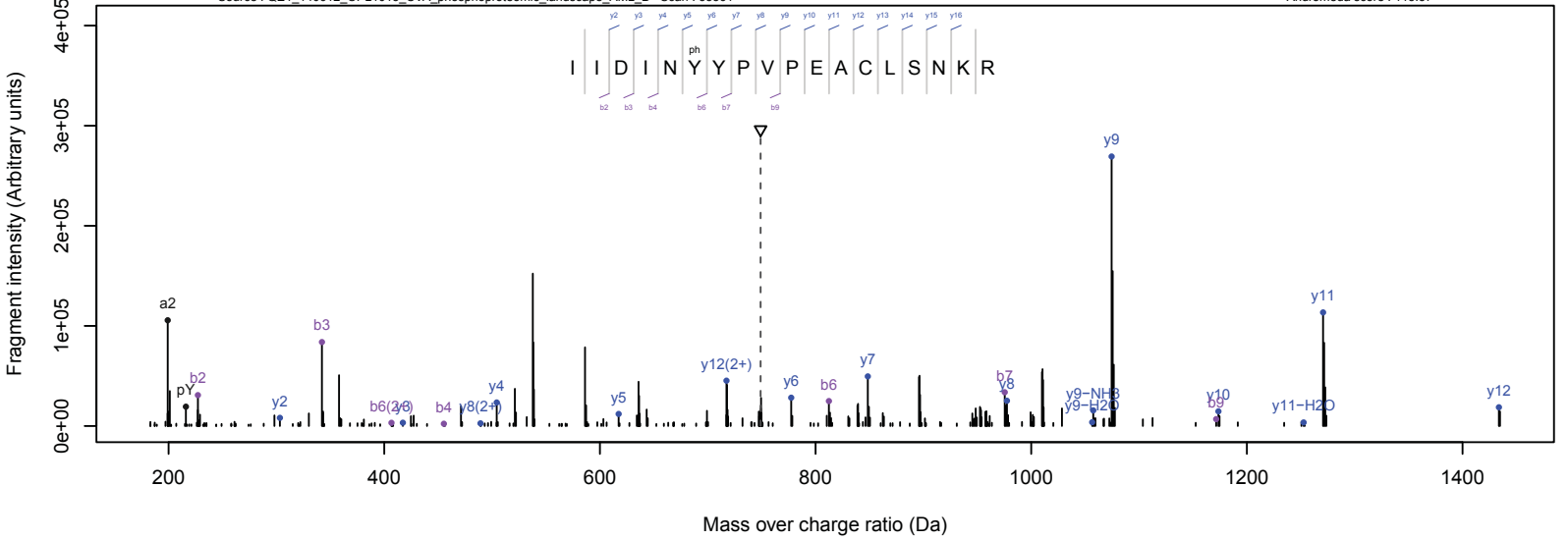
Andromeda score : 115.66



**IIDINY(ph)YPVPEACLSNKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35561

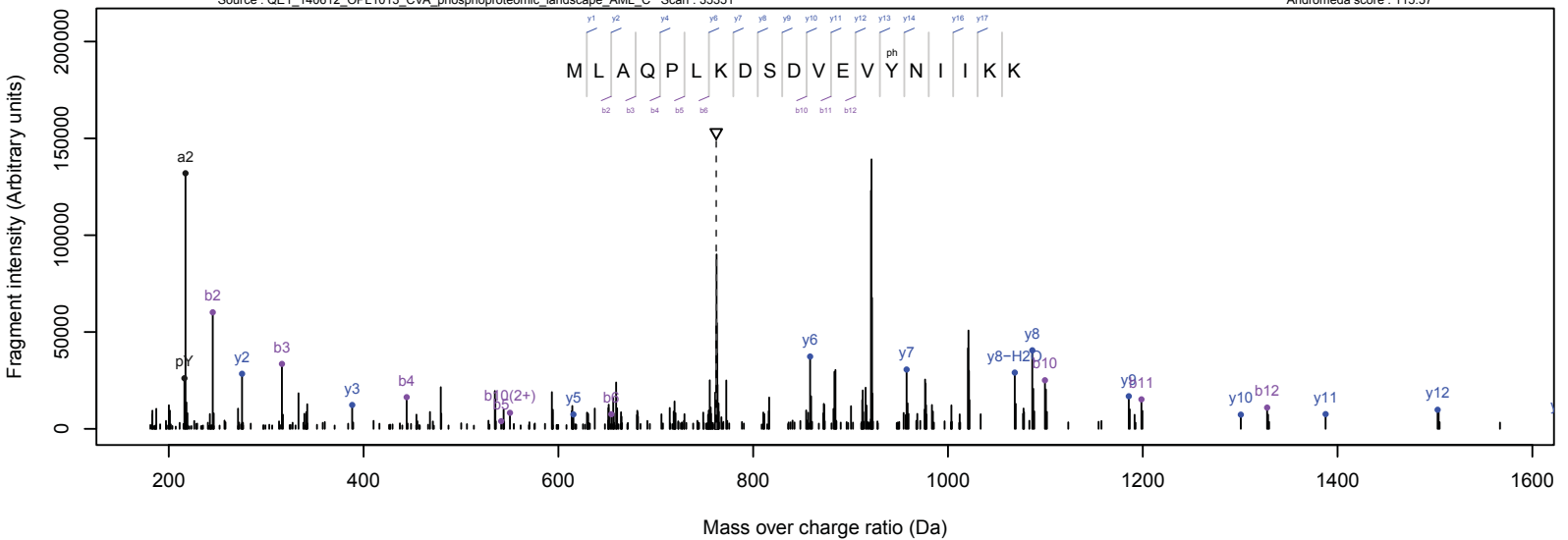
Andromeda score : 115.57



**MLAQLKDSDEVY(ph)NIKK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35351

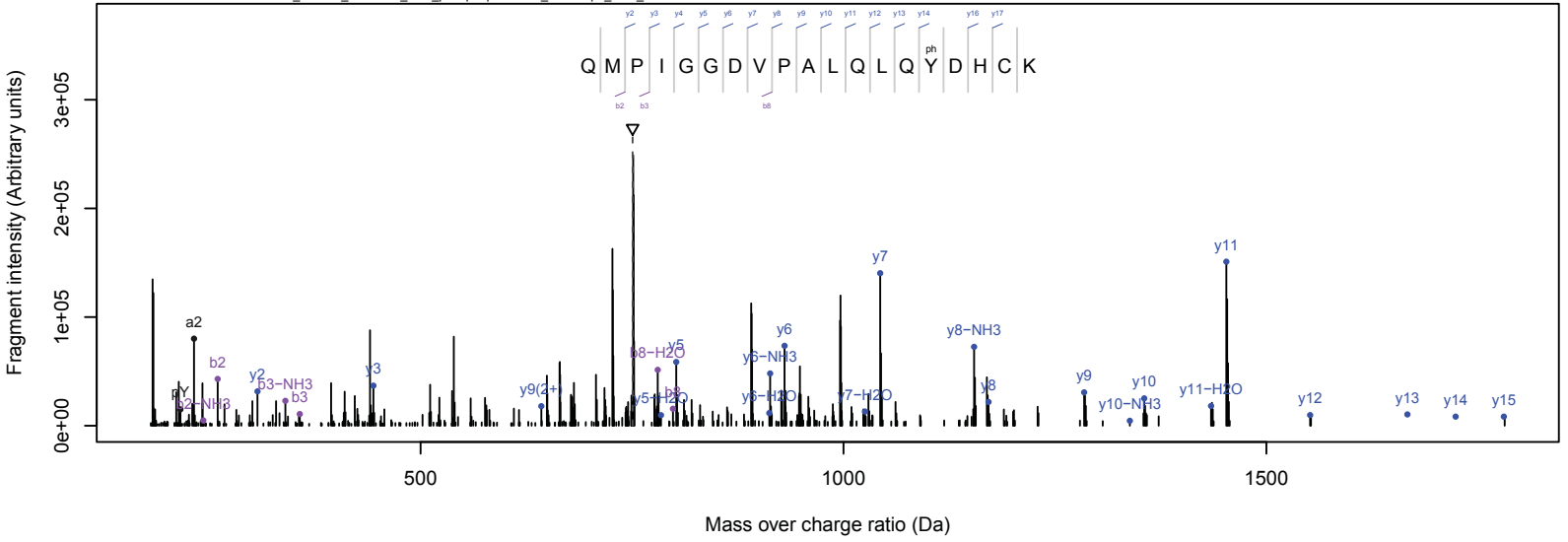
Andromeda score : 115.57



QMPIGGDVPALQLQY(ph)DHCK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34614

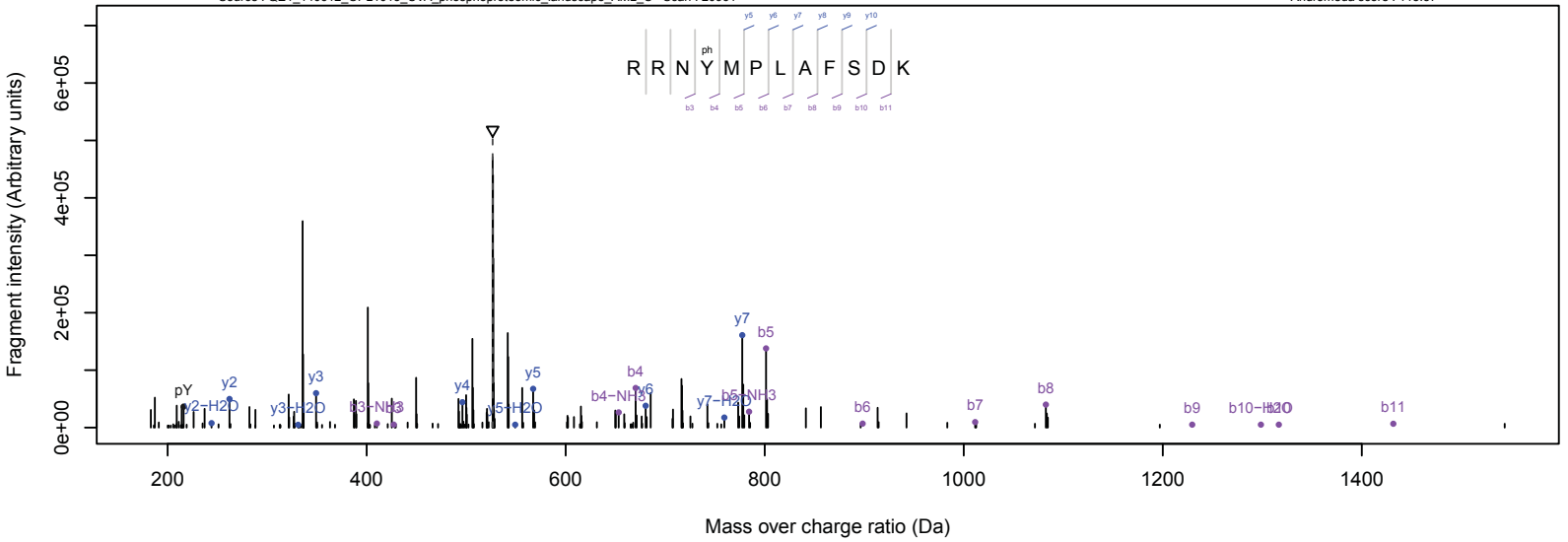
Andromeda score : 115.57



RRNY(ph)MPLAFSDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20961

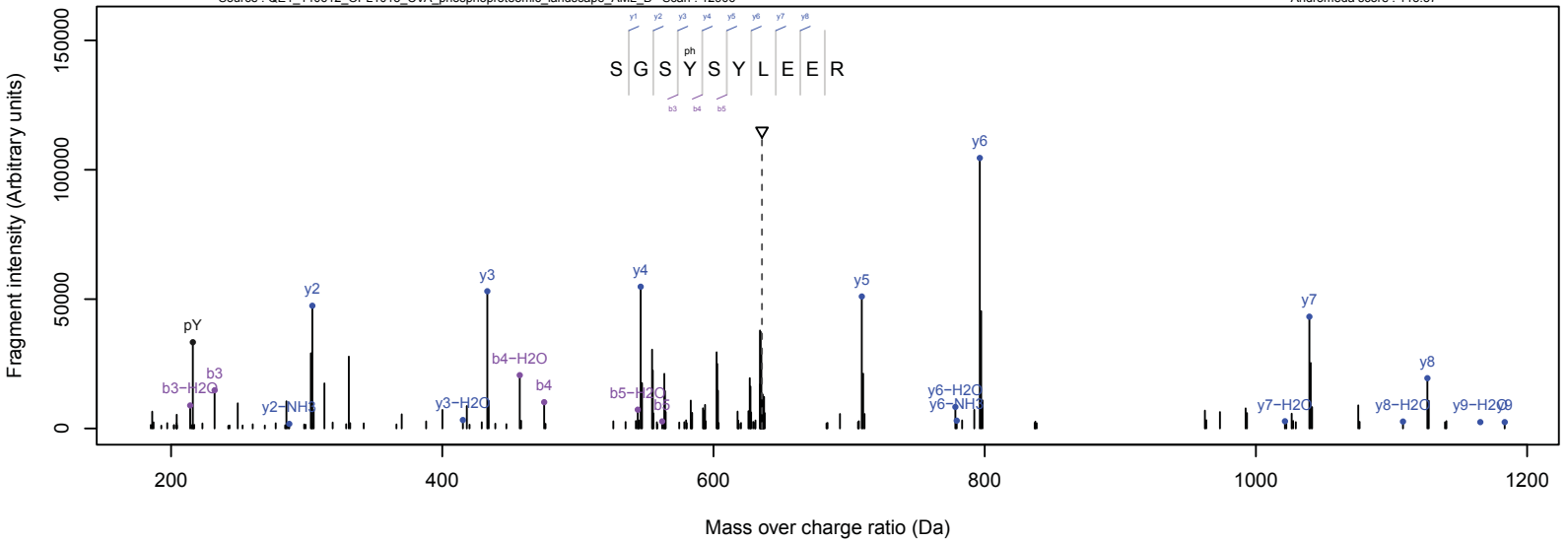
Andromeda score : 115.57



SGSY(ph)SYLEER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12906

Andromeda score : 115.57

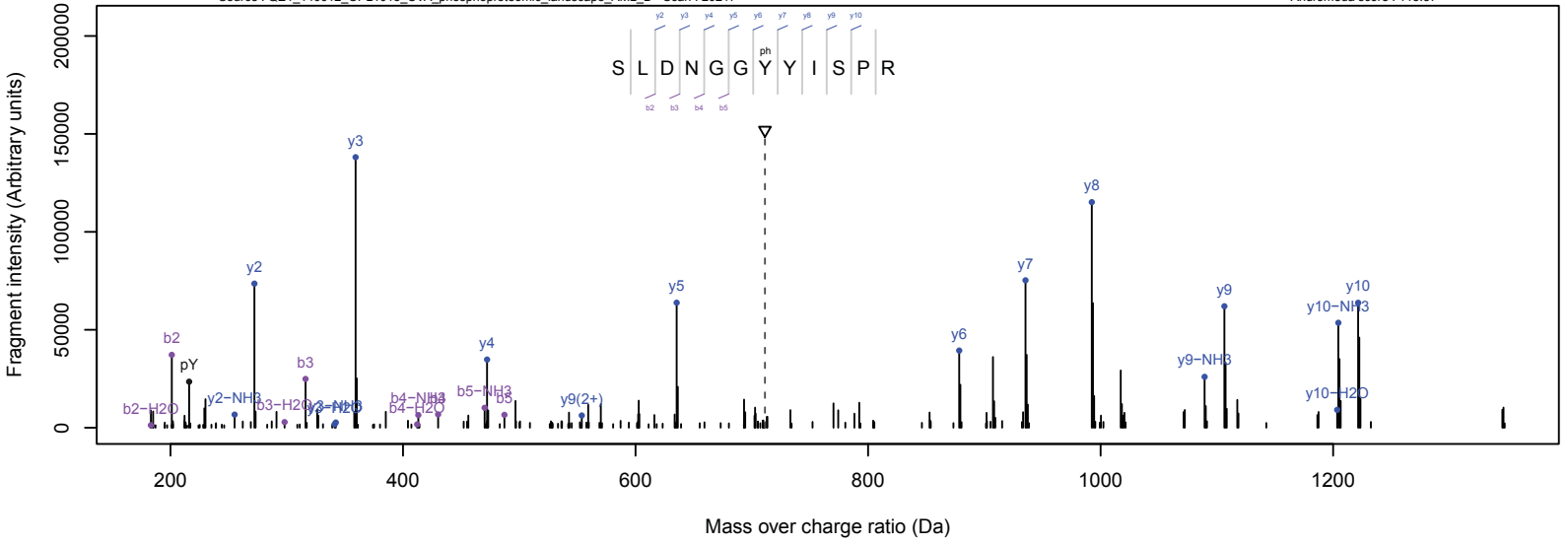




SLDNGGY(ph)YISPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 20217

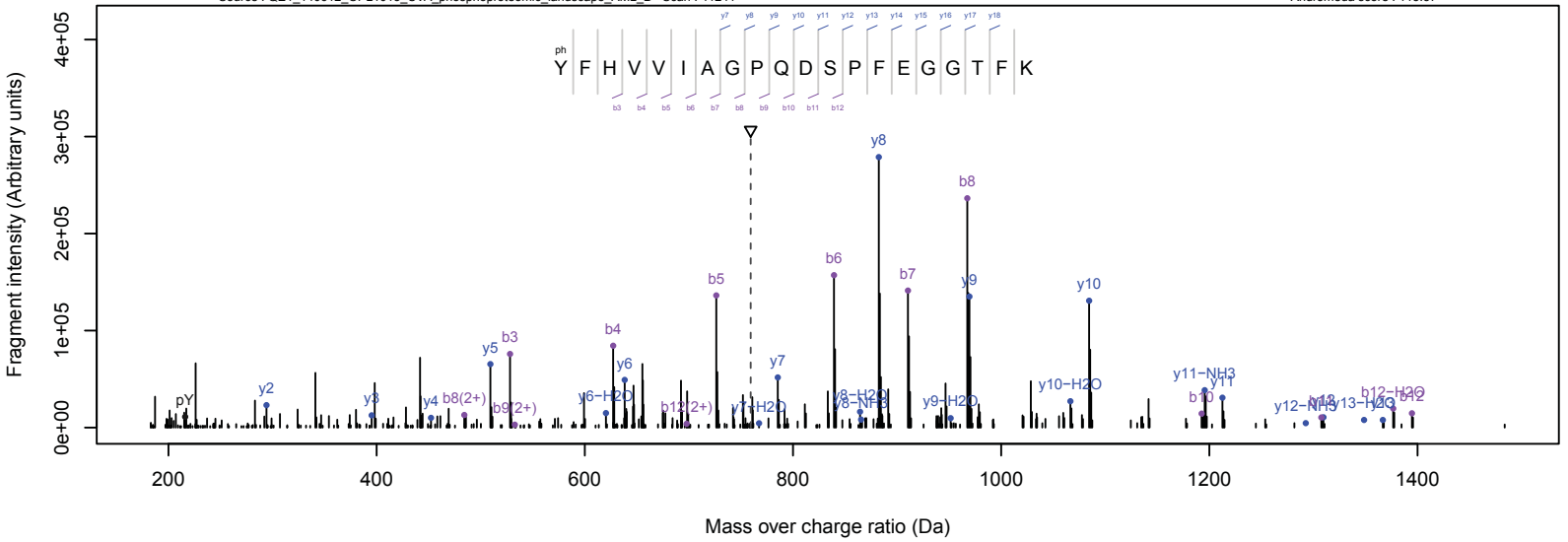
Andromeda score: 115.57



Y(ph)FHVVIAGPQDSPFEGGTFK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 41241

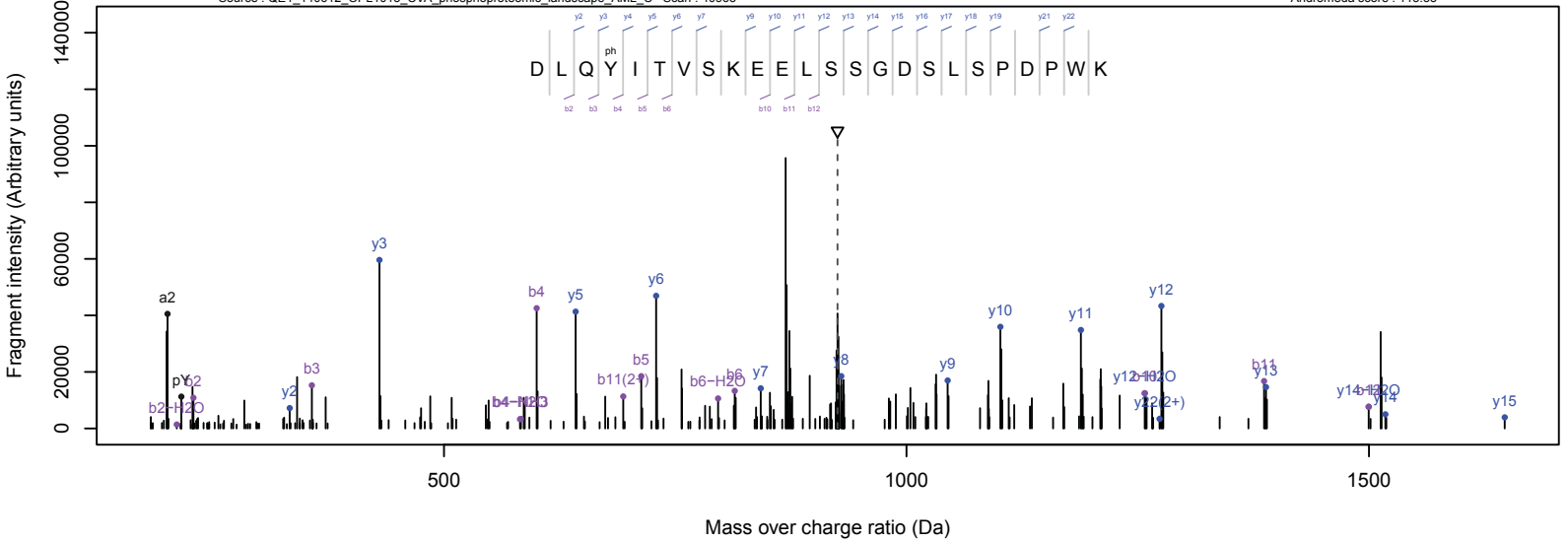
Andromeda score: 115.57



DLQY(ph)ITVSKHEELSSGDSLSPDPWK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 40963

Andromeda score: 115.55

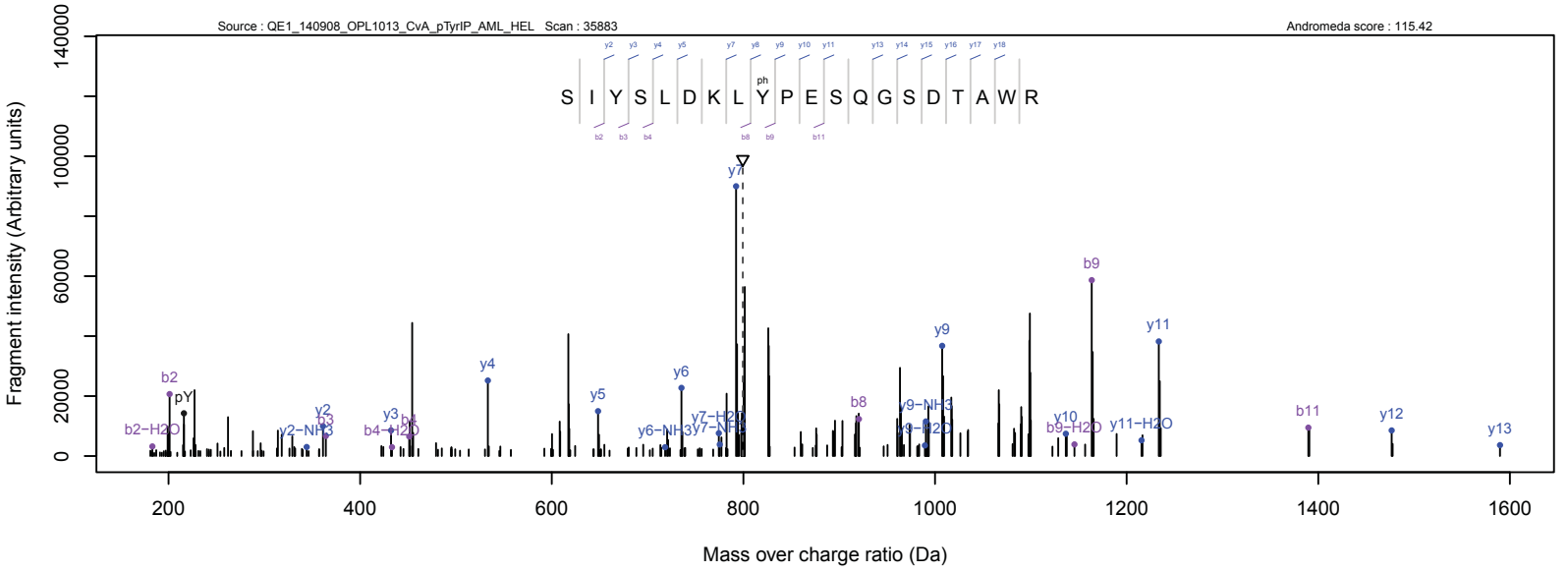




SIYSLDKLY(ph)PESQGSDTAWR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35883

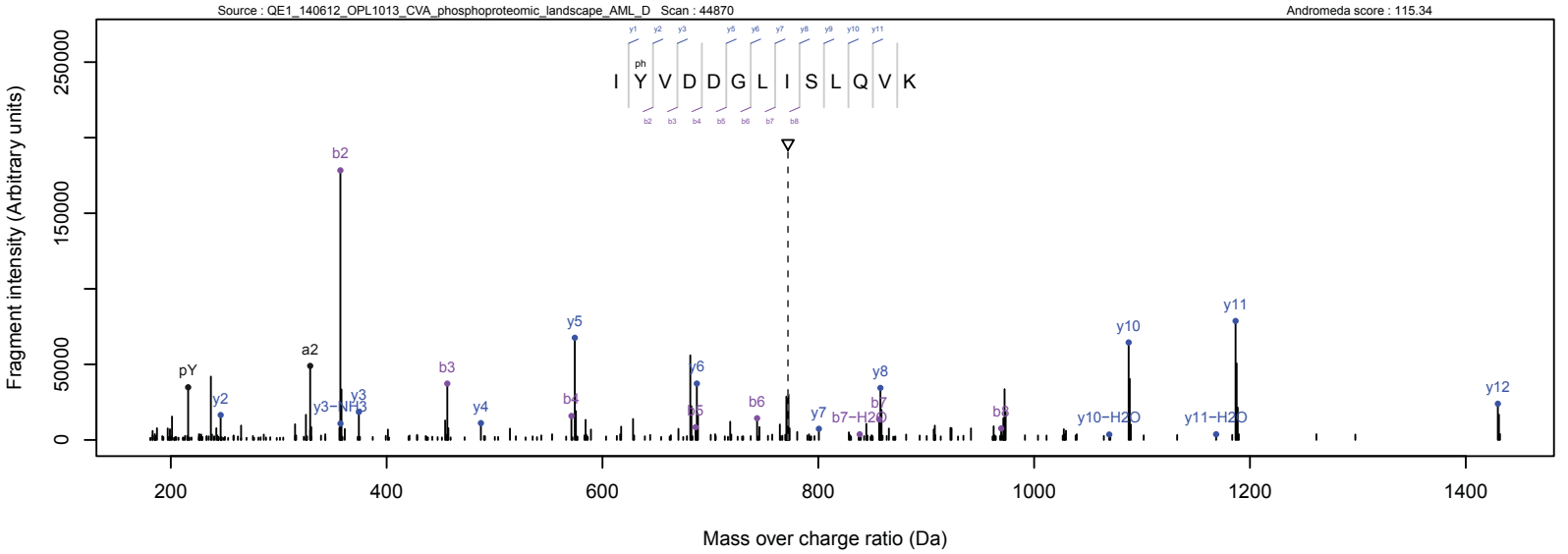
Andromeda score : 115.42



IY(ph)VDDGLISLQVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 44870

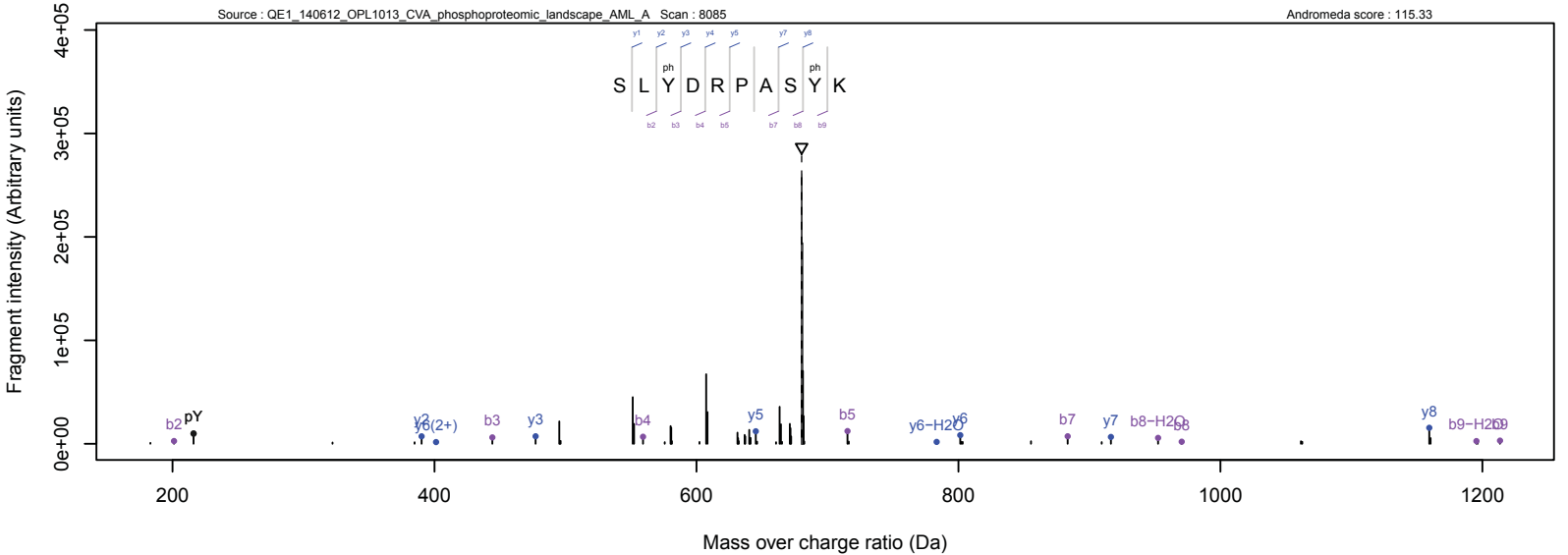
Andromeda score : 115.34



SLY(ph)DRPASY(ph)K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 8085

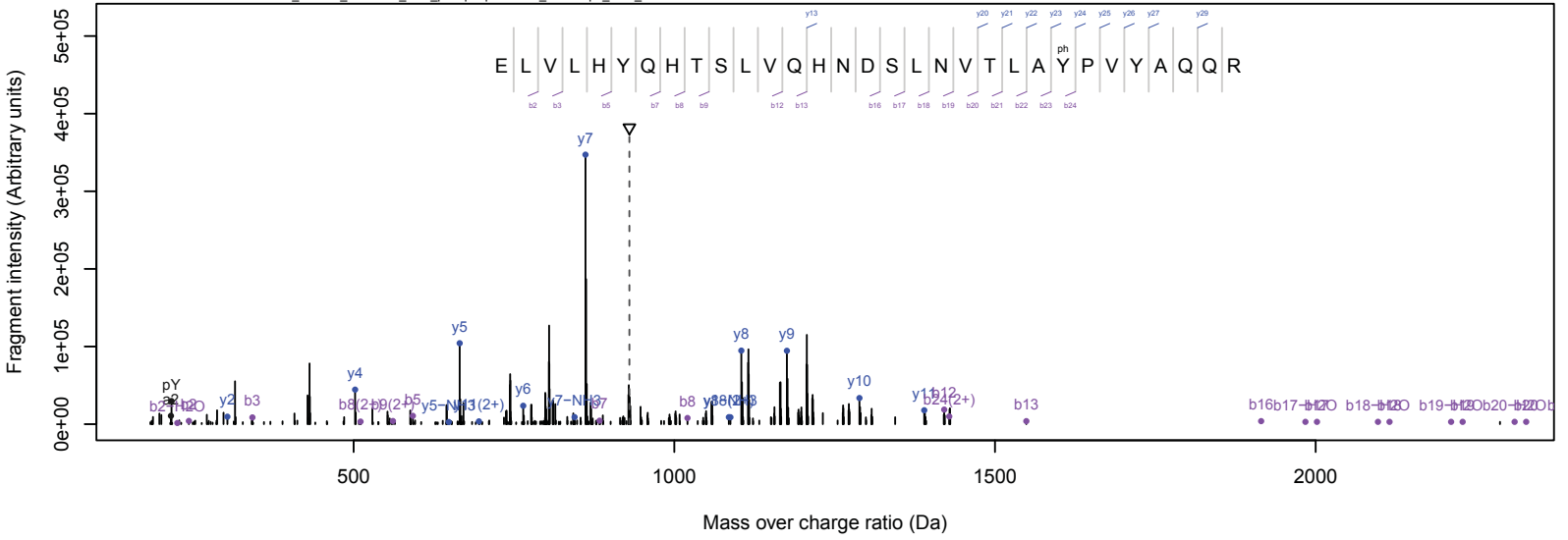
Andromeda score : 115.33



ELVLHYQHTSLVQHNDSLNVTLAY(ph)PVYAQR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37339

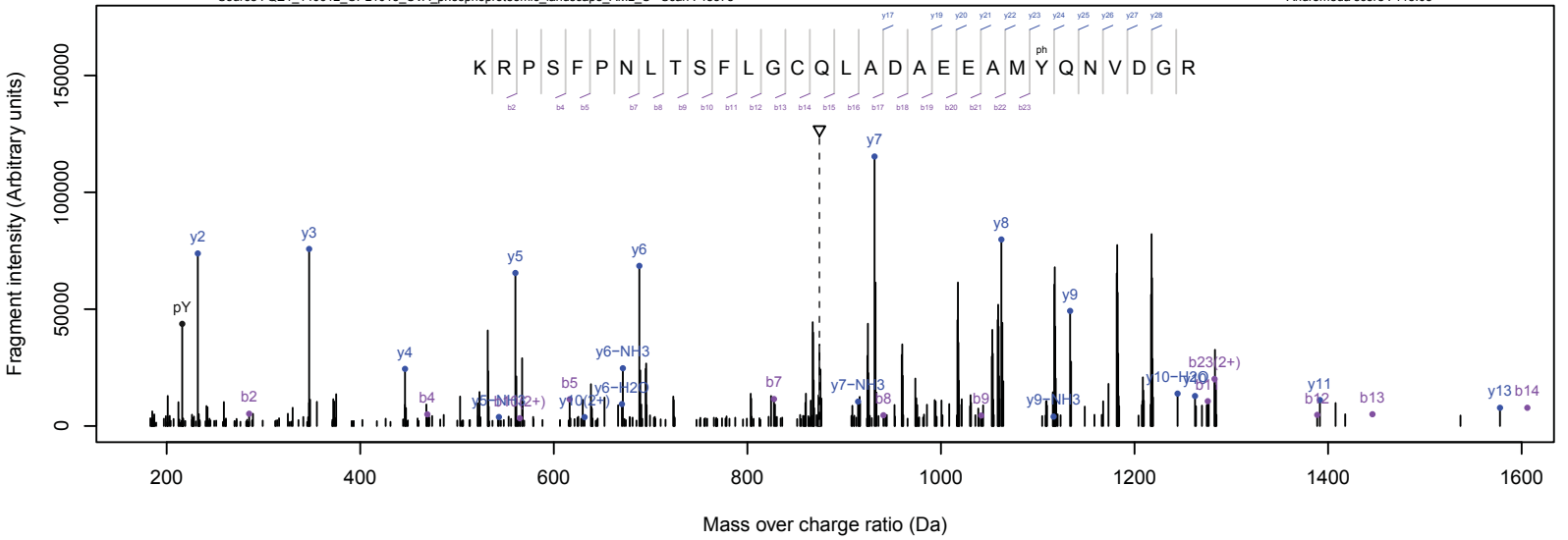
Andromeda score : 115.06



KRPSFPNLTSLGCLADAEEMY(ph)QNV DGR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 48378

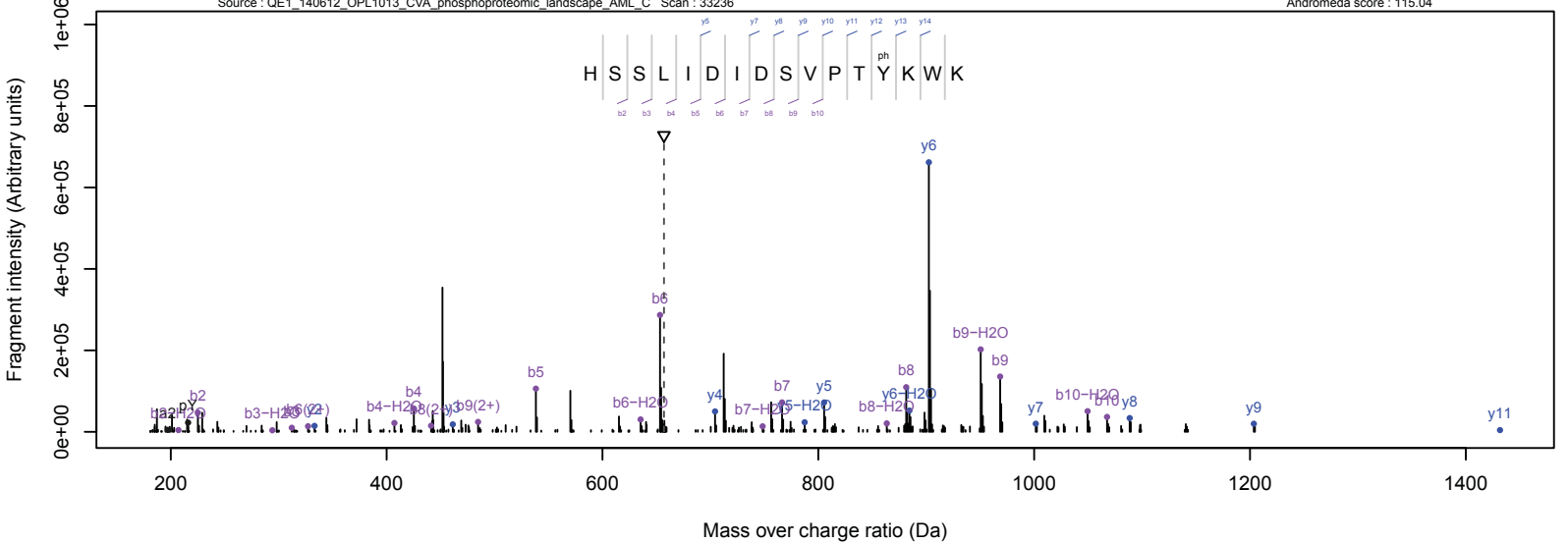
Andromeda score : 115.05



HSSLIDISVPTY(ph)KWK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33236

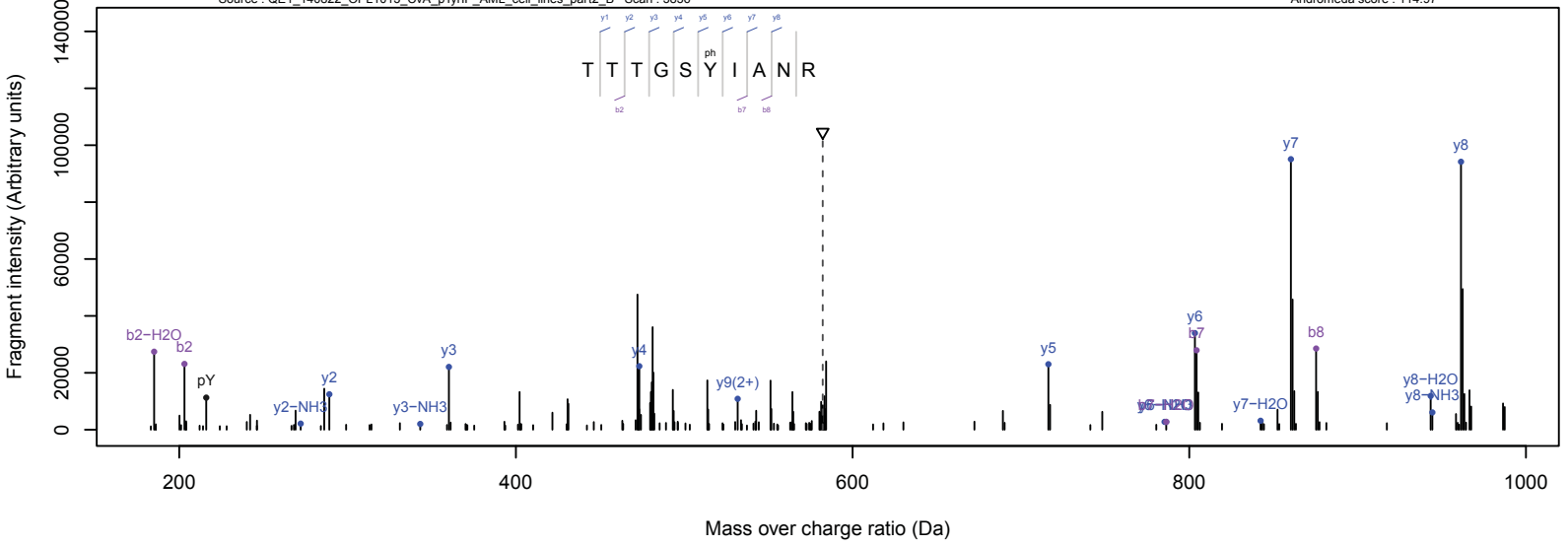
Andromeda score : 115.04



TTTGSY(ph)IANR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3836

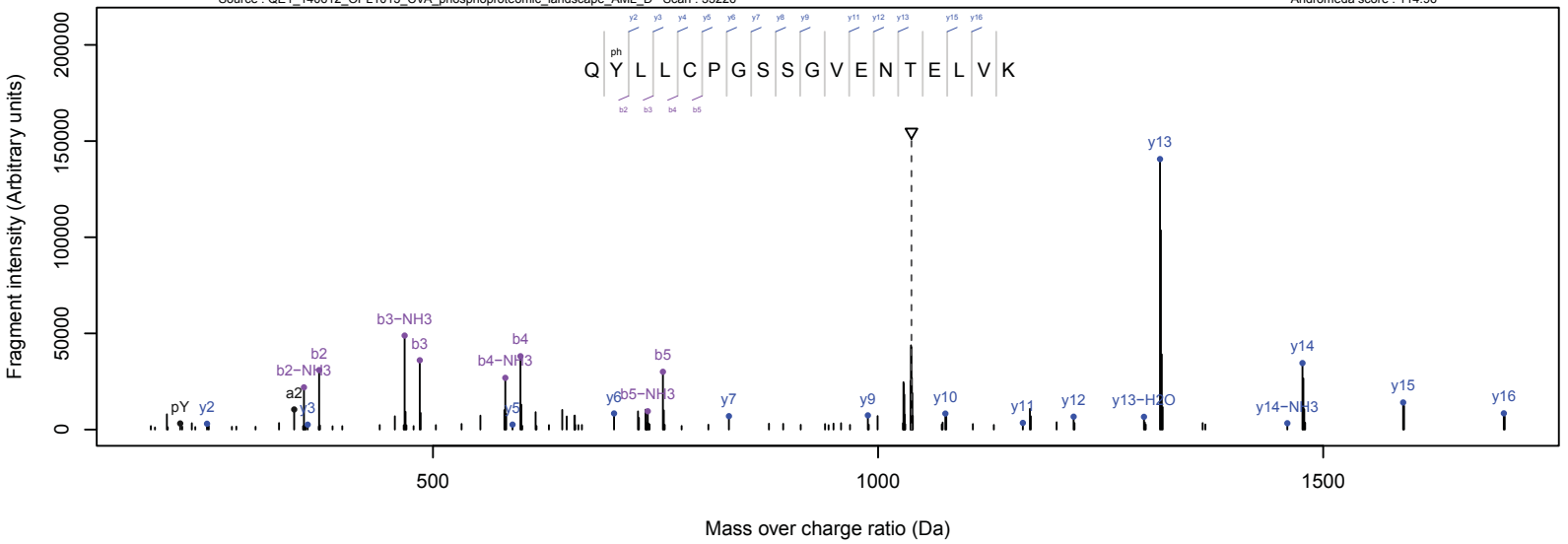
Andromeda score : 114.97



QY(ph)LLCPGSSGVENTELVK-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan : 33226

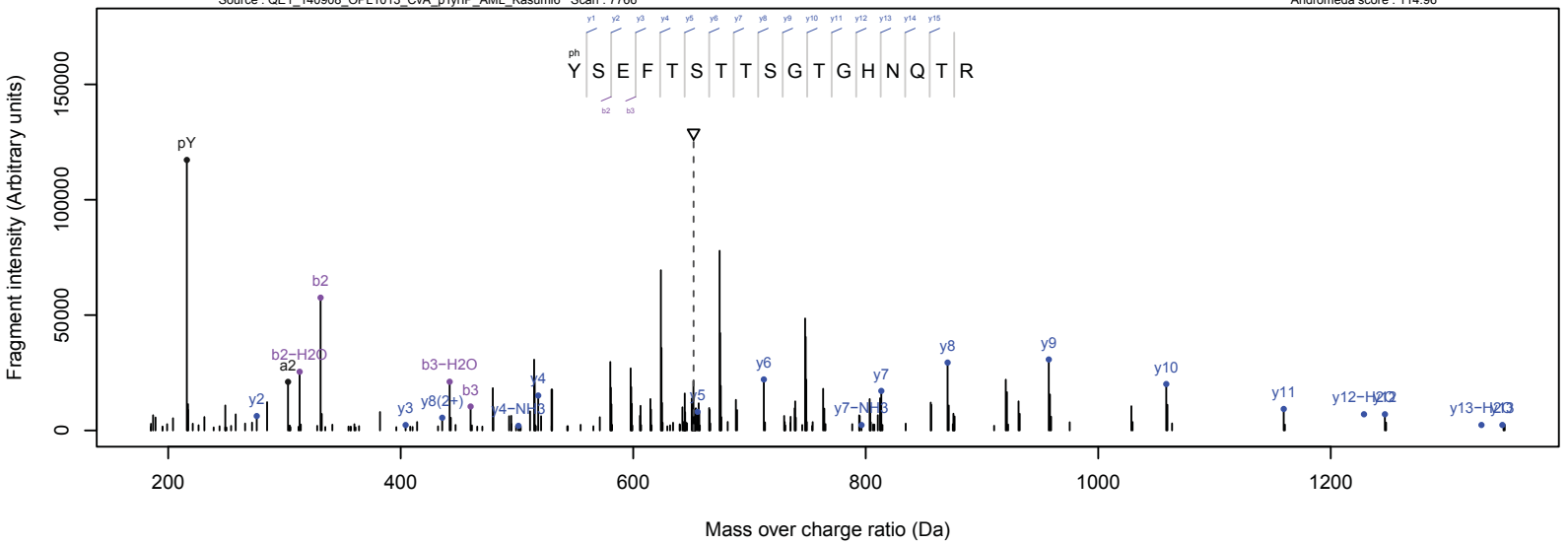
Andromeda score : 114.96



Y(ph)SEFTTSTGTHNQTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 7766

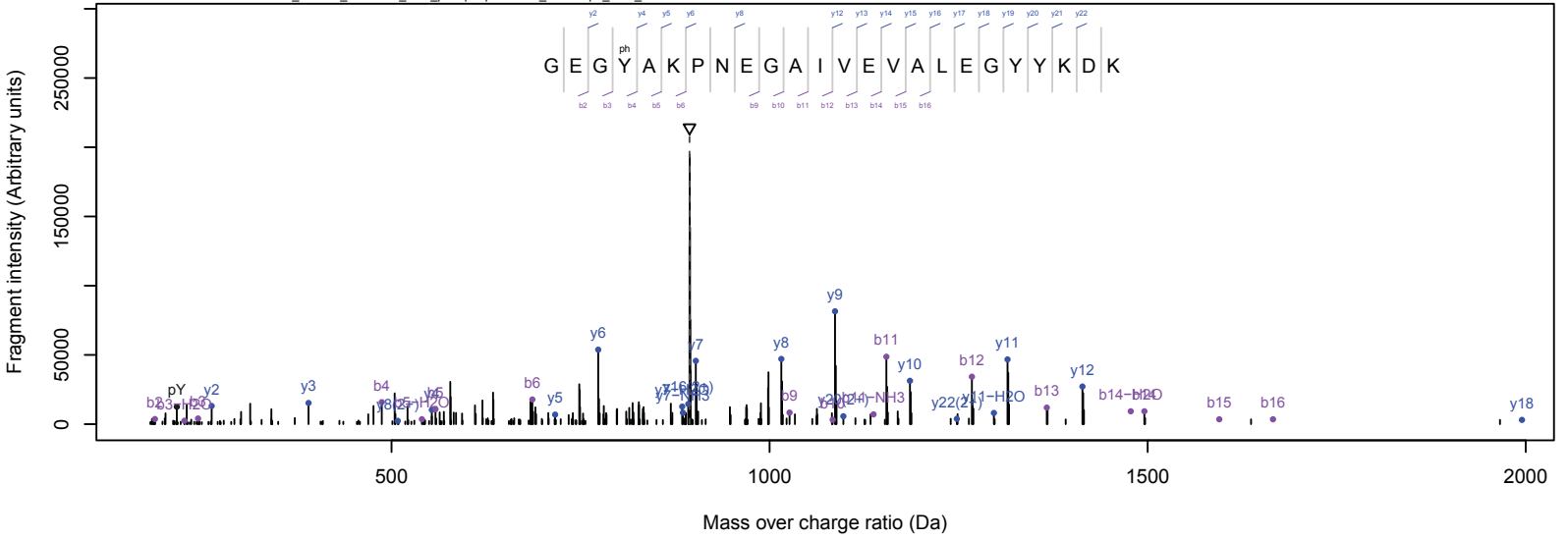
Andromeda score : 114.96



GEGY(ph)AKPNEGAIVEVALEGGYKDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38254

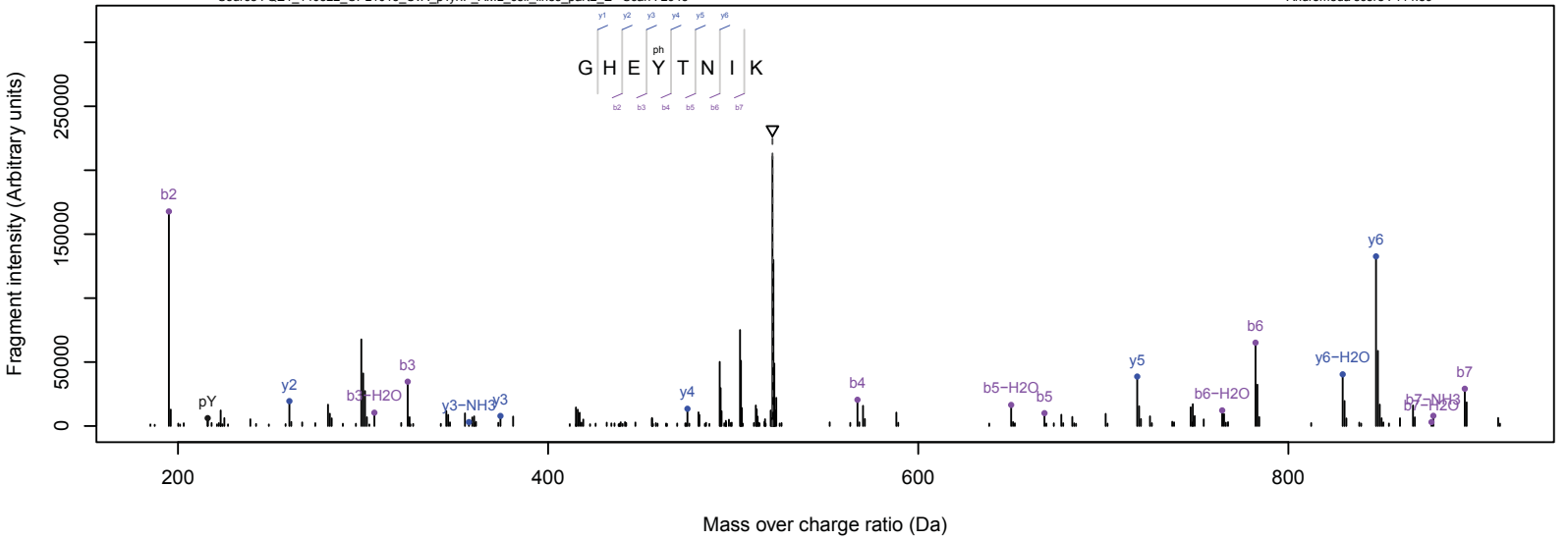
Andromeda score : 114.9



GHEY(ph)TNIK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 2943

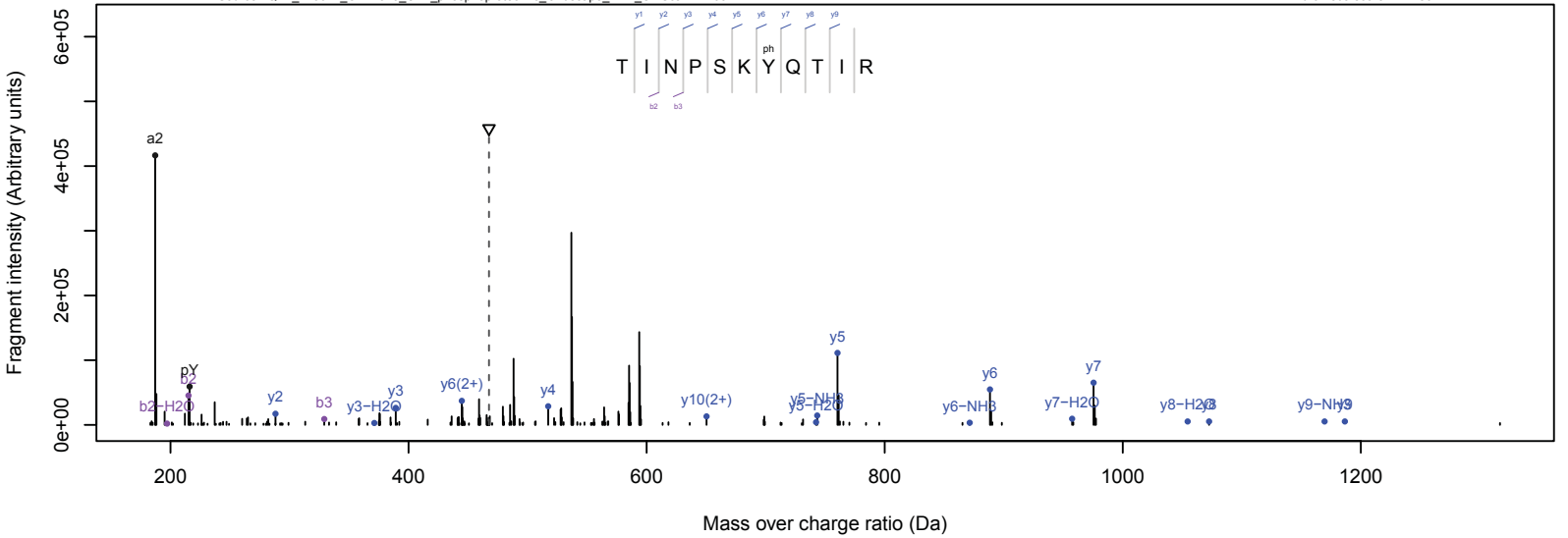
Andromeda score : 114.89



TINPSKY(ph)QTIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11032

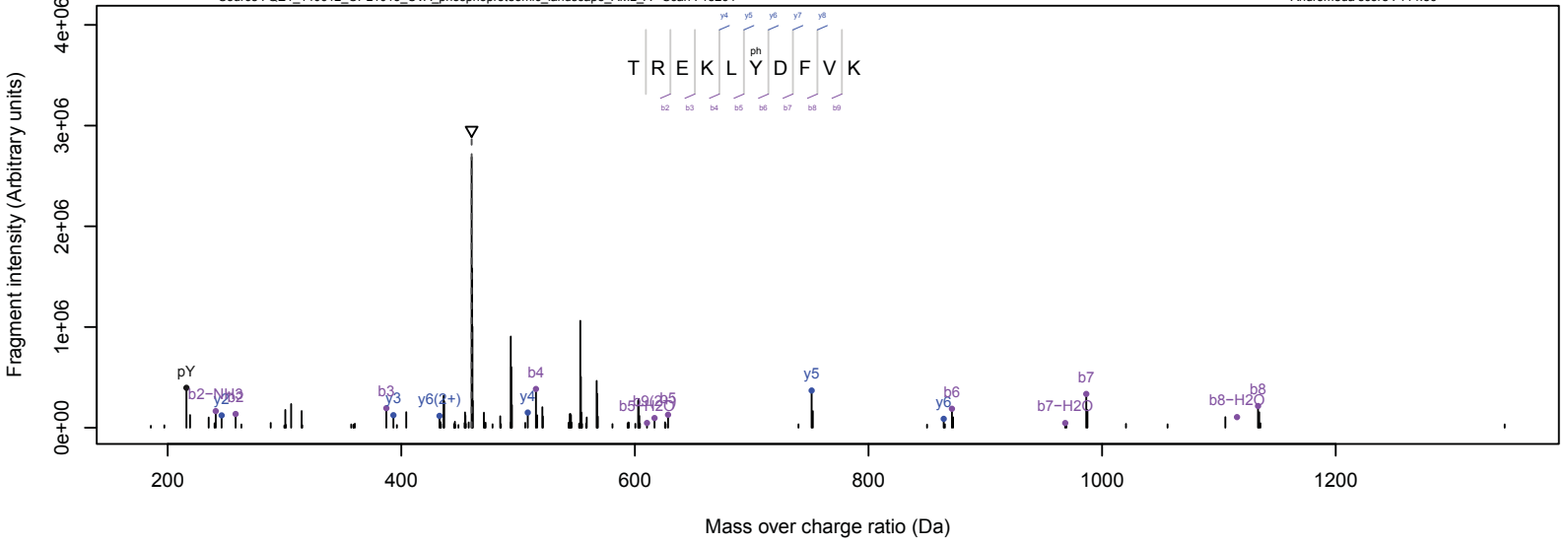
Andromeda score : 114.86



TREKLY(ph)DFVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 18264

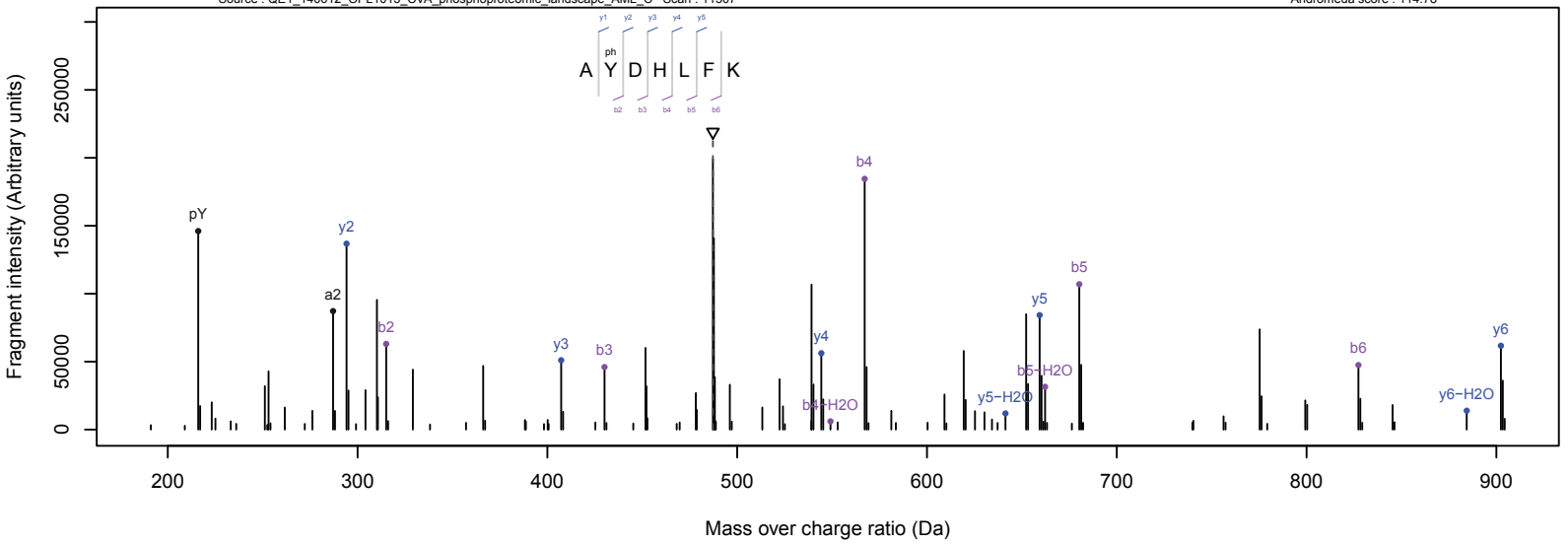
Andromeda score : 114.86



AY(ph)DHLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11507

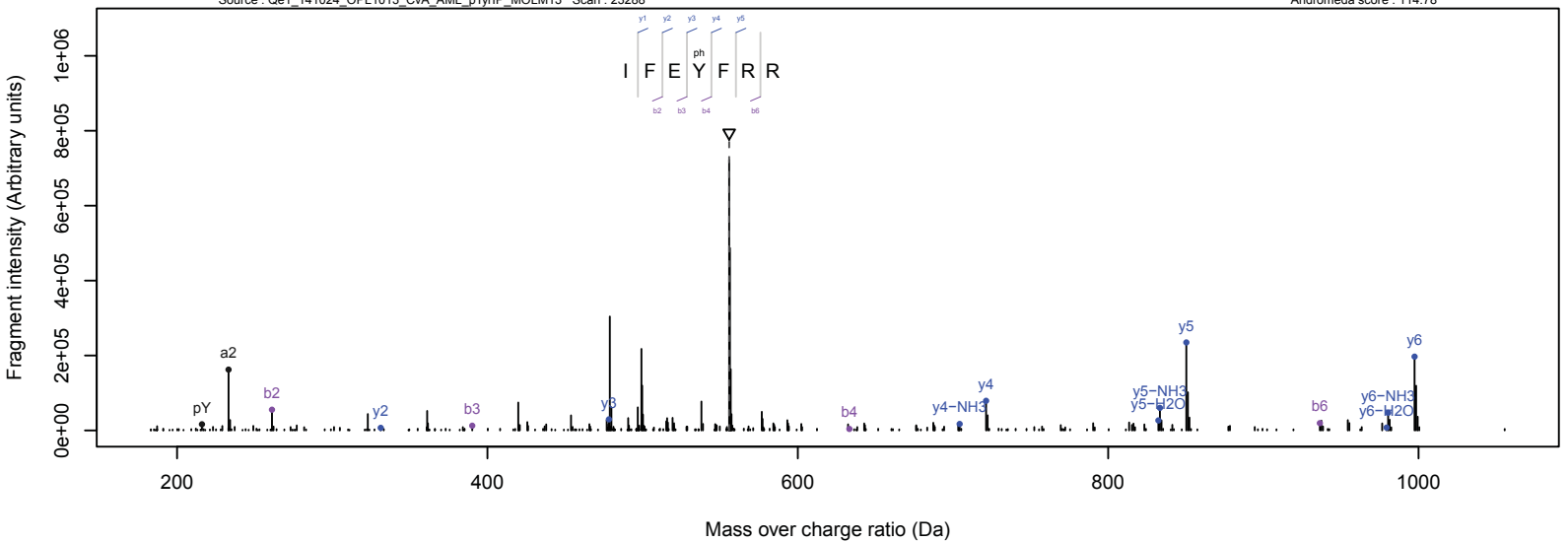
Andromeda score : 114.78



IFEY(ph)FRR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 23288

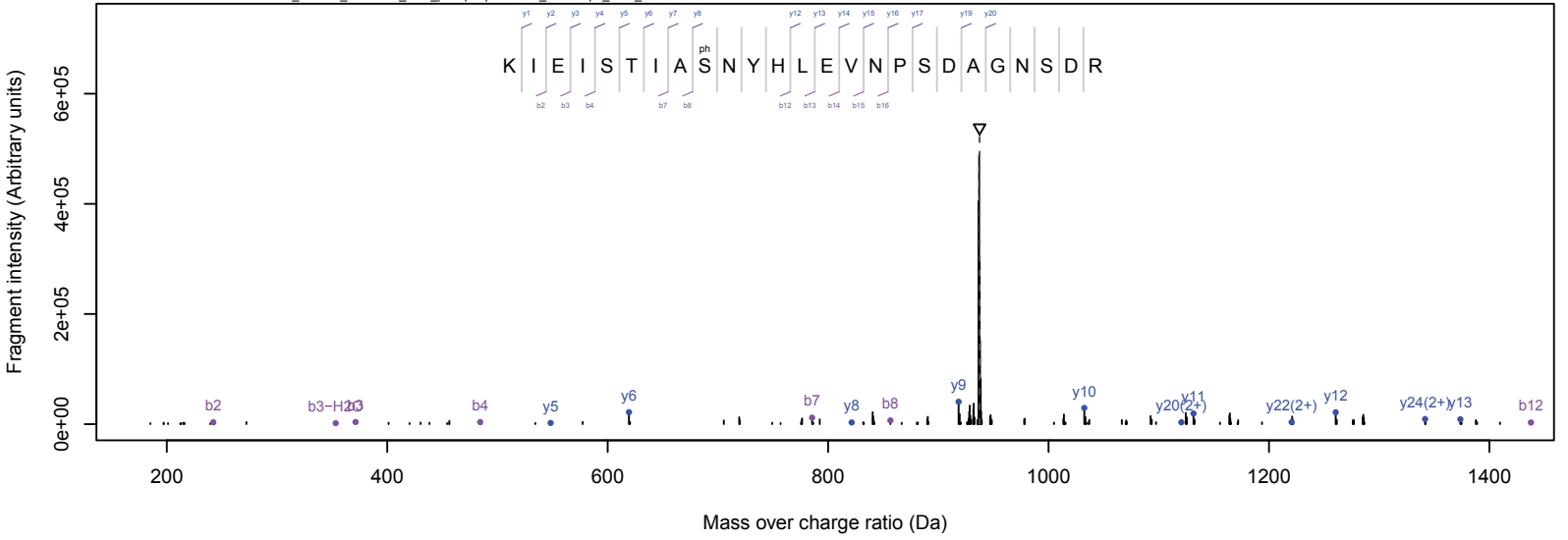
Andromeda score : 114.78



**KIEISTIAS(ph)NYHLEVNPSDAGNSDR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29777

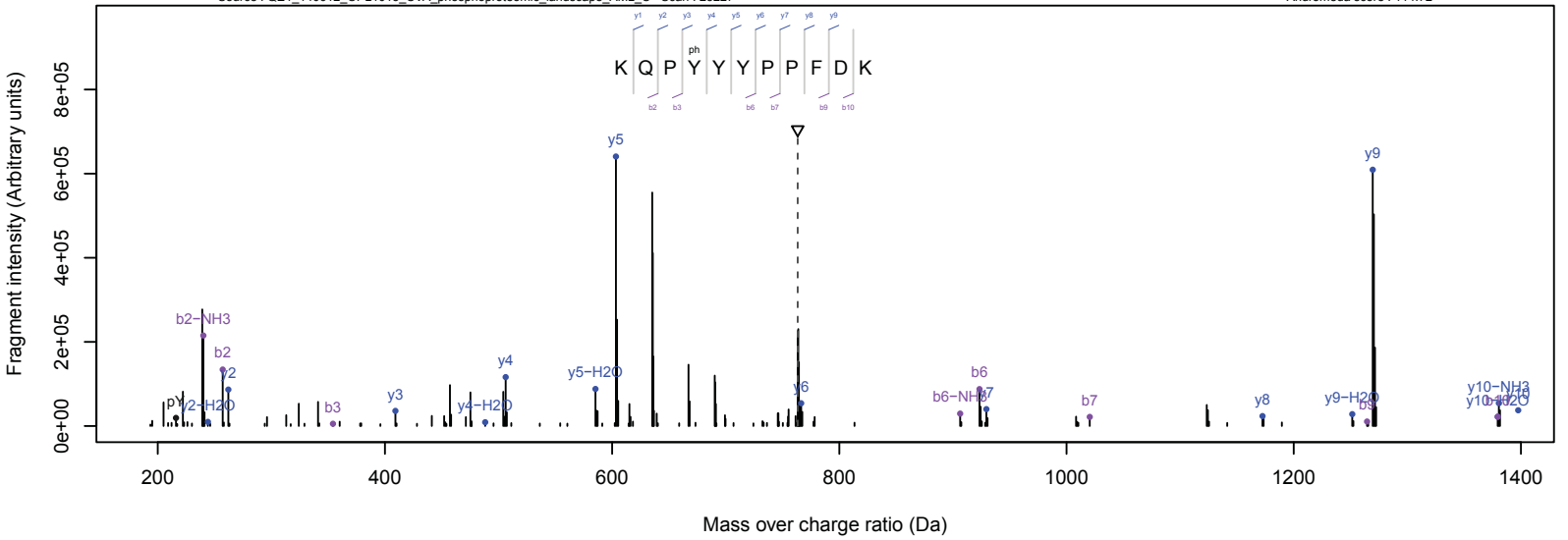
Andromeda score : 114.74



**KQPY(ph)YPPFDK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23227

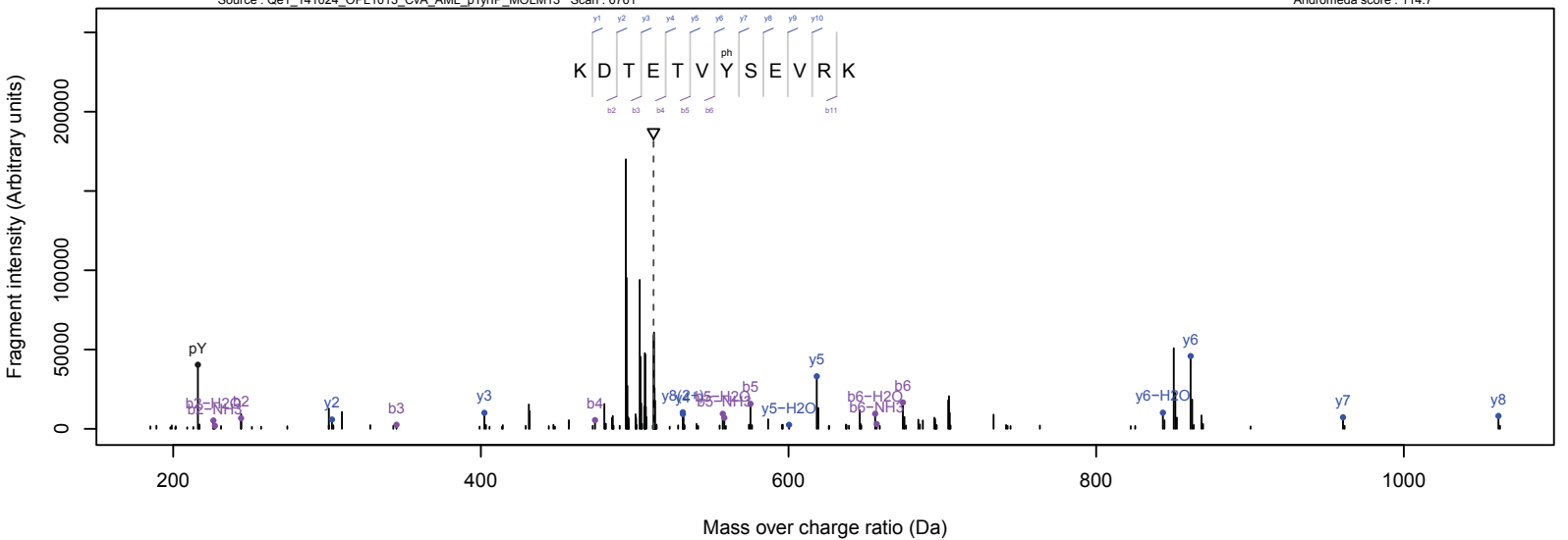
Andromeda score : 114.72



**KDTETVY(ph)SEVRK-3+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 6761

Andromeda score : 114.7

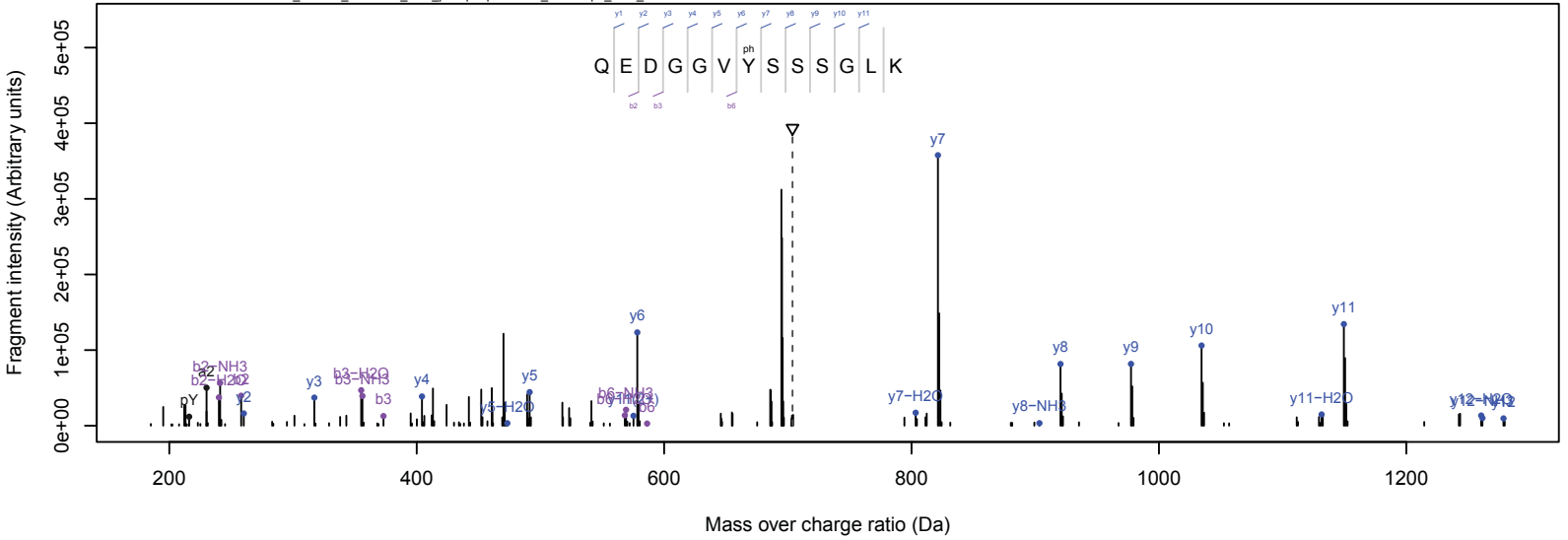




**QEDGGVY(ph)SSGLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 6903

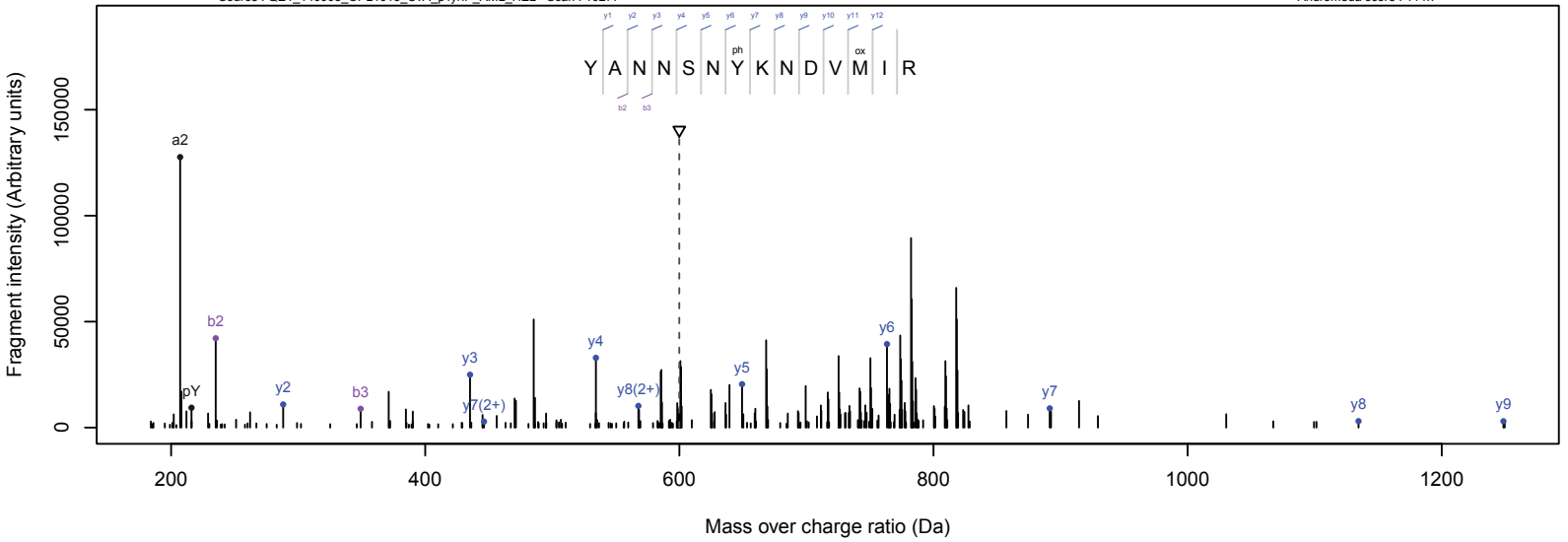
Andromeda score : 114.7



**YANNSNY(ph)KNDVM(ox)IR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 10277

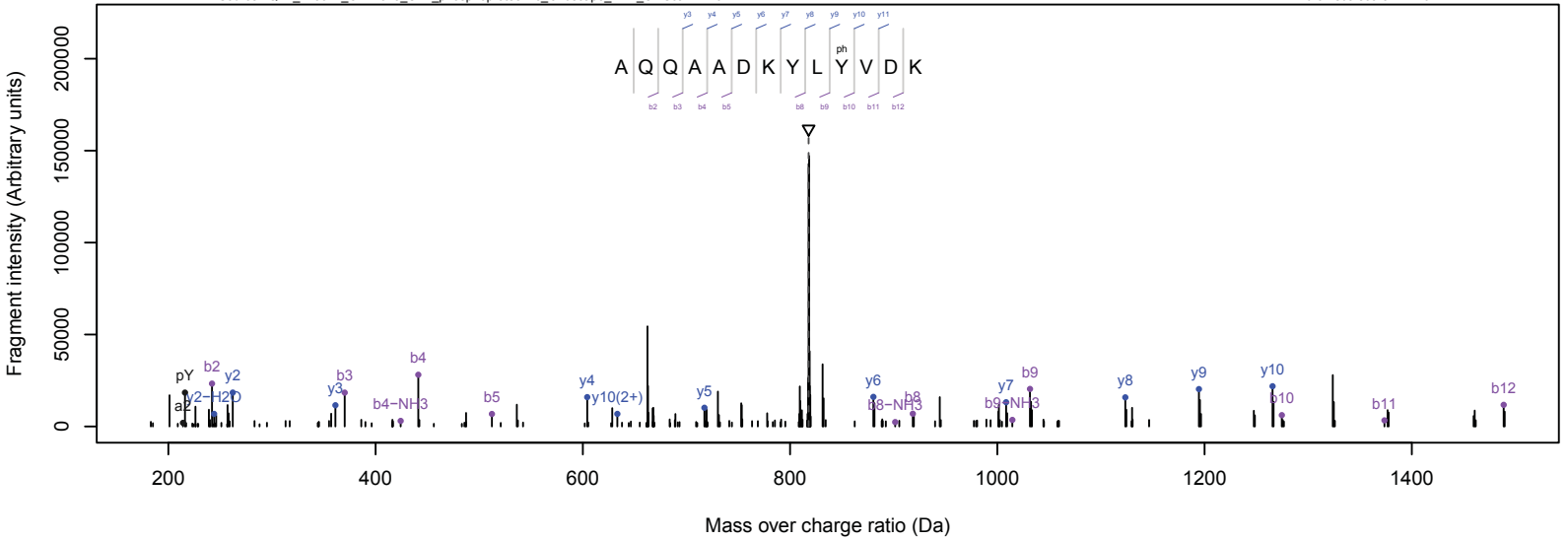
Andromeda score : 114.7



**(ac)AQQAADKYL(ph)VDK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27072

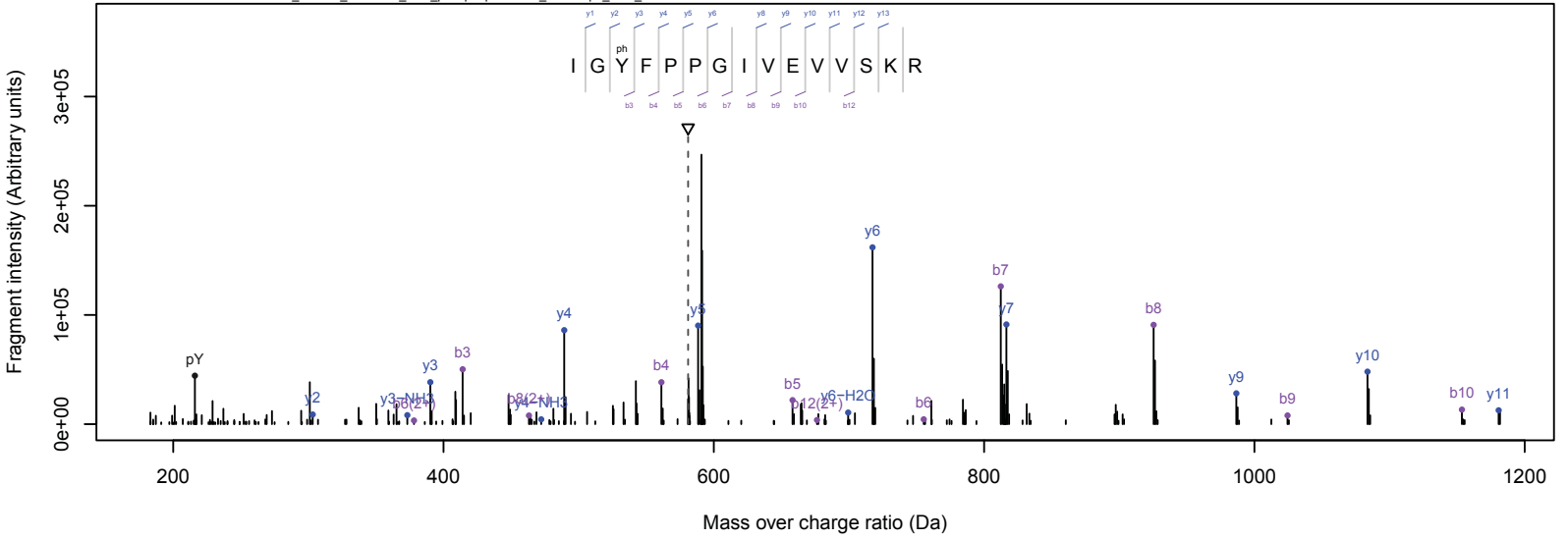
Andromeda score : 114.64



IGY(ph)FPPGIVEVSKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41916

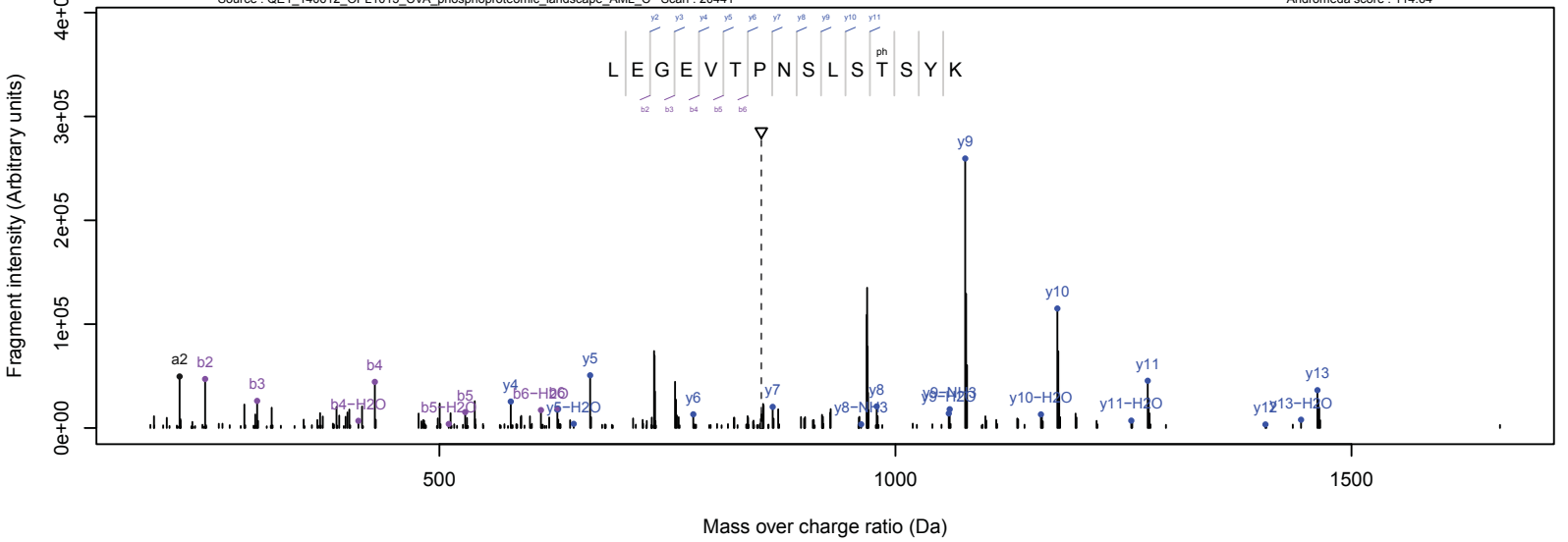
Andromeda score : 114.64



LEGEVTPNSLST(ph)SYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20441

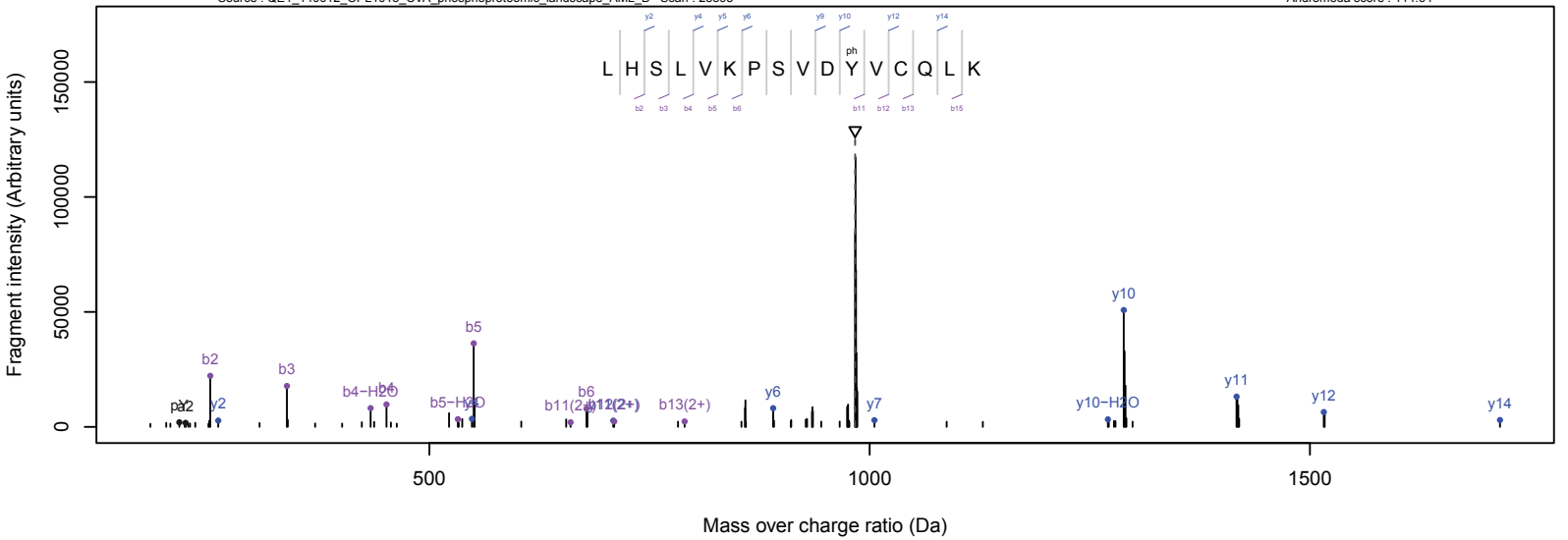
Andromeda score : 114.64



LHSLVKPSVDY(ph)VCQLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25893

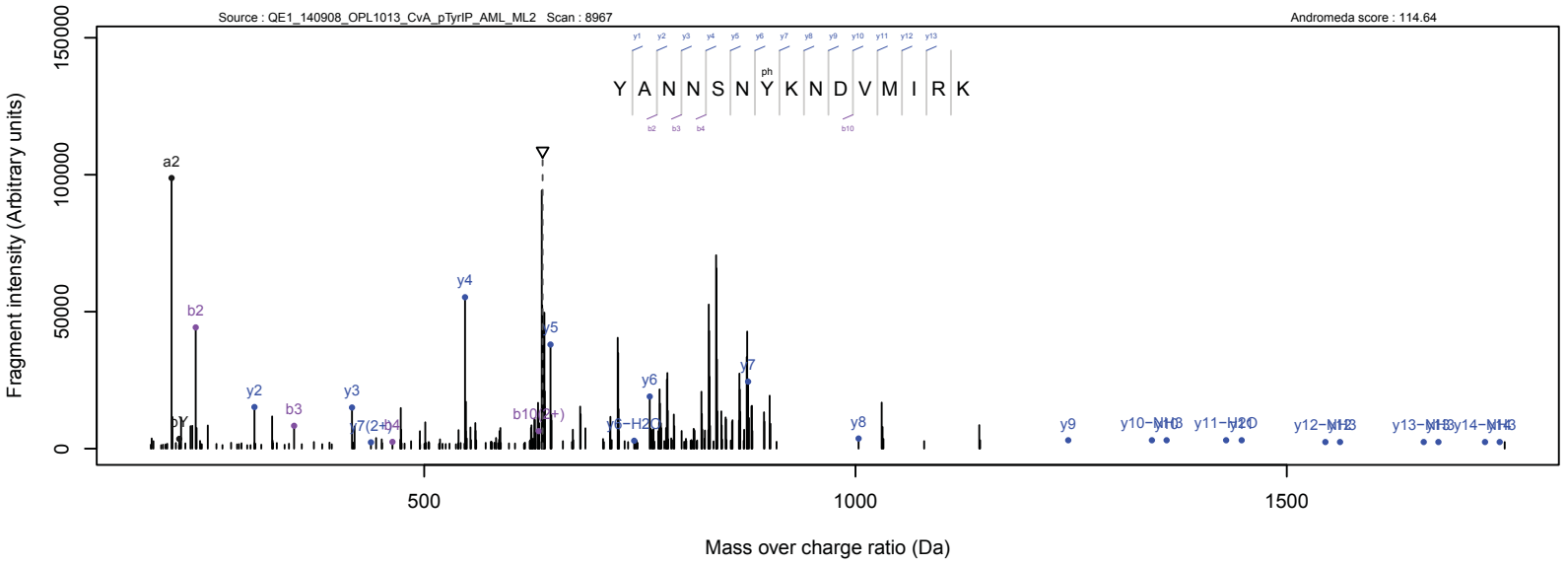
Andromeda score : 114.64



**YANNSNY(ph)KNDVMIRK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 8967

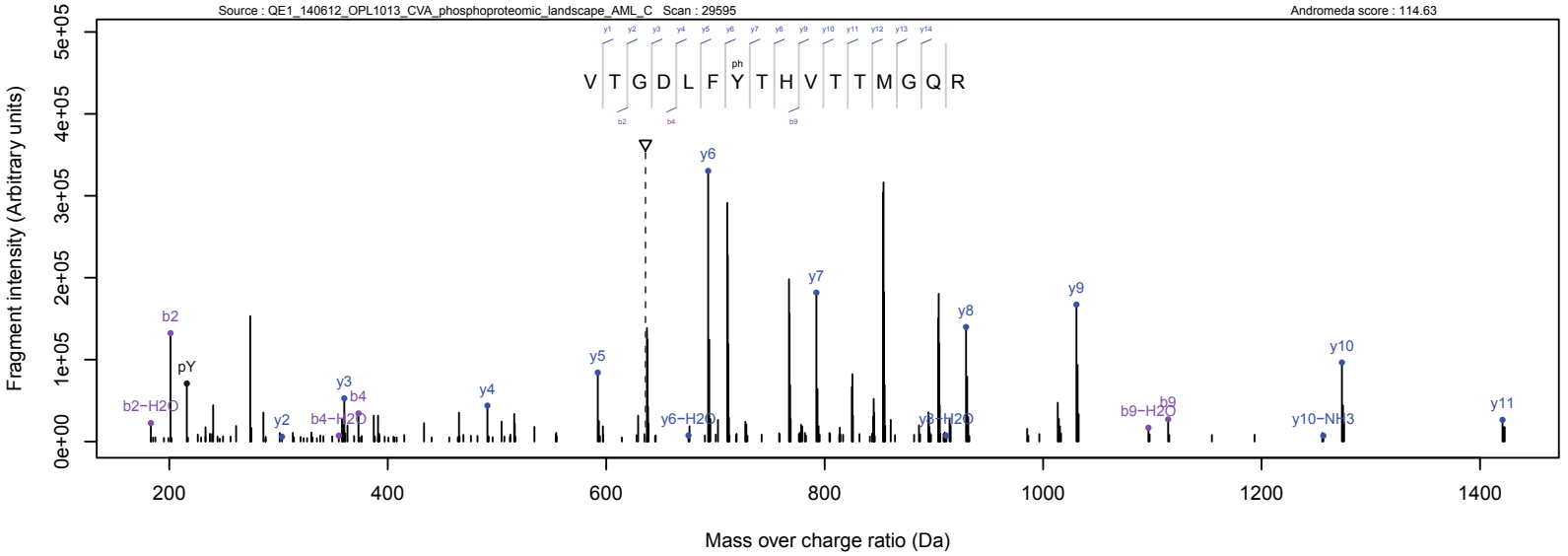
Andromeda score : 114.64



**VTGDLFY(ph)THVTTMGQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29595

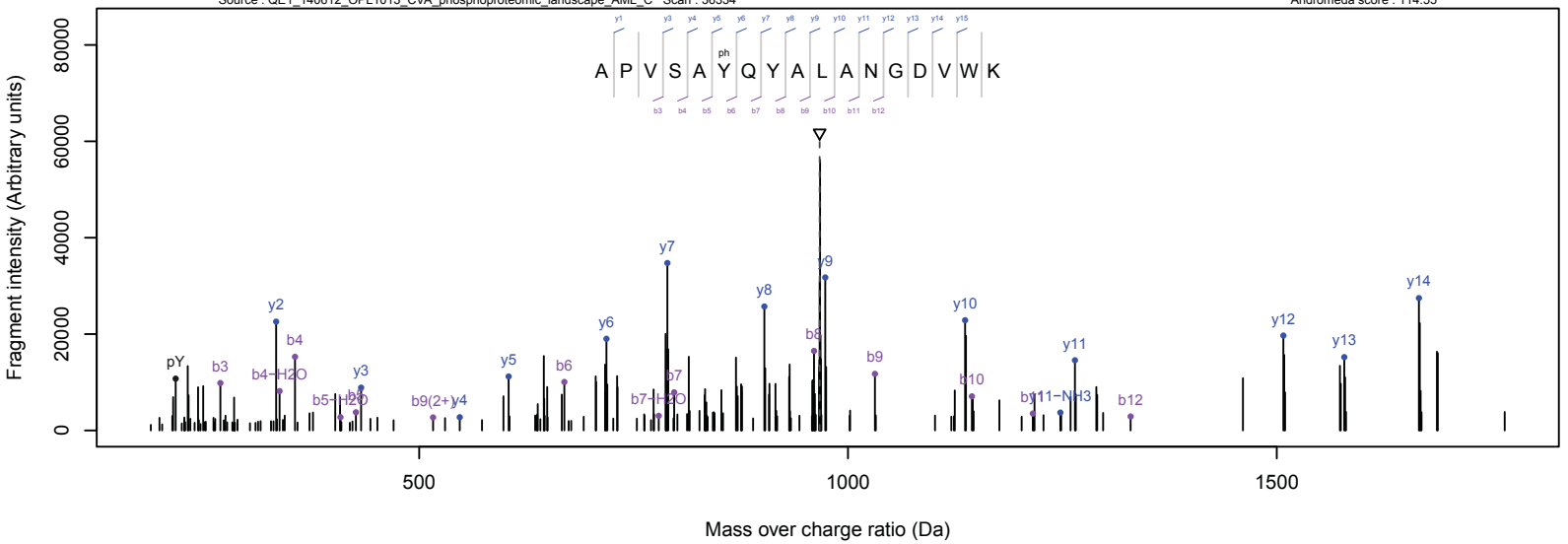
Andromeda score : 114.63



**APVSAY(ph)QYALANGDVWK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36334

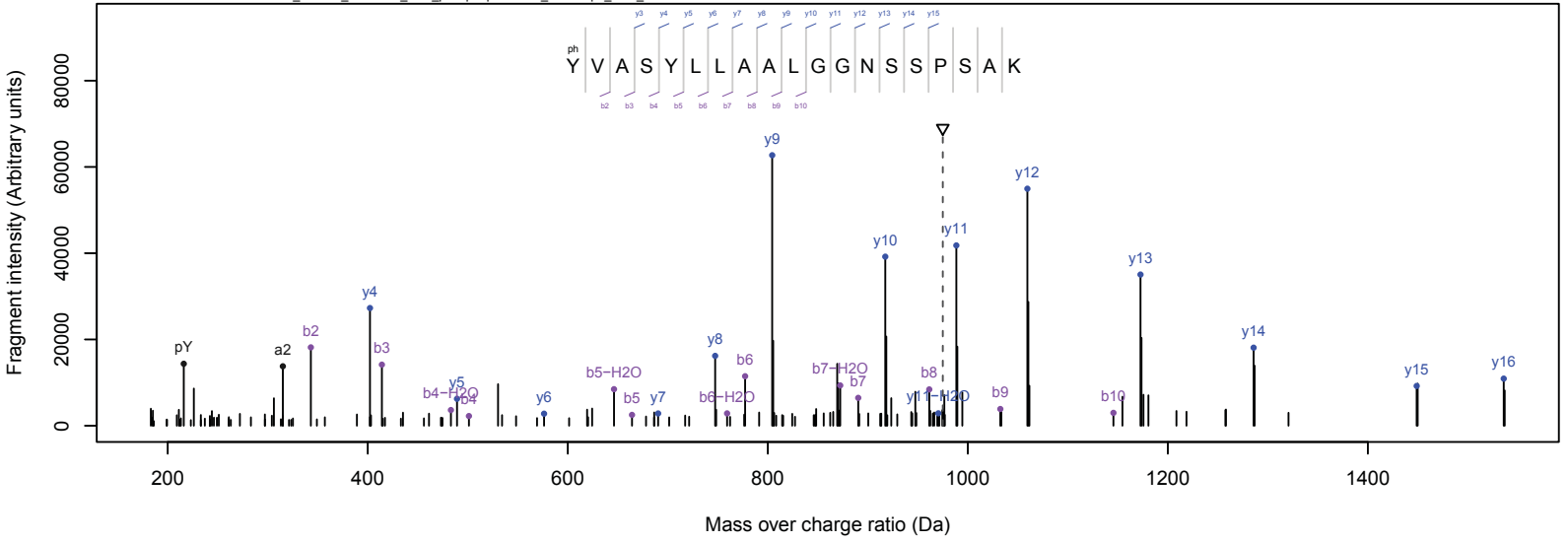
Andromeda score : 114.55



Y(ph)VASYLLAALGGNSSPSAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 46359

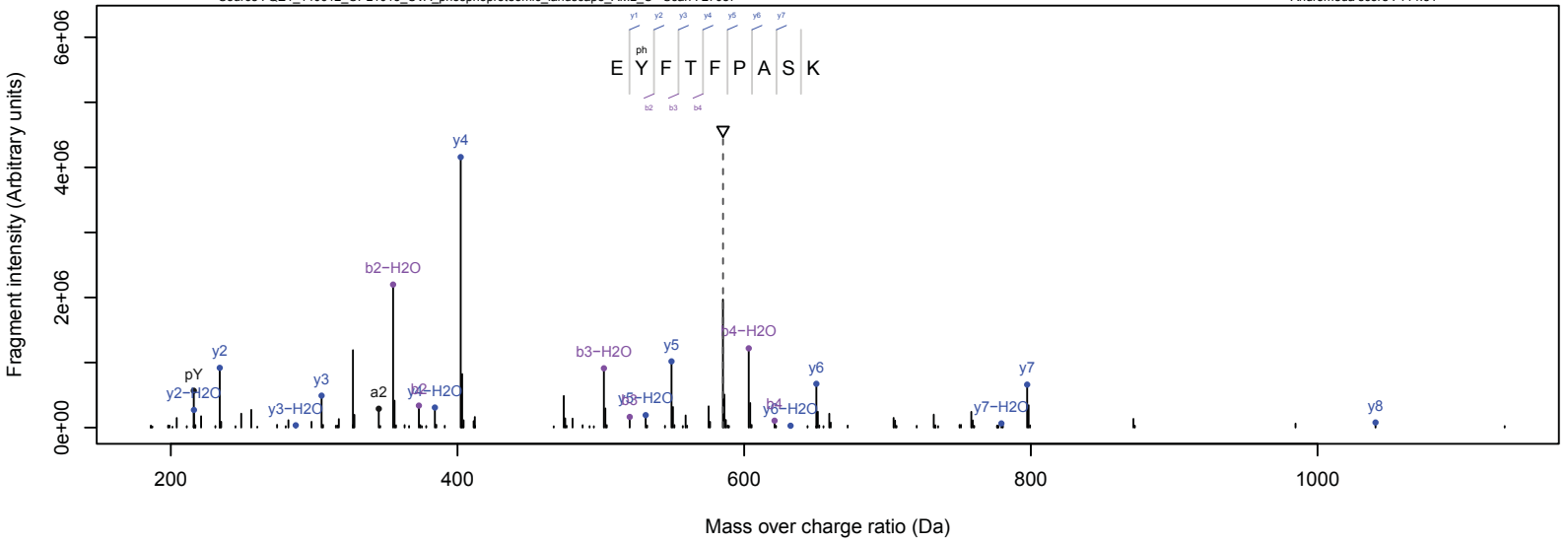
Andromeda score: 114.55



EY(ph)FTFPASK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 27087

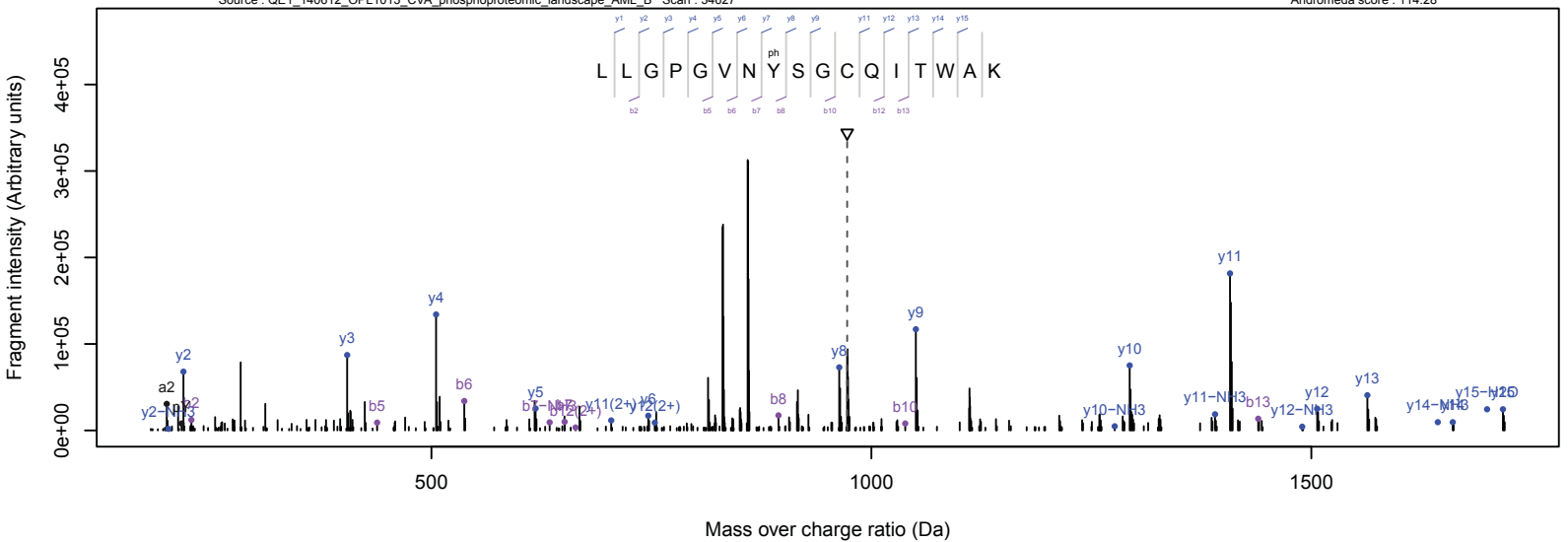
Andromeda score: 114.51



LLGPGVNY(ph)SGCQITWAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34627

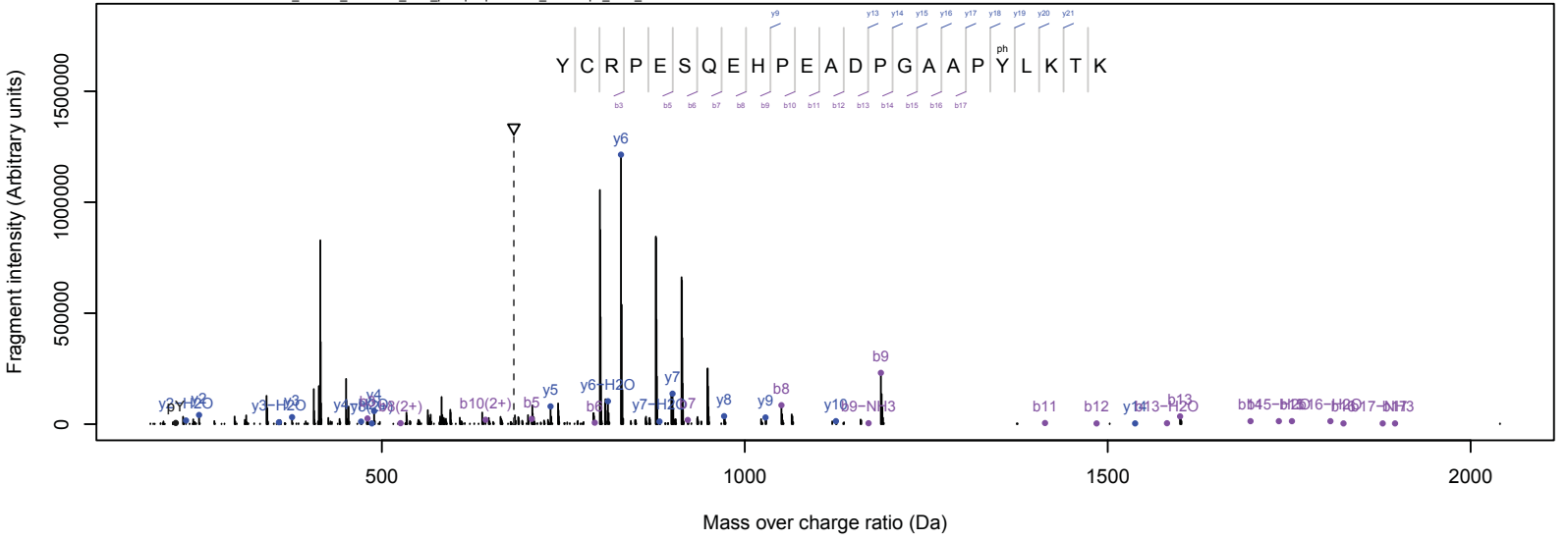
Andromeda score: 114.28



YCRPESQEHPEADPGAAPY(ph)LKTK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 13618

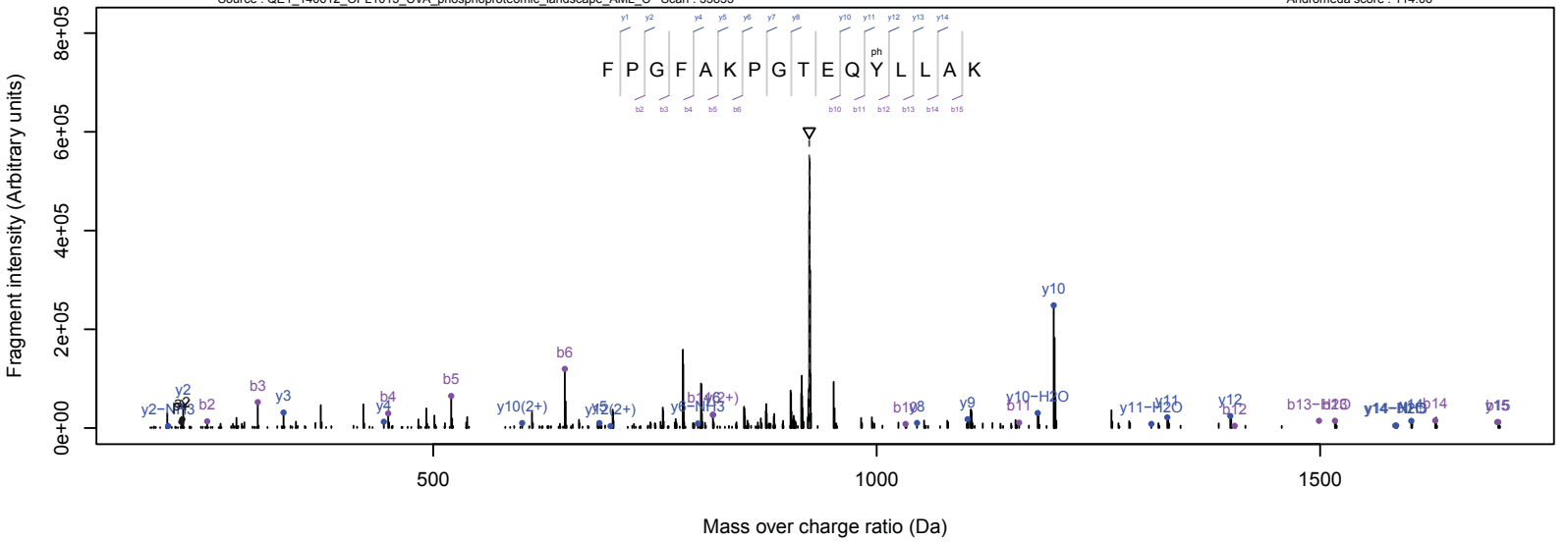
Andromeda score: 114.15



FPGFAKPGTEQY(ph)LLAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 33853

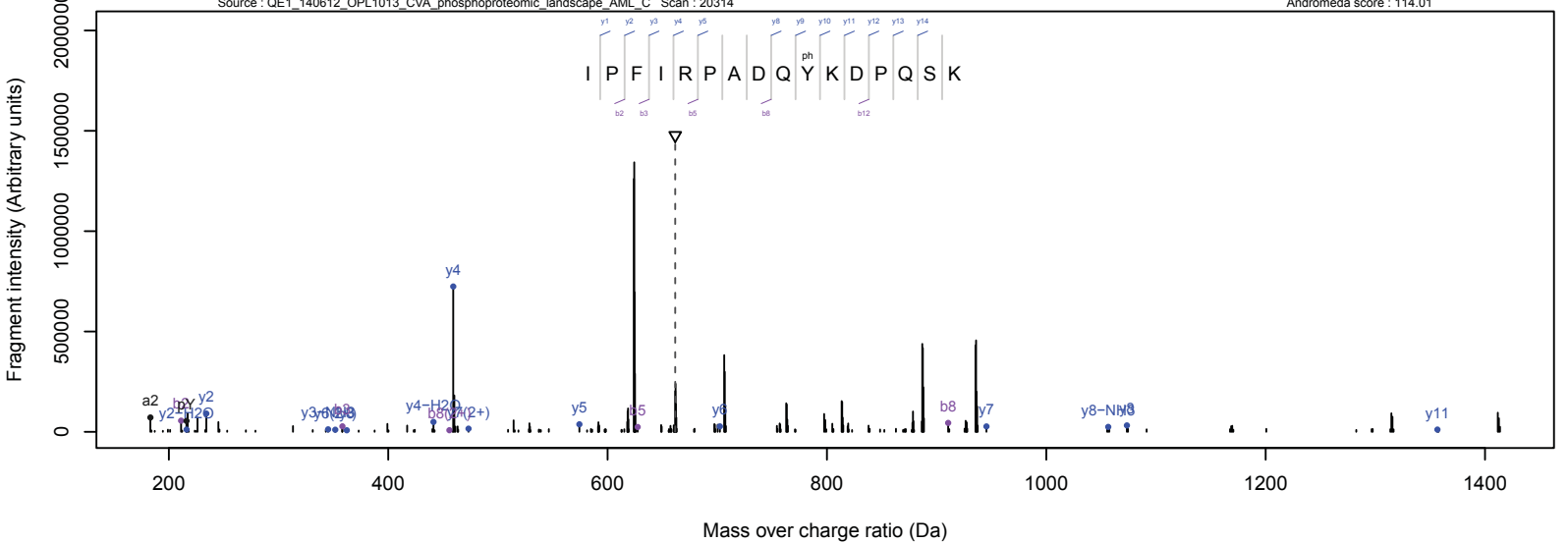
Andromeda score: 114.06



IPFIRPADQYKDPQSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 20314

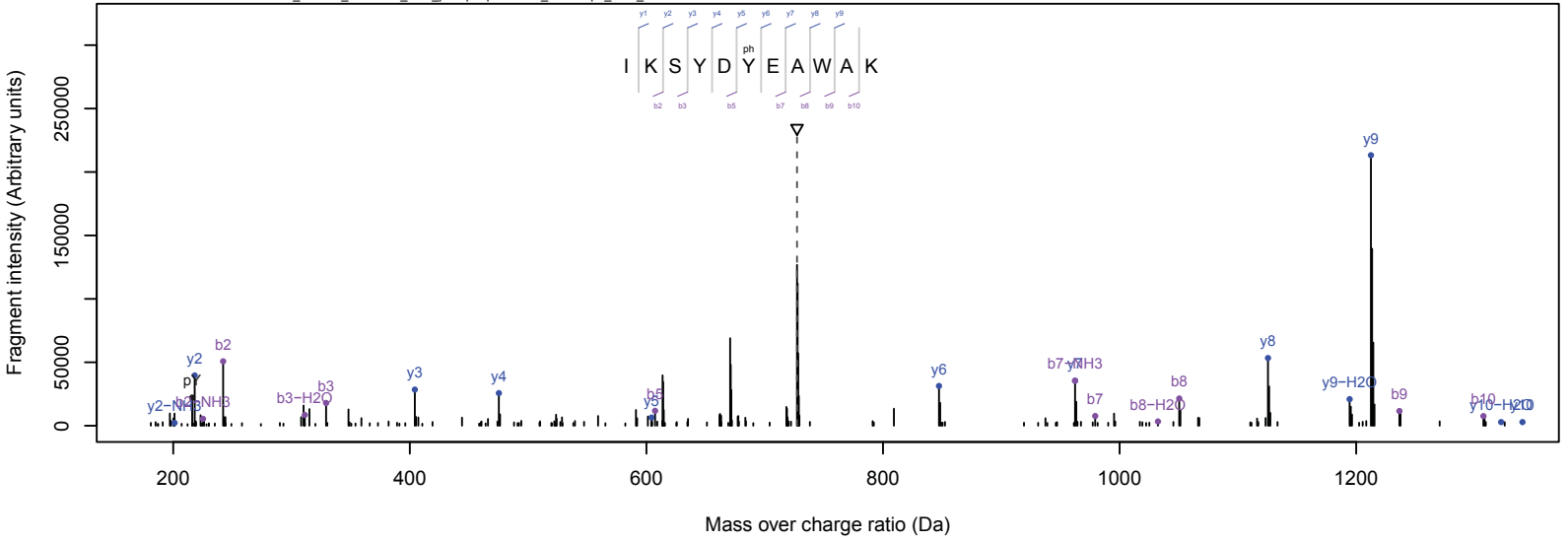
Andromeda score: 114.01



**IKSYDY(ph)EAWAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18056

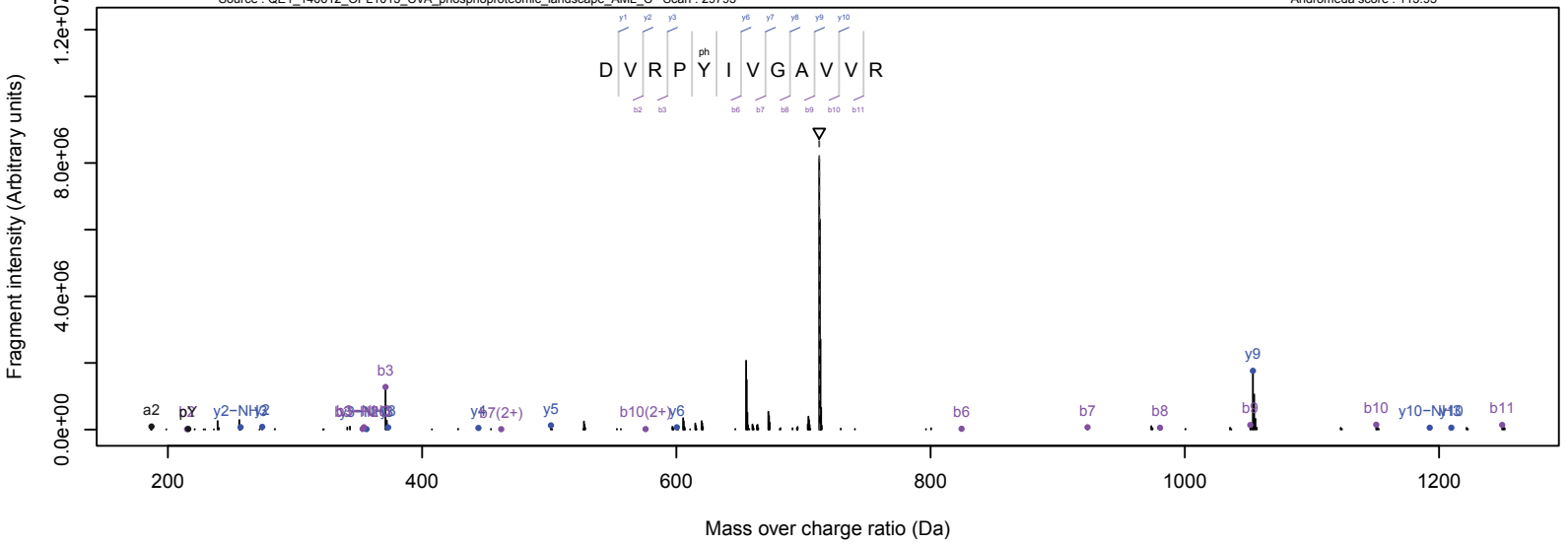
Andromeda score : 113.99



**DVRPY(ph)IVGAVR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29793

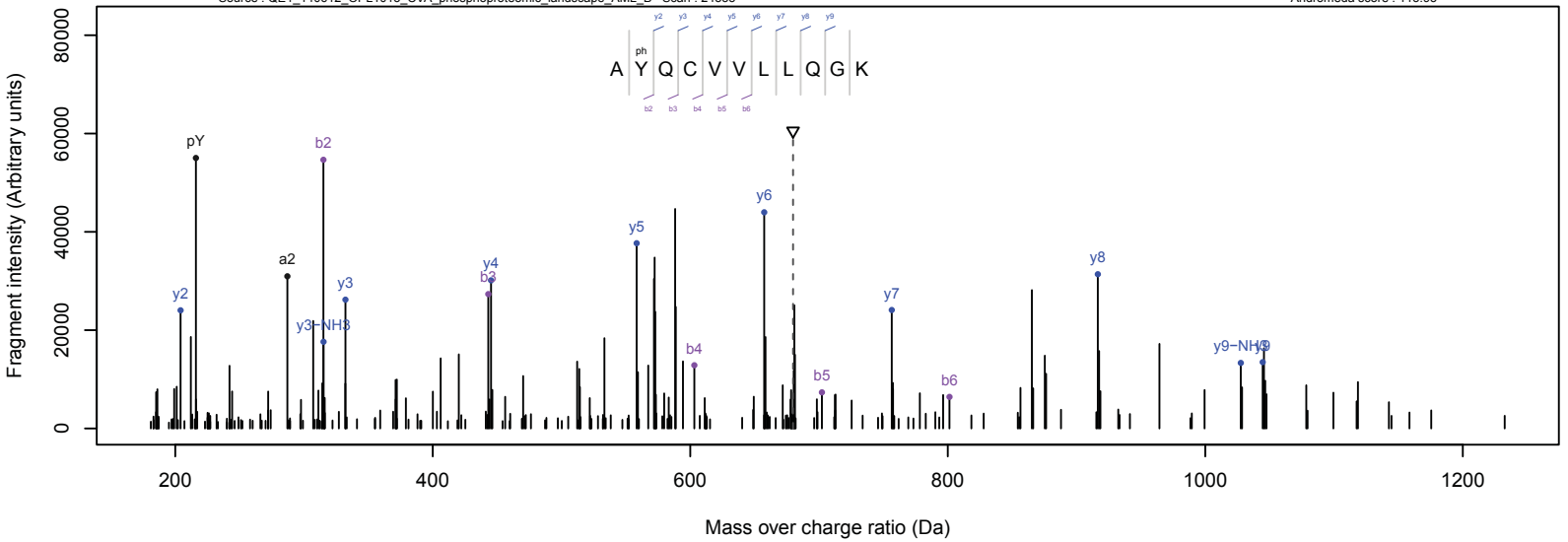
Andromeda score : 113.95



**AY(ph)QCVLLQ GK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24335

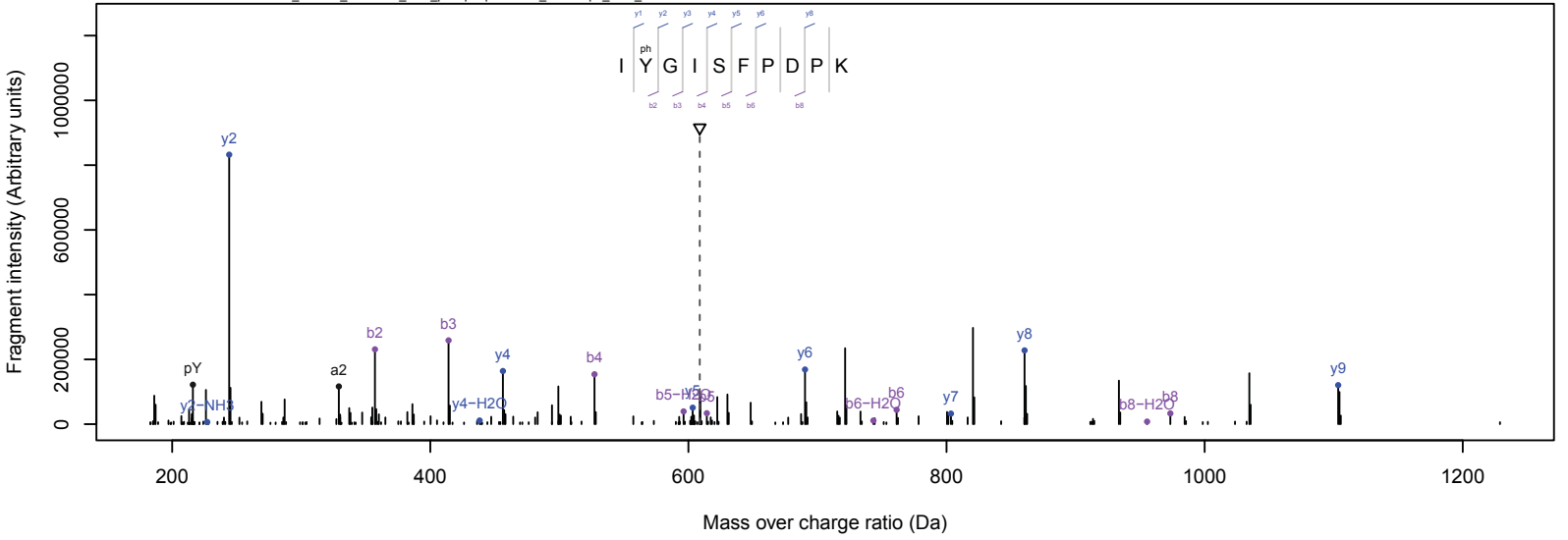
Andromeda score : 113.93



IY(ph)GISFPDPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33183

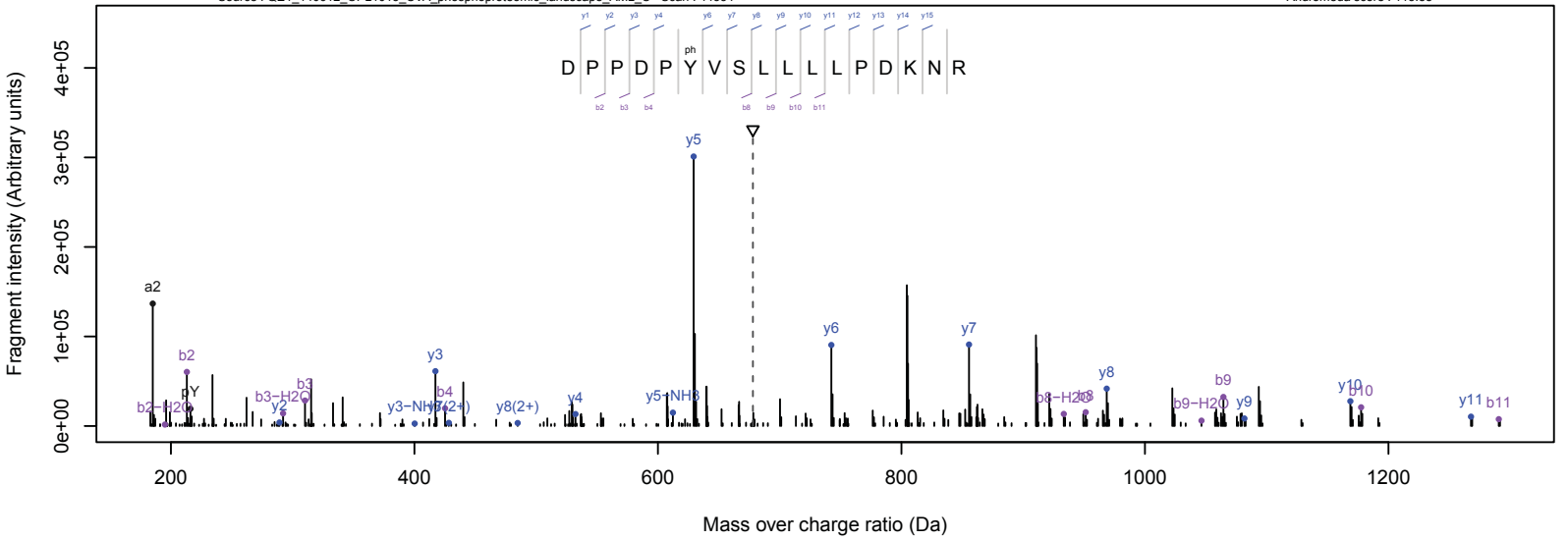
Andromeda score : 113.93



DPPDPY(ph)VSLLLPDKNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44334

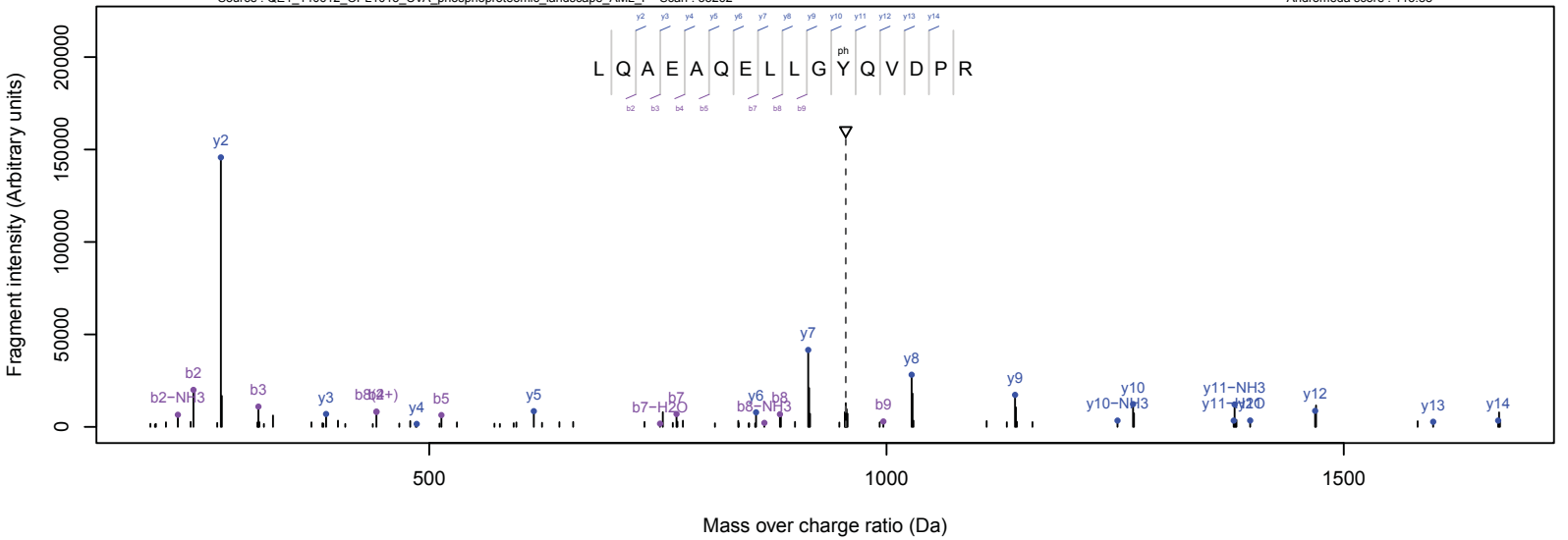
Andromeda score : 113.88



LQAEAQELLY(ph)QVDPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 33262

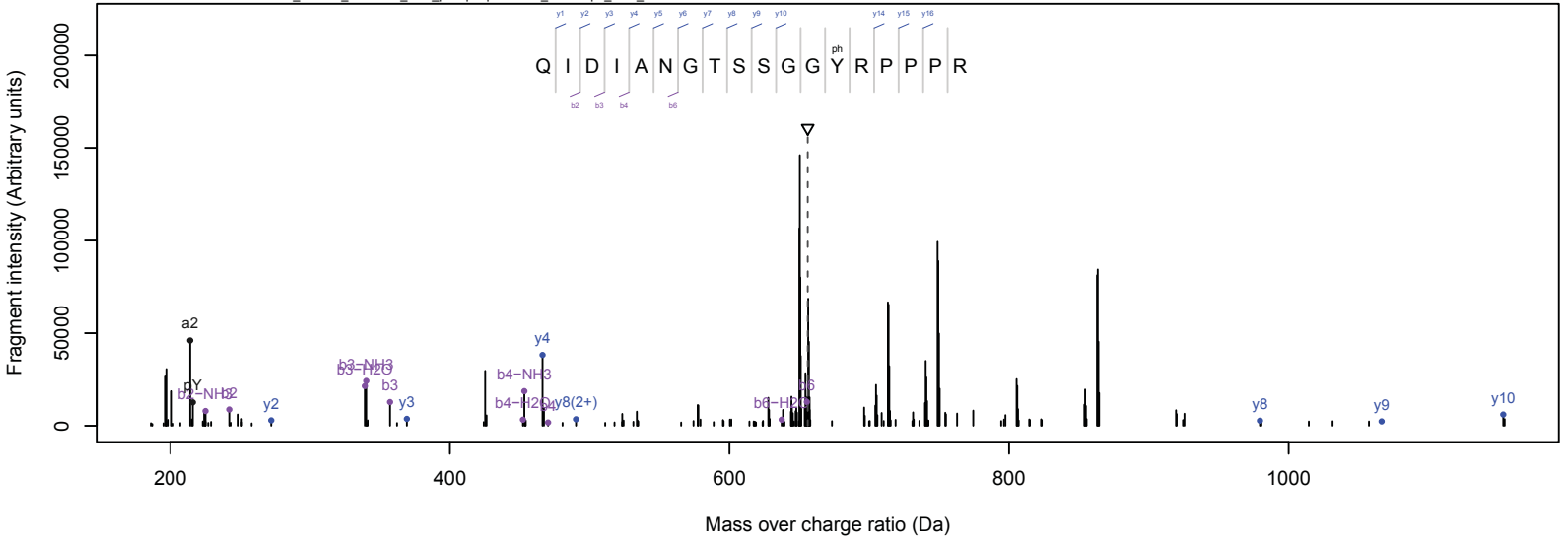
Andromeda score : 113.88



**QIDIANGTSSGGY(ph)RPPPR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14063

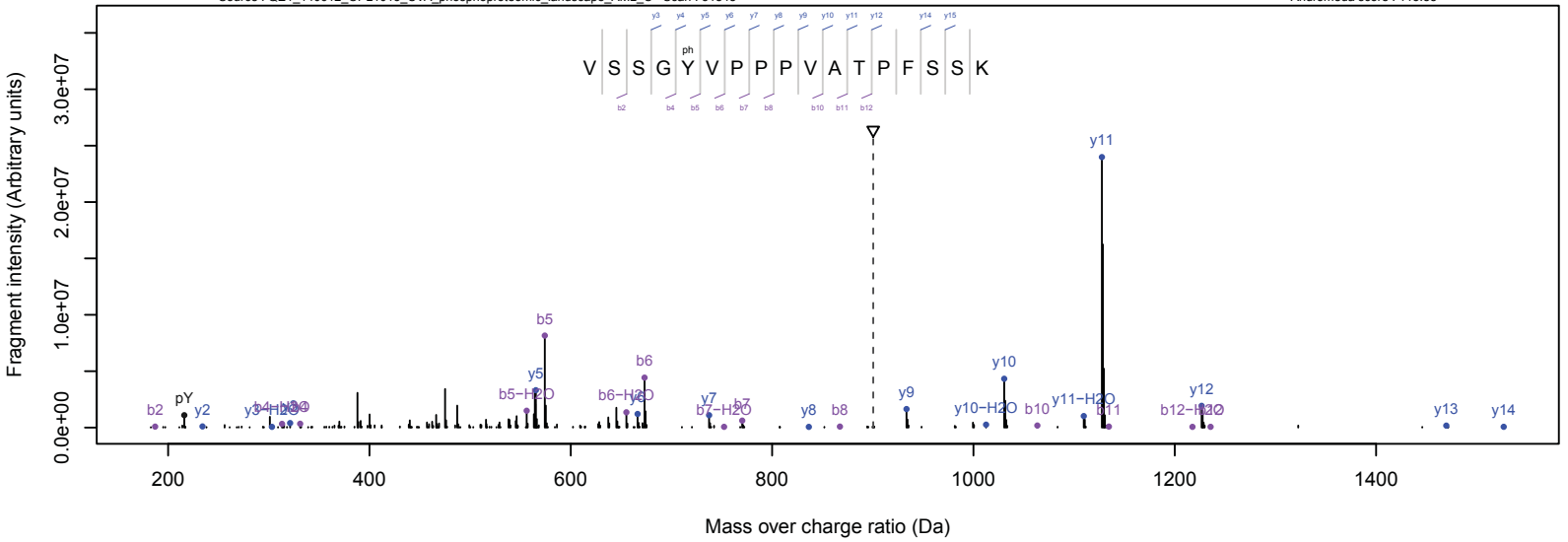
Andromeda score: 113.88



**VSSGY(ph)VPPPVATPFSSK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31848

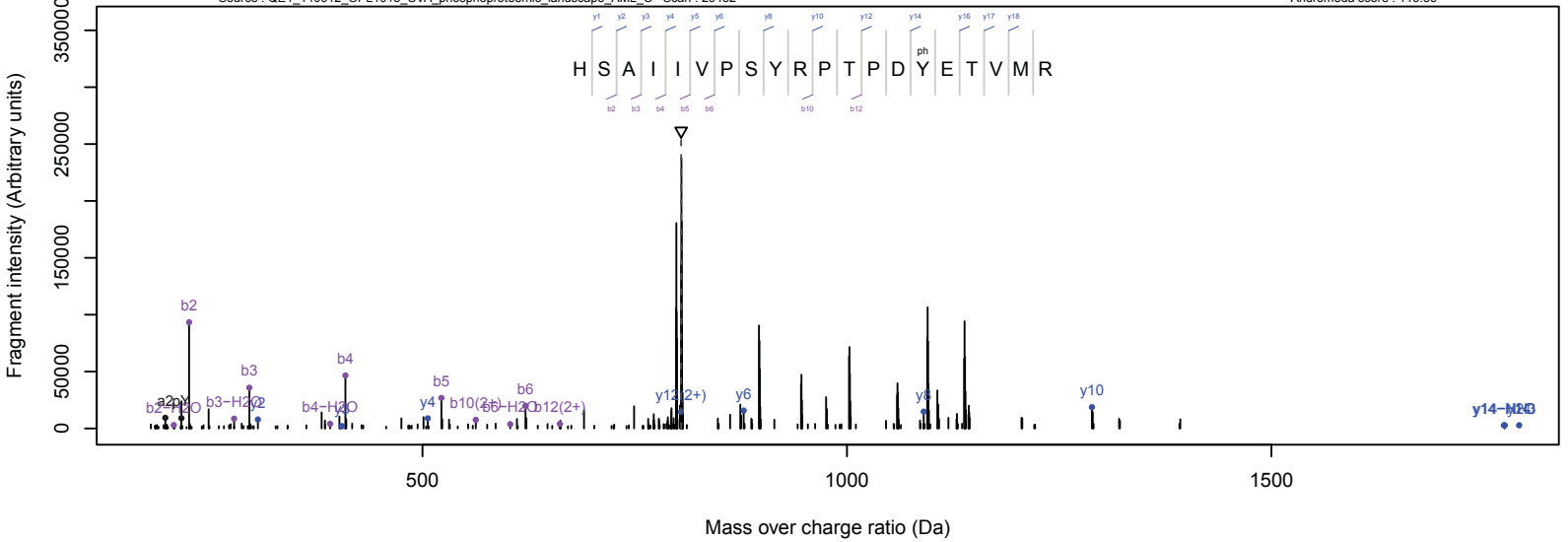
Andromeda score: 113.88



**HSAILVPSYRPTPDY(ph)ETVMR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 29432

Andromeda score: 113.86

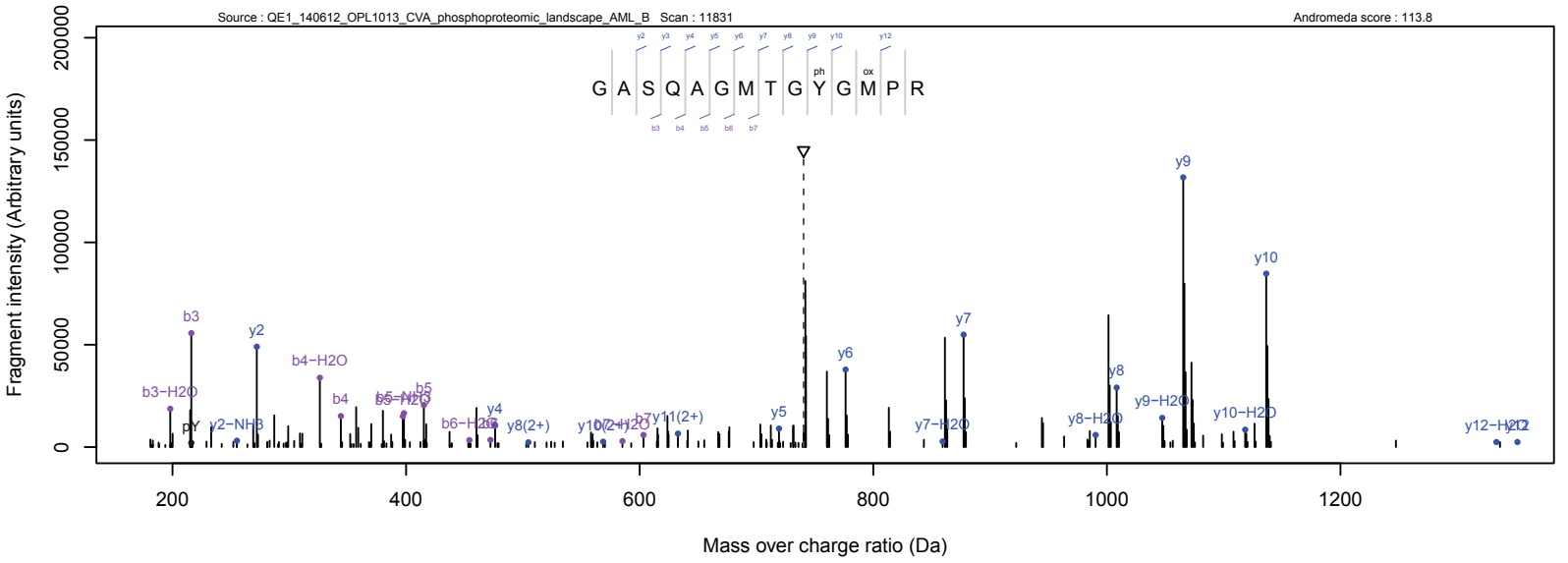




**GASQAGMTGYGM<sup>ox</sup>PR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 11831

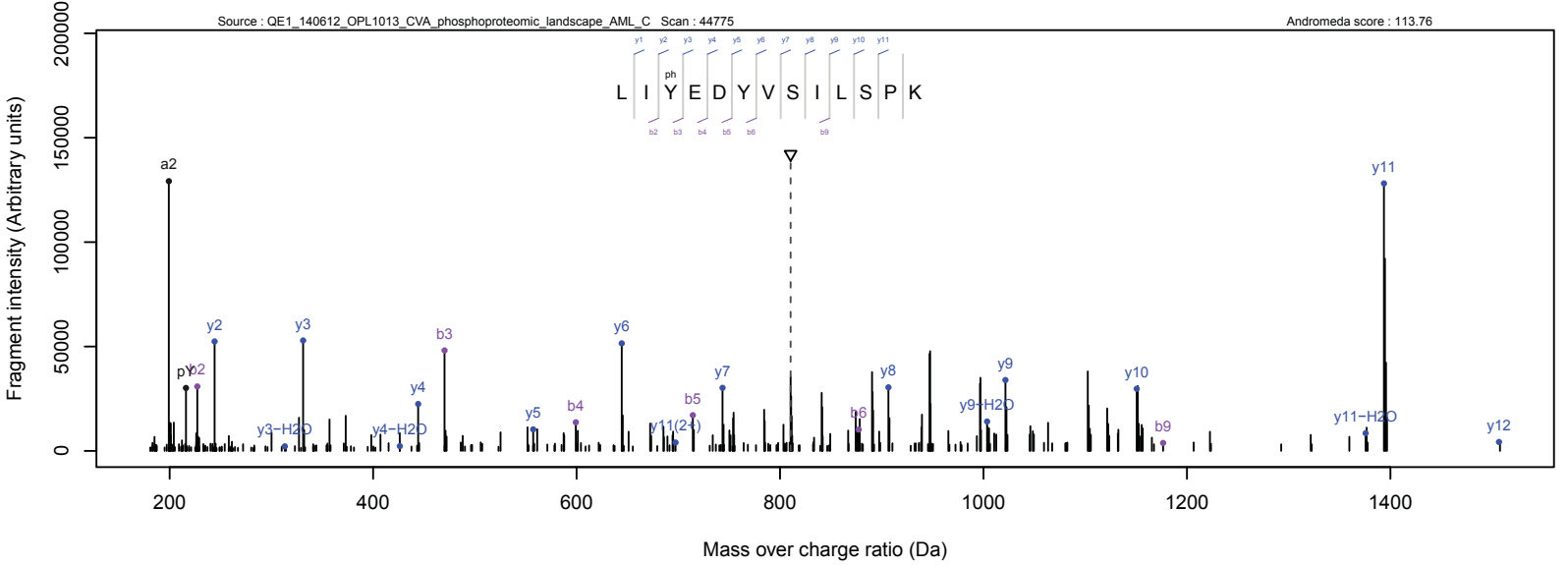
Andromeda score: 113.8



**LIY(ph)EDYVSILSPK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 44775

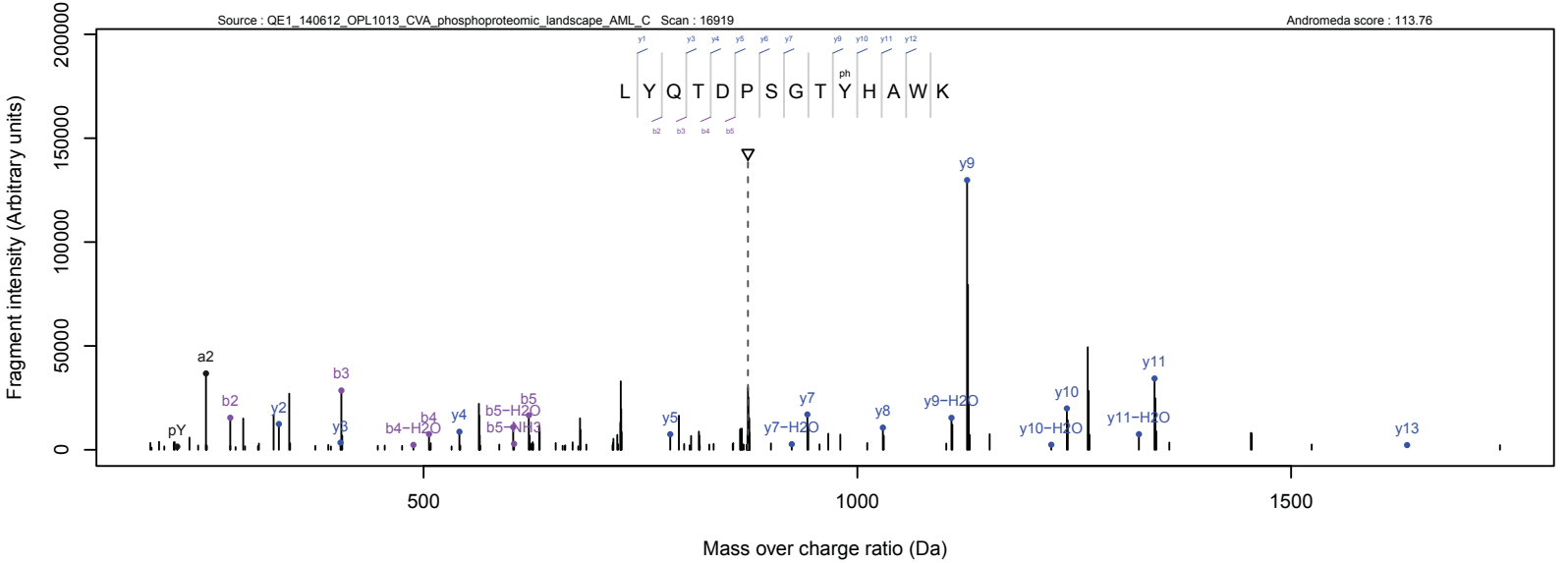
Andromeda score: 113.76



**LYQTDPSTGYHAWK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16919

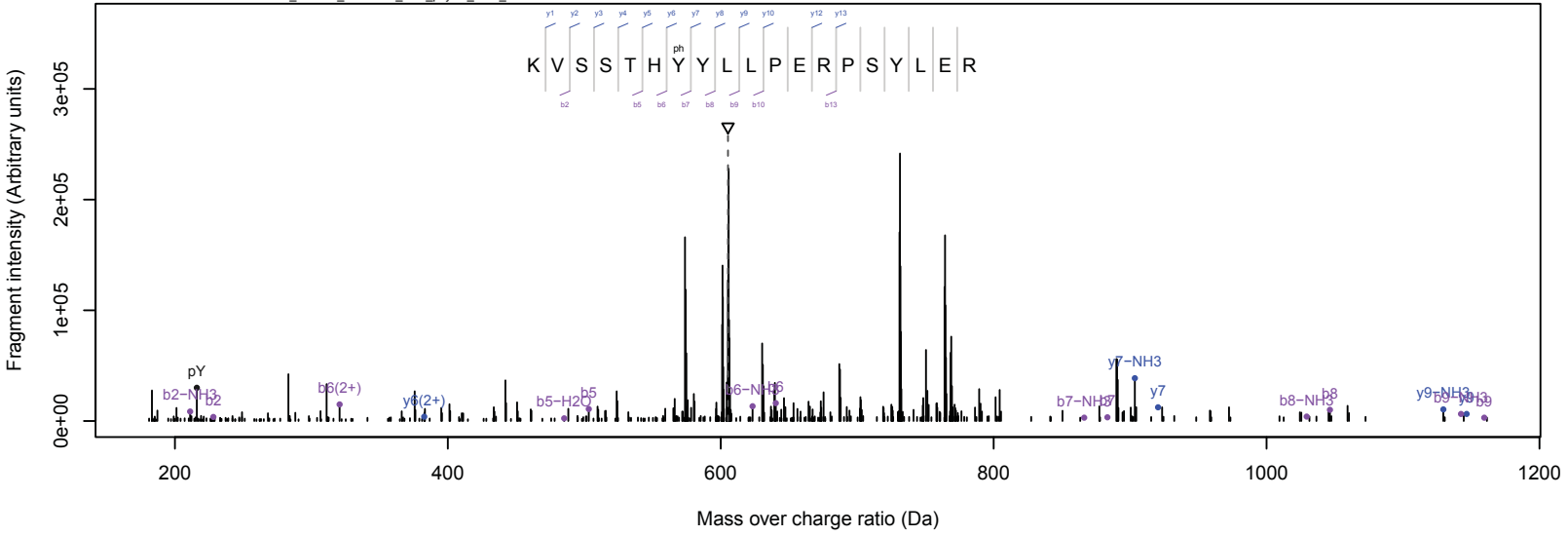
Andromeda score: 113.76



**KVSSTHY(ph)YLLPERPSYLER-4+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 25486

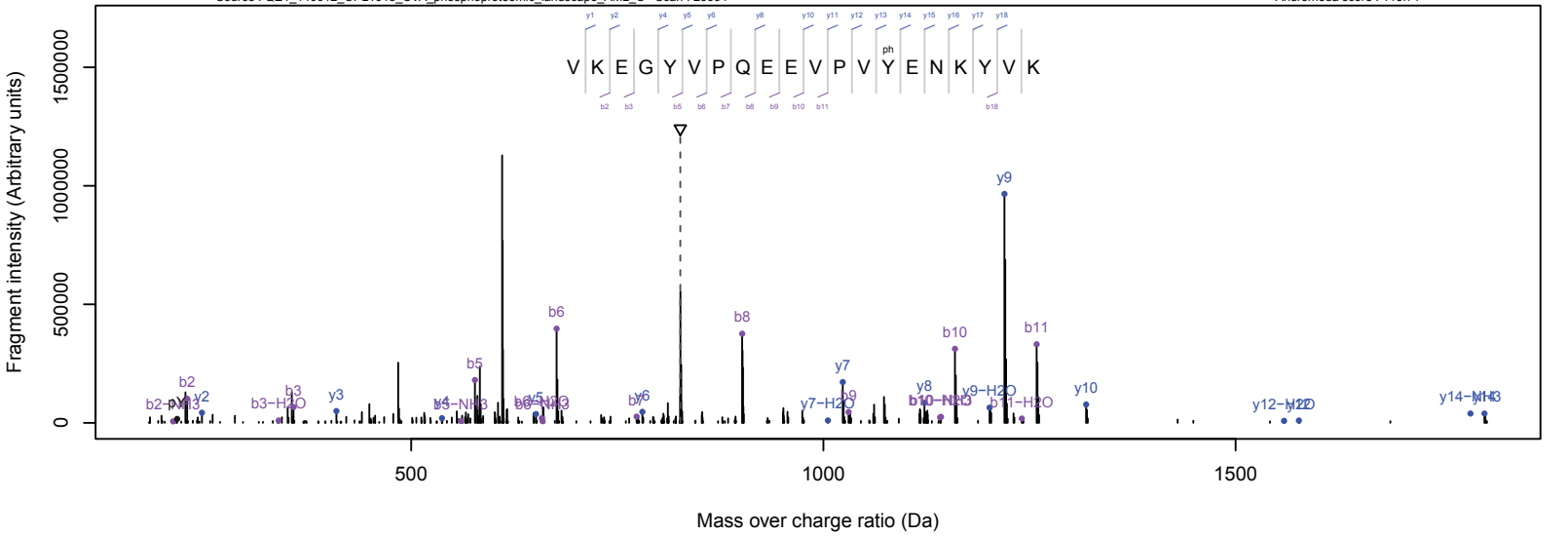
Andromeda score : 113.73



**VKEGYVPQEEVPV(ph)ENKYVK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23334

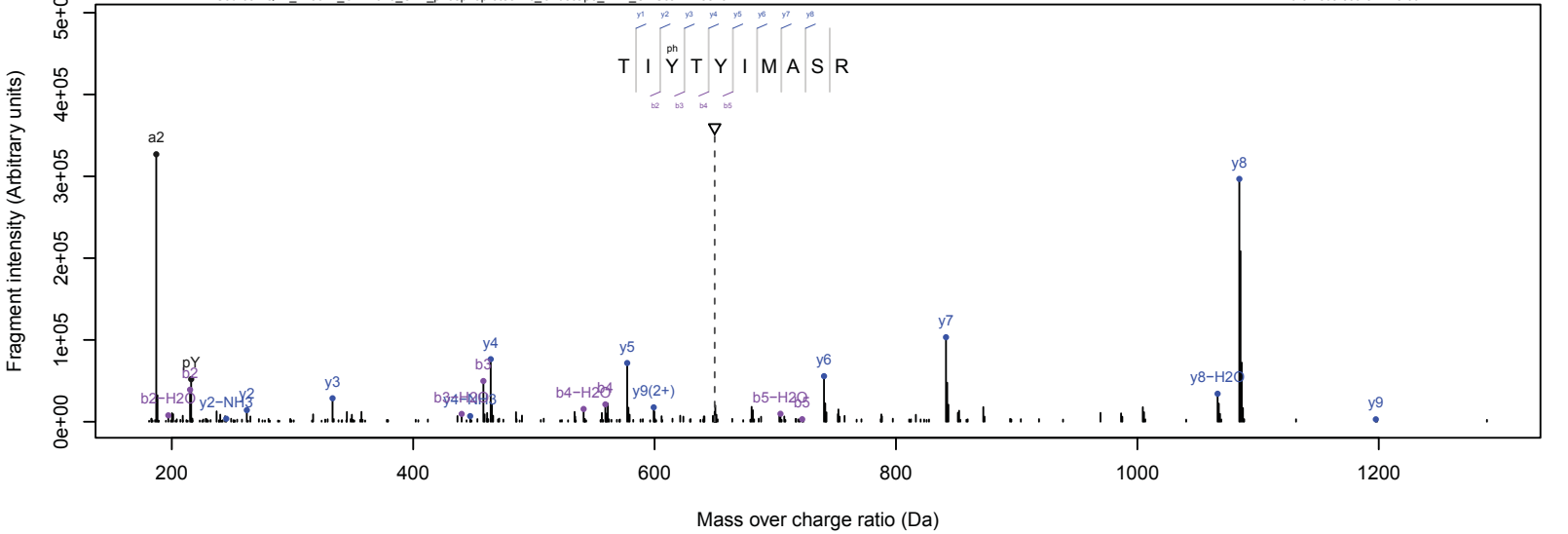
Andromeda score : 113.71



**TIY(ph)TYIMASR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28843

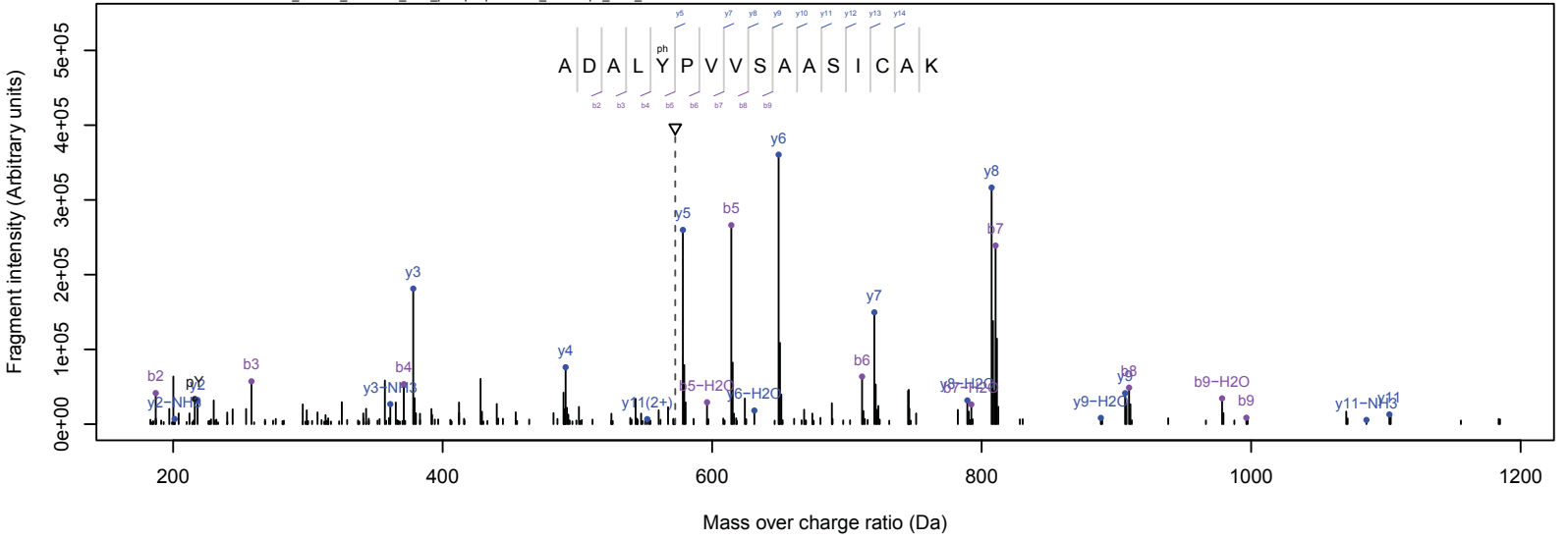
Andromeda score : 113.69



ADALY(ph)PVVSAASICAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35161

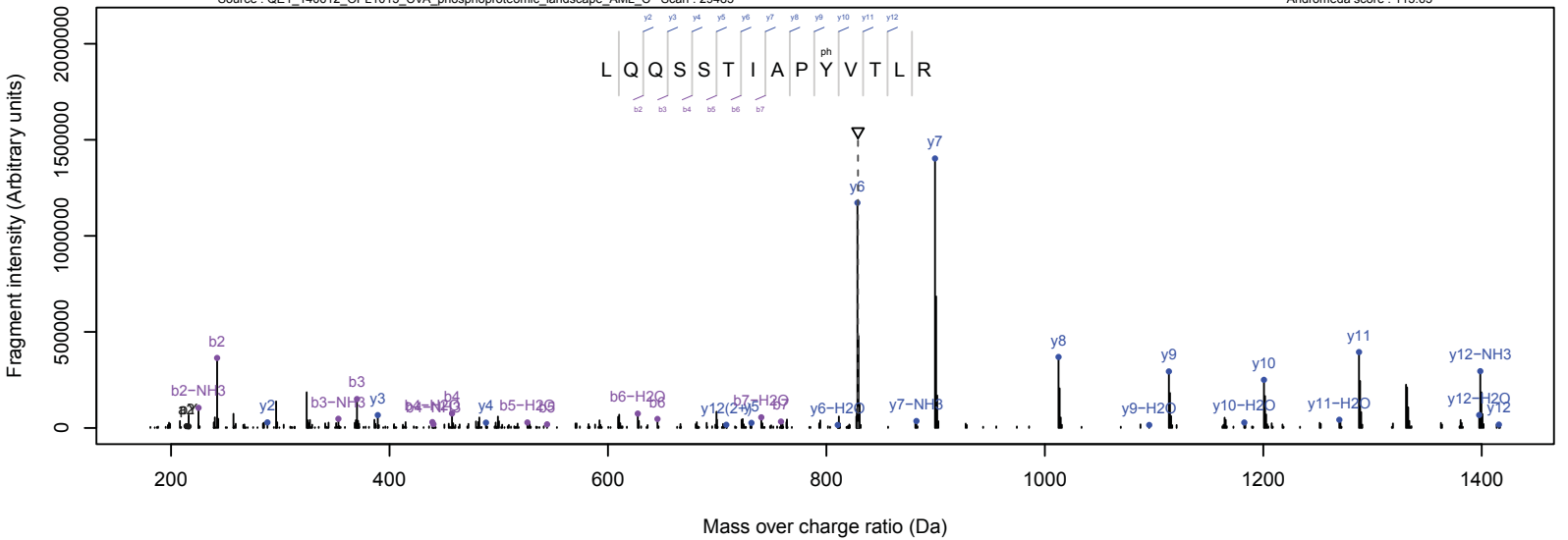
Andromeda score : 113.65



LQQSSTIAPY(ph)VTLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29485

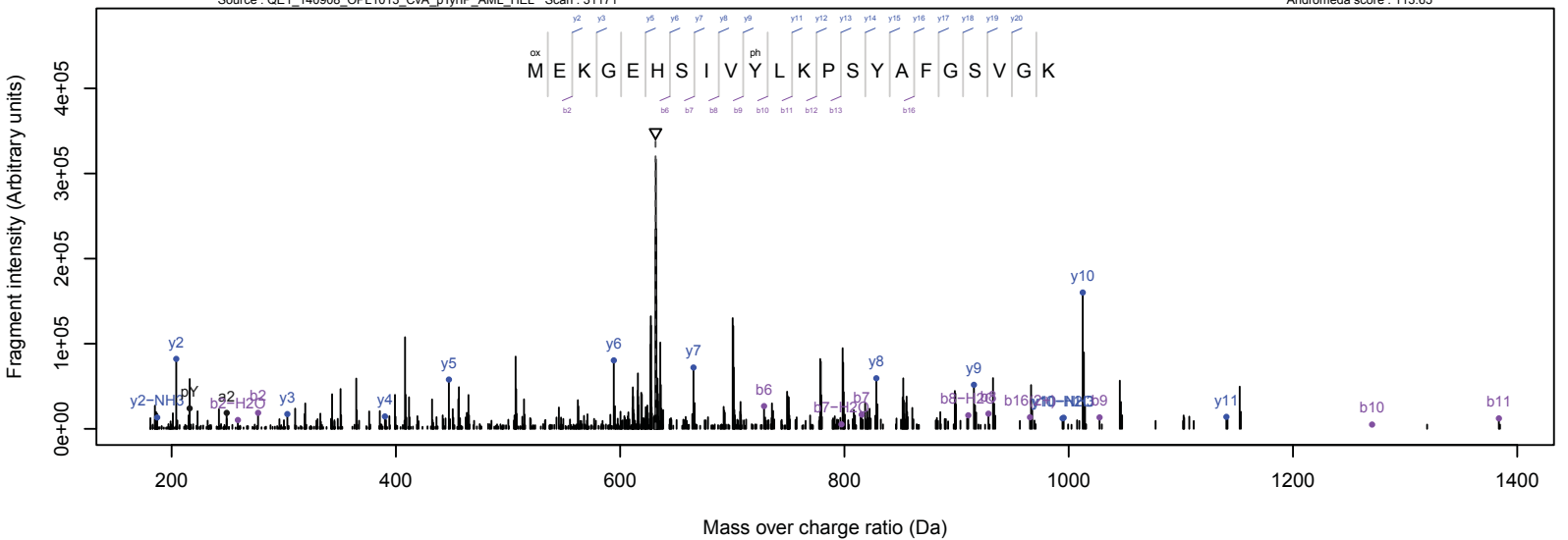
Andromeda score : 113.65



M(ox)EKGEHSIVYLPYSYAFGSVGK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31171

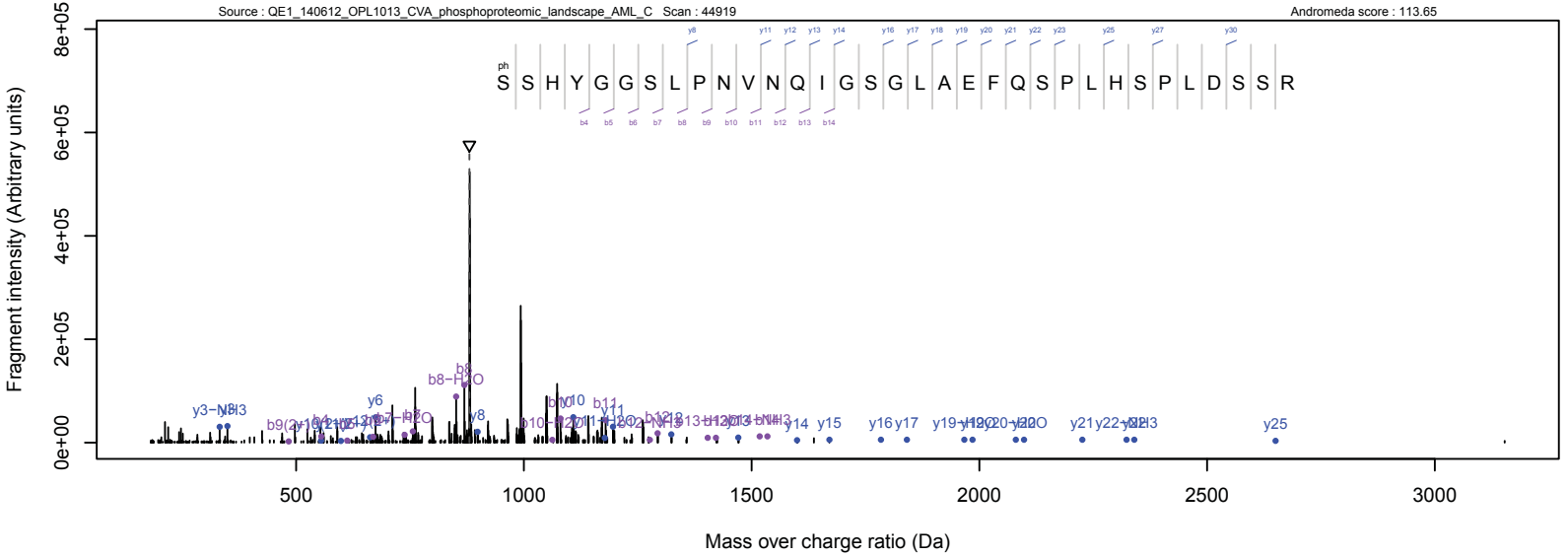
Andromeda score : 113.65



**S(ph)SHYGGSLPNVNQIGSGLAEFQSP LHSPLDSSR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44919

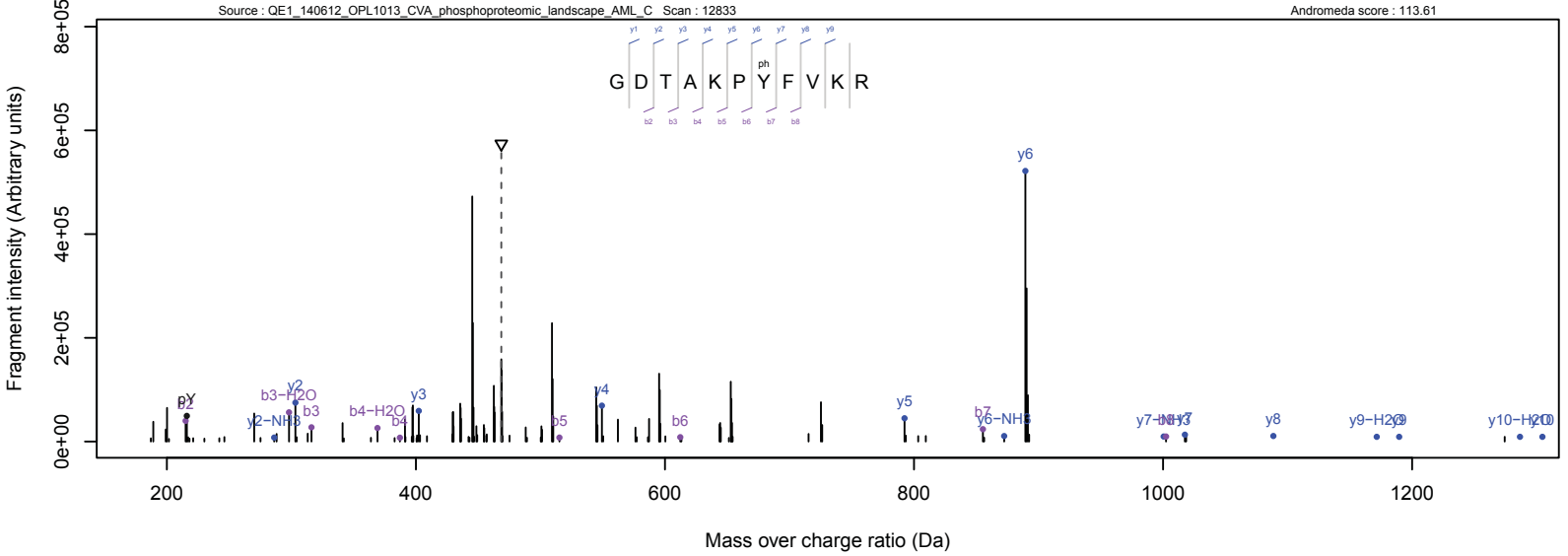
Andromeda score : 113.65



**(ac)GDTAKPY(ph)FVKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12833

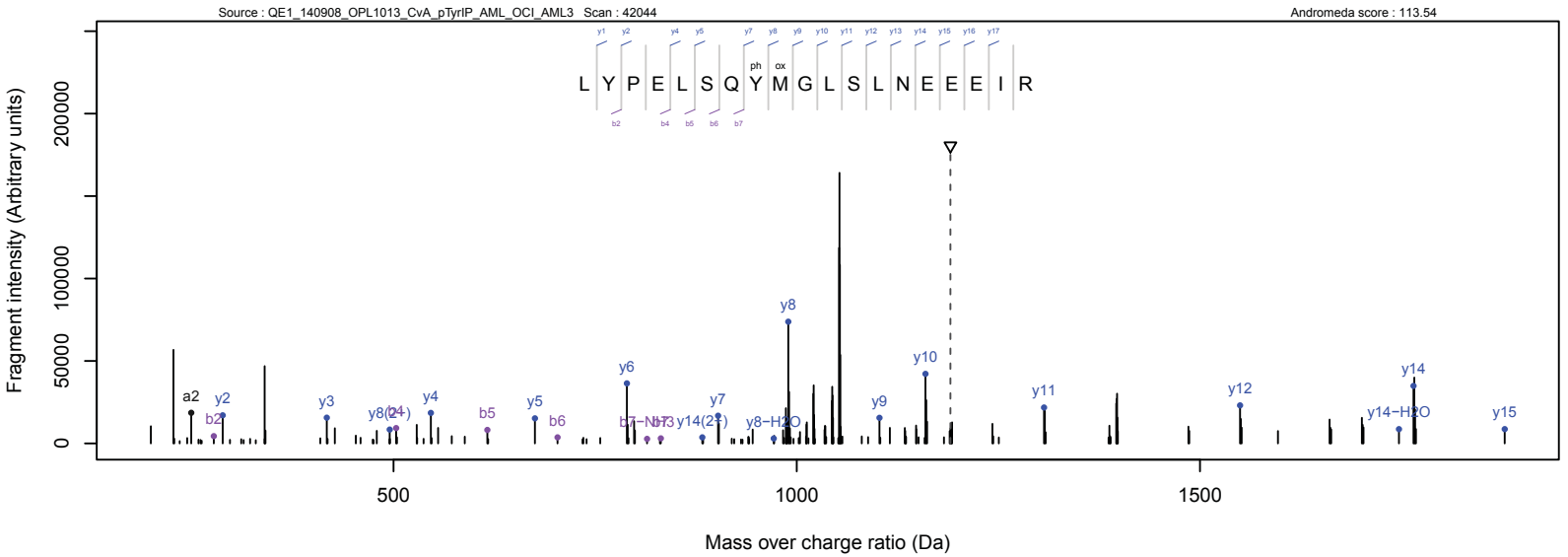
Andromeda score : 113.61



**LYPELSQY(ph)M(ox)GLSLNEEEIR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 42044

Andromeda score : 113.54

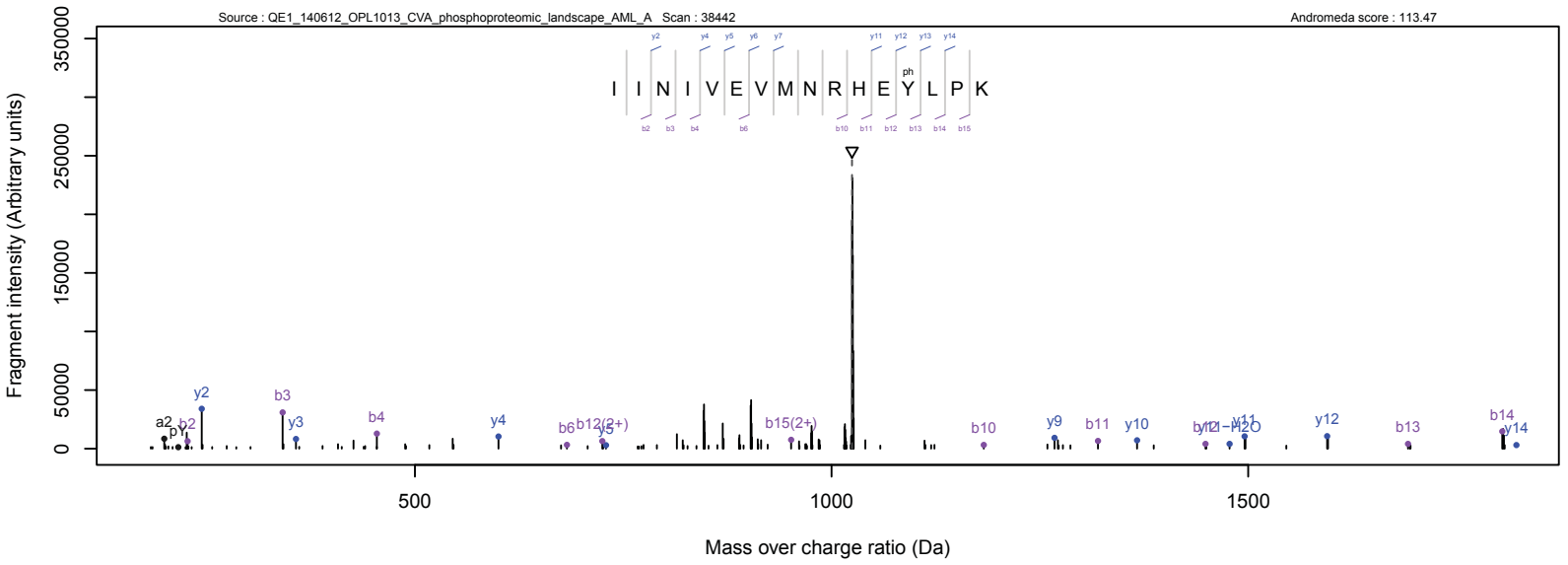




INIVEVMNRHEYLPHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 38442

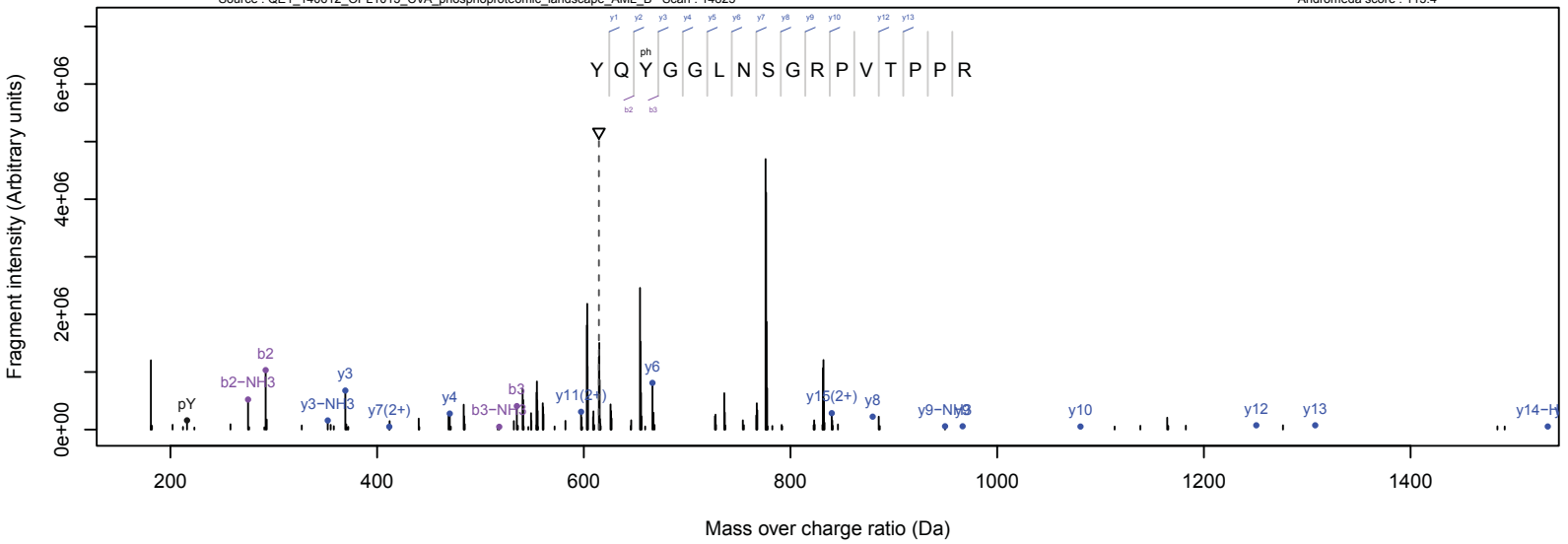
Andromeda score : 113.47



YQY(ph)GGLNSGRPVTPPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14825

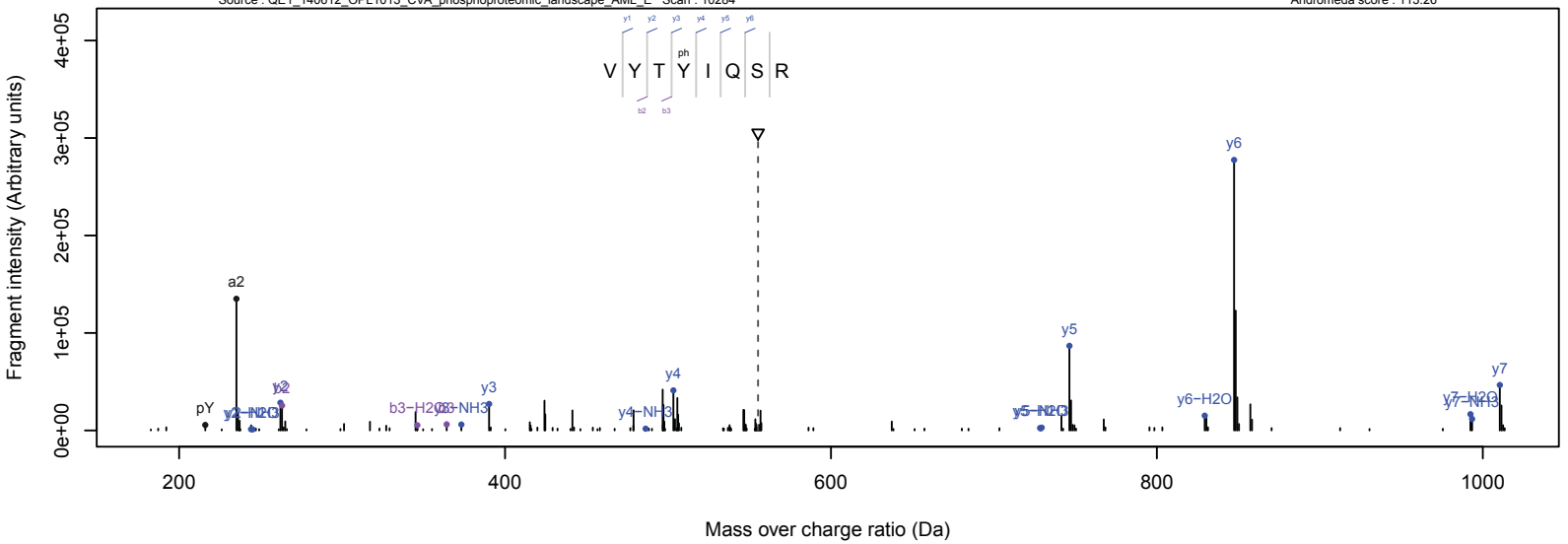
Andromeda score : 113.4



VYTY(ph)IQSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 10284

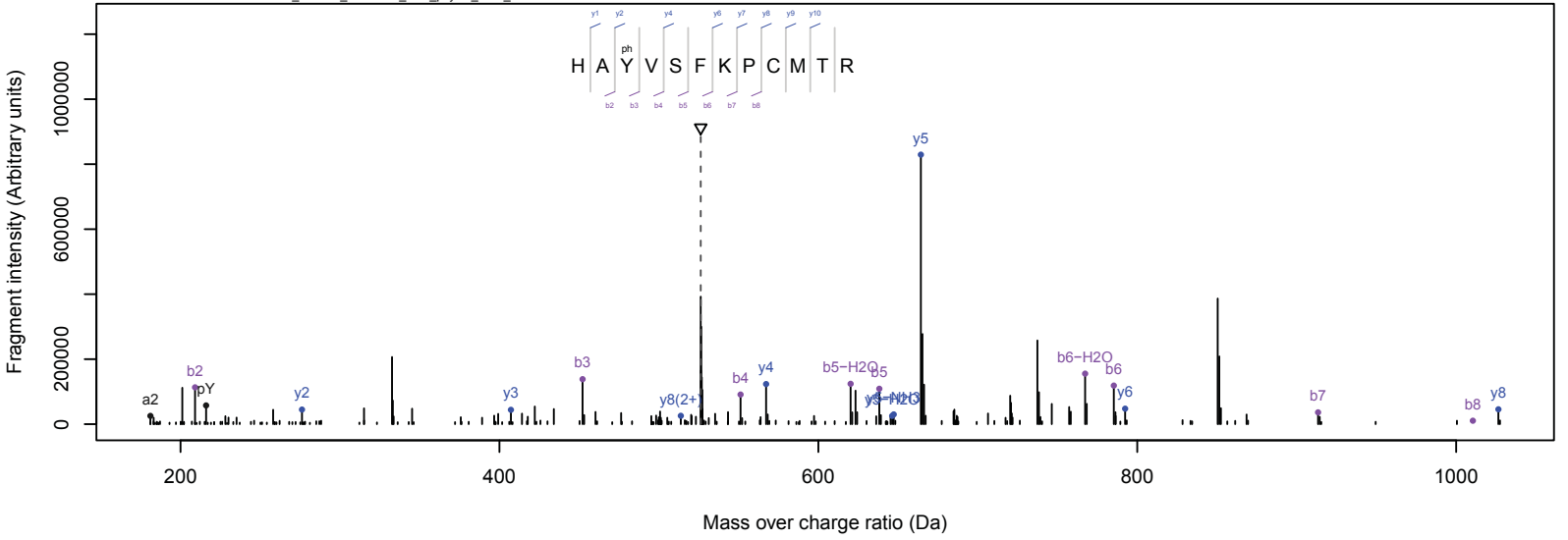
Andromeda score : 113.26



HAY(ph)VSKPCMR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 17814

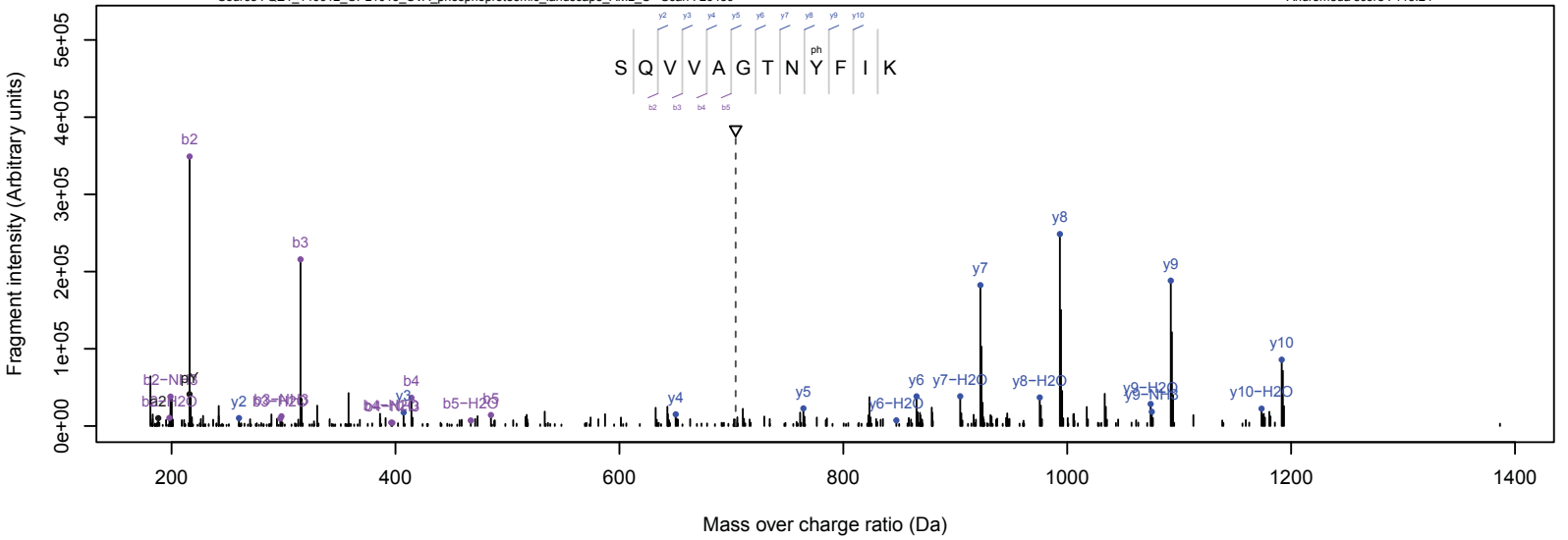
Andromeda score : 113.24



SQVVAGTNY(ph)FIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26489

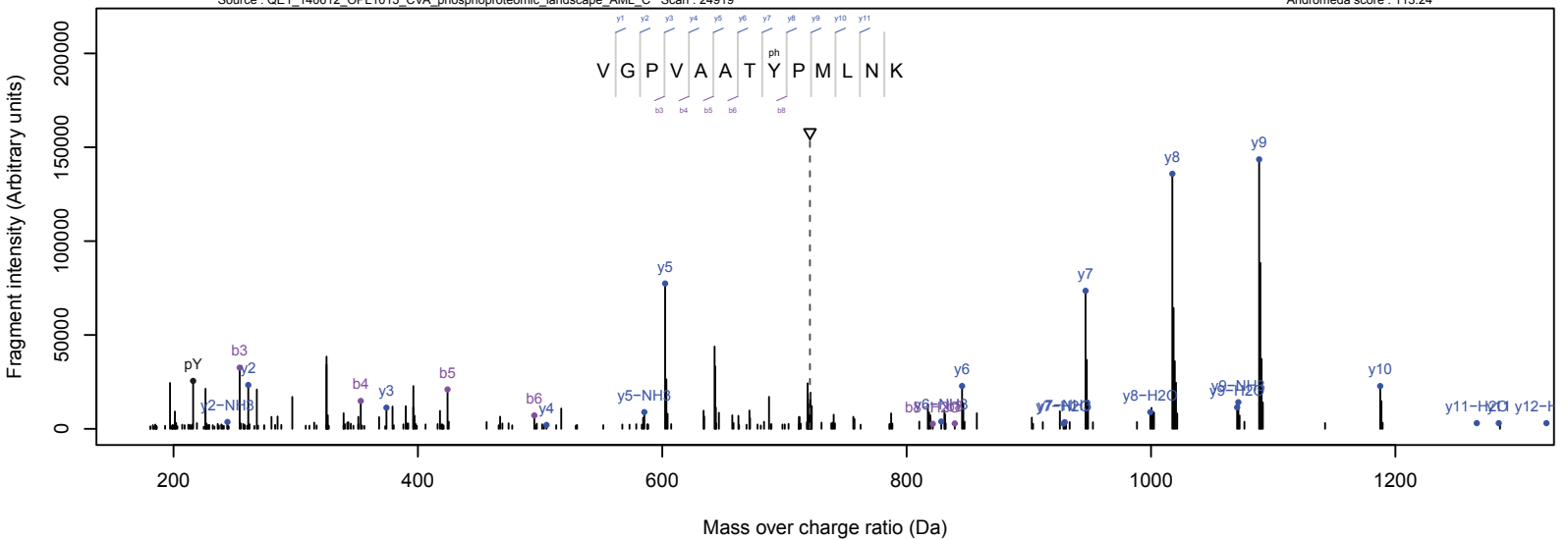
Andromeda score : 113.24



VGPVAATY(ph)PMLNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24919

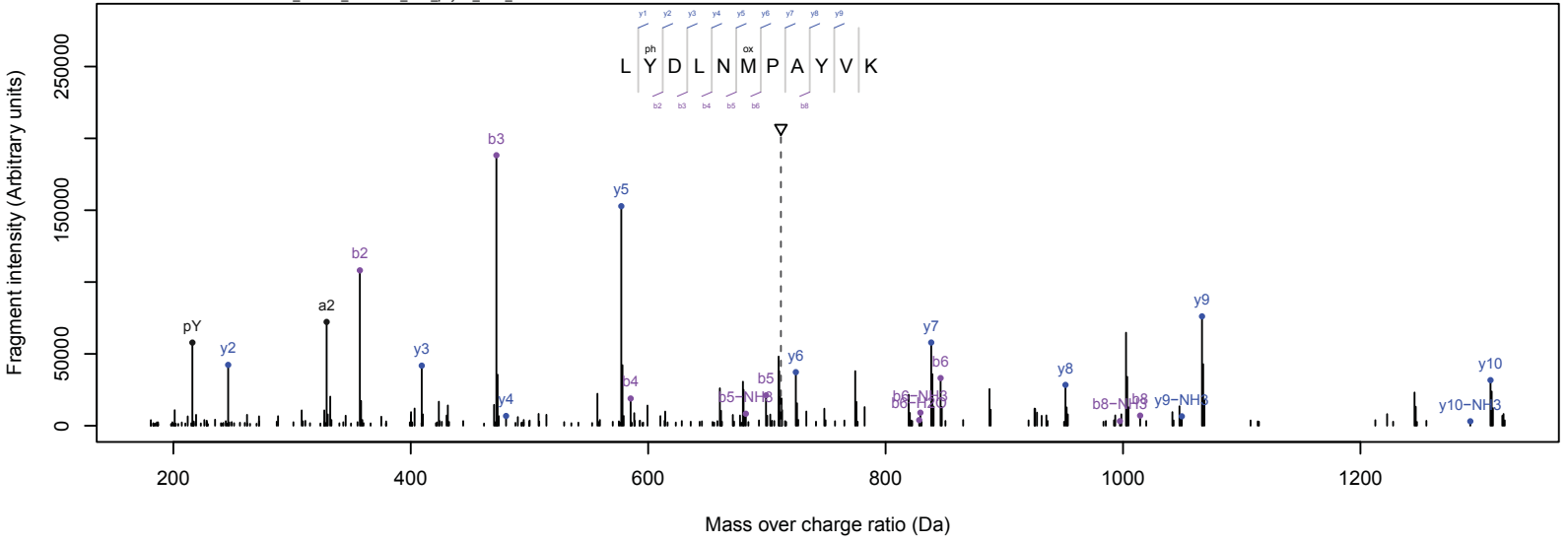
Andromeda score : 113.24



LY(ph)DLNM(ox)PAYVK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23970

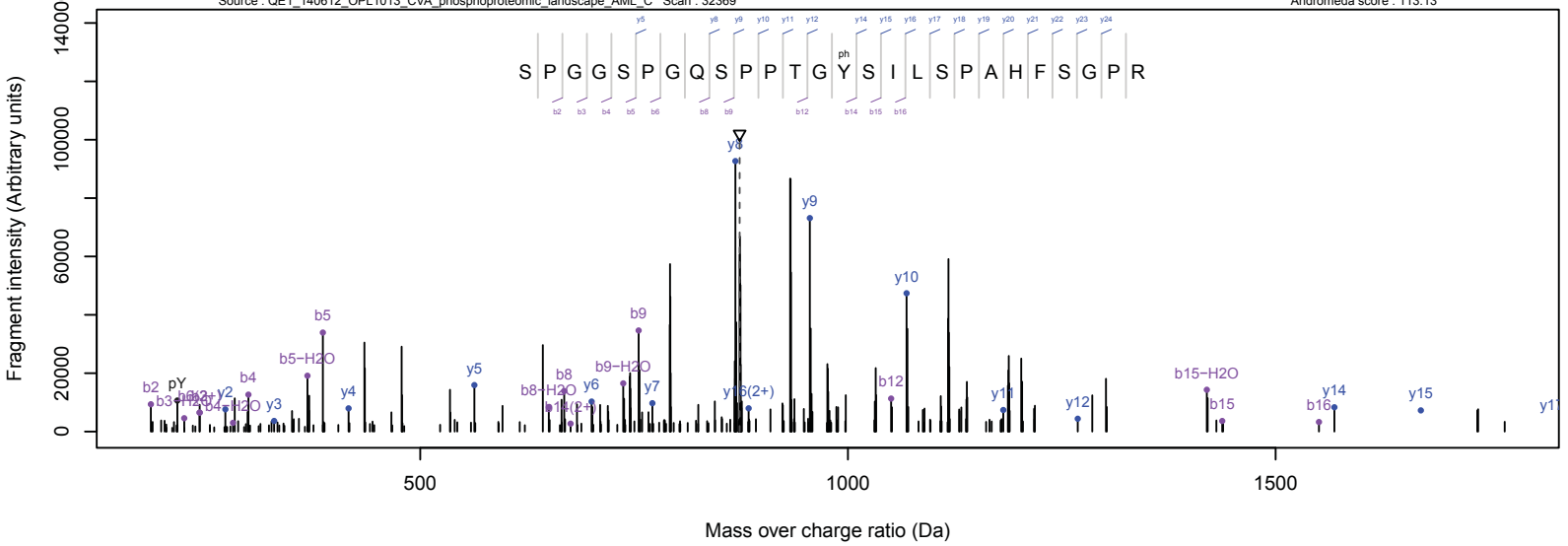
Andromeda score : 113.22



SPGGSPGQSPPTGY(ph)SILSPAHFSGPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32369

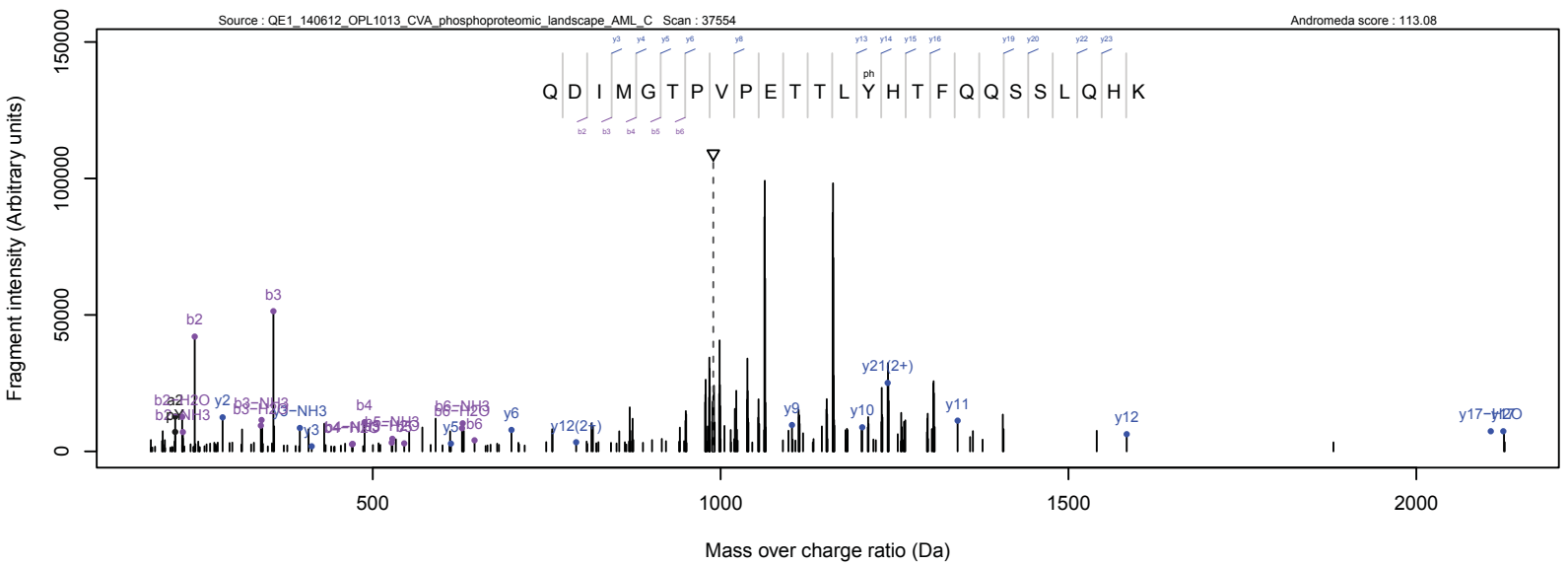
Andromeda score : 113.13



QDIMGTPVPE(TT)LYHTFQQSSLQHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37554

Andromeda score : 113.08

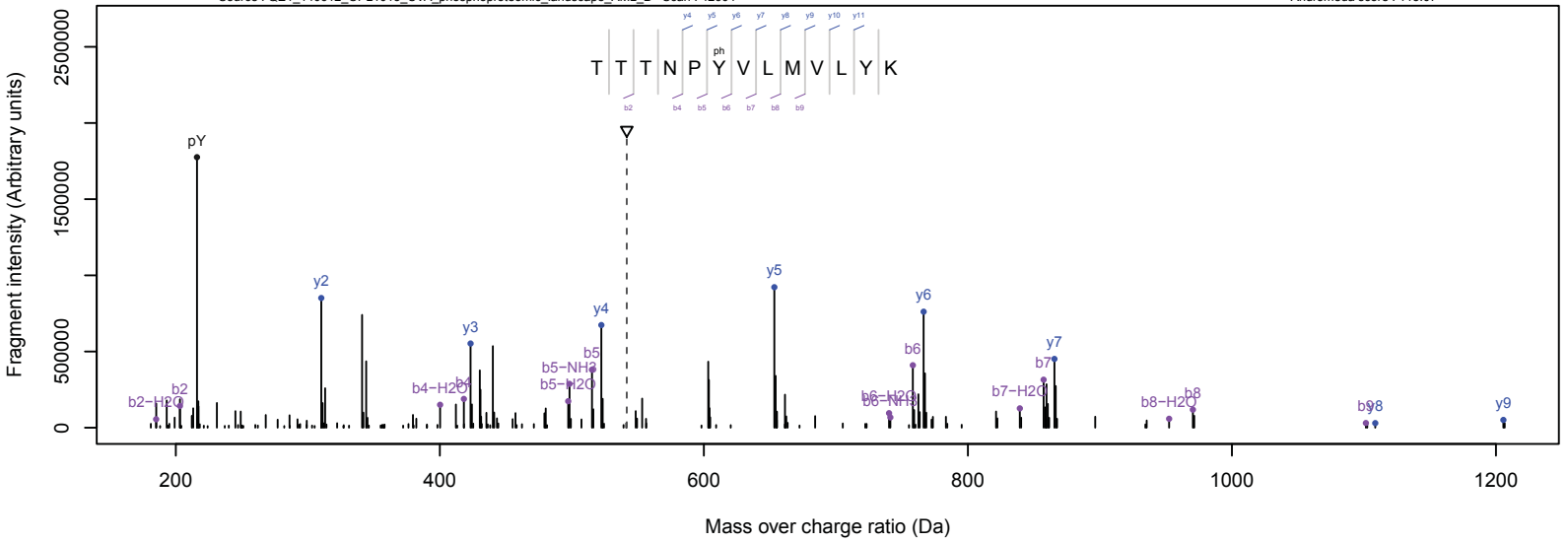




TTTNPY(ph)VLMVLYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42604

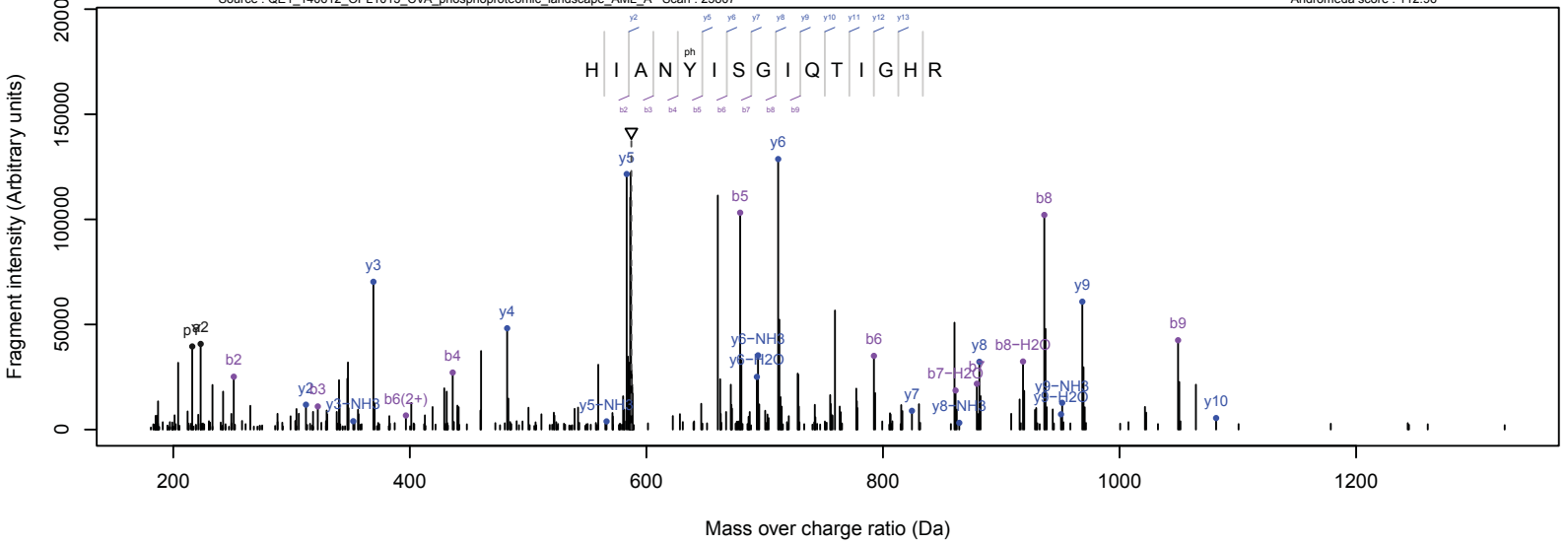
Andromeda score : 113.07



HIANY(ph)ISGIQTIGHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25807

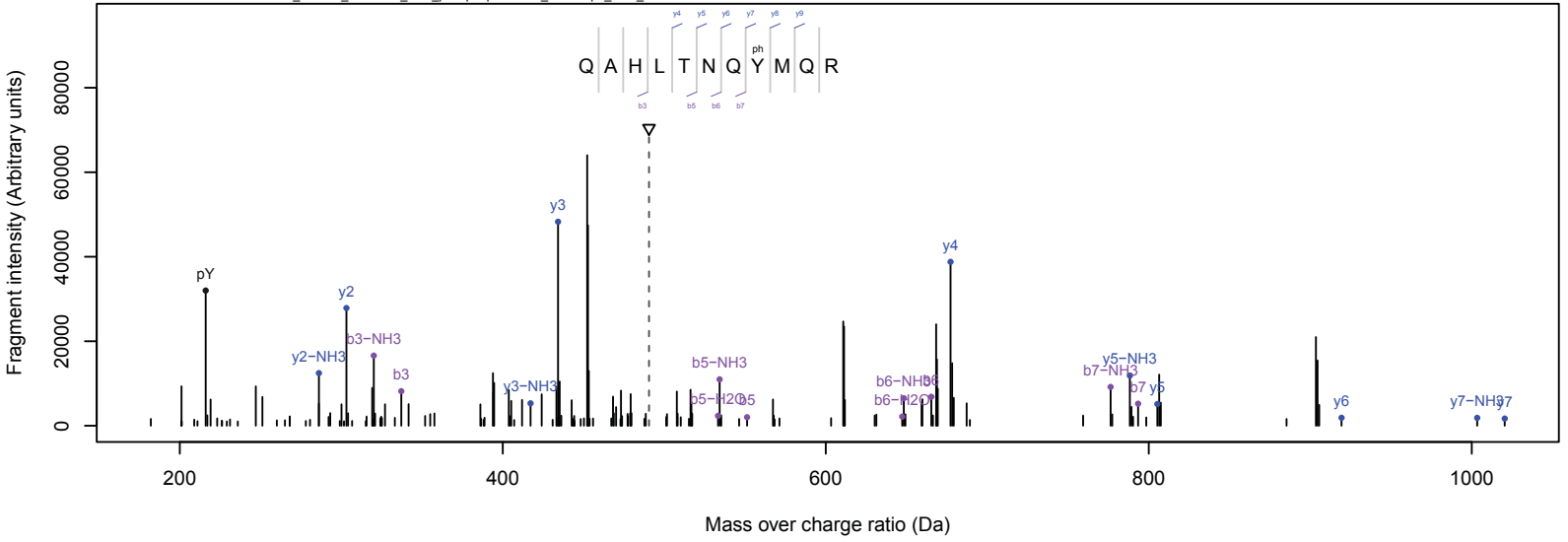
Andromeda score : 112.96



QAHLTNQ(ph)MQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6499

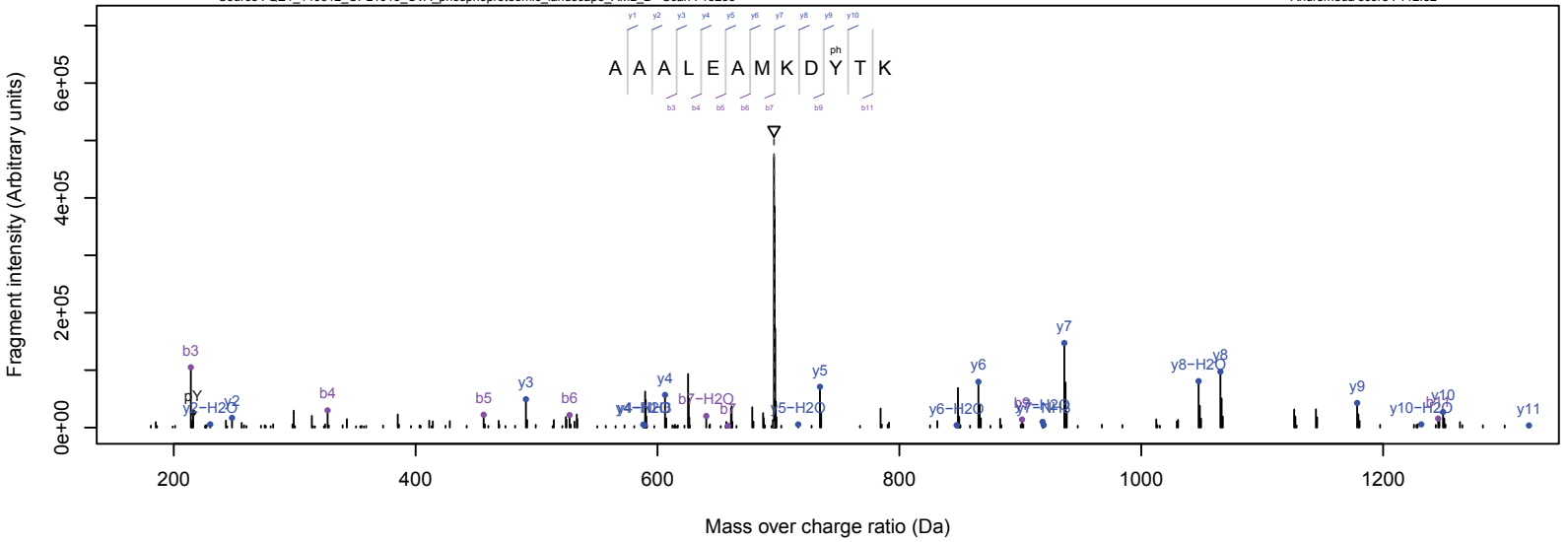
Andromeda score : 112.85



AAALEAMKDYTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13235

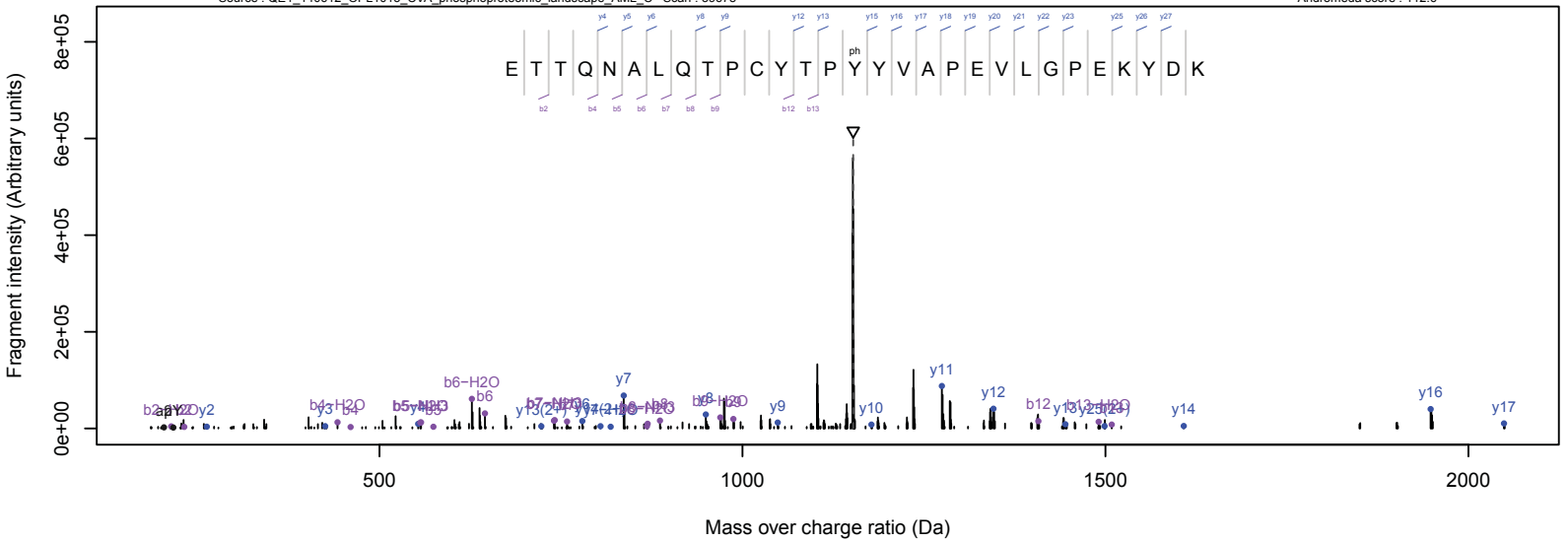
Andromeda score : 112.82



ETTQNALQTPCYPYVAVPEVLGPEKYDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39678

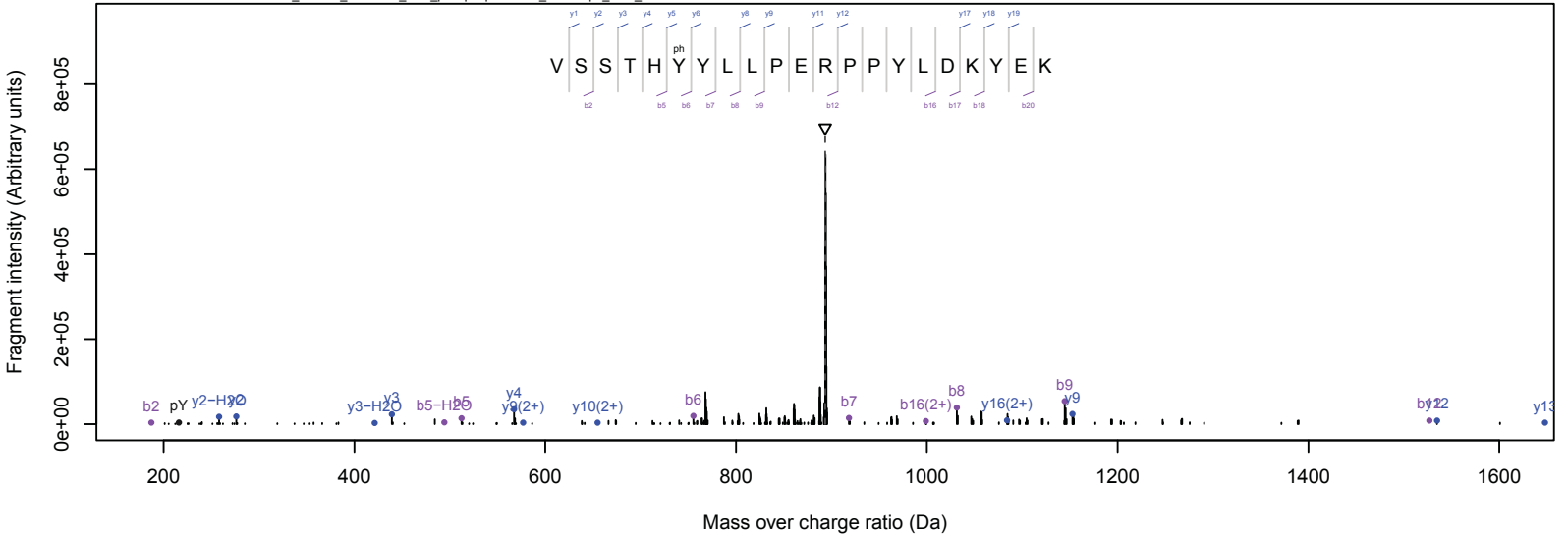
Andromeda score : 112.6



**VSSTHY(ph)YLLPERPPYLDKYEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31276

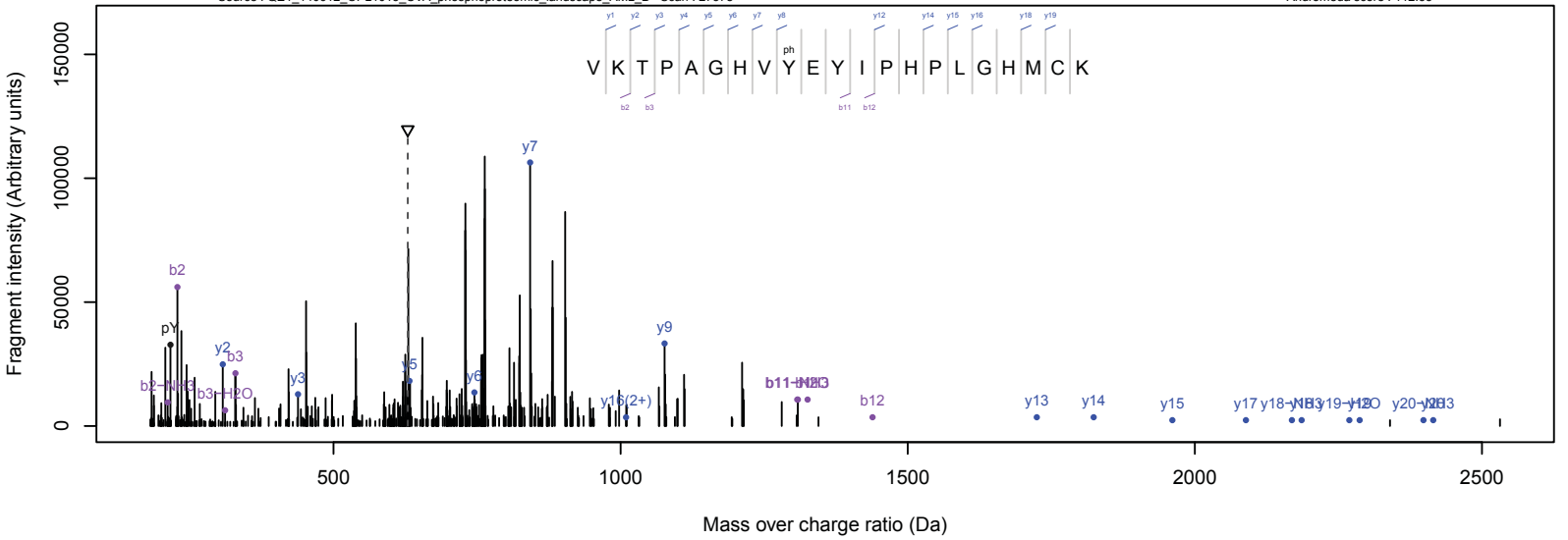
Andromeda score : 112.58



**VKTPAGHVY(ph)EYIPHLGHMCK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27378

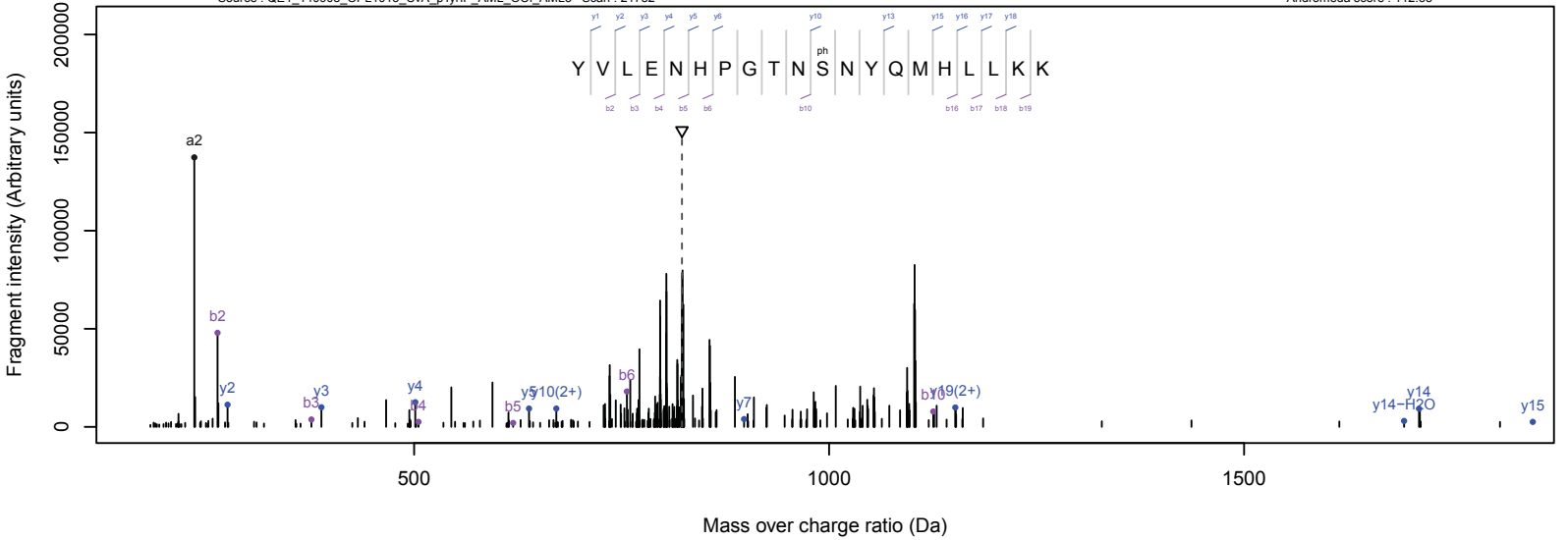
Andromeda score : 112.55



**YVLENHPGTS(ph)NYQMHLKK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 21782

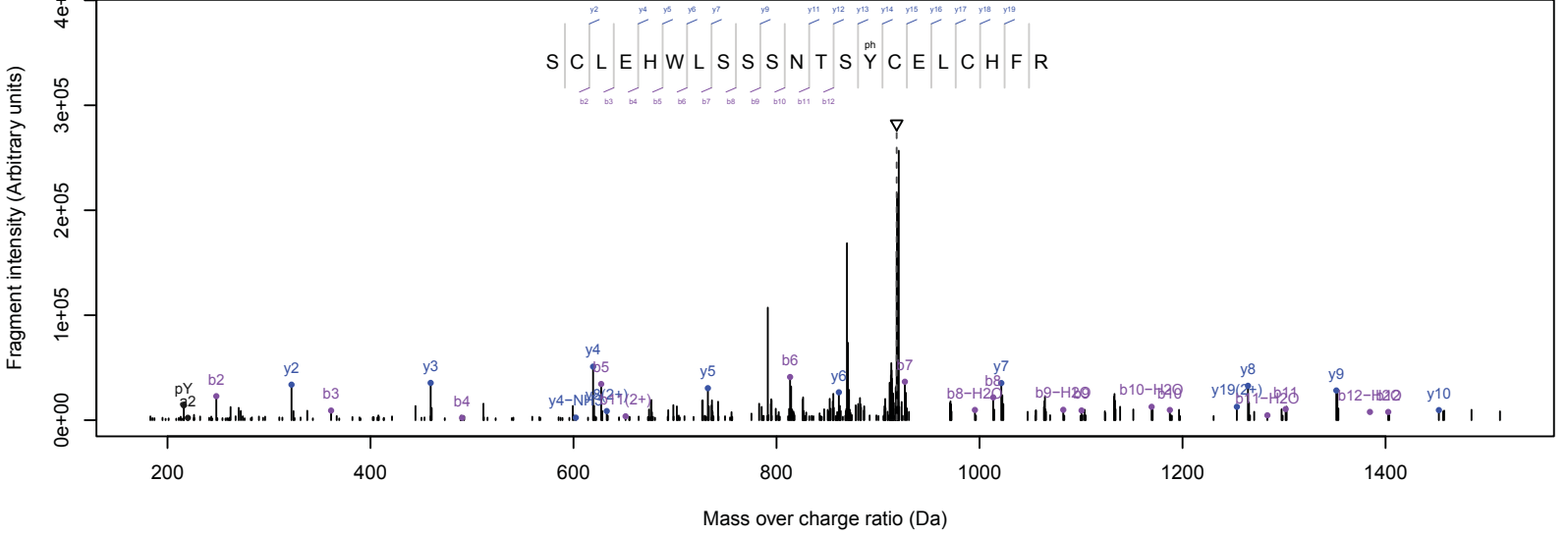
Andromeda score : 112.55



**SCLEHWLSSNTSY(ph)CELCHFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35141

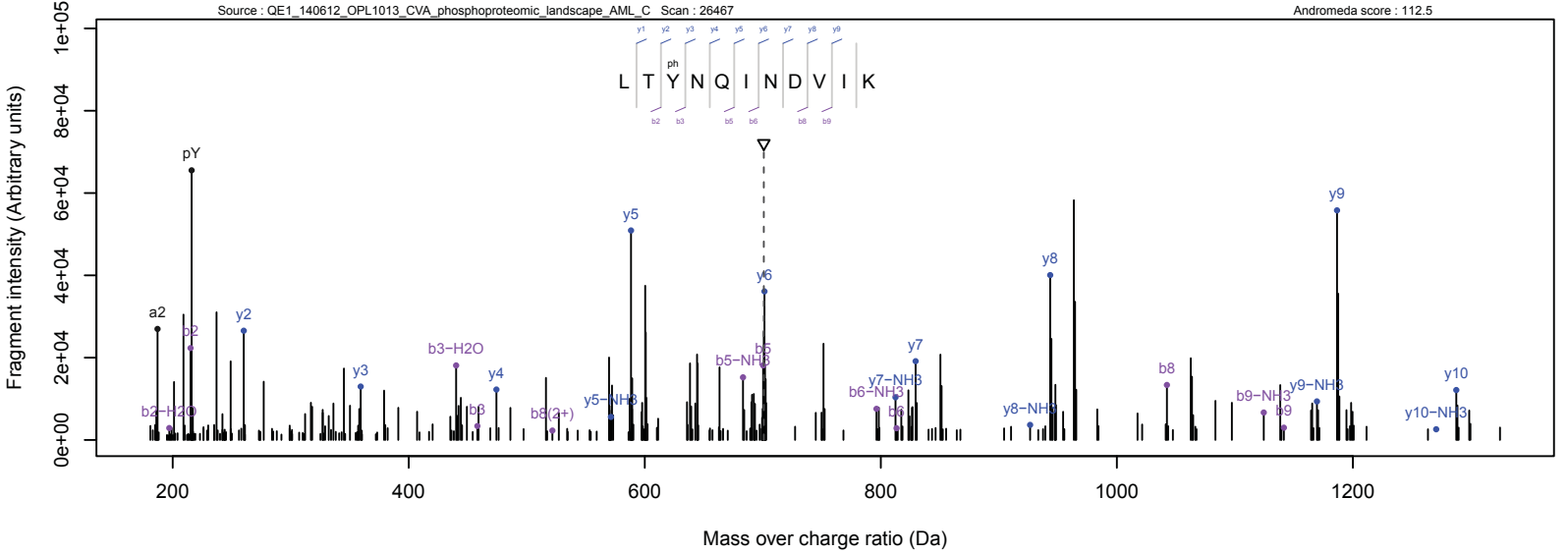
Andromeda score : 112.53



**LTY(ph)NQINDVIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26467

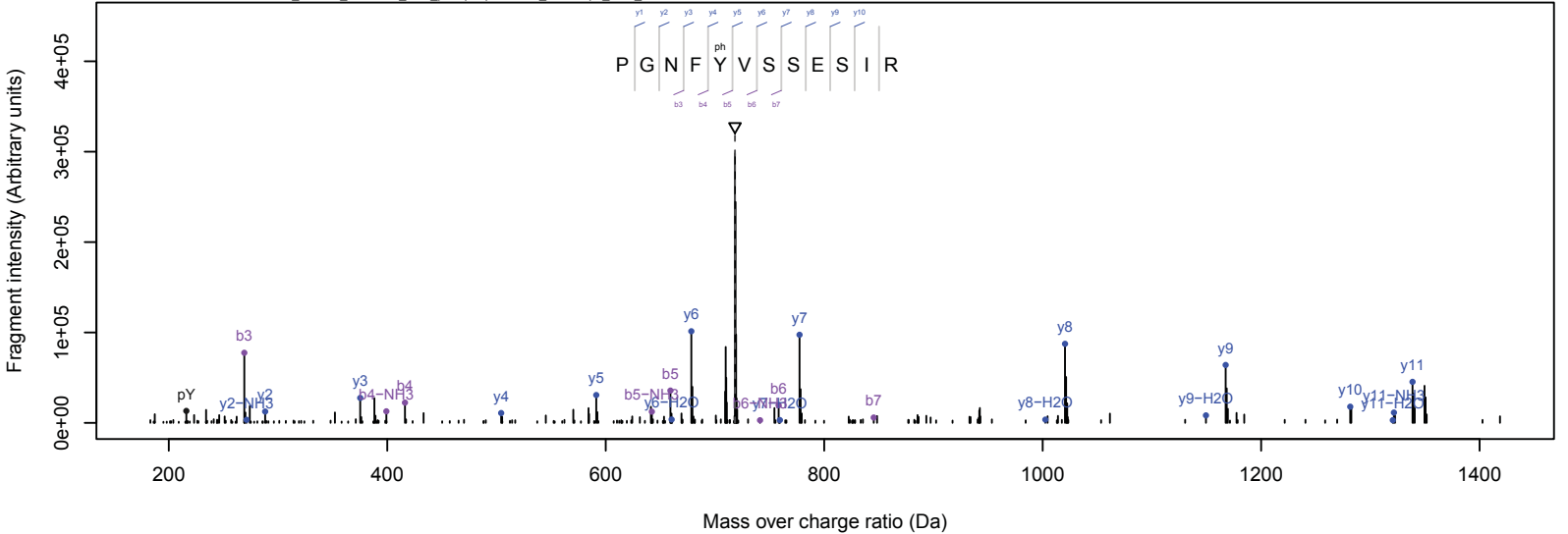
Andromeda score : 112.5



**PGNFY(ph)VSSESISIR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 22977

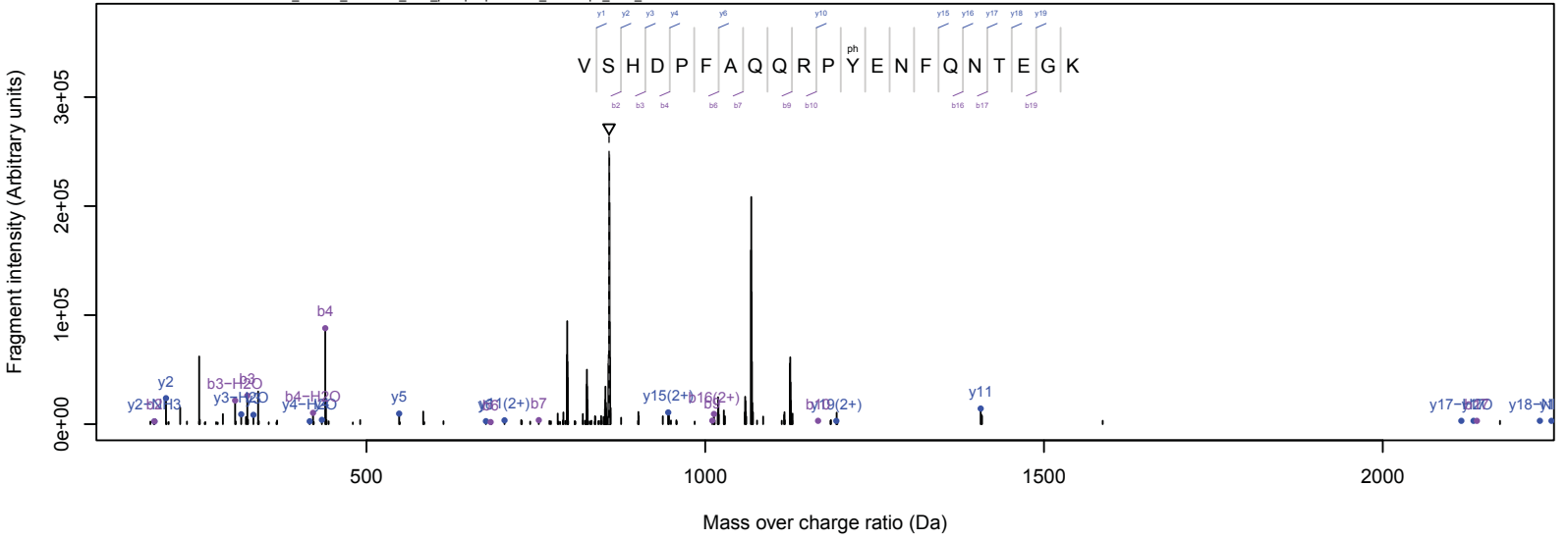
Andromeda score : 112.44



VSHDPFAQRPY(ph)ENFQNTGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18984

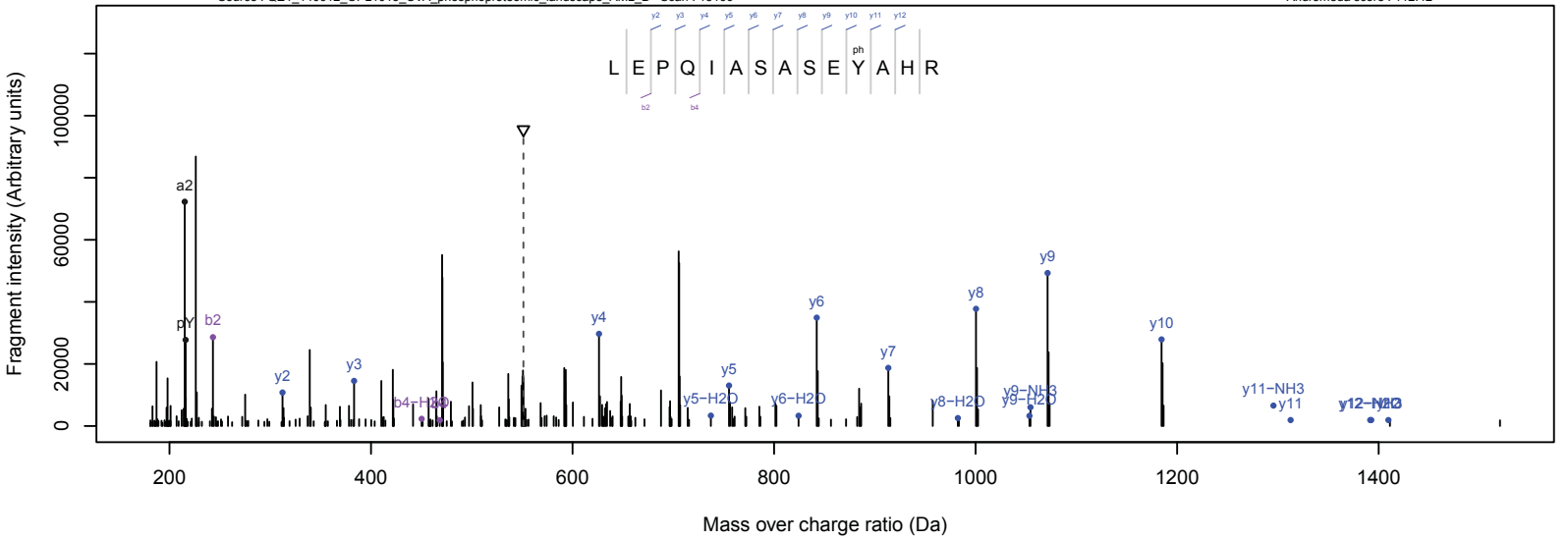
Andromeda score : 112.44



LEPQIASASEY(ph)AHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15166

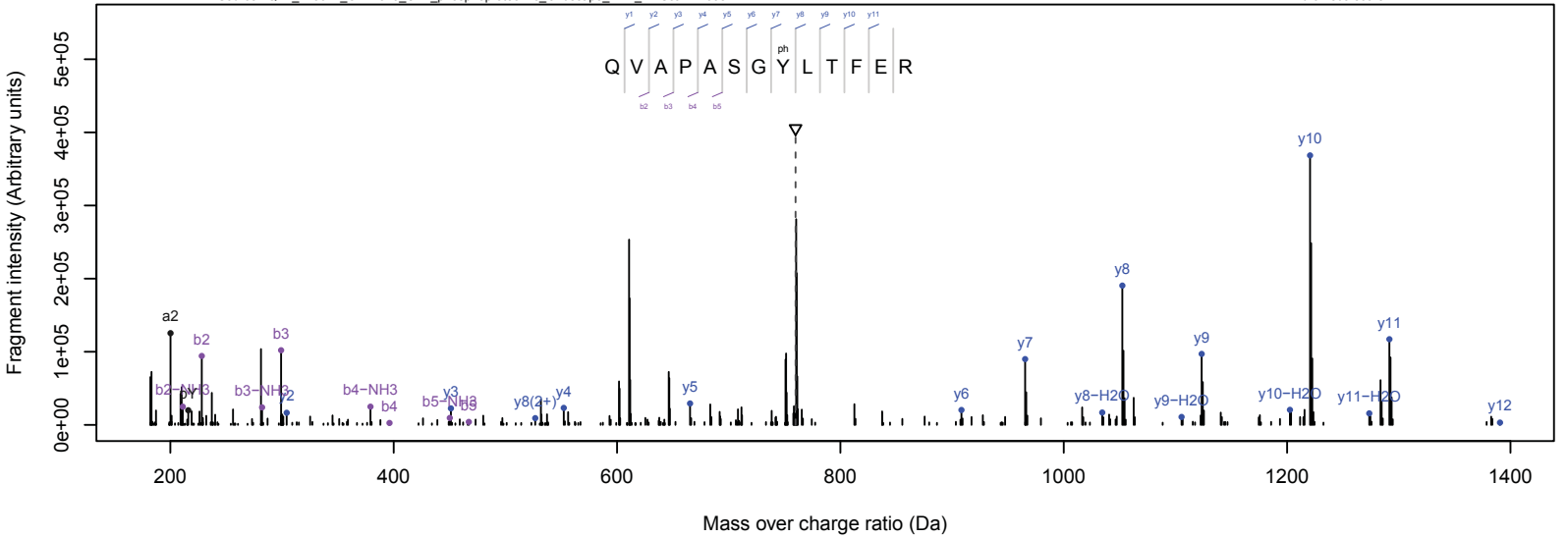
Andromeda score : 112.42



QVAPASGY(ph)LTFER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26382

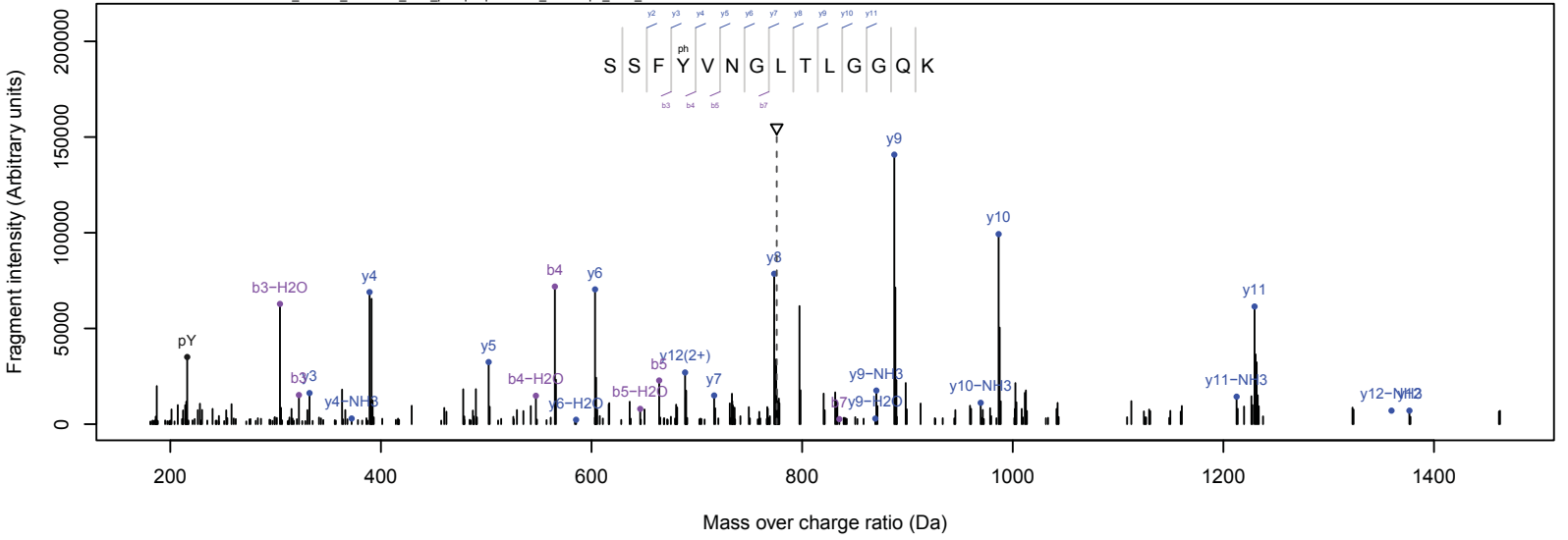
Andromeda score : 112.42



SSFY(ph)VNGLTGGQK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34426

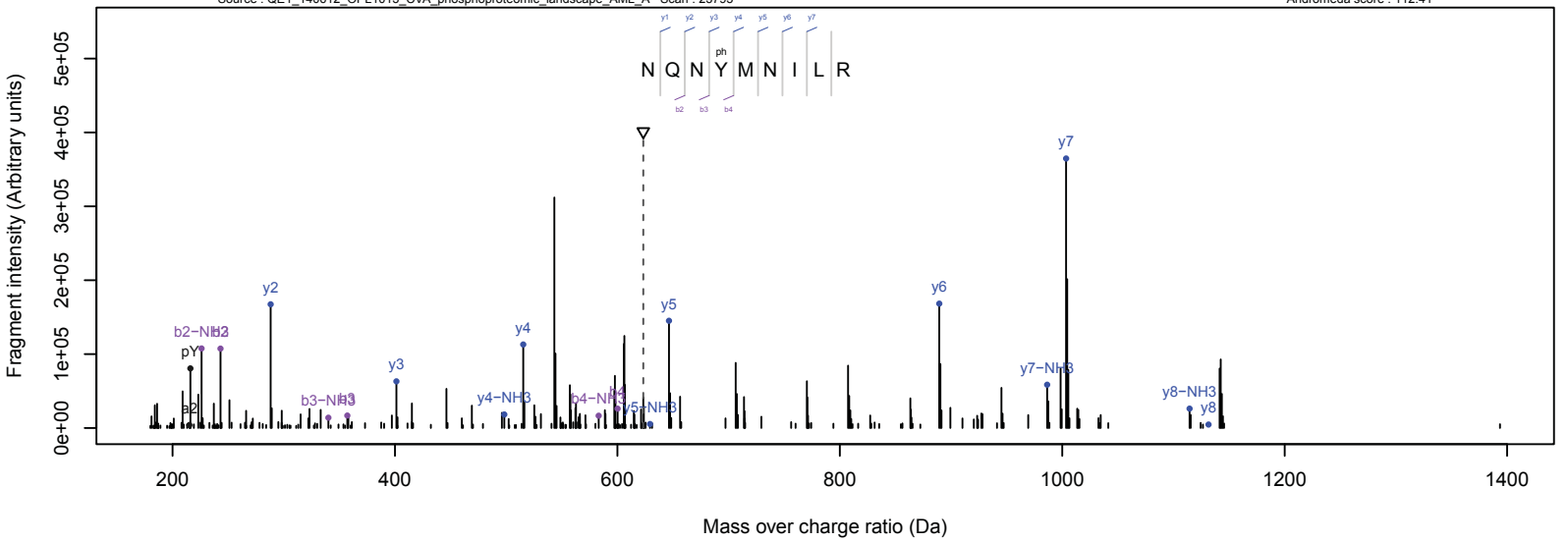
Andromeda score: 112.42



NQNY(ph)MNILR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 23793

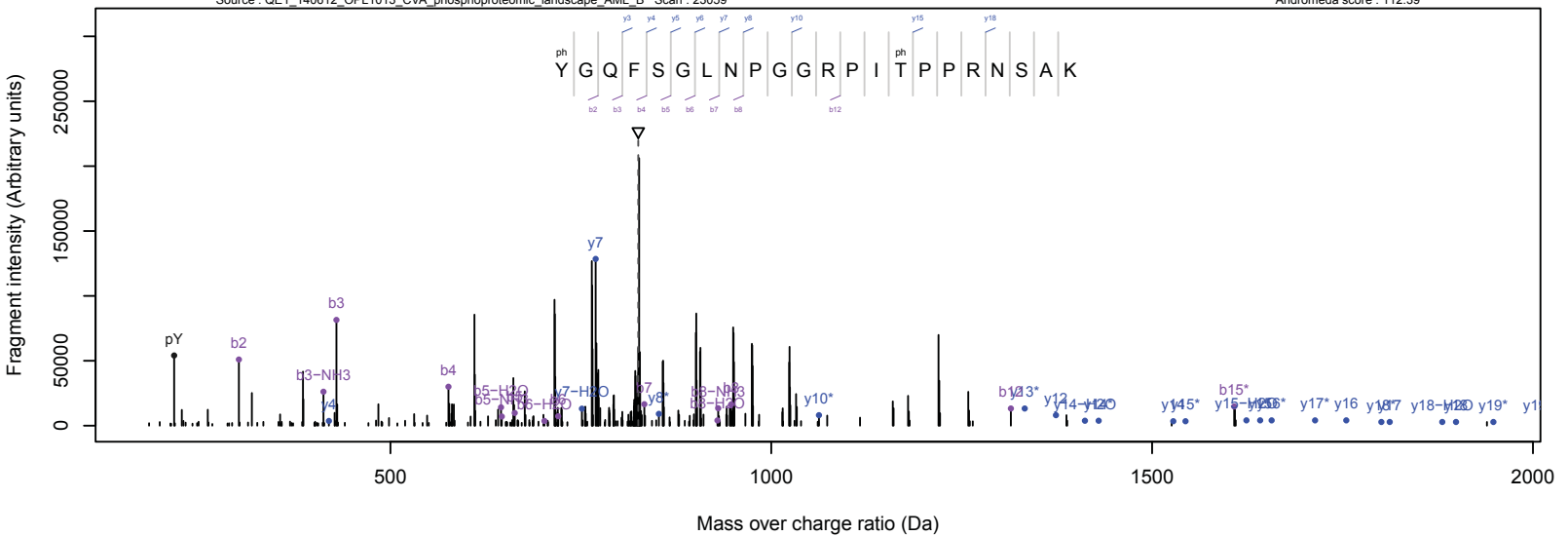
Andromeda score: 112.41



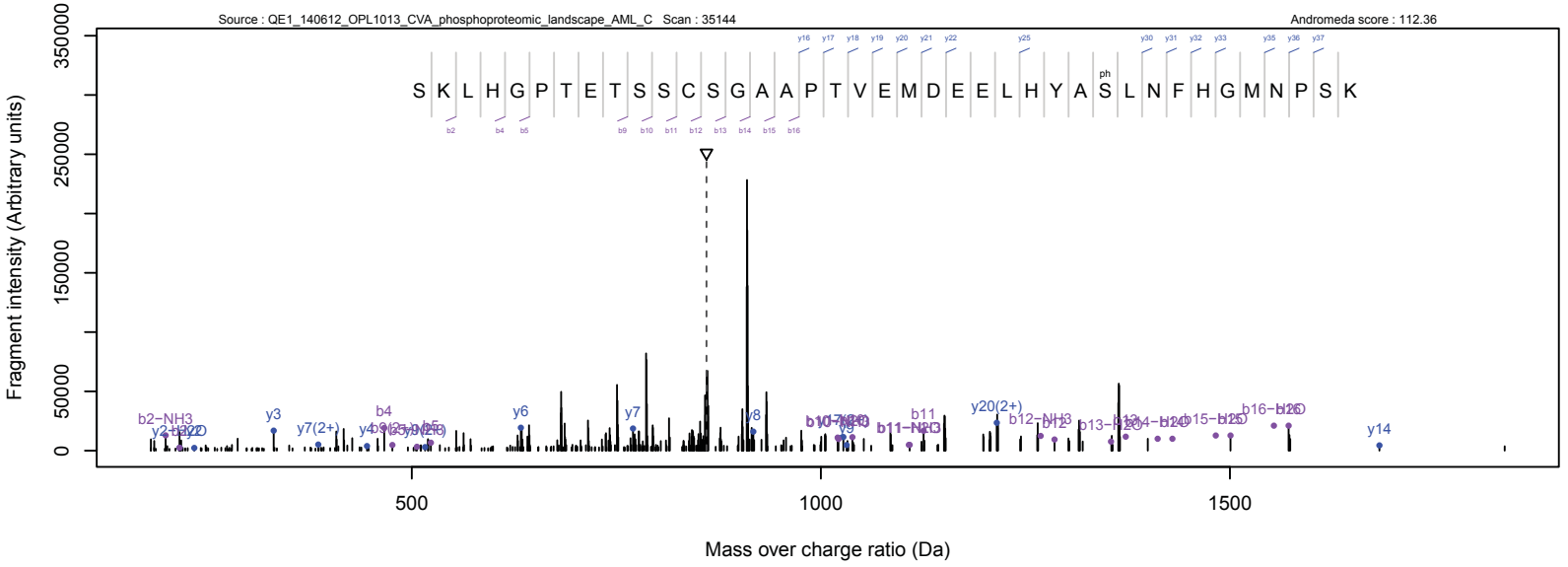
Y(ph)GQFSLNPGGRPI(ph)PPRSAK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 23059

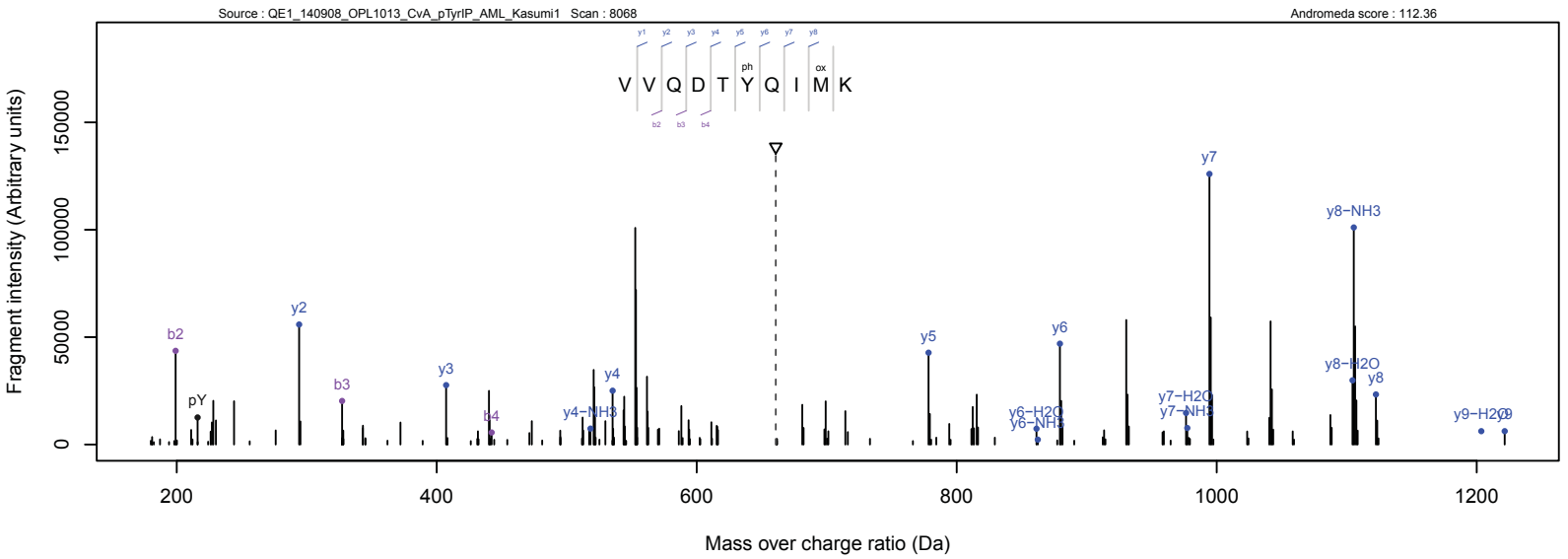
Andromeda score: 112.39



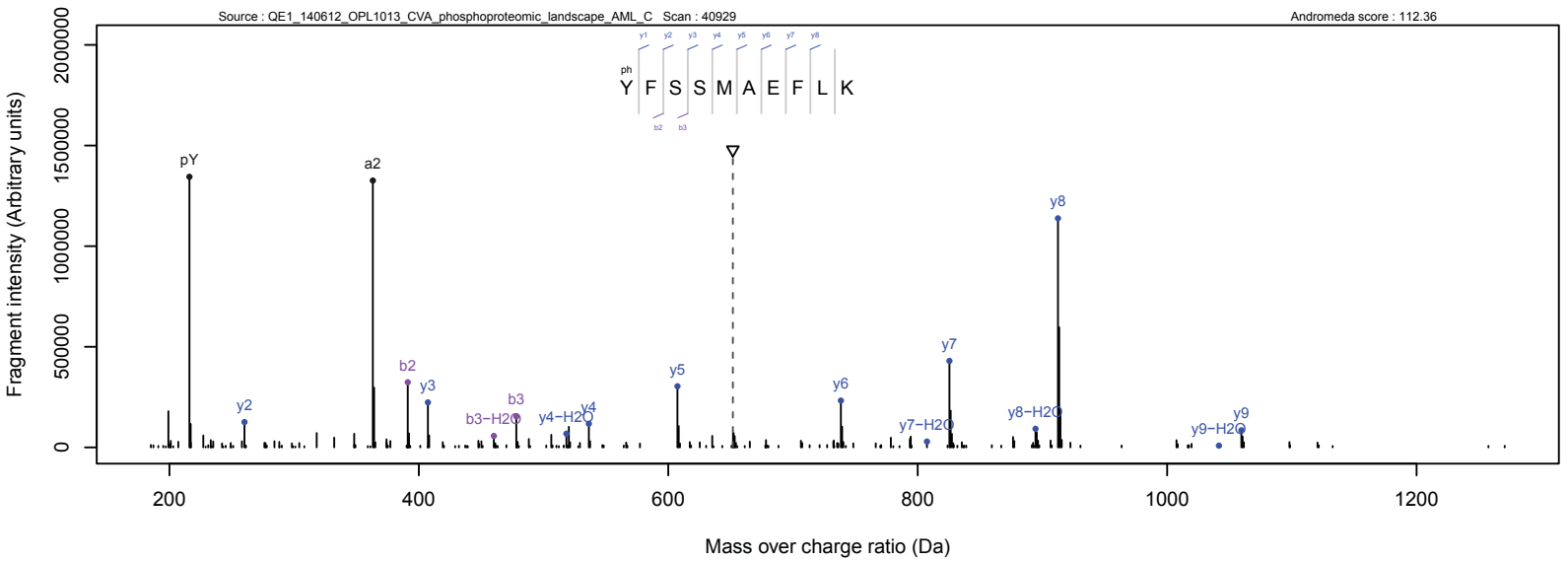
SKLHGPTETSSCSGAAPTVEEMDEELHYASLNFHGMNPSK-5+



VVQDTYQIMK-2+



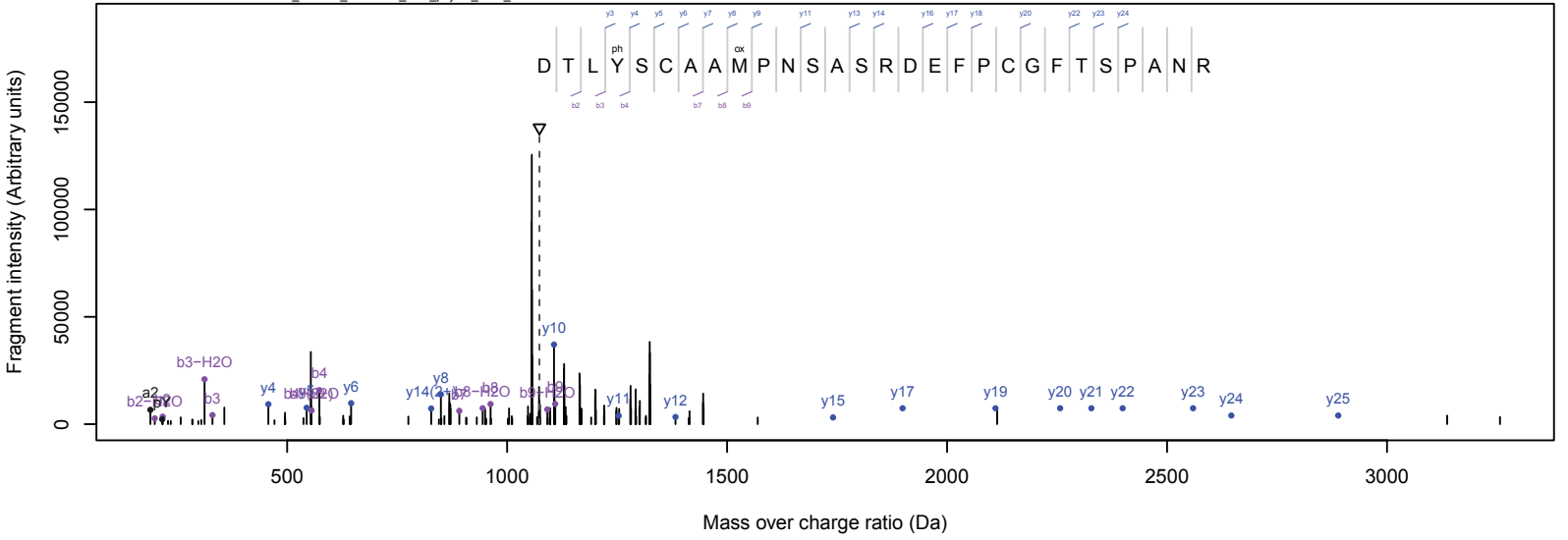
Y(ph)FSSMAEFLK-2+



DTLY(ph)SCAAM(ox)PNSASRDEFPCGFTSPANR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31983

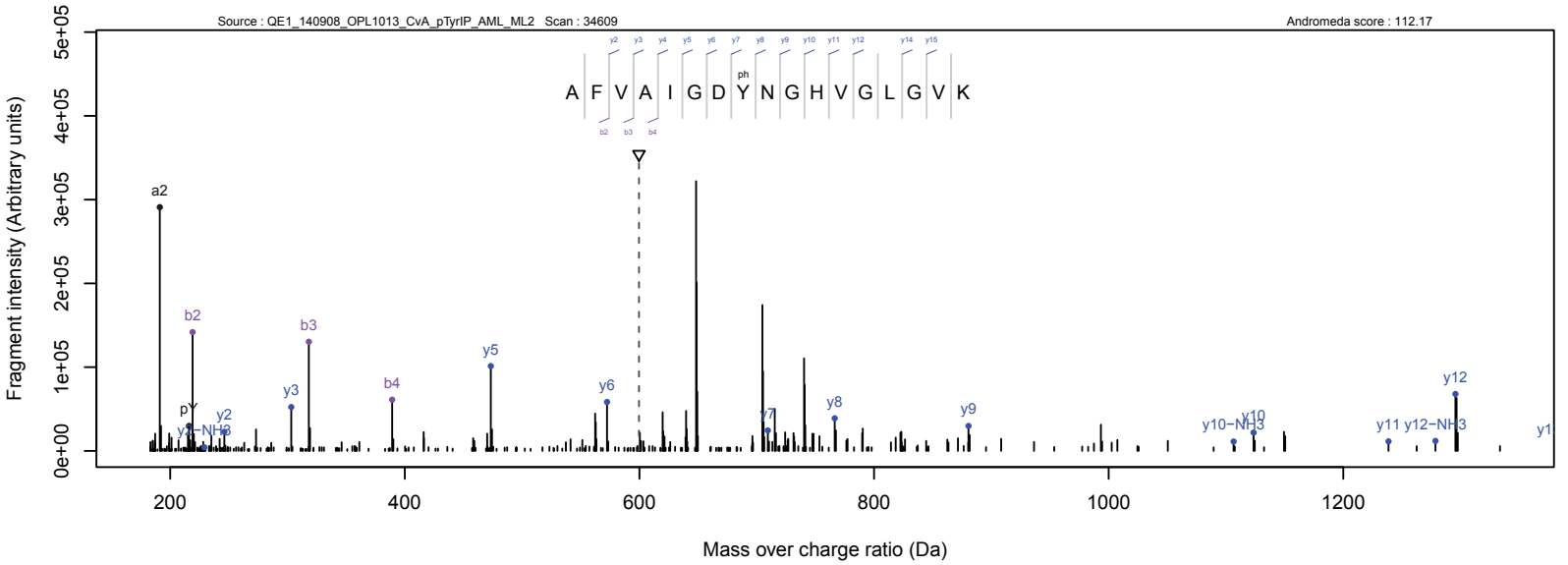
Andromeda score : 112.34



AFVAIGDY(ph)NGHVLGVK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 34609

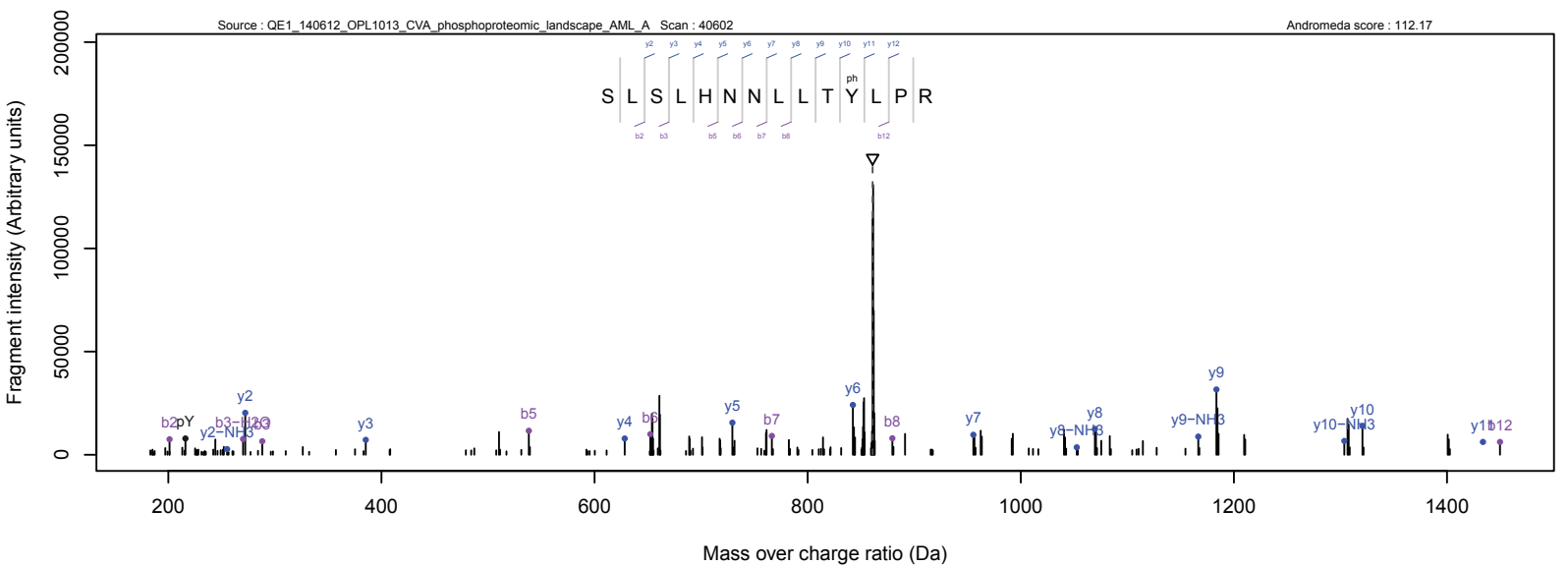
Andromeda score : 112.17



SLSLHNNLLTY(ph)LPR-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_A Scan : 40602

Andromeda score : 112.17

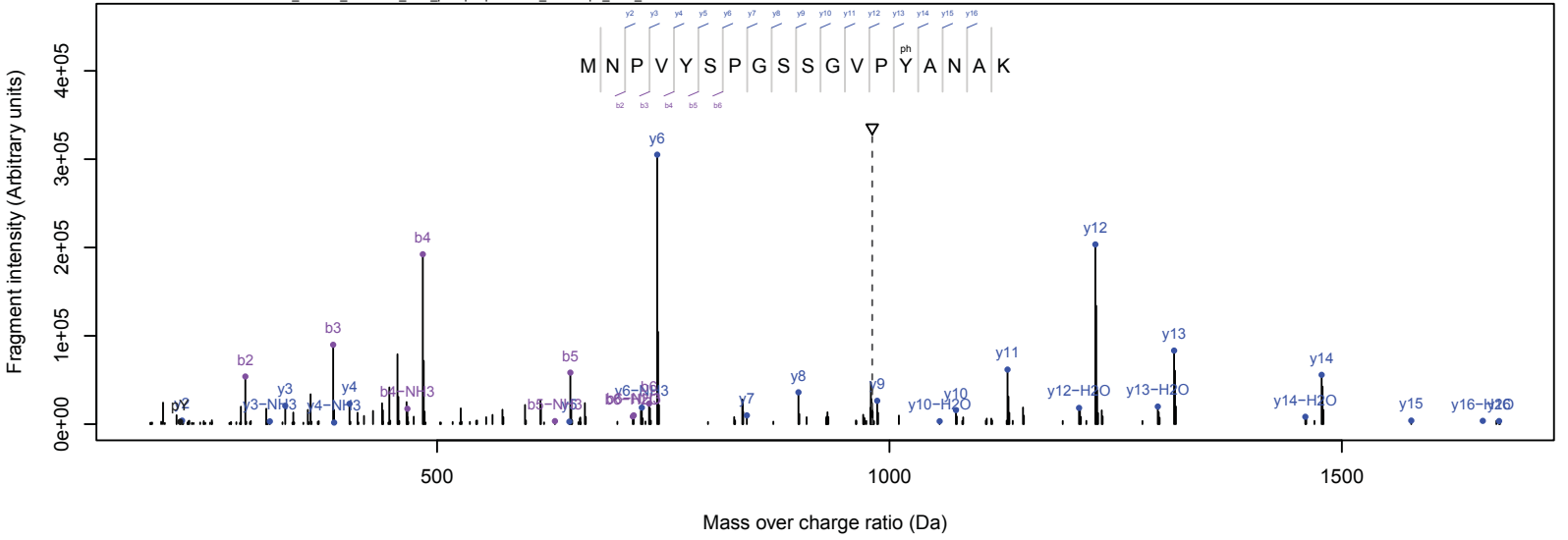




(ac)MNPVYSPGSSGVYPYANA K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31612

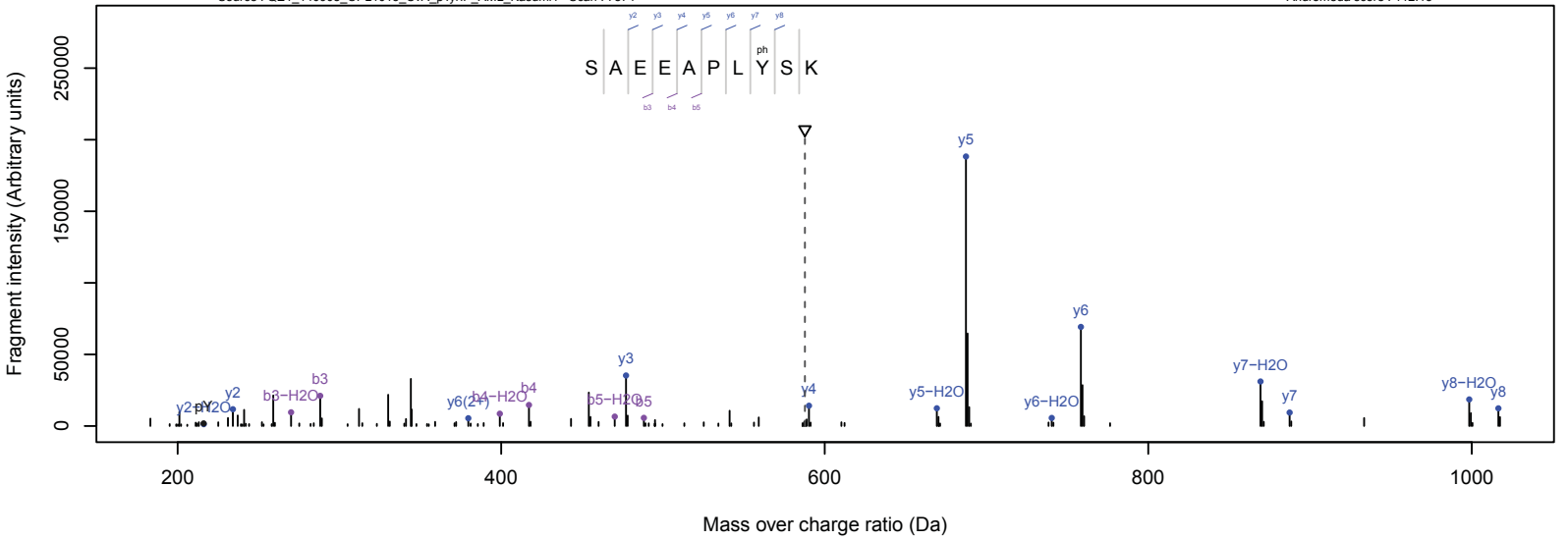
Andromeda score : 112.16



SAEEAPLYSK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 7874

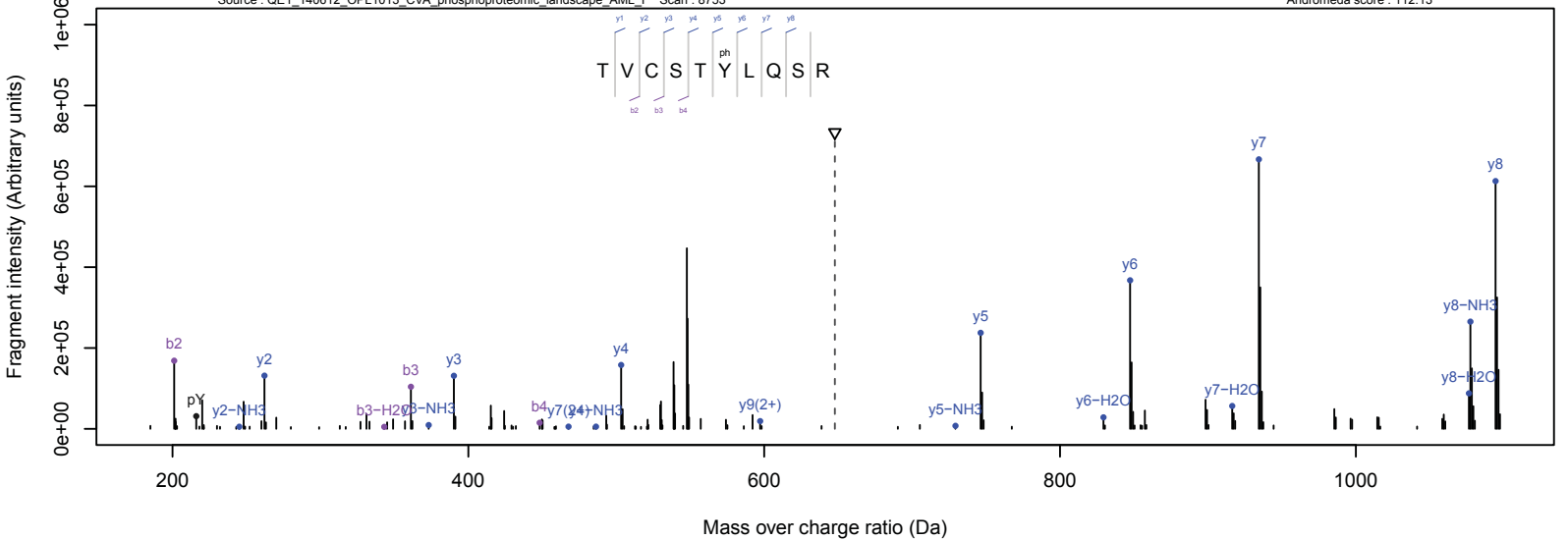
Andromeda score : 112.13



TVCS TYLQSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 8753

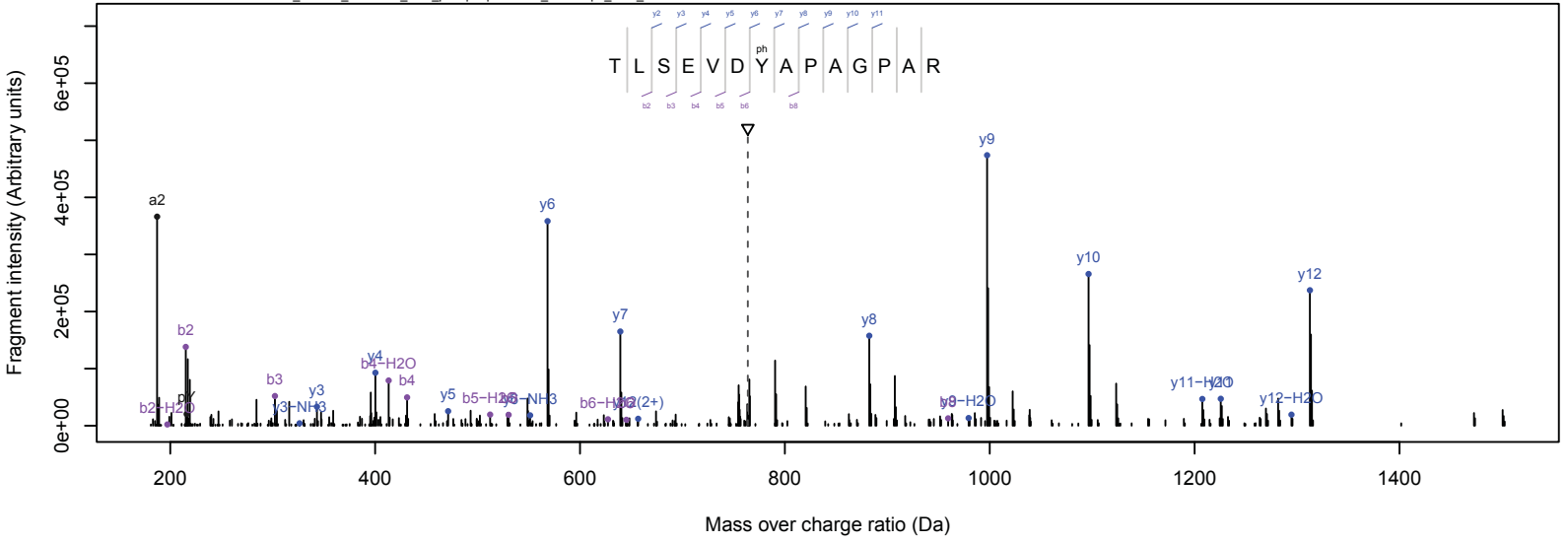
Andromeda score : 112.13



TLSEVDY(ph)APAGPAR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17568

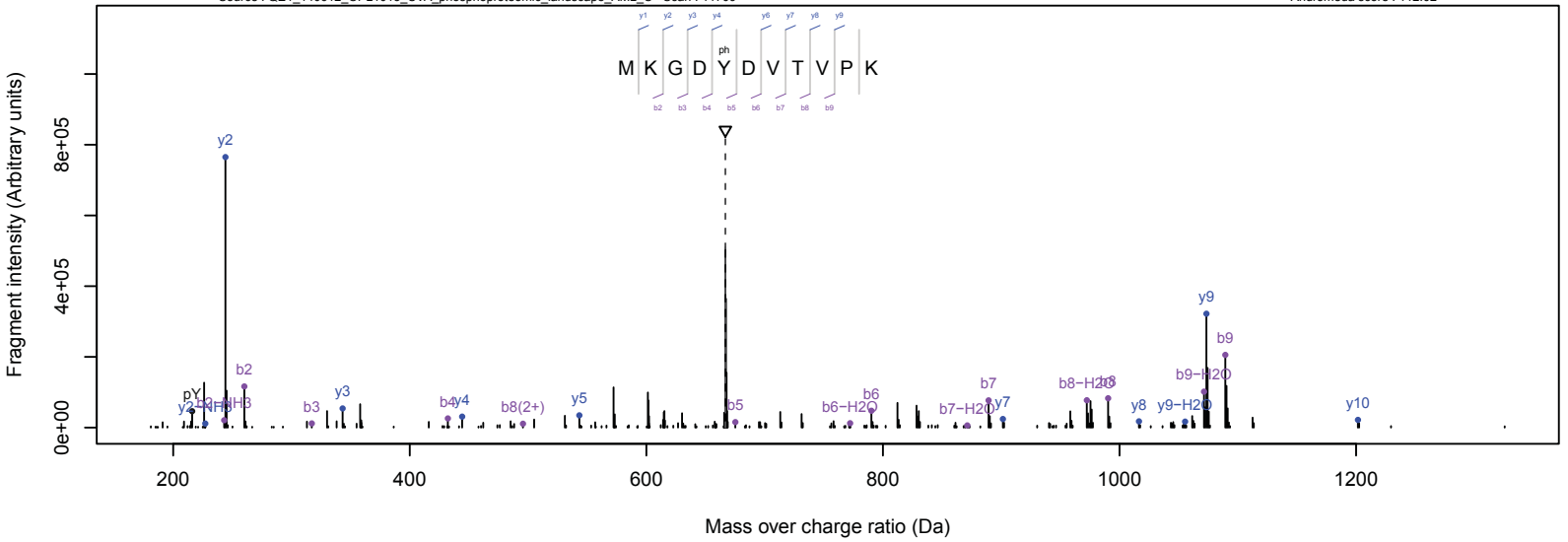
Andromeda score: 112.08



MKGDY(ph)DVTVPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11709

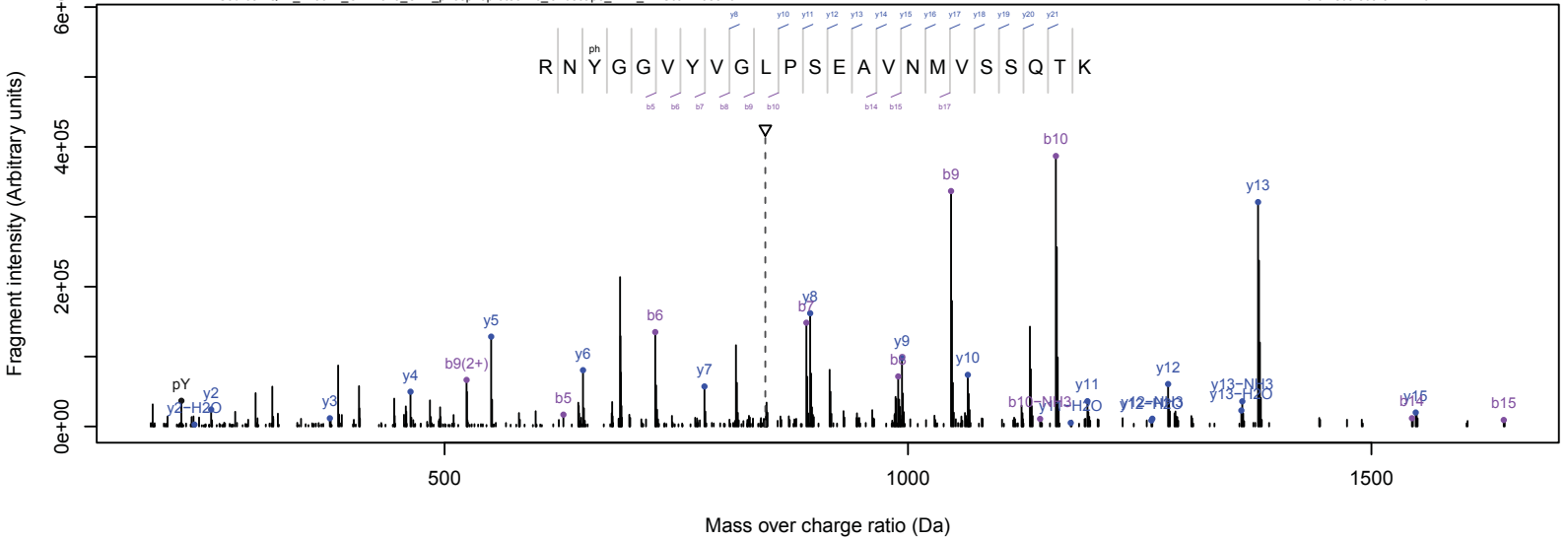
Andromeda score: 112.02



RNY(ph)GGVYVGLPSEAVNMVSSQTK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 38876

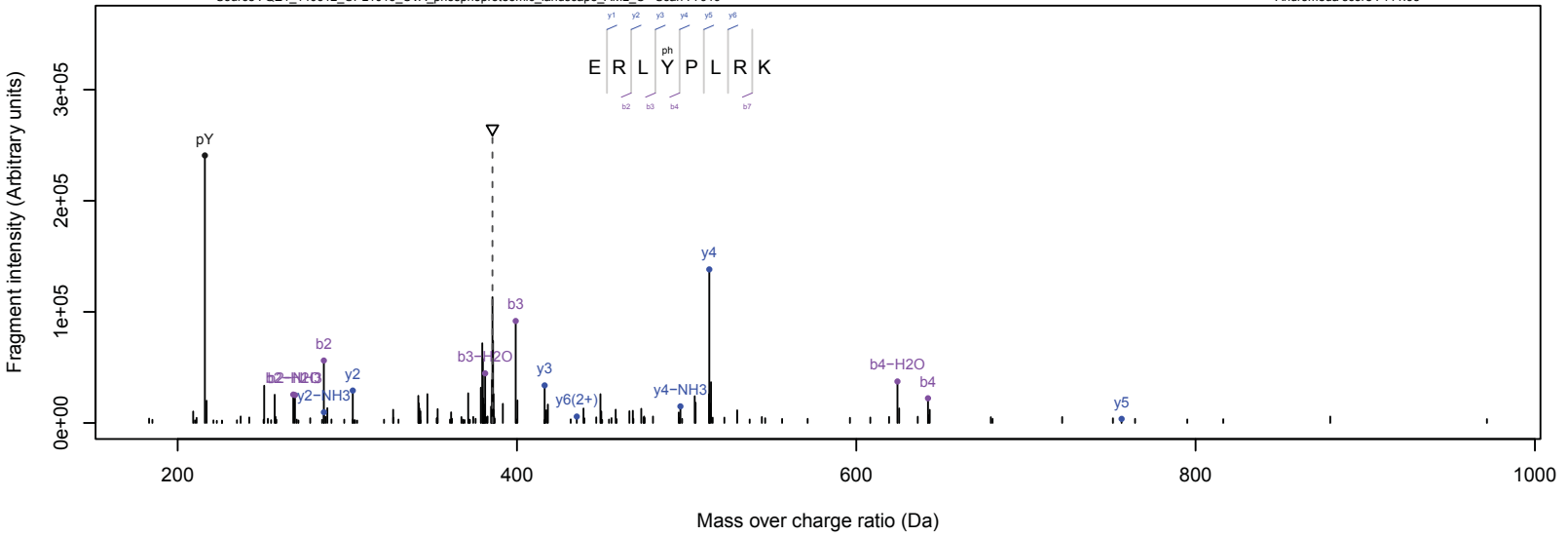
Andromeda score: 111.97



ERLY(ph)PLRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7919

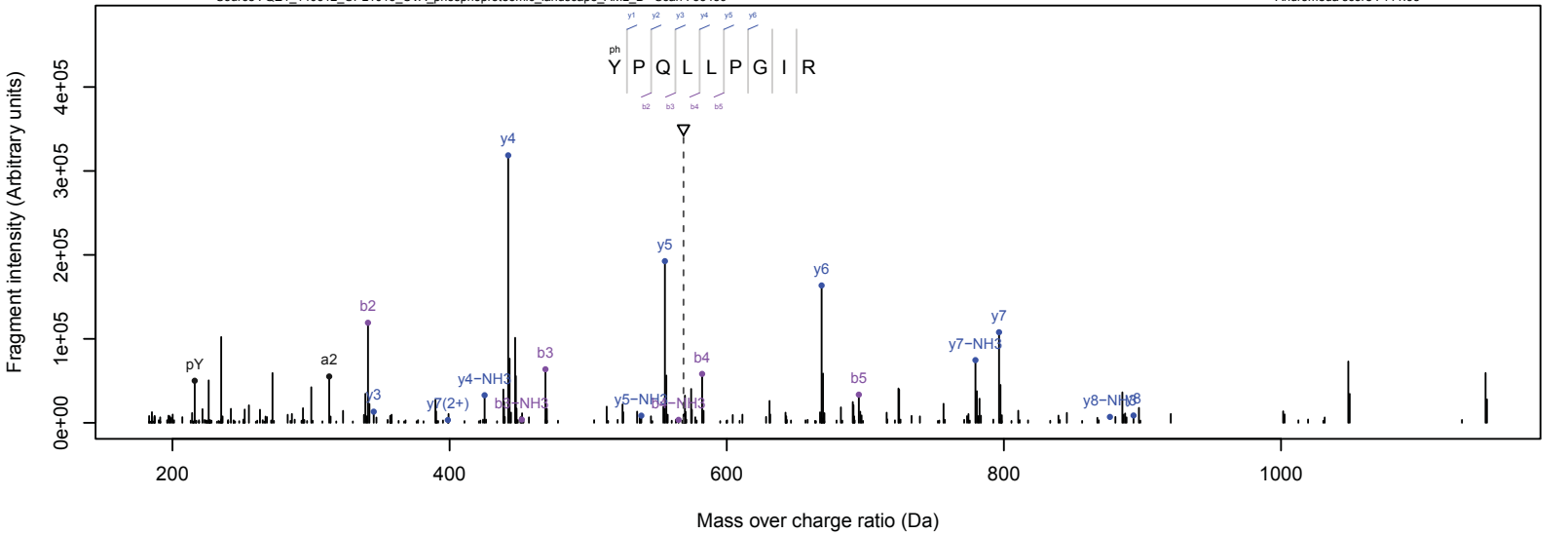
Andromeda score : 111.95



Y(ph)PQLLPGIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33439

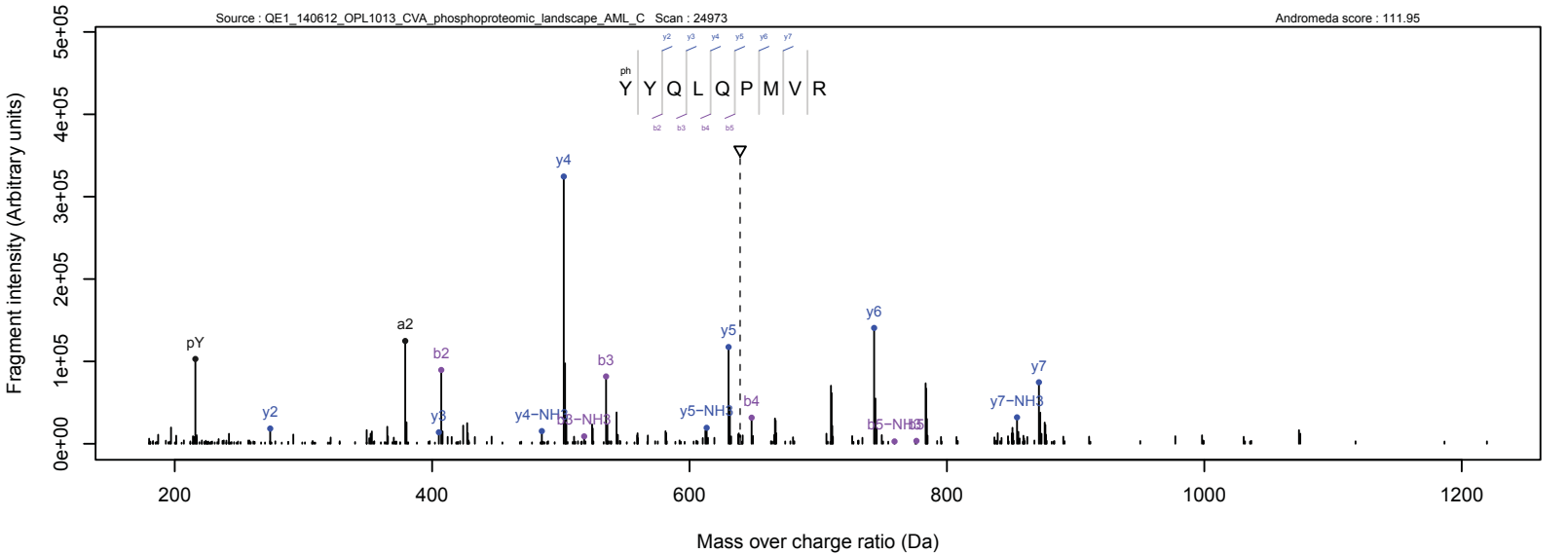
Andromeda score : 111.95



Y(ph)YQLQPMVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24973

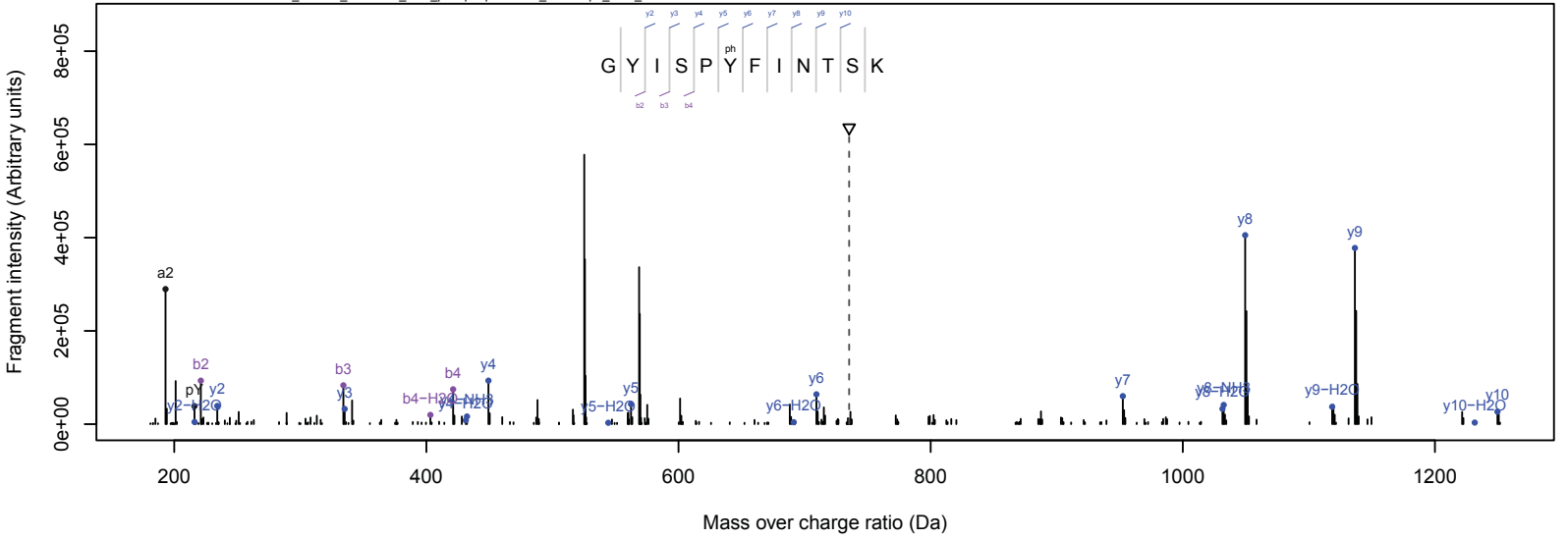
Andromeda score : 111.95



**GYISPY(ph)FINTSK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 31292

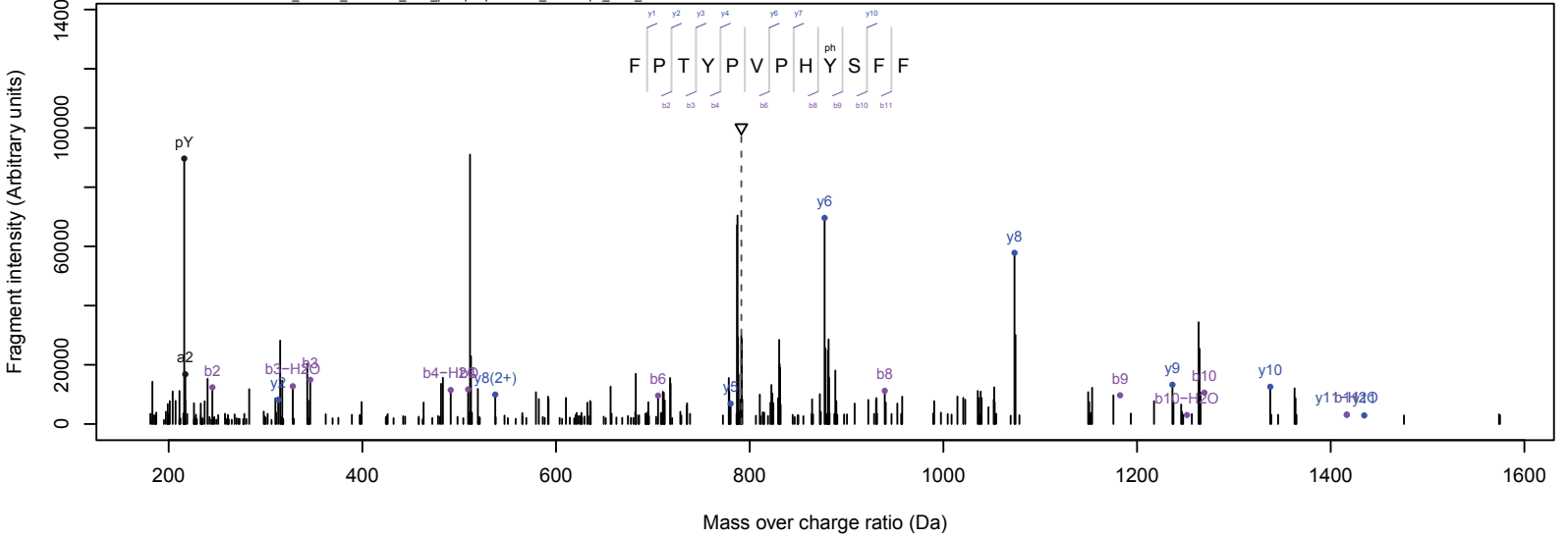
Andromeda score : 111.94



**FPTYVPVPH(ph)SFF-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47759

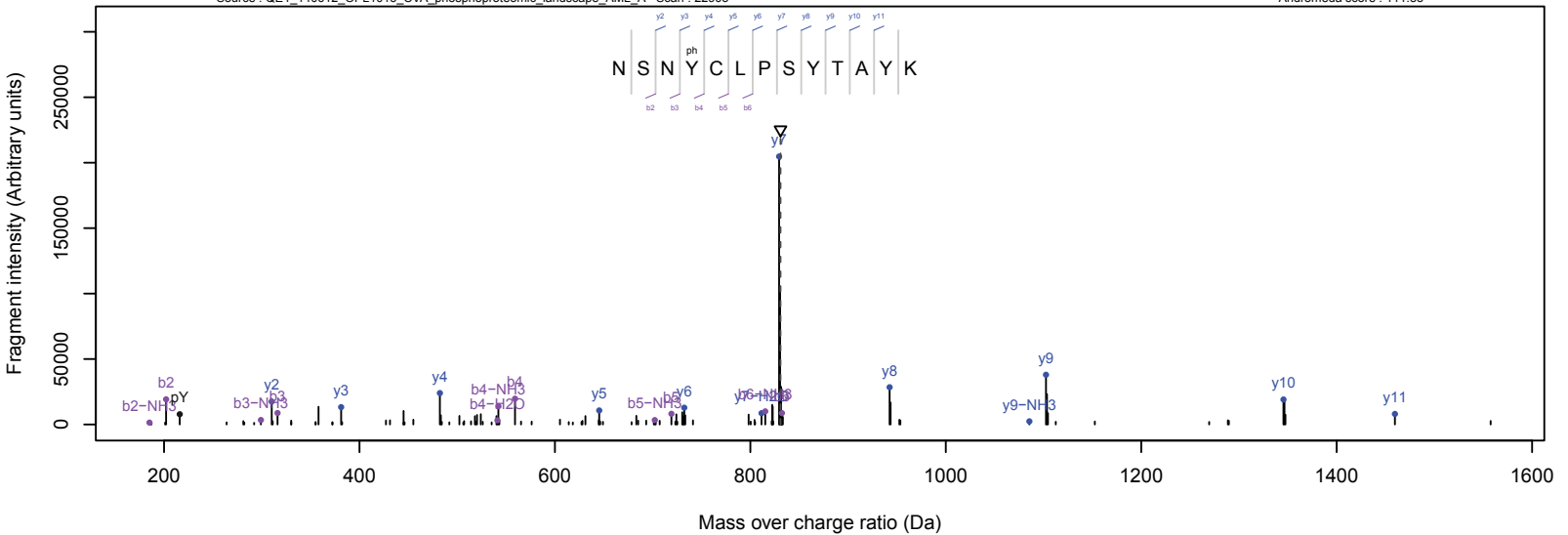
Andromeda score : 111.86



**NSNY(ph)CLPSYTAYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 22903

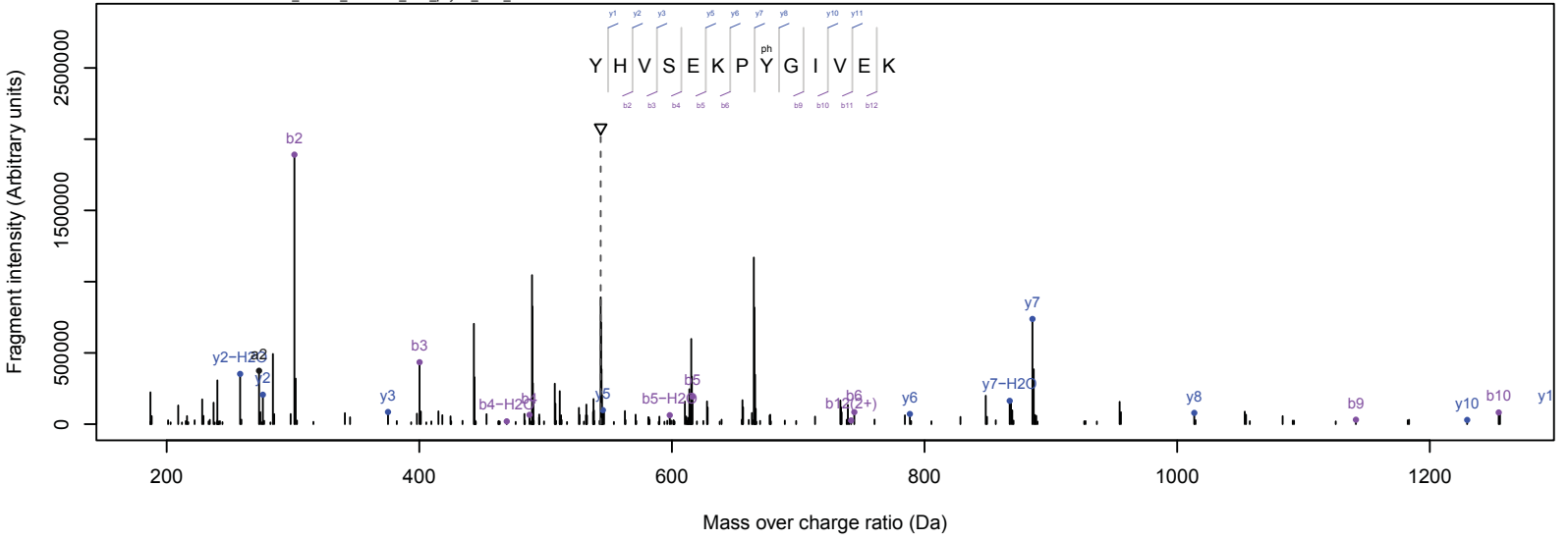
Andromeda score : 111.83



**YHVSEKPY(ph)GIVEK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 13763

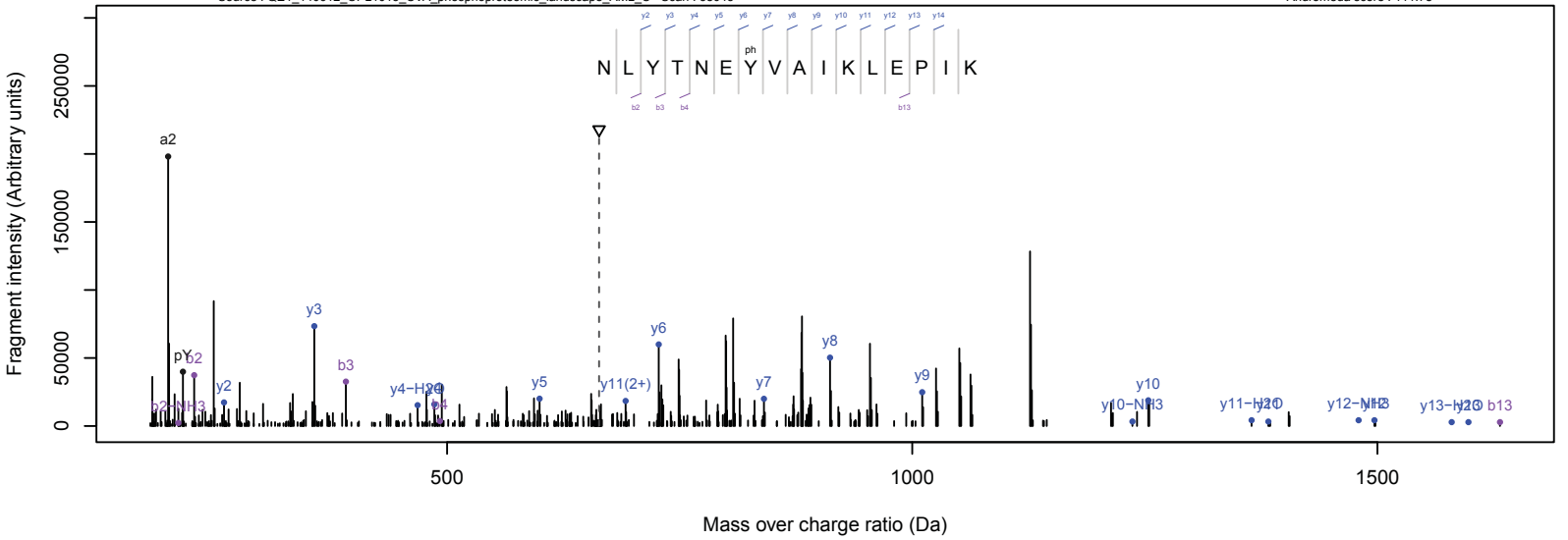
Andromeda score : 111.83



**NLYTNEY(ph)VAIKLEPIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38043

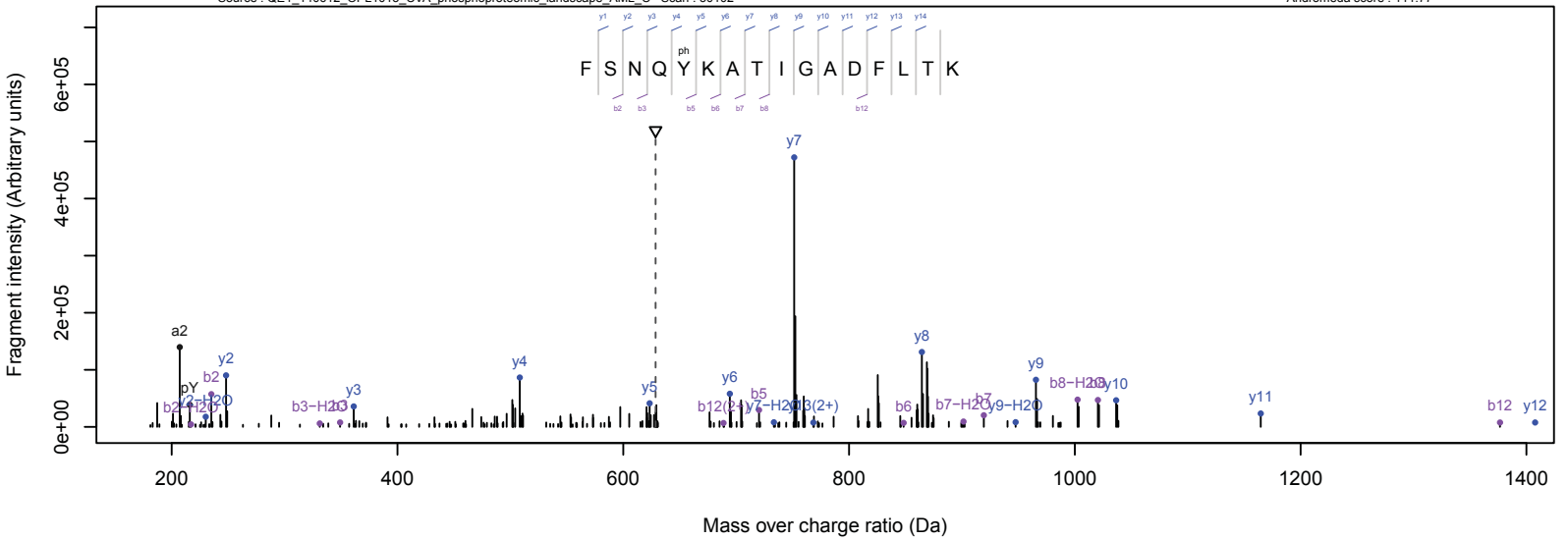
Andromeda score : 111.78



**FSNQY(ph)KATIGADFLTK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30102

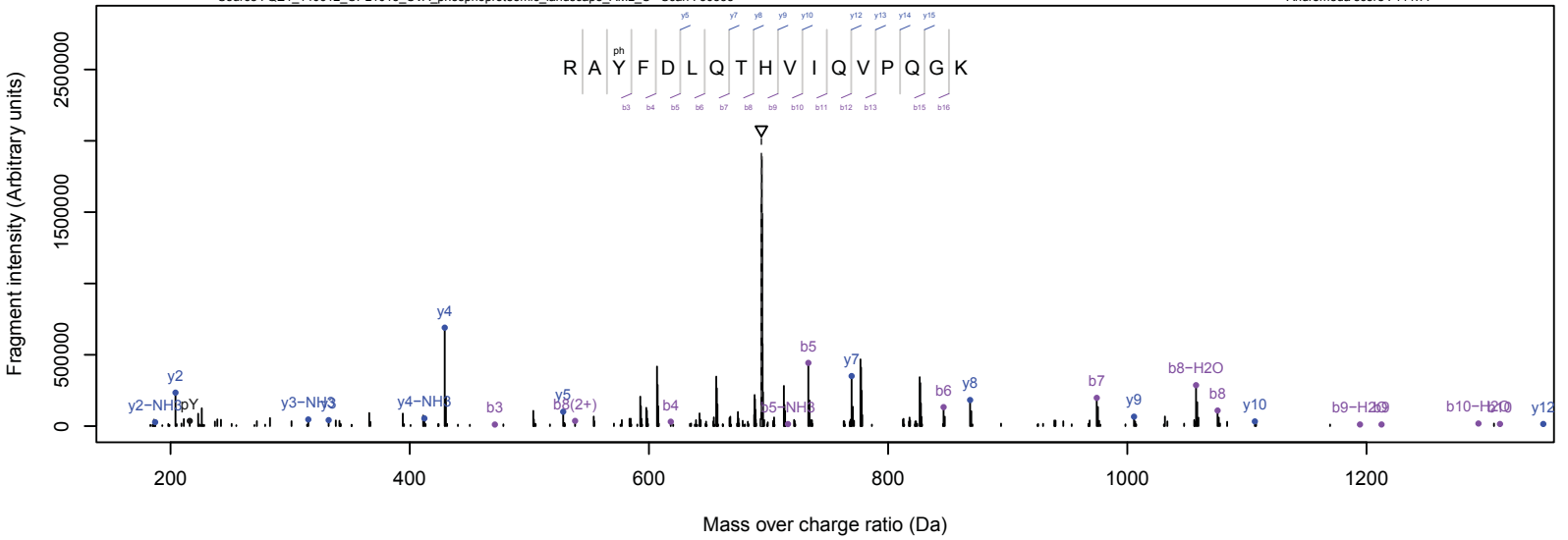
Andromeda score : 111.77



RAY(ph)FDLQTHVIQVPQGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30553

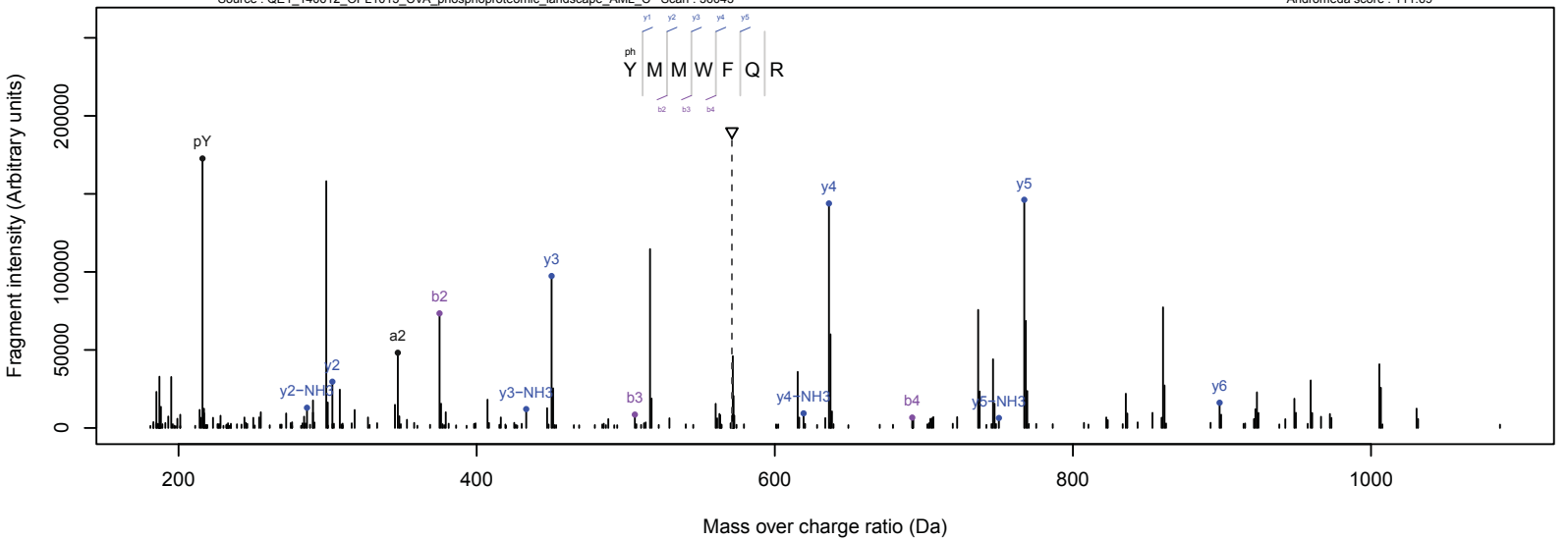
Andromeda score : 111.77



Y(ph)MMWFQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36043

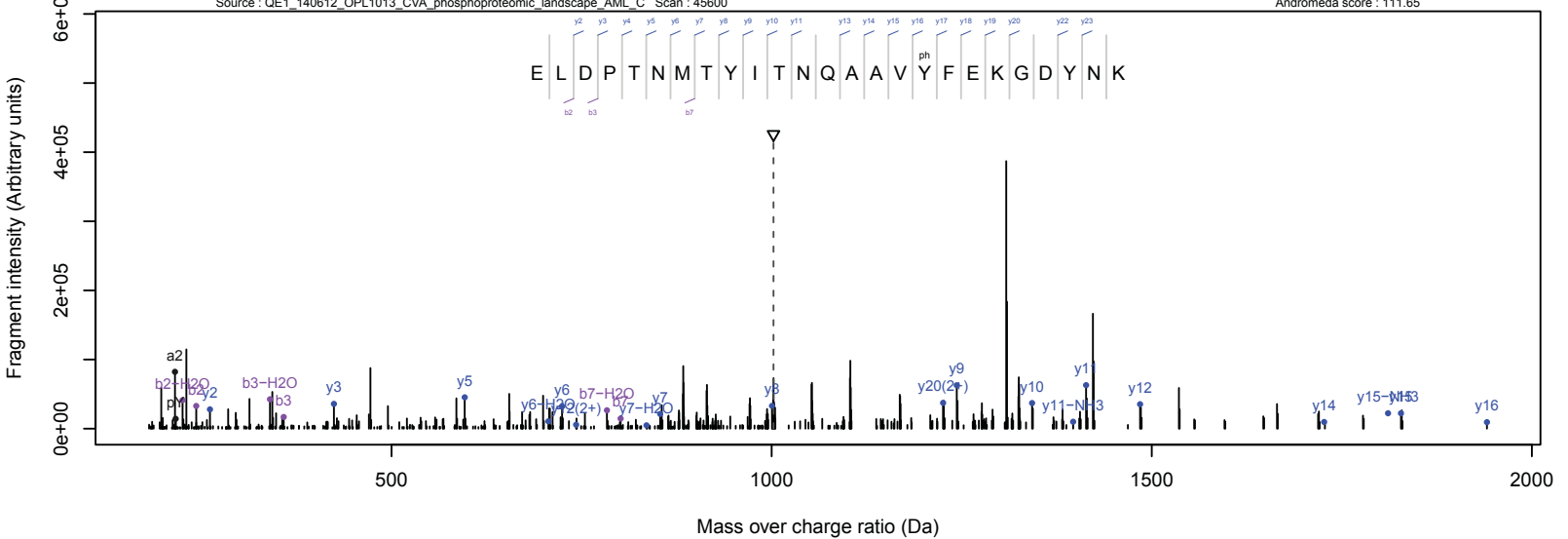
Andromeda score : 111.69



ELDPTNMTYITNQAAVY(ph)FEKGDYNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45600

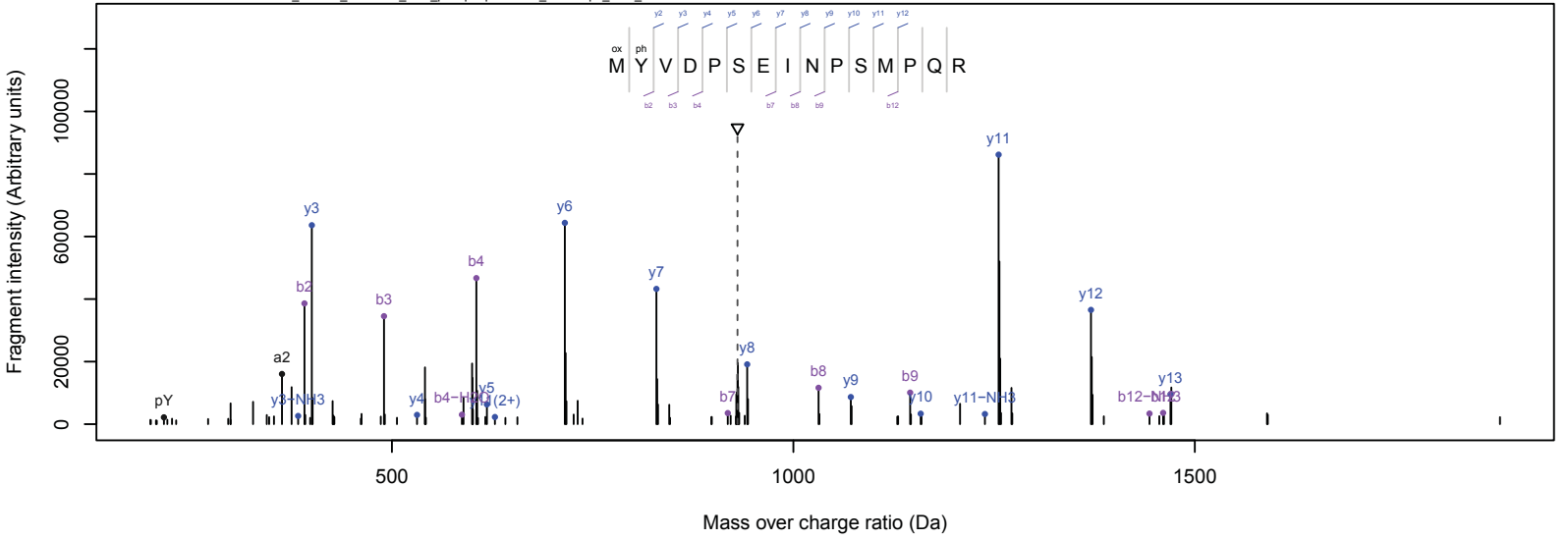
Andromeda score : 111.65



**M(ox)Y(ph)VDPSEINPSPMPQR**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 24276

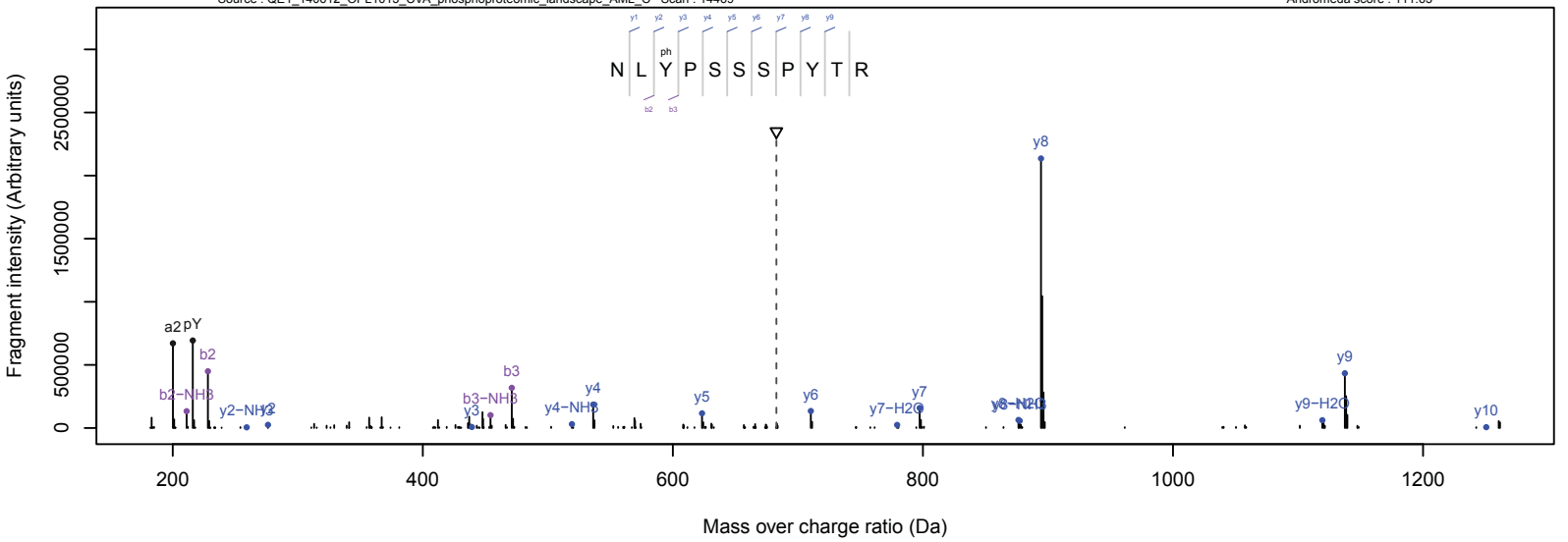
Andromeda score: 111.64



**NLY(ph)PSSSPYTR**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14469

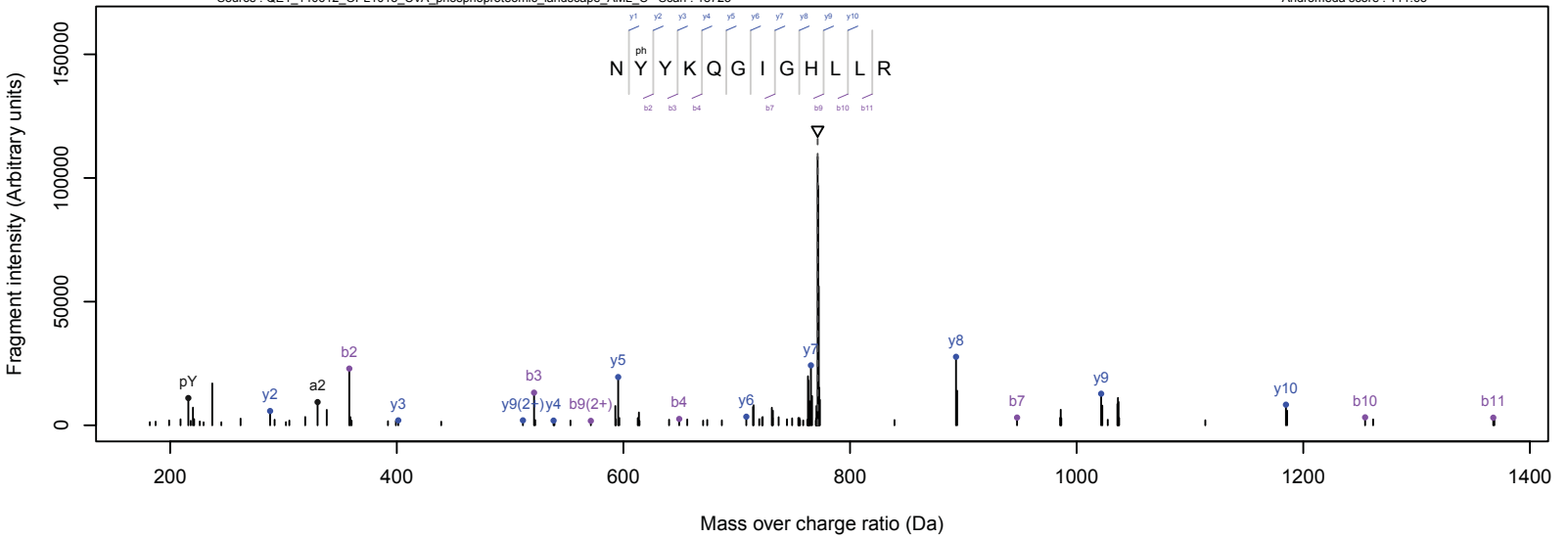
Andromeda score: 111.63



**NY(ph)YKQGIGHLLR**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18723

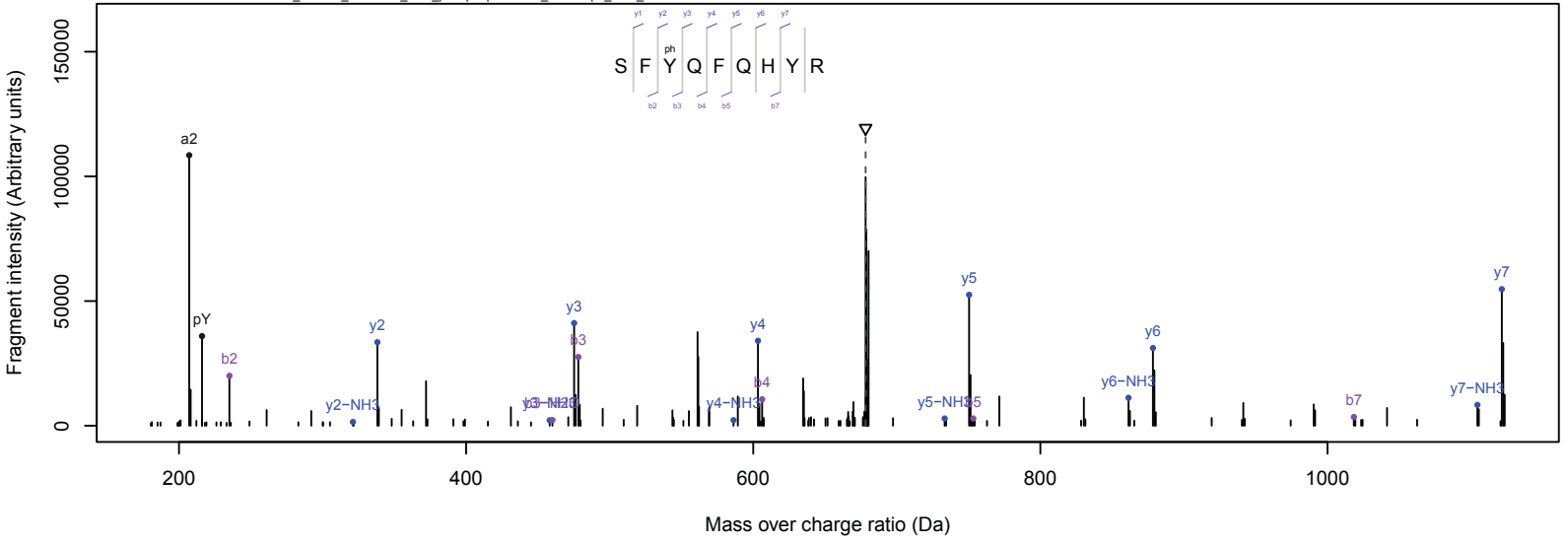
Andromeda score: 111.63



SFY(ph)QFQHYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 20144

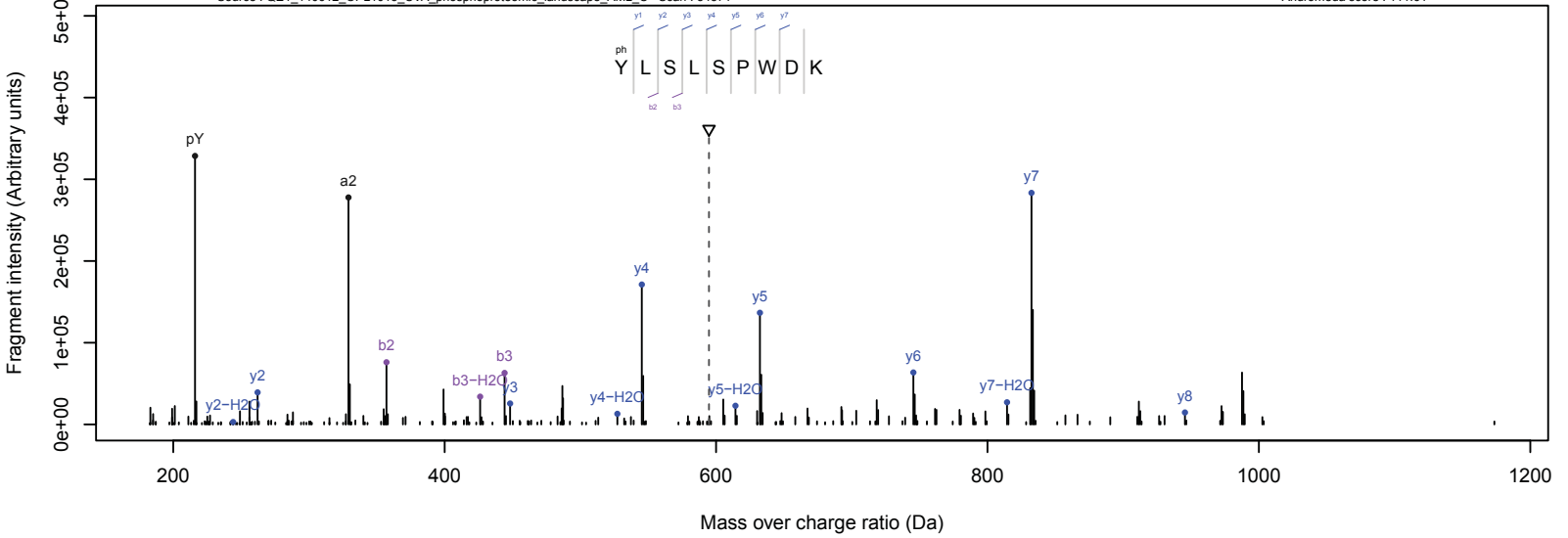
Andromeda score : 111.61



Y(ph)LSLSPWDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34571

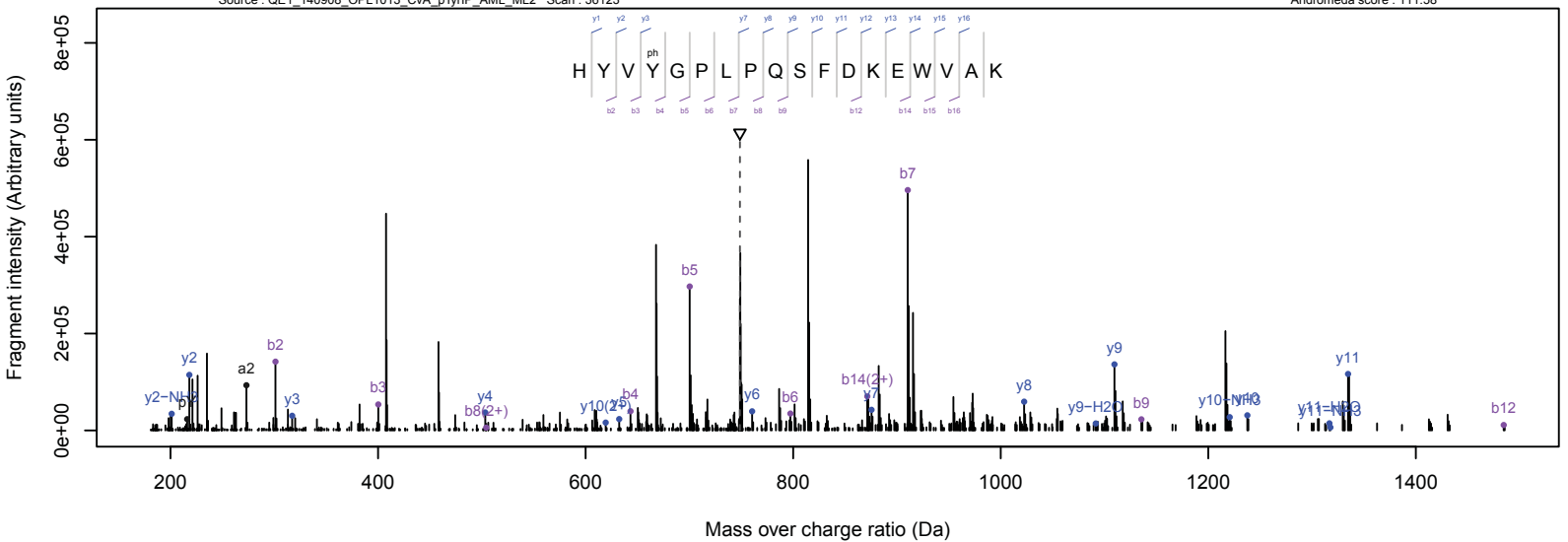
Andromeda score : 111.61



HYVY(ph)GPLQSFQK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 36123

Andromeda score : 111.58

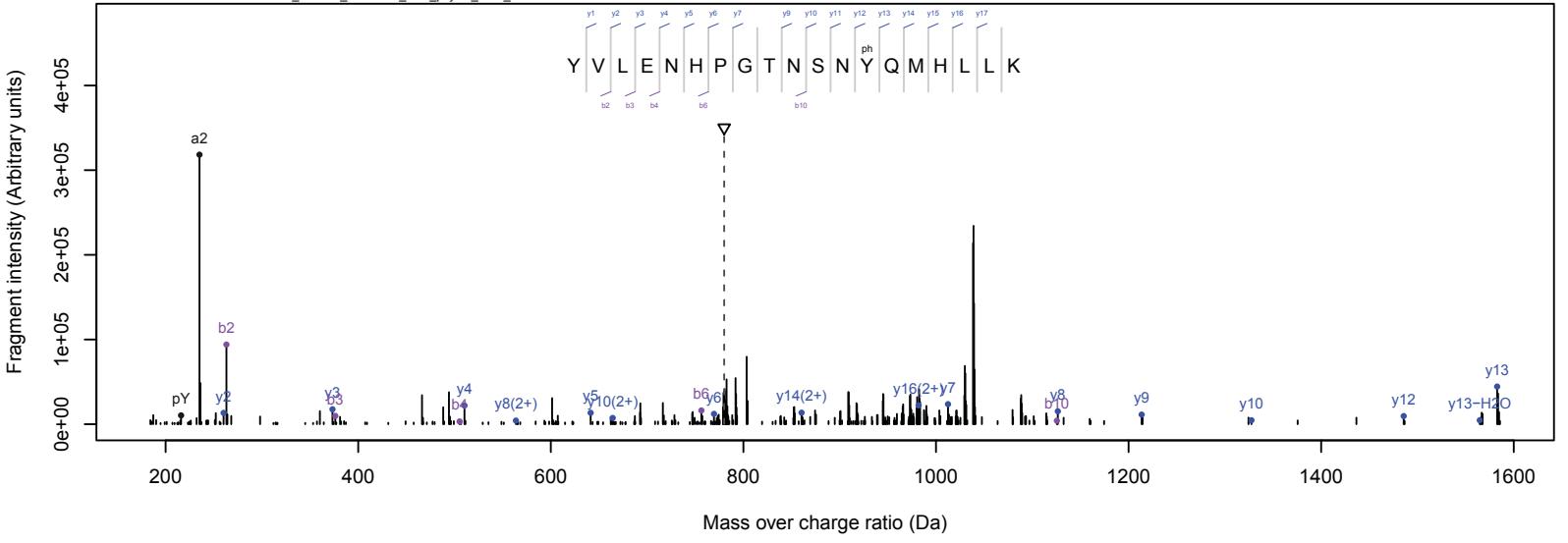




**YVLEHPTNSNY(ph)QMHLK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 25703

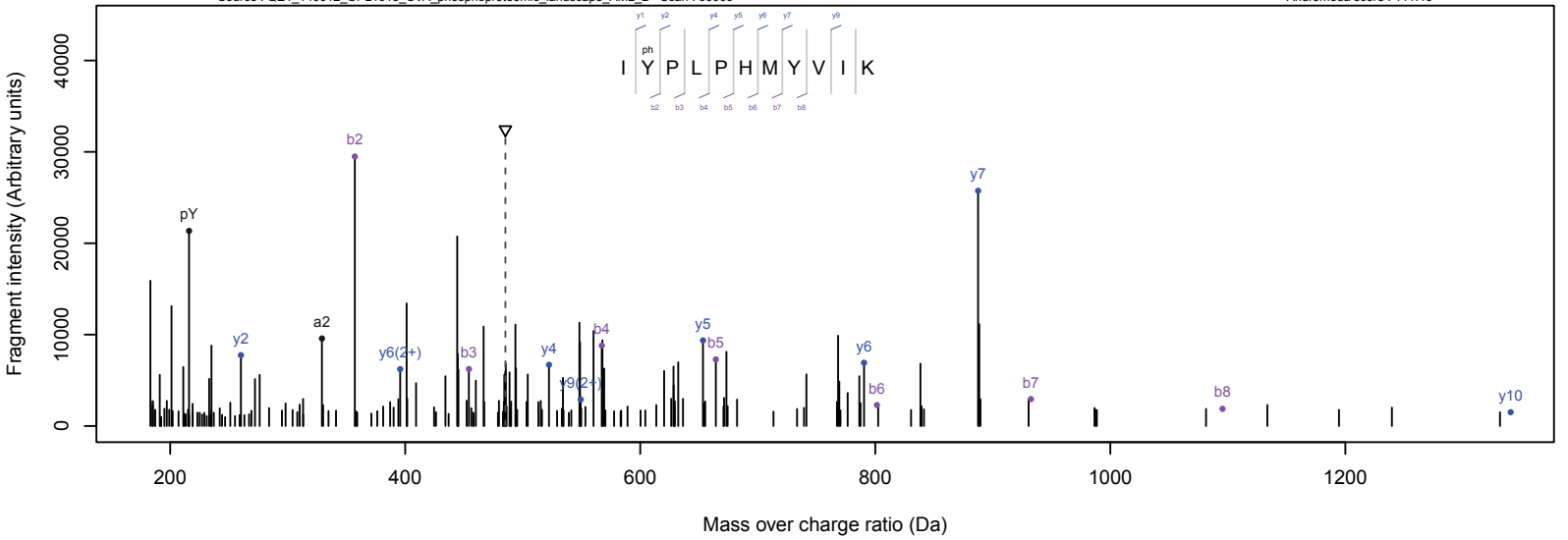
Andromeda score : 111.58



**IY(ph)PLPHMYVIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33969

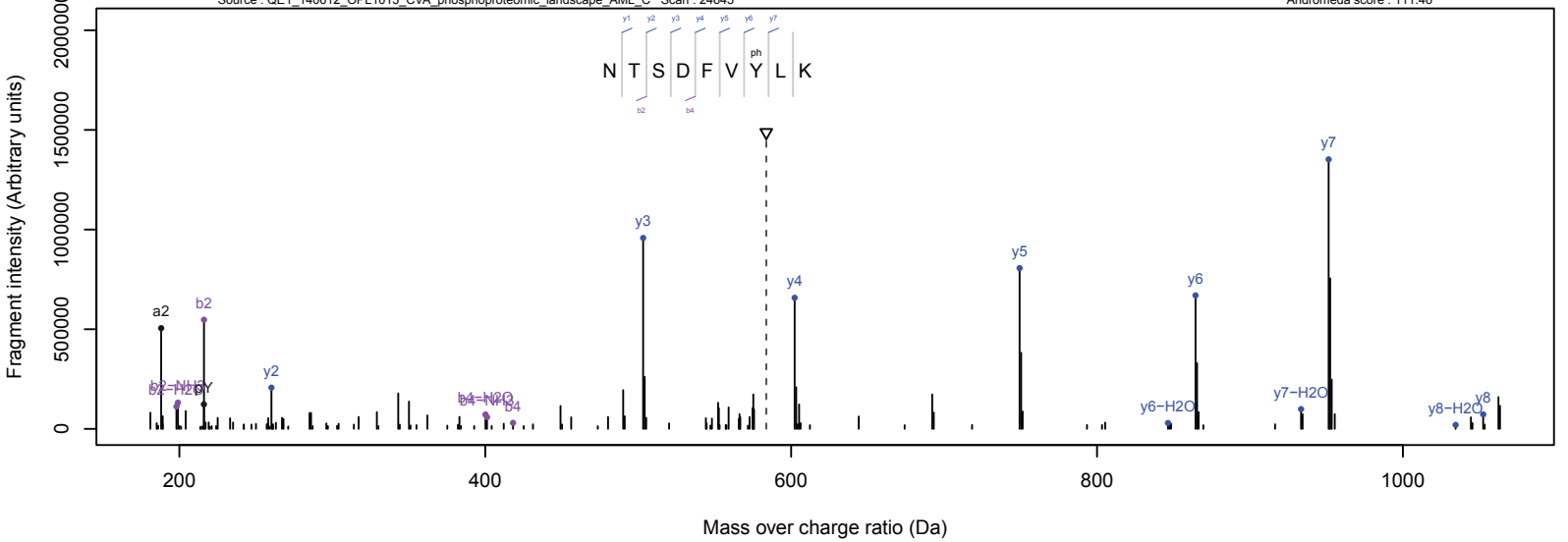
Andromeda score : 111.46



**NTSDFVY(ph)LK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24645

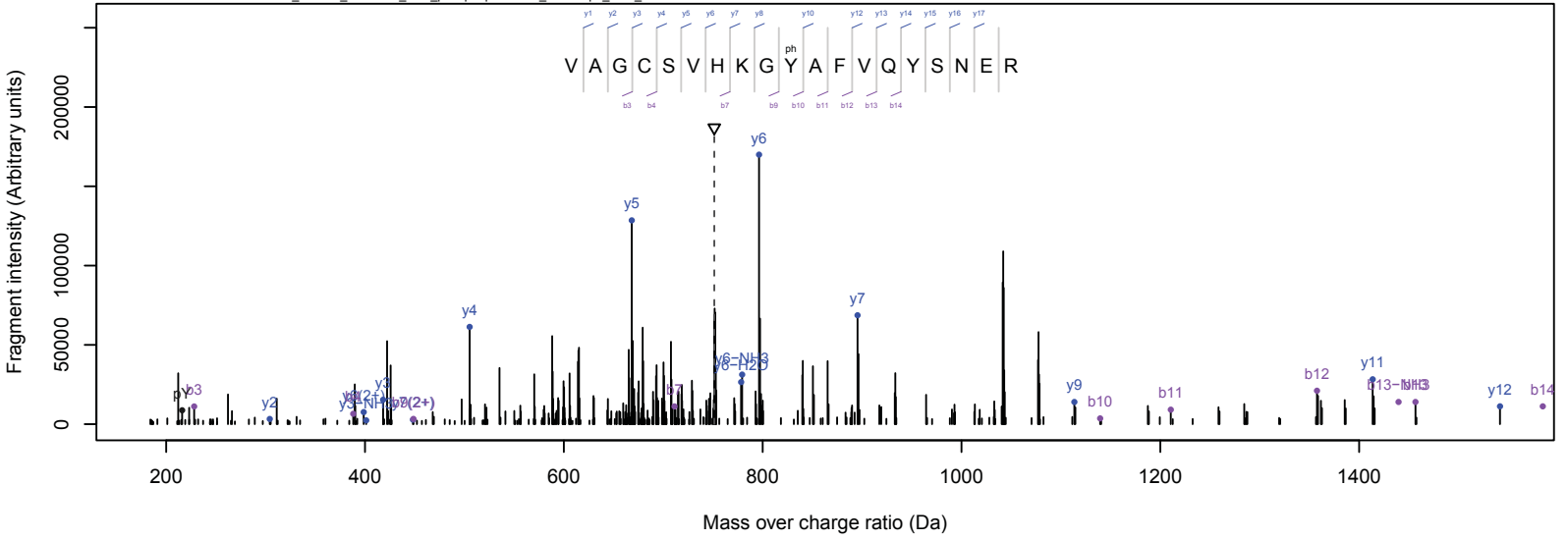
Andromeda score : 111.46



VAGCSVHKGY(ph)AFVQYSNER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15681

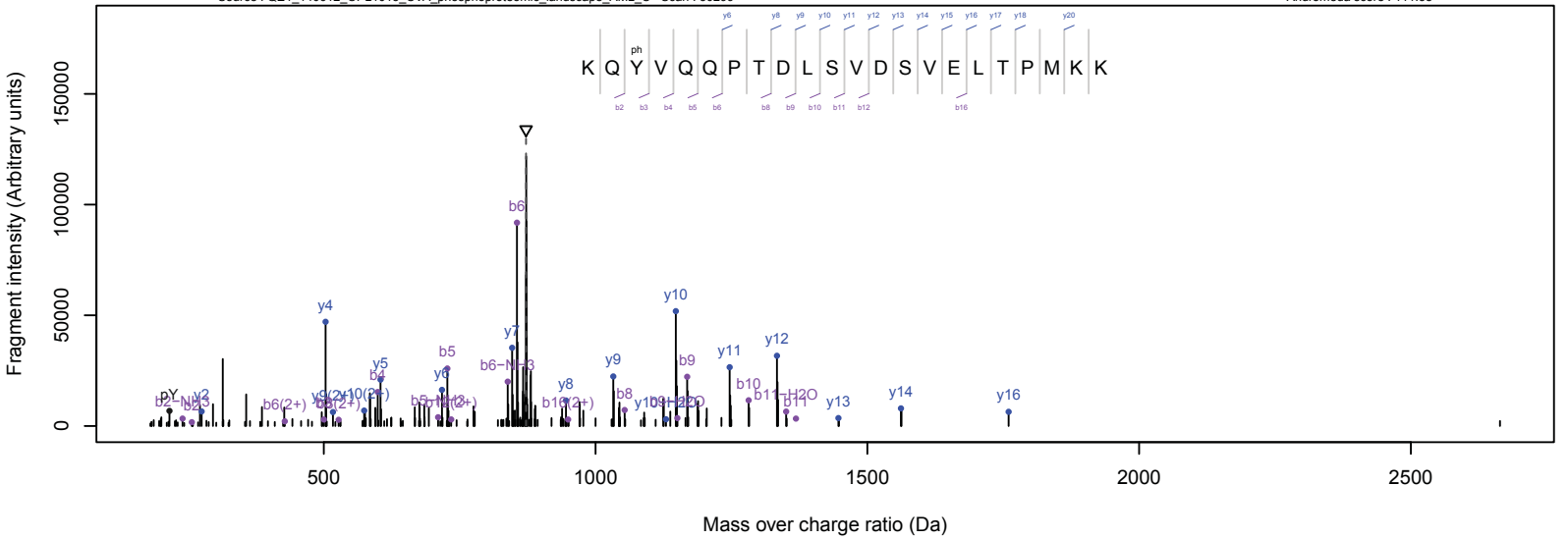
Andromeda score : 111.4



KQY(ph)VQQPTDLSVDSVELTPMKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30299

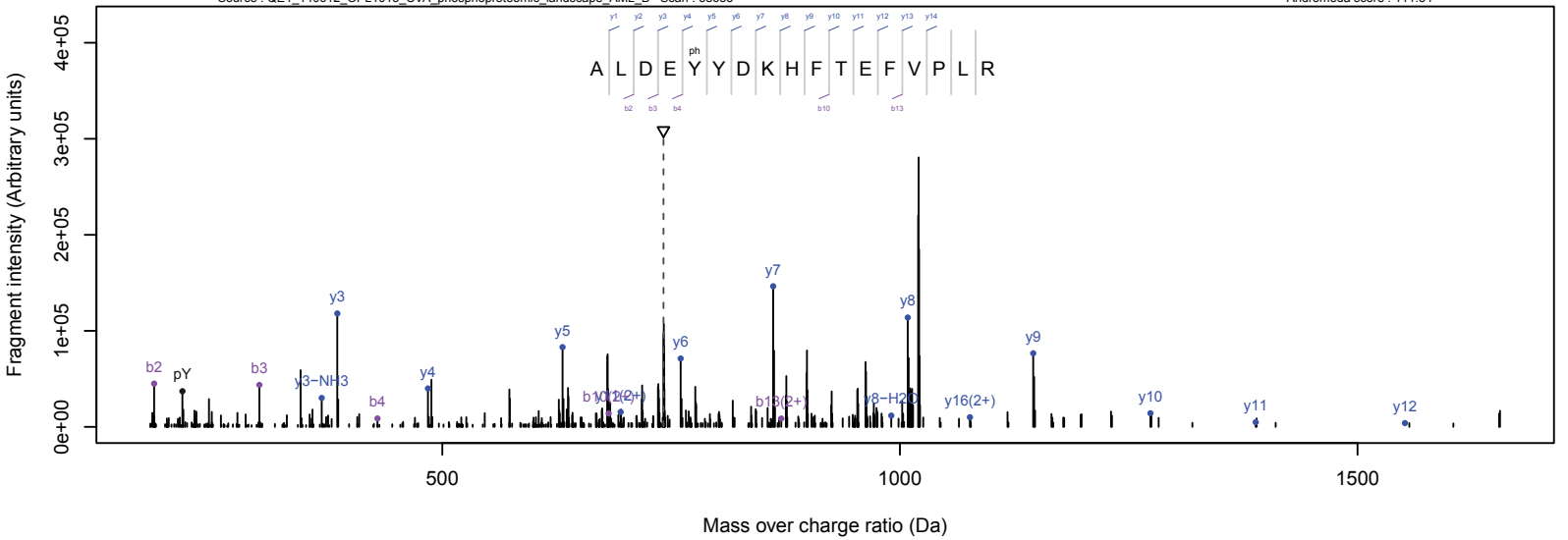
Andromeda score : 111.38



ALDEY(ph)YDKHFTEFVPLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38635

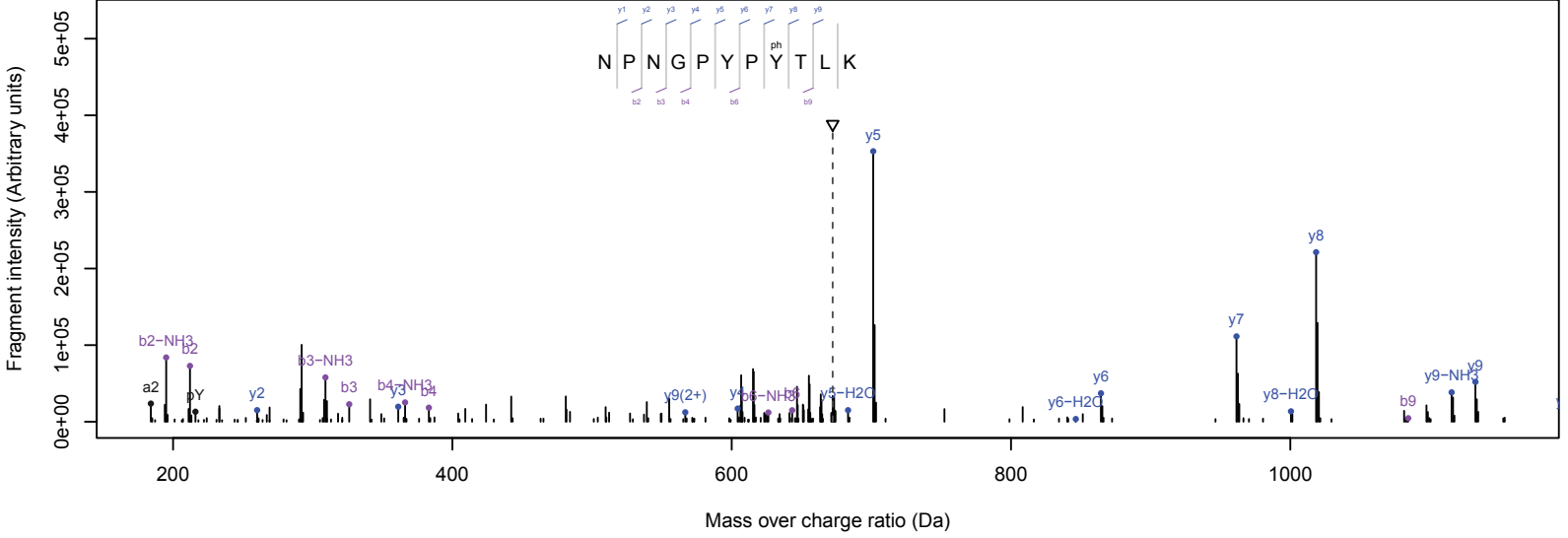
Andromeda score : 111.34



**NPNGPYPY(ph)TLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16672

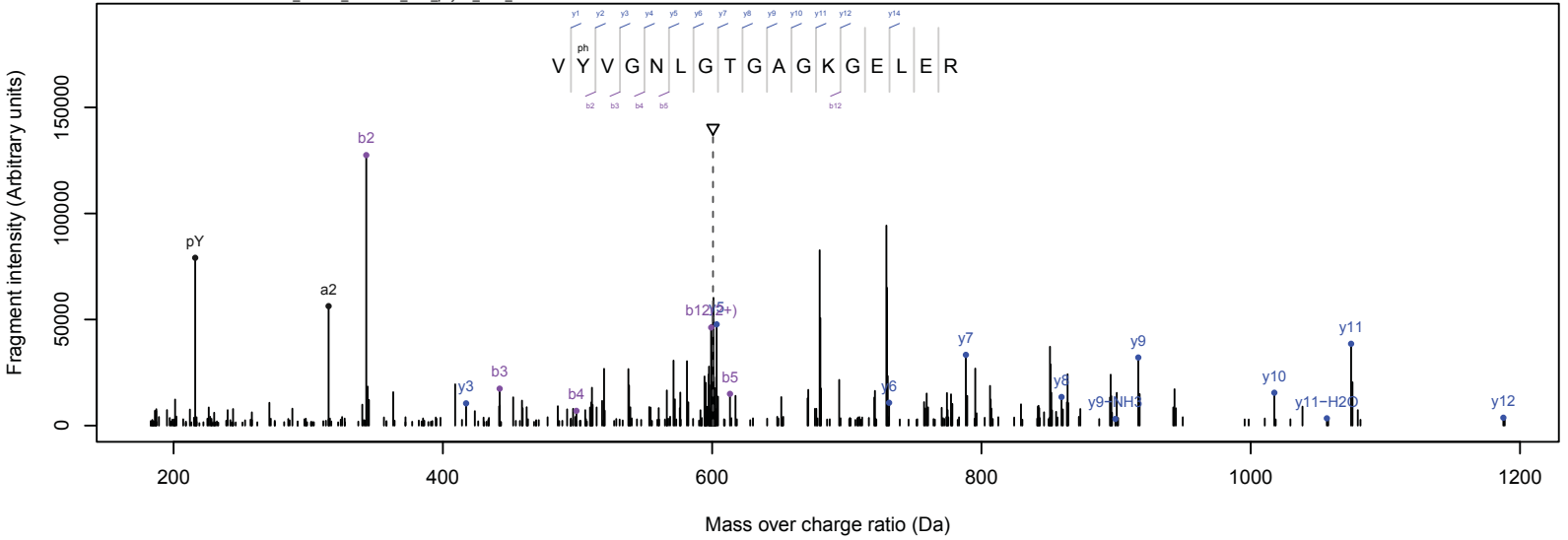
Andromeda score : 111.26



**VY(ph)VGNLGTGAGKGELEER-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 21499

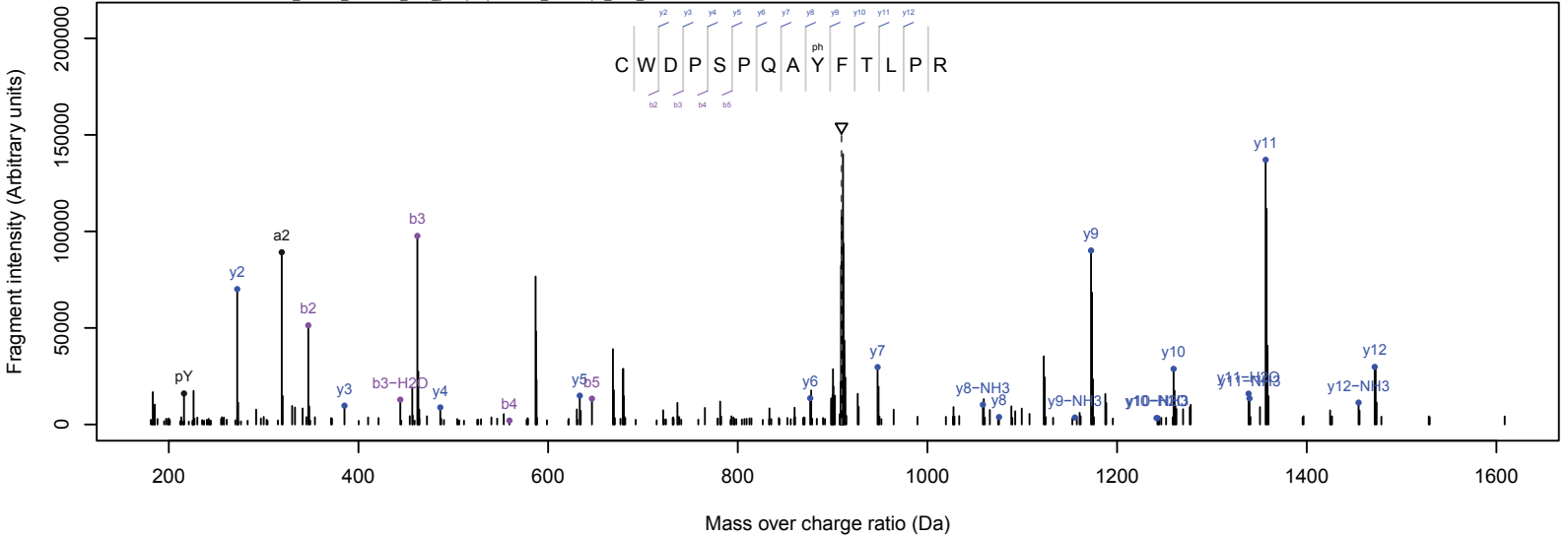
Andromeda score : 111.22



**CWDPSQAY(ph)FTLPR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40318

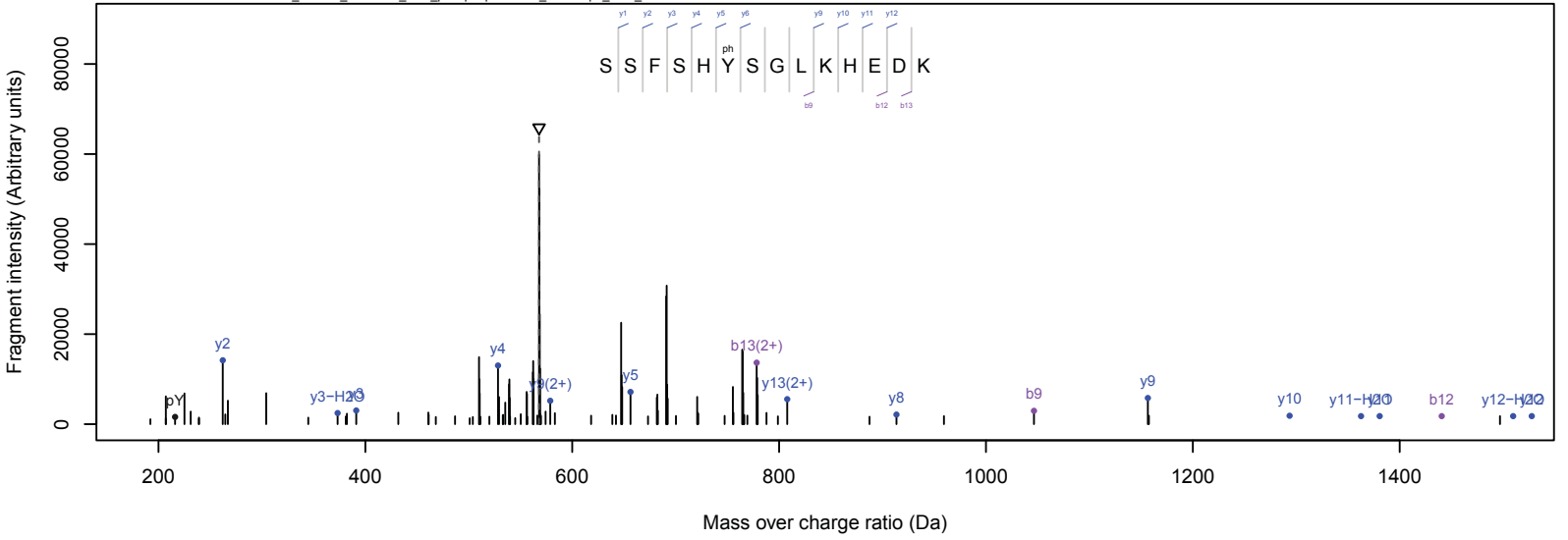
Andromeda score : 111.12



**SSFSHY(ph)SGLKHEDK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6010

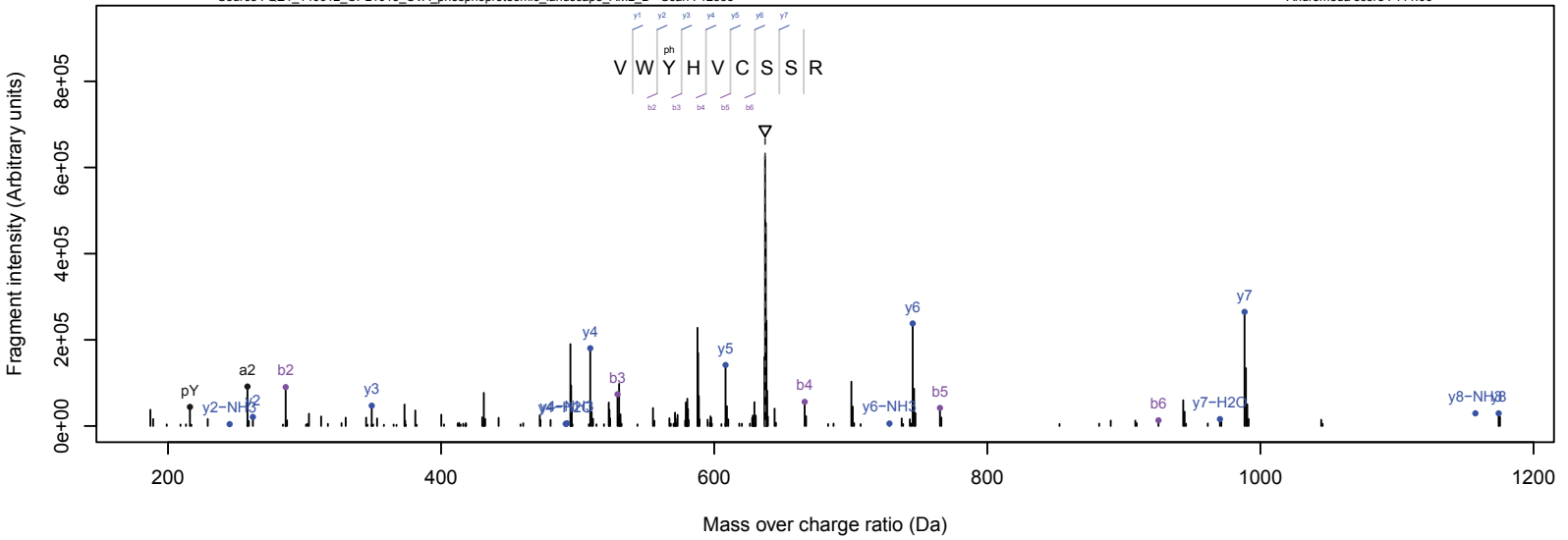
Andromeda score : 111.12



**VWY(ph)HVCSSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12335

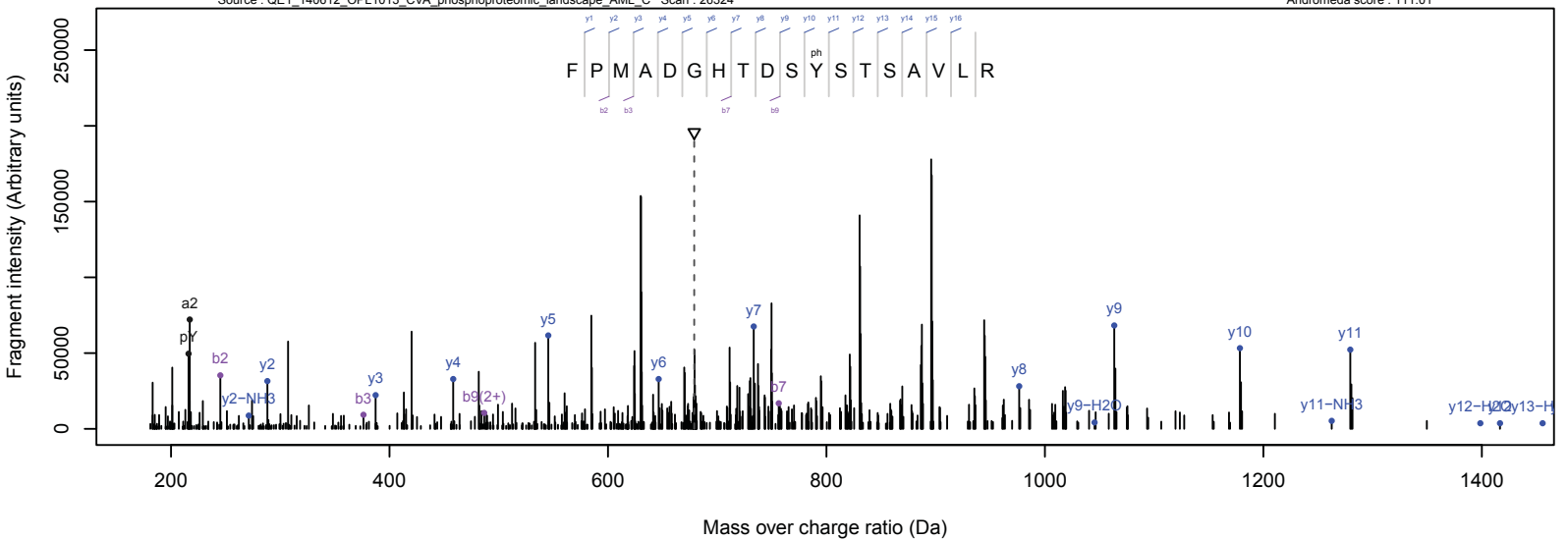
Andromeda score : 111.06



**FPMADGHTDSY(ph)STSAVLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26324

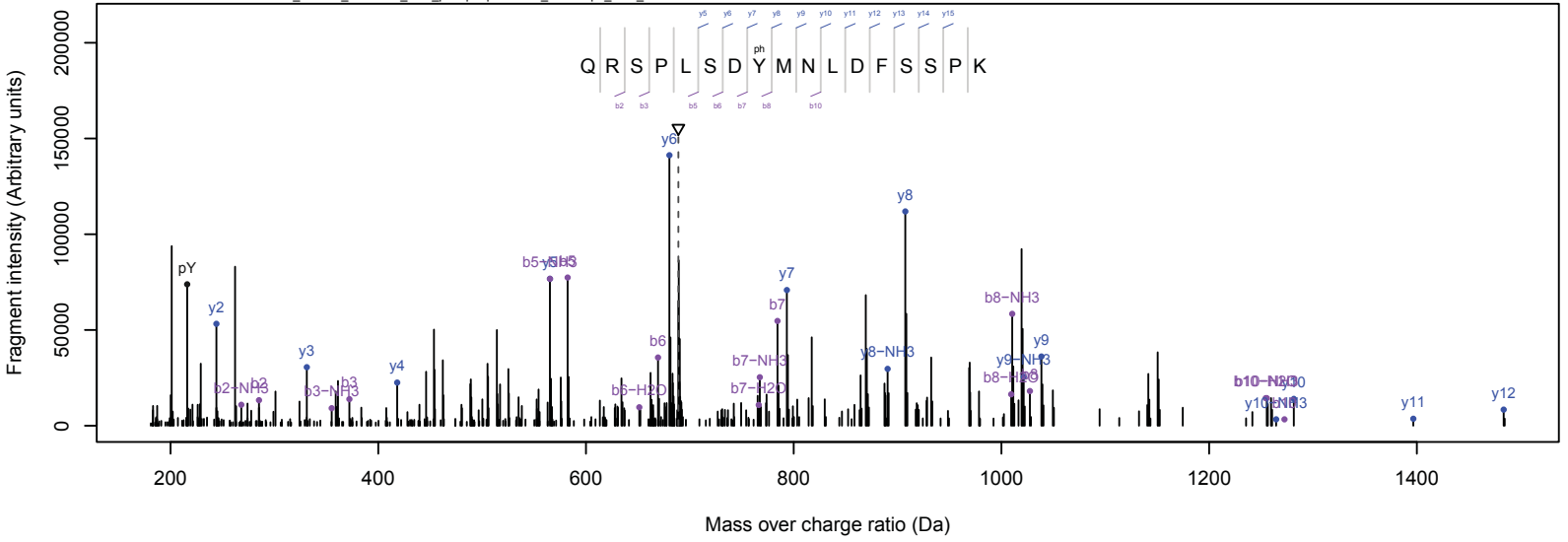
Andromeda score : 111.01



QRSPLSDY(ph)MNLDFSSPK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36507

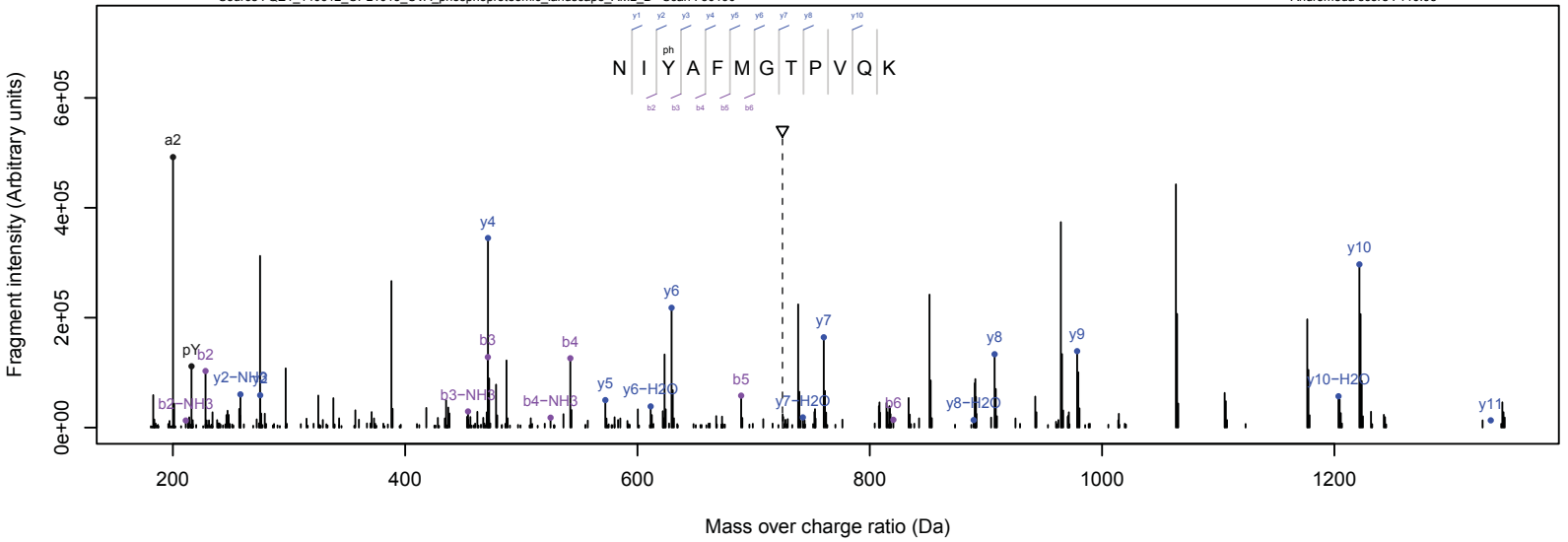
Andromeda score: 111.01



NIY(ph)AFMGTPVQK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 30190

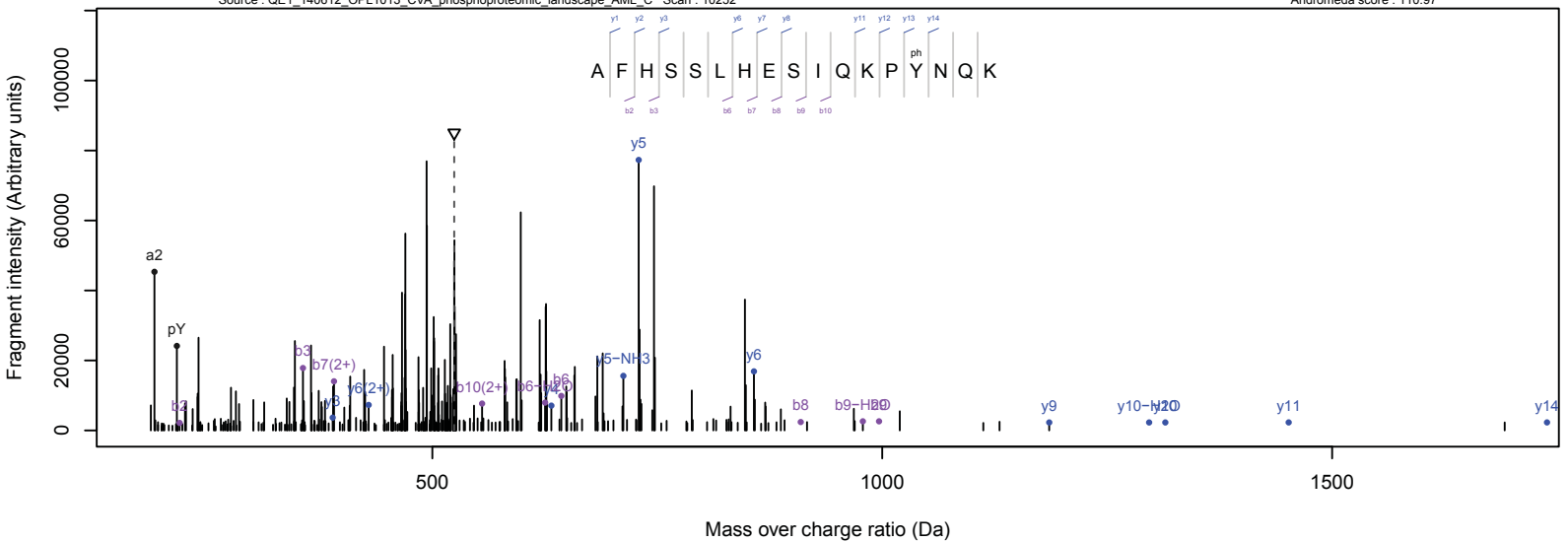
Andromeda score: 110.98

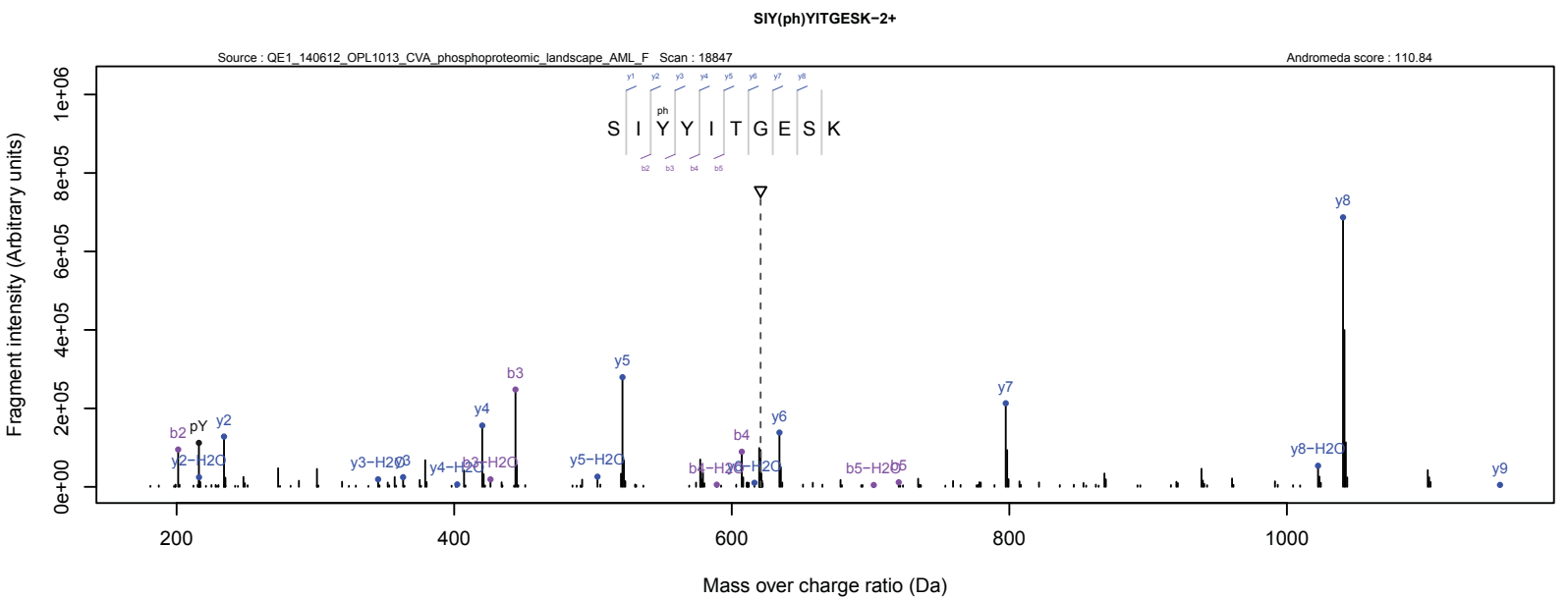
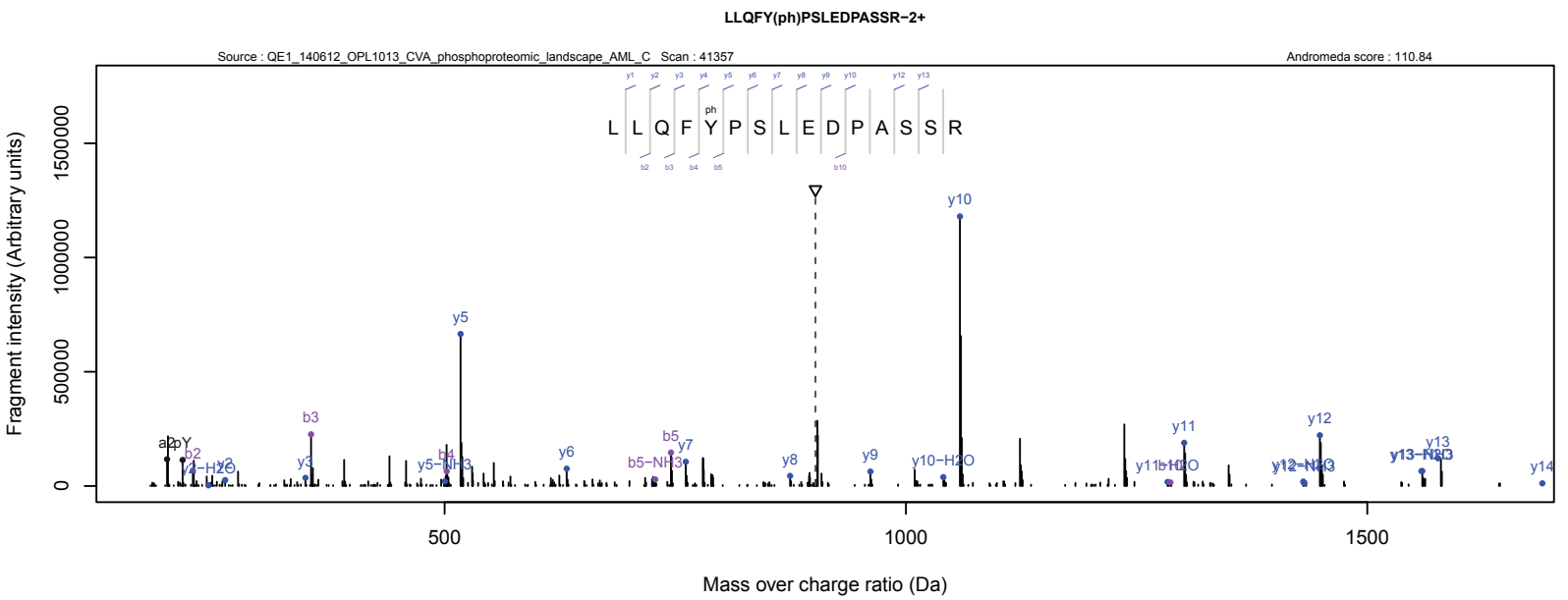
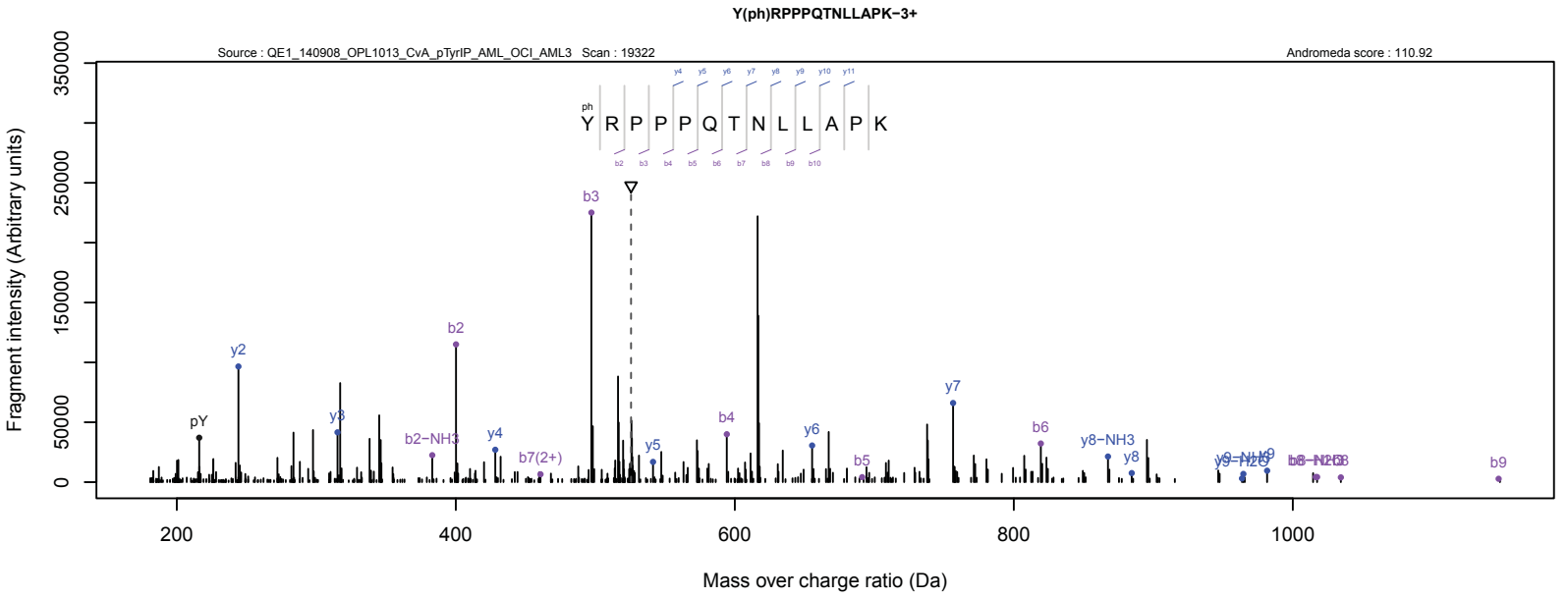


AFHSSLHESI QKPY NQK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10252

Andromeda score: 110.97

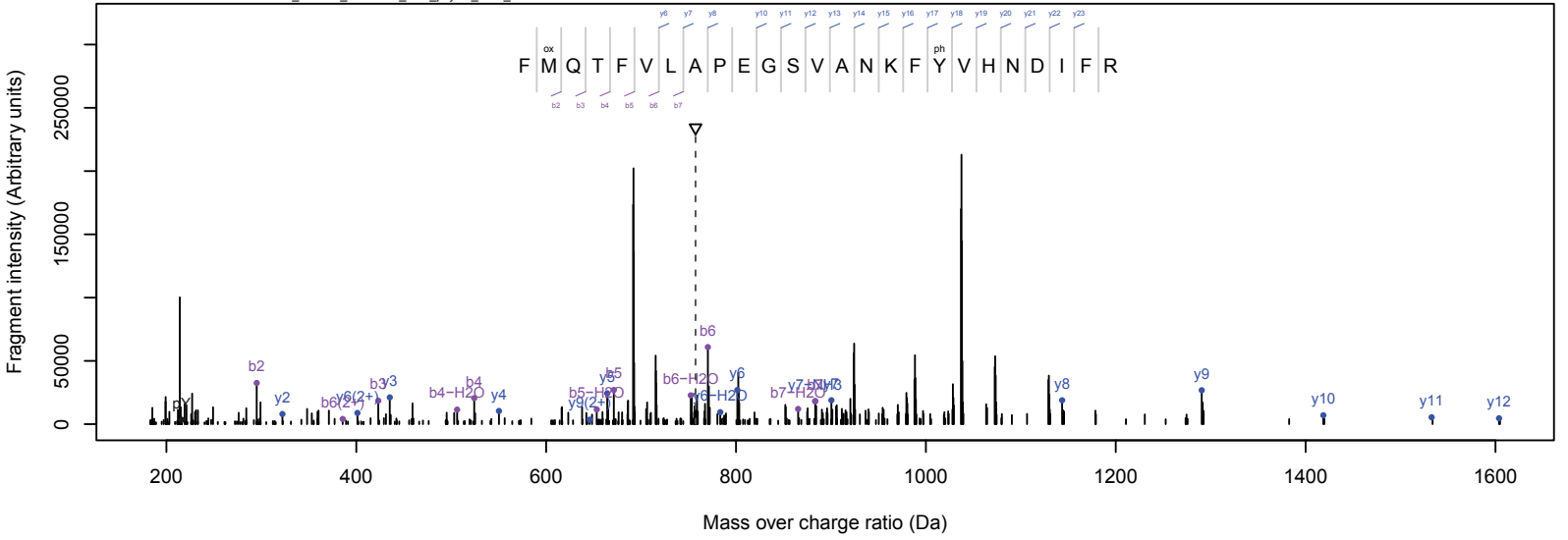




FM(ox)QTFVLAPEGSVANKFY(ph)VHNDIFR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 46377

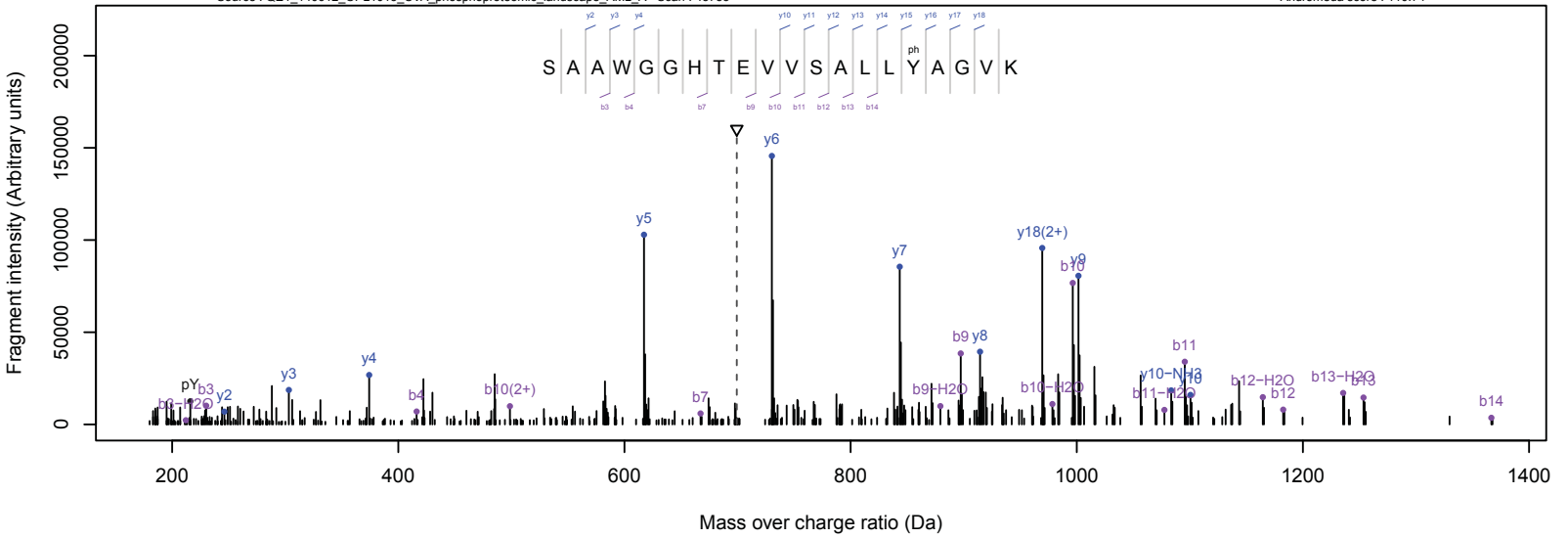
Andromeda score : 110.83



SAAWGGHTEVVSALLYAGVK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_A Scan : 43755

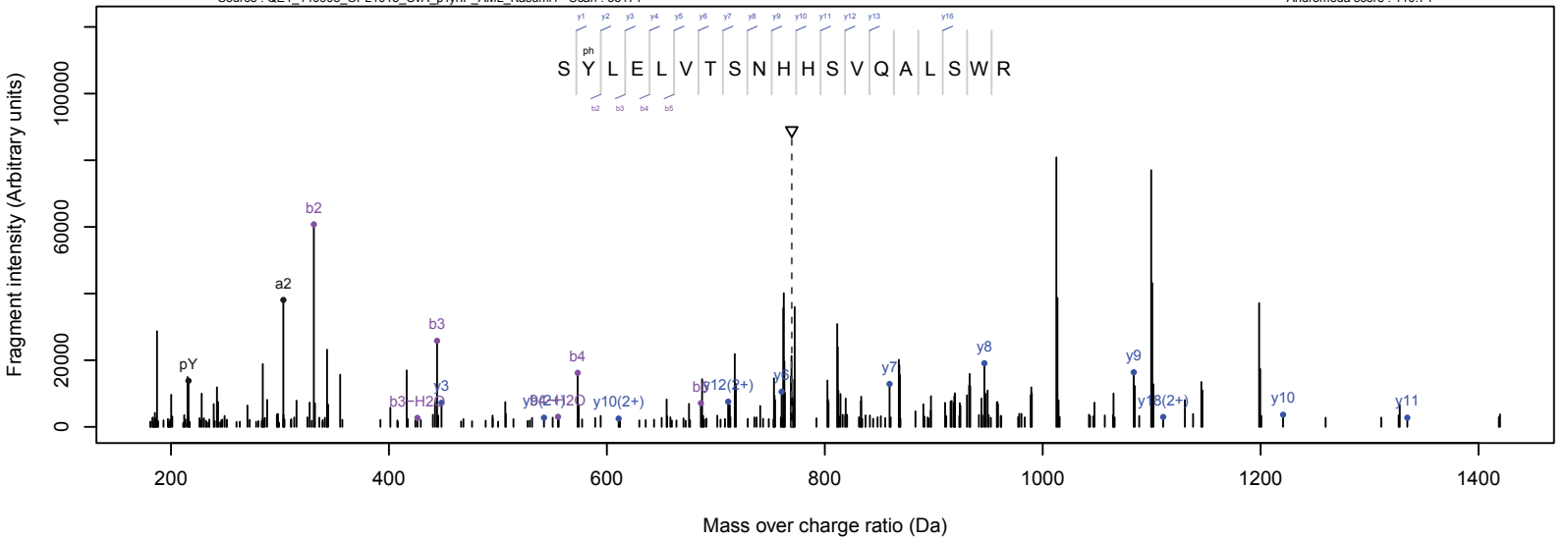
Andromeda score : 110.74



SY(ph)LELVTSNHHSVQALSWR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 33171

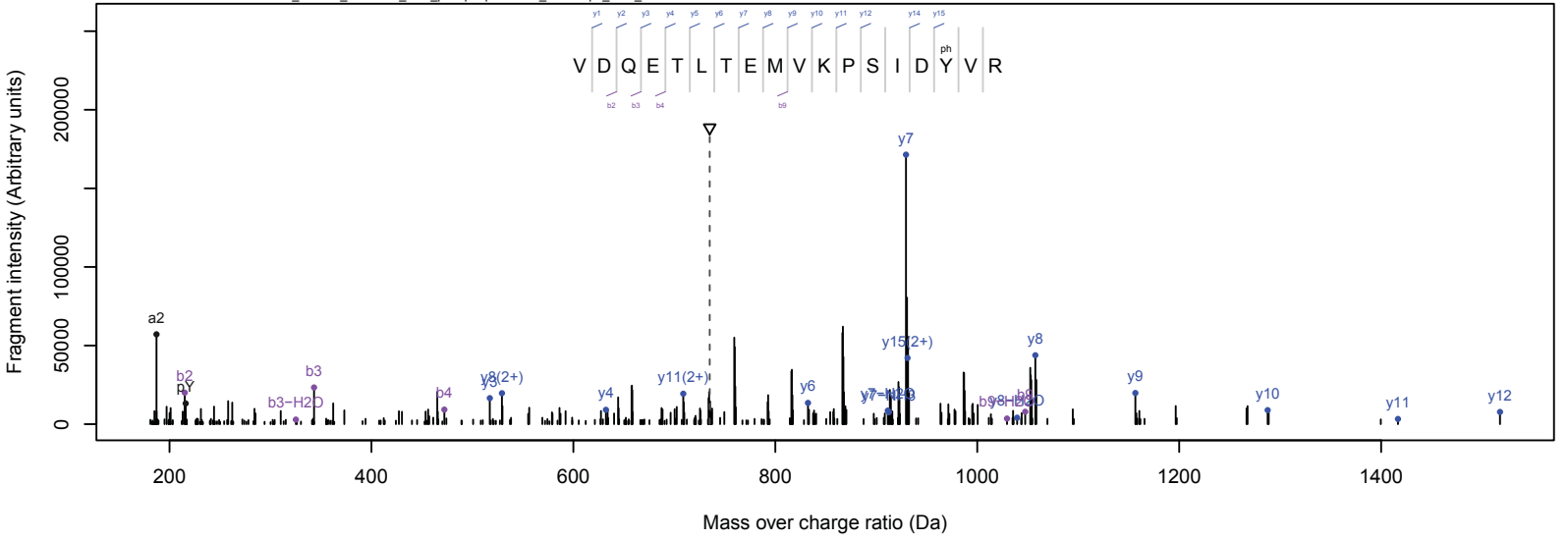
Andromeda score : 110.74



**VDQETLTEMVKPSIDYVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36503

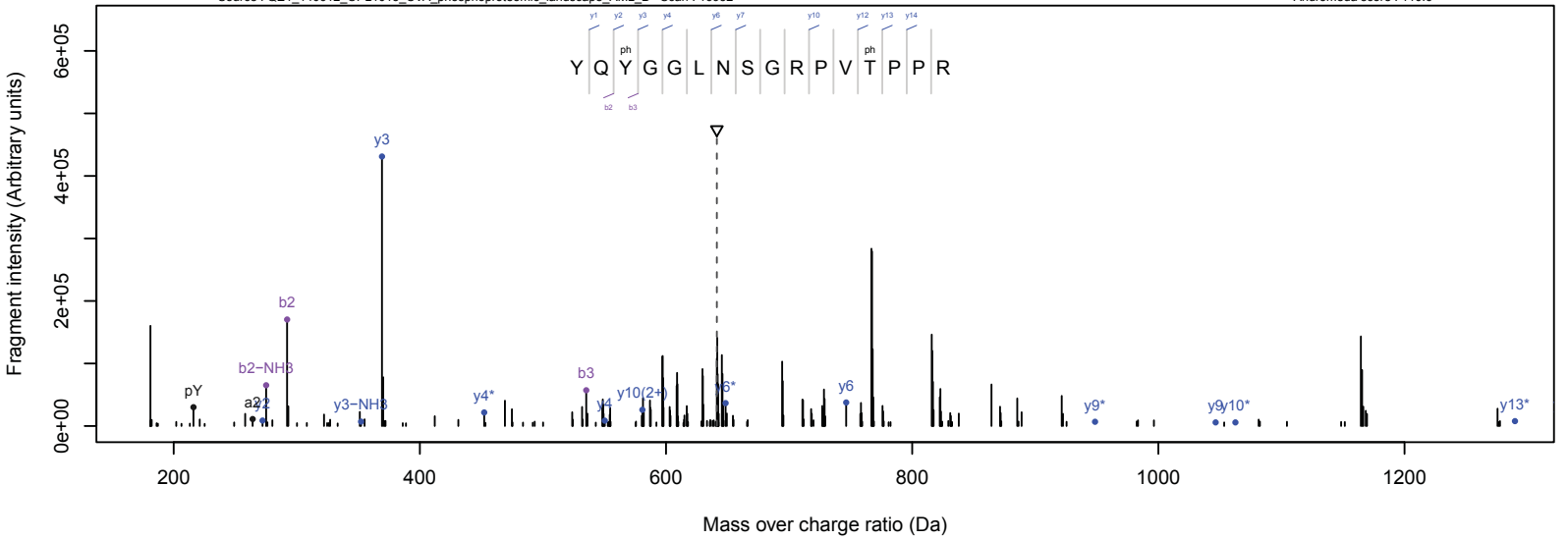
Andromeda score : 110.74



**YQY(ph)GGLNSGRPVT(ph)PPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16952

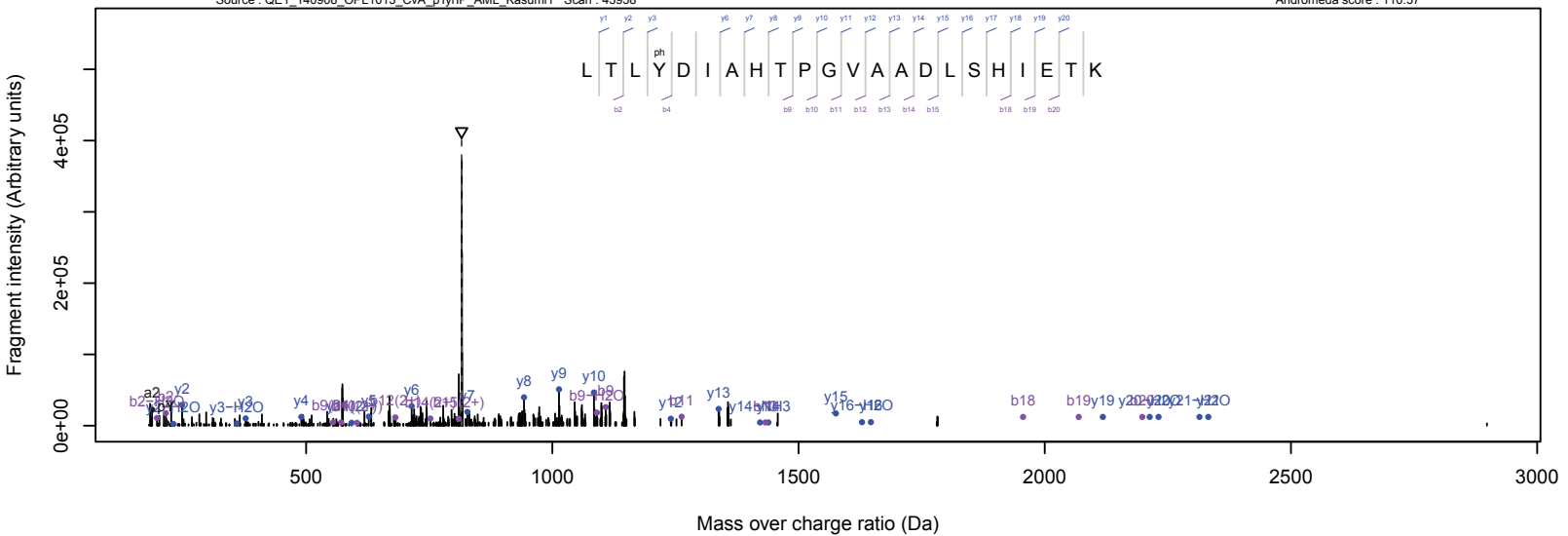
Andromeda score : 110.6



**LTLY(ph)DIAHTPGVAADLSHIETK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 43938

Andromeda score : 110.57

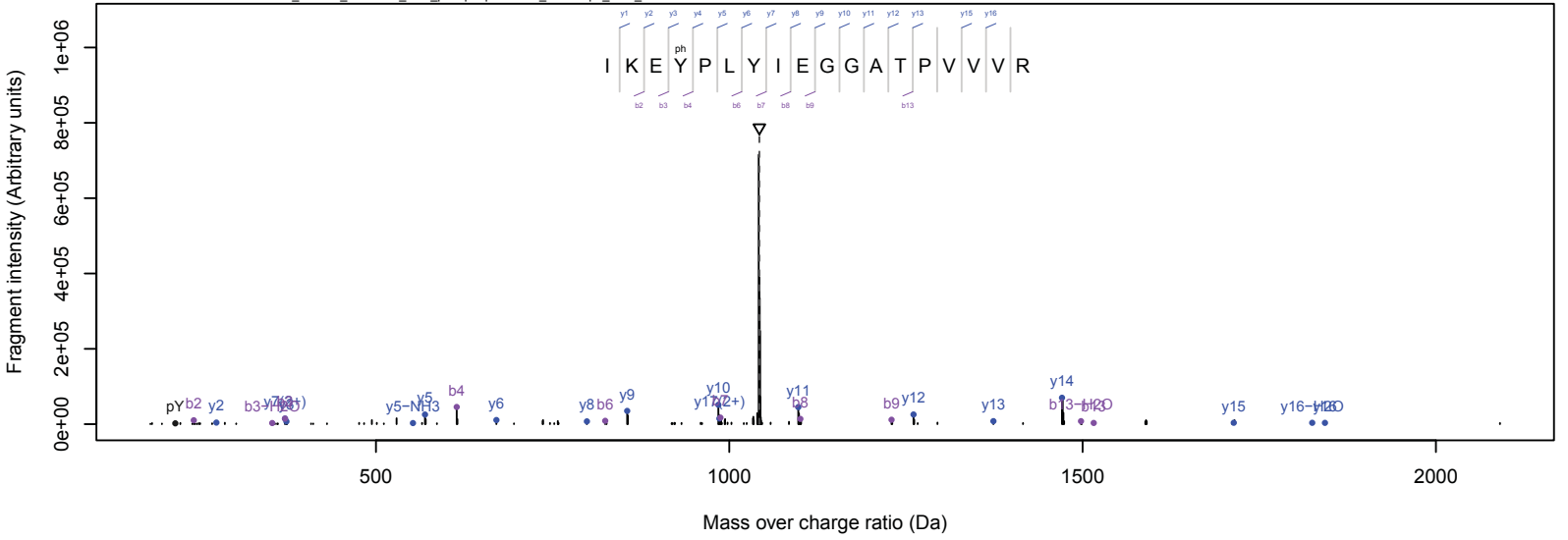




IKEY(ph)PLYIEGGATPVVVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34355

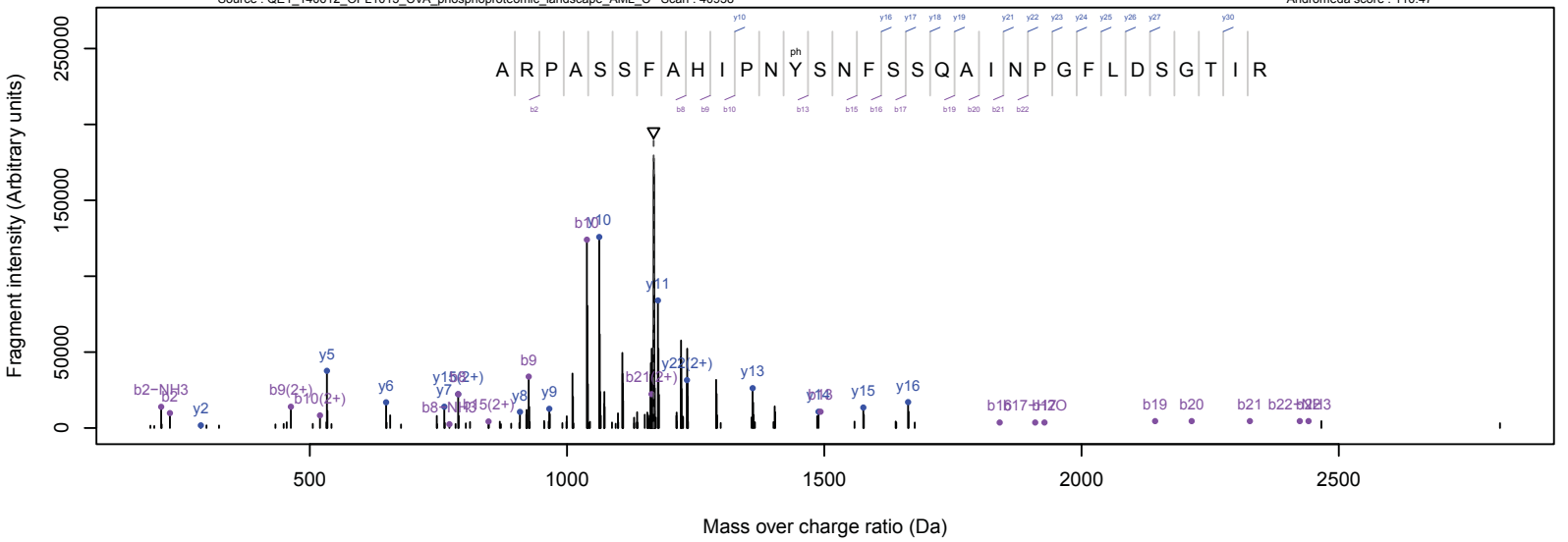
Andromeda score: 110.56



ARPASSFAHIPNY(ph)SNFSSQAINPGFLDSGTIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 40958

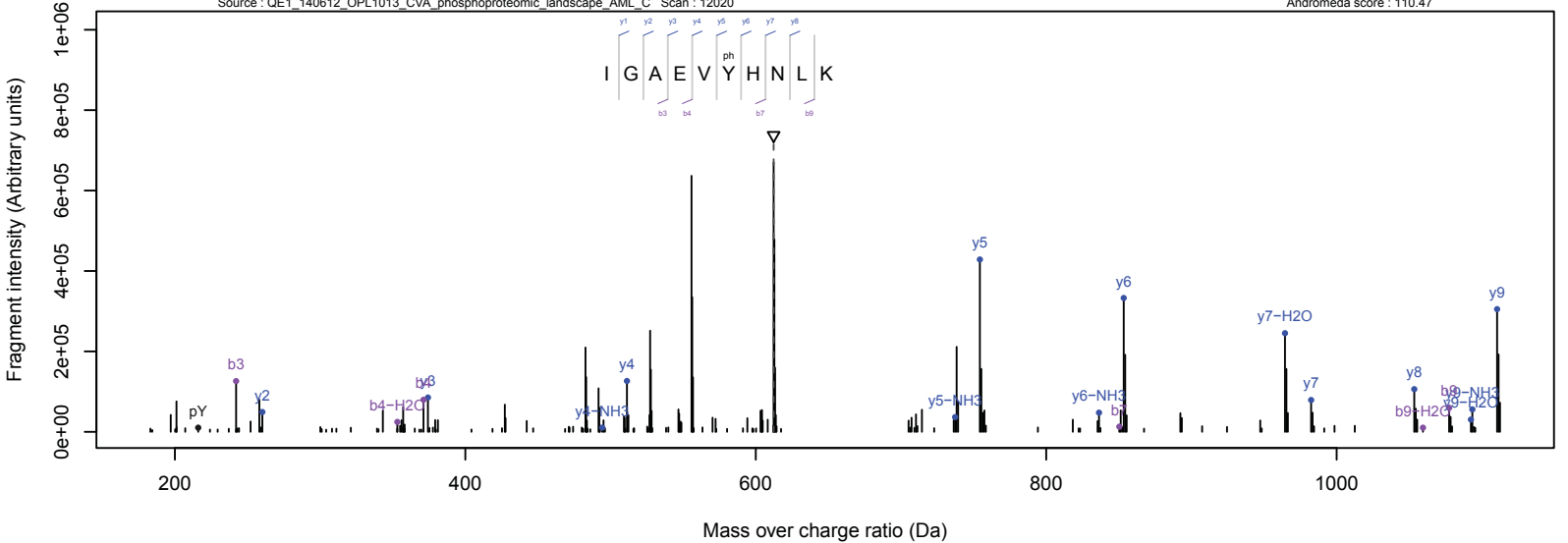
Andromeda score: 110.47



IGAEVY(ph)HNLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12020

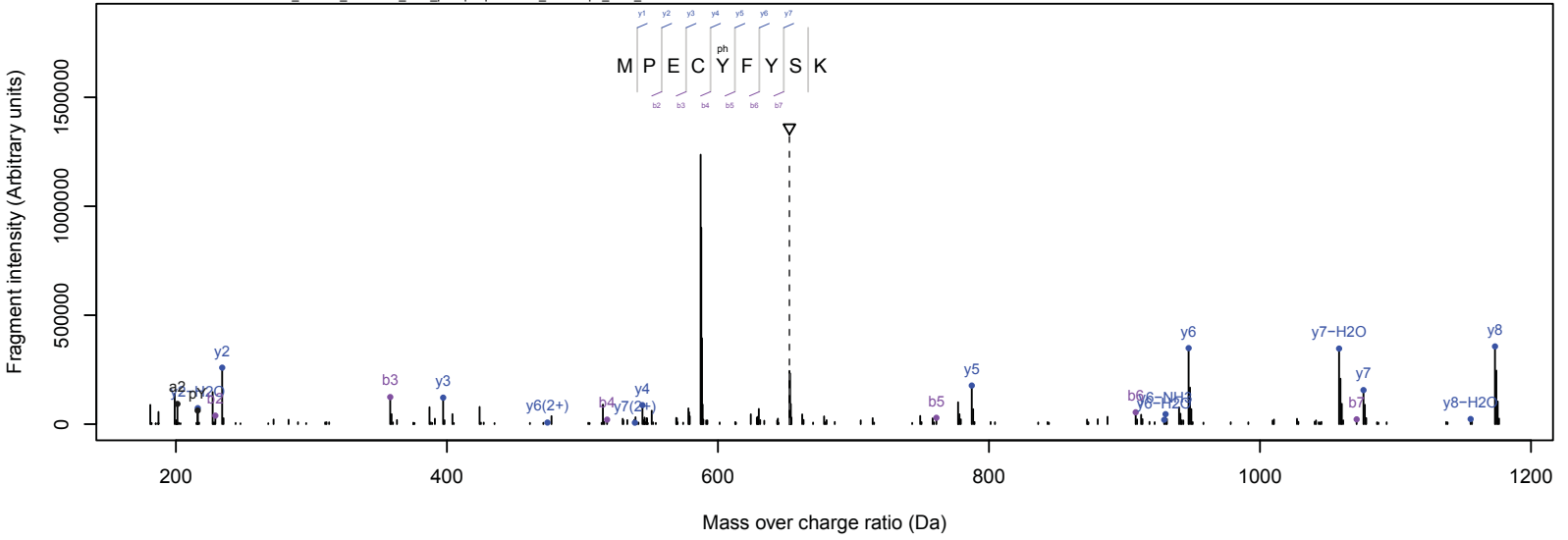
Andromeda score: 110.47



MPECY(ph)FYSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 20210

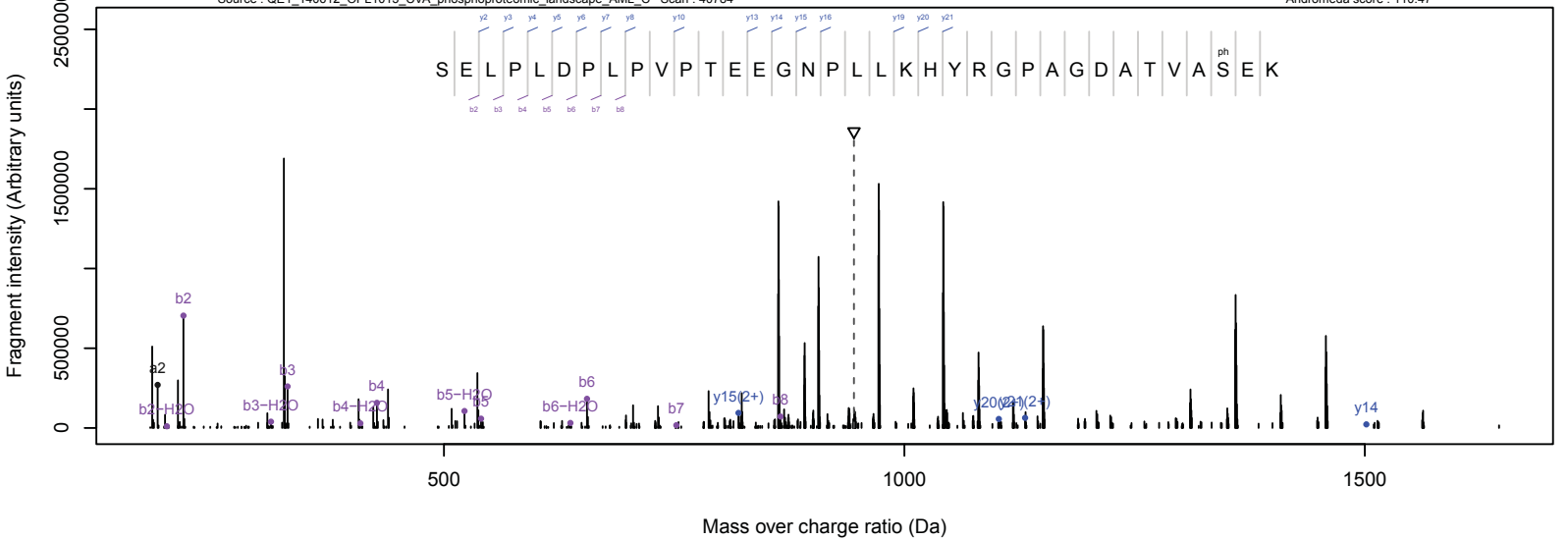
Andromeda score : 110.47



SELPLDPLPVPTEEGNPLLKHRYRGPAGDATVAS(ph)EK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40784

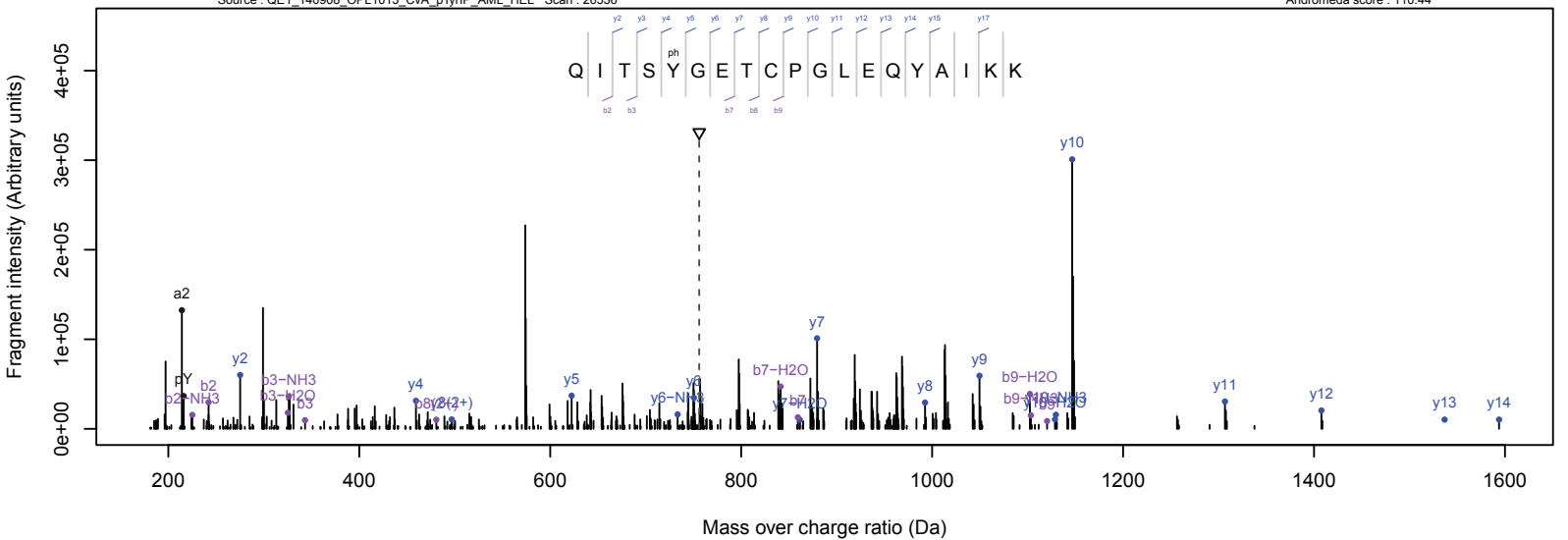
Andromeda score : 110.47



QITSY(ph)GETCPGLEQYAIKK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 26536

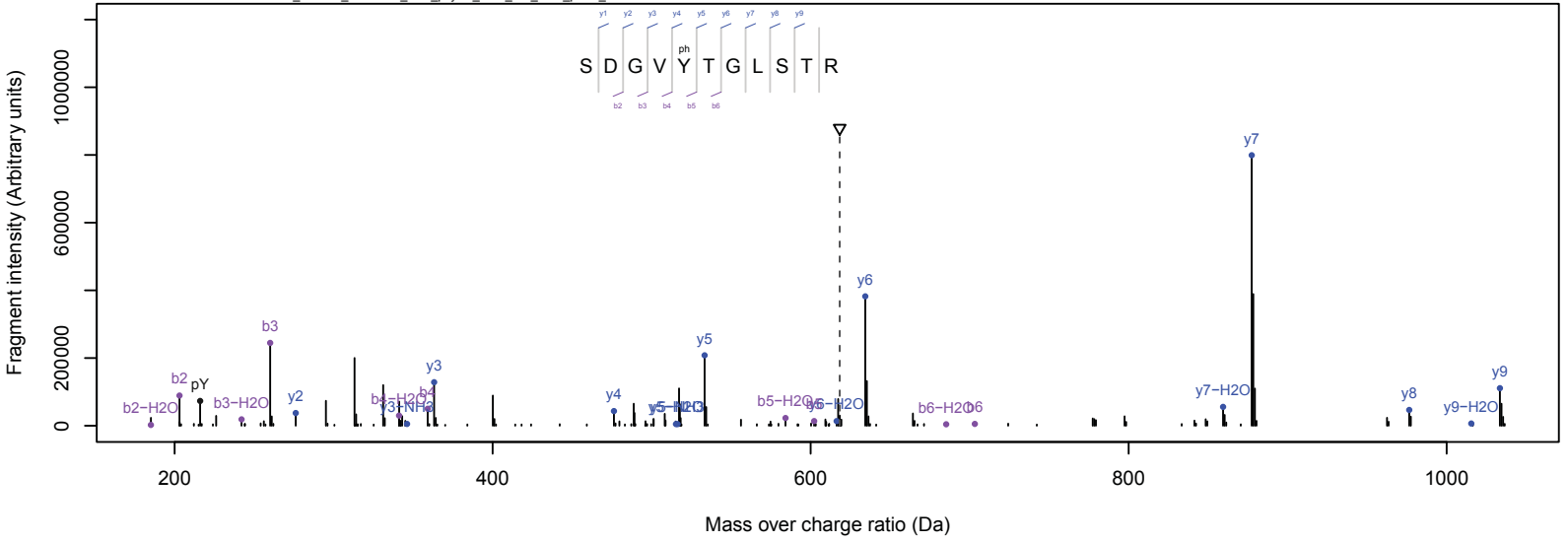
Andromeda score : 110.44



SDGVY(ph)TGLSTR-2+

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 8491

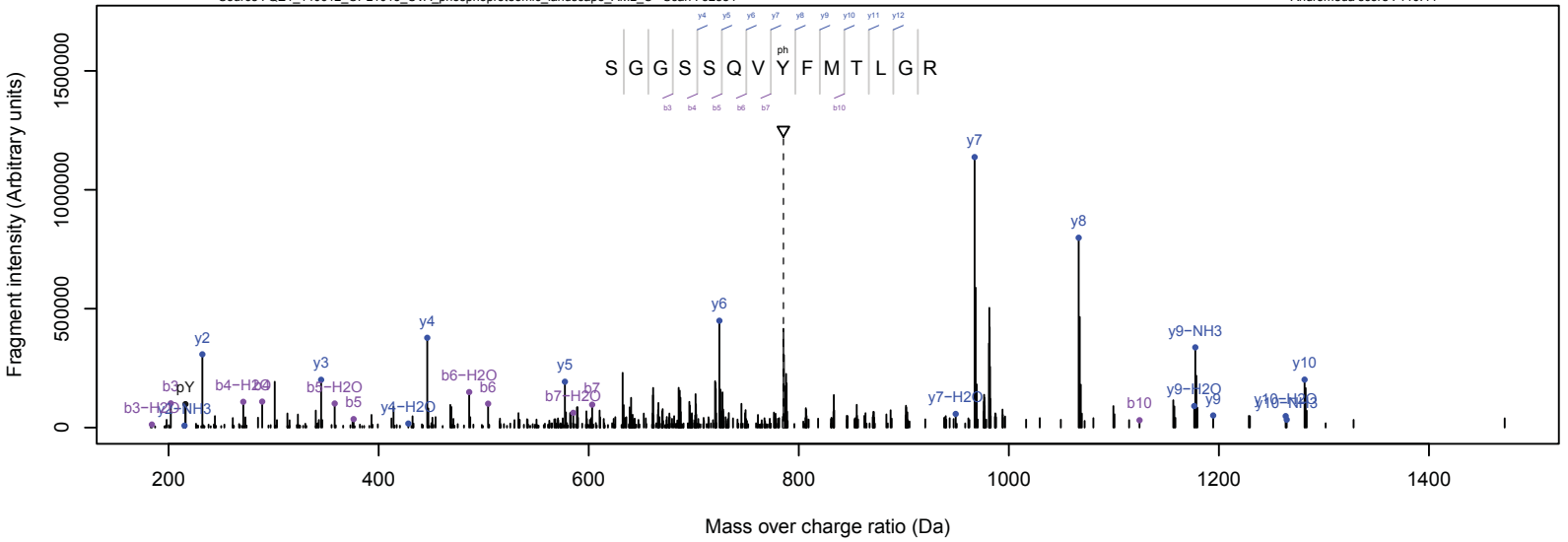
Andromeda score: 110.44



SGGSSQVY(ph)FMLTGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 32884

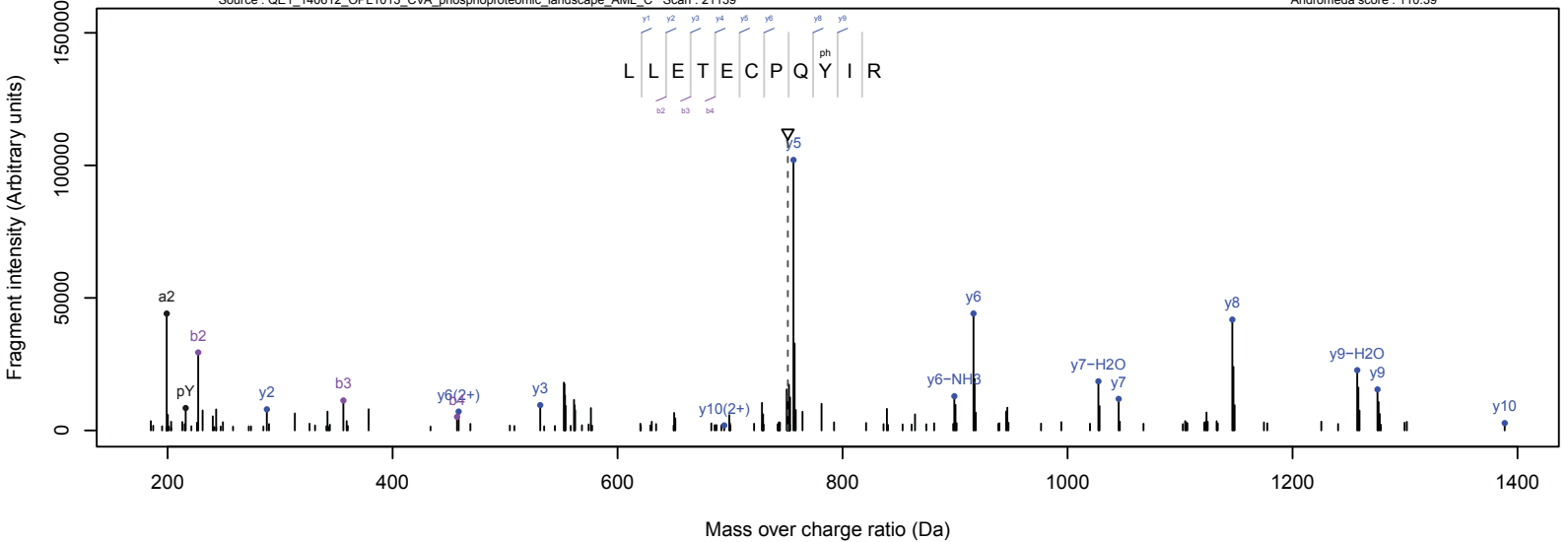
Andromeda score: 110.44



LLETCPQY(ph)IR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21139

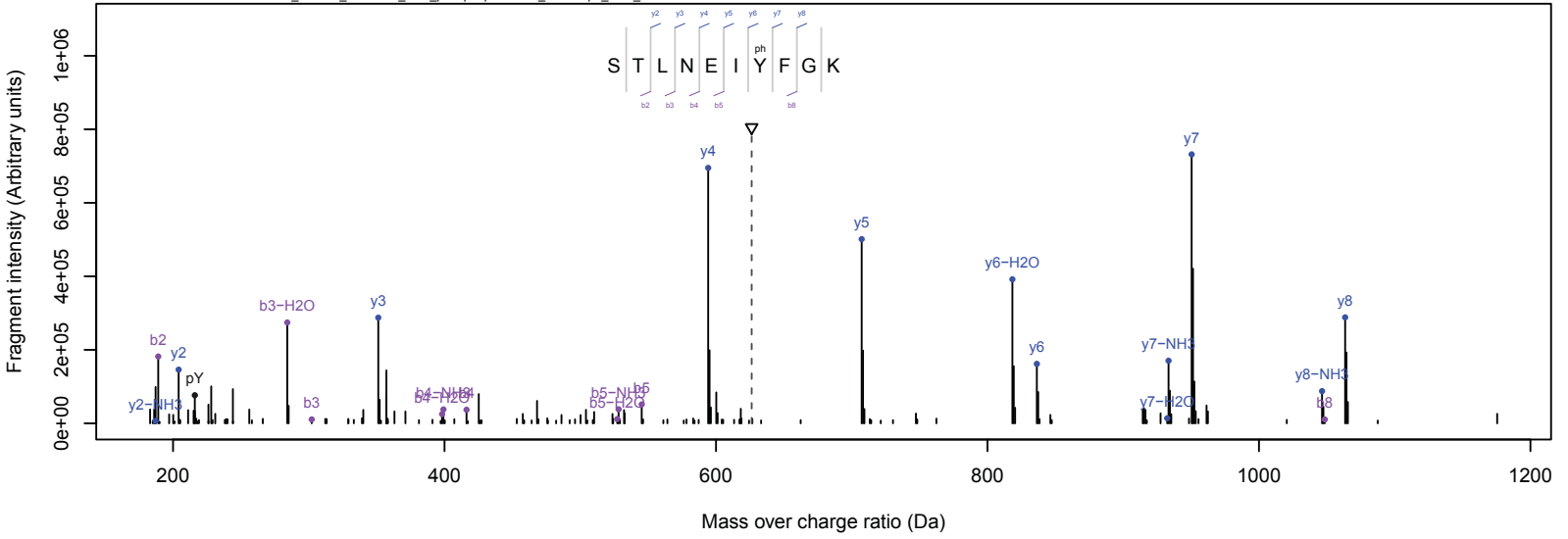
Andromeda score: 110.39



STLNEIY(ph)FGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27570

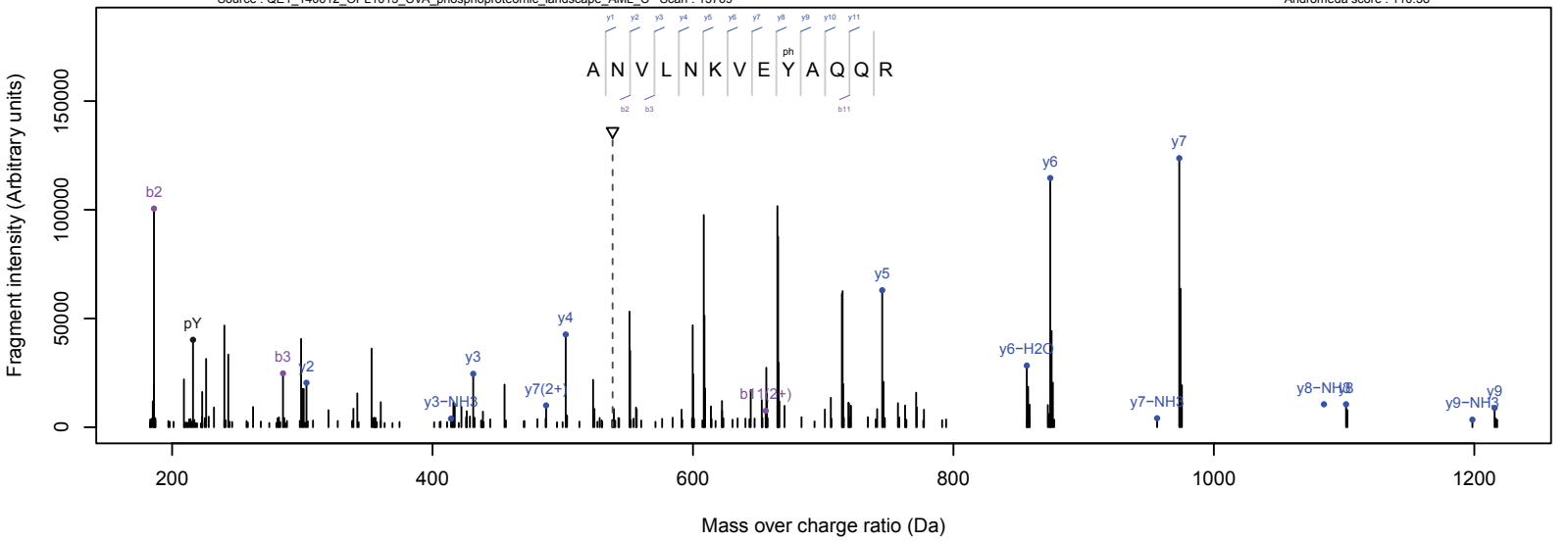
Andromeda score : 110.39



ANVLNKVEY(ph)AQQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15769

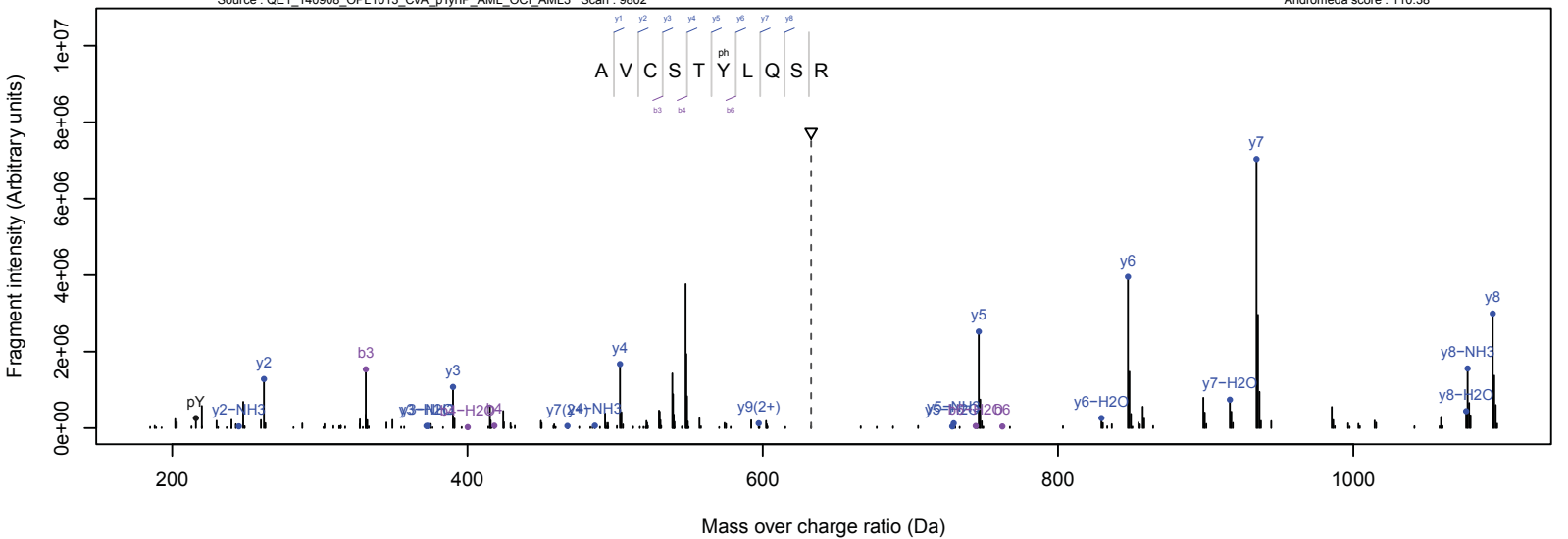
Andromeda score : 110.38



AVCSTY(ph)LQSR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 9802

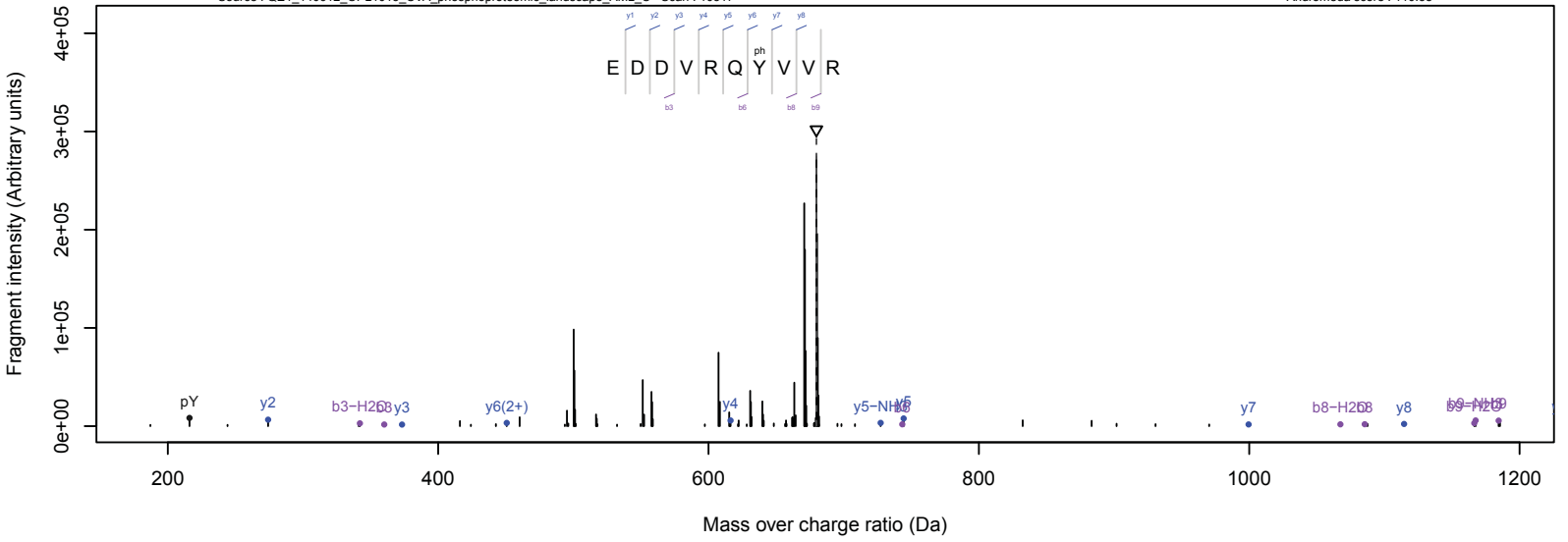
Andromeda score : 110.38



EDDVRQ(ph)VVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10517

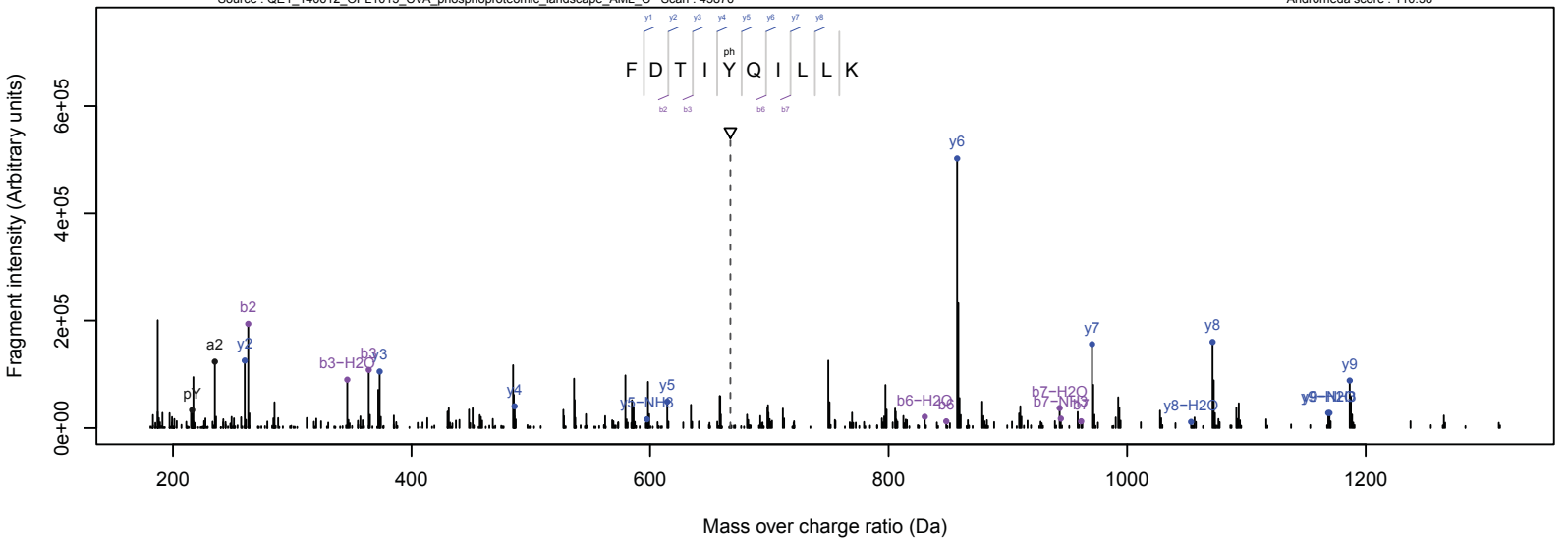
Andromeda score: 110.38



FDTIY(ph)QILLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 45876

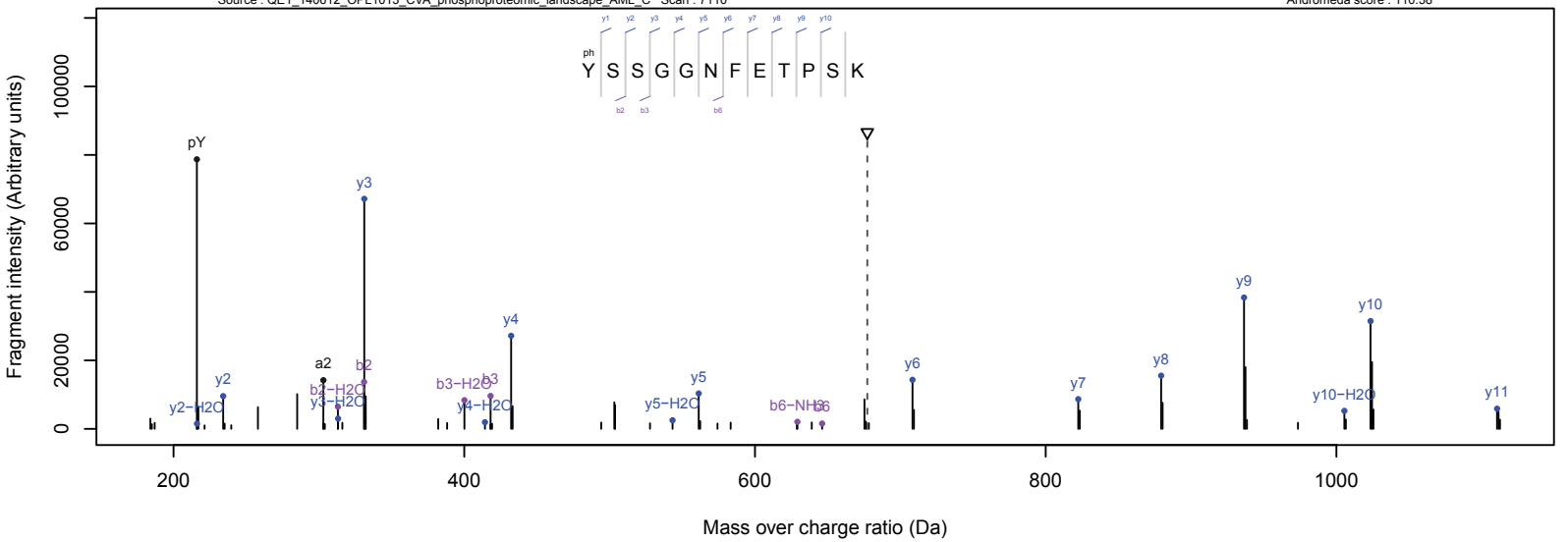
Andromeda score: 110.38



Y(ph)SSGGNFETPSK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 7110

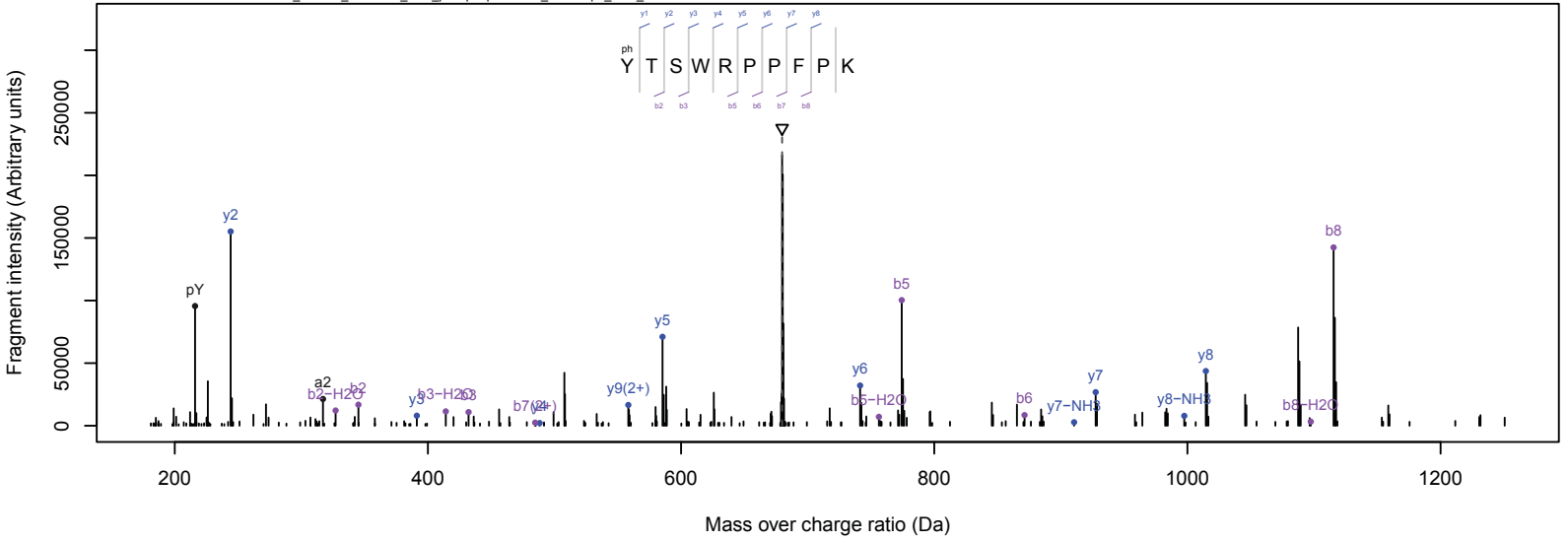
Andromeda score: 110.38



Y(ph)TSWRPPFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27148

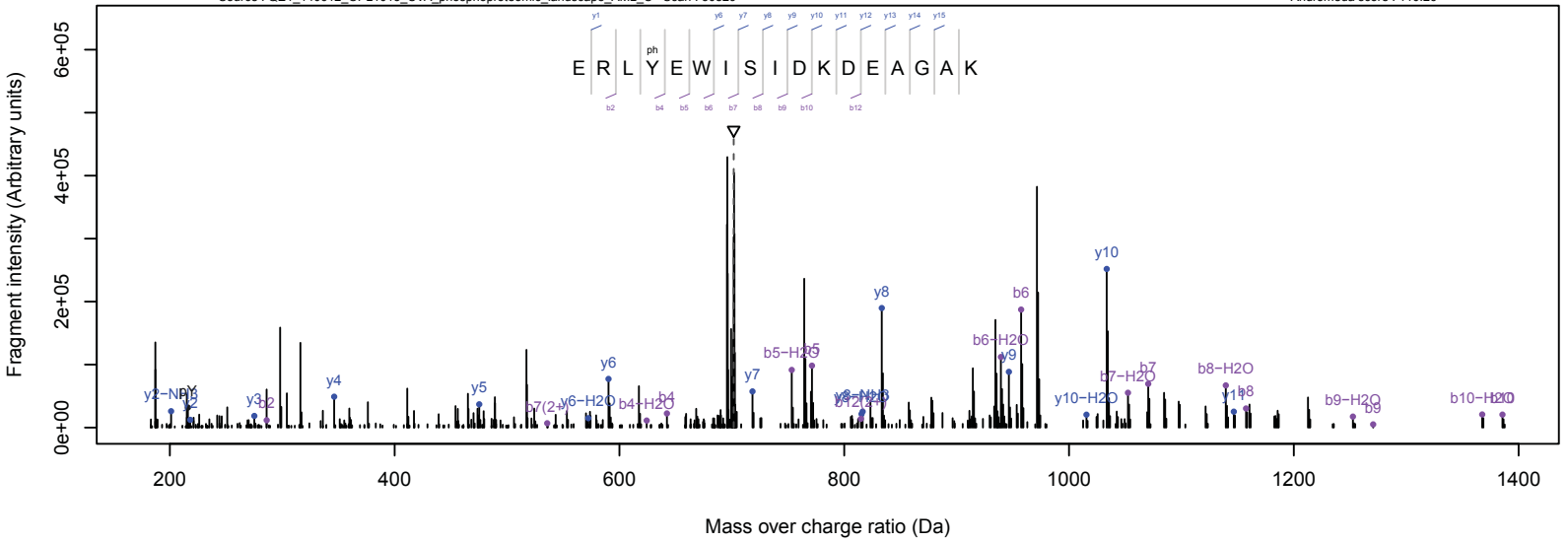
Andromeda score : 110.38



ERLY(ph)EWISIDKDEAGAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30829

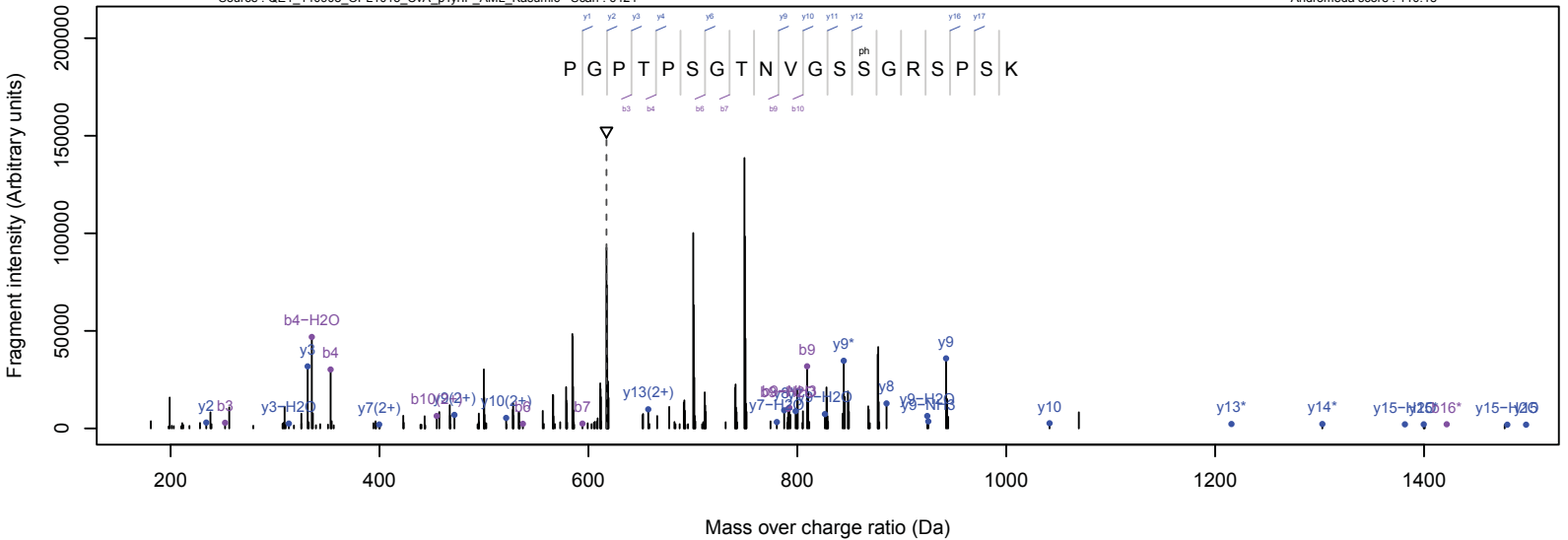
Andromeda score : 110.26



PGPTPSGTNVGSS(ph)GRSPSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 6424

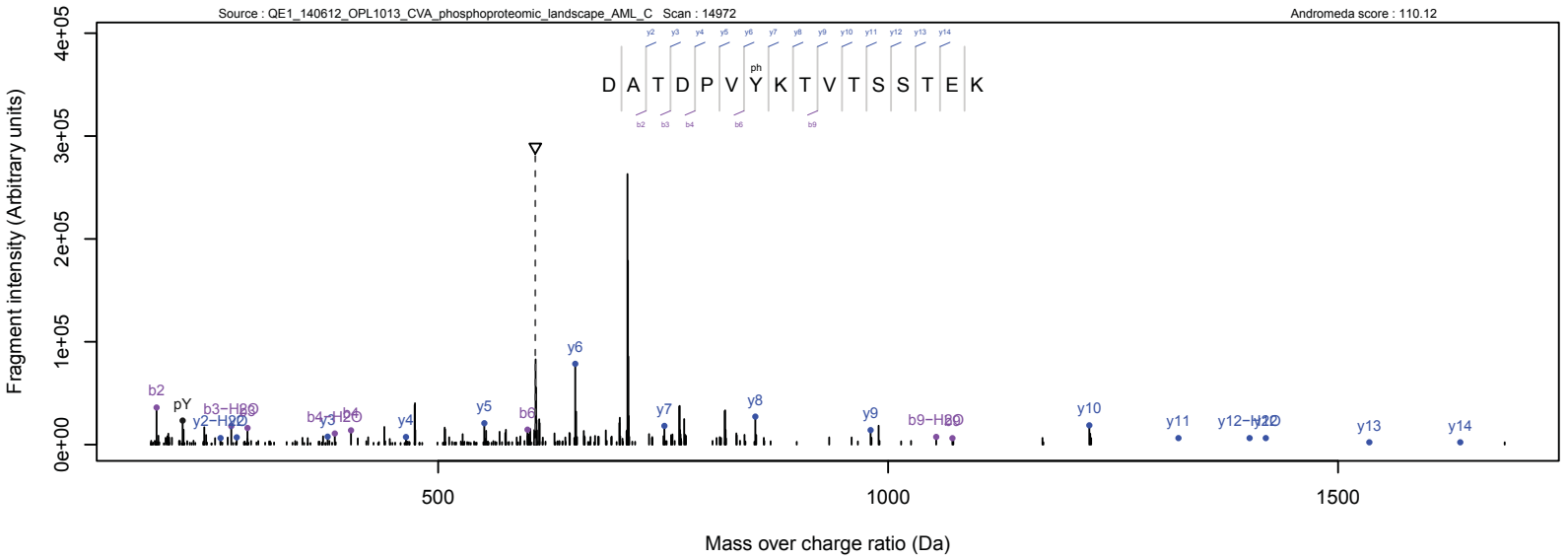
Andromeda score : 110.18



**DATDPVY(ph)KTVTSSTEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14972

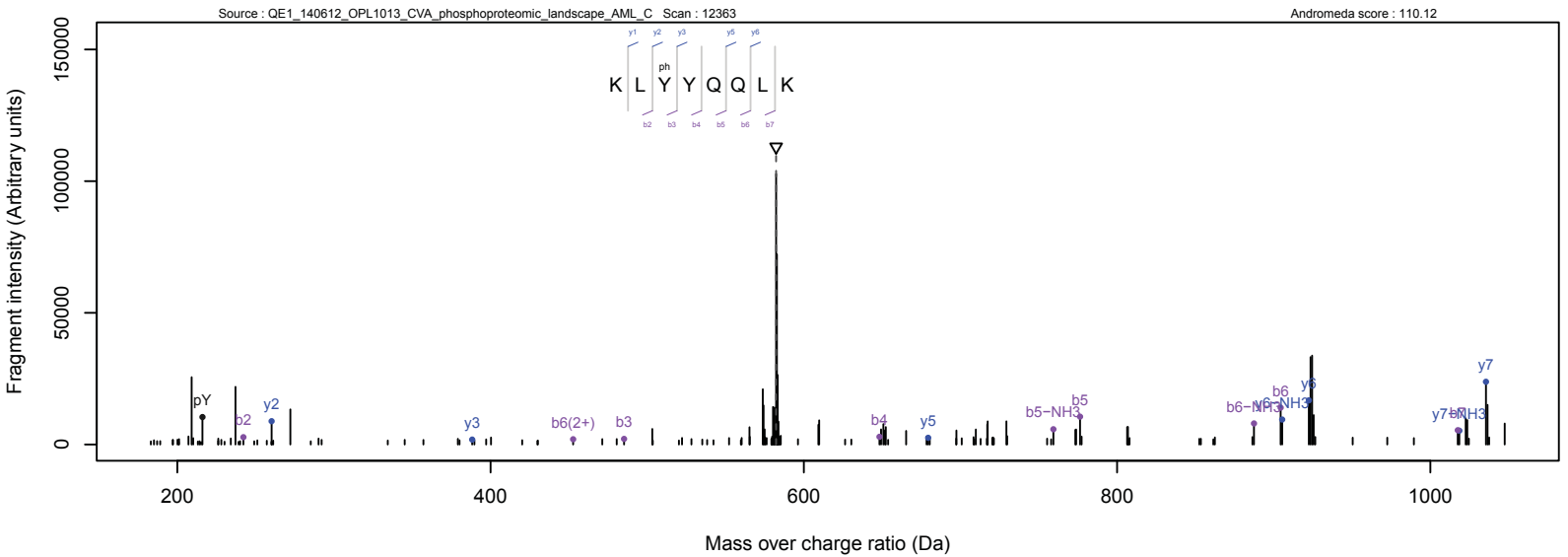
Andromeda score : 110.12



**KLY(ph)YQQLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12363

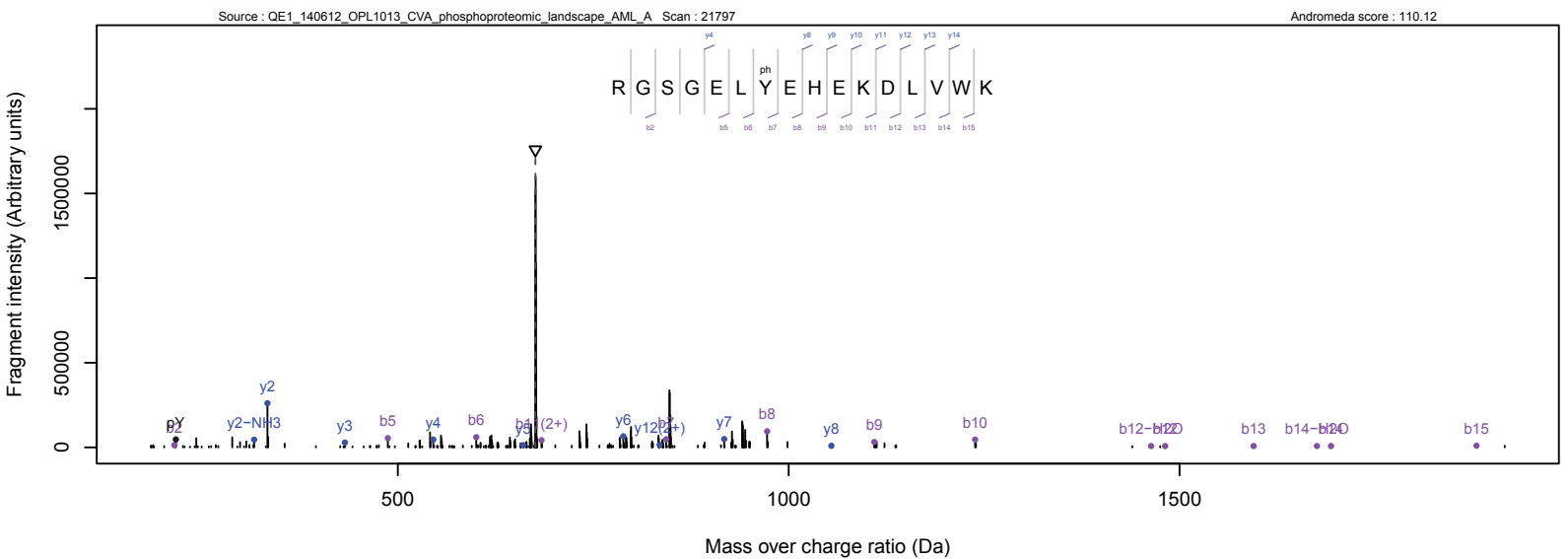
Andromeda score : 110.12



**RGSGELY(ph)EHEKDLVWK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 21797

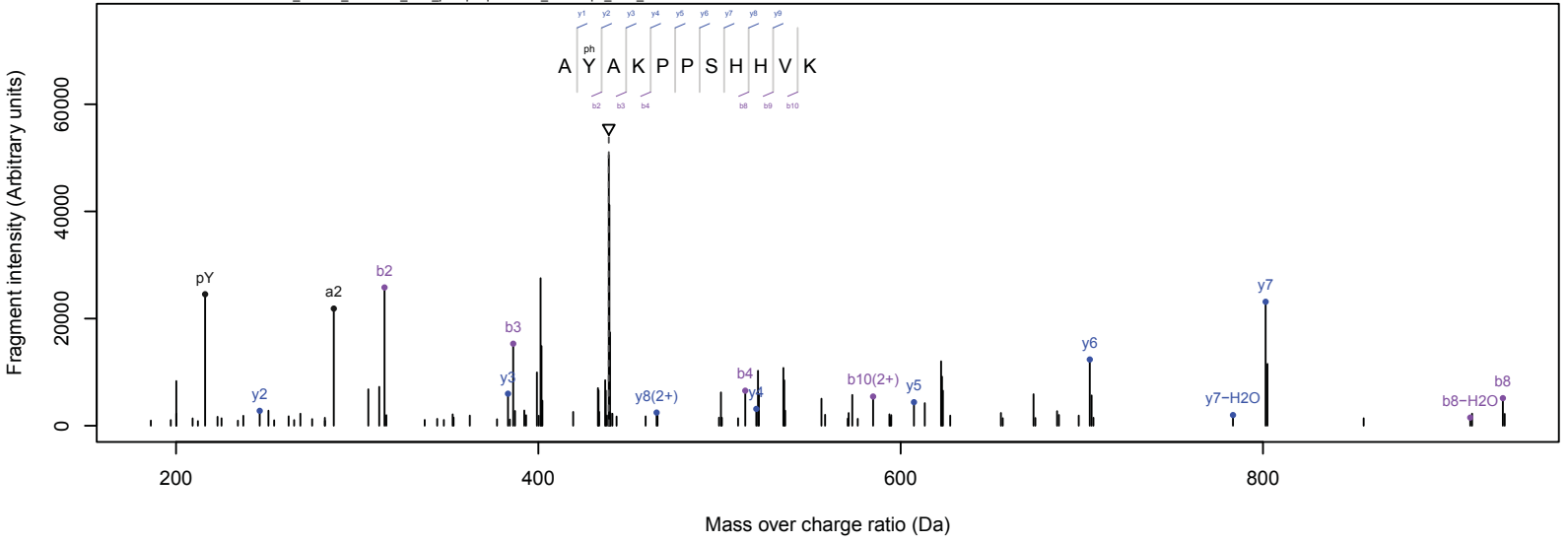
Andromeda score : 110.12



AY(ph)AKPPSHHVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 3388

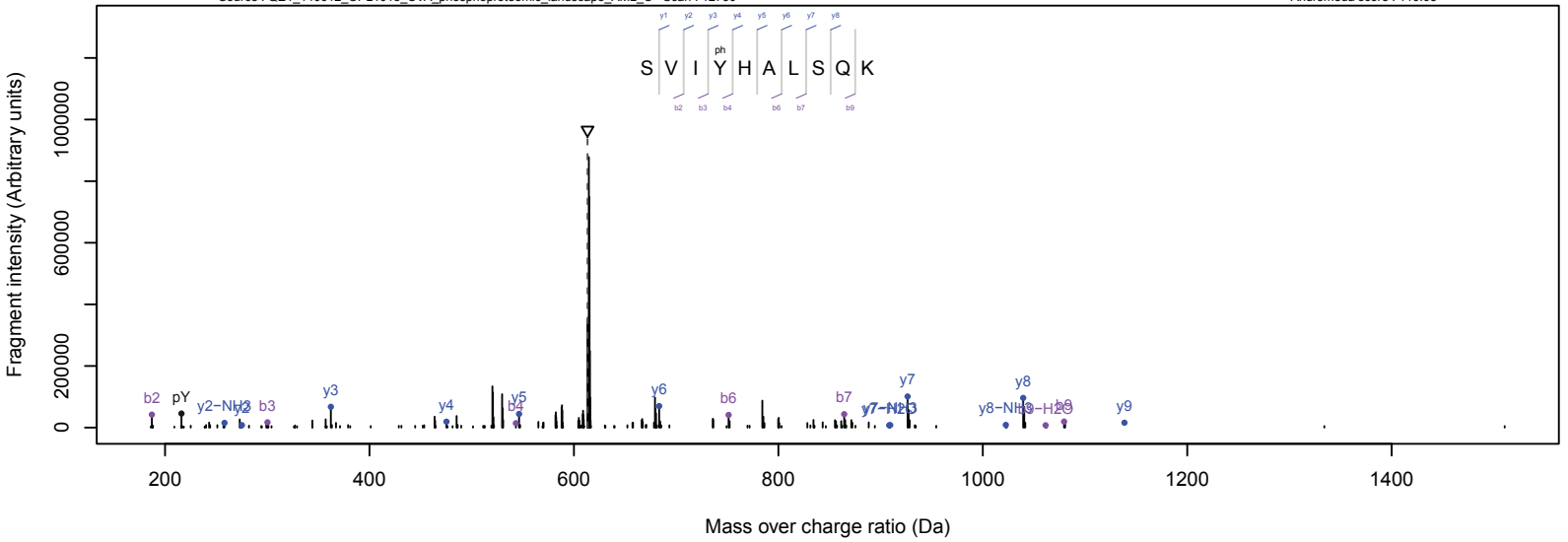
Andromeda score : 110.08



SVIY(ph)HALSQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12750

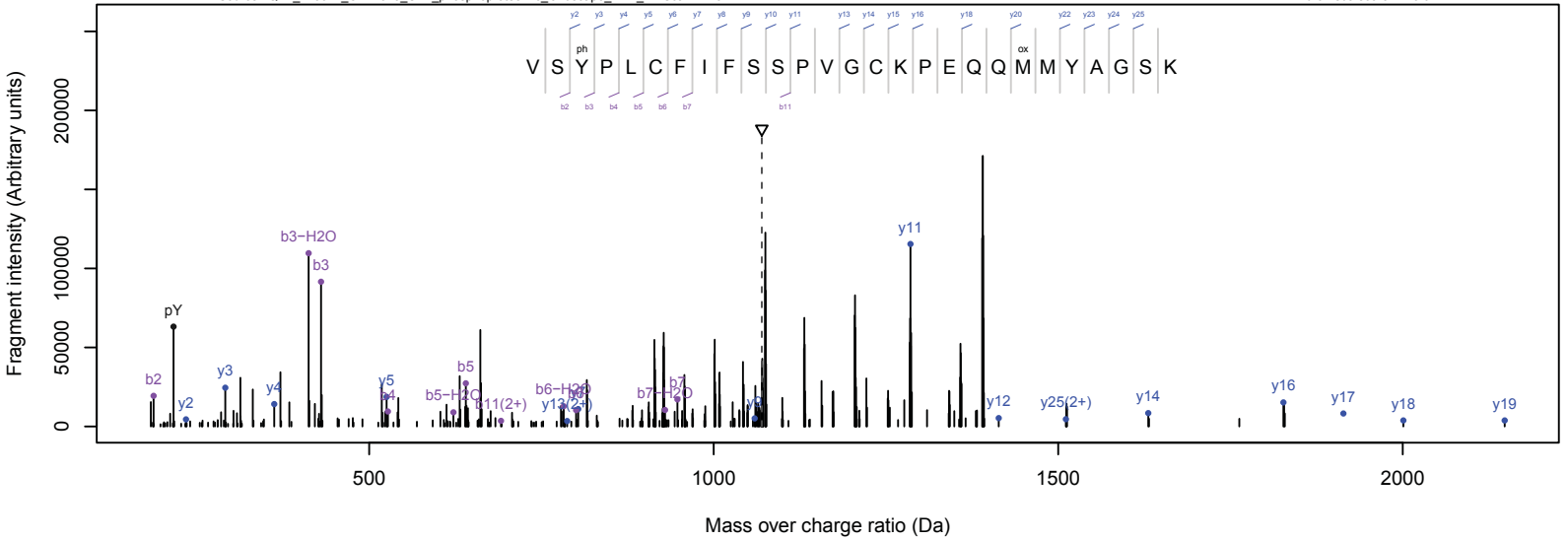
Andromeda score : 110.08



VSY(ph)PLCFIFSSPVGCKPEQMQMMYAGSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 47347

Andromeda score : 110.01





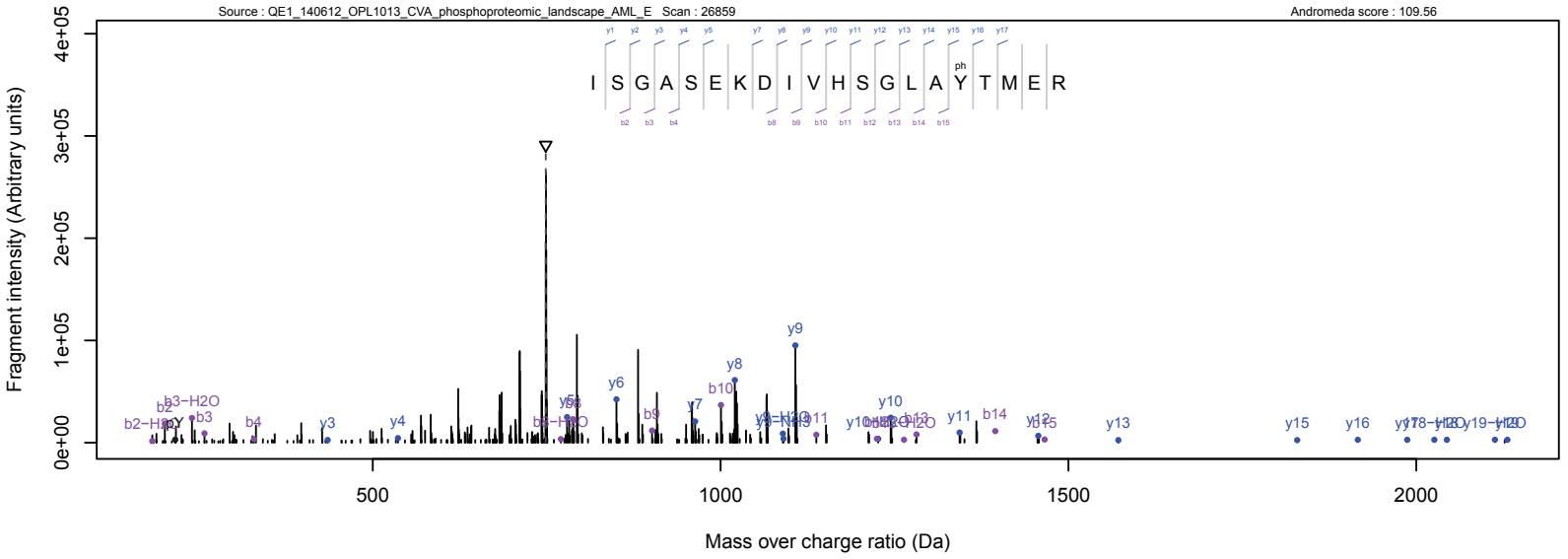




ISGASEKDIVHSGLAY(ph)TMER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 26859

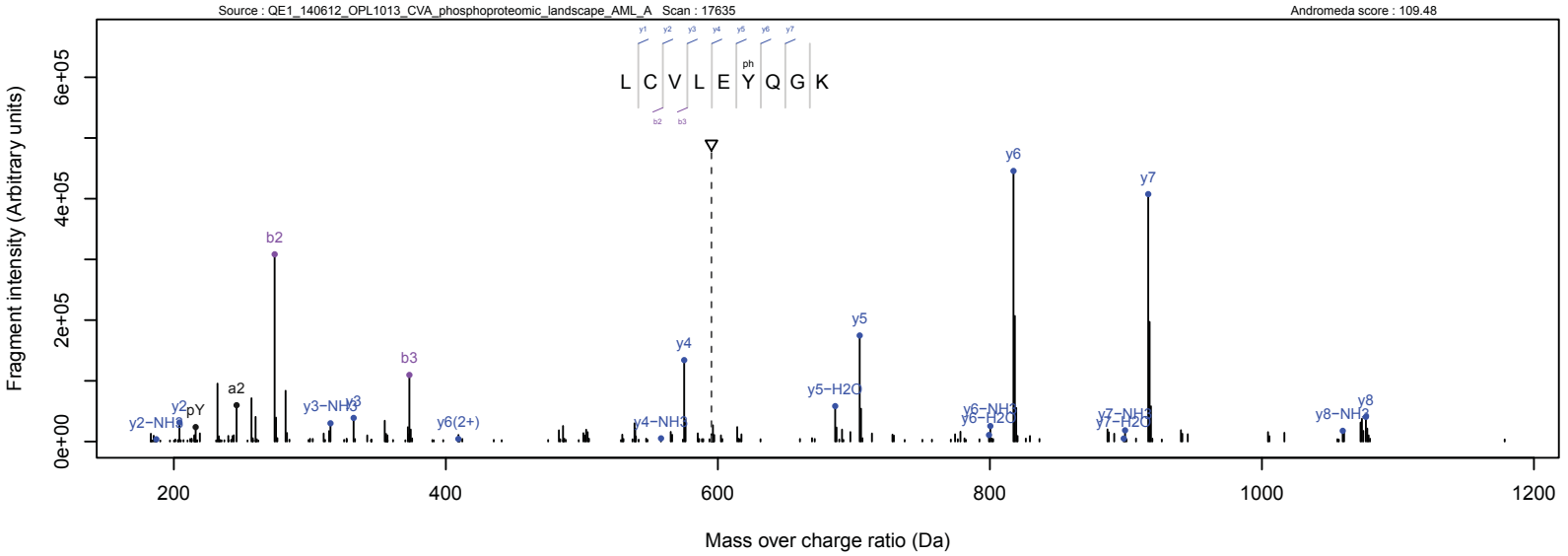
Andromeda score : 109.56



LCVLEY(ph)QGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17635

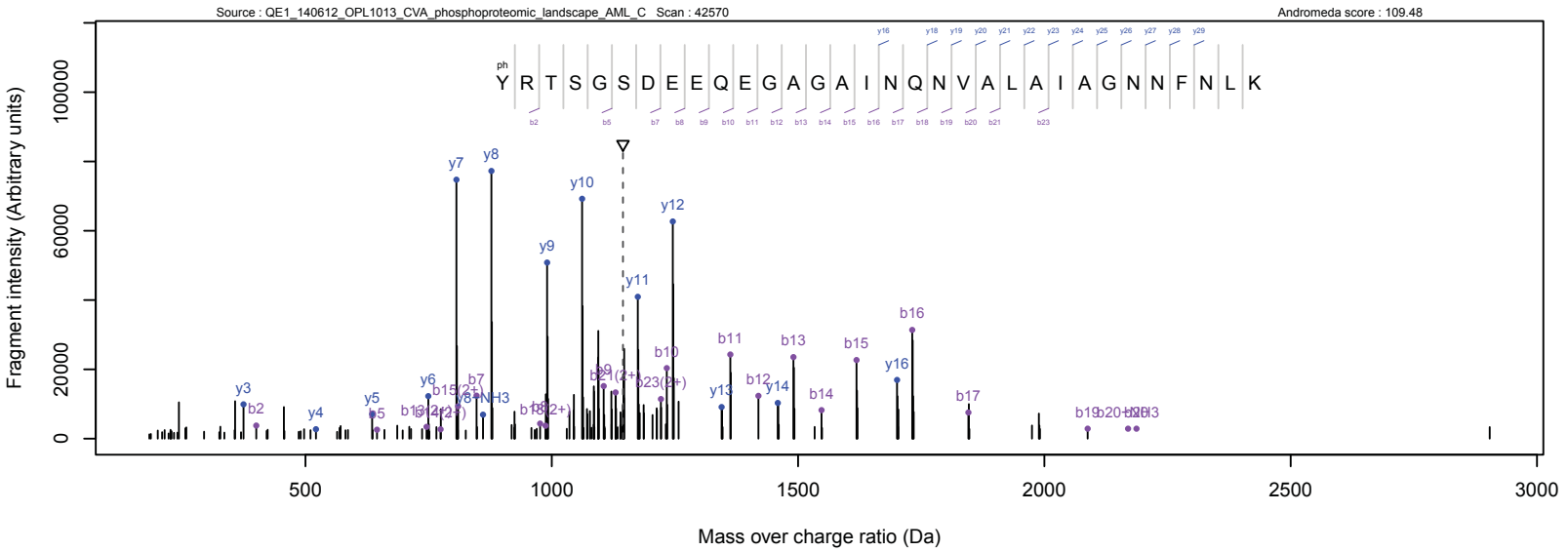
Andromeda score : 109.48



Y(ph)RTSGSDEEQEGAGAINQNVALAIAGNNFNLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42570

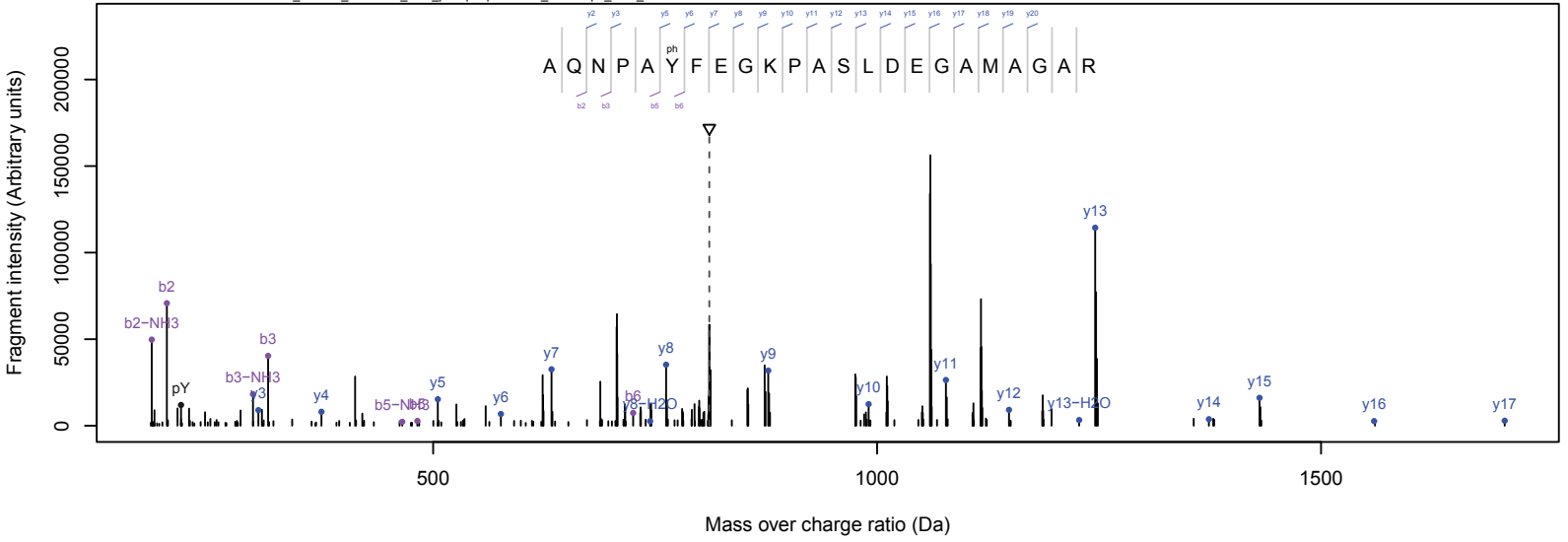
Andromeda score : 109.48



AQNPAY(ph)FEGKPSLDEGAMAGAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25767

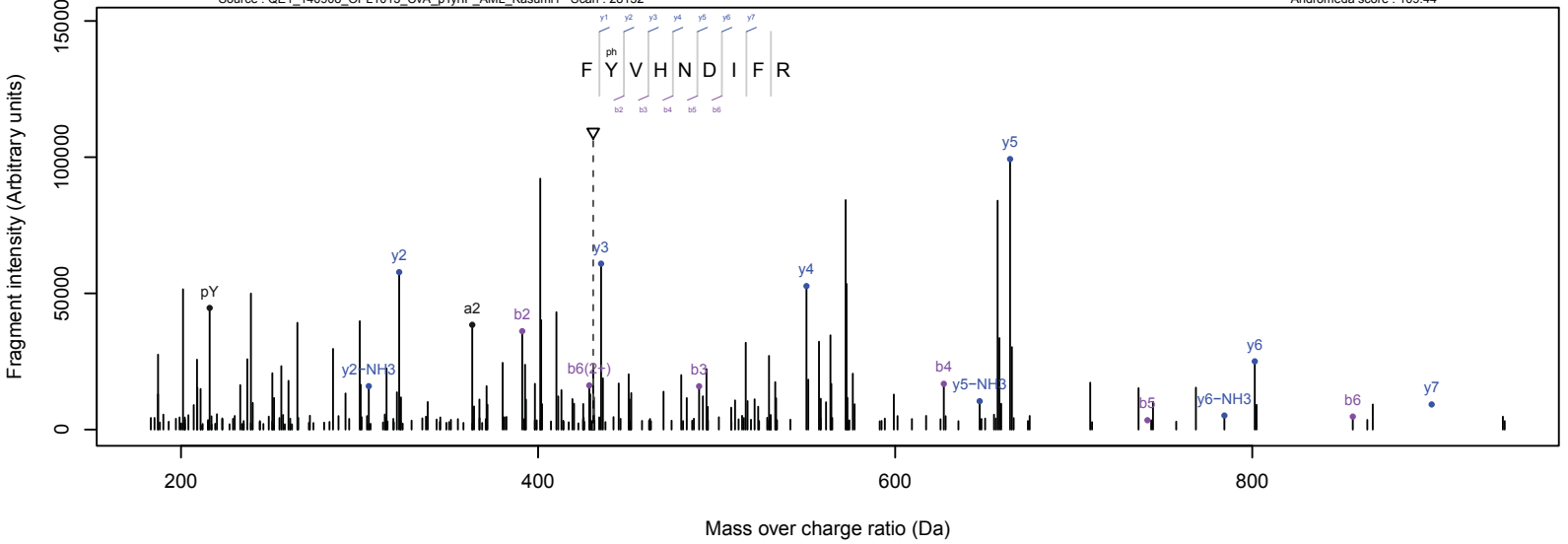
Andromeda score : 109.44



FY(ph)VHNDIFR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 28152

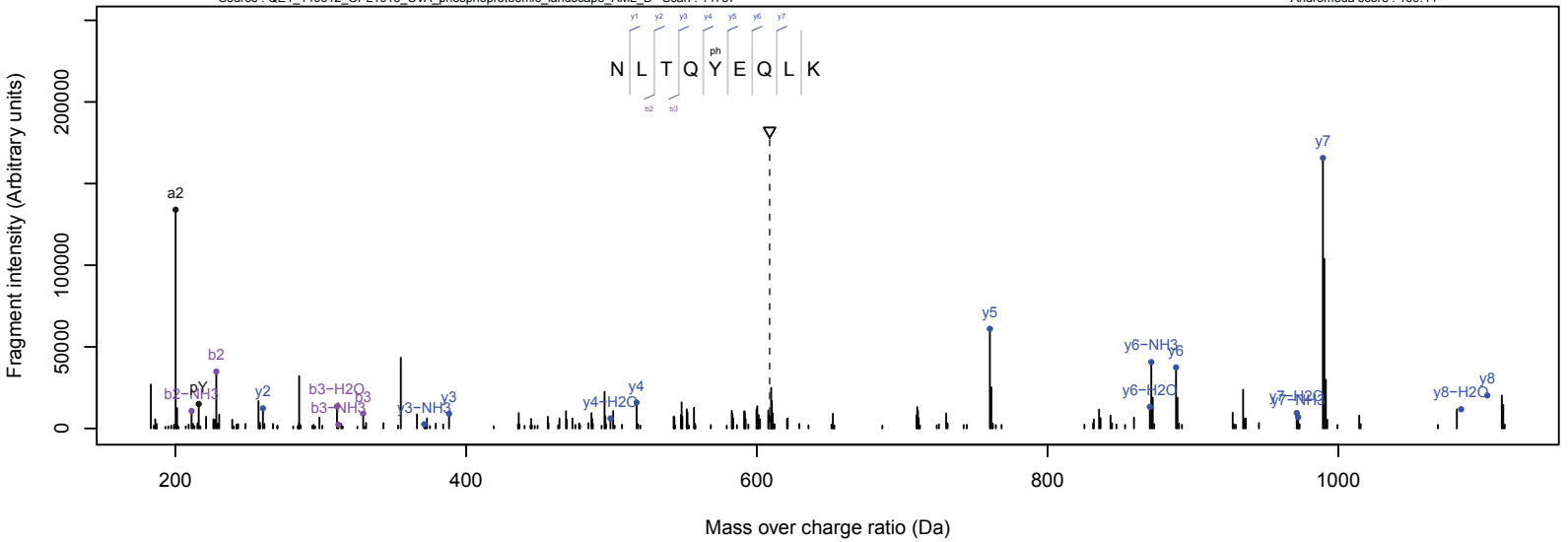
Andromeda score : 109.44



NLTQY(ph)EQLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14757

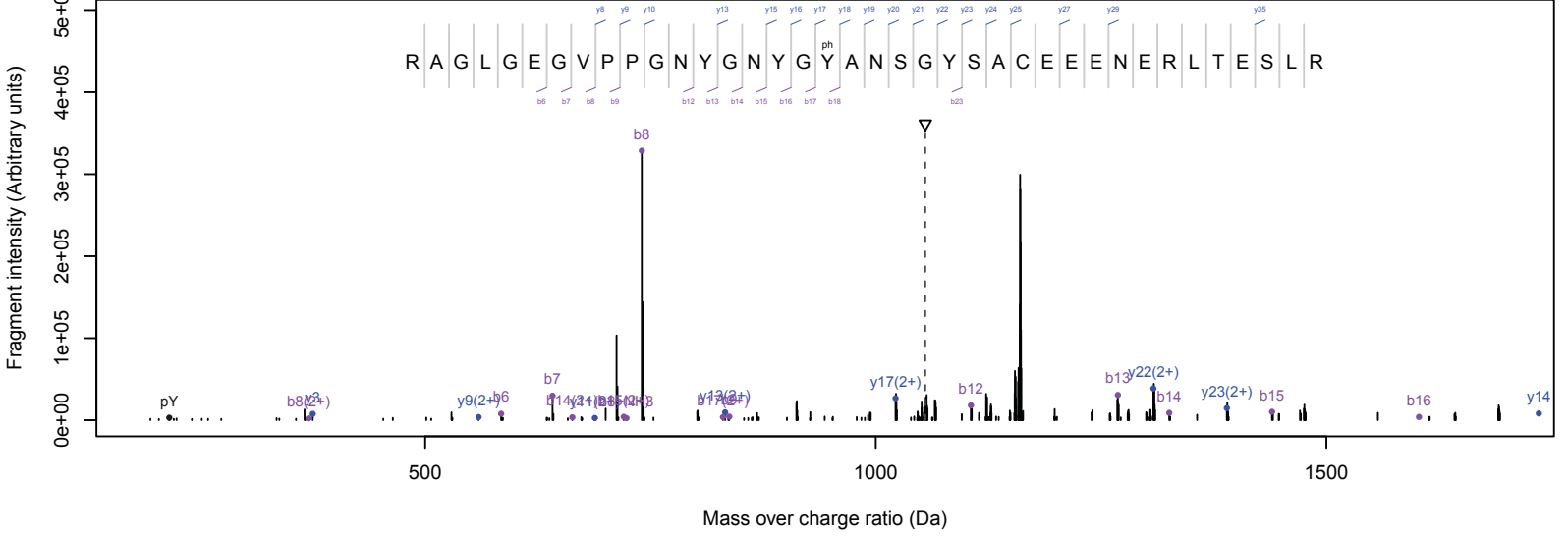
Andromeda score : 109.44



RAGLGEGVPPGNYGNYGY(ph)ANSYGSACEEENERLTESLR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31782

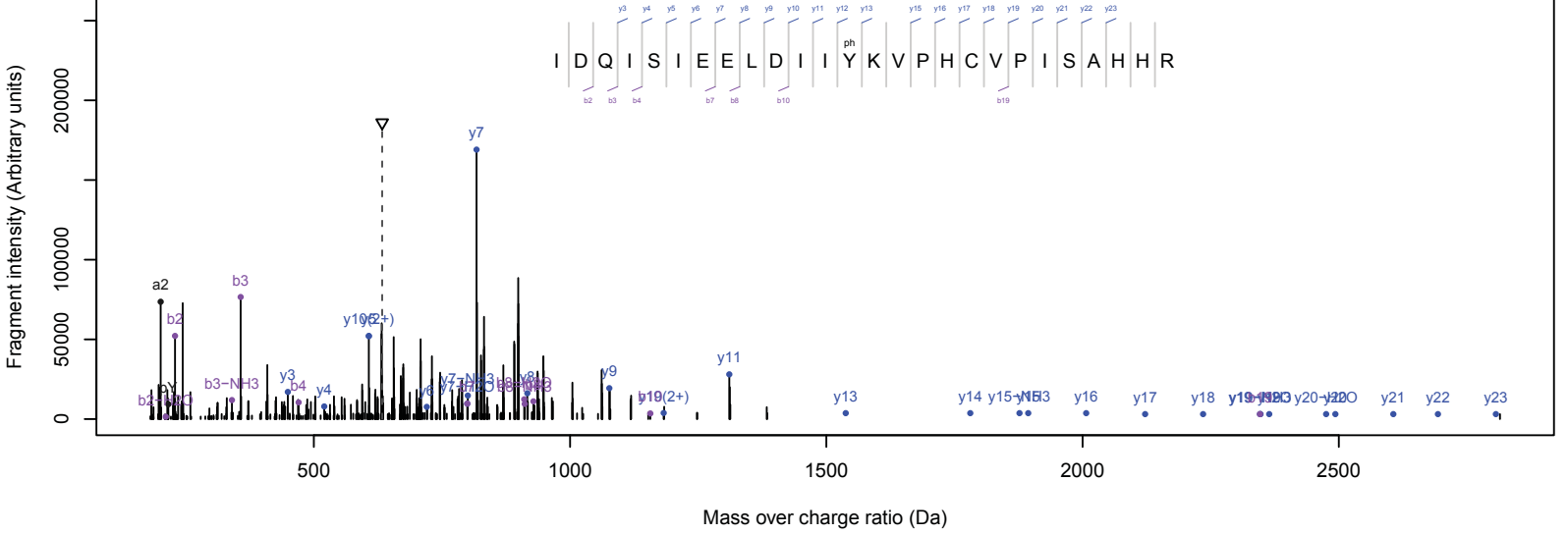
Andromeda score : 109.27



IDQISIEELDIY(ph)KVPHCVPISAHHR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44194

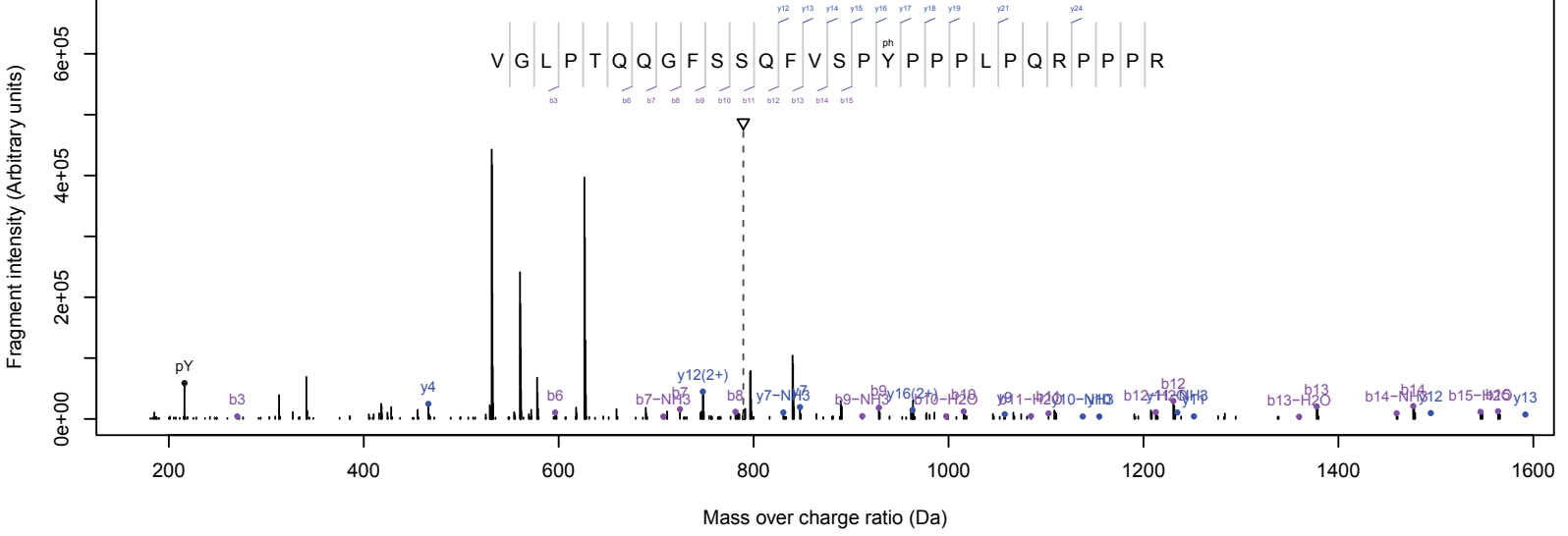
Andromeda score : 109.25



VGLPTQQGFSSQFVSPYPPLPQRPPR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40303

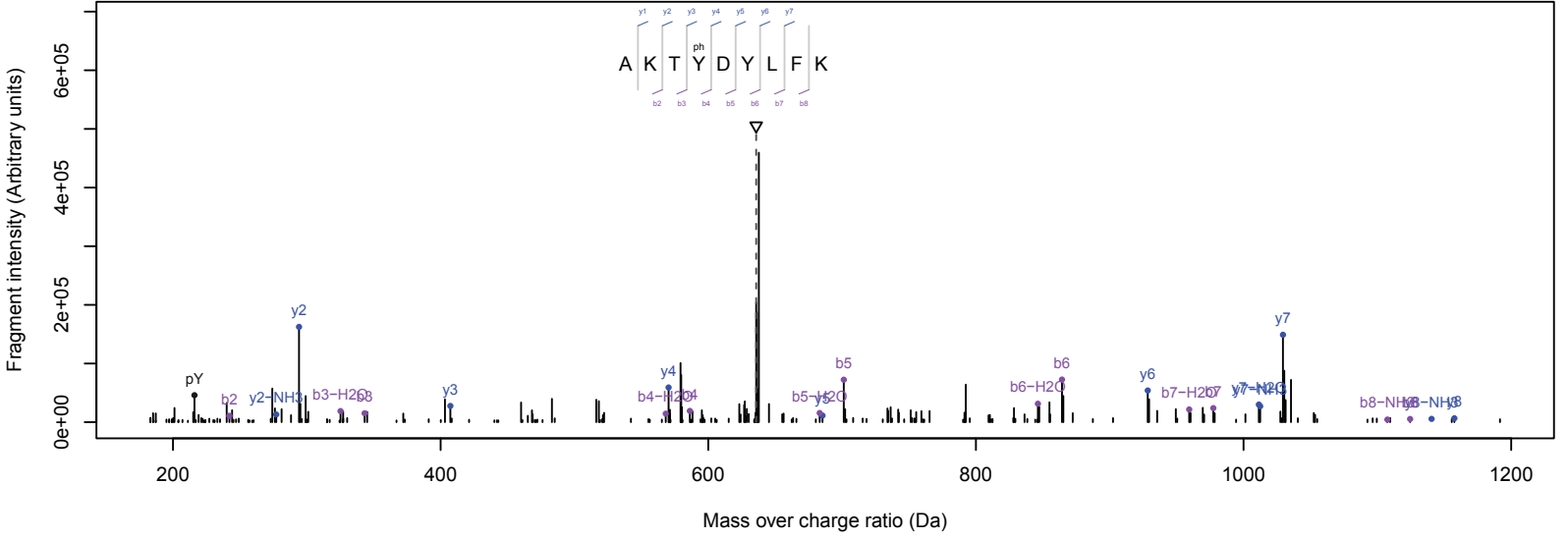
Andromeda score : 109.23



(ac)AKTY(ph)DYLFK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31223

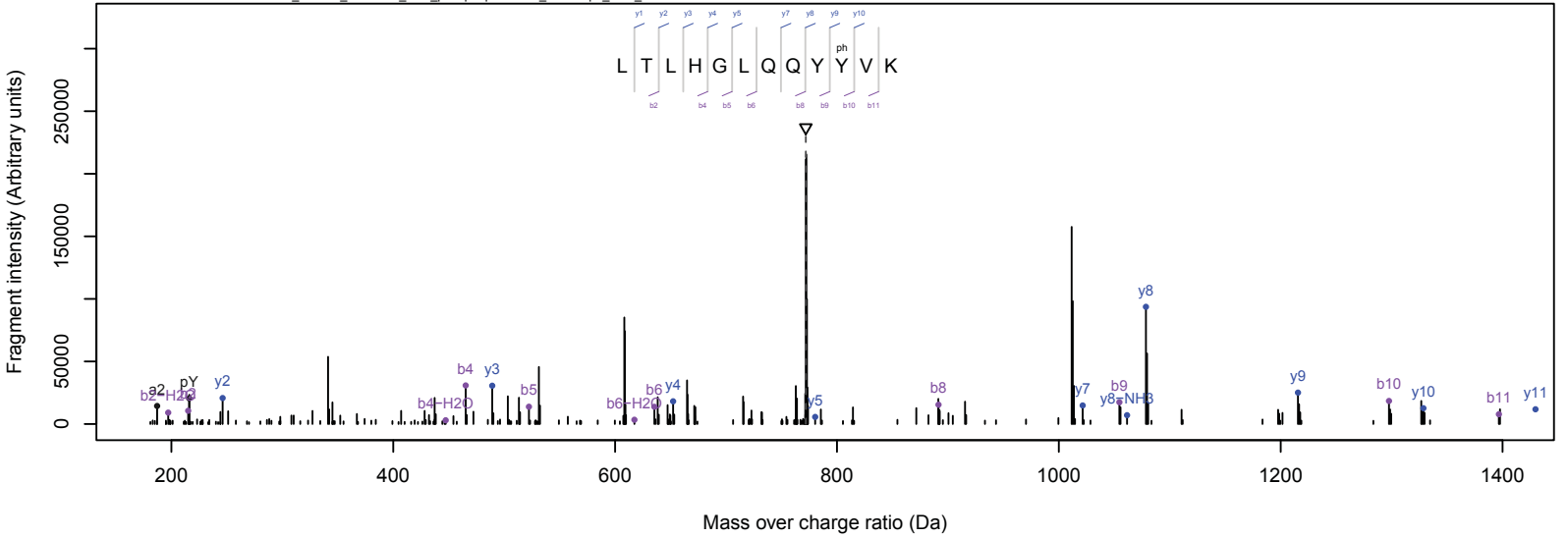
Andromeda score: 109.16



LTLHGLQQYY(ph)VK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 27143

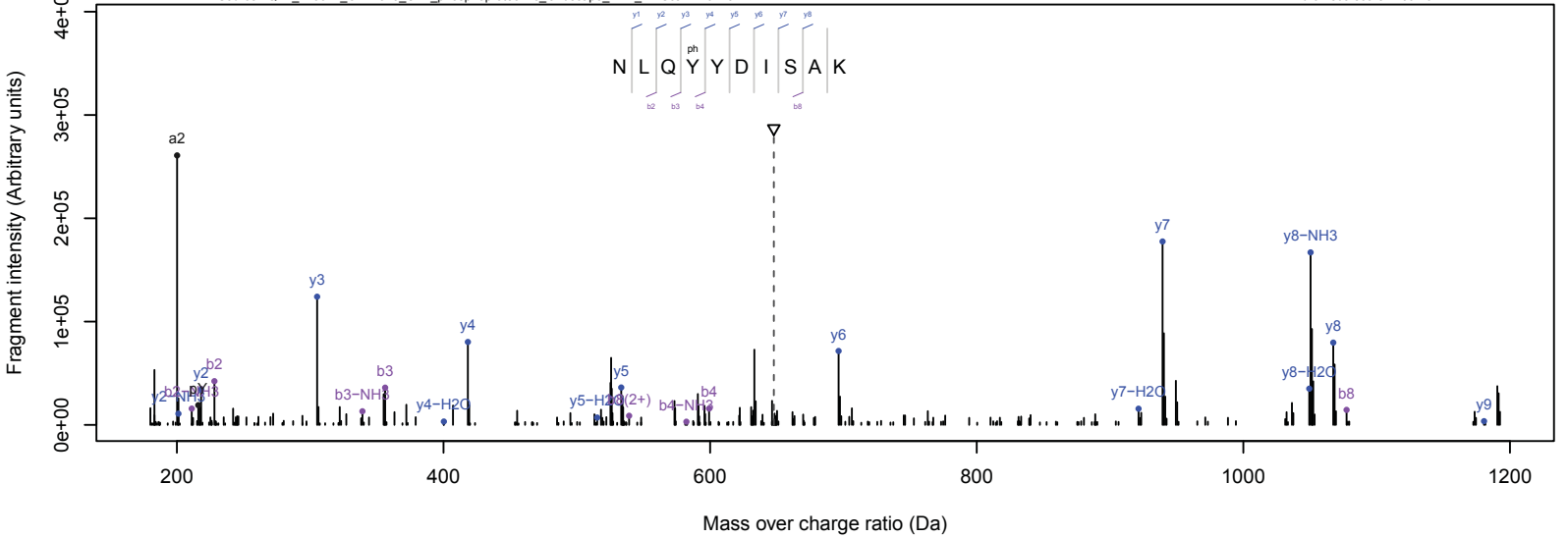
Andromeda score: 109.16



NLQY(ph)YDISAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 20413

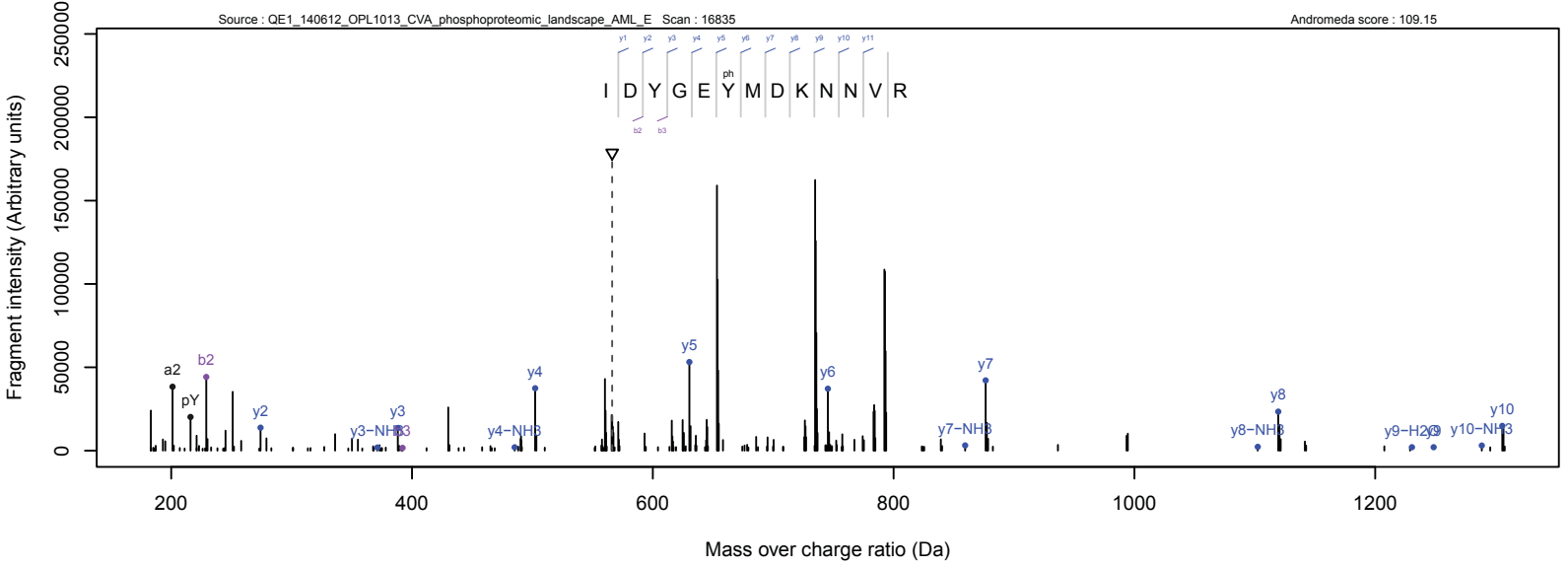
Andromeda score: 109.16



IDYGEY(ph)MDKNNVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 16835

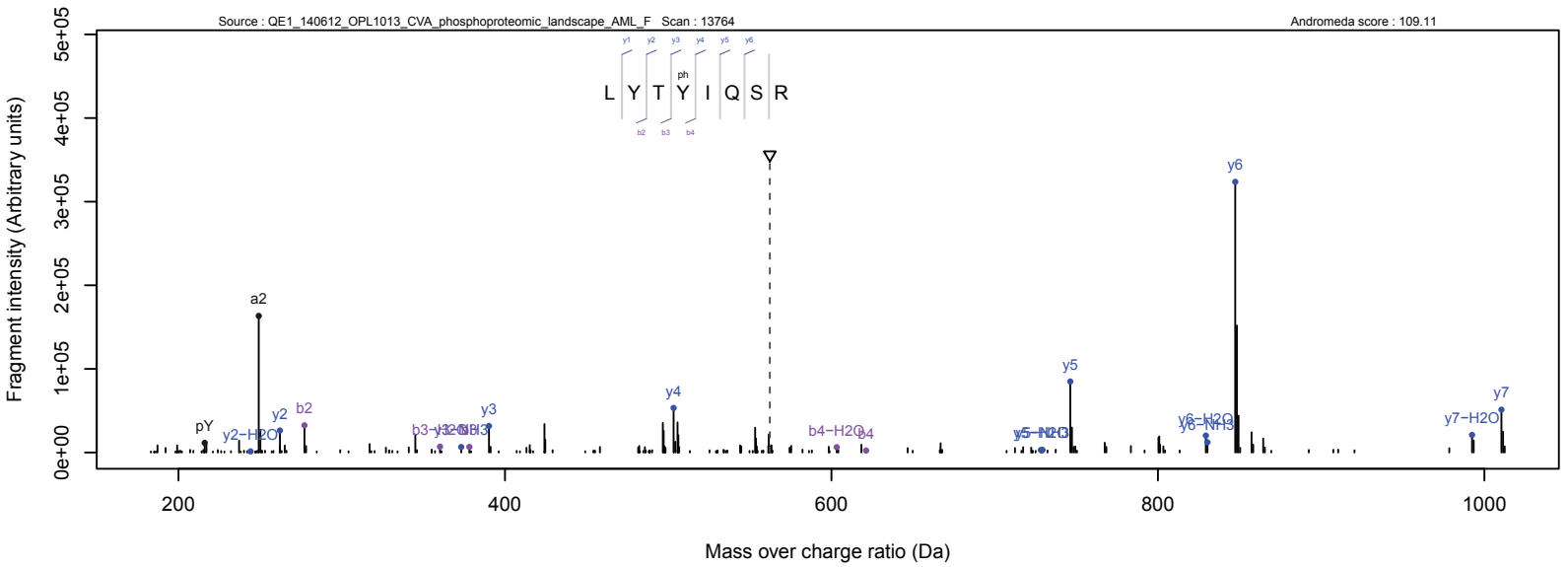
Andromeda score: 109.15



LYTY(ph)QSR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan: 13764

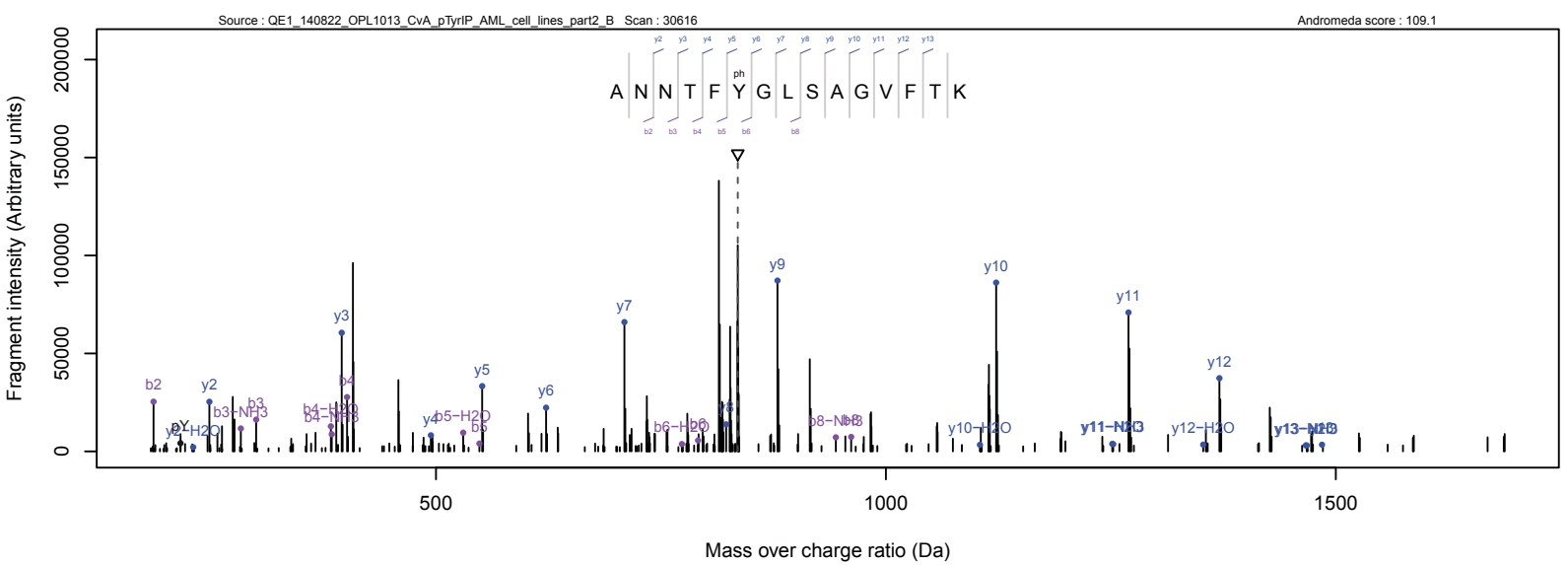
Andromeda score: 109.11



ANNTFY(ph)GLSAGVFTK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 30616

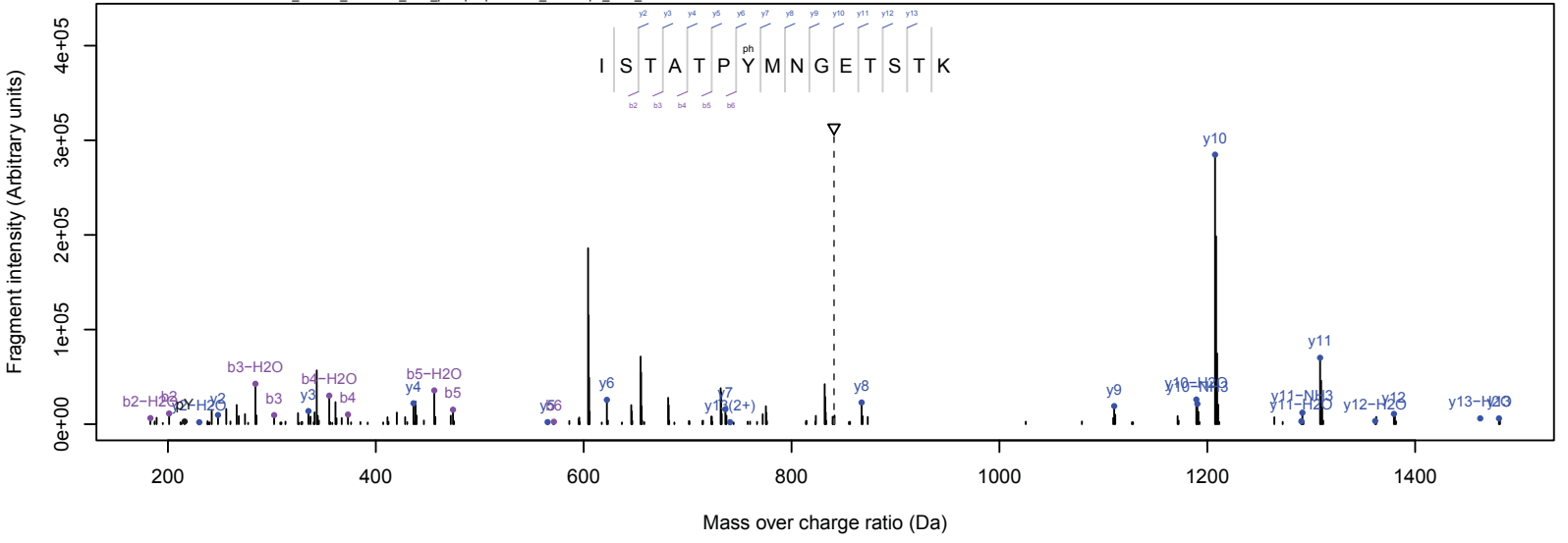
Andromeda score: 109.1



ISTATPY(ph)MNGETSTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12564

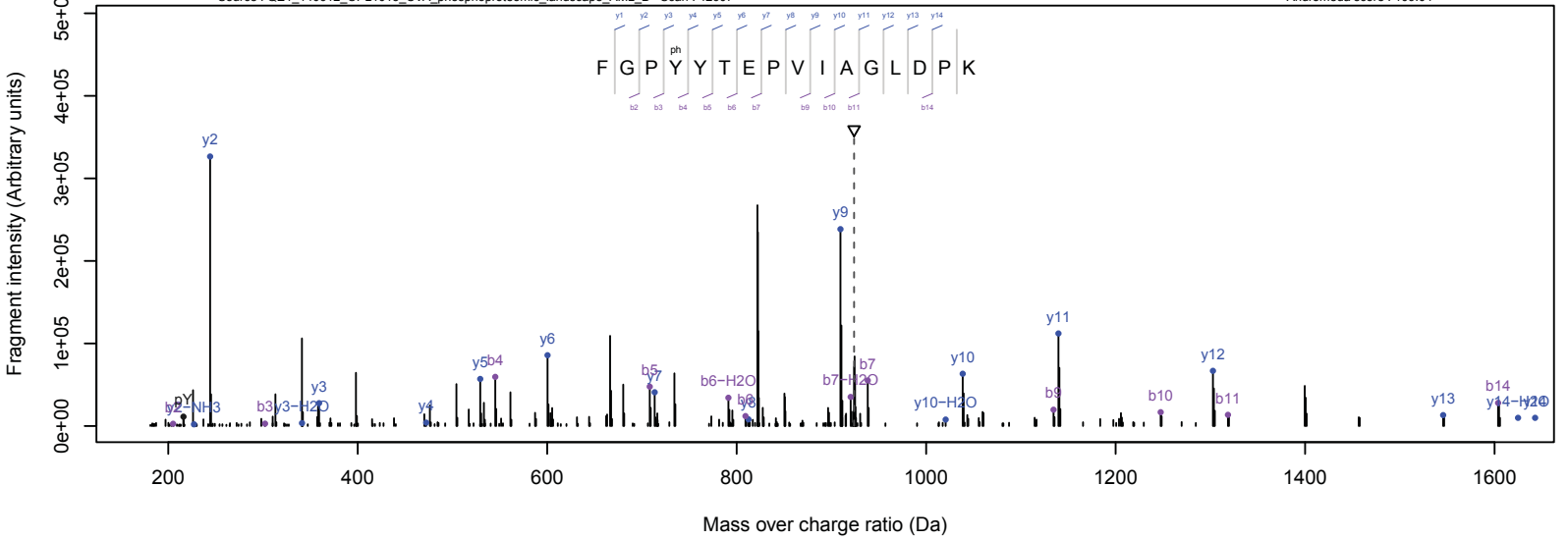
Andromeda score : 109.07



FGPY(ph)YTEPVIAGLDPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42607

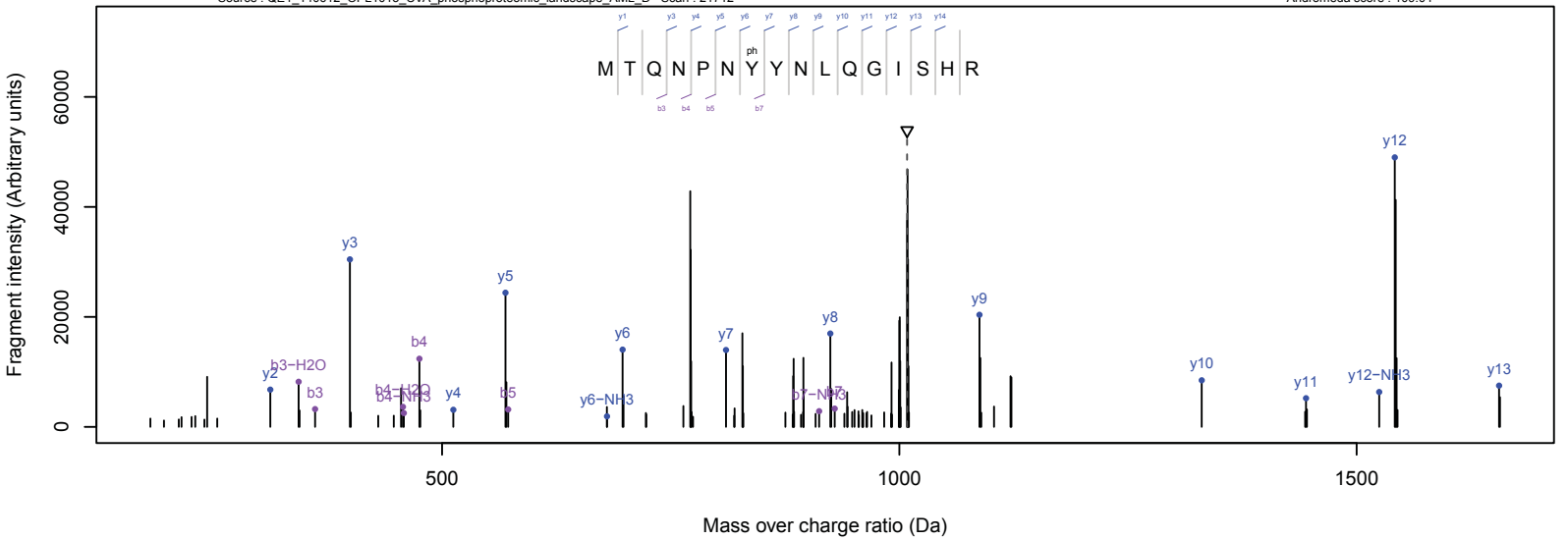
Andromeda score : 109.04



MTQNPNY(ph)YNLQGISHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 21712

Andromeda score : 109.04

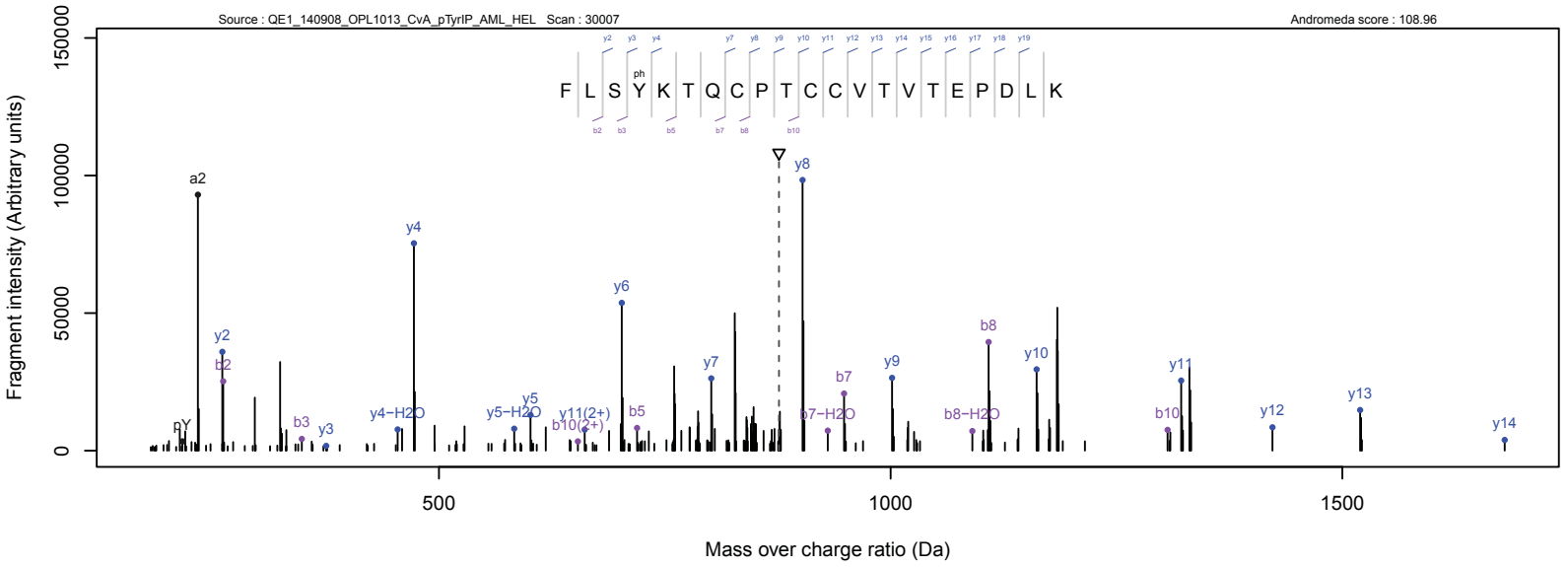




**FLSY(ph)KTQCPTCCVTVTEPDLK-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 30007

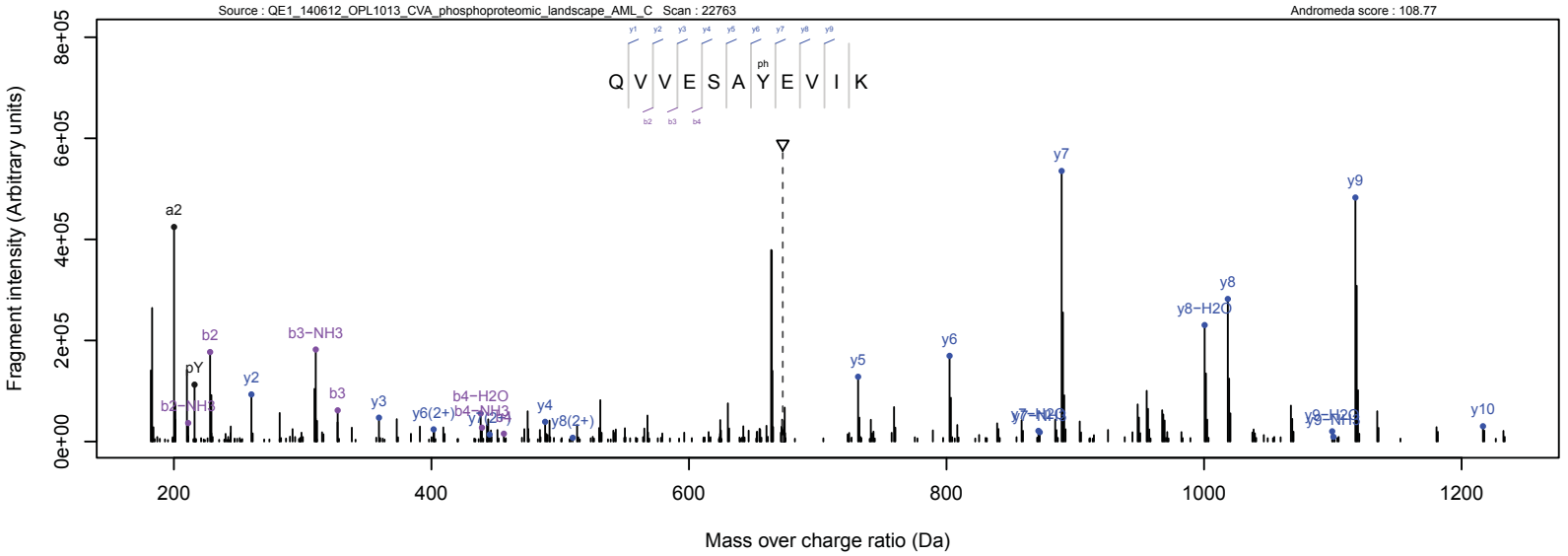
Andromeda score : 108.96



**QVVESAY(ph)EVIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22763

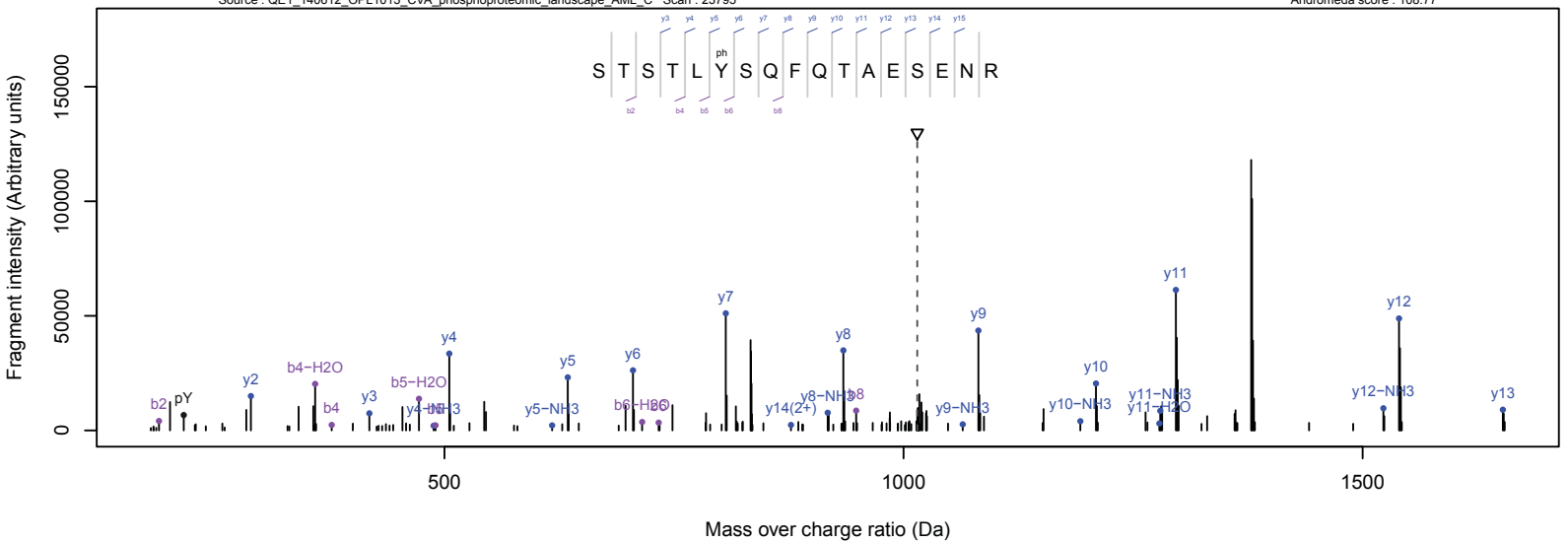
Andromeda score : 108.77



**STSTLY(ph)SQFQTAESNR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23795

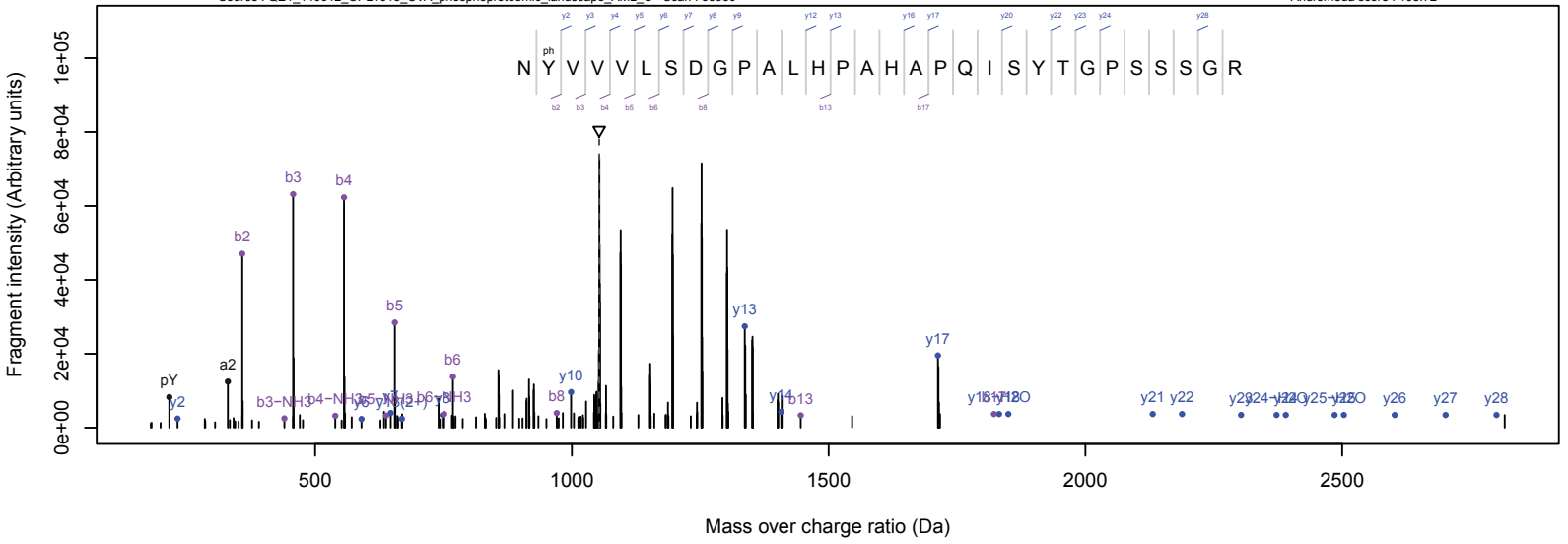
Andromeda score : 108.77



**NY(ph)VVLSDGPAALHPAHAPQISYTGPSSSGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33959

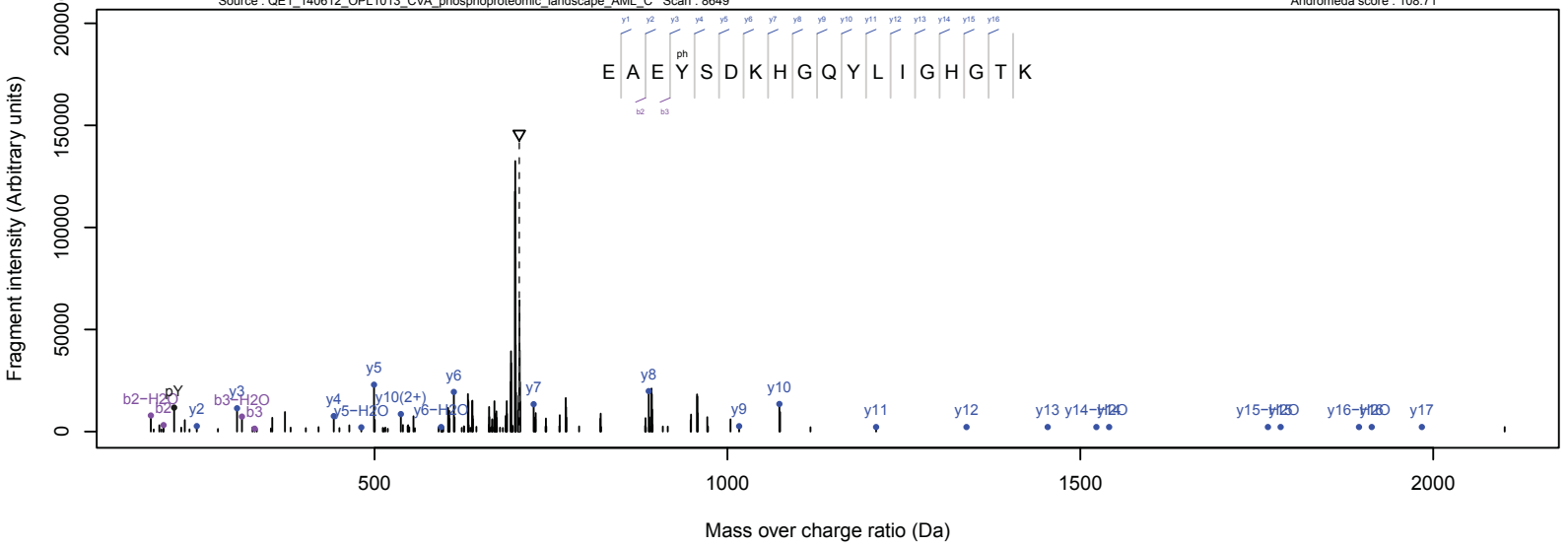
Andromeda score : 108.72



**EAEY(ph)SDKHGQYLIGHGTK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8649

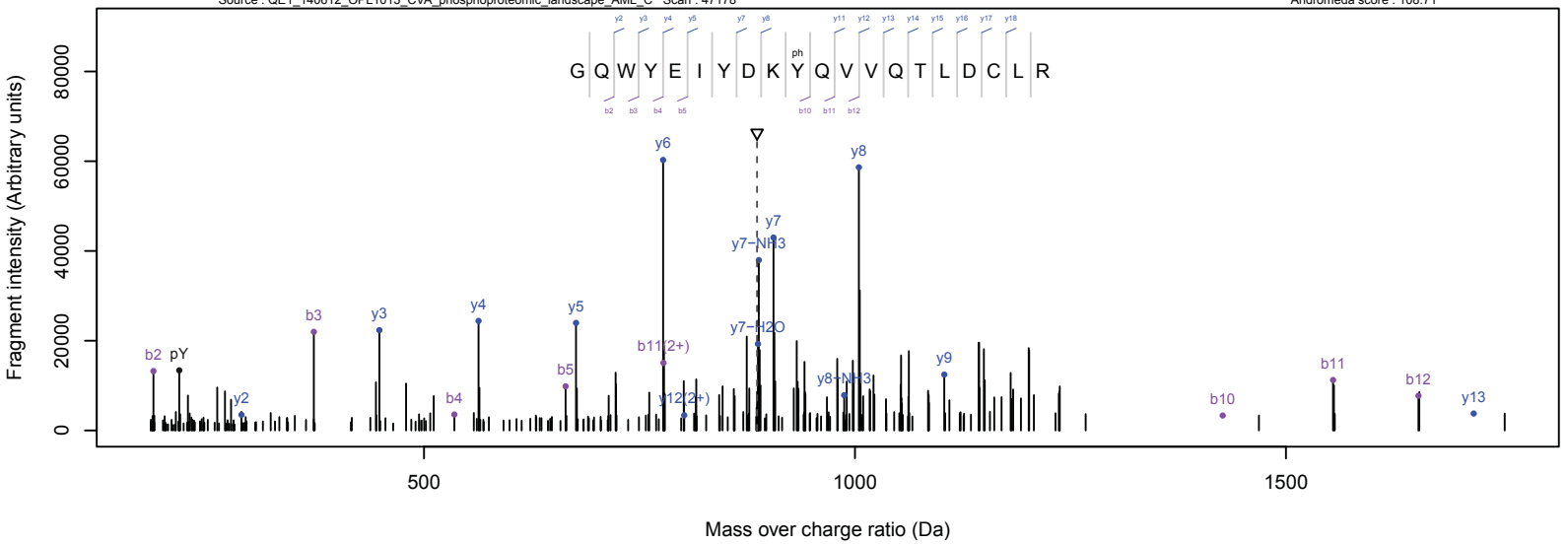
Andromeda score : 108.71



**GQWYEYDKY(ph)QVVQTLDCLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47178

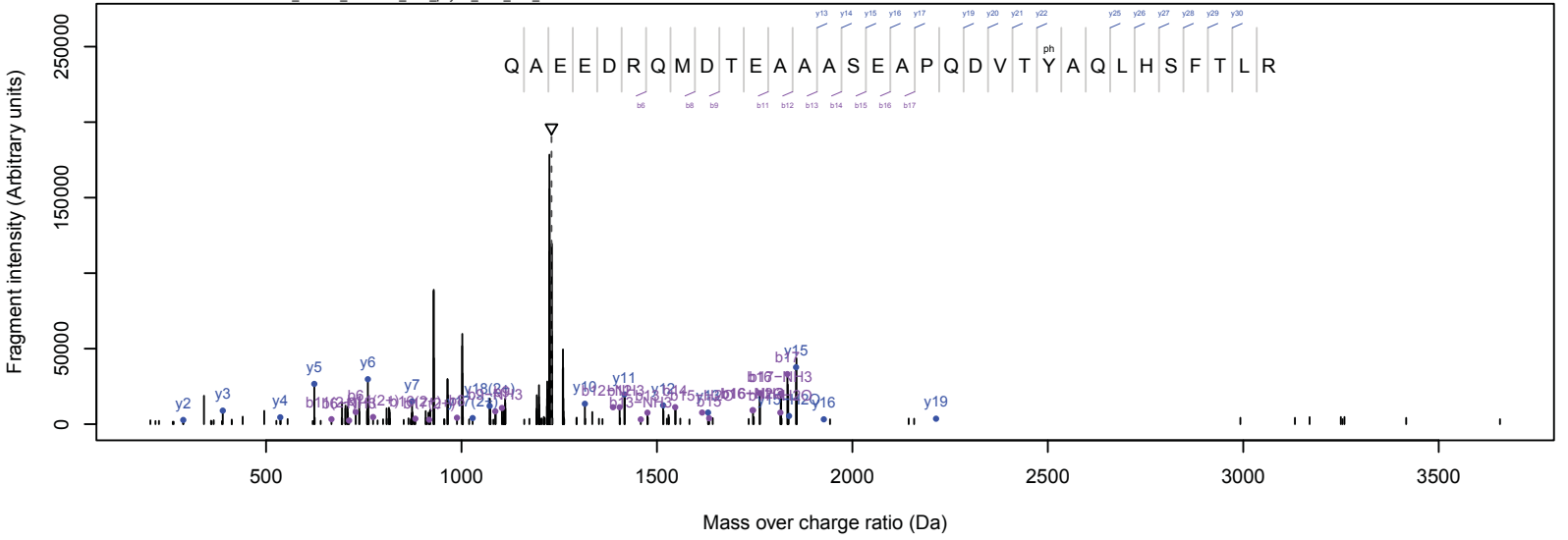
Andromeda score : 108.71



QA E E D R Q M D T E A A A S E A P Q D V T Y A Q L H S F T L R

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 38546

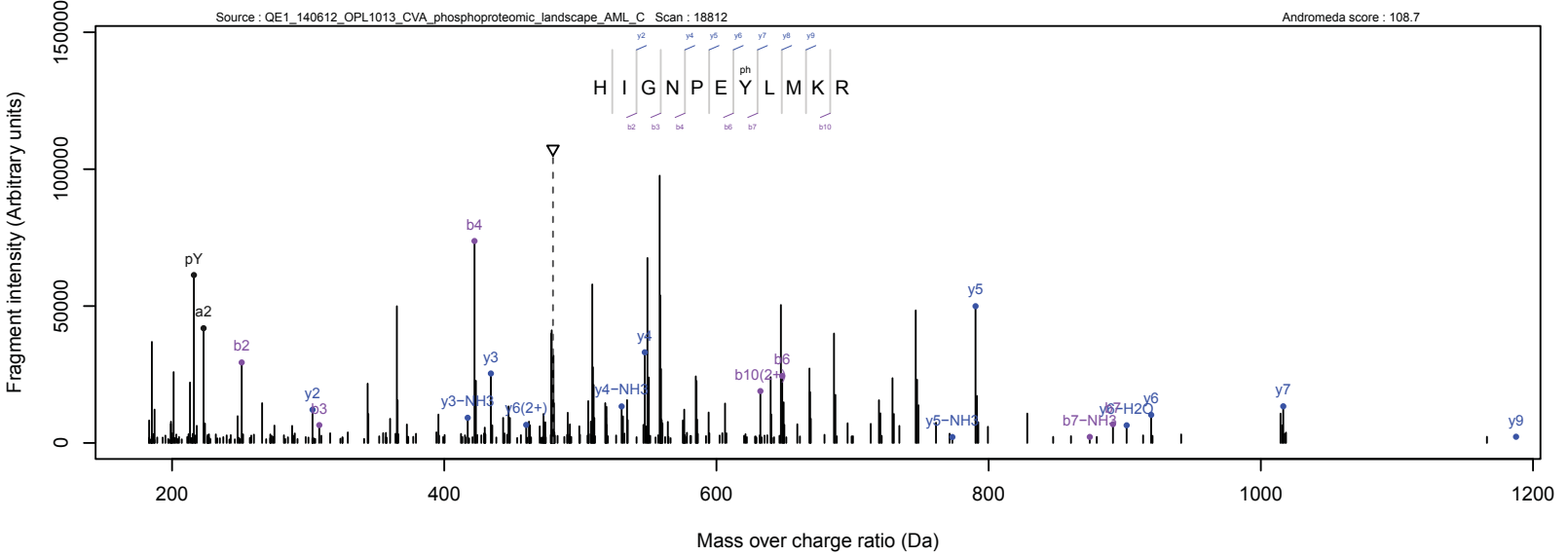
Andromeda score: 108.71



H I G N P E Y L M K R

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 18812

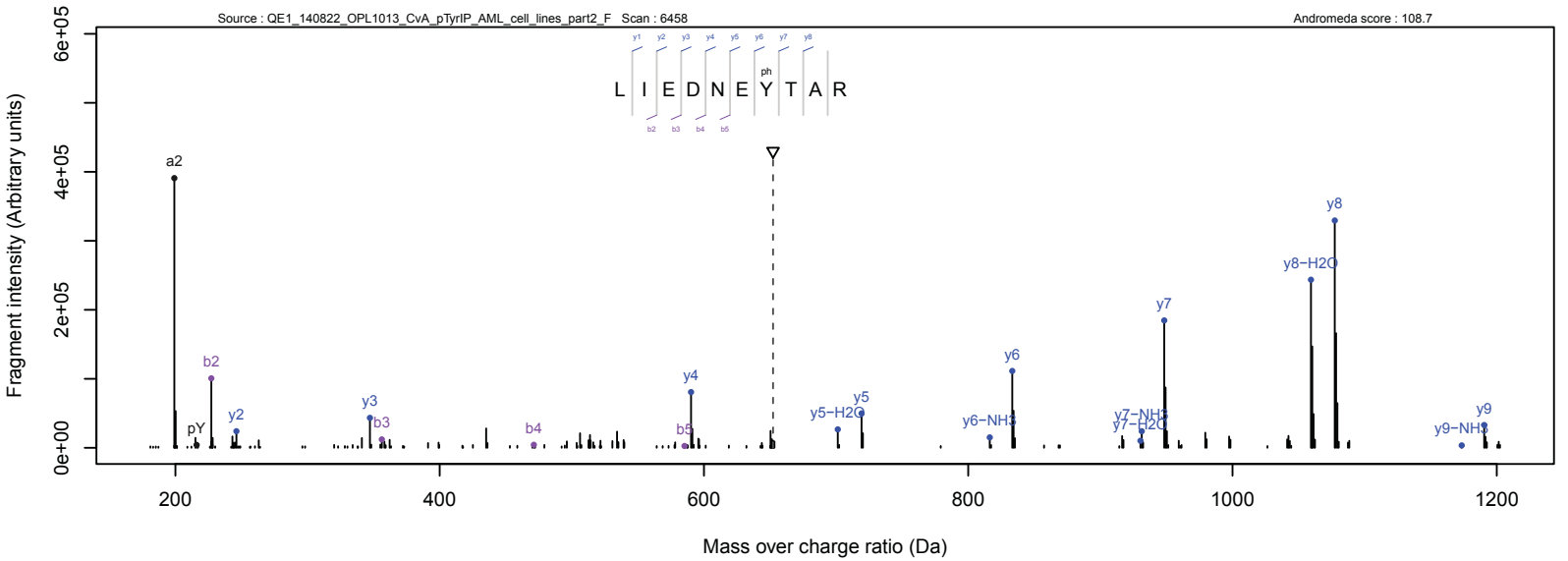
Andromeda score: 108.7



L I E D N E Y T A R

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan: 6458

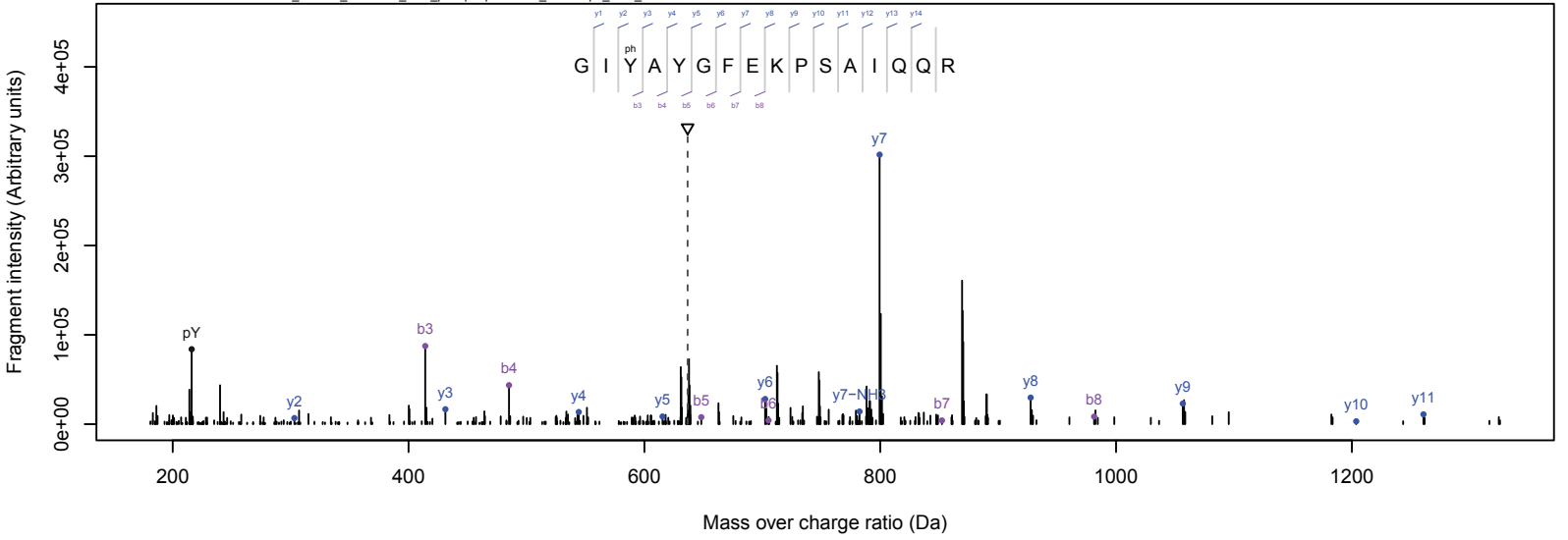
Andromeda score: 108.7



**GIY(ph)AYGFEKPSAIQQR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26718

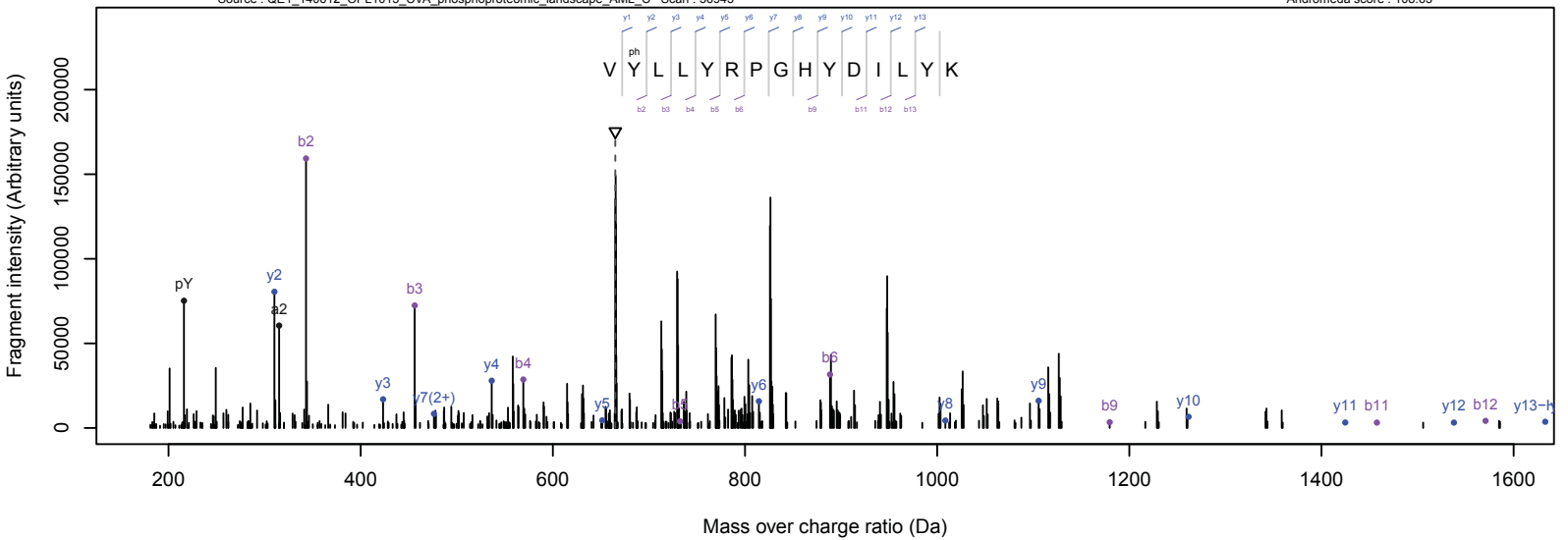
Andromeda score: 108.63



**VY(ph)LLYRPGHYDILYK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36943

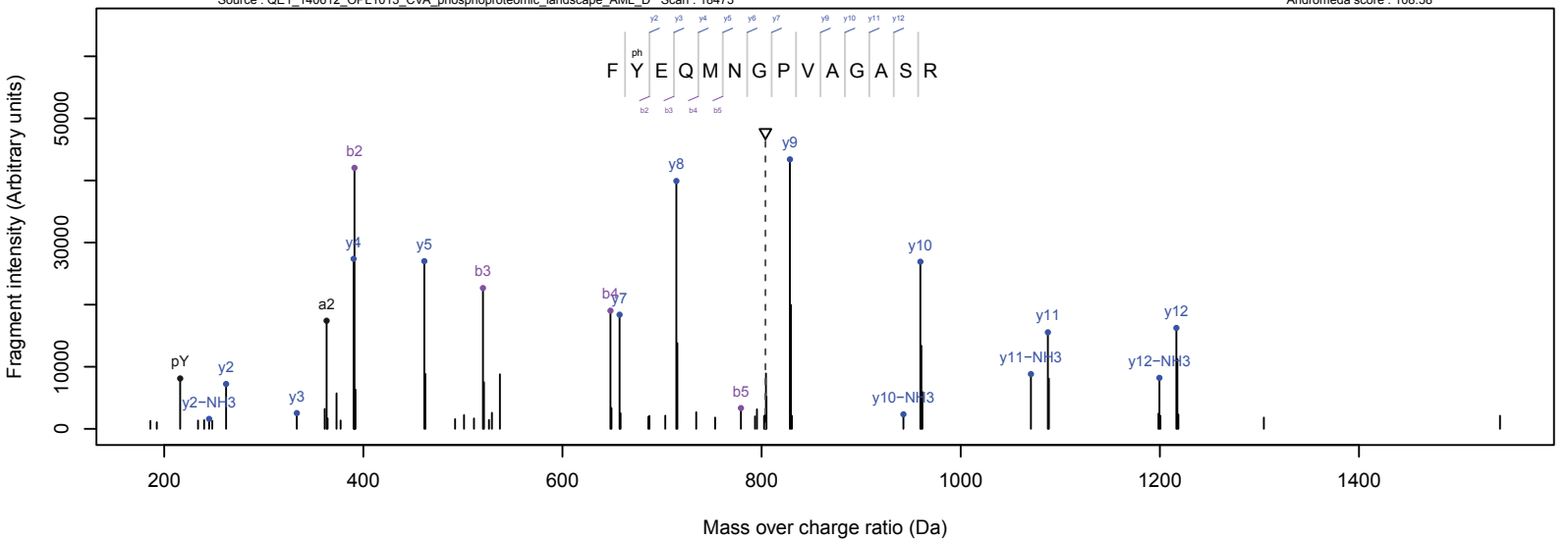
Andromeda score: 108.63



**FY(ph)EQMNGPVAGASR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 18473

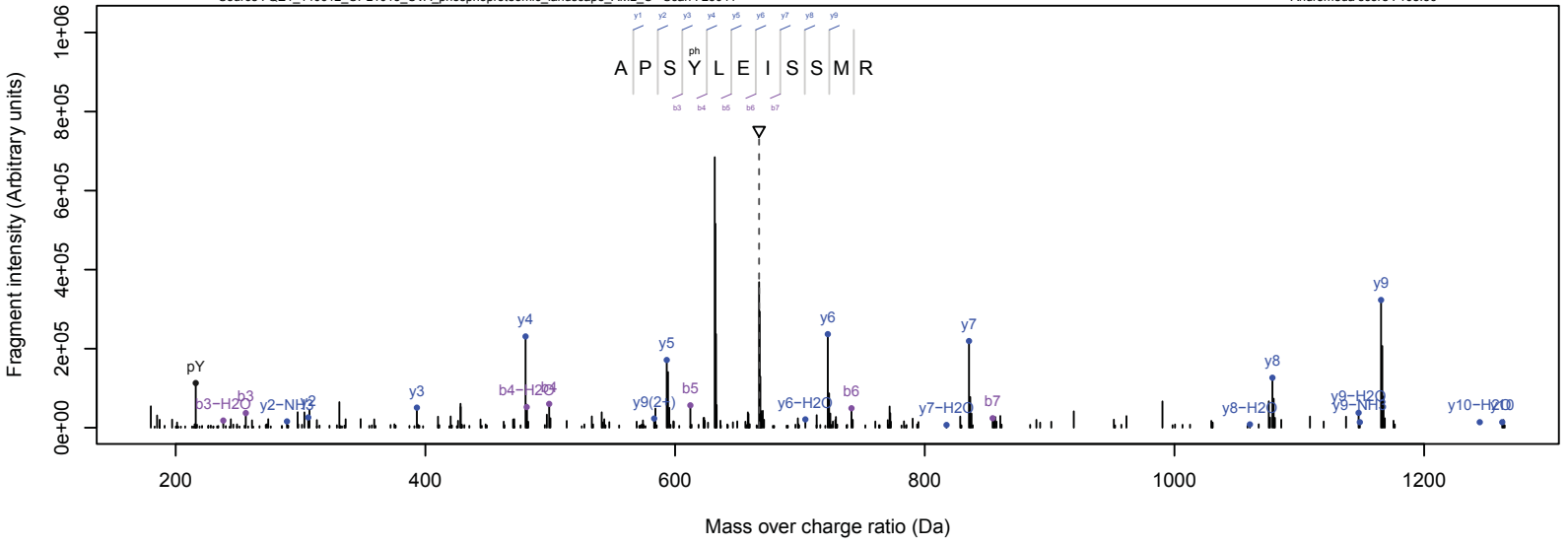
Andromeda score: 108.58



APSY(ph)LEISSMR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25041

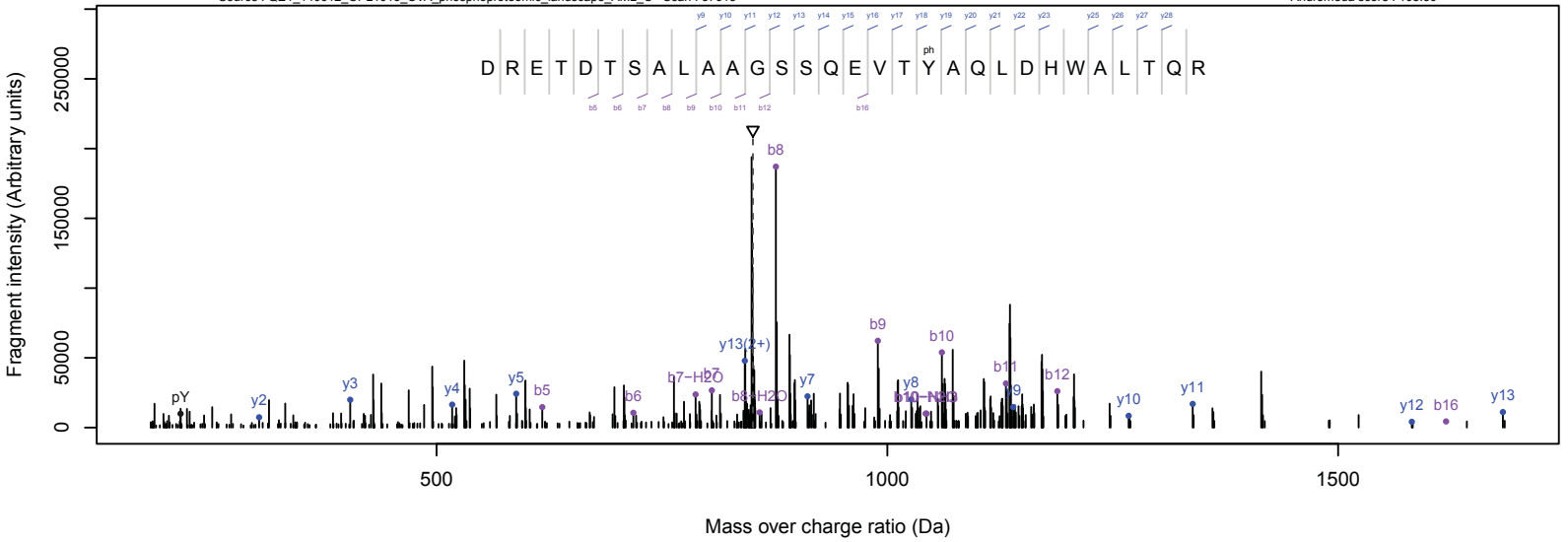
Andromeda score : 108.56



DRETDTSALAAGSSQEVY(ph)AQLDHWALTQR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37615

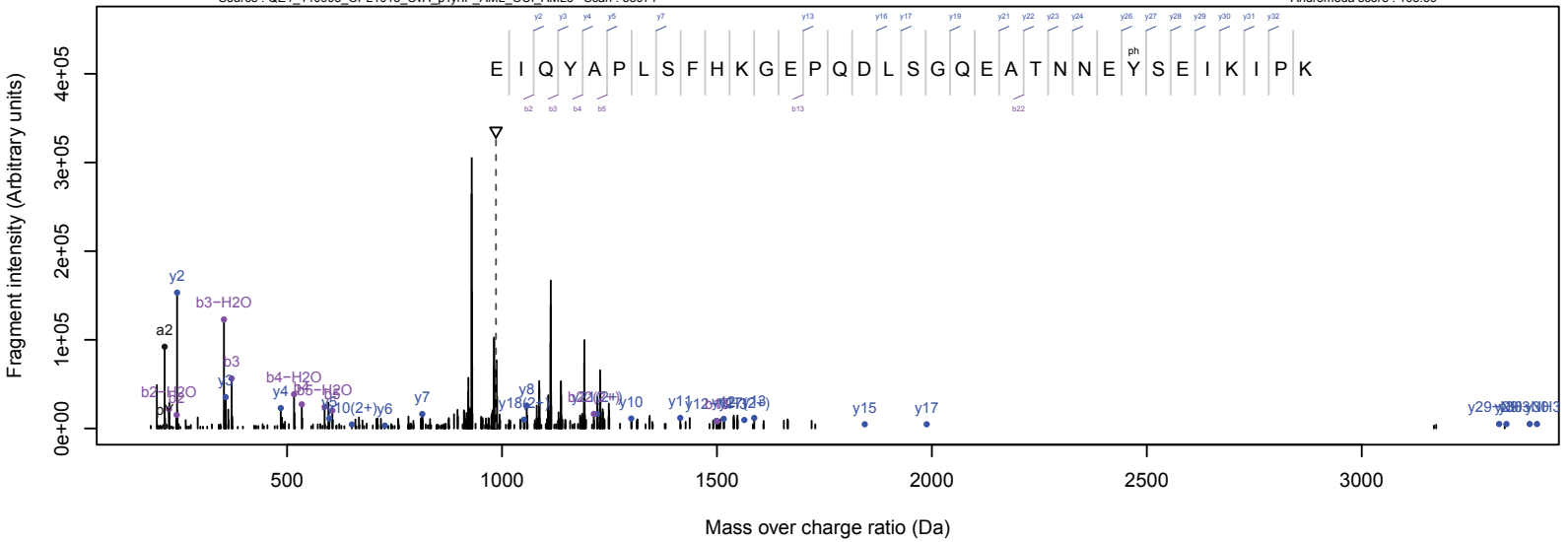
Andromeda score : 108.39



EIQYAPLSFHKGEPQDLSGQEATNNEY(ph)SEIKIPK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 35971

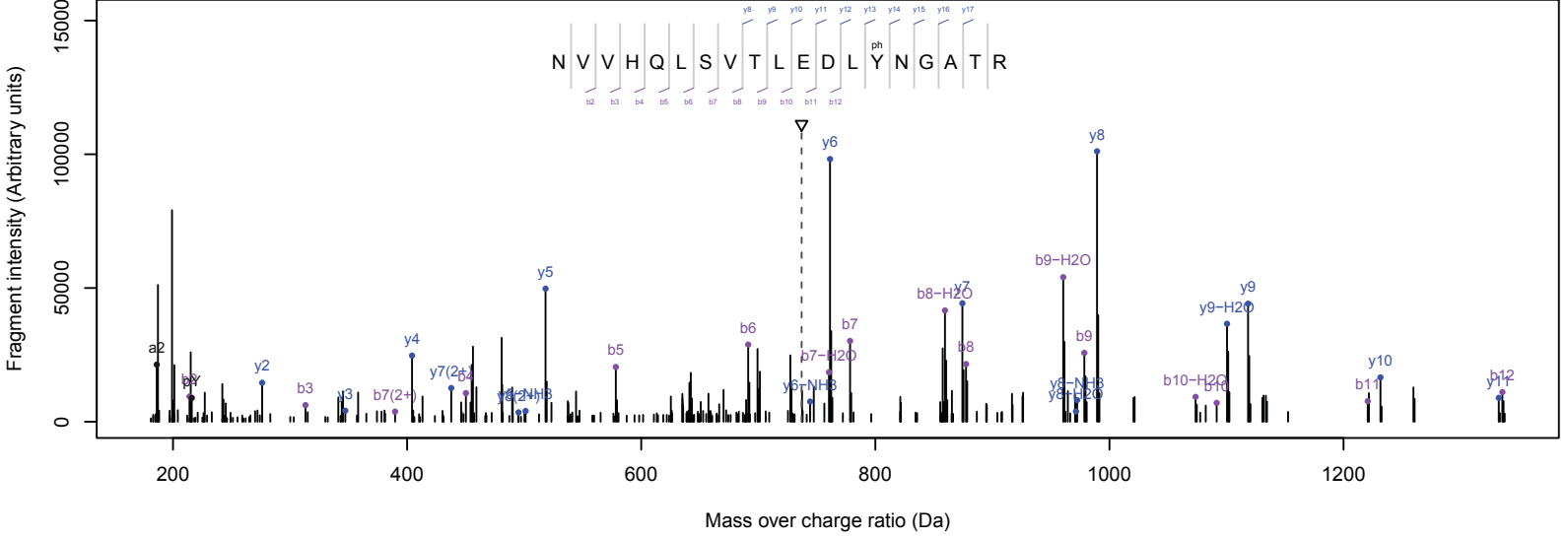
Andromeda score : 108.39



NVVHQLSVTLEDLYNGATR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 44538

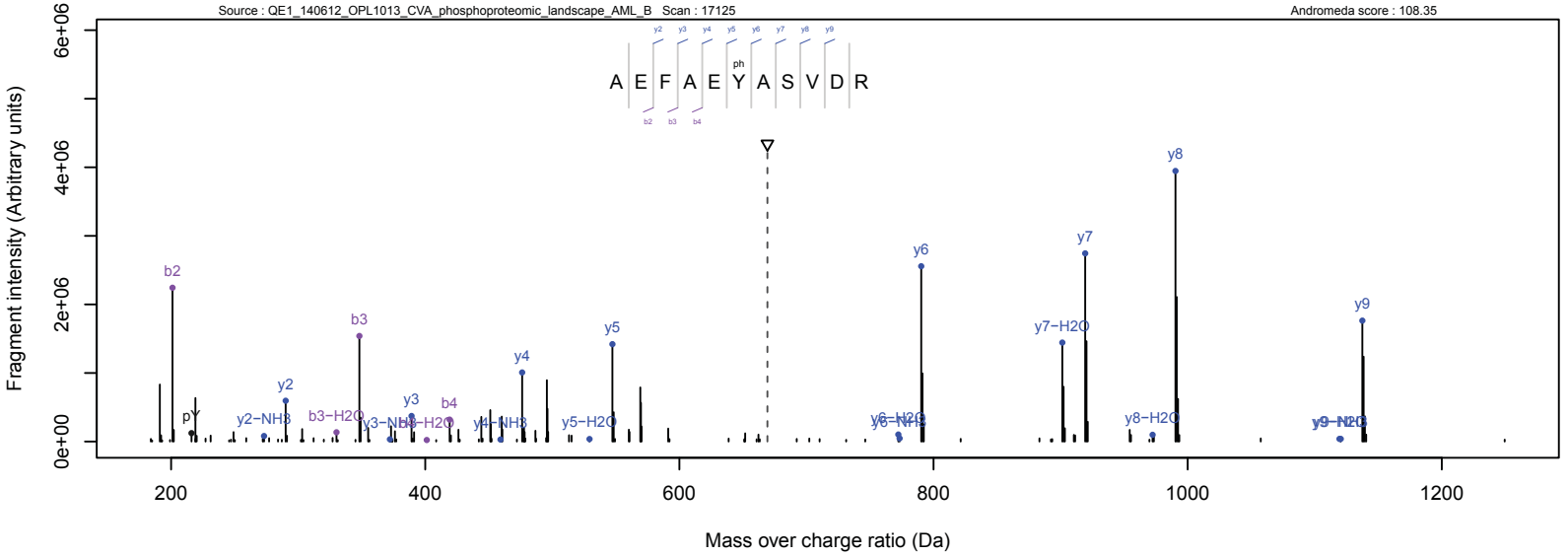
Andromeda score : 108.39



AEFAEY(ph)ASVDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17125

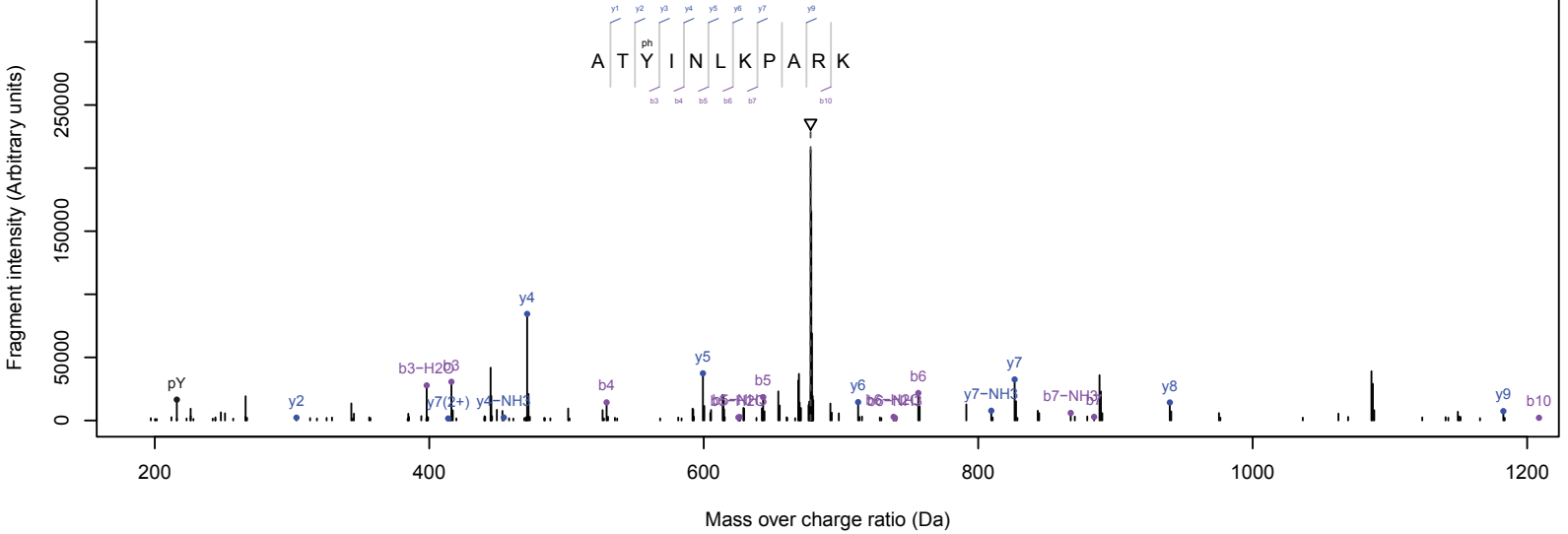
Andromeda score : 108.35



ATY(ph)INLKPAARK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10161

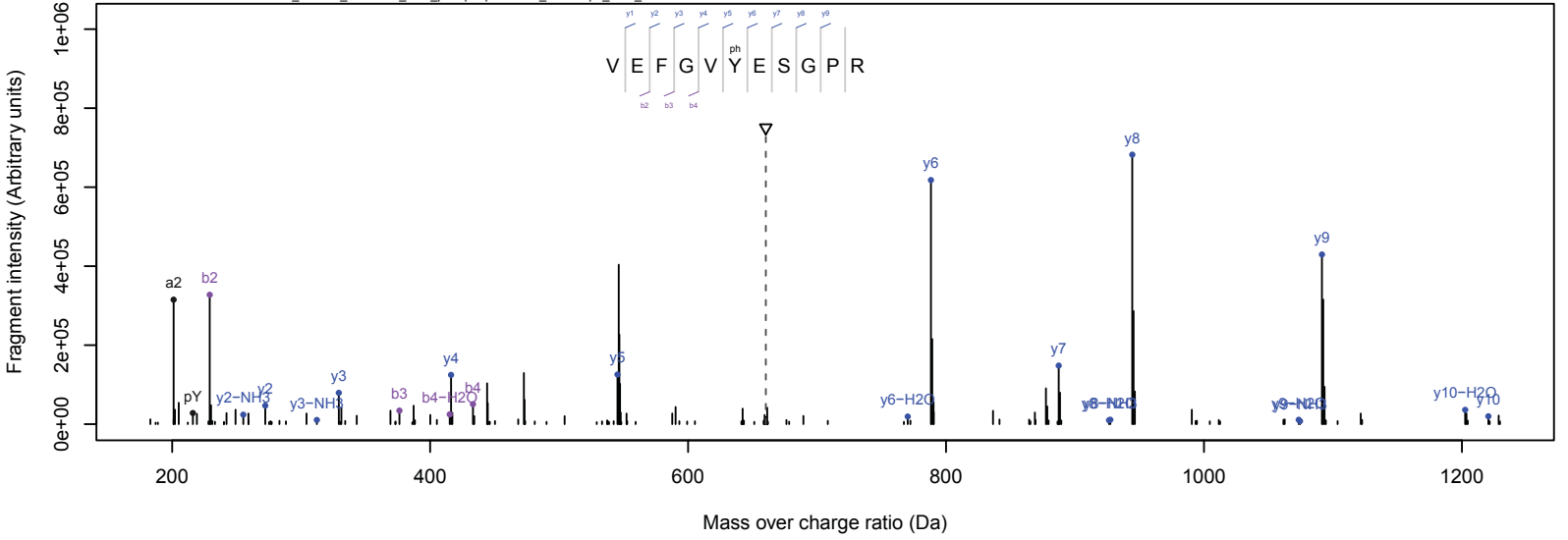
Andromeda score : 108.35



VEFGVY(ph)ESGPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 20875

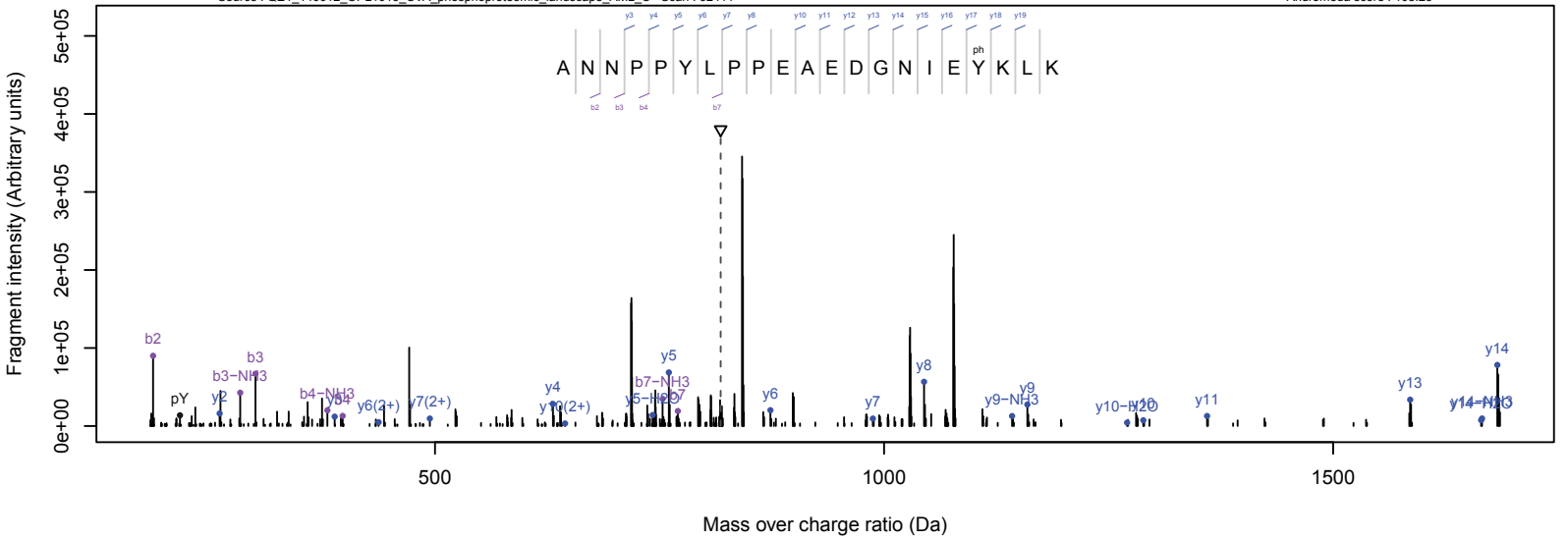
Andromeda score : 108.35



ANNPPYLPPEAEDGNIEY(ph)KLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32411

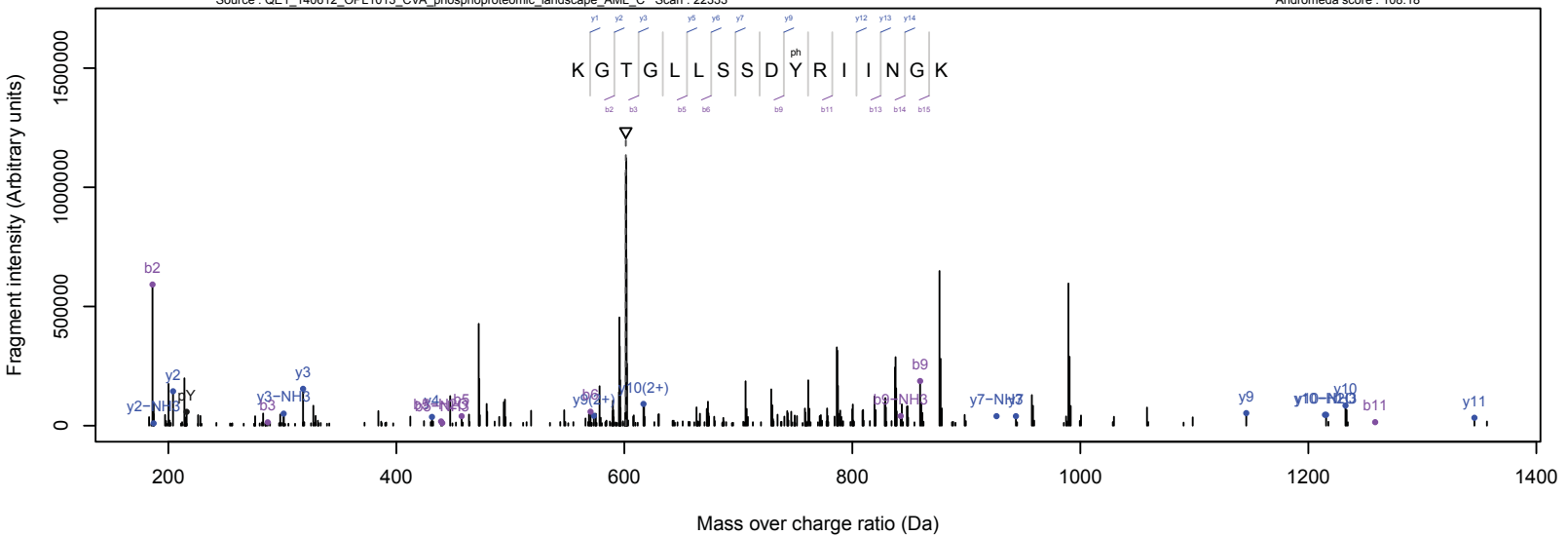
Andromeda score : 108.25



KGTGLSSDYR(ph)IINGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22333

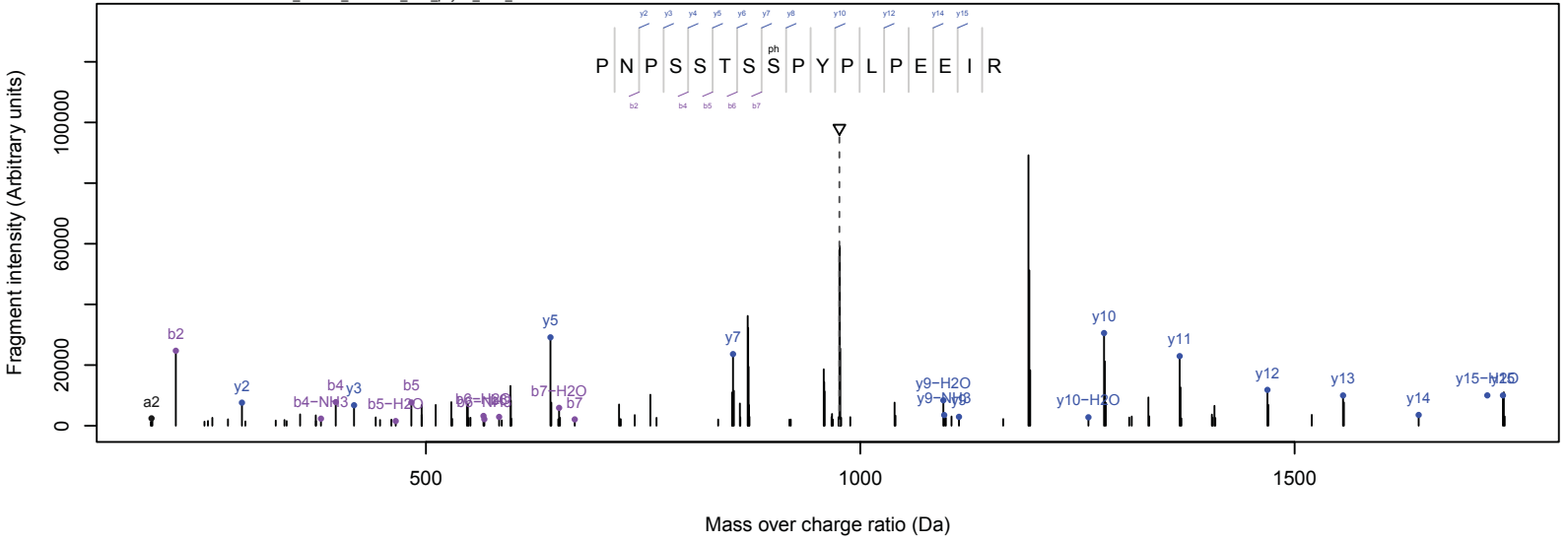
Andromeda score : 108.18



PNPSTSS(ph)PYPLPEEIR-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 26090

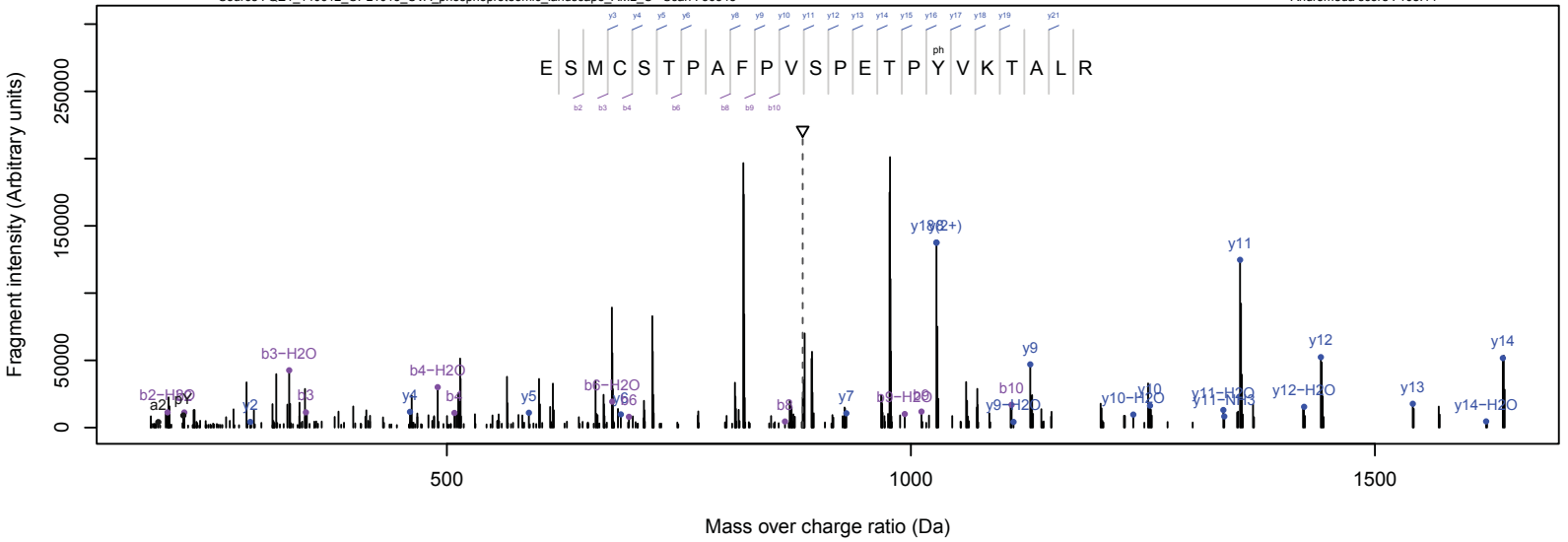
Andromeda score : 108.18



ESMCSTPAFPVSPETPY(ph)VKTALR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36548

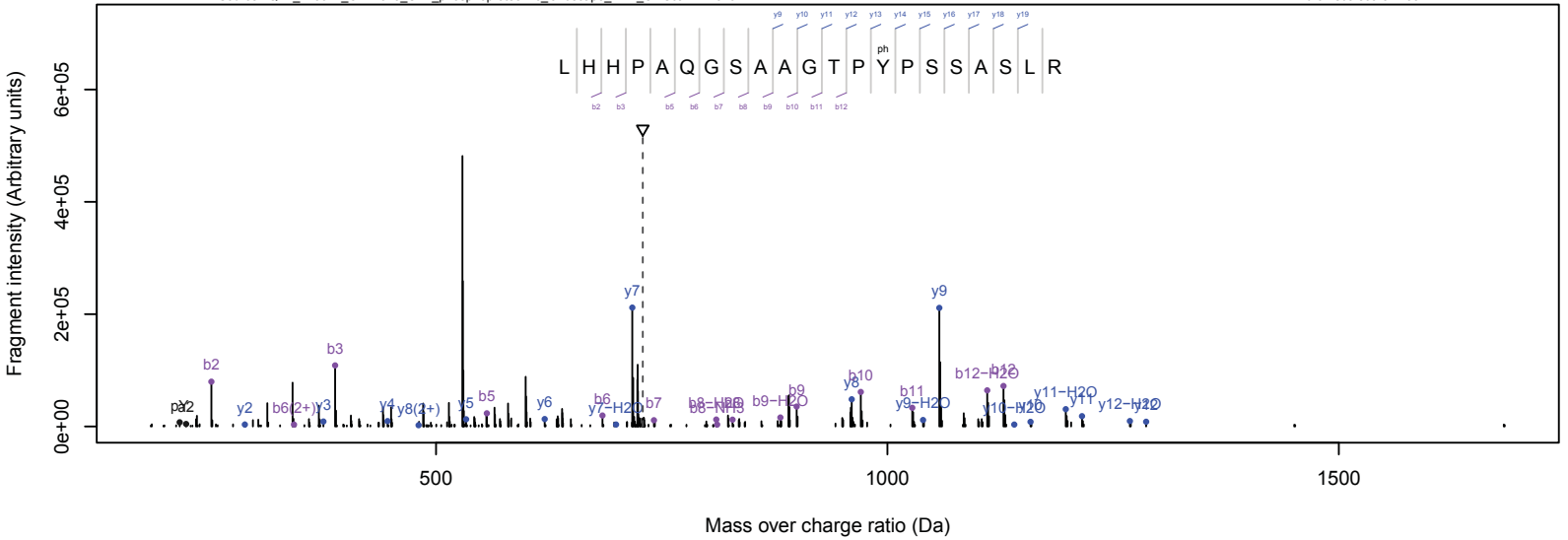
Andromeda score : 108.14



LHHPAQGSAAGTPY(ph)PSSASLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11329

Andromeda score : 108.12

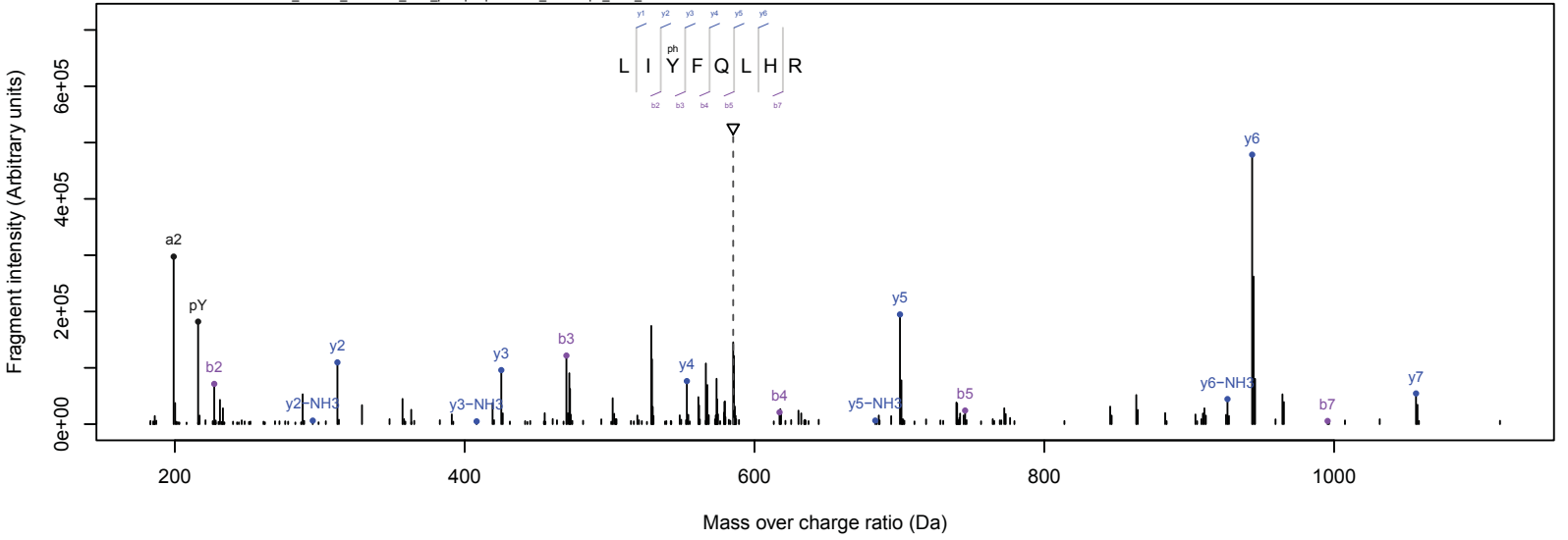




LIY(ph)FQLHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29542

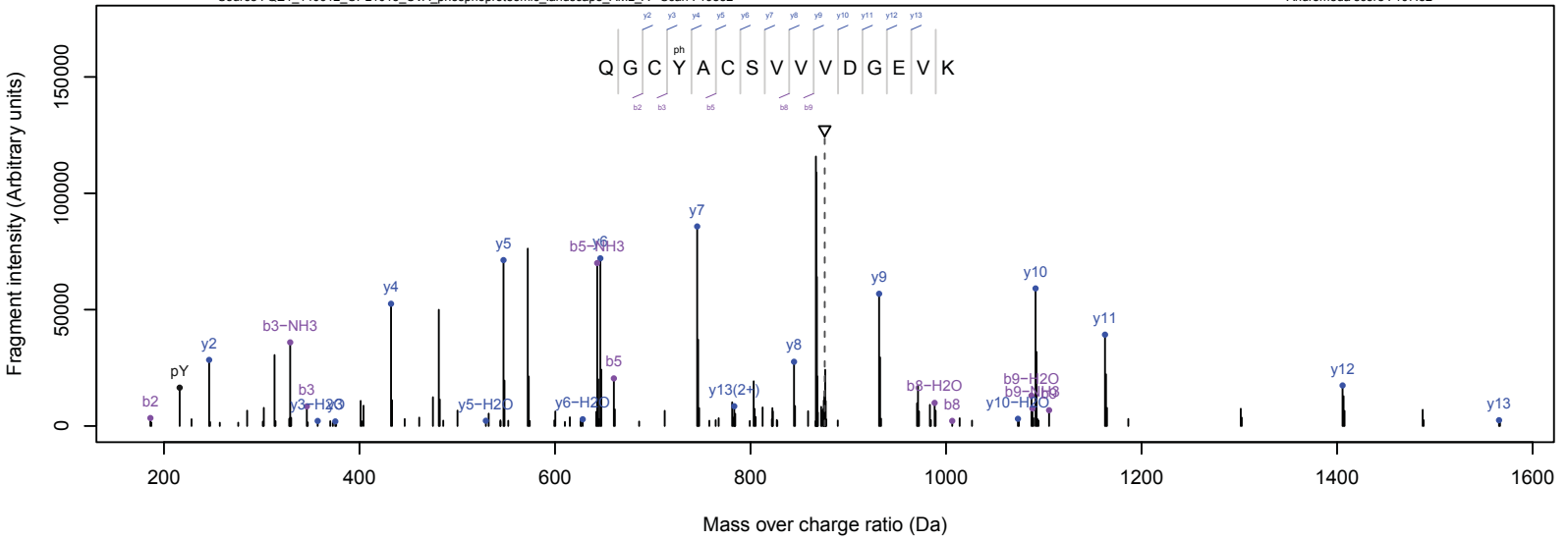
Andromeda score : 108.09



QGCV(ph)ACSVVDGEVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 19882

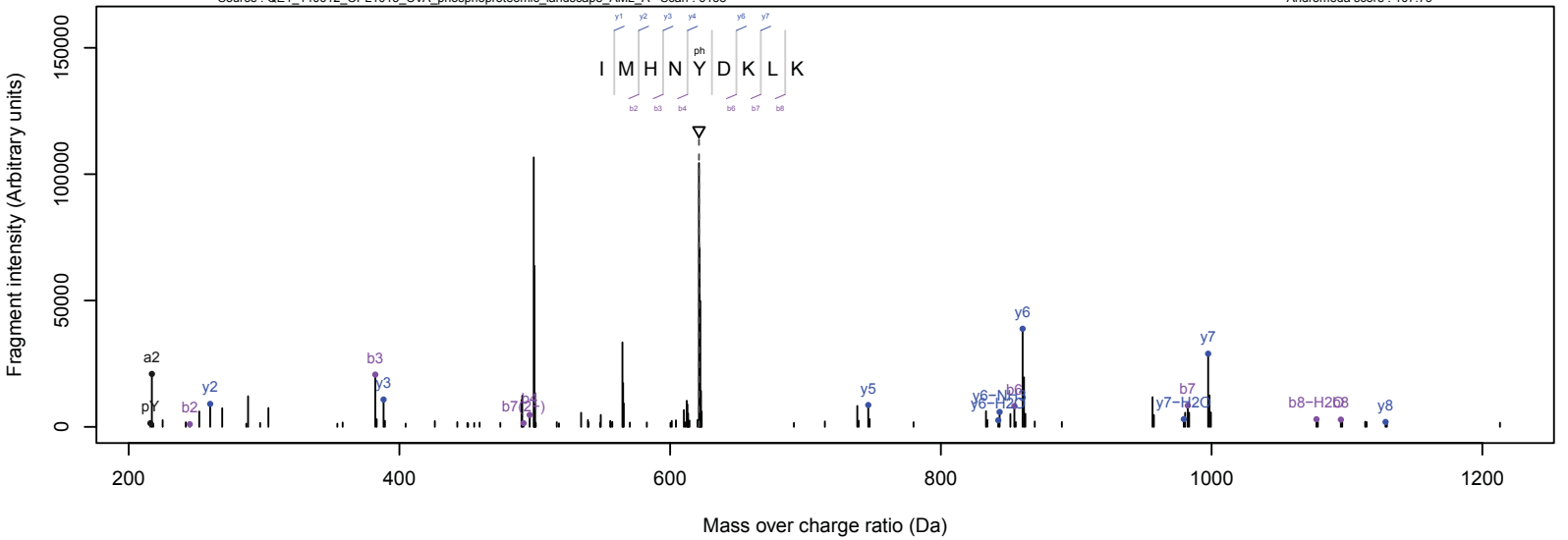
Andromeda score : 107.82



IMHNY(ph)DKLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 6158

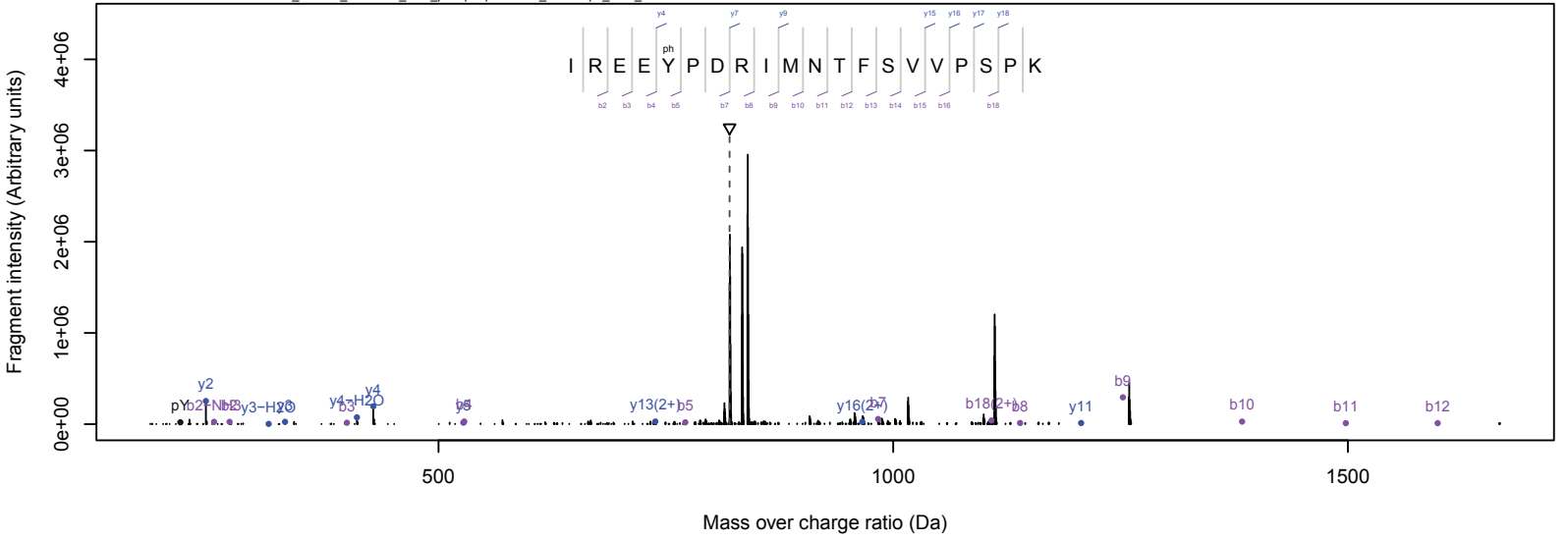
Andromeda score : 107.79



IREEY(ph)PDRIMNTFSVVPSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34941

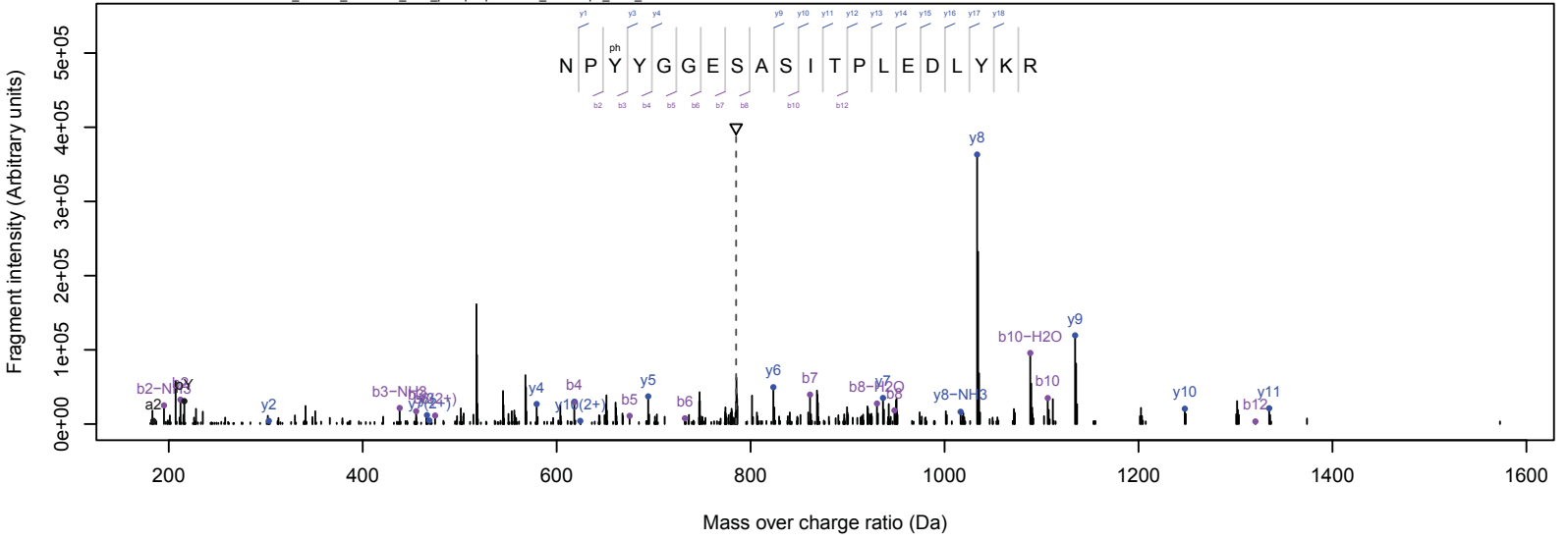
Andromeda score: 107.78



NPY(ph)YGGESASITPLEDLYKR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 37803

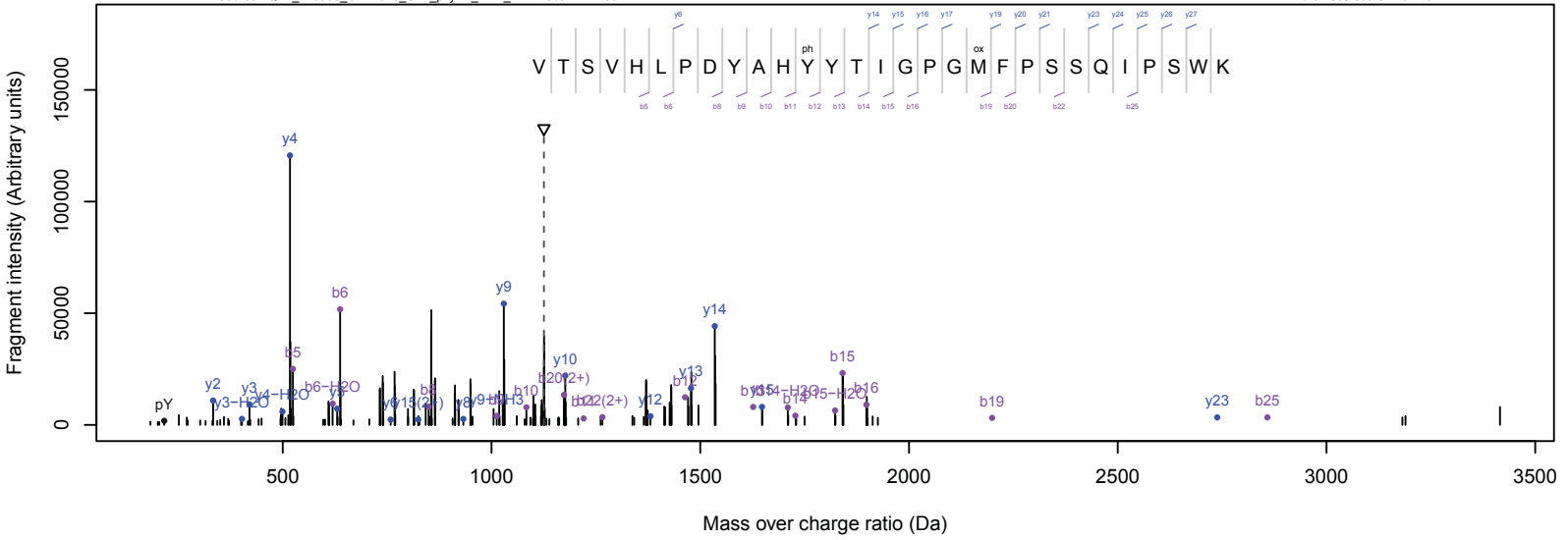
Andromeda score: 107.78



VTSVHLPDYAHY(ph)YTIGPM(ox)FPSSQIPSWK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 47180

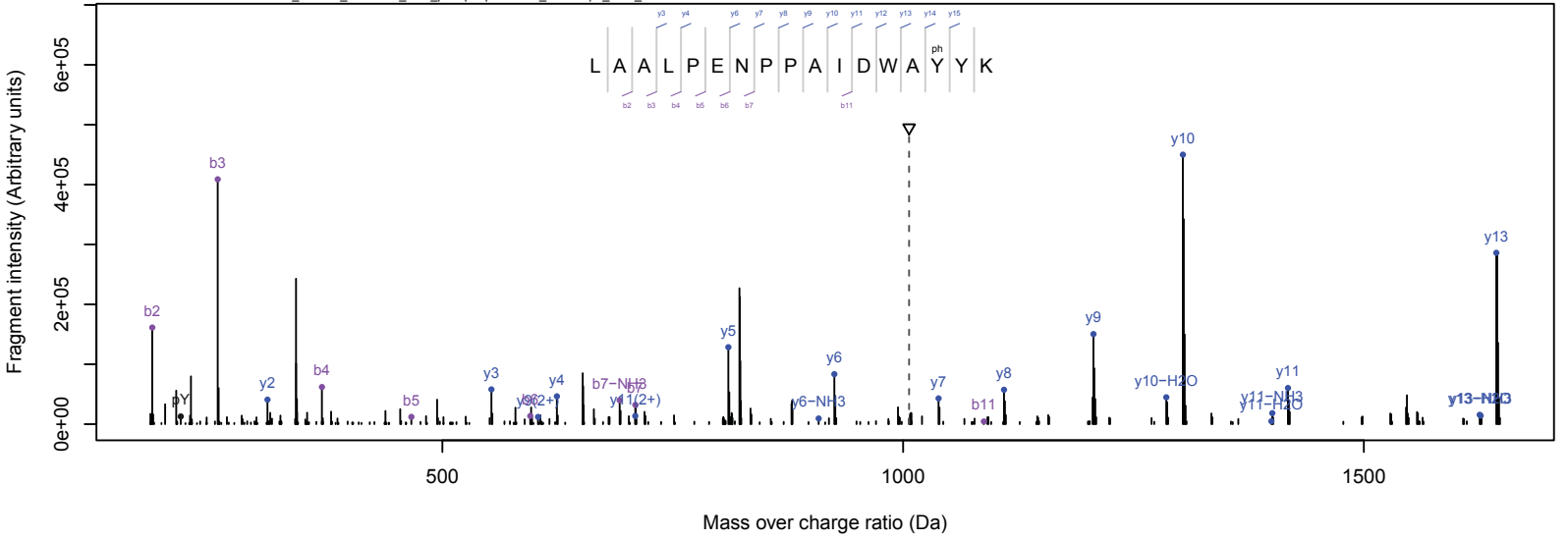
Andromeda score: 107.75



LAALPENPPAIDWAY(ph)YK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44289

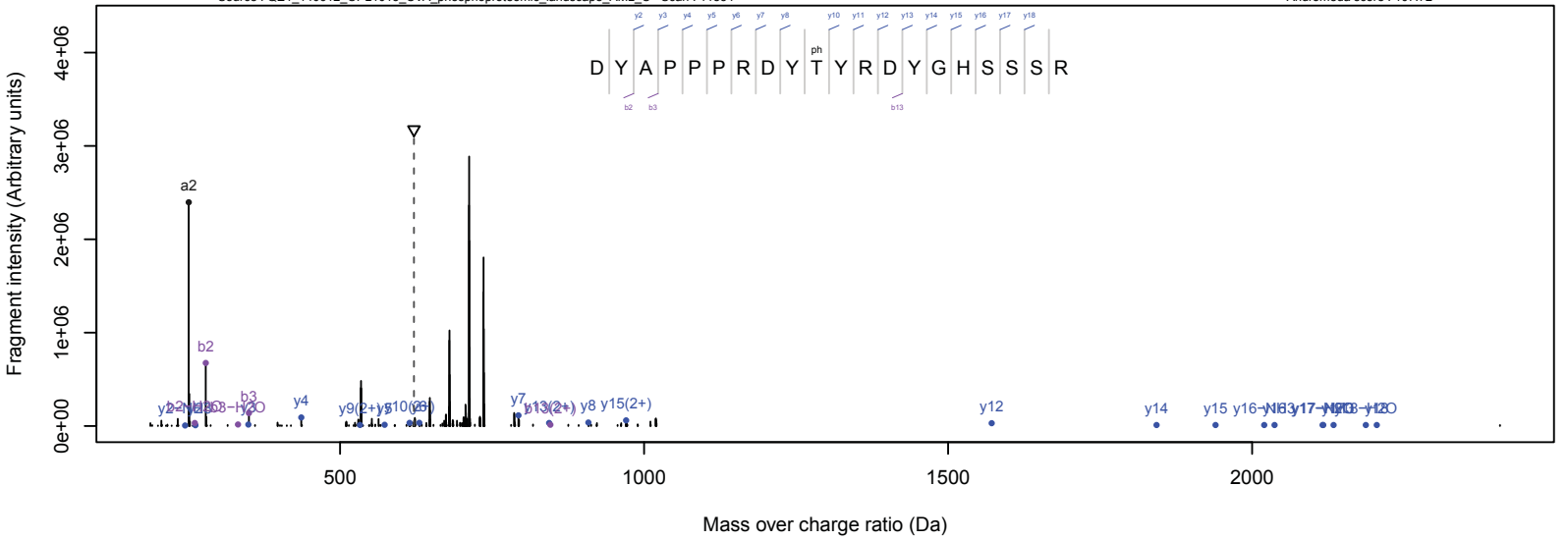
Andromeda score : 107.73



DYAPPPRDYT(ph)YRDYGHSSSR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11854

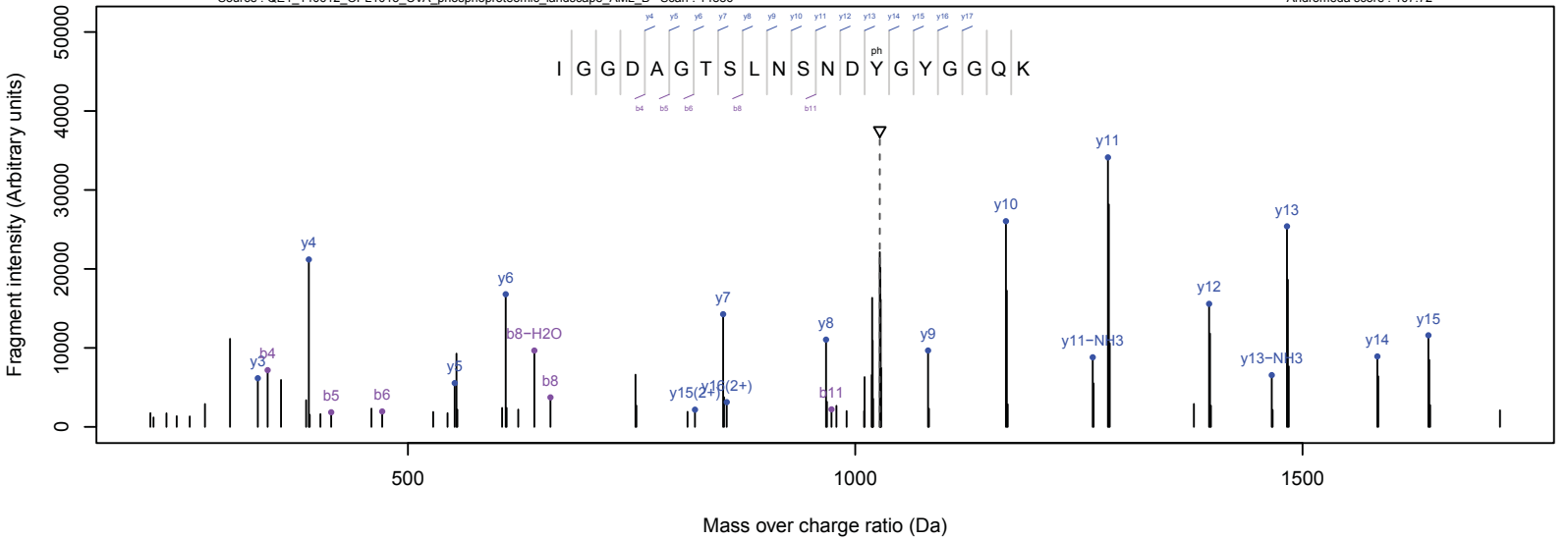
Andromeda score : 107.72



IGGDAGTSLNSNDY(ph)YGGQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14886

Andromeda score : 107.72

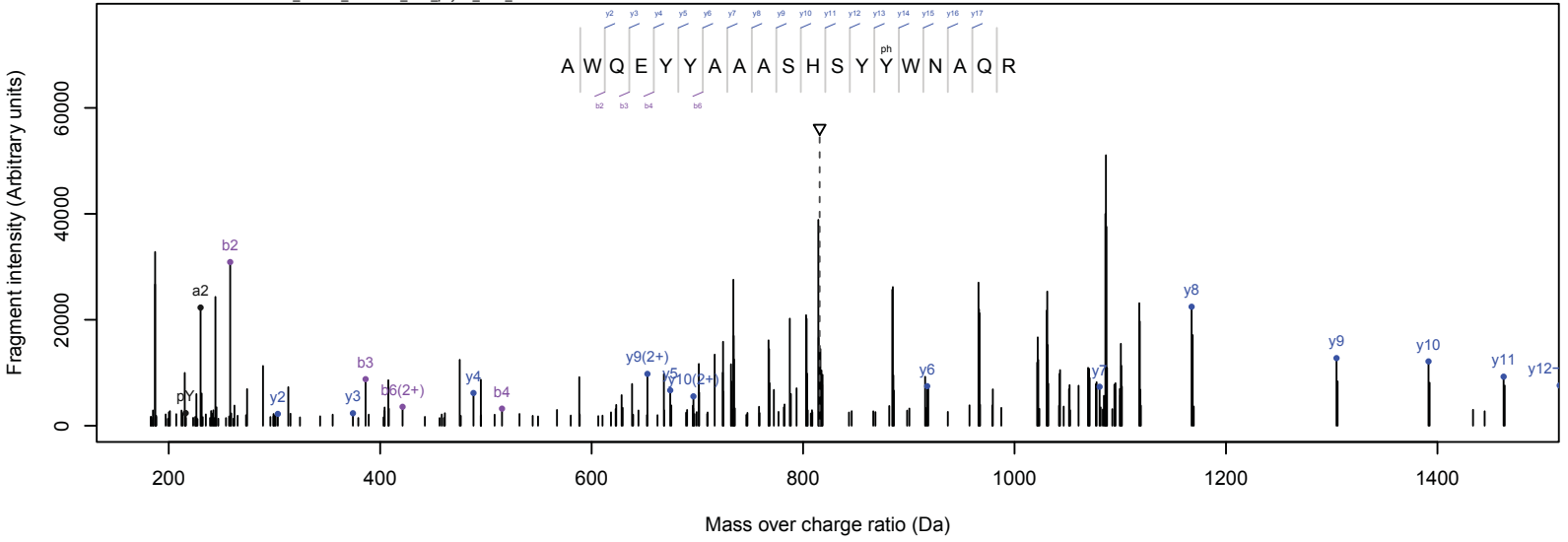




**AWQEYAAASHSY(ph)WNAQR-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 36741

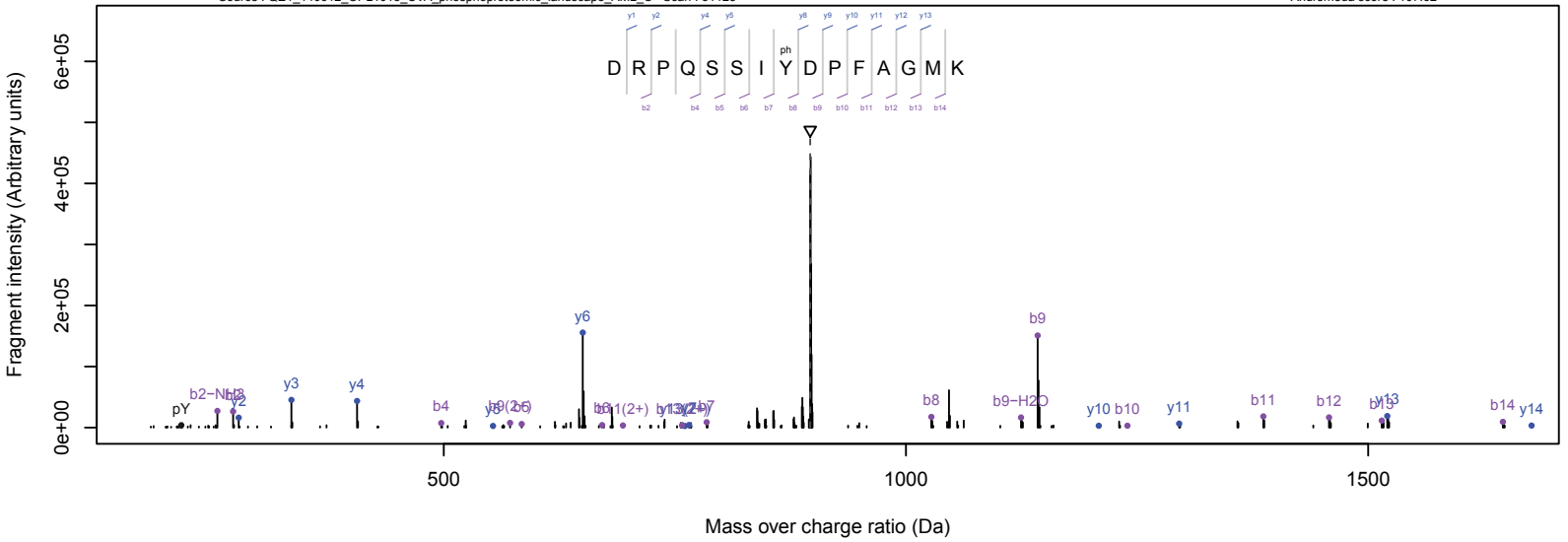
Andromeda score : 107.65



**DRPQSSIIY(ph)DPFAGMK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31125

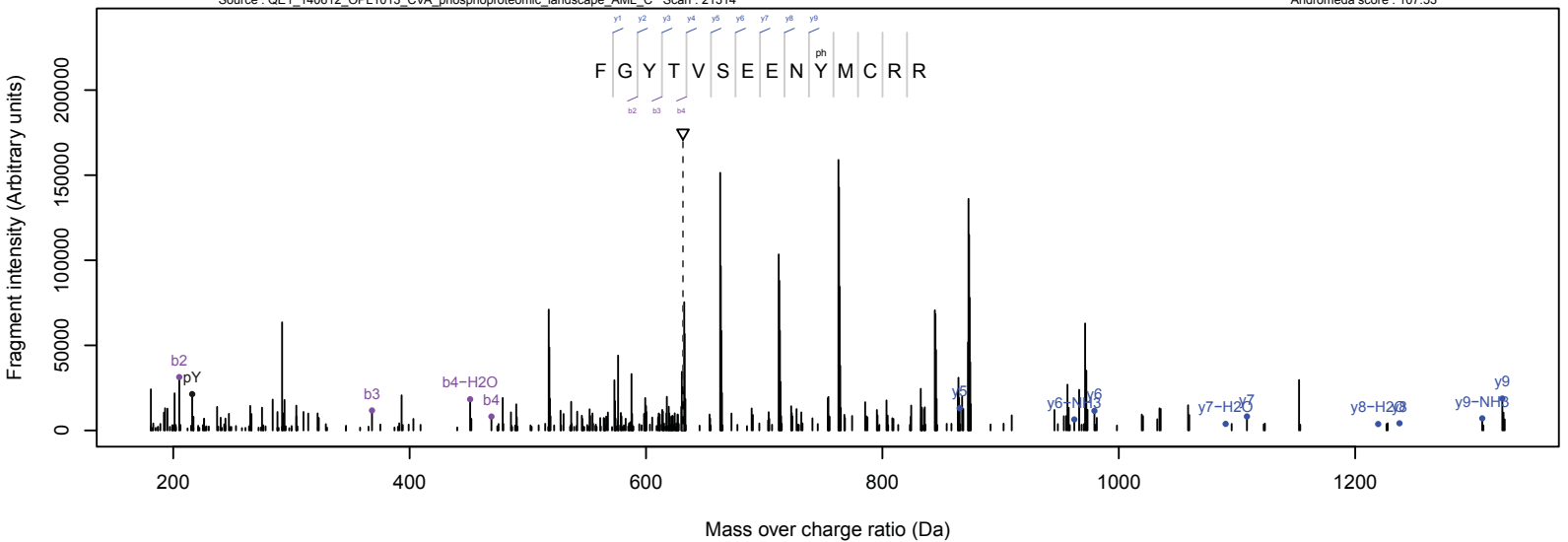
Andromeda score : 107.62



**FGYTVSEENY(ph)MCR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21314

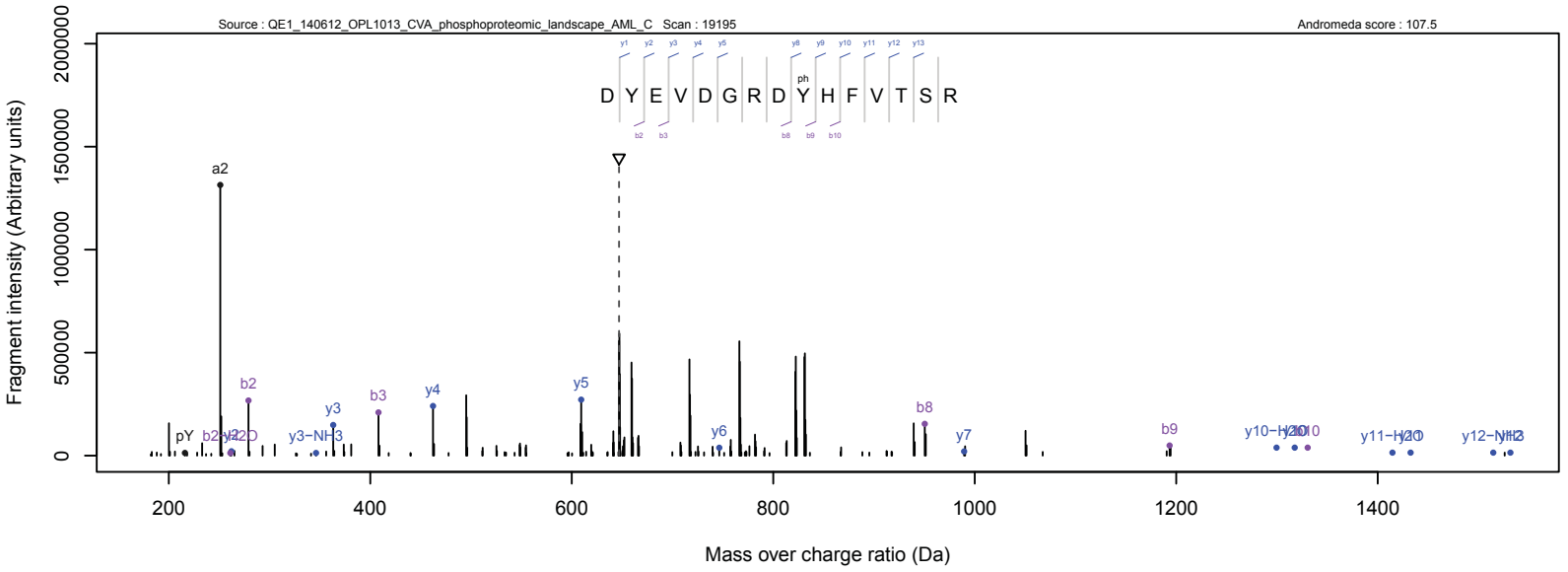
Andromeda score : 107.53



DYEVDGRDY(ph)HFVTSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19195

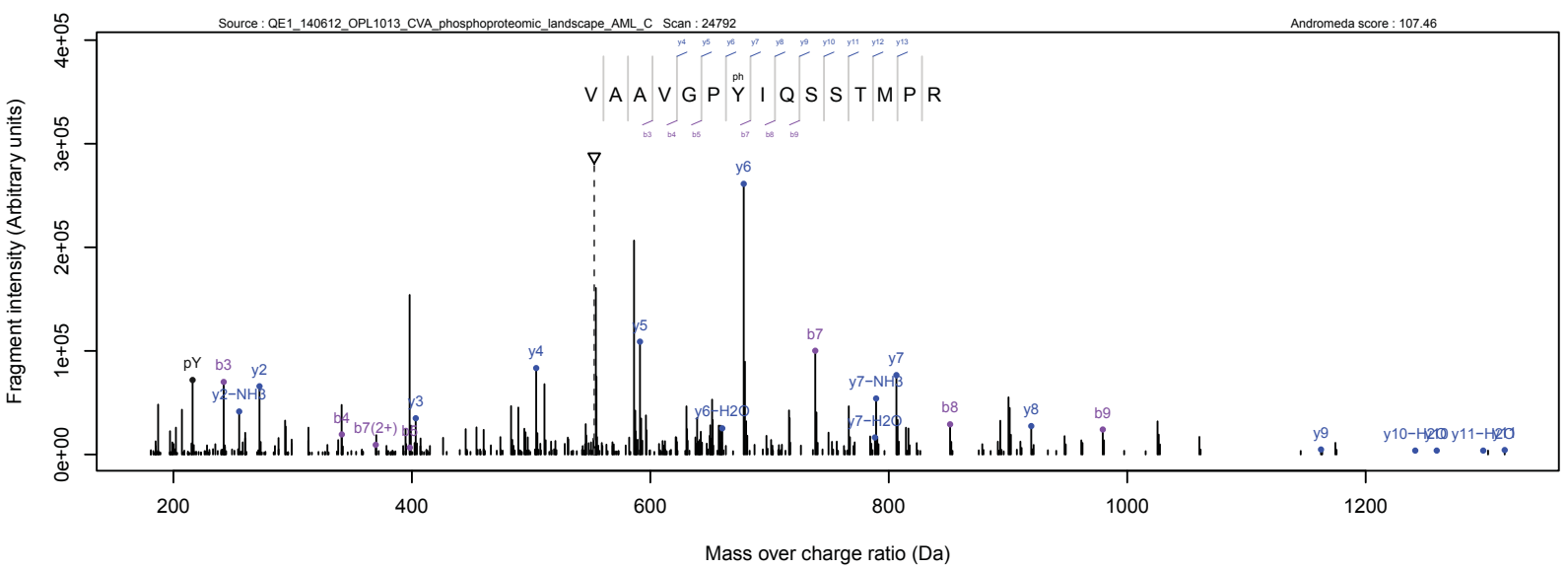
Andromeda score : 107.5



VAAVGPHYIQSSTMPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24792

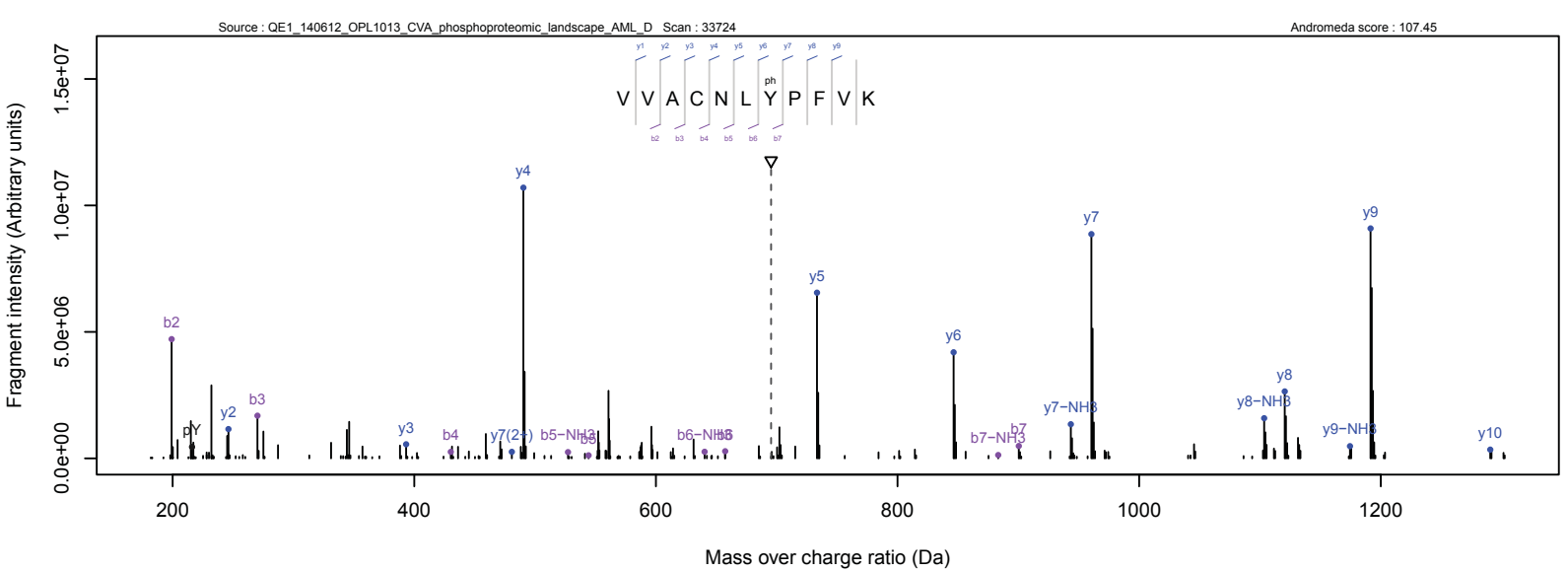
Andromeda score : 107.46



VVACNLYP(V)FVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33724

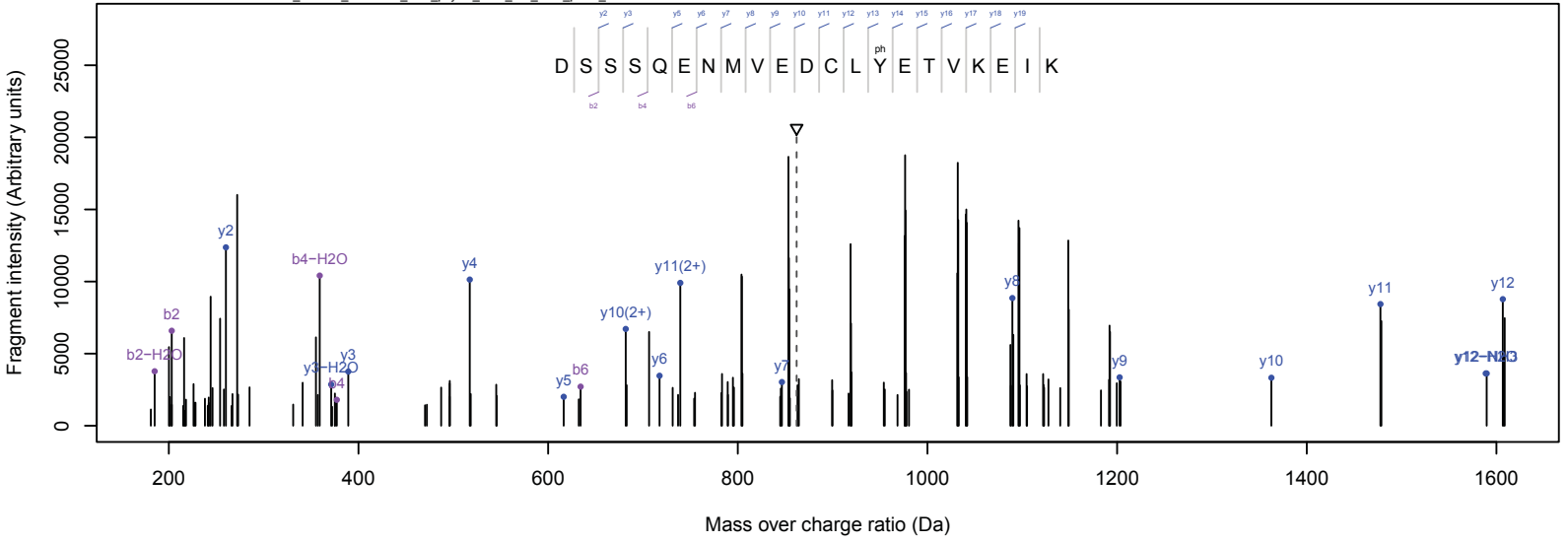
Andromeda score : 107.45



DSSSQENMVEDCLY(ph)ETVKEIK-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 39790

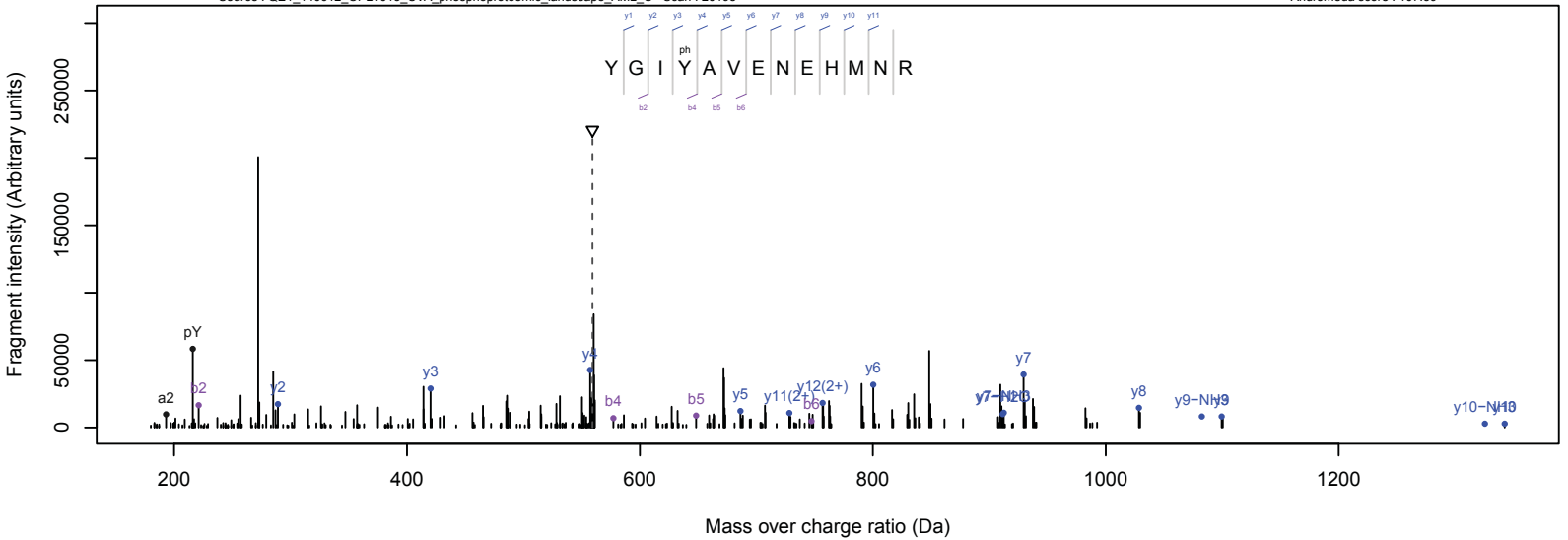
Andromeda score : 107.39



YGIY(ph)AVENEHMNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20198

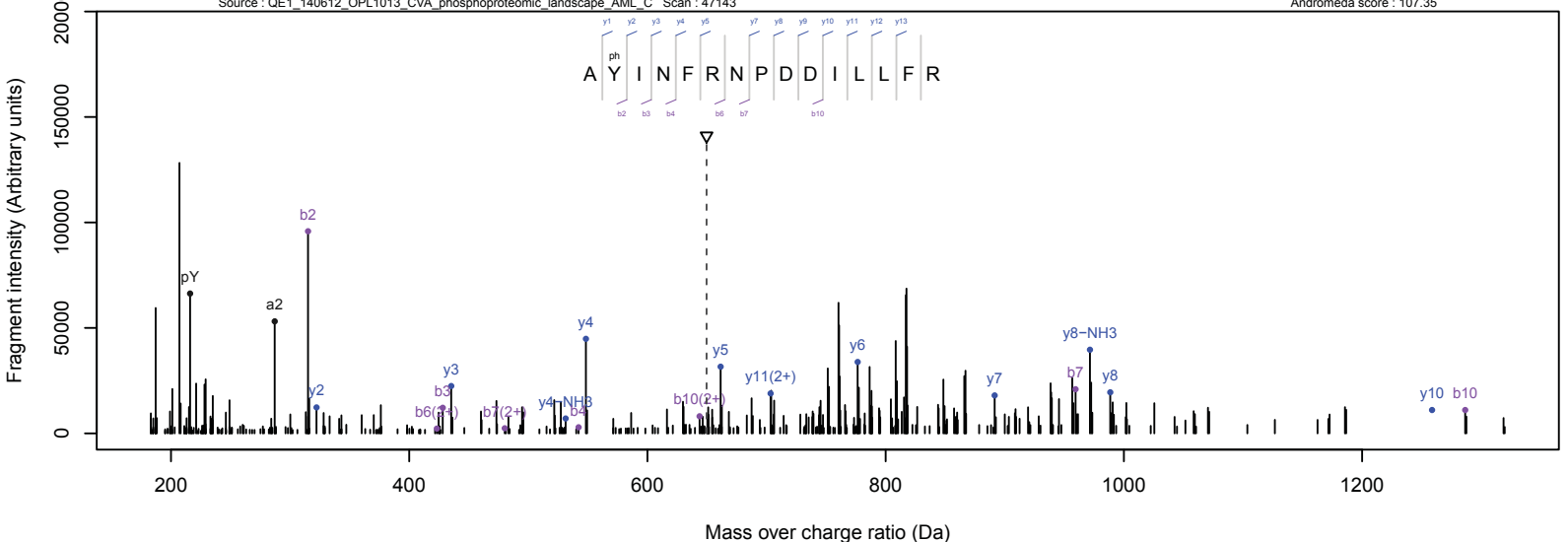
Andromeda score : 107.39



AY(ph)INFRNPDDILLFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47143

Andromeda score : 107.35



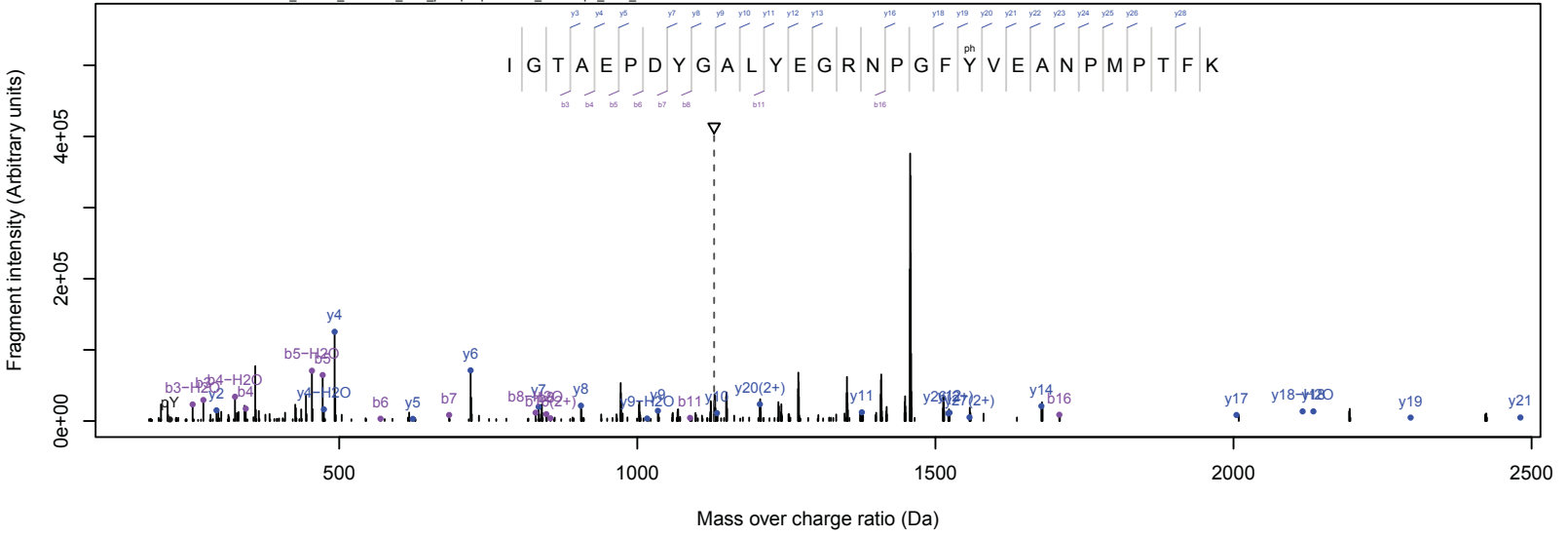




IGTAEPDYGALYEGRNPGFY(ph)VEANPMPFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44431

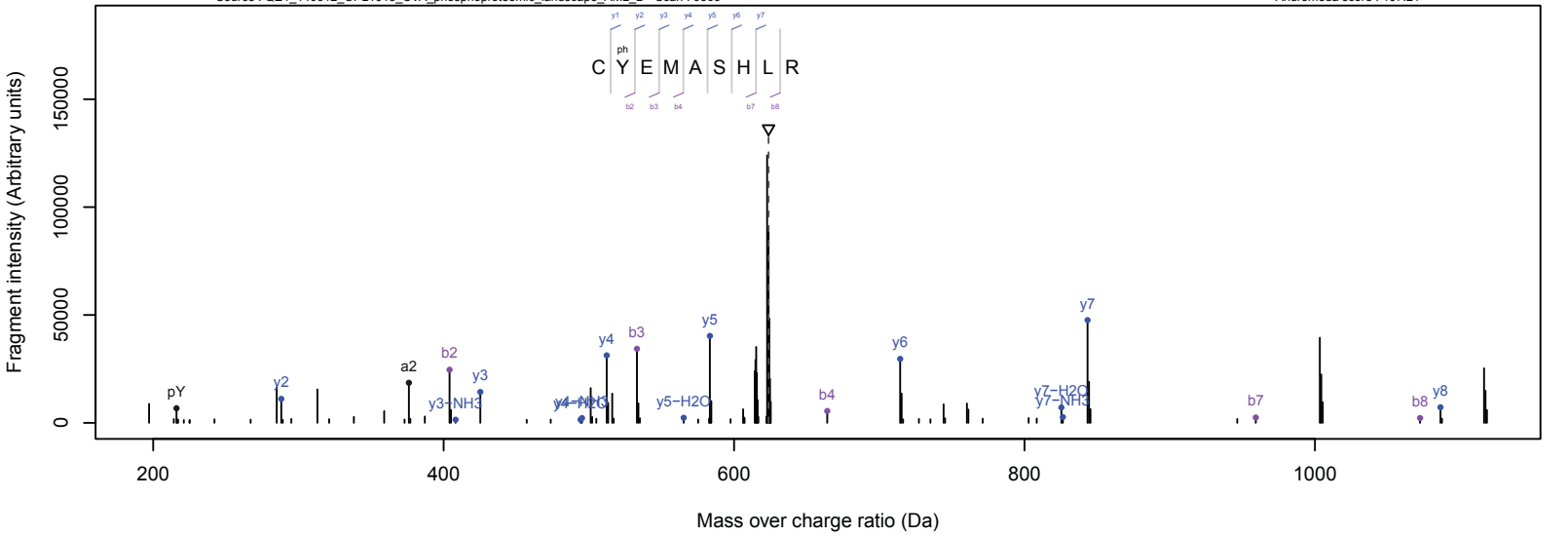
Andromeda score : 107.3



CY(ph)EMASHLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9363

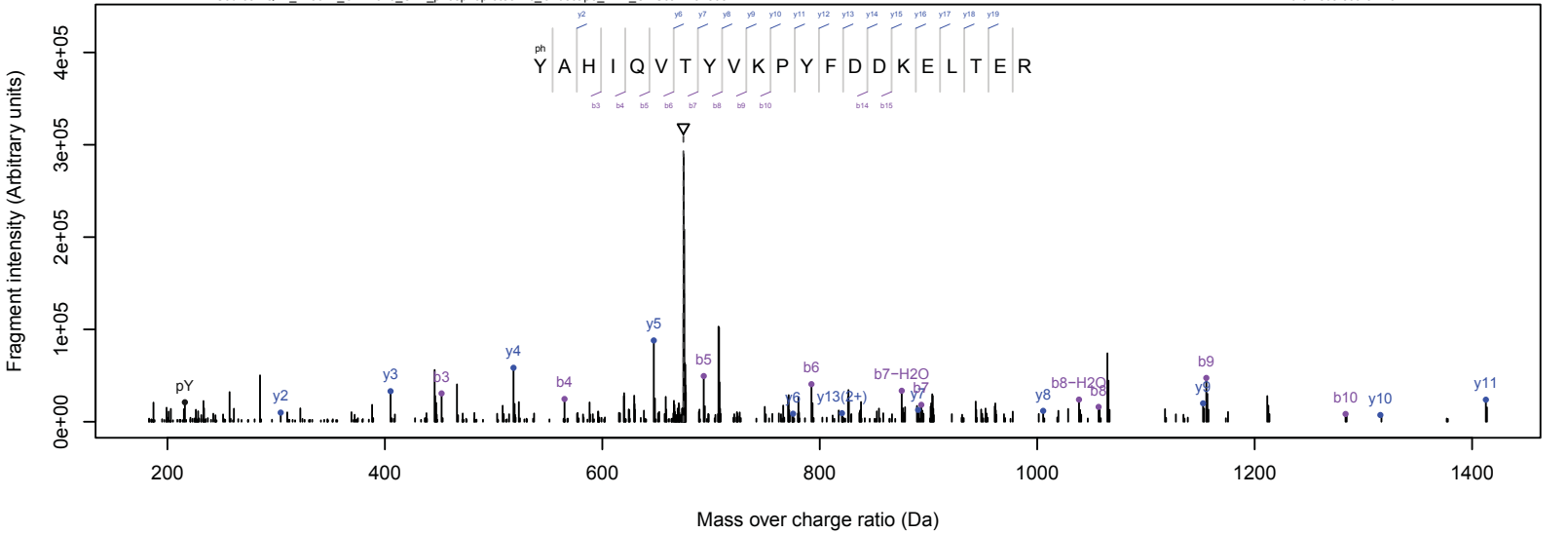
Andromeda score : 107.21



Y(ph)AHIQVTYVKPYFDDKELTER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31808

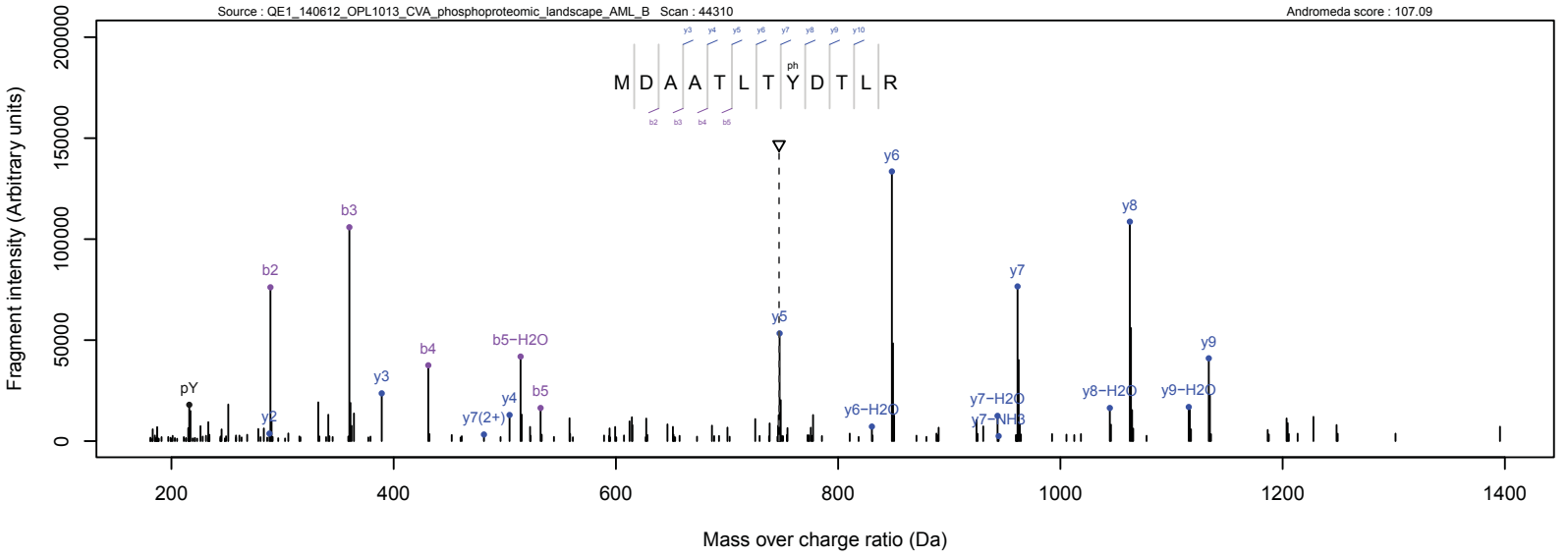
Andromeda score : 107.21



(ac)MDAATLTY(ph)DTLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 44310

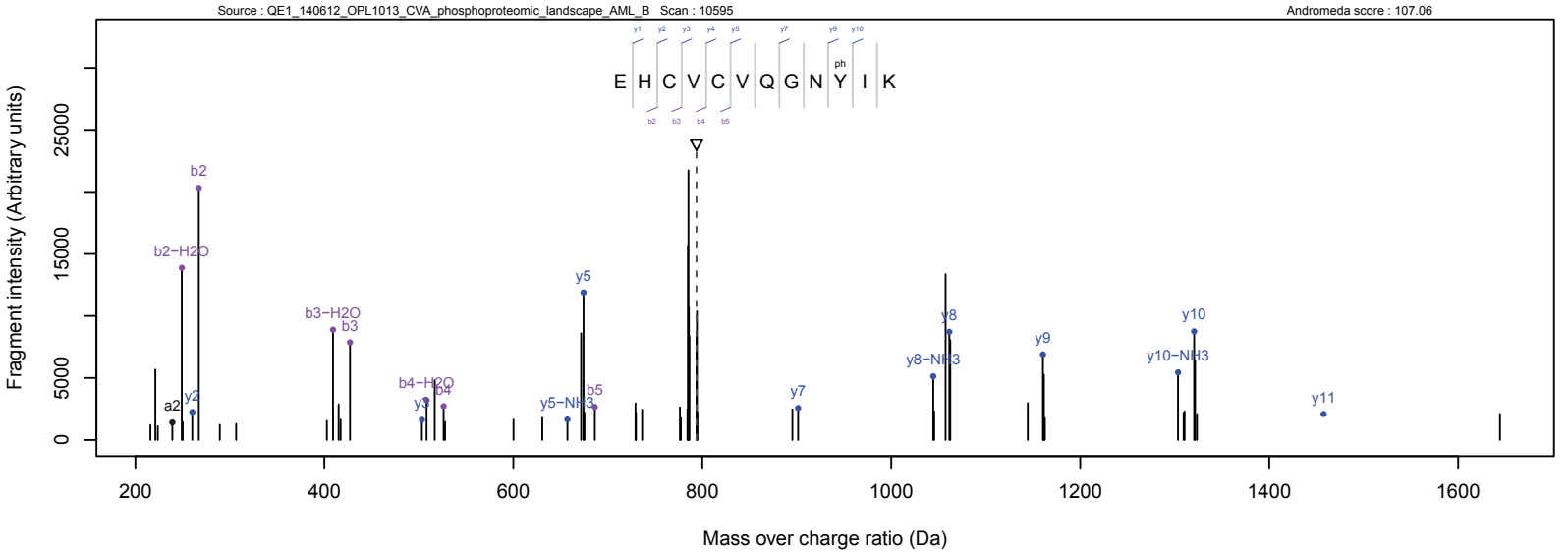
Andromeda score: 107.09



EHCVCVQNY(ph)IK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 10595

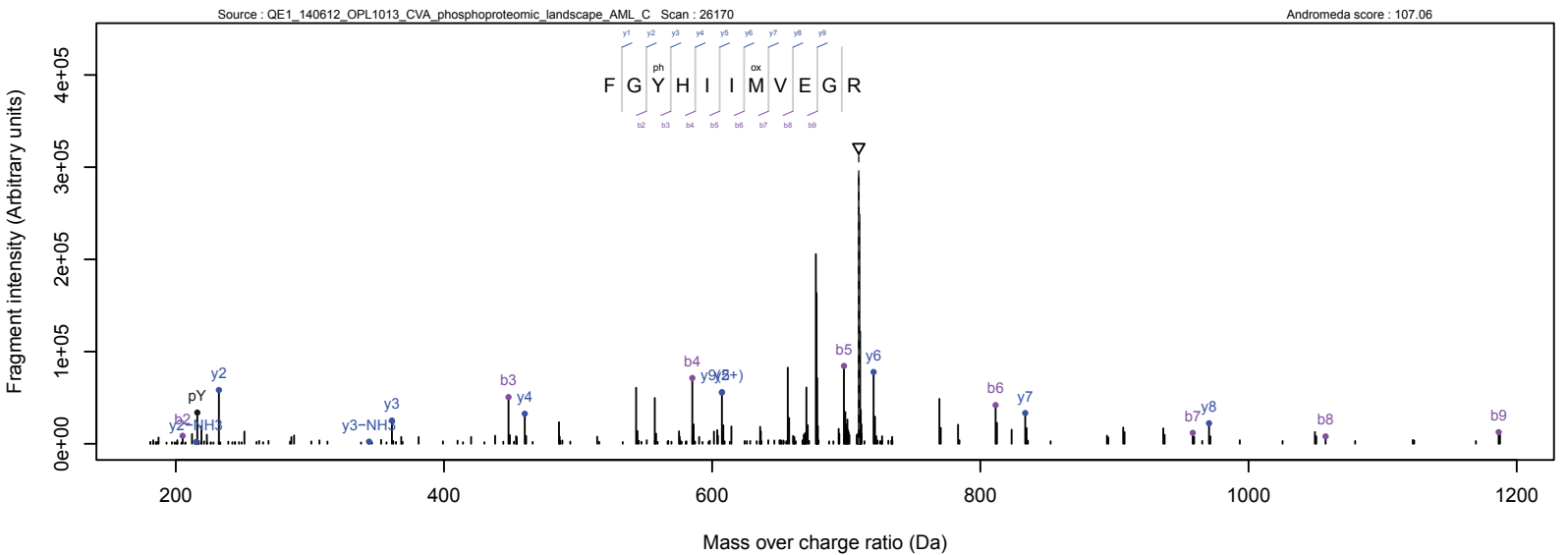
Andromeda score: 107.06



FGYHIIIMV(ox)EGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26170

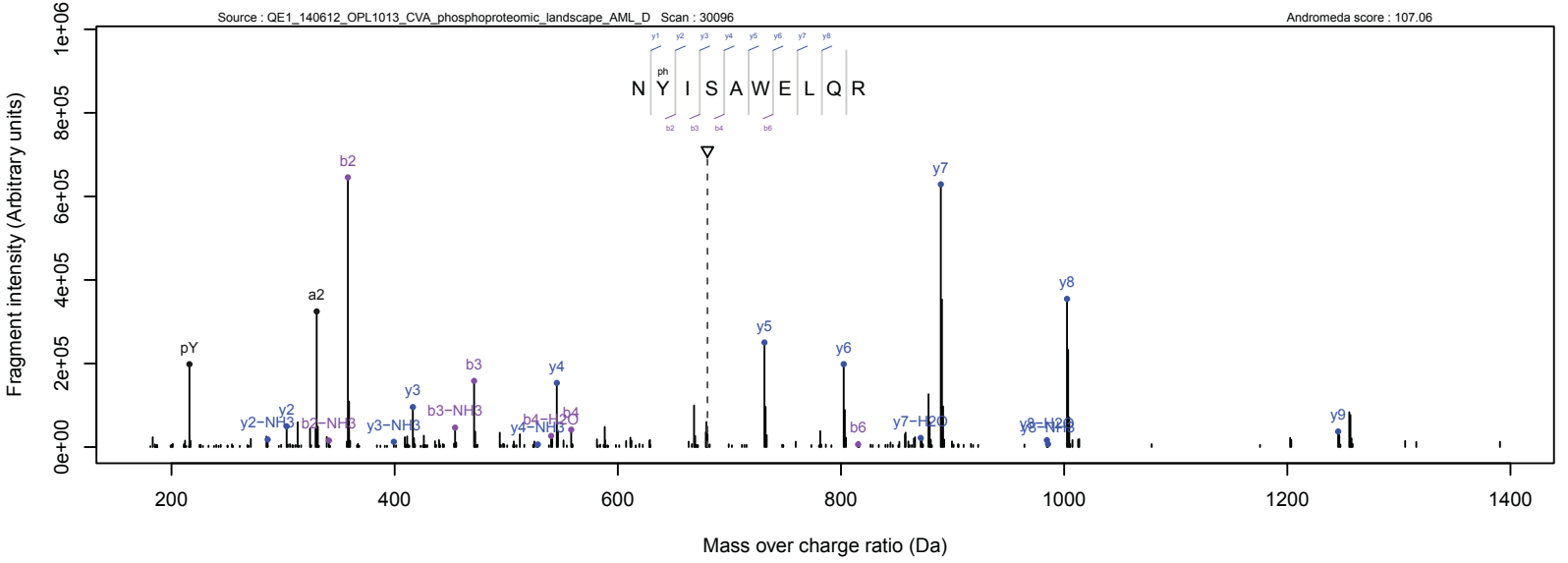
Andromeda score: 107.06



NY(ph)ISAWELQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 30096

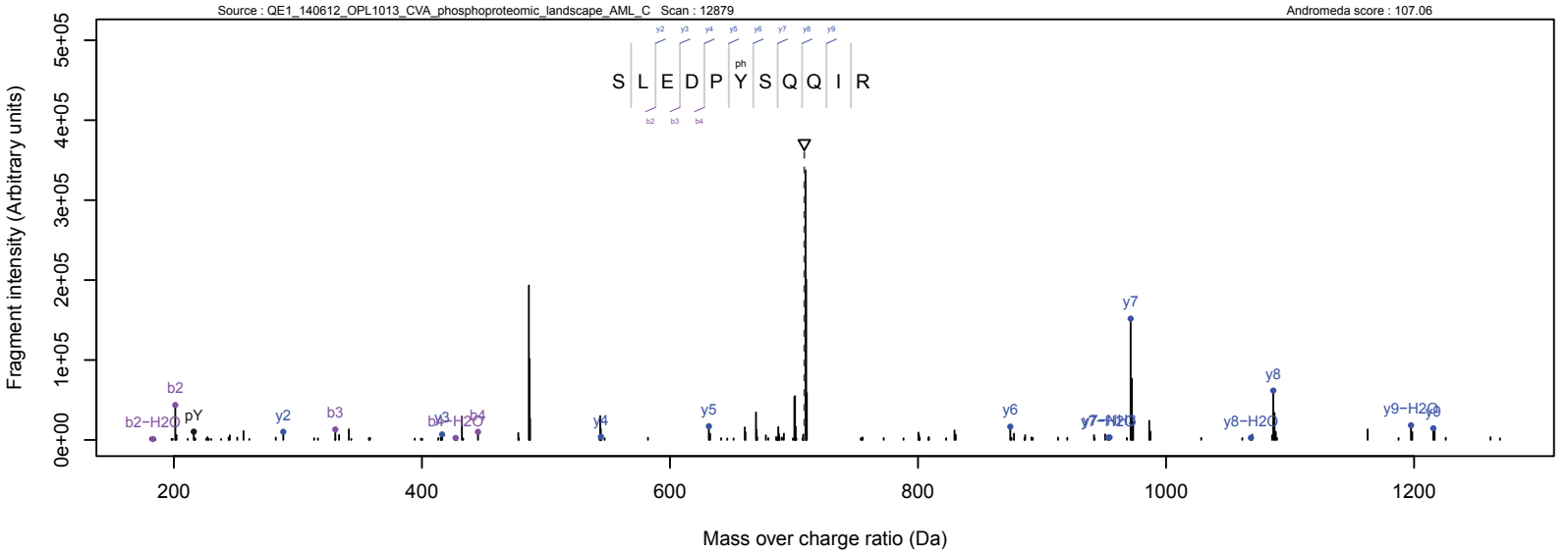
Andromeda score : 107.06



SLEDPY(ph)SQQIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12879

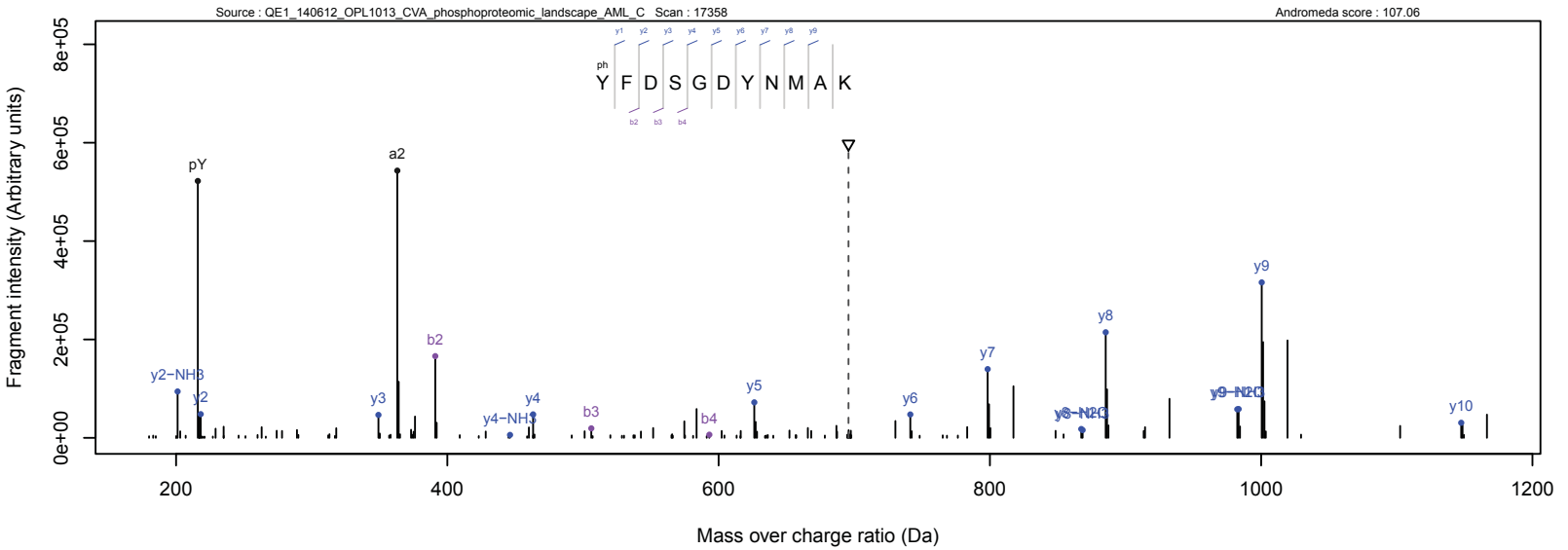
Andromeda score : 107.06



Y(ph)FDSGDYNAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17358

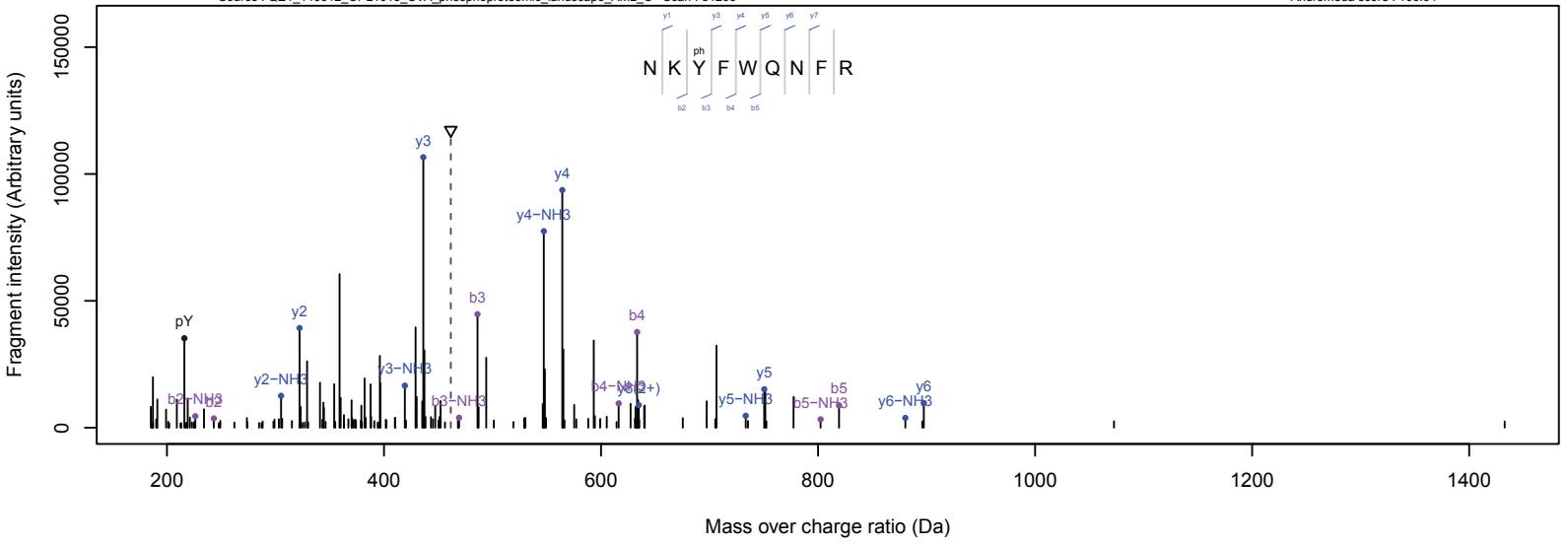
Andromeda score : 107.06



NKY(ph)FWQNFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31286

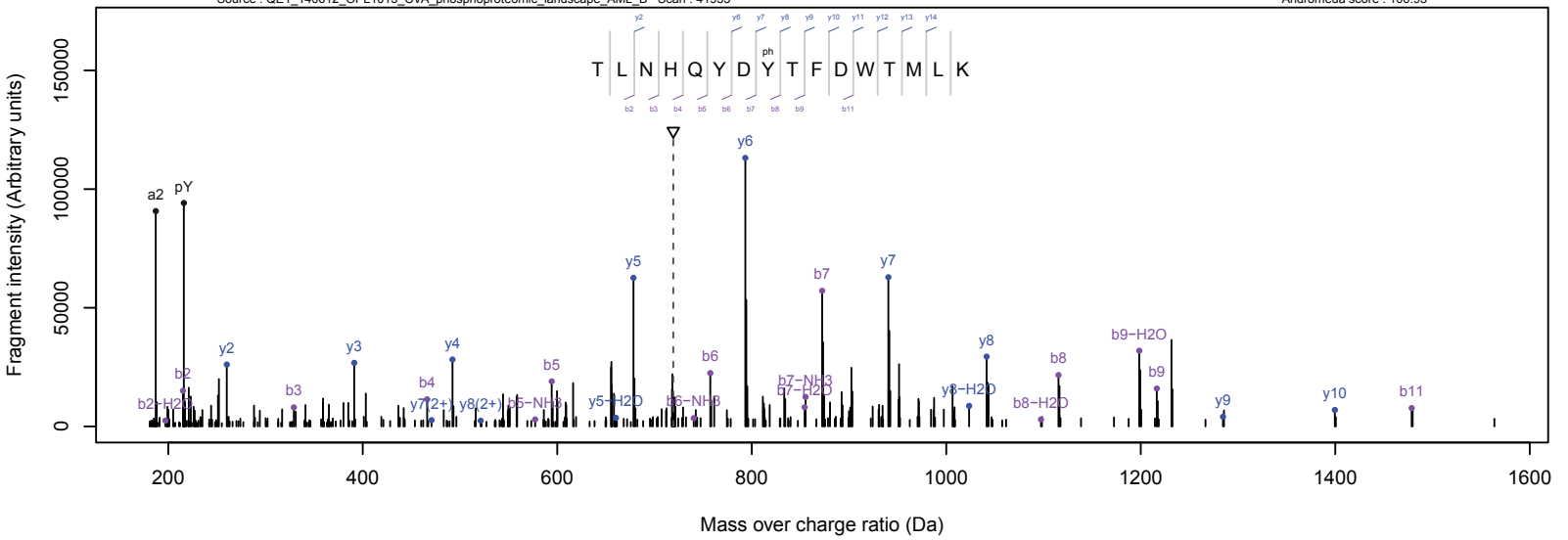
Andromeda score : 106.94



TLNHQYD(ph)TFDWTMLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41955

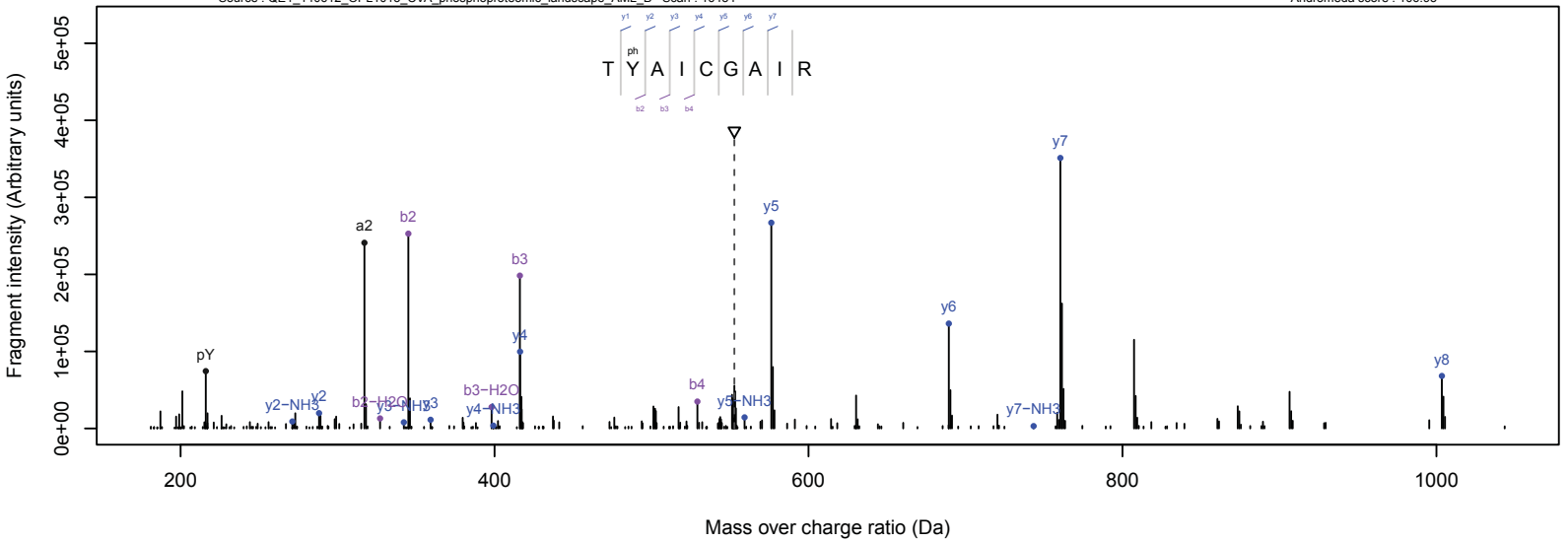
Andromeda score : 106.93



TY(ph)AICGAIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16434

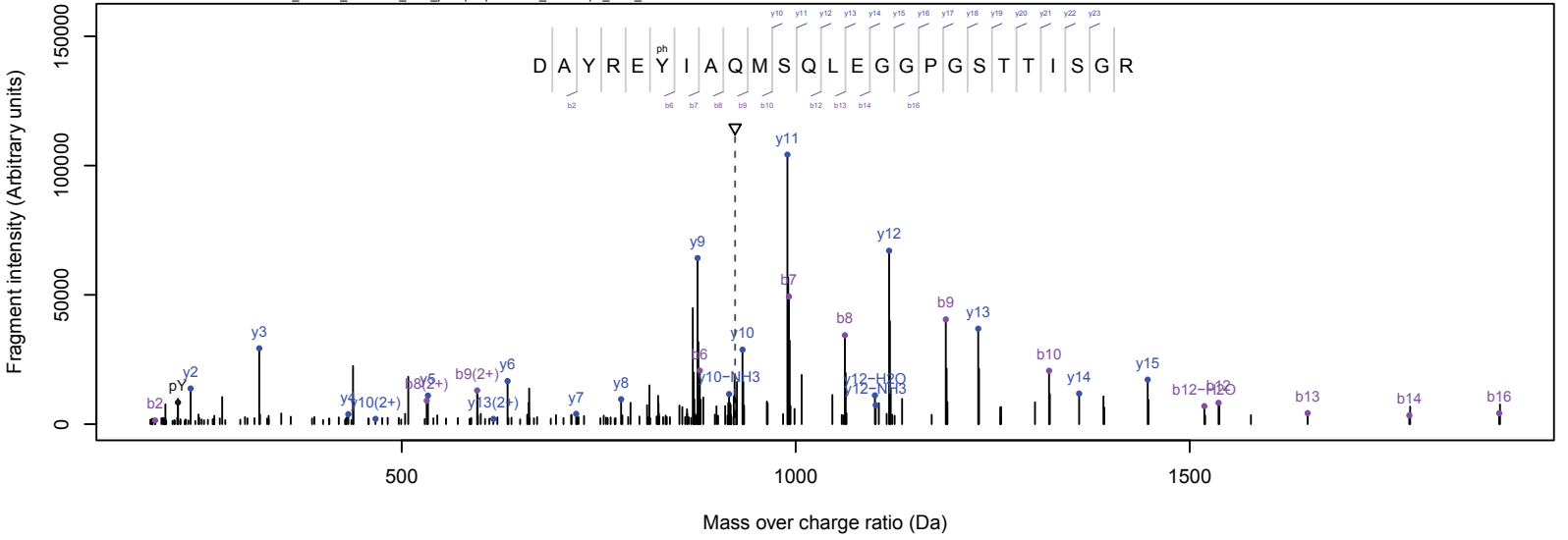
Andromeda score : 106.93



DAYREY(ph)IAQMSQLEGGPGSTTISGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37882

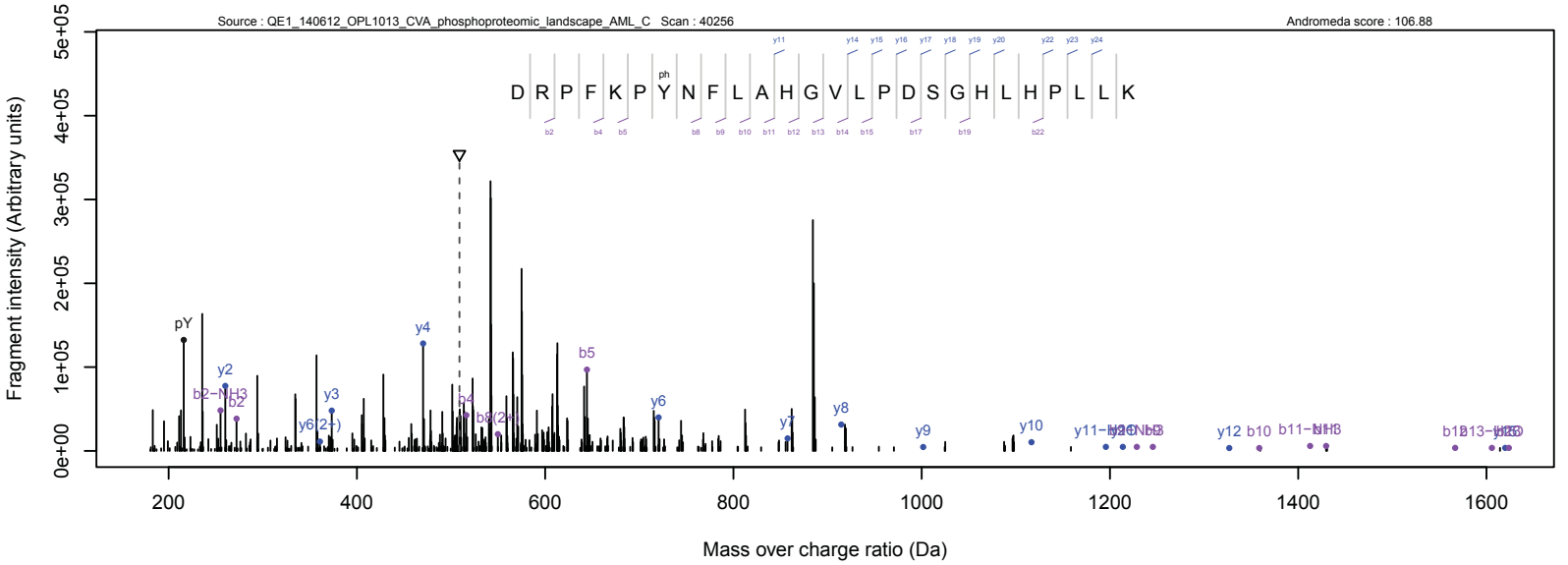
Andromeda score: 106.88



DRPFKPY(ph)NFLAHGVLPDSGHLHPLLK-6+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 40256

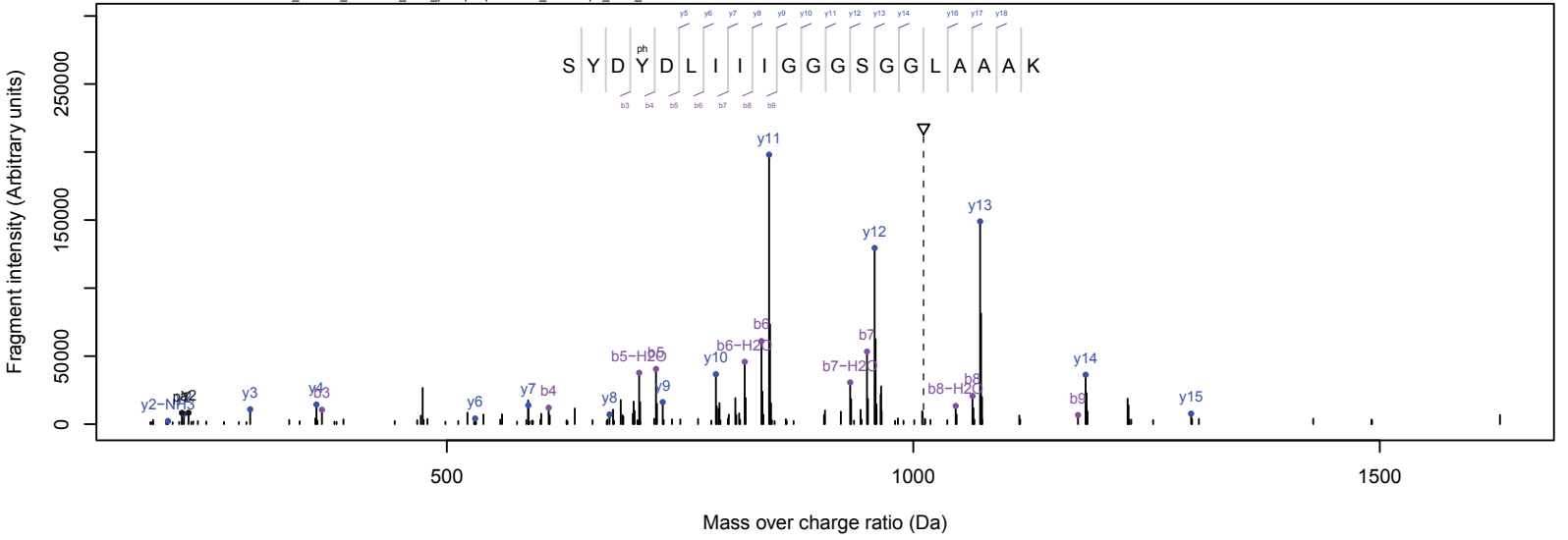
Andromeda score: 106.88



**SYDY(ph)DLIIIGGGGGLAAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43980

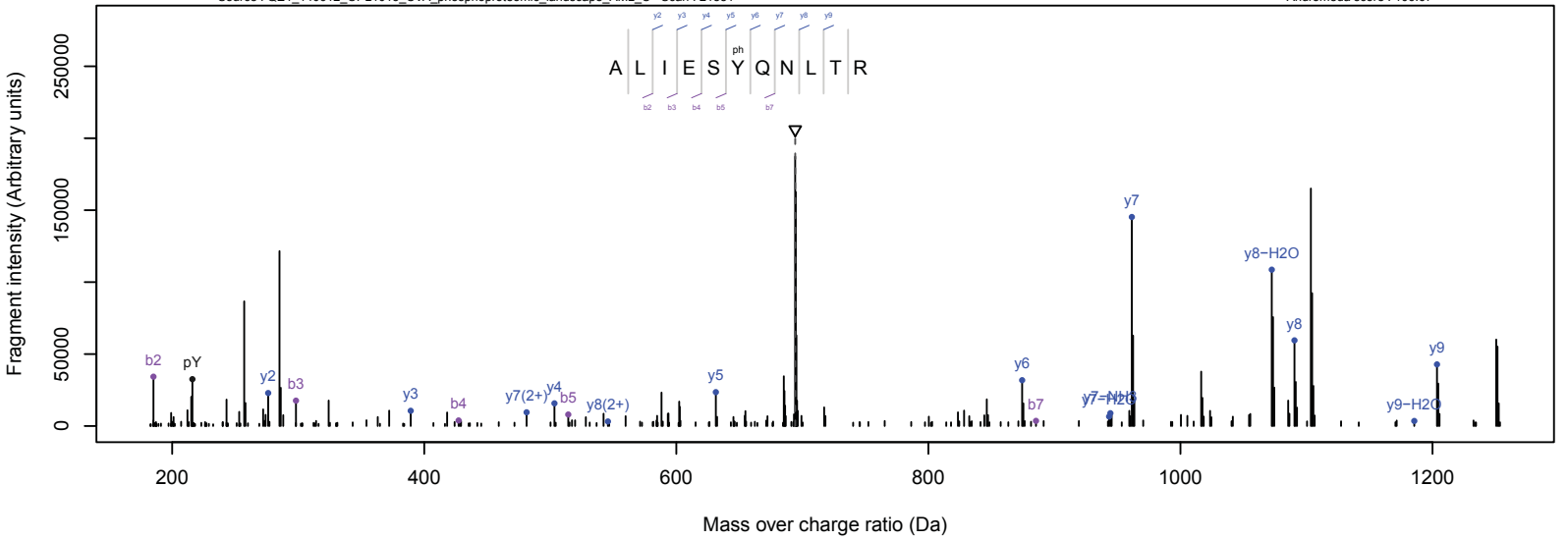
Andromeda score : 106.75



**ALIESY(ph)QLNTR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21881

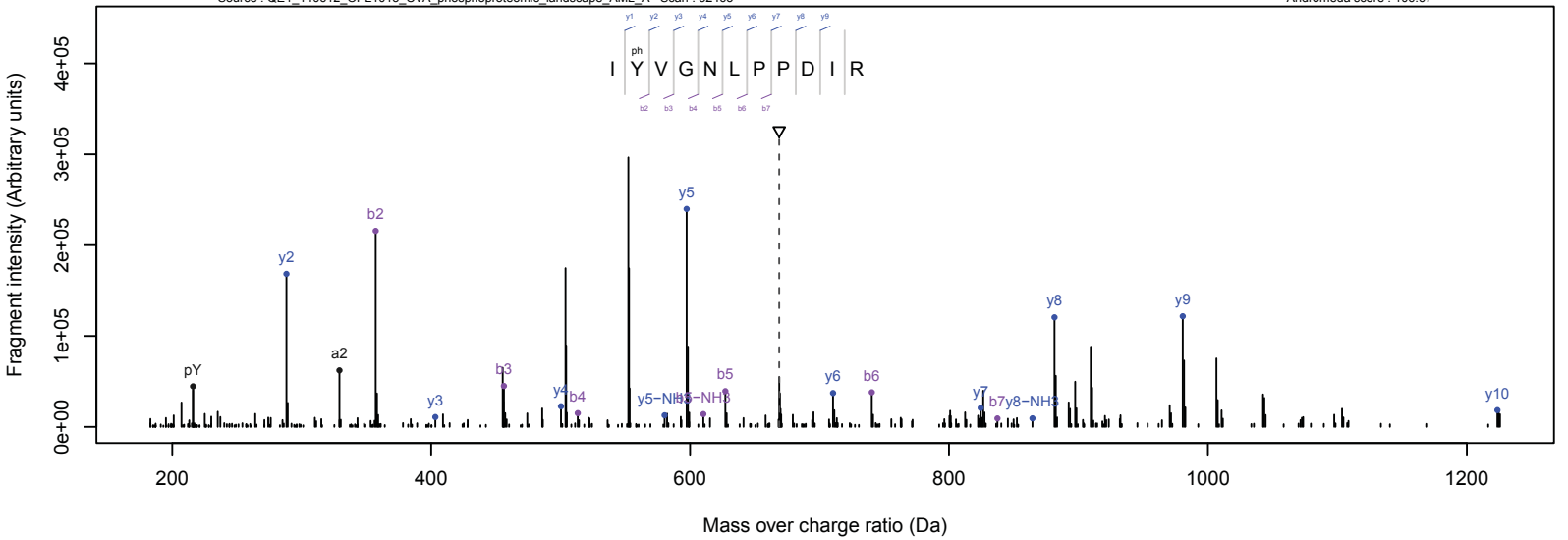
Andromeda score : 106.67



**IY(ph)VGNLPPDIR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32463

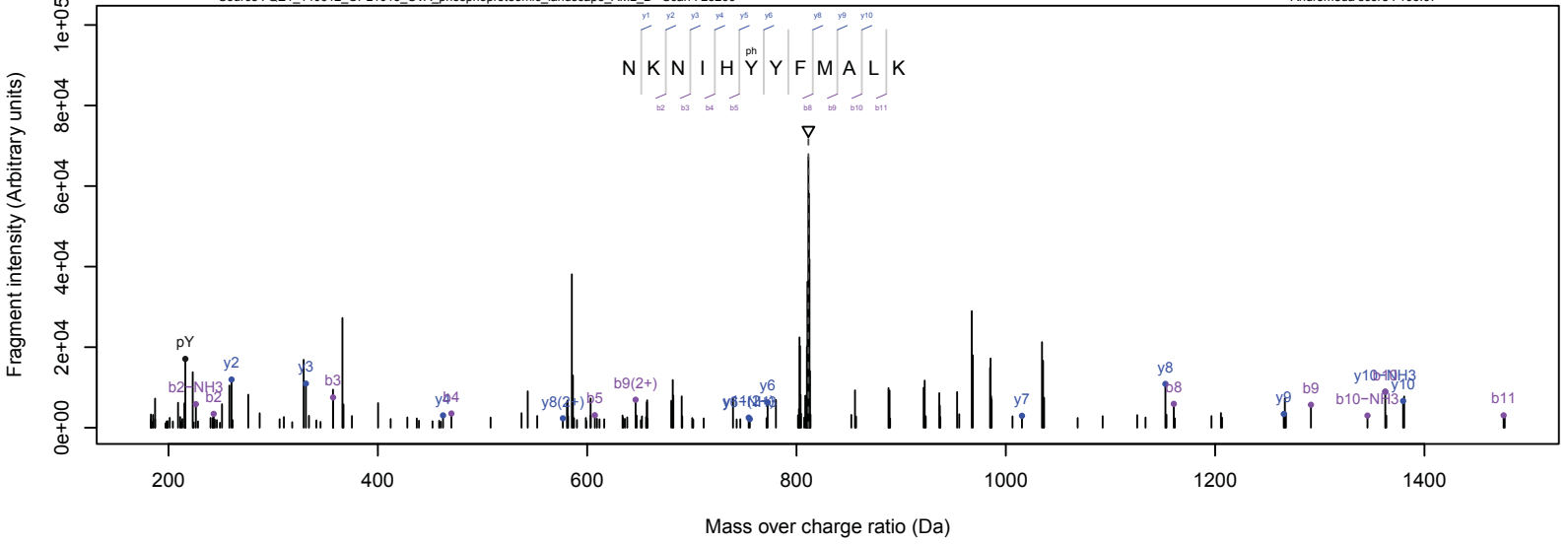
Andromeda score : 106.67



**NKNIHY(ph)YFMALK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28266

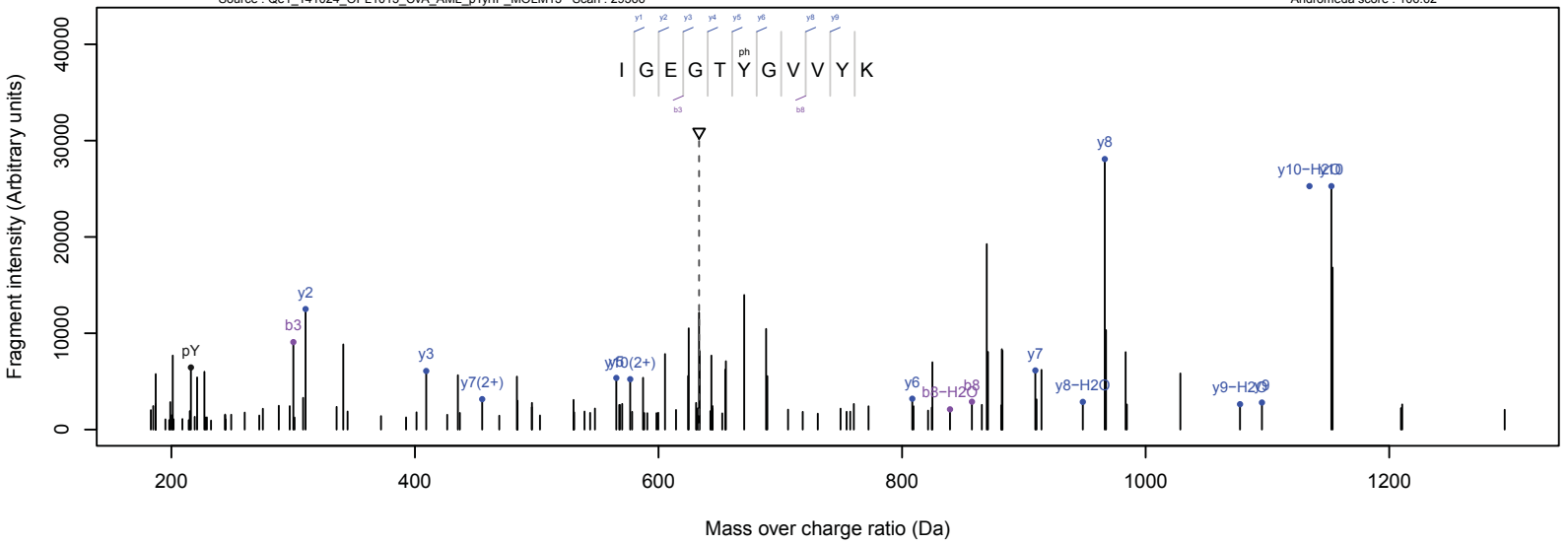
Andromeda score : 106.67



**IGEGTY(ph)GVVYK-2+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 29366

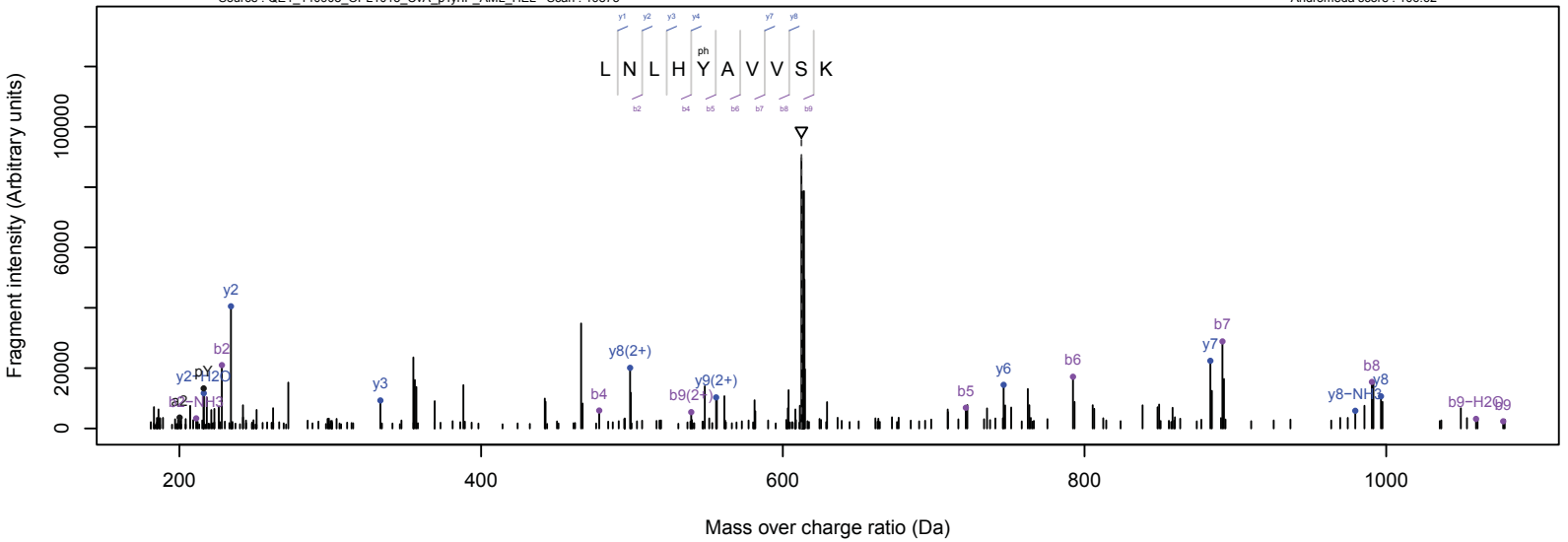
Andromeda score : 106.62



**LNLHY(ph)AVVSK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 19873

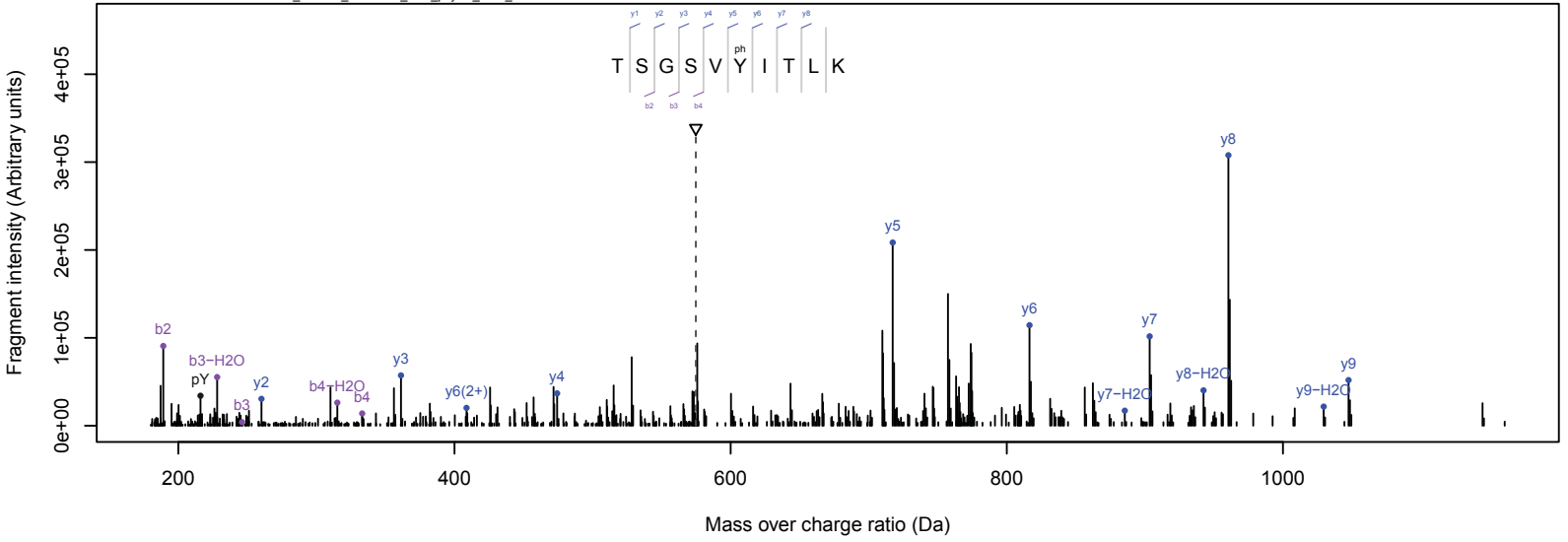
Andromeda score : 106.62



TSGSVY(ph)TLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24327

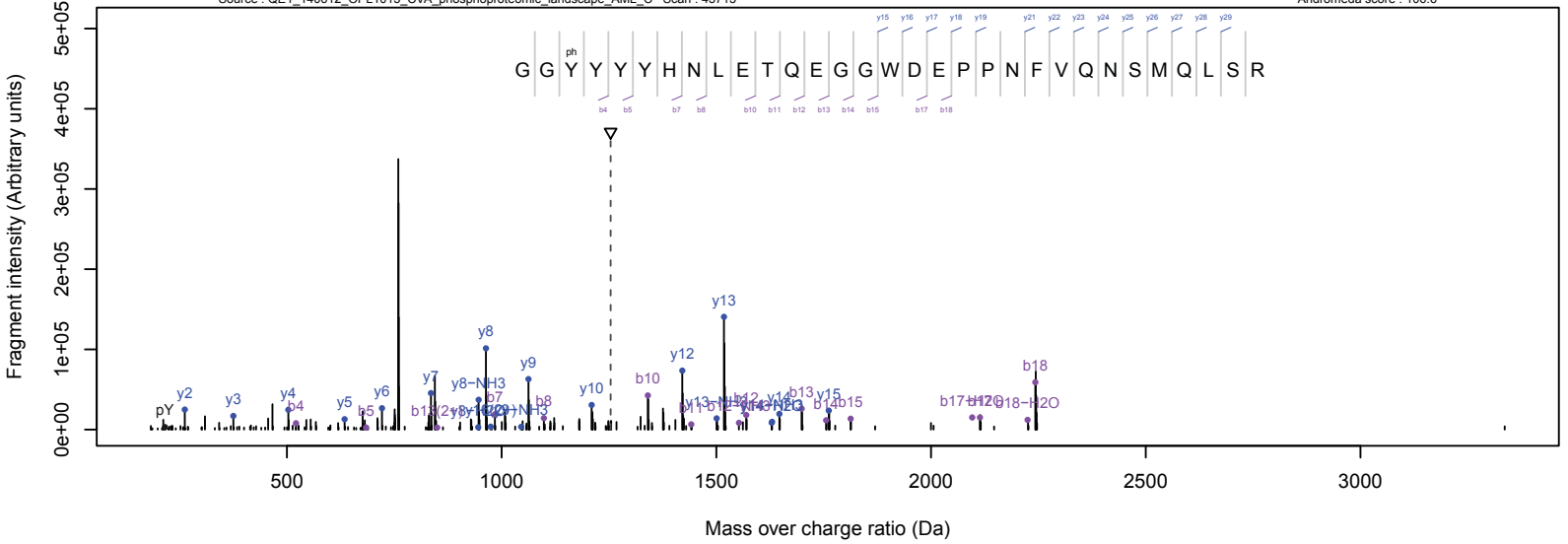
Andromeda score : 106.62



GGY(ph)YYYHNLETQEGGWDEPPNFVQNSMQLSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43713

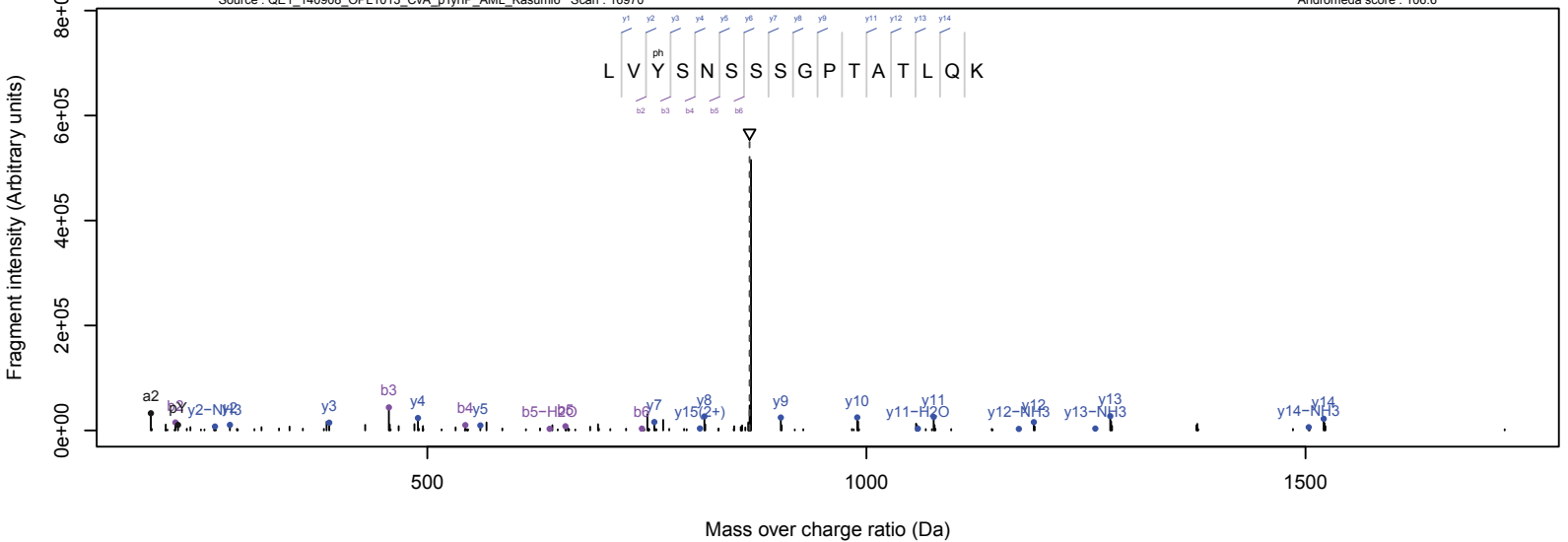
Andromeda score : 106.6



LVY(ph)SNSSSGPTATLQK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 16970

Andromeda score : 106.6

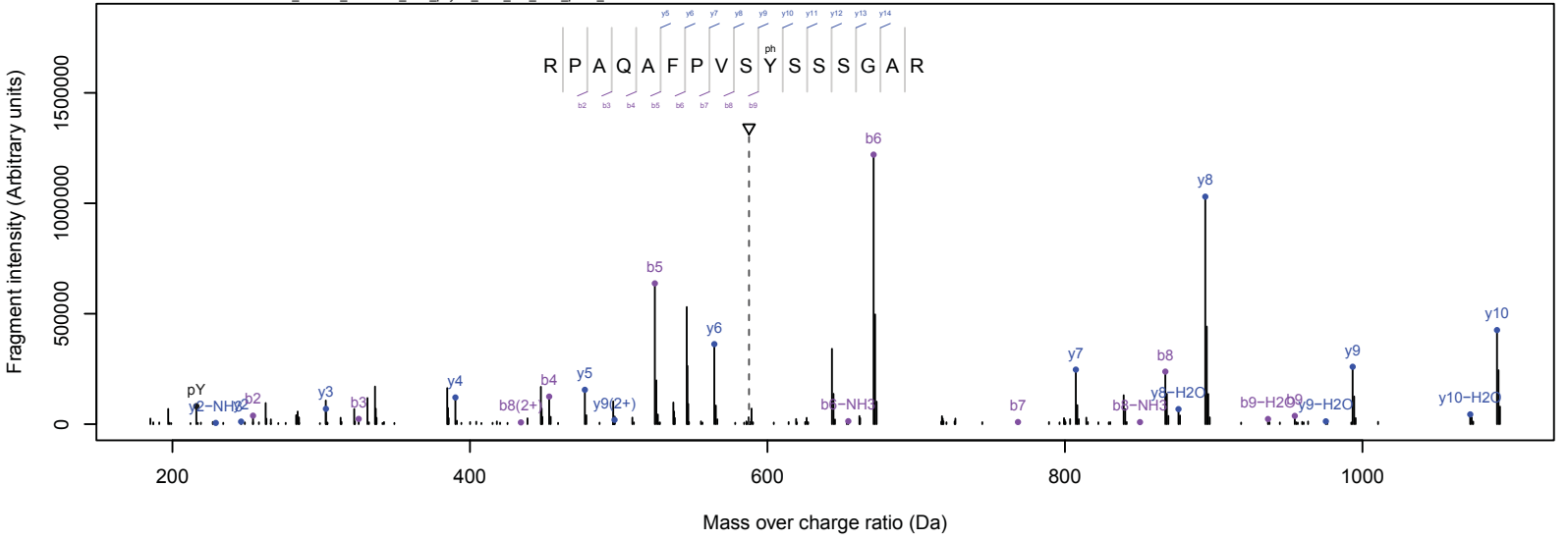




RPAQAFPVSY(ph)SSSGAR-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 10550

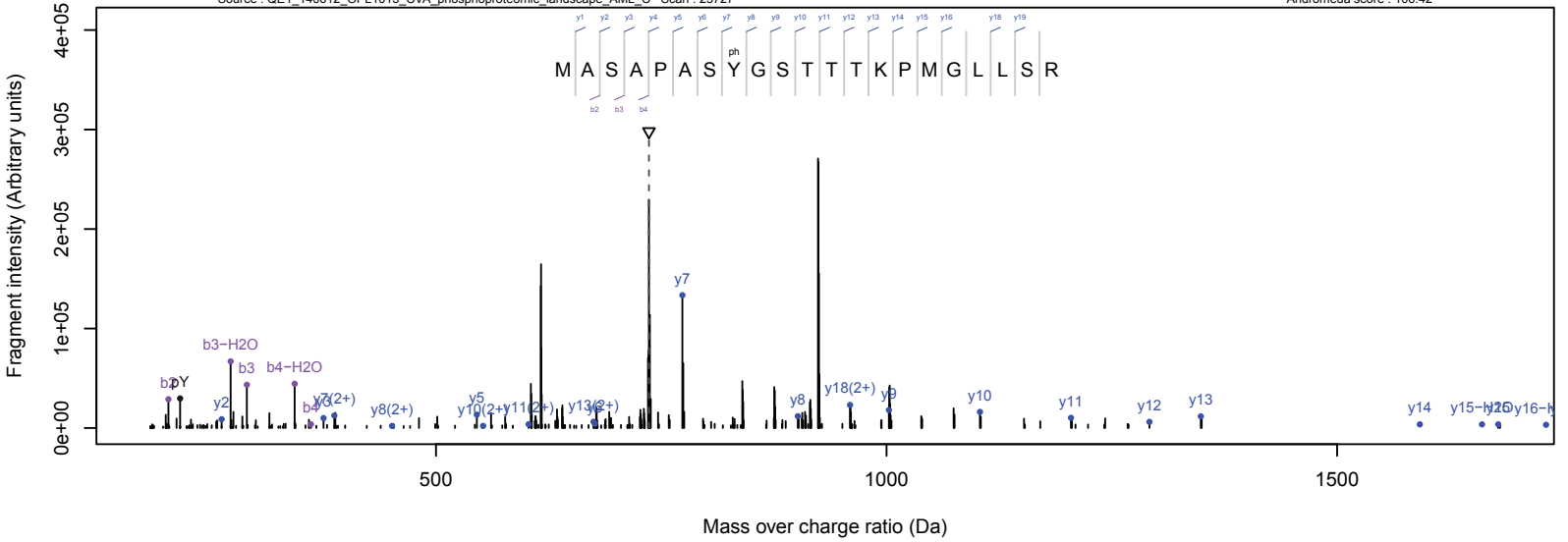
Andromeda score : 106.6



MASAPASY(ph)GSTTKPMGLLSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25727

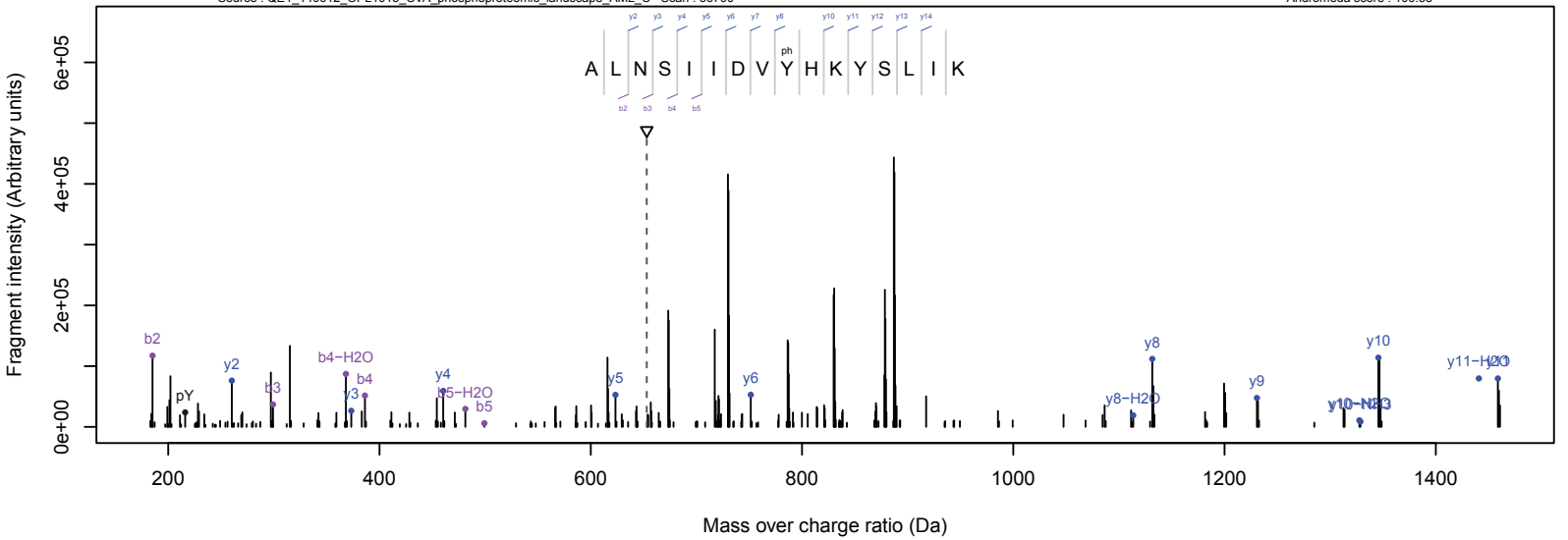
Andromeda score : 106.42



ALNSIDVY(ph)HKYSLIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36790

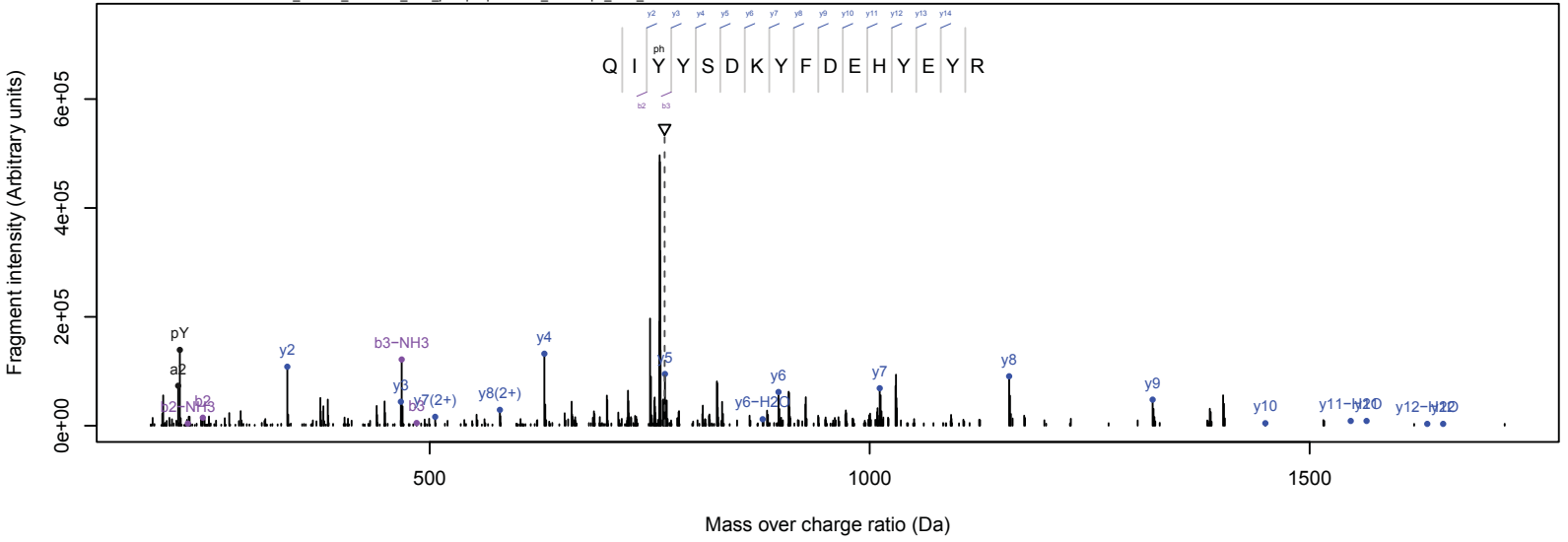
Andromeda score : 106.38



QIY(ph)YSDKYFDEHYEYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27603

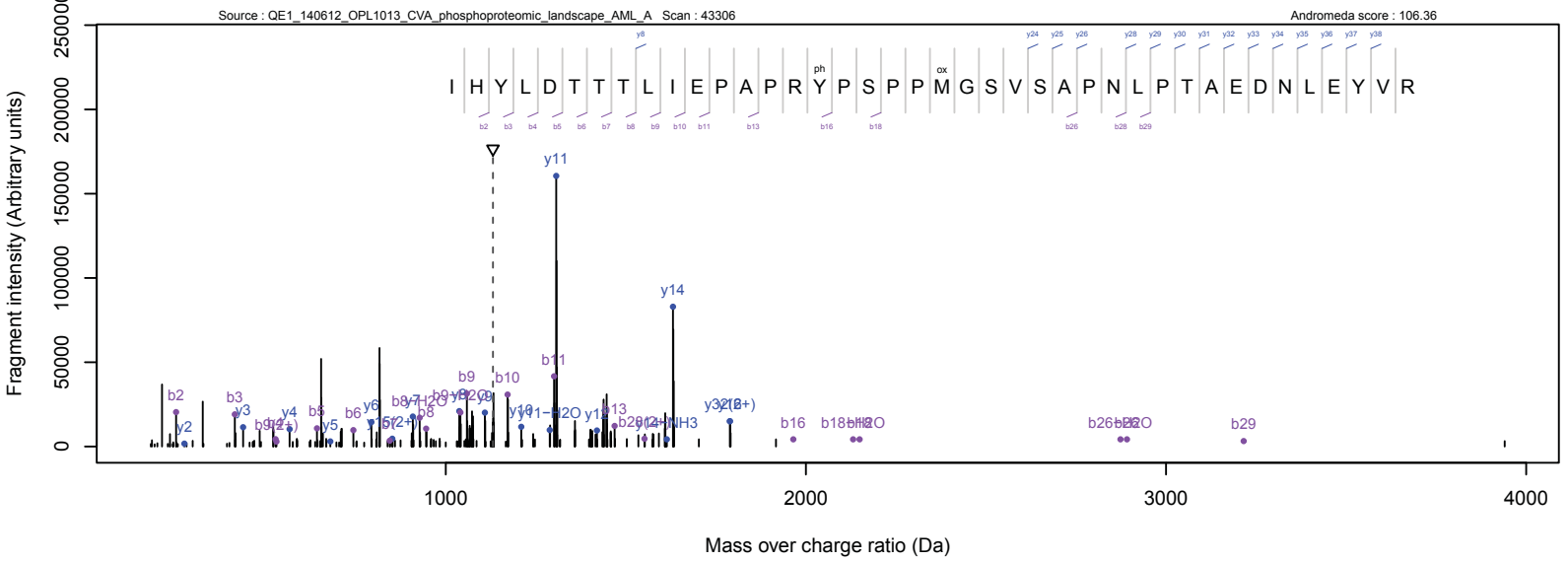
Andromeda score : 106.38



IHYLDTTLLIEPAPRY(ph)PSPPM(ox)GSVSAPNLPPTAEDNLEYVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 43306

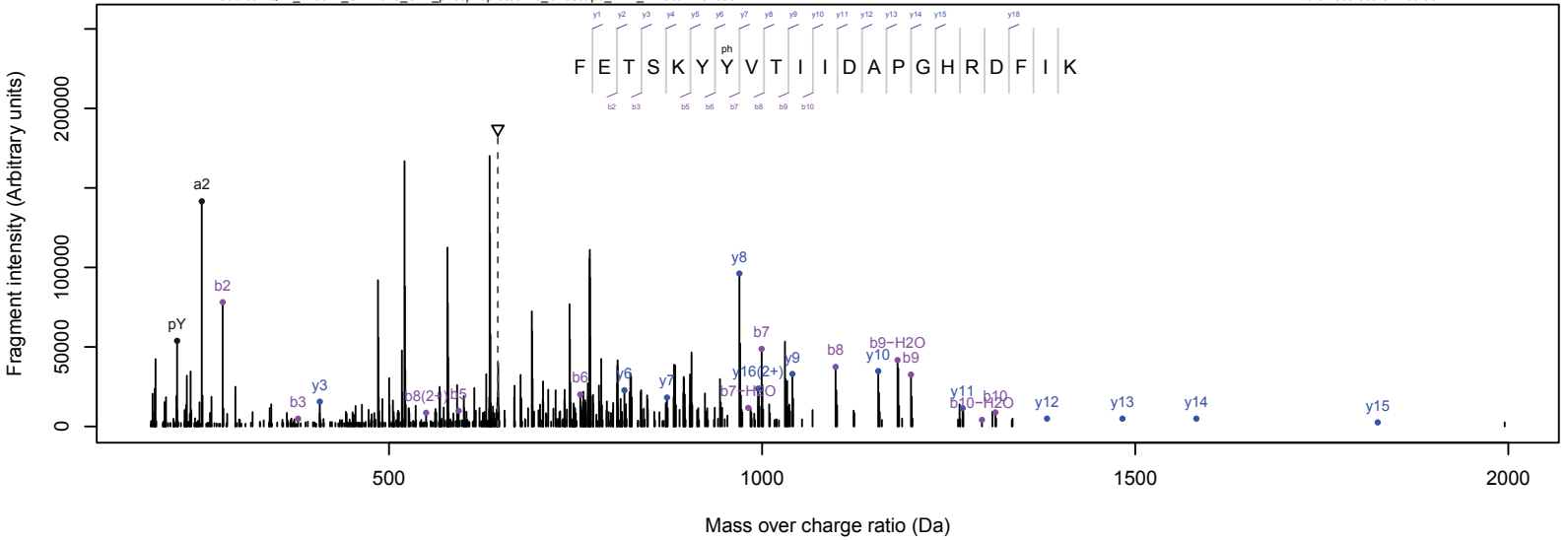
Andromeda score : 106.36



FETSKYY(ph)VTIIDAPGHRDFIK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32650

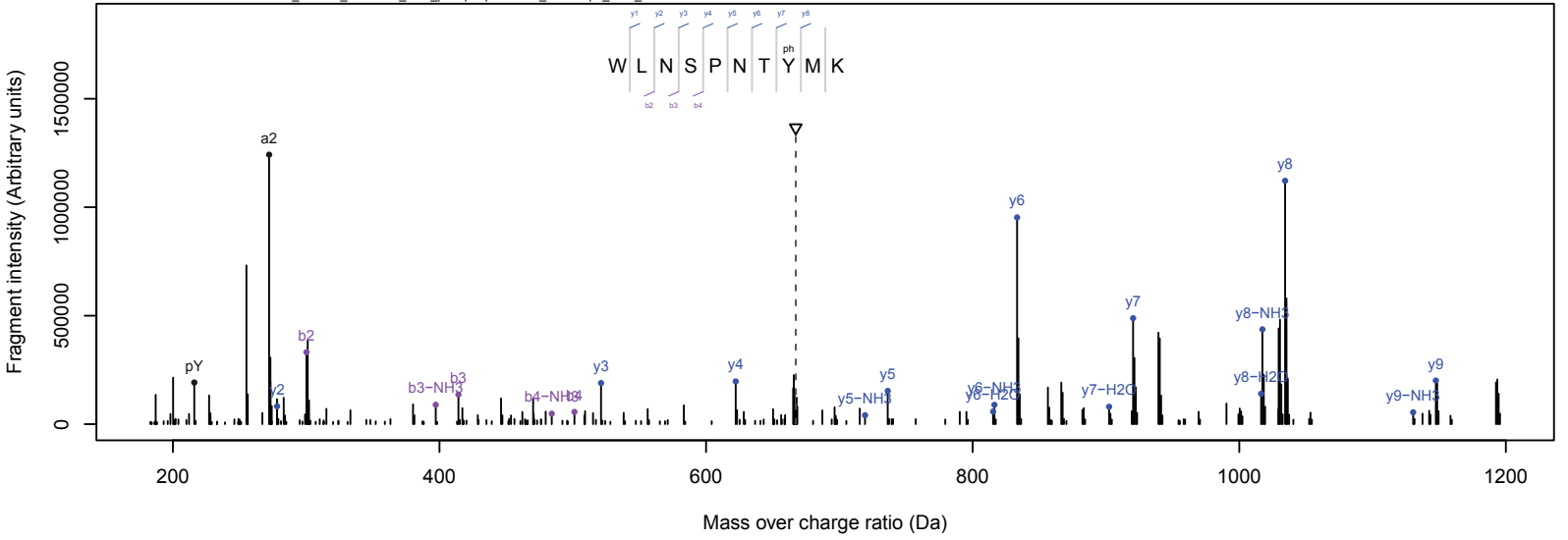
Andromeda score : 106.35



**WLNSPNTY(ph)MK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22841

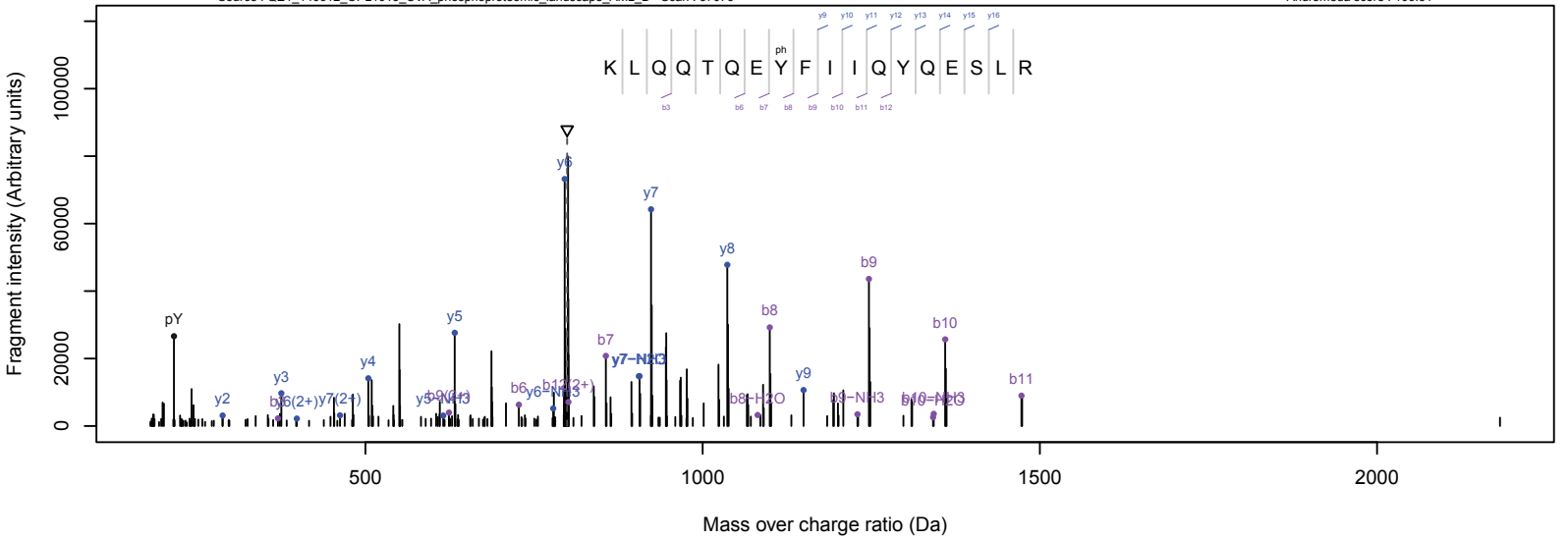
Andromeda score : 106.35



**KLQQTQEY(ph)FIQYQESLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 37076

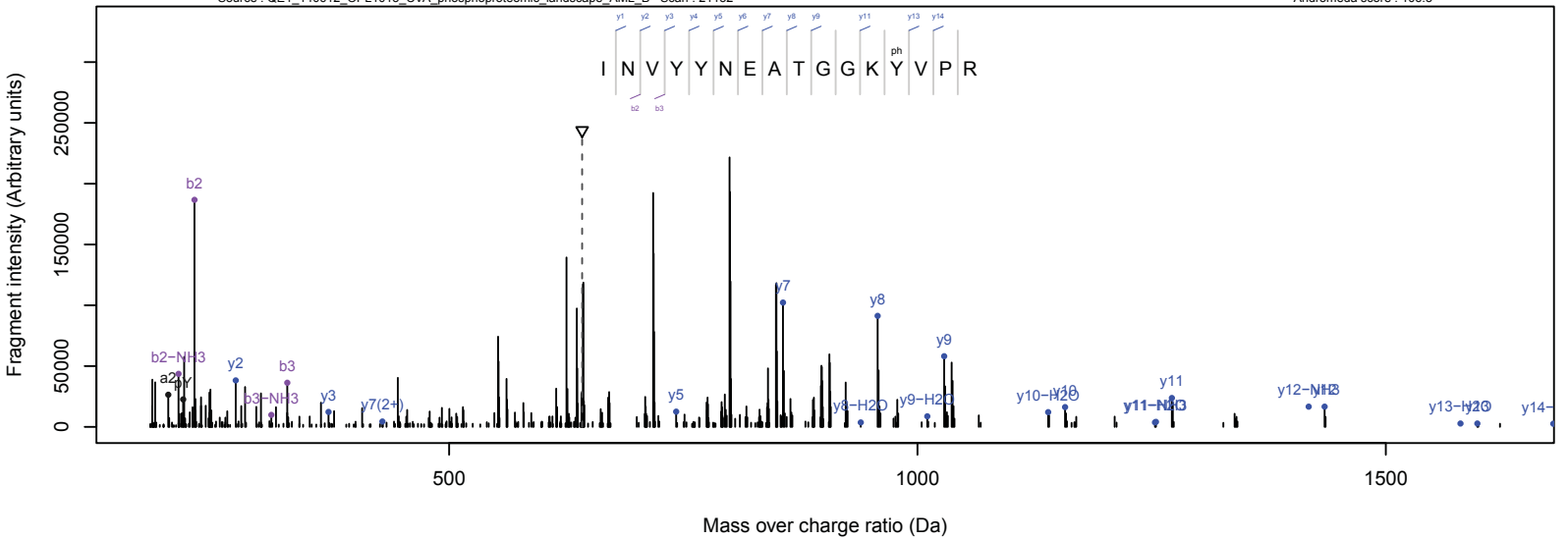
Andromeda score : 106.31



**INVYNEATGGKY(ph)VPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21132

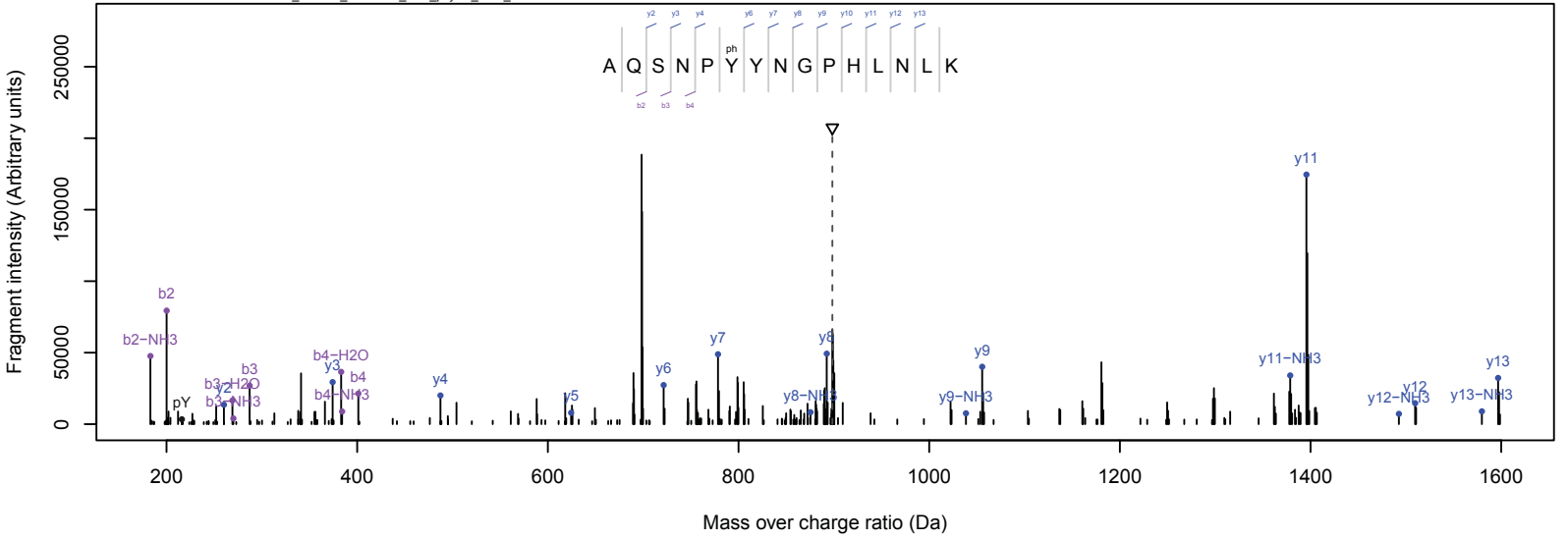
Andromeda score : 106.3



**AQSNPY(ph)YNGPHLNLK-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan: 23188

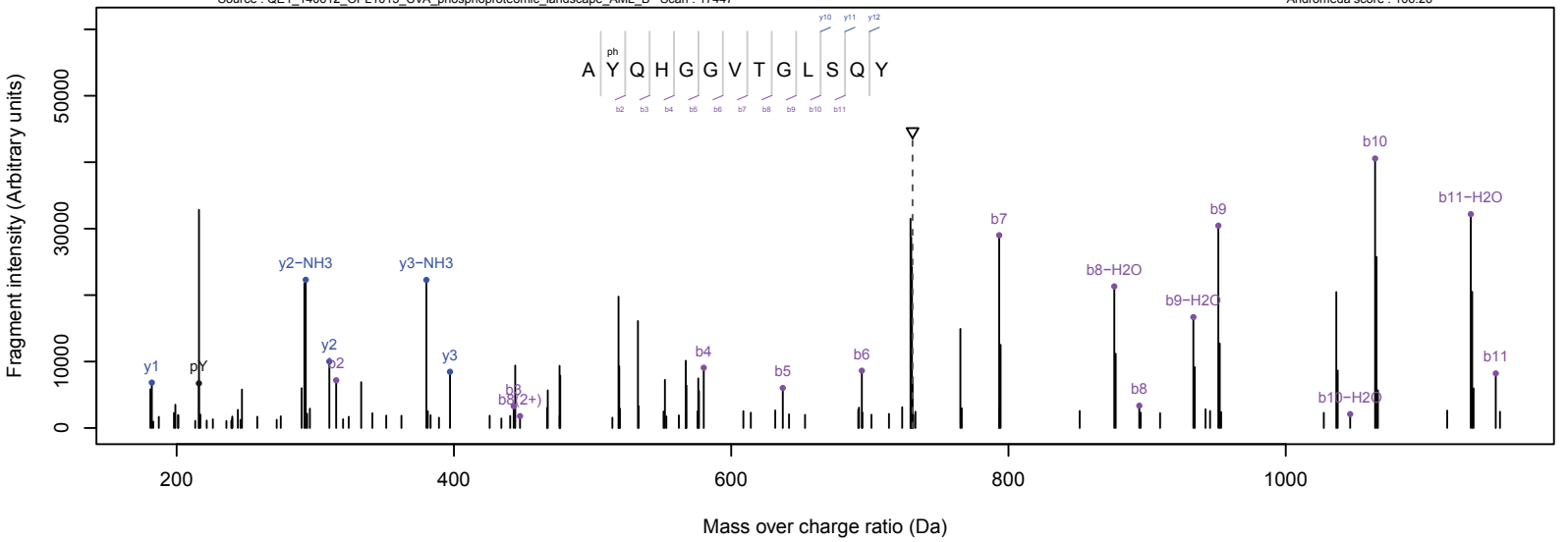
Andromeda score: 106.29



**AY(ph)QHGGVTGLSQY-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 17447

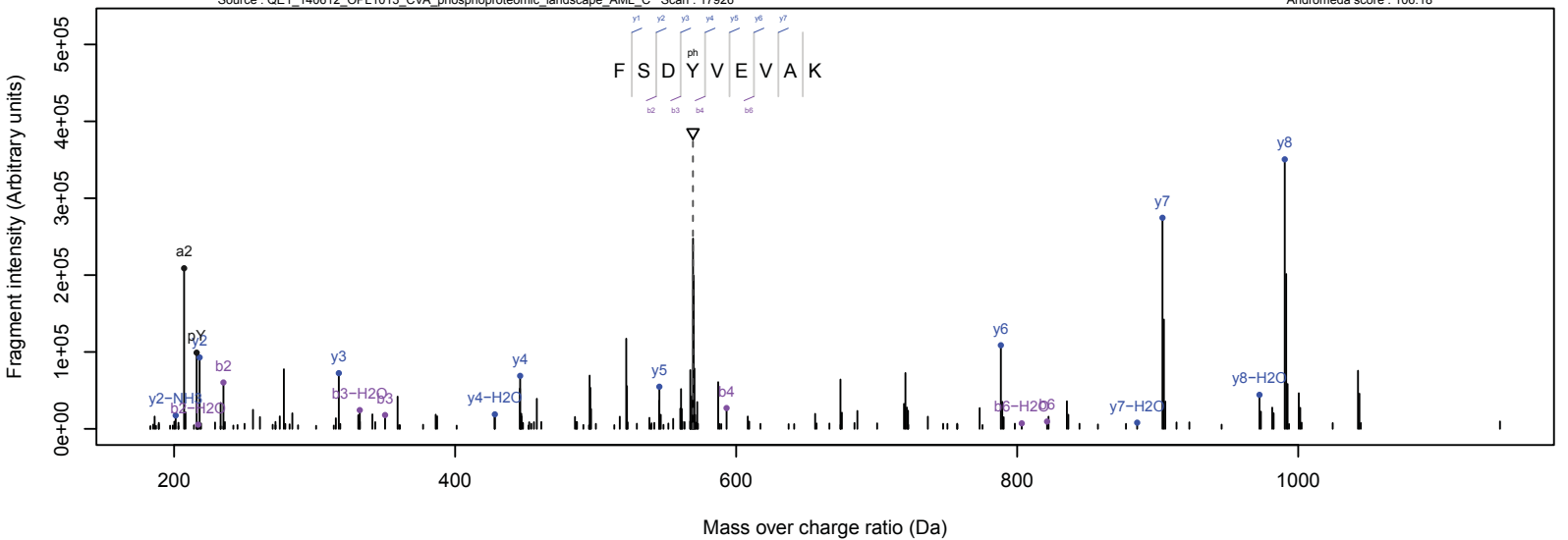
Andromeda score: 106.26



**FSDY(ph)VEVAK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17926

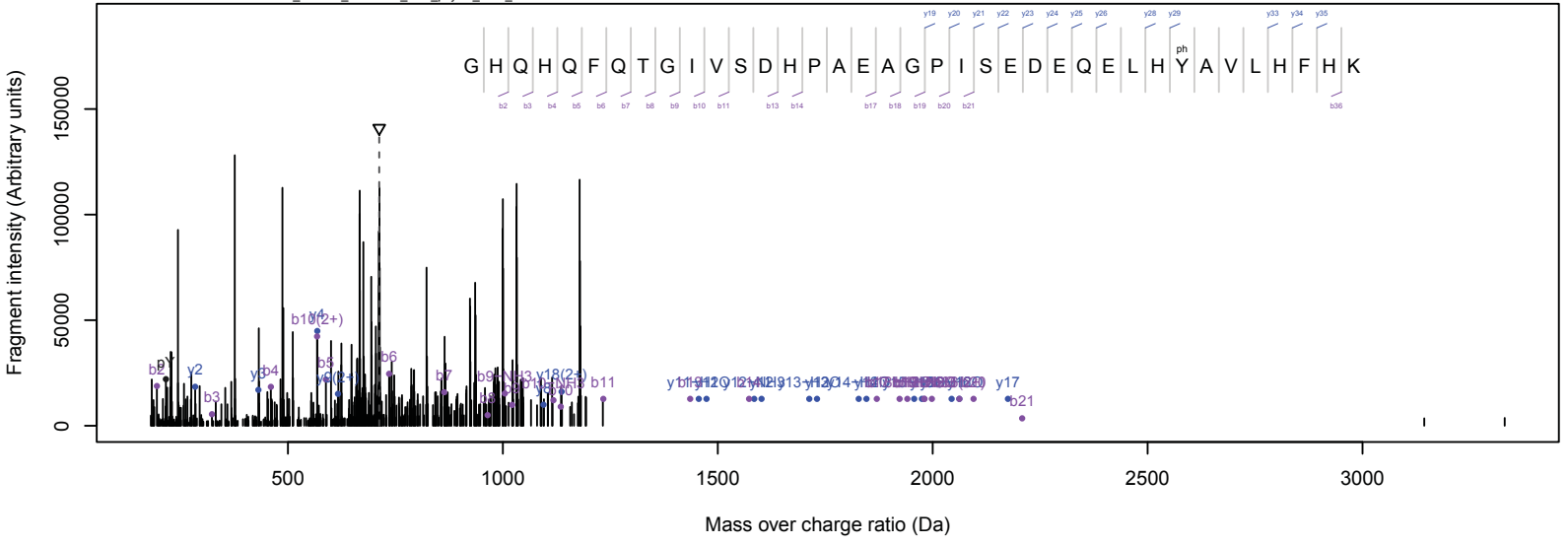
Andromeda score: 106.18



**GHQHQTGIVSDHPAEAGPISEDEQELHY(ph)AVLHFHK-6+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 37657

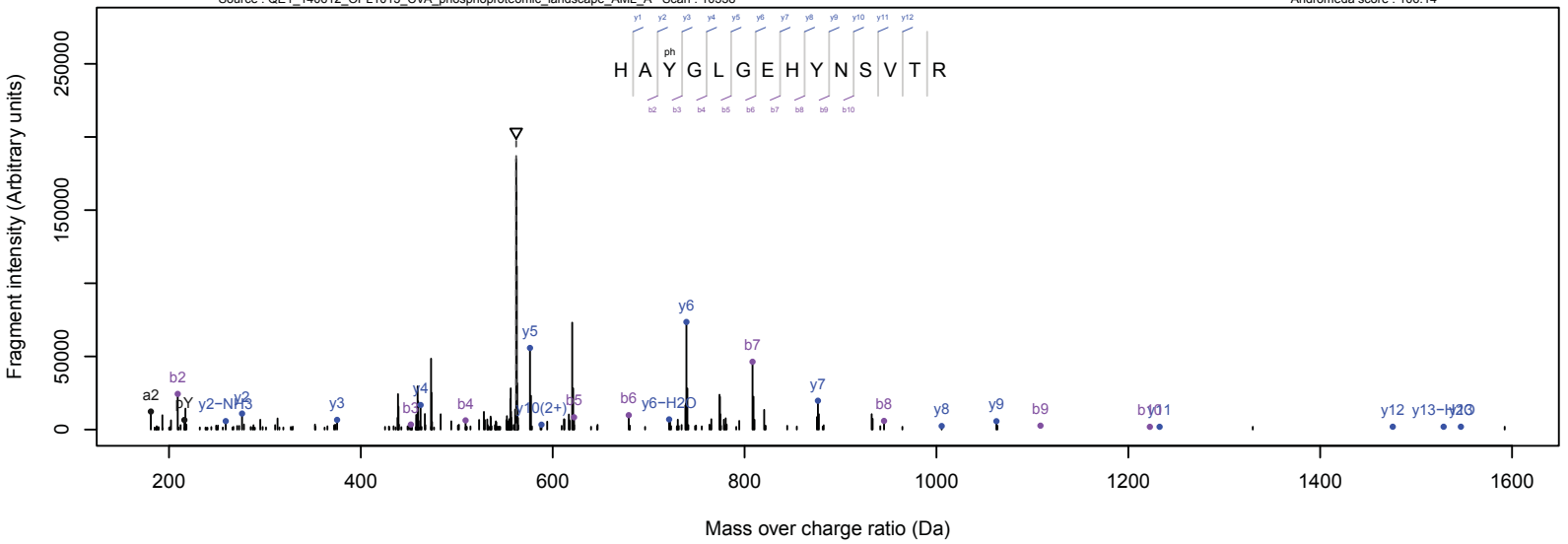
Andromeda score : 106.18



**HAY(ph)GLGEHYSVTR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 10338

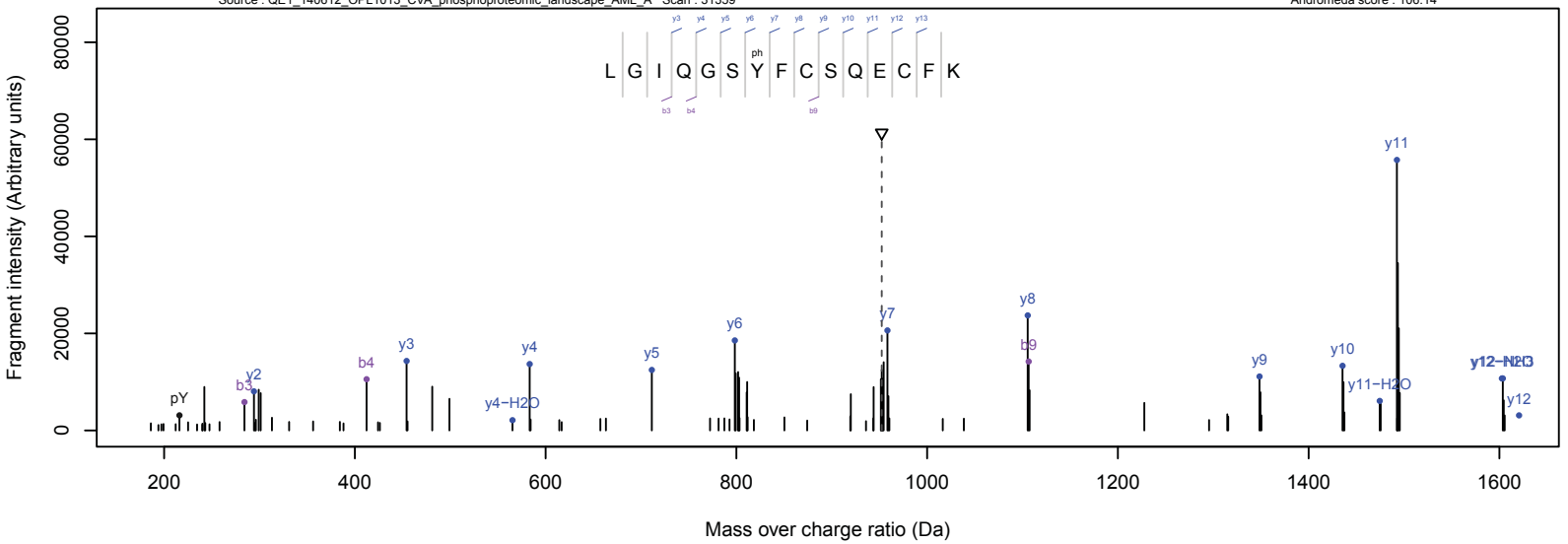
Andromeda score : 106.14



**LGIQGSY(ph)FCSQECFK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 31359

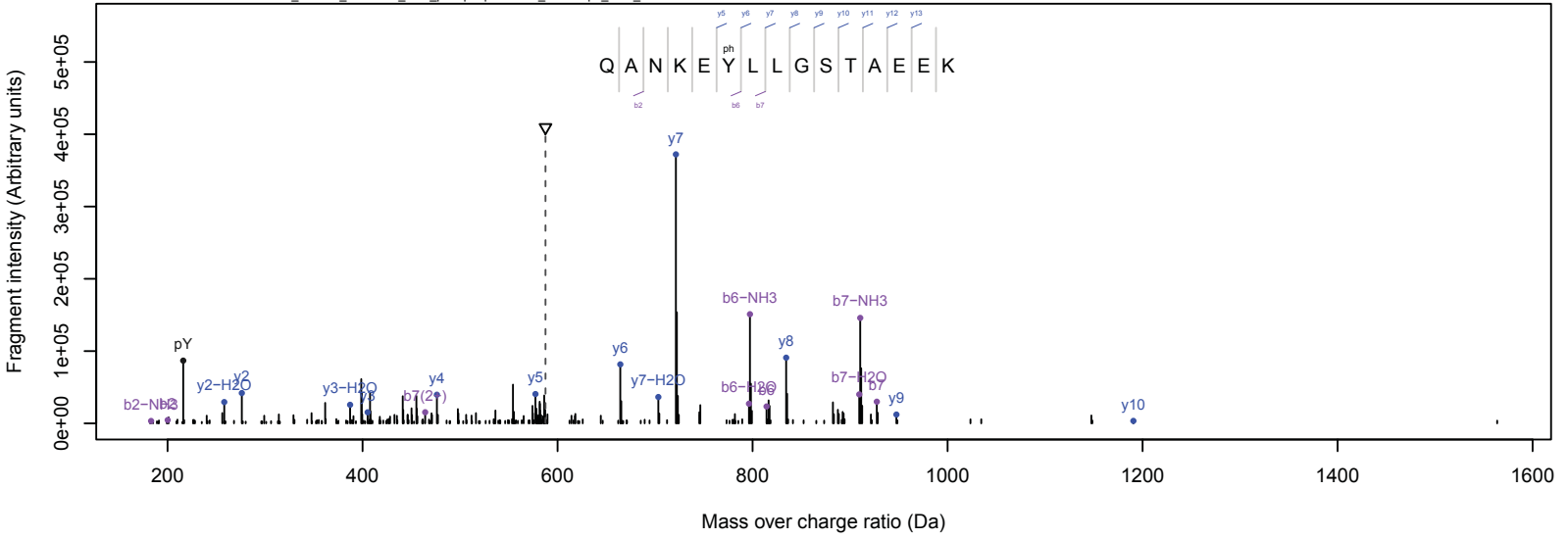
Andromeda score : 106.14



QANKEY(ph)LLGSTAEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12973

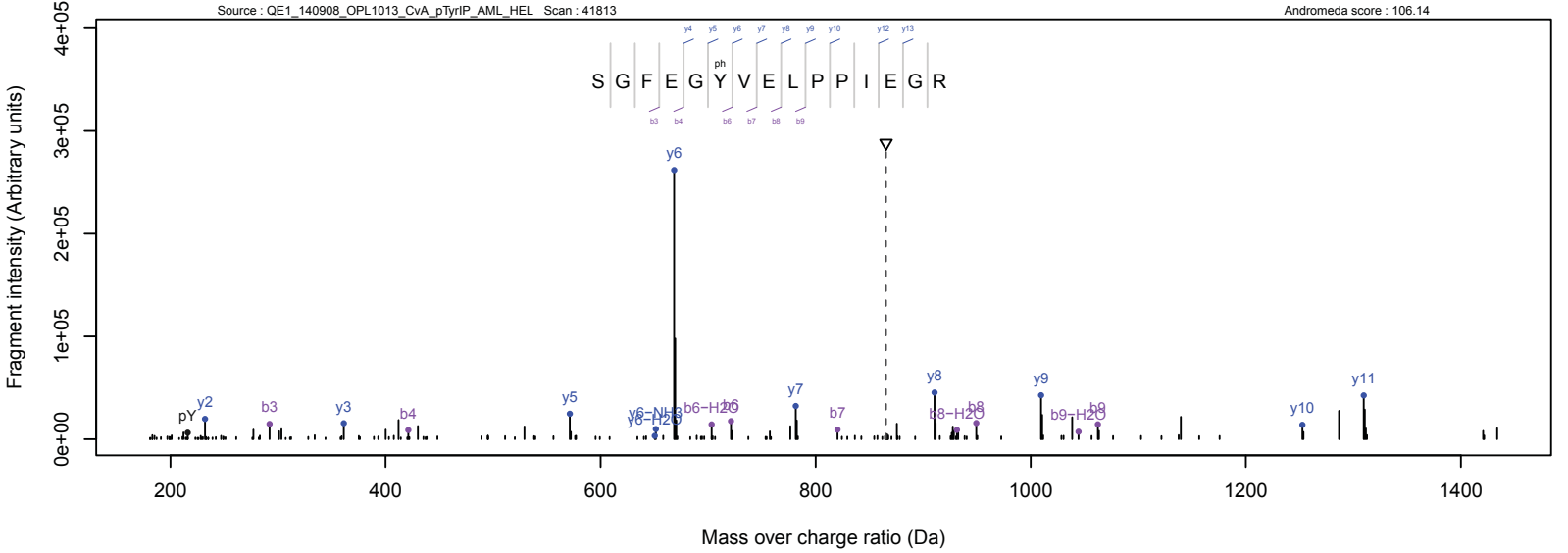
Andromeda score : 106.14



SGFEGY(ph)VELPPIEGR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 41813

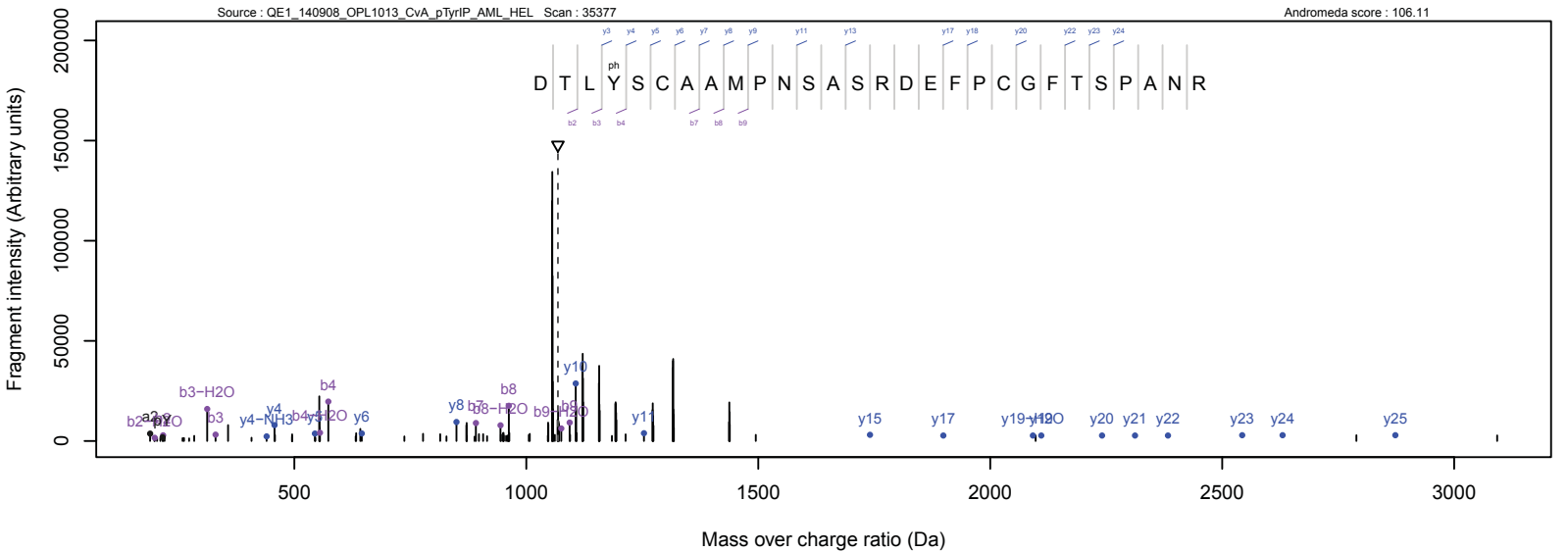
Andromeda score : 106.14



DTLY(ph)SCAAMPNSASRDEFPCGFTSPANR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35377

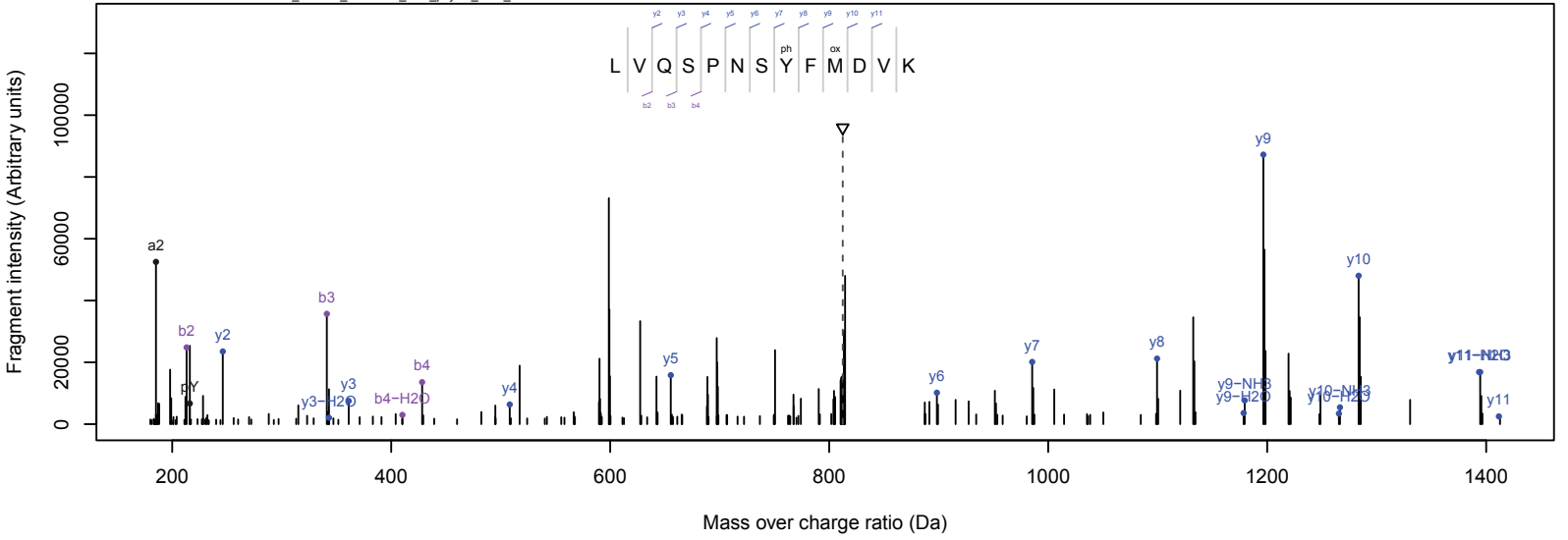
Andromeda score : 106.11



LVQSPNSY(ph)FM(ox)DVK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 24732

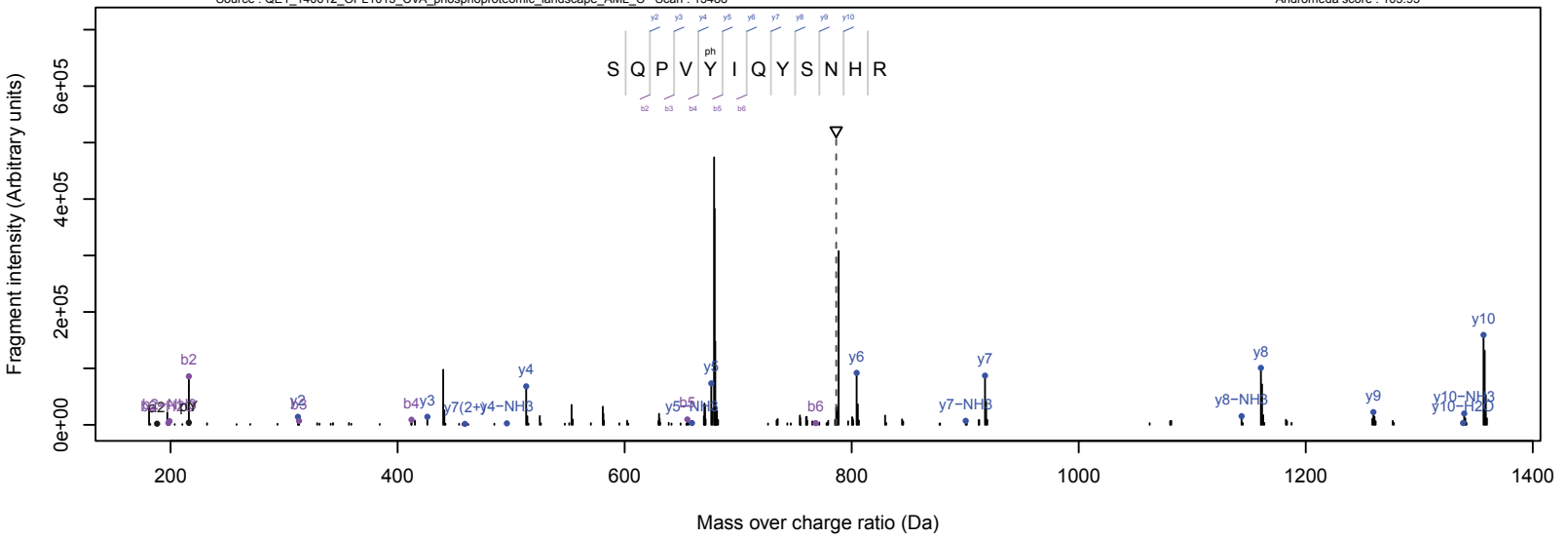
Andromeda score : 105.95



SQPVY(ph)IQYSNHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13488

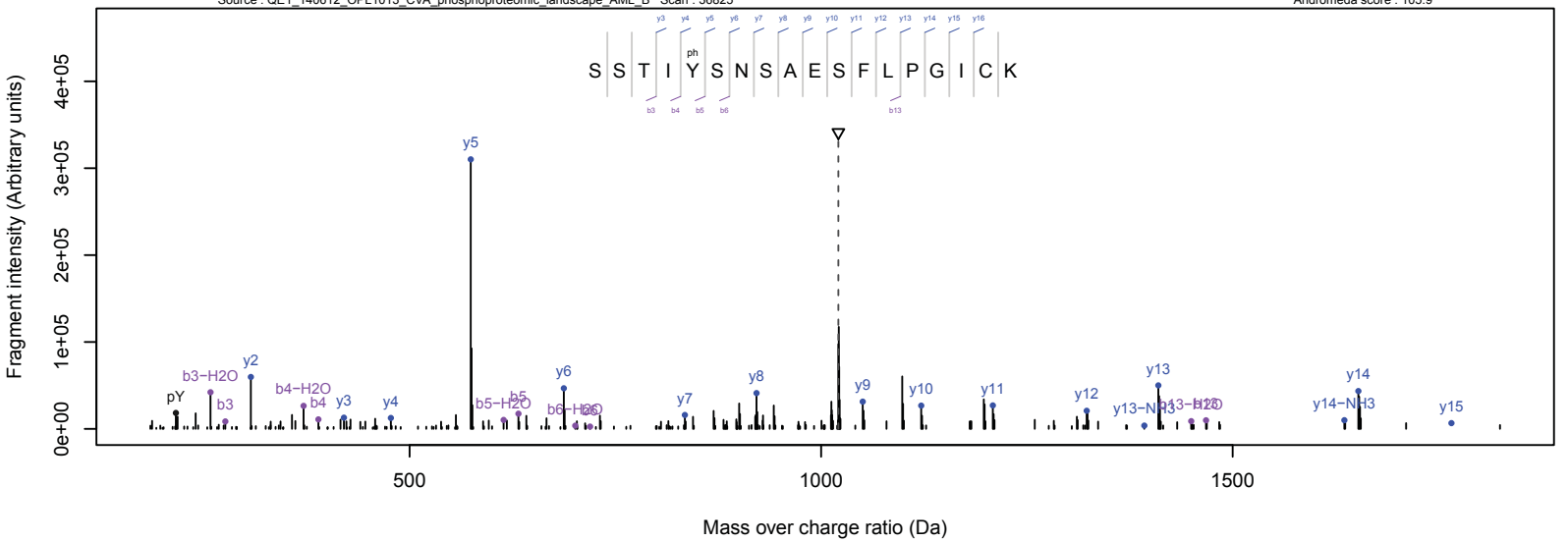
Andromeda score : 105.95



SSTIY(ph)SNAESFLPGICK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36825

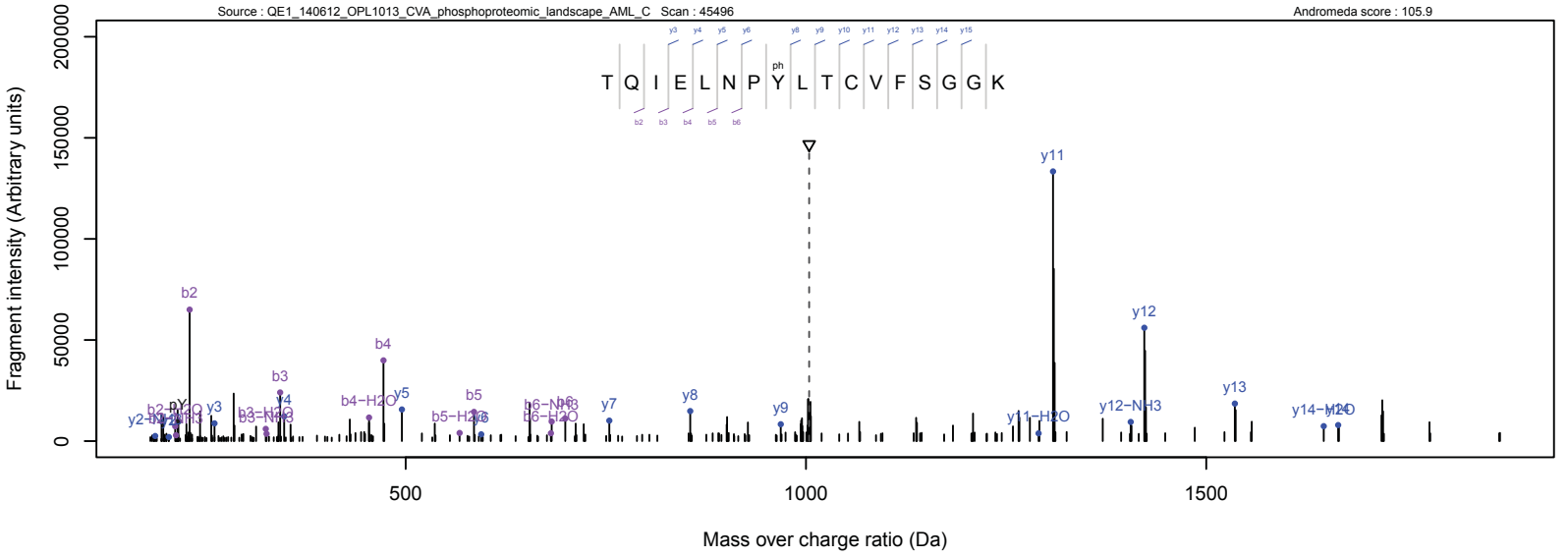
Andromeda score : 105.9



TQIELNPY(ph)LTCVFSGGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45496

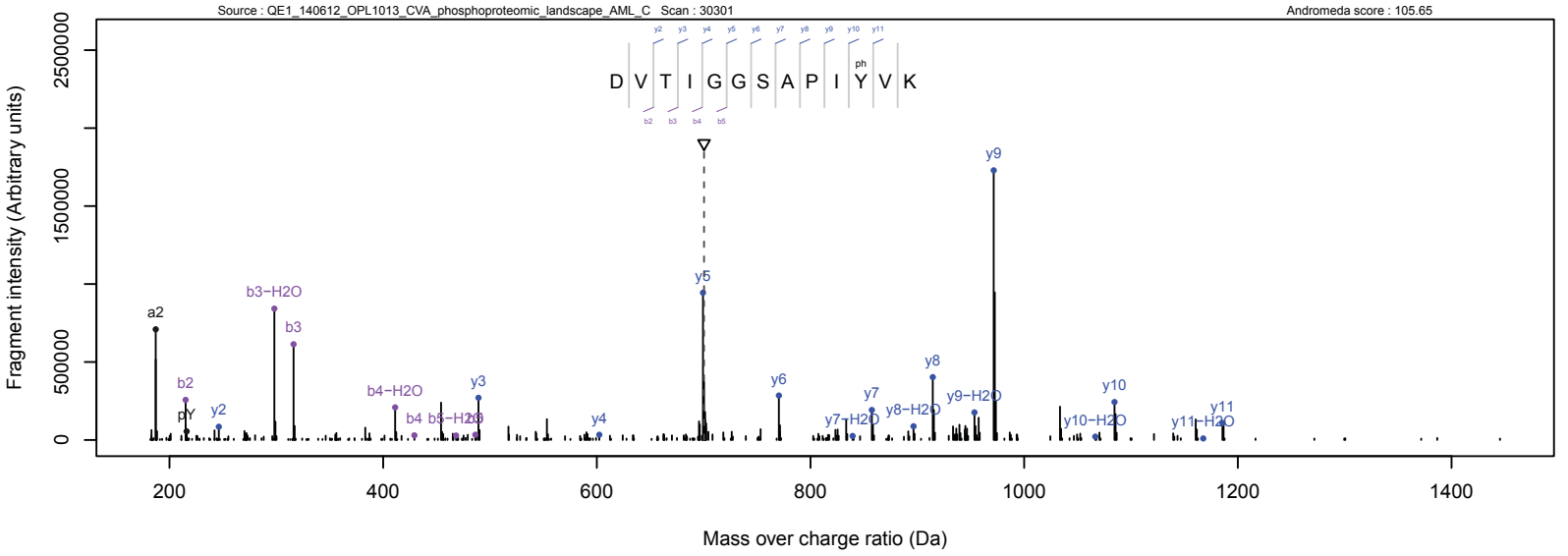
Andromeda score : 105.9



DVTIGGSAPIY(ph)VK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30301

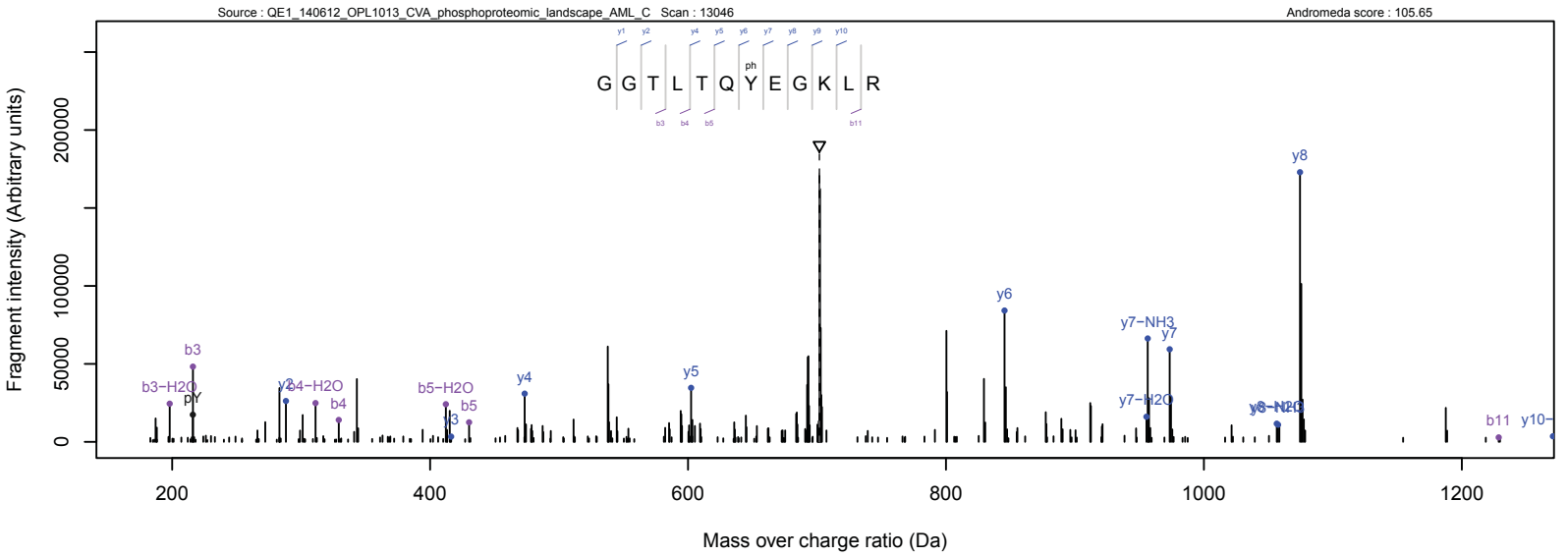
Andromeda score : 105.65



GGTLTQY(ph)EGKLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13046

Andromeda score : 105.65

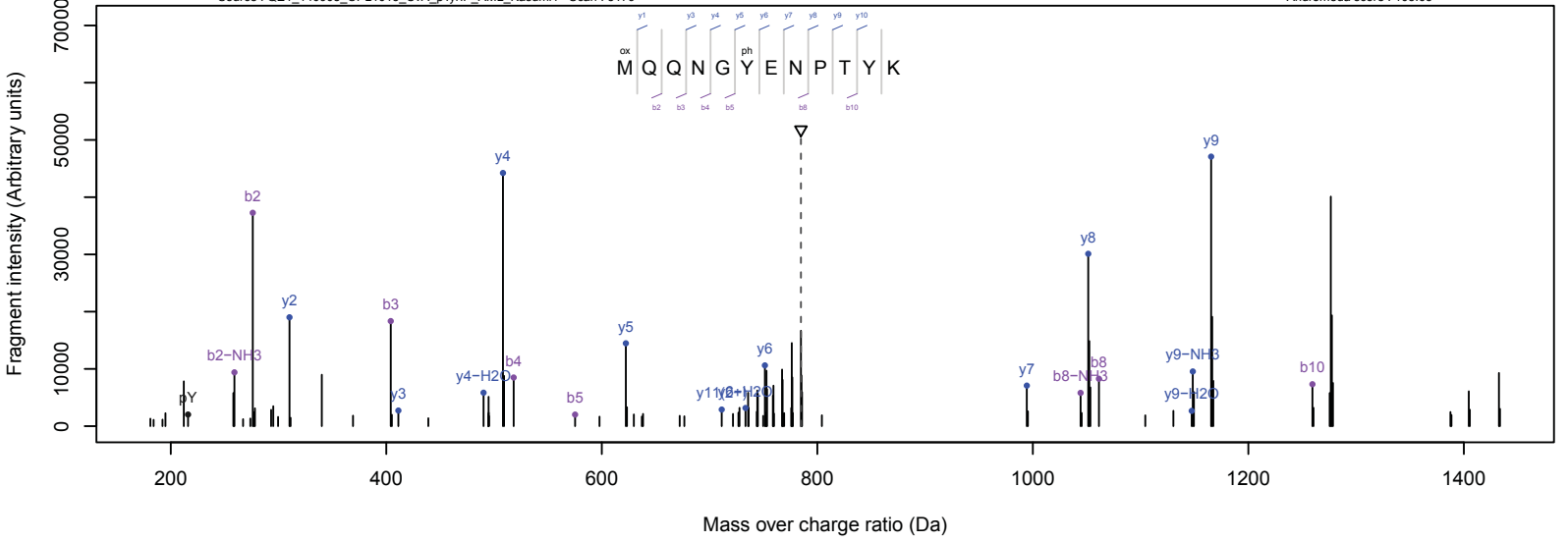




**M(ox)QQNGY(ph)ENPTYK-2+**

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan: 5179

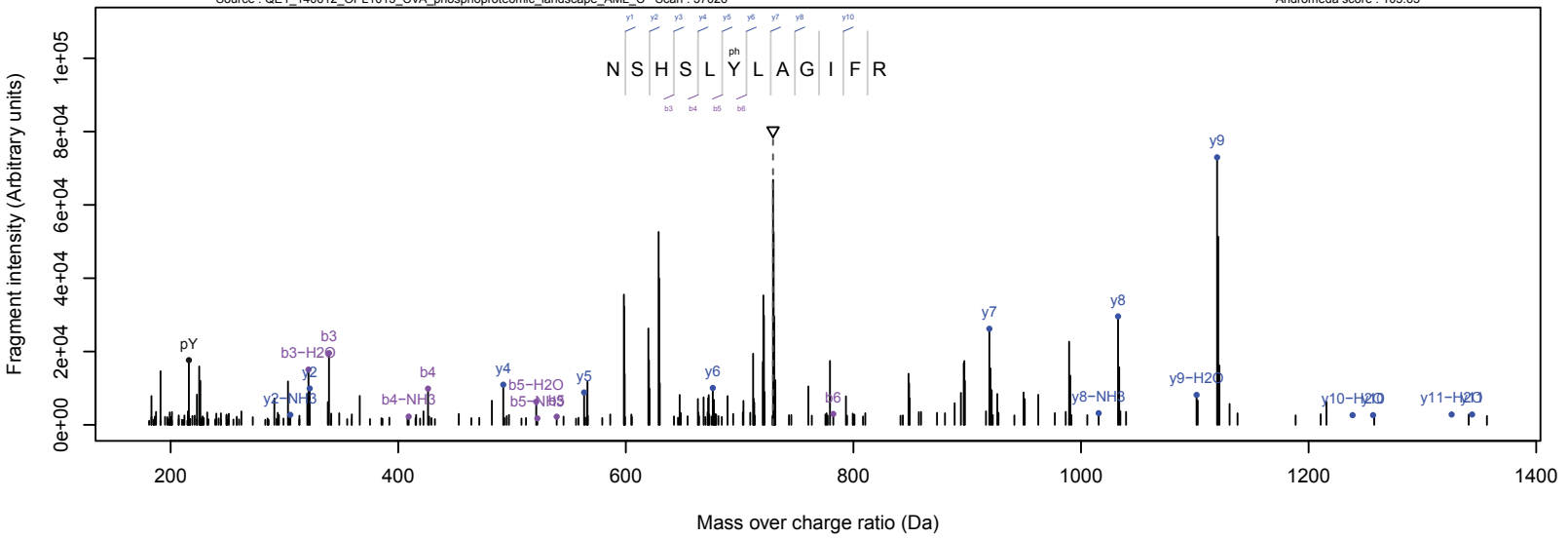
Andromeda score: 105.65



**NSHSLY(ph)LAGIFR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37026

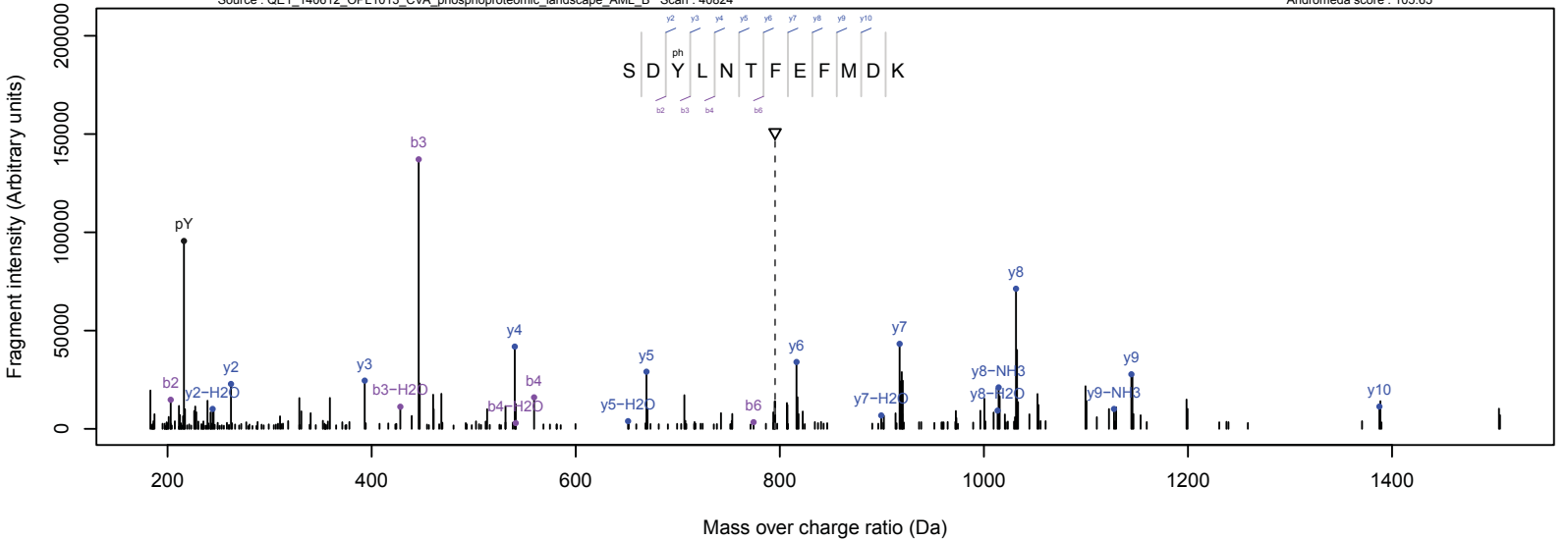
Andromeda score: 105.65



**SDY(ph)LNTFEFMDK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 40824

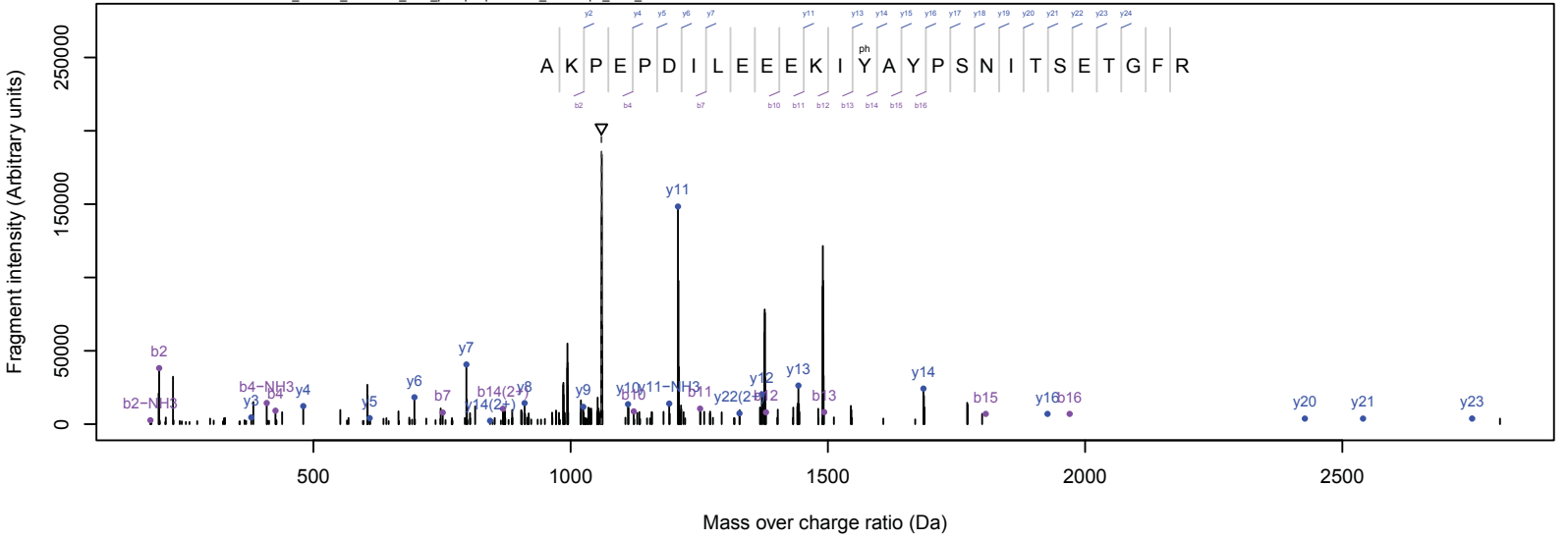
Andromeda score: 105.65



**AKPEPDILEEEKIY(ph)AYPSNITSETGFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38026

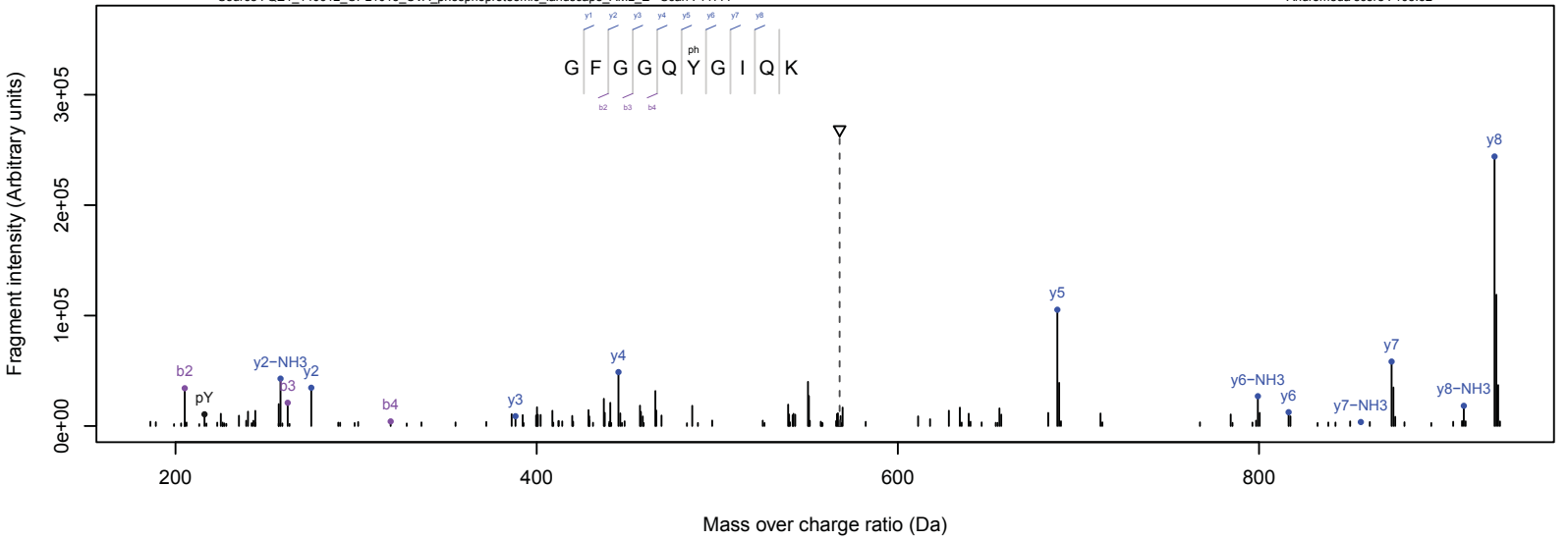
Andromeda score : 105.55



**GFGGQY(ph)GIQK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 11777

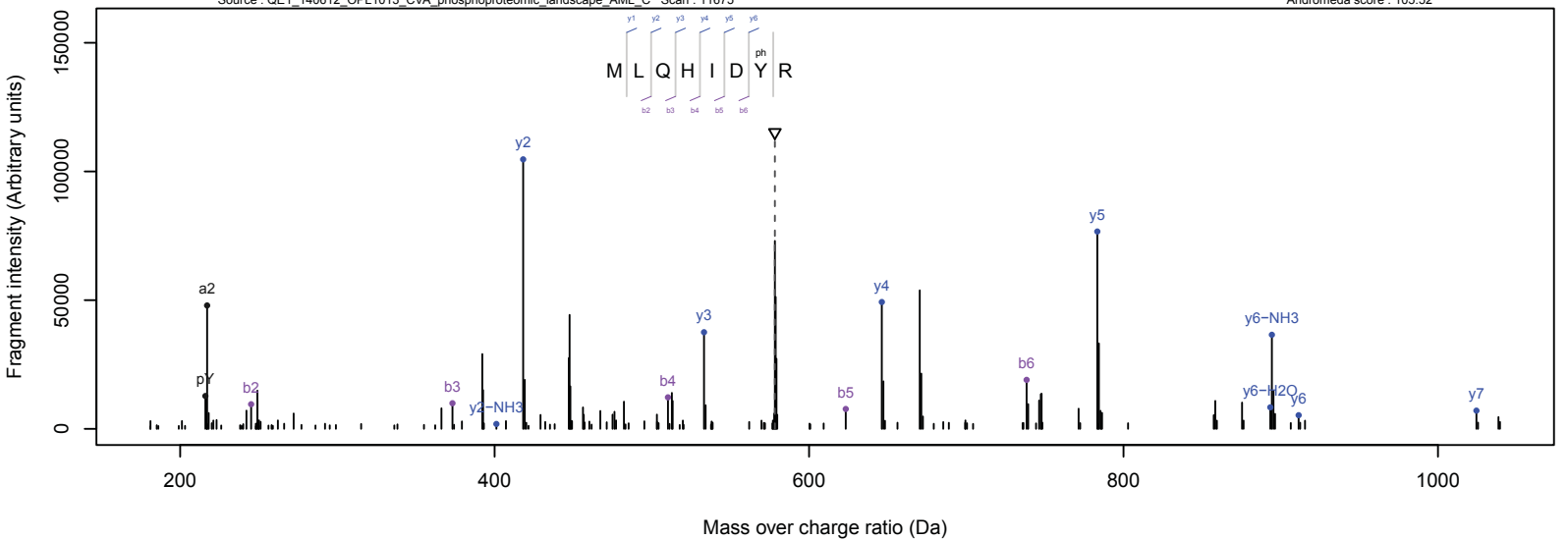
Andromeda score : 105.52



**MLQHIDY(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11675

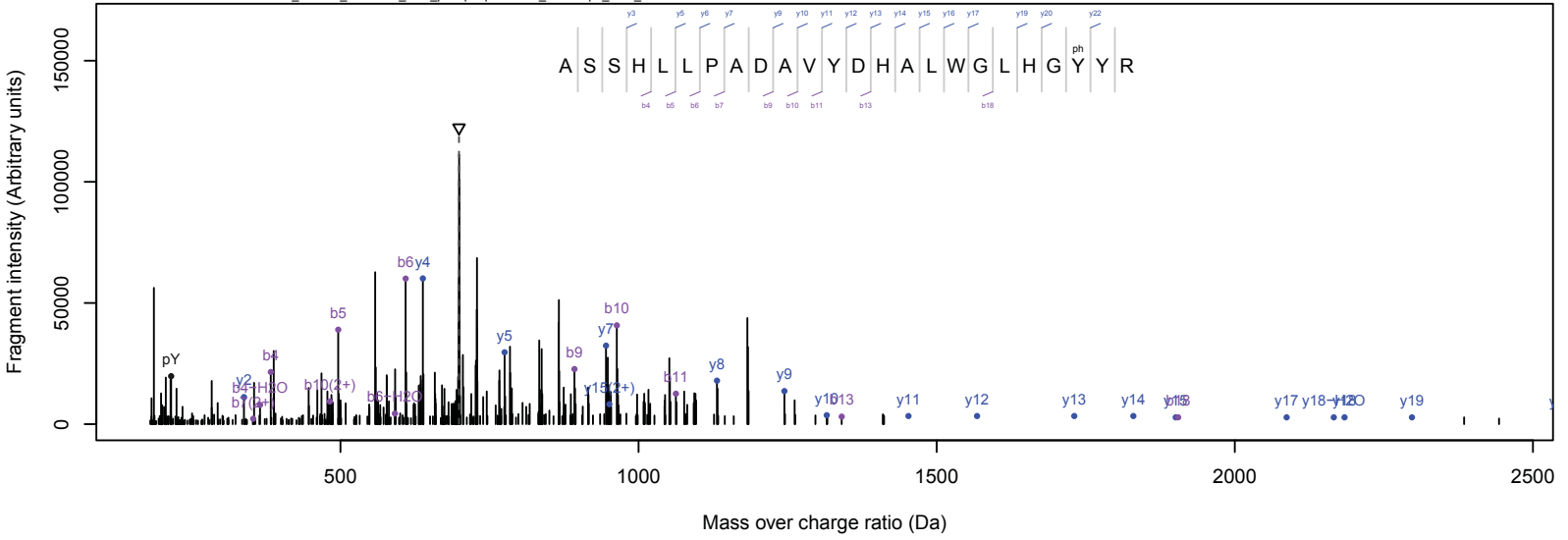
Andromeda score : 105.52



ASSHLLPADAVYDHALWGLHGYP-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39751

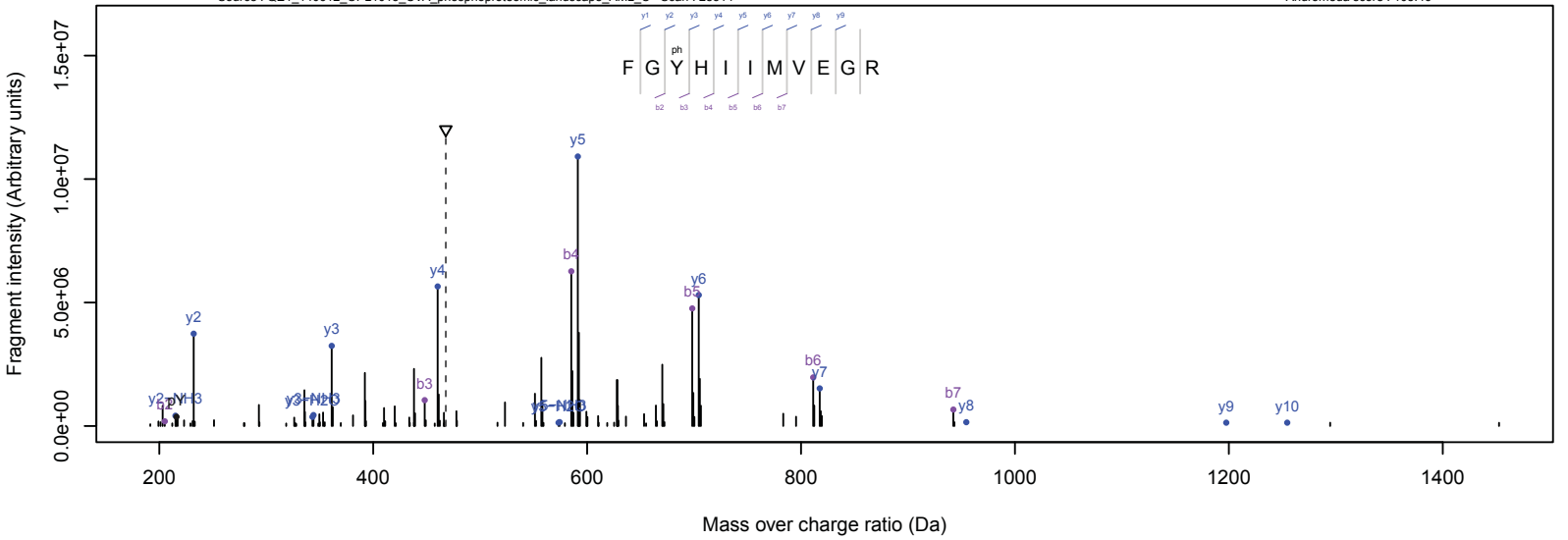
Andromeda score : 105.51



FGY(ph)HIMVEGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28514

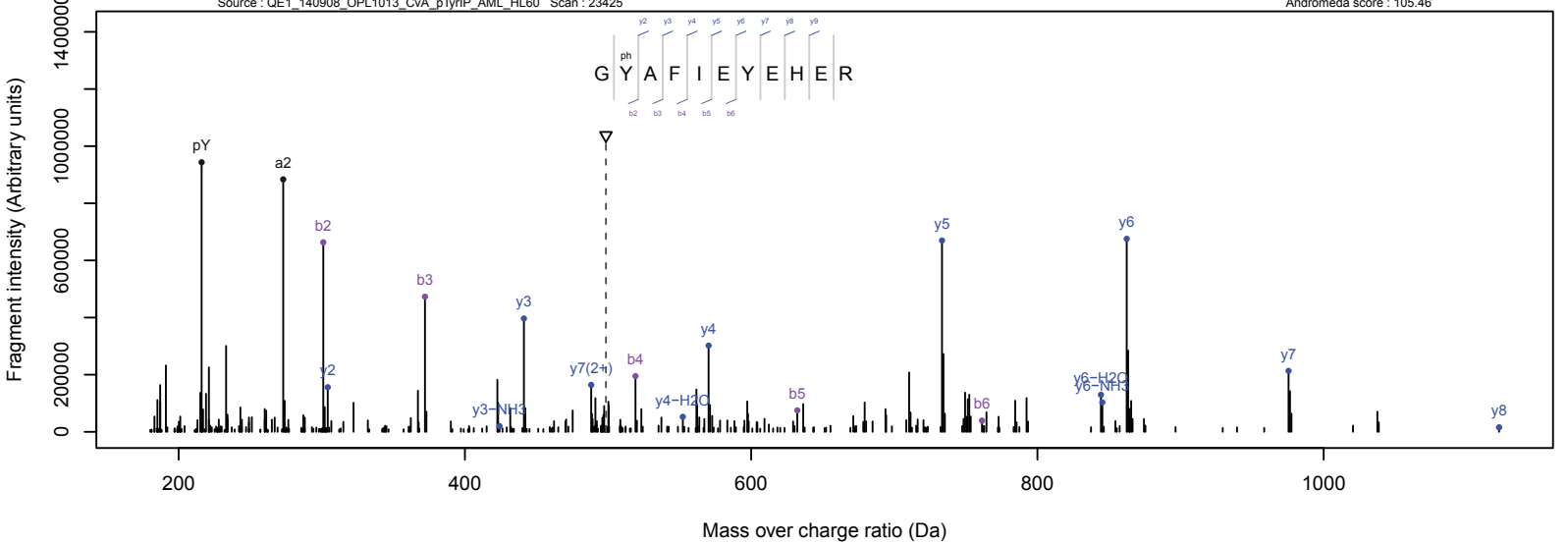
Andromeda score : 105.46



GY(ph)AFIEYEH-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 23425

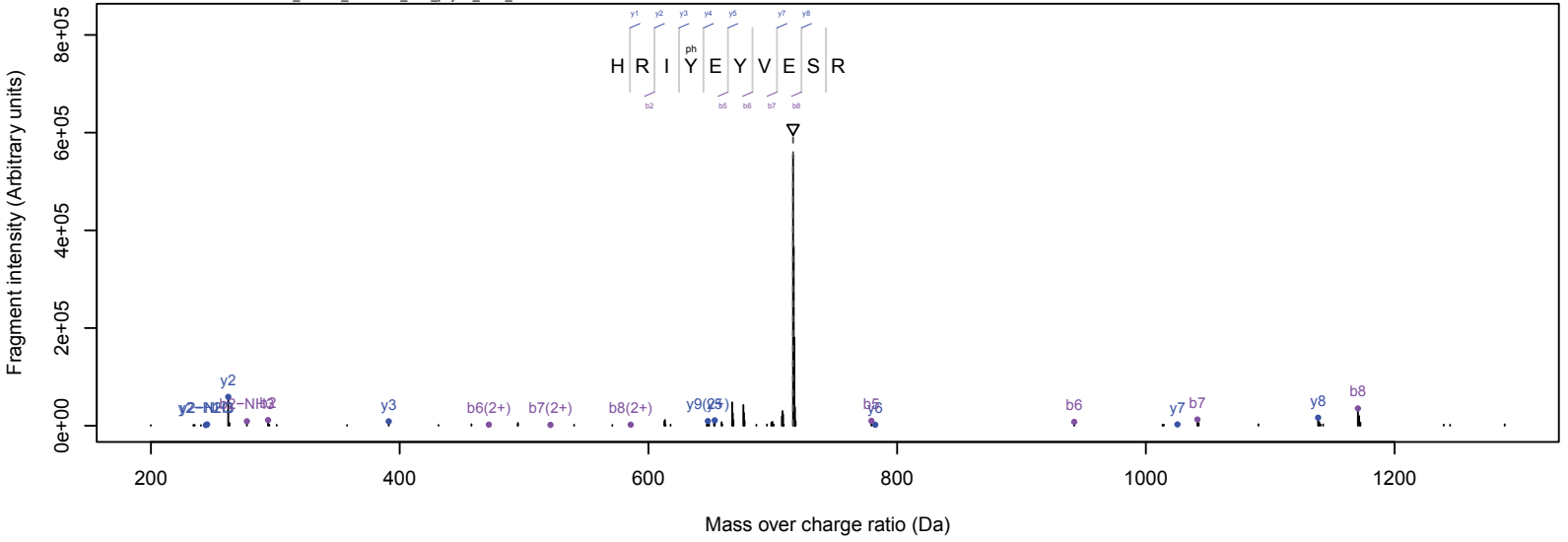
Andromeda score : 105.46



HRIY(ph)EYVESR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 9780

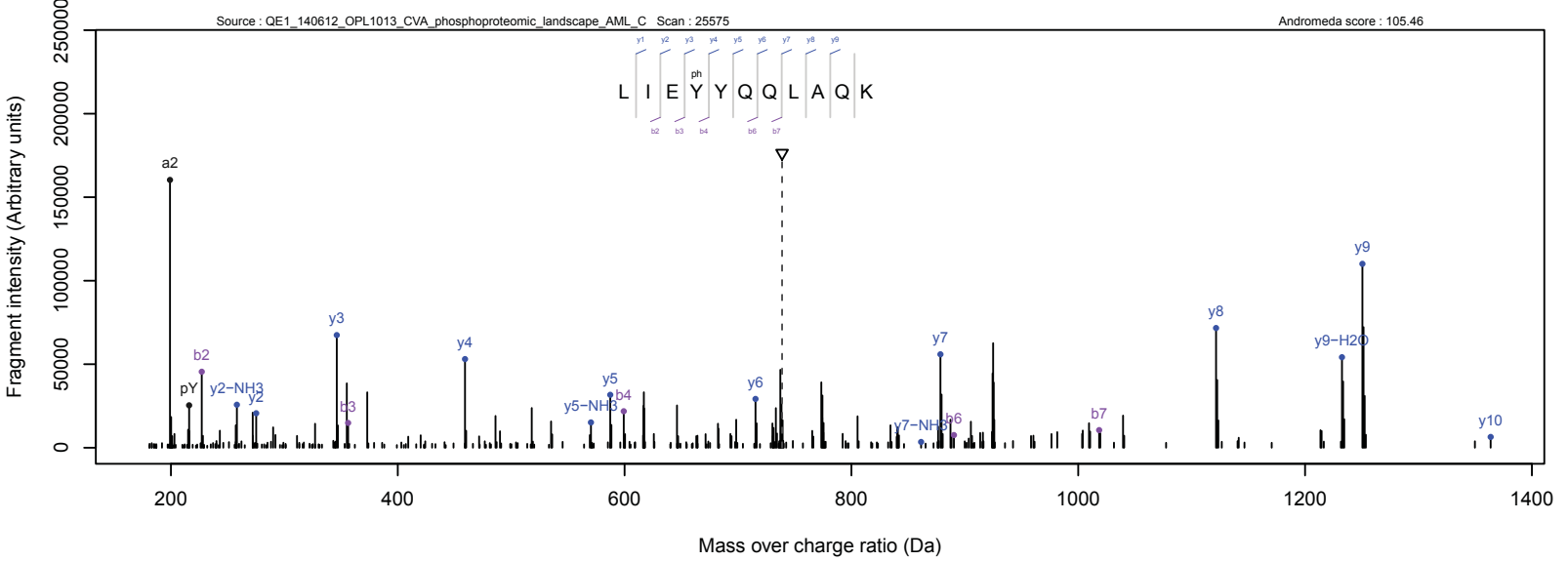
Andromeda score: 105.46



LIEY(ph)YQQLAQK-2+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 25575

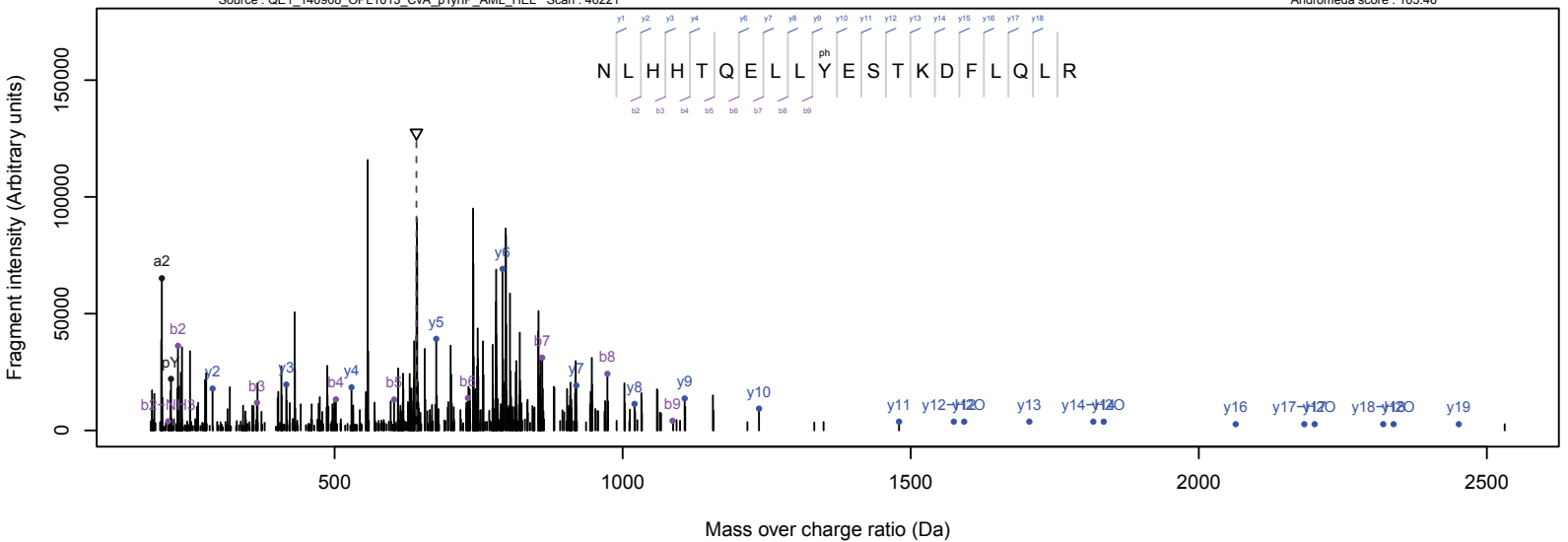
Andromeda score: 105.46



NLHHTQELLY(ph)ESTKDFLQLR-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 46221

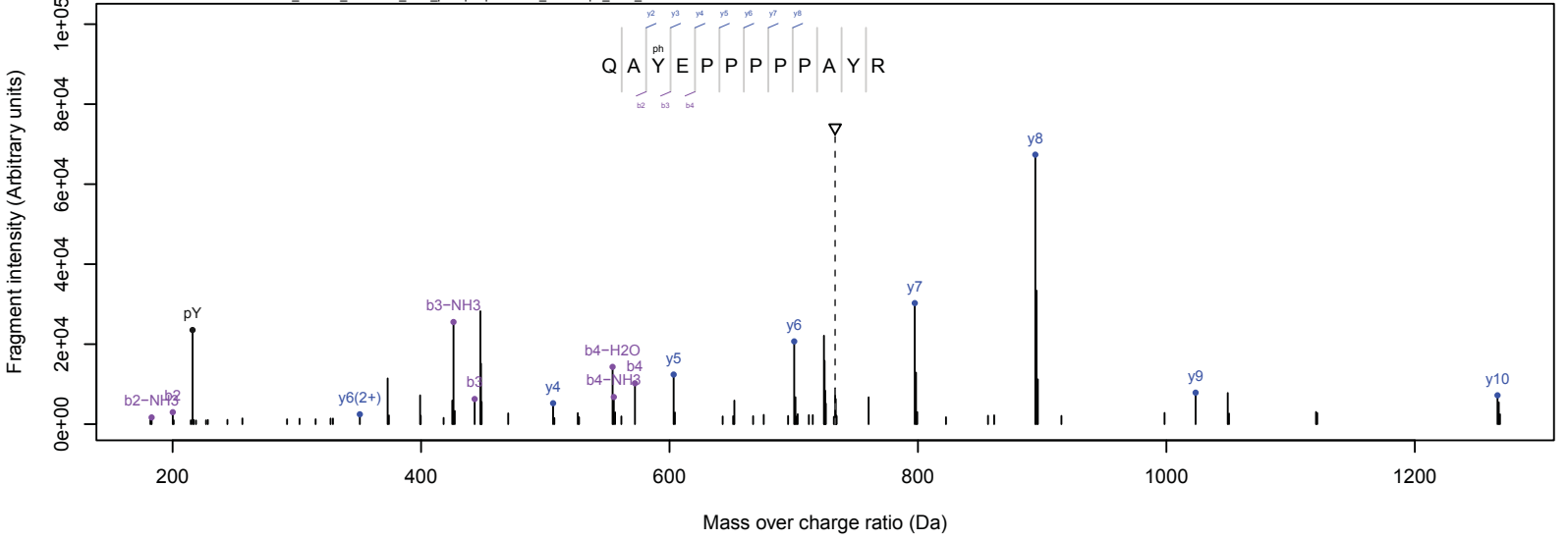
Andromeda score: 105.46



QAY(ph)EPPPPAYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14001

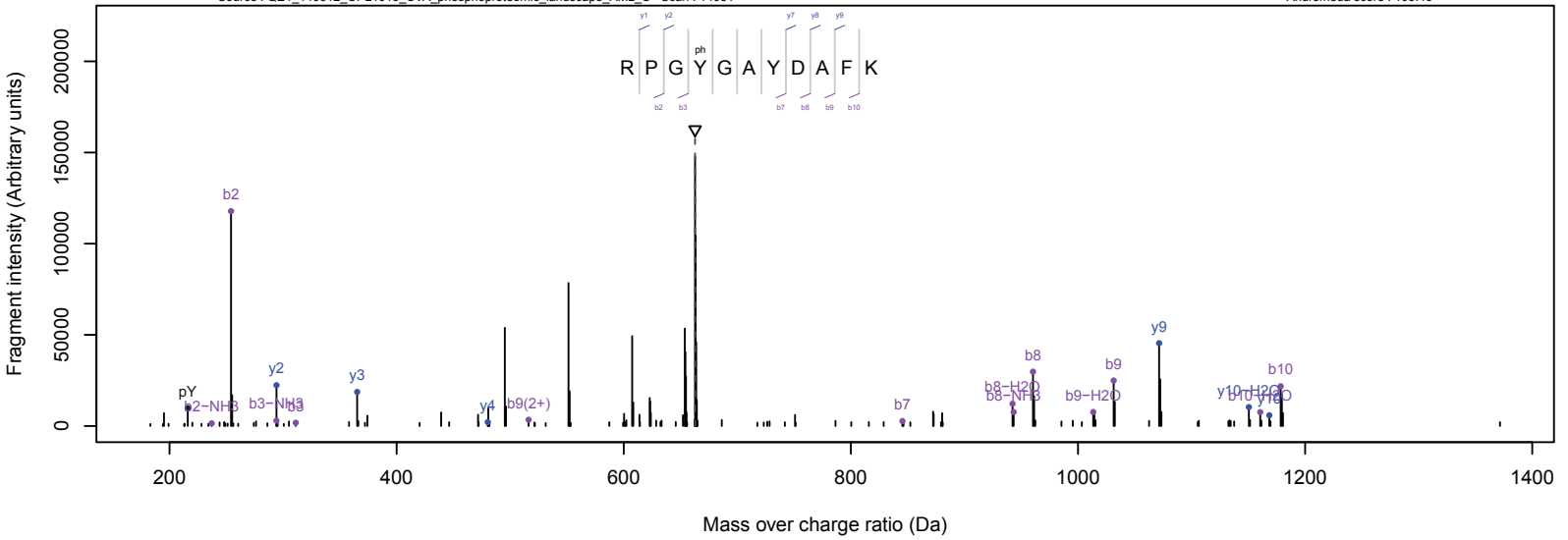
Andromeda score : 105.46



RPGY(ph)GAYDAFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14081

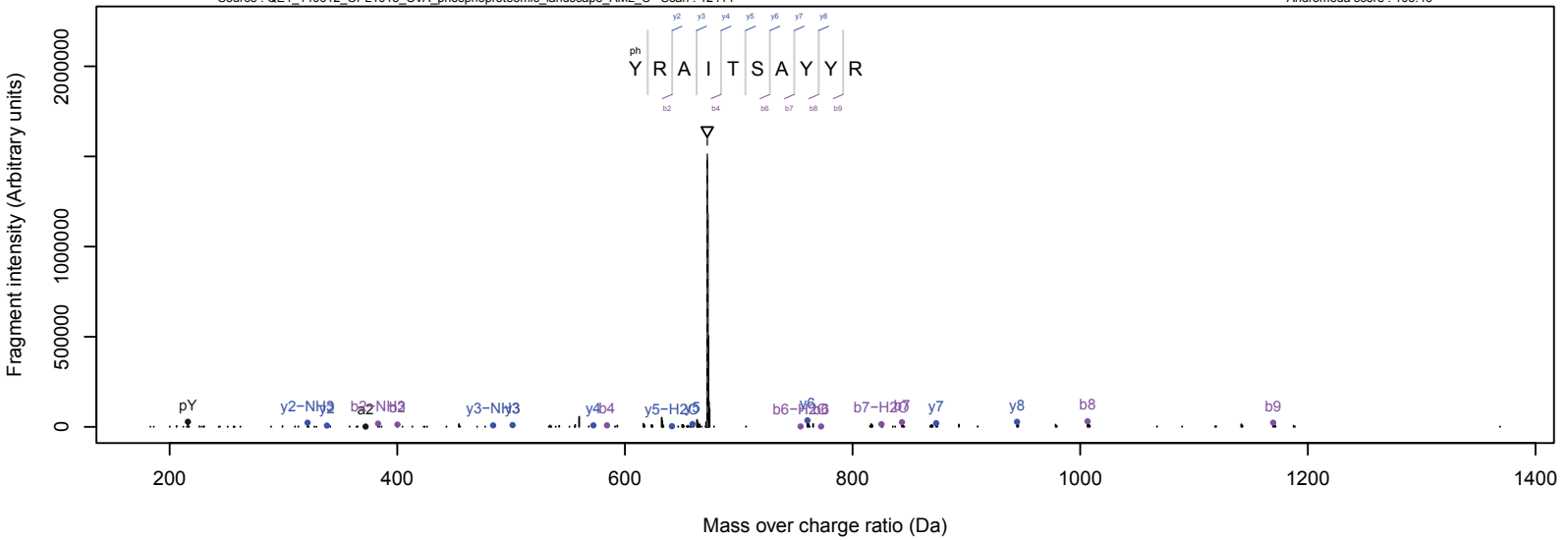
Andromeda score : 105.46



Y(ph)RAITSAYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12414

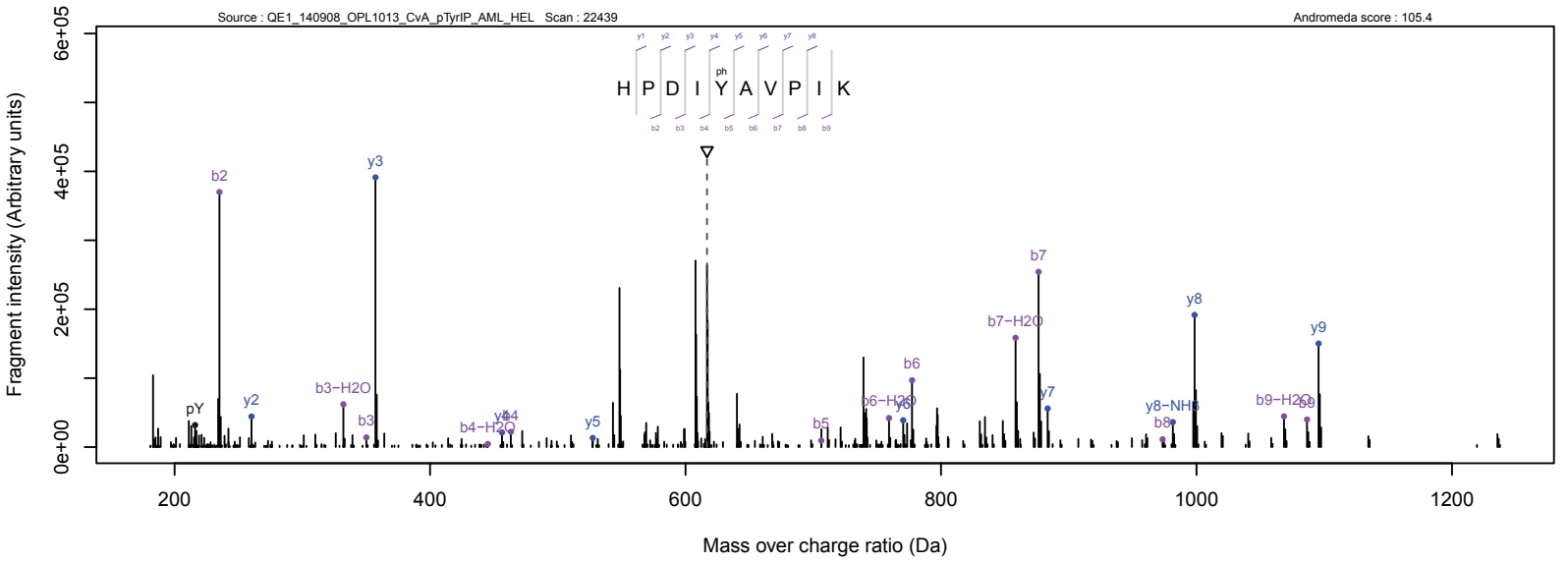
Andromeda score : 105.46



HPDIY(ph)AVPIK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 22439

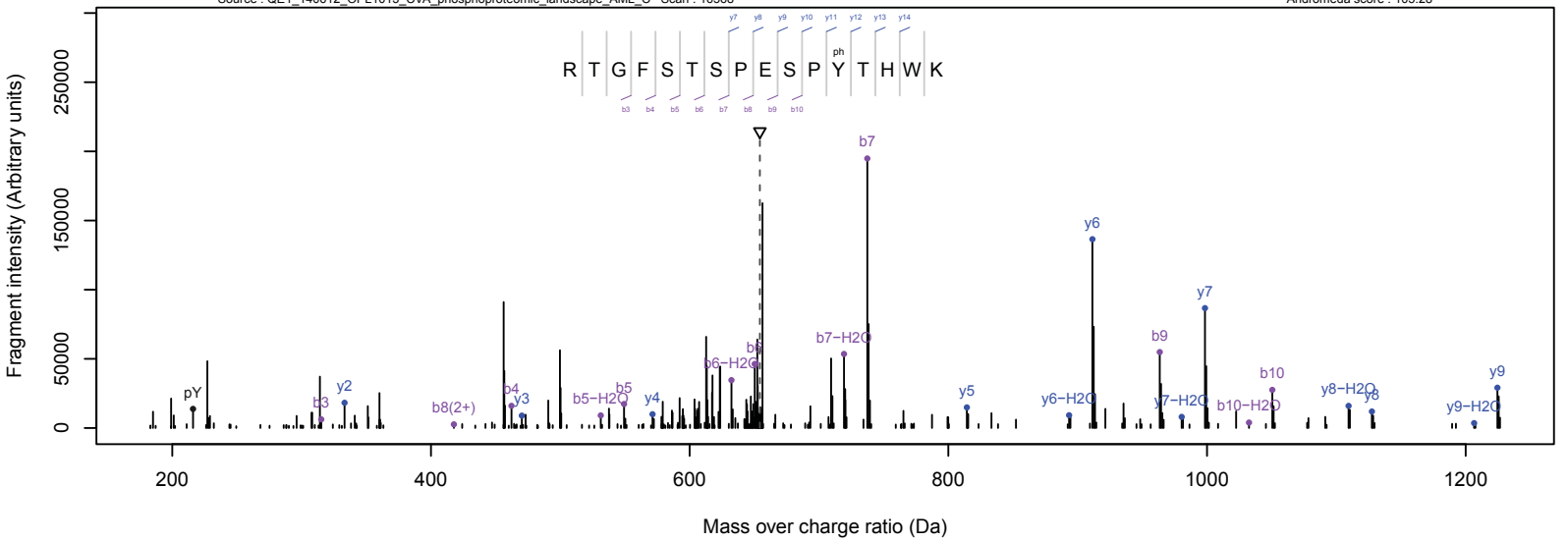
Andromeda score: 105.4



RTGFSTSPESPY(ph)THWK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16568

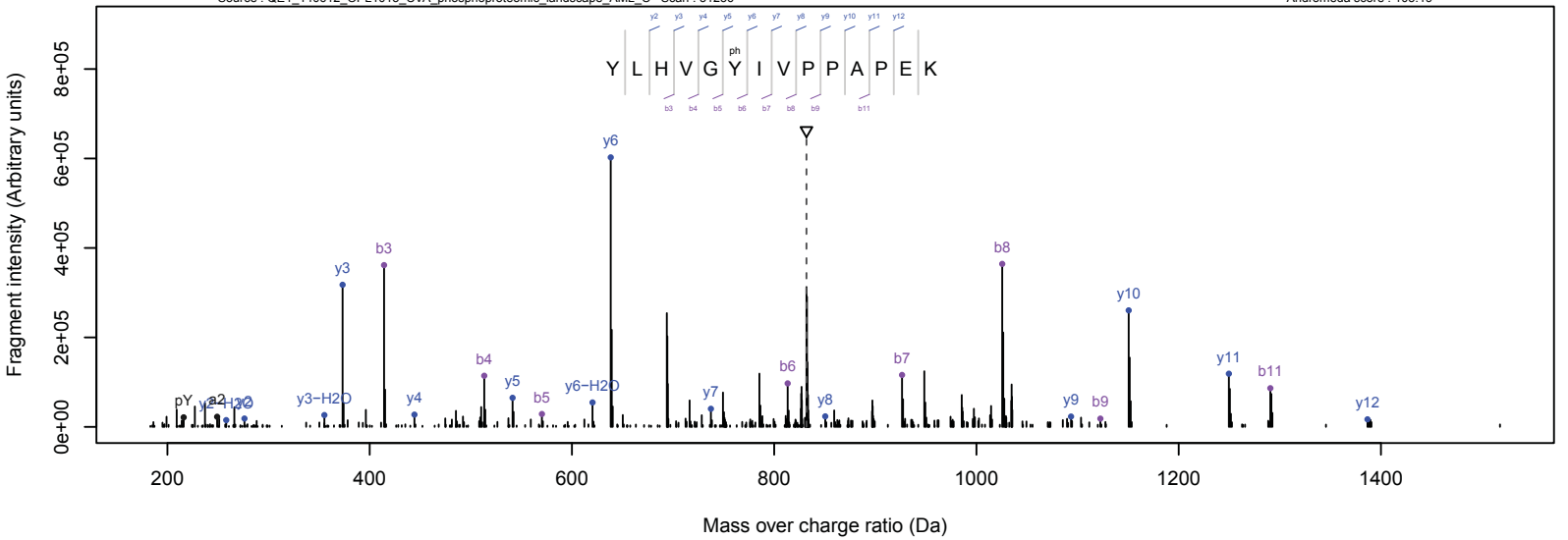
Andromeda score: 105.28



YLHVGYP(ph)IVPPAPEK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31256

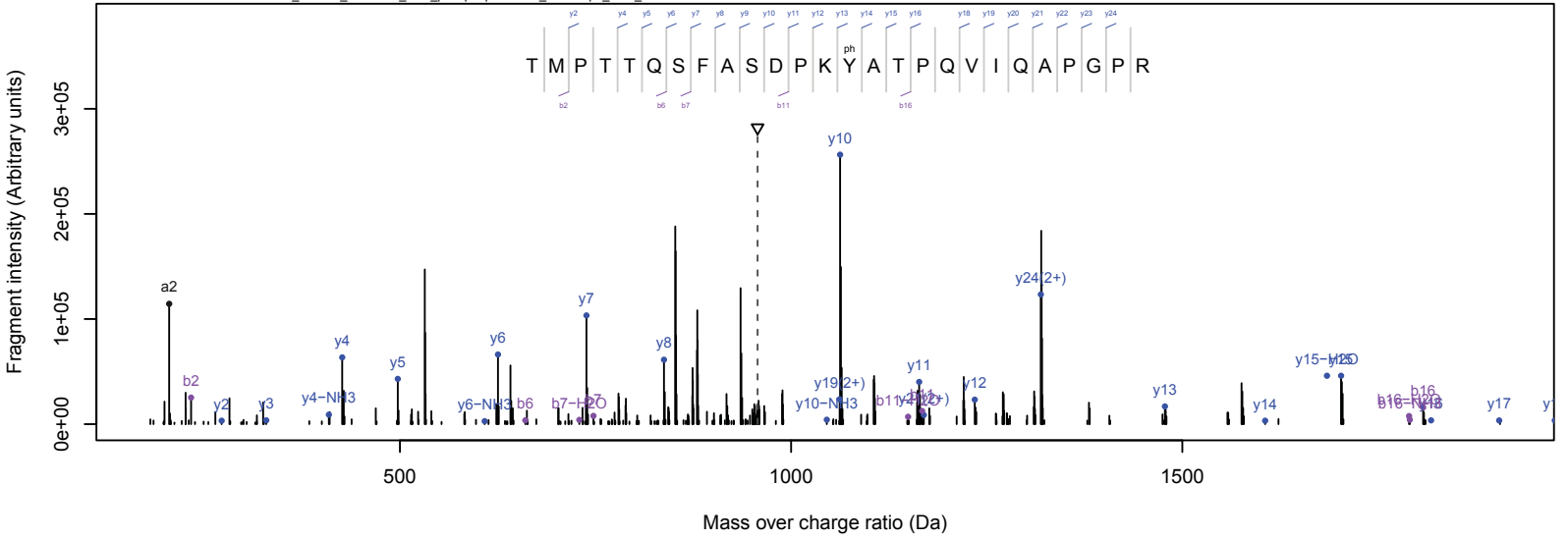
Andromeda score: 105.19



**TMPTTQSFASDPKY(ph)ATPQVIQAPGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 31853

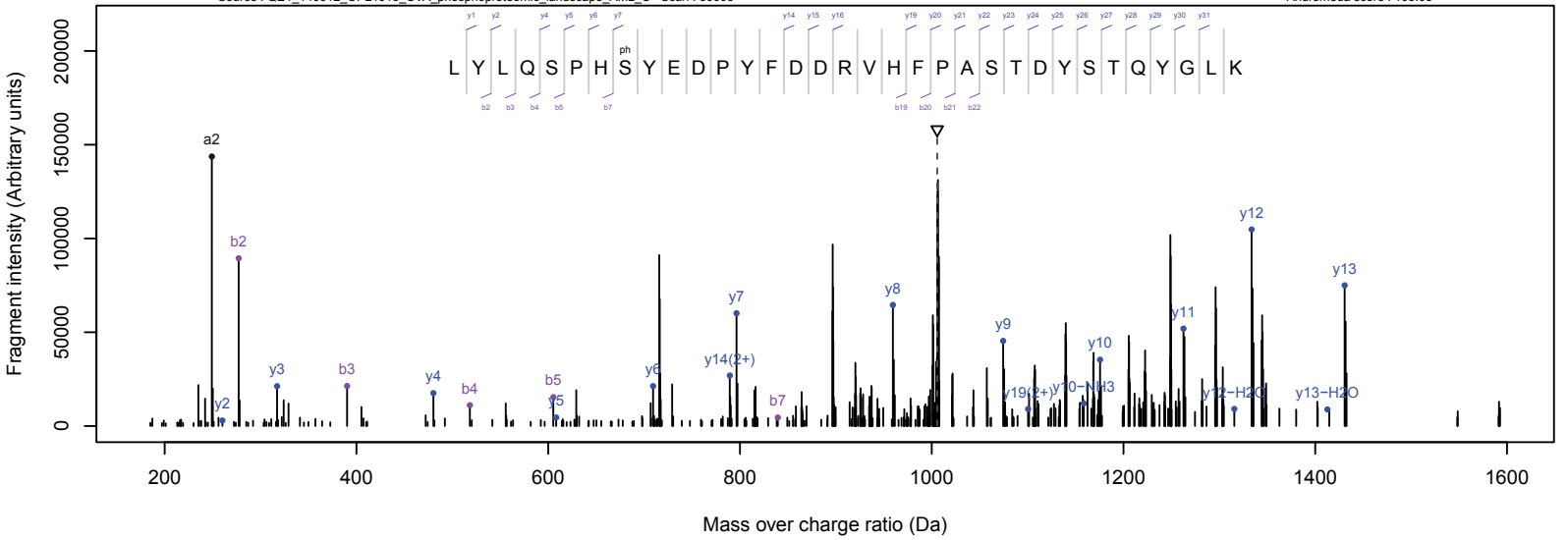
Andromeda score : 105.17



**LYLQSPHS(ph)YEDPYFDDR VHFPASTDYSTQYGLK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39095

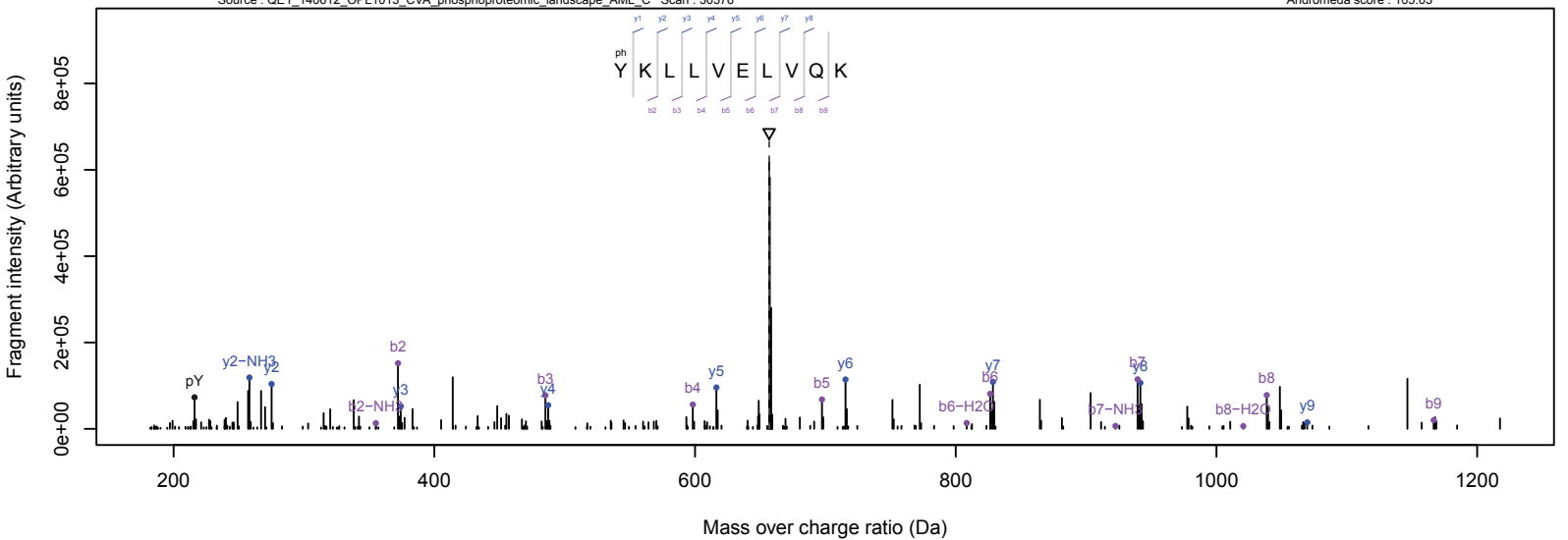
Andromeda score : 105.05



**Y(ph)KLLVELVQK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30576

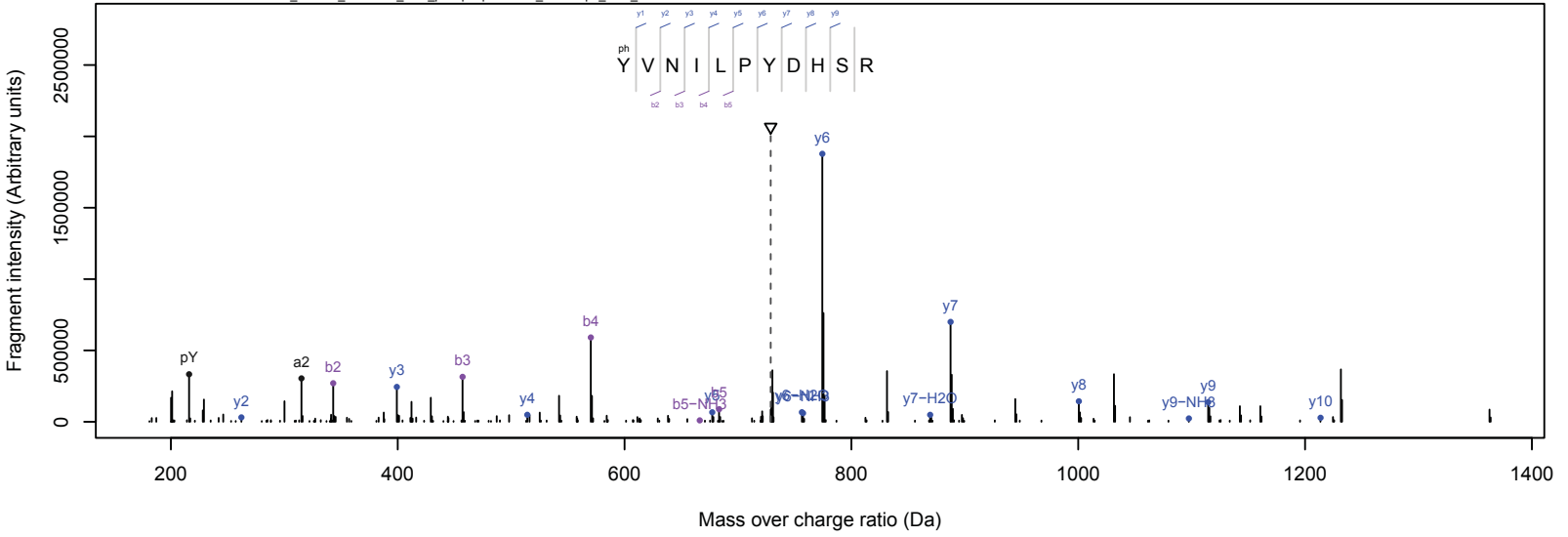
Andromeda score : 105.03



Y(ph)VNILPYDHSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25251

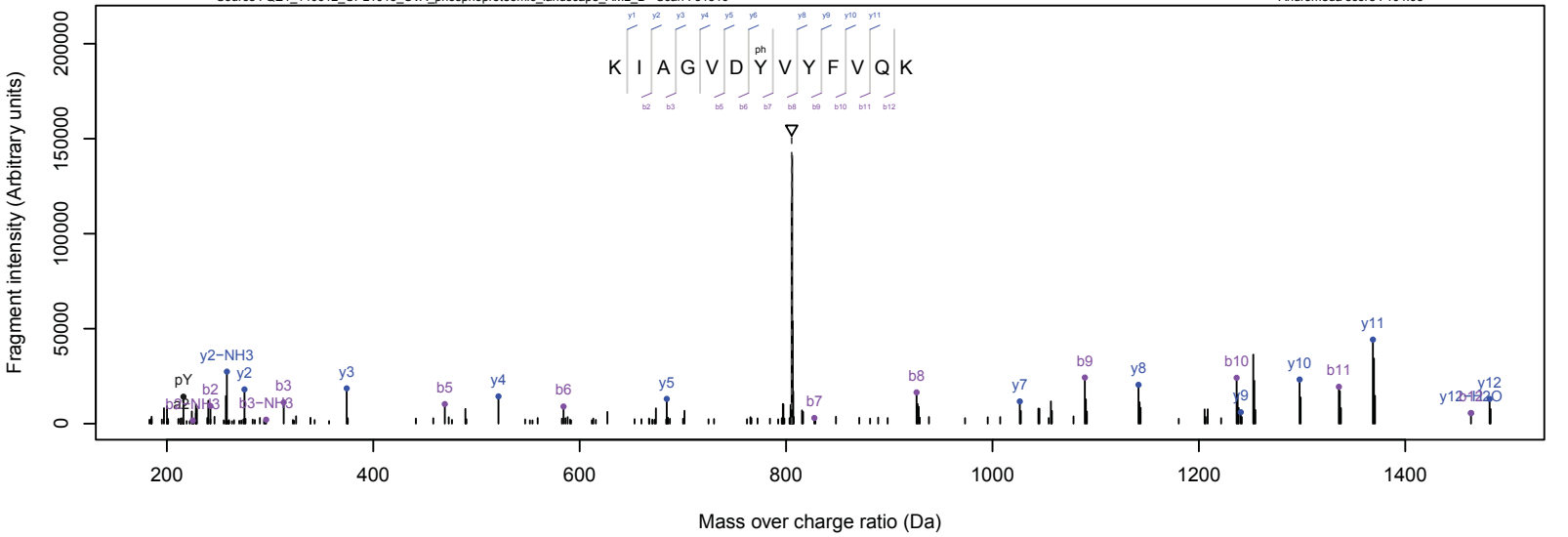
Andromeda score : 105.03



KIAGVDY(ph)VYFVQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31510

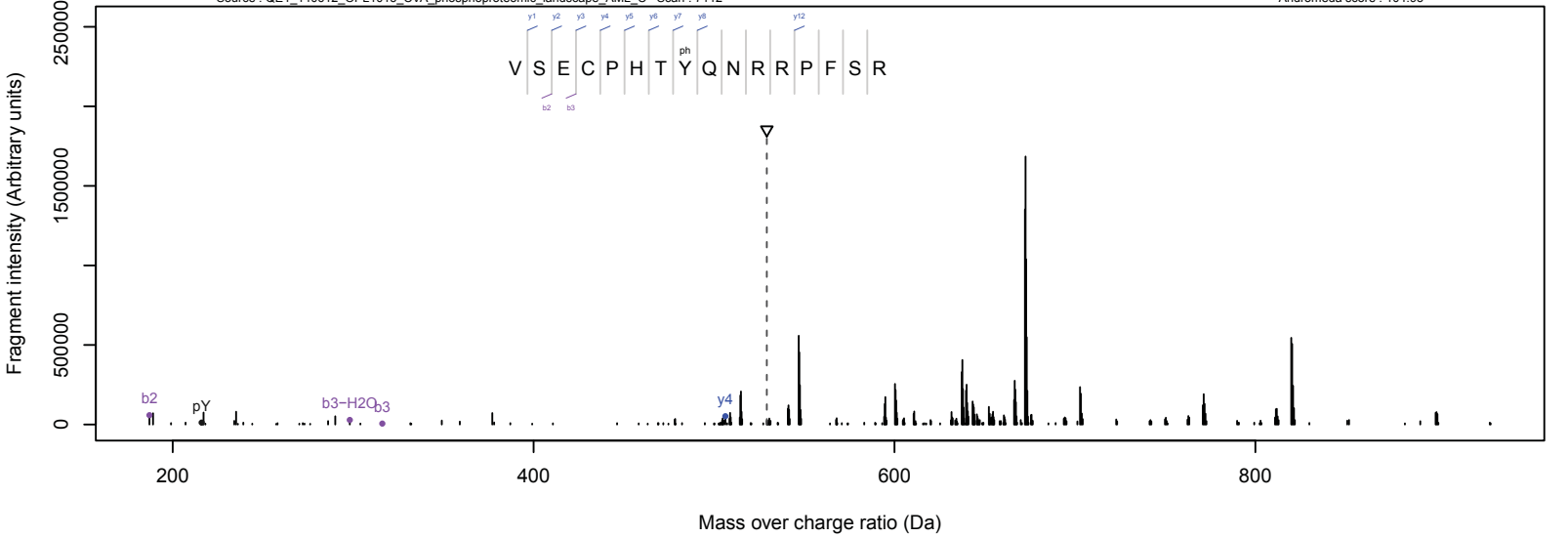
Andromeda score : 104.95



VSECPHTY(ph)QNRPF SR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7112

Andromeda score : 104.95

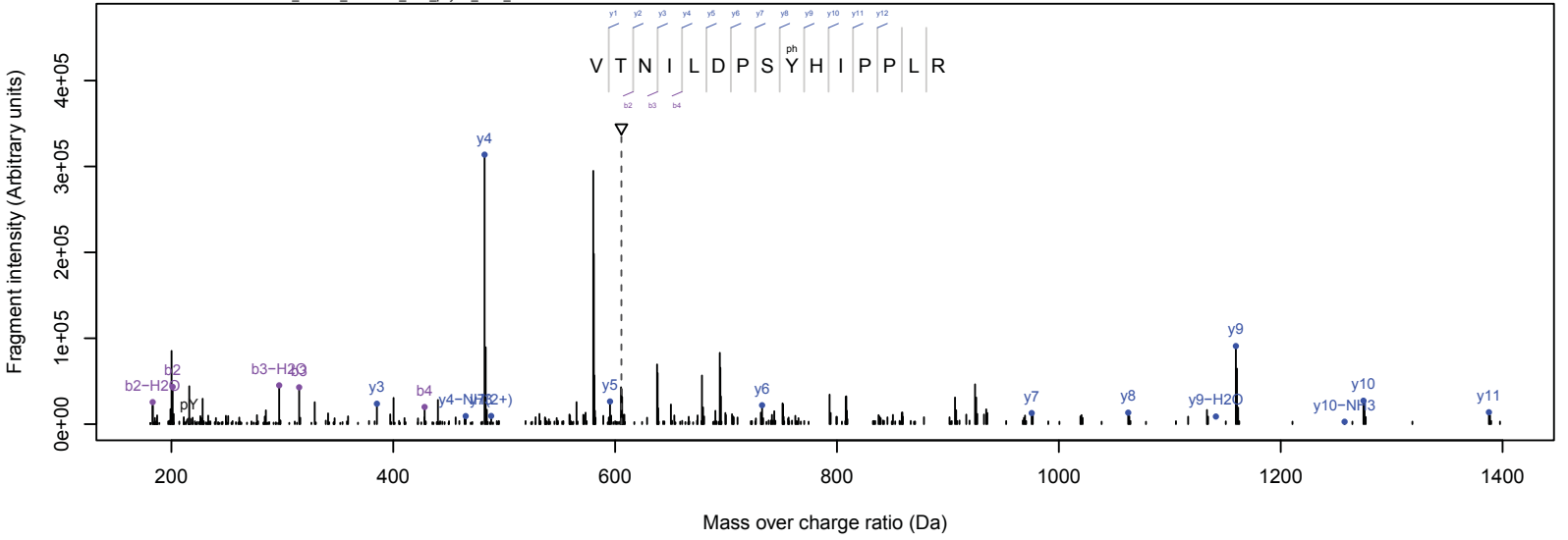




**VTNILDPSY(ph)HIPPLR-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 38046

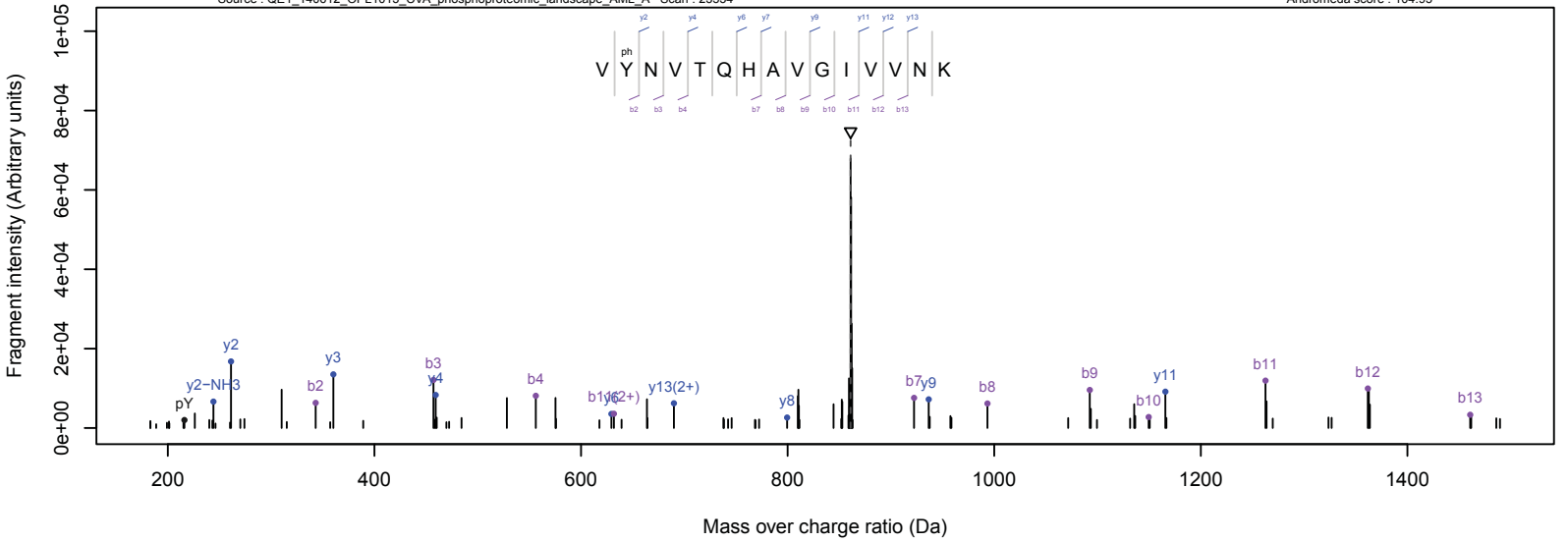
Andromeda score: 104.95



**VY(ph)NVTQHAVGIVVNK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 23534

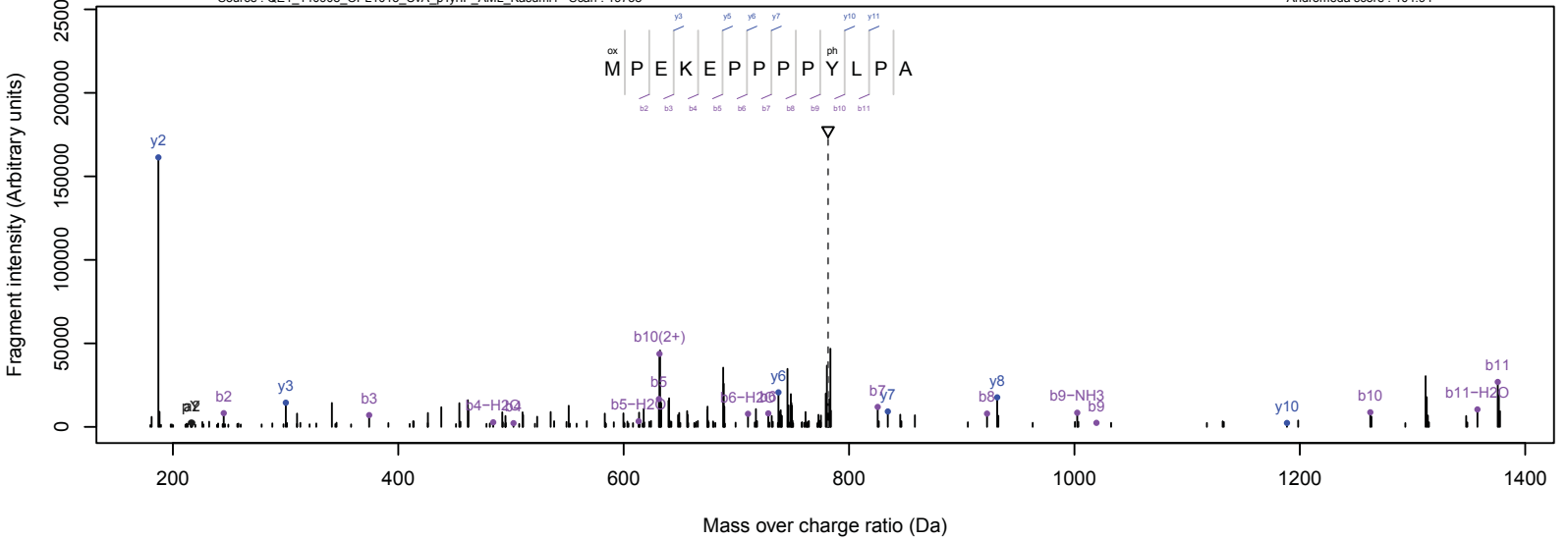
Andromeda score: 104.95



**M(ox)PEKEPPPY(ph)LPA-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 19738

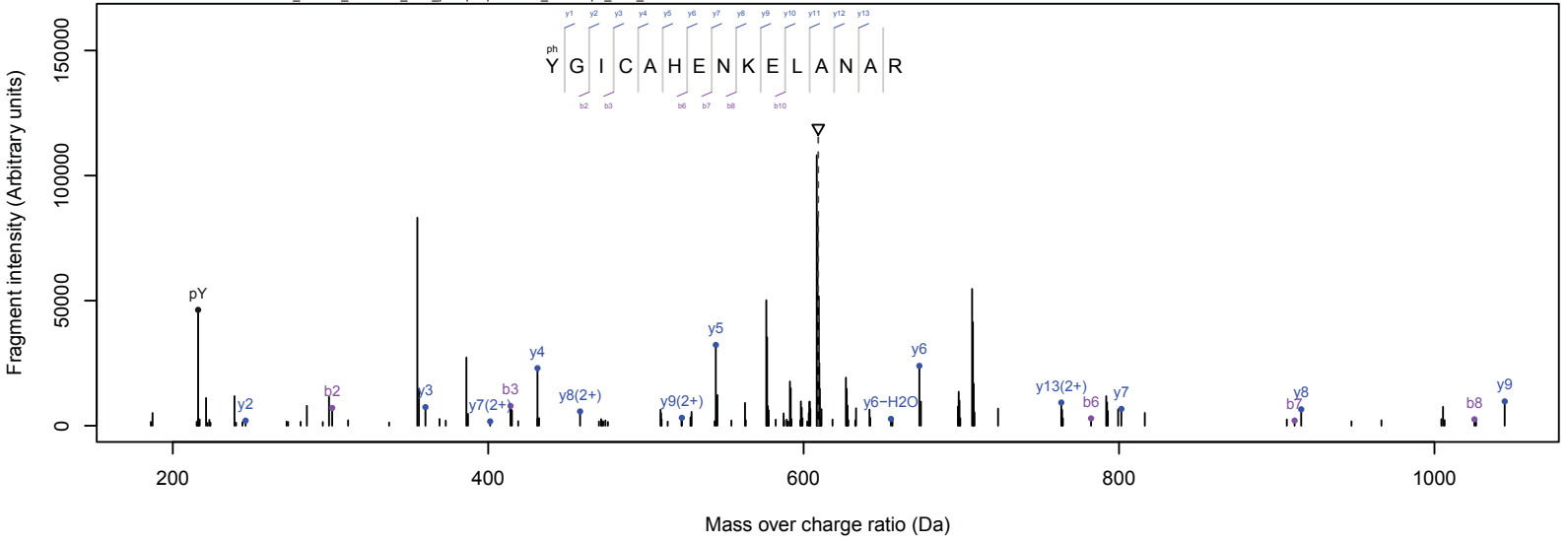
Andromeda score: 104.94



**Y(ph)GICAHENKELANAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9076

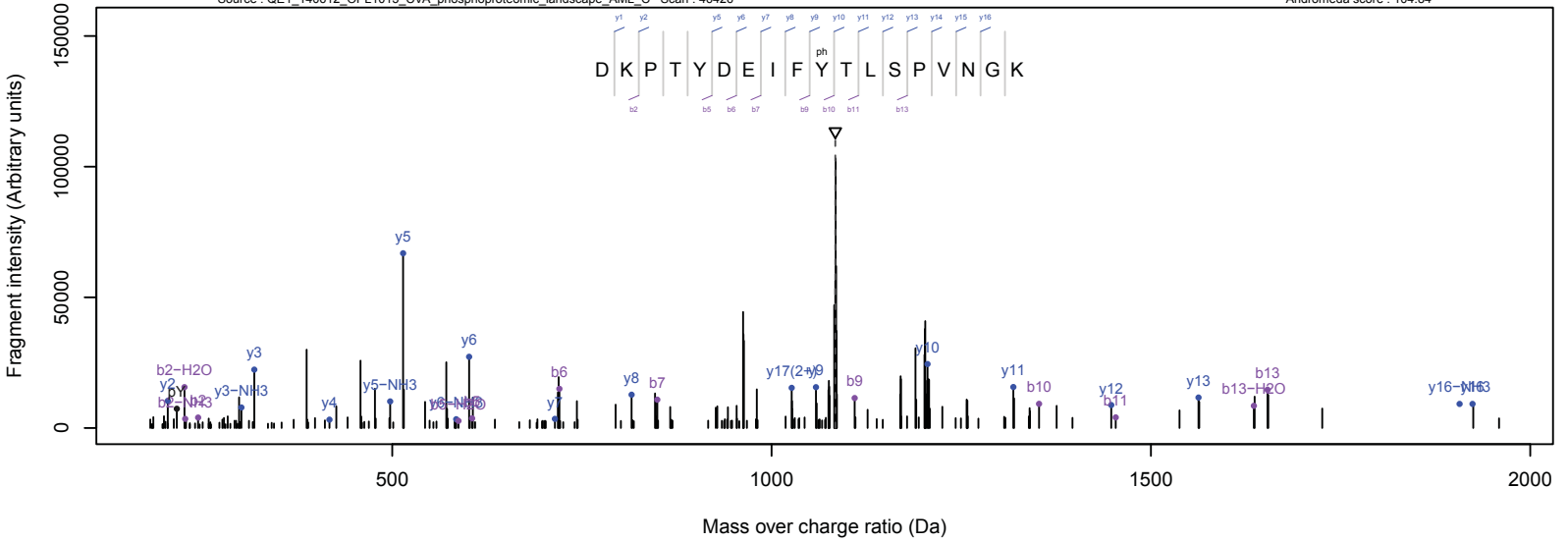
Andromeda score : 104.94



**DKPTYDEIFY(ph)TLSPVNGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46426

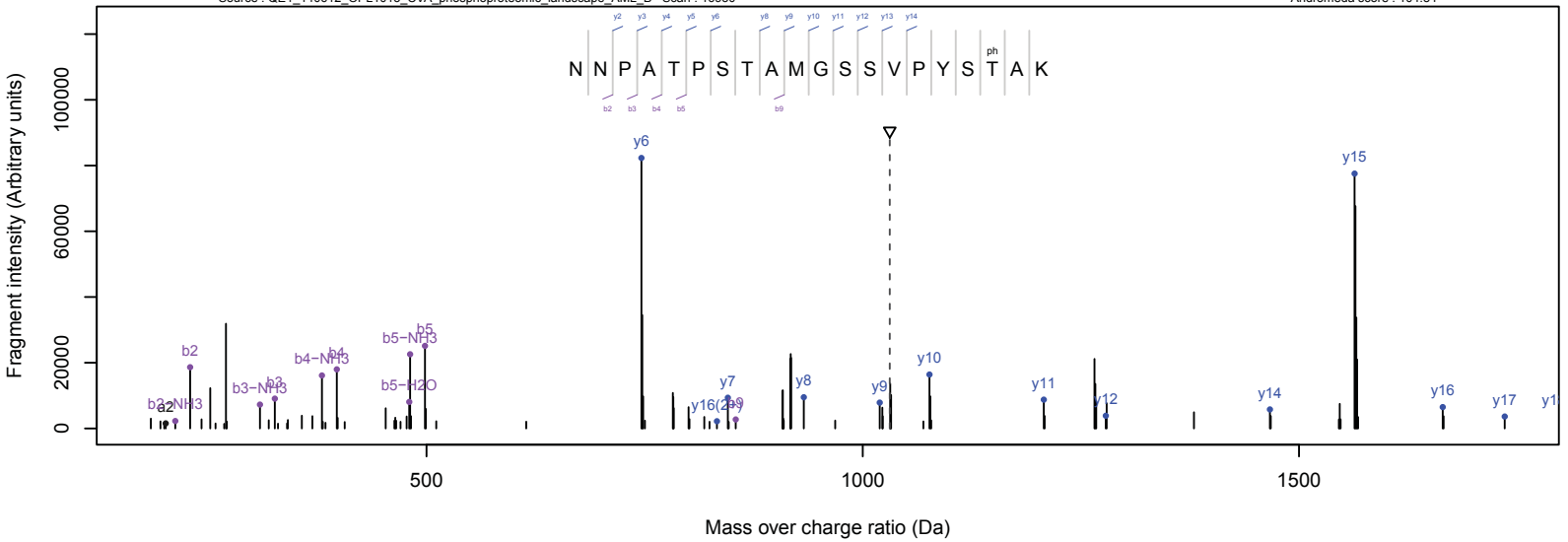
Andromeda score : 104.84



**NNPATPSTAMGSSVPYSTAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16930

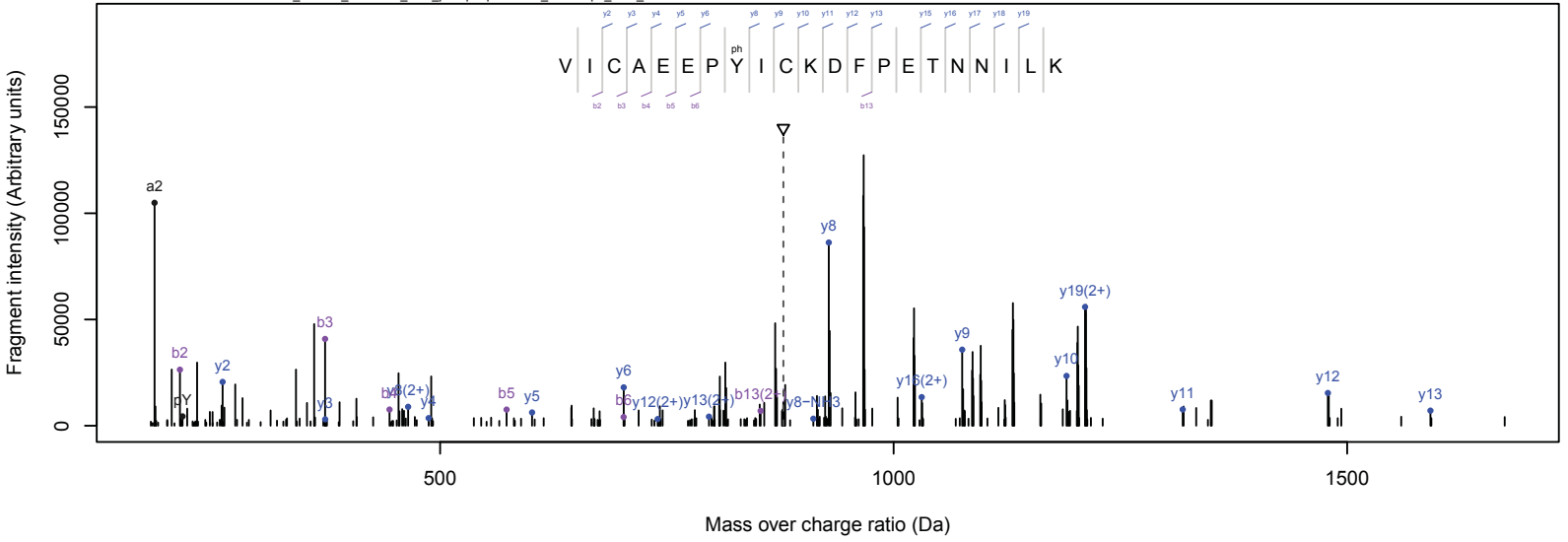
Andromeda score : 104.84



VICAEEPY(ph)ICKDFPETNNILK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 38409

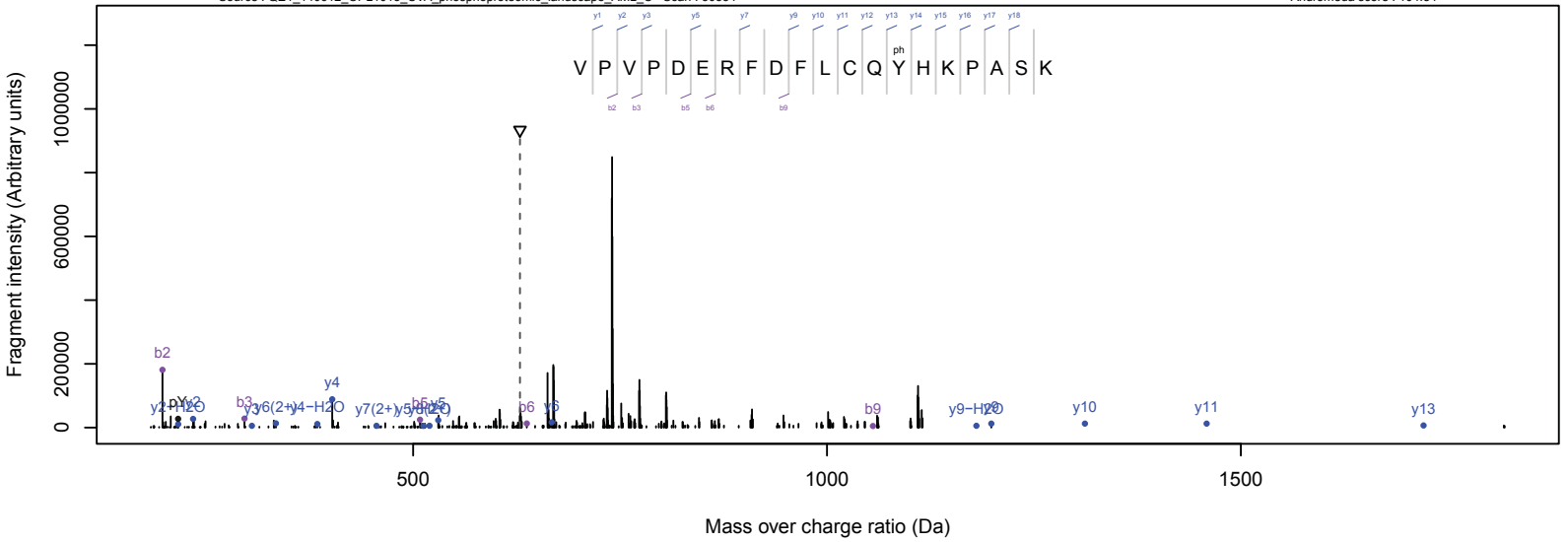
Andromeda score : 104.84



VPVDERFDLFCQ(ph)HKPASK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30354

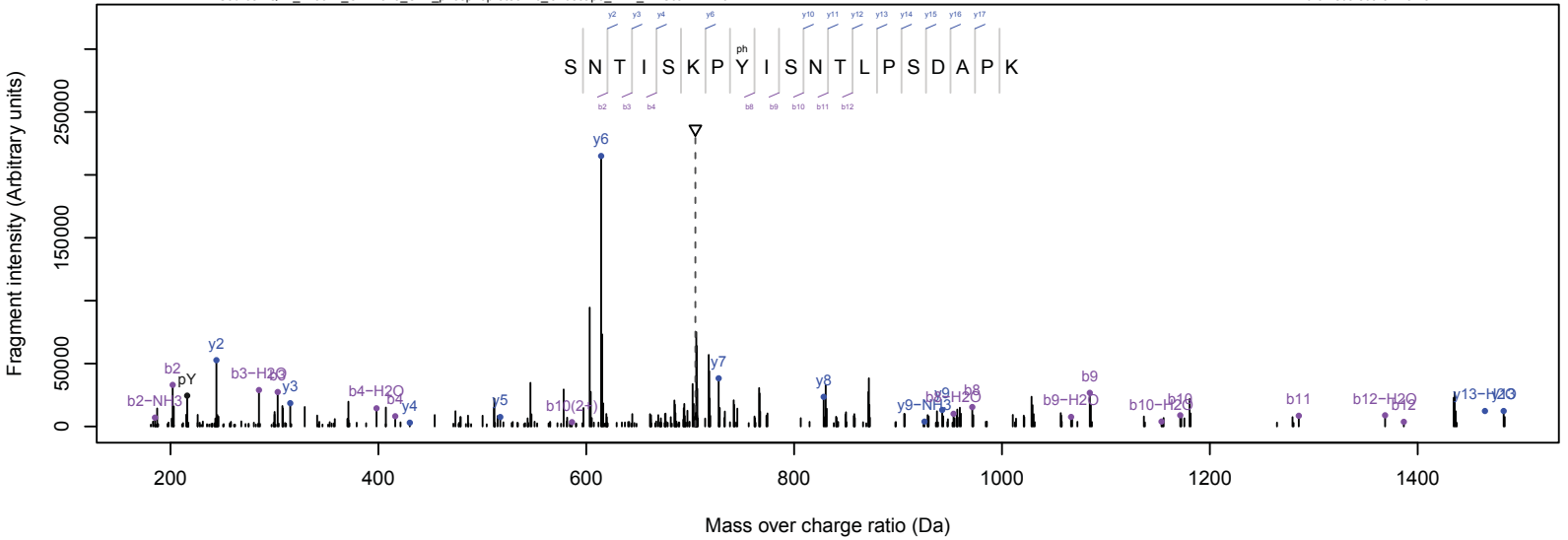
Andromeda score : 104.84



SNTISKPY(ph)ISNTLPSDAPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22702

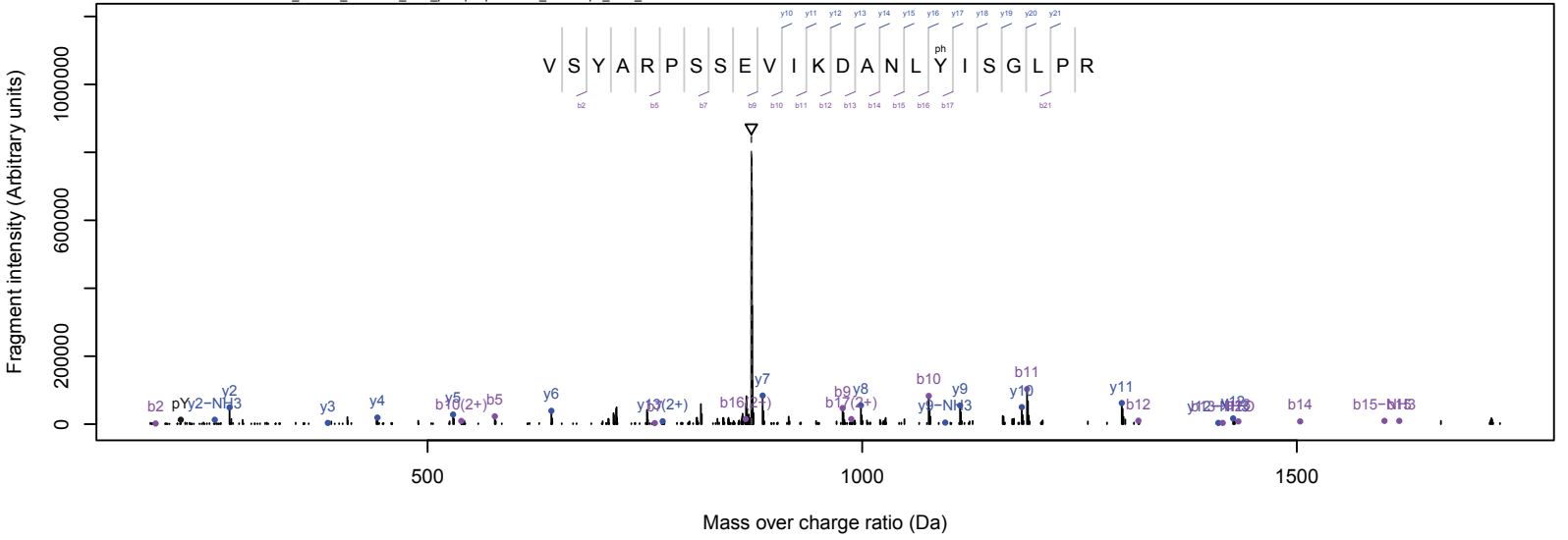
Andromeda score : 104.81



VSYARPSSEVIK(DANLY)ISGLPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34596

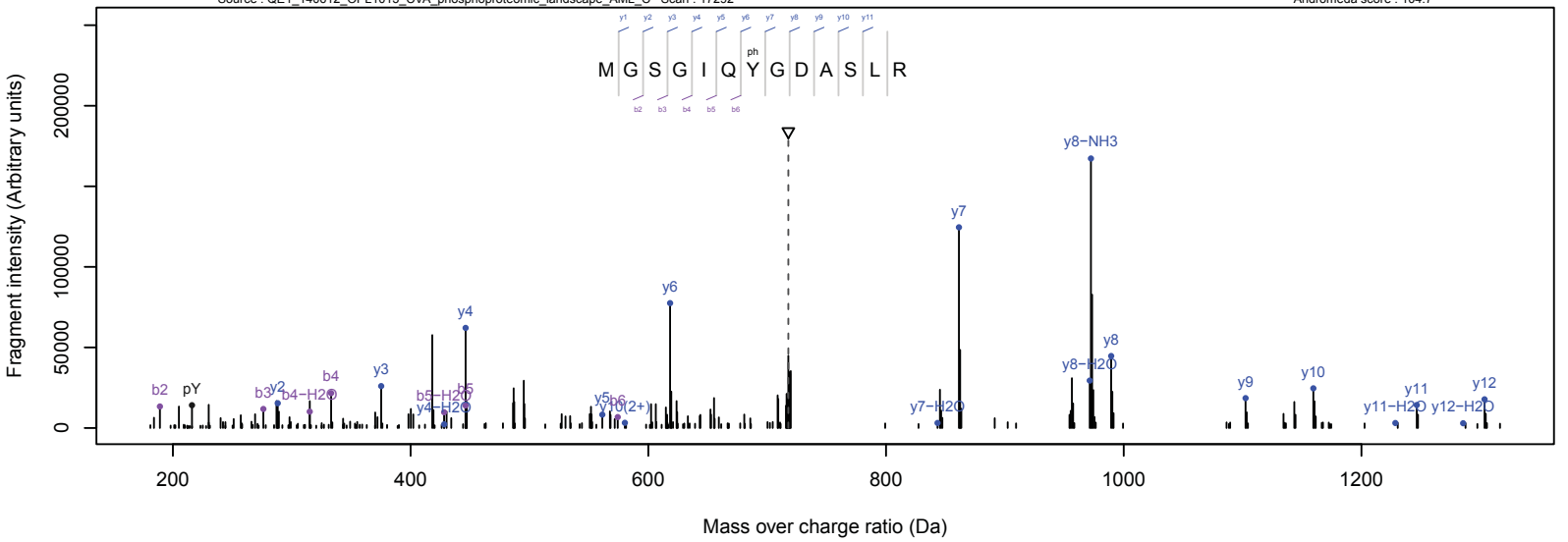
Andromeda score : 104.81



MGSGIQY(GDASLR)-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17292

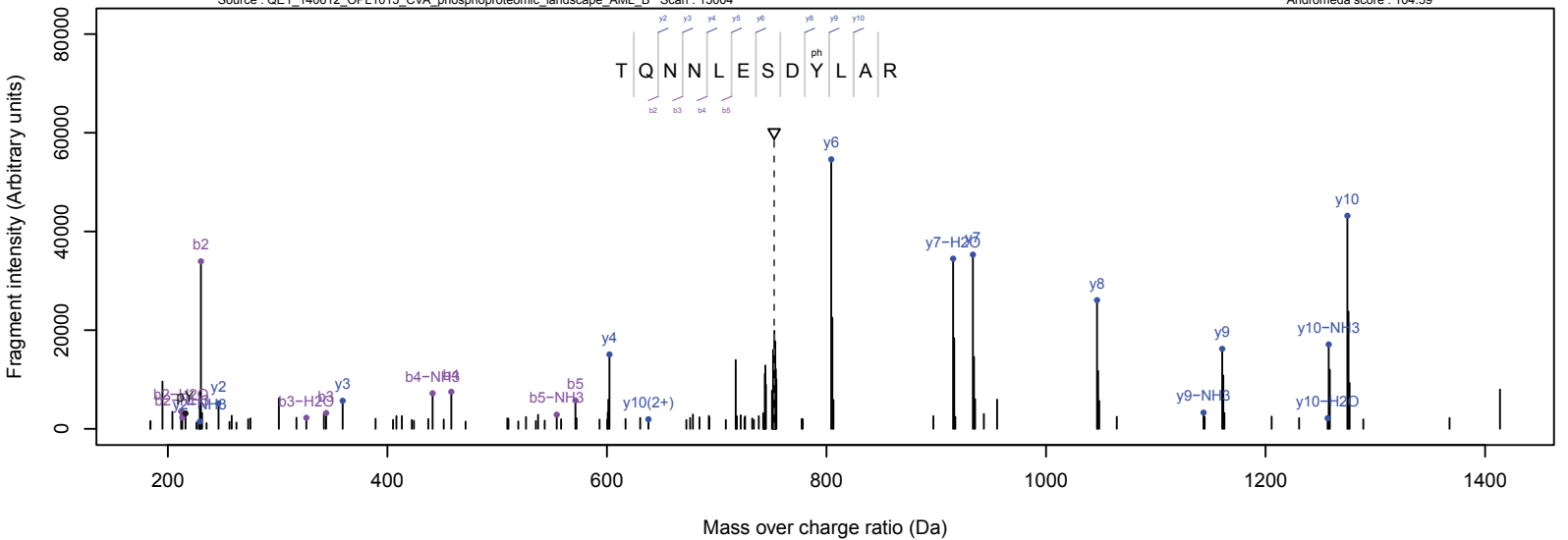
Andromeda score : 104.7



TQNNLESDY(LAR)-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15004

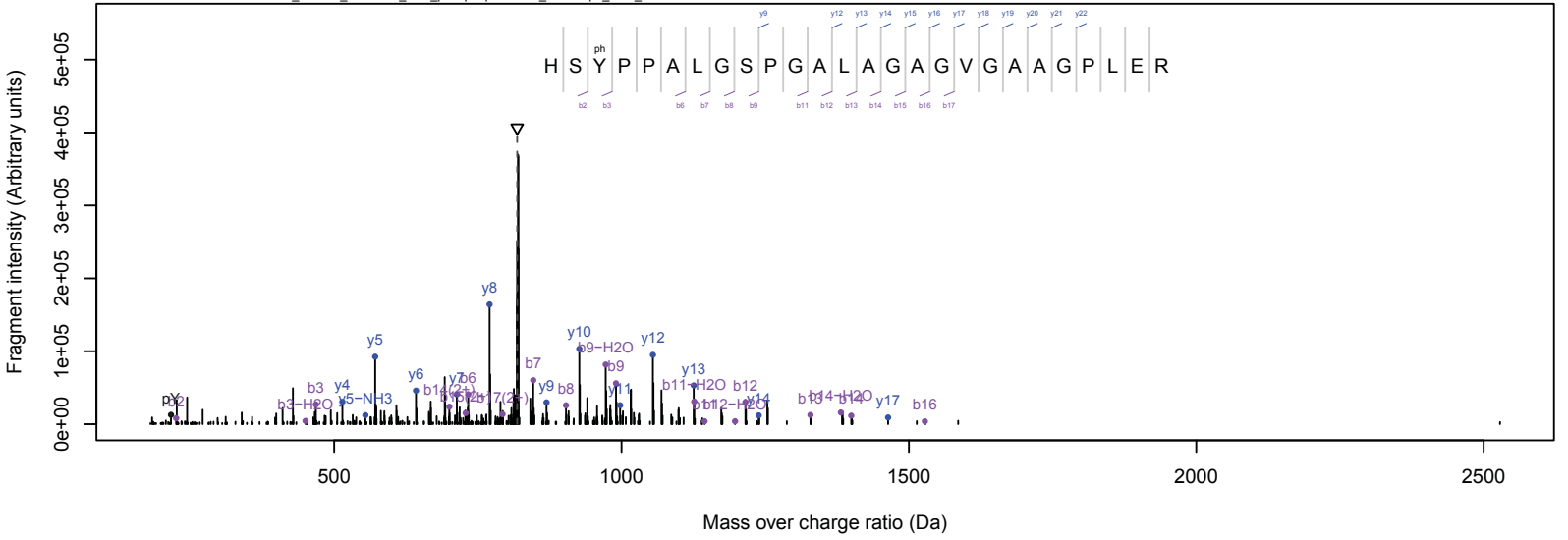
Andromeda score : 104.59



HSY(ph)PPALGSPGALAGAGVGAAGPLER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35576

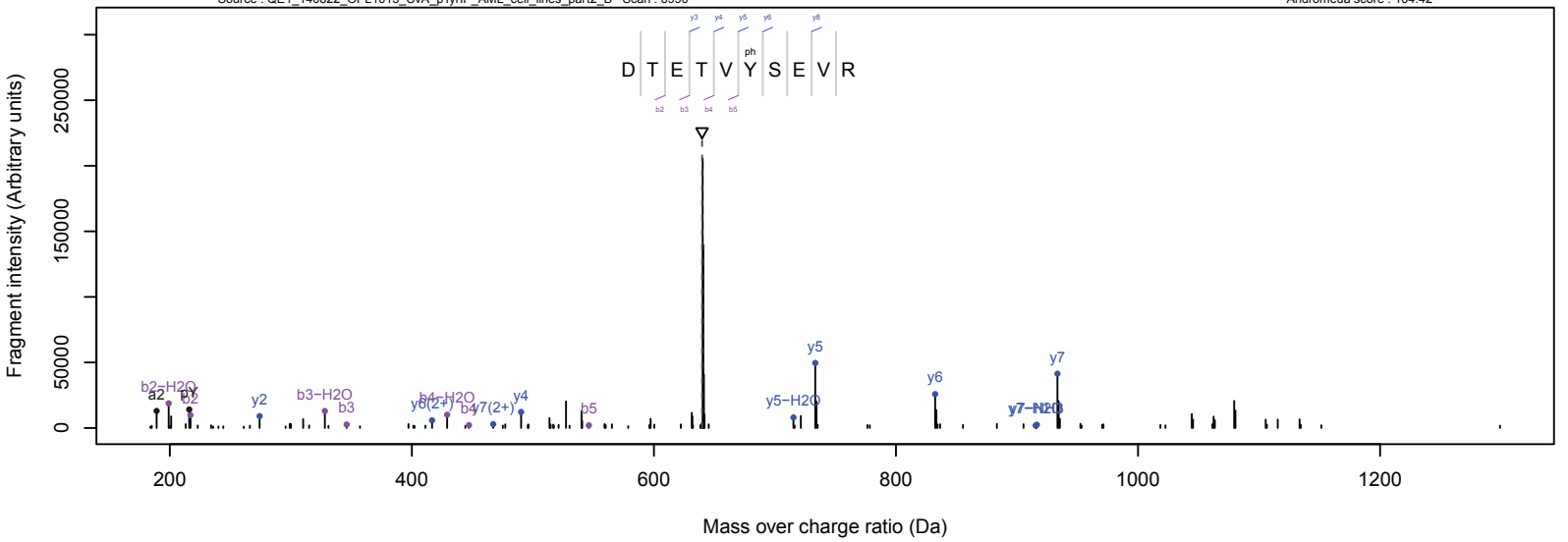
Andromeda score : 104.53



DTETVY(ph)SEVR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 6990

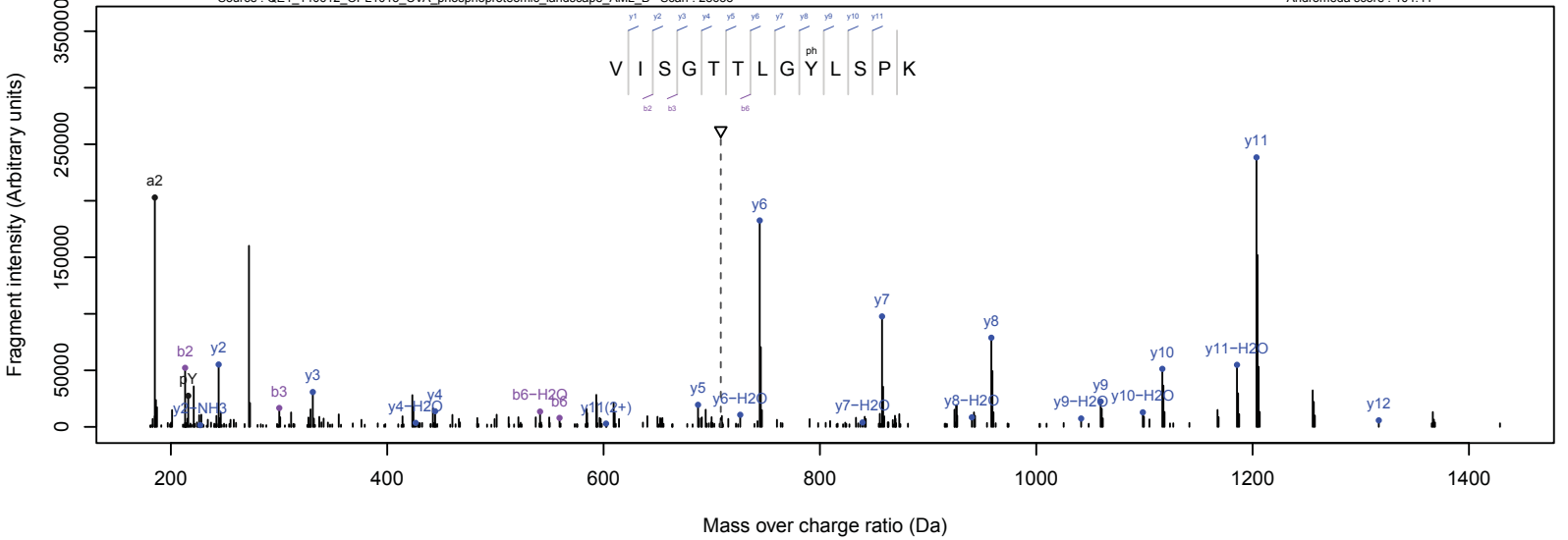
Andromeda score : 104.42



VISGTTLGYPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28658

Andromeda score : 104.41

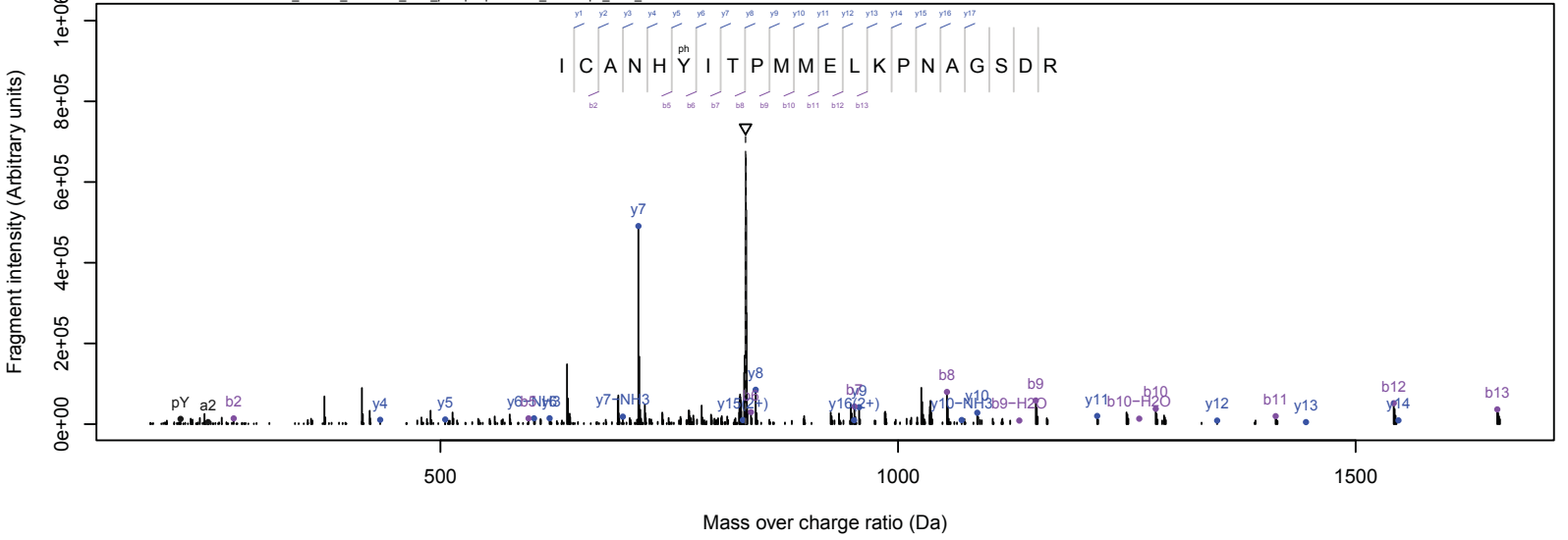




ICANH<sup>(ph)</sup>ITPMMELKPNAGSDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31544

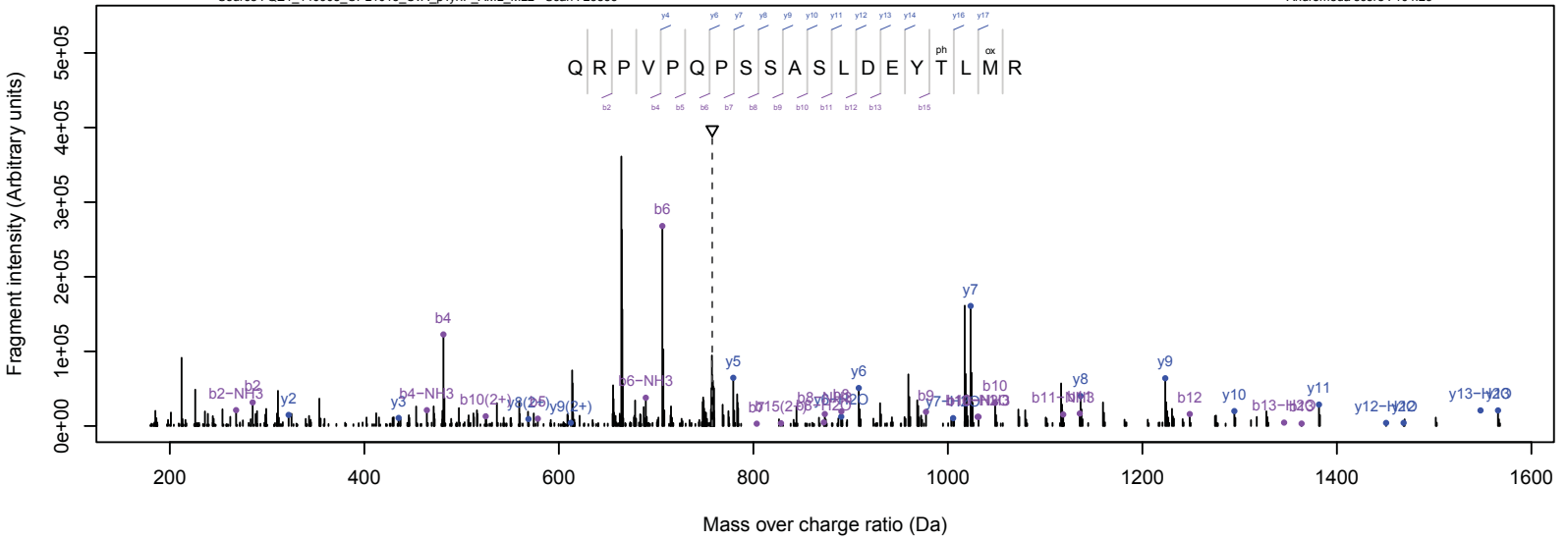
Andromeda score : 104.23



QRVPQPSSASLDEY<sup>(ph)</sup>LM<sup>(ox)</sup>R-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 23855

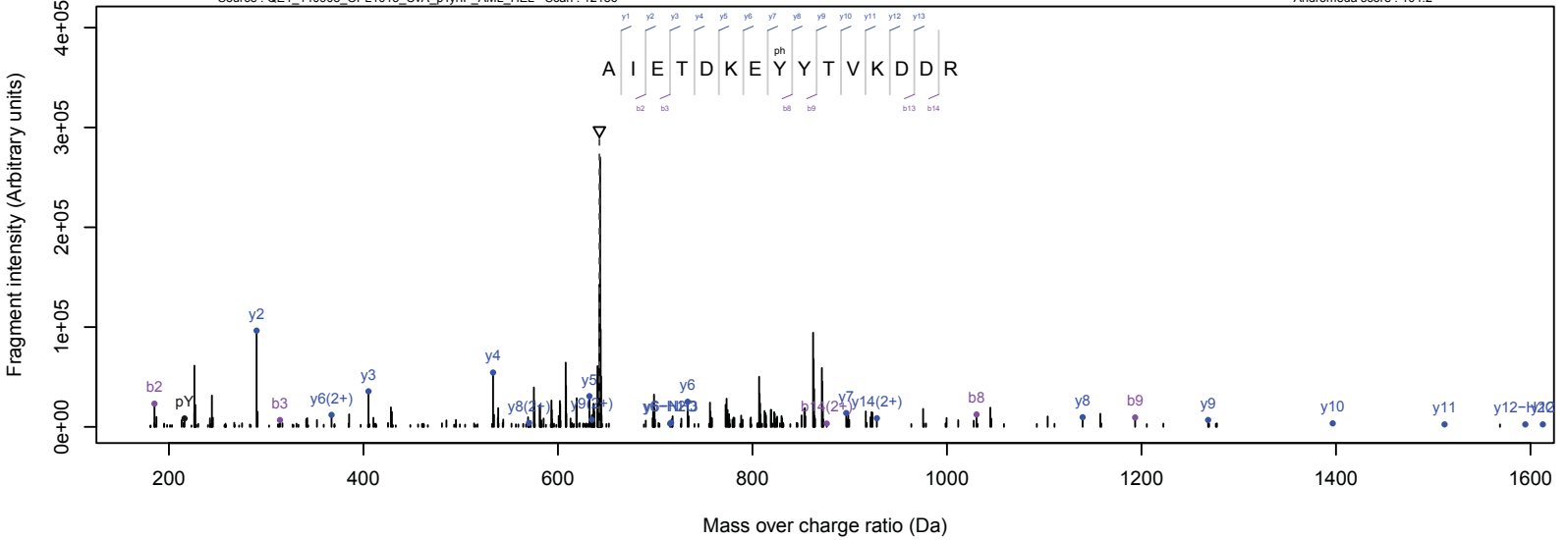
Andromeda score : 104.23



AIETDKEY<sup>(ph)</sup>YTVKDDR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 12186

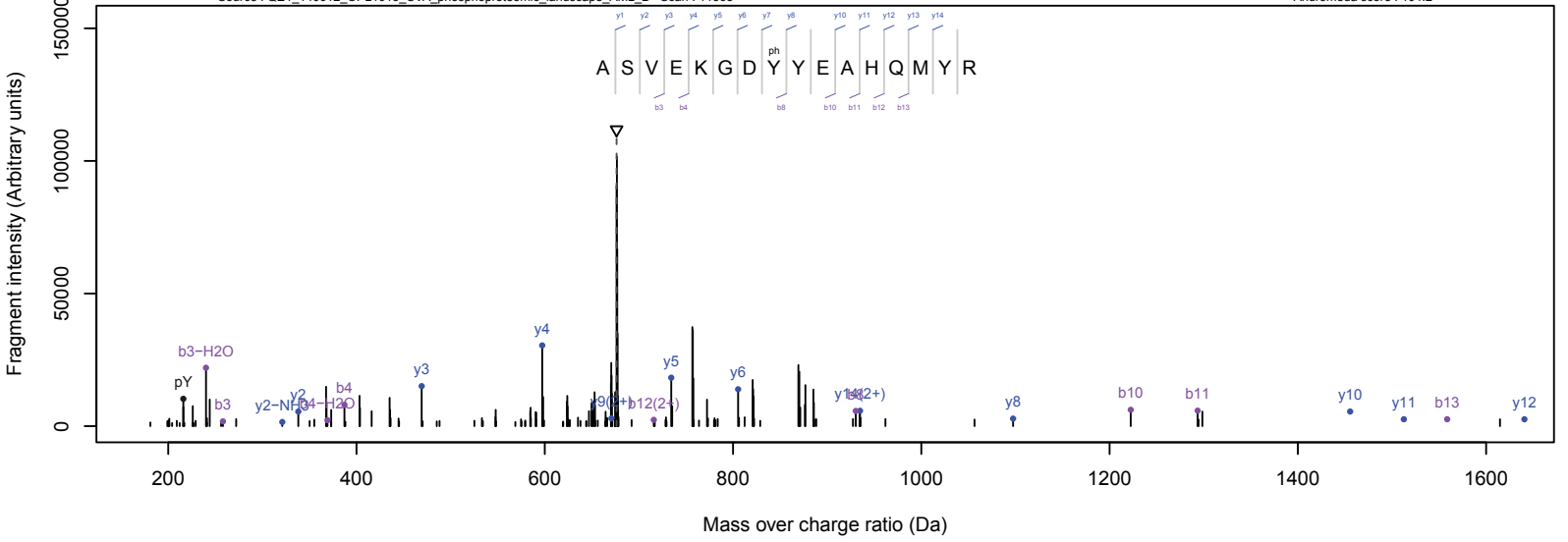
Andromeda score : 104.2



ASVEKGDY(ph)YEAHQMYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11568

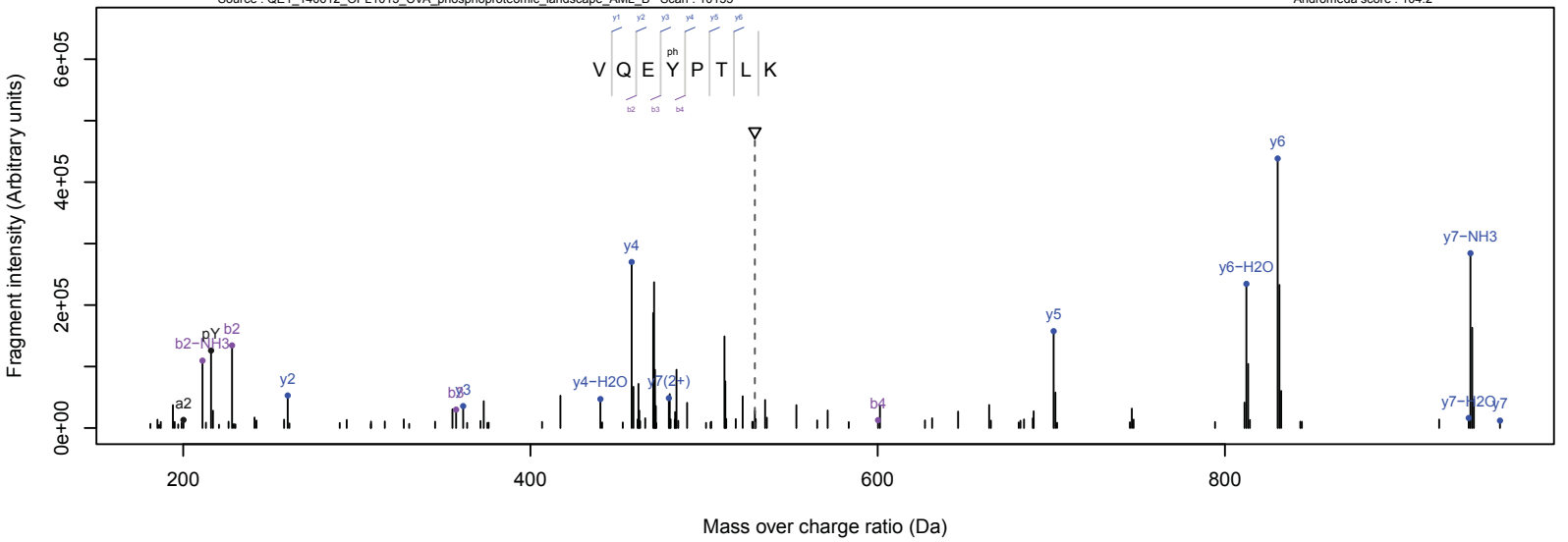
Andromeda score : 104.2



VQEY(ph)PTLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10155

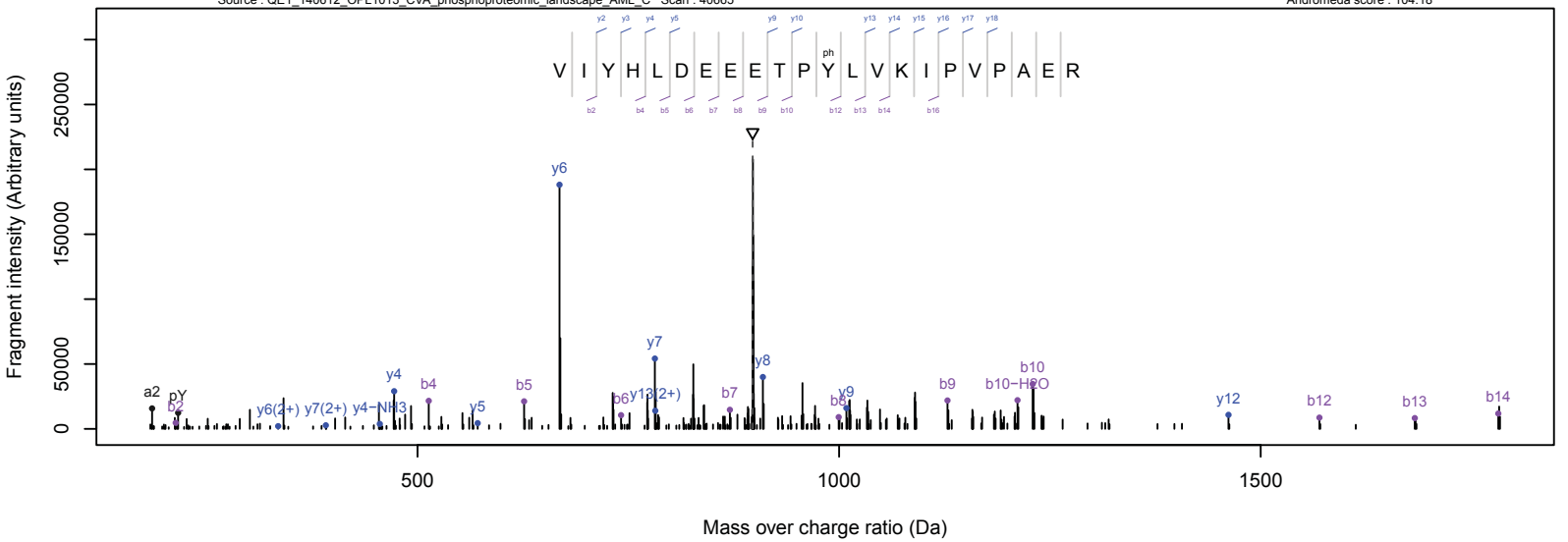
Andromeda score : 104.2



VIYHLDEEETPY(ph)LVKIPVPAER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40665

Andromeda score : 104.18

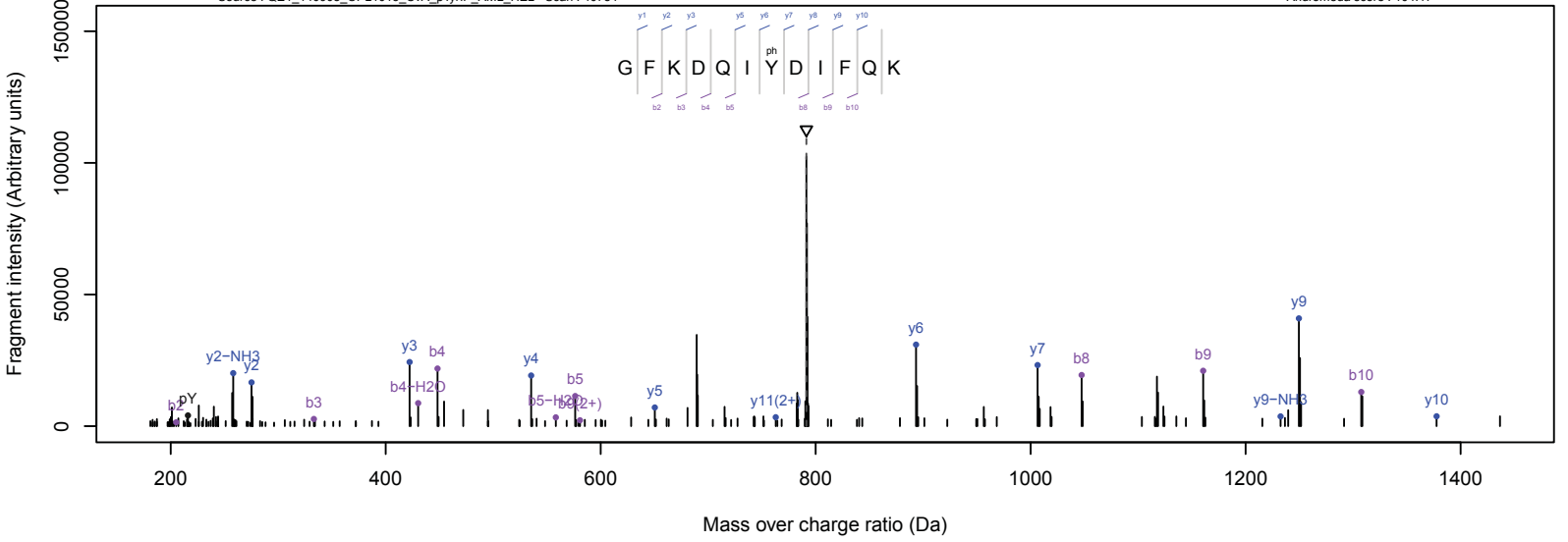




GFKDQIY(ph)DIFQK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 40784

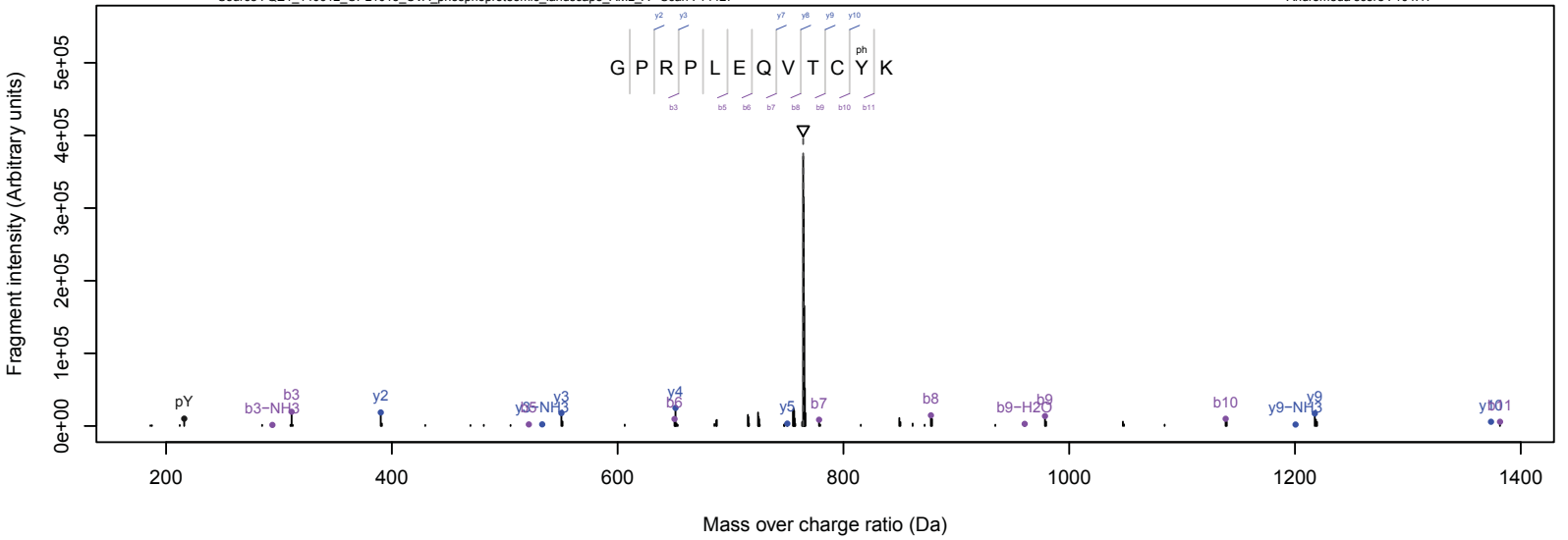
Andromeda score : 104.17



GPRPLEQVTCYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 14427

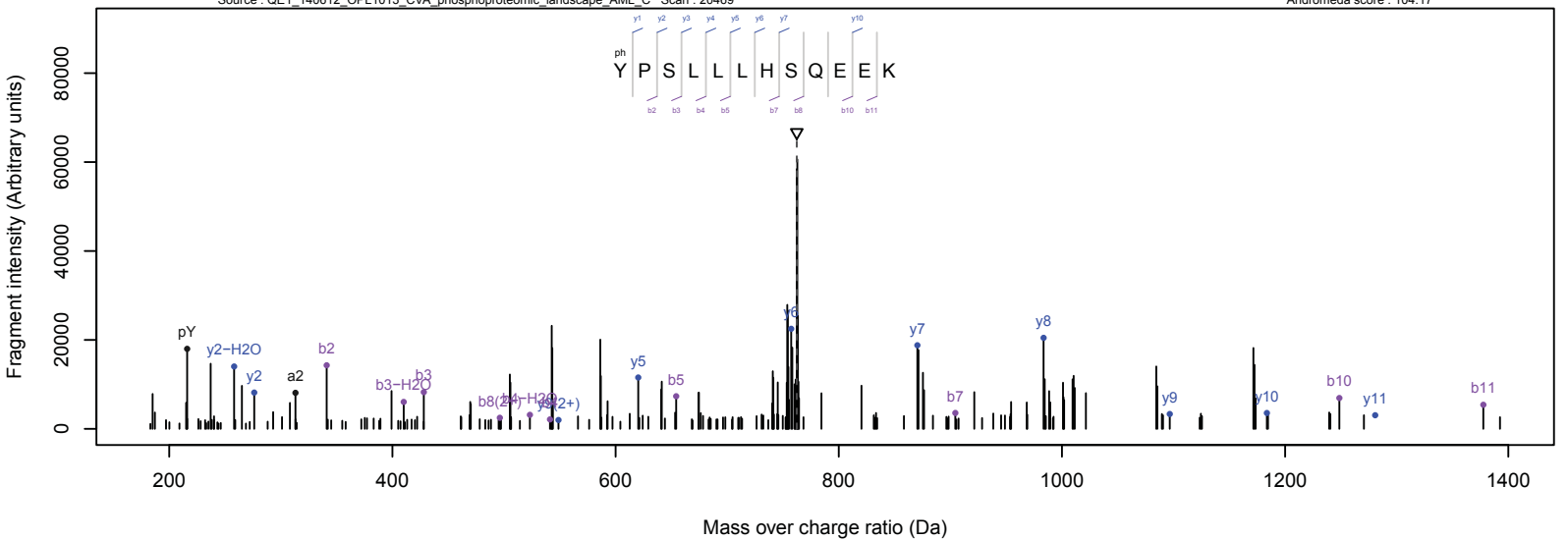
Andromeda score : 104.17



Y(ph)PSLLLSQEEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20469

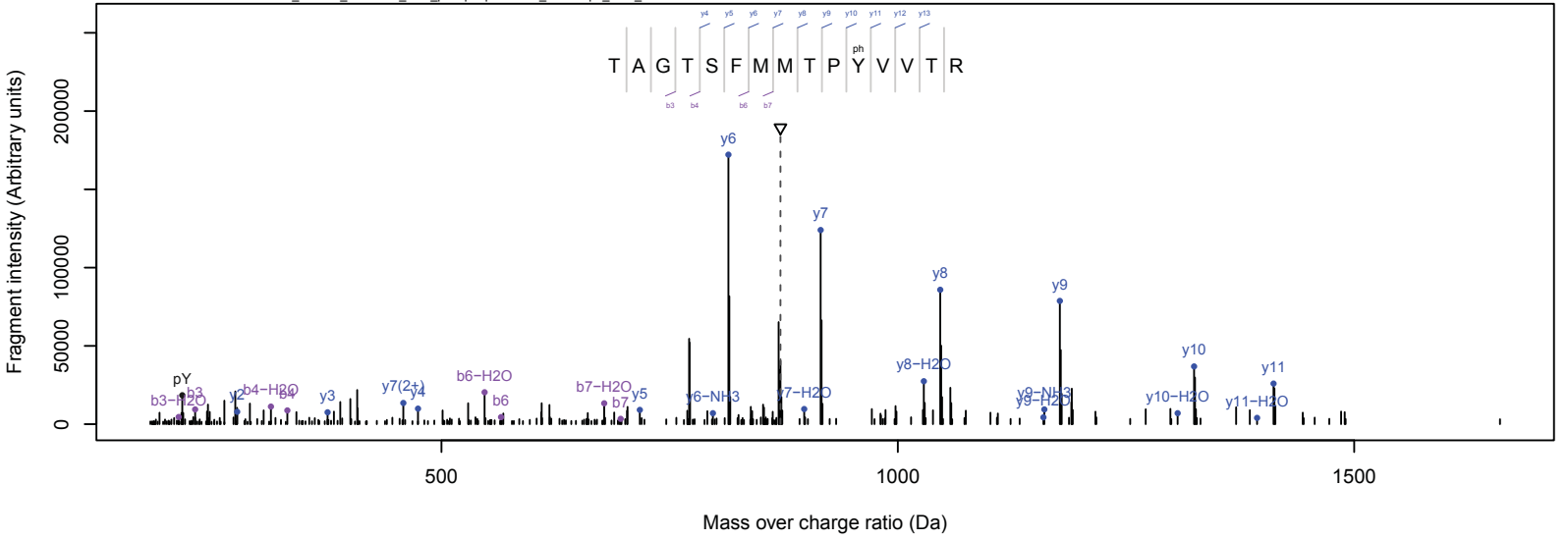
Andromeda score : 104.17



**TAGTSMMPY(ph)VVTR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36815

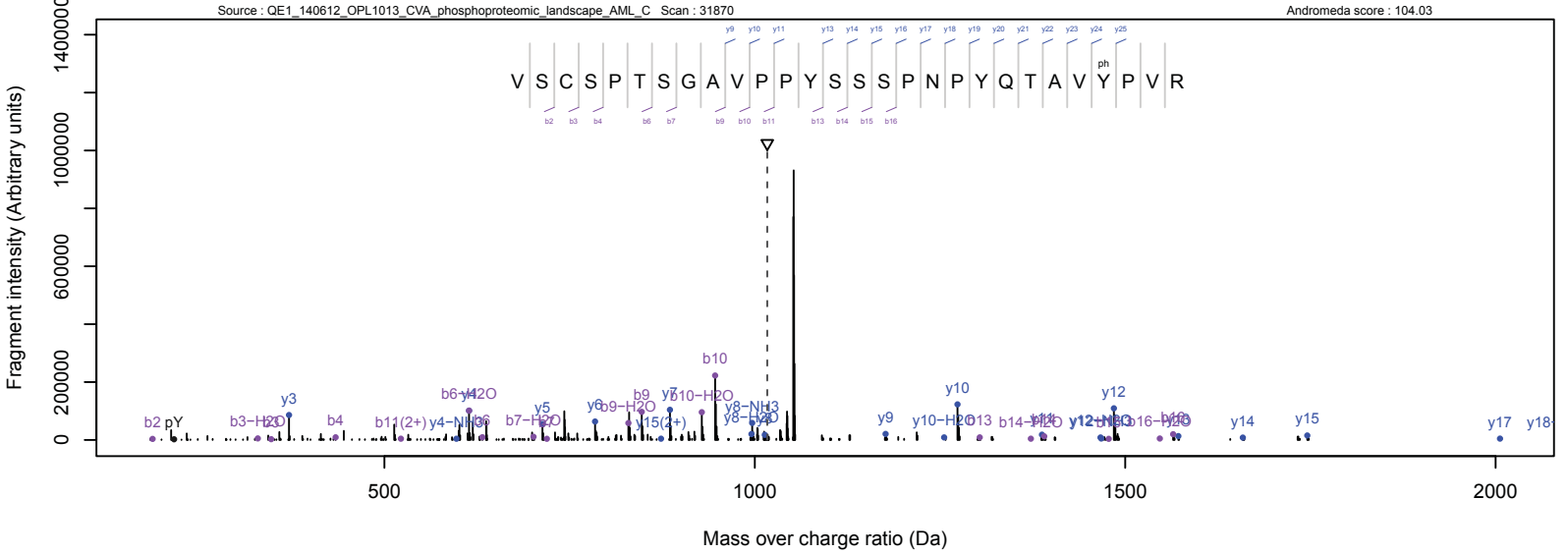
Andromeda score : 104.09



**VSCSPTSGAVPPYSSSPNPYQTAVY(ph)PVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31870

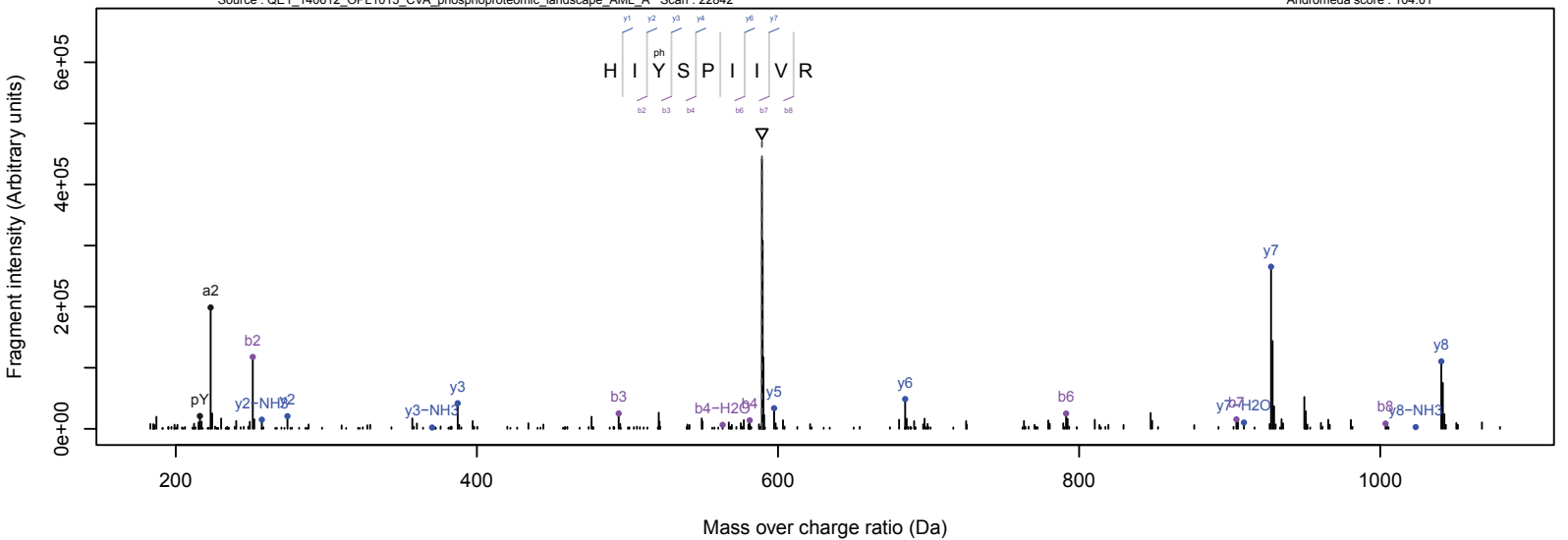
Andromeda score : 104.03



**HIY(ph)SPIIVR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 22842

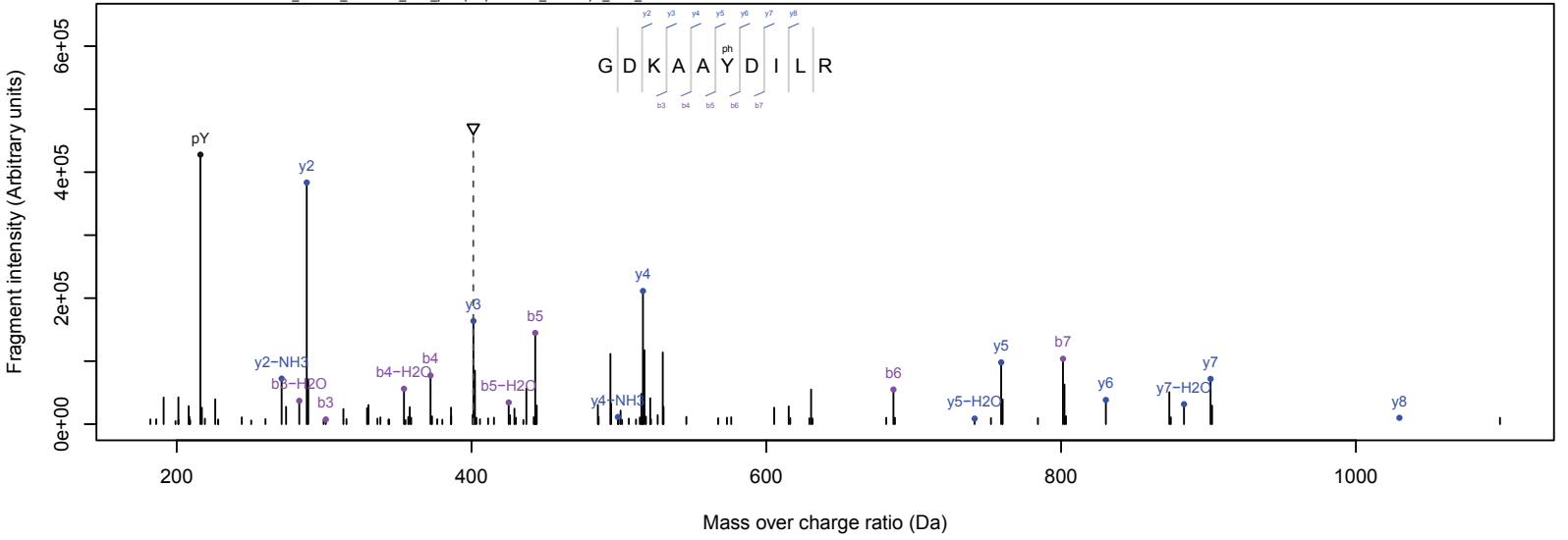
Andromeda score : 104.01



**GDKAAY(ph)DILR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15847

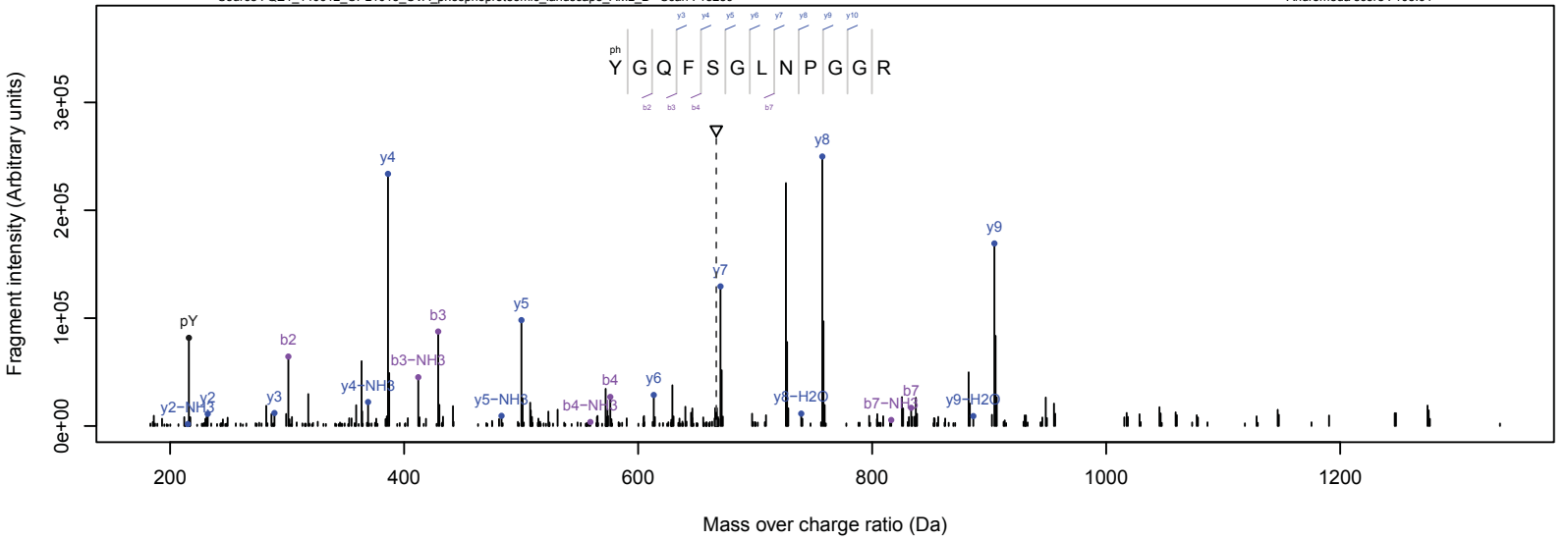
Andromeda score: 103.91



**Y(ph)GQFSLNPGGR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18289

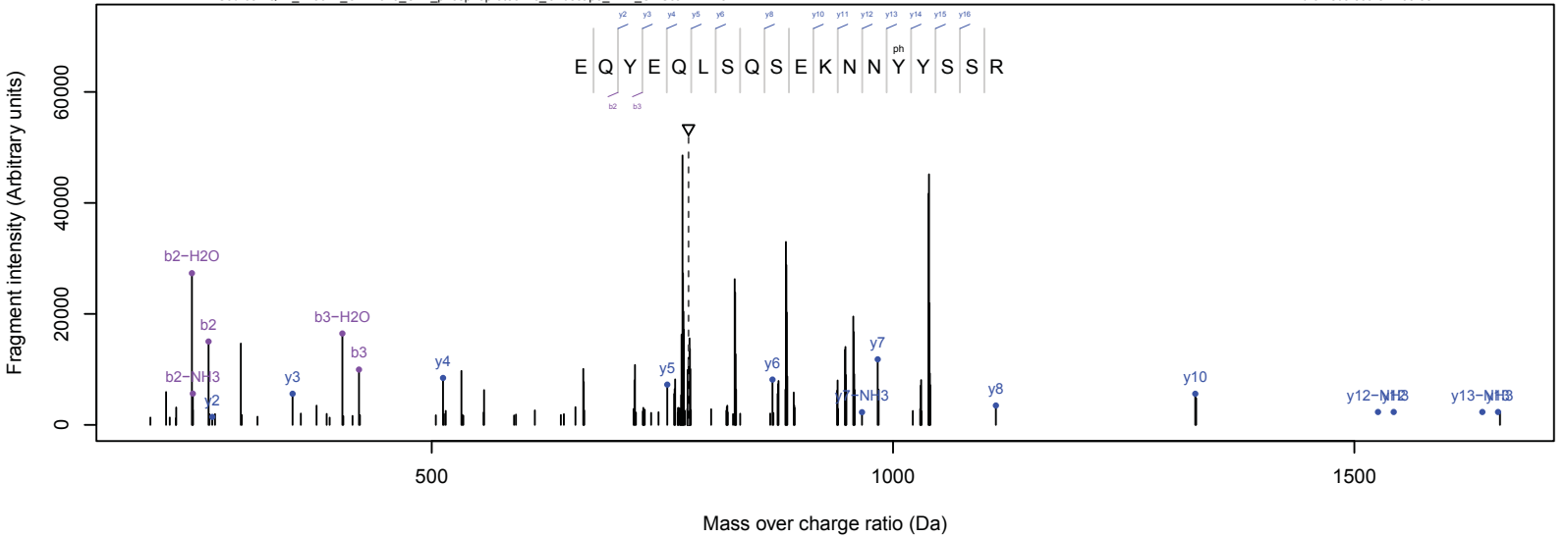
Andromeda score: 103.91



**EQYEQLSQSEKNNYYSR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12282

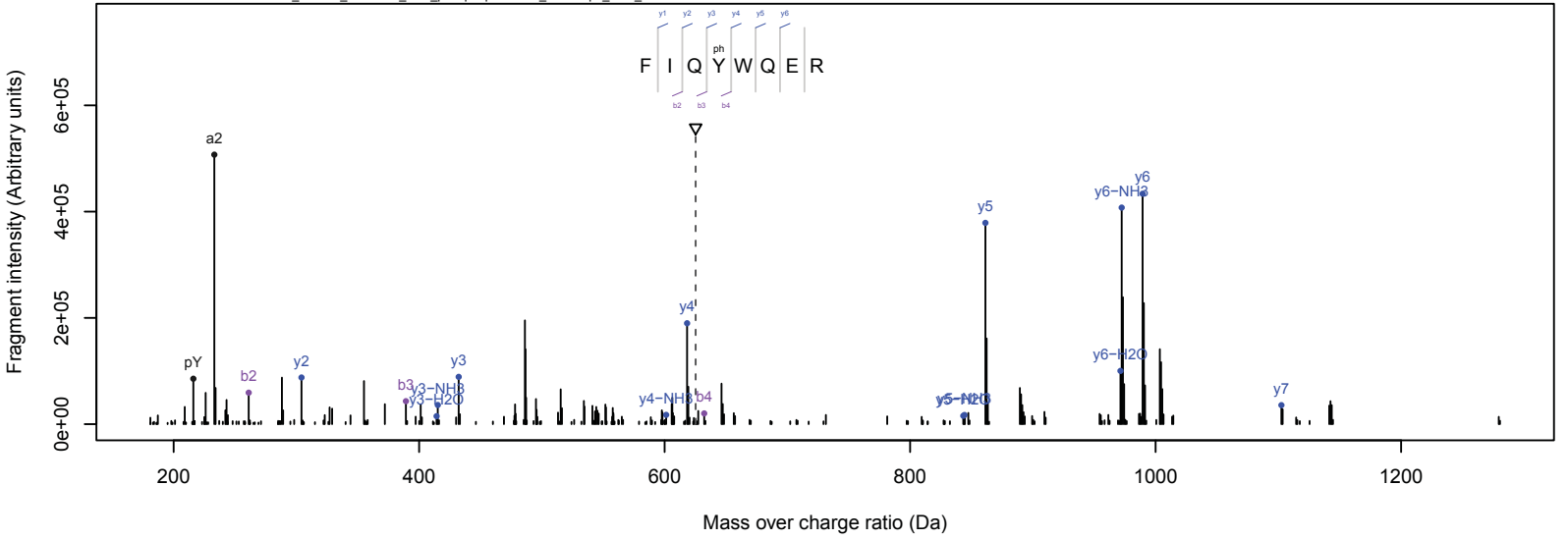
Andromeda score: 103.88



FIQY(ph)WQER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23789

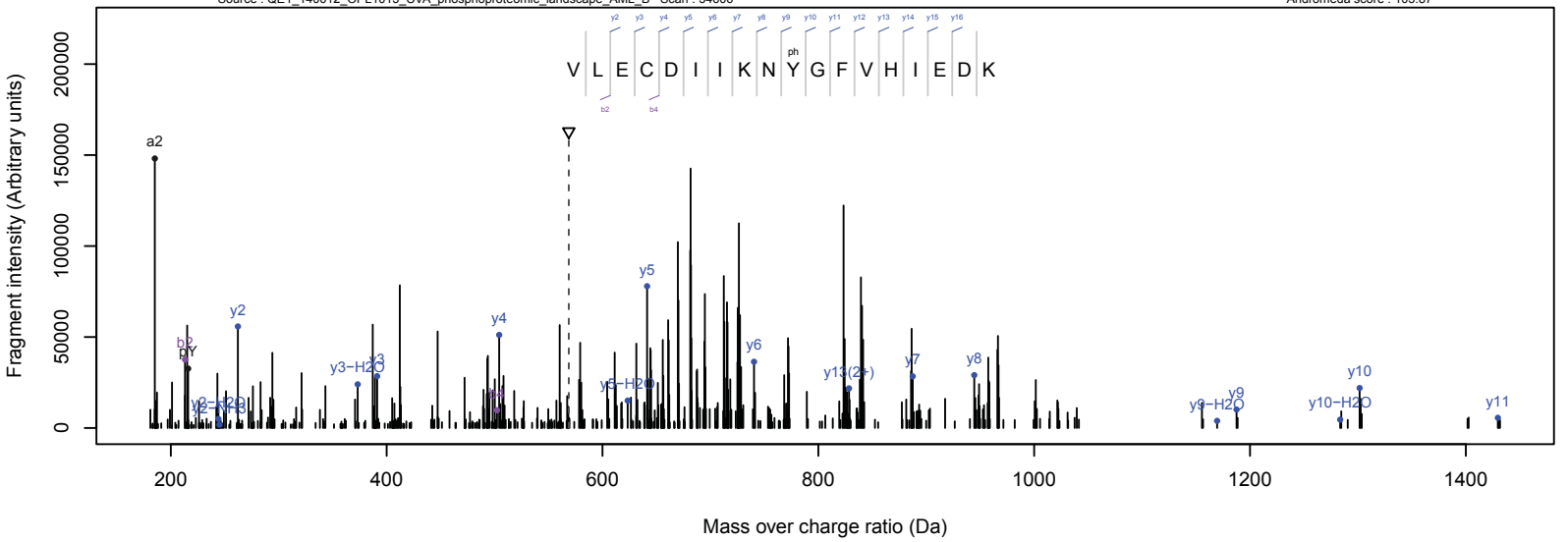
Andromeda score : 103.87



VLECDIKNY(ph)GFVHIEDK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34606

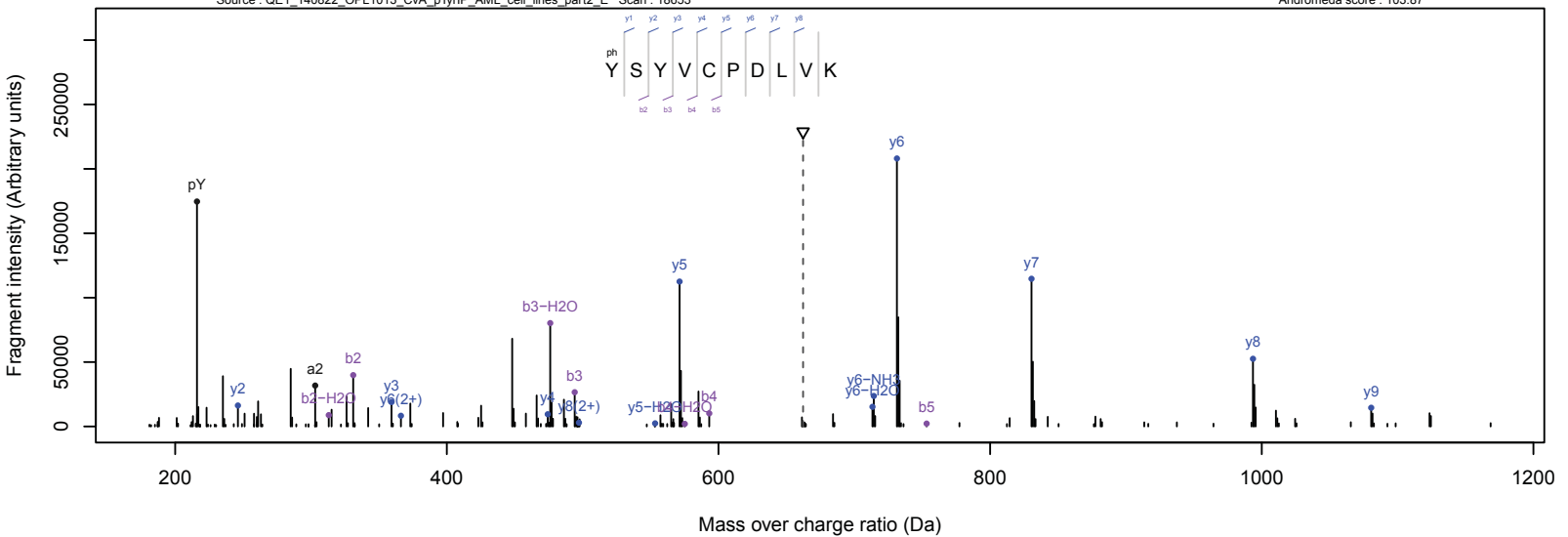
Andromeda score : 103.87



Y(ph)SYVCPDLVK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 18653

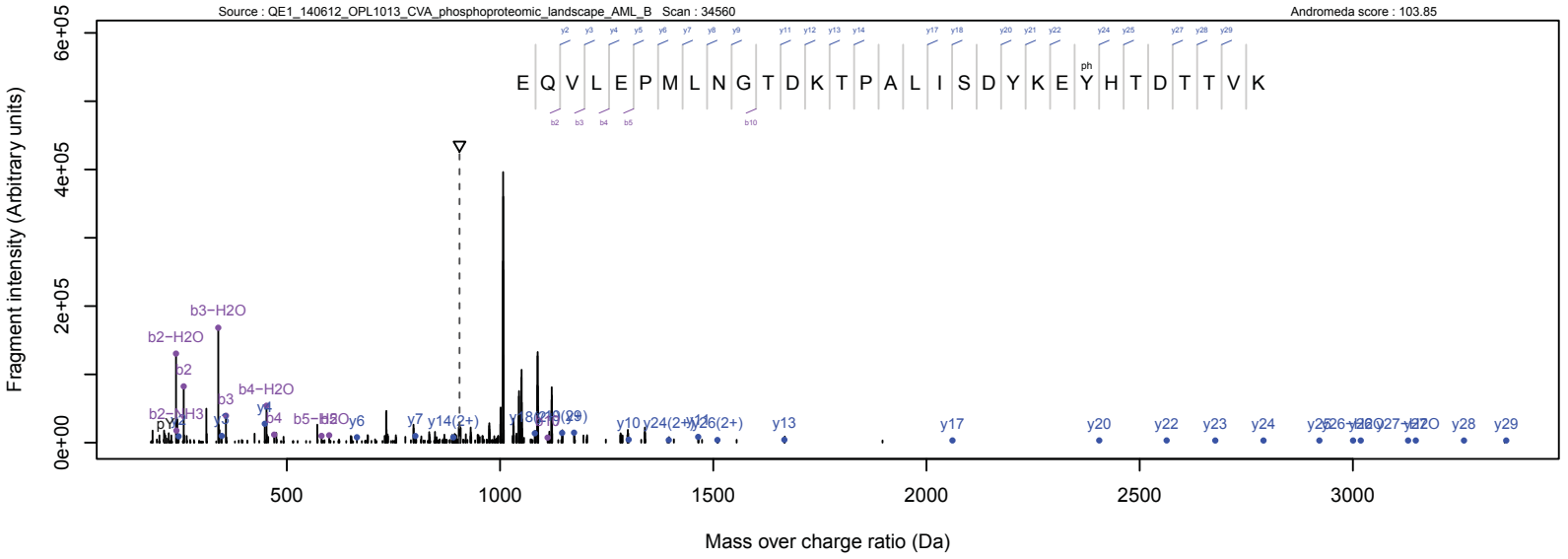
Andromeda score : 103.87



**EQVLEPMLNGTDKTPALISDYKEY(ph)HTDITTK-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34560

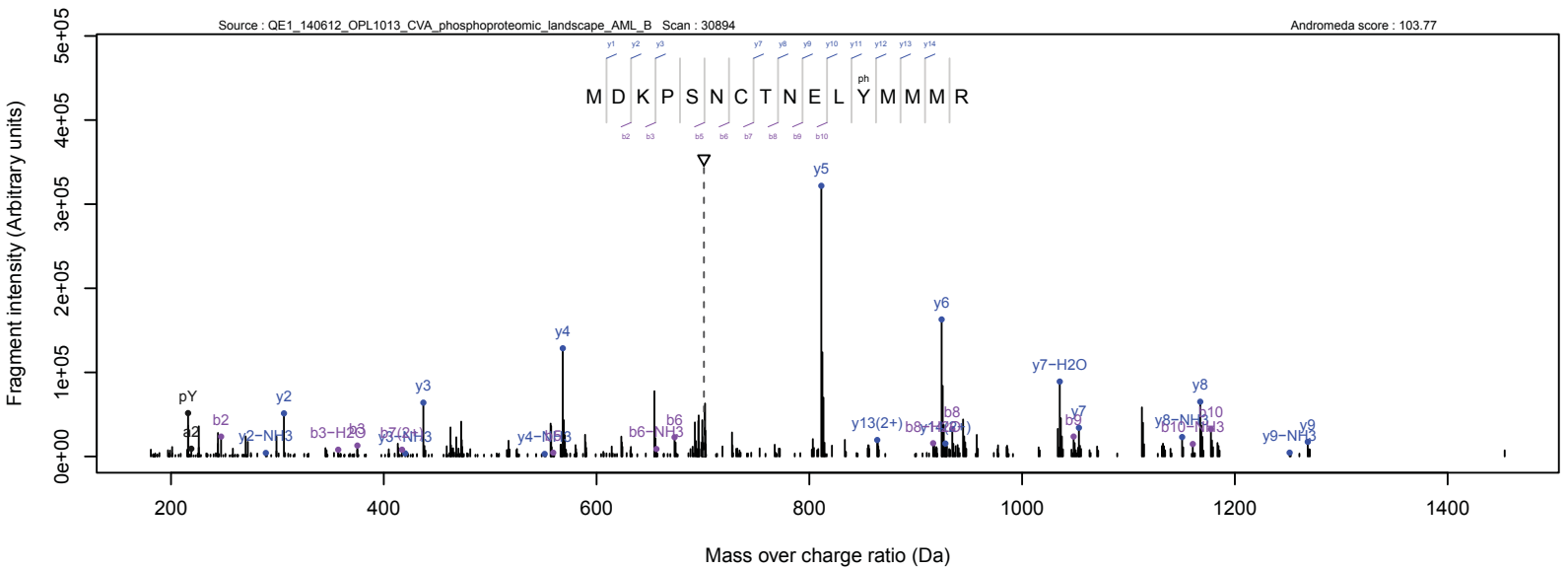
Andromeda score: 103.85



**MDKPSNCTNELY(MMMR)-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 30894

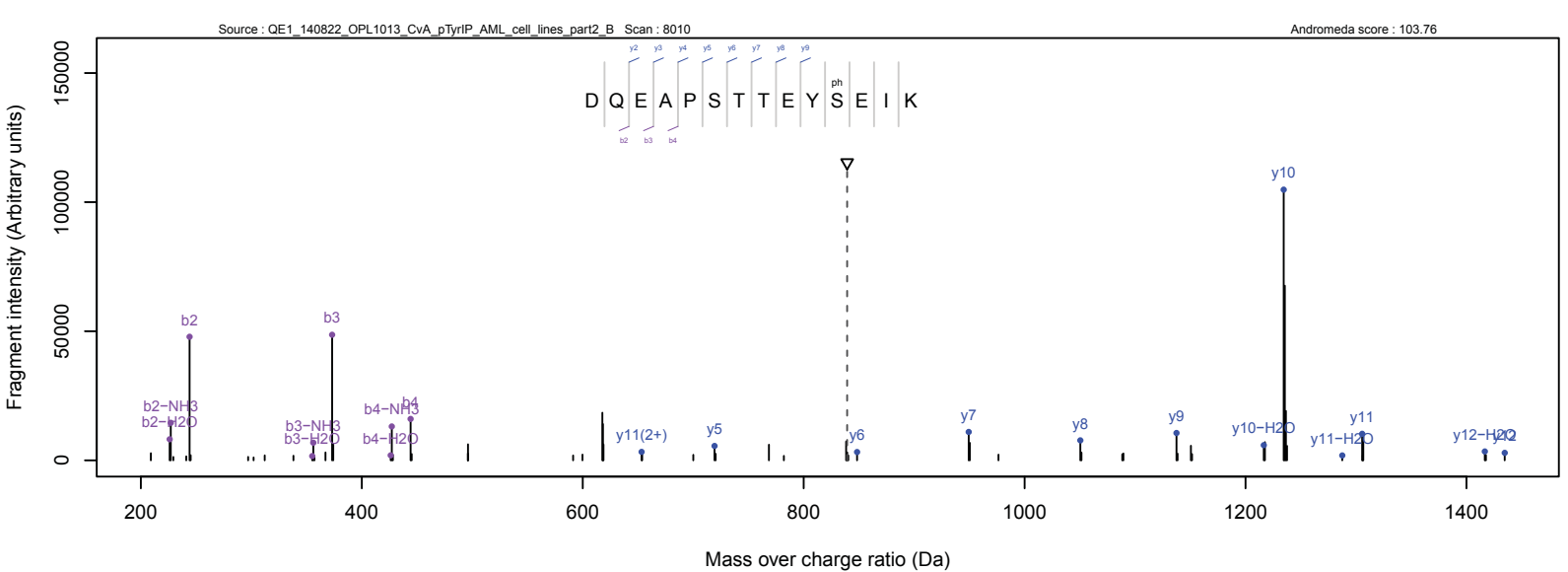
Andromeda score: 103.77



**DQEAPSTTEYS(ph)EIK-2+**

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 8010

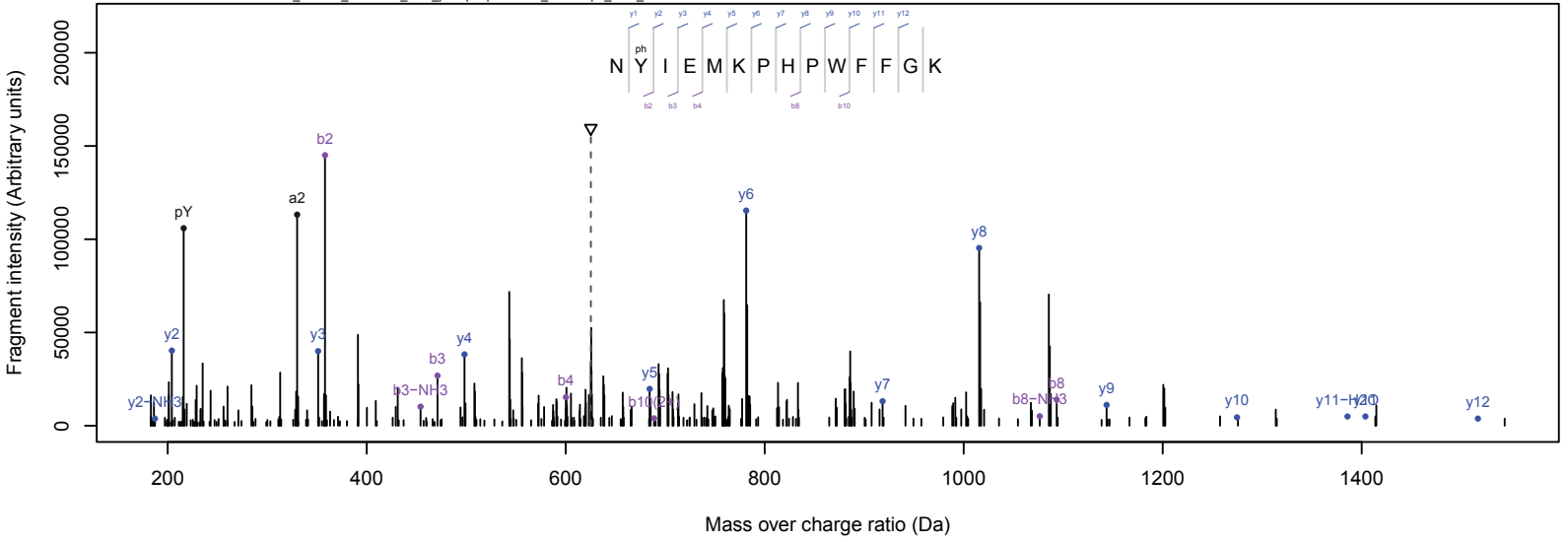
Andromeda score: 103.76



**NY(ph)IEMKHPWFFGK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 35812

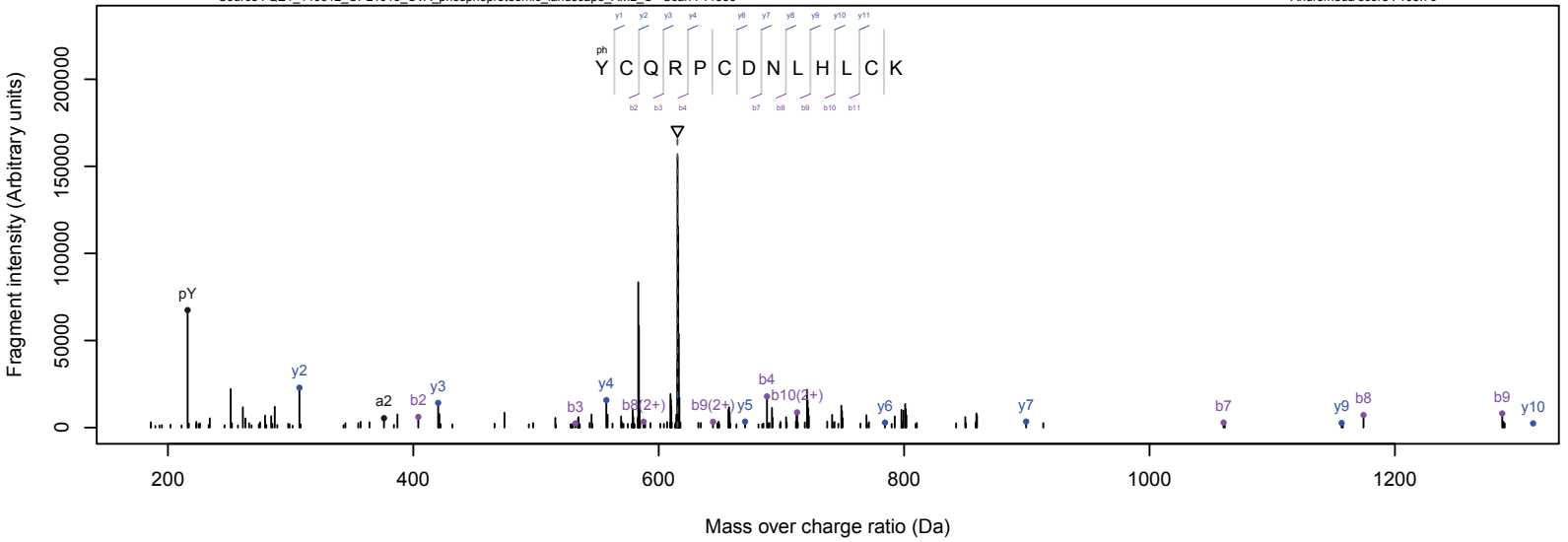
Andromeda score: 103.76



**Y(ph)CQRPCDNLHLCK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11553

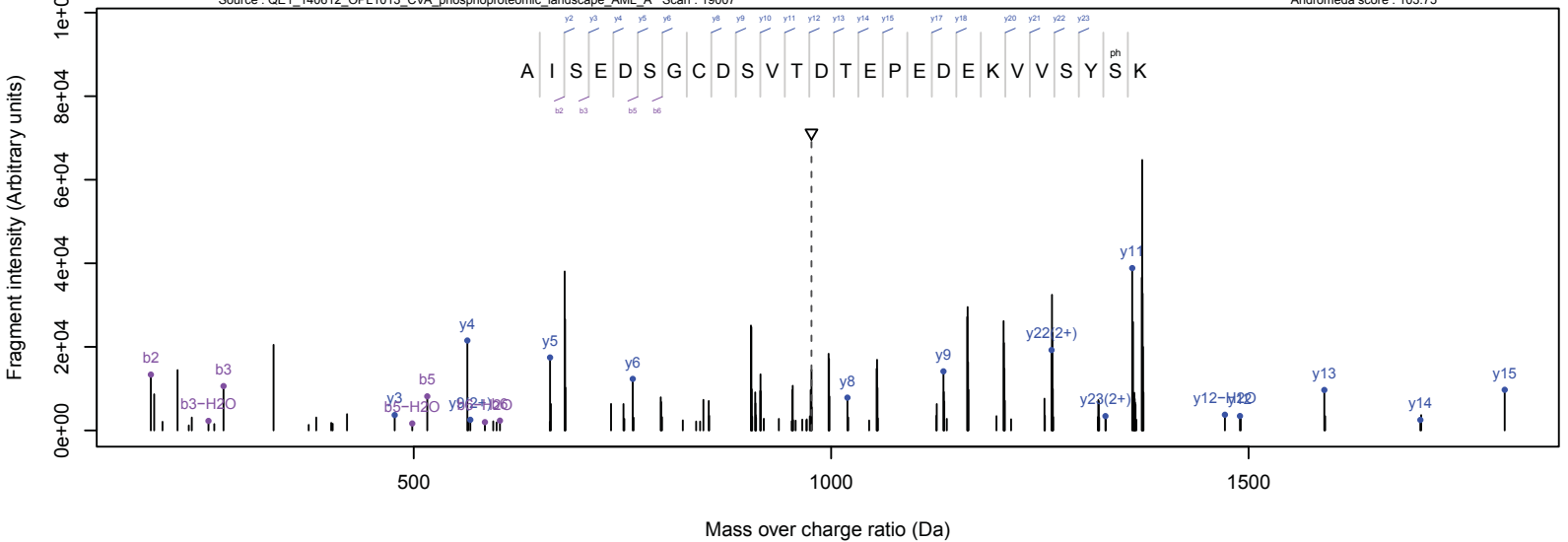
Andromeda score: 103.76



**AISESGCDSVTDTEPEDEKVVSY(ph)K-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 19007

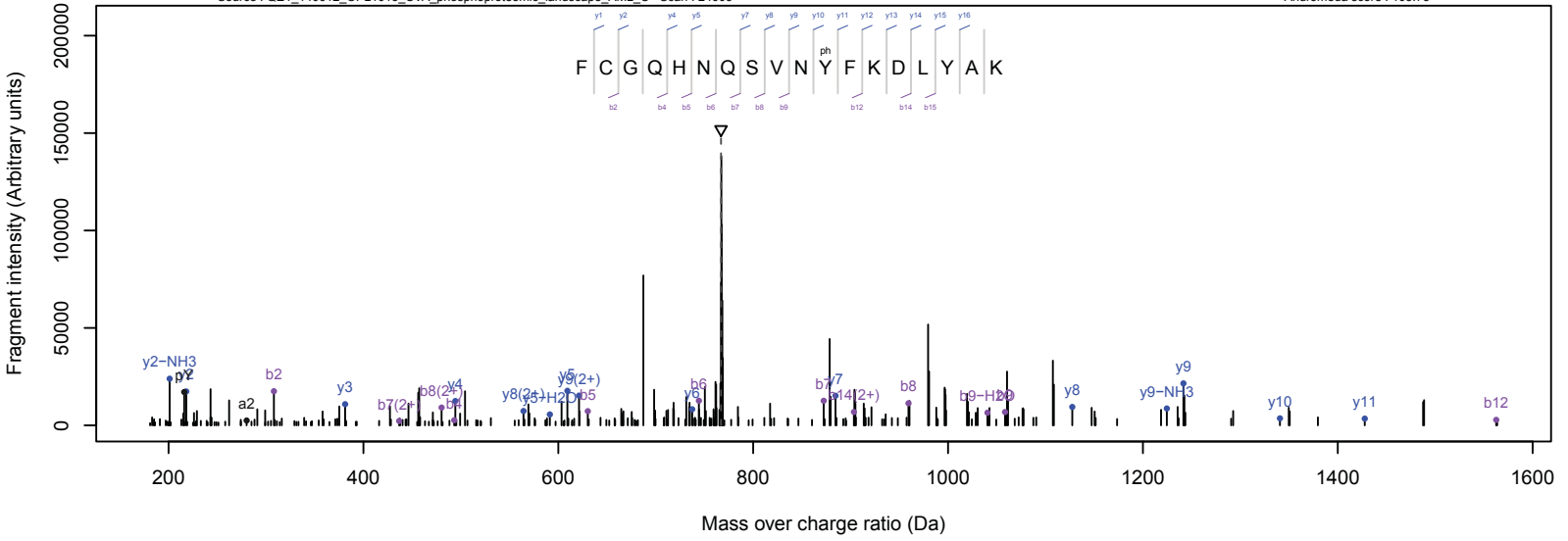
Andromeda score: 103.75



FCGQHNQSVNY(ph)FKDLYAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24003

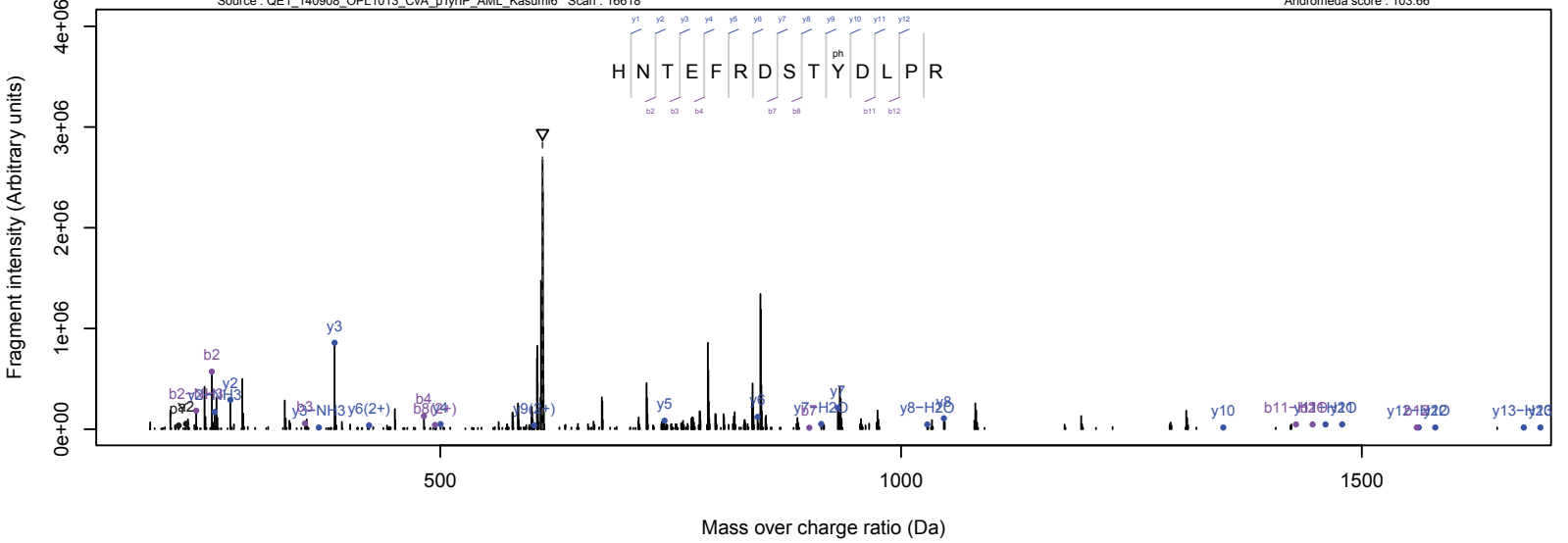
Andromeda score : 103.73



HNTEFRDSTY(ph)DLPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 16618

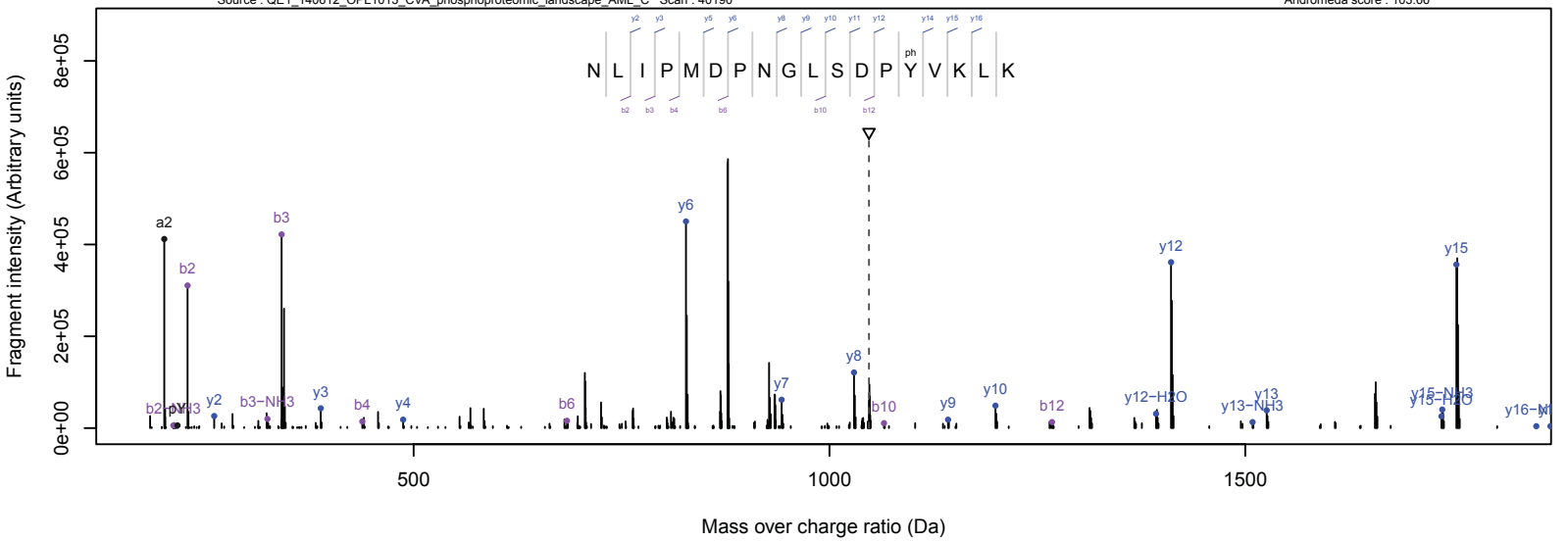
Andromeda score : 103.66



NLIPMDP NGLSDPY(ph)VKLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40190

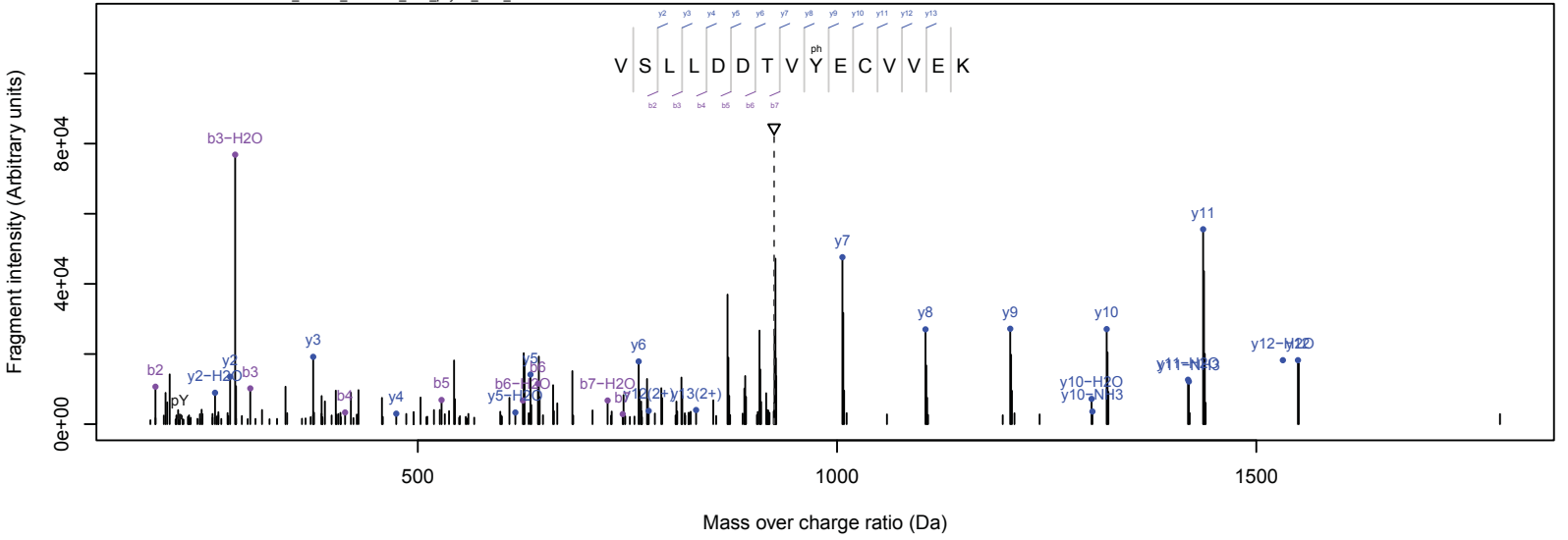
Andromeda score : 103.66



VSLDDTVY(ph)ECVVEK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 37163

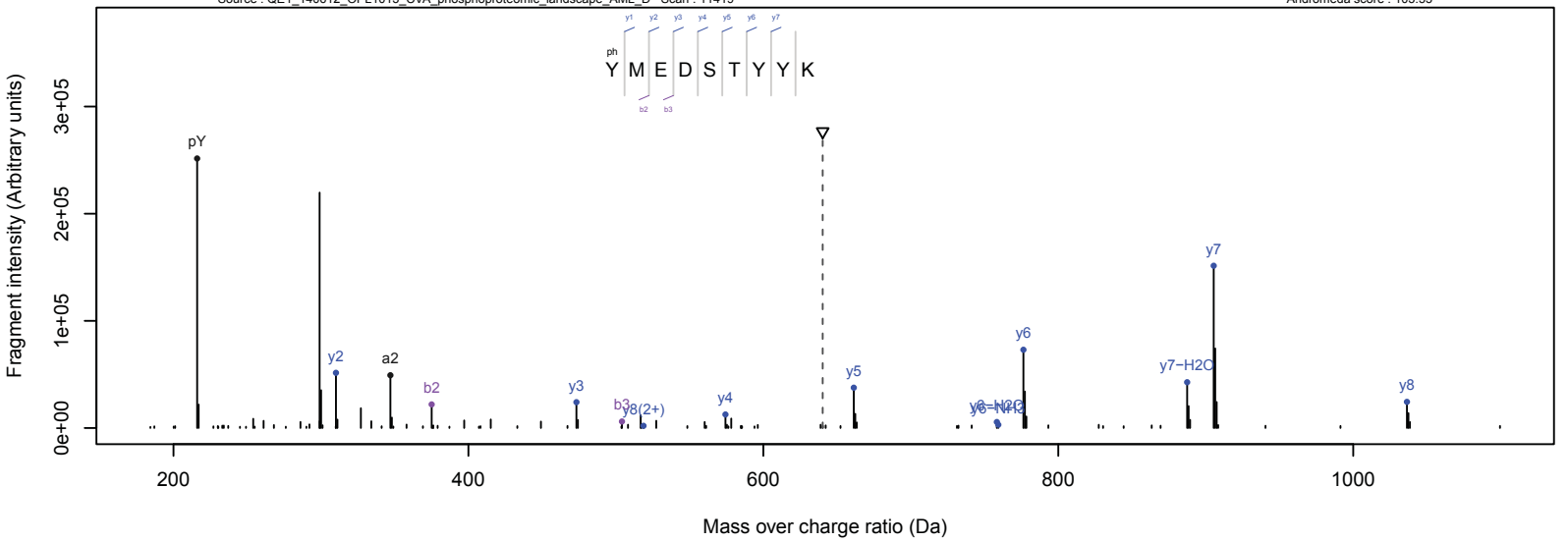
Andromeda score : 103.6



Y(ph)MEDSTYYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 11419

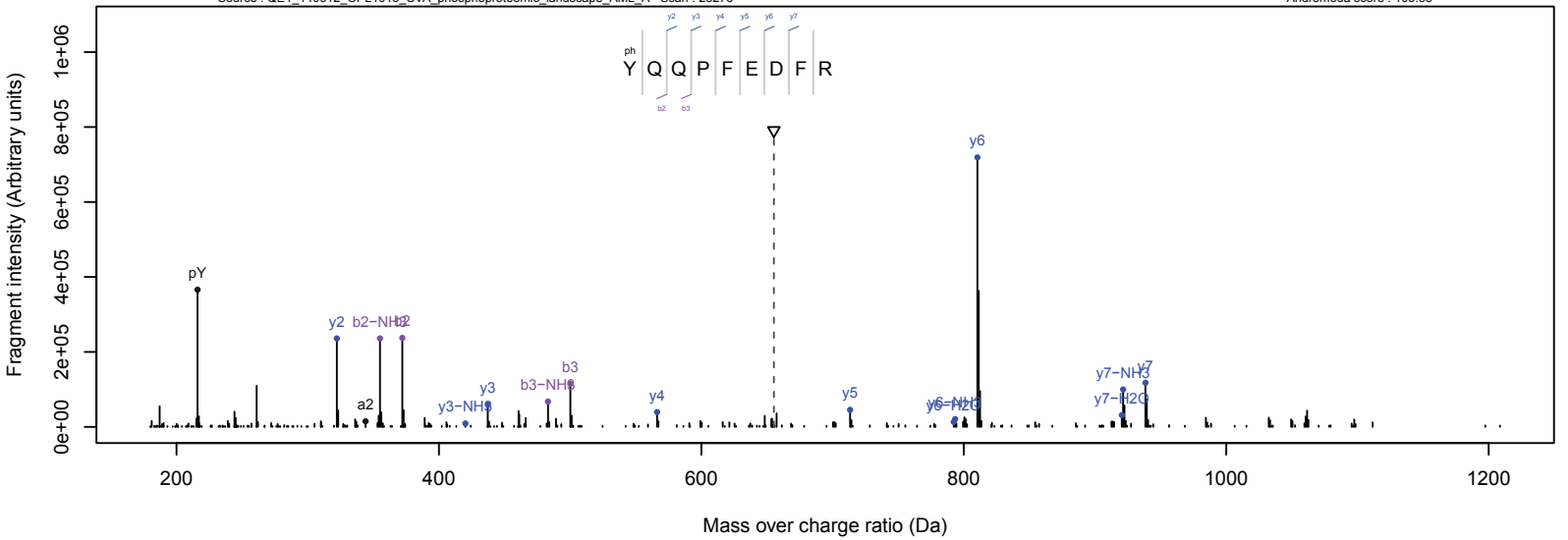
Andromeda score : 103.55



Y(ph)QQPFEDFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25275

Andromeda score : 103.55

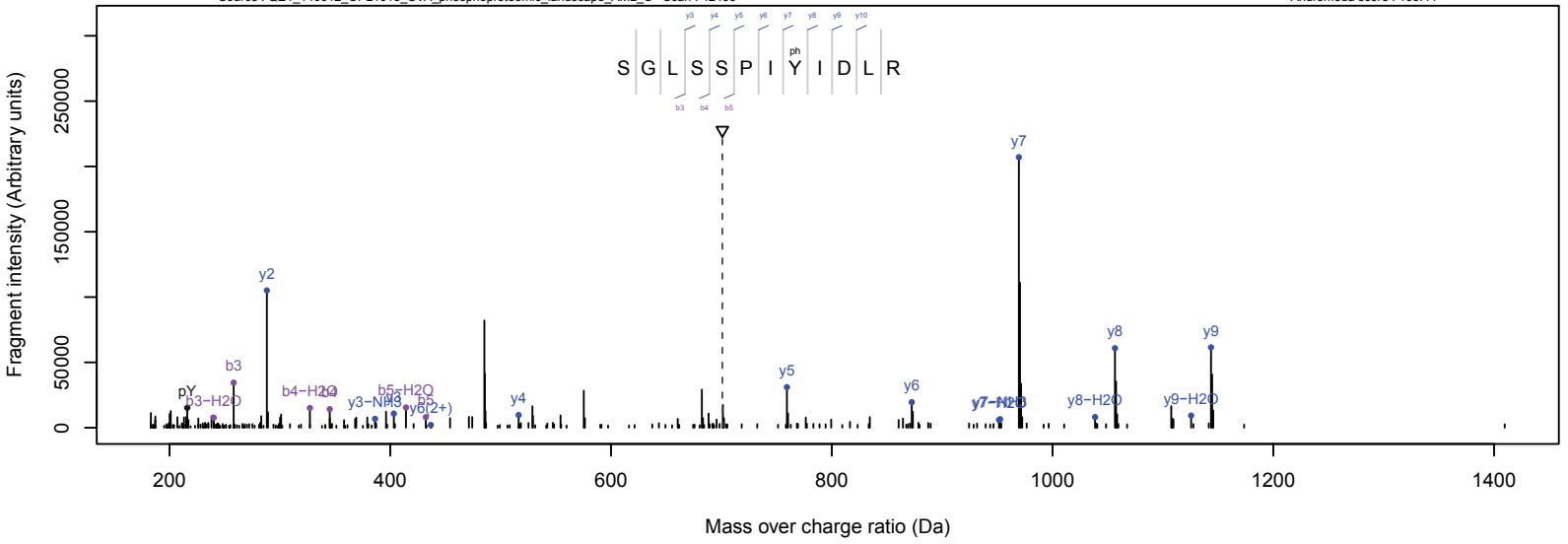




**SGLSSPIY(ph)IDLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42455

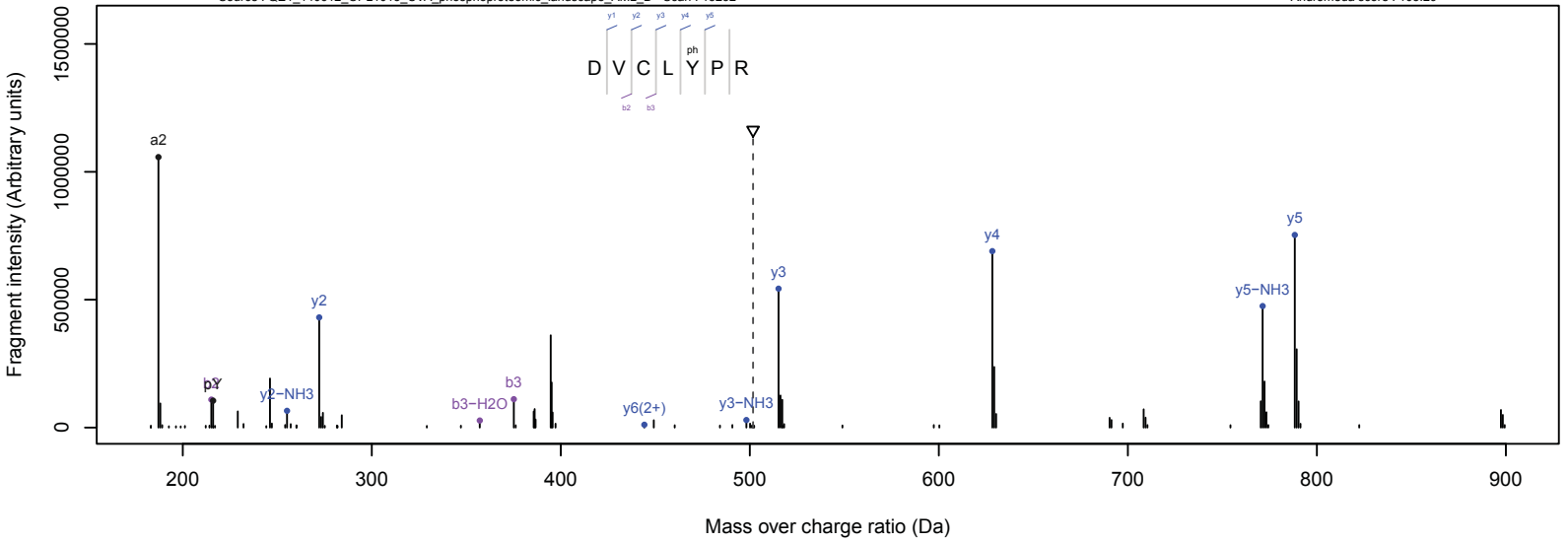
Andromeda score : 103.44



**DVCLY(ph)PR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 15232

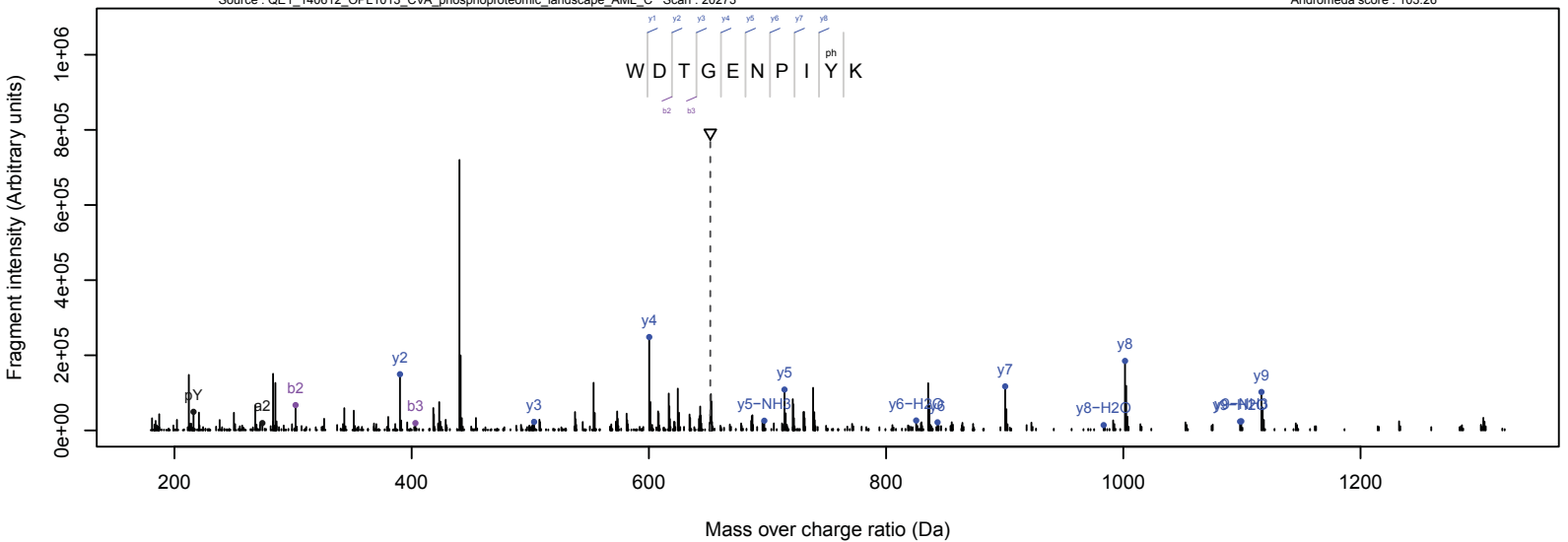
Andromeda score : 103.29



**WDTGENPIYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20273

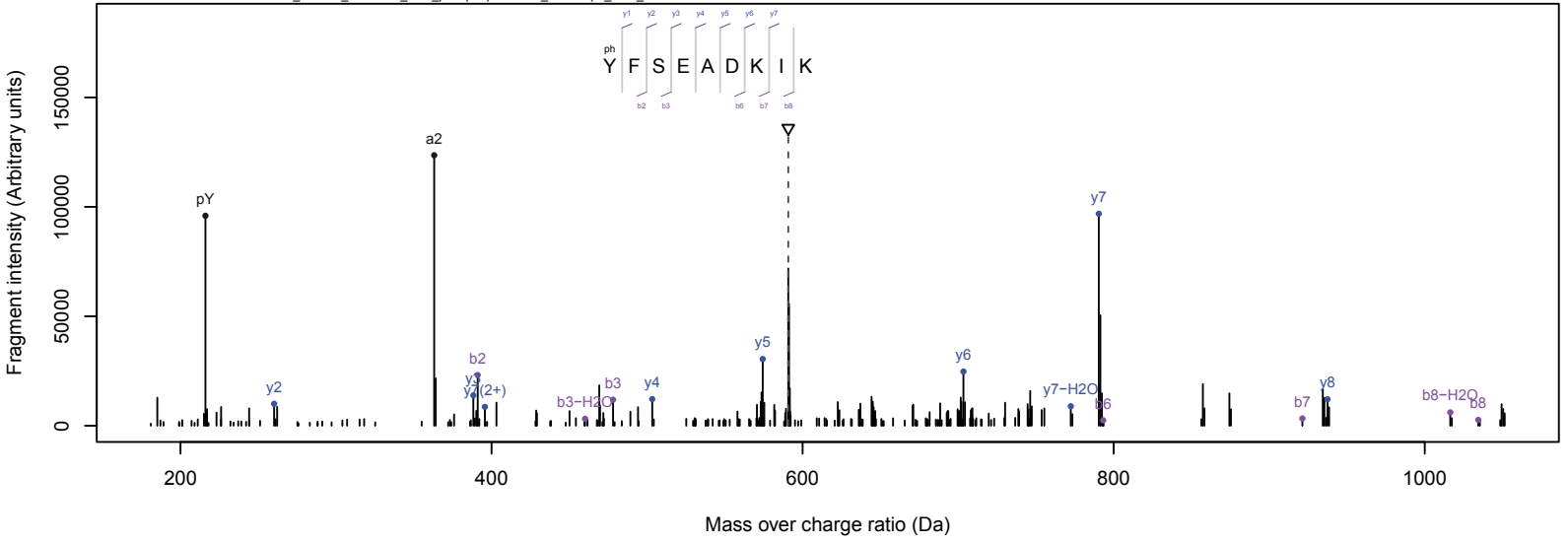
Andromeda score : 103.26



Y(ph)FSEADKIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10088

Andromeda score : 103.26

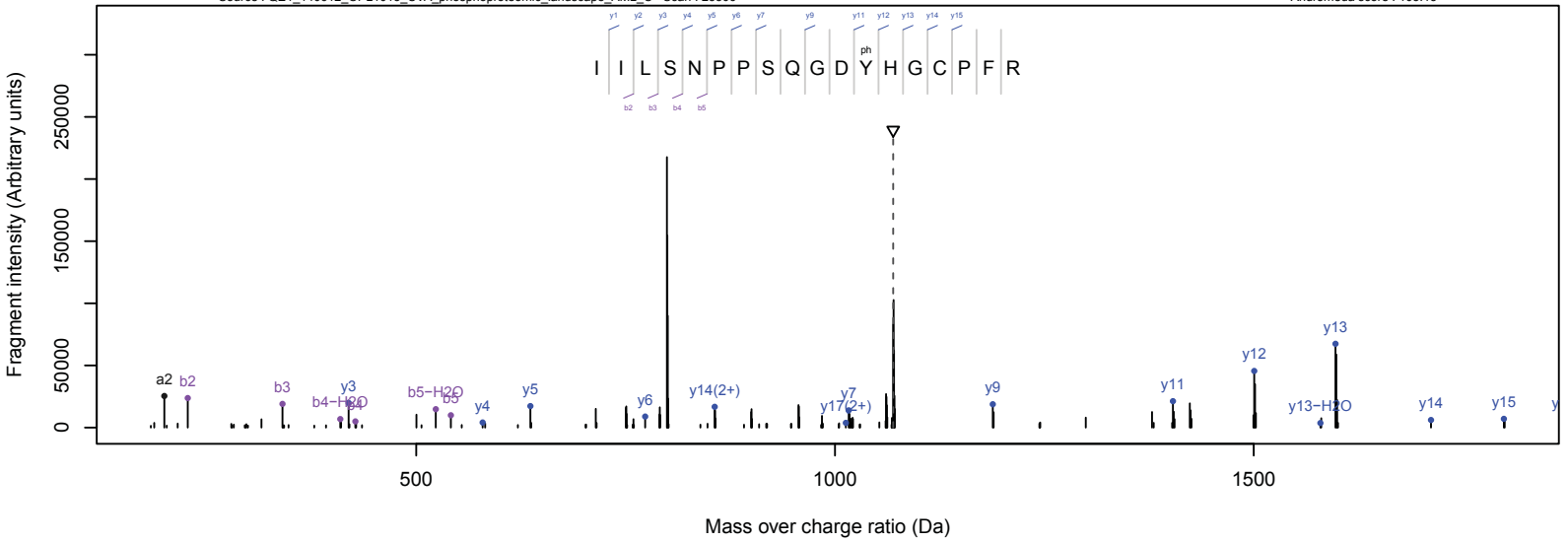


Mass over charge ratio (Da)

IILSNPPSQGDYHGCPFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25506

Andromeda score : 103.13

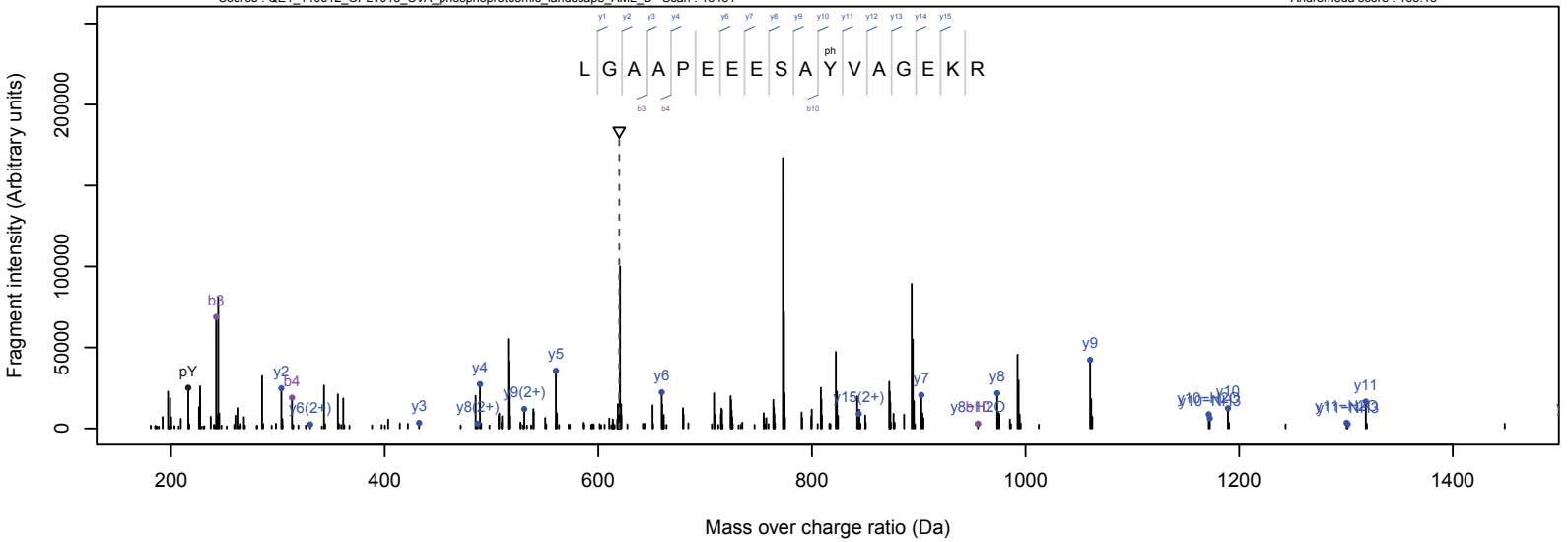


Mass over charge ratio (Da)

LGAPEEESAYVAGEKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13451

Andromeda score : 103.13

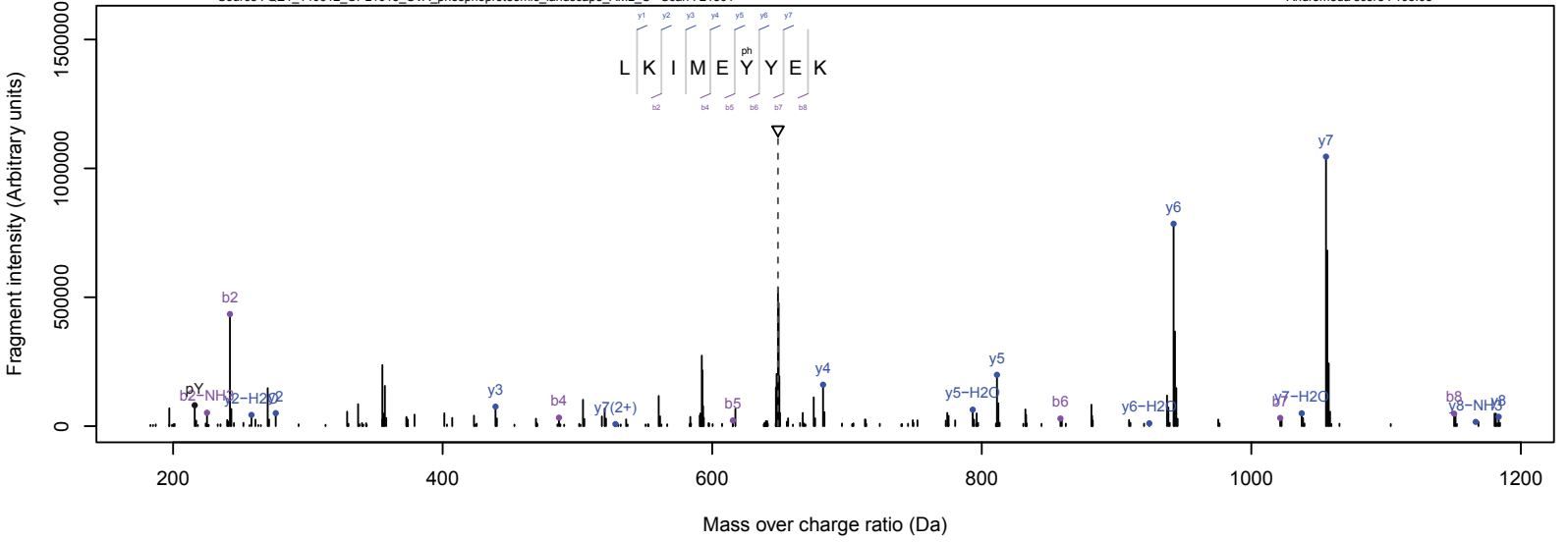


Mass over charge ratio (Da)

LKIMEY(ph)YEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21801

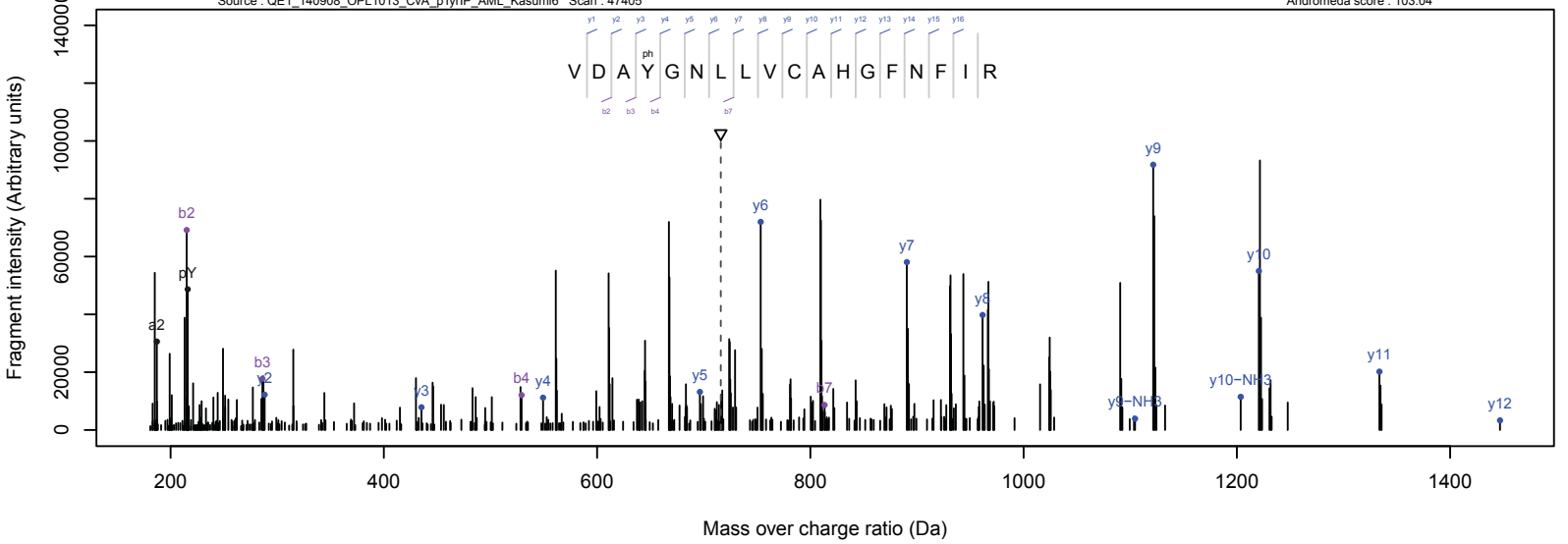
Andromeda score : 103.08



VDAY(ph)GNLLVCAHGFNFI R-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 47405

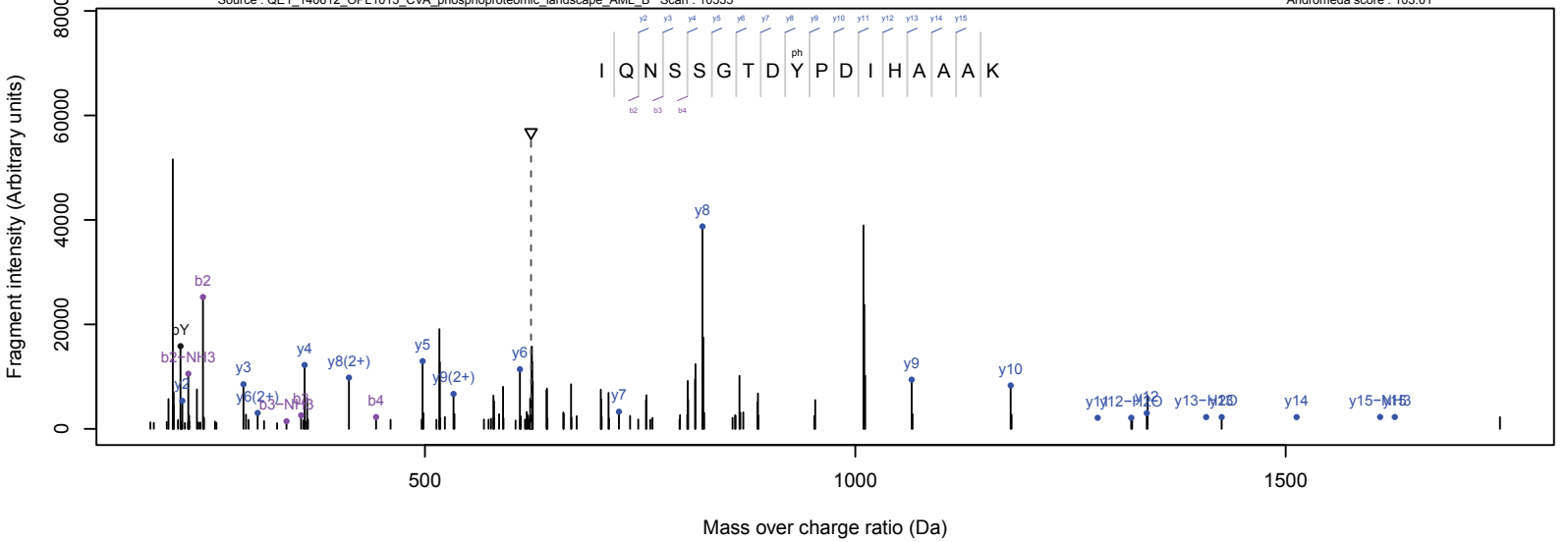
Andromeda score : 103.04



IQSSGTDY(ph)PDIHAAAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10535

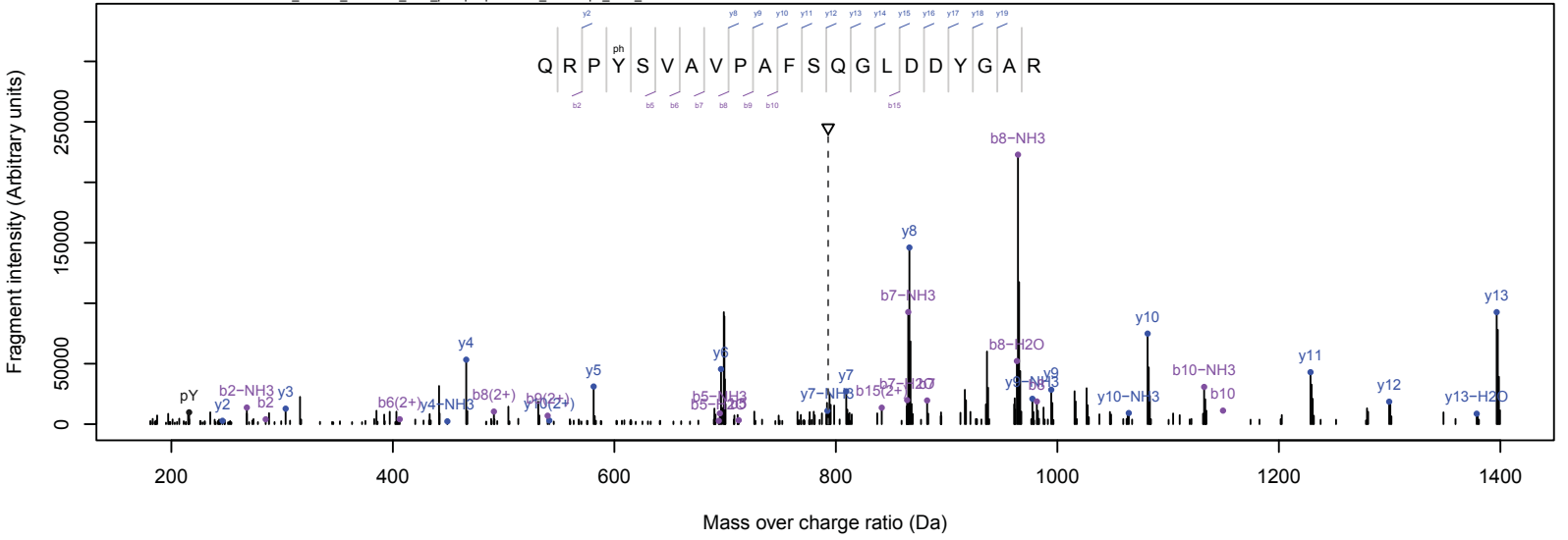
Andromeda score : 103.01



QRPY(ph)SVAVPAFSQGLDDYGAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34810

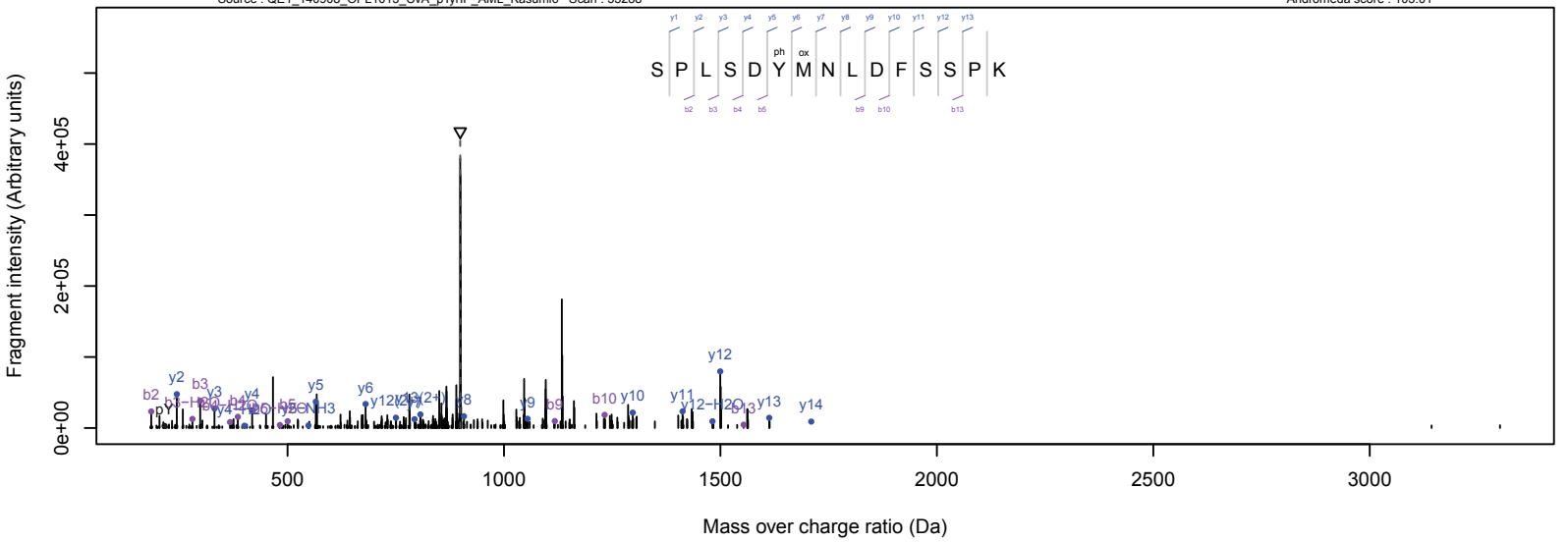
Andromeda score : 103.01



SPLSDY(ph)M(ox)NLDFSSPK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 33288

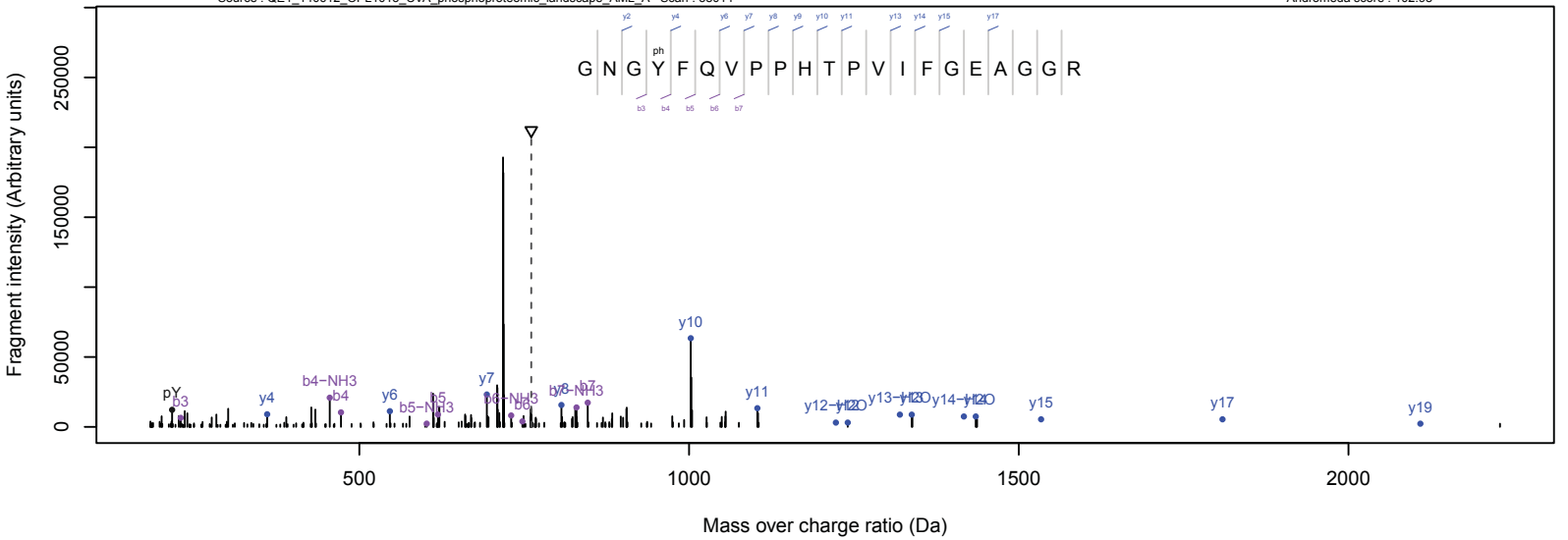
Andromeda score : 103.01



NGY(ph)FQVPHTPVIFGEAGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 38014

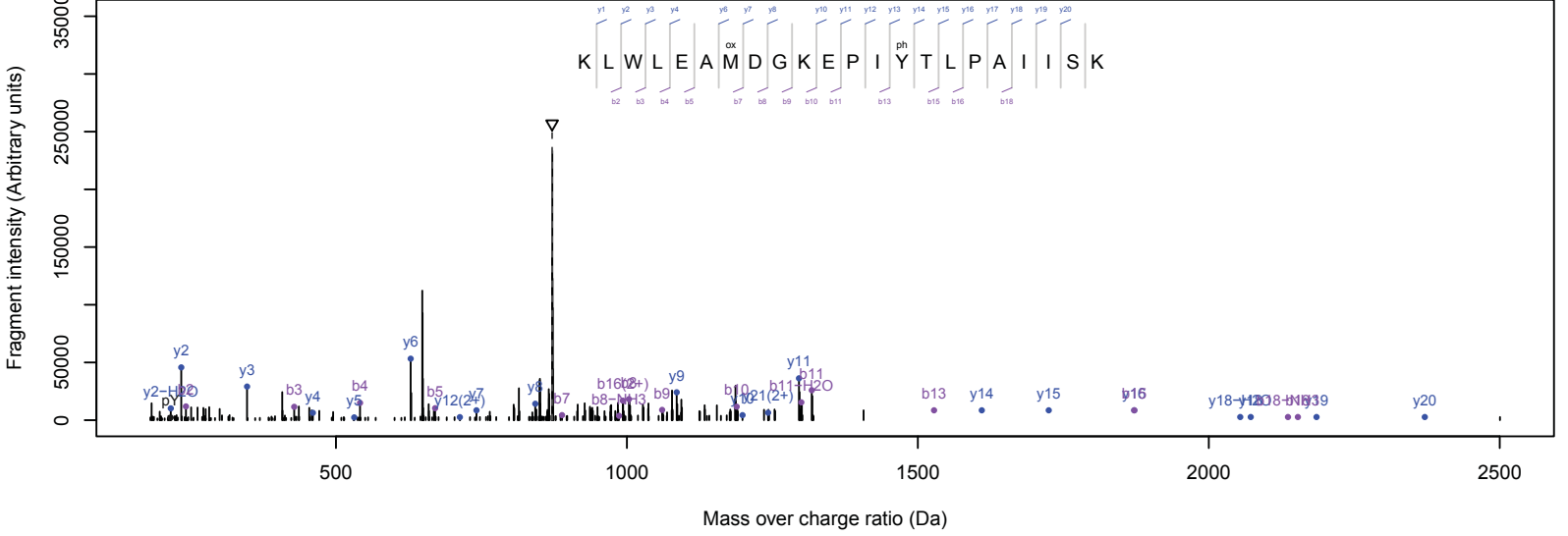
Andromeda score : 102.98



**KLWLEAM(ox)DGKEPIY(ph)TLPAISK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 46423

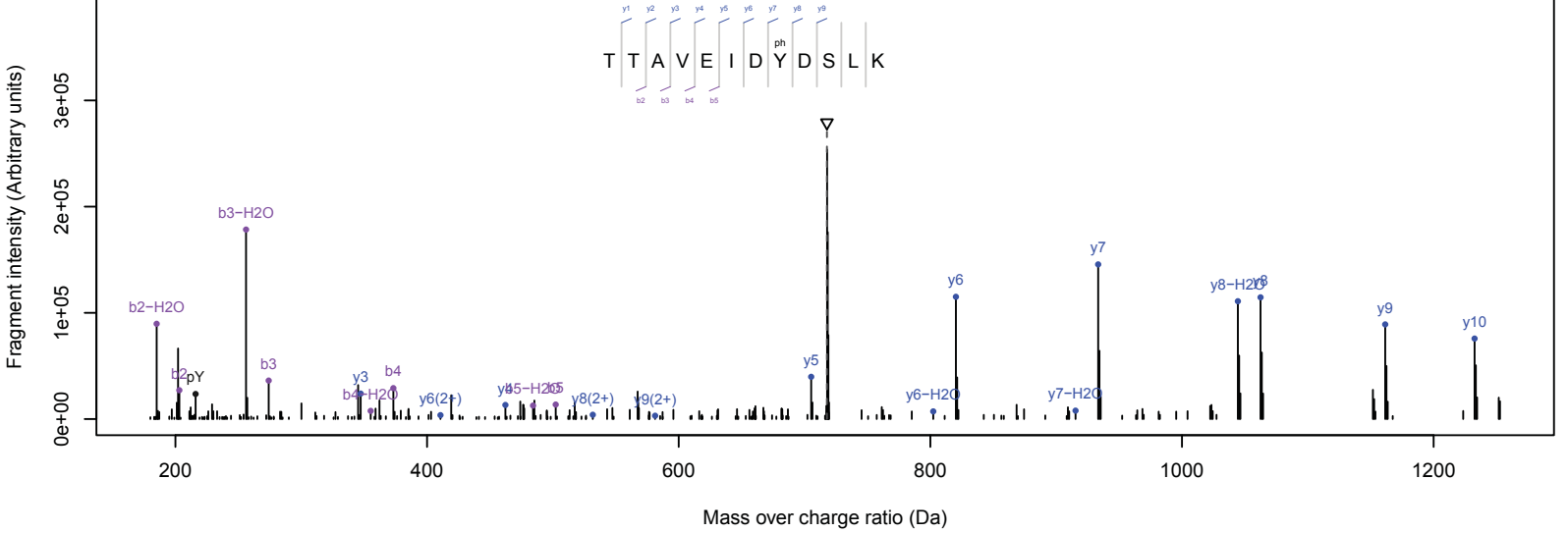
Andromeda score : 102.97



**TTAVEIDY(ph)DSLK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 26696

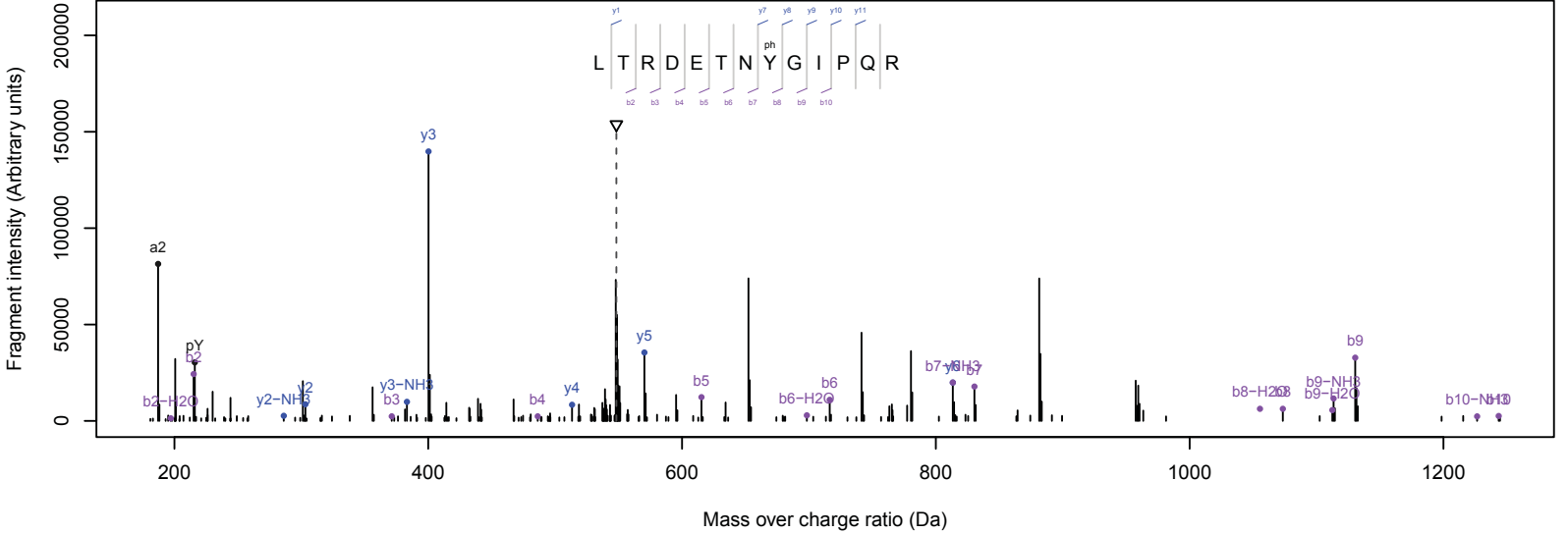
Andromeda score : 102.97



**LTRDETNY(ph)GIPQR-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 6844

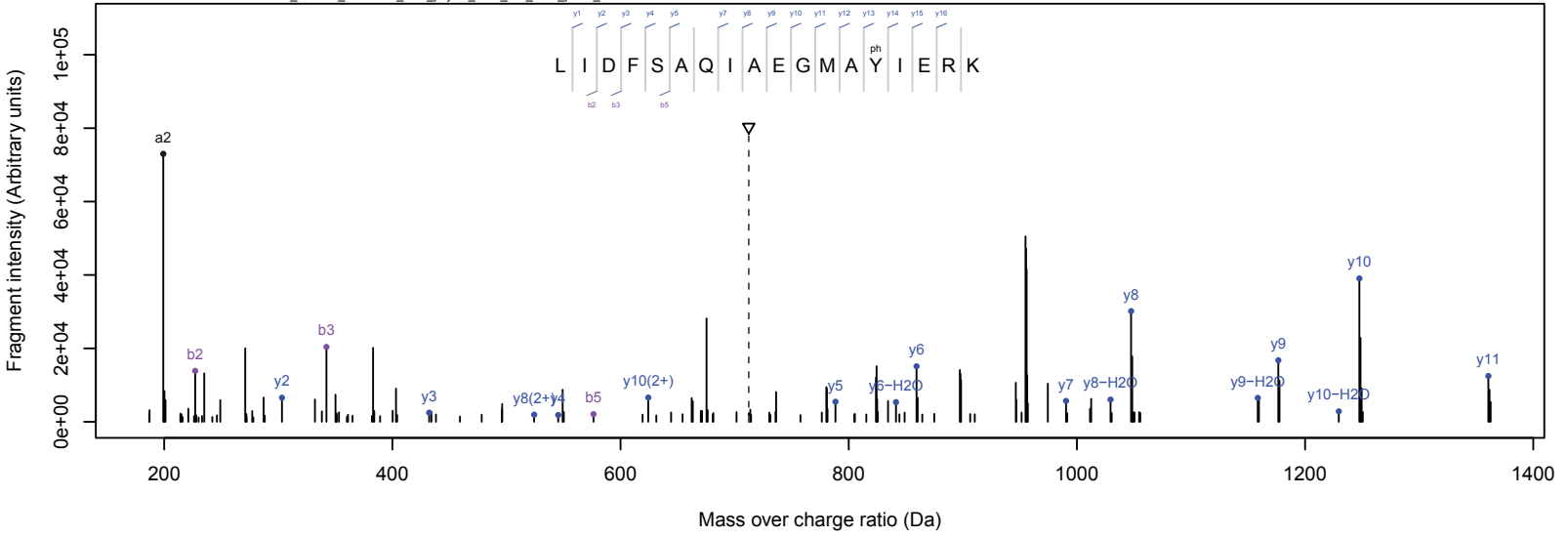
Andromeda score : 102.9



LIDFSAQI AEGMAY IERK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 44071

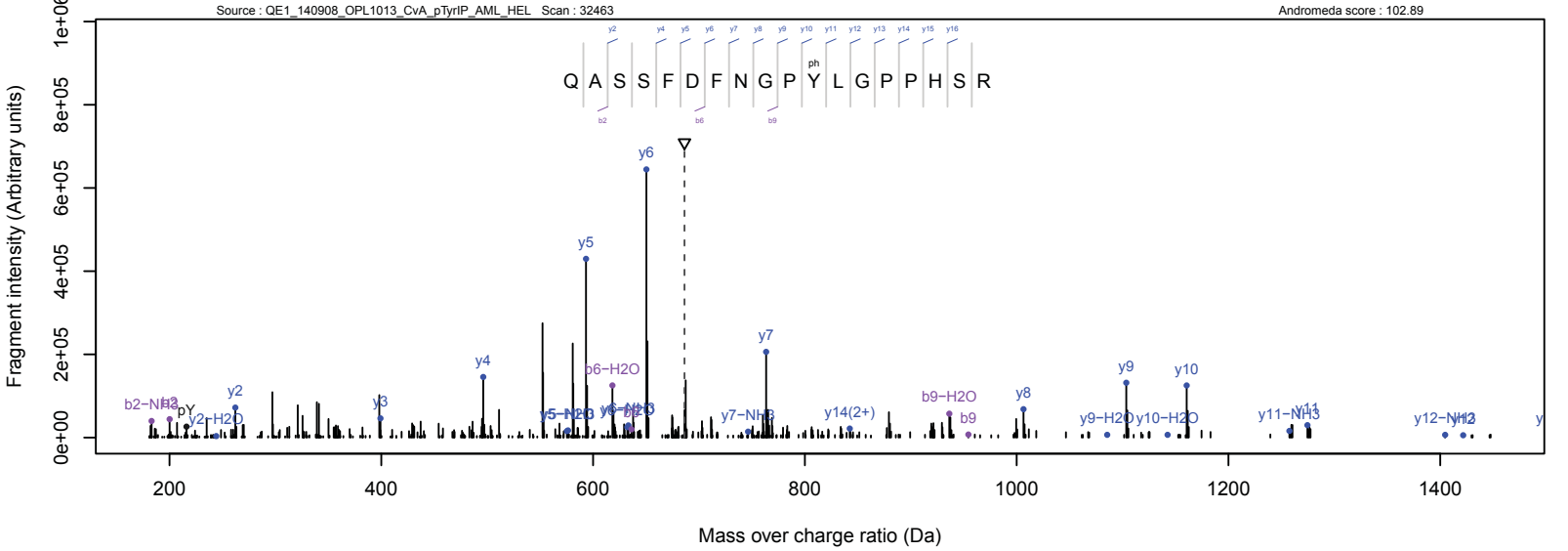
Andromeda score : 102.89



QASSFDNGPY LGPPHSR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32463

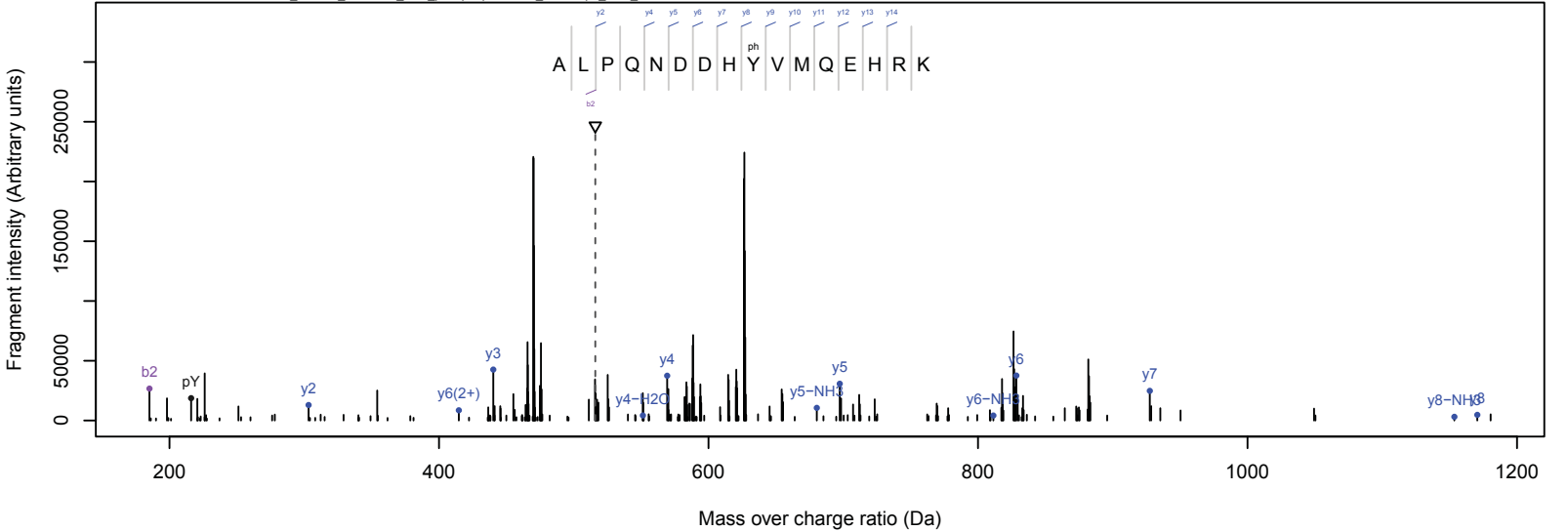
Andromeda score : 102.89



ALPQNDDHY VMQEH RK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 9705

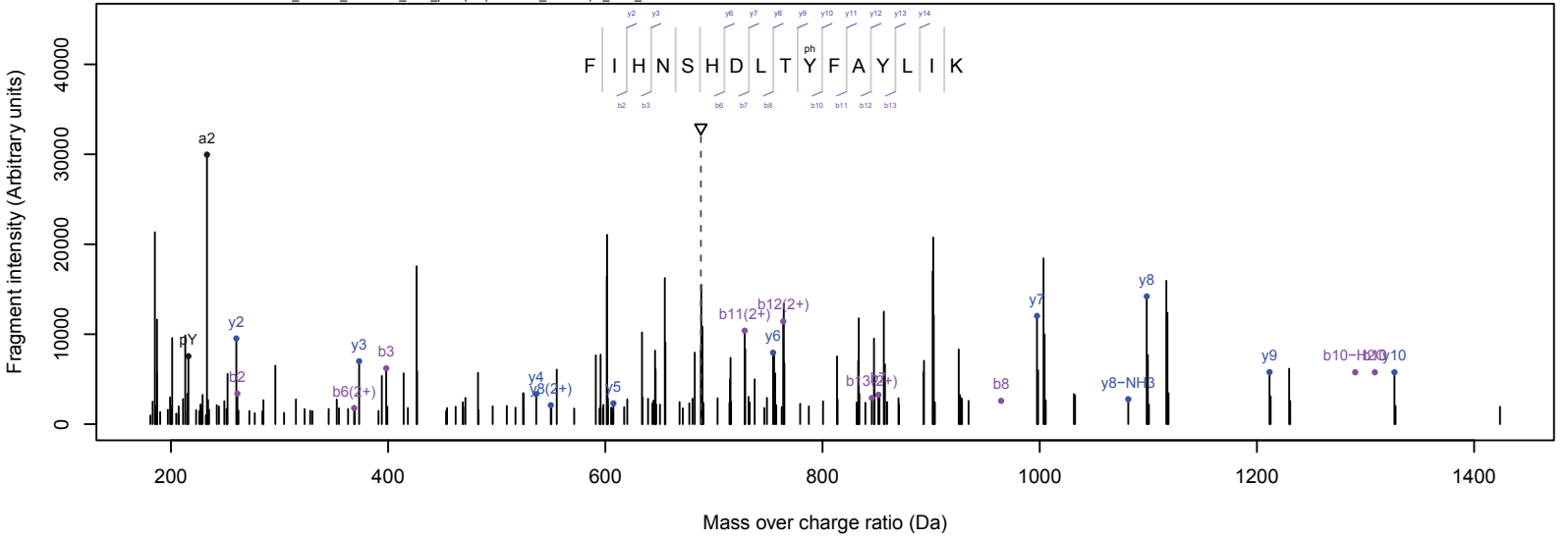
Andromeda score : 102.65



FIHNSHDLTY(ph)FAYLIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 43855

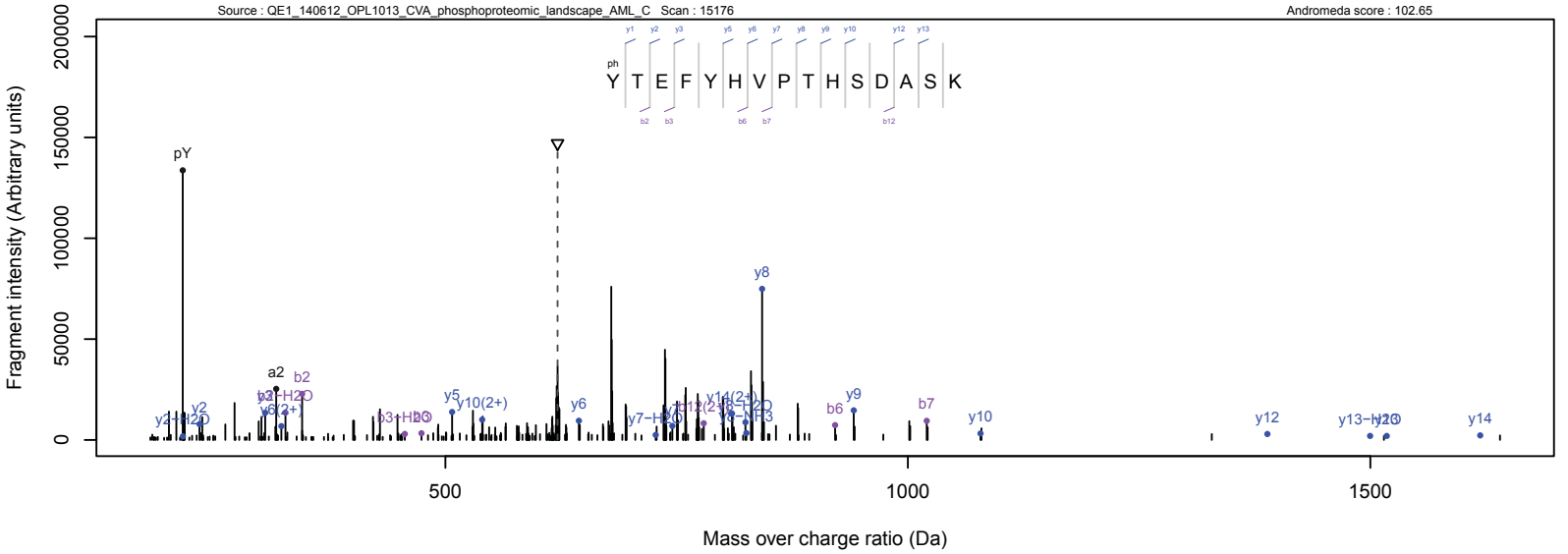
Andromeda score : 102.65



Y(ph)TEFYHVPHTSDASK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15176

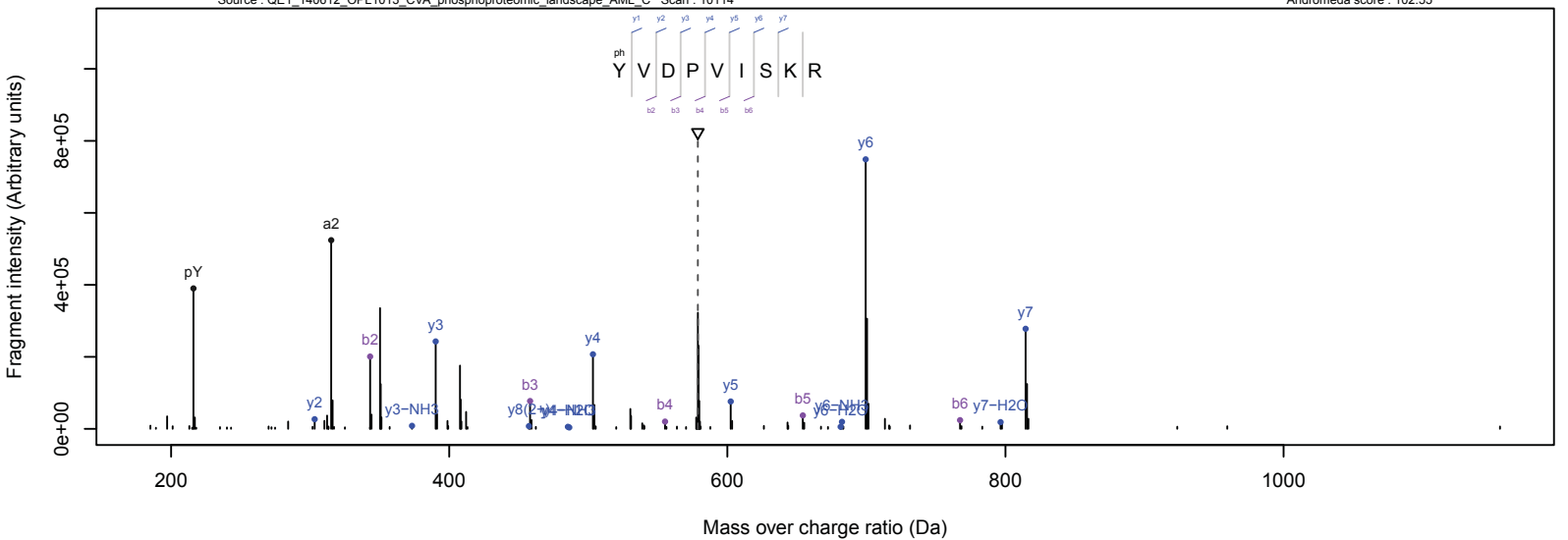
Andromeda score : 102.65



Y(ph)VDPVISKR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10114

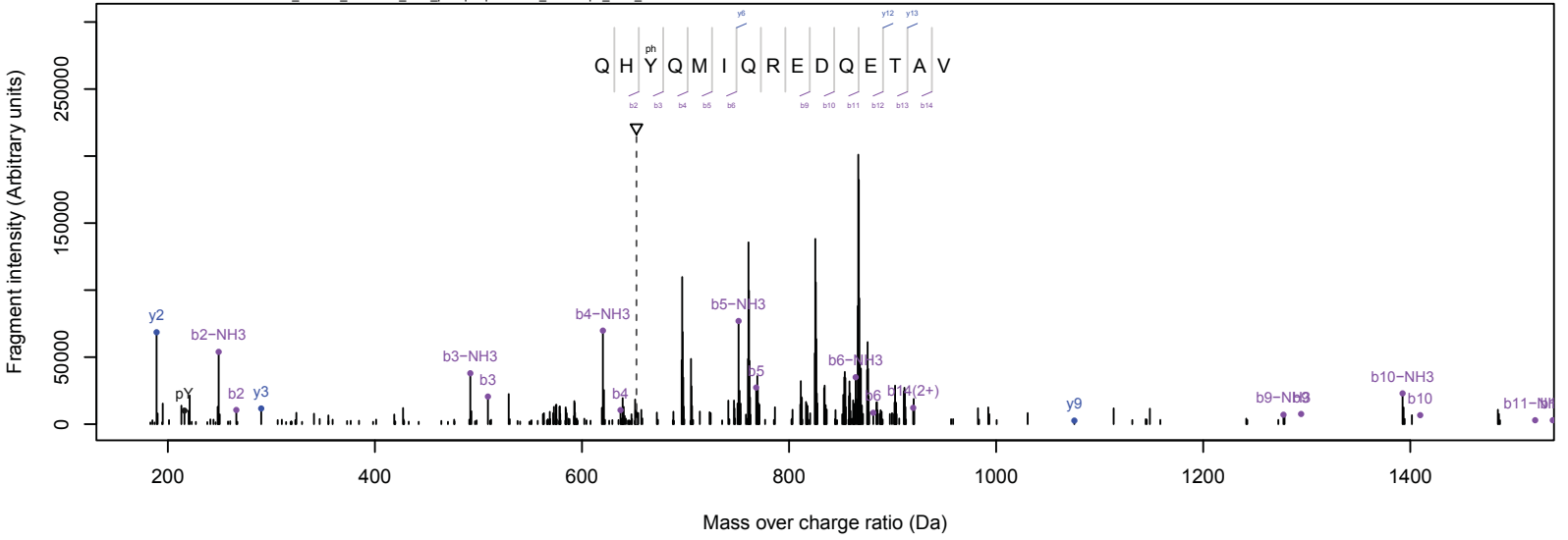
Andromeda score : 102.55



**QHY(ph)QMIQREDQETA V-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13693

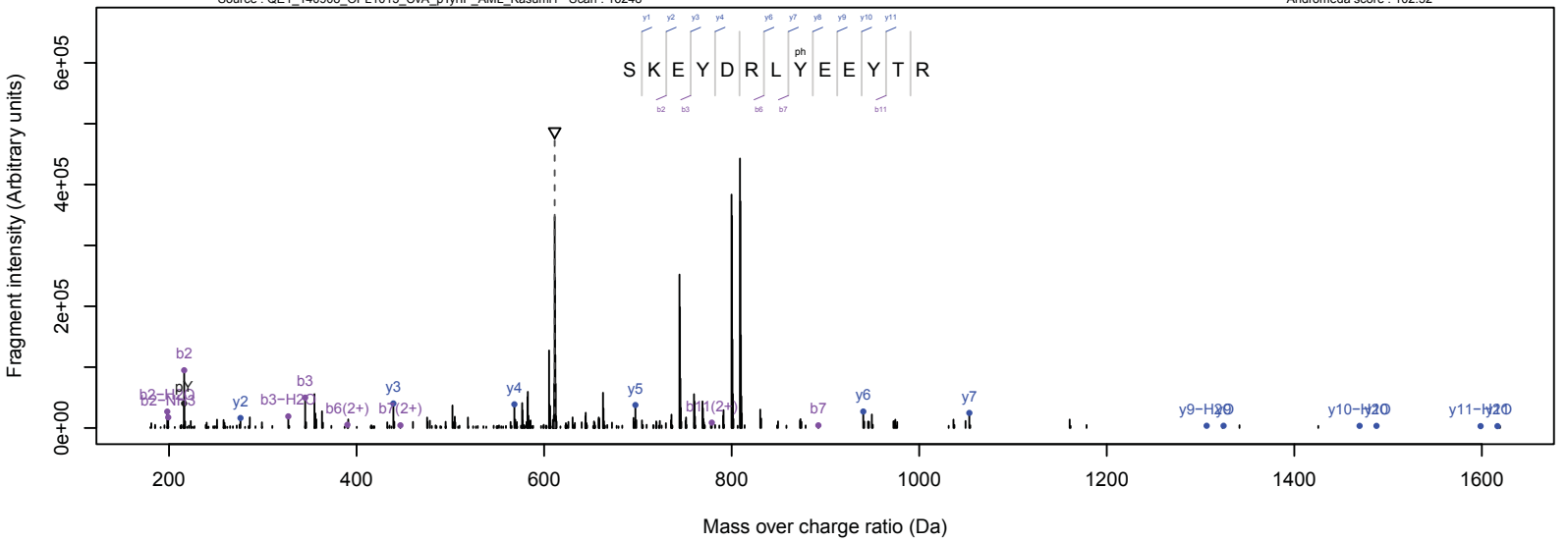
Andromeda score : 102.52



**SKEYDRLY(ph)EEYTR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 16248

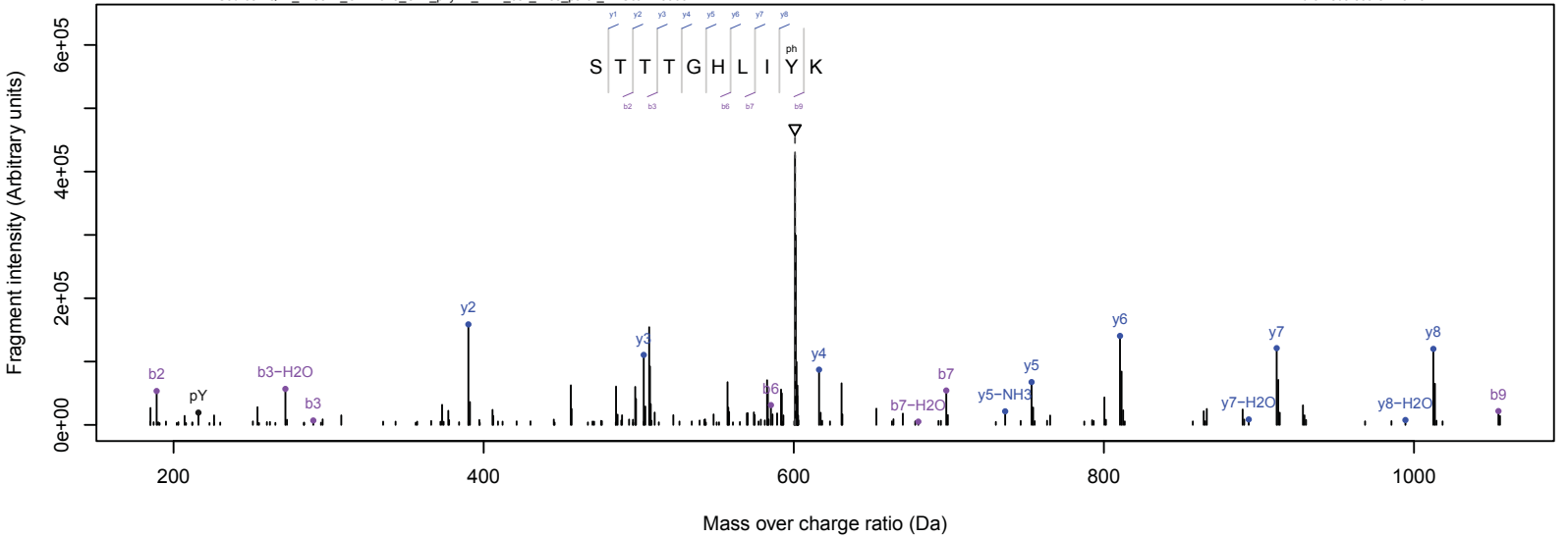
Andromeda score : 102.52



**STTTGHLIY(ph)K-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 5966

Andromeda score : 102.52

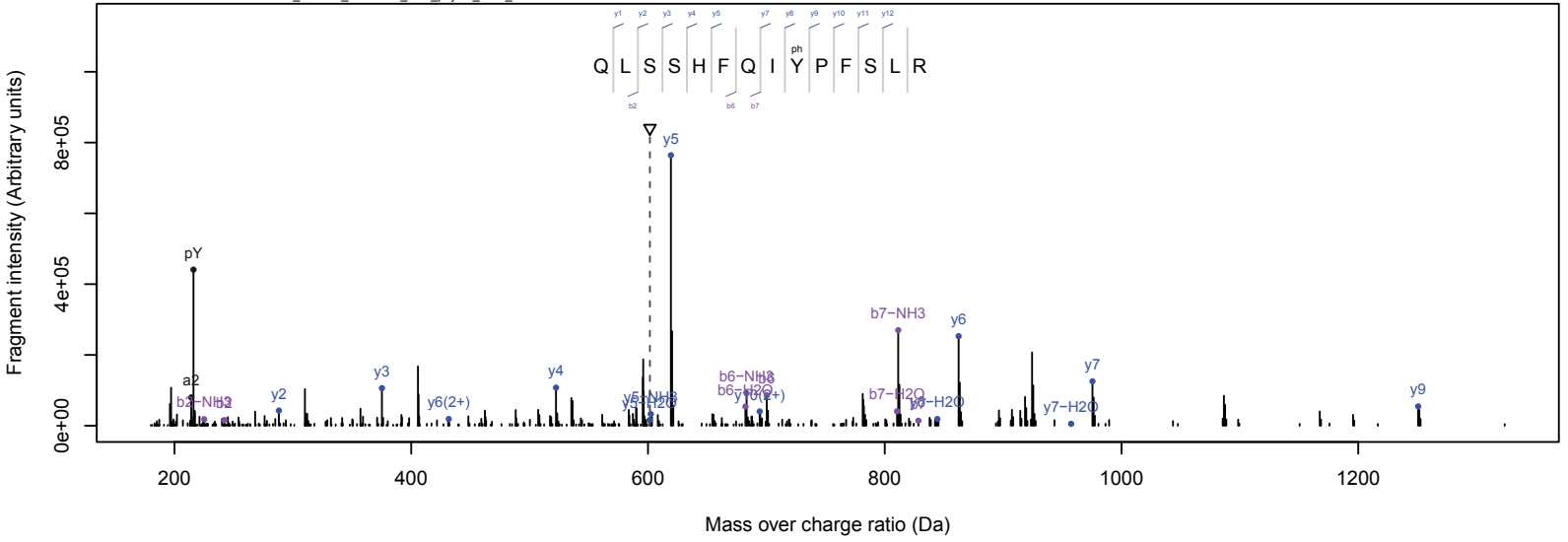




QLSSHFIY(ph)PFSLR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 40245

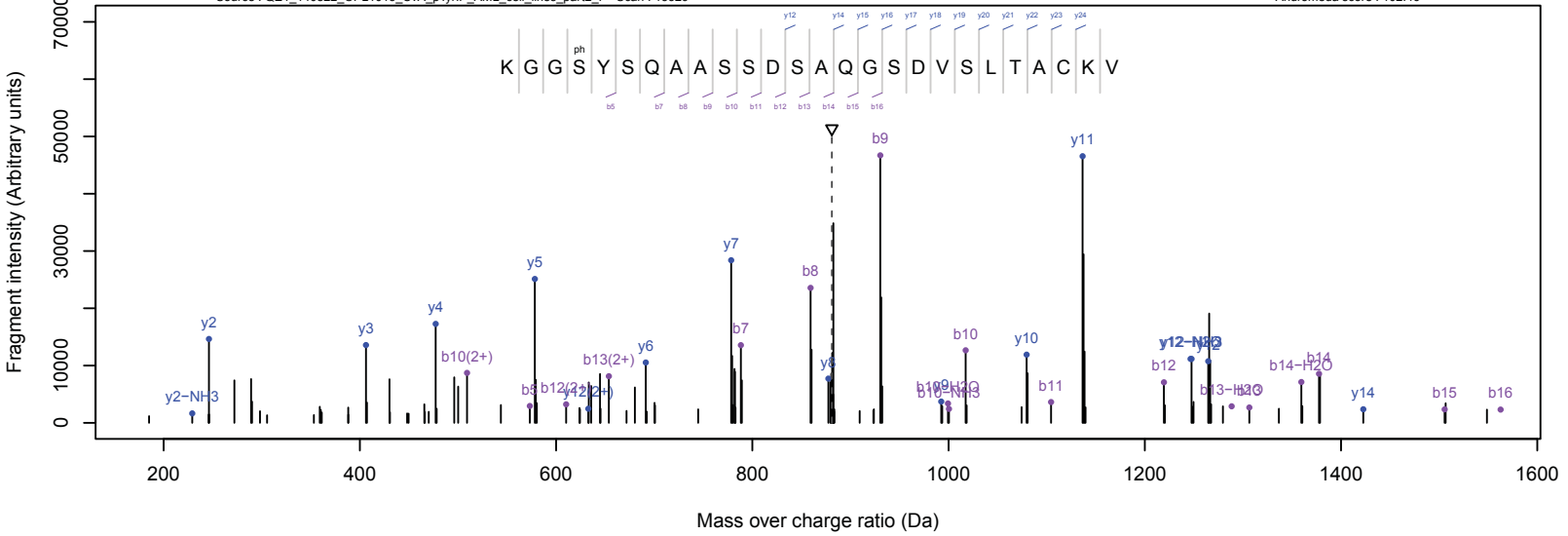
Andromeda score : 102.5



KGGS(ph)YSQAASSDSAQGS DVS LTACKV-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 13329

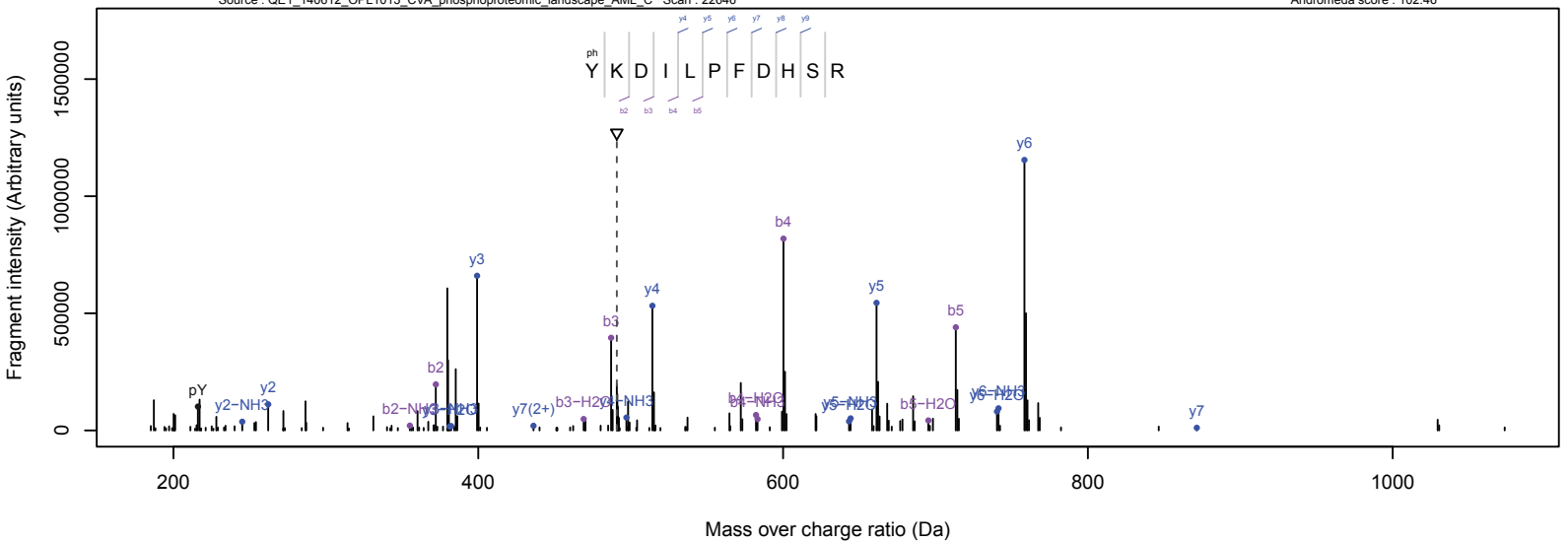
Andromeda score : 102.46



Y(ph)KDILPFDHSR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 22646

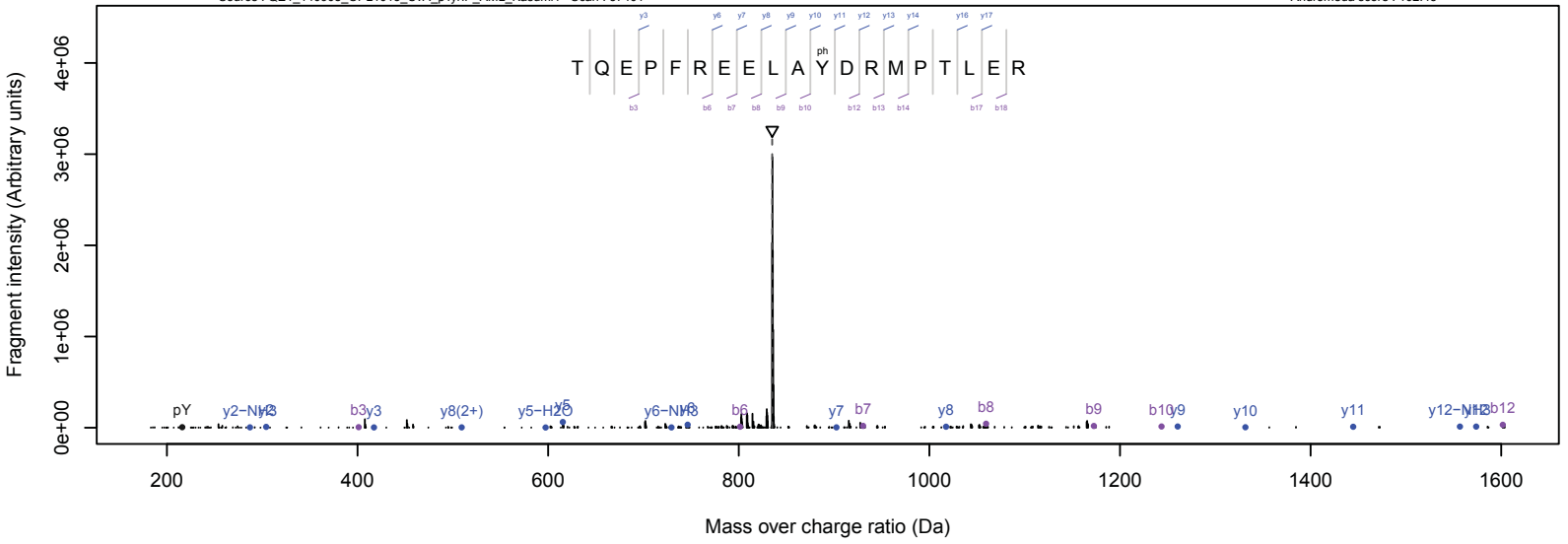
Andromeda score : 102.46



(ac)TQEPFREELAY(ph)DRMPTLER-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 37401

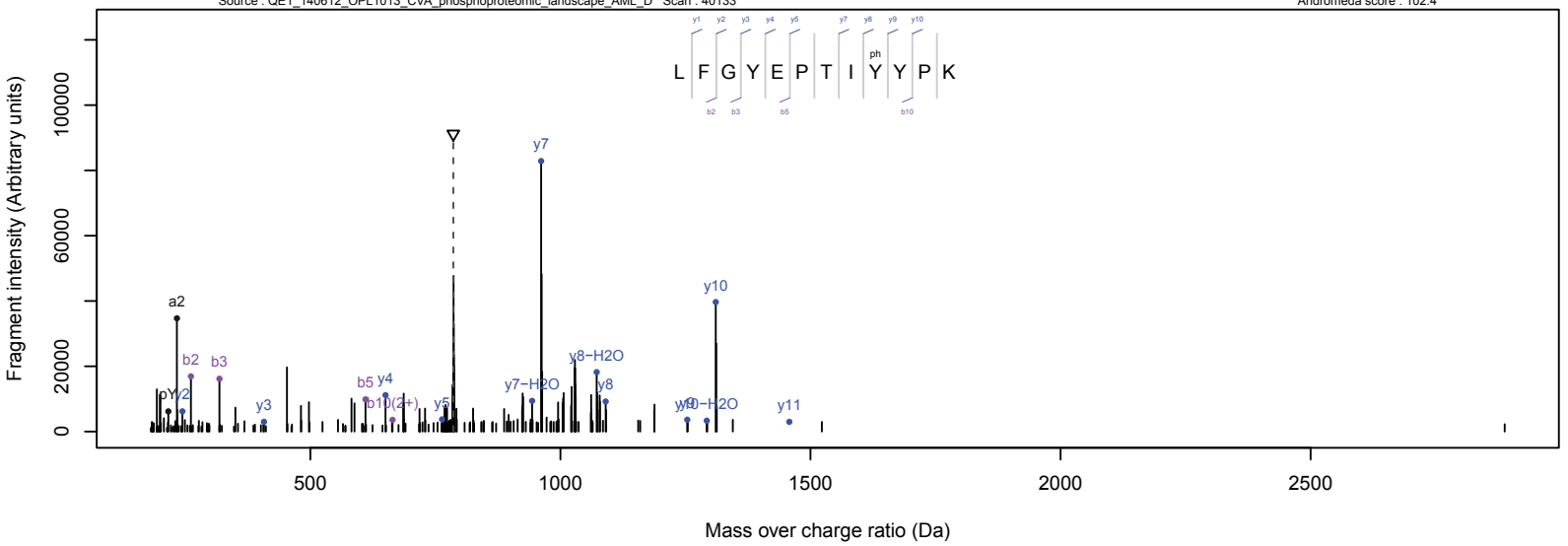
Andromeda score : 102.45



LFGYEPTIYYPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 40133

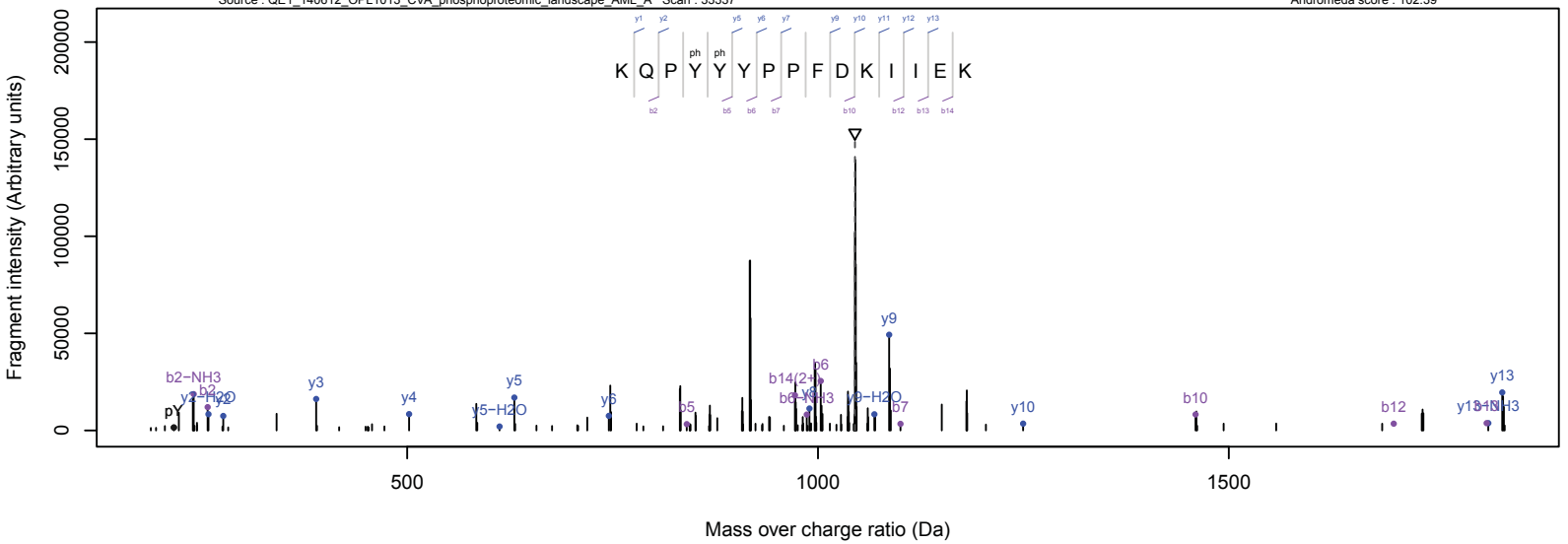
Andromeda score : 102.4

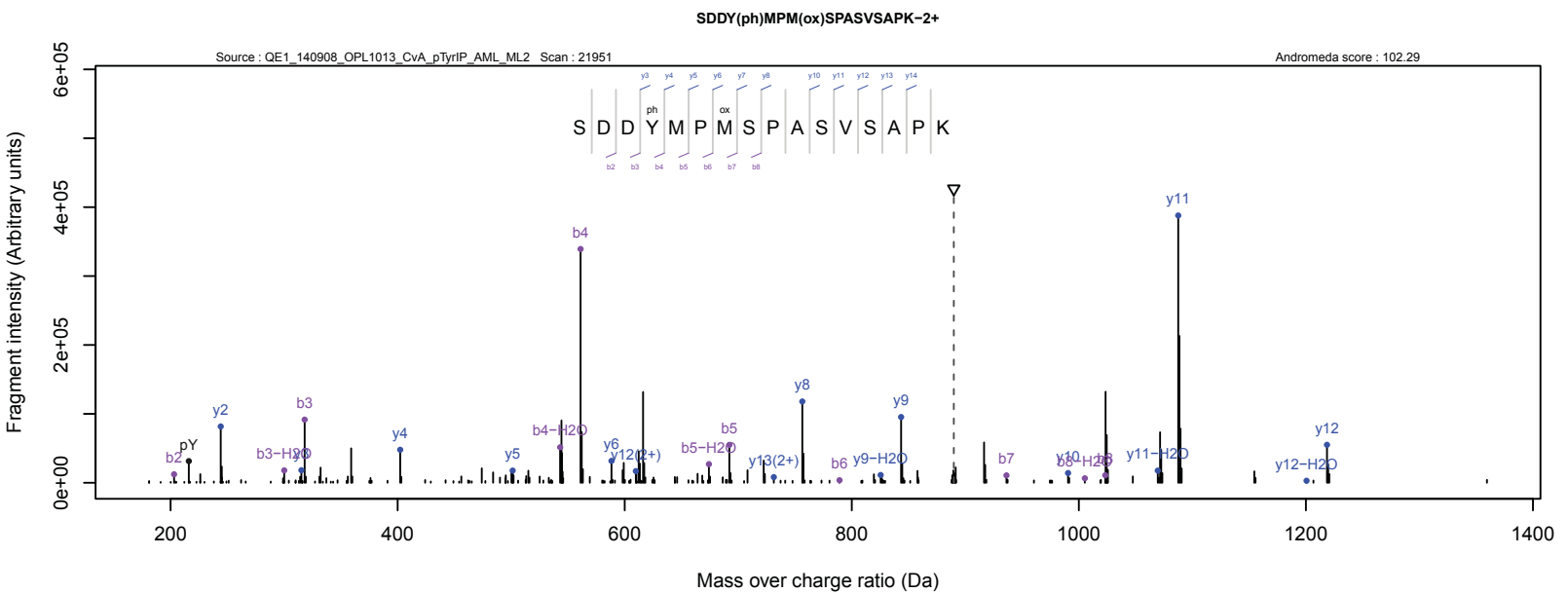
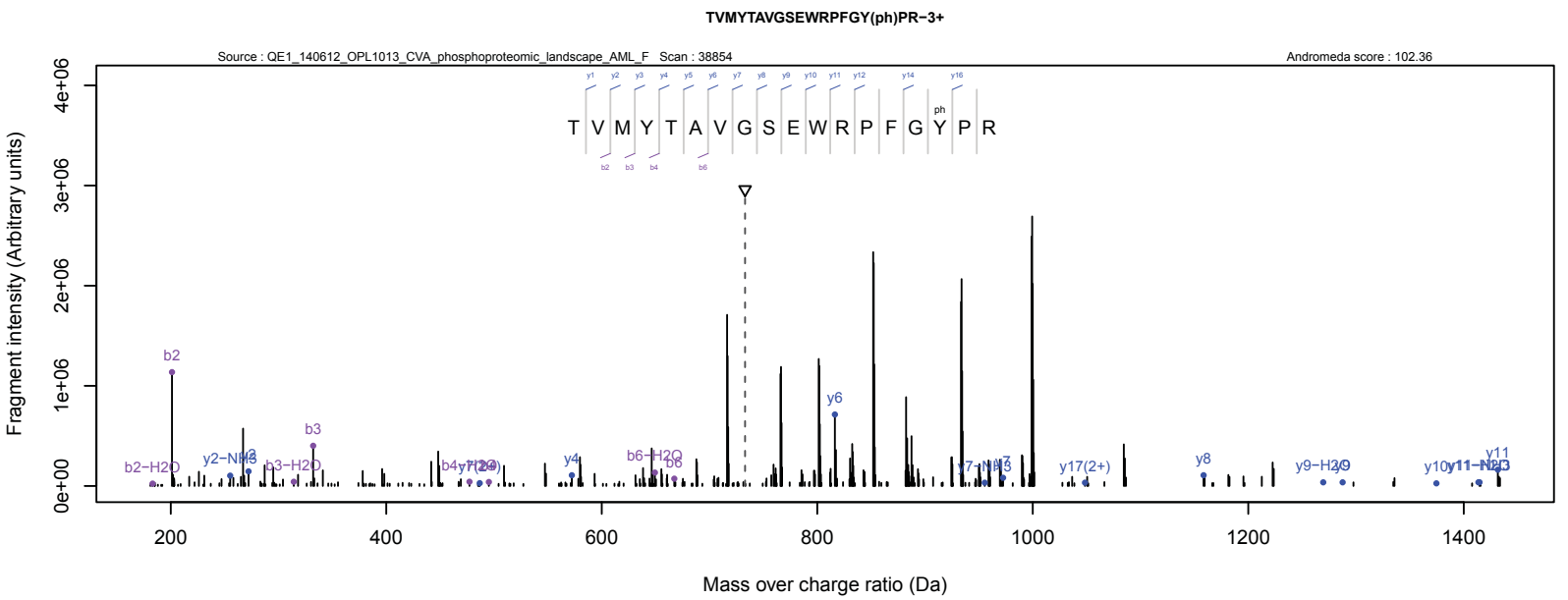
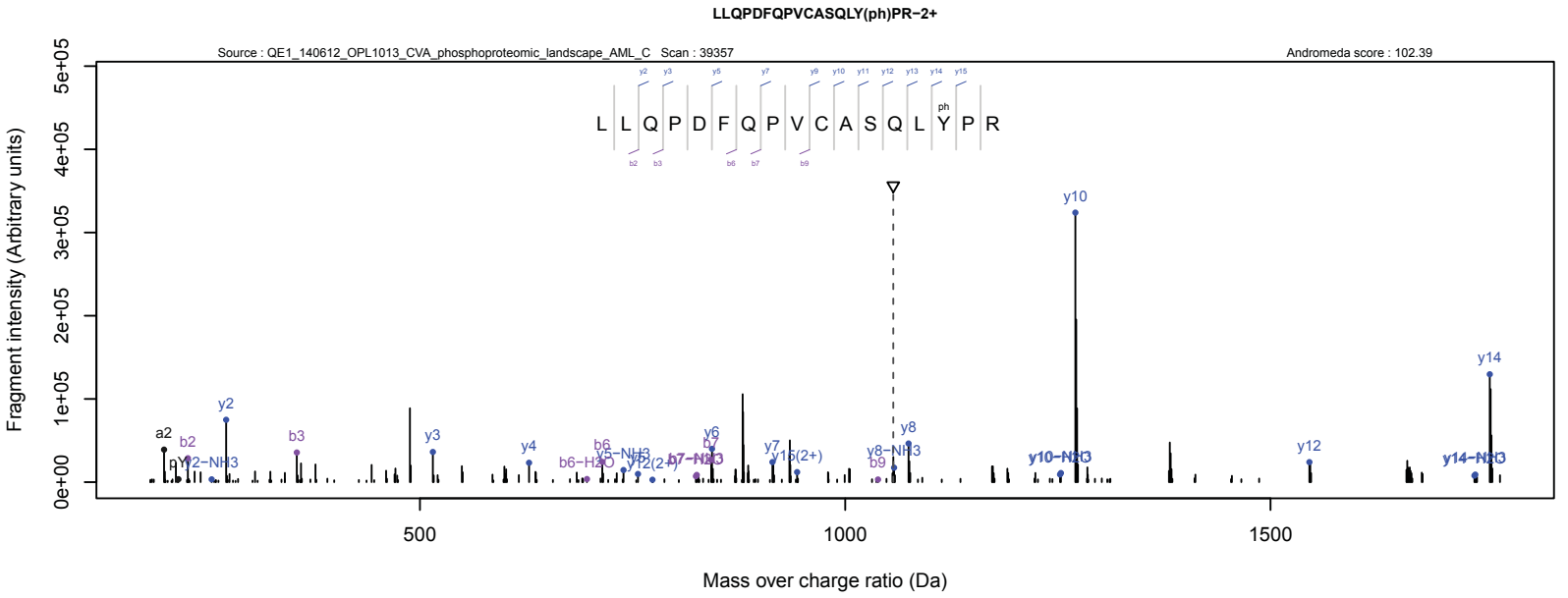


KQPY(ph)YPPFDKIIEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33337

Andromeda score : 102.39

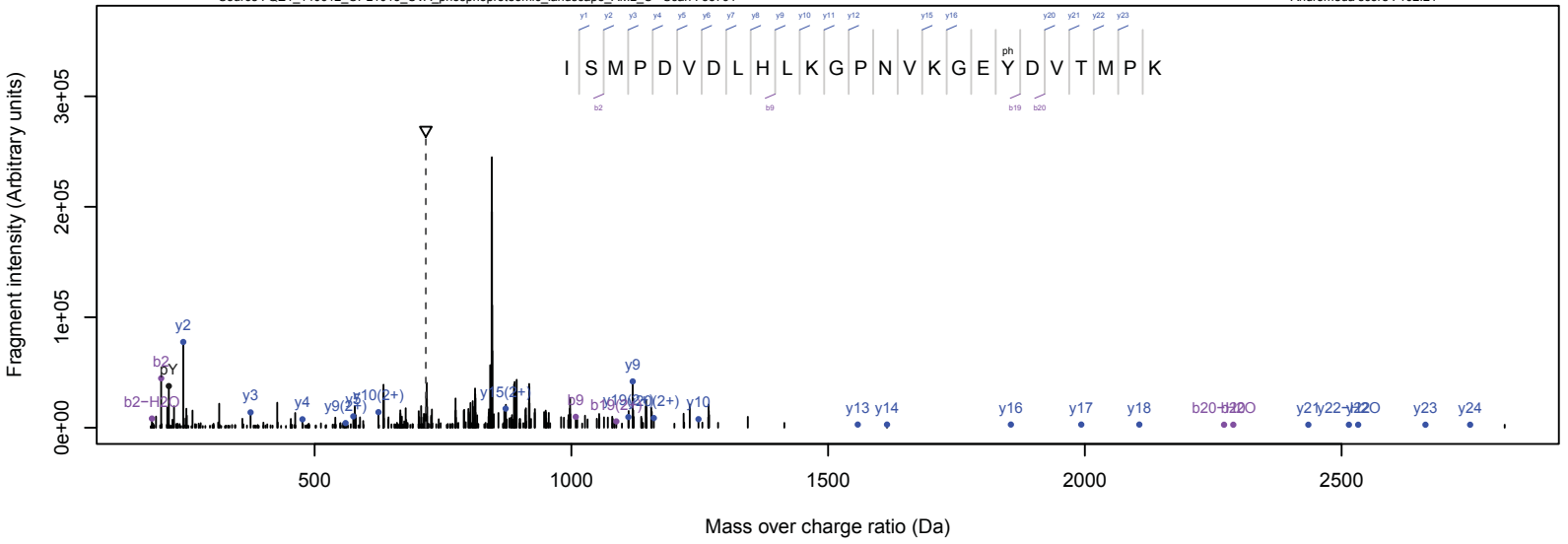




ISMPDVLHLKGNVKGEY(ph)DVTMPK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35791

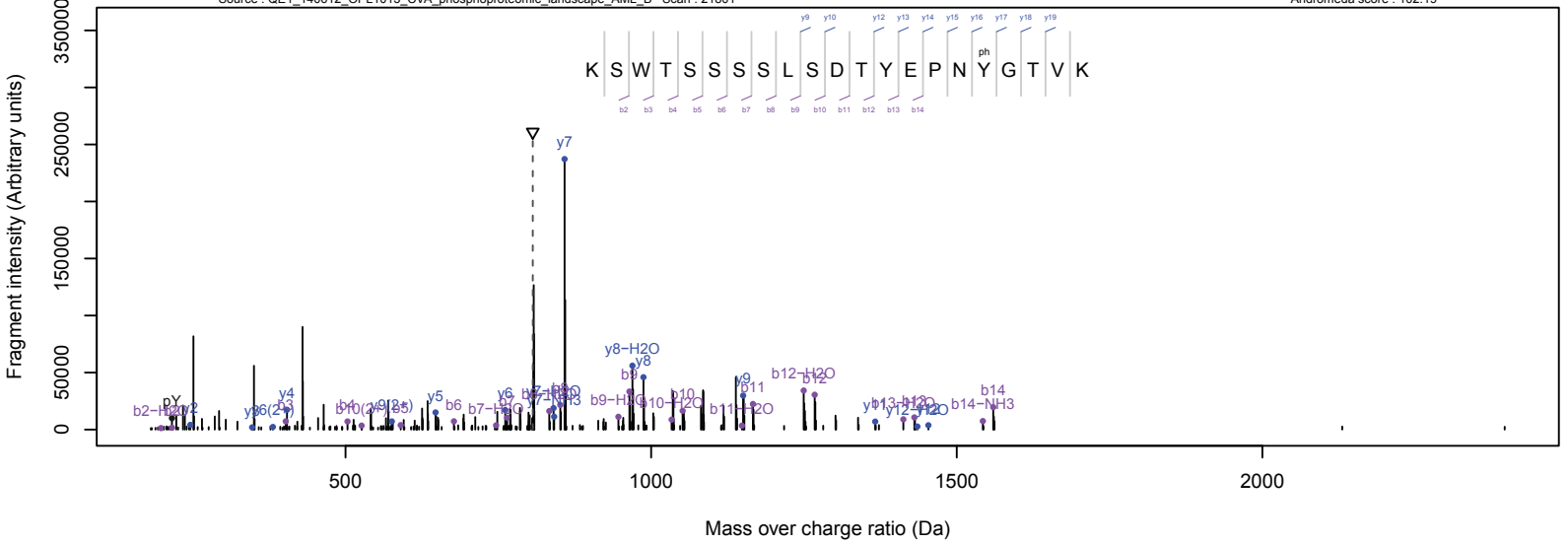
Andromeda score : 102.21



KSWTSSSLSDTYEPNY(ph)GTVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21801

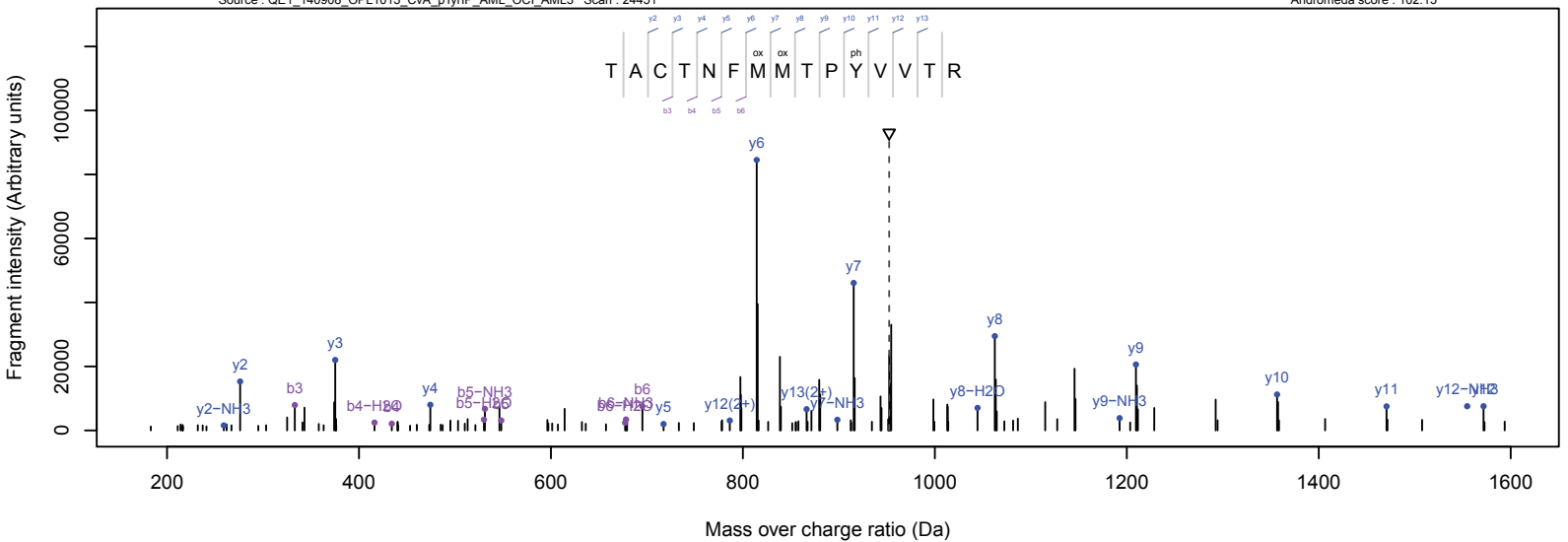
Andromeda score : 102.19



TACTNFM(ox)M(ox)TPY(ph)VVTR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 24451

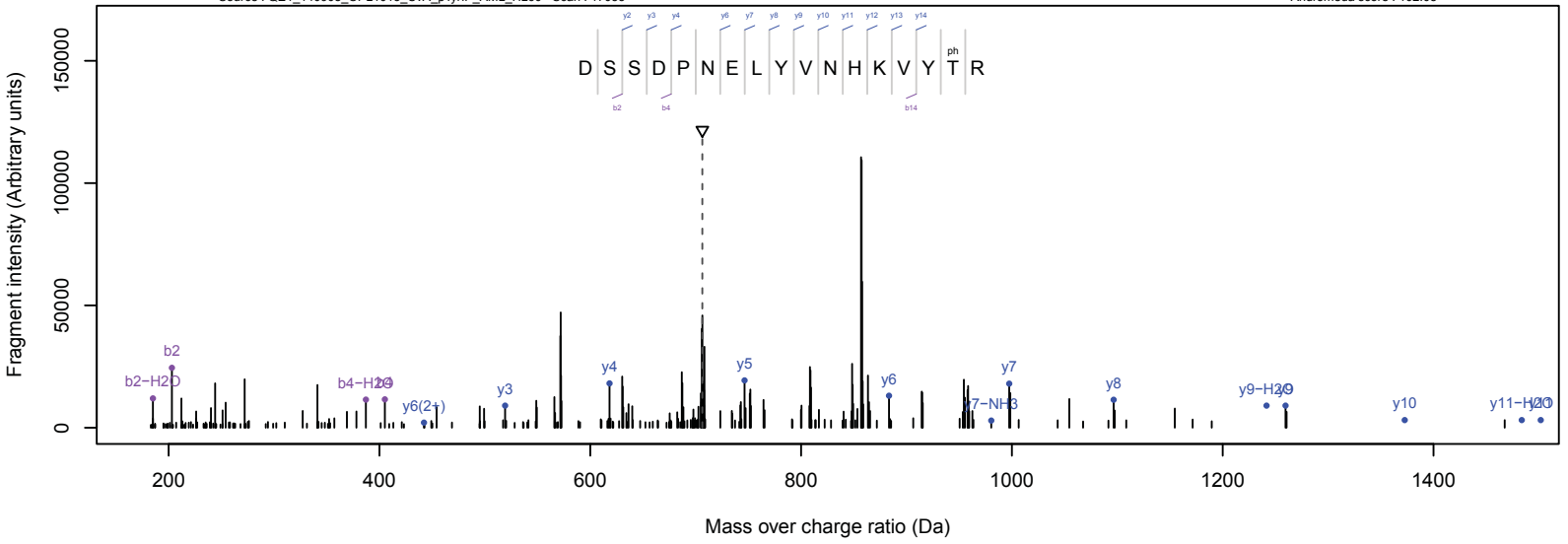
Andromeda score : 102.15



DSSDPNELYVNHKVYTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 17053

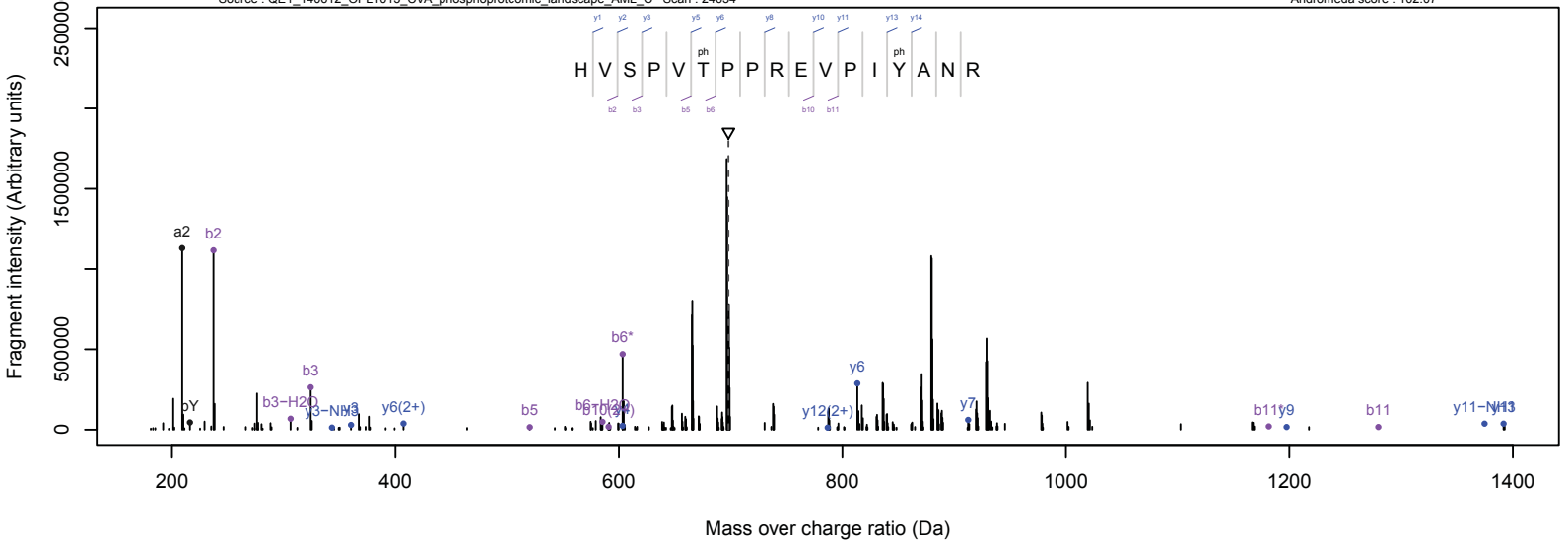
Andromeda score : 102.08



HVSPVT(ph)PPREVPIY(ph)ANR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 24634

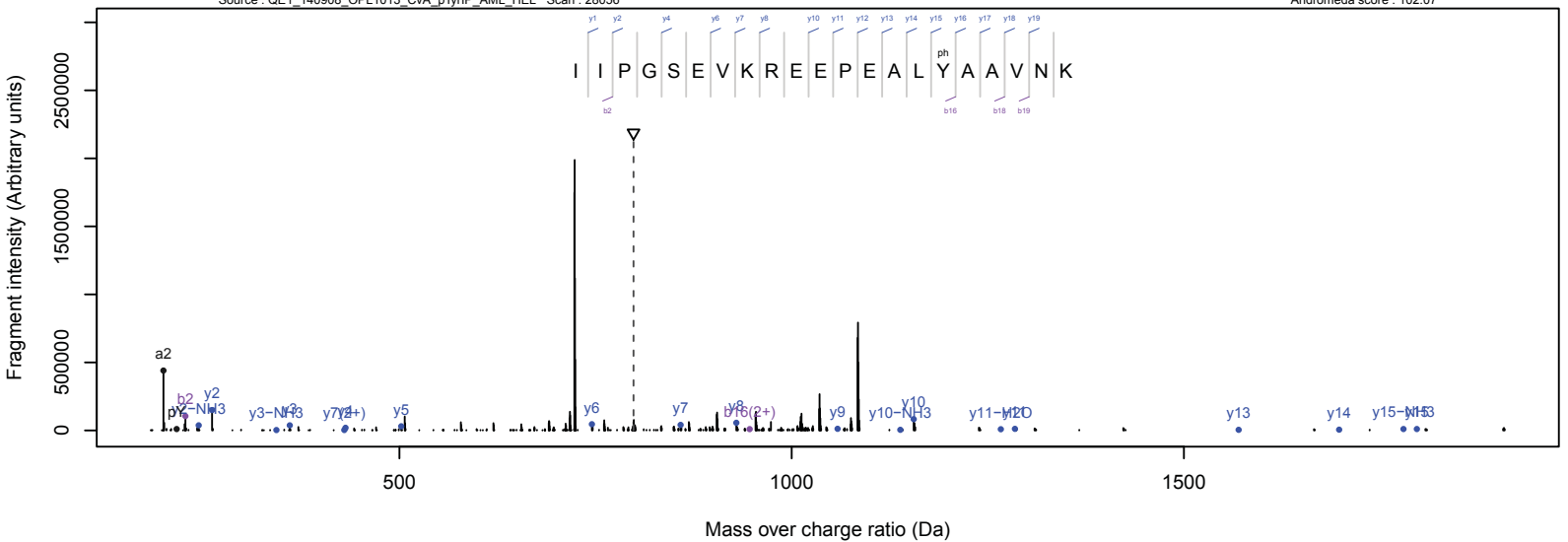
Andromeda score : 102.07



IIPGSEVKREEPEALYAAVNK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 28056

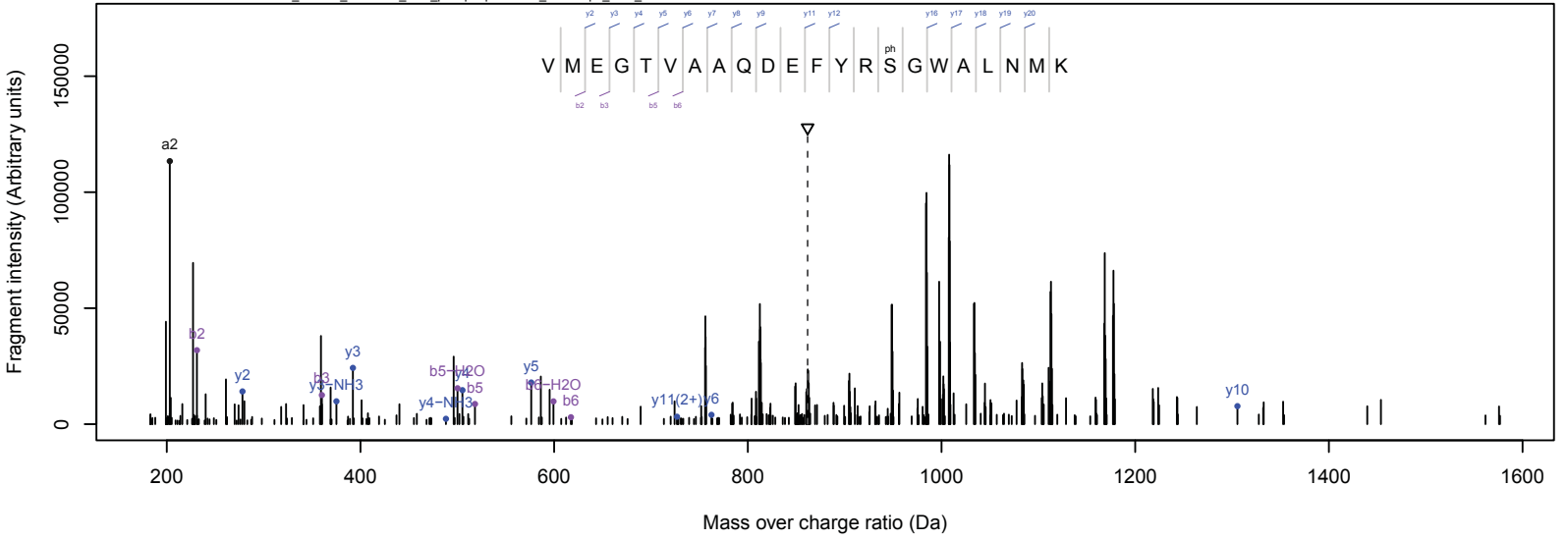
Andromeda score : 102.07



VMEGTVA A Q D E F Y R S G W A L N M K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38545

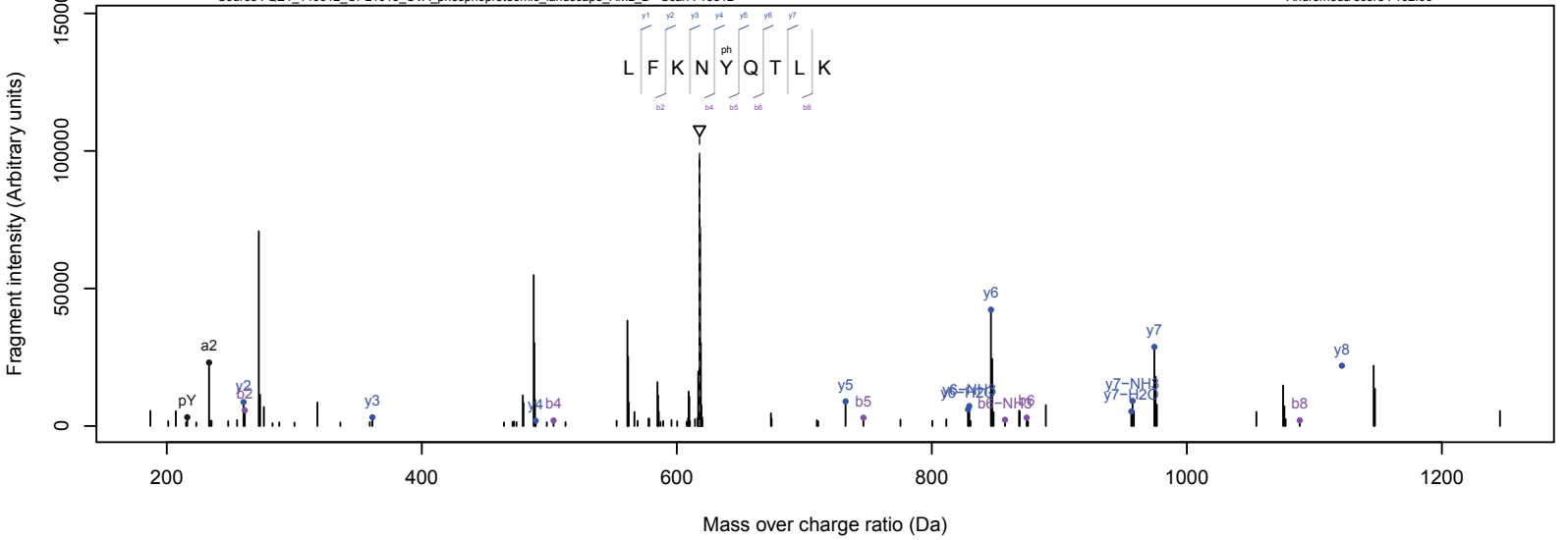
Andromeda score : 102.07



LFKNY(ph)QTLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 13812

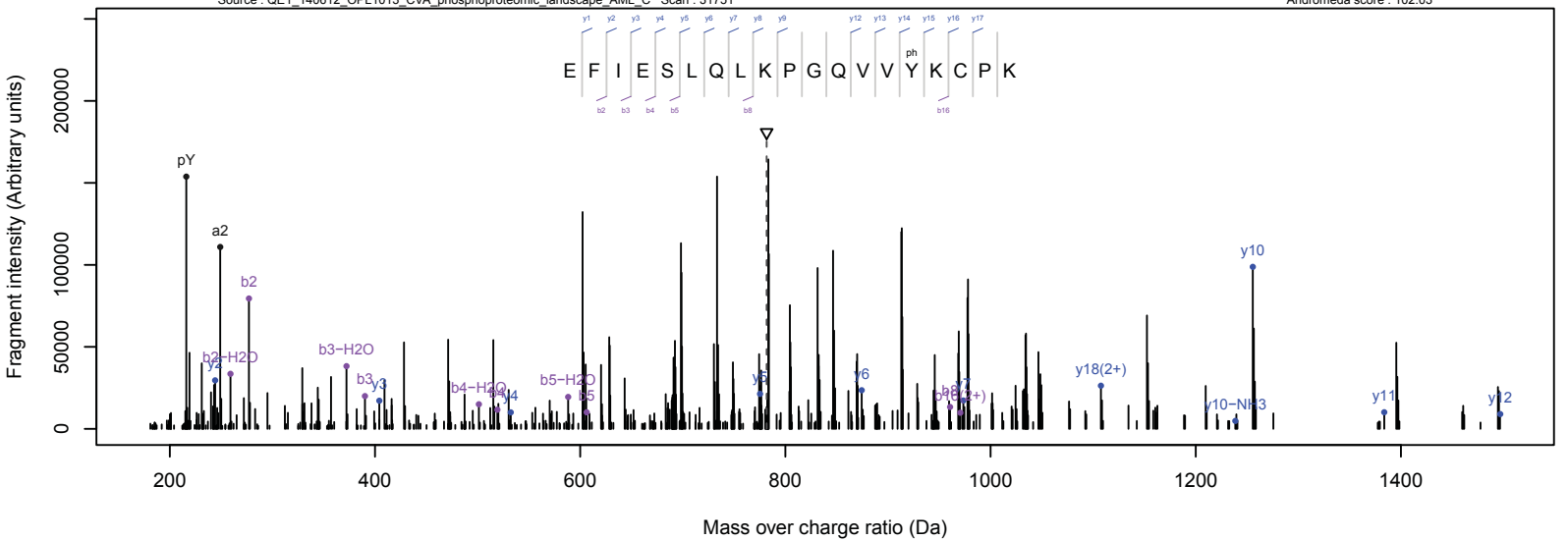
Andromeda score : 102.06



EFIESLQLKPGVVY(ph)KCPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31751

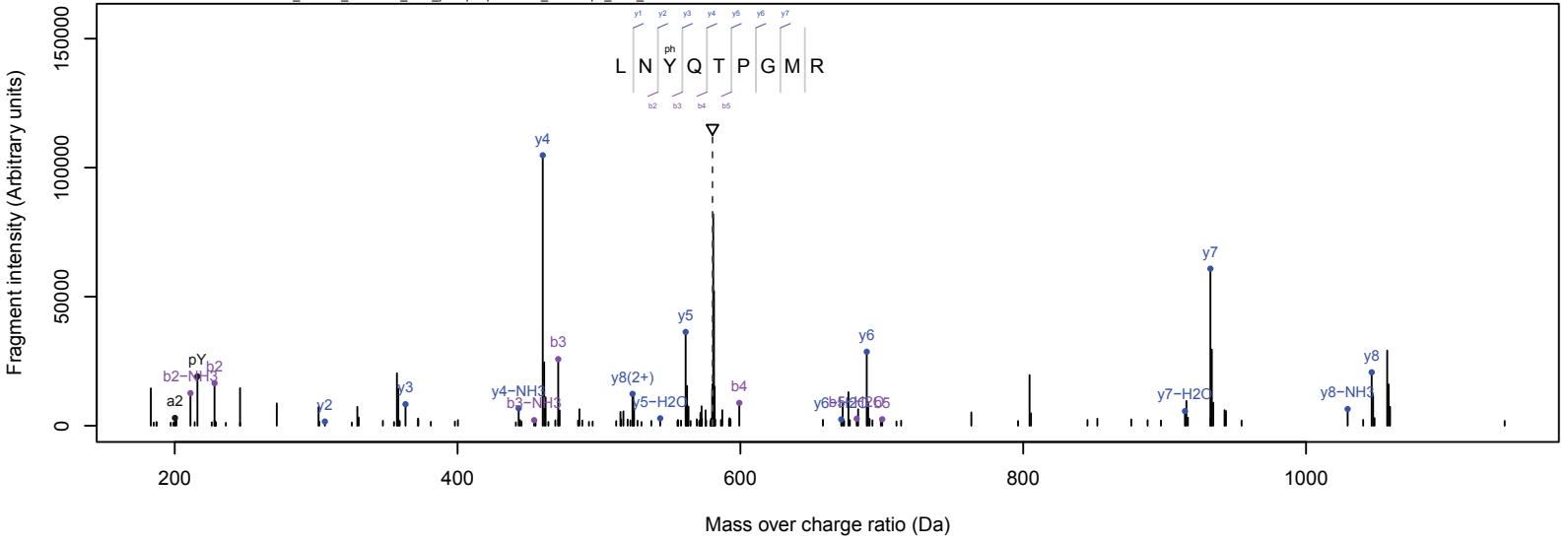
Andromeda score : 102.03



LNY(ph)QTPGMR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9775

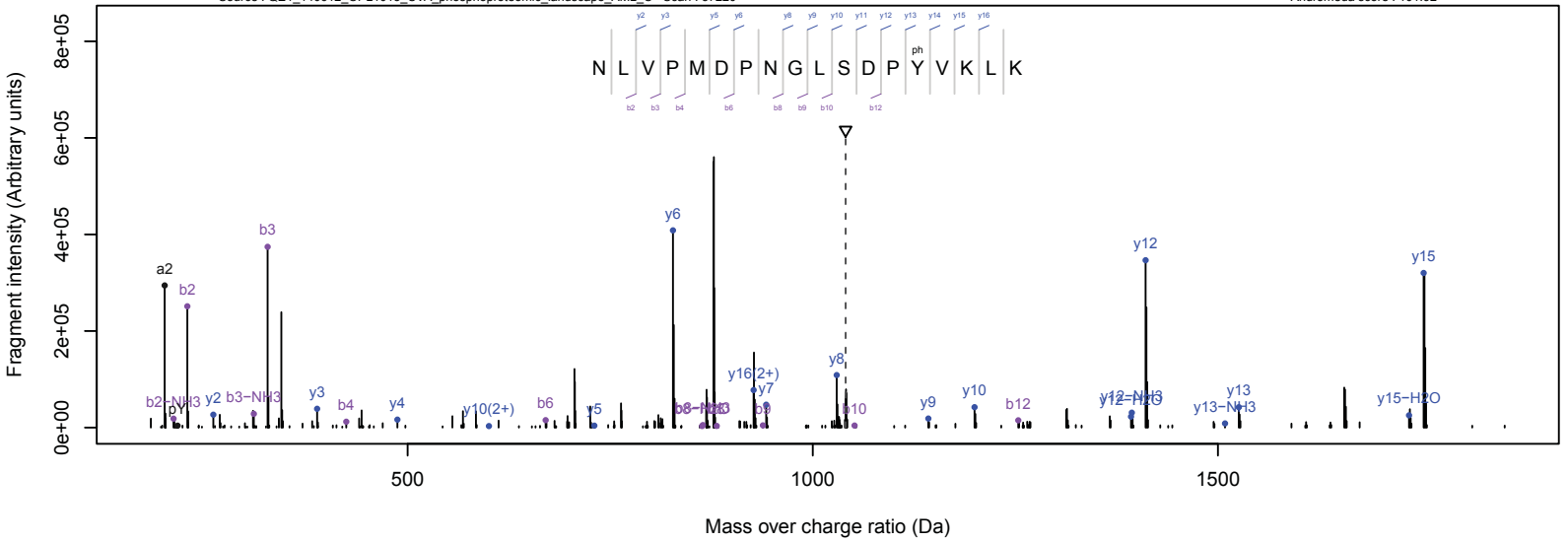
Andromeda score : 102.01



NLVMPDPNGLSDPY(ph)VKLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37220

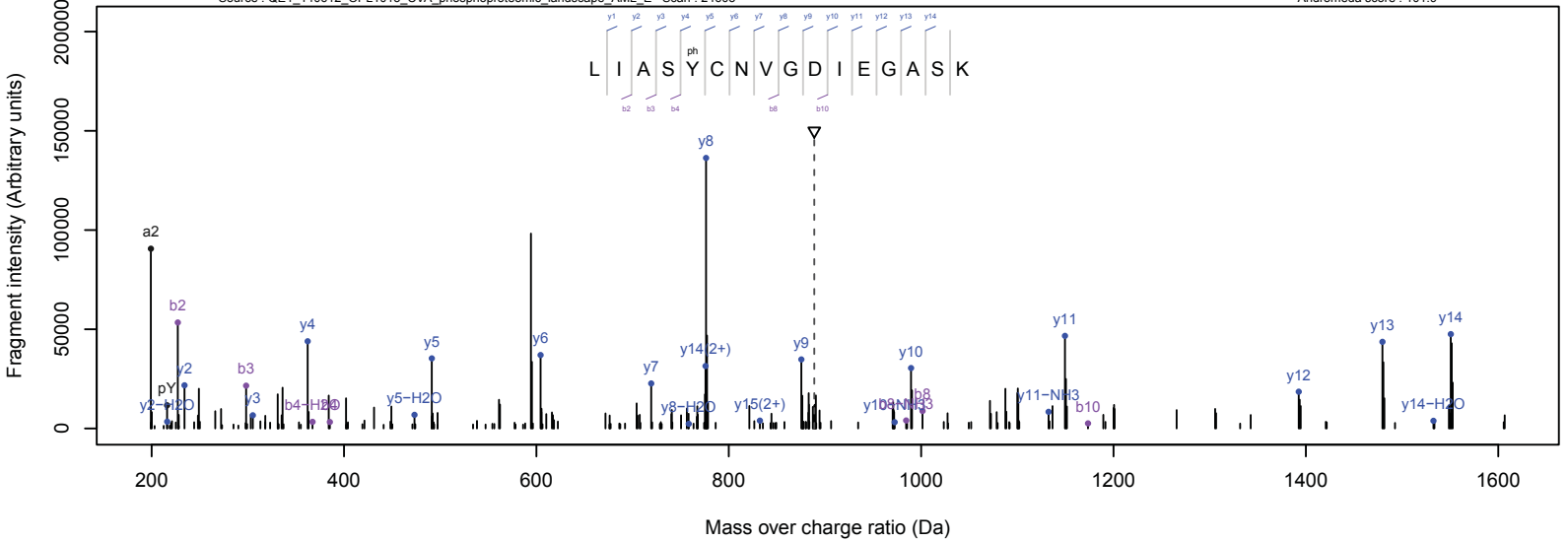
Andromeda score : 101.92



LIASY(ph)CNVGDIEGASK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 24603

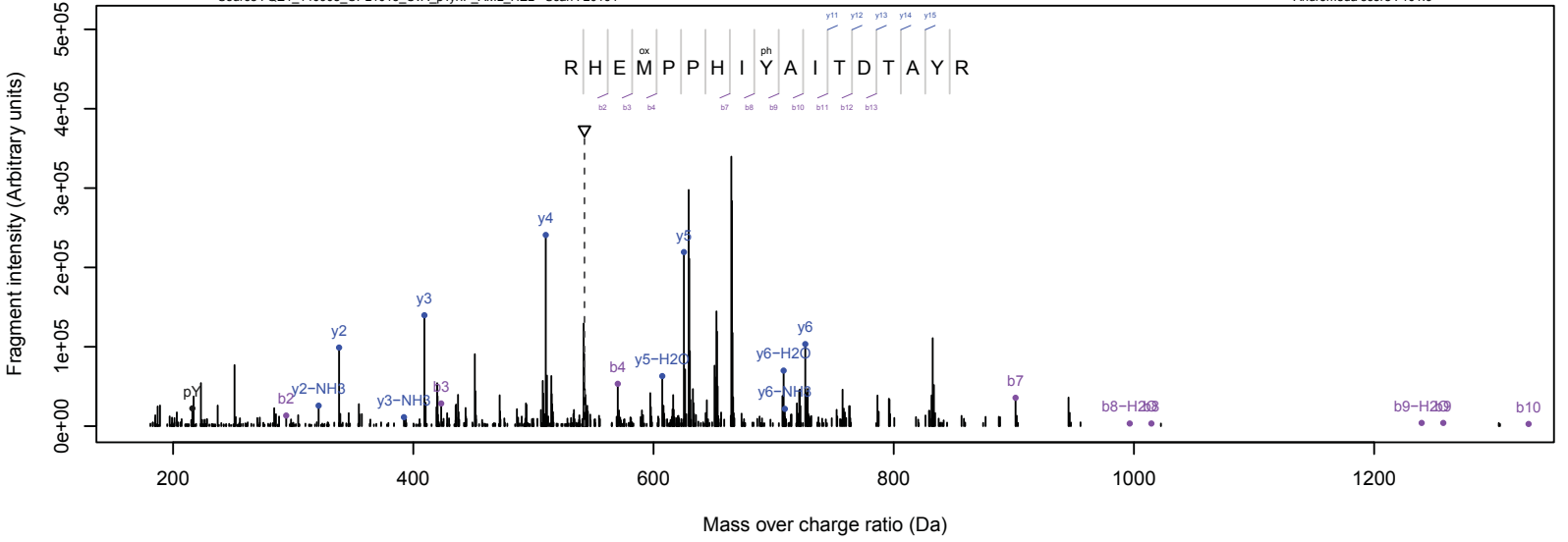
Andromeda score : 101.9



RHEM(ox)PPIHY(ph)AITDTAYR-4+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 20164

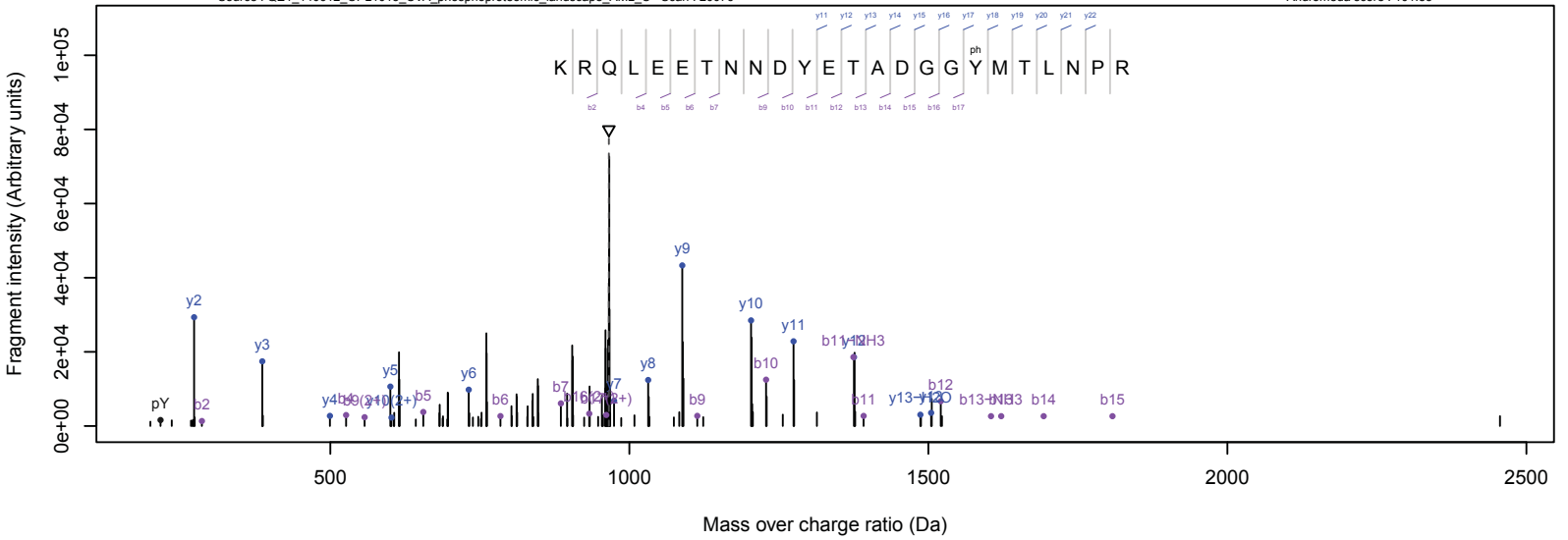
Andromeda score : 101.9



KRQLEETNNDYETADGGY(ph)MTLNPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20076

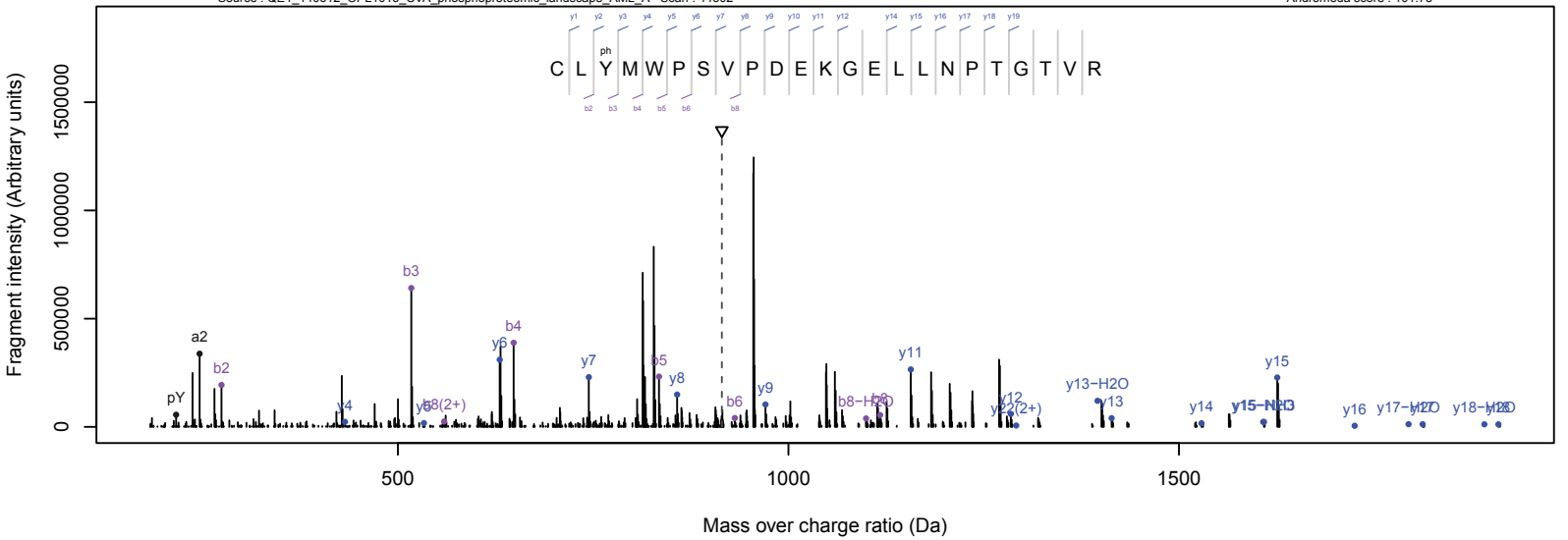
Andromeda score : 101.85



CLY(ph)MWPSVPDEKGE LLNPTGTVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 44302

Andromeda score : 101.75

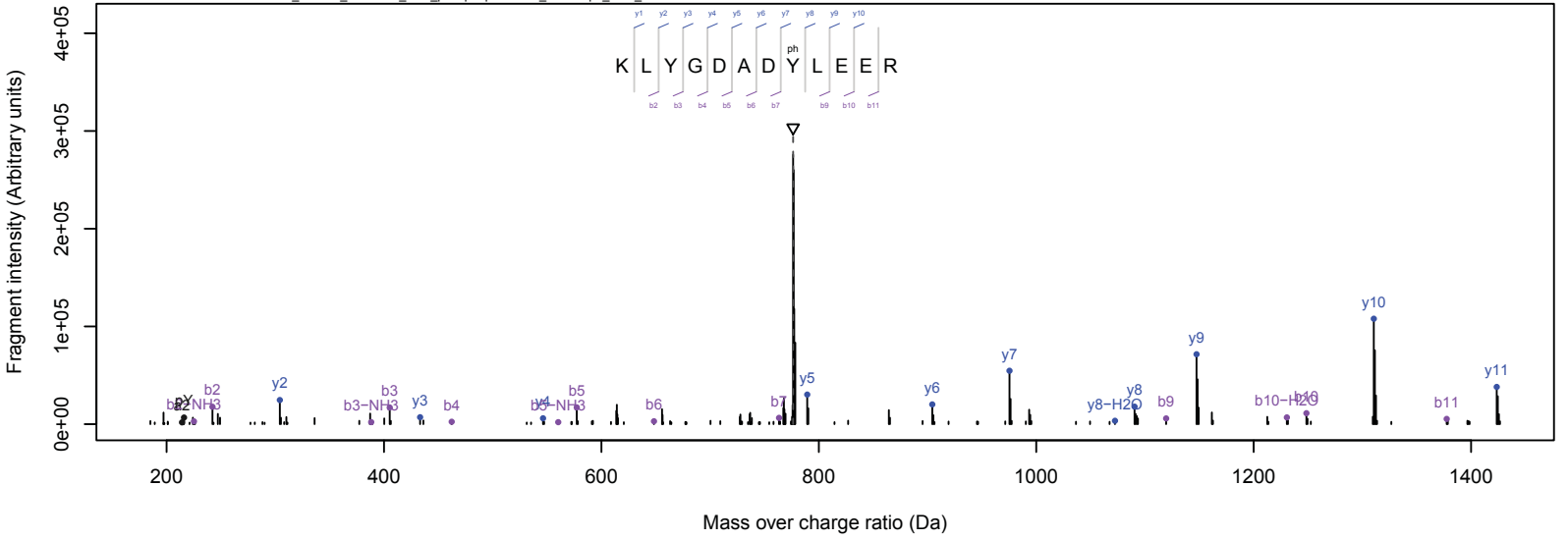




KLYGDADY(ph)LEER-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18242

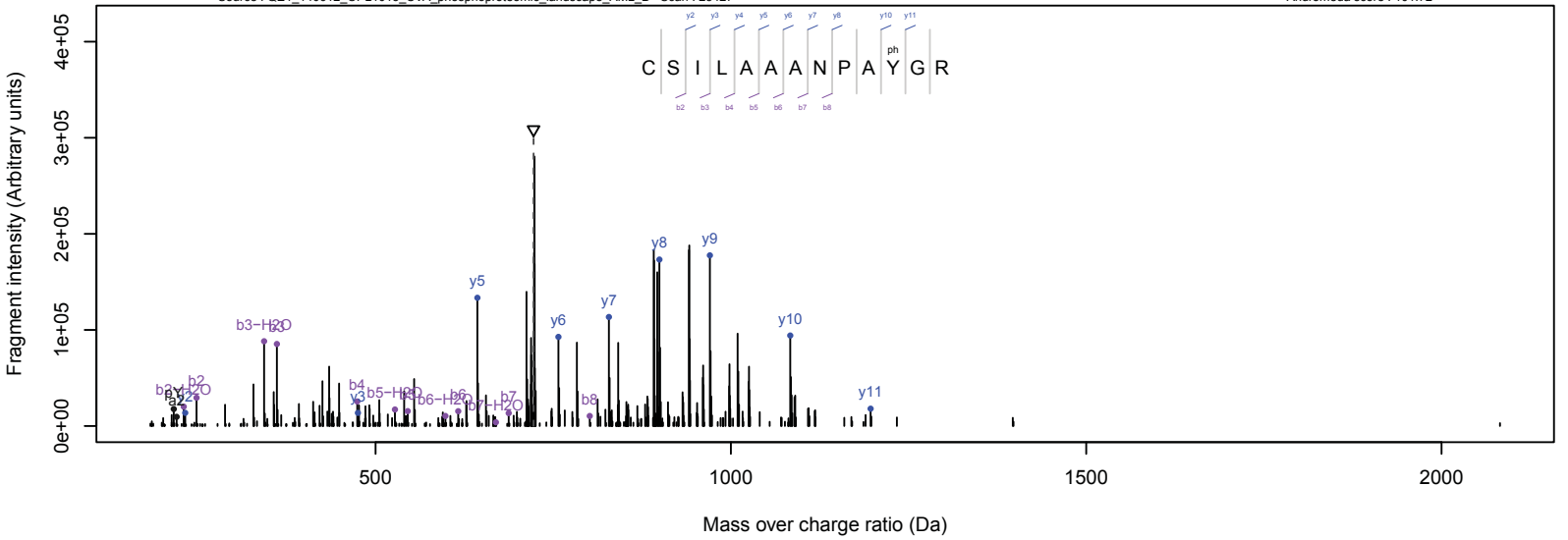
Andromeda score: 101.75



CSILAAANPAY(ph)GR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 25427

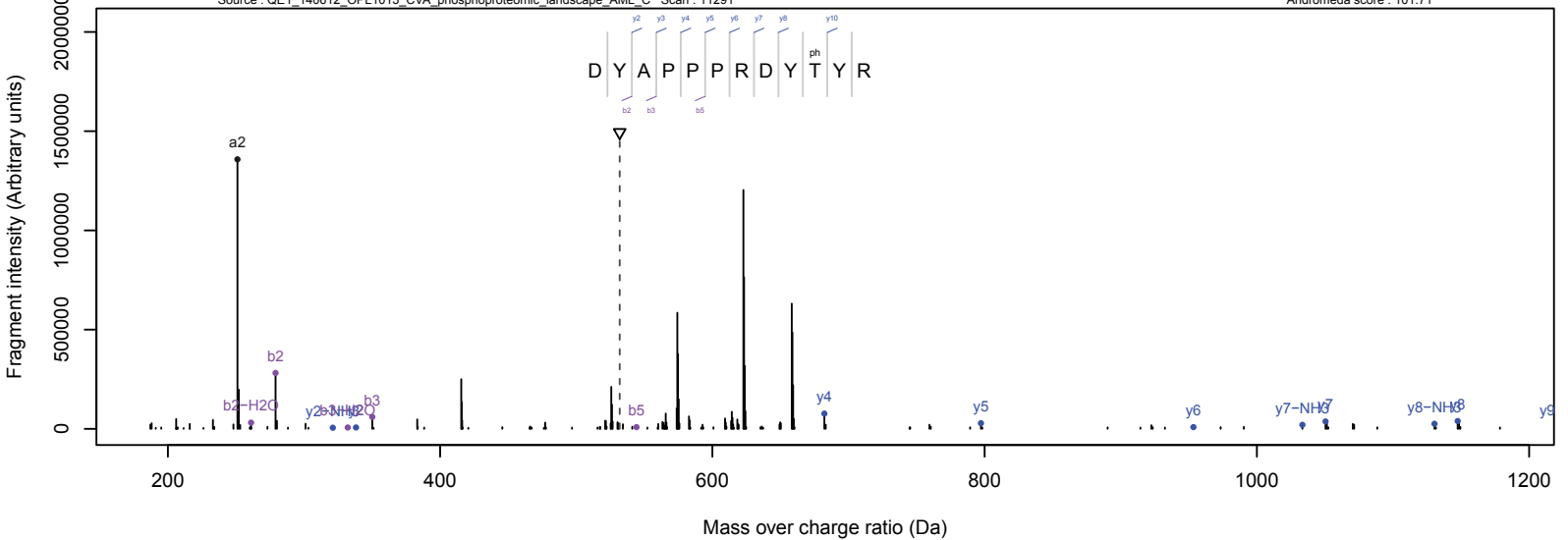
Andromeda score: 101.72



DYAPPPRDY(ph)YR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11291

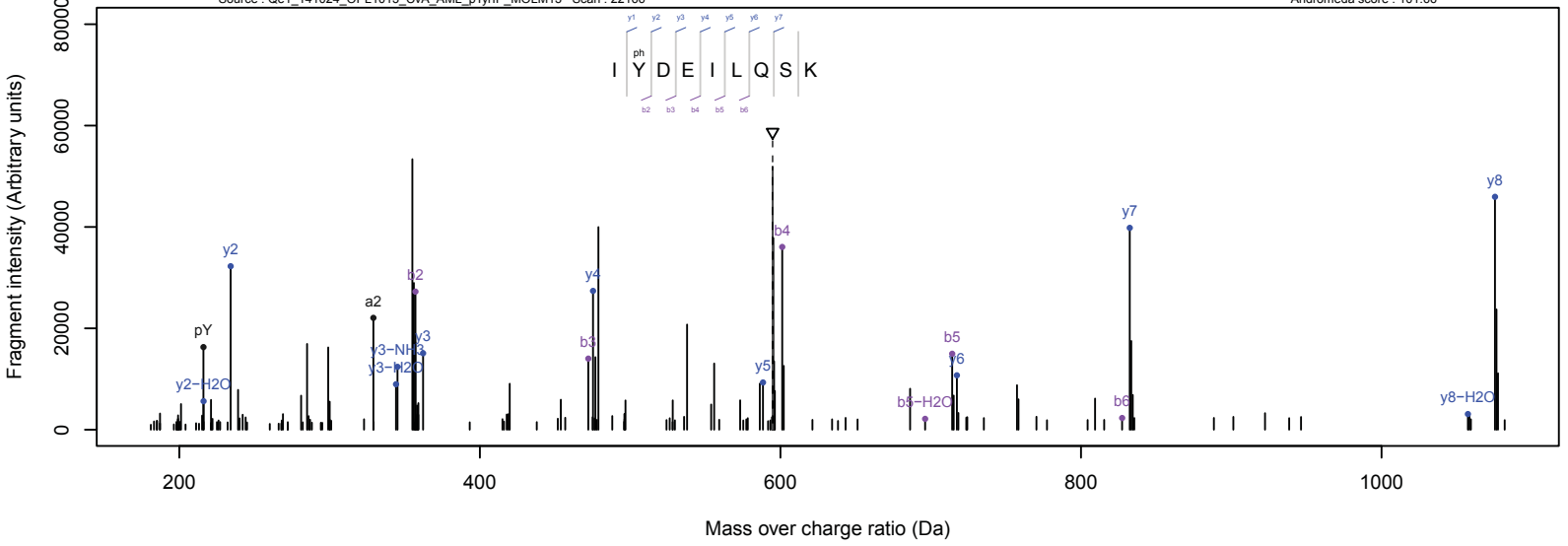
Andromeda score: 101.71



IY(ph)DEILQSK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 22166

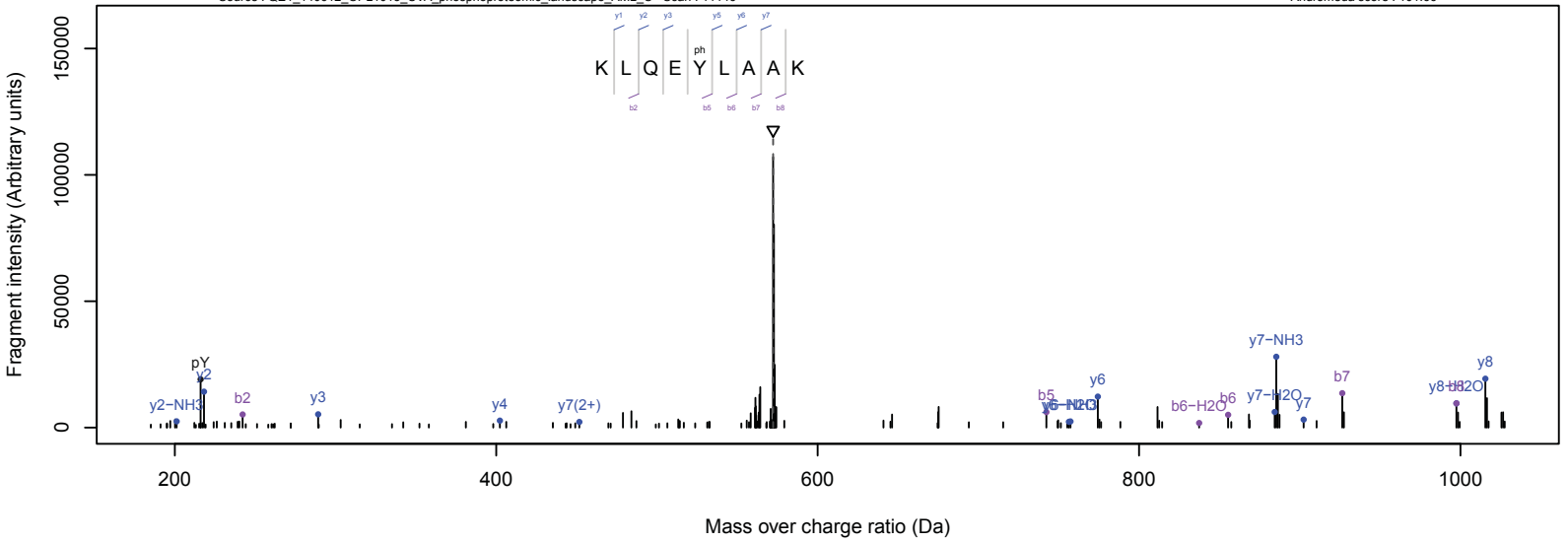
Andromeda score : 101.66



KLQEY(ph)LAAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11146

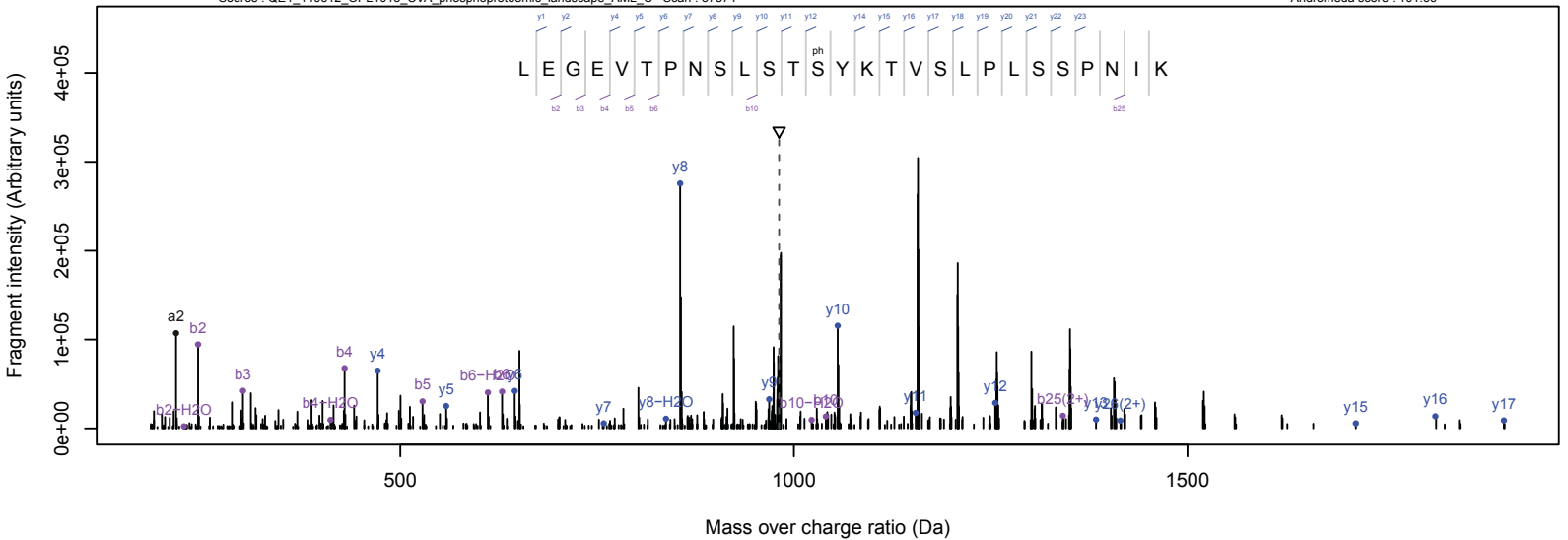
Andromeda score : 101.66



LEGEVTPNSLSTYSKTVSLPLSSPNIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37874

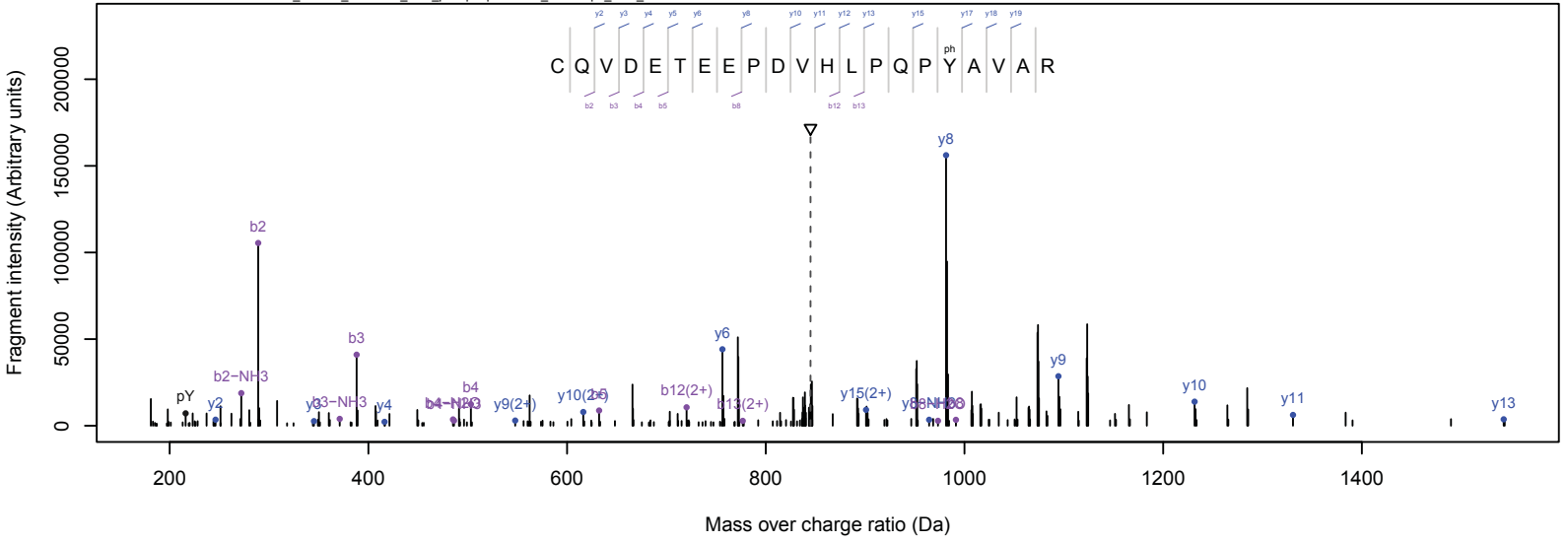
Andromeda score : 101.66



**CQVDETEEPDVHL P QPYAVAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29206

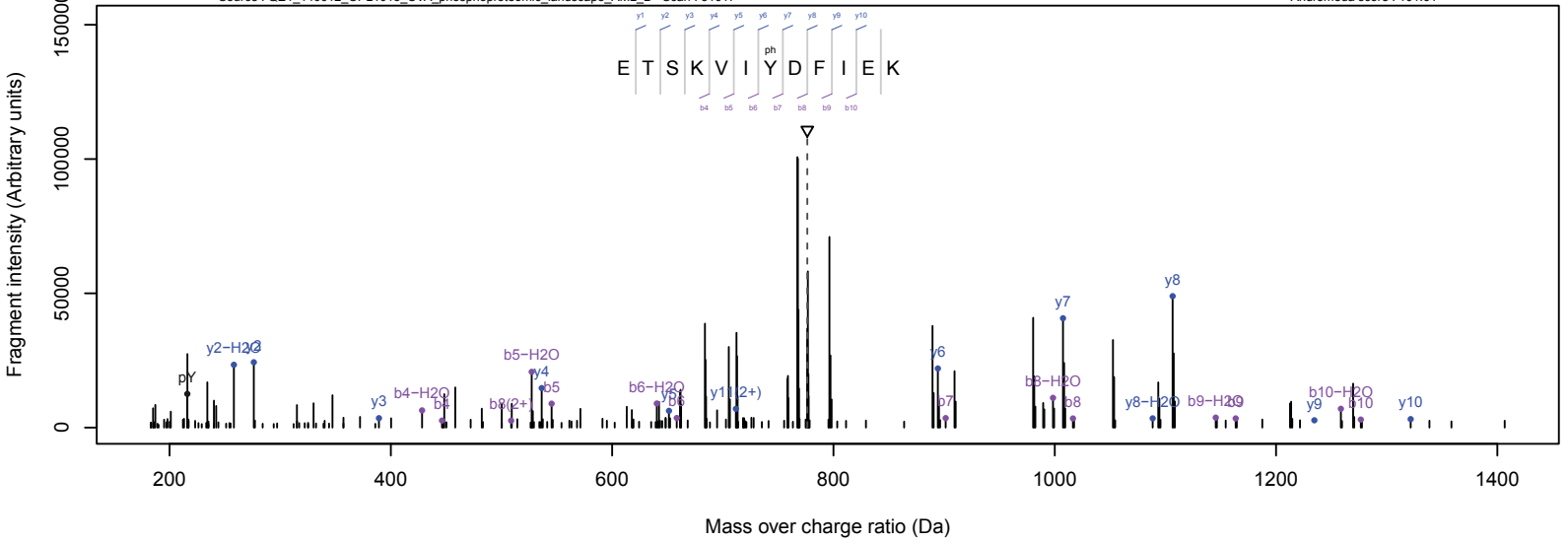
Andromeda score : 101.61



**ETSKVIY(ph)DFIEK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31617

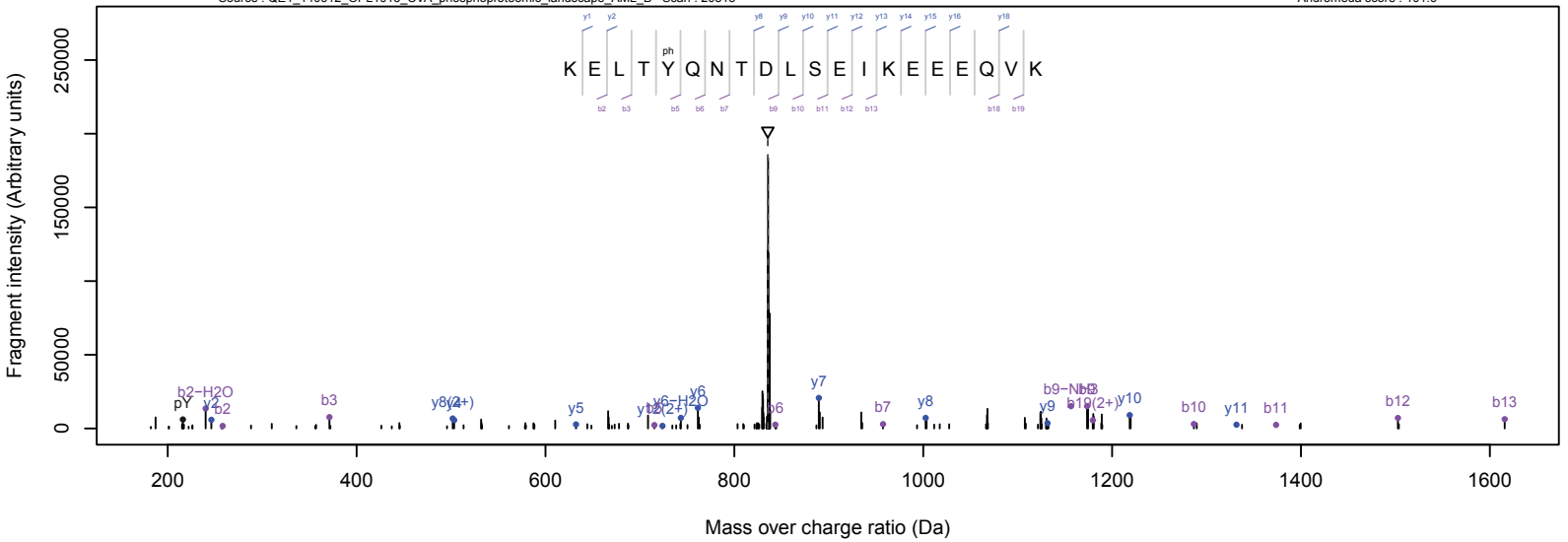
Andromeda score : 101.61



**KELTY(ph)QNTDLSEIK EEEQVK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20815

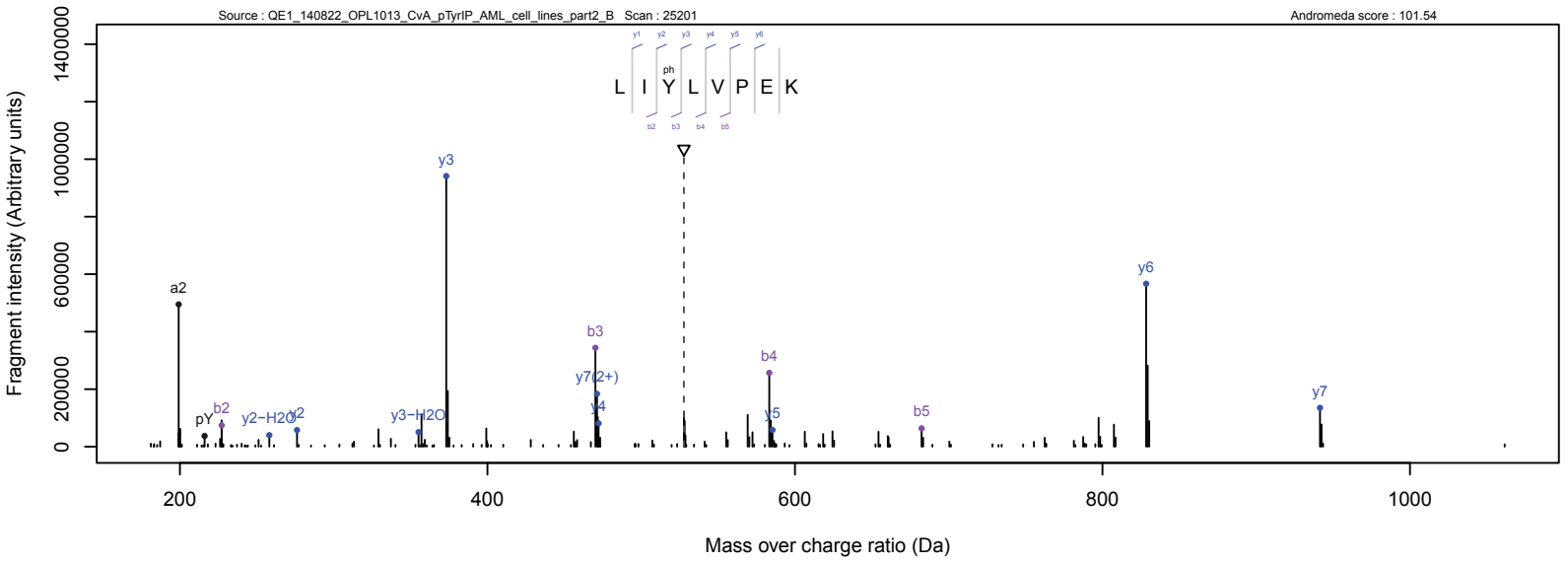
Andromeda score : 101.6



LIY(ph)LVPEK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 25201

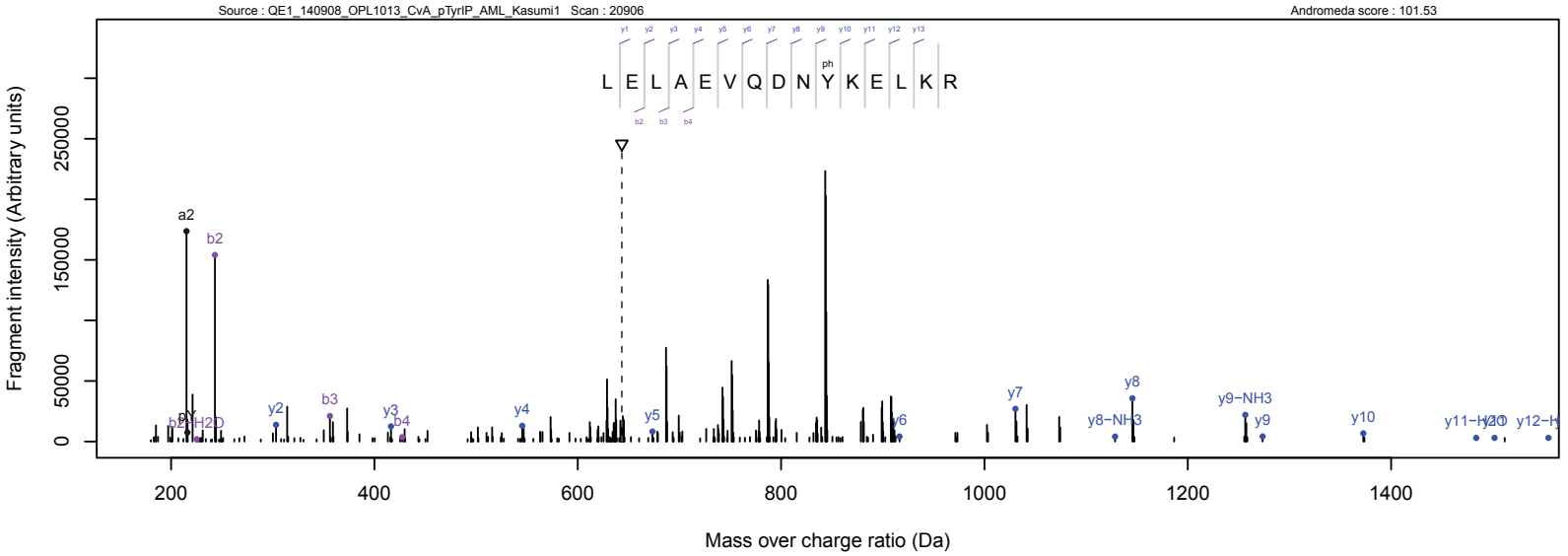
Andromeda score: 101.54



LELAEQDNY(ph)KELKR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 20906

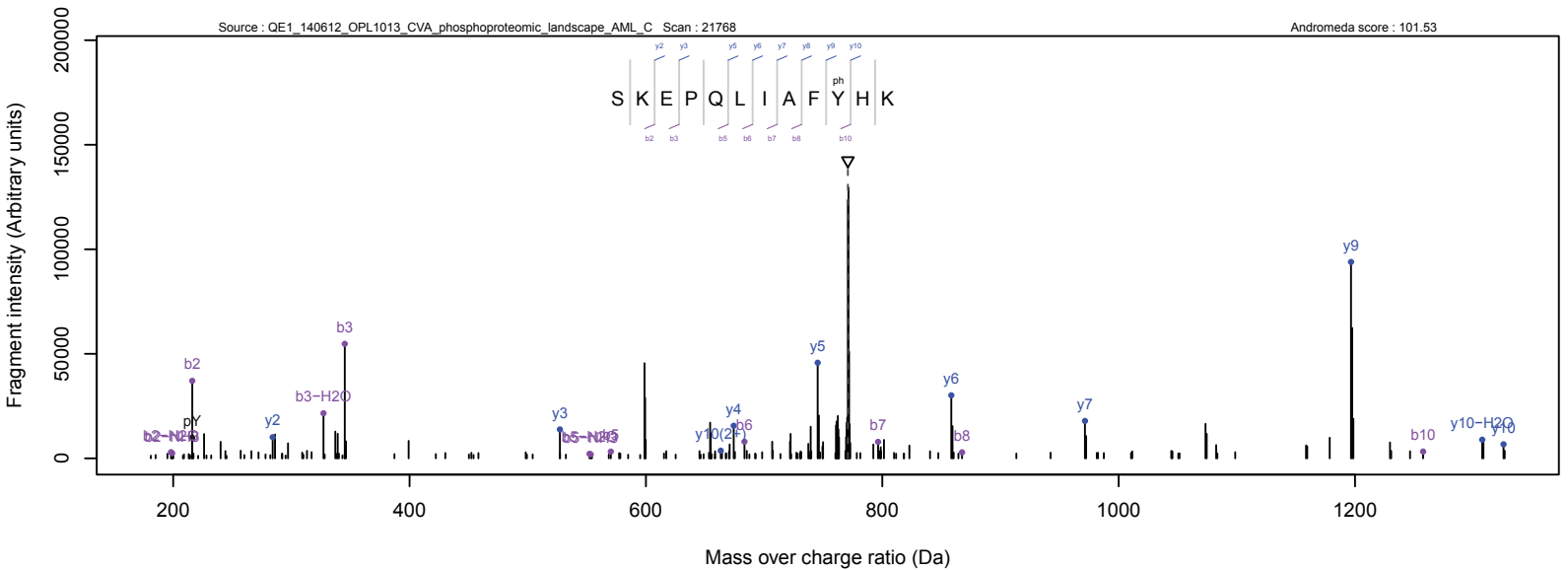
Andromeda score: 101.53



SKEPQLIAFY(ph)HK-2+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 21768

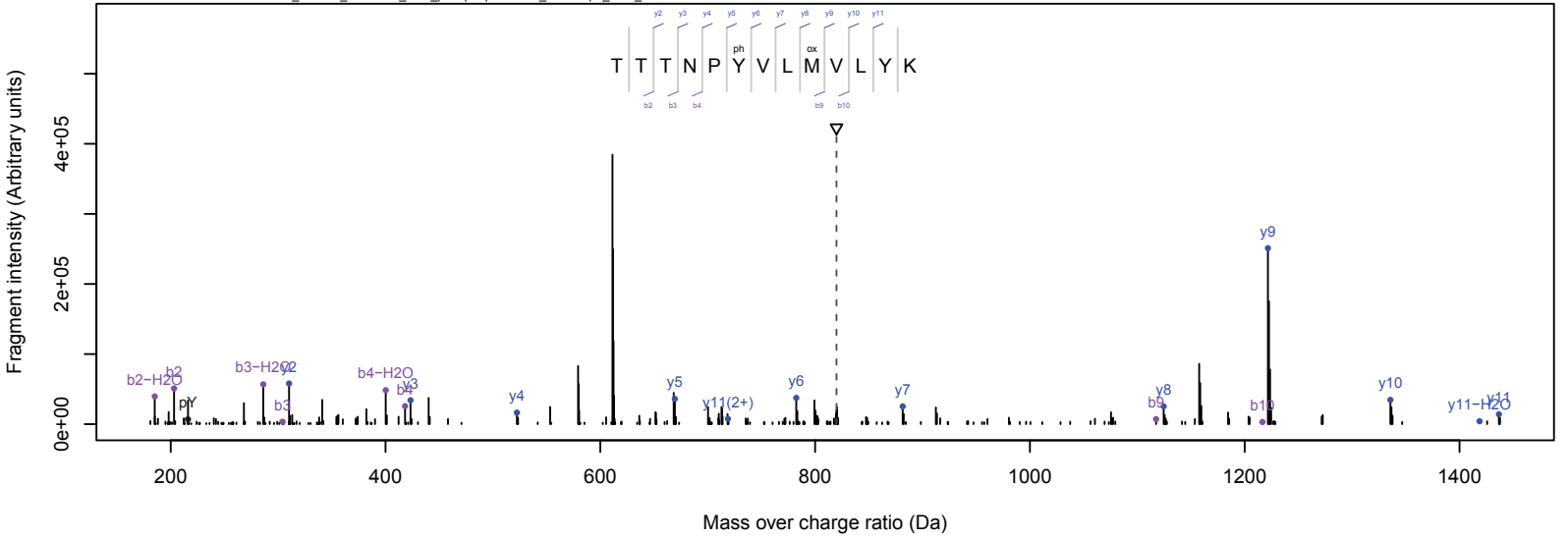
Andromeda score: 101.53



TTTNPY(ph)VLM(ox)VLYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 38015

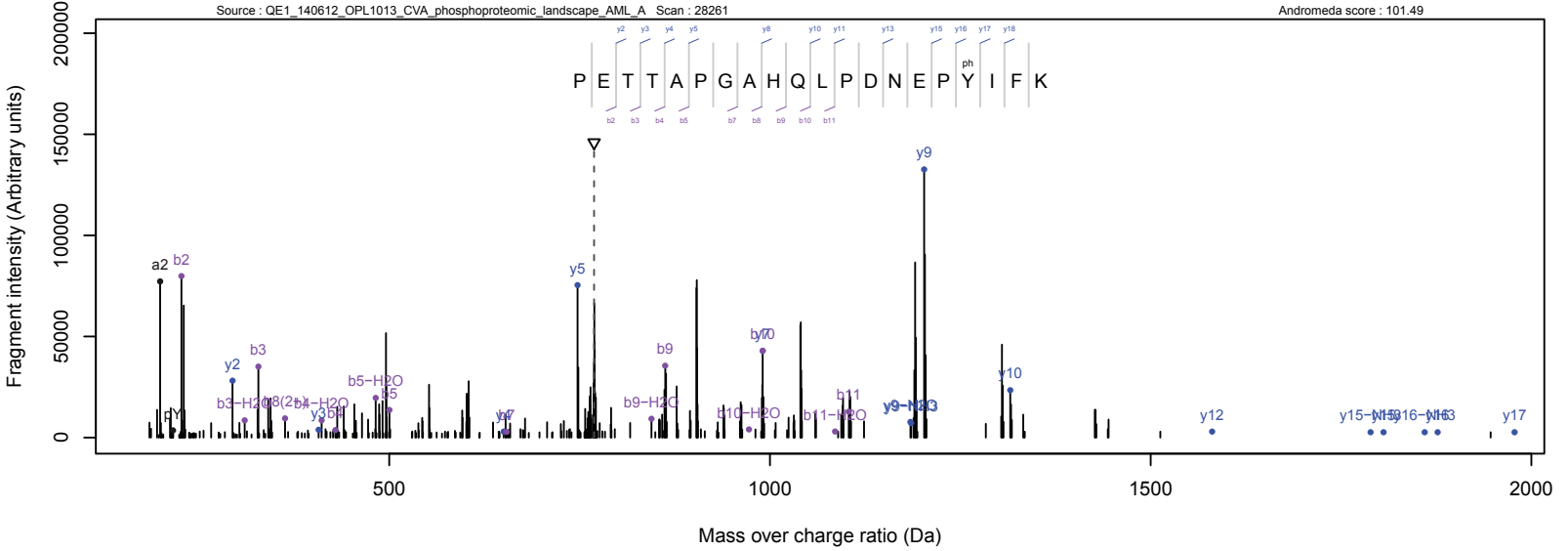
Andromeda score : 101.53



PETTAPGAHQLPDNEP(ph)YIFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 28261

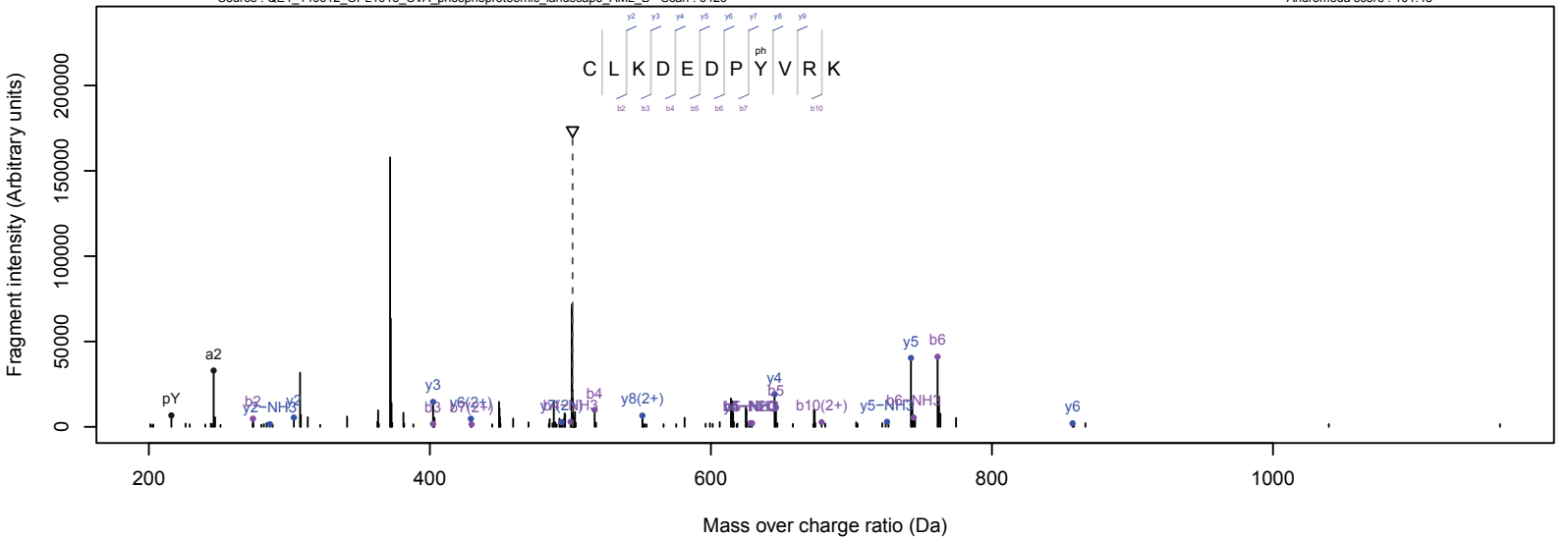
Andromeda score : 101.49



CLKDEDPYVRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 6425

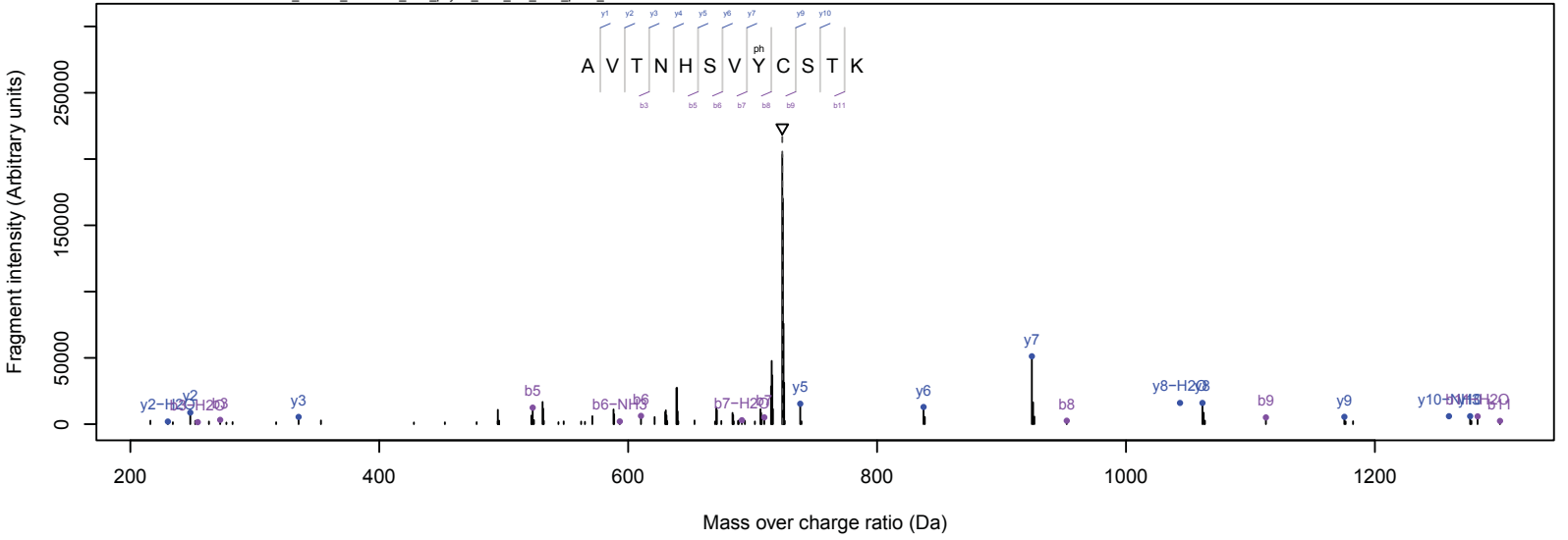
Andromeda score : 101.48



AVTNHSVY(ph)CSTK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 2960

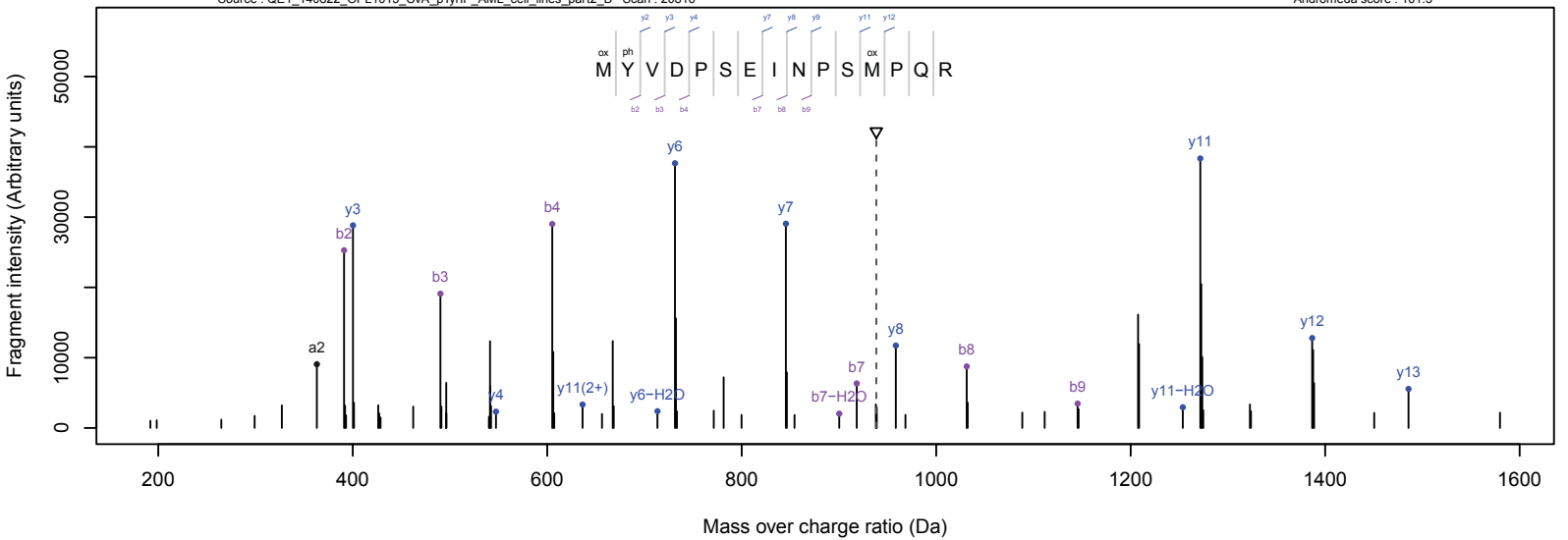
Andromeda score : 101.3



M(ox)Y(ph)VDPSEINP(ox)SMPQR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 20816

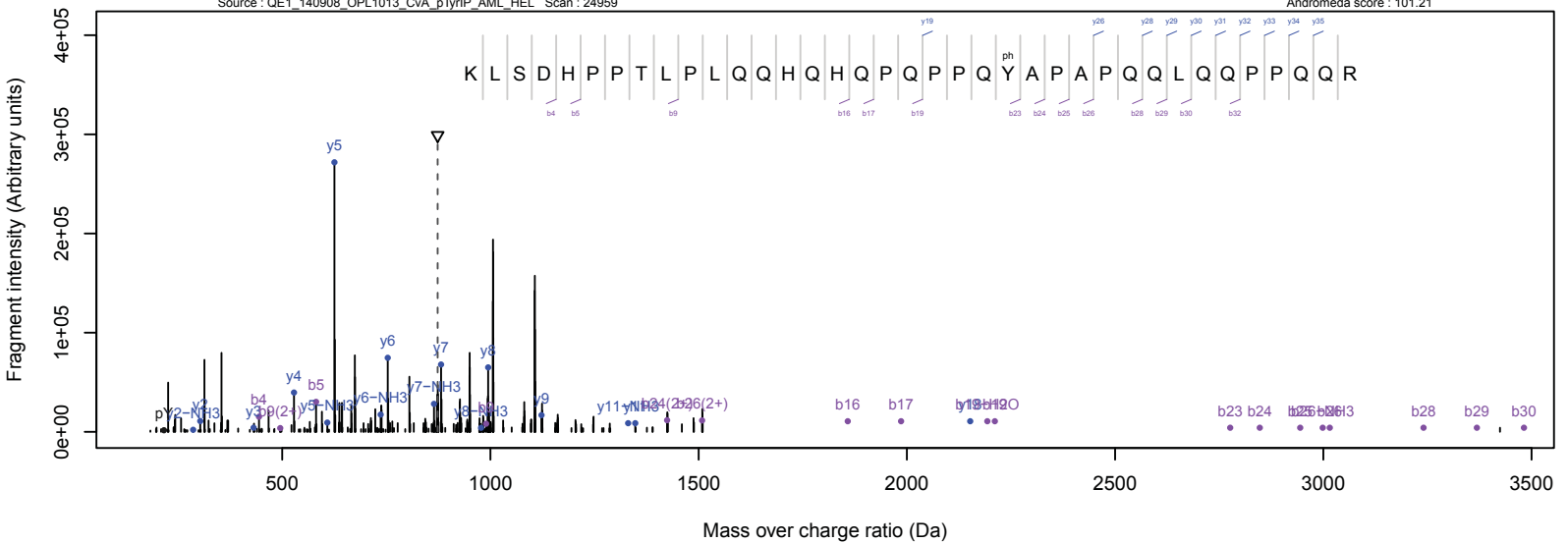
Andromeda score : 101.3

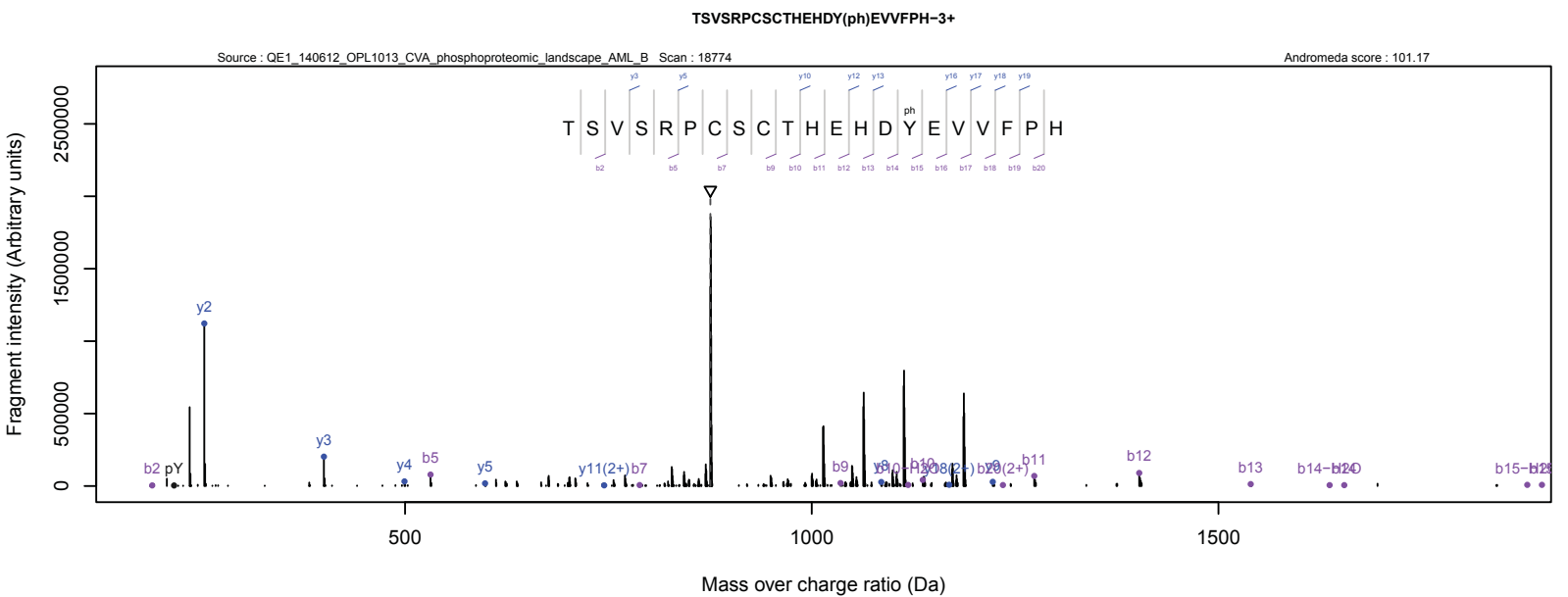
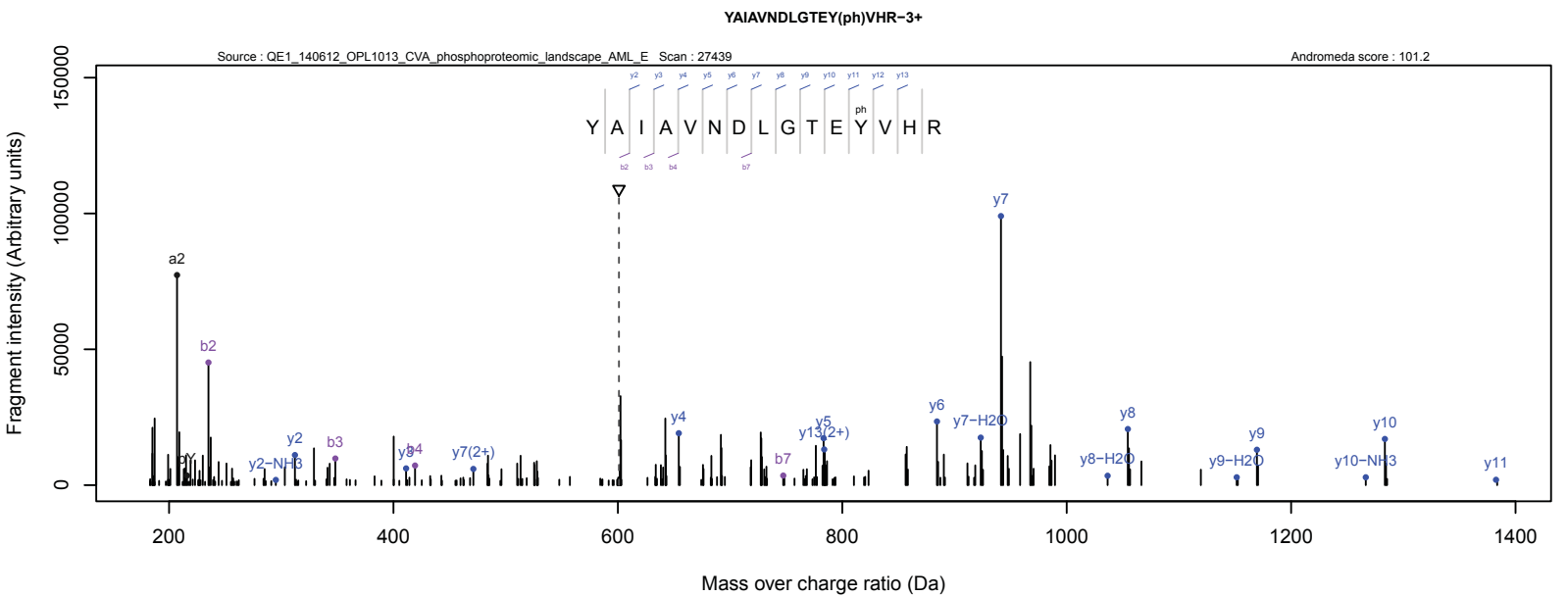
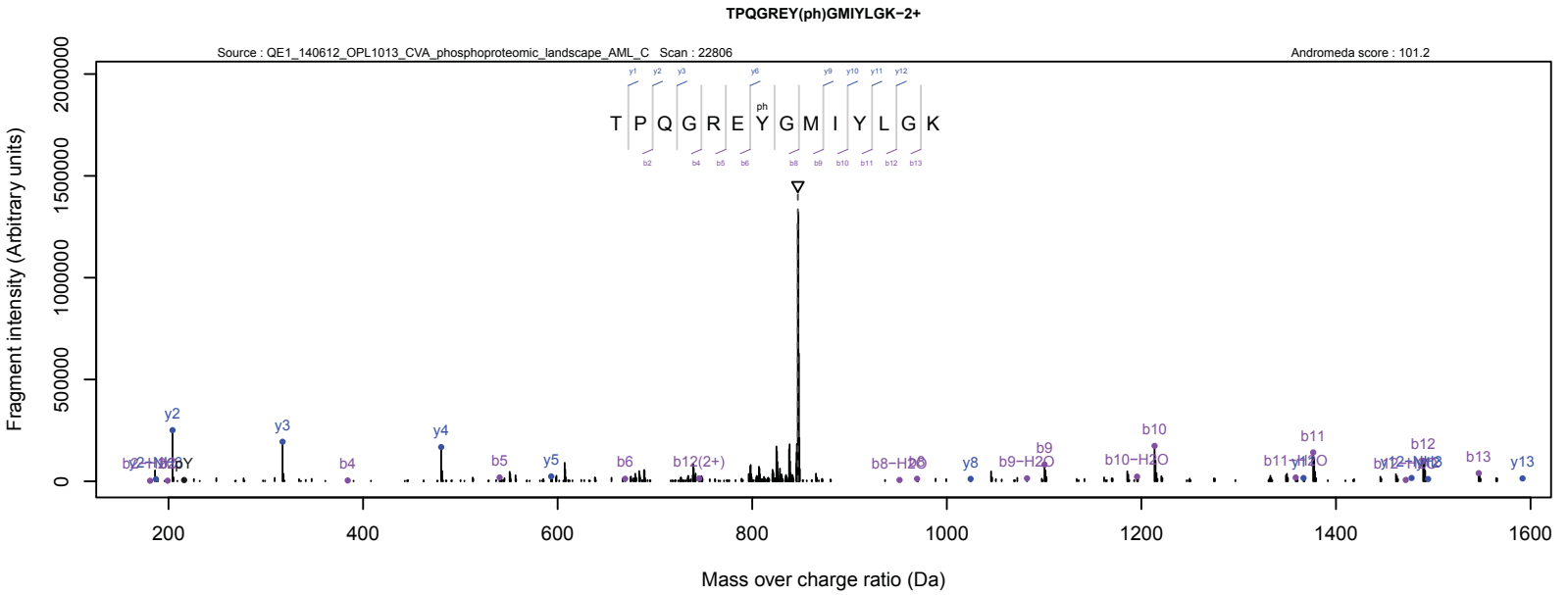


KLSDHPPTLPLQHQHQPPQY(ph)APAPQLQPPQQR-5+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24959

Andromeda score : 101.21

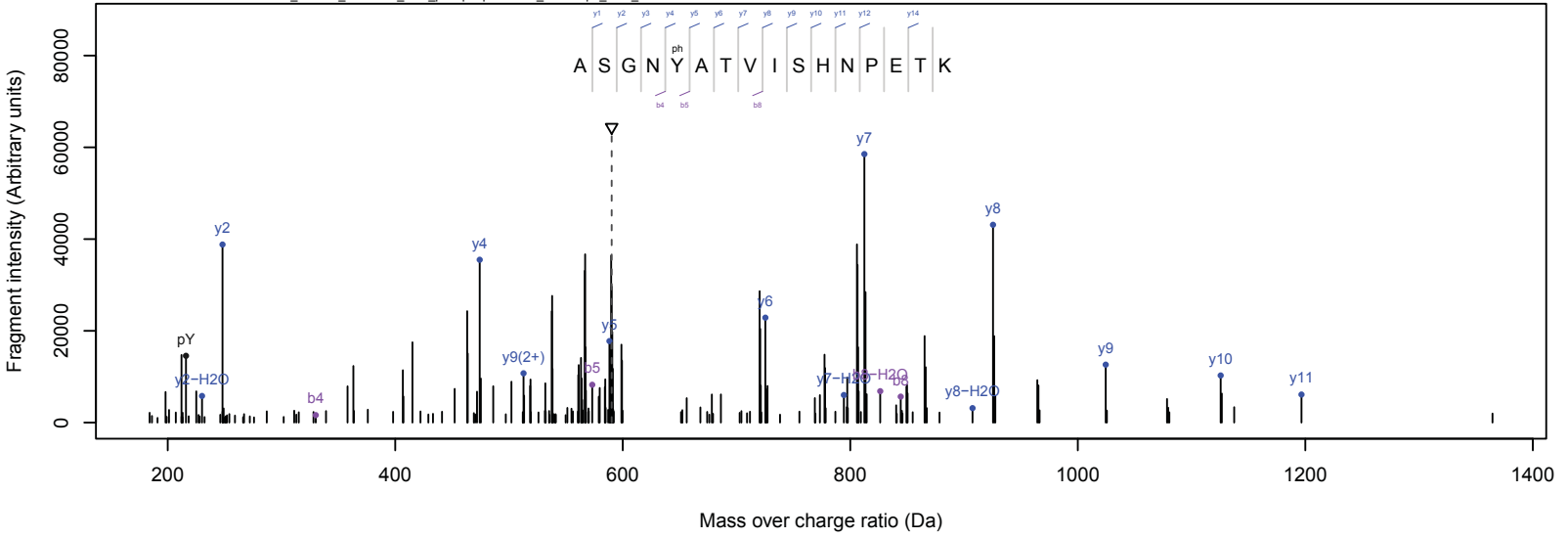




**ASGNY(ph)ATVISHNPETK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11219

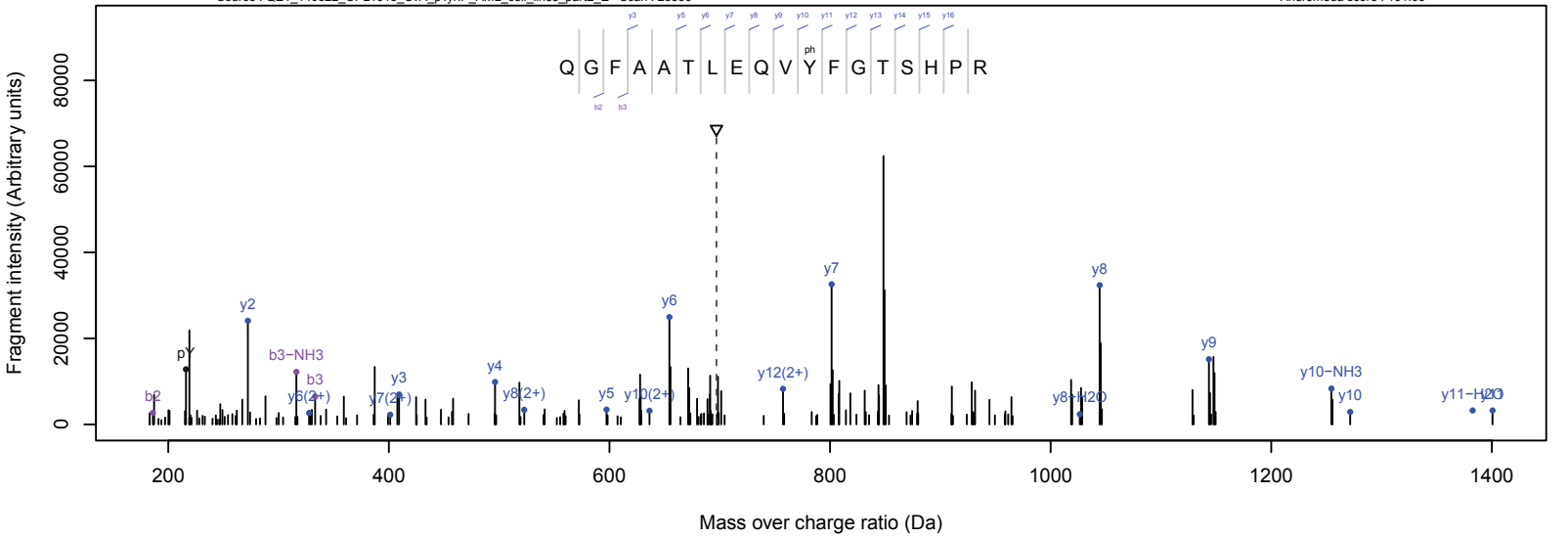
Andromeda score : 101.05



**QGFAATLEQVY(ph)FGTSHPR-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 28556

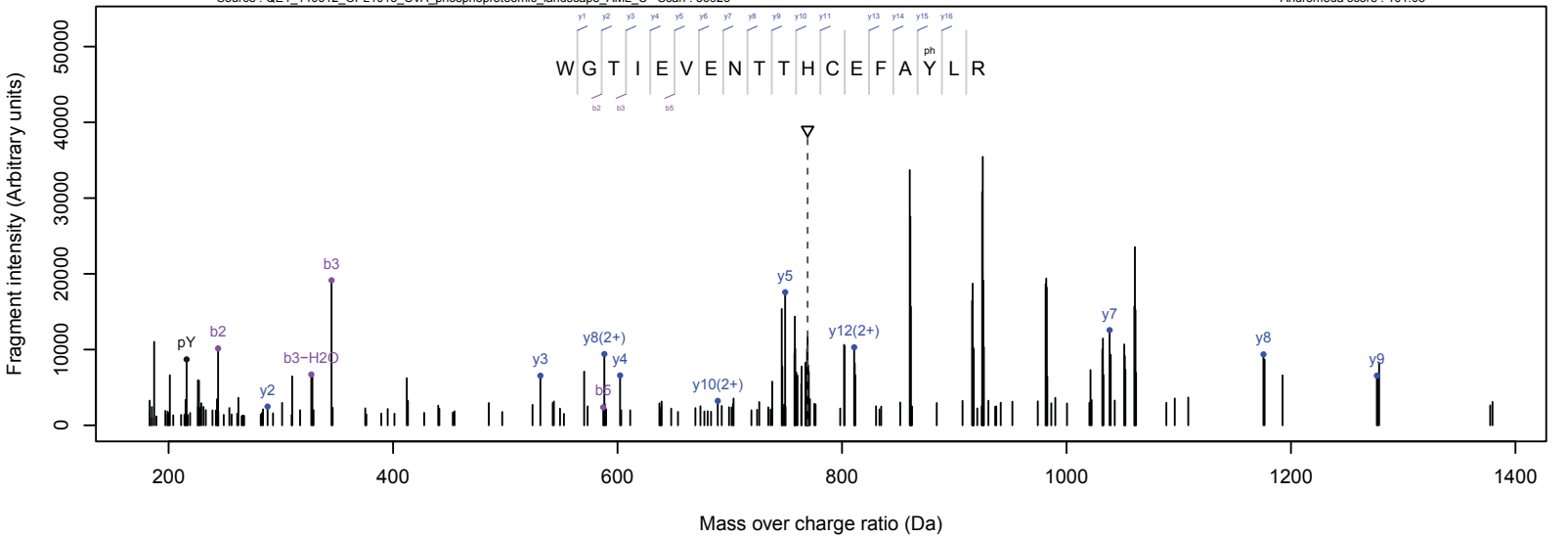
Andromeda score : 101.05



**WG TIEVENTTHCEFA Y L R-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36926

Andromeda score : 101.05

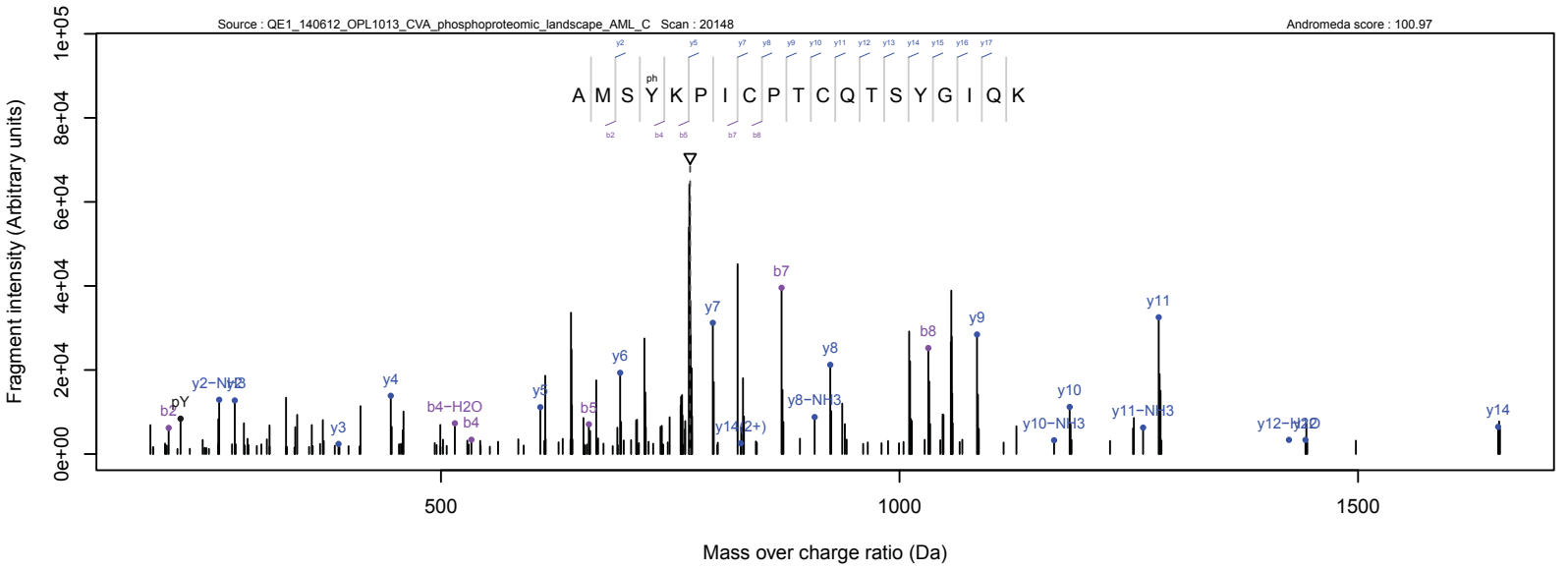




**AMSY(ph)KPICPTCQTSYGIQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20148

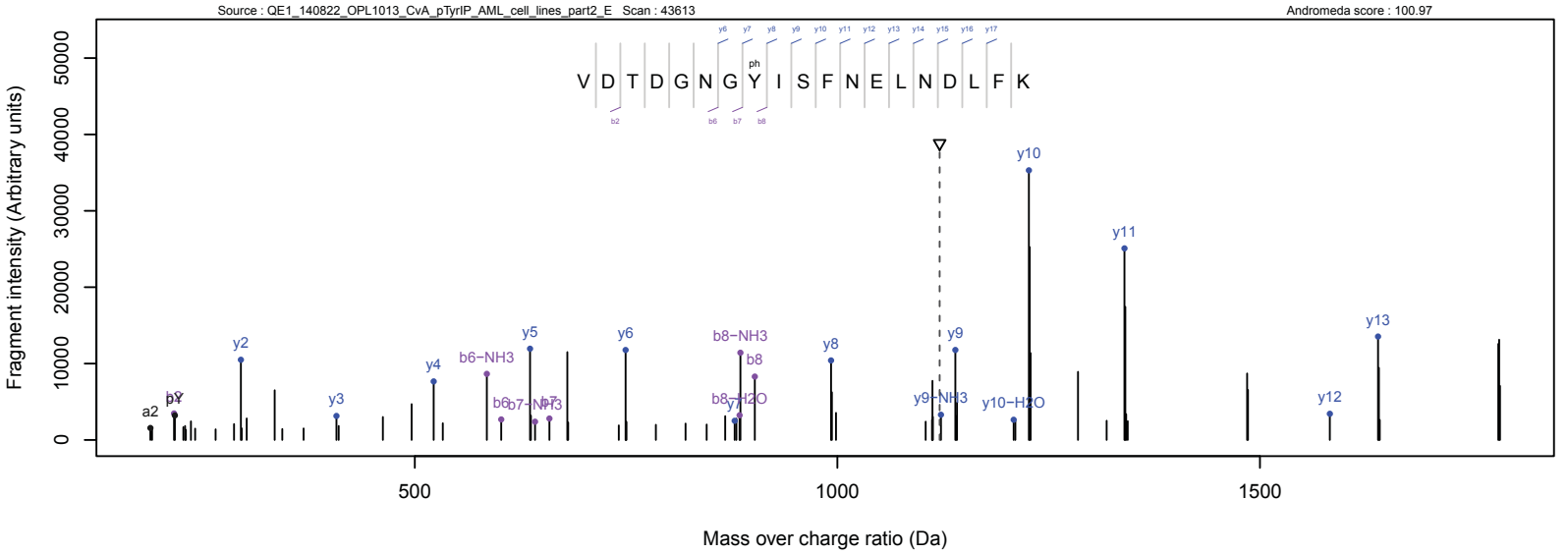
Andromeda score : 100.97



**VDTDNGY(ph)ISFNELDLFK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 43613

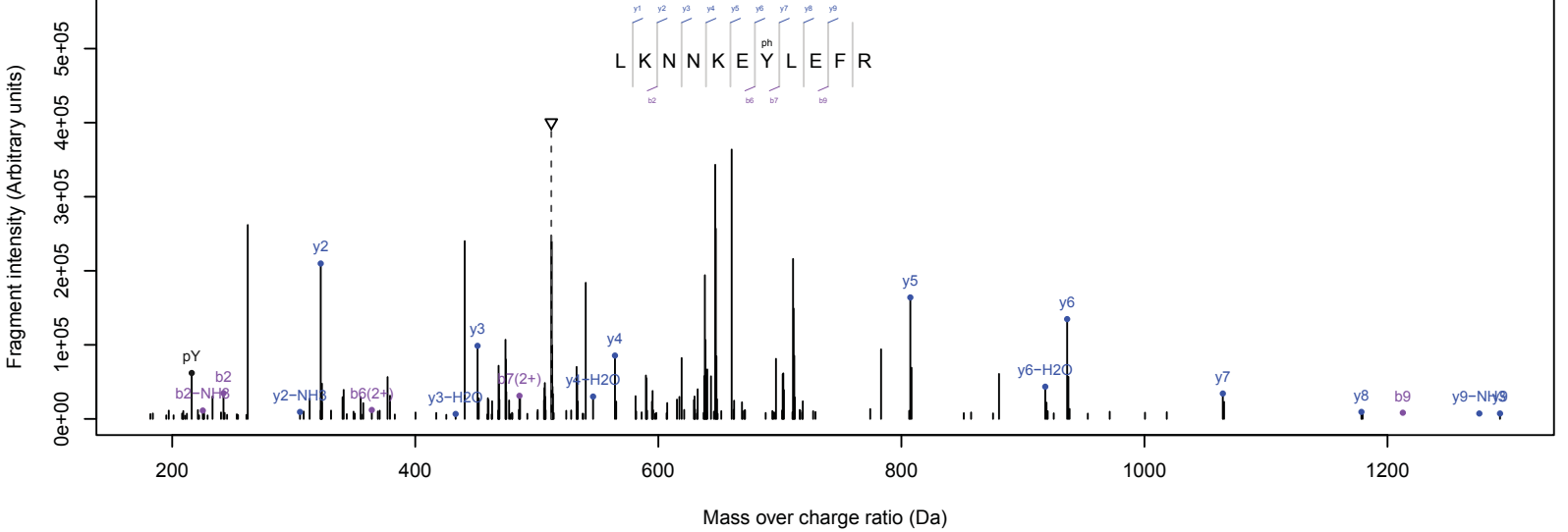
Andromeda score : 100.97



**LKNNKEY(ph)LEFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13136

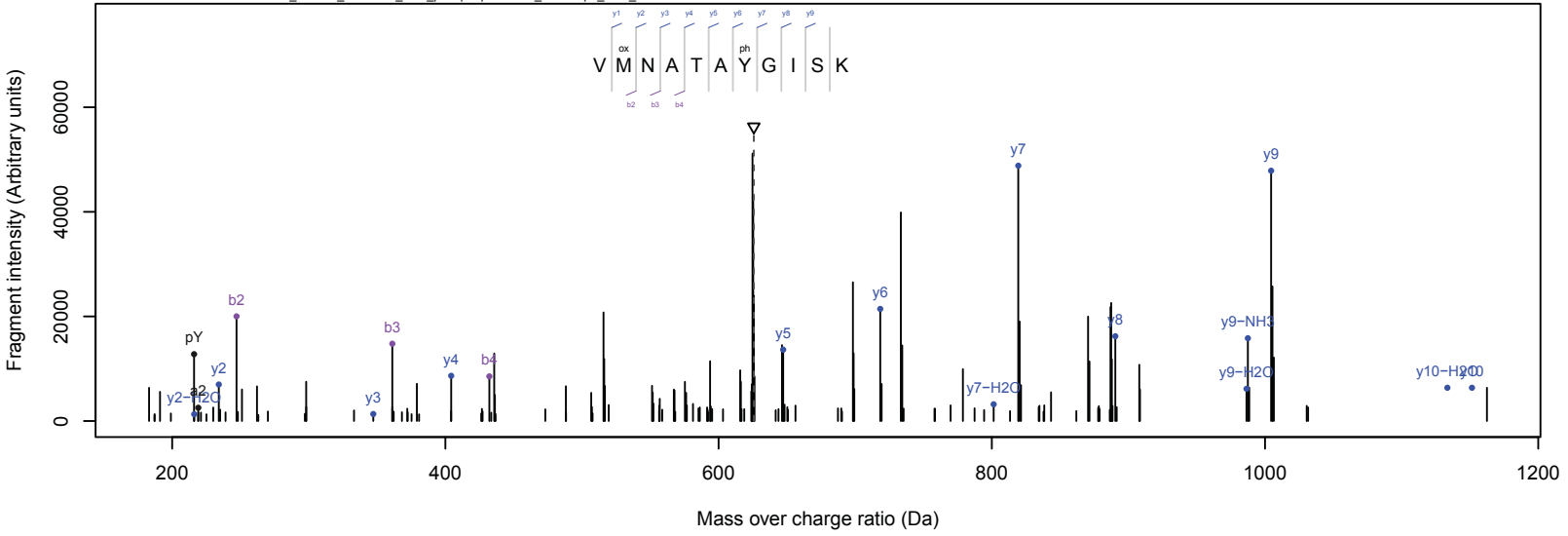
Andromeda score : 100.93



VM(ox)NATAY(ph)GISK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11531

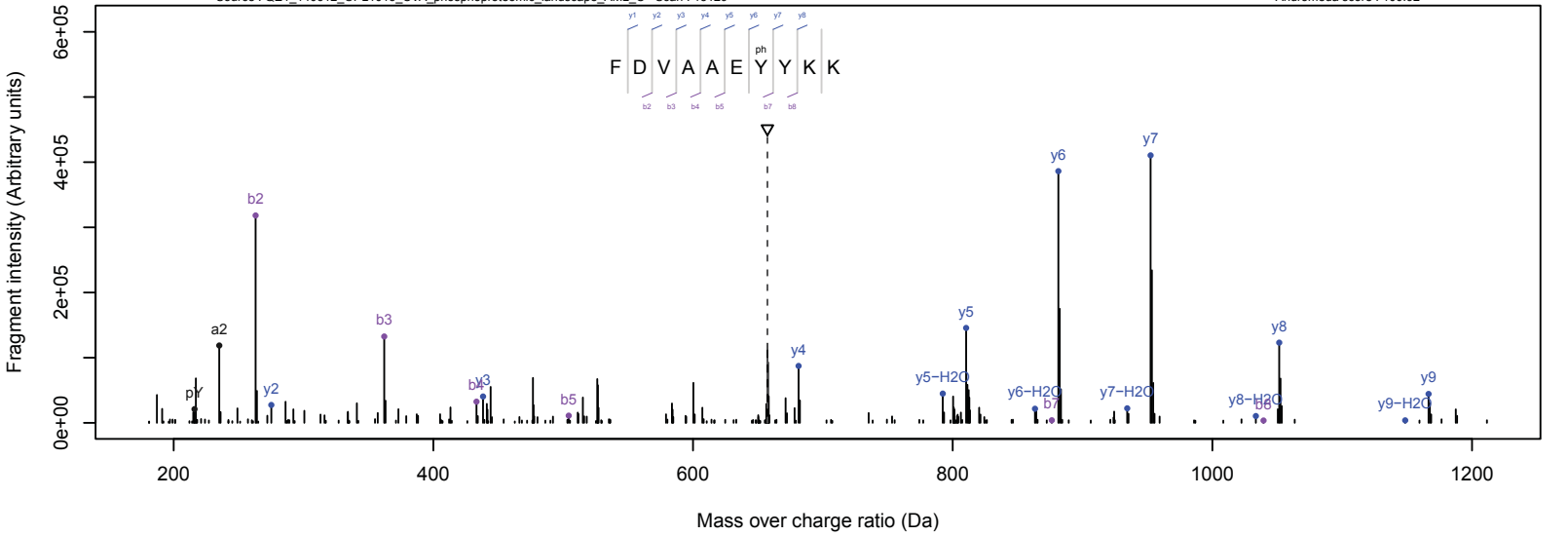
Andromeda score : 100.93



FDVAAEY(ph)YKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15129

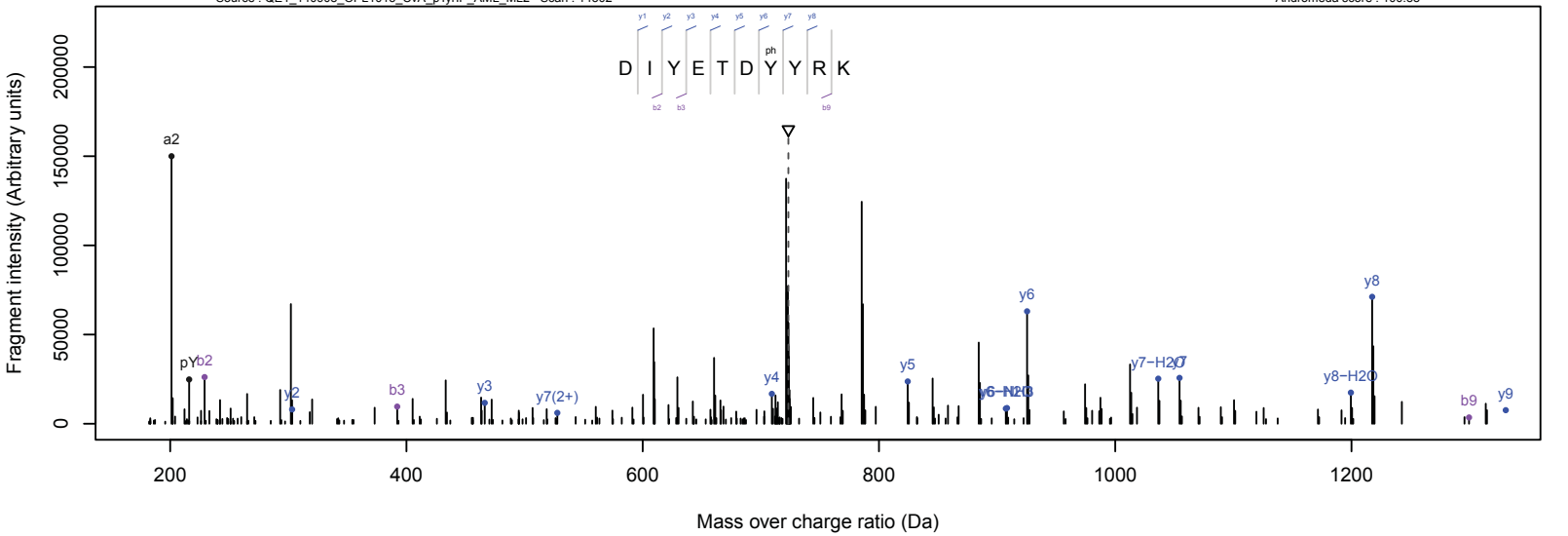
Andromeda score : 100.92



DIYETDY(ph)YRK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 14502

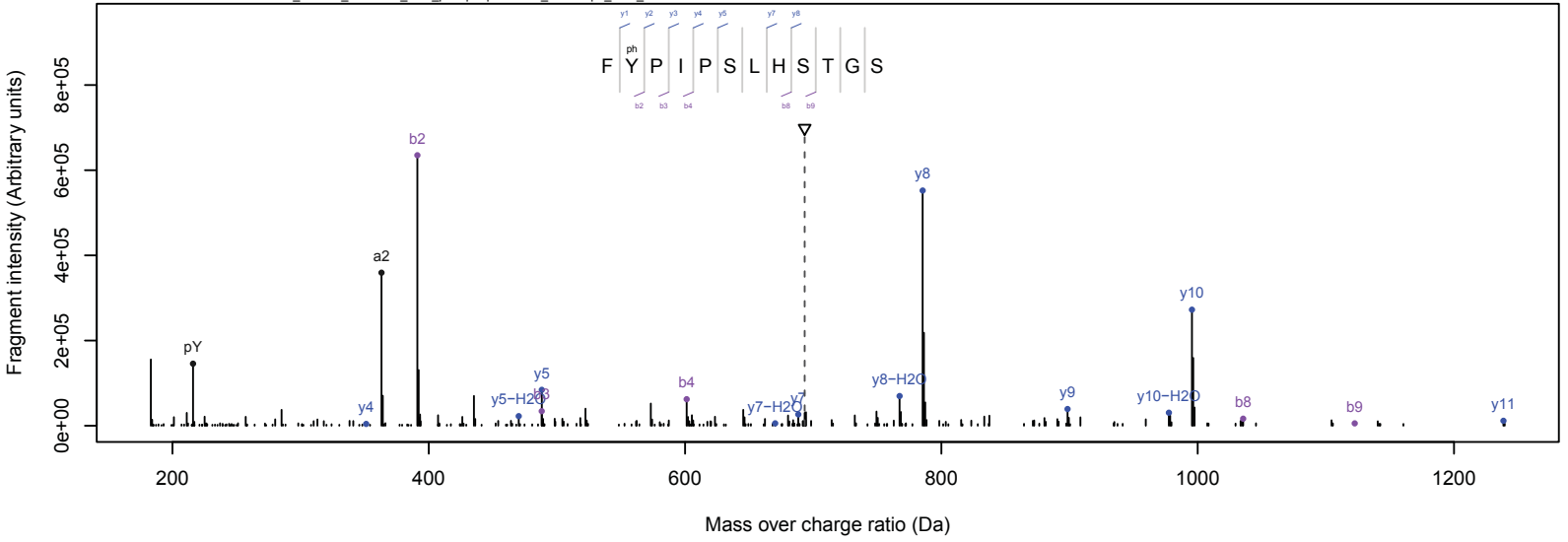
Andromeda score : 100.88



**FY(ph)PIPSLHSTGS-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31749

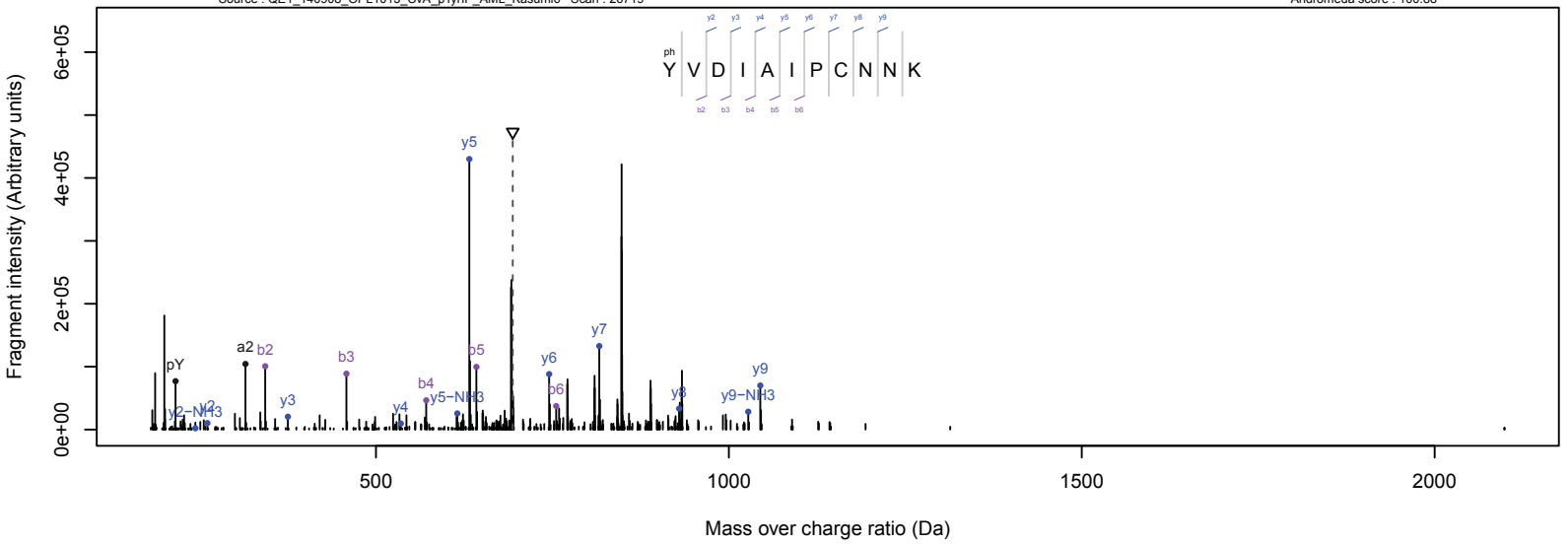
Andromeda score: 100.88



**Y(ph)VDIAIPCNNK-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 26719

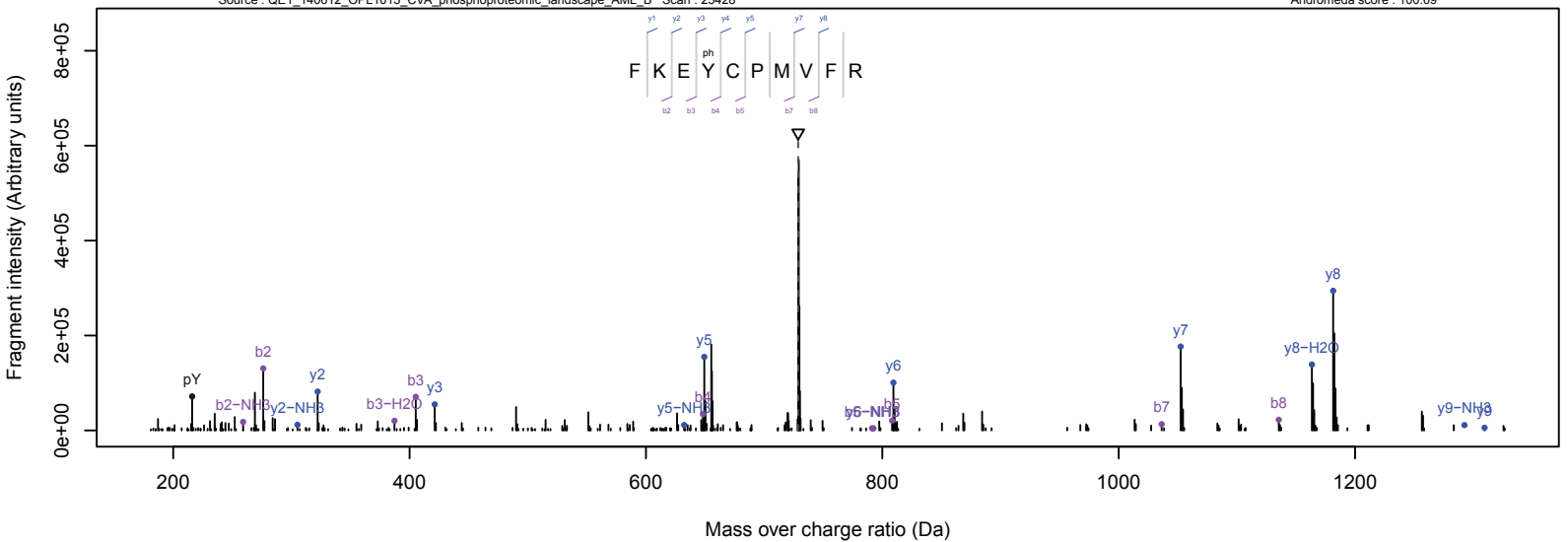
Andromeda score: 100.88



**FKEY(ph)CPMVFR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 25428

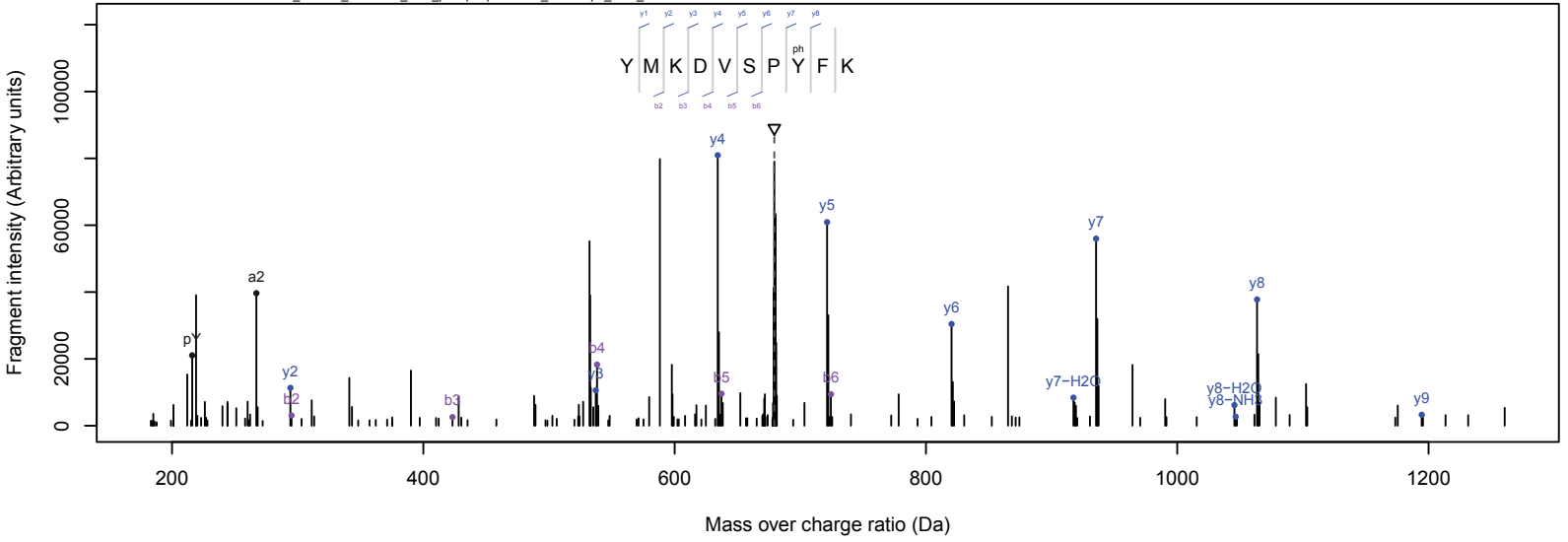
Andromeda score: 100.69



**YMKDVSPY(ph)FK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18166

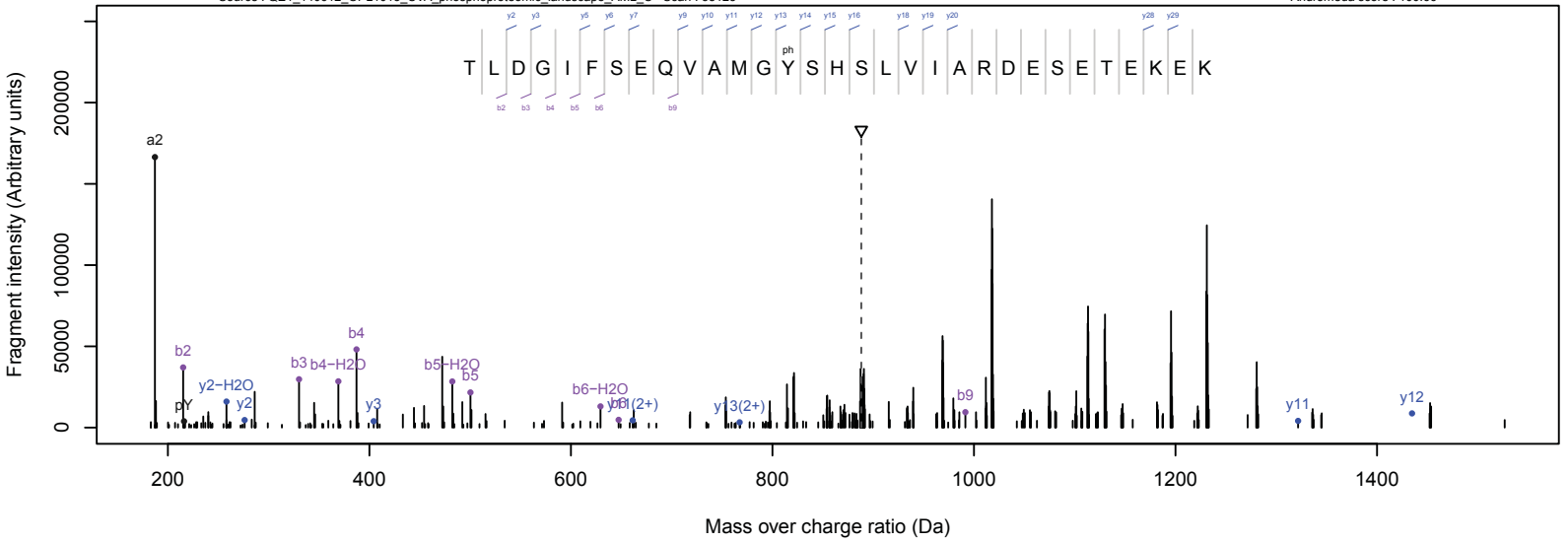
Andromeda score : 100.69



**TLDGIFSEQVAMGY(ph)SHSLVIARDESETEKEK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38125

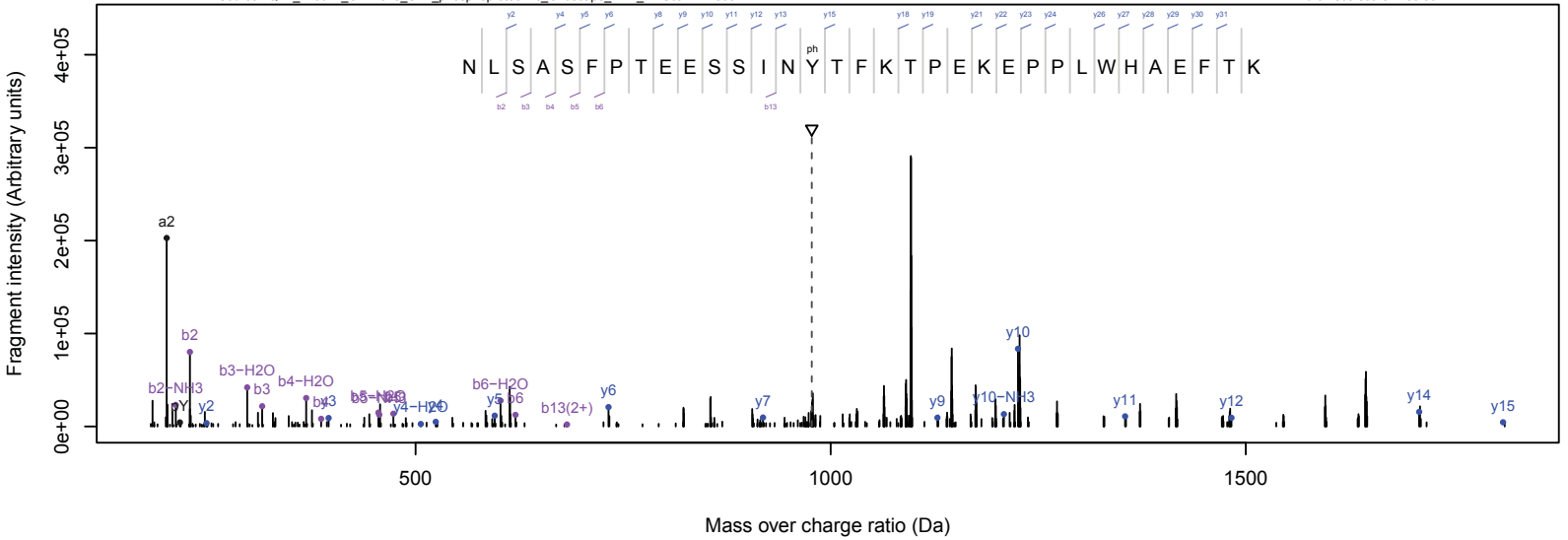
Andromeda score : 100.66



**NLSASFTEESSIN(ph)YTFKTPKEKPLWHAFTK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41065

Andromeda score : 100.59

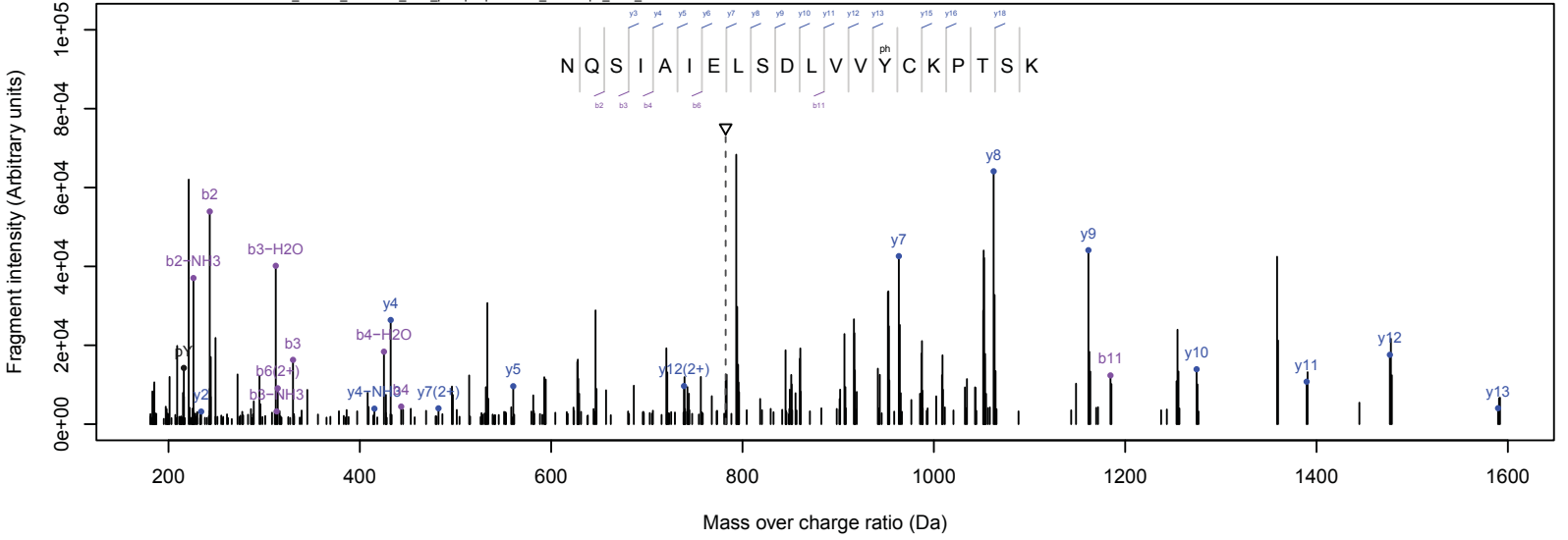




**NQSIAIELSDLVY(ph)CKPTSK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 44414

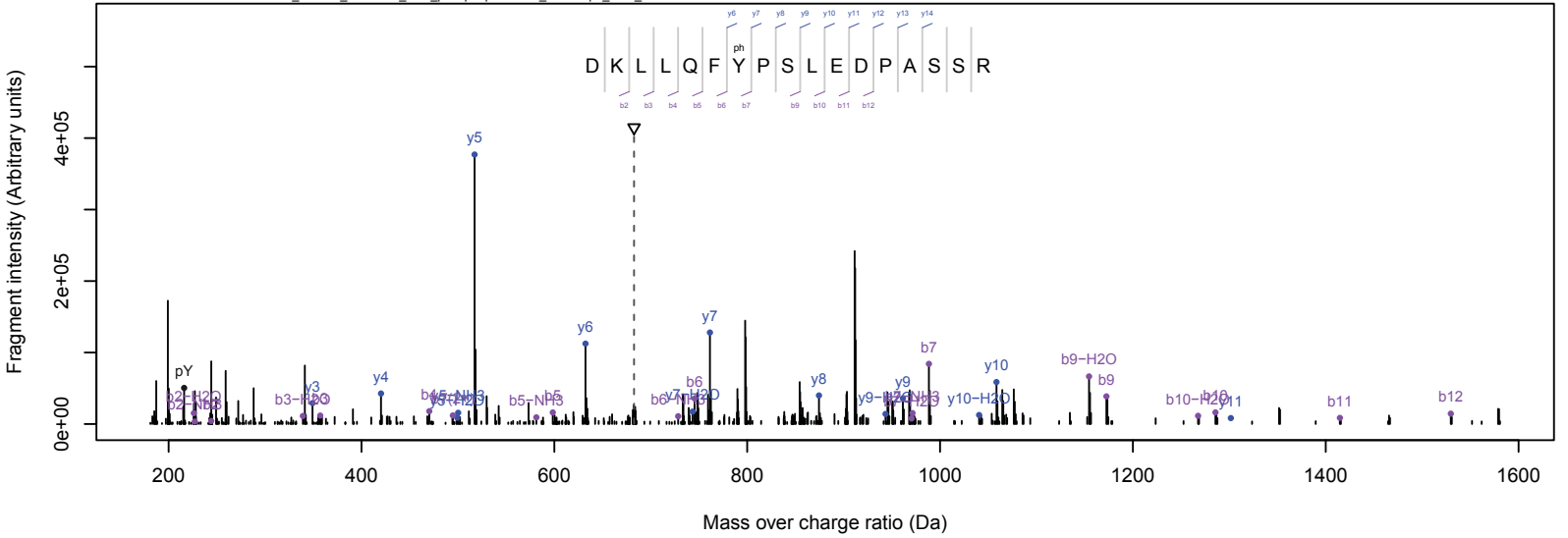
Andromeda score: 100.28



**DKLLQFY(ph)PSLEDPASSR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38706

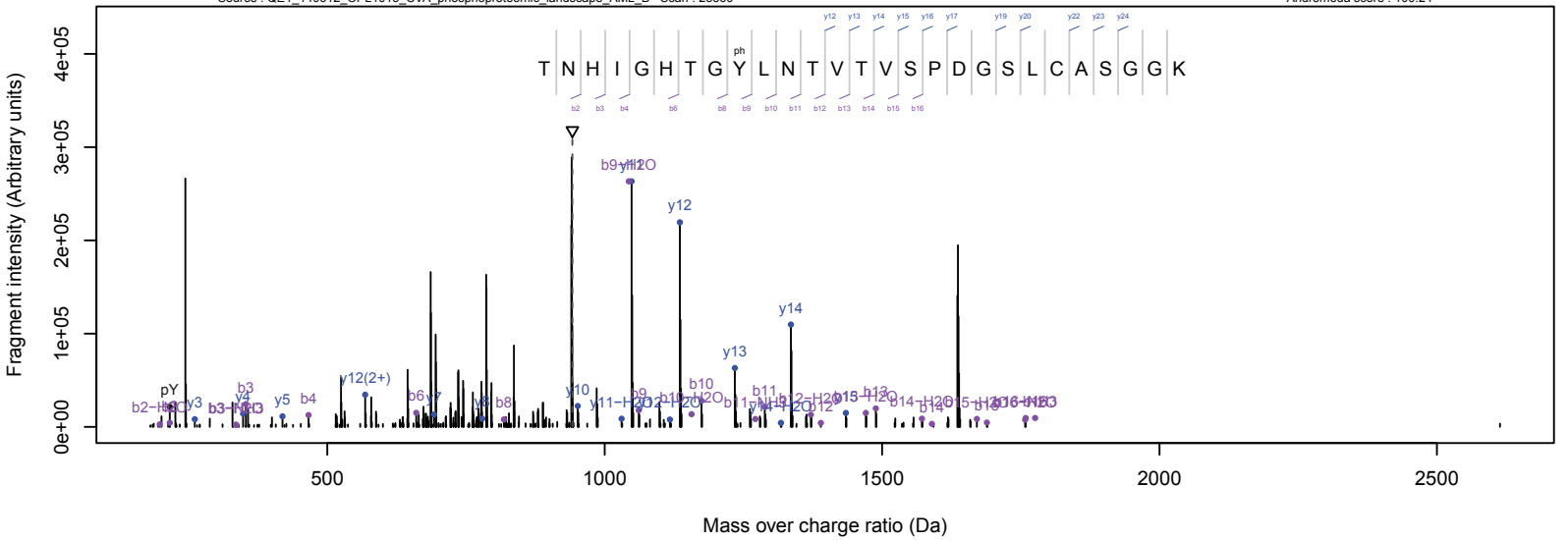
Andromeda score: 100.27



**TNHIGHTGY(ph)LNTVTVPDGLCASGGK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 23860

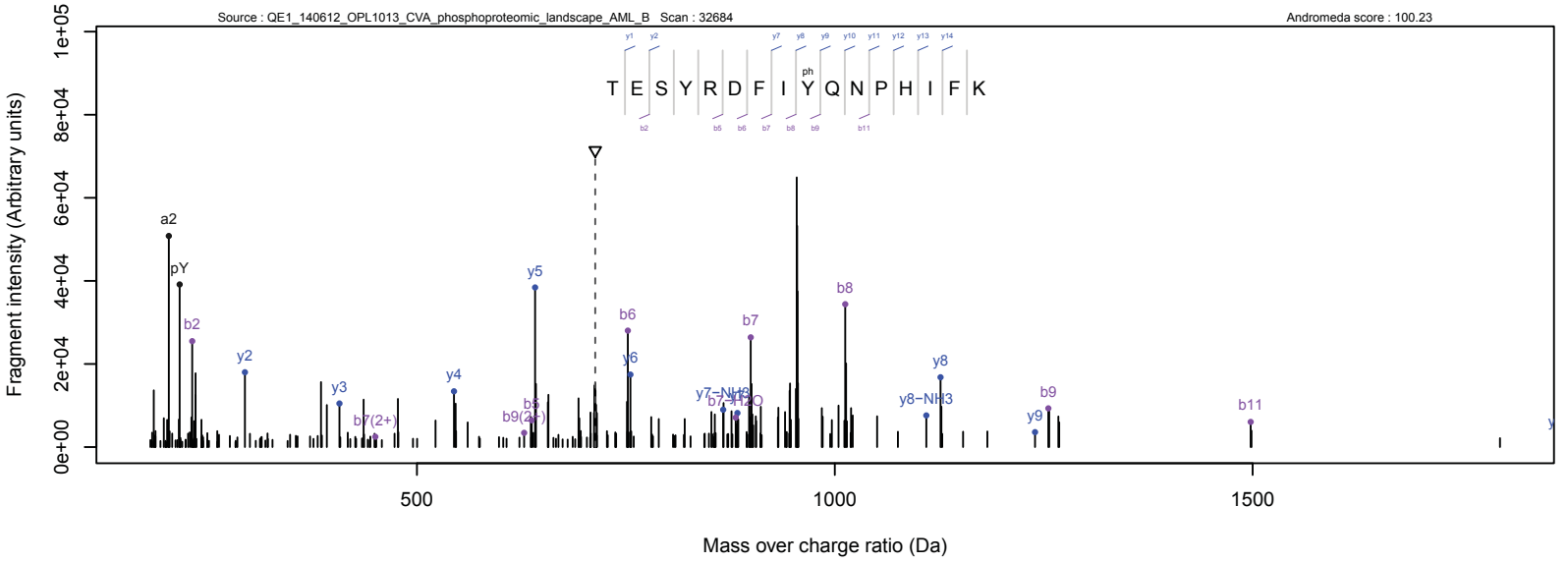
Andromeda score: 100.24



TESYRDFIYQNP<sup>ph</sup>HI<sup>ph</sup>FK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32684

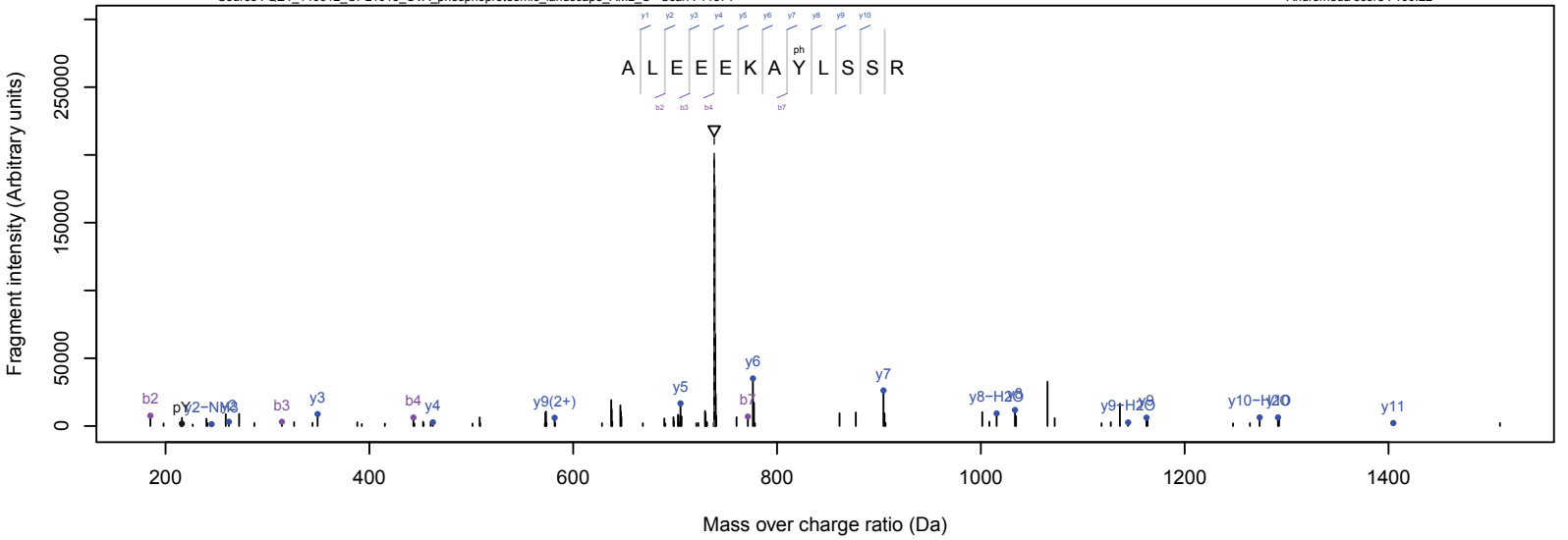
Andromeda score : 100.23



ALEEEKAYLSSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11371

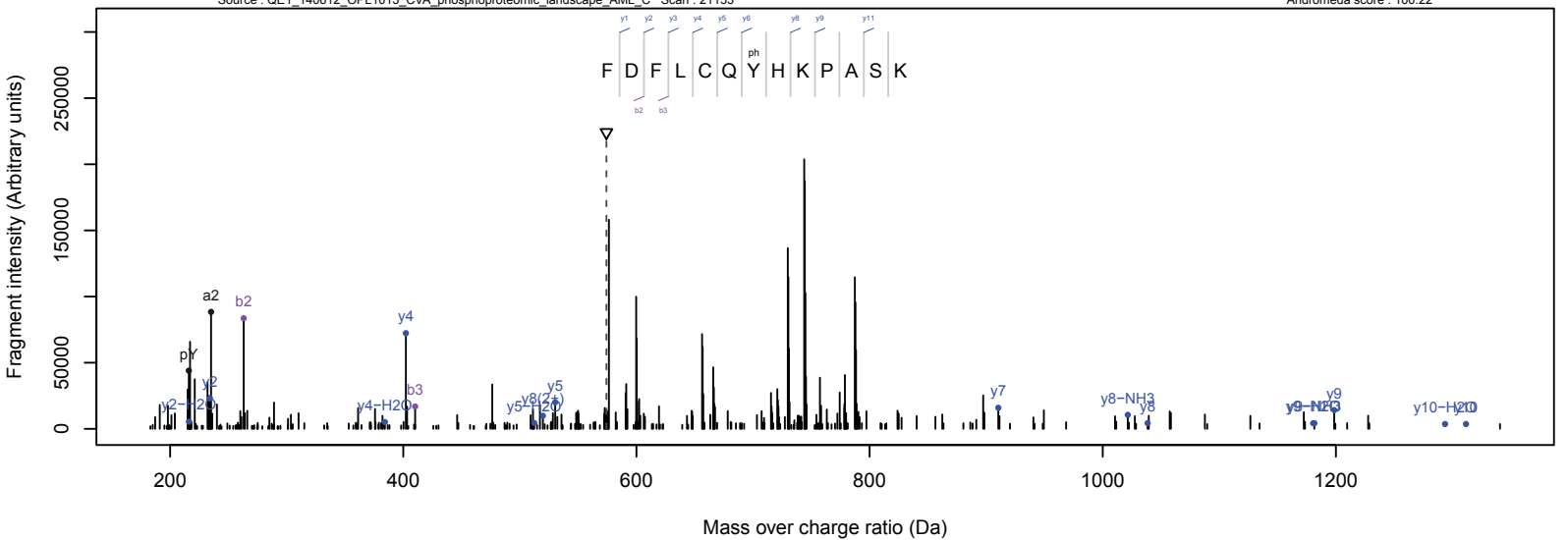
Andromeda score : 100.22



FD<sup>ph</sup>FLCQYHKPASK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21153

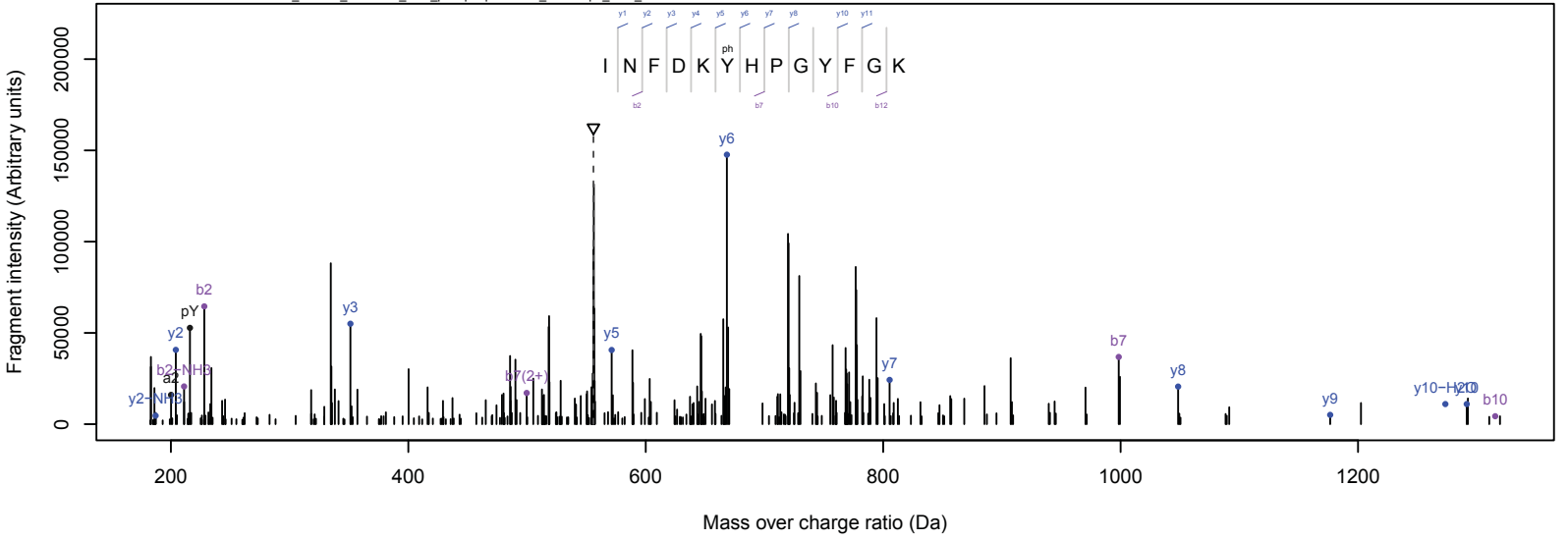
Andromeda score : 100.22



INFDKY(ph)HPGYFGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22690

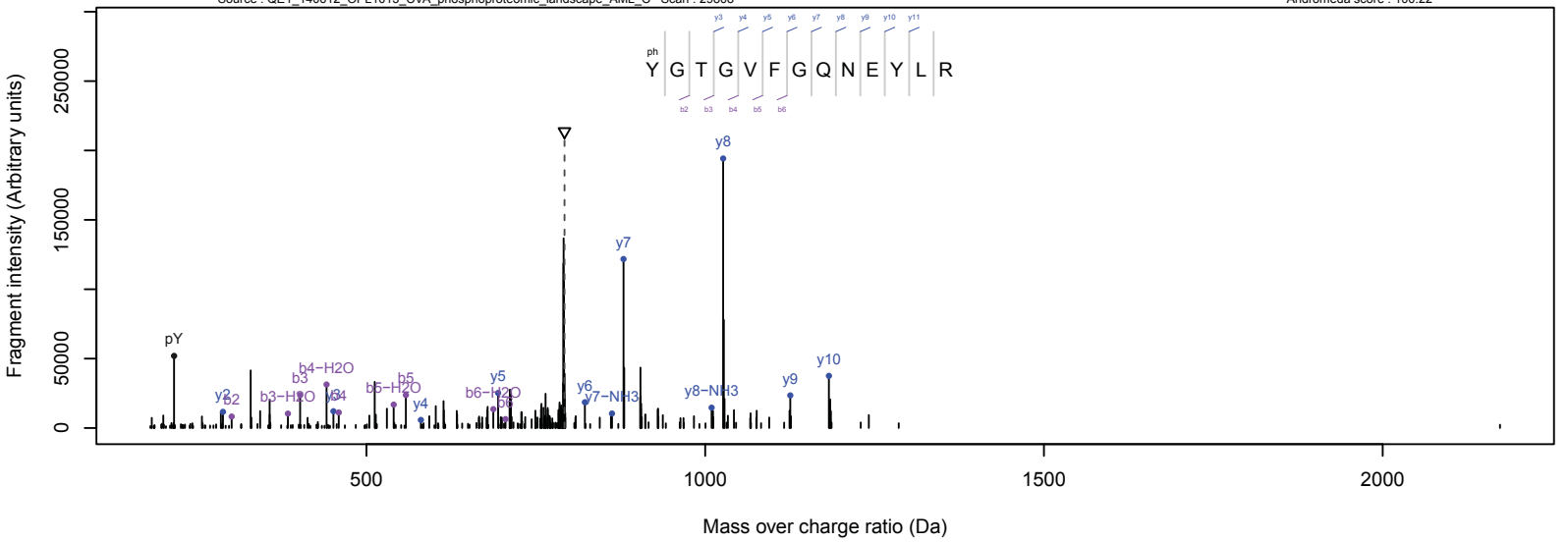
Andromeda score : 100.22



Y(ph)GTGVFGQNEYLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29808

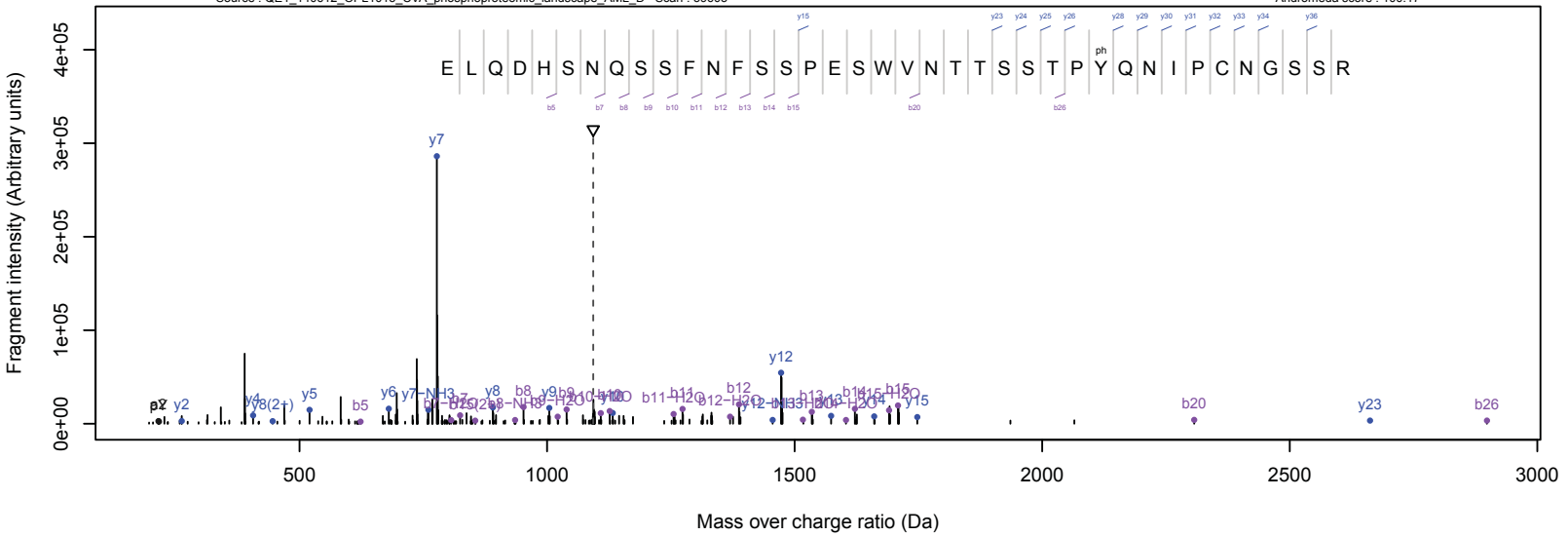
Andromeda score : 100.22



ELQDHSNQSSFNFSSPEWVNTTSSTPY(ph)QNIPCGSSR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 39603

Andromeda score : 100.17

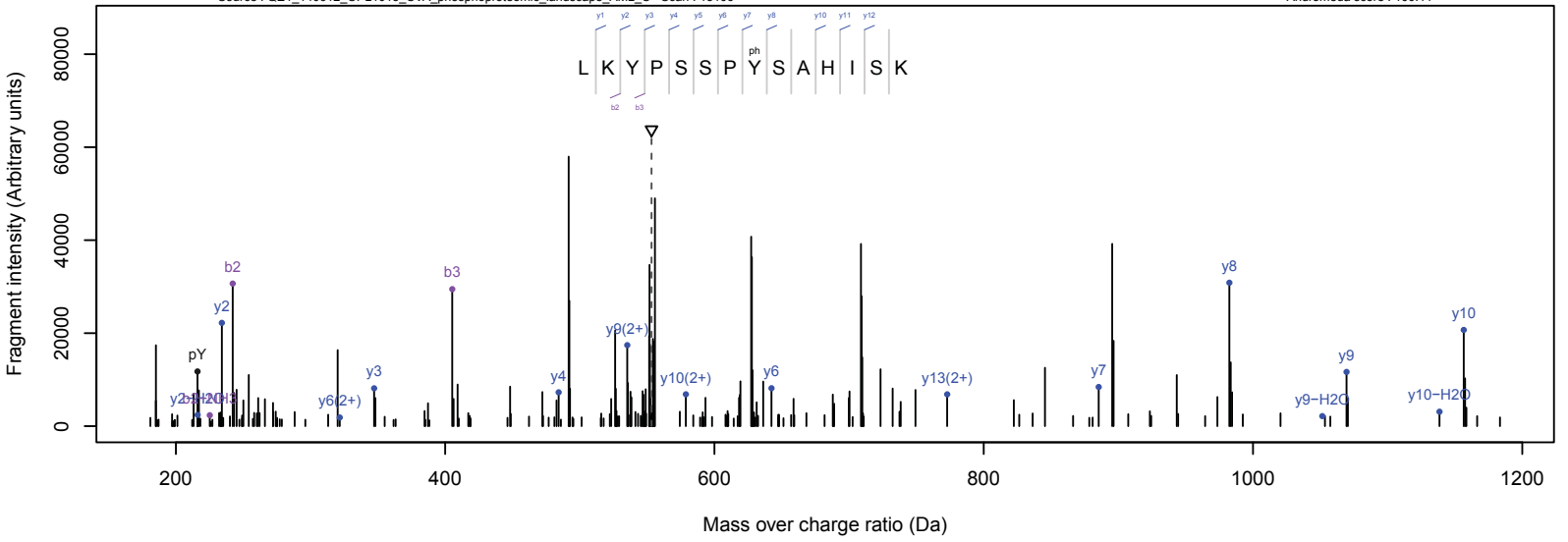




LKYPSPY(ph)SAHISK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13106

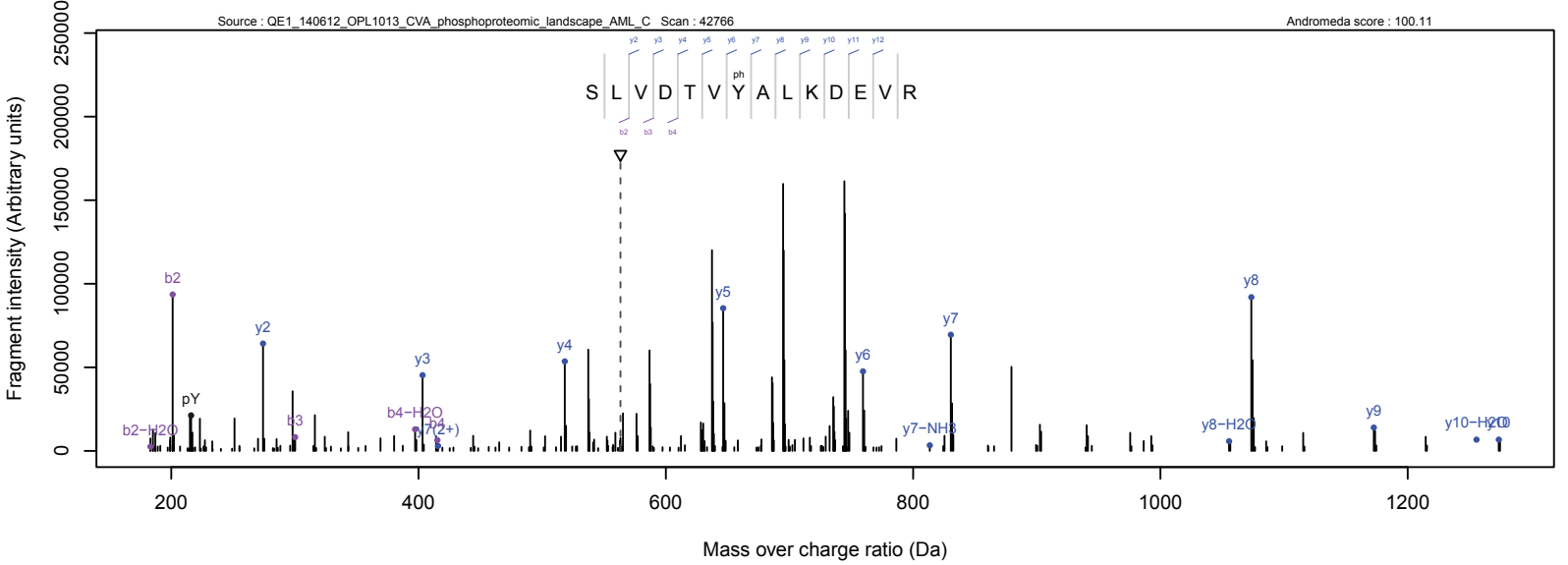
Andromeda score : 100.11



SLVDTVY(ph)ALKDEVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42766

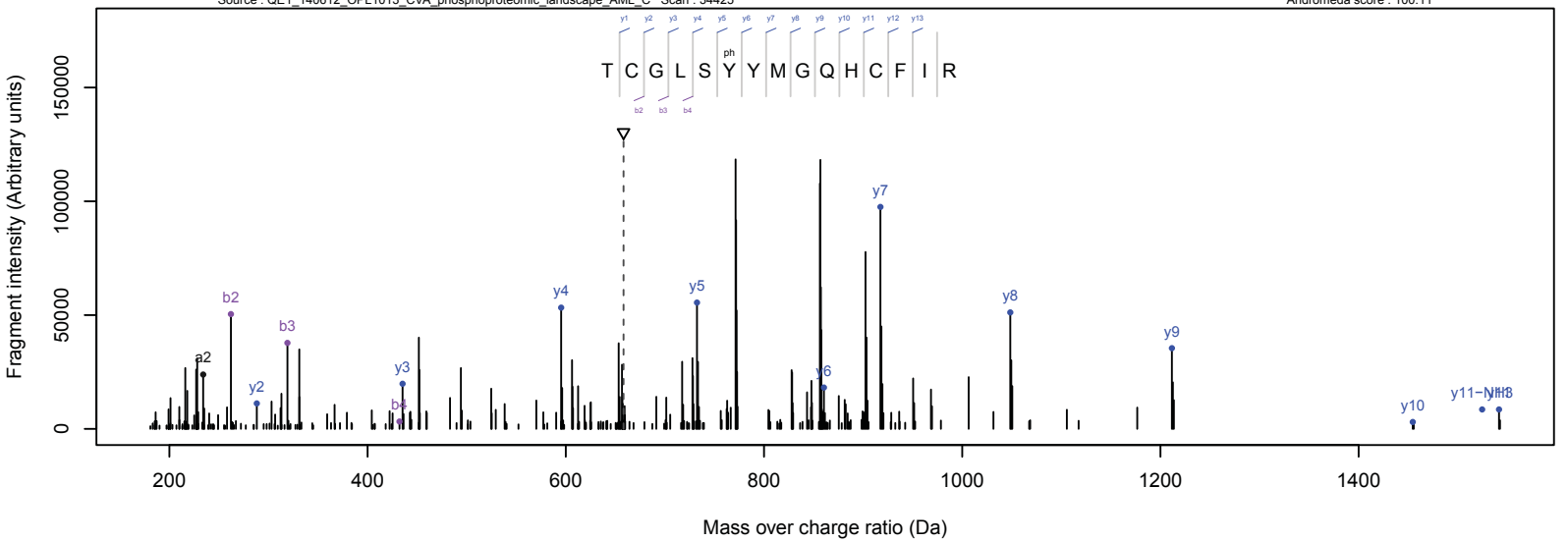
Andromeda score : 100.11

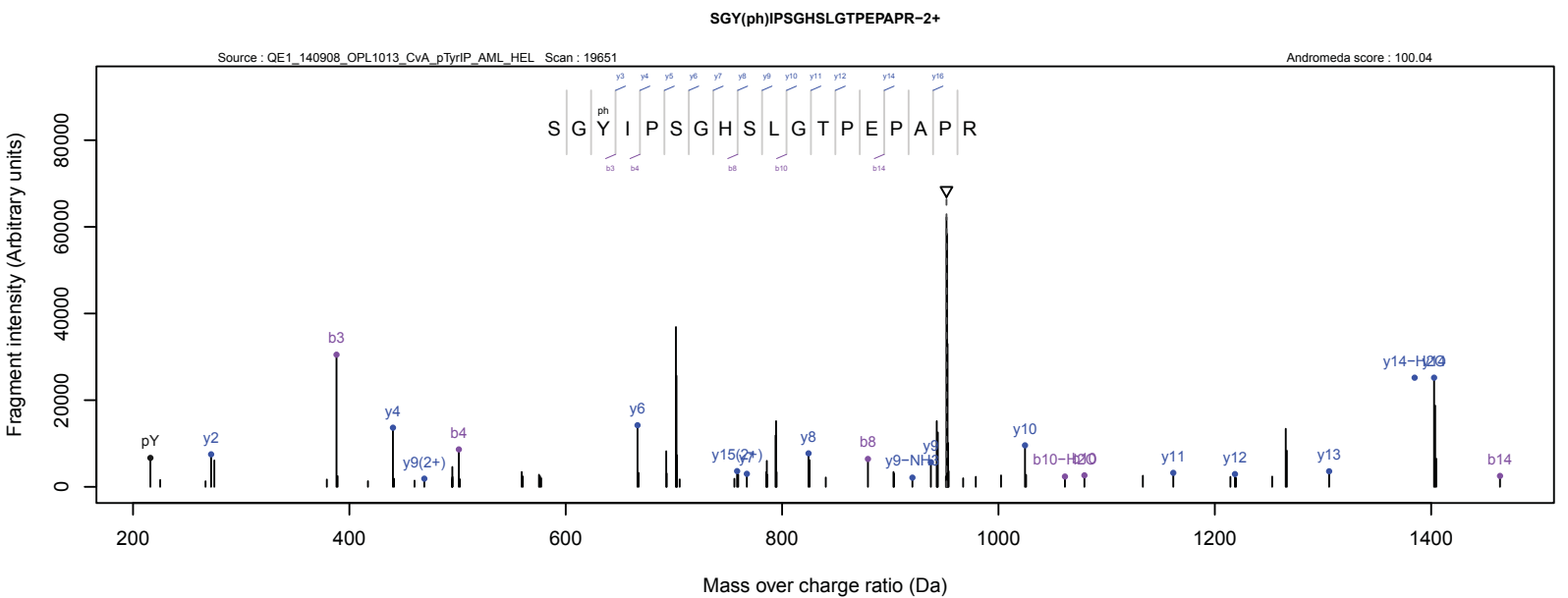
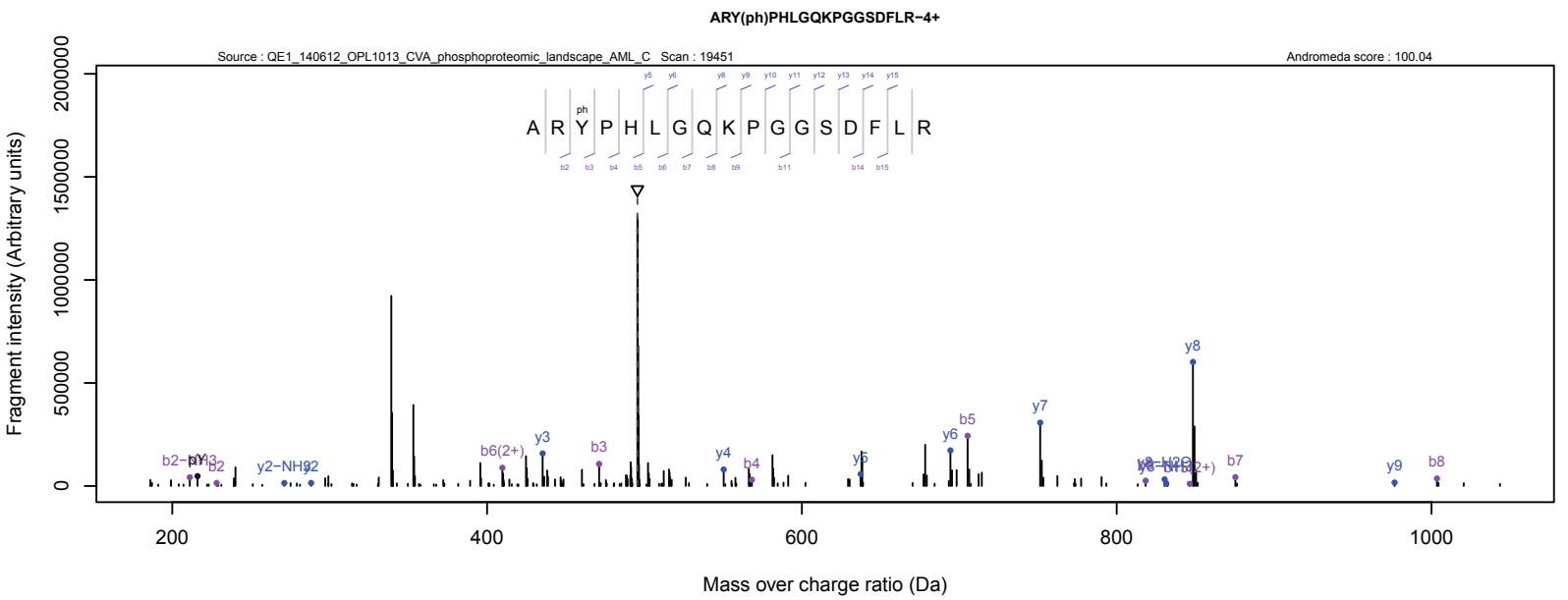
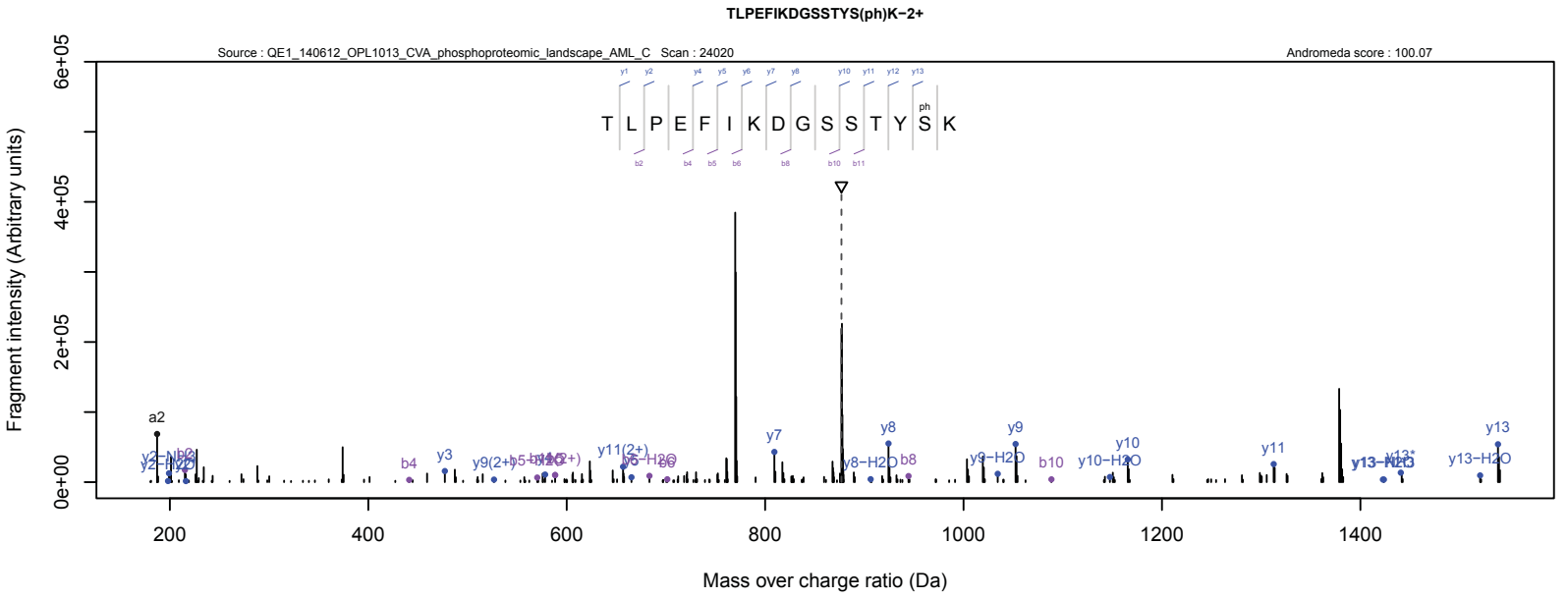


TCGLSY(ph)YMGQHC FIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34425

Andromeda score : 100.11

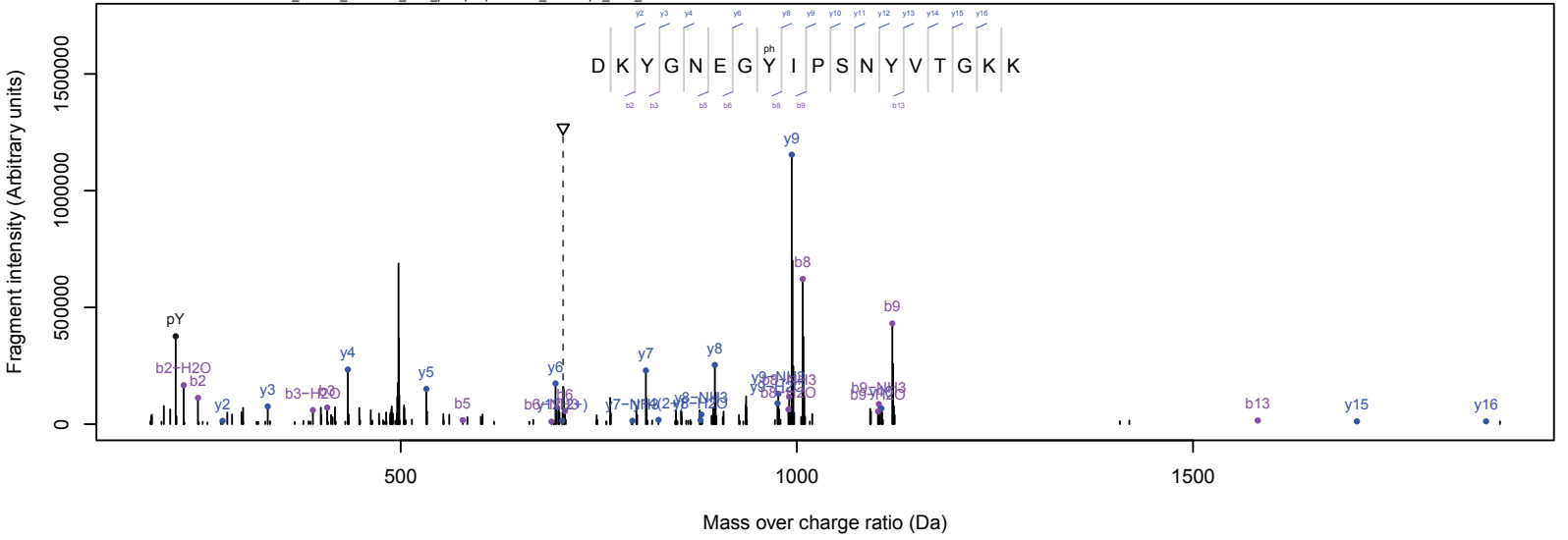




DKYGNEGY(ph)IPSNYVTGKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18134

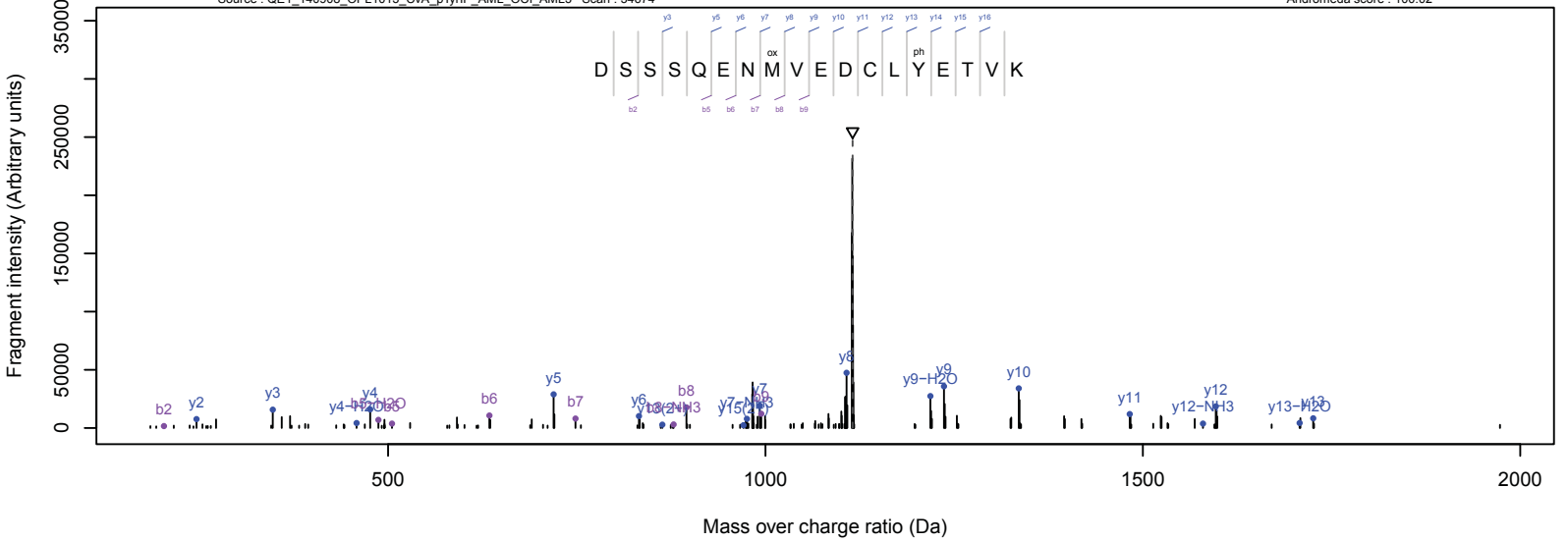
Andromeda score : 100.02



DSSSQENM(ox)VEDCLY(ph)ETVK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 34674

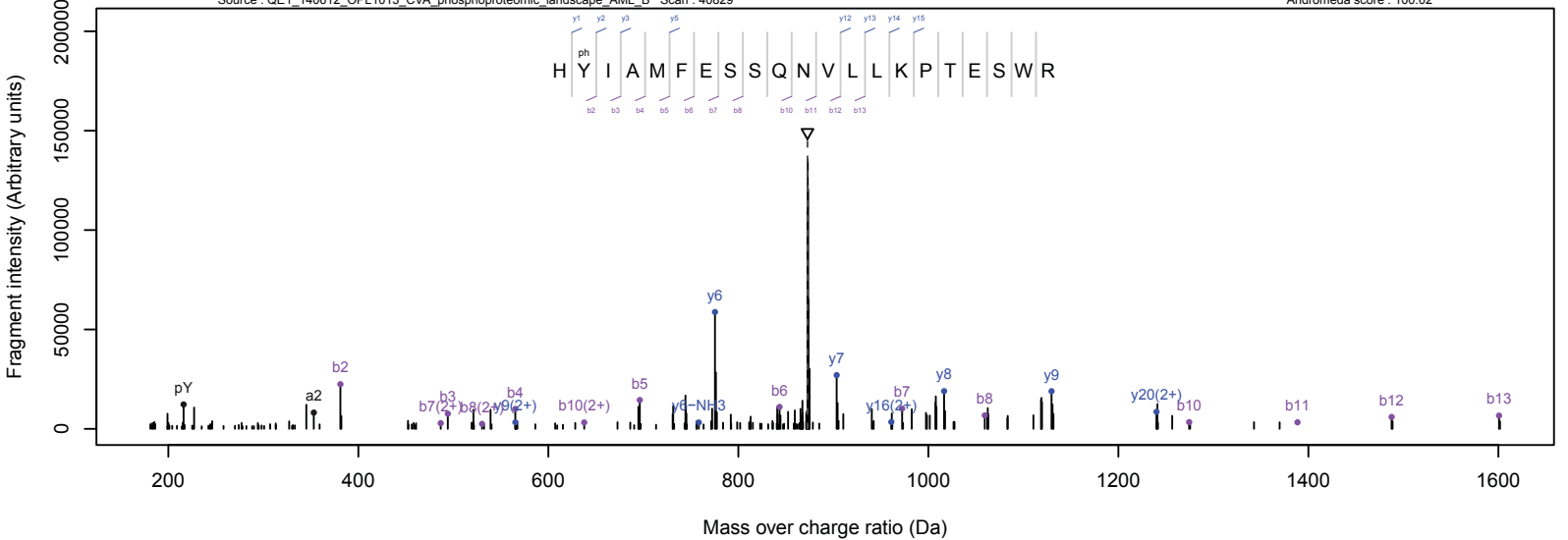
Andromeda score : 100.02



HY(ph)IAMFESSQNVLKPTESWR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40829

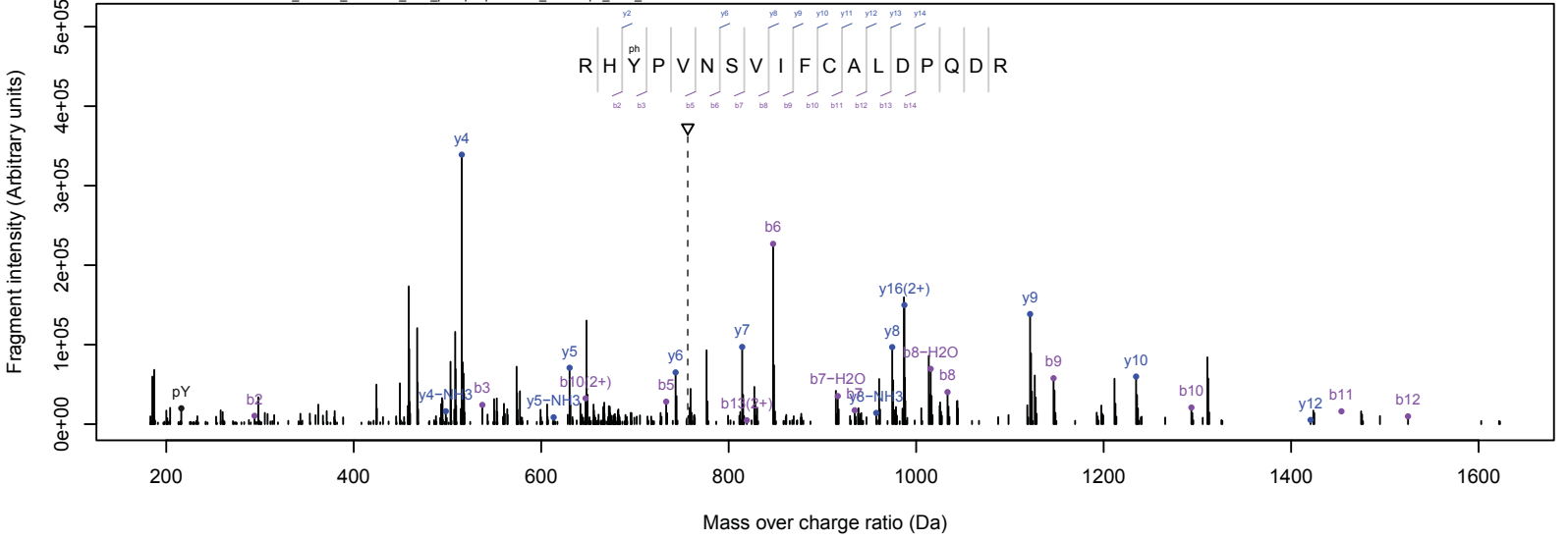
Andromeda score : 100.02



RHY(ph)PVNSVIFCALDPQDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35966

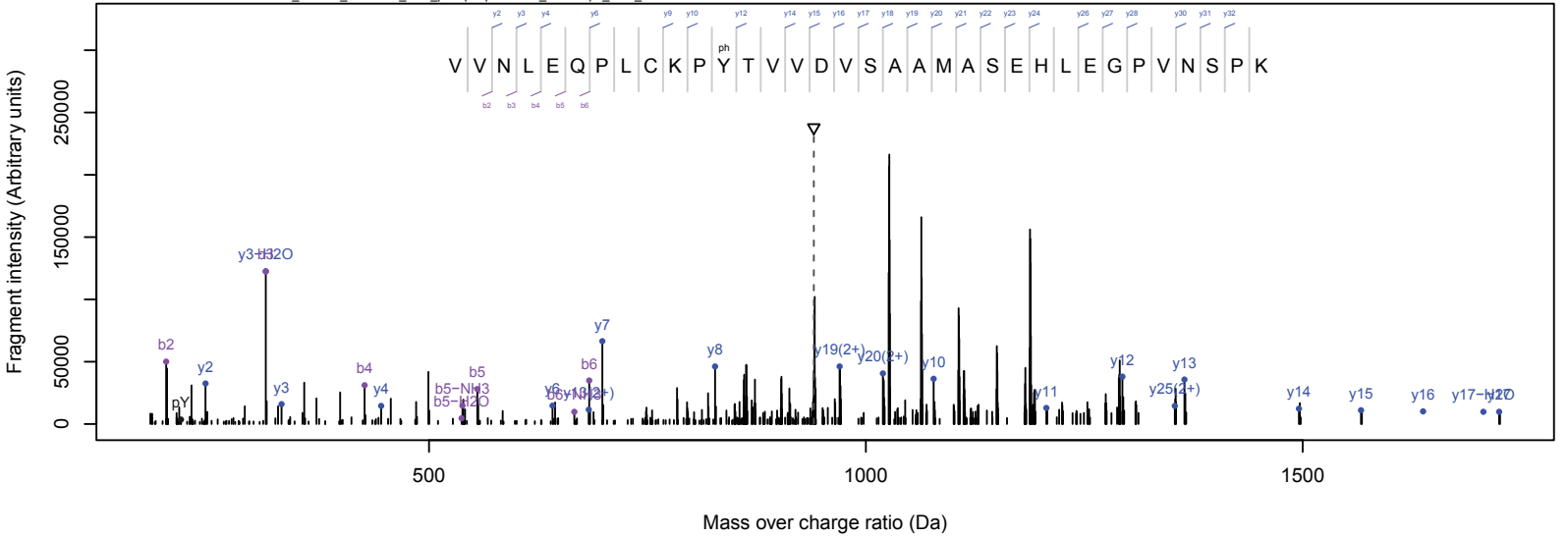
Andromeda score : 100.02



VVNLEQLCKPY(ph)TVVDVSAAMASEHLEGPVNSPK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44656

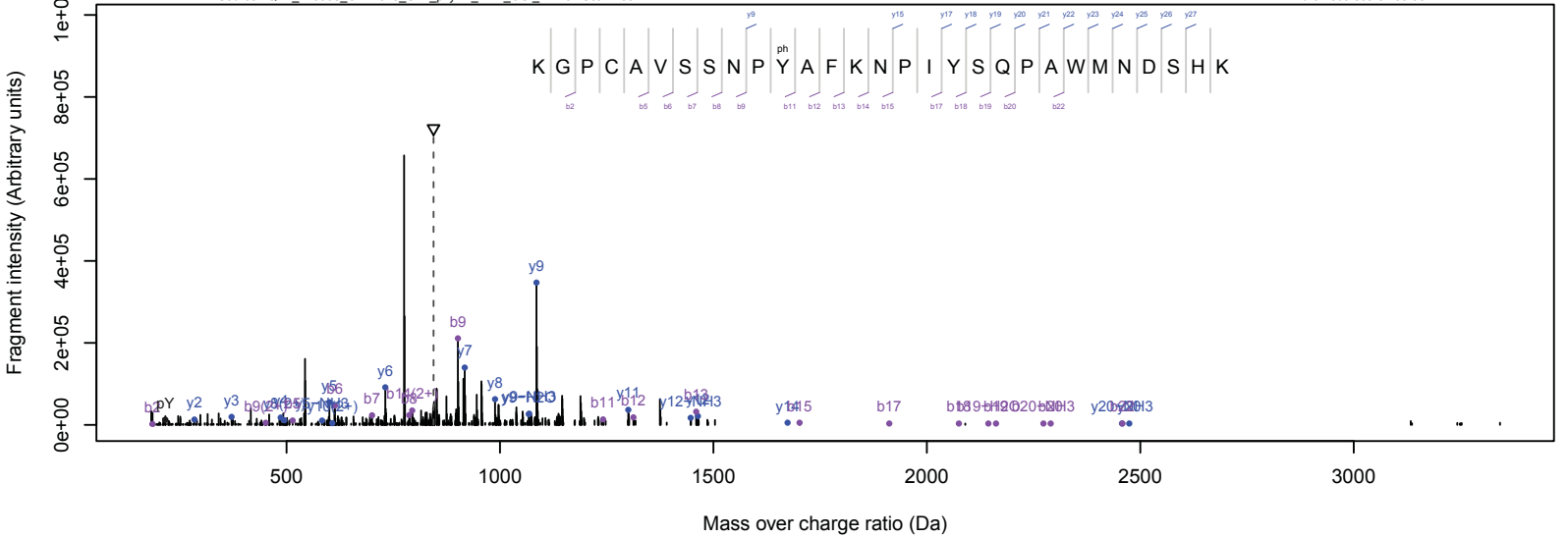
Andromeda score : 99.997



KGPCAVSSNPY(ph)YAFKNPIYSQPAWMNDSHK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 33447

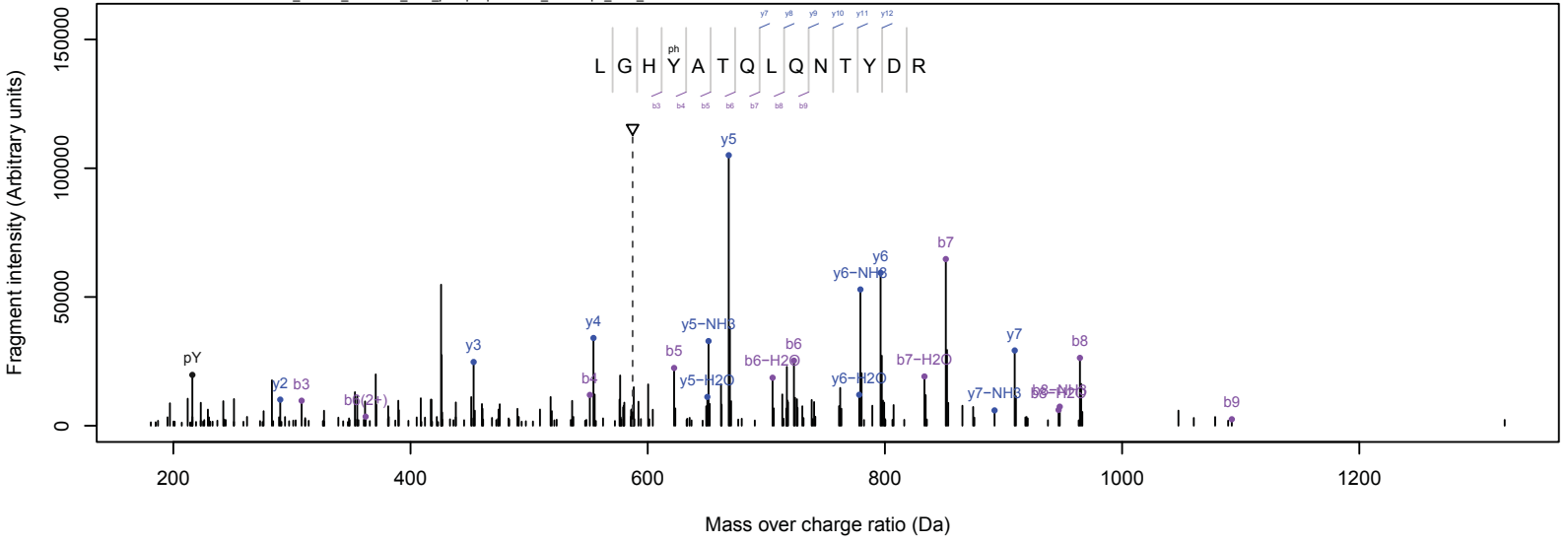
Andromeda score : 99.987



LGHY(ph)ATQLQNTYDR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15592

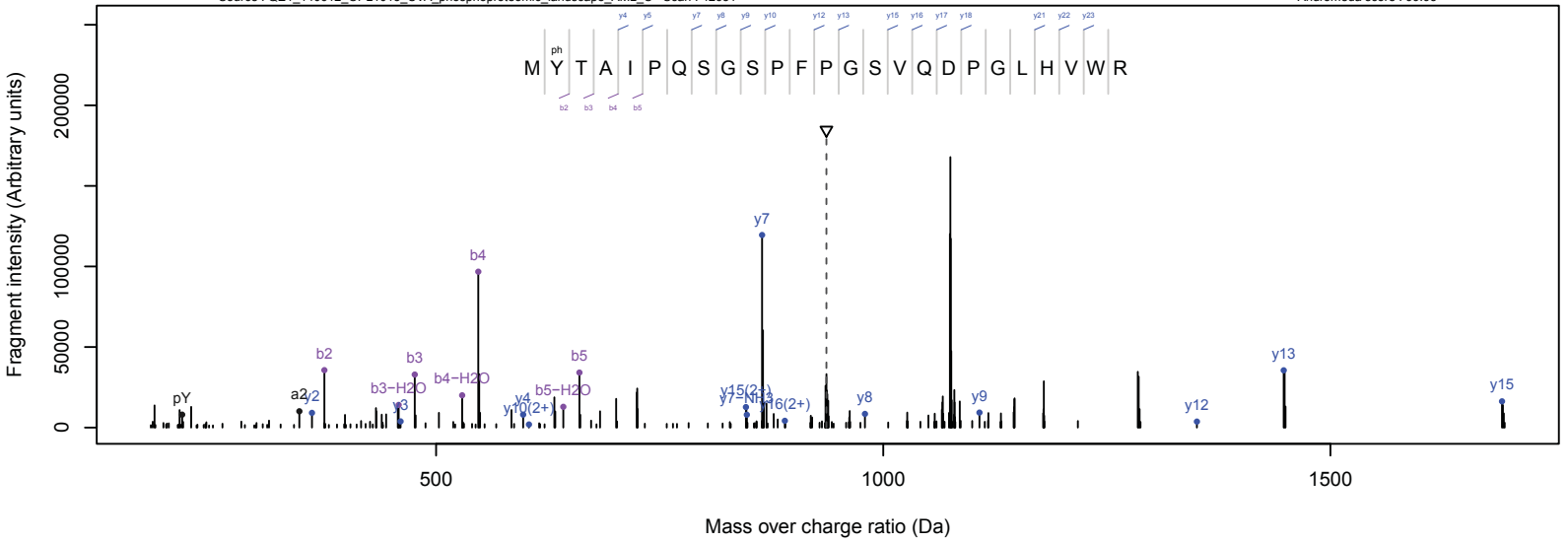
Andromeda score: 99.941



MY(ph)TAIPQSGSPFPQSGVQDPLHVWR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 42631

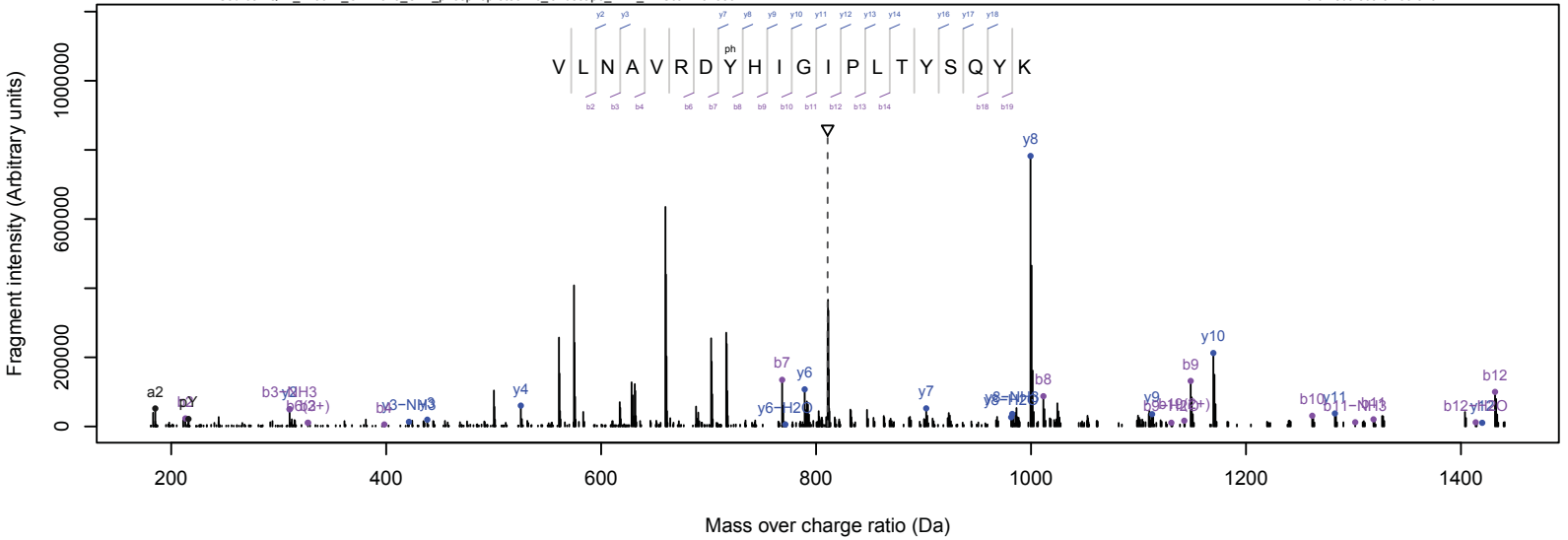
Andromeda score: 99.93



VLNAVRDY(ph)HIGIPLTYSQYK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34580

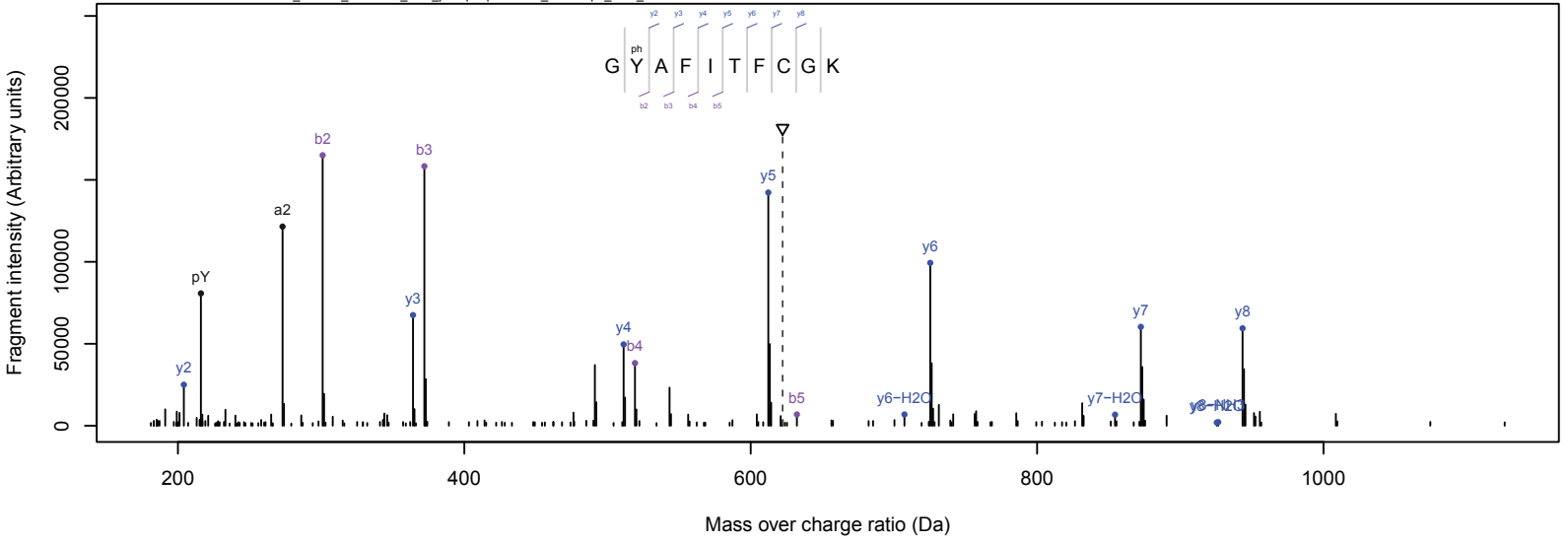
Andromeda score: 99.873



**GY(ph)AFITCGK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 33485

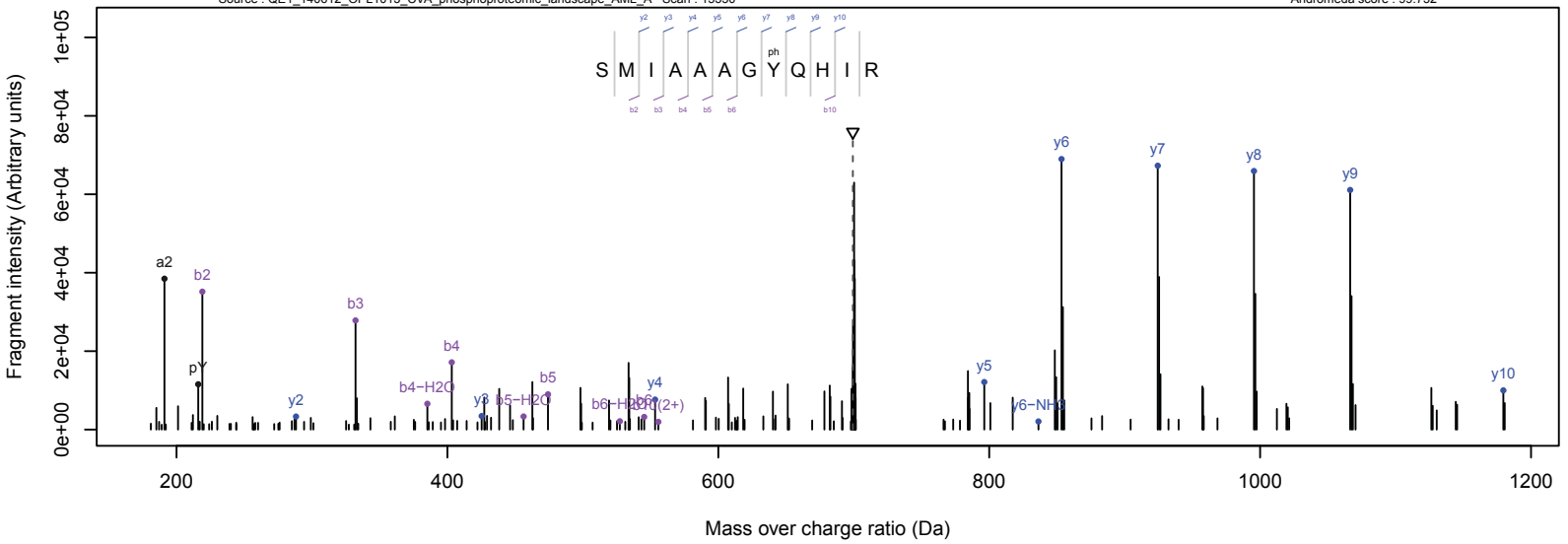
Andromeda score: 99.815



**SMIAAGY(ph)QHIR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 15350

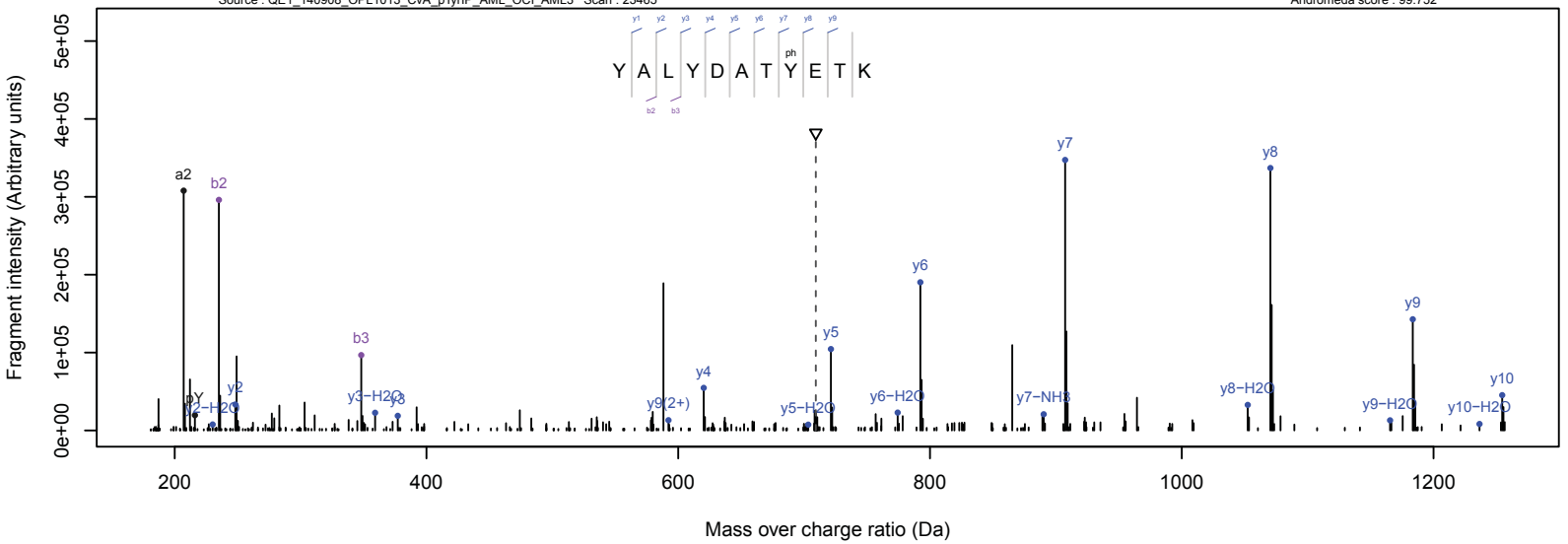
Andromeda score: 99.752



**YALYDATY(ph)ETK-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 23465

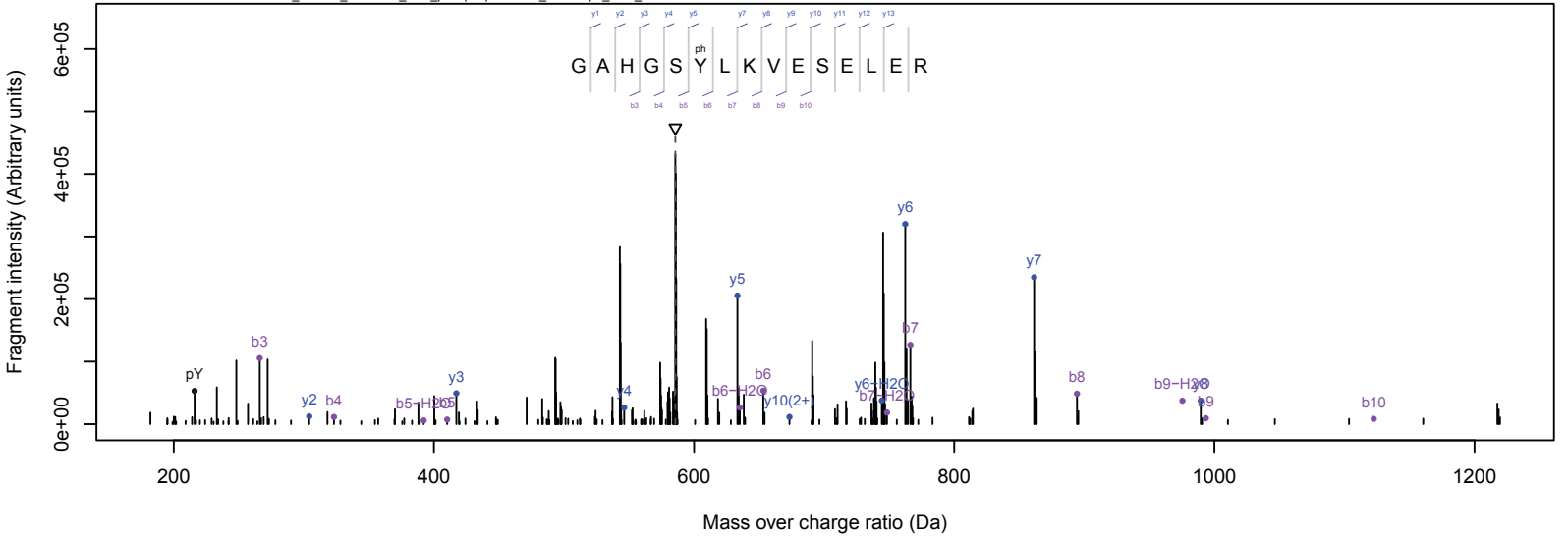
Andromeda score: 99.752



**GAHGSY(ph)LKVESELER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17355

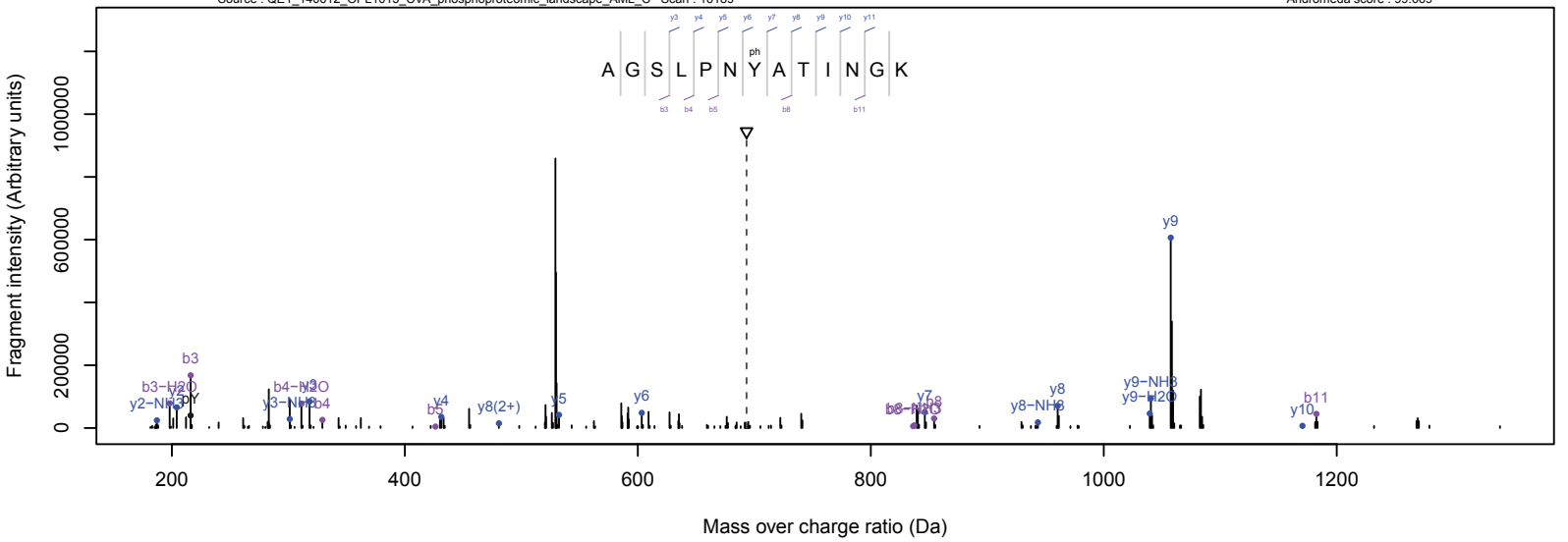
Andromeda score : 99.747



**AGSLPNY(ph)ATINGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16189

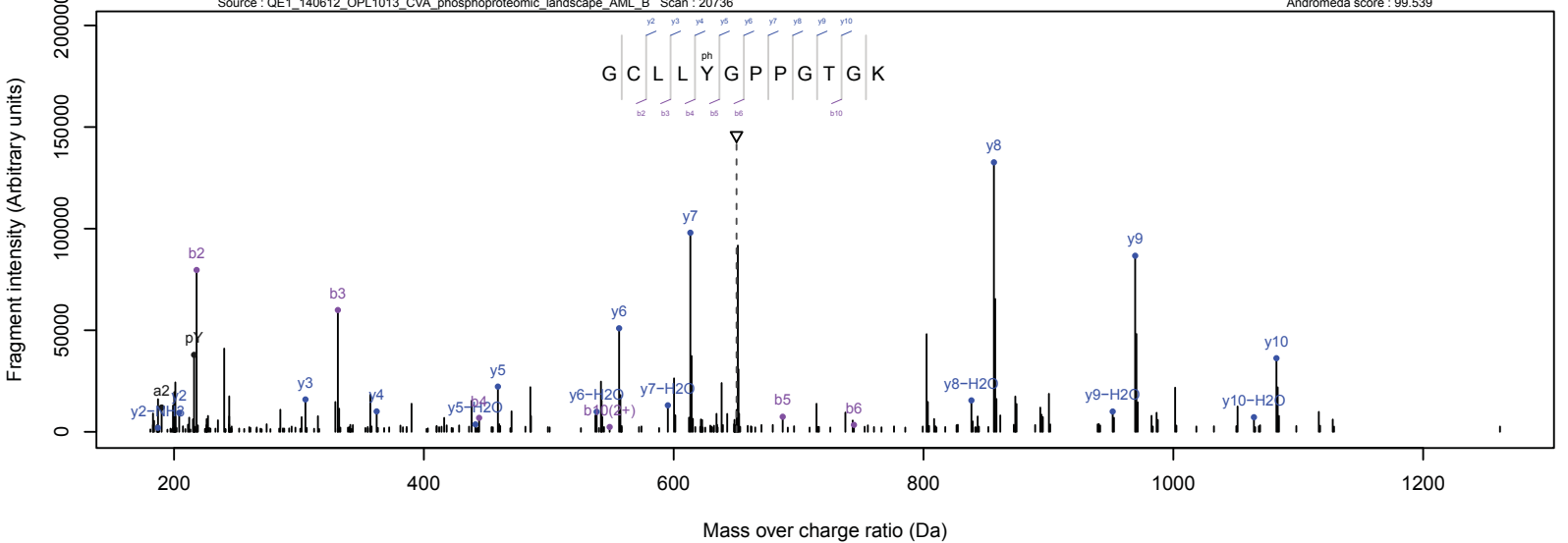
Andromeda score : 99.669



**GCLLY(ph)GPPGTGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20736

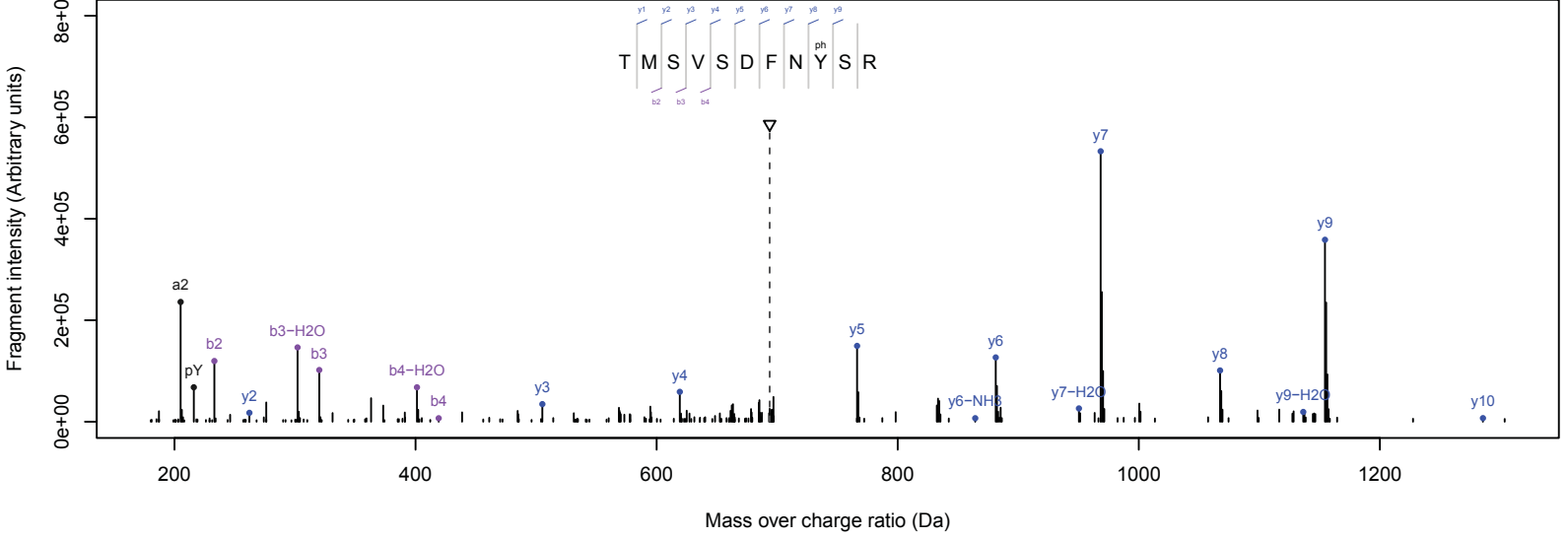
Andromeda score : 99.539



TMSVDFNY(ph)SR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19365

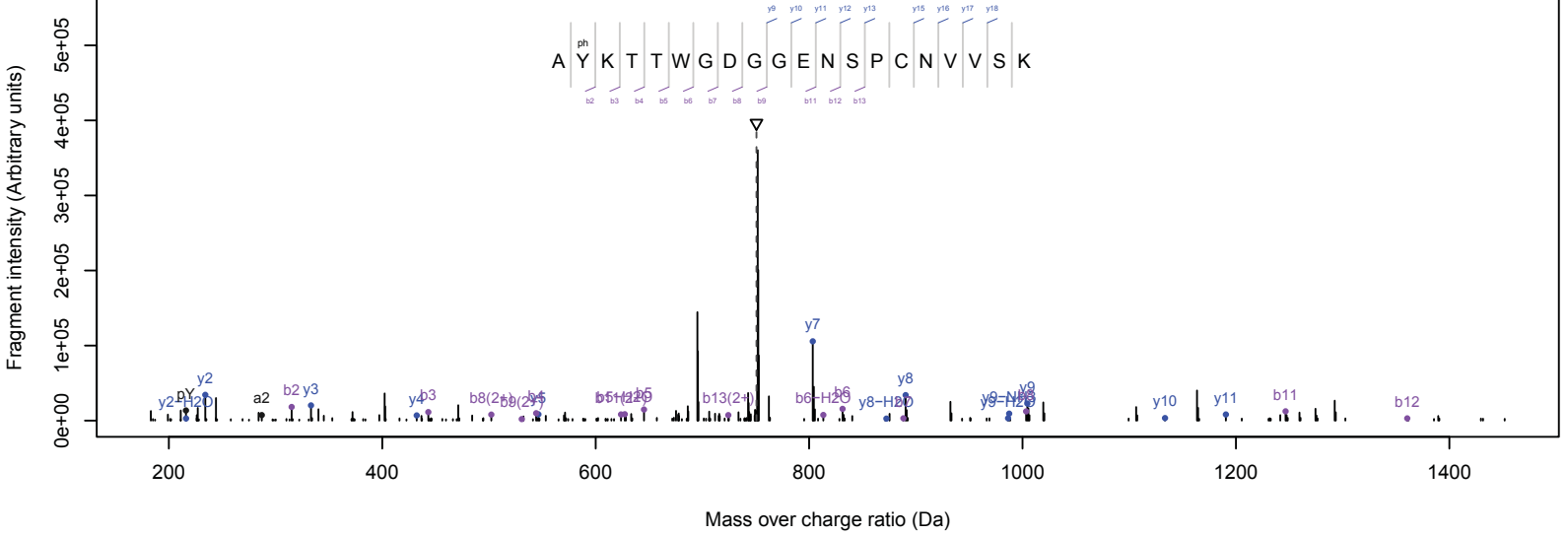
Andromeda score : 99.5



AY(ph)KTTWGDGGENSPCNVSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16636

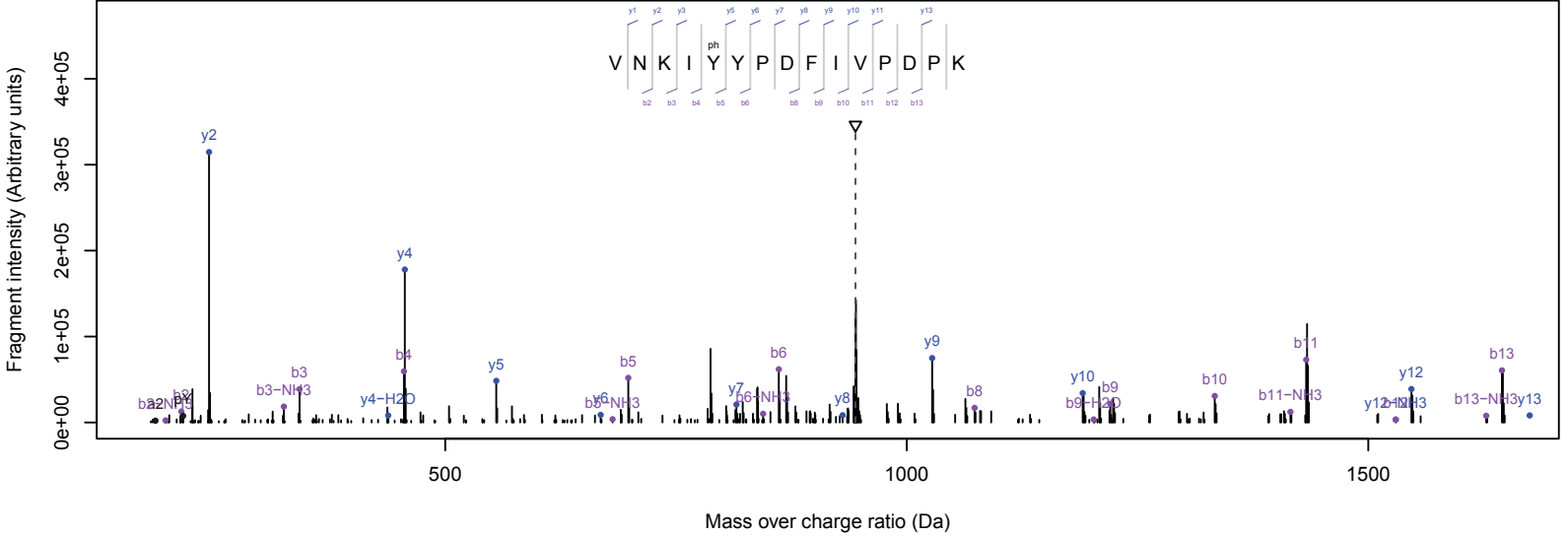
Andromeda score : 99.409



VNKIYYPDFIVDPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 38819

Andromeda score : 99.392

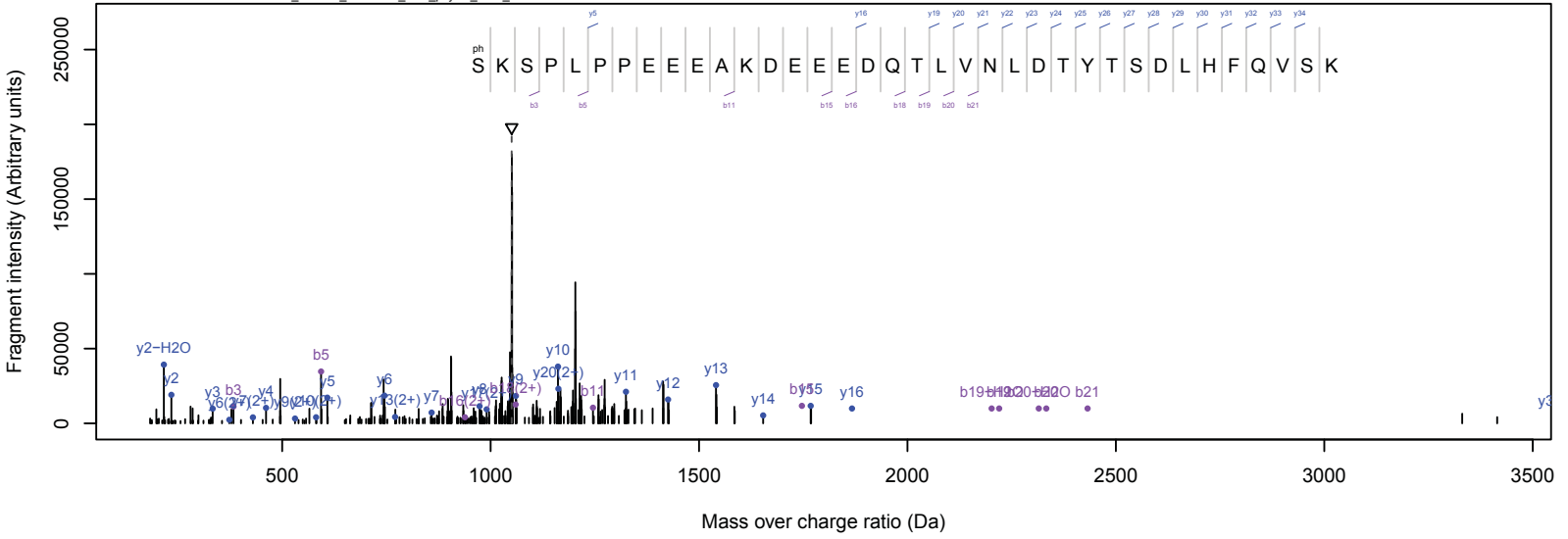




**S(ph)KSPLPPEEEAKDEEEDQLVNLDTYTSDLHFQVSK-4+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan: 45989

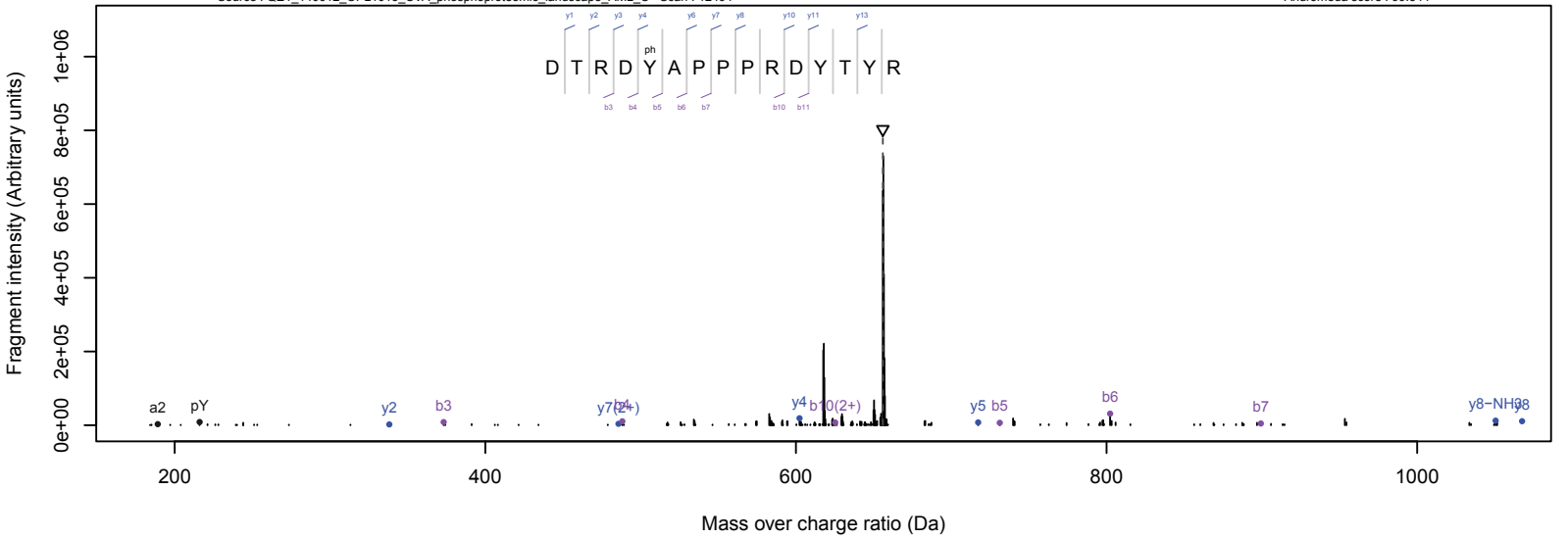
Andromeda score: 99.357



**DTRDY(ph)APPRDYTYR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12494

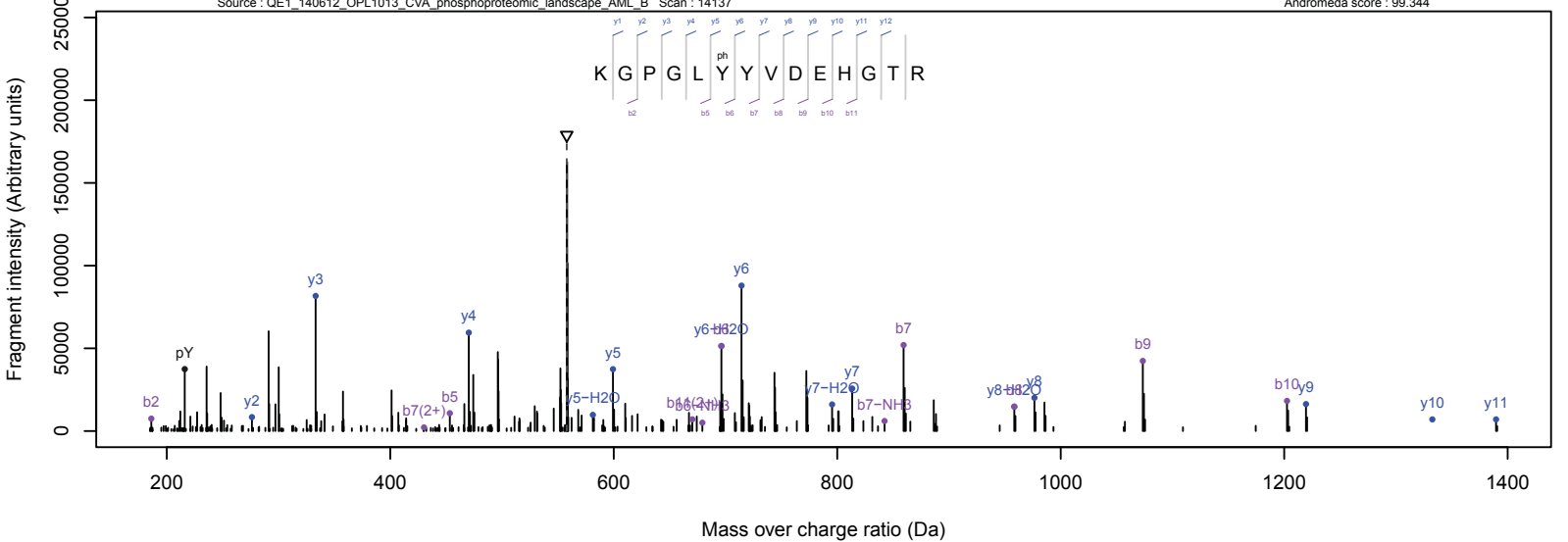
Andromeda score: 99.344



**KGPLY(ph)YVDEHGTR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14137

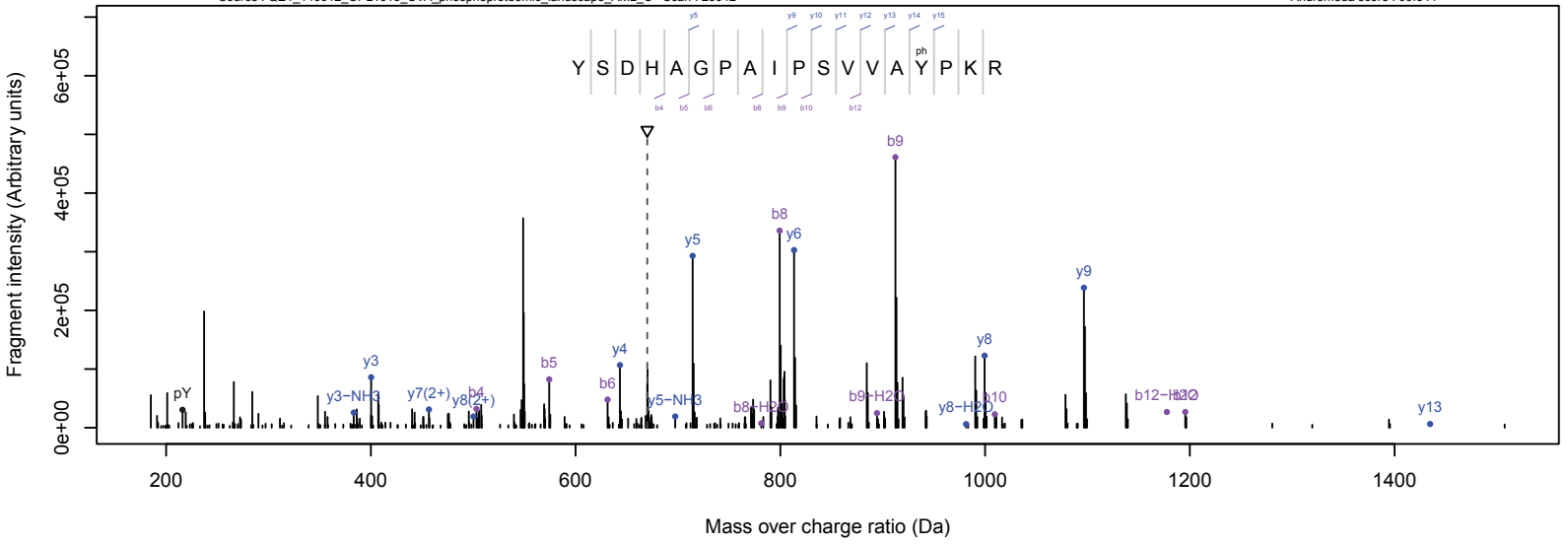
Andromeda score: 99.344



YSDHAGPAIPSVVAY<sup>ph</sup>PKR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23042

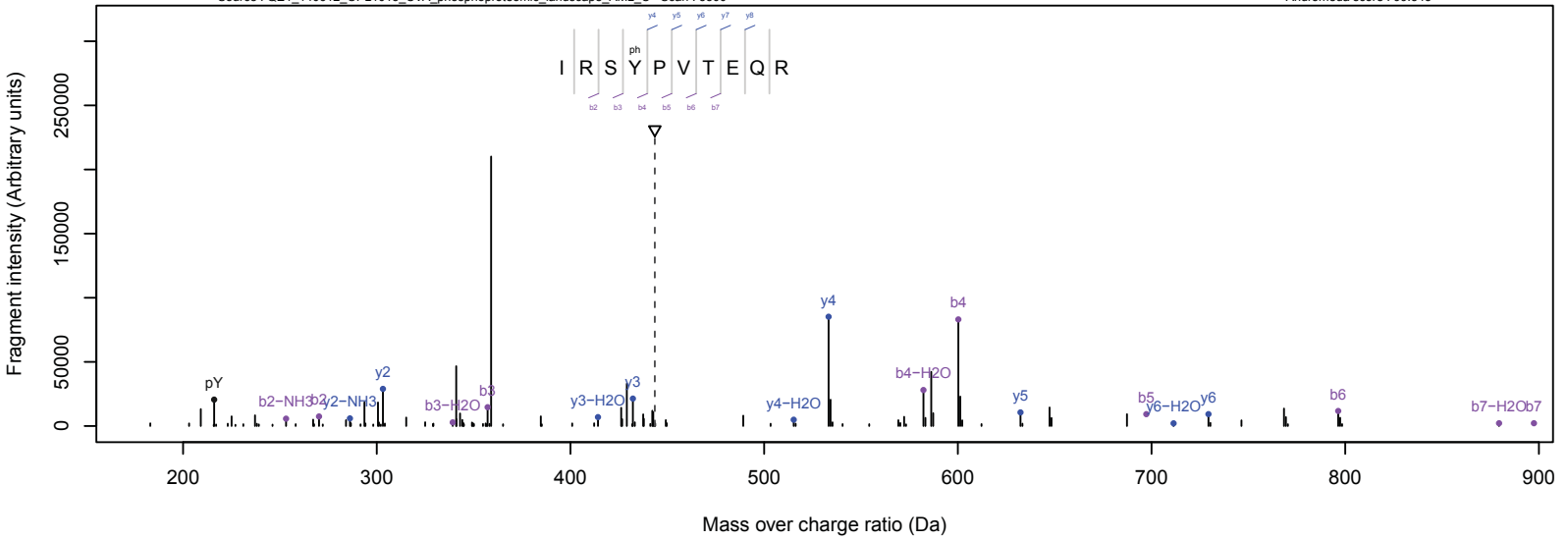
Andromeda score: 99.344



IRSY<sup>ph</sup>PVTEQR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 5300

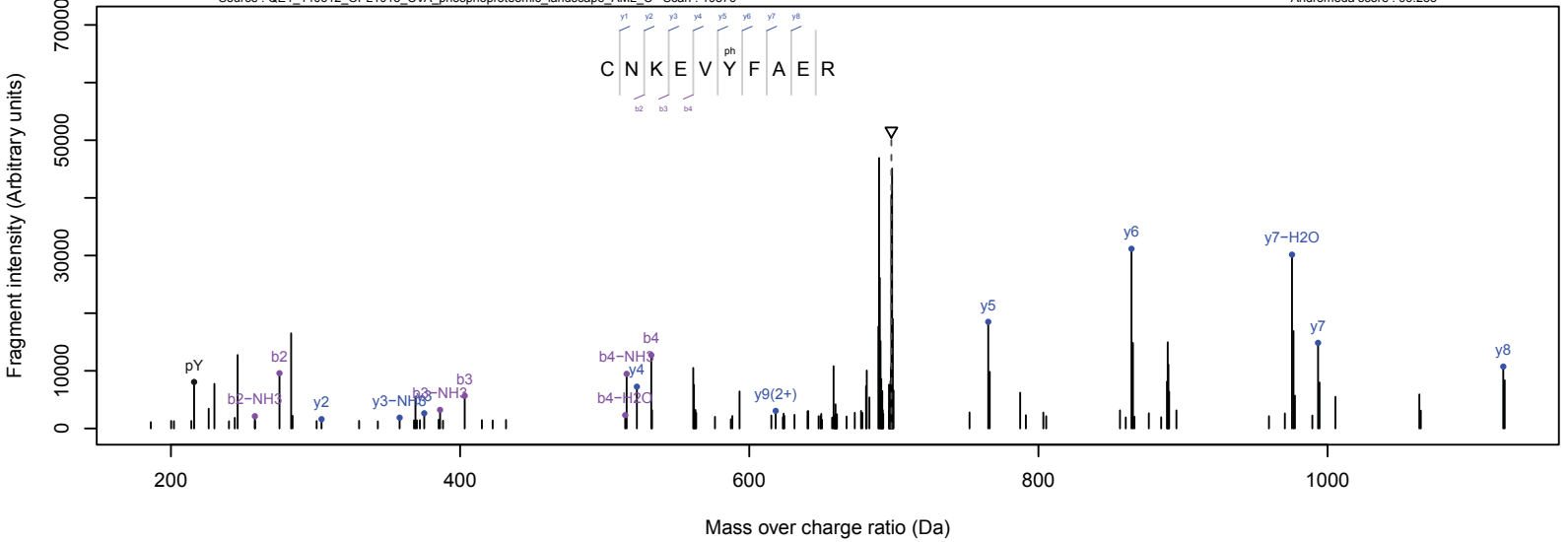
Andromeda score: 99.343



CNKEVY<sup>ph</sup>FAER-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10876

Andromeda score: 99.283



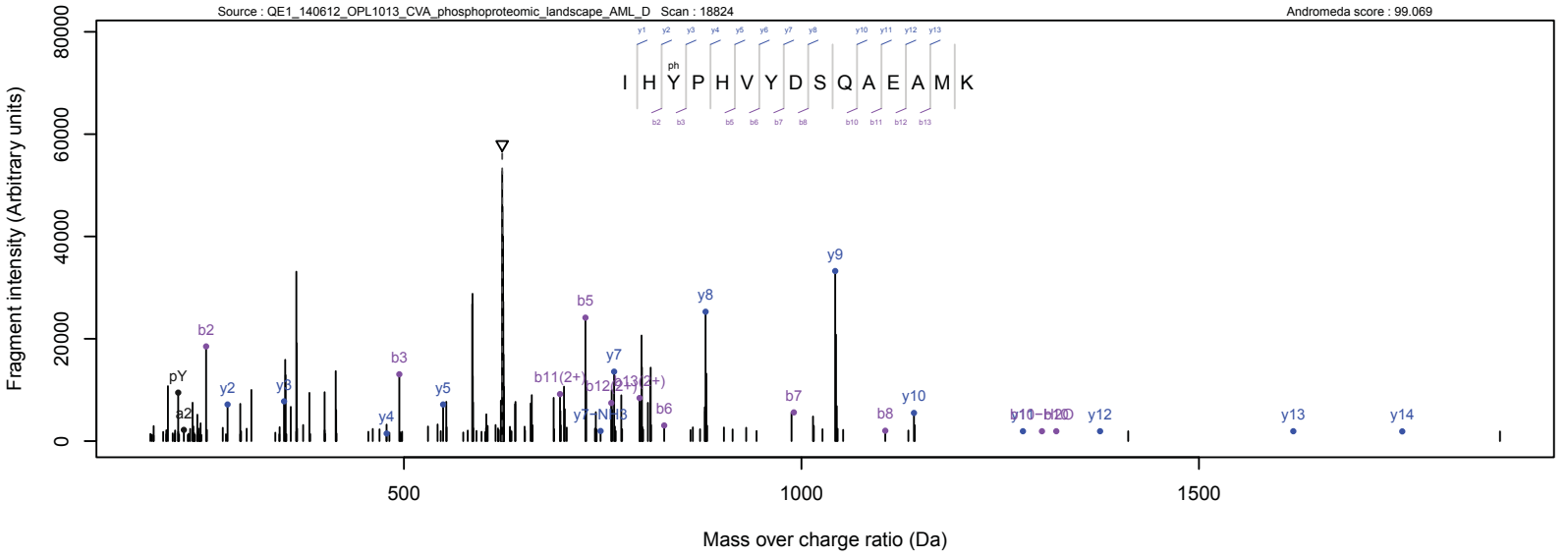




IHY(ph)PHVYDSQAEAMK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 18824

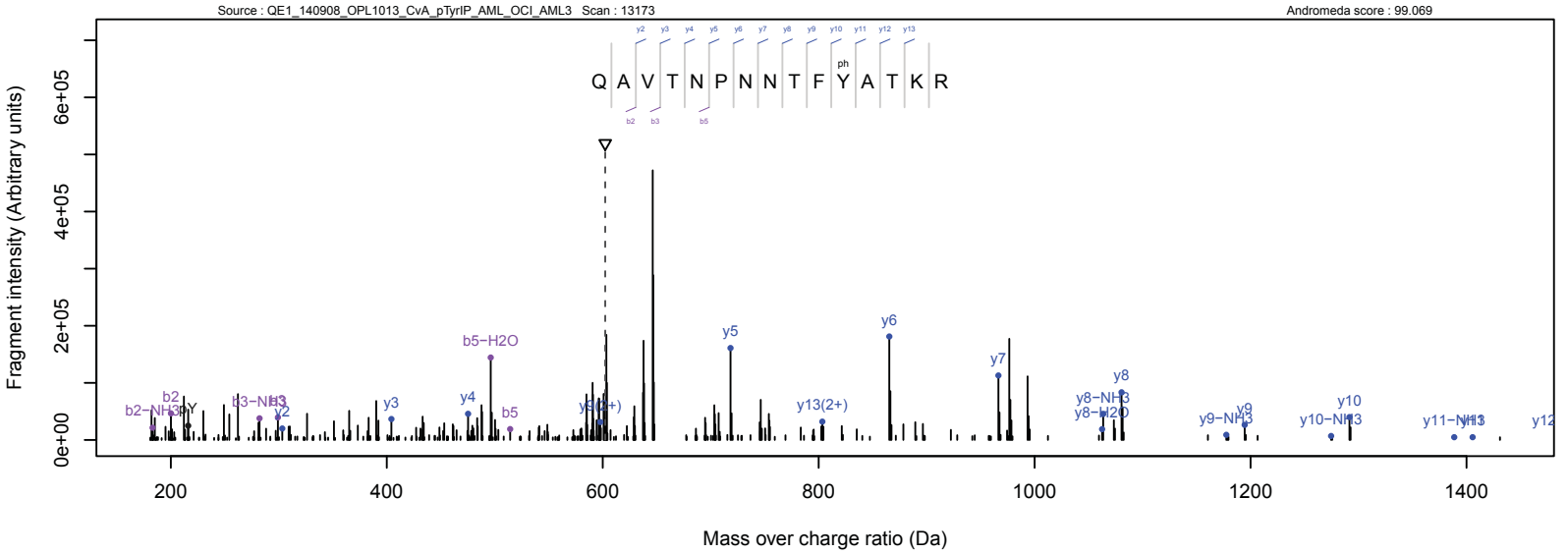
Andromeda score : 99.069



QAVTNPNTFY(ph)ATKR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 13173

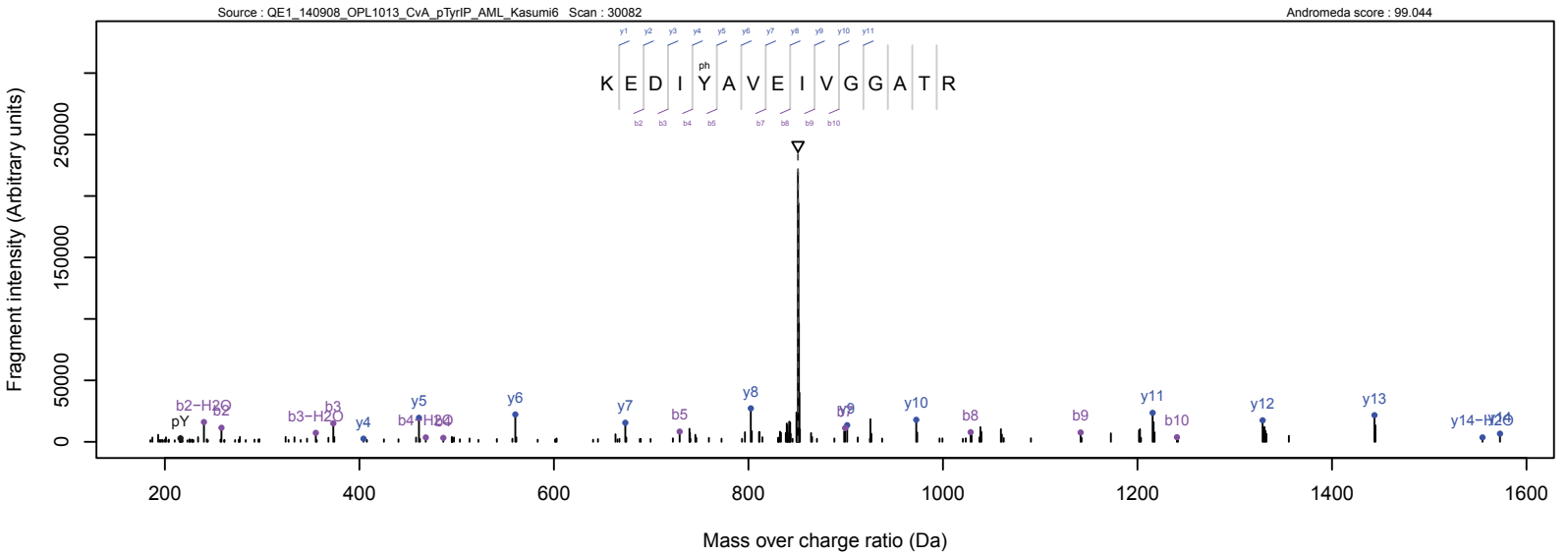
Andromeda score : 99.069



KEDIY(ph)AVEIVGGATR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 30082

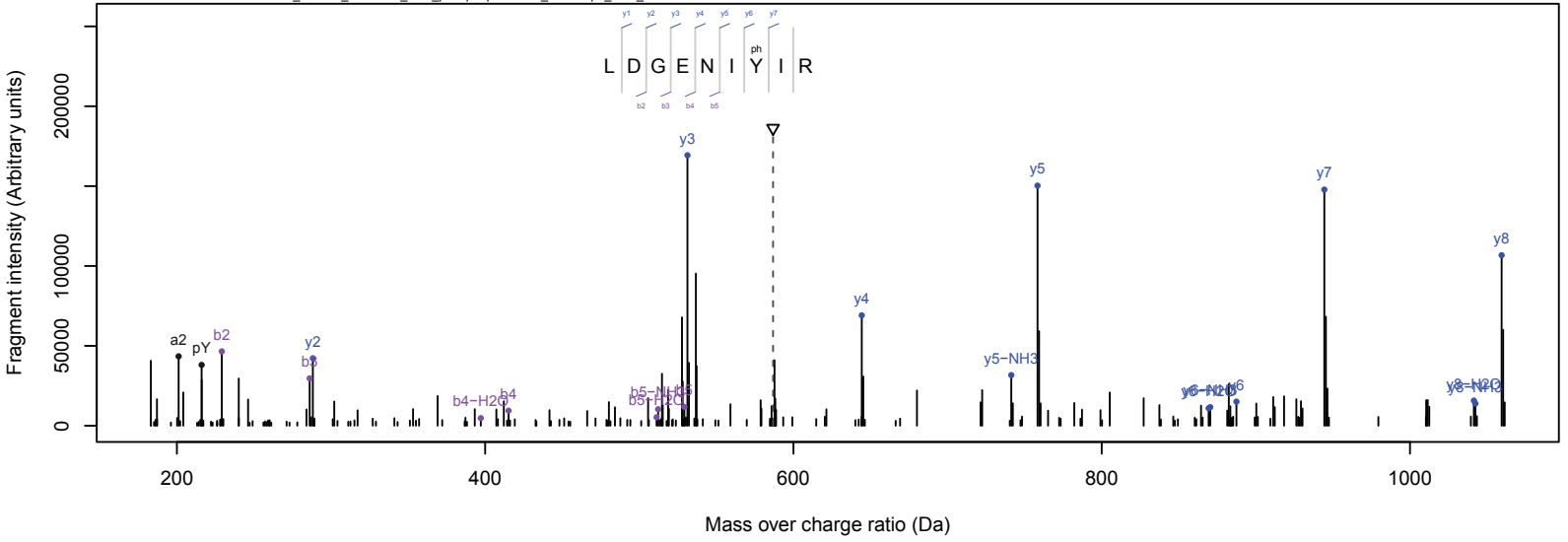
Andromeda score : 99.044



LDGENIY(ph)IR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21211

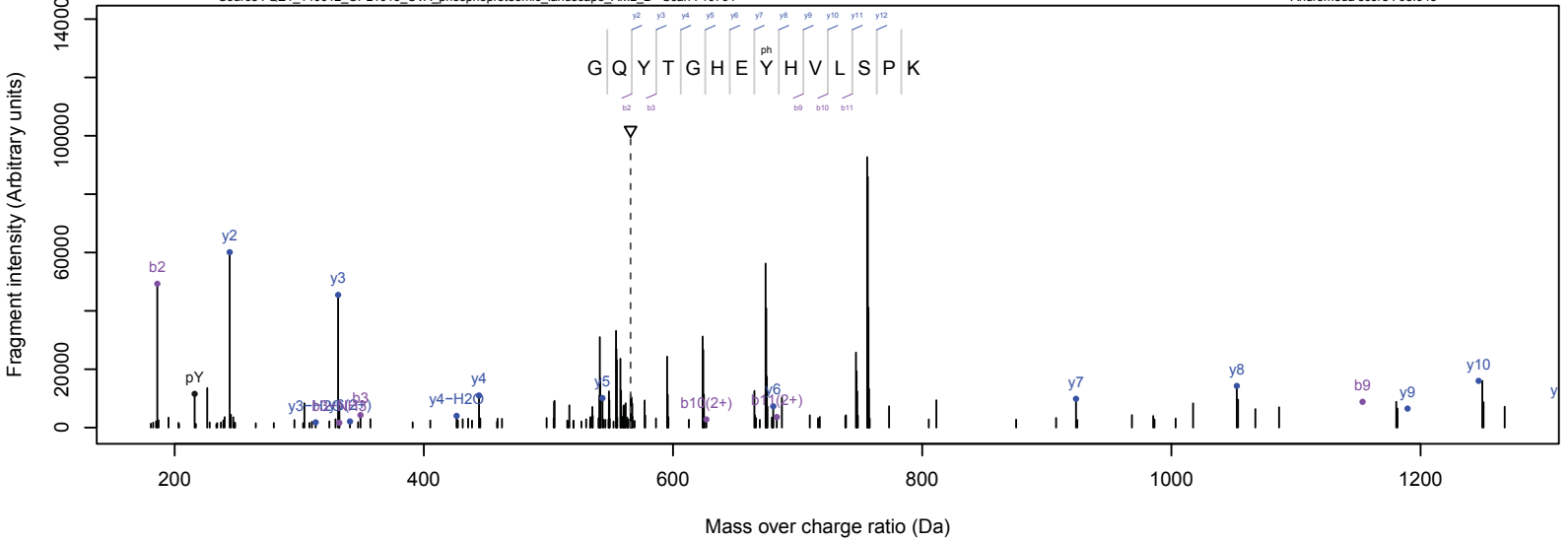
Andromeda score : 99.013



GQYTGHEY(ph)HVLSPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10764

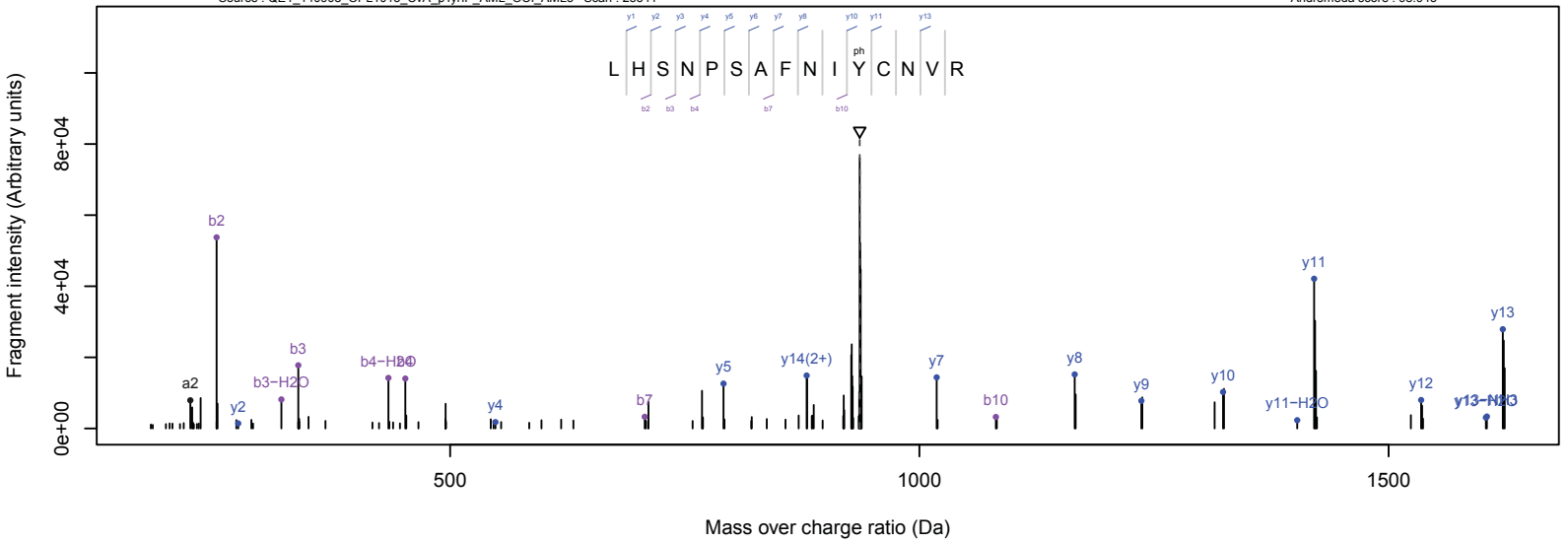
Andromeda score : 98.943



LHSNPSAFNIY(ph)CNVR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 25341

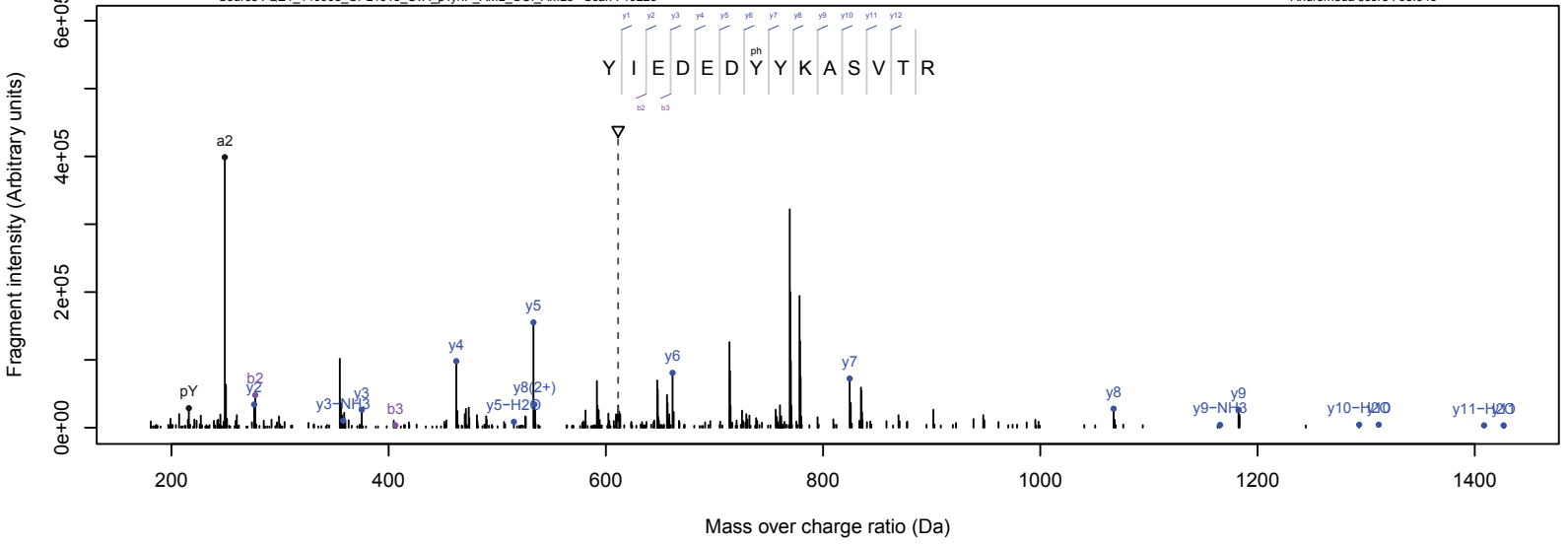
Andromeda score : 98.943



**YIEDEDY(ph)YKASVTR-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 19223

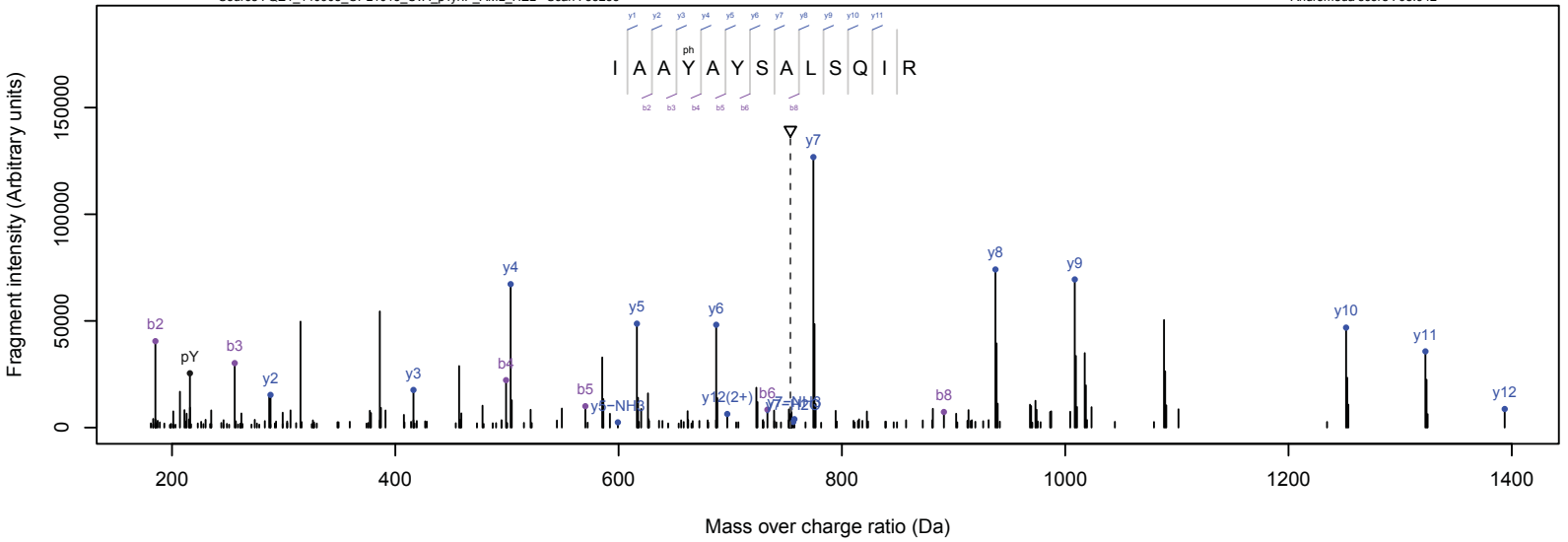
Andromeda score: 98.943



**IAAY(ph)AYSALSQIR-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 35255

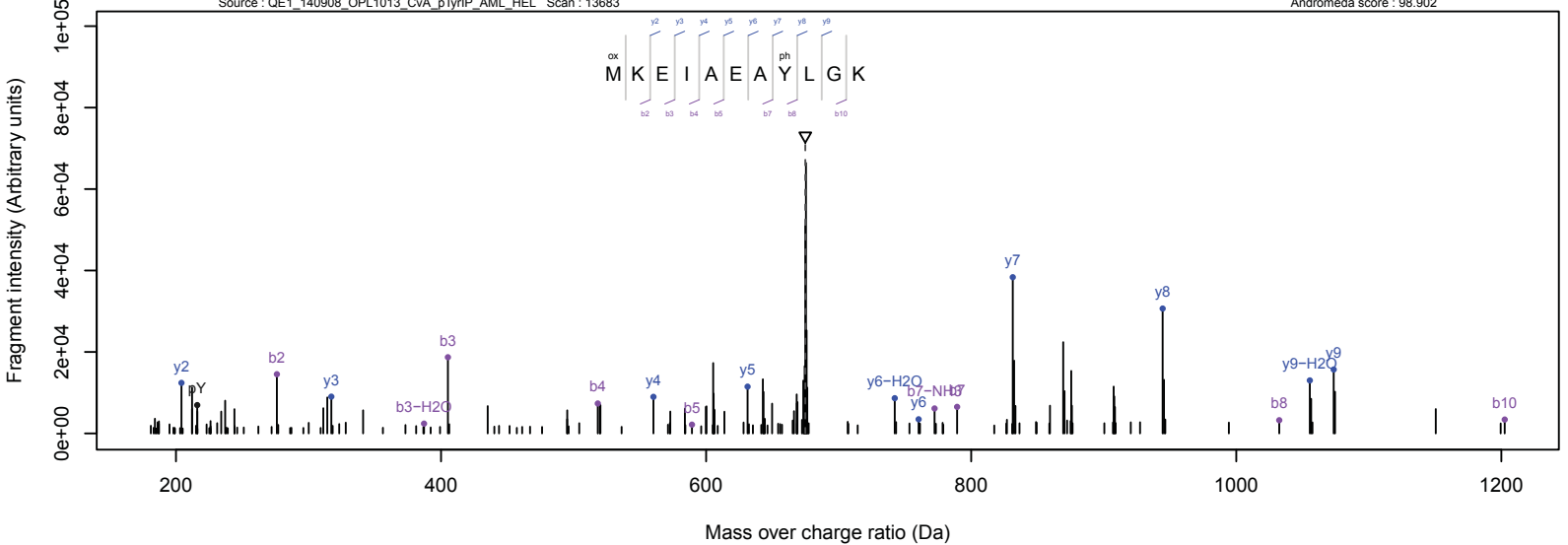
Andromeda score: 98.942



**M(ox)KEIAEAY(ph)LGK-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 13683

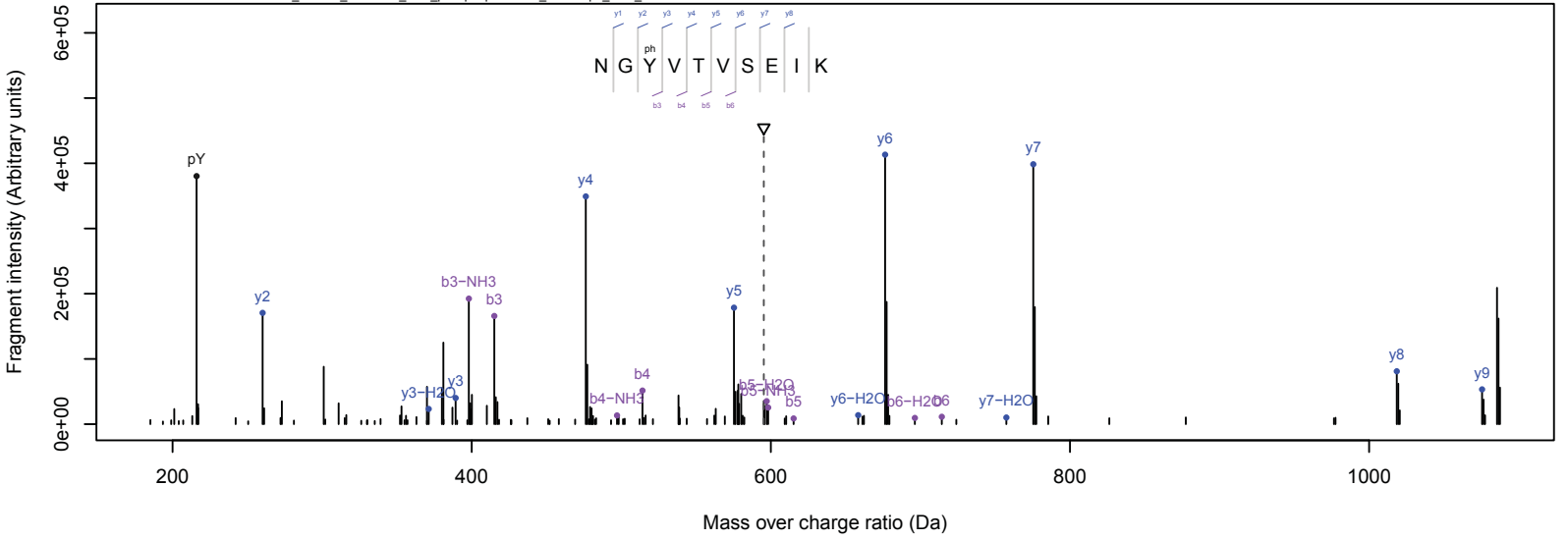
Andromeda score: 98.902



NGY(ph)VTVSEIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16136

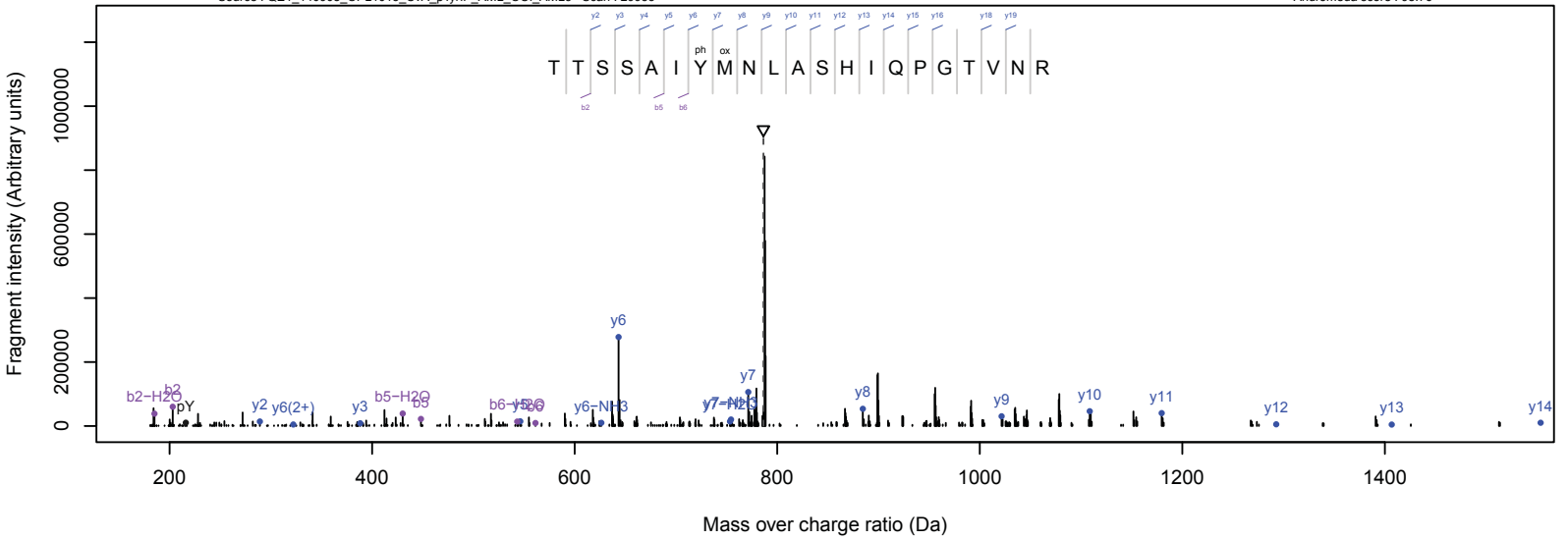
Andromeda score : 98.898



TTSSAIY(ph)M(ox)NLASHIQPGTVNR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 29953

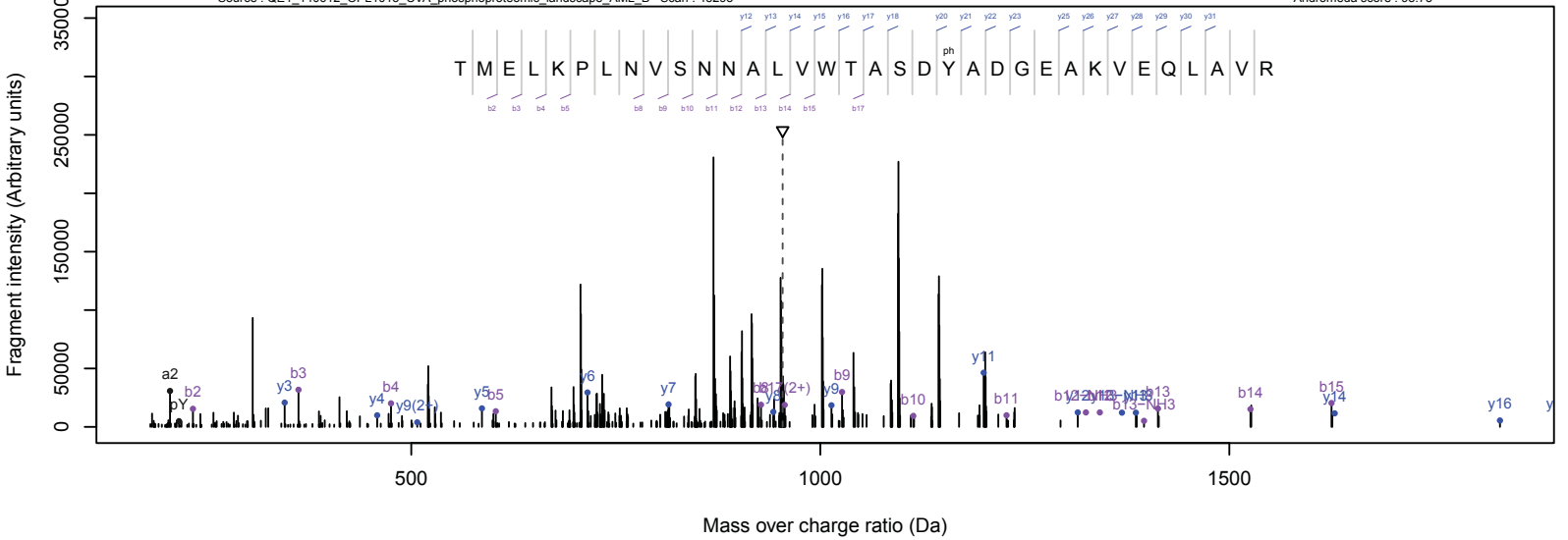
Andromeda score : 98.76



TMELKPLNVSNNALVWVTASDY(ph)ADGEAKVEQLAVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45295

Andromeda score : 98.75

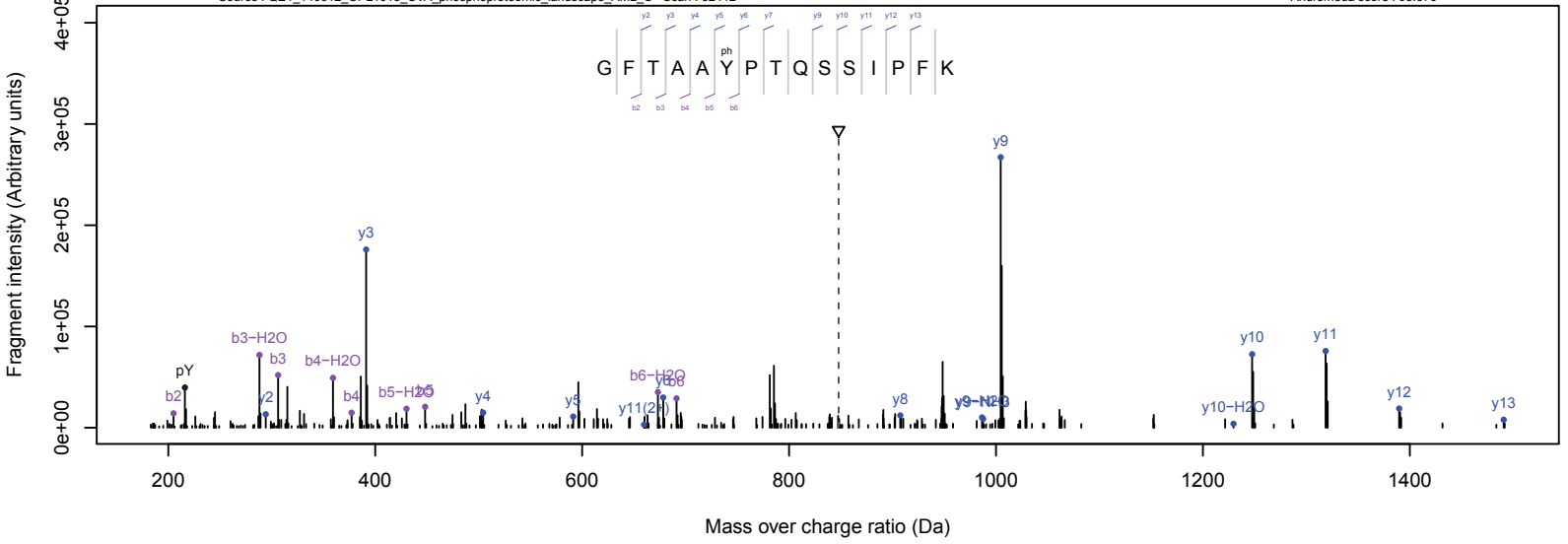




GFTAAY(ph)PTQSSIPFK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 32412

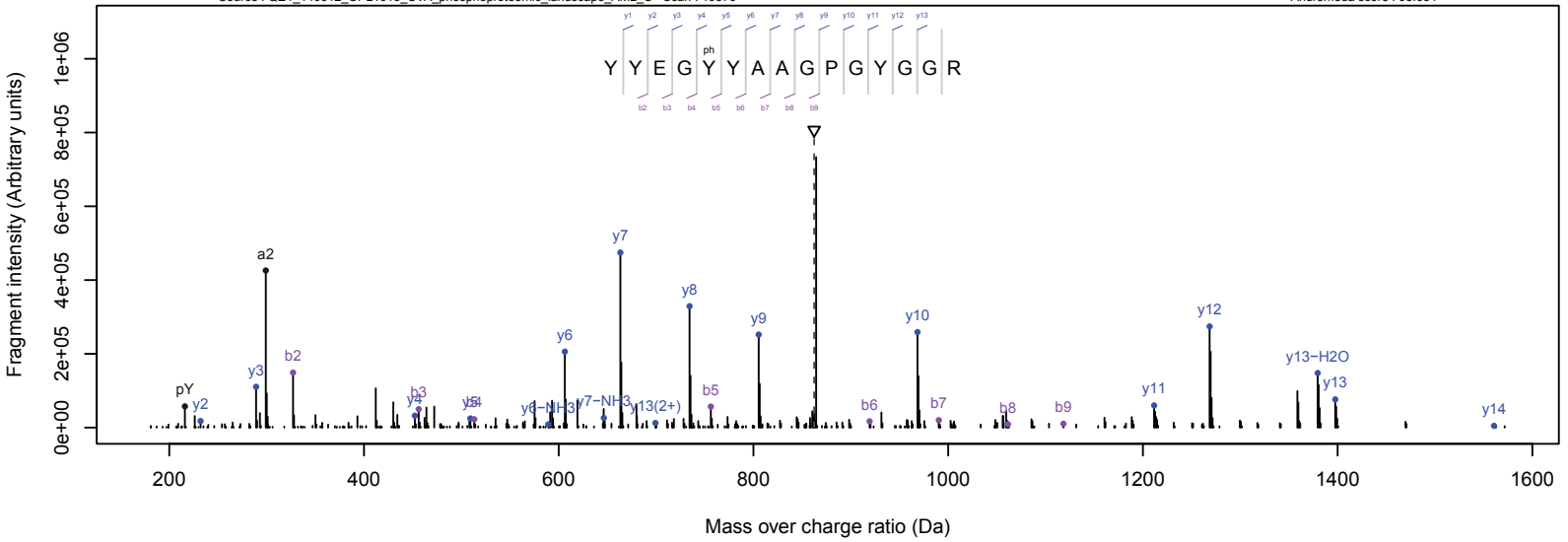
Andromeda score: 98.676



YYEGY(ph)YAAGPGYGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19376

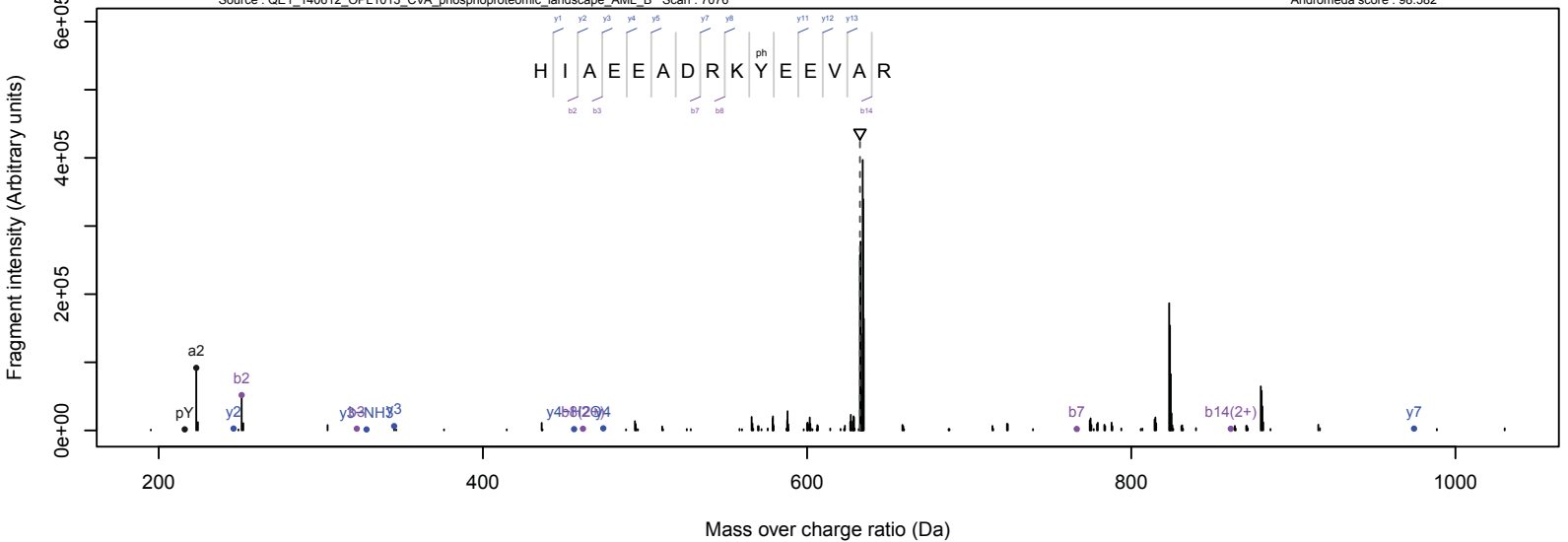
Andromeda score: 98.654



HIAEEADRKY(ph)EEVAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 7076

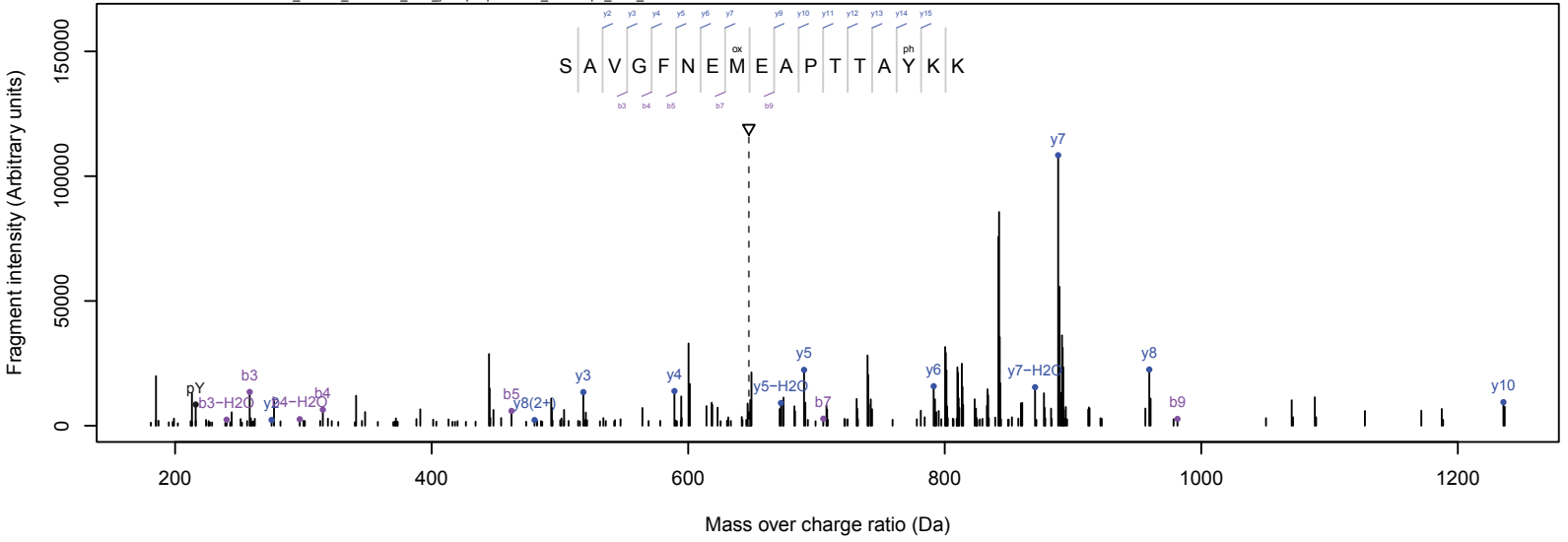
Andromeda score: 98.582



SAVGFNEM(ox)EAPTTAY(ph)KK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13909

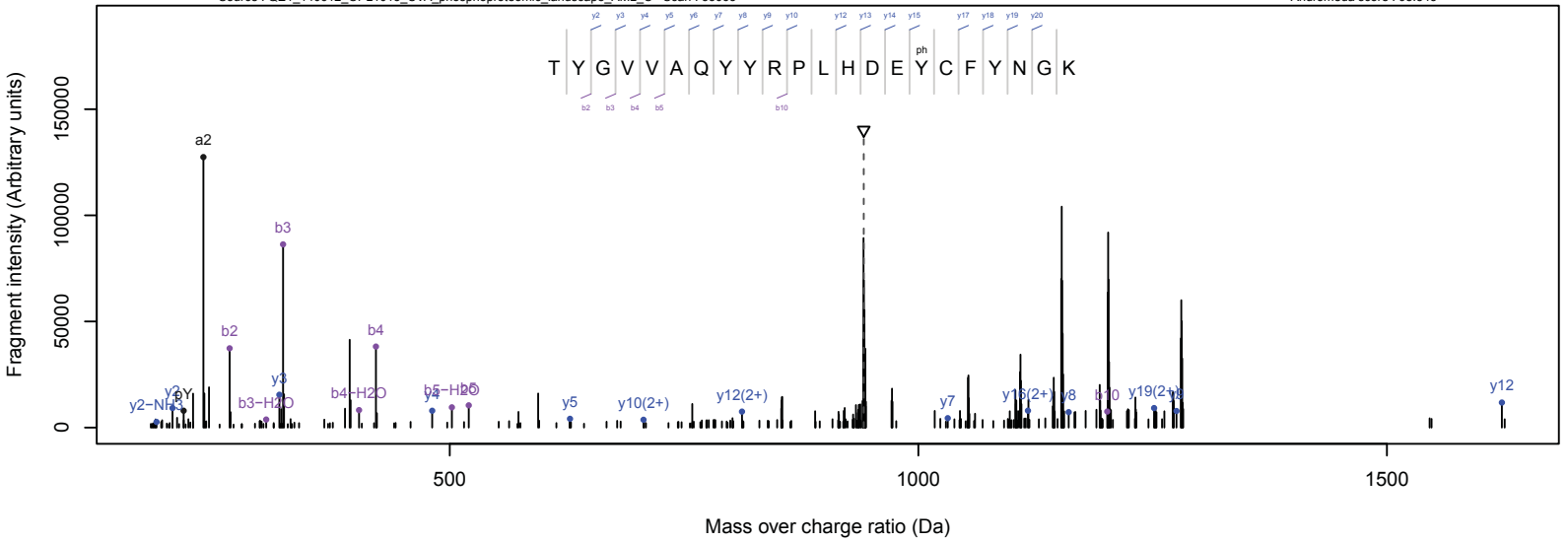
Andromeda score : 98.573



TYGVVAQYYRPLHDEY(ph)CFYNGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33063

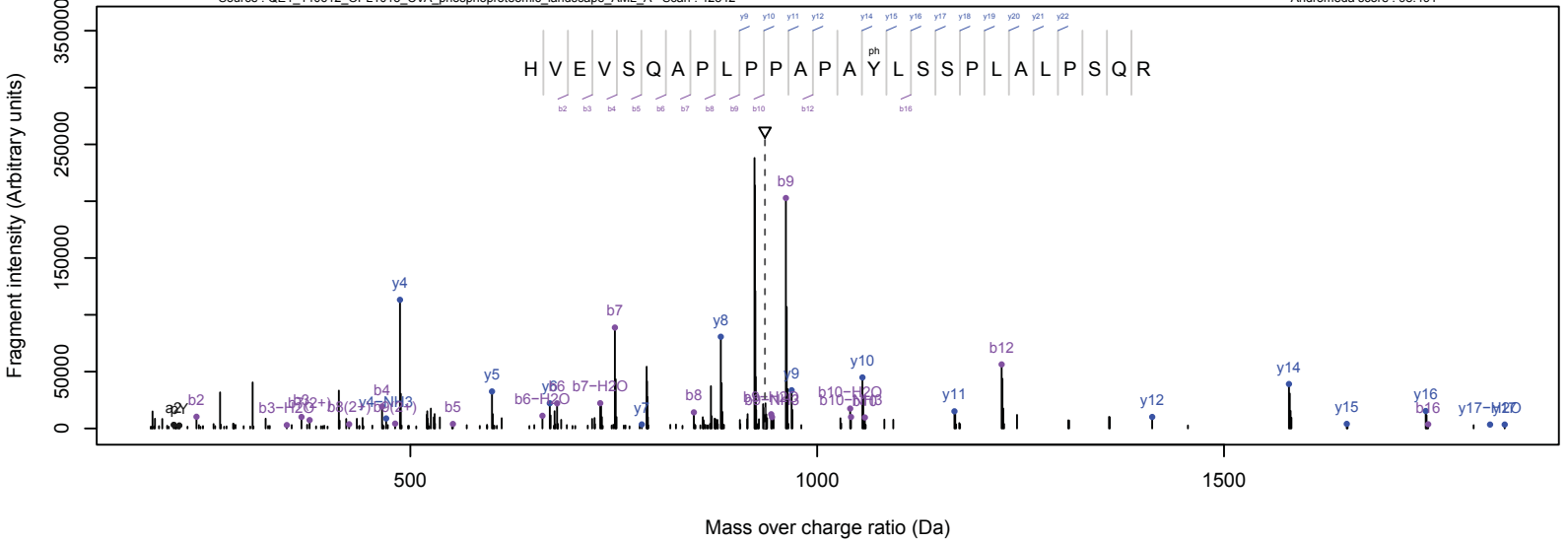
Andromeda score : 98.543



HVEVSQAPLPAPAY(ph)LSSPLALPSQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 42542

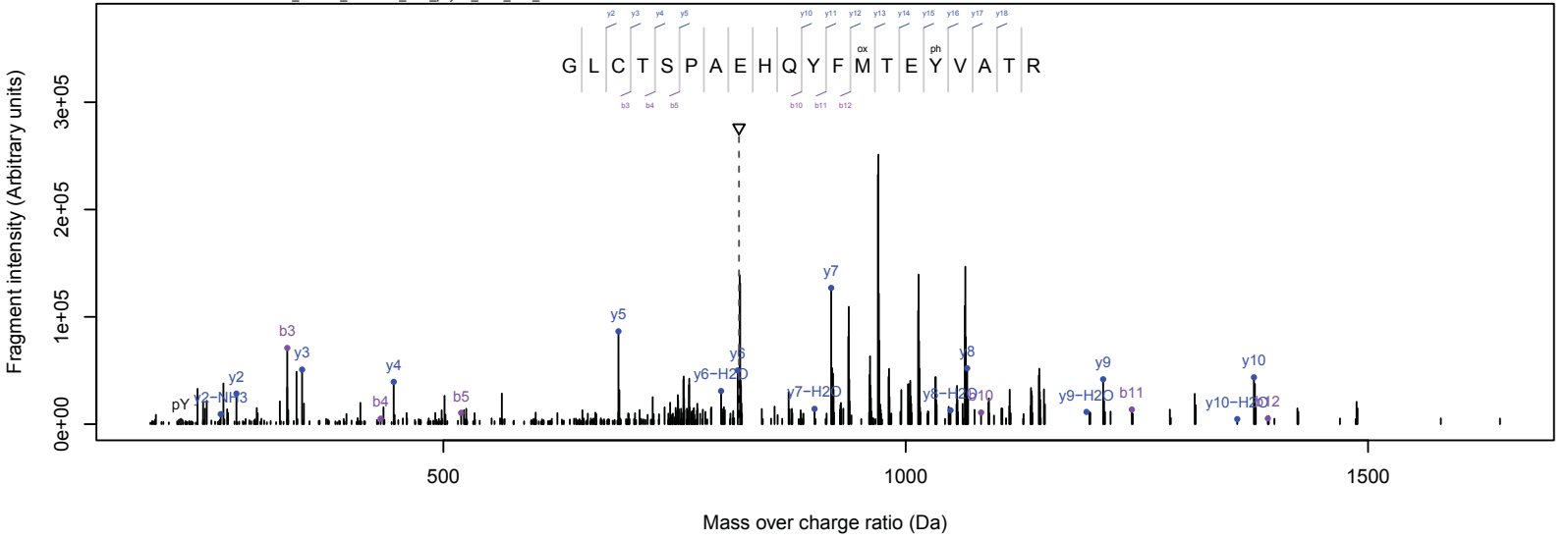
Andromeda score : 98.491



GLCTSPA<sup>ox</sup>EHQYFMTEYVATR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 28079

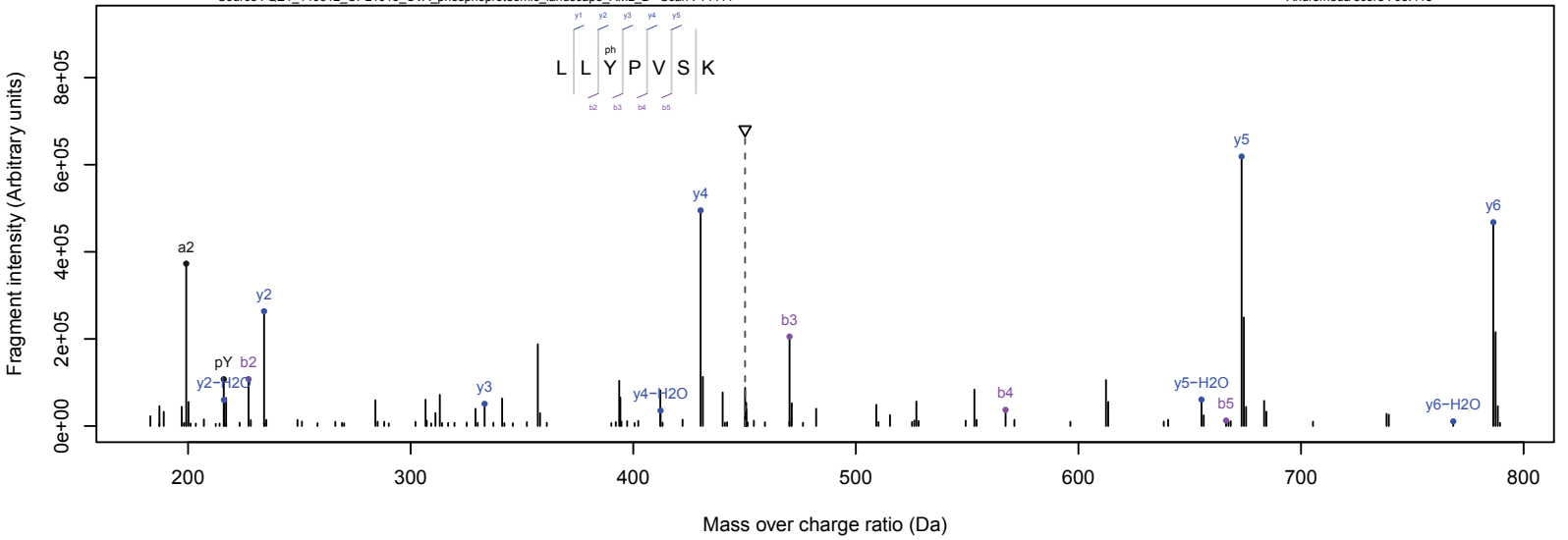
Andromeda score : 98.443



LLY(ph)PVS<sup>K</sup>-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14411

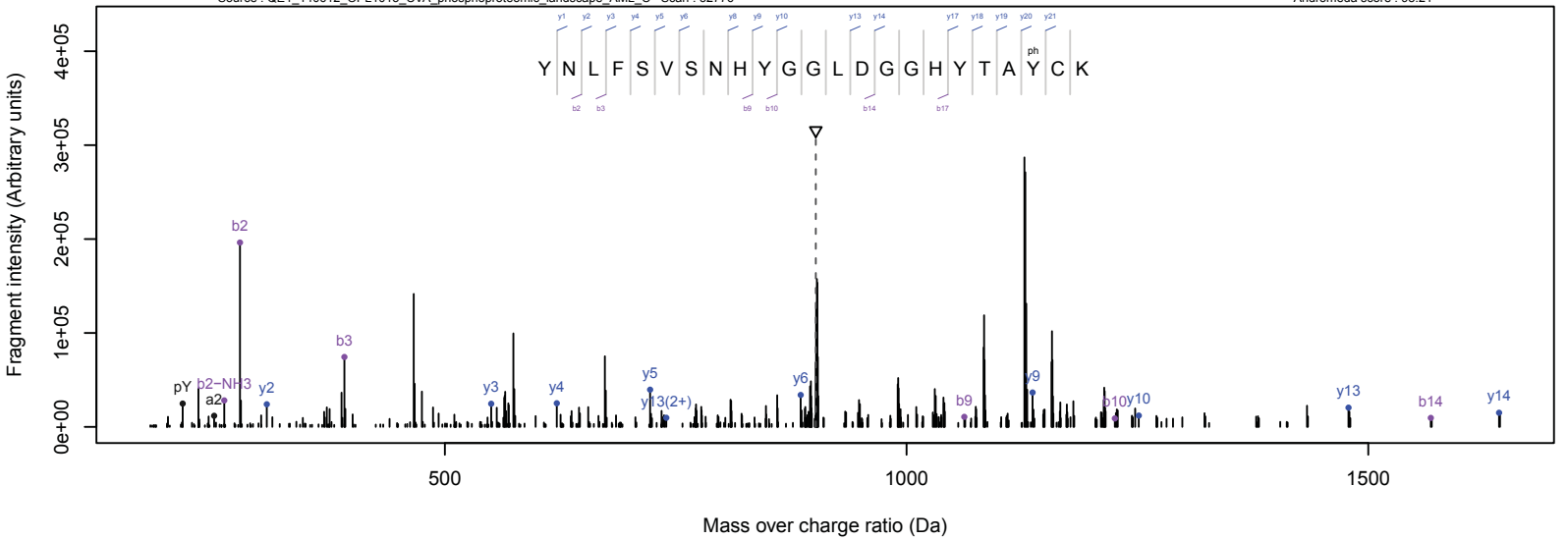
Andromeda score : 98.418



YNLFSVSNHYGGLDGGHYTAY<sup>ph</sup>CK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32776

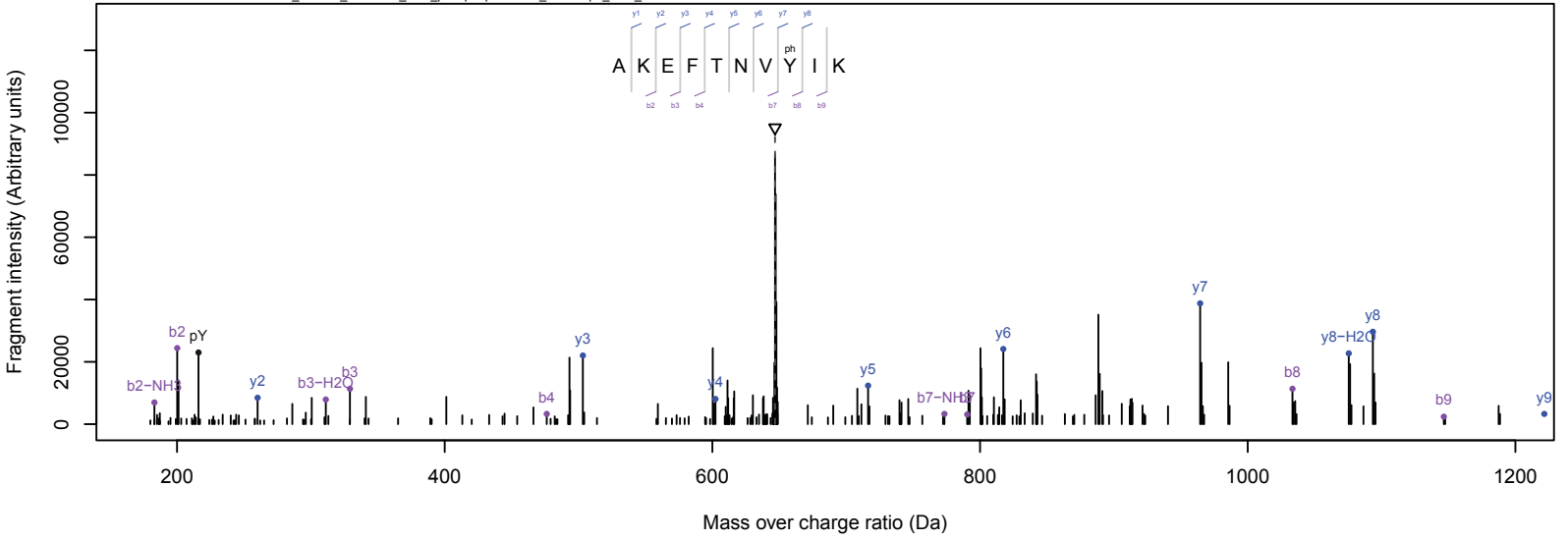
Andromeda score : 98.21



**AKEFTNVY(ph)IK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17457

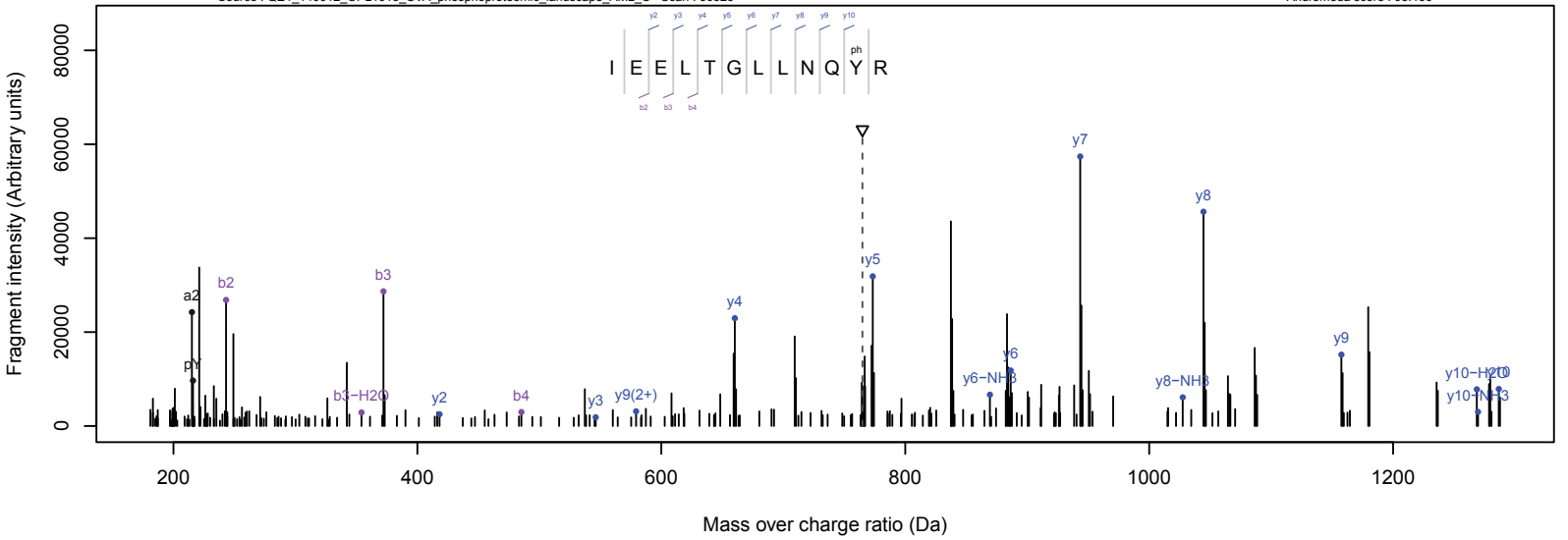
Andromeda score : 98.156



**IEELTGLLNQY(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36523

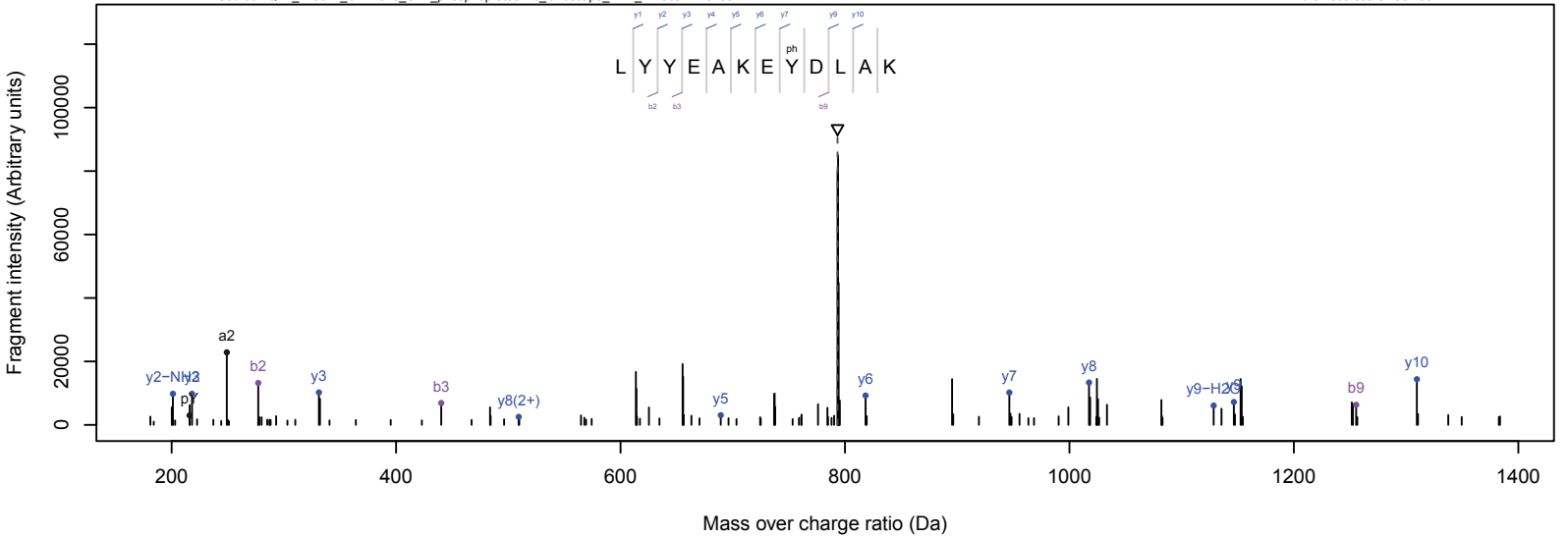
Andromeda score : 98.156



**LYYEAKY(ph)DLAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18155

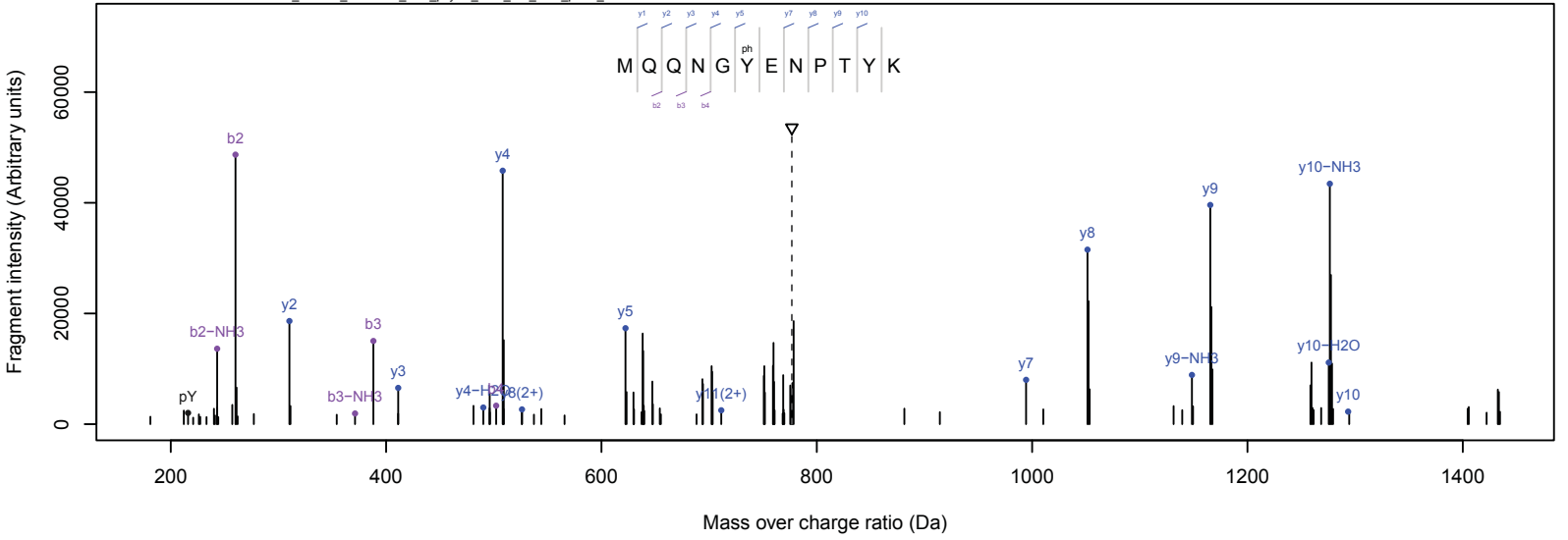
Andromeda score : 98.105



**MQQNGY(ph)ENPTYK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 4489

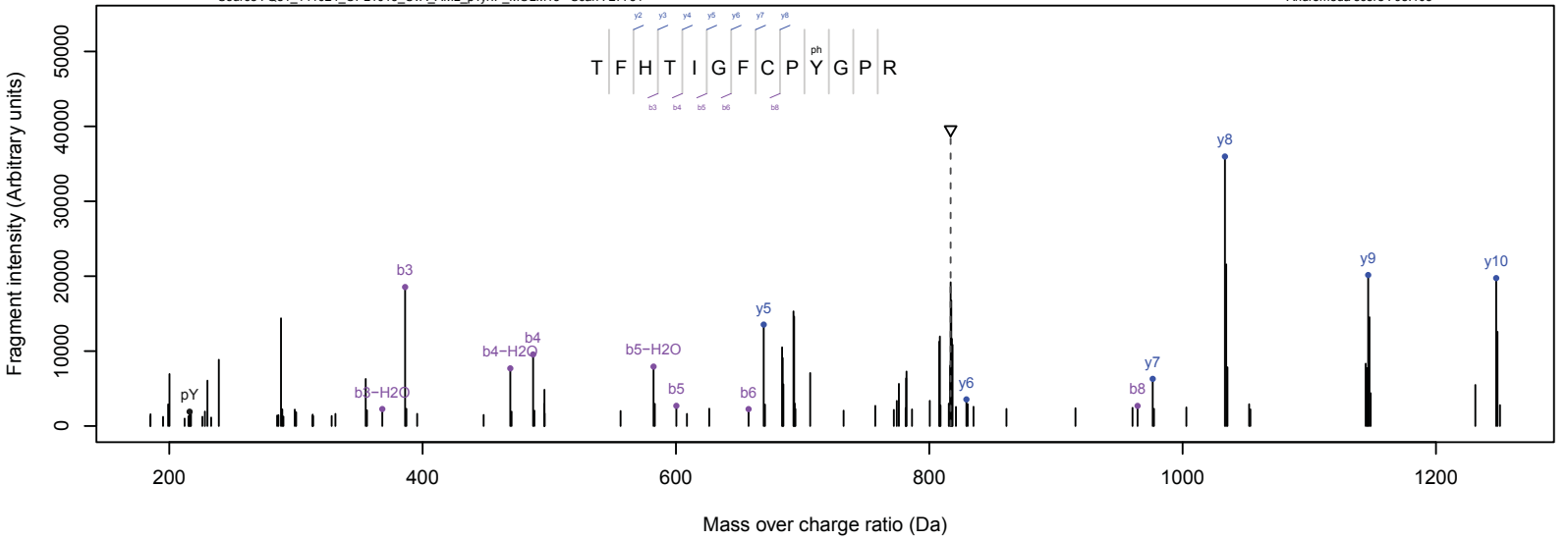
Andromeda score : 98.105



**TFHTIGFCPY(ph)GPR-2+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 27751

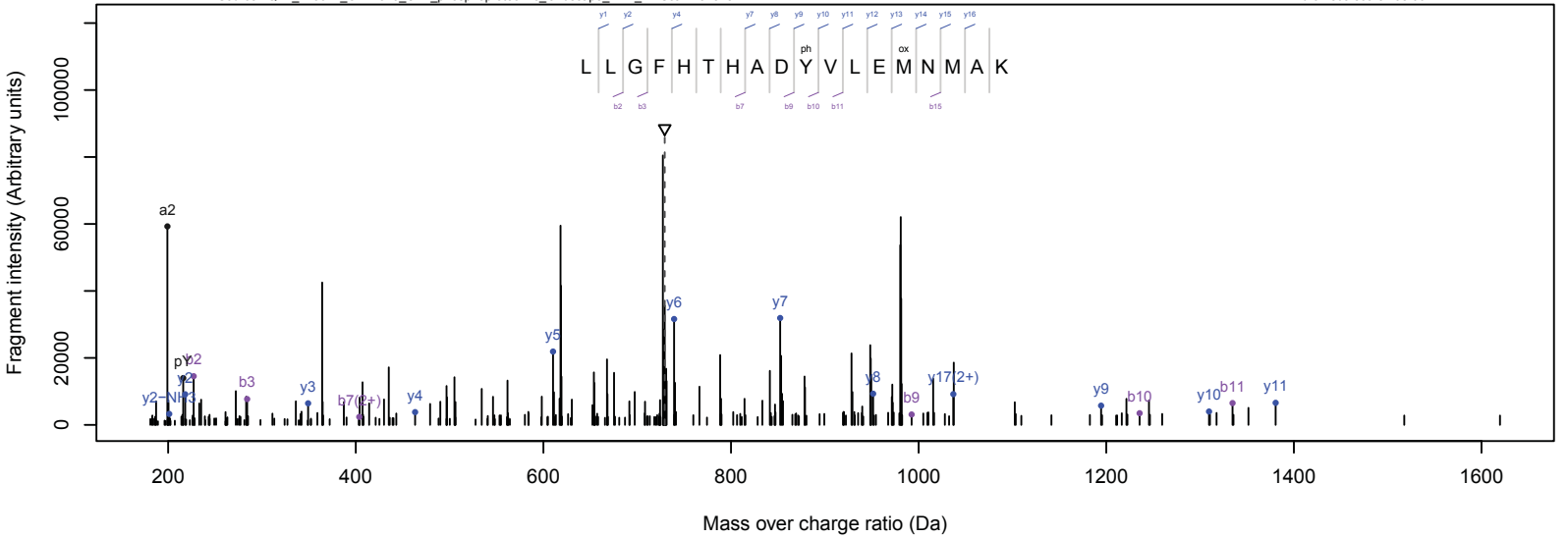
Andromeda score : 98.105



**LLGFHTHADY(ph)VLEMNMAK-3+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan : 34828

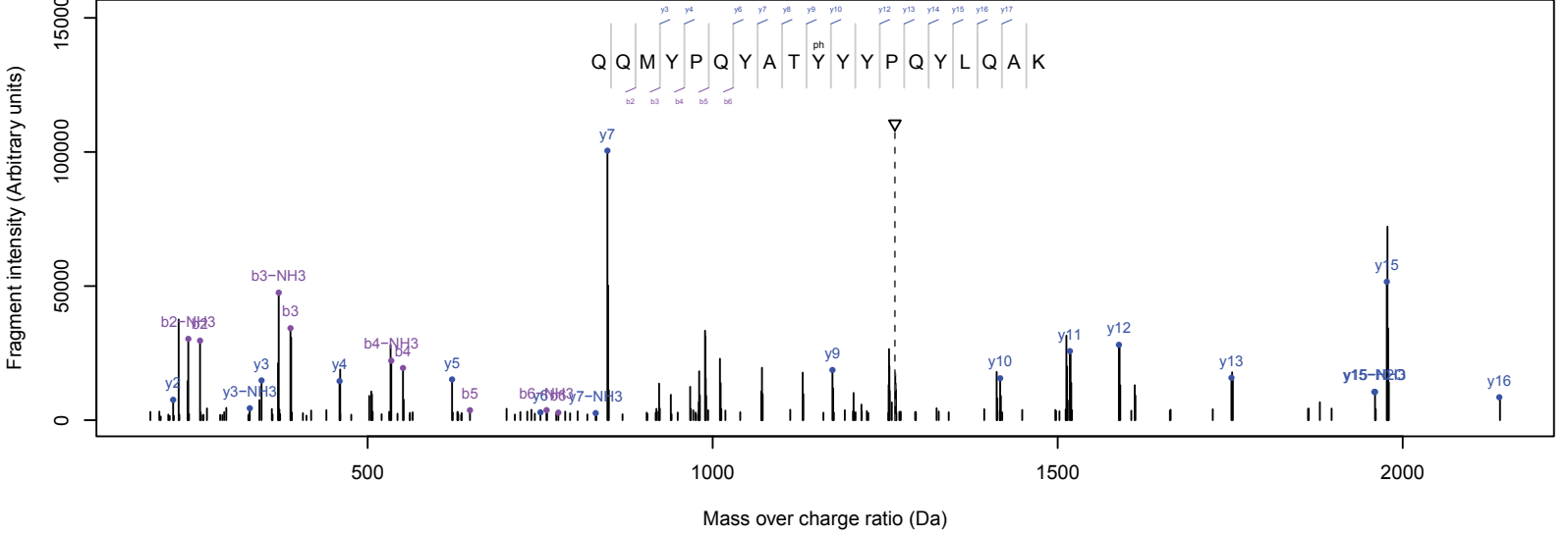
Andromeda score : 98.062



QQMYPQYATYYPQLQAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39249

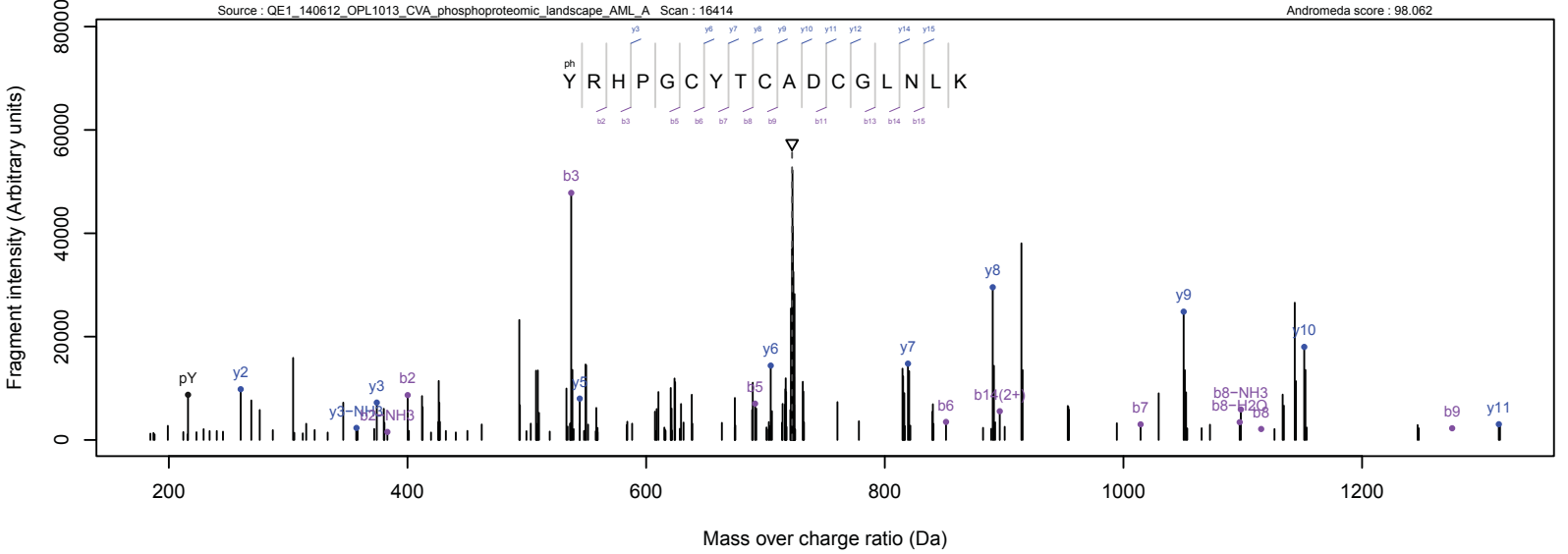
Andromeda score : 98.062



Y(pH)RHPGICYTCADCGLNLIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16414

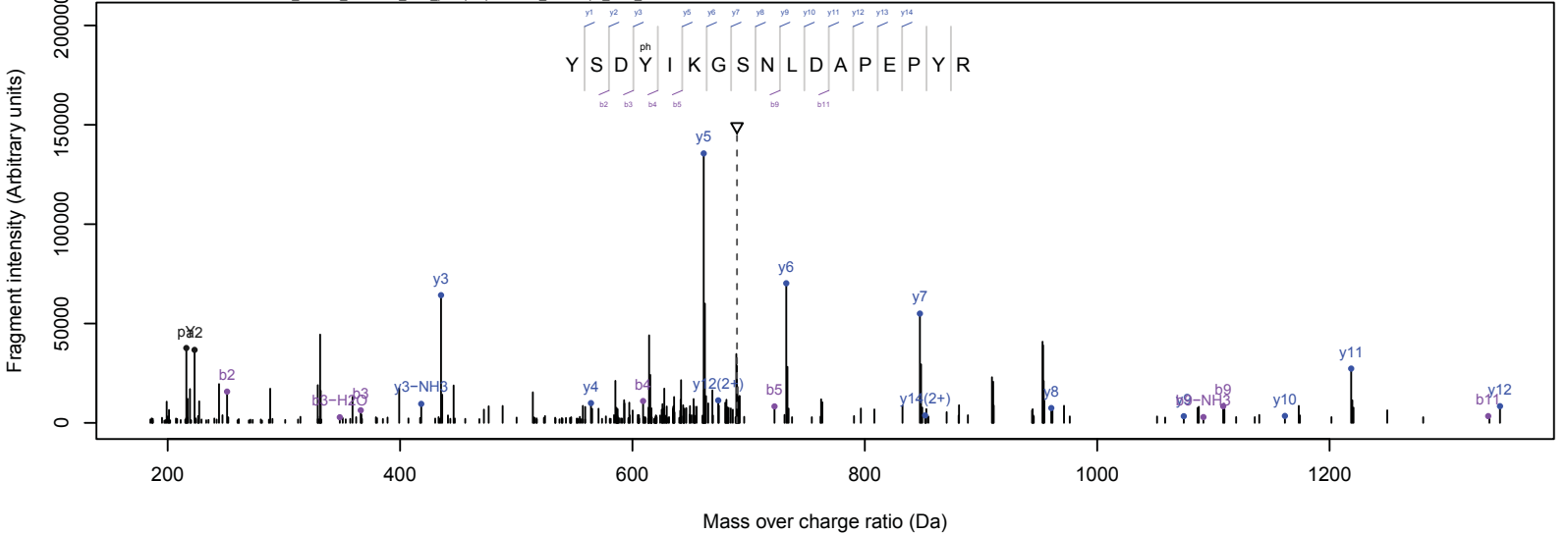
Andromeda score : 98.062



YSDY(pH)KGSNLDAPEPYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22781

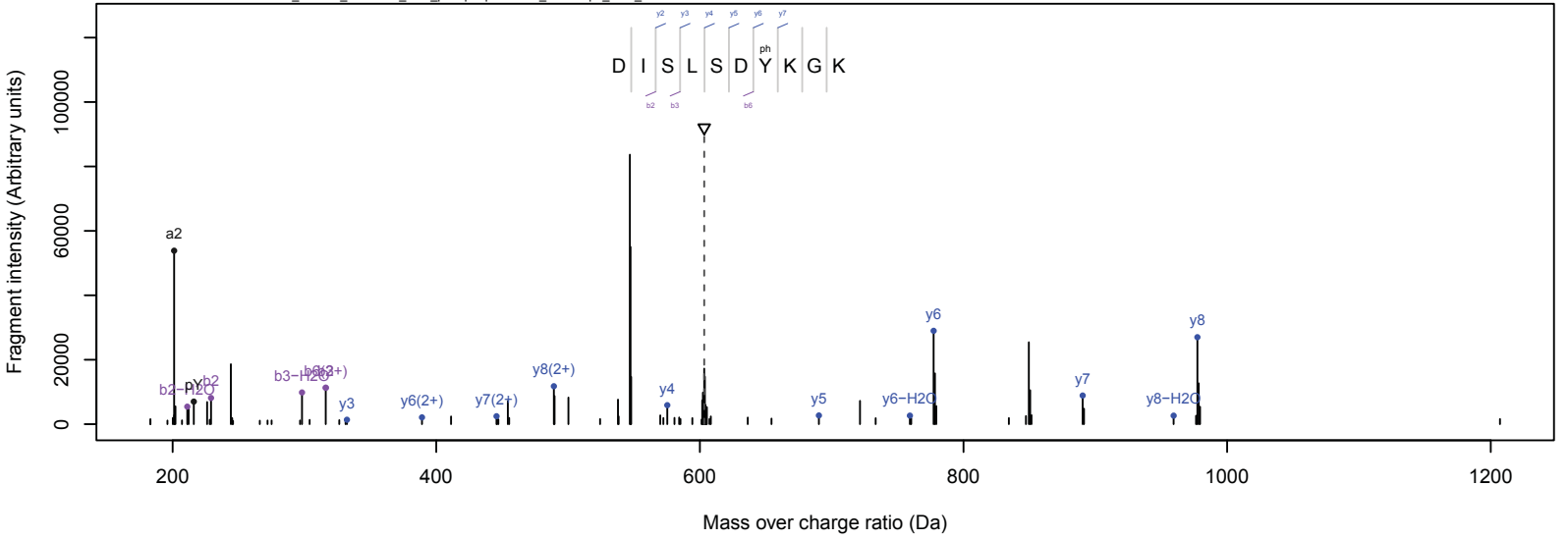
Andromeda score : 98.062



DISLSDY(ph)KGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10253

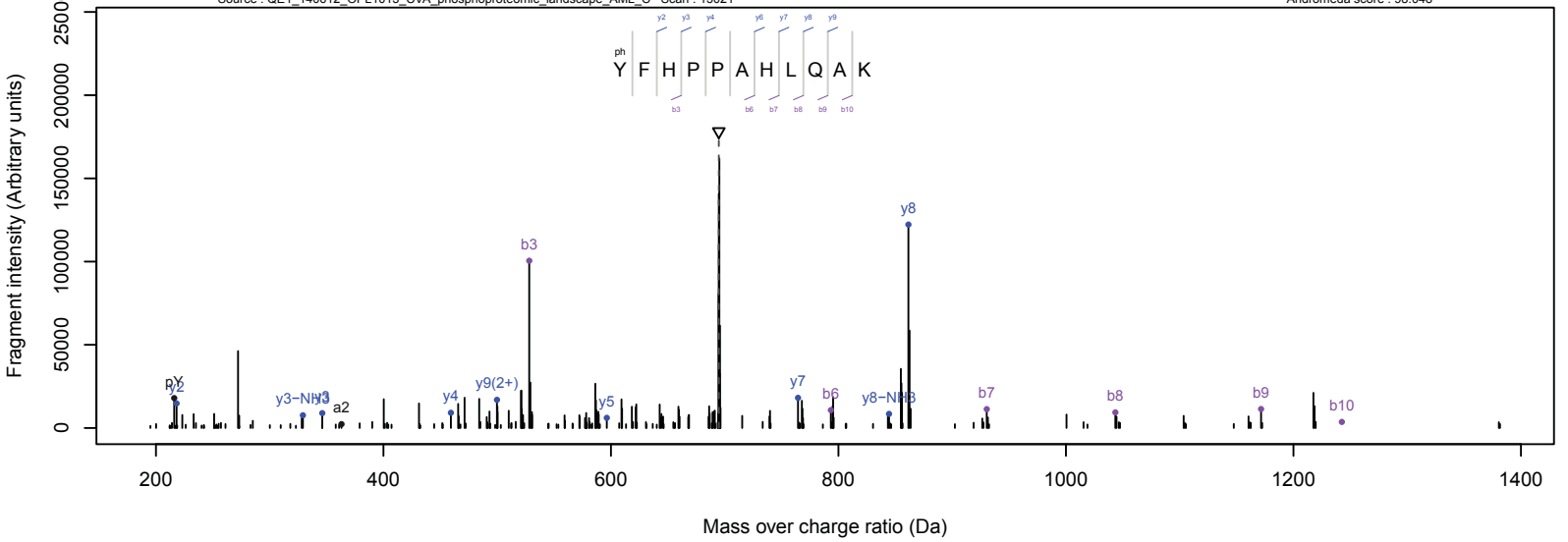
Andromeda score : 98.048



Y(ph)FHPPAHLQAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13021

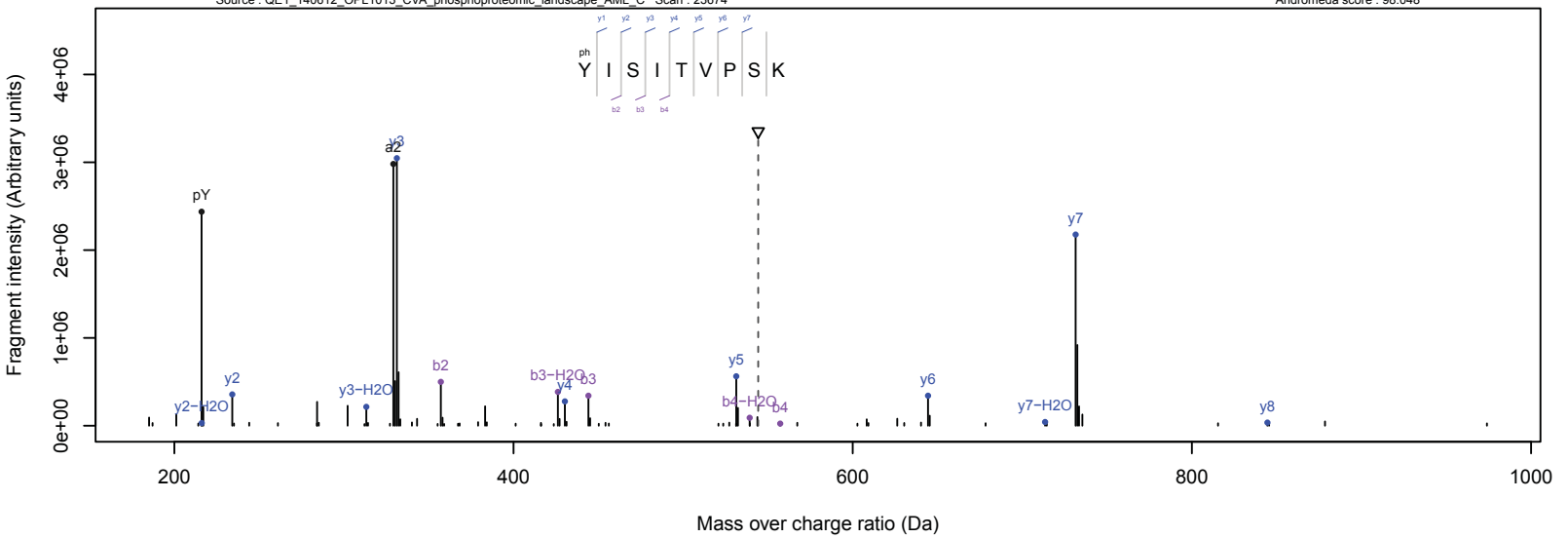
Andromeda score : 98.048



Y(ph)ISITVPSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23674

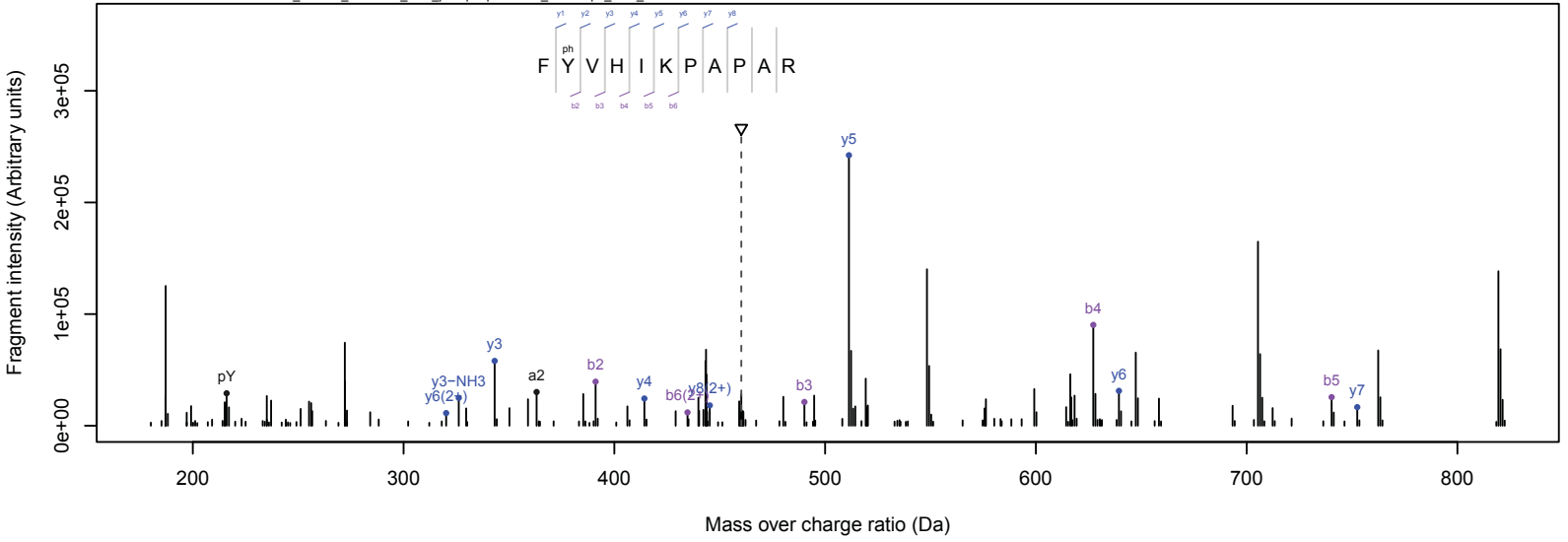
Andromeda score : 98.048



FY(ph)VHIKPAPAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17748

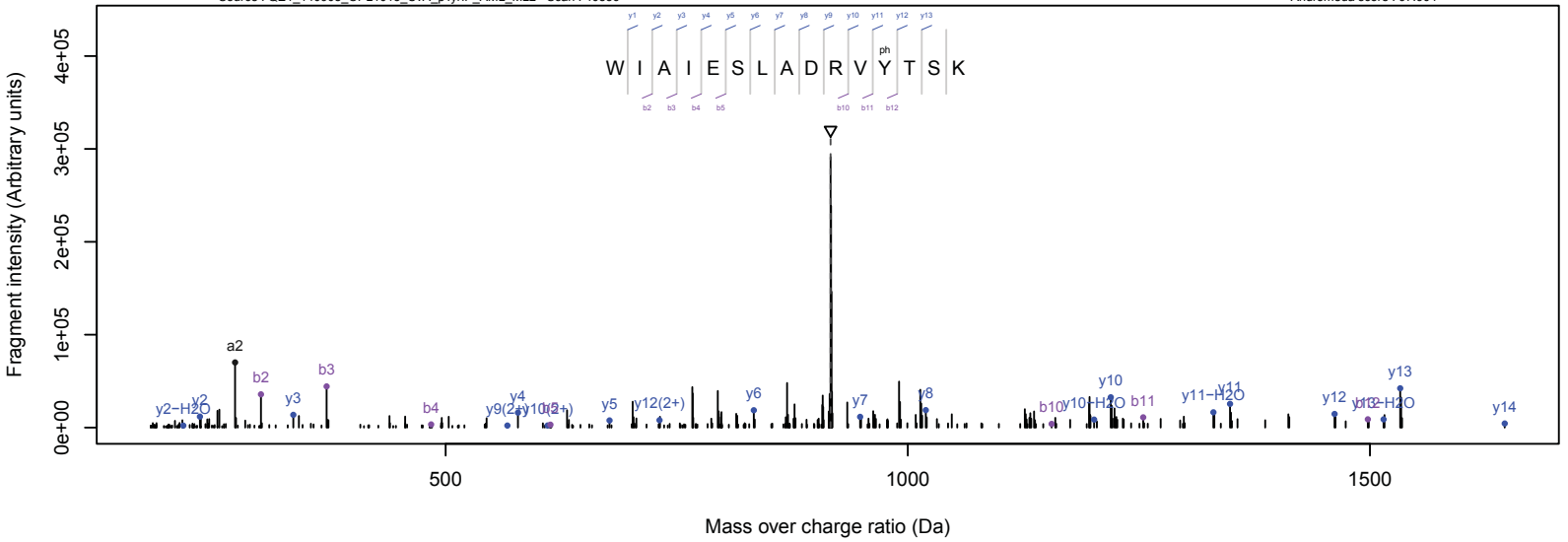
Andromeda score : 98.044



WIAIESLADR VY(ph)TSK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 40836

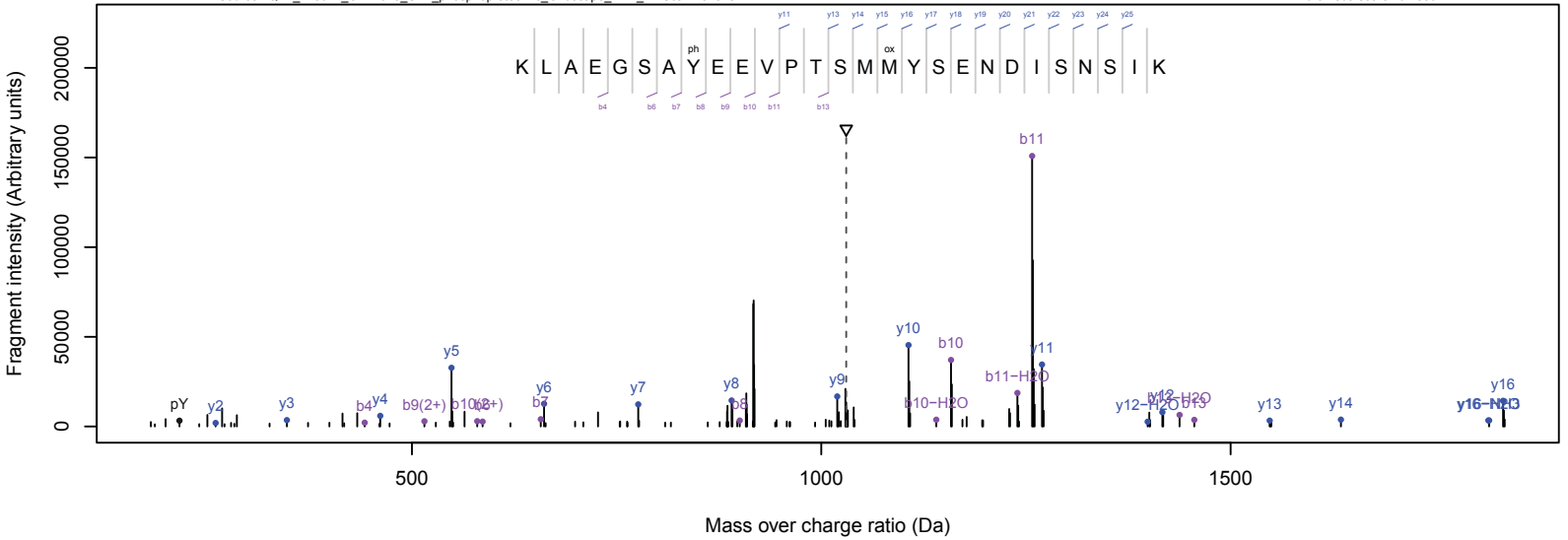
Andromeda score : 97.904



KLAEGSAYE E V P T S M M Y S E N D I S N S I K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34375

Andromeda score : 97.839

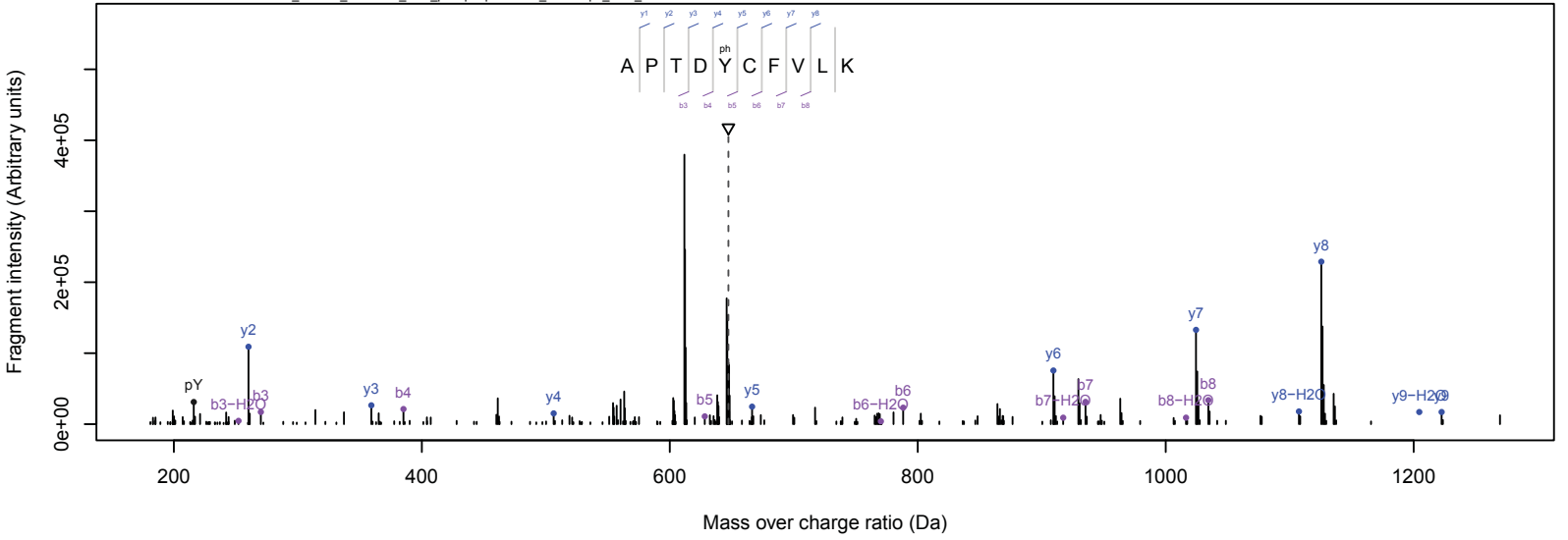




APT<sup>ph</sup>DYCFVLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28773

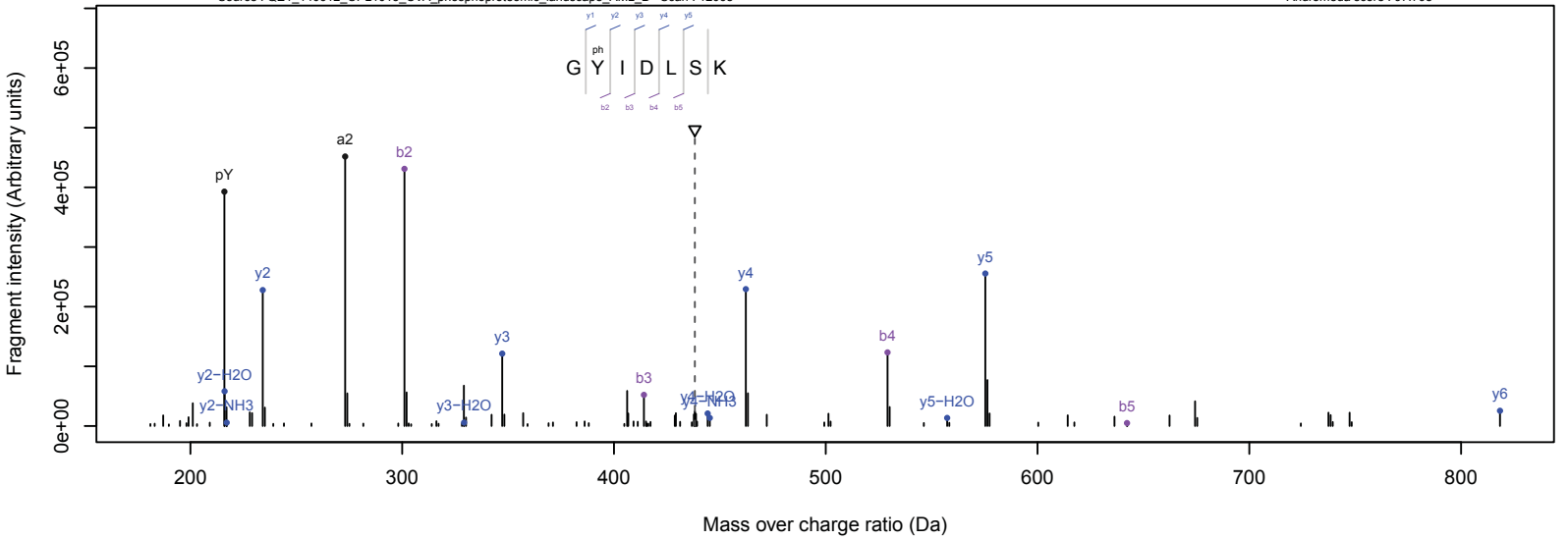
Andromeda score : 97.813



GY<sup>ph</sup>IDLSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12068

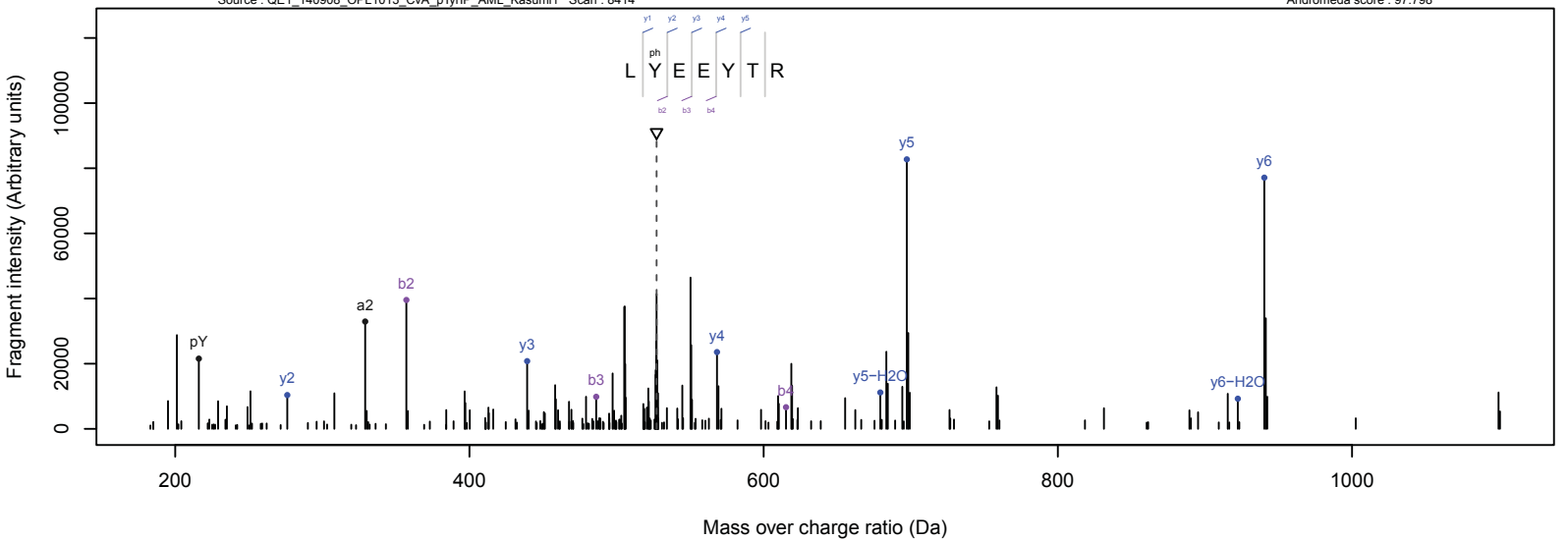
Andromeda score : 97.798



LY<sup>ph</sup>EEYTR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 8414

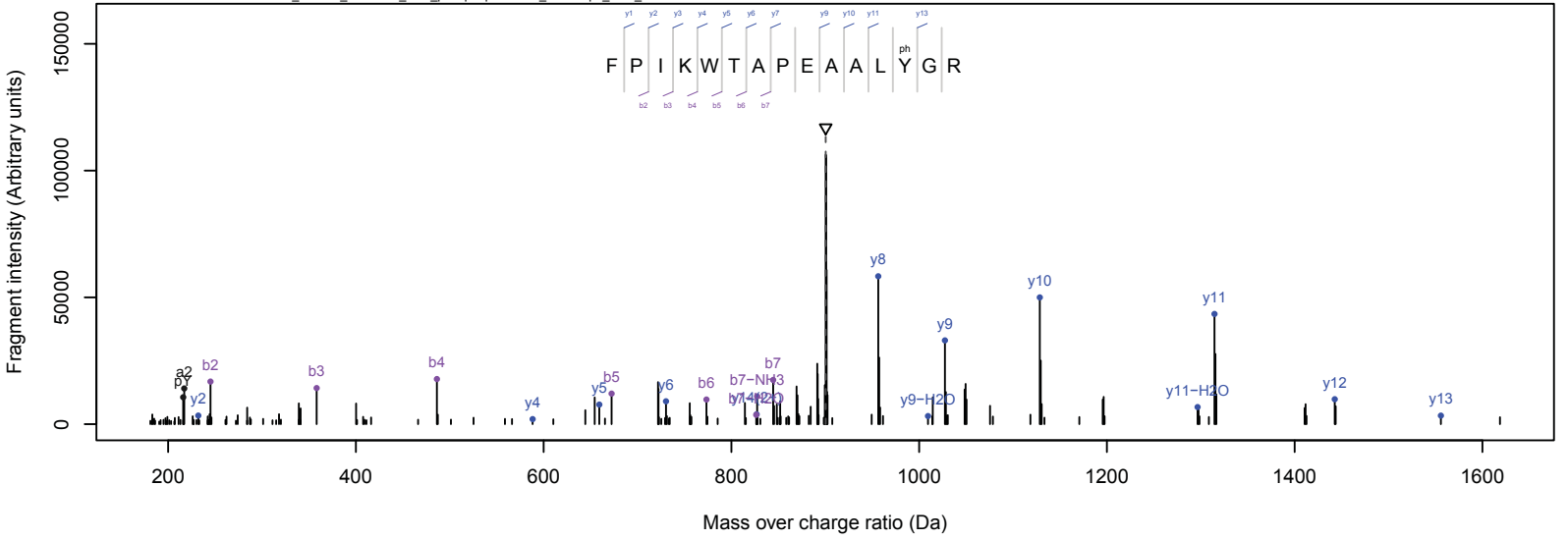
Andromeda score : 97.798



FPIKWTAEALYGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38316

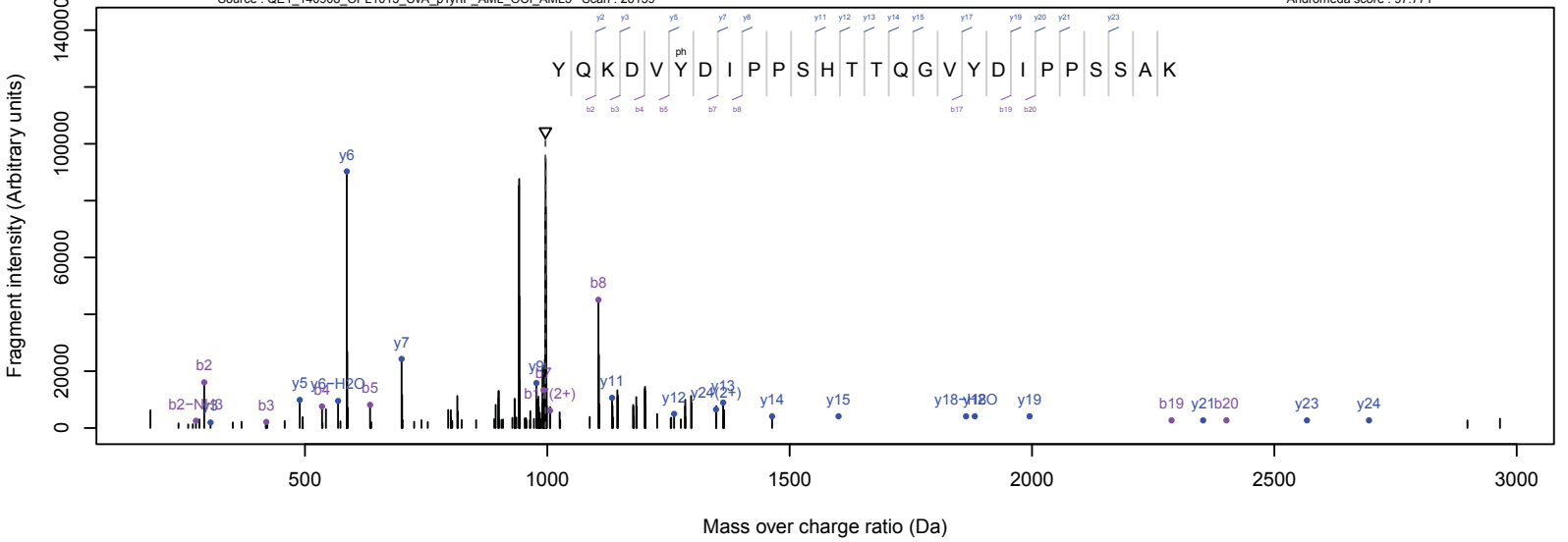
Andromeda score: 97.797



YQKDVY(ph)DIPPSHTTQGVYDIPSSAK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 26159

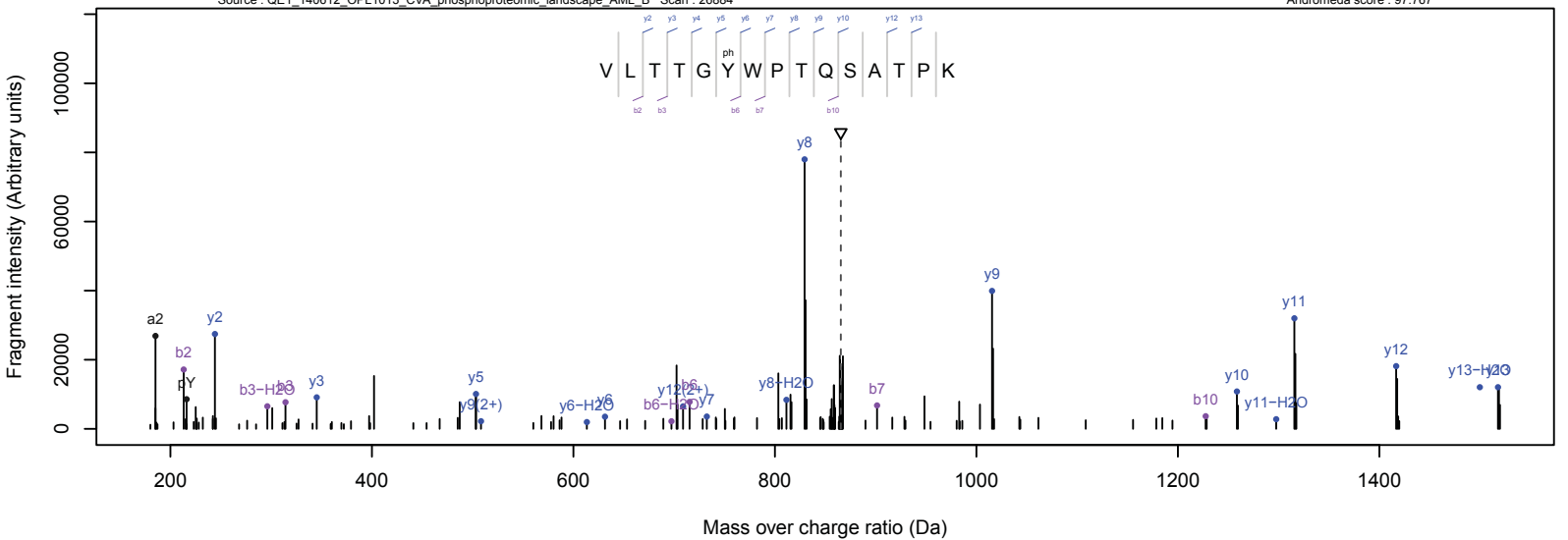
Andromeda score: 97.771



VLTTGYWPTQSATPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 26884

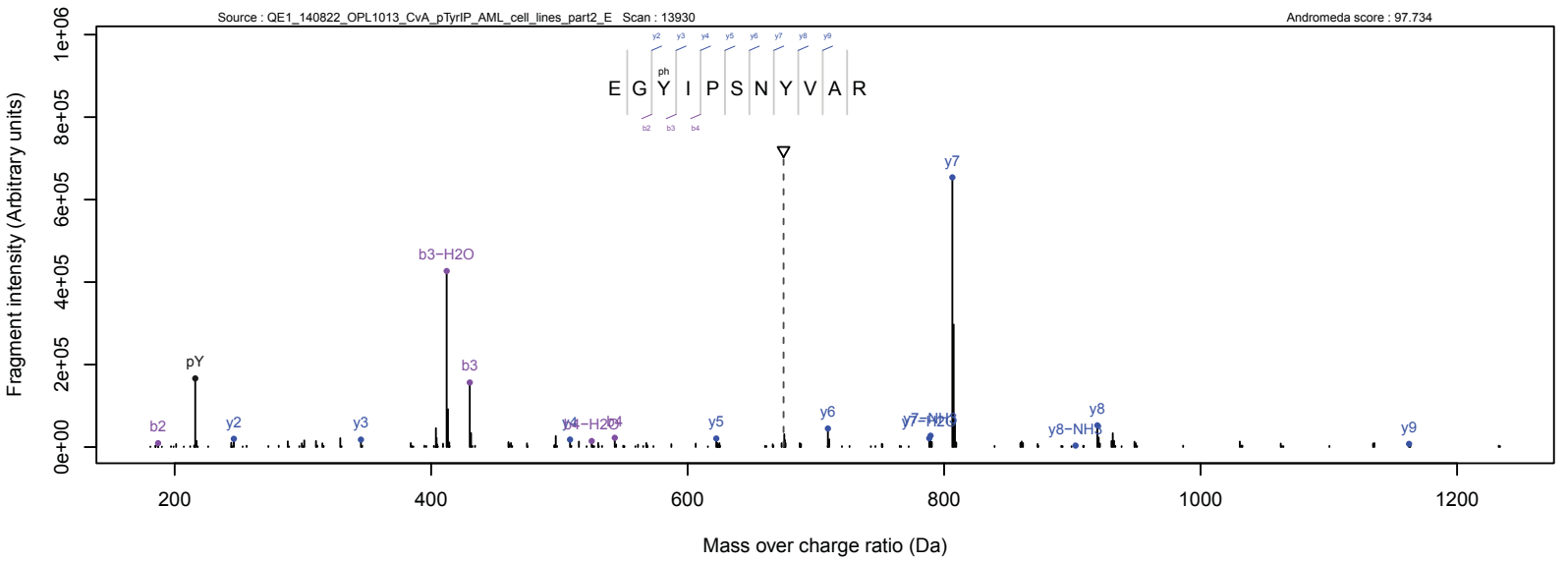
Andromeda score: 97.767



EGY(ph)IPSNYVAR-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 13930

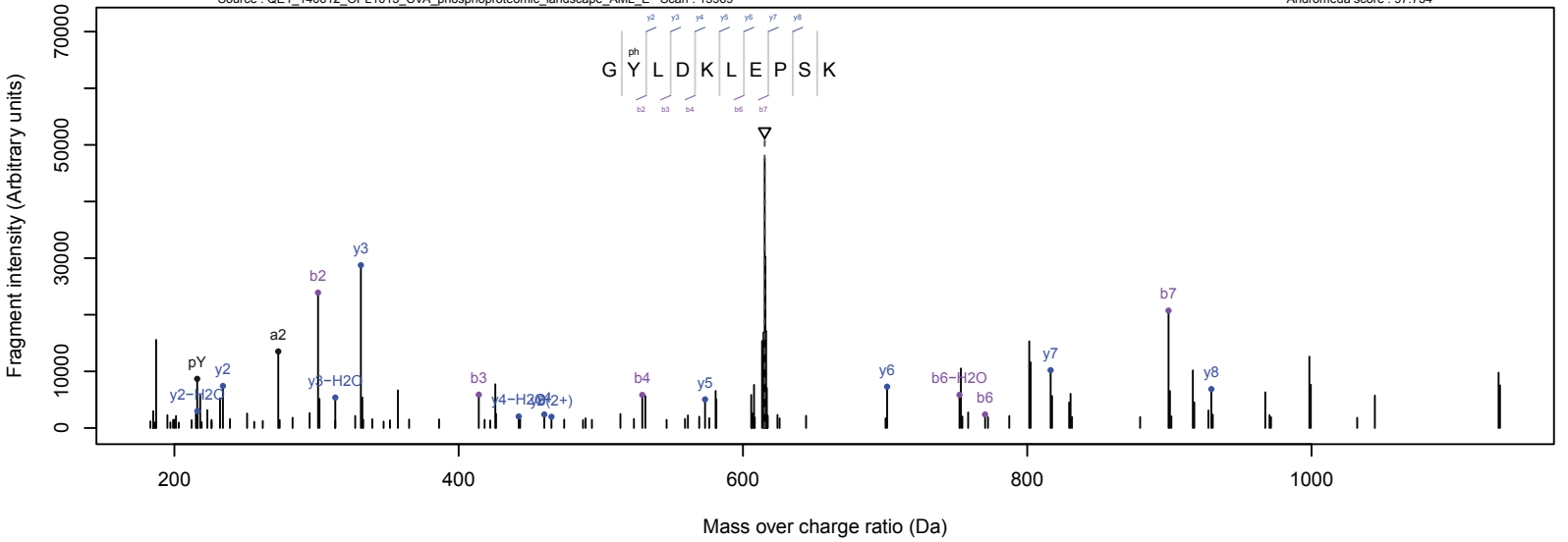
Andromeda score : 97.734



GY(ph)LDKLEPSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 15969

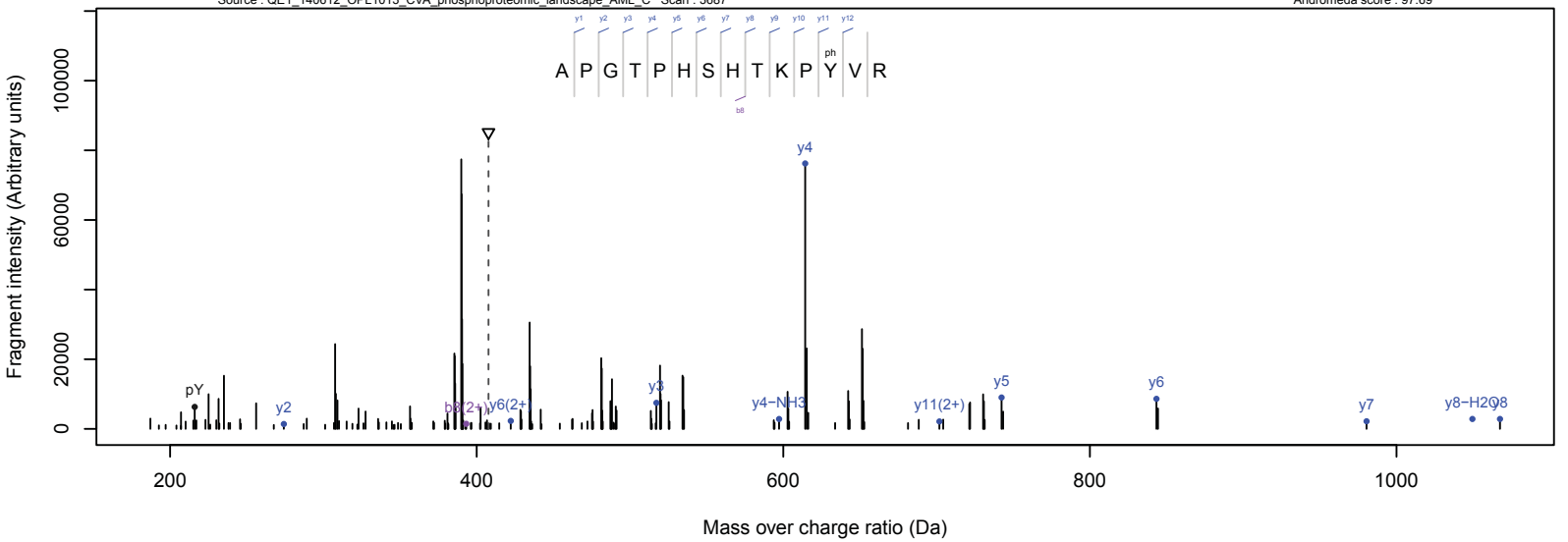
Andromeda score : 97.734



APGTPHSHTKPYVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3687

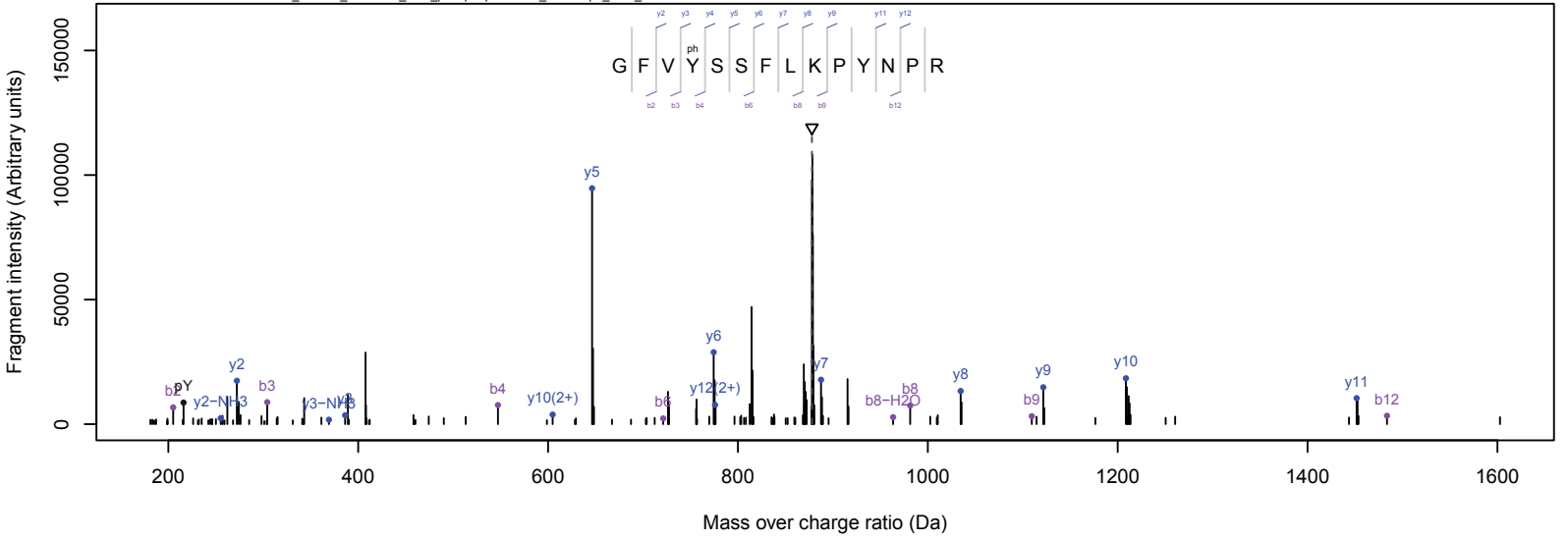
Andromeda score : 97.69



GFVY(ph)SSFLKPYNPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35555

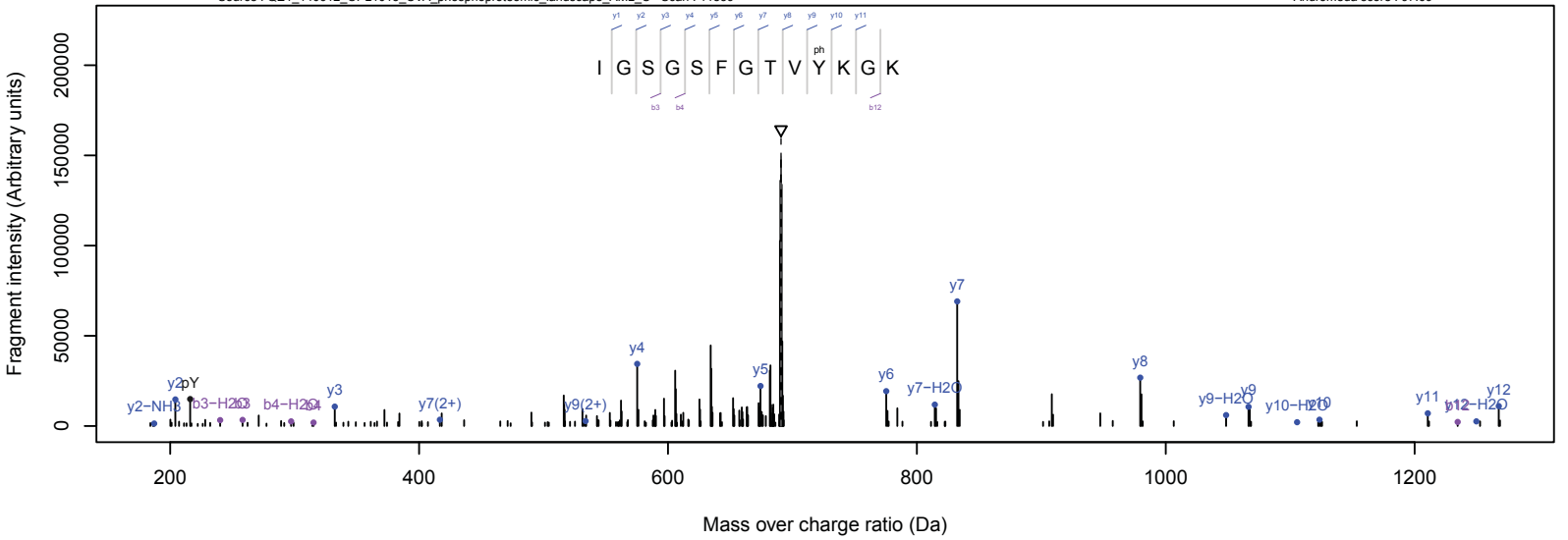
Andromeda score : 97.69



IGSGSFGTVY(ph)KGGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11839

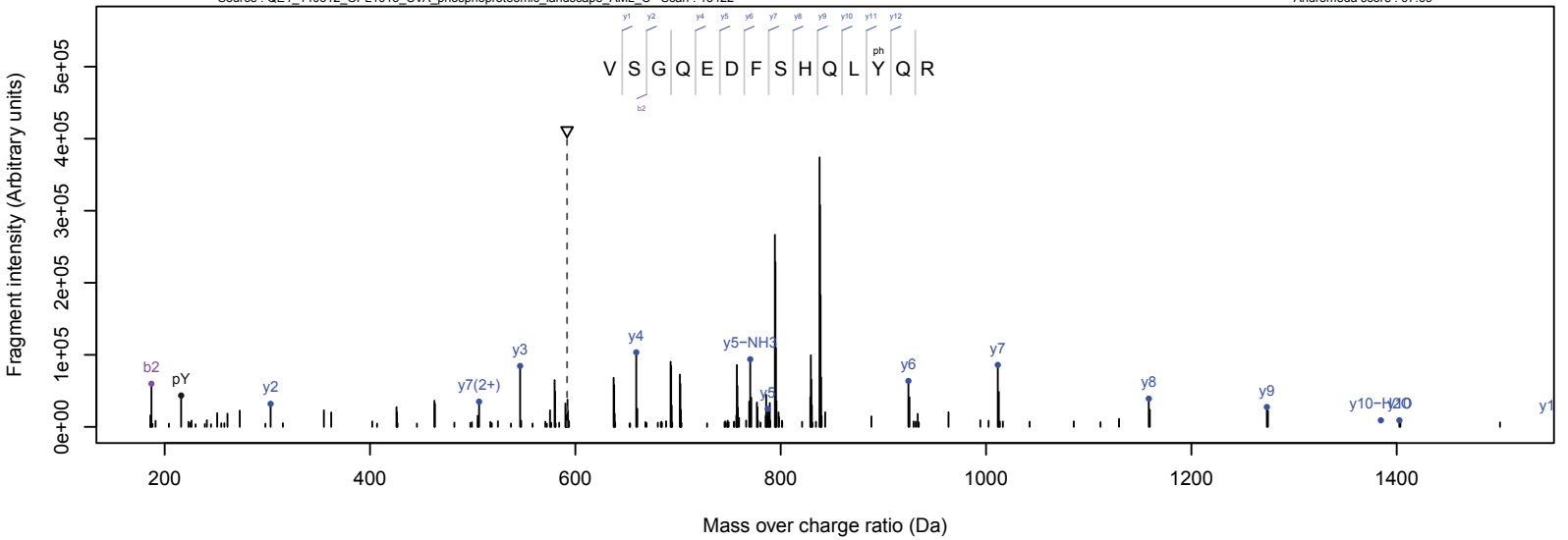
Andromeda score : 97.69



VSGQEDFSHQLY(ph)QR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15422

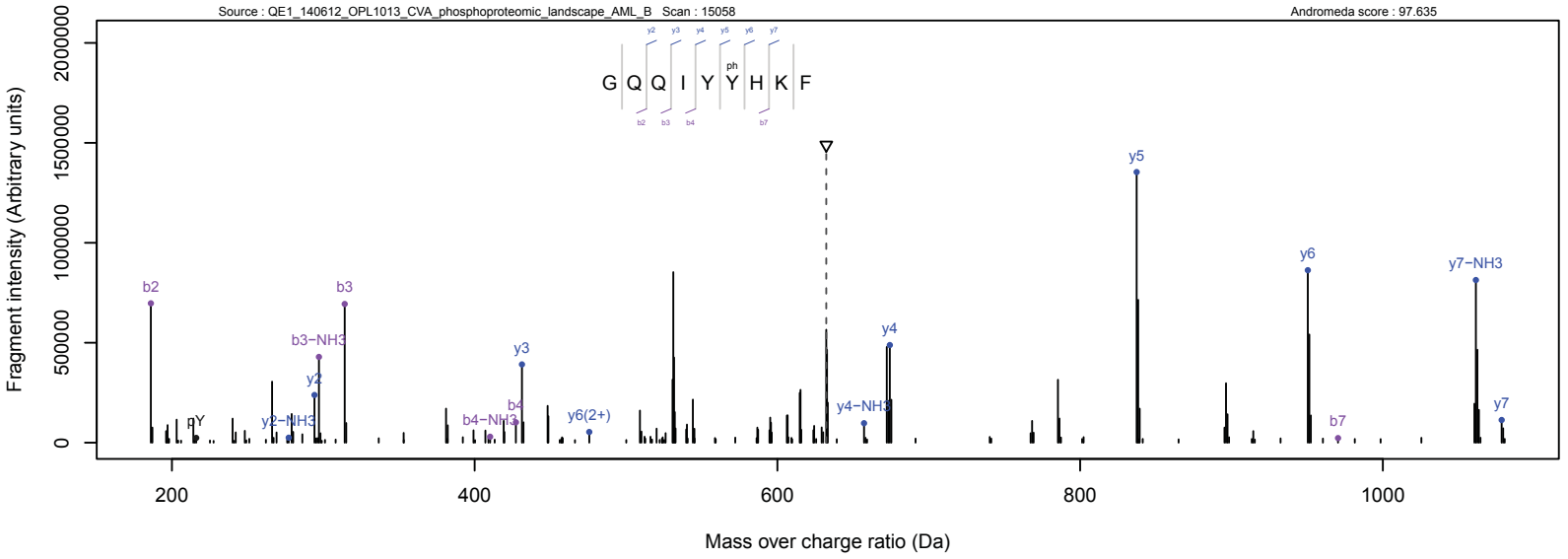
Andromeda score : 97.69



GQIQY(ph)HKF-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15058

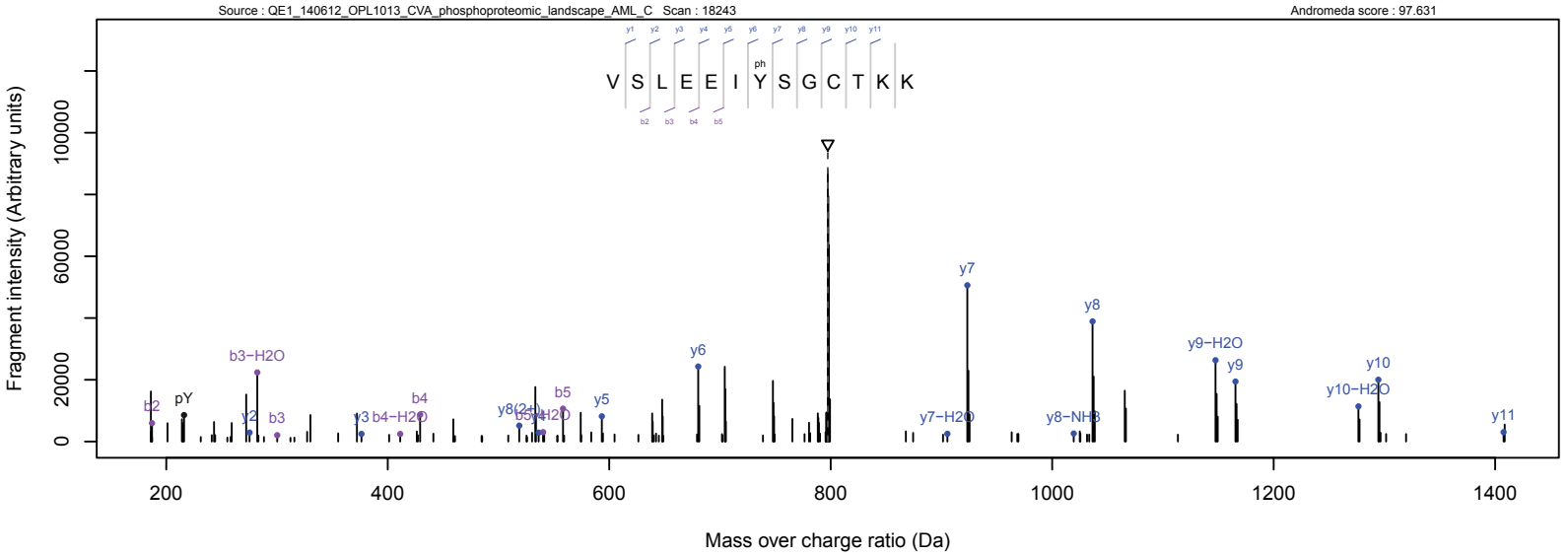
Andromeda score : 97.635



VSLEEIY(ph)SGCTKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18243

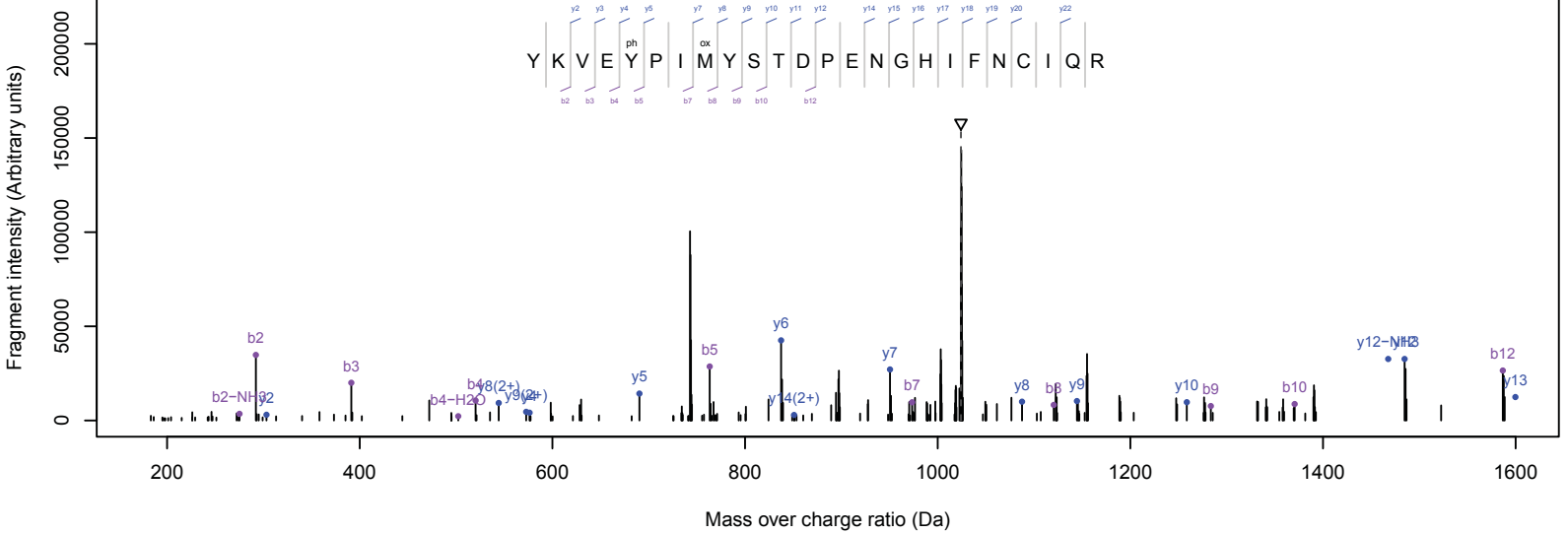
Andromeda score : 97.631



YKVEEY(ph)PIM(ox)YSTDPENGHIFNCIQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 34347

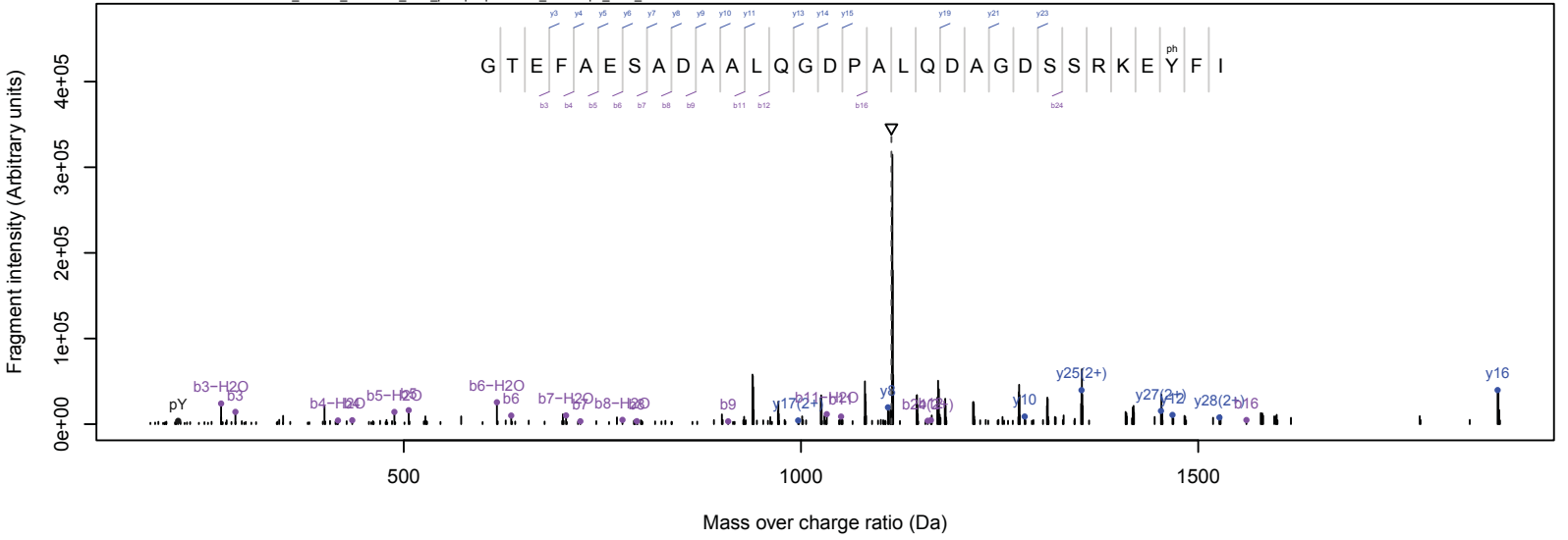
Andromeda score : 97.542



GTEFAESADAALQGDPALQDAGDSSRKEY(ph)FI-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41692

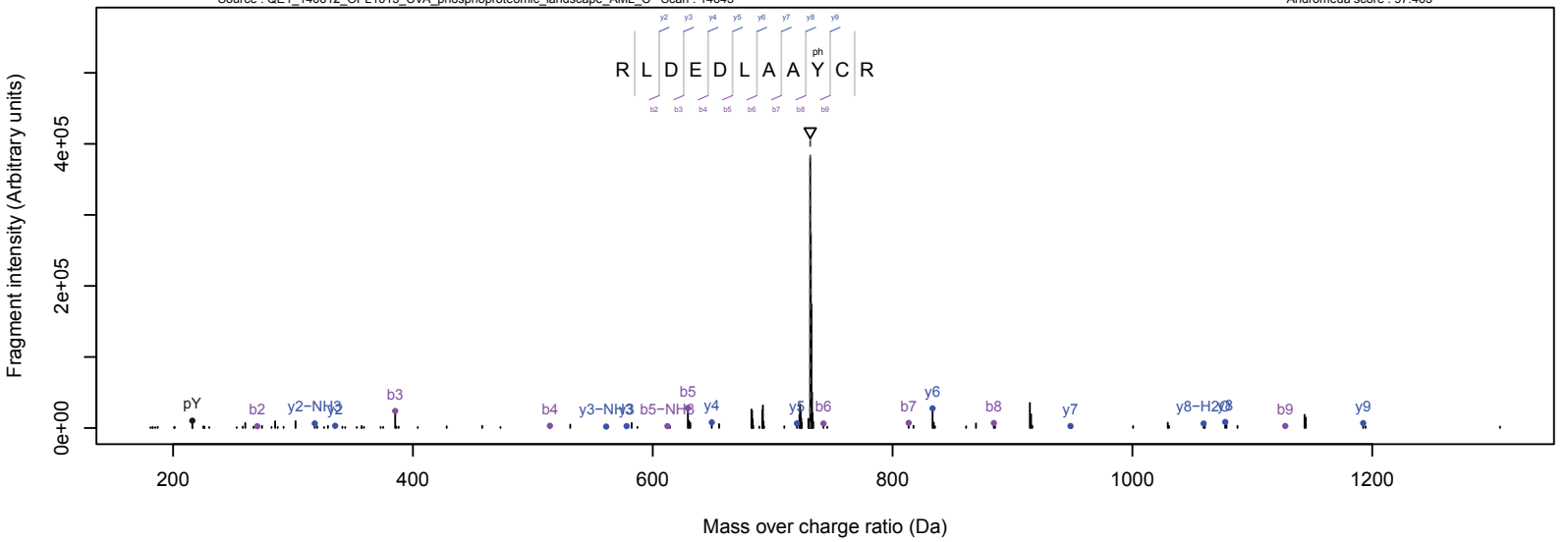
Andromeda score : 97.469



RLDEDLAAY(ph)CR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14643

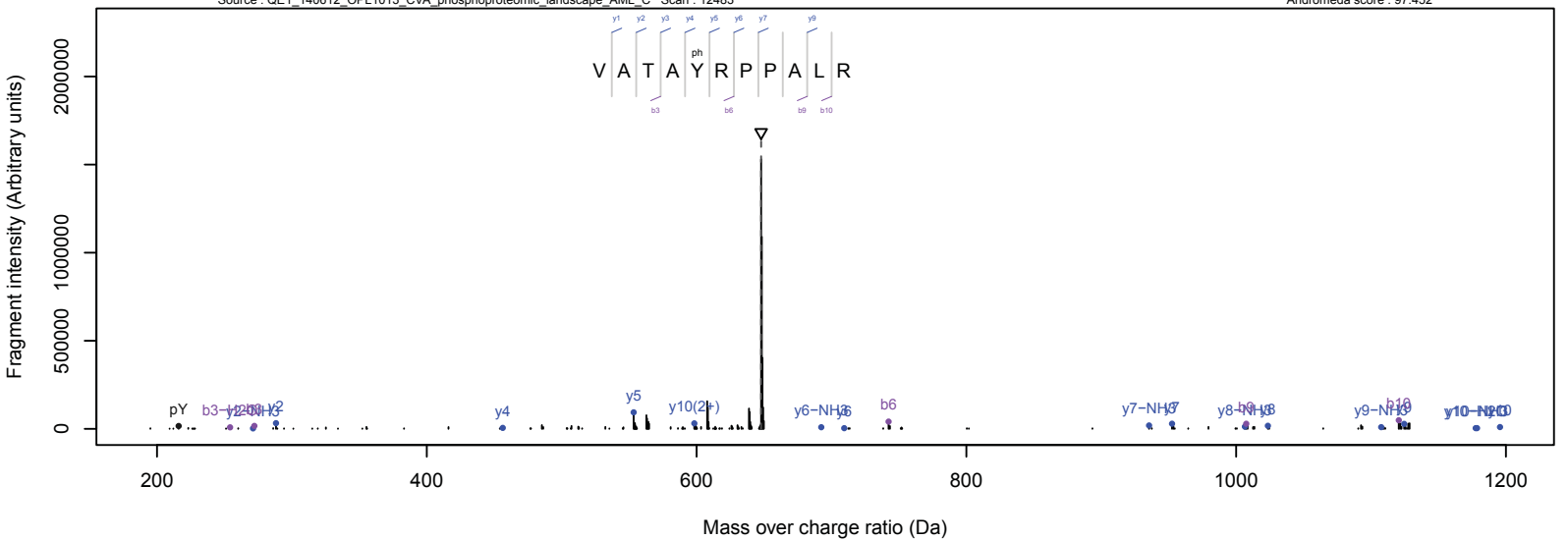
Andromeda score : 97.463



VATAY(ph)RPPALR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12483

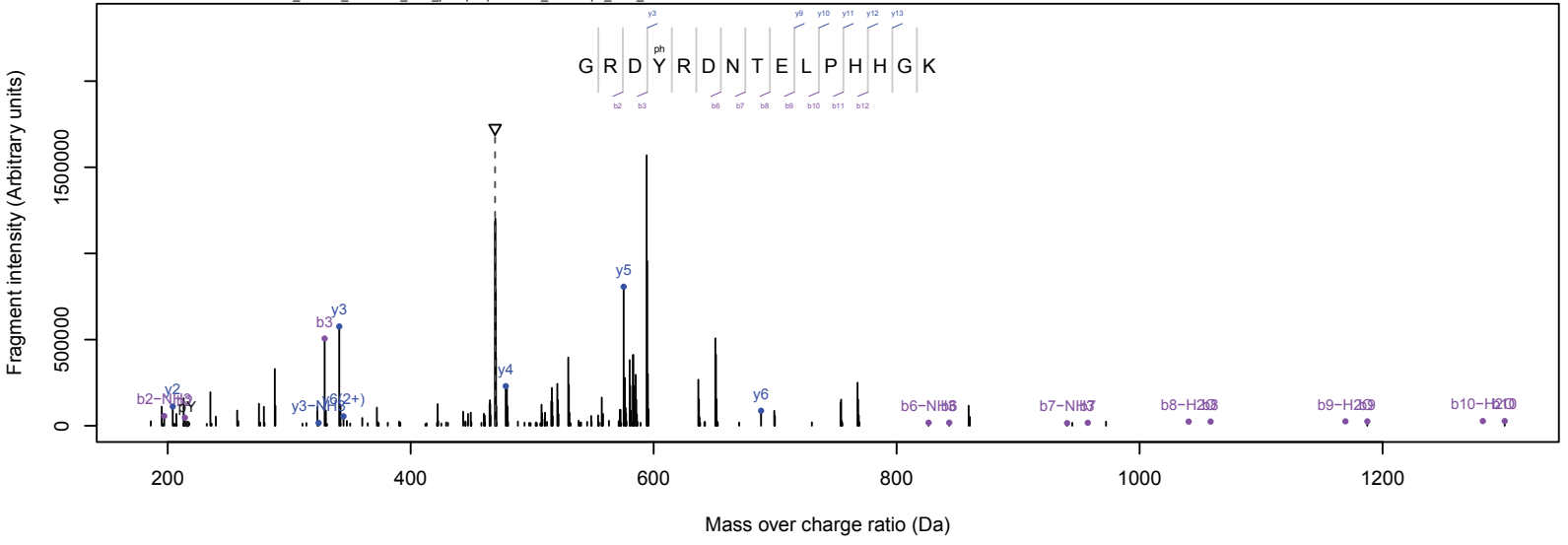
Andromeda score : 97.452



**GRDY(ph)RDNTLPHHGK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 3890

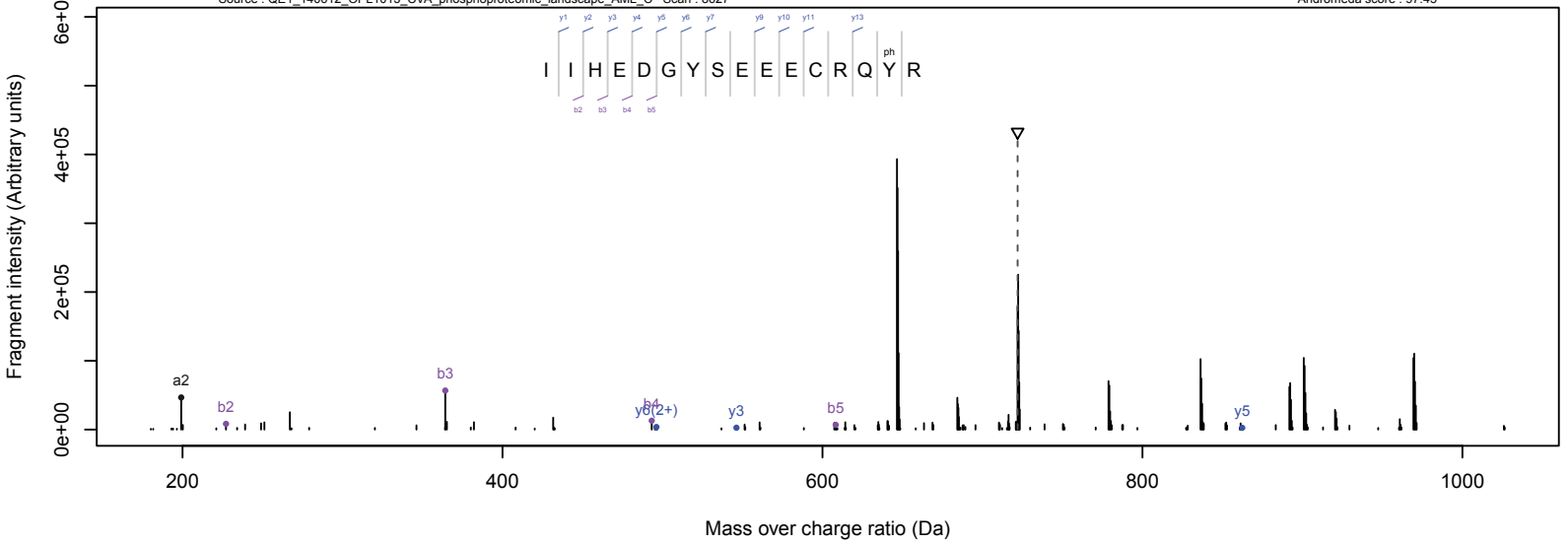
Andromeda score : 97.45



**I IHEDGYSEEECRQY(ph)R-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8627

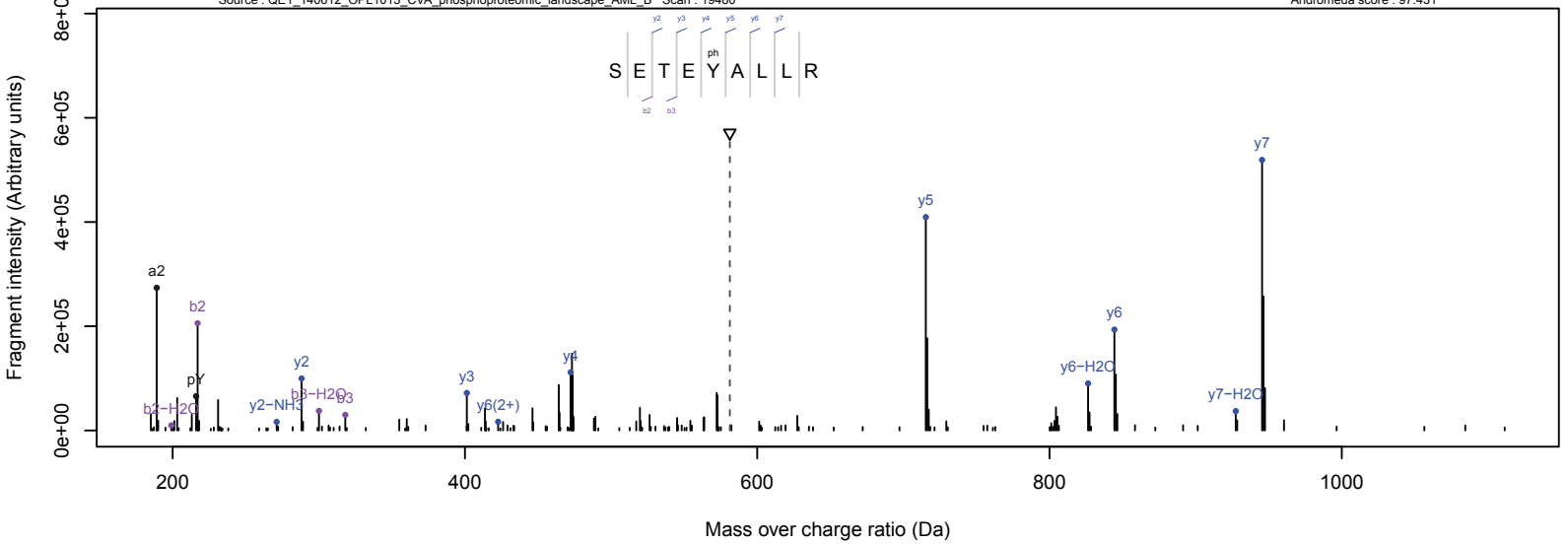
Andromeda score : 97.45



**SETEY(ph)ALLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19480

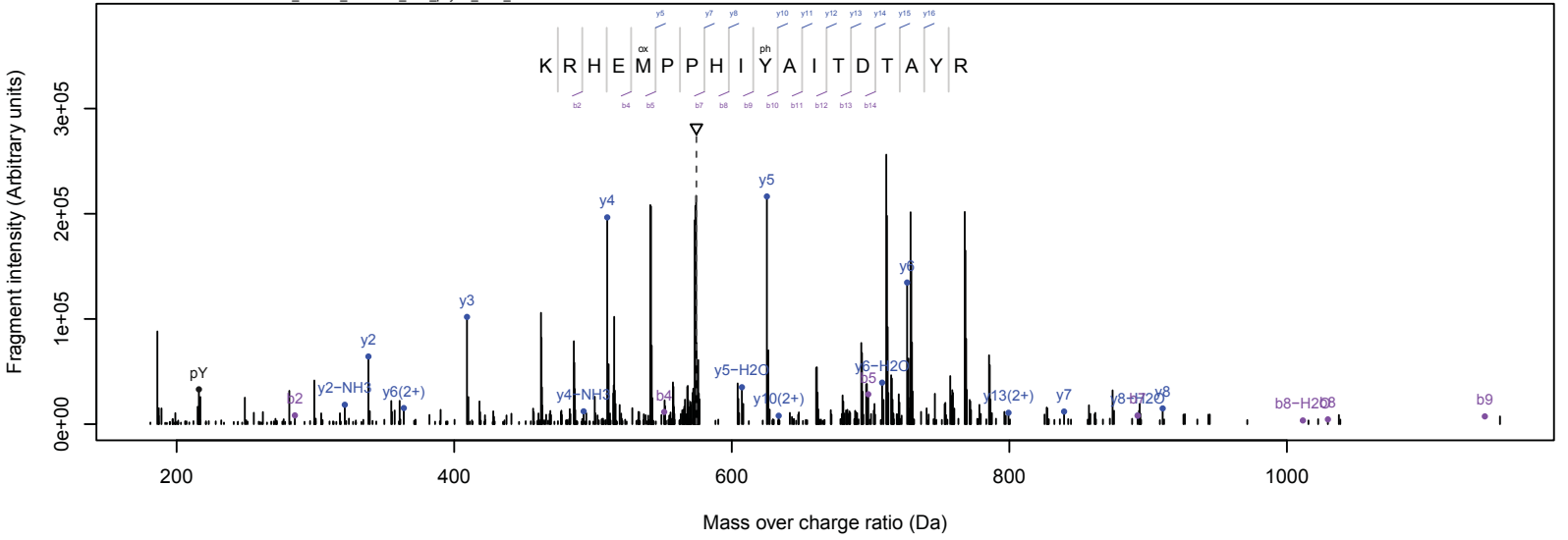
Andromeda score : 97.431



KRHEM(ox)PPIHY(ph)AITDTAYR-4+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 17143

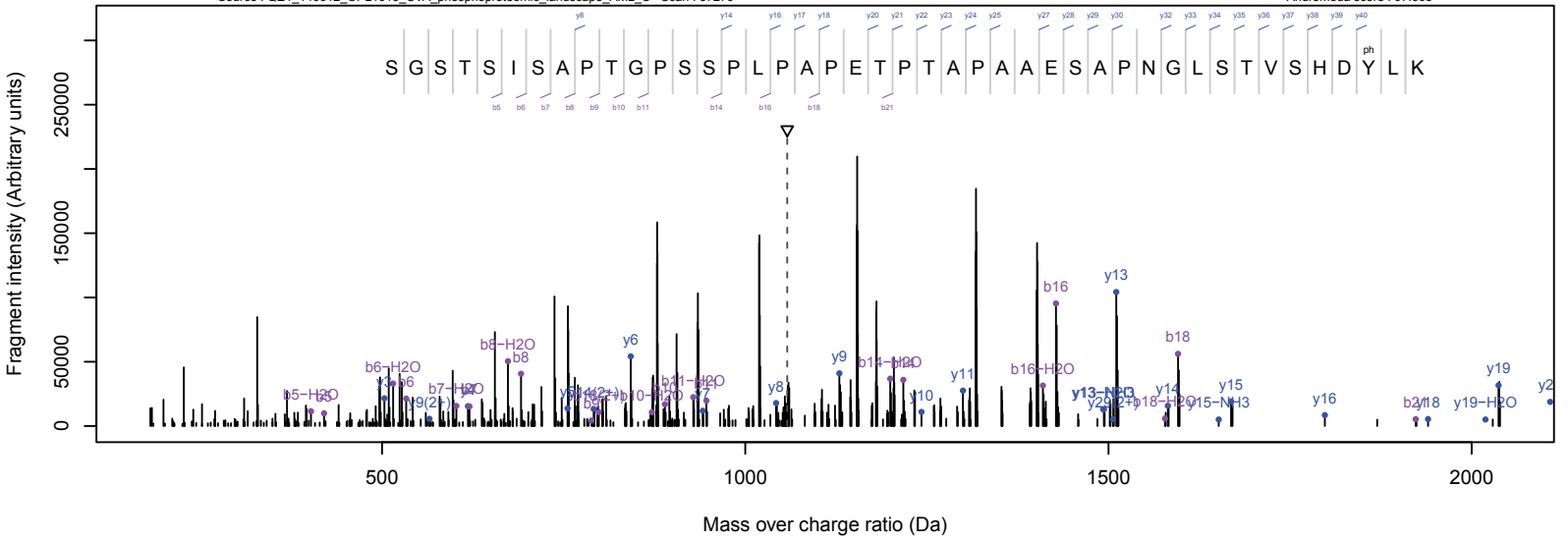
Andromeda score : 97.384



SGSTSISAPTGPSSPLPAPETPTAPAAESAPNGLSTVSHDY(ph)LK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37279

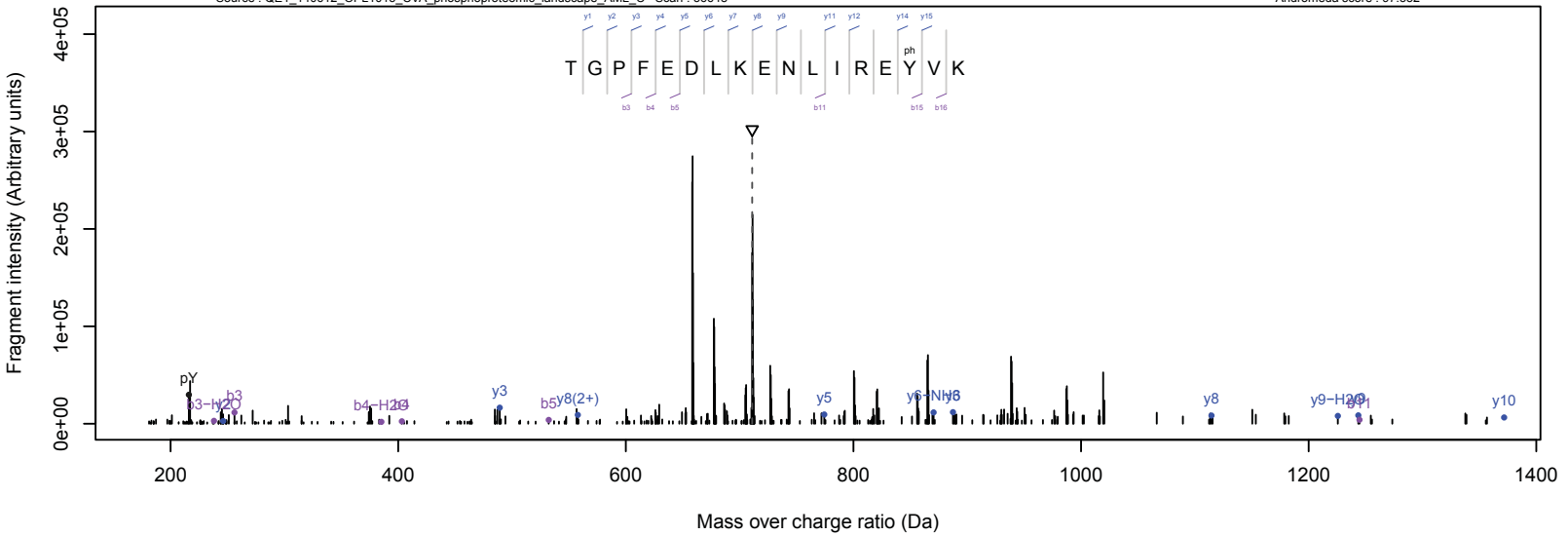
Andromeda score : 97.363



TGPFEDLKENLIREYV K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36648

Andromeda score : 97.352

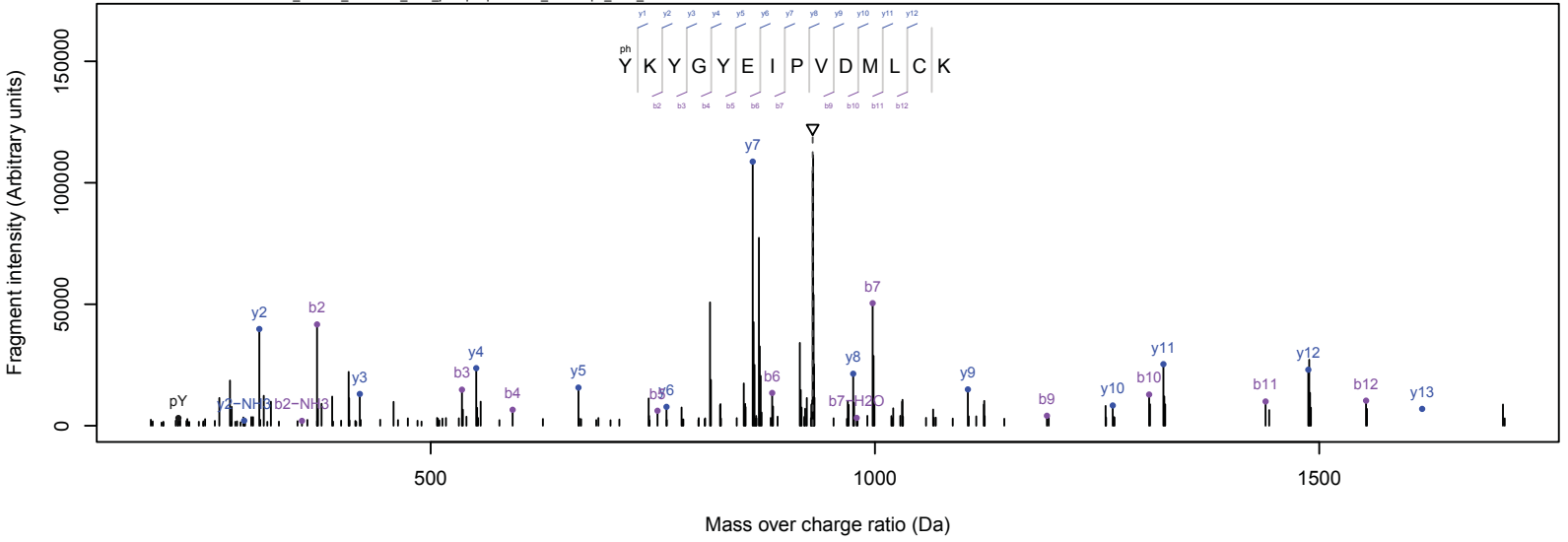




**Y(ph)KYG YEIPVDMLCK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 36645

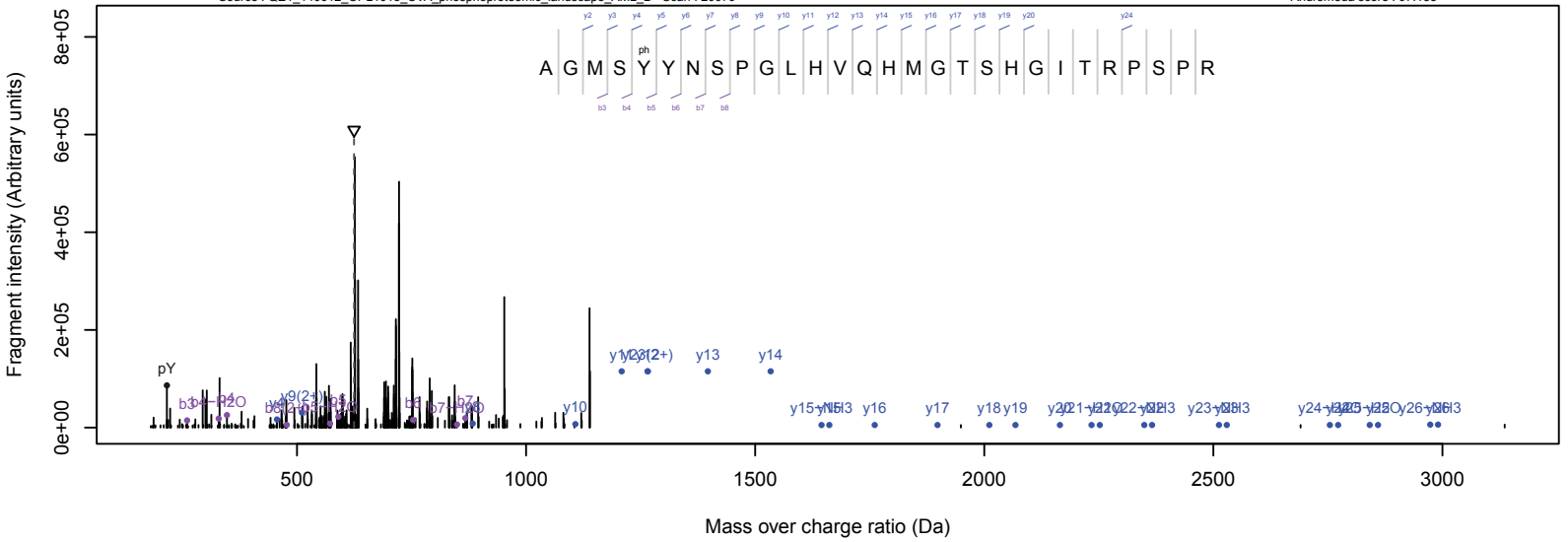
Andromeda score : 97.291



**AGMSY(ph)YNSPGLHVQHMGTSHGITRPSR-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20079

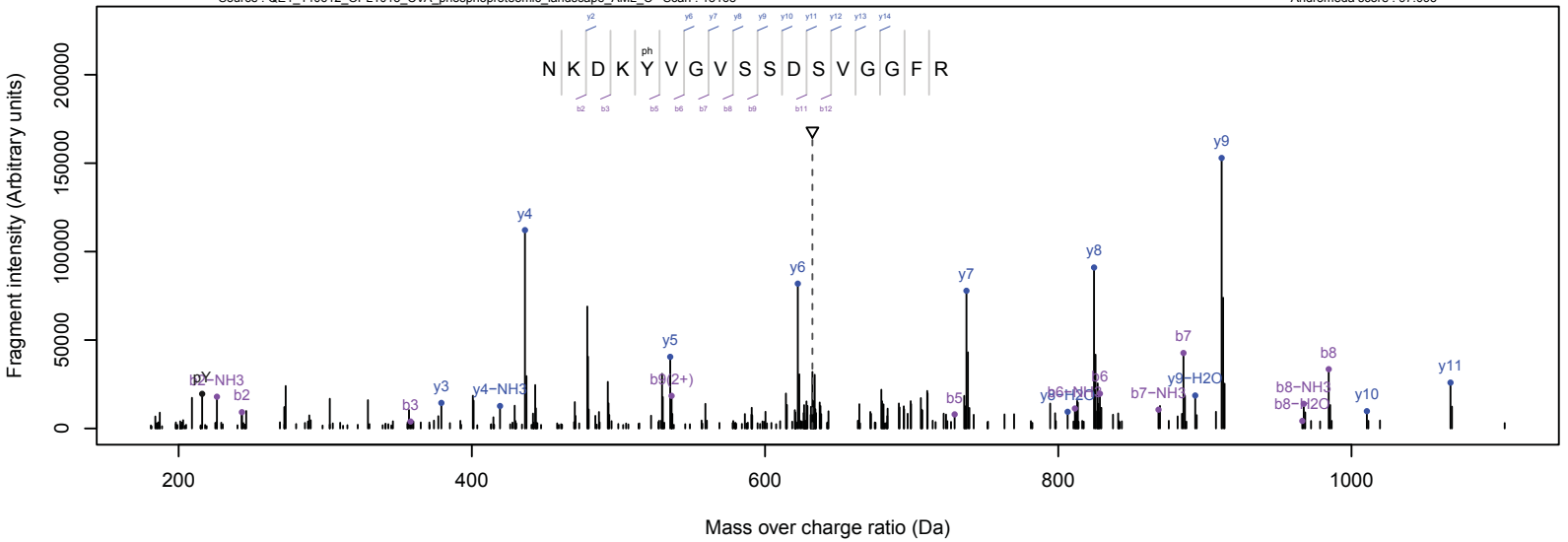
Andromeda score : 97.188



**NKDKY(ph)VGVS S D S V G G F R-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15168

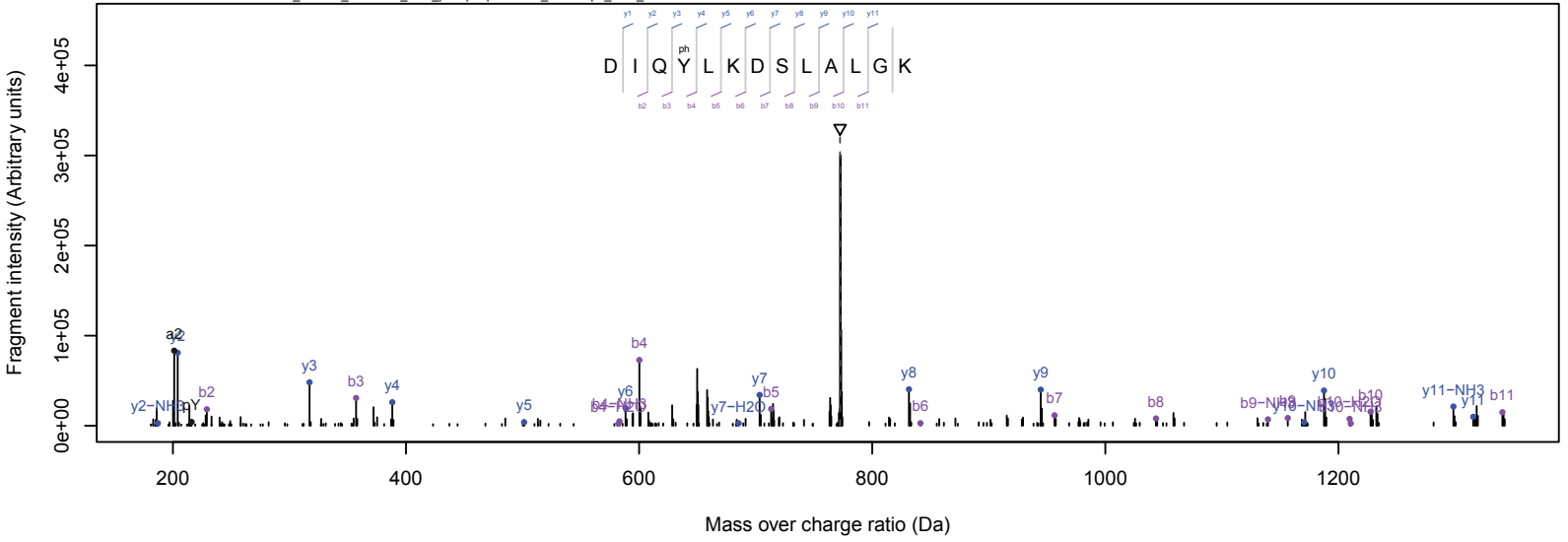
Andromeda score : 97.095



DIQY(ph)LKDSLALGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 31393

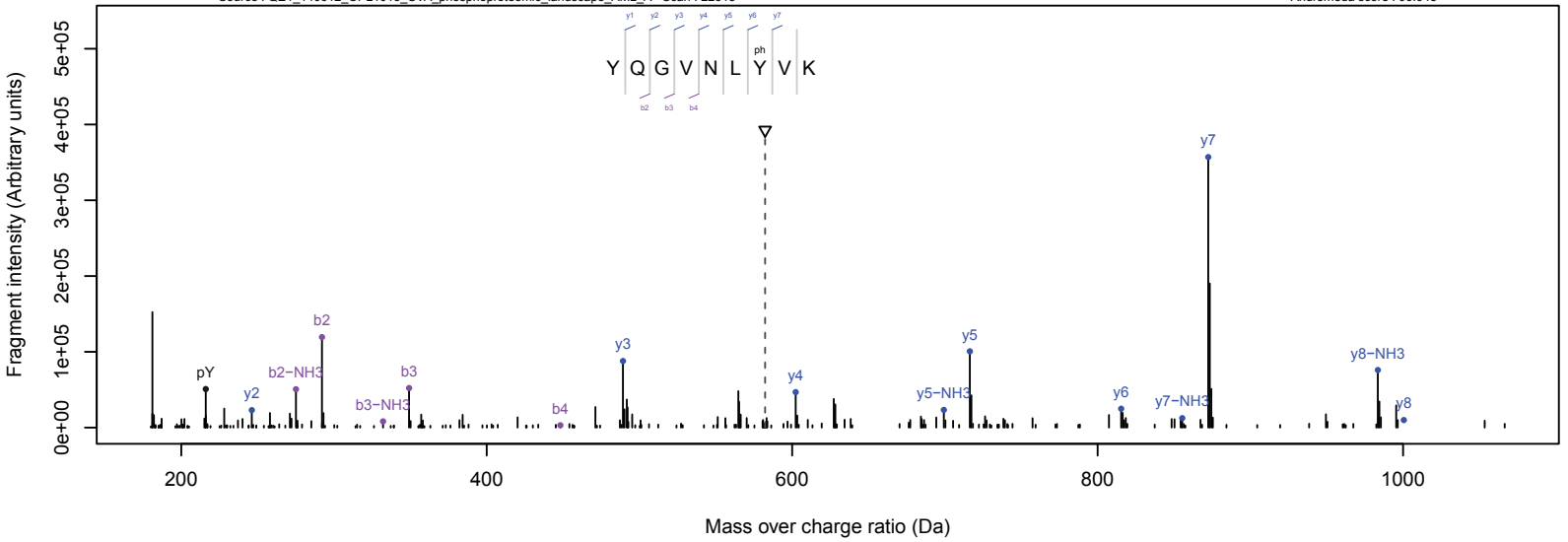
Andromeda score : 97.073



YQGVNLYVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 22318

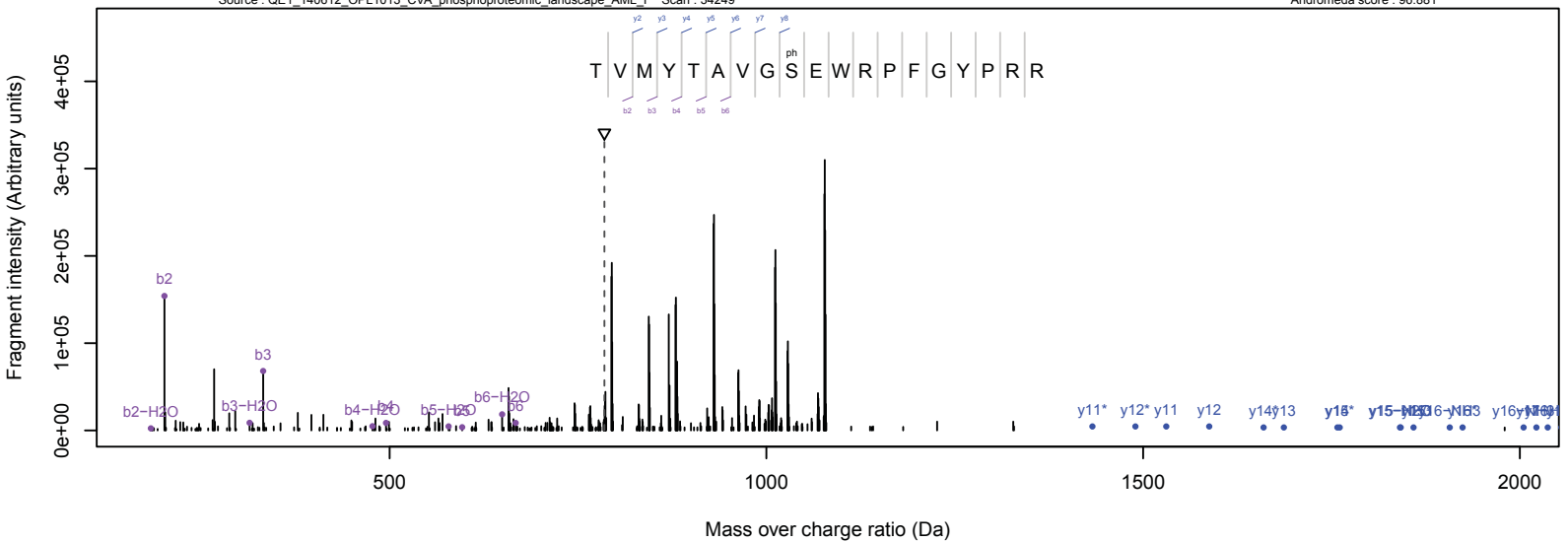
Andromeda score : 96.948



TVMYTAGVSEWRPFGYPRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 34249

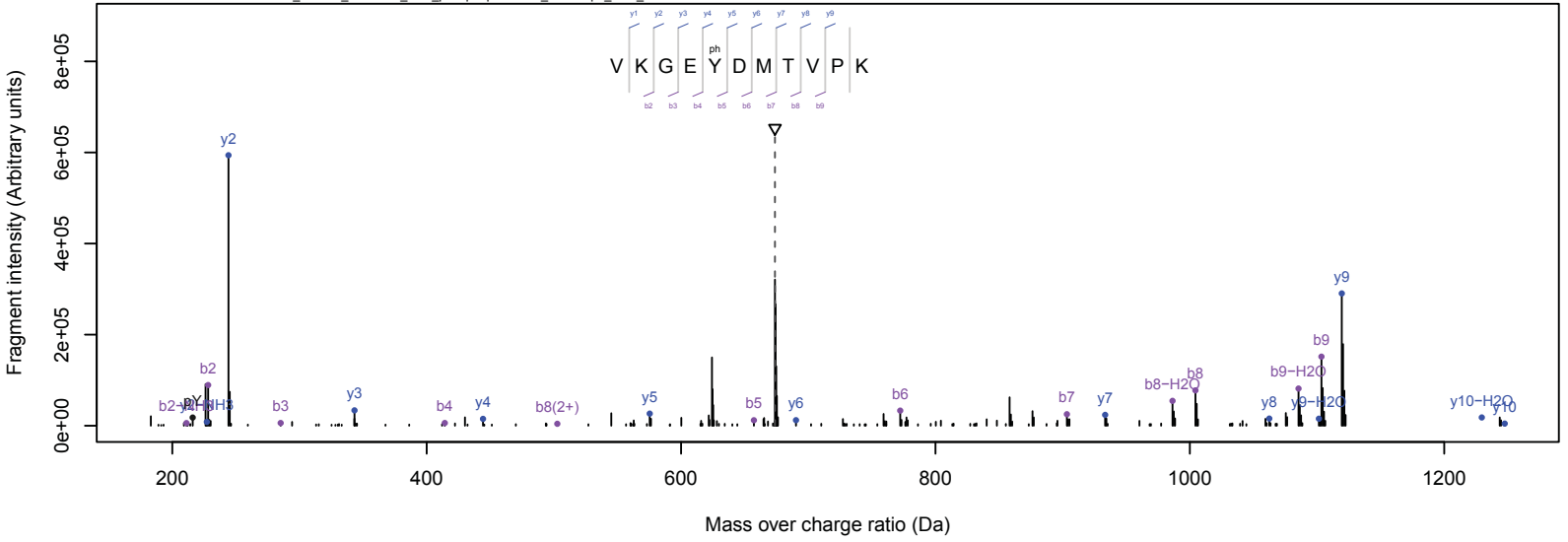
Andromeda score : 96.881



VKGEY(ph)DMTVPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10886

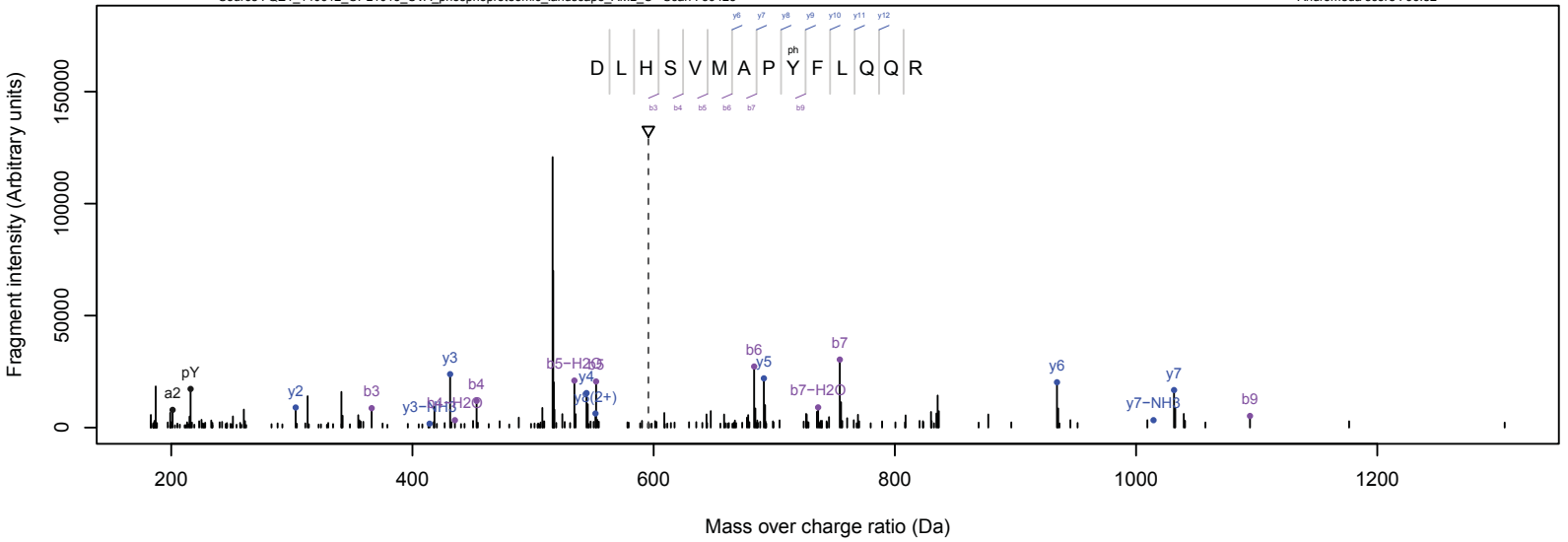
Andromeda score : 96.866



DLHSVMA<sup>ph</sup>YFLQQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39428

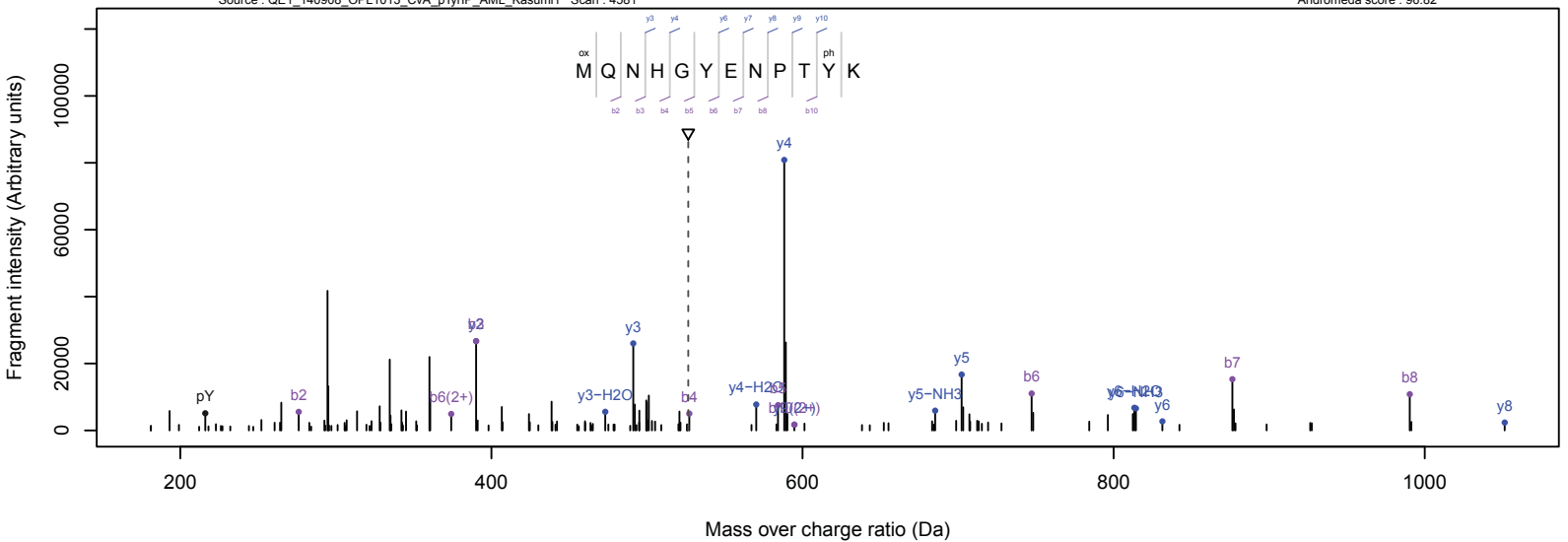
Andromeda score : 96.82



M(ox)QN<sup>H</sup>GYENPT<sup>ph</sup>YK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 4581

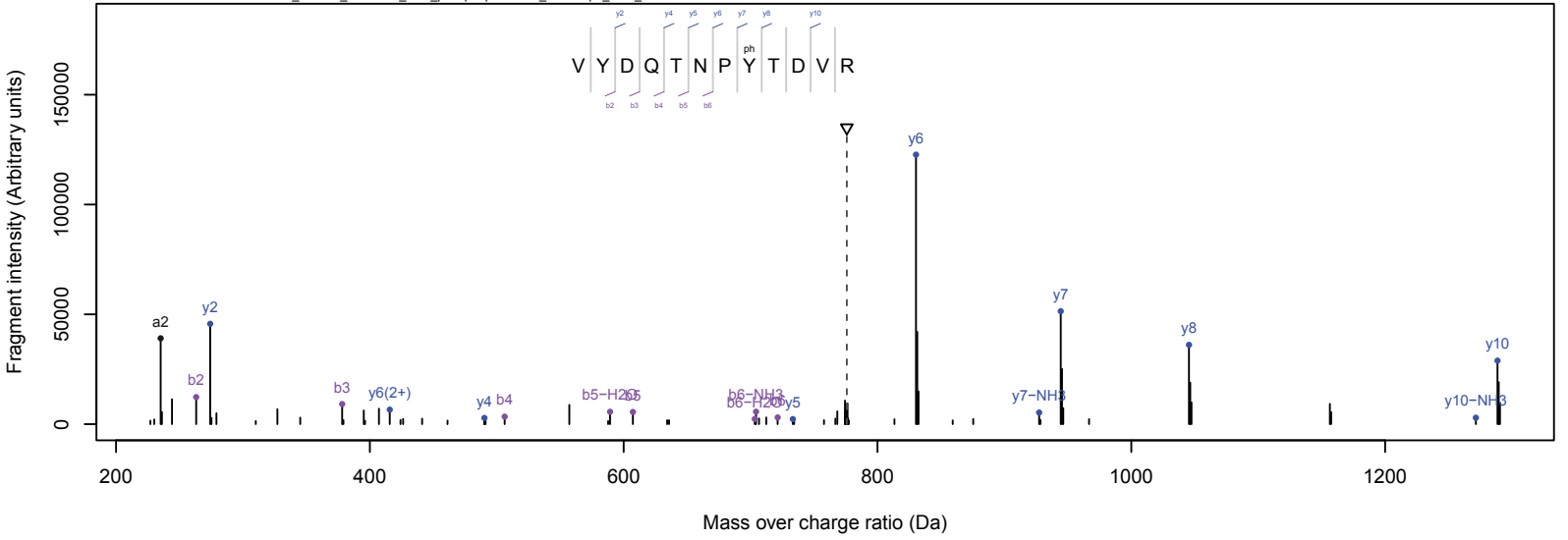
Andromeda score : 96.82



**VYDQTNPY(ph)TDVR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 10737

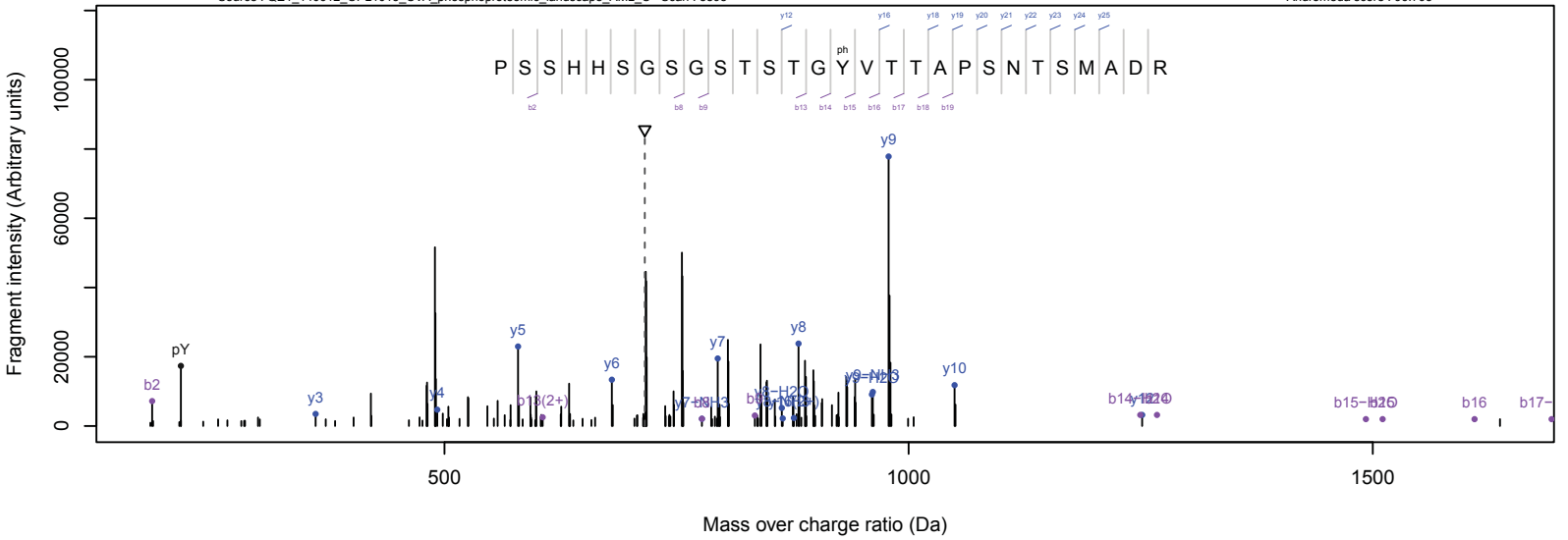
Andromeda score : 96.82



**PSSHHS GSGSTSTGY(ph)VTTAPSNNTSMADR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8898

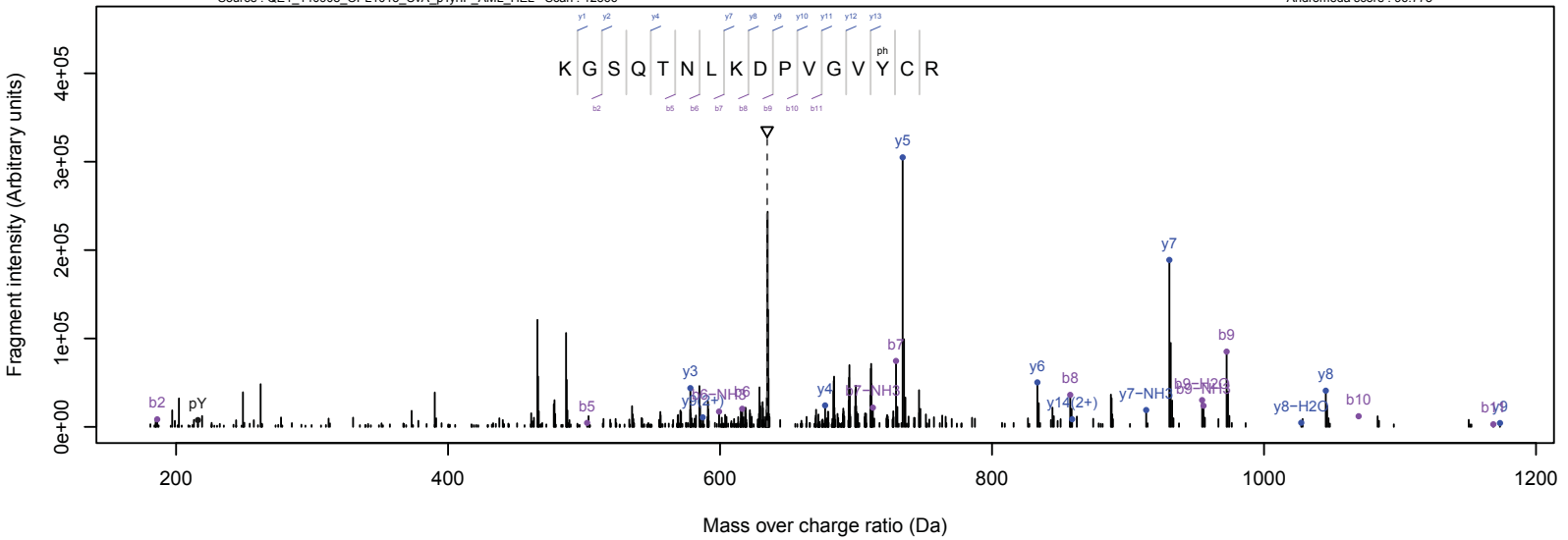
Andromeda score : 96.793



**KGSQTNLKDVPVGY(ph)CR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 12566

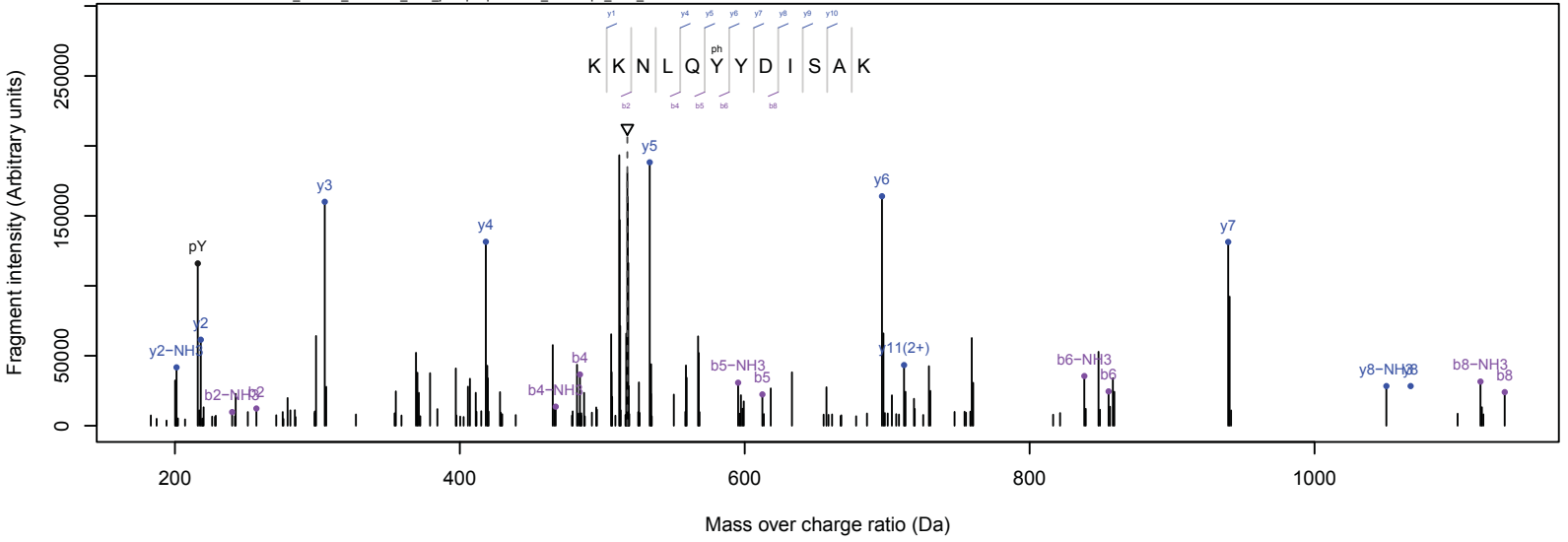
Andromeda score : 96.773



**KKNLQY(ph)YDISAK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12283

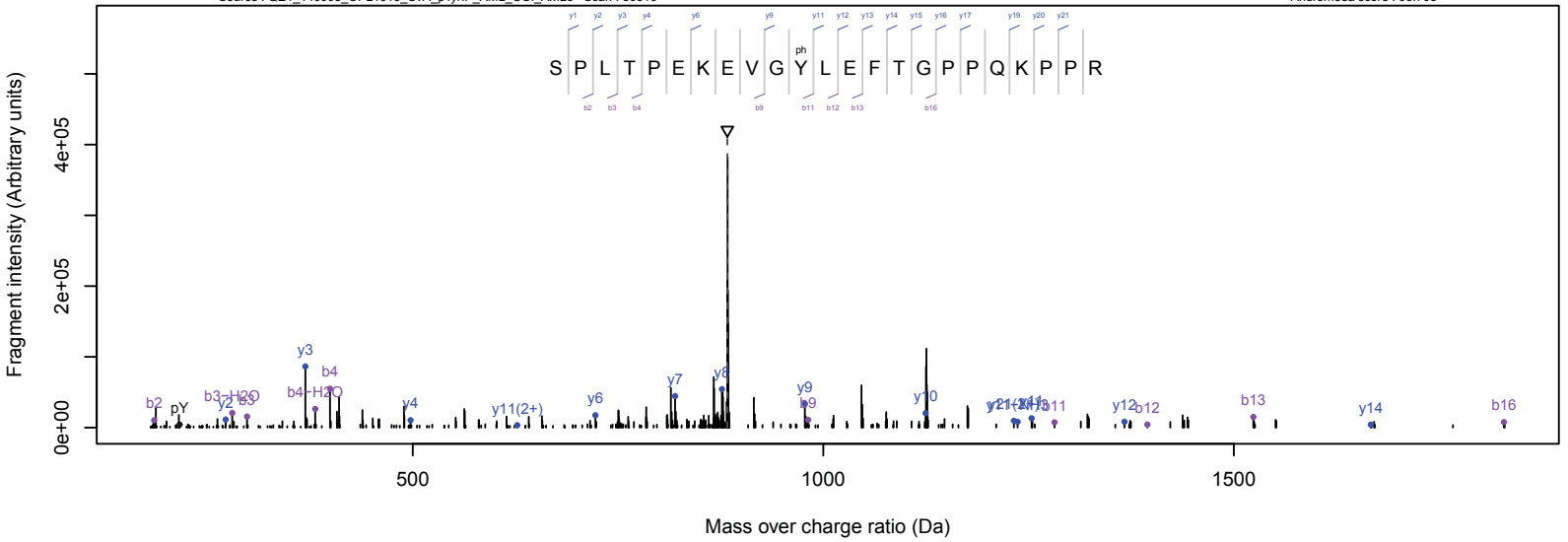
Andromeda score : 96.745



**SPLTPEKEVGY(ph)LEFTGPPQKPPR-3+**

Source : QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_OCI\_AML3 Scan : 36516

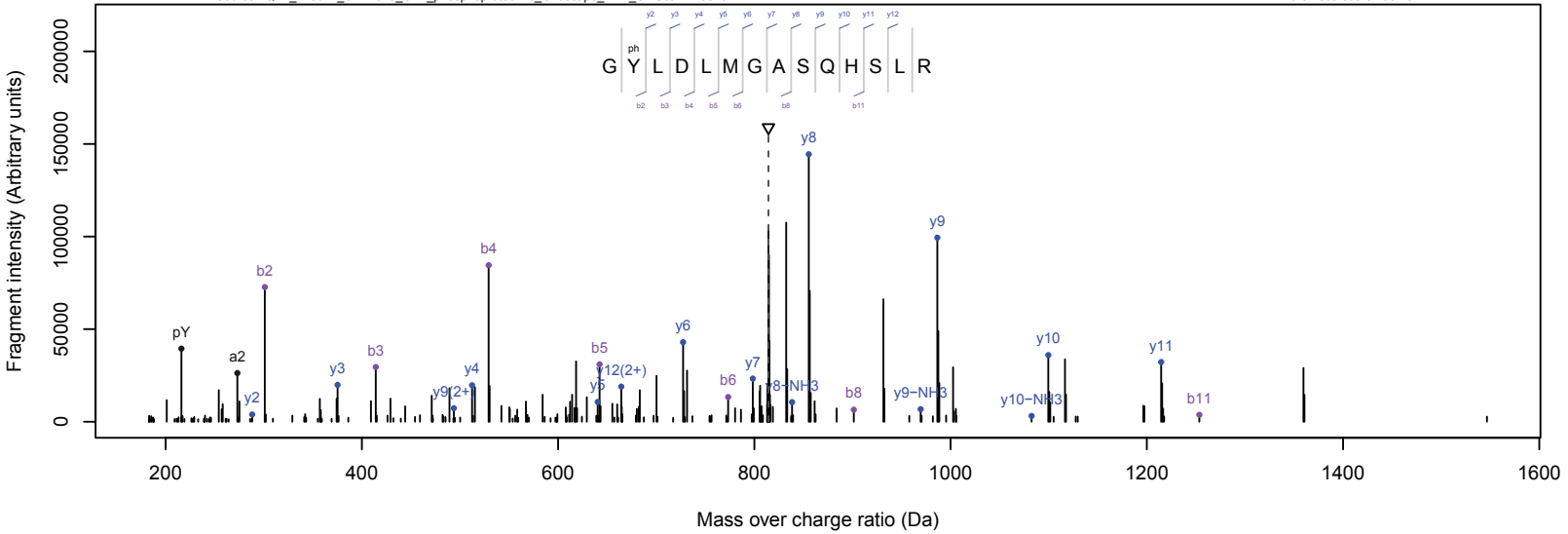
Andromeda score : 96.708



**GY(ph)LDLMGASQHSLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29070

Andromeda score : 96.707

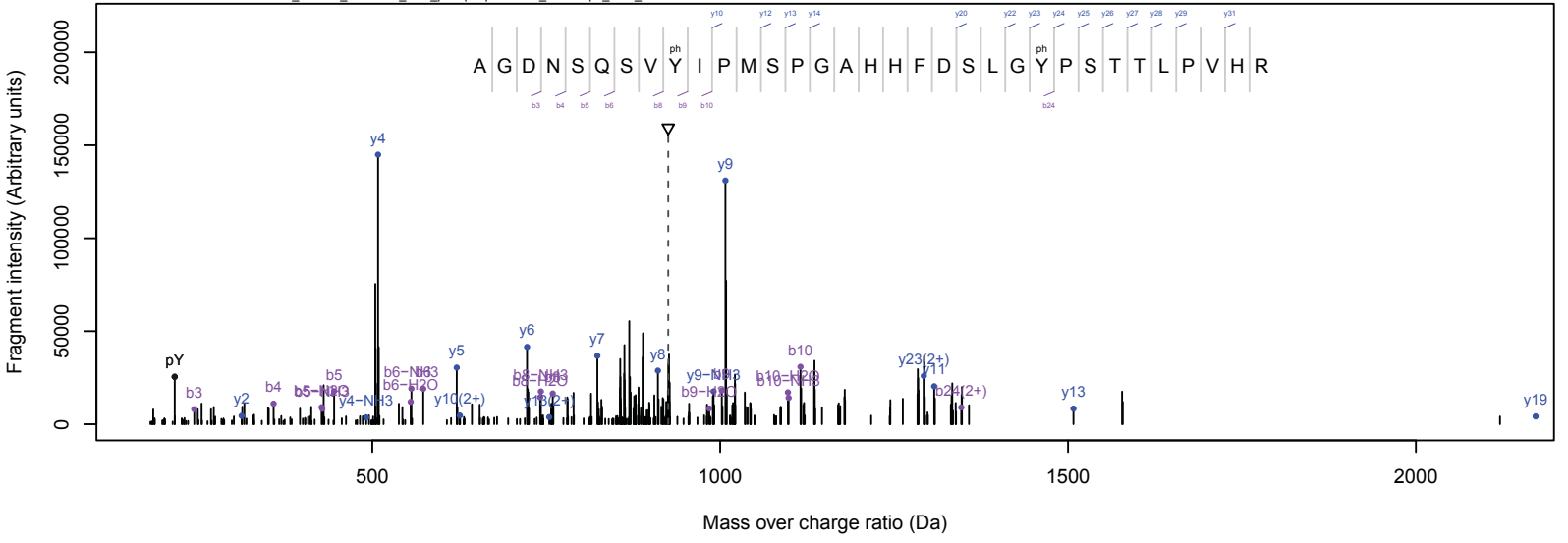




AGDNSQSVY(ph)IPMSPGAHHFDSLGY(ph)PSTTLPVHR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37915

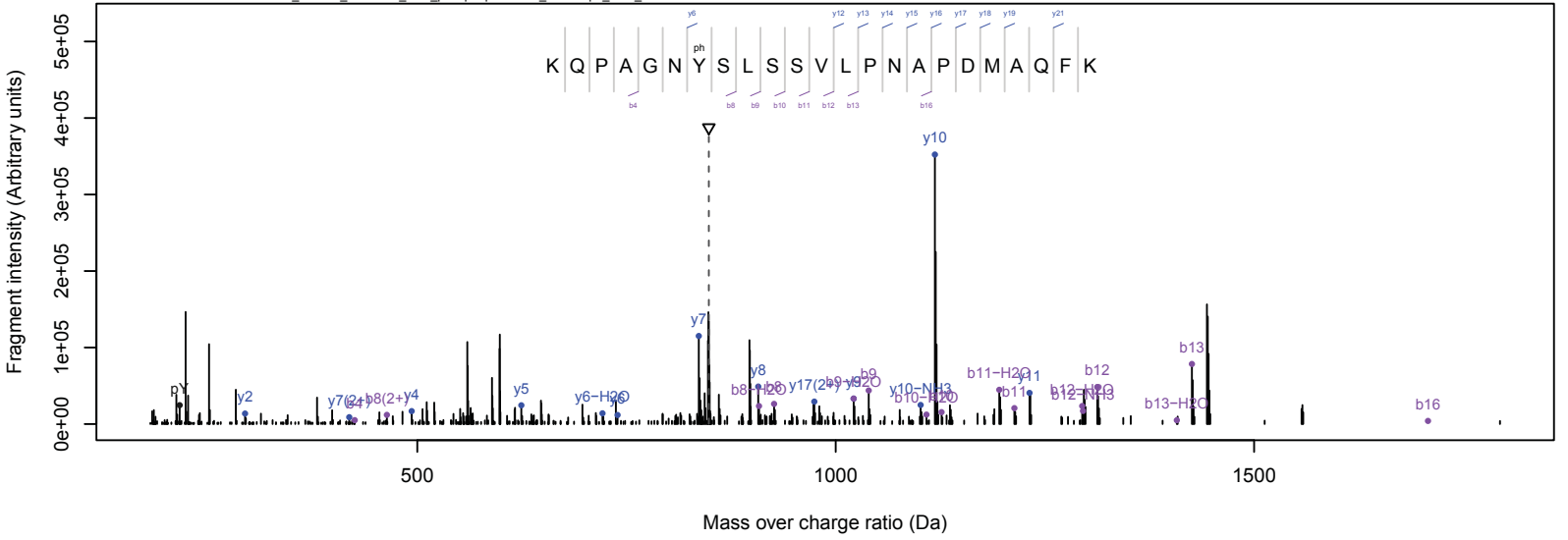
Andromeda score : 96.598



KQPAGNY(ph)SLSSVLPNAPDMAQFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39191

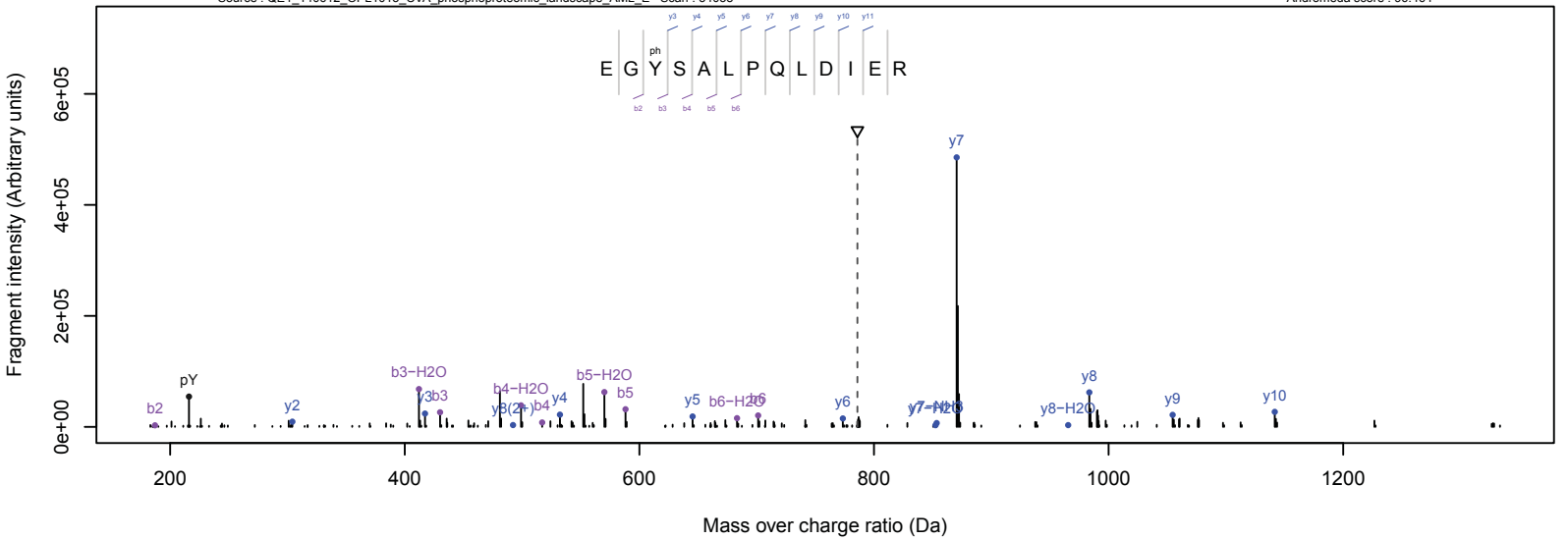
Andromeda score : 96.591



EGY(ph)SALPLQLDIER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 34038

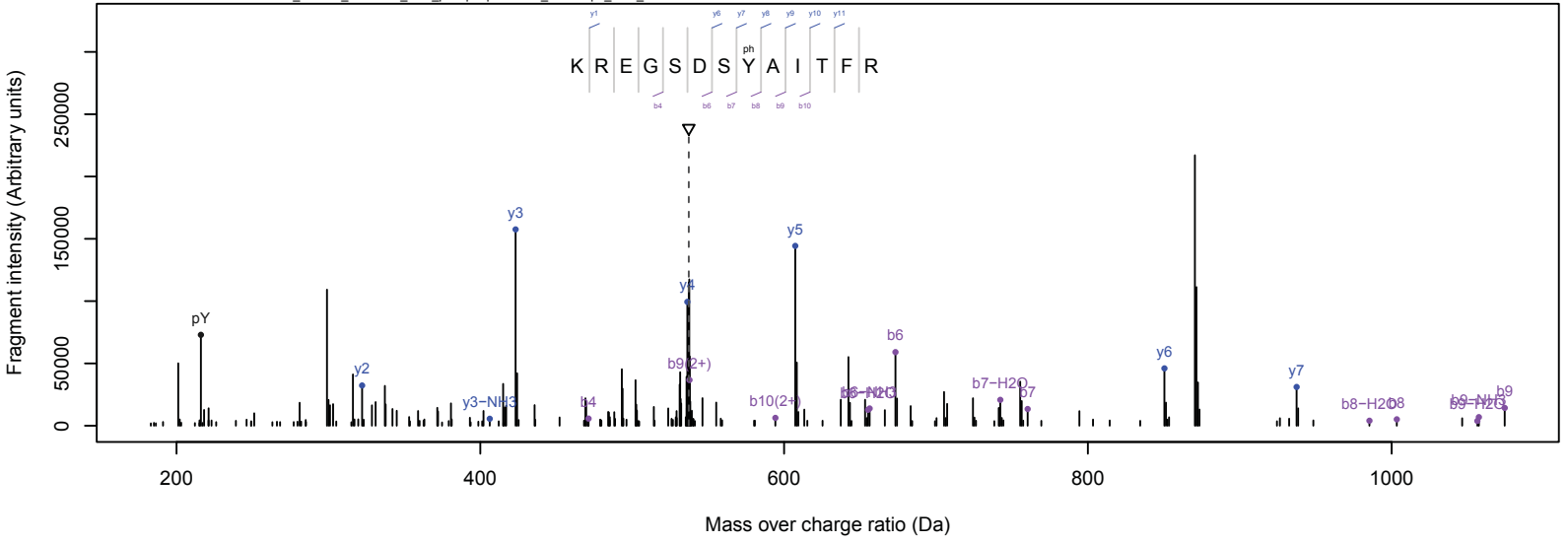
Andromeda score : 96.464



**KREGSDSY(ph)AITFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 12378

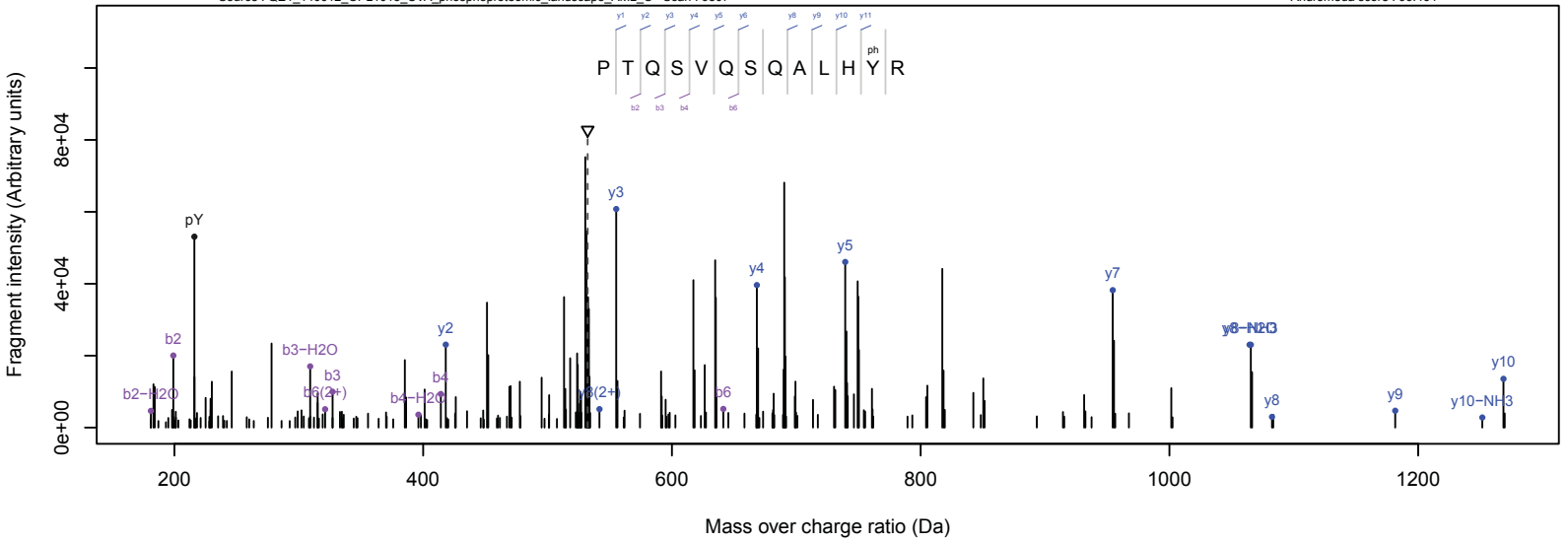
Andromeda score : 96.464



**PTQSVQSQALHY(ph)R-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9807

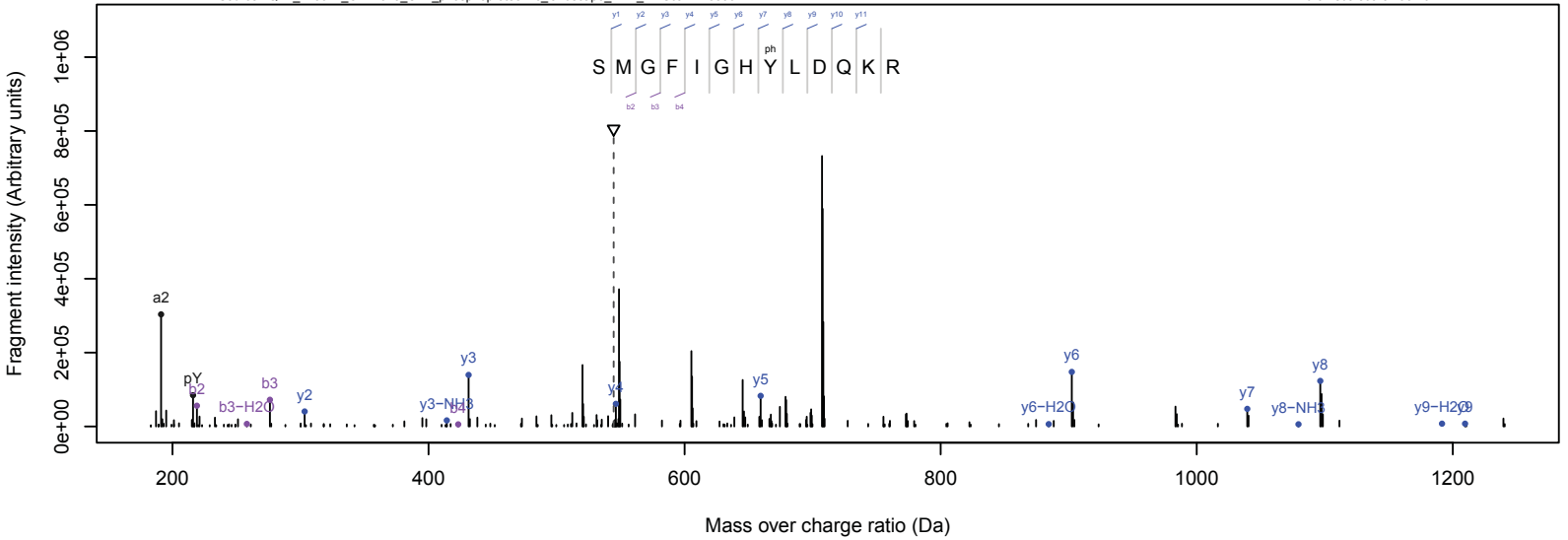
Andromeda score : 96.464



**SMGFIGHYLDQKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26036

Andromeda score : 96.464

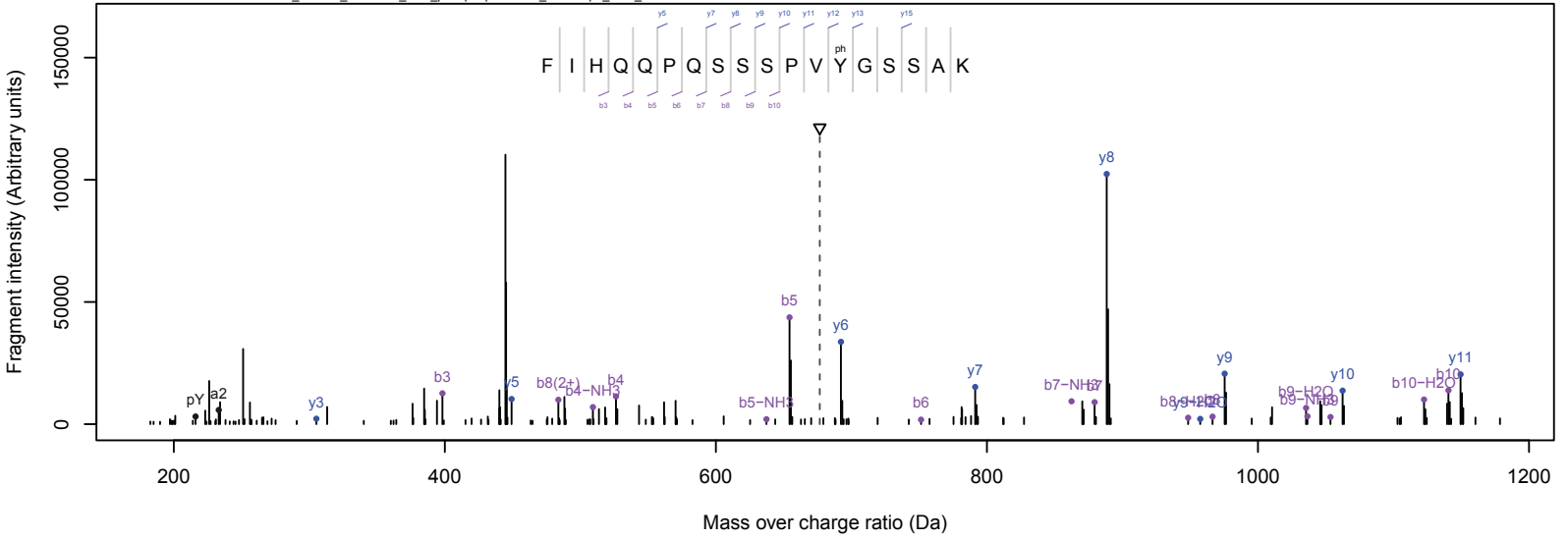




**FIHQQPSSSPV(ph)GSSAK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10015

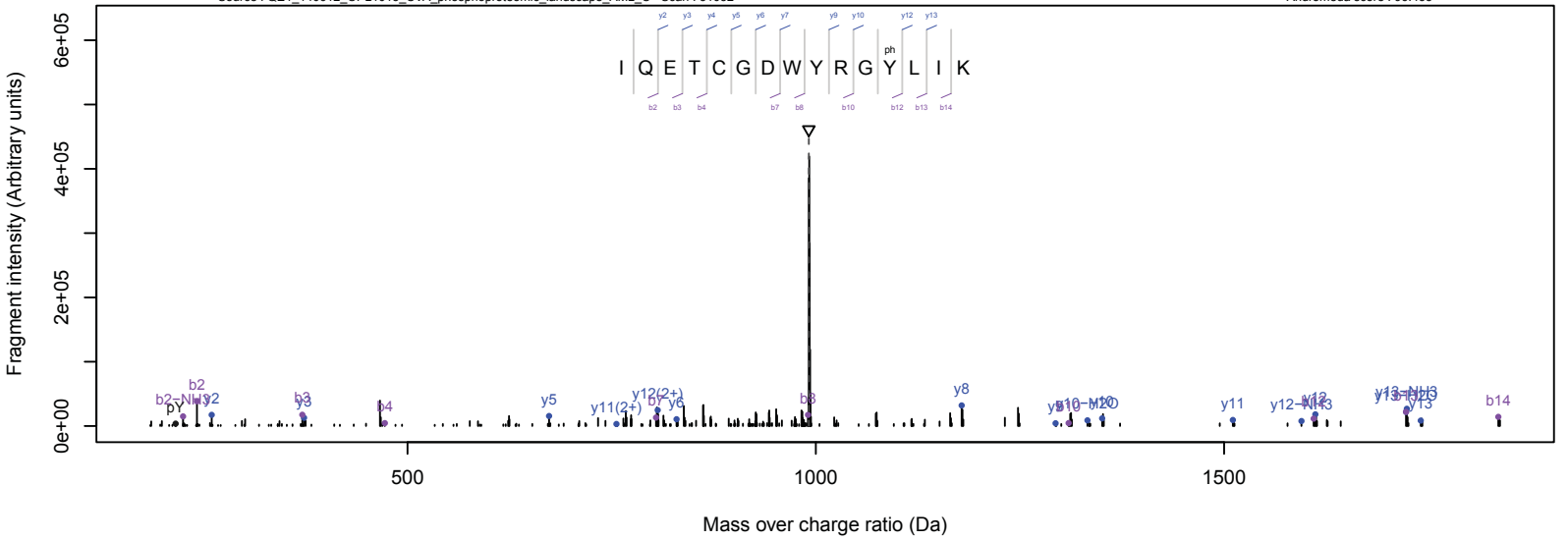
Andromeda score : 96.447



**IQETCGDWYRGY(ph)LIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31052

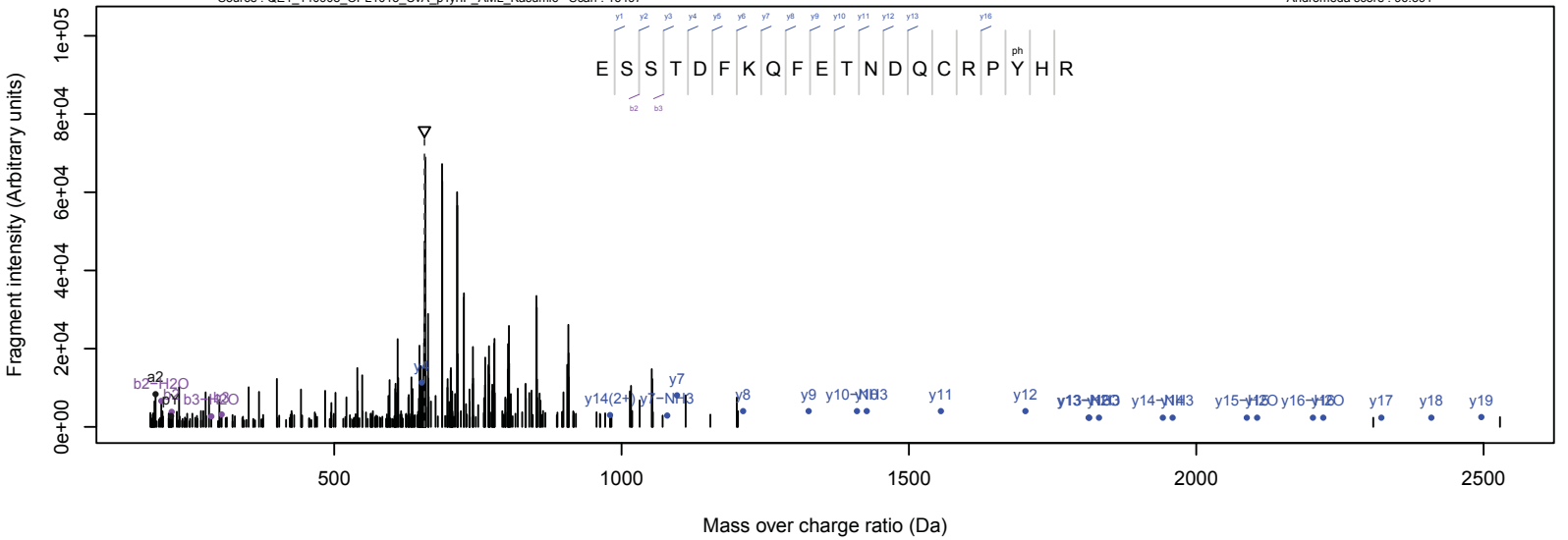
Andromeda score : 96.435



**ESSTDFKQFETNDQCRPY(ph)HR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 16497

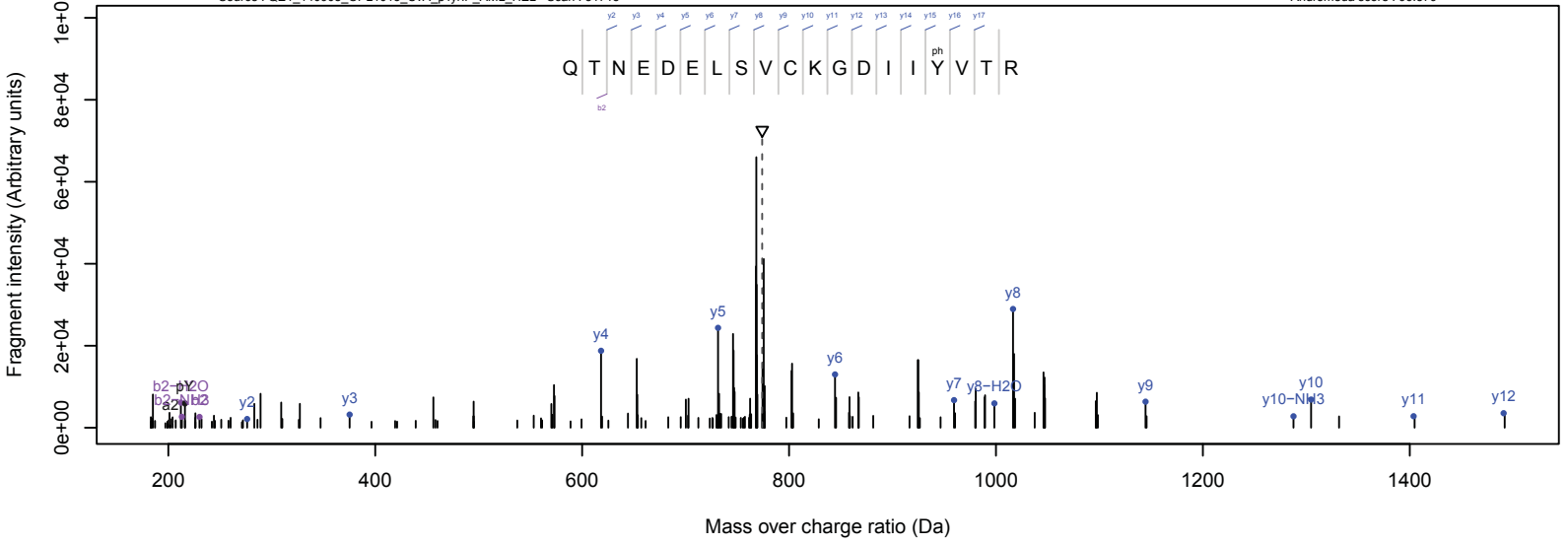
Andromeda score : 96.391



QTNEDELSVCKGDIYVTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31715

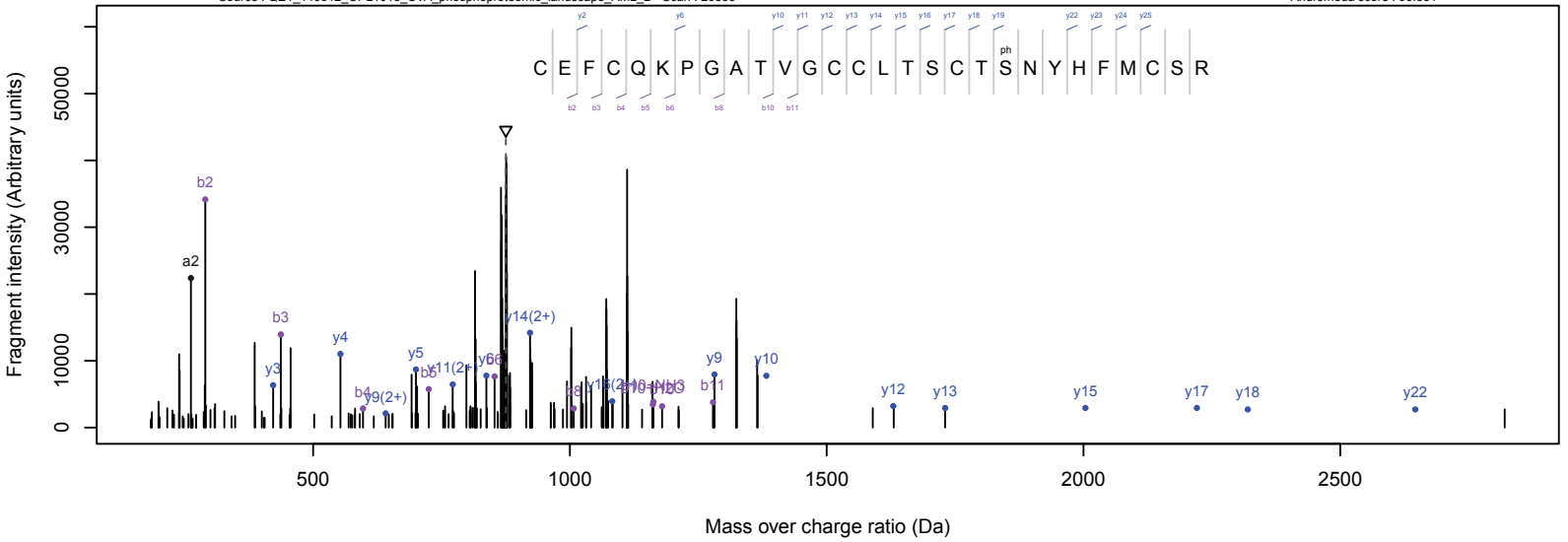
Andromeda score : 96.379



CEFCQKPGATVGCCLTSCTS(ph)NYHFMCSR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26853

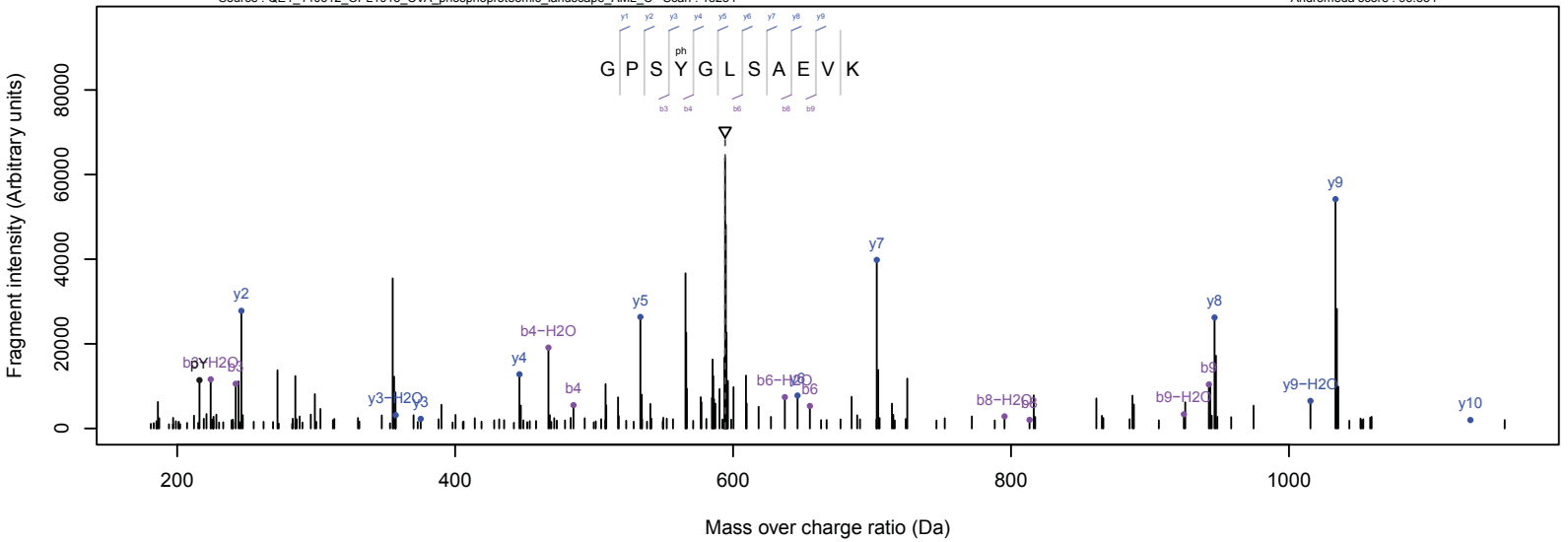
Andromeda score : 96.351



GPSY(ph)GLSAEVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13254

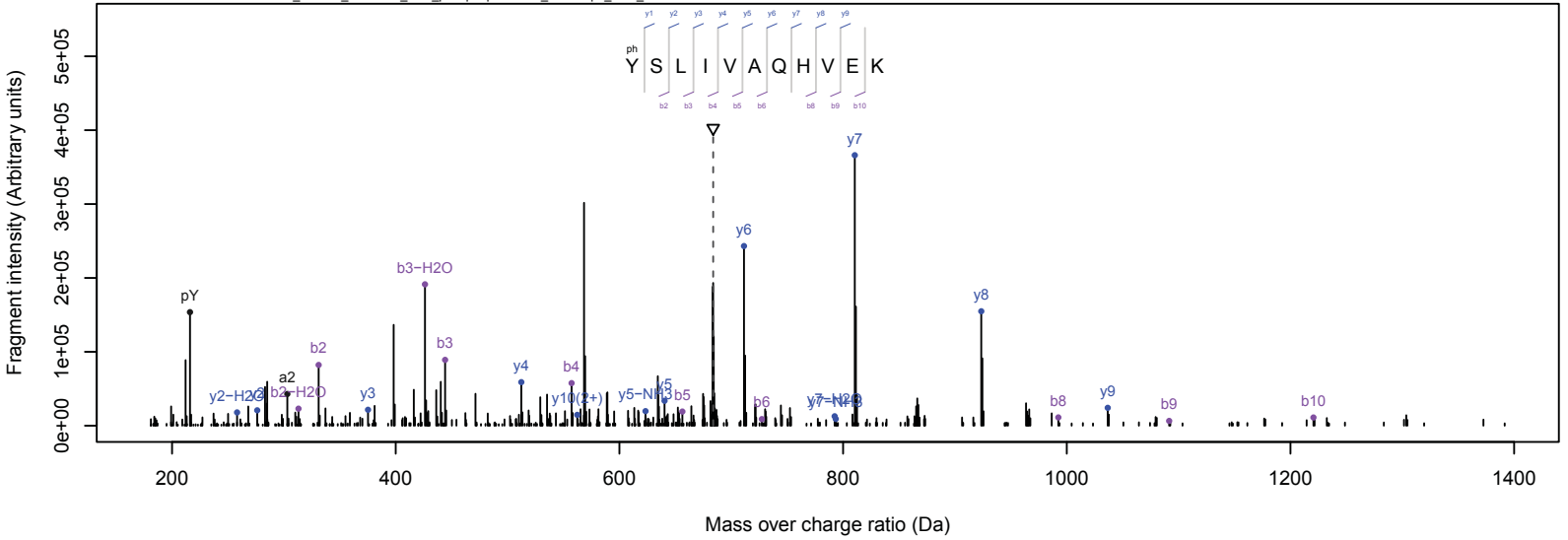
Andromeda score : 96.334



Y(ph)SLIVAQHVEK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17073

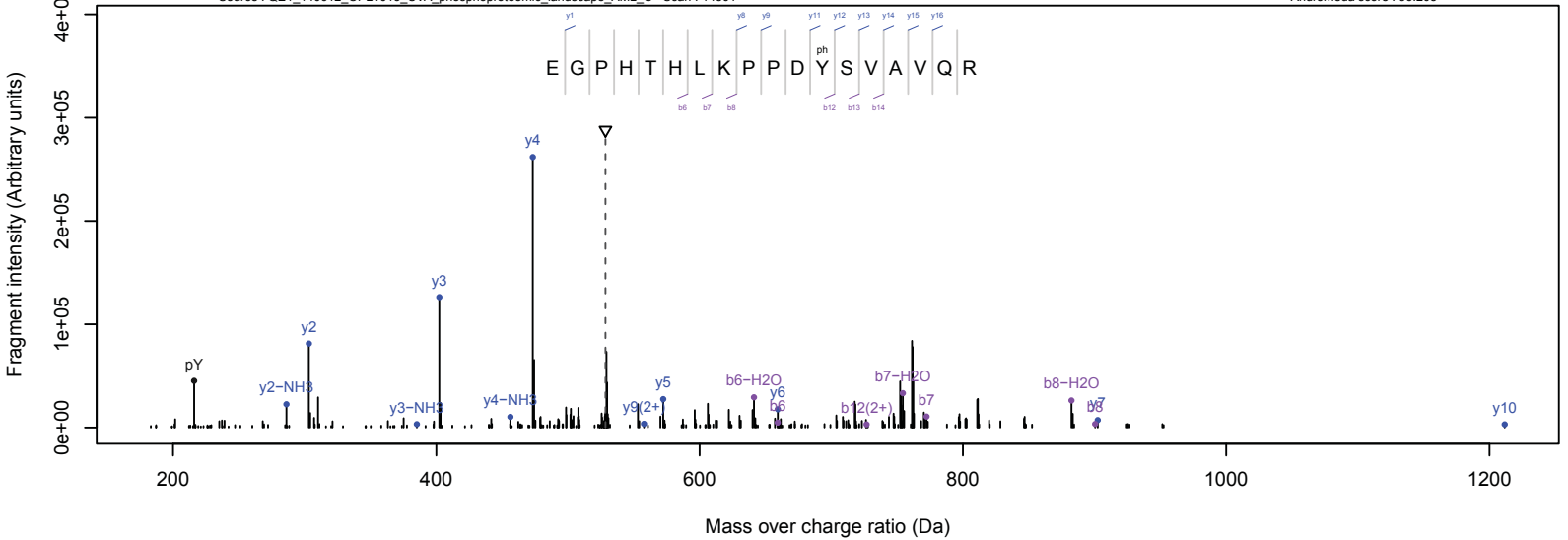
Andromeda score: 96.23



EGPHTLKPDPY(ph)SVAQR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14501

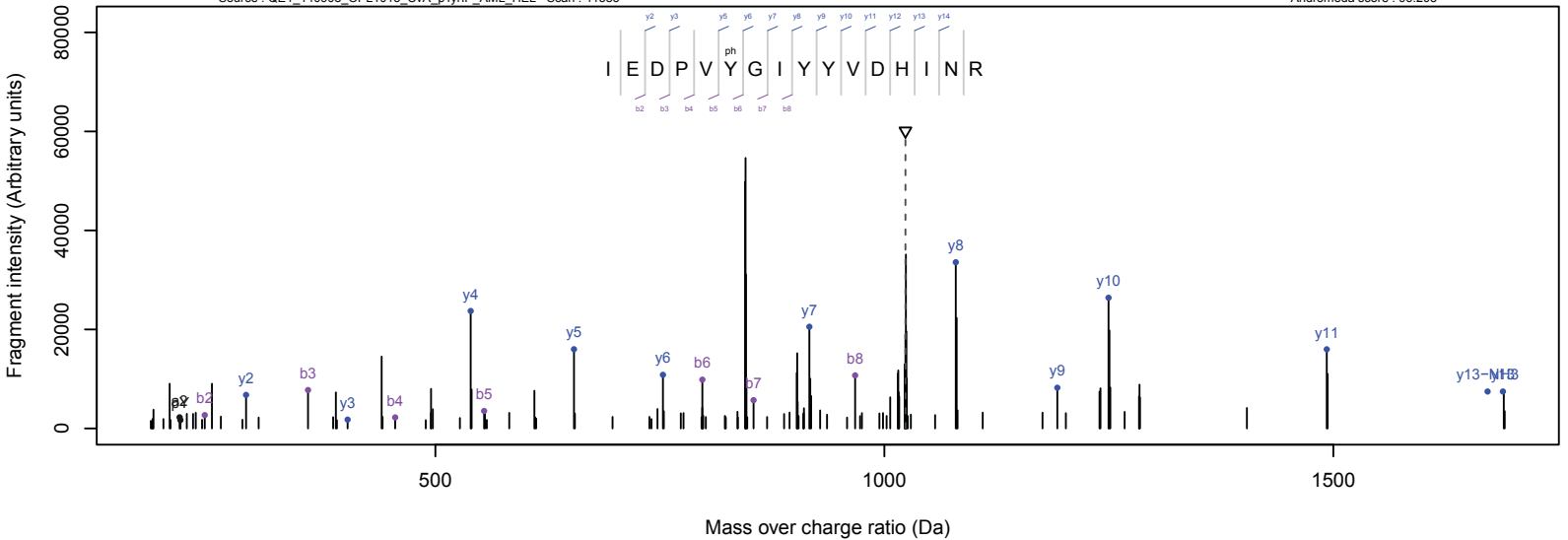
Andromeda score: 96.208



IEDPVY(ph)GIYYVDHINR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 41385

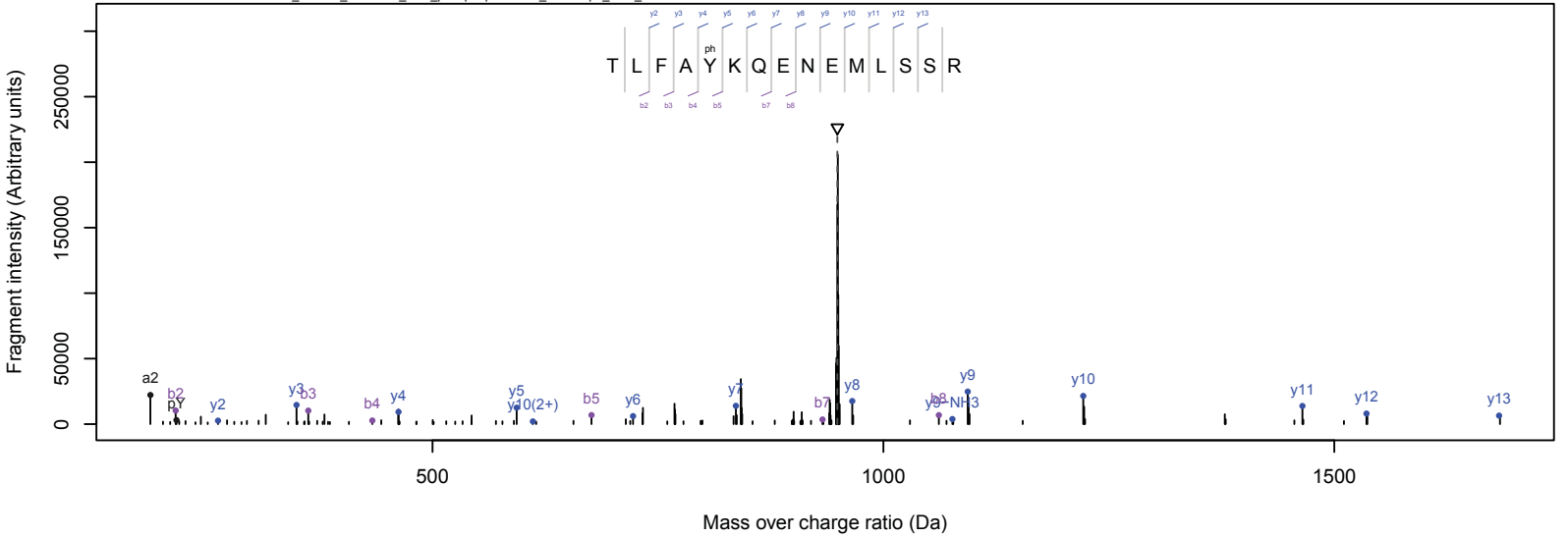
Andromeda score: 96.208



TLFAY(ph)KQENEMLSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25662

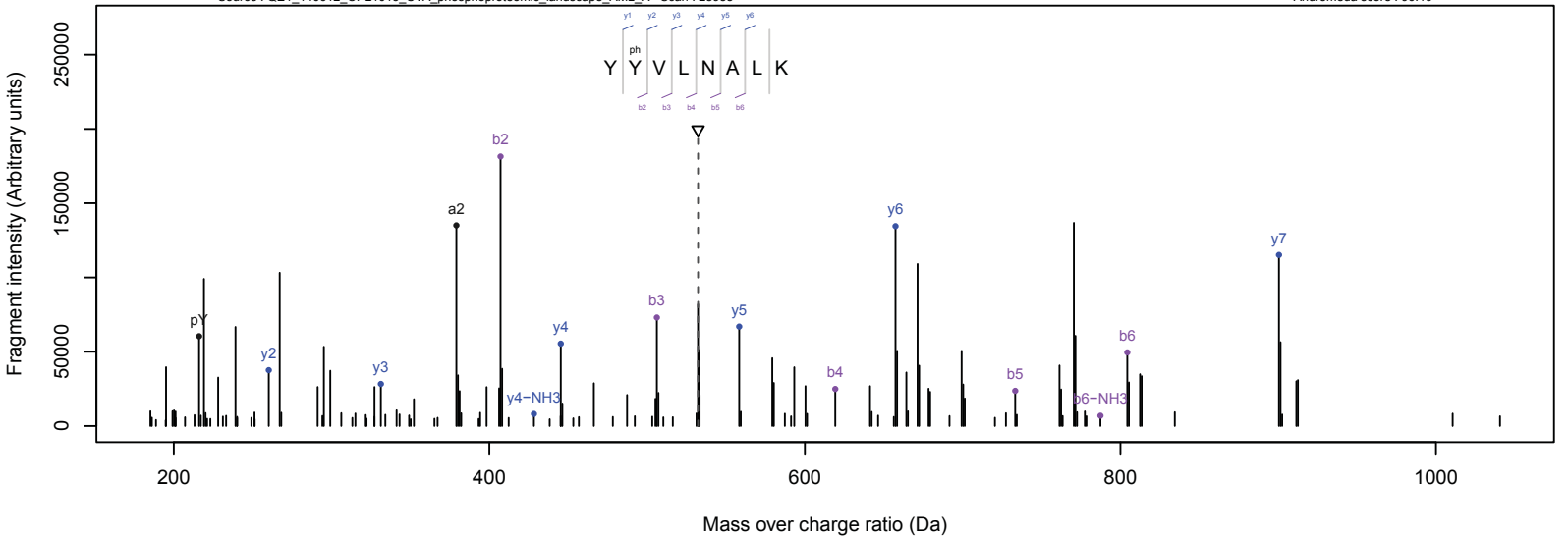
Andromeda score : 96.208



YY(ph)VLNALK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 28988

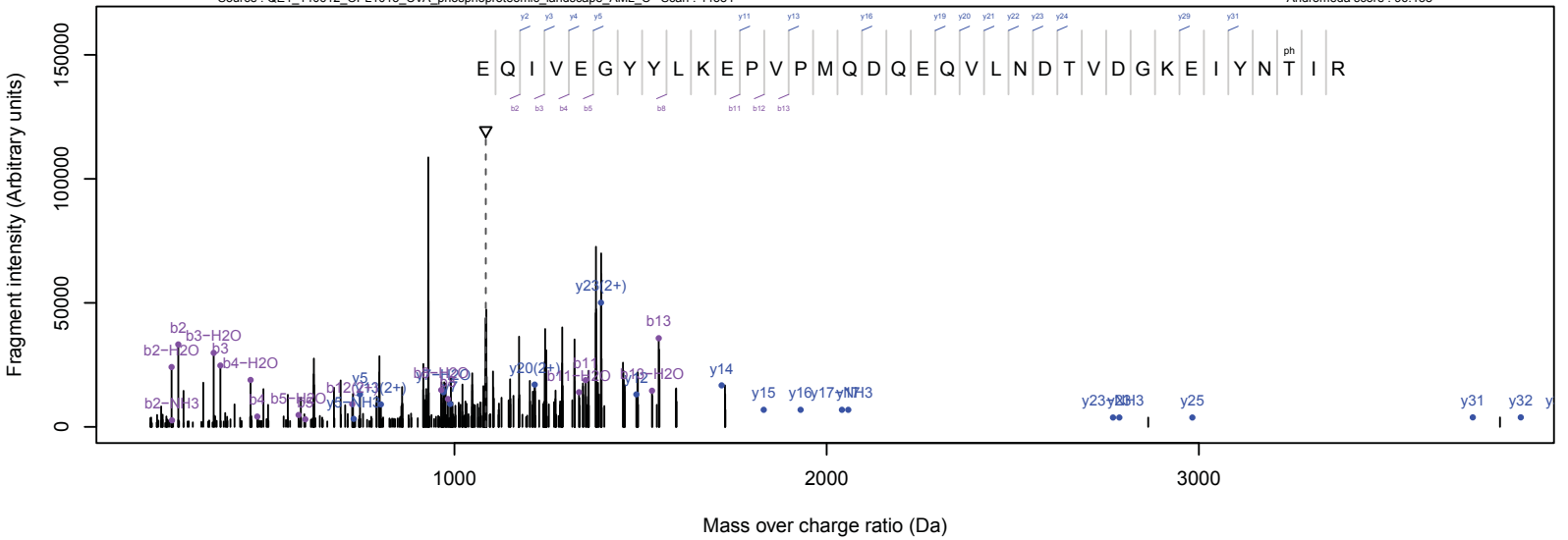
Andromeda score : 96.19



EQIVEGYLKEPVPMDQEQVLNDTVDGKEIYNT(ph)IR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44384

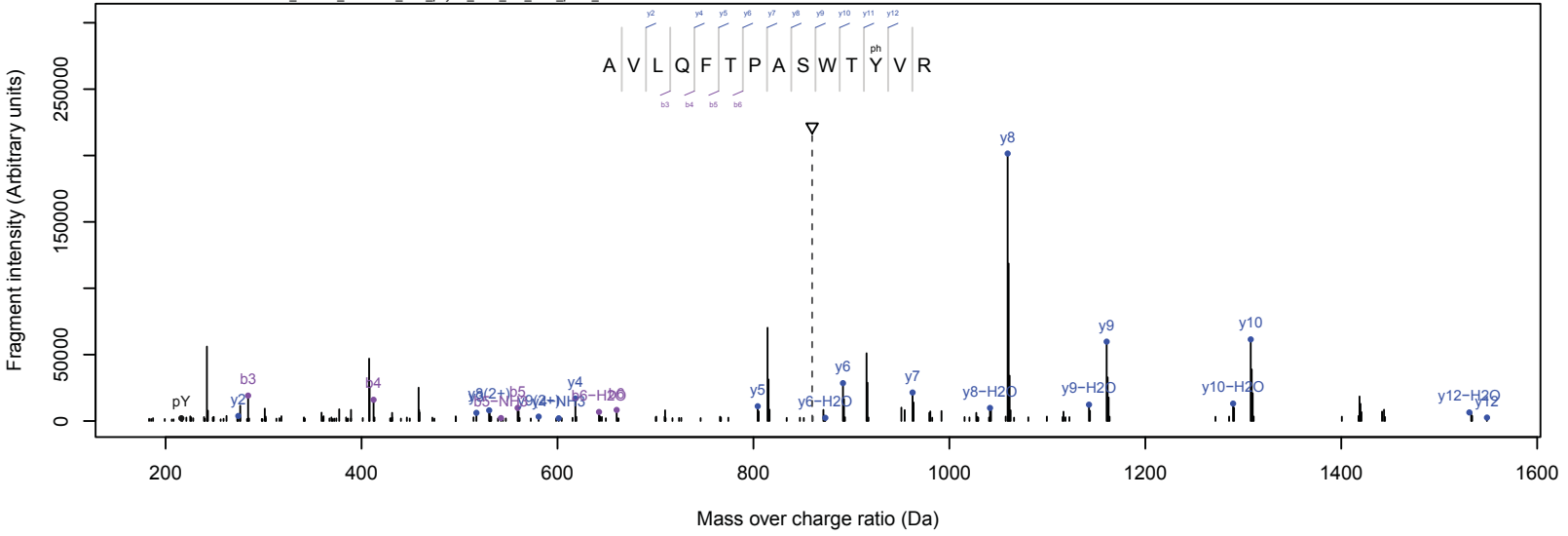
Andromeda score : 96.158



AVLQFTPASWTY(ph)VR-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 36372

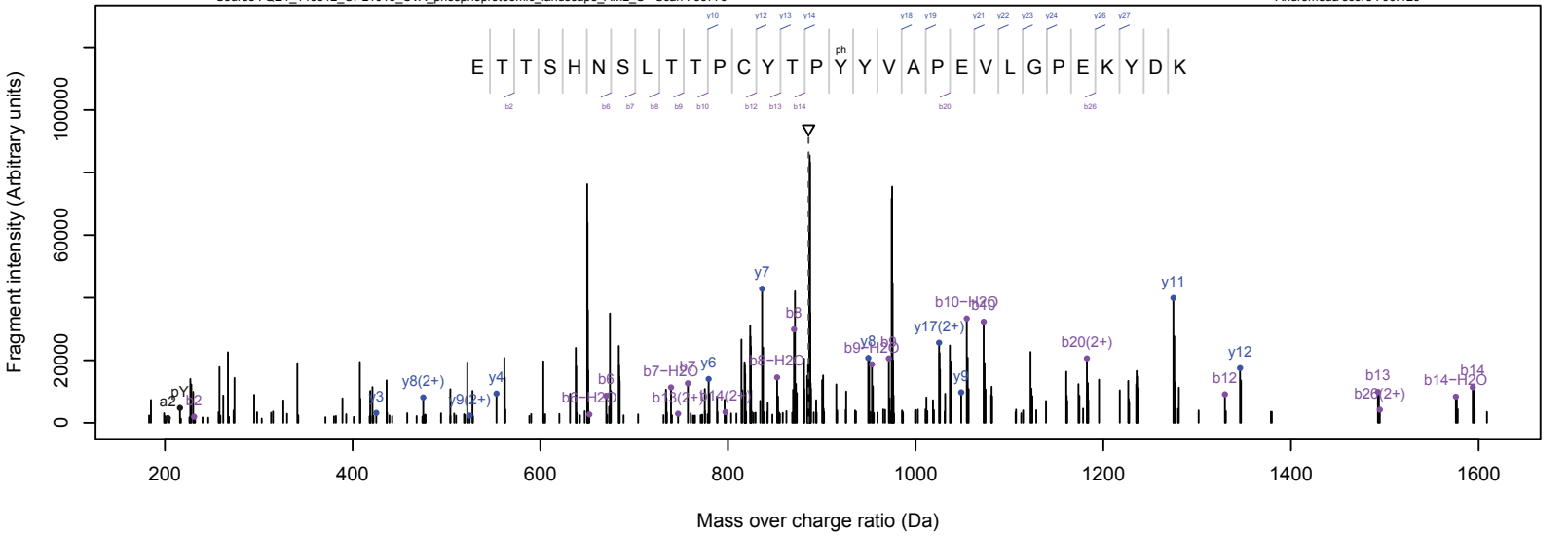
Andromeda score : 96.135



ETTSHNSLTTPCYTPYVVAPEVLGPEKYDK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35779

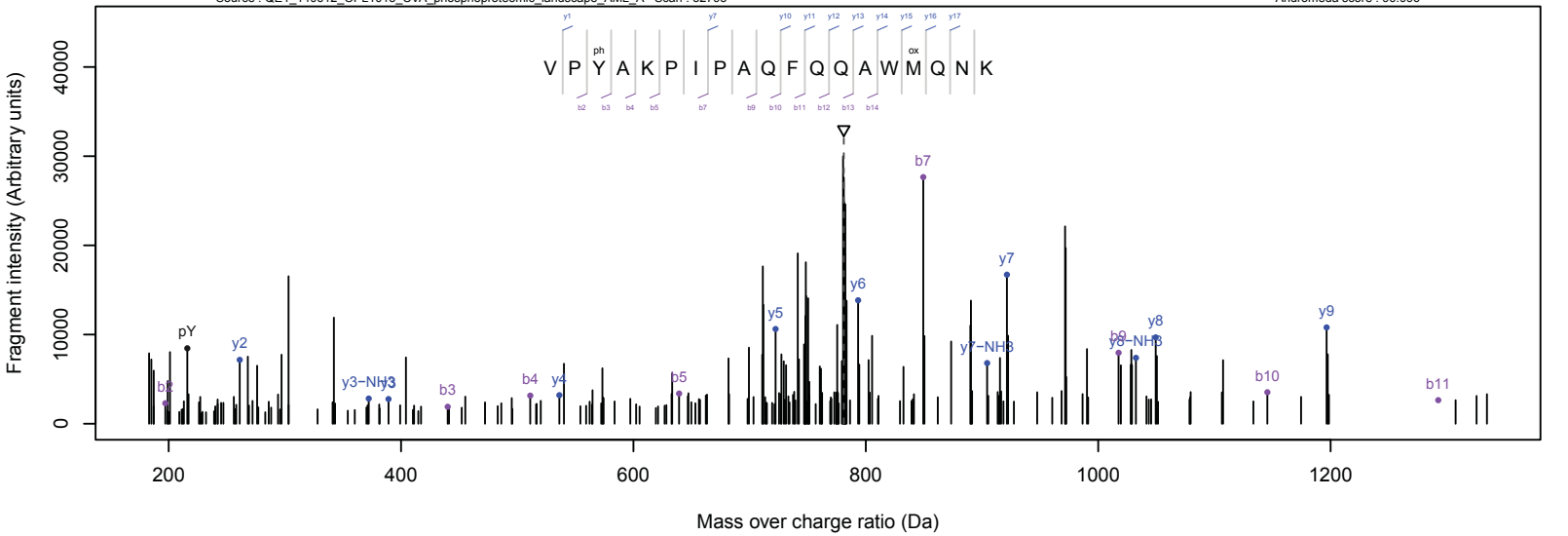
Andromeda score : 96.123



VPY(ph)AKPIPAQFQQAWM(ox)QNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32795

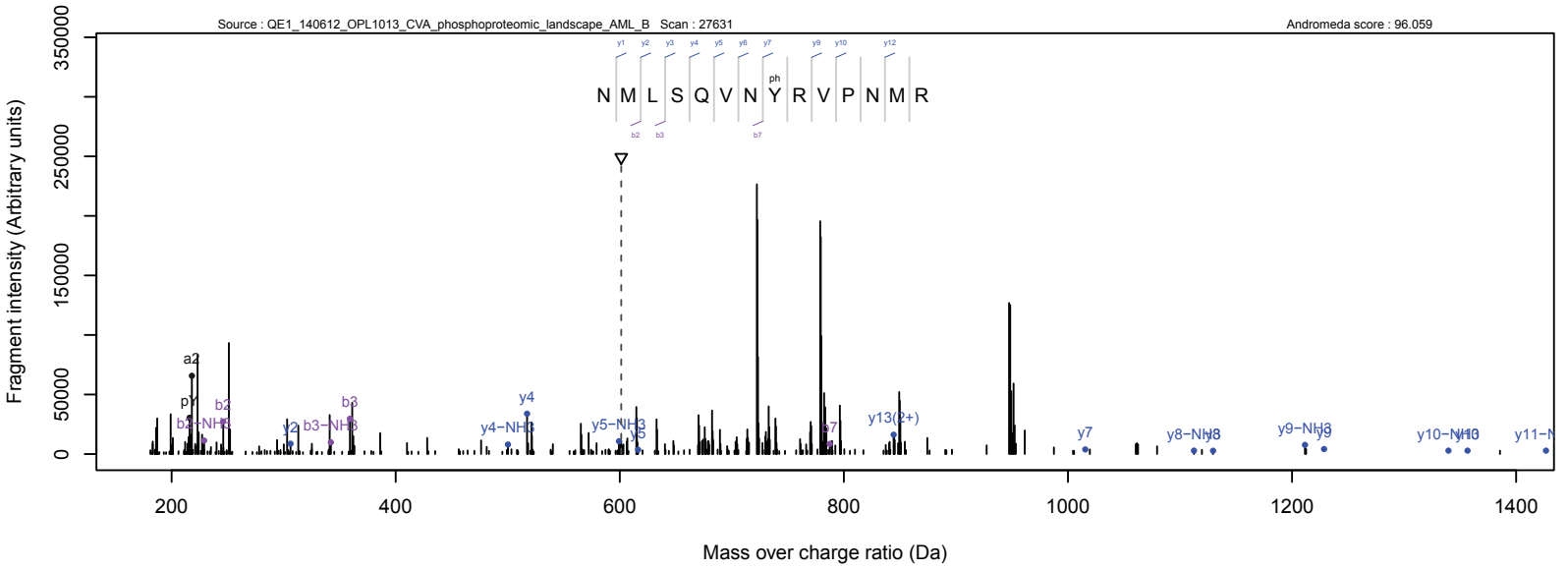
Andromeda score : 96.096



NMLSQVNY(ph)RVPNMR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27631

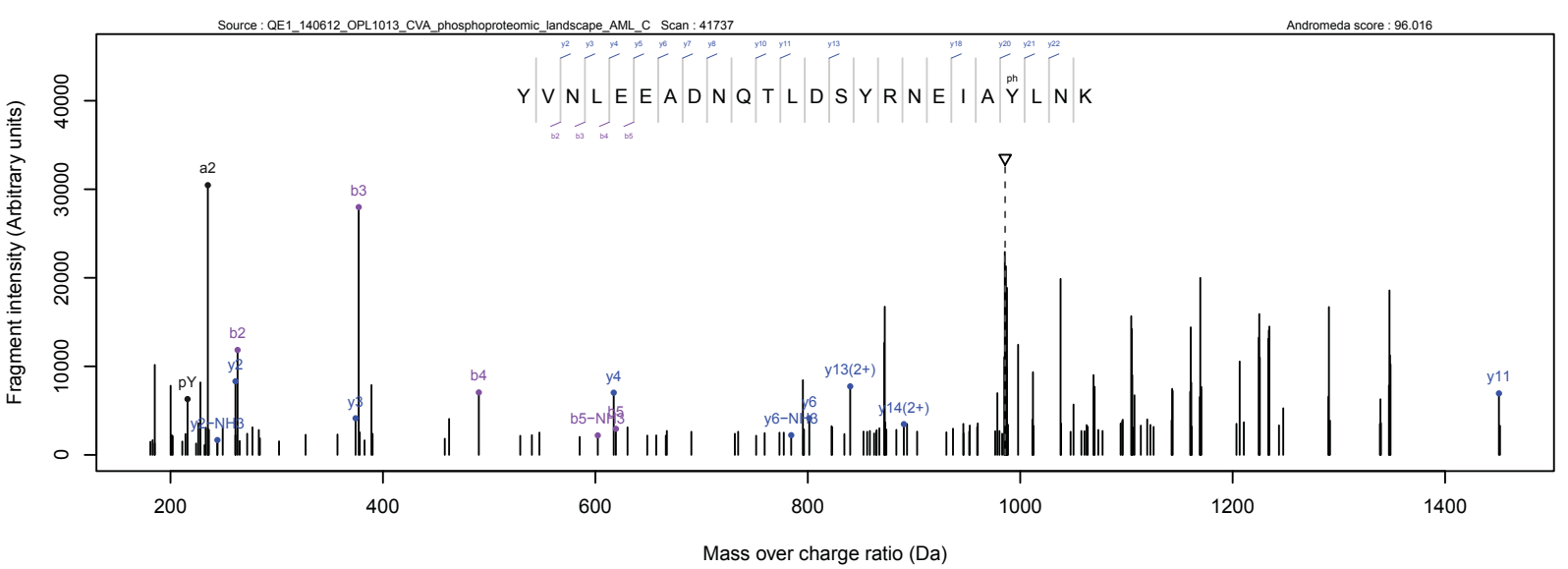
Andromeda score: 96.059



YVNLEEDNQTLDSYRNEIAY(ph)LNK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41737

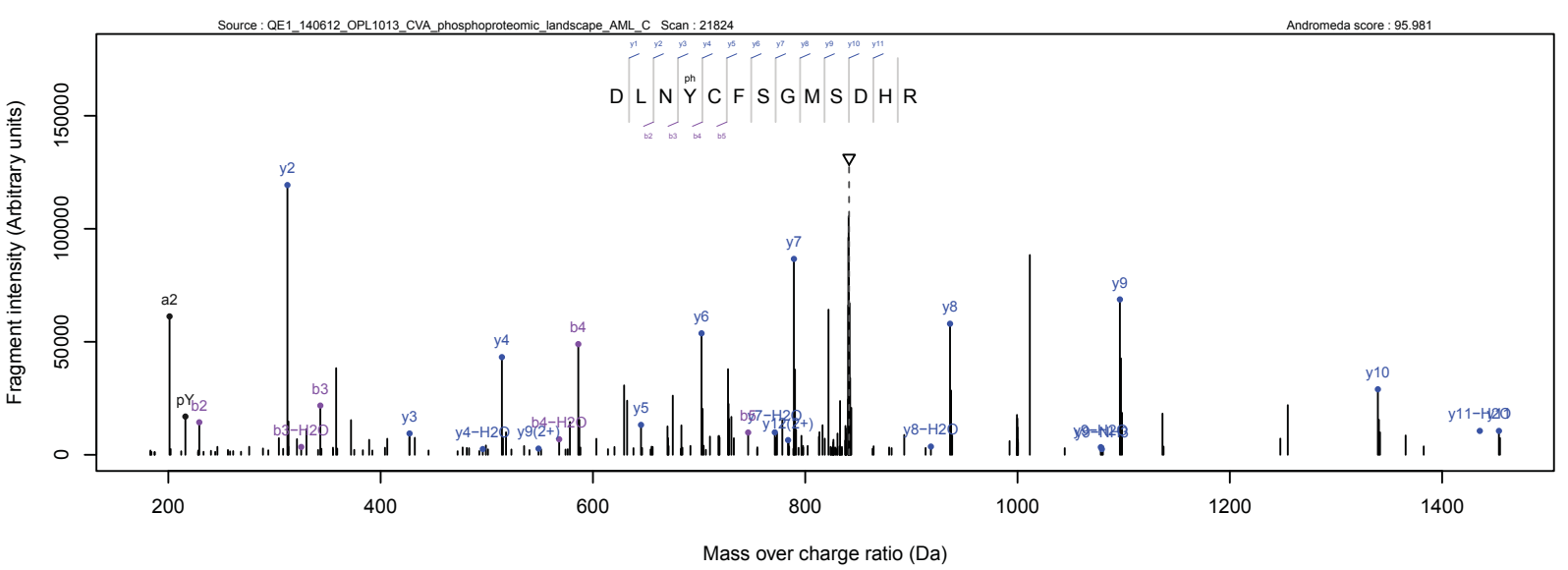
Andromeda score: 96.016



DLNY(ph)CFSGMSDHR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21824

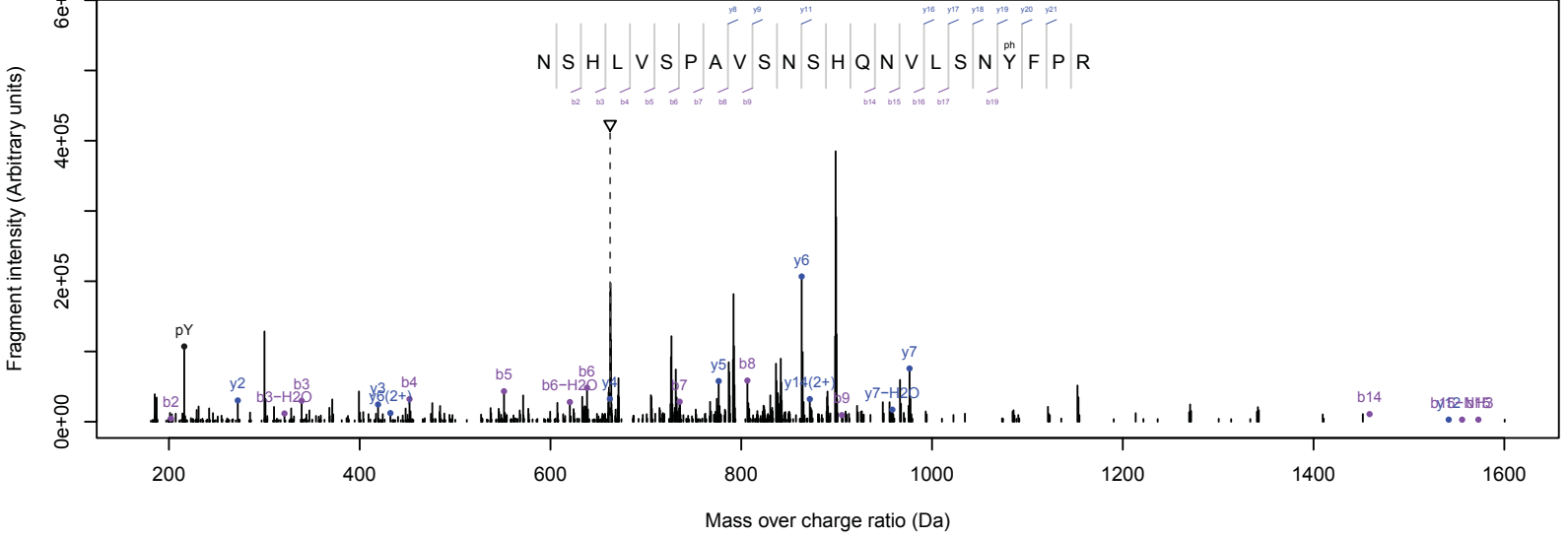
Andromeda score: 95.981



**NSHLVSPAVSNSHQNVLSNYPFR-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 28420

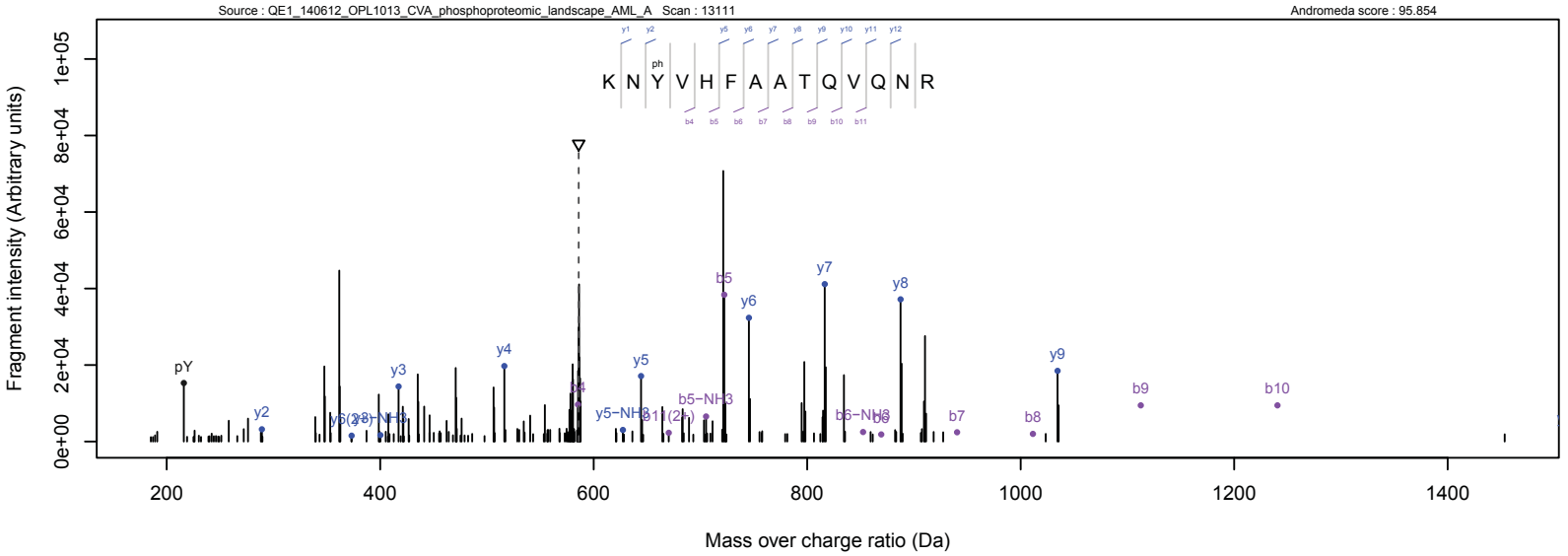
Andromeda score: 95.926



**KNYVHFAATQVQNR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 13111

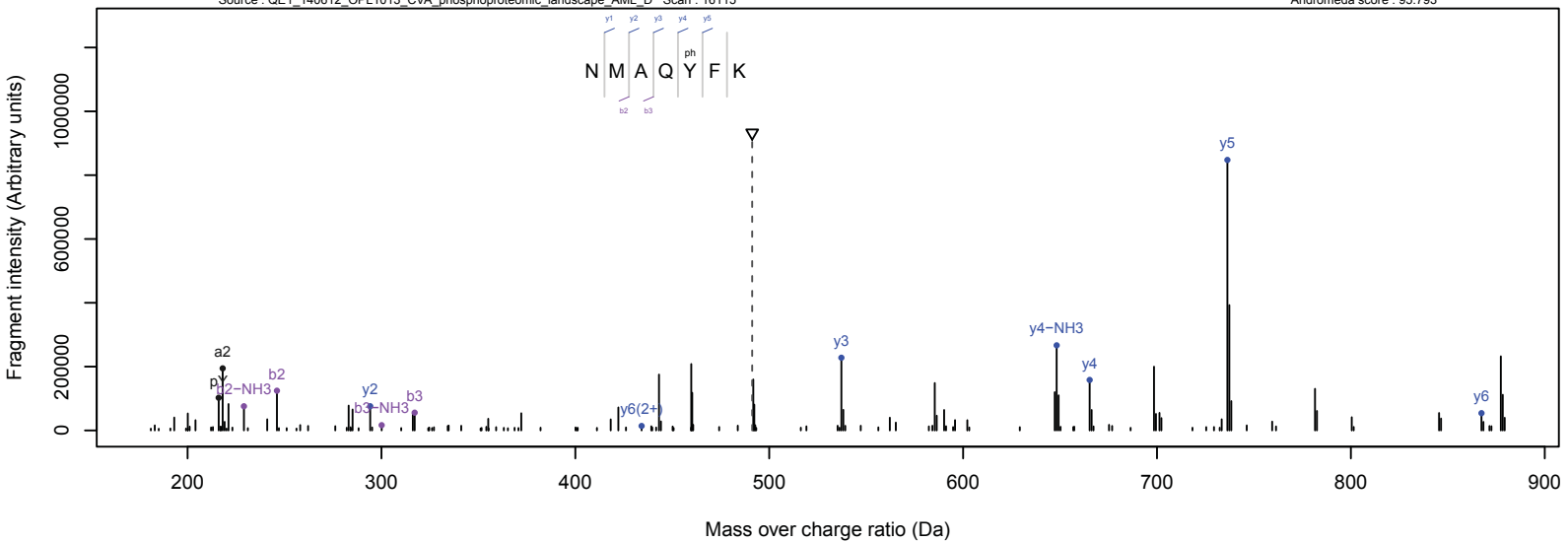
Andromeda score: 95.854



**NMAQYFK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 16115

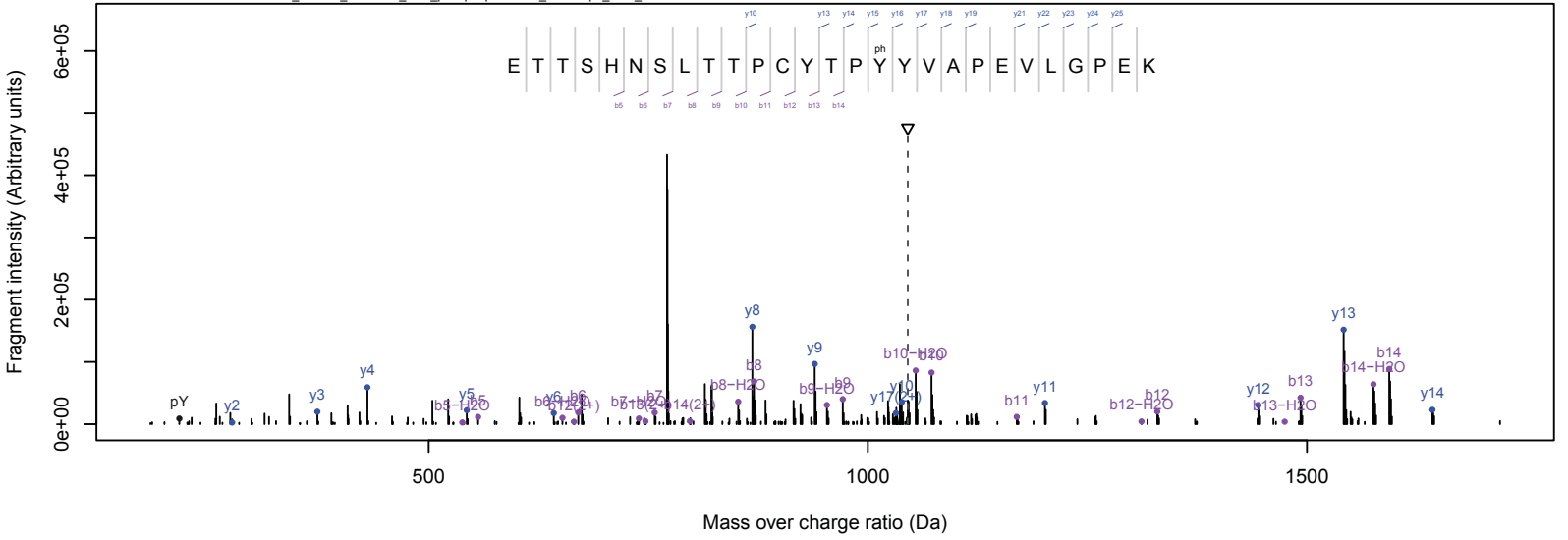
Andromeda score: 95.793



ETTSHNSLTPCYTPY(ph)YVAPEVLGPEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37387

Andromeda score : 95.652

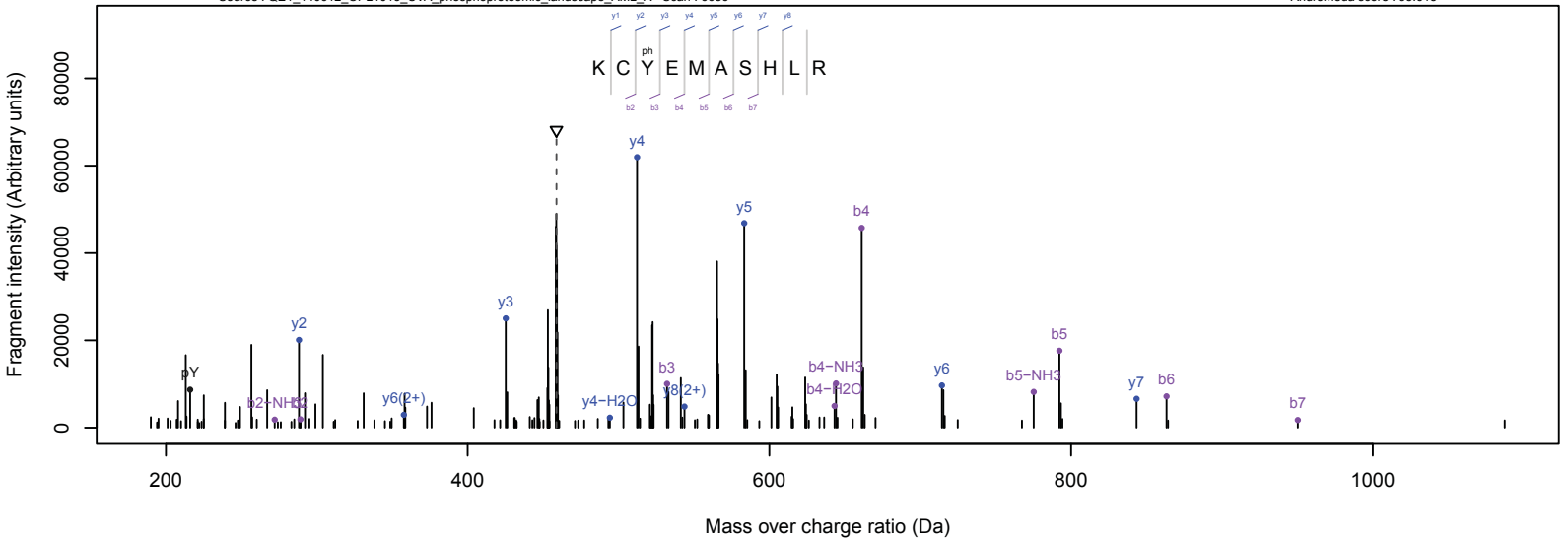




KCY(ph)EMASHLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 6686

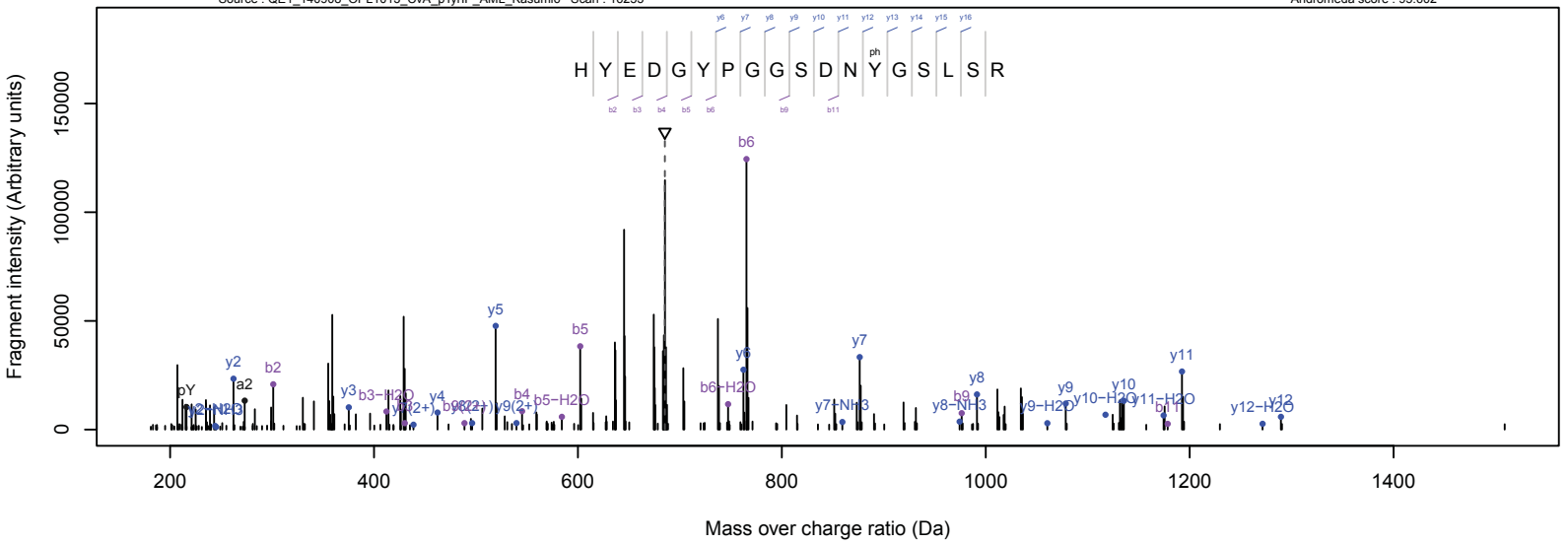
Andromeda score : 95.618



HYEDGYPGSDNY(ph)GSLSR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 16253

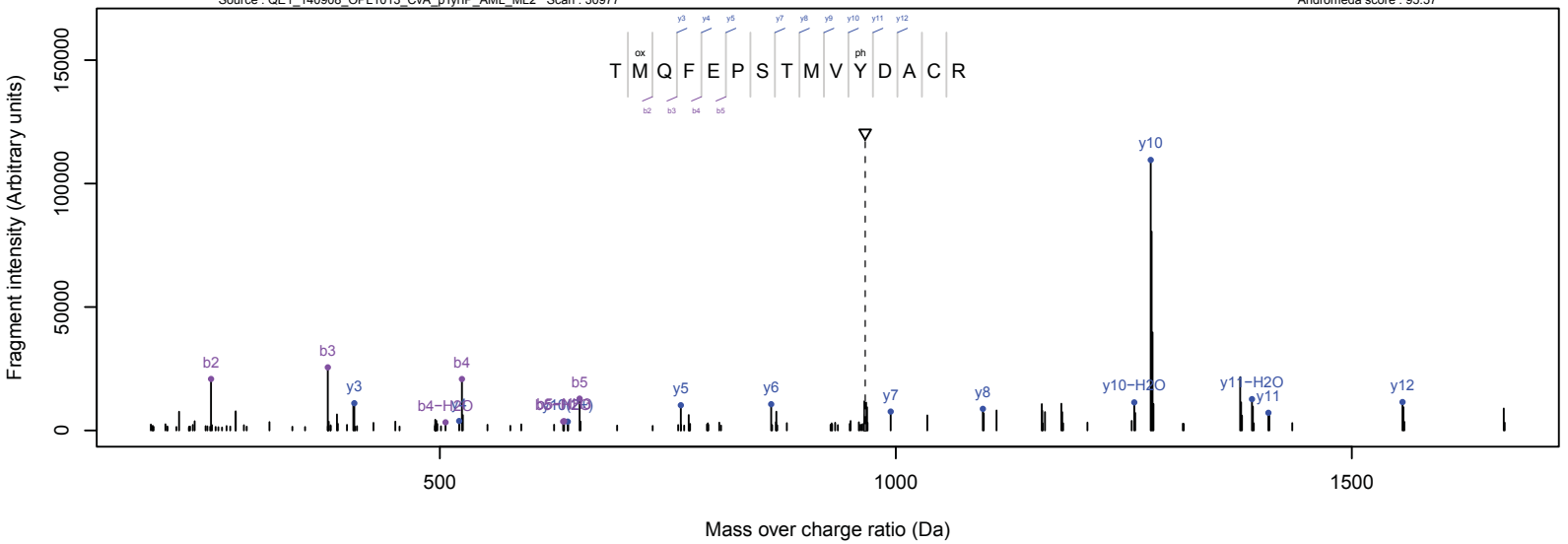
Andromeda score : 95.602



TM(ox)QFEPSTMVY(ph)DACR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 30977

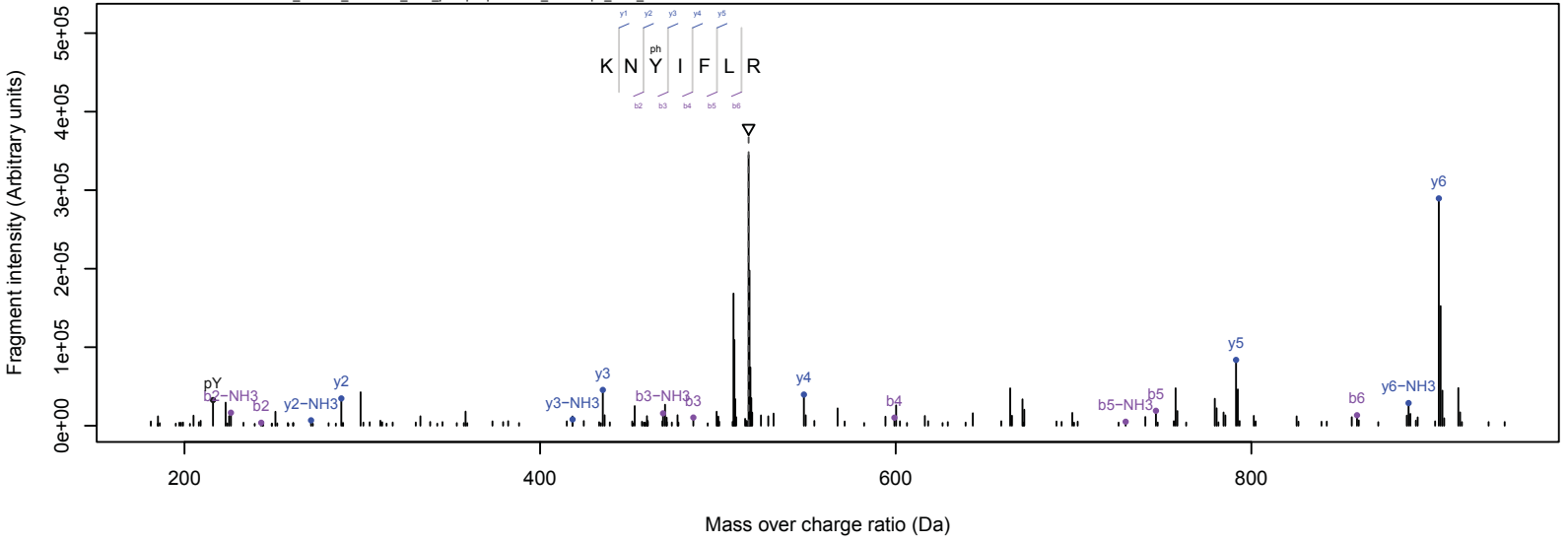
Andromeda score : 95.57



KNY(ph)IFLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 19673

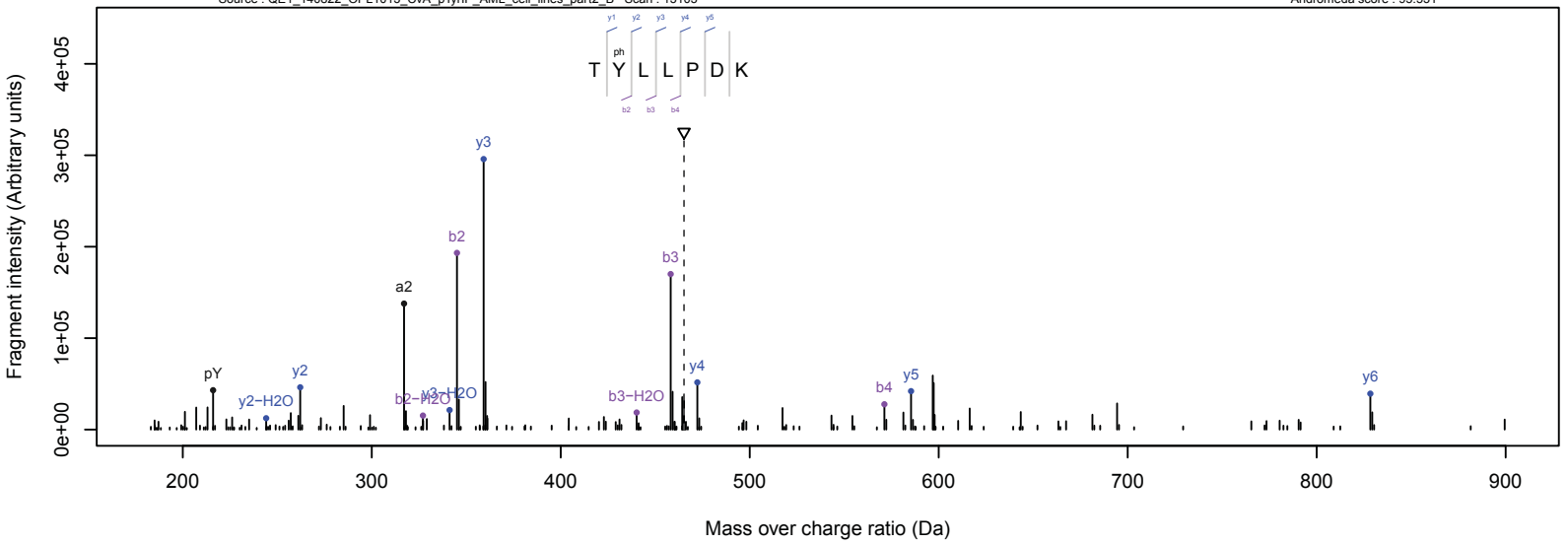
Andromeda score: 95.541



TY(ph)LLPDK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 13103

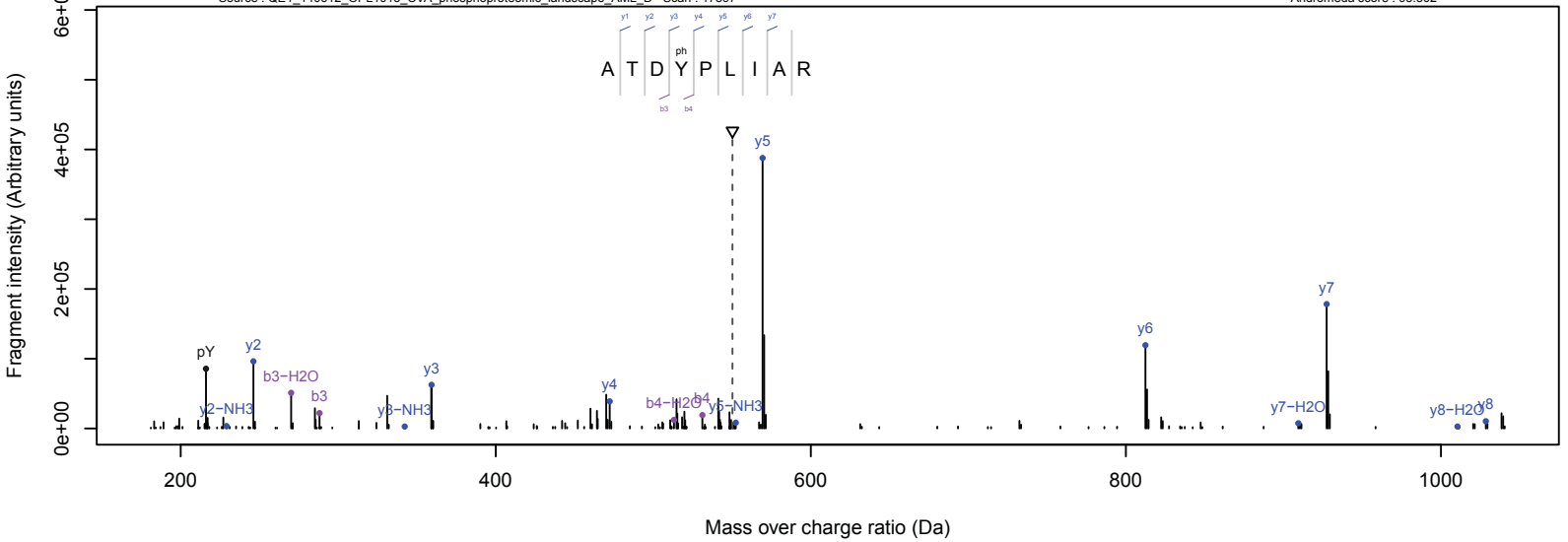
Andromeda score: 95.531



ATDY(ph)PLIAR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 17397

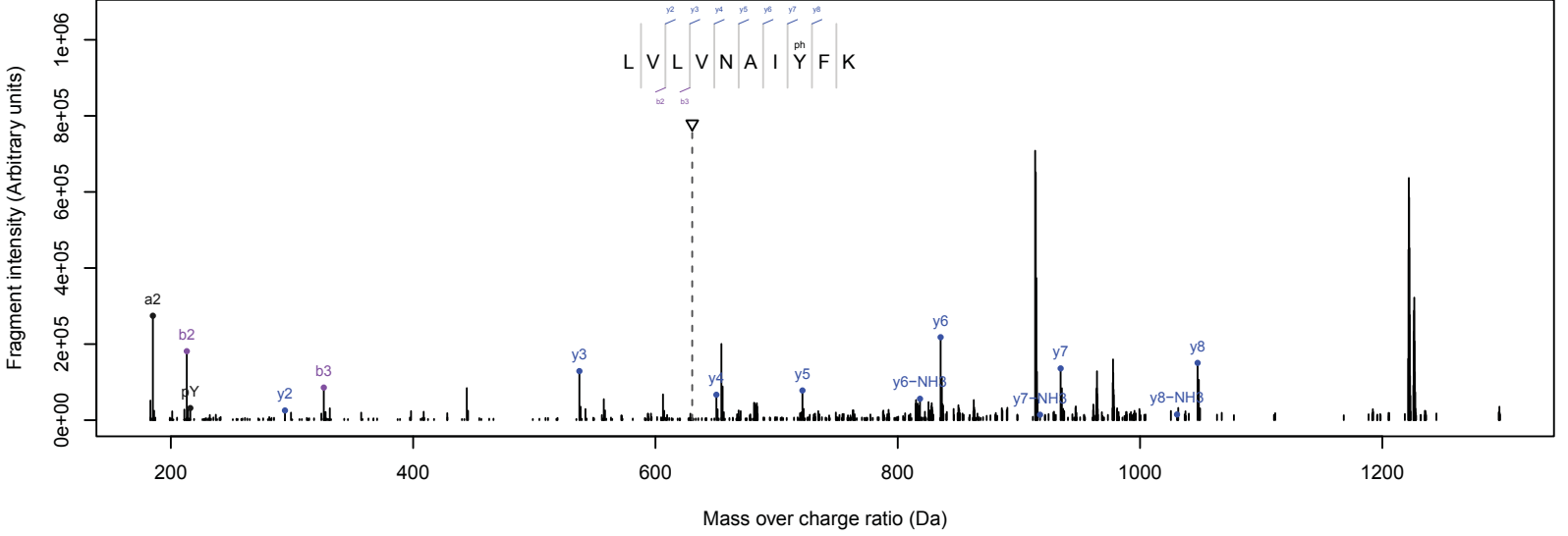
Andromeda score: 95.502



LVLVNAIY(ph)FK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44993

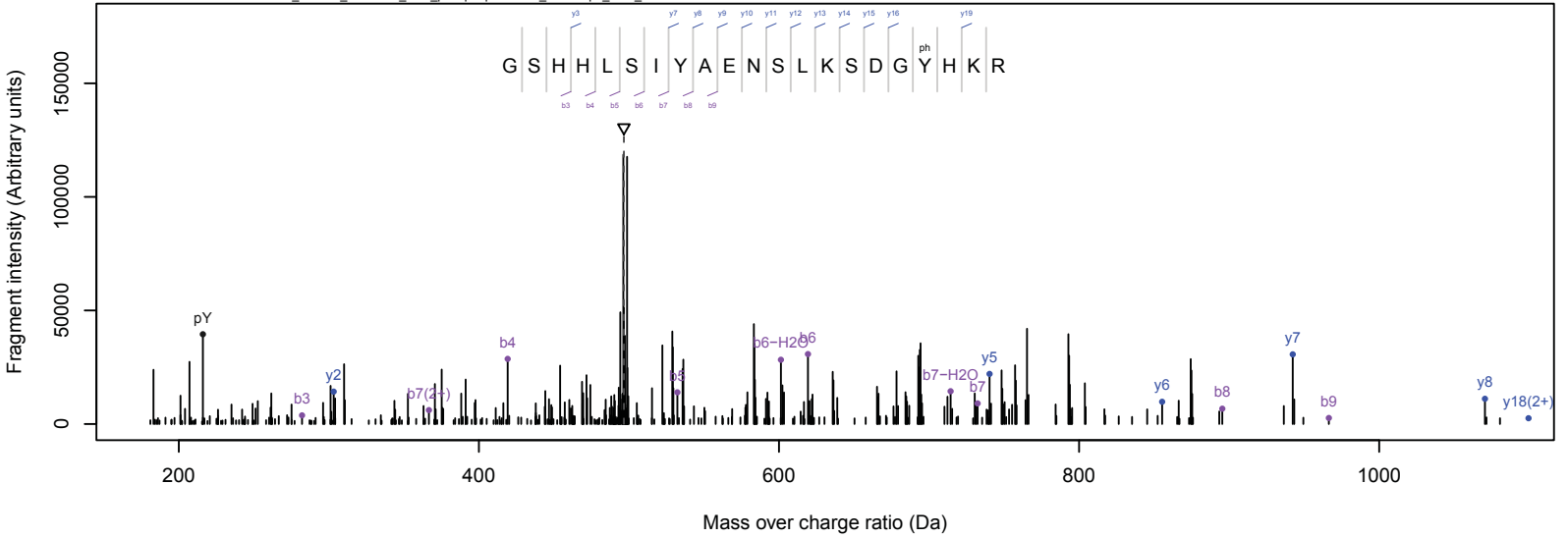
Andromeda score : 95.502



GSHLSIYAENSLKSDGY(ph)HKR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13404

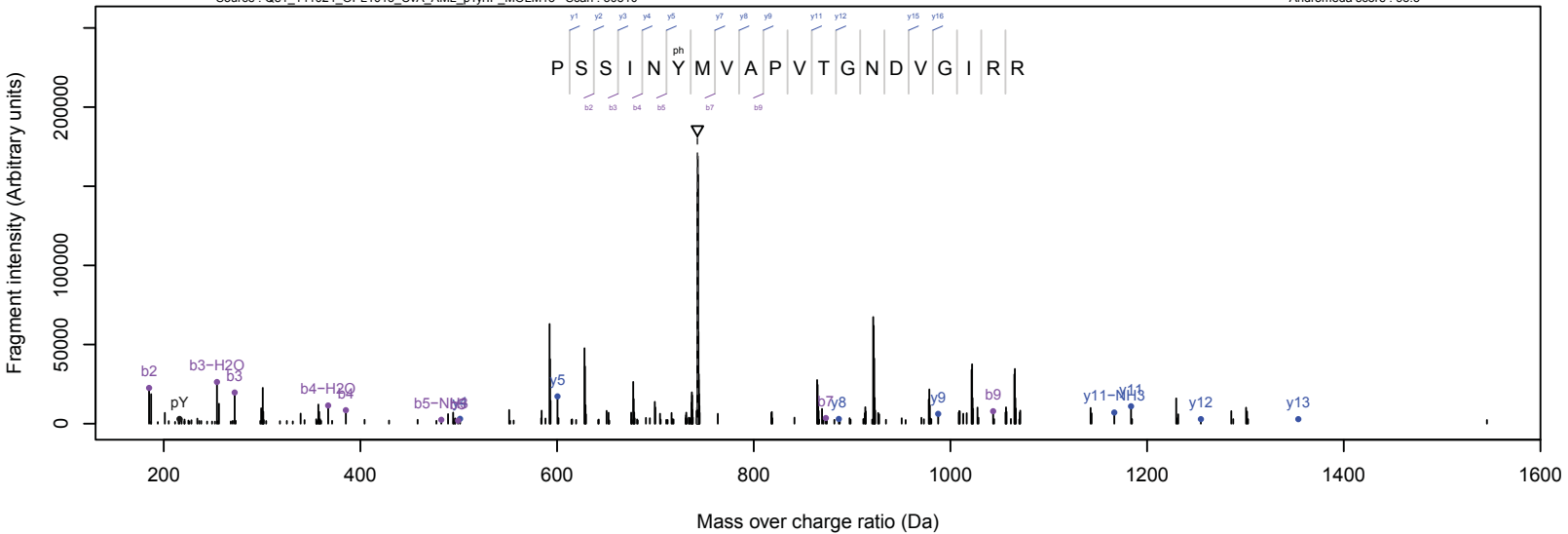
Andromeda score : 95.5



PSSINY(ph)MVA PVTGNDVGIR-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 30510

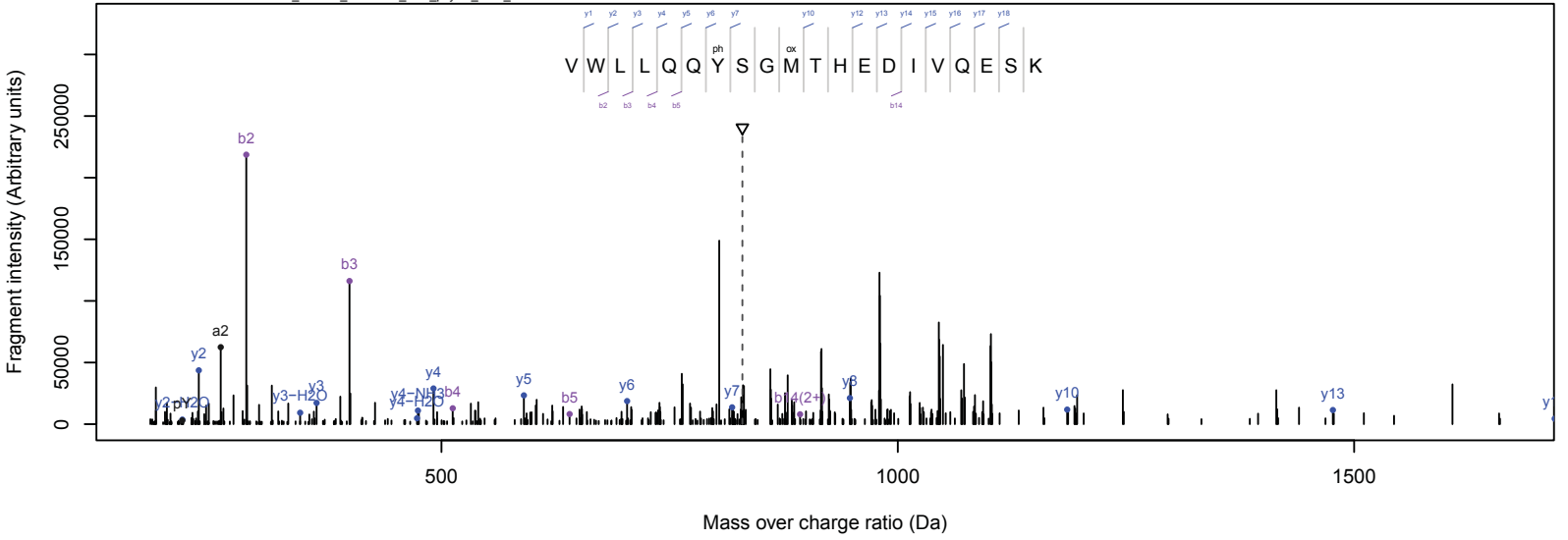
Andromeda score : 95.5



VWLLQY(ph)SGM(ox)THEDIVQESK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 36911

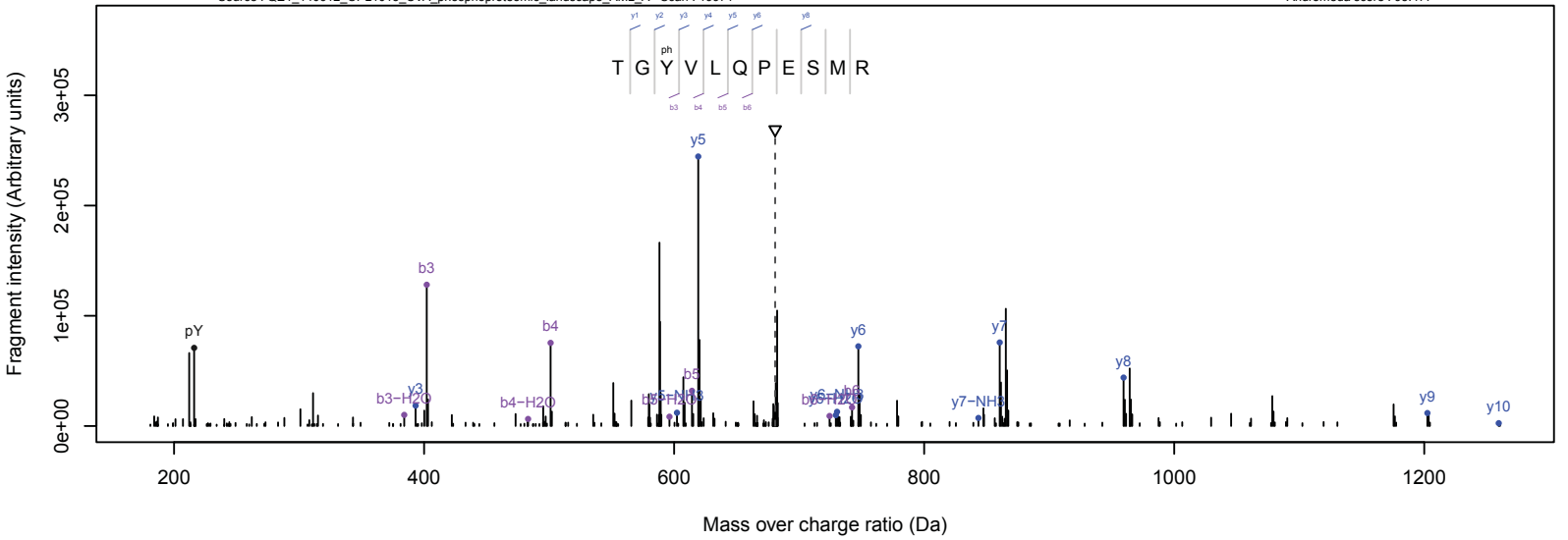
Andromeda score : 95.5



TGY(ph)VLQPESMR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 19971

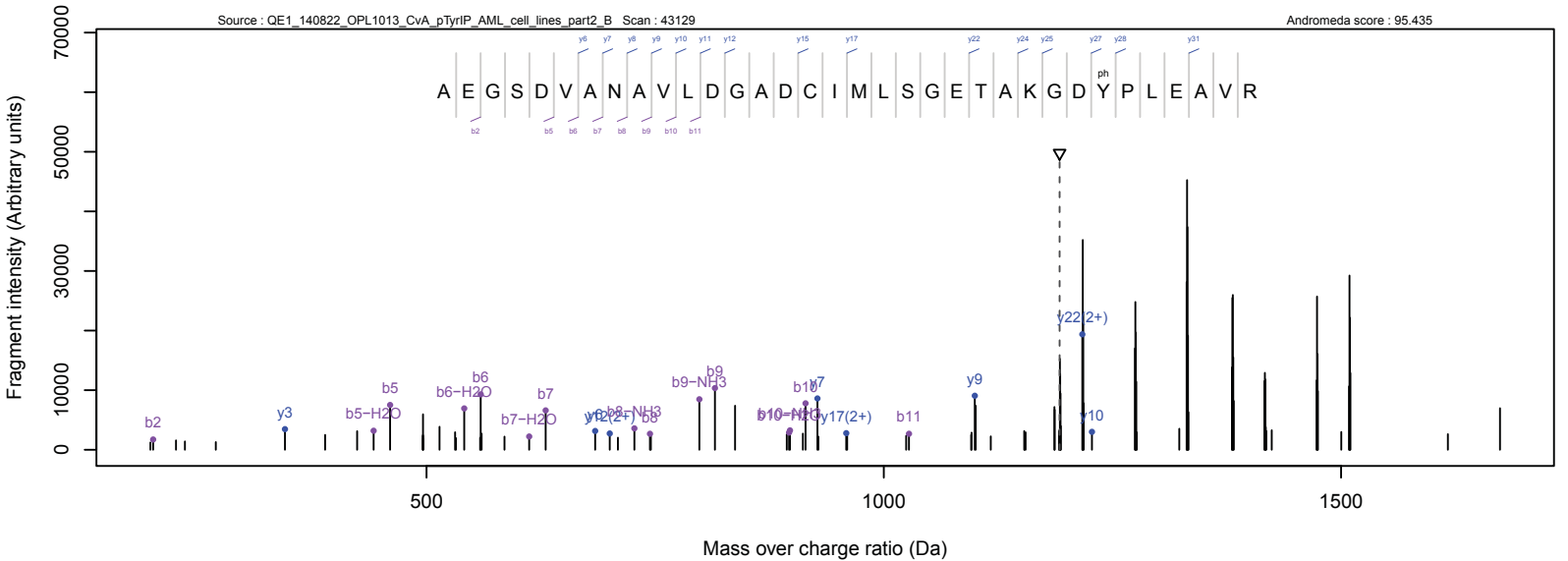
Andromeda score : 95.477



AEGSDVANAVLDGADCIMLSGETAKGDY(ph)PLEAVR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 43129

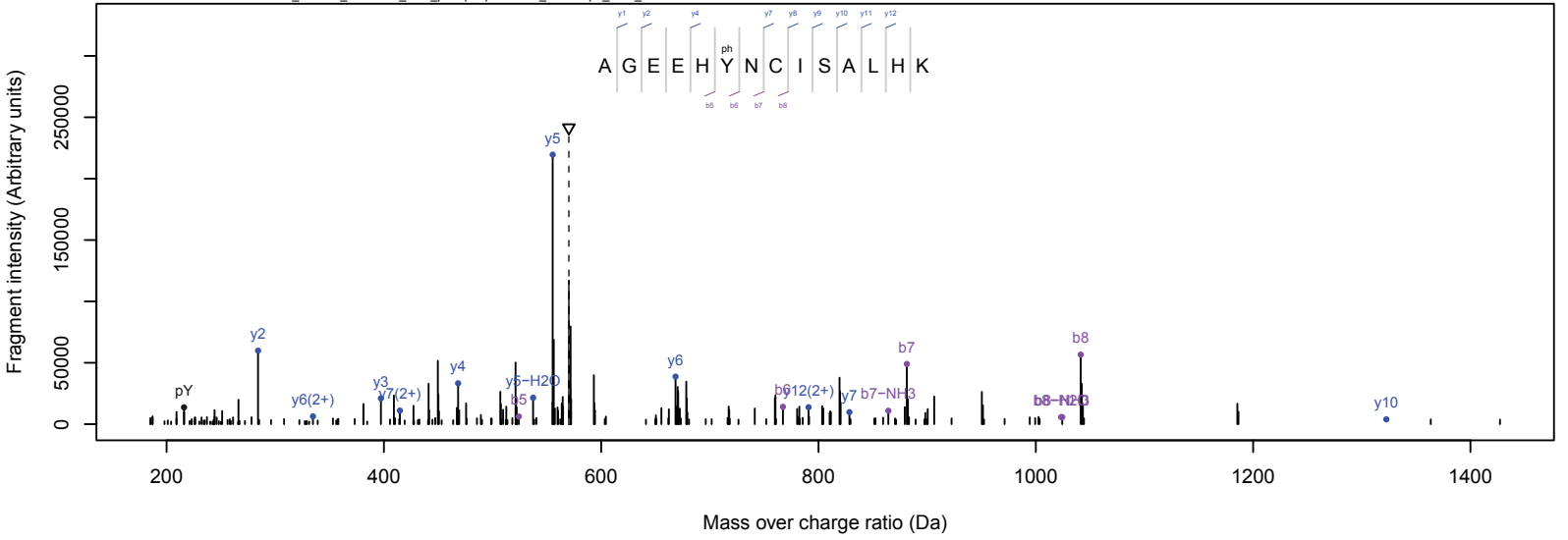
Andromeda score : 95.435



AGEEHY(ph)NCISALHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 11175

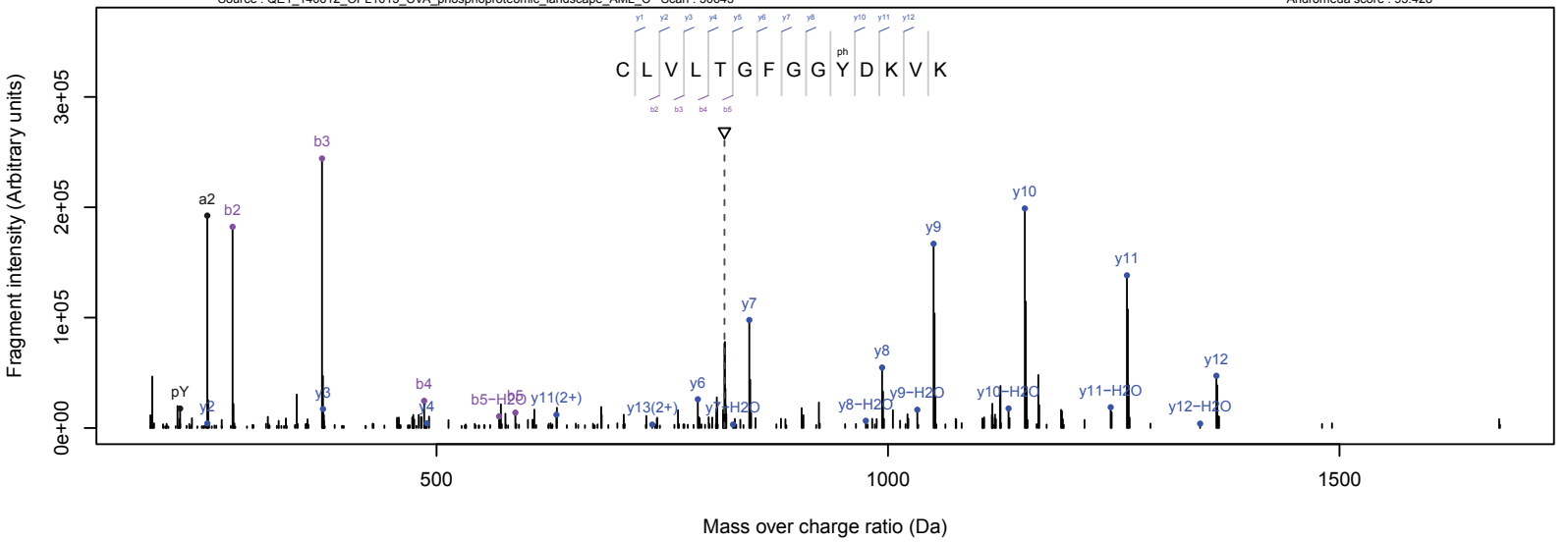
Andromeda score : 95.428



CLVLTGFGGY(ph)DKVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30645

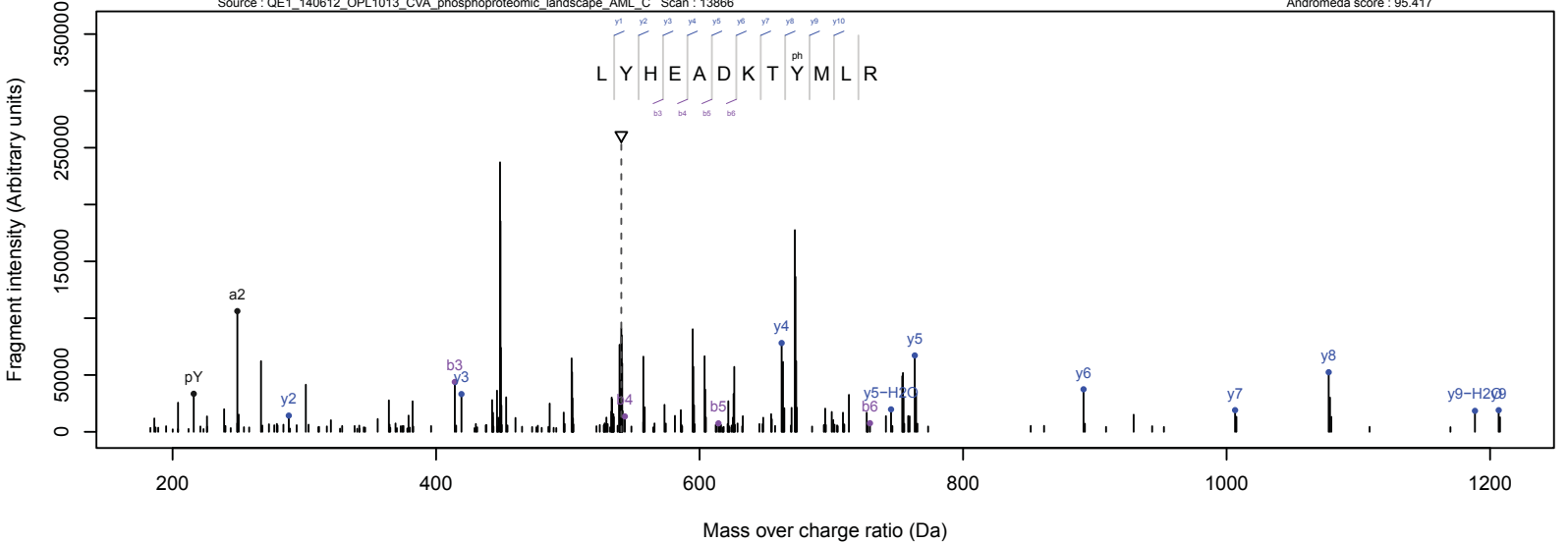
Andromeda score : 95.428



LYHEADKTY(ph)MLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13866

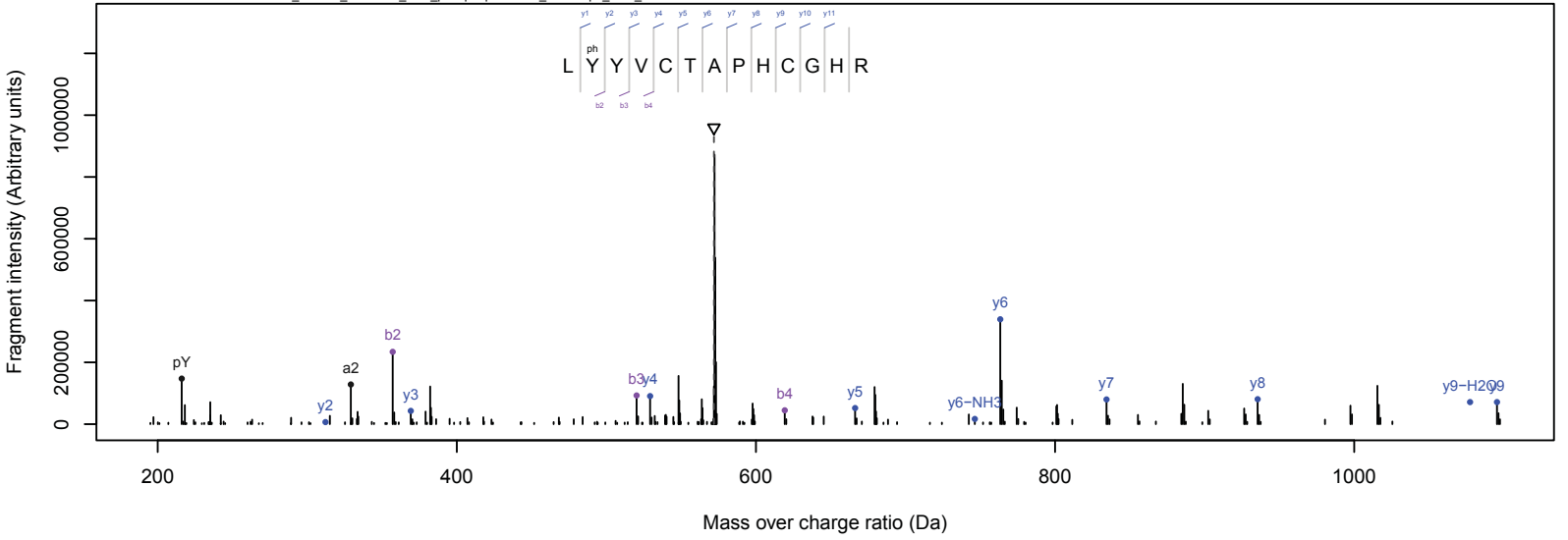
Andromeda score : 95.417



LY(ph)YVCTAPHCGHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11281

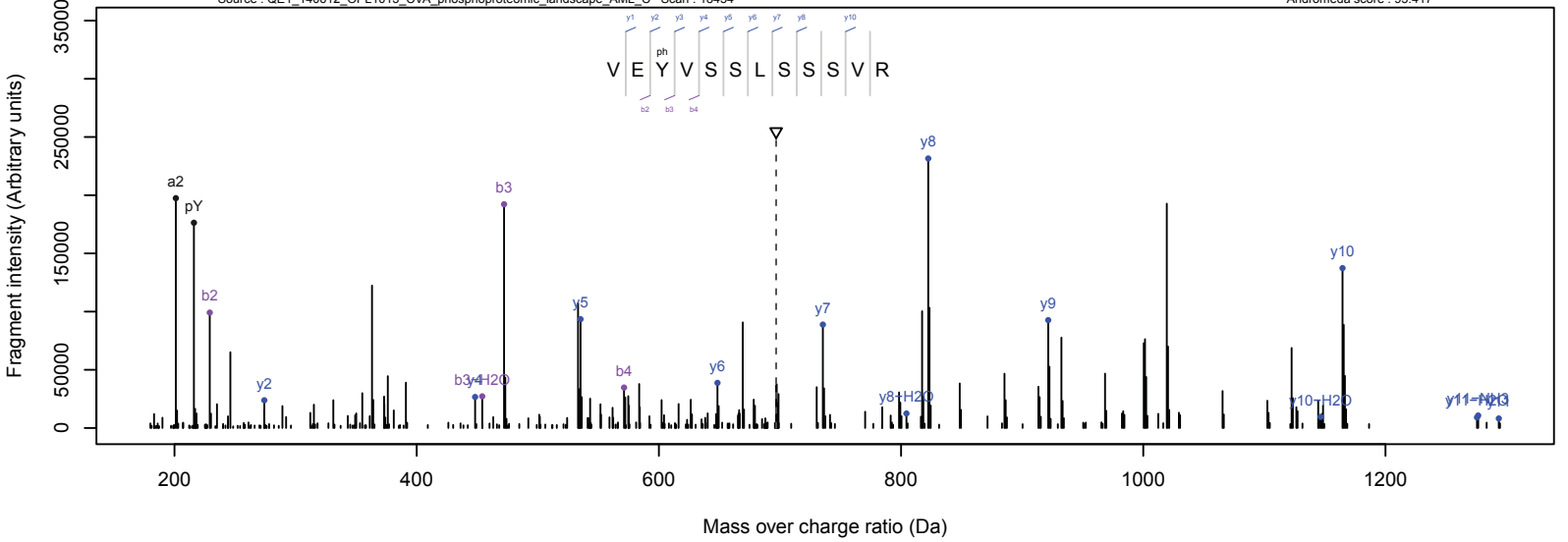
Andromeda score : 95.417



VEY(ph)VSSLSSVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18454

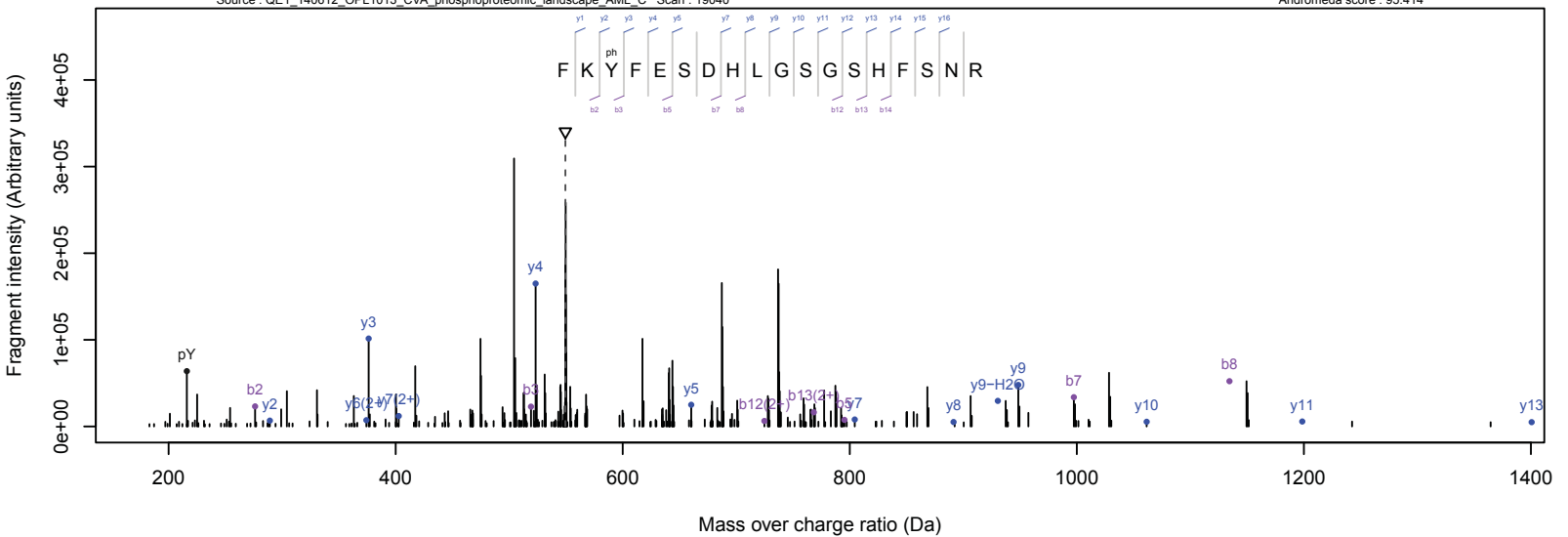
Andromeda score : 95.417



FKY(ph)FESDLGSGSHFSNR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19040

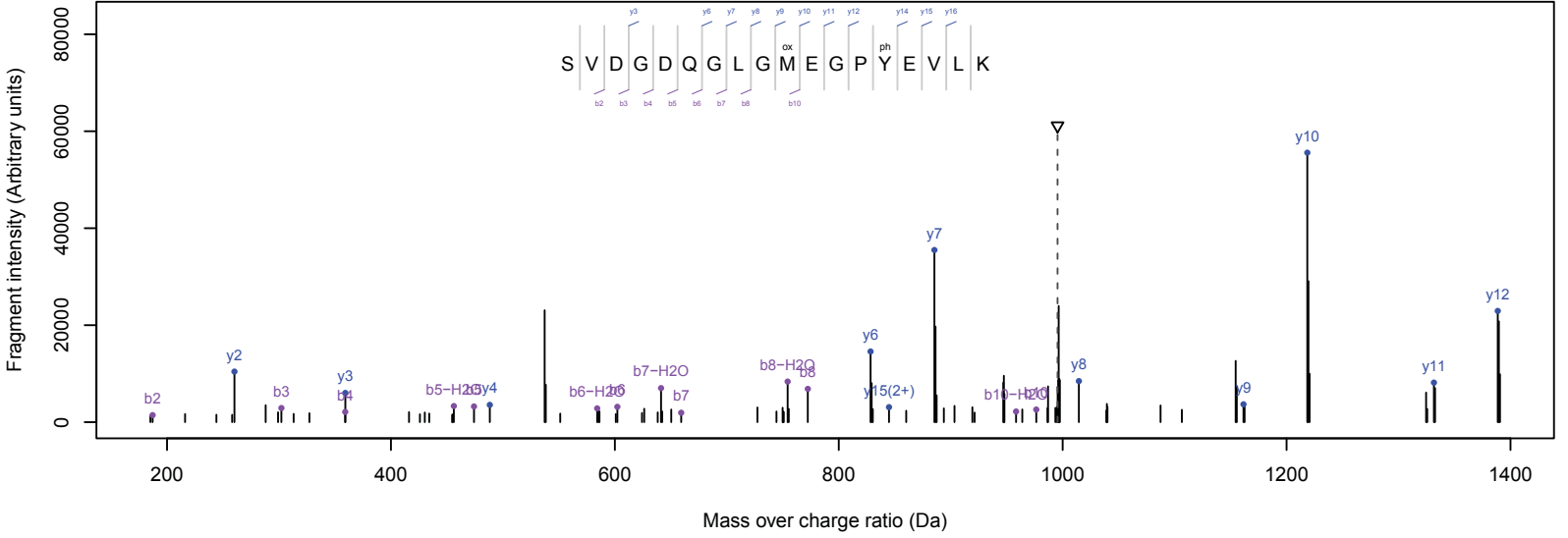
Andromeda score : 95.414



**SVDGDQGLG<sup>(ox)</sup>EGPY<sup>(ph)</sup>EVLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28191

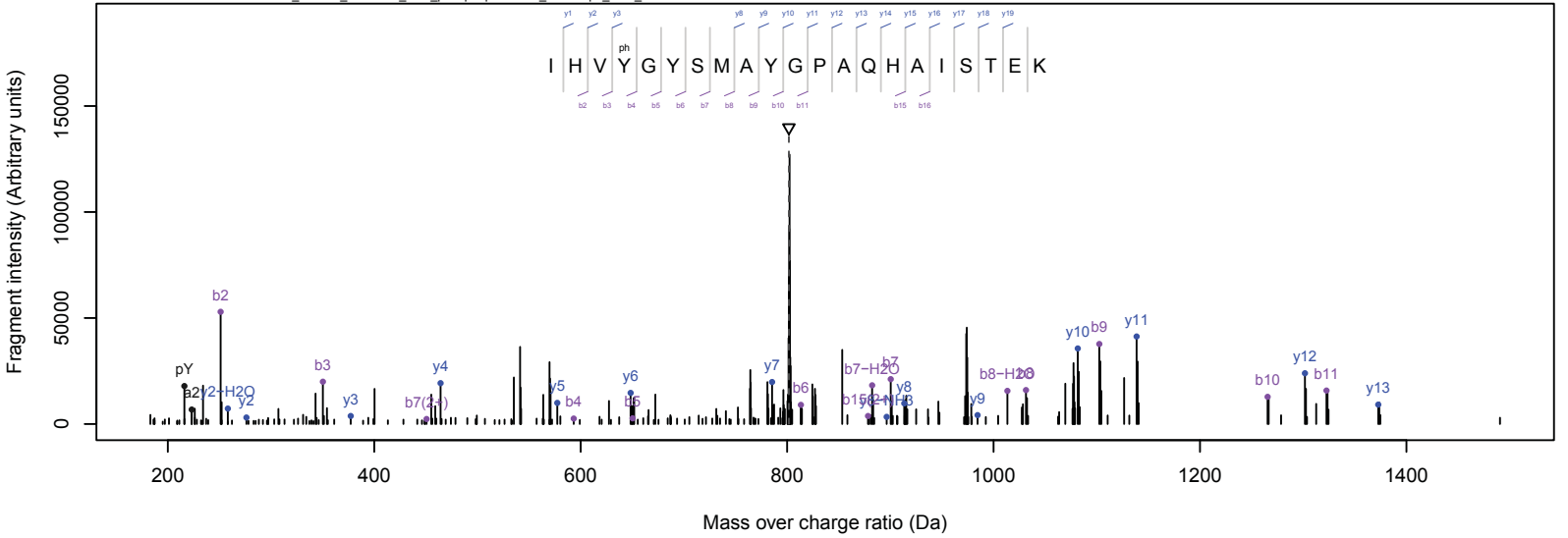
Andromeda score : 95.205



**IHVY<sup>(ph)</sup>GYSMAYGPAQHAISTEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26705

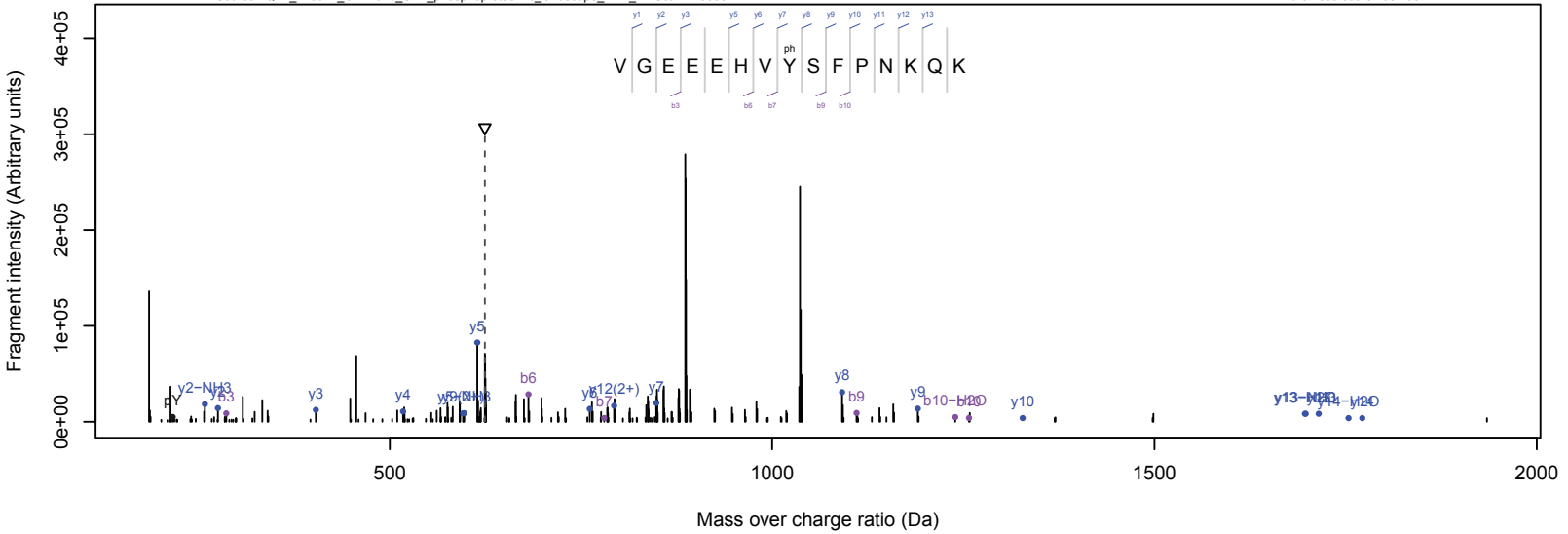
Andromeda score : 95.195



**VGEEHVY<sup>(ph)</sup>SFPNKQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 10558

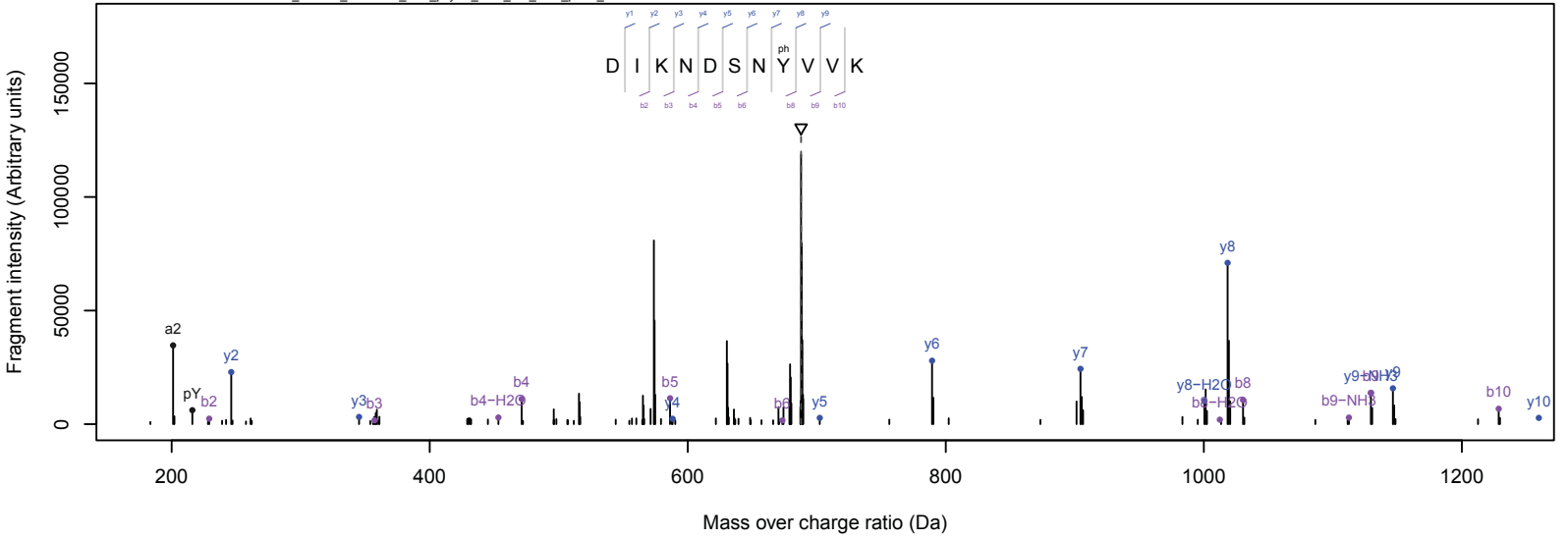
Andromeda score : 95.195



DIKNSNY(ph)VVK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 6714

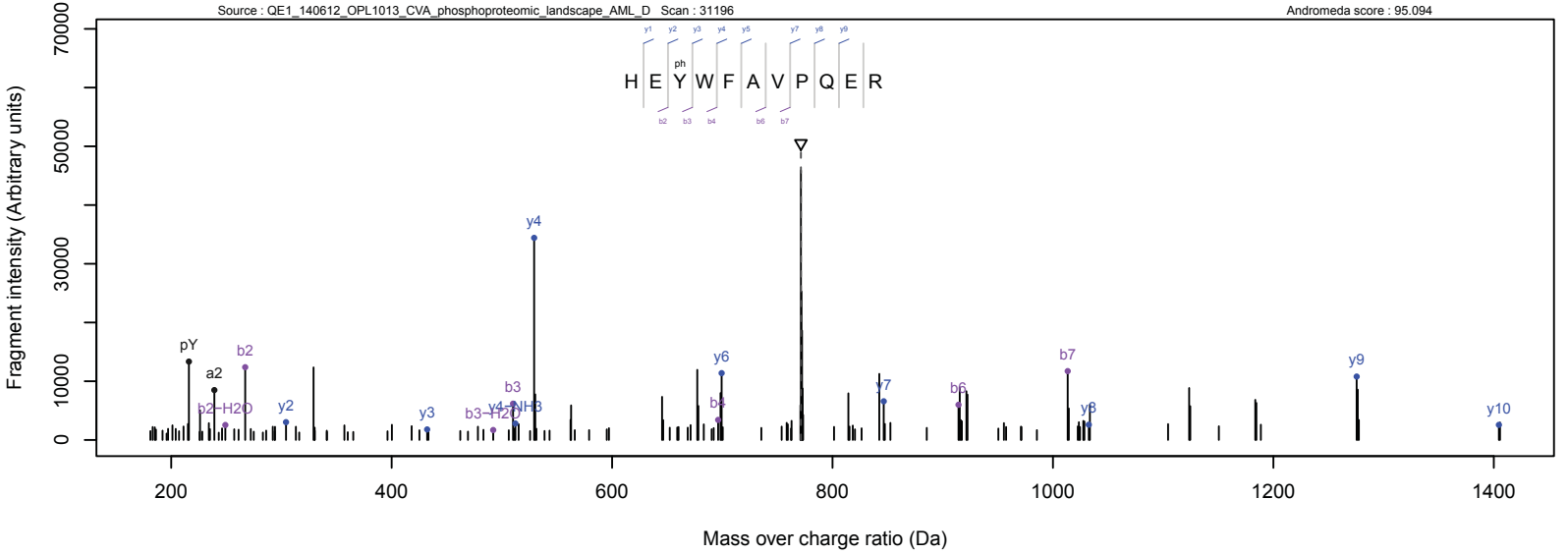
Andromeda score : 95.183



HEY(ph)WFAVPQER-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan : 31196

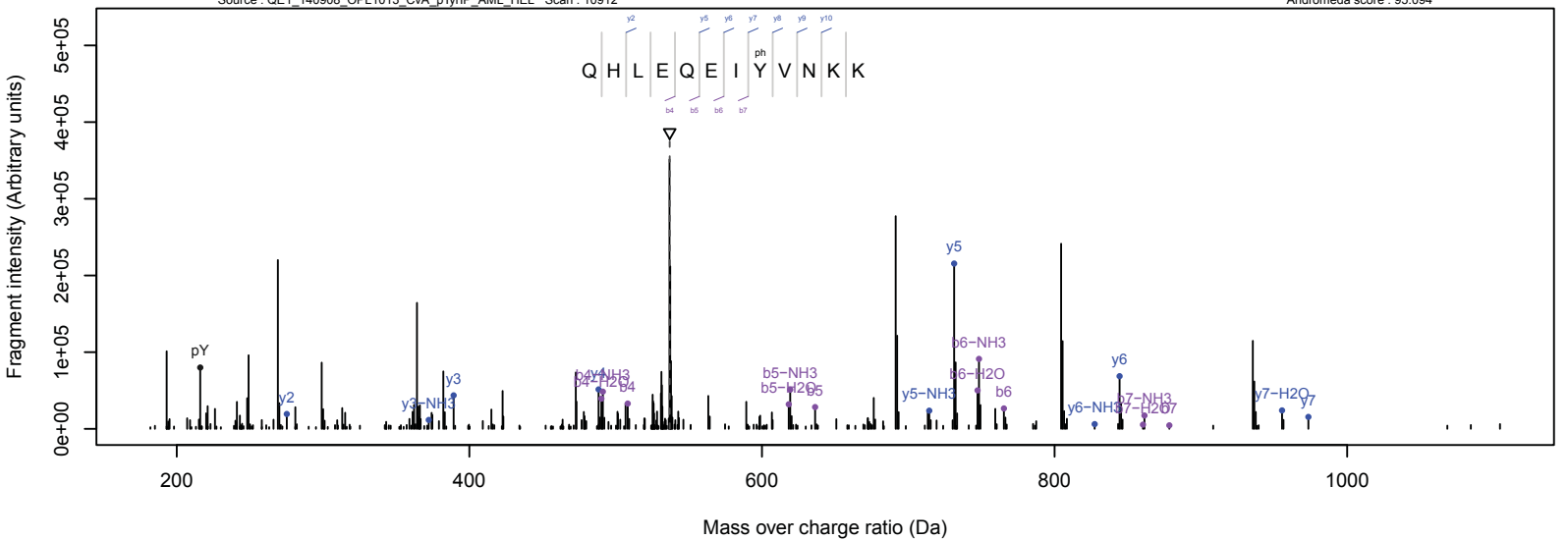
Andromeda score : 95.094



QHLEQEIYVNKK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 10912

Andromeda score : 95.094

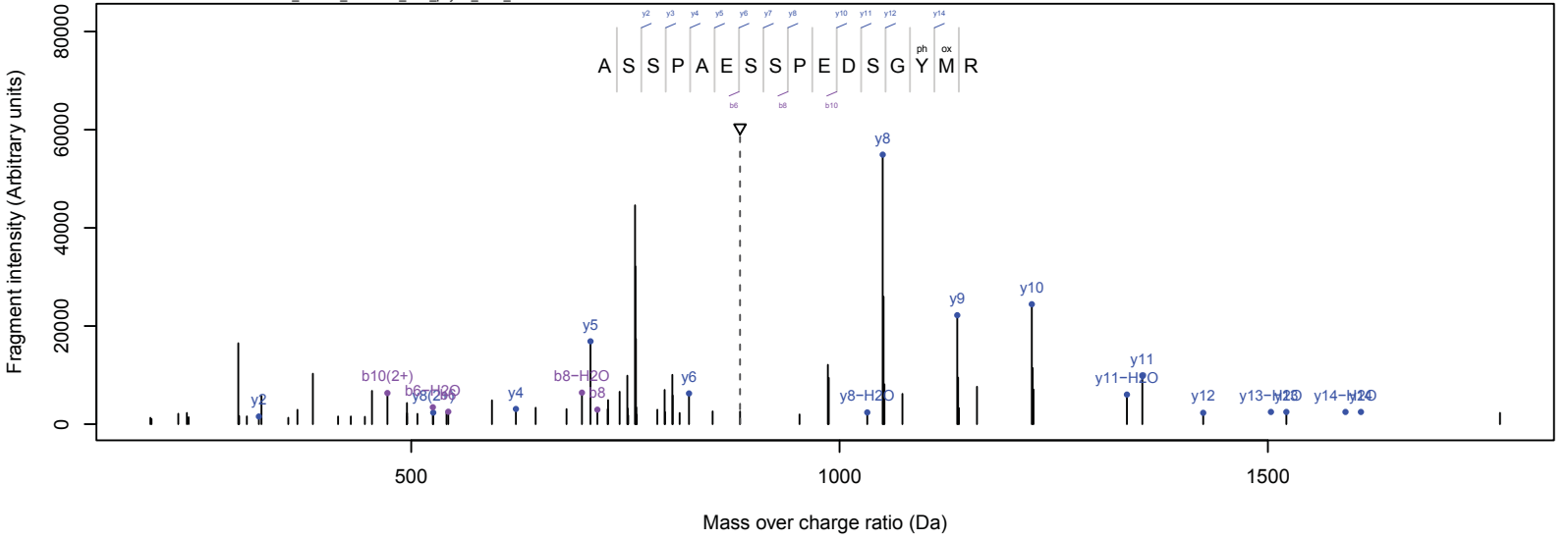




ASSPAESSPEDSGY(ph)M(ox)R-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 5850

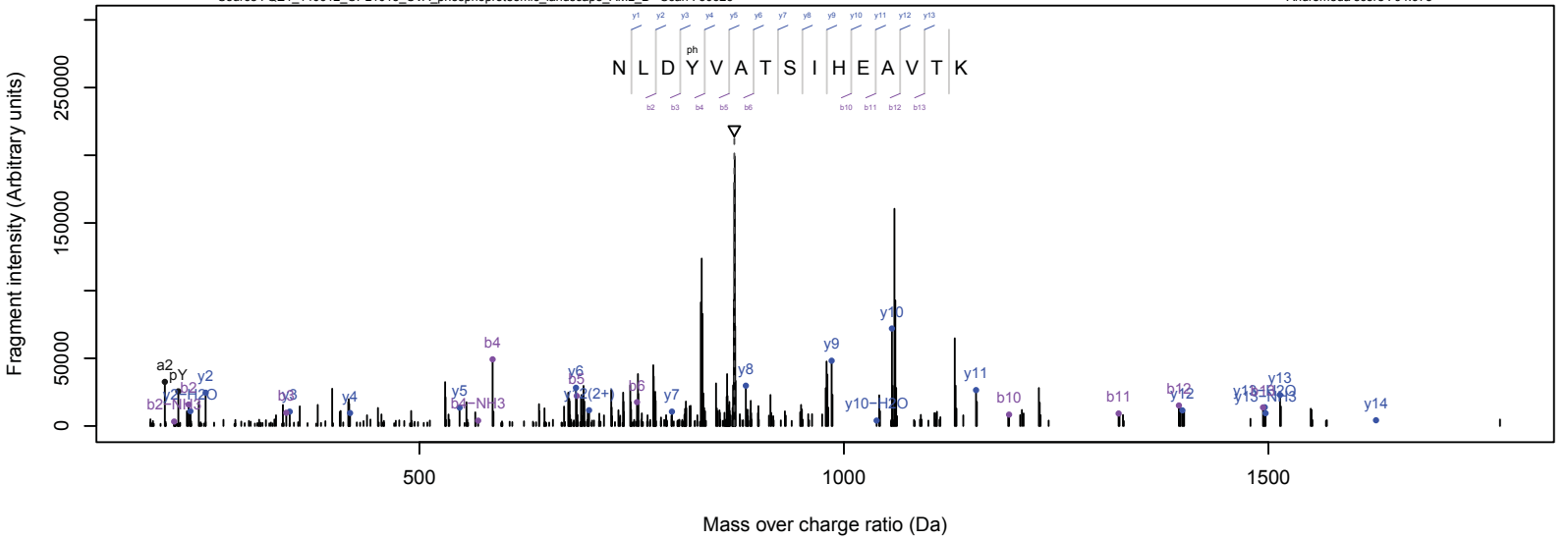
Andromeda score : 95.067



NLDY(ph)VATSIHEAVTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30626

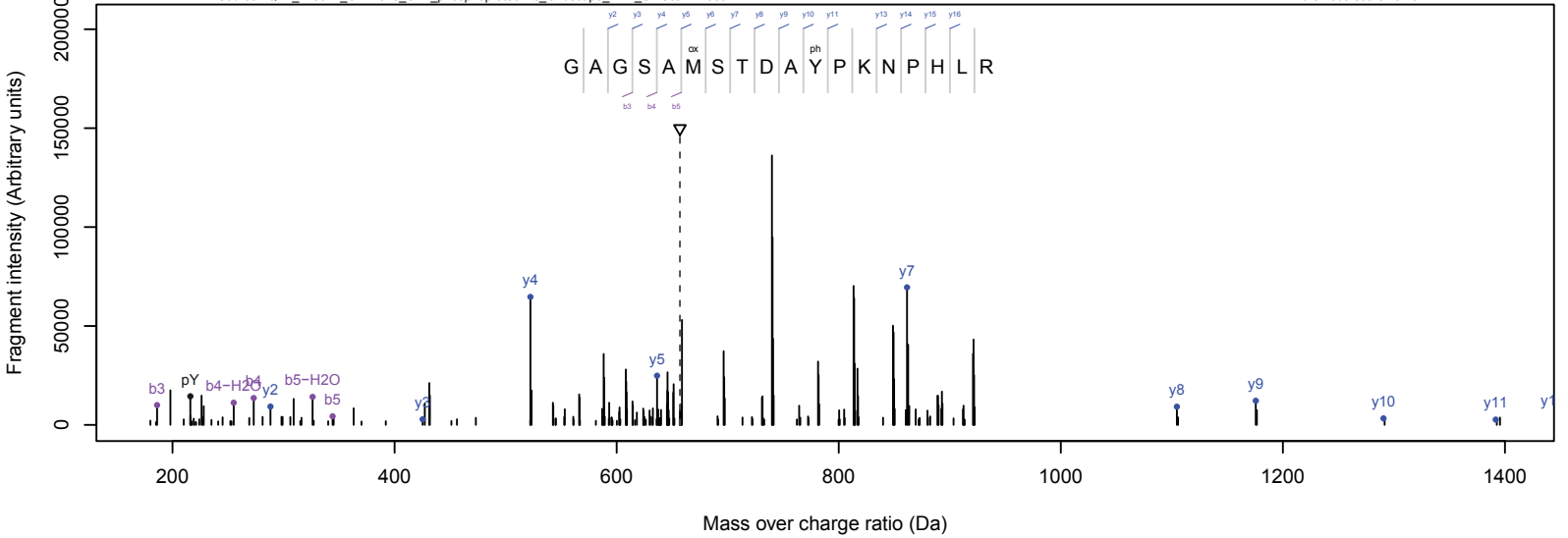
Andromeda score : 94.973



GAGSAM(ox)STDAY(ph)PKNPHLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10032

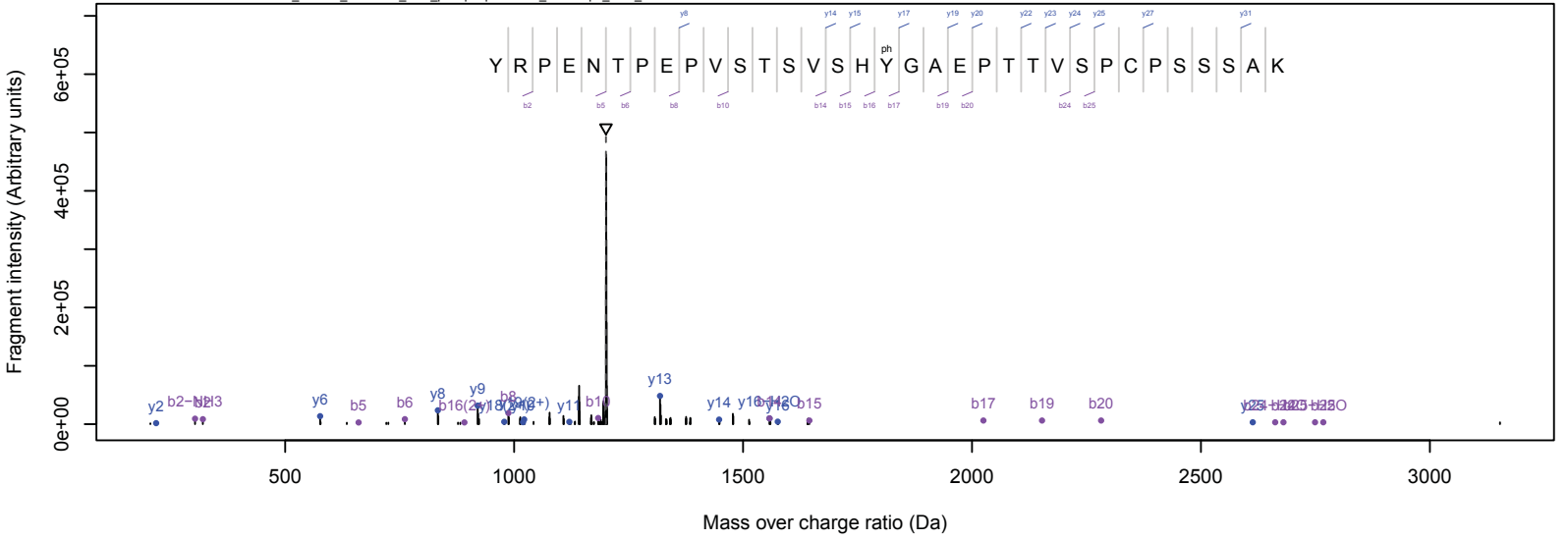
Andromeda score : 94.922



YRPENTPEPVSTSVSHY(ph)GAEPTTVSPCPSSSAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20759

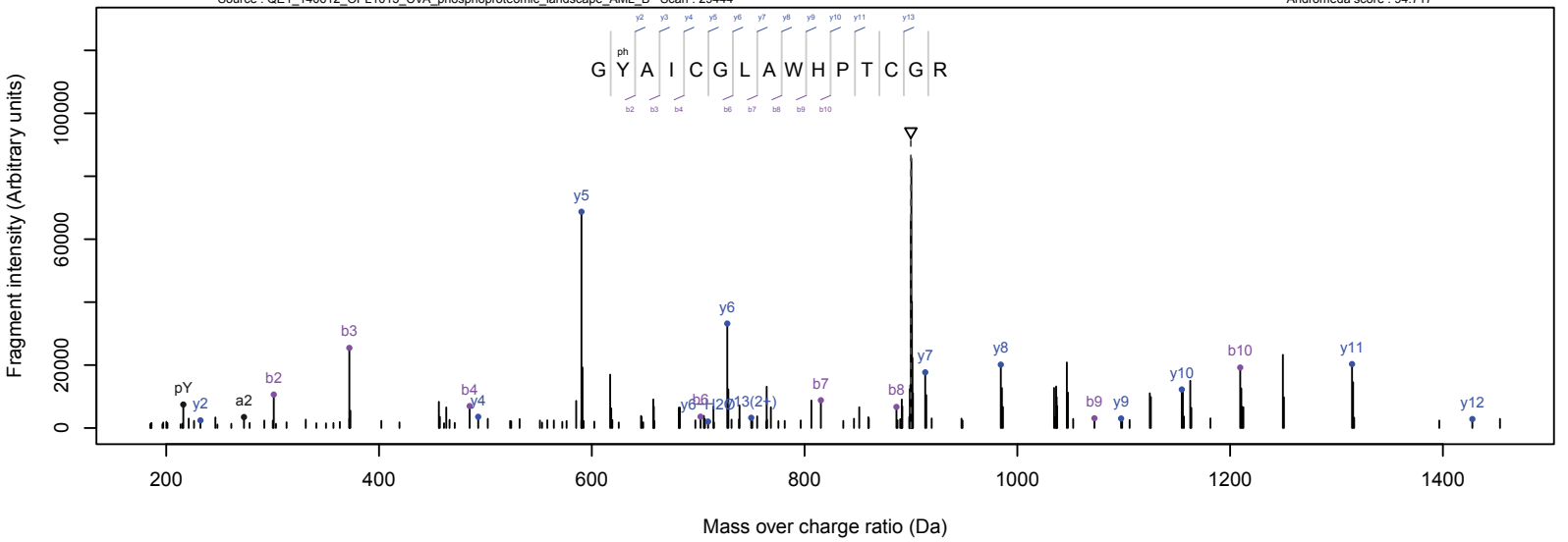
Andromeda score : 94.807



GY(ph)AICGLAWHPTCGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29444

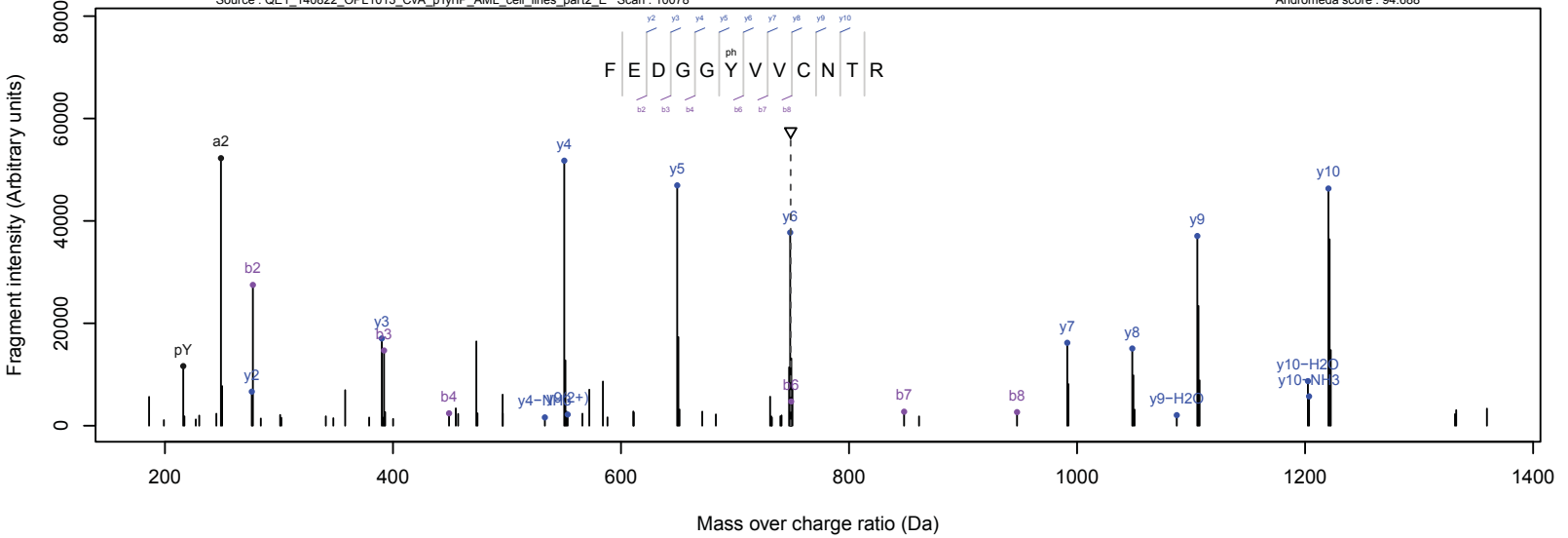
Andromeda score : 94.717



FEDGGY(ph)VVCNTR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 10078

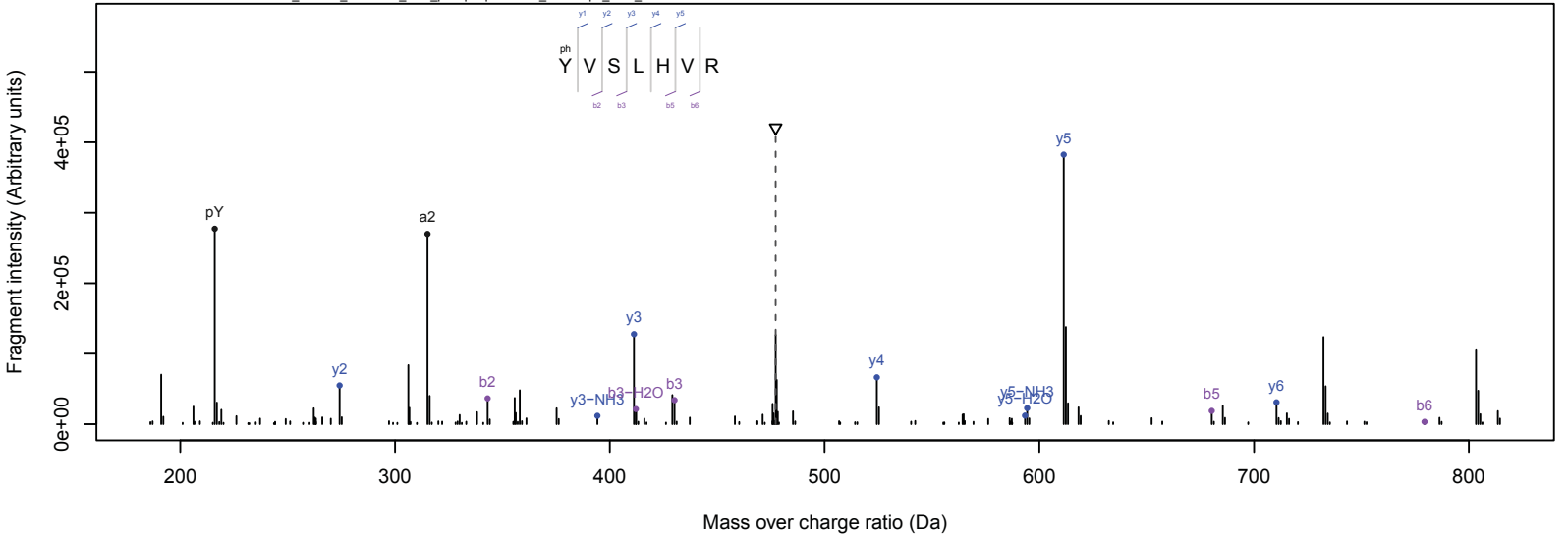
Andromeda score : 94.688



Y(ph)VSLHVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11209

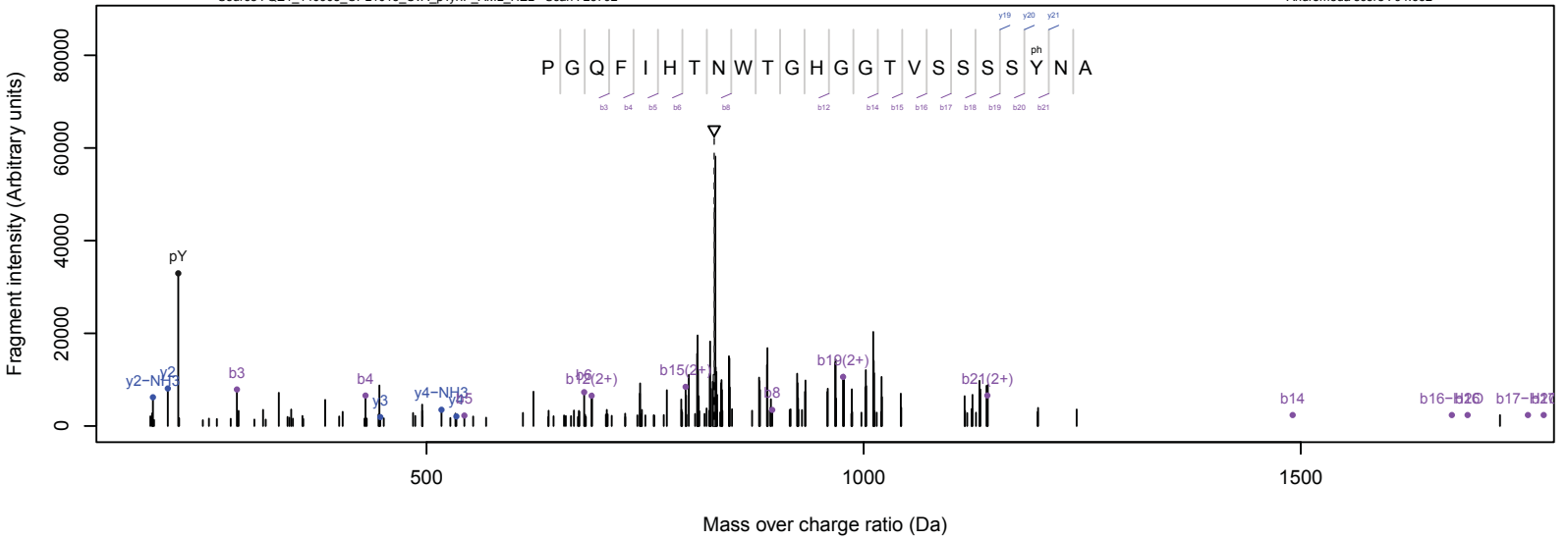
Andromeda score : 94.662



PGQFIHTNWTGHGGTVSSSY(ph)NA-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 25792

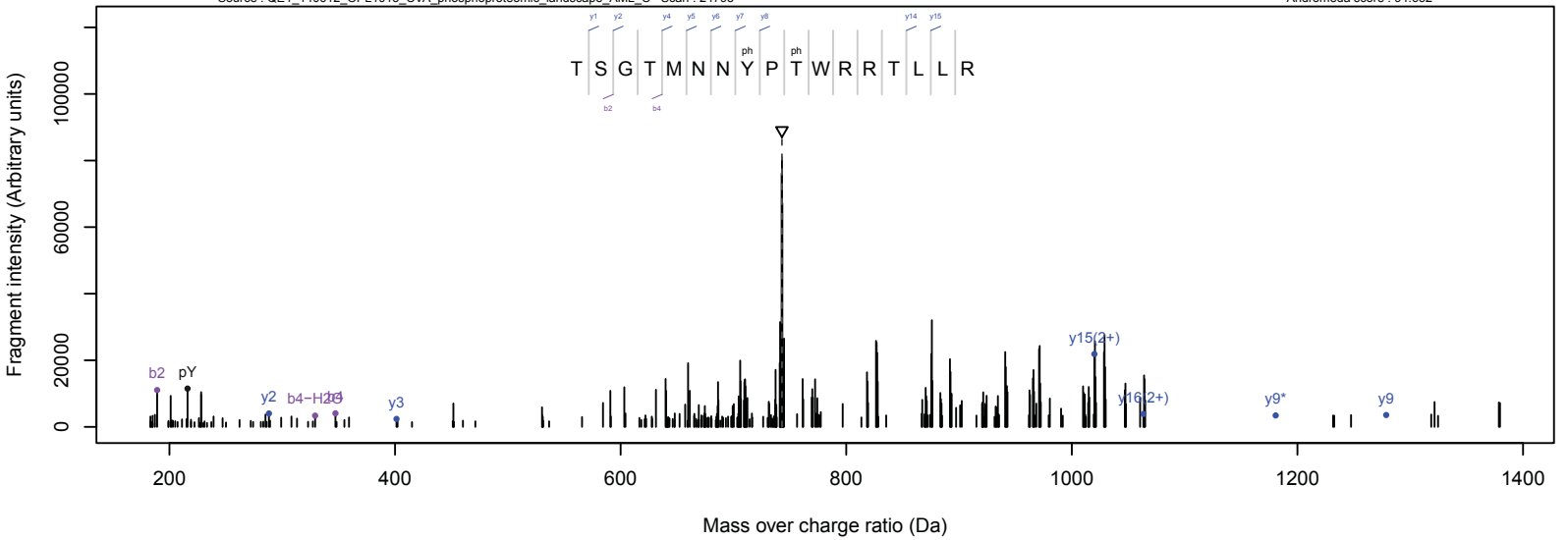
Andromeda score : 94.632



TSGTMNNY(ph)PT(ph)WRRTLLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24795

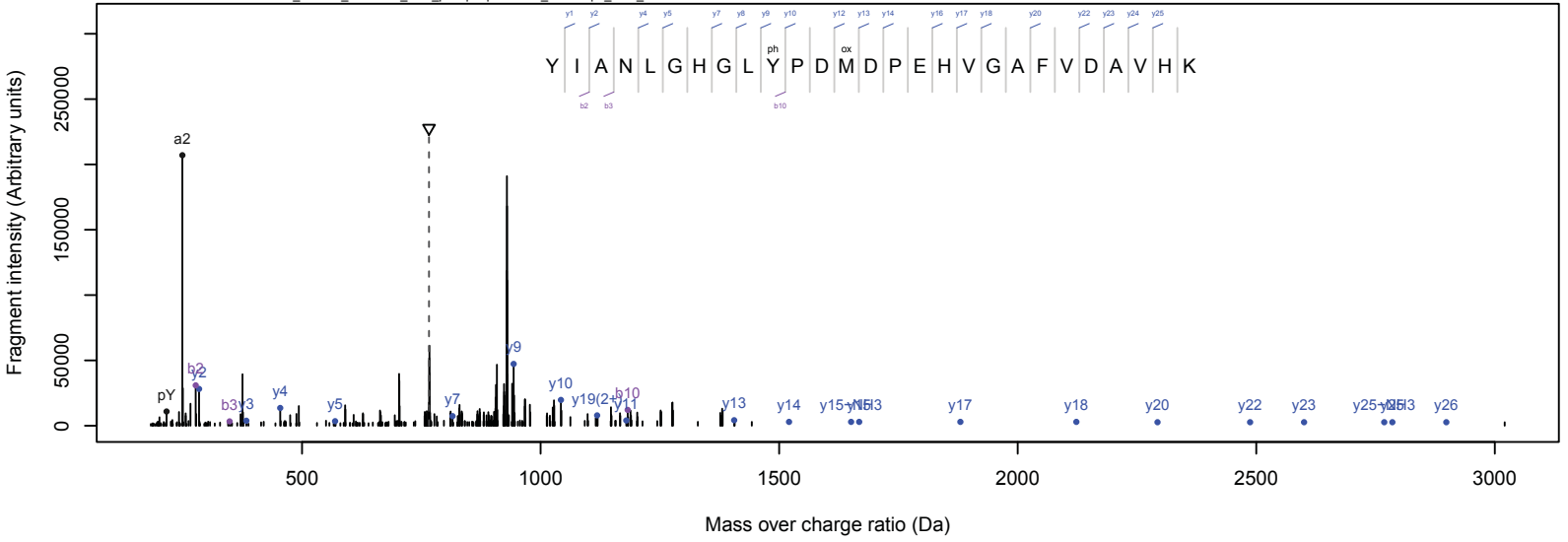
Andromeda score : 94.632



YIANLGHGLY(ph)PDM(ox)DPEHVGAFVDAVHK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 36983

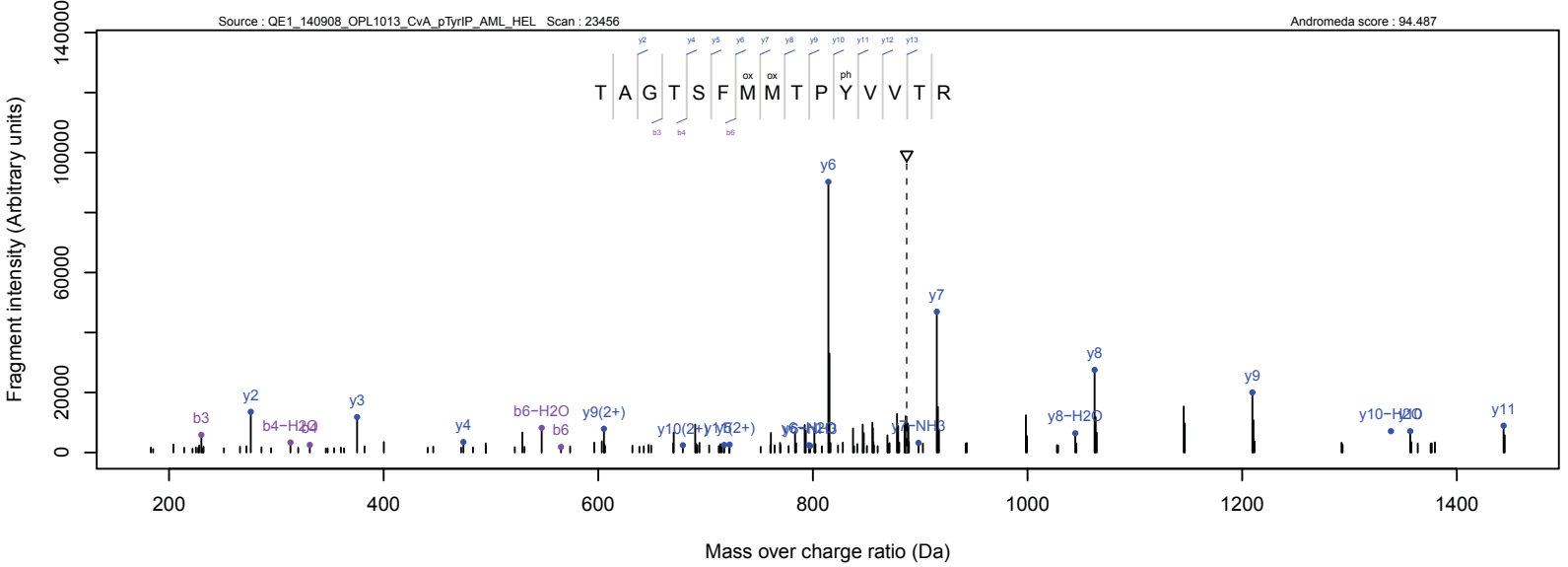
Andromeda score: 94.594



TAGTSFM(ox)M(ox)TPY(ph)VVTR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 23456

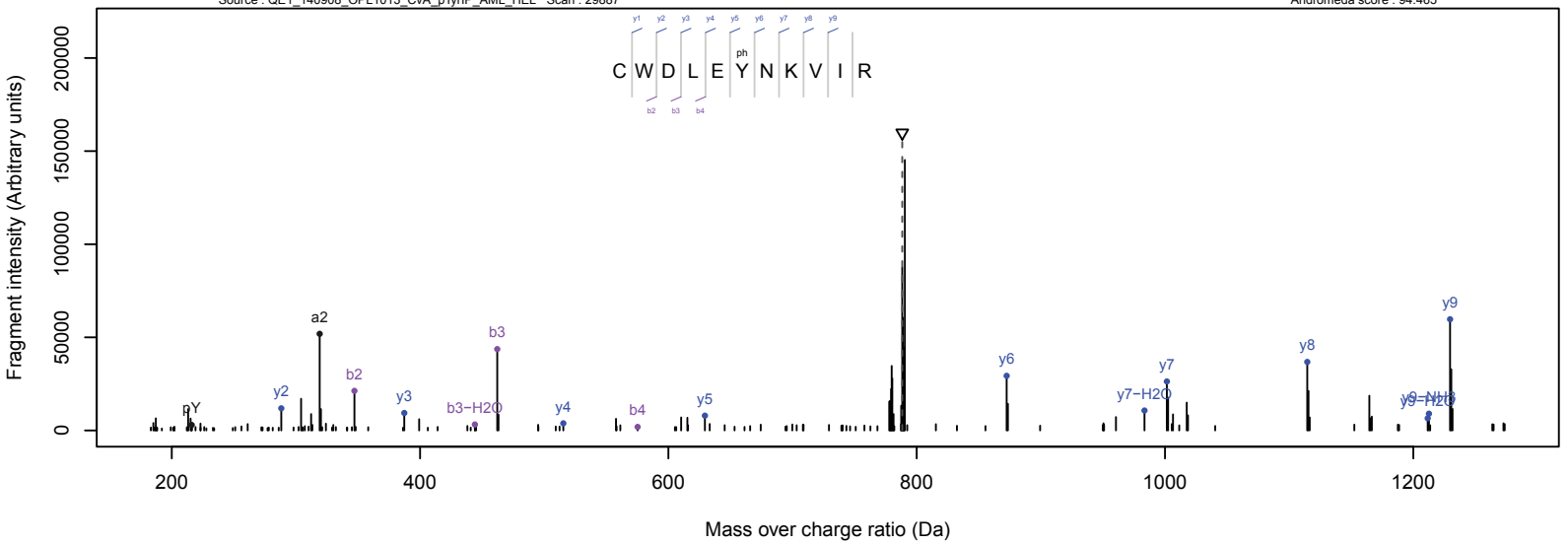
Andromeda score: 94.487



CWDLEY(ph)NKVIR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 29887

Andromeda score: 94.465

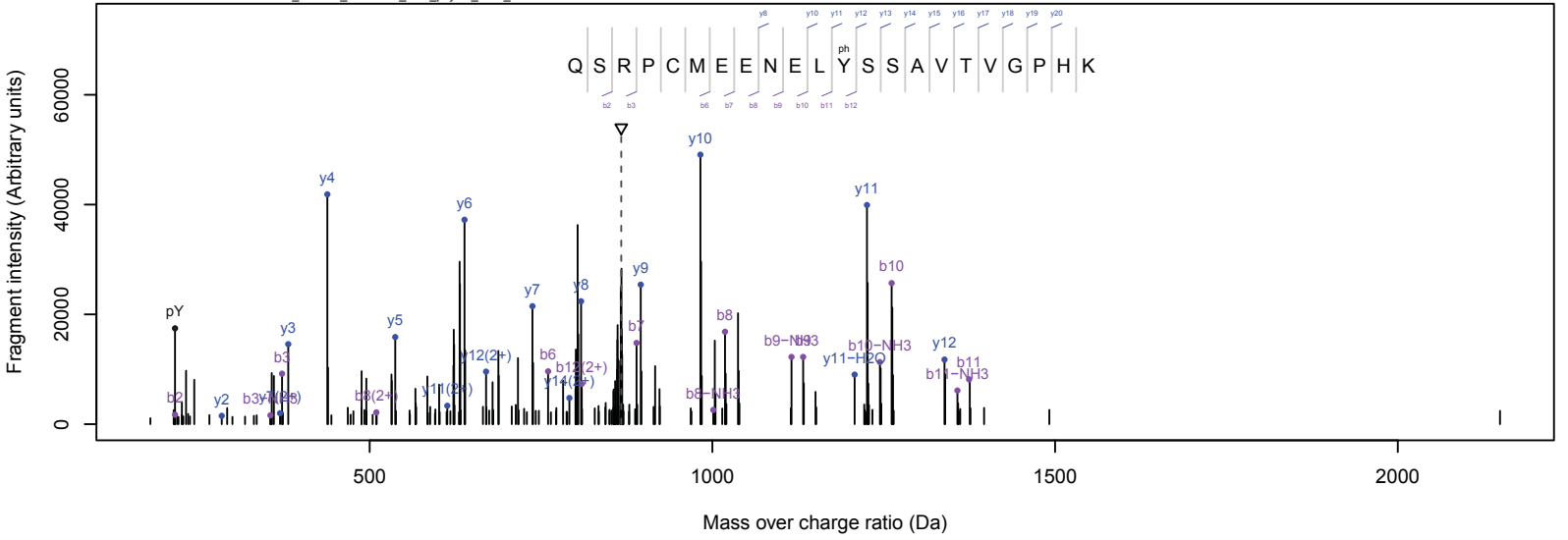




QSRPCMEENELYSSAVTVGPHK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 22024

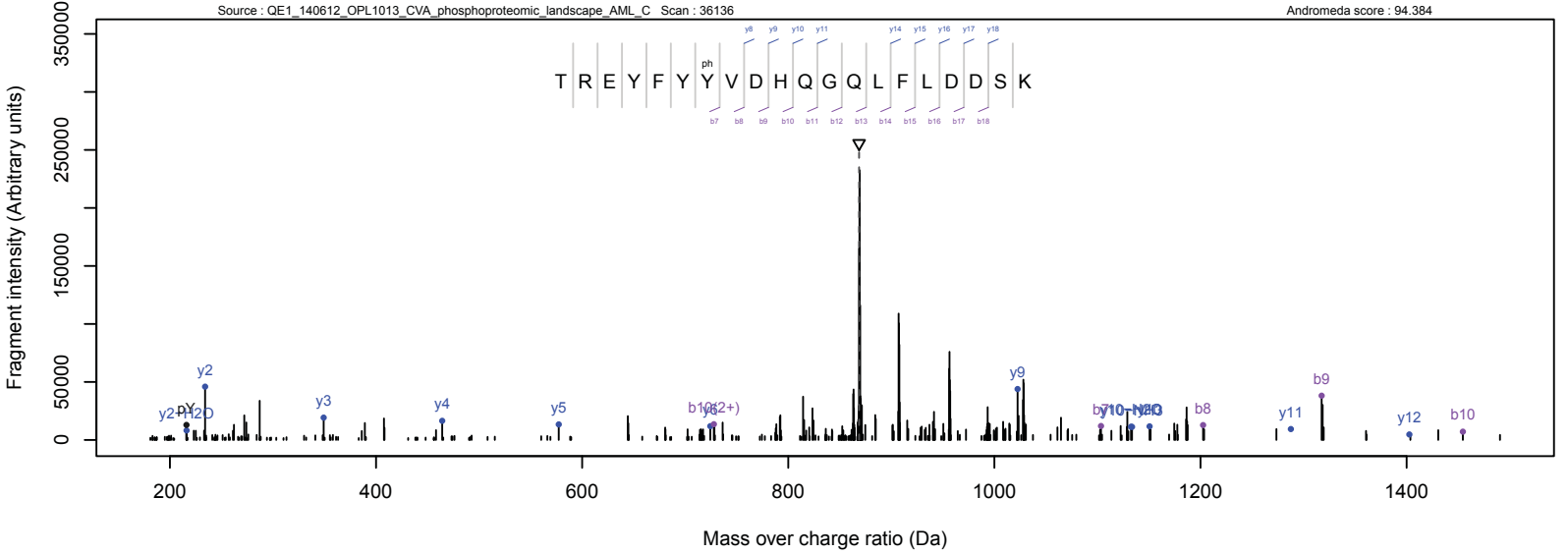
Andromeda score : 94.384



TREYFY(ph)VDHQQLFLDDSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36136

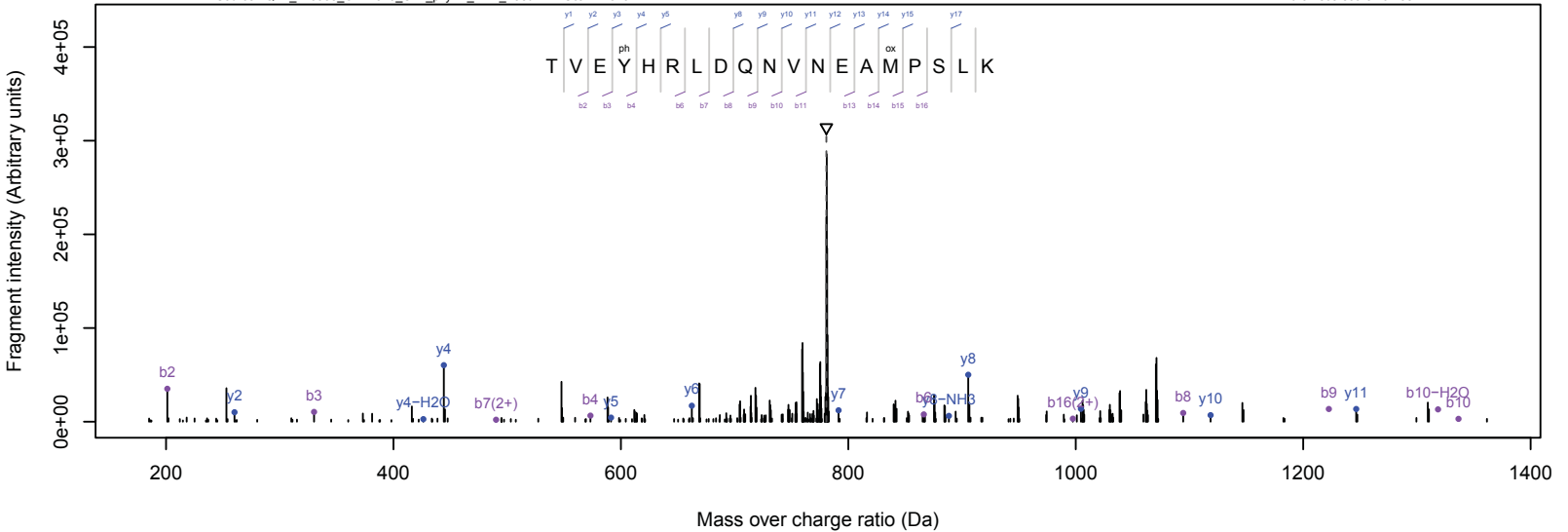
Andromeda score : 94.384



TVEY(ph)HRLDQNVNEAMP(SL)K-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 16737

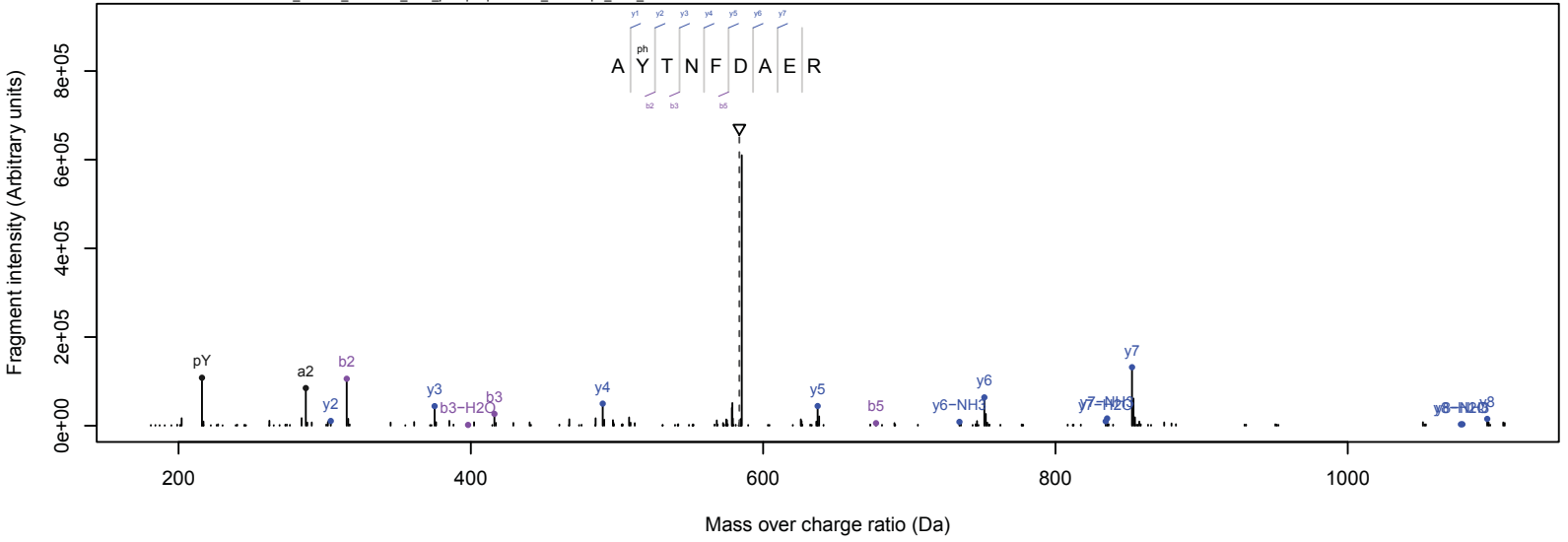
Andromeda score : 94.384



AY(ph)TNFDAER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9108

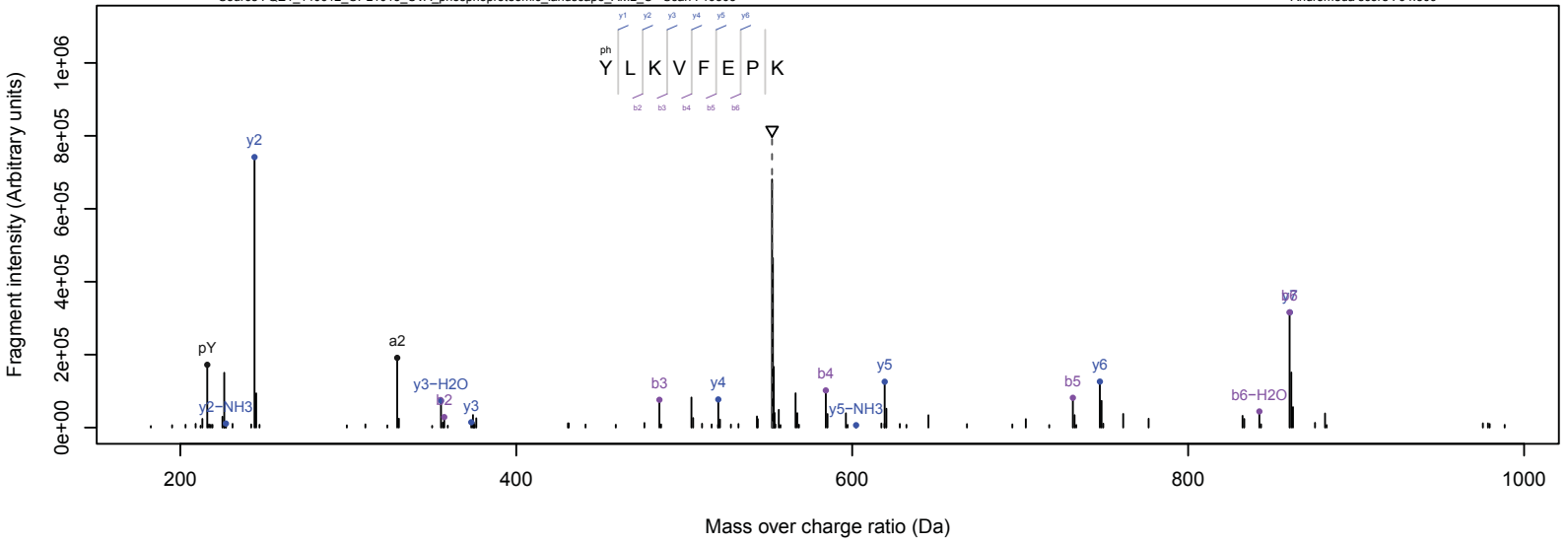
Andromeda score : 94.309



Y(ph)LKVFEPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19303

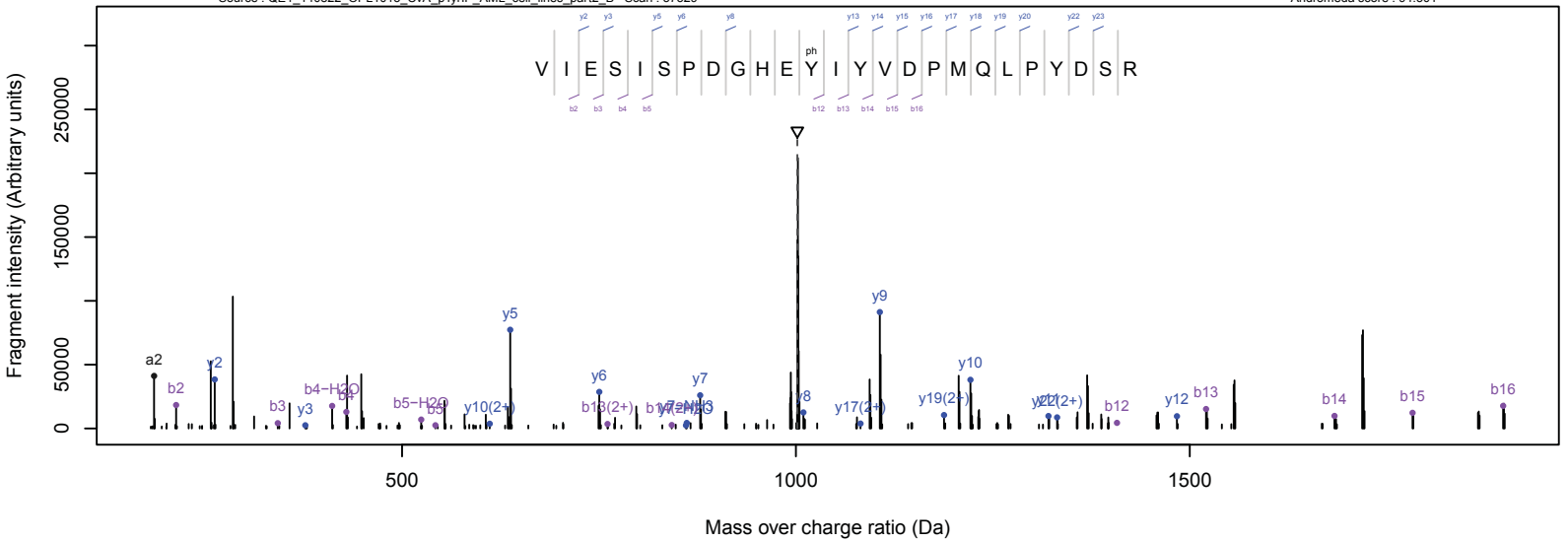
Andromeda score : 94.309



VIESISPDGHEY(ph)IYVDPMQLPYDSR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 37329

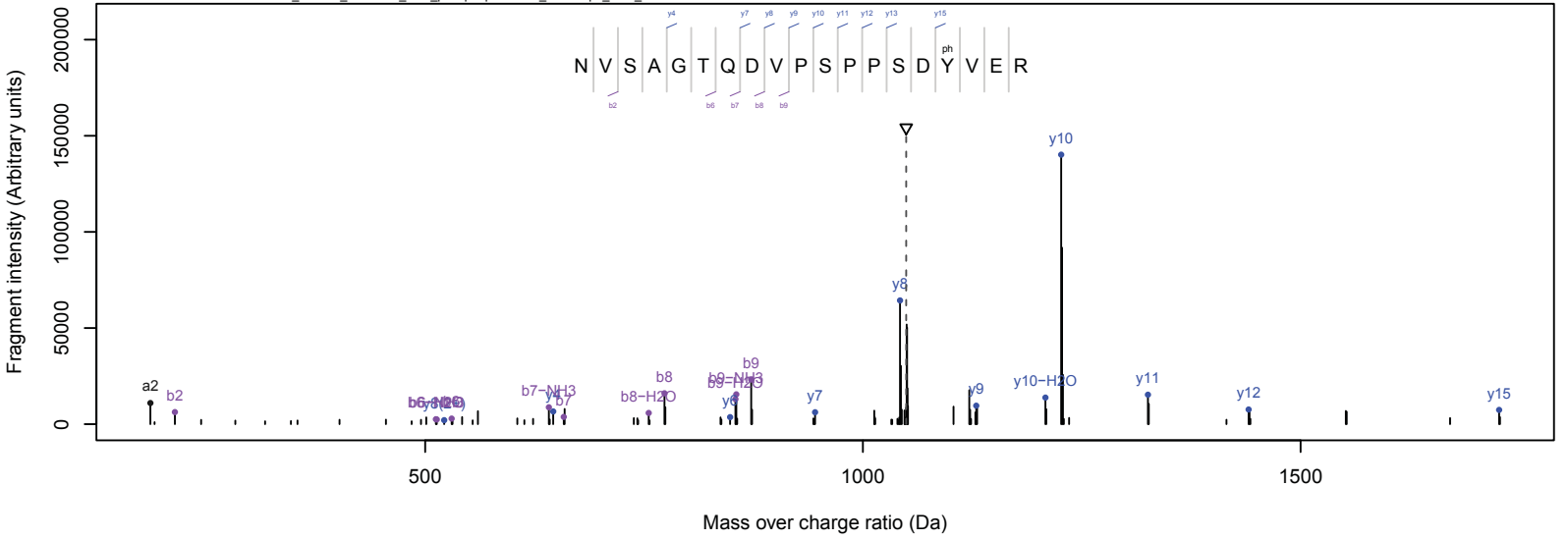
Andromeda score : 94.301



NVSAGTQDVPSPPSDY(ph)VER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21141

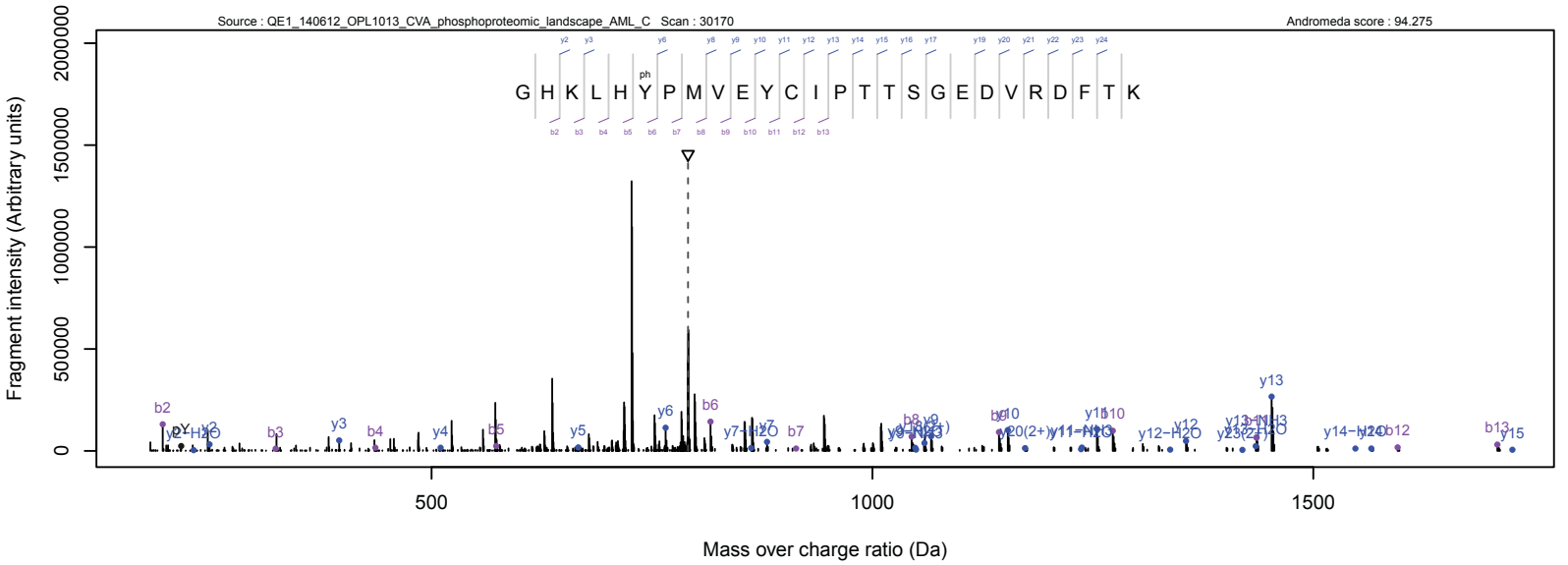
Andromeda score : 94.281



GHKLHY(ph)PMVEYCIPTTSGEDVRDFTK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30170

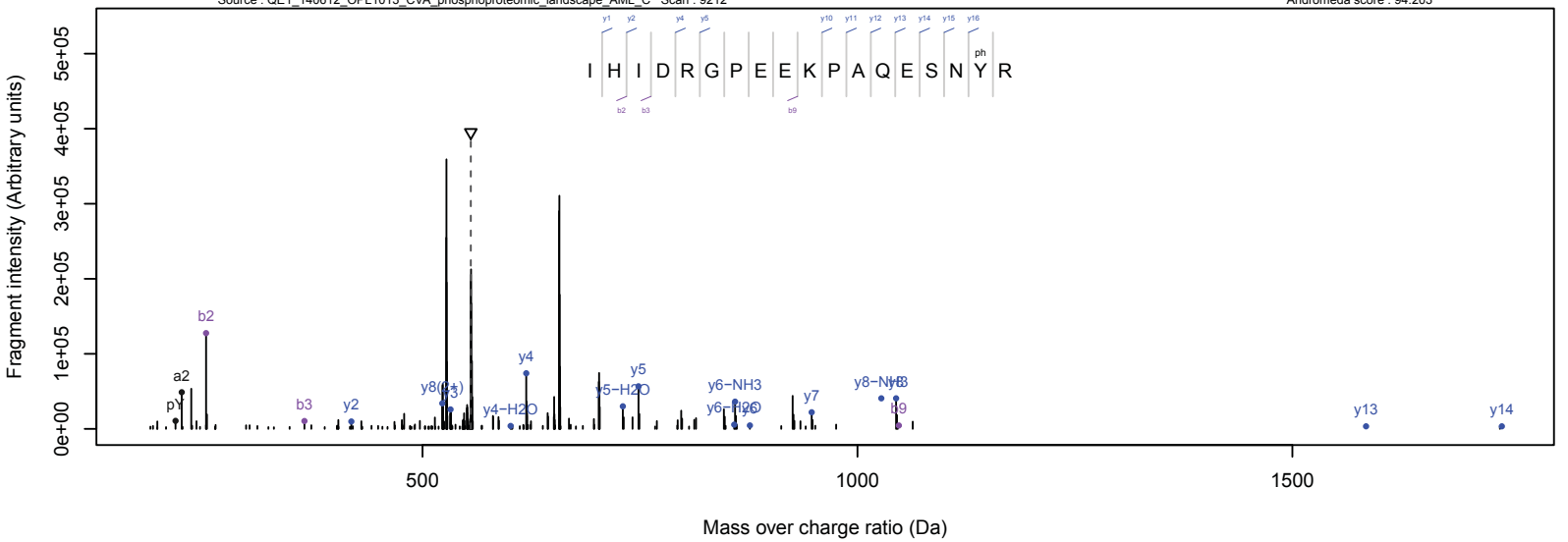
Andromeda score : 94.275



IHDRGPEEKPAQESNY(ph)R-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9212

Andromeda score : 94.203

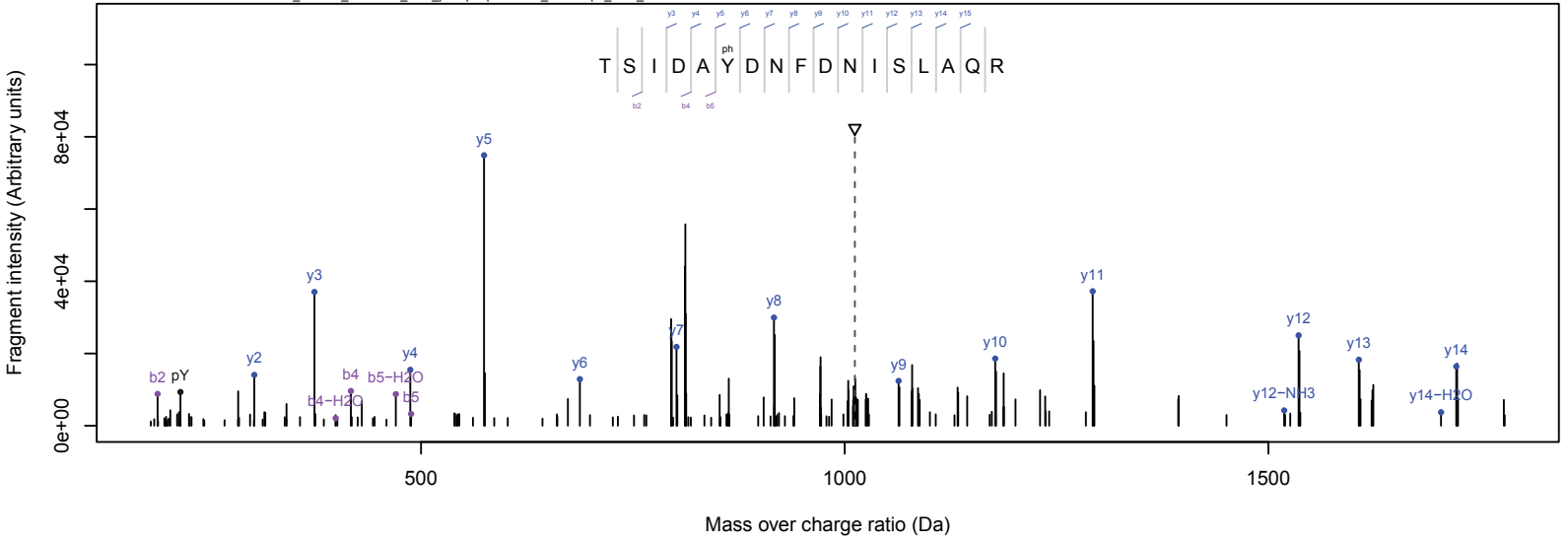




TSIDAY(ph)DNFDNISLAQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34948

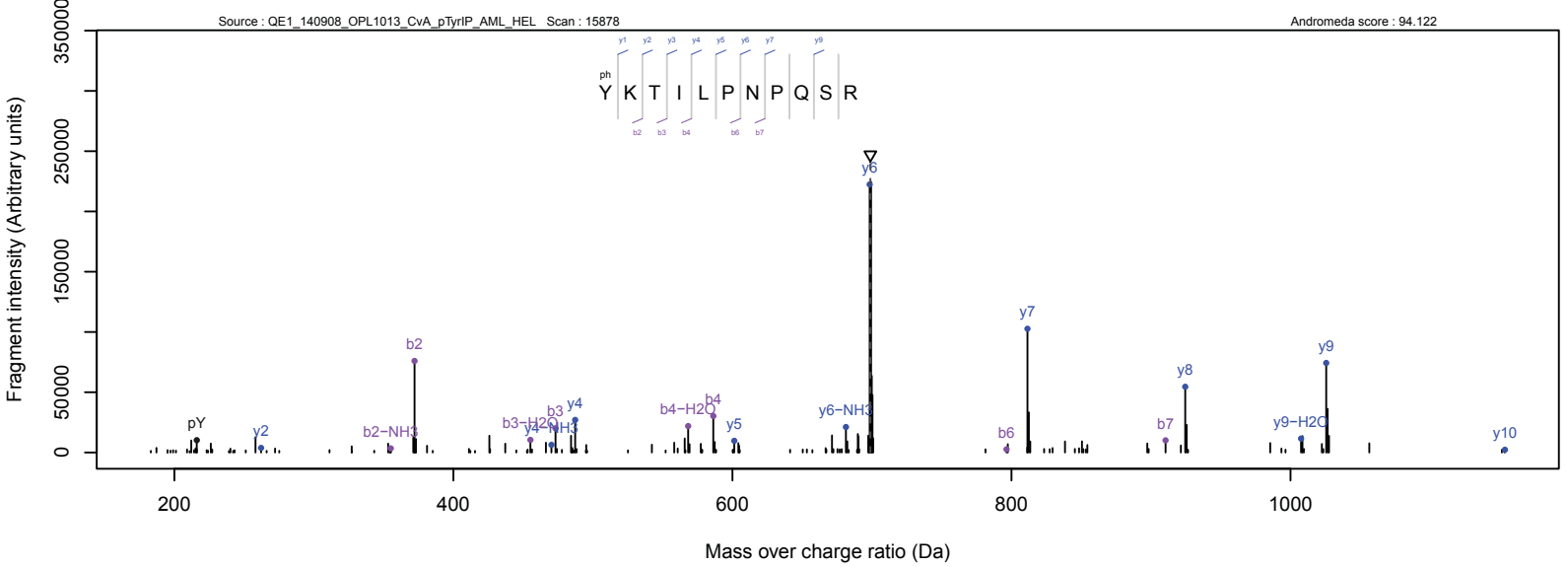
Andromeda score : 94.203



Y(ph)KTILPNPQSR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 15878

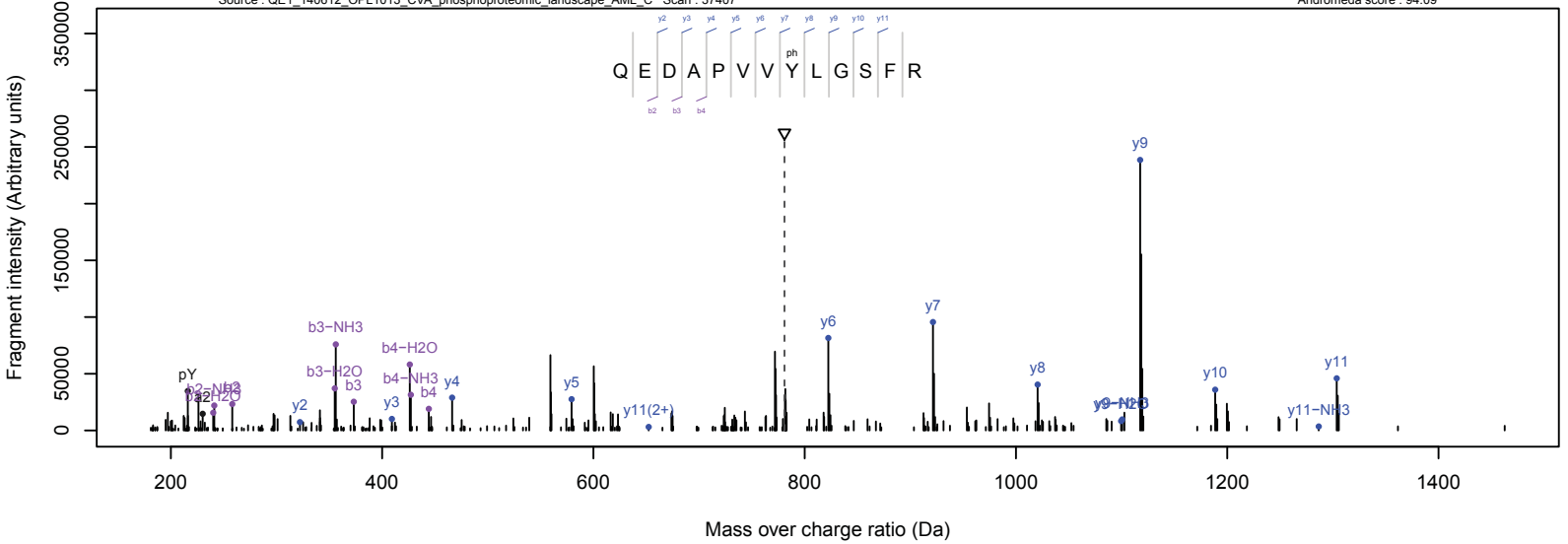
Andromeda score : 94.122



QEDAPVY(ph)LGSFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37407

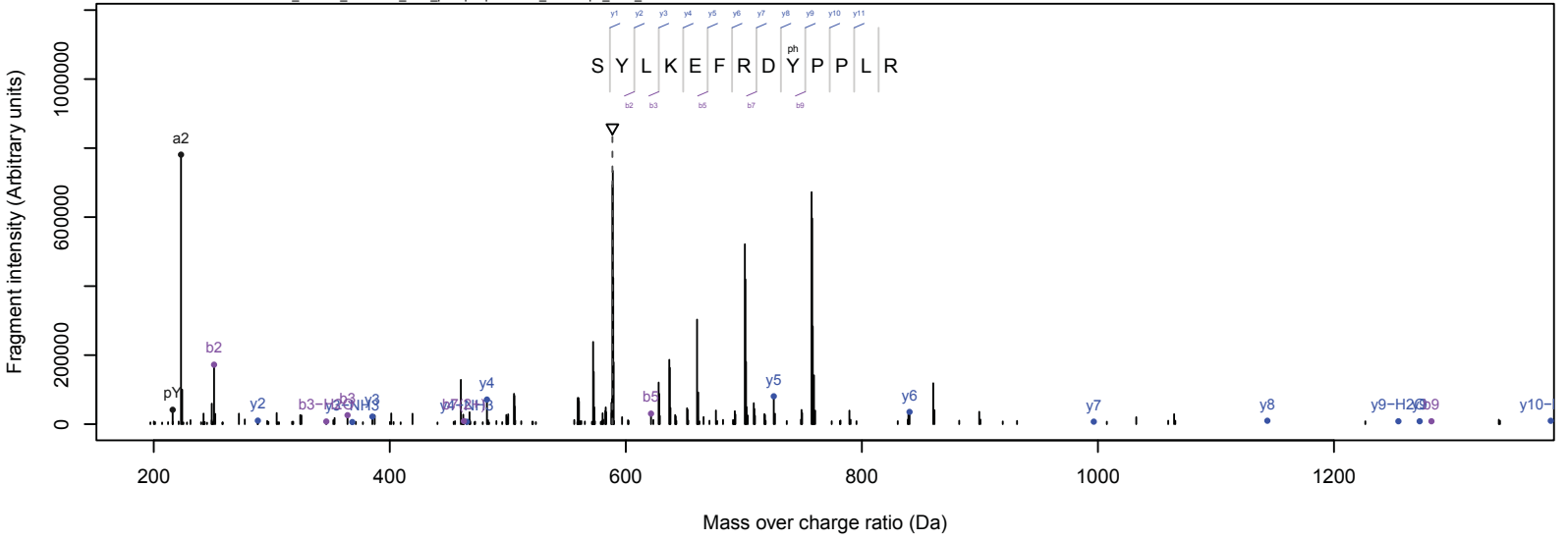
Andromeda score : 94.09



SYLKEFRDY(ph)PPLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26192

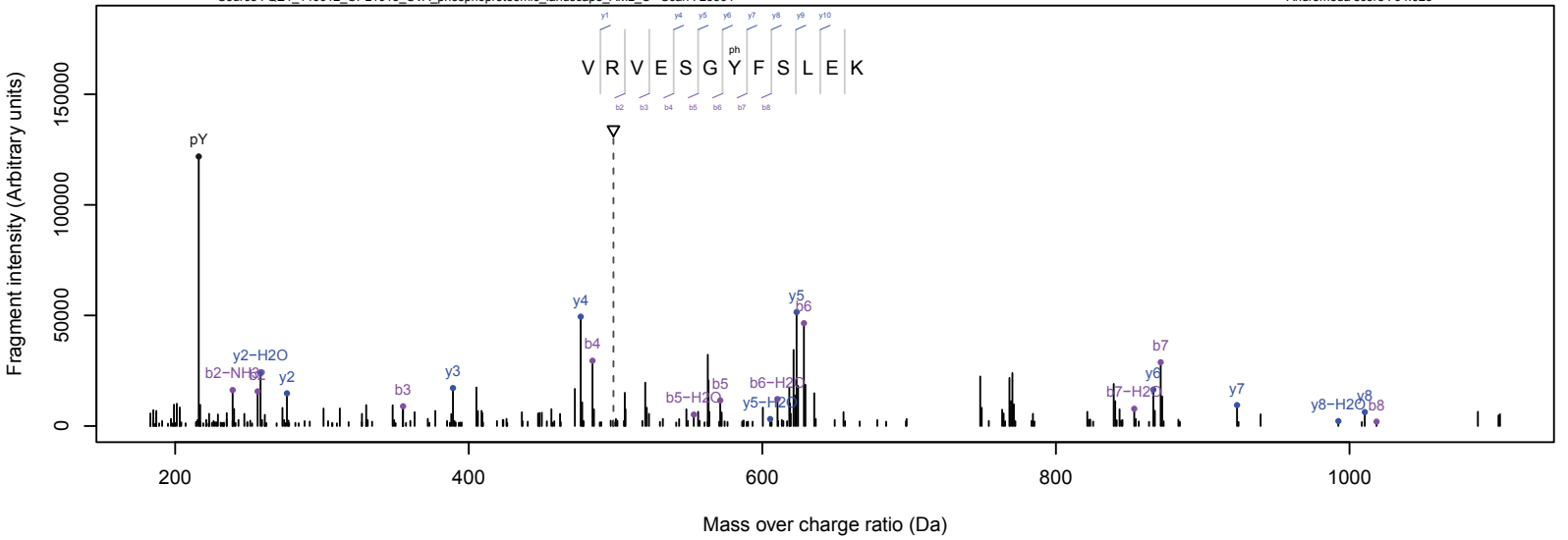
Andromeda score : 94.023



VRVESGY(ph)FSLEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23364

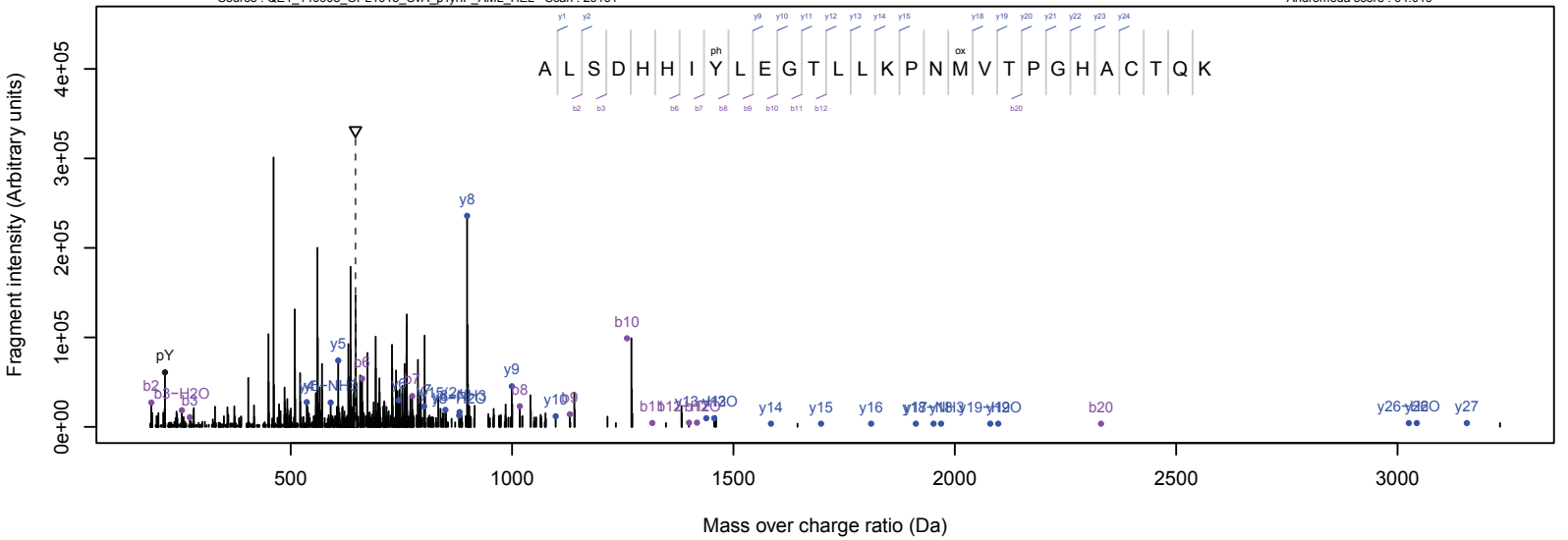
Andromeda score : 94.023



ALSDHHIY(ph)LEGTLKPNMVTTPGHACTQK-5+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29161

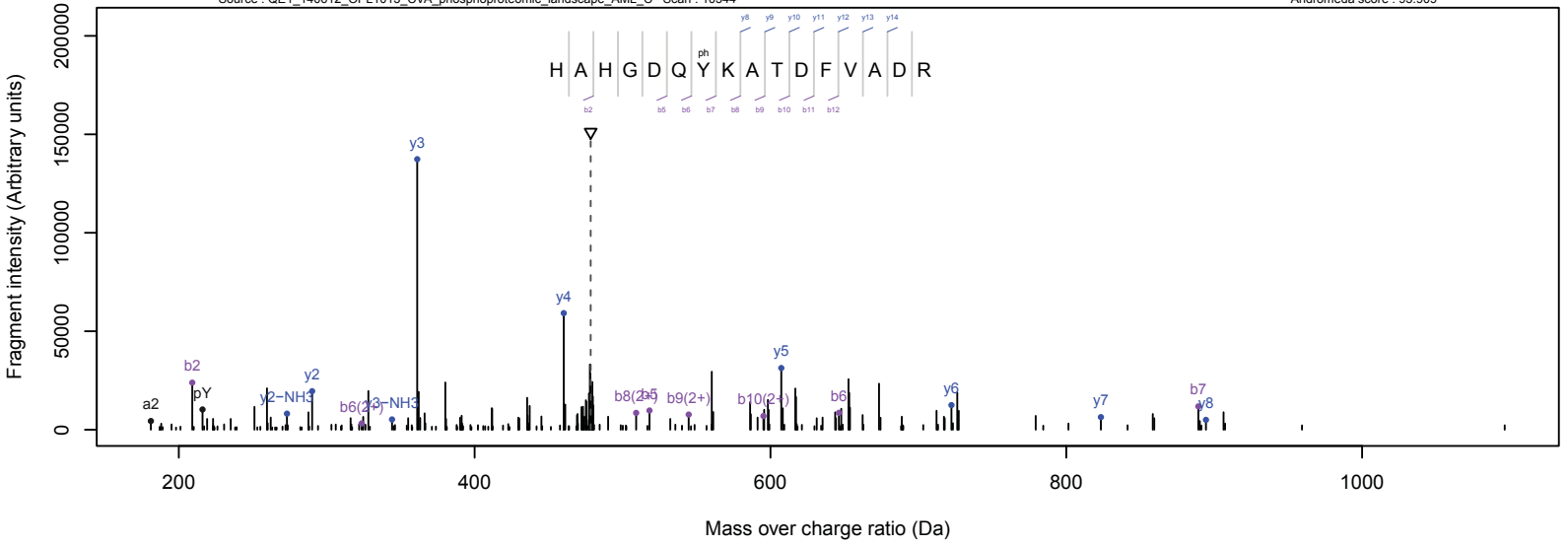
Andromeda score : 94.019



HAHGDQY(ph)KATDFVADR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10344

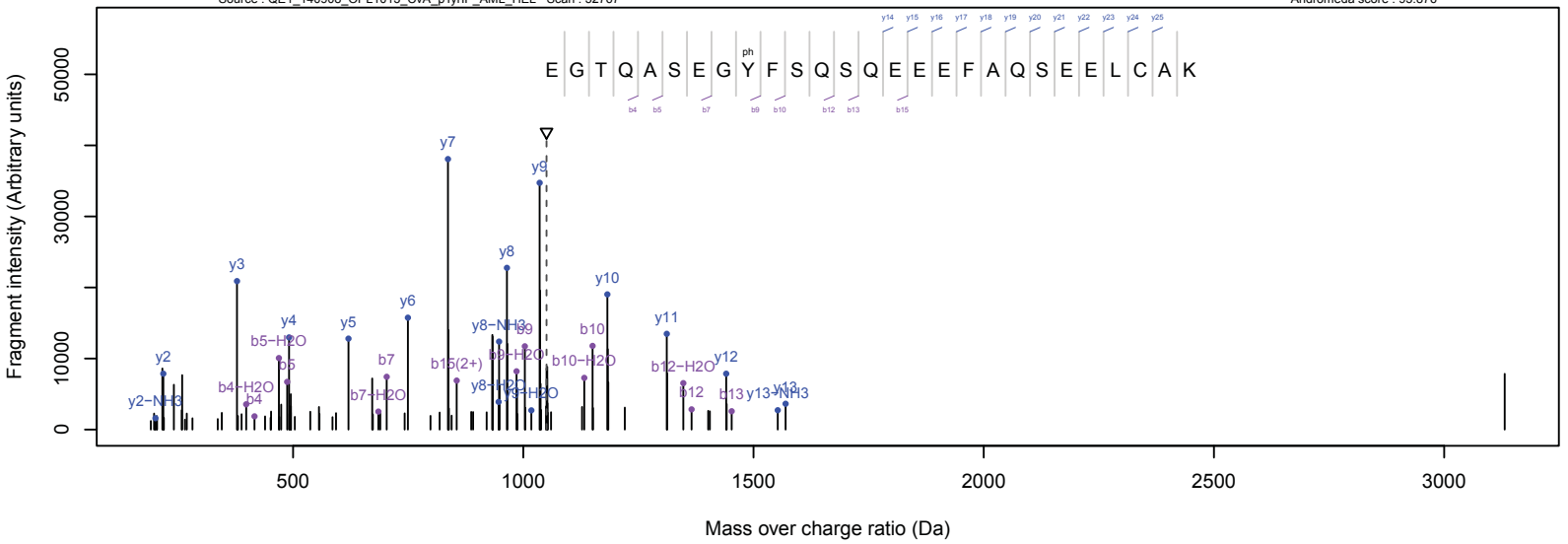
Andromeda score: 93.909



EGTQASEGY(ph)FSQSQEEFAQSEELCAK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 32767

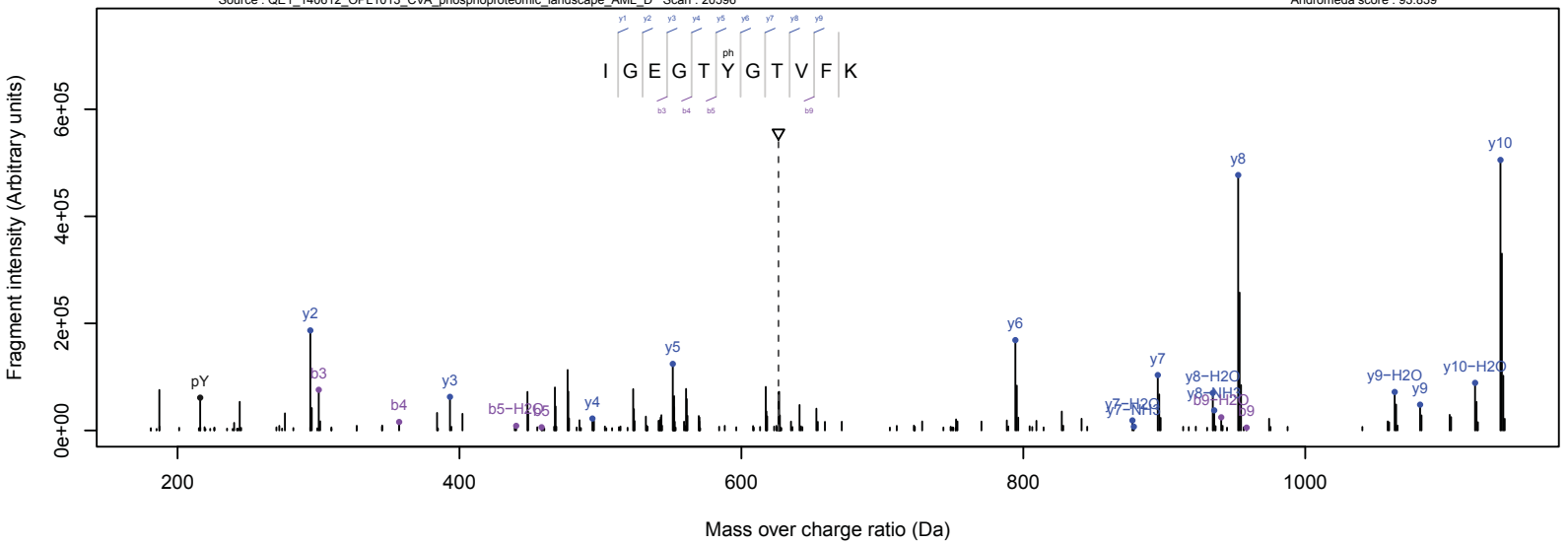
Andromeda score: 93.876



IGEGTY(ph)GTVFK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 20596

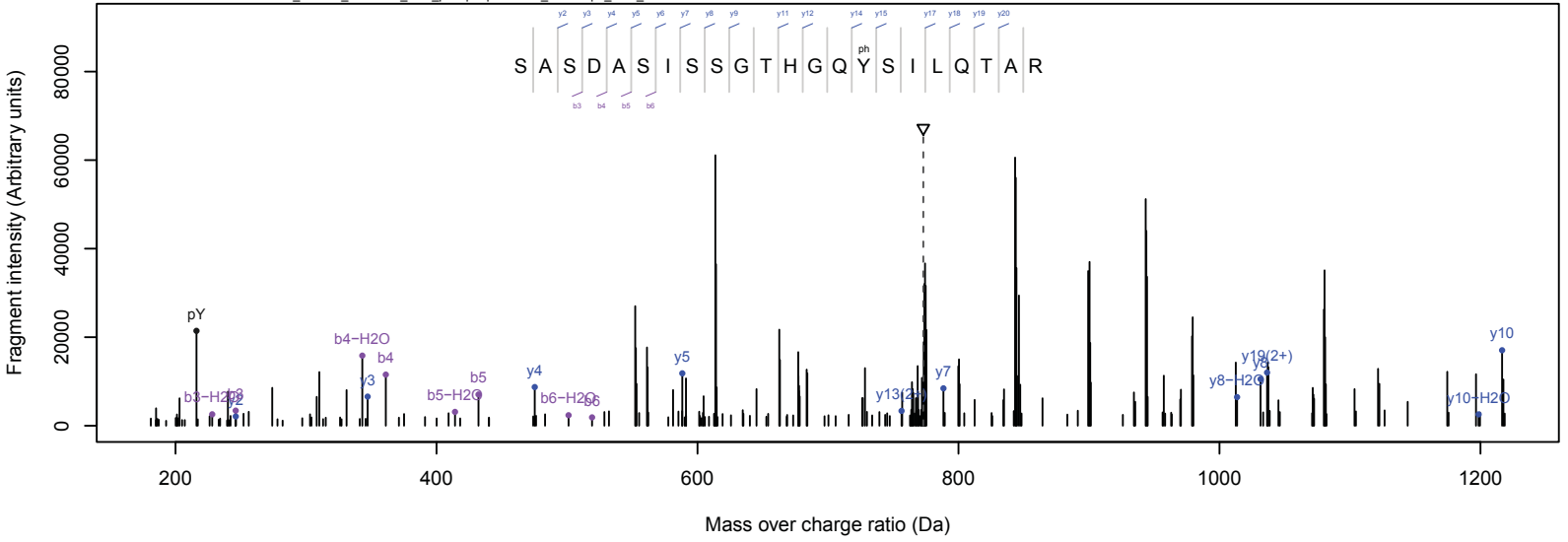
Andromeda score: 93.839



SASDASISSGTHGQY(ph)SILQ TAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20857

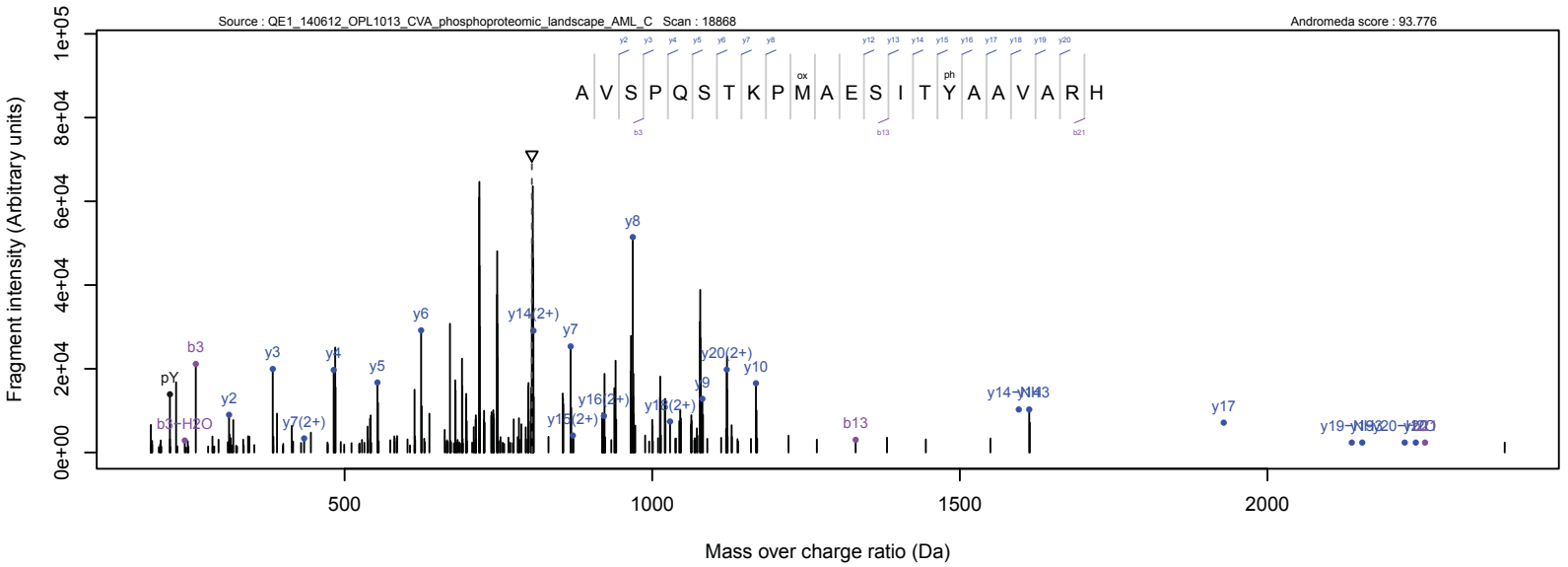
Andromeda score : 93.804



AVSPQSTKPM(ox)AESITY(ph)AAVARH-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18868

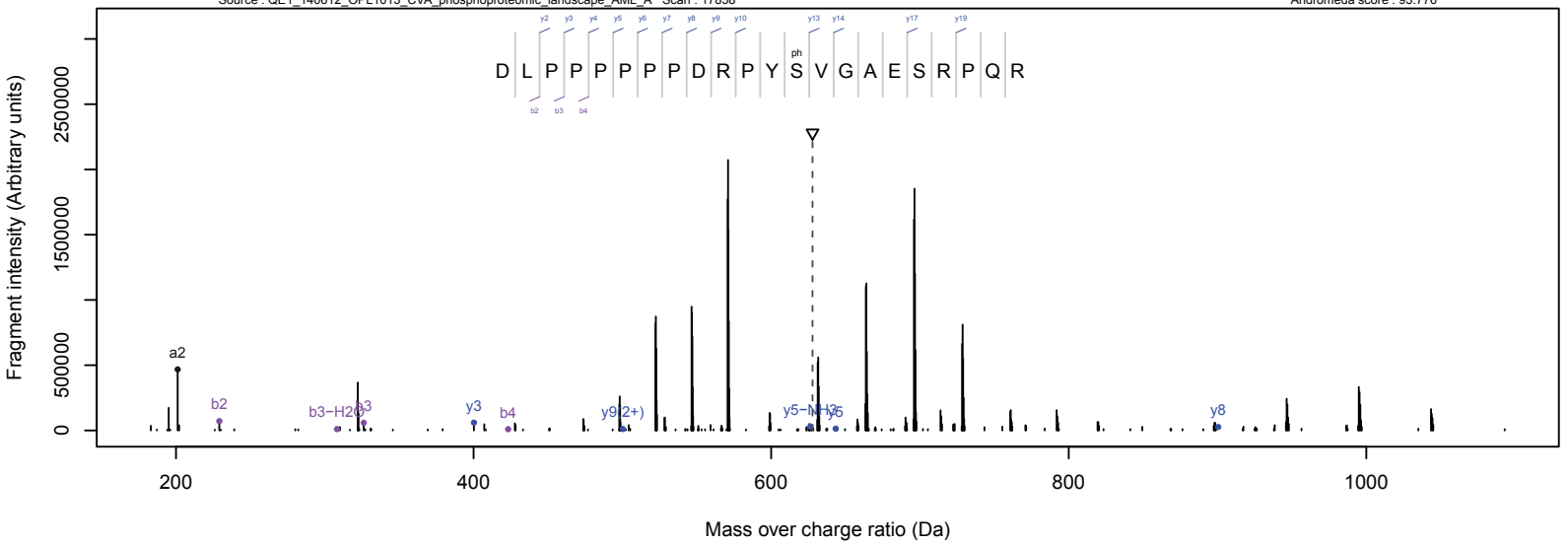
Andromeda score : 93.776



DLPPPPDRPYS(ph)VGAESRPQR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17858

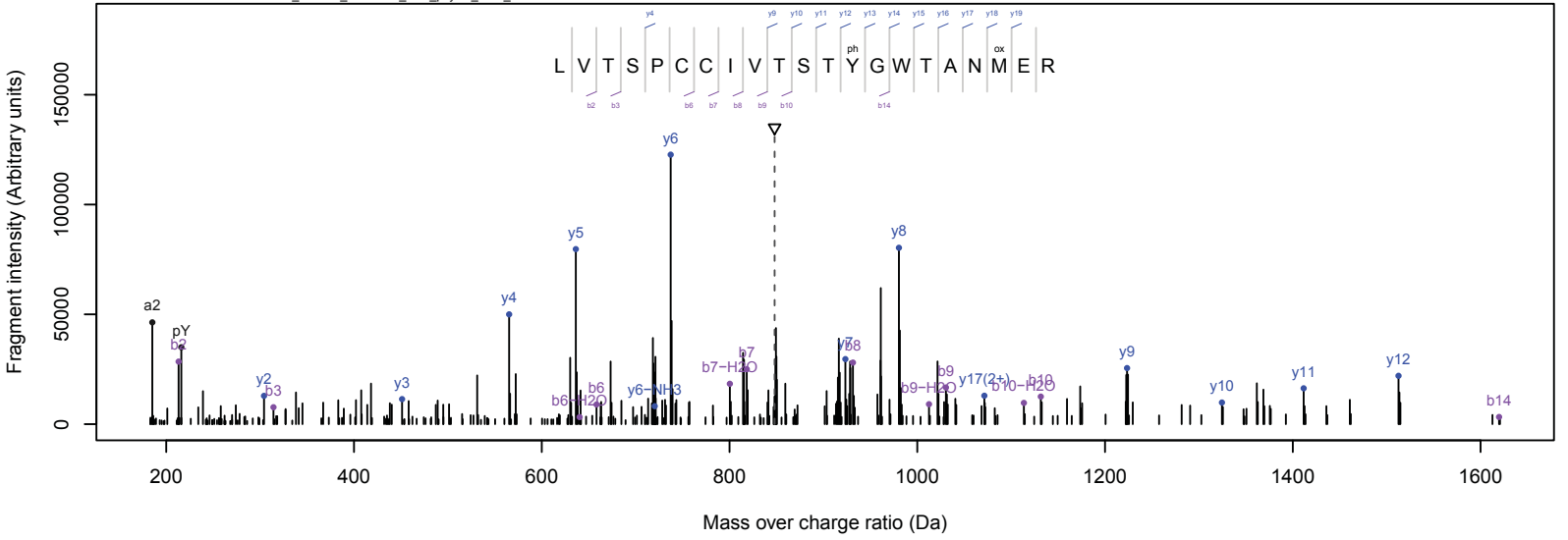
Andromeda score : 93.776



LVTSPPCCIVTSTY(ph)GWTANM(ox)ER-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 36116

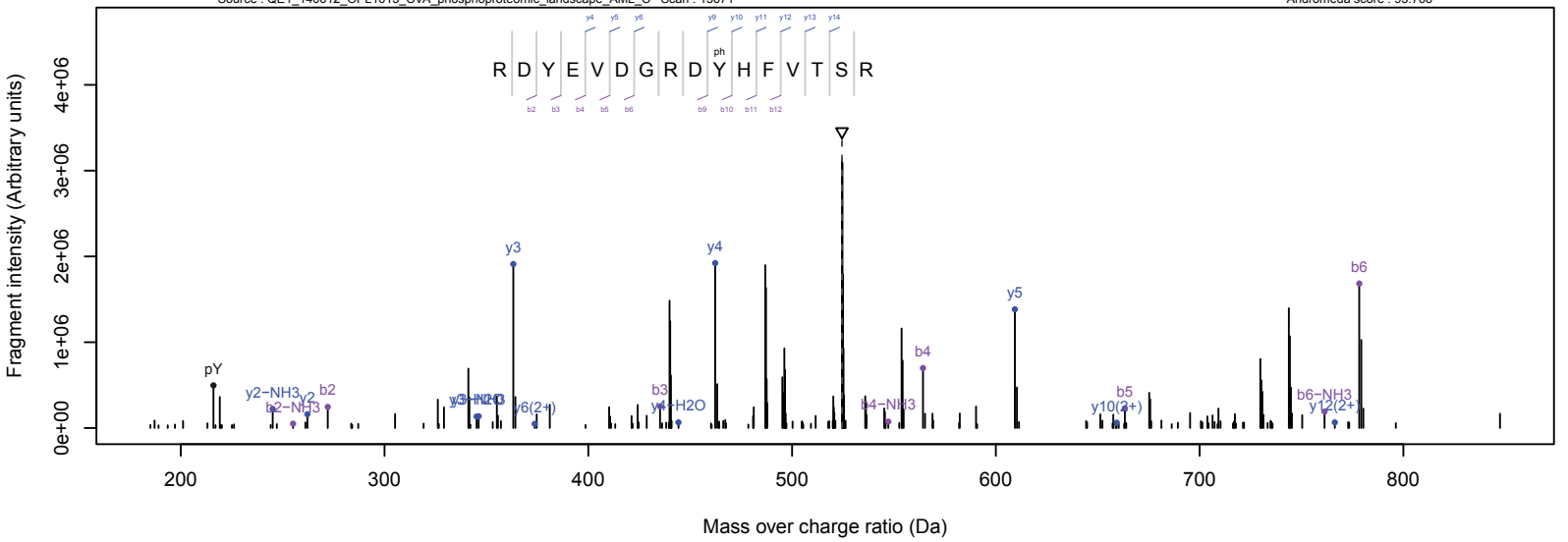
Andromeda score : 93.776



RDYEVDRDY(ph)HFVTSR-4+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 15671

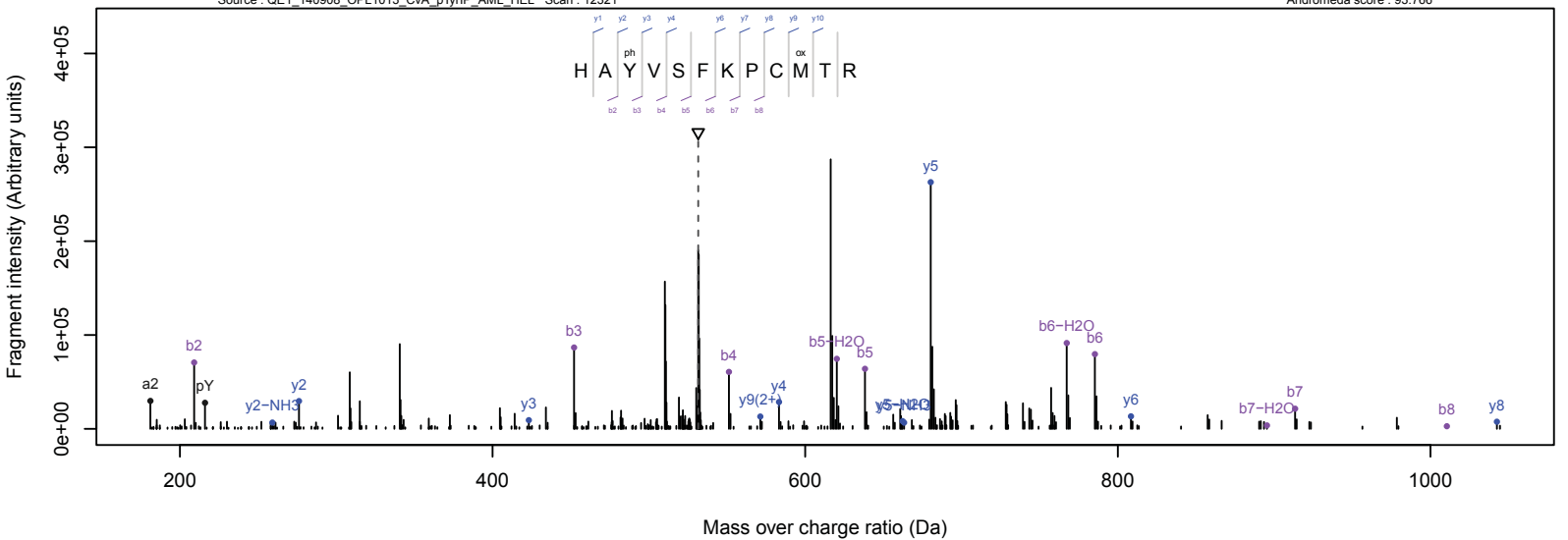
Andromeda score : 93.768



HAY(ph)VSKPCMTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 12321

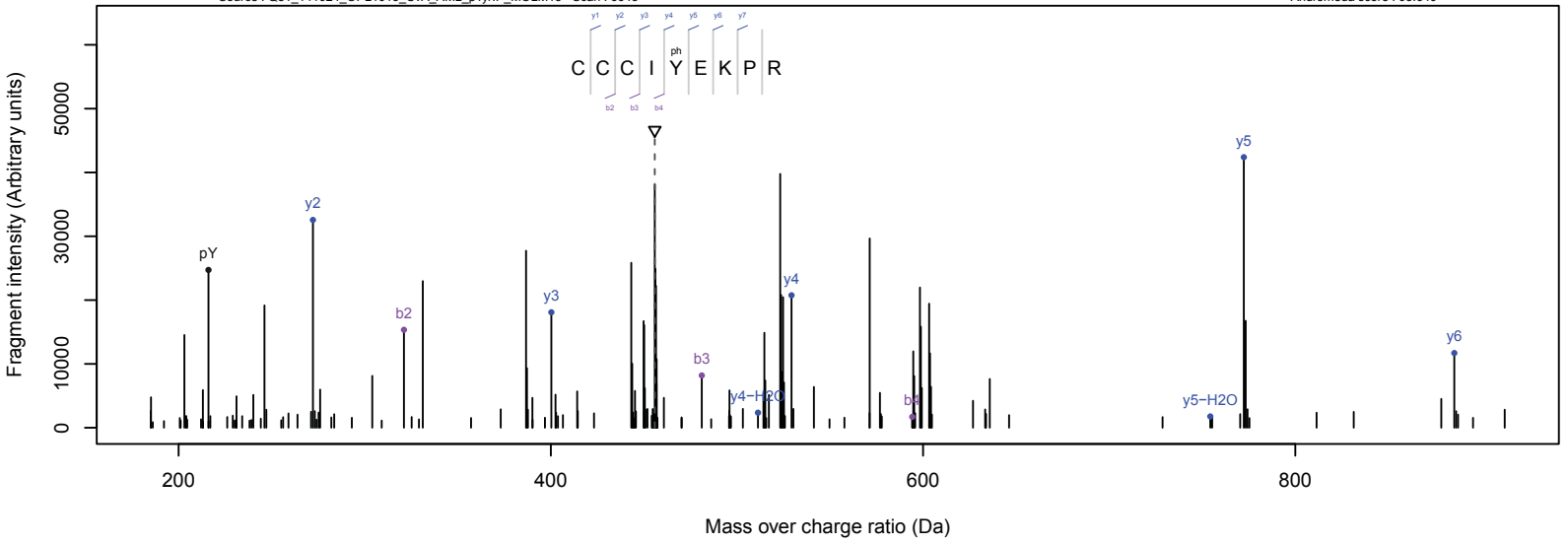
Andromeda score : 93.766



CCCIY(ph)EKPR-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 6643

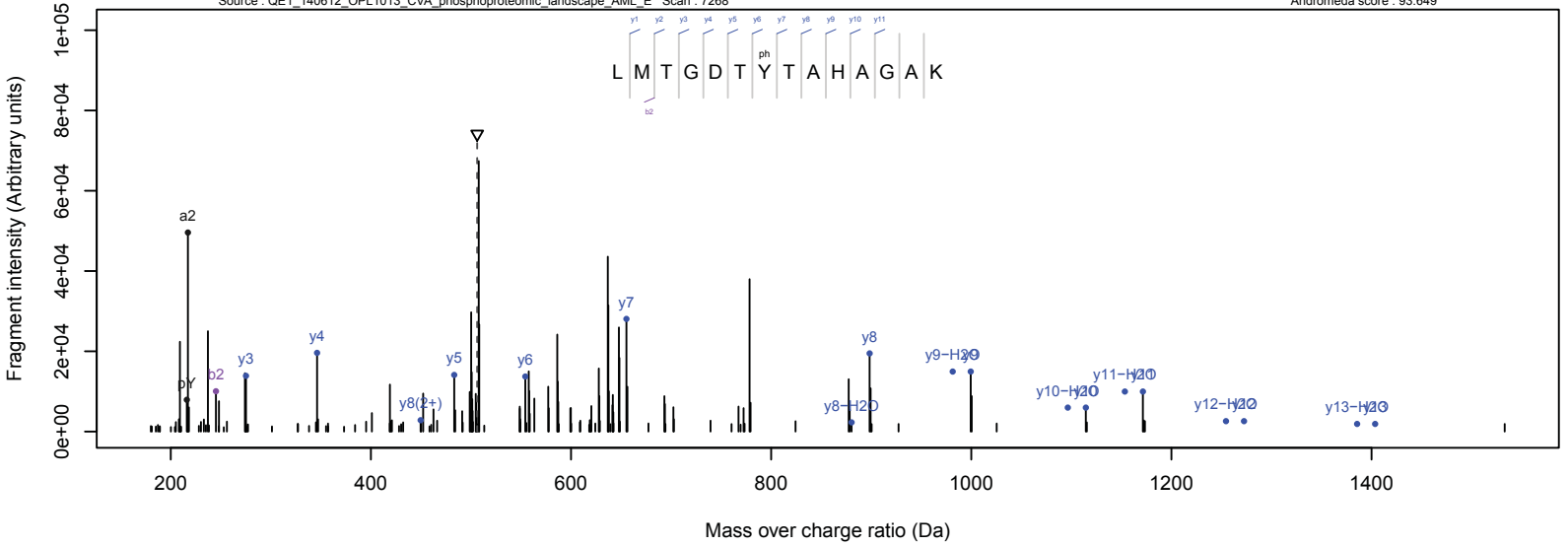
Andromeda score : 93.649



LMTGDTY(ph)TAHAGAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 7268

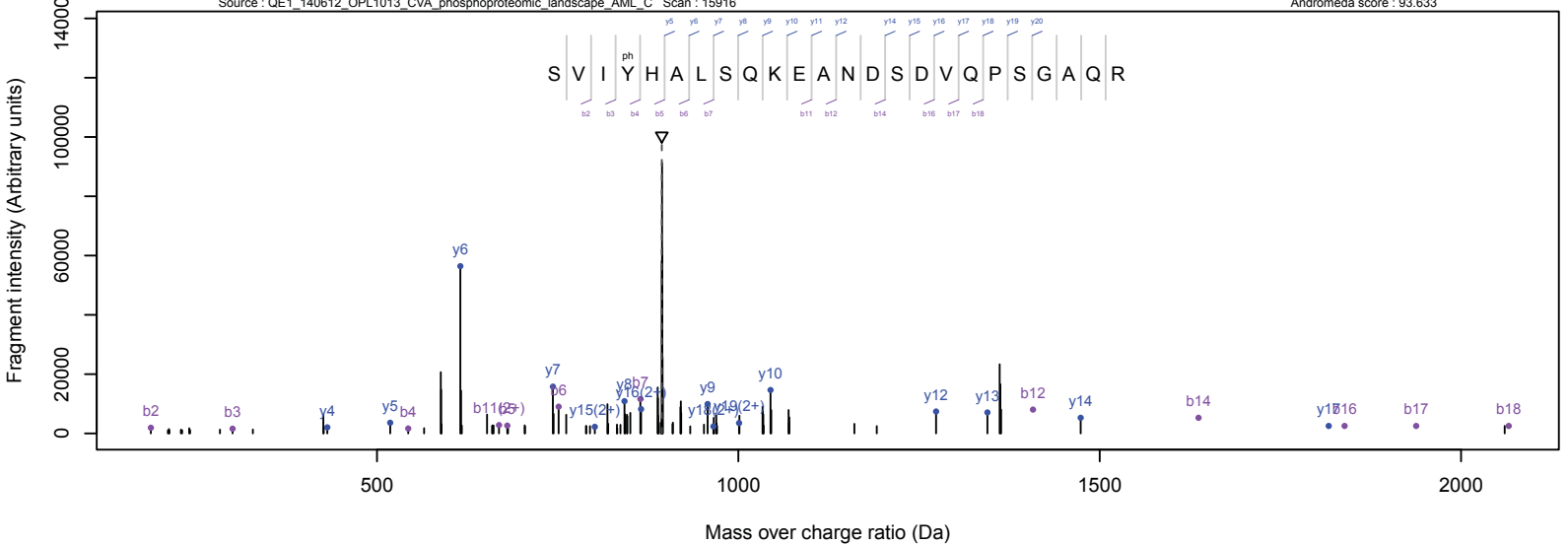
Andromeda score : 93.649



SVIY(ph)HALSQKEANDSDVQPSGAQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15916

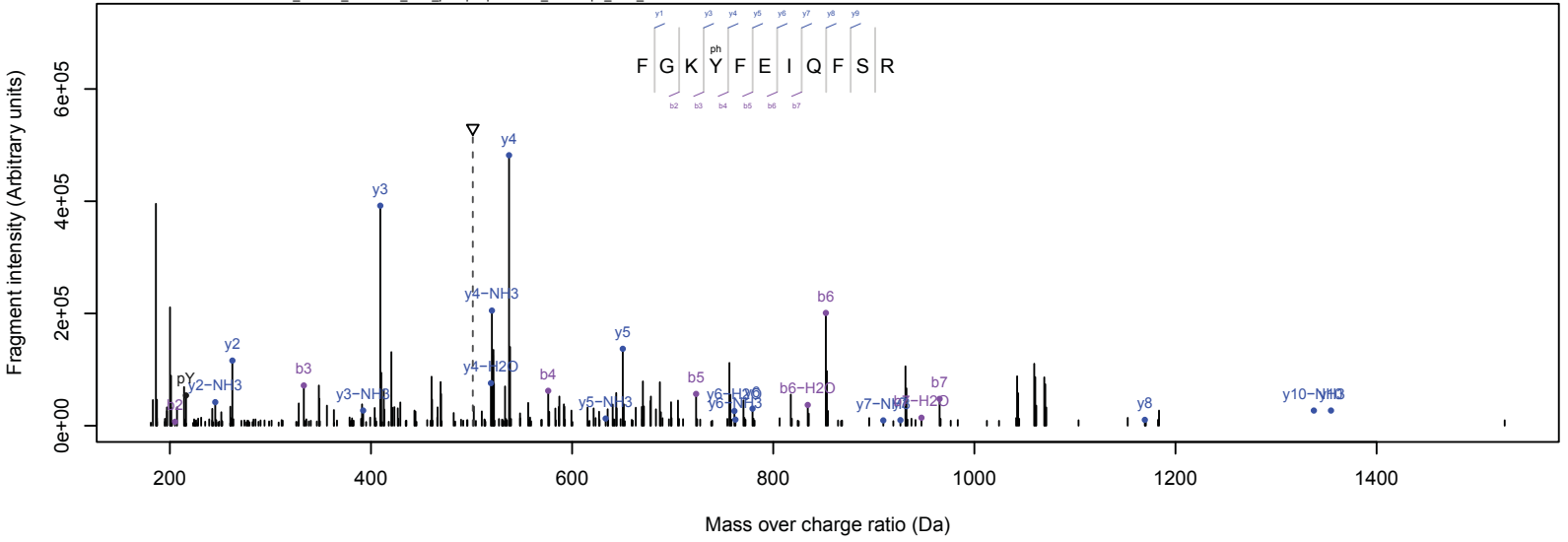
Andromeda score : 93.633



**FGKY(ph)FEIQFSR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34030

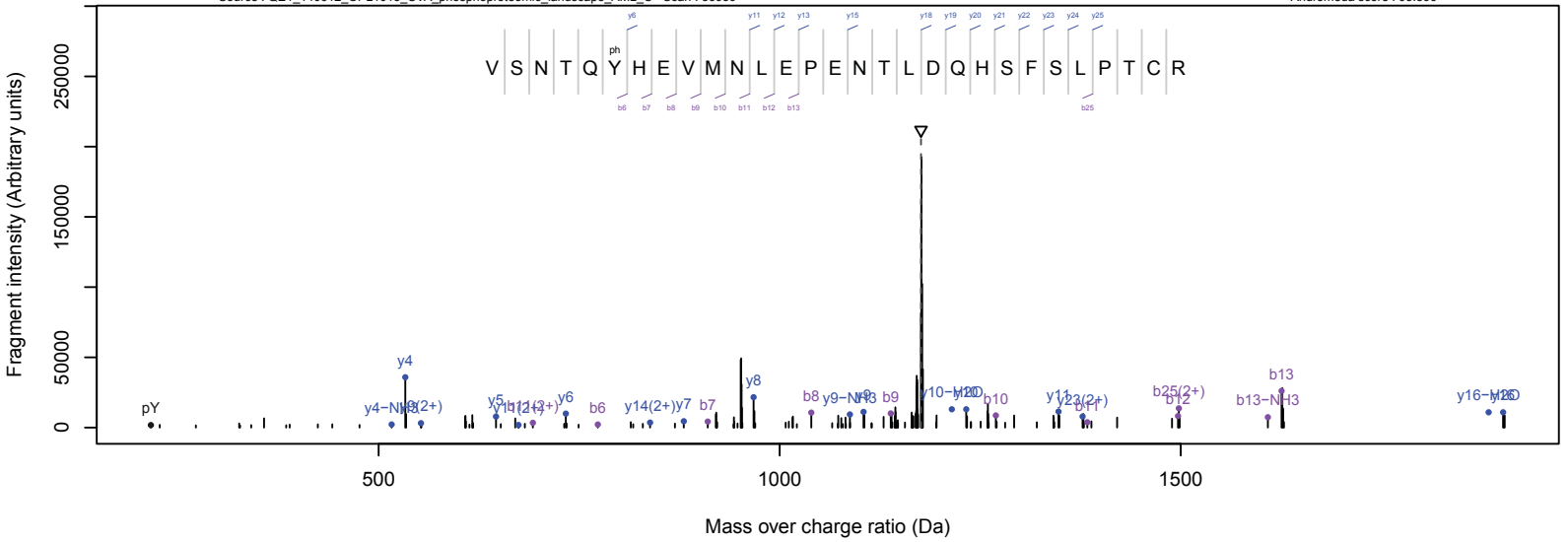
Andromeda score: 93.614



**VSNTQY(ph)HEVMNLEPENTLDQHSFSLPTCR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36939

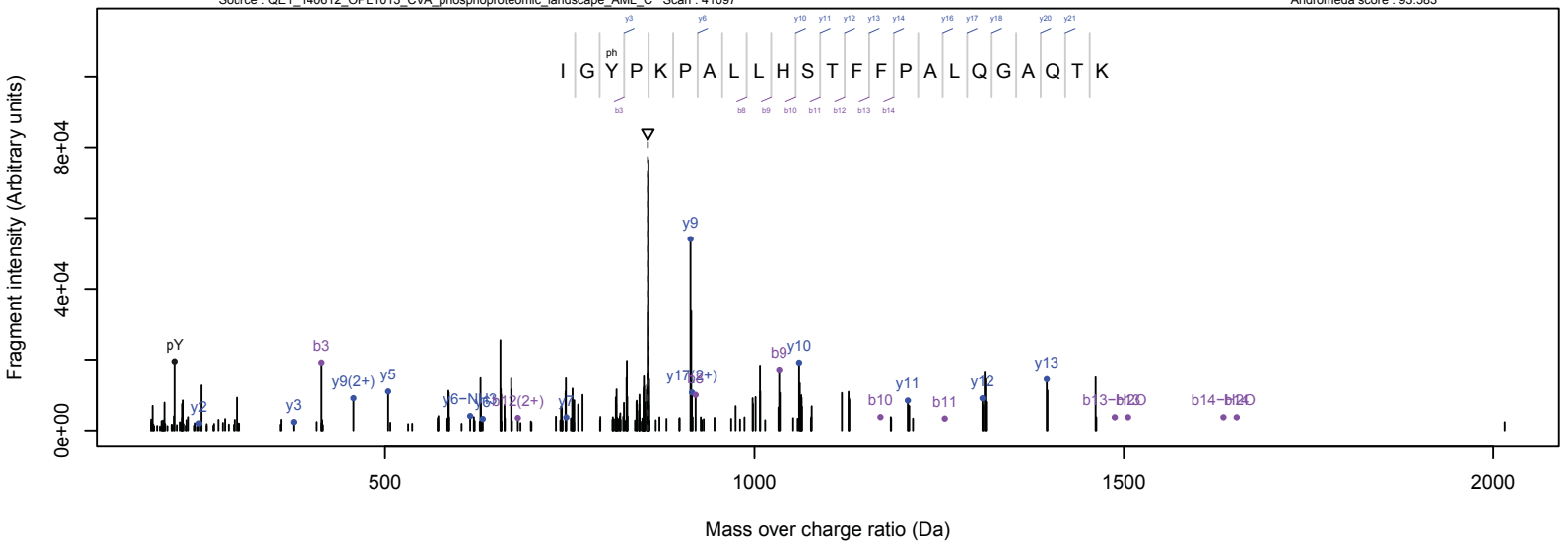
Andromeda score: 93.596



**IGY(ph)PKPALLHSTFFPALQGAQTK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41097

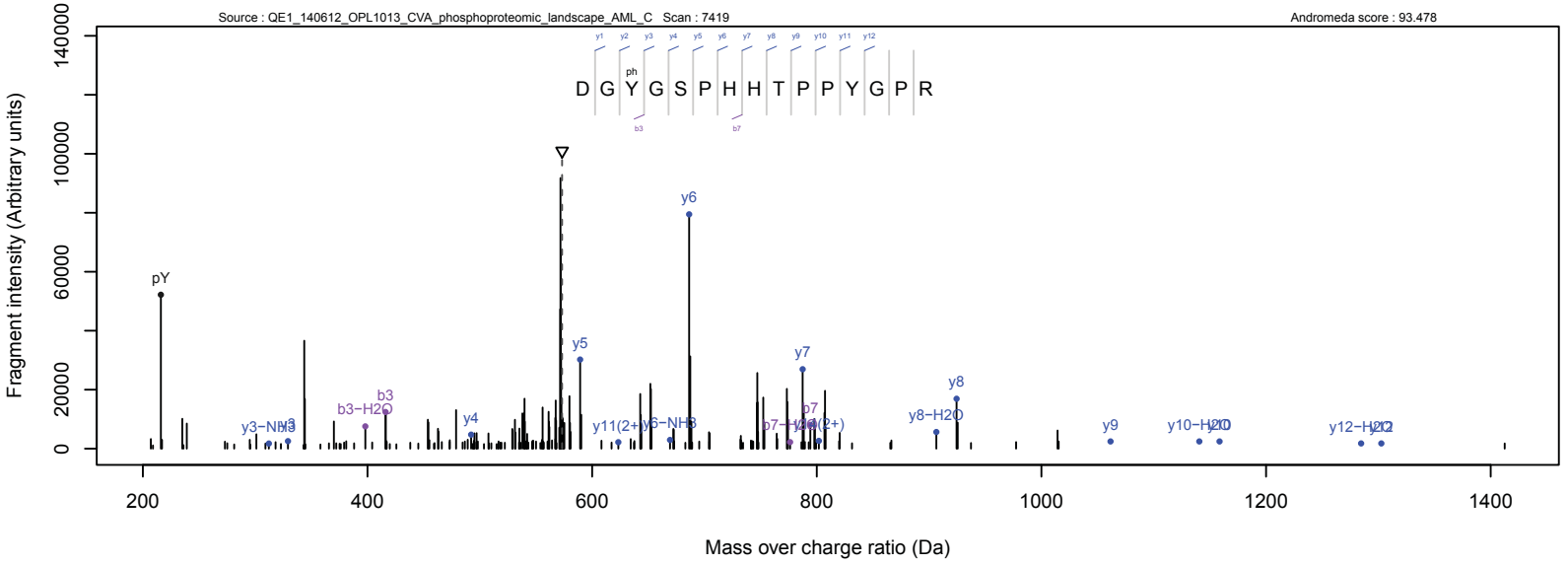
Andromeda score: 93.583



DGY(ph)GSPHHTPPYGPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 7419

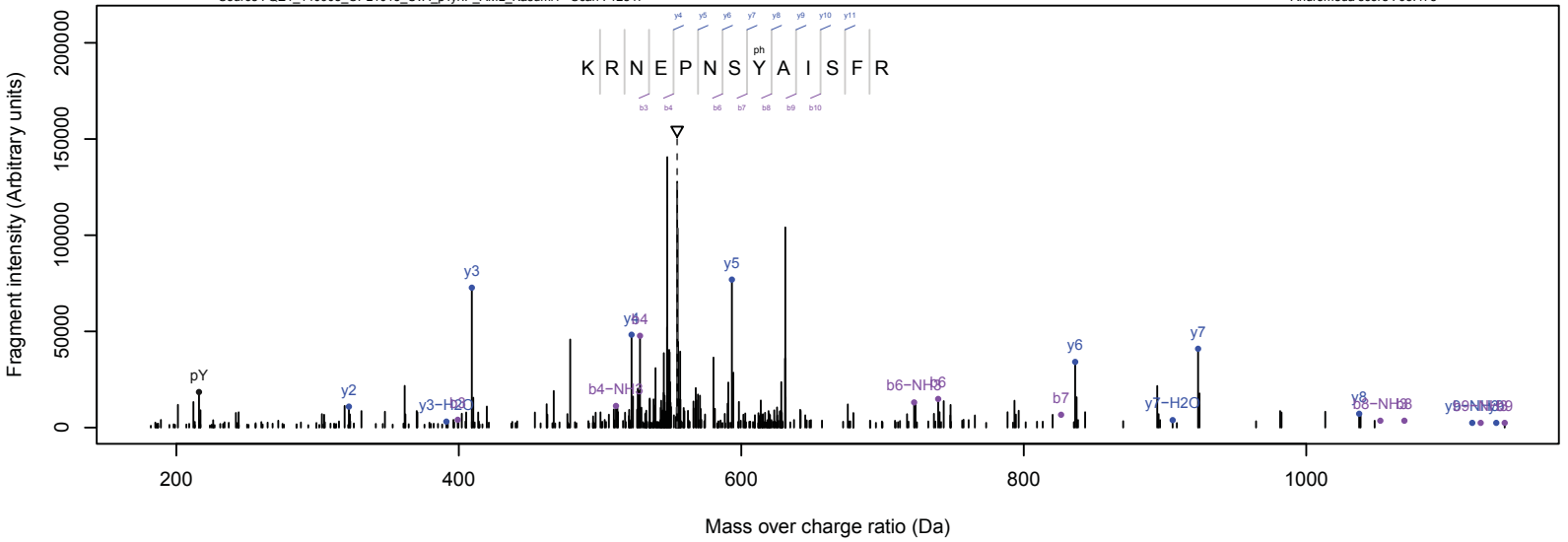
Andromeda score: 93.478



KRNEPNS(ph)AISFR-3+

Source: QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_Kasumi1 Scan: 12847

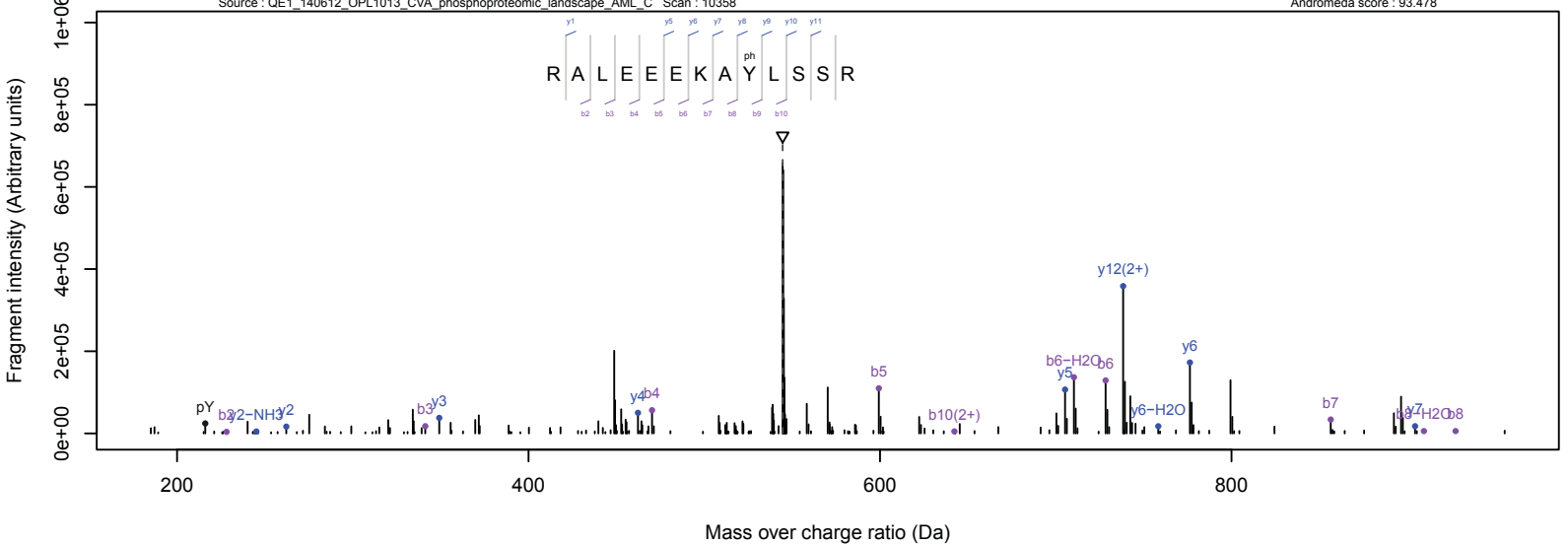
Andromeda score: 93.478



RALEEEKAY(ph)LSSR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10358

Andromeda score: 93.478

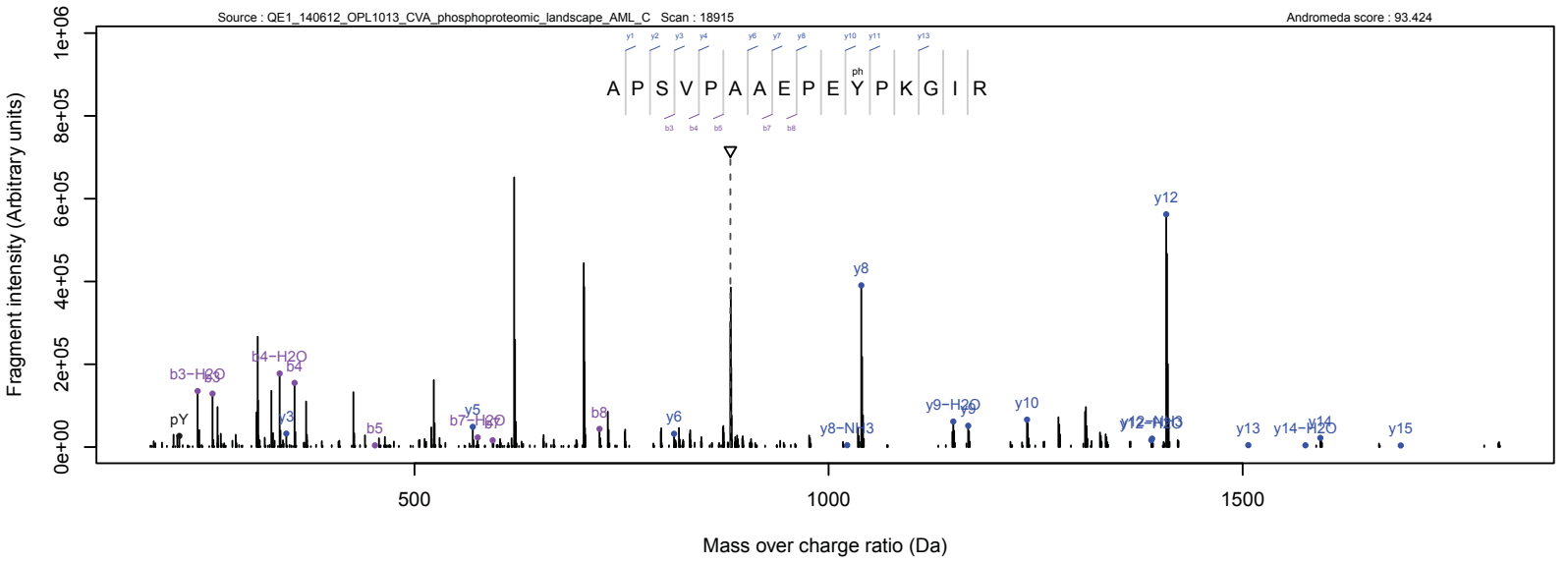




APSVPAEPEY(ph)PKGIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18915

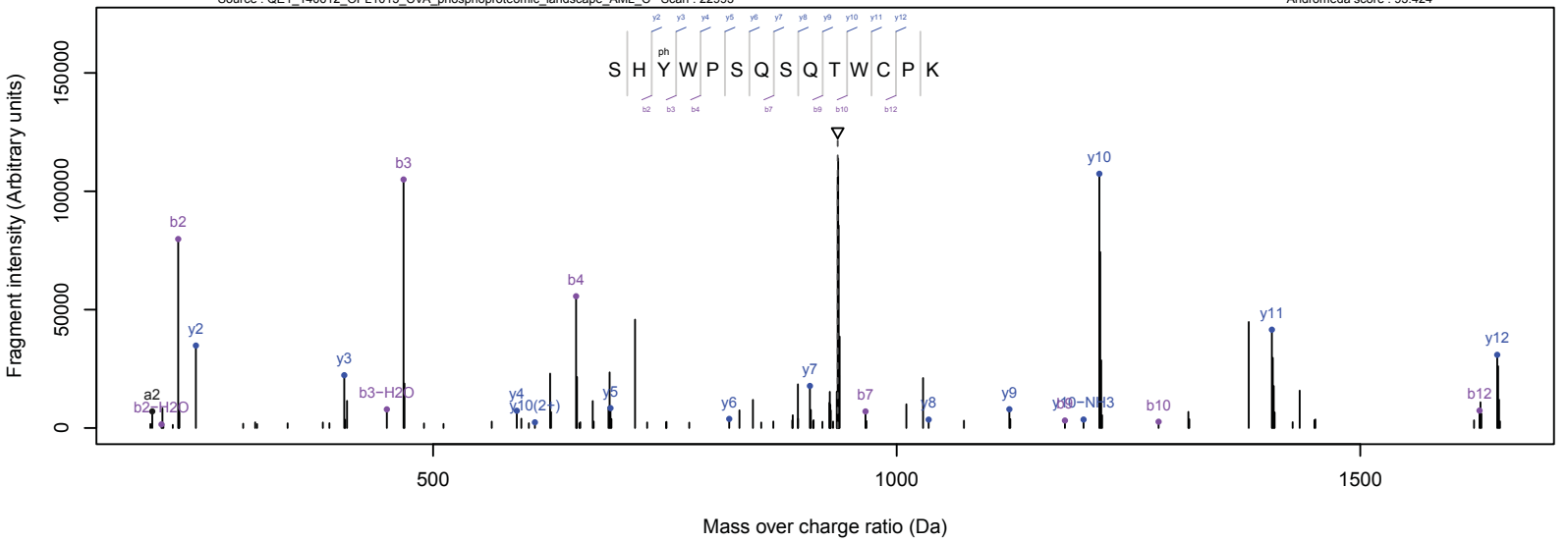
Andromeda score : 93.424



SHY(ph)WPSQQTWCPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22993

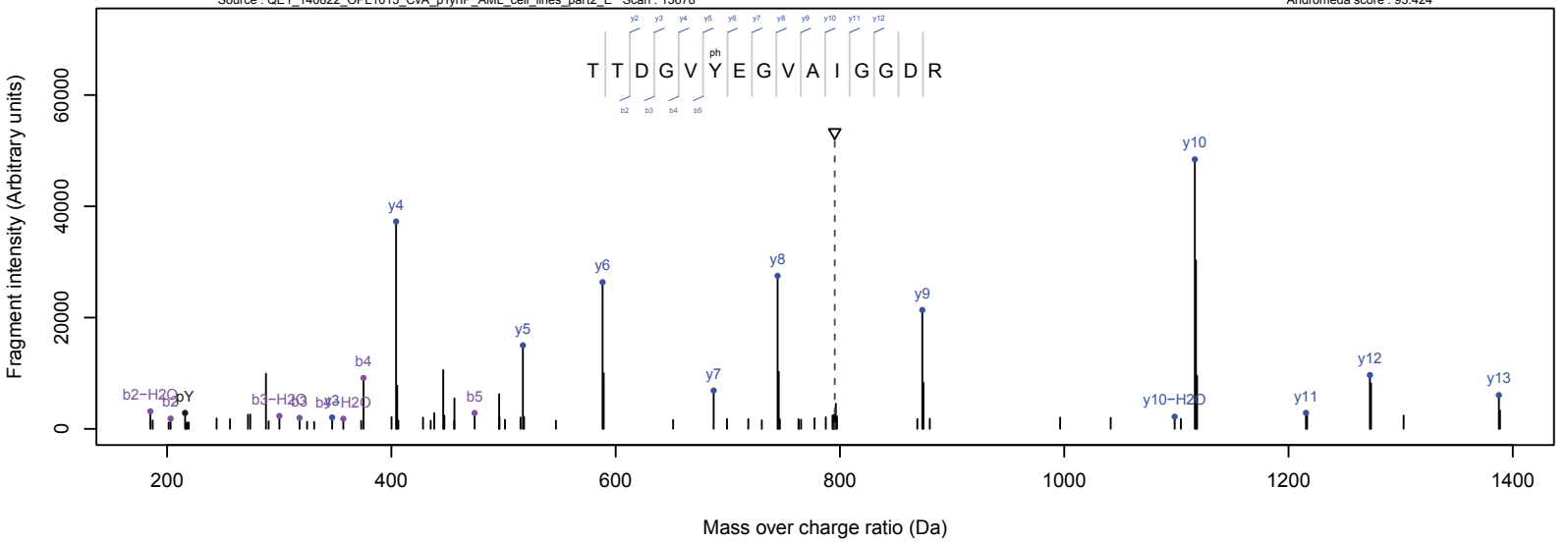
Andromeda score : 93.424



TTDGVY(ph)EGVAIGDR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 13678

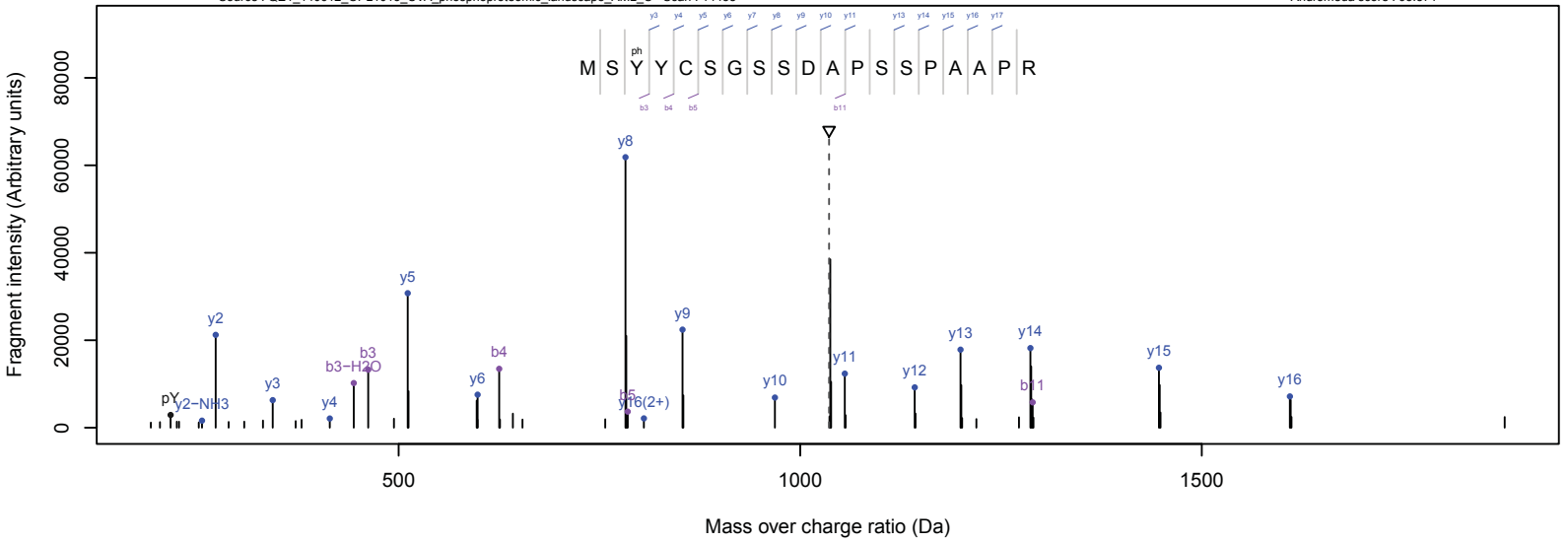
Andromeda score : 93.424



**MSY(ph)YCSGSSDAPSSPAAPR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14458

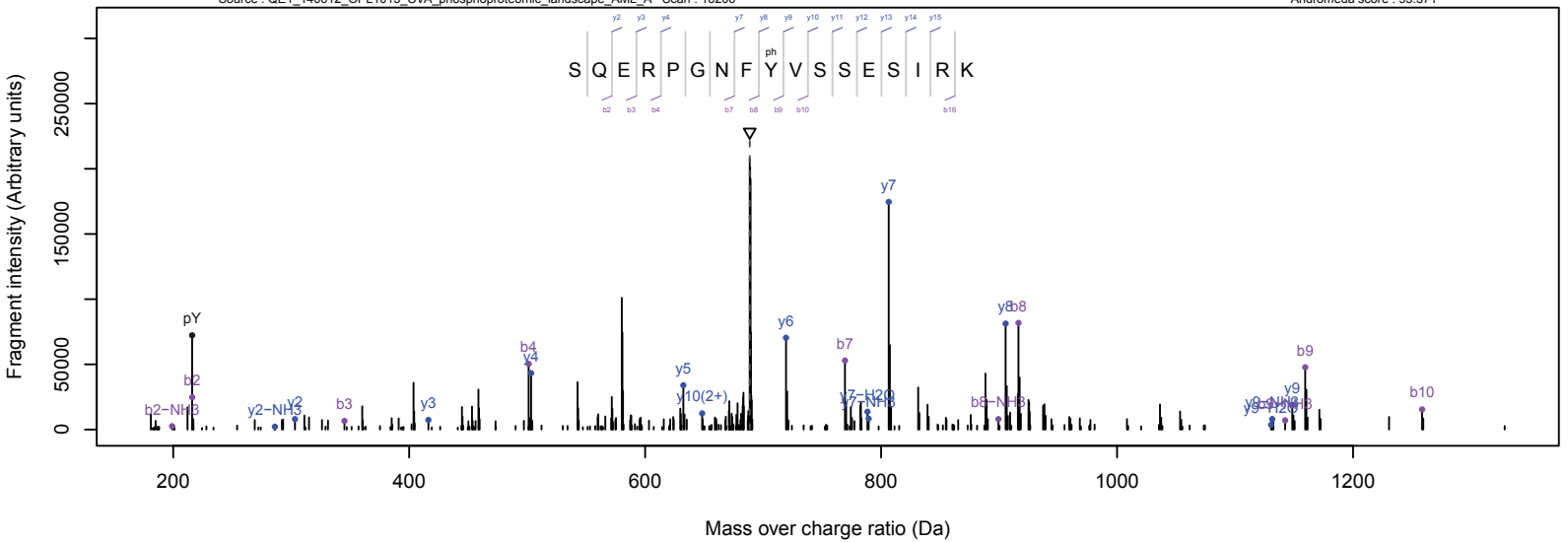
Andromeda score : 93.371



**SQERPGNFY(ph)VSSSIRK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16206

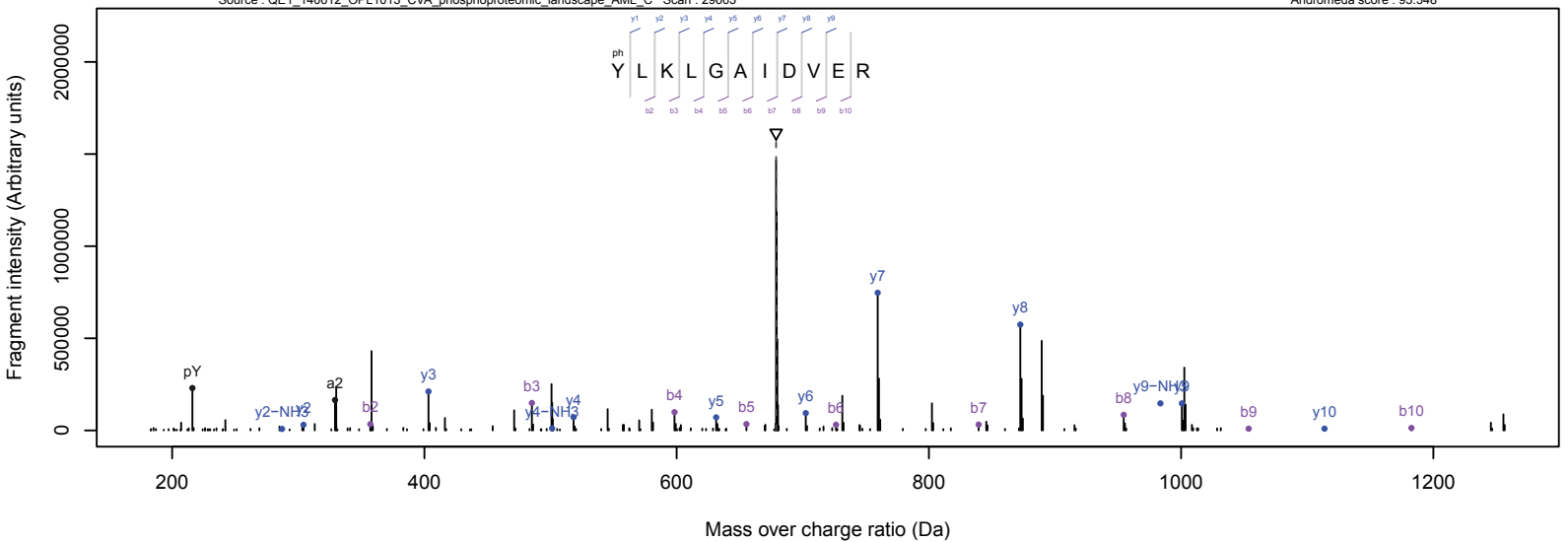
Andromeda score : 93.371



**Y(ph)LKLGAI DVER-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29663

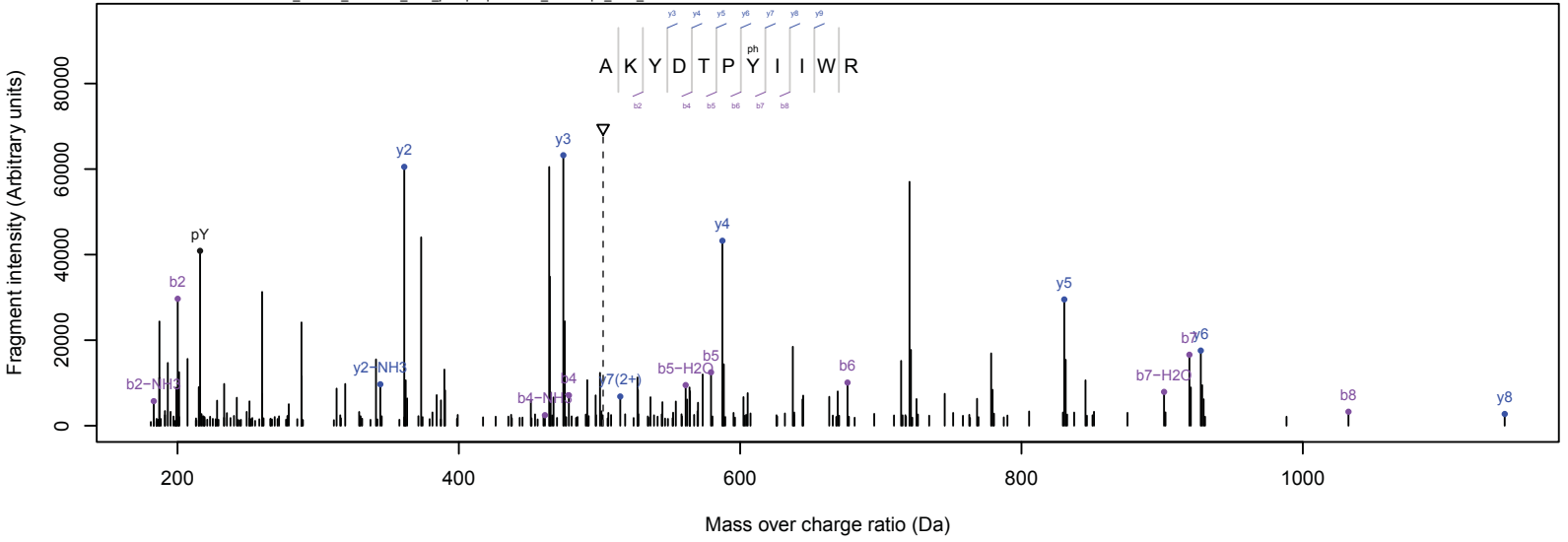
Andromeda score : 93.348



**AKYDTPY(ph)IWR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31697

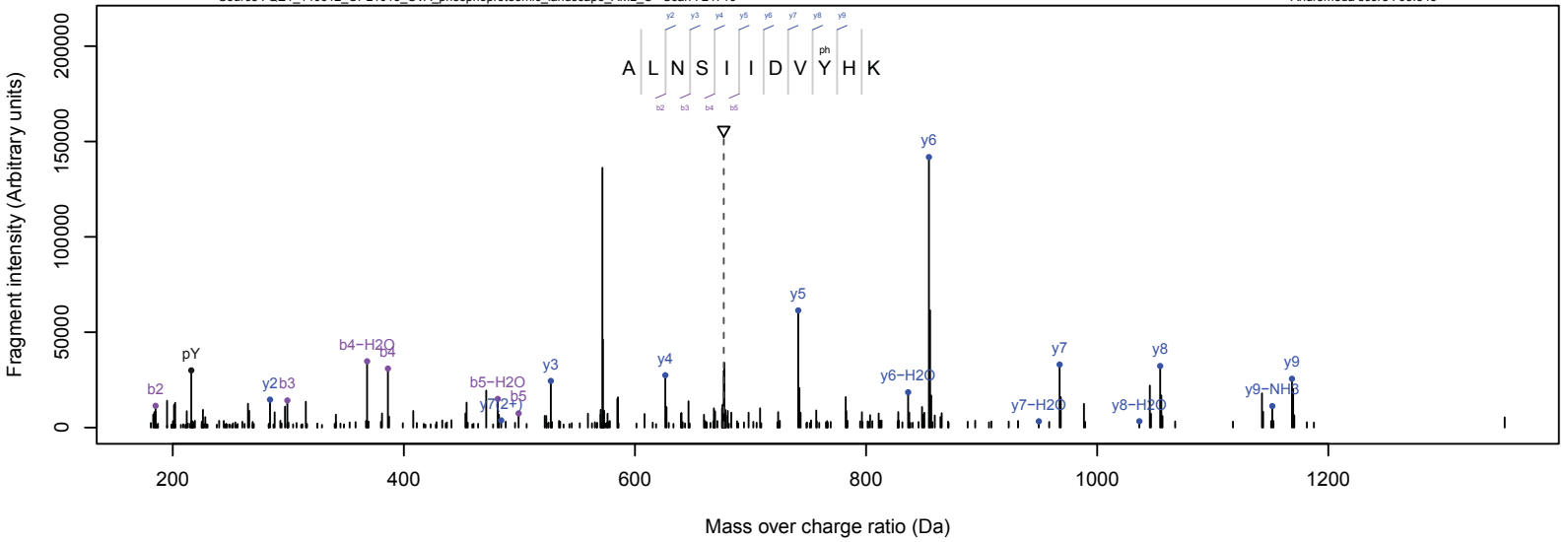
Andromeda score : 93.345



**ALNSIIDVY(ph)HK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24716

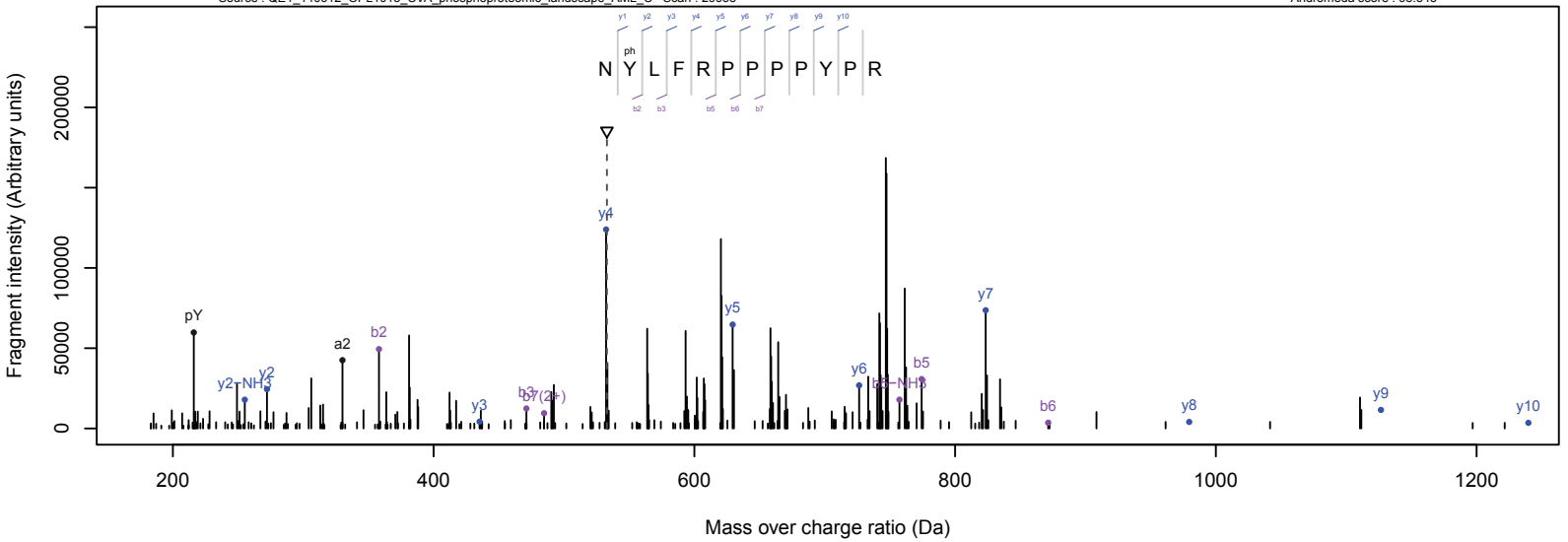
Andromeda score : 93.345



**NY(ph)LFRPPPPYR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29058

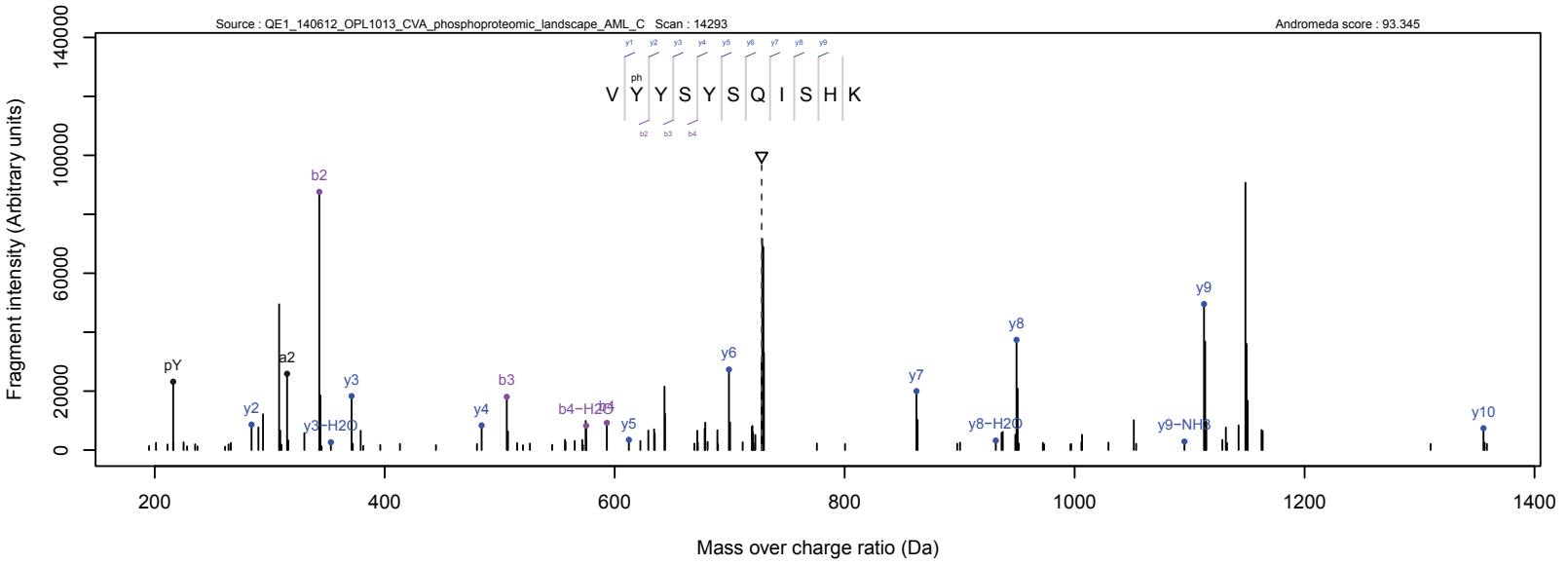
Andromeda score : 93.345



VY(ph)YSYSQISHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14293

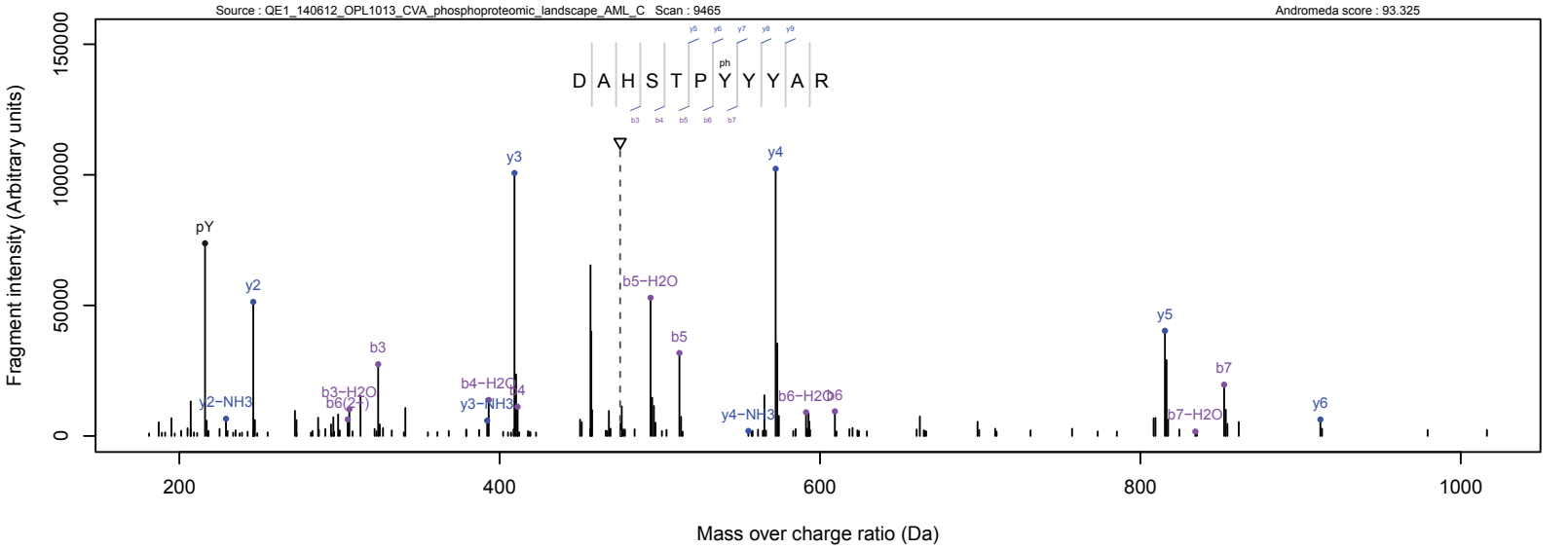
Andromeda score : 93.345



DAHSTPY(ph)YYAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9465

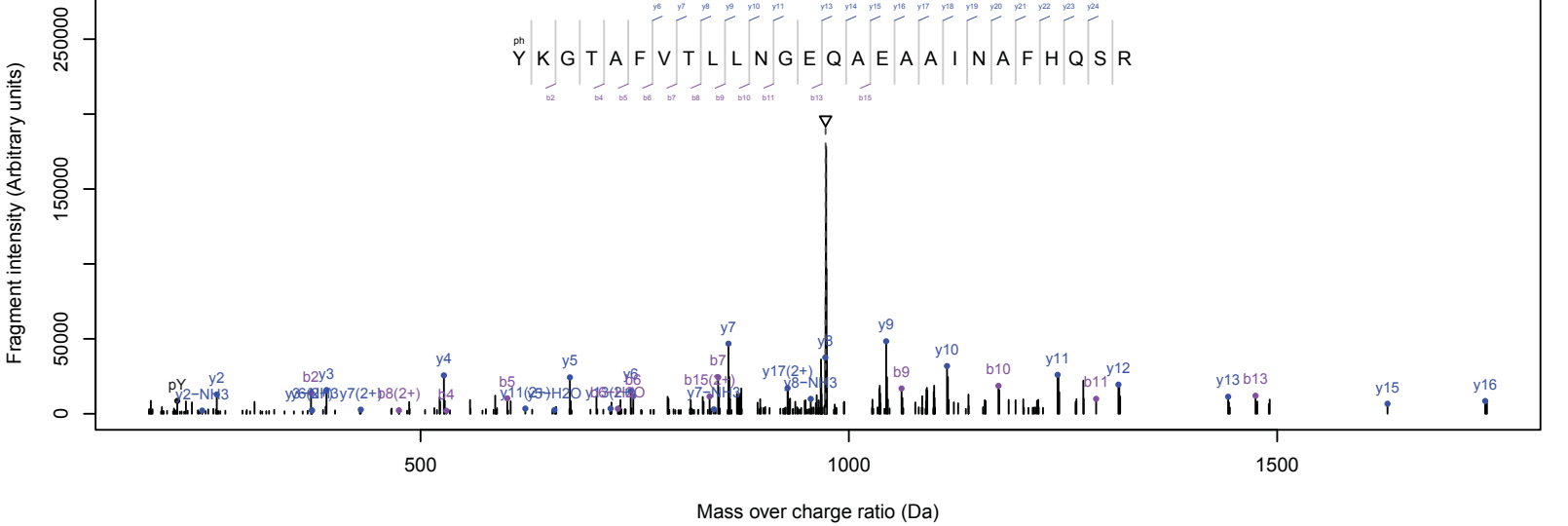
Andromeda score : 93.325



Y(ph)KGTAFTLLNGEQAEAAINAFHQSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45412

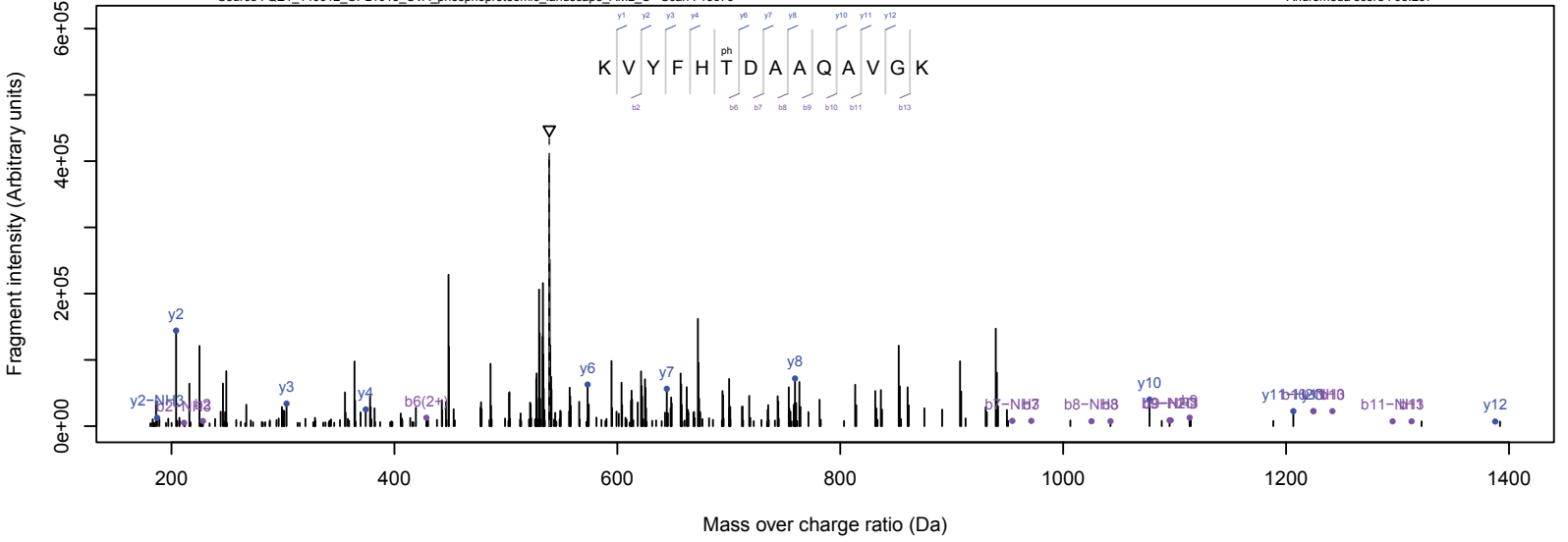
Andromeda score : 93.278



**KVYFHT(ph)DAAQAVGK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13673

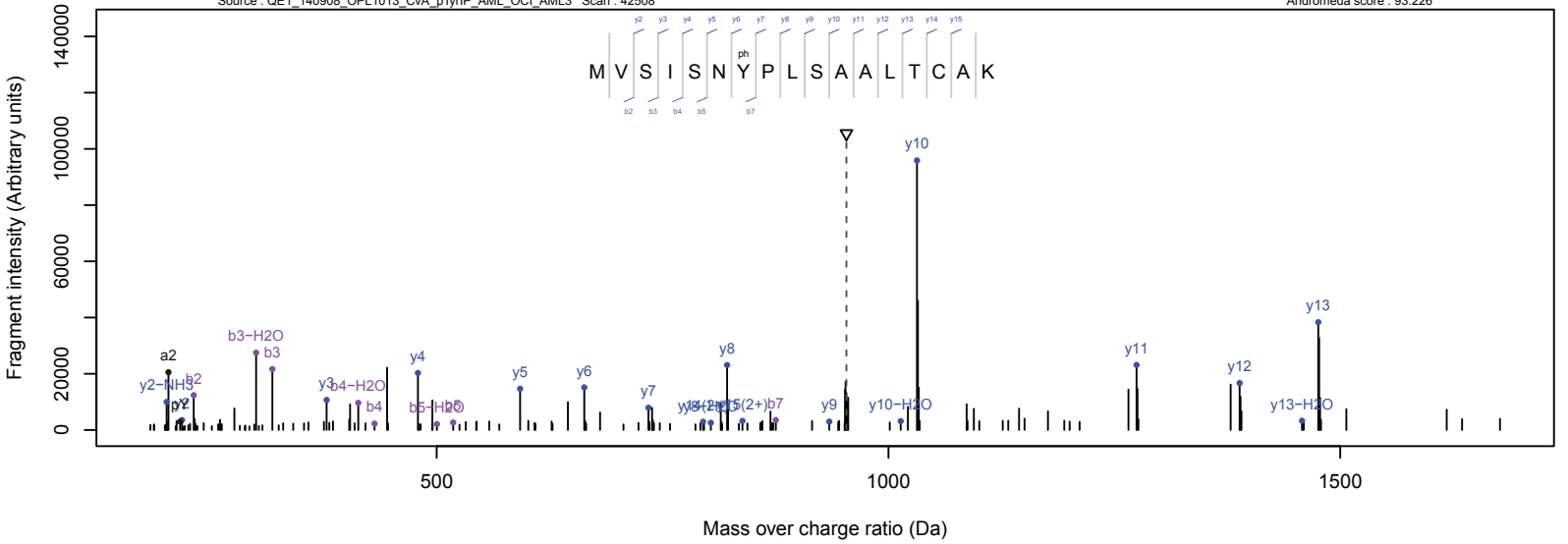
Andromeda score: 93.237



**MVSISNY(ph)PLSAALTCCK-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 42508

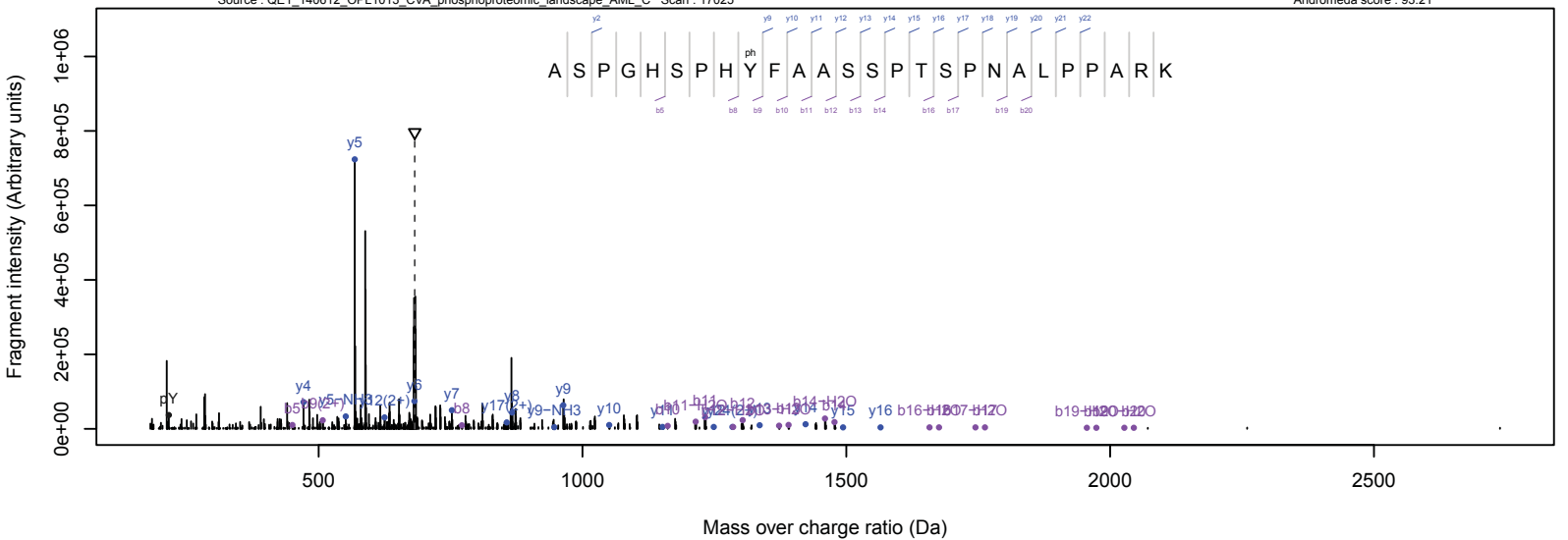
Andromeda score: 93.226



**ASPGHSPHY(ph)FAASPTSPNALPPARK-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17025

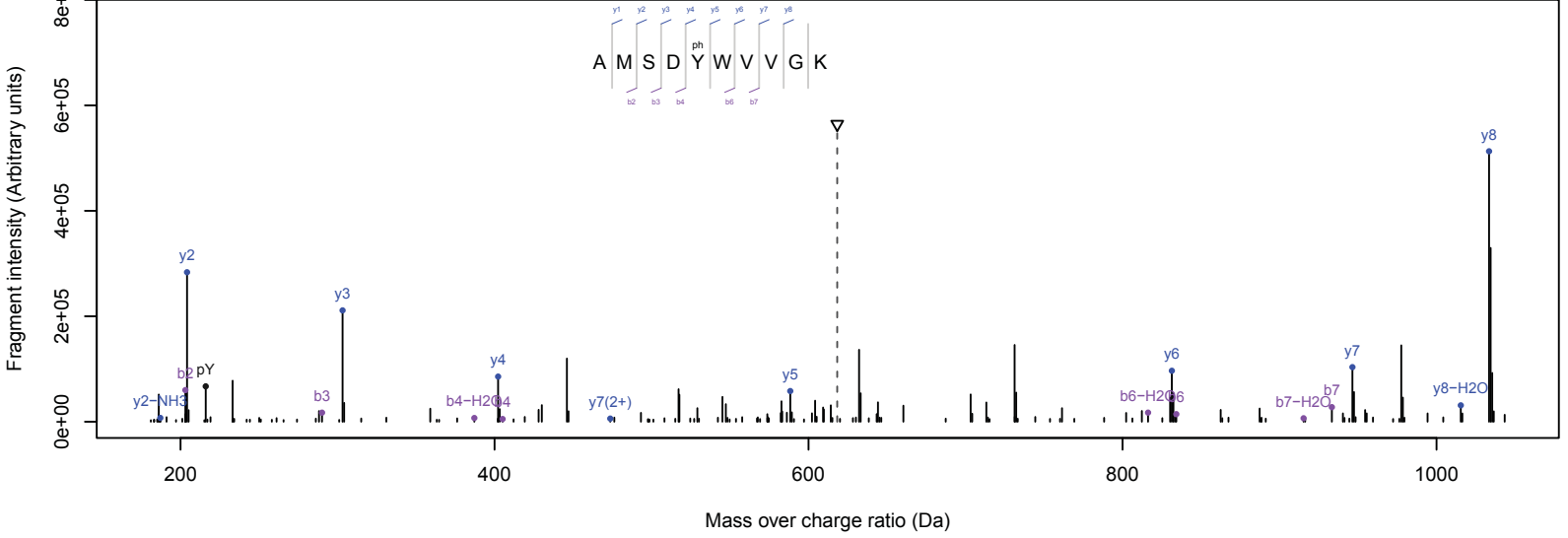
Andromeda score: 93.21



AMSDY(ph)WVVGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29476

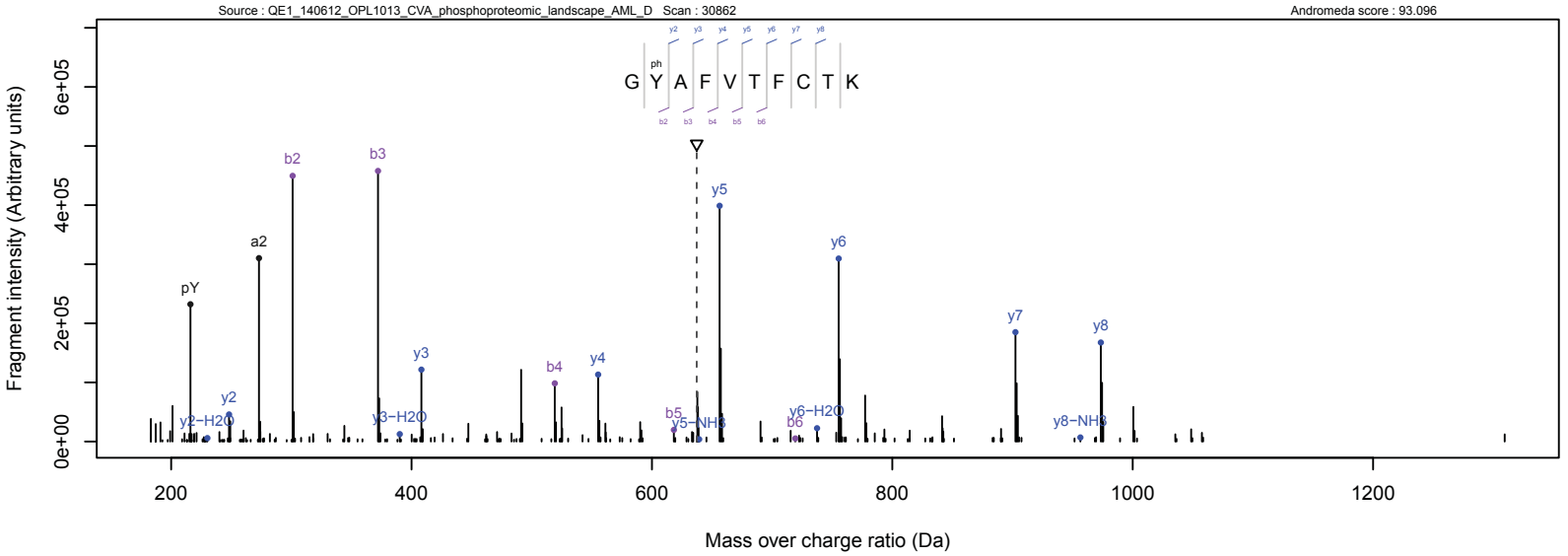
Andromeda score : 93.096



GY(ph)AFVTFCTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 30862

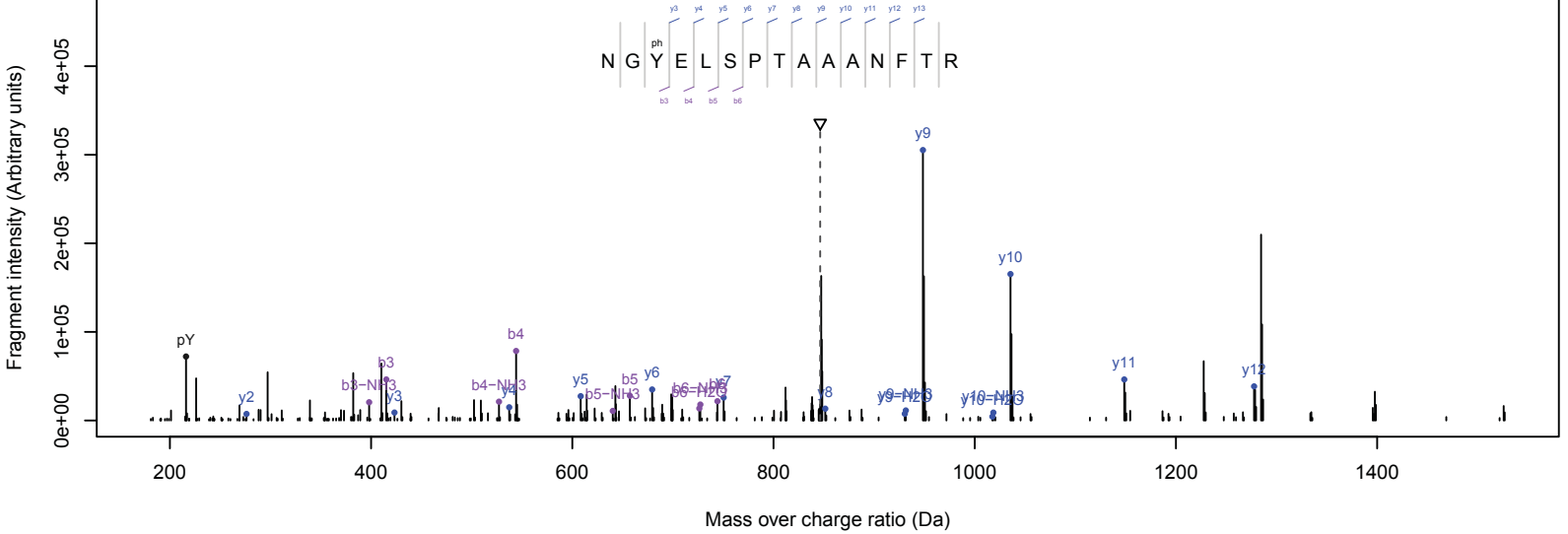
Andromeda score : 93.096



NGY(ph)ELSPAAANFTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27847

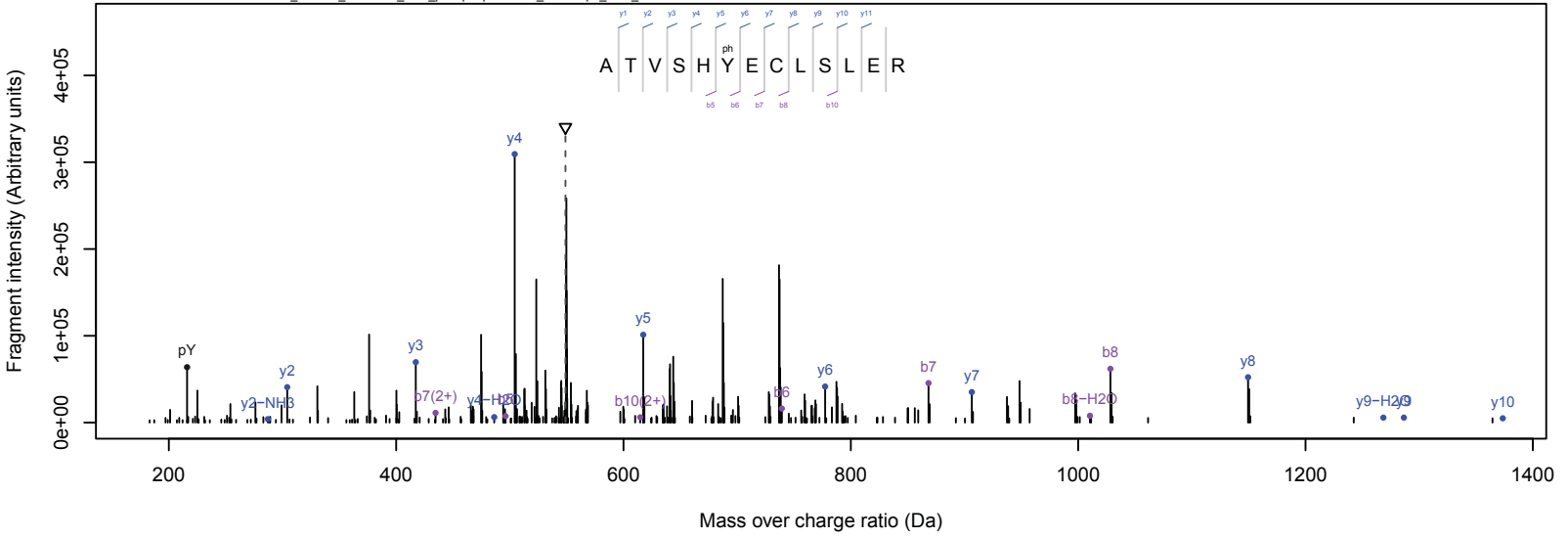
Andromeda score : 92.952



ATVSHY(ph)ECLSLER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19040

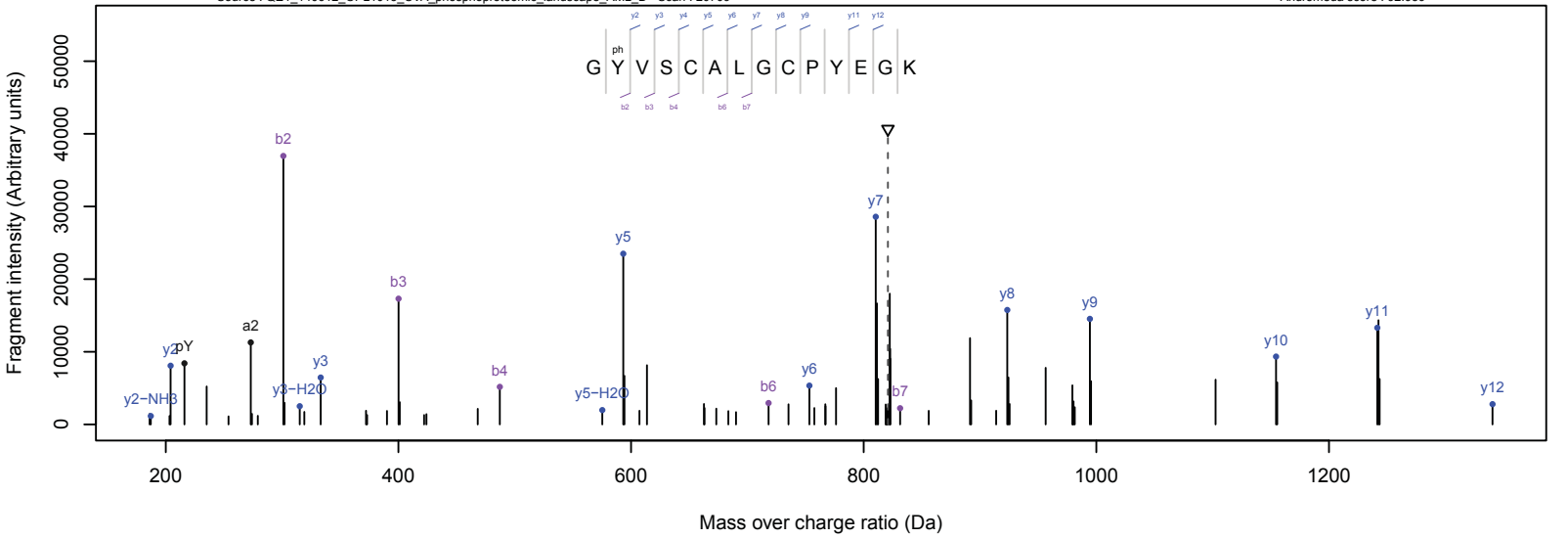
Andromeda score : 92.939



GY(ph)VSCALGCPYEGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 20709

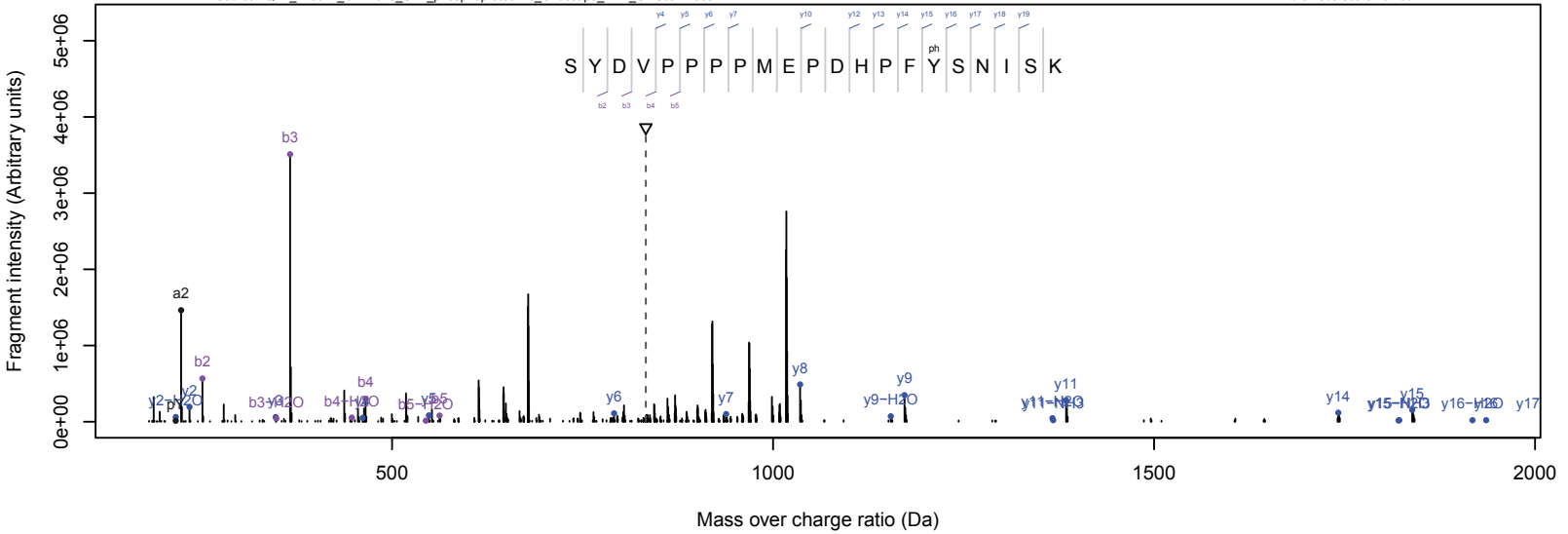
Andromeda score : 92.939



SYDVP P P P M E P D H P F Y S N I S K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33071

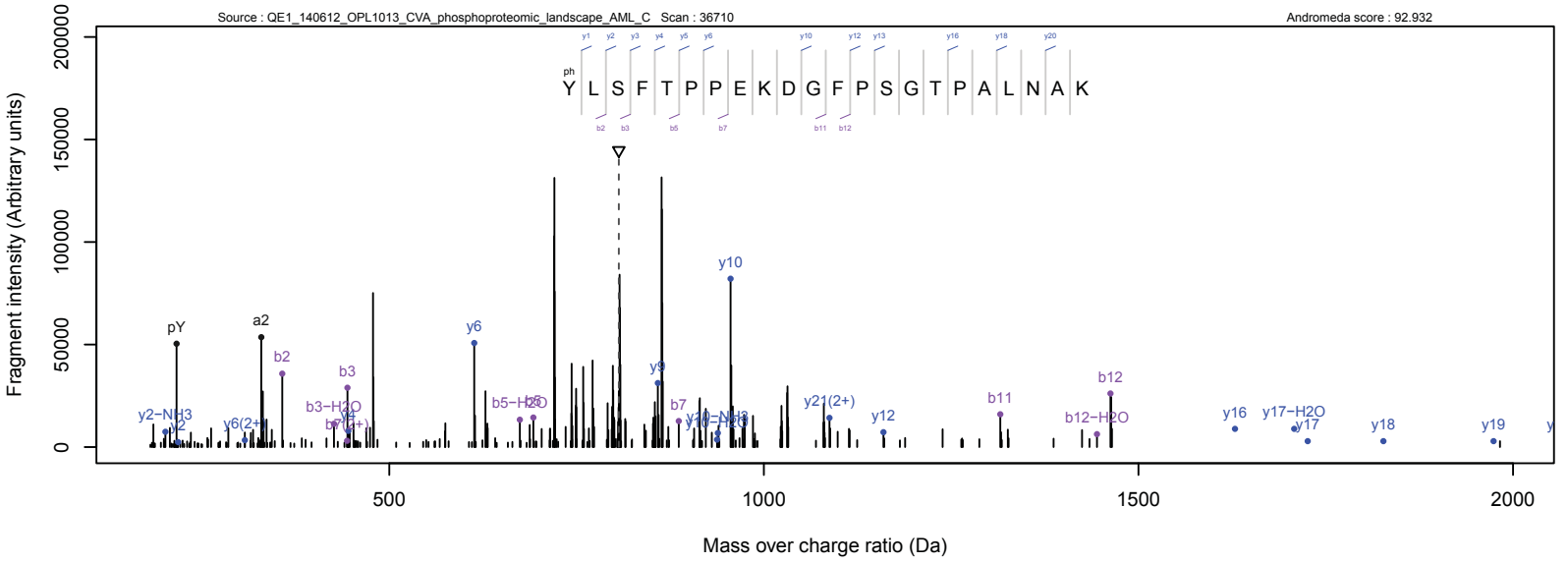
Andromeda score : 92.932



**Y(ph)LSFTPPEKDGFPSTPALNAK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36710

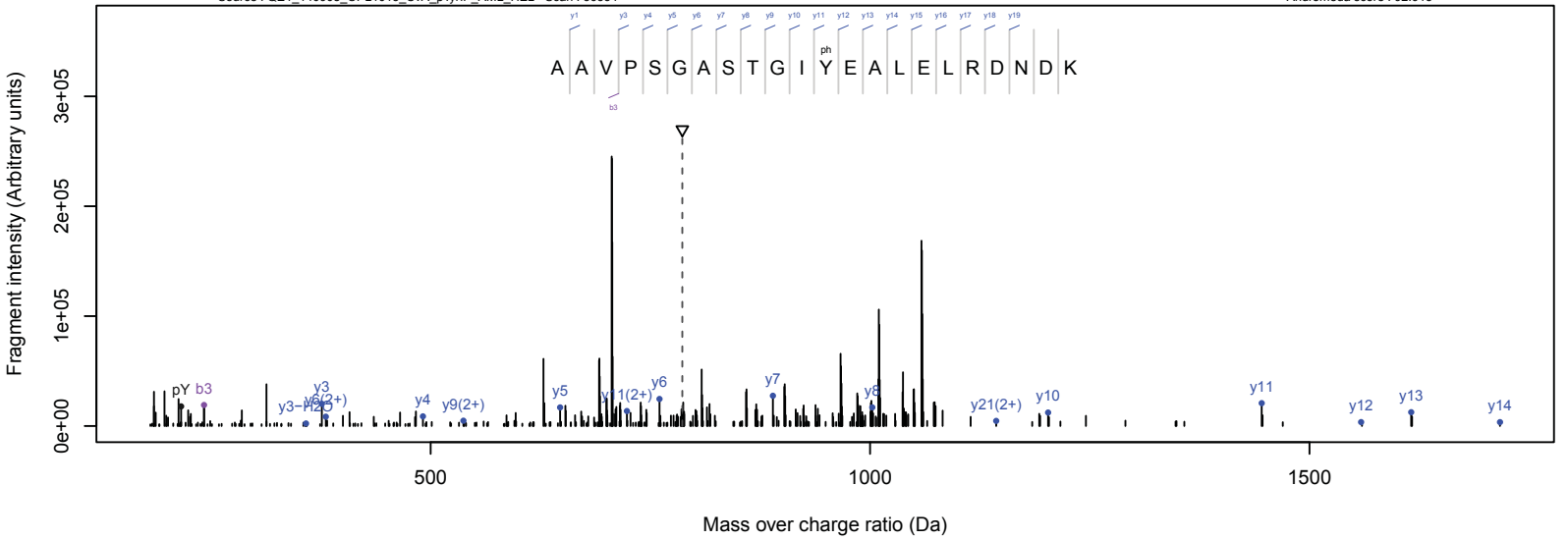
Andromeda score: 92.932



**AAVPSGASTGIY(ph)EALERDNDK-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 36334

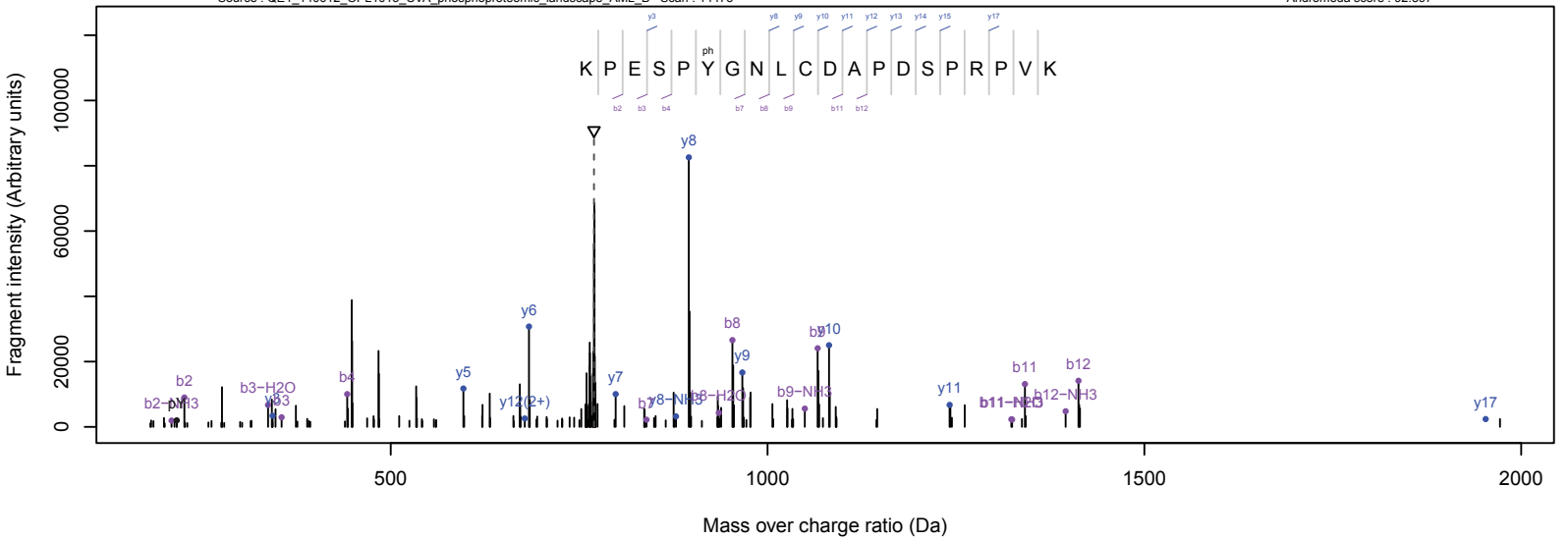
Andromeda score: 92.913



**KPESPY(ph)GNLCDAPDSPRPVK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14475

Andromeda score: 92.897

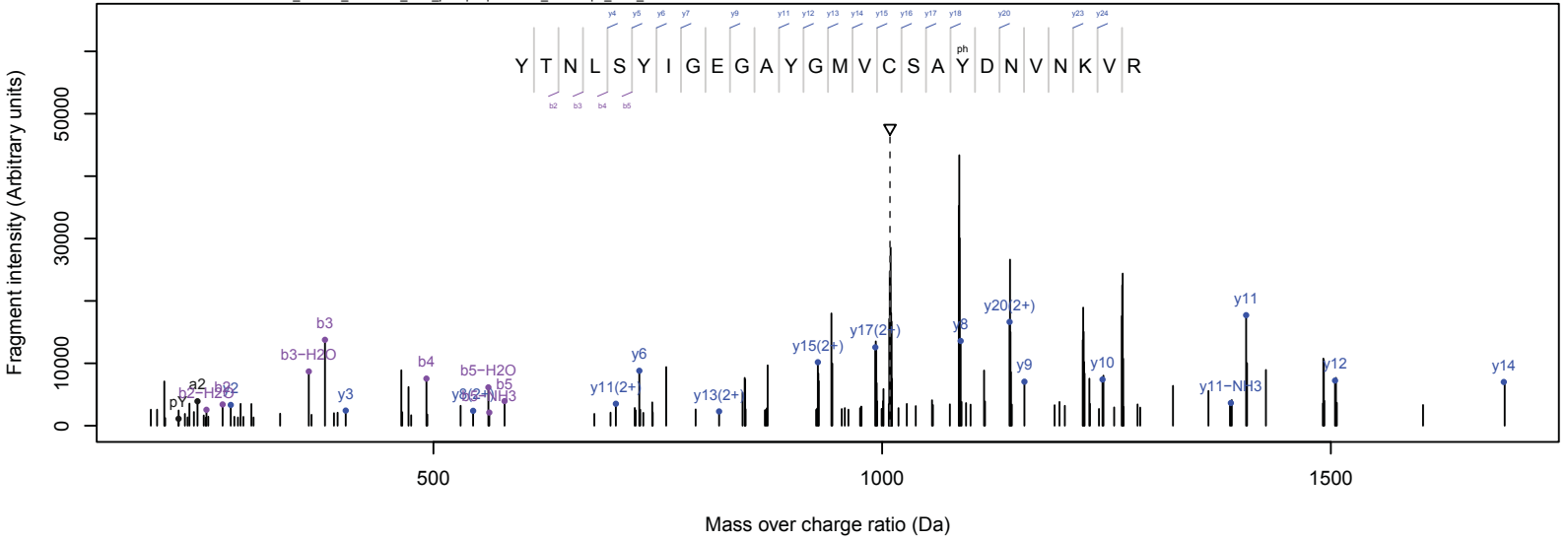




YTNL<sup>ph</sup>SYIGEGAYGMVCSAYD<sup>ph</sup>NVNKVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 42578

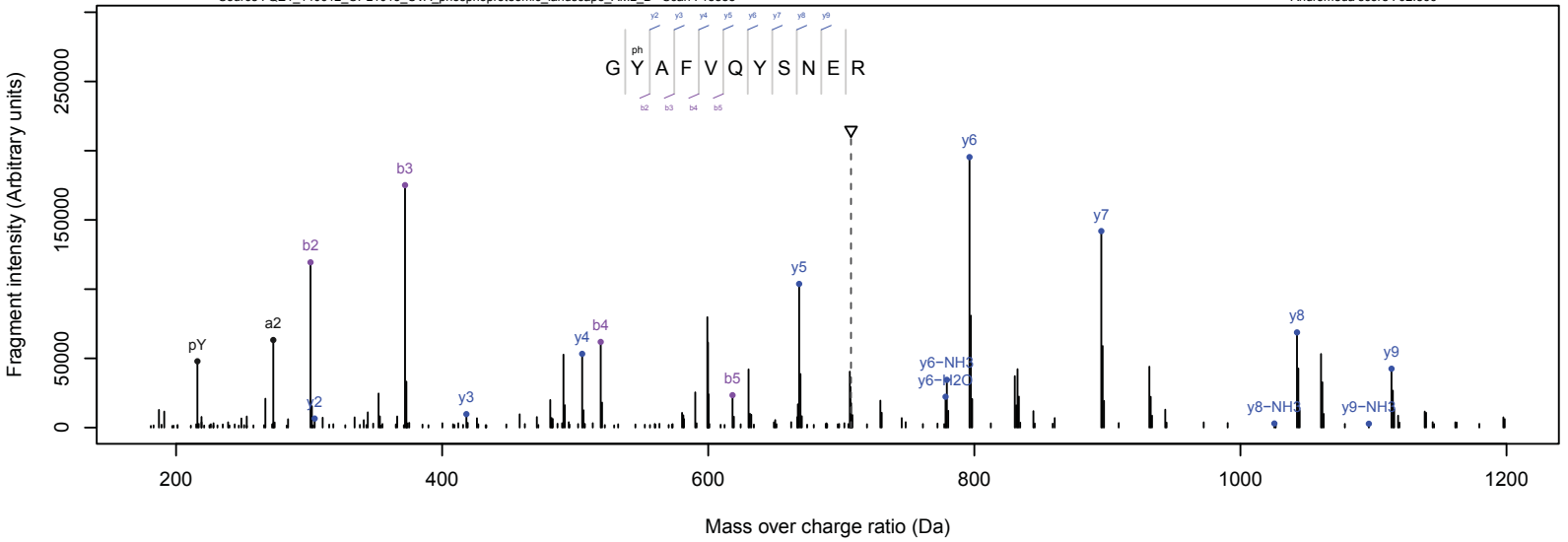
Andromeda score : 92.886



GY<sup>ph</sup>AFVQYSNER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18888

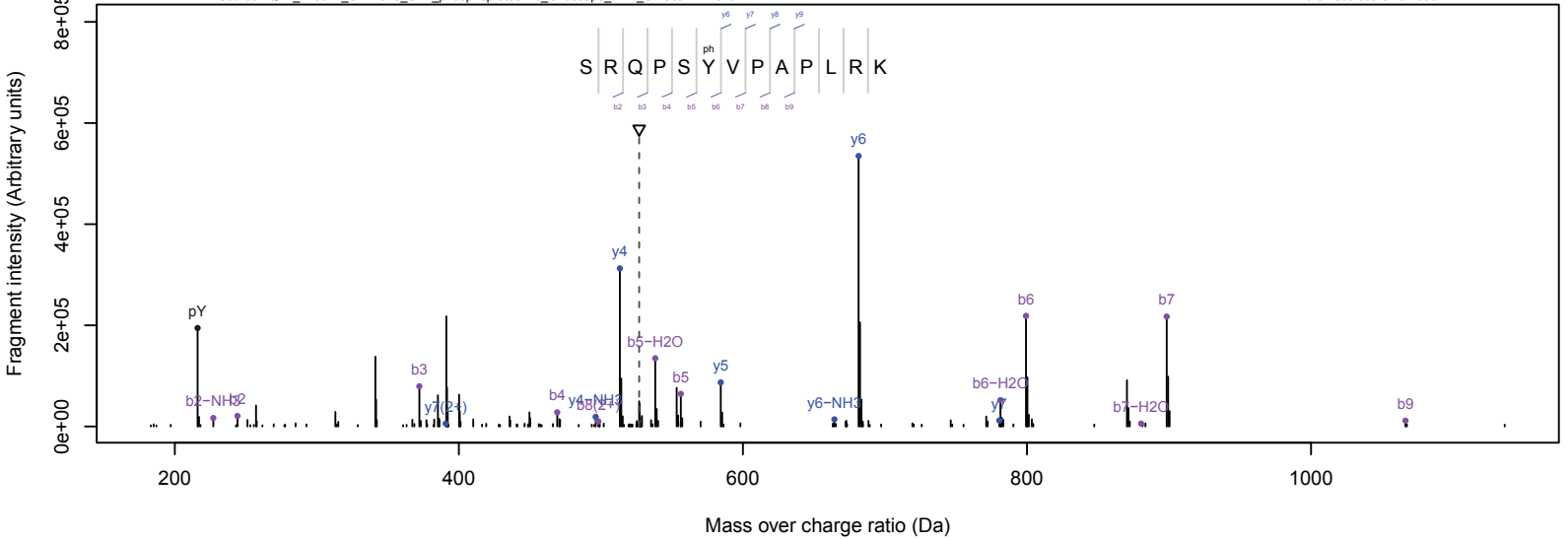
Andromeda score : 92.866



SRQPSY<sup>ph</sup>VPAPLRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12029

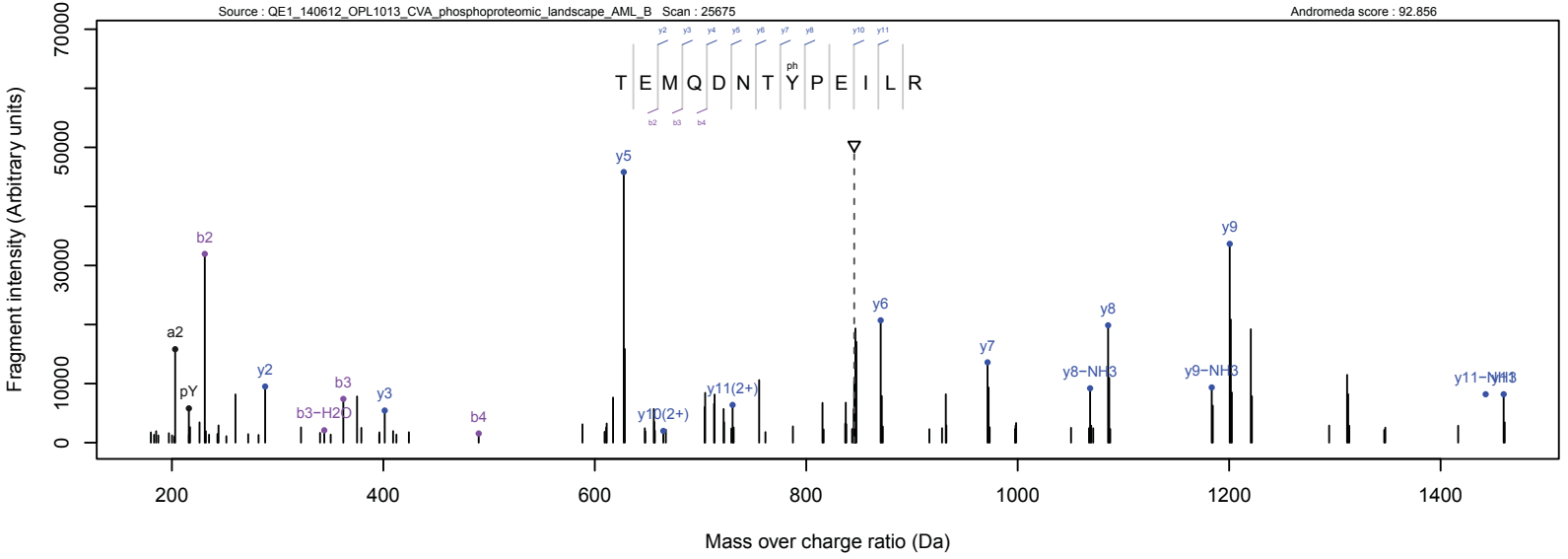
Andromeda score : 92.856



TEMQDNTY(ph)PEILR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25675

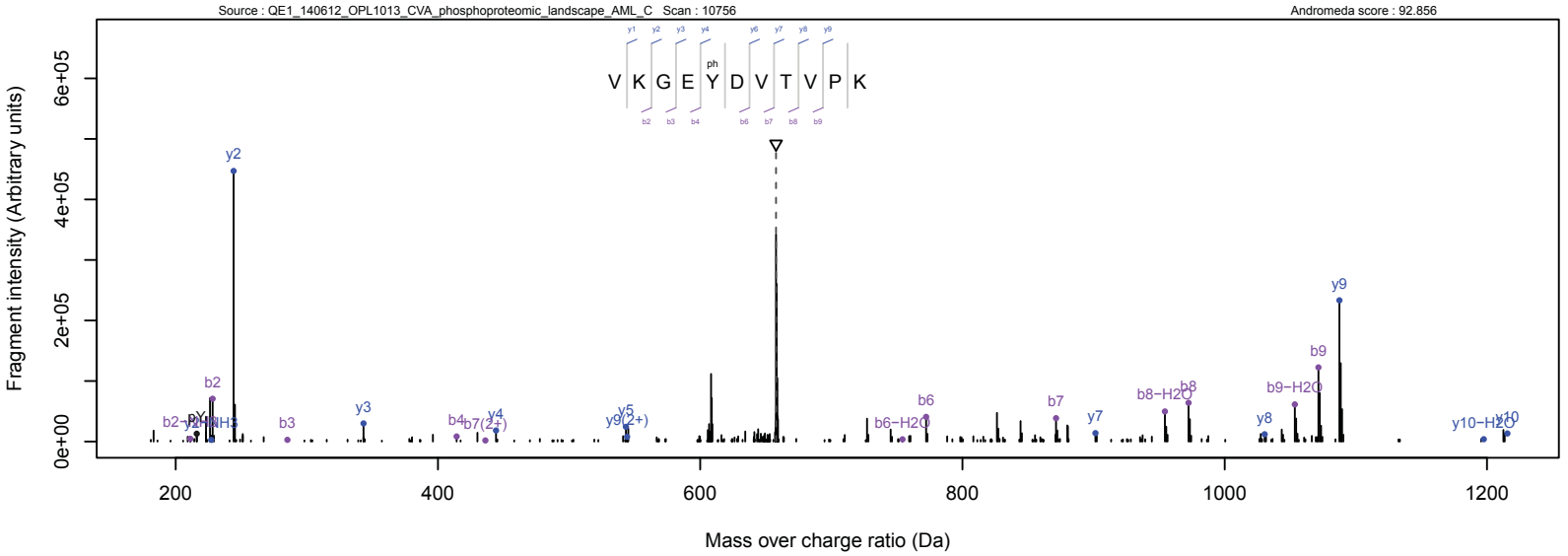
Andromeda score : 92.856



VKGEY(ph)DVTVPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10756

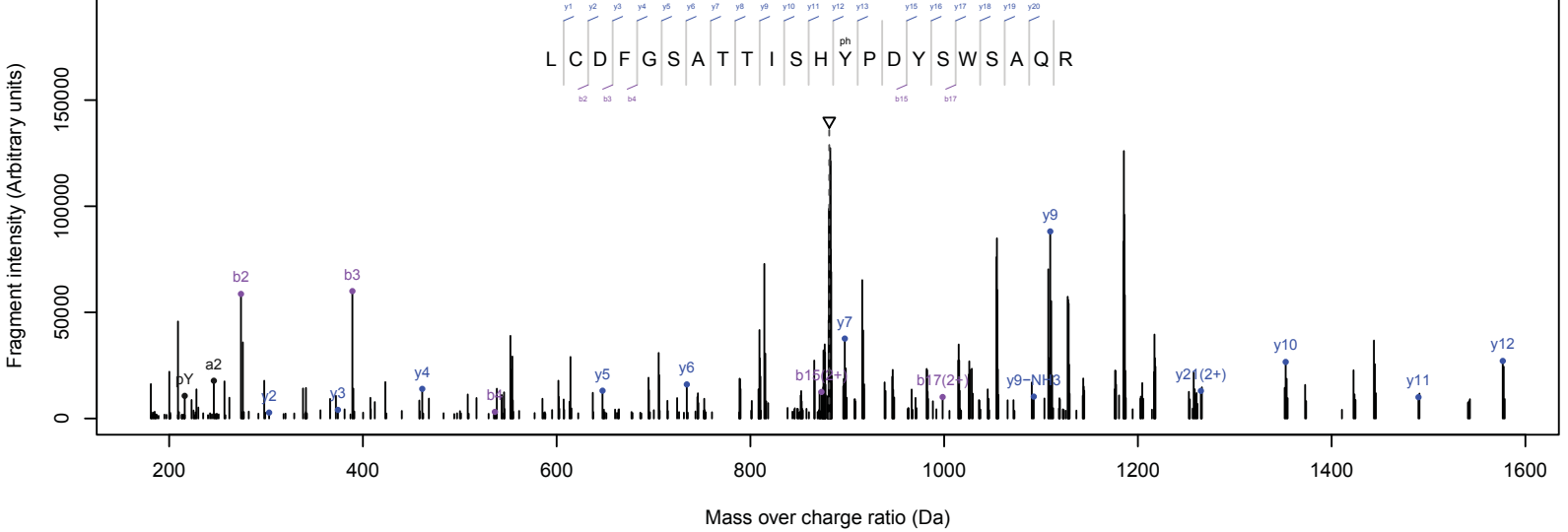
Andromeda score : 92.856



LCDFGSATTISHY(ph)PDYSWSAQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33132

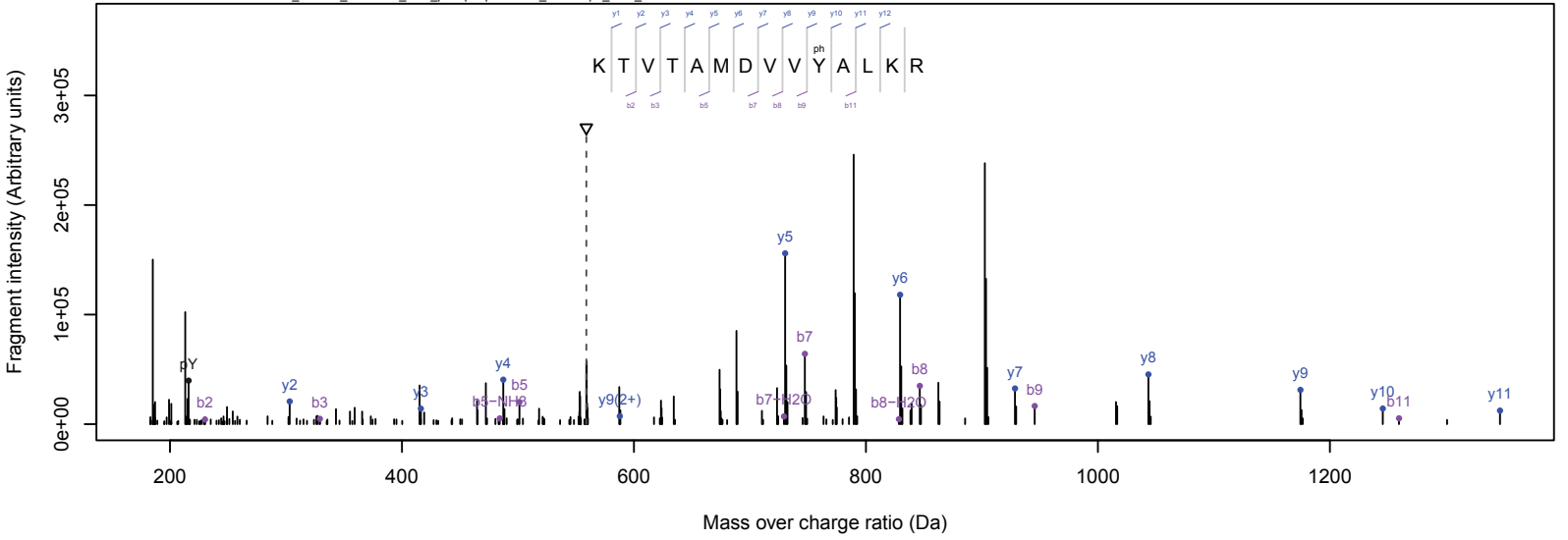
Andromeda score : 92.792



**KTVTAMDVVY(ph)ALKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35240

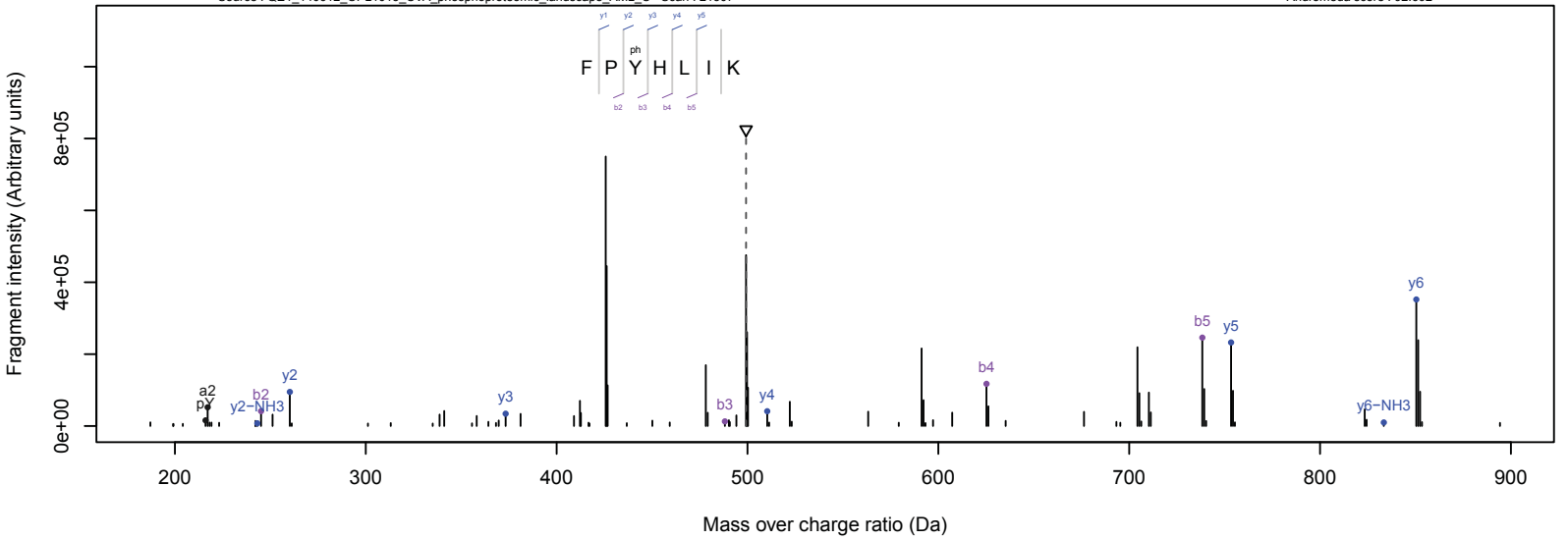
Andromeda score : 92.773



**FPY(ph)HLIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21507

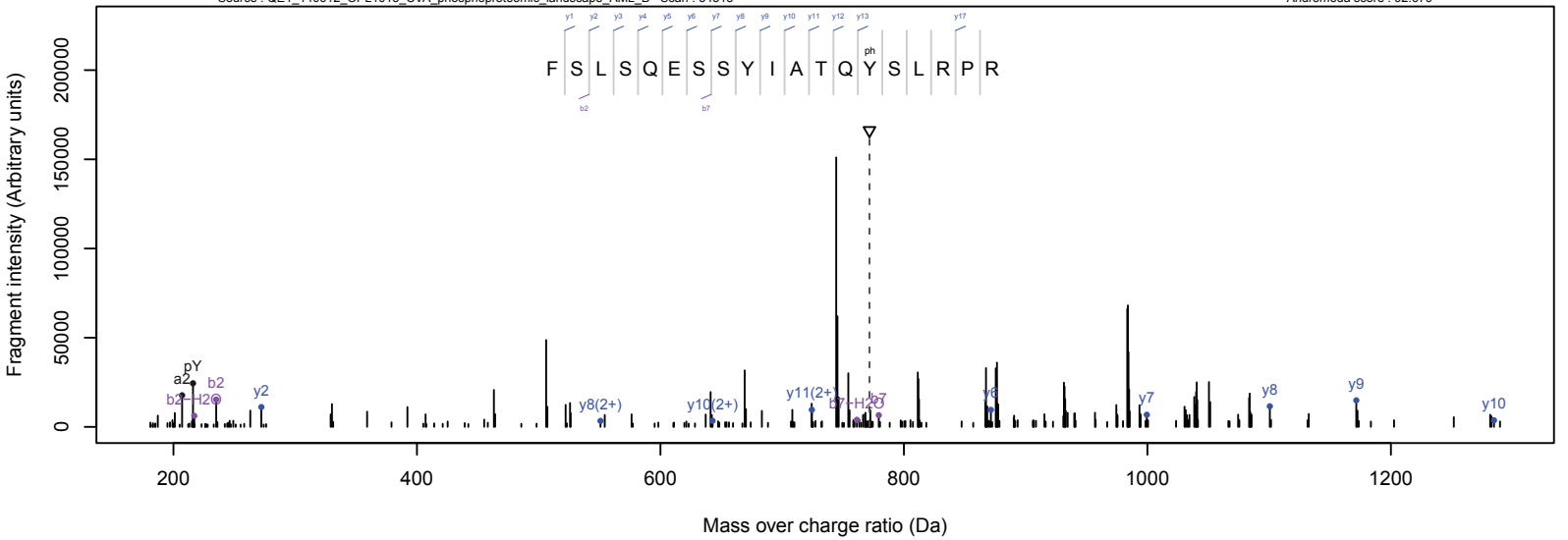
Andromeda score : 92.692



**FSLSQESSYIATQY(ph)SLRPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34518

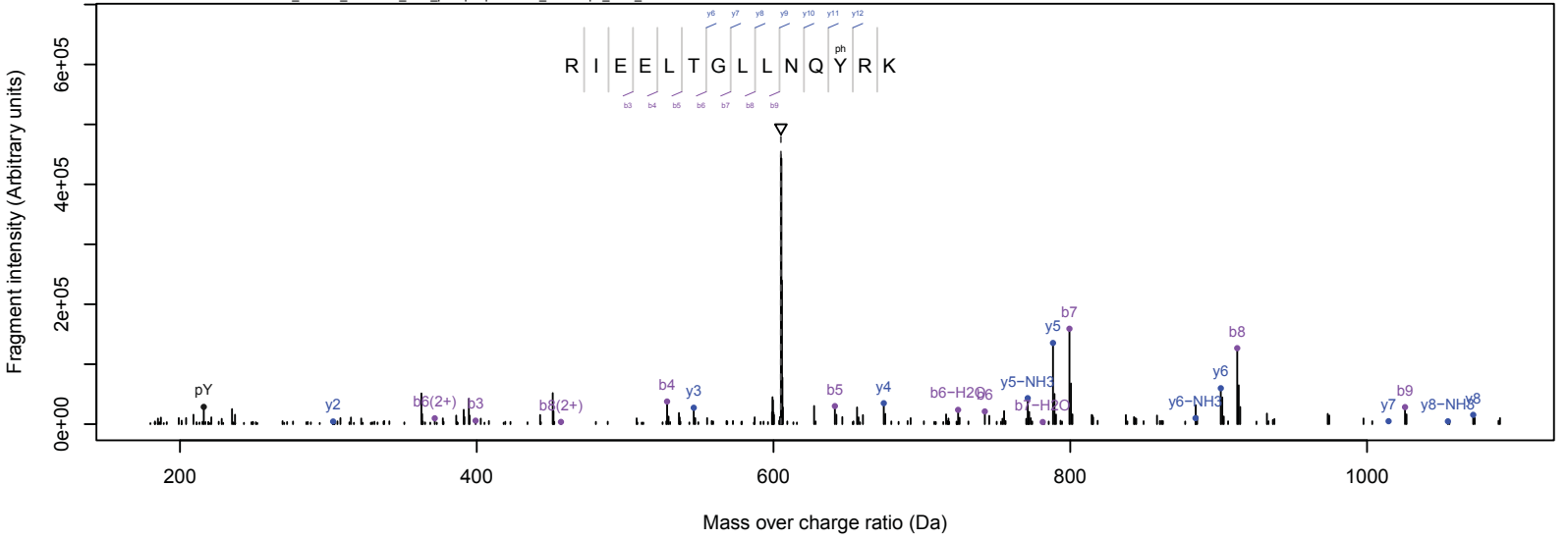
Andromeda score : 92.679



**RIEELGLLNQY(ph)RK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28471

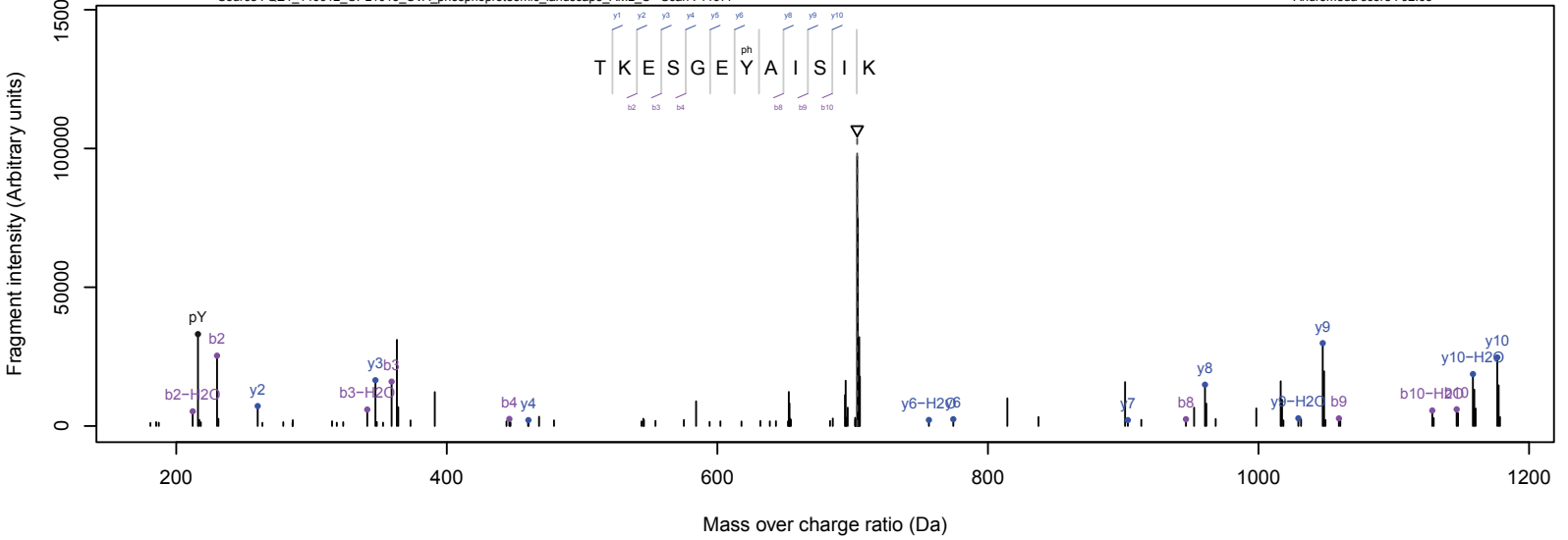
Andromeda score : 92.65



**TKESGEY(ph)AISIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11977

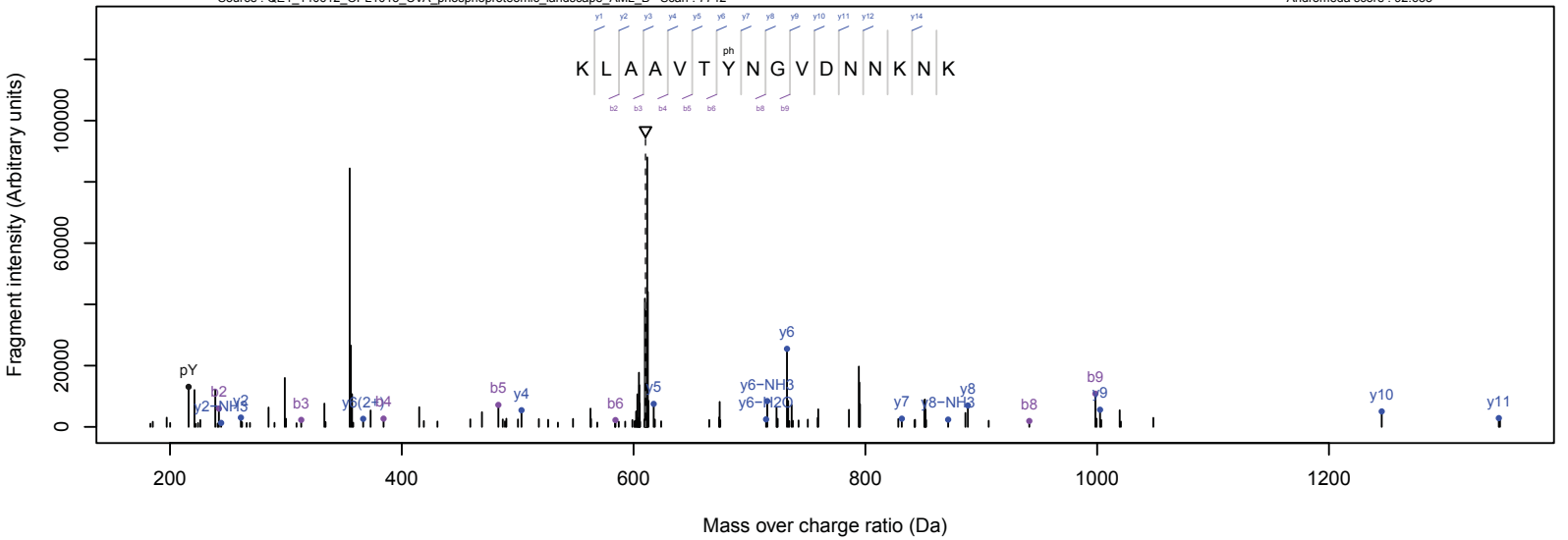
Andromeda score : 92.65



**KLAAVTY(ph)NGVDNKNK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7742

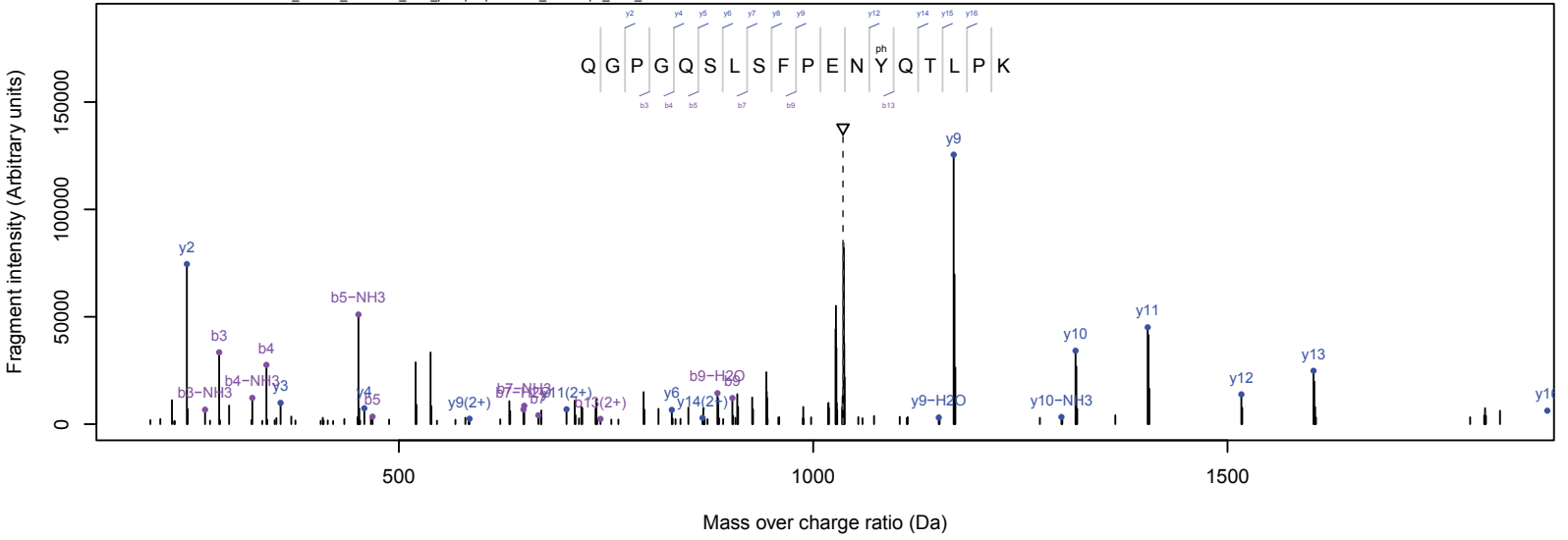
Andromeda score : 92.633



**QGGQQLSFPENY(ph)QTLPK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 30912

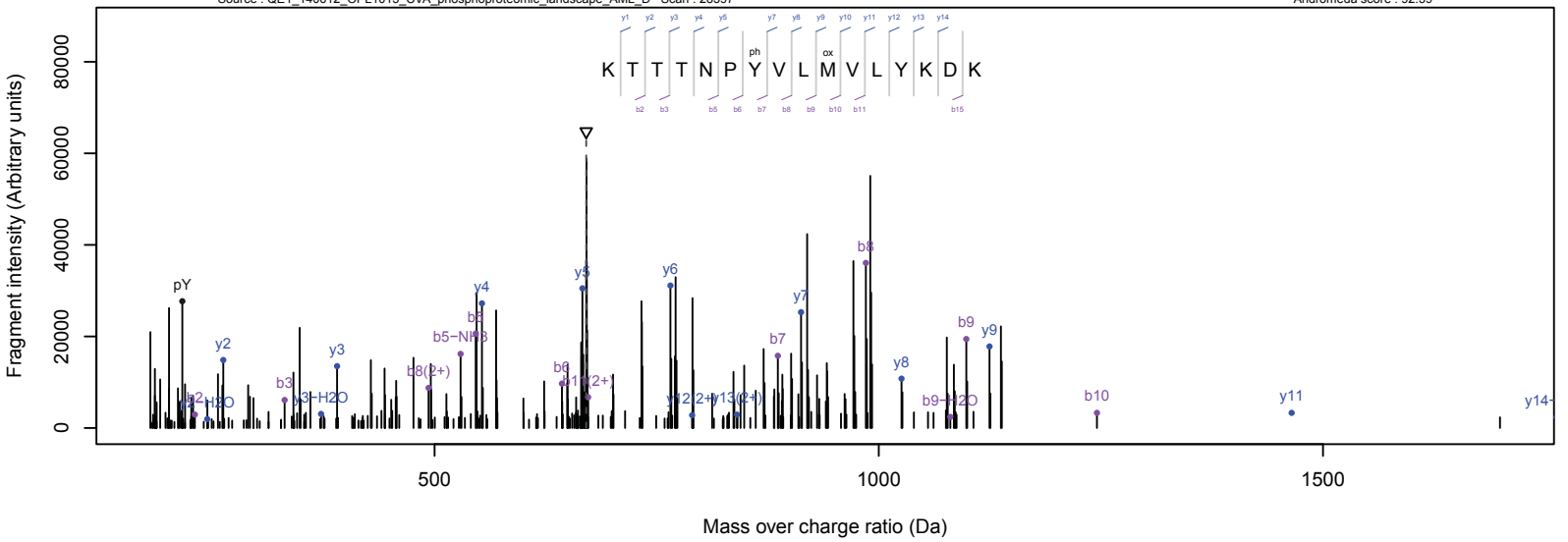
Andromeda score: 92.633



**KTTTNPY(ph)VLM(ox)VLYKDK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 28397

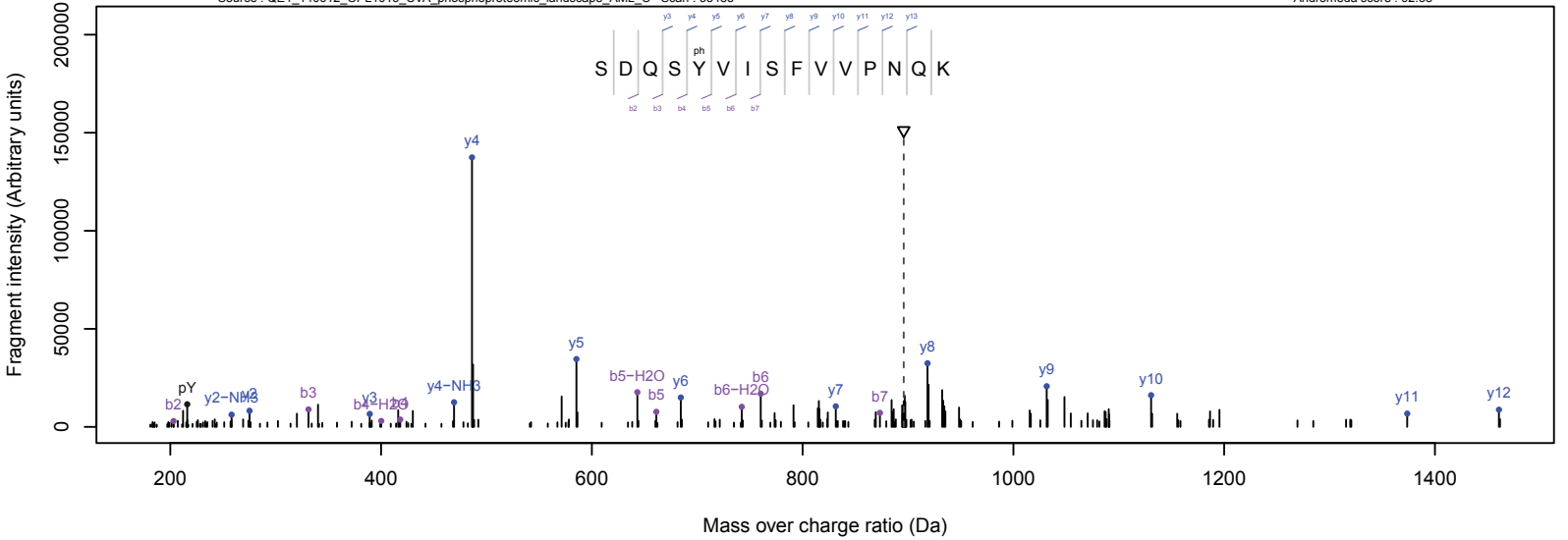
Andromeda score: 92.39



**SDQSY(ph)VISFVVPNQK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38483

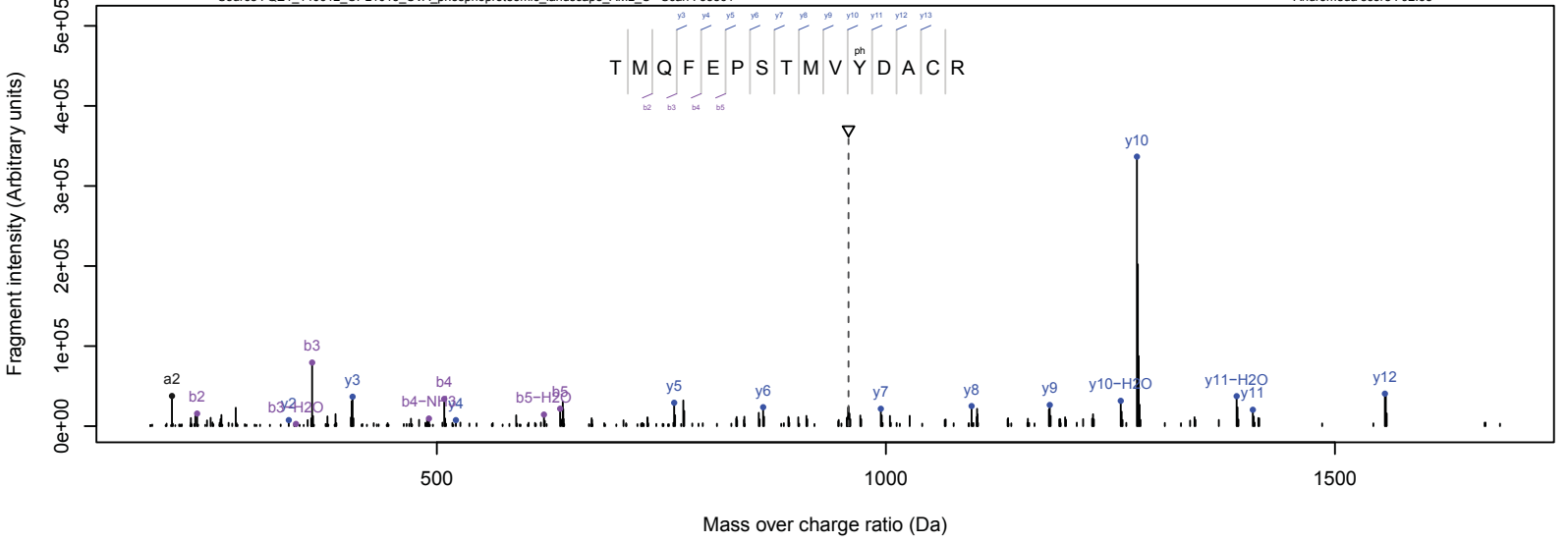
Andromeda score: 92.38



**TMQFEPSTMVY(ph)DACR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33861

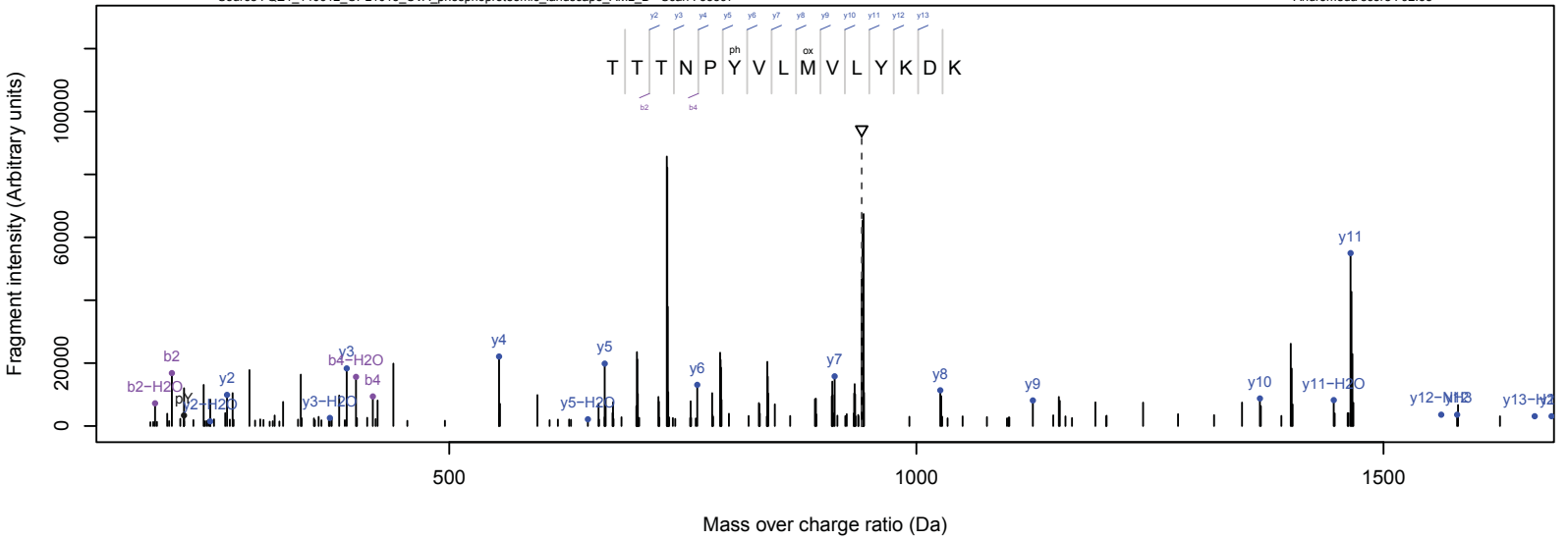
Andromeda score : 92.38



**TTTNPY(ph)VLM(ox)VLYKDK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33597

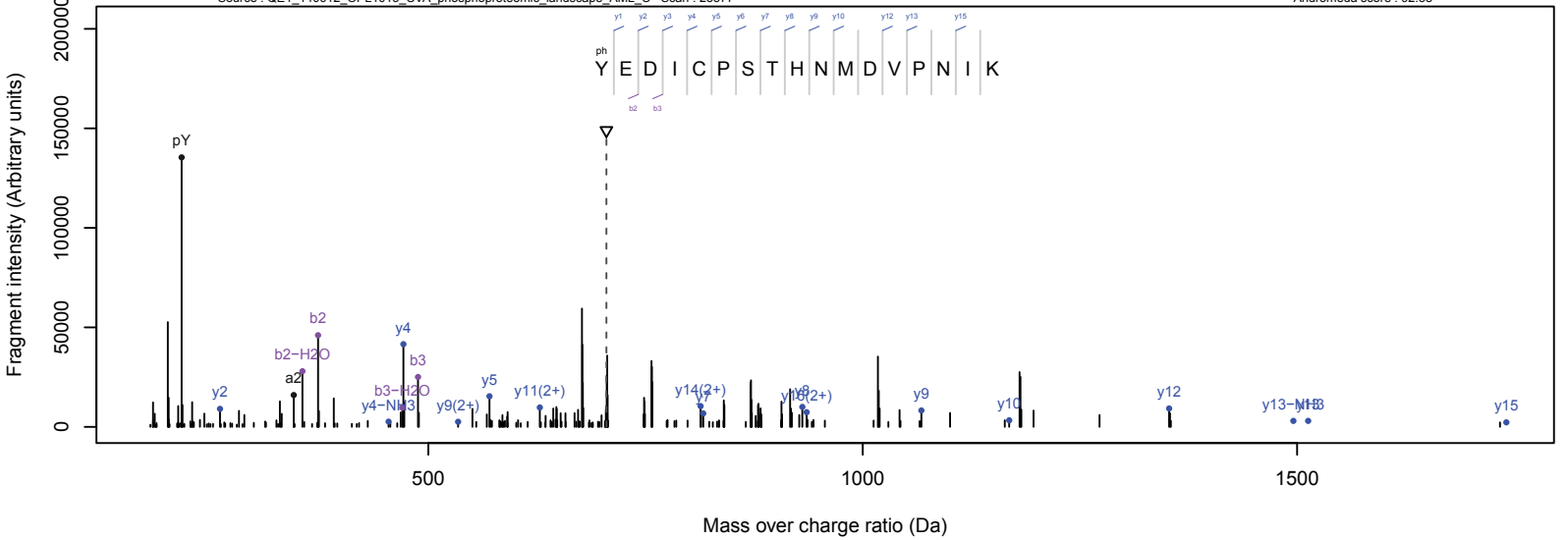
Andromeda score : 92.38



**Y(ph)EDICPSTHNM DVPNIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25577

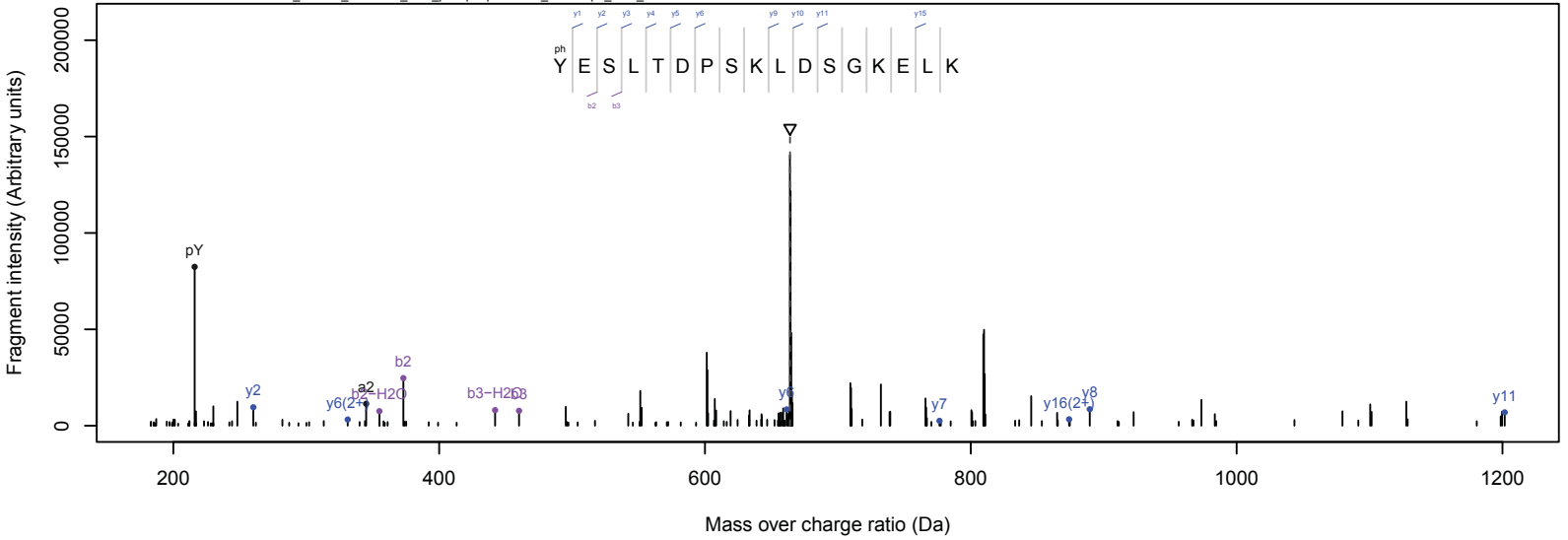
Andromeda score : 92.38



Y(ph)ESLTDP SKLD S G K E L K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 19537

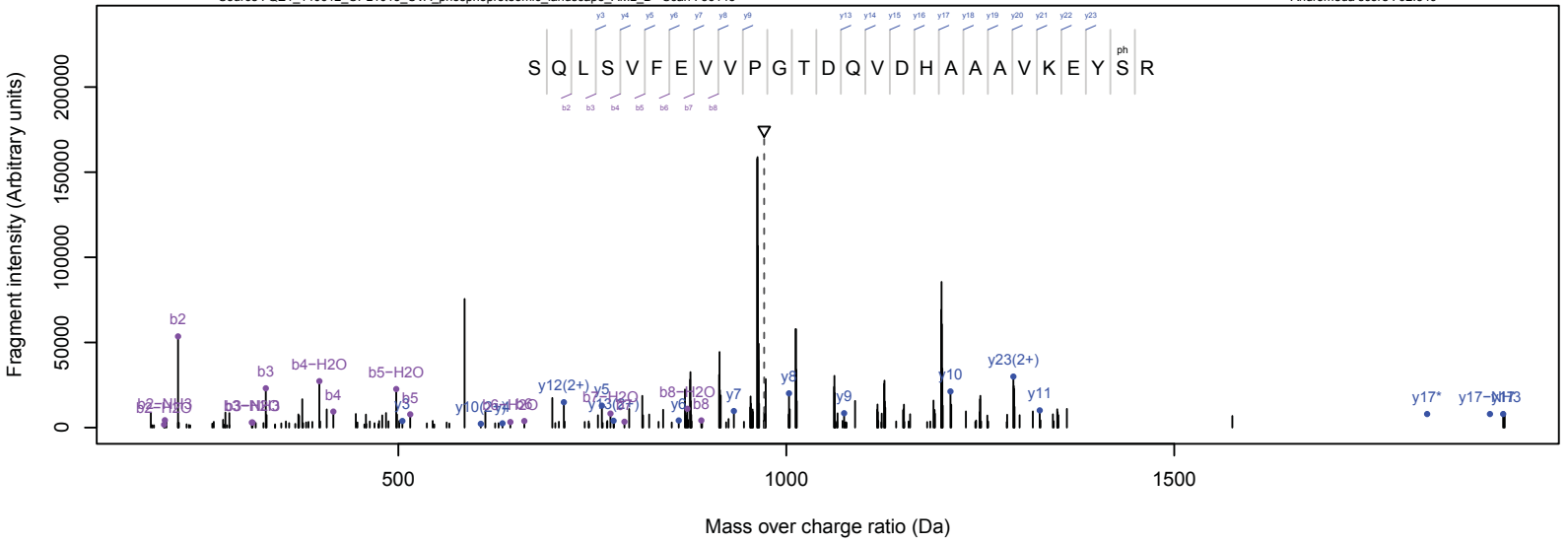
Andromeda score : 92.38



SQLSVFEVVP GT D Q V D H A A V K E Y S R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36145

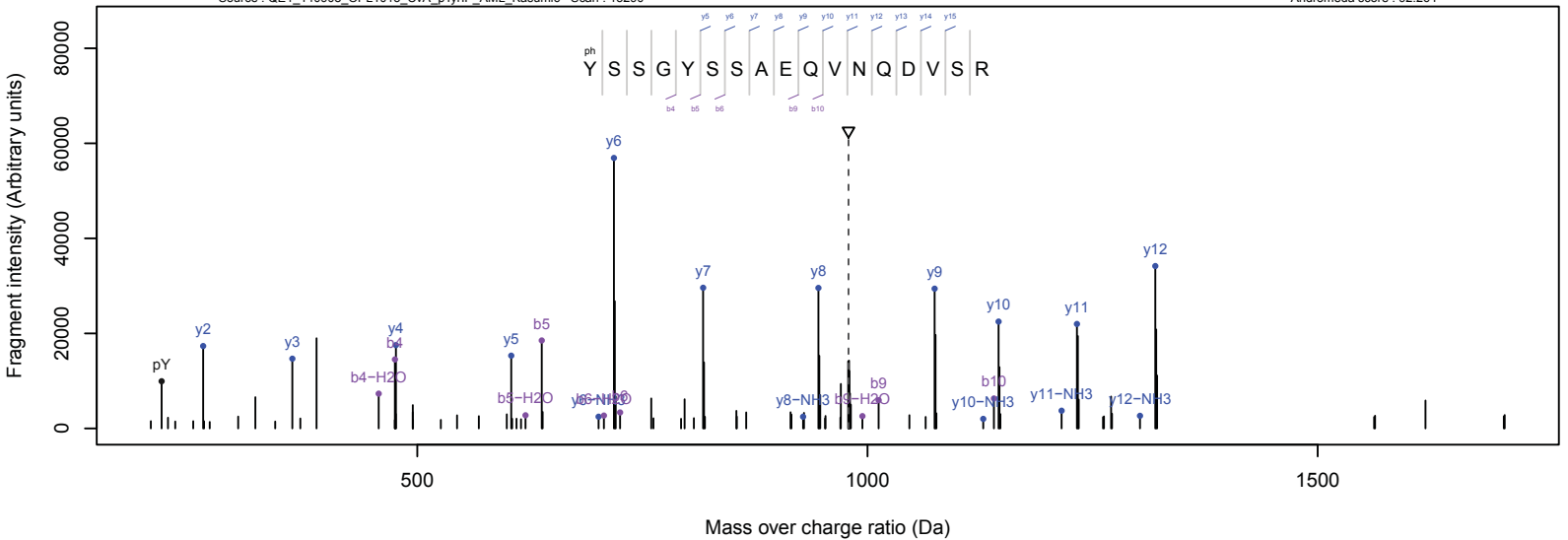
Andromeda score : 92.349



Y(ph)SSGYSSAEQ V N Q D V S R-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 15290

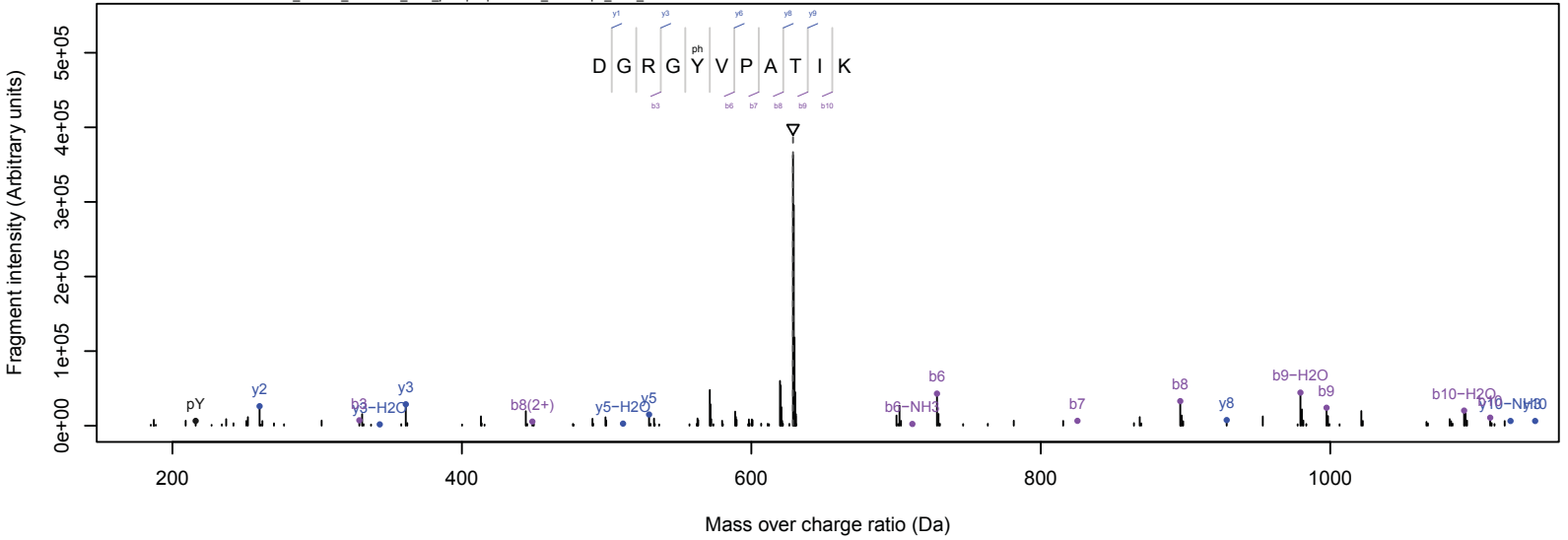
Andromeda score : 92.264



DGRGY(ph)VPATIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10923

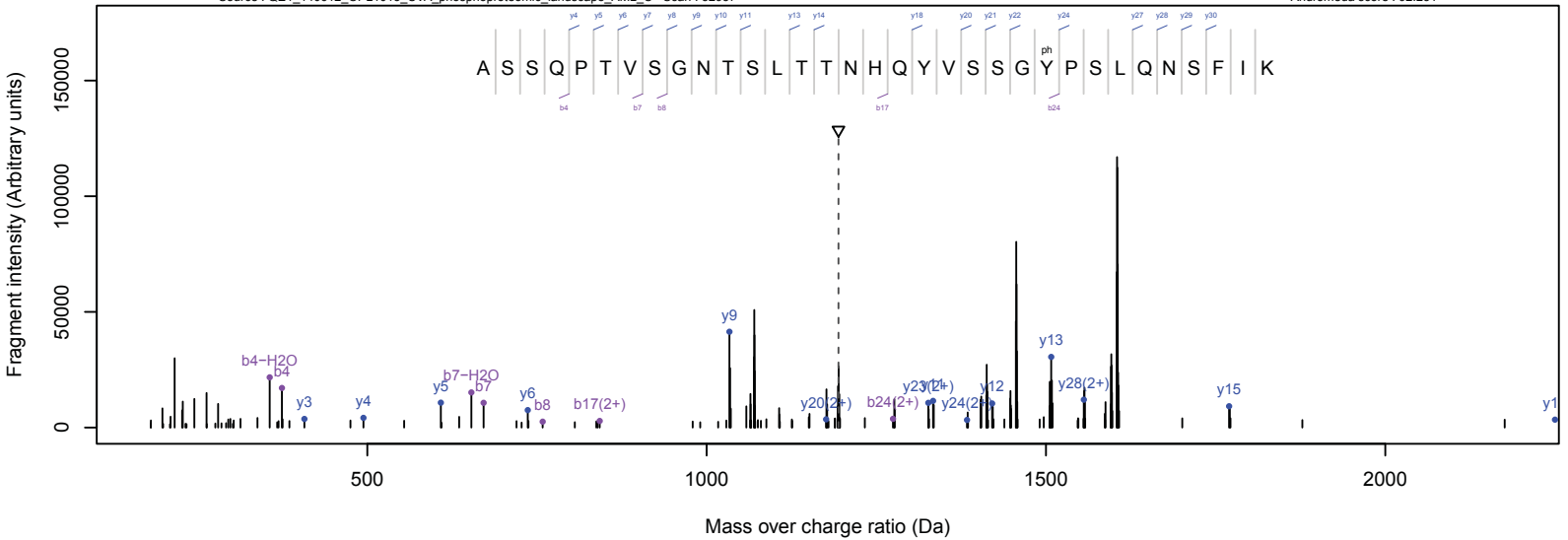
Andromeda score : 92.247



ASSQPTVSGNTSLTTHQYVSSG(ph)PSLQNSFIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32957

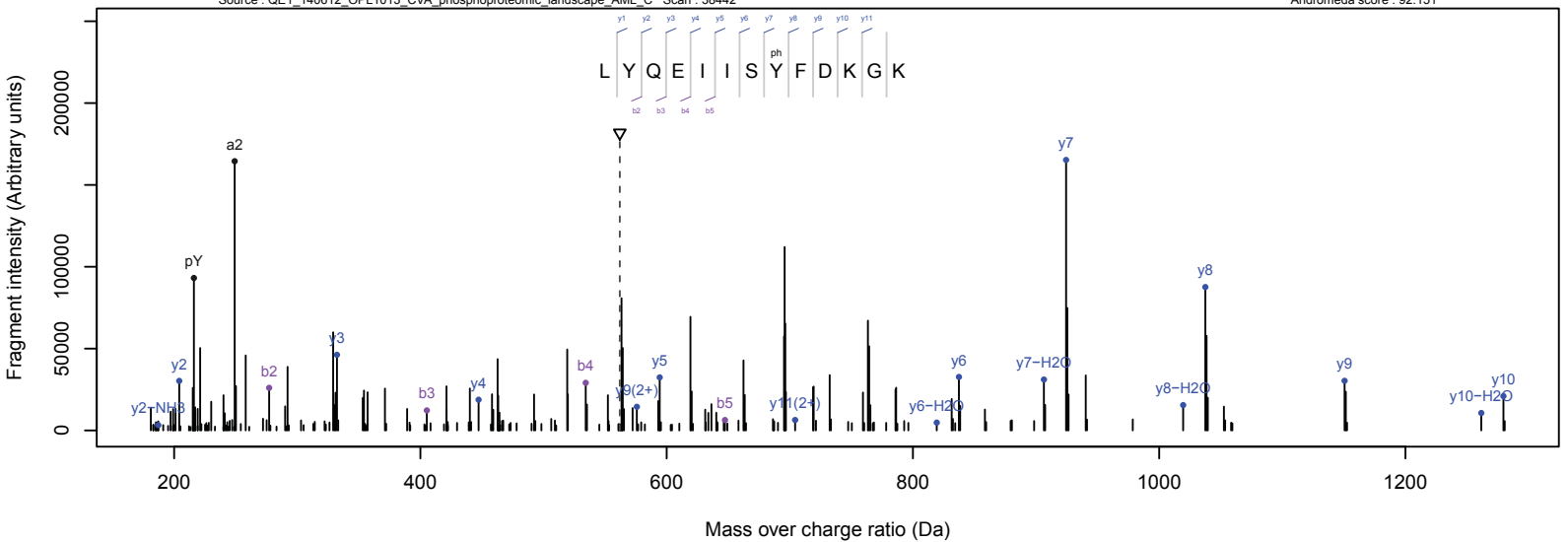
Andromeda score : 92.231



LYQEIIISYF(DK)GK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38442

Andromeda score : 92.151

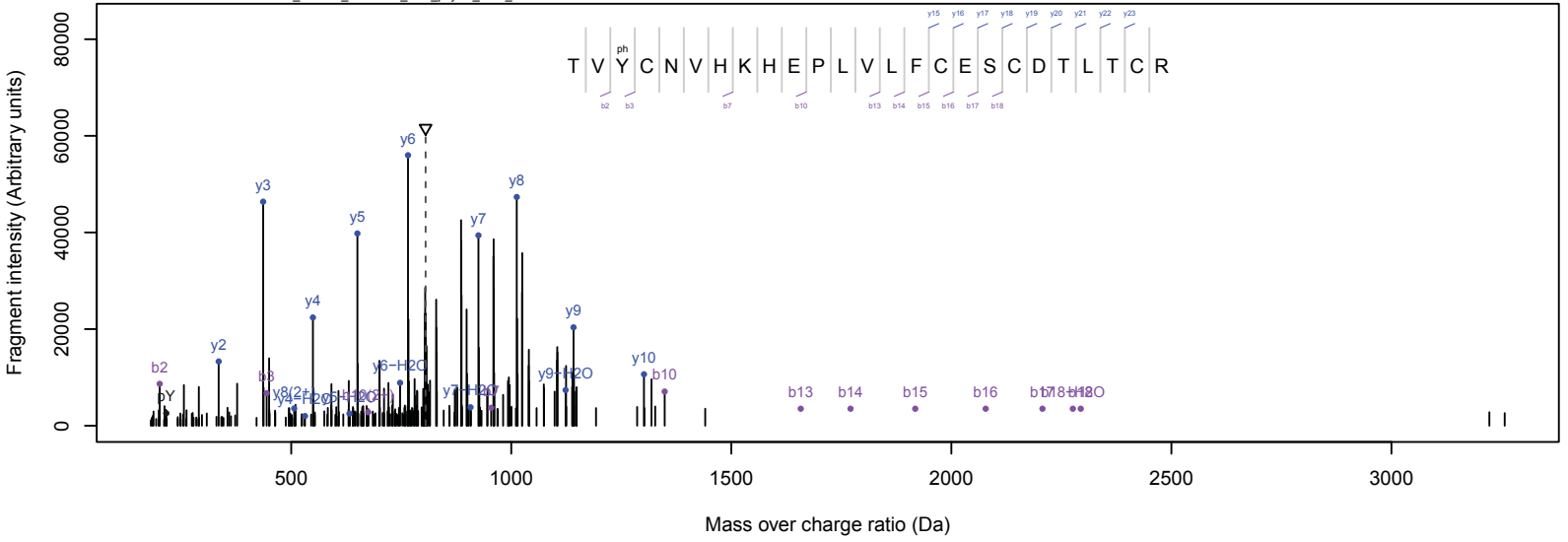




TVY(ph)CNVHKHEPLVLFCE~~S~~CDTLTCR-4+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 33013

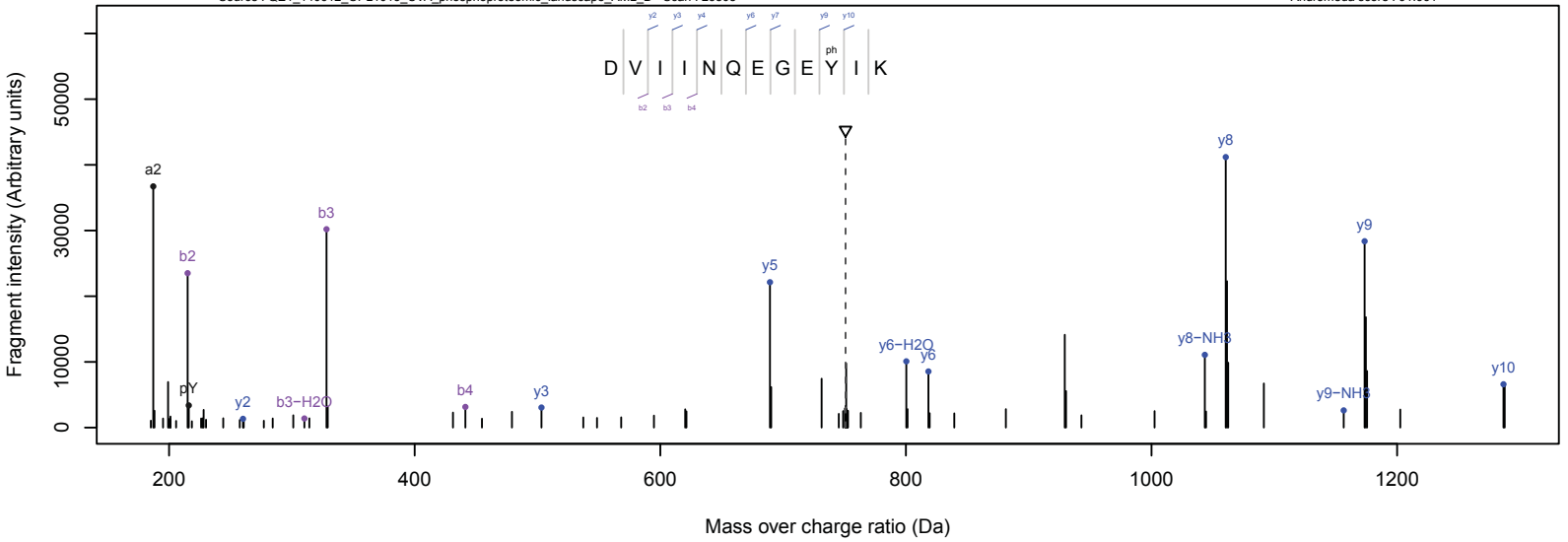
Andromeda score : 92.147



DVIINQEGEY(ph)IK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23808

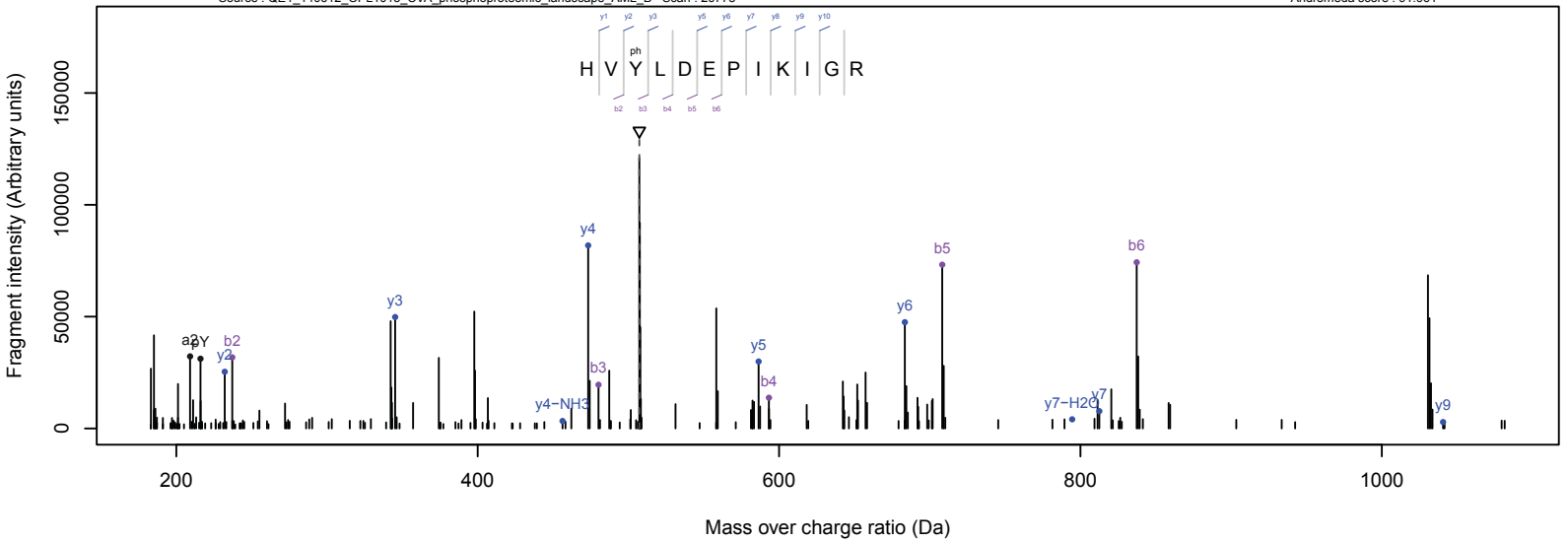
Andromeda score : 91.961



HVY(ph)LDEPIKIGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26778

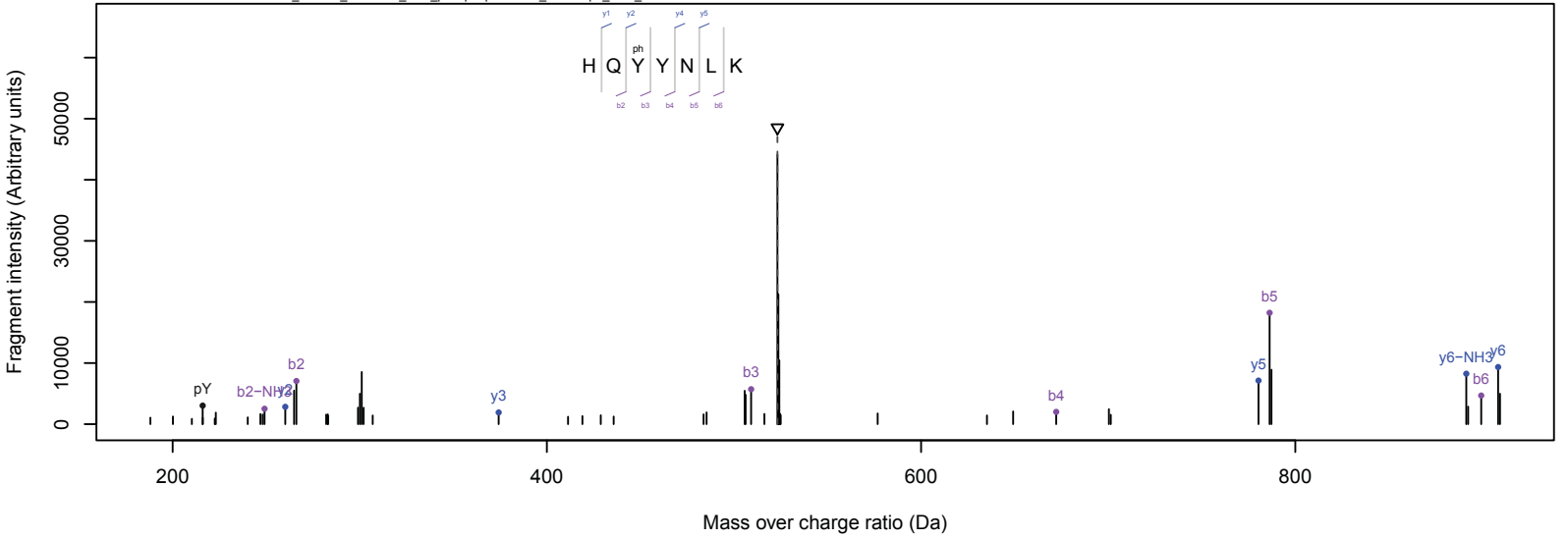
Andromeda score : 91.961



HQY(ph)YNLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 7603

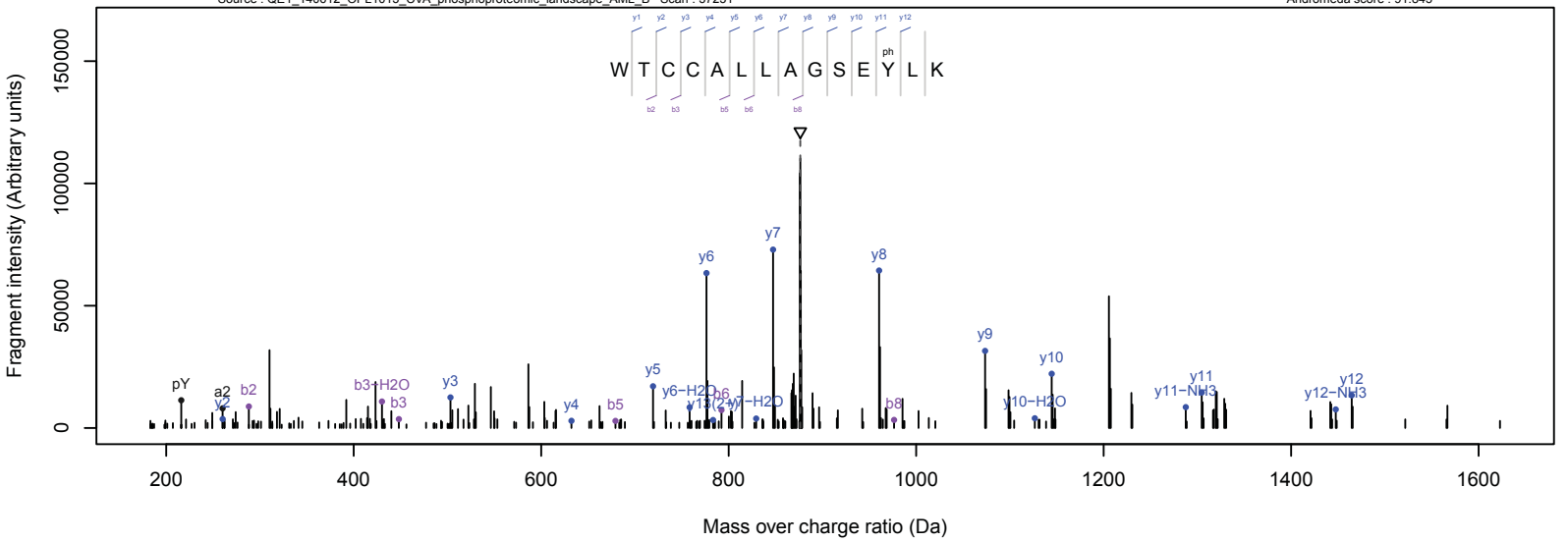
Andromeda score : 91.906



WTCCALLAGSEY(ph)LK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37231

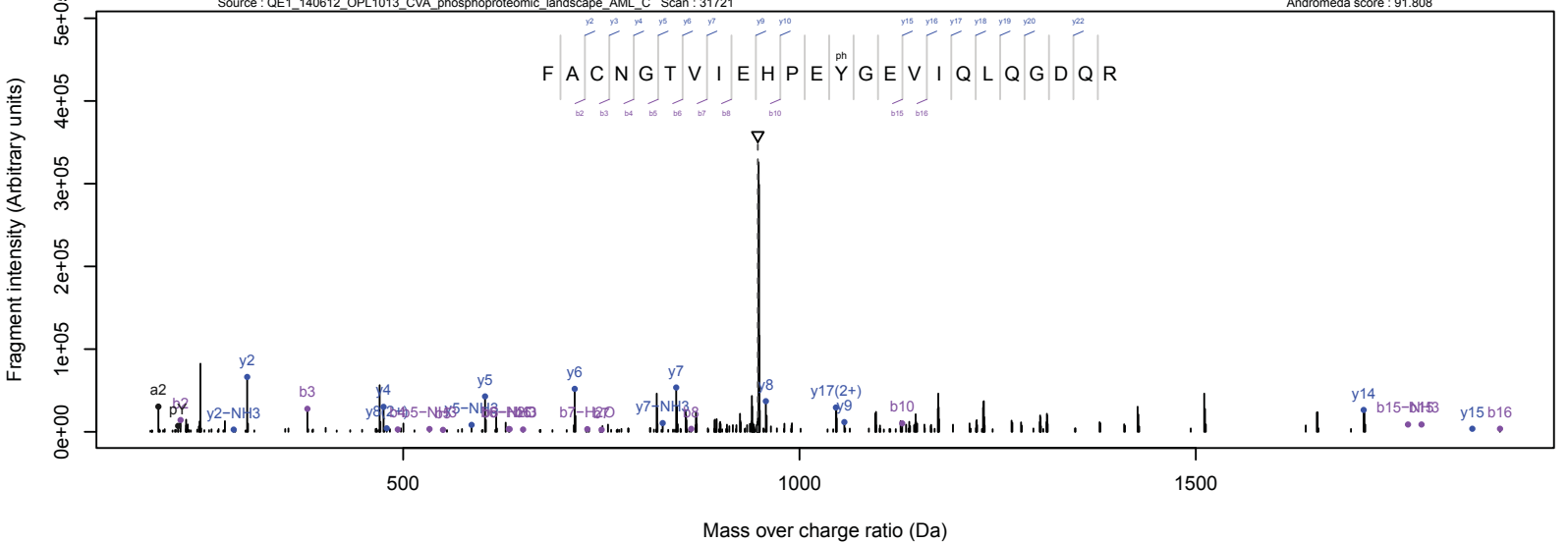
Andromeda score : 91.845



FACNGTVIEHPEY(ph)GEVIQLQGDQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31721

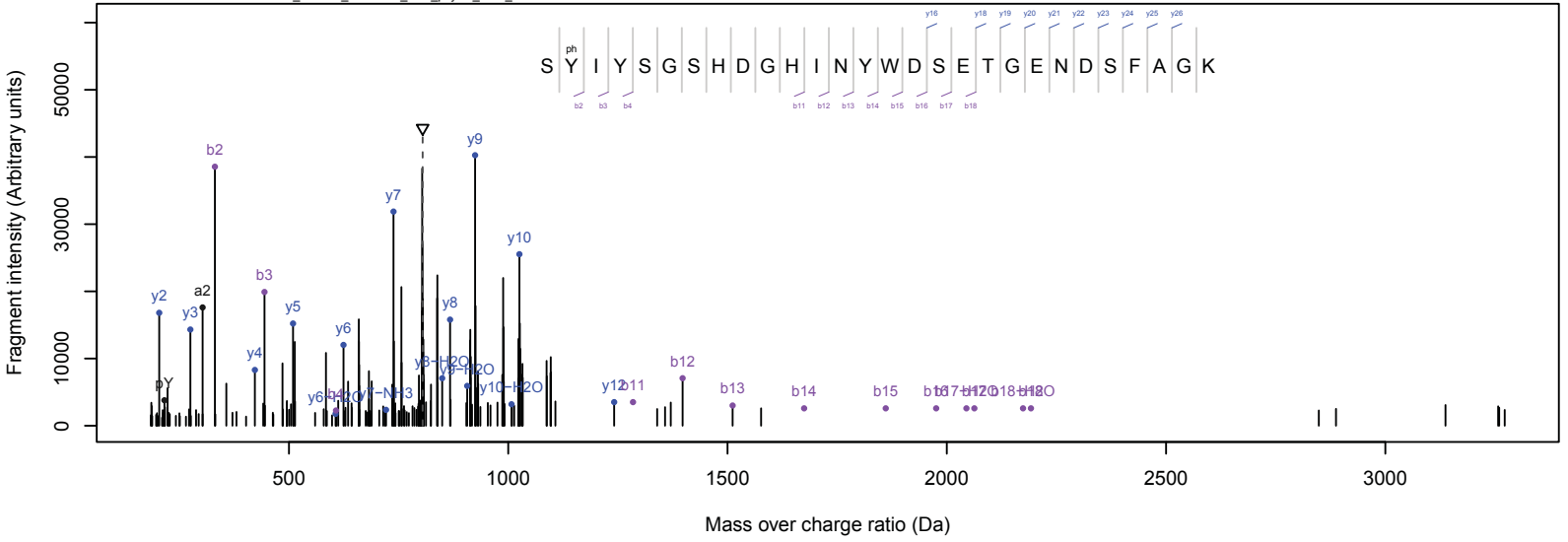
Andromeda score : 91.808



**SY(ph)IYSGSHDGHINIWDSSETGENDSFAGK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 30399

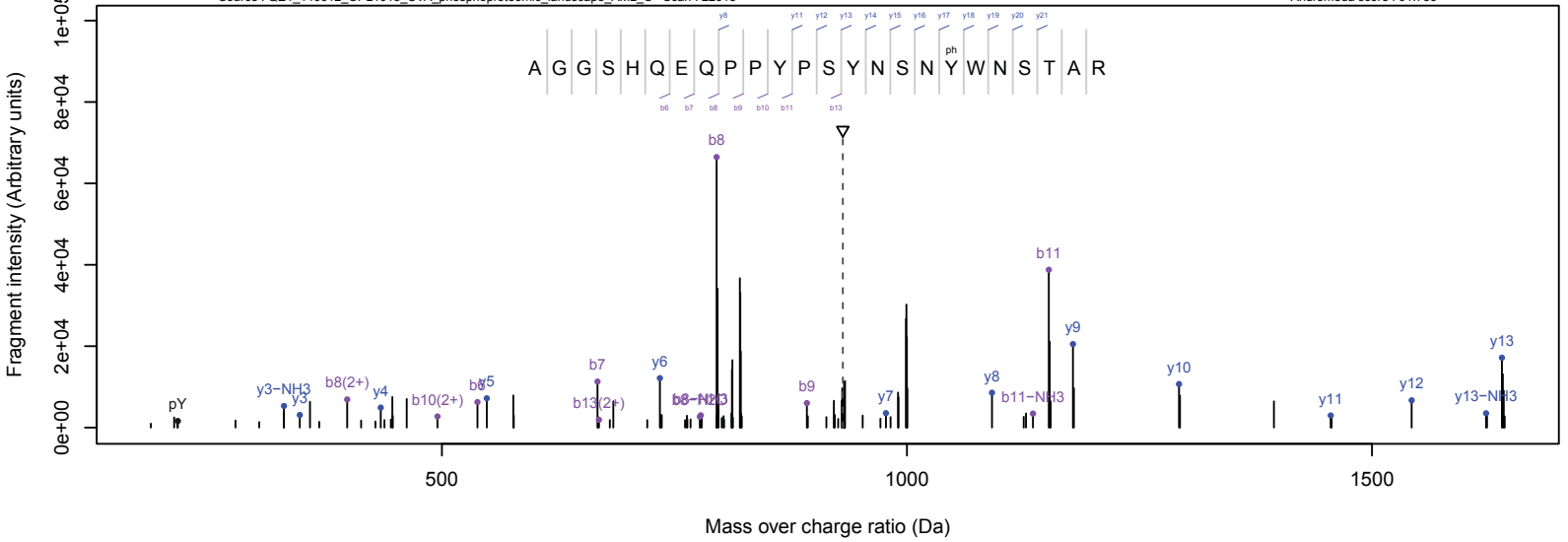
Andromeda score : 91.767



**AGGSHQEQQPPYPSYNSNY(ph)WNSTAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22618

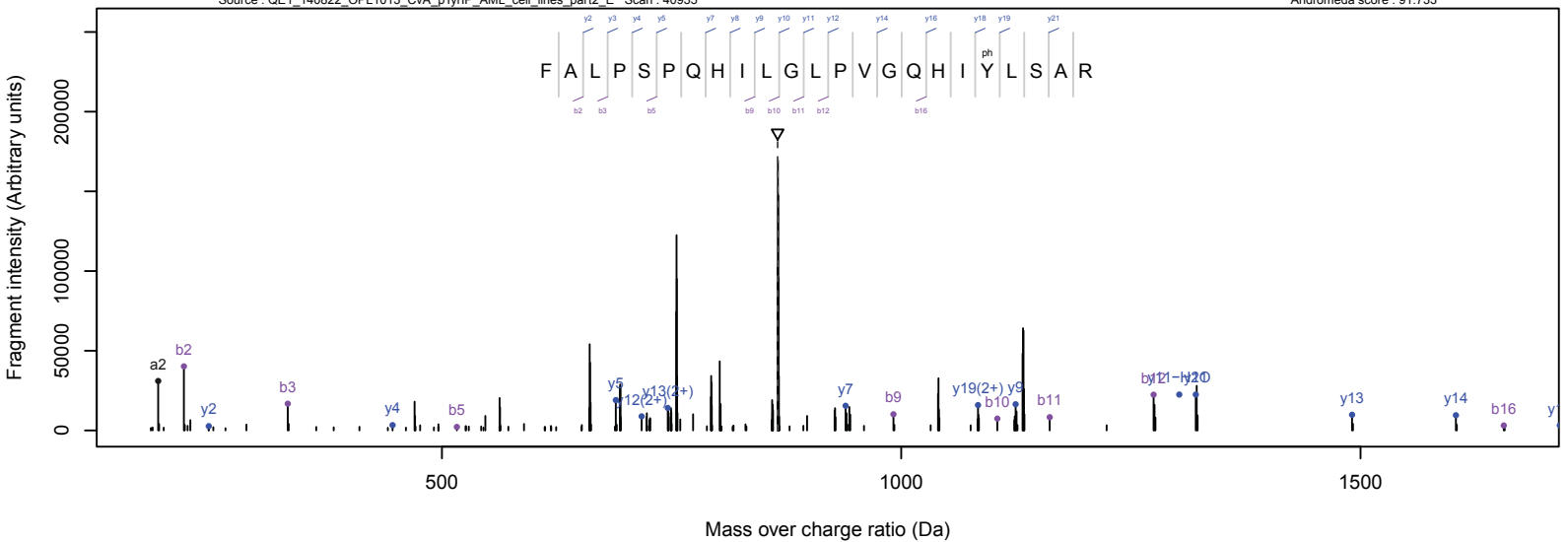
Andromeda score : 91.733



**FALPSPQHILGLPLVQGHIYLSAR-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 40935

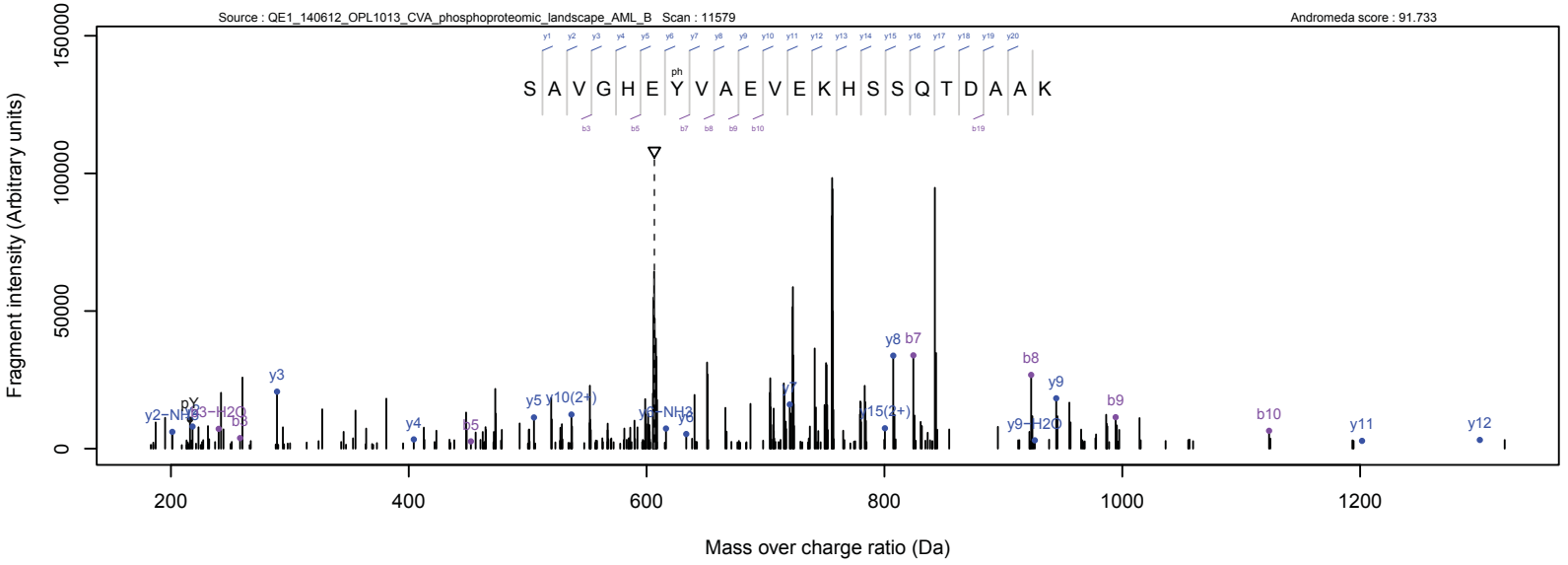
Andromeda score : 91.733



SAVGHEY(ph)VAEVEKHSSTDAAK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11579

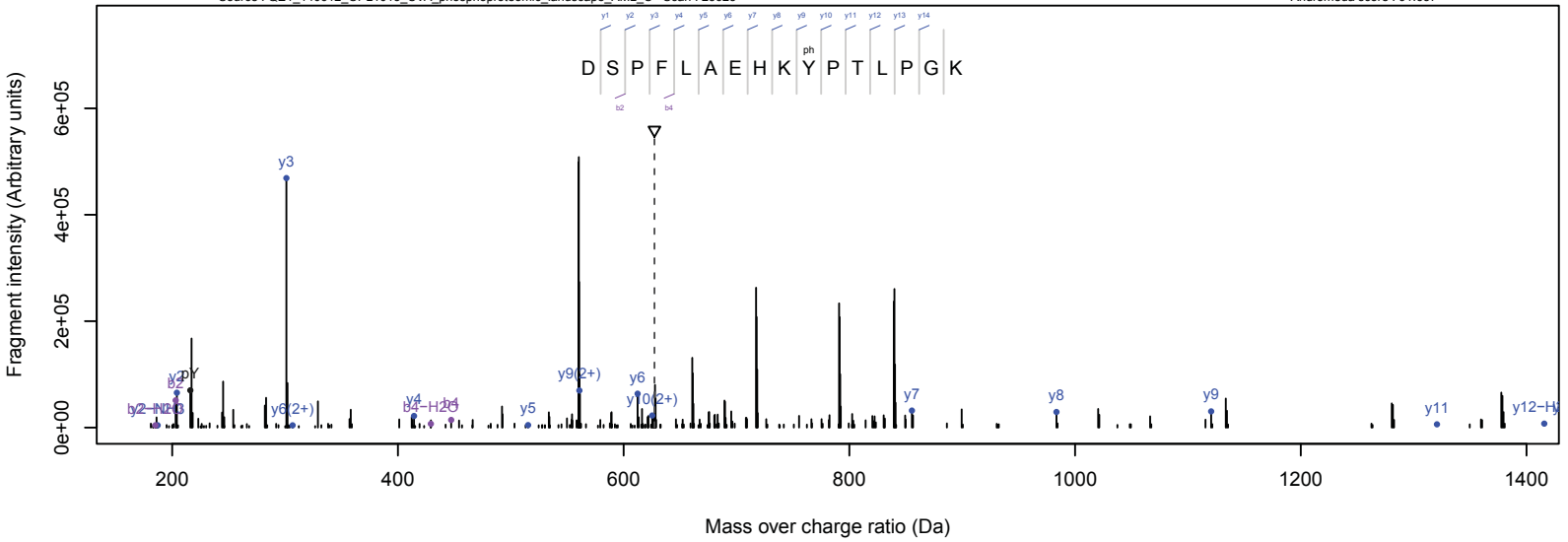
Andromeda score : 91.733



DSPFLAEHKY(ph)PTLP GK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25625

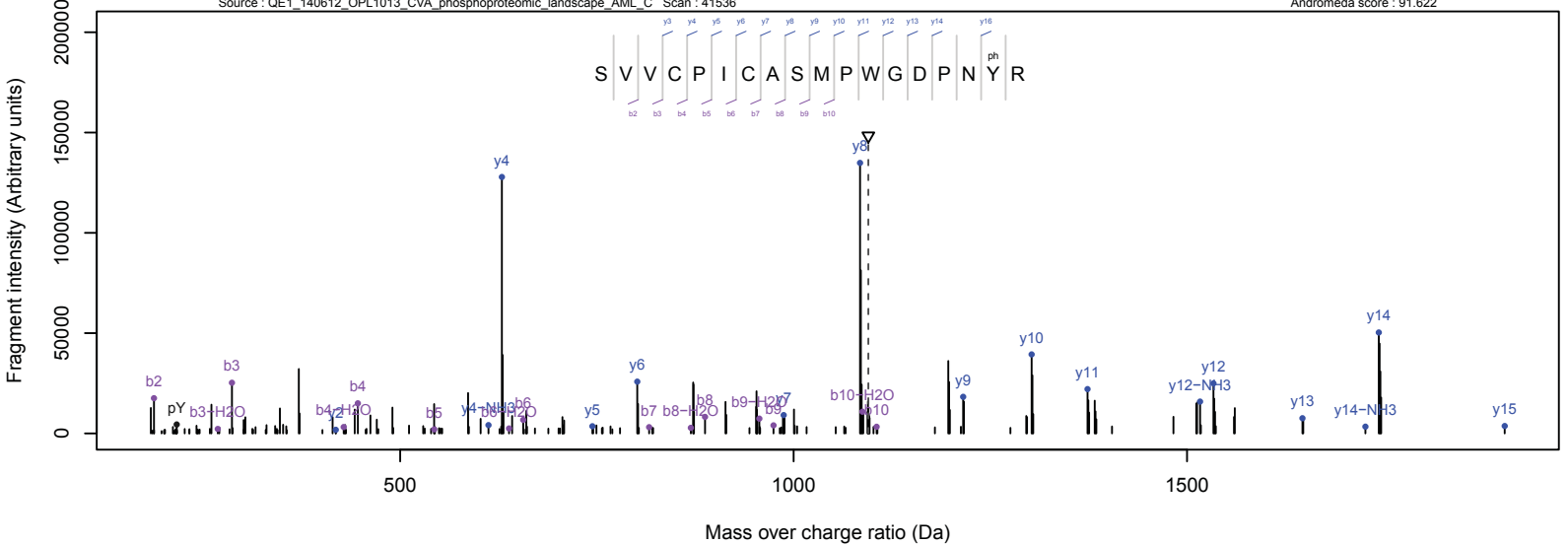
Andromeda score : 91.657



SVVCPCASMPWGP NY(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41536

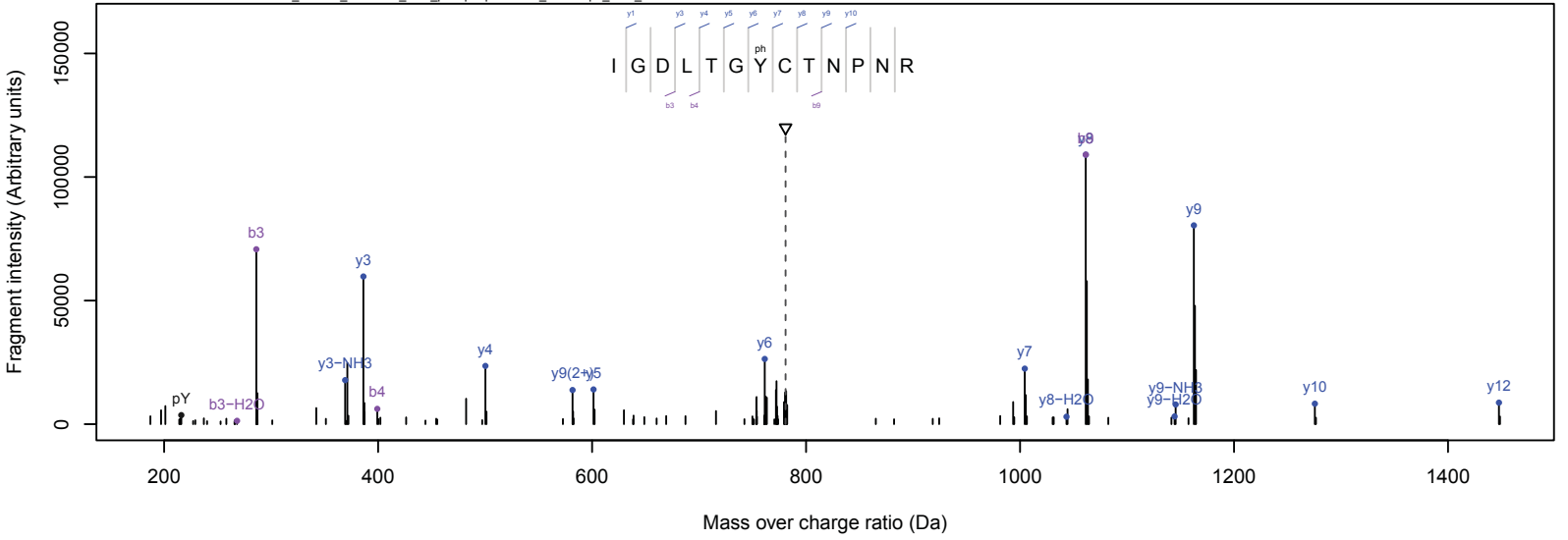
Andromeda score : 91.622



**IGDLTGY(ph)CTNPNR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14756

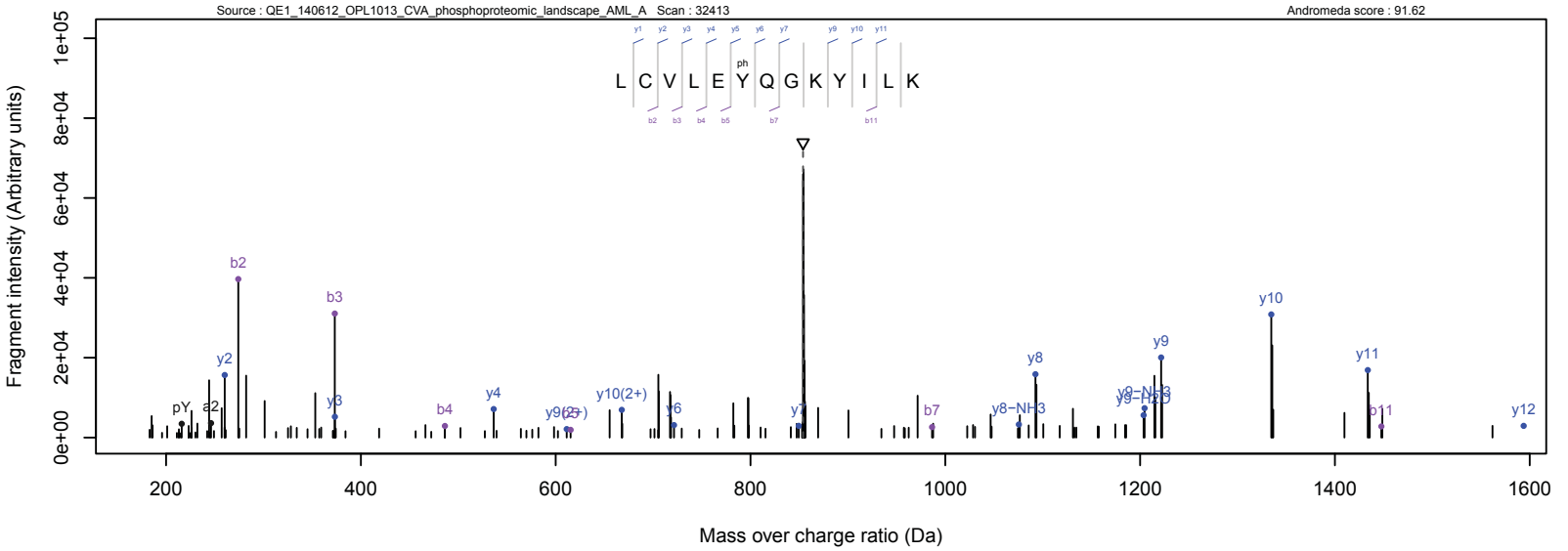
Andromeda score : 91.62



**LCVLEY(ph)QGKYILK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32413

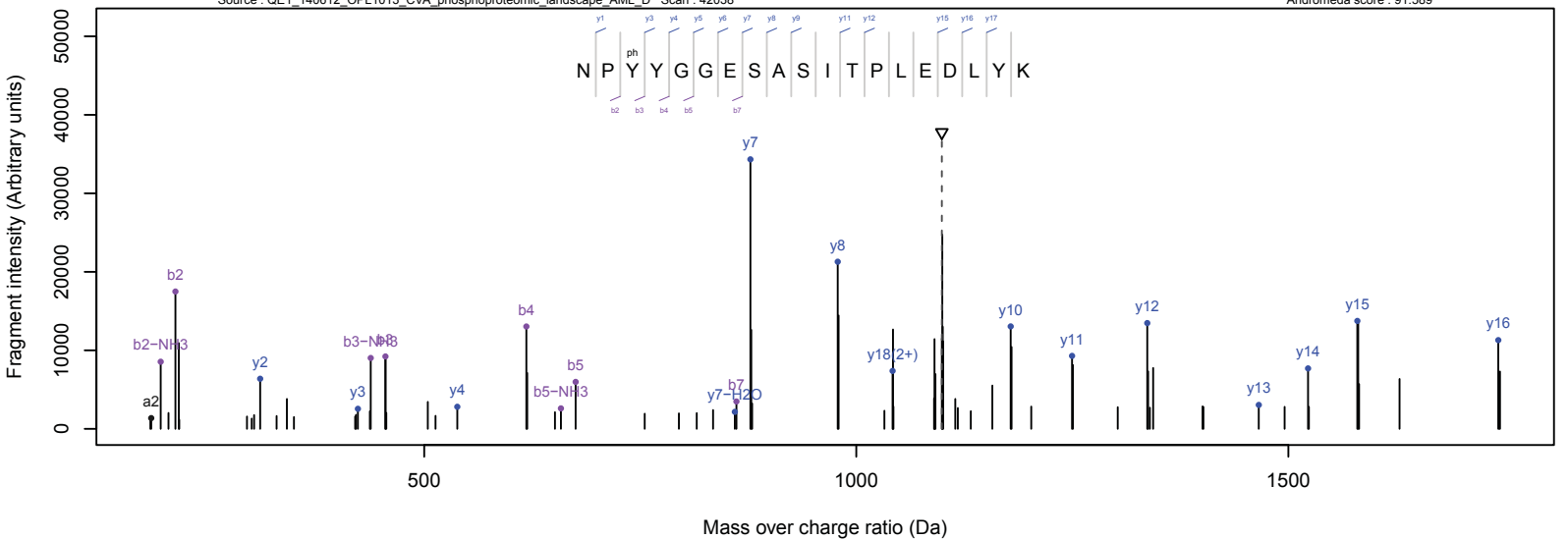
Andromeda score : 91.62



**NPY(ph)YGGESASITPLEDLYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 42038

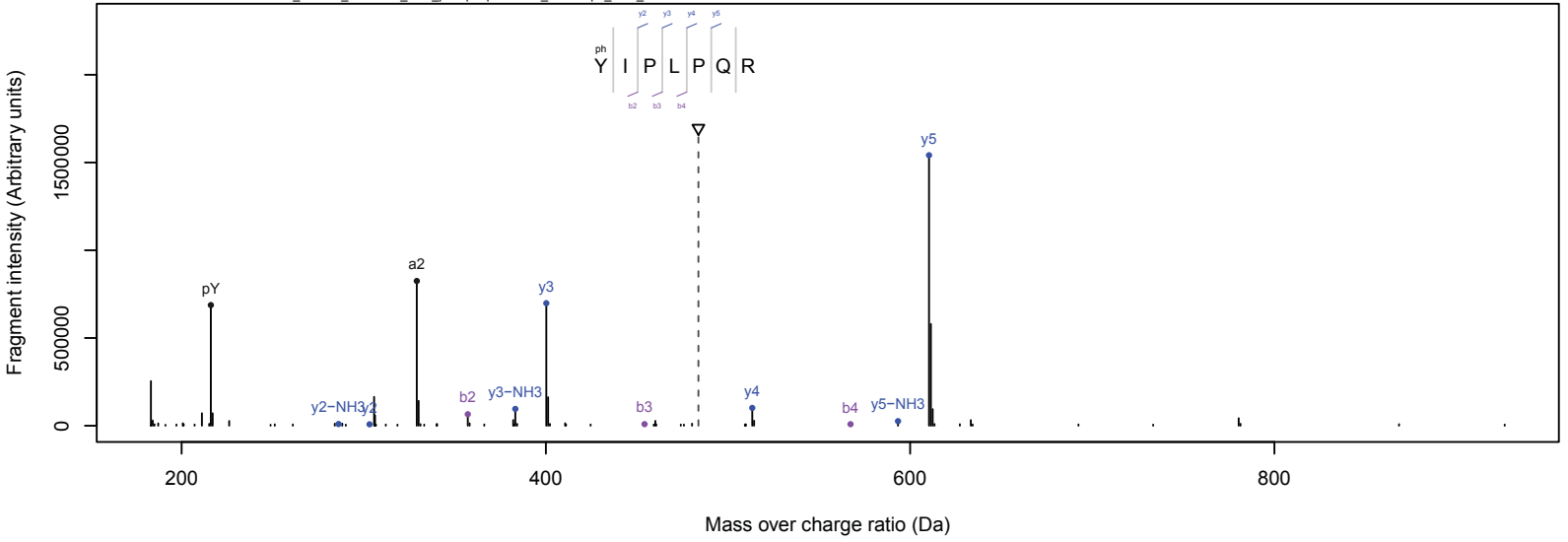
Andromeda score : 91.589



Y(ph)IPLPQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17775

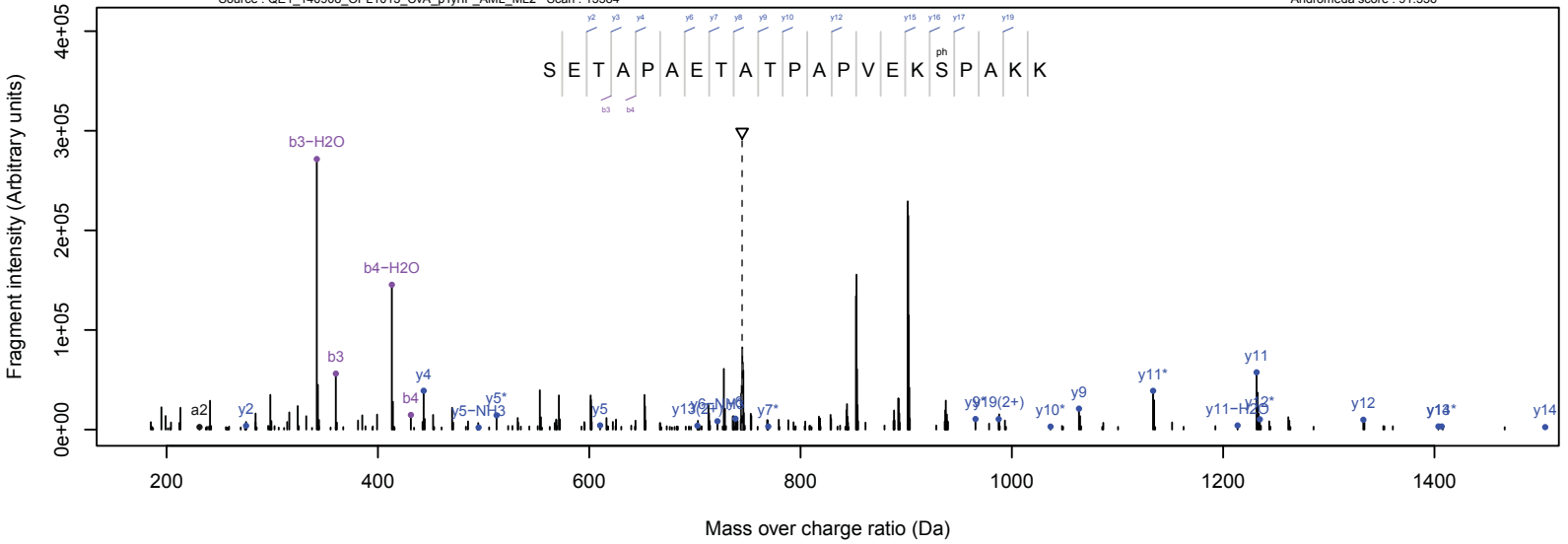
Andromeda score : 91.584



(ac)SETAPAETATPAPVEKS(ph)PAKK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 13584

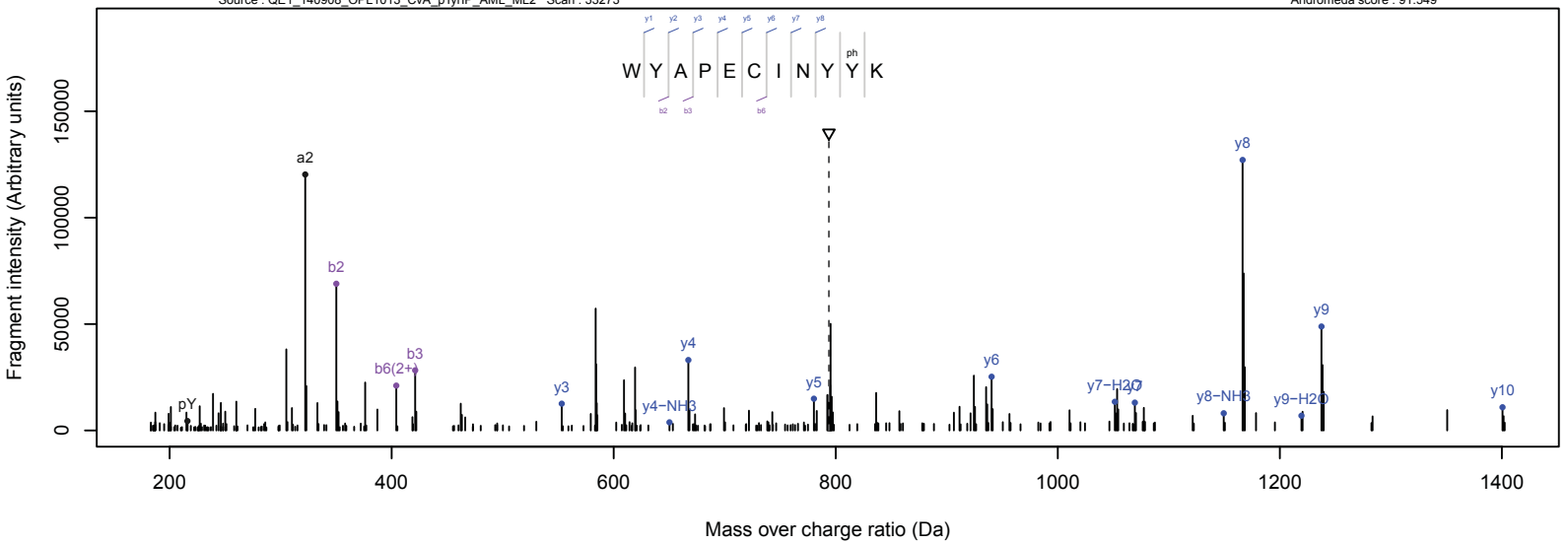
Andromeda score : 91.556



WYAPECINYY(ph)K-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 33273

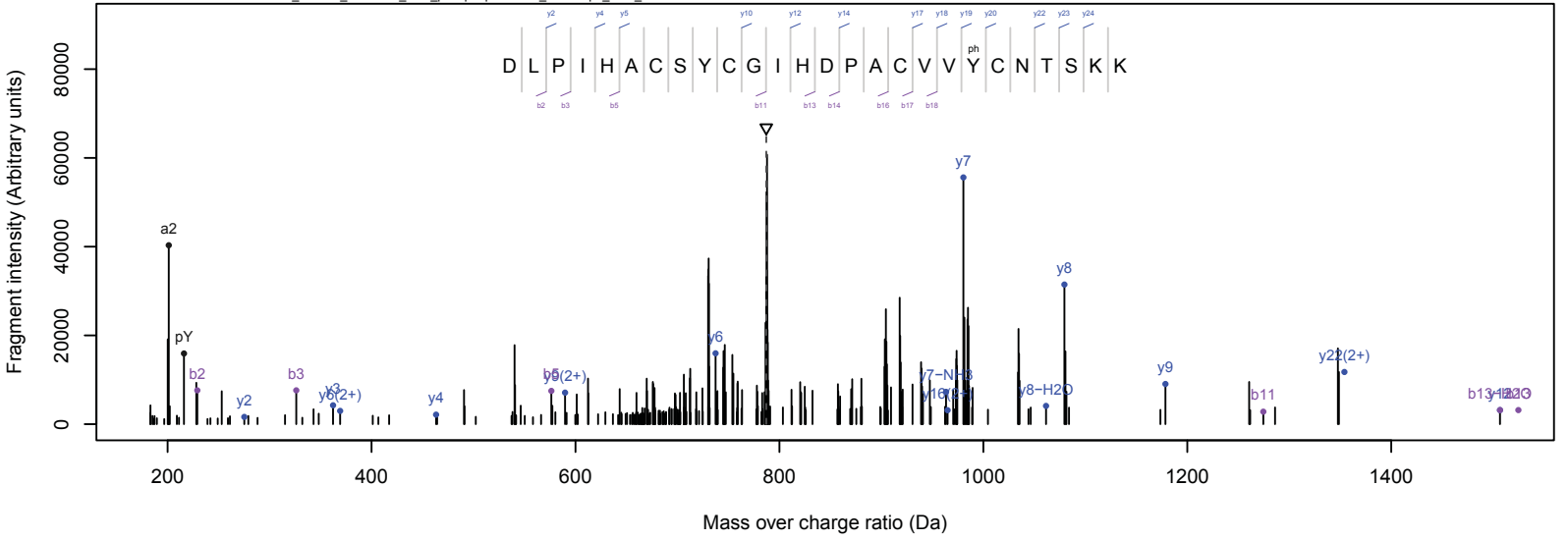
Andromeda score : 91.549



**DLPIHACSYGIIHDPACVVY(ph)CNTSKK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29036

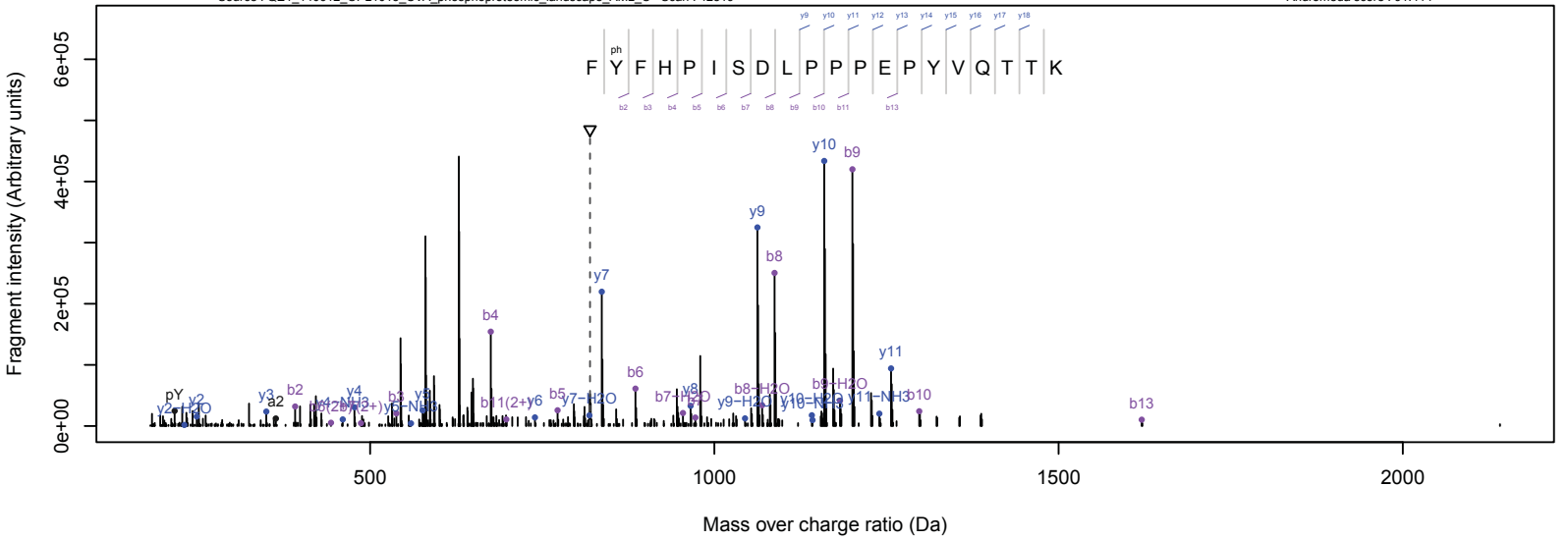
Andromeda score : 91.416



**FY(ph)FHPISDLPPPEPYVQTTK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42819

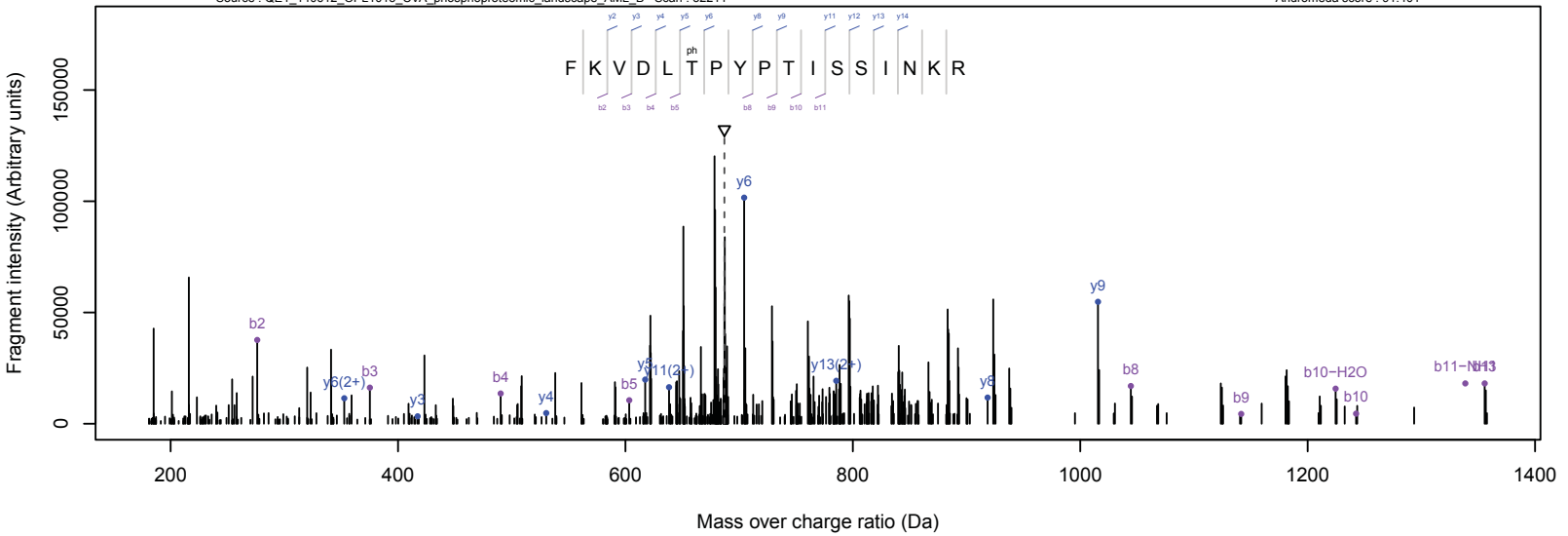
Andromeda score : 91.414



**FKVDLT(ph)PYPTISSINKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32211

Andromeda score : 91.401



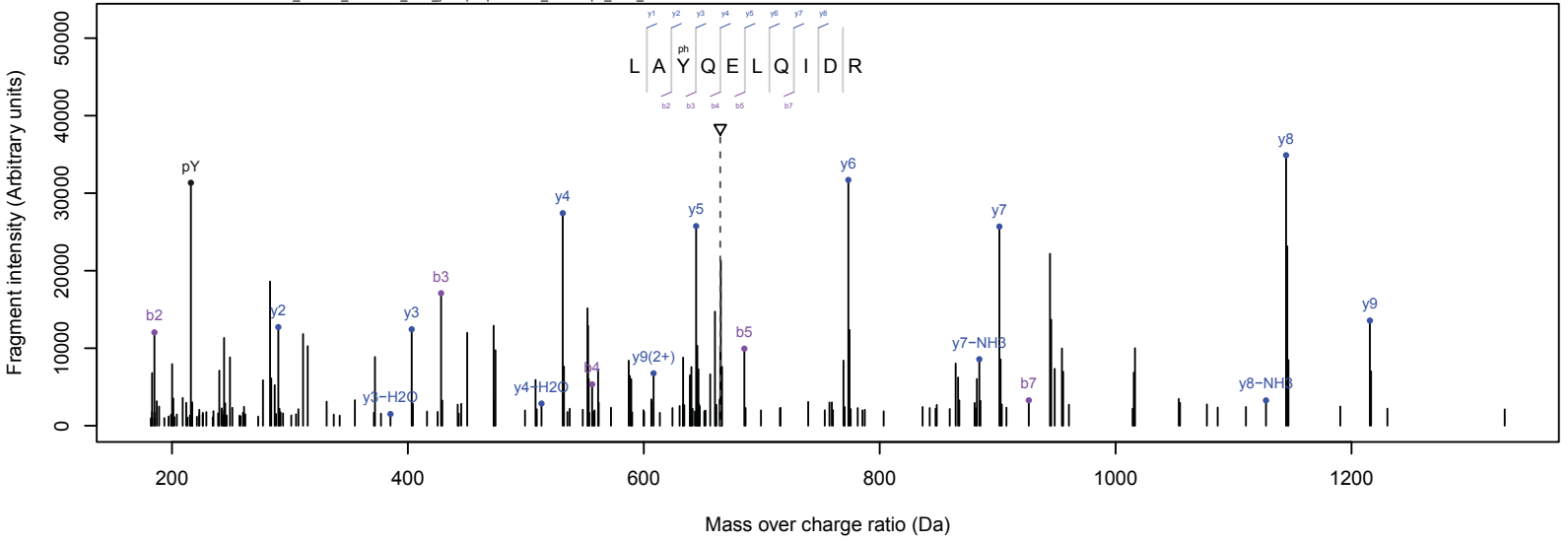




LAY(ph)QELQIDR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 21308

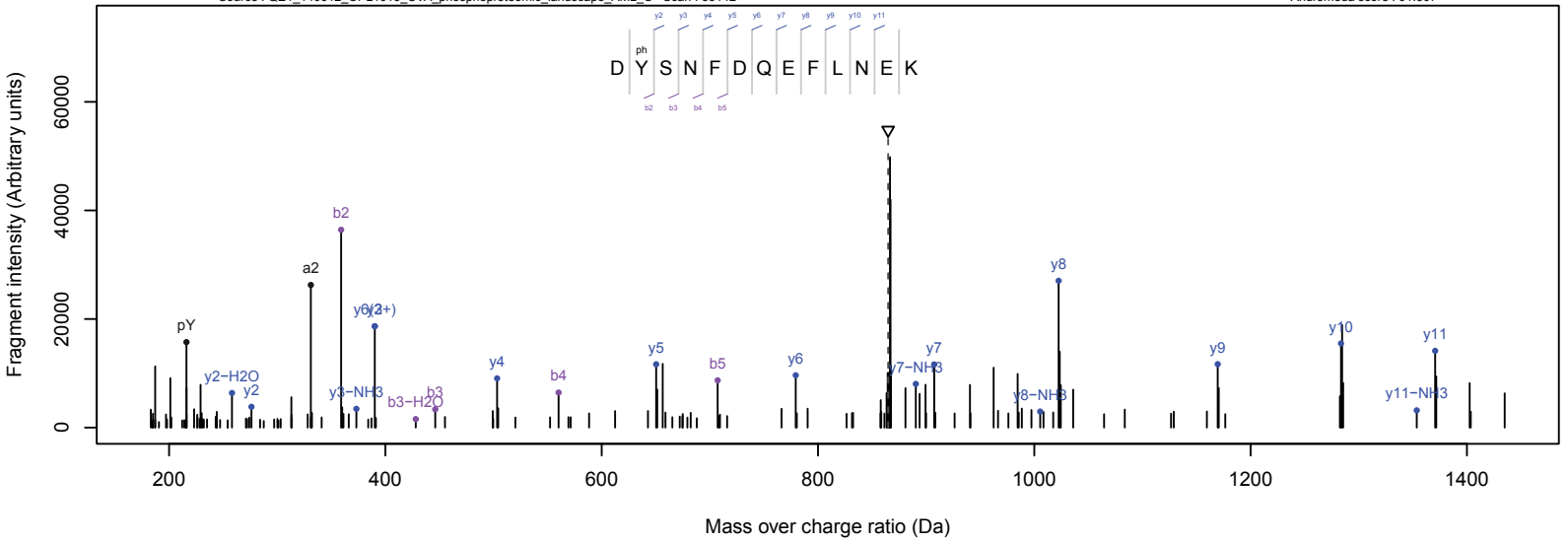
Andromeda score: 91.313



DY(ph)SNFDQEFLEK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 33142

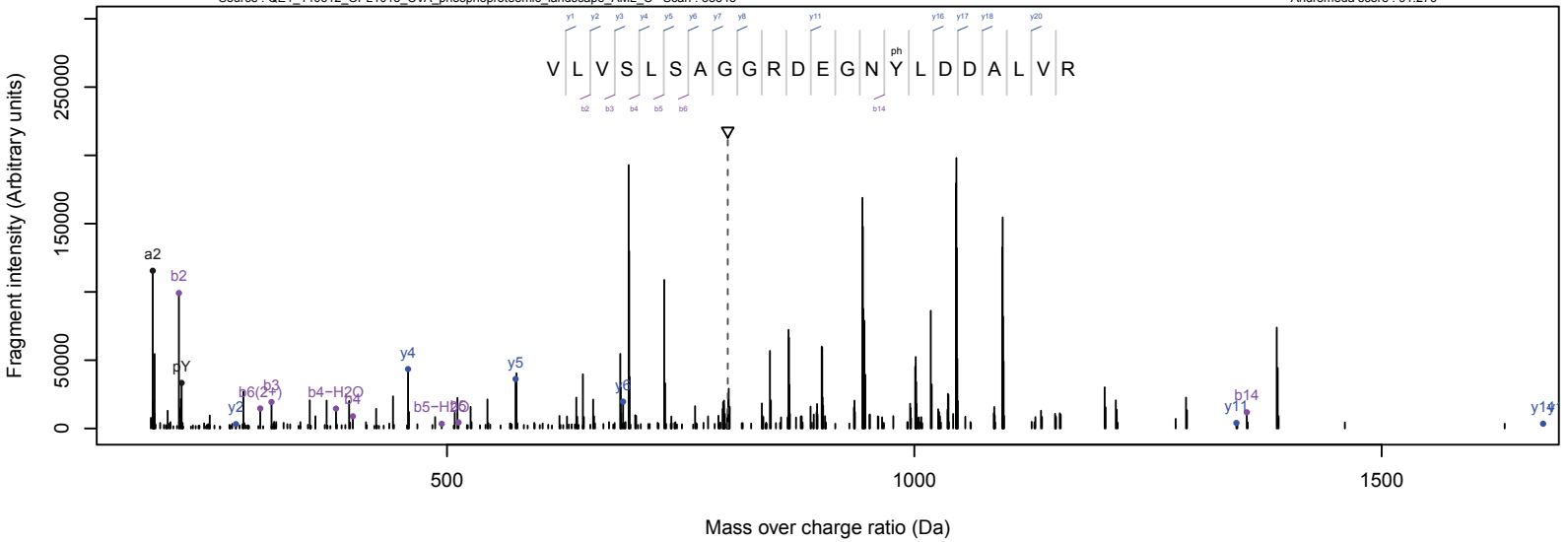
Andromeda score: 91.307



VLVLSAGGRDEGNY(ph)LDDALVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38343

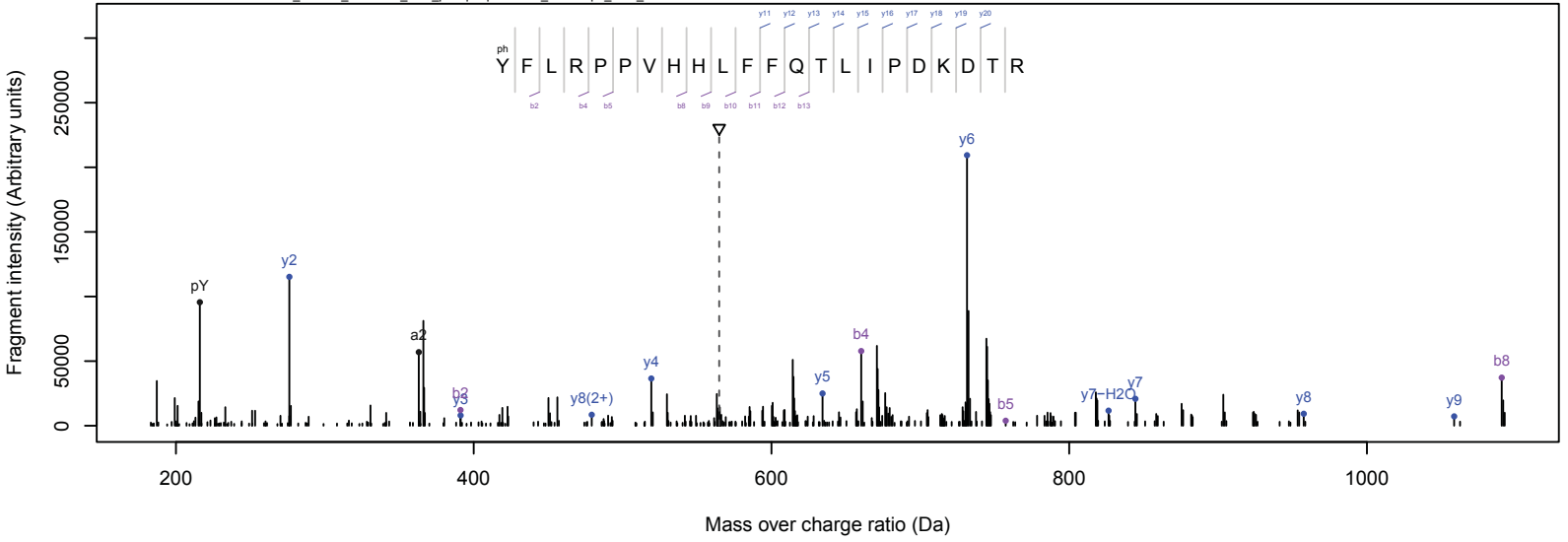
Andromeda score: 91.276



Y(ph)FLRPPVHHLFFQTLIPDKDTR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43797

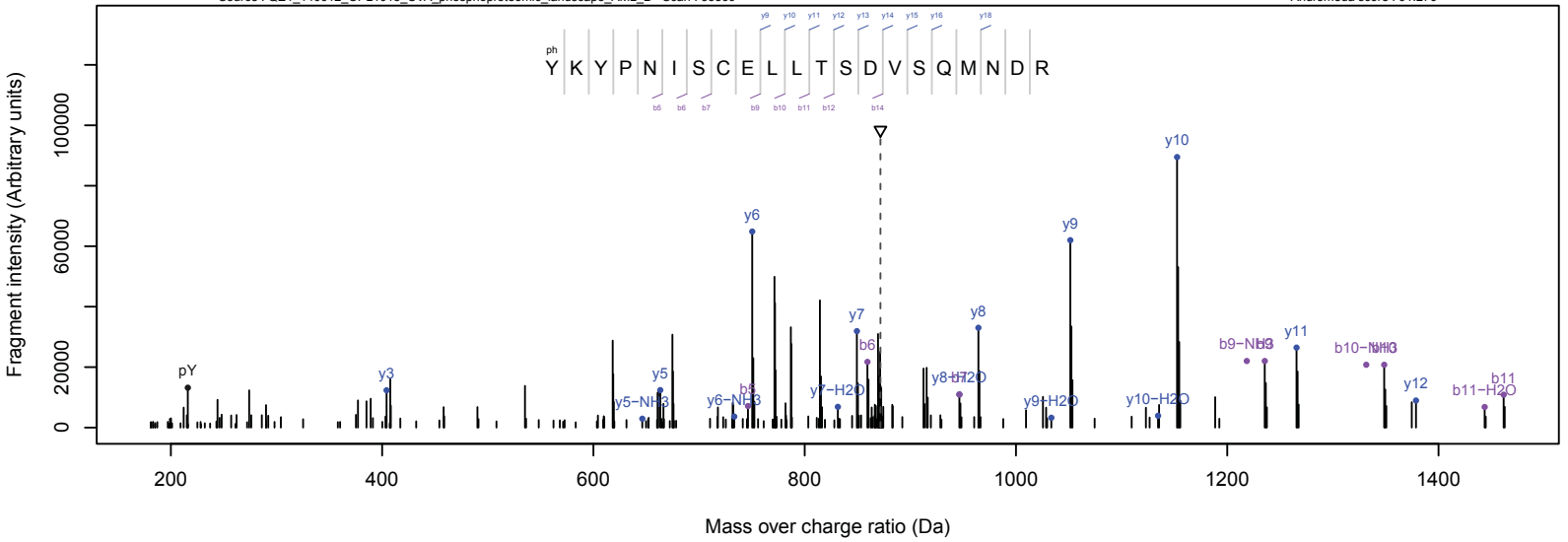
Andromeda score : 91.276



Y(ph)KYPNISCELLTSDV SQMND R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35869

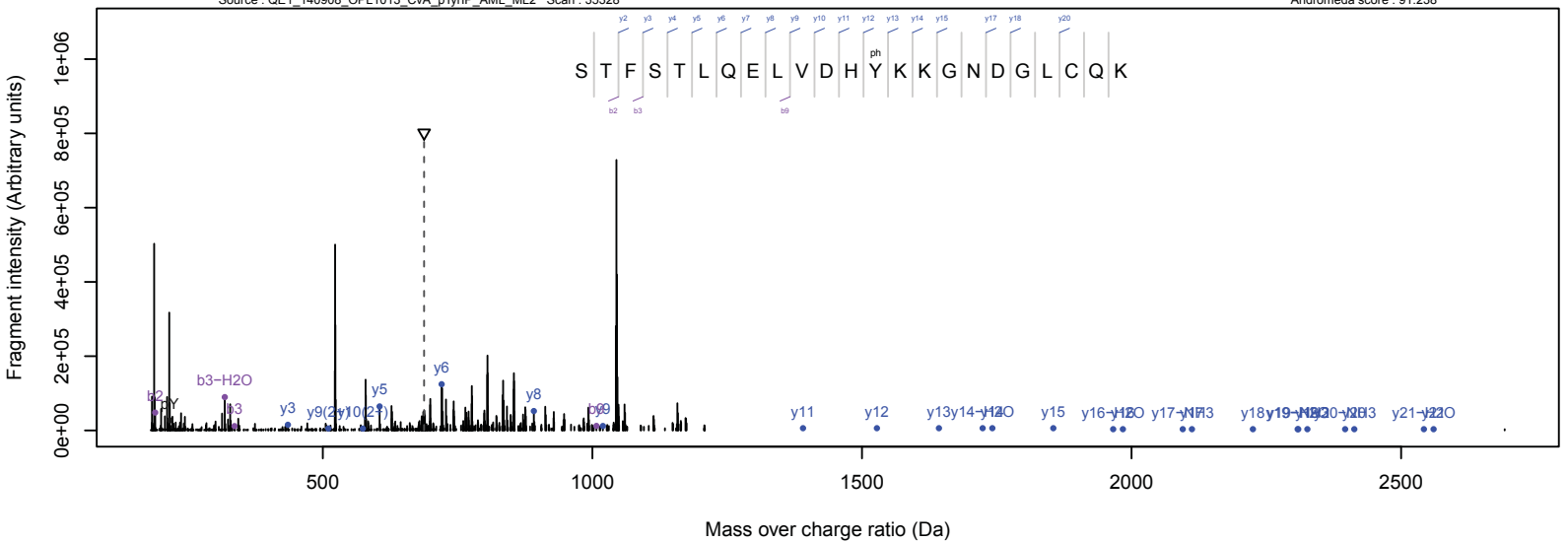
Andromeda score : 91.276



STFSTLQELVDHY(ph)KKGNDGLCQK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 35328

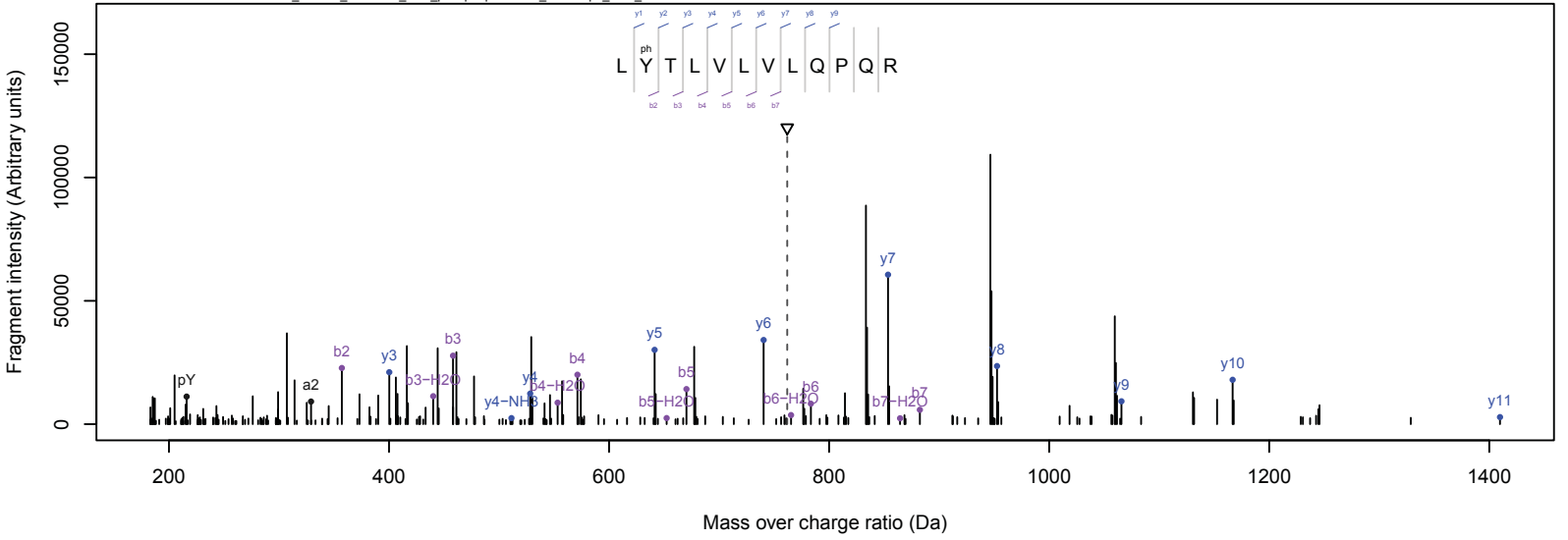
Andromeda score : 91.238



LY(ph)TLVLVLPQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 46858

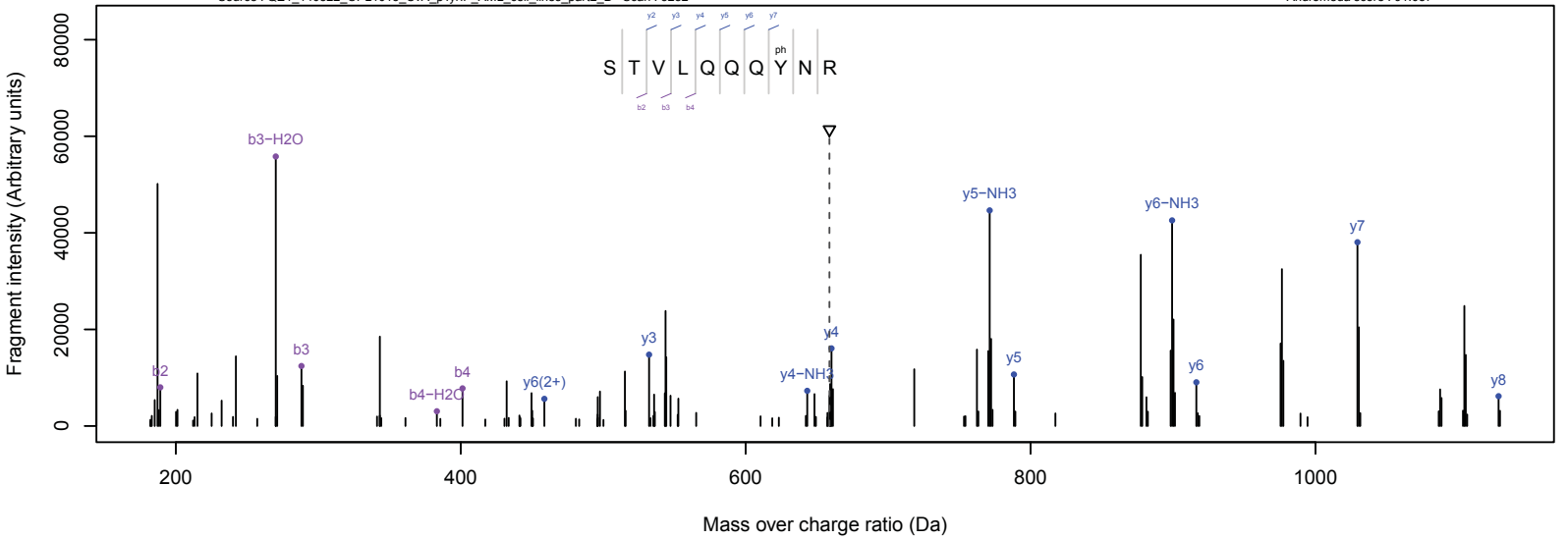
Andromeda score : 91.123



STVLQQY(ph)NR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 5282

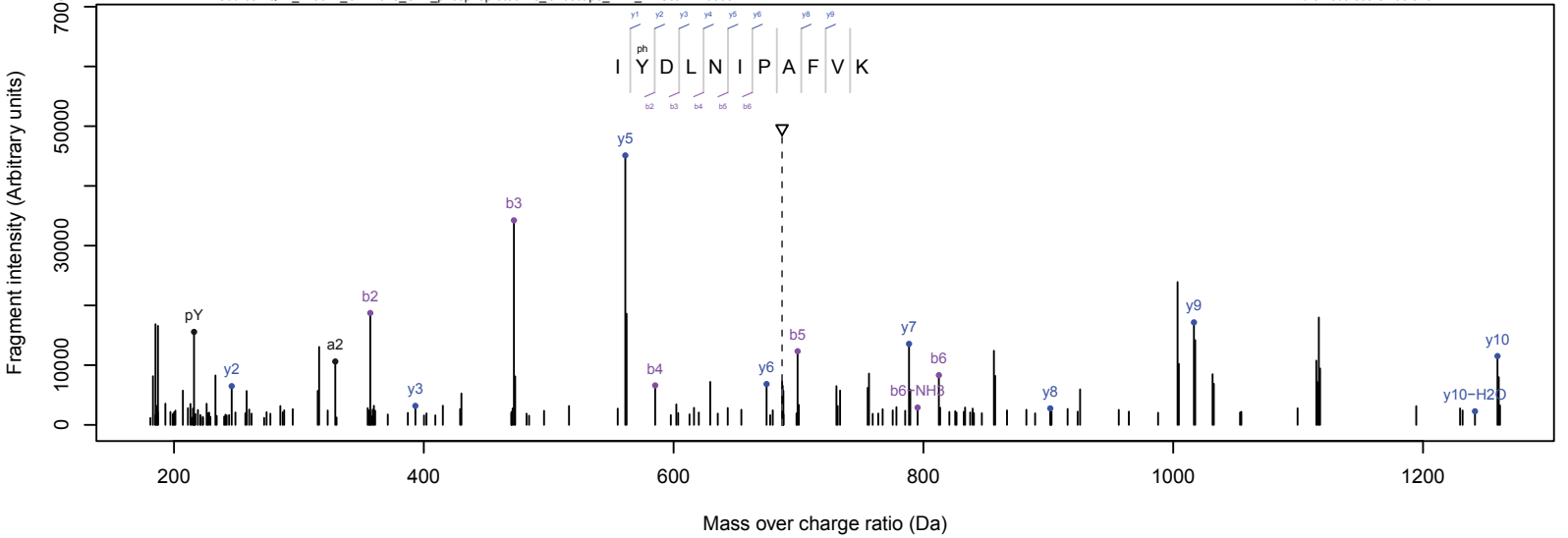
Andromeda score : 91.087



IY(ph)DLNIPAFVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43355

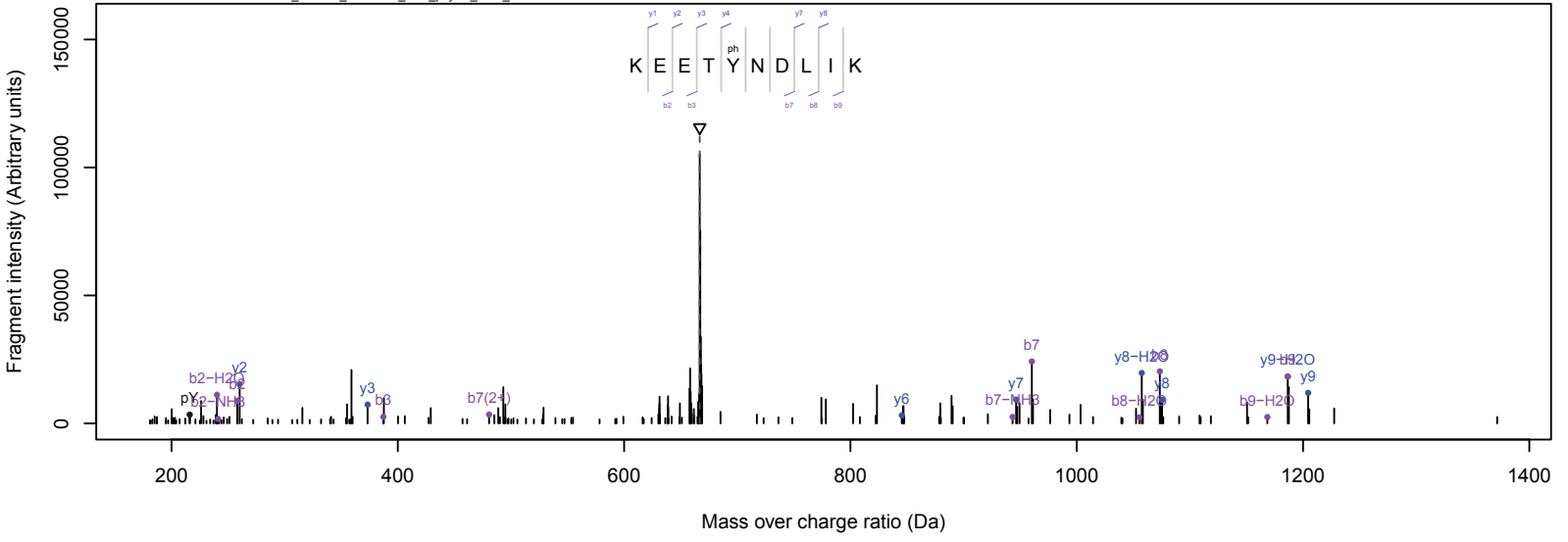
Andromeda score : 90.913



KEETY(ph)NDLIK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 13993

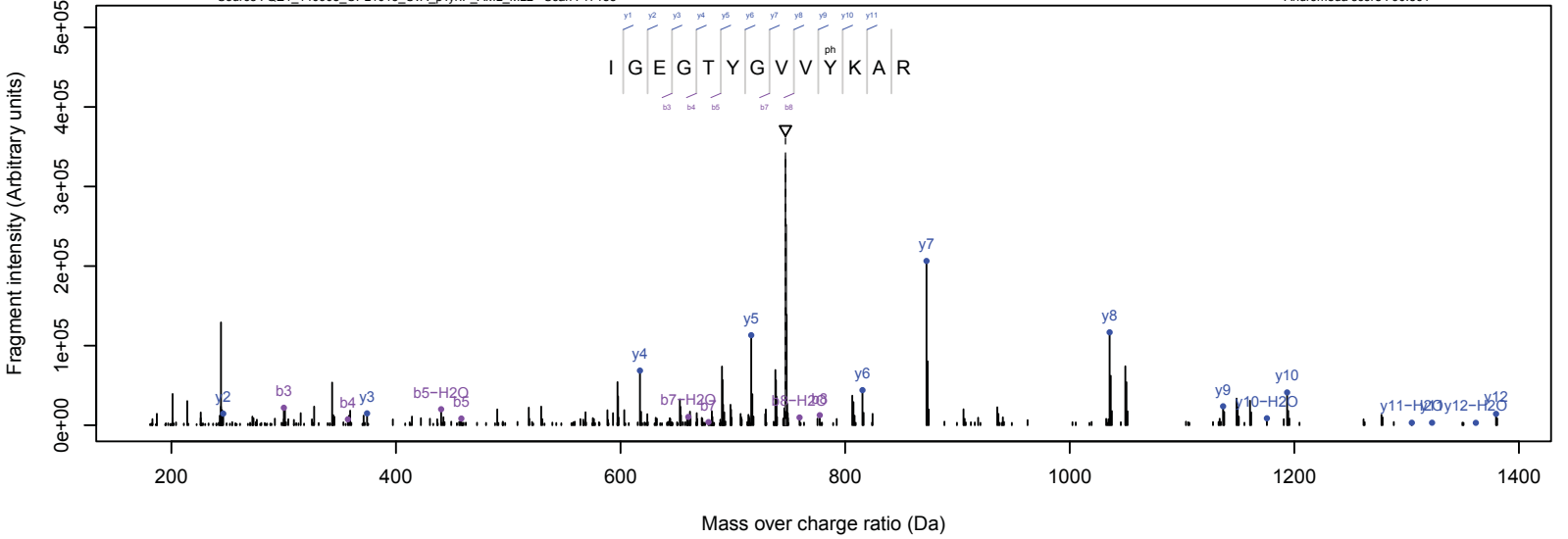
Andromeda score : 90.913



IGEGTYGVVY(ph)KAR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 17188

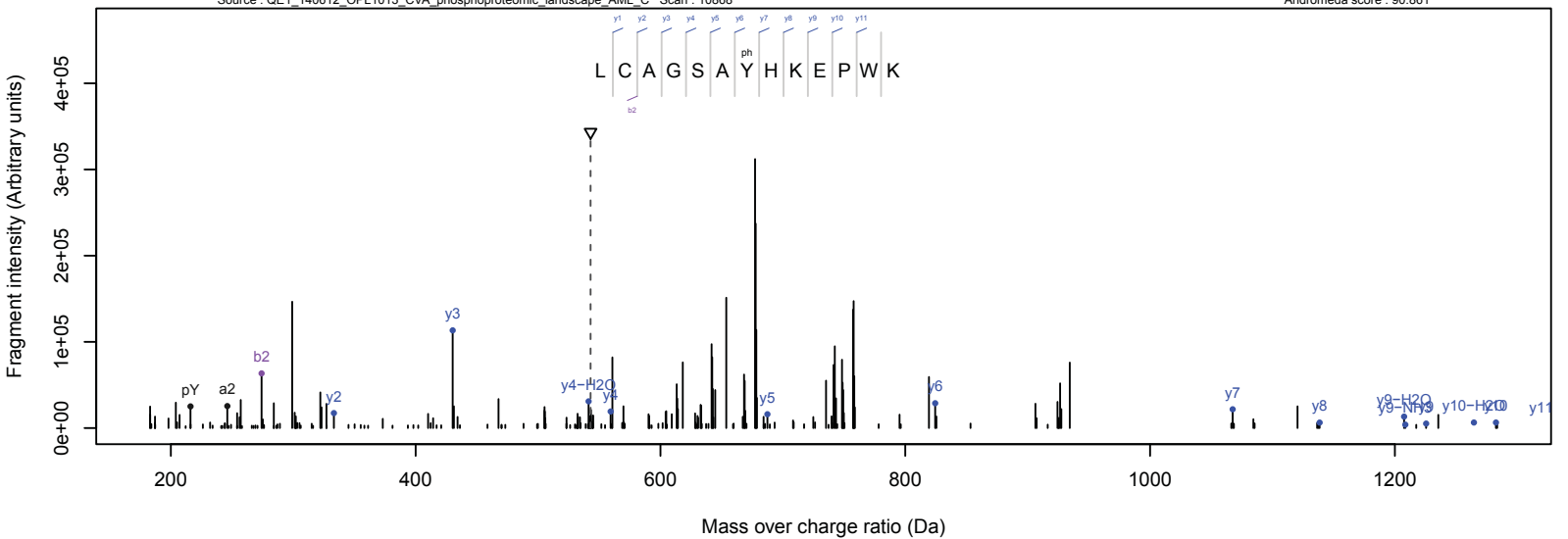
Andromeda score : 90.861



LCAGSAY(ph)HKEPWK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10868

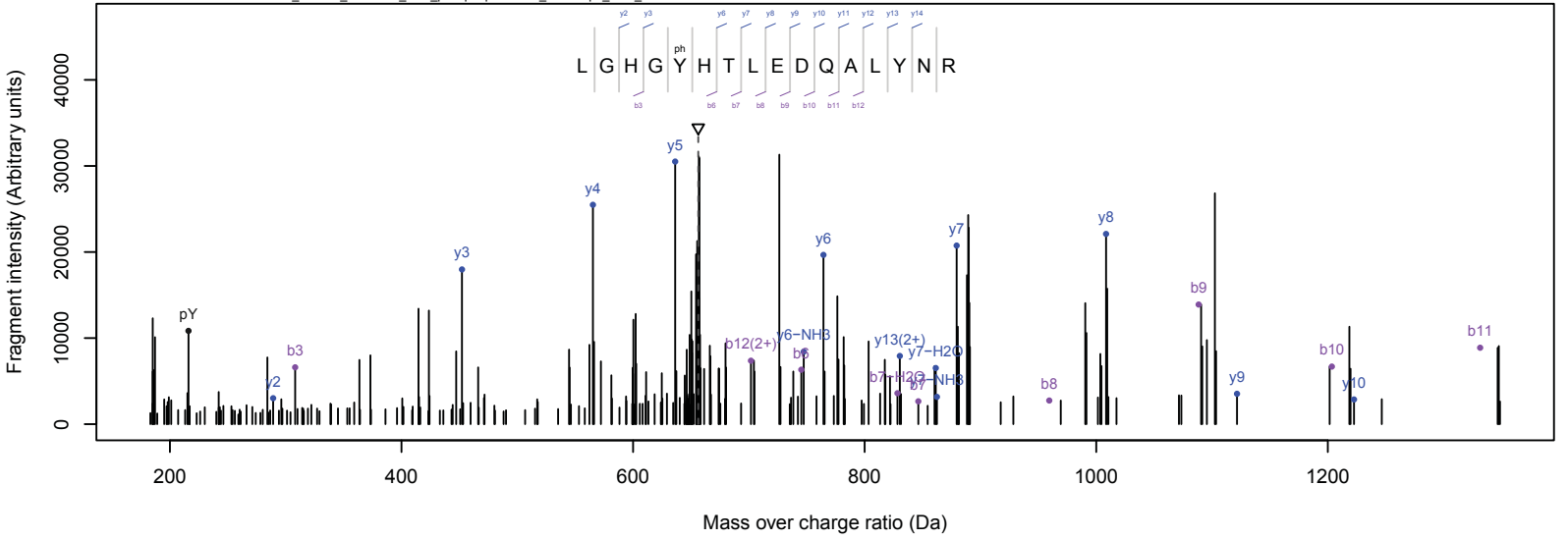
Andromeda score : 90.861



**LGHGY(ph)HTLEDQALYNR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18989

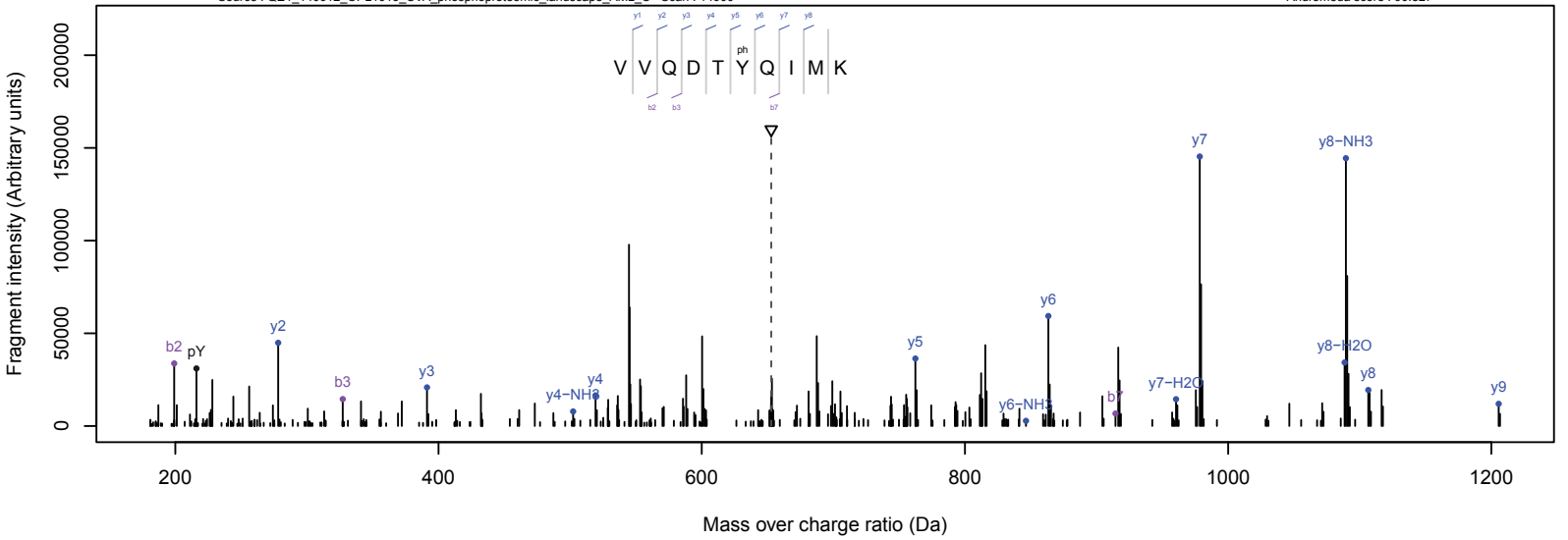
Andromeda score : 90.853



**VVQDTY(ph)QIMK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14900

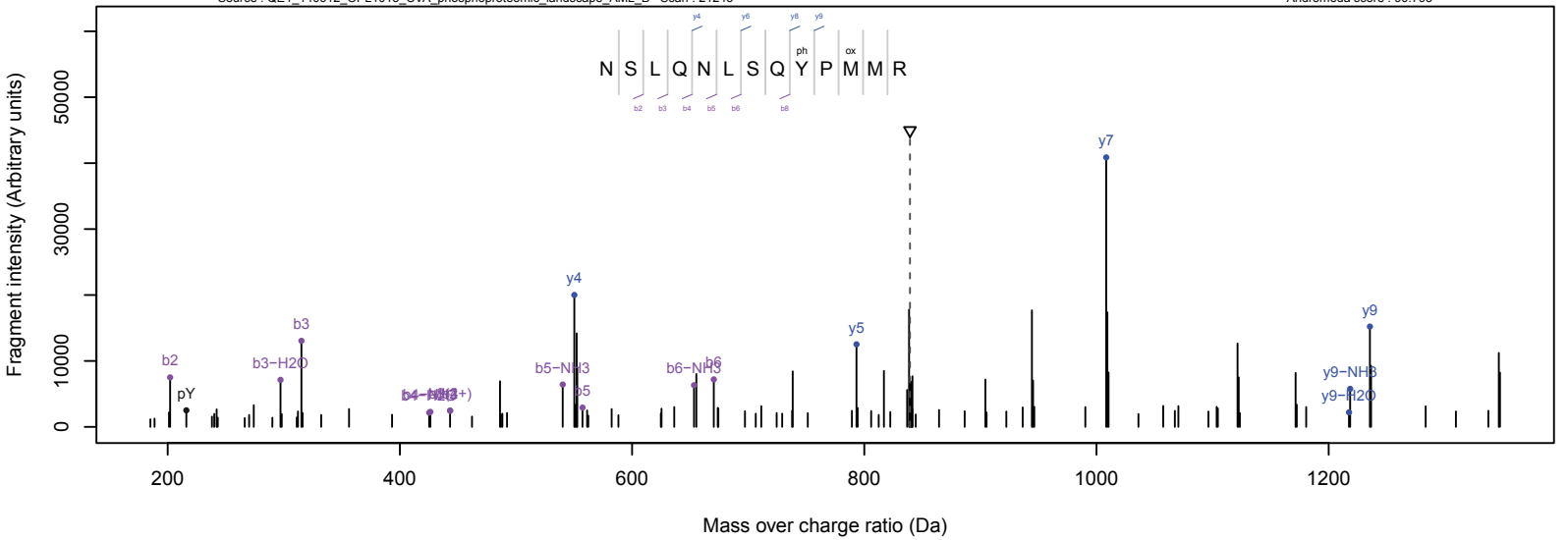
Andromeda score : 90.827



**NSLQNL S Q Y P M M R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21245

Andromeda score : 90.793



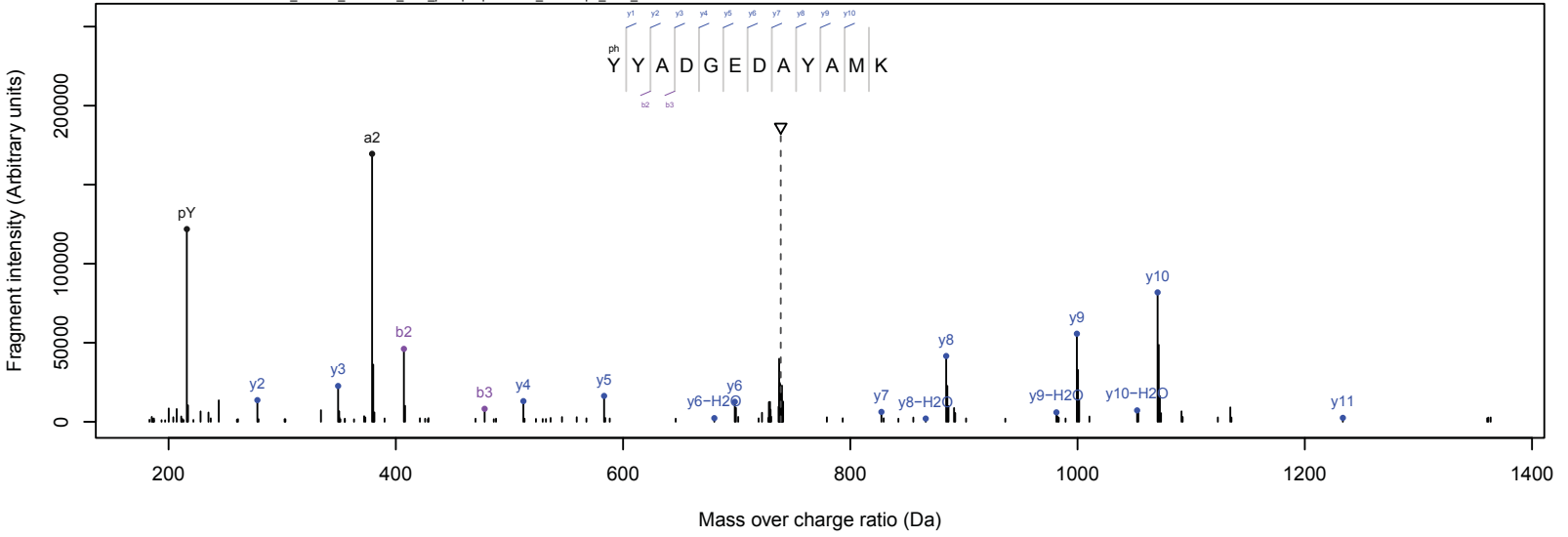




Y(ph)YADGEDAYAMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15389

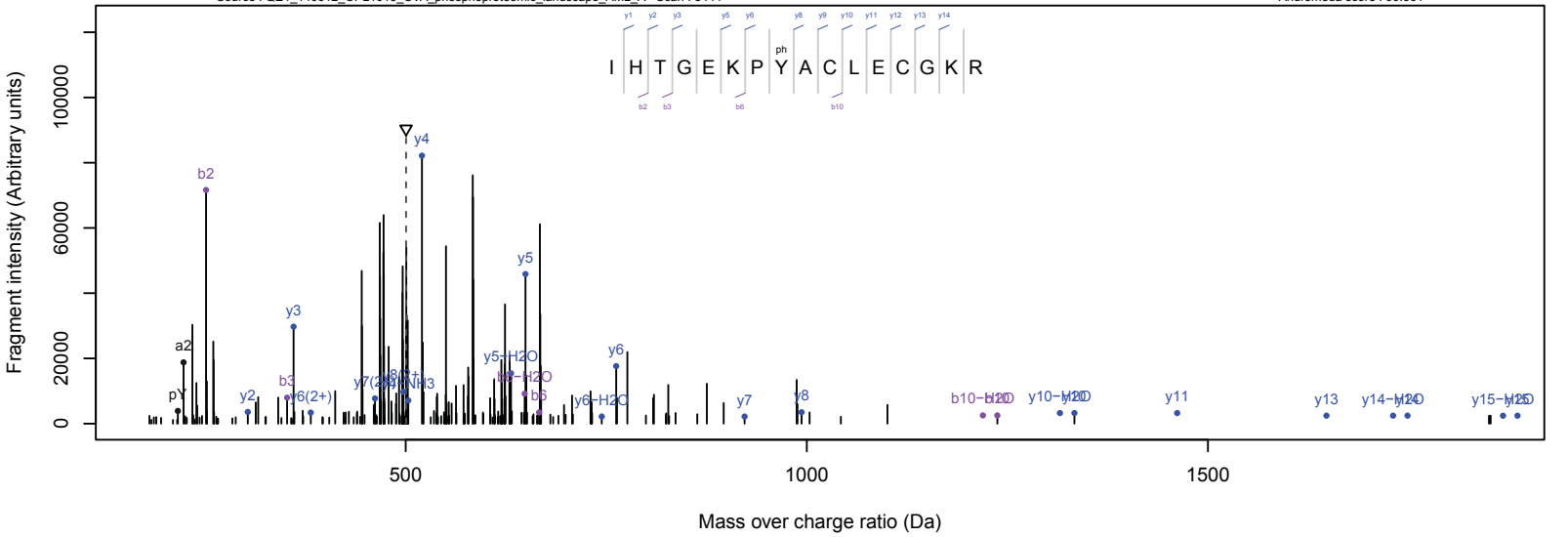
Andromeda score : 90.614



IHTGKPY(ph)ACLECGKR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 8444

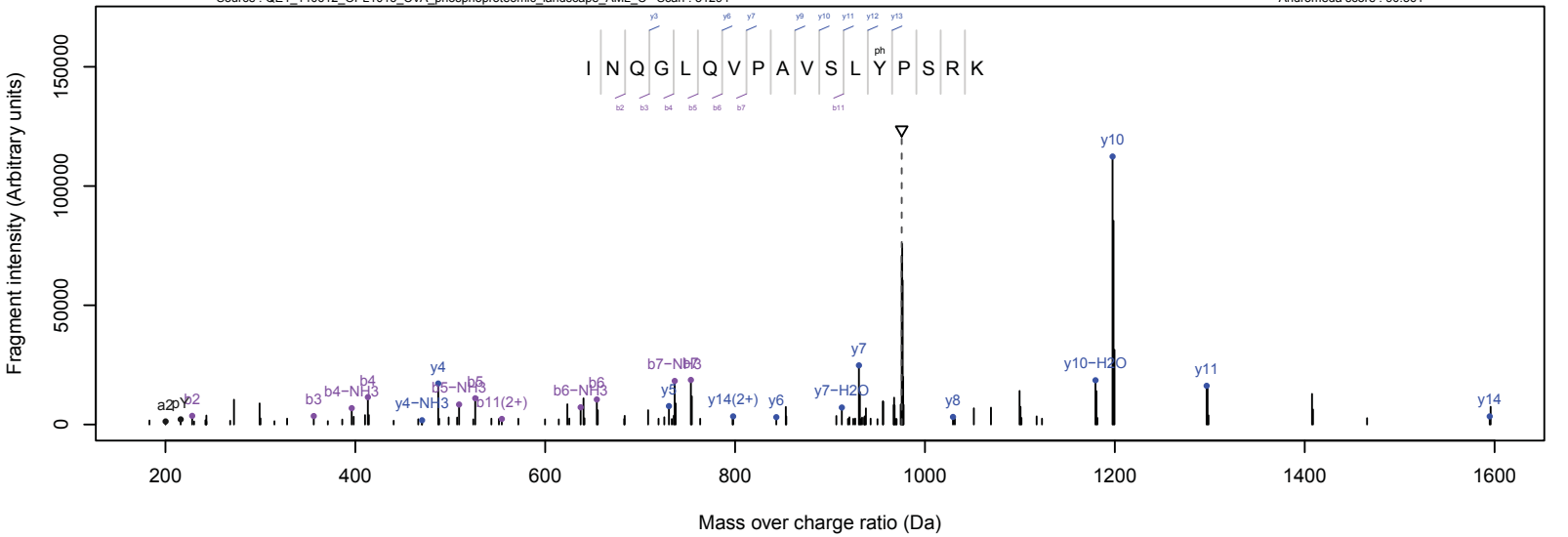
Andromeda score : 90.561



INQLQVPAVSLYP SRK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31294

Andromeda score : 90.561

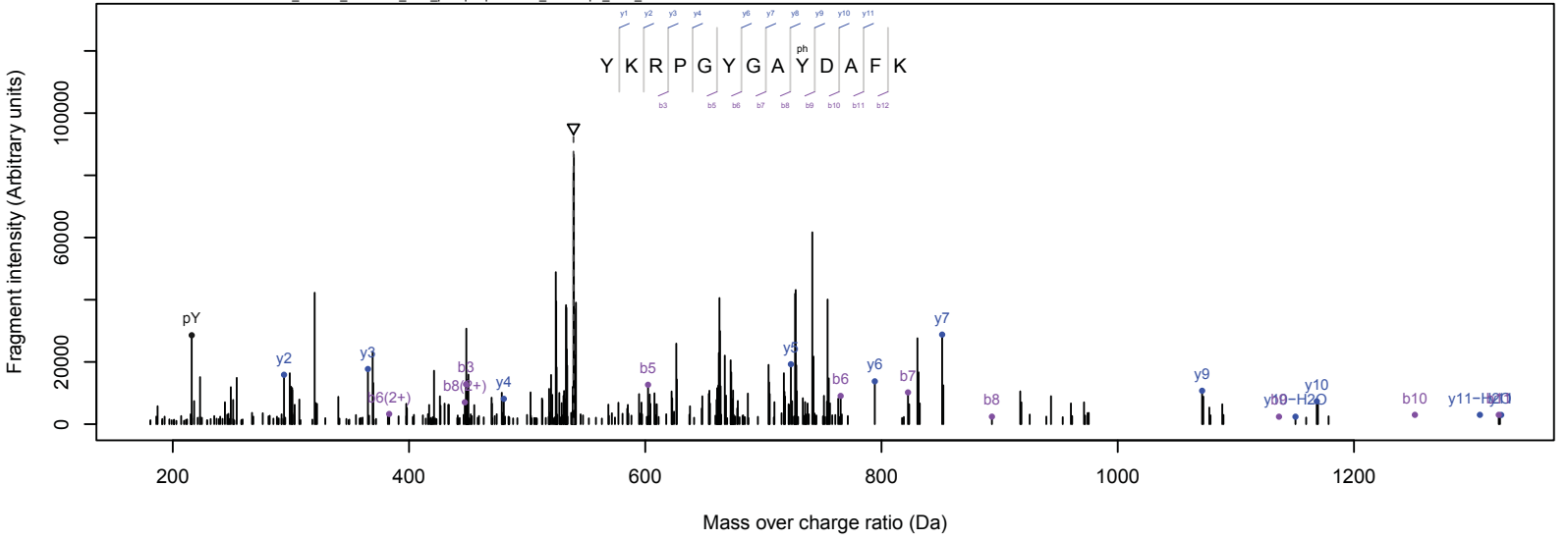




YKRPGYGAY(ph)DAFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 14291

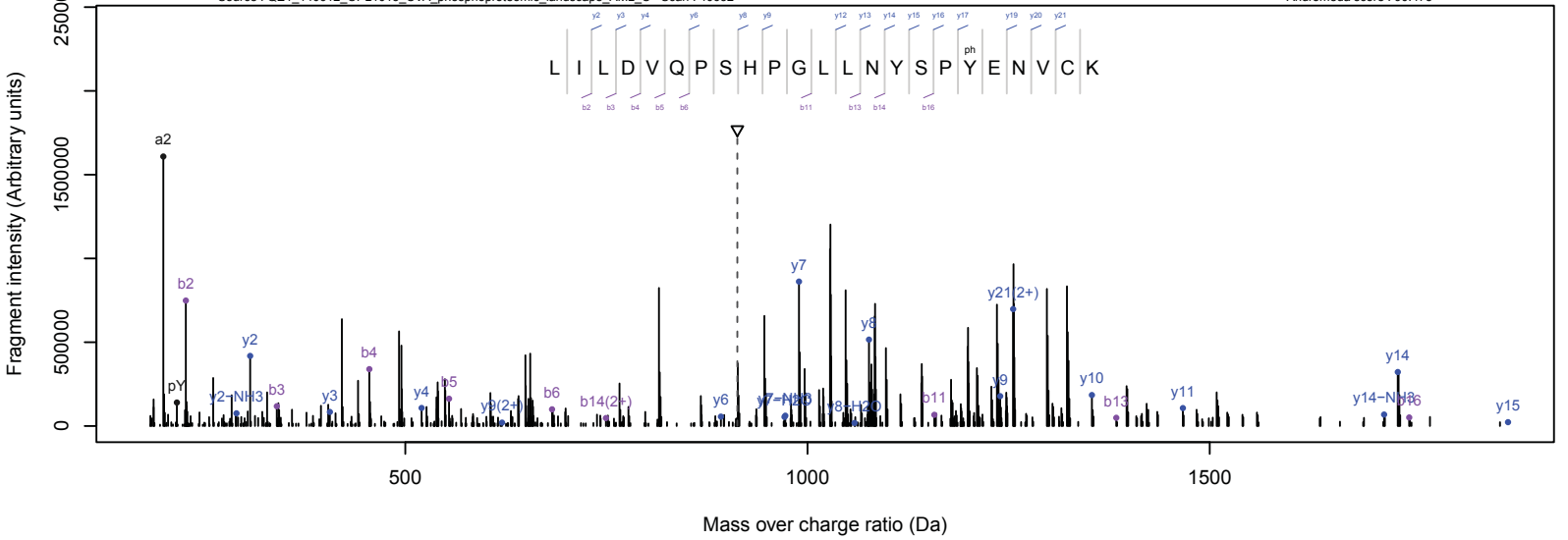
Andromeda score : 90.476



LILDVQSPHPLLNSPY(ph)ENVCK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40632

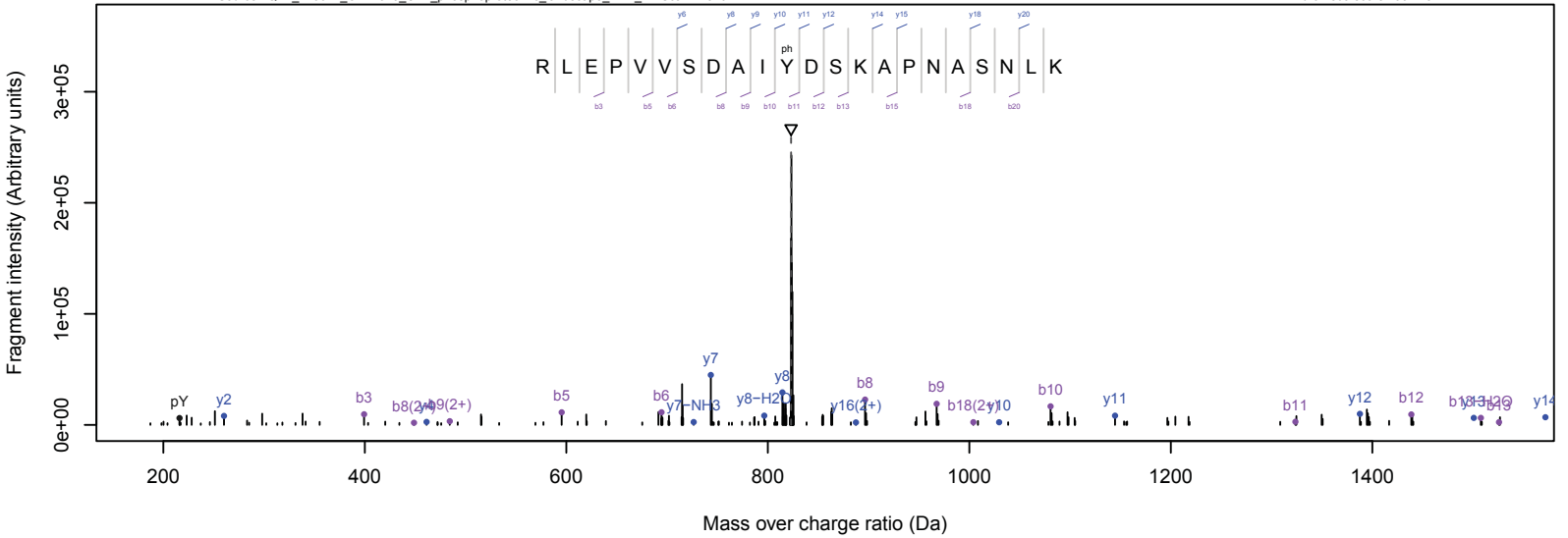
Andromeda score : 90.475



RLEPVSDAIY(ph)DSKAPNASNLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28201

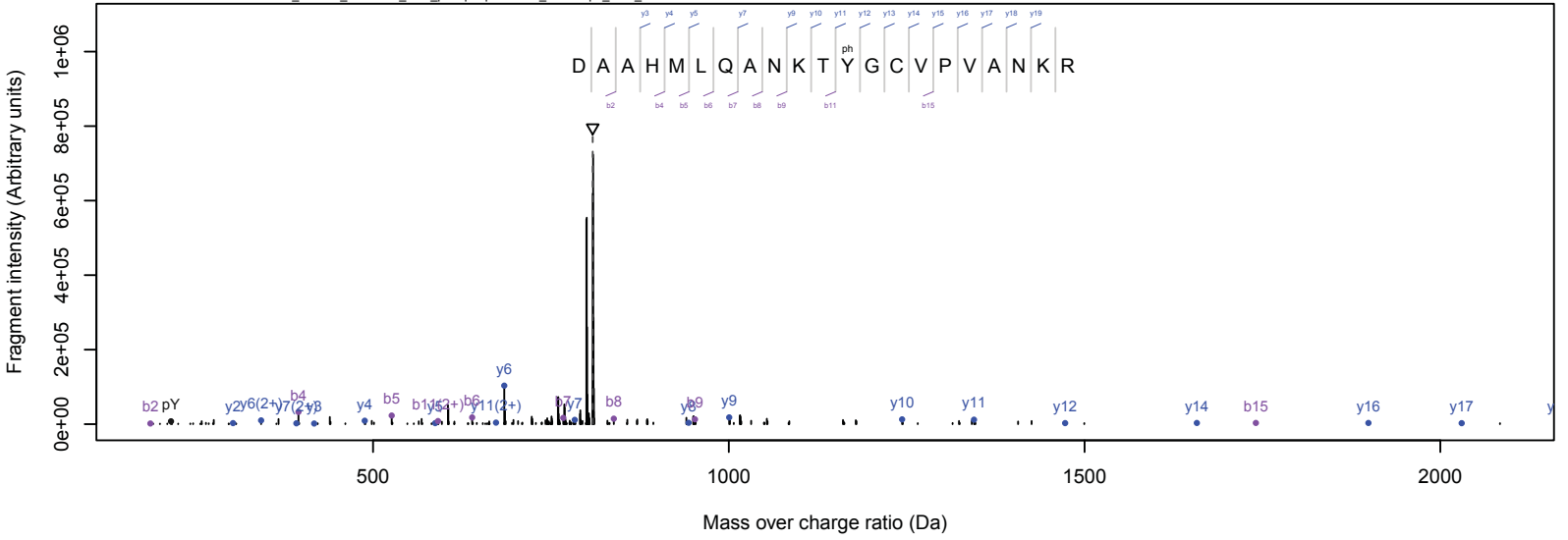
Andromeda score : 90.475



**DAAHMLQANKTY(ph)GCVPVANKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17183

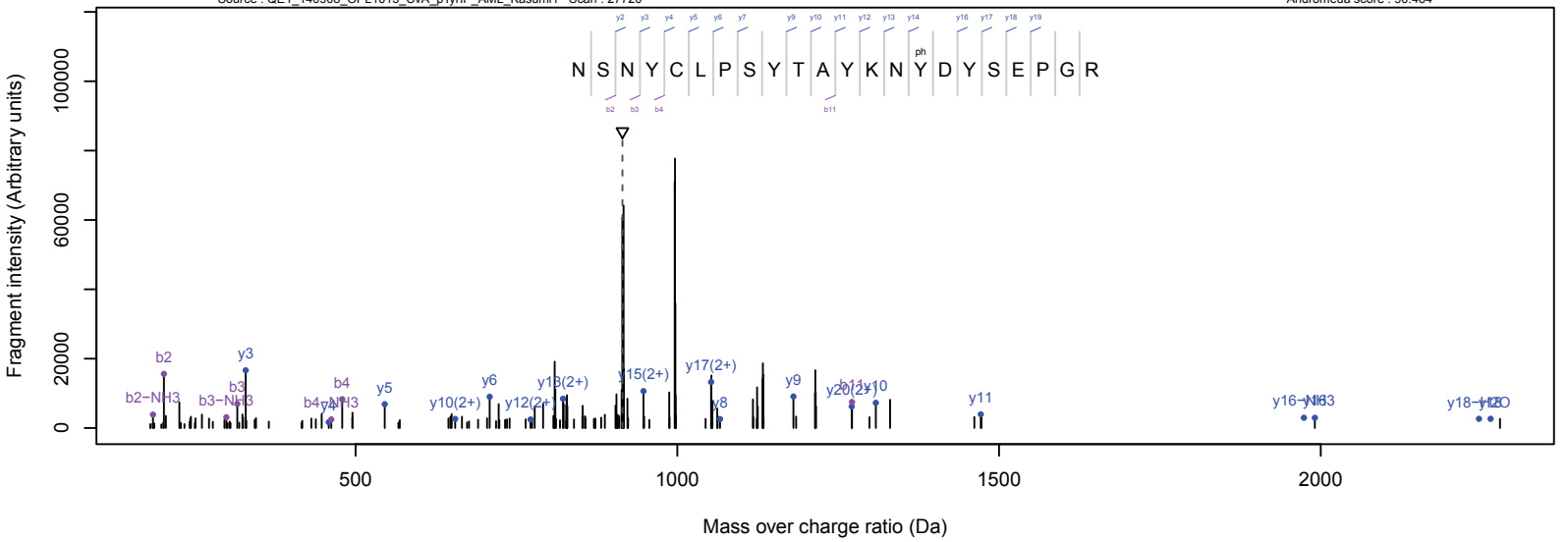
Andromeda score : 90.464



**NSNYCLPSYATYKNYDYSEPR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27720

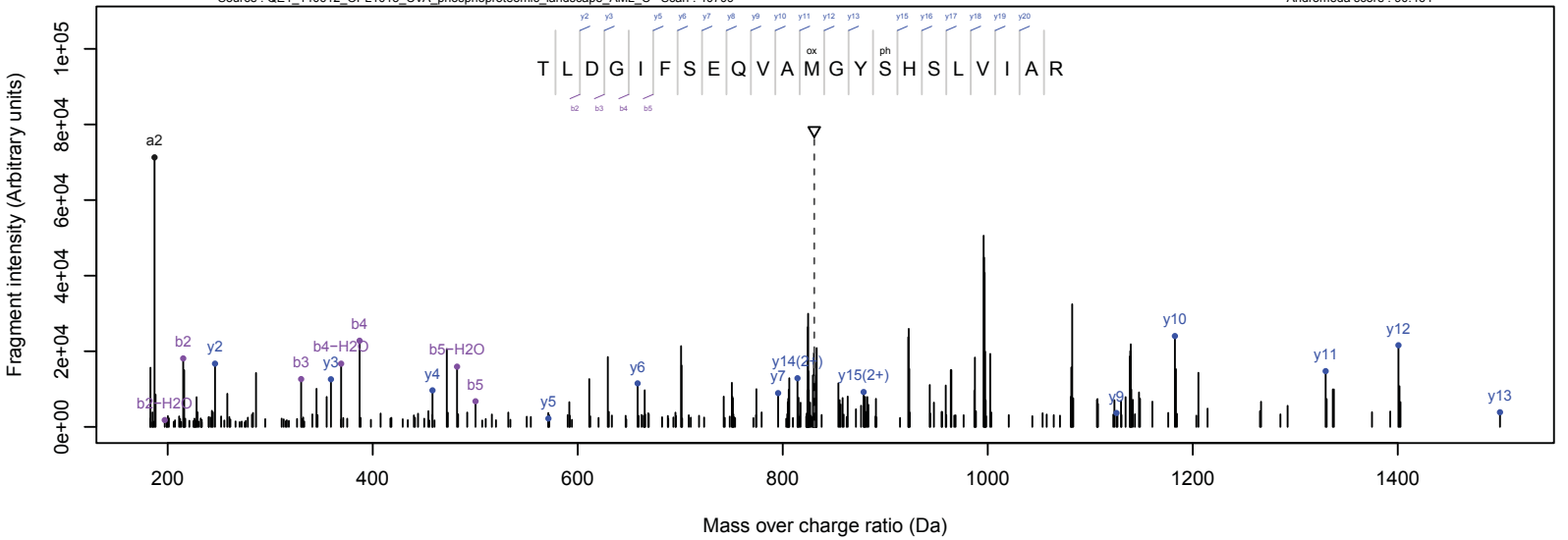
Andromeda score : 90.464



**TLDGIFSEQVAMGYS(ph)HSLVIAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40709

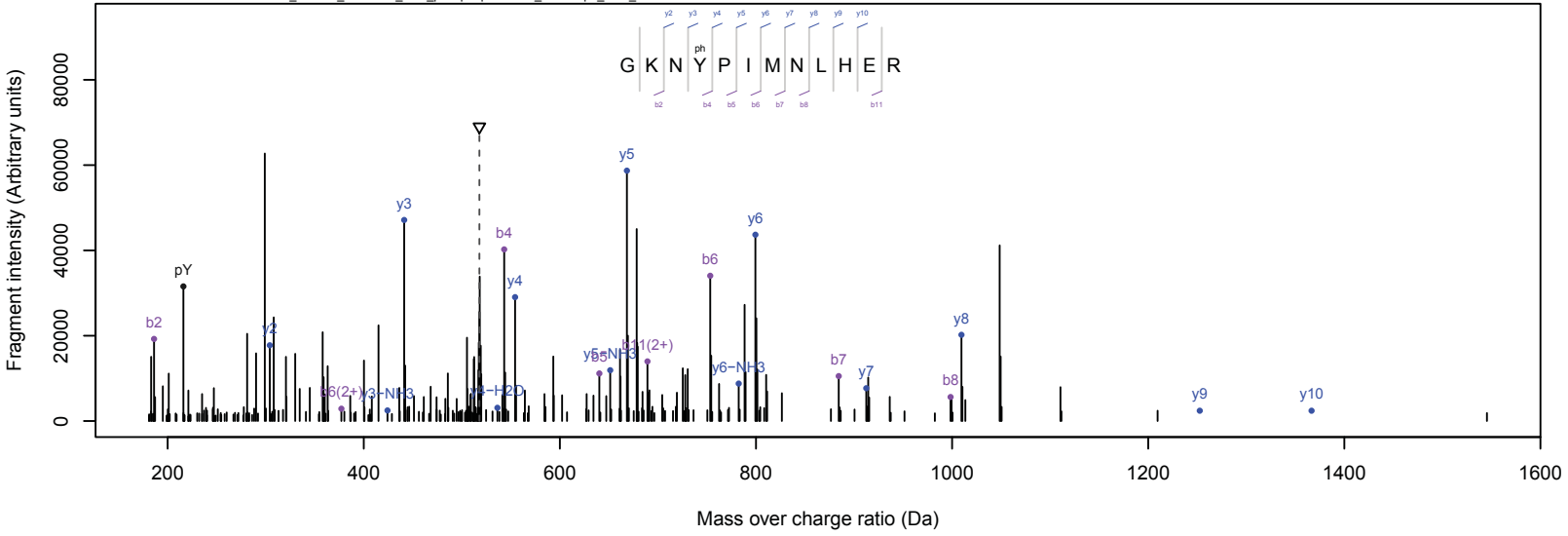
Andromeda score : 90.464



GKNY(ph)PIMNLHER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13768

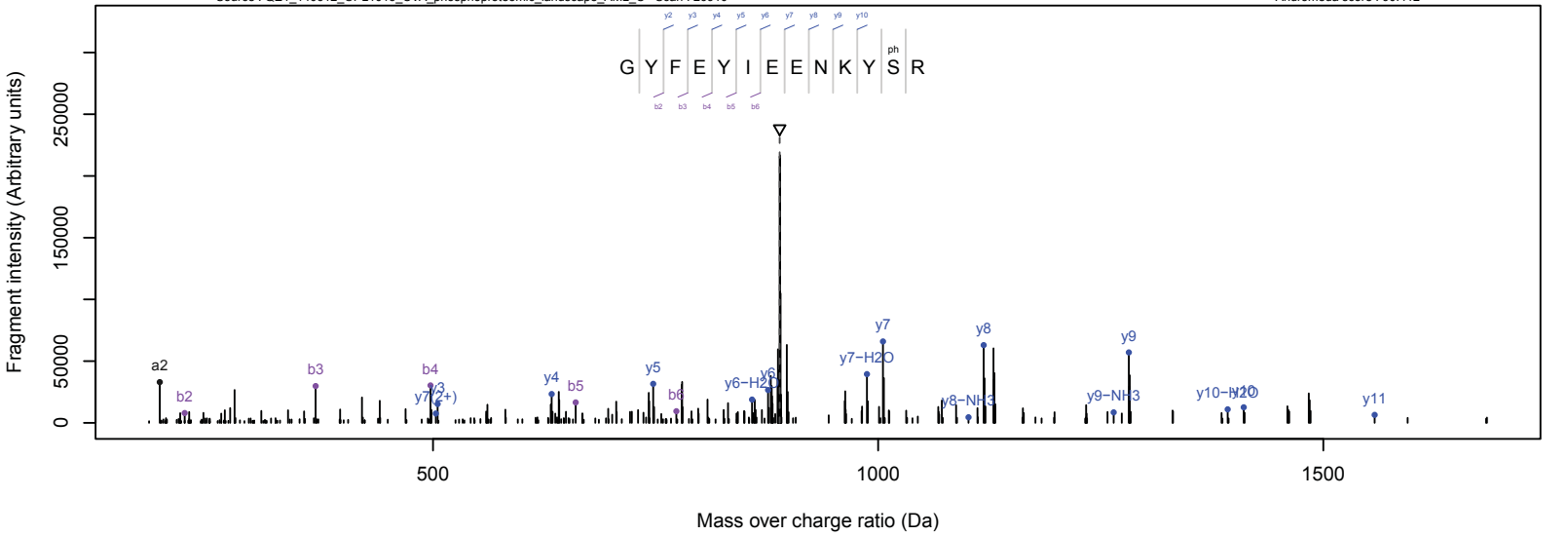
Andromeda score : 90.412



GYFEYIEENKYS(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26910

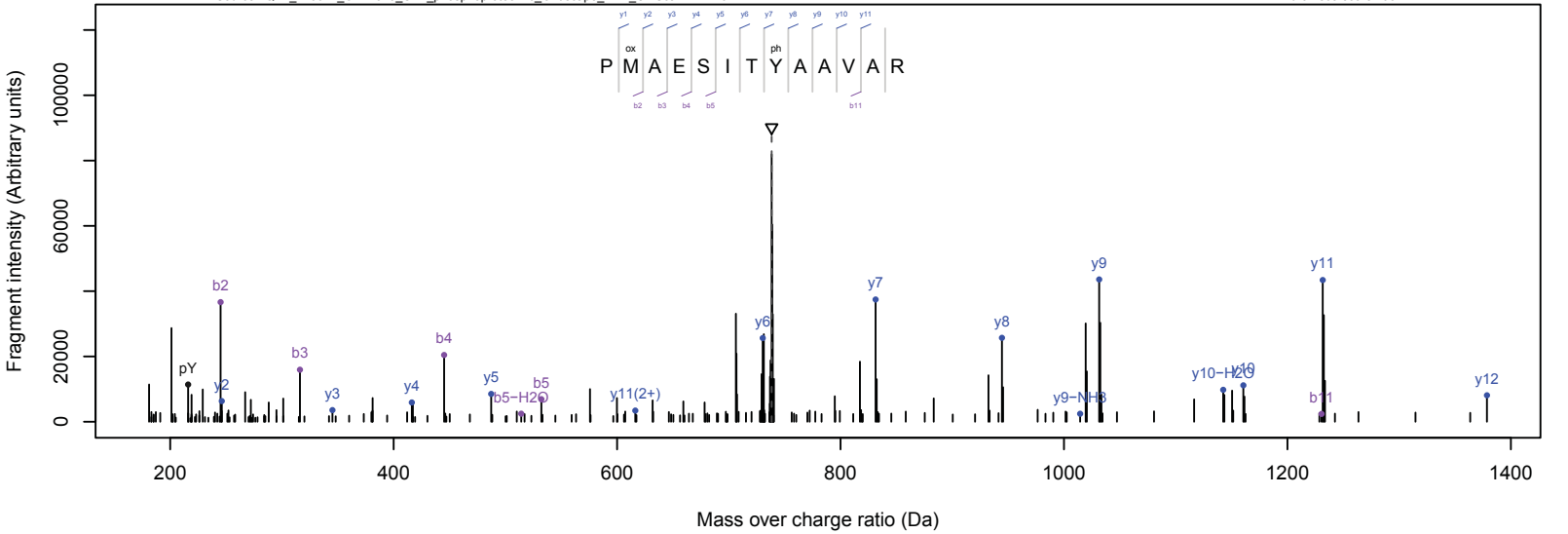
Andromeda score : 90.412



PM(ox)AESITY(ph)AAVAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24110

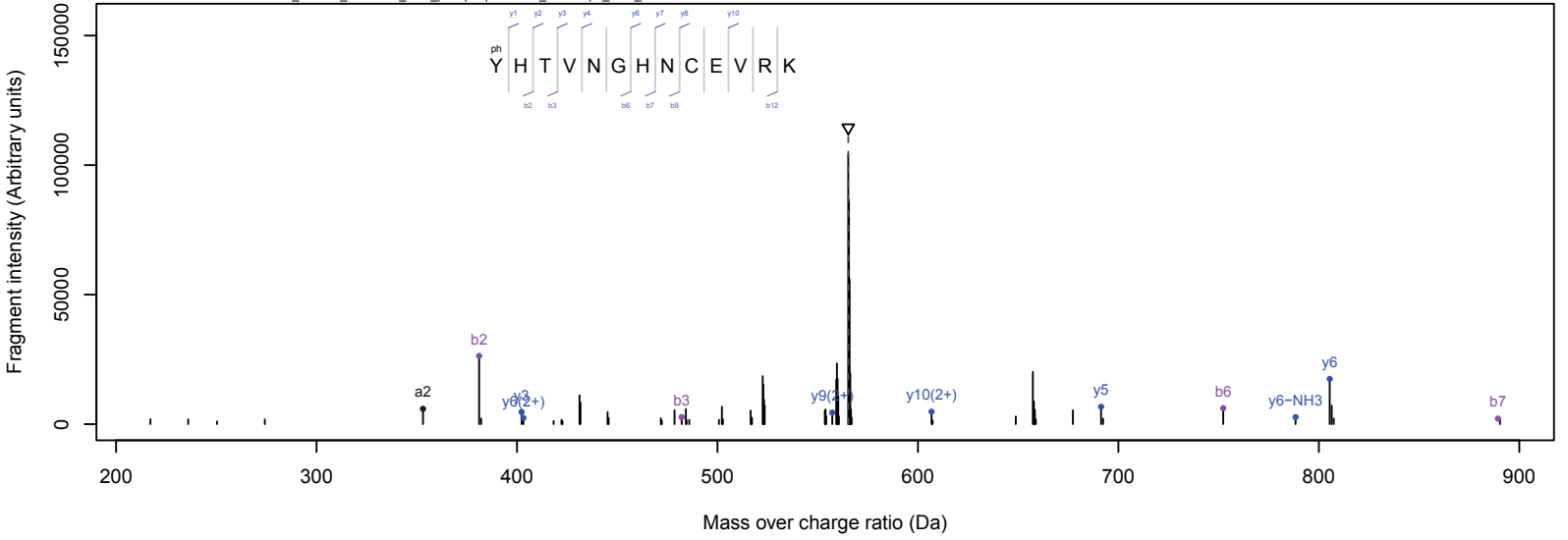
Andromeda score : 90.412



Y(ph)HTVNGHNCEVRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 3436

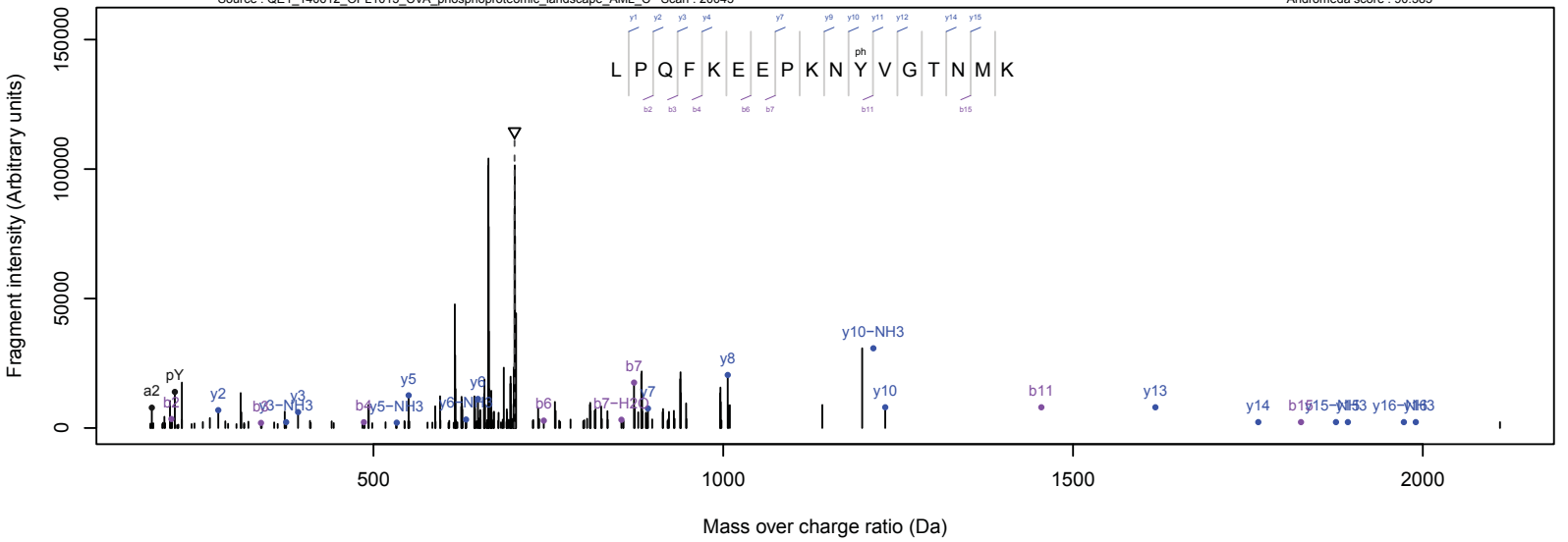
Andromeda score : 90.412



LPQFKEEPPKNYVGTNMG-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20045

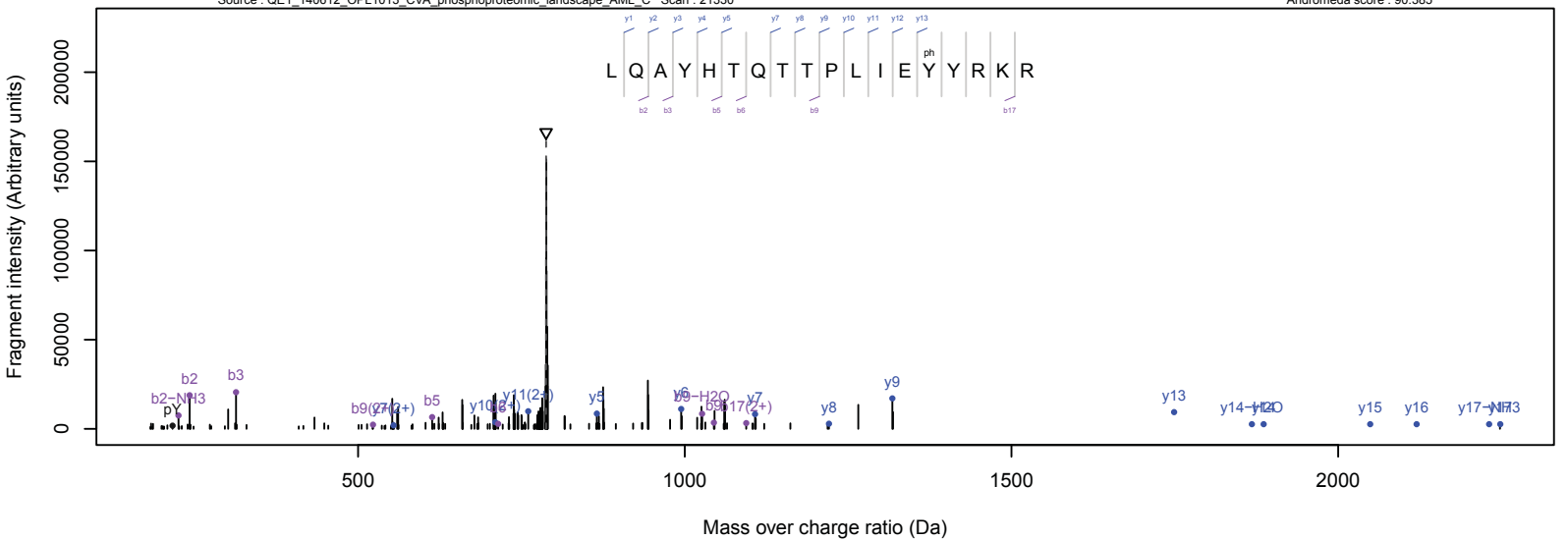
Andromeda score : 90.385



LQAYHTQTTPLEIYRKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21330

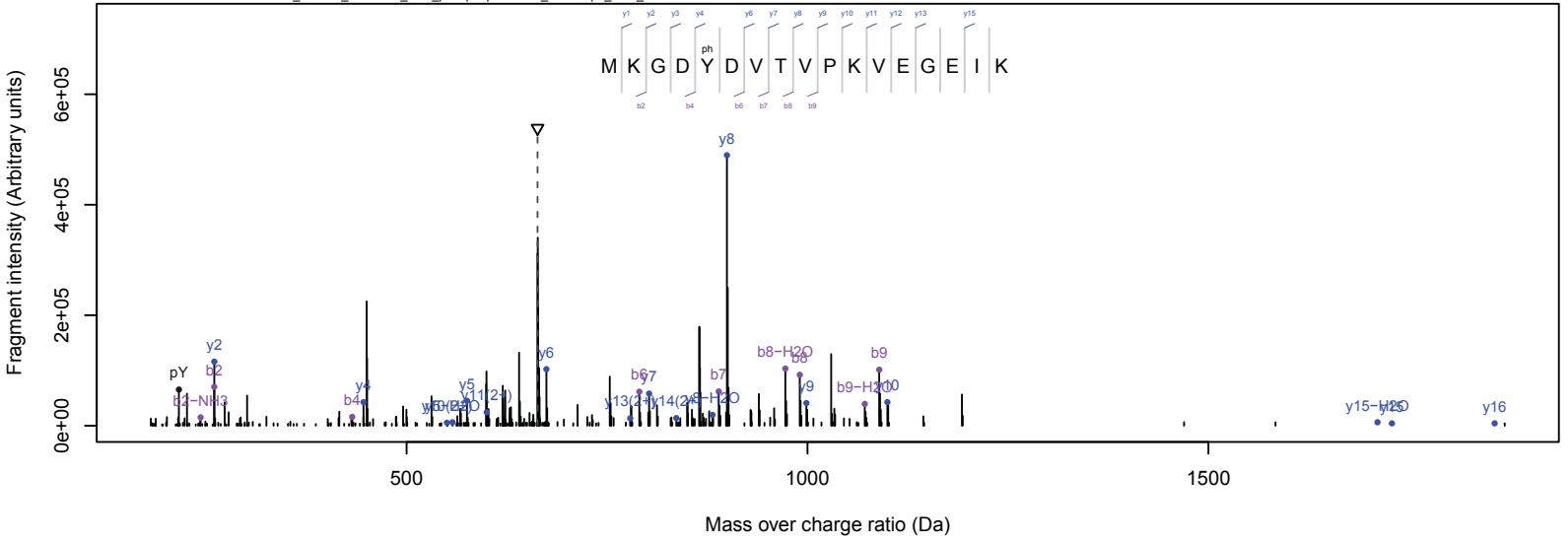
Andromeda score : 90.385



MKGDY(ph)DVTVPKVEGEIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22655

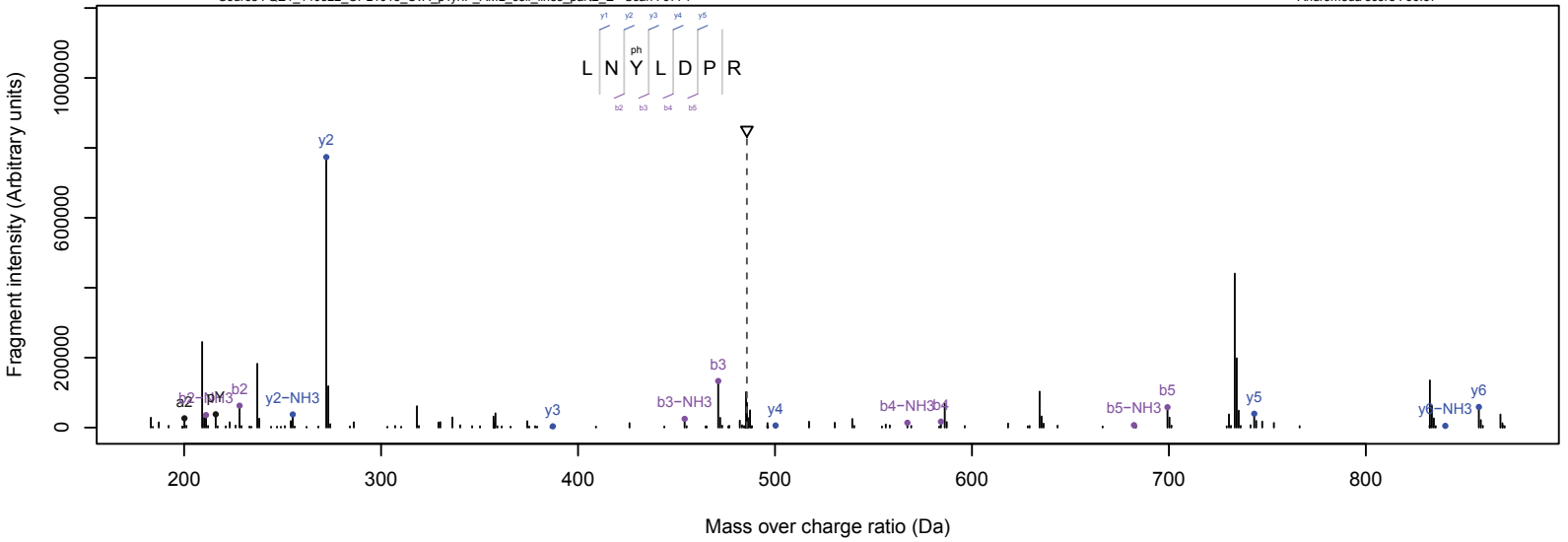
Andromeda score : 90.385



LNY(ph)LDPR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 9774

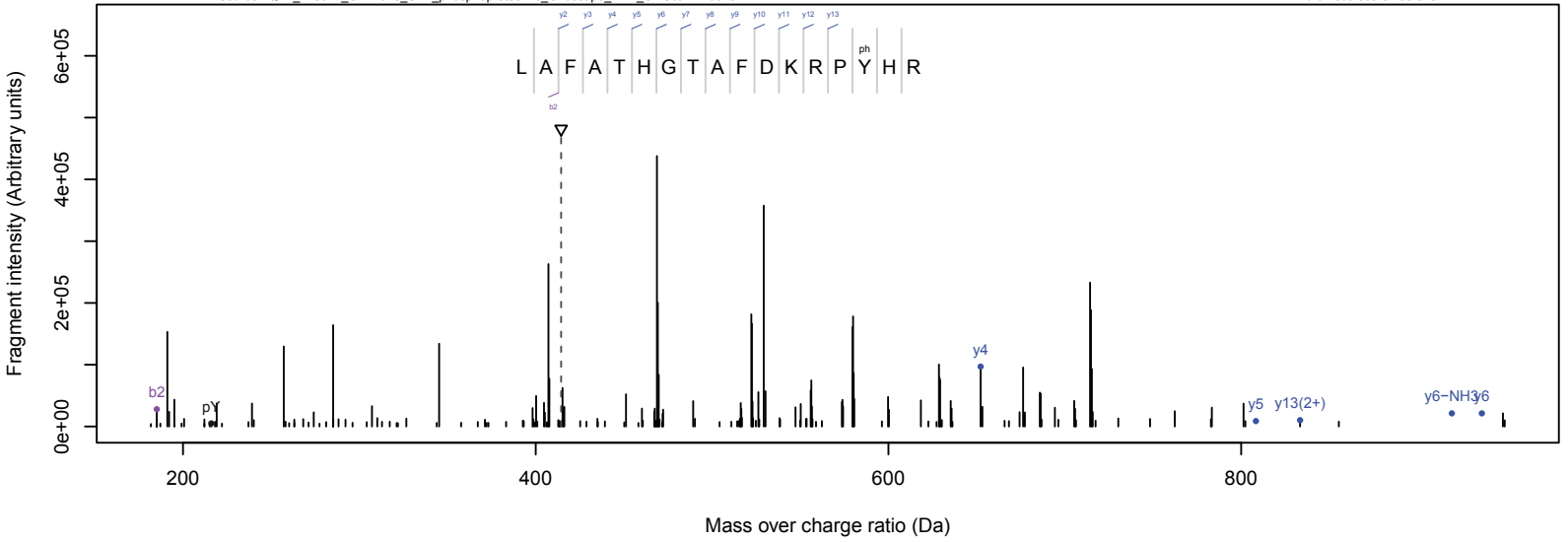
Andromeda score : 90.37



LAFATHGTAFDKRPY<sup>ph</sup>HR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15323

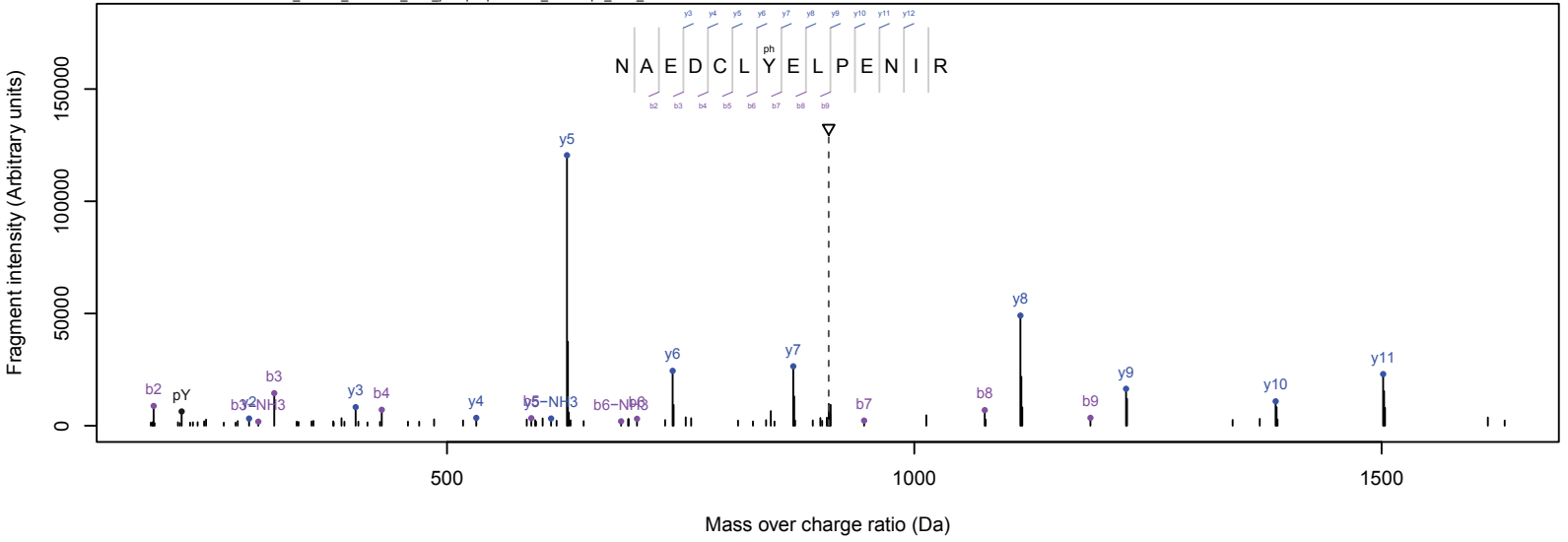
Andromeda score : 90.348



NAEDCLY(ph)ELPENIR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 32679

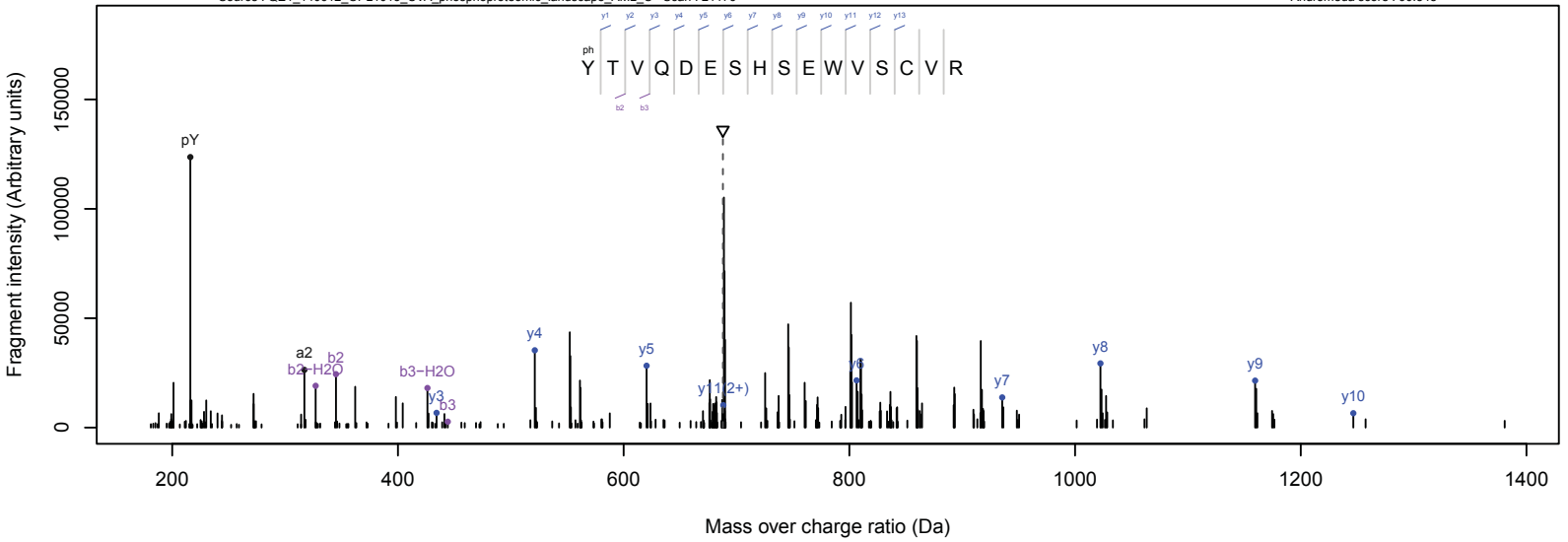
Andromeda score: 90.348



Y(ph)TVQDESHSEWVSCVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21176

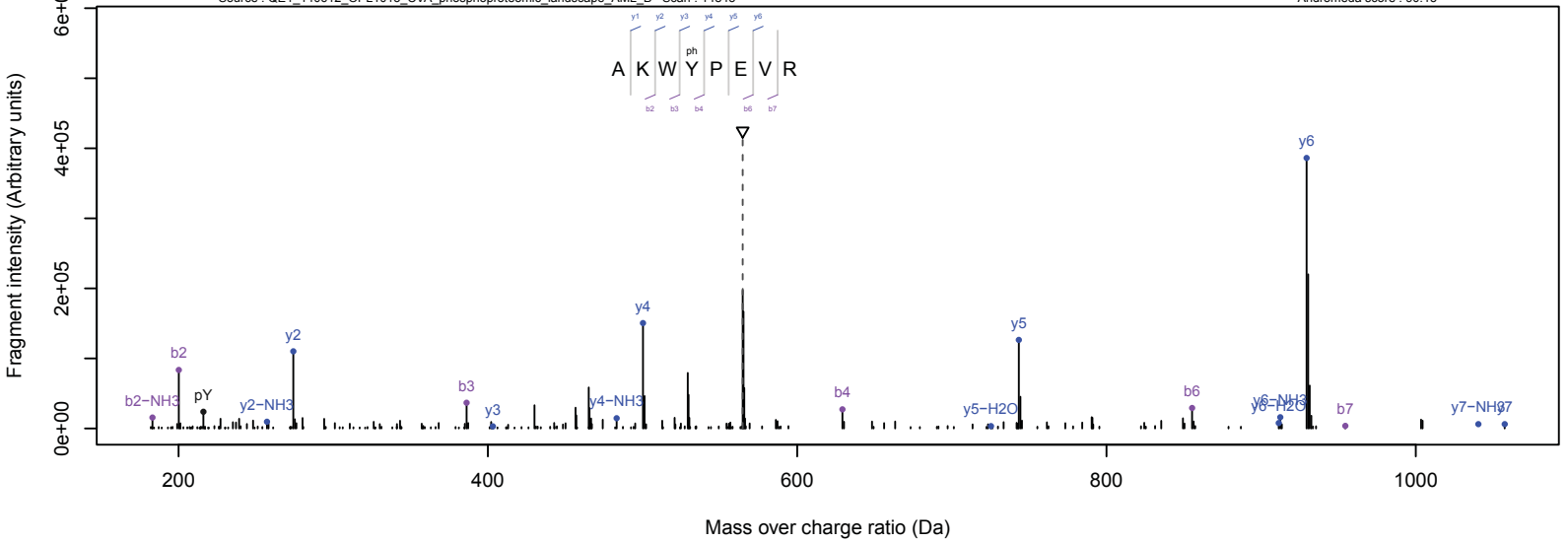
Andromeda score: 90.348



AKWY(ph)PEVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14845

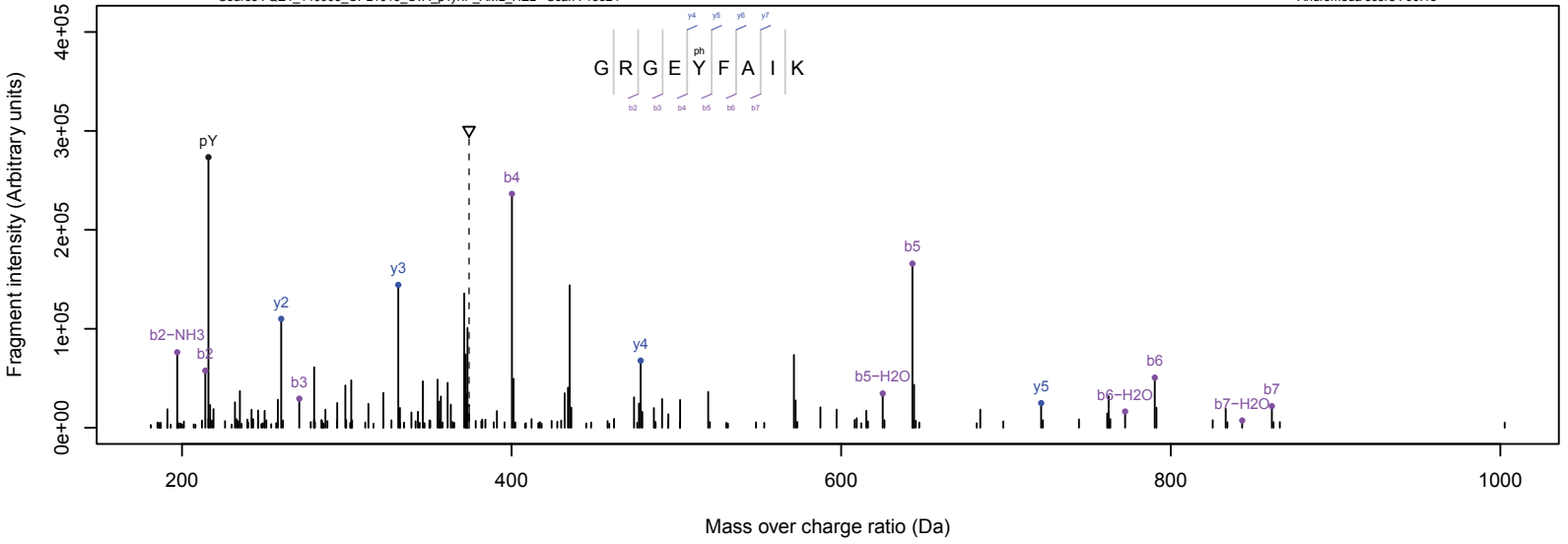
Andromeda score: 90.15



GRGEY(ph)FAIK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 15824

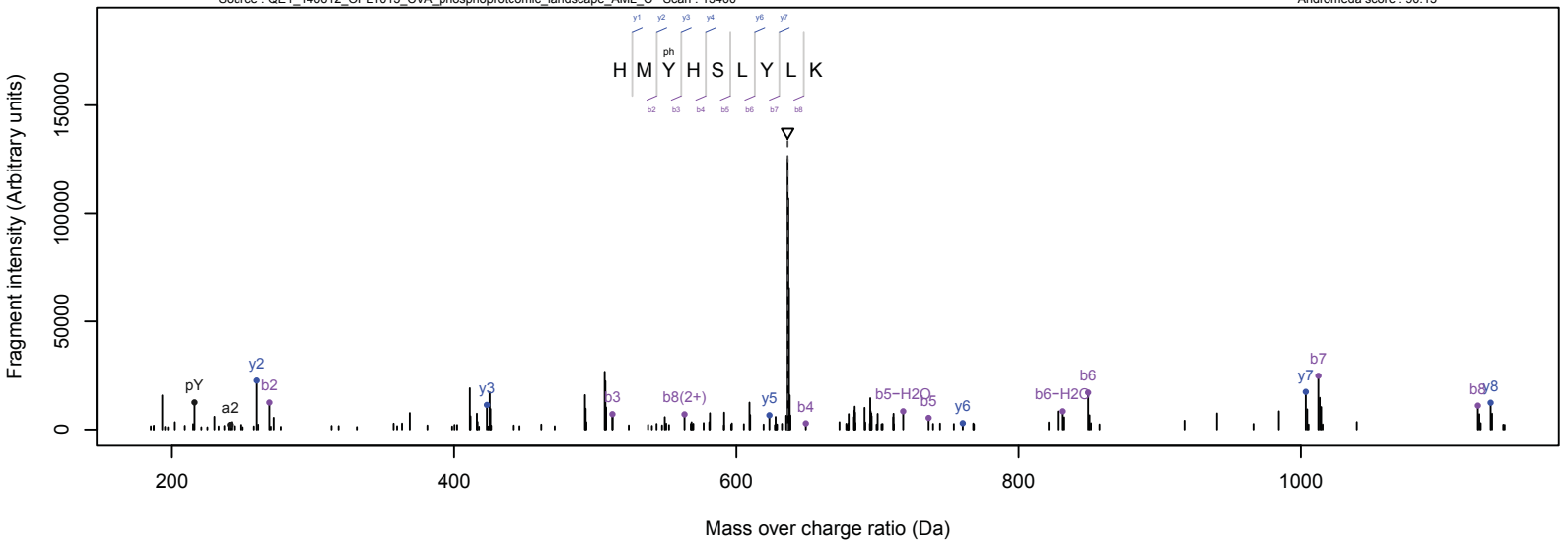
Andromeda score : 90.15



HMY(ph)HSLYLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13400

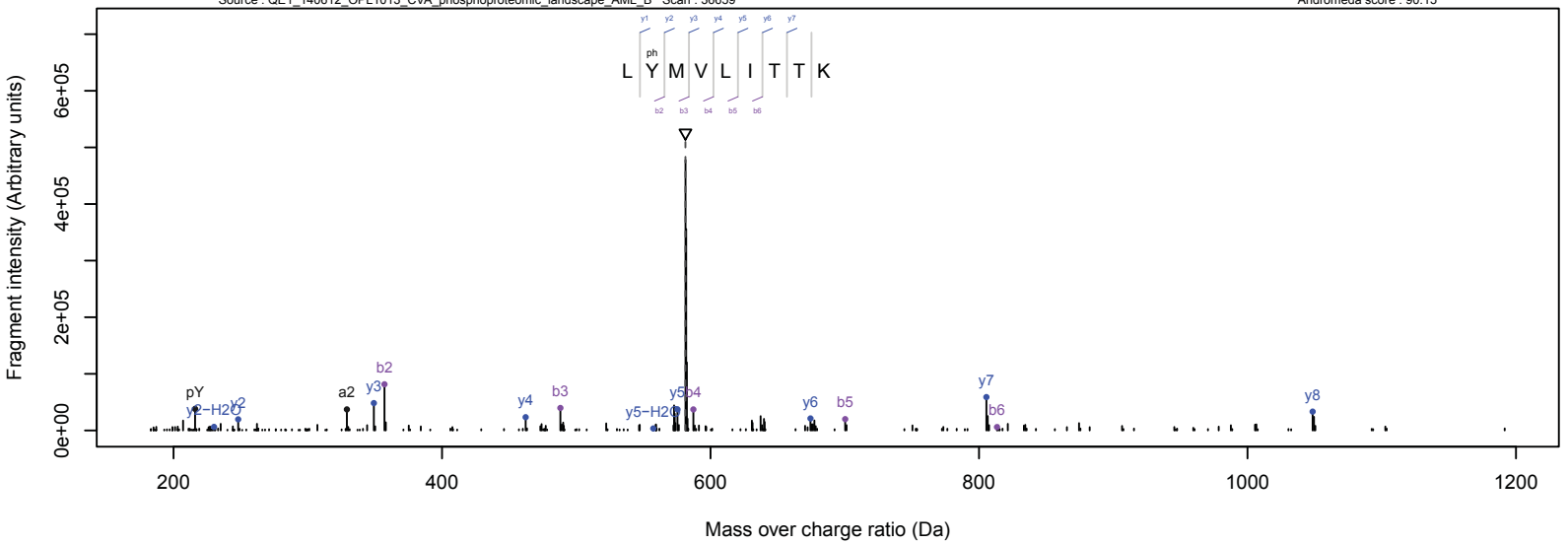
Andromeda score : 90.15



LY(ph)MVLITTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36659

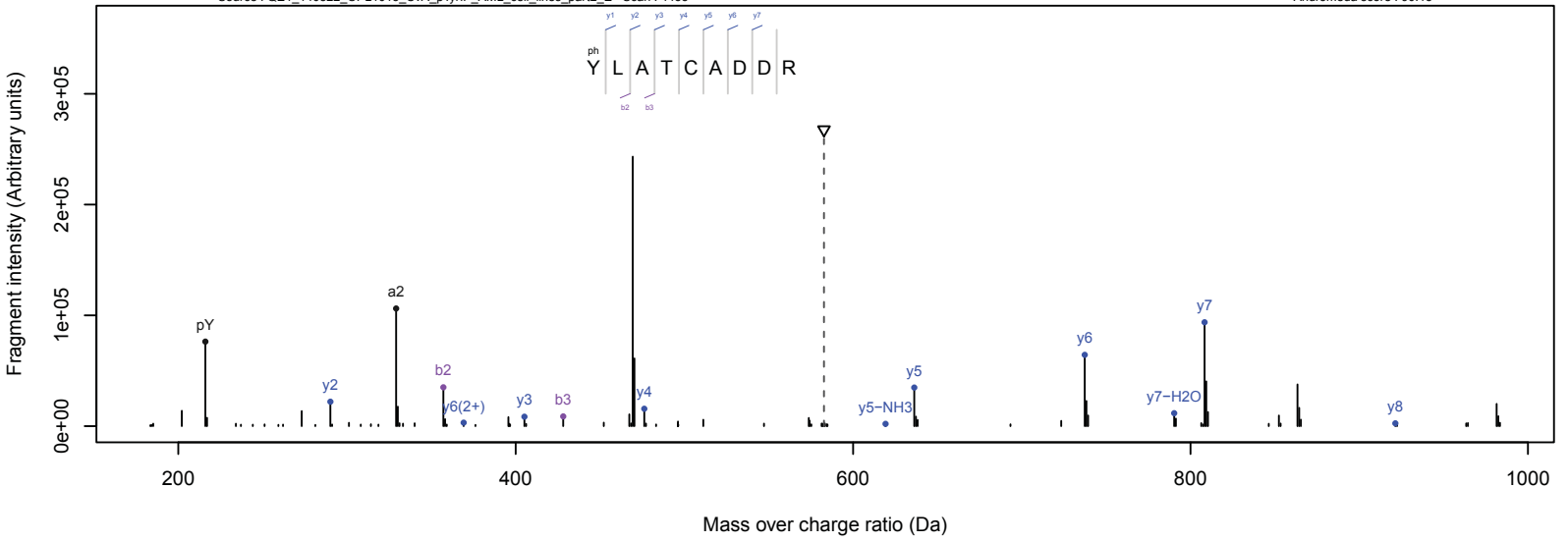
Andromeda score : 90.15



Y(ph)LATCADDR-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 4453

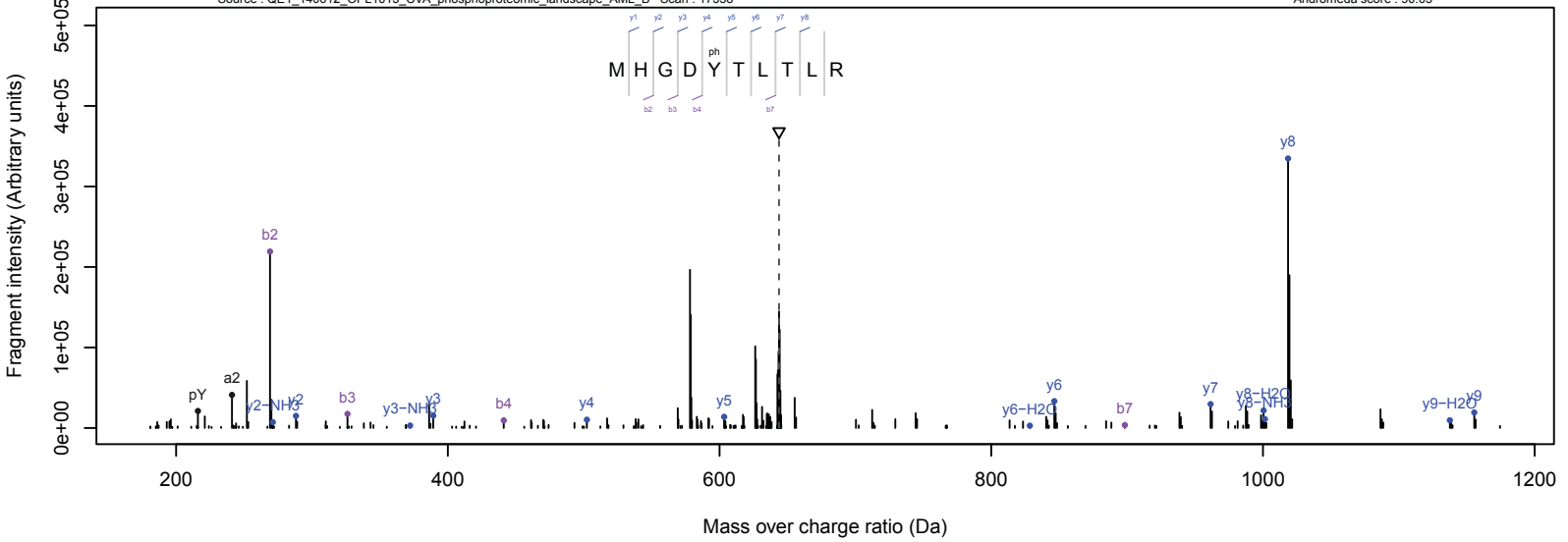
Andromeda score : 90.15



MHGDY(ph)TLTLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17358

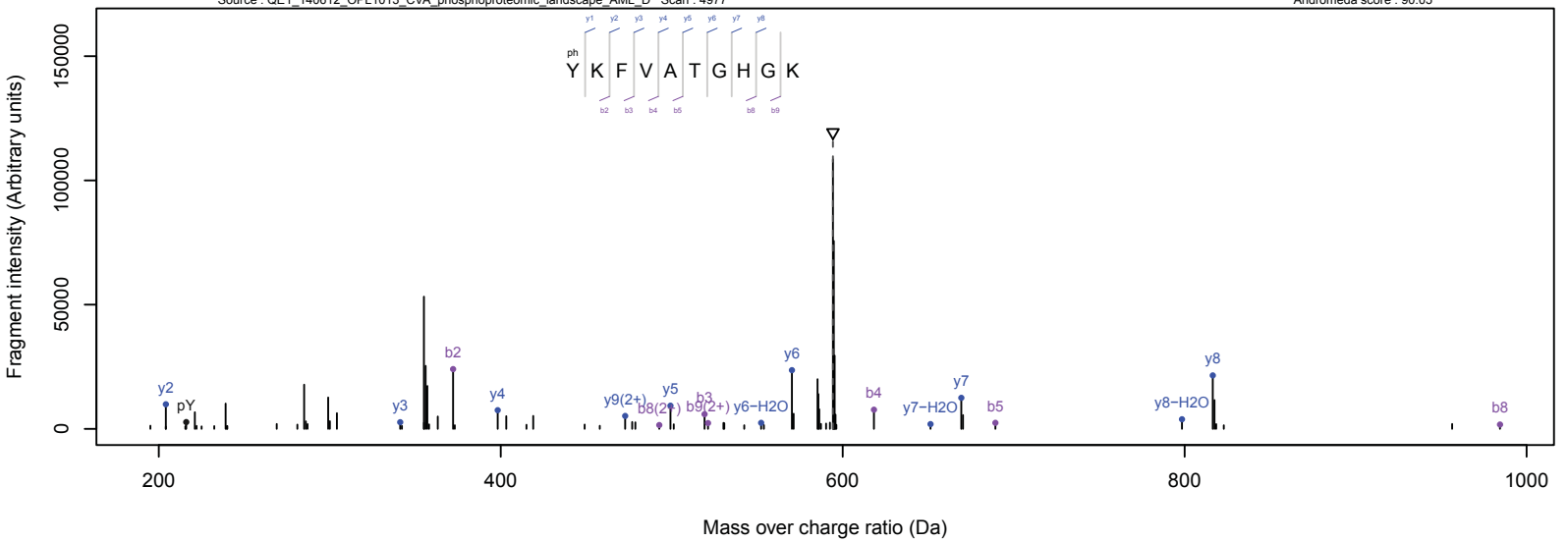
Andromeda score : 90.05



Y(ph)KFVATGHGK-2+

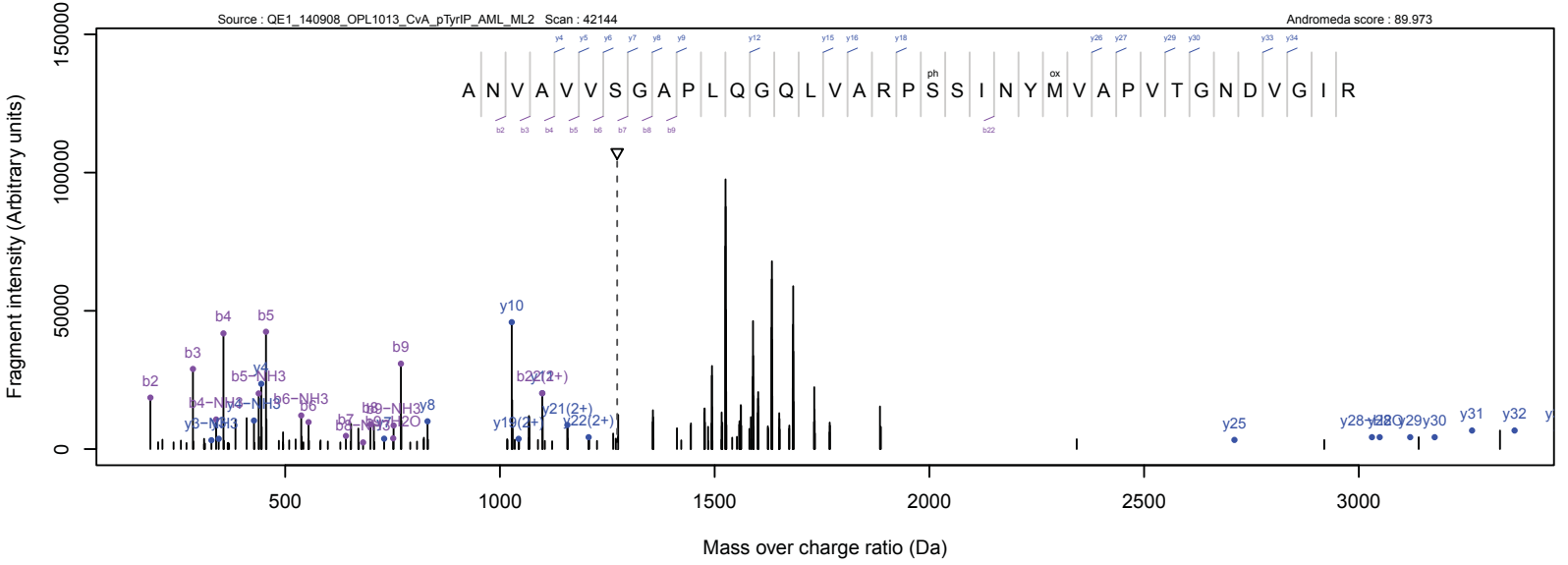
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 4977

Andromeda score : 90.05

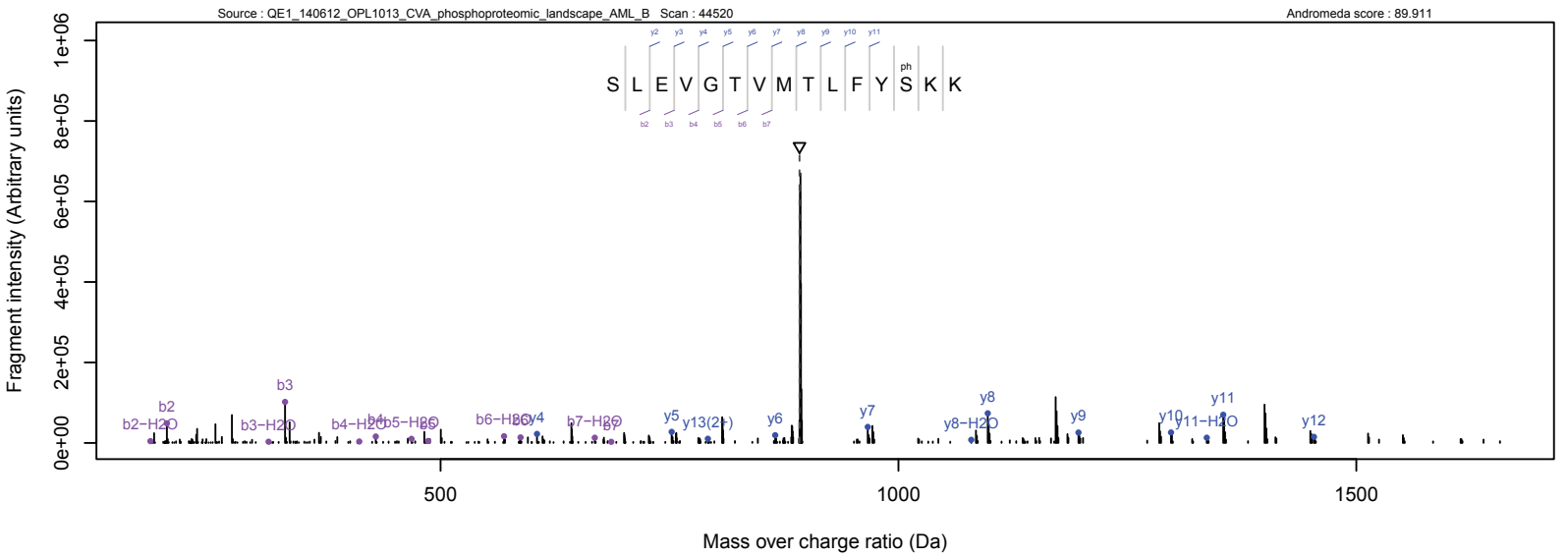




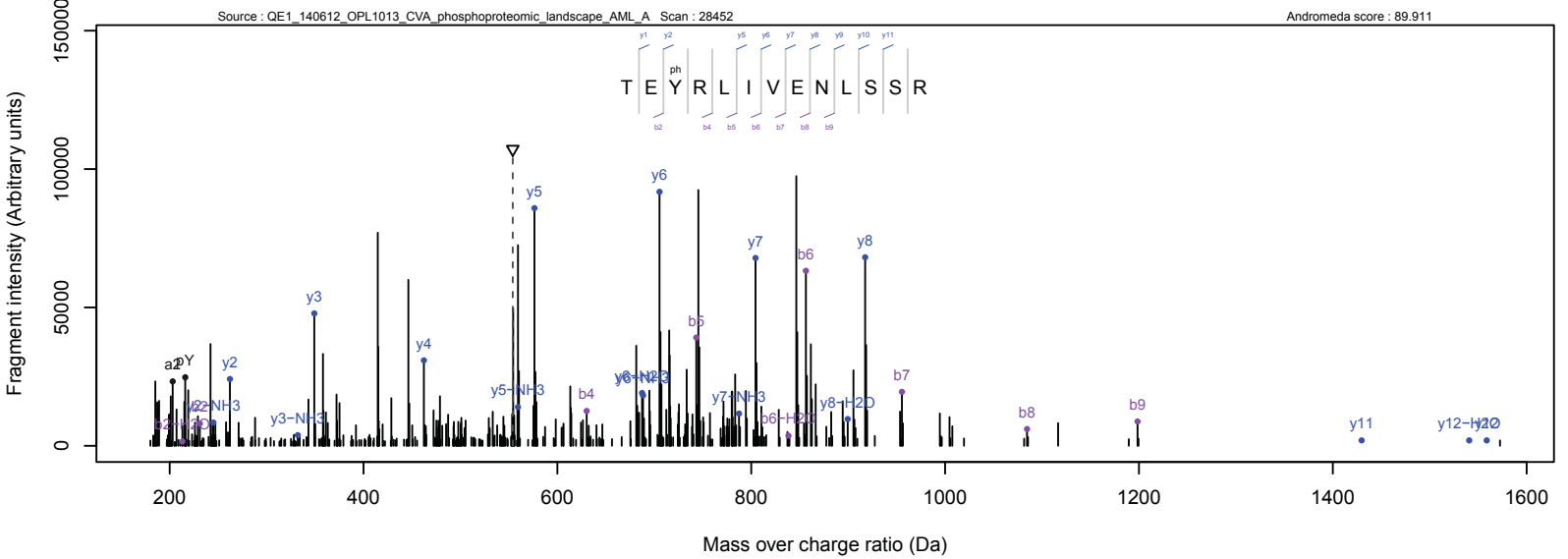
ANVAVVSGAPLQGQLVARPS(ph)SINYM(ox)VAPVTGNDVGIR-3+



SLEVGTVMTLFY(ph)SKK-2+



TEY(ph)RLIVENLSSR-3+

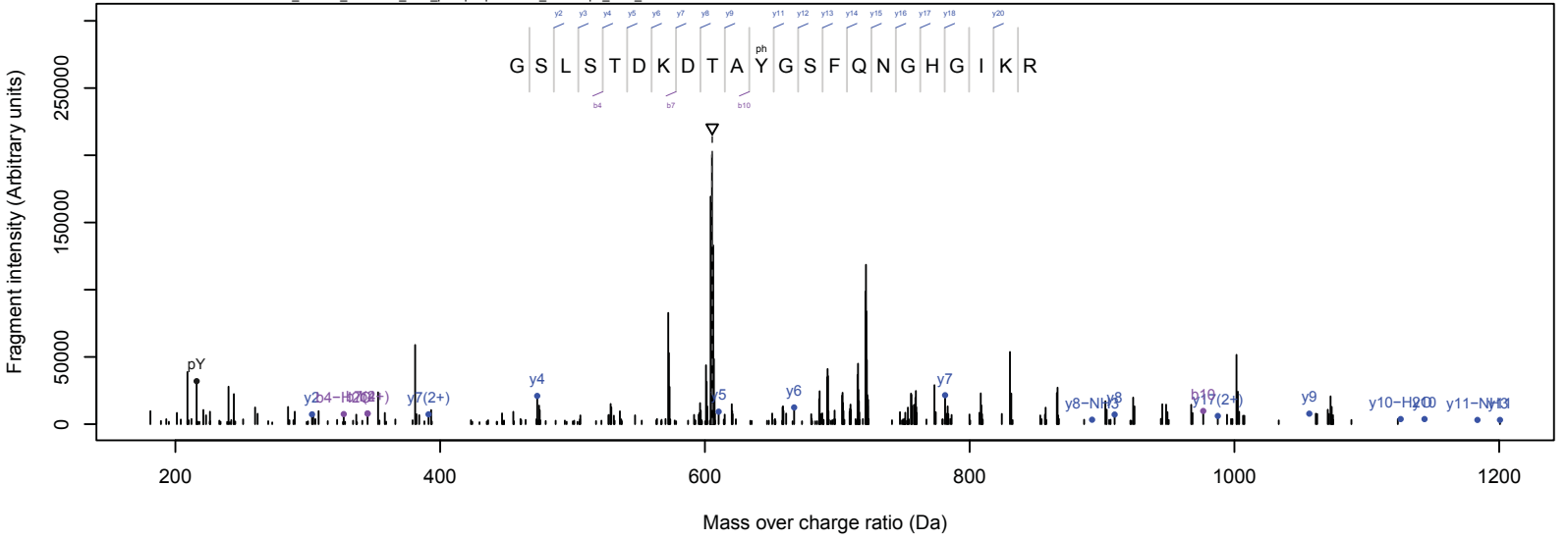




GSLSTDKDTAY(ph)GSFQNGHGIKR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11806

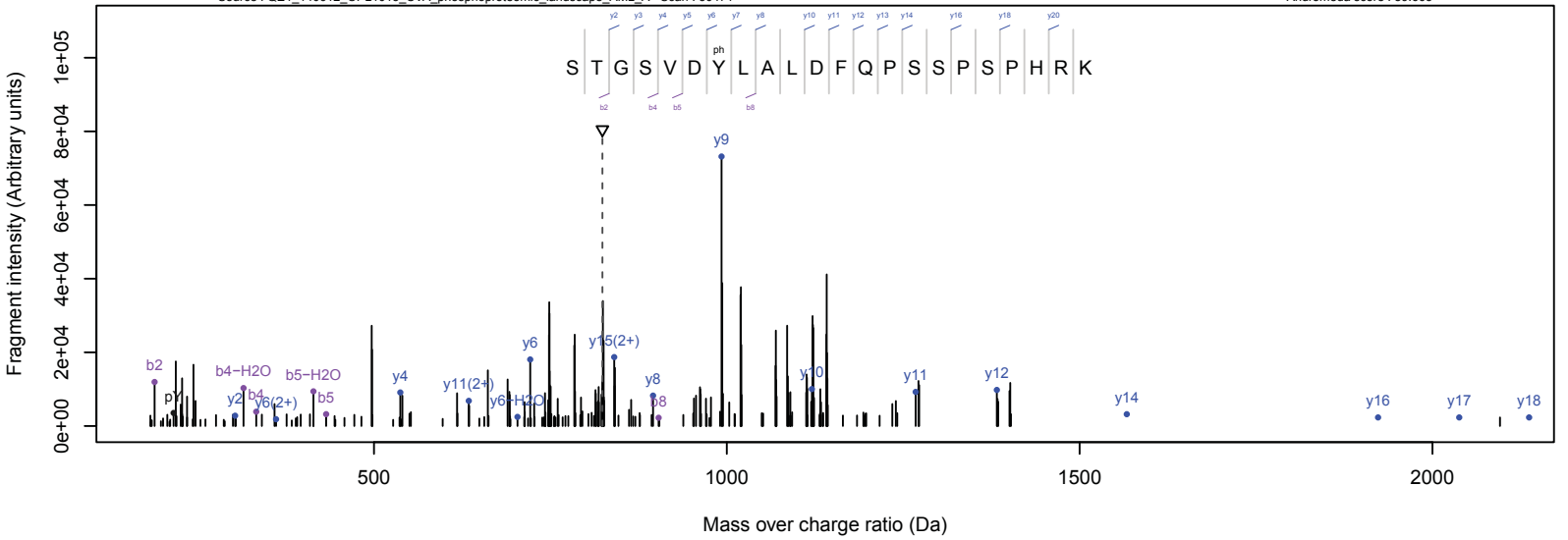
Andromeda score : 89.663



STGSVDY(ph)LALDFQPSSPSPHRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 30471

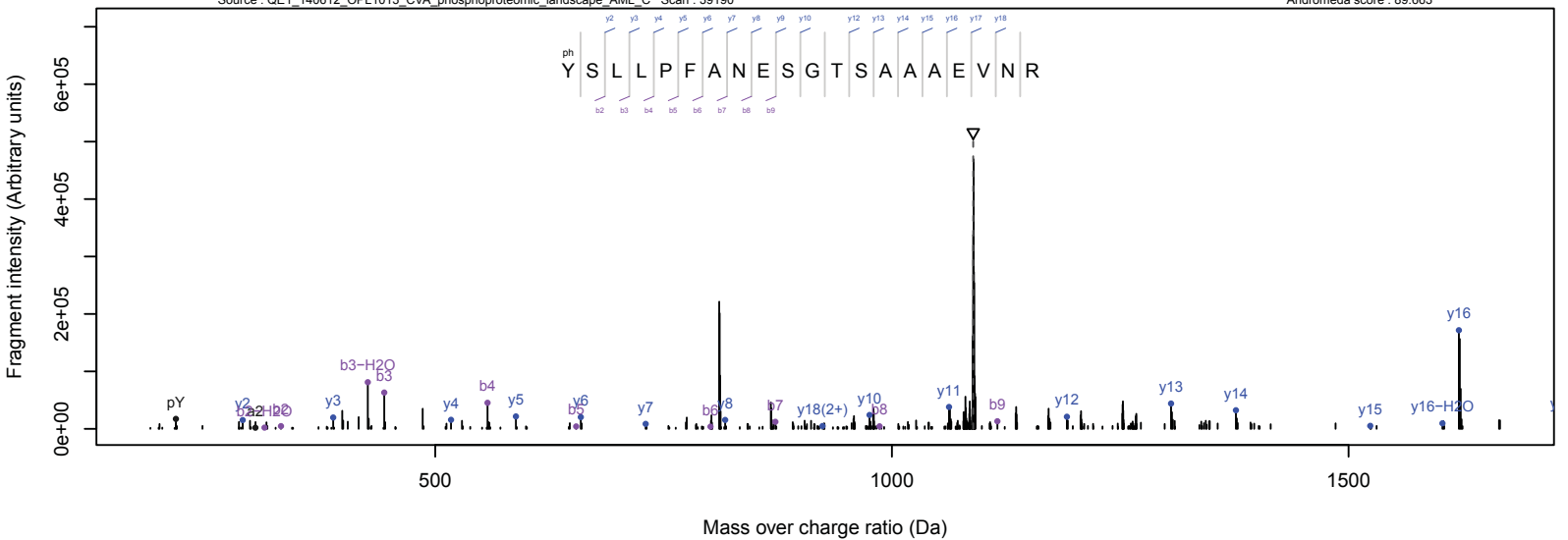
Andromeda score : 89.663



Y(ph)SLLPFANESGTSAAAEVNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39190

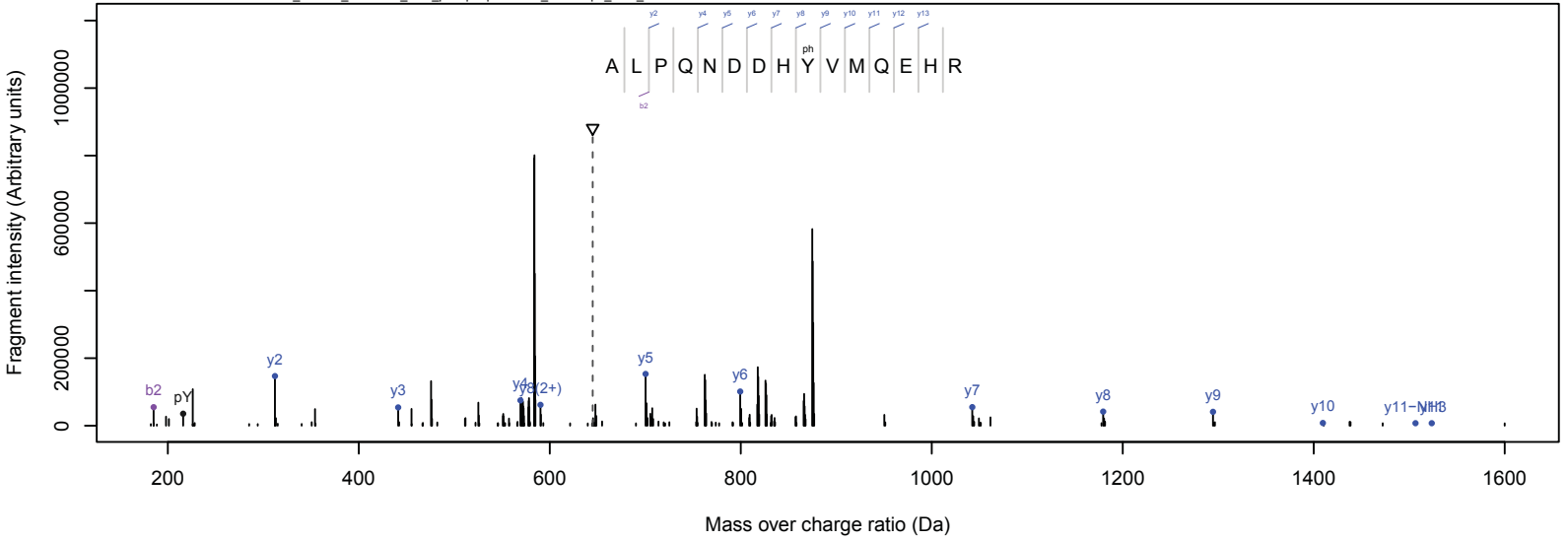
Andromeda score : 89.663



ALPQDDHY(ph)VMQEHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12025

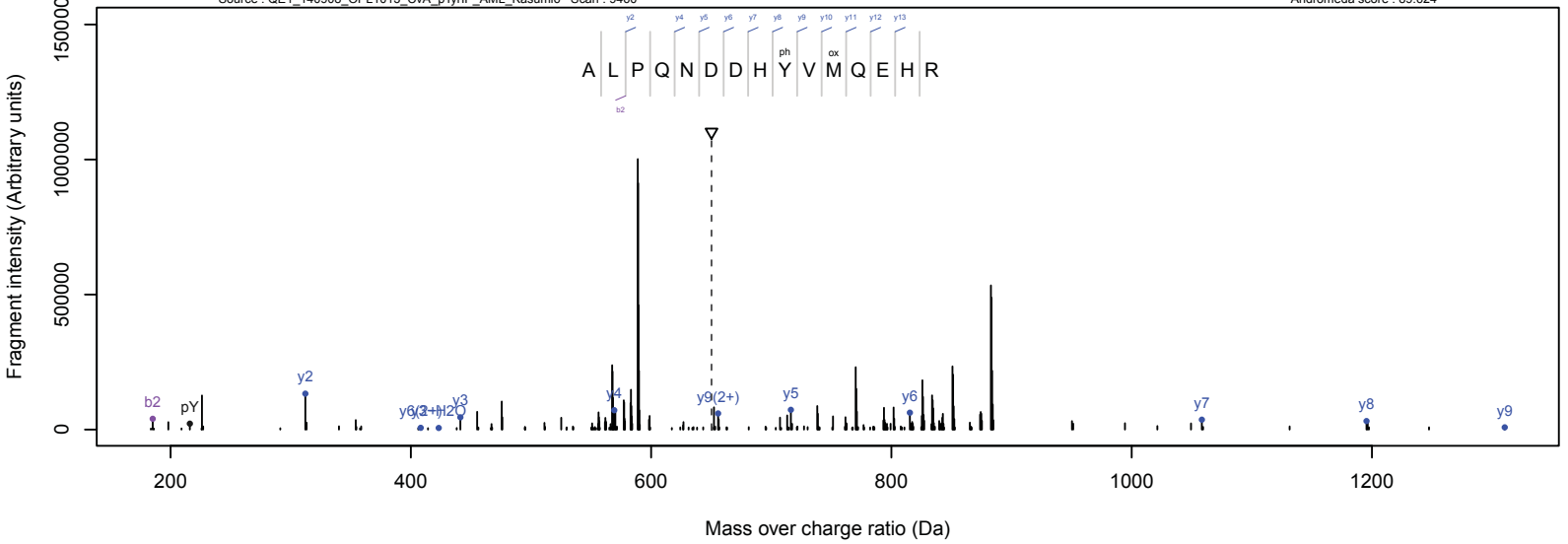
Andromeda score : 89.624



ALPQDDHY(ph)VM(ox)QEHR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 9460

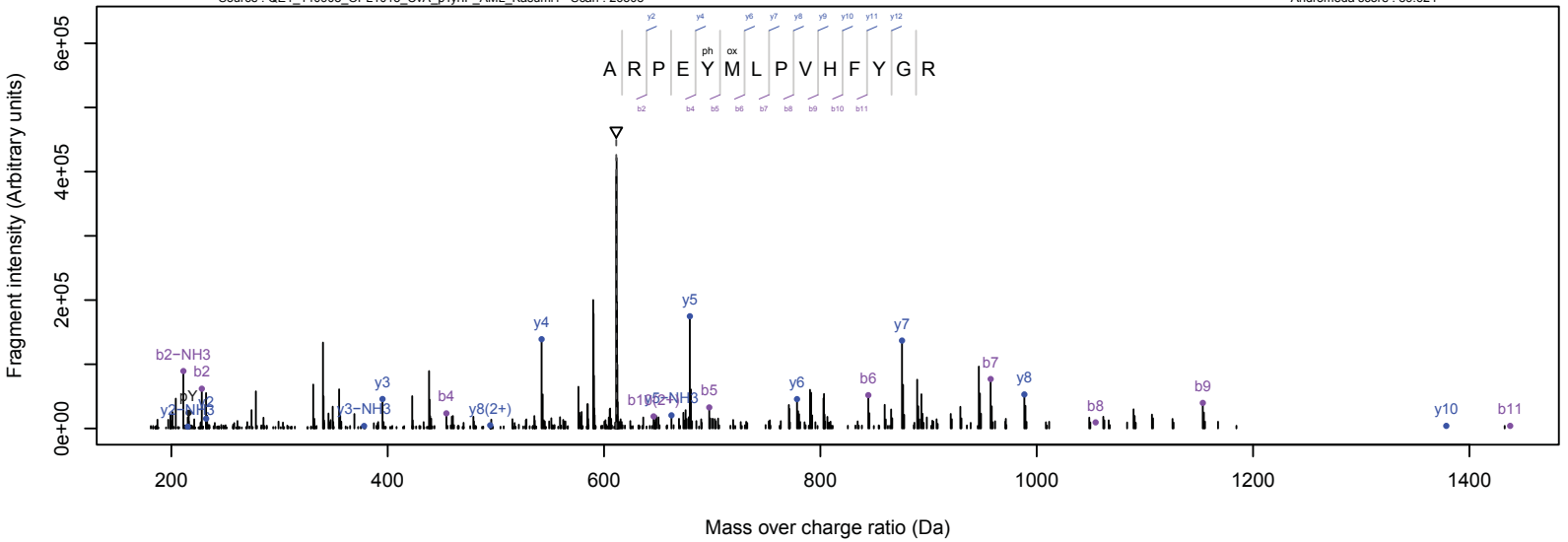
Andromeda score : 89.624



ARPEY(ph)M(ox)LPVHFYGR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 26308

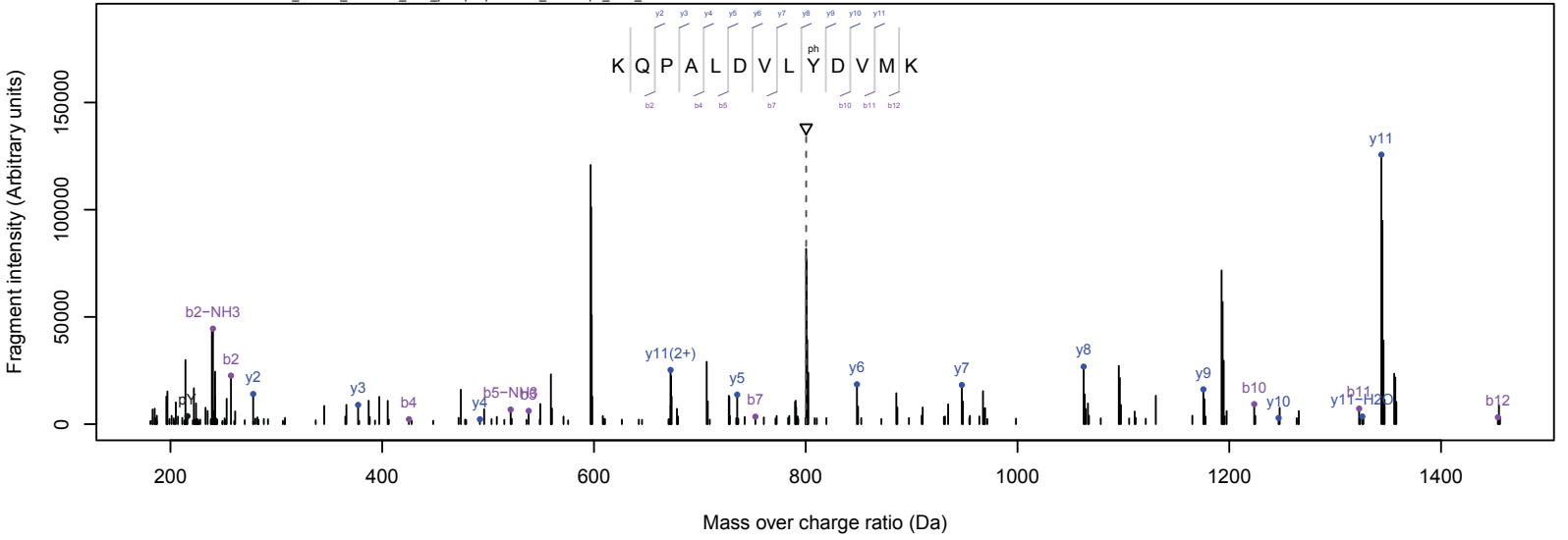
Andromeda score : 89.624



KQPALDVL(ph)DVMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 43654

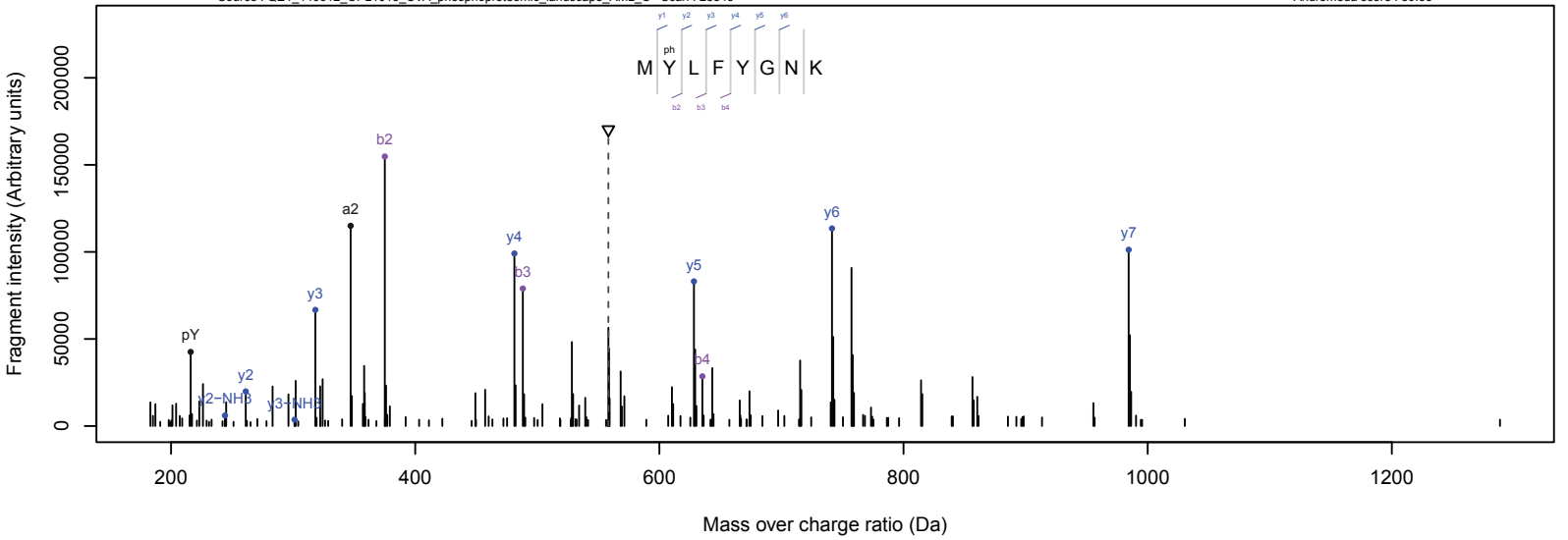
Andromeda score : 89.624



MY(ph)LFYGNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28845

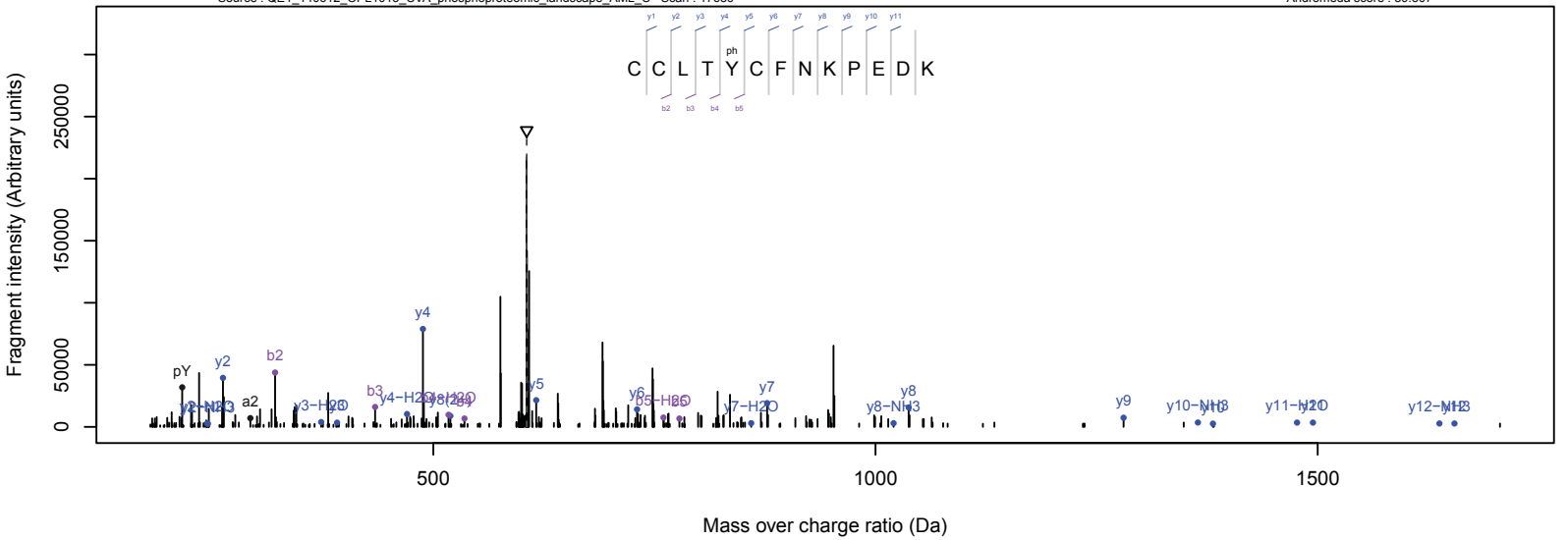
Andromeda score : 89.55



CCLTY(ph)CFNKPEDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17689

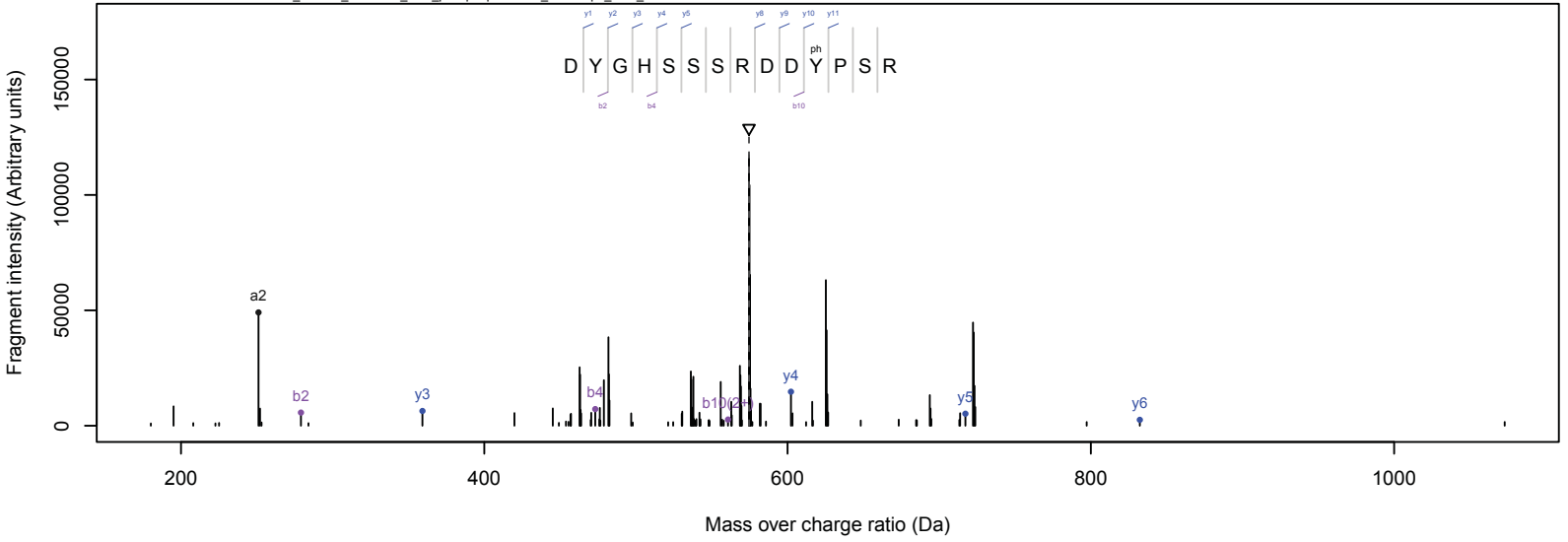
Andromeda score : 89.507



DYGHSSRDDY(ph)PSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3583

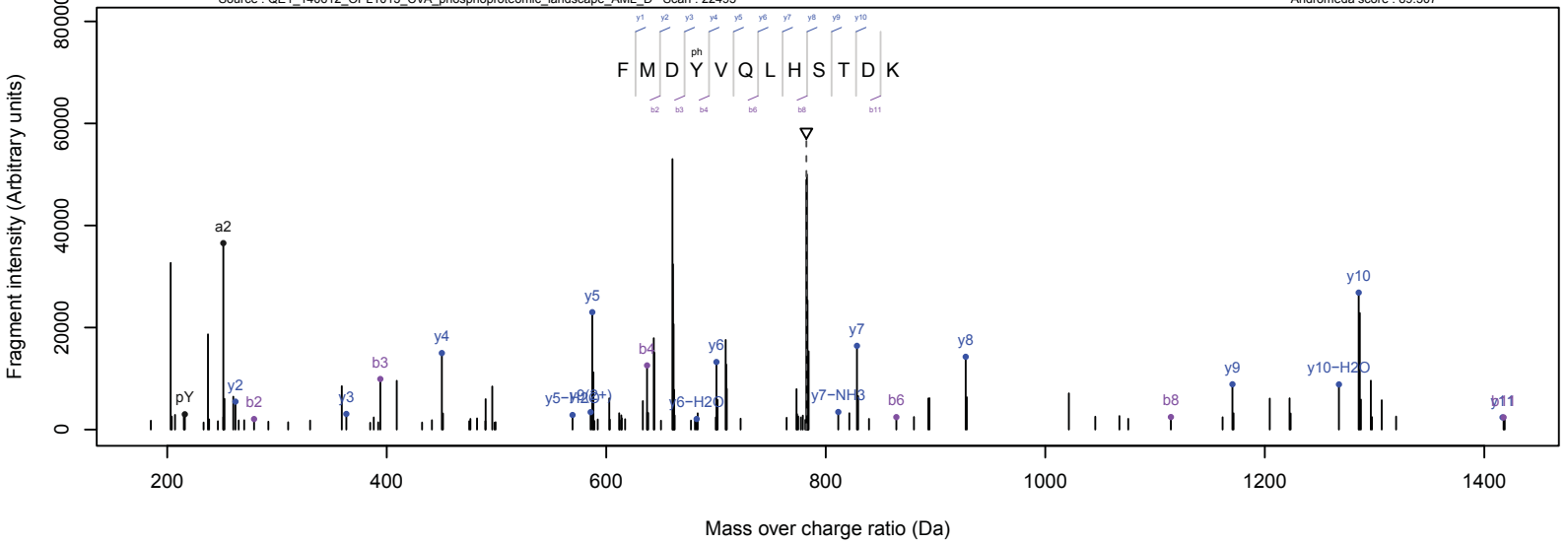
Andromeda score : 89.507



FMDY(ph)VQLHSTDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22495

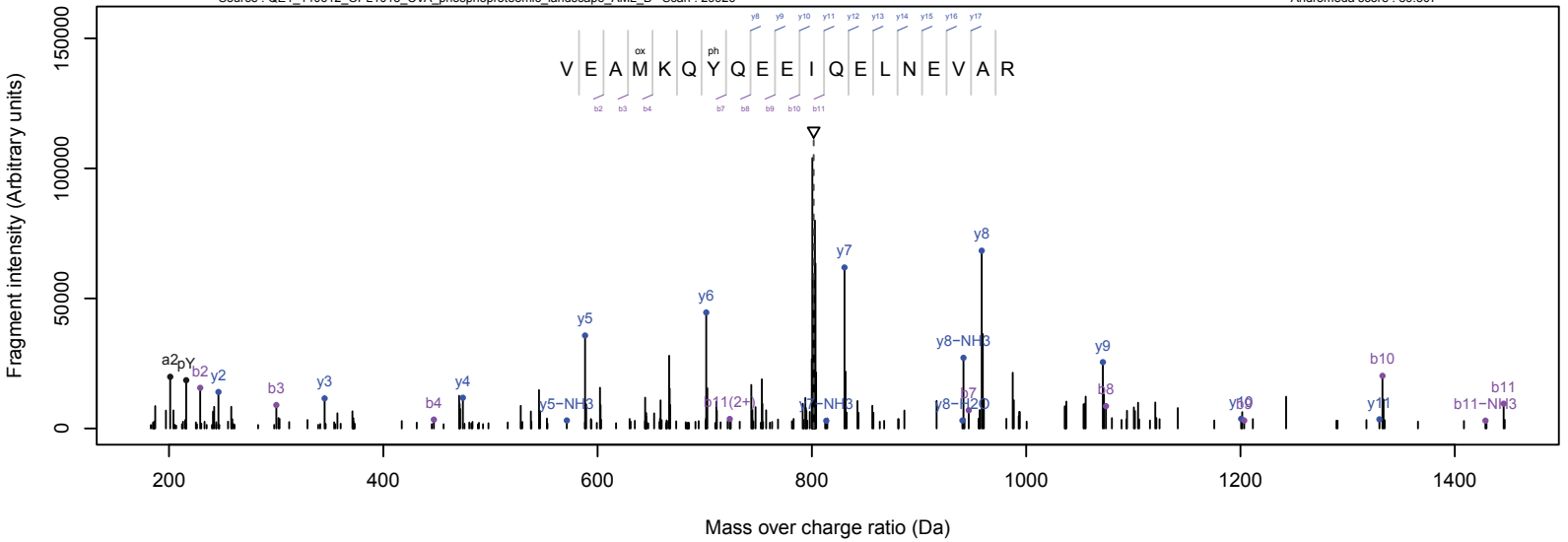
Andromeda score : 89.507



VEAM(ox)KQY(ph)QEEIQELNEVAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29926

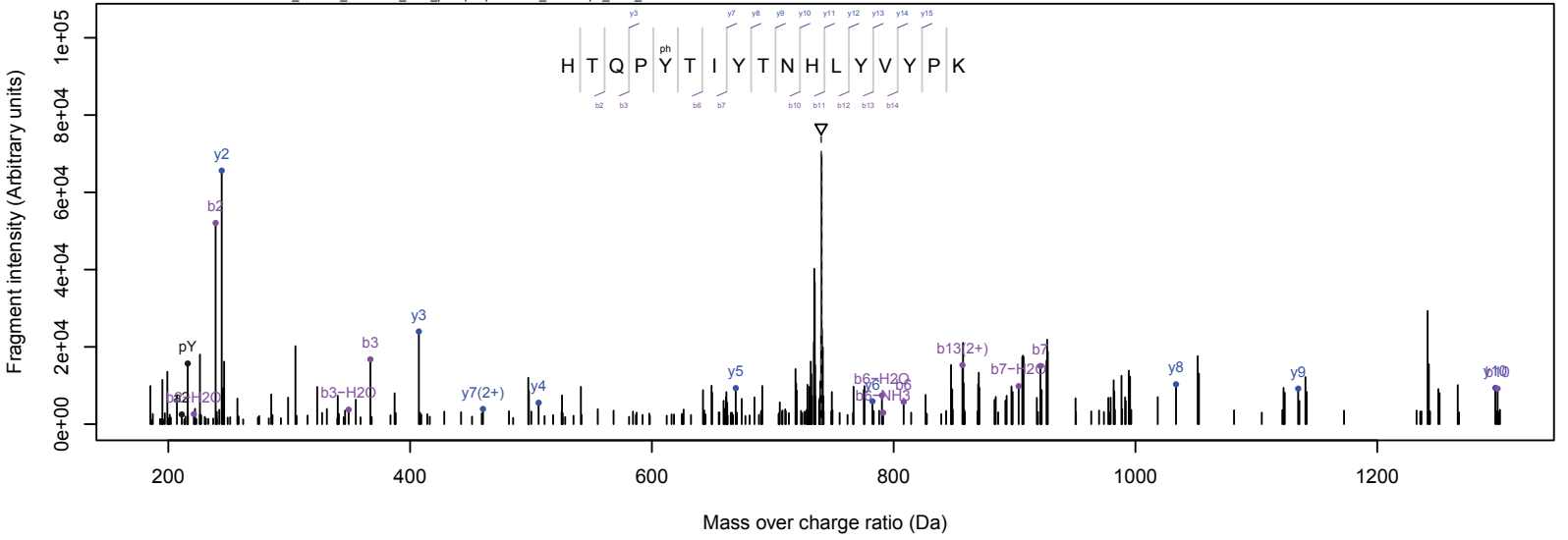
Andromeda score : 89.507



HTQPY(ph)TIYTNHLYVYPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26450

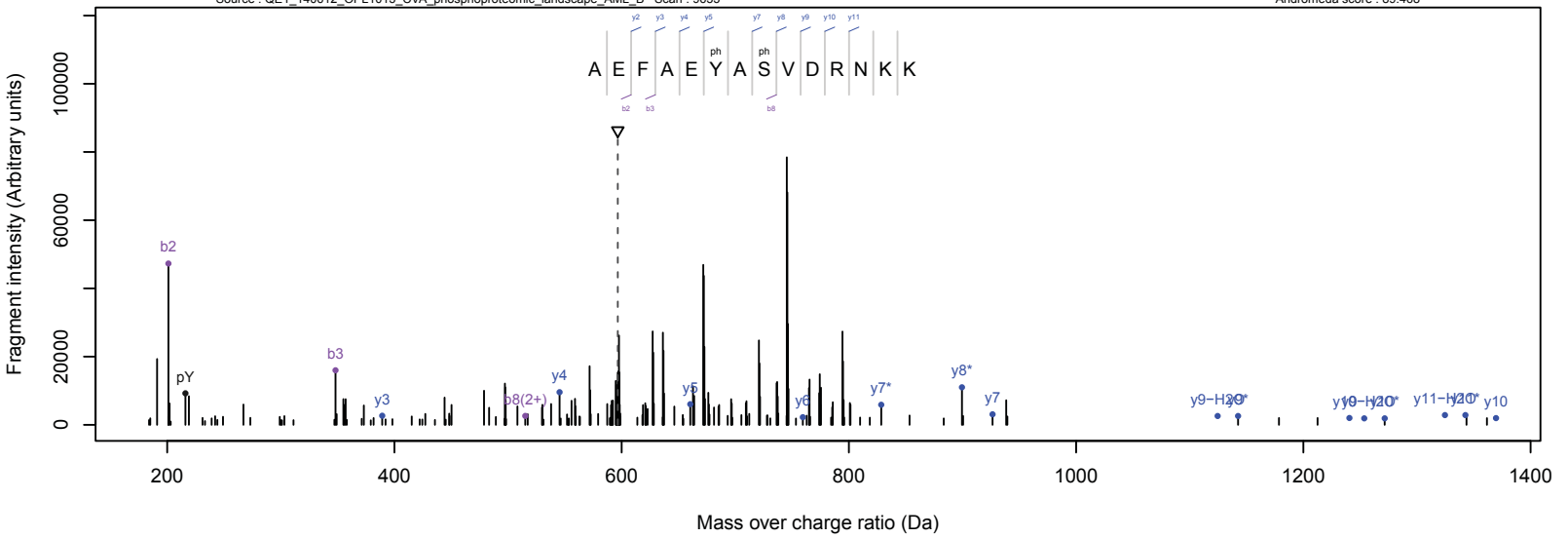
Andromeda score : 89.502



AEFAEY(ph)AS(ph)VDRNKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9633

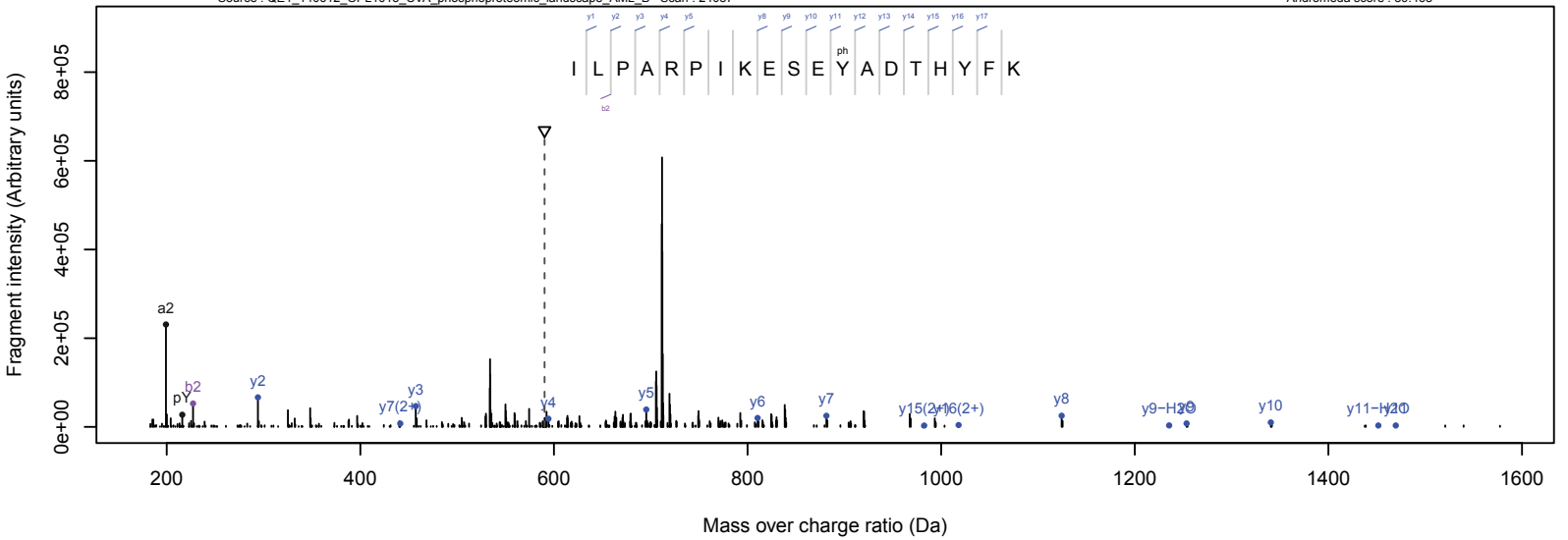
Andromeda score : 89.468



ILPARPIKESEY(ph)ADTHYFK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24087

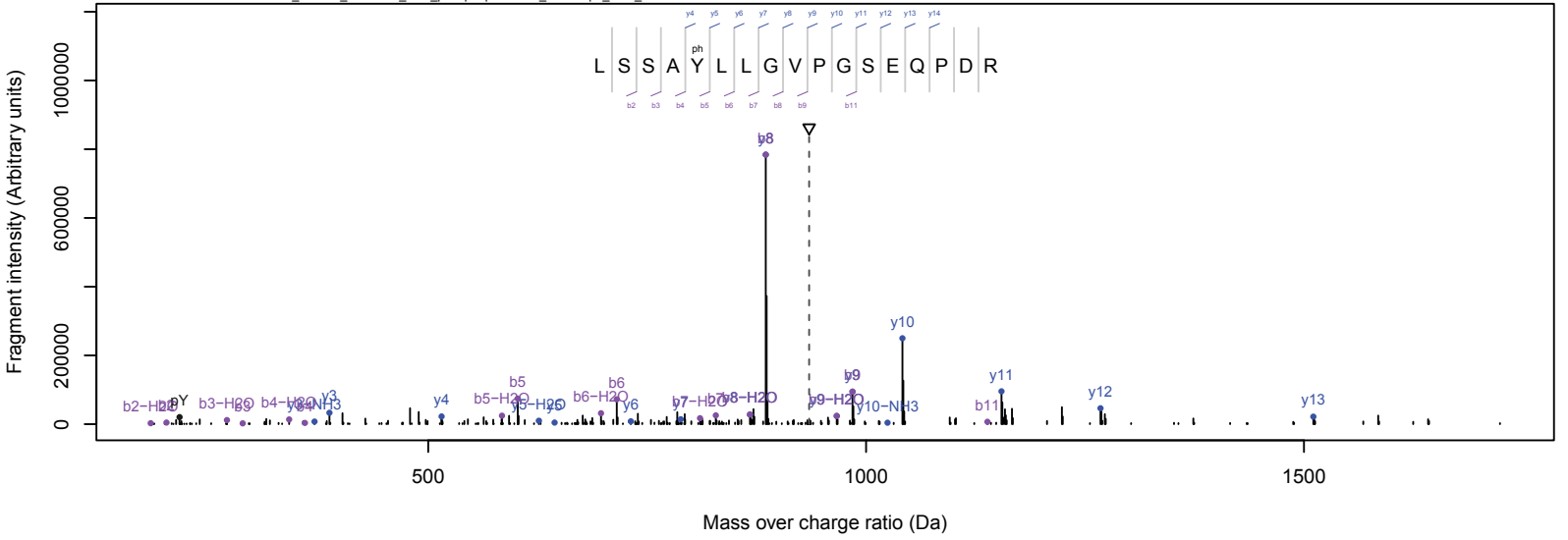
Andromeda score : 89.468



LSSAY(ph)LLGVPGEQPD<sup>2+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33916

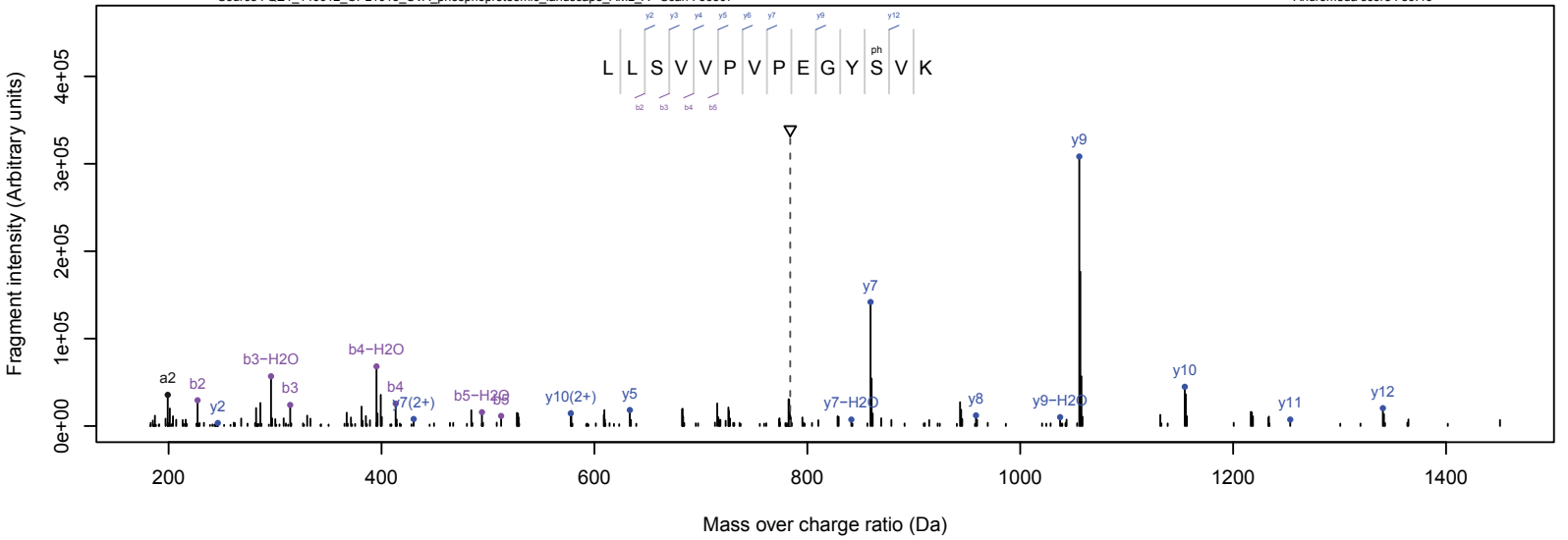
Andromeda score : 89.468



LLSVVPEGYS(ph)VK<sup>2+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36307

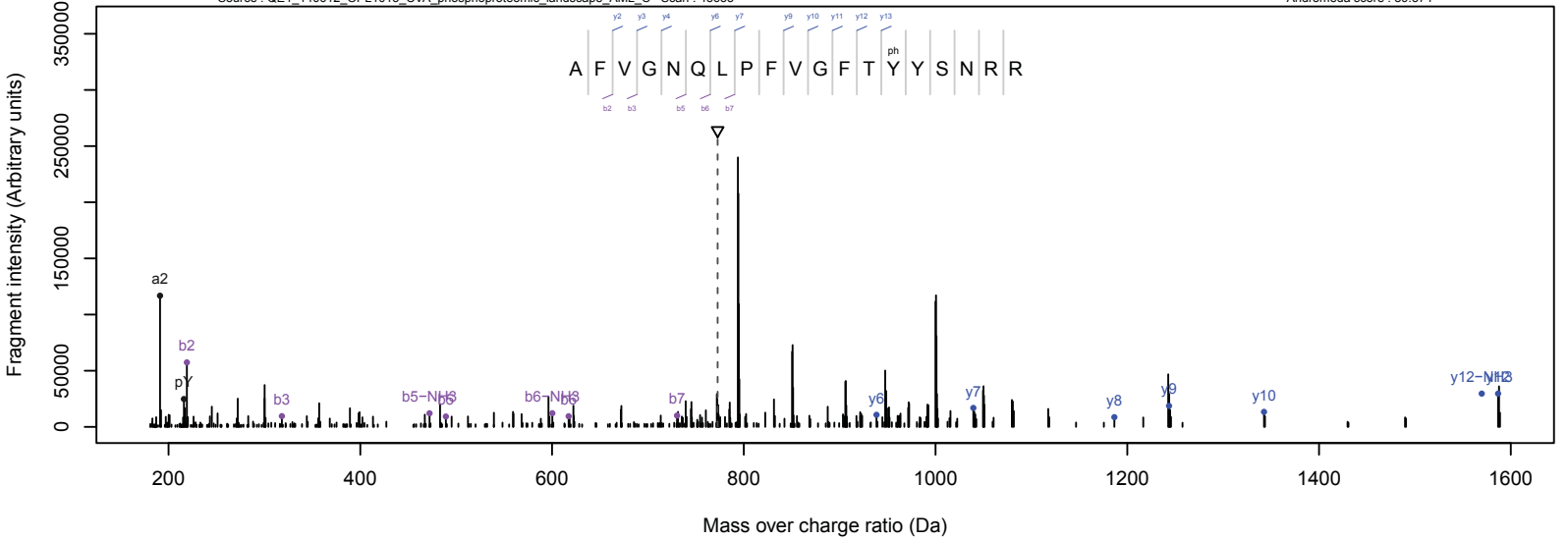
Andromeda score : 89.46



AFVGNQLPFGFTY(ph)YSNRR<sup>3+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45655

Andromeda score : 89.374

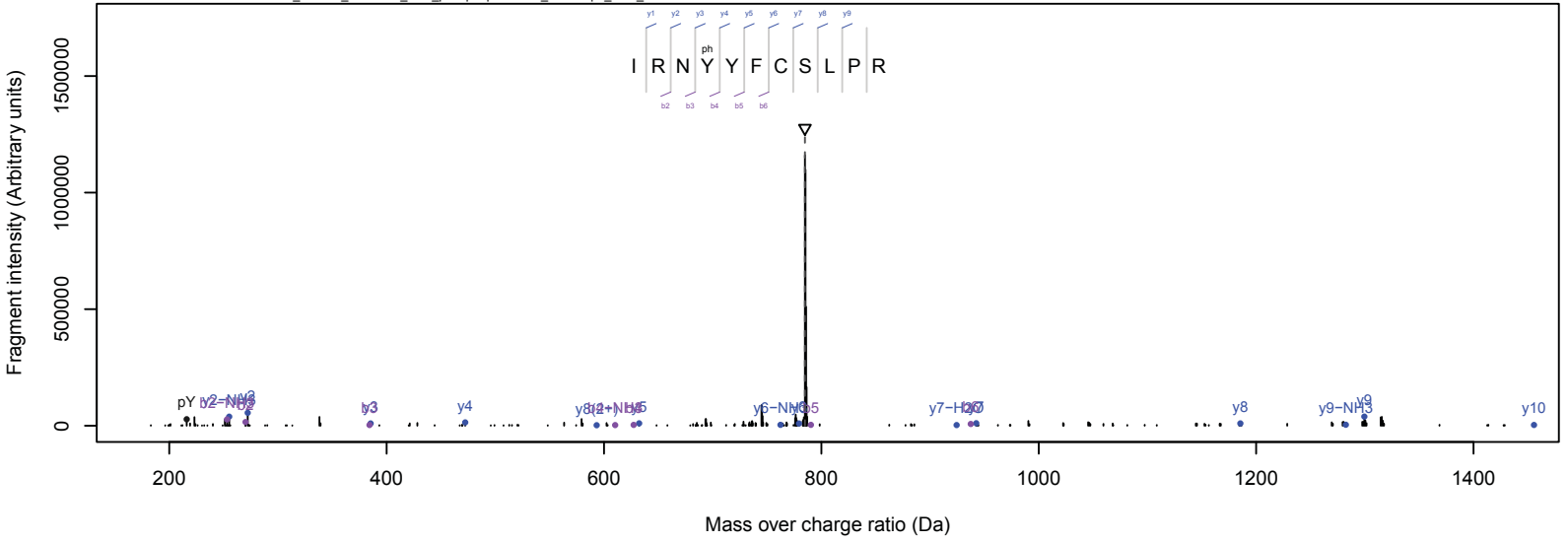




IRNY(ph)YFCSLPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26610

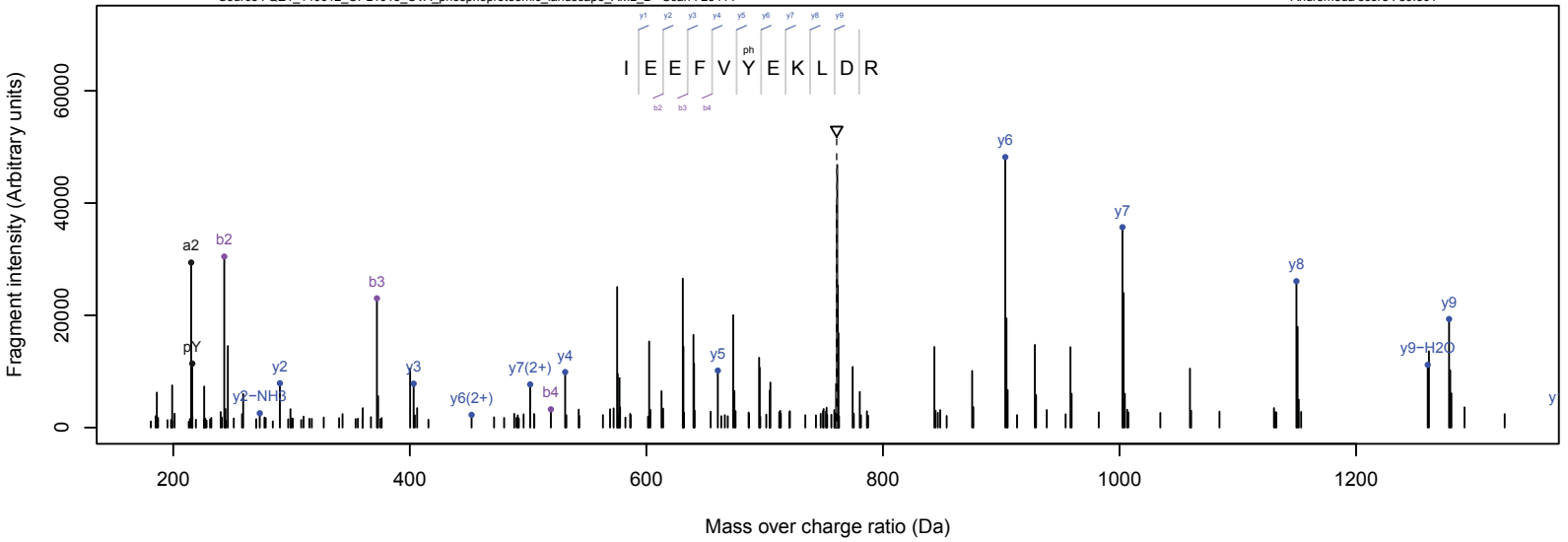
Andromeda score : 89.356



IEEFVY(ph)EKLDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23414

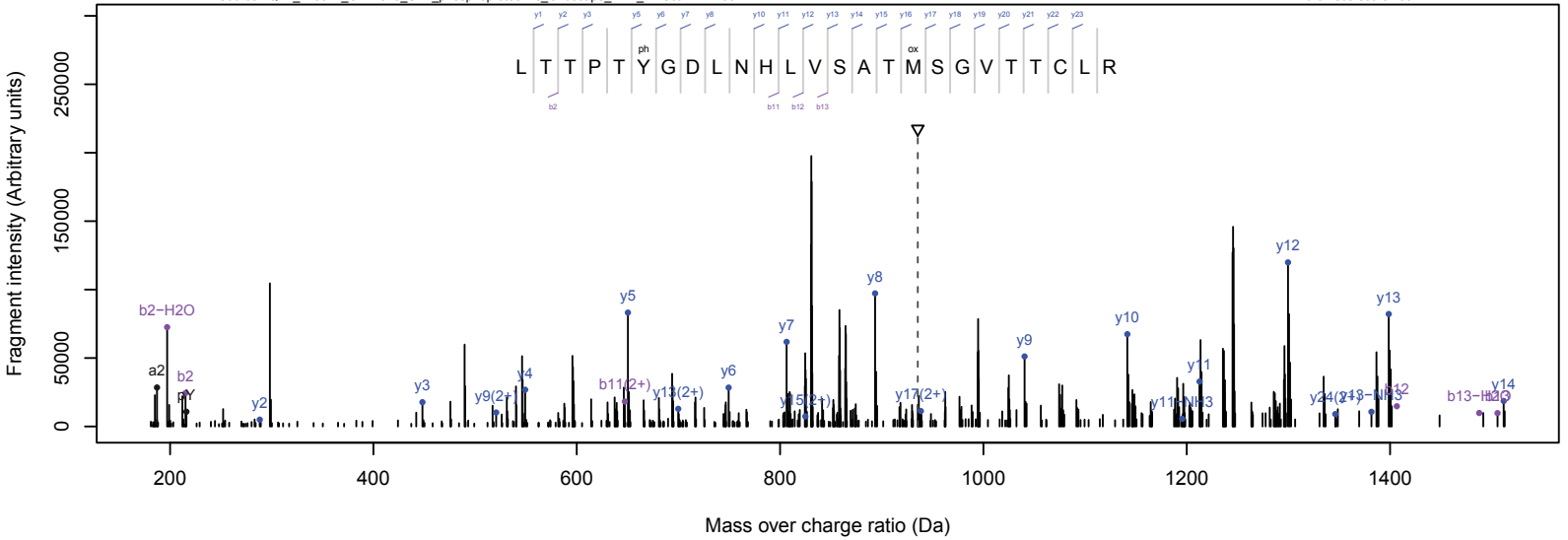
Andromeda score : 89.301



LTTPY(ph)GDLNHLVSATM(ox)SGVTCLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41296

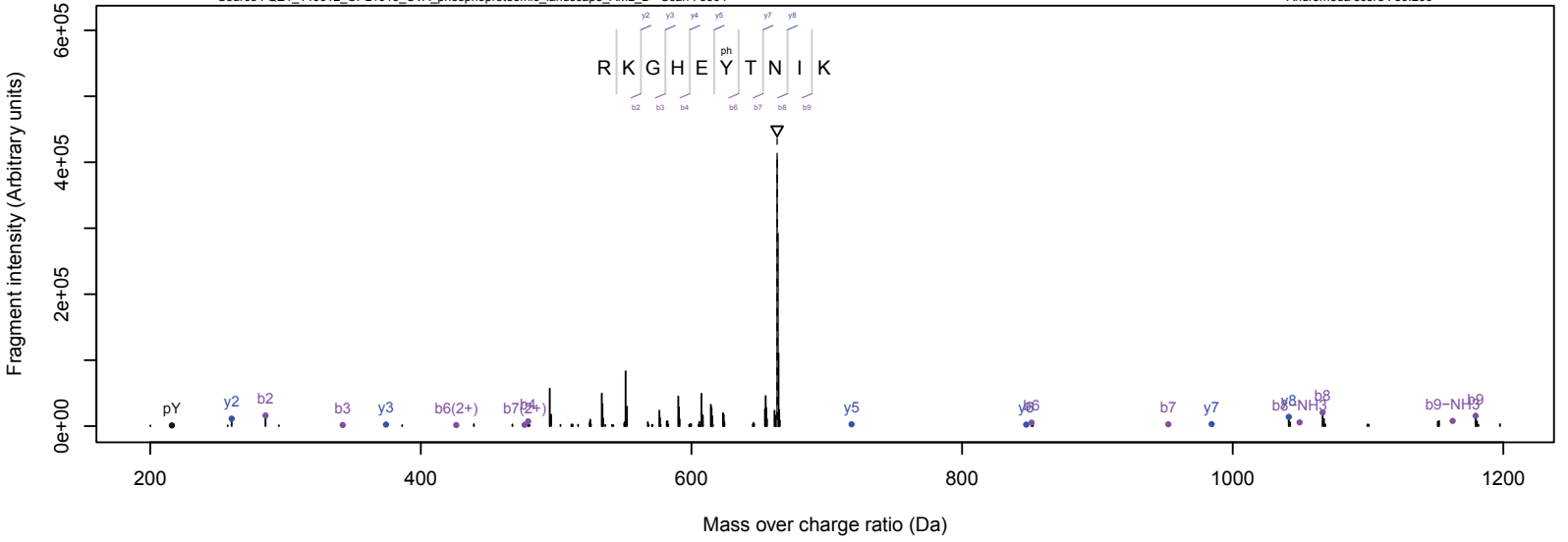
Andromeda score : 89.272



**RGKHEY(ph)TNIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 3504

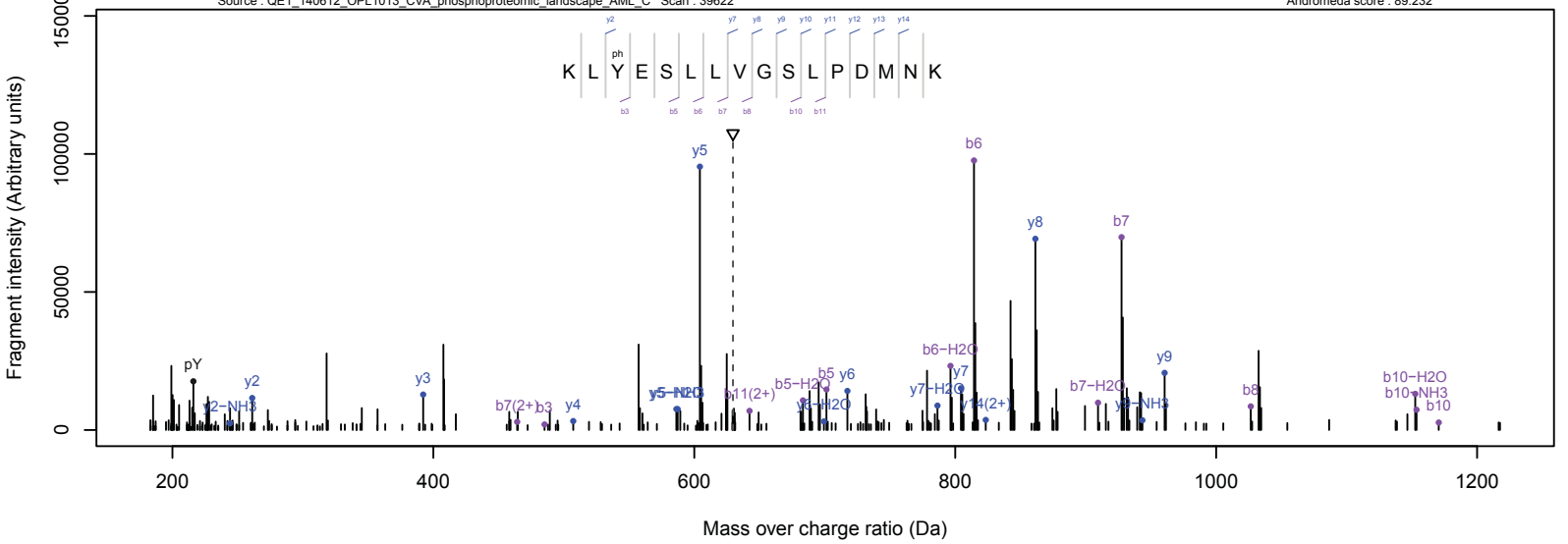
Andromeda score : 89.266



**KLY(ph)ESLLVGS L P D M N K-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39622

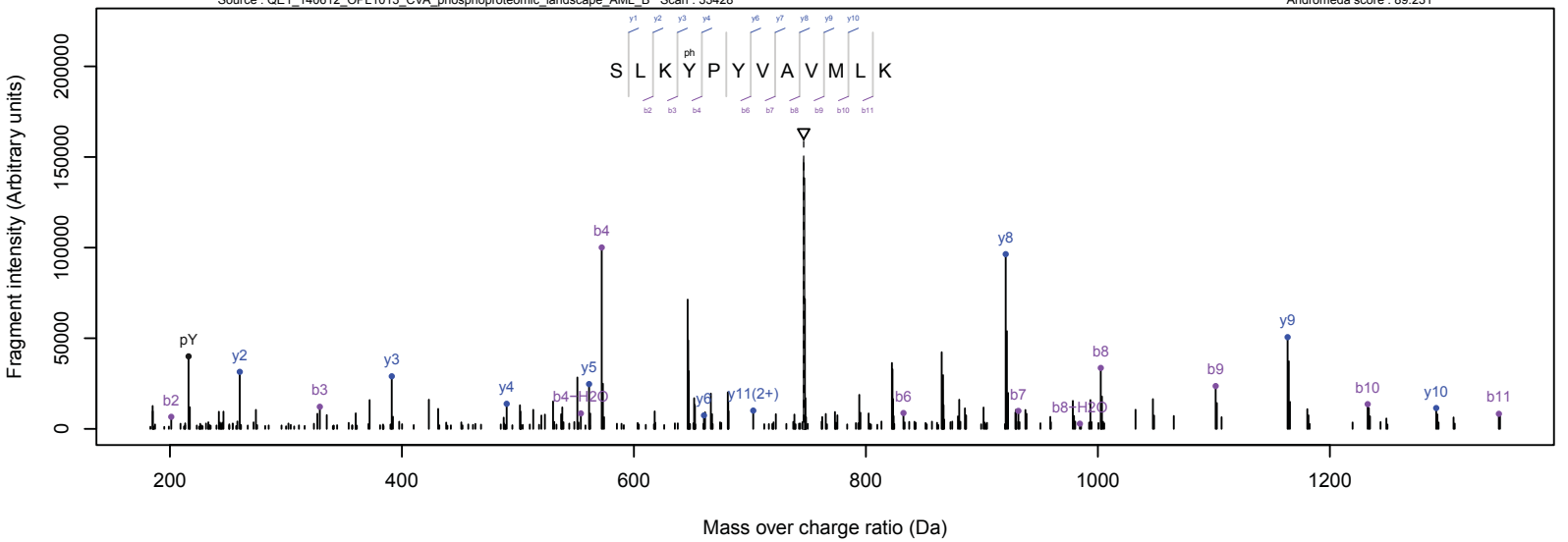
Andromeda score : 89.232



**SLKY(ph)PYVAVMLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33428

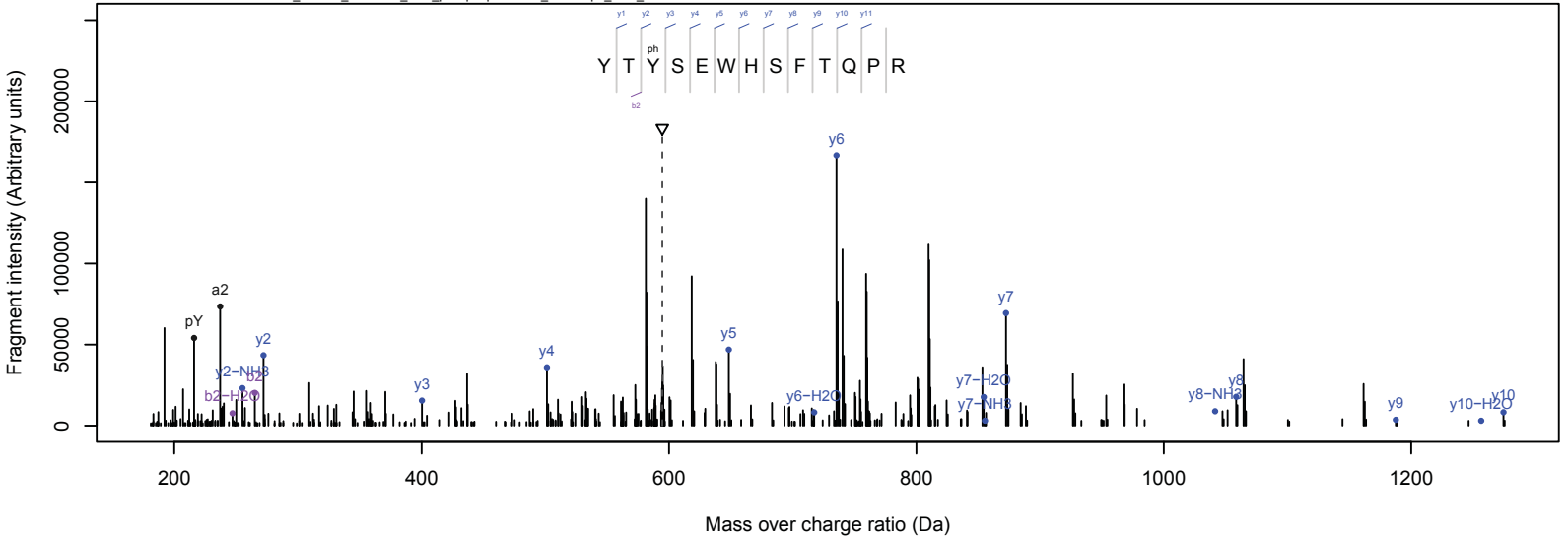
Andromeda score : 89.231



YTY(ph)SEWHSFTQPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23975

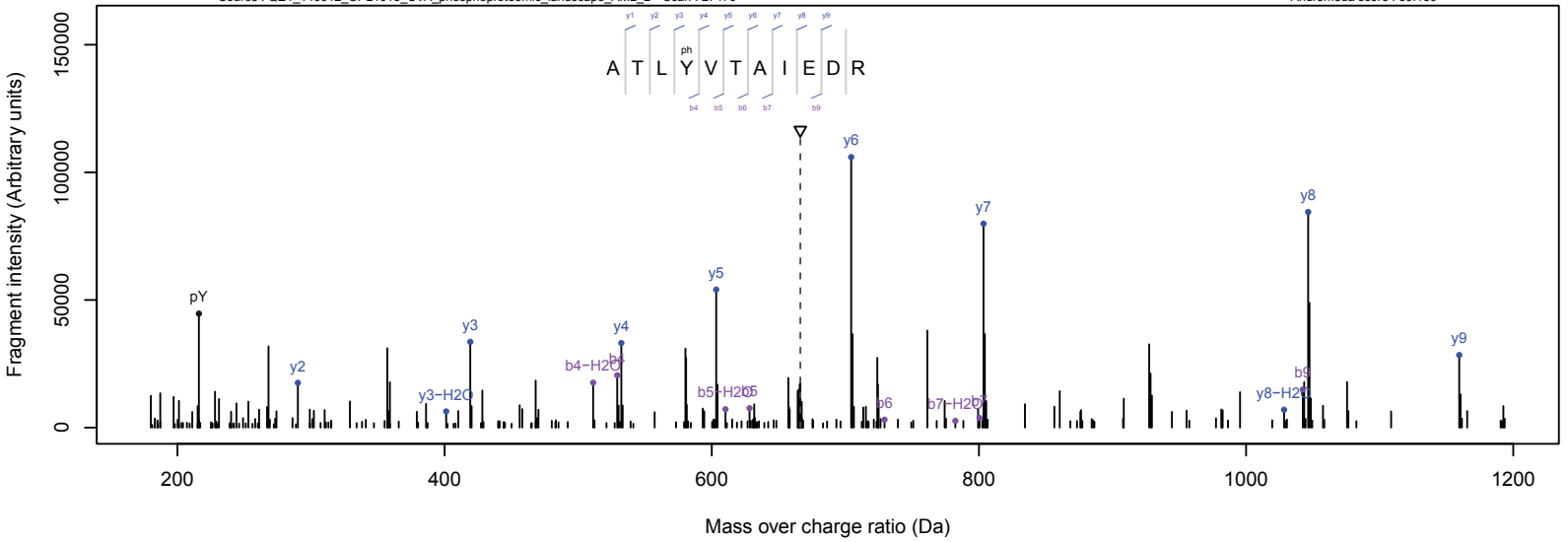
Andromeda score : 89.231



ATLY(ph)VTAIEDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27476

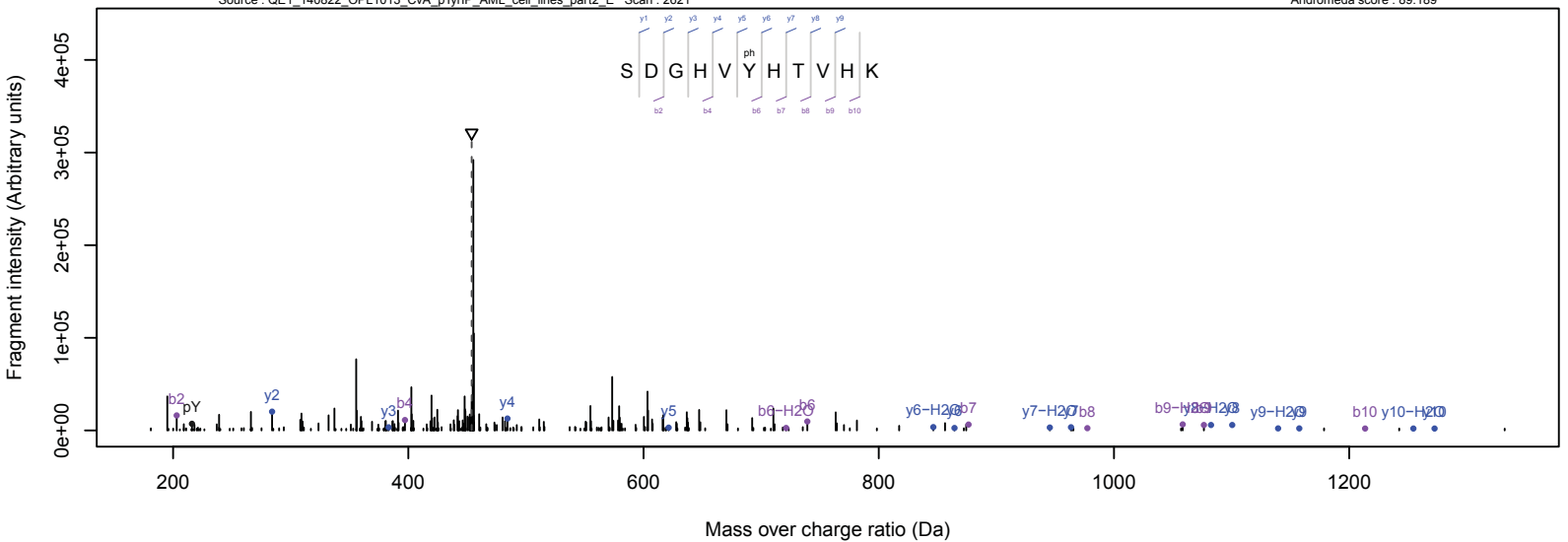
Andromeda score : 89.189



SDGHVY(ph)HTVHK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 2621

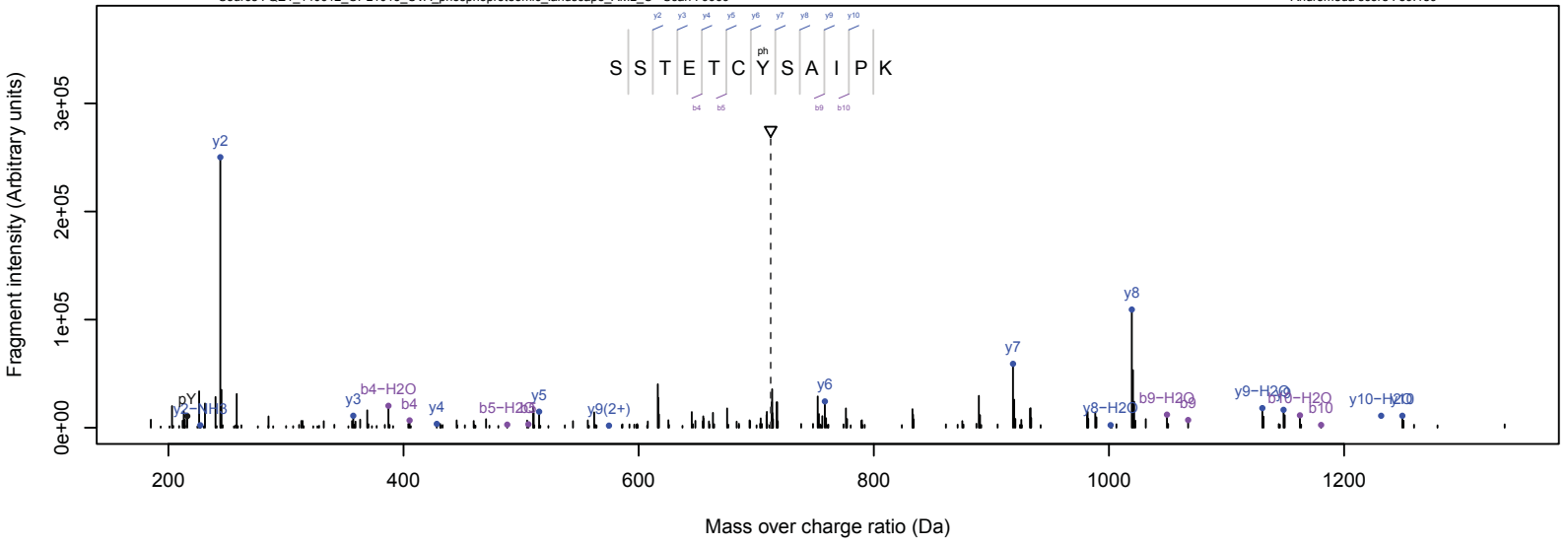
Andromeda score : 89.189



SSTETCY(ph)SAIPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9969

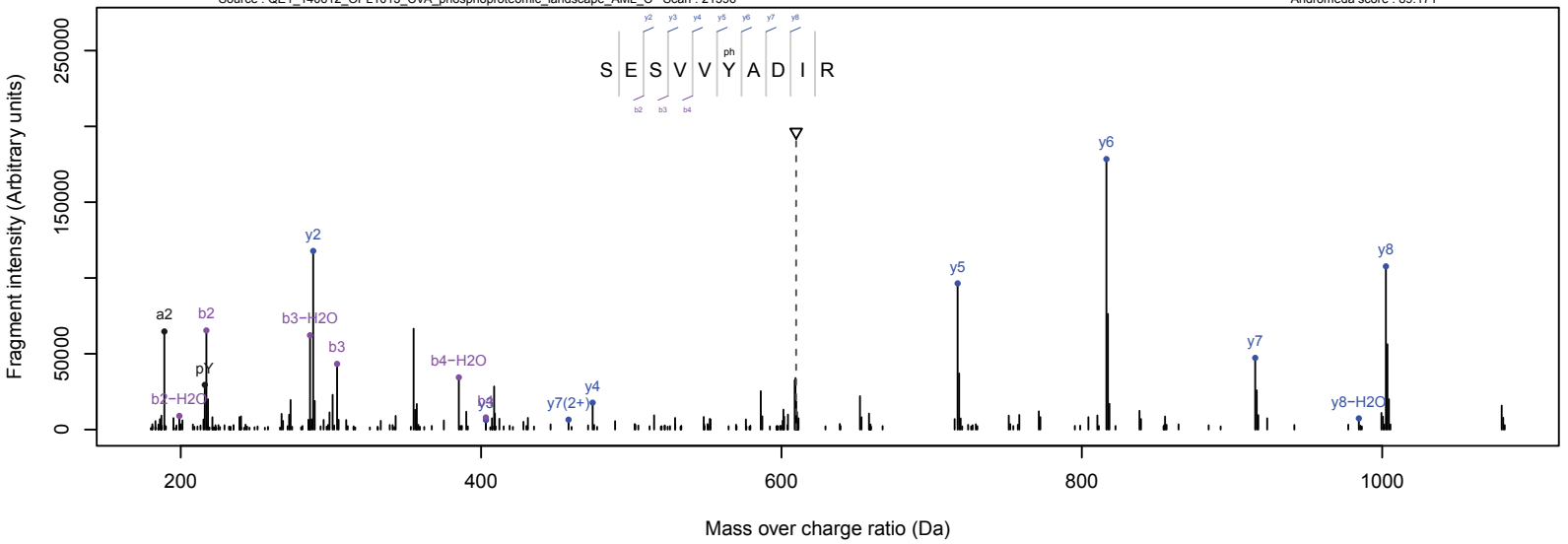
Andromeda score : 89.189



SESVVY(ph)ADIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21596

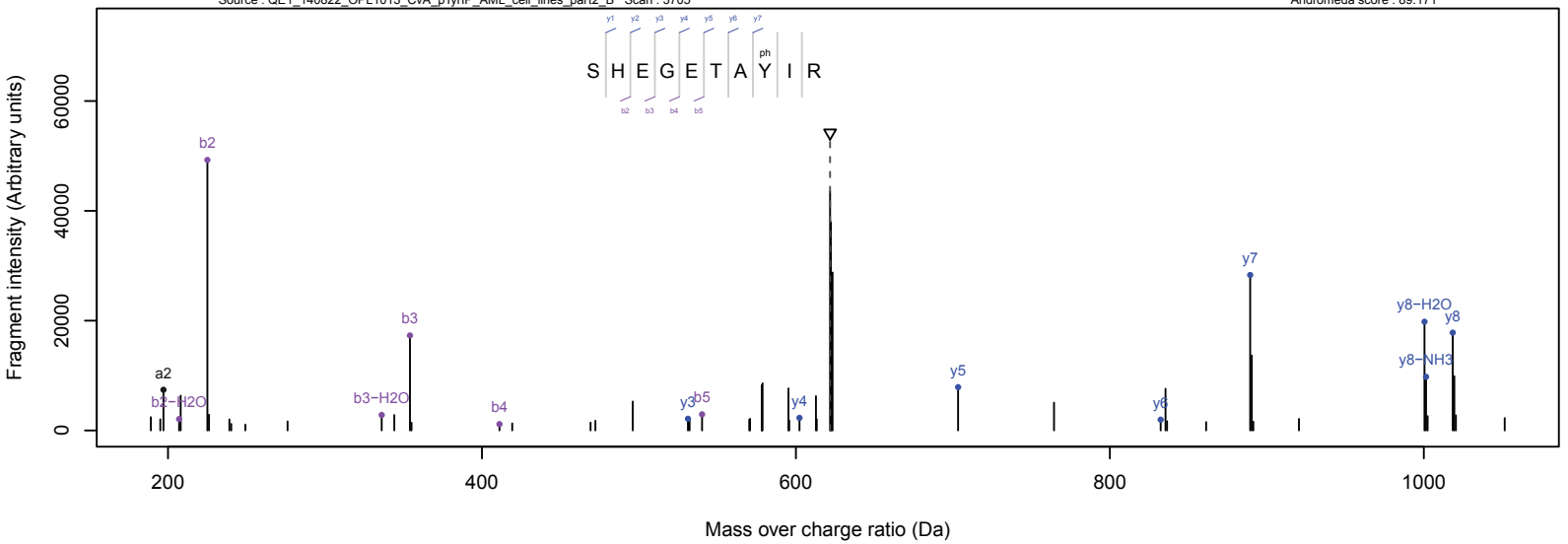
Andromeda score : 89.171



SHEGETAY(ph)IR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3705

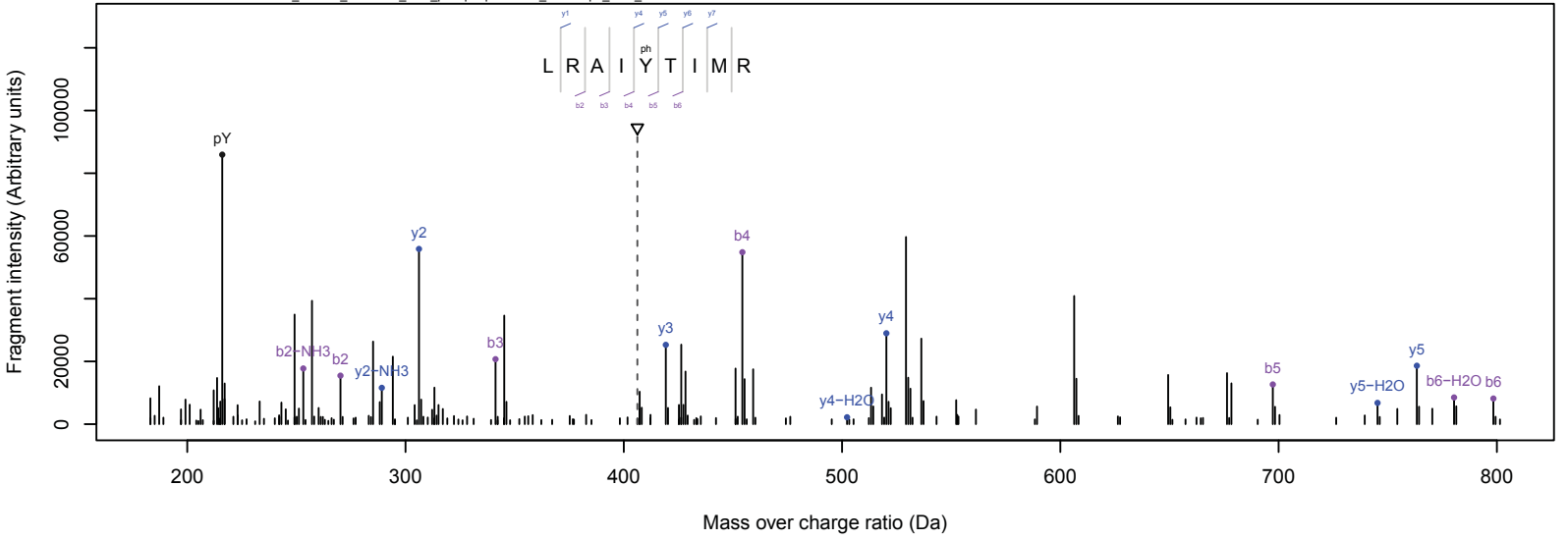
Andromeda score : 89.171



LRAIY(ph)TIMR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25414

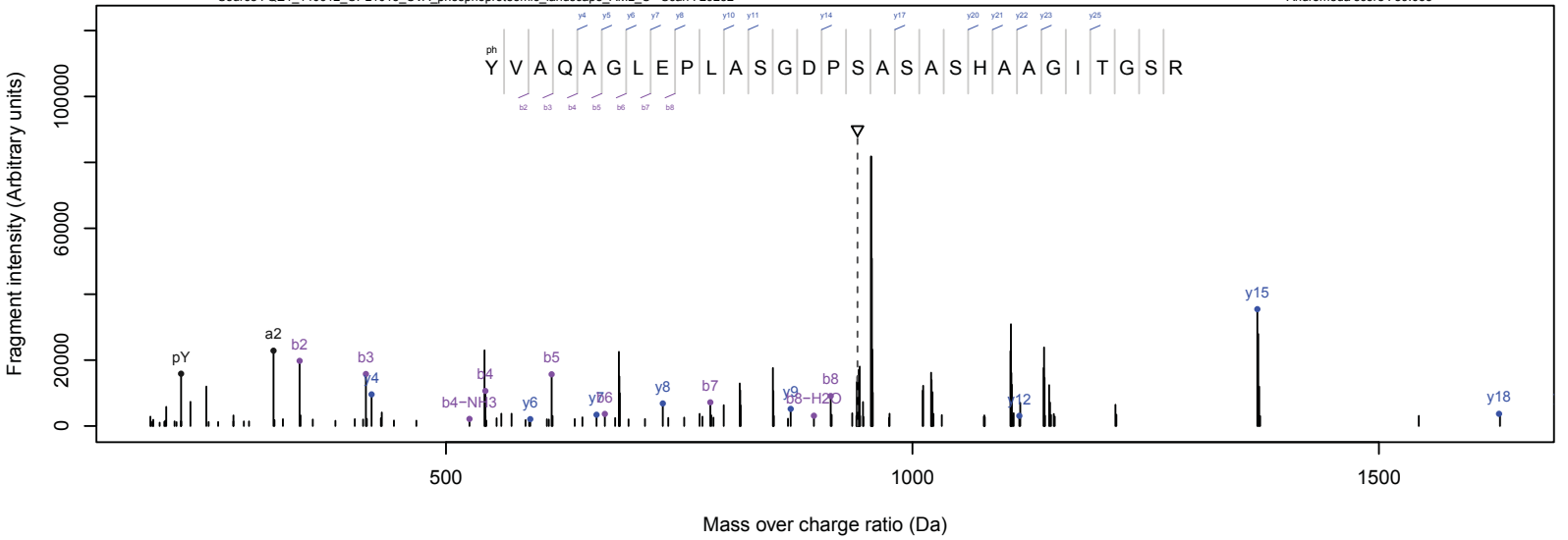
Andromeda score : 89.08



Y(ph)VAQAGLEPLASGDPSASASHAGITGSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29282

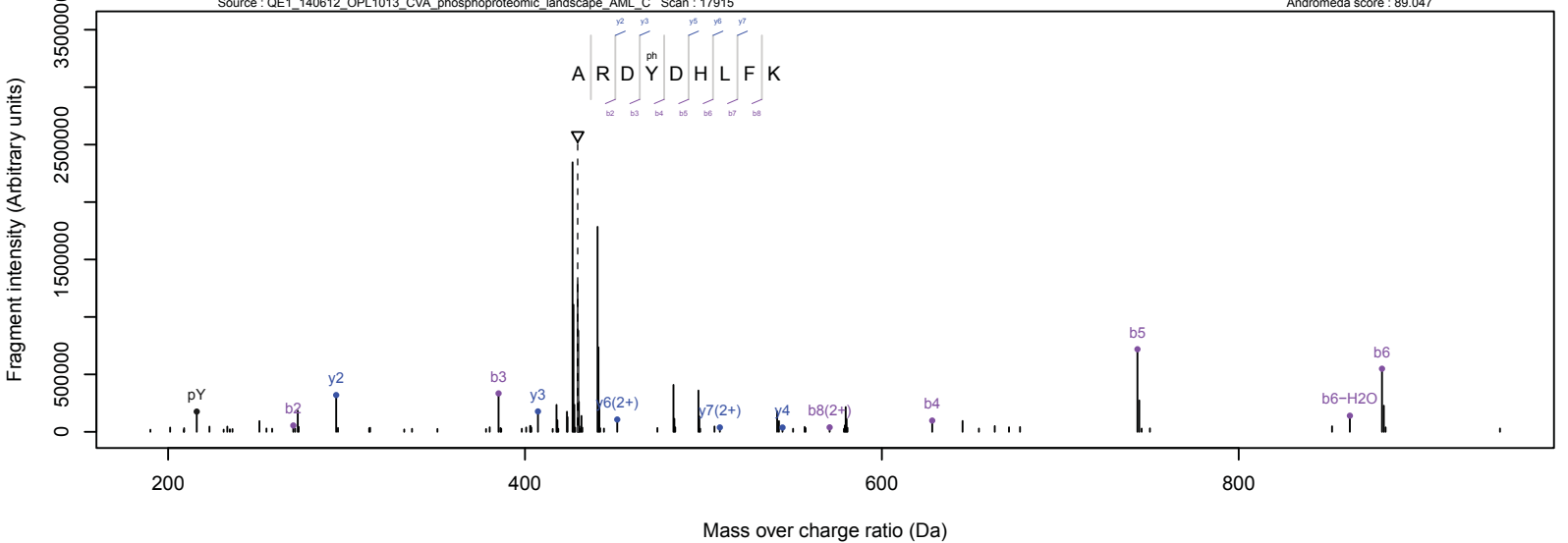
Andromeda score : 89.055



(ac)ARDY(ph)DHLFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17915

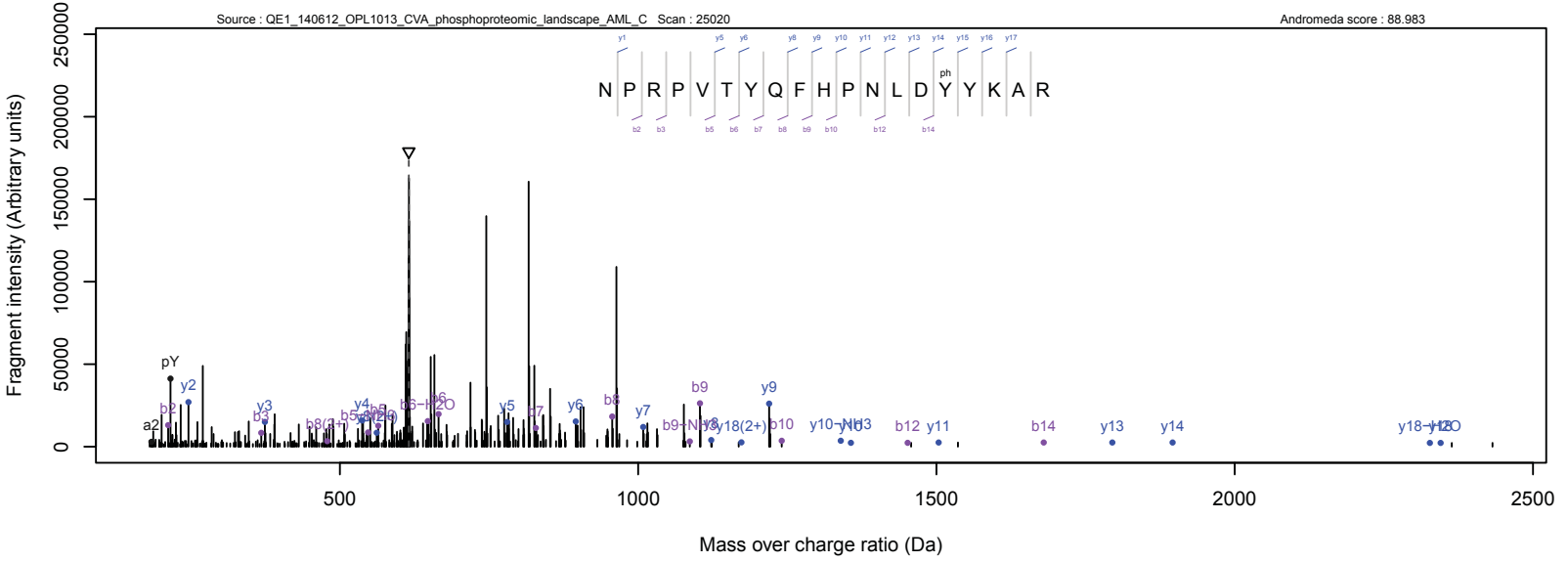
Andromeda score : 89.047



**NPRPVTYQFHPNLDYYKAR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25020

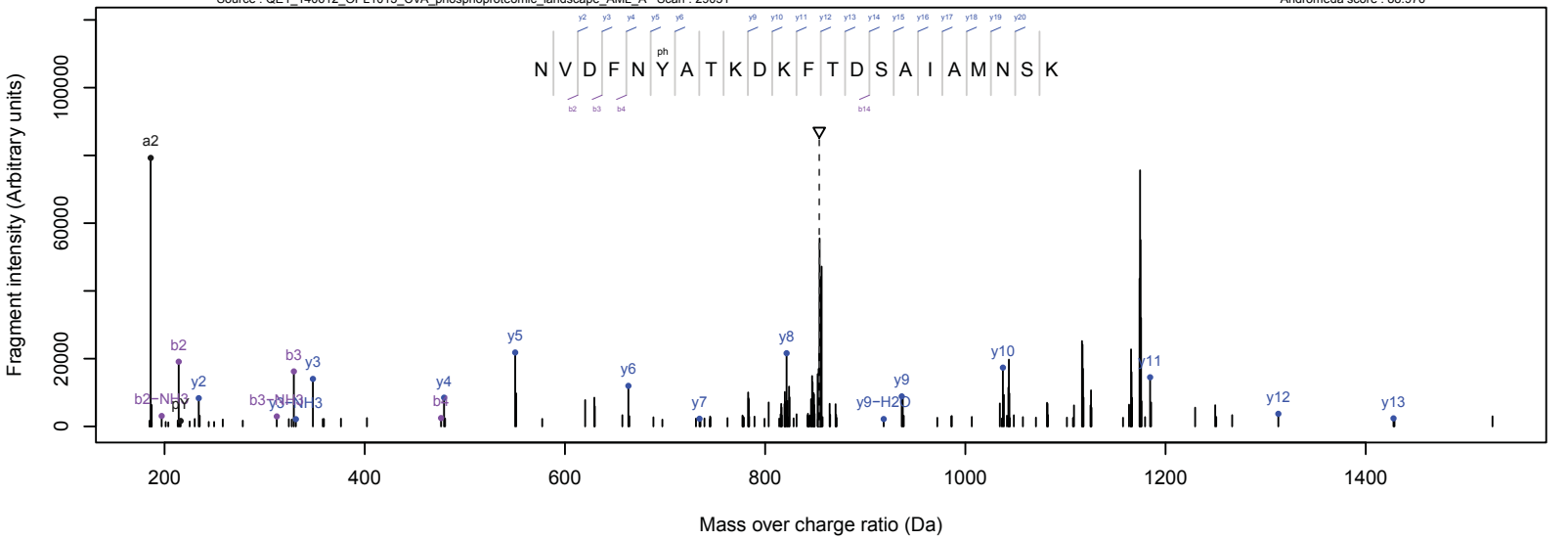
Andromeda score : 88.983



**NVDFNY(ph)ATKDKFTDSAIAMNSK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29051

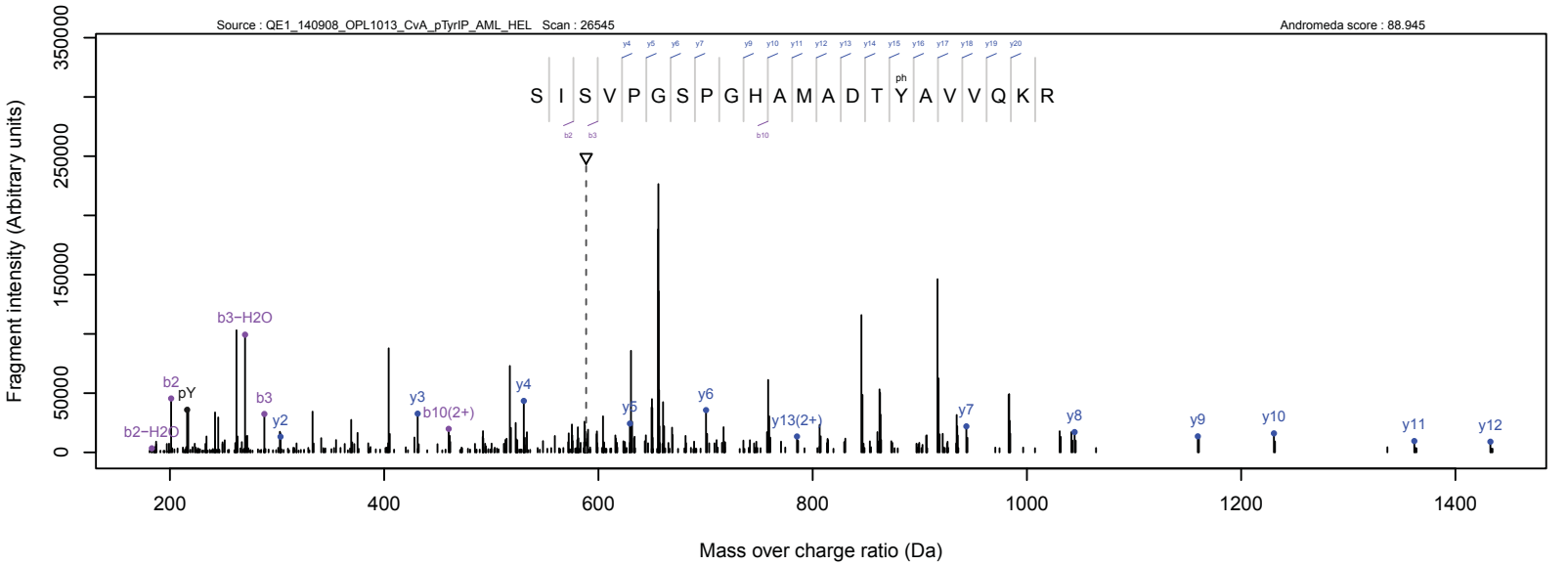
Andromeda score : 88.976



**SISVPGSGHAMADTY(ph)AVVQKR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 26545

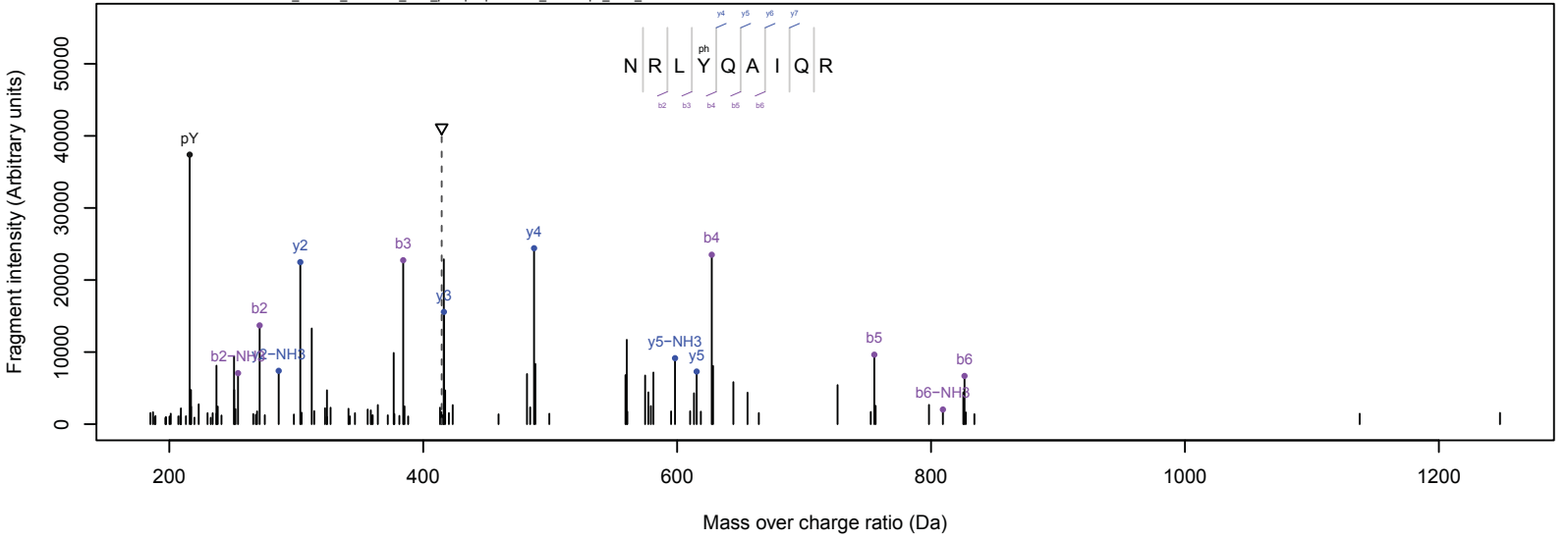
Andromeda score : 88.945



**NRLY(ph)QAIQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9430

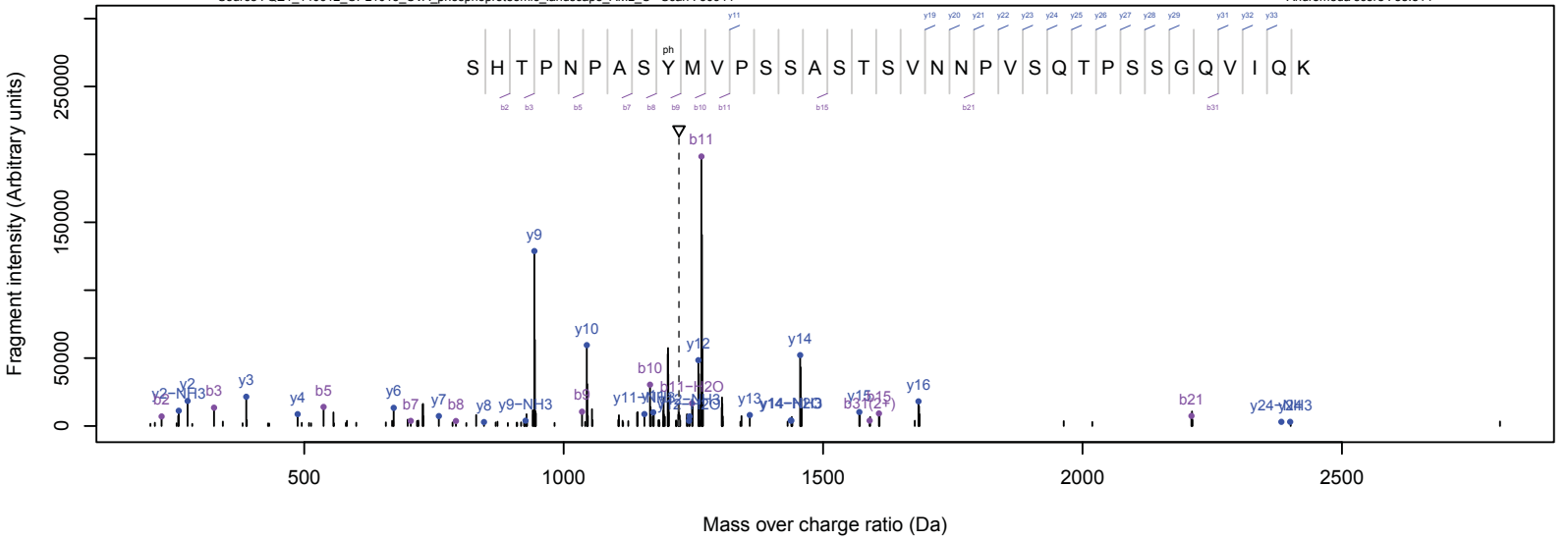
Andromeda score : 88.819



**SHTPNPASY(ph)MVPSSASTSVNNPVSQTPSSGQVIQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30944

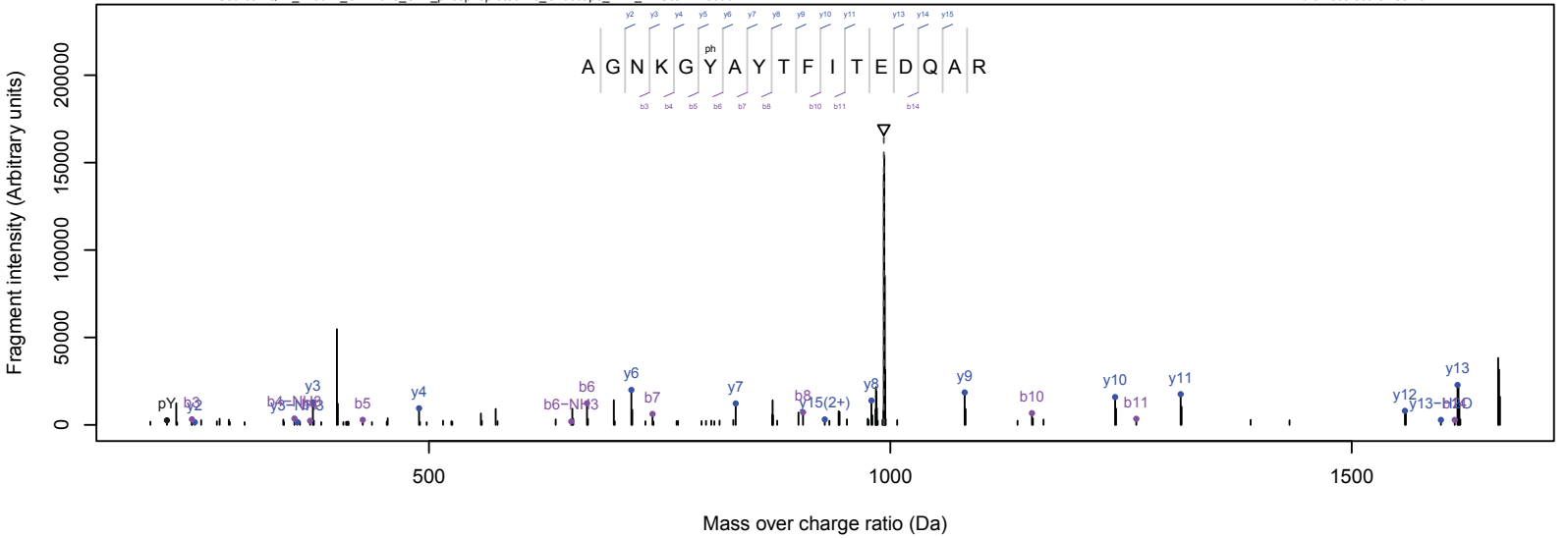
Andromeda score : 88.814



**AGNKGY(ph)AYTFITEDQAR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23596

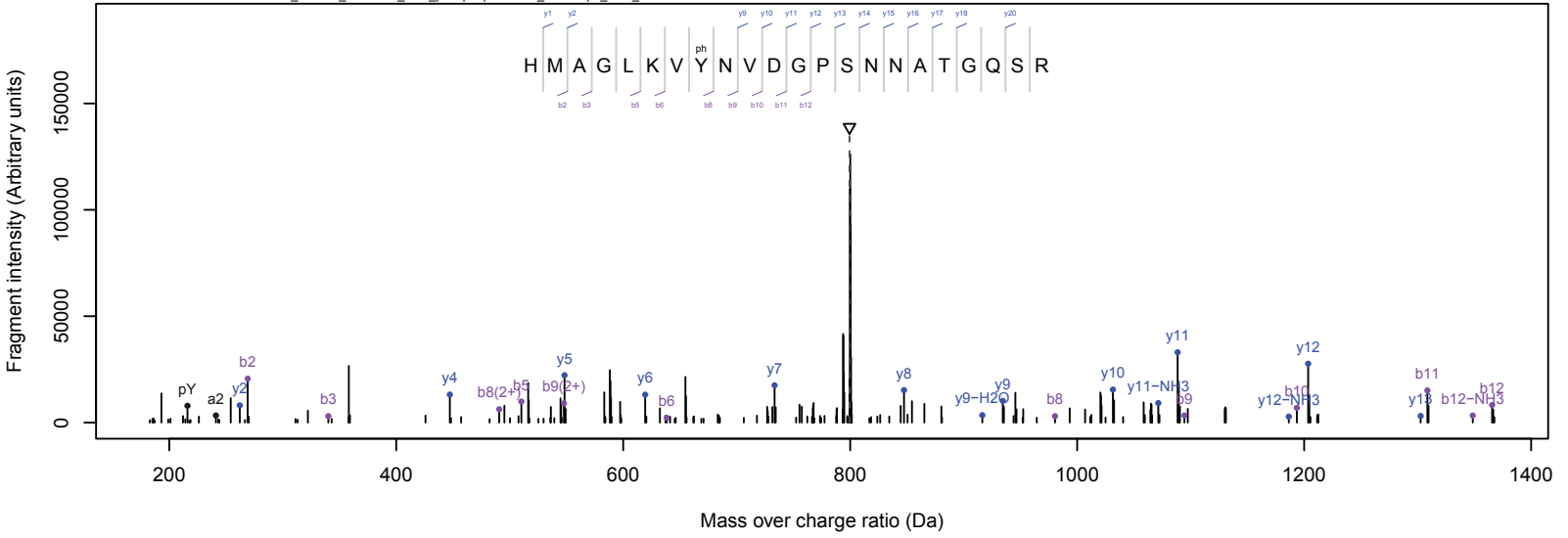
Andromeda score : 88.781



**HMAGLKVY(ph)NVDGPSNNATGQSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15983

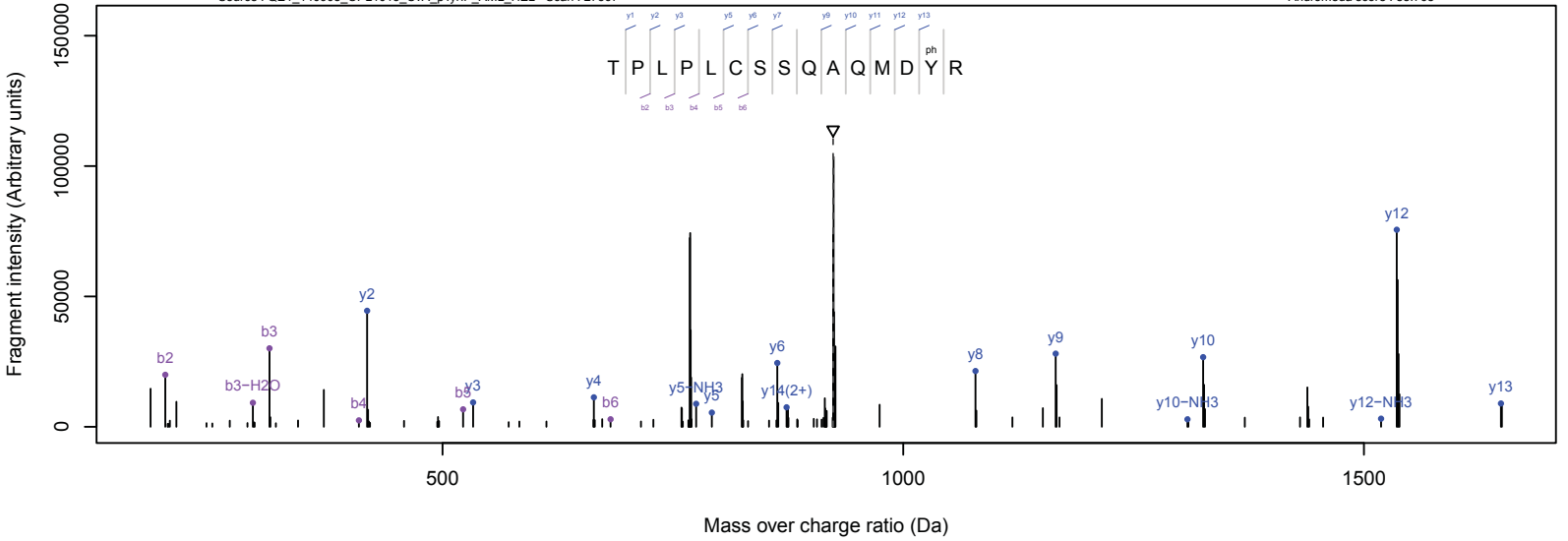
Andromeda score : 88.768



**TPLPLCQQAMDY(ph)R-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 27567

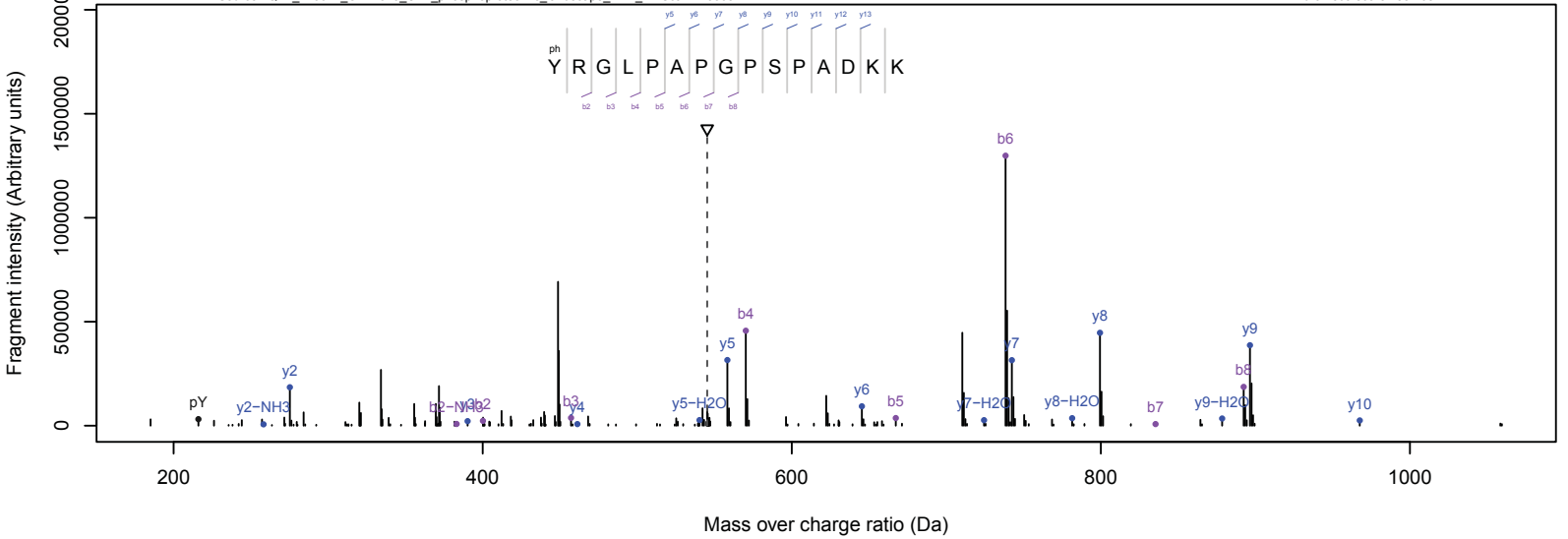
Andromeda score : 88.768



**Y(ph)RGLPAPGSPADKK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10065

Andromeda score : 88.768

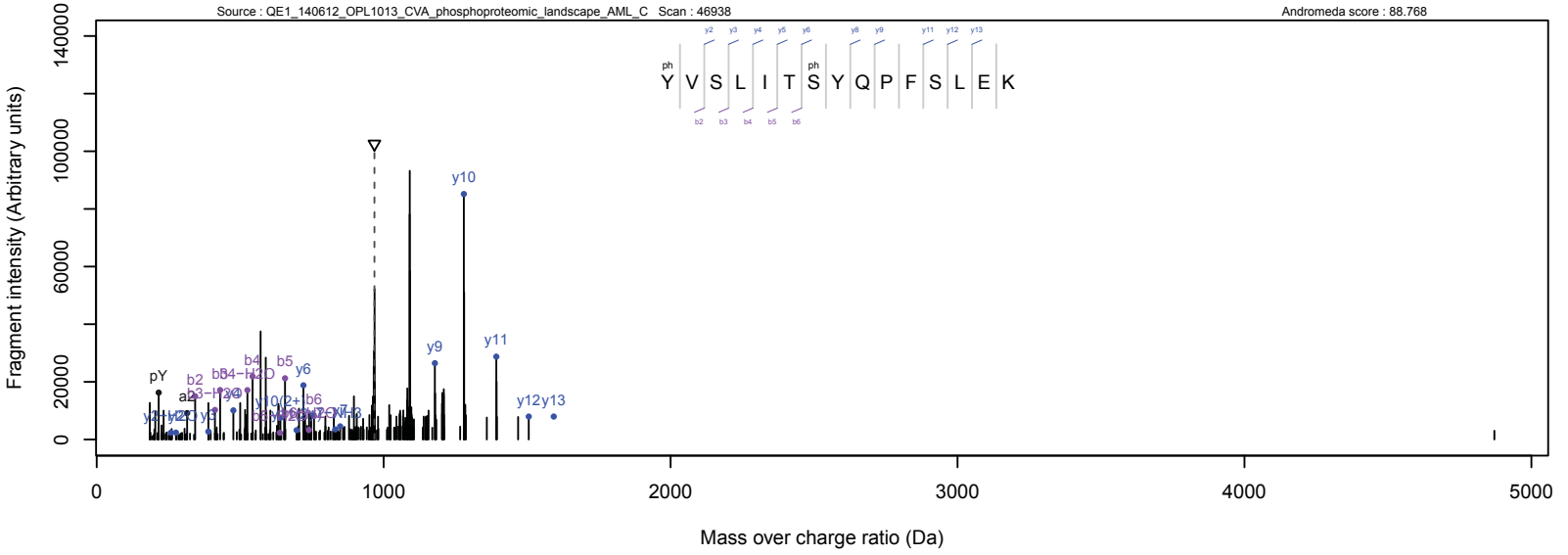




Y(ph)VSLITS(ph)YQPFSLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46938

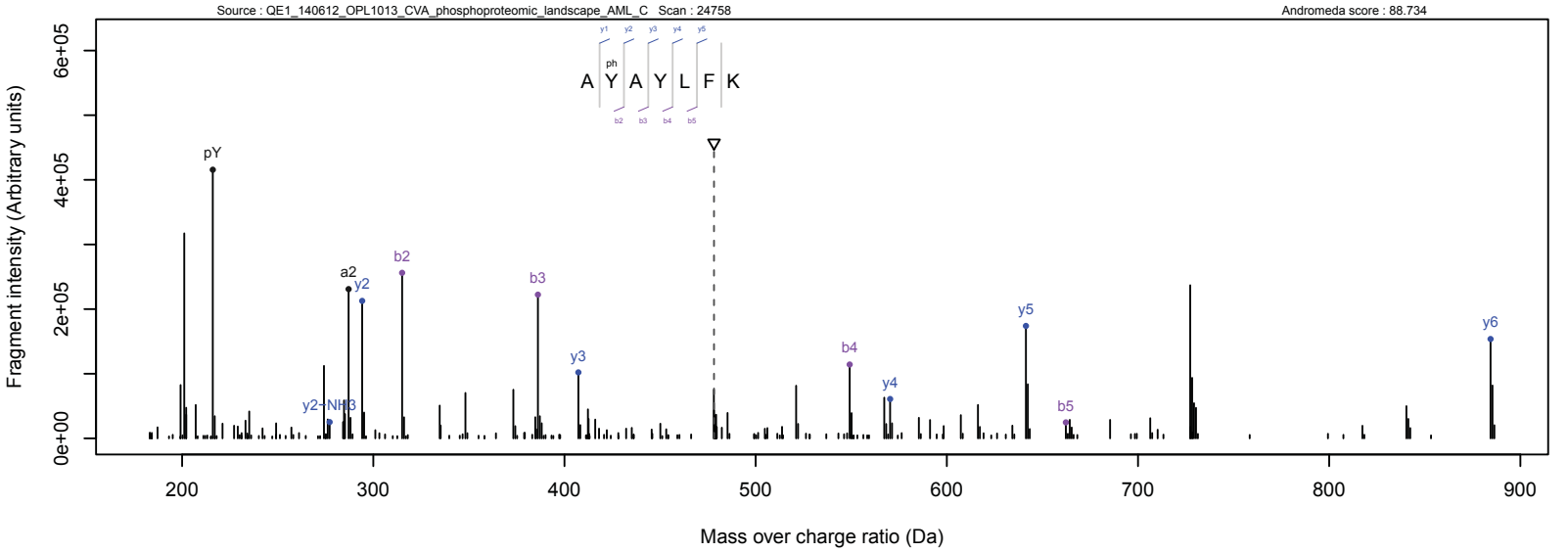
Andromeda score : 88.768



AY(ph)AYLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24758

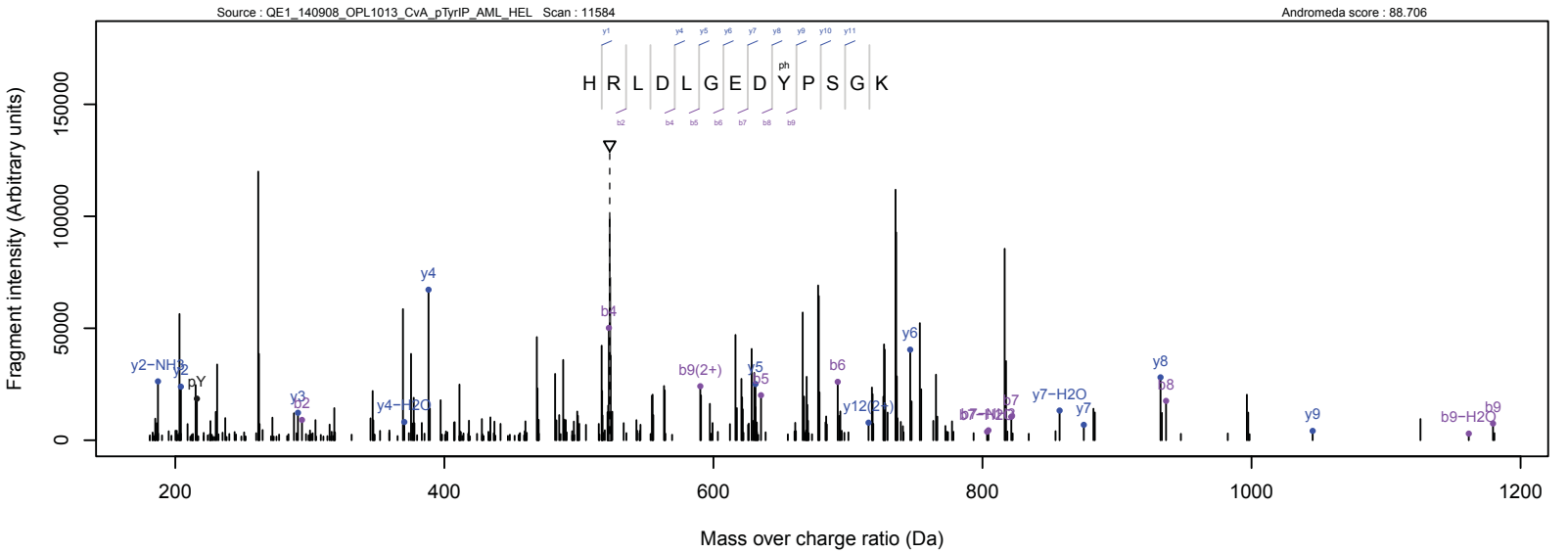
Andromeda score : 88.734



HRLDLGEDY(ph)PSGK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 11584

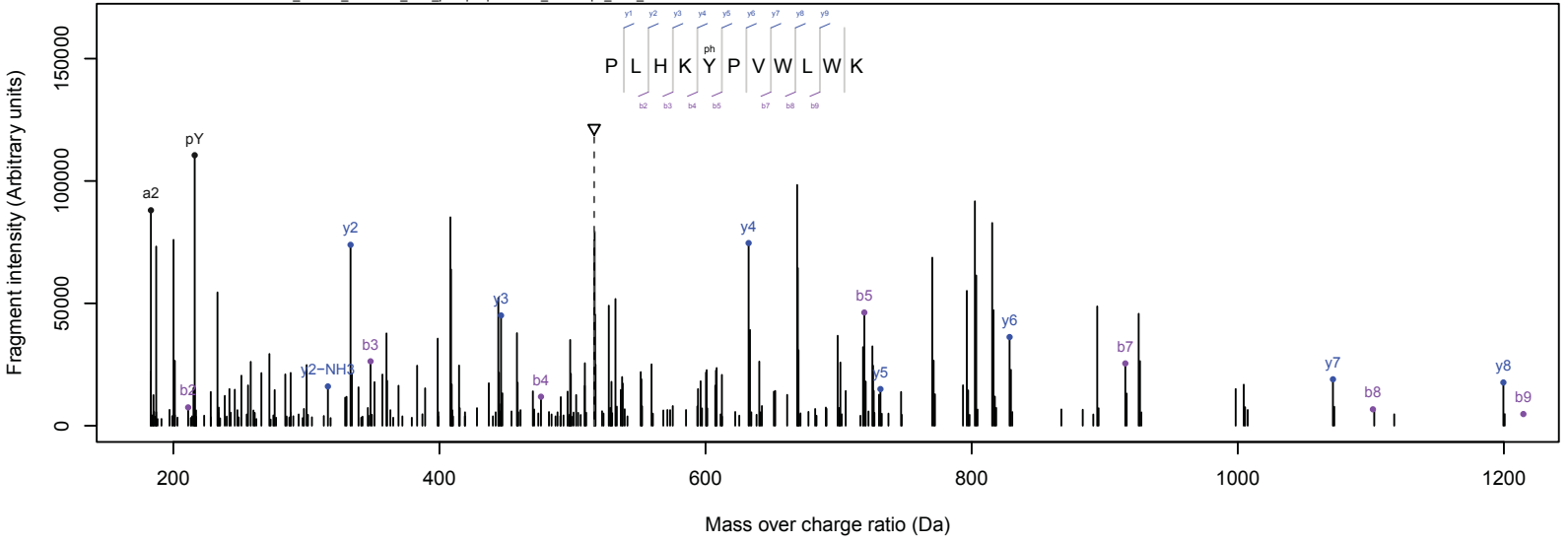
Andromeda score : 88.706



PLHKY(ph)PVWLWK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31696

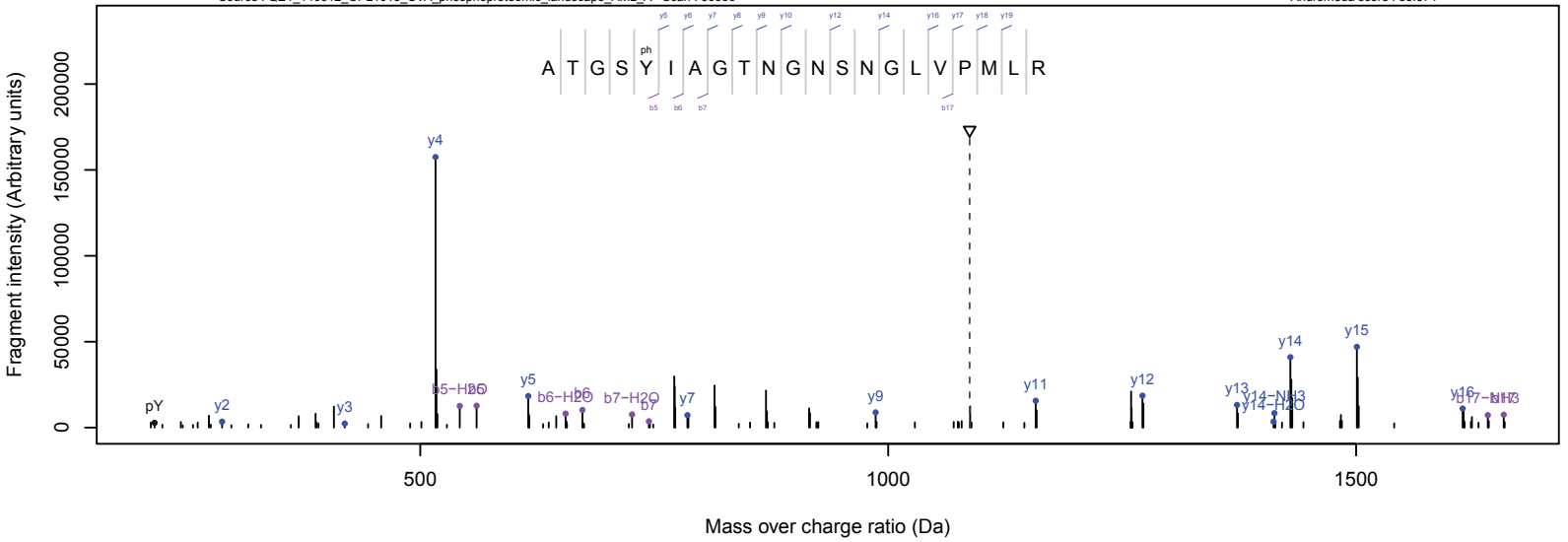
Andromeda score : 88.596



ATGSY(ph)IAGTNGNSNGLVPMRL-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36556

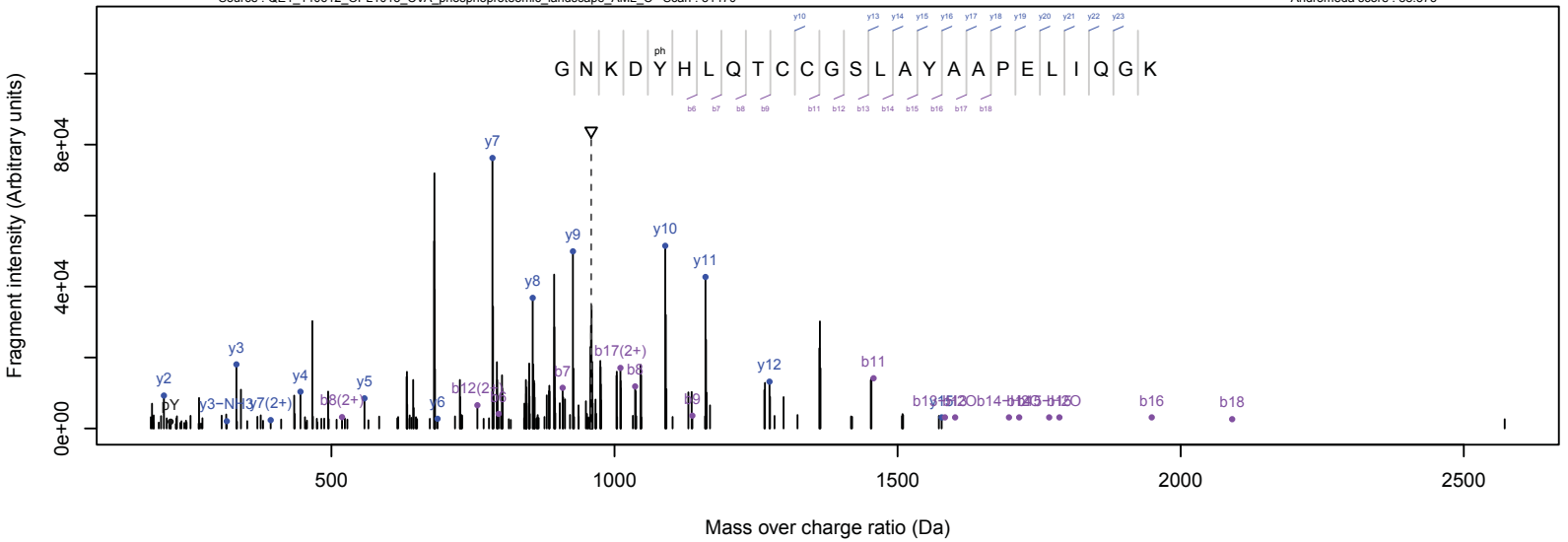
Andromeda score : 88.574



GNKDY(ph)HLQTCGSLAYAAPELIQGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31470

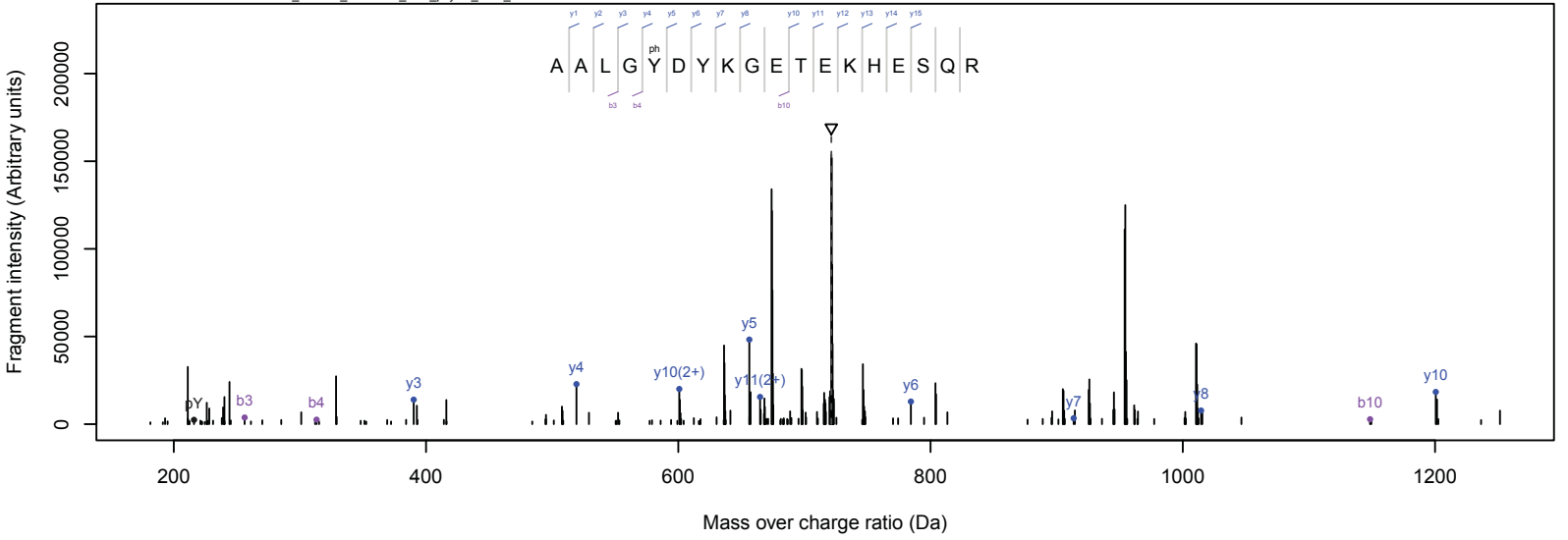
Andromeda score : 88.573



**AALGY(ph)DYKGETEKHESQR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 8673

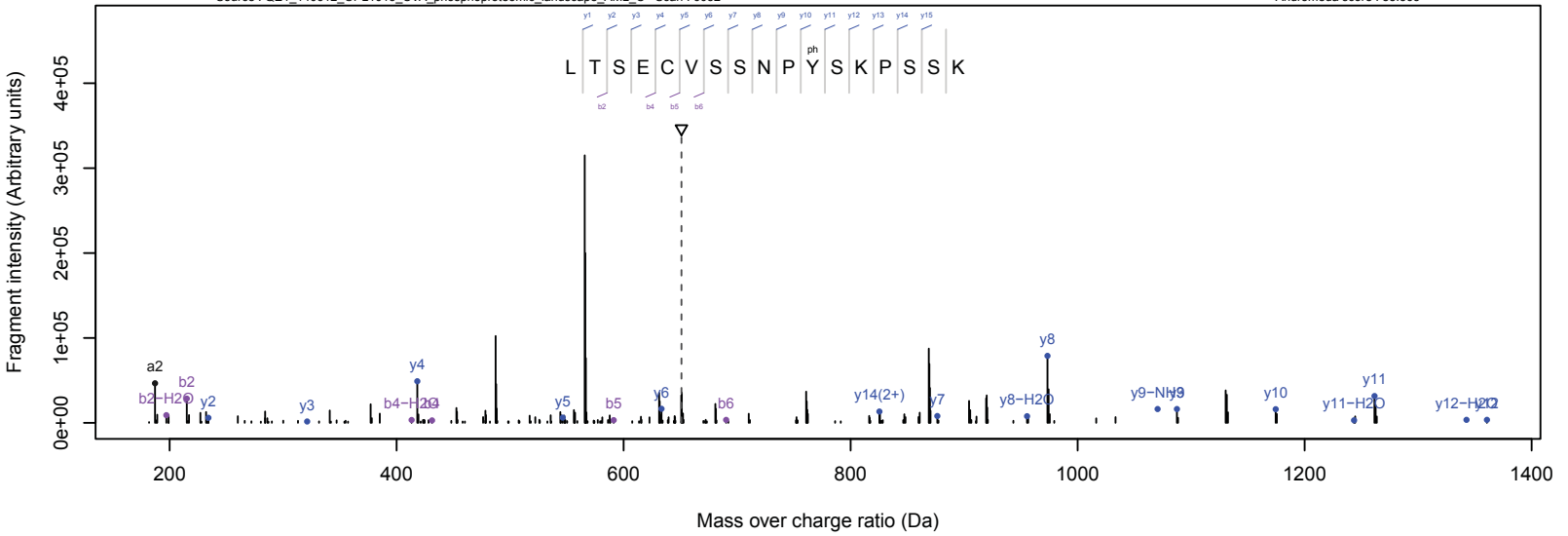
Andromeda score : 88.565



**LTSECVSSNPY(ph)SKPSSK-3+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 6052

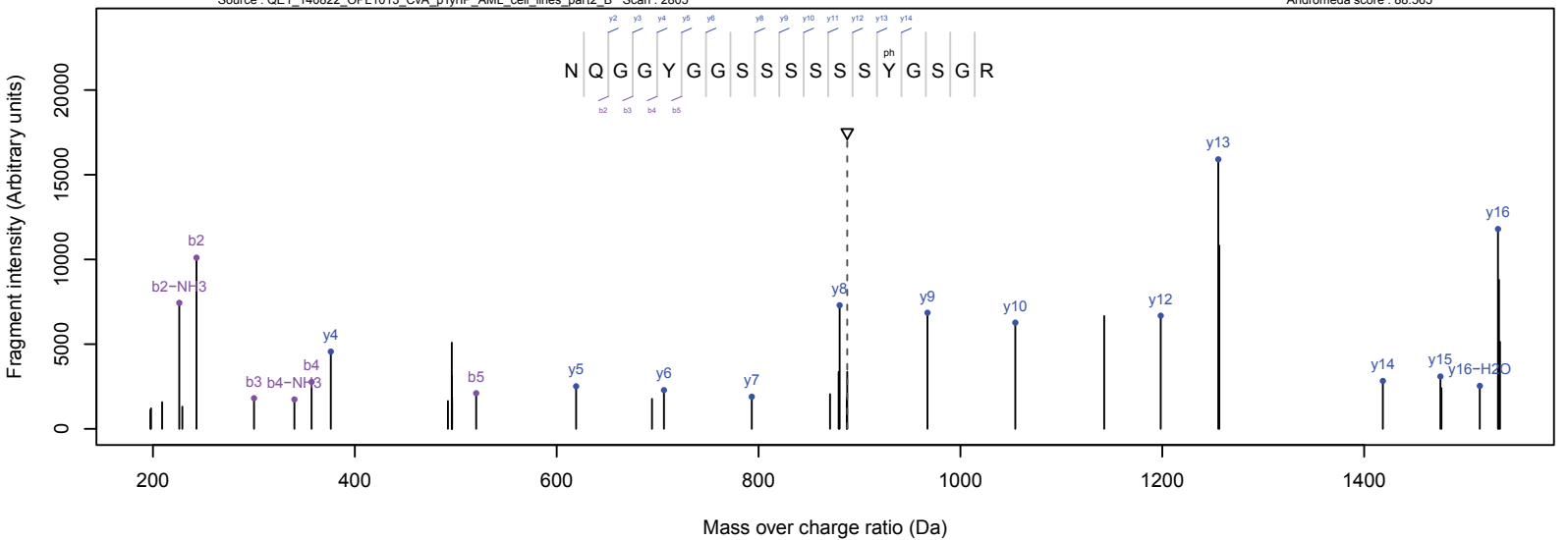
Andromeda score : 88.565



**NQGGYGGSSSSSY(ph)GSGR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 2805

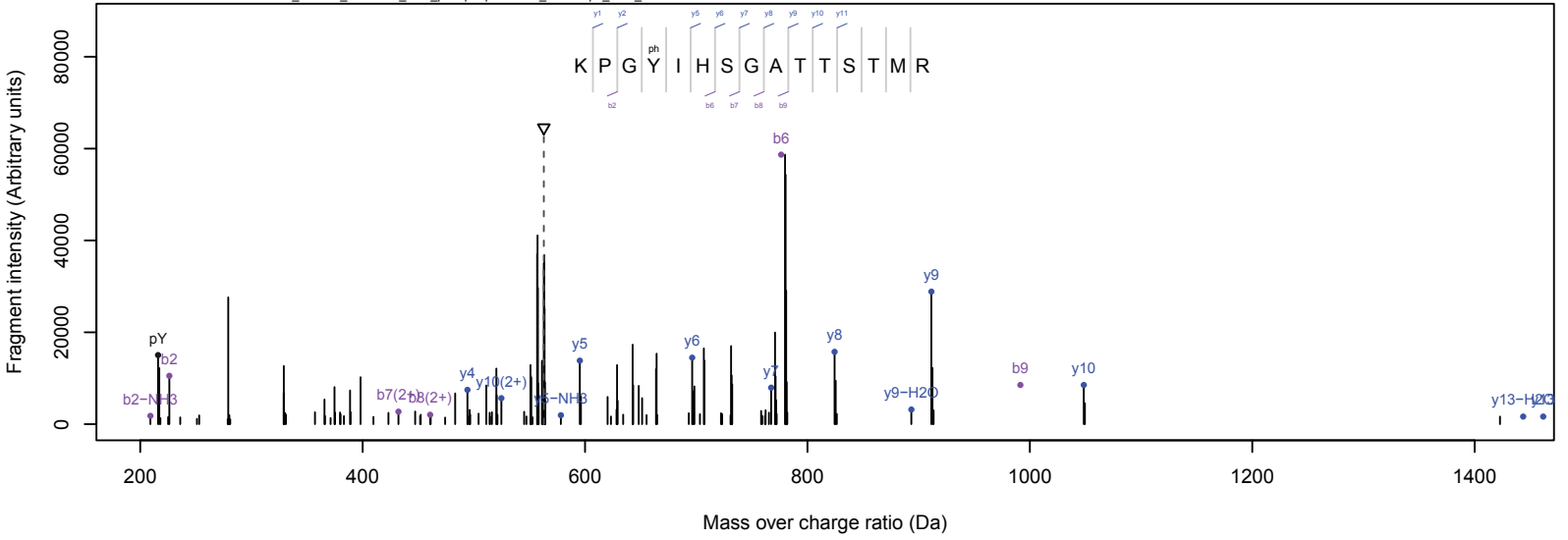
Andromeda score : 88.565



KPGY(ph)IHSGATTSTM<sup>3+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 5888

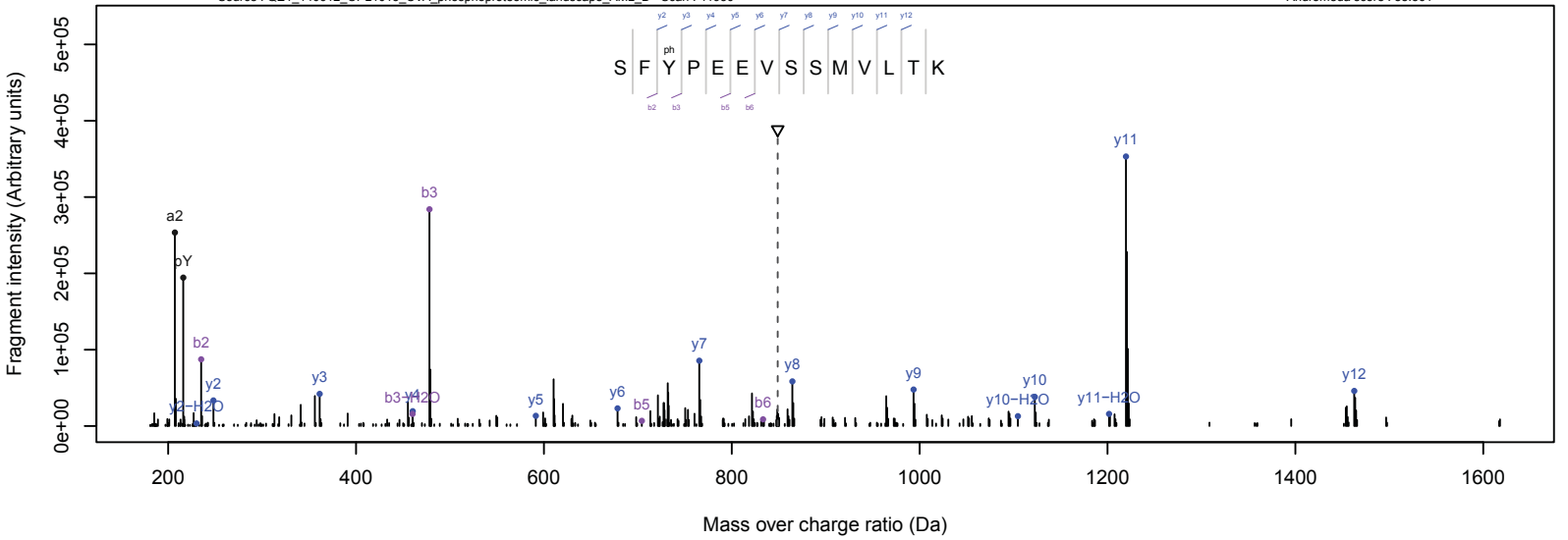
Andromeda score : 88.561



SFY(ph)PEEVSSMLTK<sup>2+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41930

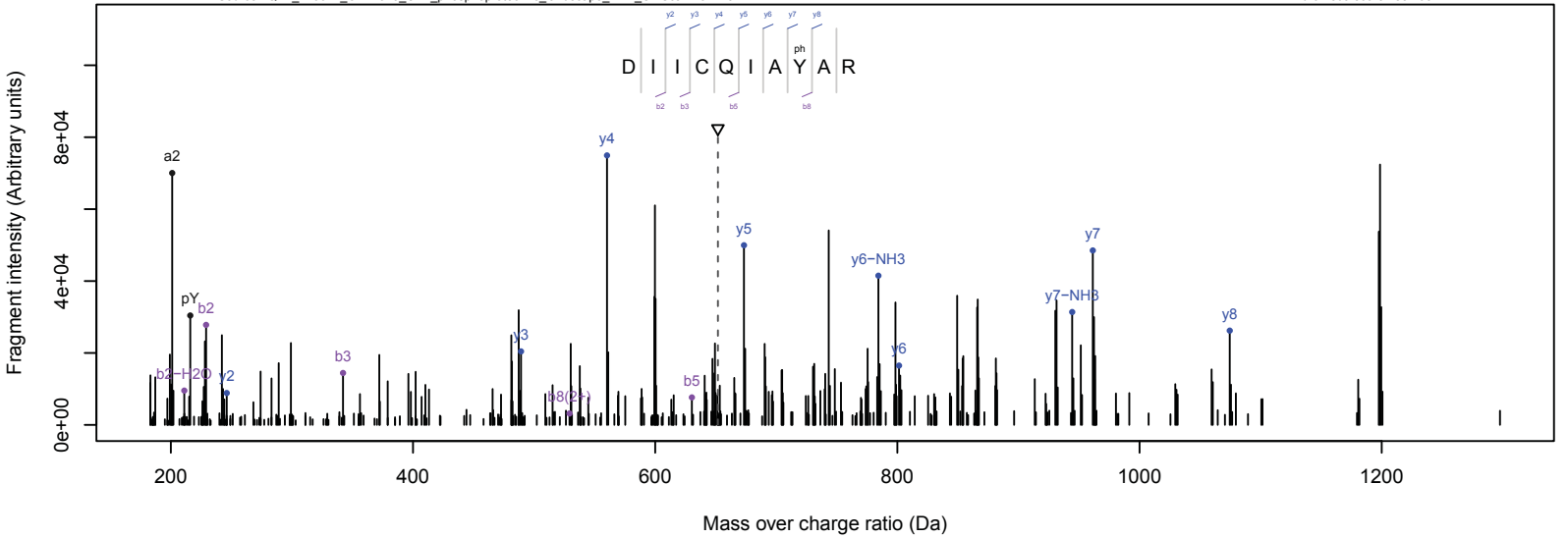
Andromeda score : 88.561



DIICQIAY(ph)AR<sup>2+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31415

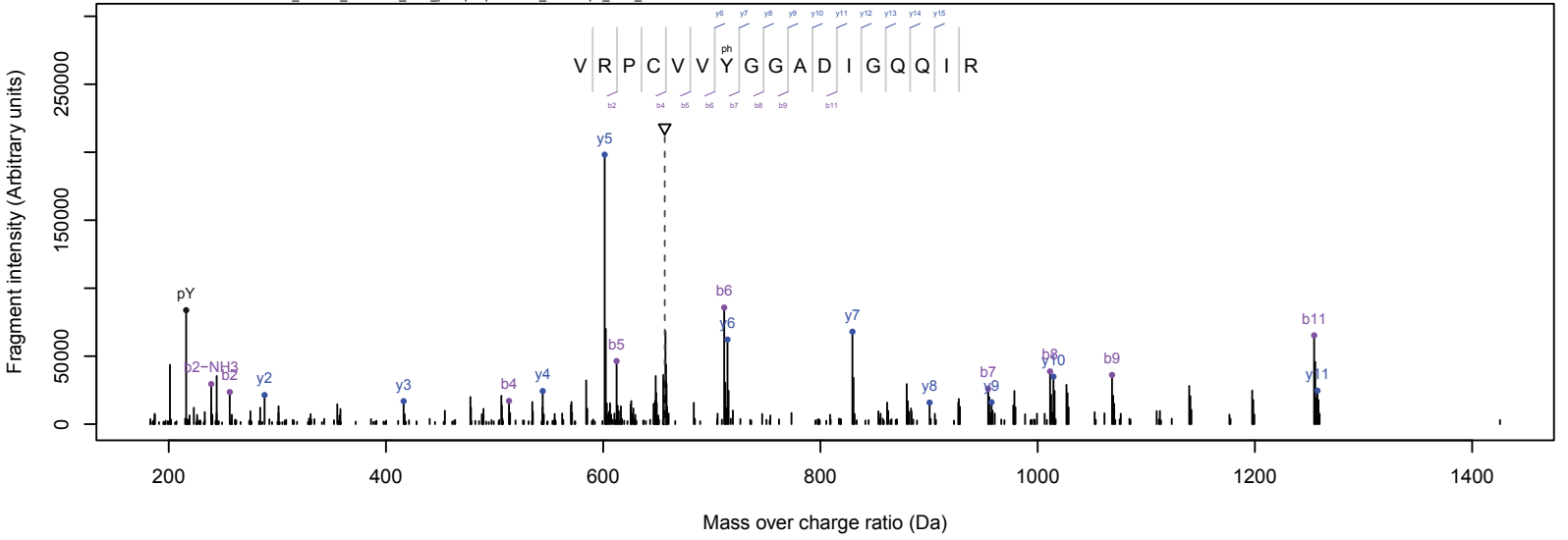
Andromeda score : 88.495



VRPCVV(ph)GGADIGQQIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23956

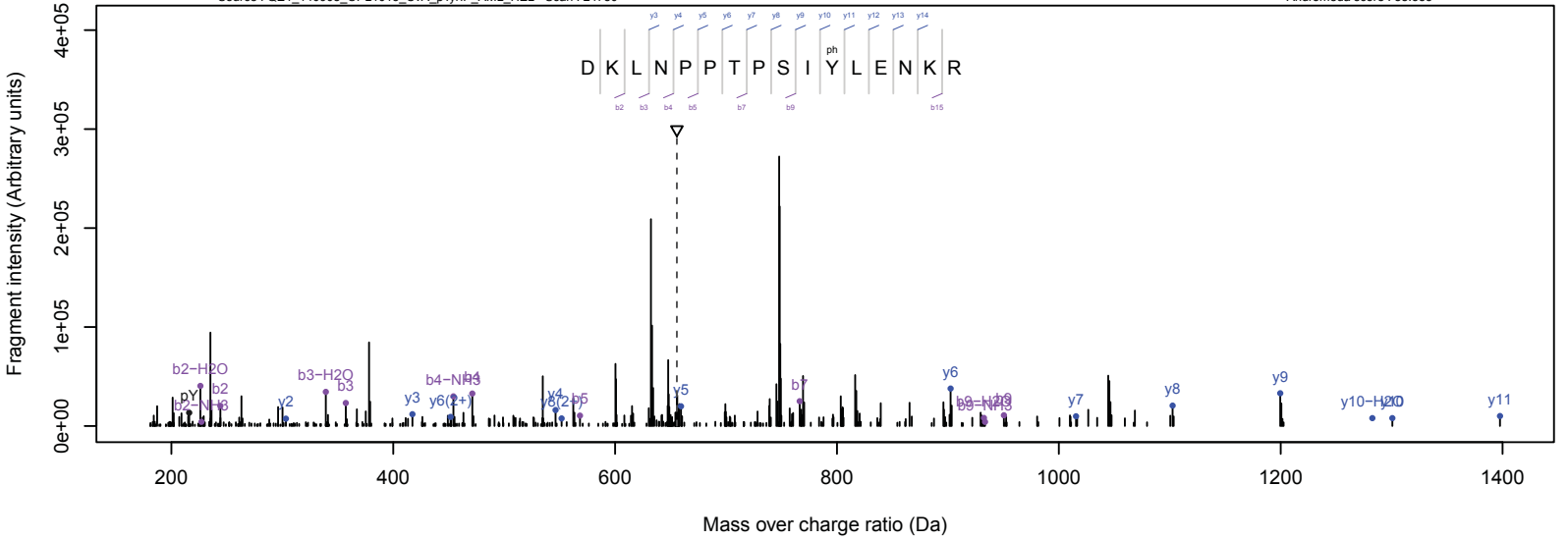
Andromeda score : 88.427



DKLNPTPSIY(ph)LENKR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24780

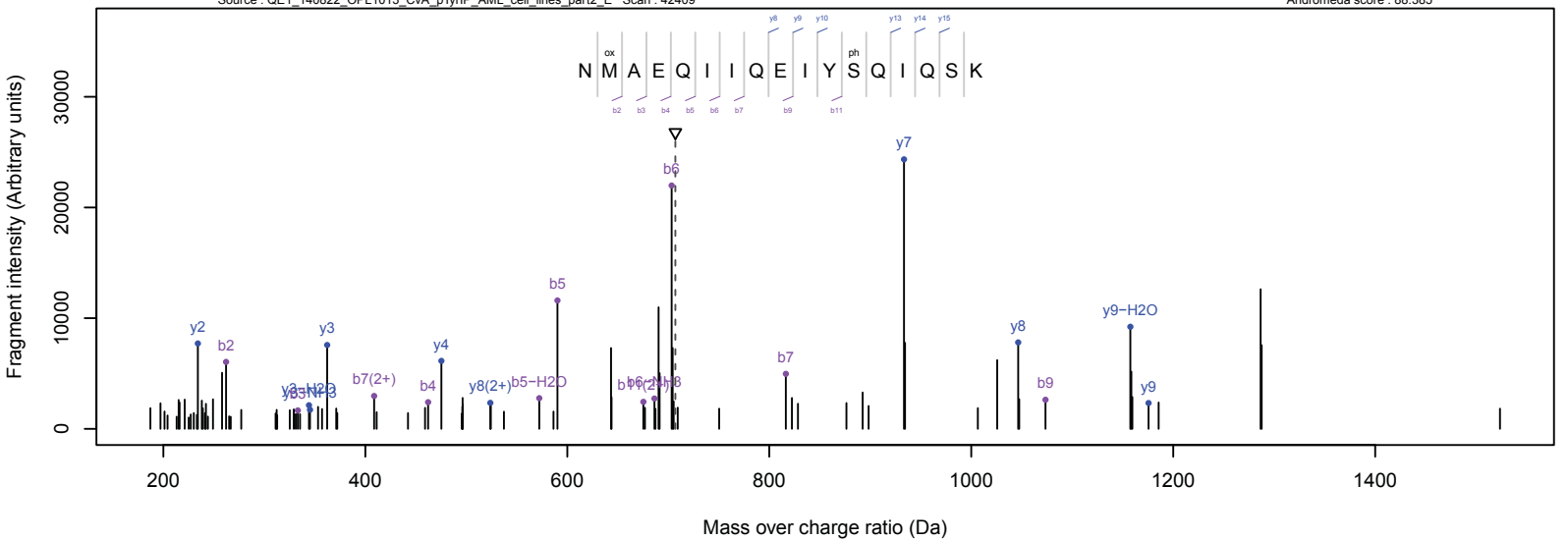
Andromeda score : 88.385



NM(ox)AEQIIQEIYS(ph)QIQSK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 42409

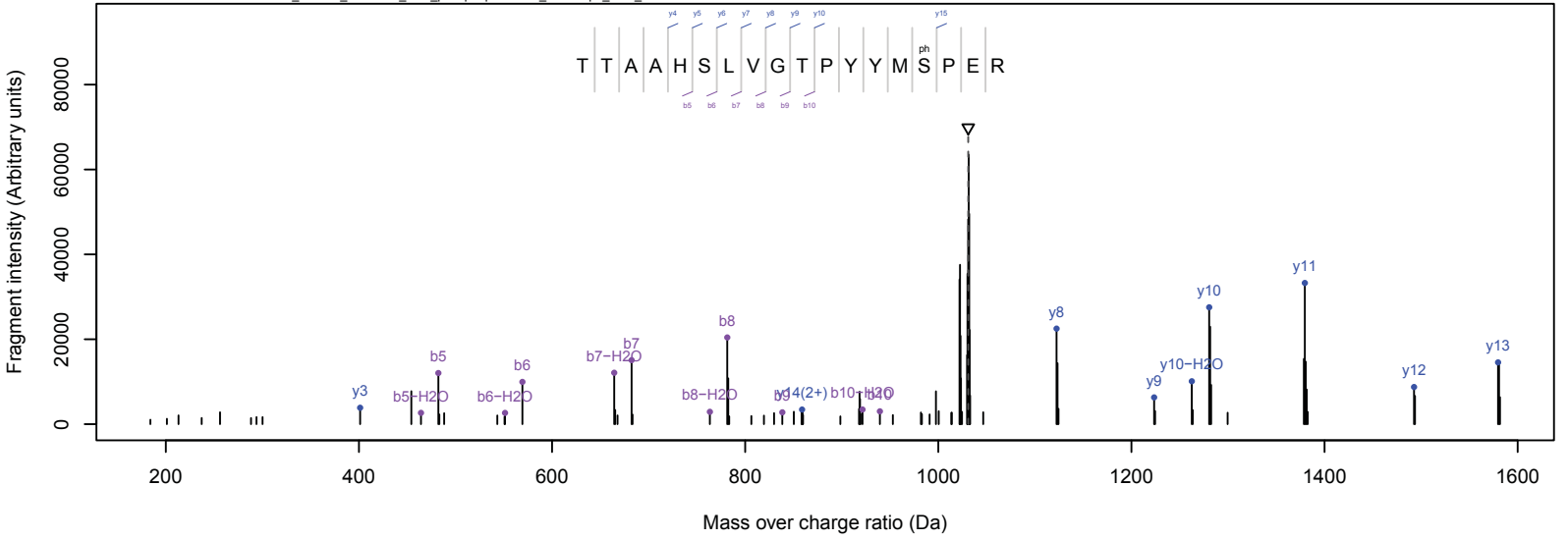
Andromeda score : 88.385



TTAAHSLVGTPTYMSPER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24786

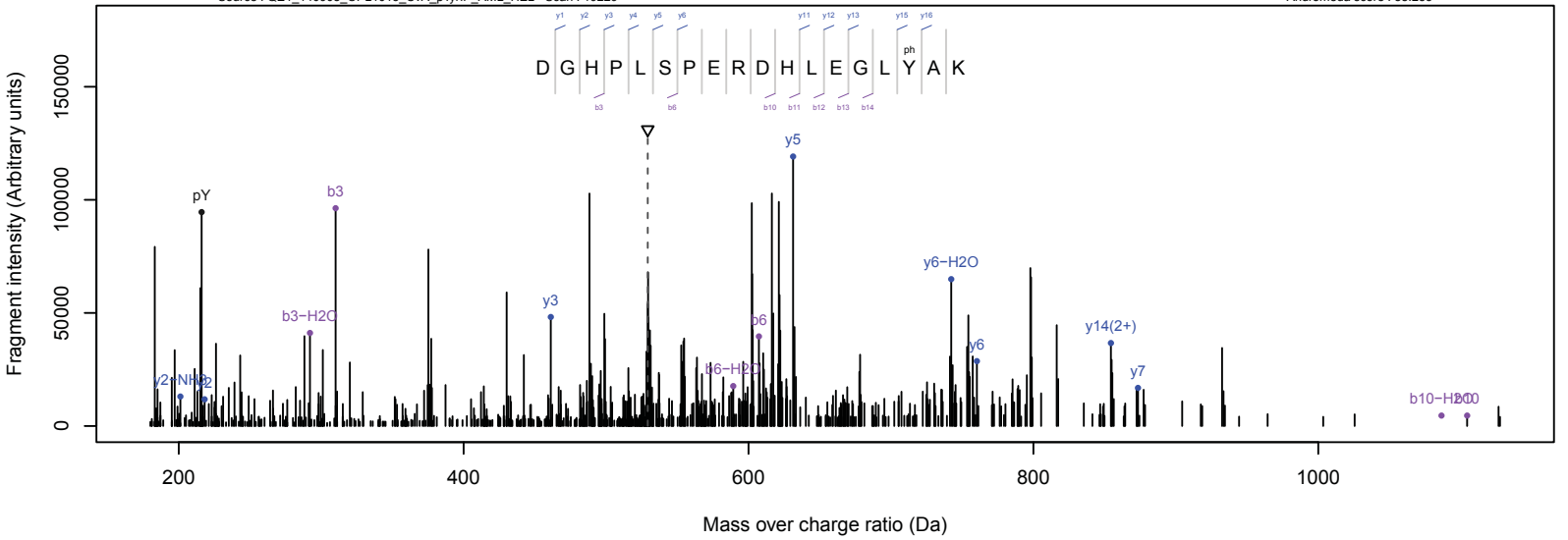
Andromeda score : 88.385



DGHLSPERDHLLEGLYAK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 19229

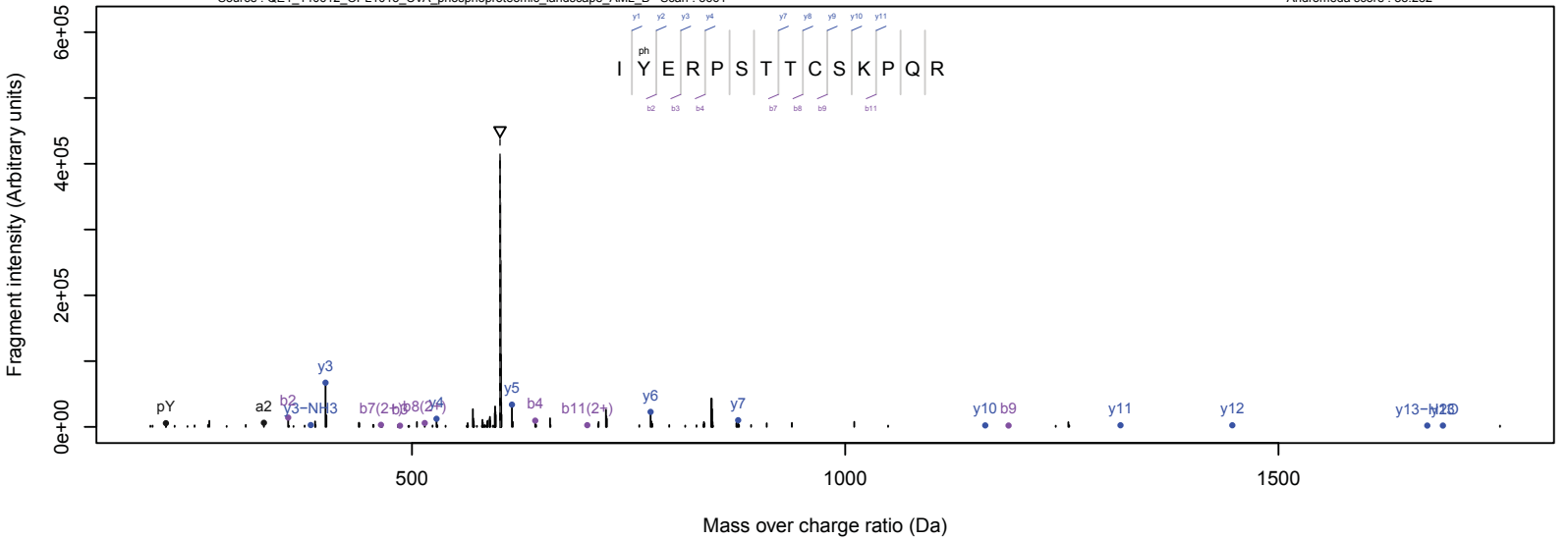
Andromeda score : 88.283



IY(ph)ERPSTTCSKPKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 3661

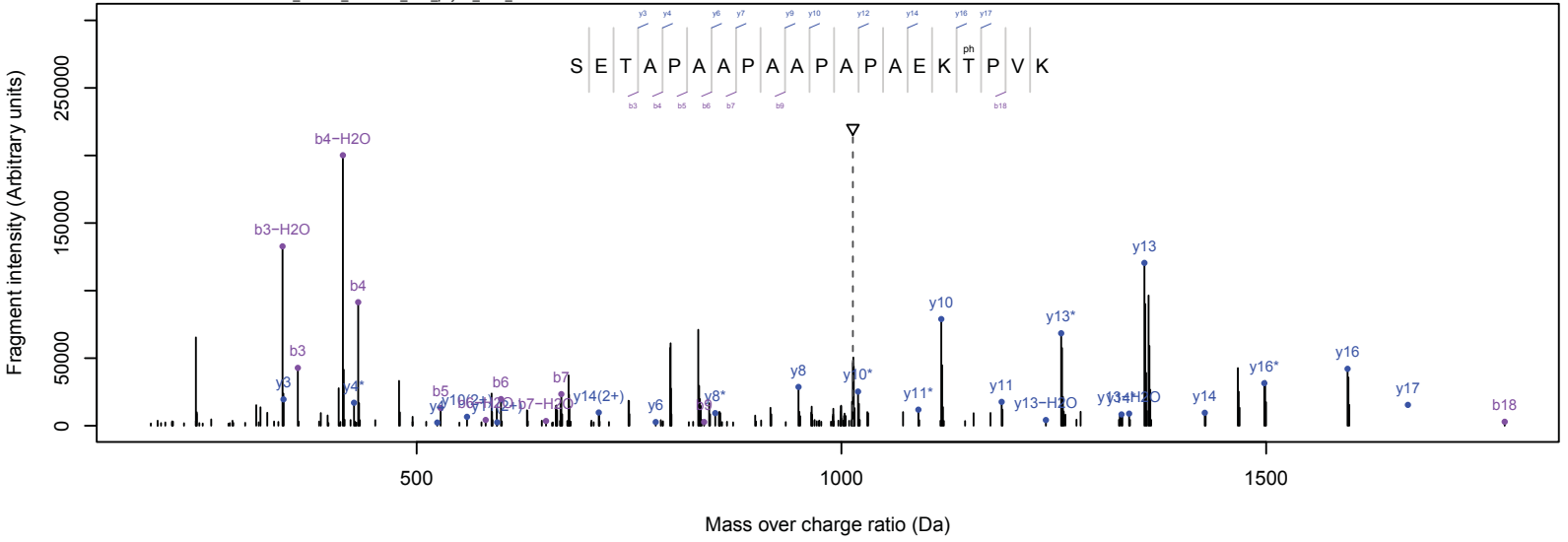
Andromeda score : 88.282



(ac)SETAPAAPAAPAEK(ph)PVK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 19415

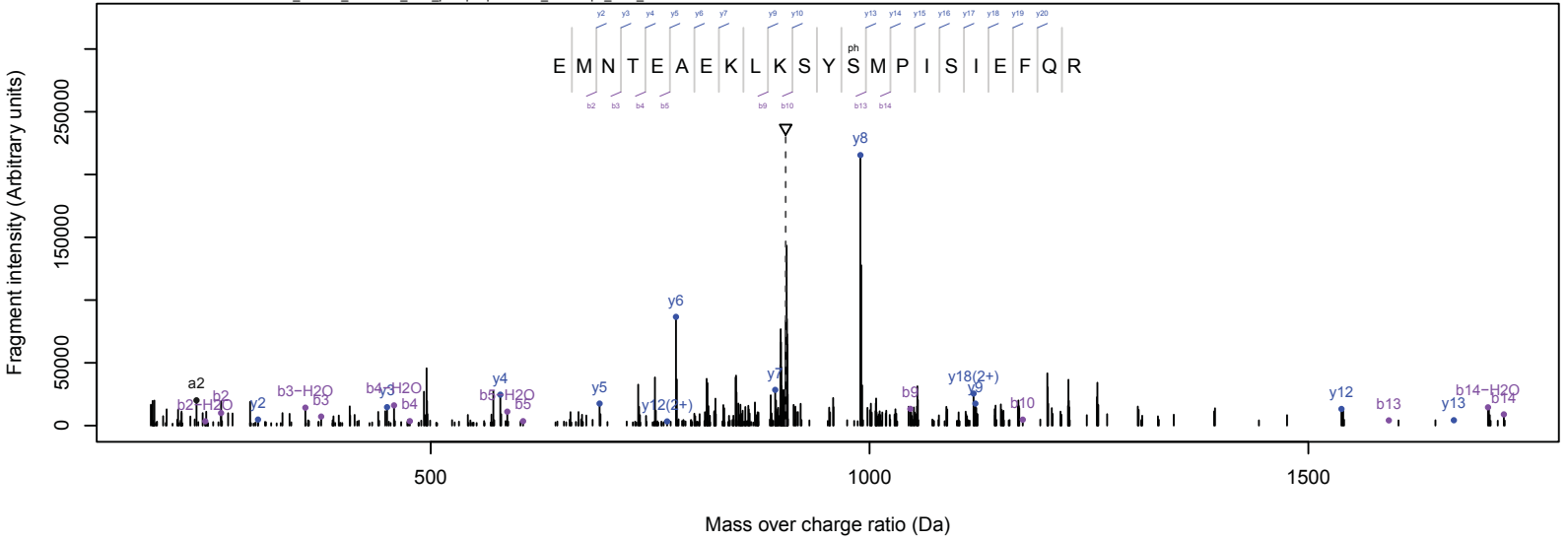
Andromeda score : 88.244



EMNTEAEKLSYS(ph)MPISIEFQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35622

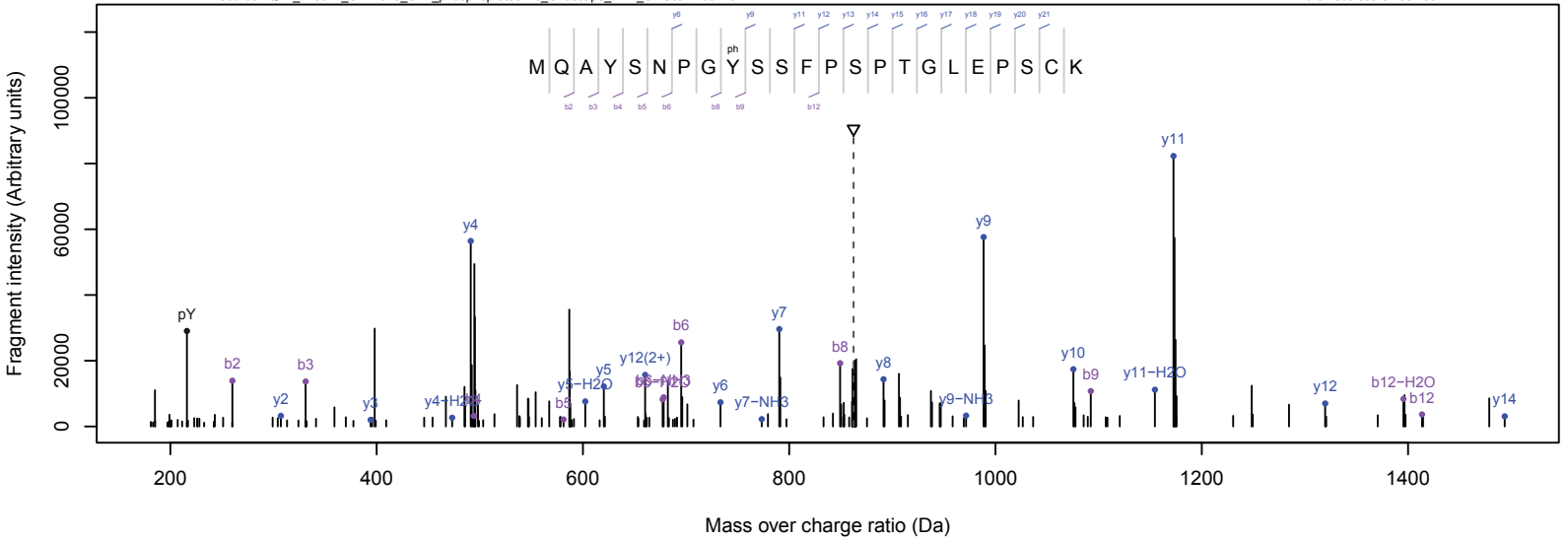
Andromeda score : 88.239



MQAYSNPGYS(ph)SSFPSTGLEPSCK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33446

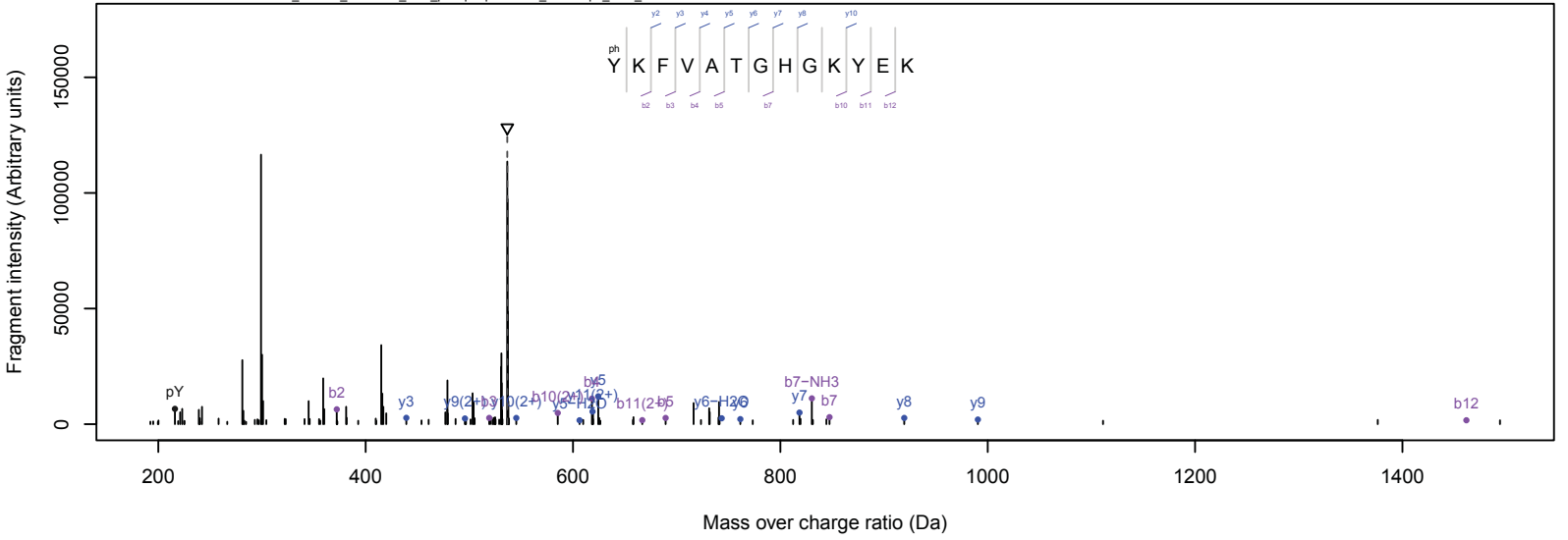
Andromeda score : 88.239



Y(ph)KVFATGHGKYEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 6281

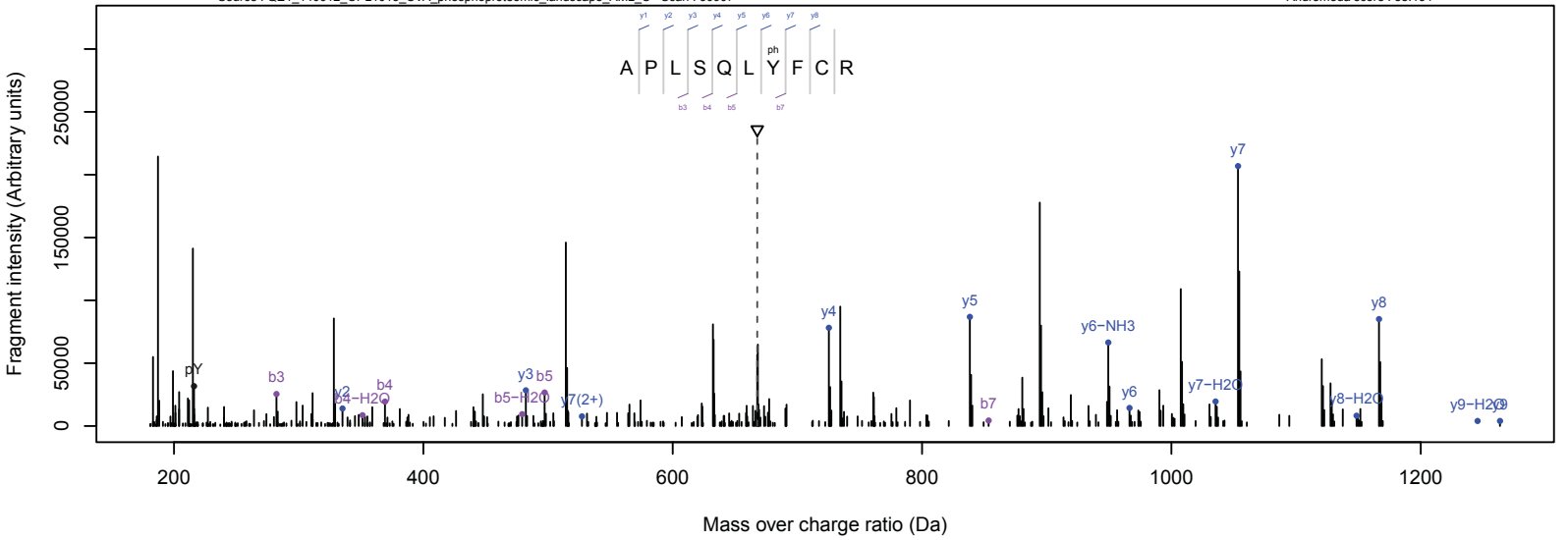
Andromeda score : 88.163



APLSQLY(ph)FCR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30907

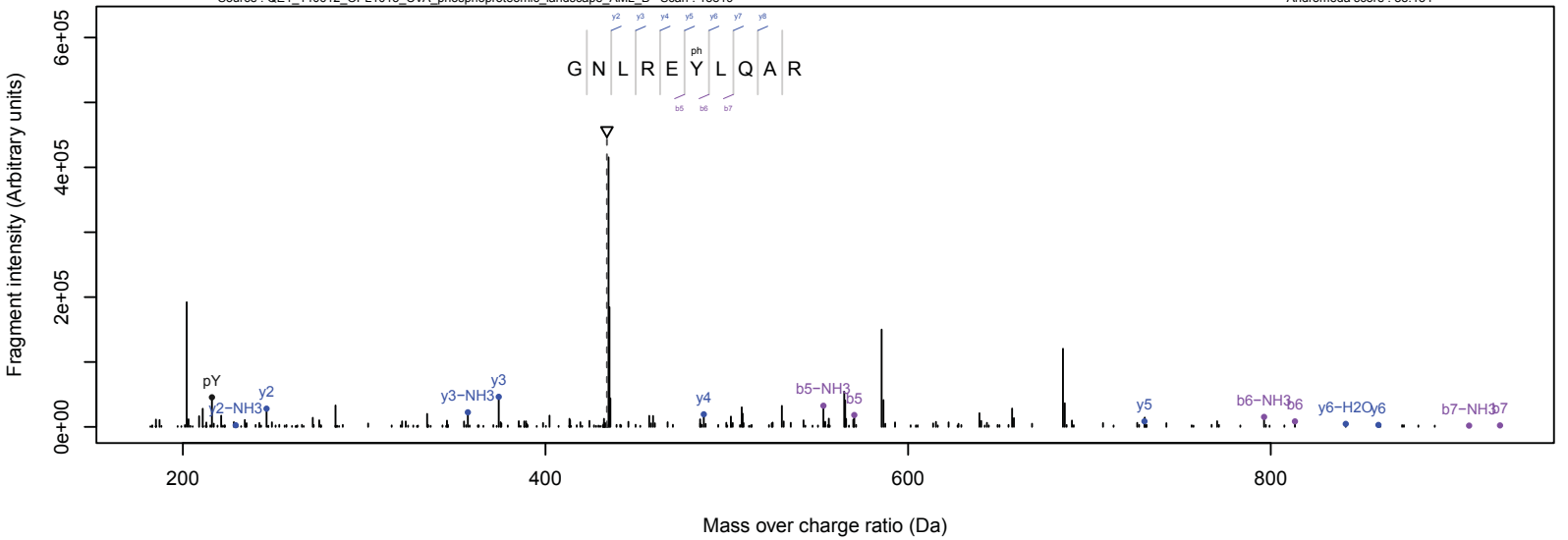
Andromeda score : 88.134



GNLREY(ph)LQAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15819

Andromeda score : 88.134

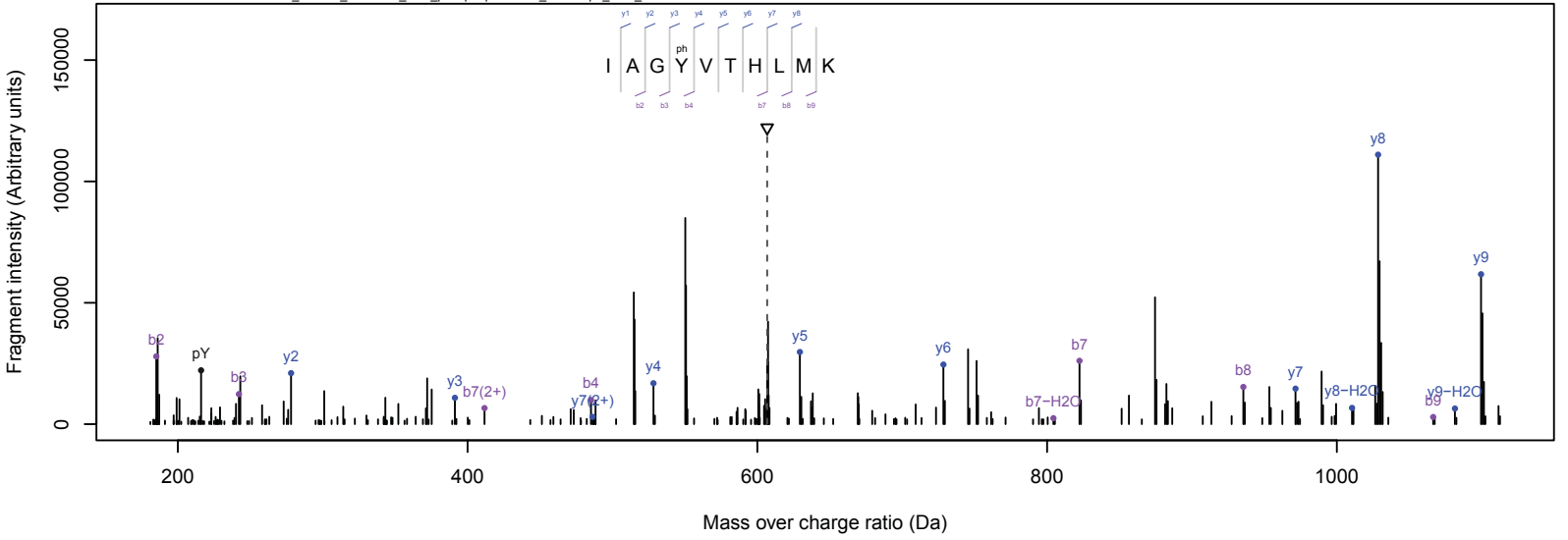




IAGY(ph)VTHLMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 19059

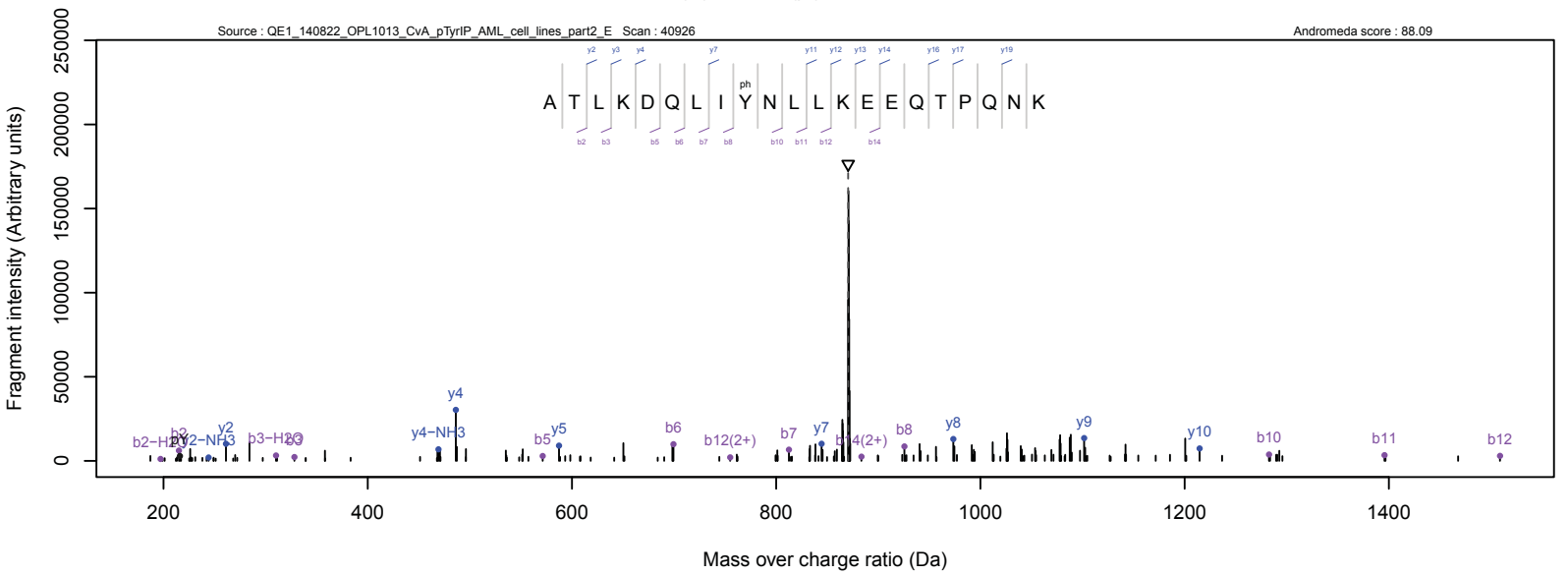
Andromeda score : 88.134



(ac)ATLKDQLIY(ph)NLLKEEQTPQNK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 40926

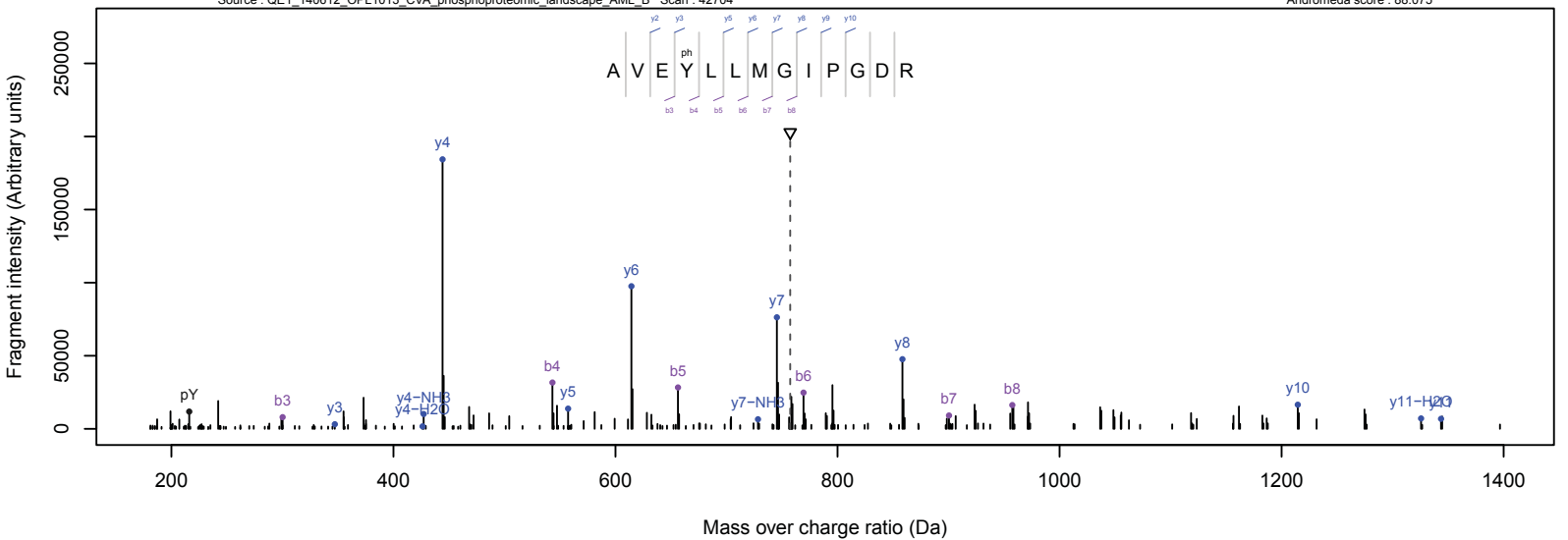
Andromeda score : 88.09



AVEY(ph)LLMGIPGDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42704

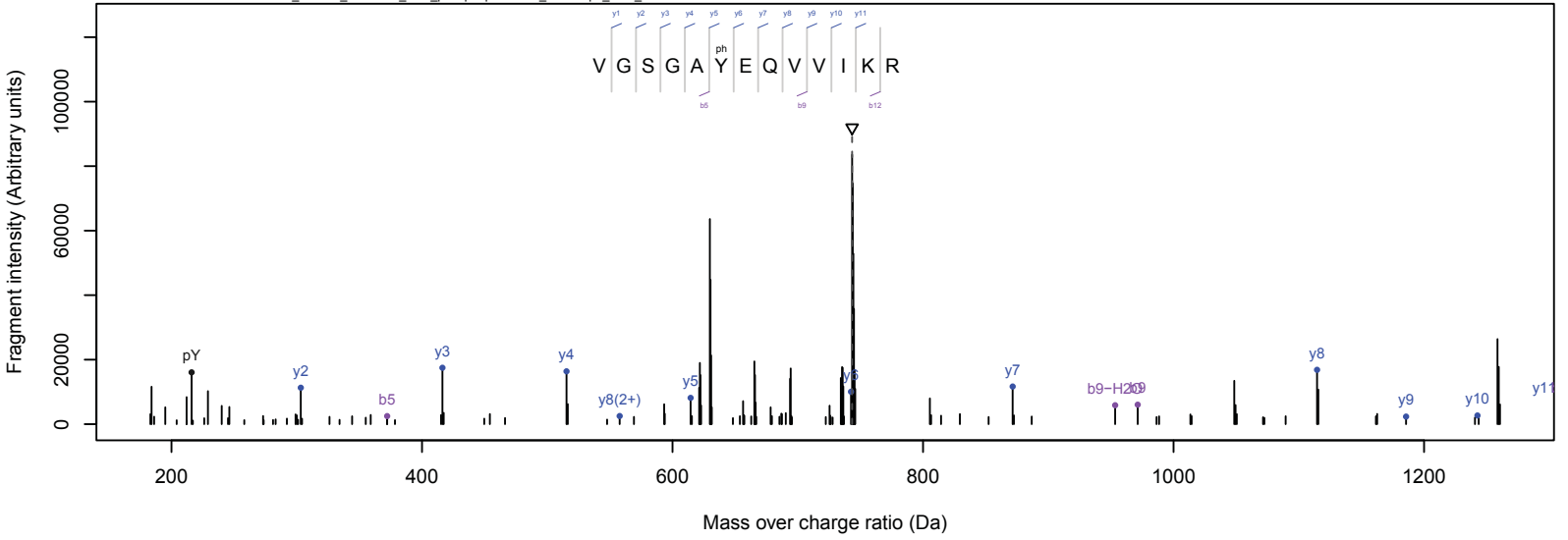
Andromeda score : 88.075



VGSGAY(ph)EQVVIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13513

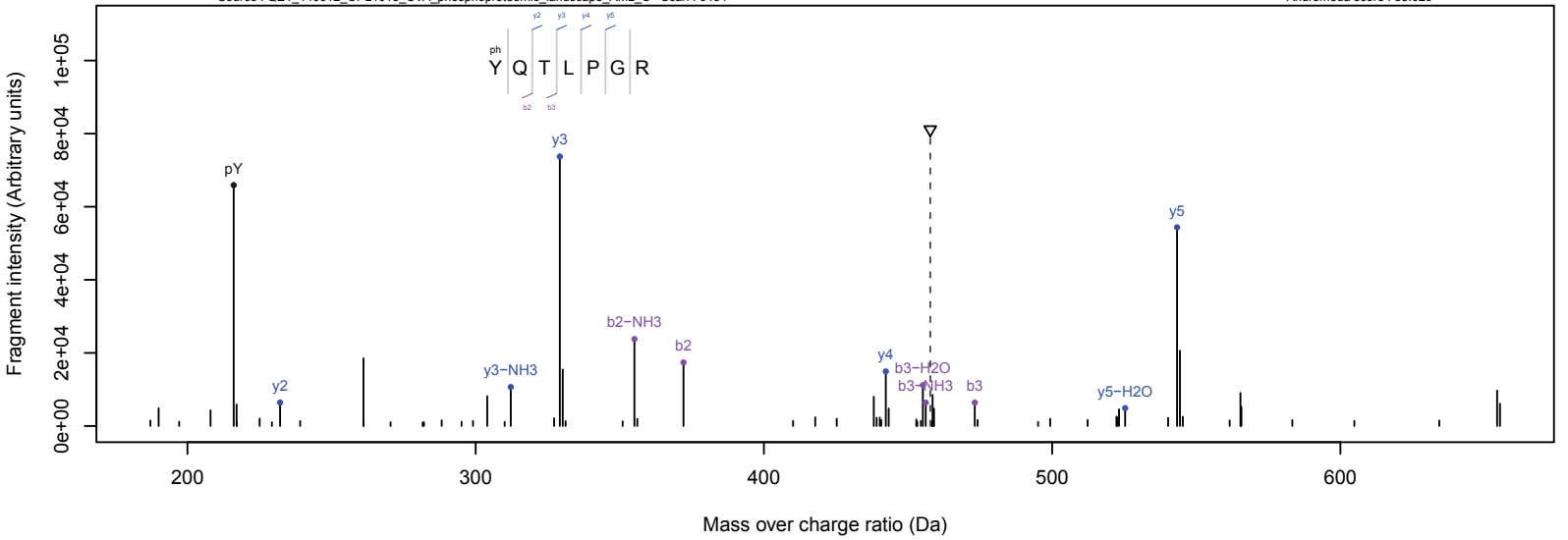
Andromeda score : 88.075



Y(ph)QTLPLGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6434

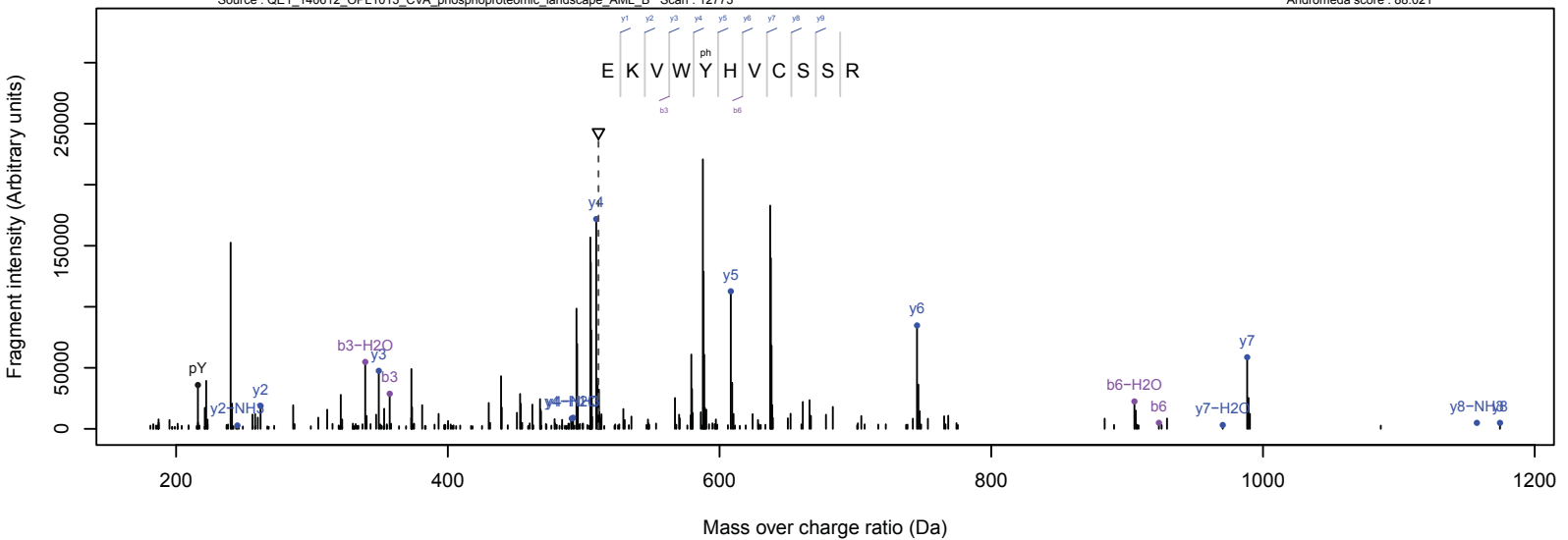
Andromeda score : 88.029



EKVWY(ph)HVCSSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12773

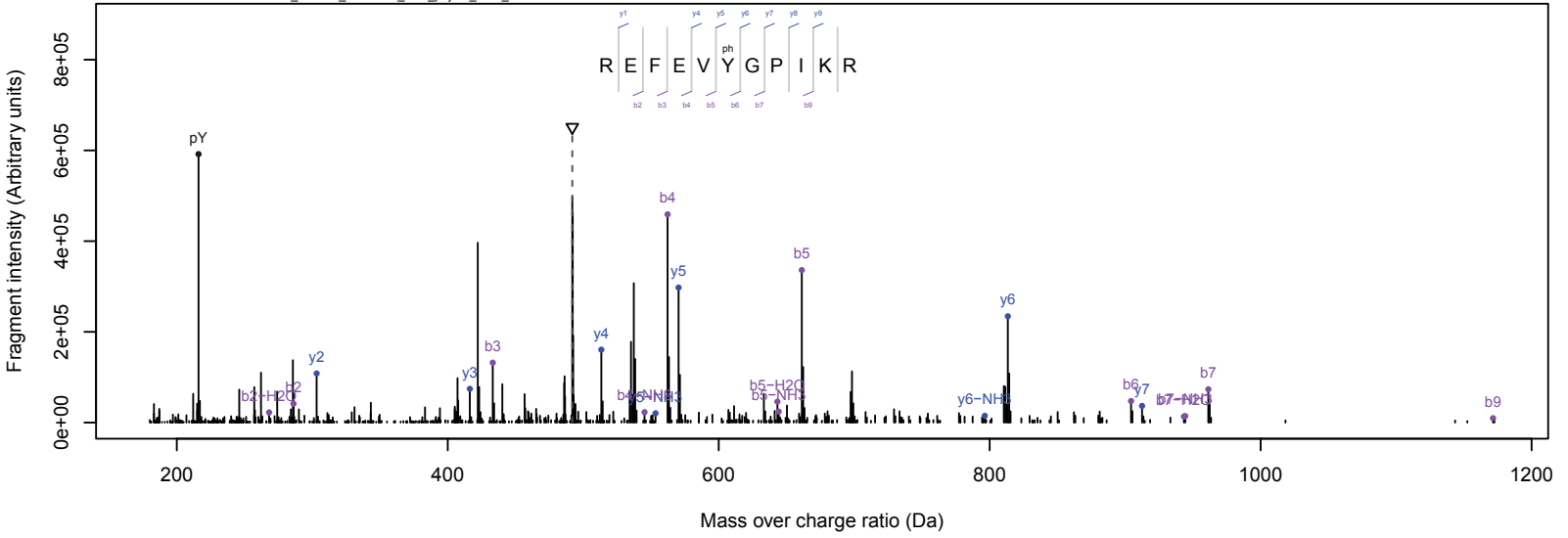
Andromeda score : 88.021



REFEVY(ph)GPIKR-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan: 18786

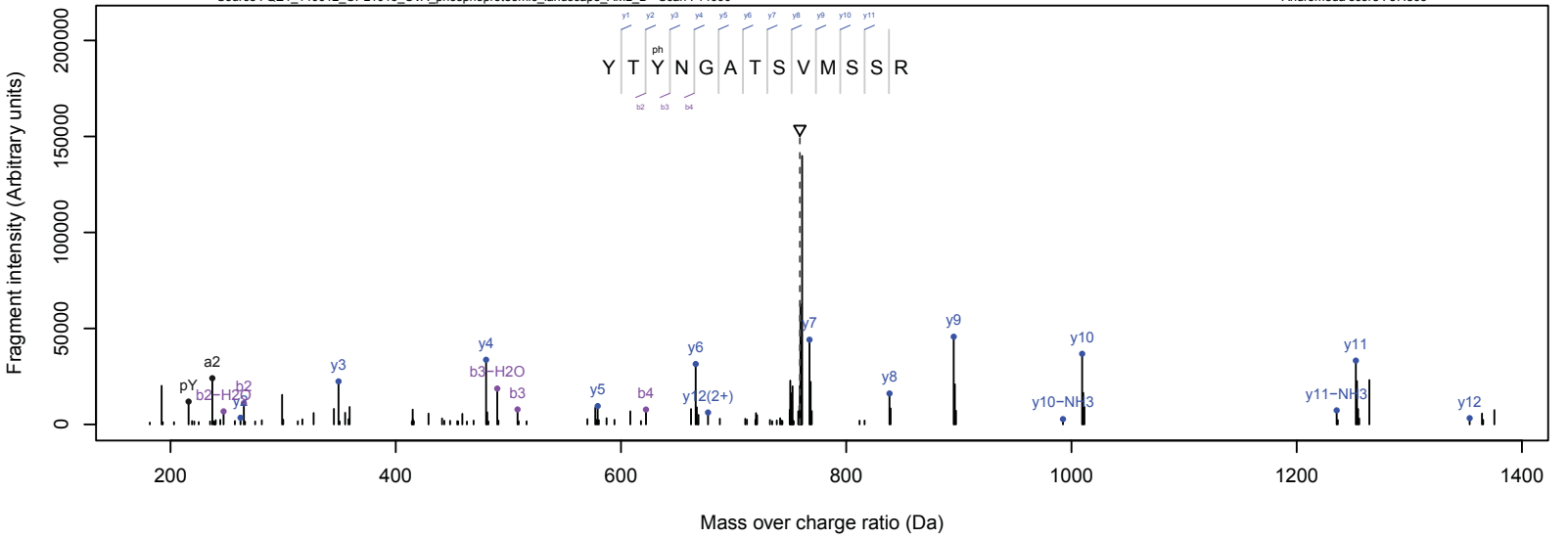
Andromeda score: 88.021



YTY(ph)NGATSVMSR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 14090

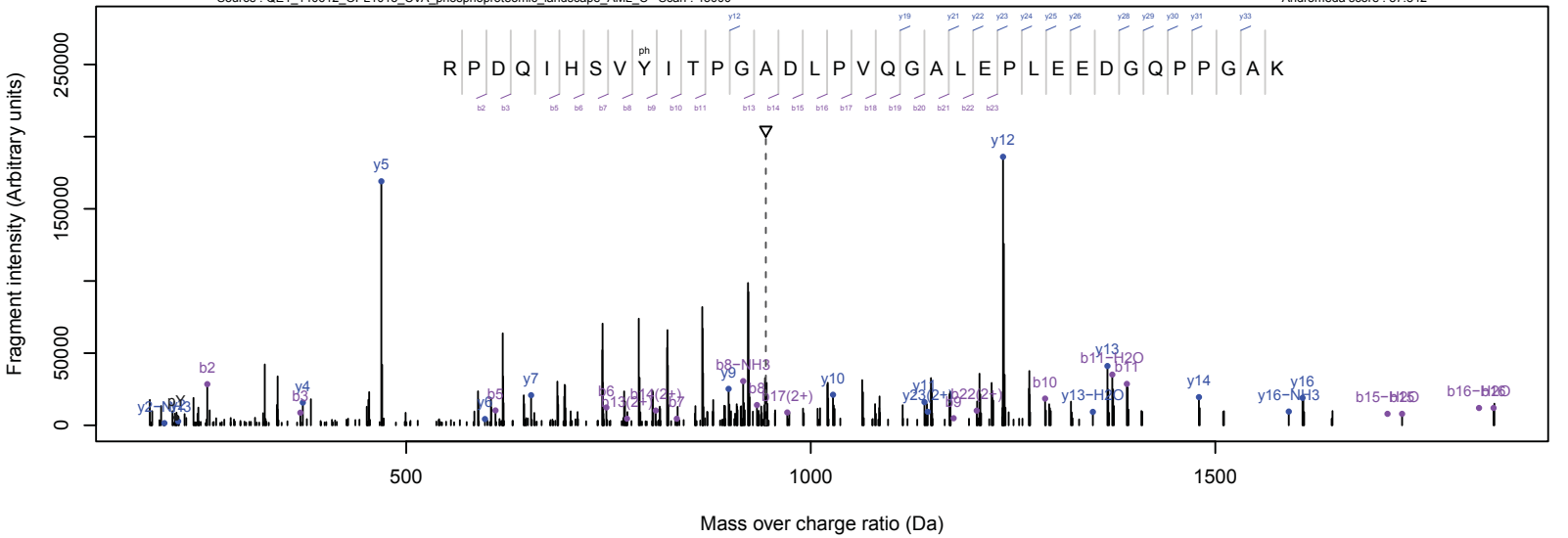
Andromeda score: 87.863



RPDQIHSVY(ph)ITPGADLPVQGALEPLEEDGQPPGAK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 43000

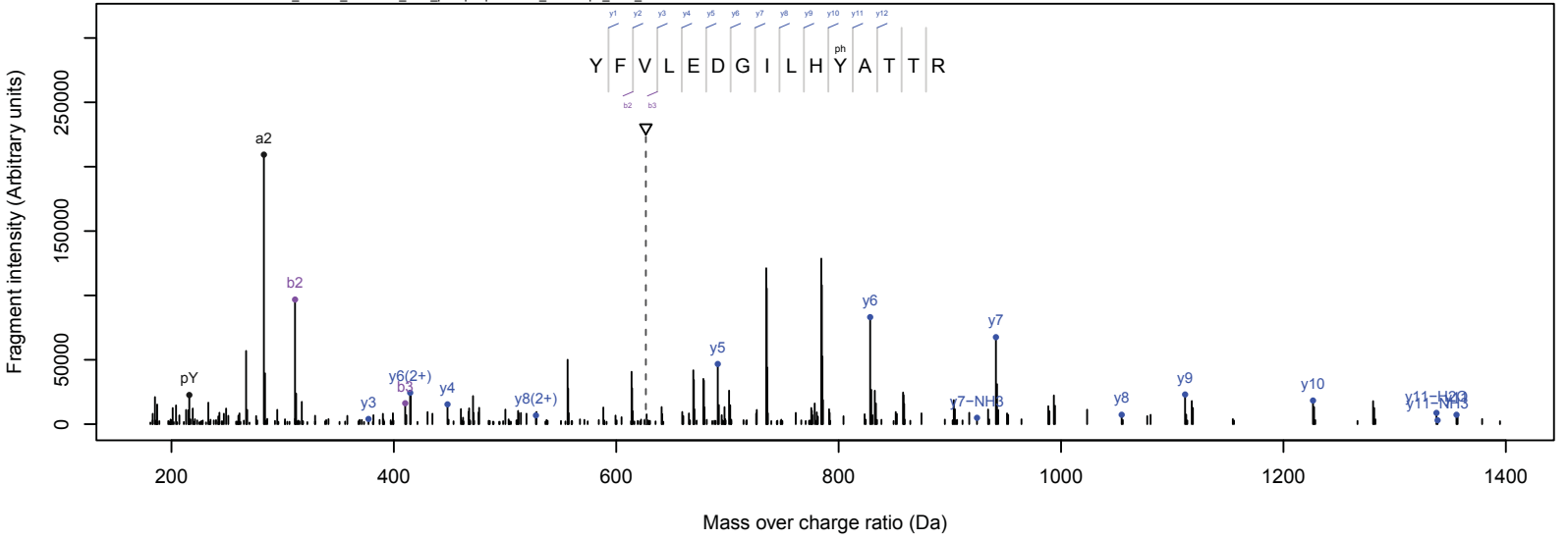
Andromeda score: 87.842



**YFVLEDGILHY(ph)ATTR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42874

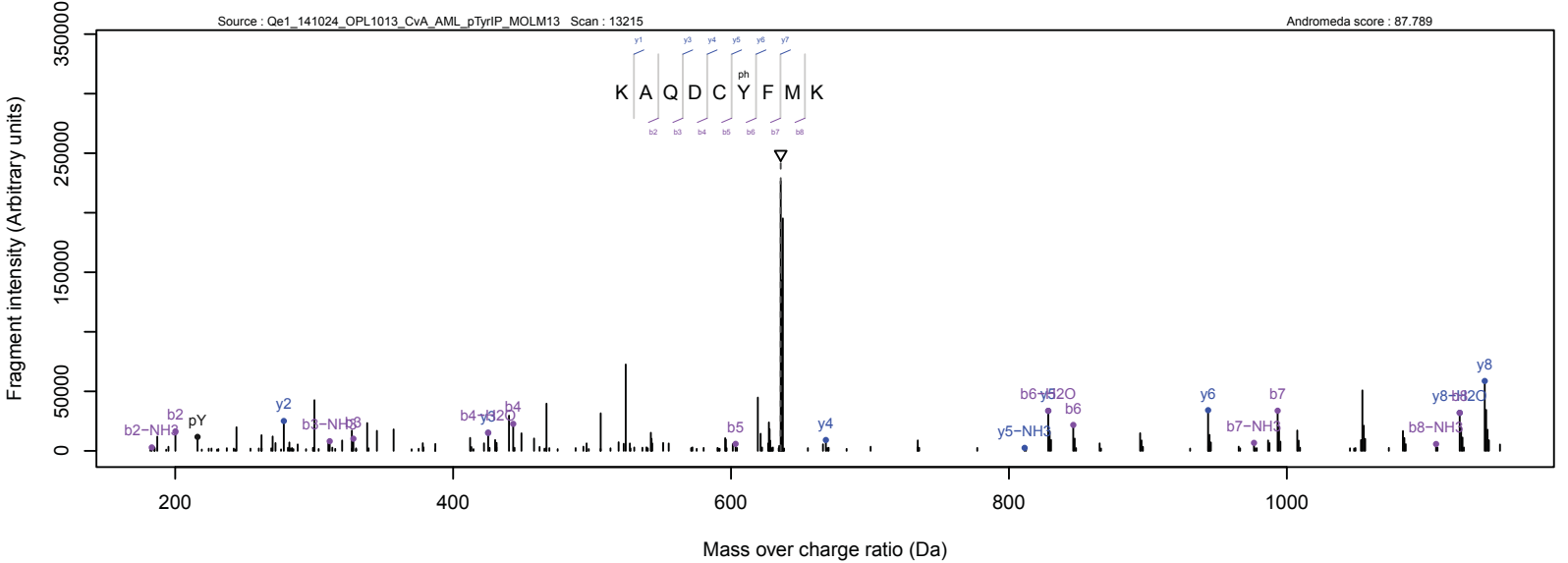
Andromeda score : 87.824



**KAQDCY(ph)FMK-2+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 13215

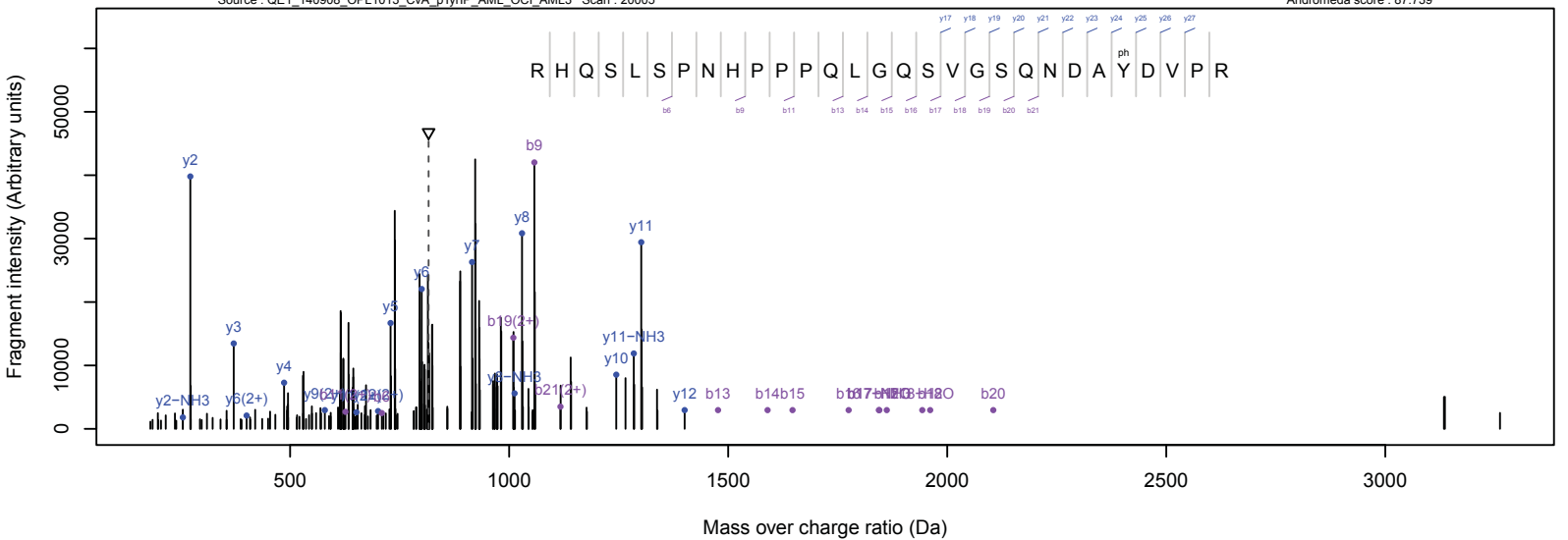
Andromeda score : 87.789



**RHQSLSPNHPPQLGQSVGSQNDAY(ph)DVPR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 20005

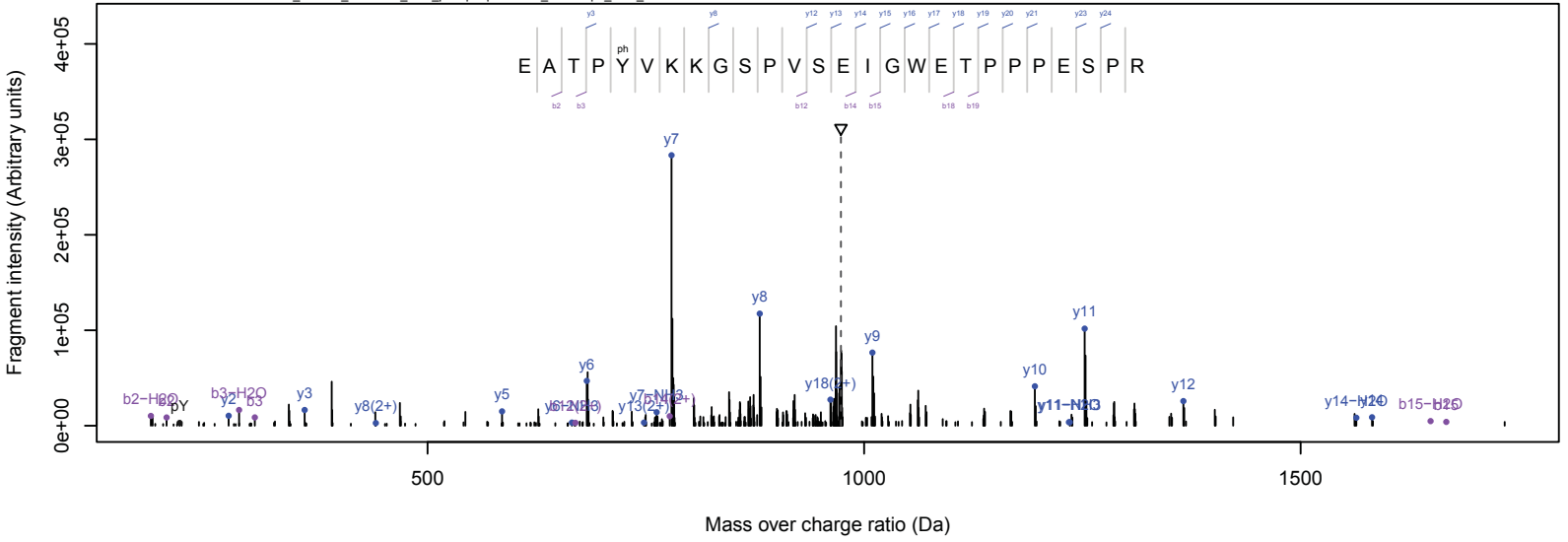
Andromeda score : 87.739



EATPY(ph)VKKGSPVSEIGWETPPPEspr-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28638

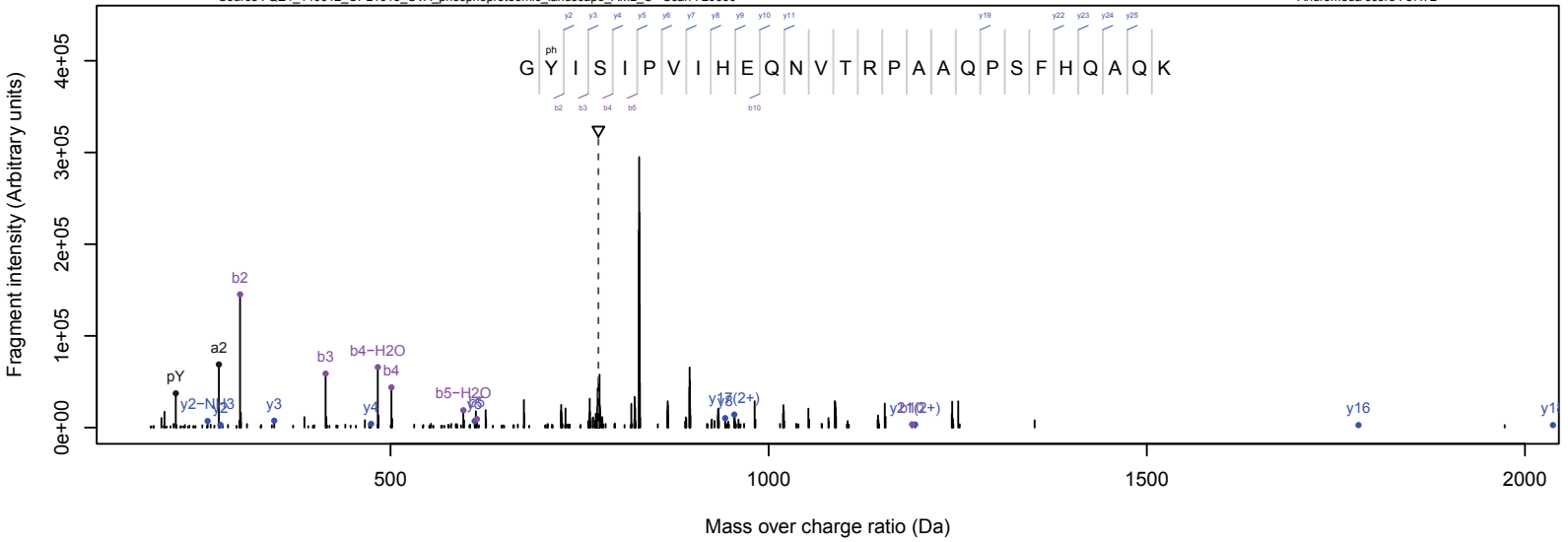
Andromeda score : 87.72



GY(ph)ISIPVIHEQNVTRPAAQPSFHQAQK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29650

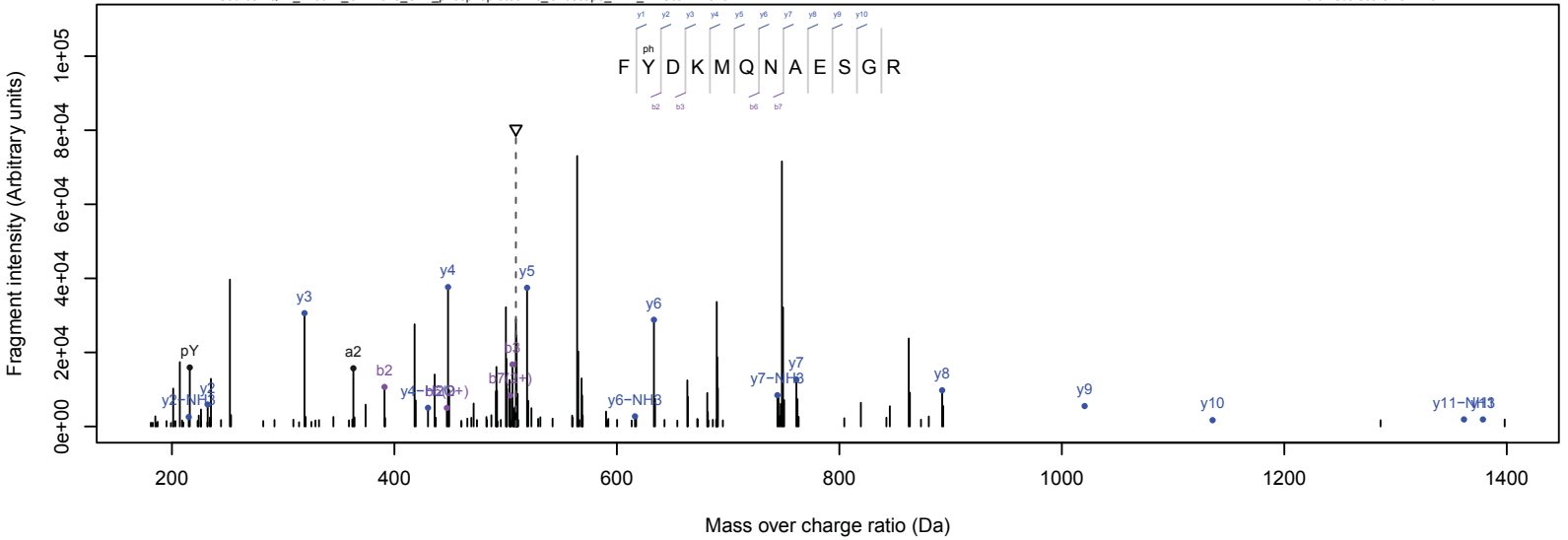
Andromeda score : 87.72



FY(ph)DKMQNAESGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7348

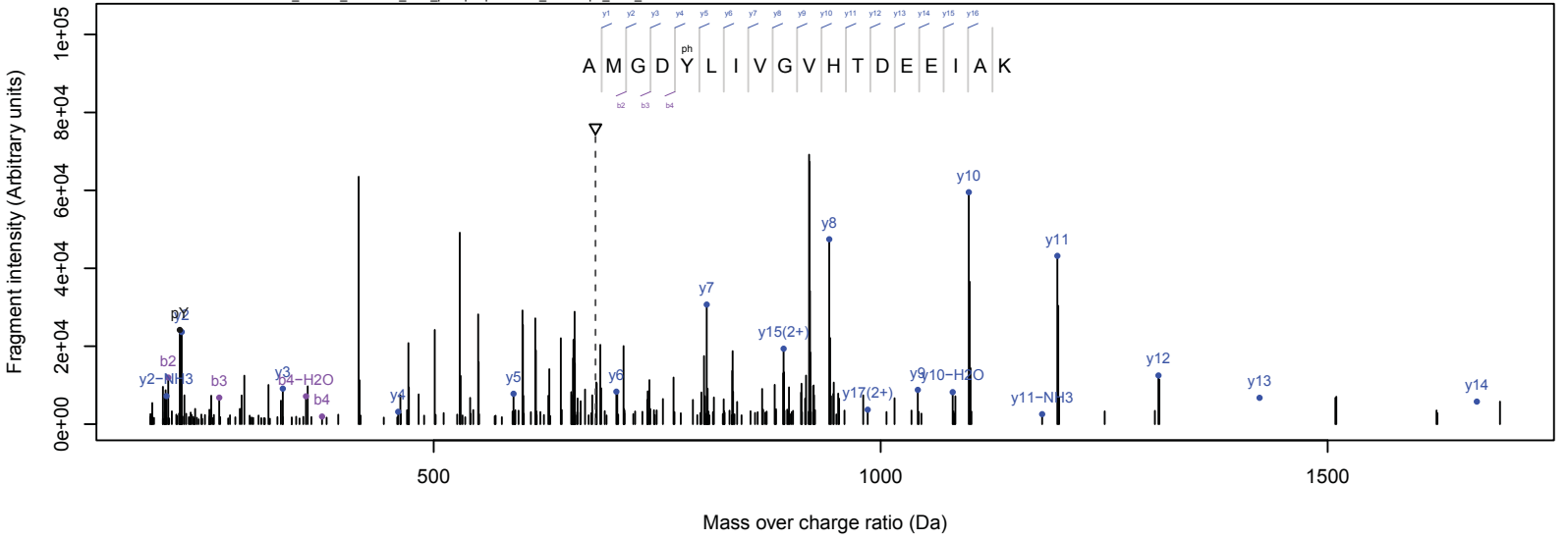
Andromeda score : 87.719



AMGDY(ph)LIVGVHTDEEIAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 47888

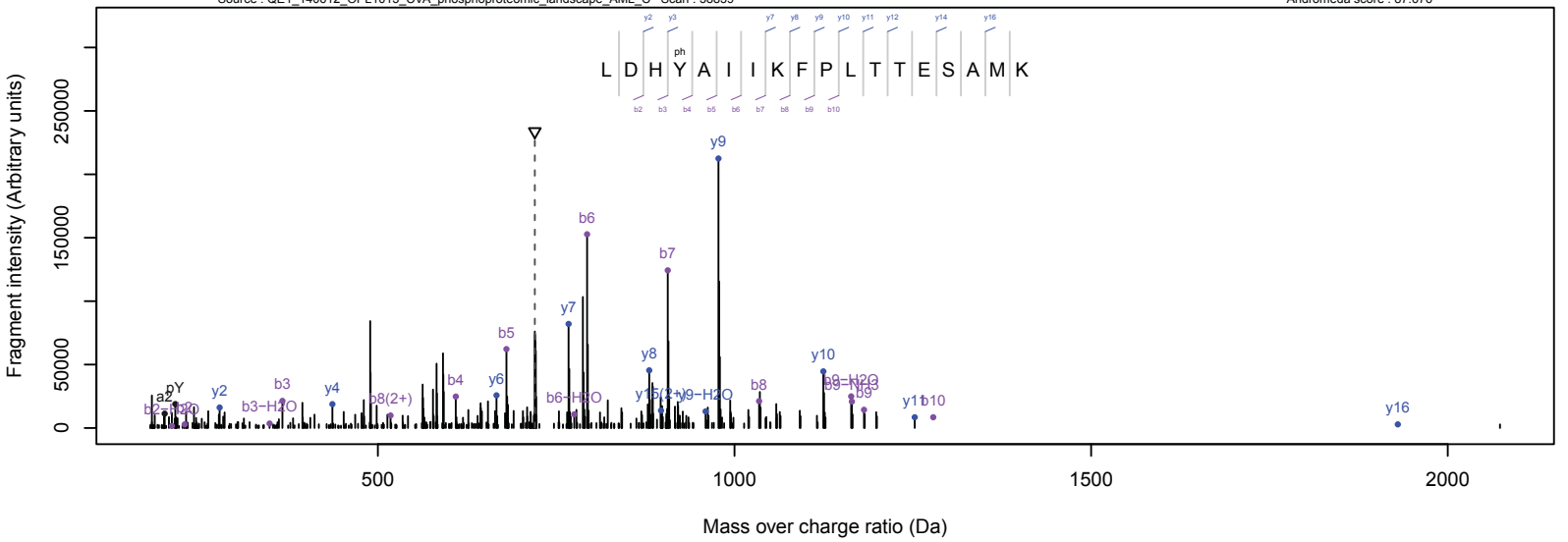
Andromeda score : 87.676



LDHY(ph)AIKFPLTTESAMK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38839

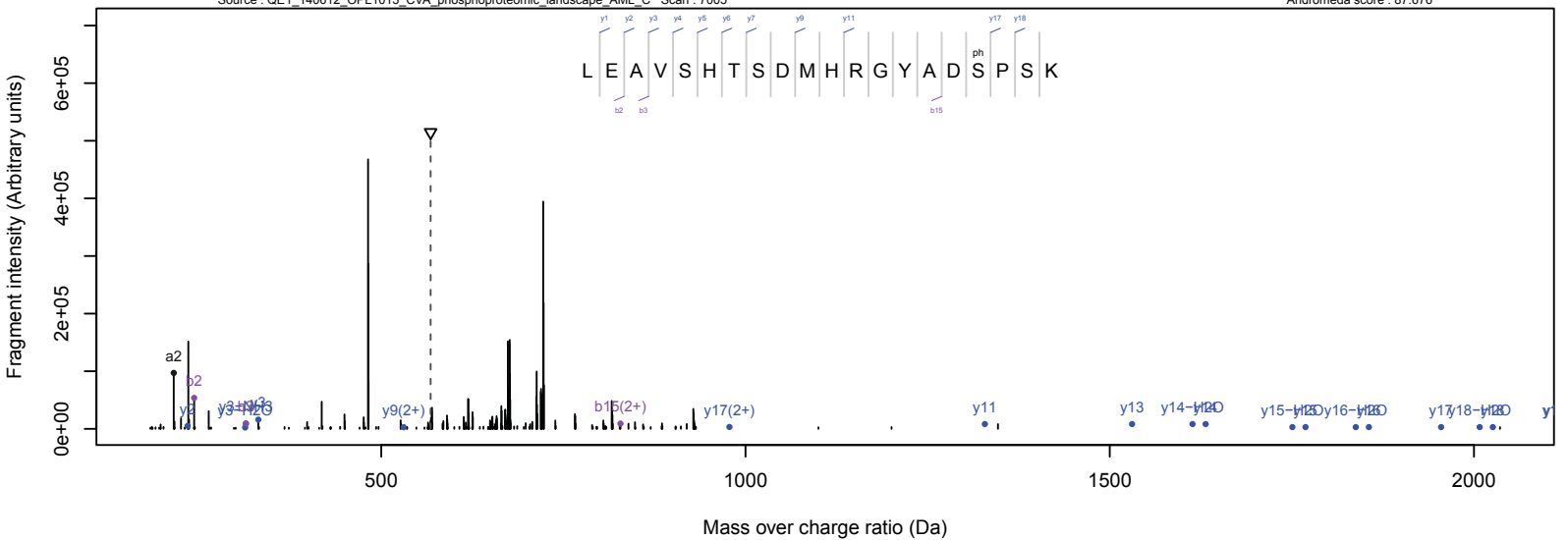
Andromeda score : 87.676



LEAVSHTSDMHRGYADSPSK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7005

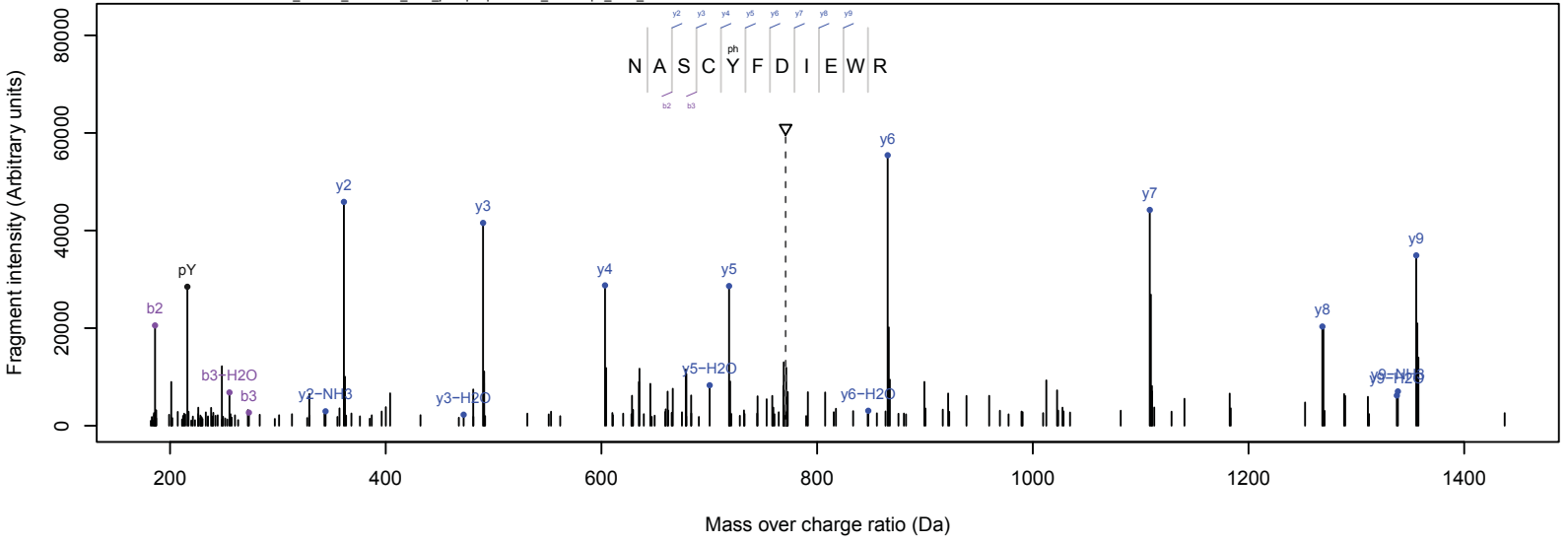
Andromeda score : 87.676



NASCY(ph)FDIEWR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39296

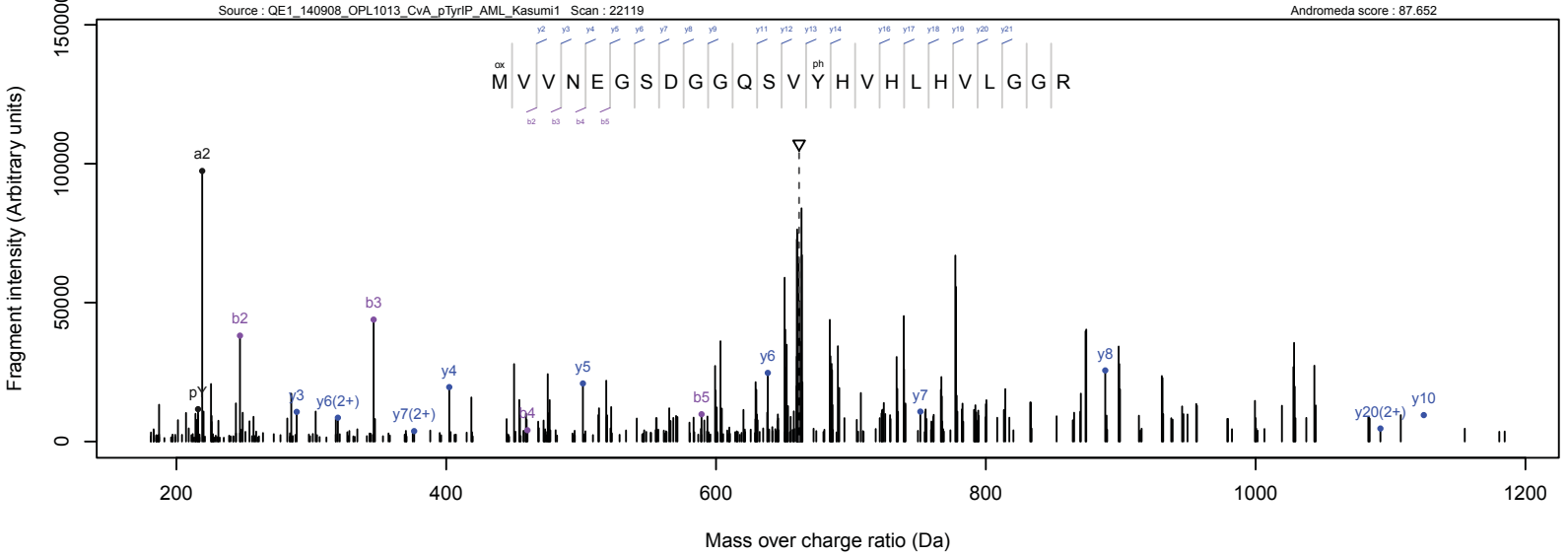
Andromeda score : 87.667



M(ox)VVNEGSDDGGQSVY(ph)HVLHLVLR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 22119

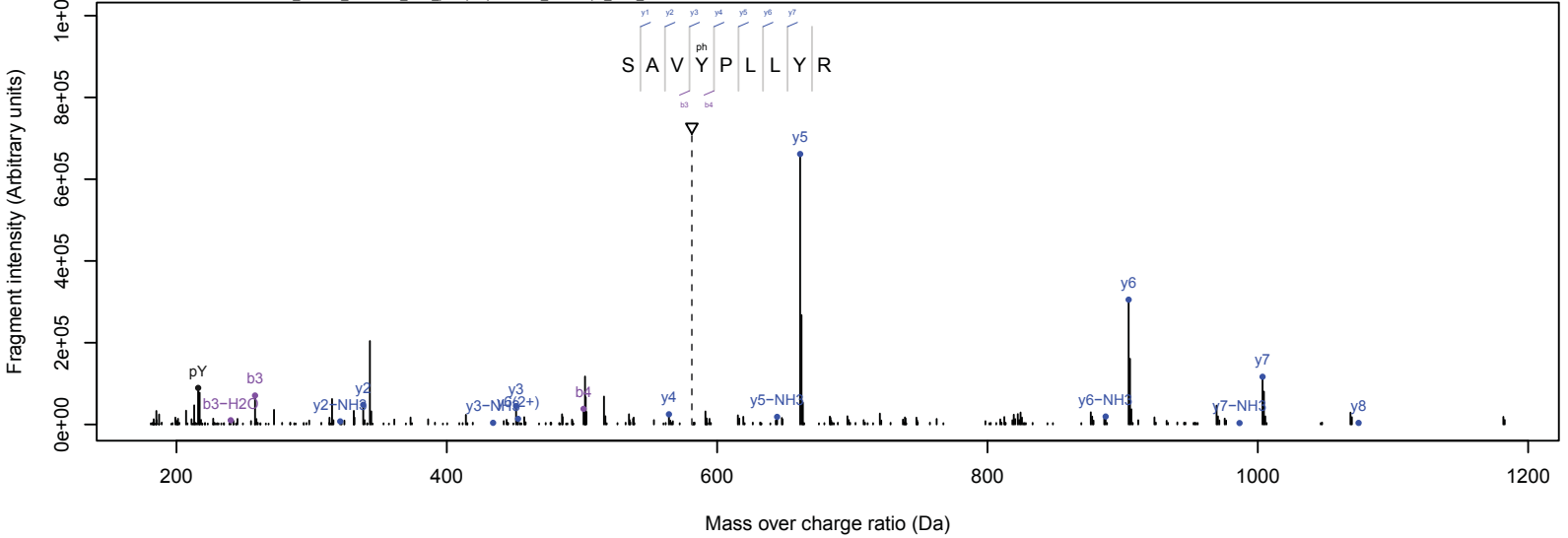
Andromeda score : 87.652



SAVY(ph)PLLYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28568

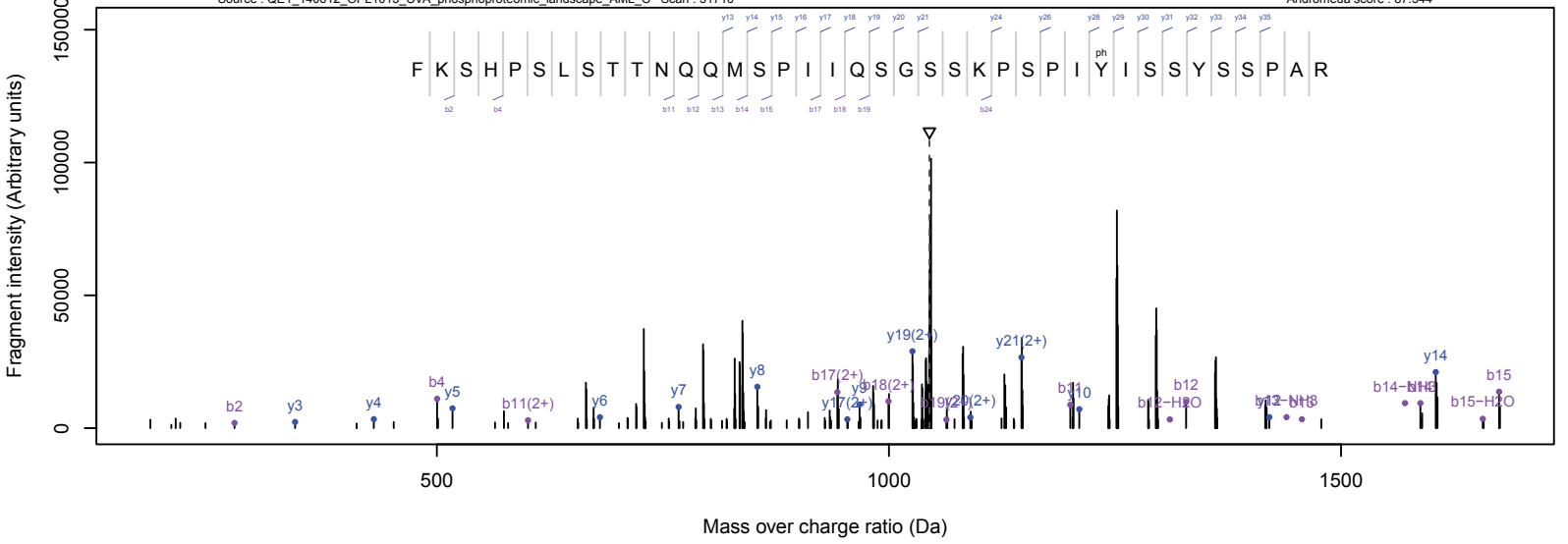
Andromeda score : 87.639



**FKSHPSLSTTNQQMSPIIQSGSSKPSPIYISSYSSPAR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31710

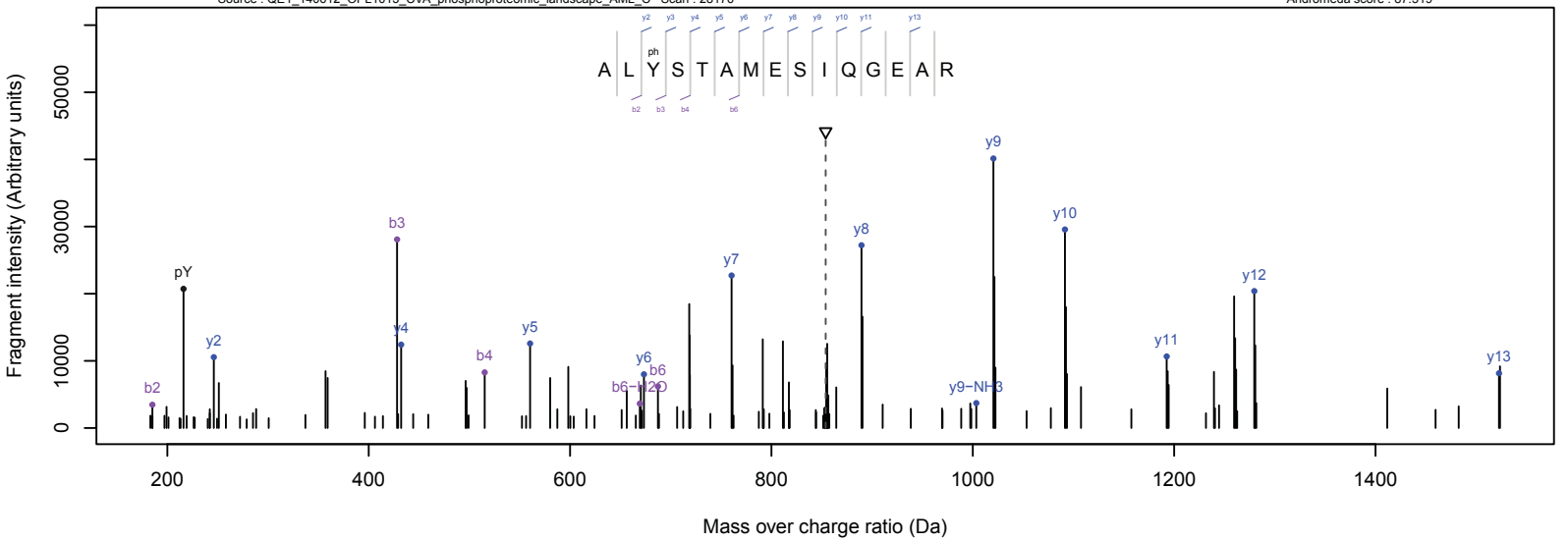
Andromeda score : 87.544



**ALY(ph)STAMESIQGEAR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28176

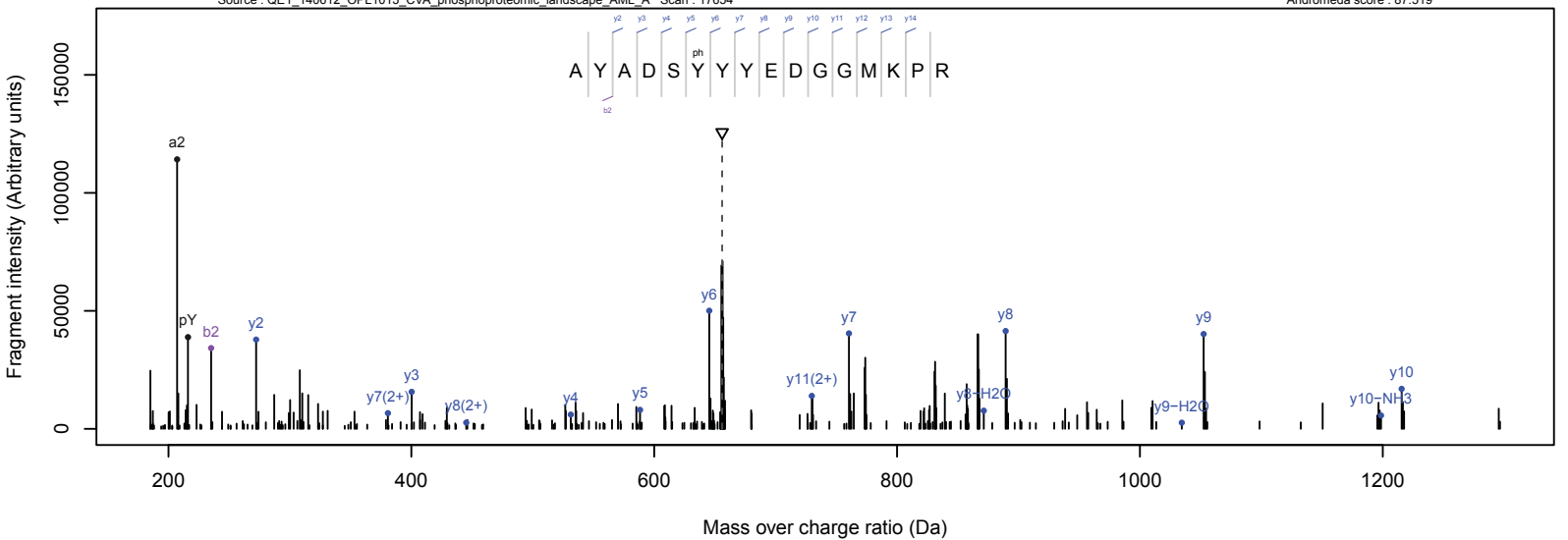
Andromeda score : 87.519



**AYADSY(ph)YYEDGGMKPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17654

Andromeda score : 87.519

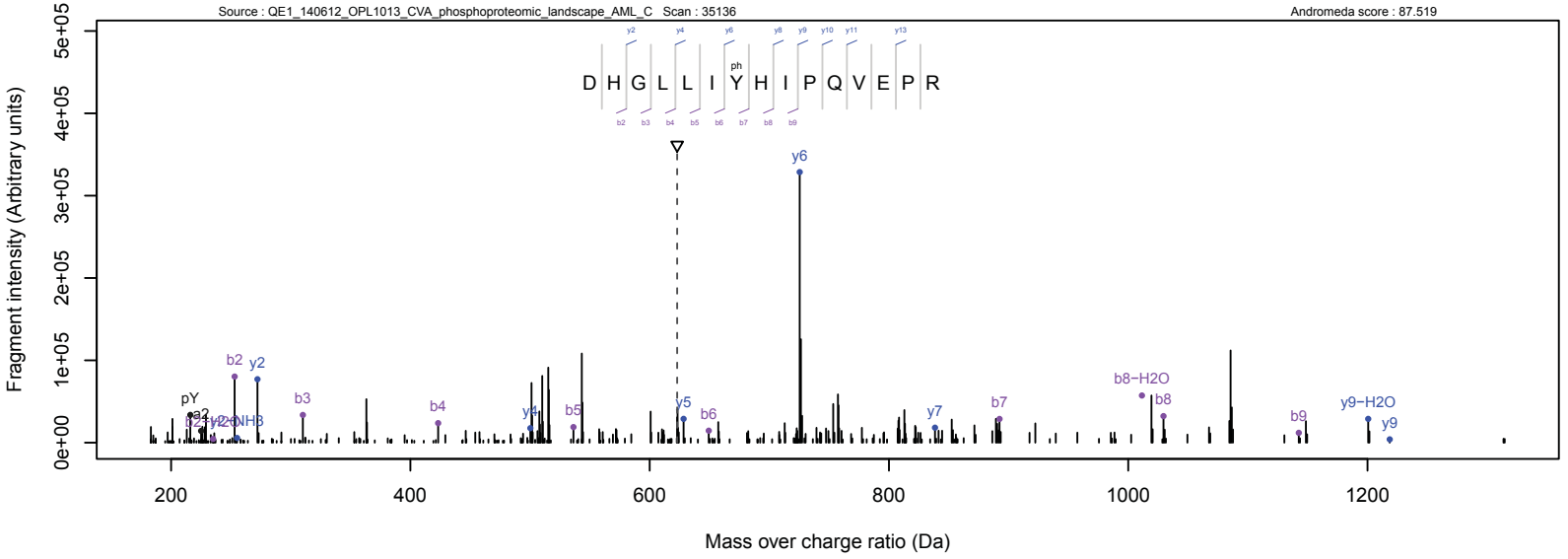




DHGLLIY(ph)HIPQVEPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 35136

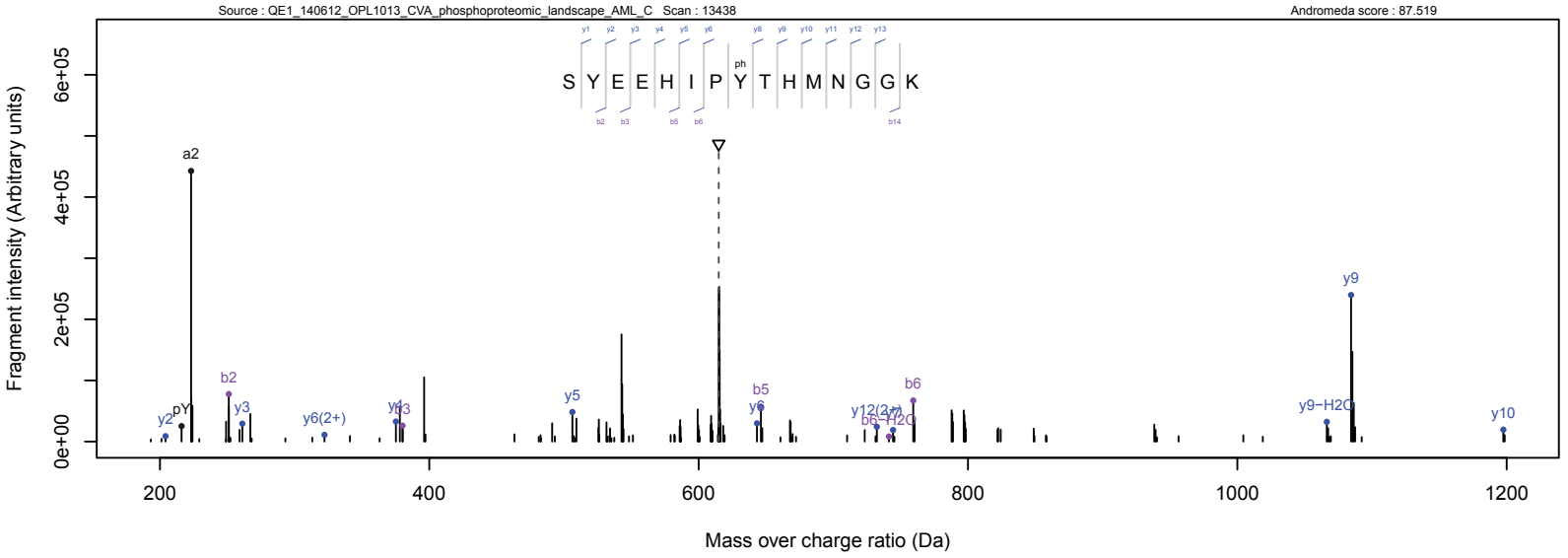
Andromeda score: 87.519



SYEEHIPY(ph)THMNGGK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13438

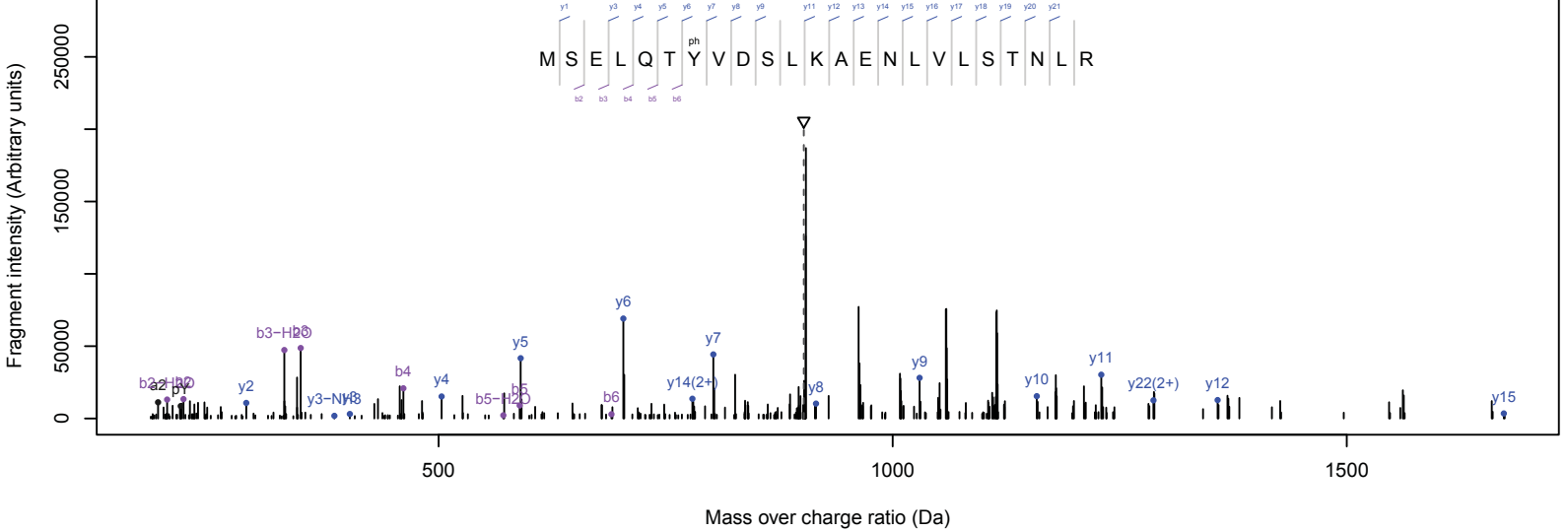
Andromeda score: 87.519



MSELQTY(ph)VDSLKAENLVLSSTNLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 46951

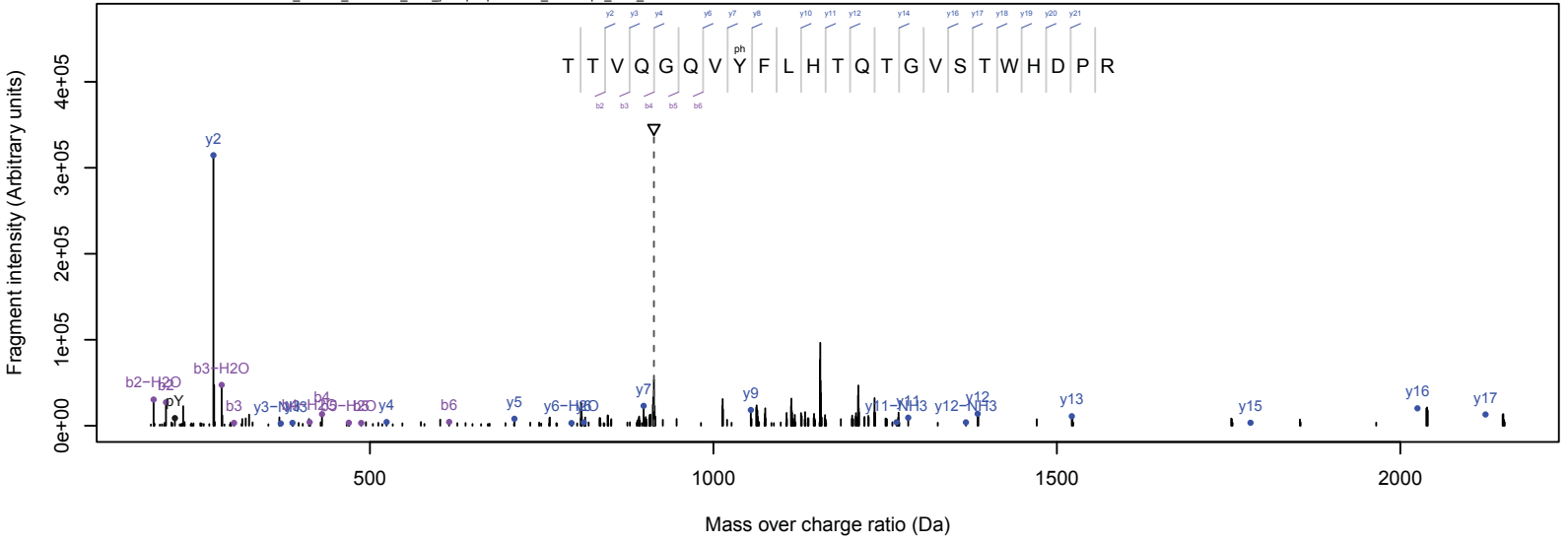
Andromeda score: 87.511



**TTVQGQVY(ph)FLHTQTGVSTWHDPR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31973

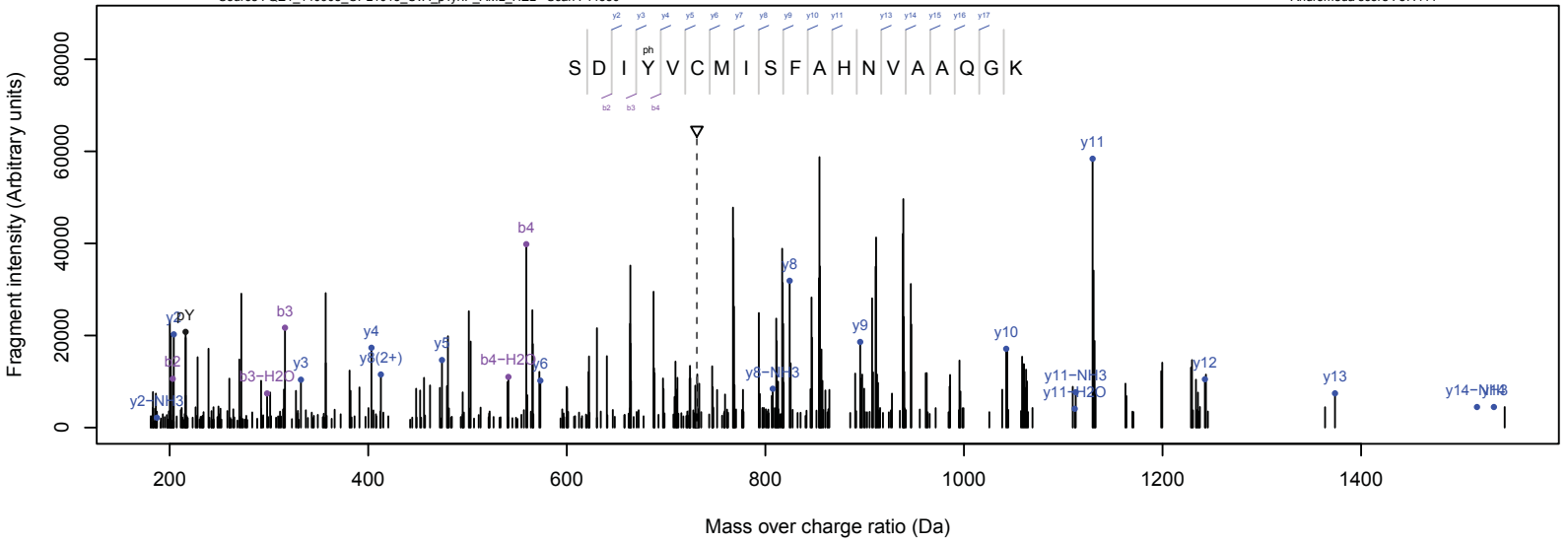
Andromeda score: 87.511



**SDIY(ph)VCMISFAHNVAAGK-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 44350

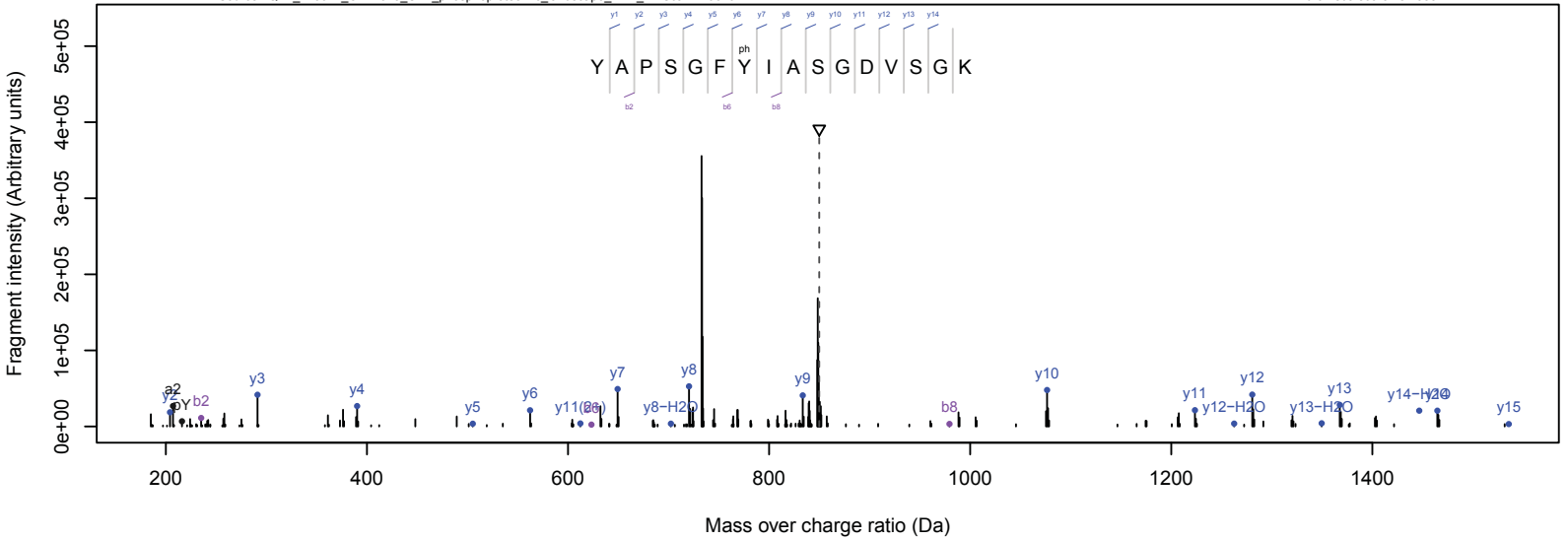
Andromeda score: 87.411



**YAPSGFY(ph)IASGDVSGK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 29629

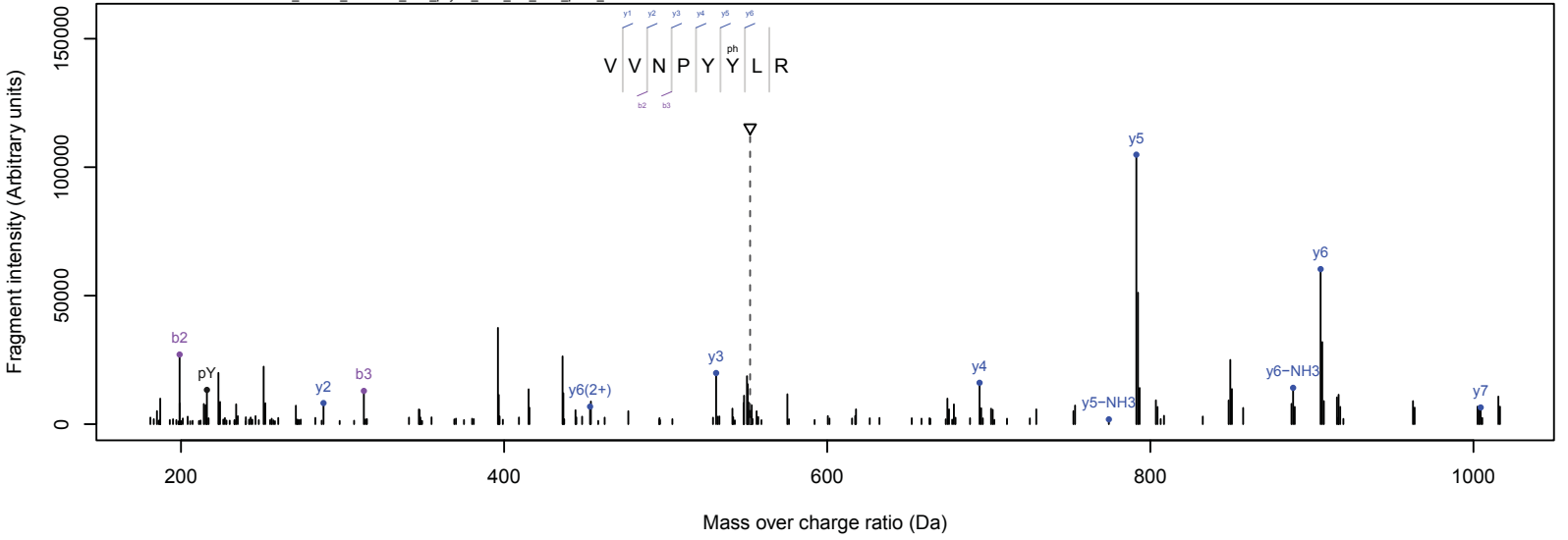
Andromeda score: 87.388



VVNPYY(ph)LR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 16129

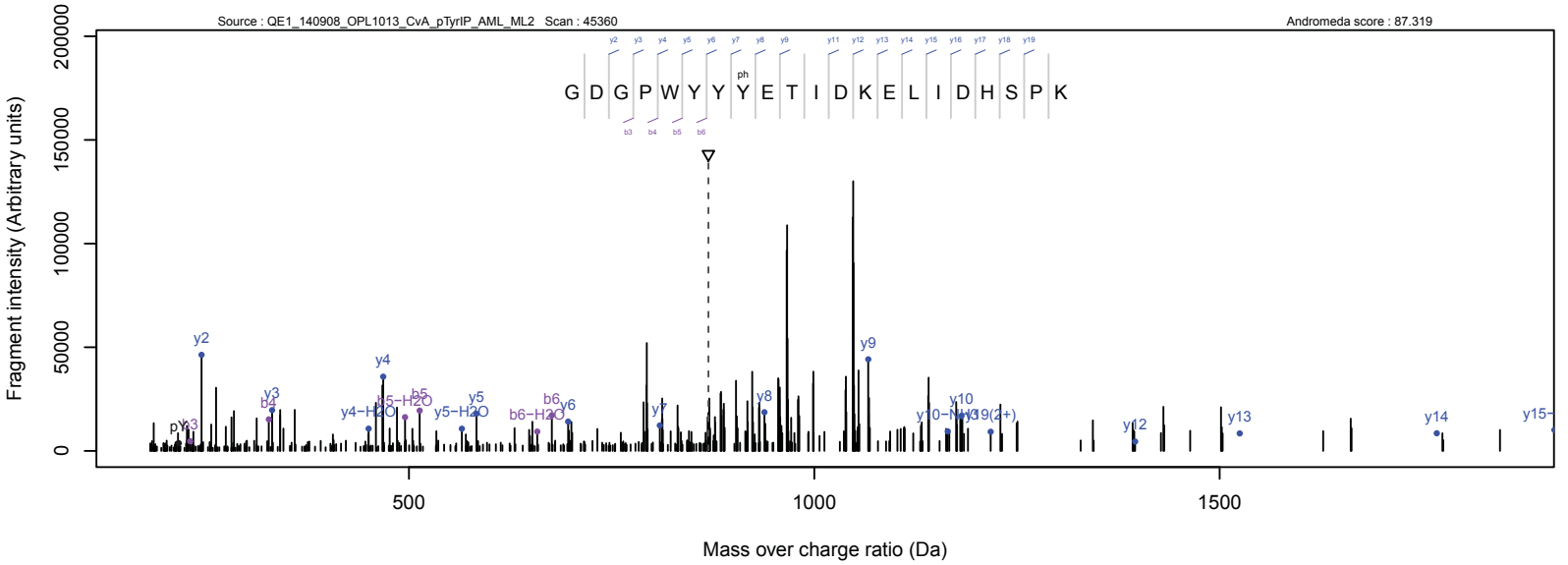
Andromeda score : 87.323



GDGPWYY(ph)ETIDKELIDHSPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 45360

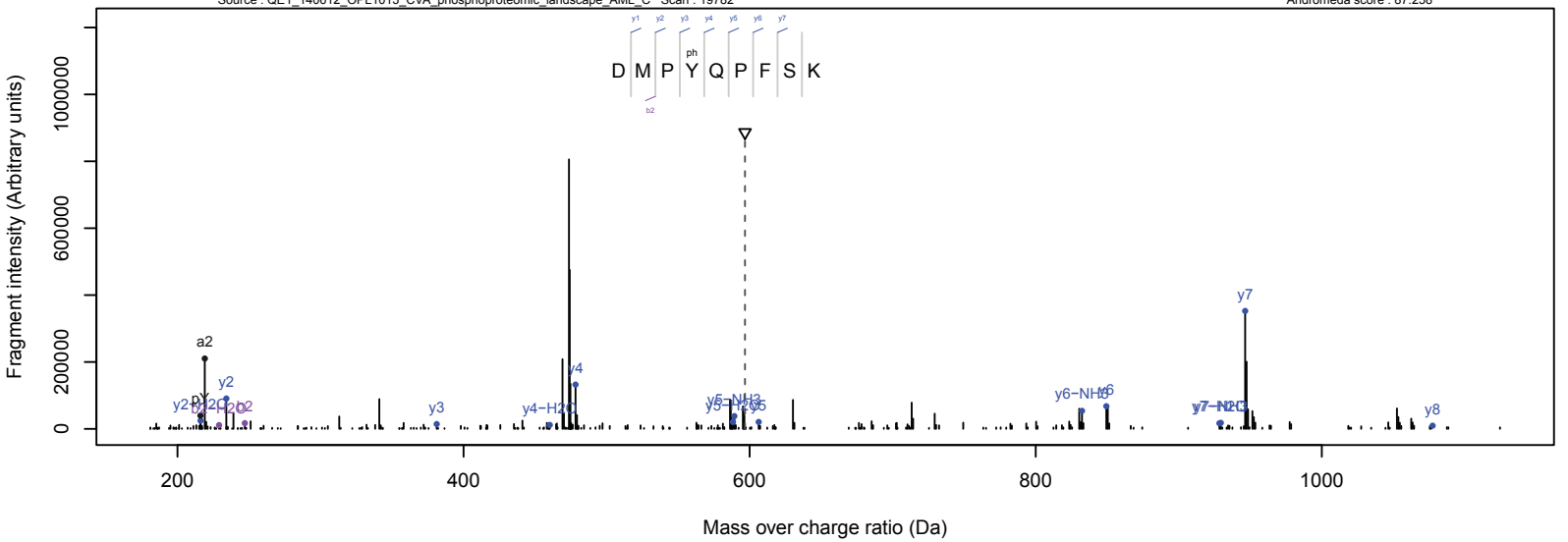
Andromeda score : 87.319



DMPY(ph)QPFISK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19782

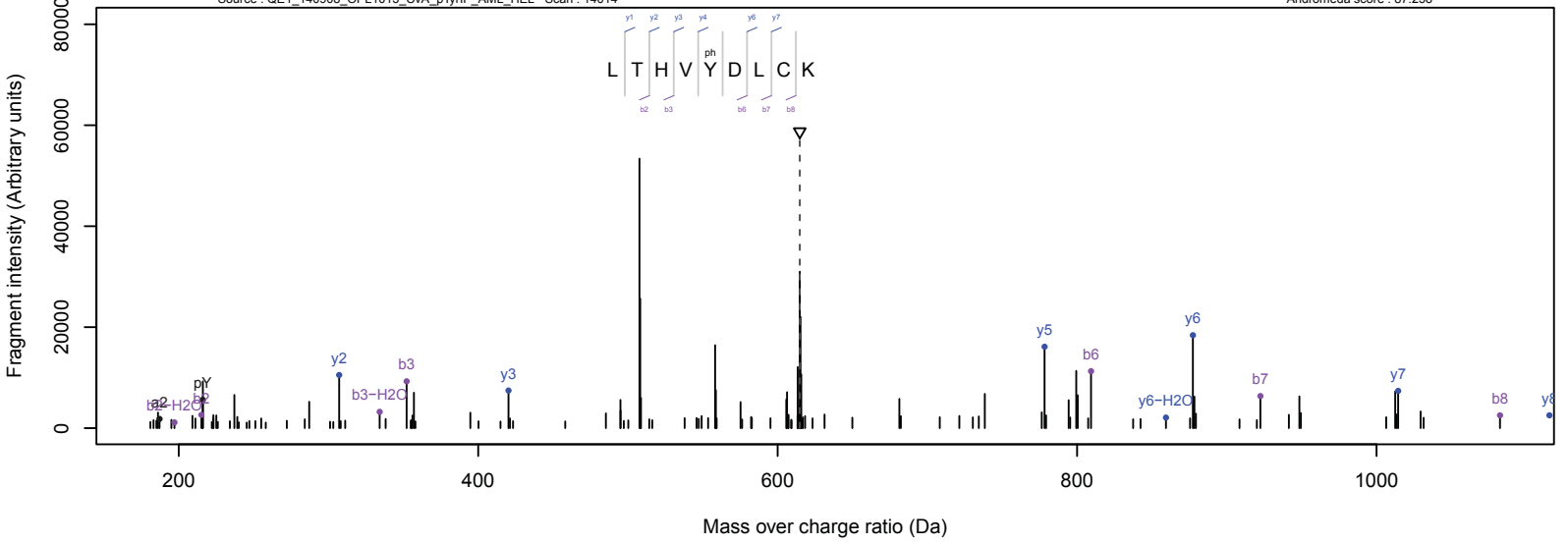
Andromeda score : 87.258



LTHVY(ph)DLCK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 14014

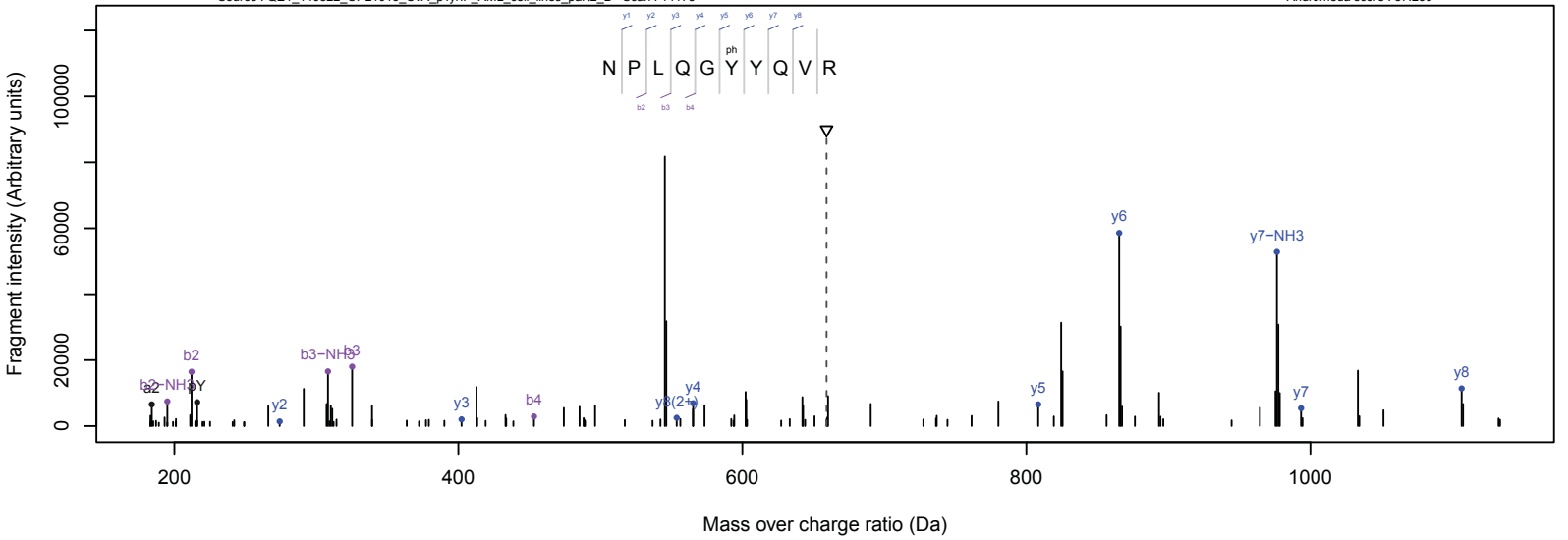
Andromeda score : 87.258



NPLQGY(ph)YQVR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 14178

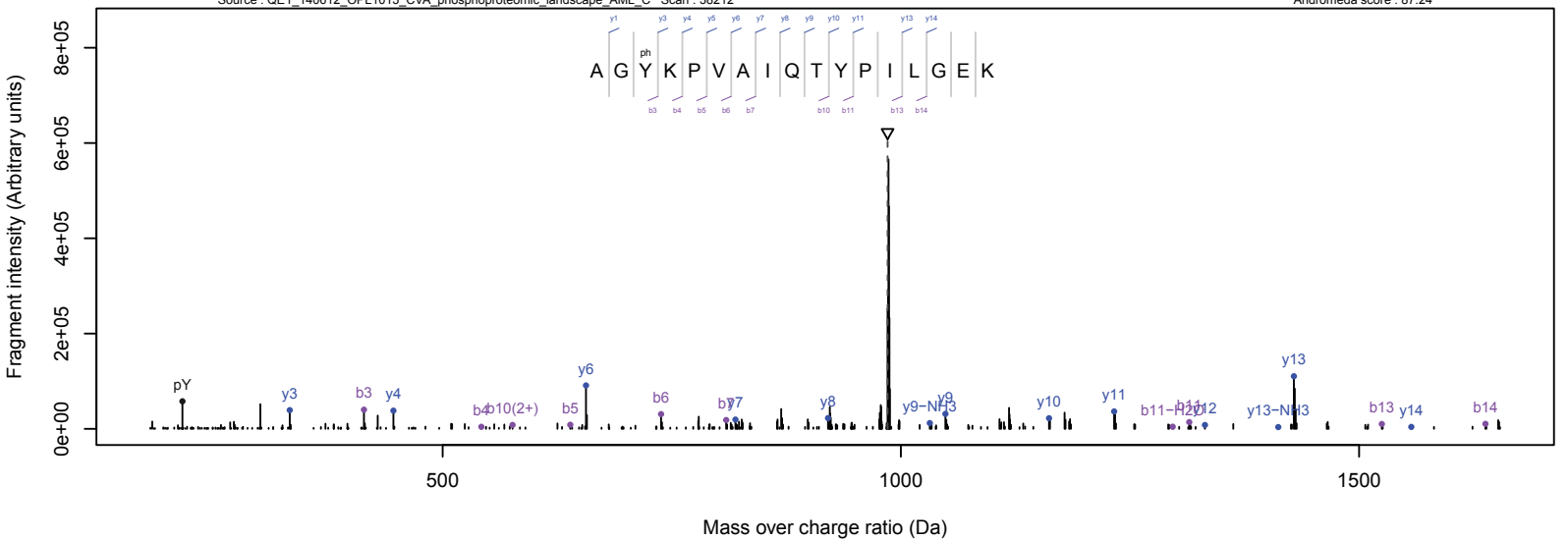
Andromeda score : 87.258



(ac)AGY(ph)KPVAIQTYPI LG EK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38212

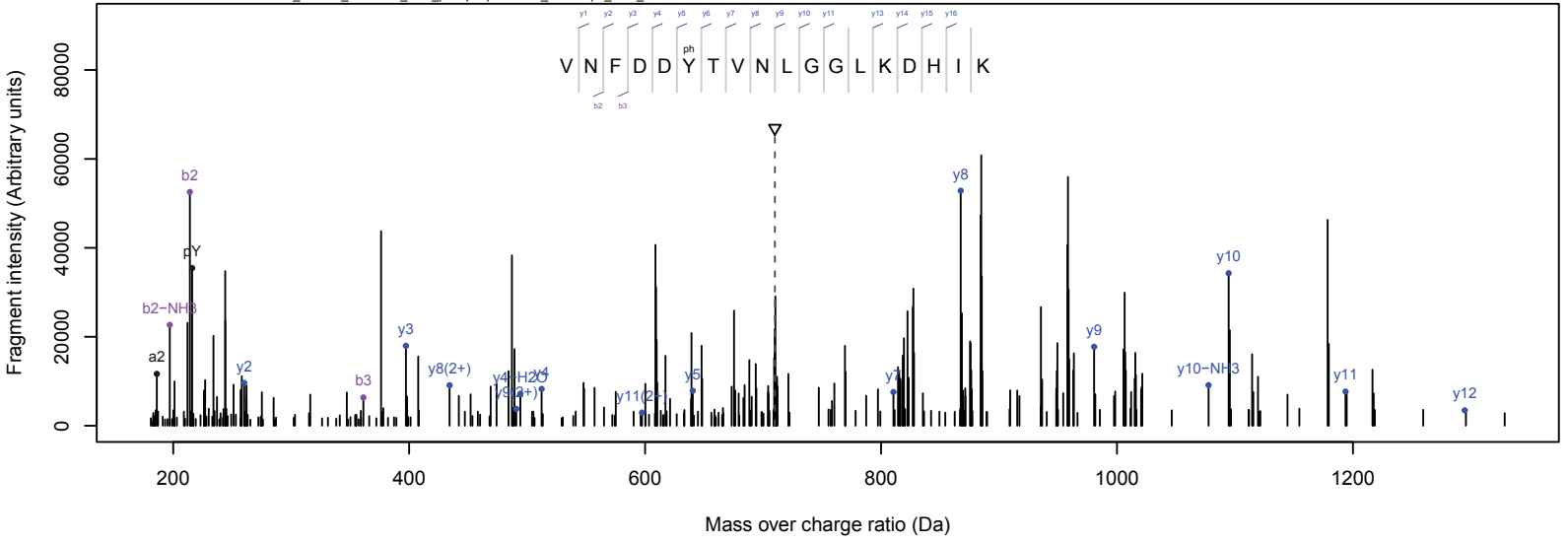
Andromeda score : 87.24



VNFDDY(ph)TVNLGGLKDHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34696

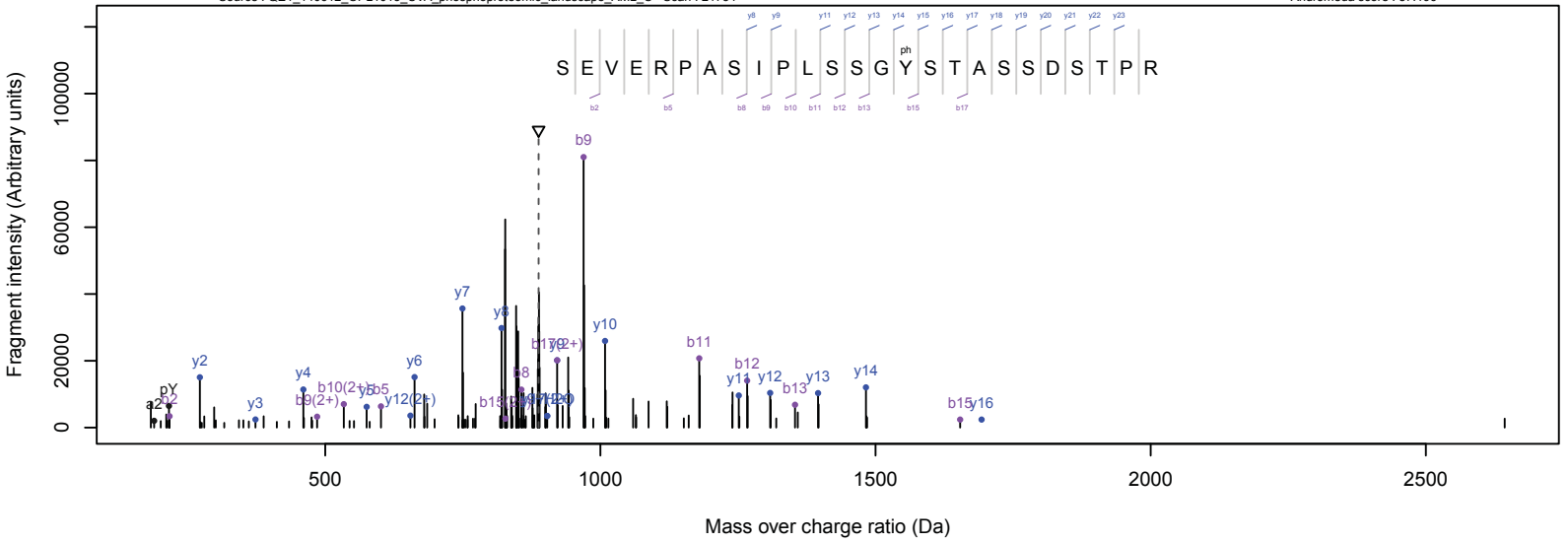
Andromeda score : 87.24



SEVERPASIPLSSGY(ph)STASSDSTPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21754

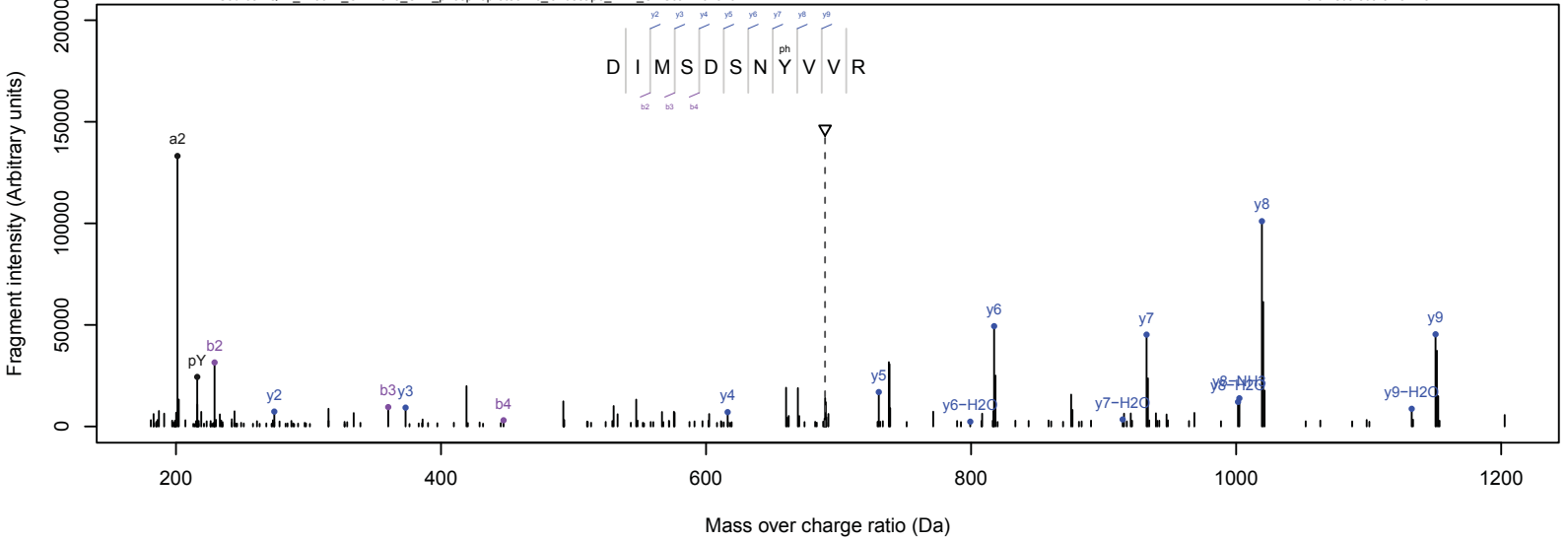
Andromeda score : 87.196



DIMSDSNY(ph)VVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34340

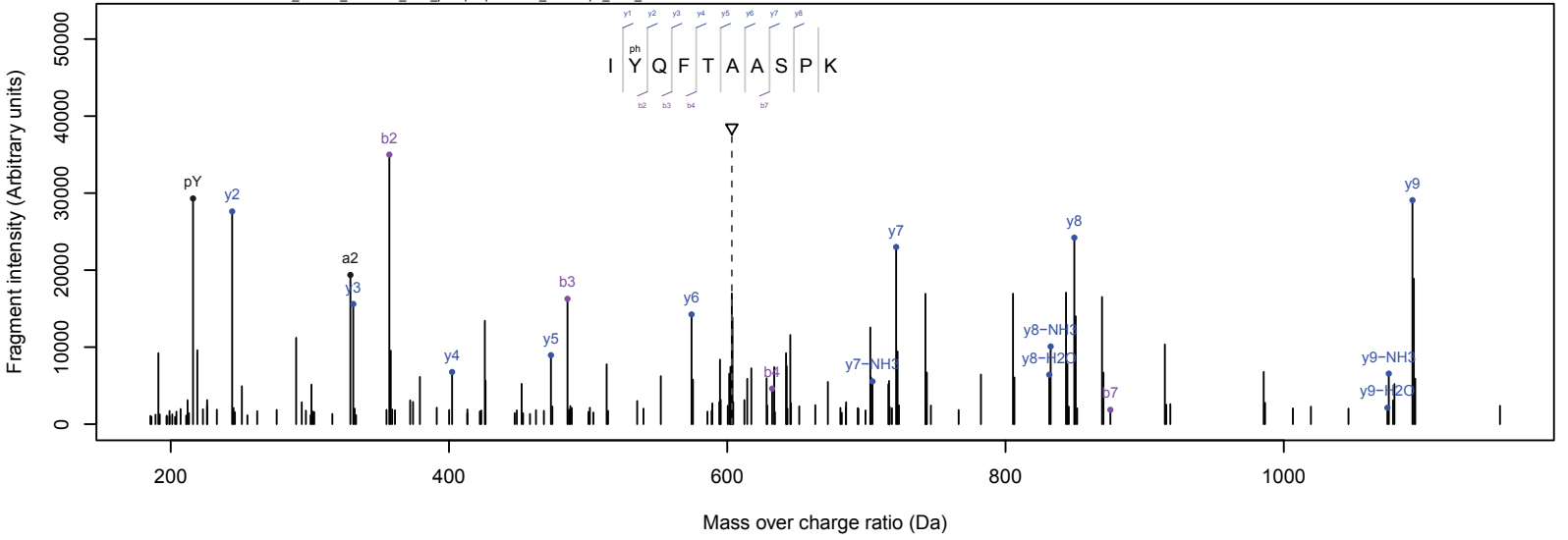
Andromeda score : 87.184



IY(ph)QFTAASPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15488

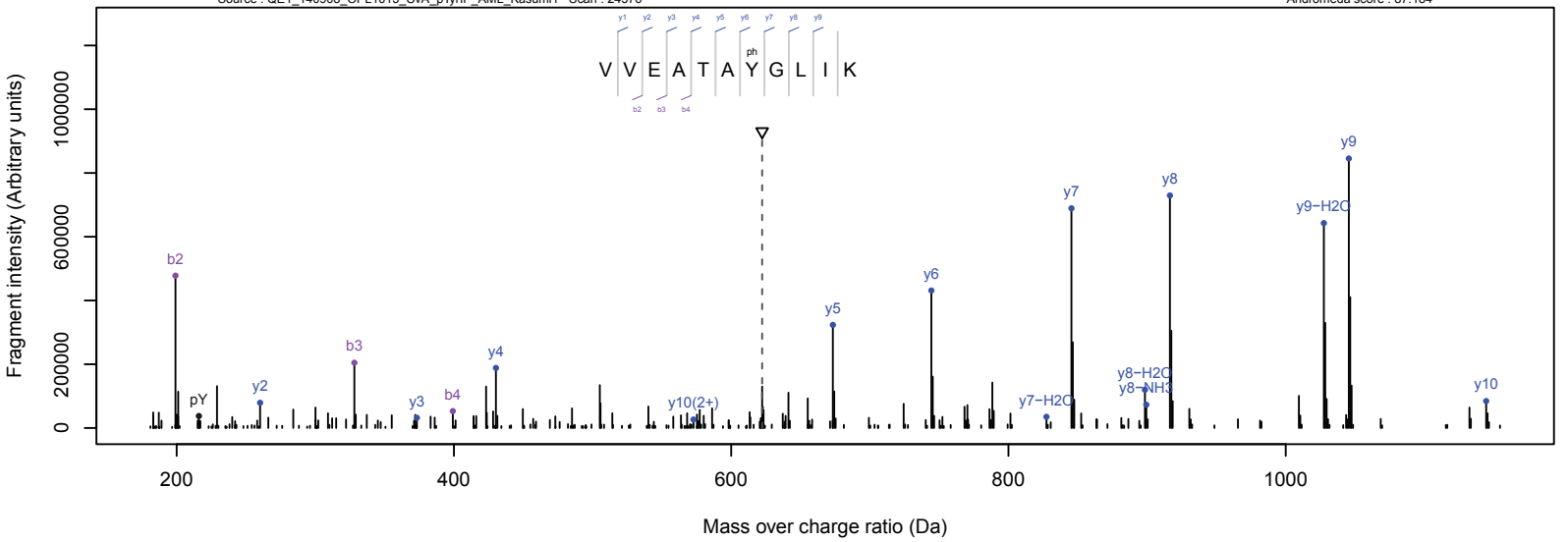
Andromeda score : 87.184



VVEATAY(ph)GLIK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 24576

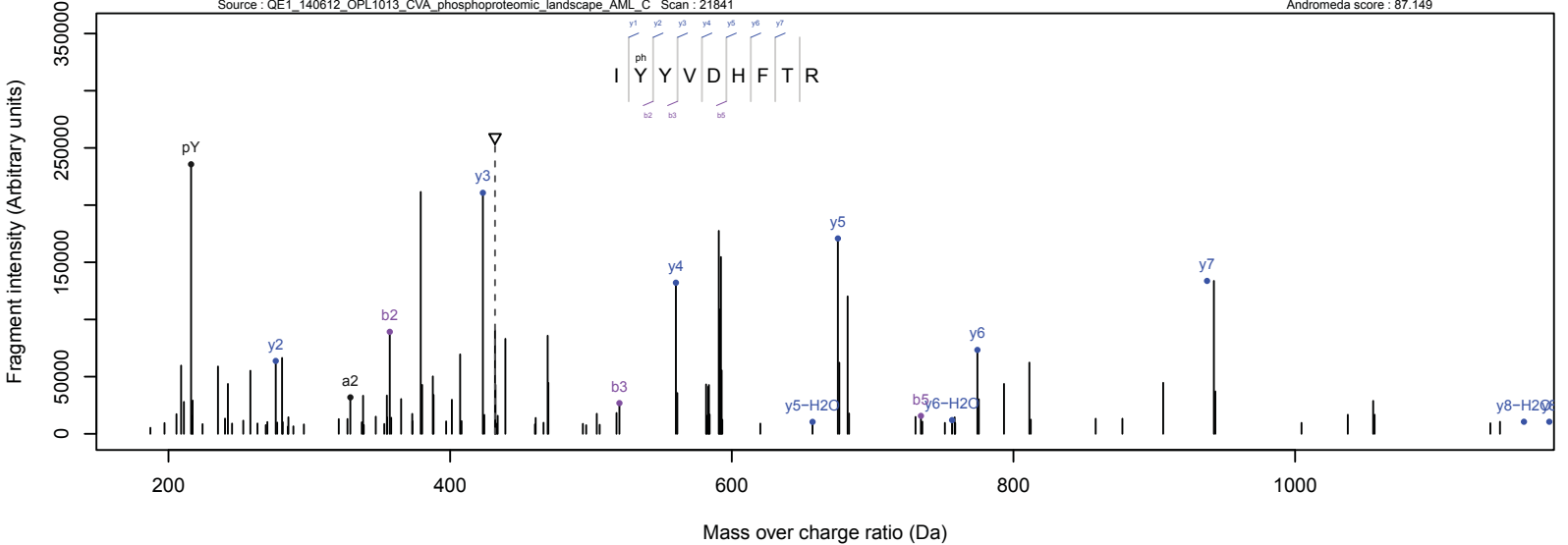
Andromeda score : 87.184



IY(ph)YVDHFTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21841

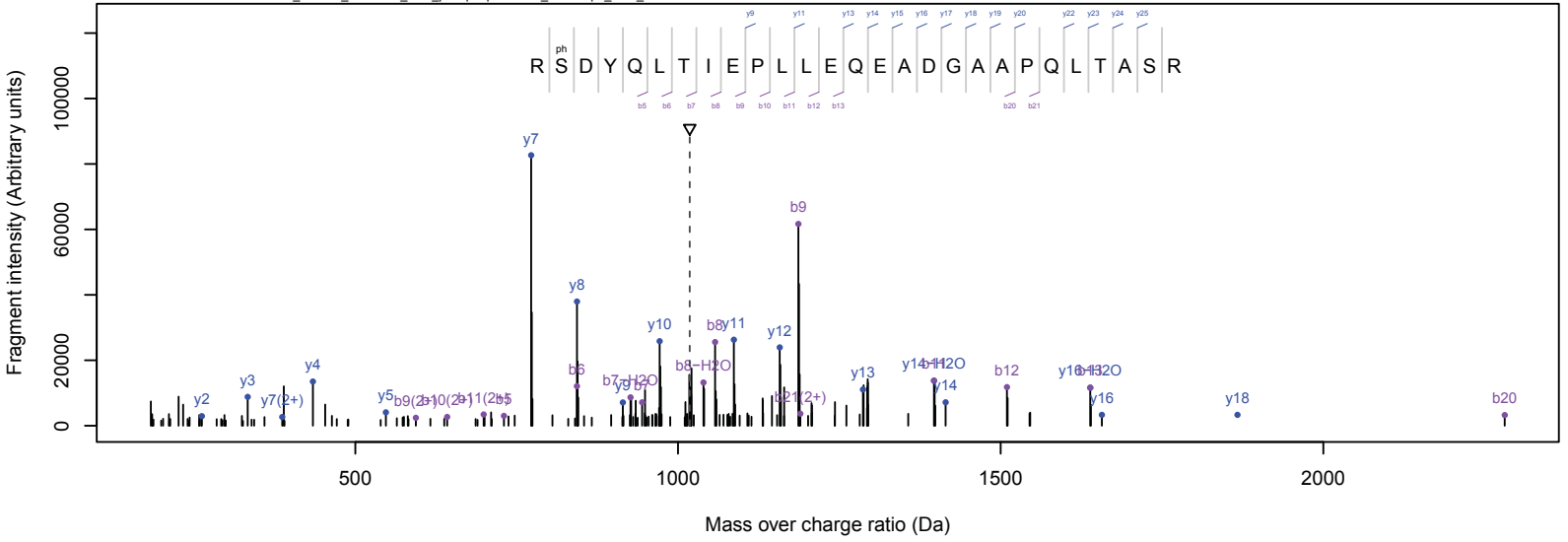
Andromeda score : 87.149



RS(ph)DYQLTIEPLLEQEAADGAAPQLTASR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44285

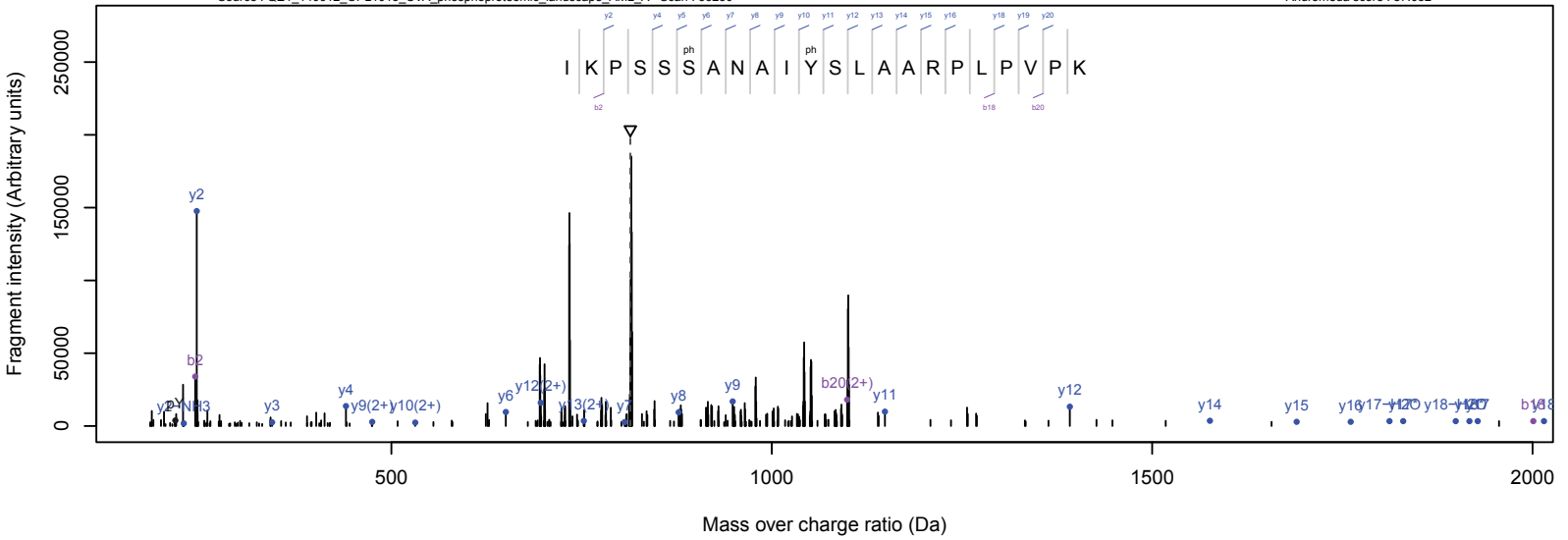
Andromeda score : 87.091



IKPSSS(ph)ANAAY(ph)SLAARPLVPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36250

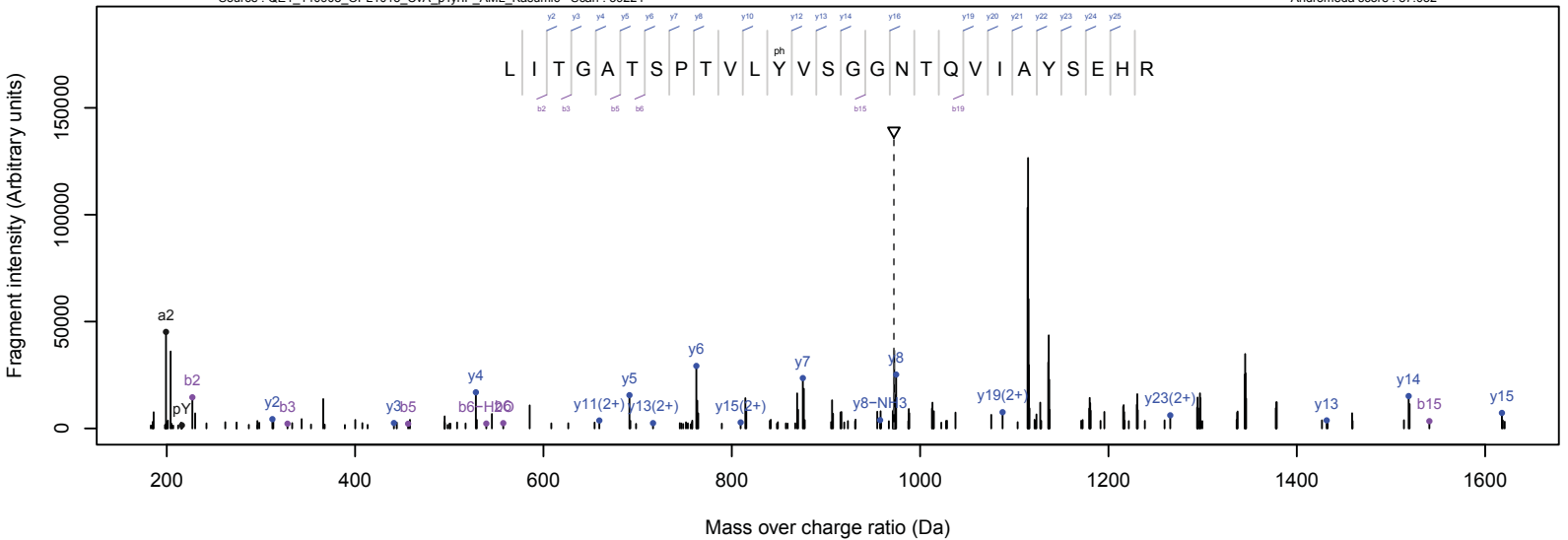
Andromeda score : 87.052



LITGATSPTVLY(ph)VSGGNTQVIAYSEHR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 39224

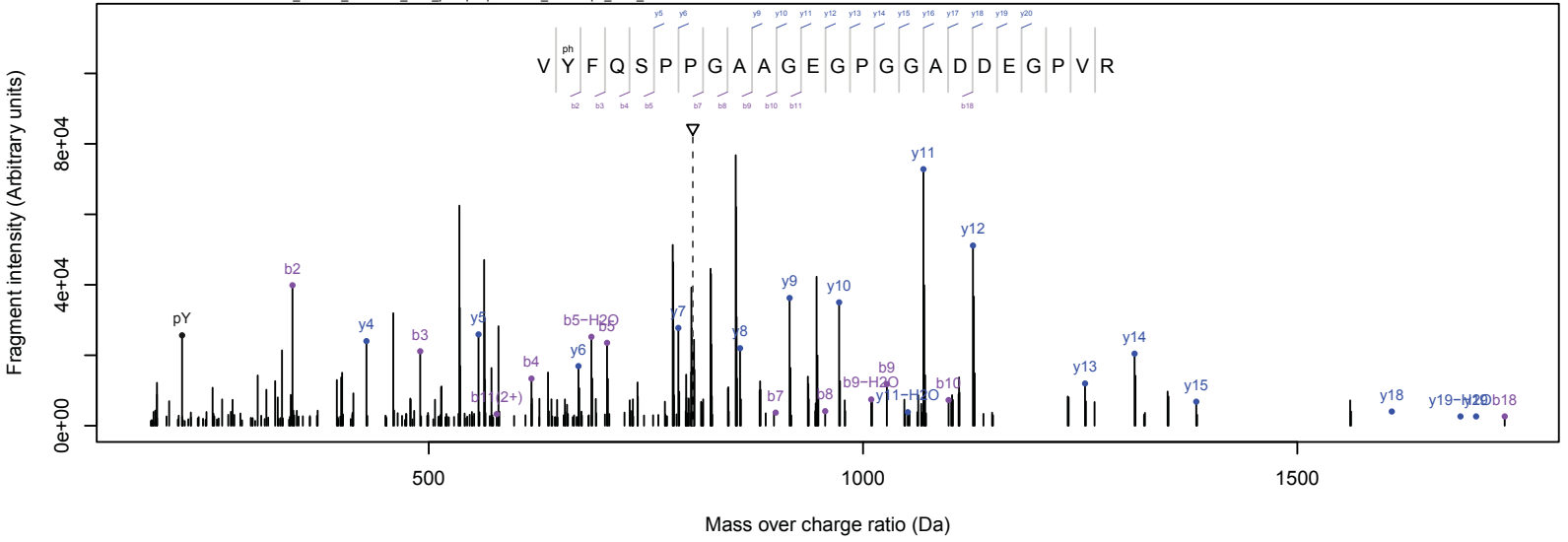
Andromeda score : 87.052



VY(ph)FQSPPGAAGEGPGADDEGPVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26634

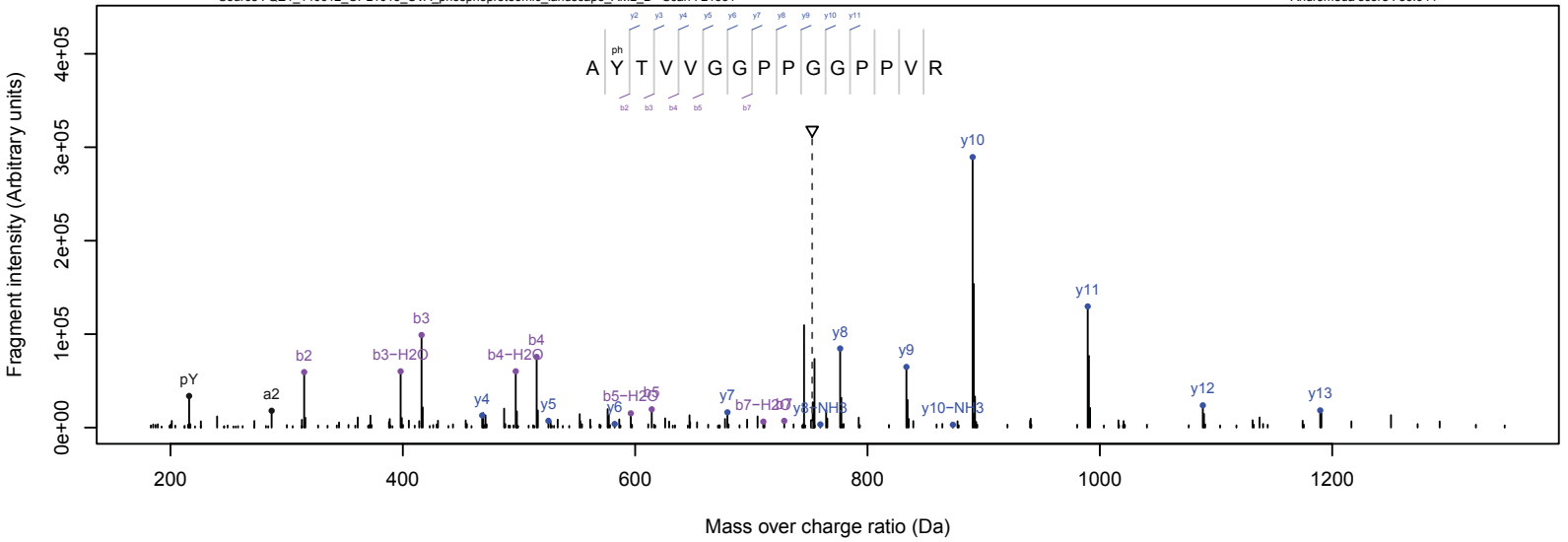
Andromeda score : 86.946



AY(ph)TVVGGPPGPPVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21331

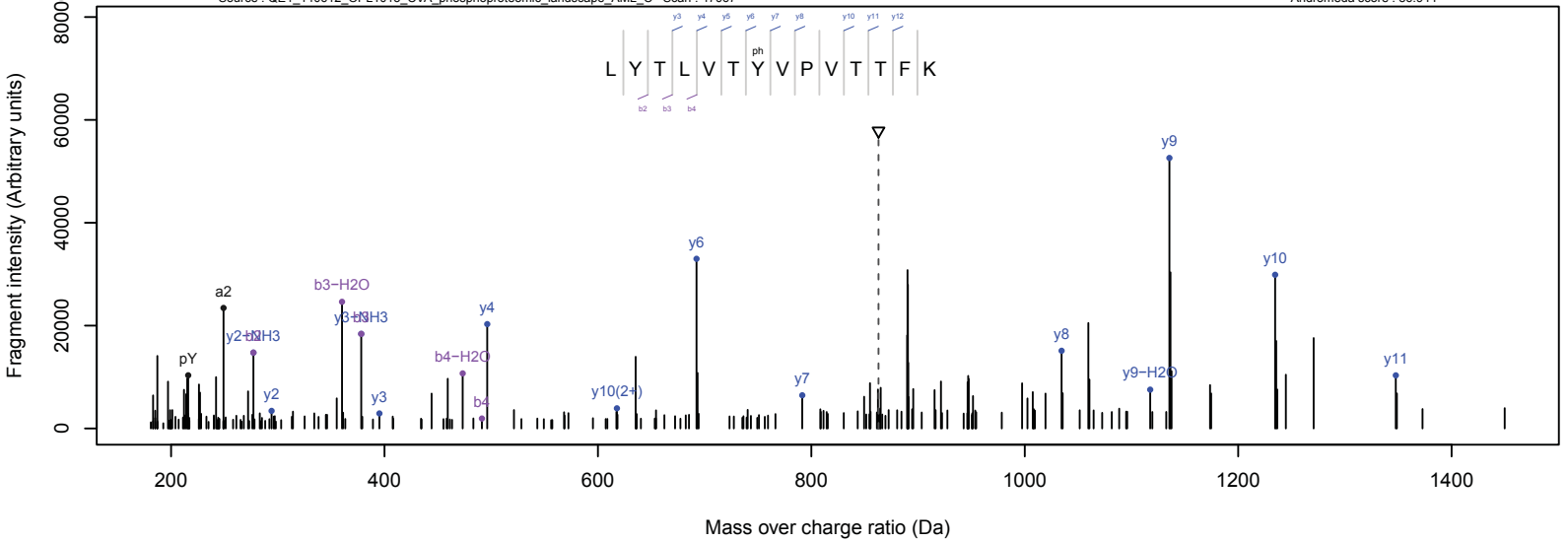
Andromeda score : 86.944



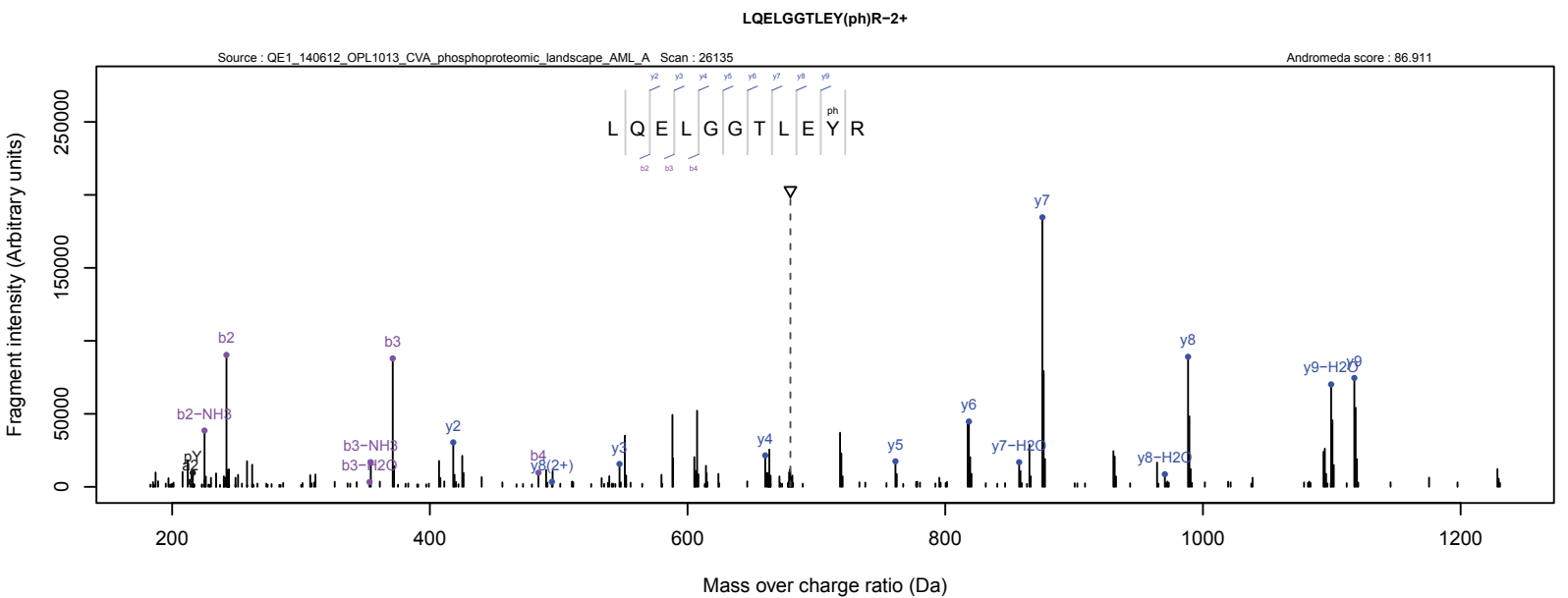
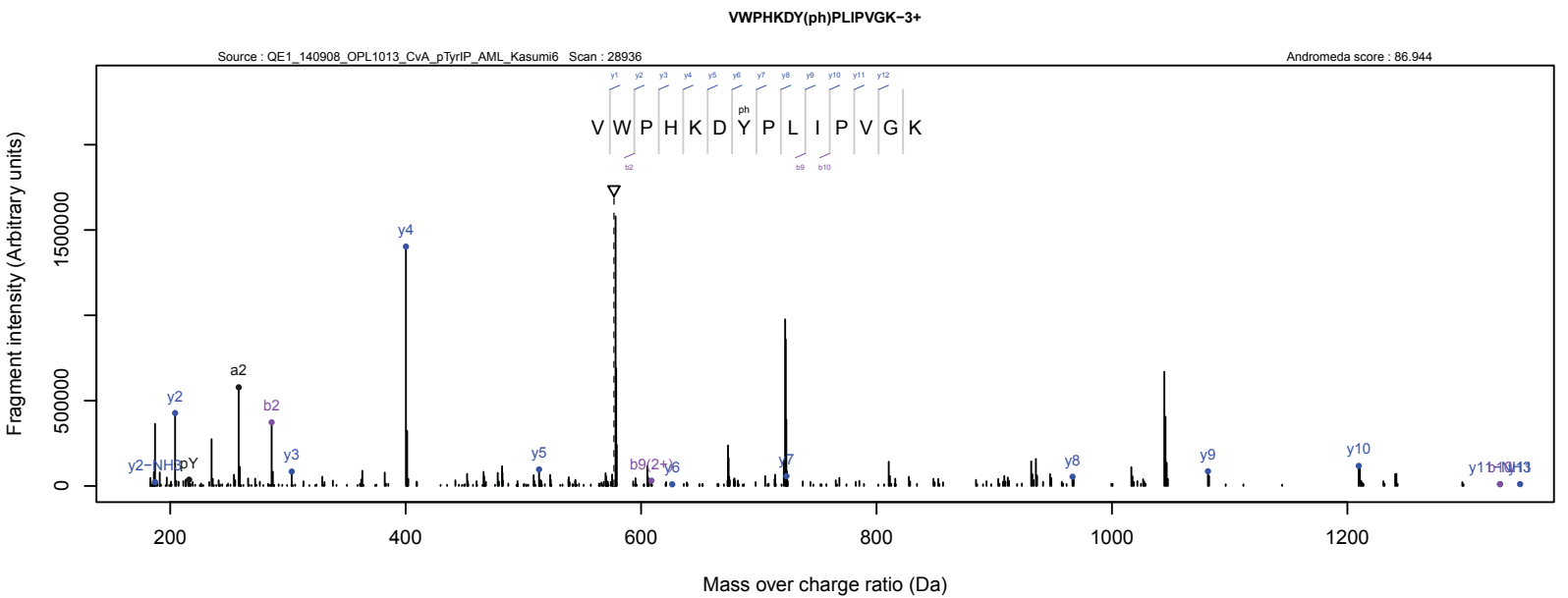
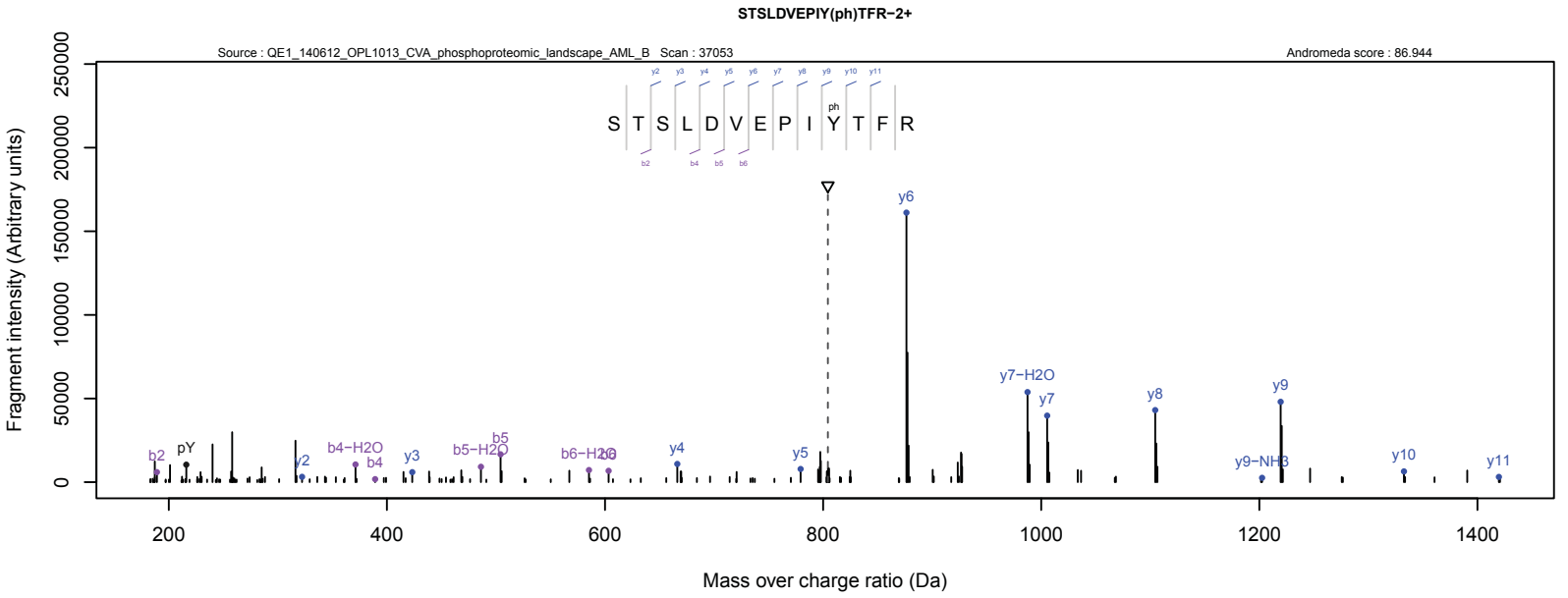
LYTLVY(ph)VPVTFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47067

Andromeda score : 86.944



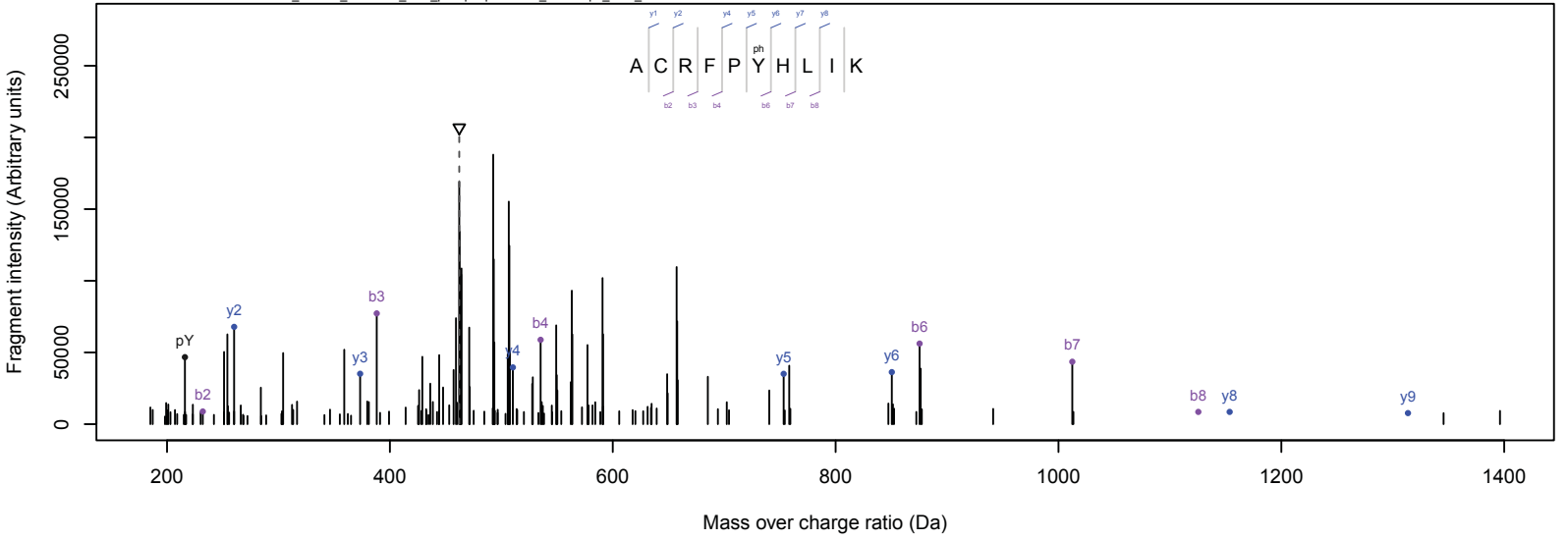




ACRFPY(ph)HLIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20535

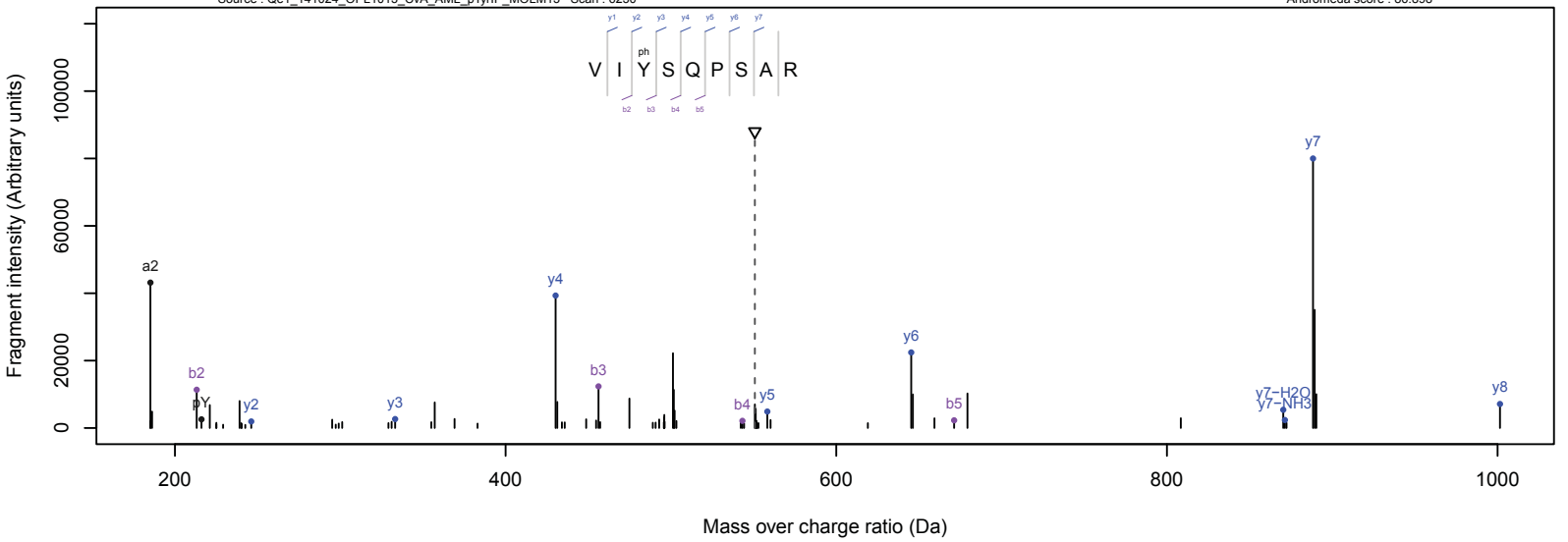
Andromeda score : 86.898



VIY(ph)SQPSAR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 6230

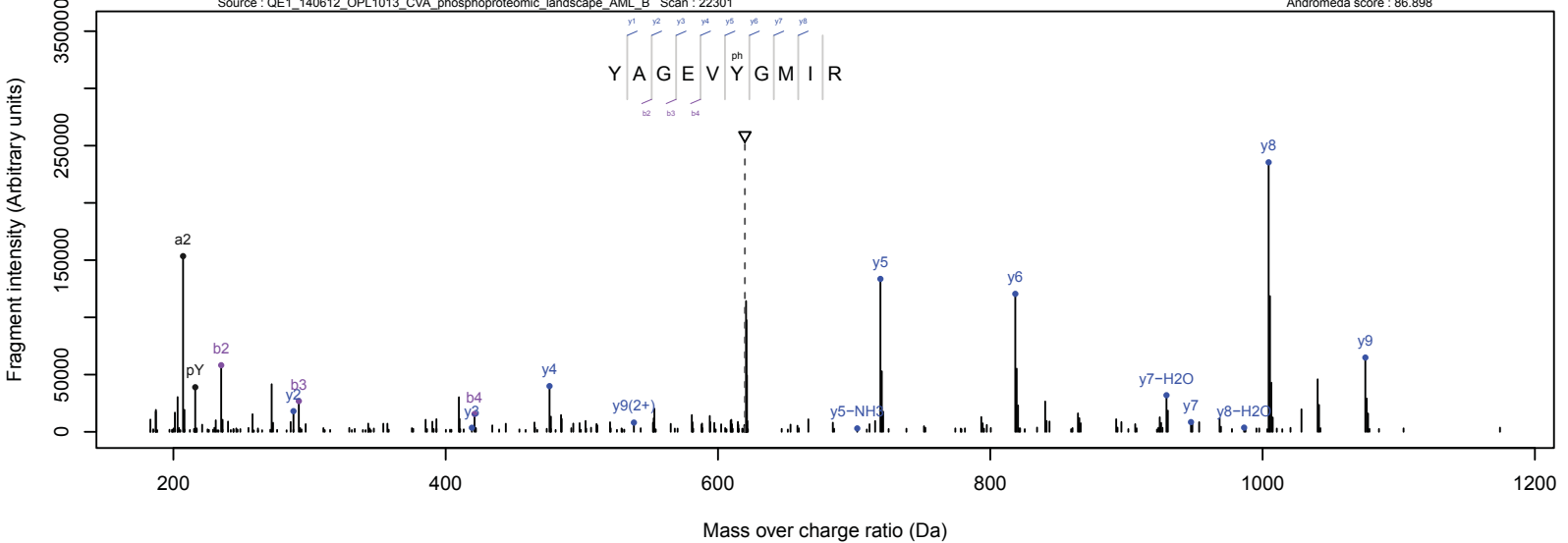
Andromeda score : 86.898



YAGEVY(ph)GMIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22301

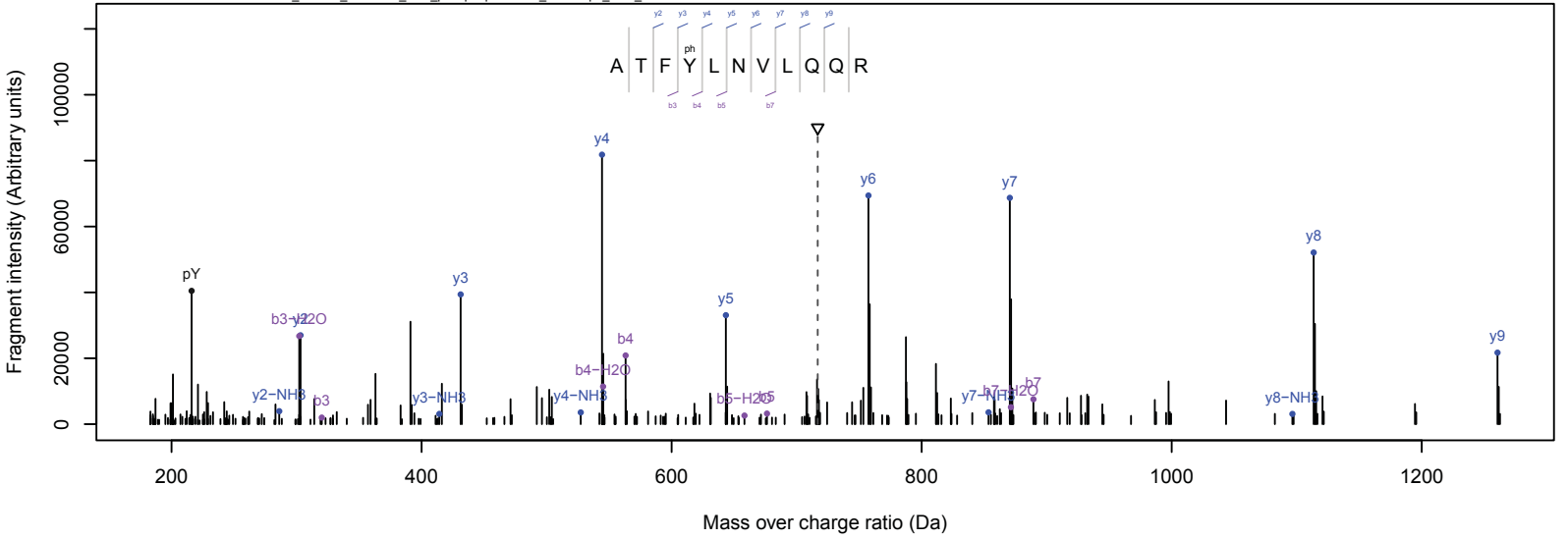
Andromeda score : 86.898



ATFY(ph)LNVLQQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38941

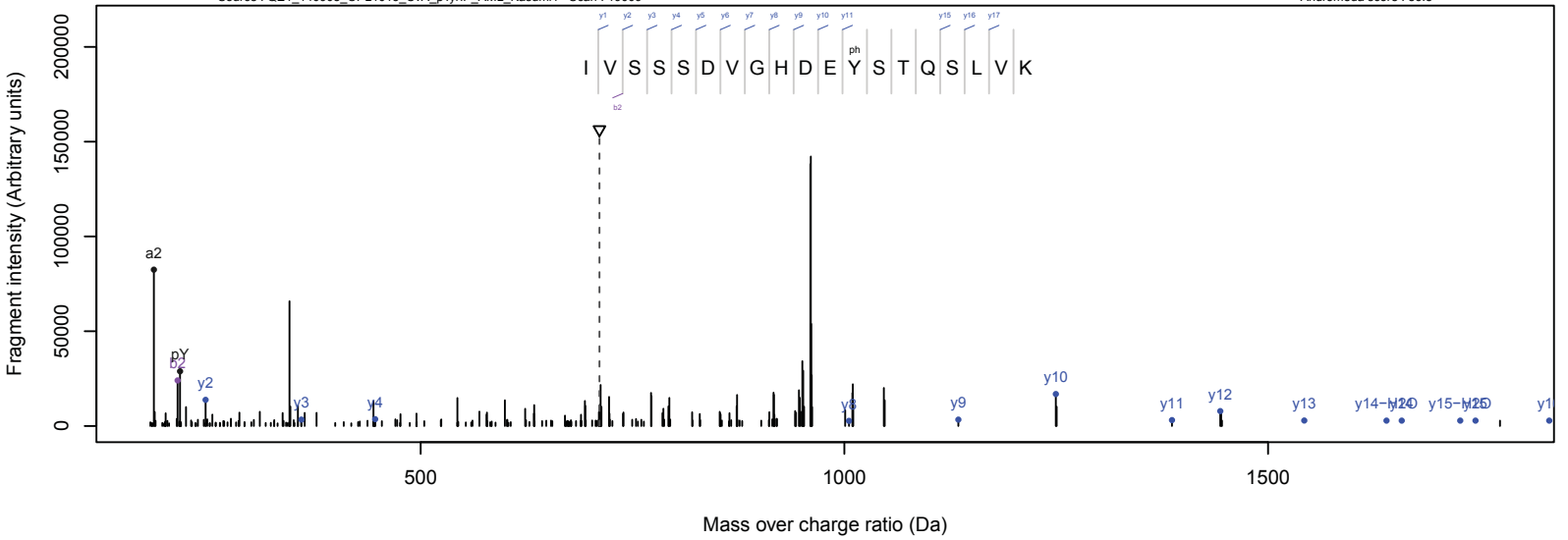
Andromeda score : 86.879



IVSSSDVGHDEY(ph)STQSLVK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 19009

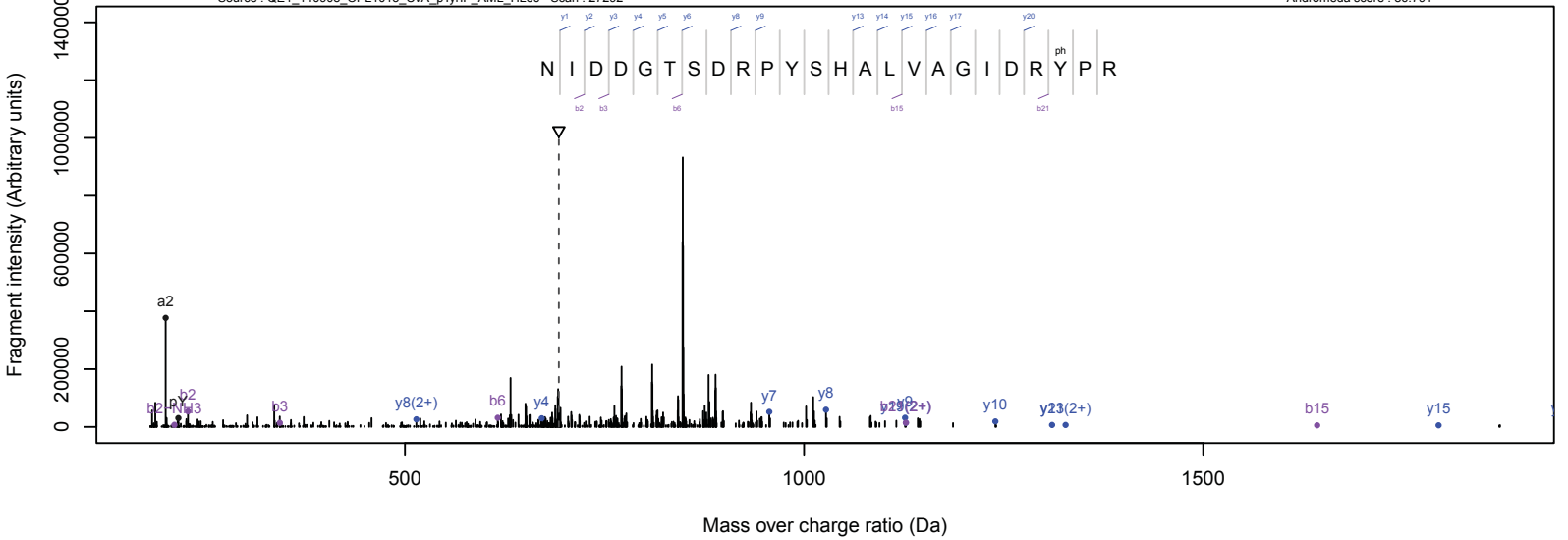
Andromeda score : 86.8



NIDDGTSRYPYSHALVAGIDRY(ph)PR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 27292

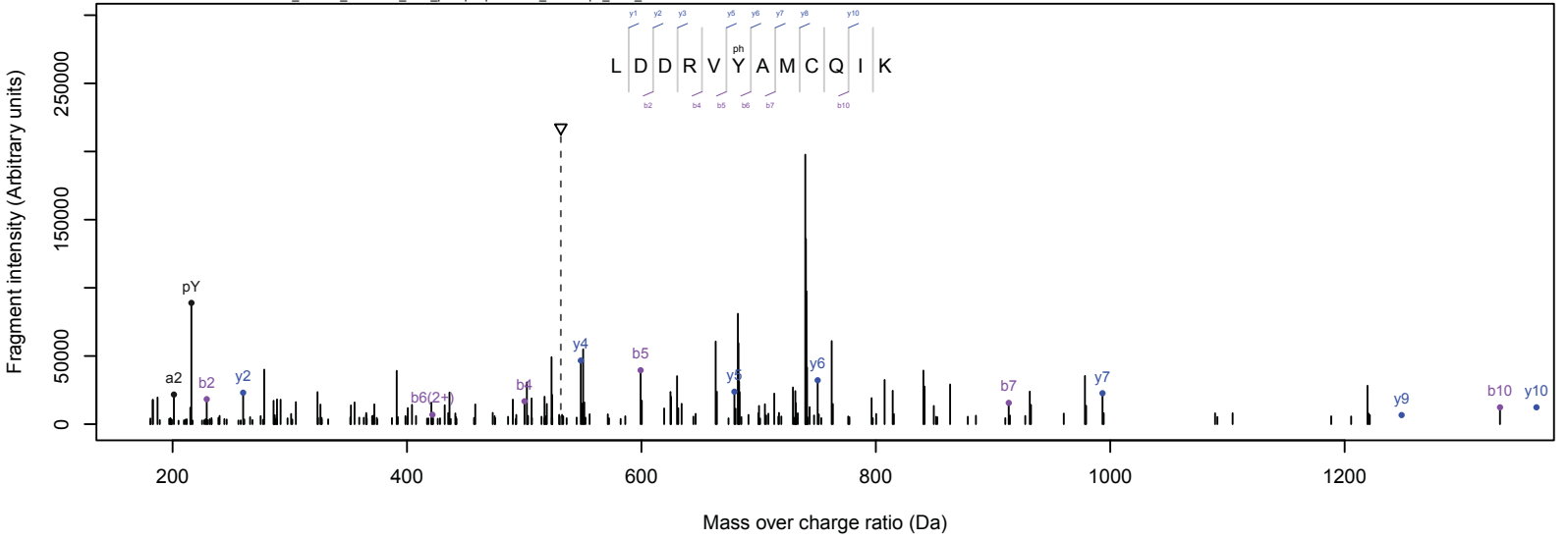
Andromeda score : 86.791



**LDDR<sup>VY</sup>(ph)AMCQIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 21238

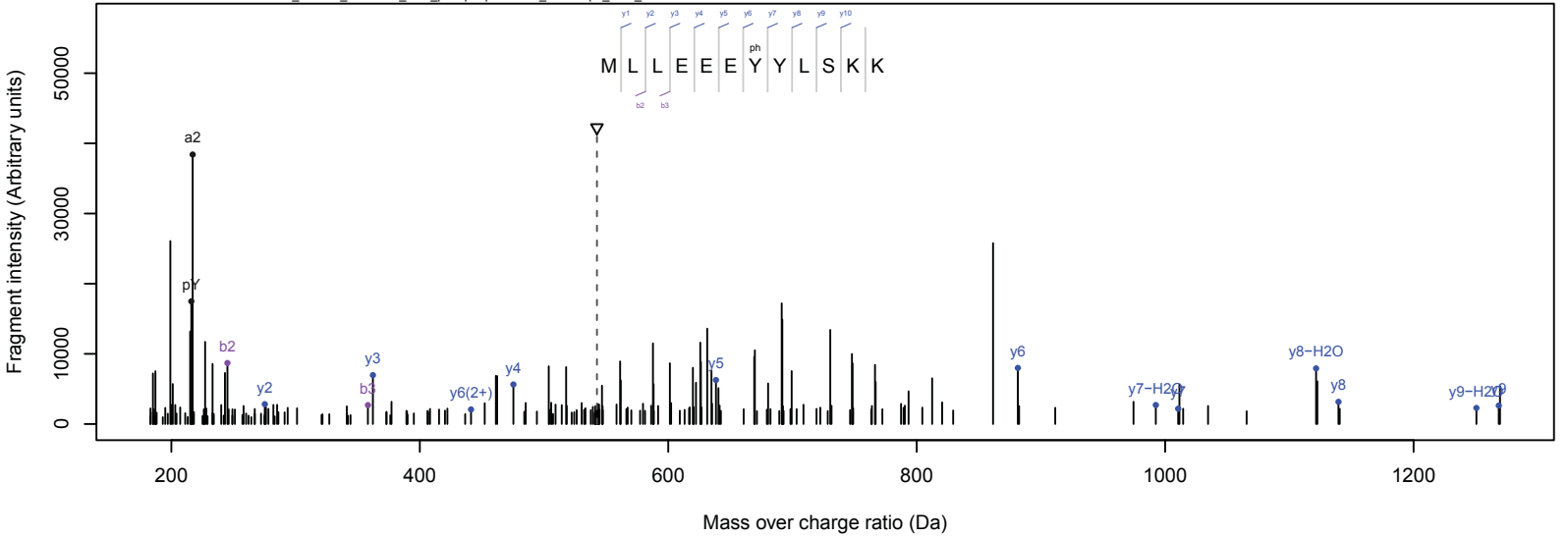
Andromeda score : 86.772



**MLLEEE<sup>Y</sup>YL<sup>SK</sup>K-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30032

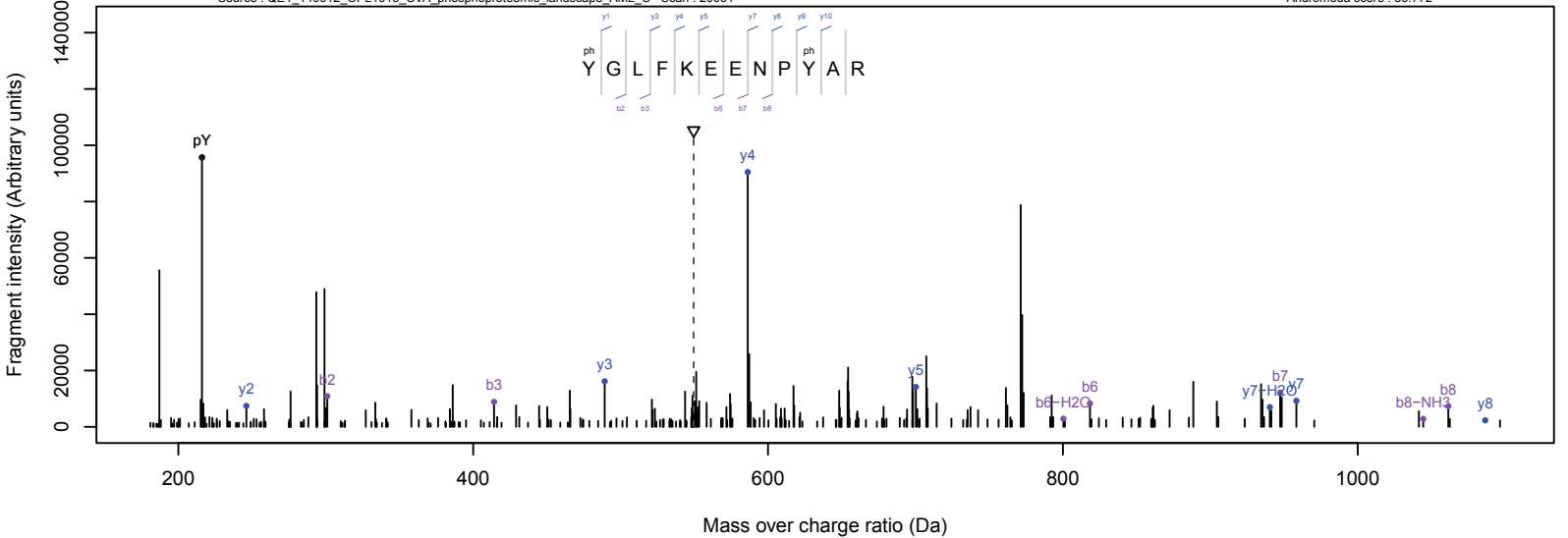
Andromeda score : 86.772



**Y(ph)GLFKEEN<sup>P</sup>YA<sup>R</sup>-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20061

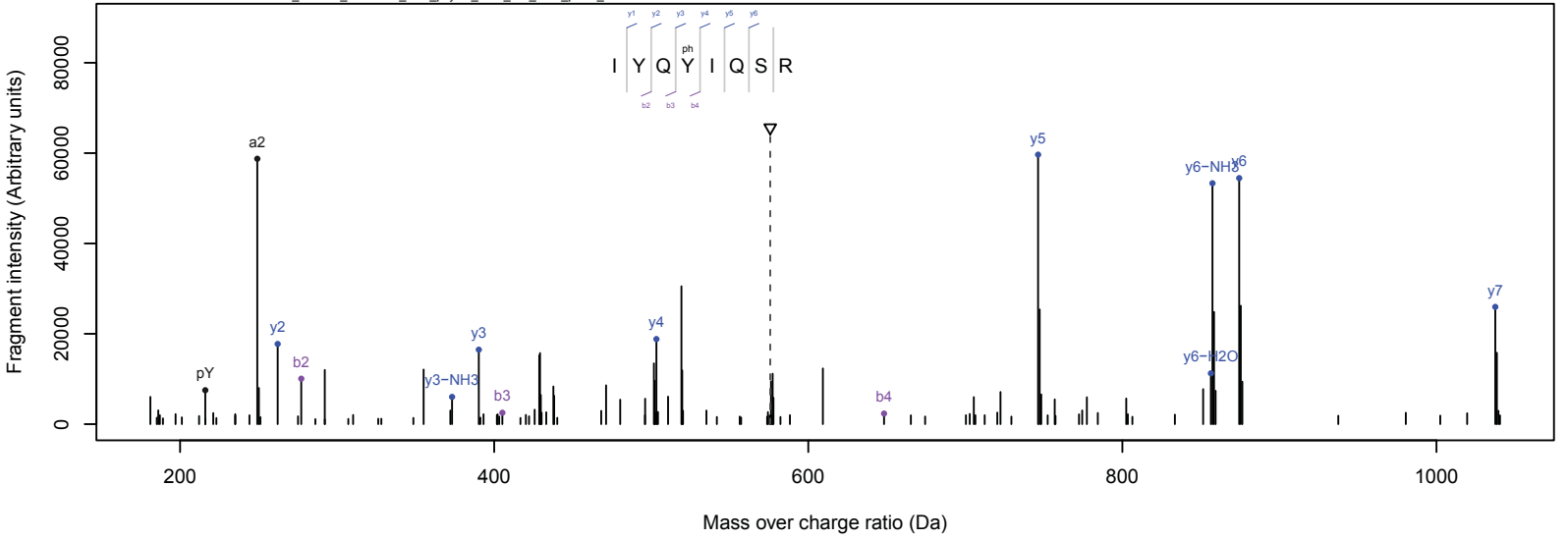
Andromeda score : 86.772



IYQY(ph)QSR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 11704

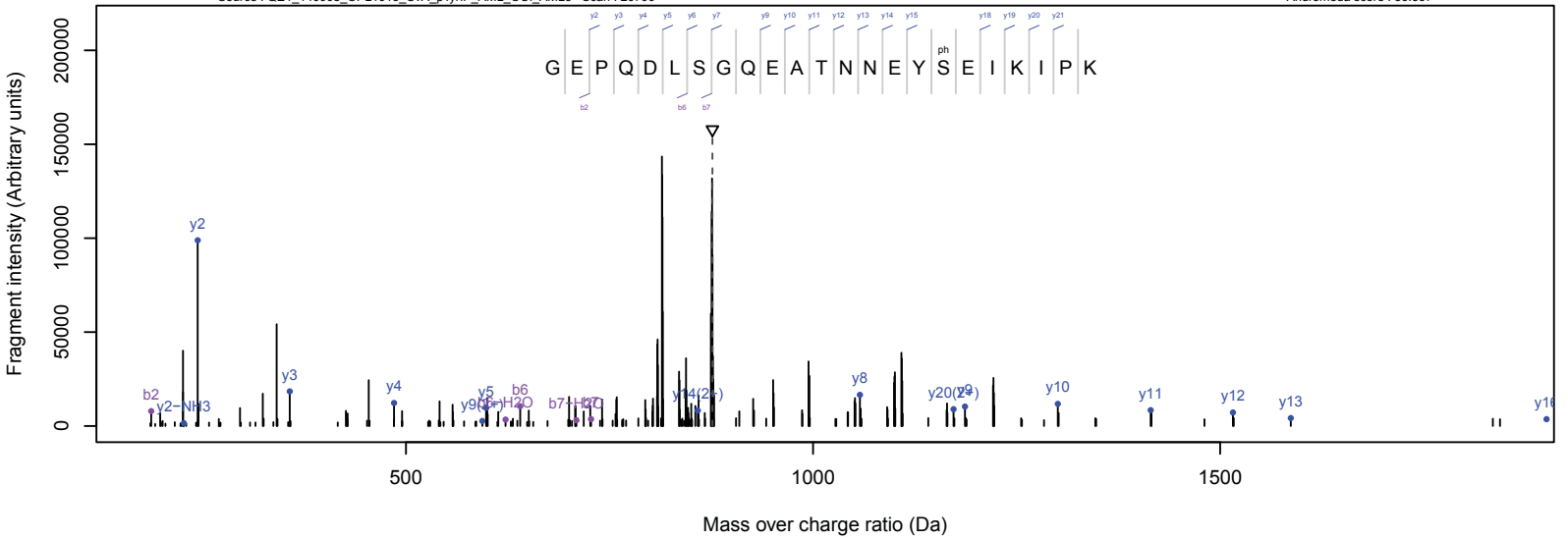
Andromeda score : 86.67



GEPQDLSGQEATNNEY(ph)SEIKIPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 26796

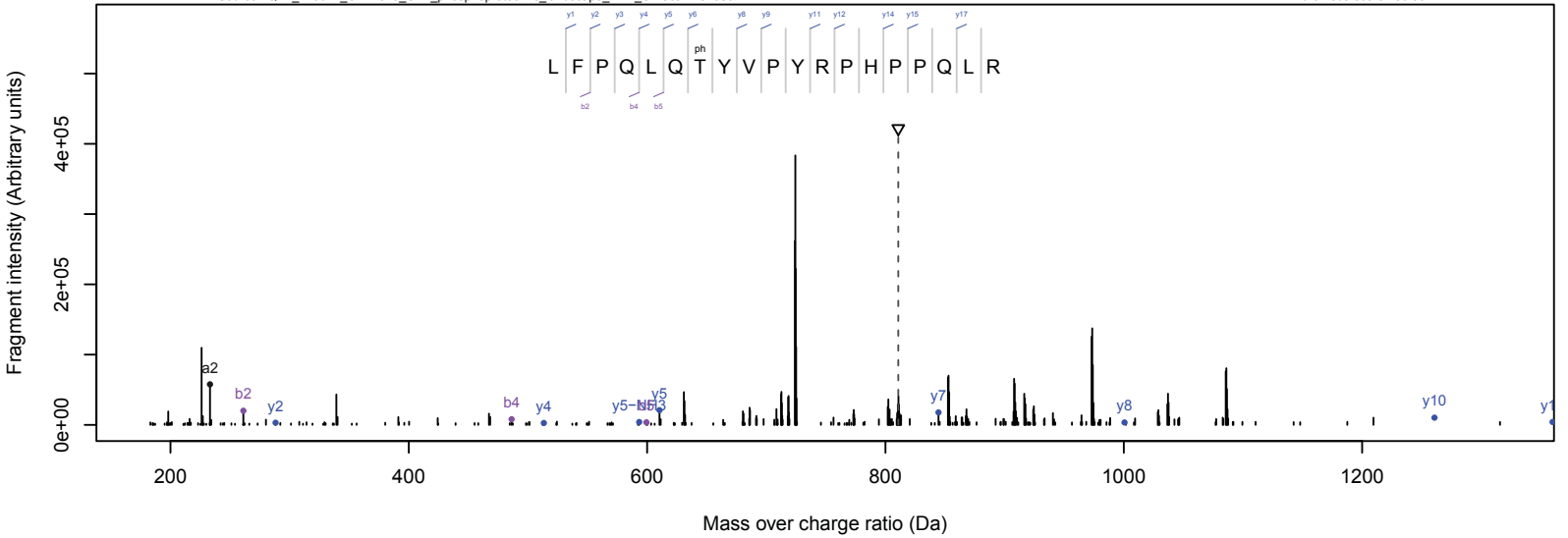
Andromeda score : 86.557



LFPQLQT(ph)YVPYRPHPPQLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37386

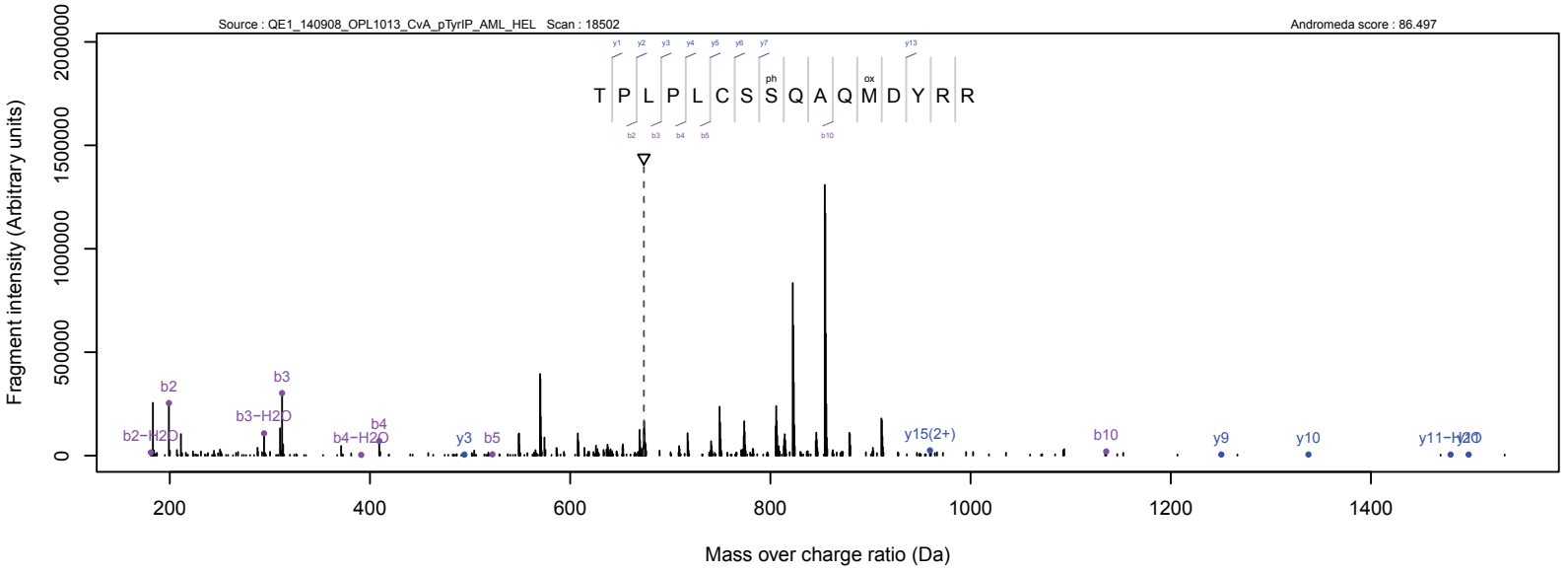
Andromeda score : 86.557



TPLPLCSS(ph)QAQM DYRR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 18502

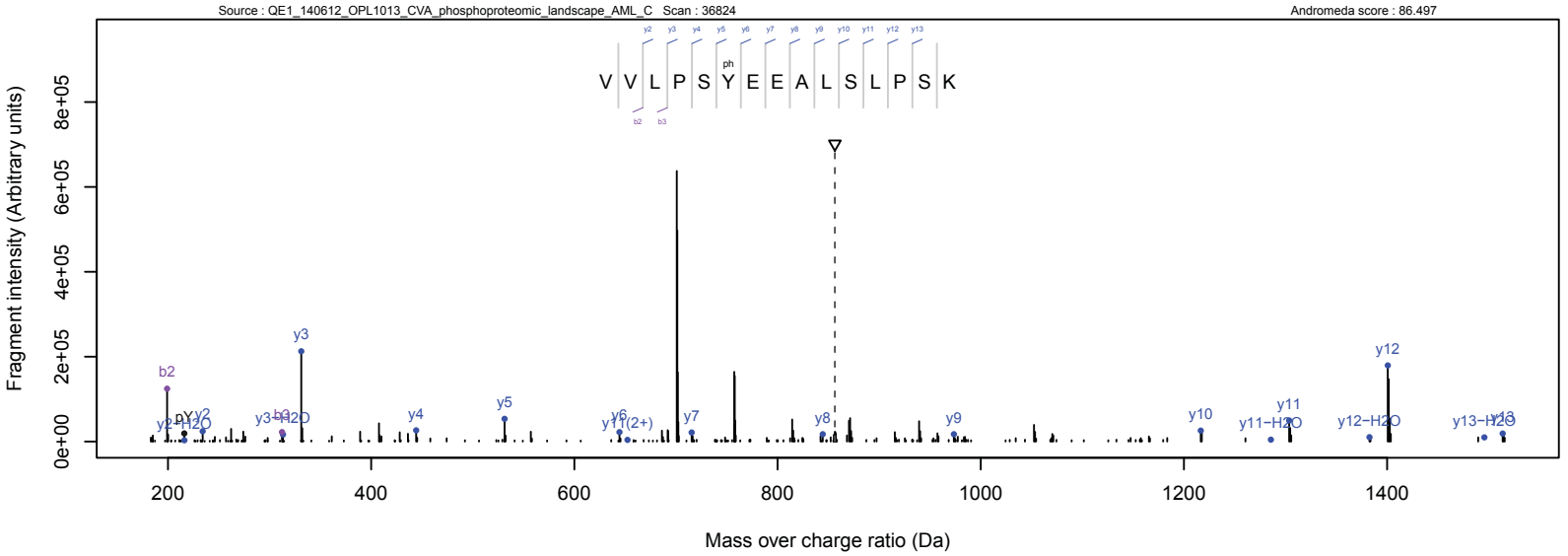
Andromeda score : 86.497



VVLPSY(ph)EEALSLPSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36824

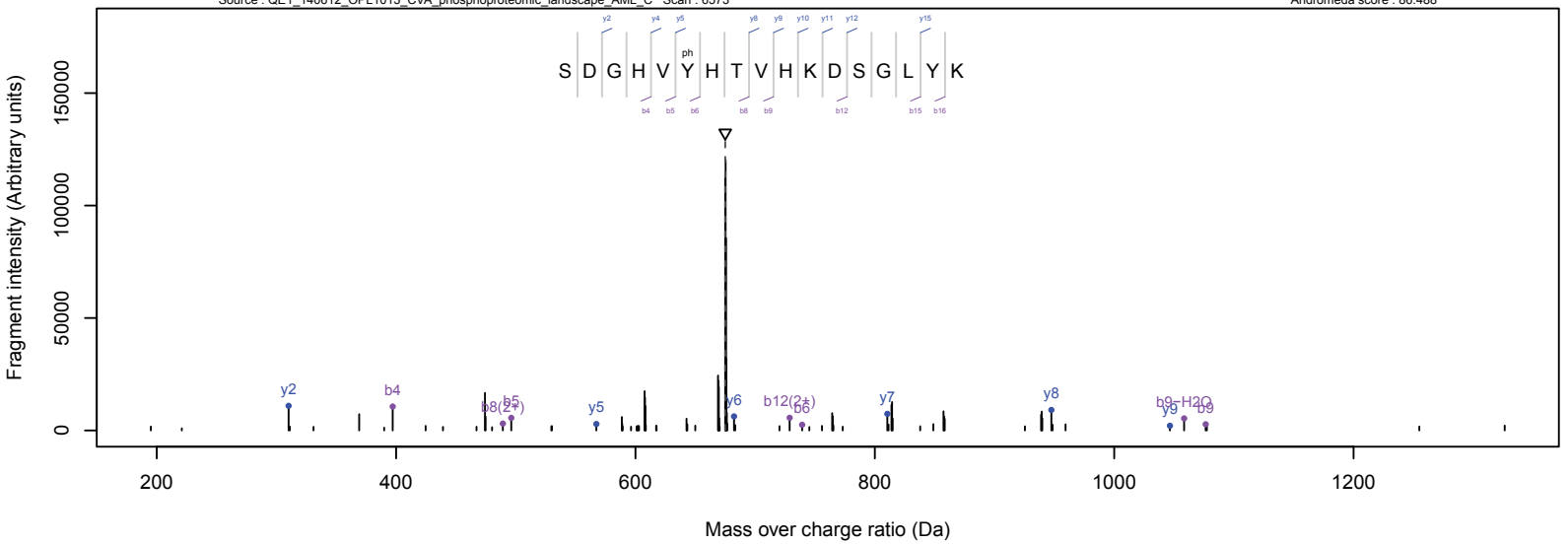
Andromeda score : 86.497



SDGHVY(ph)HTVHKDSGLYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6573

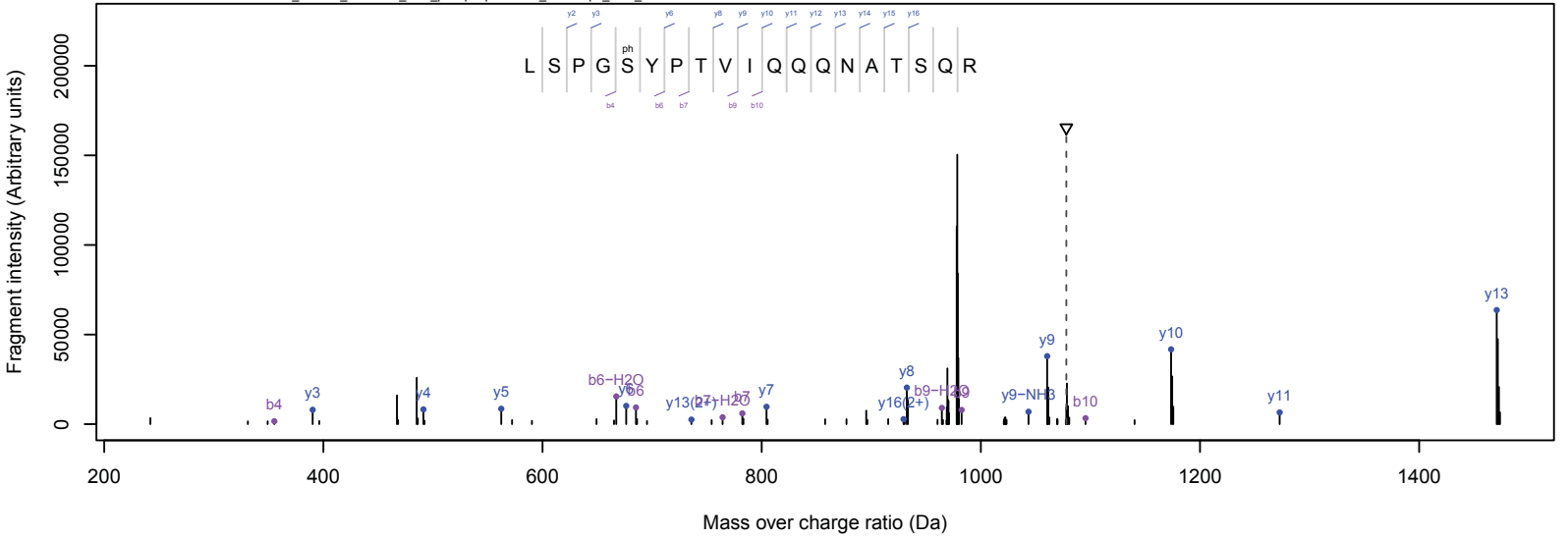
Andromeda score : 86.488



LSPGS(ph)YPTVIQQQNATSQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20120

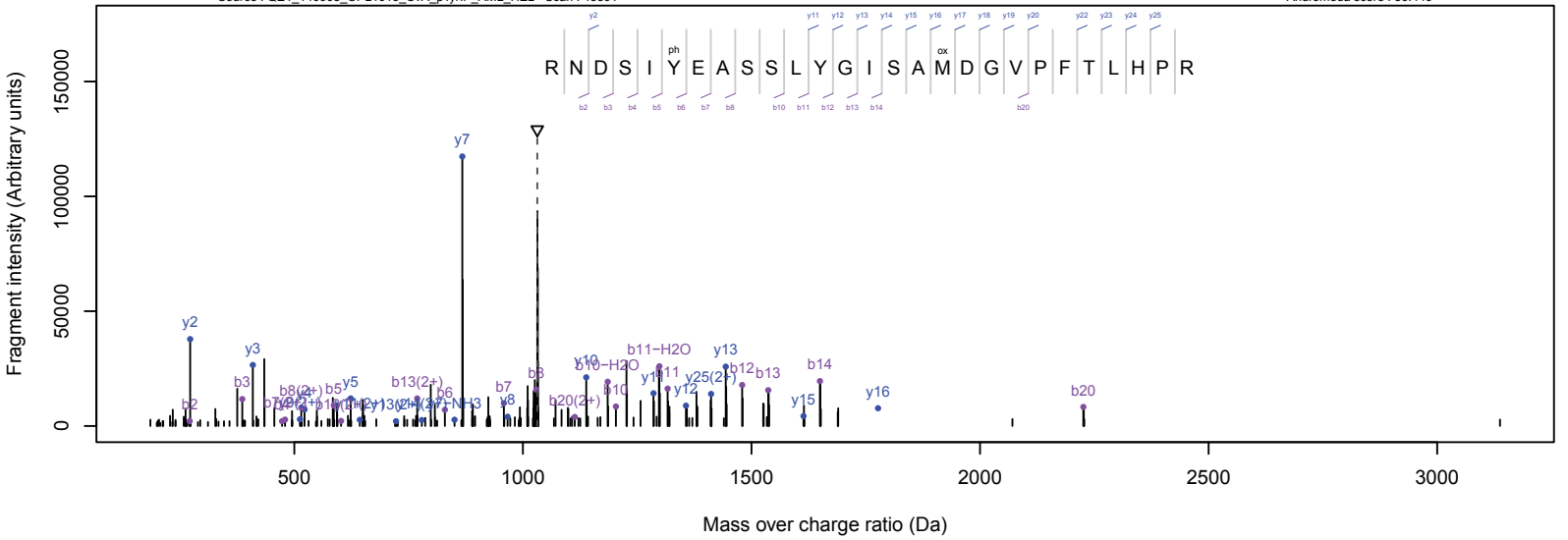
Andromeda score : 86.453



RNSIY(ph)EASSLYGISAM(x)DGVPTLHPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 43891

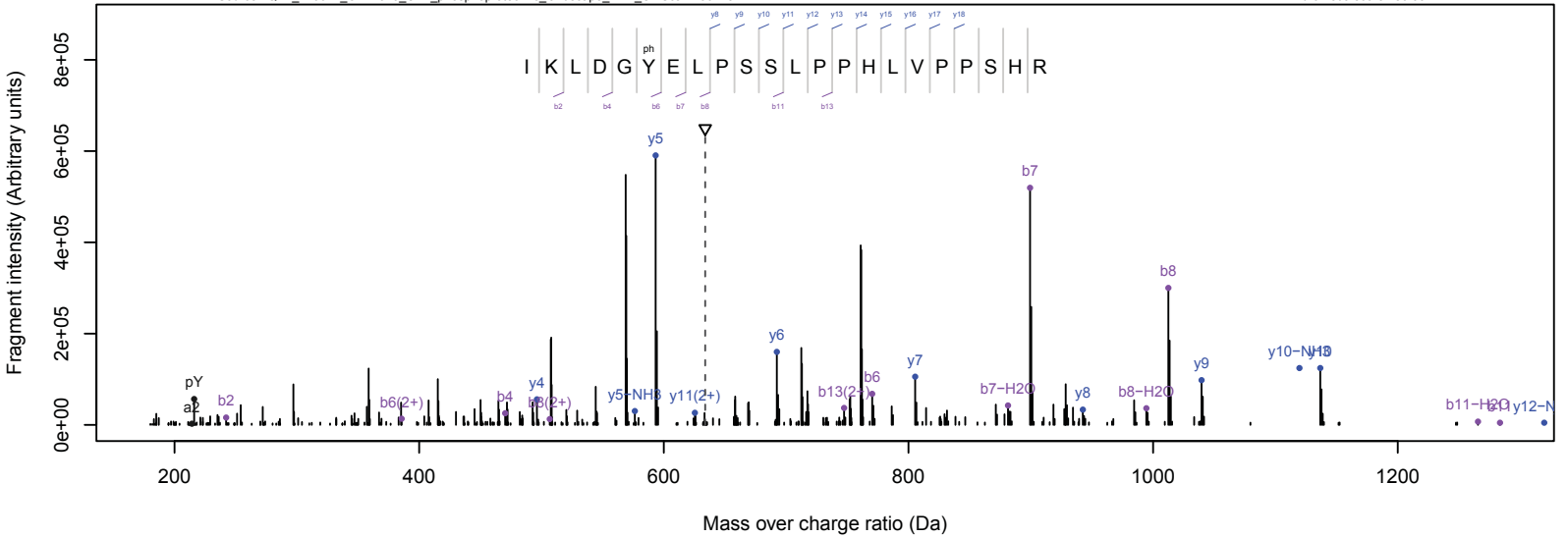
Andromeda score : 86.449



IKLDGY(ph)ELPSSLPVPPSHR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36218

Andromeda score : 86.357



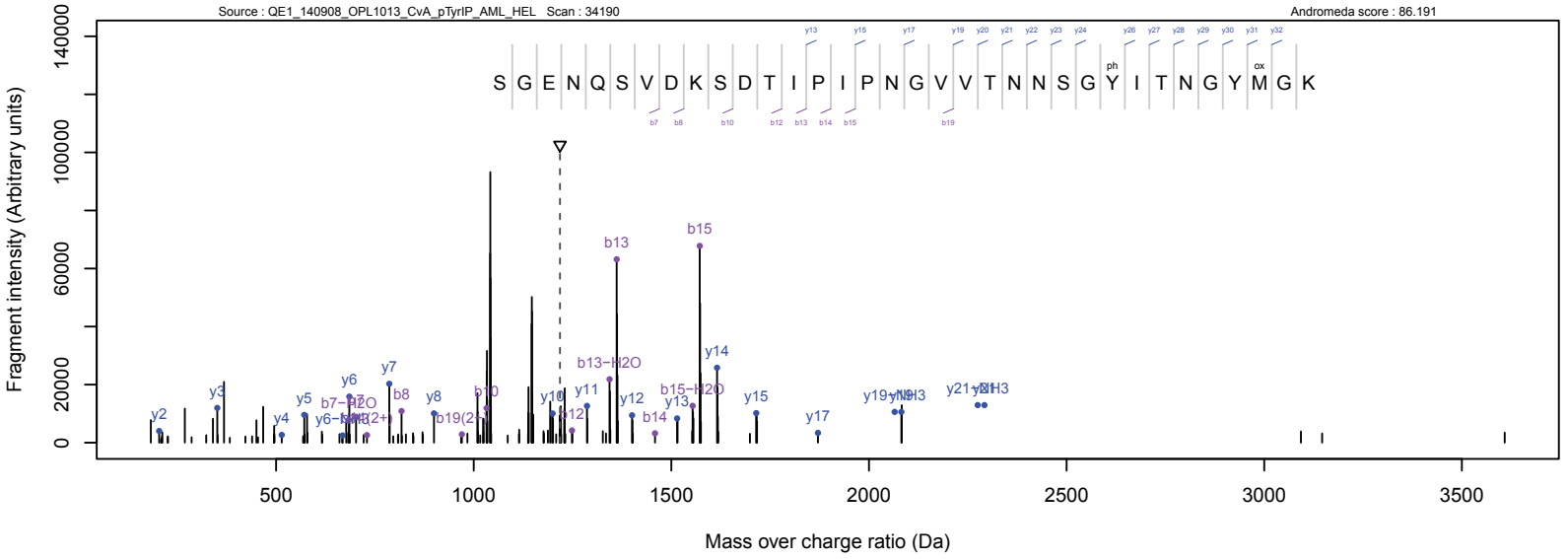




SGENQSVDKKSDTIPIPIPNGVVVTNNSGY(ph)ITNGYM(ox)GK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 34190

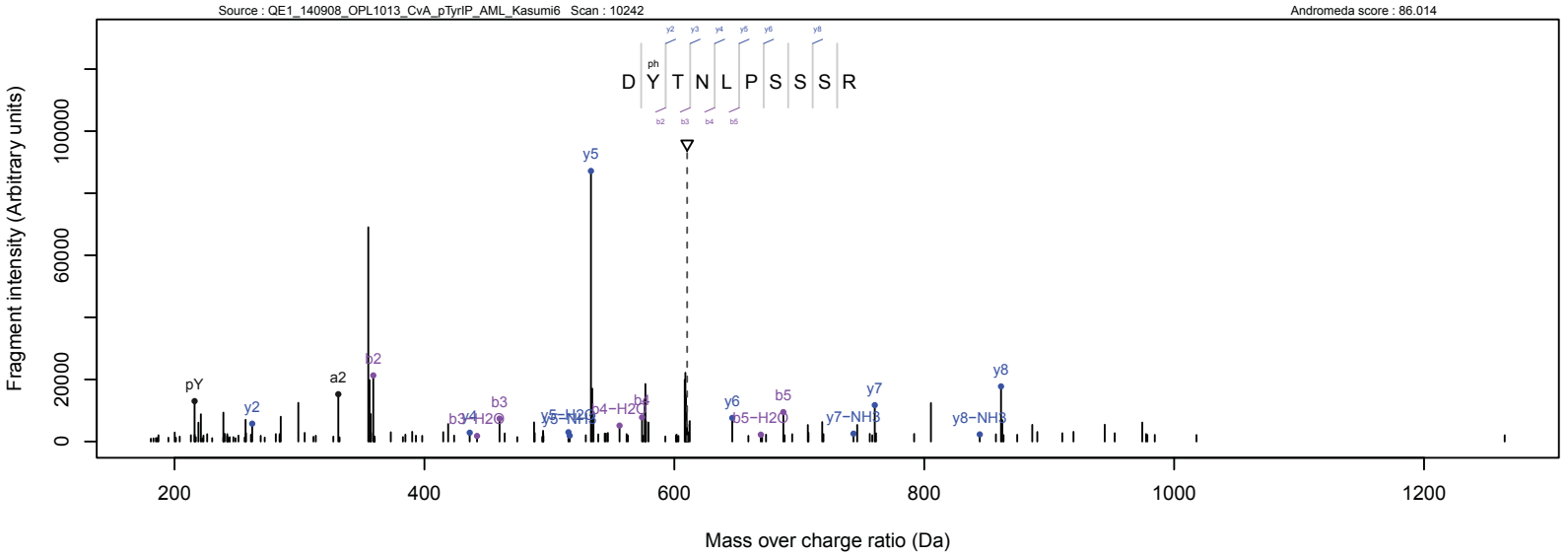
Andromeda score : 86.191



DY(ph)TNLPSSSR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 10242

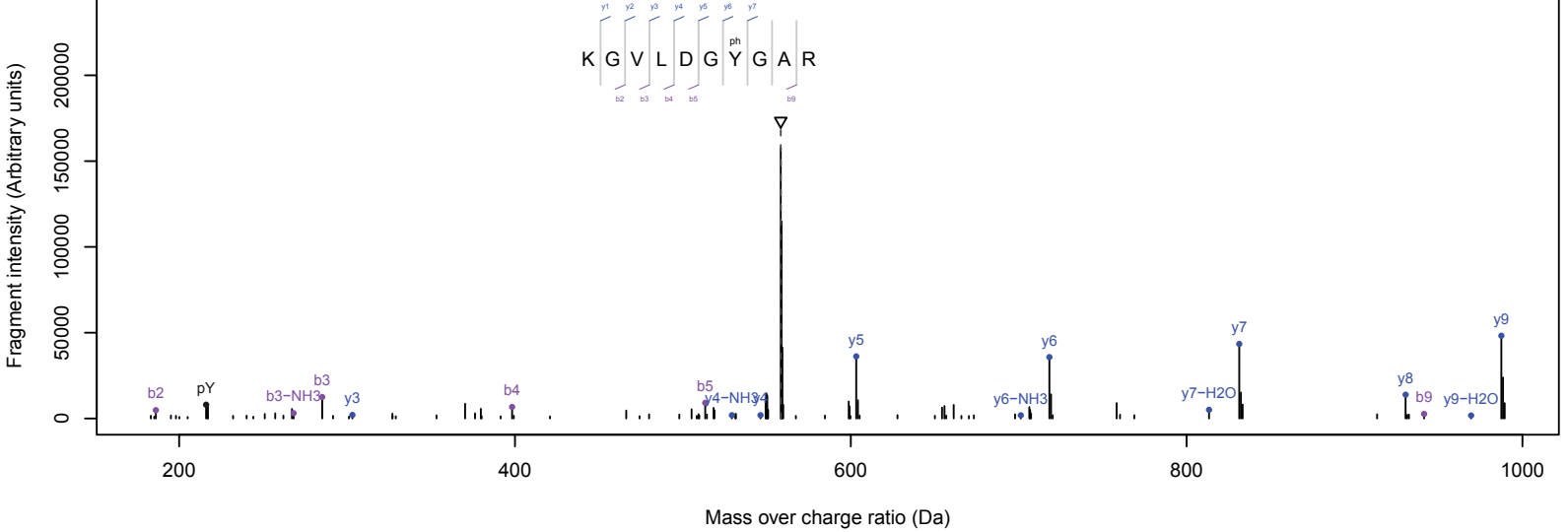
Andromeda score : 86.014



KGVL DGYGAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7976

Andromeda score : 86.014

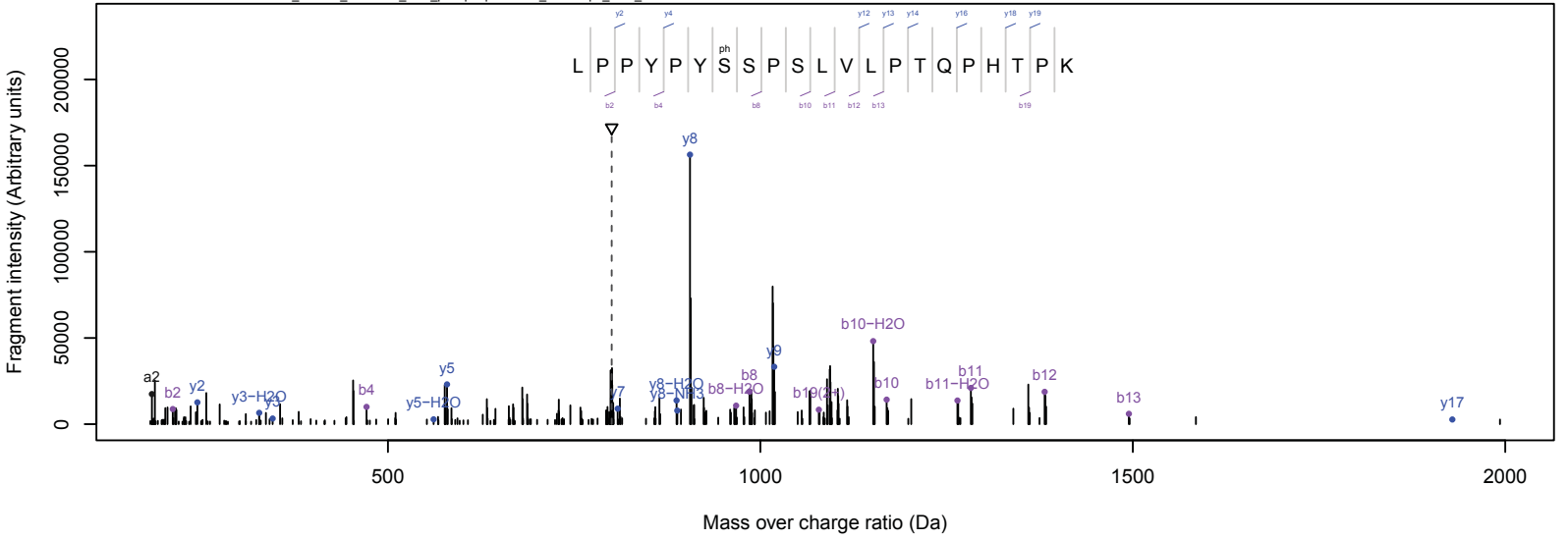




LPPYPYS(ph)SPSLVLPTQPHTPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35674

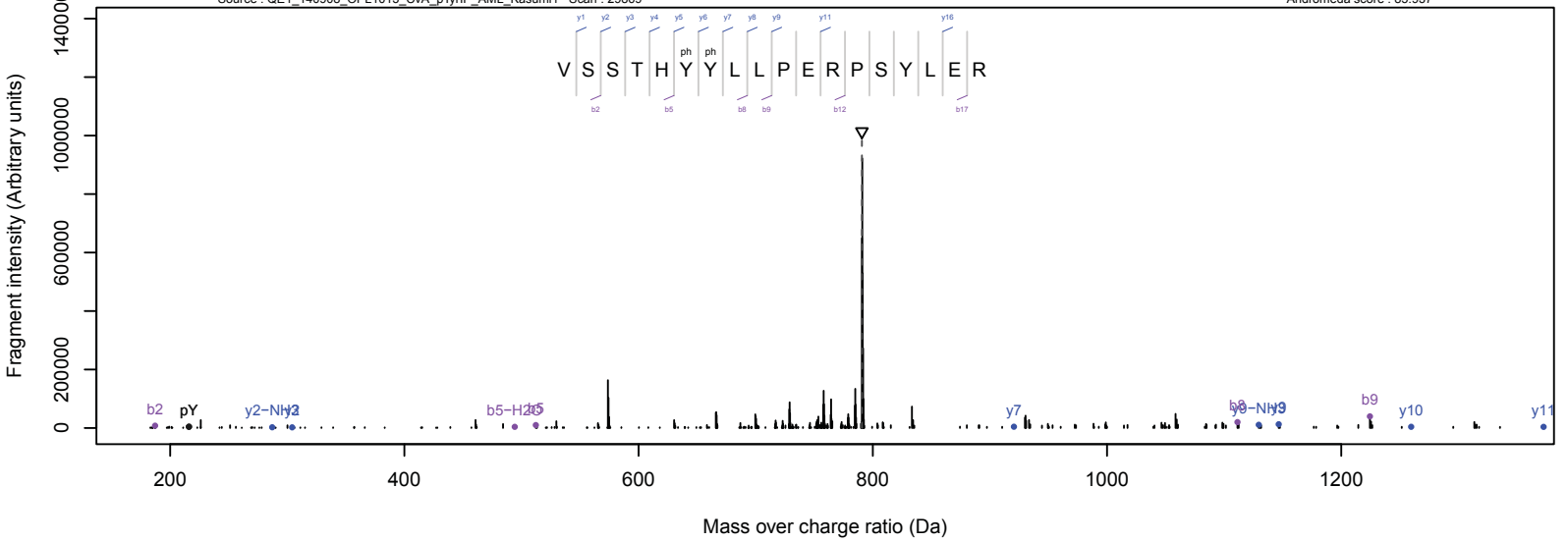
Andromeda score : 85.937



VSSTHY(ph)Y(ph)LLPERPSYLER-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 29869

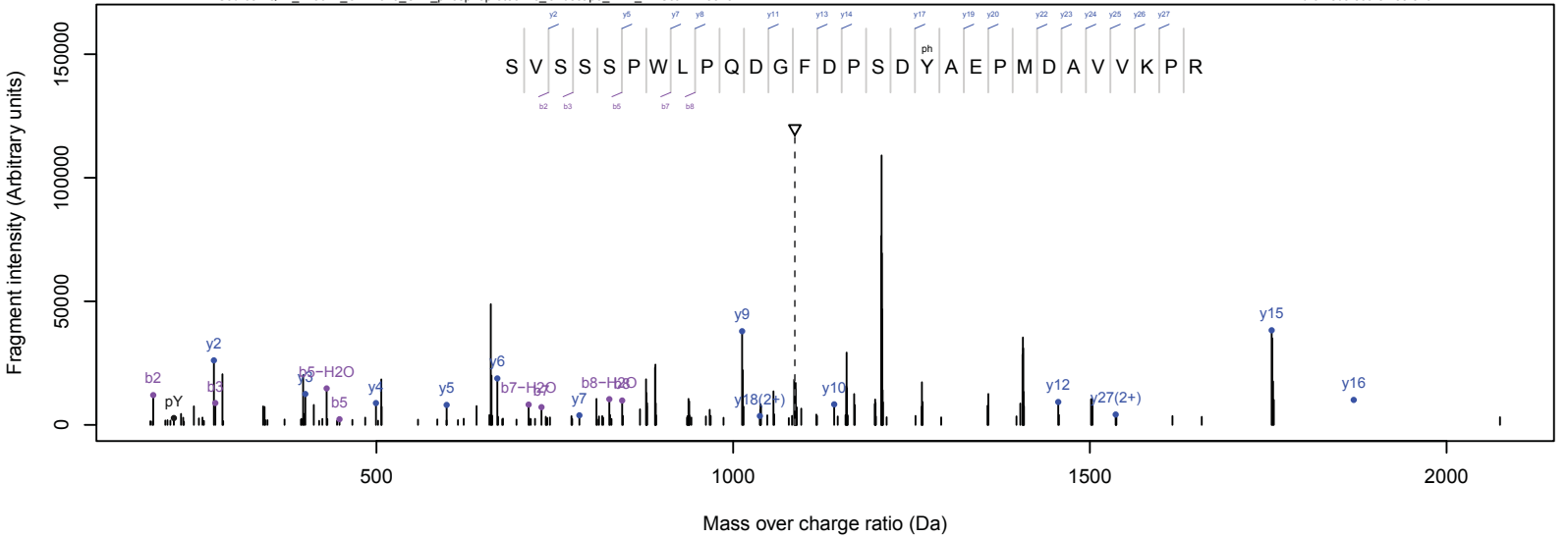
Andromeda score : 85.937



SVSSPWLPQDGFDPSPDY(ph)AEPMDAVVKPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45845

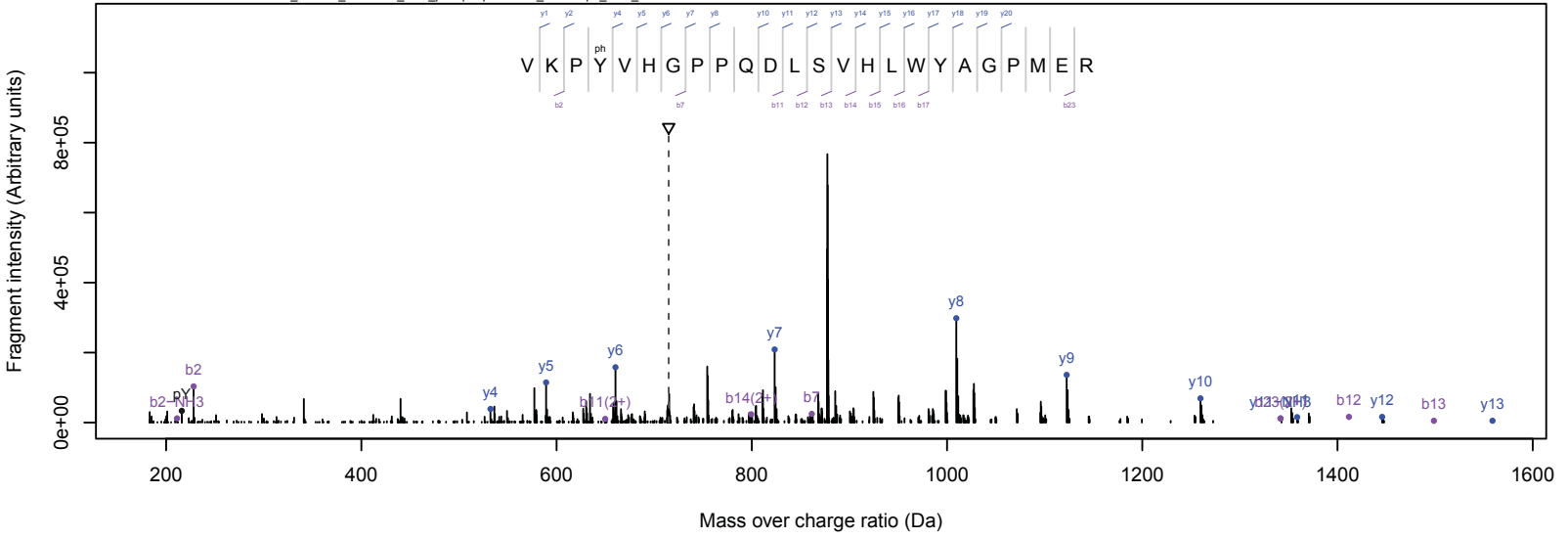
Andromeda score : 85.849



VKPY(ph)VHGGPPQDLSVHLWYAGPMER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36633

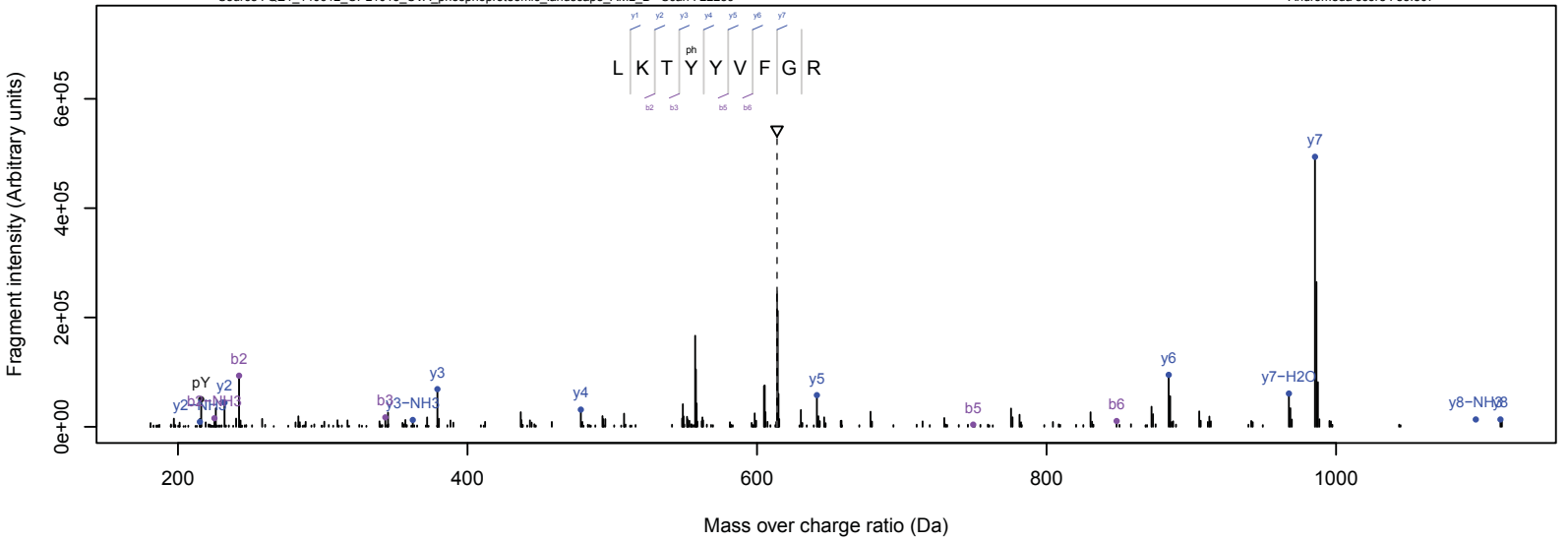
Andromeda score : 85.849



LKTY(ph)YVFG R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22289

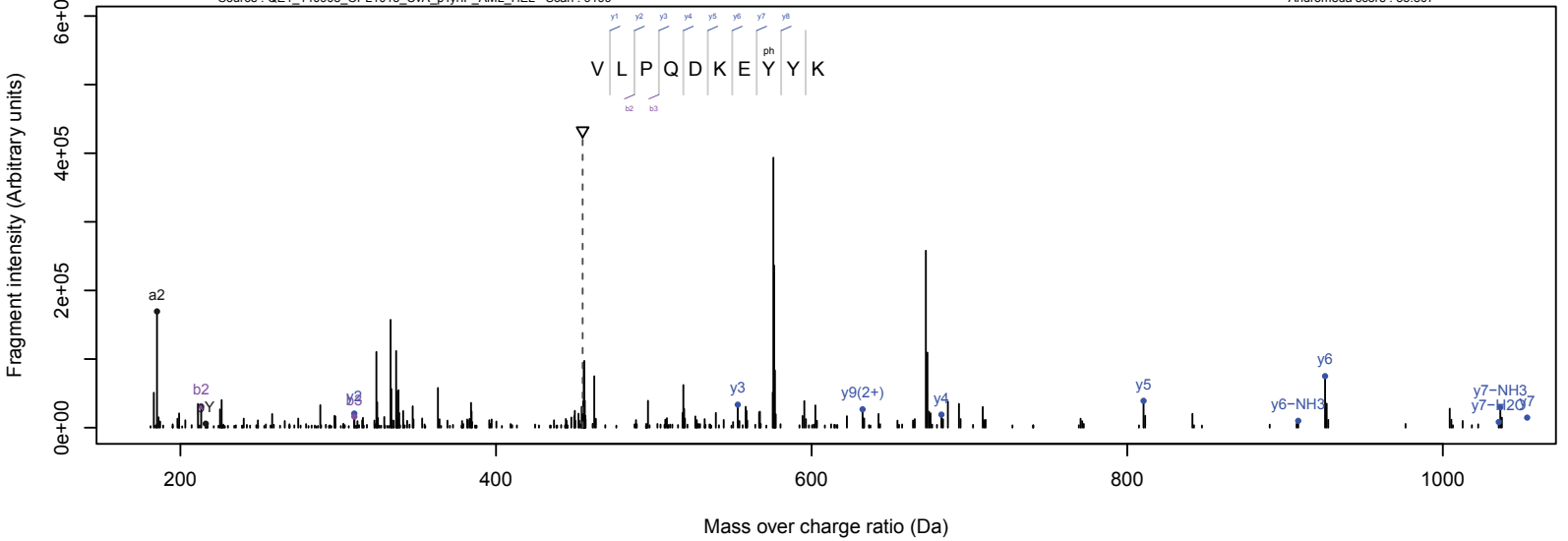
Andromeda score : 85.807



VLPQDKEY(ph)YK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 9199

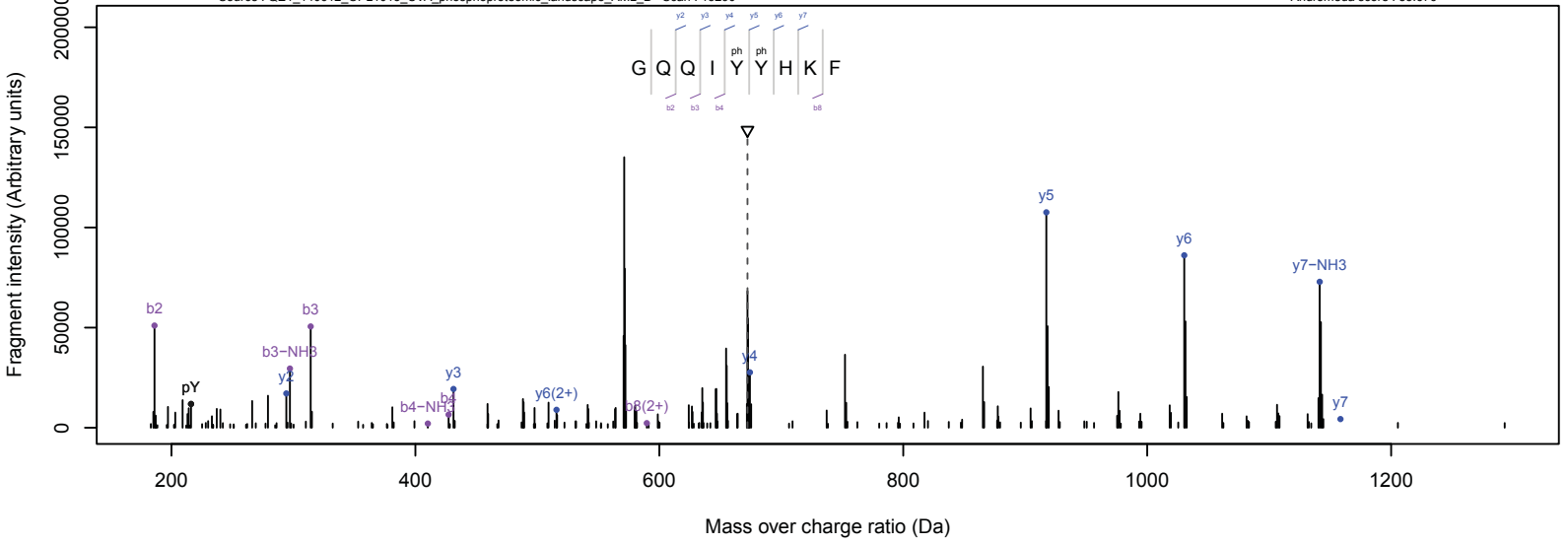
Andromeda score : 85.807



**GQQIY(ph)YHKF-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 13290

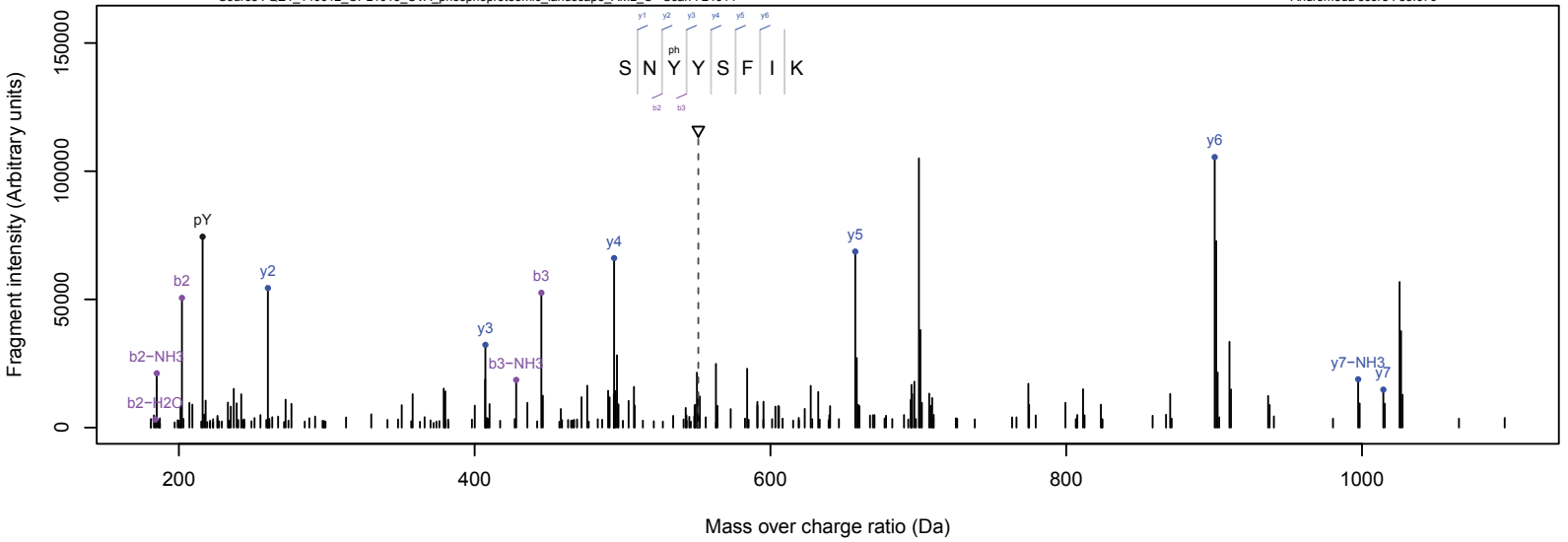
Andromeda score: 85.676



**SNY(ph)YSFIK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24614

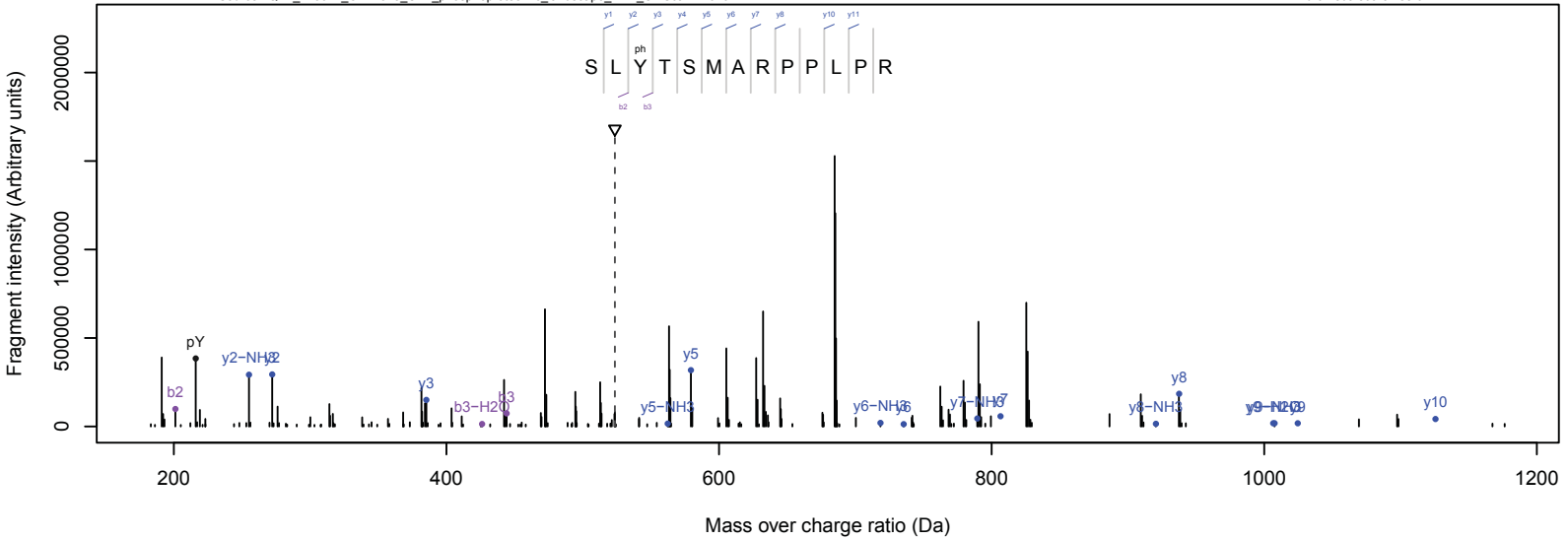
Andromeda score: 85.676



**SLY(ph)TSMARPLPR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26401

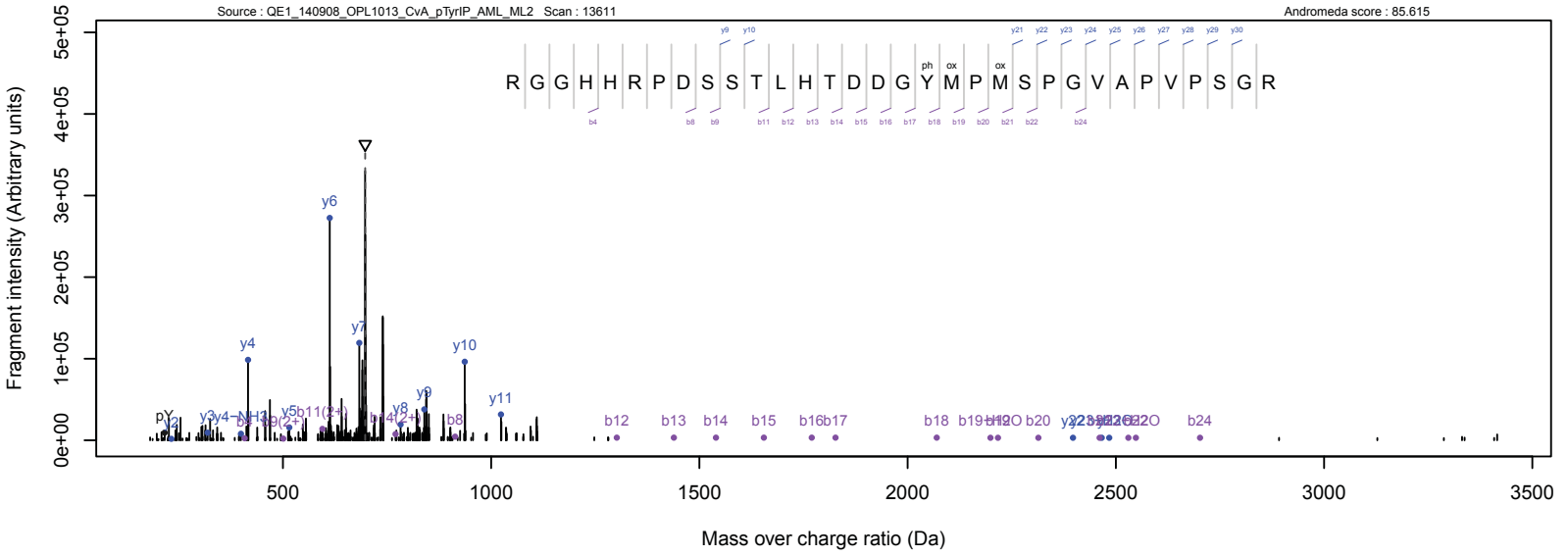
Andromeda score: 85.672



**RGGHRRPDSSTLHTDDGYM(ox)PM(ox)SPGVAPVPSGR-5+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 13611

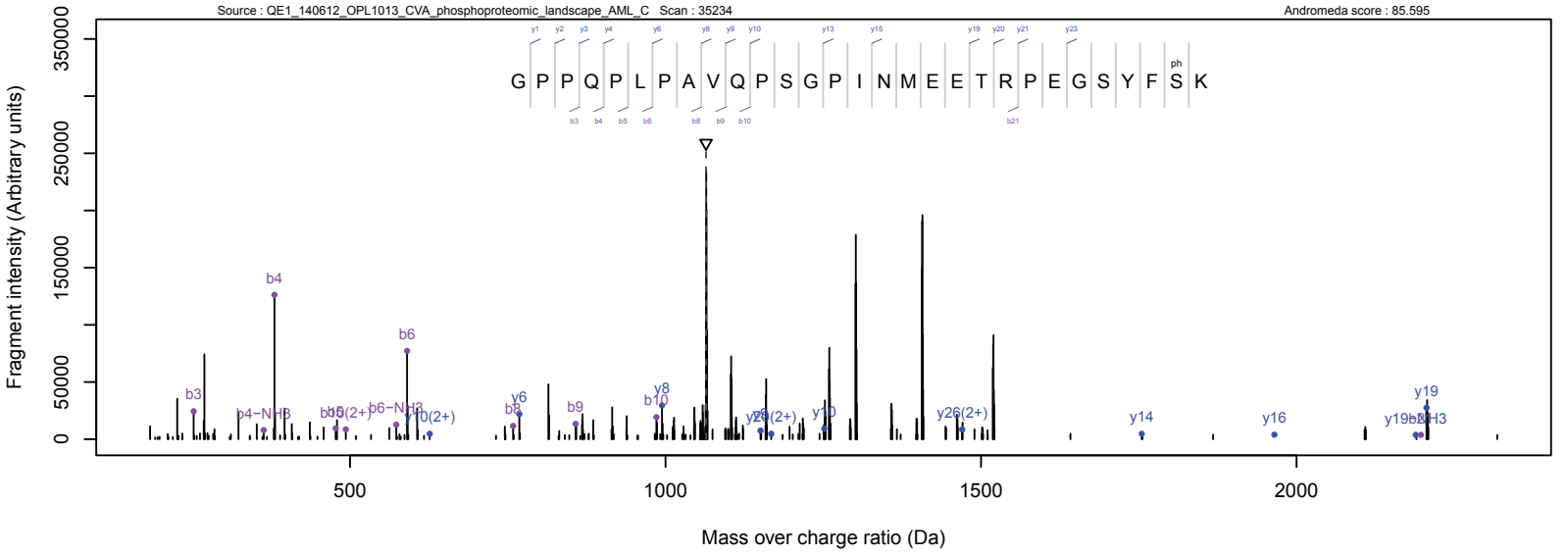
Andromeda score : 85.615



**GPPQLPAVQPSGPIINMEETRPEGSYFS(ph)K-3+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 35234

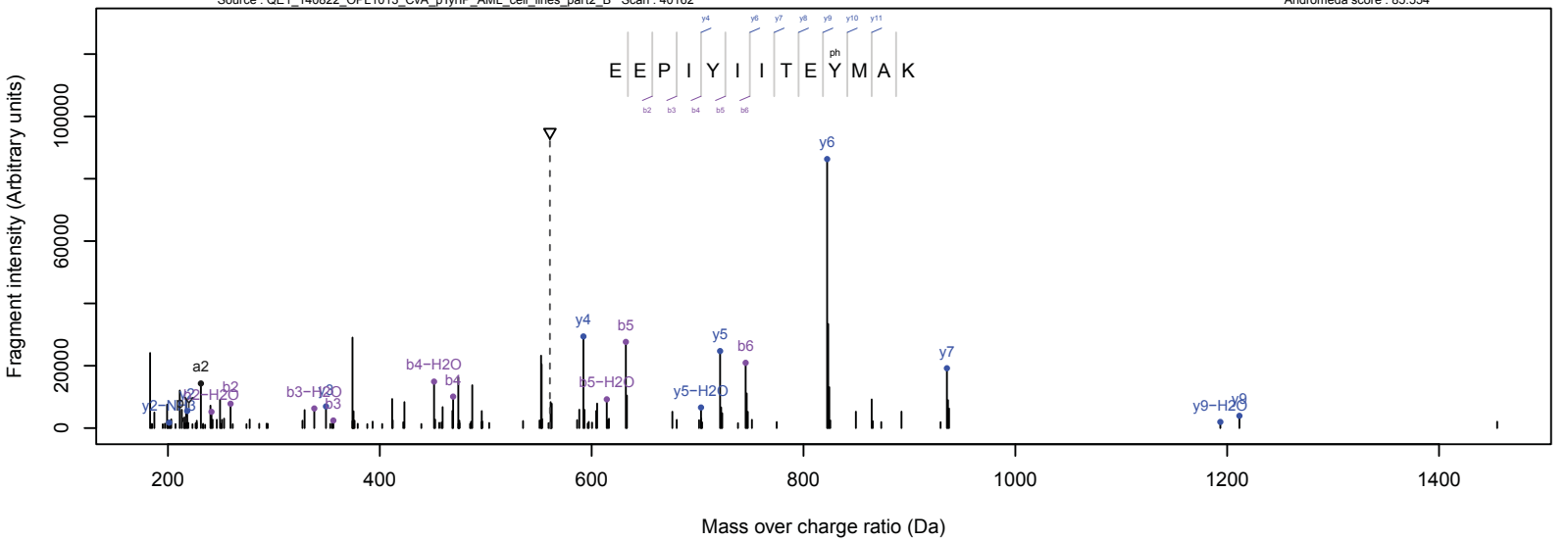
Andromeda score : 85.595



**EEPIIITEYMAK-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 40162

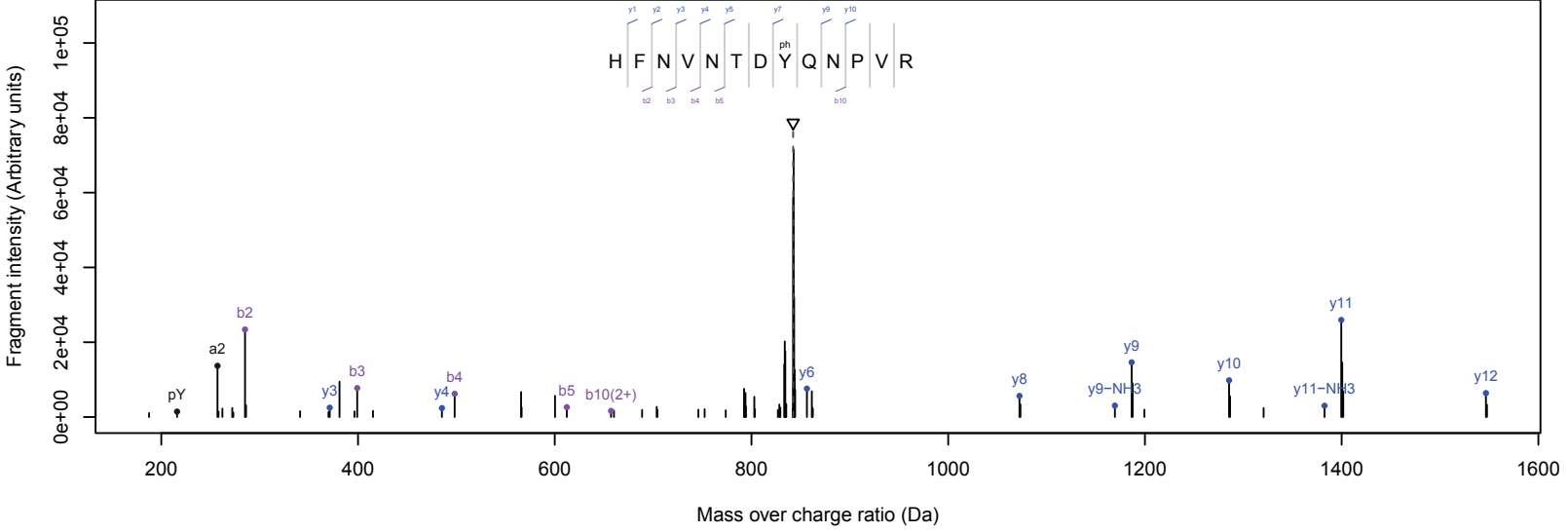
Andromeda score : 85.554



HFNVNTDY(ph)QNPVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15030

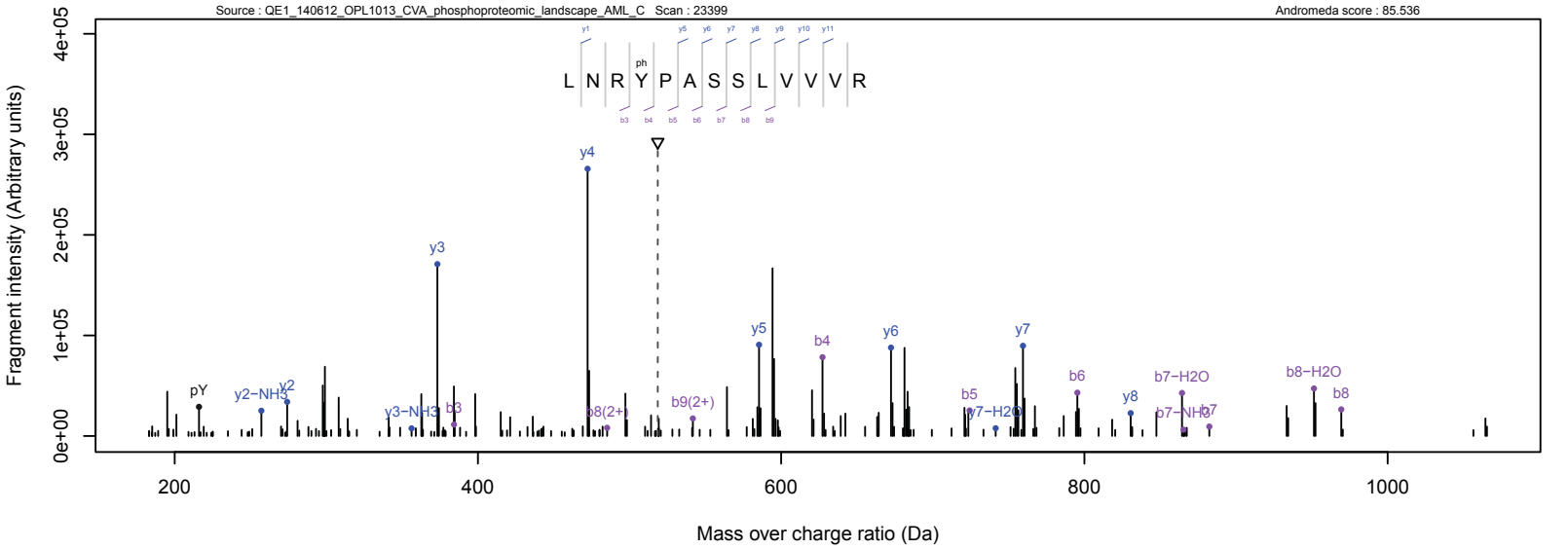
Andromeda score : 85.554



LNRY(ph)PASSLVVVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23399

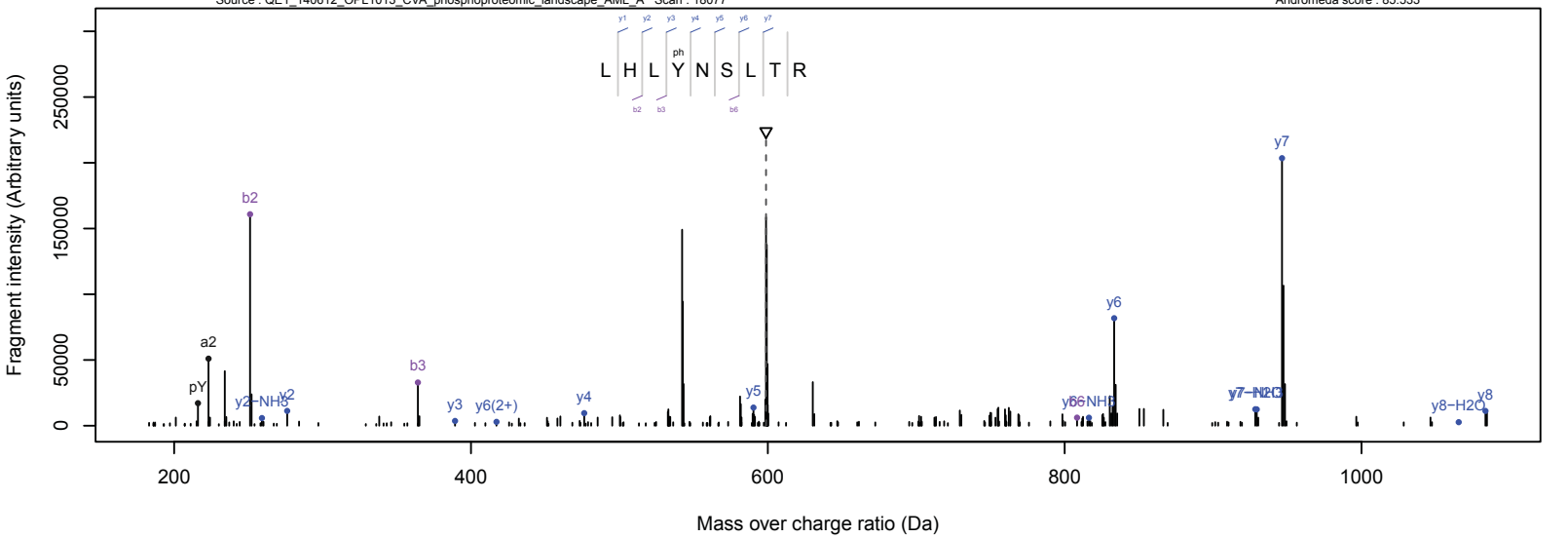
Andromeda score : 85.536



LHLY(ph)NSLTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 18077

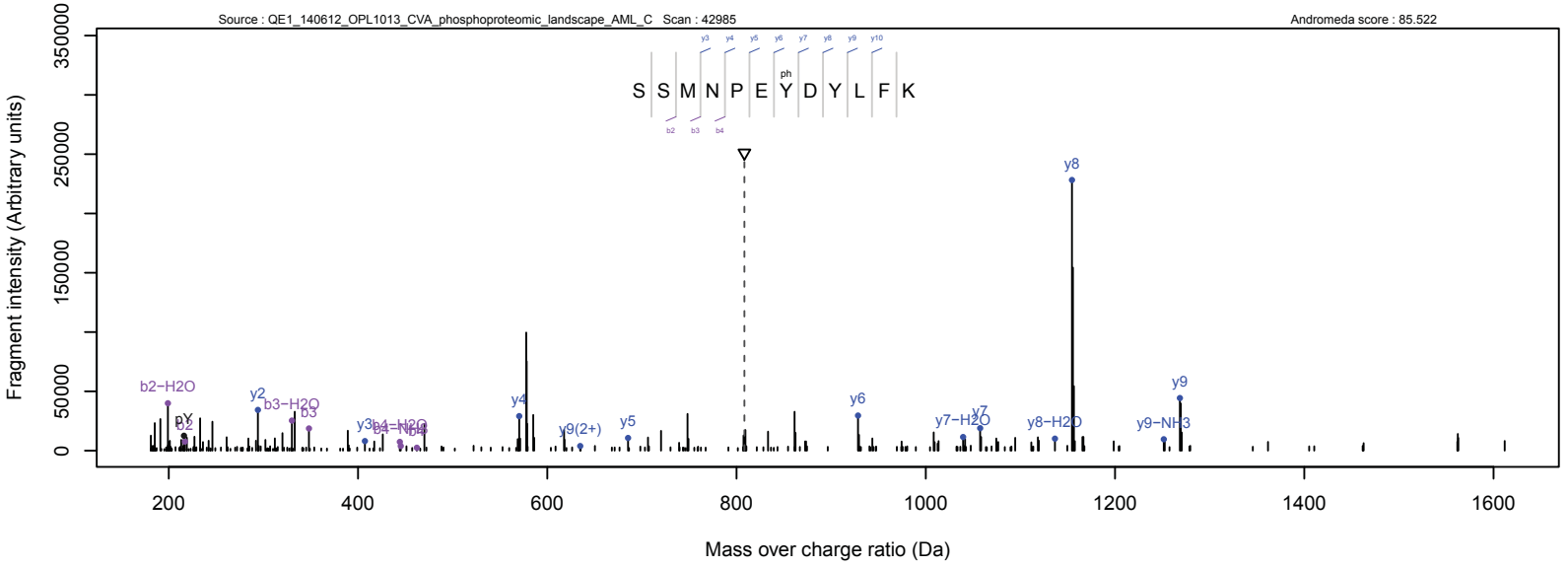
Andromeda score : 85.533



(ac)SSMNPEY(ph)DYLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42985

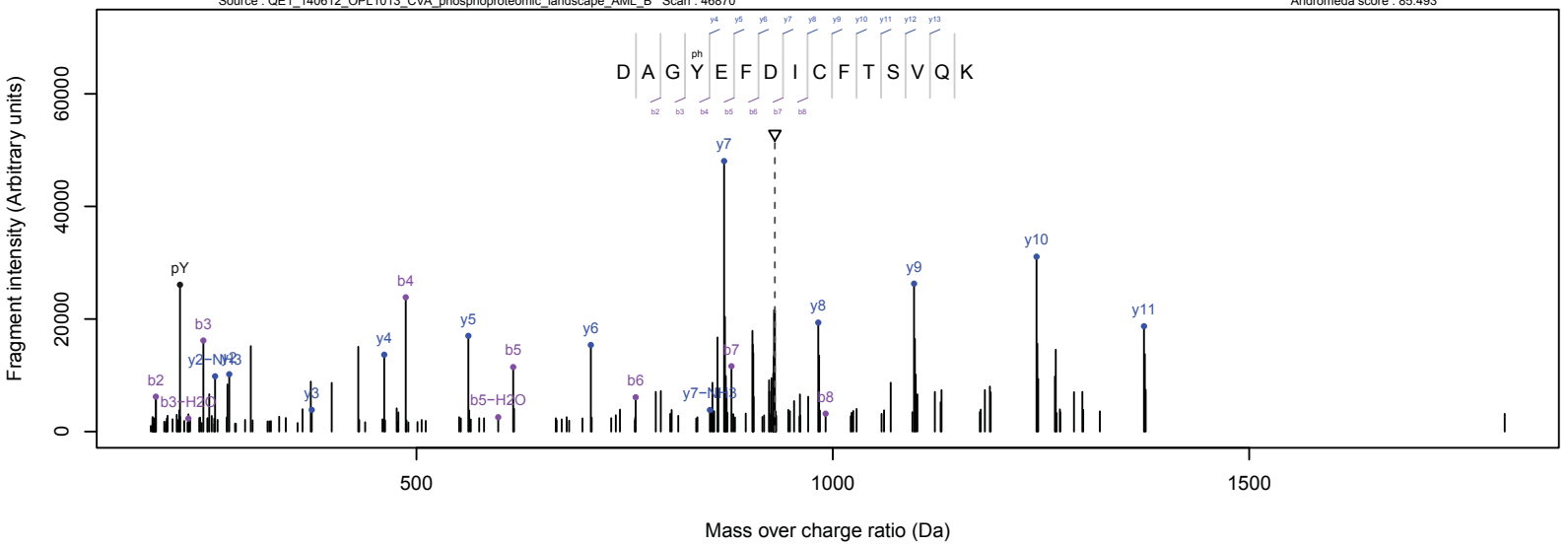
Andromeda score : 85.522



DAGY(ph)EFDICTSVQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46870

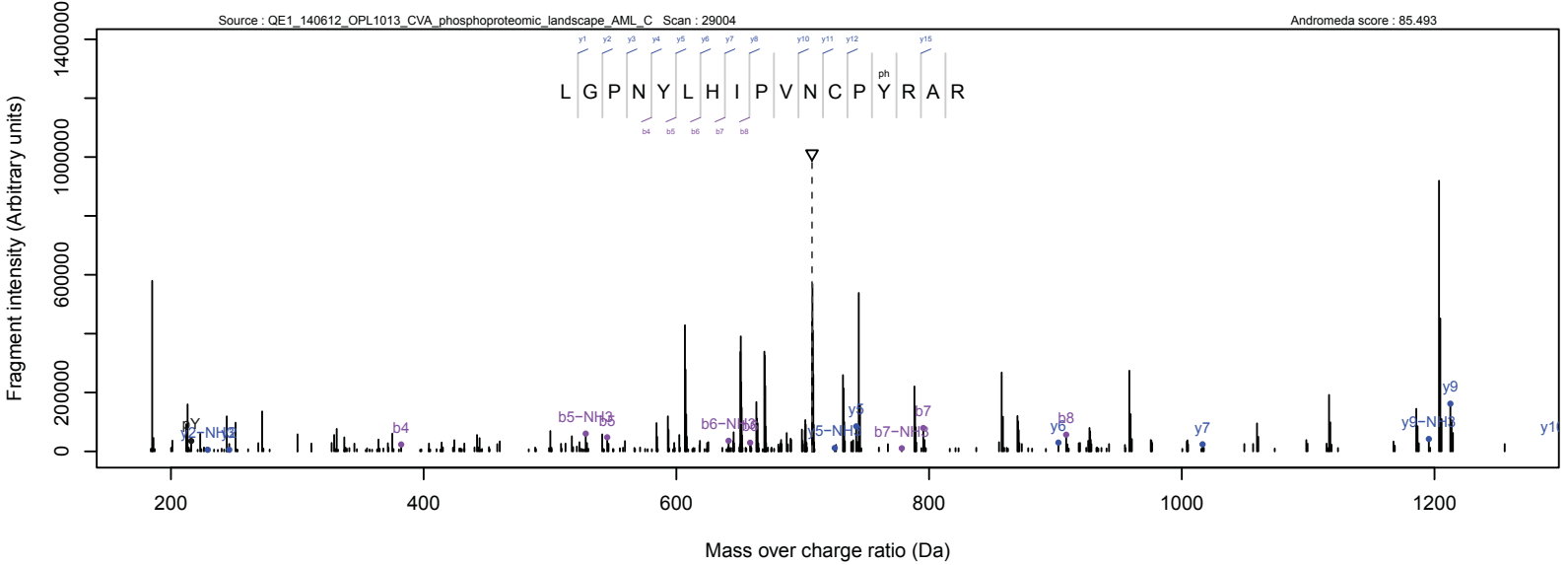
Andromeda score : 85.493



LGPNYLHIPVNC(ph)PYRAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29004

Andromeda score : 85.493

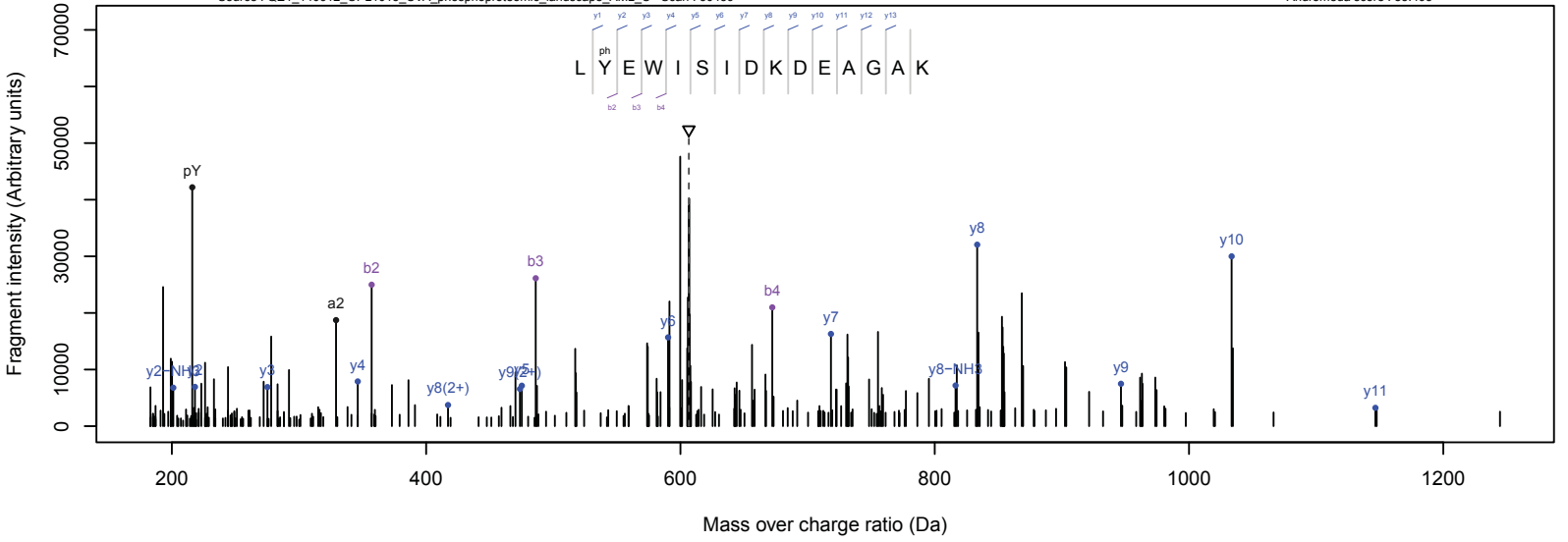




LY(ph)EWISIDKDEAGAK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 30459

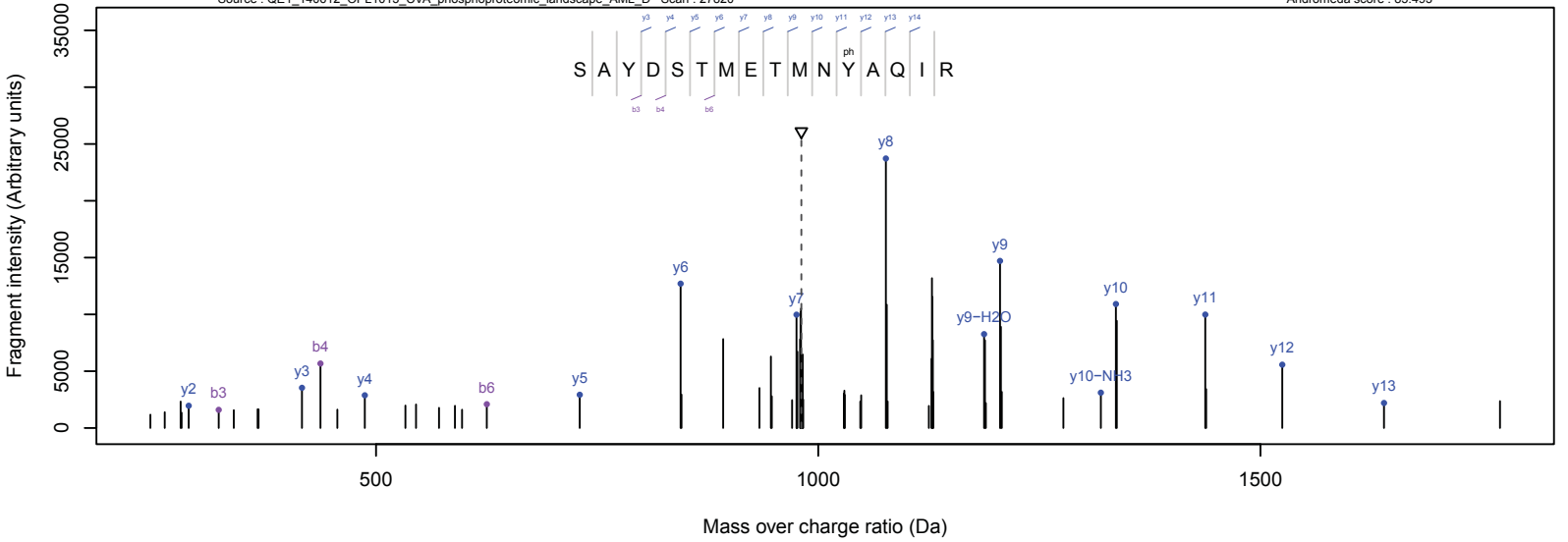
Andromeda score: 85.493



SAYDSTMETMNY(ph)AQIR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 27820

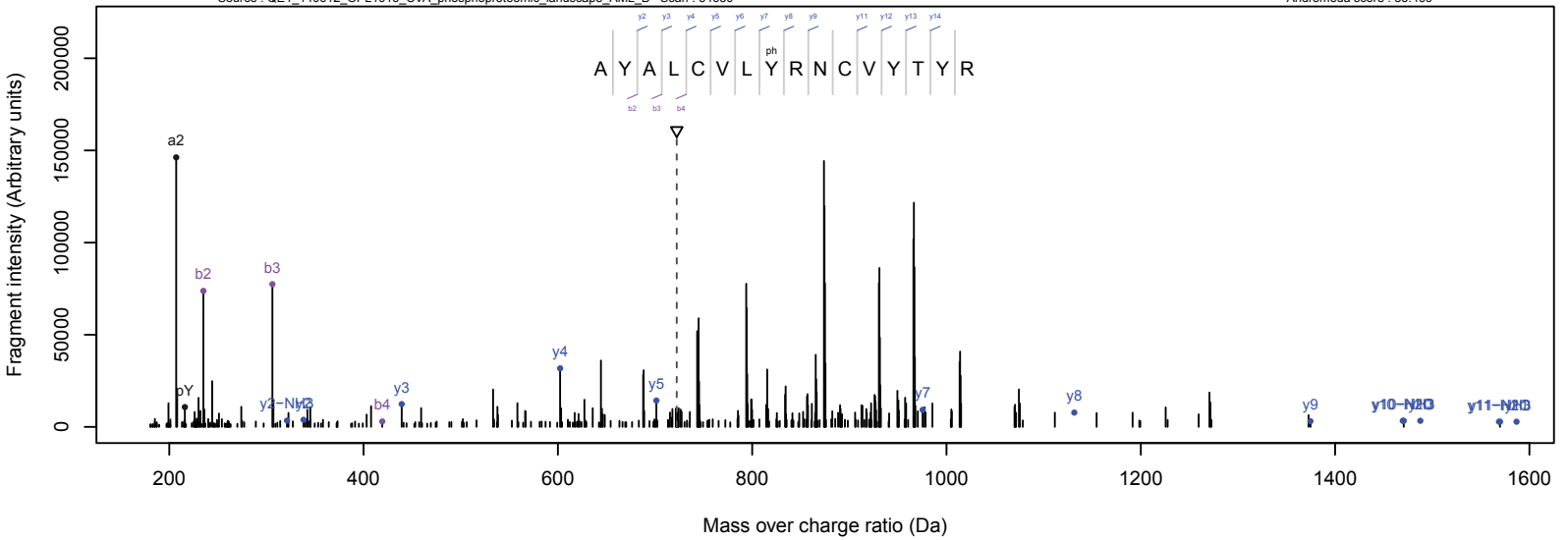
Andromeda score: 85.493



AYALCVLY(ph)RNCVYTYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34680

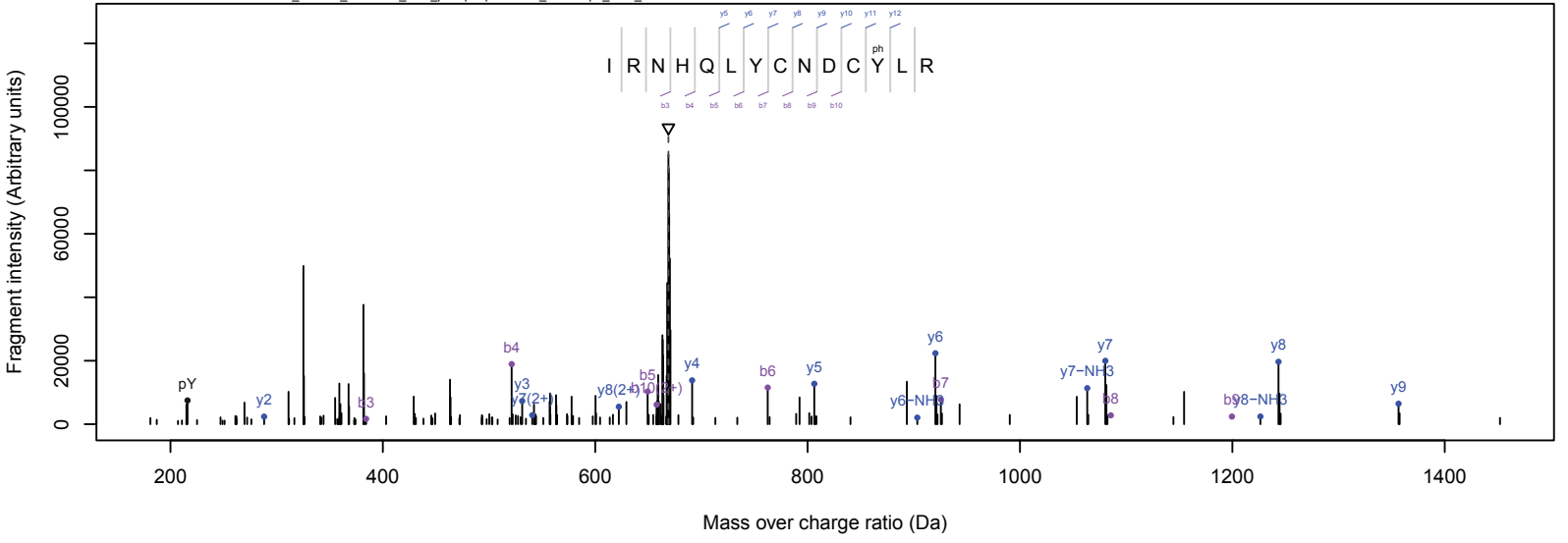
Andromeda score: 85.469



**IRNHQLYCNDCY(ph)LR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13420

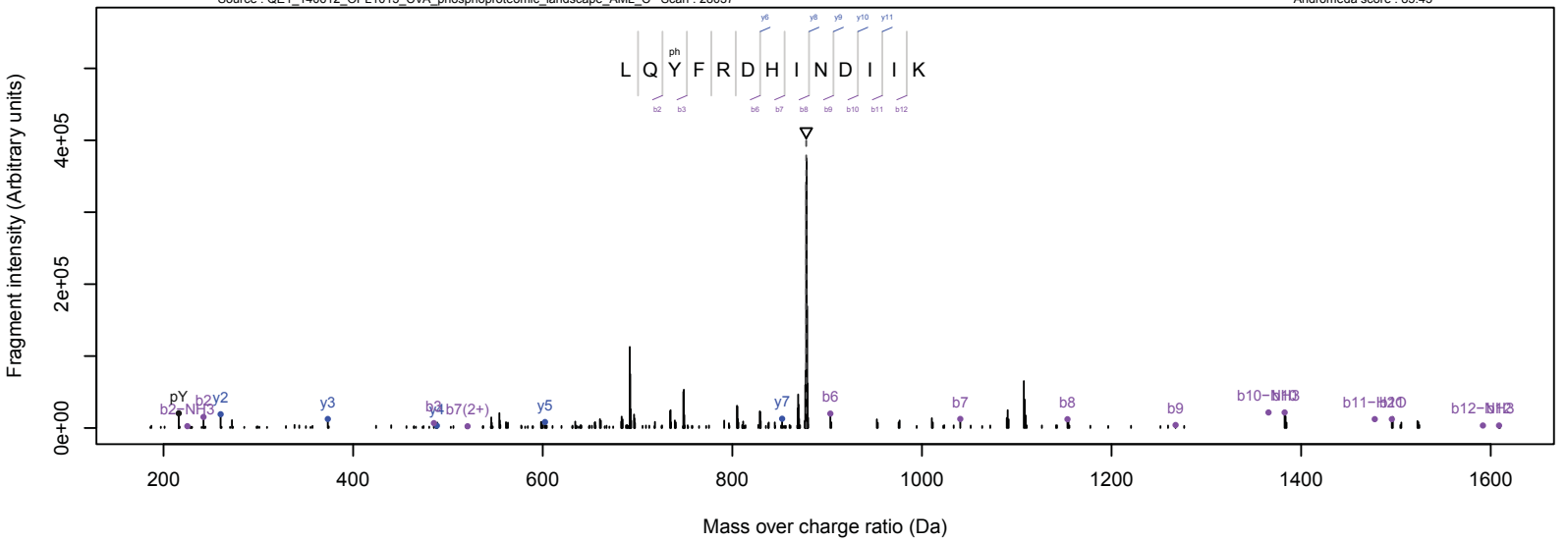
Andromeda score : 85.45



**LQY(ph)FRDHINDIIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28037

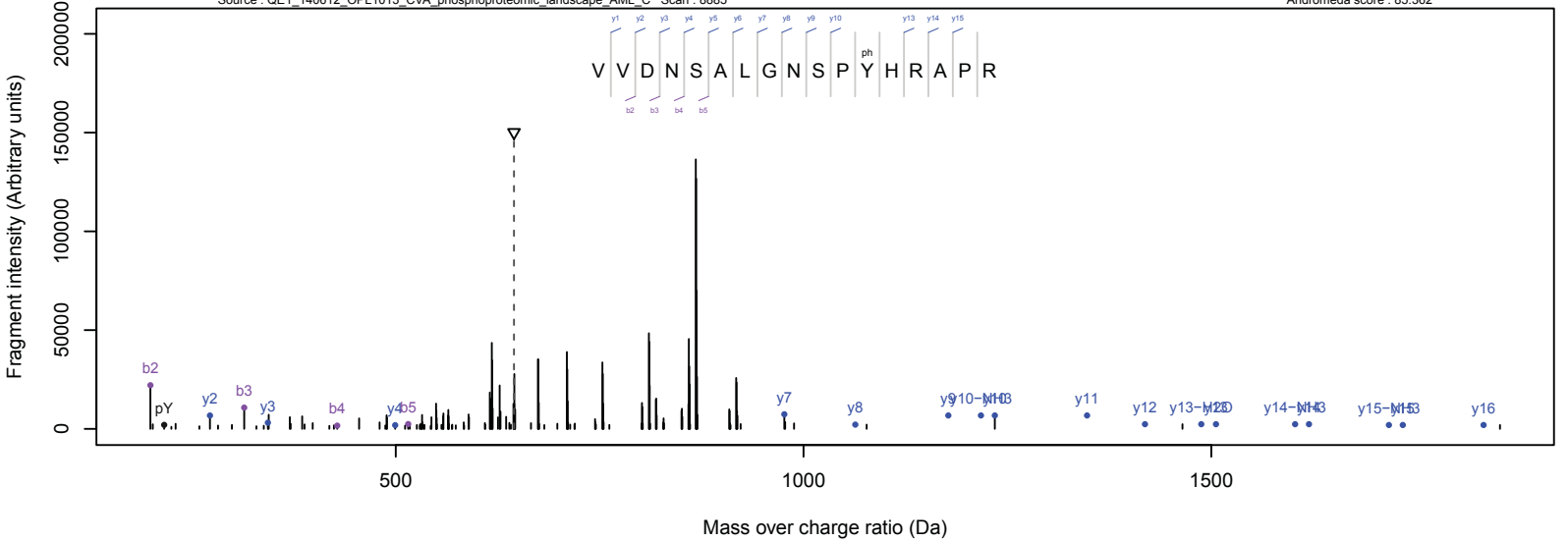
Andromeda score : 85.45



**VVDNSALGNSPY(ph)HRAPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8885

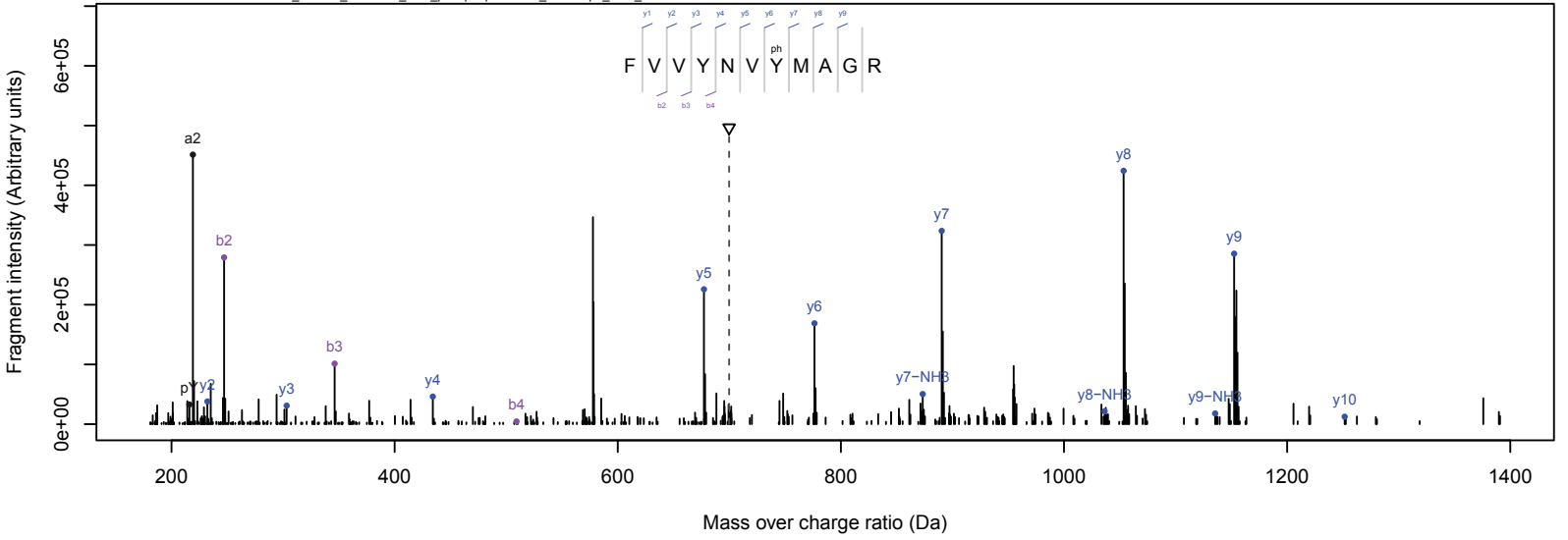
Andromeda score : 85.362



**FVVYNY(ph)MAGR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 32985

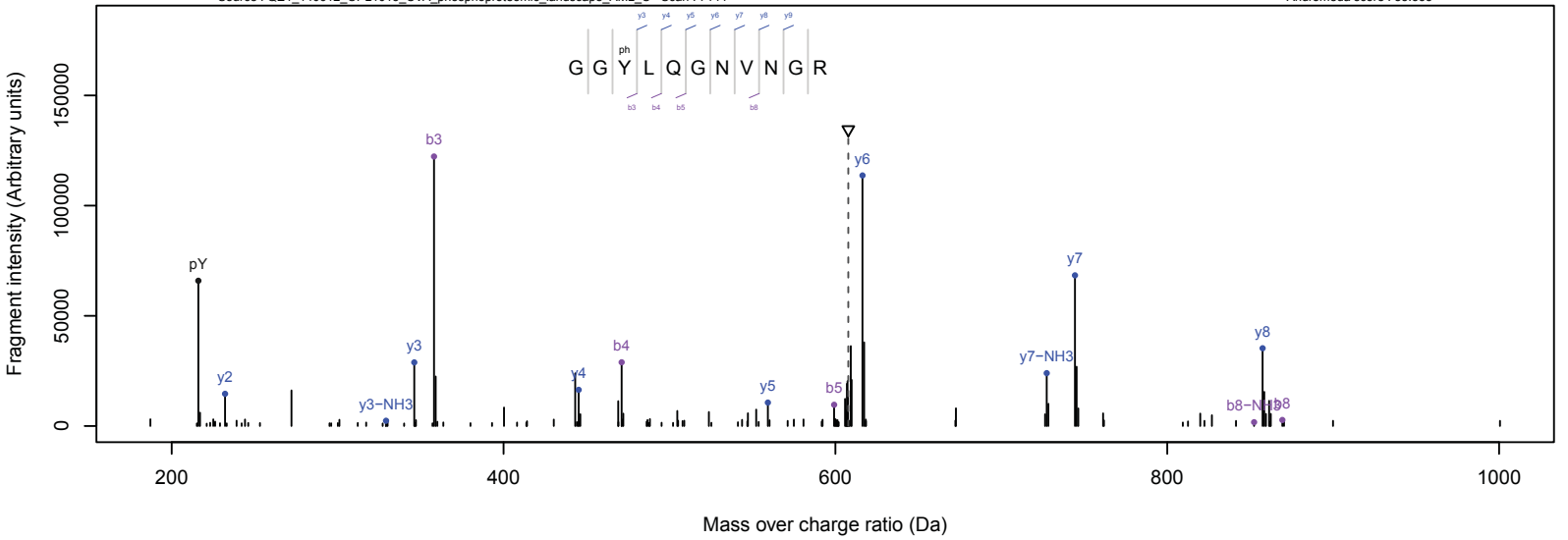
Andromeda score: 85.355



**GGY(ph)LQGNVNGR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 7144

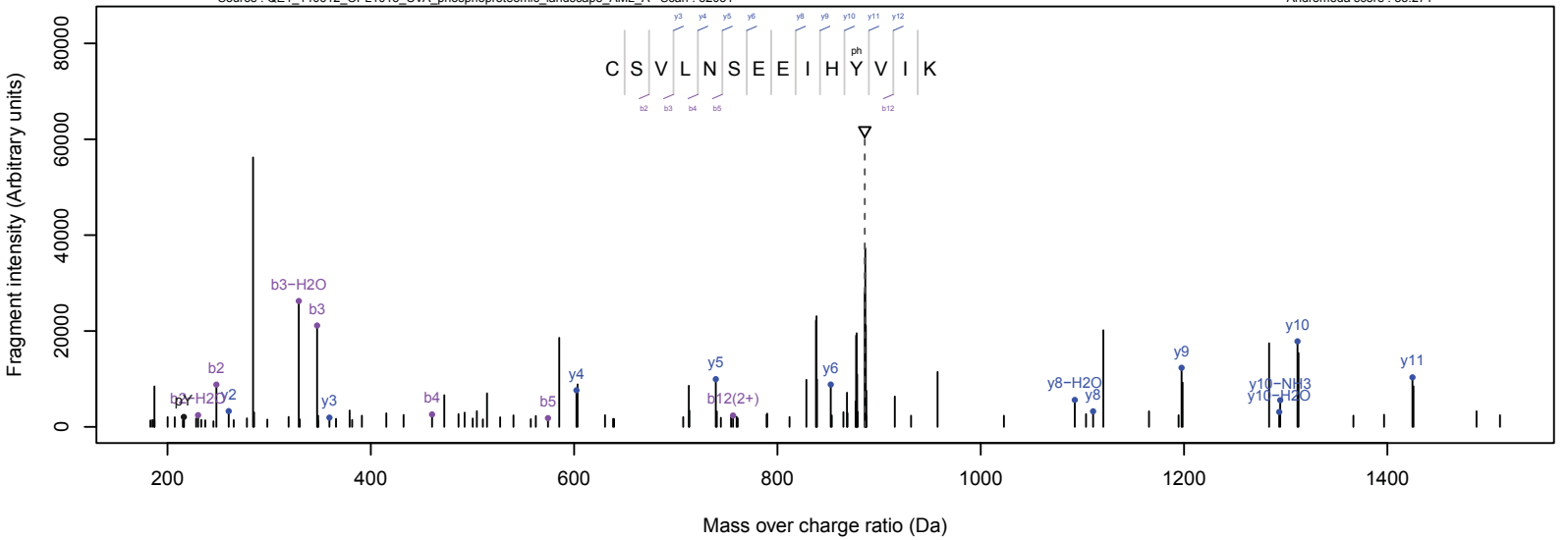
Andromeda score: 85.355



**CSVLNSEEIHY(ph)VIK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 32051

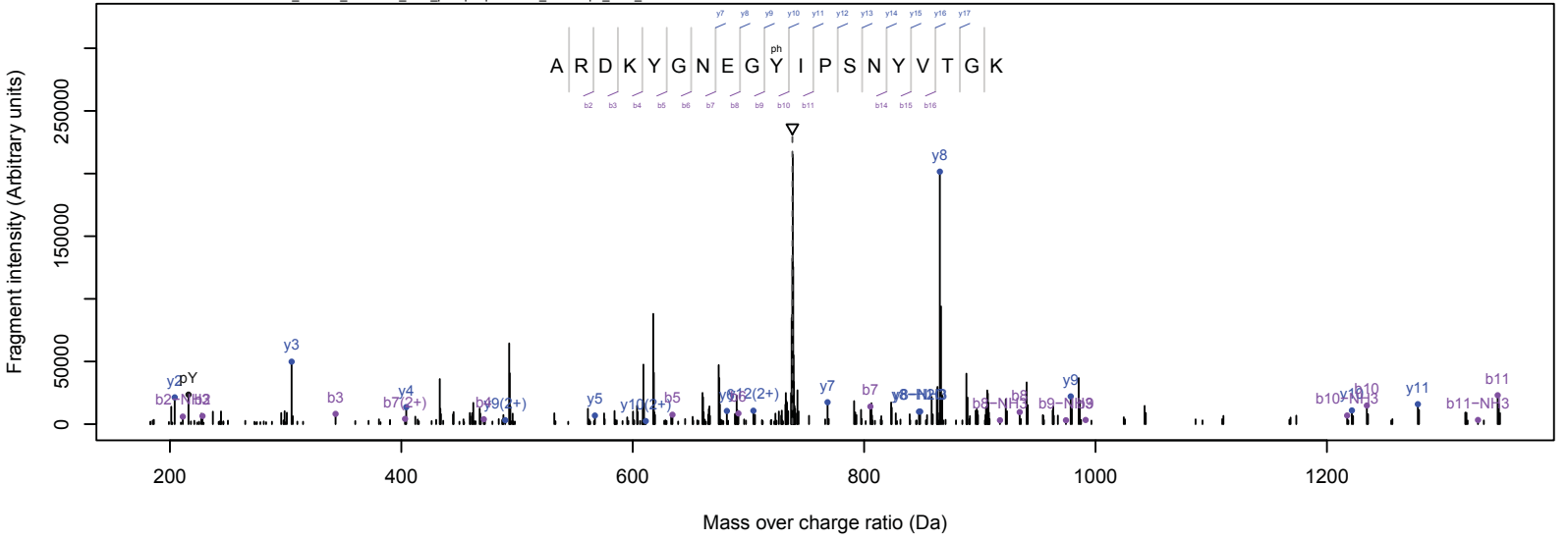
Andromeda score: 85.271



ARDKYGNEGY(ph)IPSNYVTGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17594

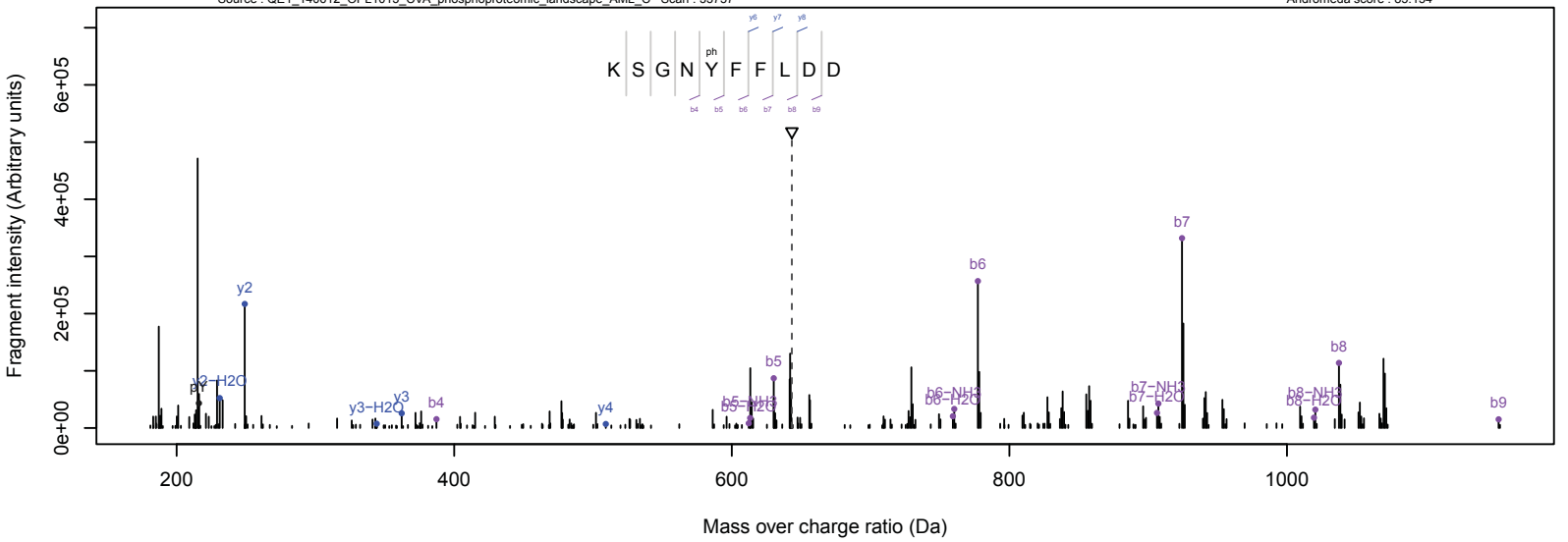
Andromeda score : 85.185



KSGNY(ph)FFLDD-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33757

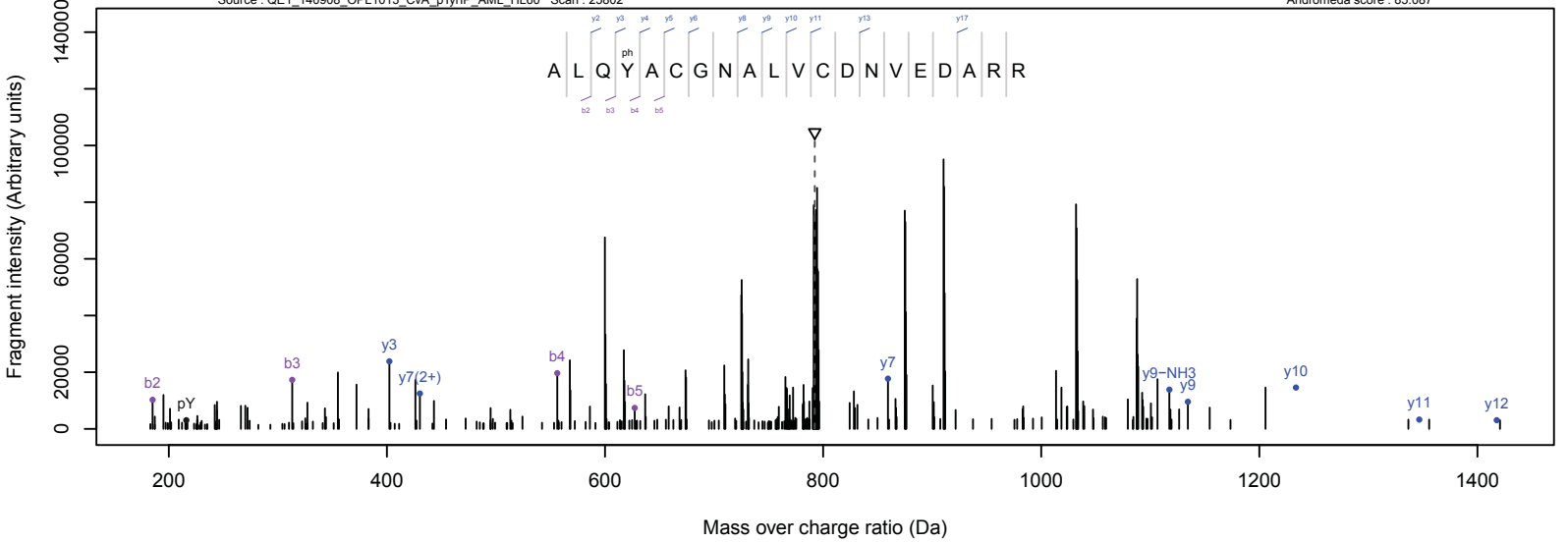
Andromeda score : 85.154



ALQY(ph)ACGNALVCDNVEDARR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 25802

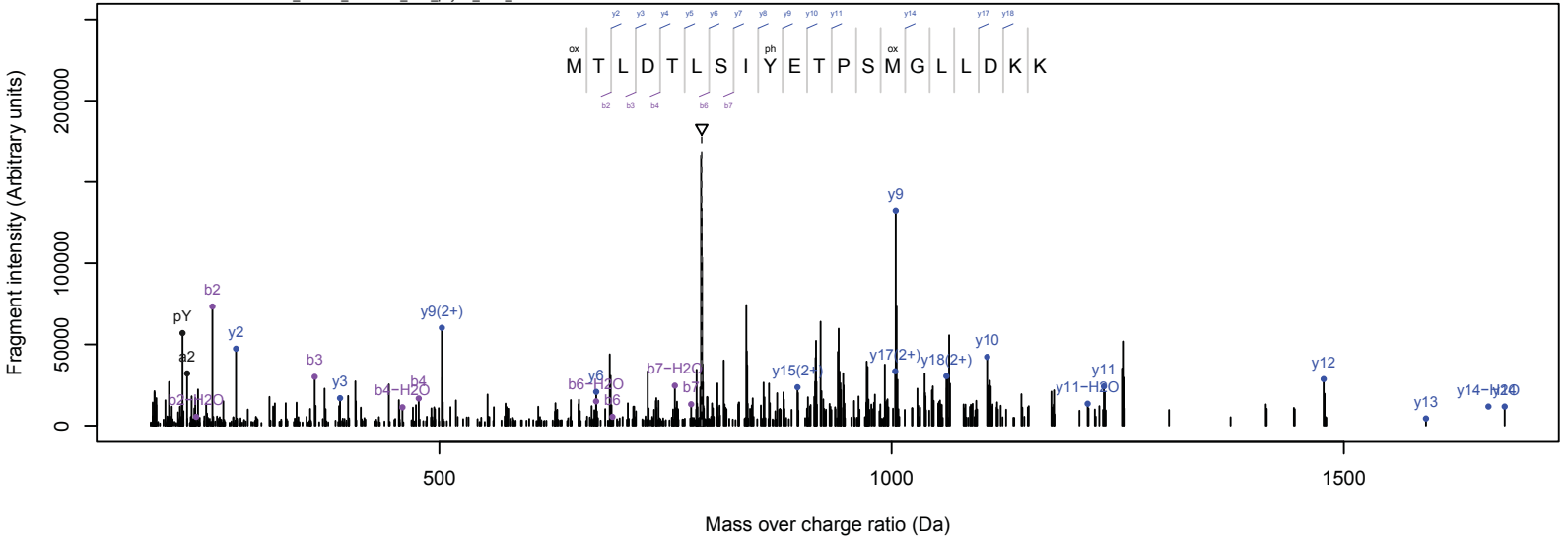
Andromeda score : 85.087



**M(ox)TLDLSIY(ph)ETPSM(ox)GLLDKK-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan : 38398

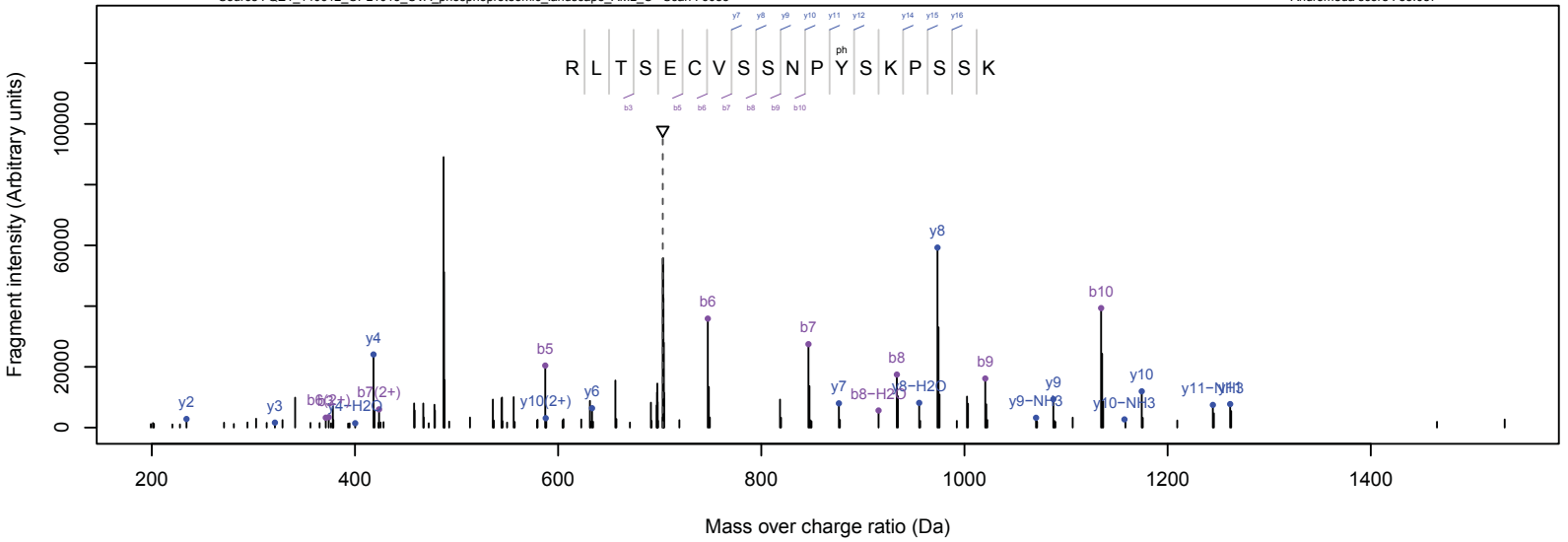
Andromeda score : 85.087



**RLTSECVSSNPY(ph)SKPSSK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6085

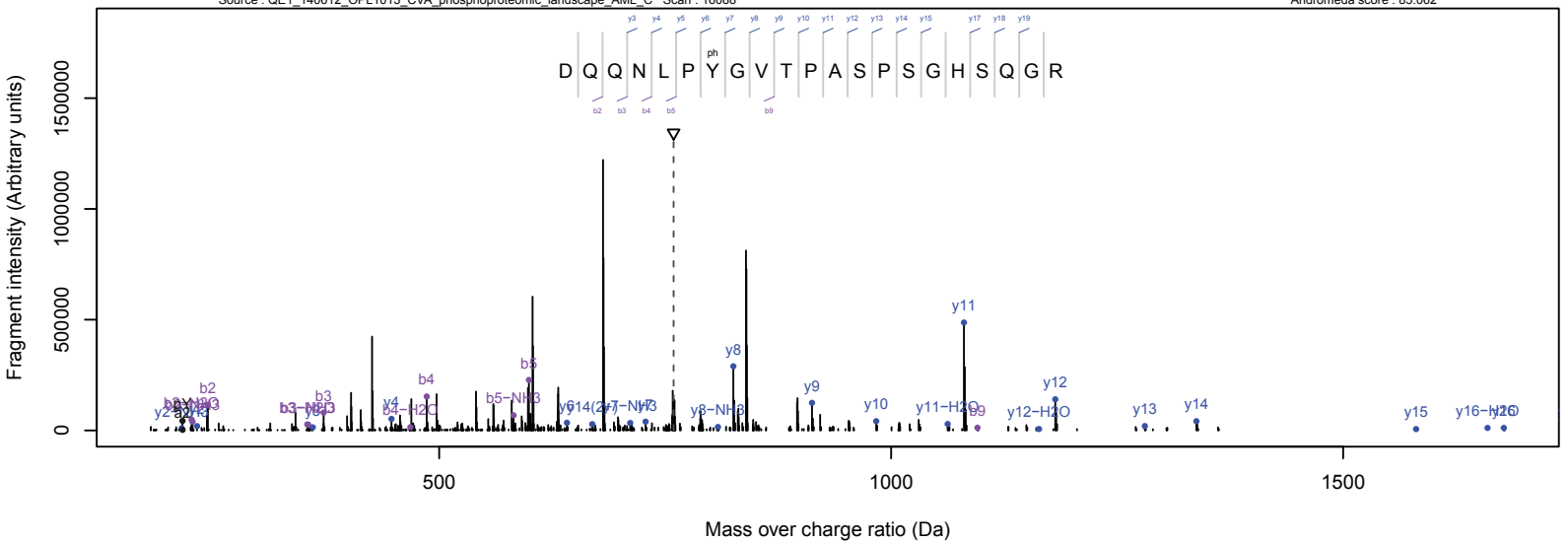
Andromeda score : 85.087



**DQQNLPY(ph)GVTPASPSGHSQGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16088

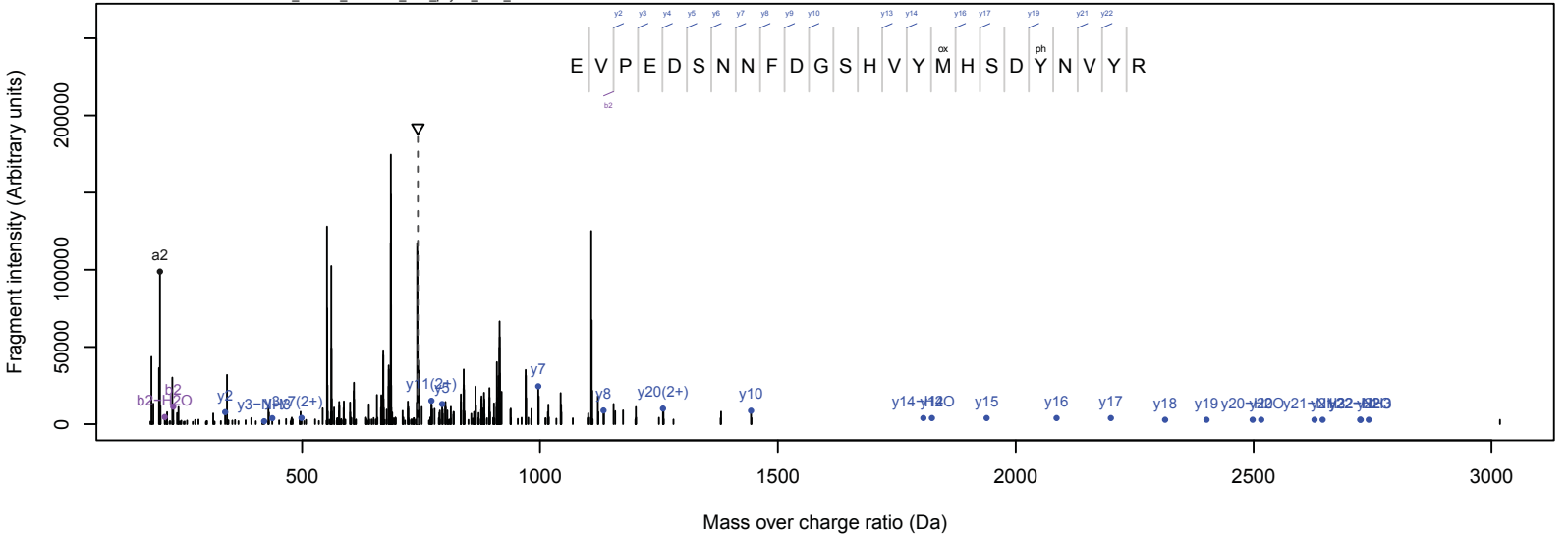
Andromeda score : 85.062



EVPEDSNNFDGSHVYM(ox)HSDY(ph)NVYR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 22840

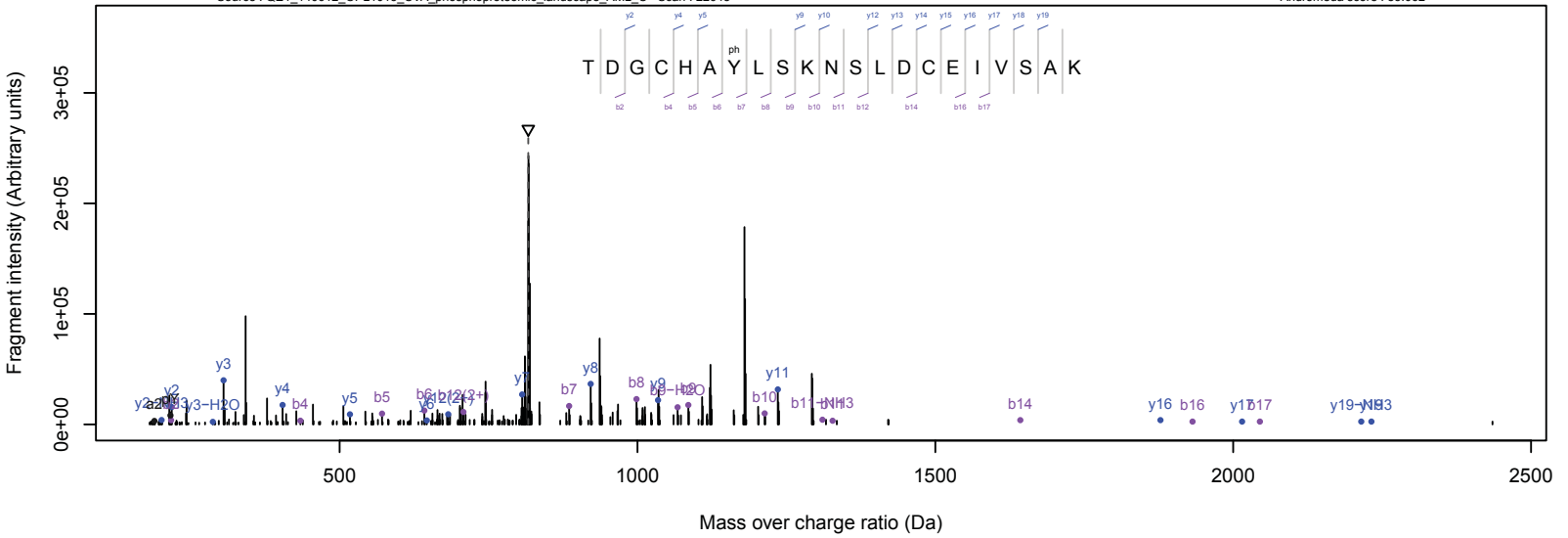
Andromeda score : 85.062



TDGCHAY(ph)LSKNSLDCEIVSAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22648

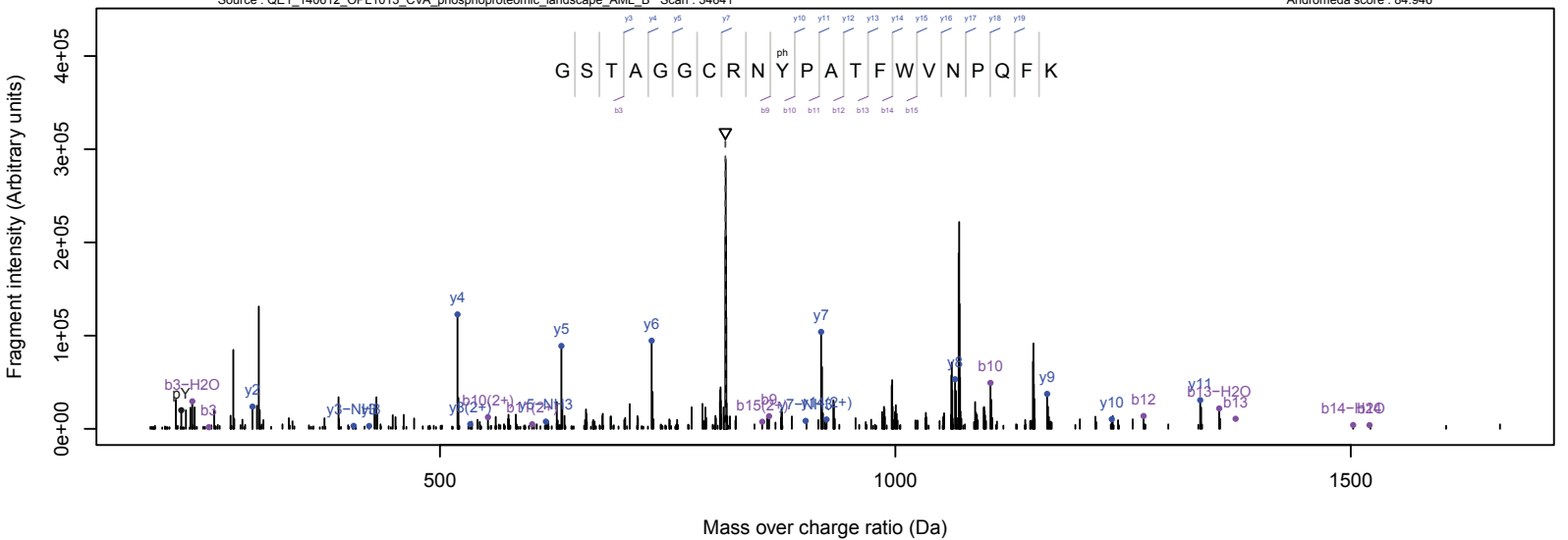
Andromeda score : 85.062



GSTAGGCRNY(ph)PATFWVNPQFK-3+

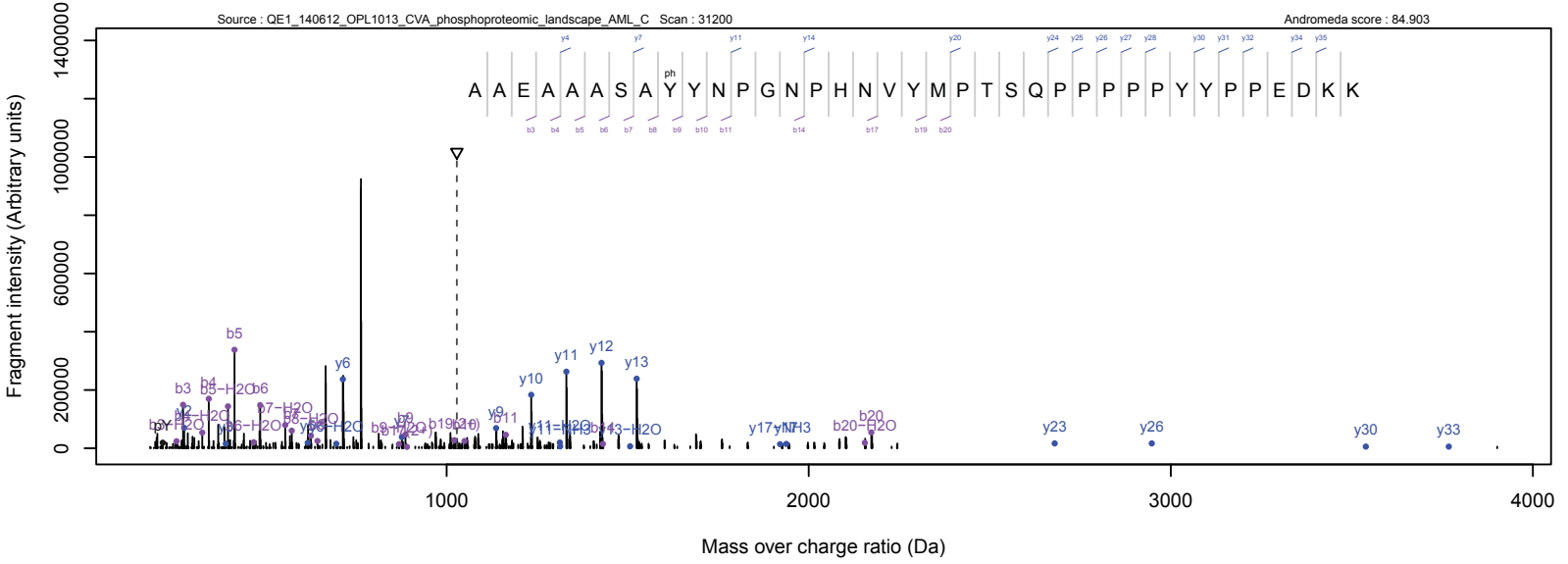
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34641

Andromeda score : 84.946

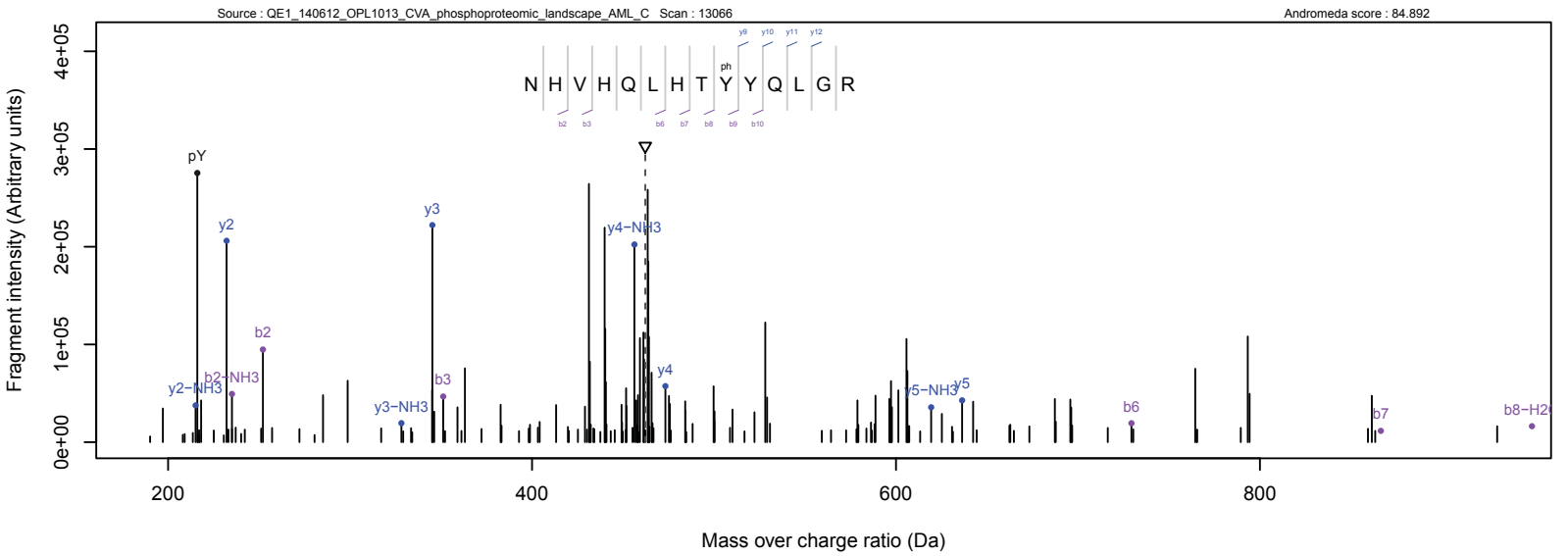




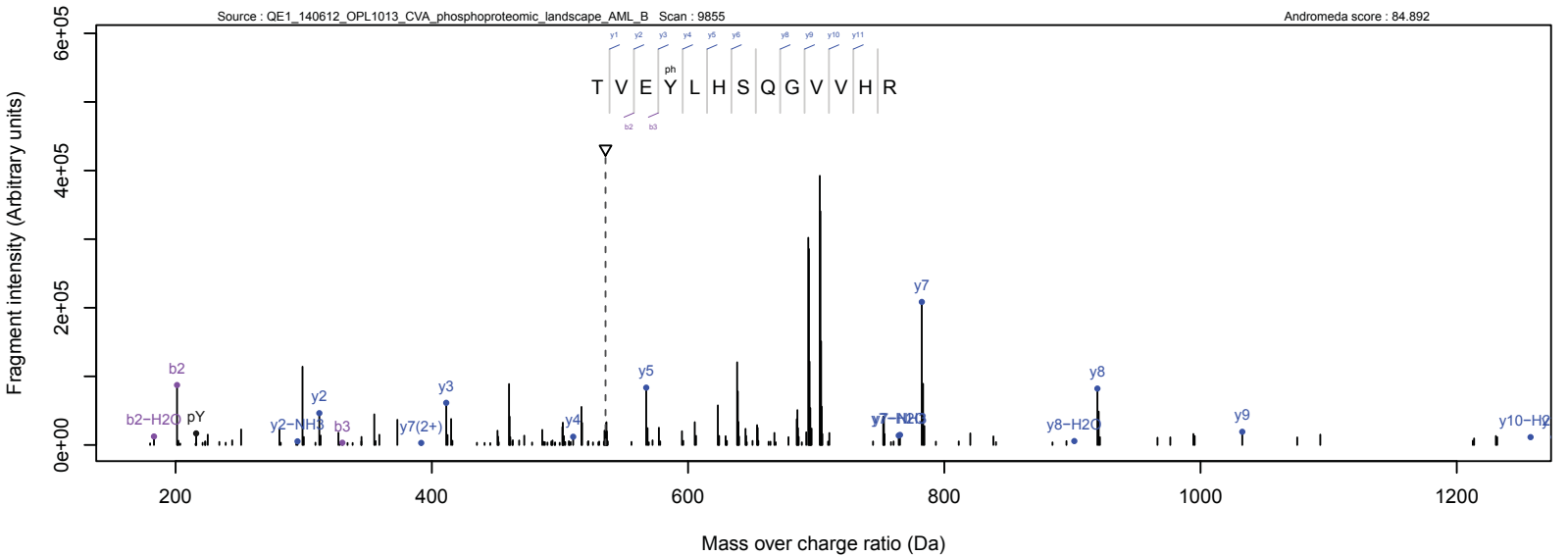
**AEEAAASAY(ph)YNPGNPHNVYMPPTSQPPPPYPPEDEKK-4+**



**NHVHQLHTY(ph)YQLGR-4+**



**TVEY(ph)LHSQGVVHR-3+**

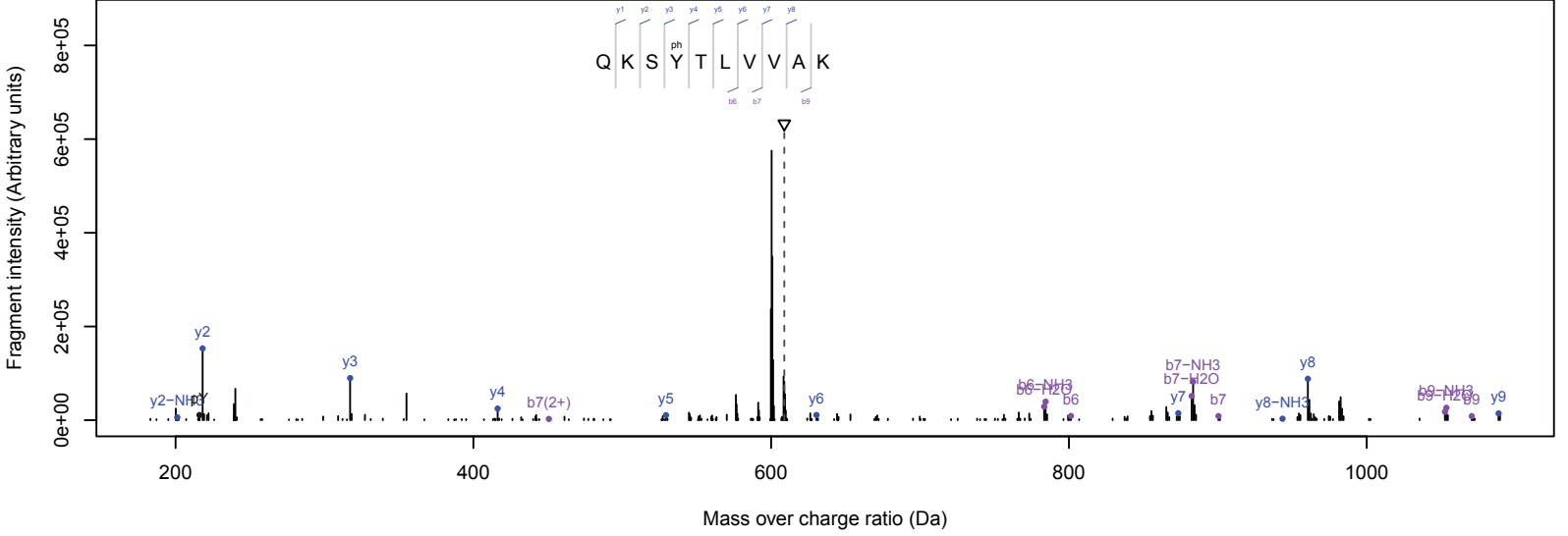




**QKSY(ph)TLVVAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10001

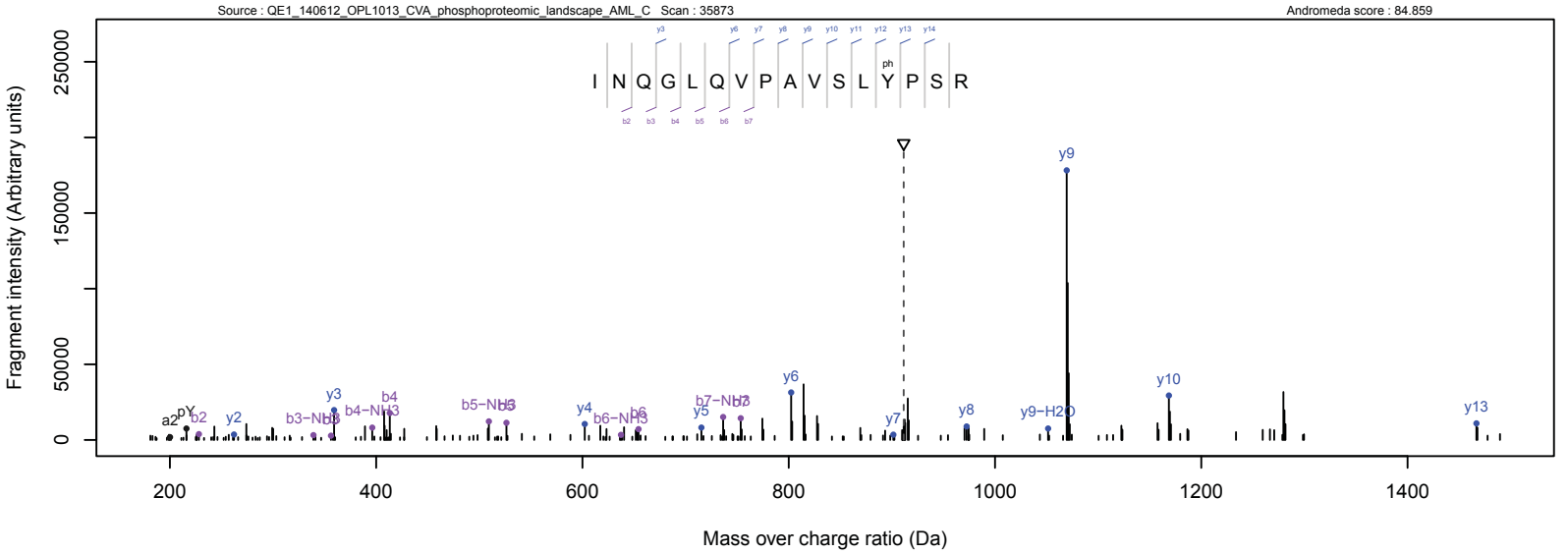
Andromeda score : 84.882



**INQLQVPAVSLY(ph)PSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35873

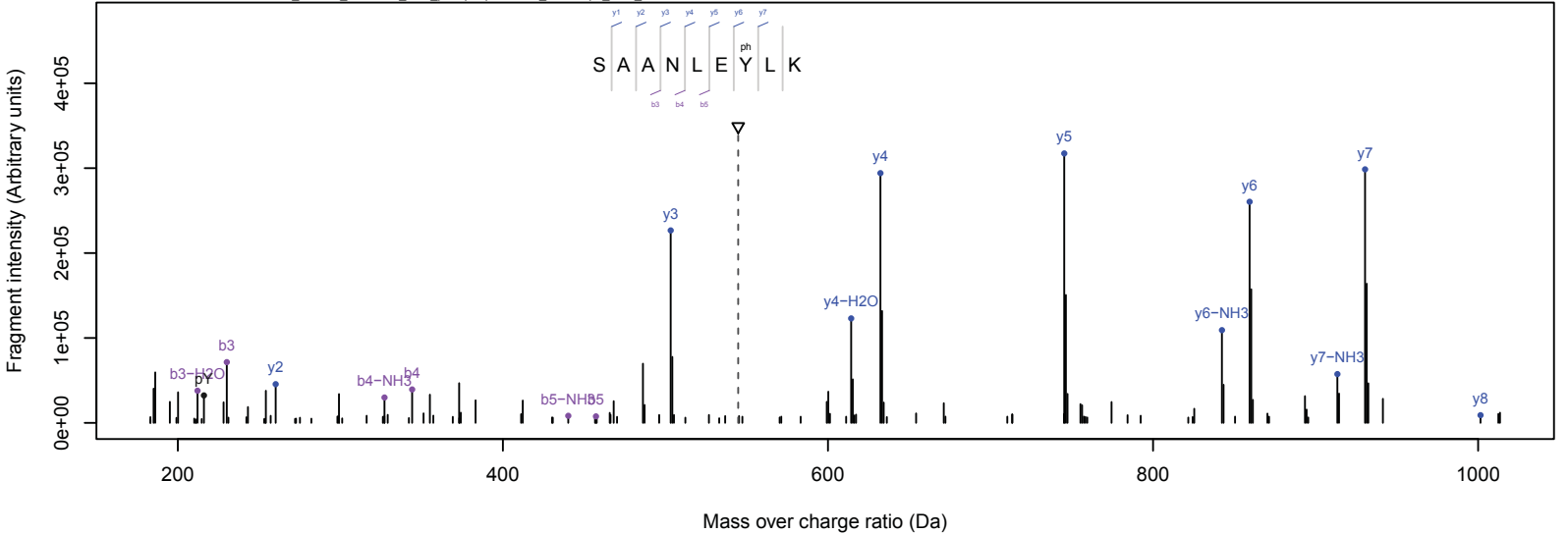
Andromeda score : 84.859



**SAANLEY(ph)LK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16673

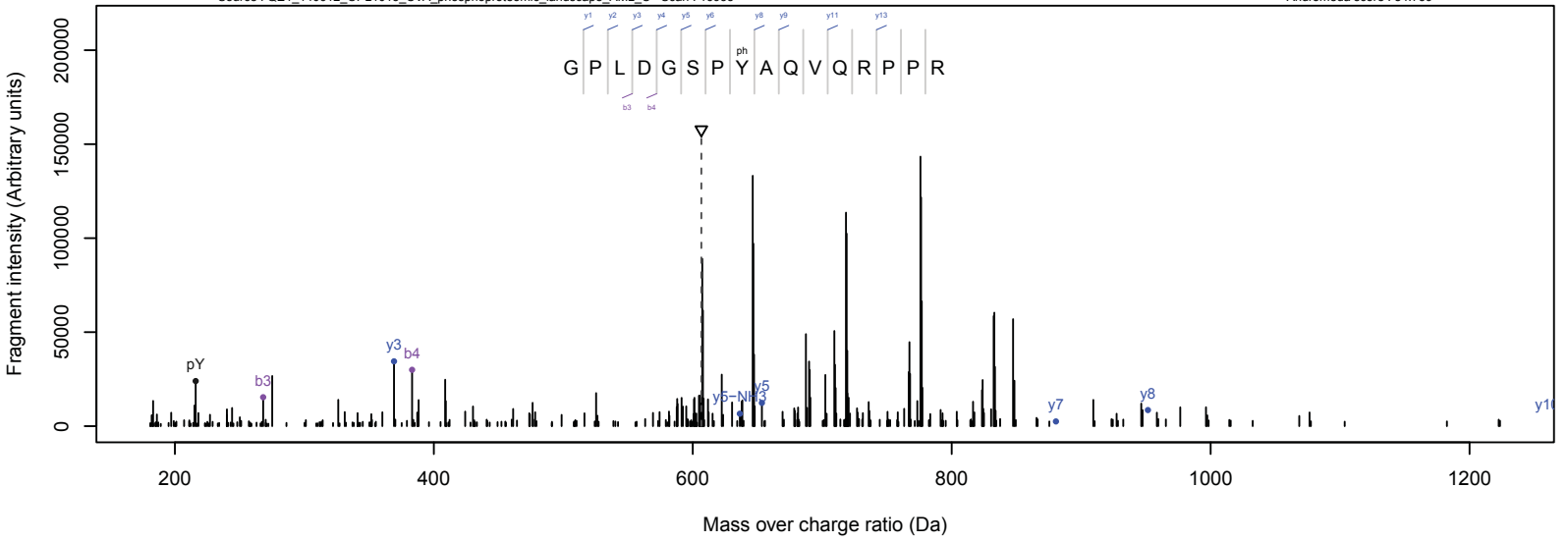
Andromeda score : 84.817



**GPLDGSFY(ph)AQVQRPPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16933

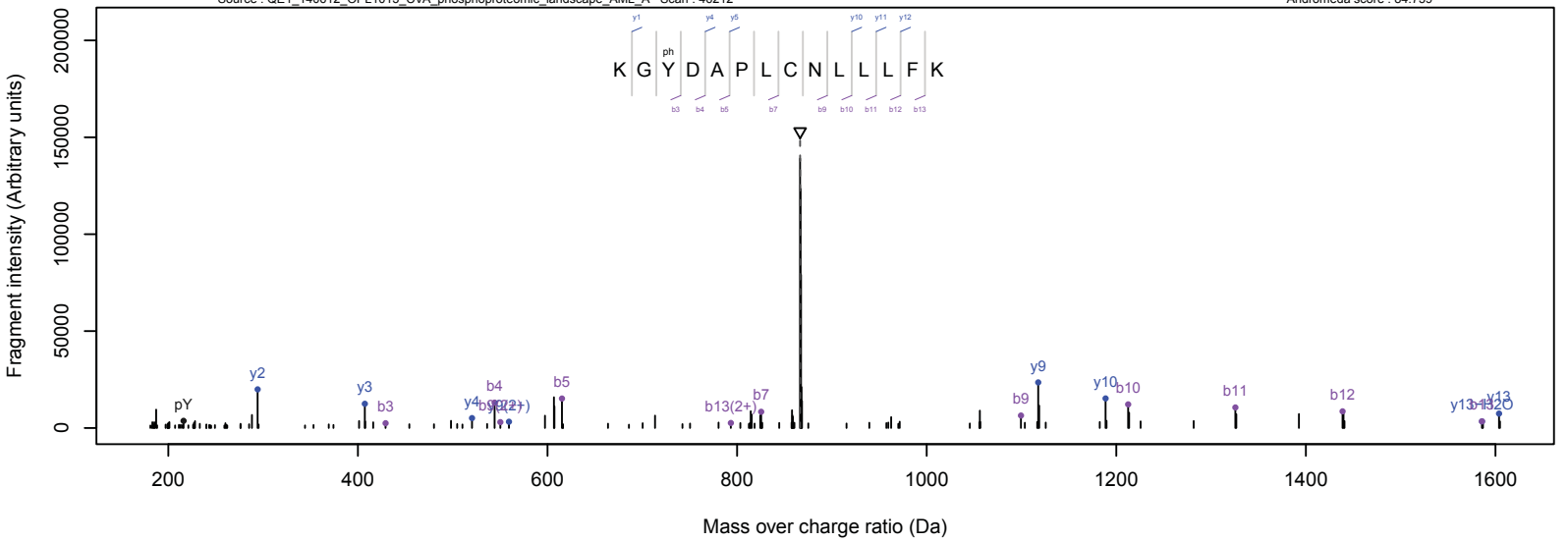
Andromeda score : 84.759



**KGY(ph)DAPLCNLLLFK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 46212

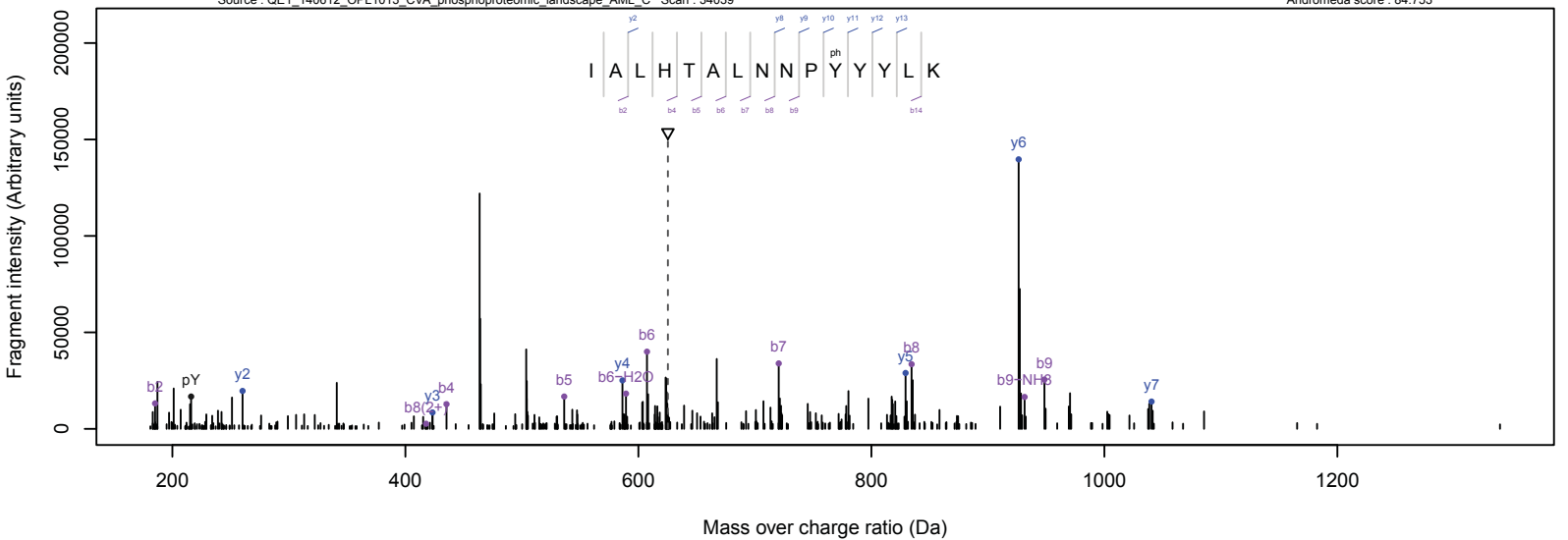
Andromeda score : 84.759



**IALHTALNNP(ph)YYLK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34039

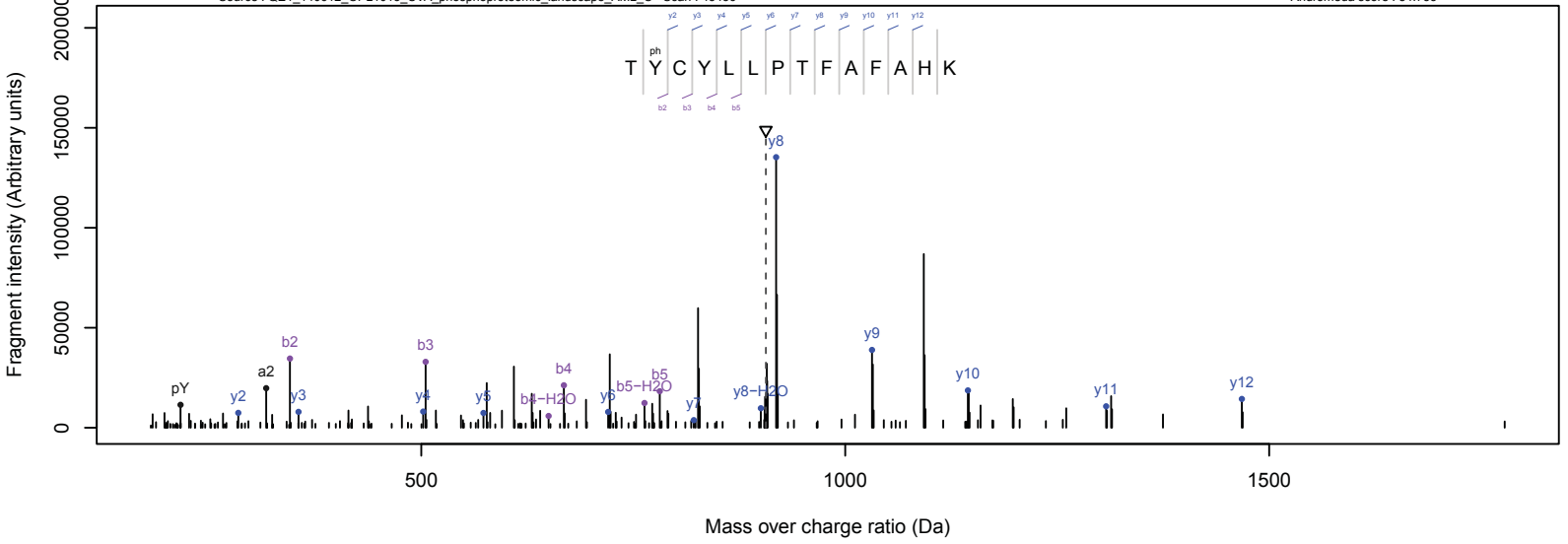
Andromeda score : 84.753



**TY(ph)CYLLPTFAFAHK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45436

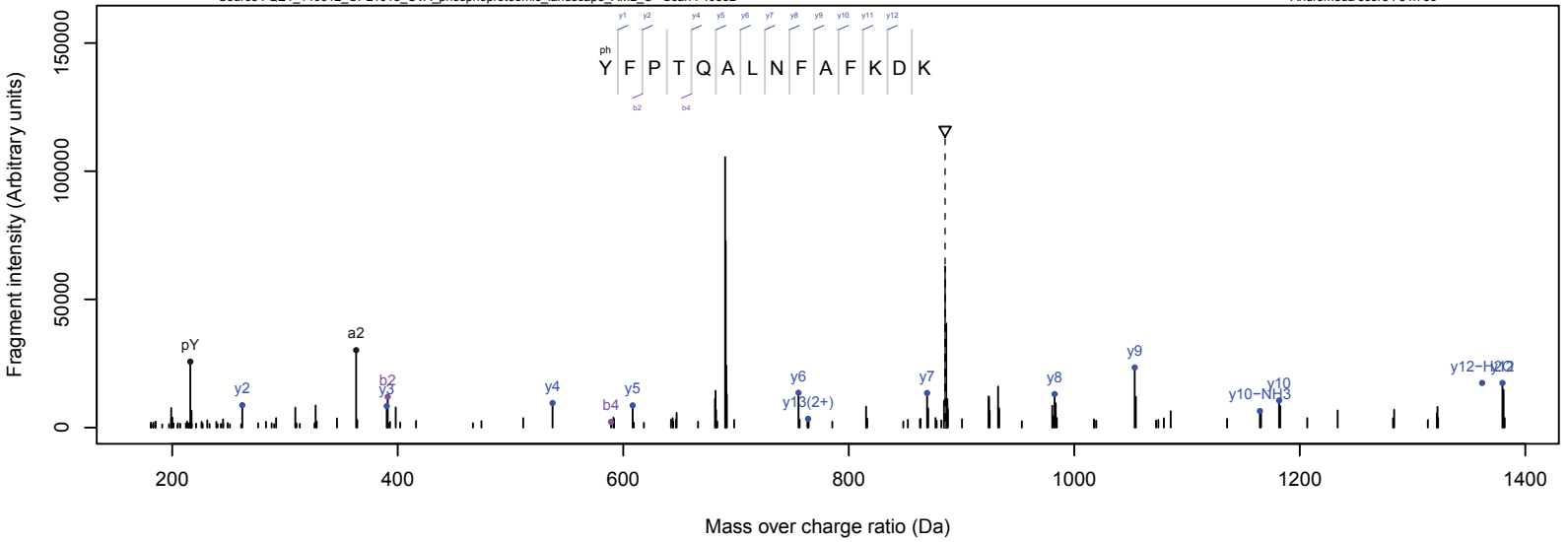
Andromeda score : 84.753



**Y(ph)FPTQALNFAFKDK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40852

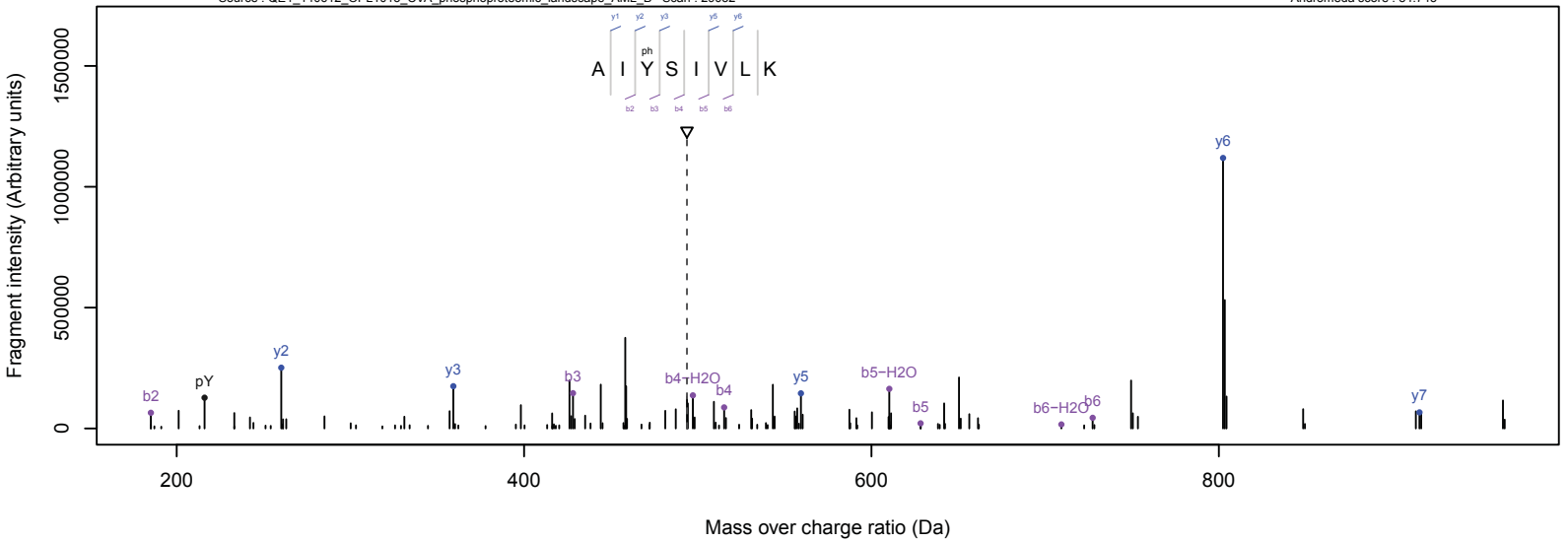
Andromeda score : 84.753



**AIY(ph)SIVLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29032

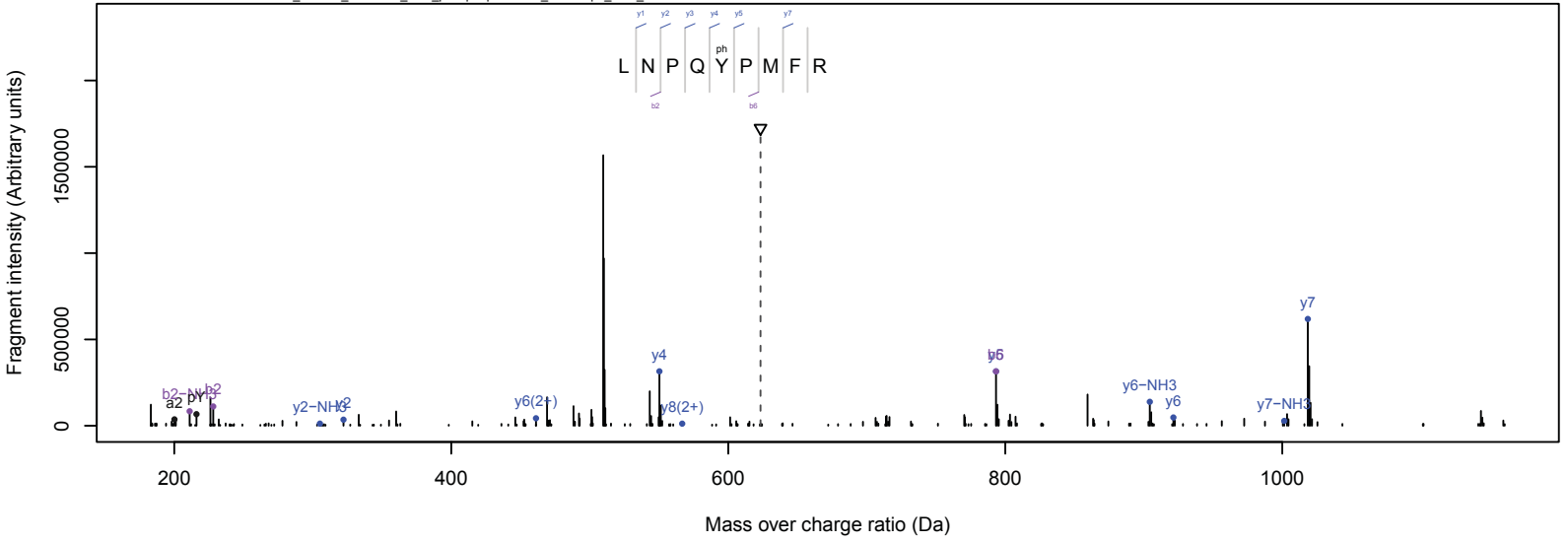
Andromeda score : 84.743



LNPQY(ph)PMFR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 24686

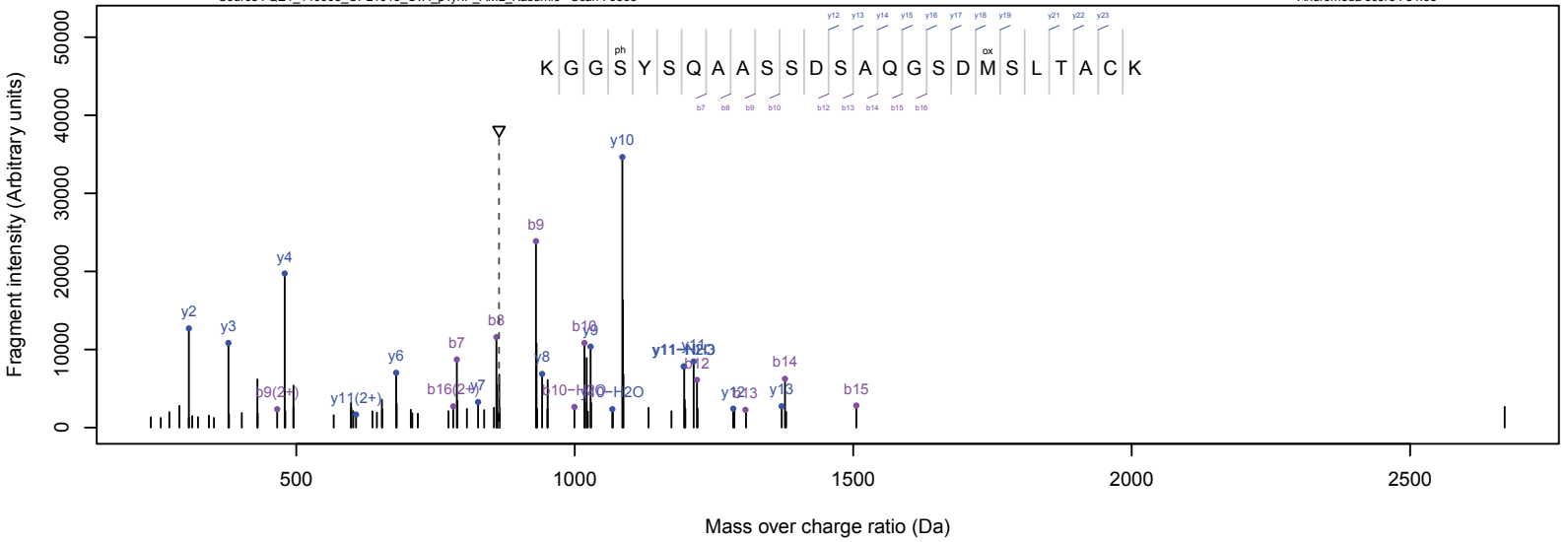
Andromeda score: 84.743



KGGS(ph)YSQAASSDSAQGS(DM)SLTACK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 9368

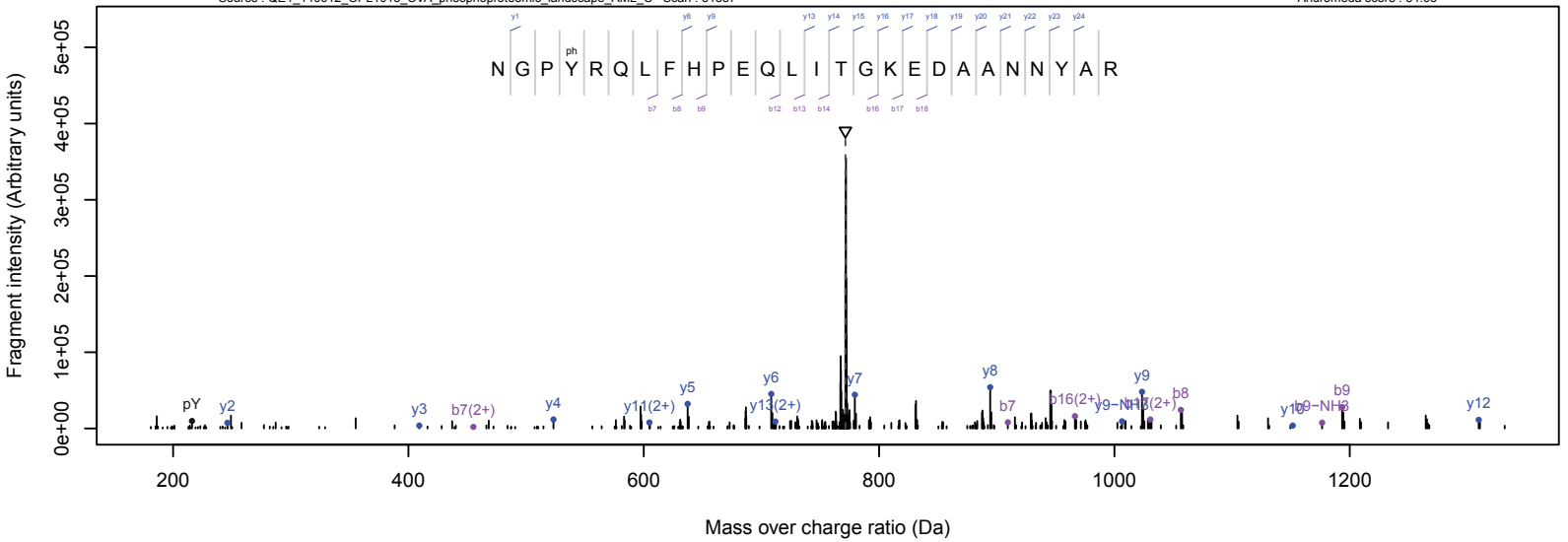
Andromeda score: 84.68



NGPY(ph)RQLFHPEQLITGKEDAANNYAR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31587

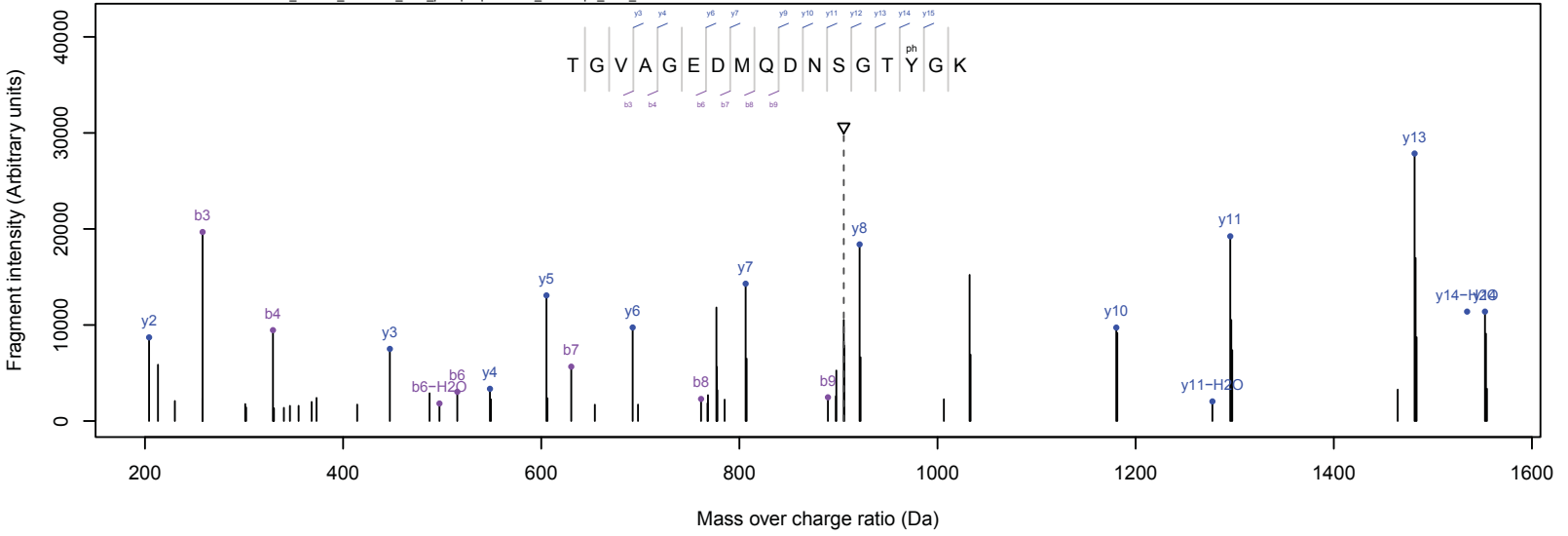
Andromeda score: 84.68



TGVAGEDM<sup>ph</sup>QD<sup>ph</sup>NSGT<sup>ph</sup>Y<sup>ph</sup>GK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7590

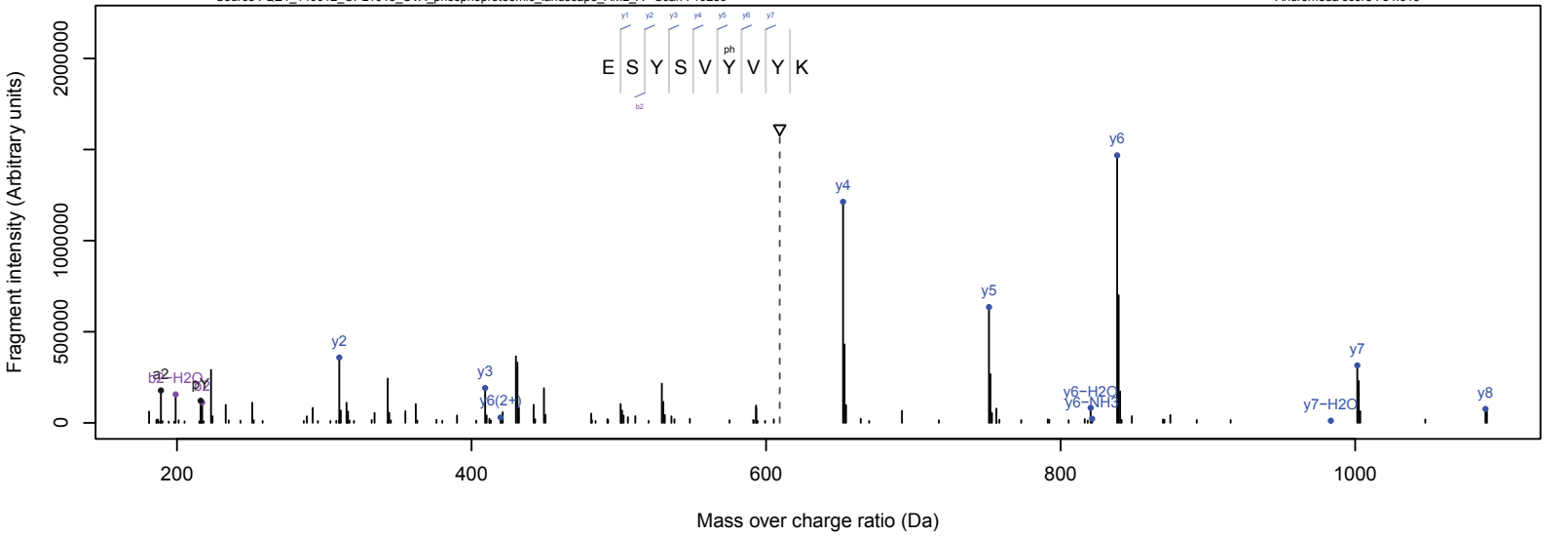
Andromeda score : 84.653



ESYSV<sup>ph</sup>Y<sup>ph</sup>VYK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 19289

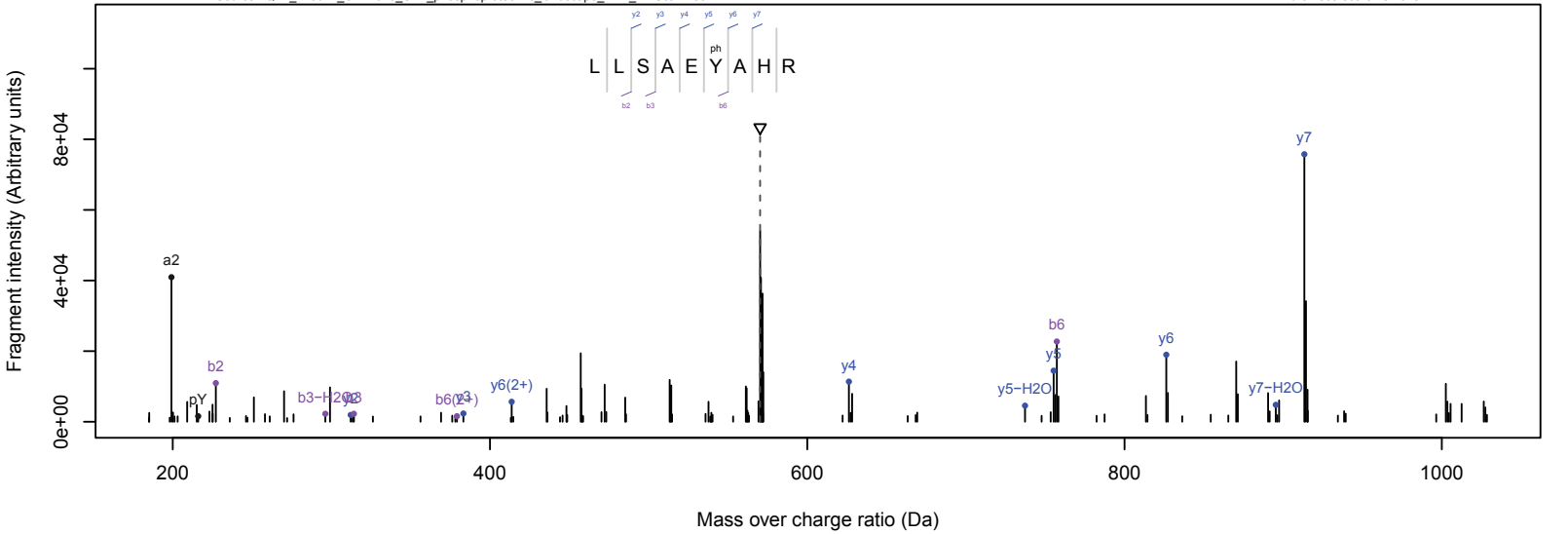
Andromeda score : 84.615



LLSAE<sup>ph</sup>Y<sup>ph</sup>AHR

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 8521

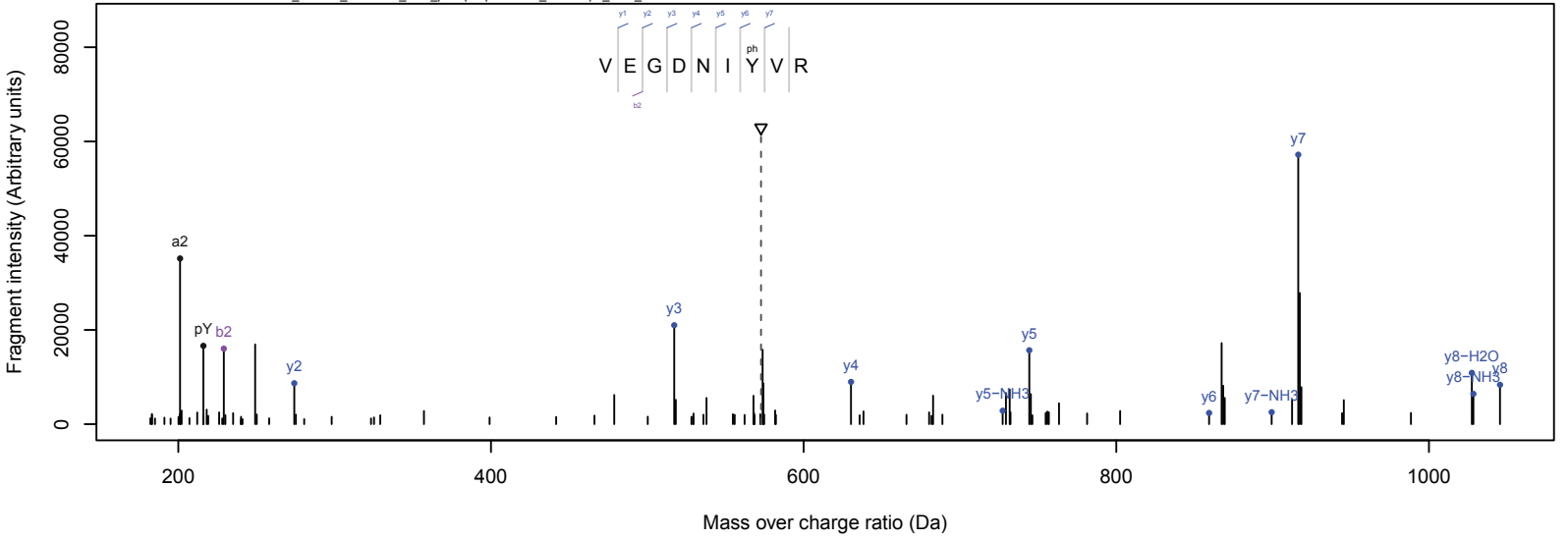
Andromeda score : 84.615



VEGDNIY(ph)VR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11956

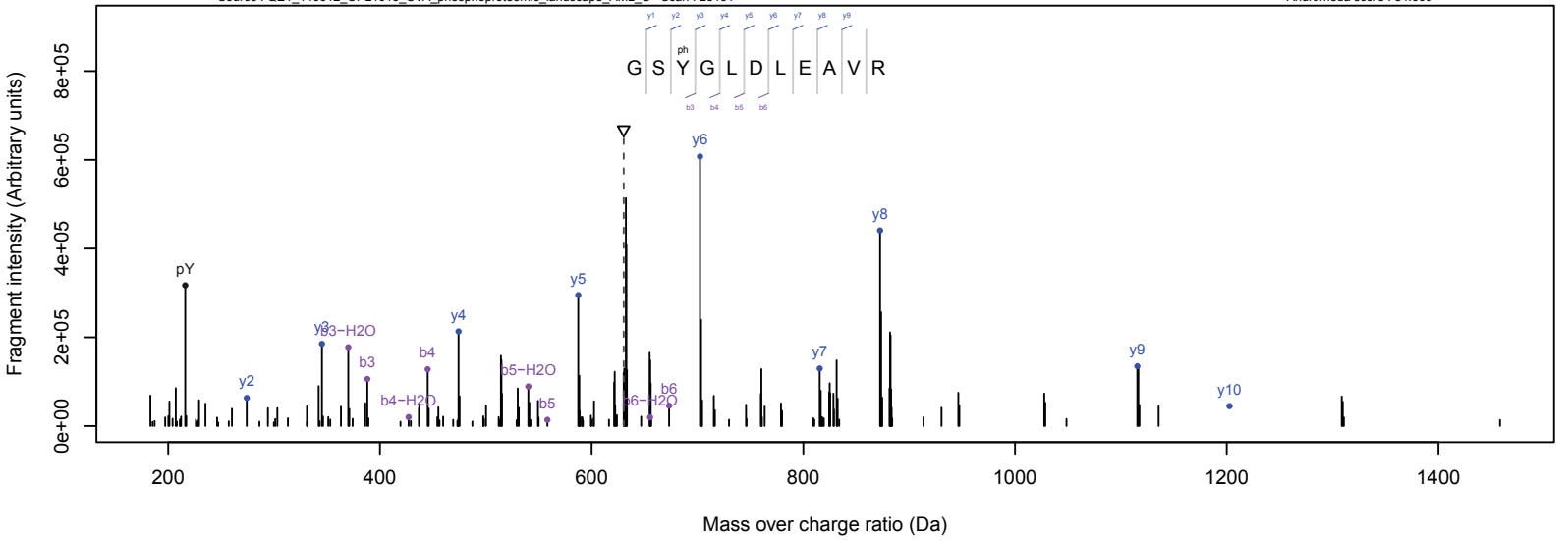
Andromeda score : 84.615



GSY(ph)GLDLEAVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28151

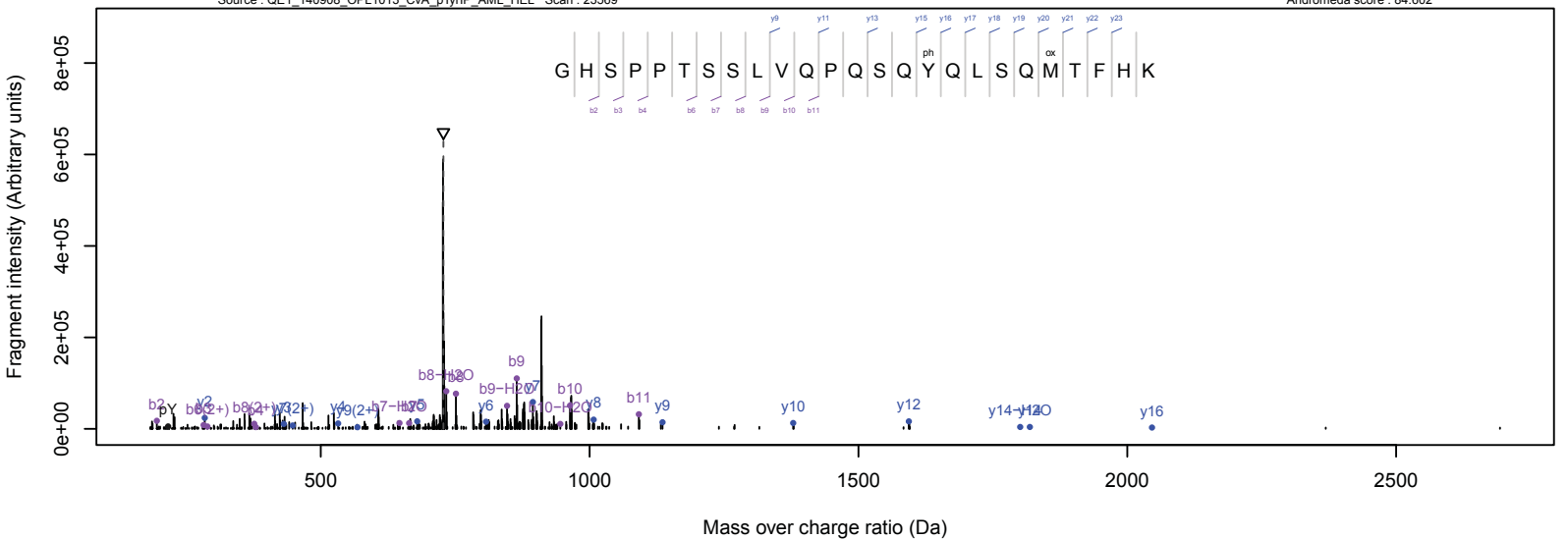
Andromeda score : 84.605



GHSPTSSLVQPSQY(ph)QLSQM(ox)TFHK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23569

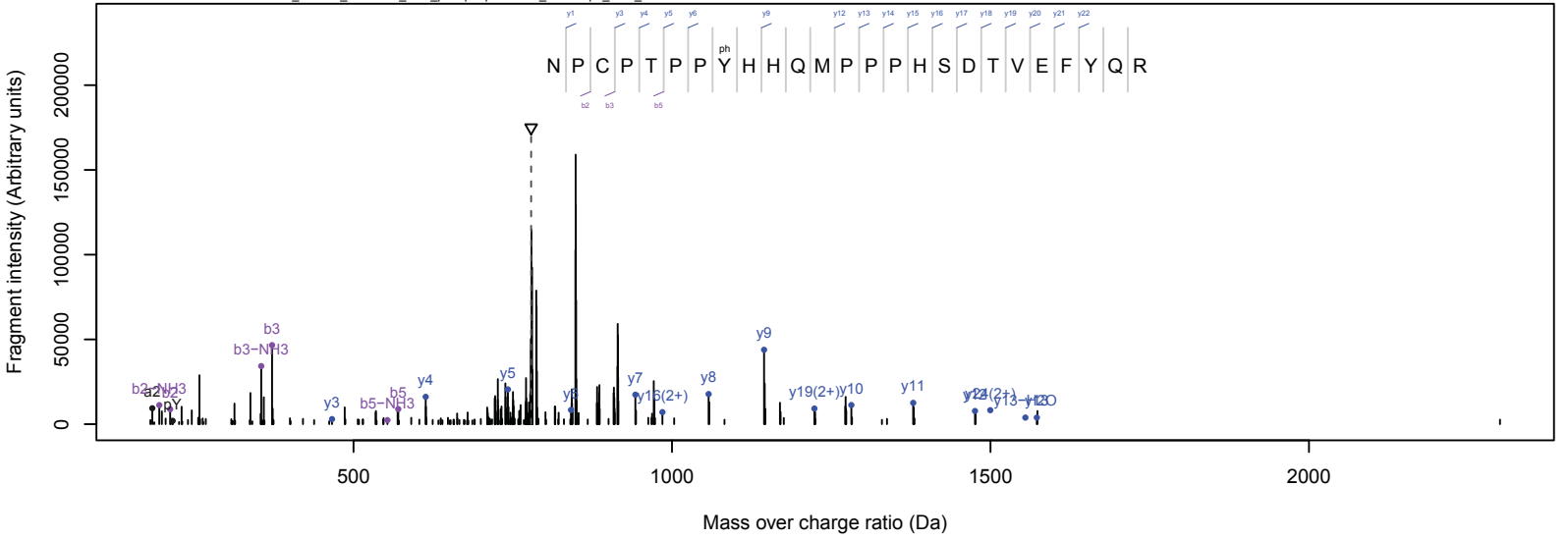
Andromeda score : 84.602



**NPCPTPY(ph)HHQMPPPHSDTVEFYQR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19878

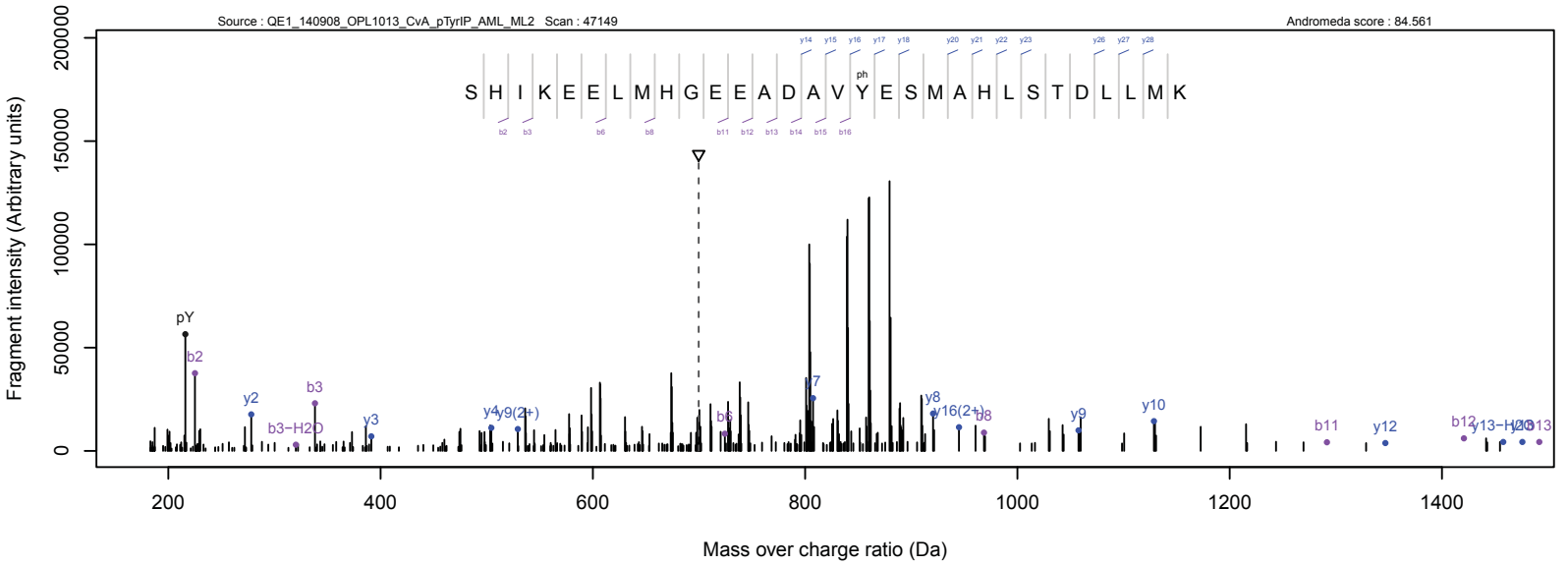
Andromeda score : 84.602



**SHIKEELMHGEEADAVY(ph)ESMAHLSTDLMLK-5+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 47149

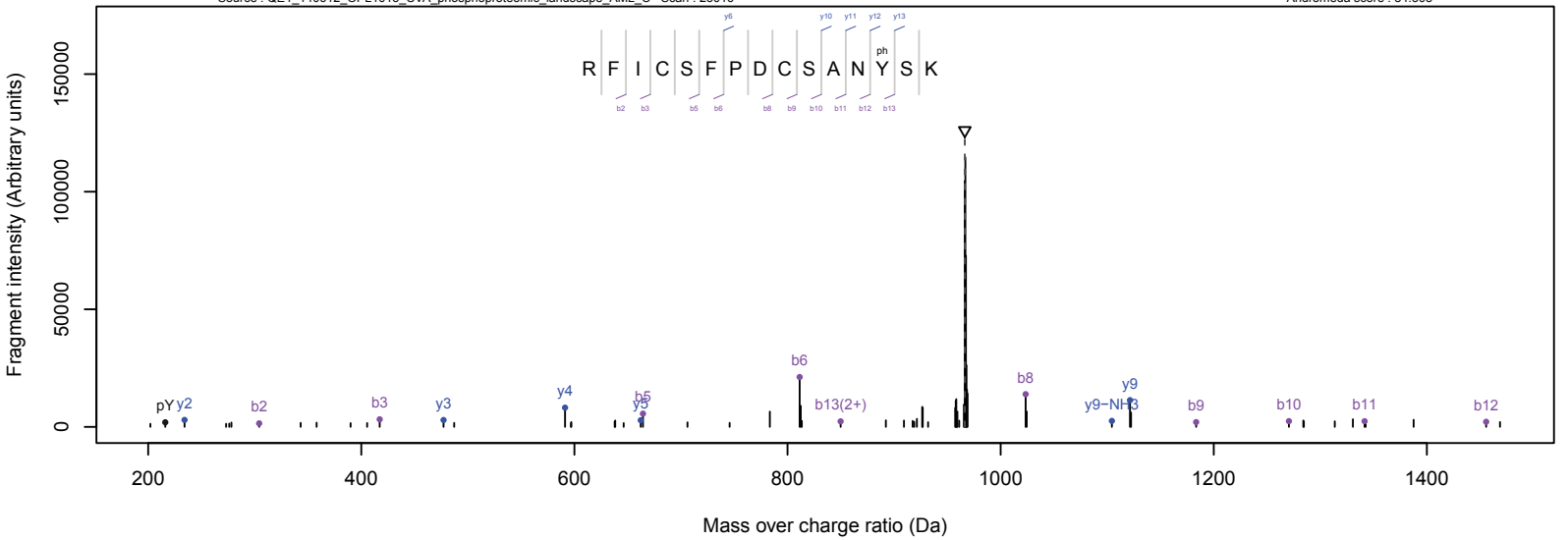
Andromeda score : 84.561



**RFICSPDCSANY(ph)SK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23016

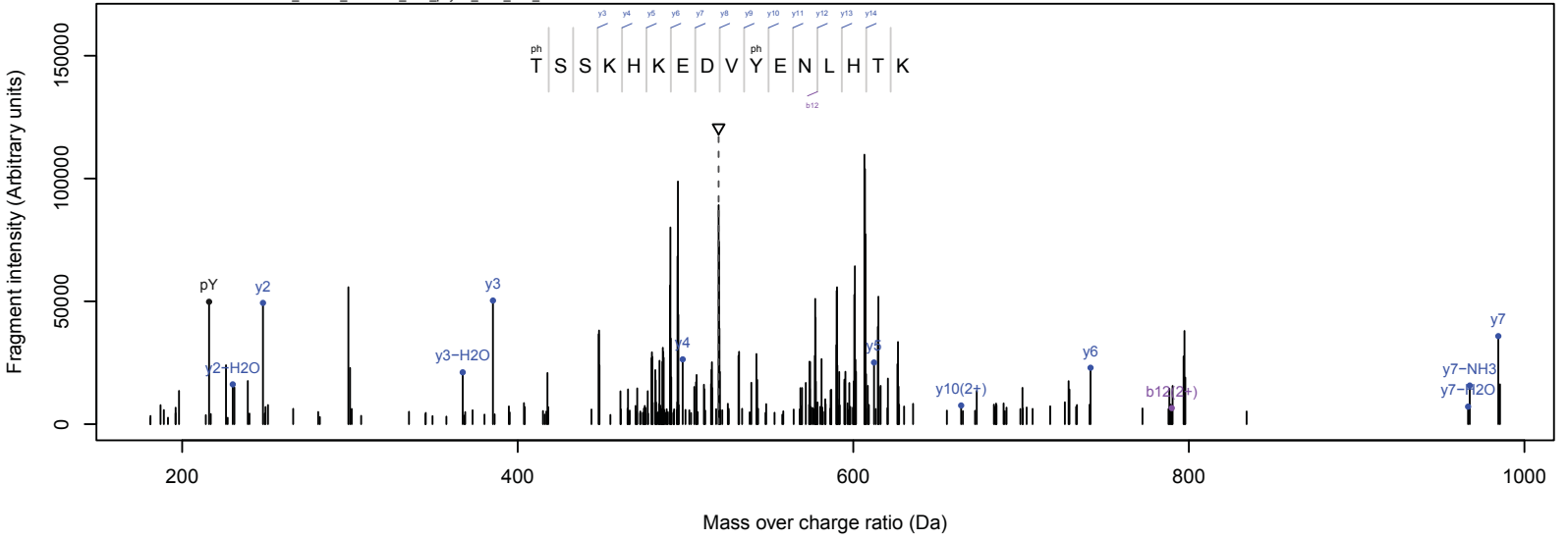
Andromeda score : 84.508



T(ph)SSKHKEDVY(ph)ENLHTK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 5578

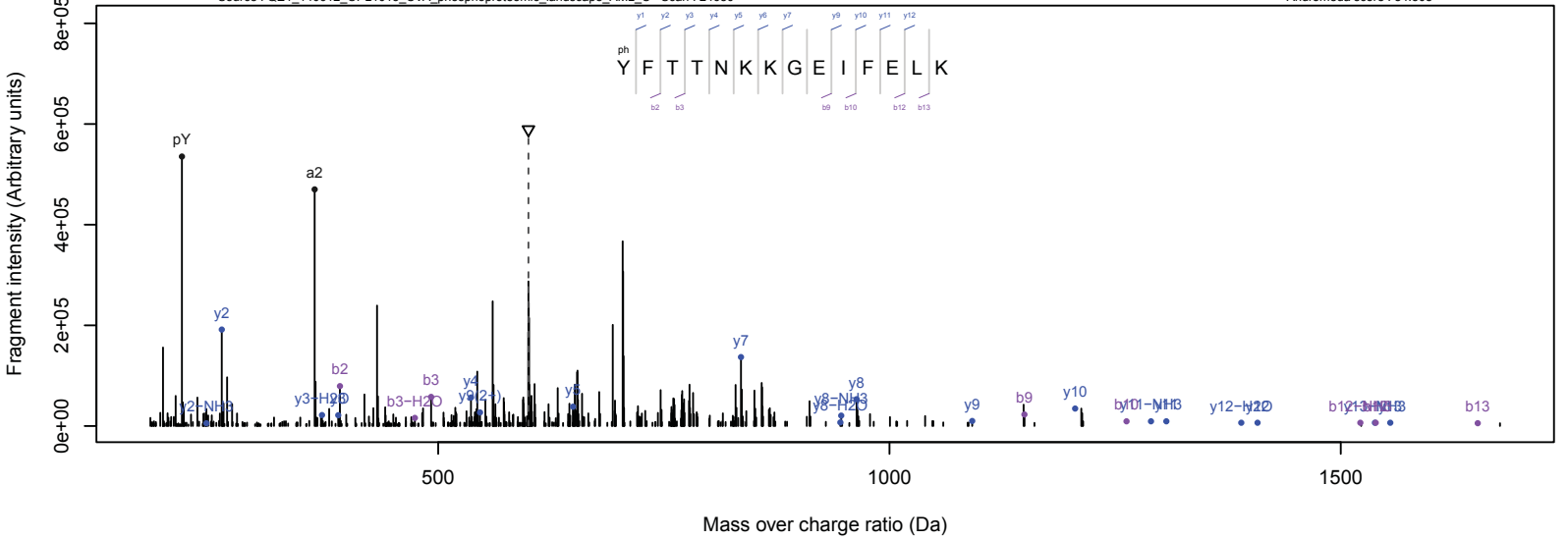
Andromeda score : 84.508



Y(ph)FTTNKGEIFELK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 24680

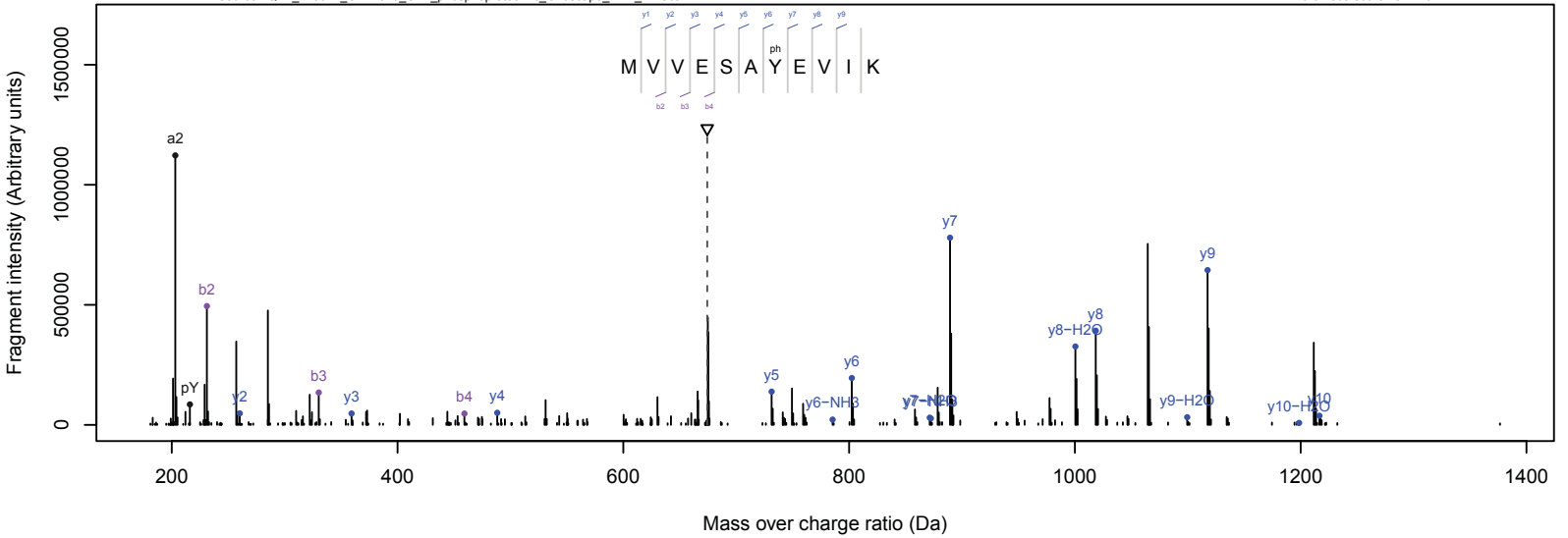
Andromeda score : 84.508



MVVESAY(ph)EVIK-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan : 27442

Andromeda score : 84.479

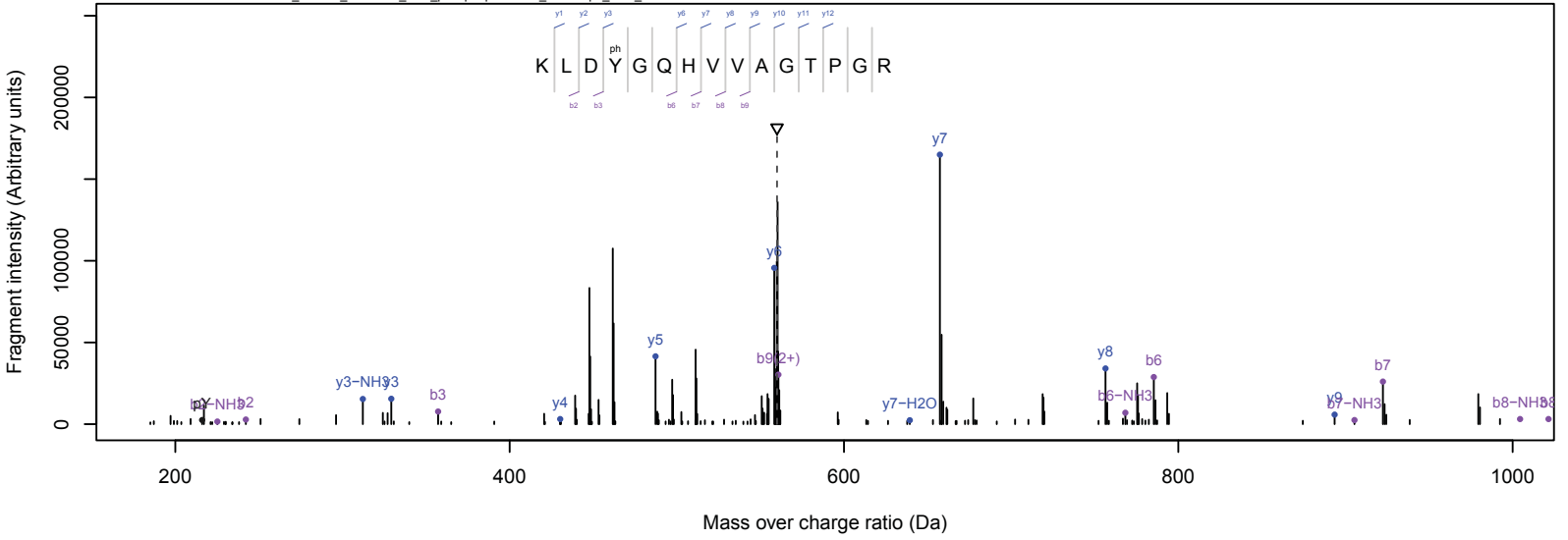




**KLDY(ph)GQHVVAGTPGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 10698

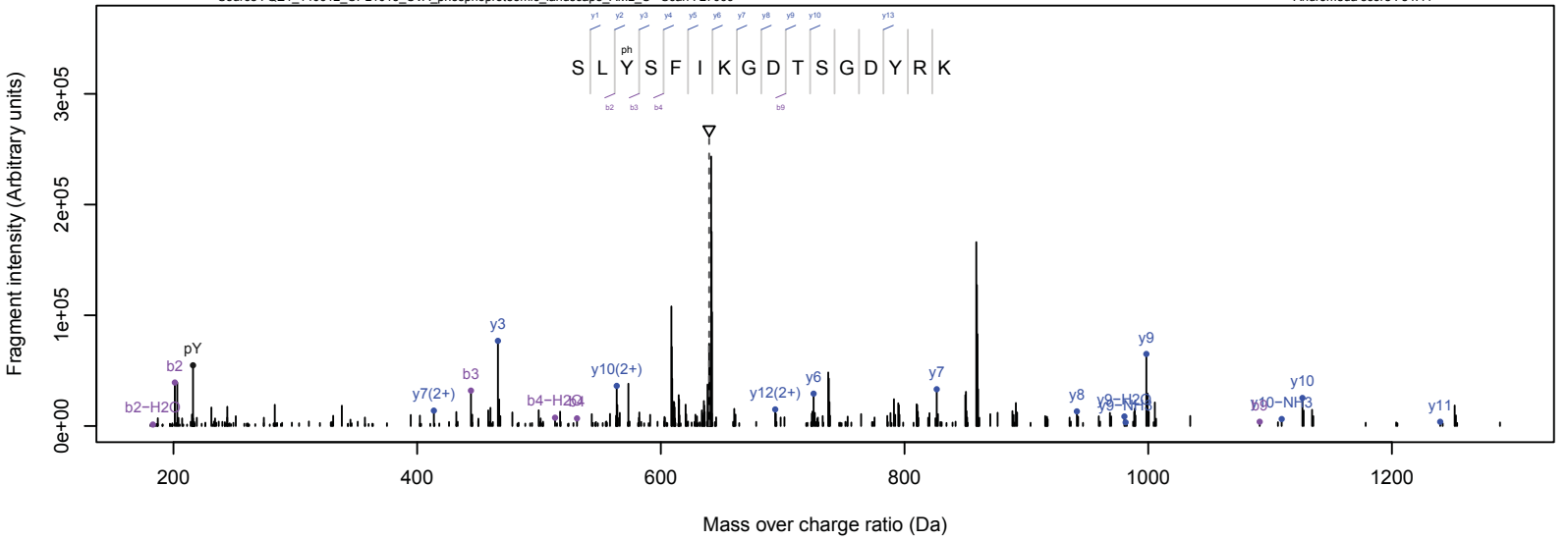
Andromeda score : 84.41



**SLY(ph)SFIKGDTSGLDYRK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27069

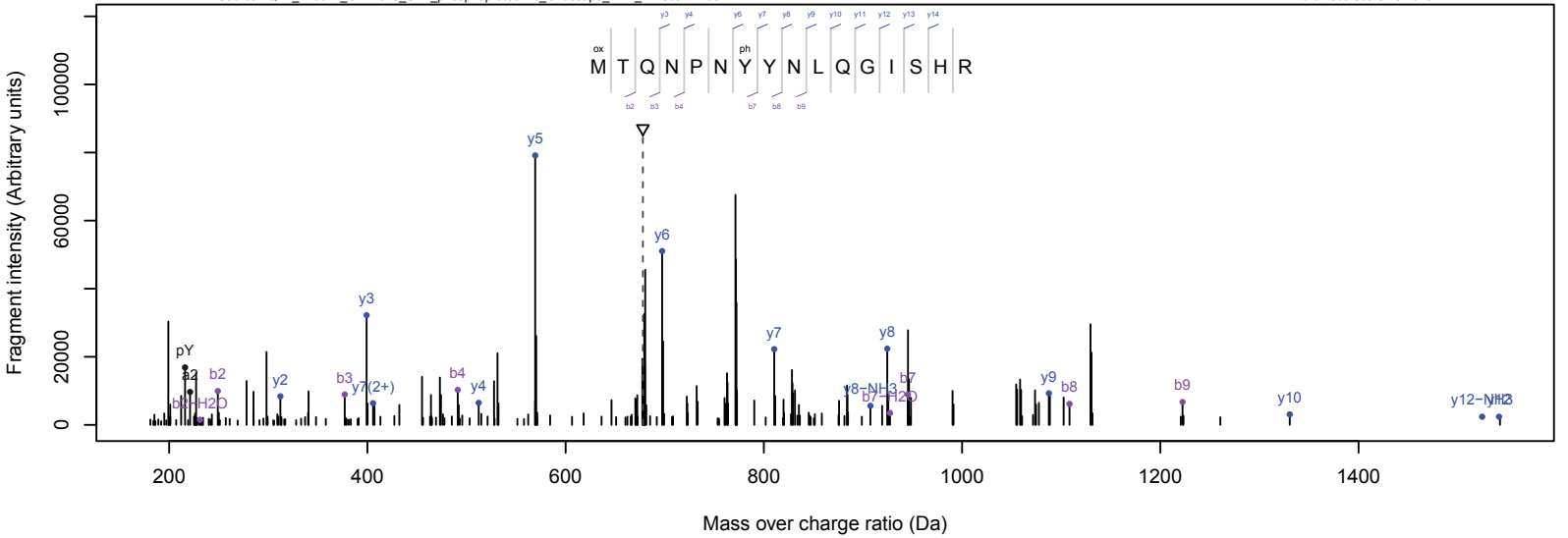
Andromeda score : 84.41



**M(ox)TQNPY(ph)YNLQGISHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 19341

Andromeda score : 84.375

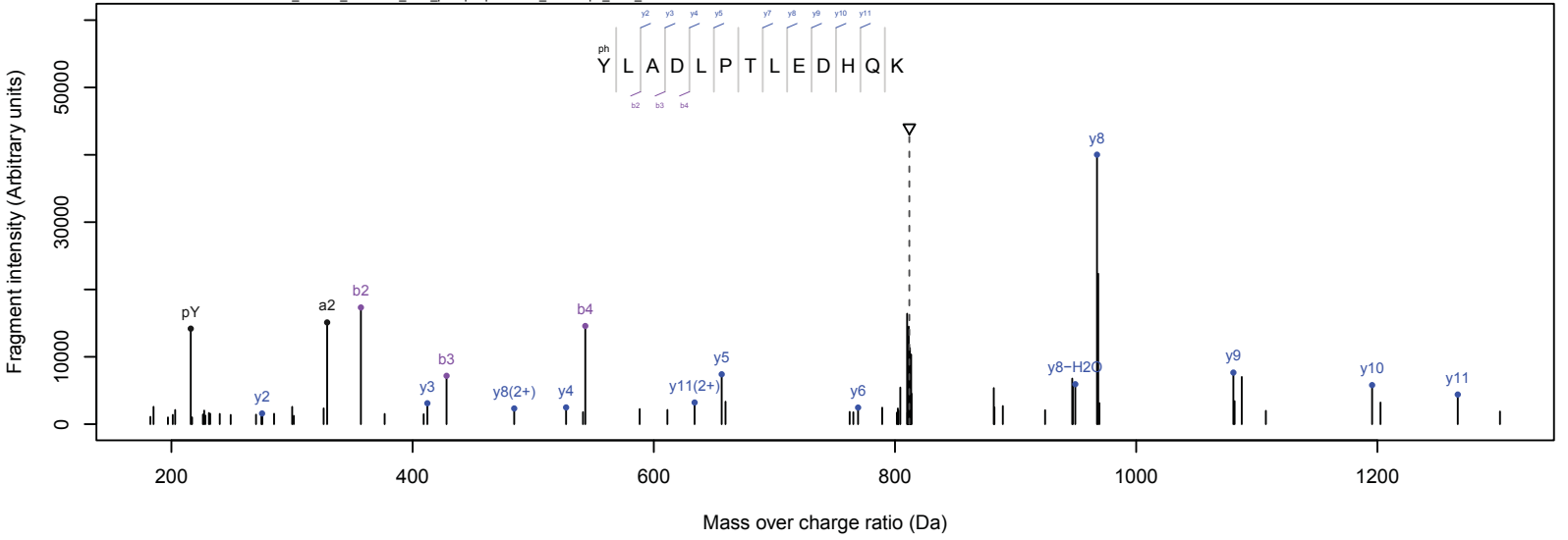




Y(ph)LADLPTLEDHQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28216

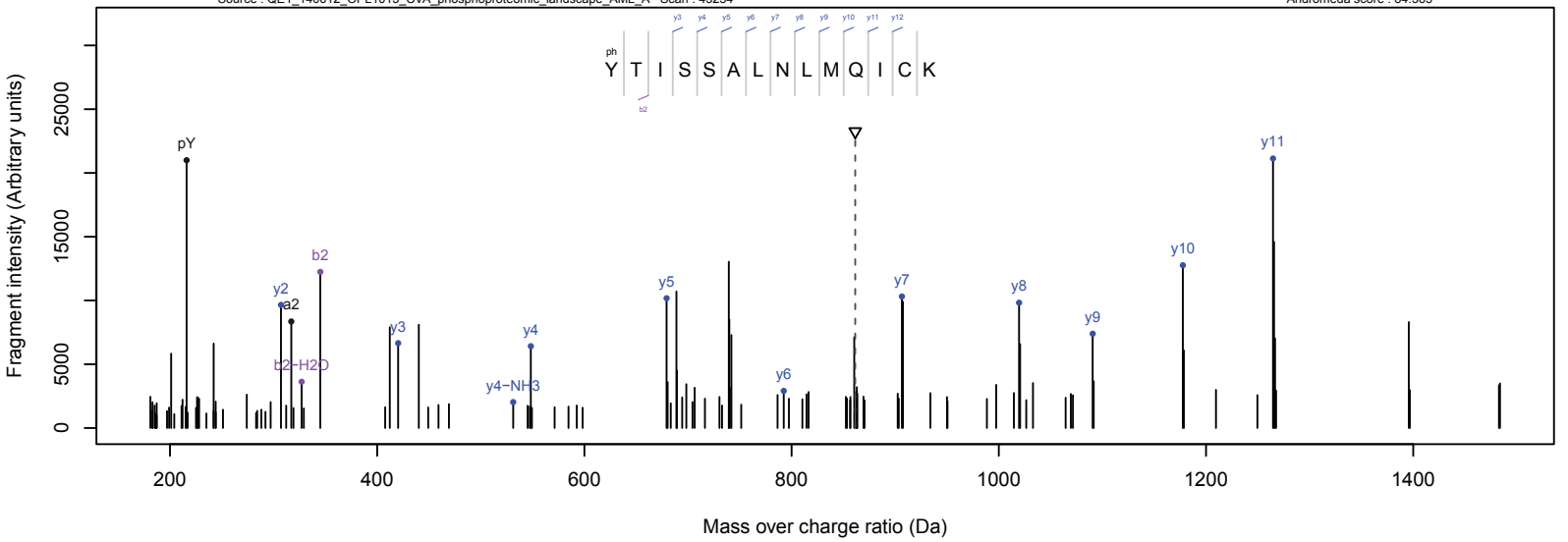
Andromeda score : 84.365



Y(ph)TISSALNLMQICK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 43254

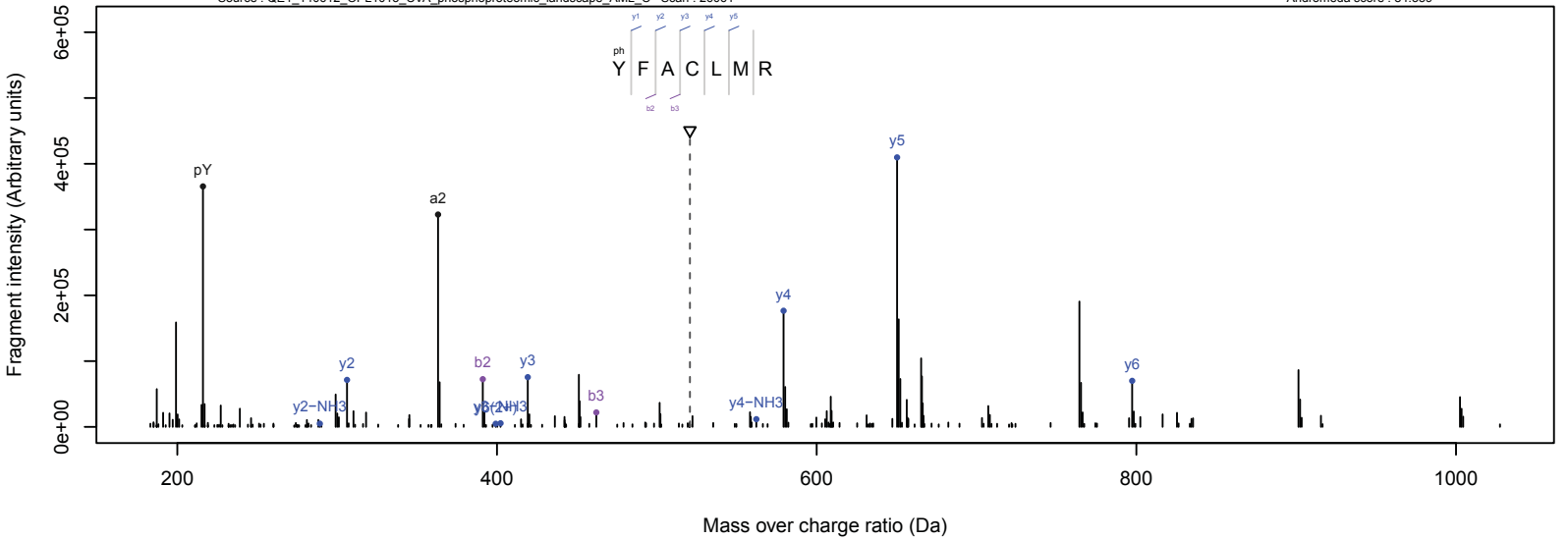
Andromeda score : 84.365



Y(ph)FACLMR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26061

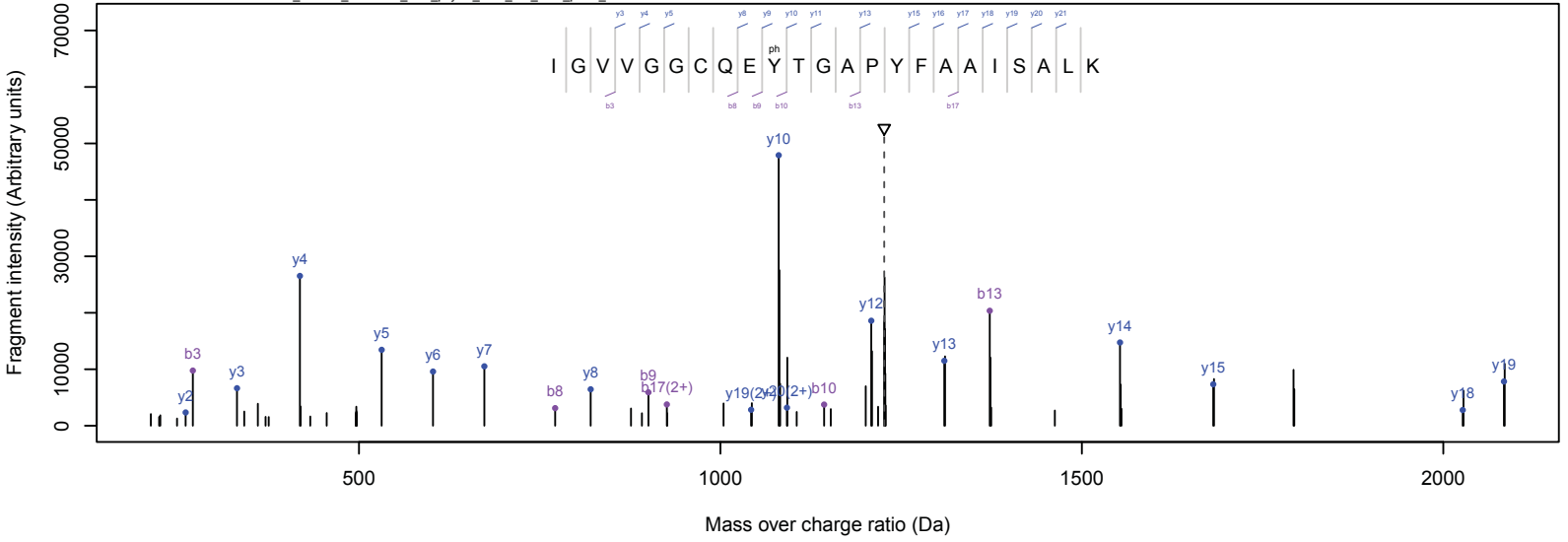
Andromeda score : 84.359



IGVVGGCQEY(ph)TGAPYFAAISALK-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 42947

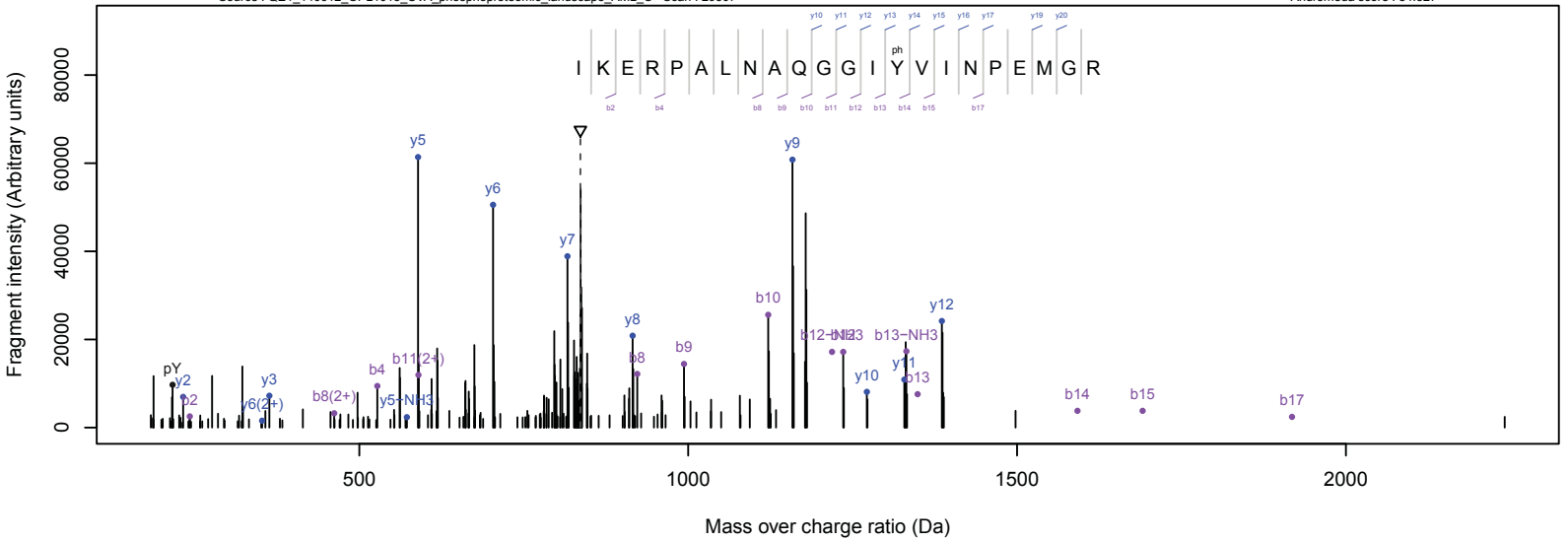
Andromeda score : 84.327



IKERPALNAQGGIY(ph)VINPEMGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29507

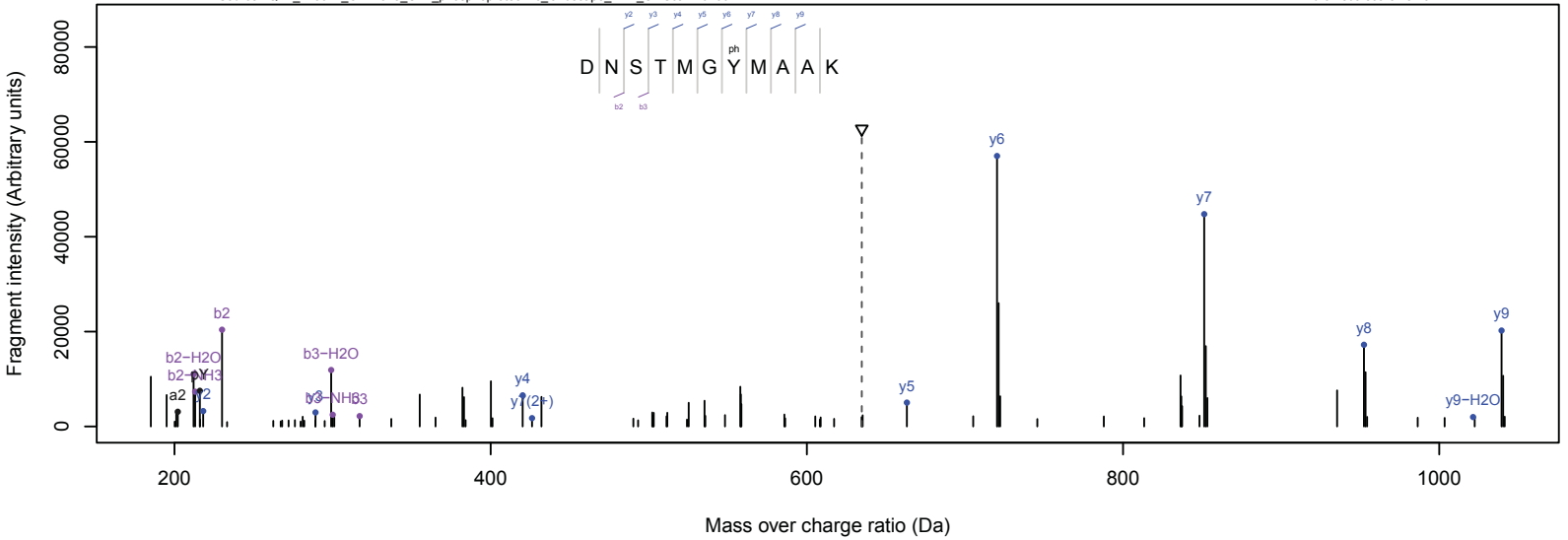
Andromeda score : 84.327



DNSTMGY(ph)MAAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9400

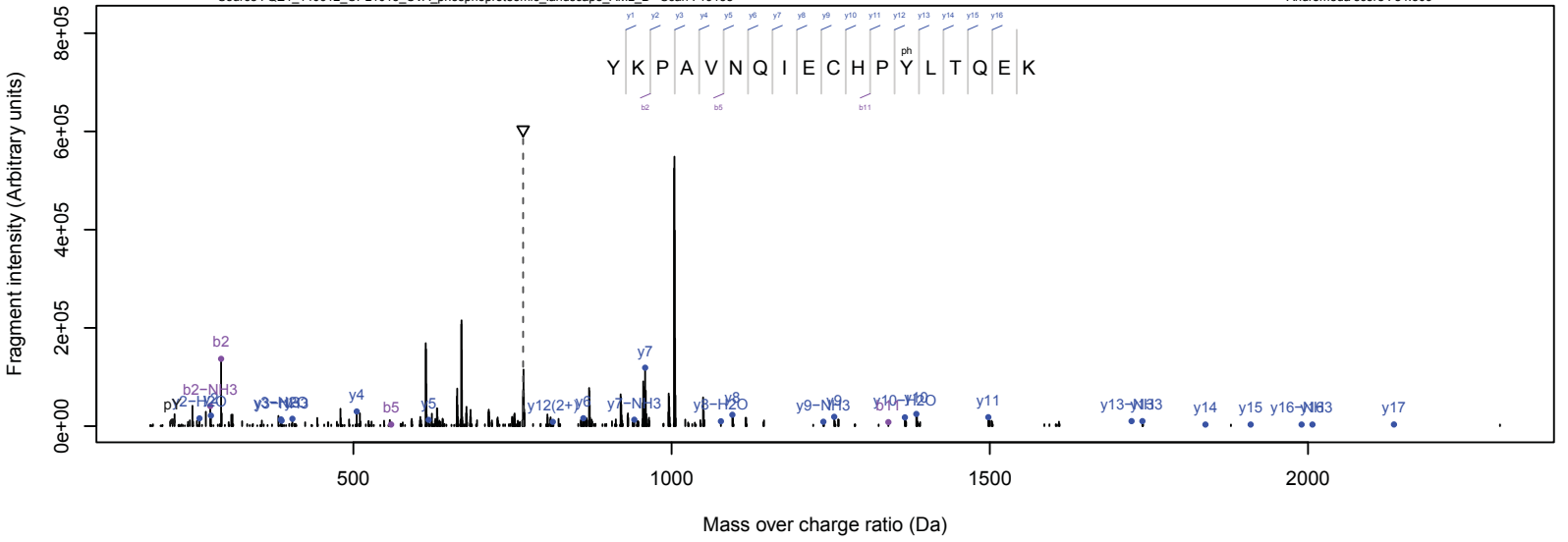
Andromeda score : 84.31



YKPAVNQIECHPY(ph)LTQEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19158

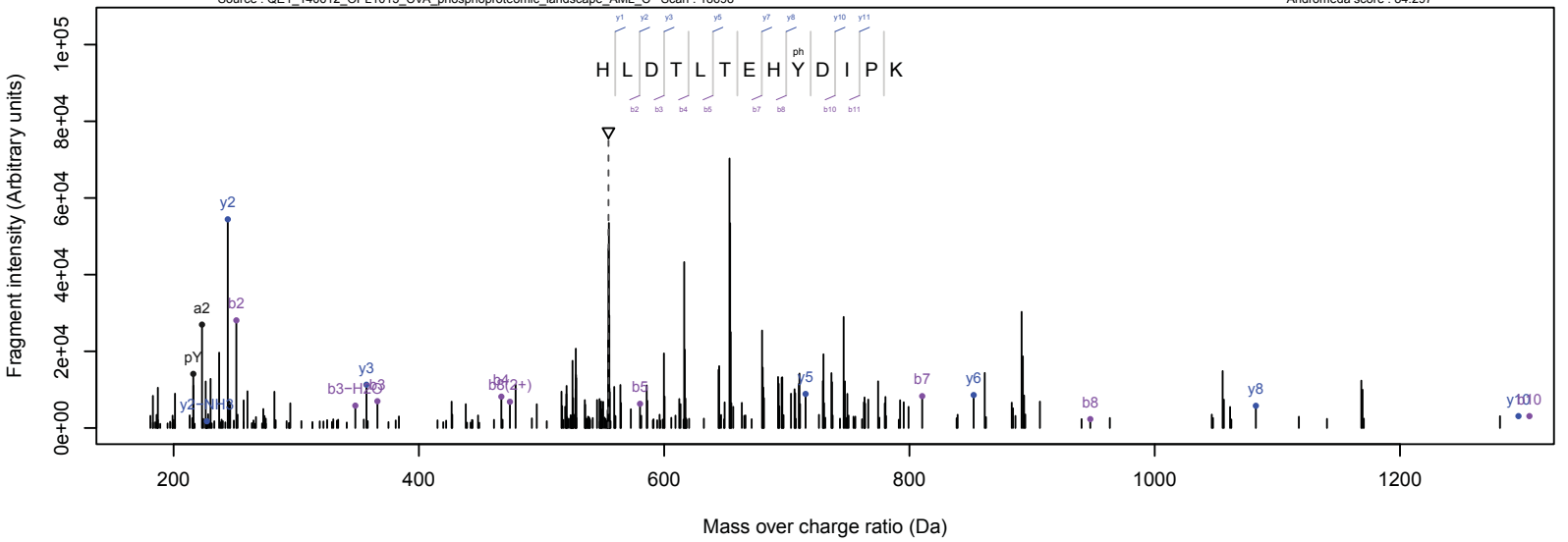
Andromeda score : 84.309



HLDLTHEY(ph)DIPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18698

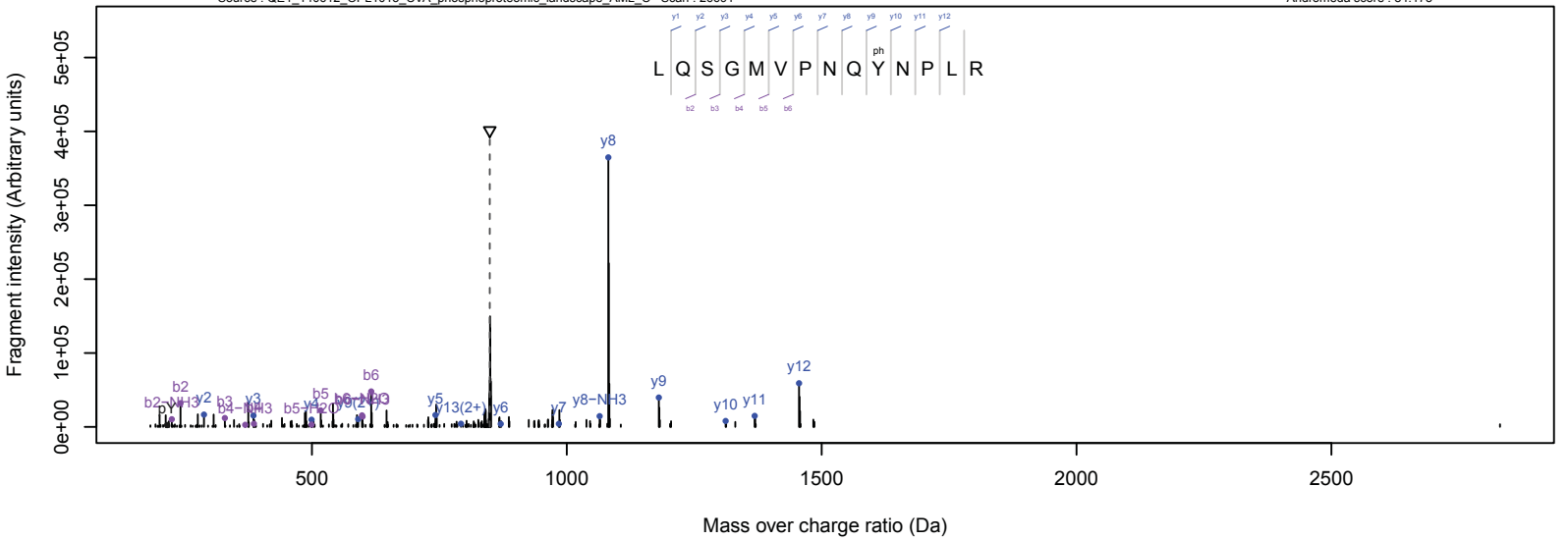
Andromeda score : 84.297



LQSGMVPNQY(ph)NPLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26691

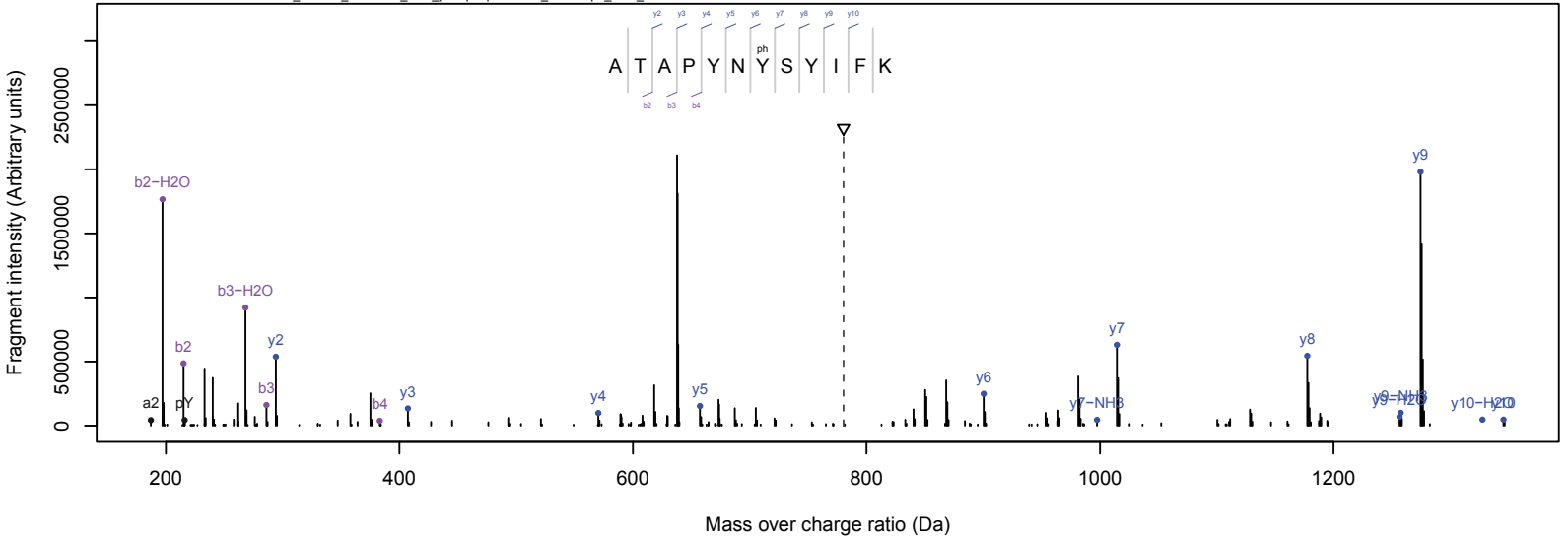
Andromeda score : 84.173



(ac)ATAPYNY(ph)SYIFK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41586

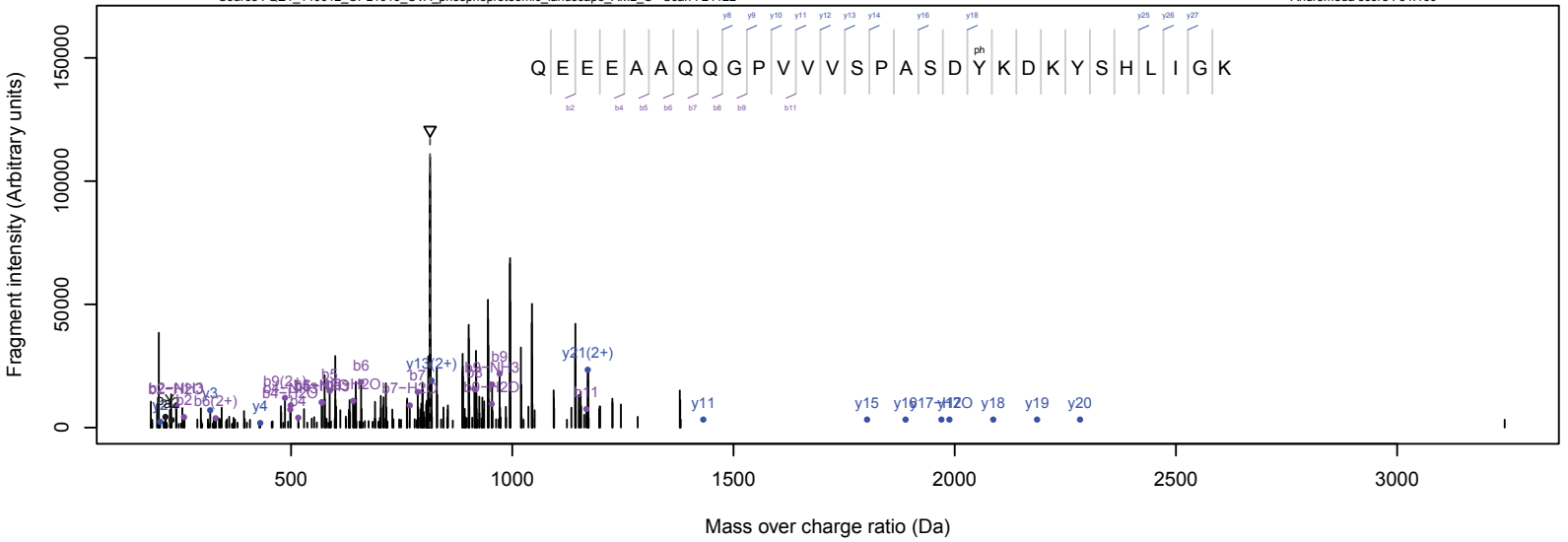
Andromeda score: 84.169



QEEEEAAQQGPVVVSPASDY(ph)KDKYSHLIGK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24122

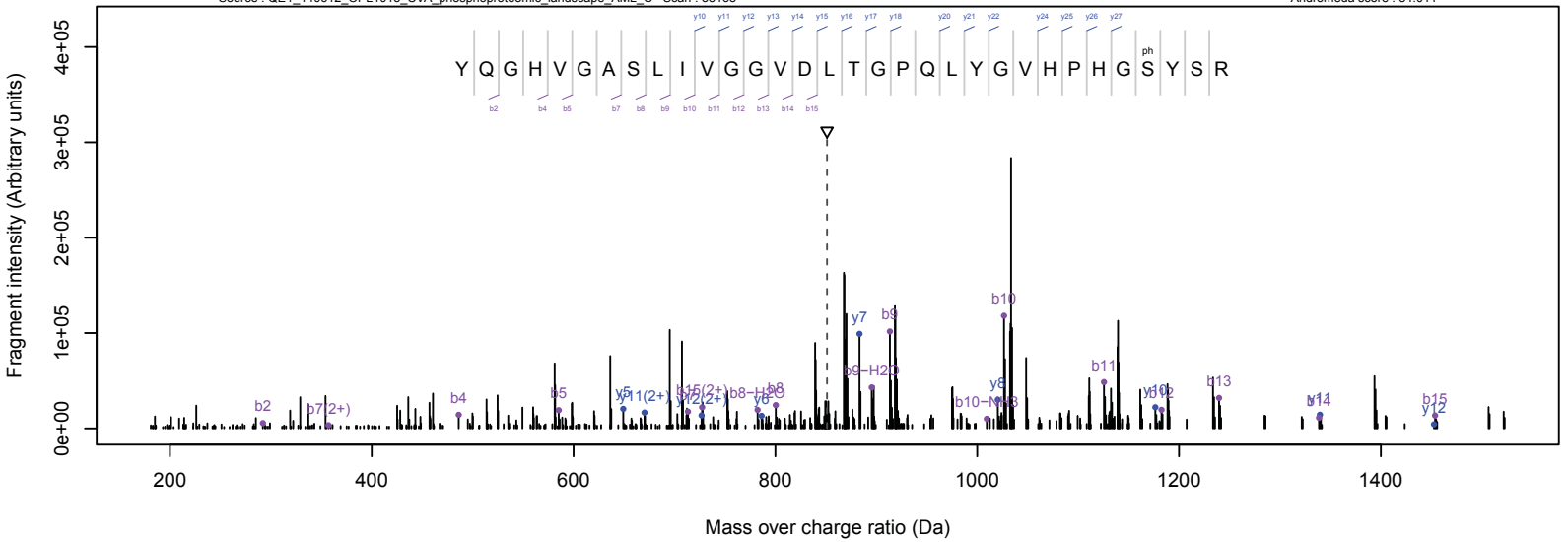
Andromeda score: 84.169



YQGHVGASLIVGGVDLTGPQLYGVHPHGS(ph)YSR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38168

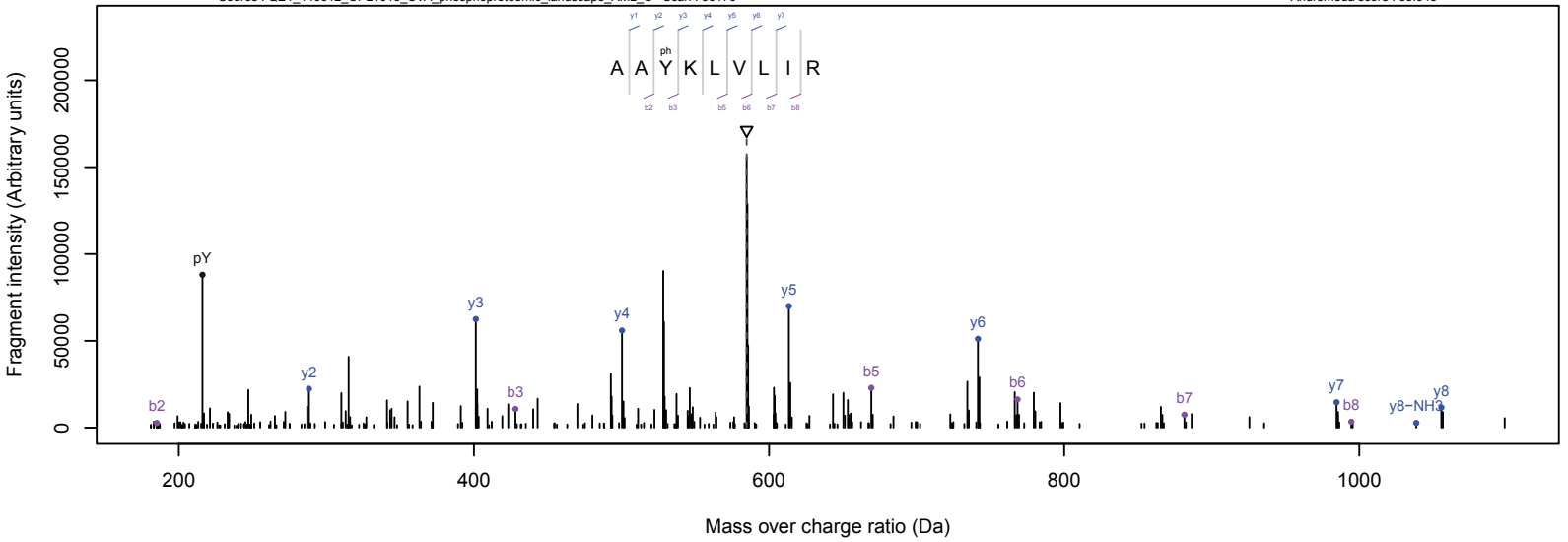
Andromeda score: 84.011



(ac)AAY(ph)KLVLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36176

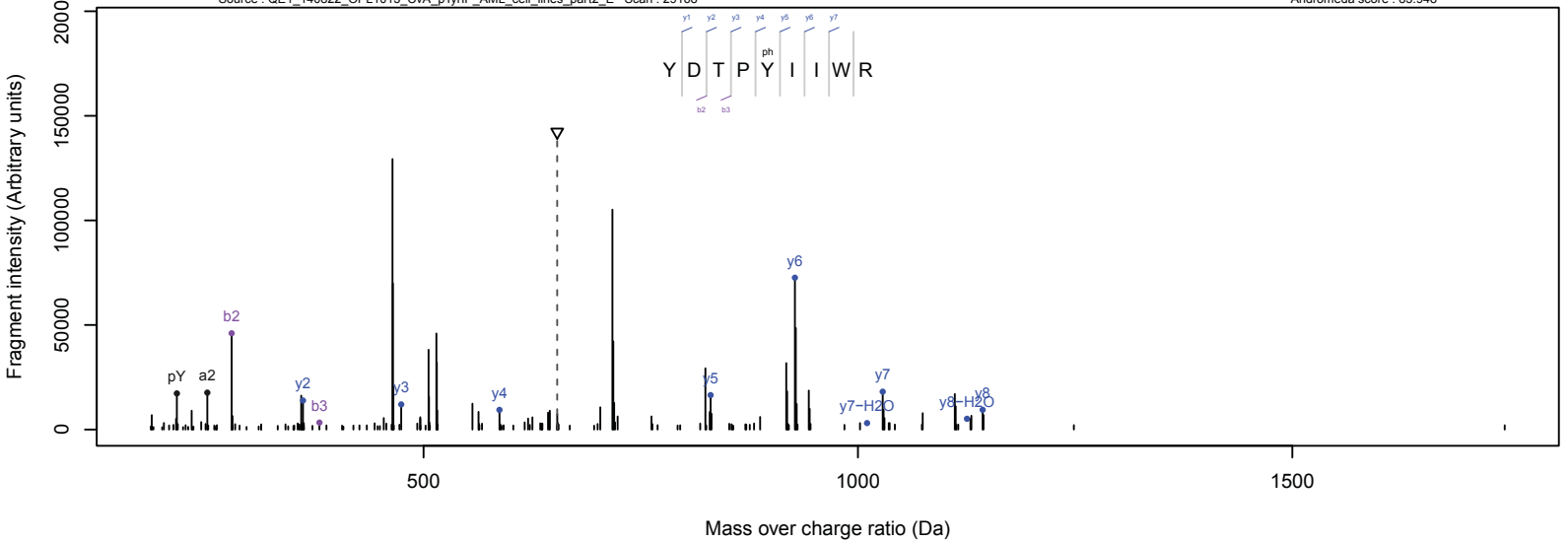
Andromeda score : 83.948



YDTPY(ph)IWR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 29168

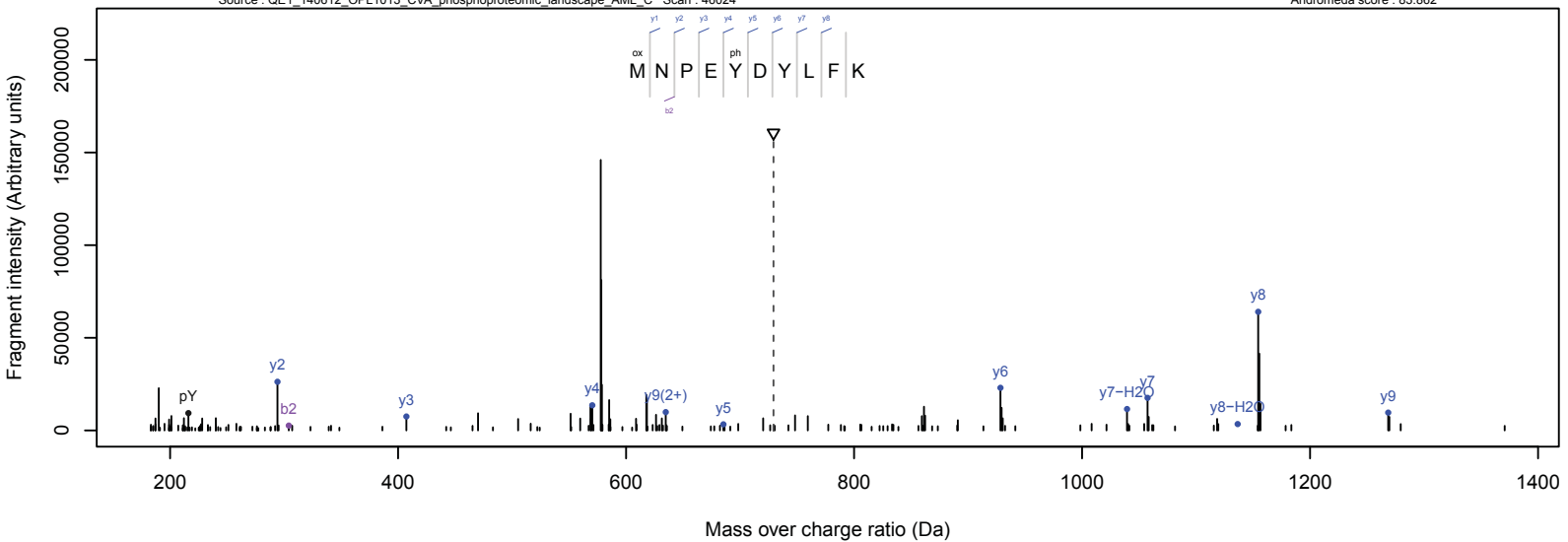
Andromeda score : 83.948



(ac)M(ox)NPEY(ph)DYLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46024

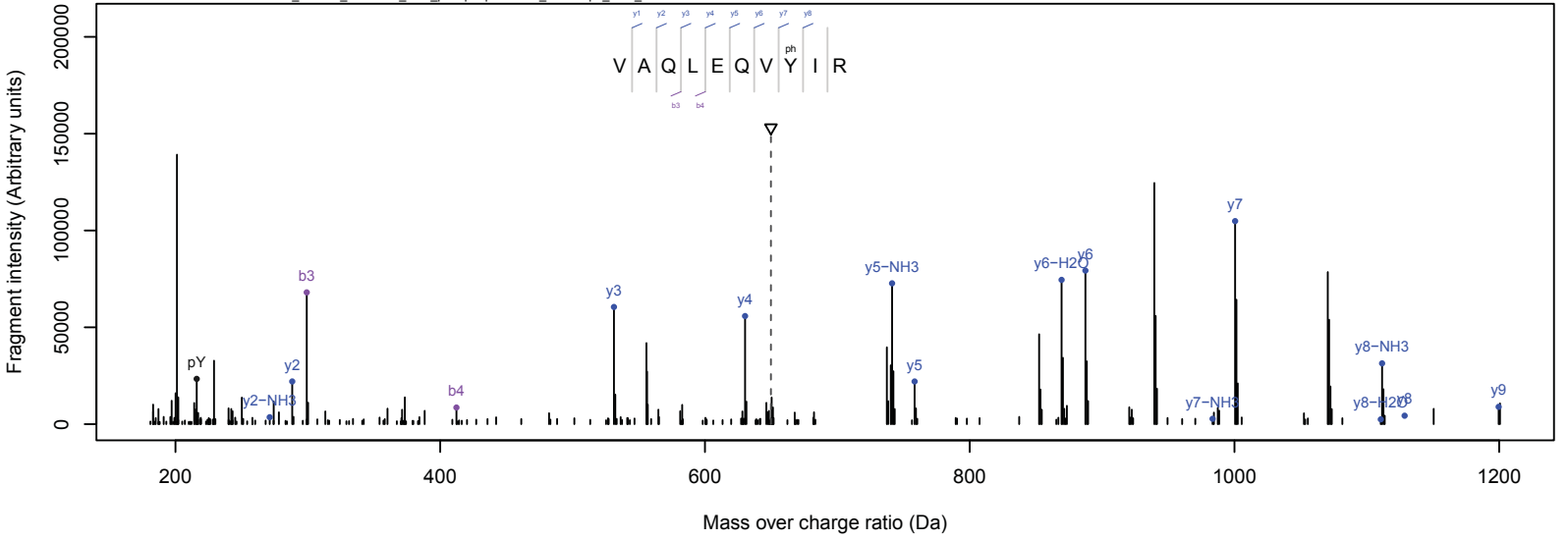
Andromeda score : 83.862



VAQLEQVY(ph)IR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24670

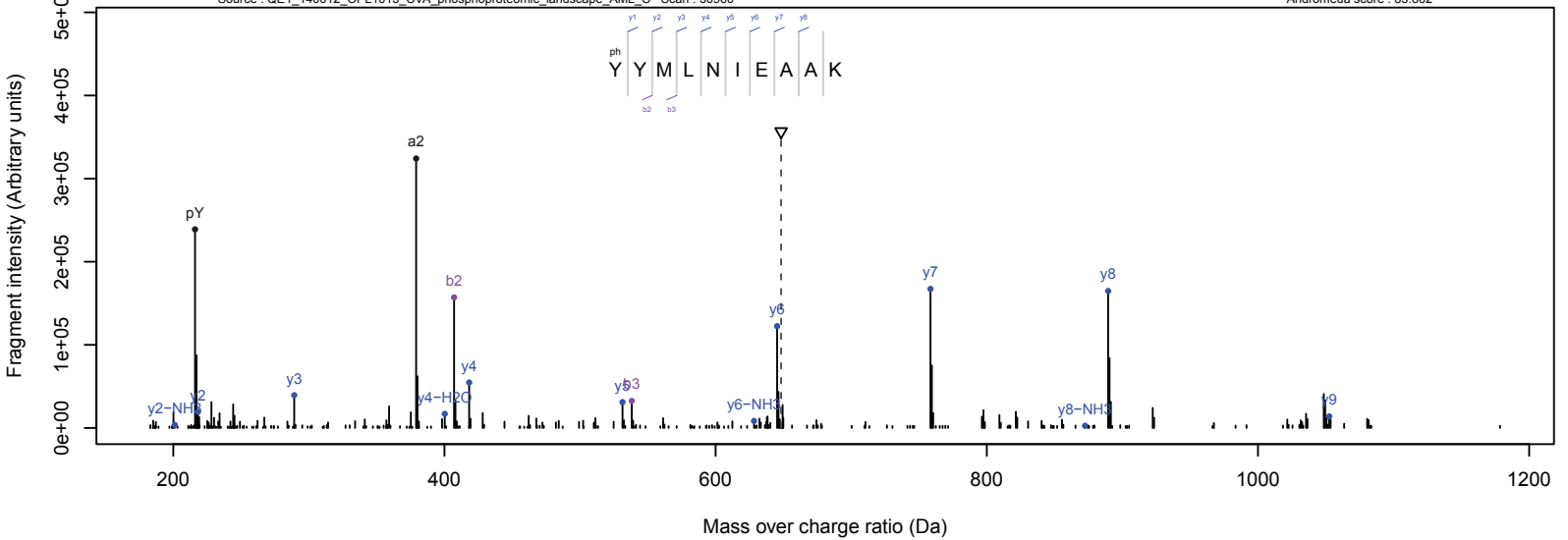
Andromeda score : 83.862



Y(ph)YMLNIEAAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30960

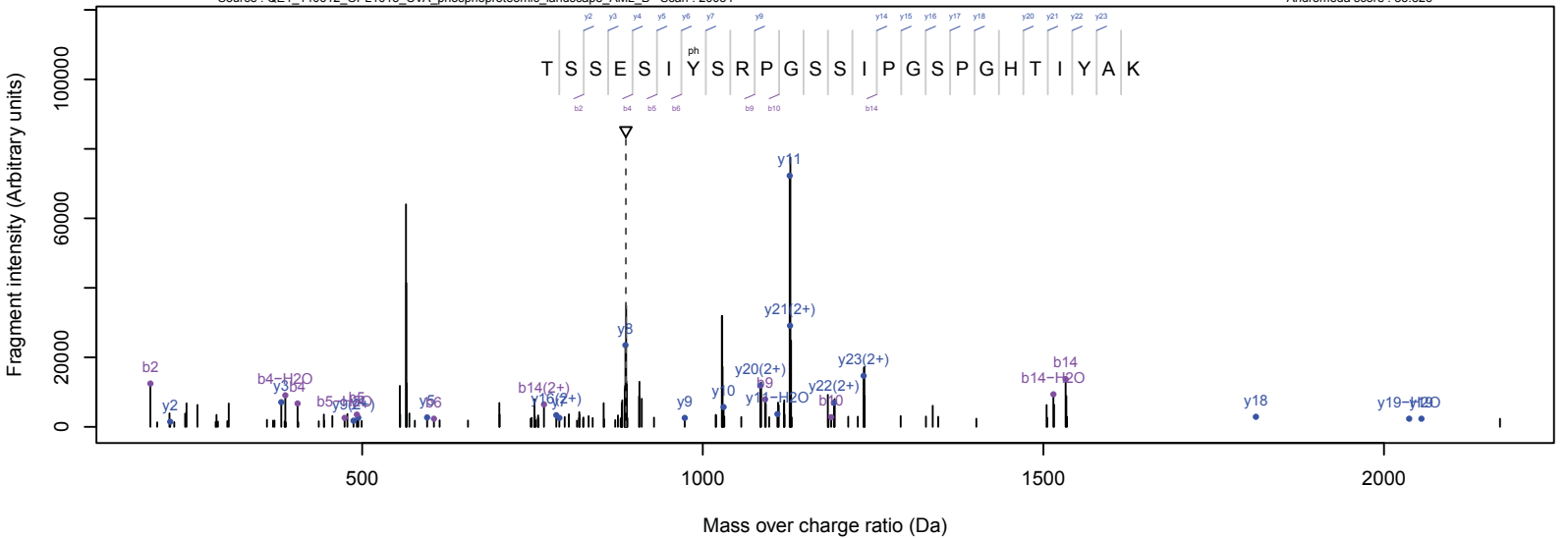
Andromeda score : 83.862



TSSESIY(ph)SRPGSSIPGSPGHTIYAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20034

Andromeda score : 83.826

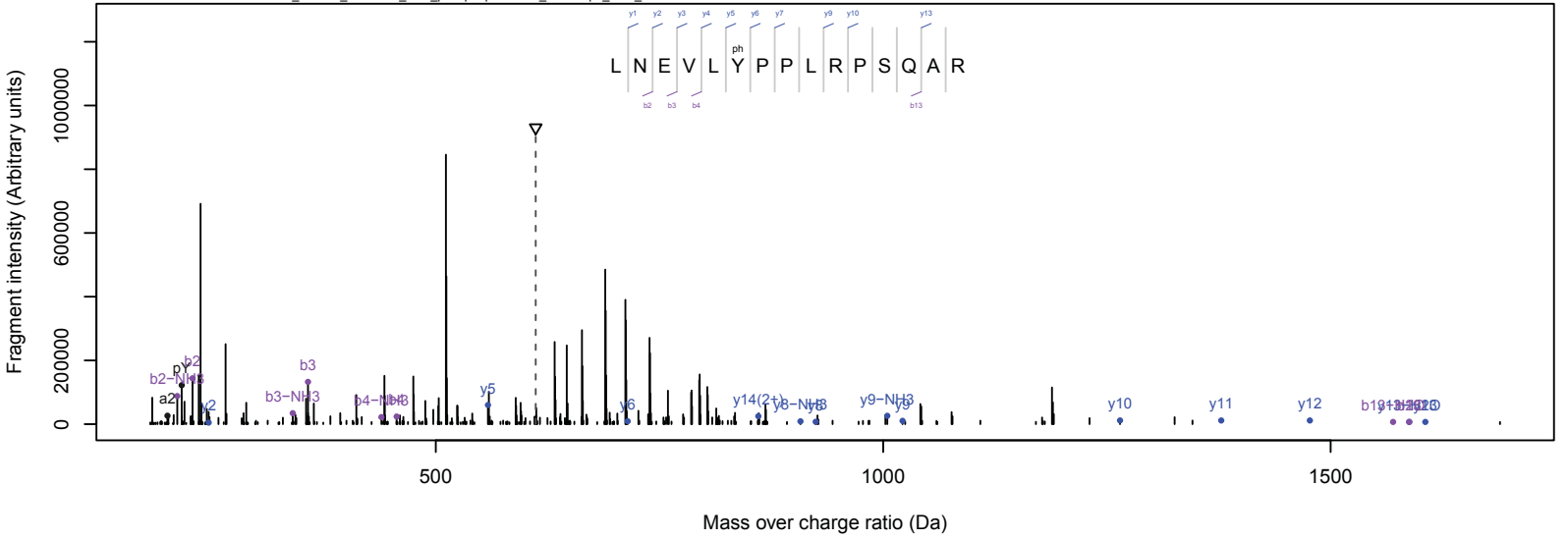




LNEVLY(ph)PPLRPSQAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28951

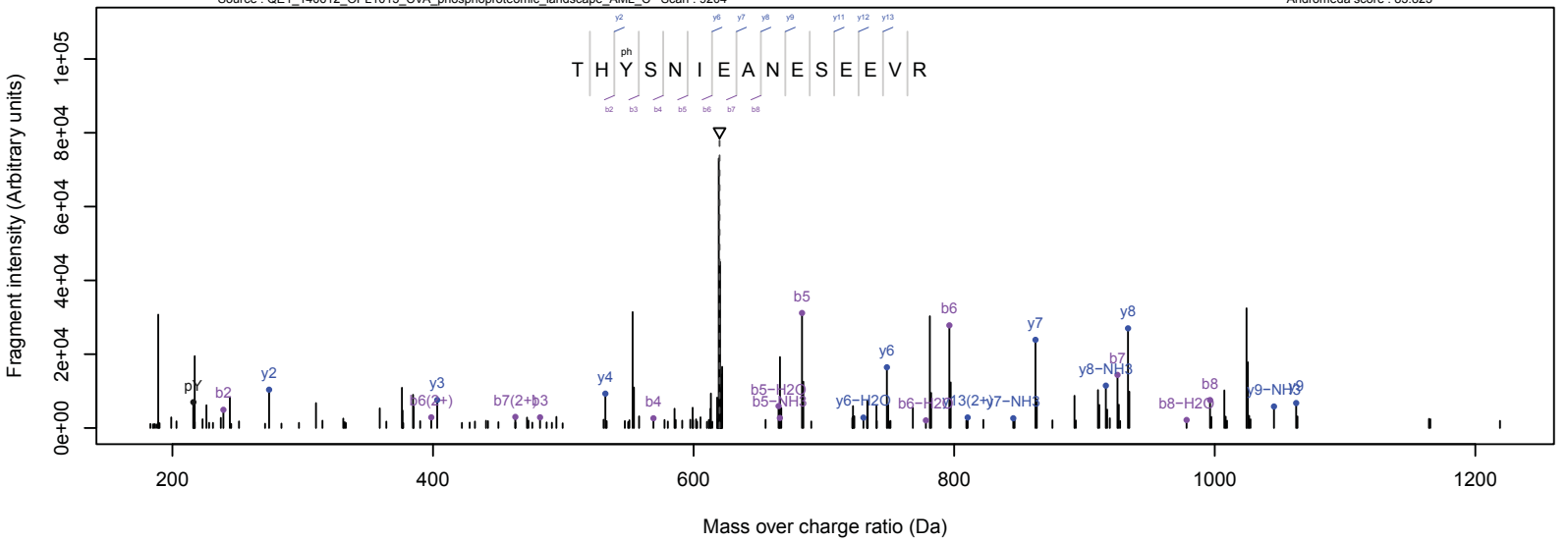
Andromeda score : 83.823



THY(ph)SNIEANESEEV-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9204

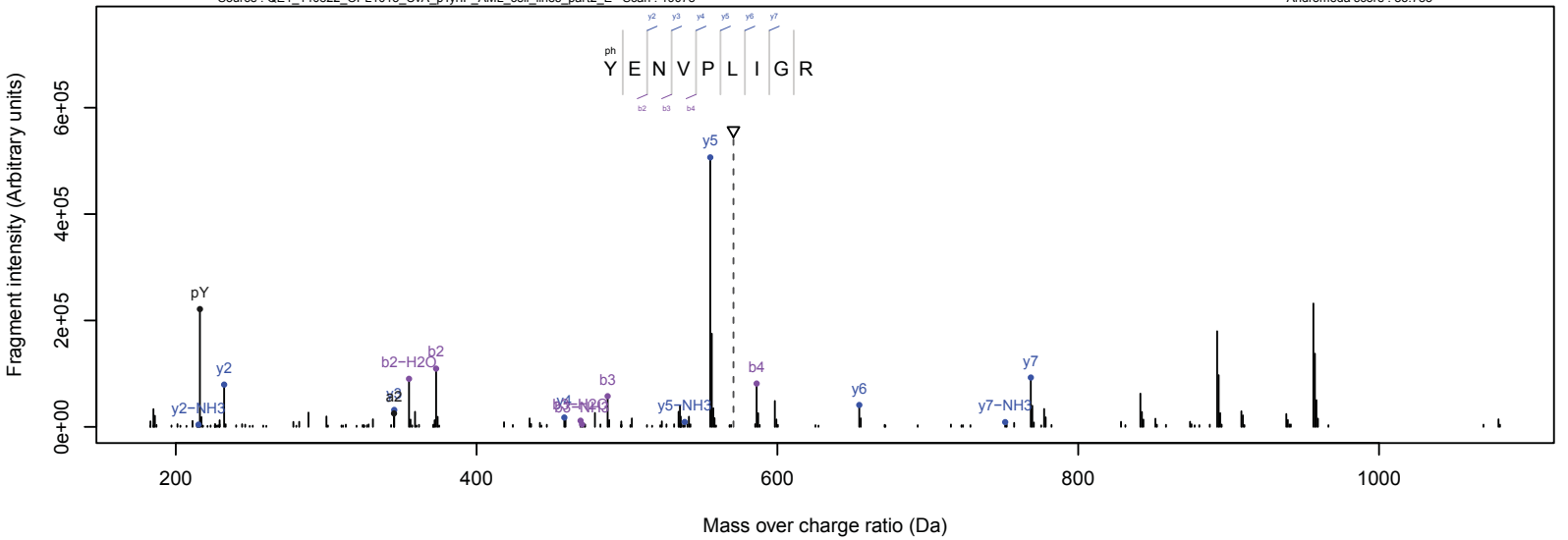
Andromeda score : 83.823



Y(ph)ENVPLIGR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 19078

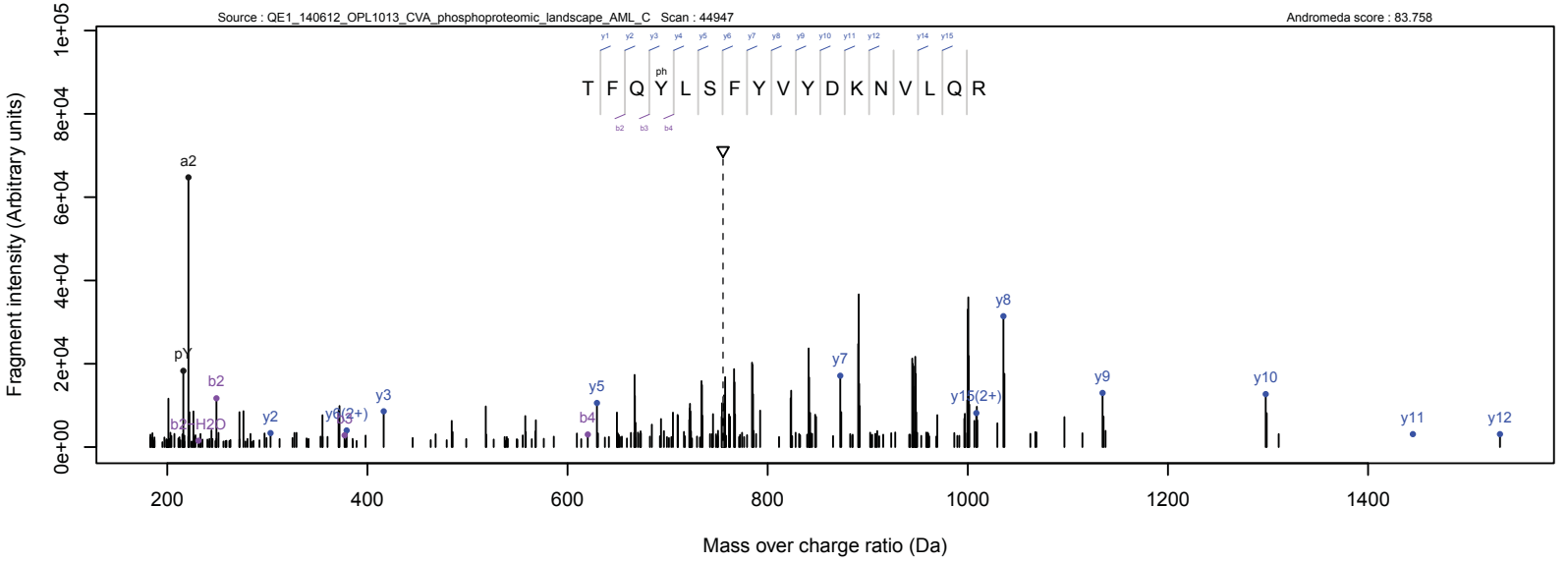
Andromeda score : 83.783



**TFQY(ph)LSFYVDKNVLQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44947

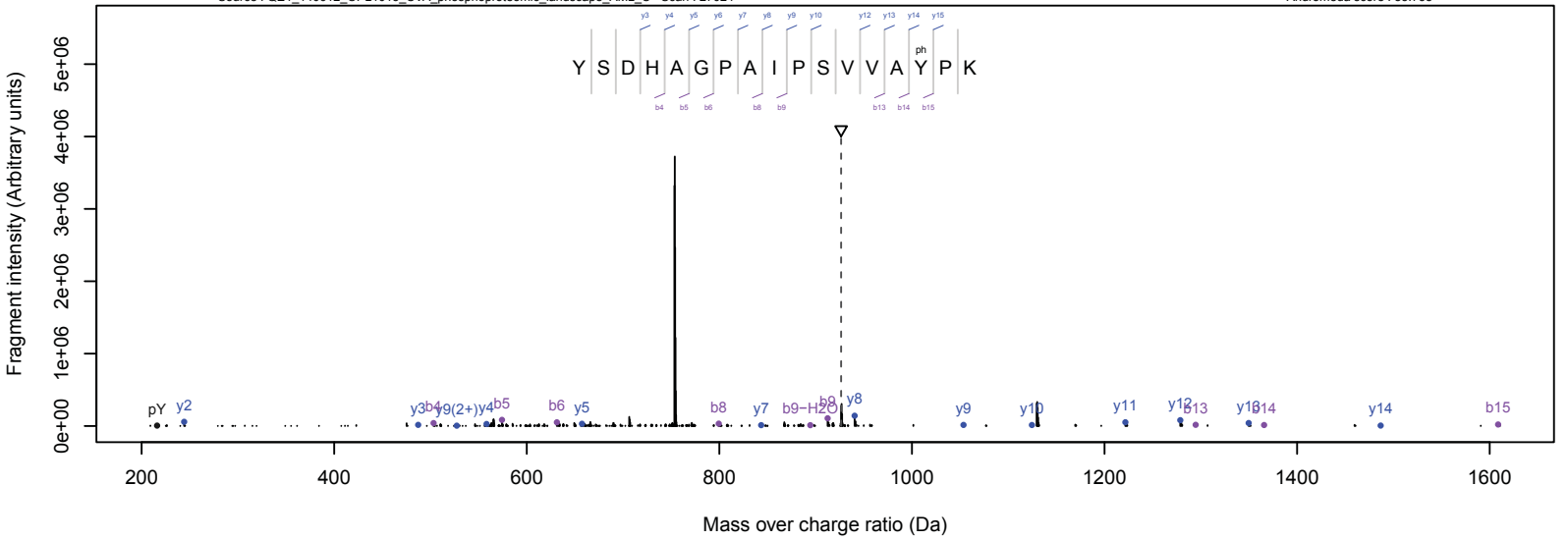
Andromeda score : 83.758



**YSDHAGPAIPSVVAY(ph)PK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27024

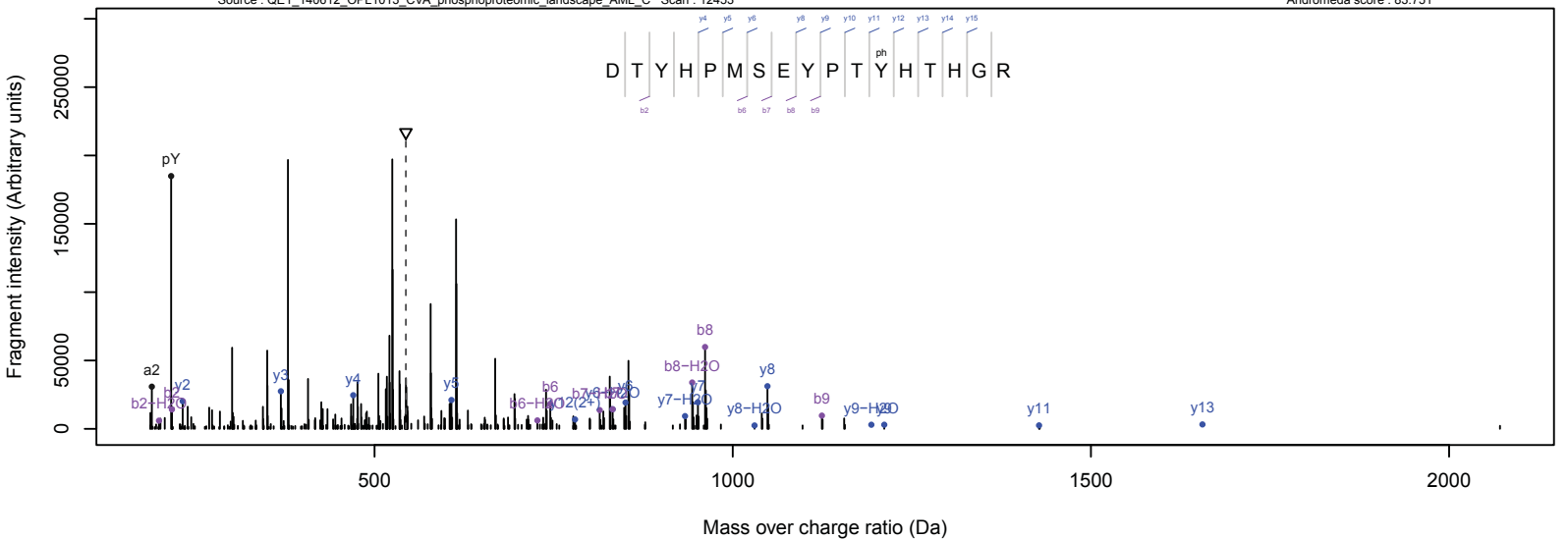
Andromeda score : 83.758



**DTYHPMSEYPTYPHTHGR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12453

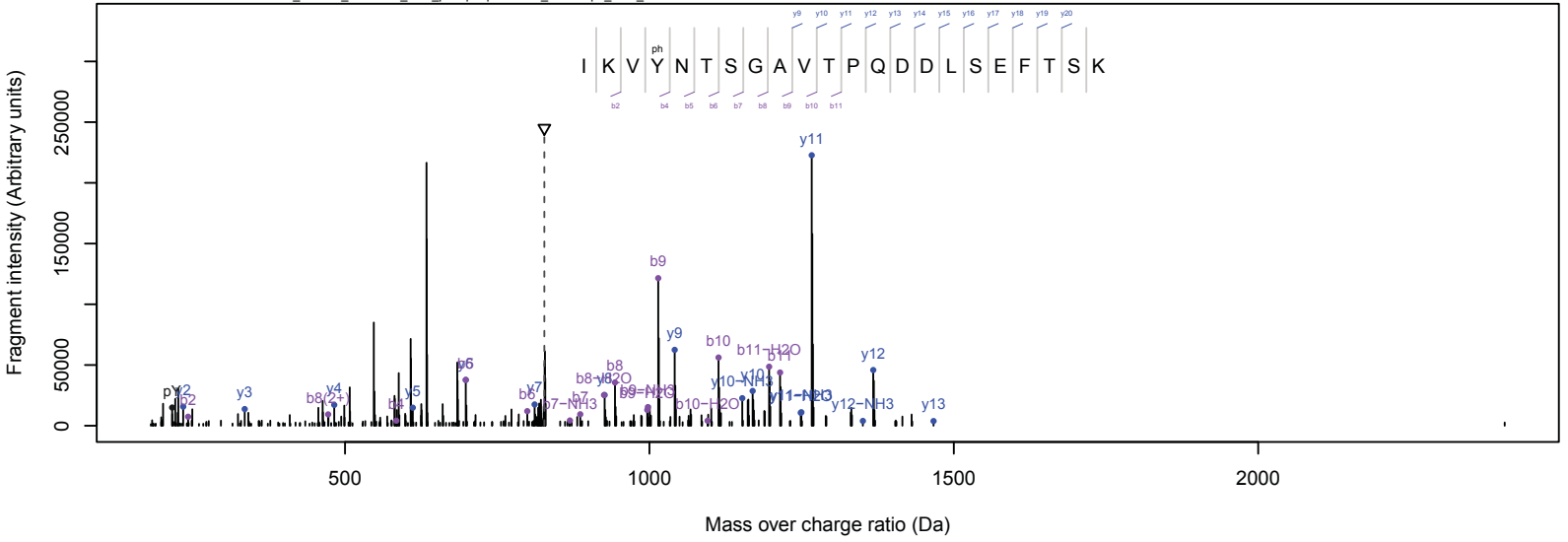
Andromeda score : 83.751



**IKVY(ph)NTSGAVTPQDDLSEFTSK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31124

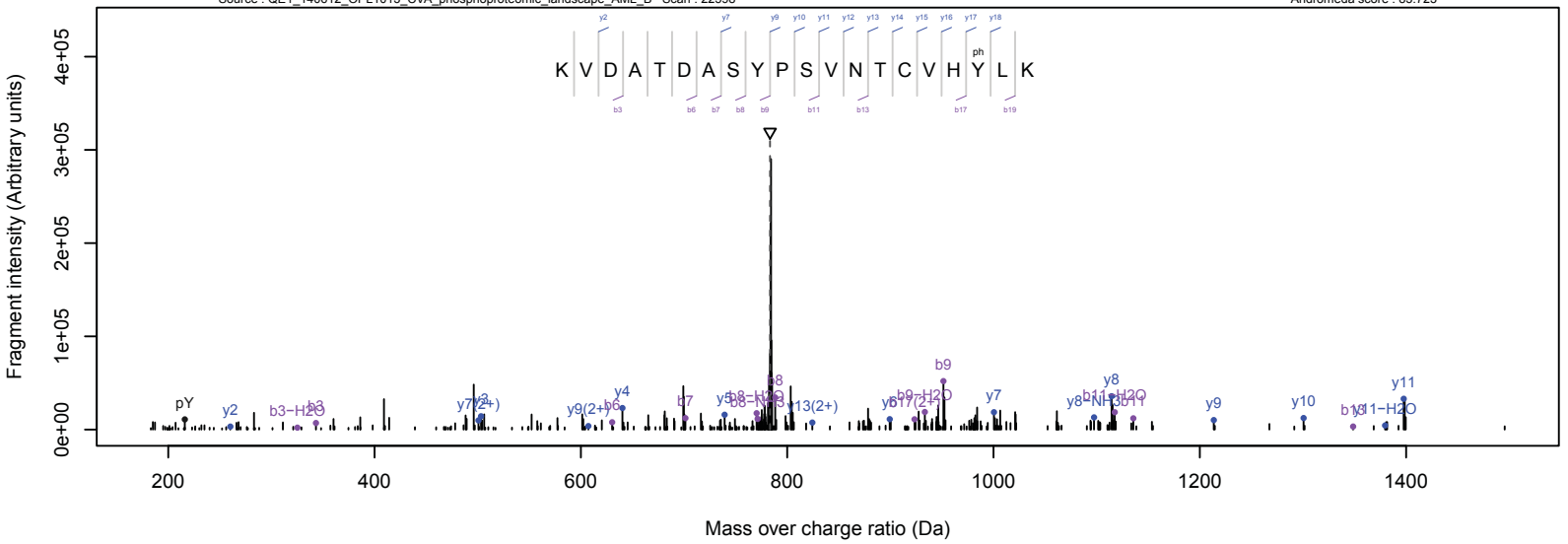
Andromeda score : 83.727



**KVDATDASYPNVTCVHY(ph)LK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22598

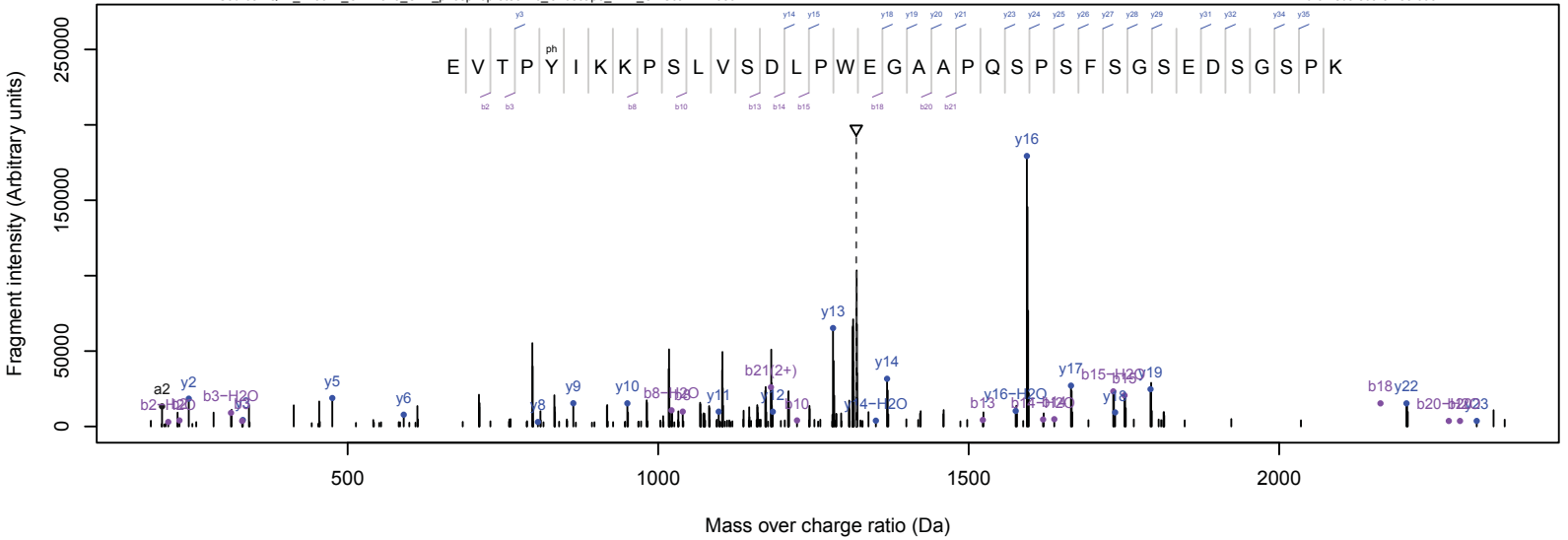
Andromeda score : 83.723



**EVTPY(ph)IKKPSLVSDLPWEGAAPQSPSFSGSEDSGSPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42089

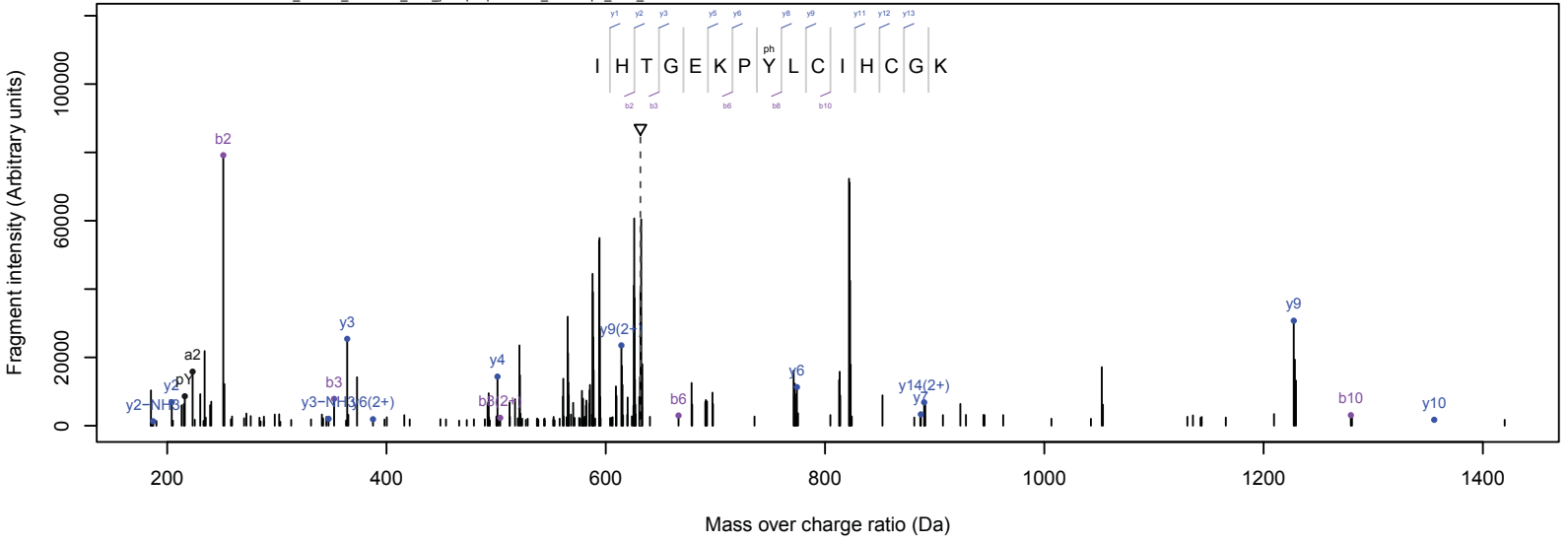
Andromeda score : 83.693



IHTGEKPYLCIHCGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 13257

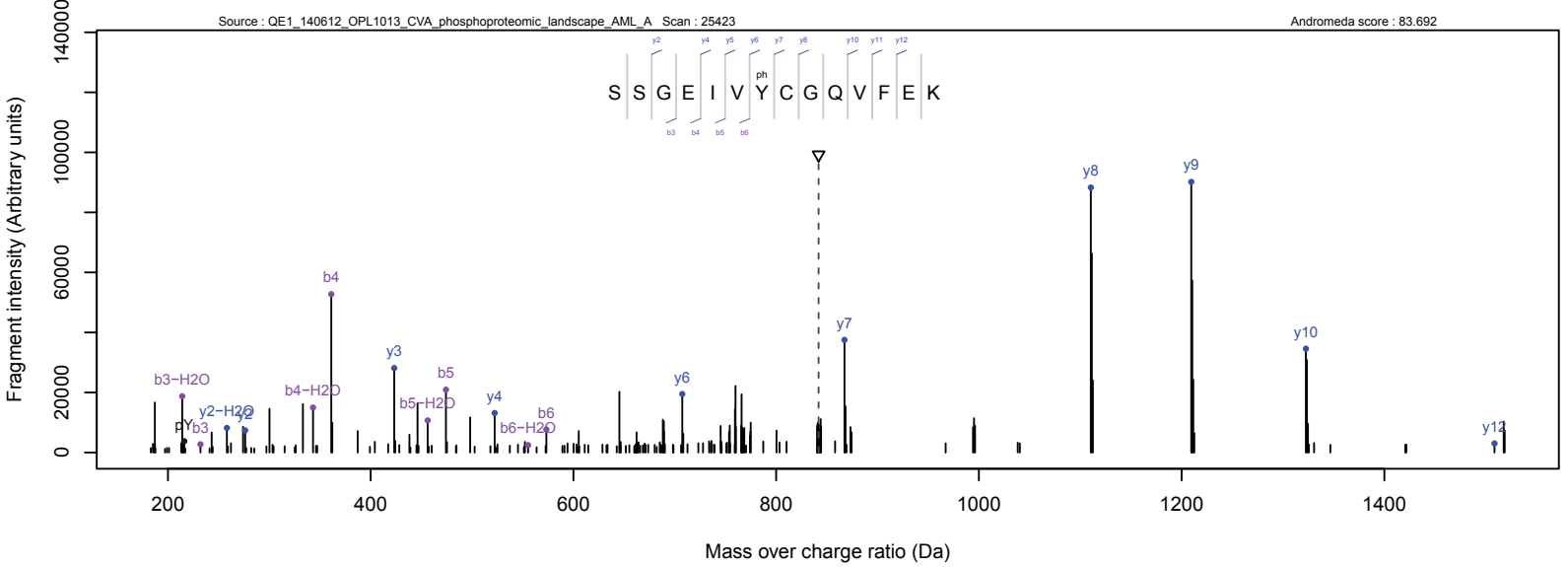
Andromeda score : 83.692



SSGEIVYCGQVFEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25423

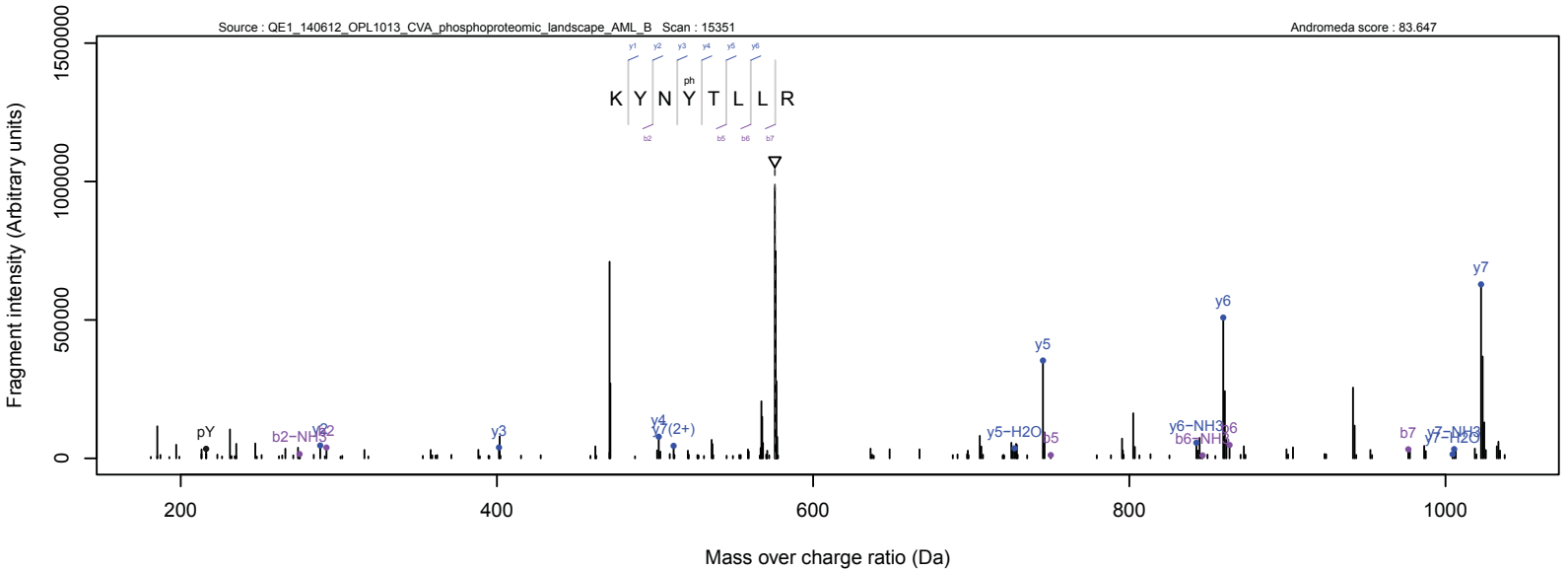
Andromeda score : 83.692

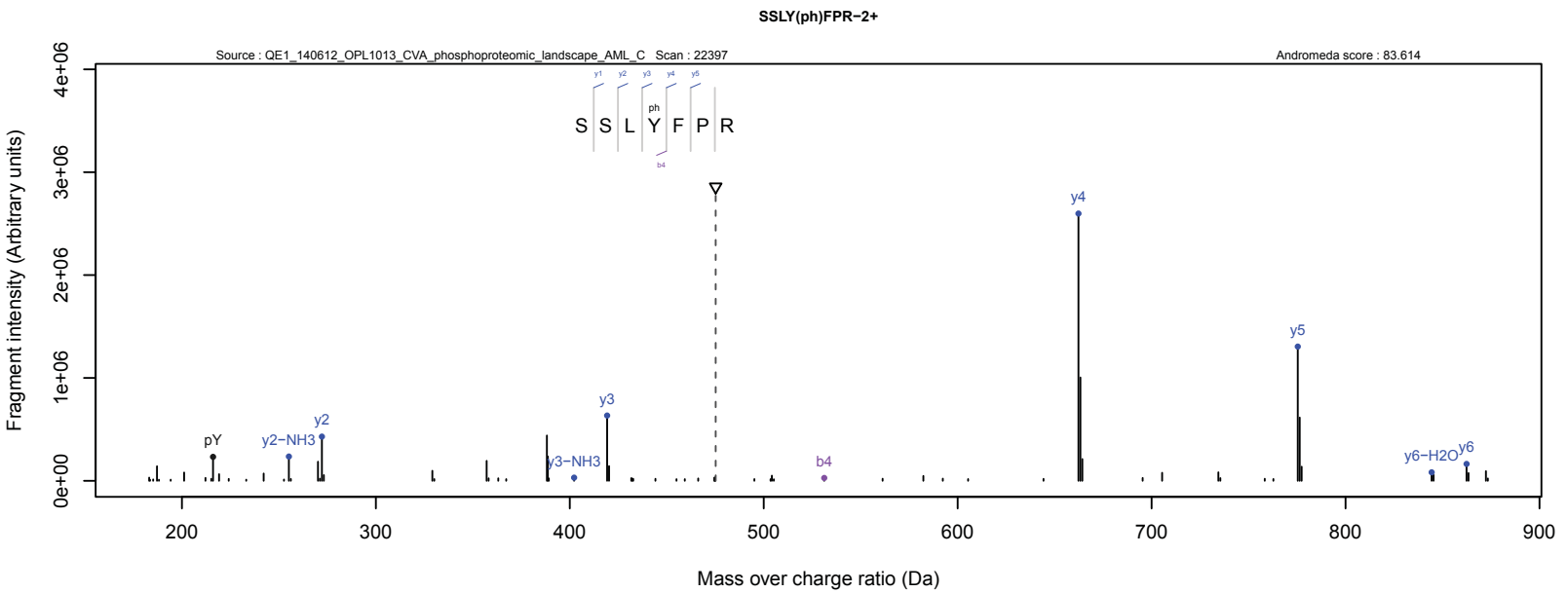
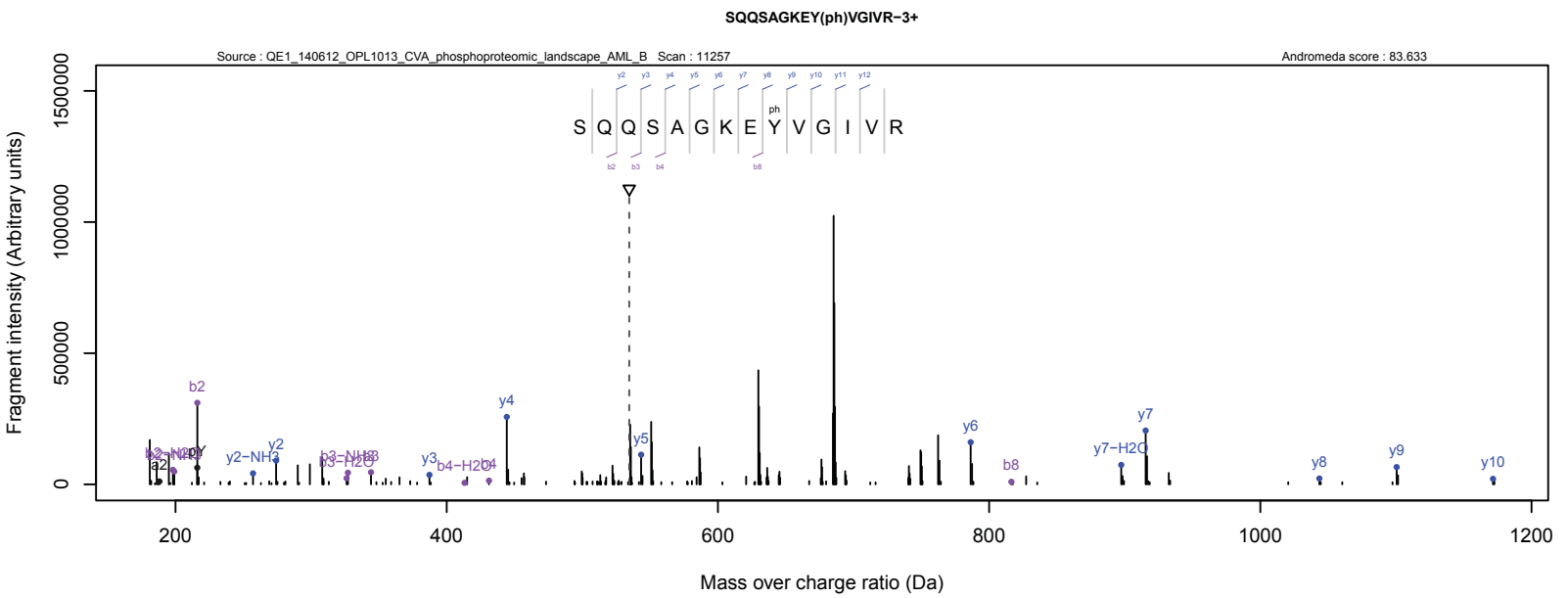
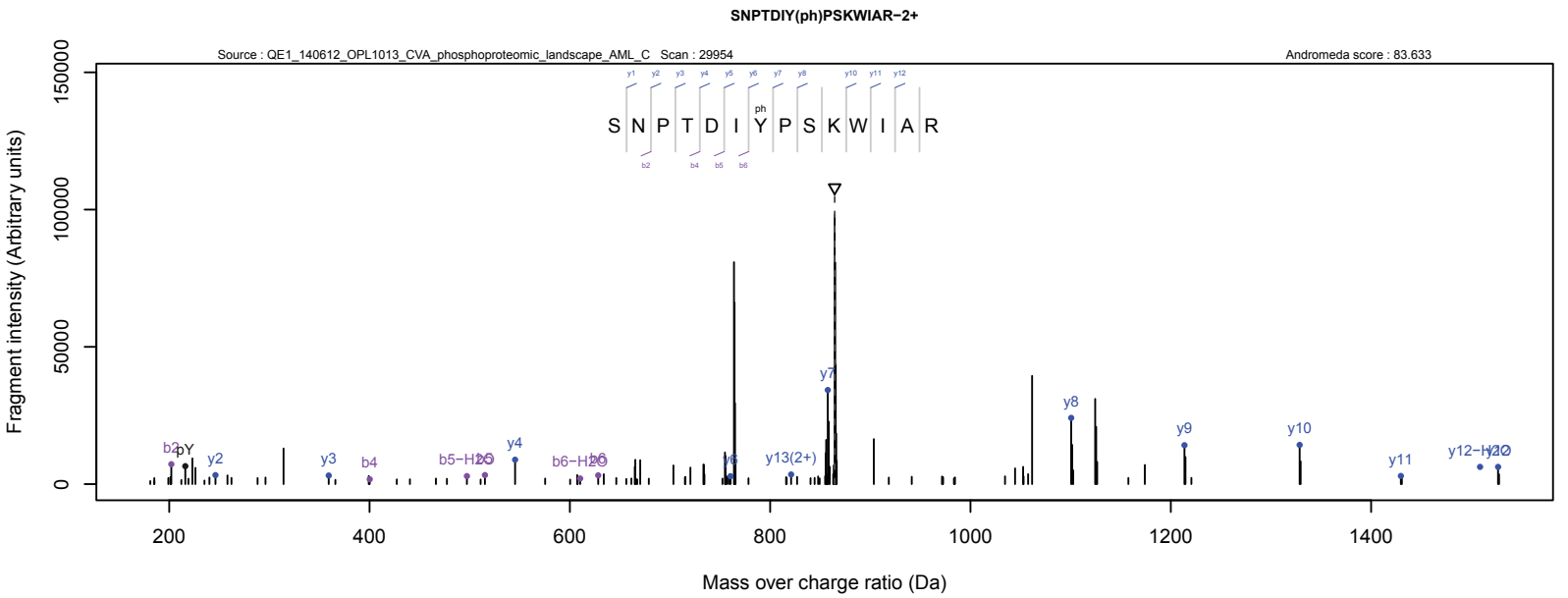


KYNYTLLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15351

Andromeda score : 83.647

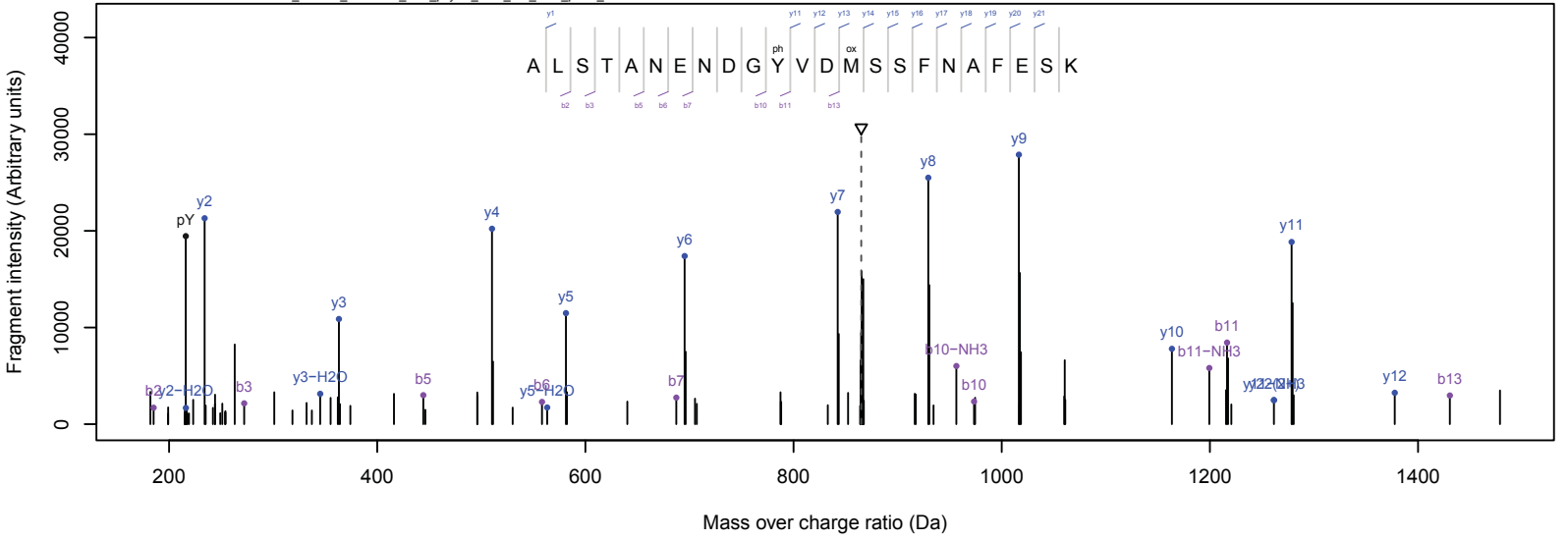




**ALSTANENDGY(ph)VDM(ox)SSFNAFESK-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 27308

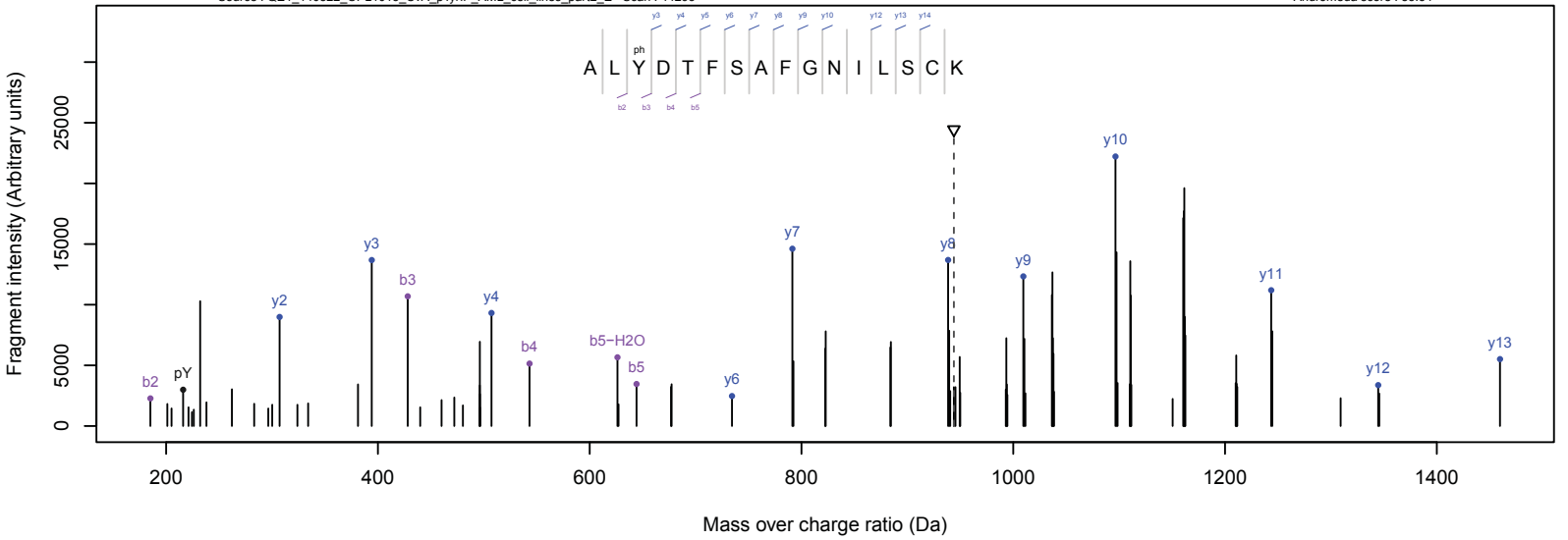
Andromeda score : 83.602



**ALY(ph)DTFSAFGNILSCK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 44293

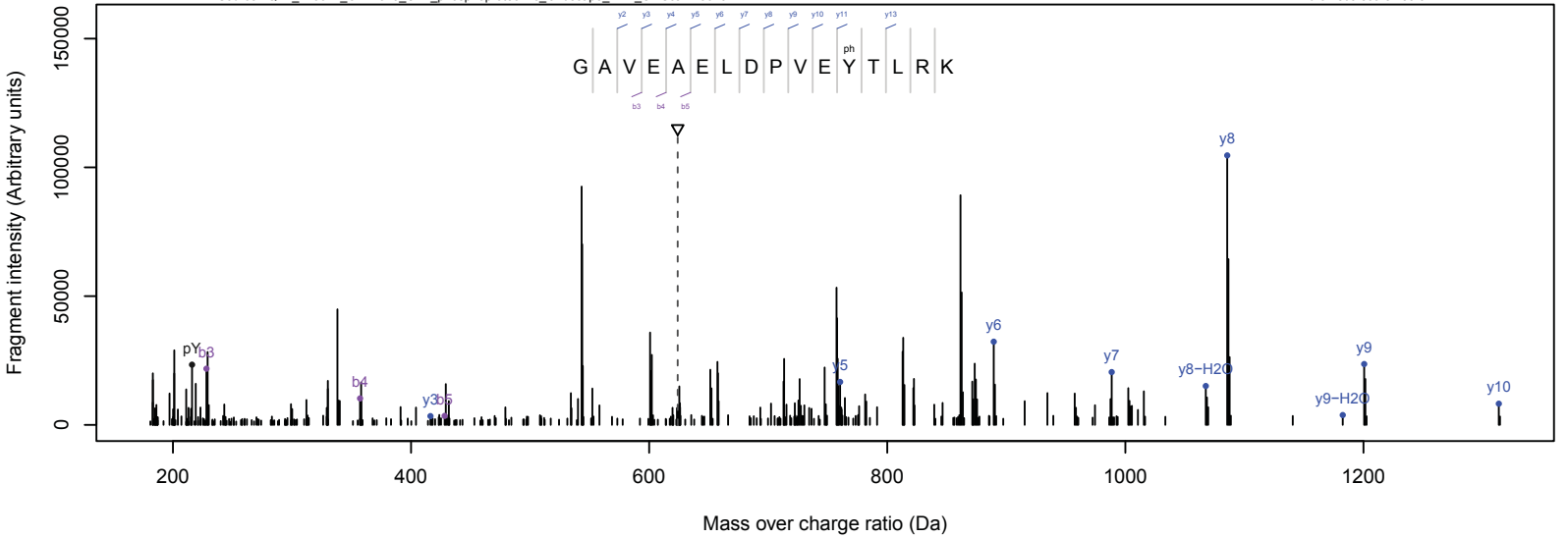
Andromeda score : 83.54



**GAVEAELDPVEY(ph)TLRK-3+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 36152

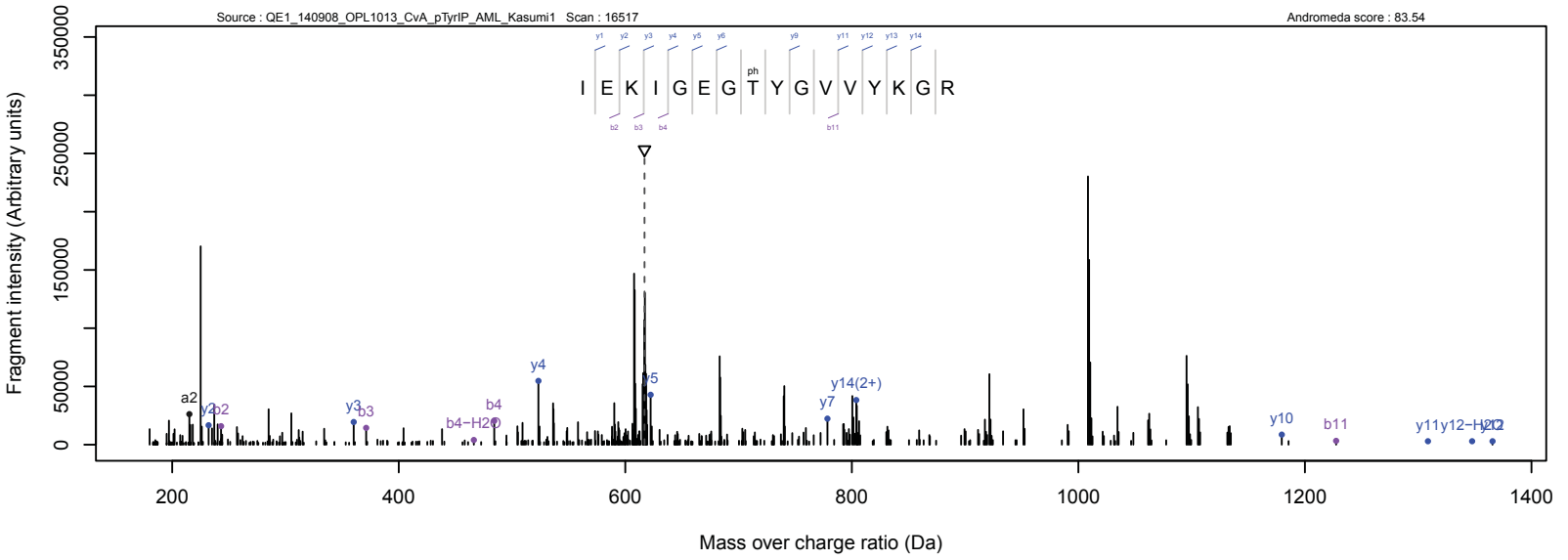
Andromeda score : 83.54



**IEKIGEGT(ph)YGVVYKGR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 16517

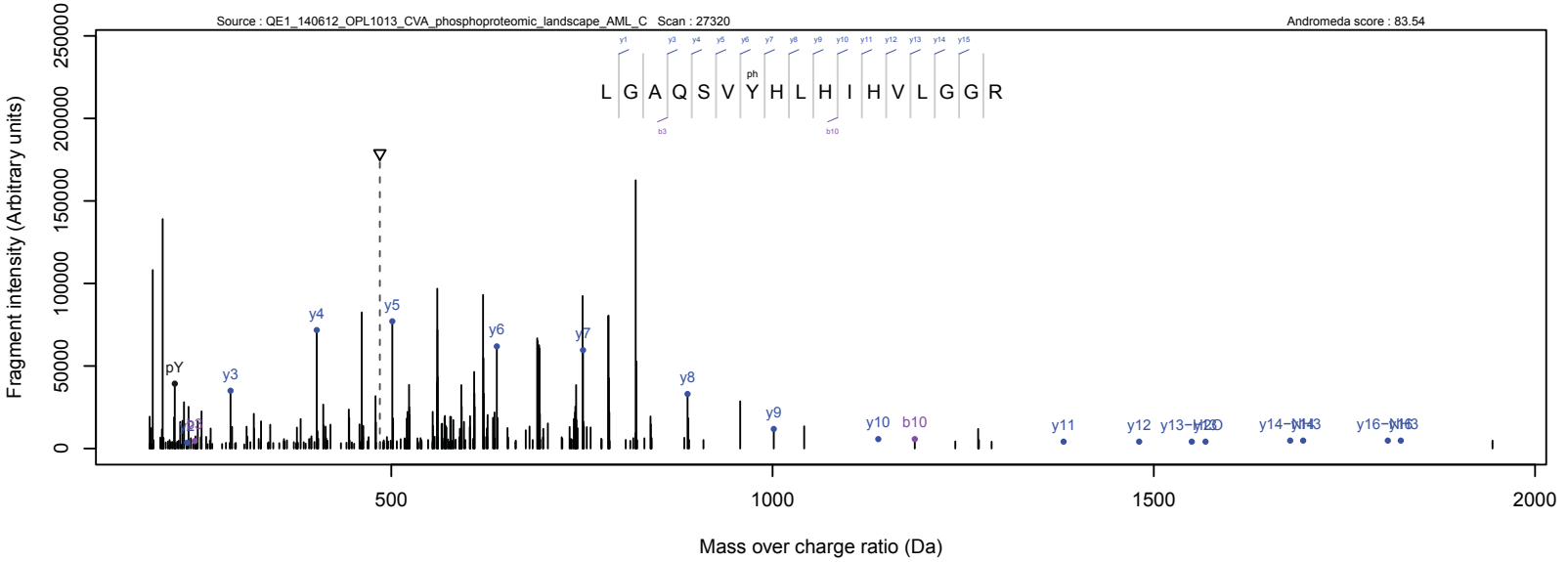
Andromeda score : 83.54



**LGAQSVY(ph)HLHIHLGGR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27320

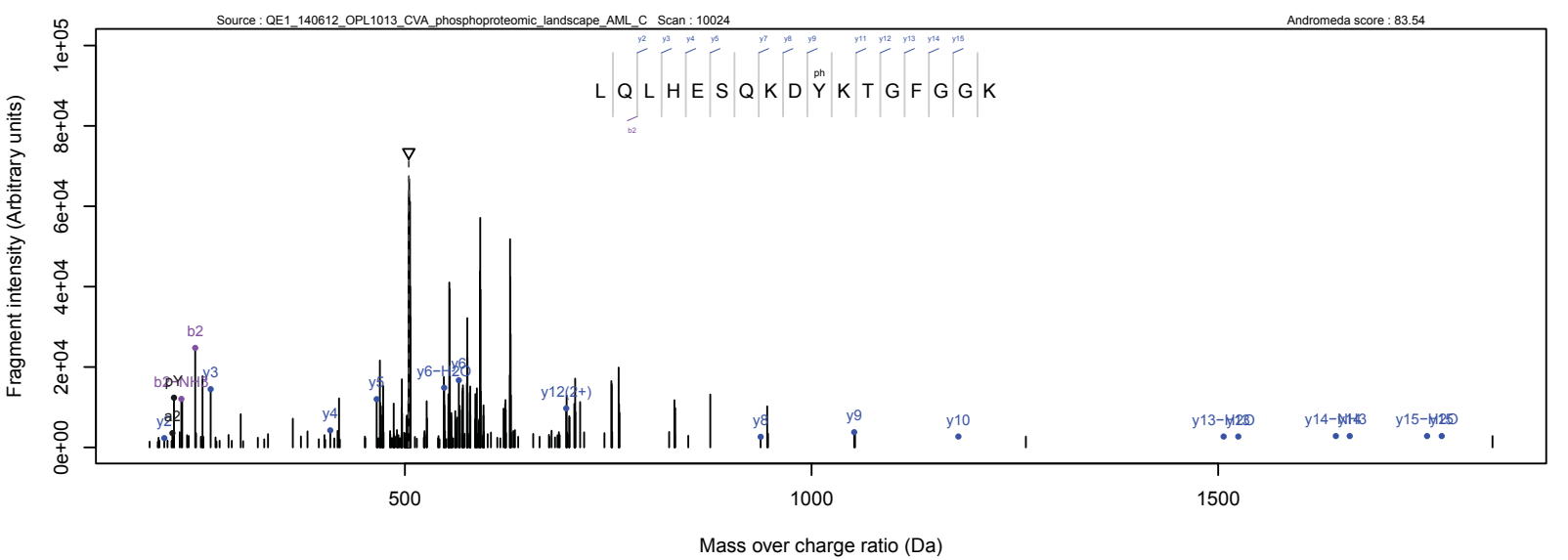
Andromeda score : 83.54



**LQLHESQKD(ph)YKTGFGGK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10024

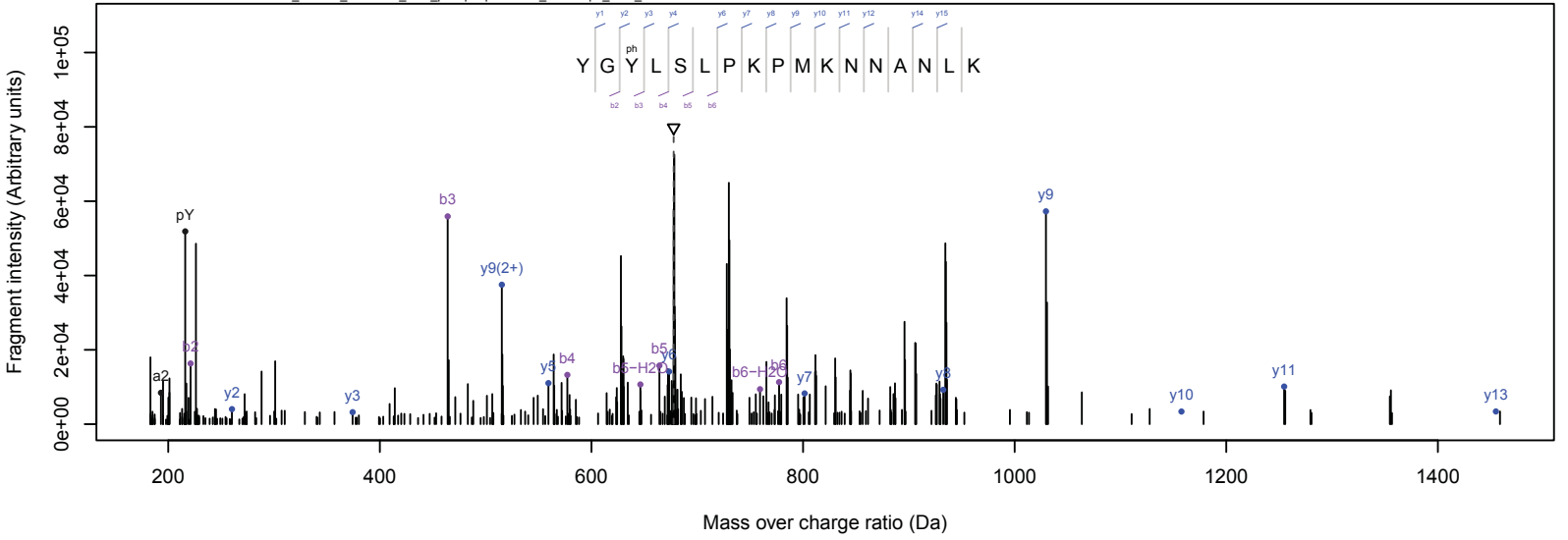
Andromeda score : 83.54



YG<sup>(ph)</sup>L<sup>(ph)</sup>S<sup>(ph)</sup>L<sup>(ph)</sup>P<sup>(ph)</sup>K<sup>(ph)</sup>M<sup>(ph)</sup>K<sup>(ph)</sup>N<sup>(ph)</sup>N<sup>(ph)</sup>A<sup>(ph)</sup>N<sup>(ph)</sup>L<sup>(ph)</sup>K<sup>(ph)</sup>-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25898

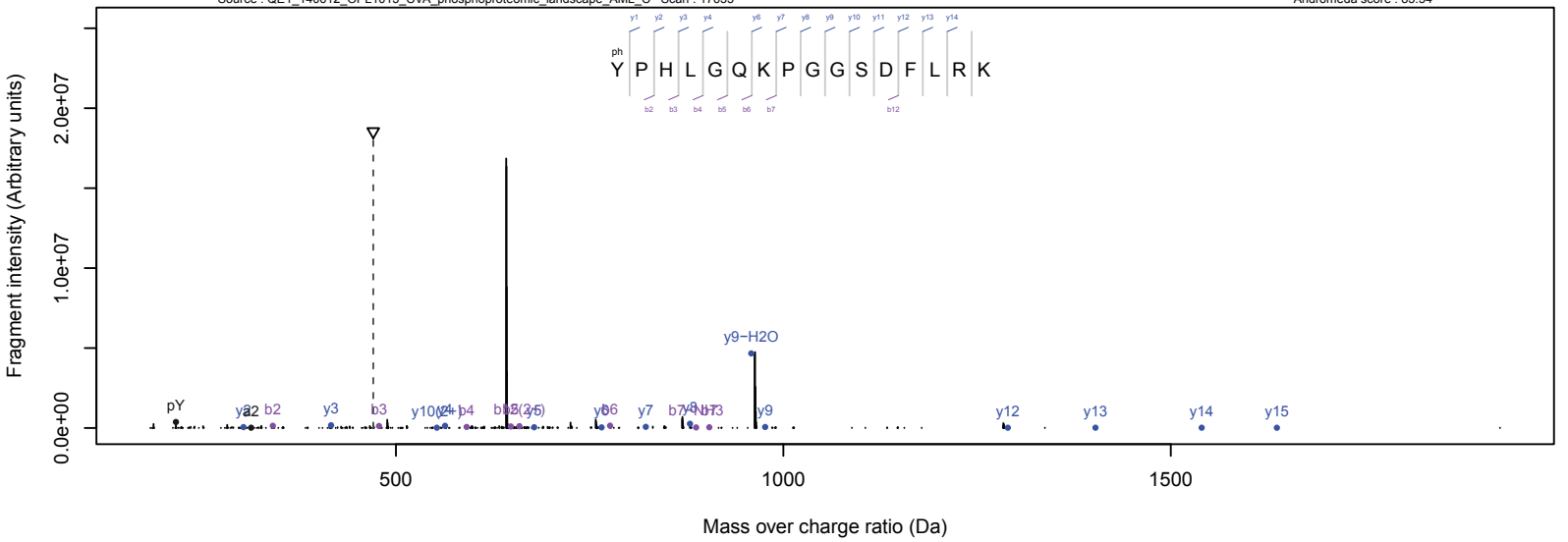
Andromeda score : 83.54



Y<sup>(ph)</sup>P<sup>(ph)</sup>H<sup>(ph)</sup>L<sup>(ph)</sup>G<sup>(ph)</sup>Q<sup>(ph)</sup>K<sup>(ph)</sup>P<sup>(ph)</sup>G<sup>(ph)</sup>G<sup>(ph)</sup>S<sup>(ph)</sup>D<sup>(ph)</sup>F<sup>(ph)</sup>L<sup>(ph)</sup>R<sup>(ph)</sup>K<sup>(ph)</sup>-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17635

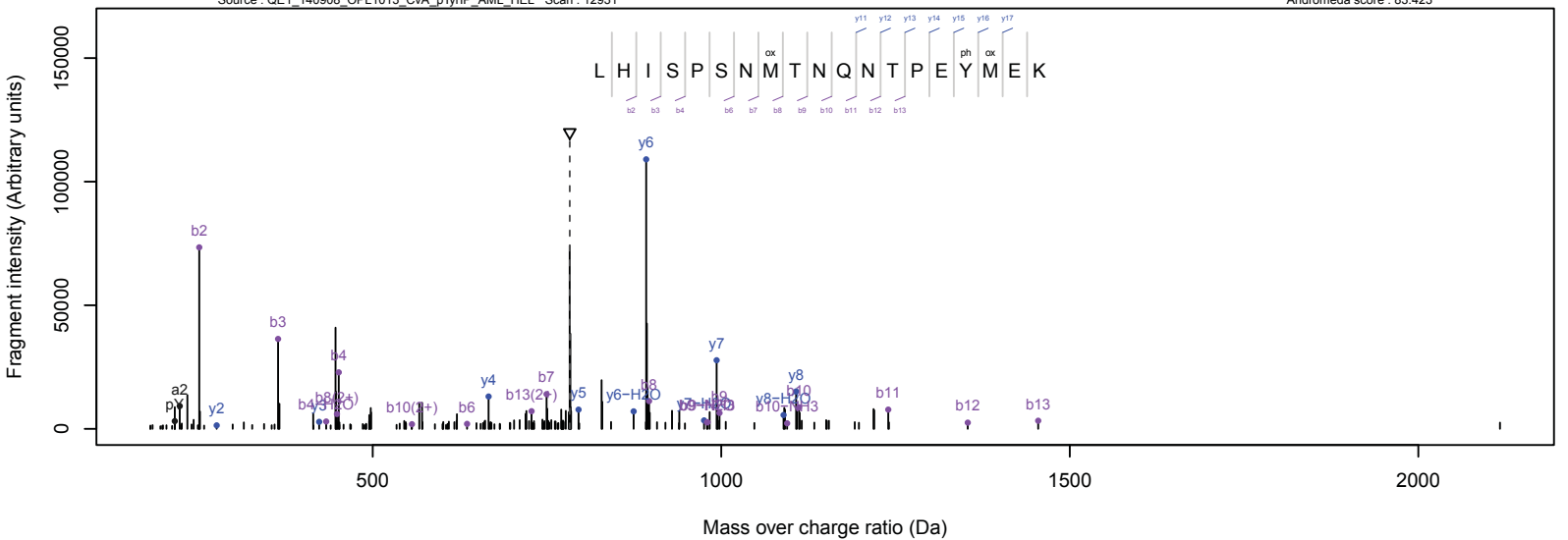
Andromeda score : 83.54



LHISPSN<sup>(ox)</sup>M<sup>(ox)</sup>TNQNTPEY<sup>(ph)</sup>M<sup>(ox)</sup>EK<sup>(ox)</sup>-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 12931

Andromeda score : 83.423

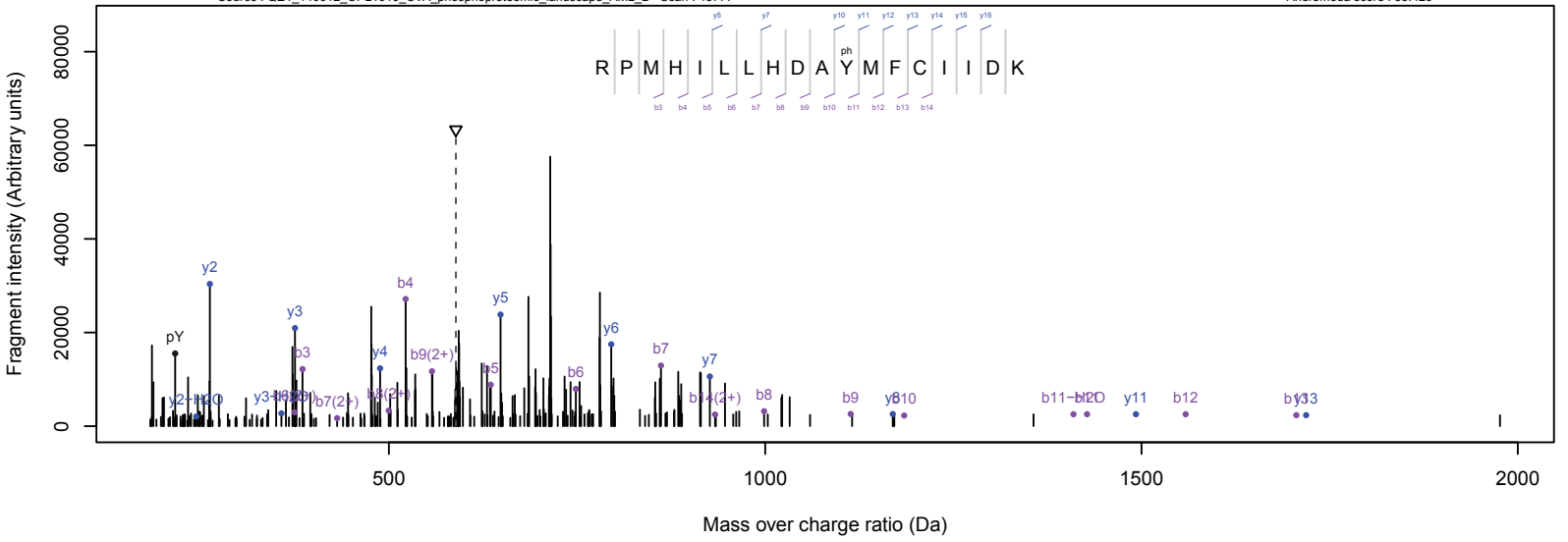




RPMHILLHDAY(ph)MFCIIDK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 43777

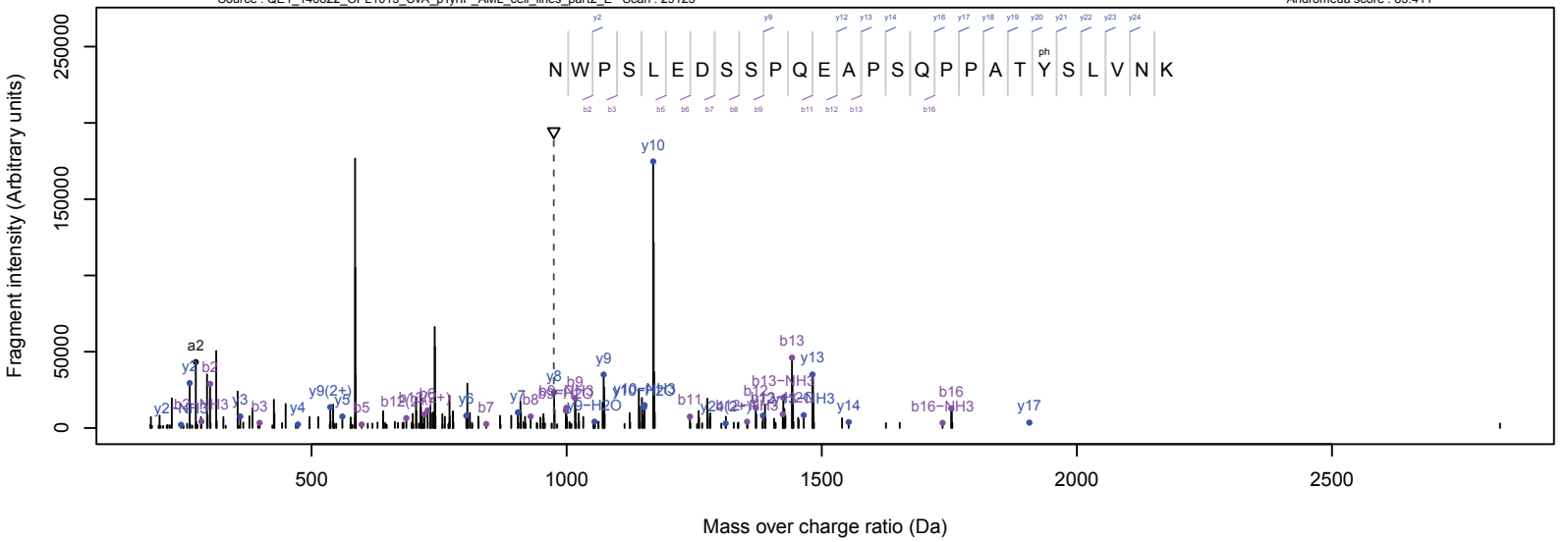
Andromeda score: 83.423



NWPSLEDSSPQEAPSQPATY(ph)SLVNK-3+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 29123

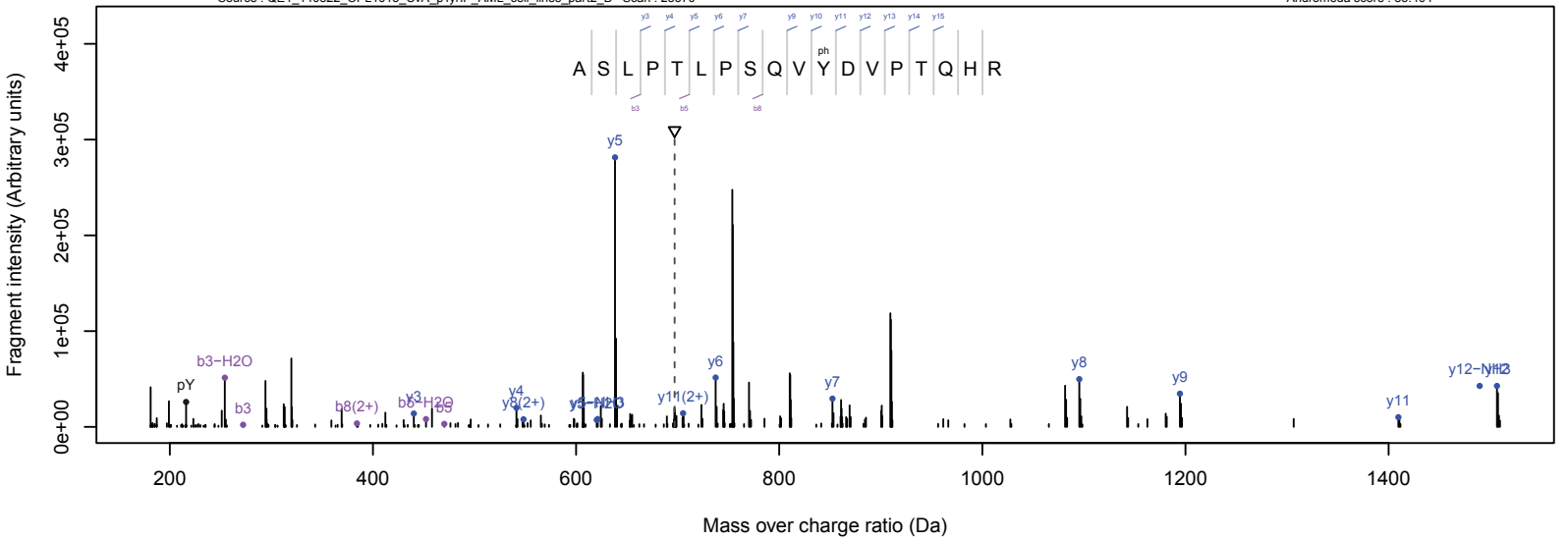
Andromeda score: 83.411



ASLPTLP SQVYDVP TQHR-3+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 25670

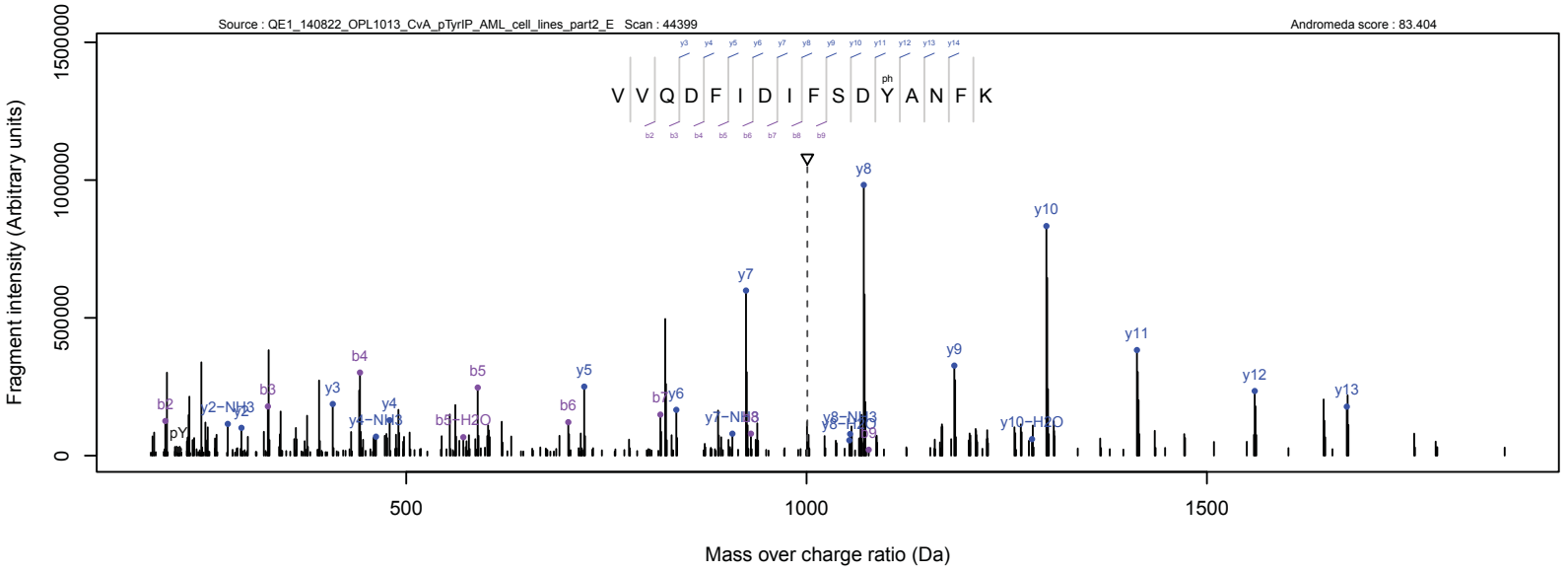
Andromeda score: 83.404



VVQDFIDIFS DY(ph)ANFK-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 44399

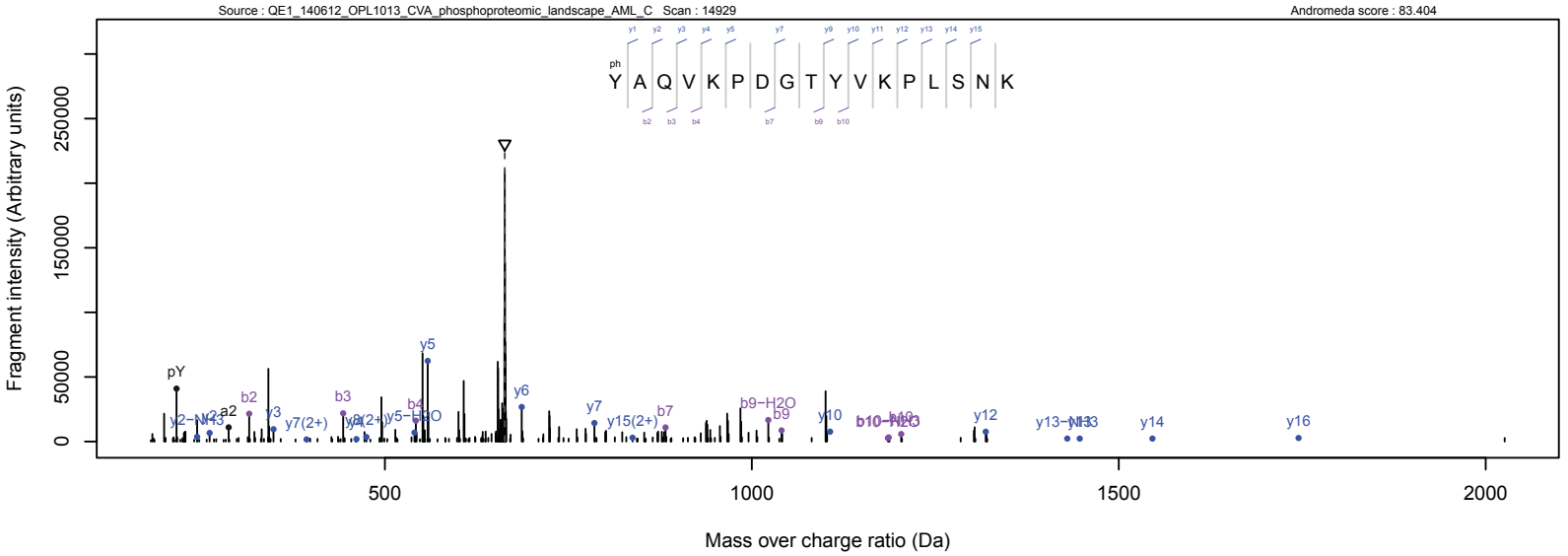
Andromeda score : 83.404



Y(ph)AQVKPDGTYVKPLSNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14929

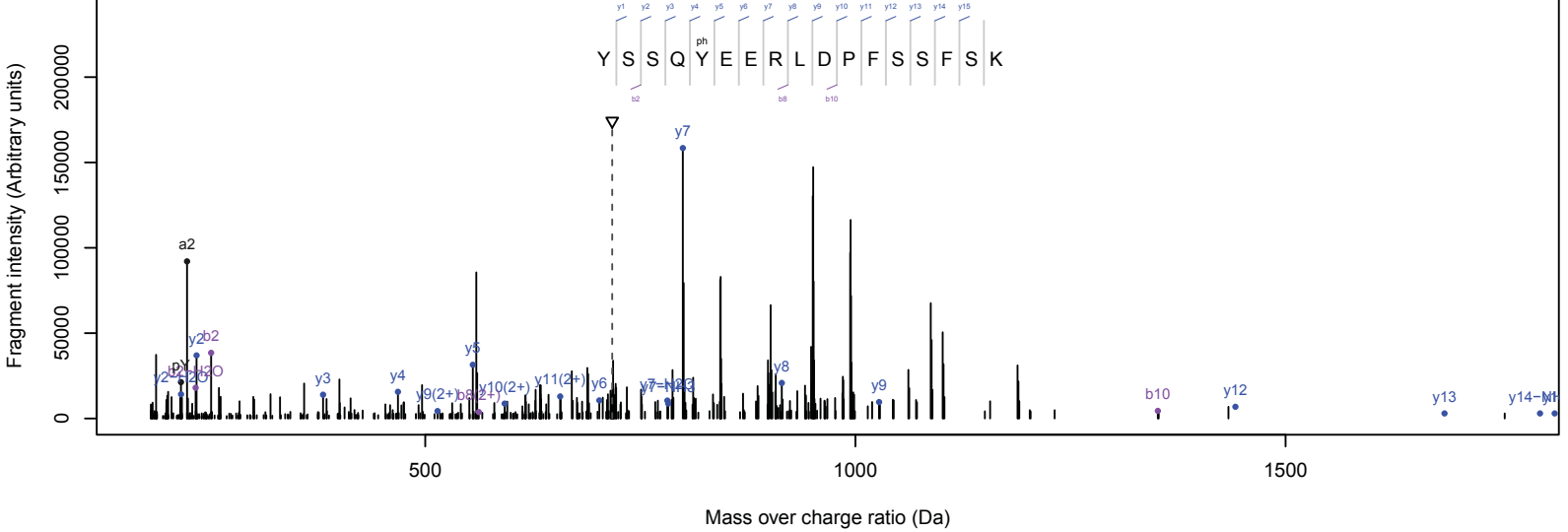
Andromeda score : 83.404



YSSQY(ph)EERLDPFSSFSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35173

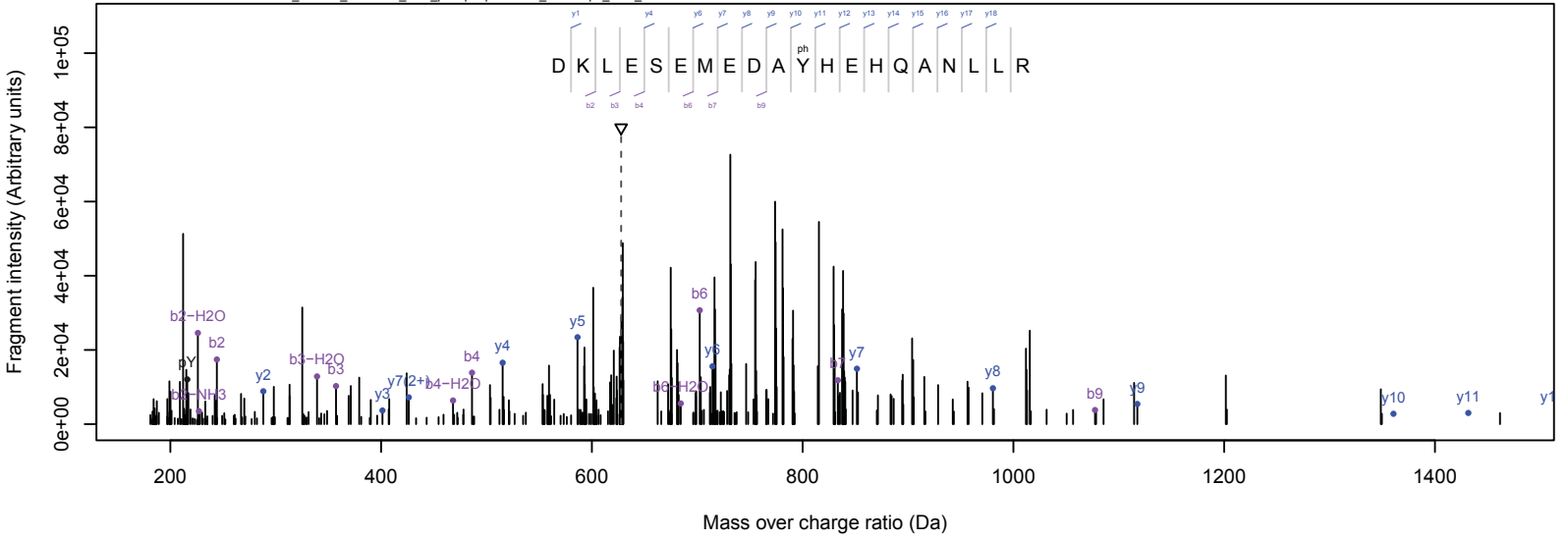
Andromeda score : 83.404



DKLESEMEDAY<sup>ph</sup>HEHQANLLR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26890

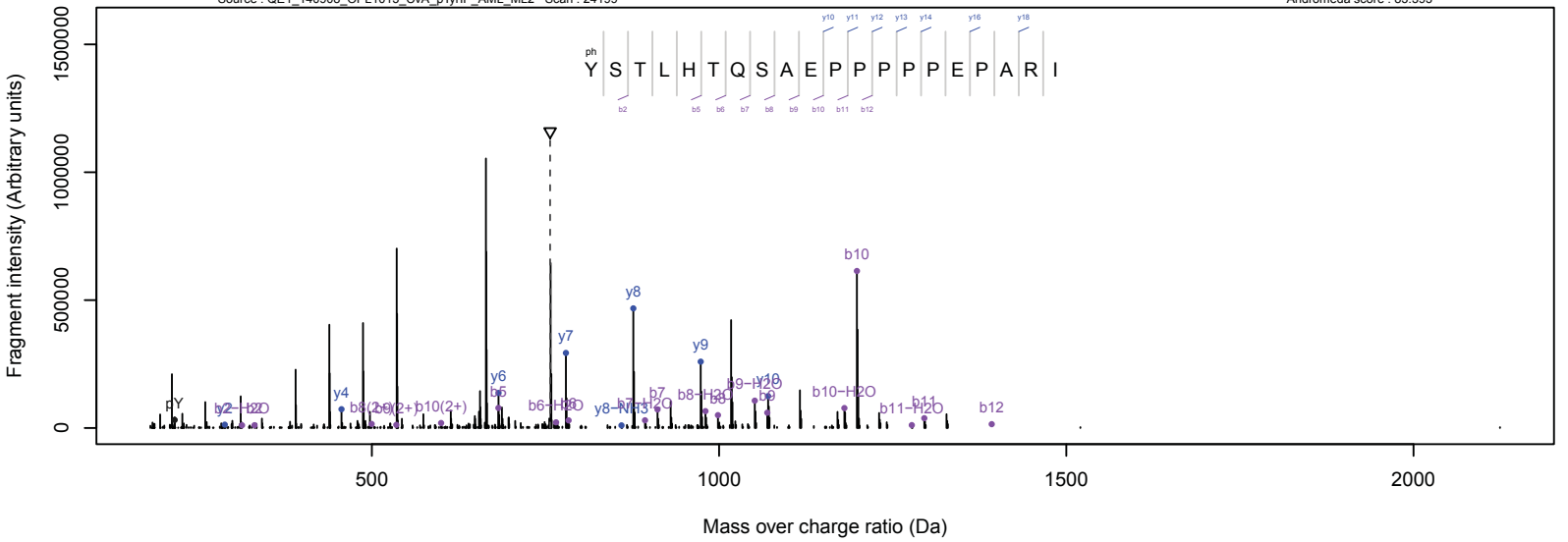
Andromeda score : 83.395



Y<sup>ph</sup>STLHTQSAEPPPPPEPARI-3+

Source : QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_ML2 Scan : 24199

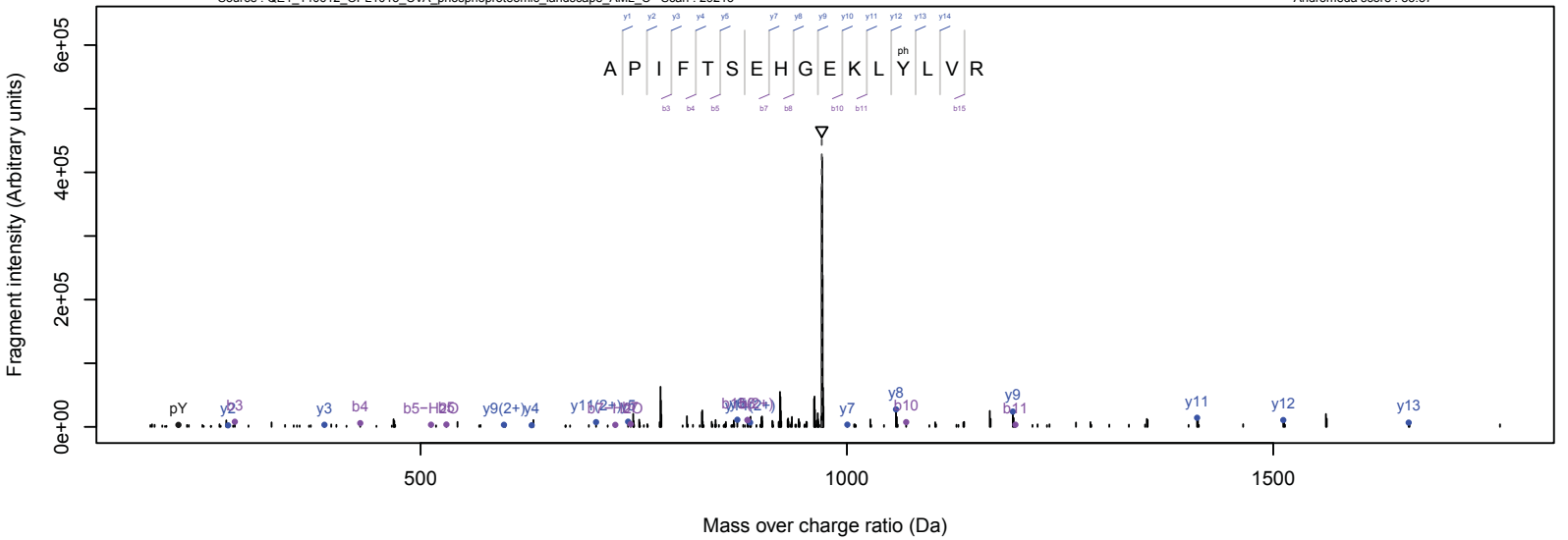
Andromeda score : 83.393



APIFTSEHGKLY<sup>ph</sup>LVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29215

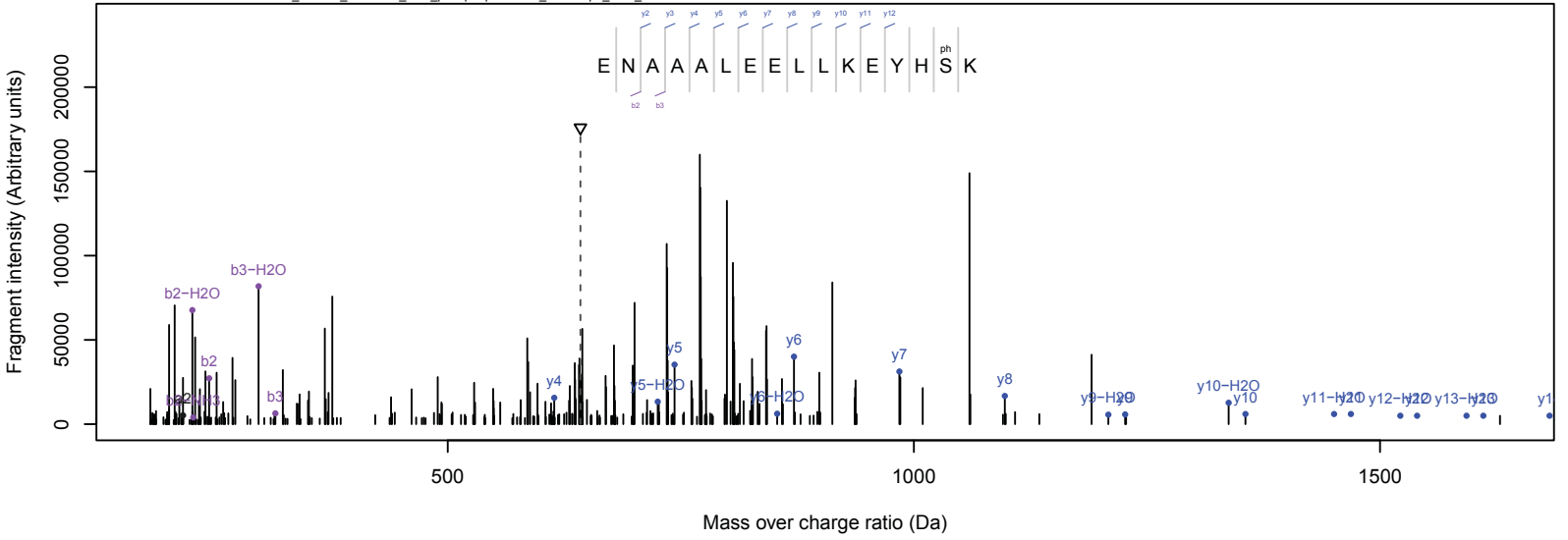
Andromeda score : 83.37



**ENAAALEELLKEYHS(ph)K-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30884

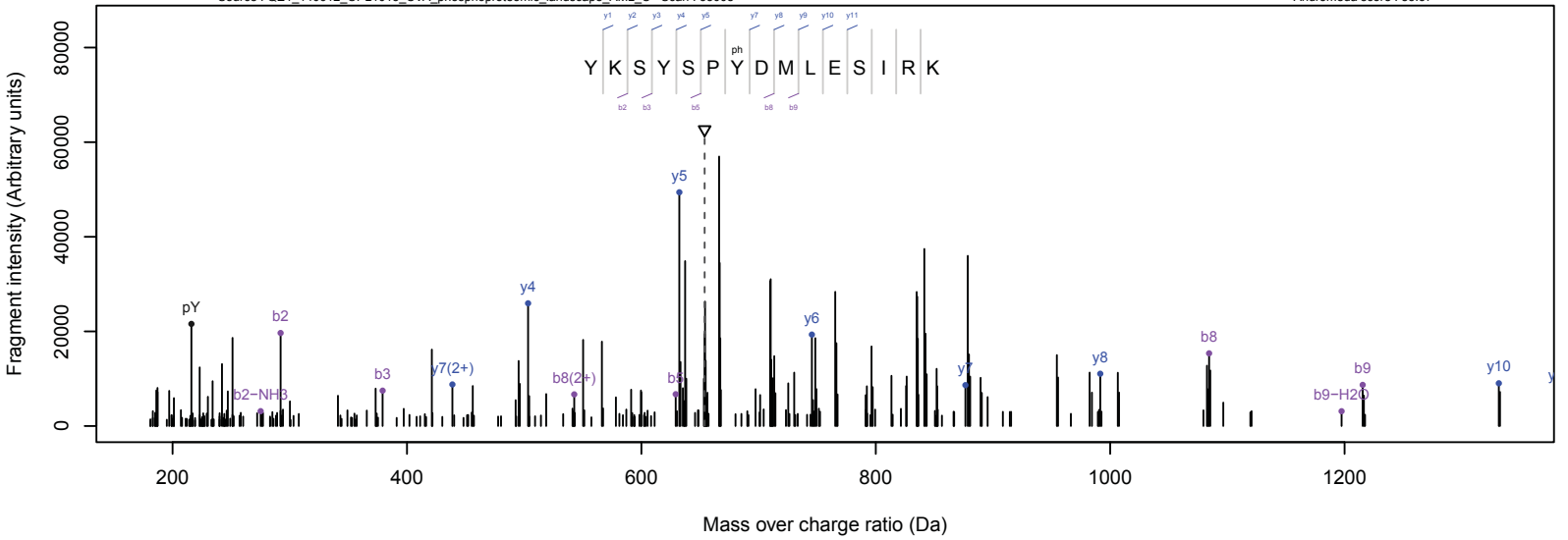
Andromeda score : 83.37



**YKSYSPY(ph)DMLESIRK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33008

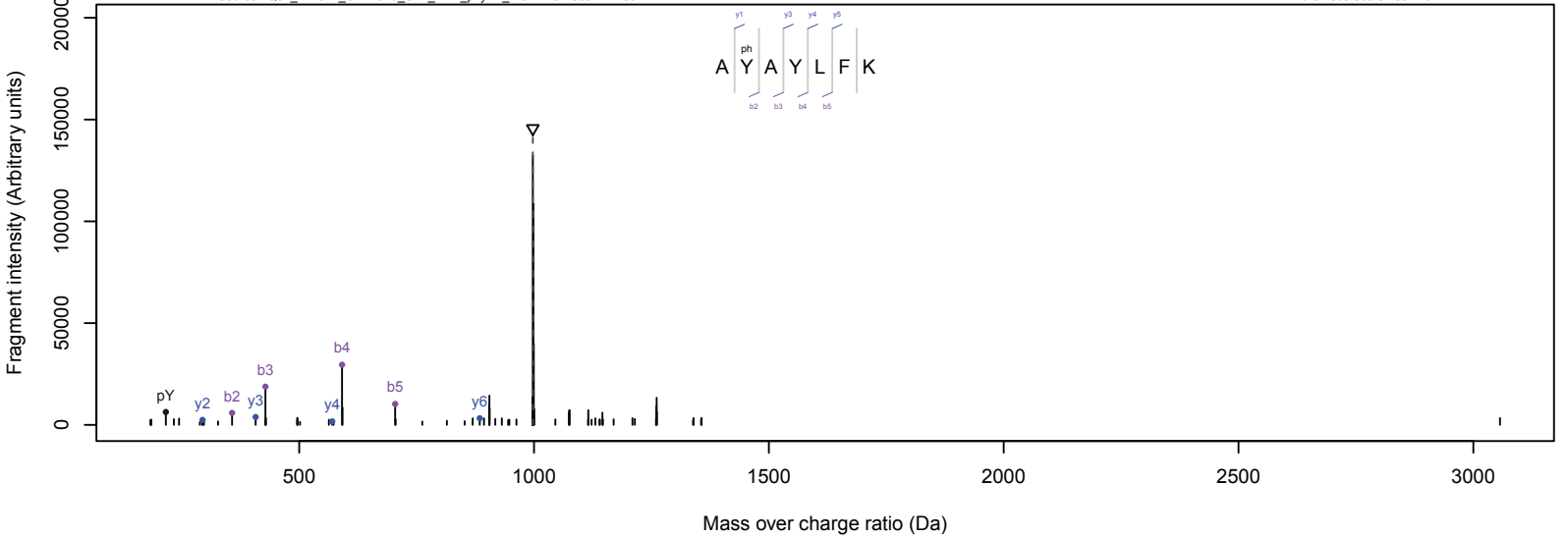
Andromeda score : 83.37



**(ac)AY(ph)AYLFK-1+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 41034

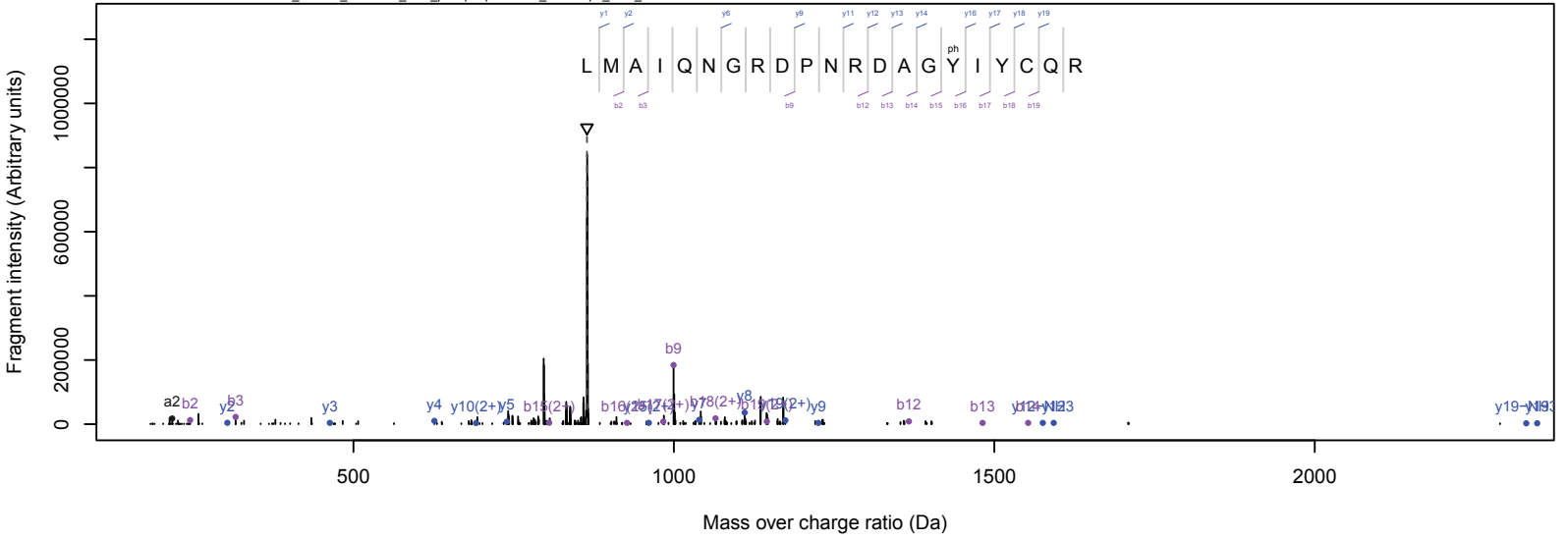
Andromeda score : 83.229



**LMAIQNGRDPNRYIYQQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22337

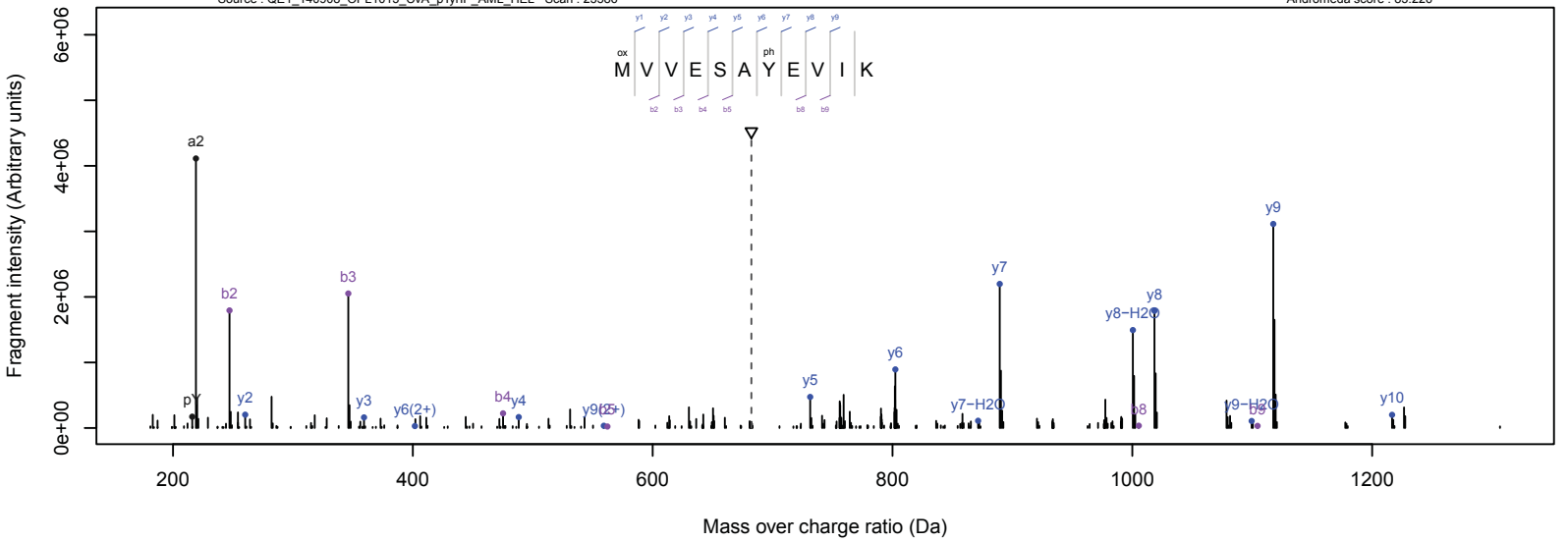
Andromeda score : 83.229



**M(ox)VVESAY(ph)EVIK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23586

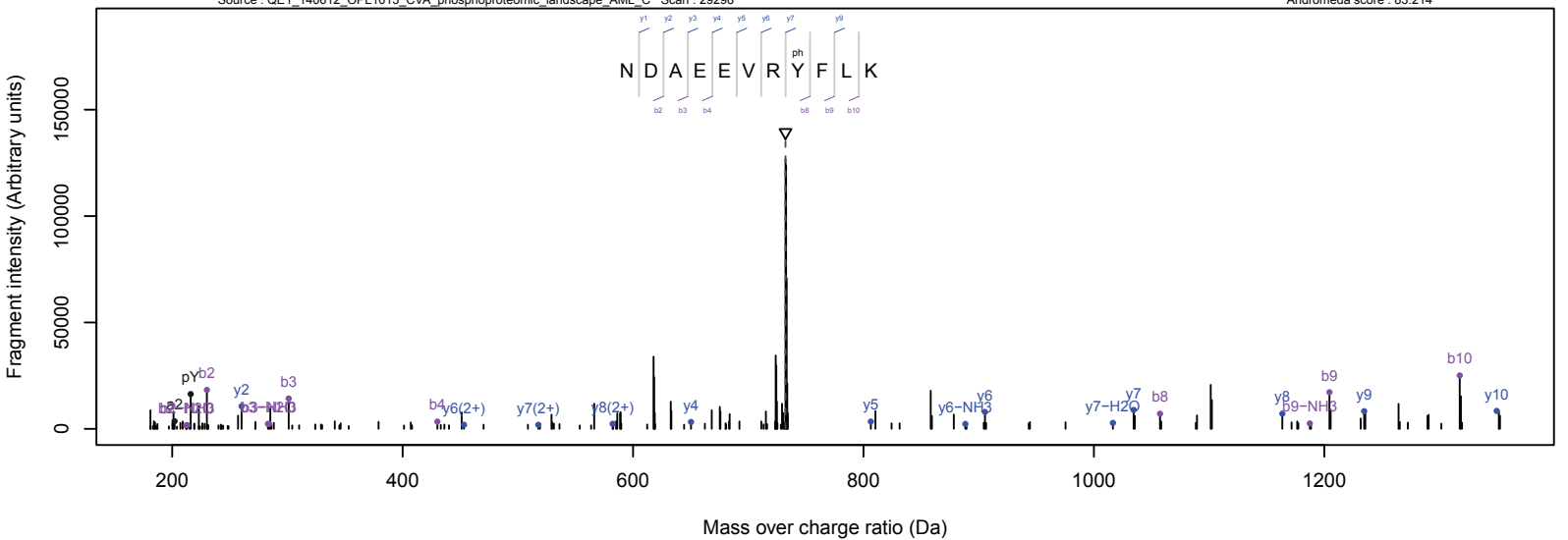
Andromeda score : 83.226



**NDAEEVRY(ph)FLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29298

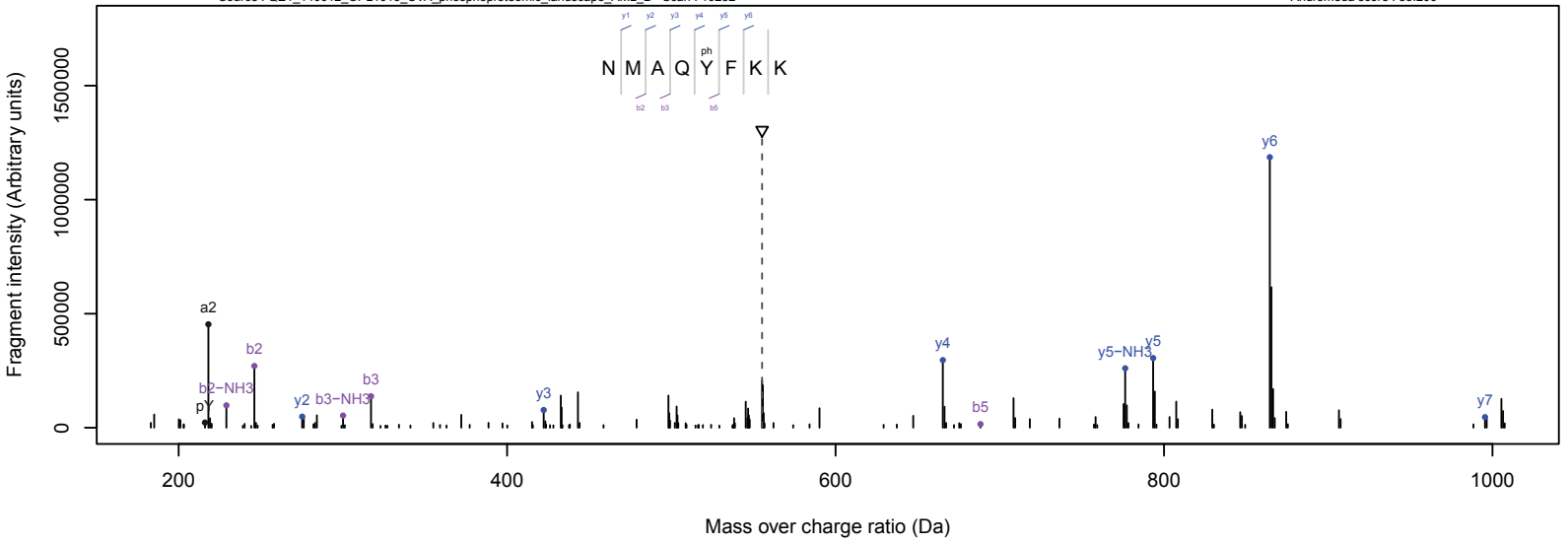
Andromeda score : 83.214



NMAQY(ph)FKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10252

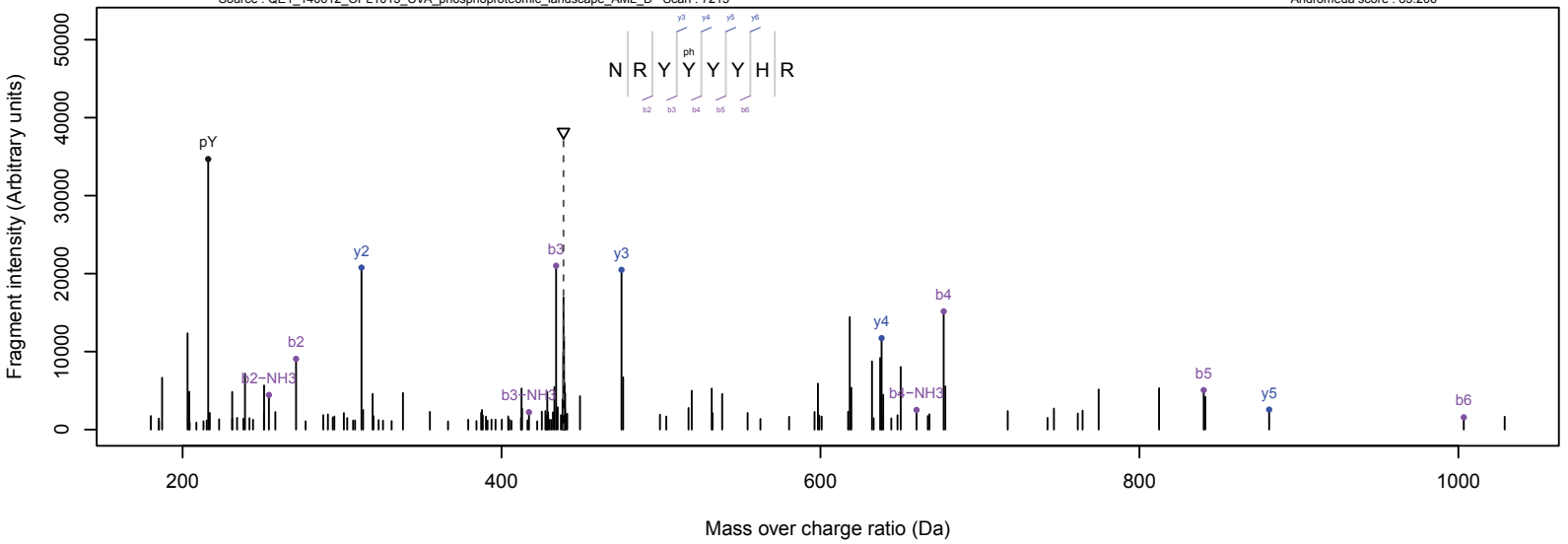
Andromeda score : 83.206



NRYY(ph)YYHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 7215

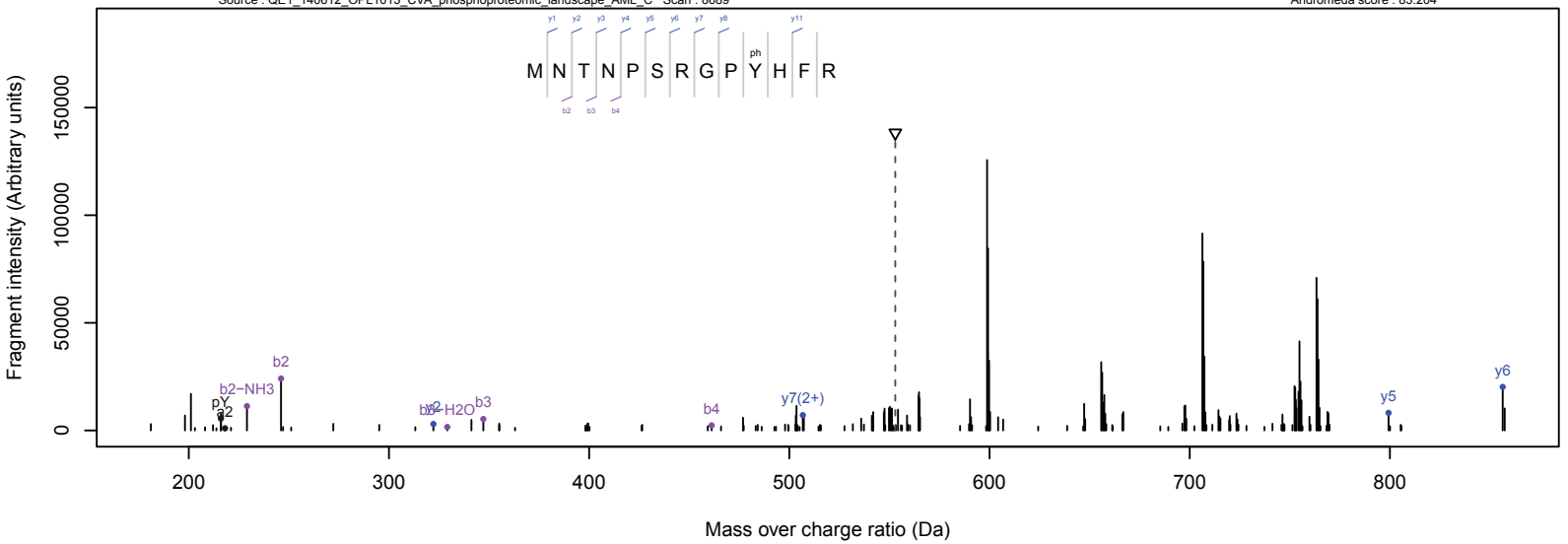
Andromeda score : 83.206



MNTNPSRGPY(ph)HFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8689

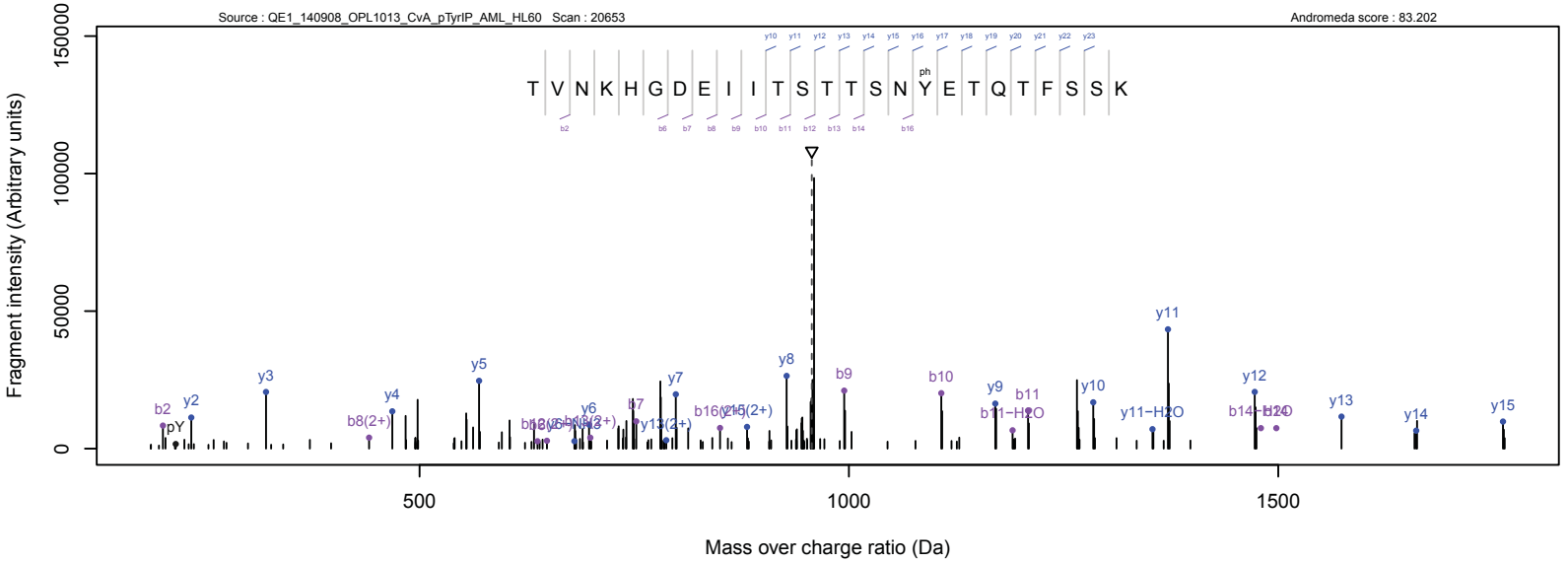
Andromeda score : 83.204



TVNKHGDEIITSTTSNY(ph)ETQTFSSK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan : 20653

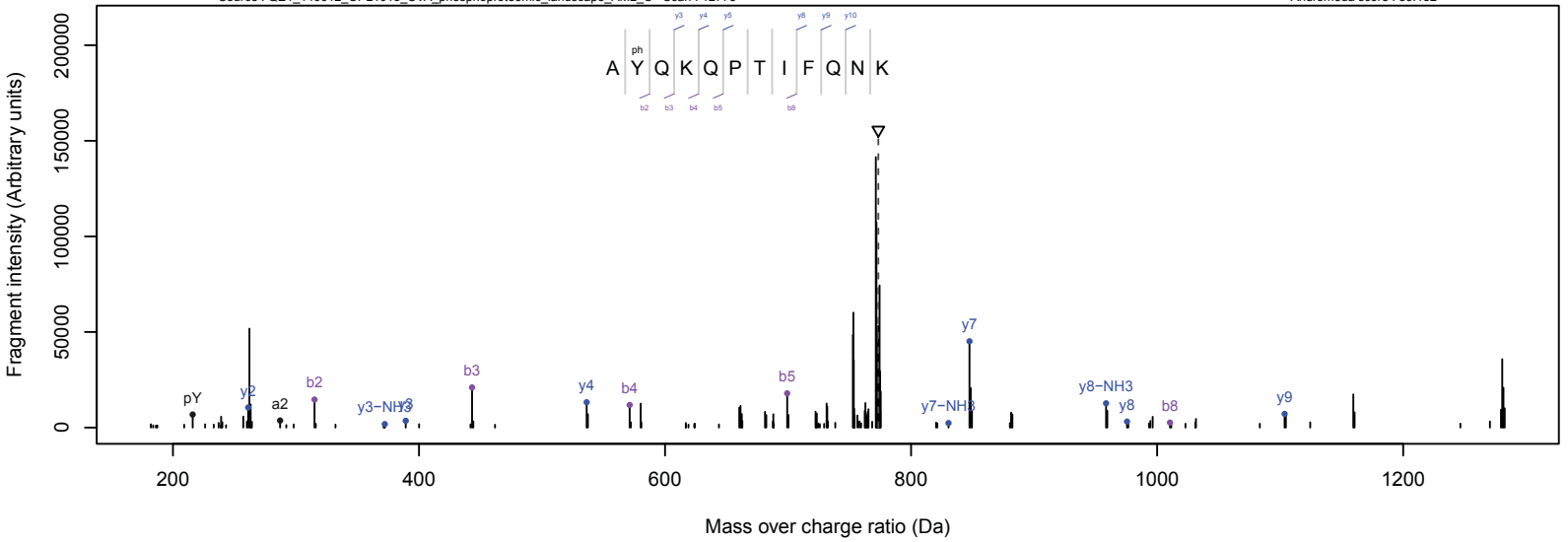
Andromeda score : 83.202



AY(ph)QKQPTIFQNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12778

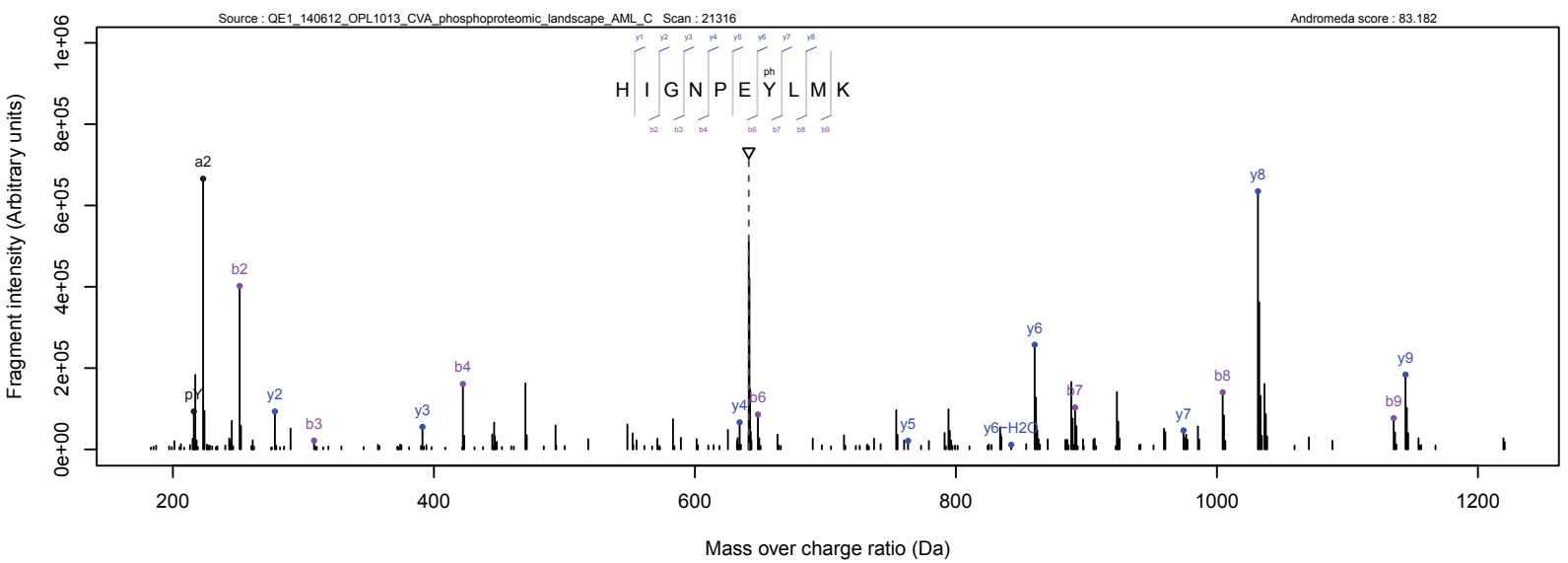
Andromeda score : 83.182



HIGNPEY(ph)LMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21316

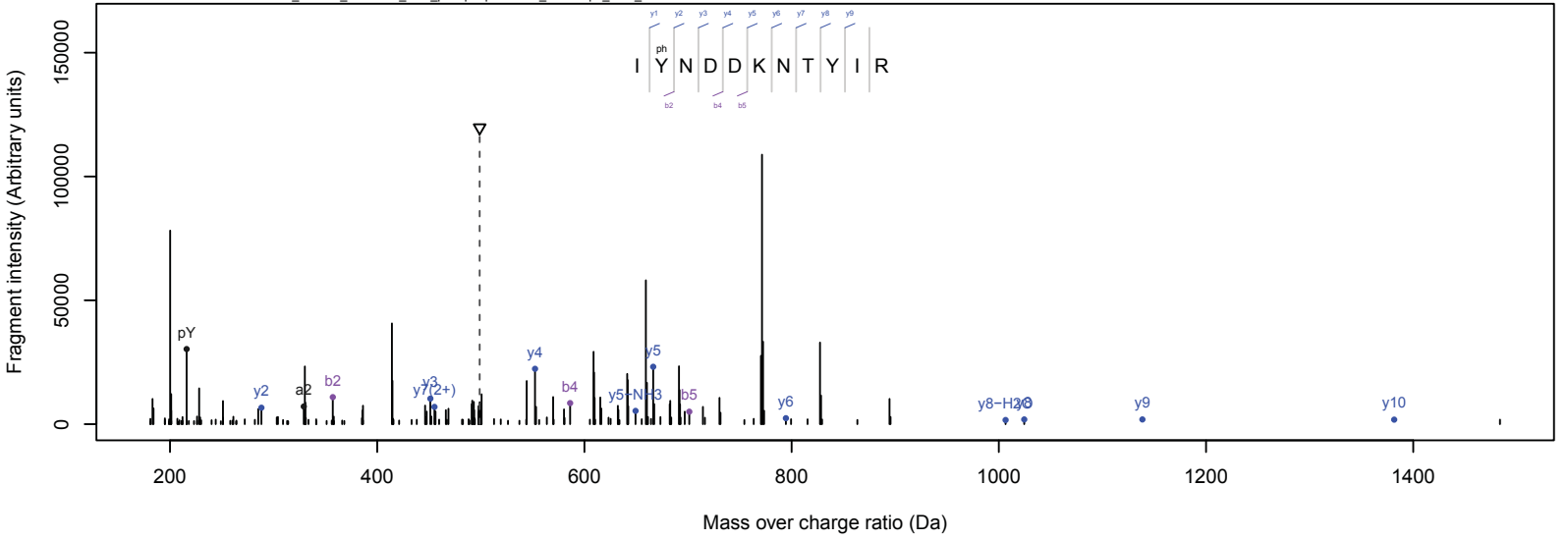
Andromeda score : 83.182



IY(ph)NDDKNTYIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9818

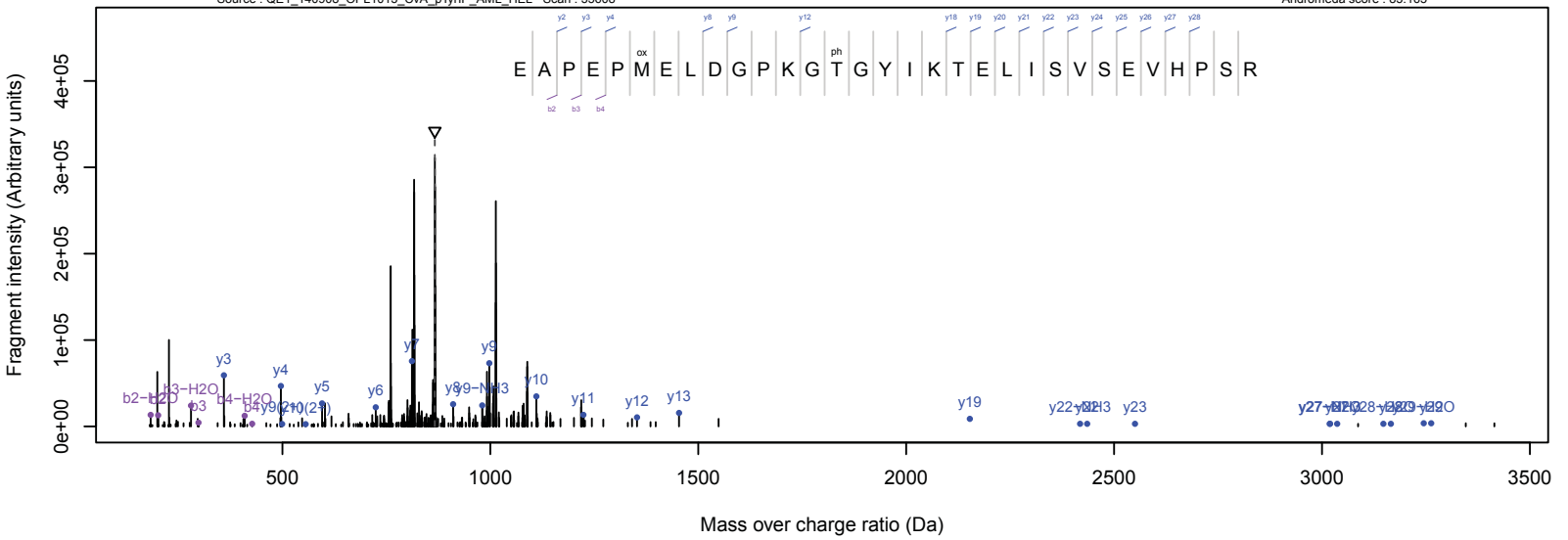
Andromeda score : 83.182



EAPEPM(ox)ELDGPKG(ph)GYIKTELISVSEVHPSR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35608

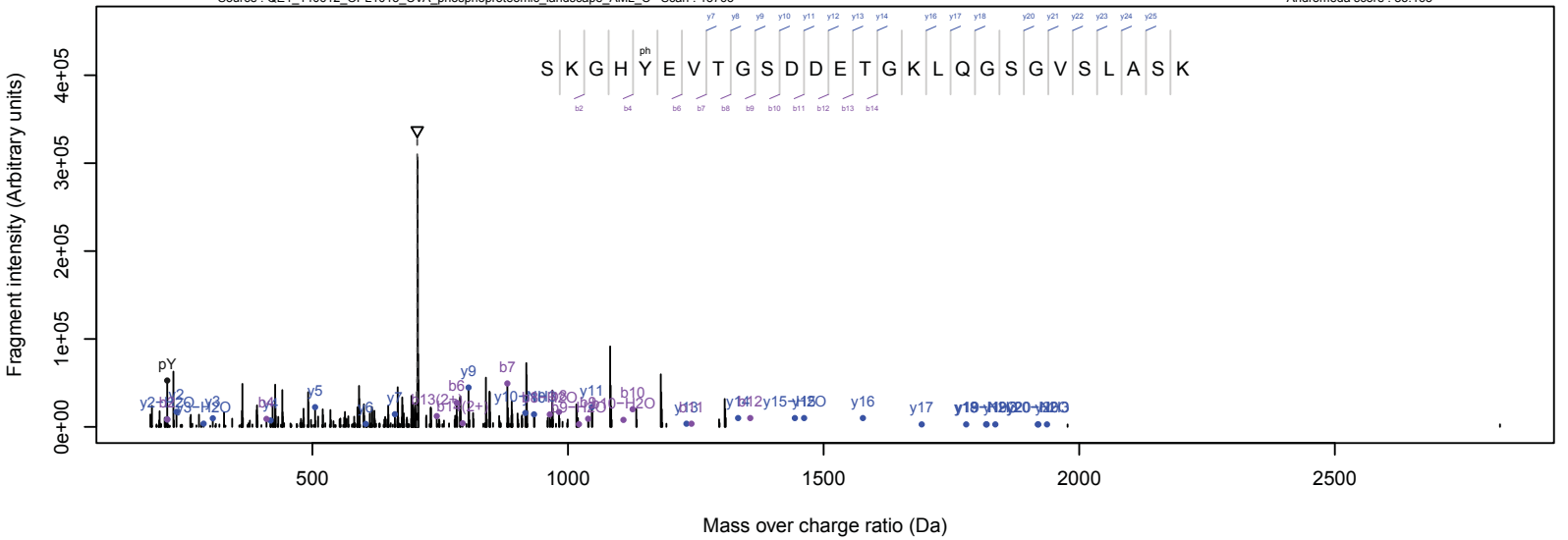
Andromeda score : 83.165



SKGHY(ph)EVTGSDDDETGKLGSGVSLASK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15763

Andromeda score : 83.165

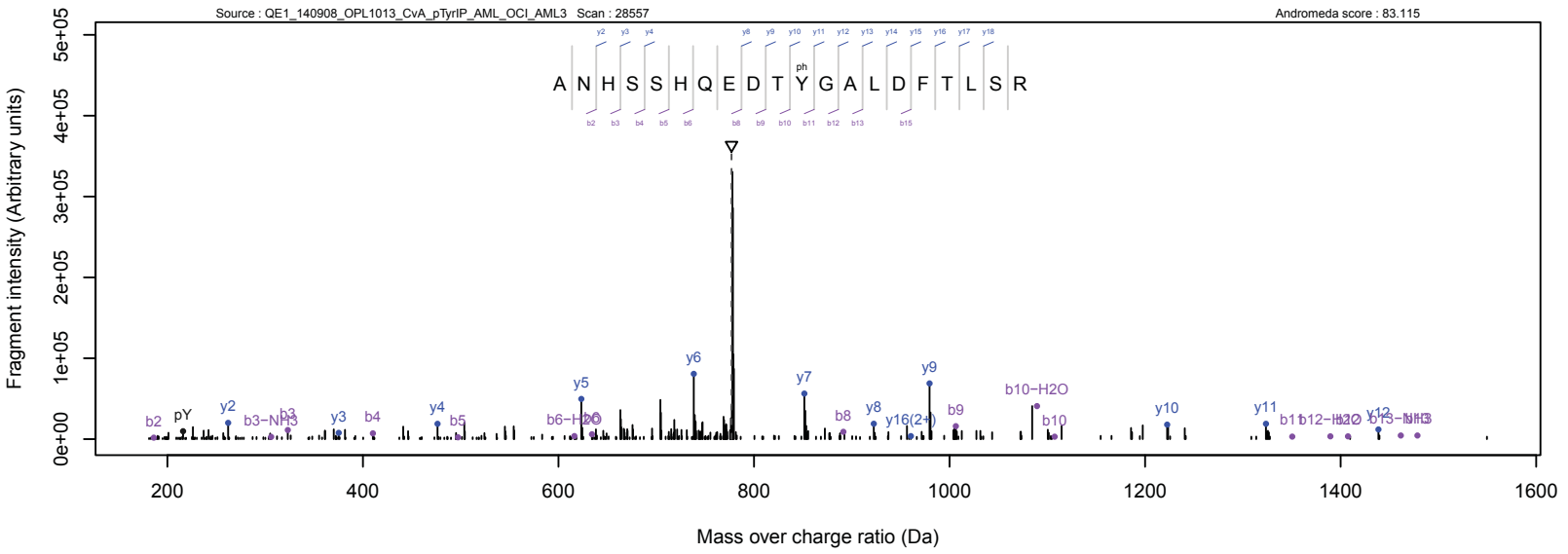




ANHSSHQEDTY(ph)GALDFTLR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 28557

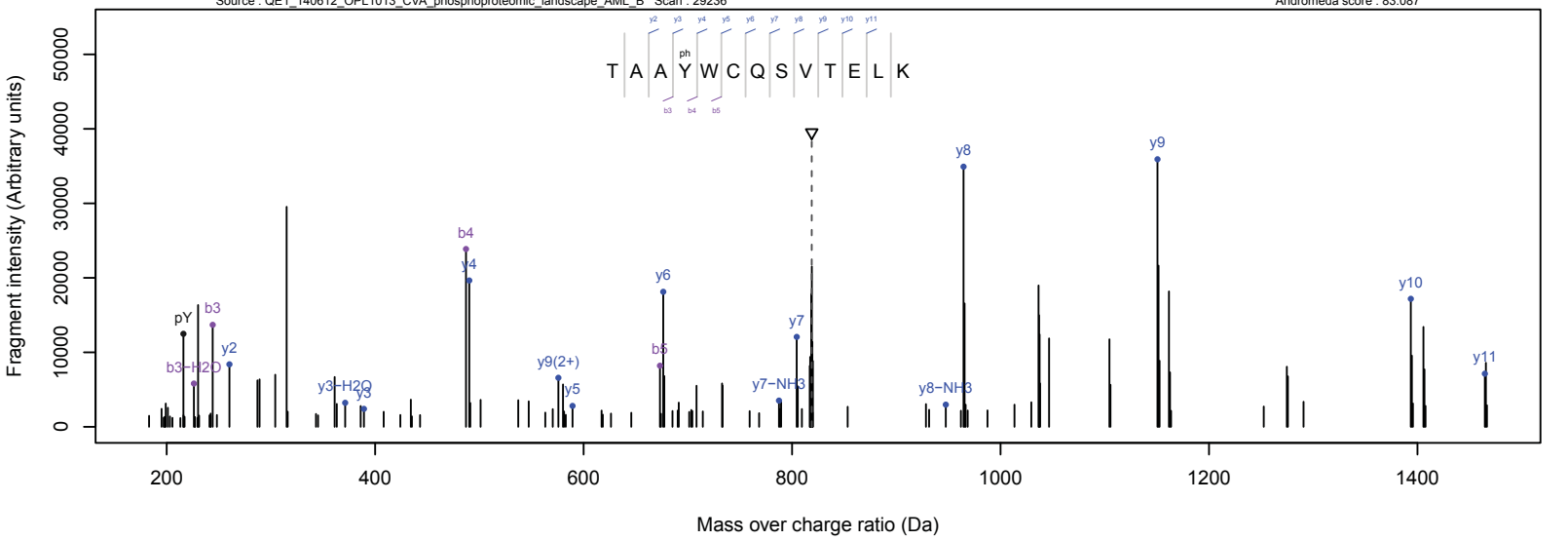
Andromeda score: 83.115



TAAY(ph)WCQSVTELK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 29236

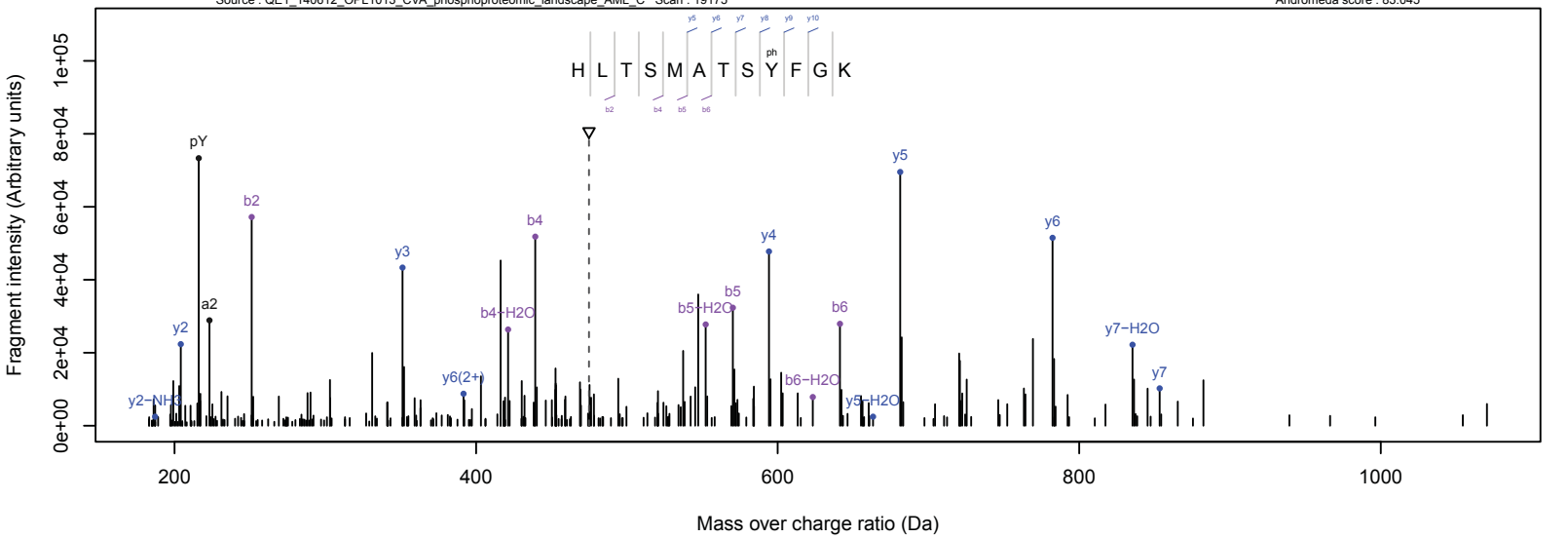
Andromeda score: 83.087



HLTSMATSY(ph)FGK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19175

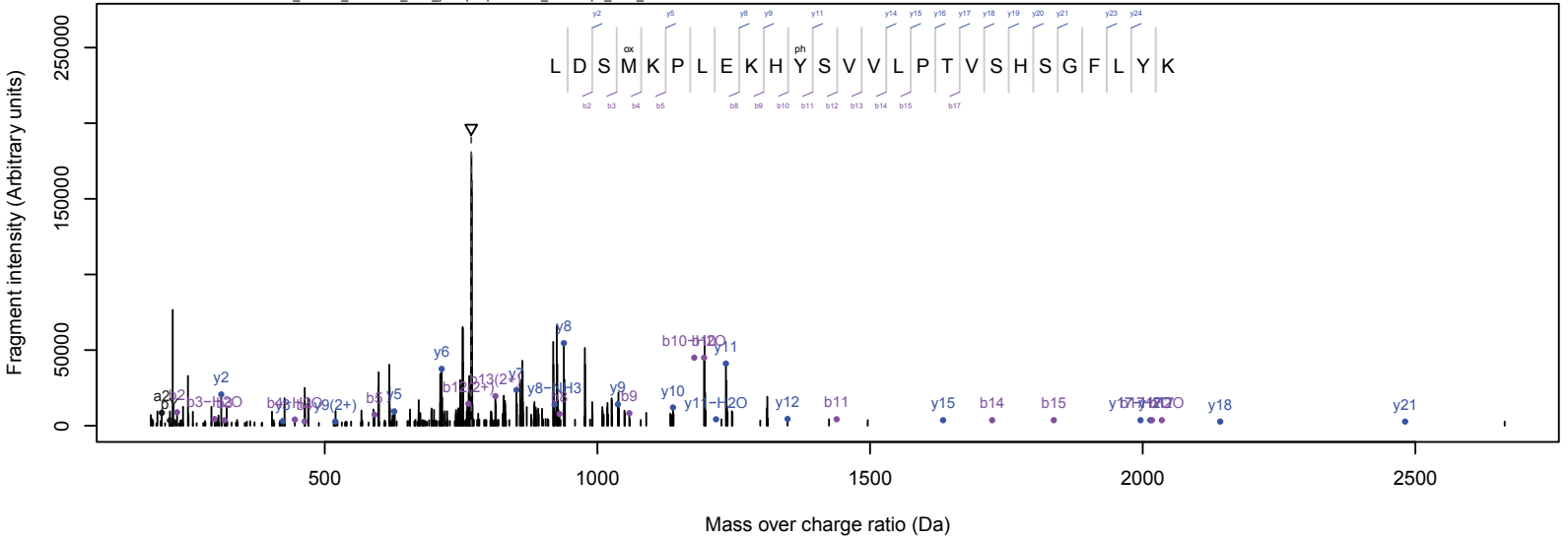
Andromeda score: 83.045



LDSM(ox)KPLEKHY(ph)SVVLPVSHSGFLYK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 31436

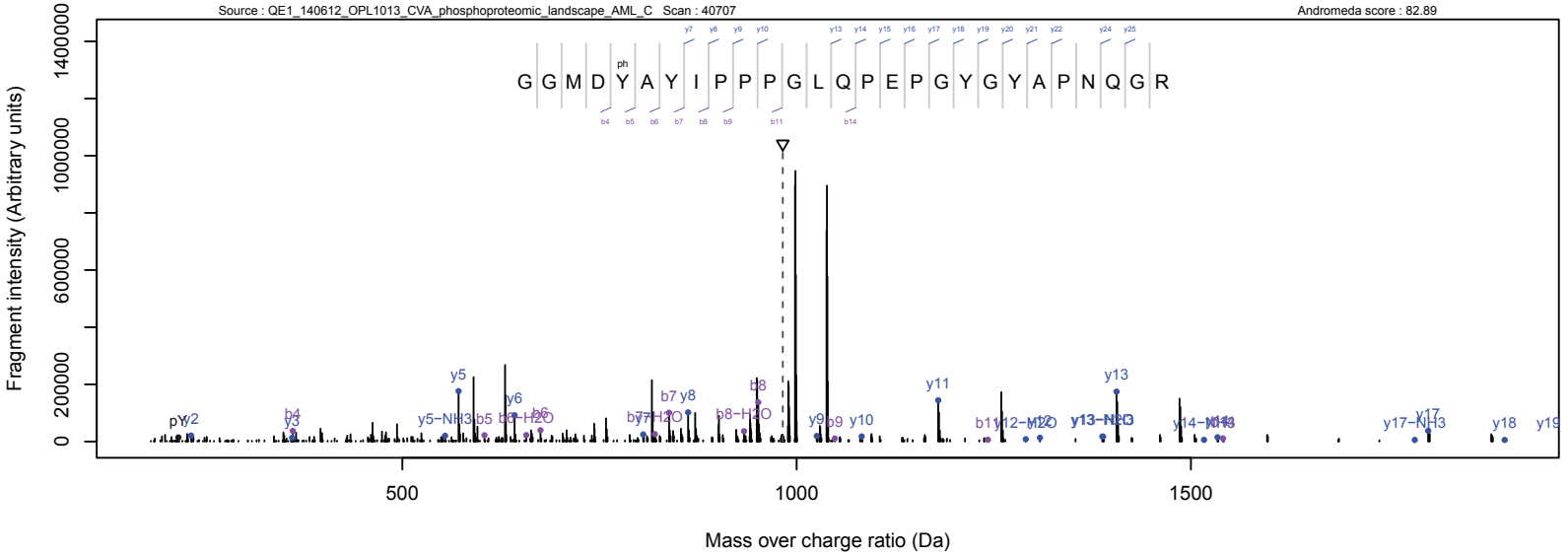
Andromeda score: 82.942



GGMDY(ph)AYIPPPGLQPEPGYAPNQGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 40707

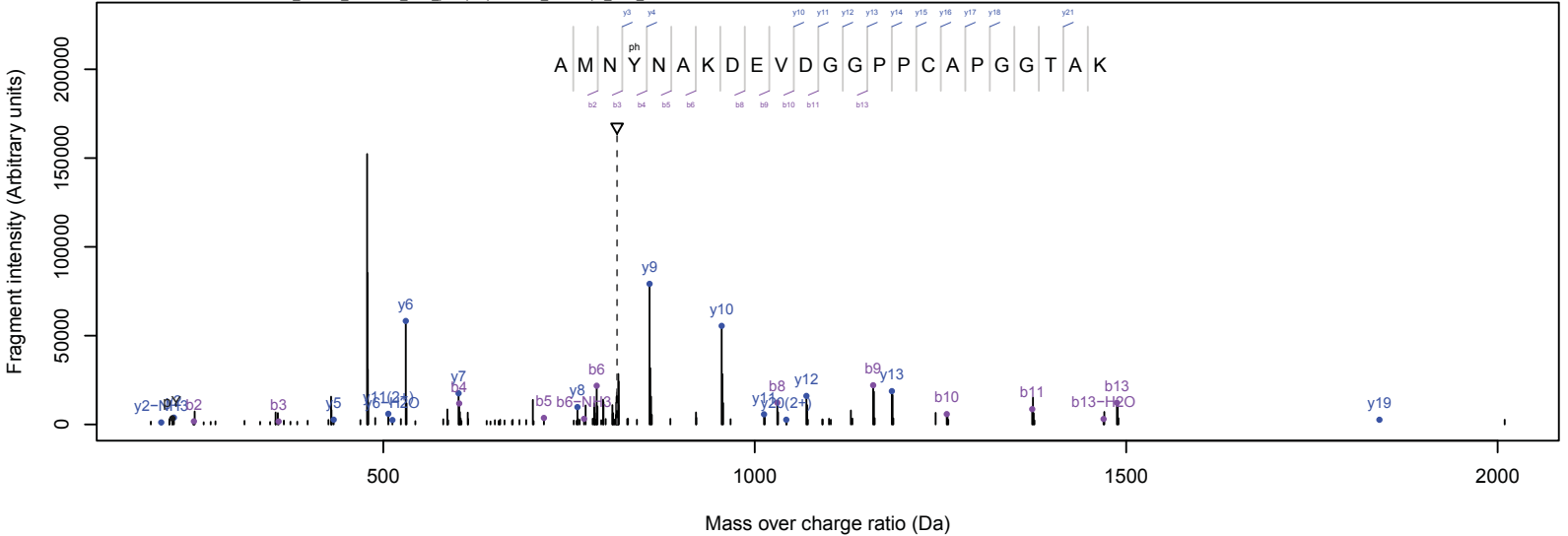
Andromeda score: 82.89



(ac)AMNY(ph)NAKDEVDGGPPCAPGGTAK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19677

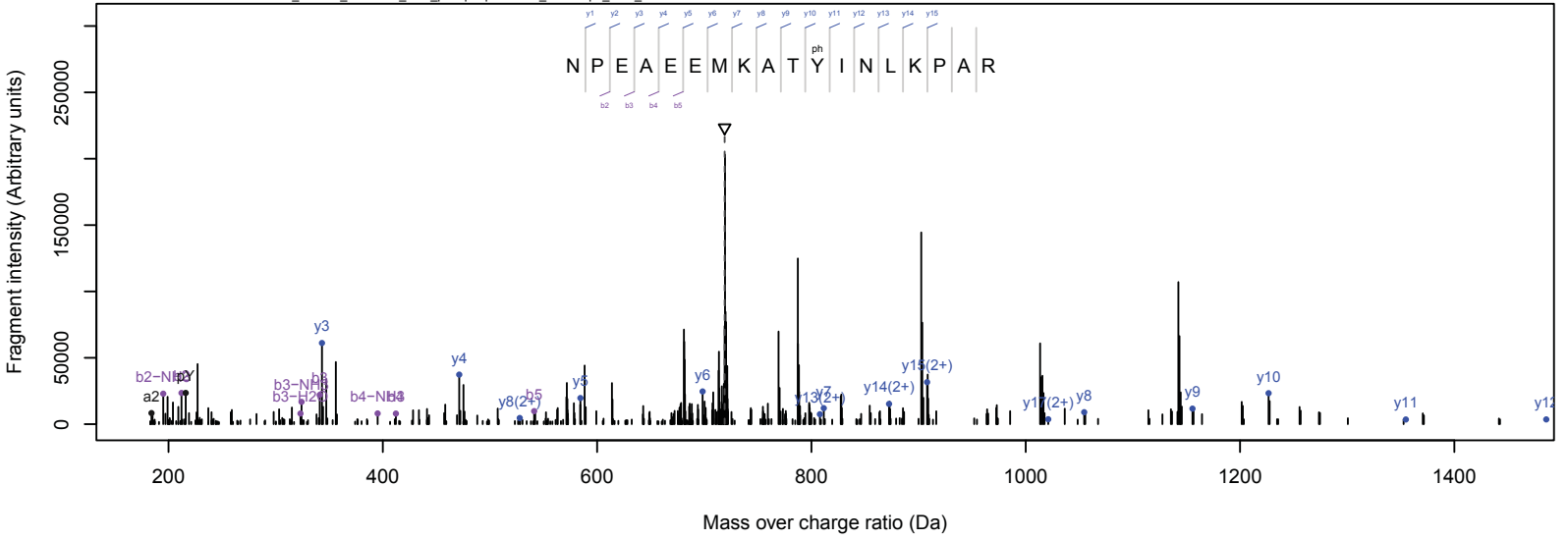
Andromeda score: 82.885



**NPEAEEMKATY(ph)INLKPAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21964

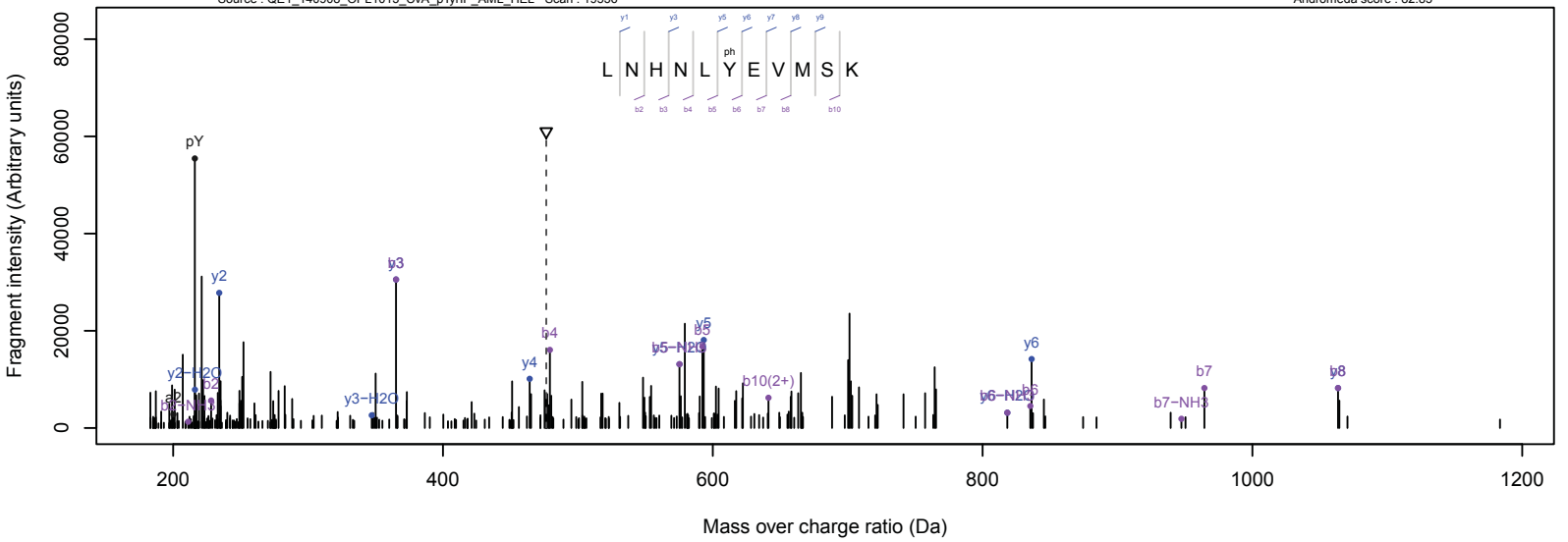
Andromeda score : 82.877



**LNHNLY(ph)EVMSK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 19536

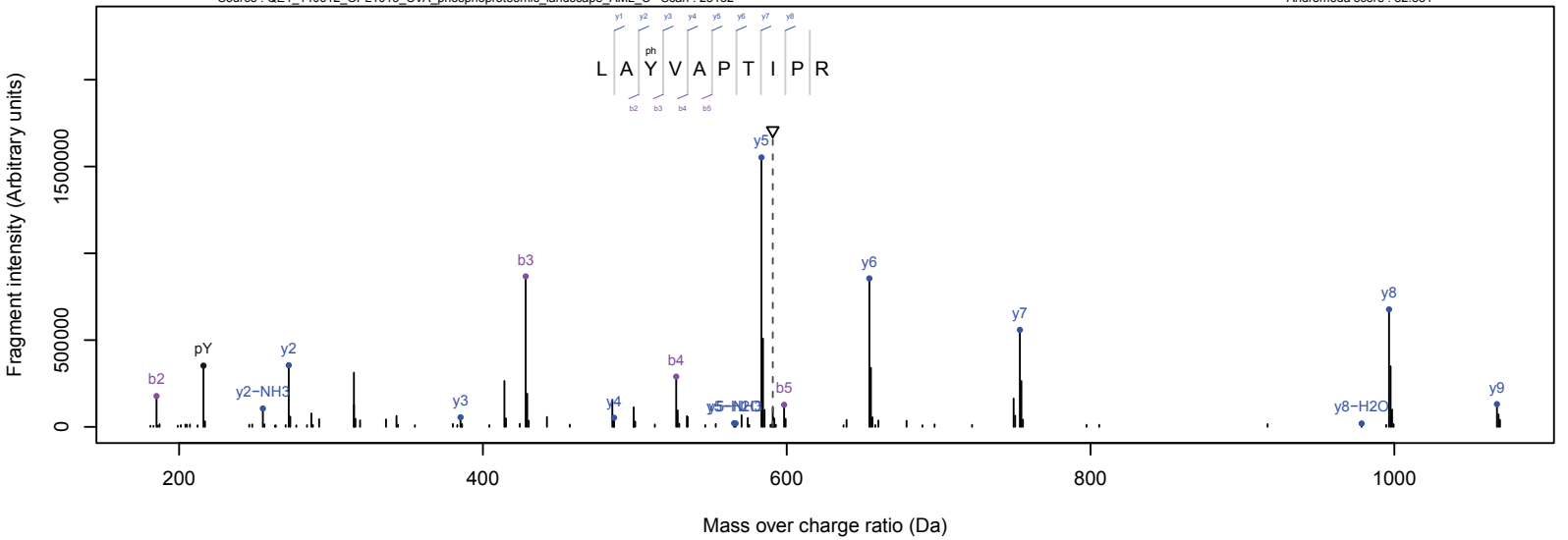
Andromeda score : 82.85



**LAY(ph)VAPTIPR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25182

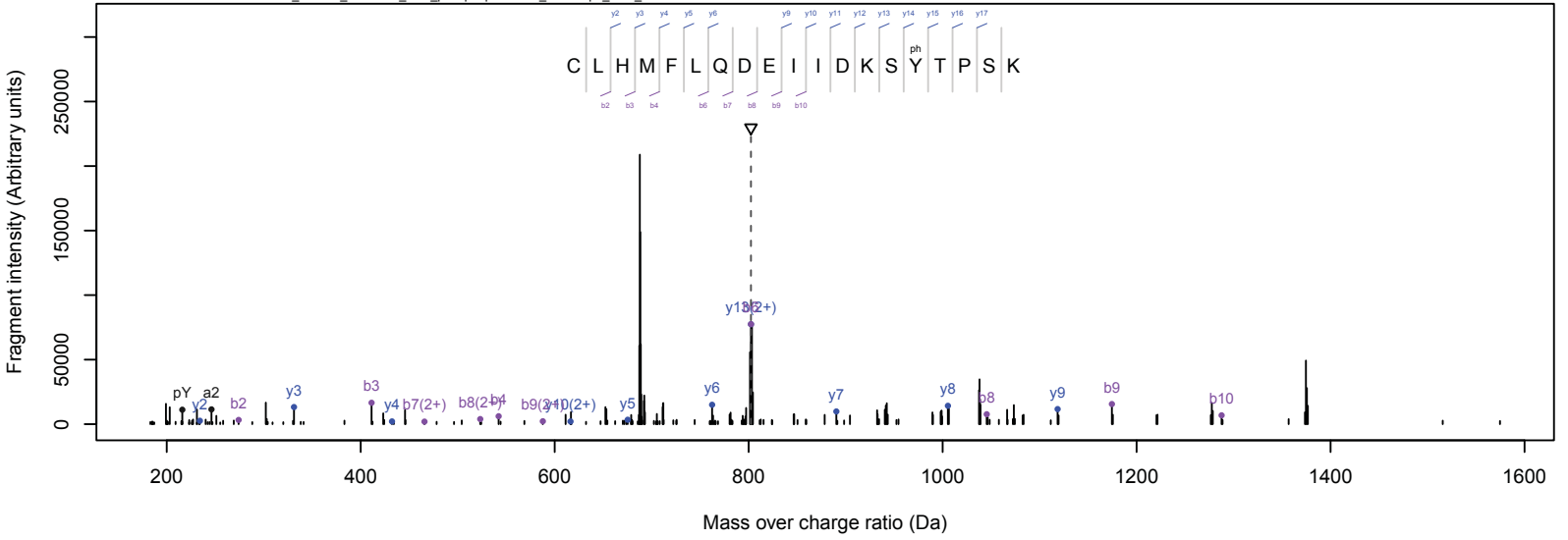
Andromeda score : 82.831



CLHMFLQDEIIDKSY(ph)TPSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34906

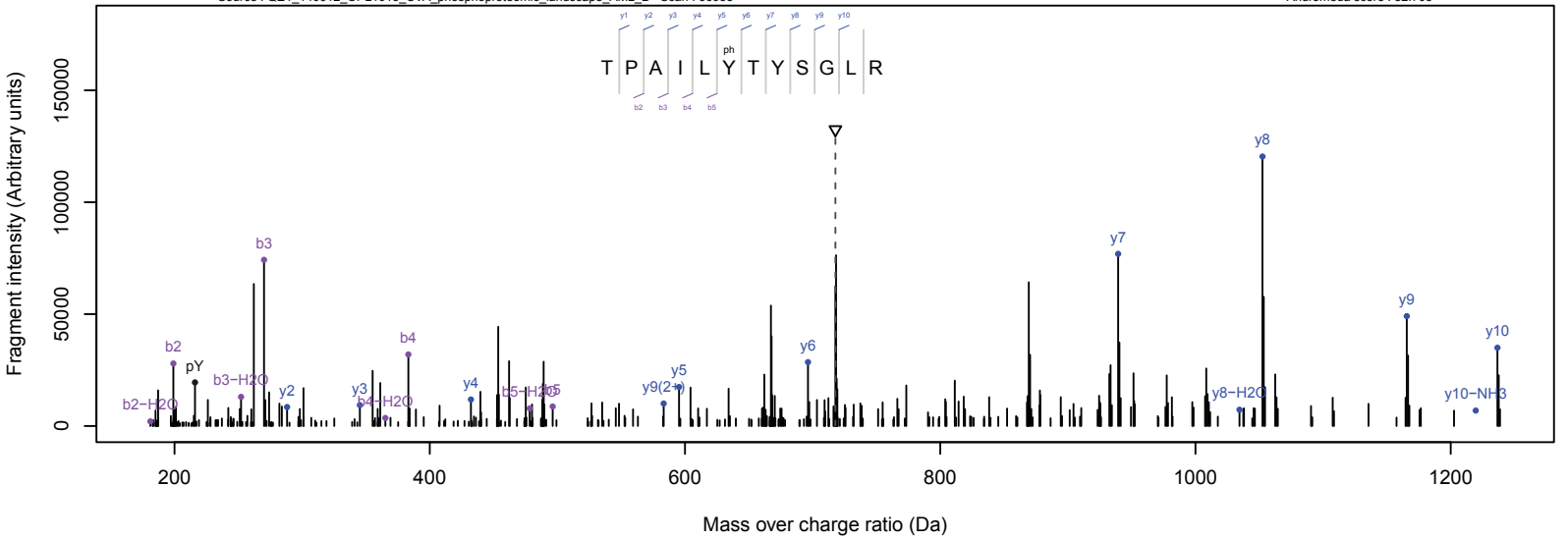
Andromeda score: 82.765



TPAILY(ph)TYSGLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 36958

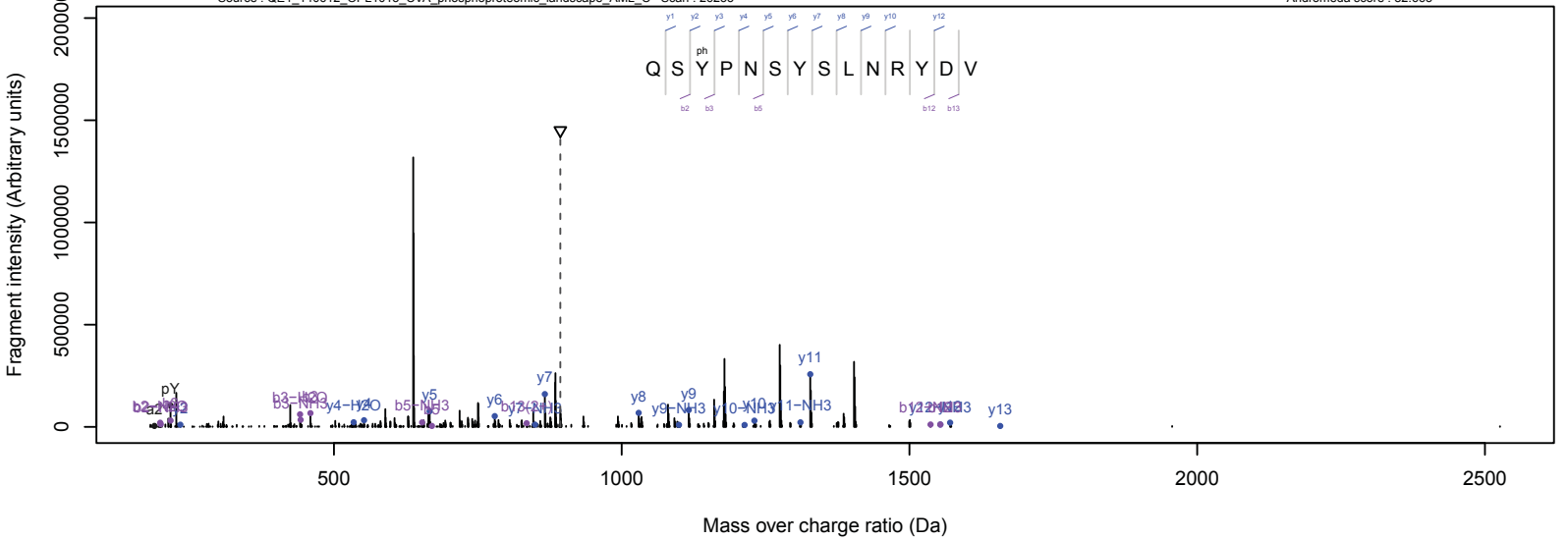
Andromeda score: 82.705



QSY(ph)PNSYSLNRYDV-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26235

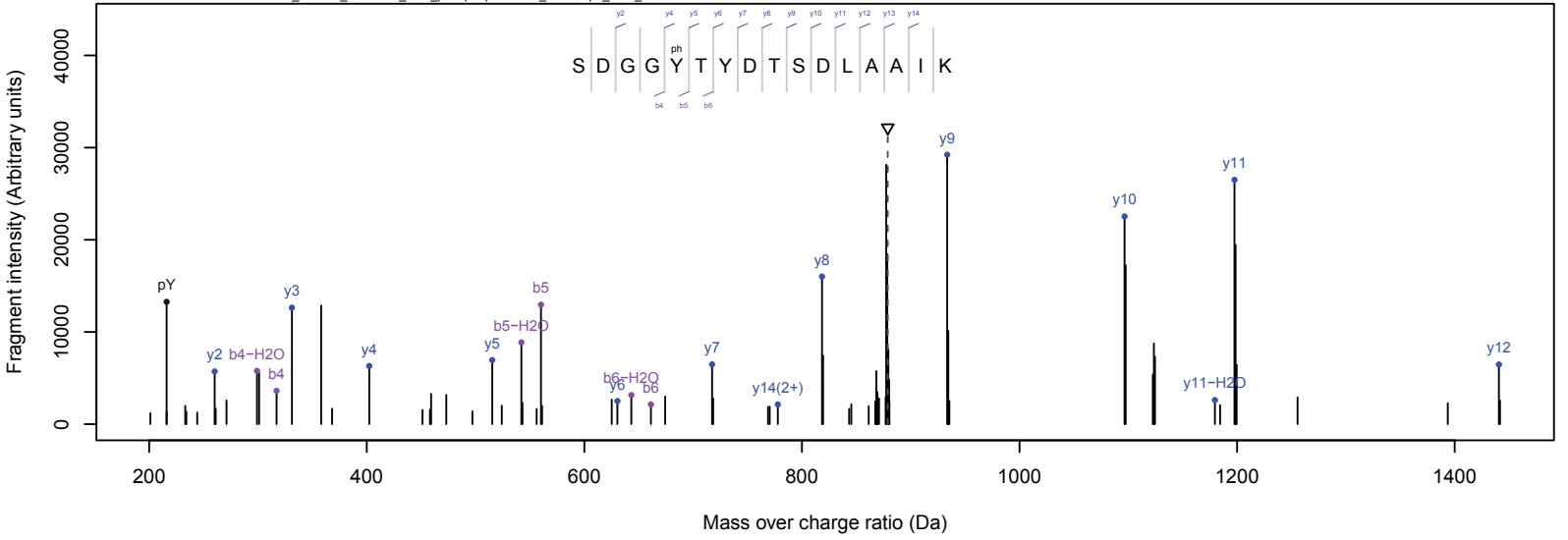
Andromeda score: 82.663



SDGGY(ph)TYDTSDLAAIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27016

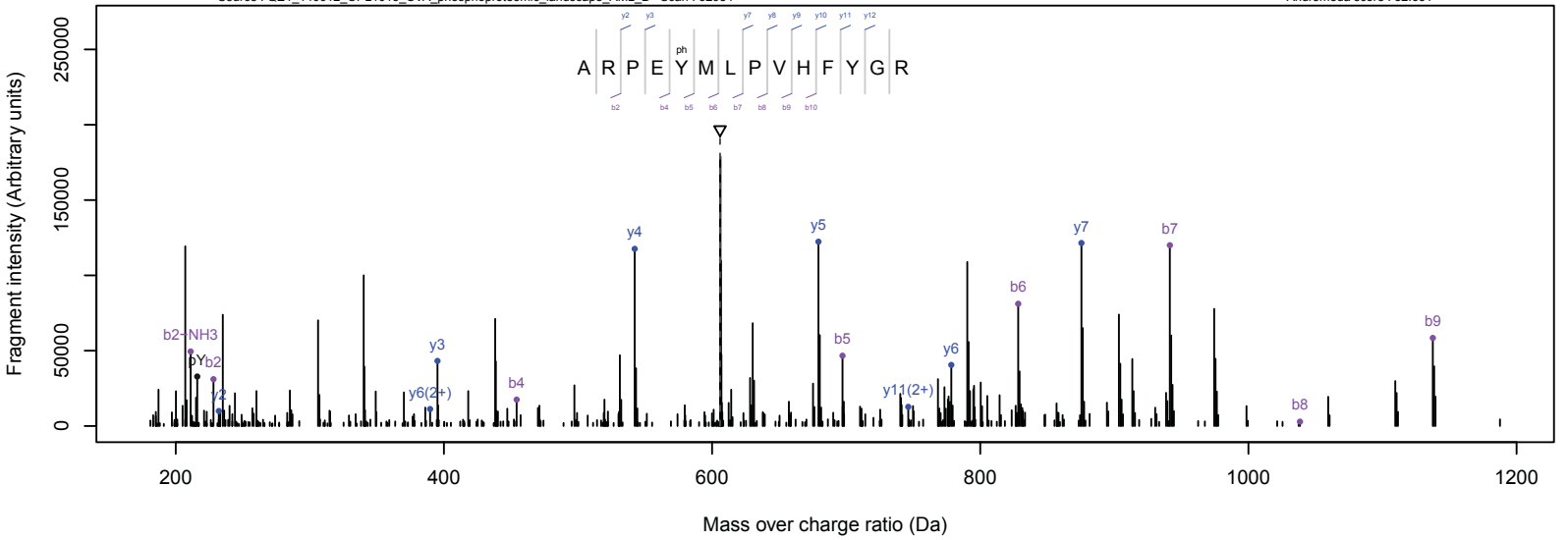
Andromeda score : 82.663



ARPEY(ph)MLPVHFYGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32984

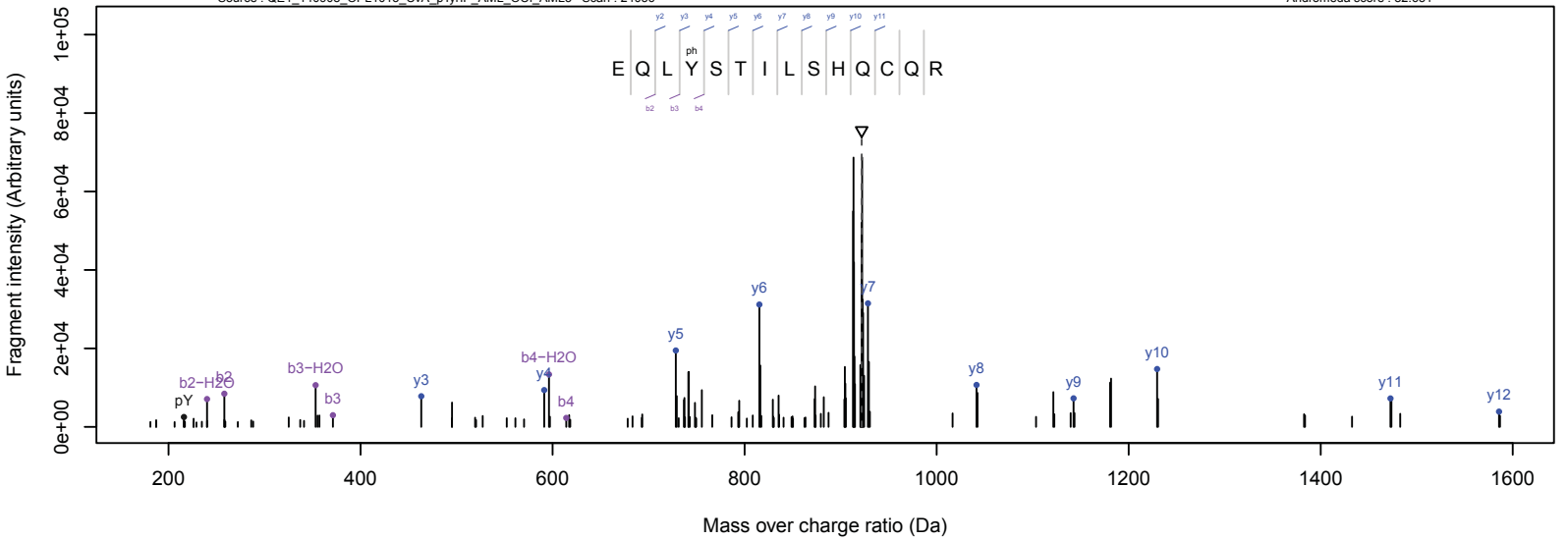
Andromeda score : 82.651



EQLY(ph)STILSHQCQR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 24056

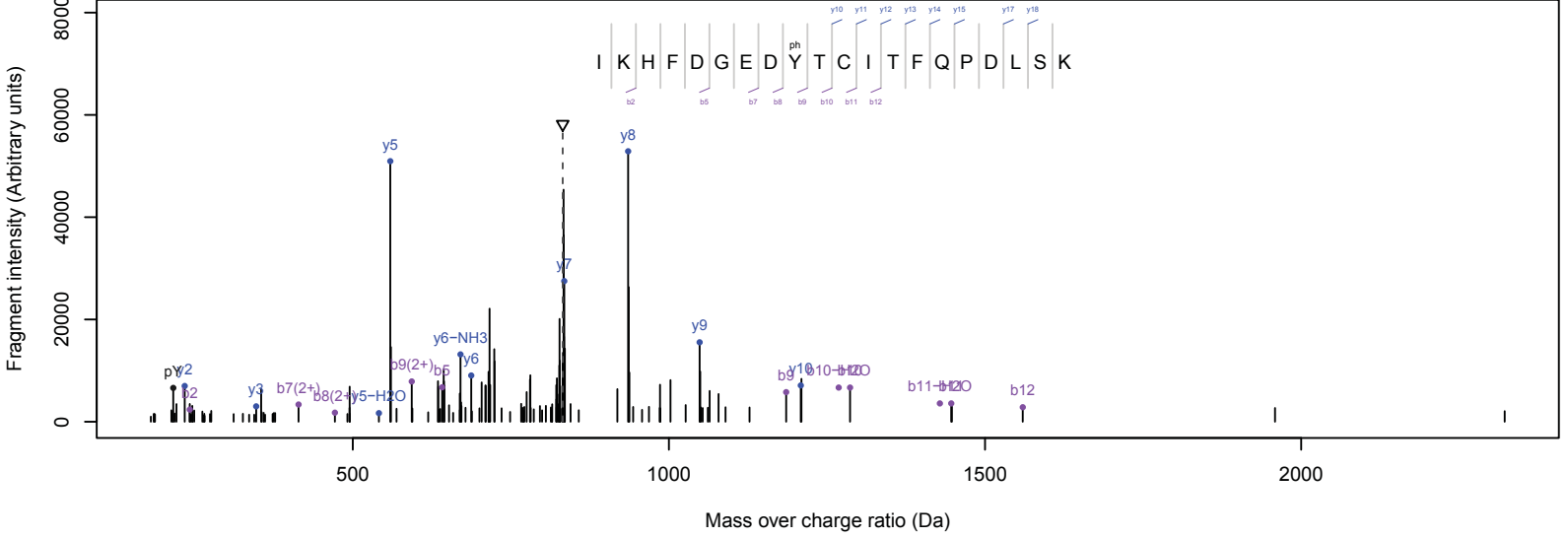
Andromeda score : 82.651



**IKHFDGEDY(ph)TCITFPDLSK-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 30510

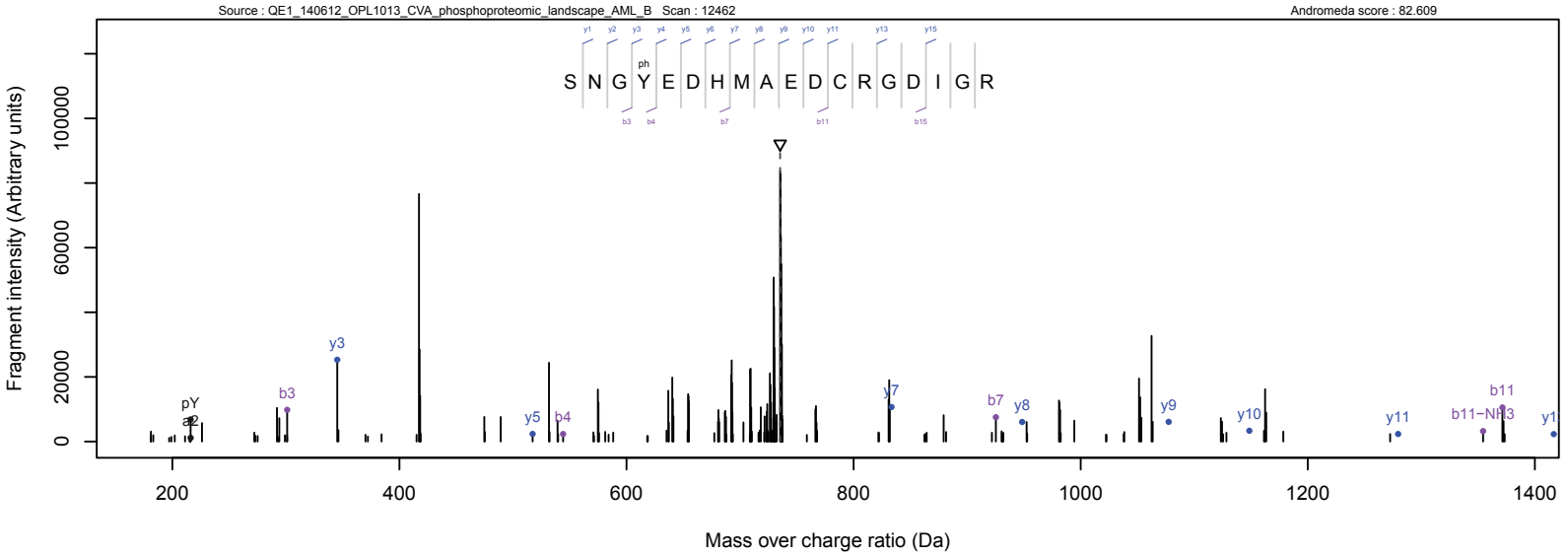
Andromeda score: 82.609



**(ac)SNGY(ph)EDHMAEDCRGDIGR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 12462

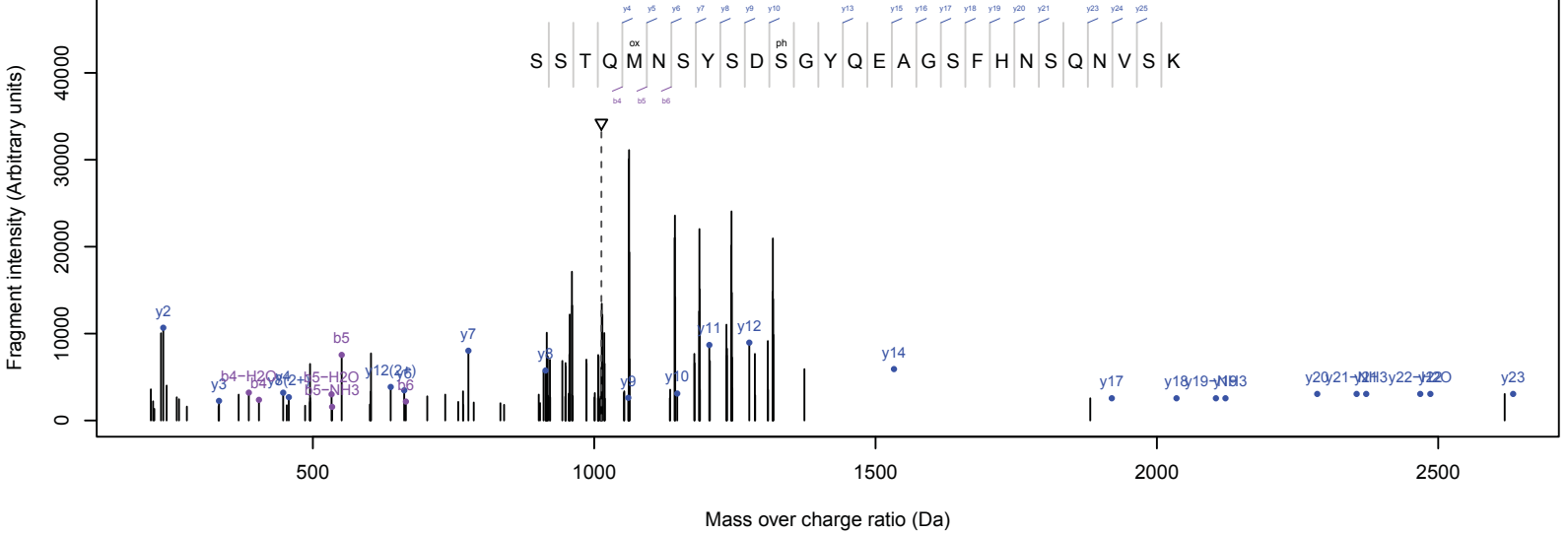
Andromeda score: 82.609



**SSTQM(ax)NSYSDS(ph)GYQEAGSFHNSQNVSK-3+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 14752

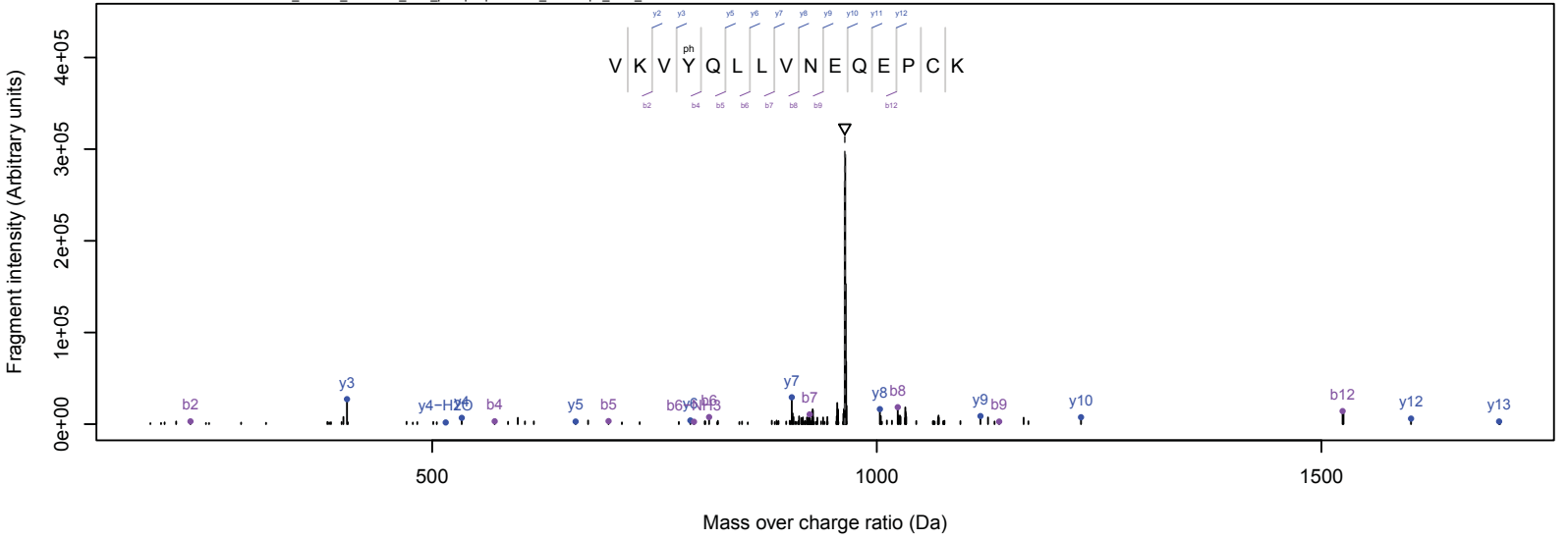
Andromeda score: 82.6



VKVY(ph)QLLVNEQEPCK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27058

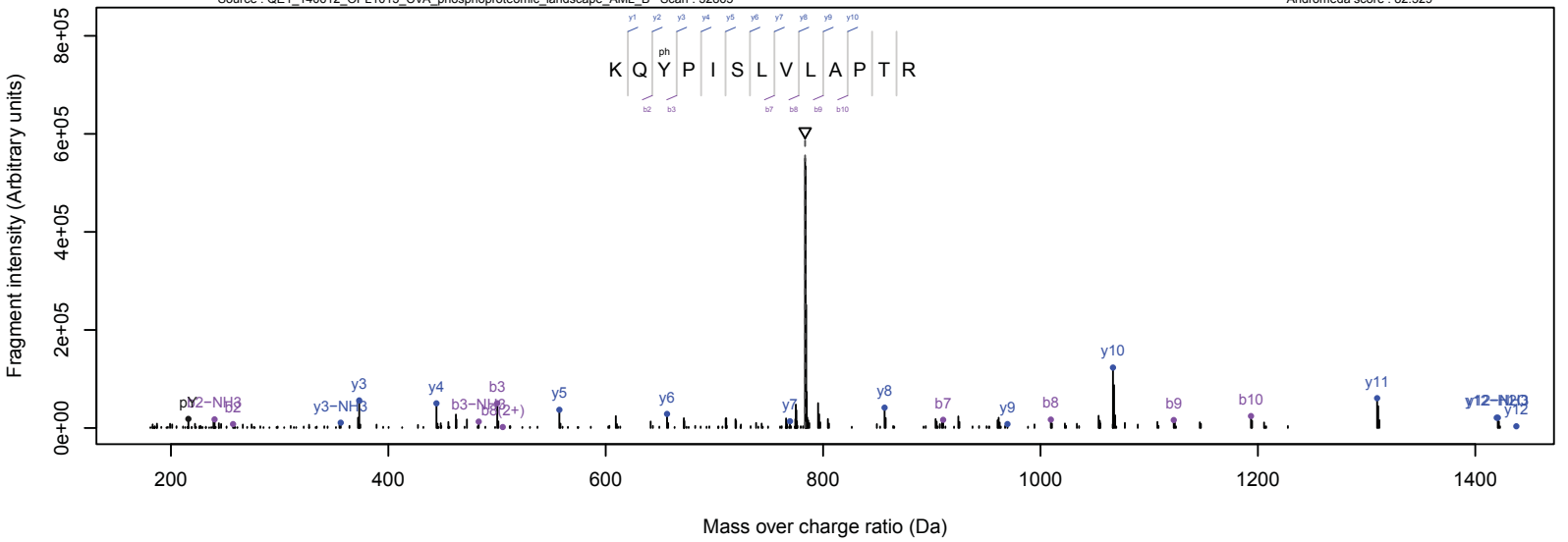
Andromeda score : 82.59



KQY(ph)PISLVLAPTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32865

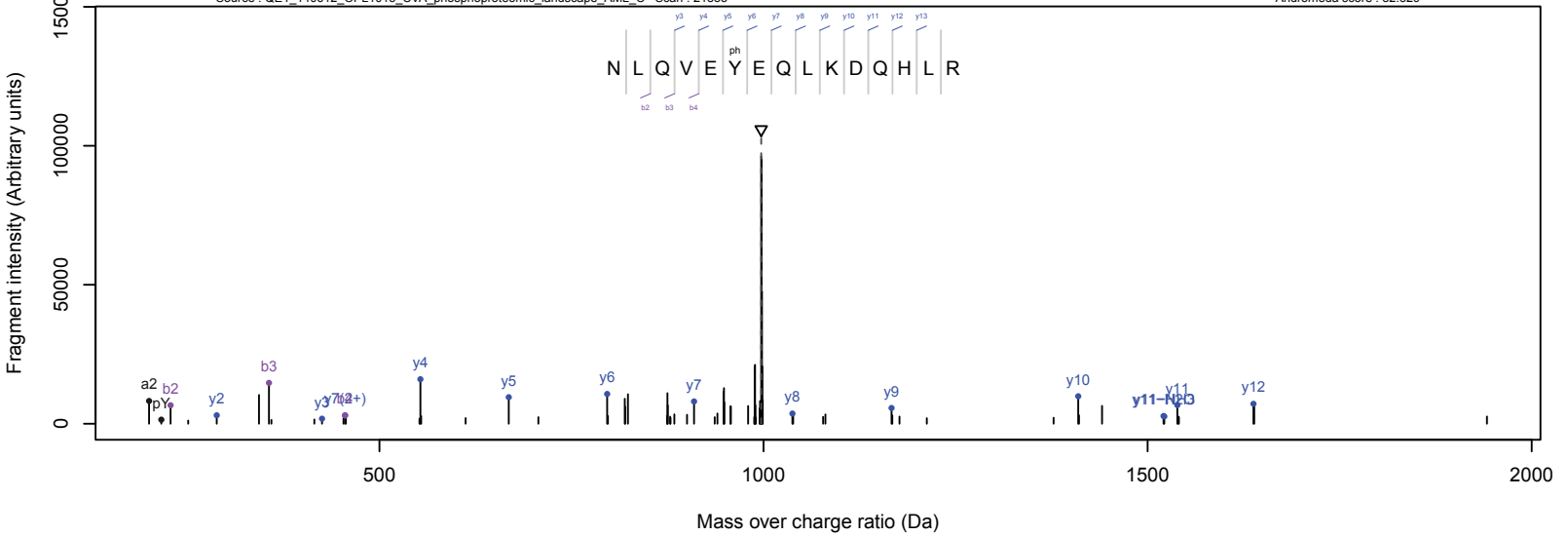
Andromeda score : 82.529



NLQVEY(ph)EQLKDQHLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21853

Andromeda score : 82.529



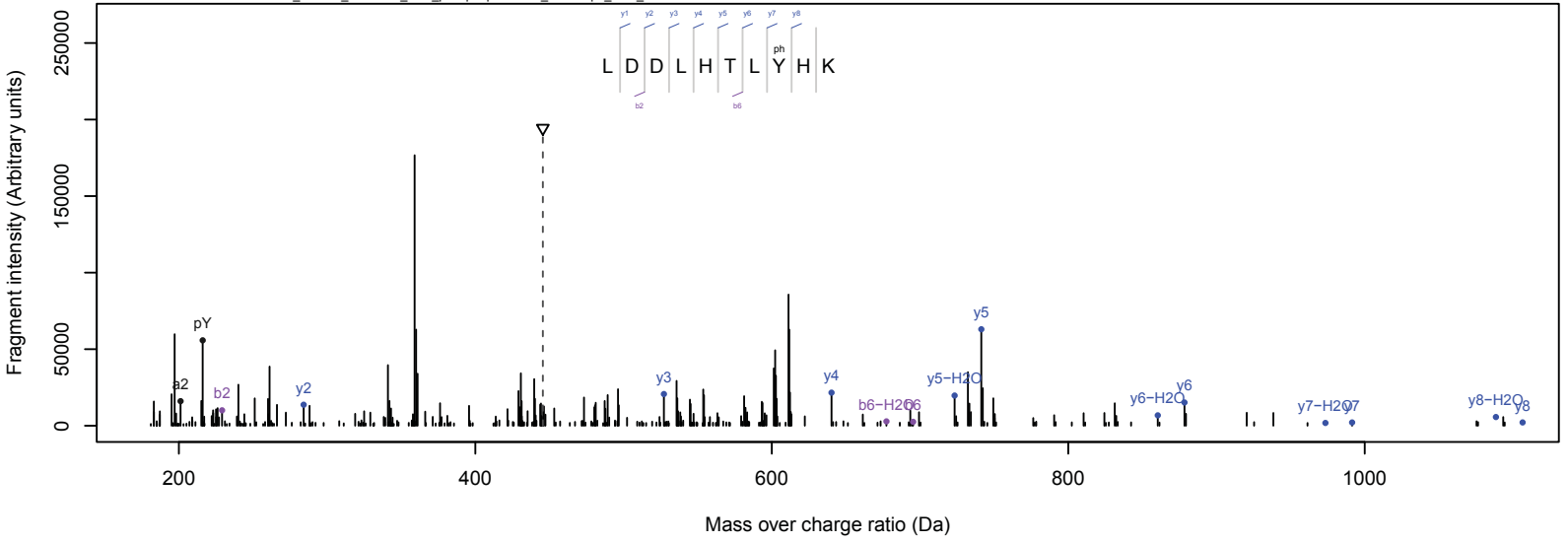




LDDLHTLYHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13165

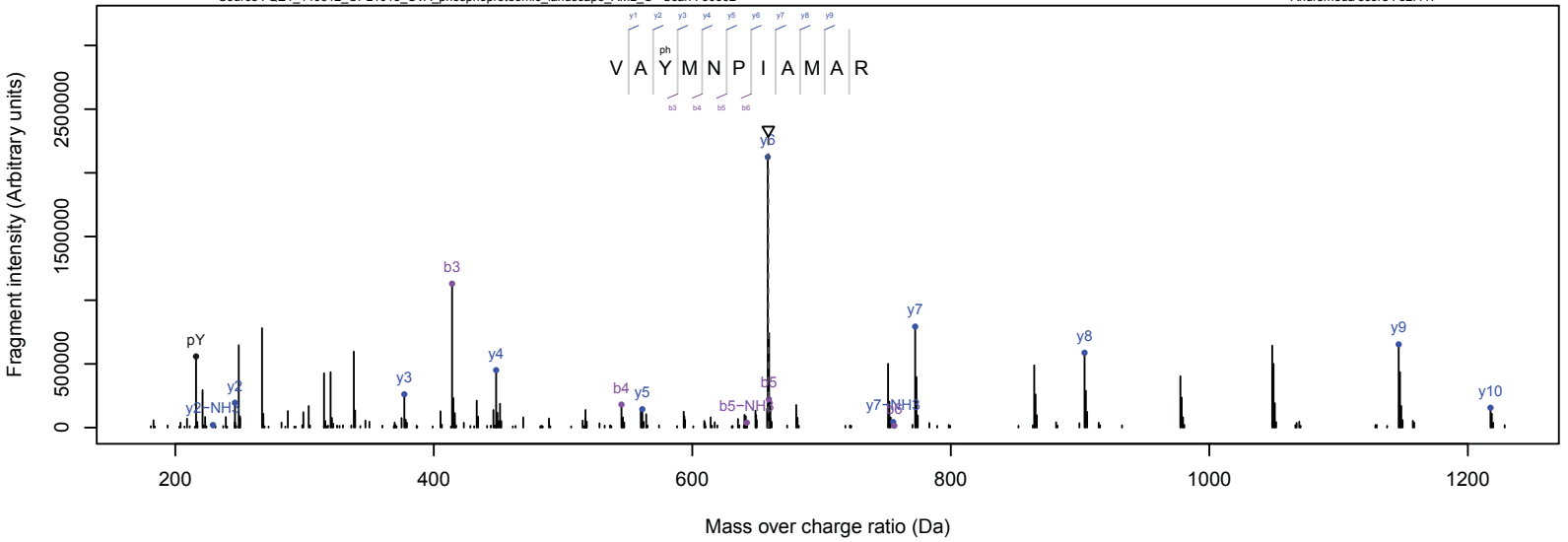
Andromeda score : 82.417



VAYMNPIMAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30302

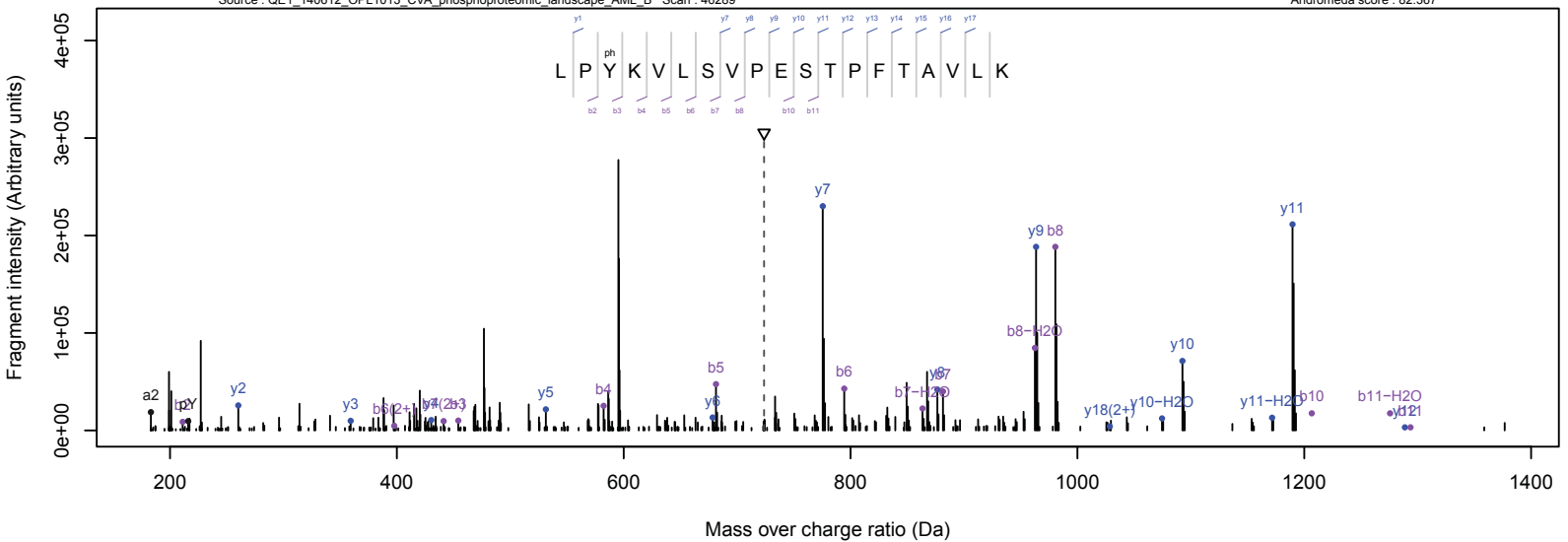
Andromeda score : 82.417



LPYKVLSPVPESTPFTAVLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46289

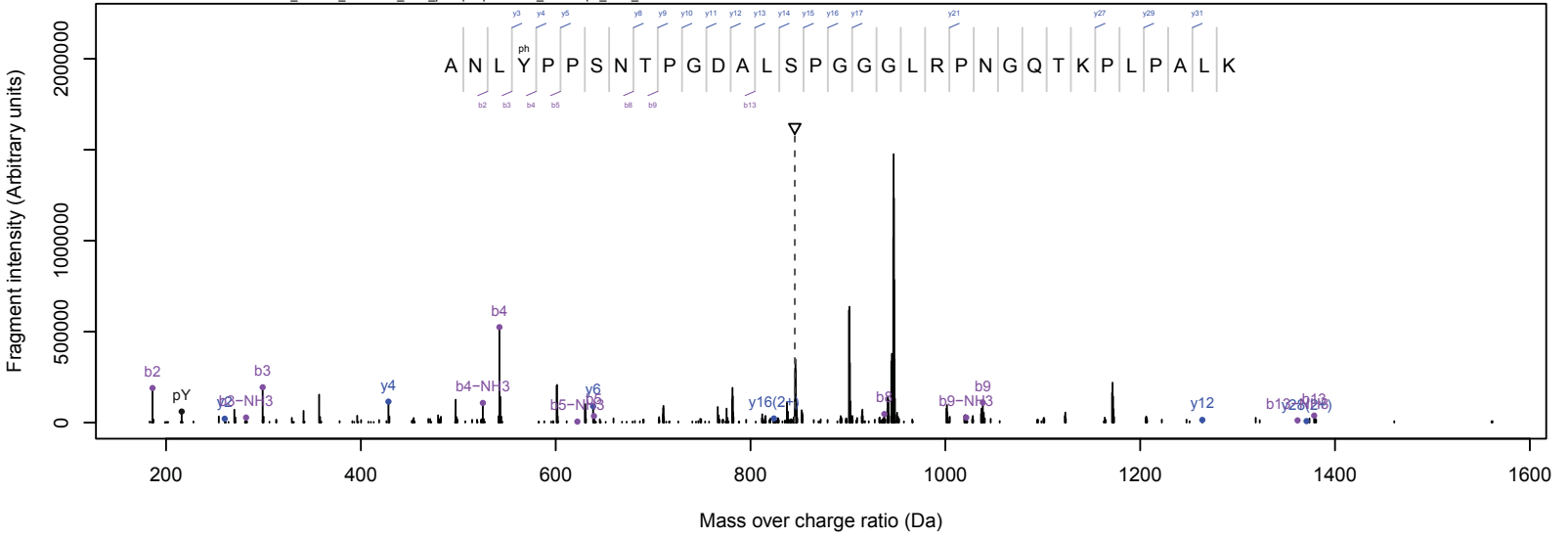
Andromeda score : 82.367



**ANLY(ph)PPSNTPGDALSPPGGGLRPNQGQTKPLPALK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32532

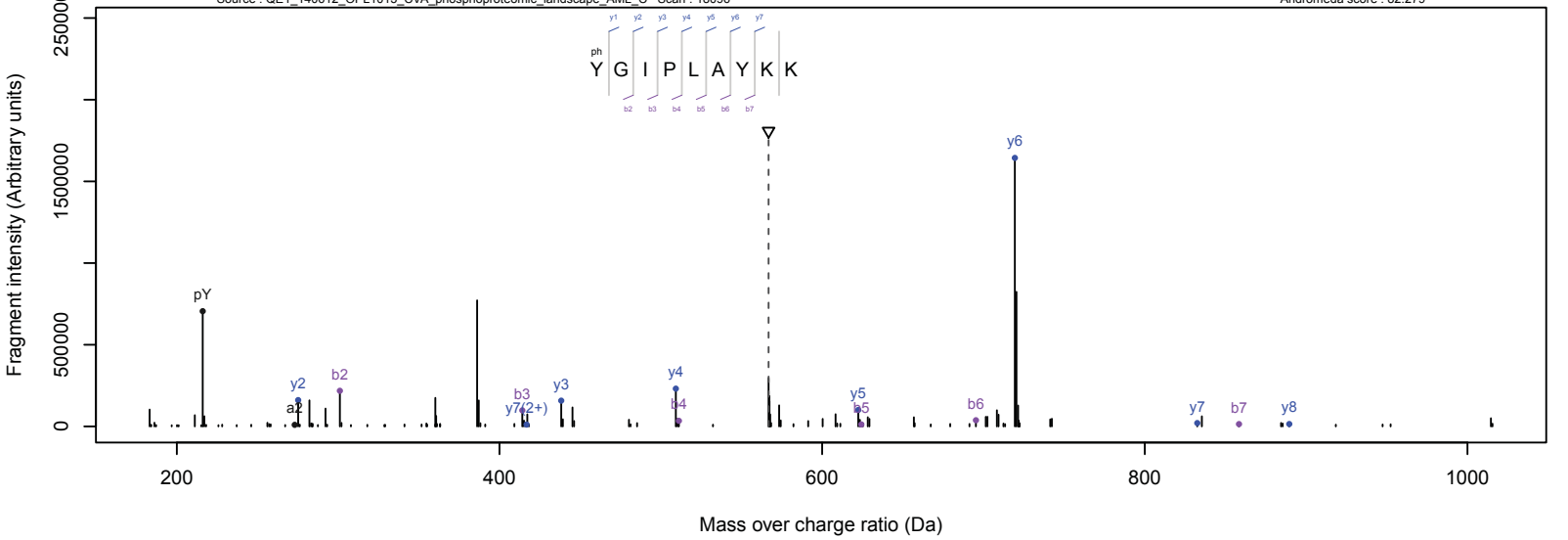
Andromeda score : 82.364



**Y(ph)GIPLAYKK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18096

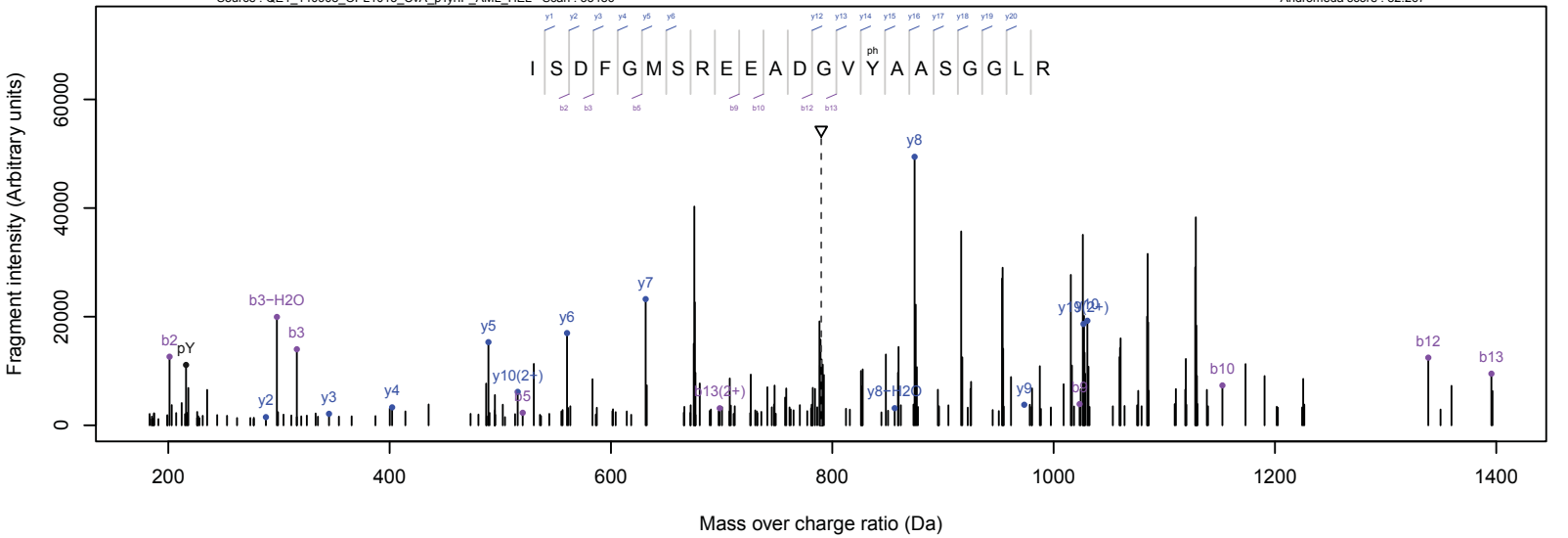
Andromeda score : 82.279



**ISDFGMSREEADGVY(ph)AASGGLR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 33486

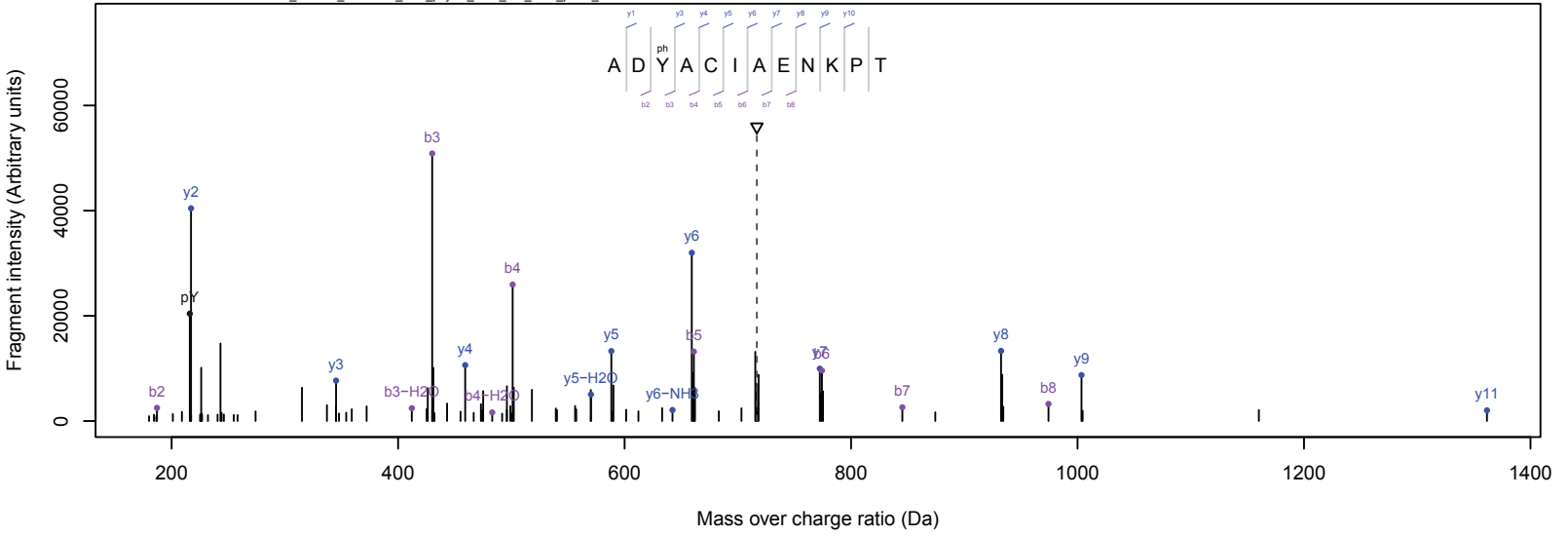
Andromeda score : 82.267



ADY(ph)ACIAENKPT-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 8186

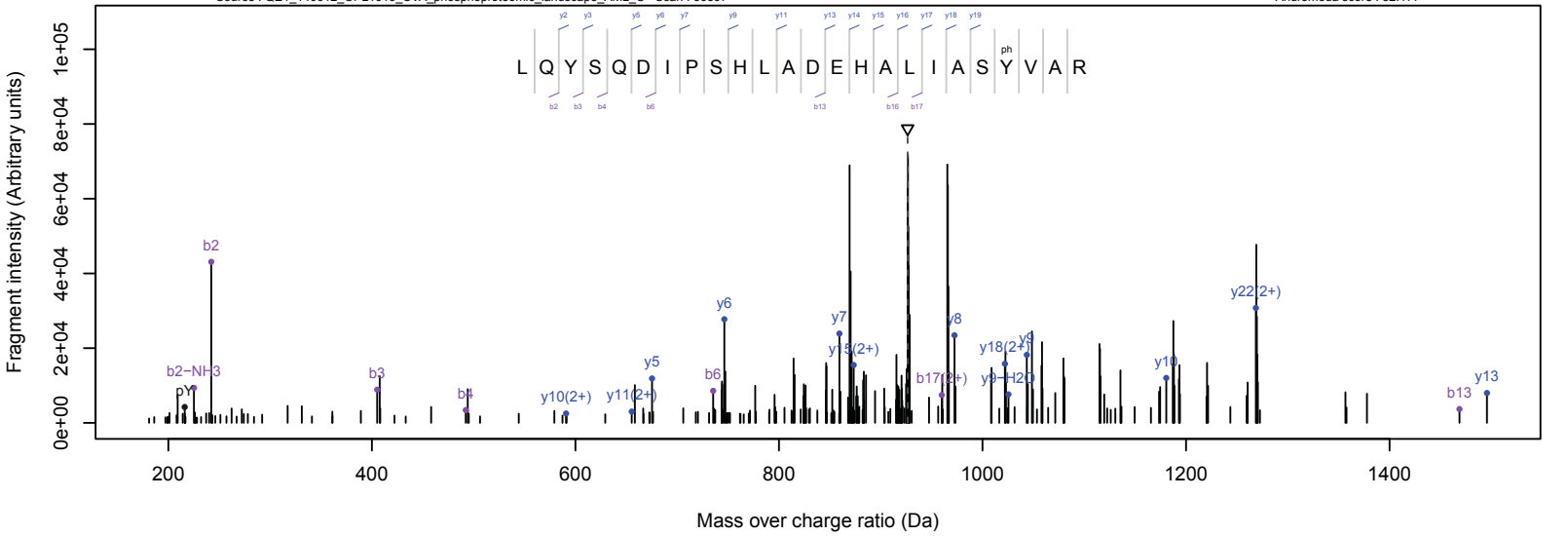
Andromeda score : 82.202



LQYSQDIPSHLADEHALIASY(ph)VAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36897

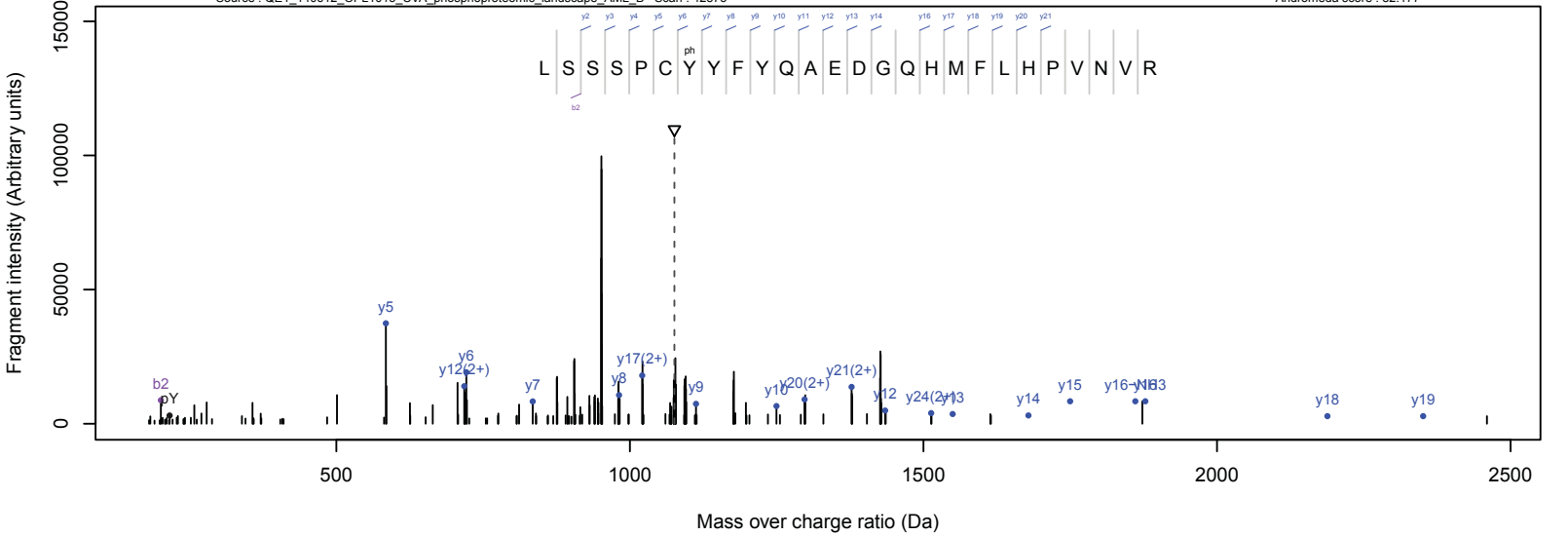
Andromeda score : 82.177



LSSSPCY(ph)YFYQAEDGQHMFHPVNVNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42873

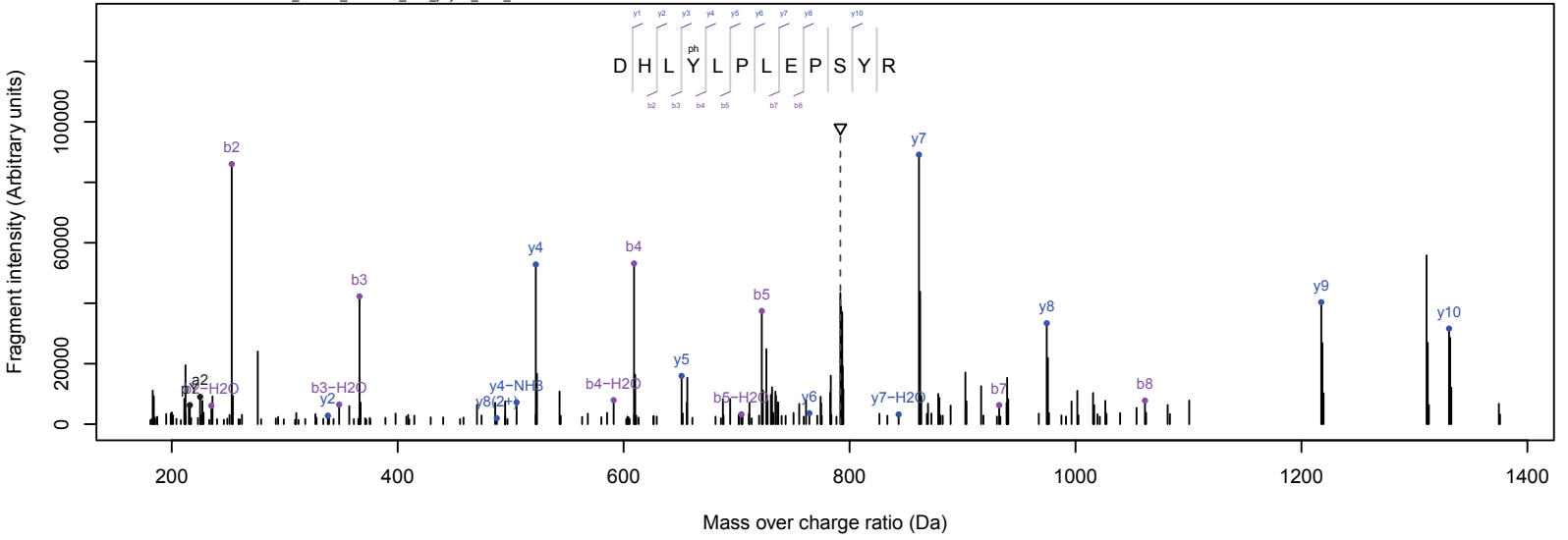
Andromeda score : 82.177



DHLY(ph)LPLEPSYR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37517

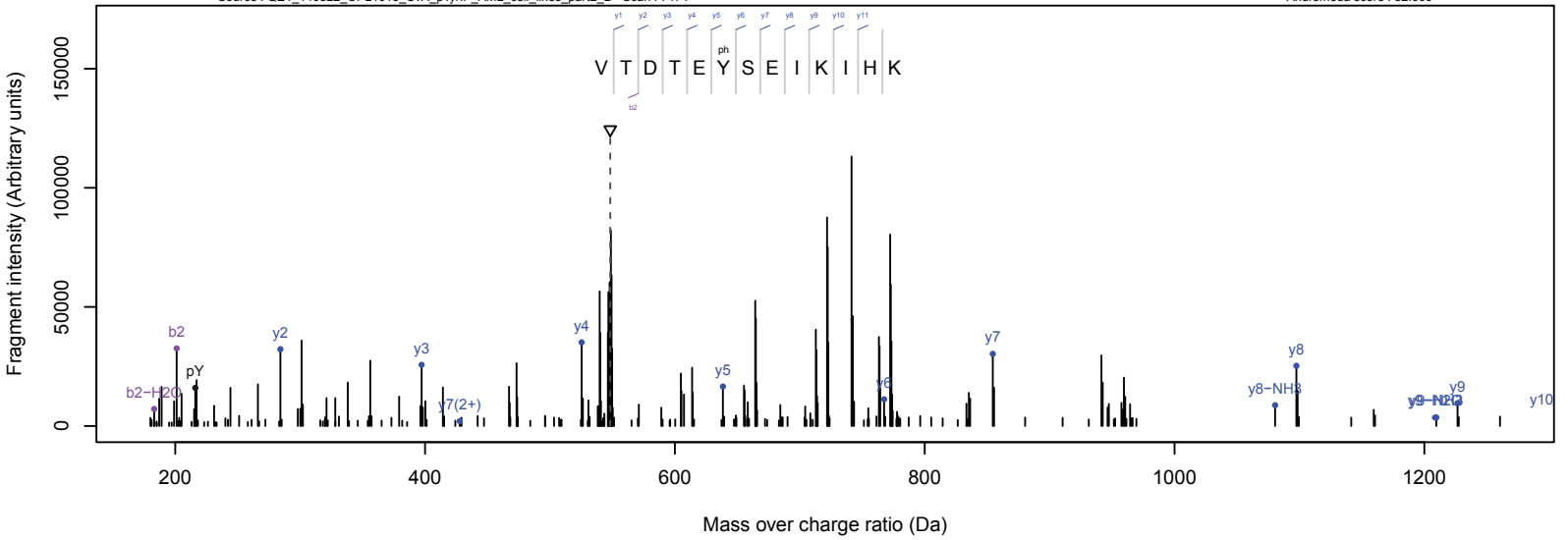
Andromeda score : 82.069



VTDTEY(ph)SEIKHK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 7474

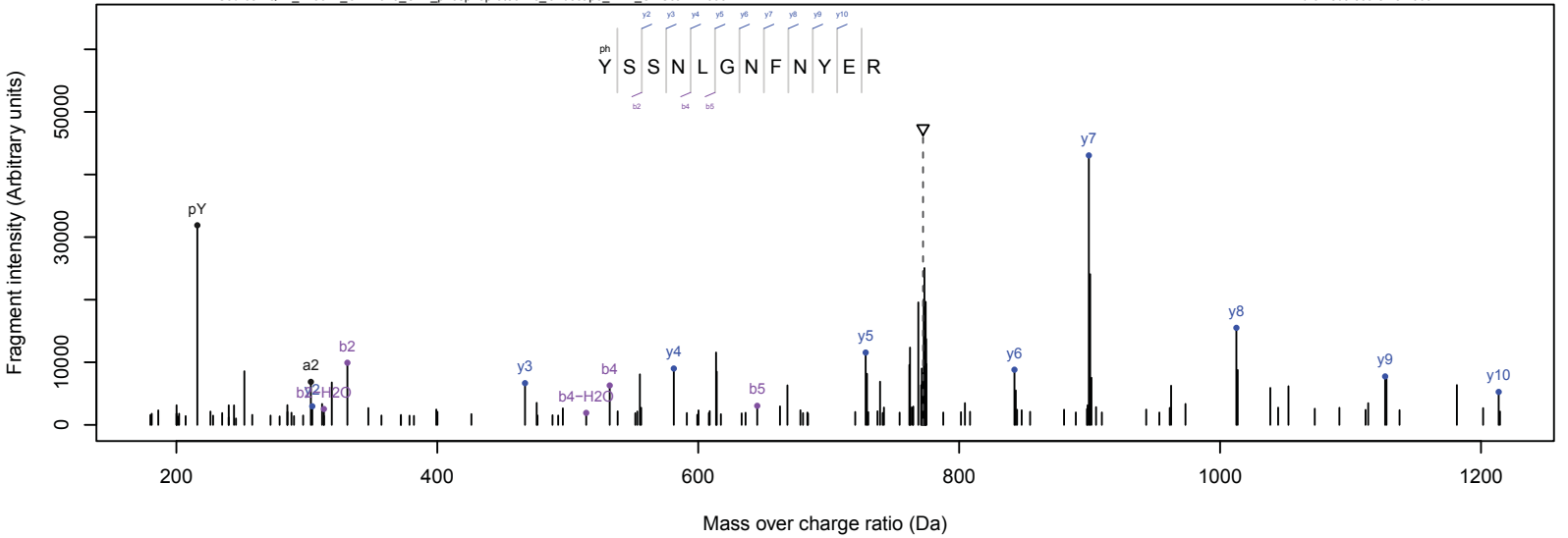
Andromeda score : 82.069



Y(ph)SSNLGNFYER-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 20932

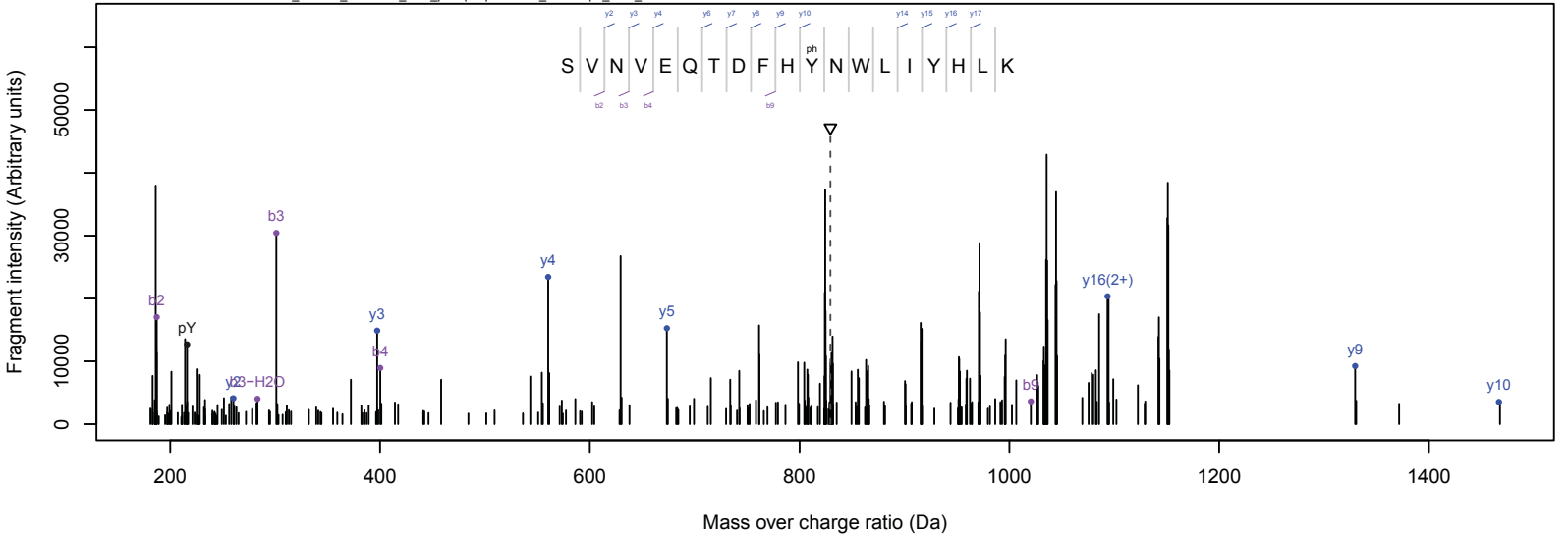
Andromeda score : 82.069



SVNVEQTFHY(ph)NWLIIYHLK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41476

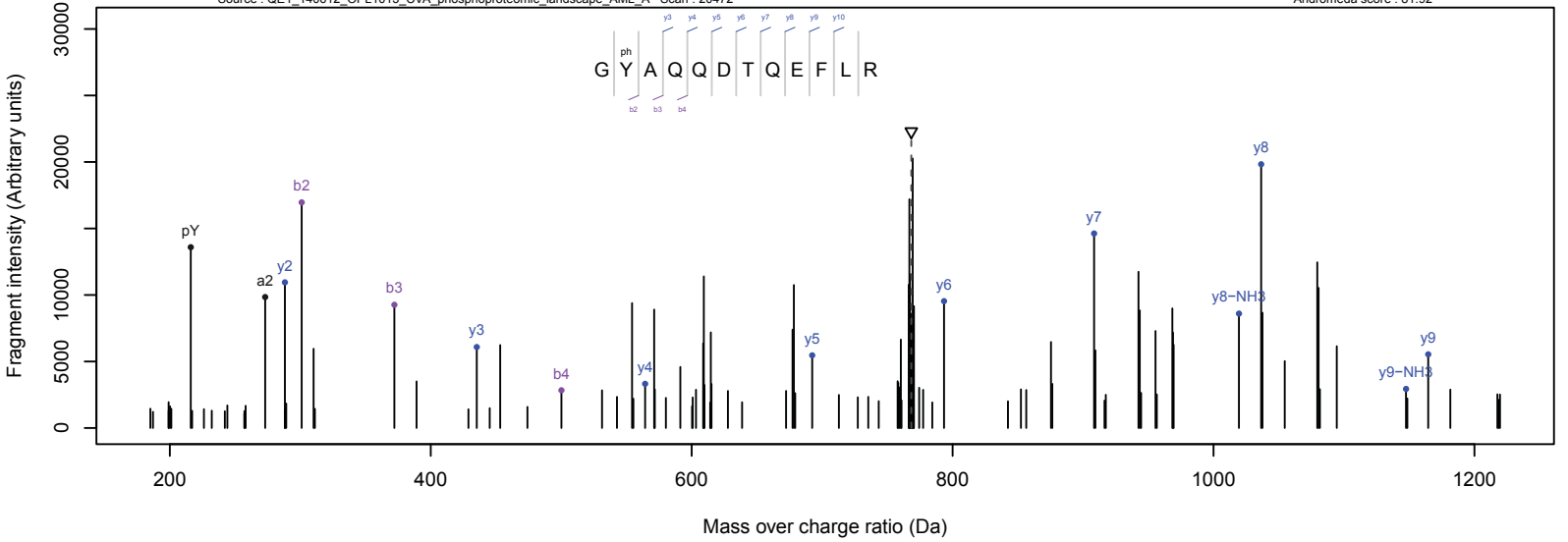
Andromeda score: 82.01



GY(ph)AQDQTQEFLLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 20472

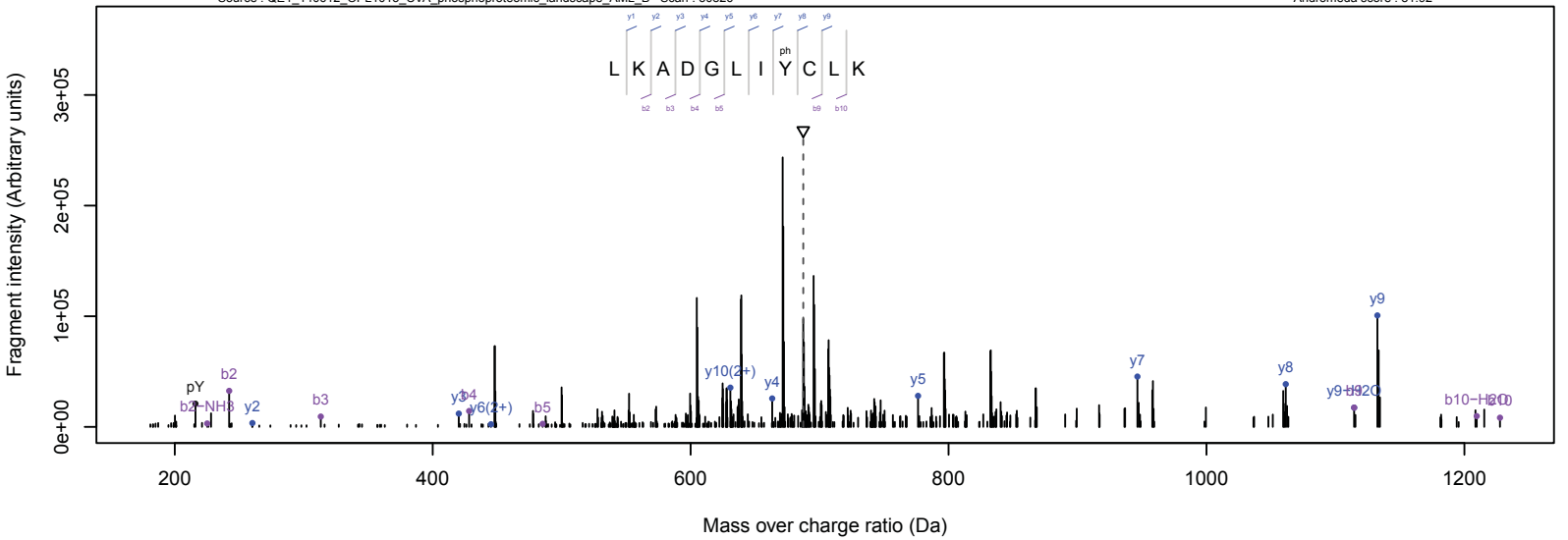
Andromeda score: 81.92



LKADGLIY(ph)CLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 30826

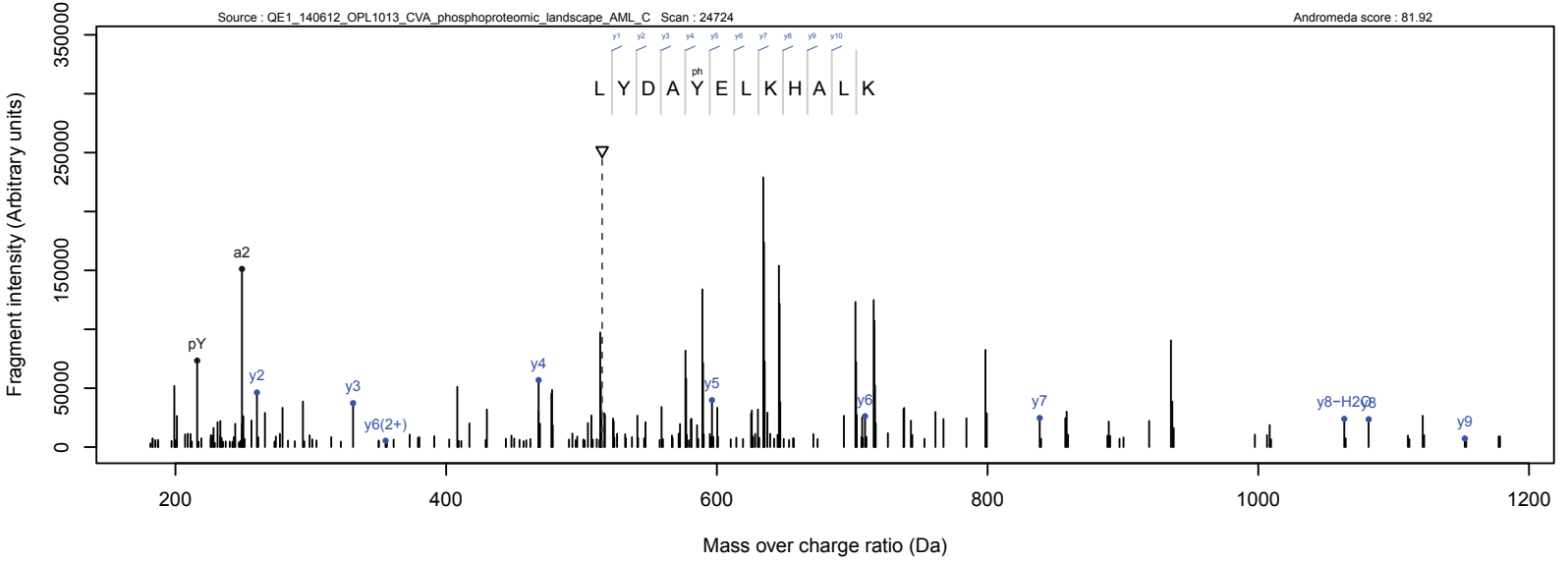
Andromeda score: 81.92



LYDAY(ph)ELKHALK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24724

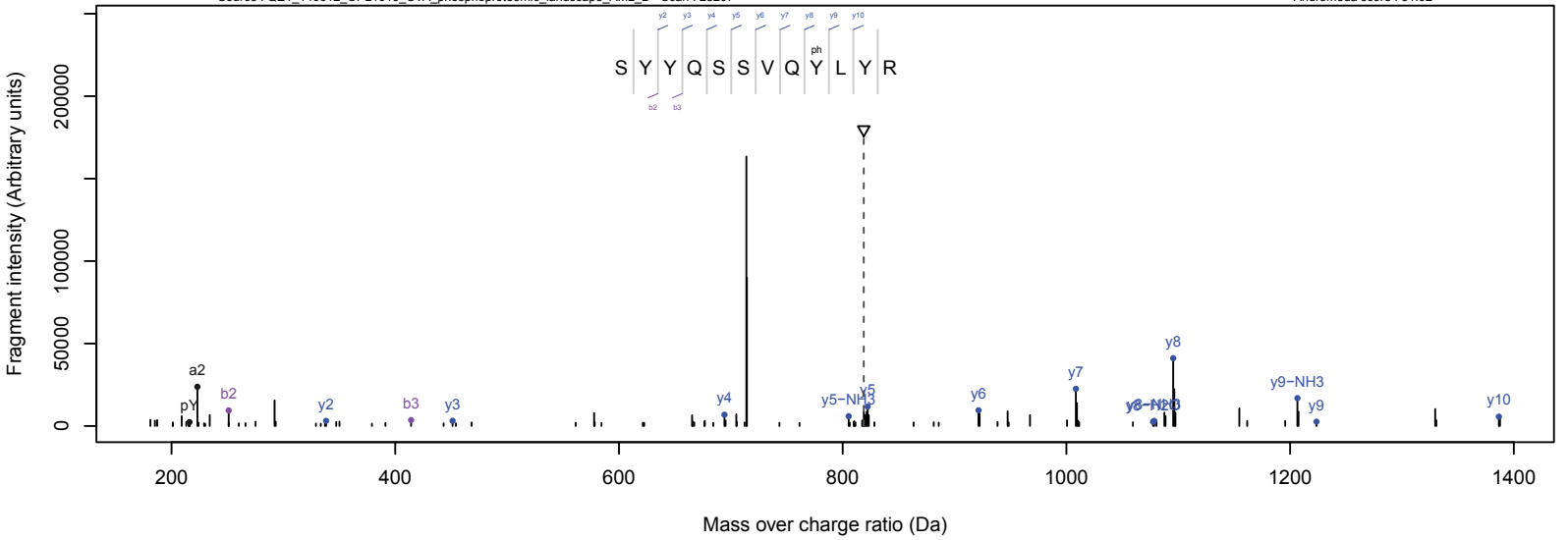
Andromeda score : 81.92



SYQQSSVQ(ph)LYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28207

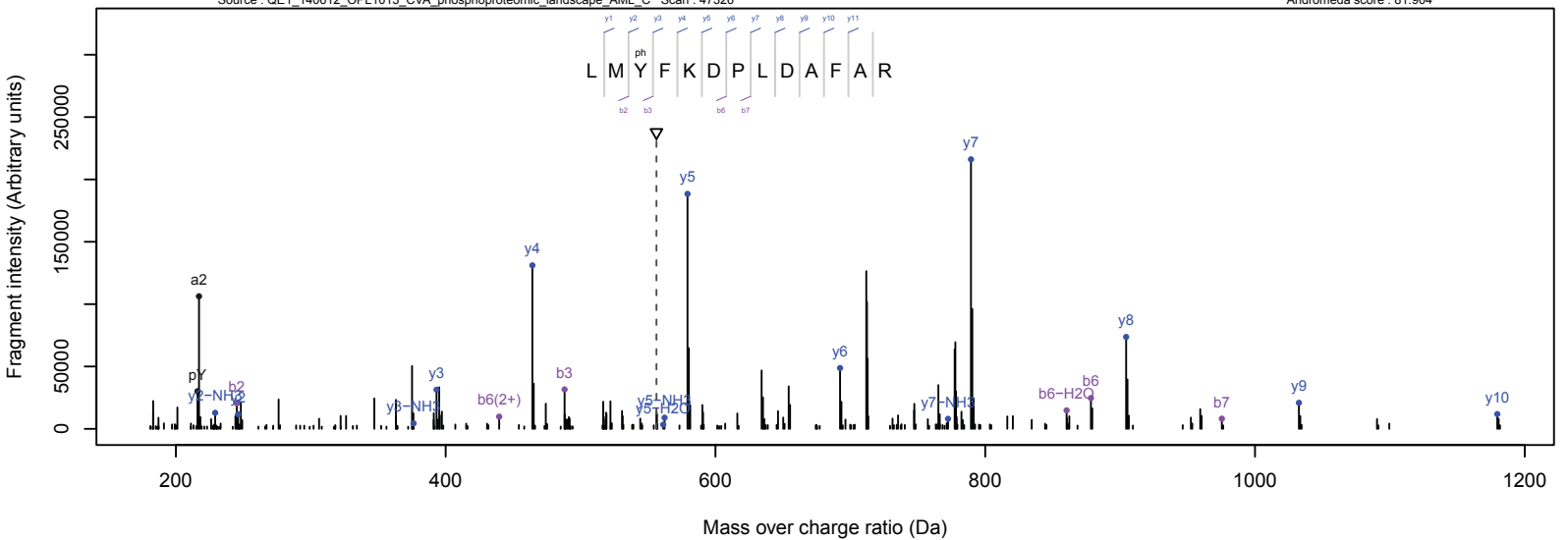
Andromeda score : 81.92



LMY(ph)FKDPLDAFAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47326

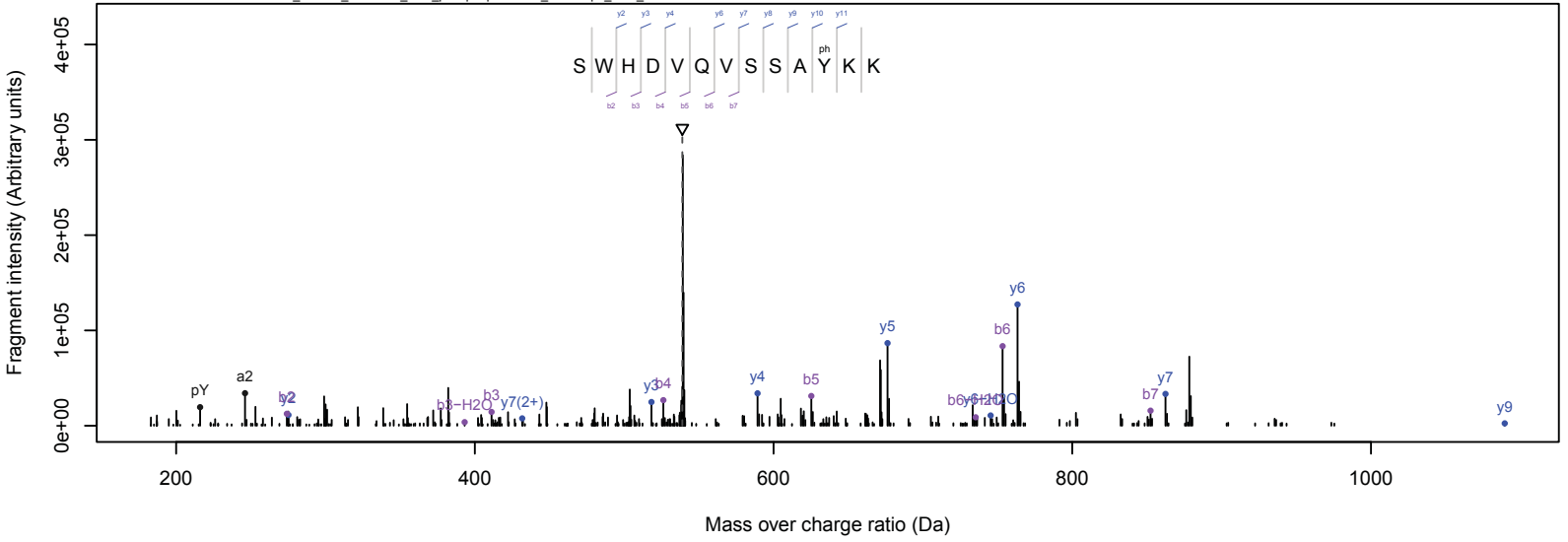
Andromeda score : 81.904



SWHDVQSSAY(ph)KK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10327

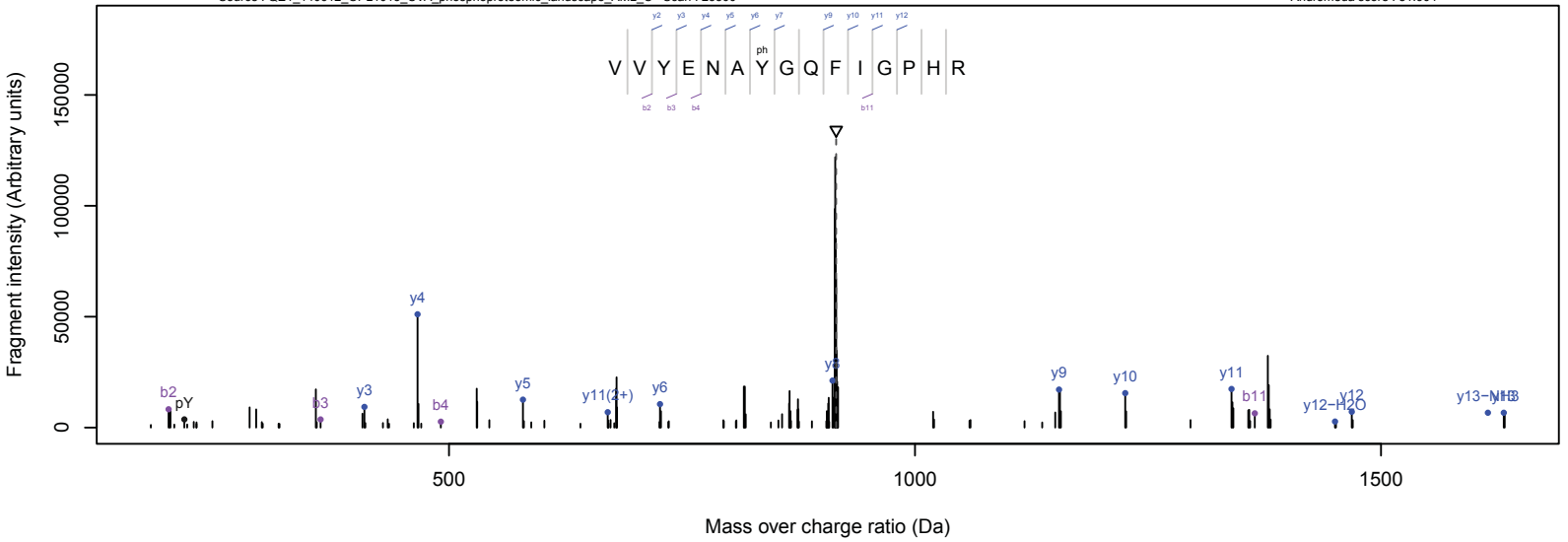
Andromeda score : 81.904



VVYENAY(ph)GQFIGPHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23500

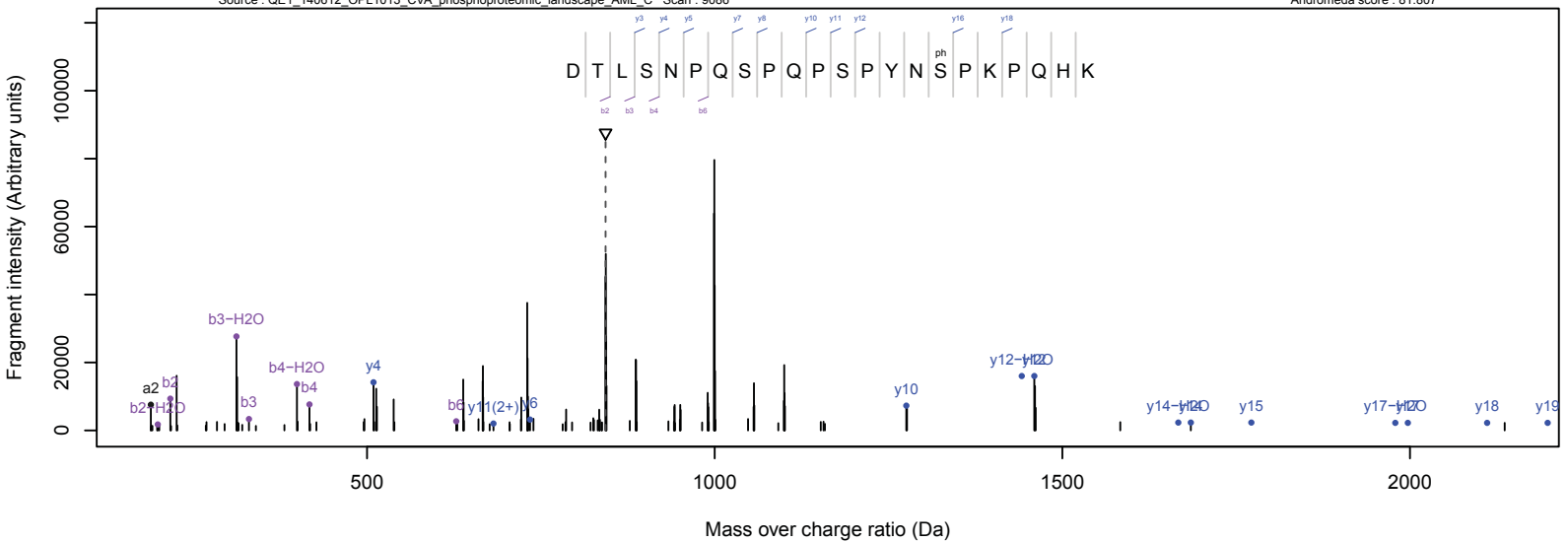
Andromeda score : 81.904



DTLSNPQSPQSPYNS(ph)PKPQHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9086

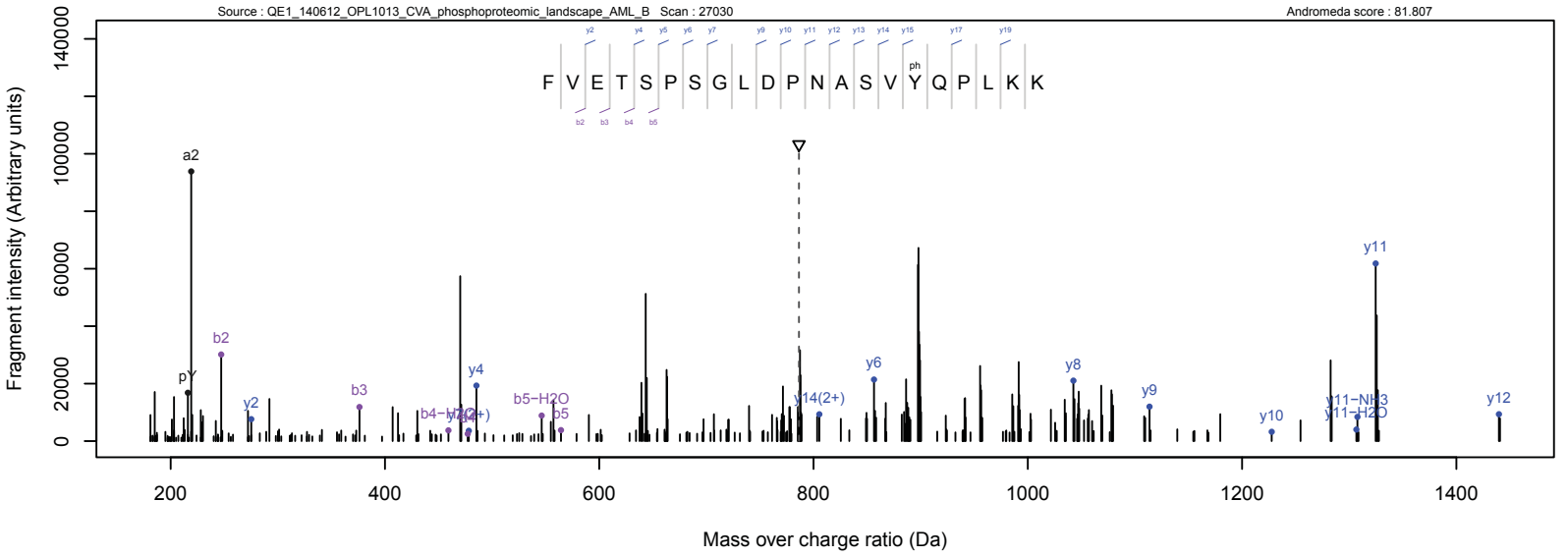
Andromeda score : 81.807



**FVETSPGLDPNASV(ph)QPLKK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27030

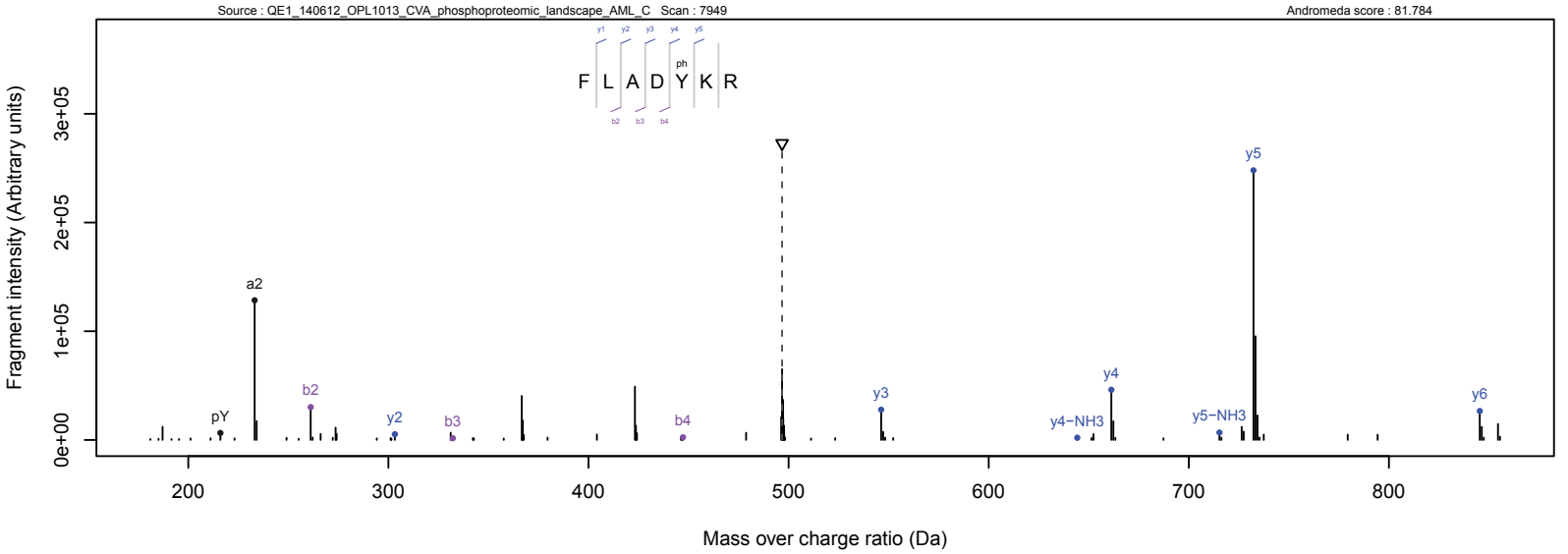
Andromeda score : 81.807



**FLADY(ph)KR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7949

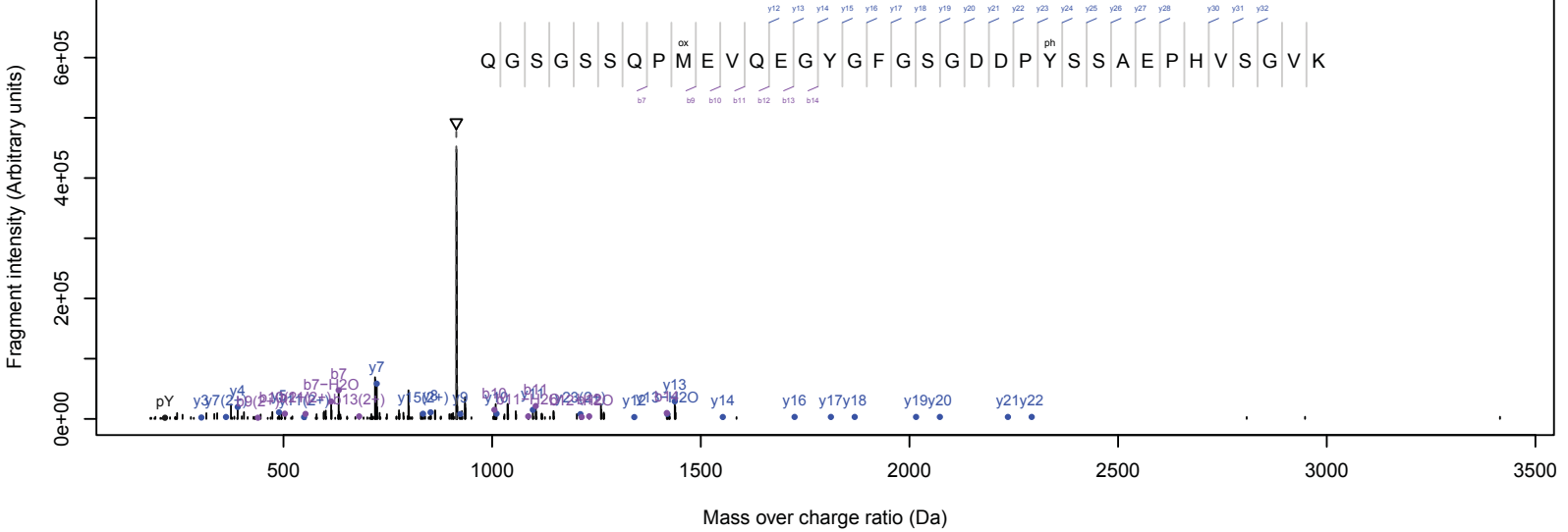
Andromeda score : 81.784



**QGS GSS QP M E V Q E G Y G F G S G D D P Y S S A E P H V S G V K-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 24695

Andromeda score : 81.718

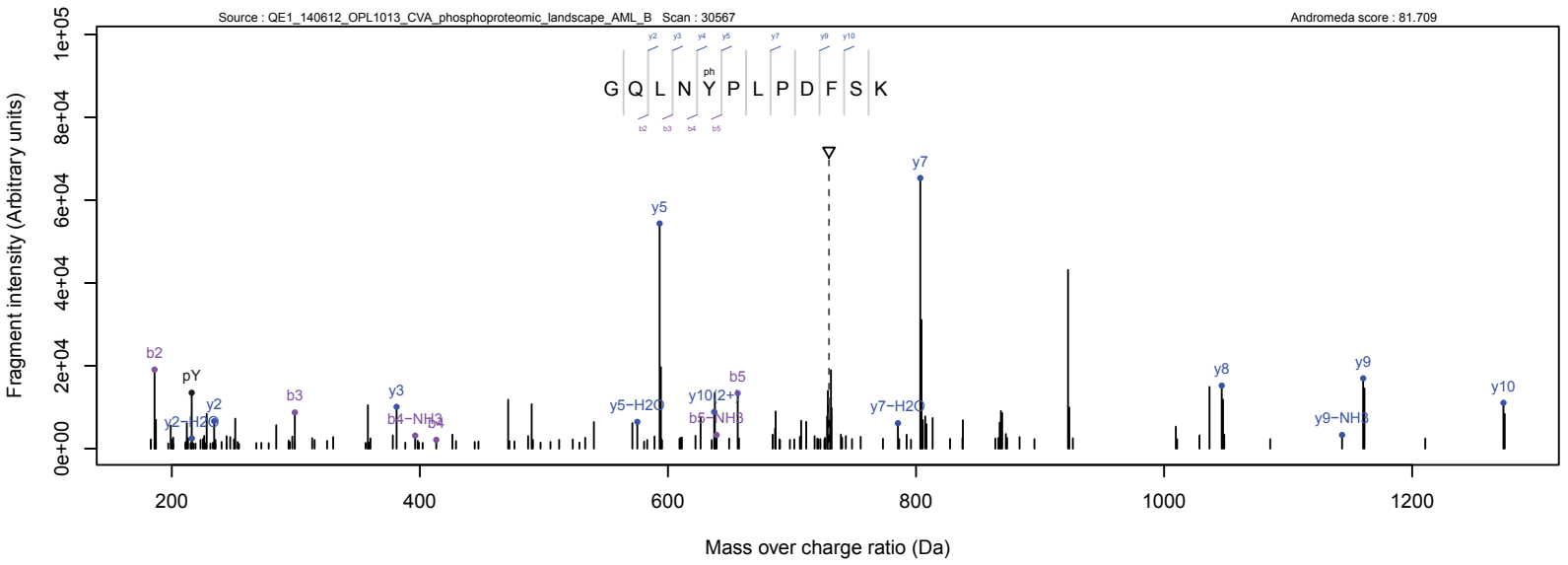




GQLNY(ph)PLPDFSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30567

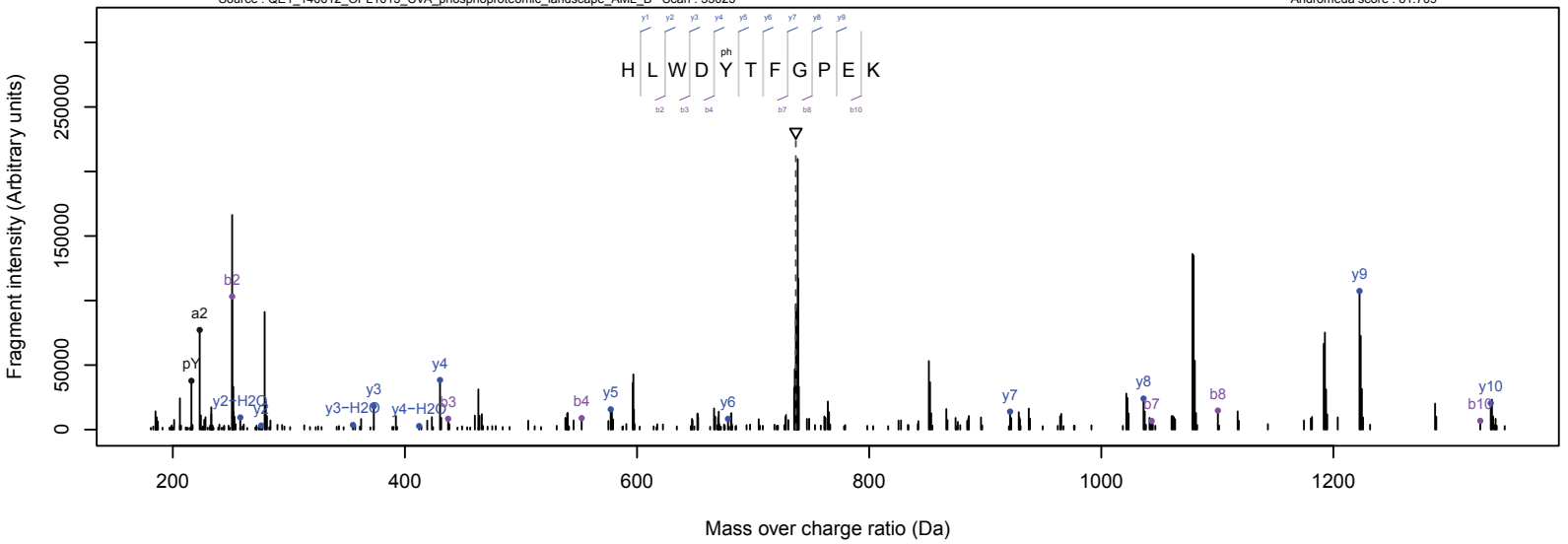
Andromeda score : 81.709



HLWDY(ph)TFGPEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33023

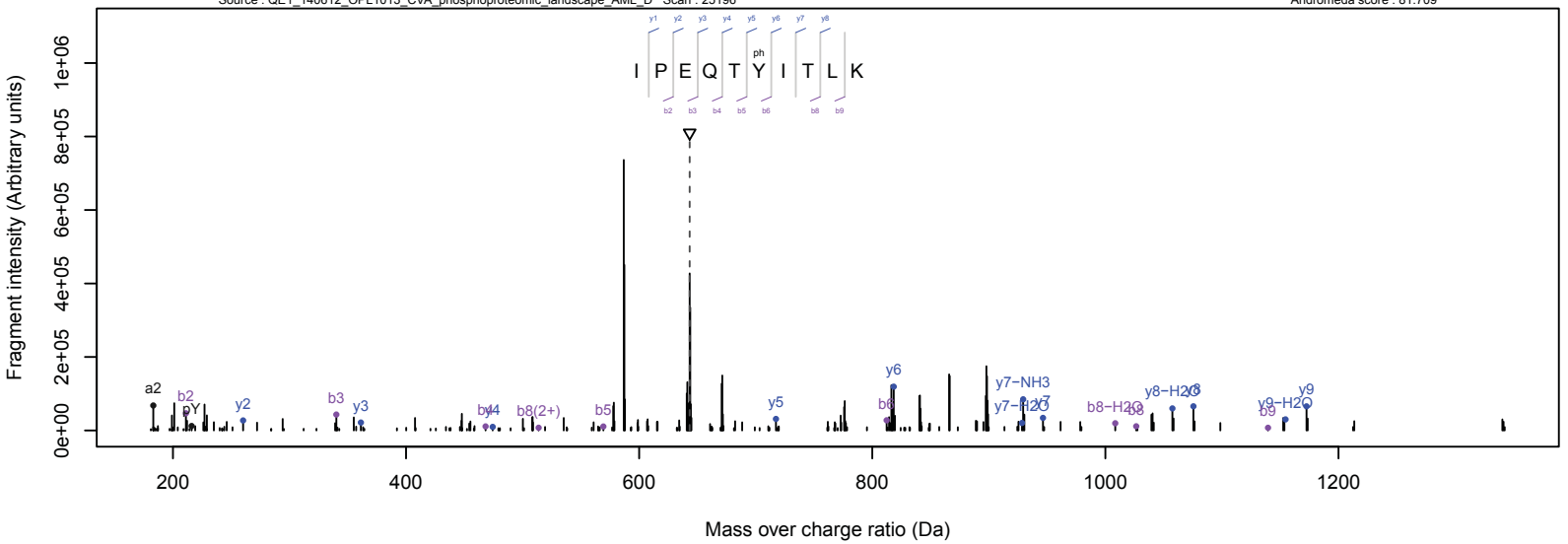
Andromeda score : 81.709



IPEQTY(ph)ITLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 25196

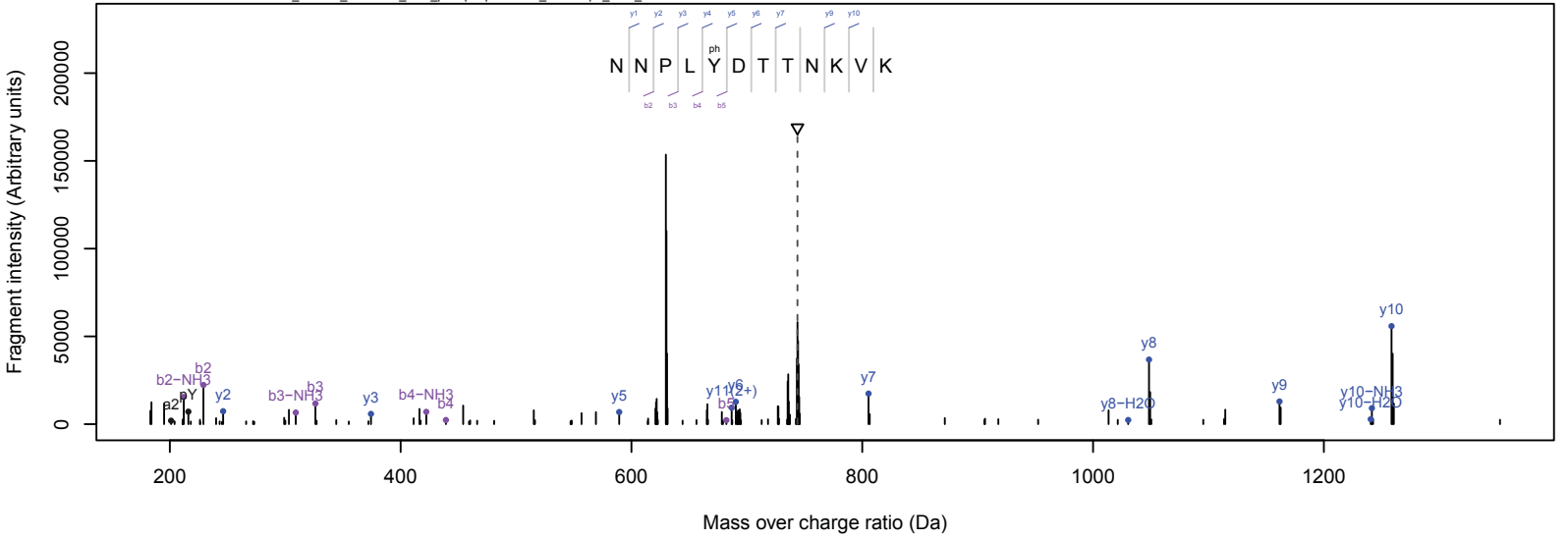
Andromeda score : 81.709



**NNPLY(ph)DTTNKVK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13429

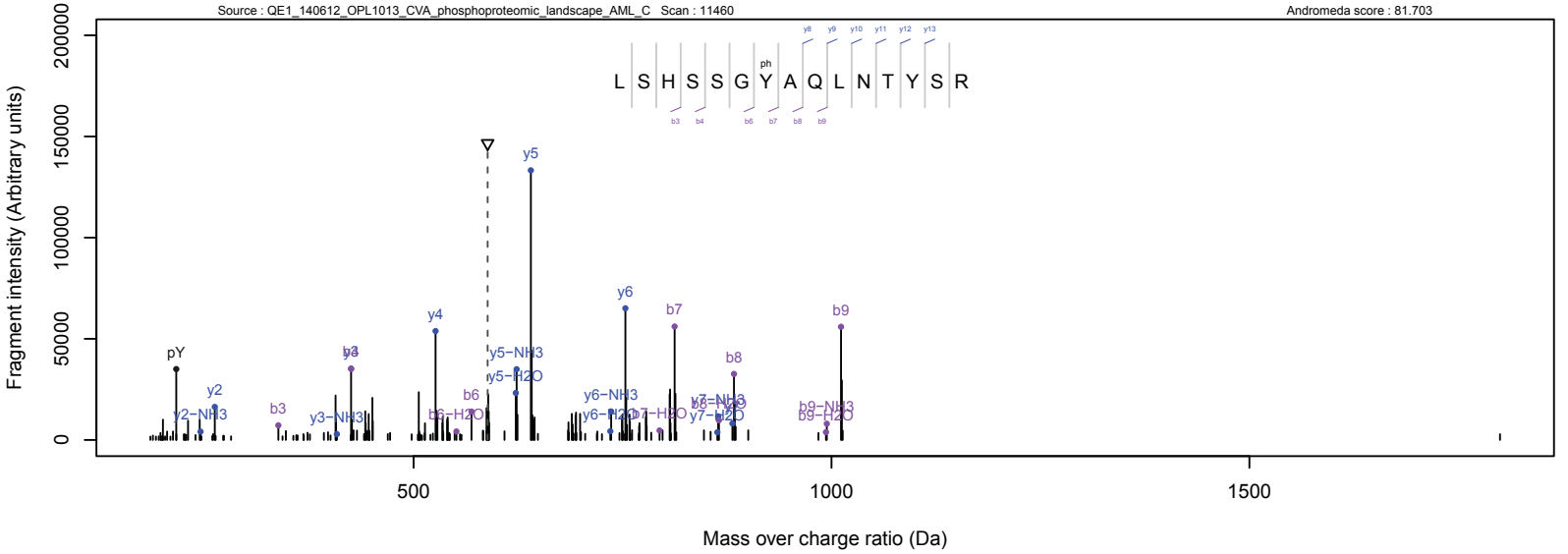
Andromeda score : 81.709



**LSHSSGY(ph)AQLNTYSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11460

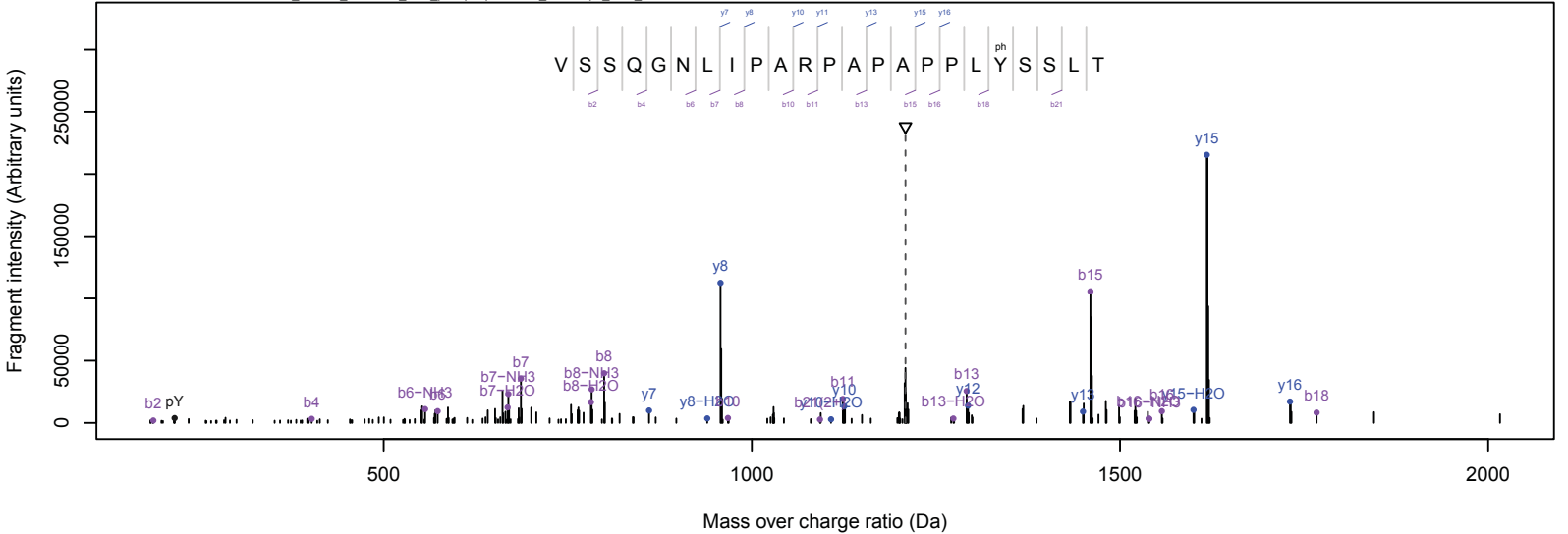
Andromeda score : 81.703



**VSSQGNLIPARPAPAPPLY(ph)SSLT-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38562

Andromeda score : 81.696

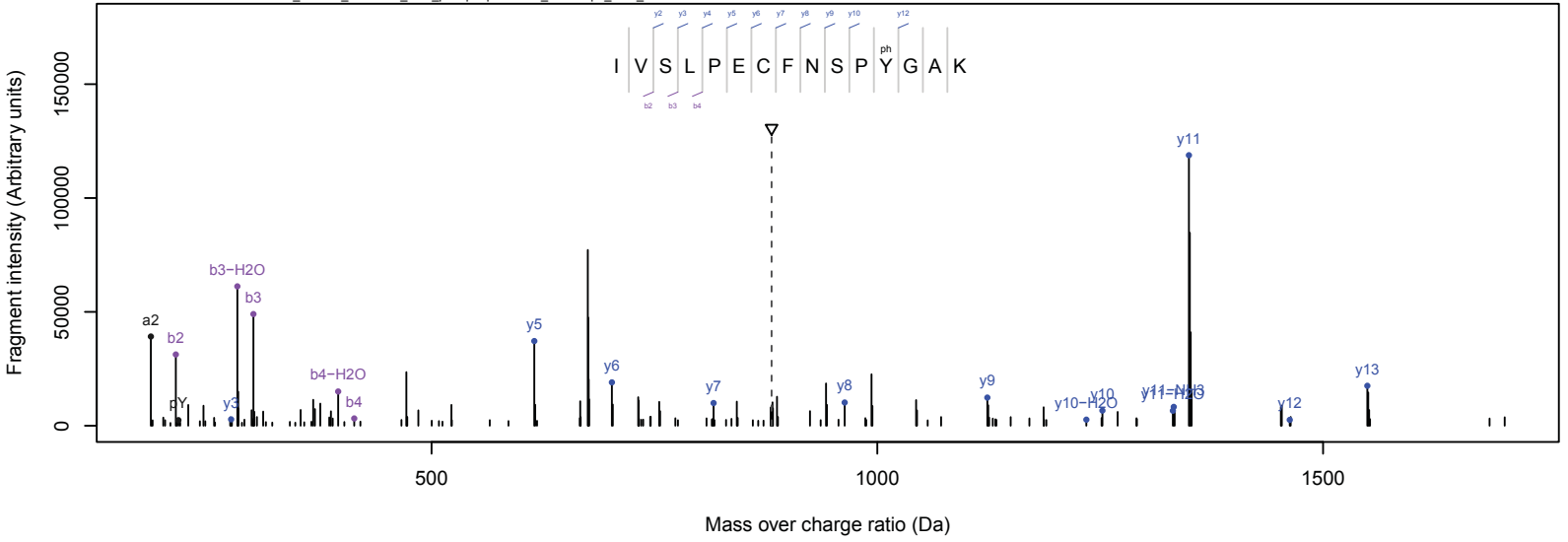




IVSLPECFNSPY(ph)GAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 31617

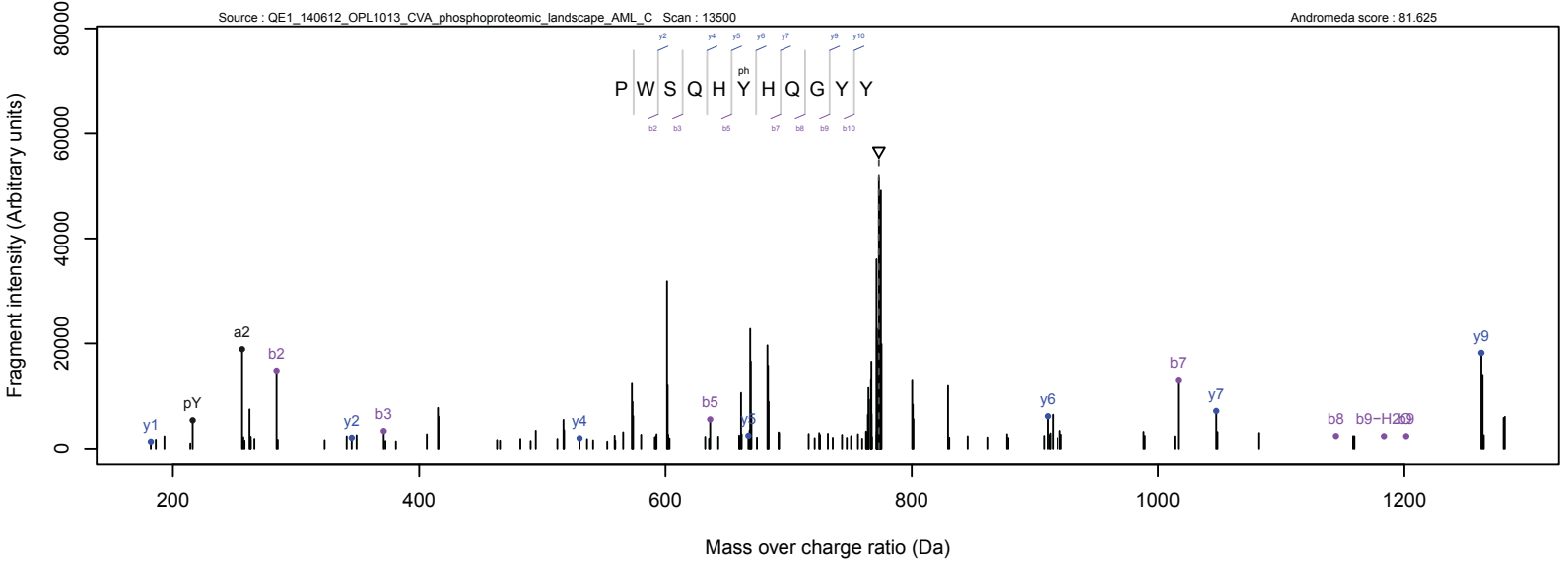
Andromeda score : 81.632



PWSQHY(ph)HQGY-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13500

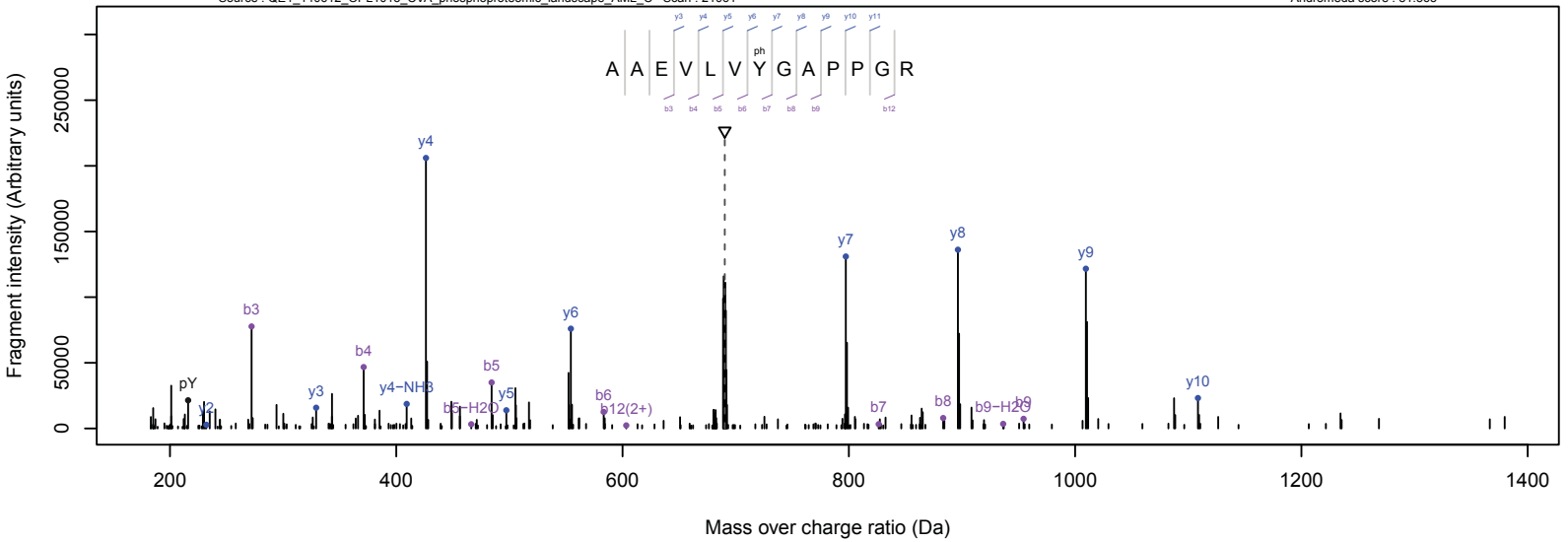
Andromeda score : 81.625



AAEVLVY(ph)GAPPGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21061

Andromeda score : 81.565

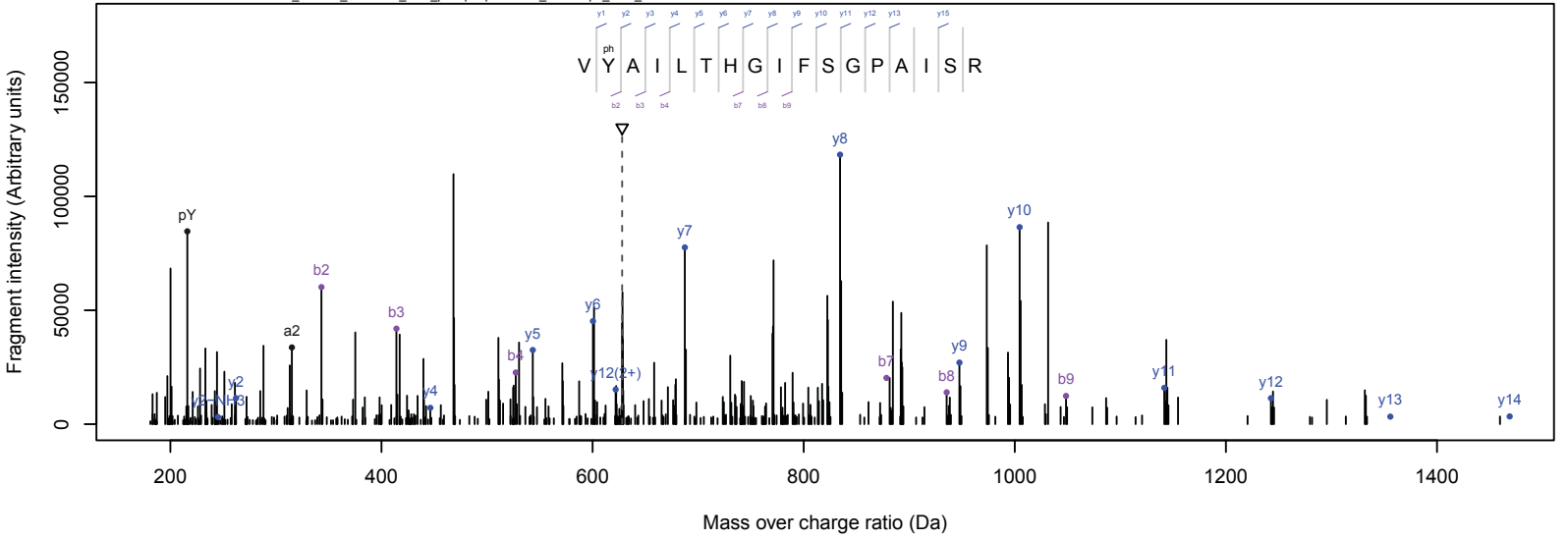




VY(ph)ALTHGIFSGPAISR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40363

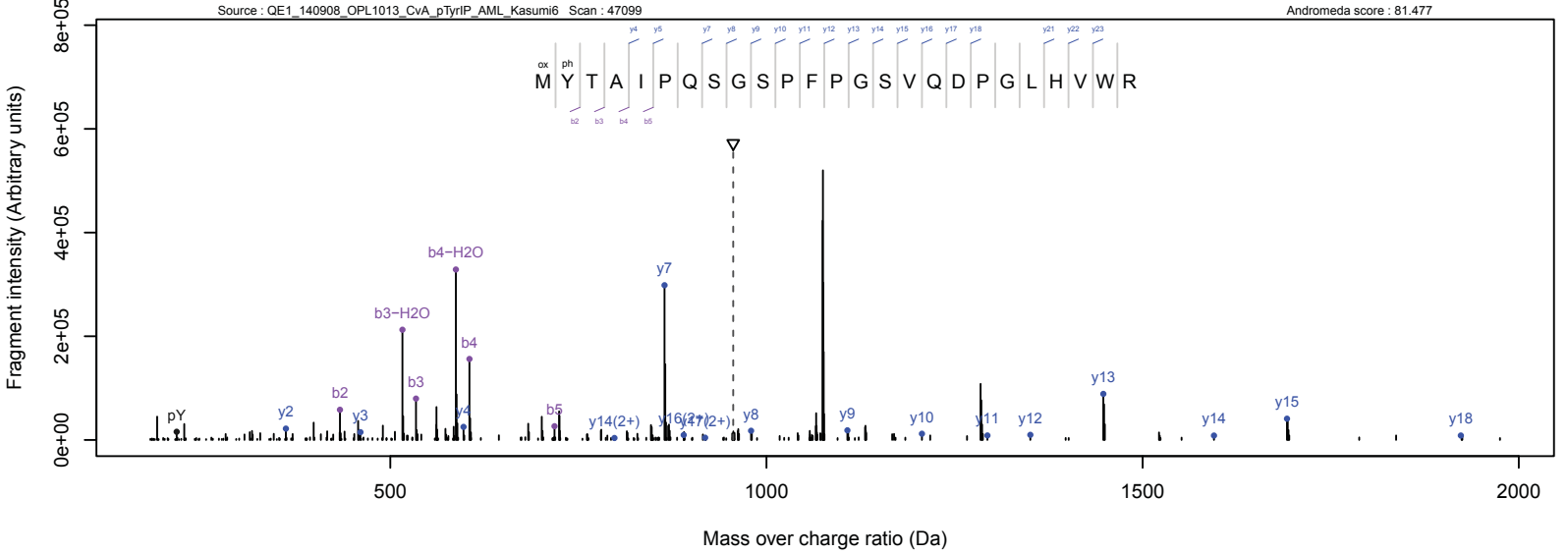
Andromeda score : 81.51



(ac)M(ox)Y(ph)TAIPQSGSPFP GSVQDPLHVWR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 47099

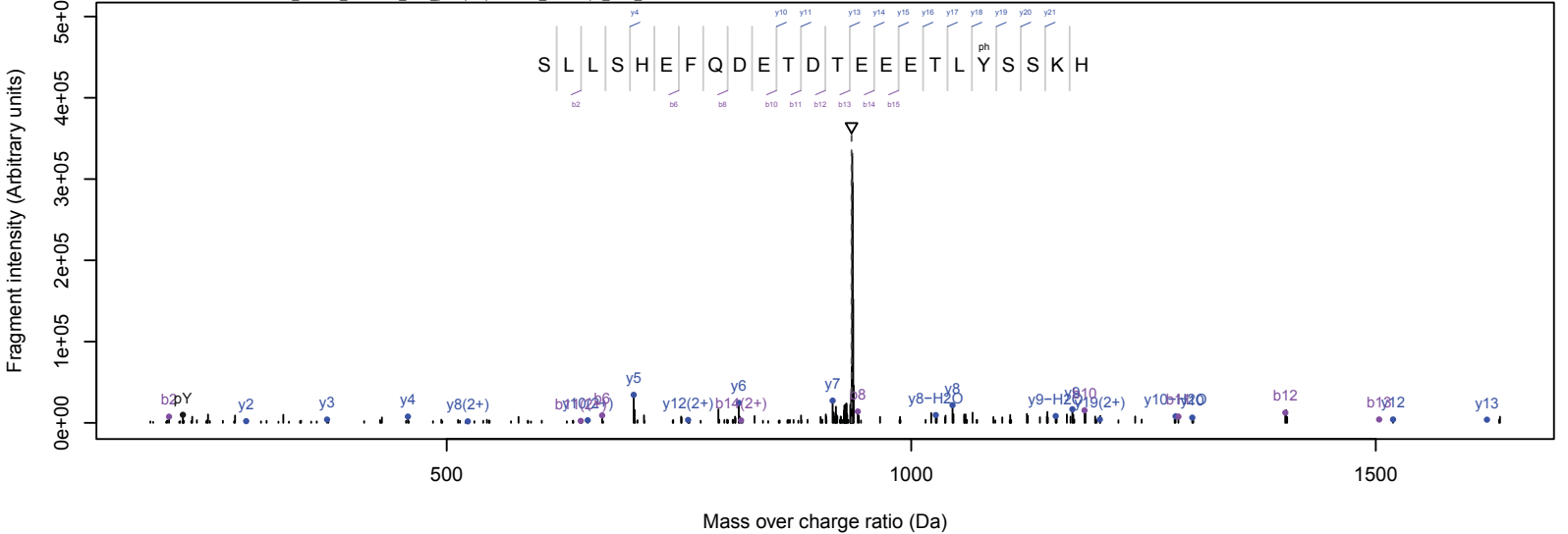
Andromeda score : 81.477



SLLSHEFQDETDEETLY(ph)SSKH-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29763

Andromeda score : 81.477

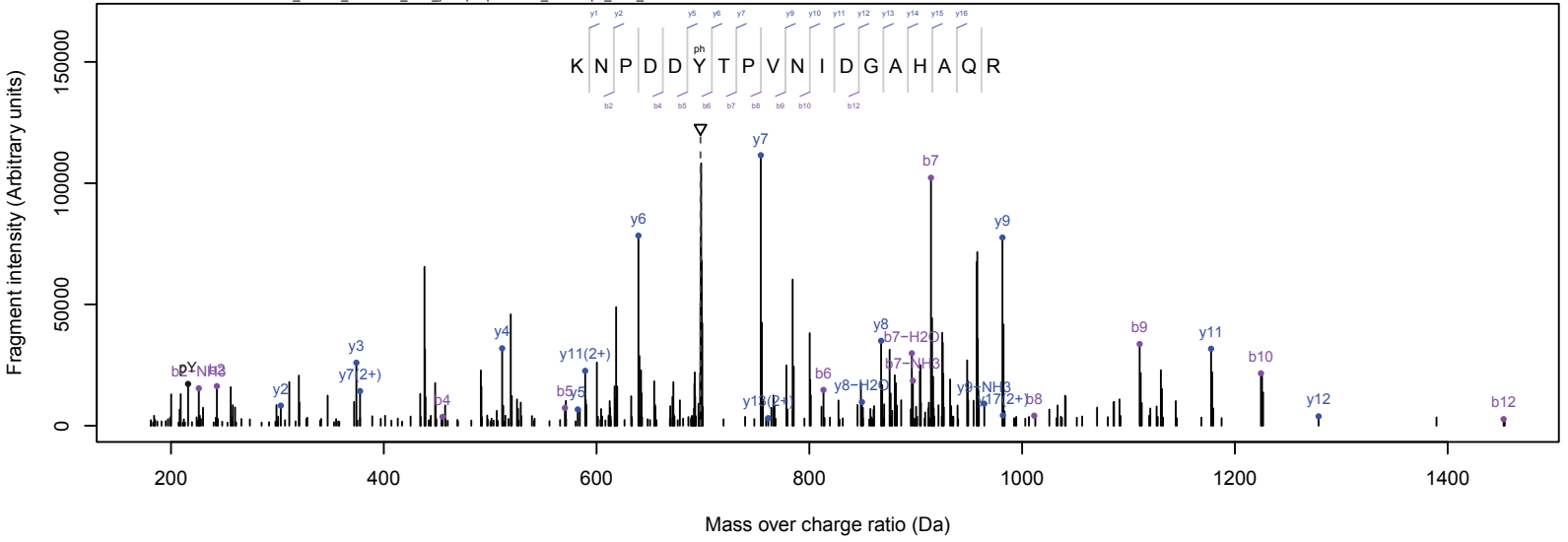




**KNPDDY(ph)TPVNIIDGAHAQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15064

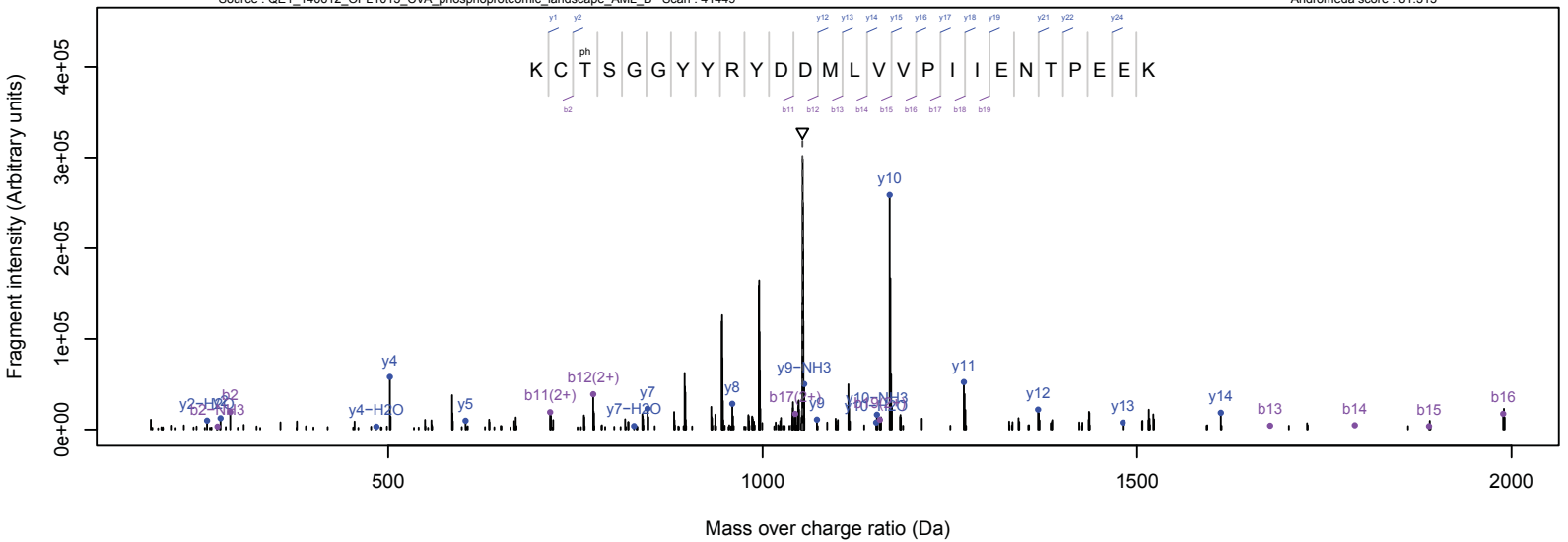
Andromeda score : 81.316



**KCT(ph)SGGYRYDDMLVVPPIIENTPEEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41449

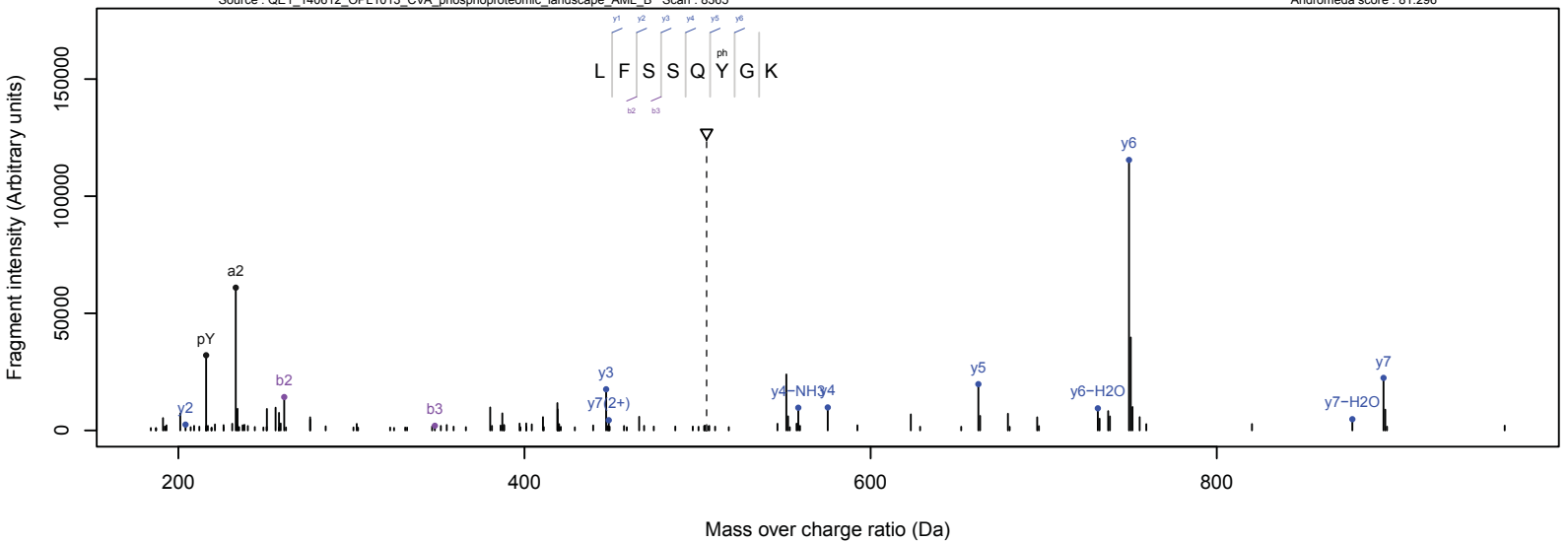
Andromeda score : 81.315



**LFSSQY(ph)GK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 8365

Andromeda score : 81.296

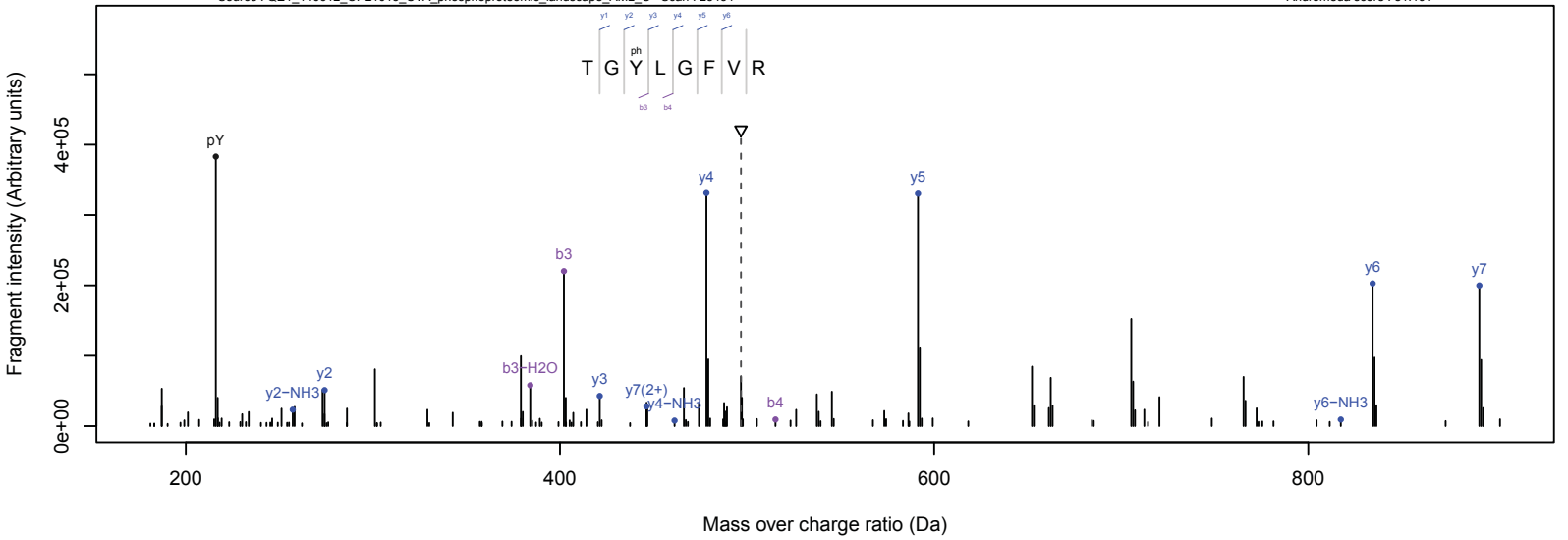




TGY(ph)LGFVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25434

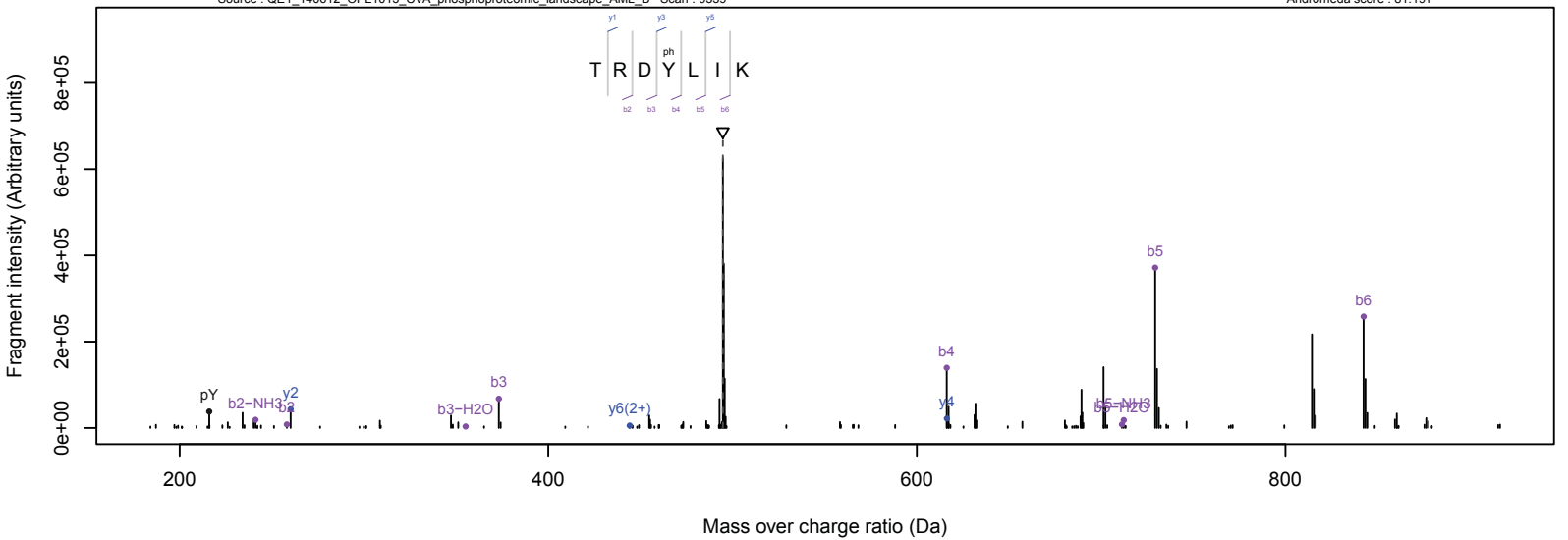
Andromeda score : 81.191



TRDY(ph)LIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9339

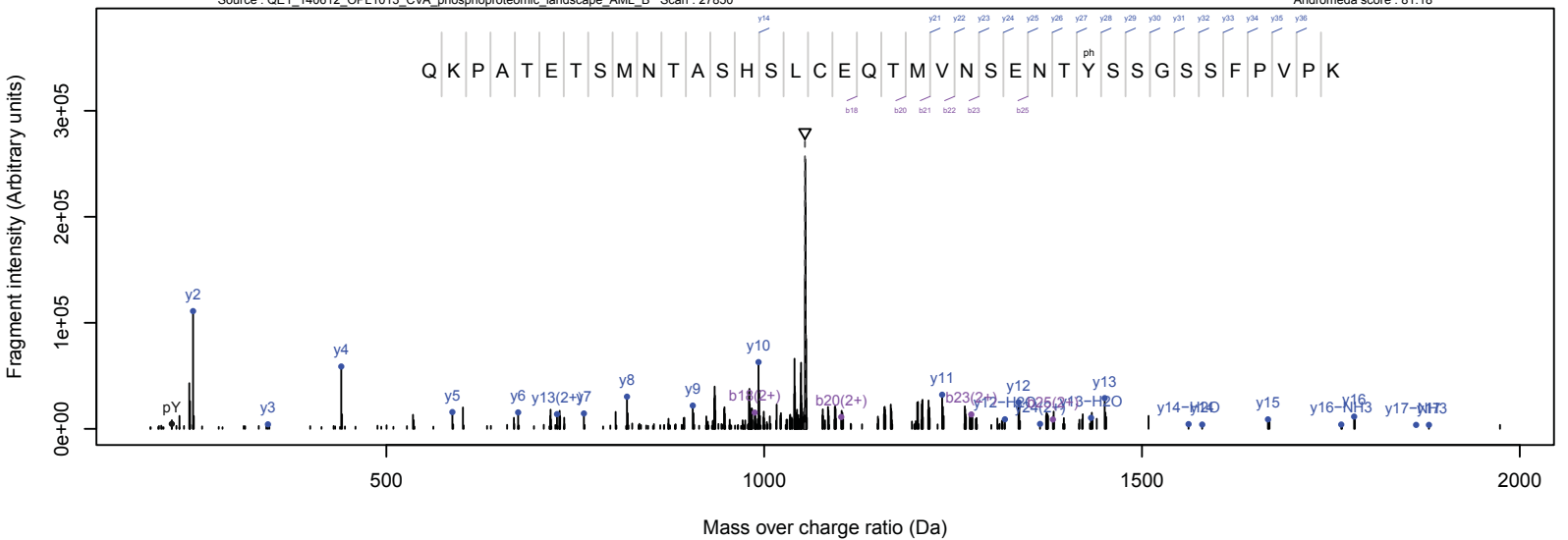
Andromeda score : 81.191



QKPATETSMNTASHSLCEQTMVNSENTY(ph)SSGSSFPVK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27850

Andromeda score : 81.18

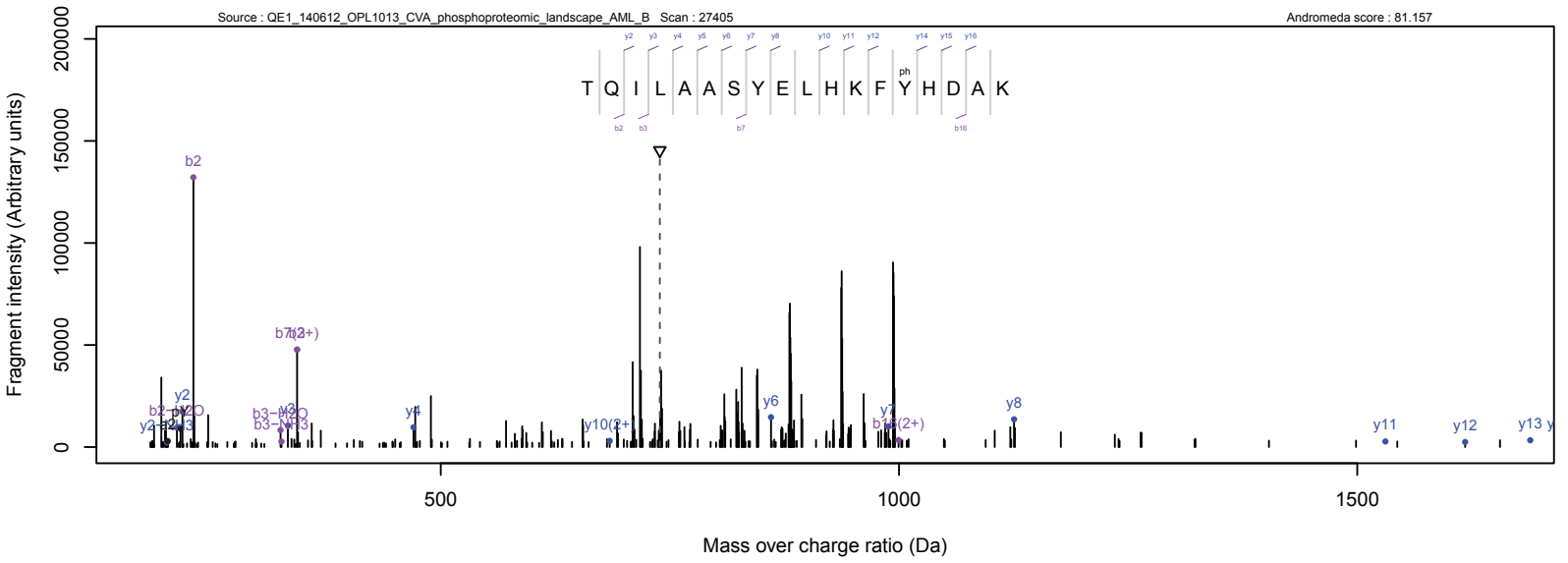




TQILAASYELHKFY(ph)HDAK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27405

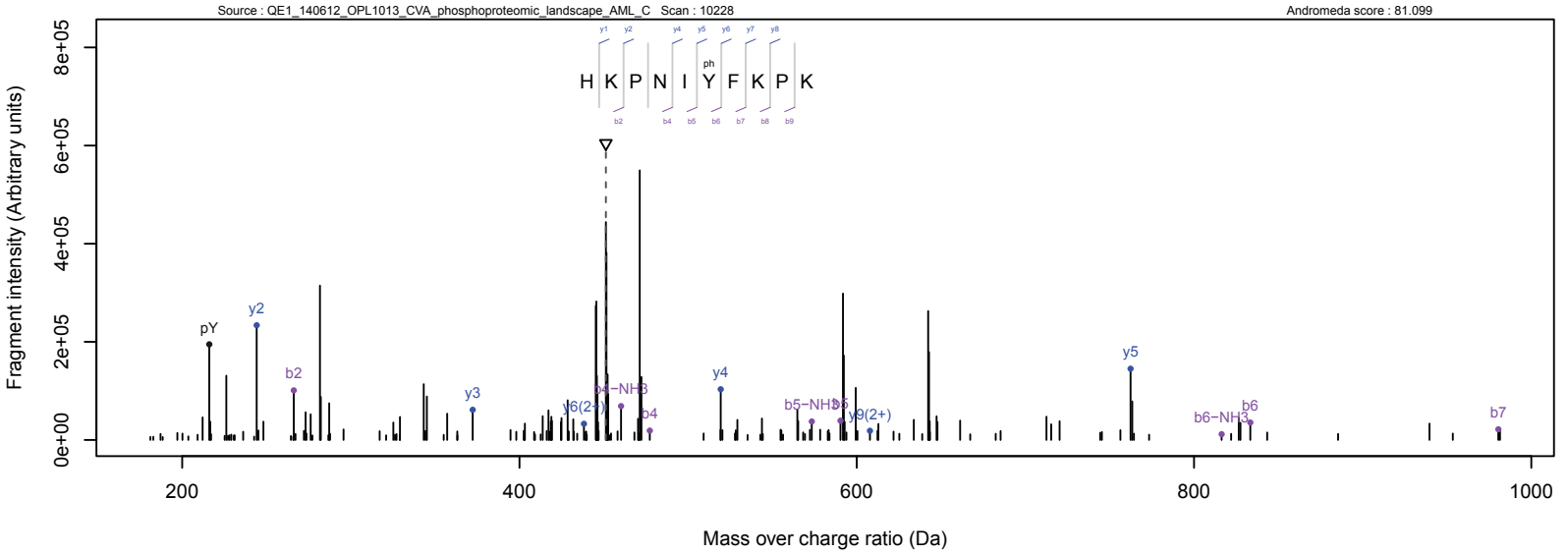
Andromeda score: 81.157



HKPNY(ph)FKPK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10228

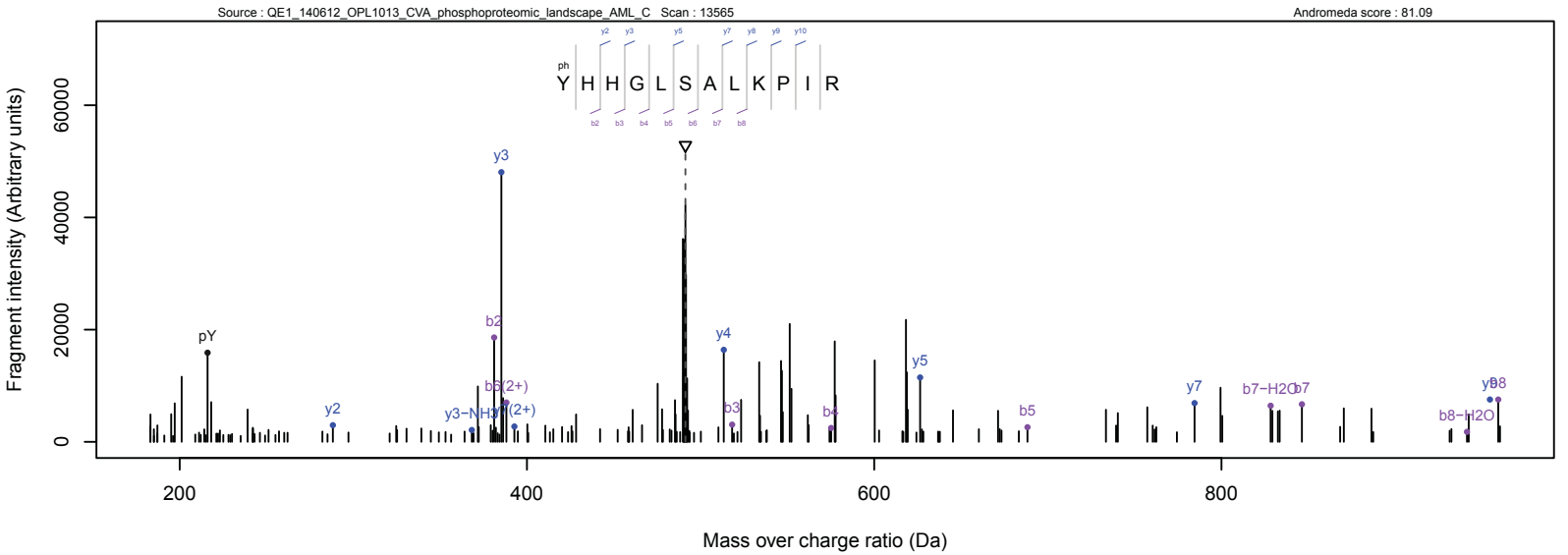
Andromeda score: 81.099



Y(ph)HHGLSALKPIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13565

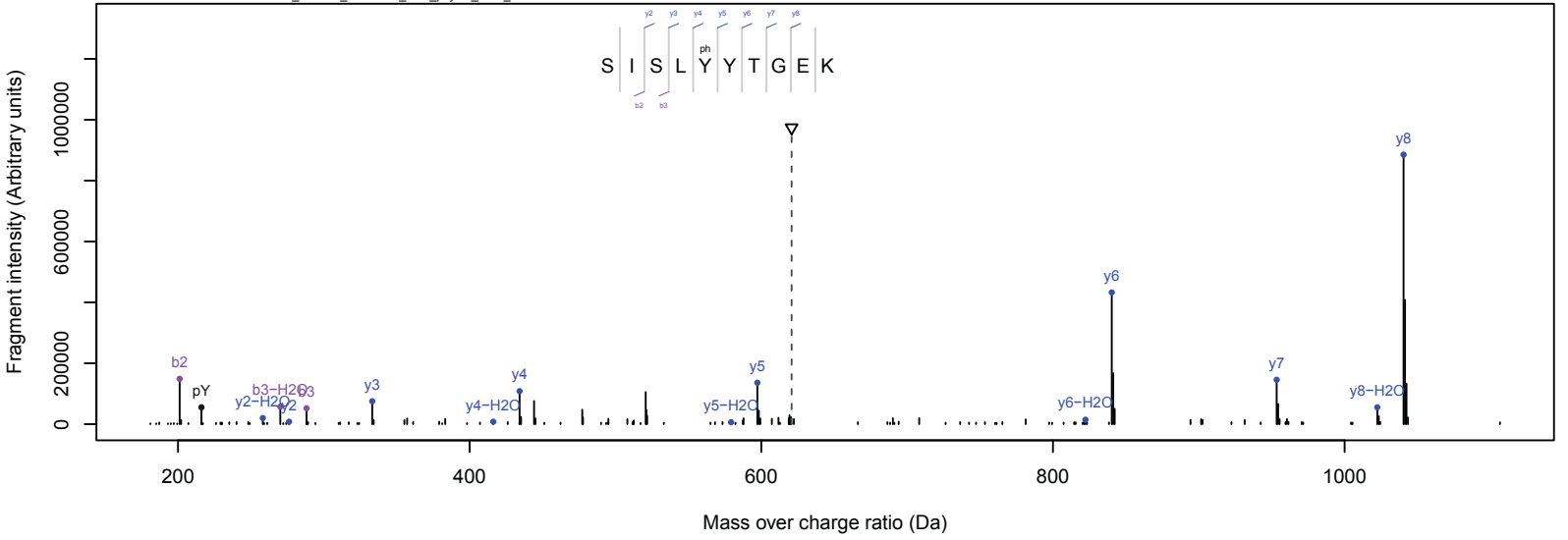
Andromeda score: 81.09



SISLY(ph)YTGEK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23890

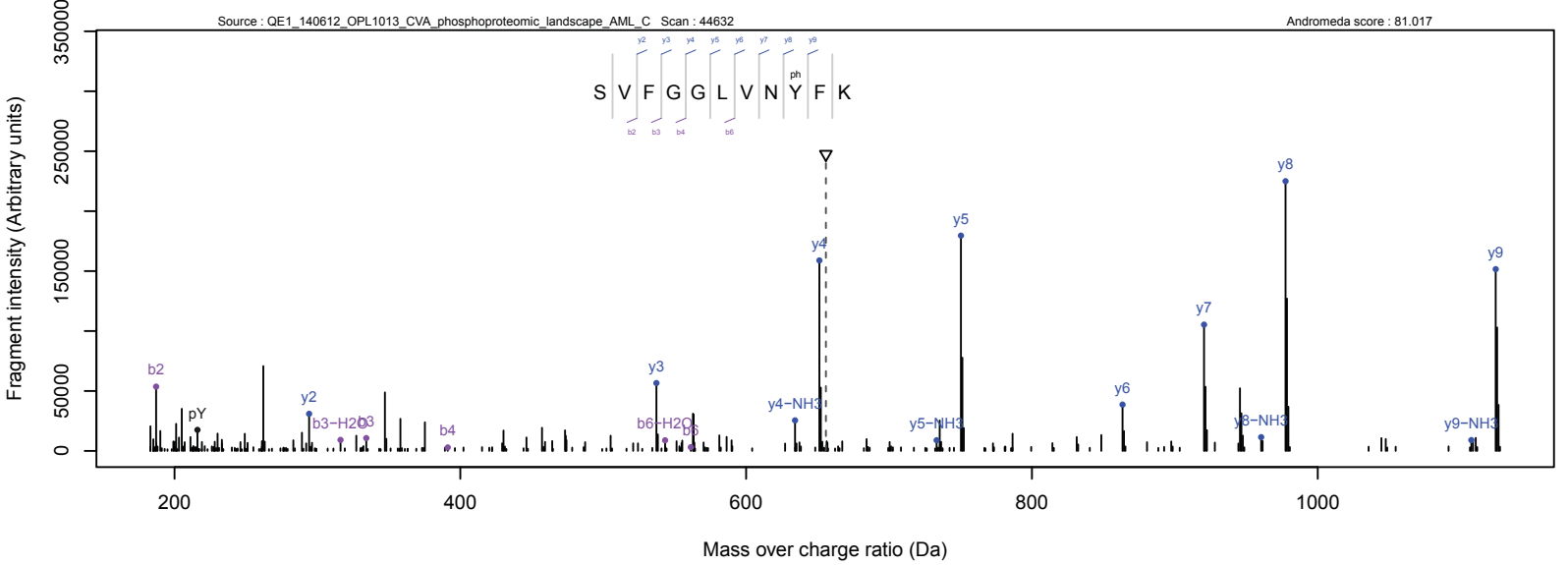
Andromeda score : 81.017



SVFGGLVNY(ph)FK-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 44632

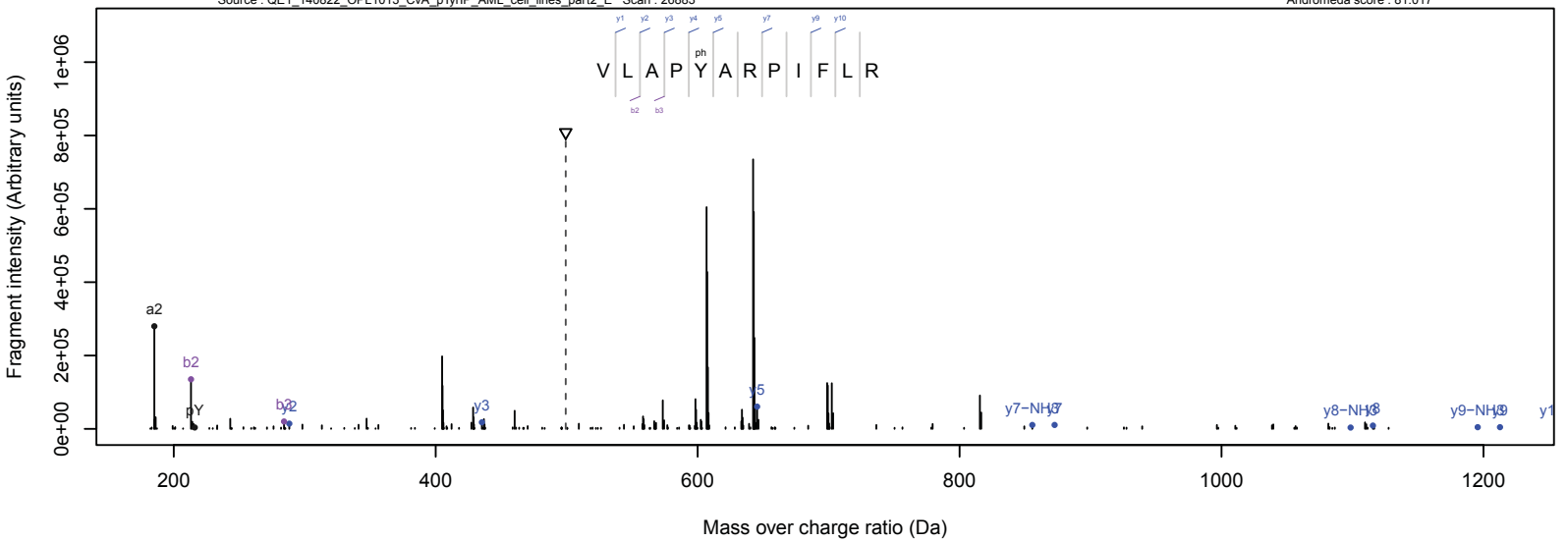
Andromeda score : 81.017



VLAPY(ph)ARPIFLR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 26883

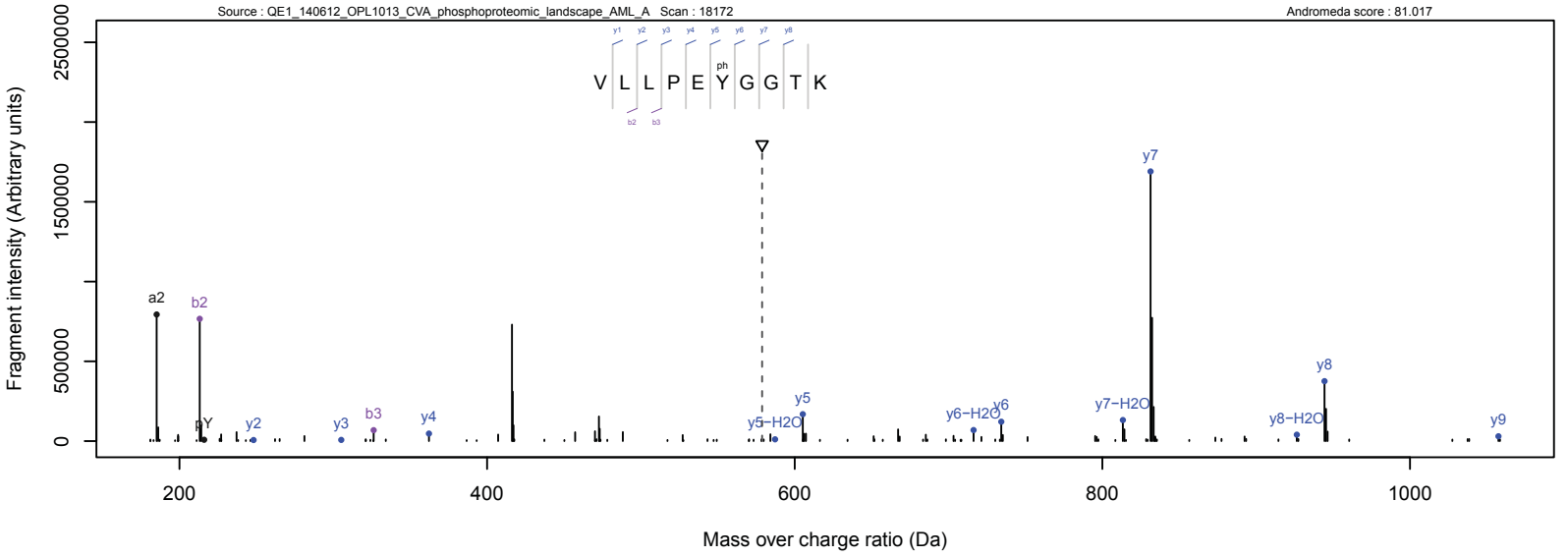
Andromeda score : 81.017



VLLPEY(ph)GGTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 18172

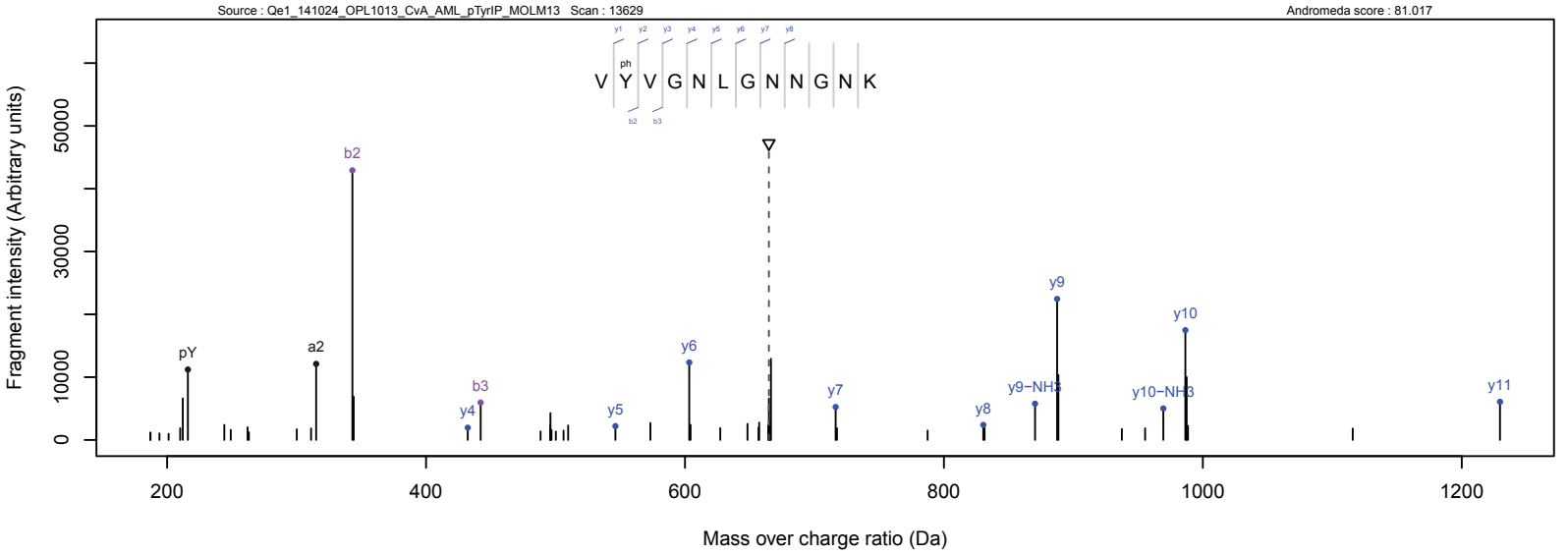
Andromeda score : 81.017



VY(ph)VGNLGNGNK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyriP\_MOLM13 Scan : 13629

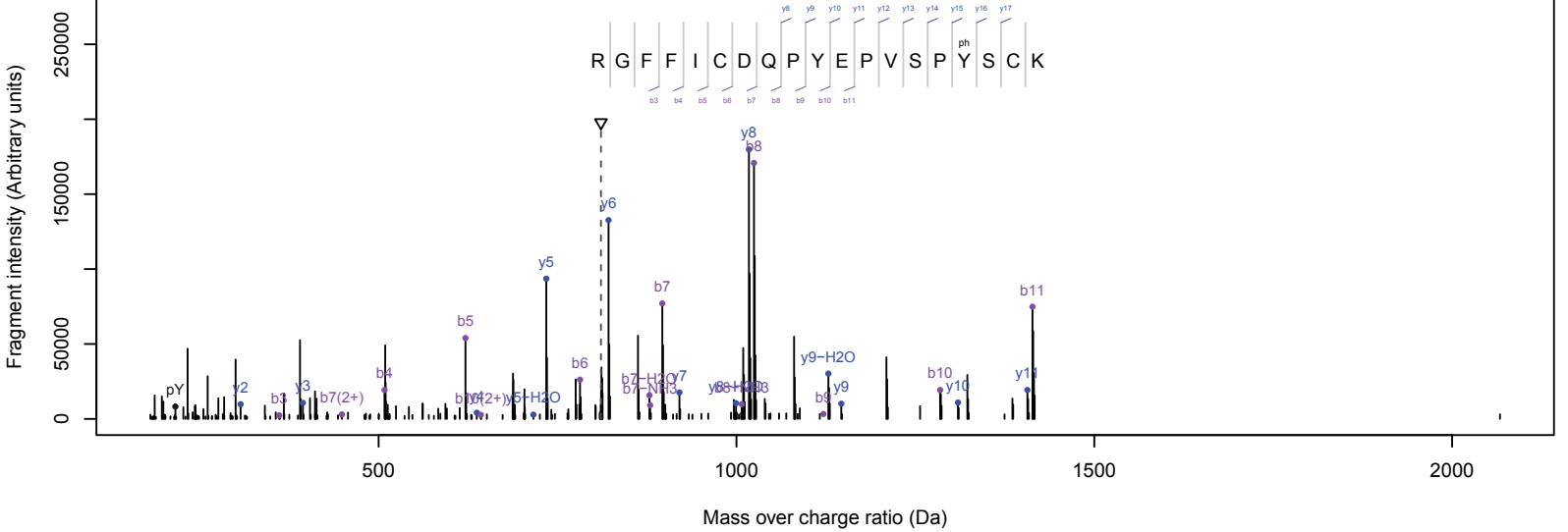
Andromeda score : 81.017



RGFFCDQPYEPVSPY(ph)SCK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34405

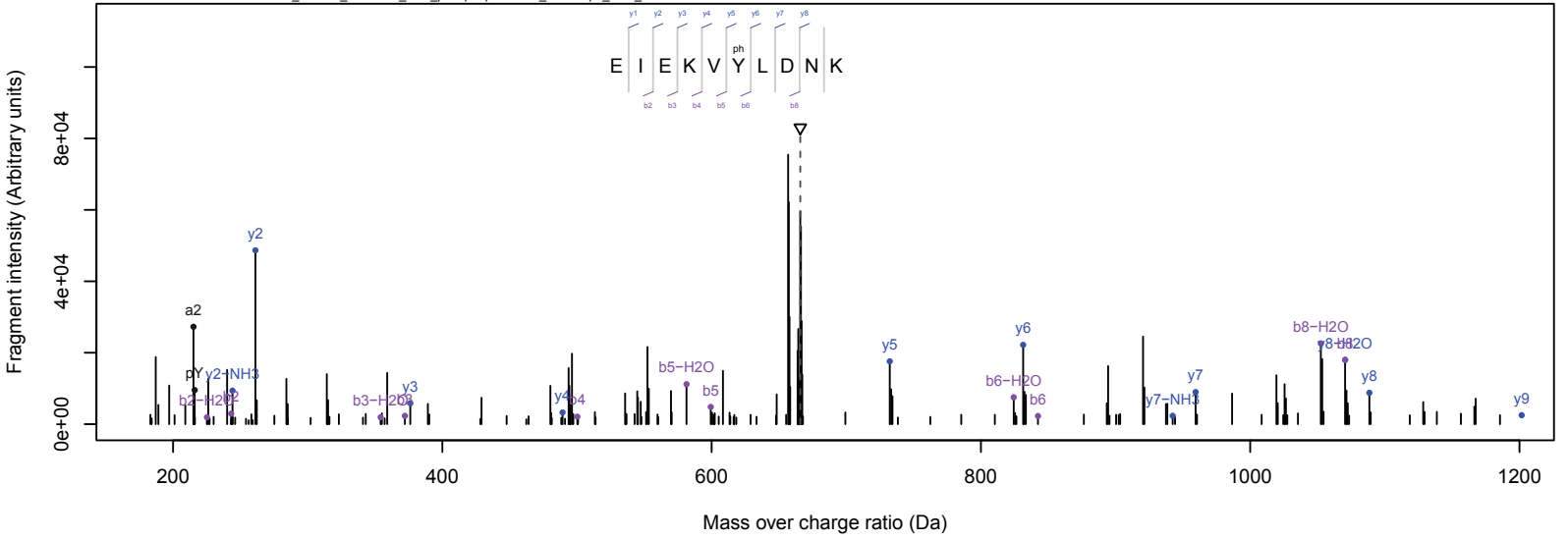
Andromeda score : 81.016



EIEKVY(ph)LDNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12927

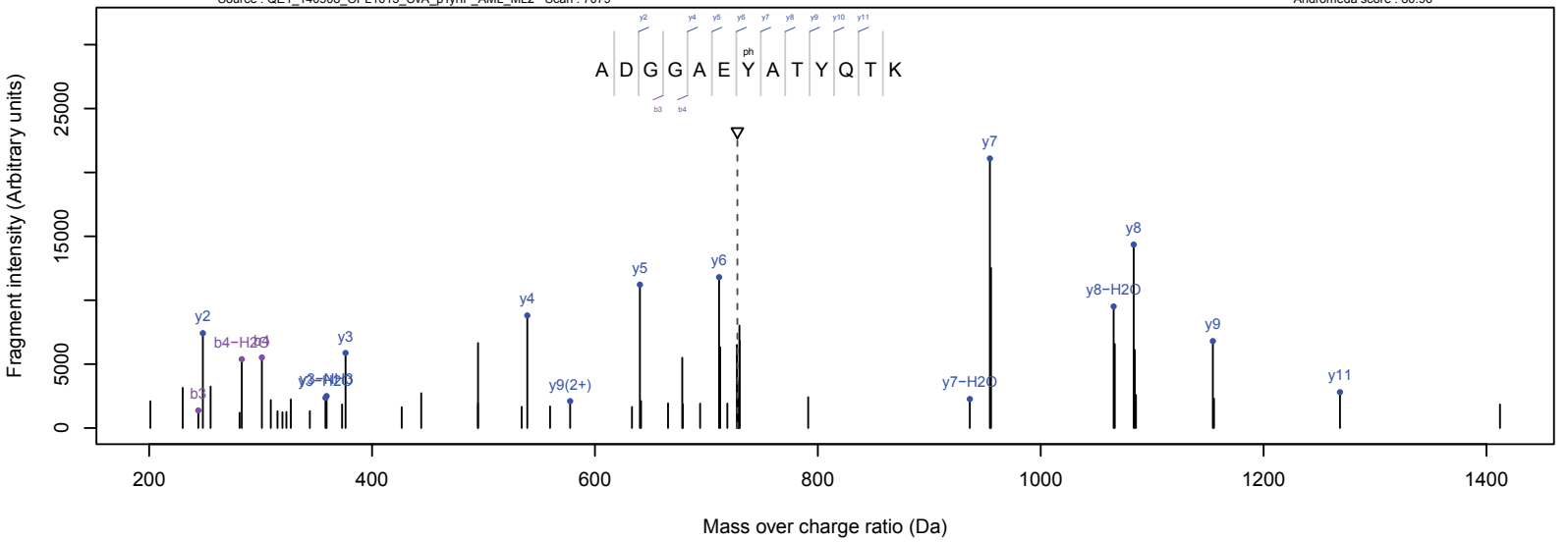
Andromeda score : 81.002



ADGGAEY(ph)ATYQTK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 7079

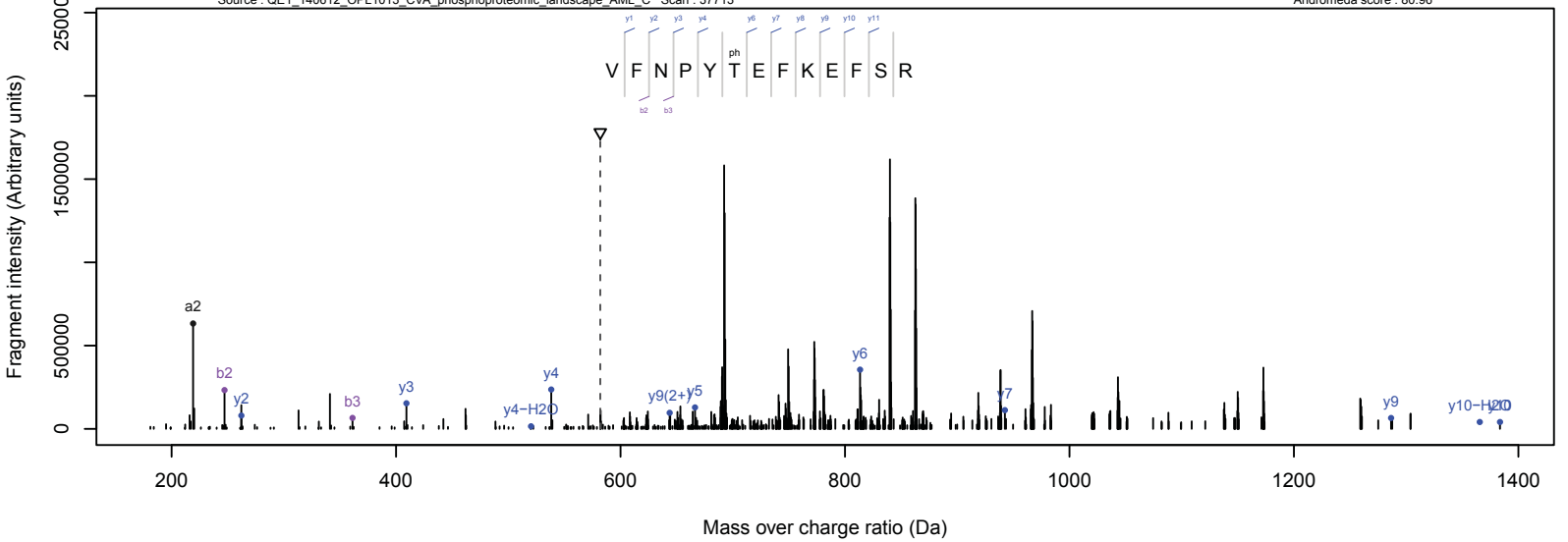
Andromeda score : 80.96



VFNPY(ph)EFKFSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37713

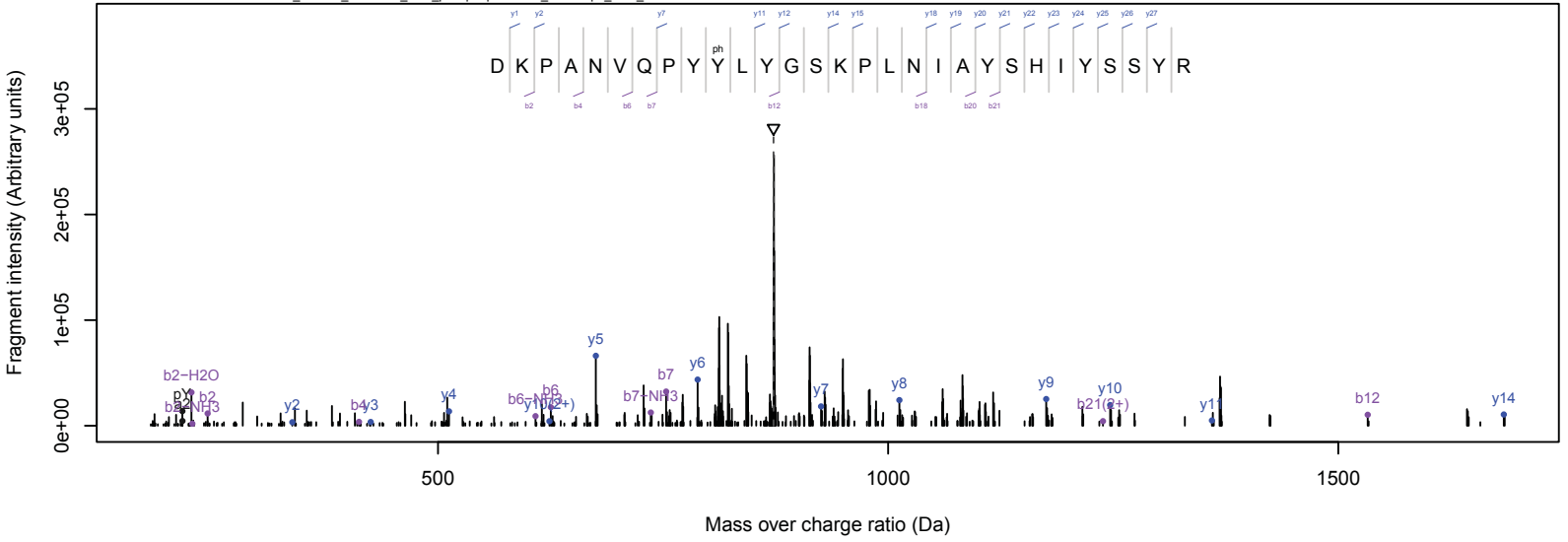
Andromeda score : 80.96



DKPANVQPY(ph)LYGSKPLNIAYSHIYSSYR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38583

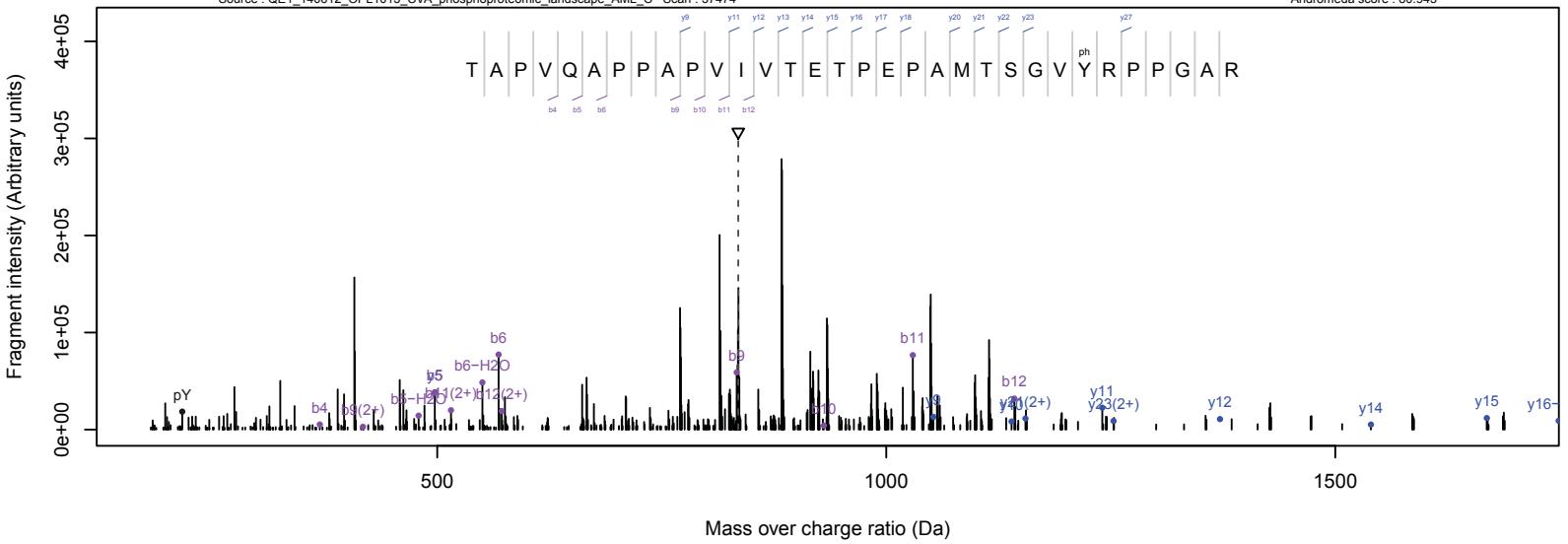
Andromeda score : 80.943



TAPVQAPPAPVIVTETPEPAMTSGVY(ph)RPPGAR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37474

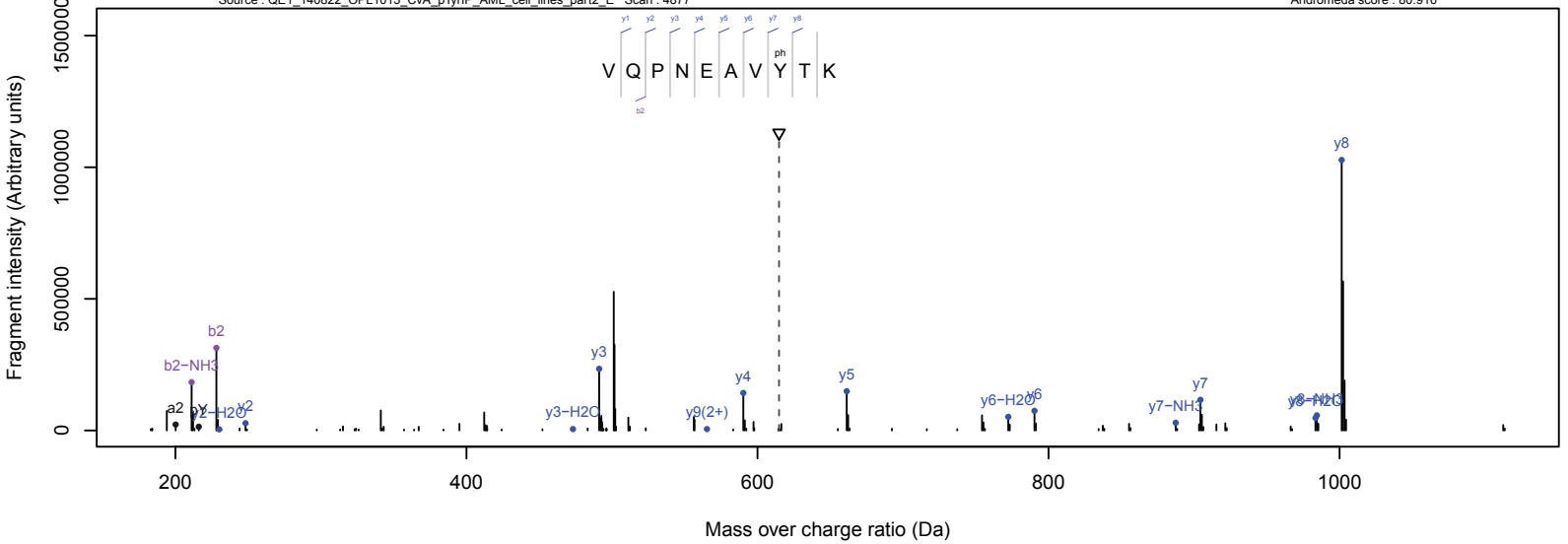
Andromeda score : 80.943

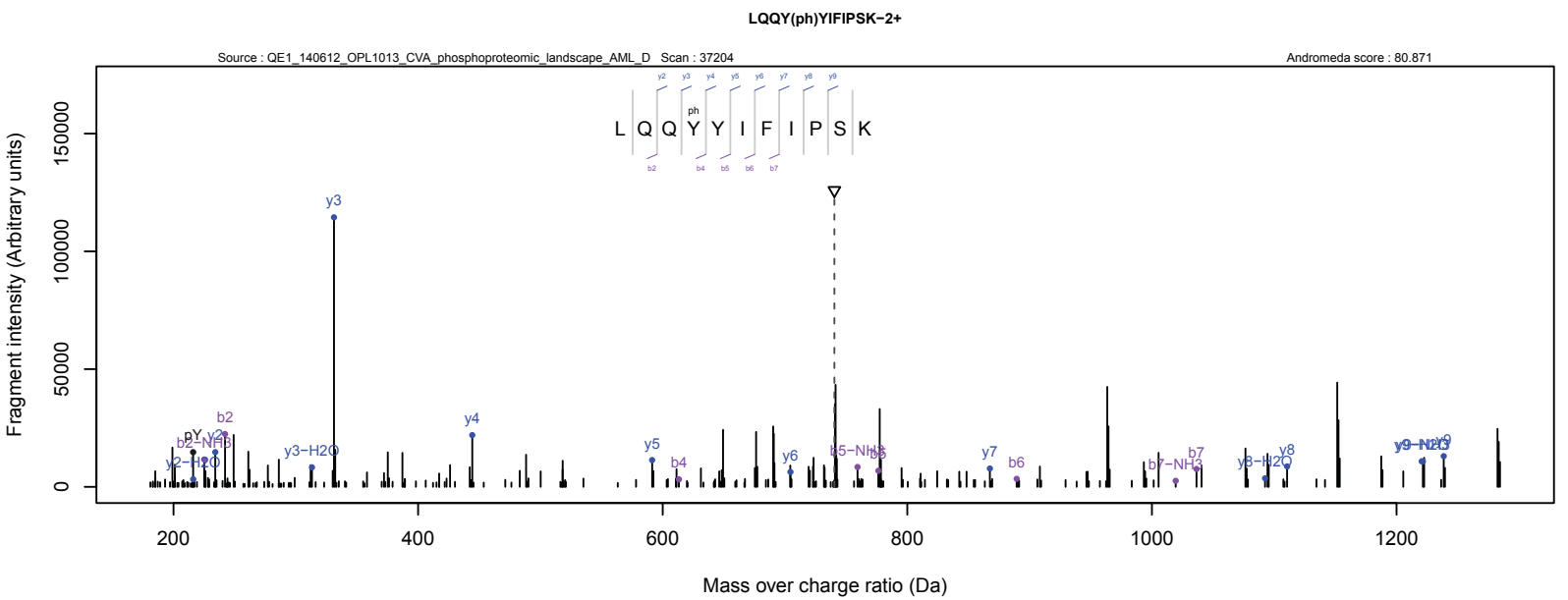
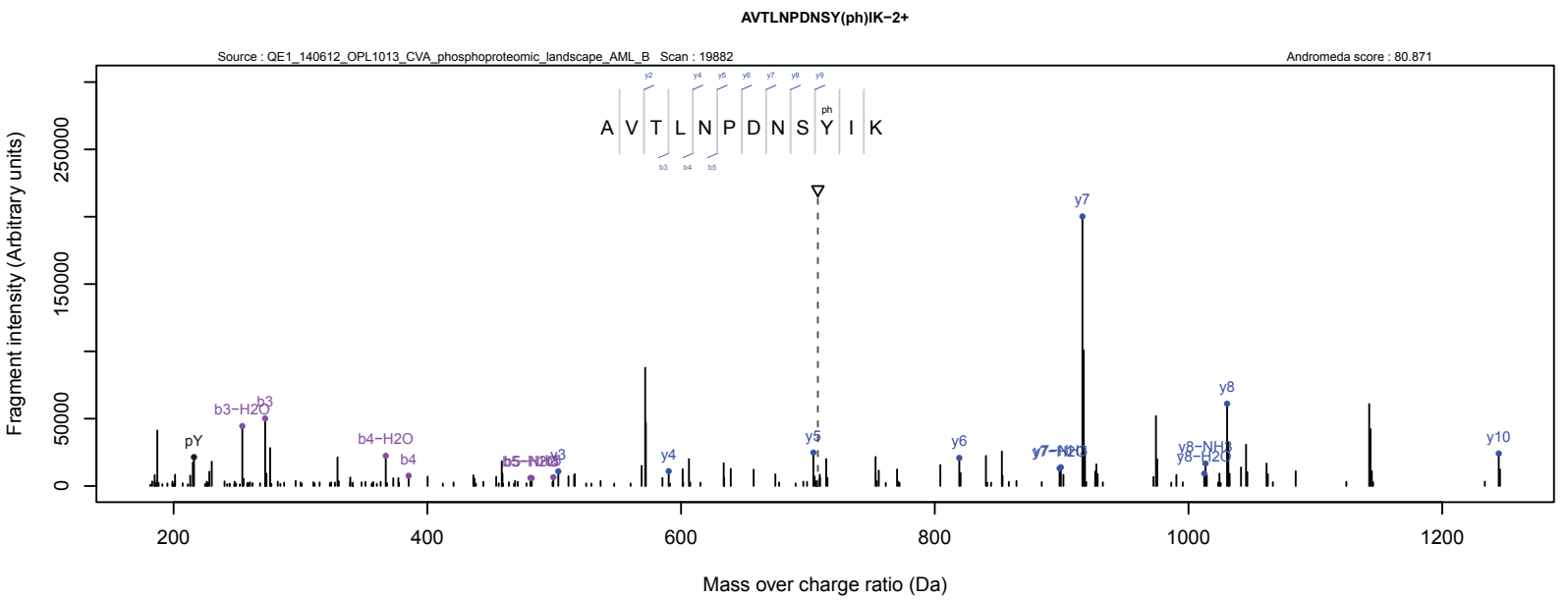
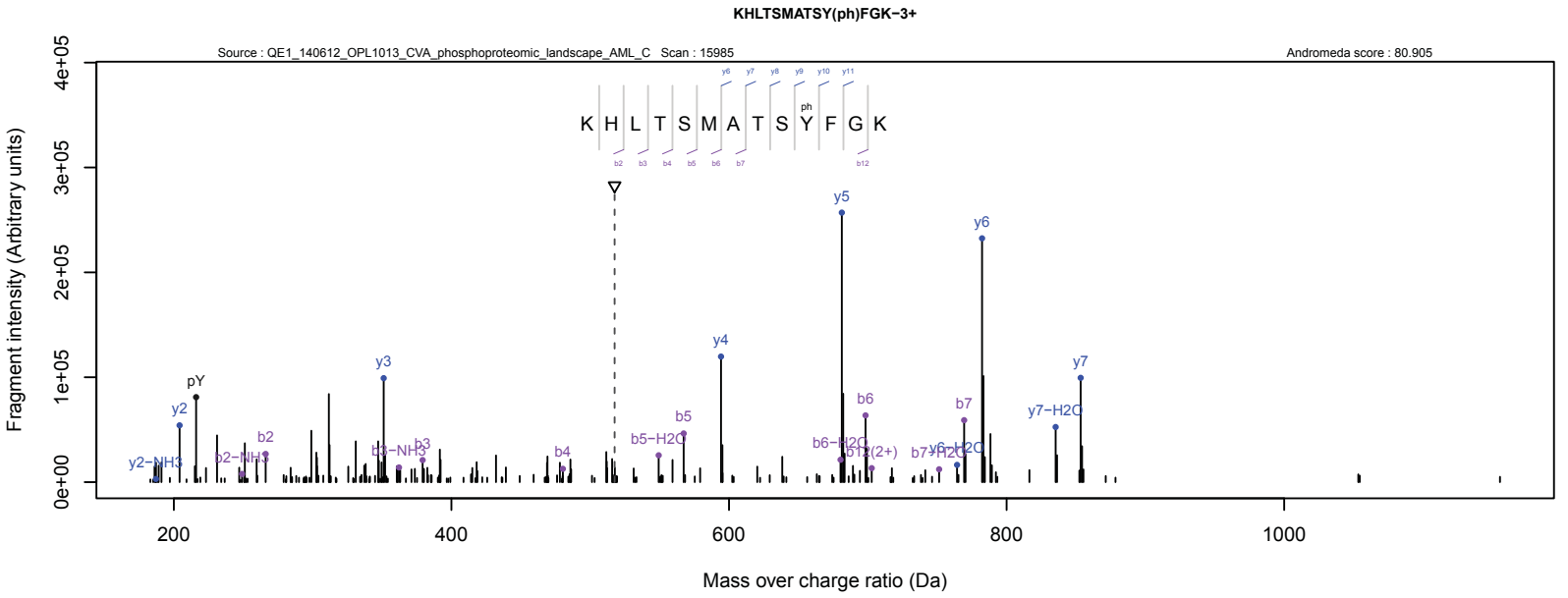


VQPNEAVYTK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 4877

Andromeda score : 80.916



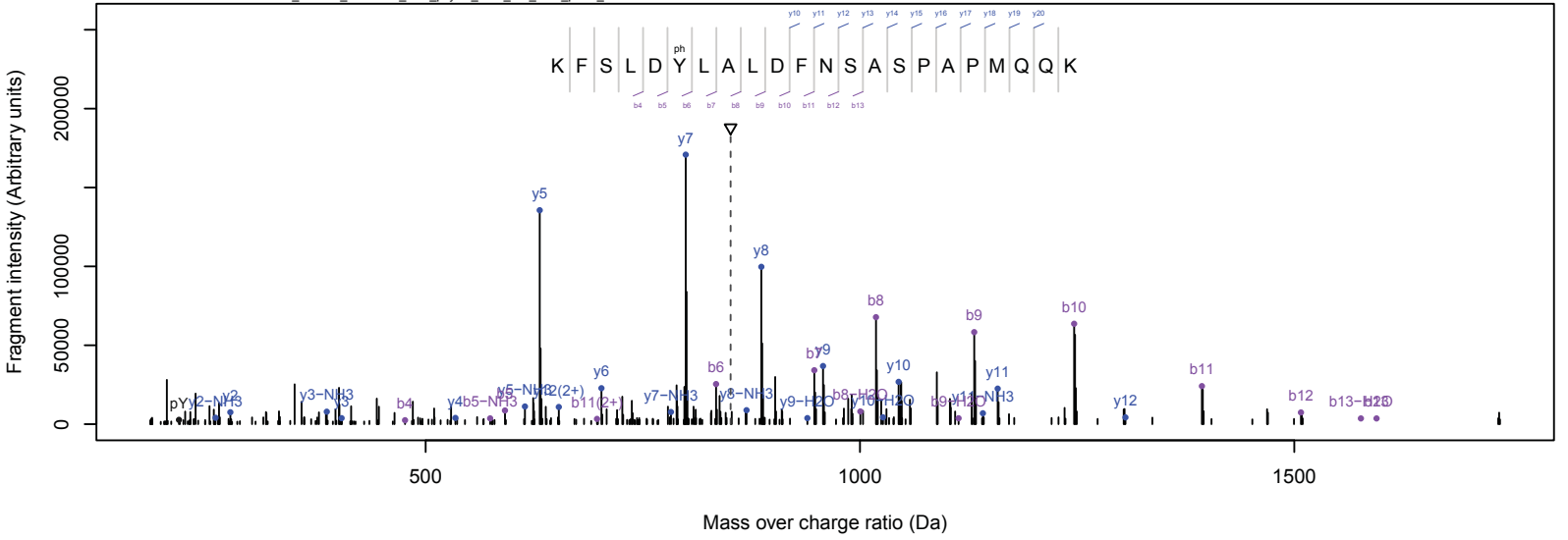




KFSLDY(ph)LALDFNSASPAPMQQK-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 42157

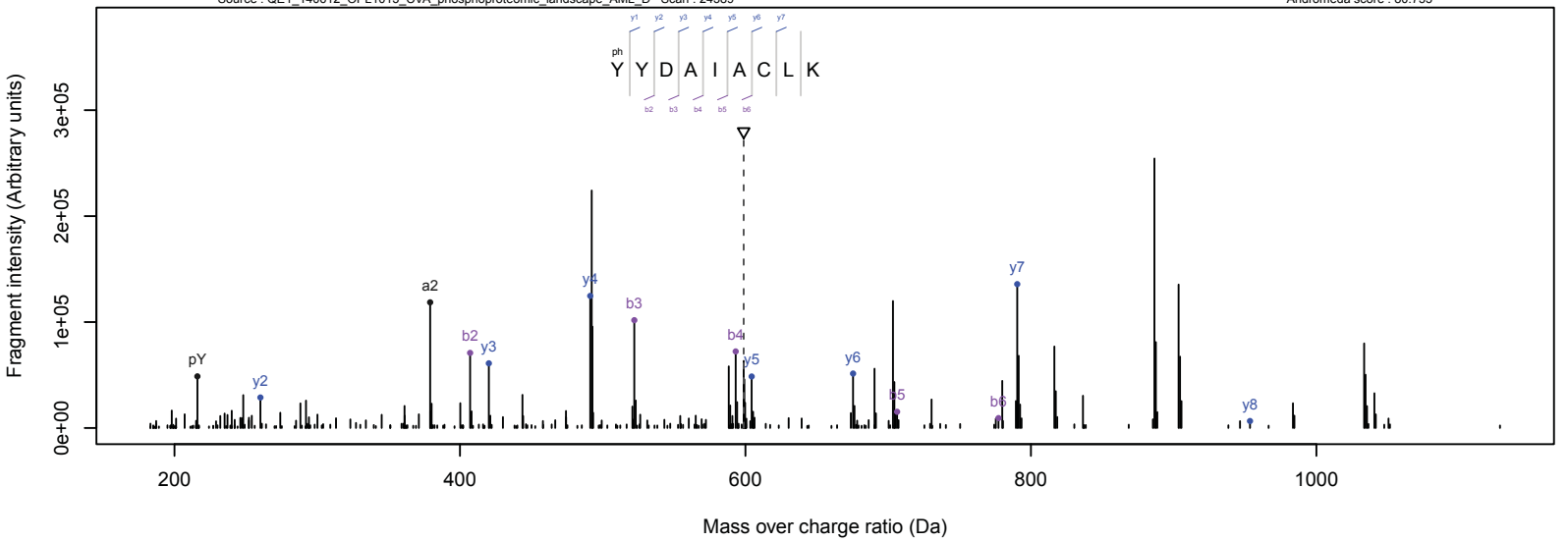
Andromeda score : 80.844



Y(ph)YDAIACLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 24389

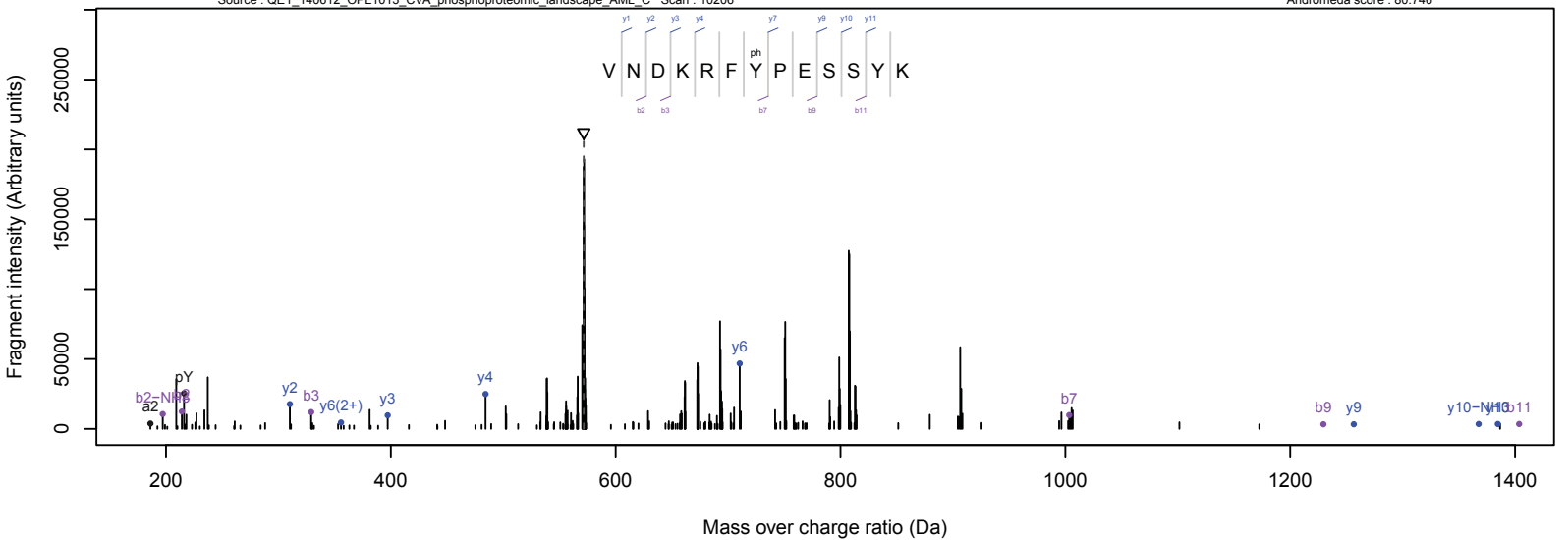
Andromeda score : 80.755



VNDKRFY(ph)PESSYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10206

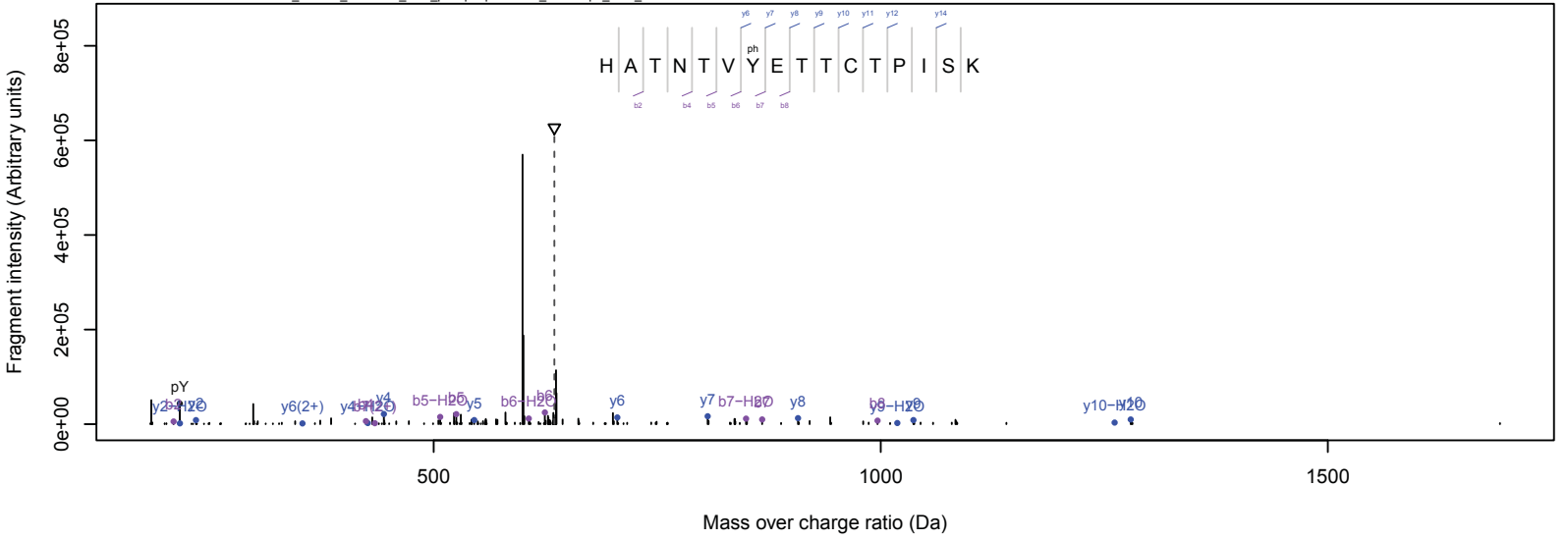
Andromeda score : 80.746



HATNTVY(ph)ETTCTPIISK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11695

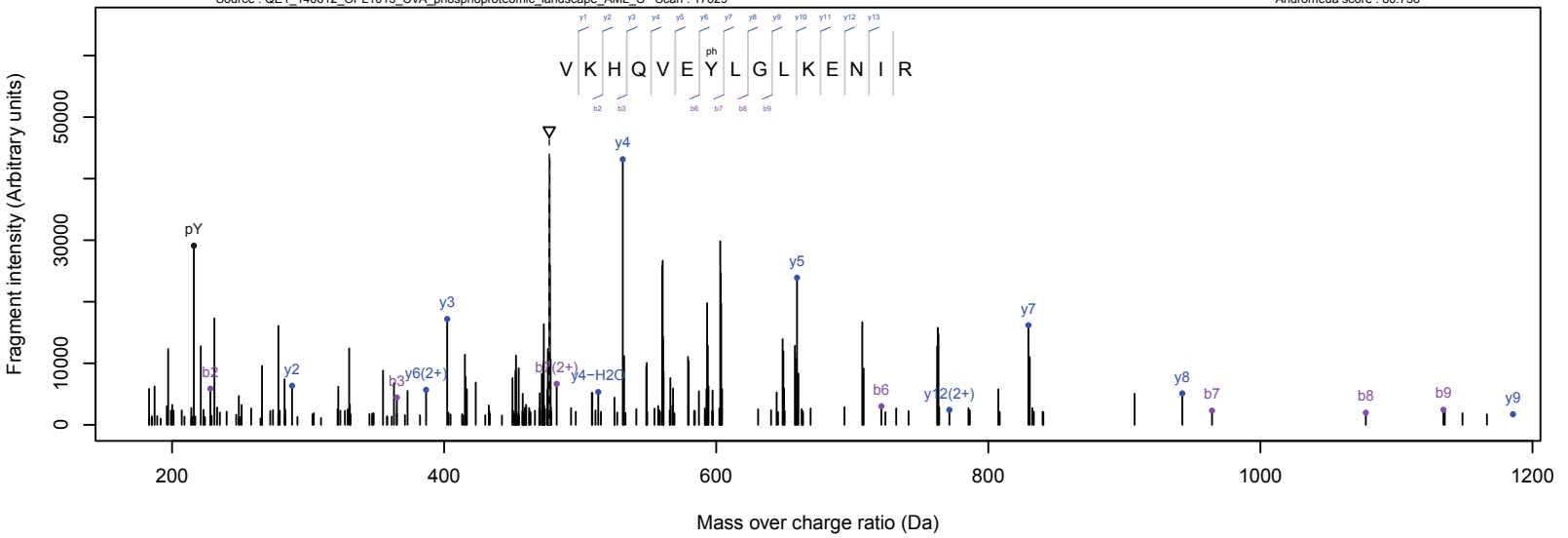
Andromeda score : 80.738



VKHQVEY(ph)LGLKENIR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17629

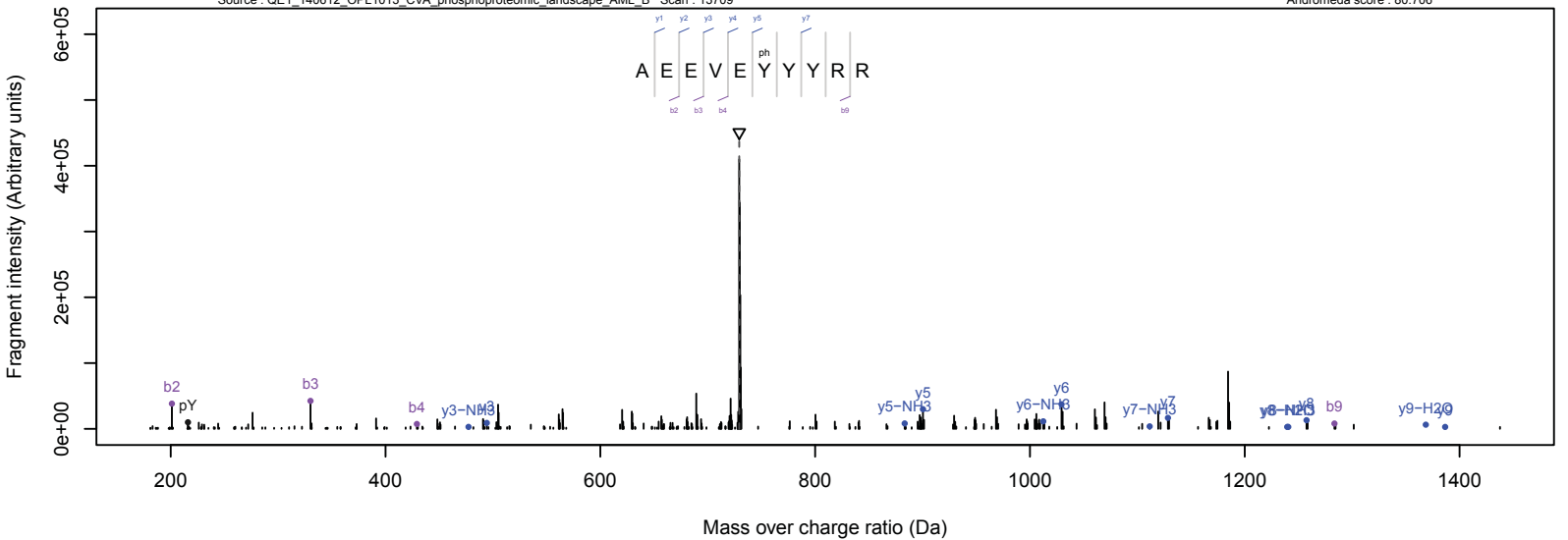
Andromeda score : 80.738



AEVEY(ph)YYRR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13709

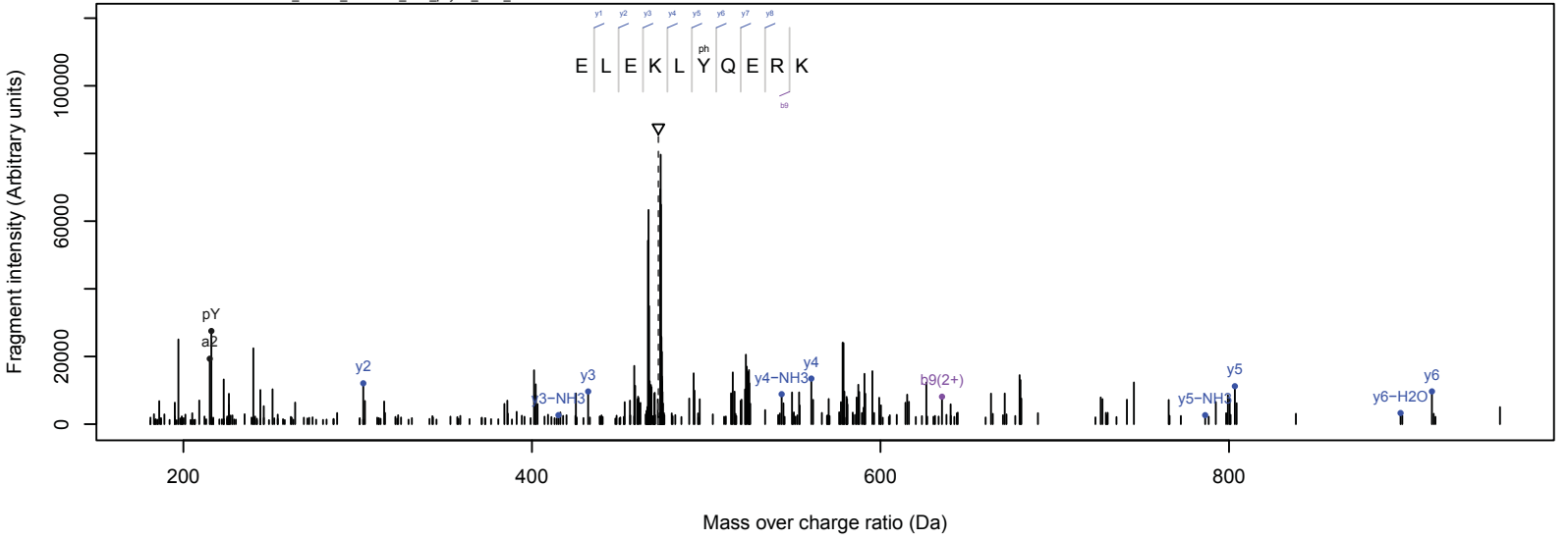
Andromeda score : 80.706



ELEKLY(ph)QERK-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HL60 Scan: 7622

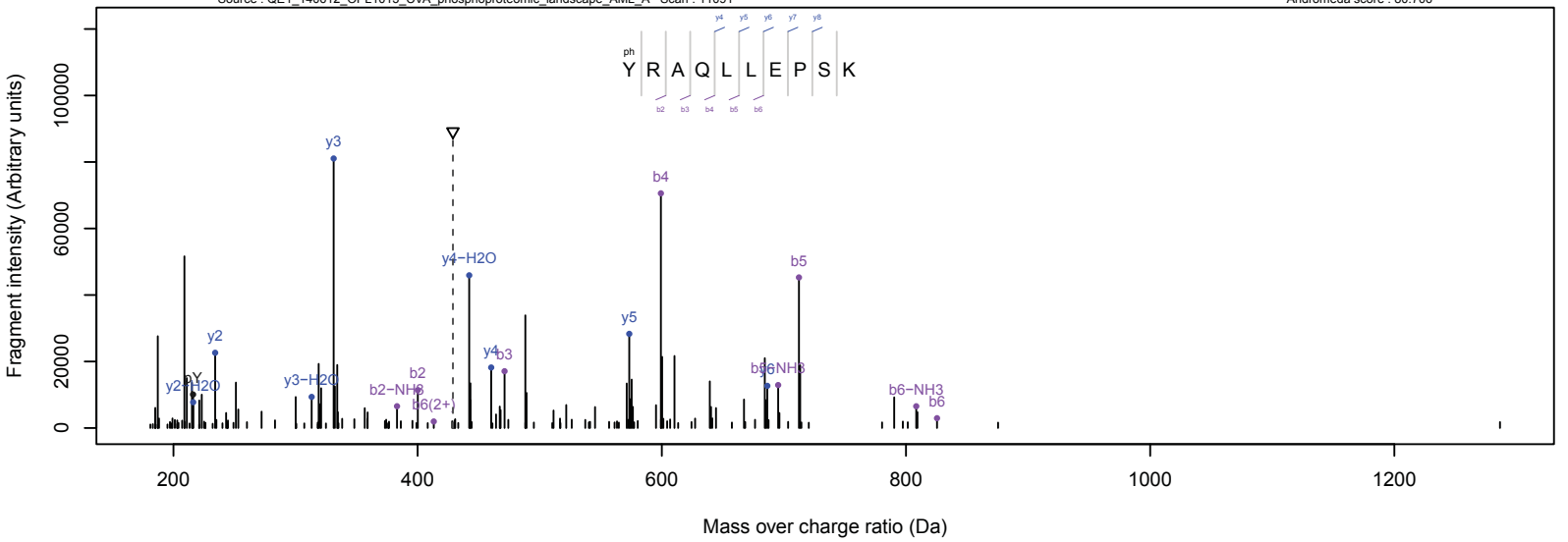
Andromeda score: 80.706



Y(ph)RAQLLEPSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 11091

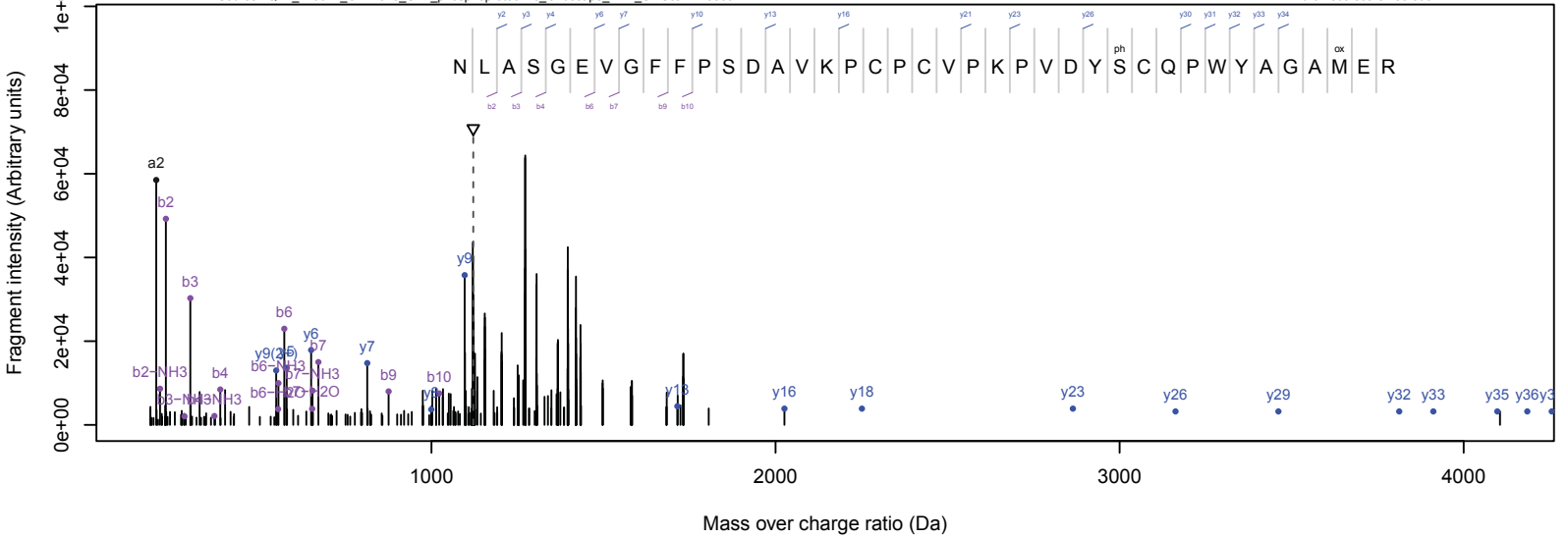
Andromeda score: 80.706



NLASGEVGFPPSDAVKPCPCVPKPVDS(ph)CQPWYAGAM(ox)ER-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 43509

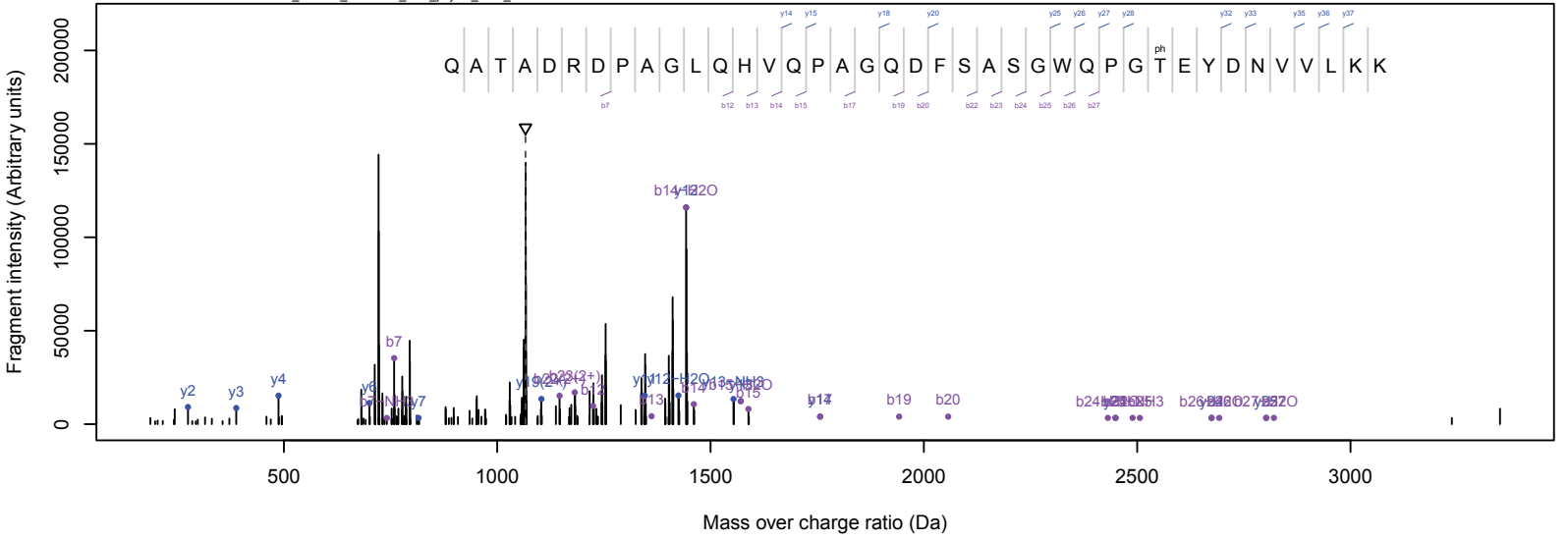
Andromeda score: 80.693



**QATADRDPAQLQHVQPAAGQDFASAGWQPGT(ph)EYDNNVVLKK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32501

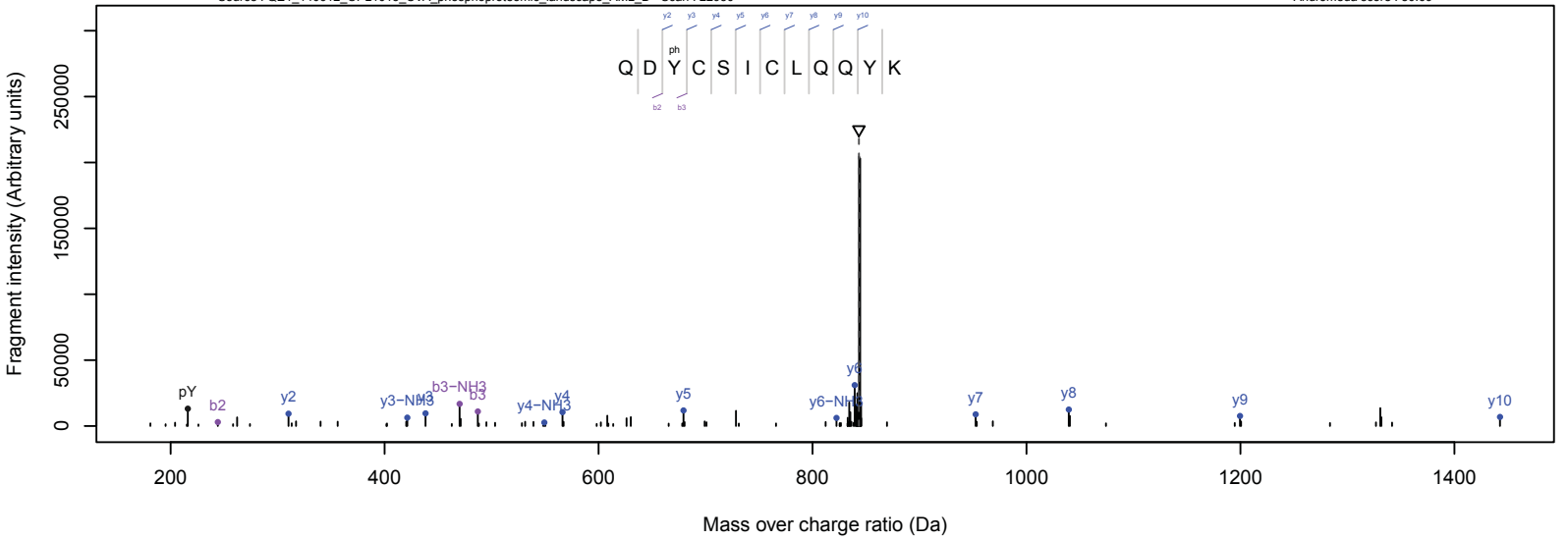
Andromeda score : 80.693



**QDY(ph)CSICLQYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22950

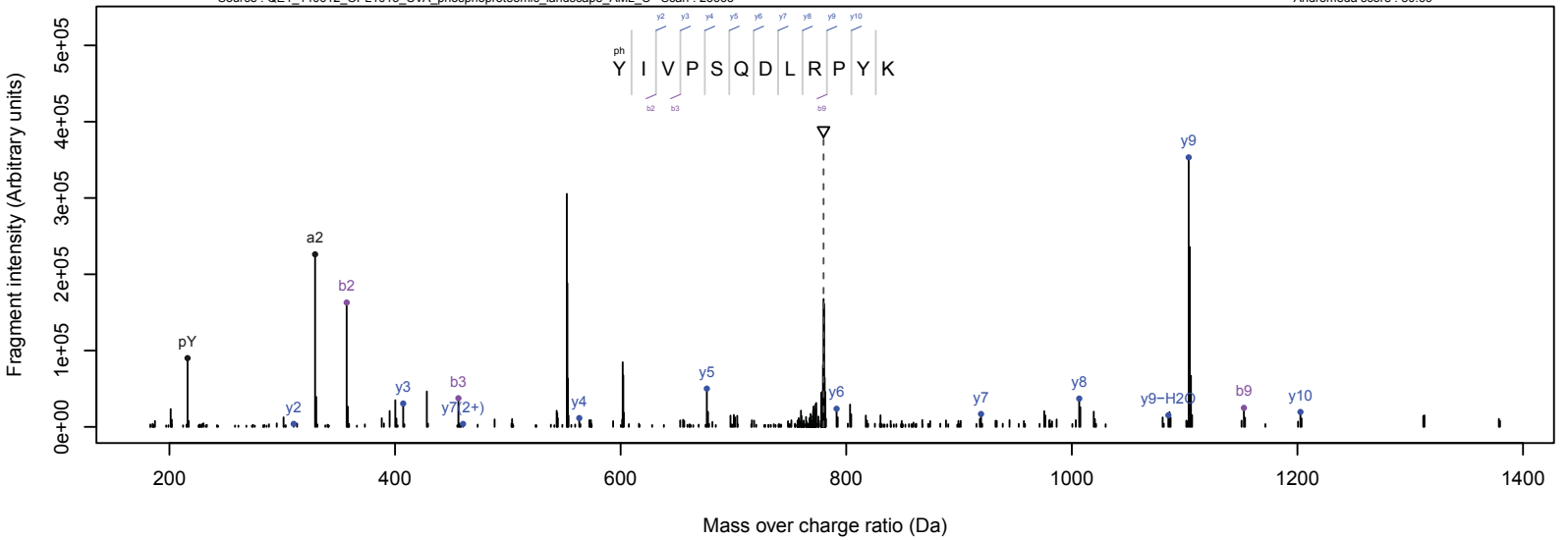
Andromeda score : 80.69



**Y(ph)IVPSQDLRPYK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23665

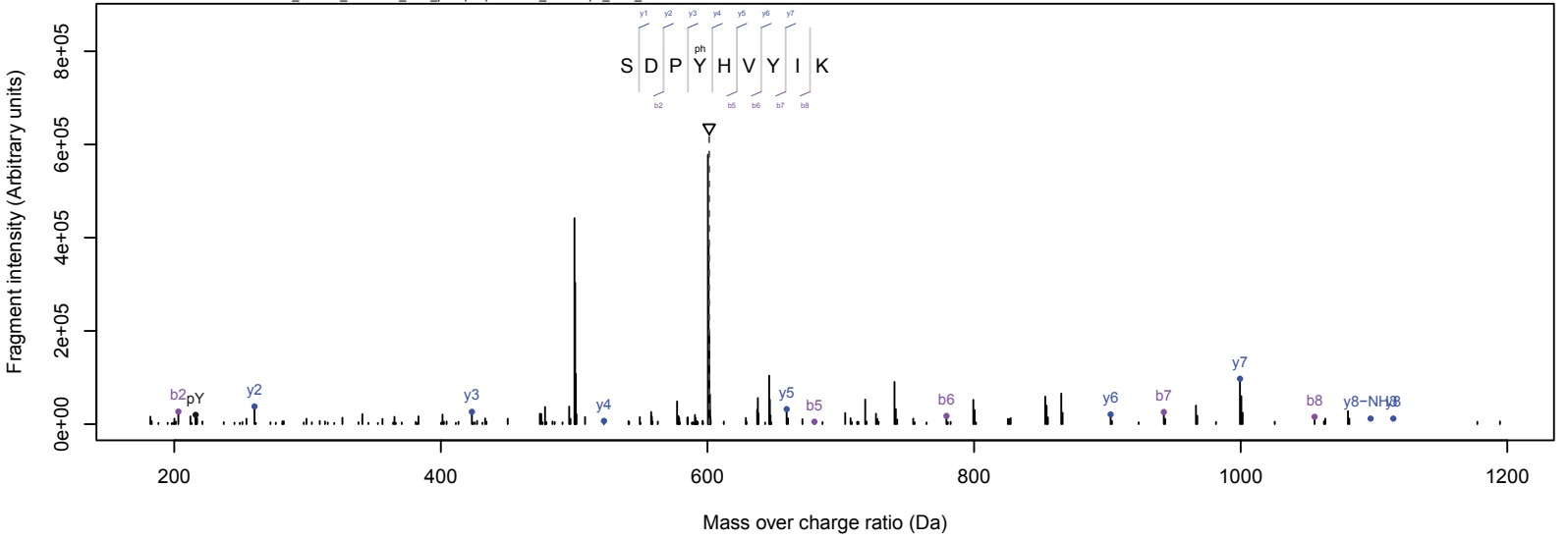
Andromeda score : 80.69



SDPY(ph)HVIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11405

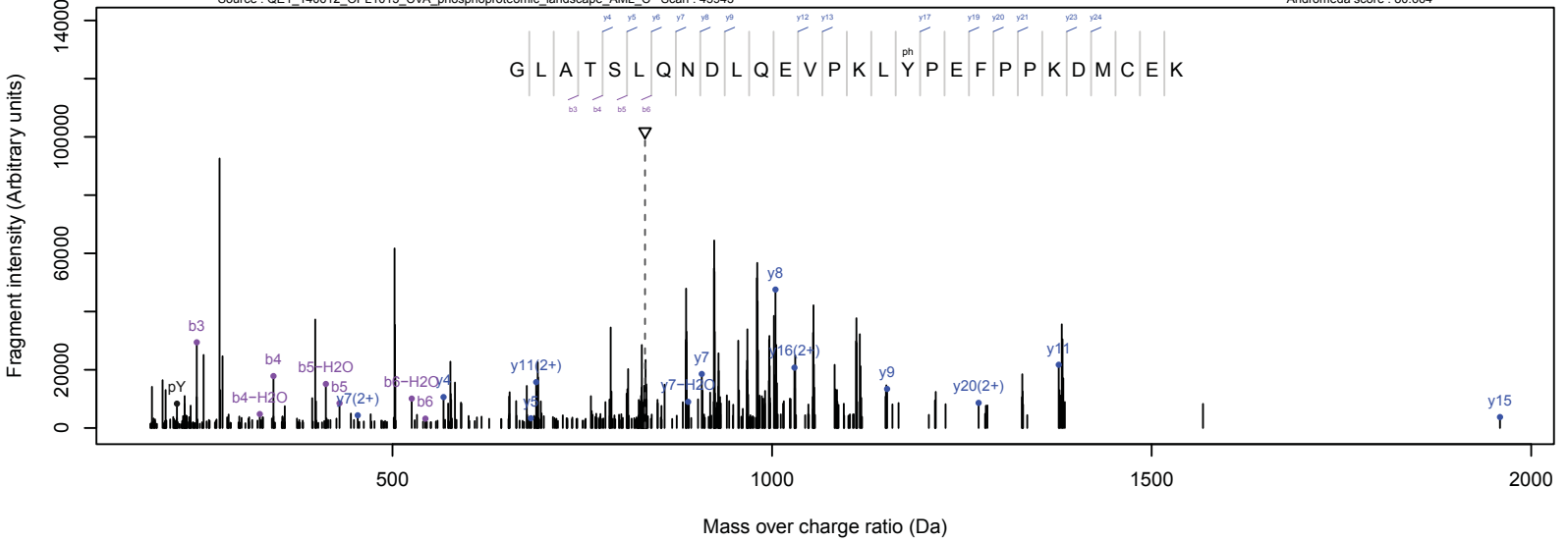
Andromeda score : 80.688



GLATSLQNDLQEVPKLYPEFPKDMCEK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45943

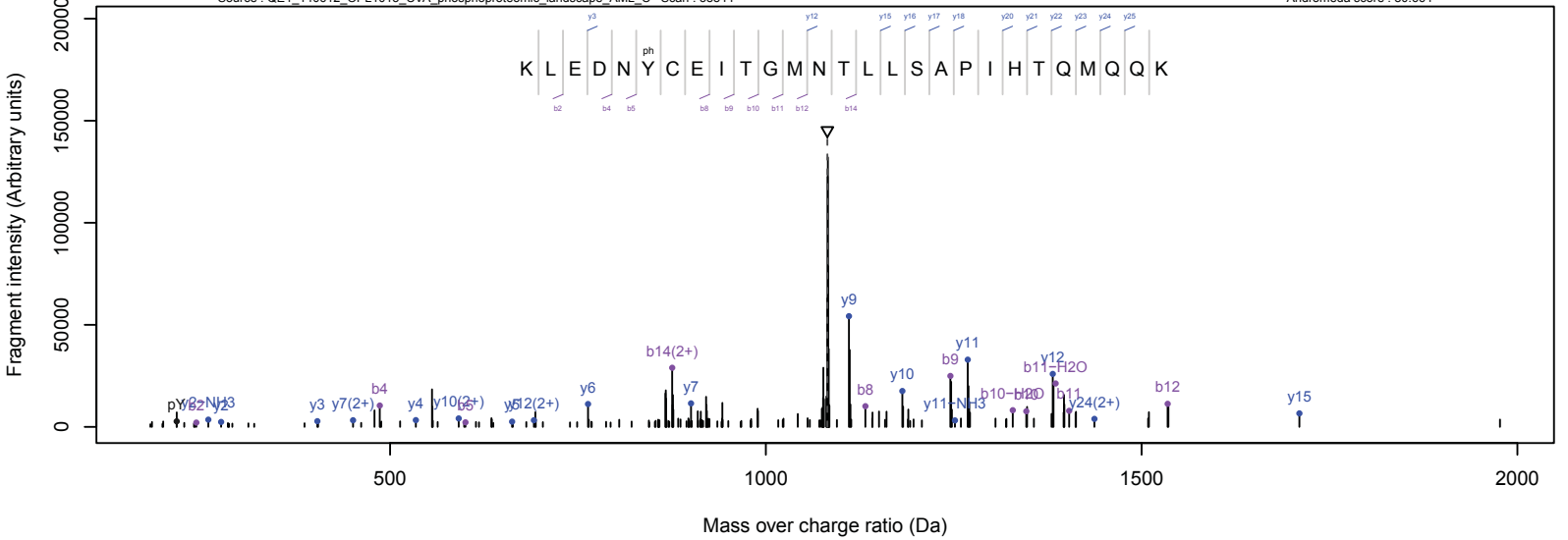
Andromeda score : 80.664



KLEDNY(ph)CEITGMNTLLSAPIHTQMQQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38844

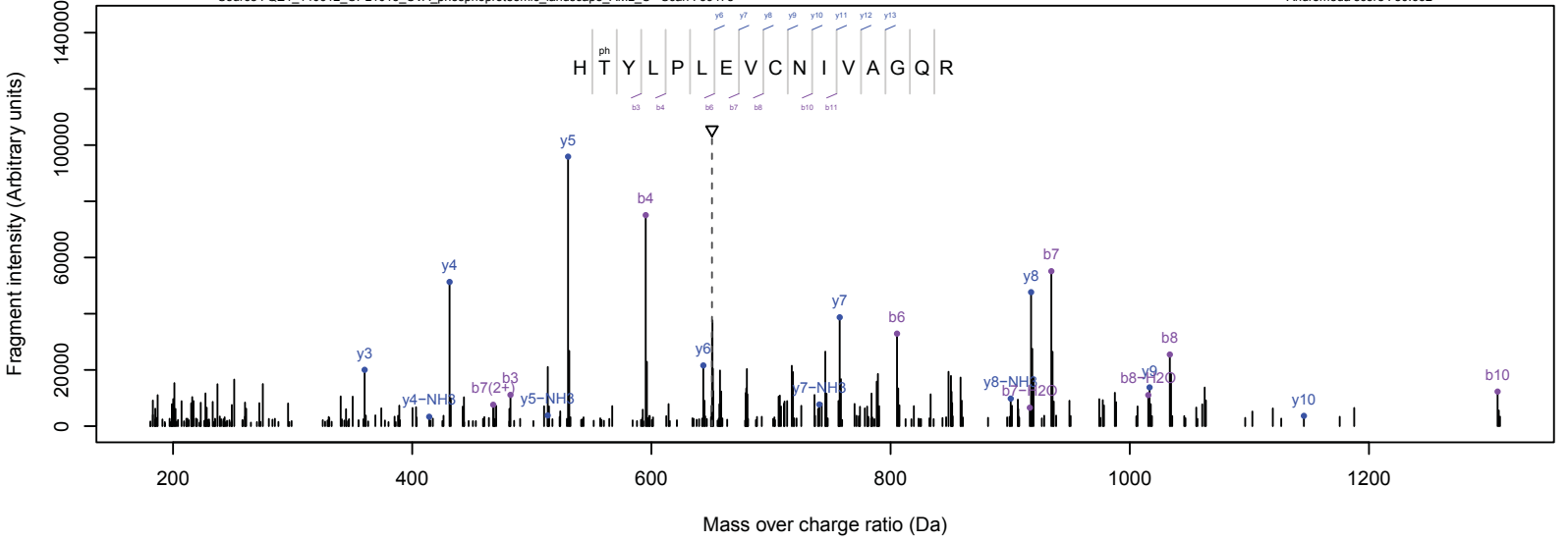
Andromeda score : 80.664



HT(ph)YLPLEVCNIVAGQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39478

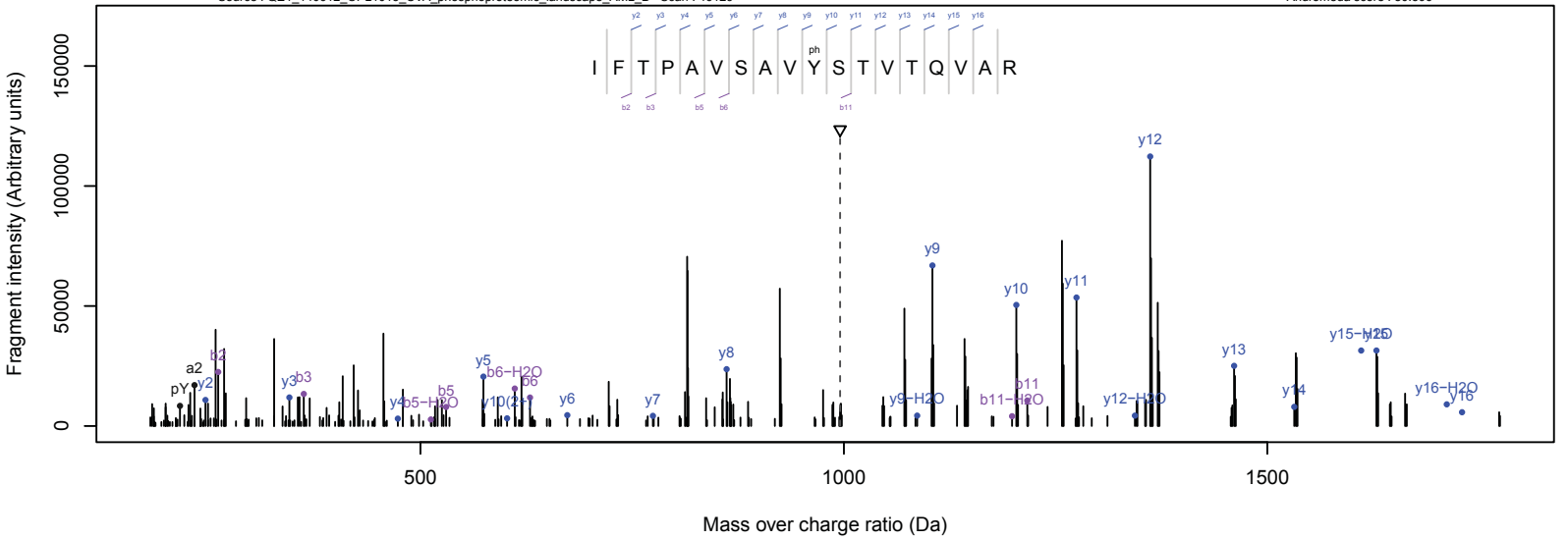
Andromeda score : 80.632



IFTPAVSAVY(ph)STVTQVAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46129

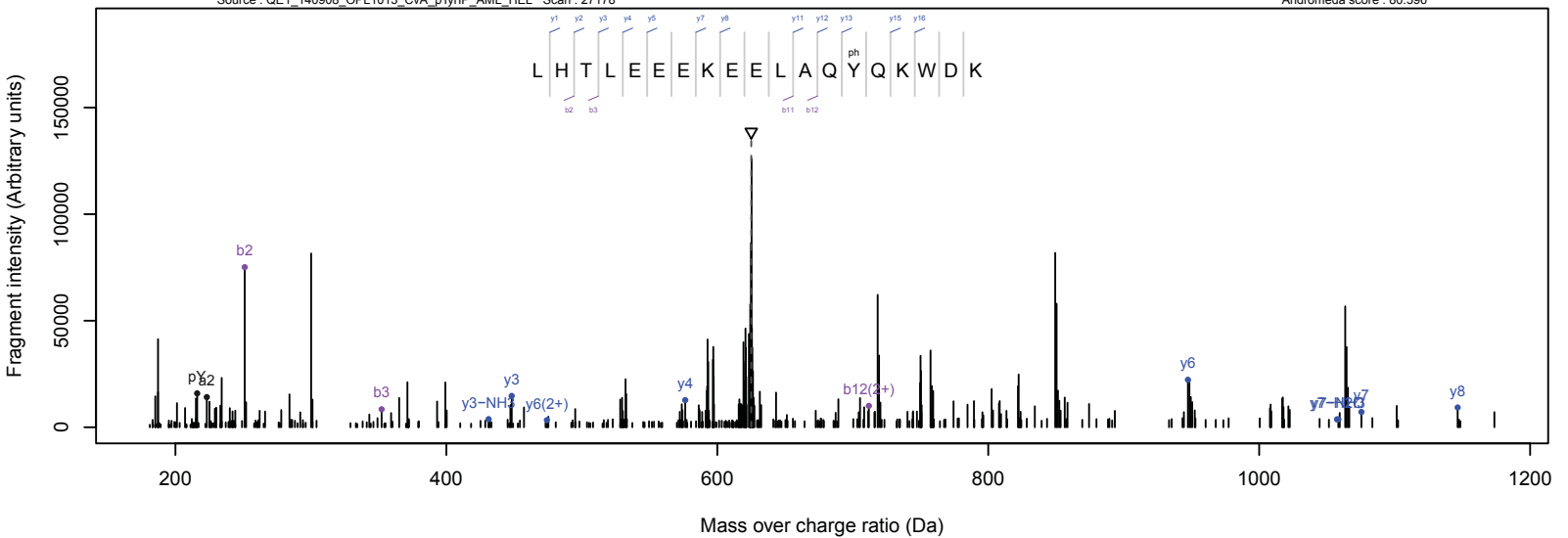
Andromeda score : 80.596



LHTLEEEKEELAQYQKWDK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 27178

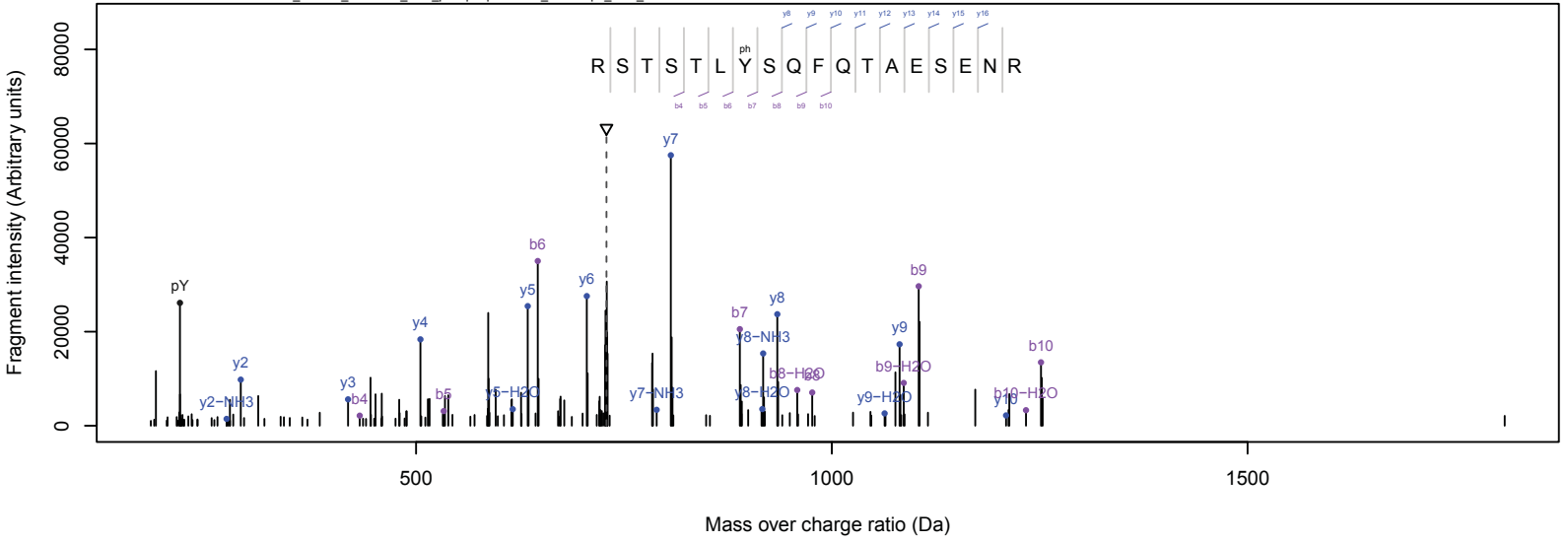
Andromeda score : 80.596



**RSTSTLY(ph)SQFQTAESNR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19555

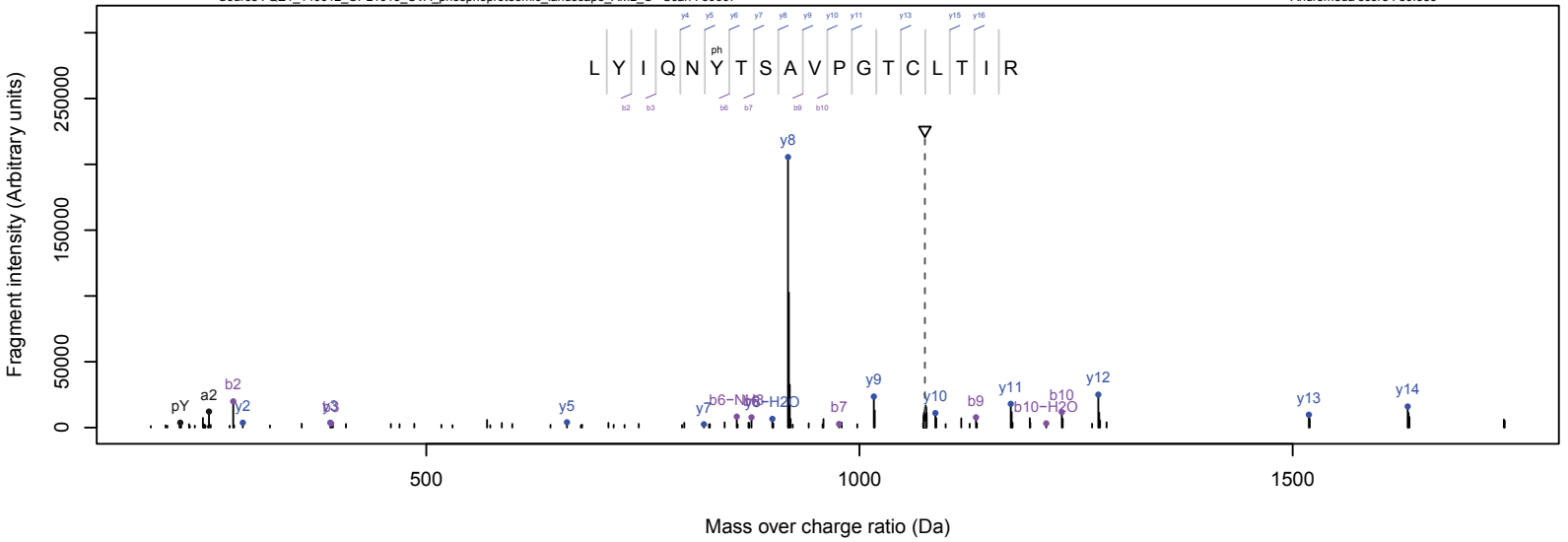
Andromeda score : 80.596



**LYIQNY(ph)TSAVPGTCLTIR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39667

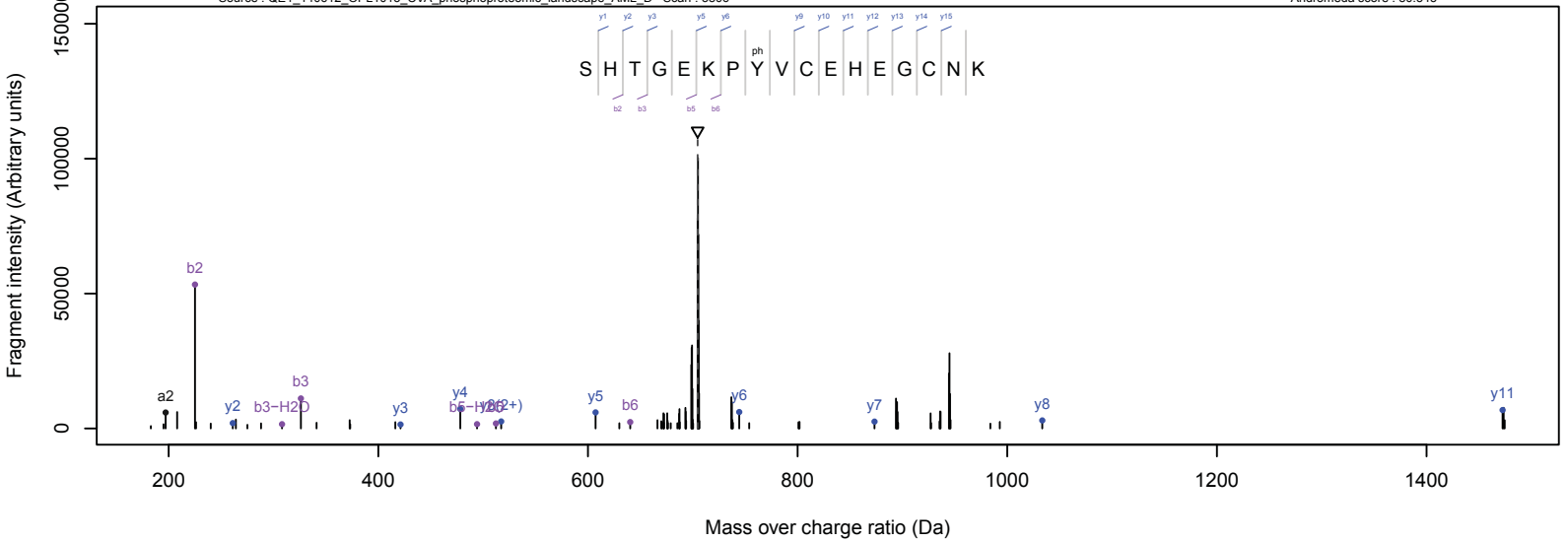
Andromeda score : 80.585



**SHTGEKPY(ph)VCEHEGCKN-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 3806

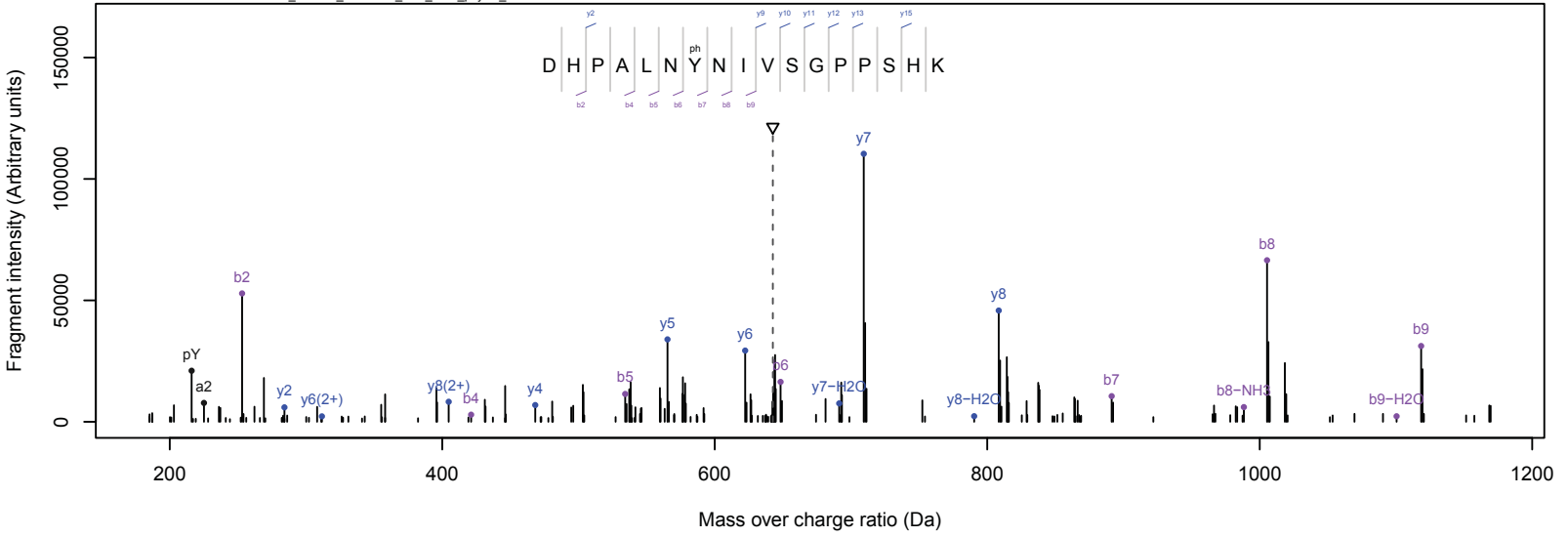
Andromeda score : 80.545



DHPALNY(ph)NIVSGPPSHK-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 20244

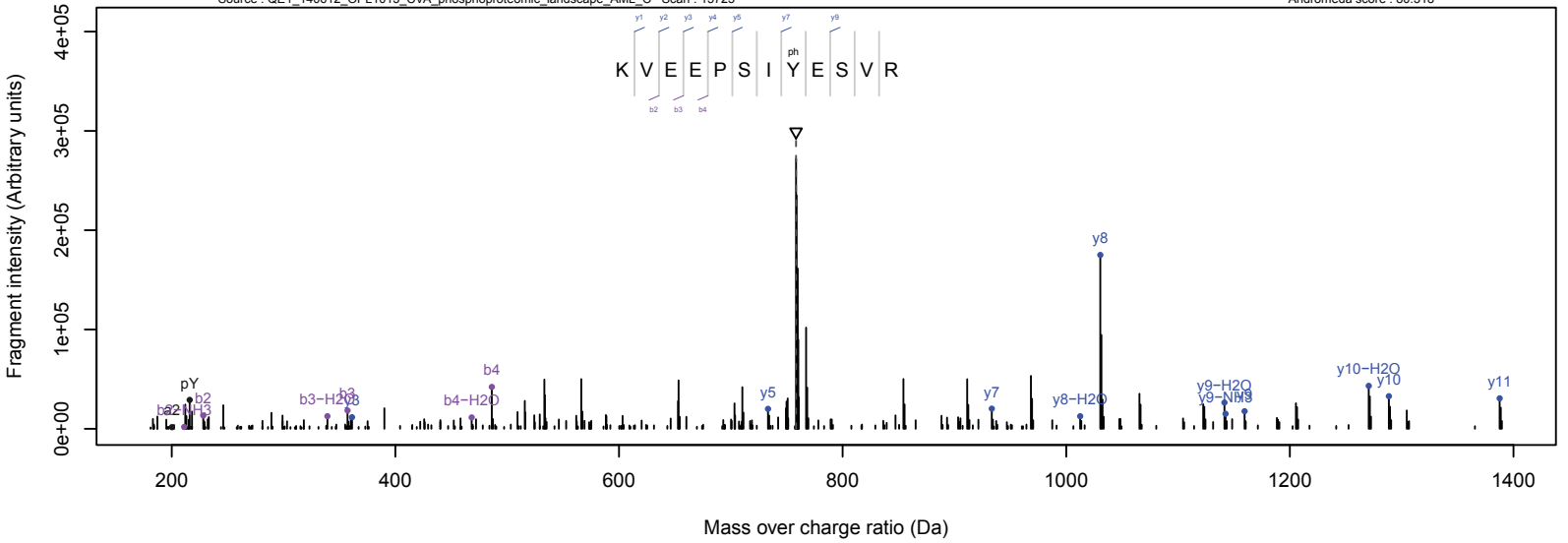
Andromeda score : 80.361



KVEEPSIY(ph)ESVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15725

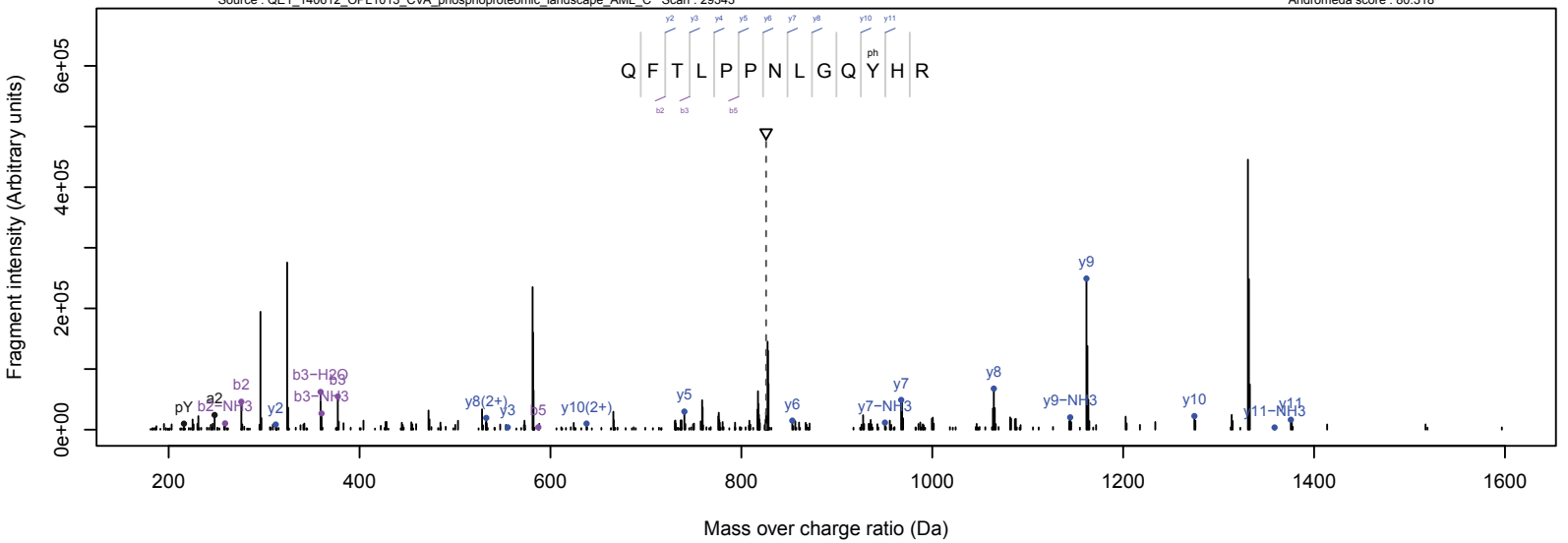
Andromeda score : 80.318



QFTLPPNLGQY(ph)HR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29343

Andromeda score : 80.318

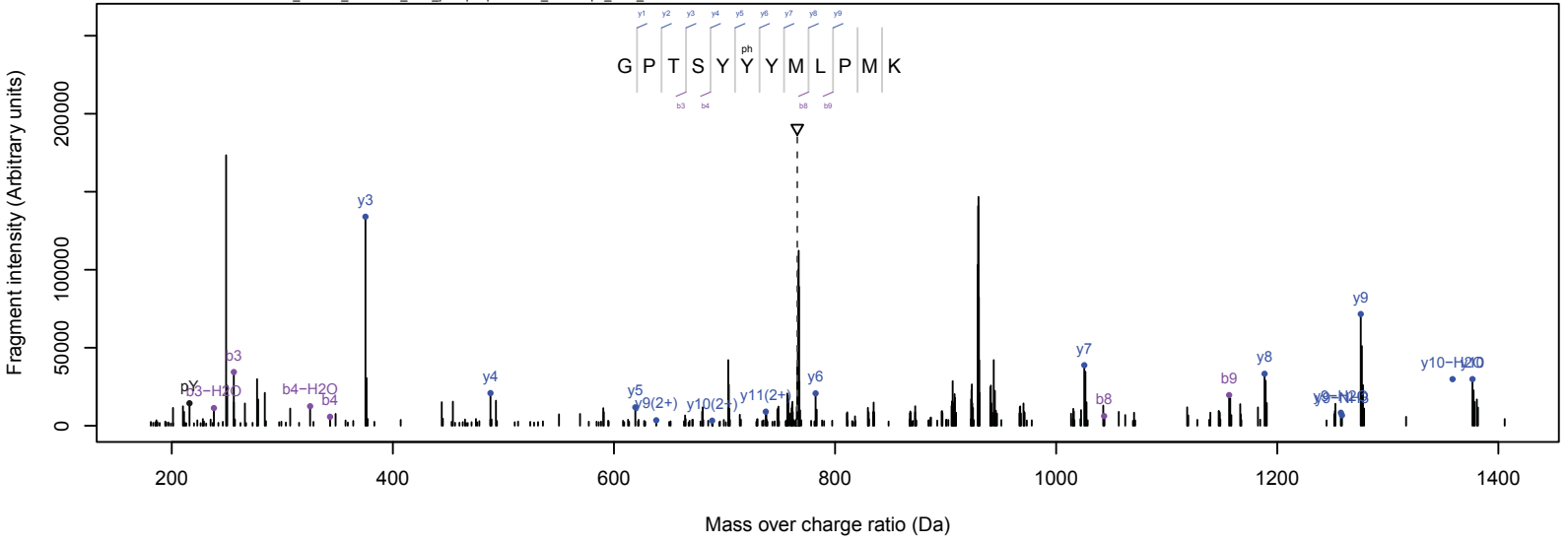




**GPTSYY(ph)YMLPMK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36426

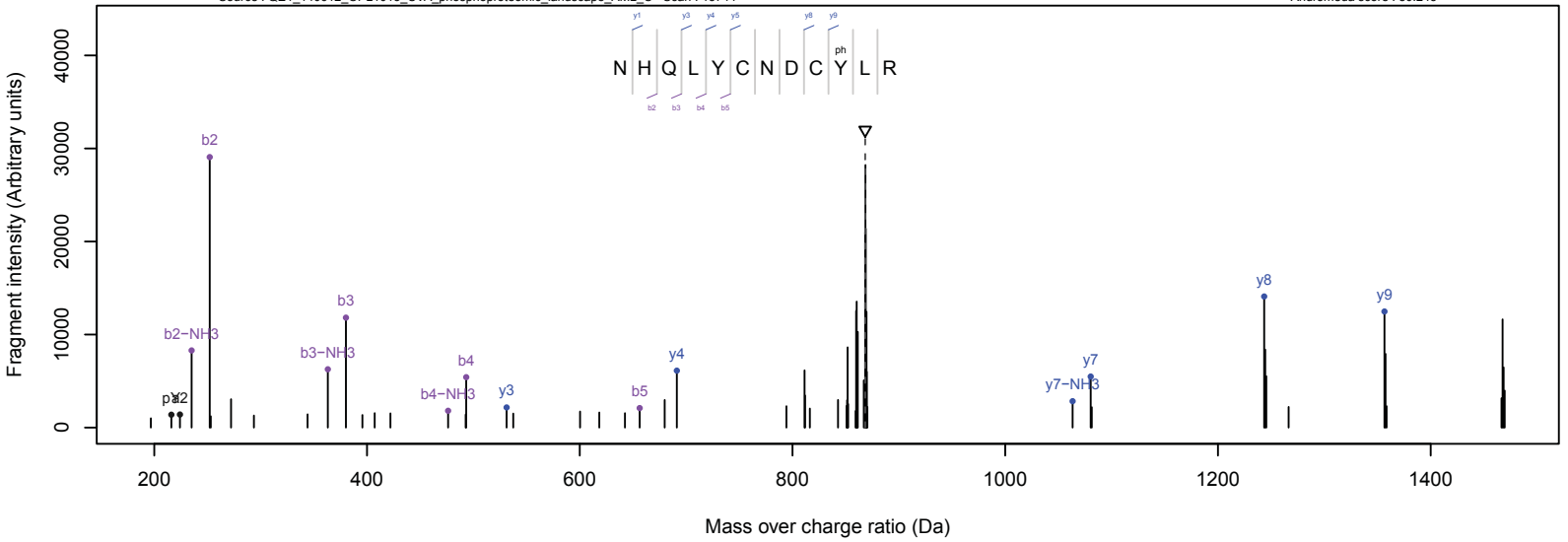
Andromeda score : 80.245



**NHQLYCNDY(ph)LR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13744

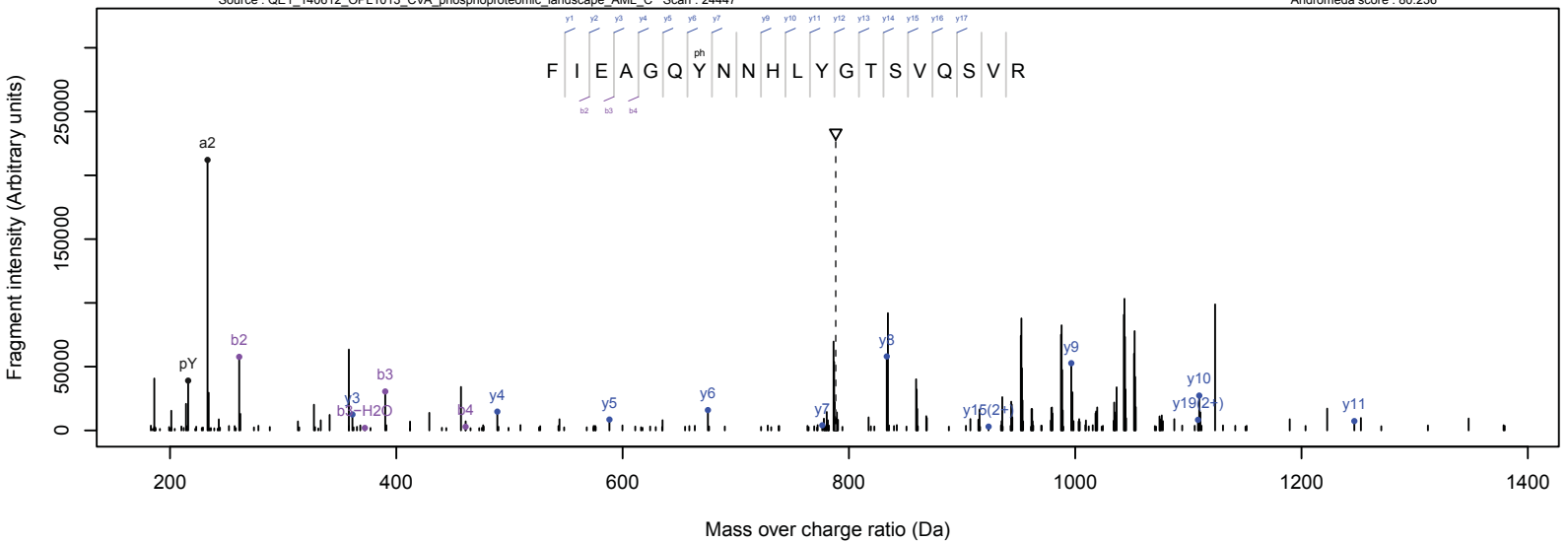
Andromeda score : 80.245



**FIEAGQY(ph)NNHLYGTSVQSVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24447

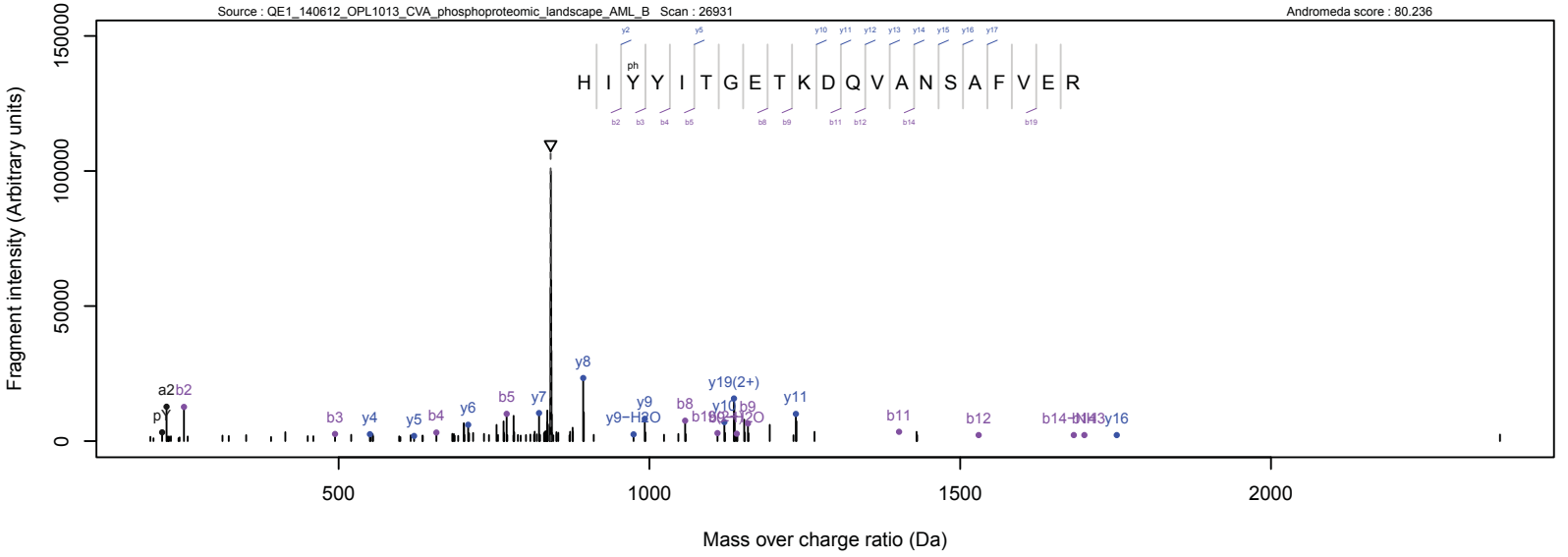
Andromeda score : 80.236



HIY(ph)YITGETKDQVANSAFVER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26931

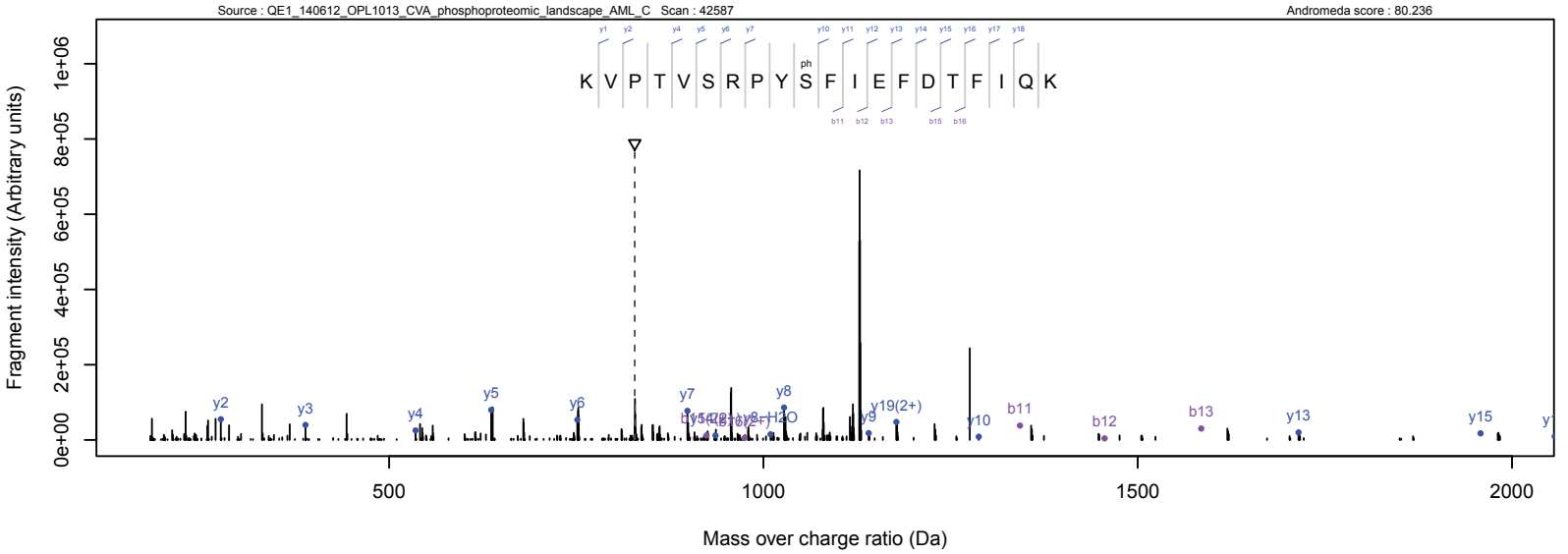
Andromeda score : 80.236



KVPTVSRPYS(ph)FIEFDTFIQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42587

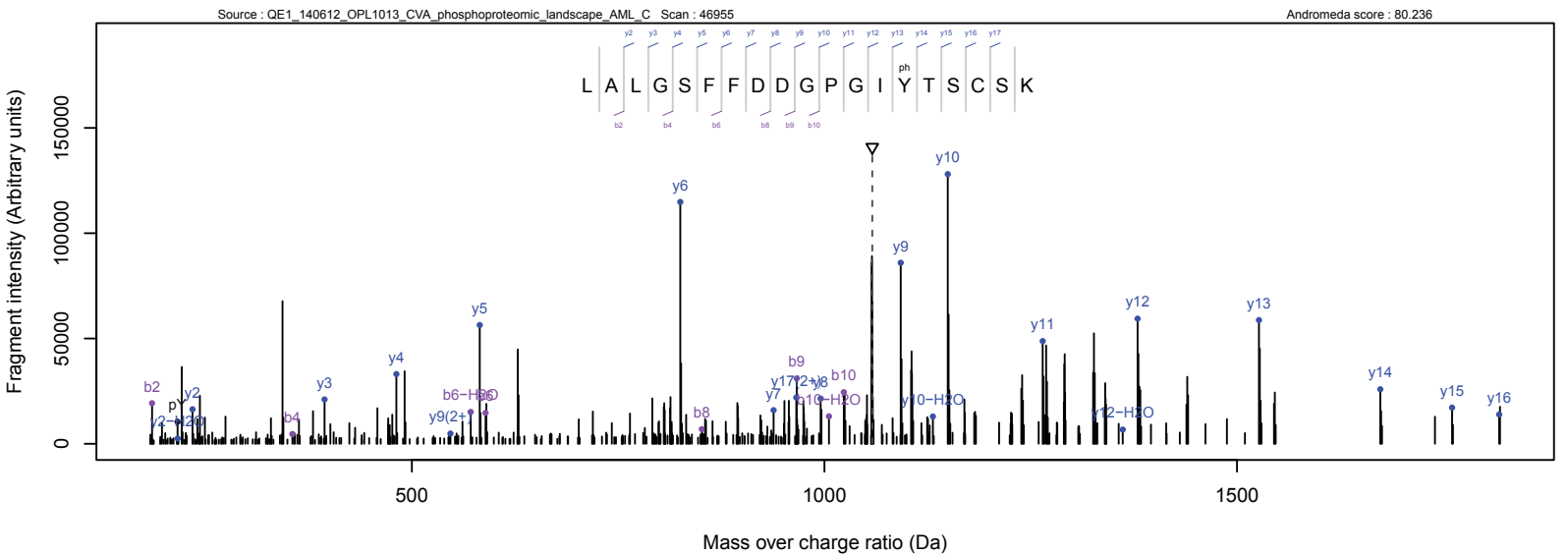
Andromeda score : 80.236



LALGSFFDDGPIY(ph)TSCSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46955

Andromeda score : 80.236

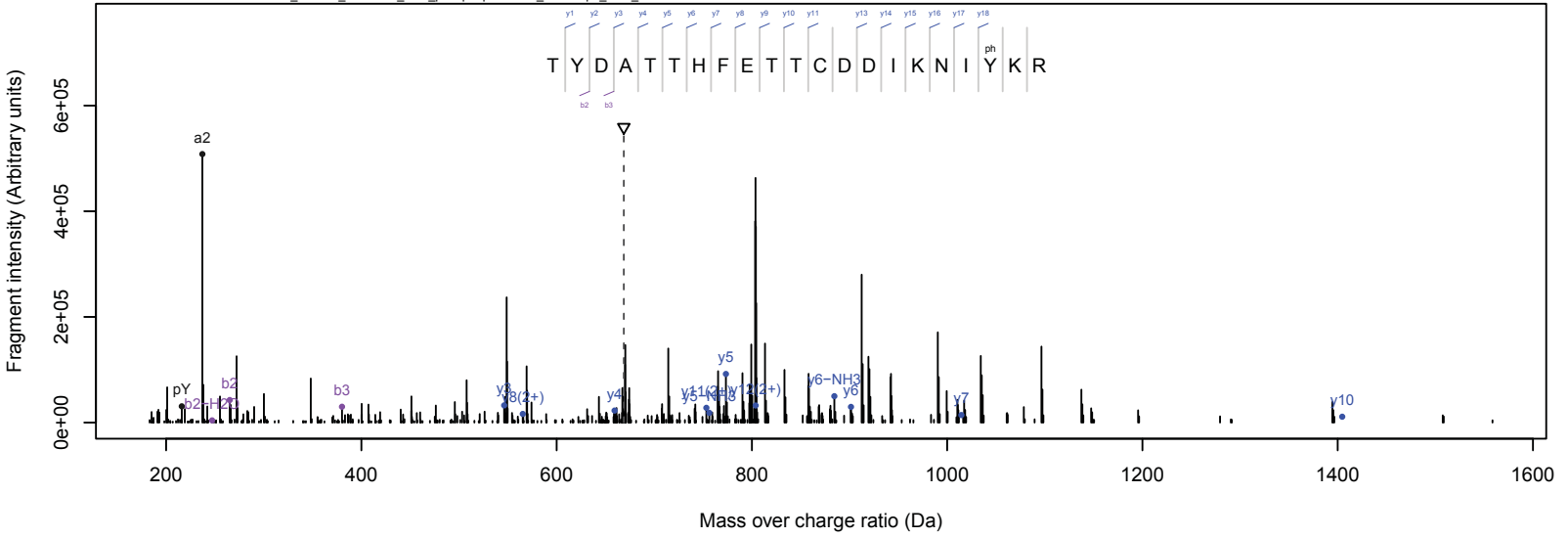




**TYDATTHFETTCDDIKNIY<sup>ph</sup>KR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23086

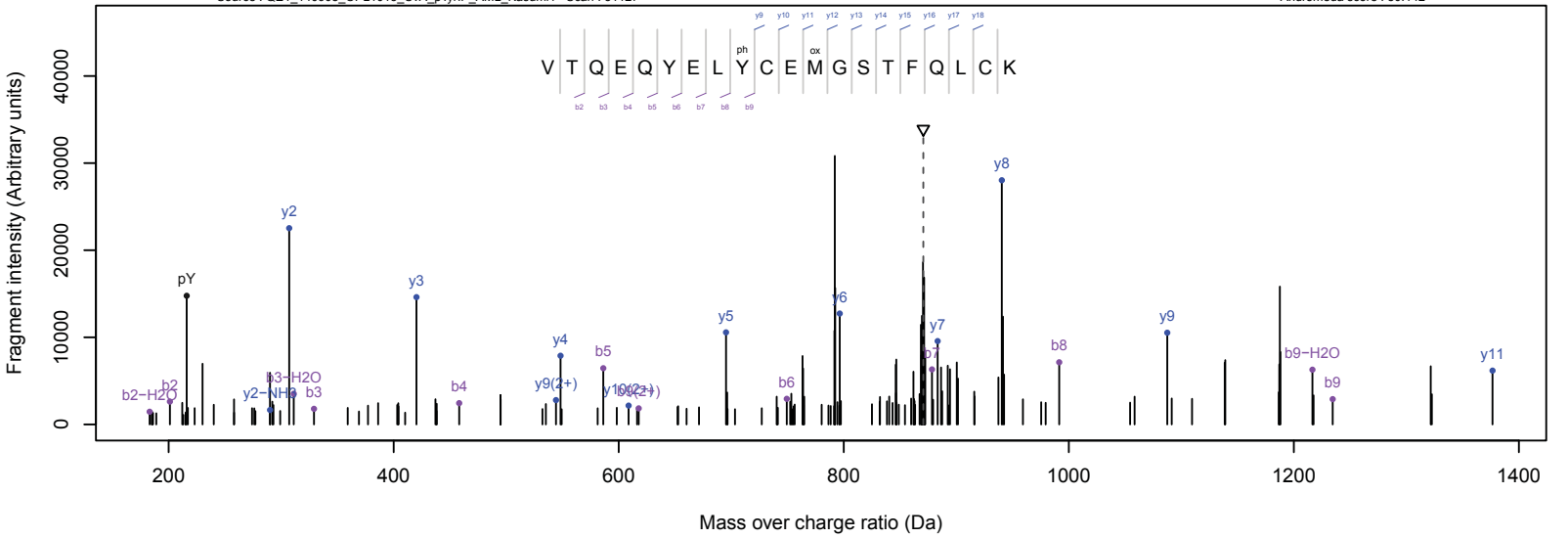
Andromeda score : 80.142



**VTQEQYELY<sup>ph</sup>CEM<sup>ox</sup>GSTFQLCK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 31127

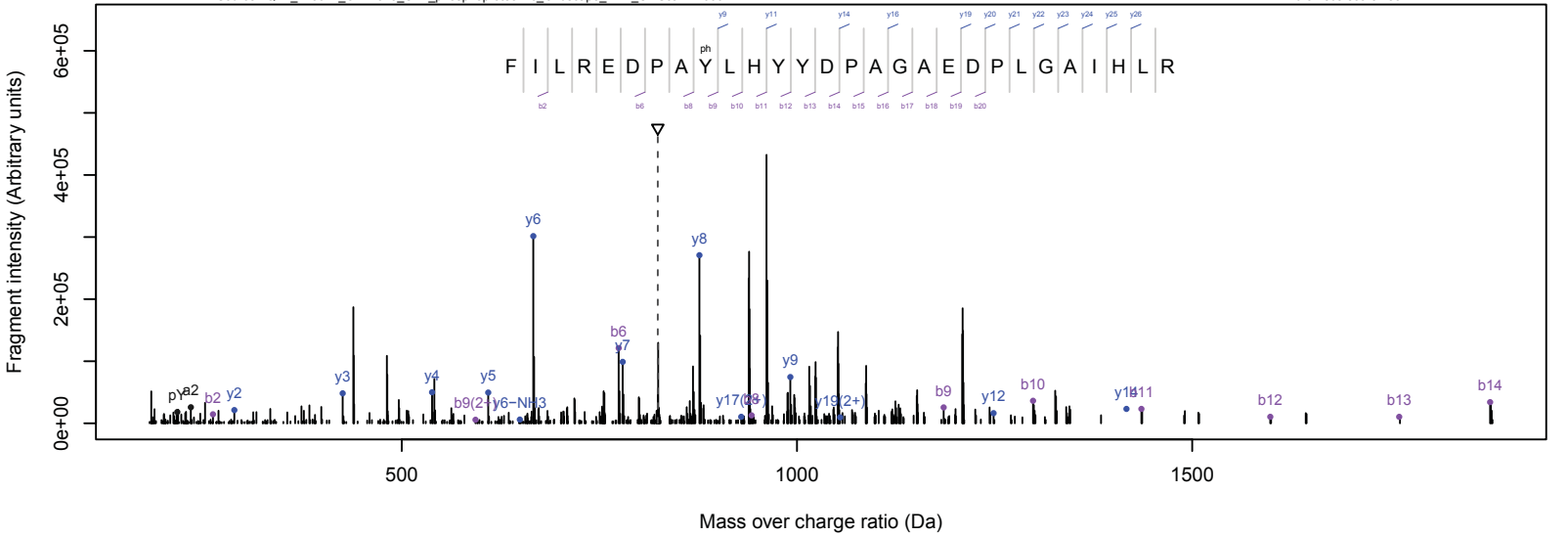
Andromeda score : 80.142



**FILREDPAY<sup>ph</sup>LHYYPAGAEDPLGAIHLR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46667

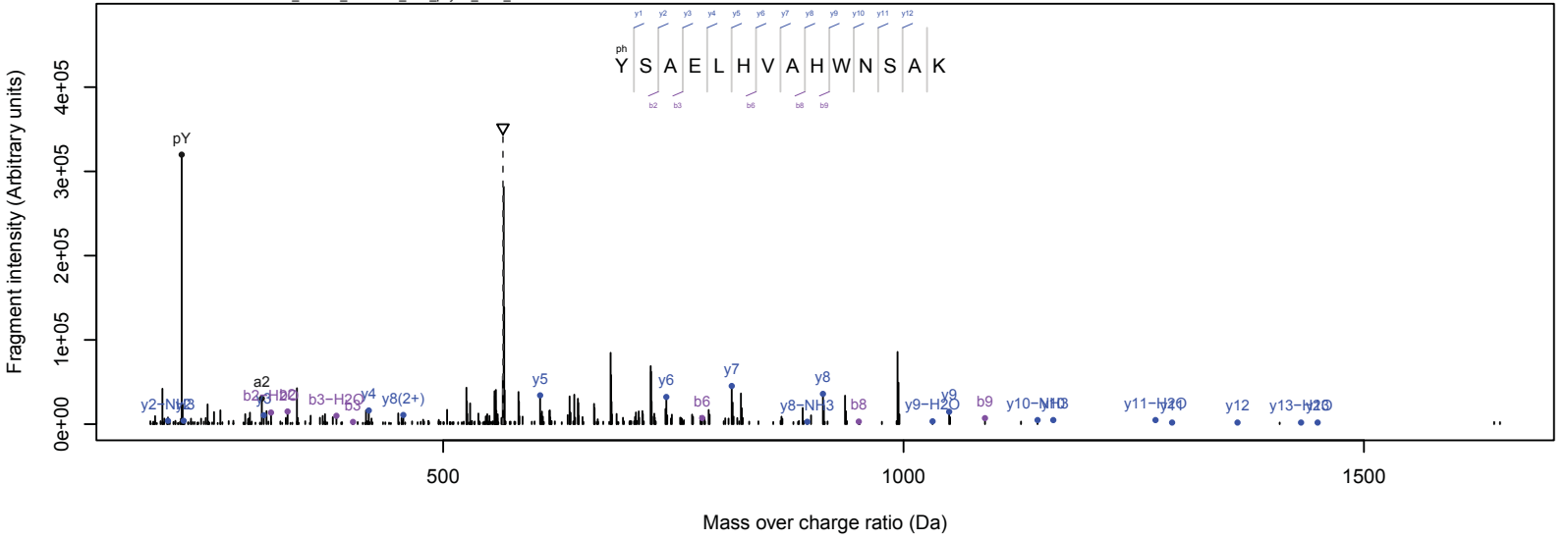
Andromeda score : 80.117



Y(ph)SAELHVAHWNSAK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 16979

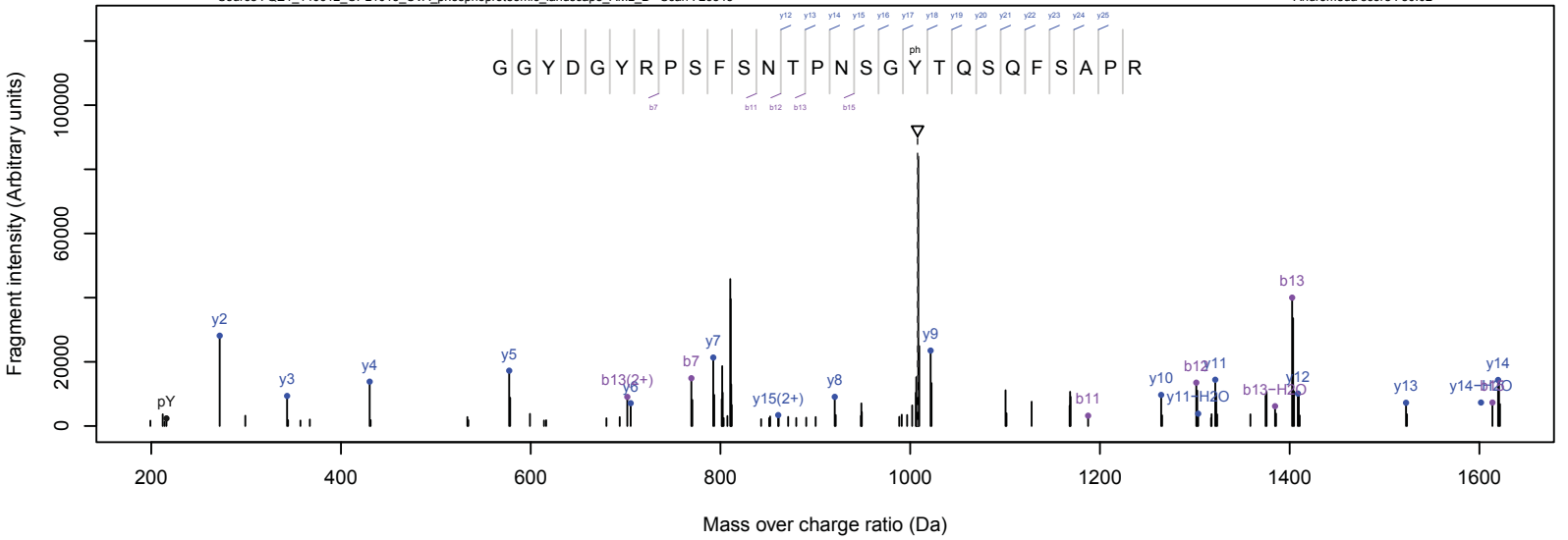
Andromeda score : 80.102



GGYDGYRPSFSNTPSG(ph)TQSQFSAPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26543

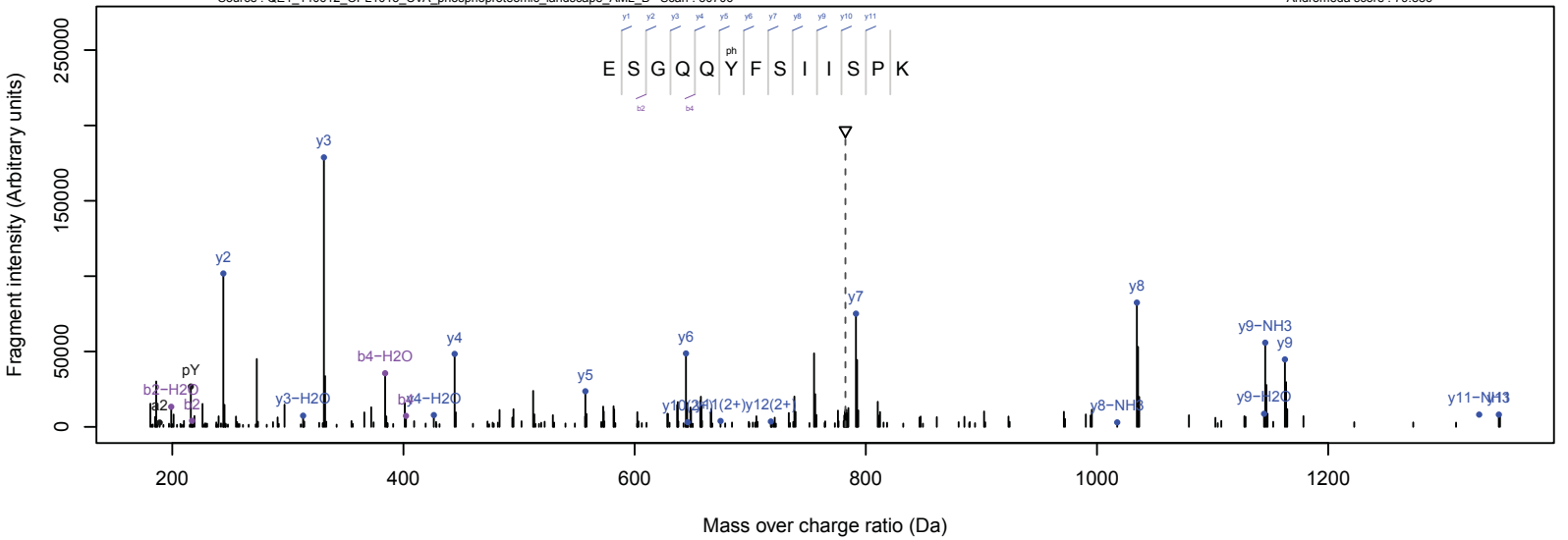
Andromeda score : 80.02



ESGQQY(ph)FSIISPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30706

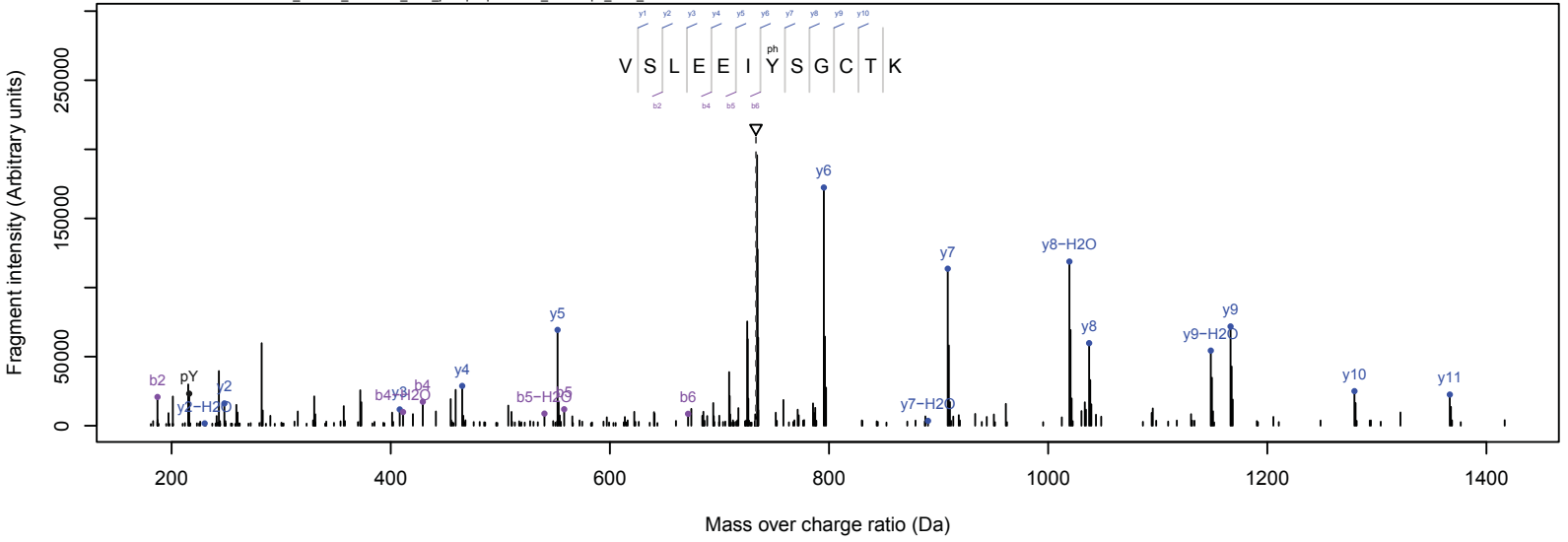
Andromeda score : 79.886



**VSLEEIY(ph)SGCTK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22194

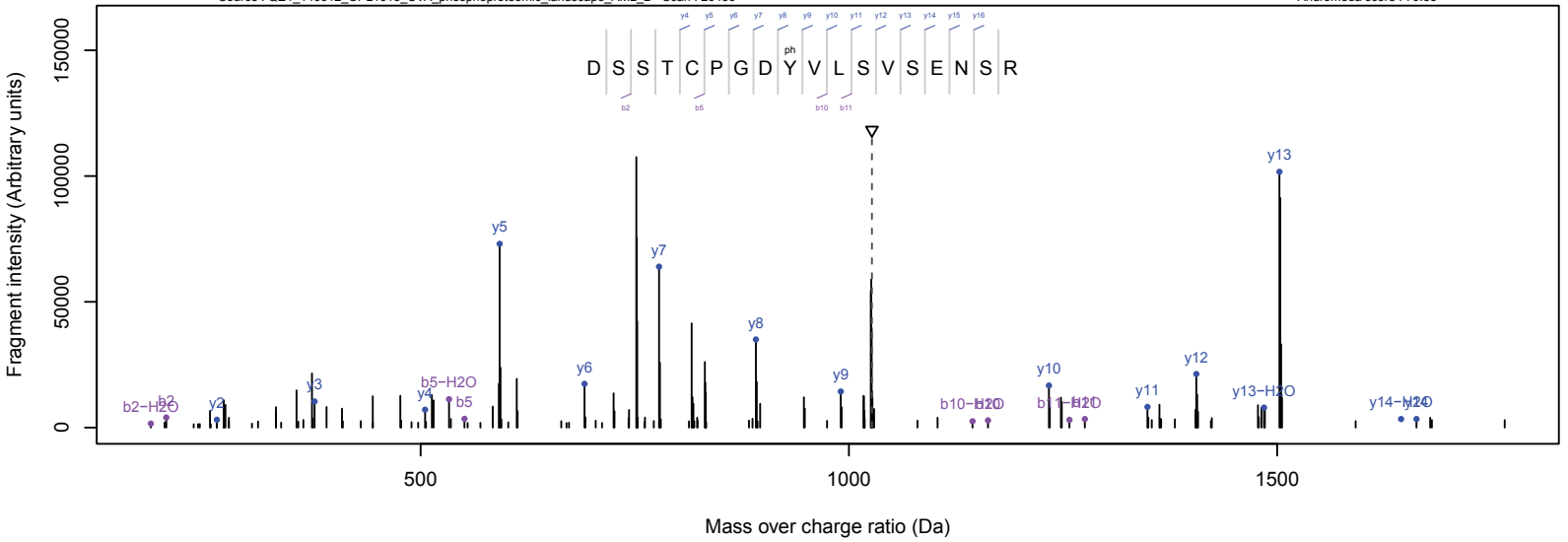
Andromeda score : 79.886



**DSSTCPGDYVLSVSENSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26436

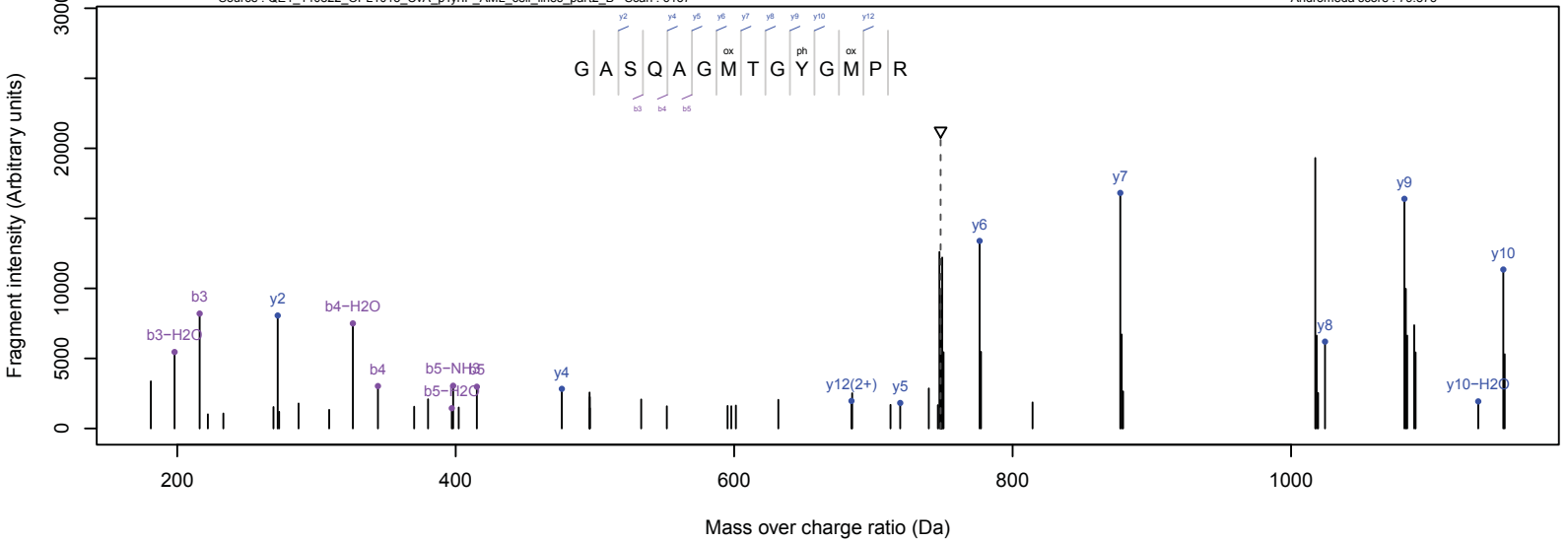
Andromeda score : 79.88



**GASQAGM(ox)TGY(ph)GM(ox)PR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 6137

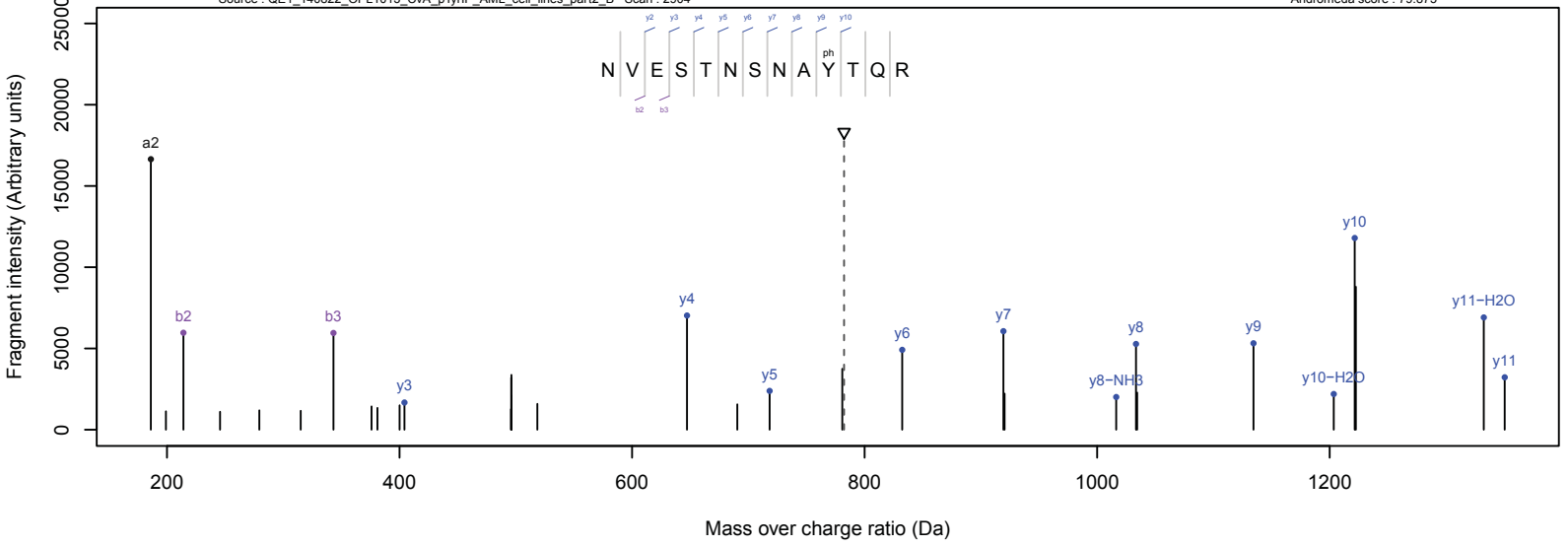
Andromeda score : 79.875



**NVESTNSNAY(ph)TQR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 2904

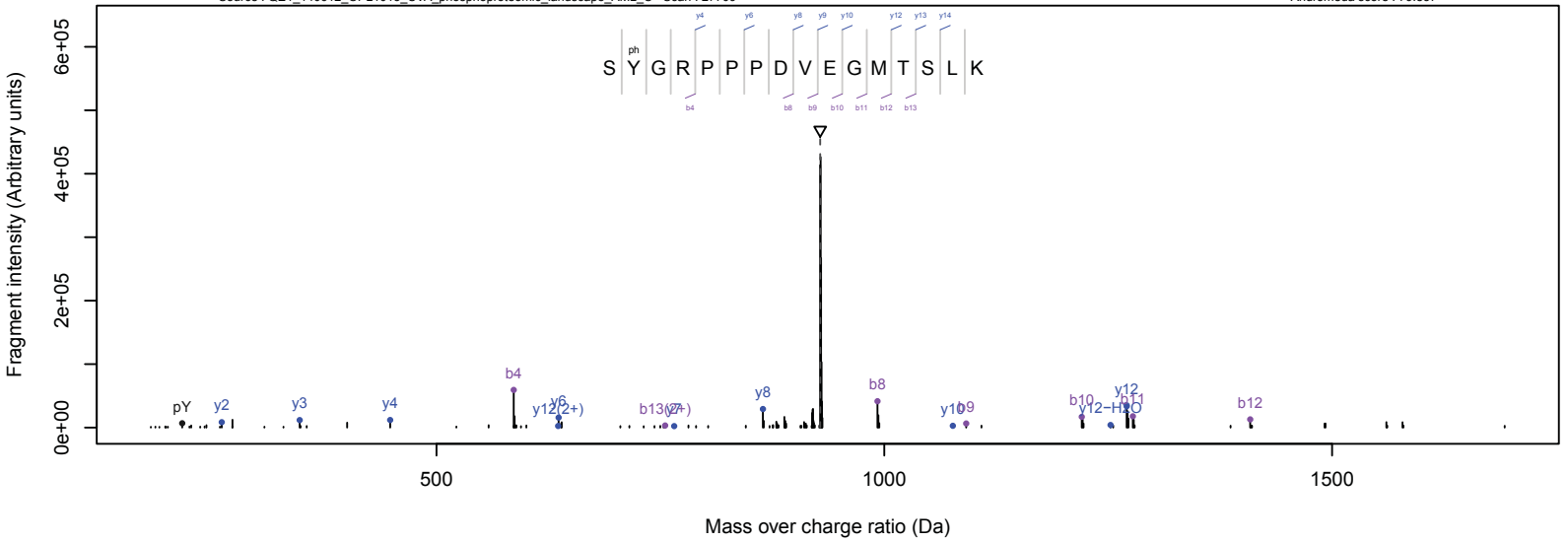
Andromeda score : 79.875



**(ac)SY(ph)GRPPDVEGMTSLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27709

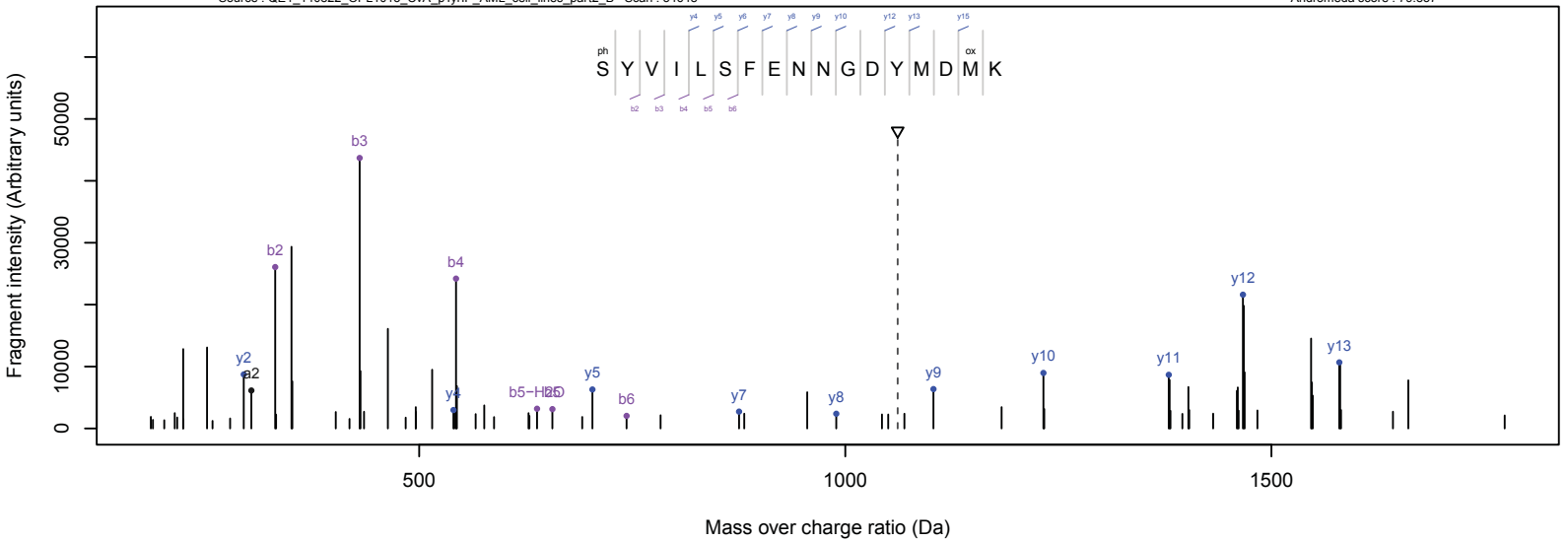
Andromeda score : 79.837



**S(ph)YVILSFENNGDYMDM(ox)K-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 34648

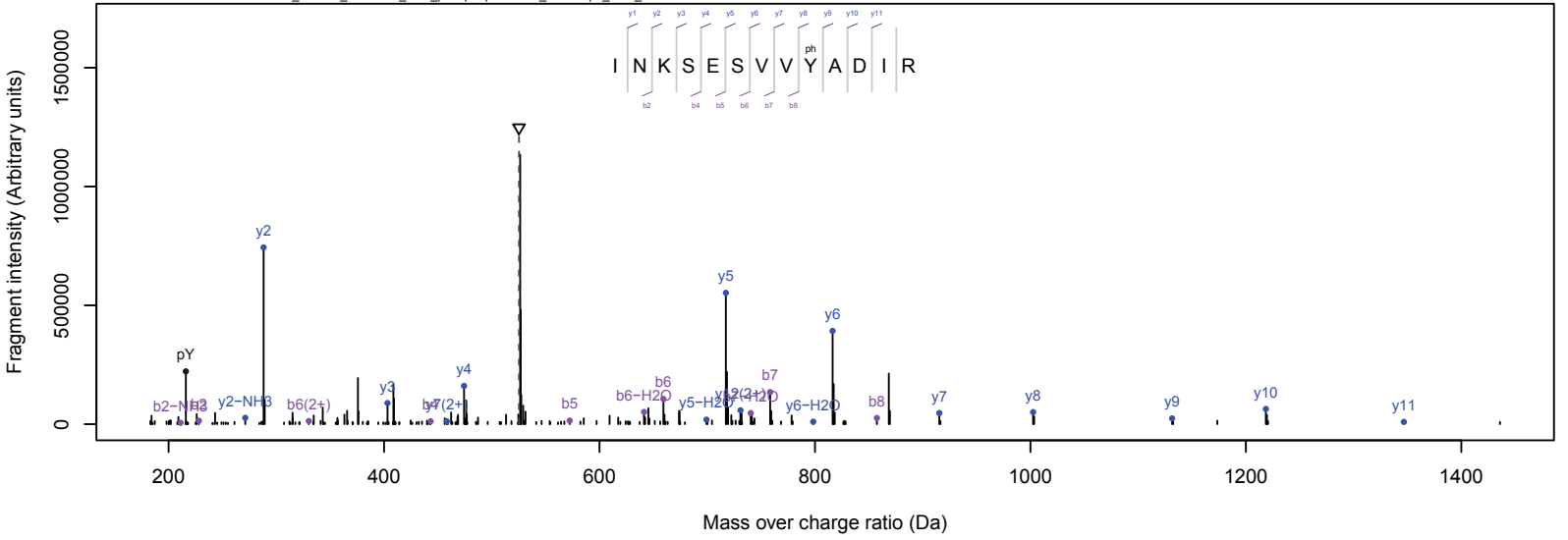
Andromeda score : 79.837



**INKSESVVY(ph)ADIR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16677

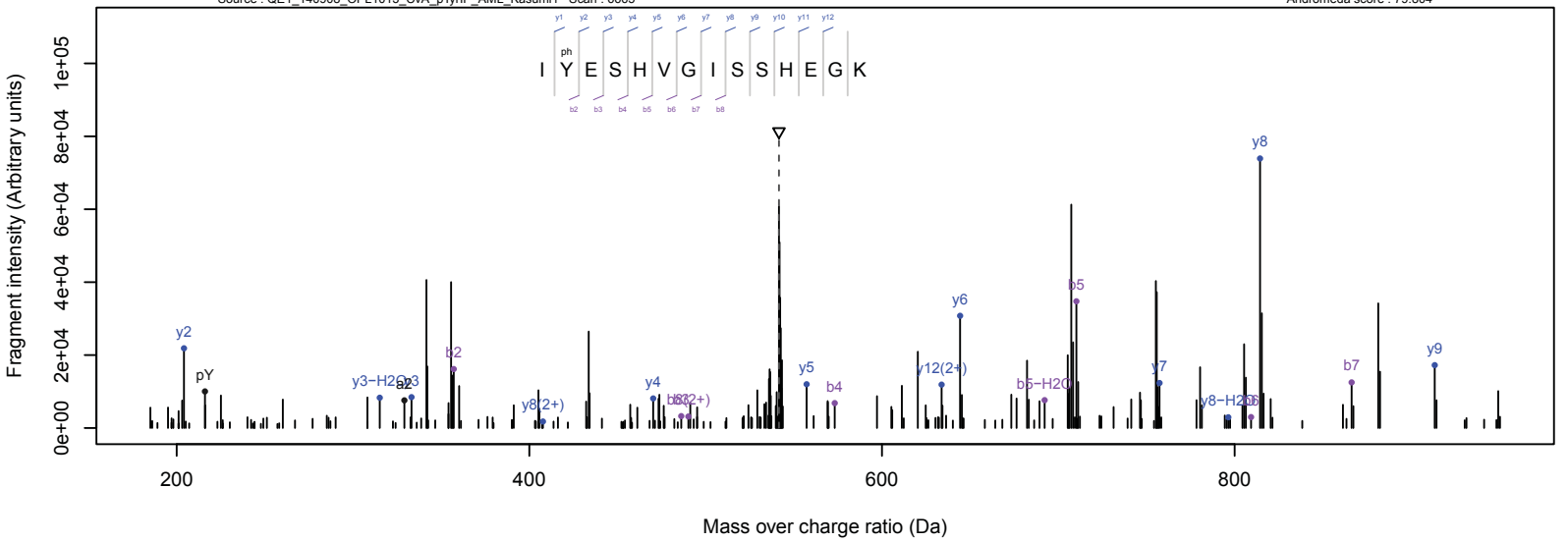
Andromeda score : 79.804



**IY(ph)ESHVGISSHEGK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 6665

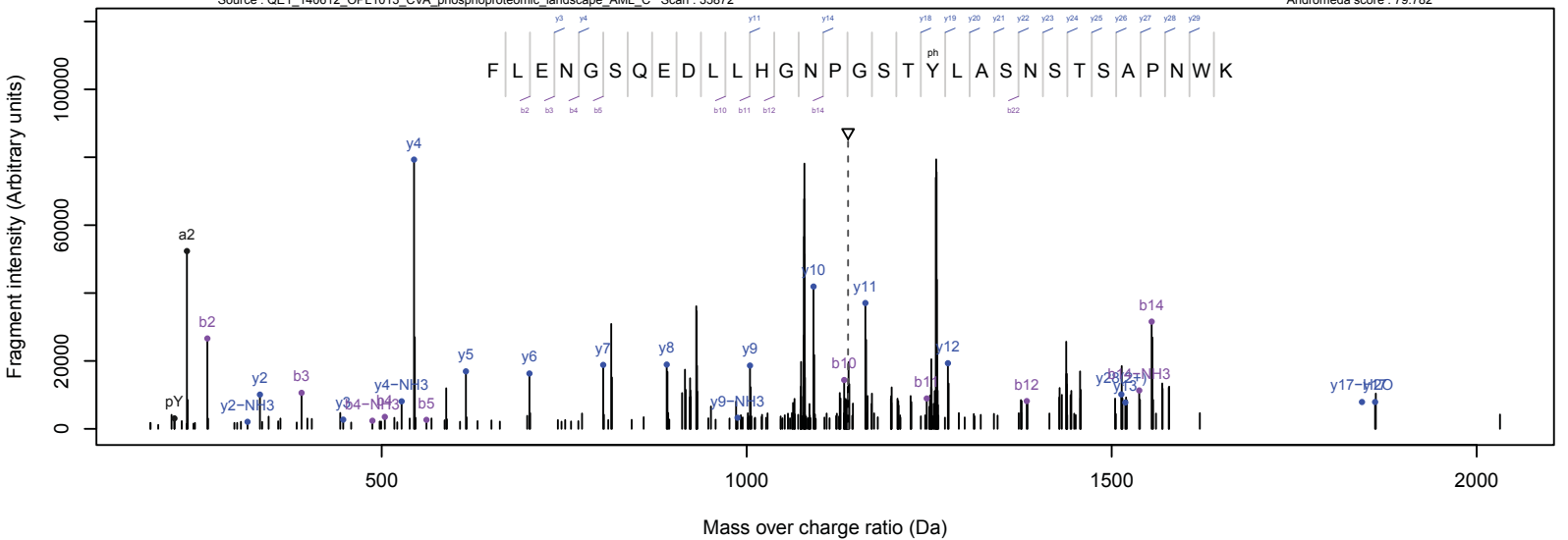
Andromeda score : 79.804



**FLENGSQEDLLHGNGSTY(ph)LASNSTSAPNWK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35872

Andromeda score : 79.782



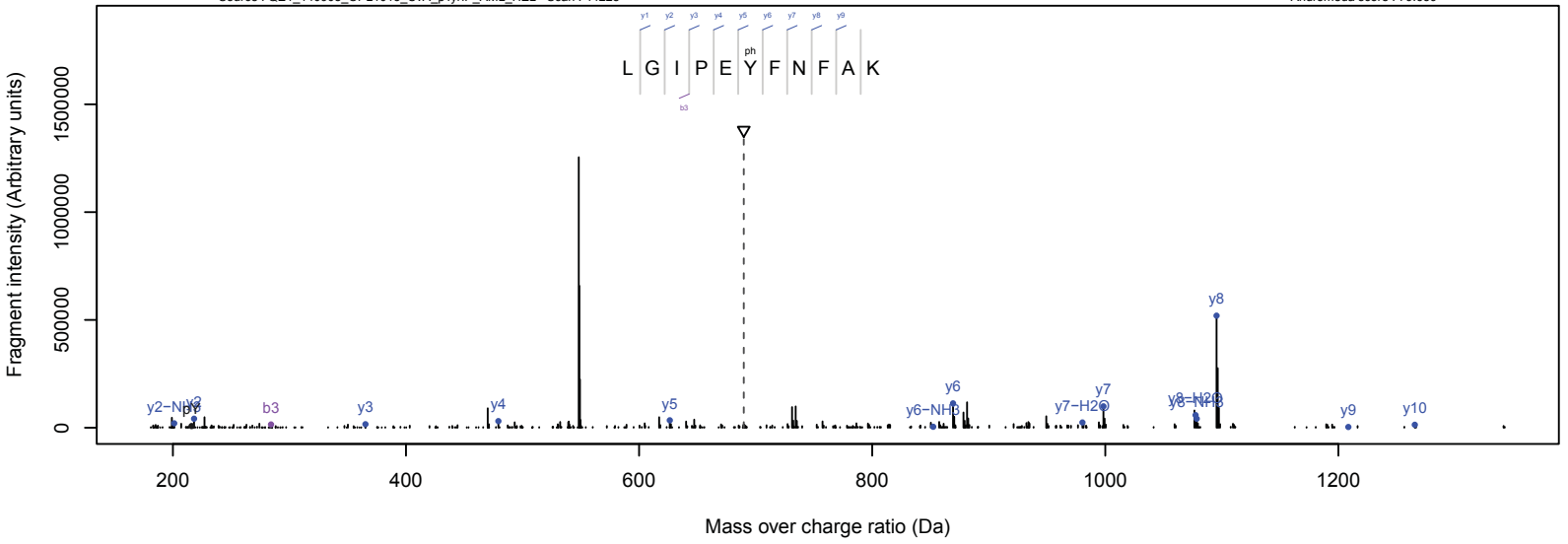




LG IPEY FNF A K

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 44228

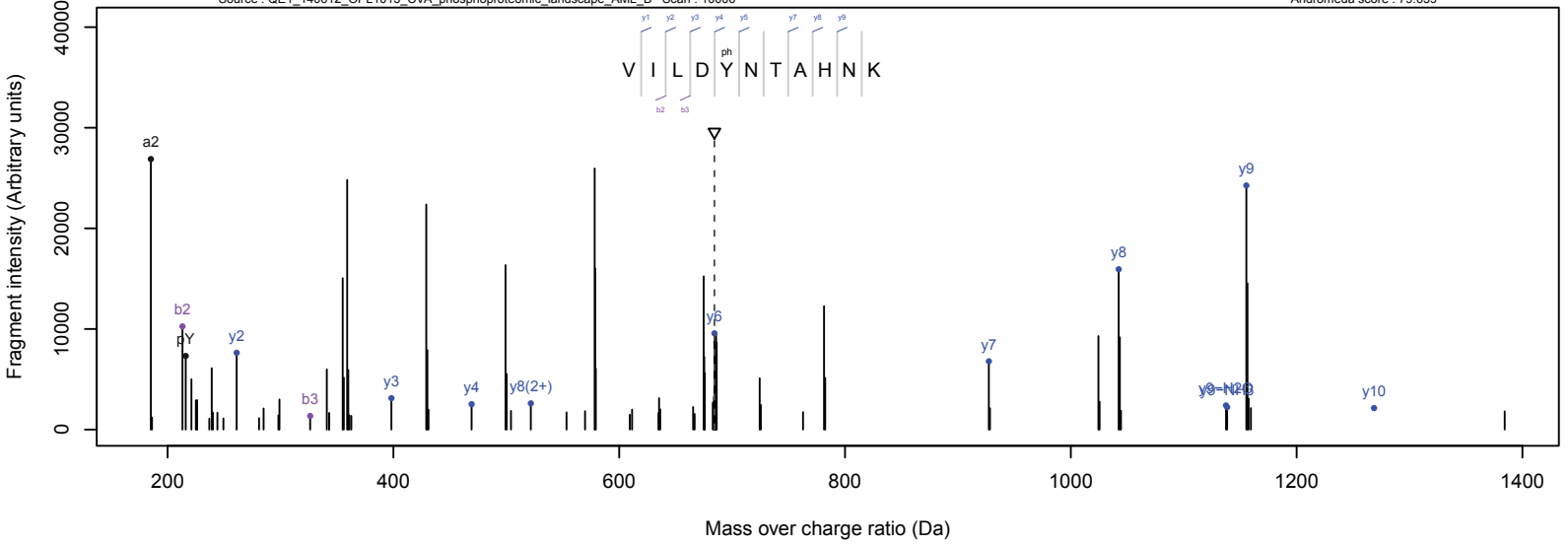
Andromeda score : 79.659



V I L D Y N T A H N K

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10006

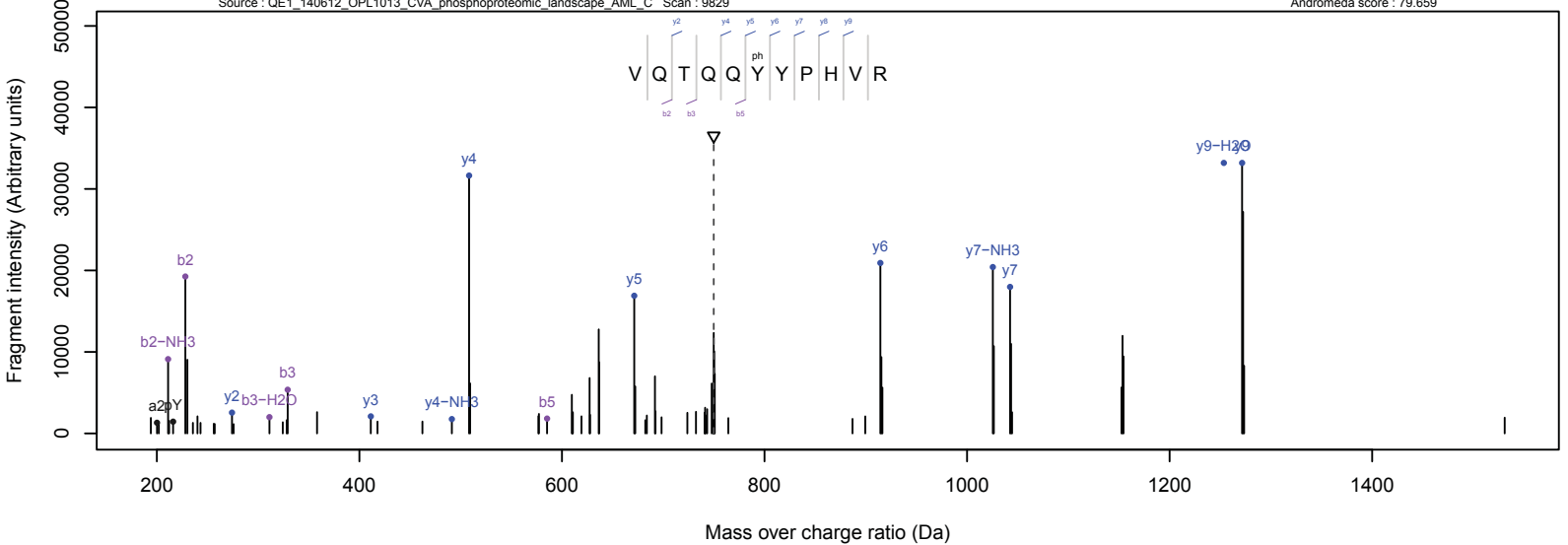
Andromeda score : 79.659



V Q T Q Q Y Y P H V R

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9829

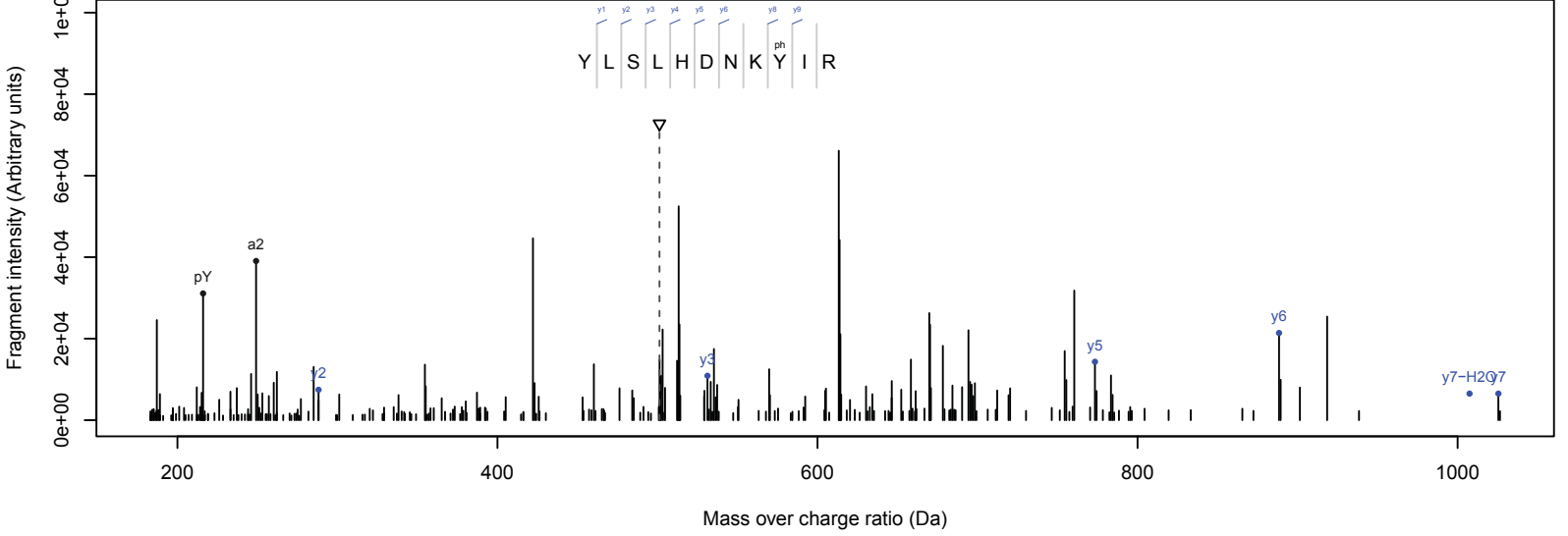
Andromeda score : 79.659



YLSLHDNKY(ph)IR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 15920

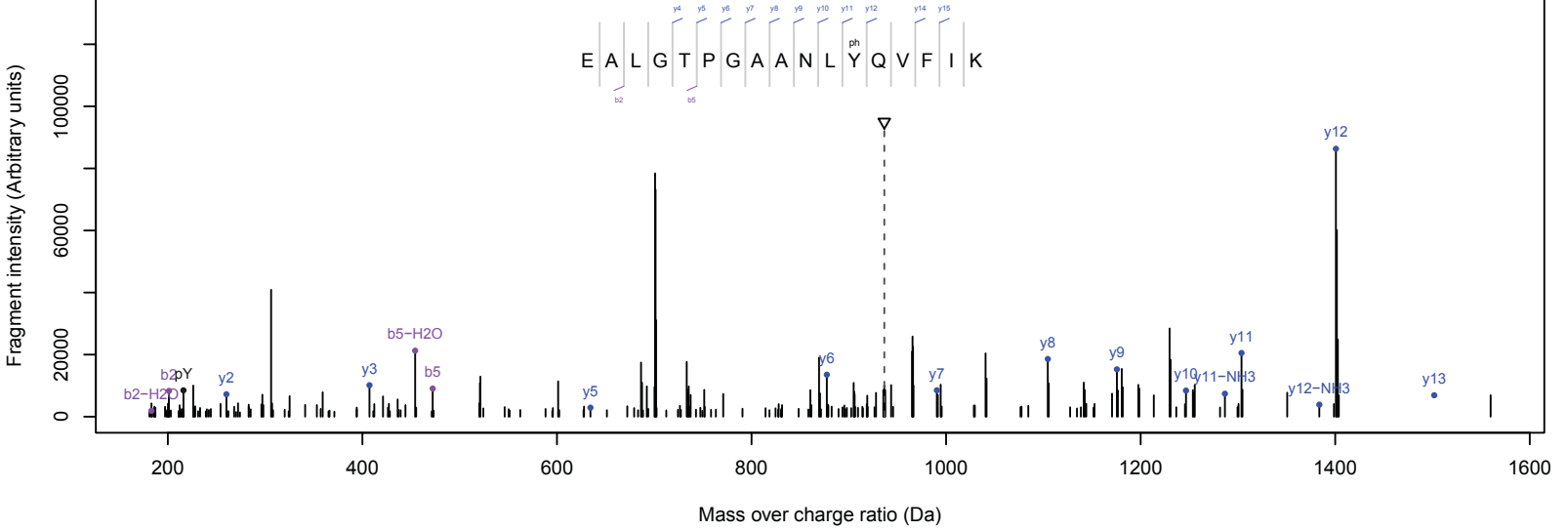
Andromeda score: 79.659



EALGTPGAANLY(ph)QVFIK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 43588

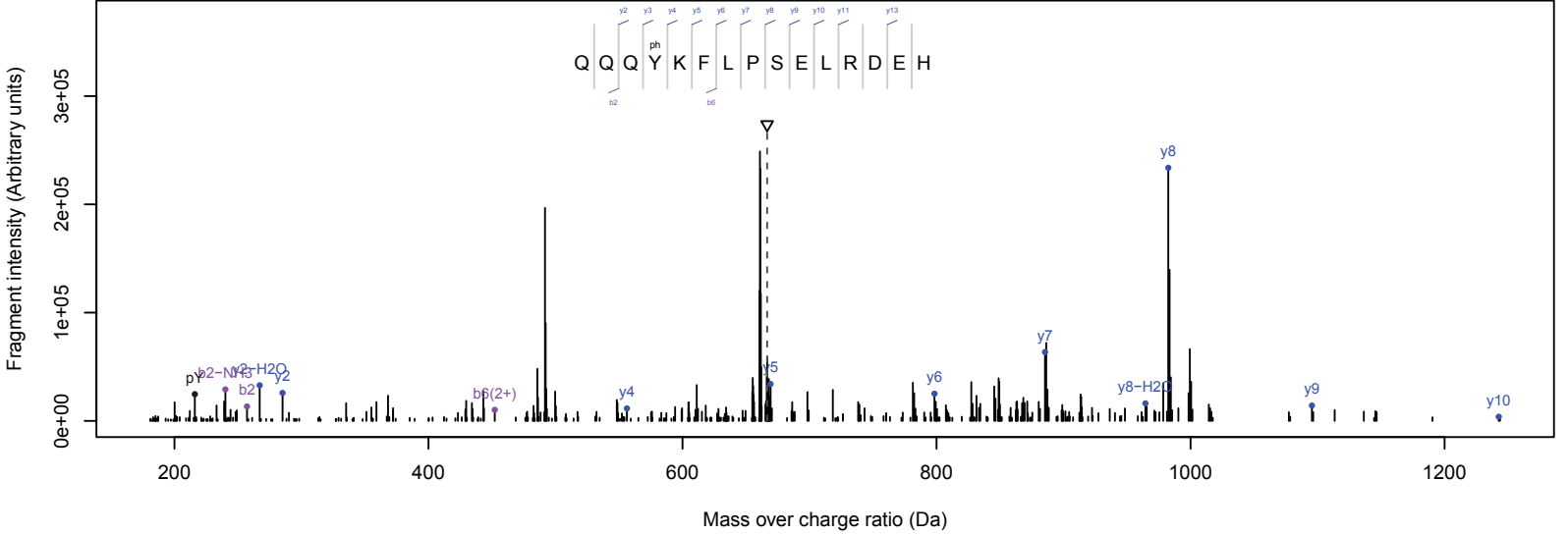
Andromeda score: 79.652



QQQY(ph)KFLPSELRDEH-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 21747

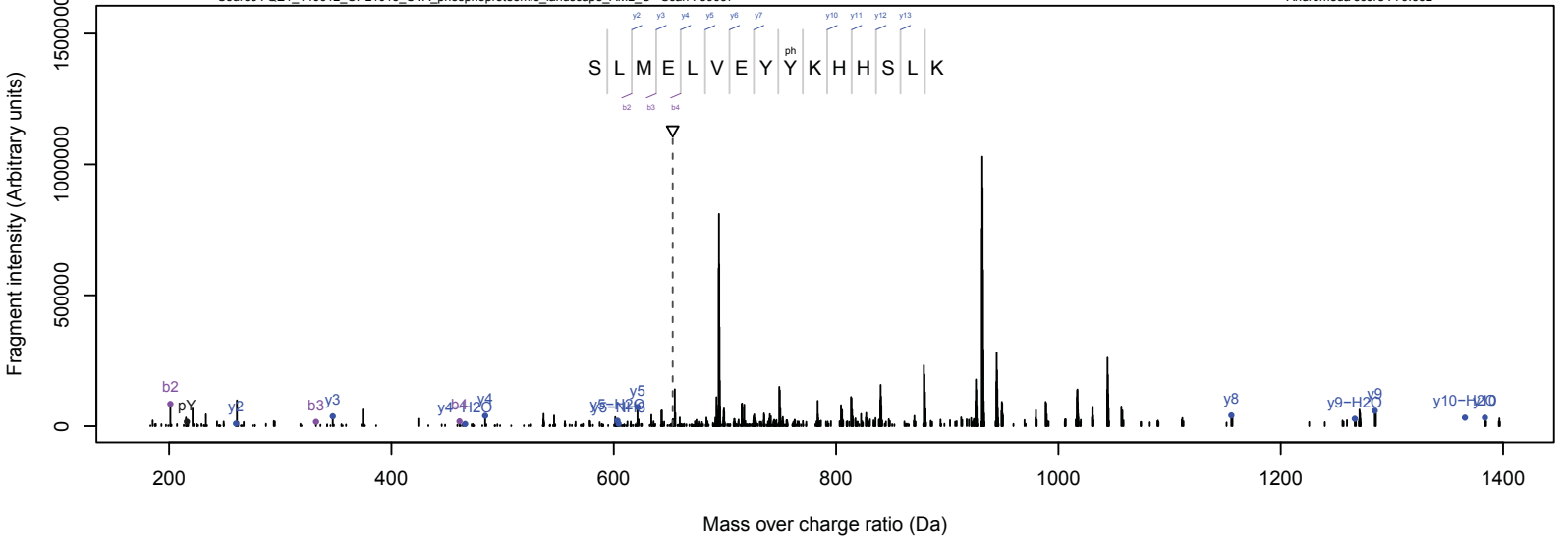
Andromeda score: 79.652



SLMELVEYY(ph)KHHSLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39057

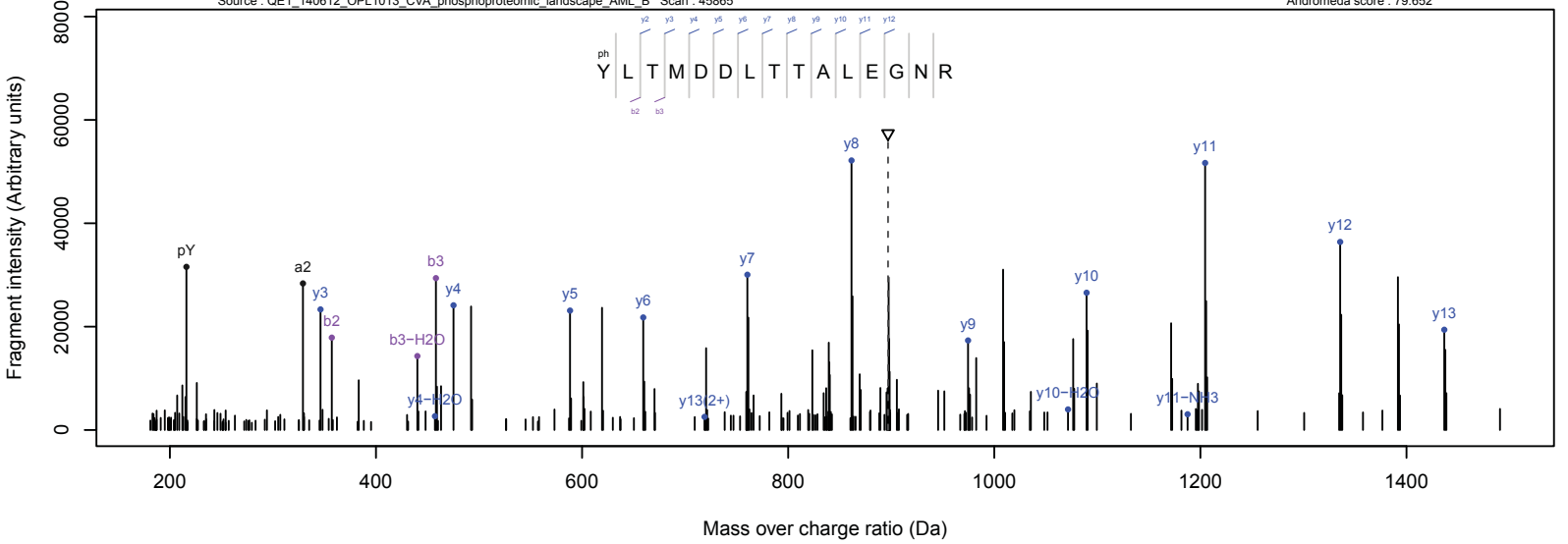
Andromeda score : 79.652



Y(ph)LTMDLLTALEGNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45865

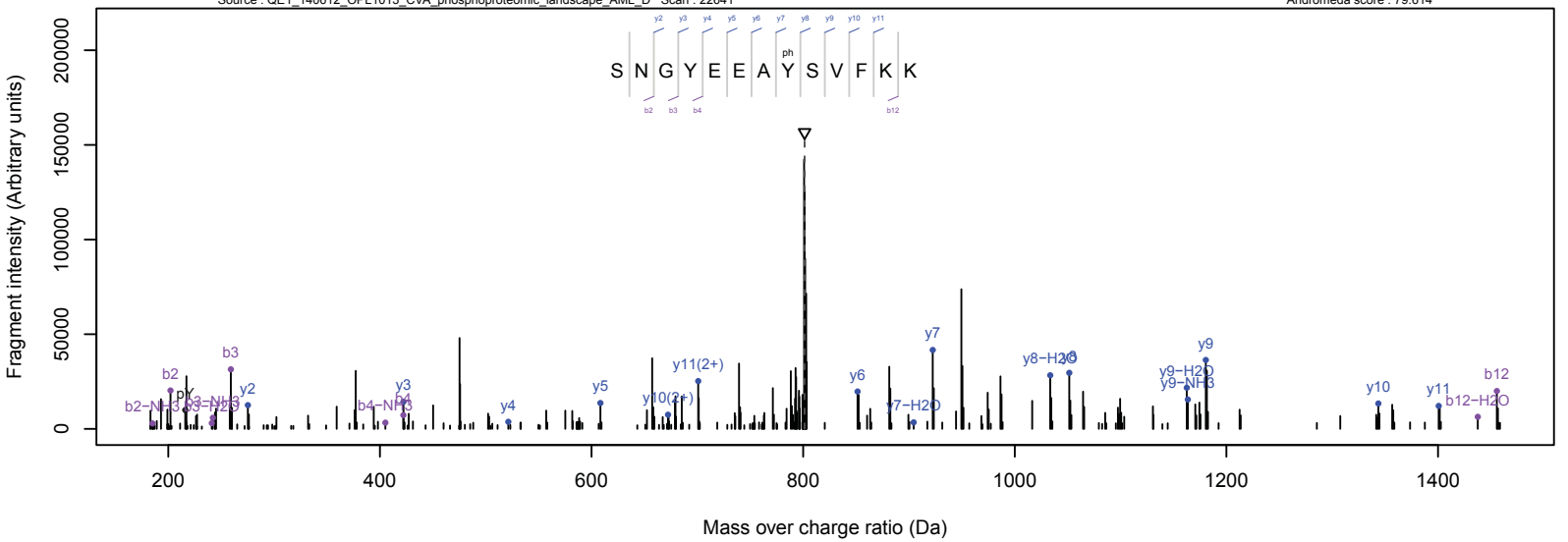
Andromeda score : 79.652



SNGYEEAY(ph)SVFKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22641

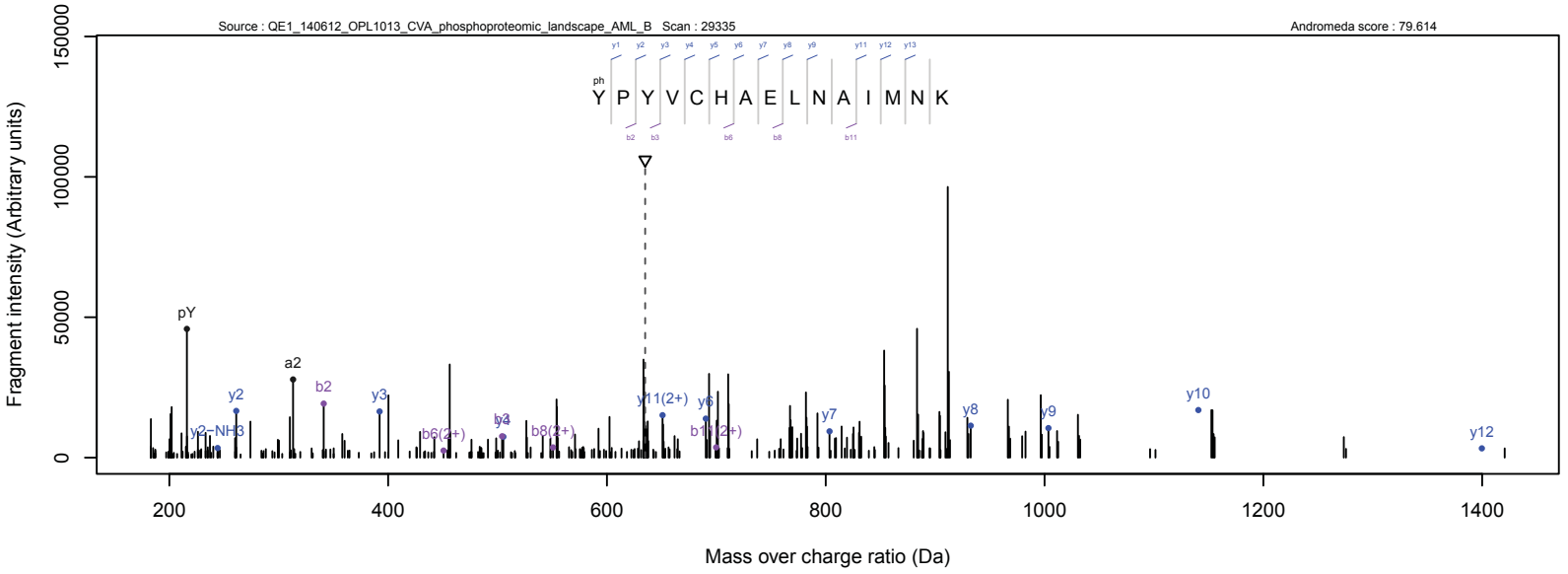
Andromeda score : 79.614



**Y(ph)PYVCHAE LNAIMNK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29335

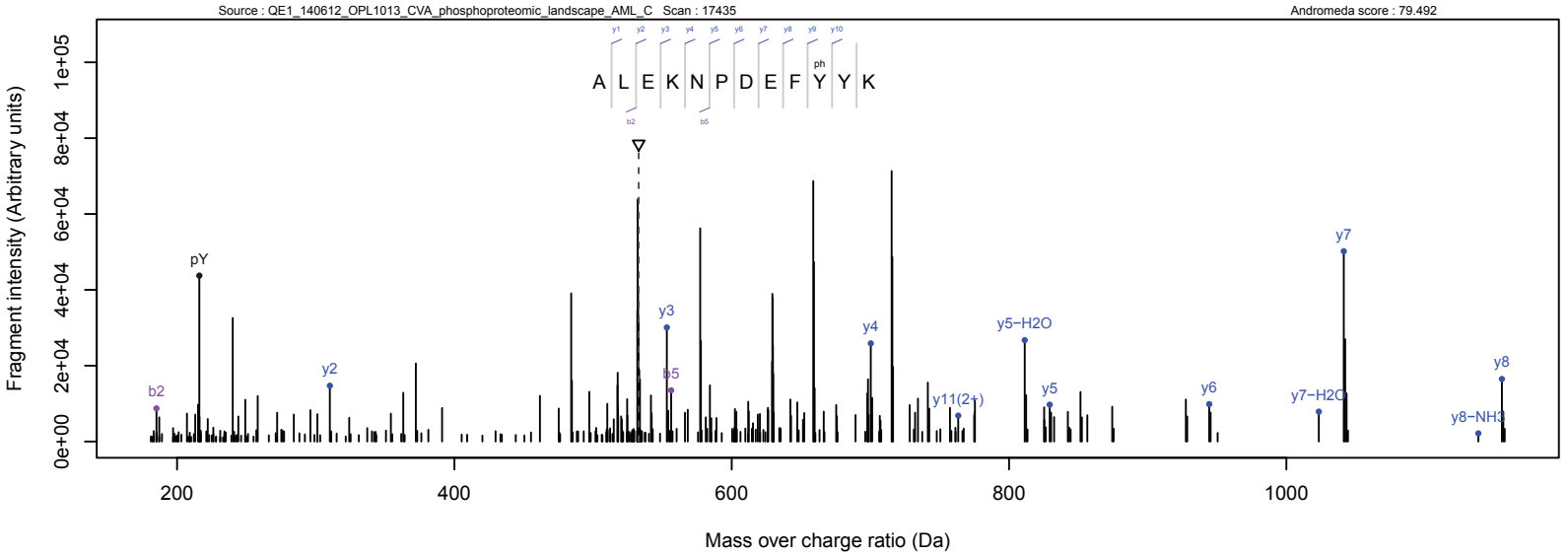
Andromeda score : 79.614



**ALEKNPDEFY(ph)YK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17435

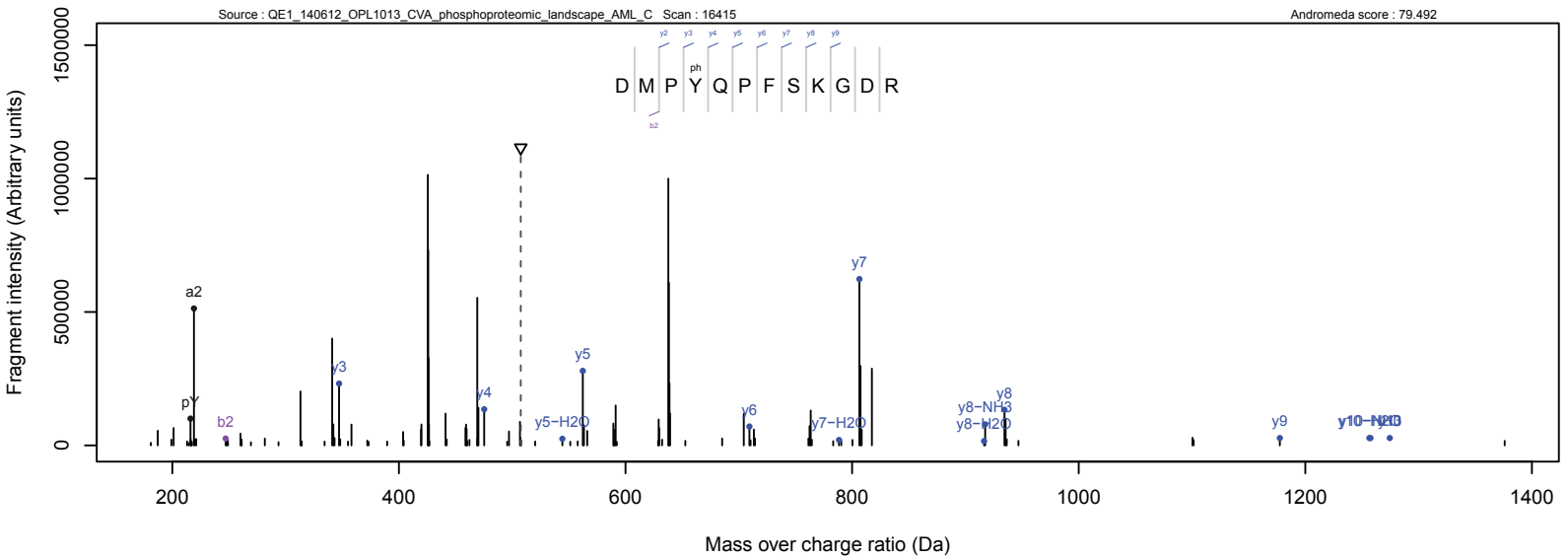
Andromeda score : 79.492



**DMPY(ph)QPFSGDR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16415

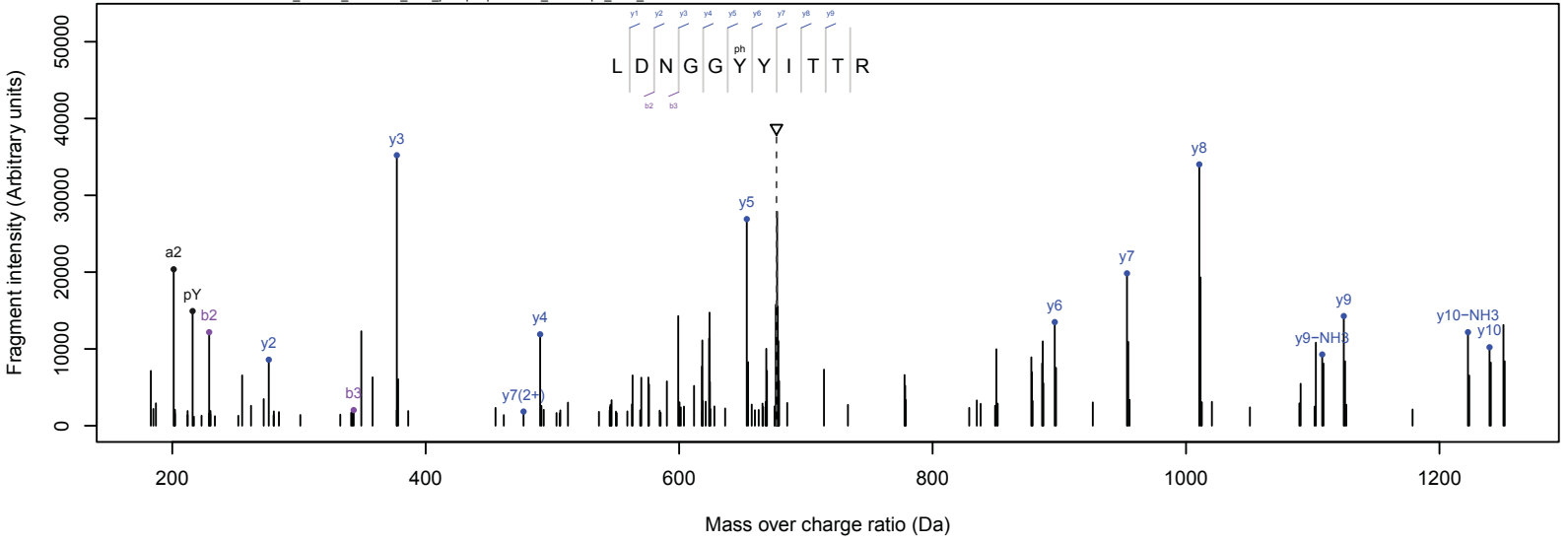
Andromeda score : 79.492



**LDNGGY(ph)YITTR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16026

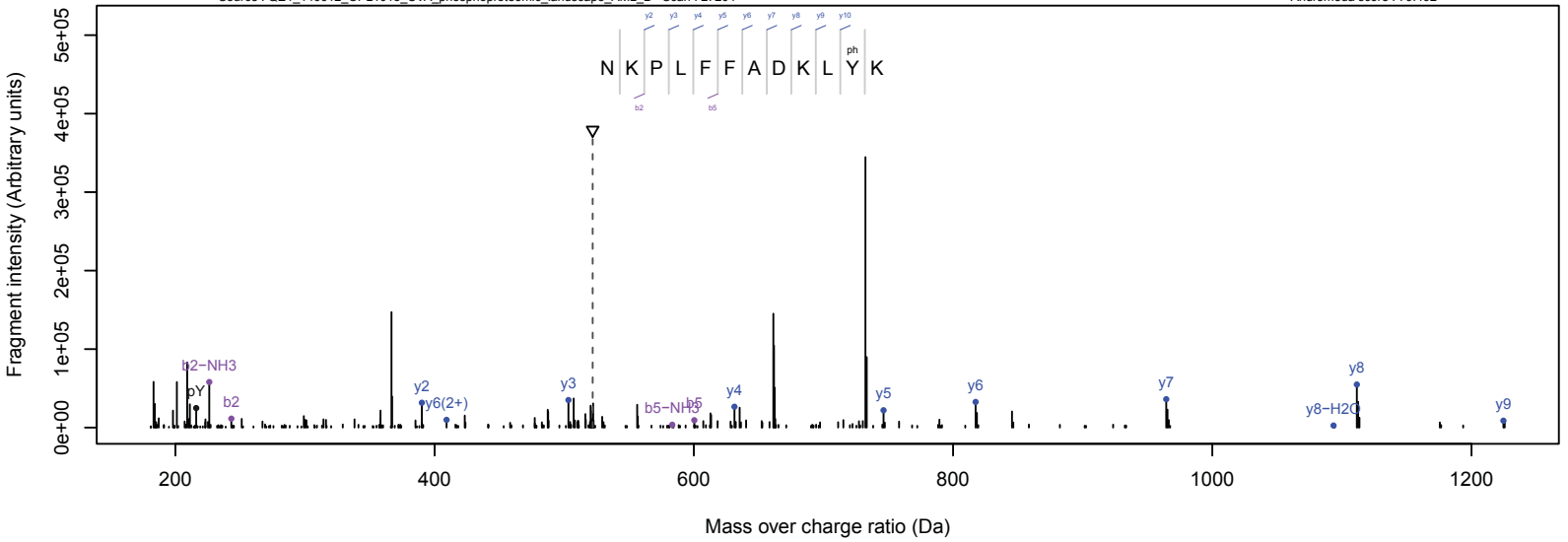
Andromeda score : 79.492



**NKPLFFADKLY(ph)K-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27204

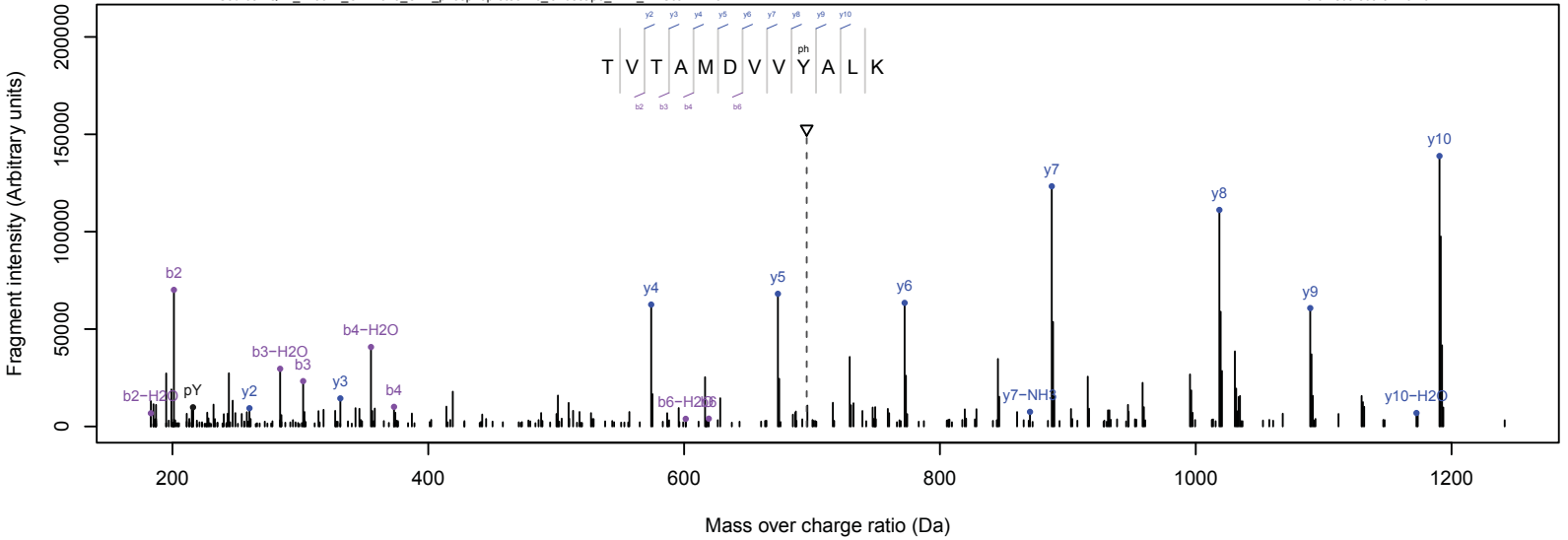
Andromeda score : 79.492



**TVTAMDVVY(ph)ALK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 44074

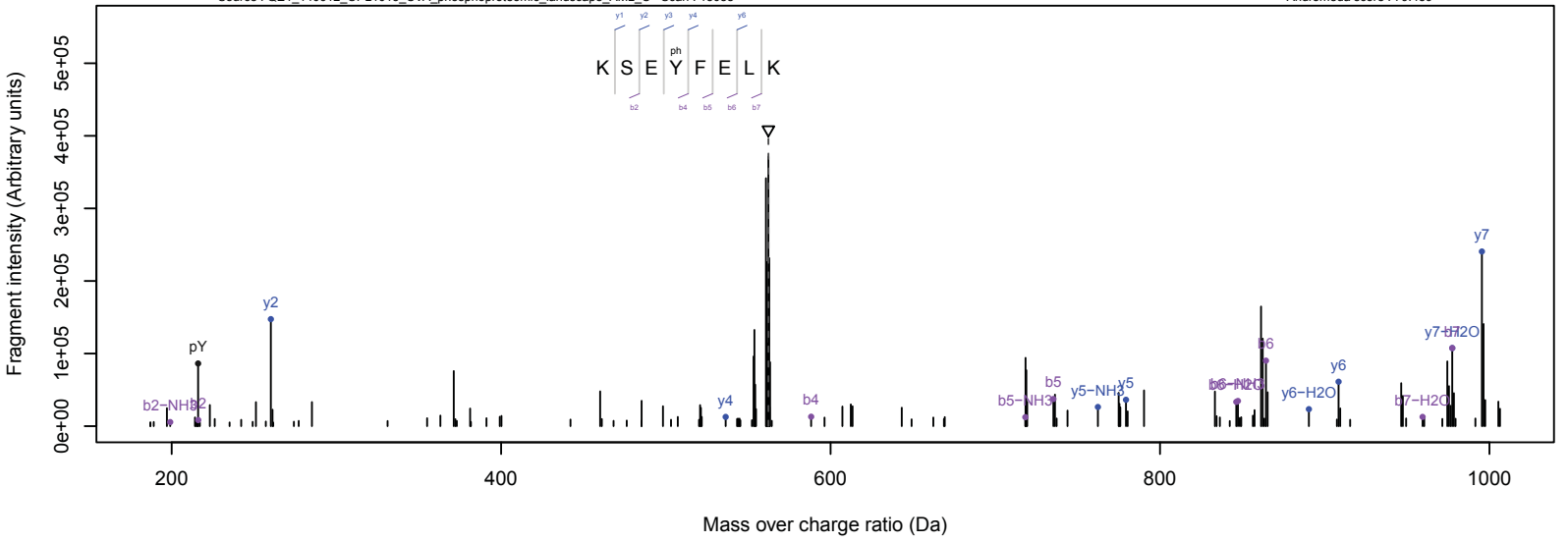
Andromeda score : 79.492



**KSEY(ph)FELK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15088

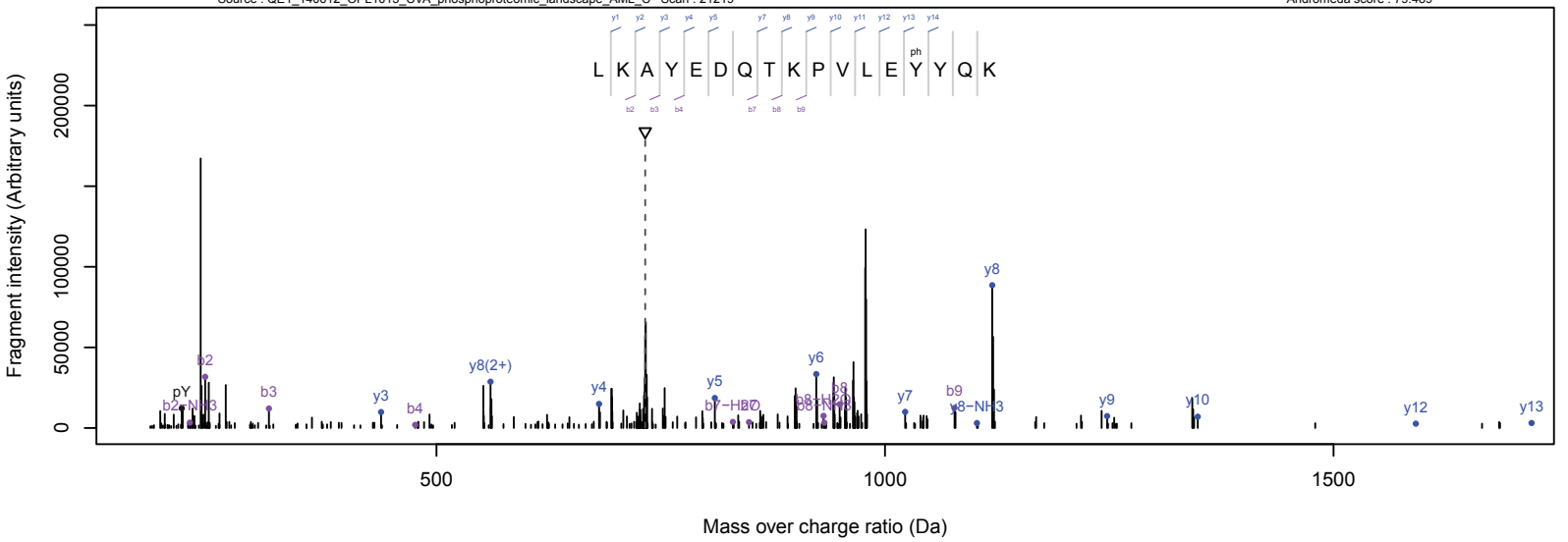
Andromeda score: 79.489



**LKAYEDQTKPVL EY YQK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21219

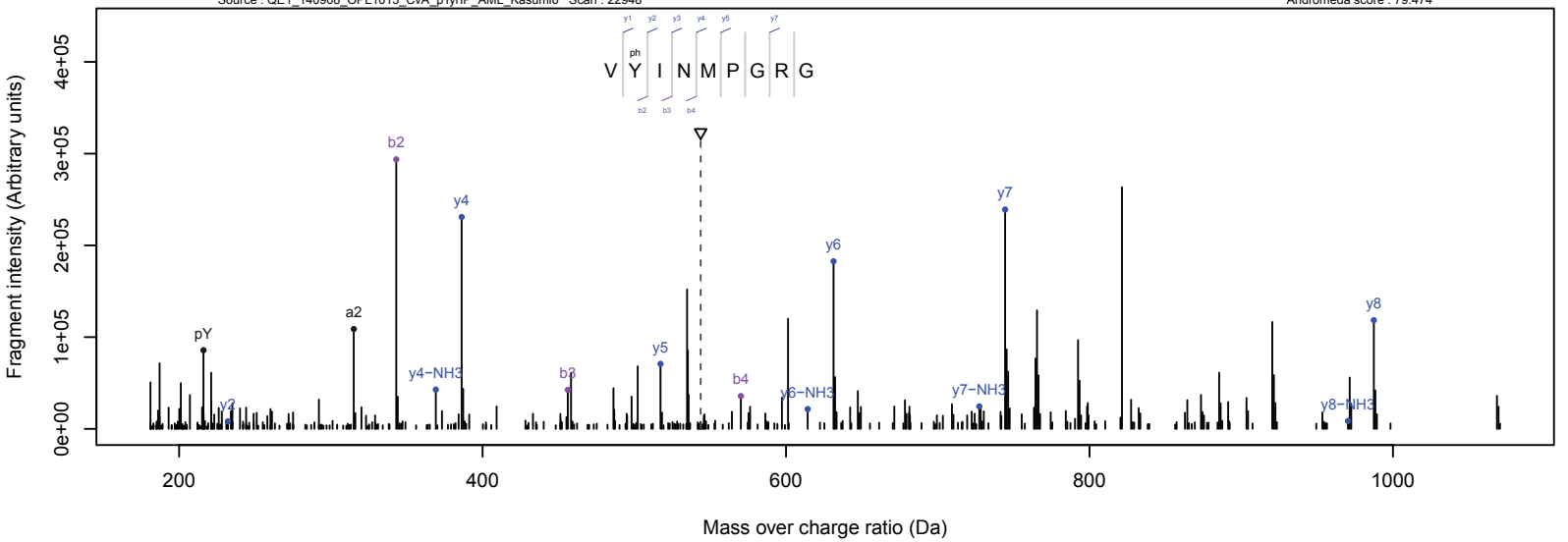
Andromeda score: 79.489



**VY(ph)INMPGRG-2+**

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 22948

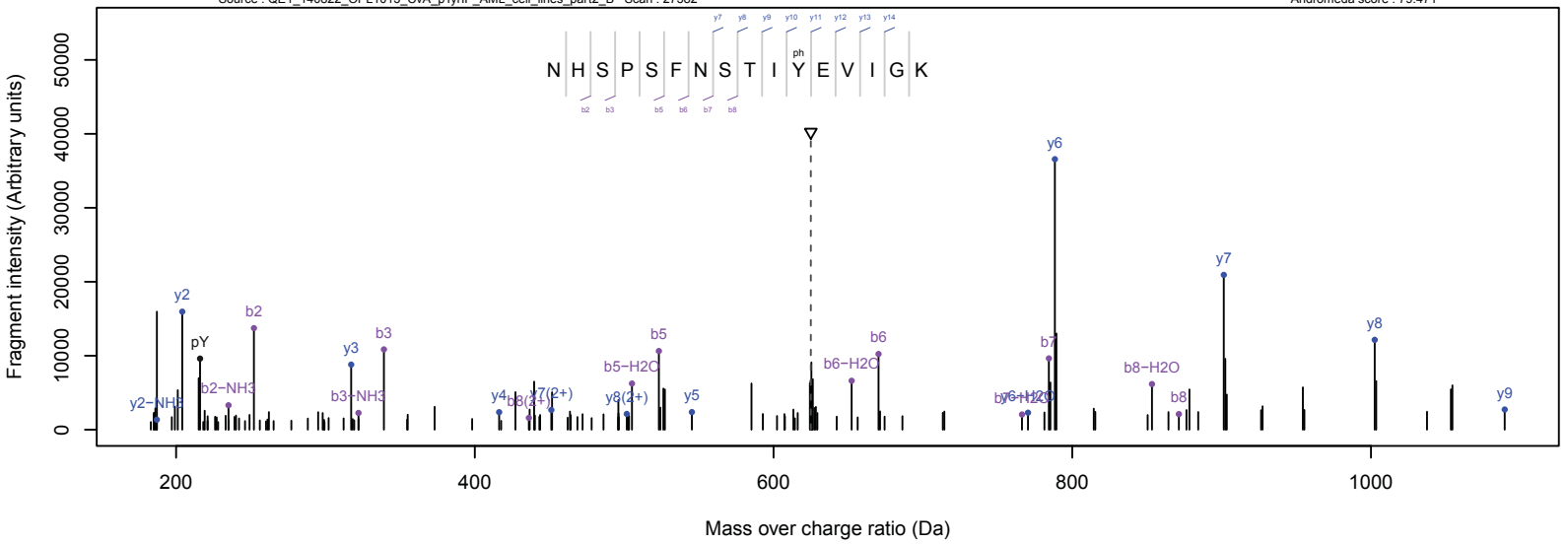
Andromeda score: 79.474



NHSPSFNSTIY(ph)EVIGK-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 27382

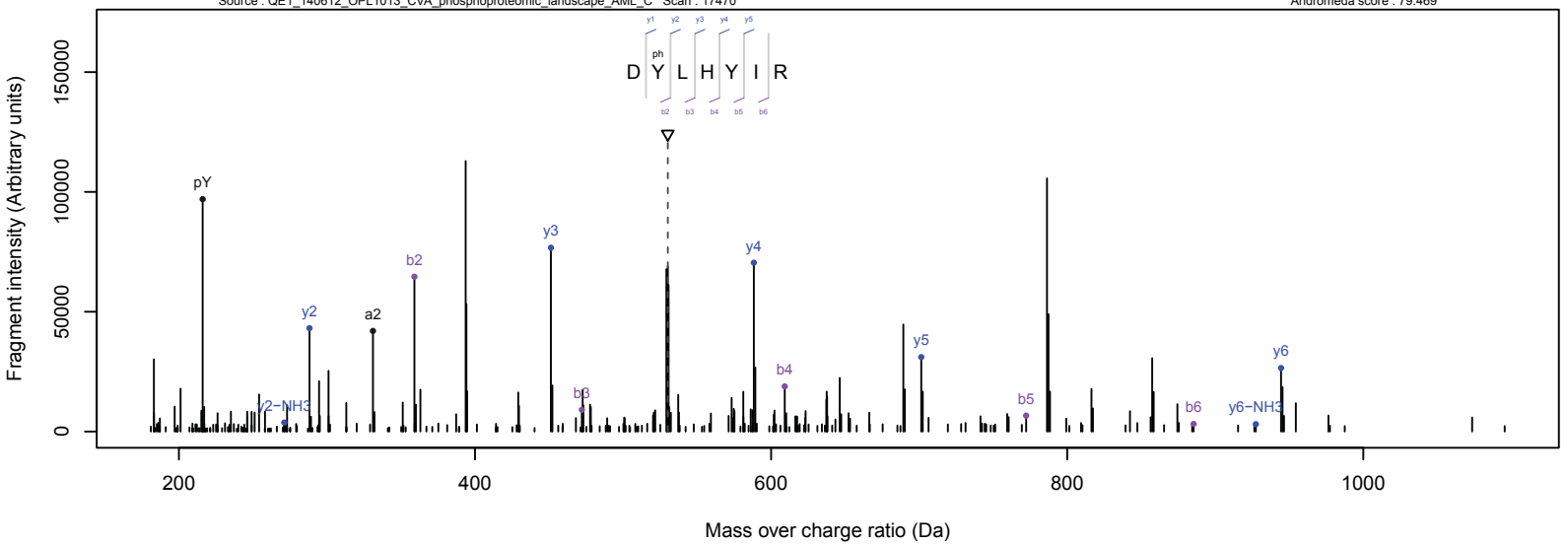
Andromeda score : 79.471



DY(ph)LHYIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17470

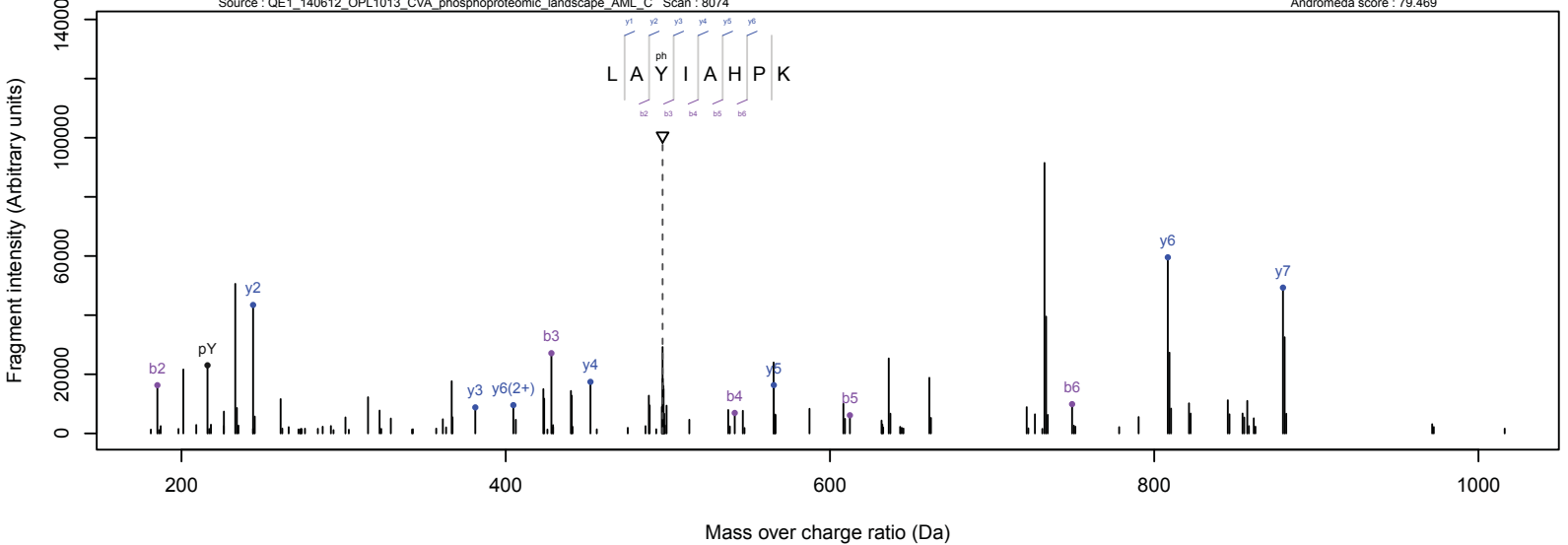
Andromeda score : 79.469



LAY(ph)IAHPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8074

Andromeda score : 79.469

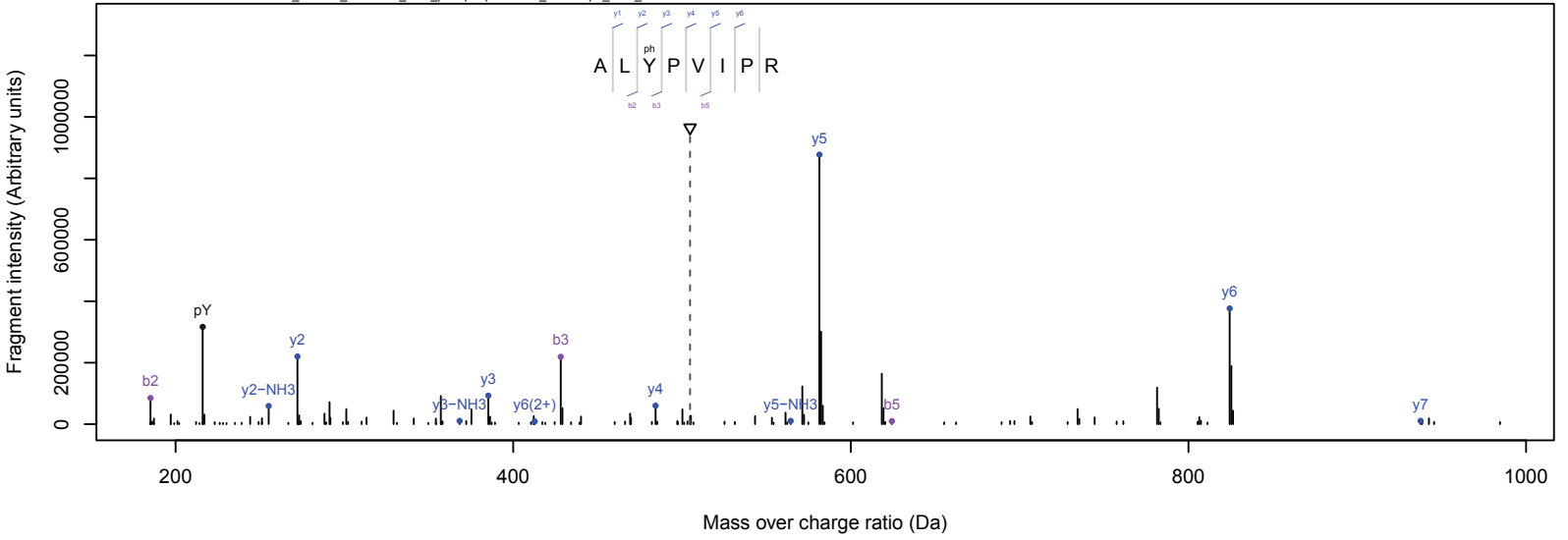




ALY(ph)PVIPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 26351

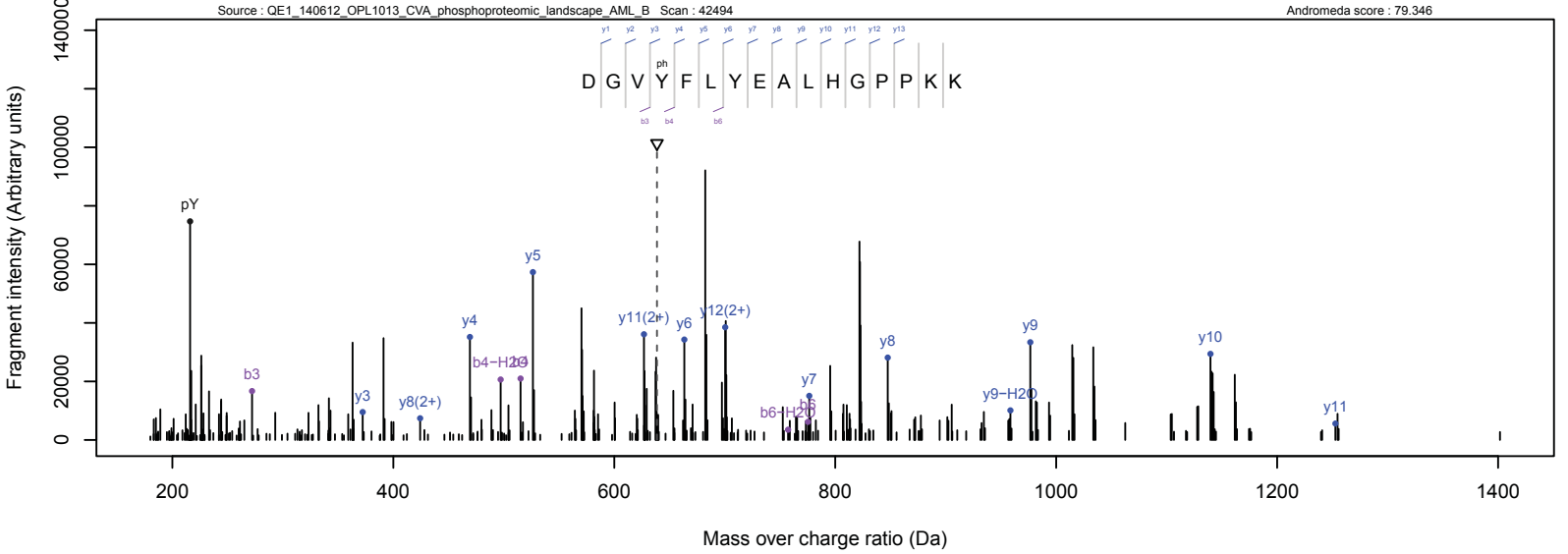
Andromeda score: 79.451



DGVY(ph)FLYEALHGPPK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 42494

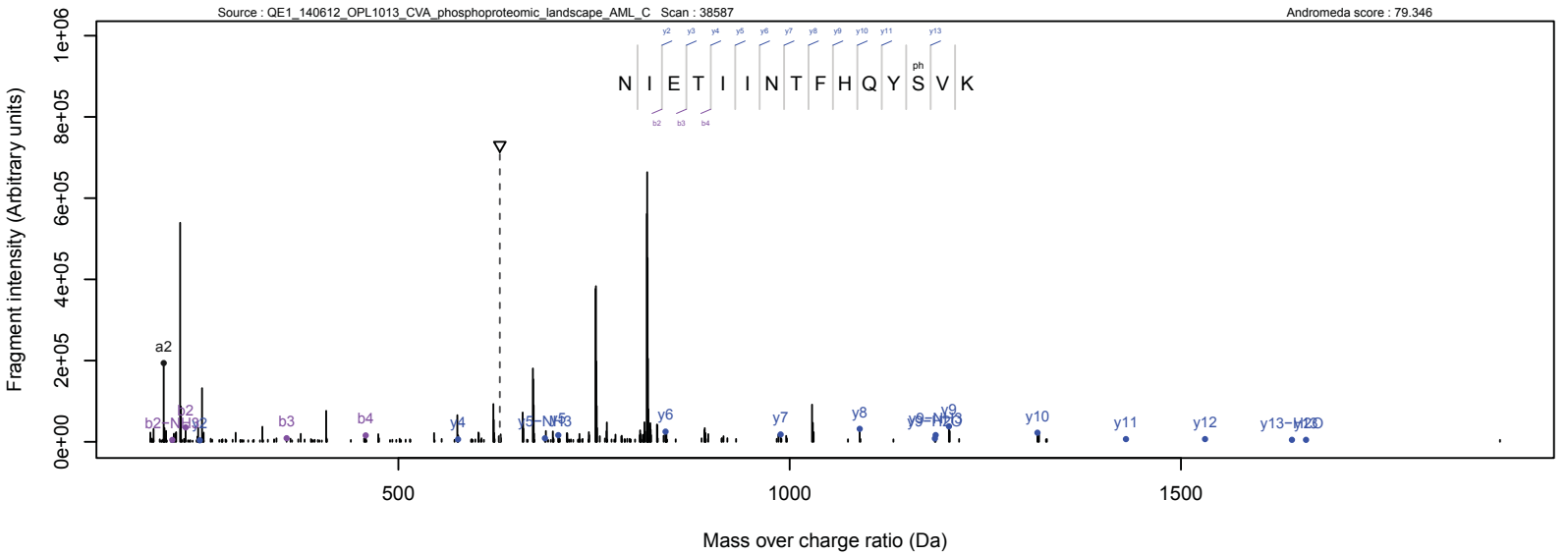
Andromeda score: 79.346



NIETINTFHQYS(ph)VK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 38587

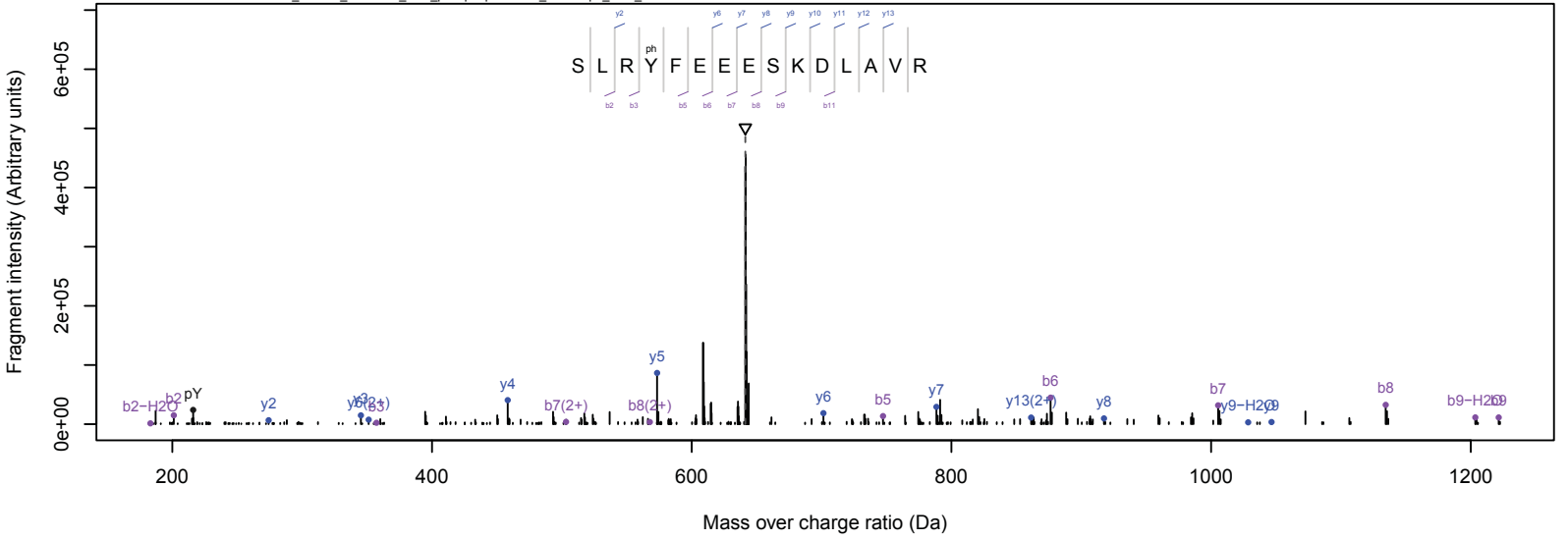
Andromeda score: 79.346



SLRY(ph)FEEESKDLAVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26964

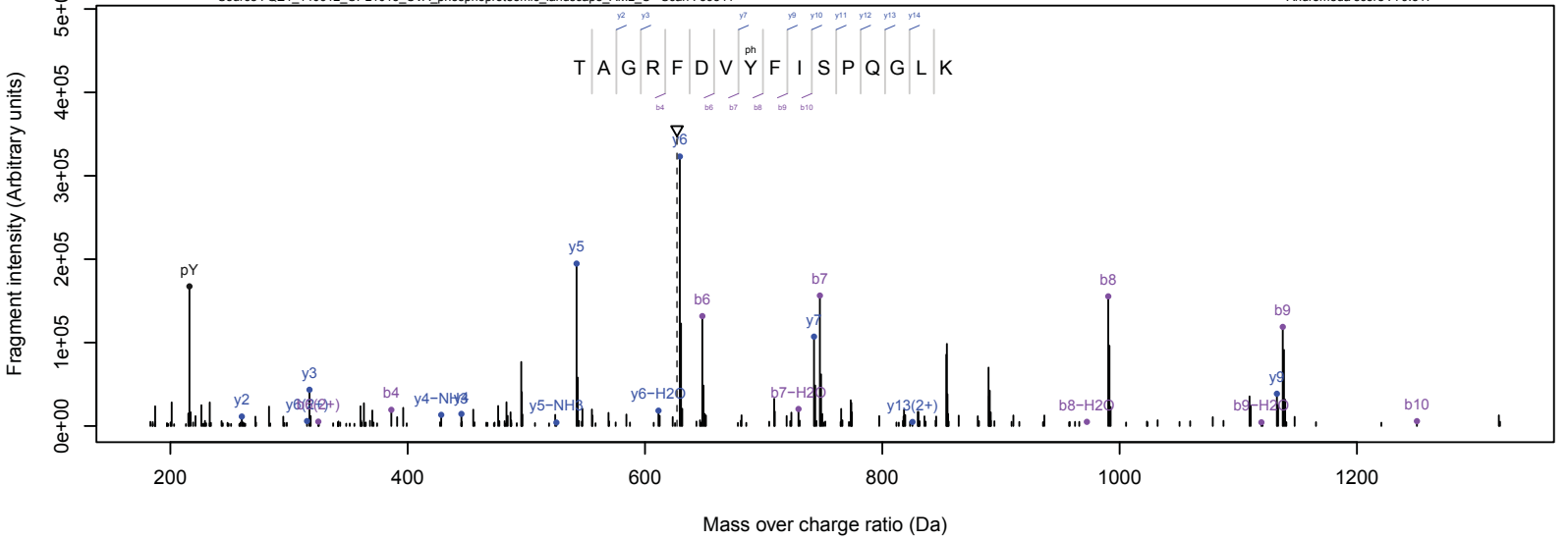
Andromeda score : 79.317



TAGRFDVY(ph)FISPQGLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39541

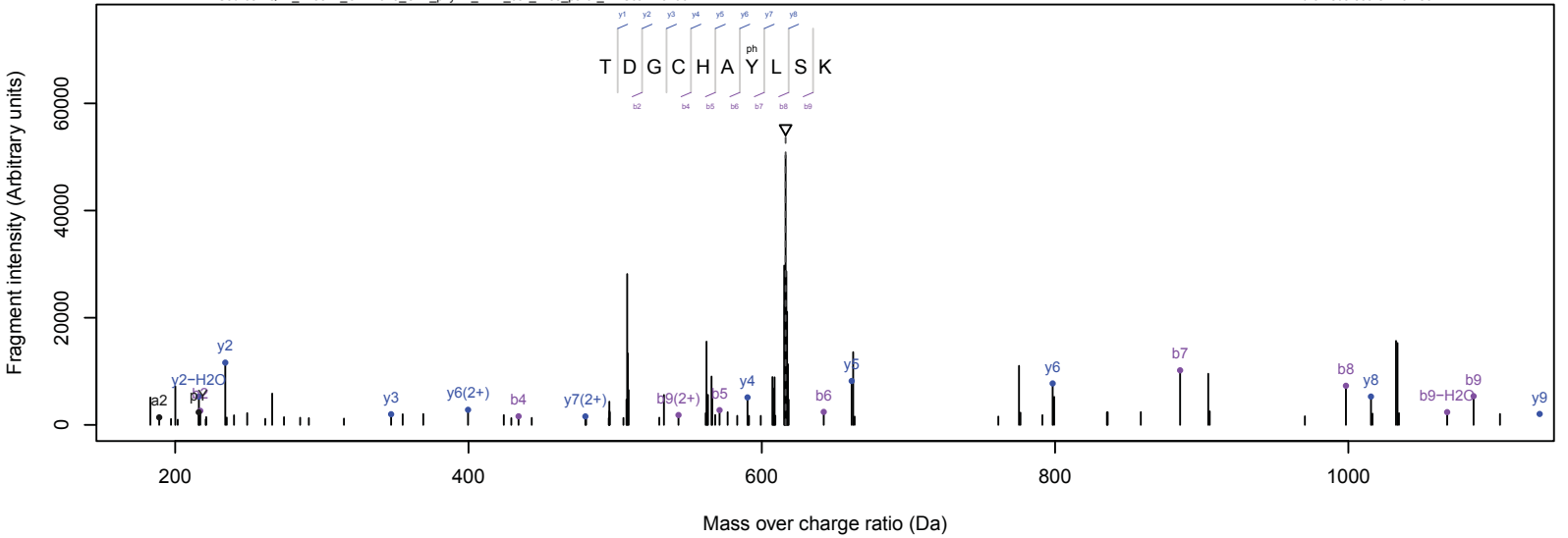
Andromeda score : 79.317



TDGCHAYLISK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3296

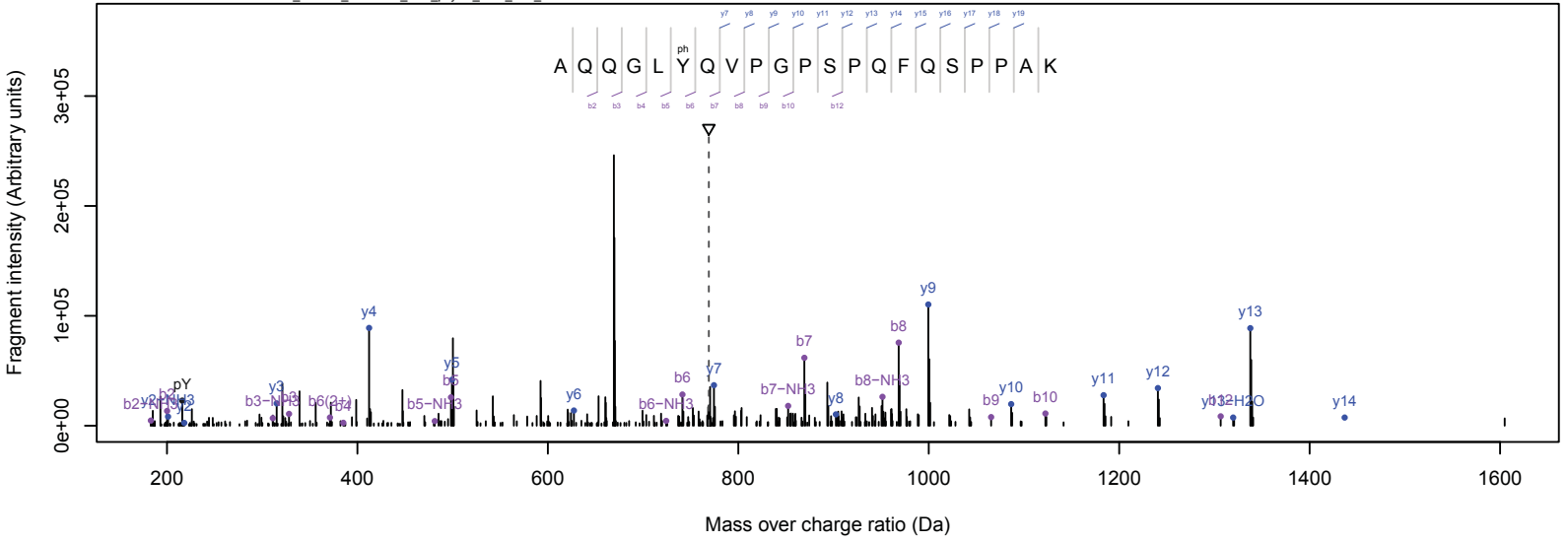
Andromeda score : 79.286



**AQQGLY(ph)QVGPSPQFQSPPAK-3+**

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_OCI\_AML3 Scan: 29216

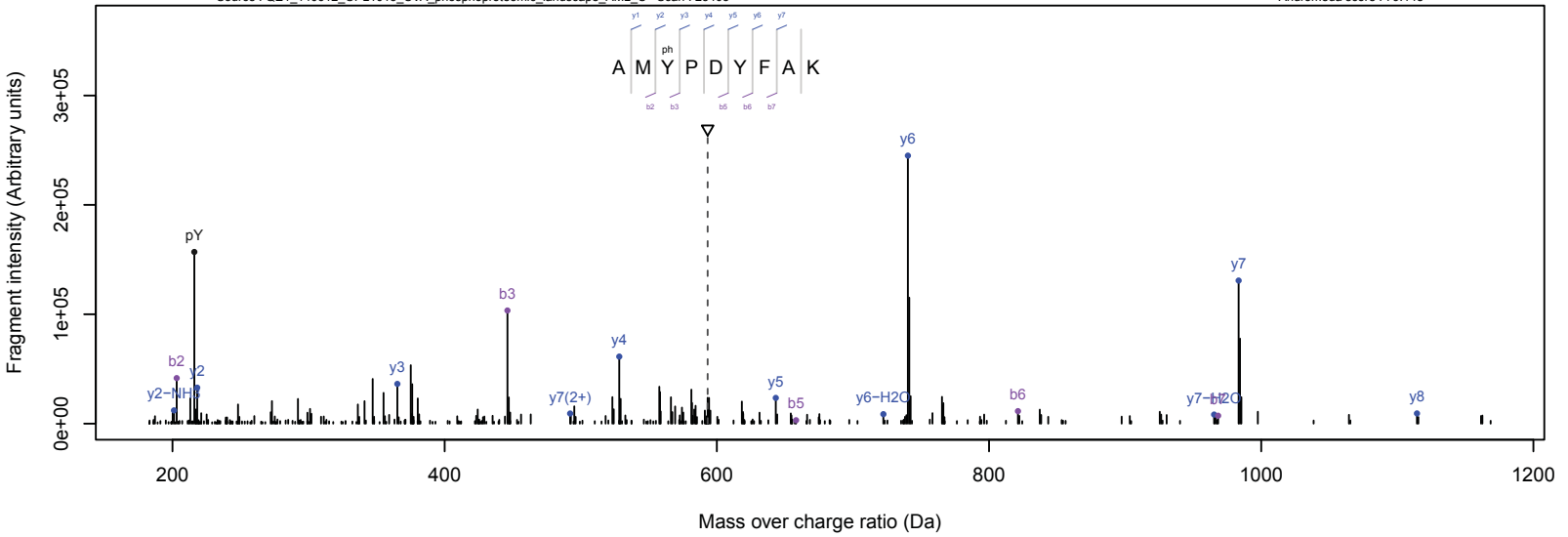
Andromeda score: 79.237



**AMY(ph)PDYFAK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25403

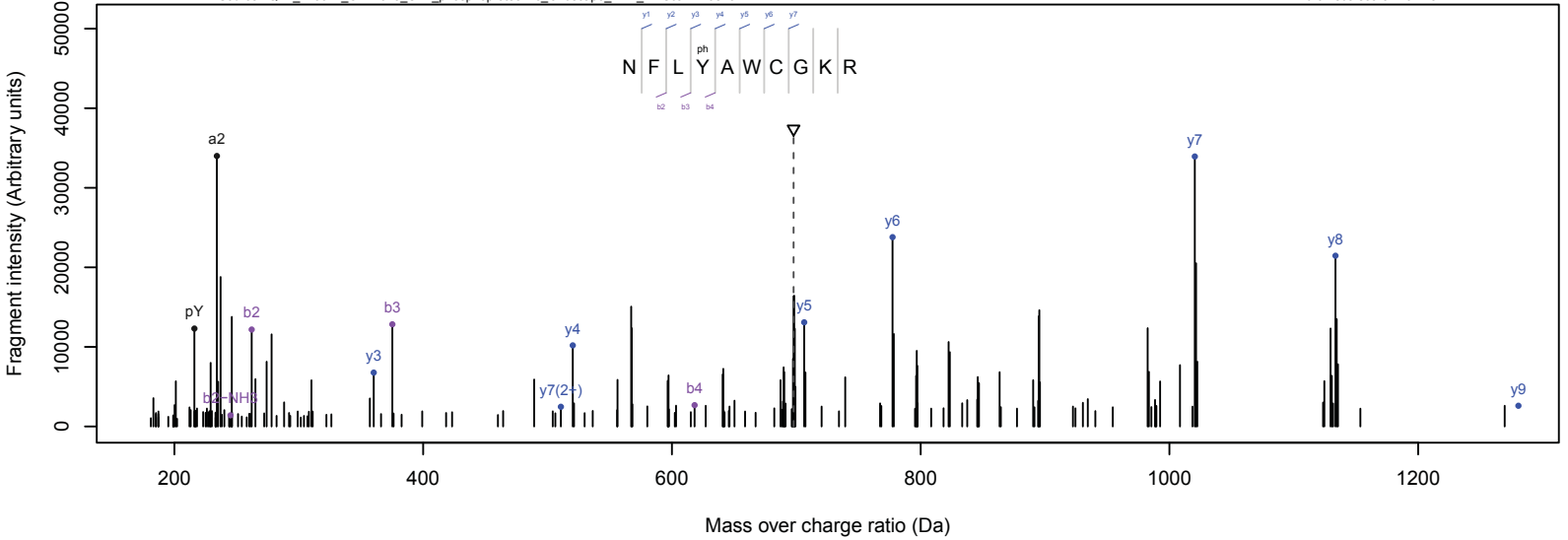
Andromeda score: 79.148



**NFLY(ph)AWCGKR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 29810

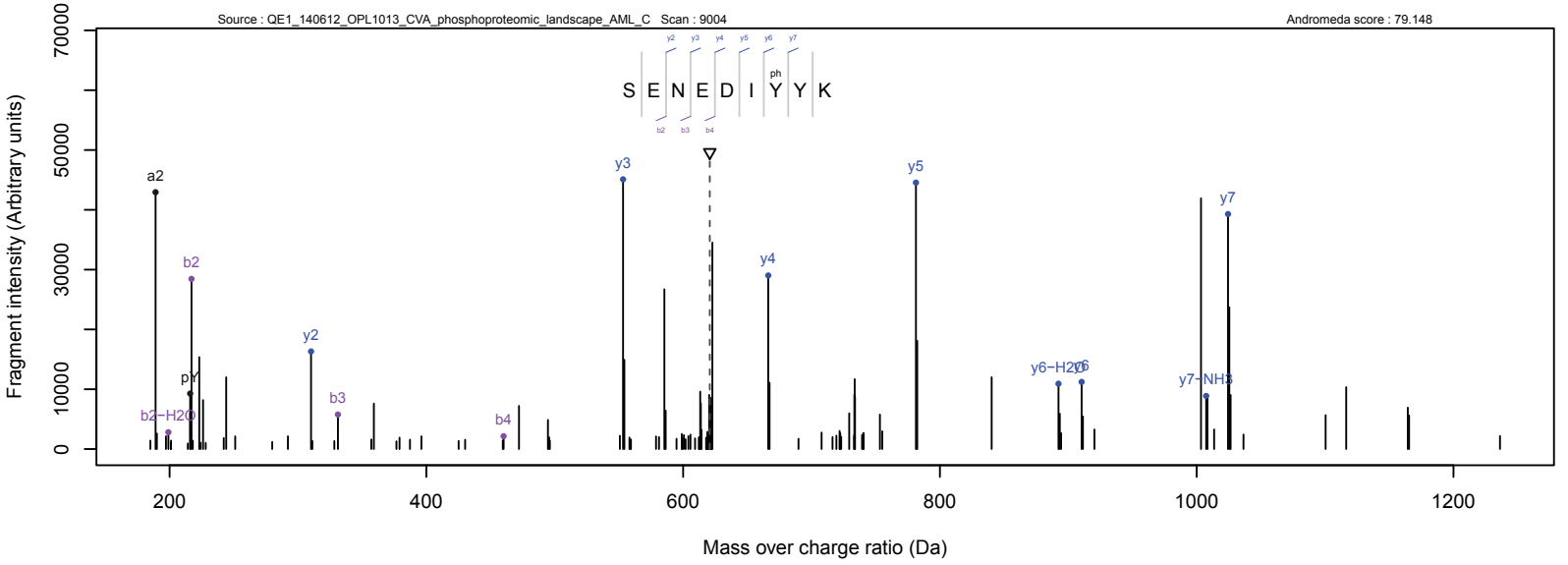
Andromeda score: 79.148



SENEDIY(ph)YK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9004

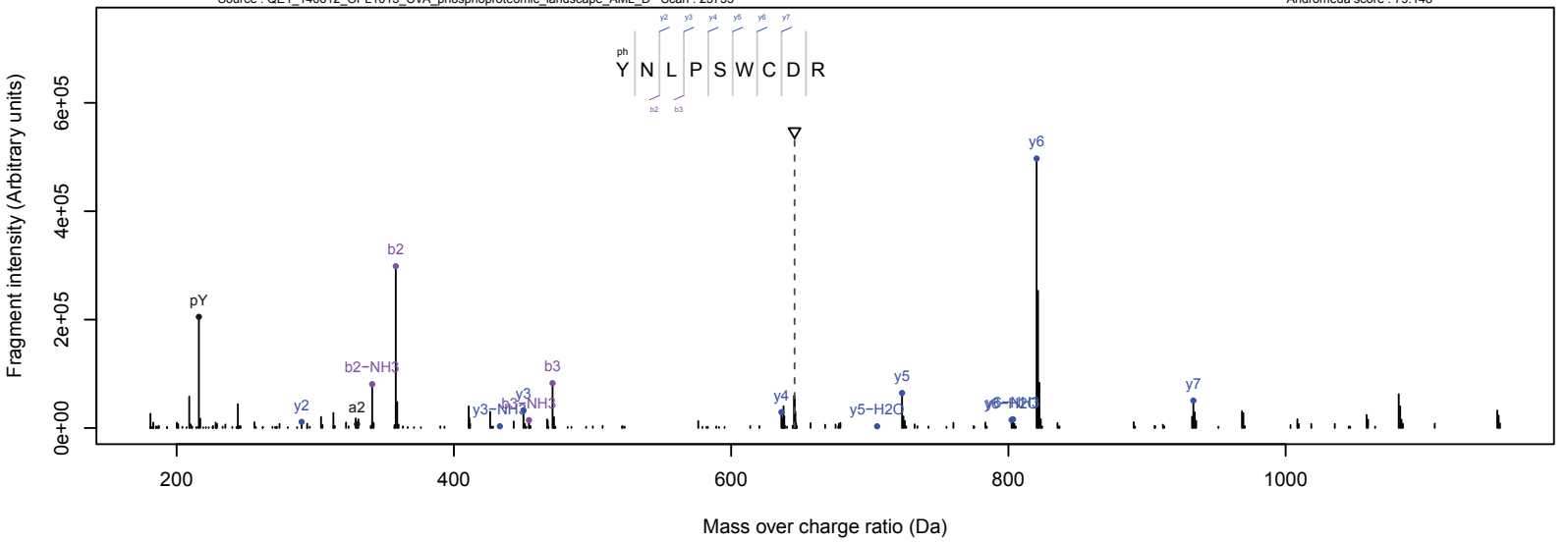
Andromeda score : 79.148



Y(ph)NLPWCDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23735

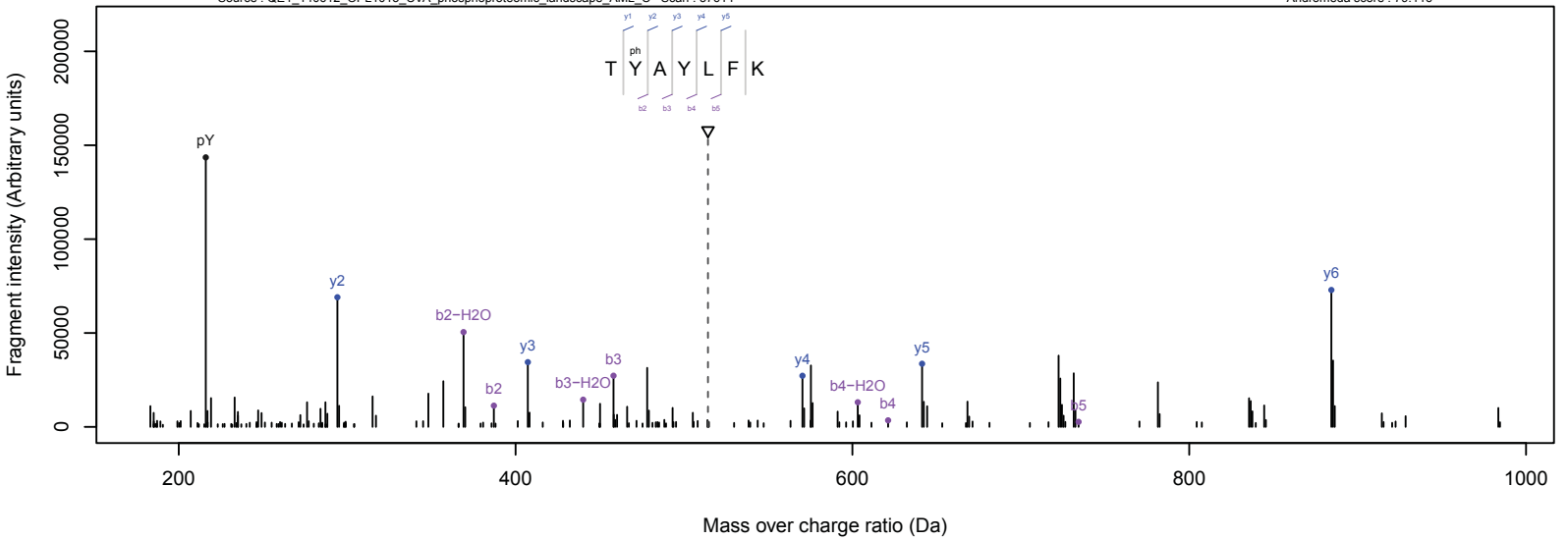
Andromeda score : 79.148



(ac)TY(ph)AYLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37614

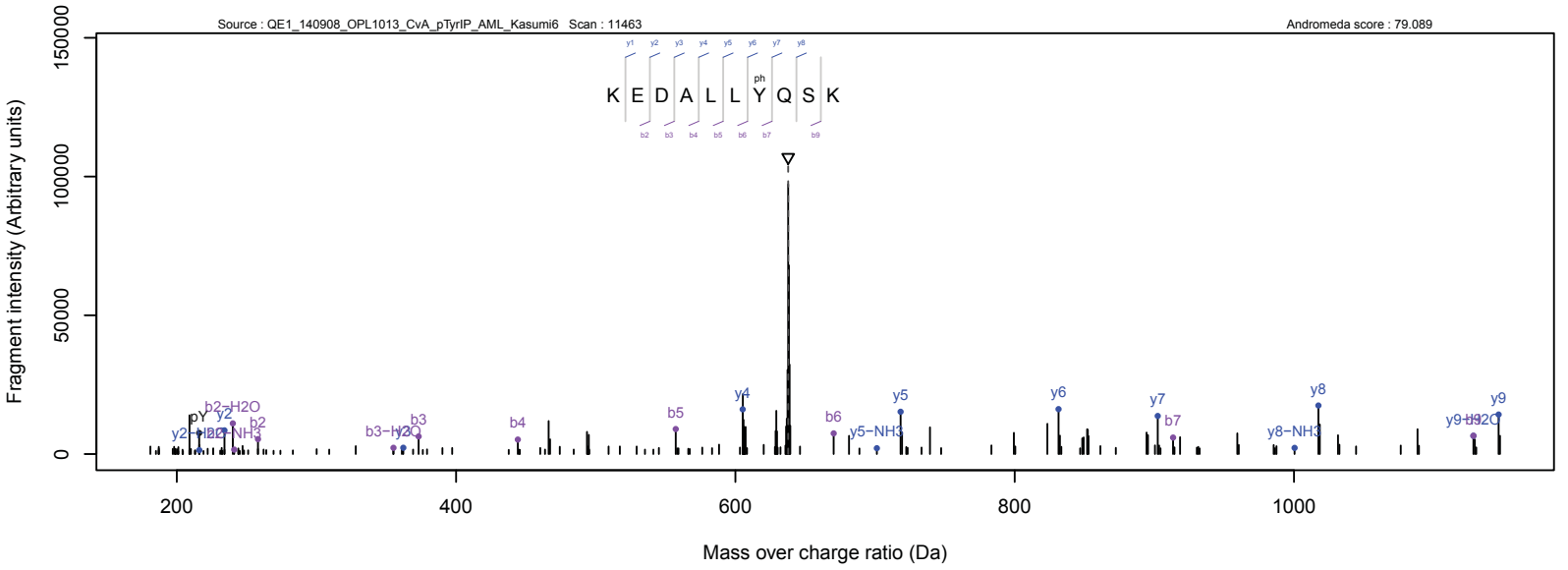
Andromeda score : 79.116



**KEDALLYQSK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 11463

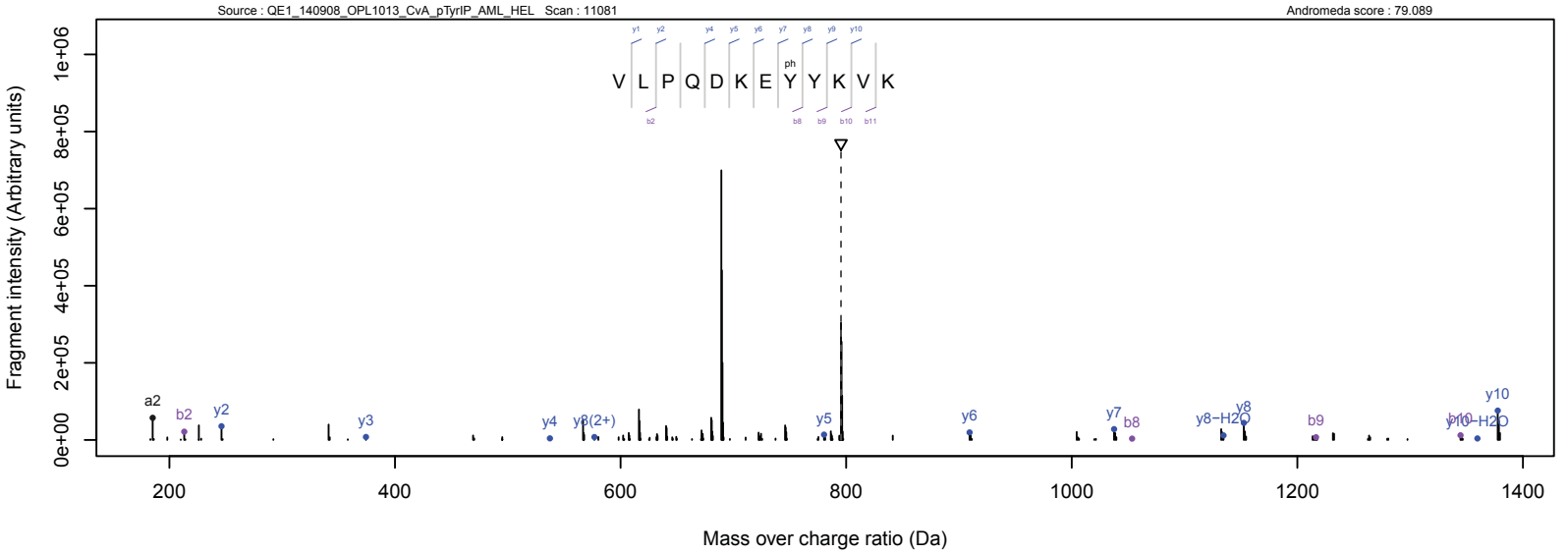
Andromeda score : 79.089



**VLPQDKEY(Y)KVK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 11081

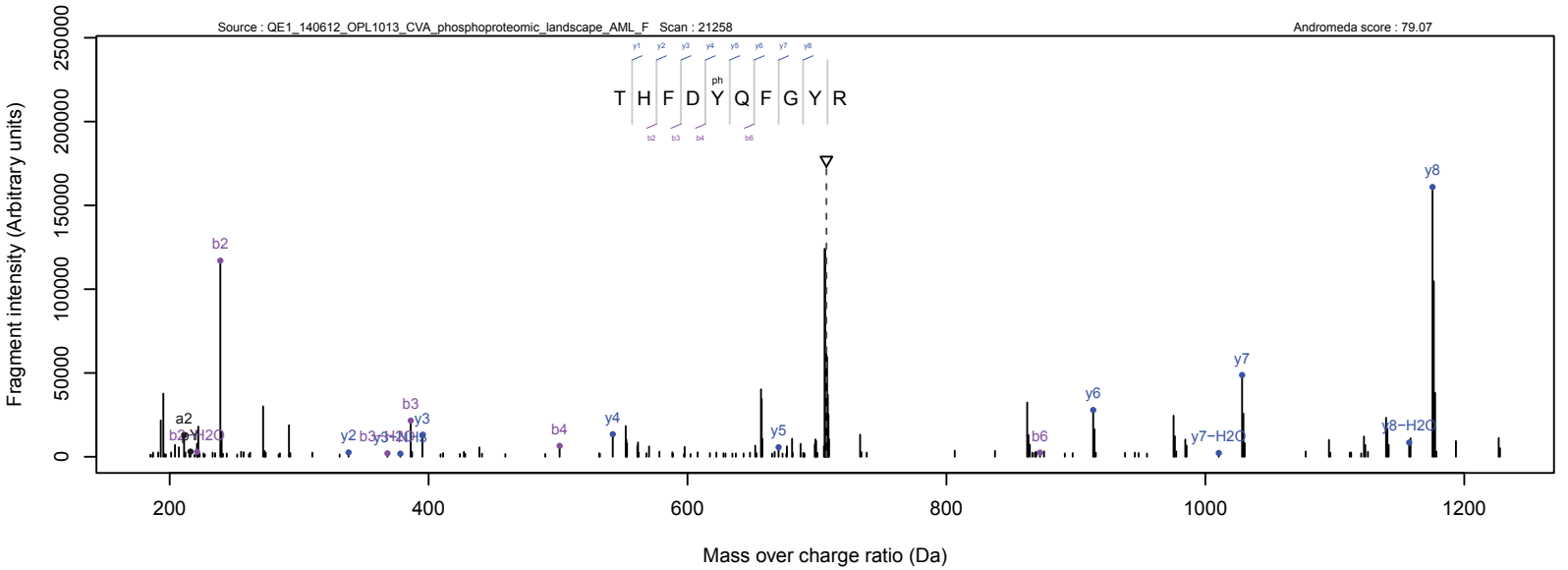
Andromeda score : 79.089



**THFDY(ph)QFGYR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 21258

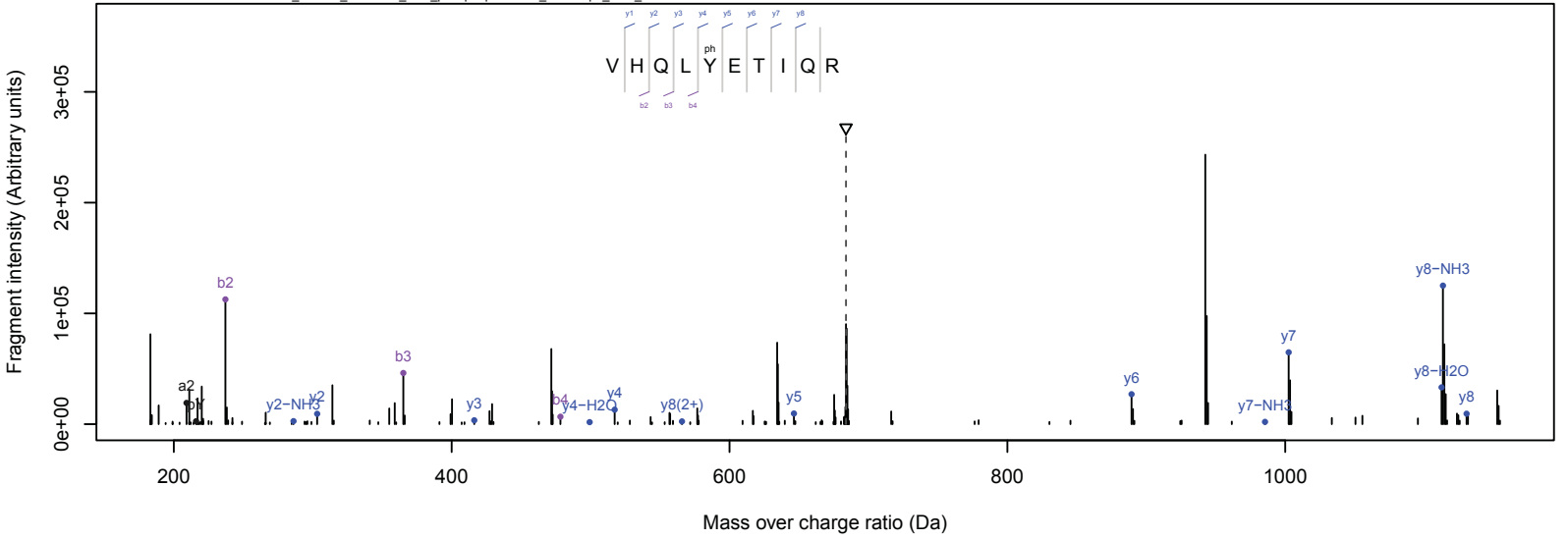
Andromeda score : 79.07



VHQLY(ph)ETIQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12138

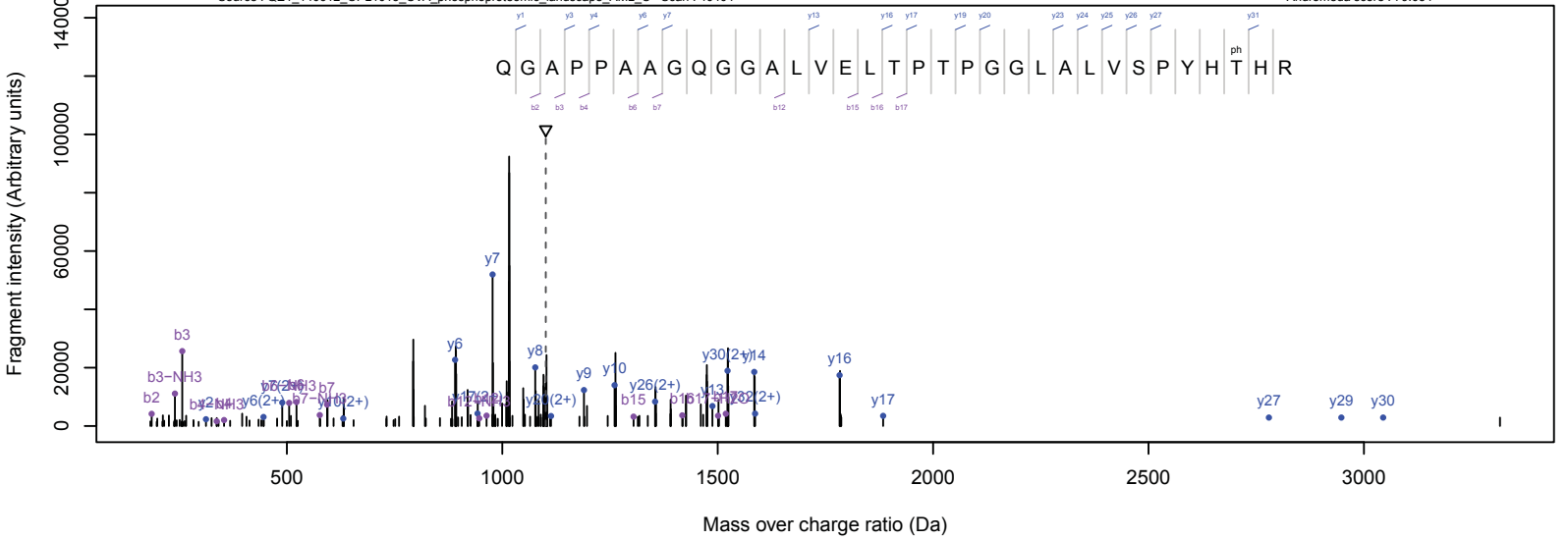
Andromeda score : 79.07



QGAPPAAGGGALVELTPTPGGLALVSPYHT(ph)HR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40404

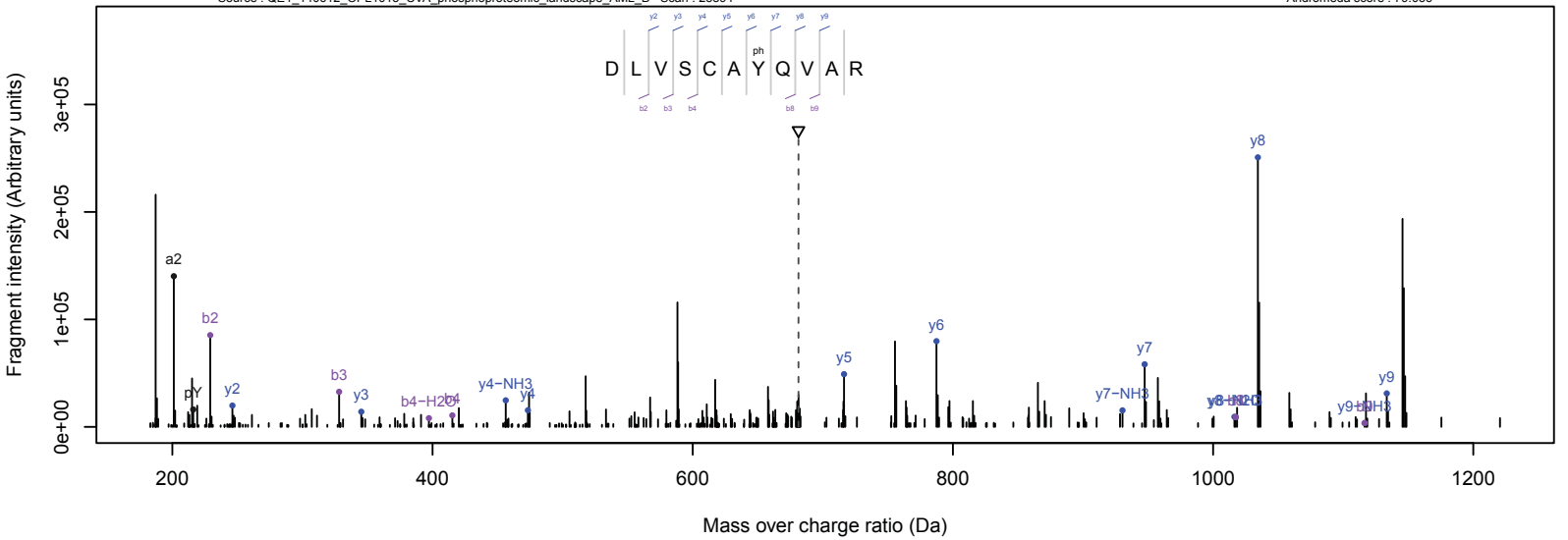
Andromeda score : 79.054



DLVSCAY(ph)QVAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25894

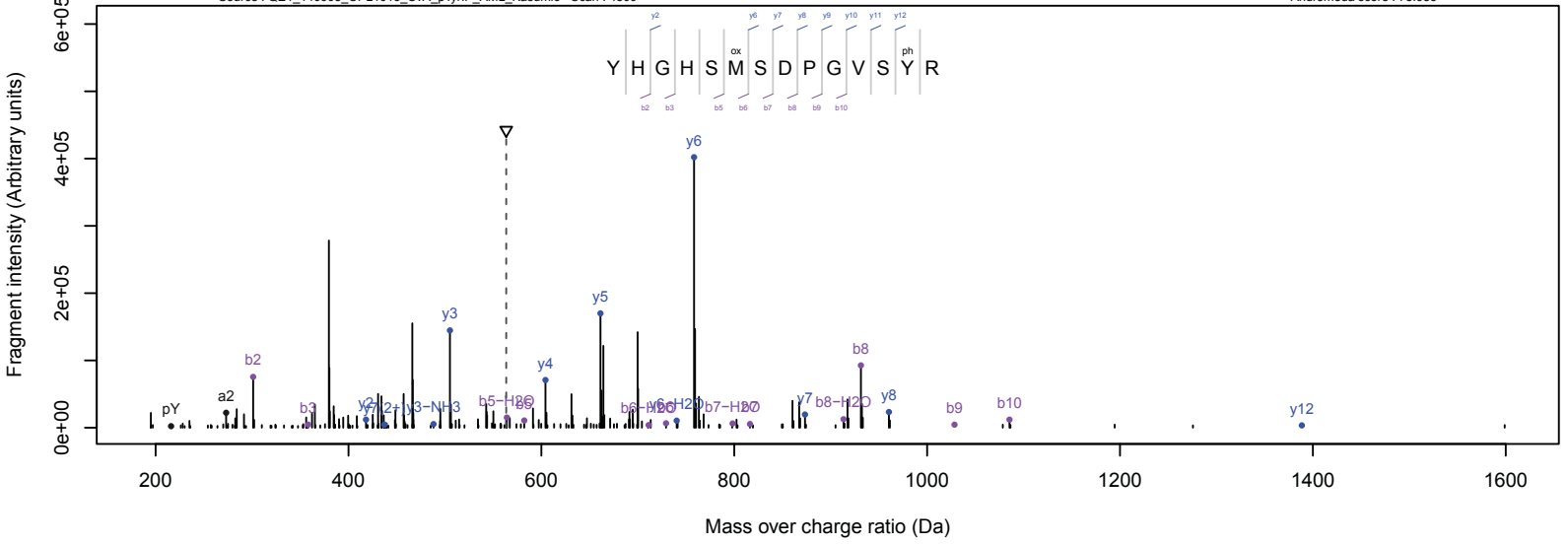
Andromeda score : 79.036



**YHGHSM(ox)SDPGVSY(ph)R-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 4509

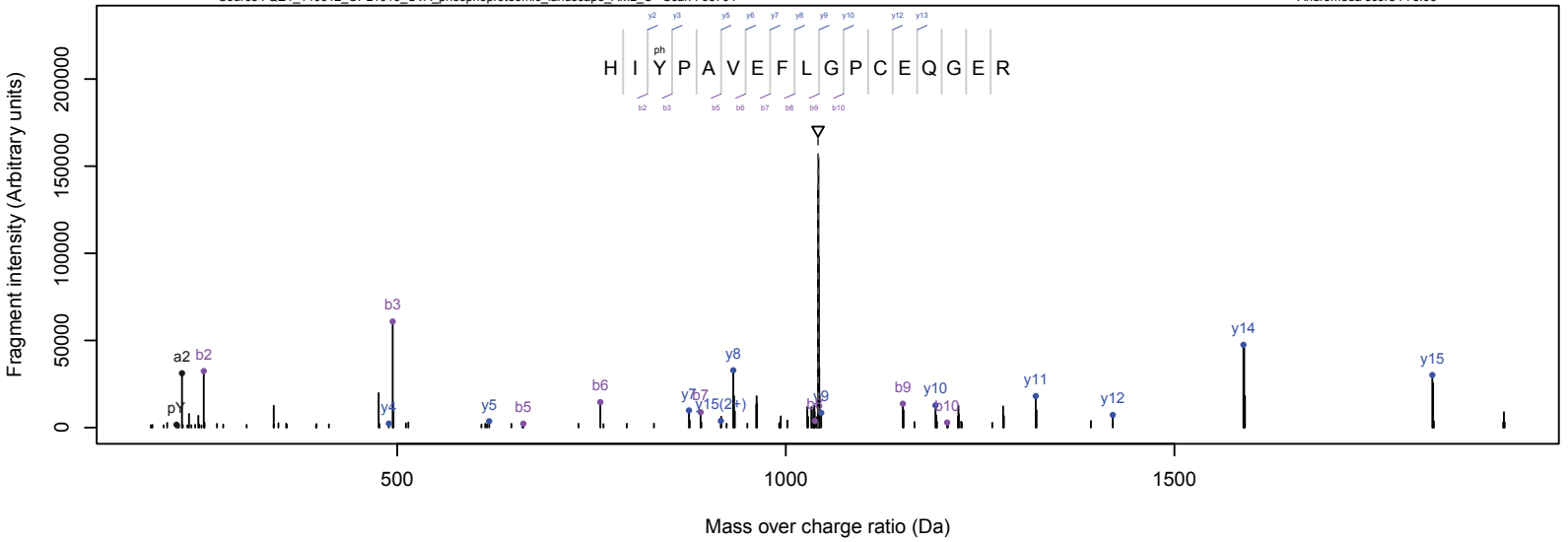
Andromeda score : 78.985



**HIY(ph)PAVEFLGPCEQGER-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33704

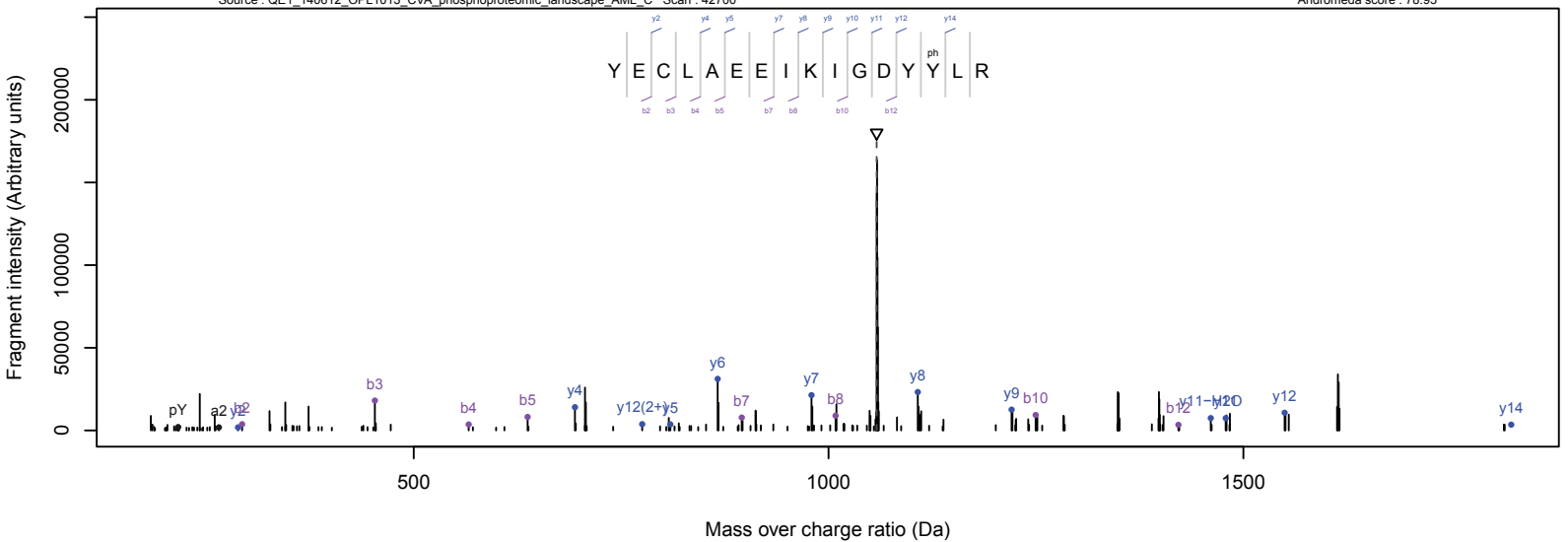
Andromeda score : 78.95



**YECLAEEIKIGDY(ph)LR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42700

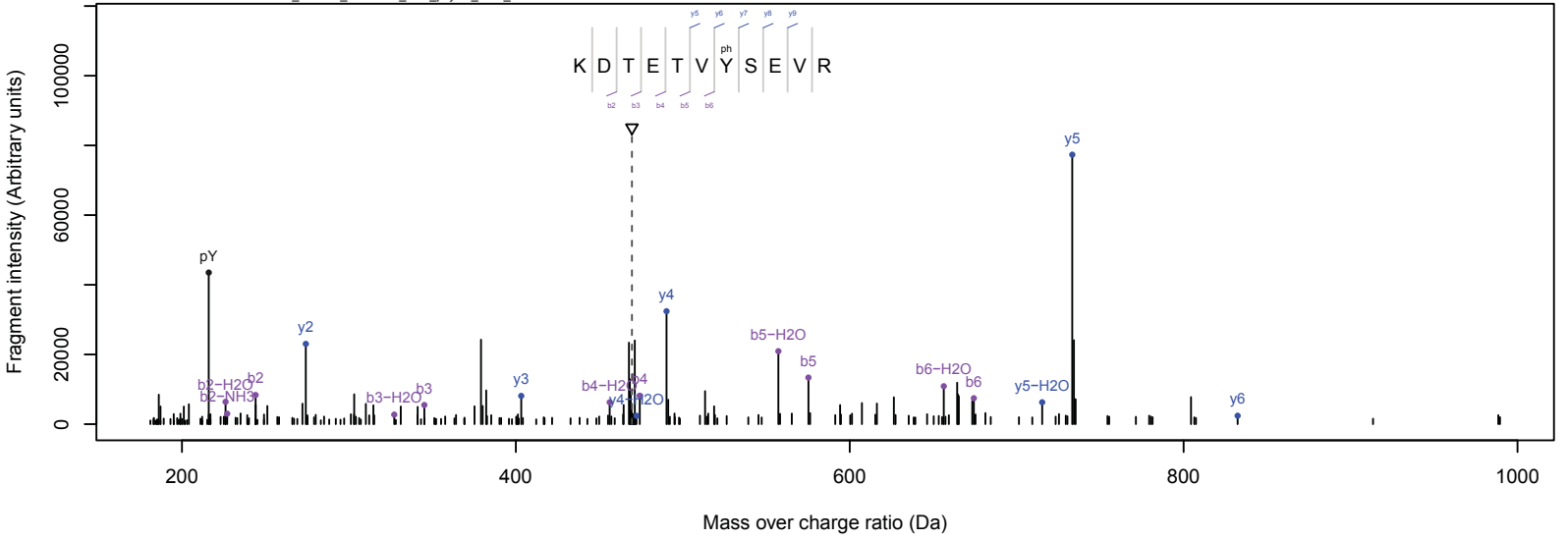
Andromeda score : 78.95



**KDTETV(ph)SEVR-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 7844

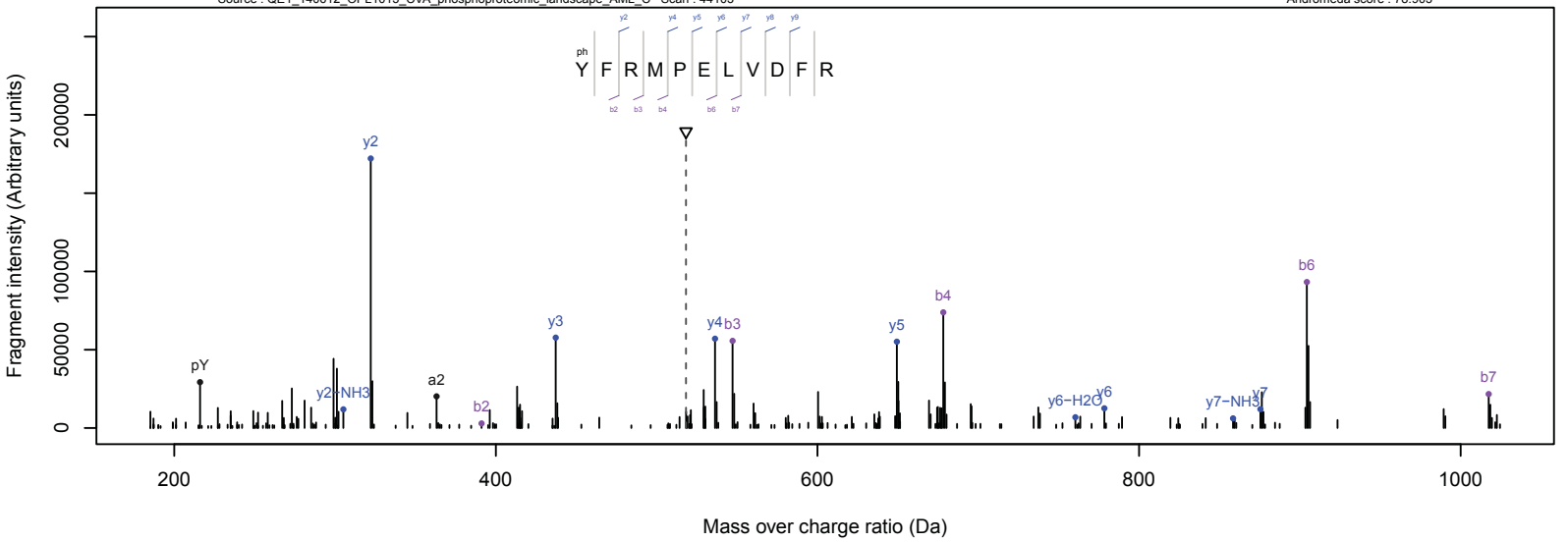
Andromeda score : 78.903



**Y(ph)FRMPELVDFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44103

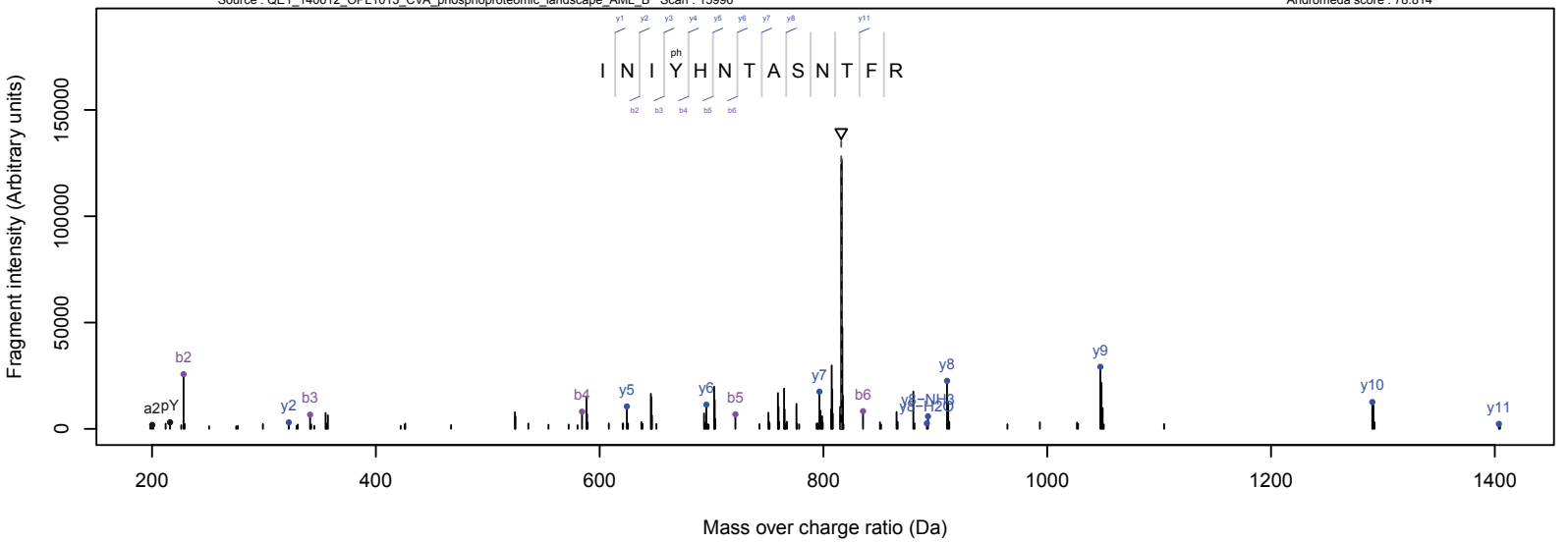
Andromeda score : 78.903



**INIIYHNTASNTFR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15996

Andromeda score : 78.814

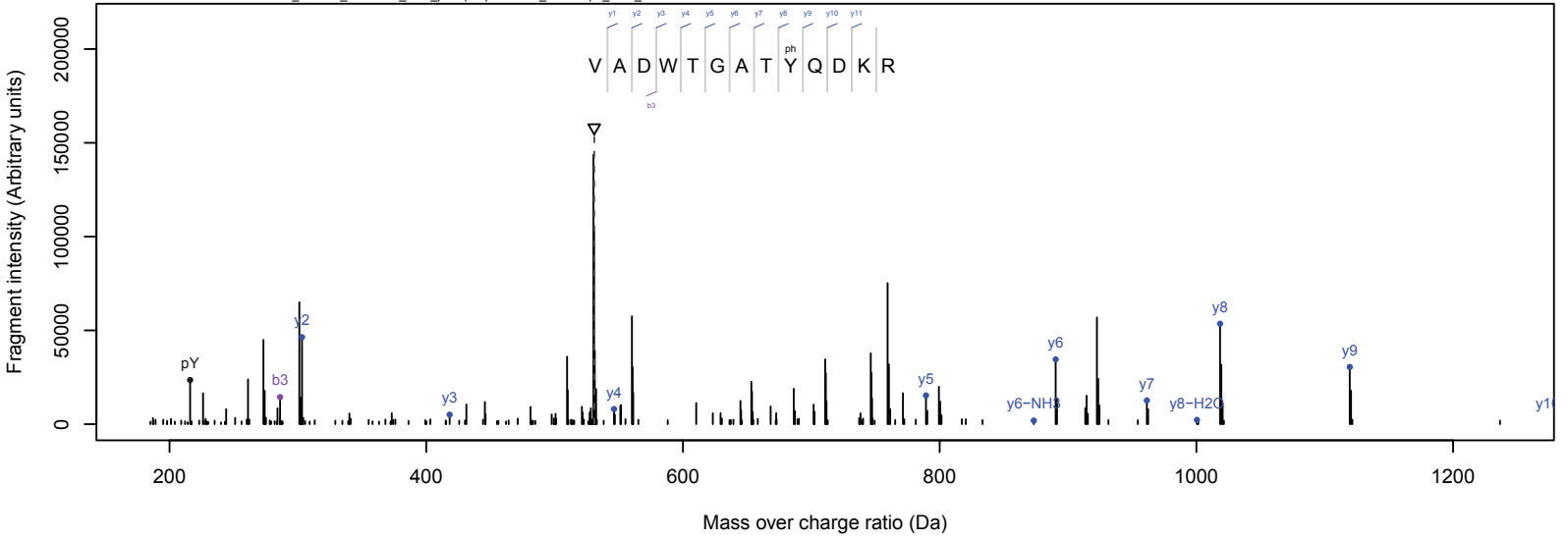




VADWTGATY(ph)QDKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10926

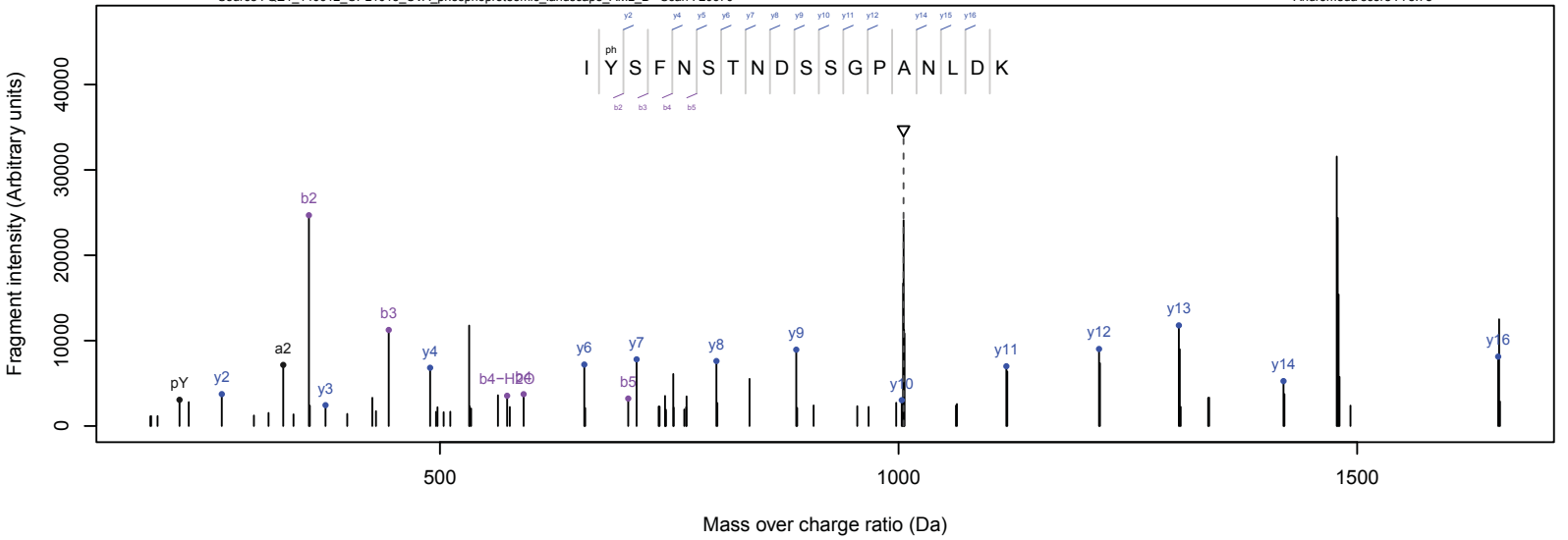
Andromeda score : 78.814



IY(ph)SFNSTNDSSGPANLDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20670

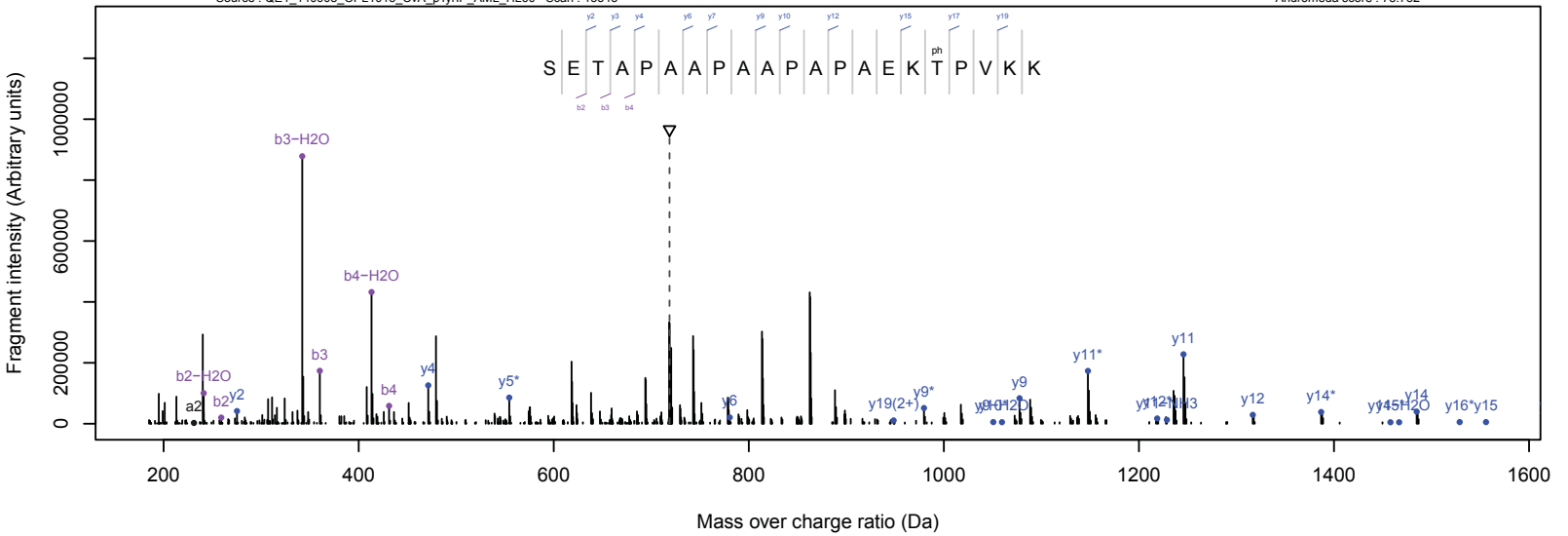
Andromeda score : 78.78



(ac)SETAPAAPAAPAEK(ph)PVKK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 15543

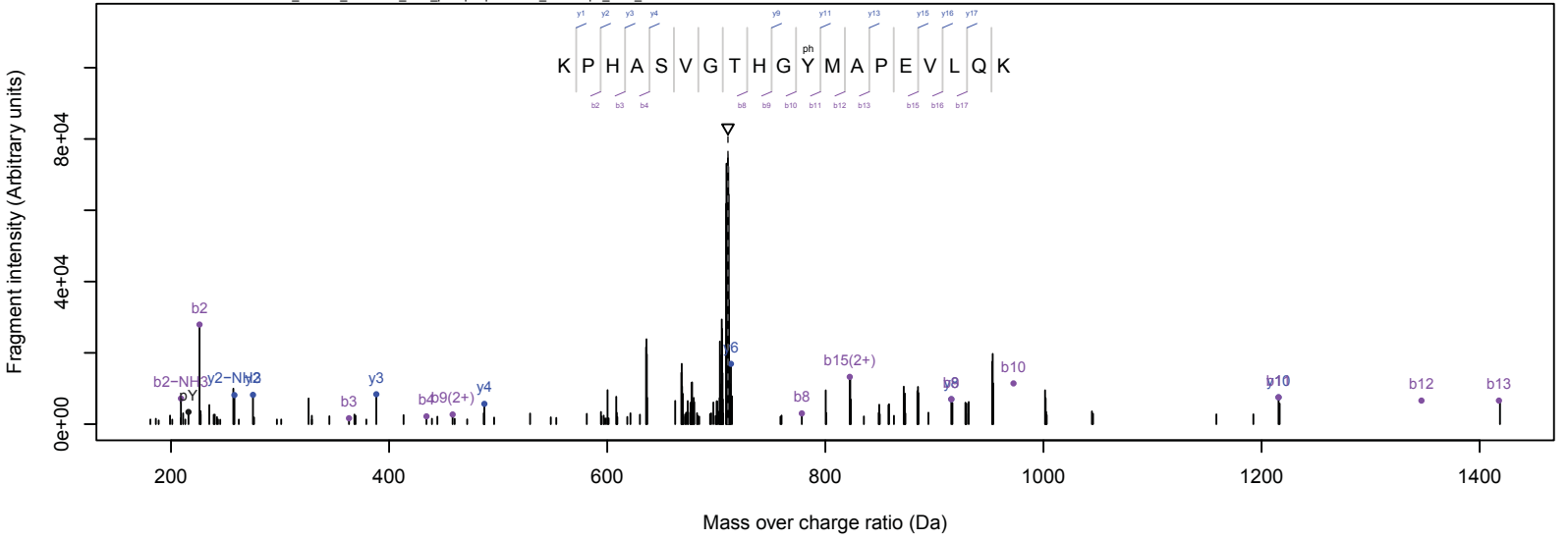
Andromeda score : 78.752



**KPHASVGT<sup>ph</sup>HGYMAPEVLQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15169

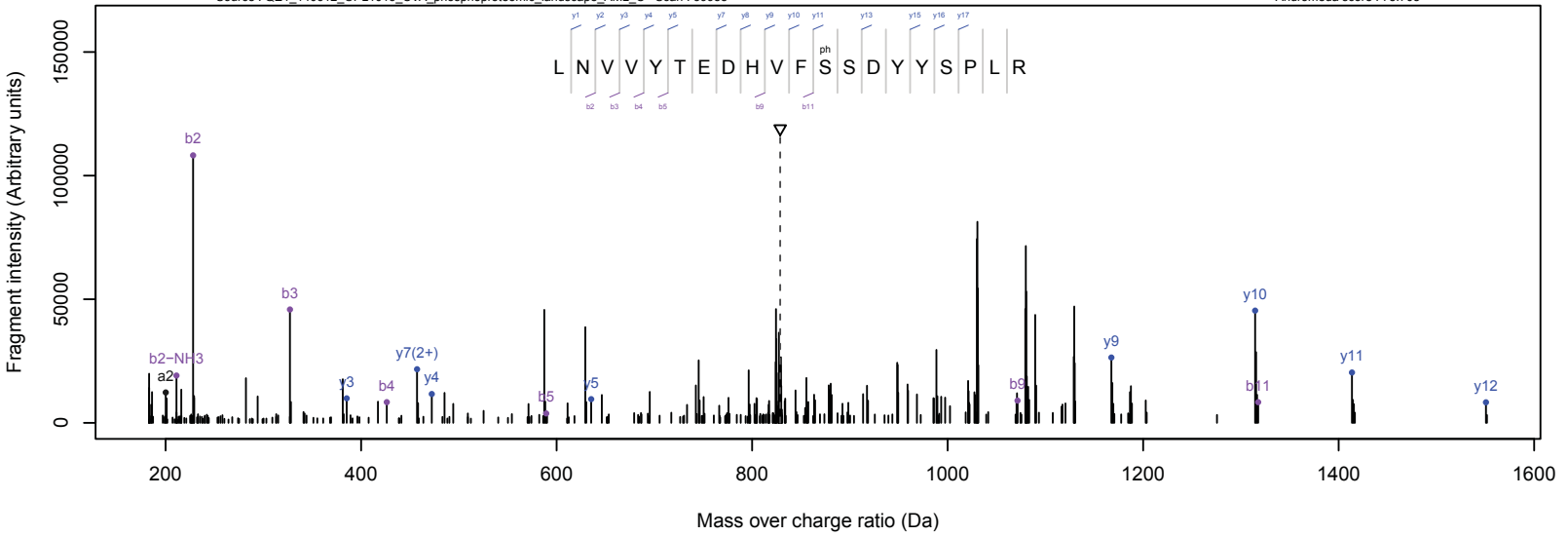
Andromeda score : 78.708



**LNVVYTEDHVFS<sup>ph</sup>SDYYSPLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39988

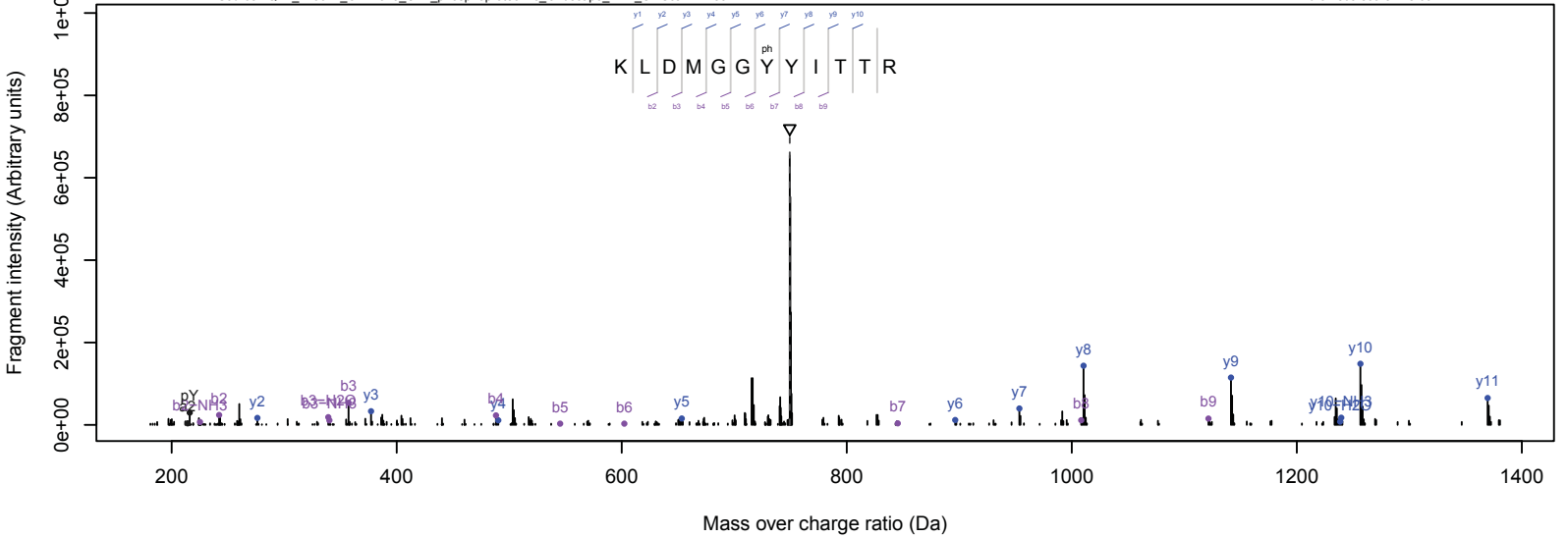
Andromeda score : 78.708



**KLDMGGY<sup>ph</sup>YITTR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22852

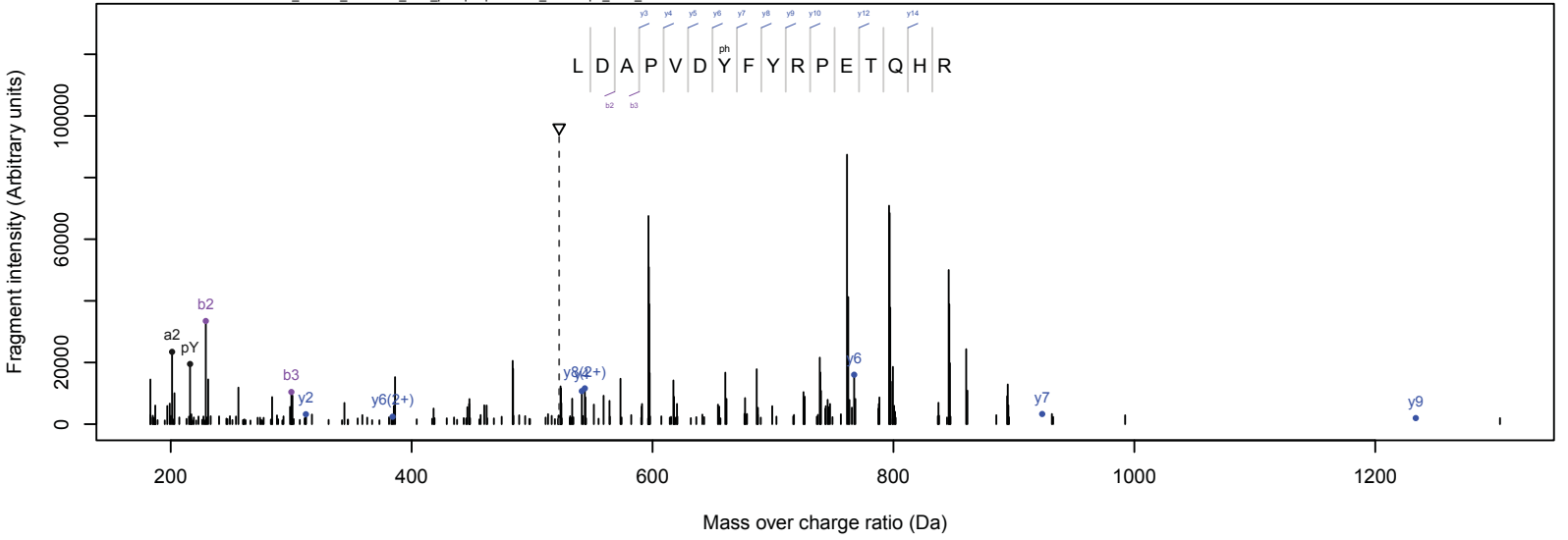
Andromeda score : 78.69



**LDAPVDY(ph)FYRPETQHR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29829

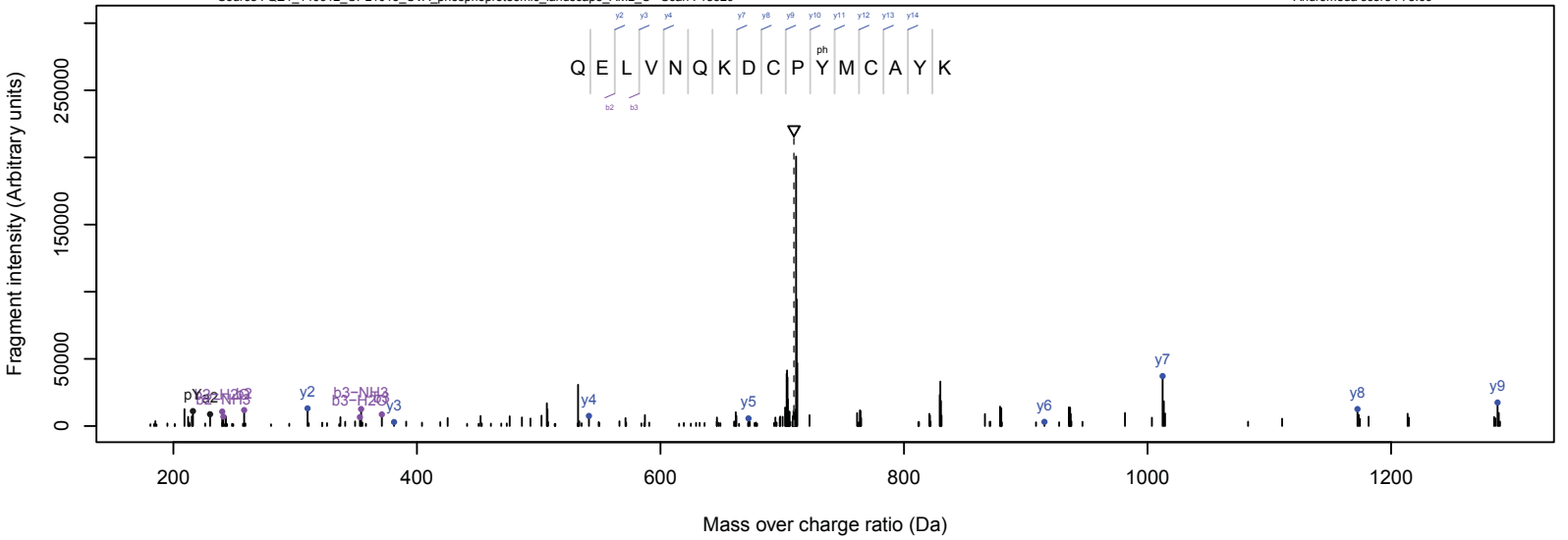
Andromeda score : 78.69



**QELVNQKDCPY(ph)MCAYK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18329

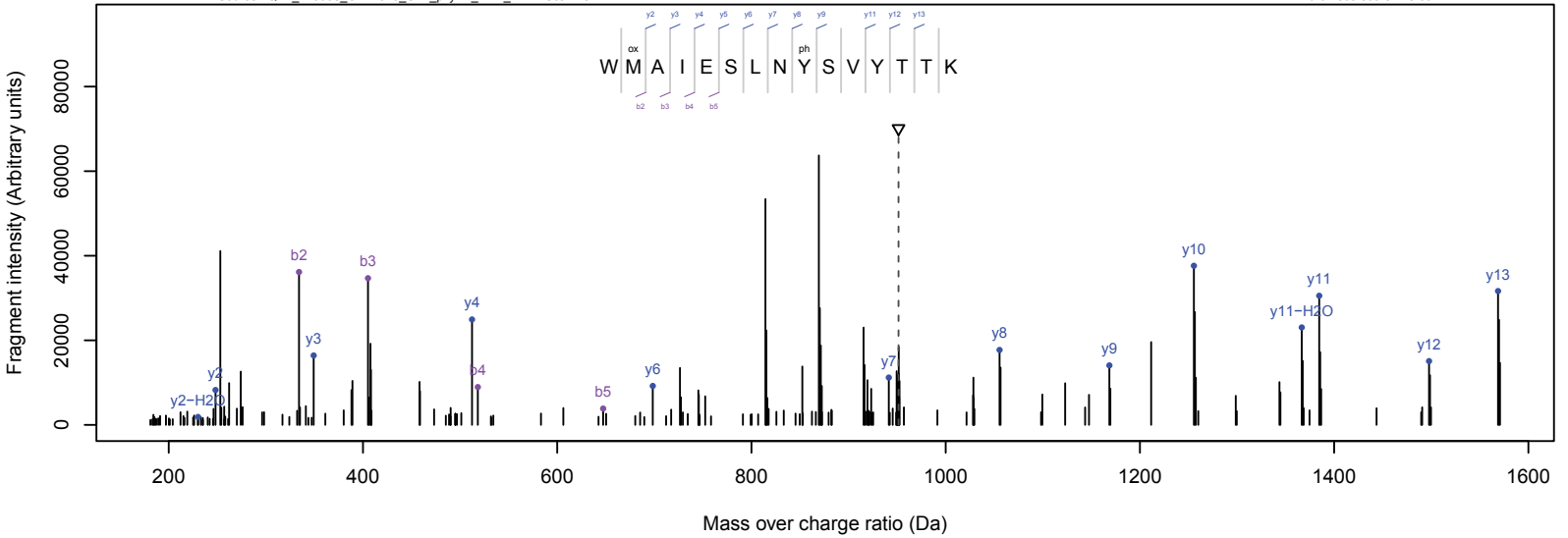
Andromeda score : 78.69



**WM(ox)AIESLNY(ph)SVYTK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37717

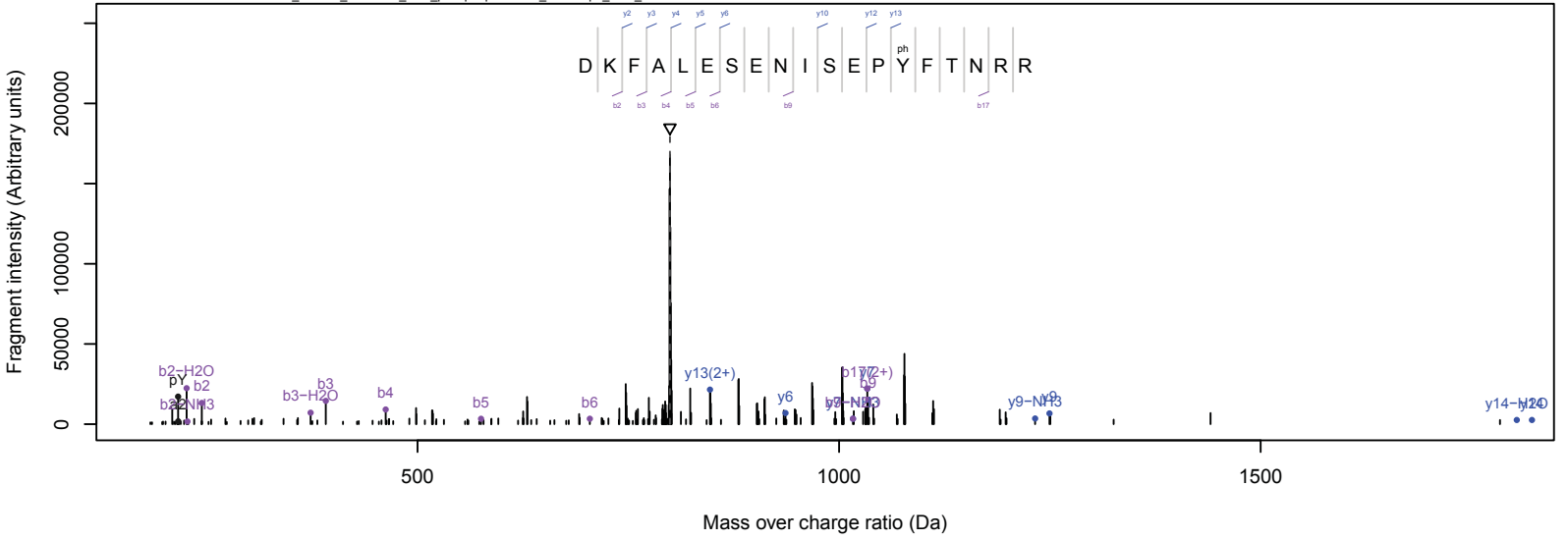
Andromeda score : 78.69



**DKFALESENISEPY(ph)FTNRR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29345

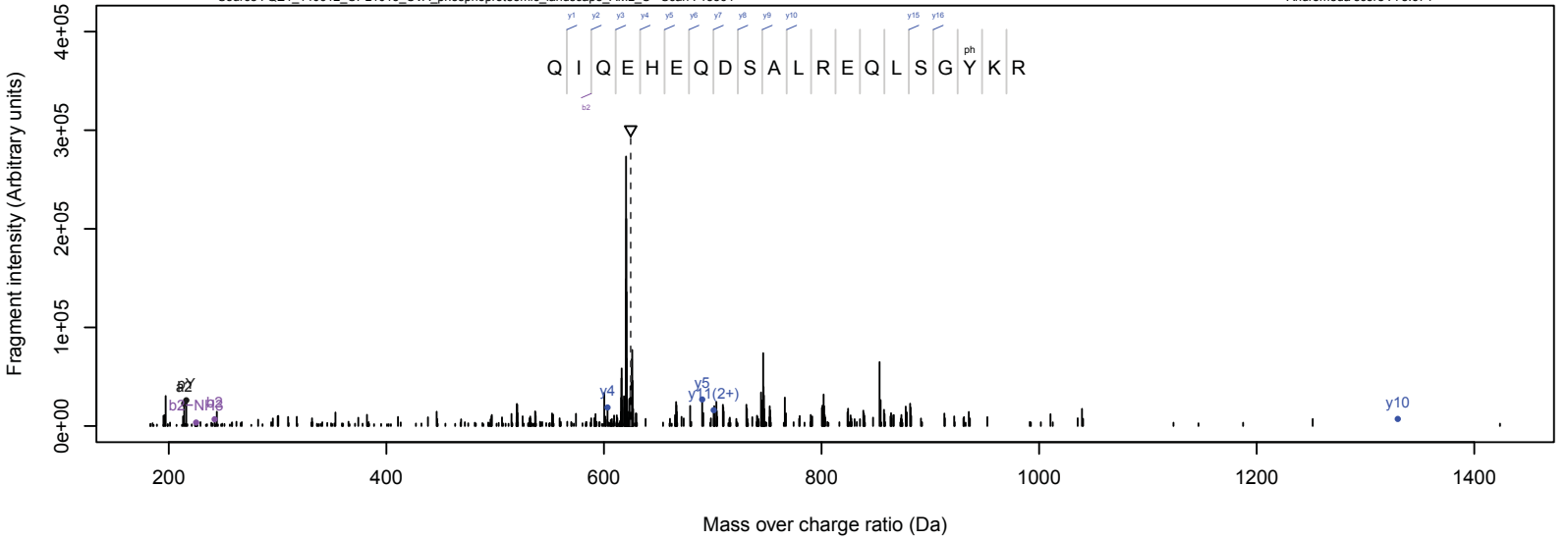
Andromeda score : 78.674



**QIQEHEQDSALREQLSGYK R-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13564

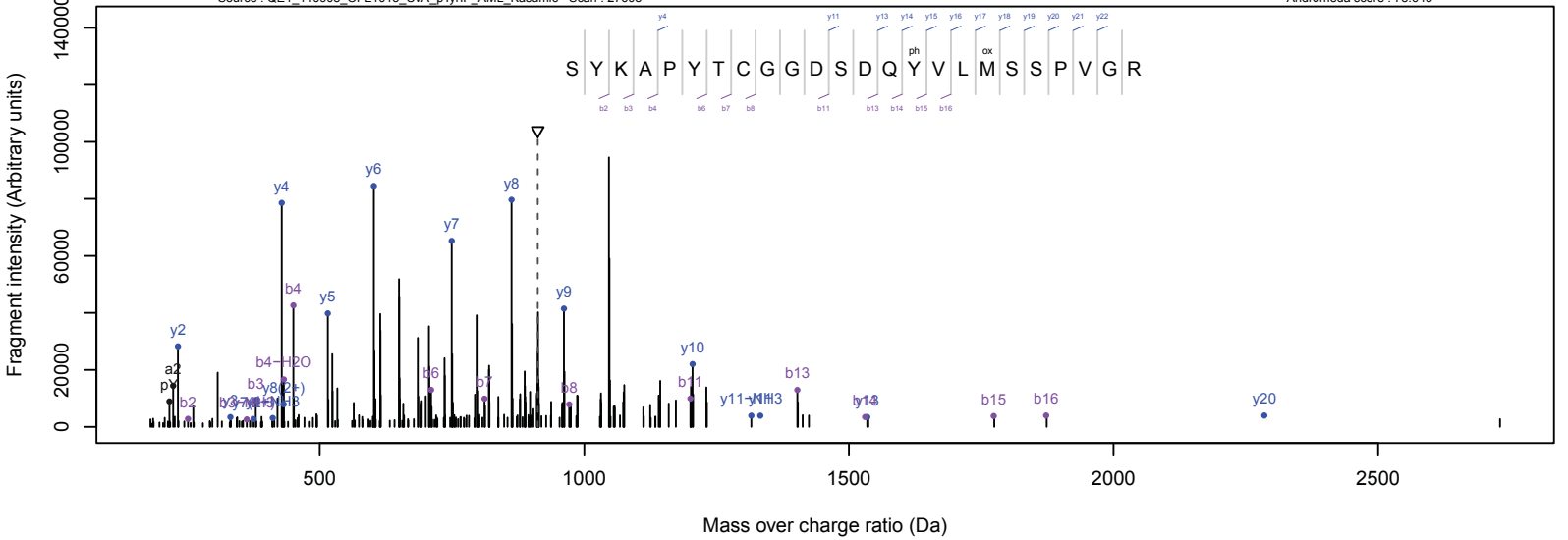
Andromeda score : 78.674



**SYKAPYTCGGDSQY(ph)VLM(ox)SSPVGR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 27605

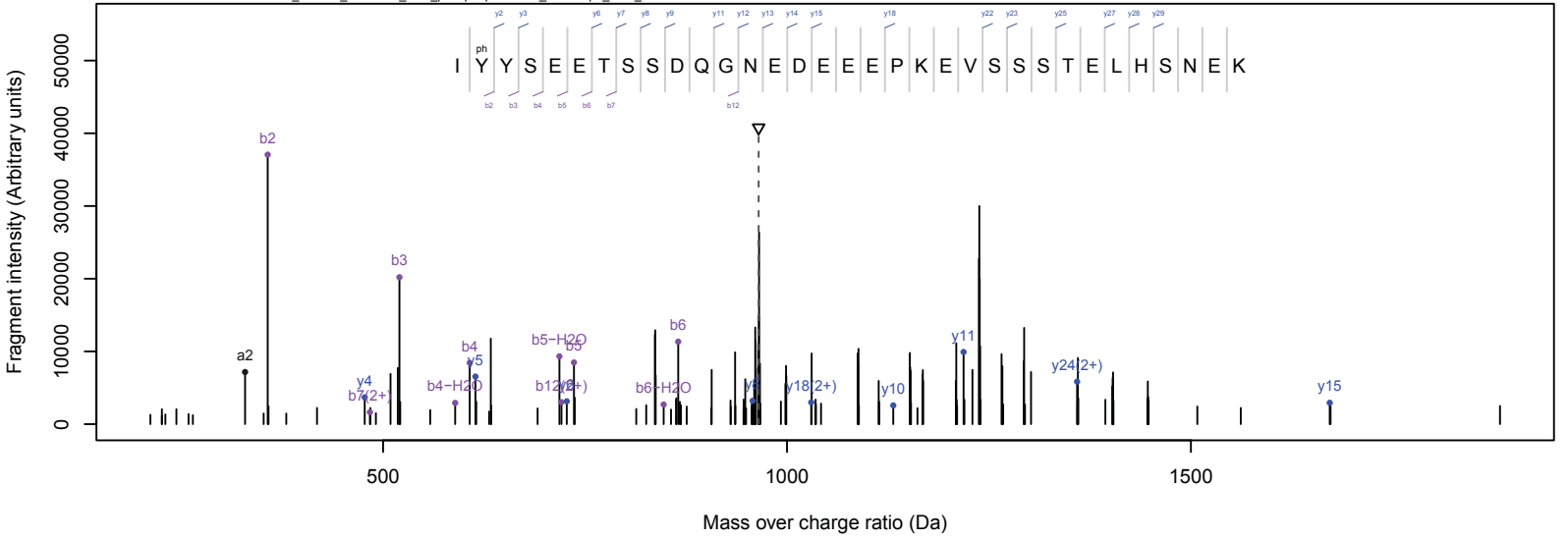
Andromeda score : 78.645



IY(ph)YSEETSSDQGNEDDEEPEKVVSSSTELHSNEK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 17709

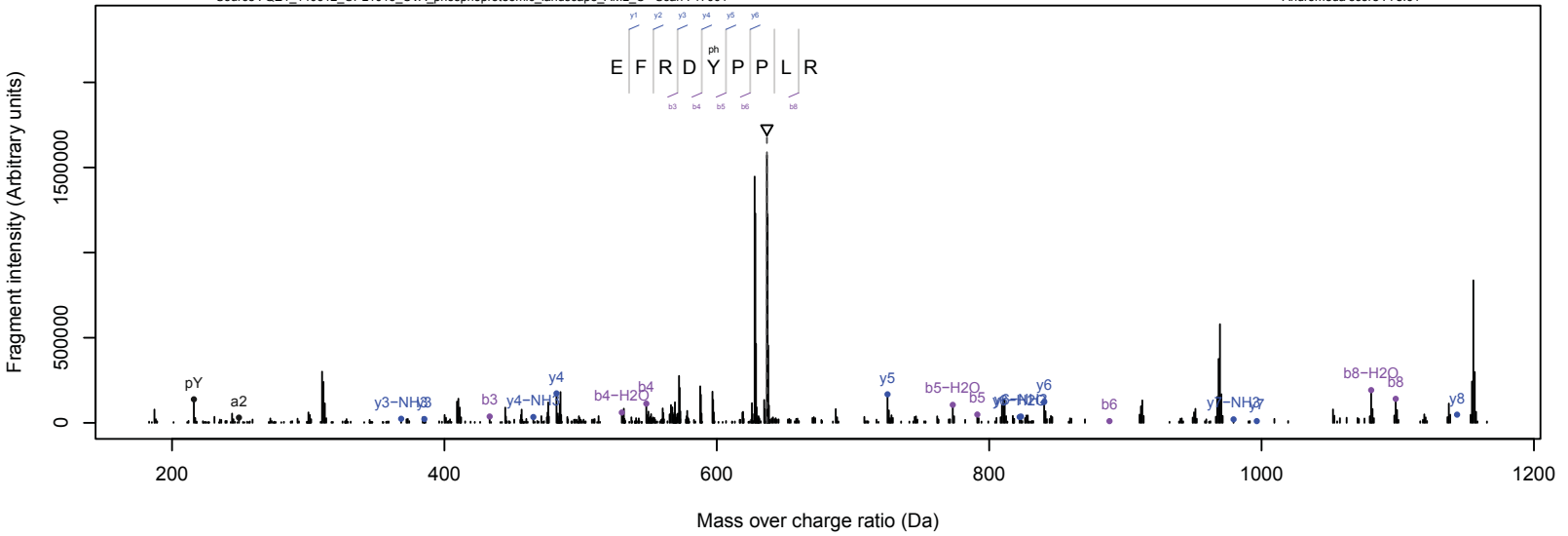
Andromeda score : 78.642



EFRDY(ph)PPLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17001

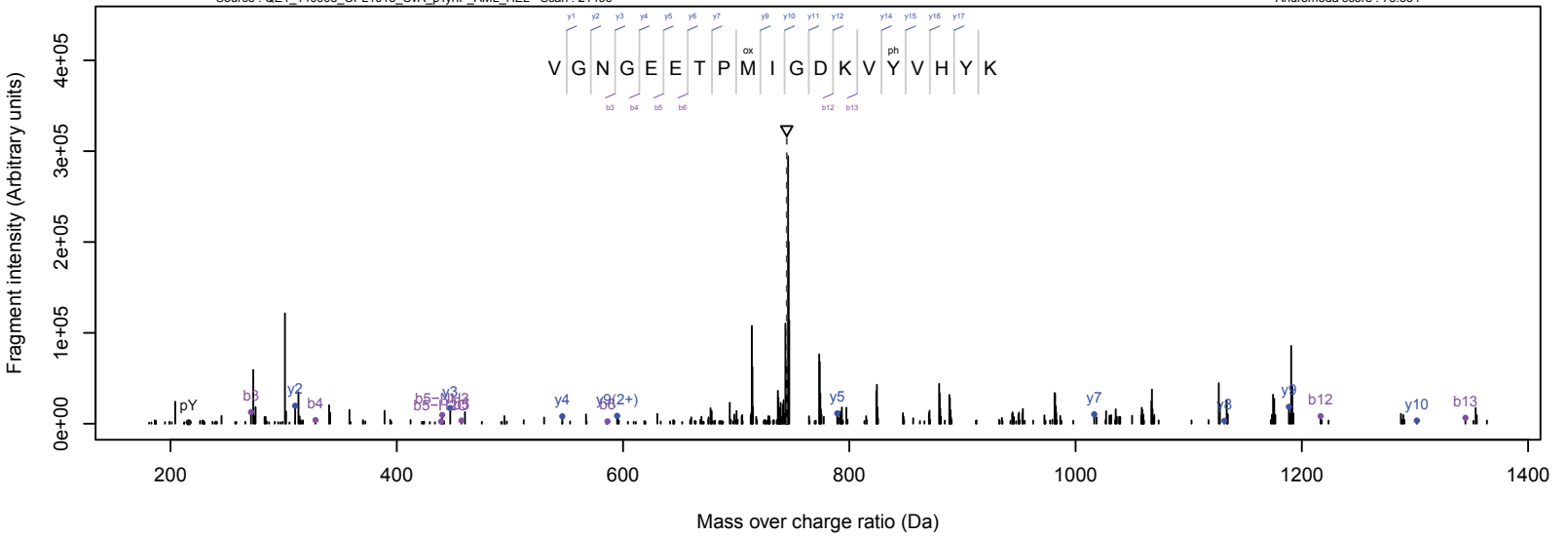
Andromeda score : 78.61



VGNGETPM(ox)IGDKVYVHYK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 21496

Andromeda score : 78.554

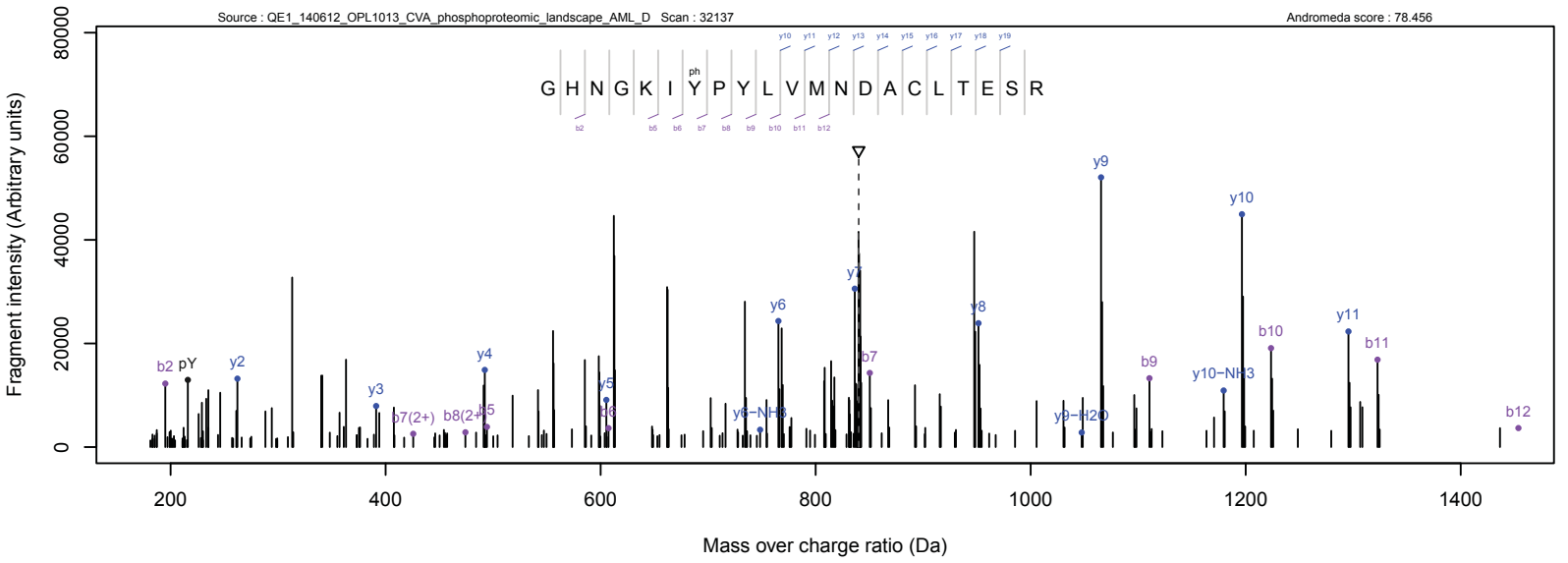




**GHNGKIY(ph)PYLVMNDACLTER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 32137

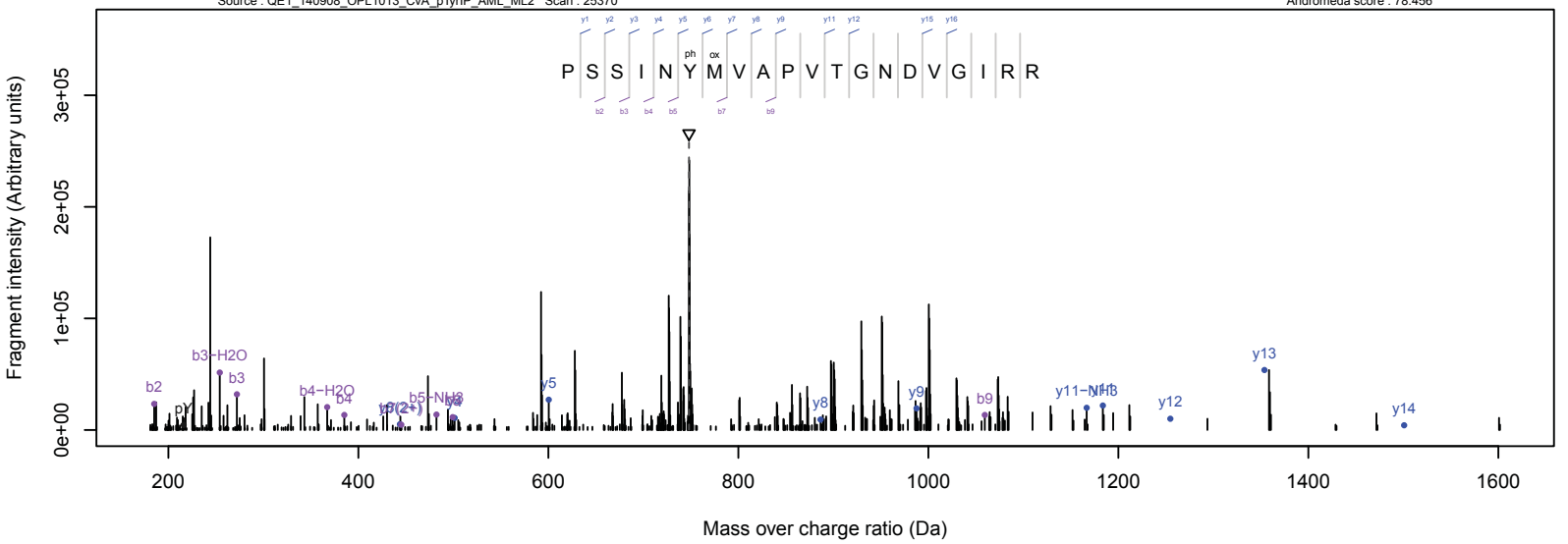
Andromeda score : 78.456



**PSSINY(ph)M(ox)VAPVTGNDVGIRR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 25370

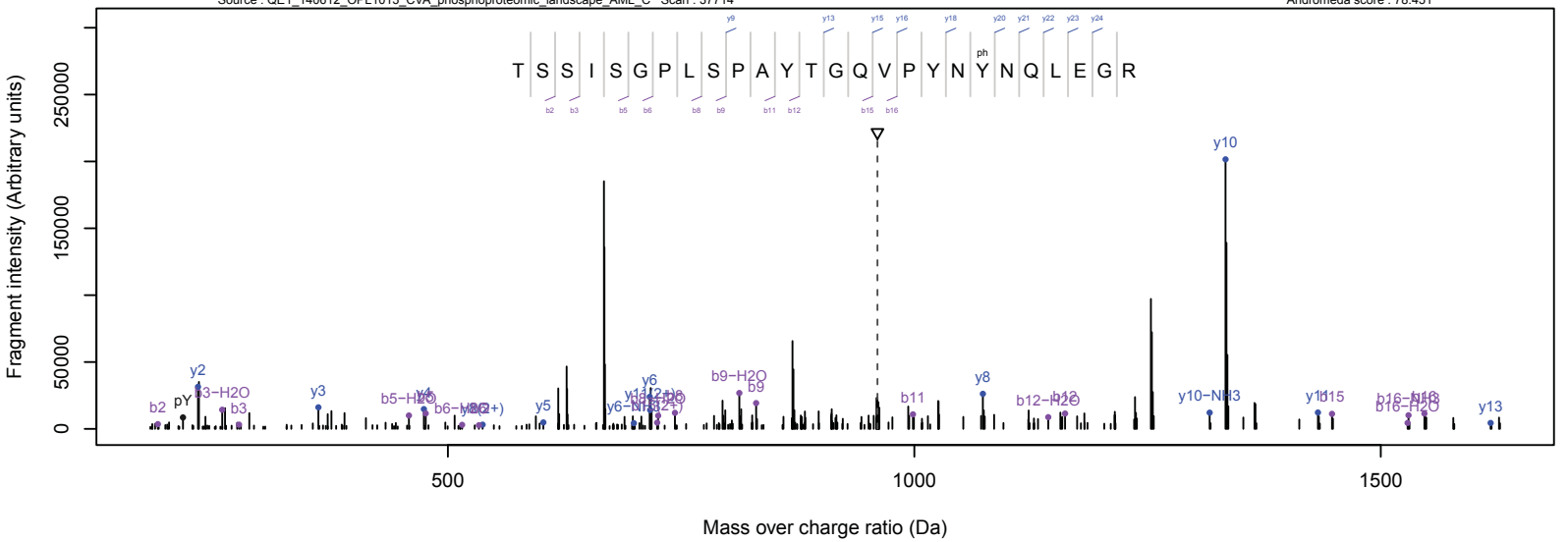
Andromeda score : 78.456



**TSSISGPLSPAYTGQVPYNYNQLLEGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37714

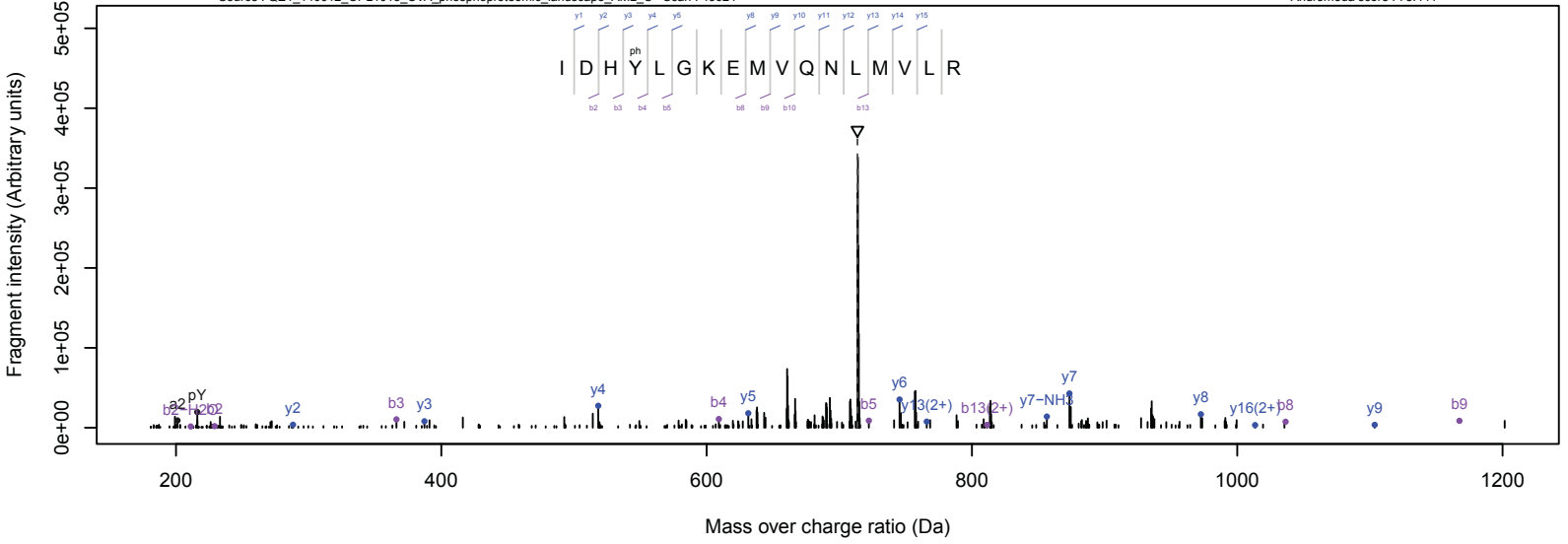
Andromeda score : 78.451



IDHY(ph)LGKEMVQLMVLRL-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43624

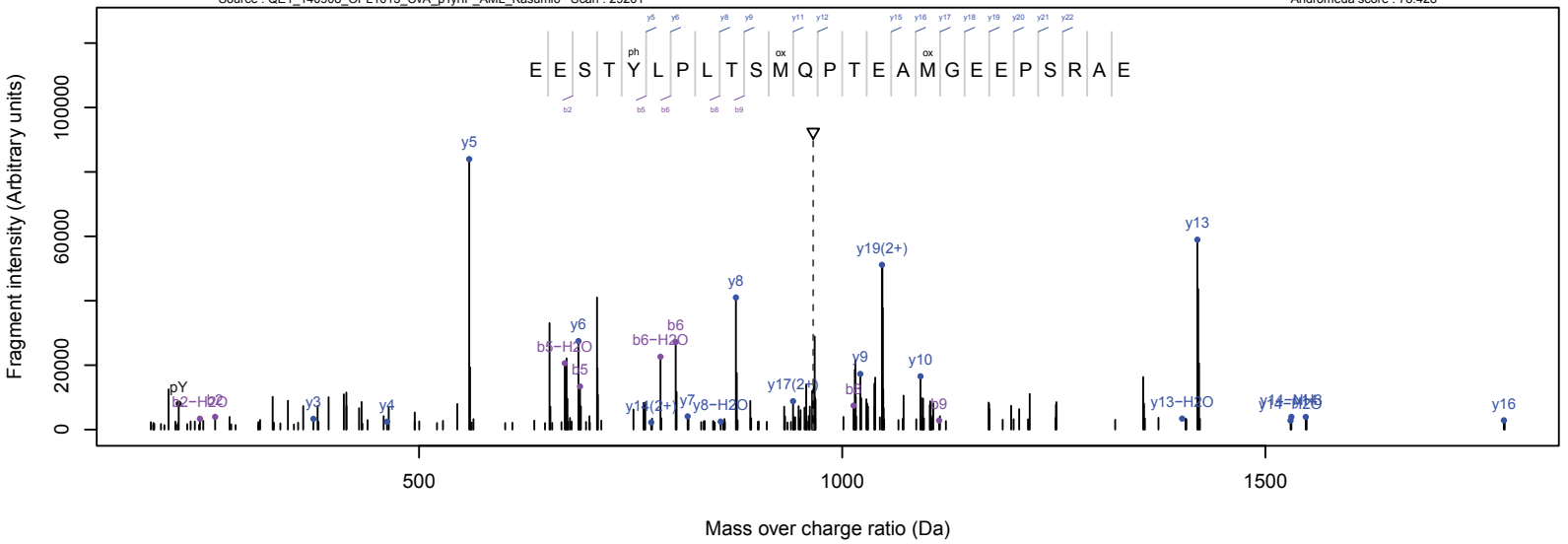
Andromeda score : 78.441



EESTY(ph)LPLTSM(ox)QPTEAM(ox)GEEPSRAE-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 29201

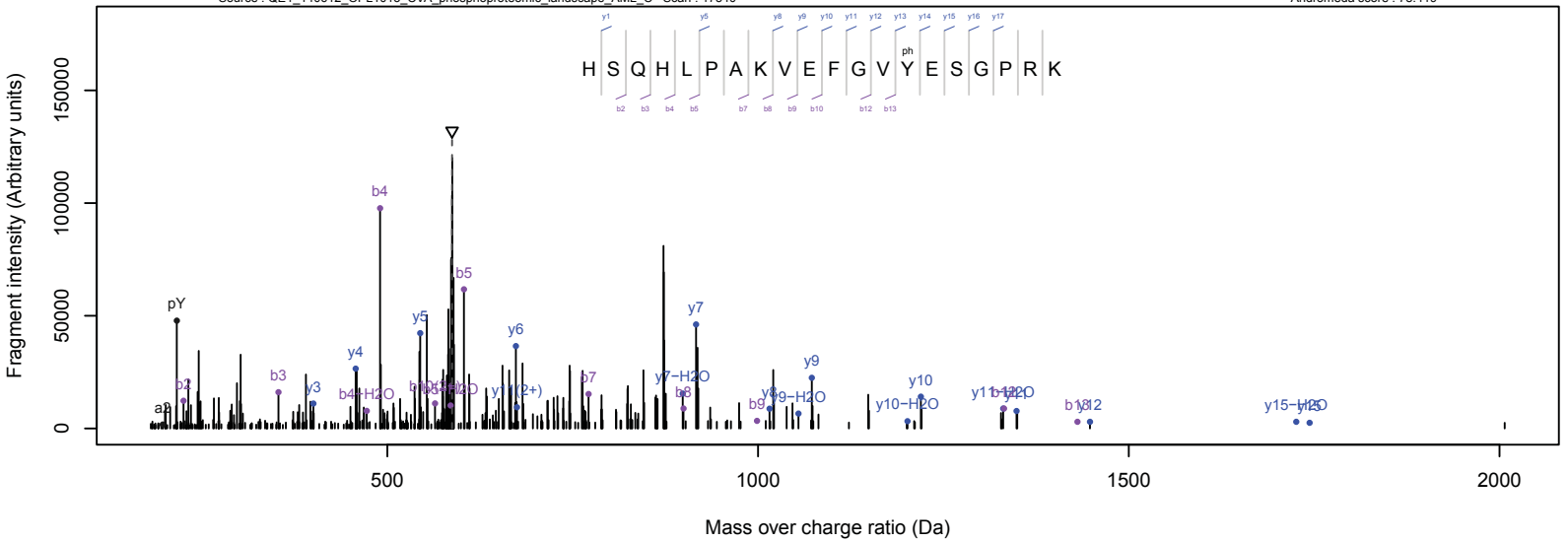
Andromeda score : 78.428



HSQHLPAKVEFGVY(ESGPRK)-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17540

Andromeda score : 78.419

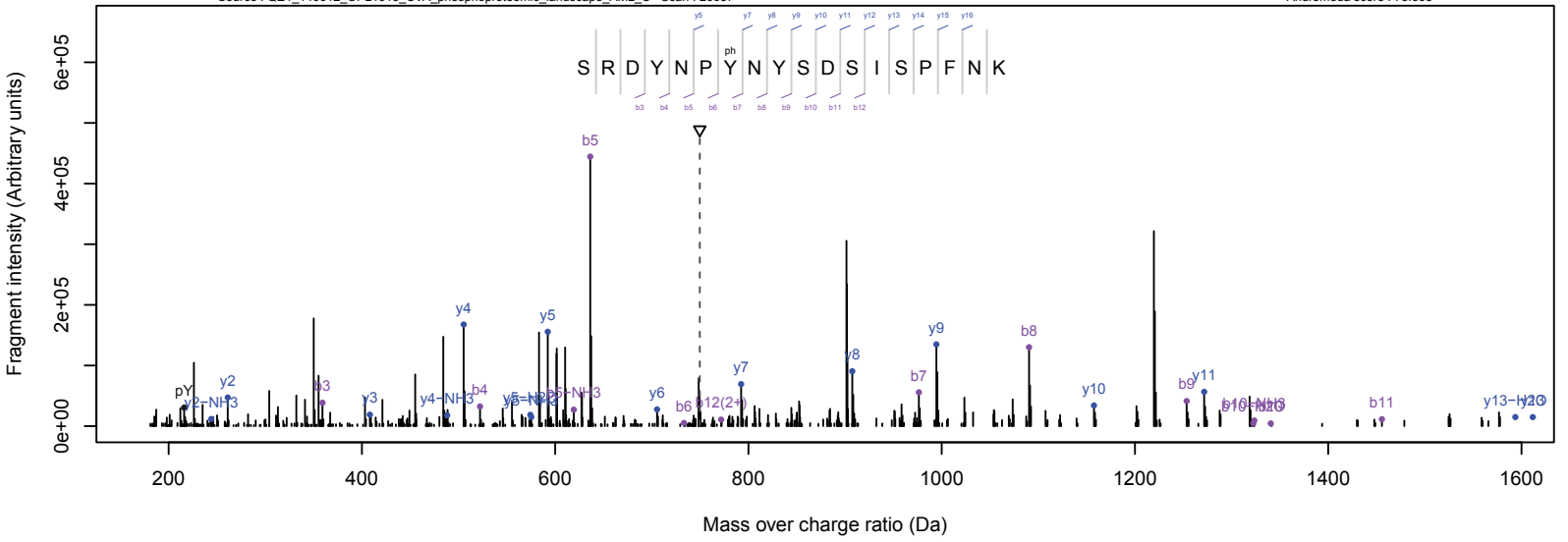




SRDYNP(ph)NYSDSISPFNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26687

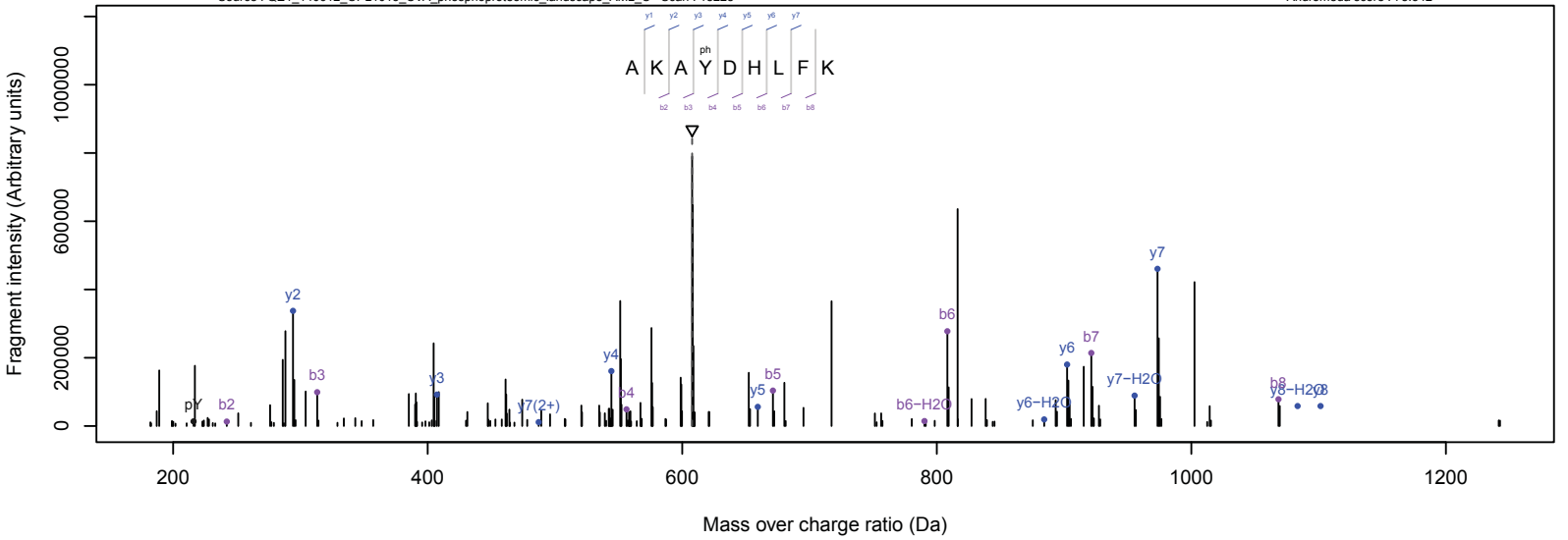
Andromeda score : 78.358



(ac)AKAY(ph)DHLFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18228

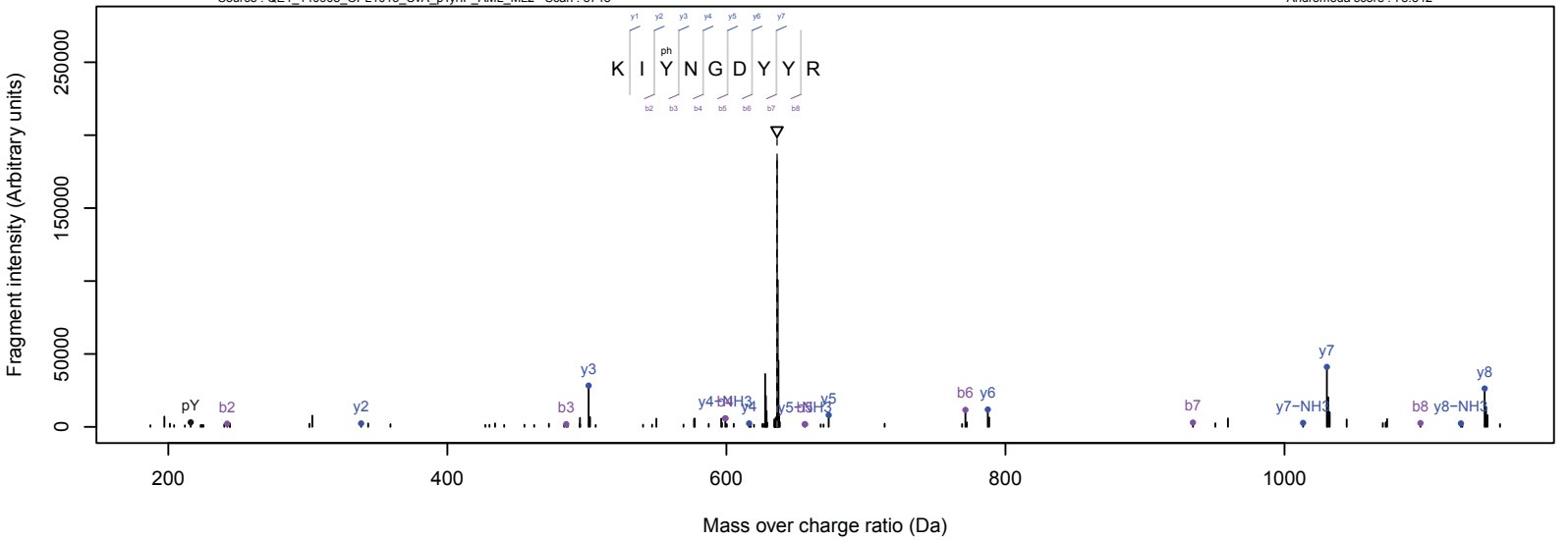
Andromeda score : 78.342



KIY(ph)NGDYR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 8748

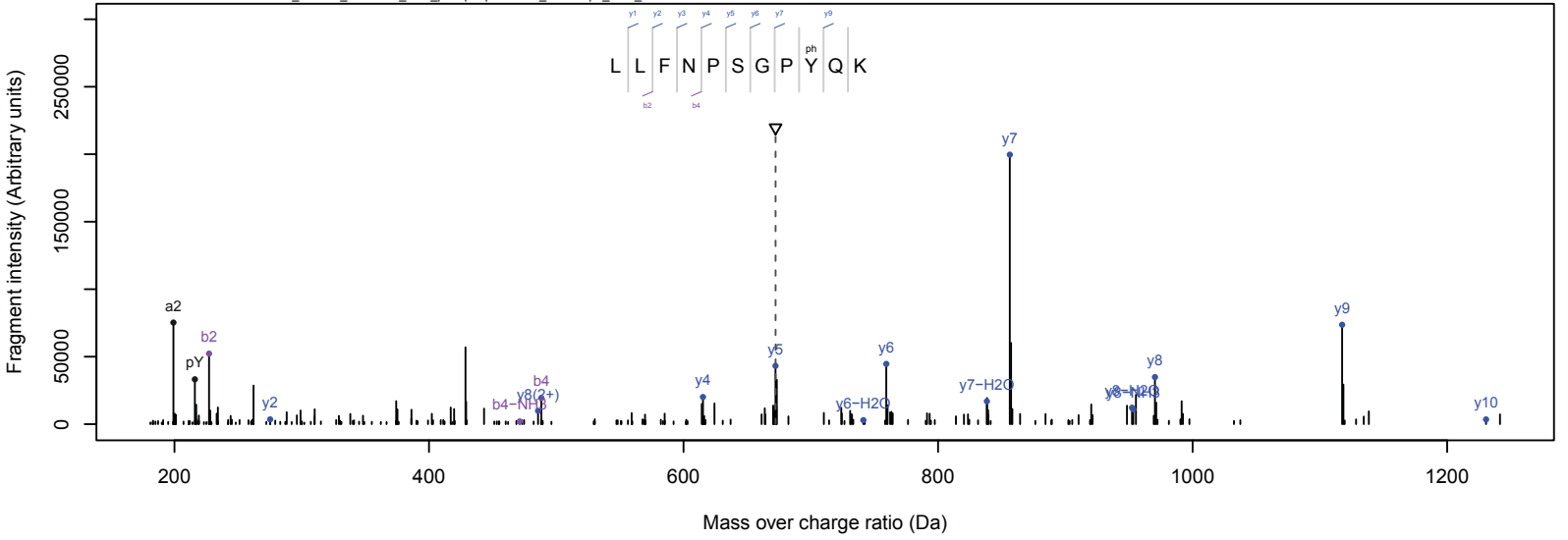
Andromeda score : 78.342



LLFNPSGPYQK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 27132

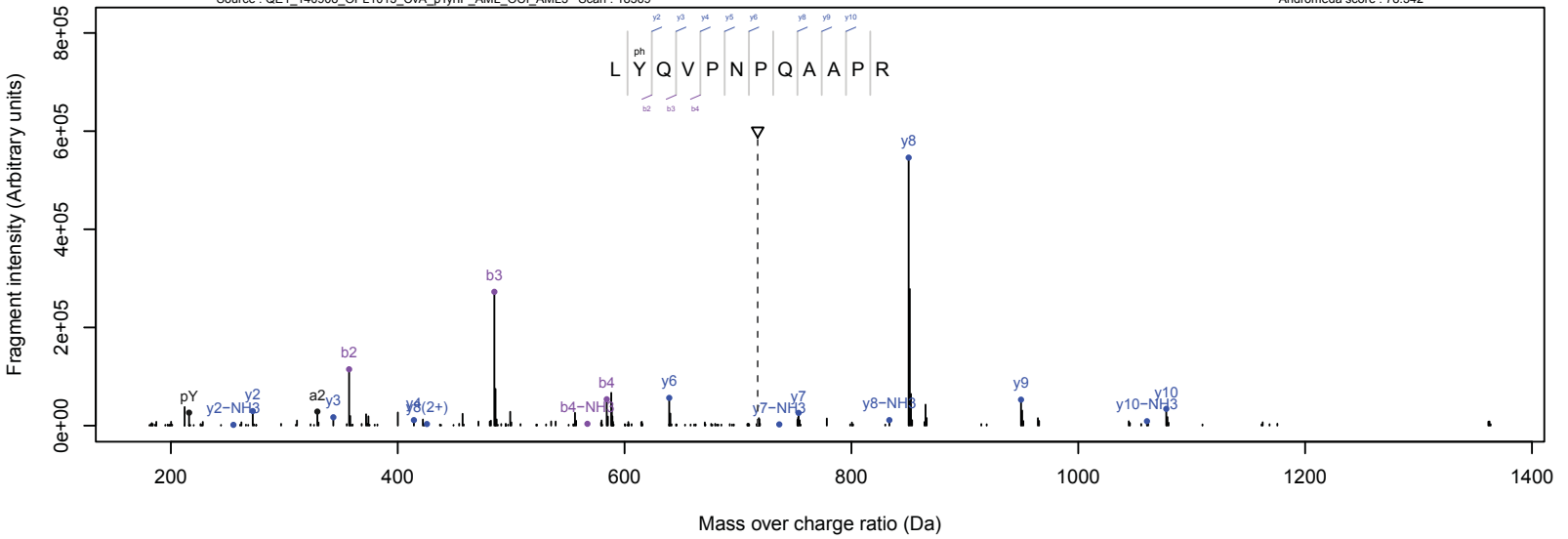
Andromeda score: 78.342



LY(ph)QVPNPQAAPR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 18569

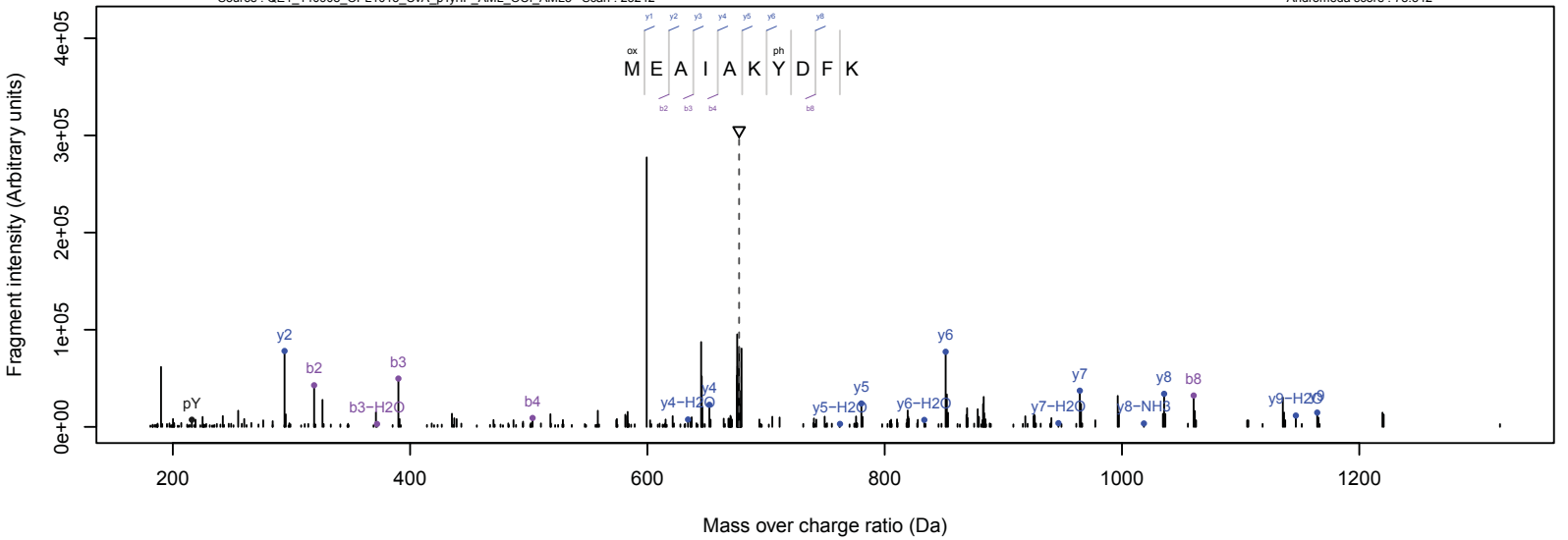
Andromeda score: 78.342



(ac)M(ox)EAIKY(ph)DFK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 23242

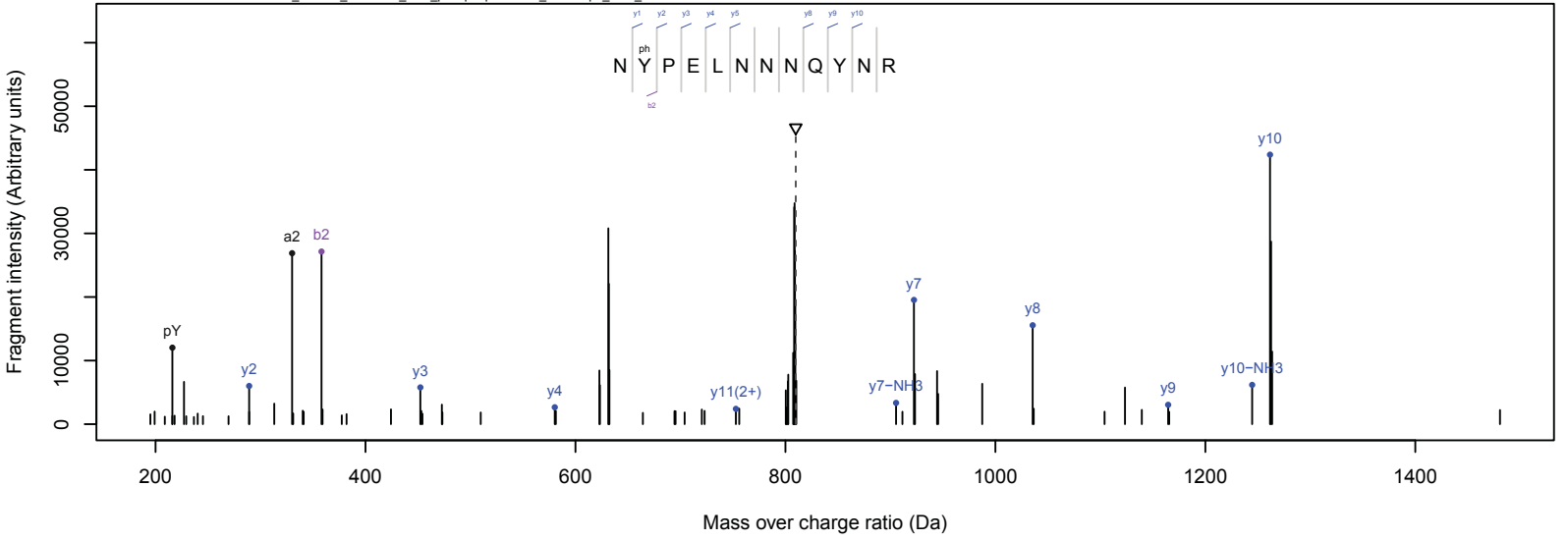
Andromeda score: 78.342



NY(ph)PELNNQYNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11567

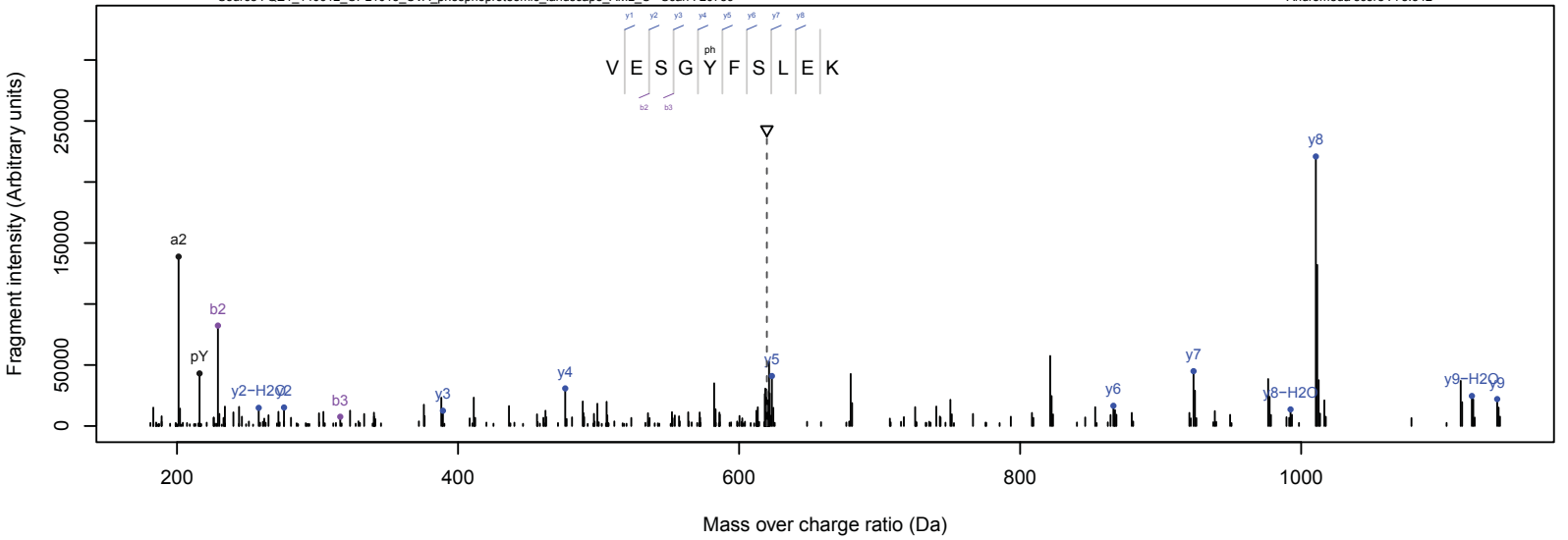
Andromeda score : 78.342



VESGY(ph)FSLEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20789

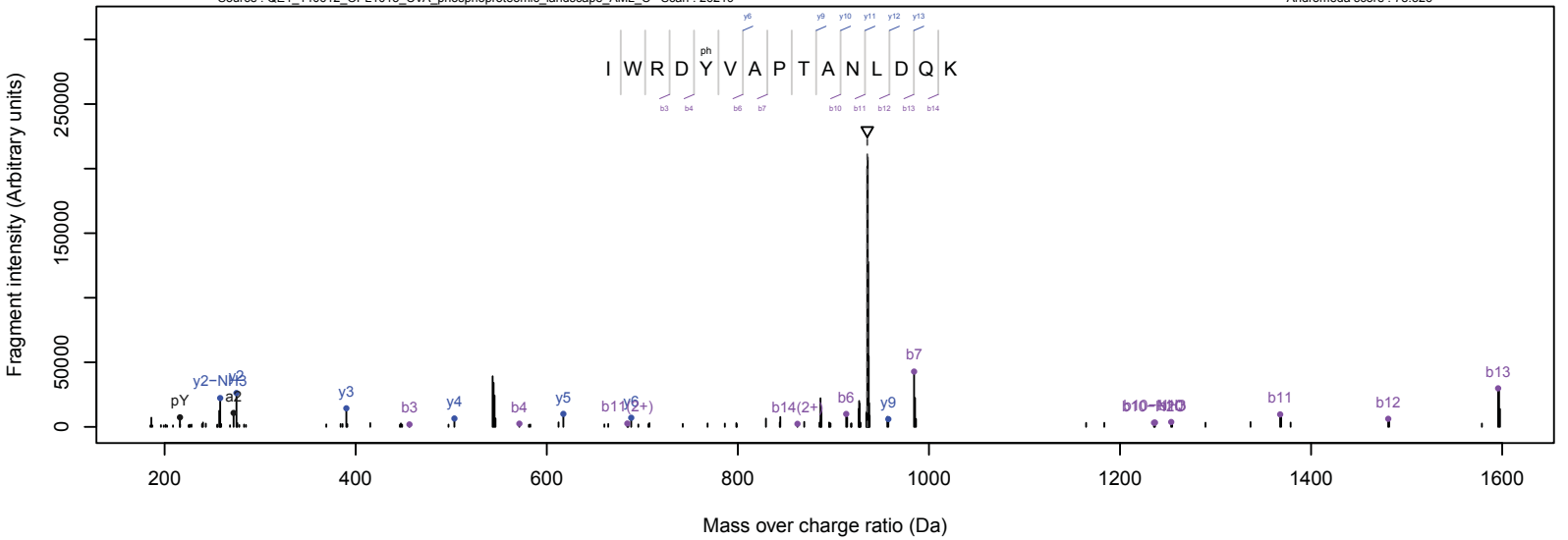
Andromeda score : 78.342



IWRDY(ph)VAPTANLDQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26219

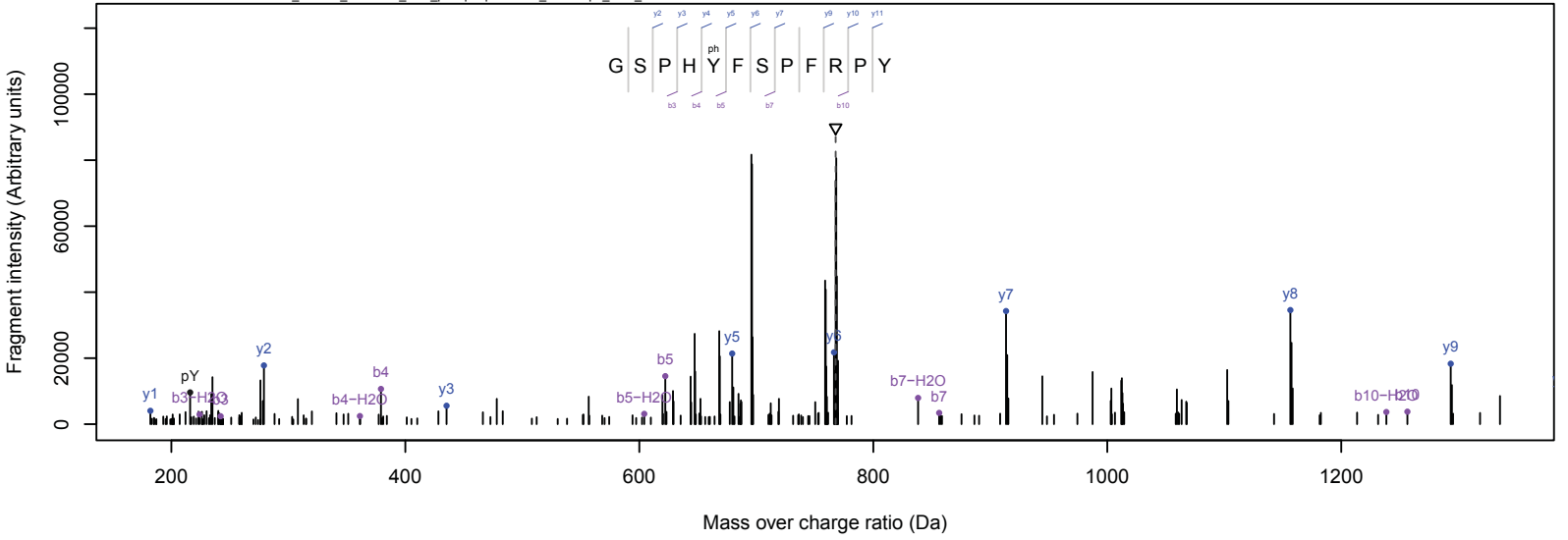
Andromeda score : 78.326



**GSPHY(ph)FSPFRPY-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30515

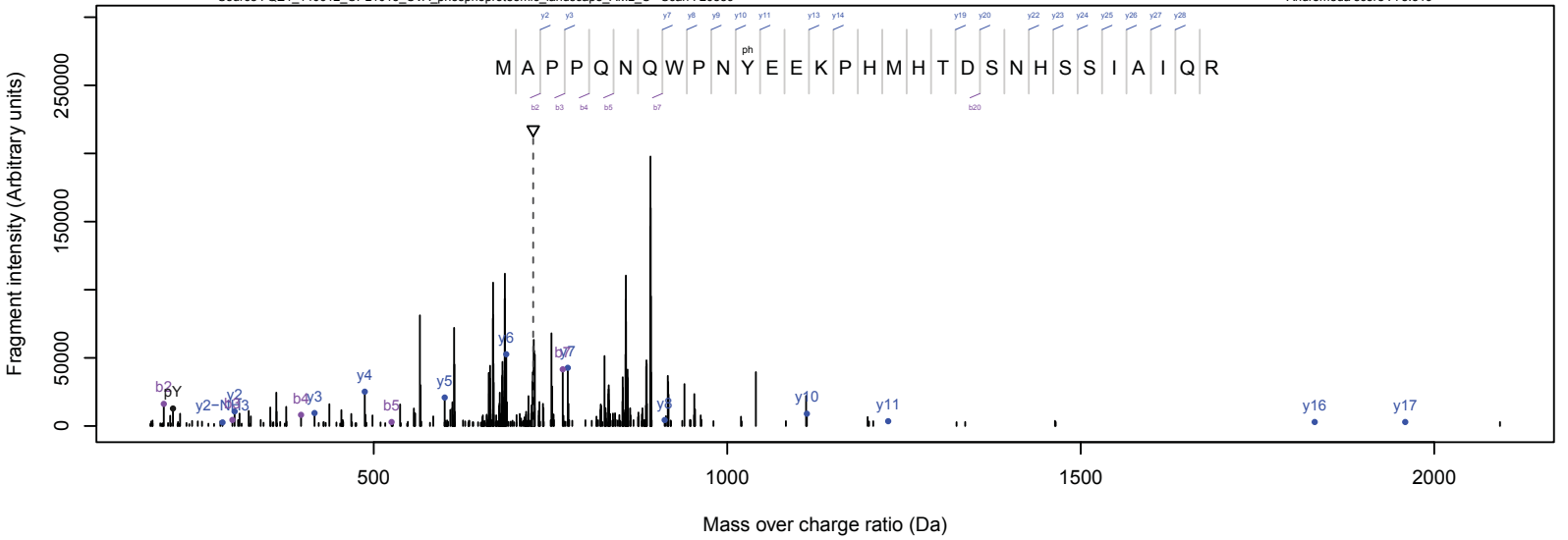
Andromeda score : 78.324



**MAPPQNQWPNY(ph)EEKPHMHTDSNHSSIAIQR-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20889

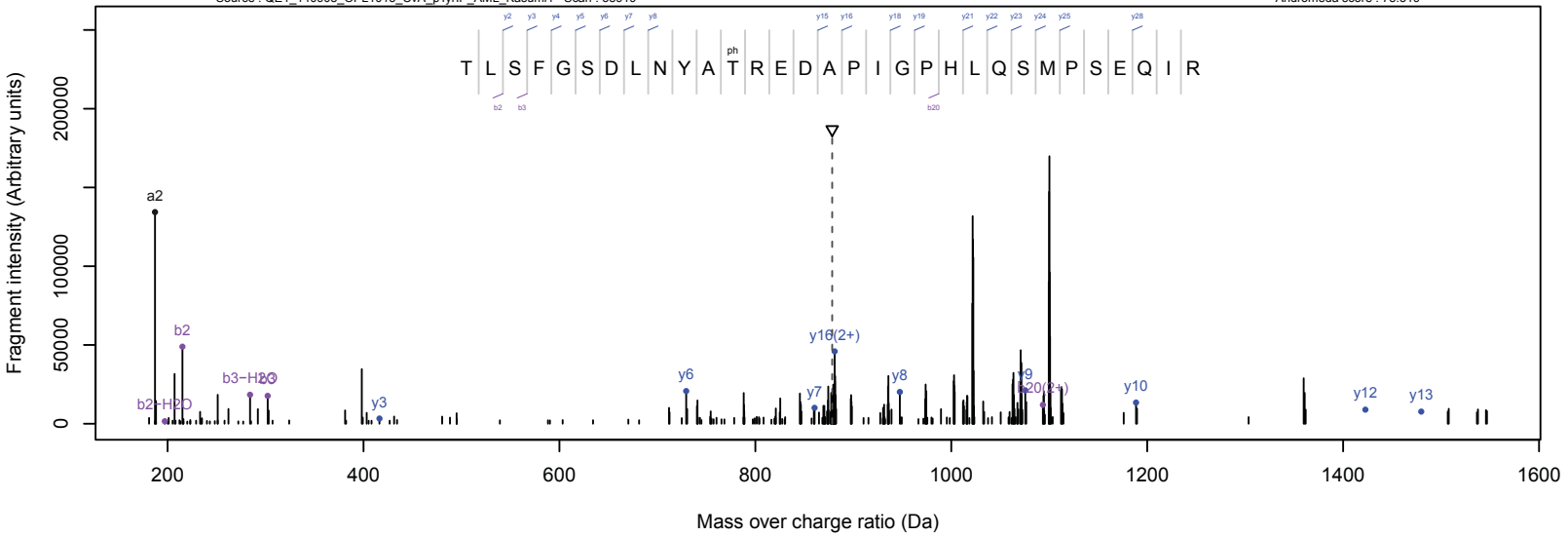
Andromeda score : 78.319



**TLSFGSDLNYAT(ph)REDAPIGPHLQSMPEQIR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 38919

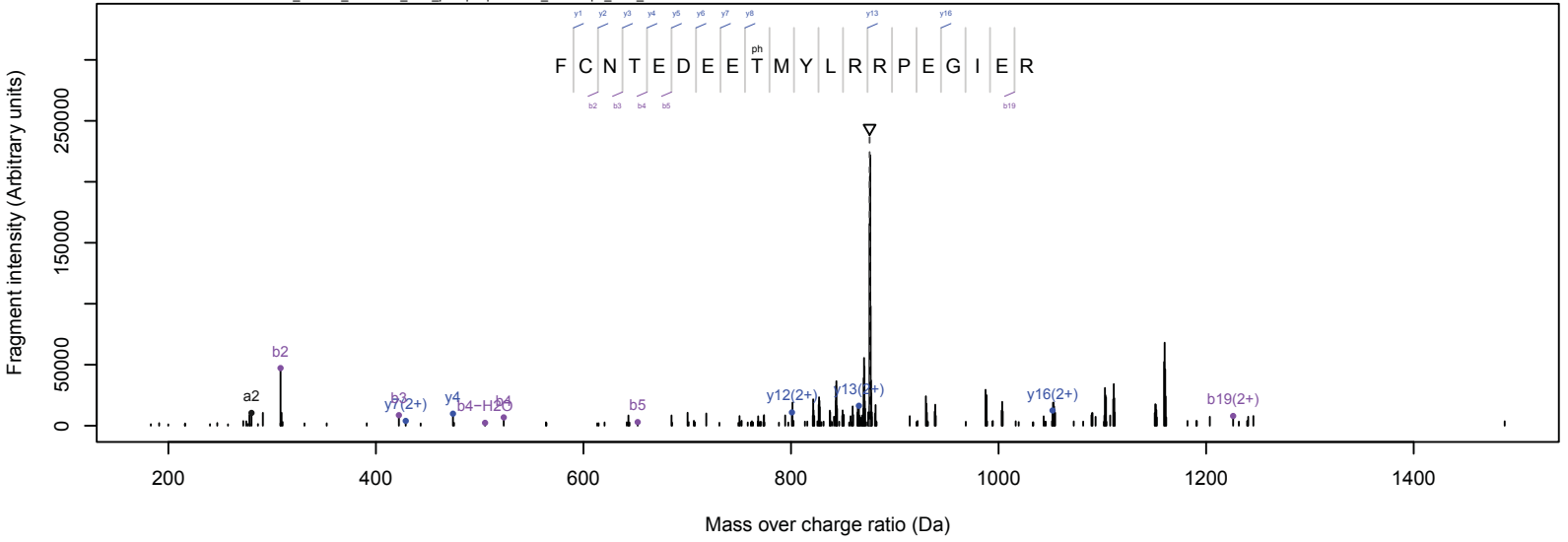
Andromeda score : 78.319



**FCNTEDEET(ph)MYLRRPEGIER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23172

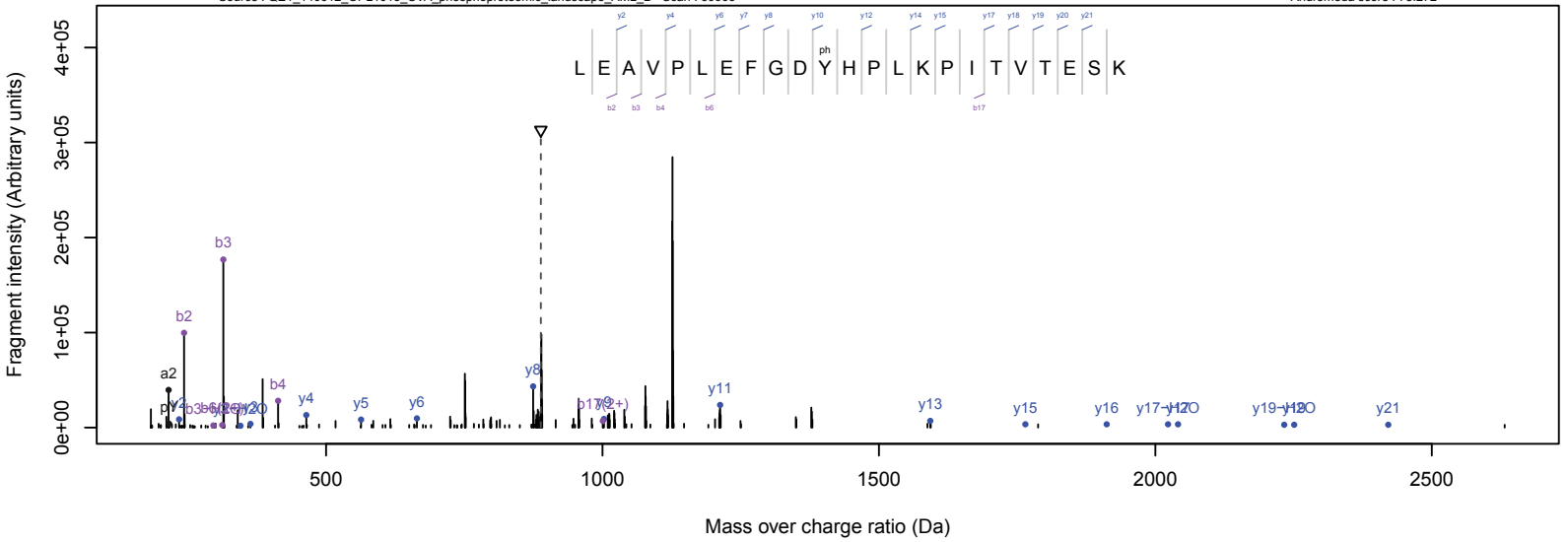
Andromeda score : 78.272



**LEAVPLEFGDY(ph)HPLKPIVTESK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39903

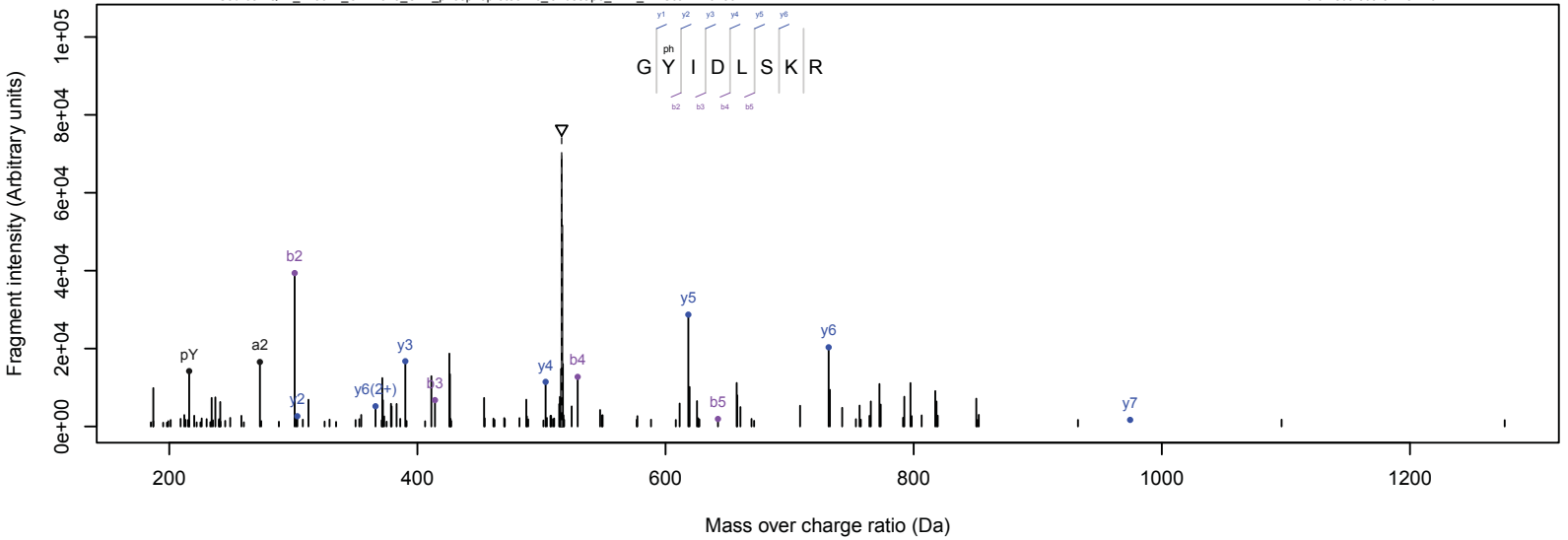
Andromeda score : 78.272



**GY(ph)IDLSKR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 10780

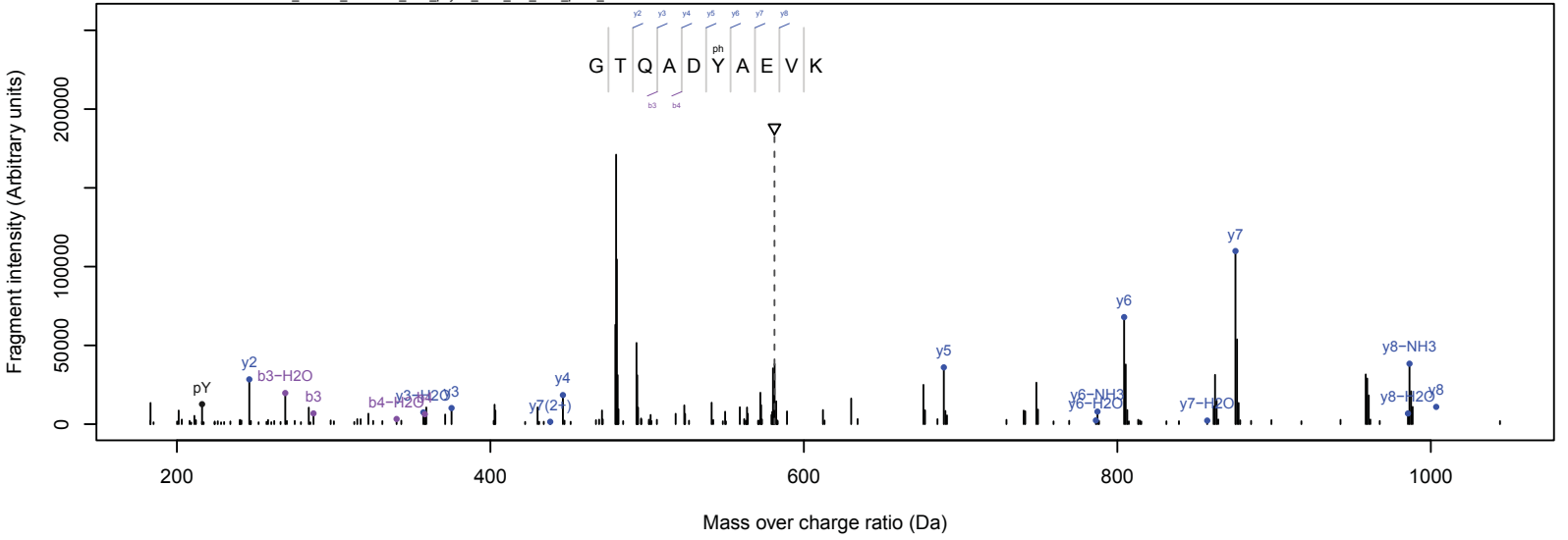
Andromeda score : 78.149



GTQADY(ph)AEVK-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3688

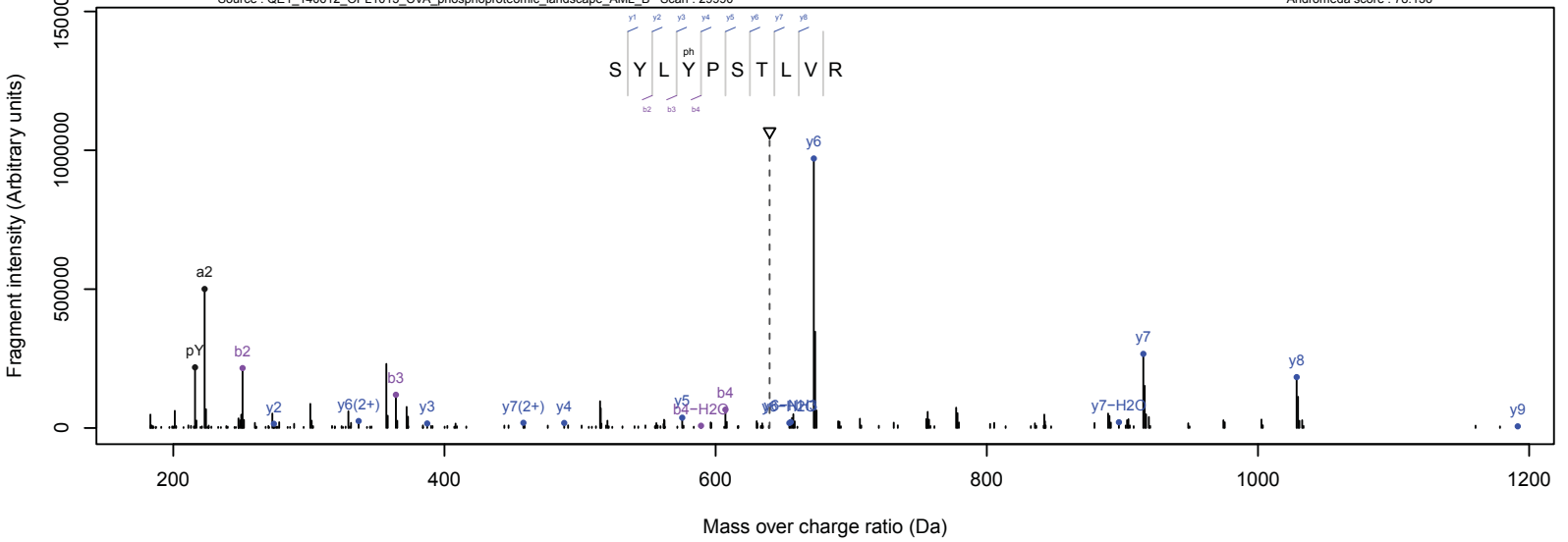
Andromeda score : 78.136



SYLY(ph)PSTLVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29990

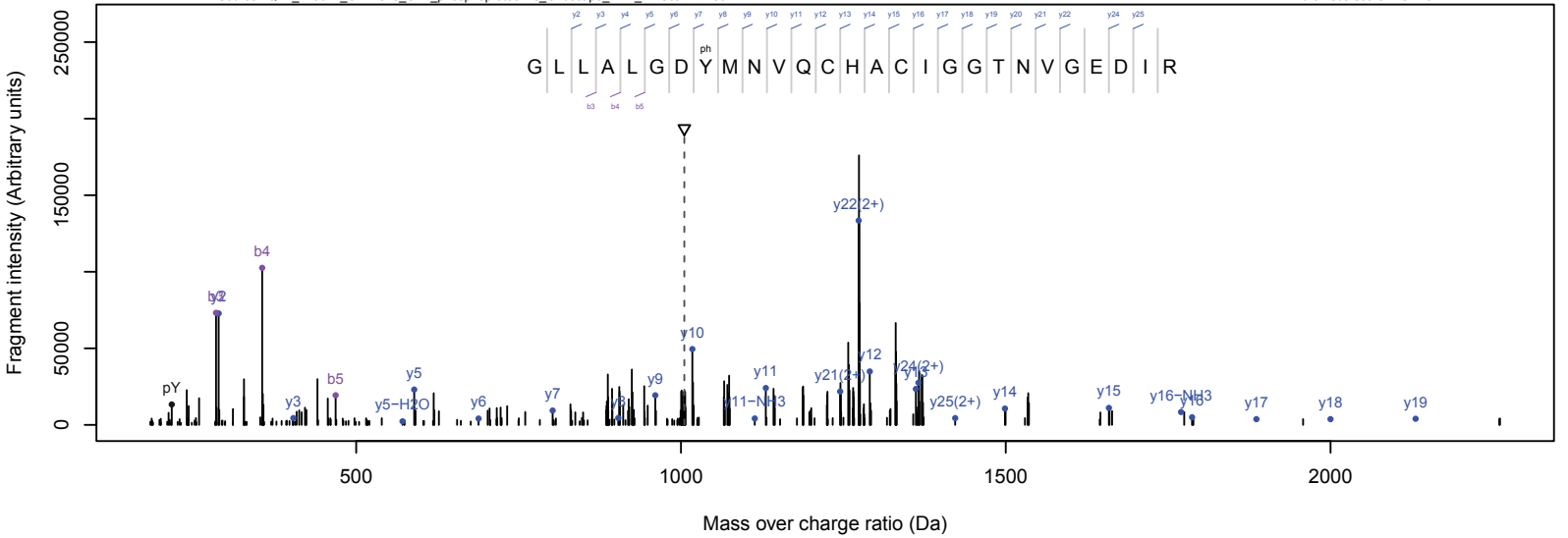
Andromeda score : 78.136



GLLALGDY(ph)MNVQCHACIGGTNVGEDIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42837

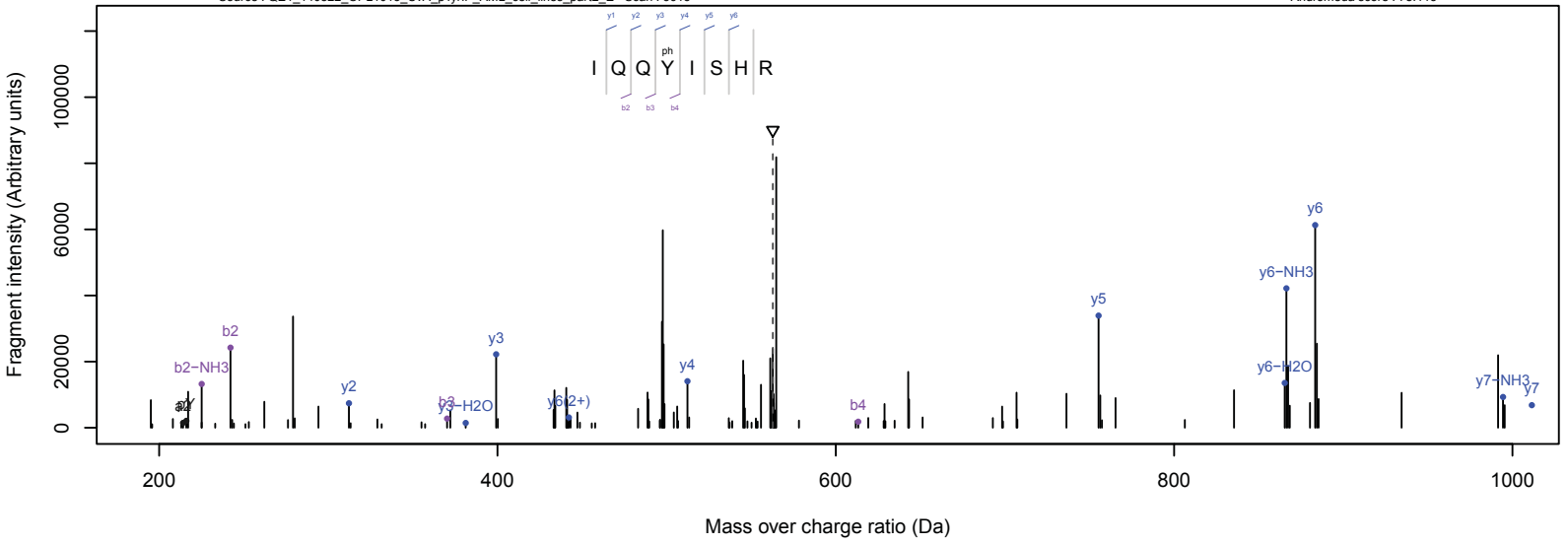
Andromeda score : 78.128



**IQQY(ph)ISHR-2+**

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 3615

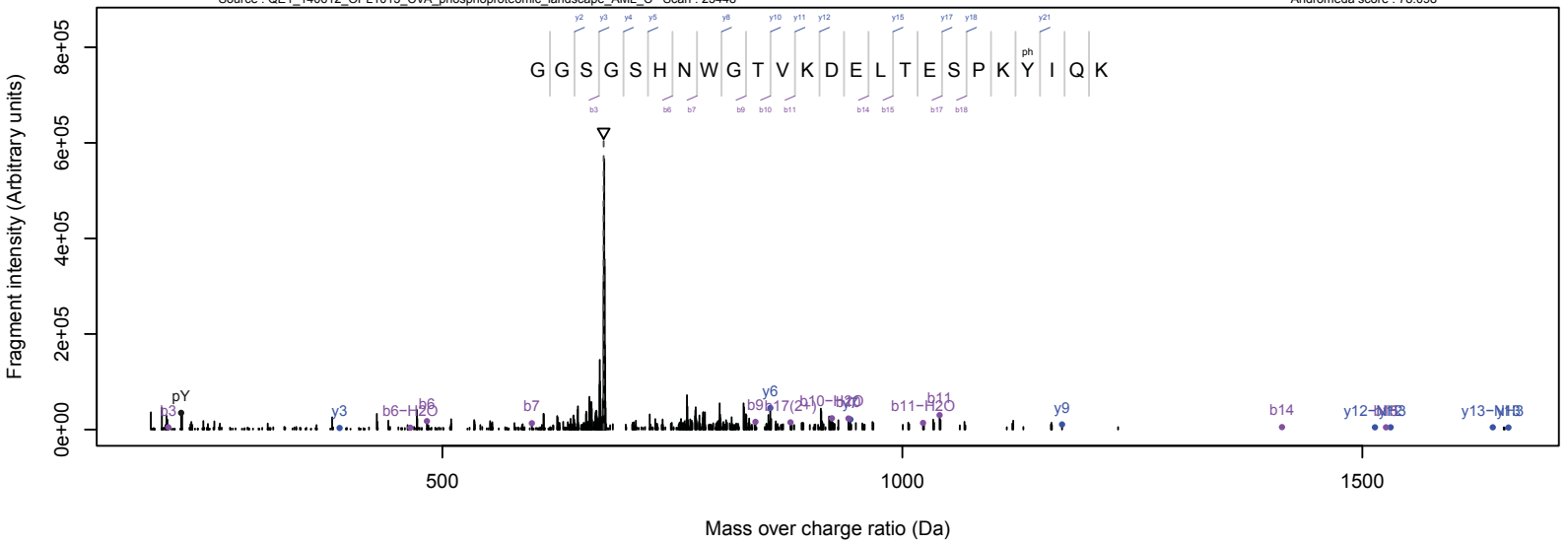
Andromeda score : 78.113



**GGSGSHNWGTVKDELTESPK(ph)IQK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23448

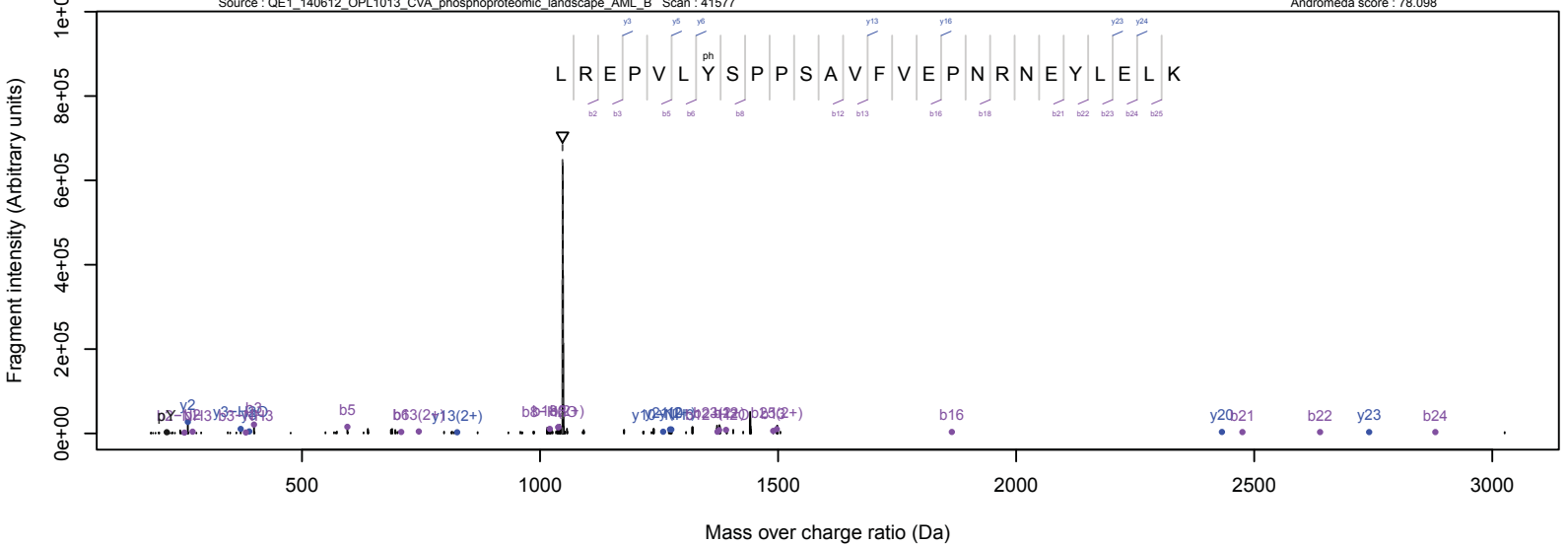
Andromeda score : 78.098



**LREPVL(ph)SPPSAVFVEPNRNEYLELK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41577

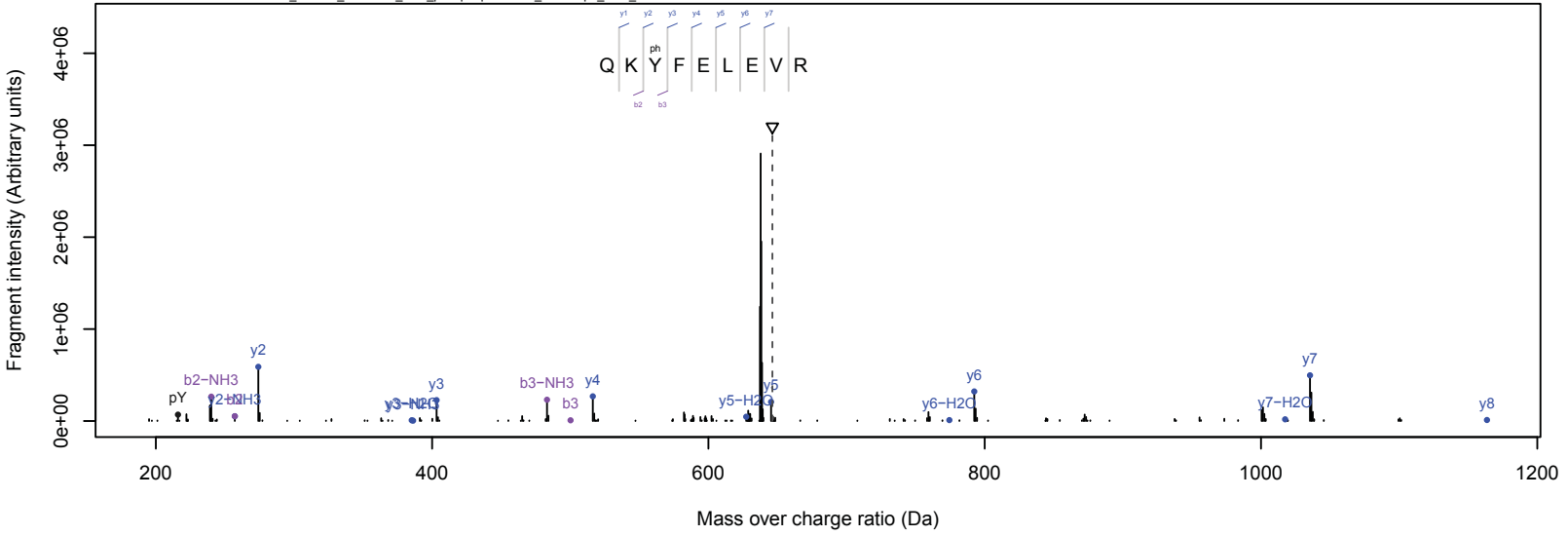
Andromeda score : 78.098



QKY(ph)FELEVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22773

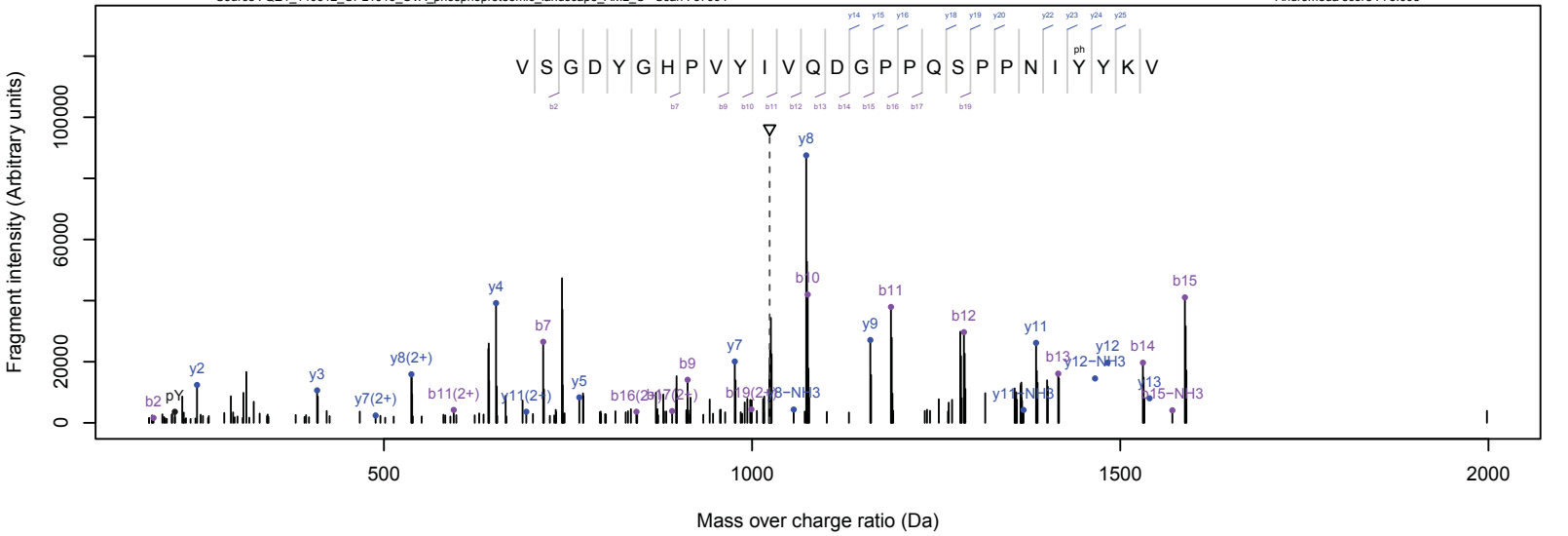
Andromeda score : 78.098



VSGDYGHPVYIQDGPQSPPIY(ph)YKV-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37994

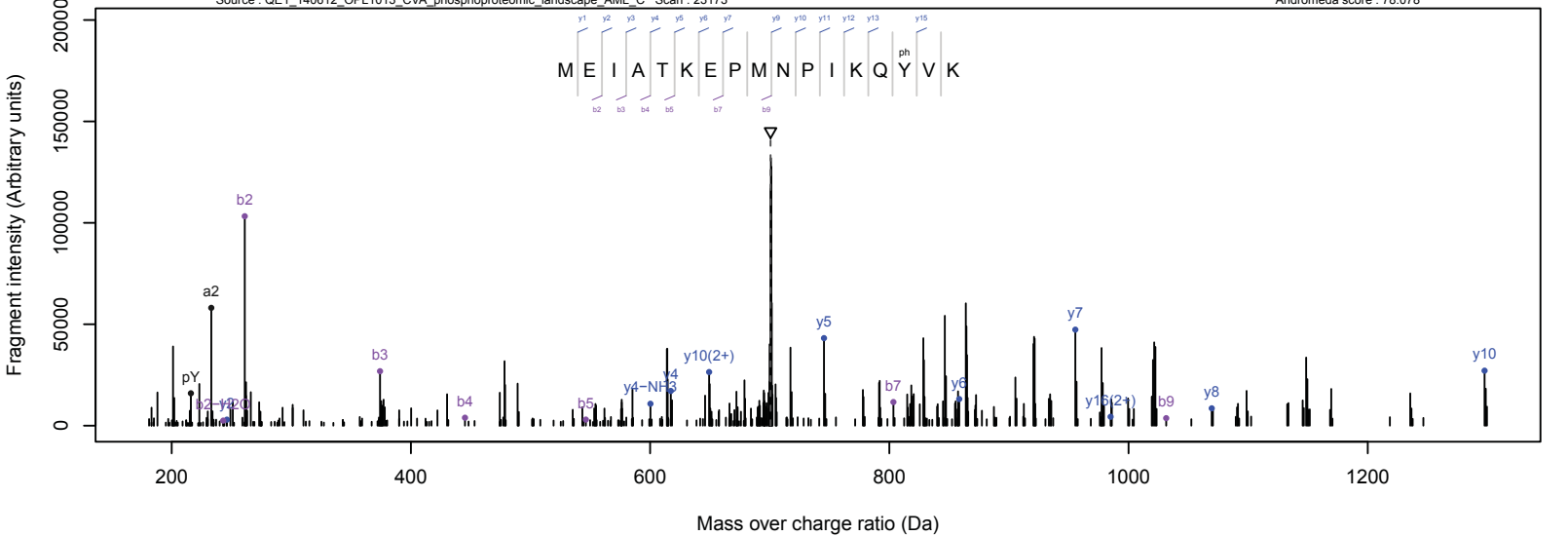
Andromeda score : 78.098



MEIATKEPMNP I K Q Y V K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23173

Andromeda score : 78.078

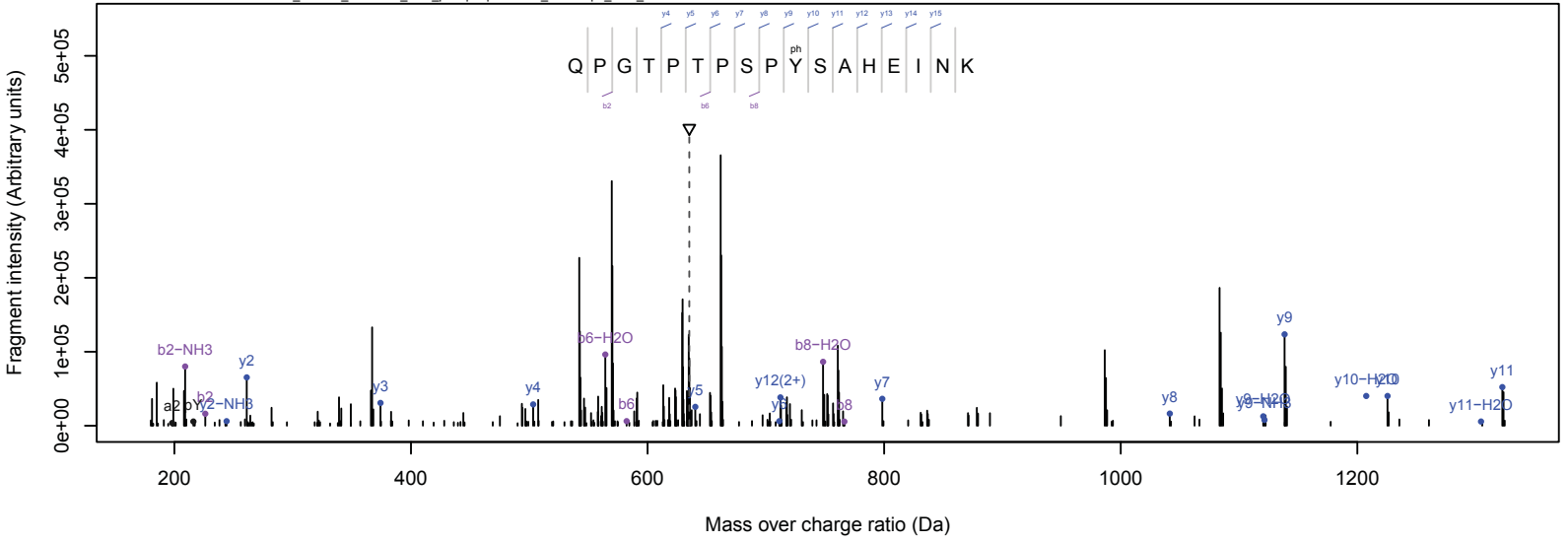




QPGTPTSPY(ph)SAHEINK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11135

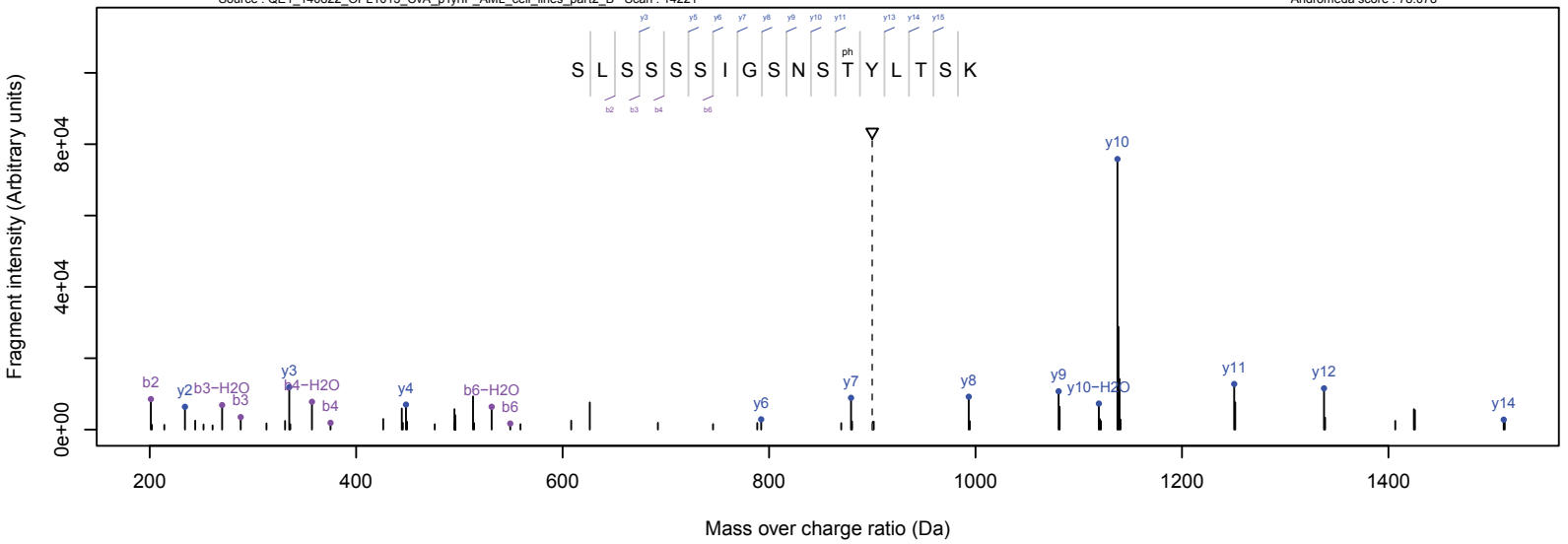
Andromeda score : 78.078



SLSSSSIGSNST(ph)YLTSK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 14221

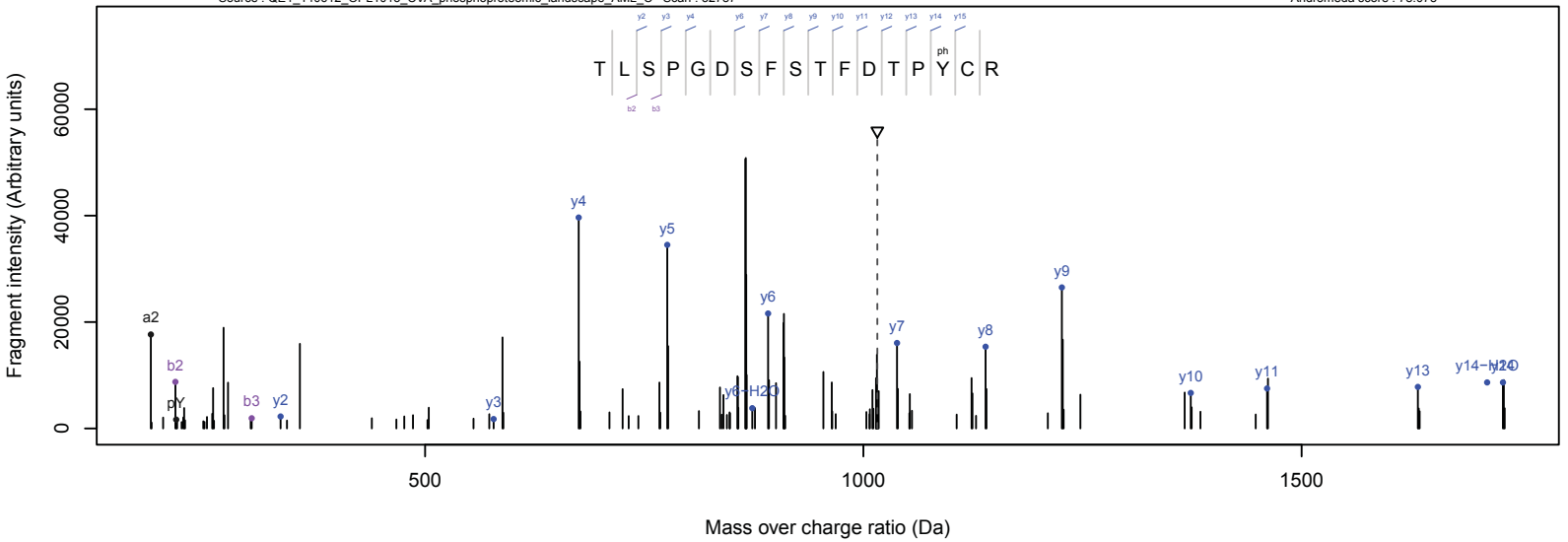
Andromeda score : 78.078



TLSPGDSFSTFDTPY(ph)CR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32787

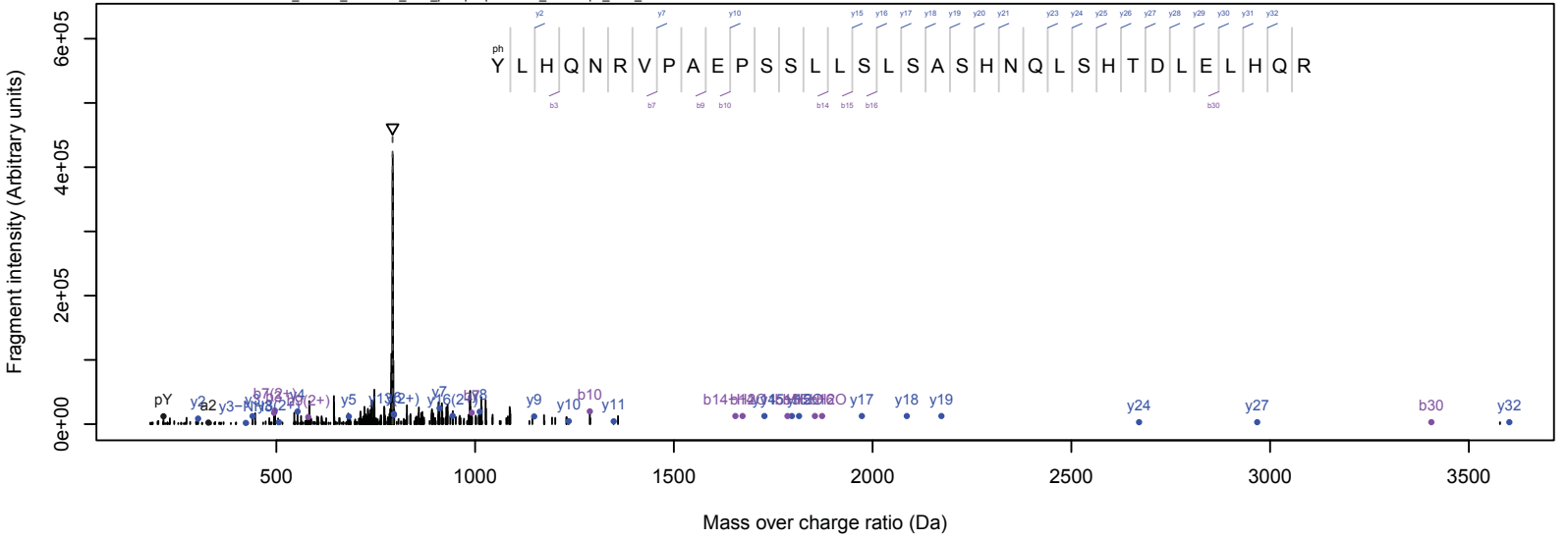
Andromeda score : 78.078



Y(ph)LHQNRVPAEPLSSLLSASHNQLSHTDLELHQR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31957

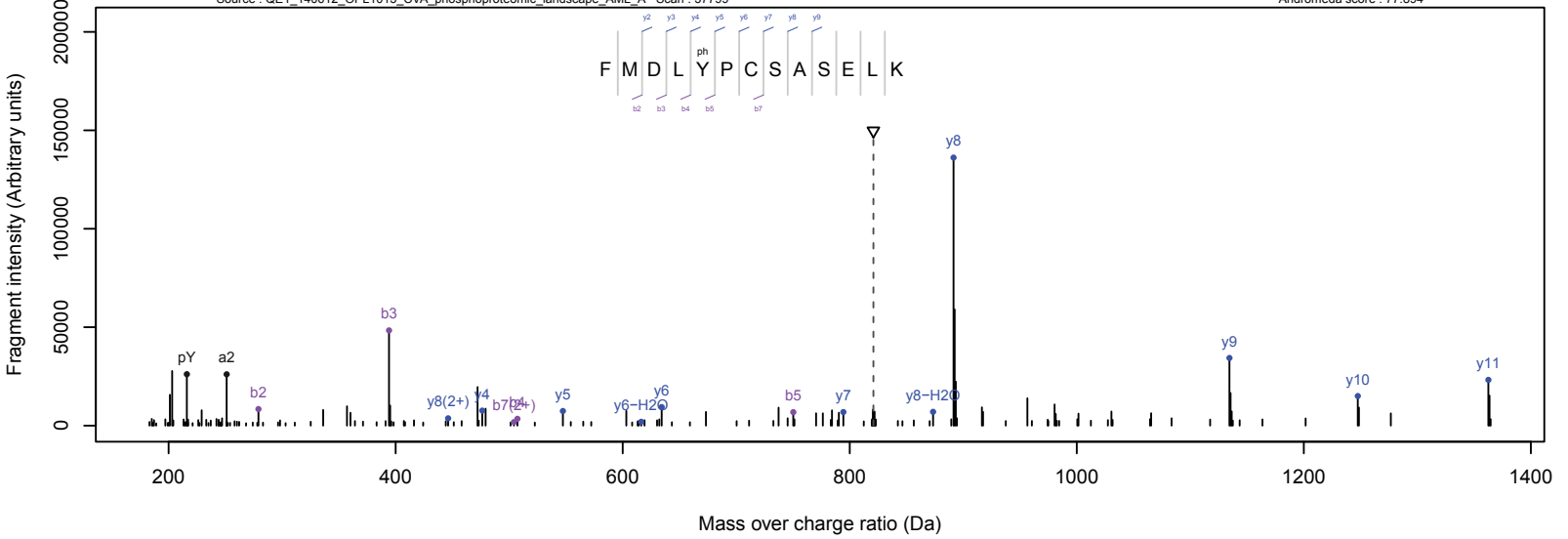
Andromeda score : 77.995



FMDLY(ph)PCSAELK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 37799

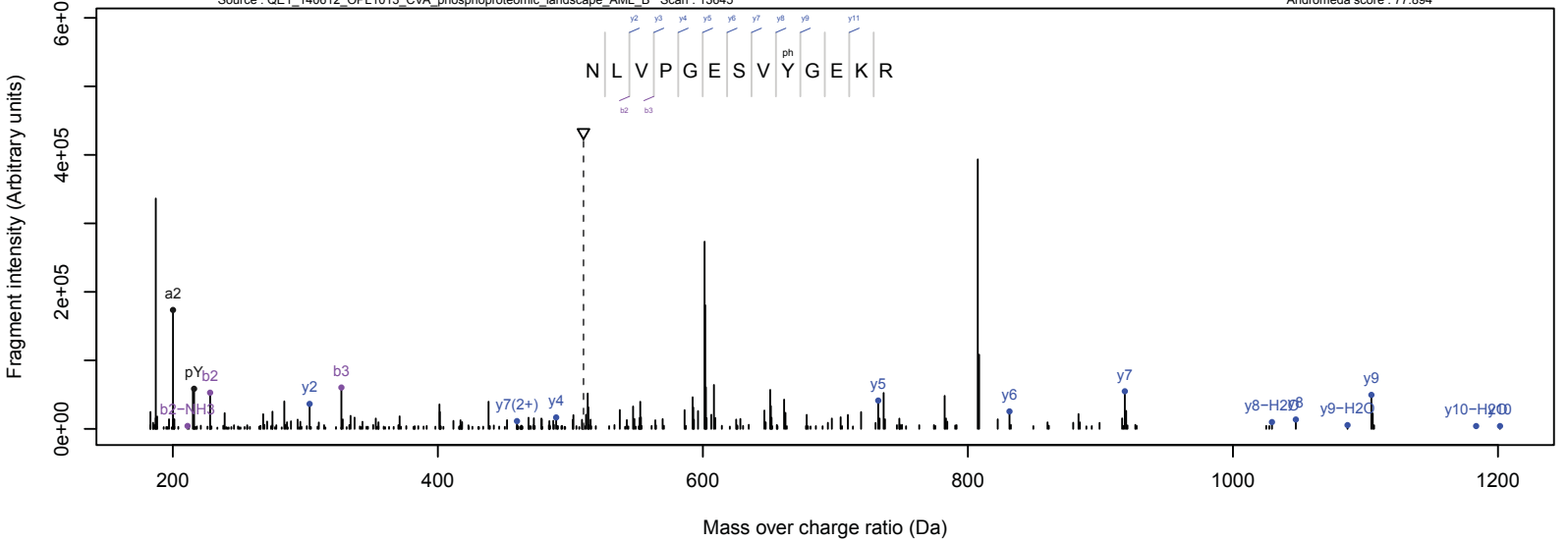
Andromeda score : 77.894



NLVPGESVYGEKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13645

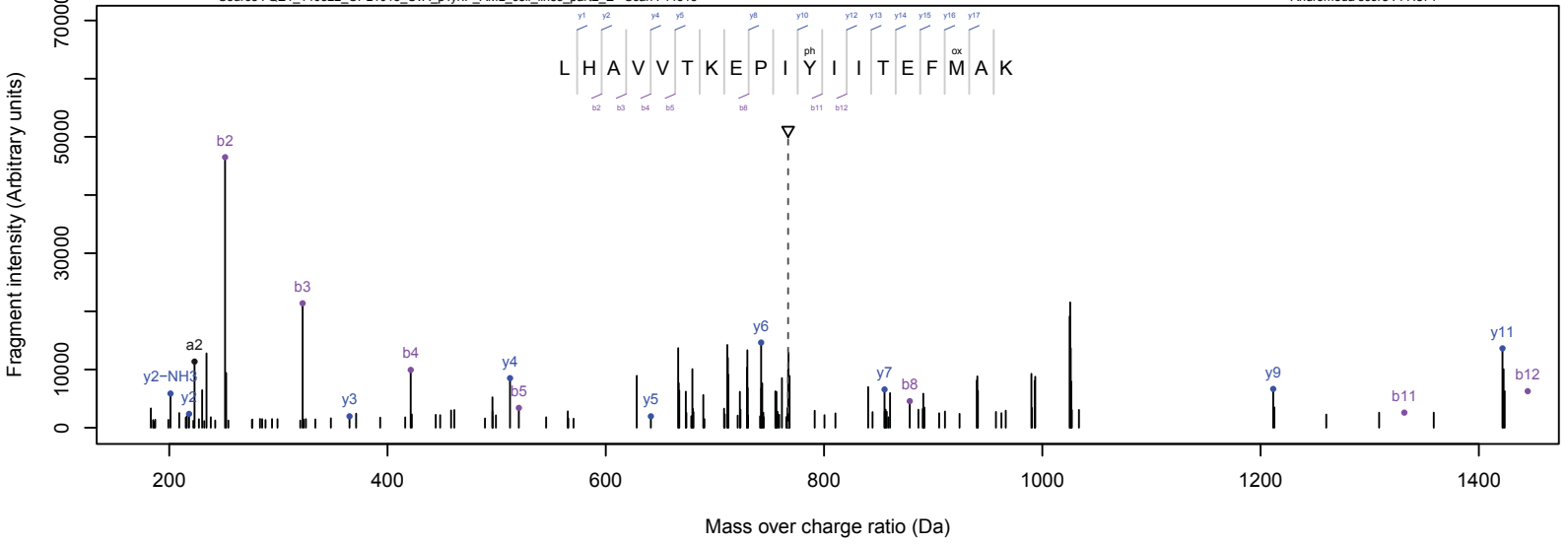
Andromeda score : 77.894



LHAVVTKEPIY(ph)ITEFM(ox)AK-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 44315

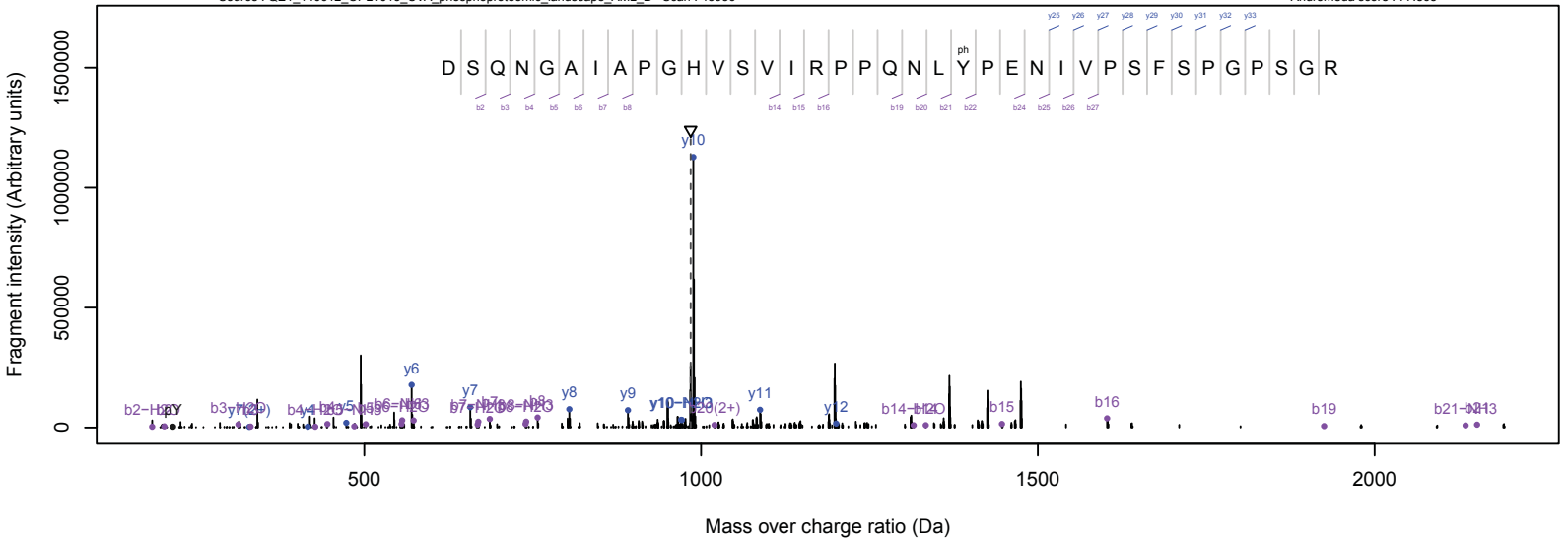
Andromeda score : 77.871



DSQNGAIAPGHVSVIRPPQNL(ph)PENIVPSFSPGSPGR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43936

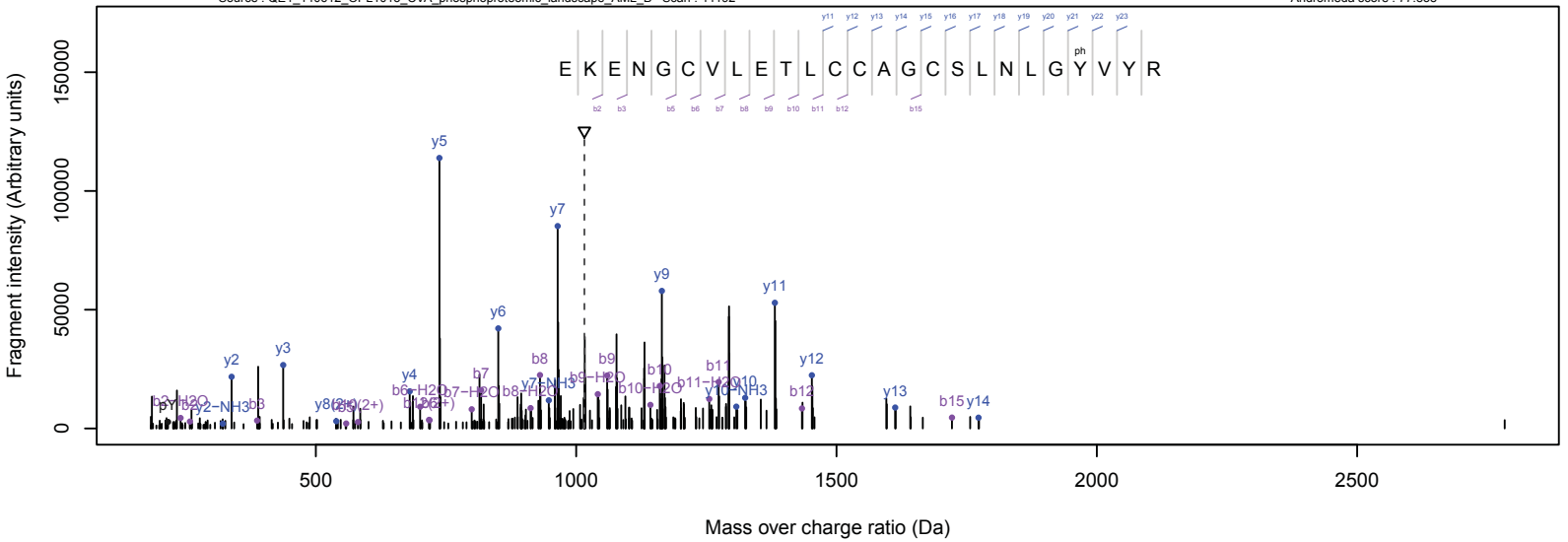
Andromeda score : 77.868



EKENGCVLETLCAGCSLNLG(ph)VYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44192

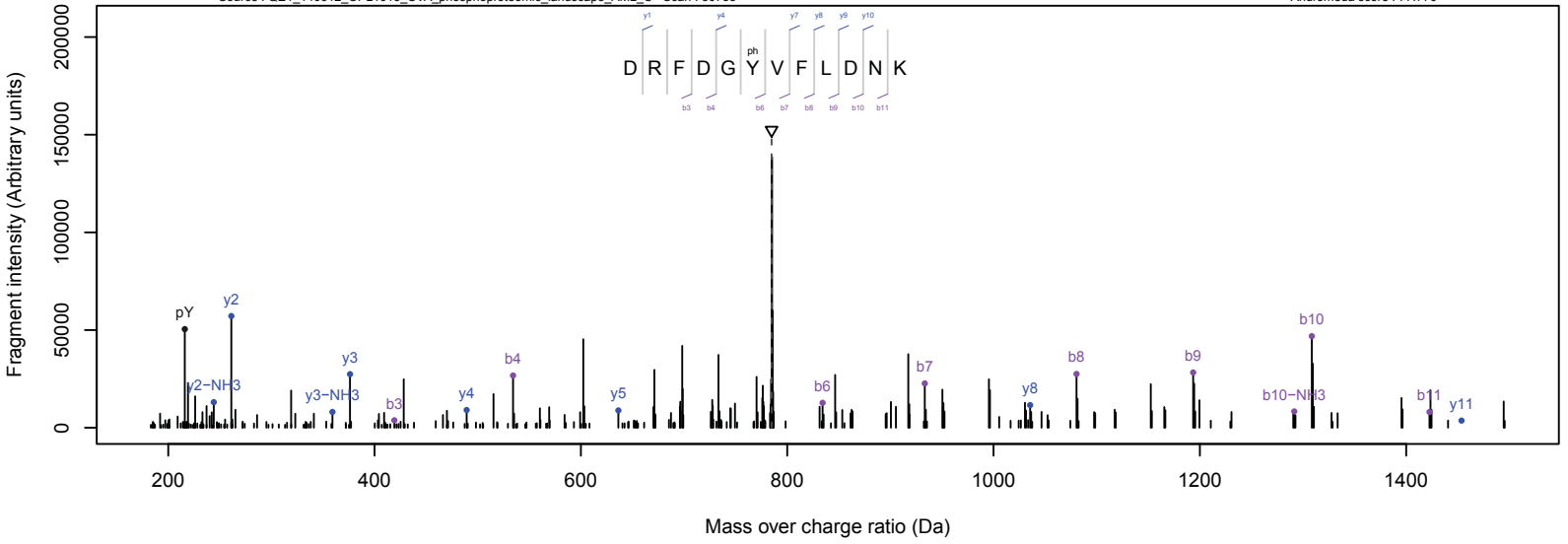
Andromeda score : 77.838



**DRFDGY(ph)VFLDNK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30783

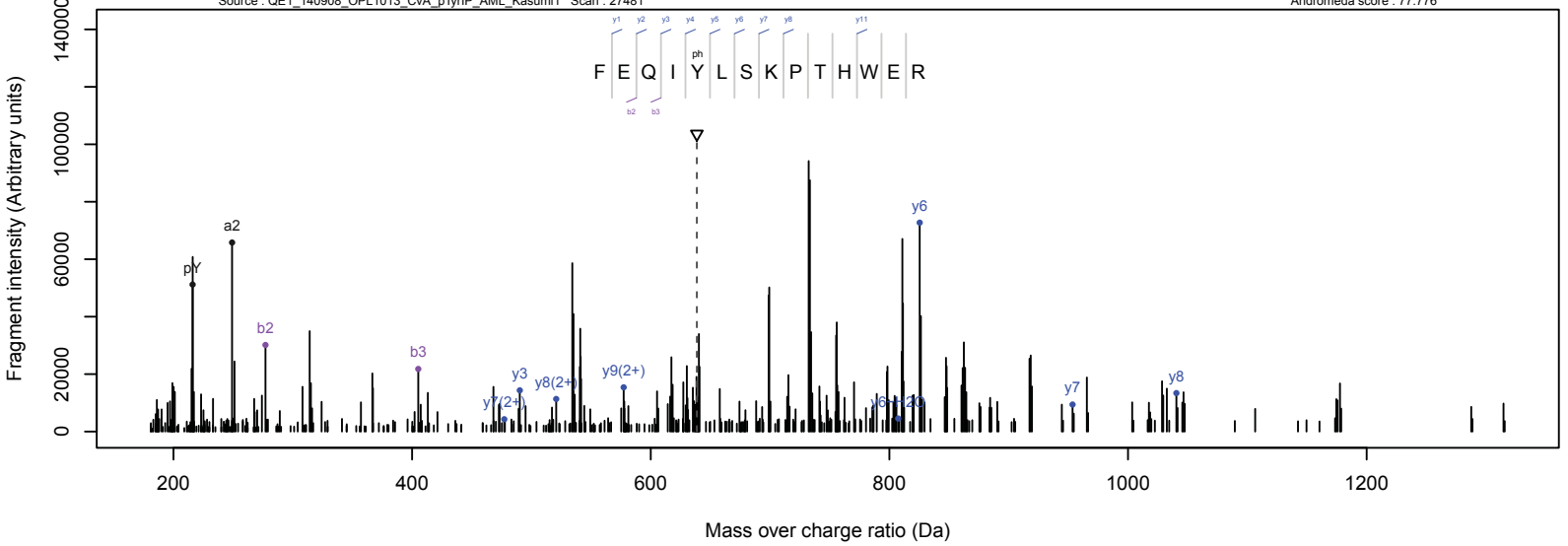
Andromeda score : 77.776



**FEQIY(ph)LSKPTHWER-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27481

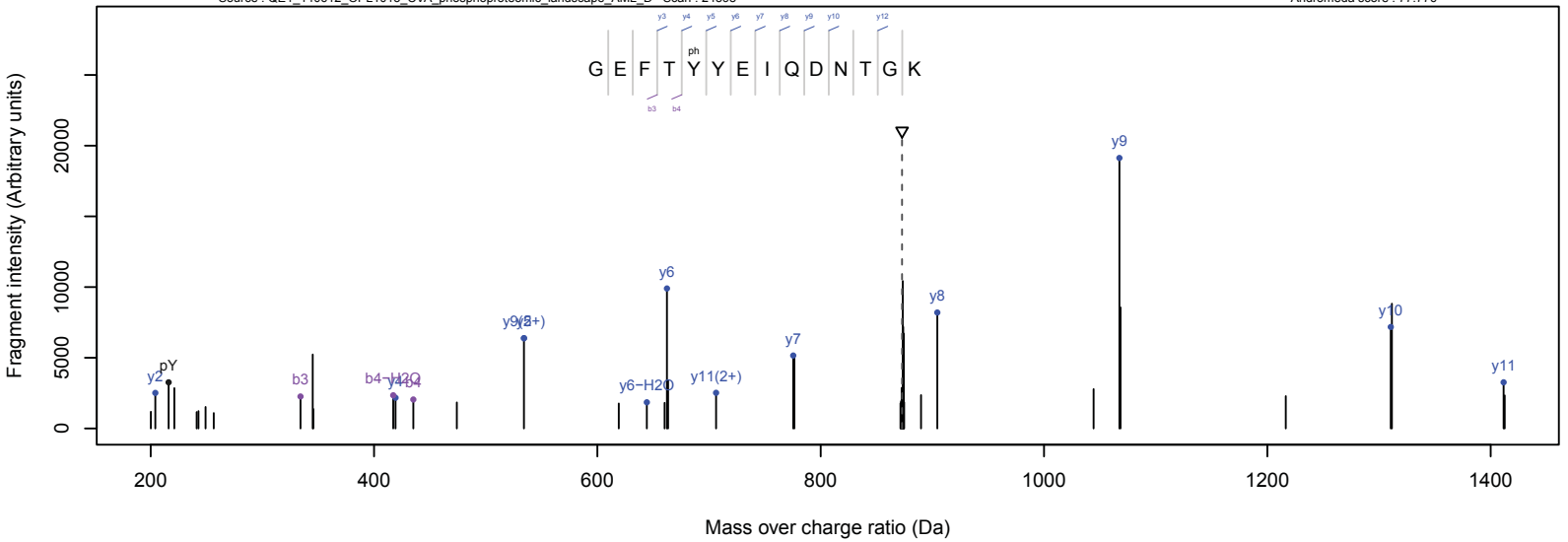
Andromeda score : 77.776



**GEFTY(ph)YEIQDNTGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 24598

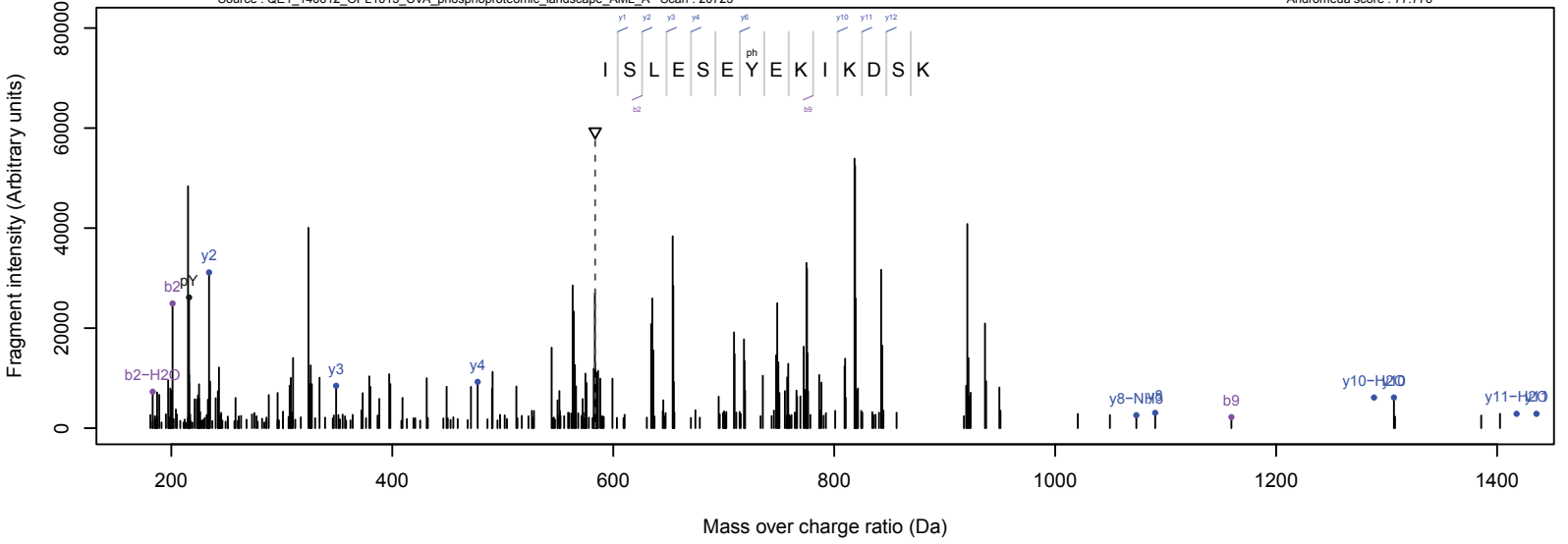
Andromeda score : 77.776



ISLESEY(ph)EKIKDSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 20723

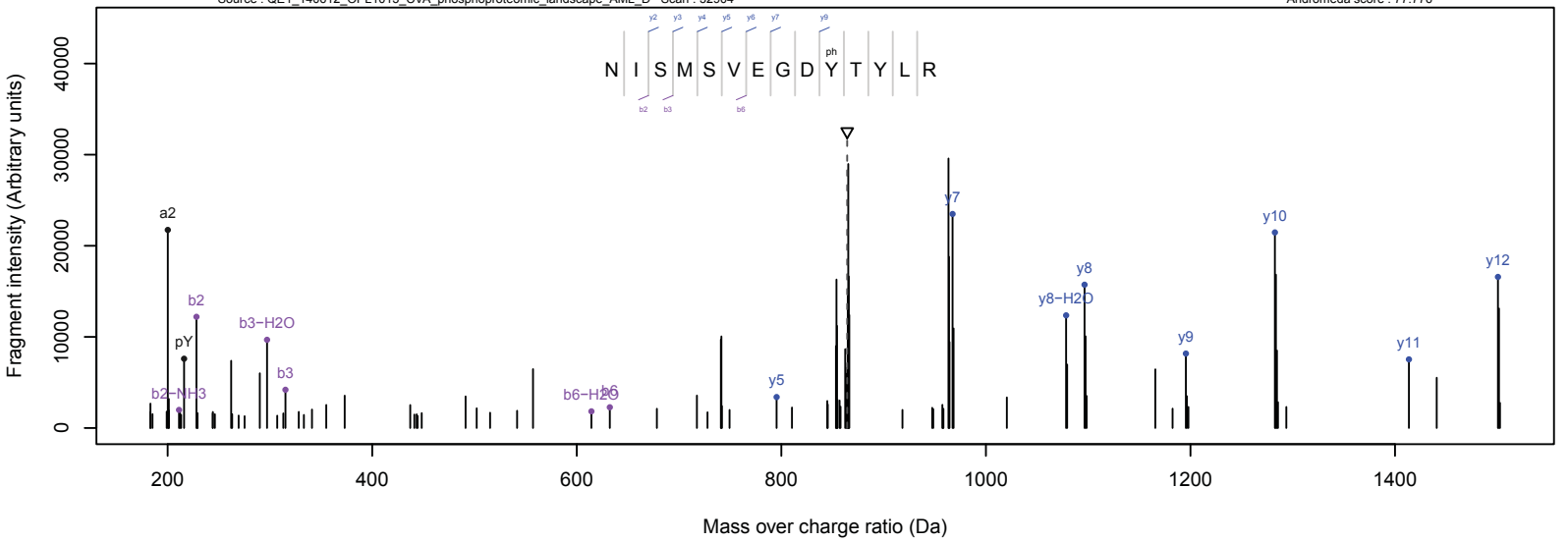
Andromeda score : 77.776



NISMSVEGDY(ph)TYLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 32904

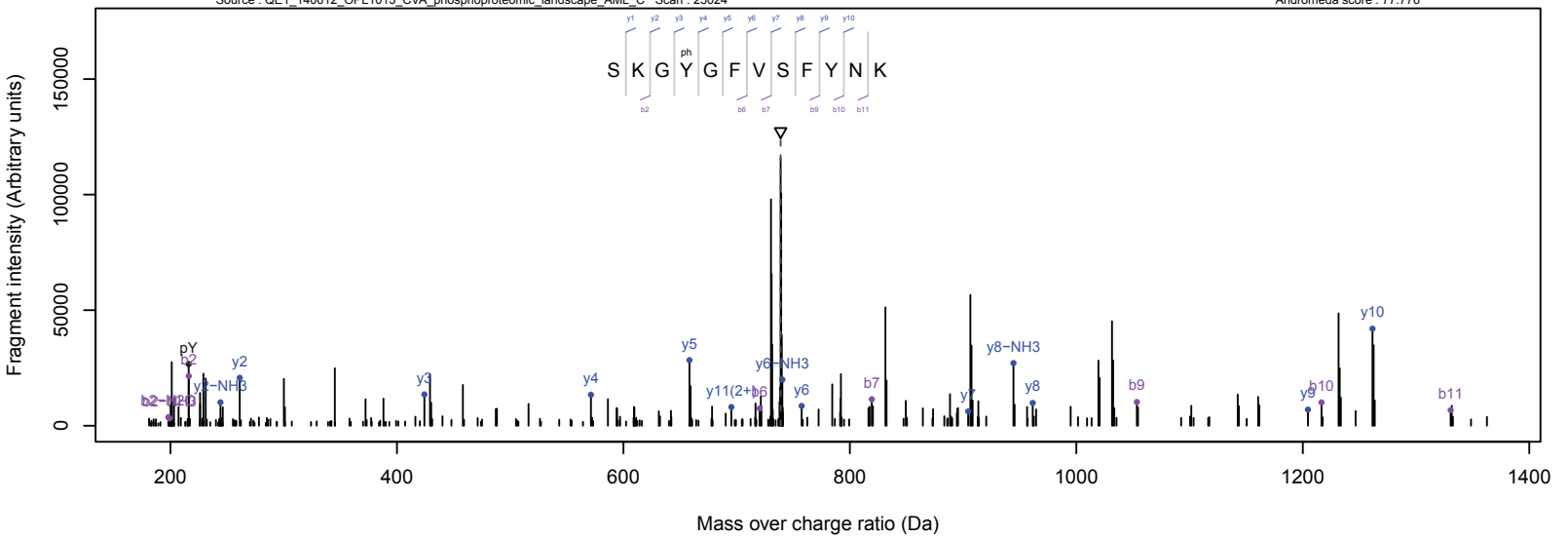
Andromeda score : 77.776



SKGY(ph)GFVSFYNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25024

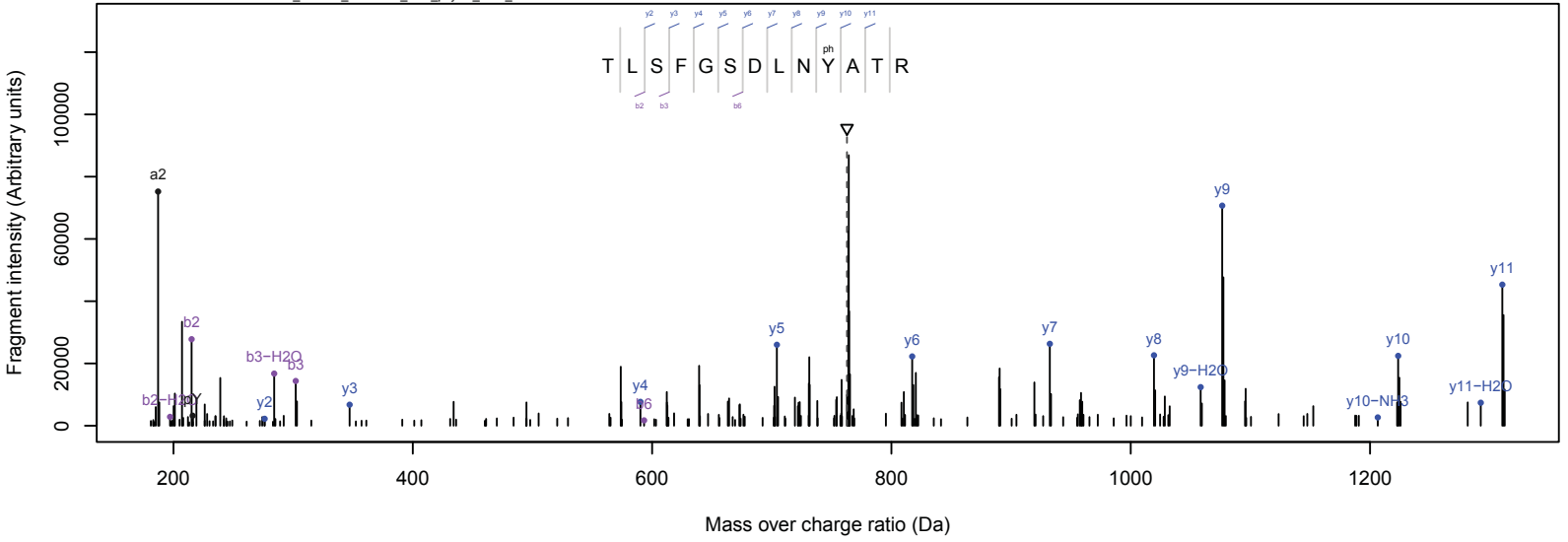
Andromeda score : 77.776



**TLSFGSDLNY(ph)ATR-2+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 30170

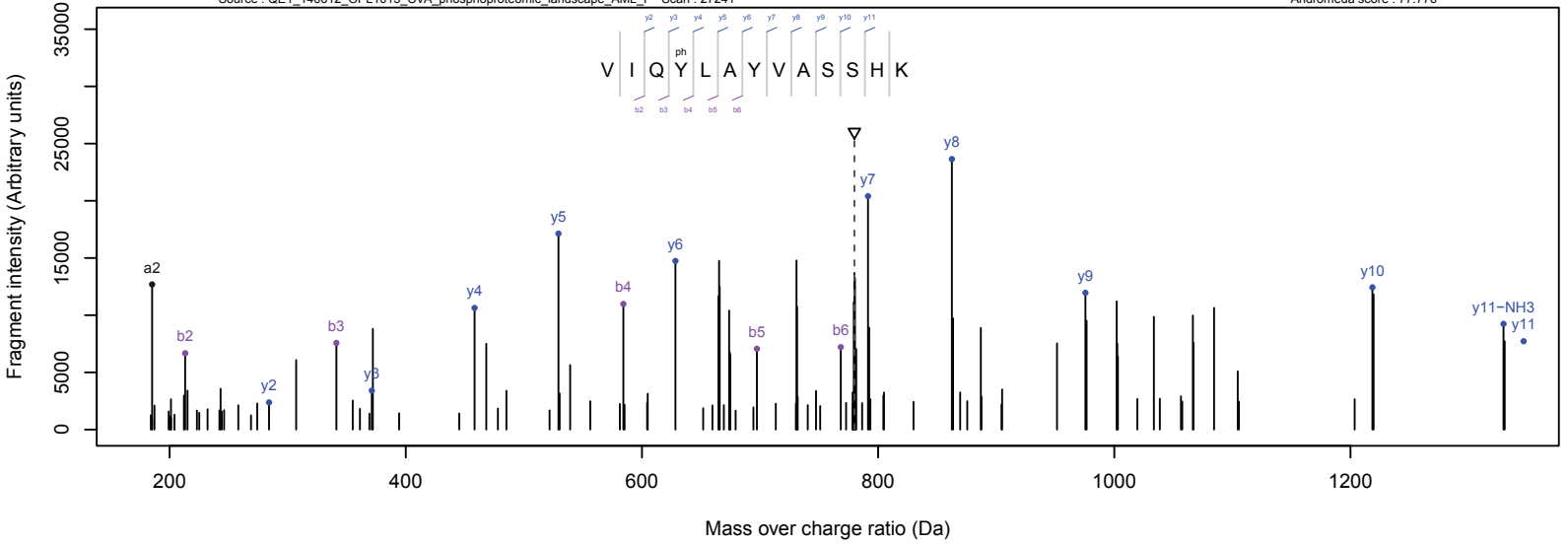
Andromeda score : 77.776



**VIQY(ph)LAYASSHK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 27241

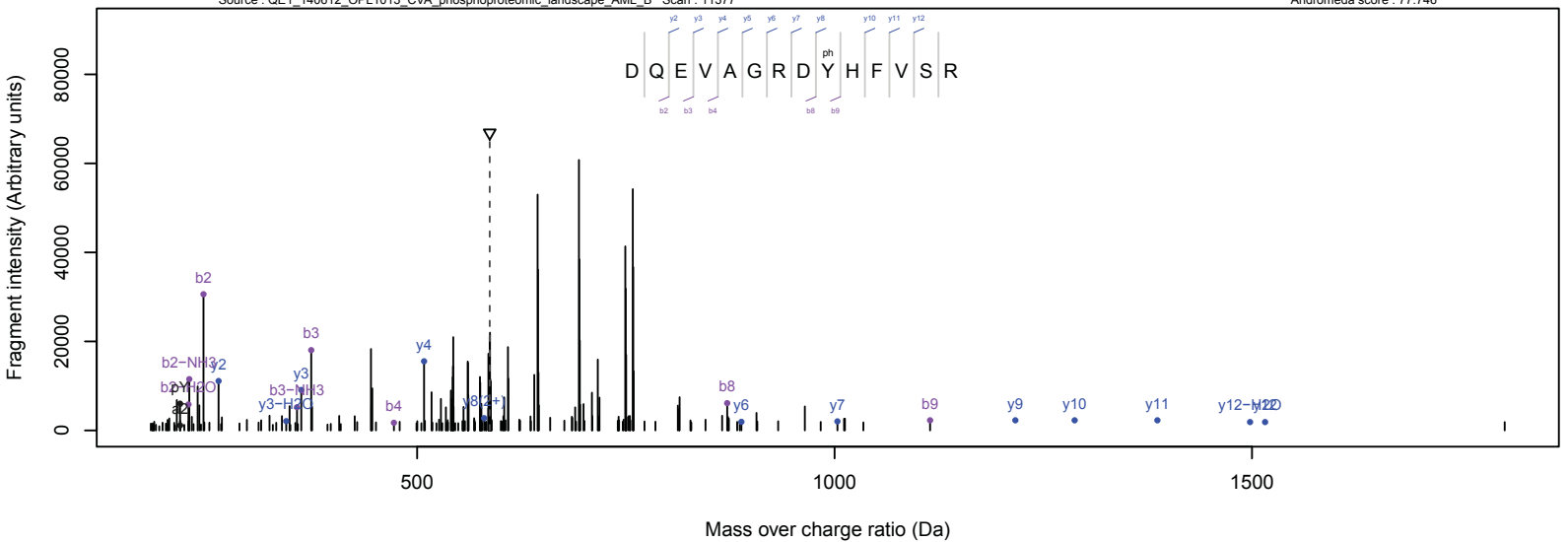
Andromeda score : 77.776



**DQEVAGRDY(ph)HFVSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11377

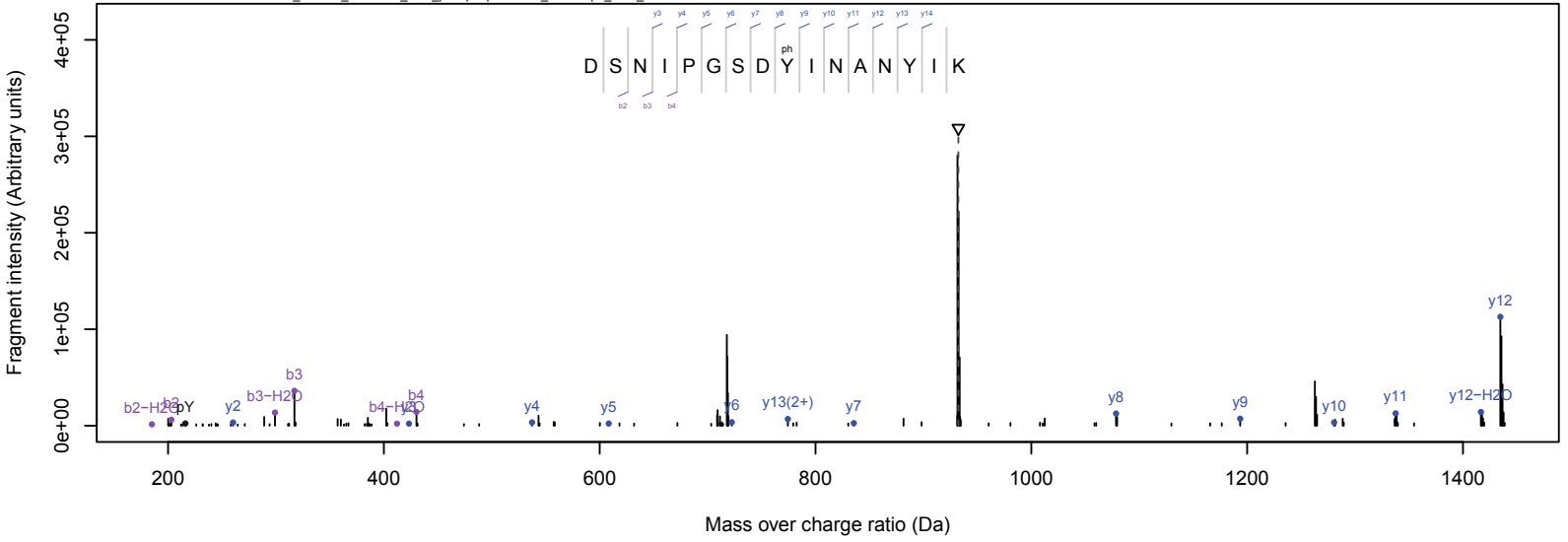
Andromeda score : 77.746



**DSNIPGSDY(ph)INANYIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32992

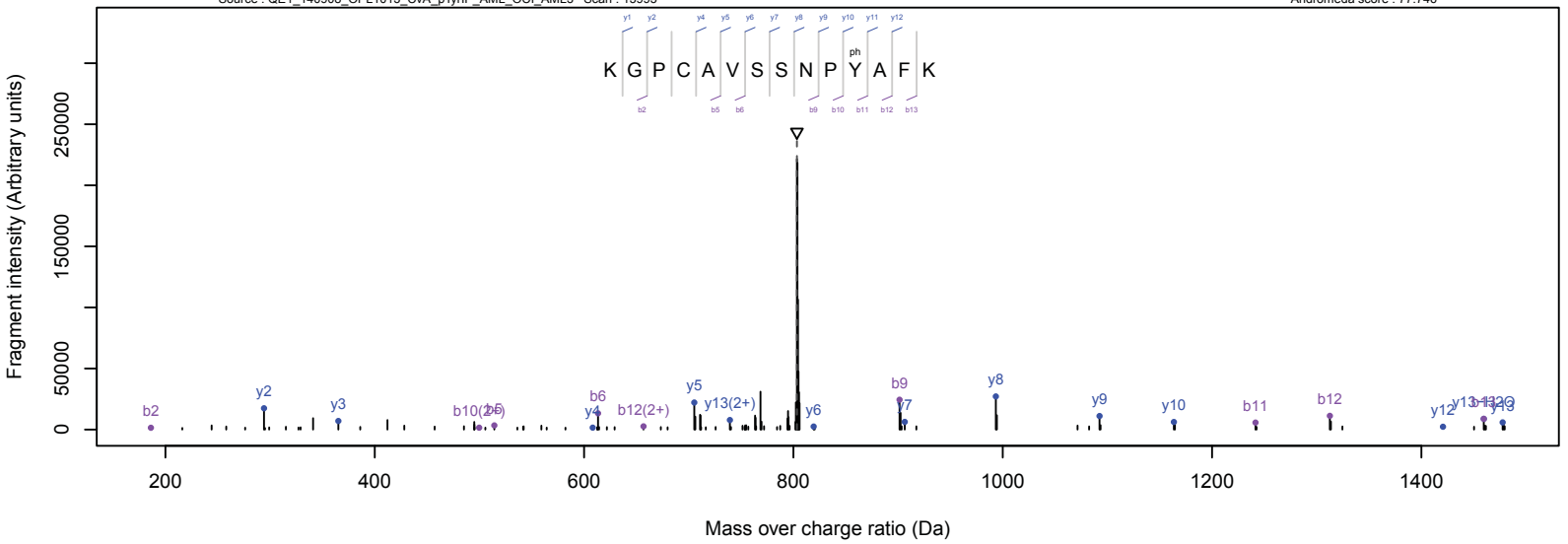
Andromeda score : 77.746



**KGPCAVSSNPY(ph)AFK-2+**

Source : QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_OCI\_AML3 Scan : 13993

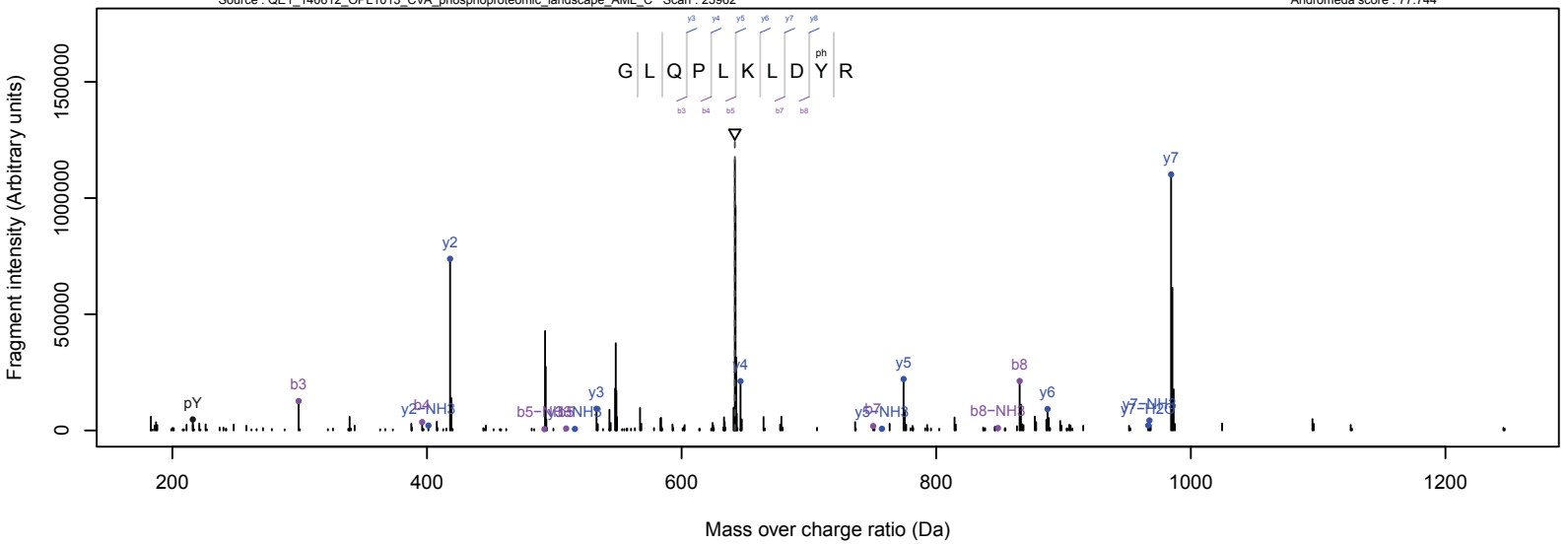
Andromeda score : 77.746



**GLQLKLDY(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23962

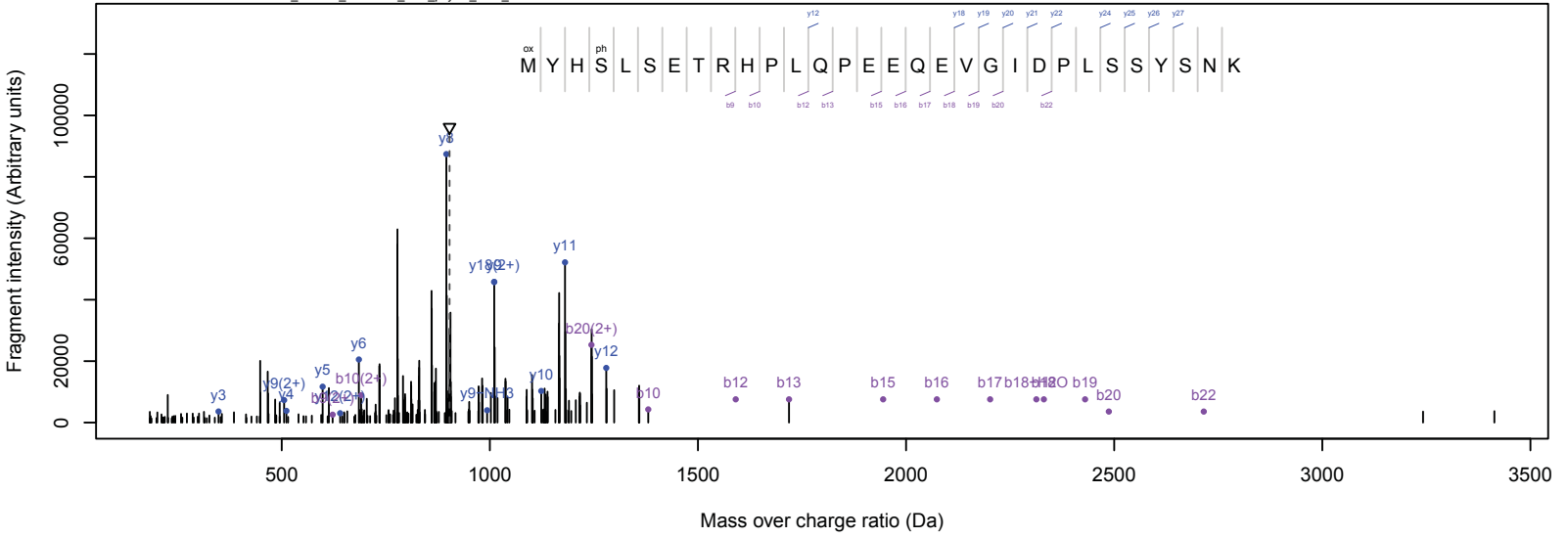
Andromeda score : 77.744



(ac)M(ox)YHS(ph)LSETRHPLQPEEQEVGIDPLSSYSNK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 33817

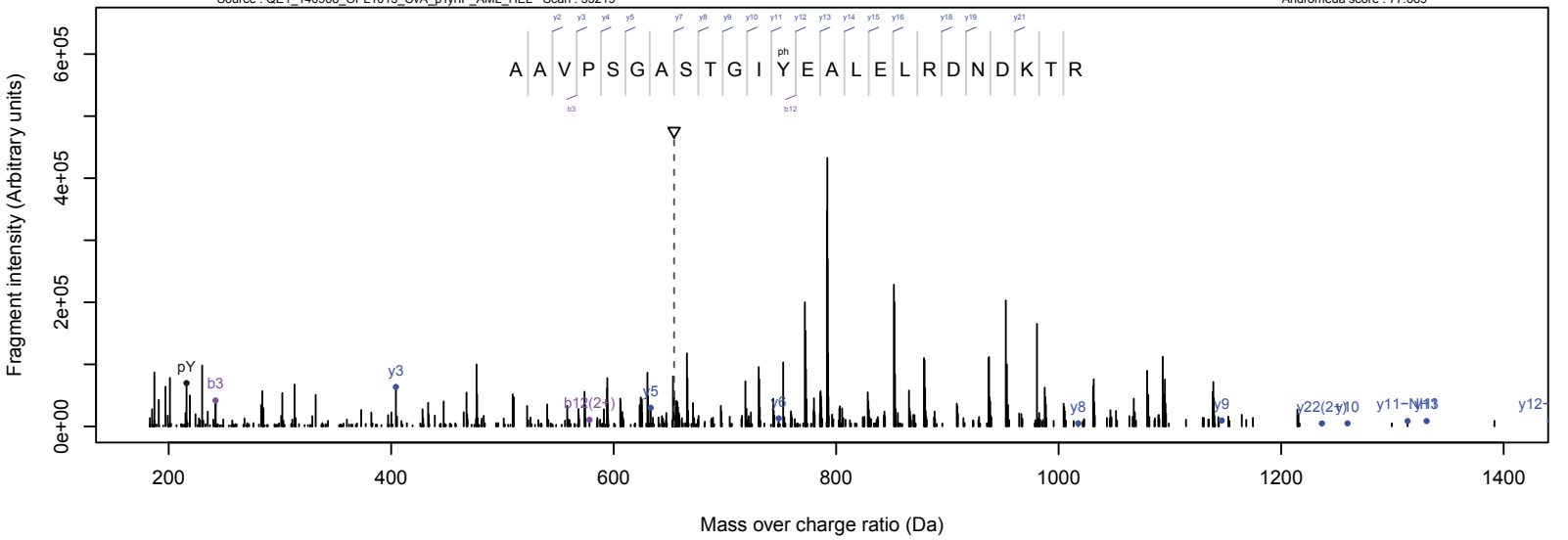
Andromeda score : 77.731



AAVPSGASTGIY(ph)EALELRDNDKTR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 33219

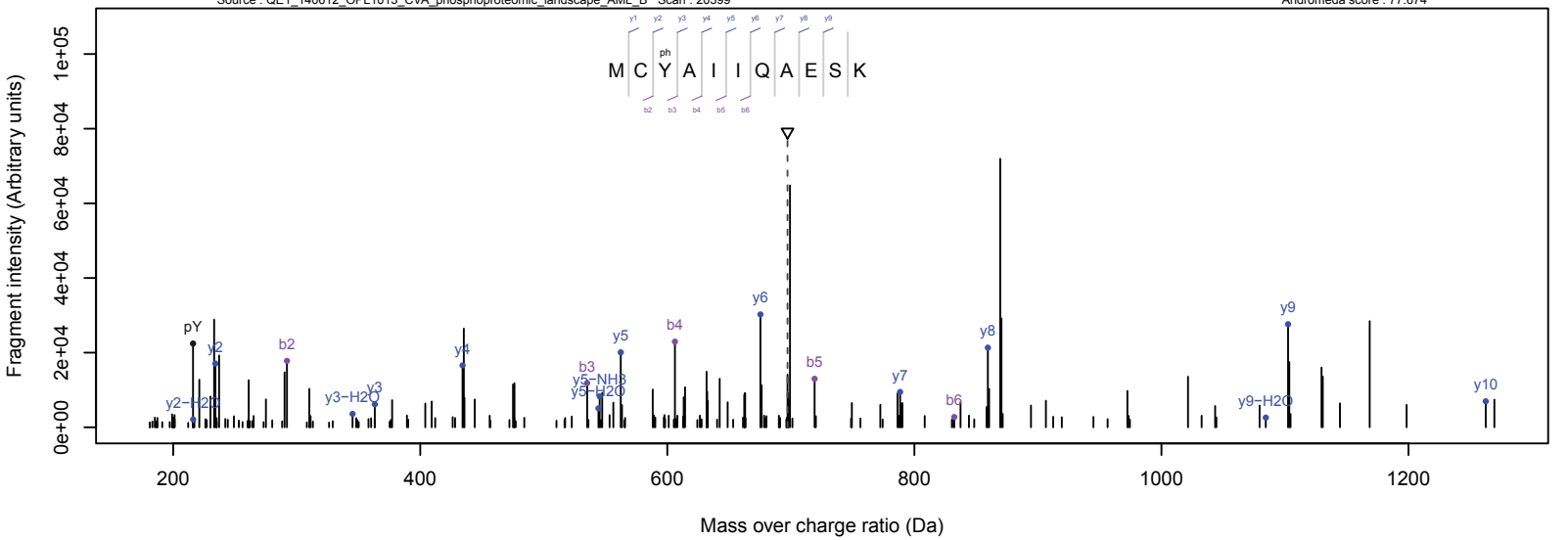
Andromeda score : 77.689



MCY(ph)AIQAESK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20399

Andromeda score : 77.674

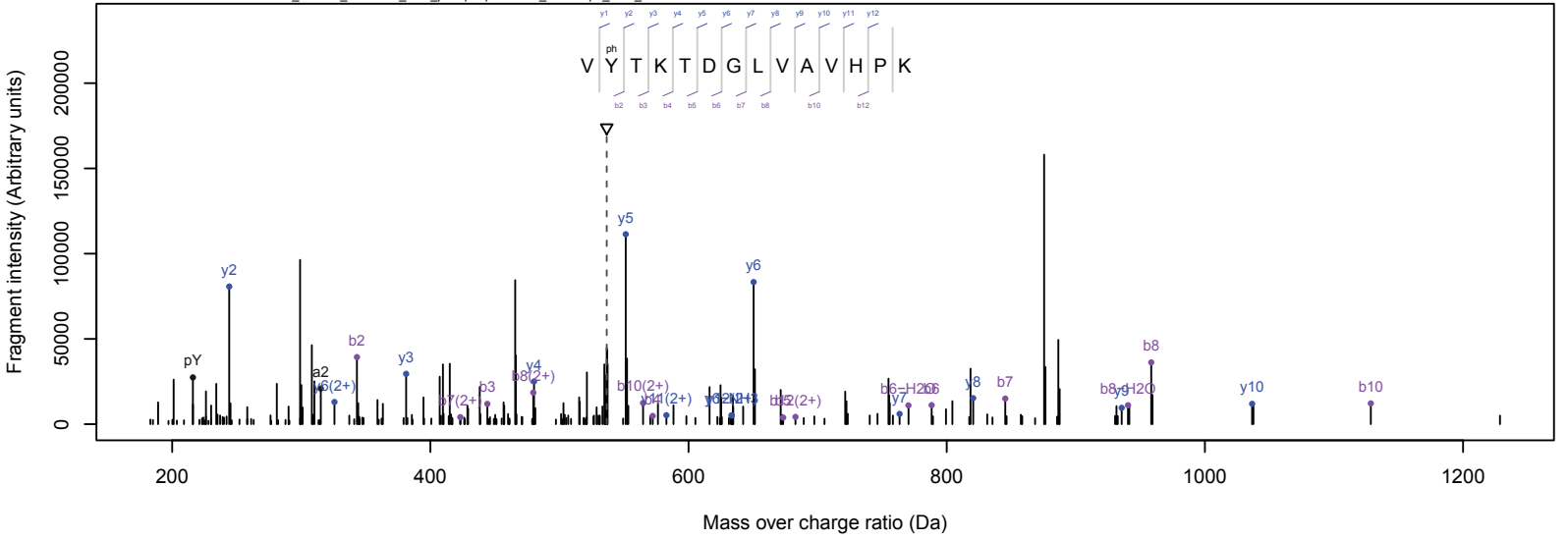




VY(ph)TKTDGLVAVHPK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 13528

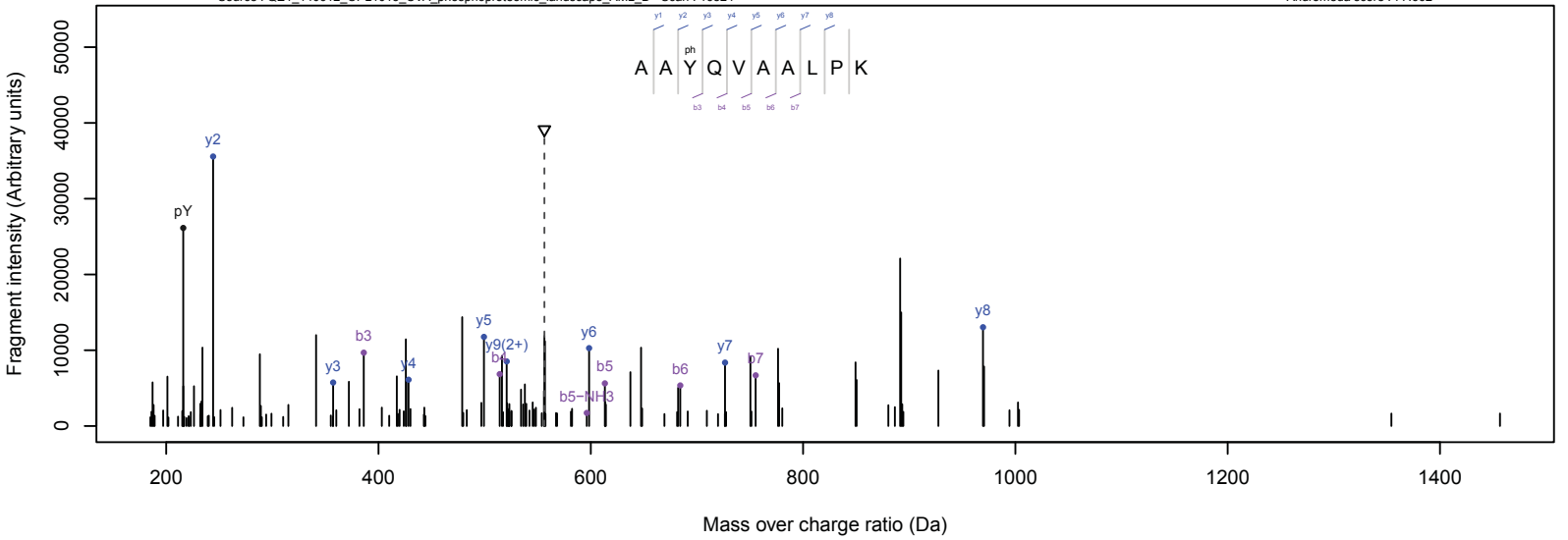
Andromeda score: 77.664



AA Y(ph)QVAALPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 16824

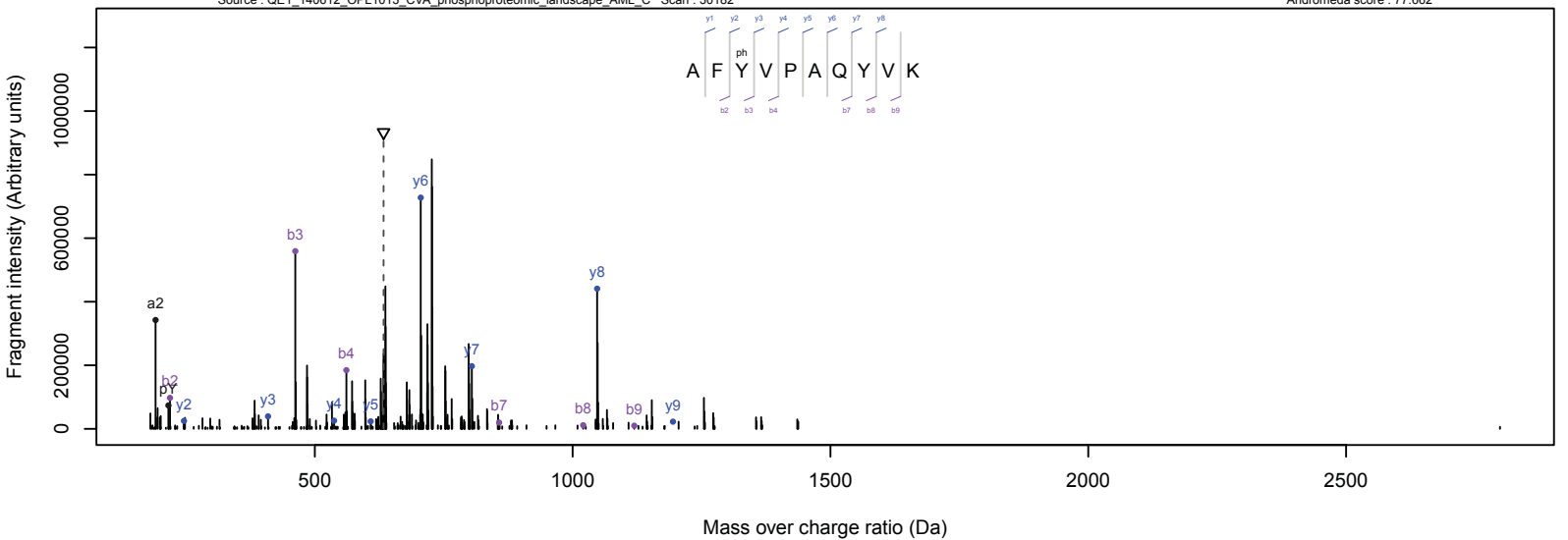
Andromeda score: 77.662



AF Y(ph)VPAQYVK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 30182

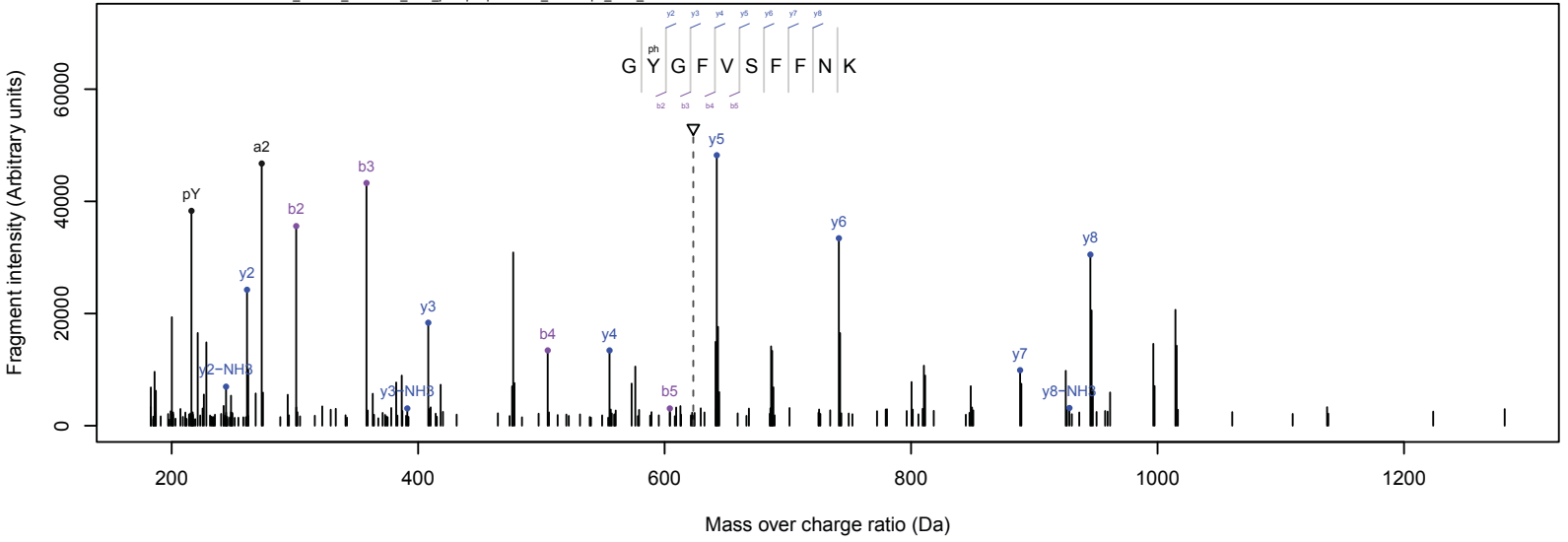
Andromeda score: 77.662



**GY(ph)GFVSFFNK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41356

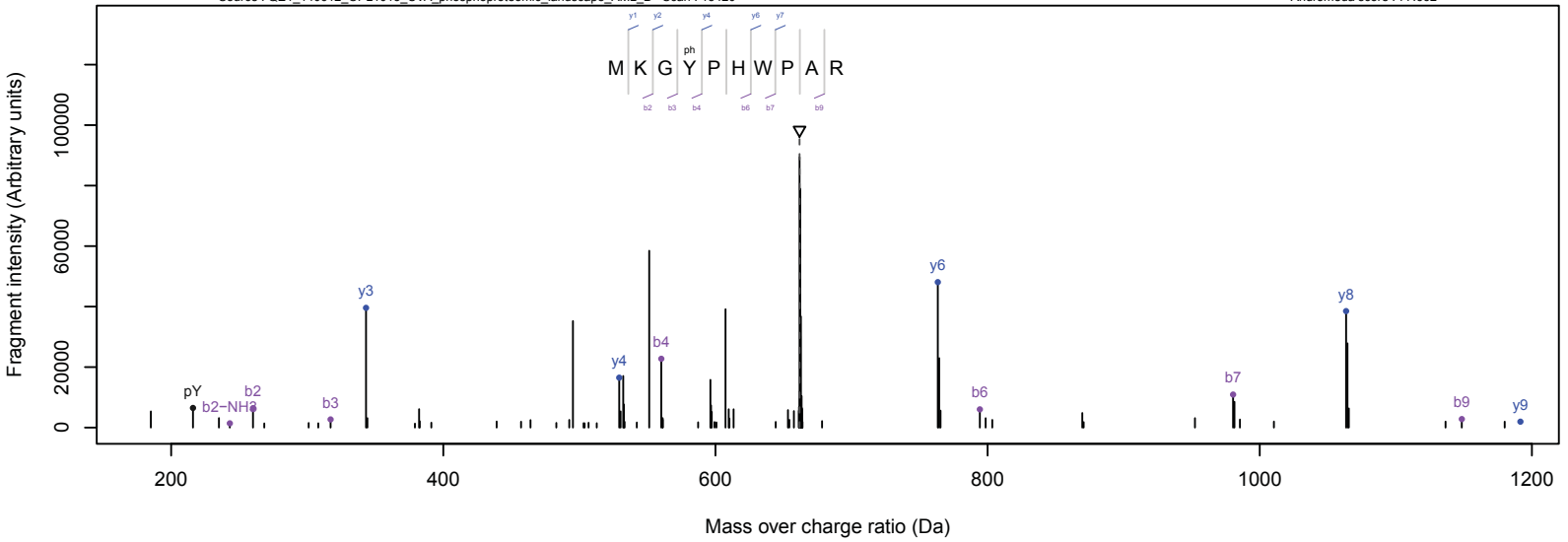
Andromeda score : 77.662



**MKGY(ph)PHWPAR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13420

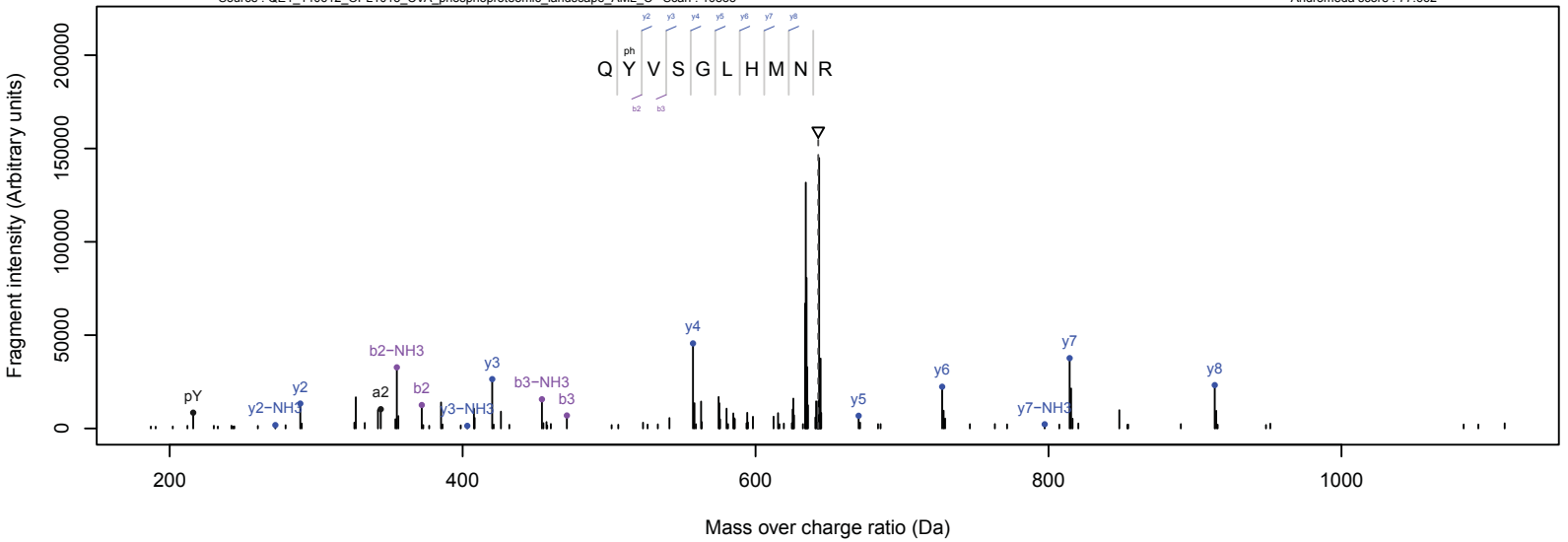
Andromeda score : 77.662



**QY(ph)VSLHMNR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10835

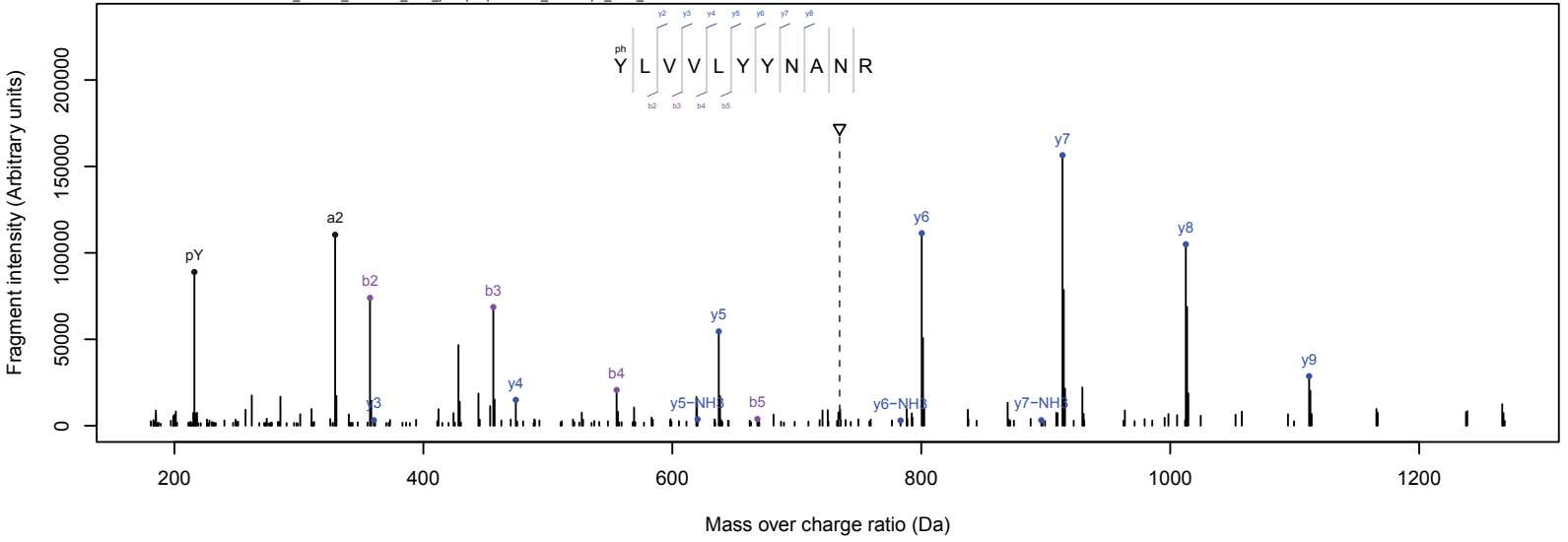
Andromeda score : 77.662



**Y(ph)LVLLYNNANR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36871

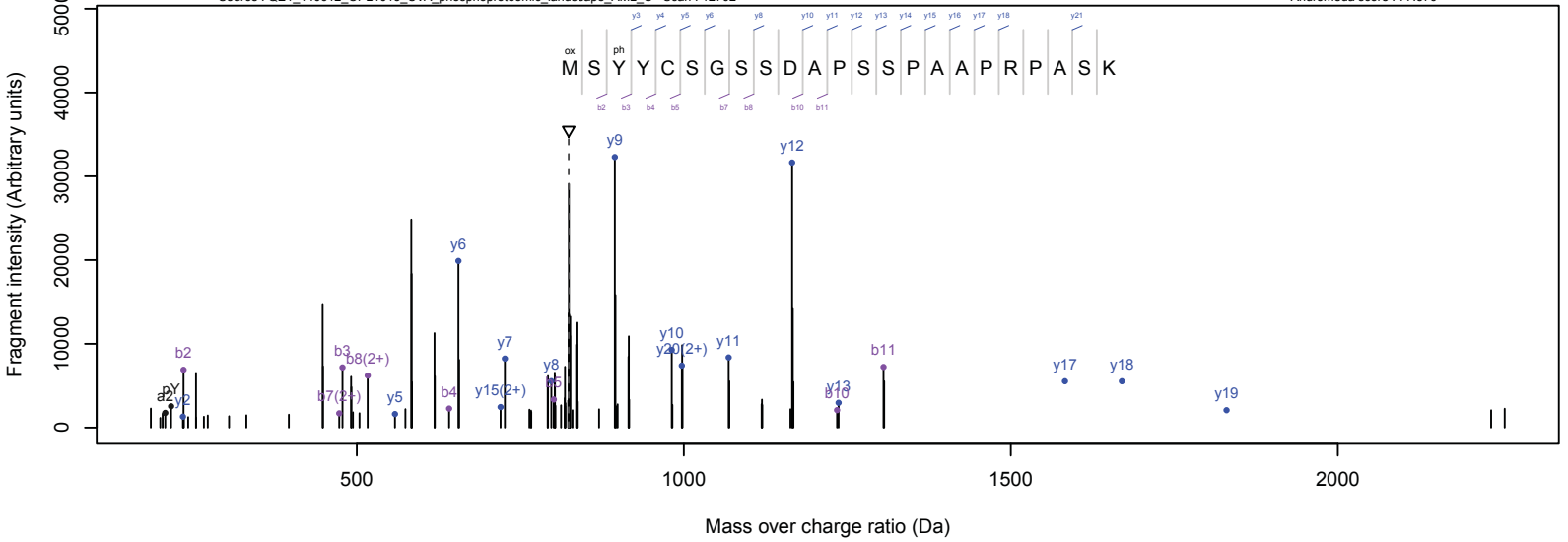
Andromeda score : 77.662



**M(ox)SY(ph)YCSGSSDAPSSPAAPRPASK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12702

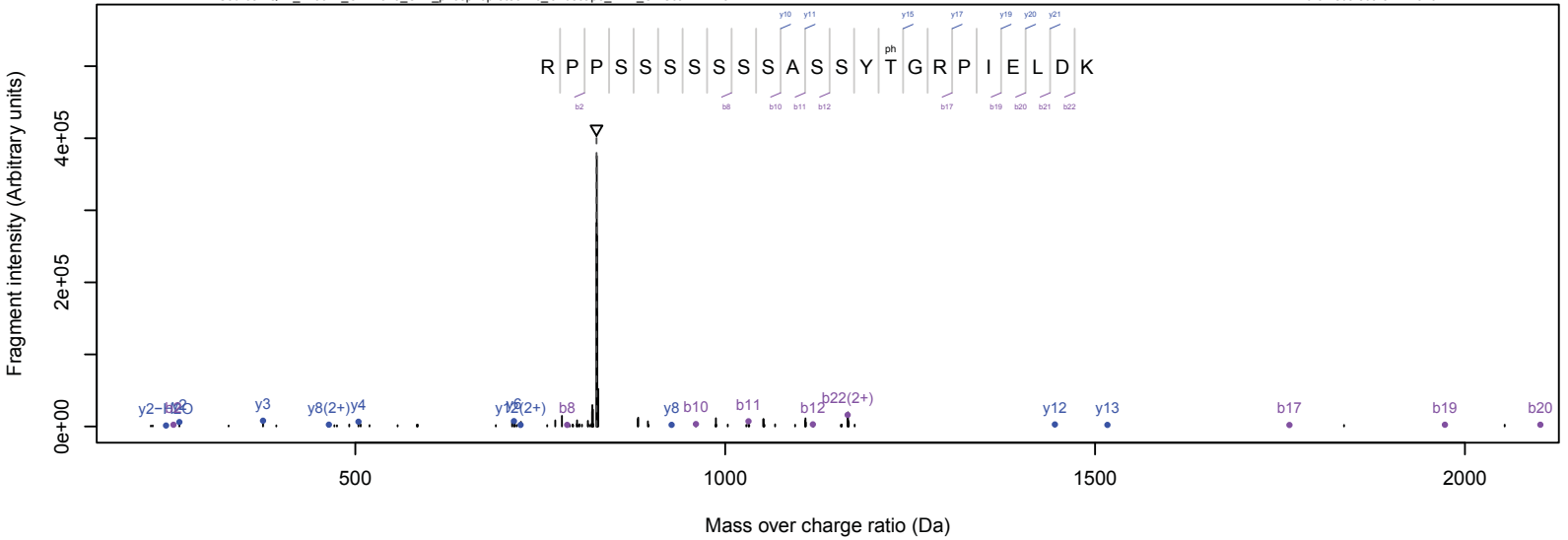
Andromeda score : 77.576



**RPPSSSSSSASSY(ph)GRPIELDK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12151

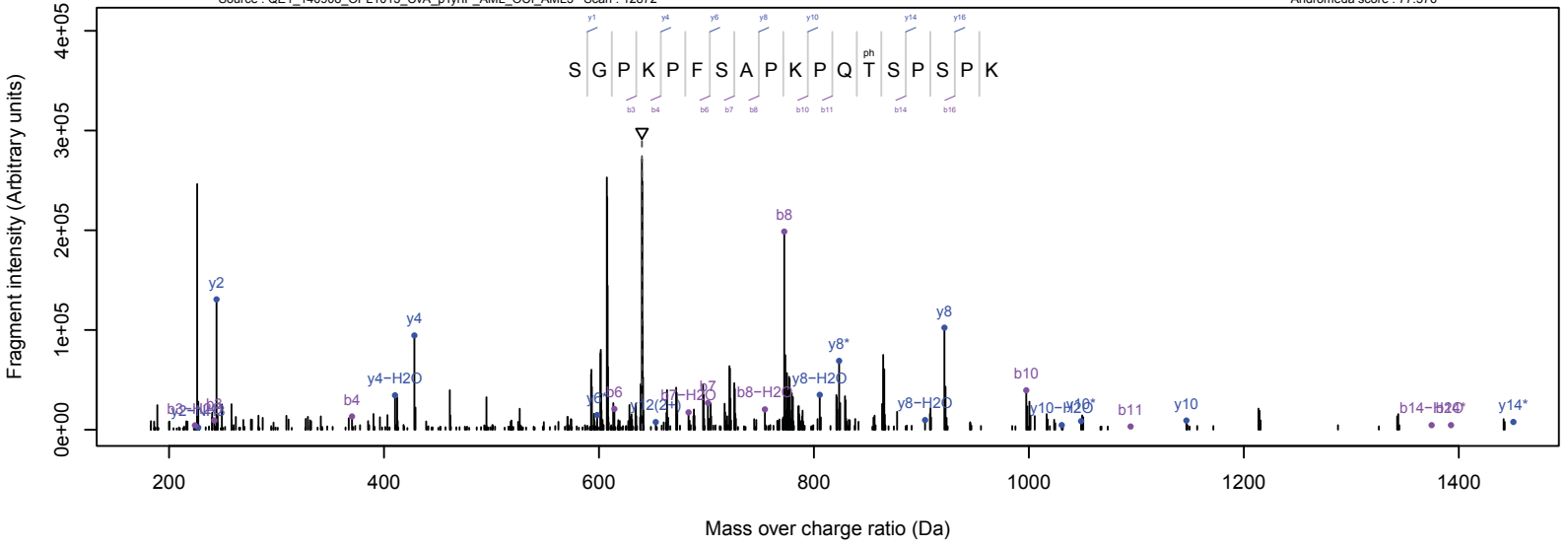
Andromeda score : 77.576



SGPKPFSAPK(ph)SPSPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 12872

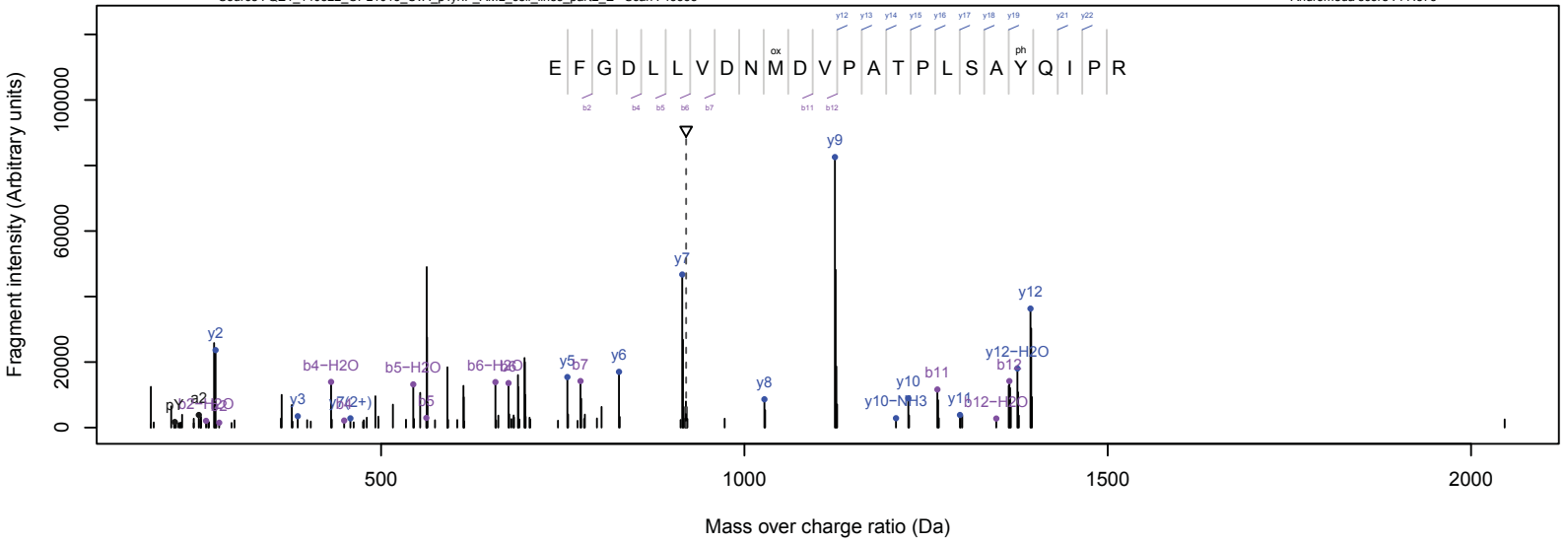
Andromeda score : 77.576



EFGDLLVDNM(ox)DVPATPLSAY(ph)QIPR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 43338

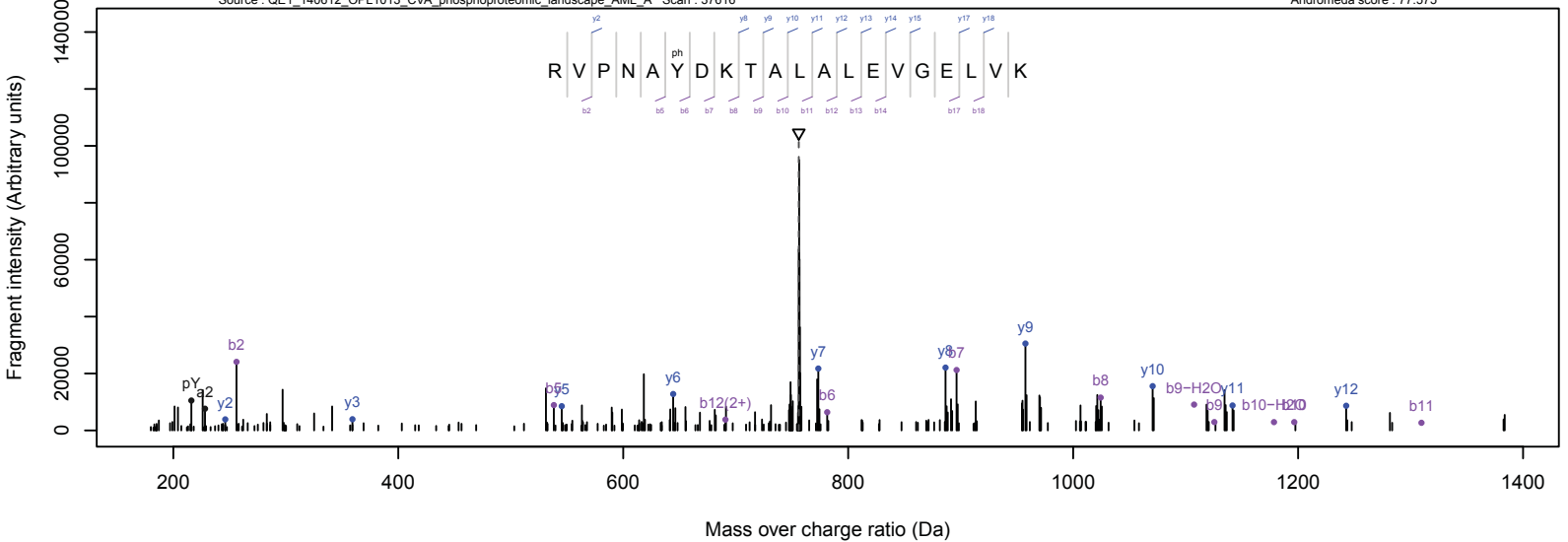
Andromeda score : 77.575



RVPNAY(ph)DKTALALEVGE L V K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 37616

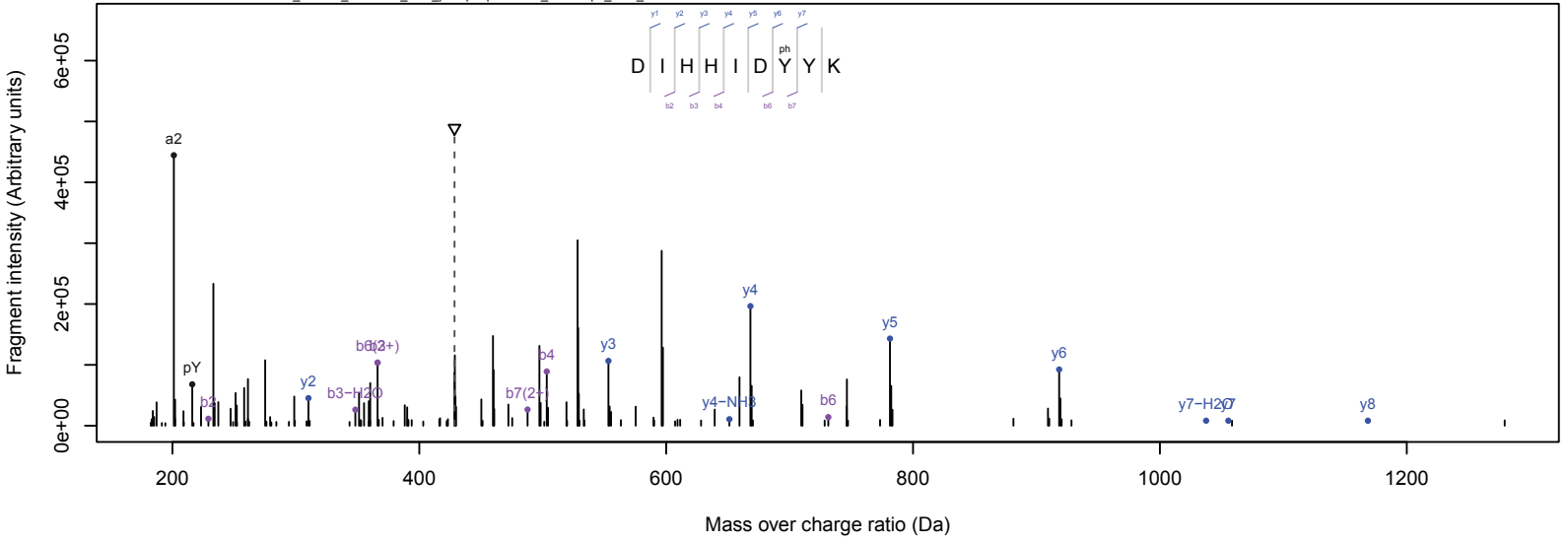
Andromeda score : 77.575



DIHHIDY(ph)YK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10213

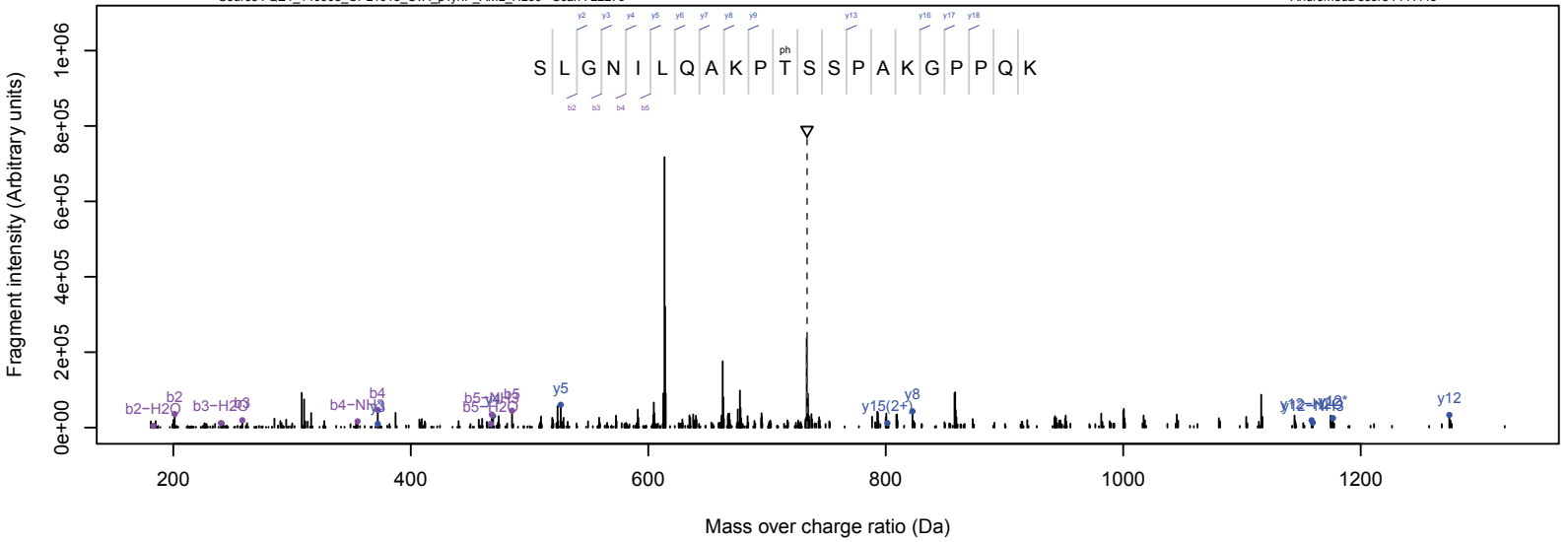
Andromeda score : 77.533



SLGNILQAKPT(ph)SSPAKGGPPQK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 22279

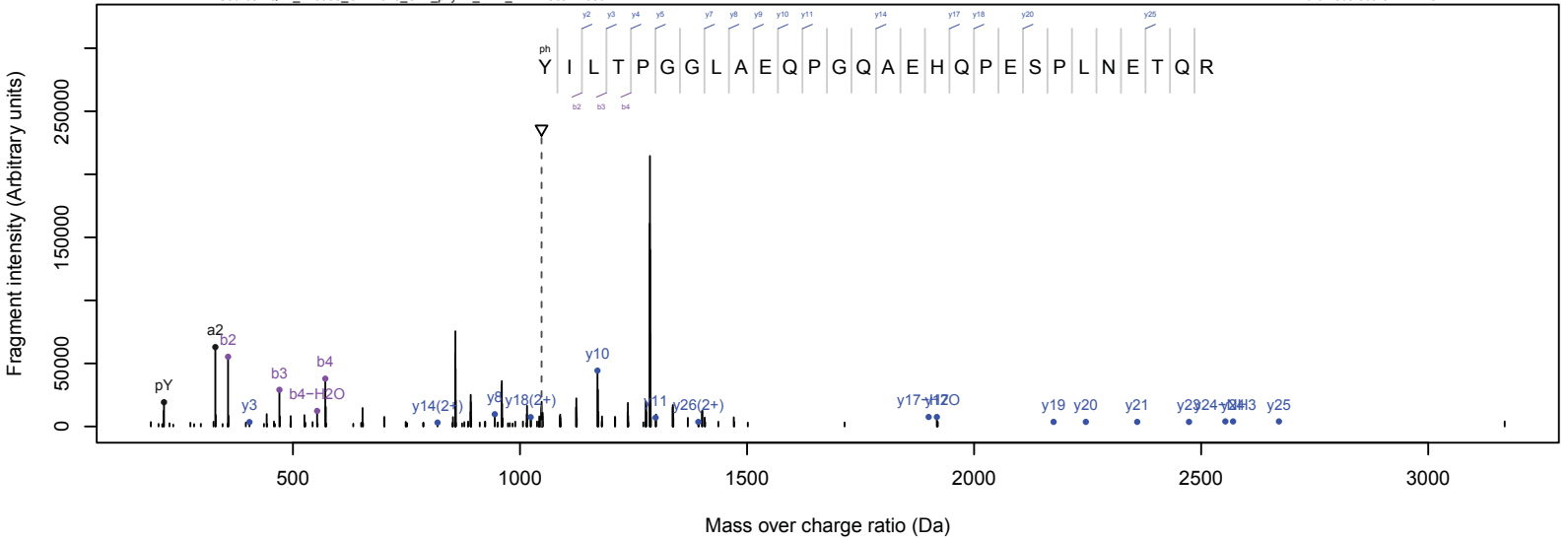
Andromeda score : 77.445



Y(ph)ILTPGLAEQPGQAEHQPEsplNETQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 33911

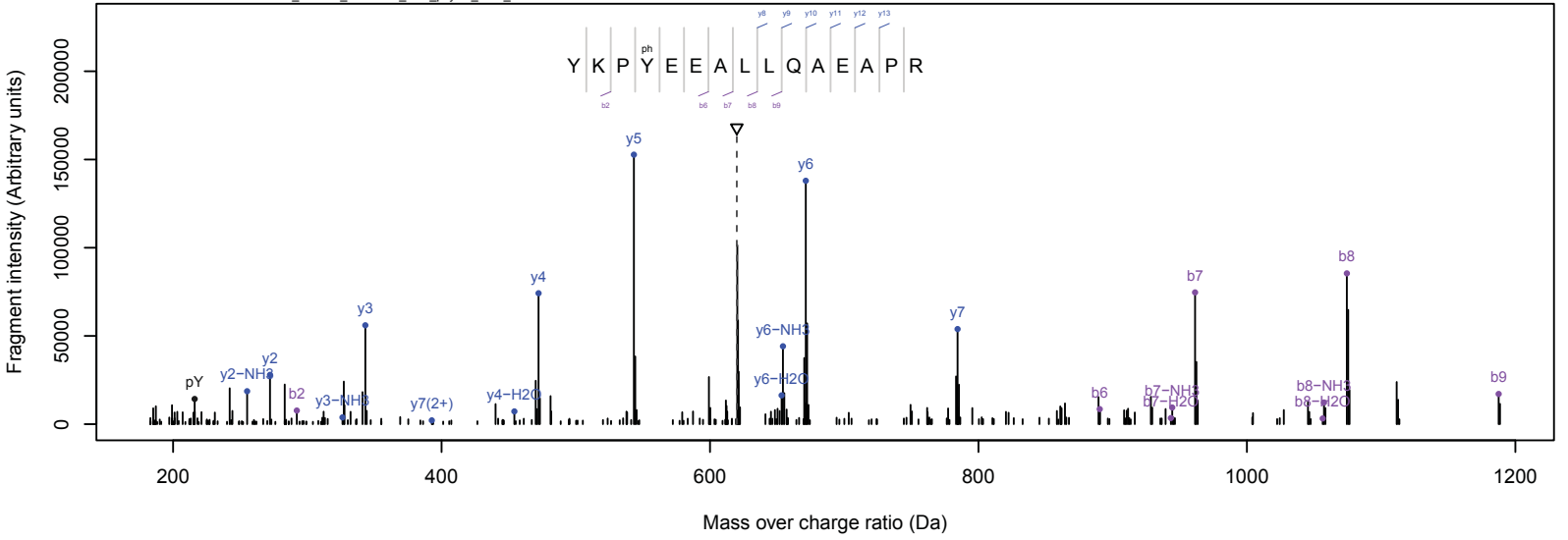
Andromeda score : 77.445



YKPY(ph)EEALLQAEAPR-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan: 26237

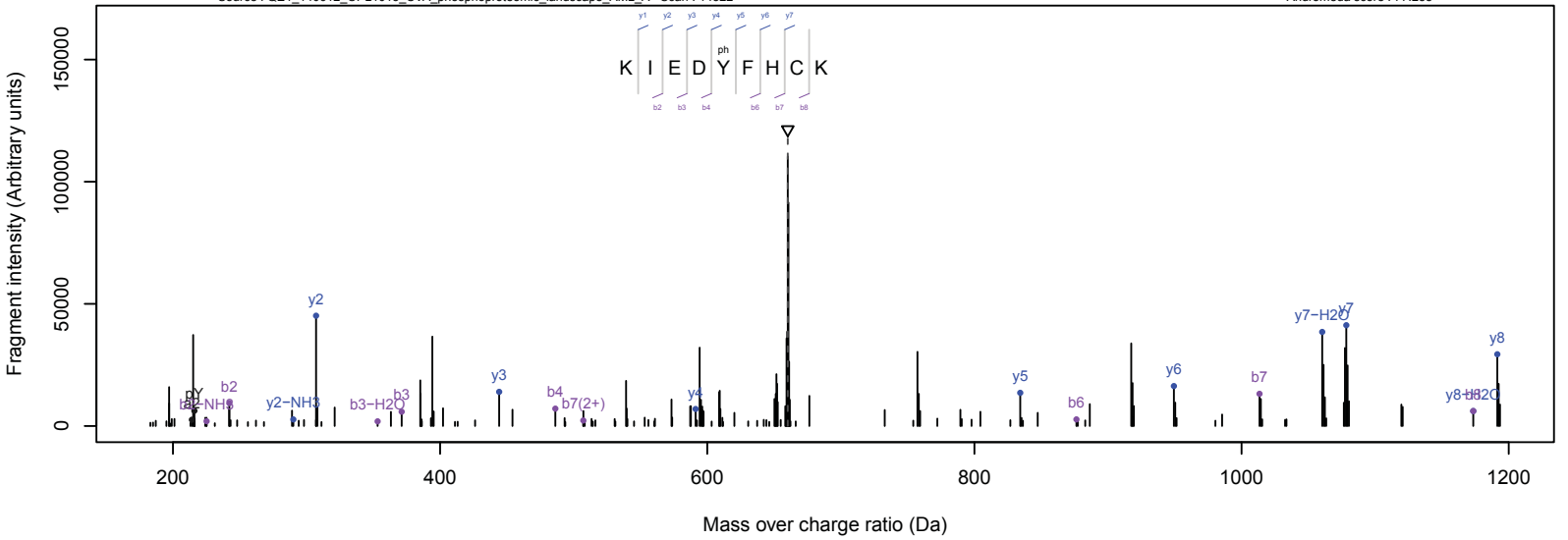
Andromeda score: 77.42



KIEDY(ph)FHCK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 14322

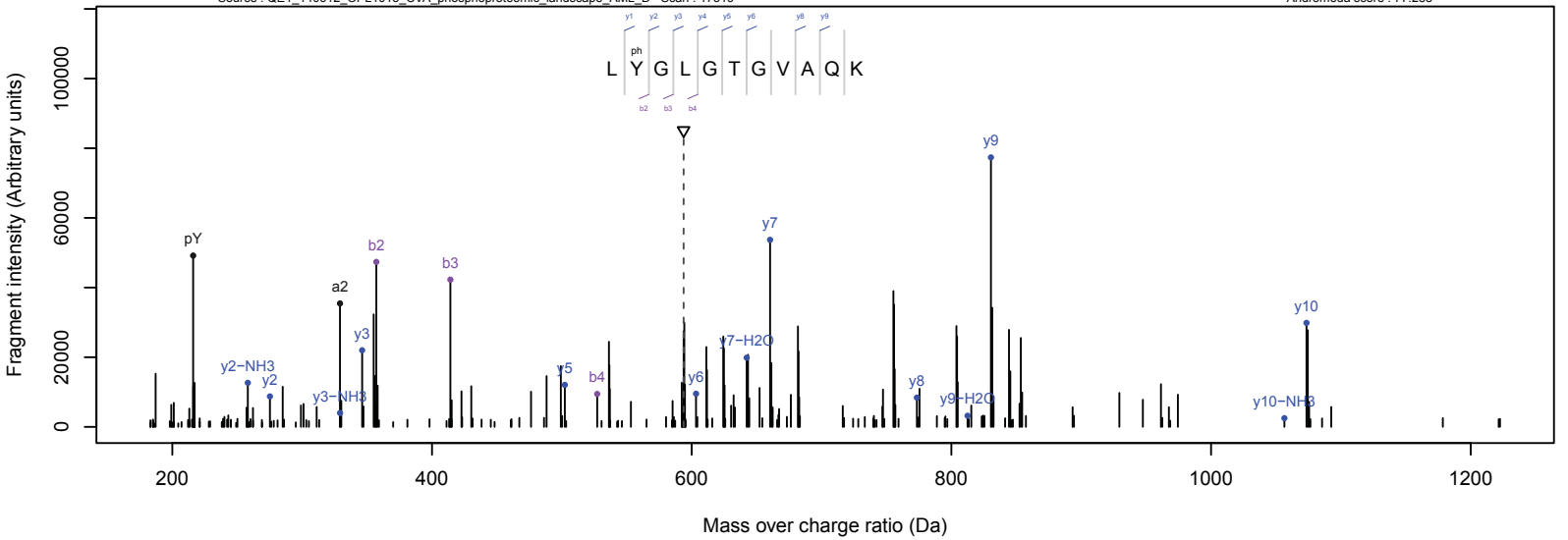
Andromeda score: 77.288



LY(ph)GLGTGVAQK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 17519

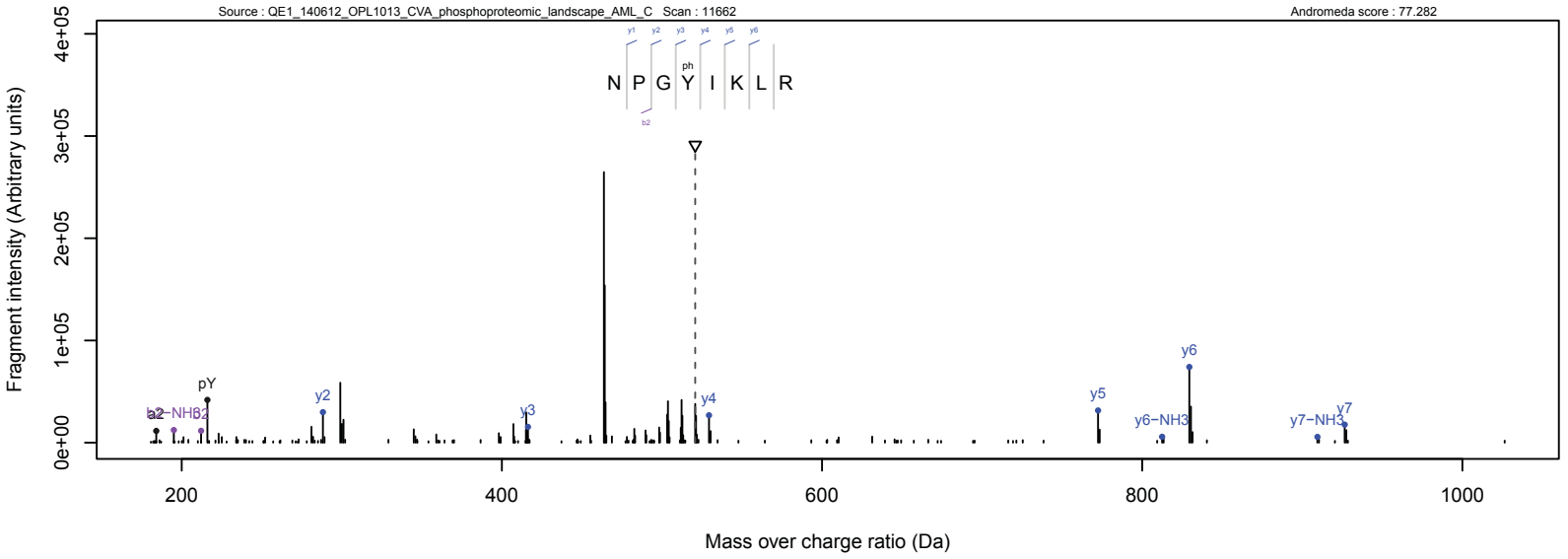
Andromeda score: 77.288



**NPGY(ph)IKLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11662

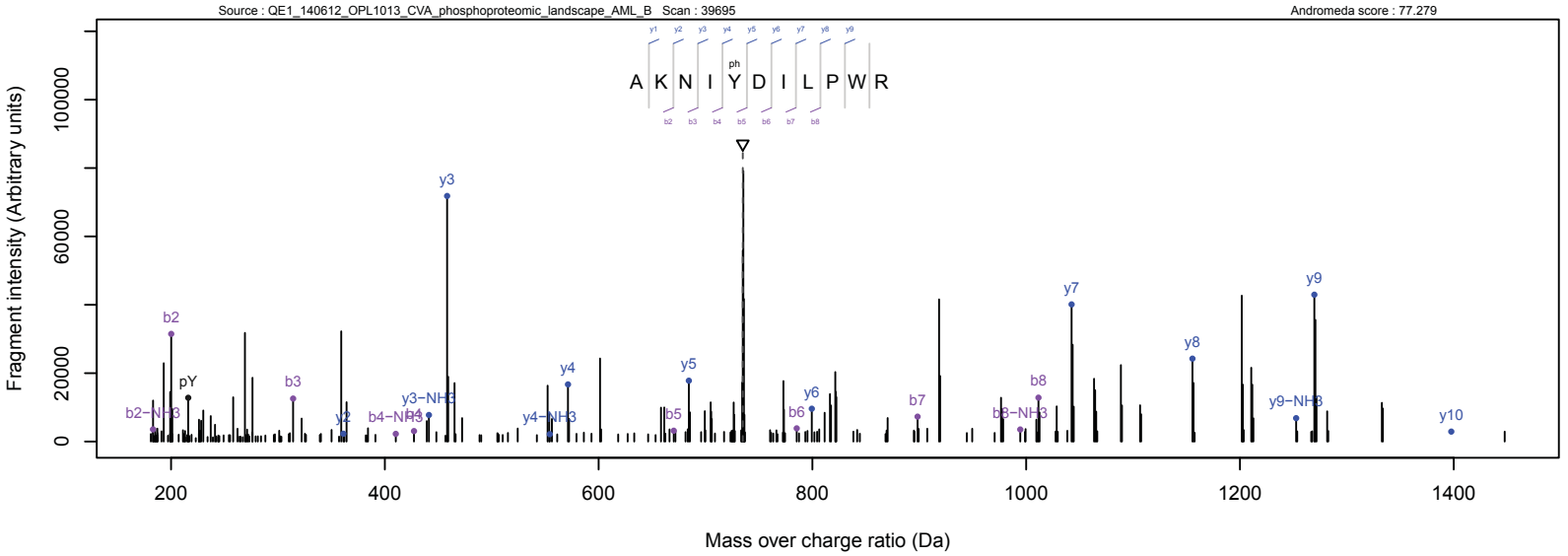
Andromeda score : 77.282



**AKNIY(ph)DILPWR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39695

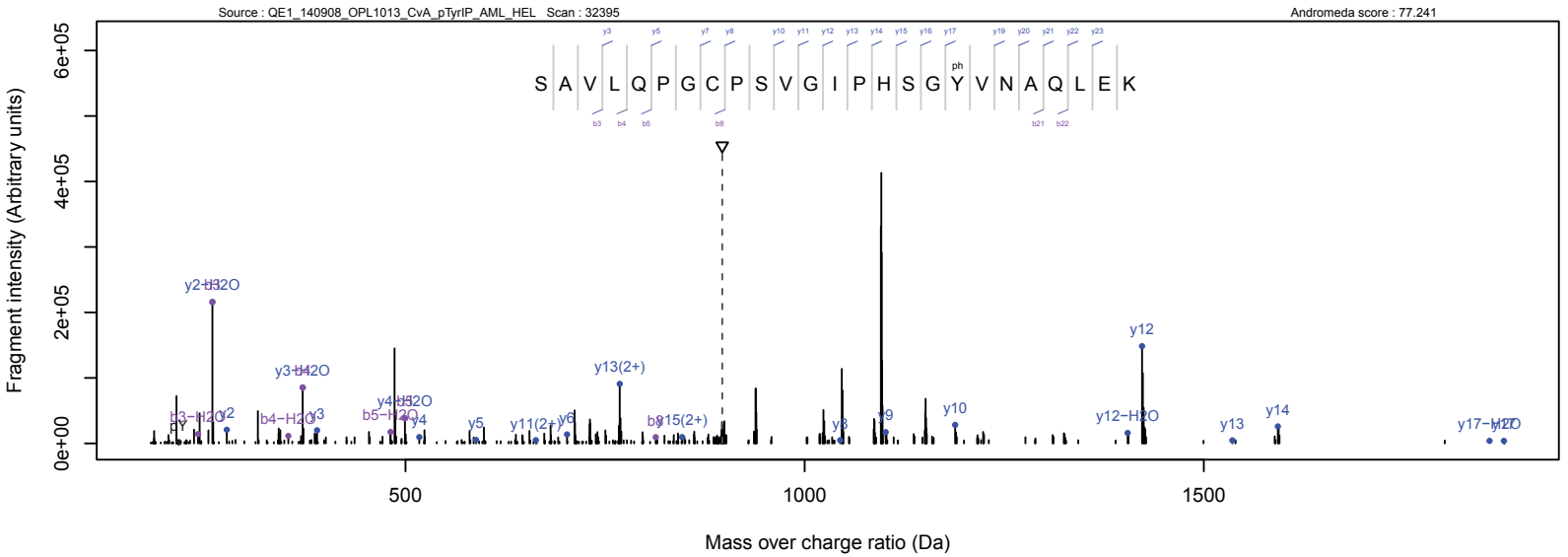
Andromeda score : 77.279



**SAVLQPGCPSVGIHSGY(ph)VNAQLEK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32395

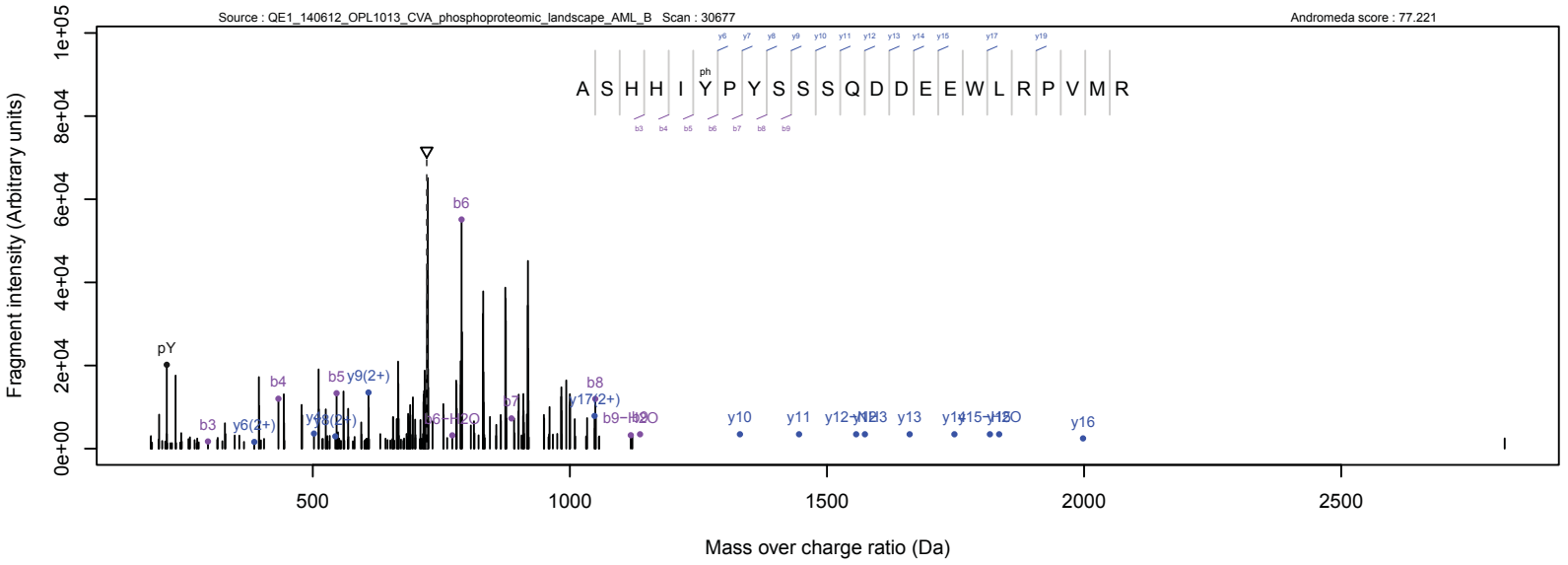
Andromeda score : 77.241



ASHHIY(ph)PYSSQDDEEWRPVMR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 30677

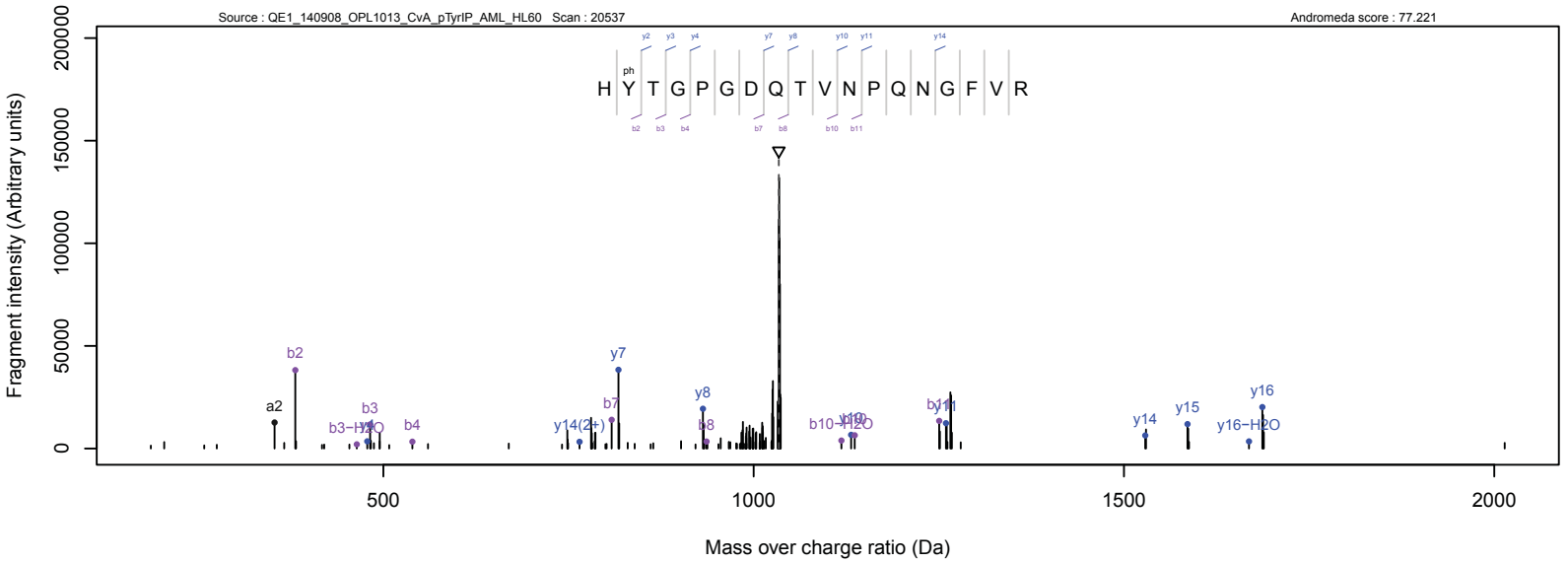
Andromeda score: 77.221



HY(ph)TGPGDQTVNPNQNGFVR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan: 20537

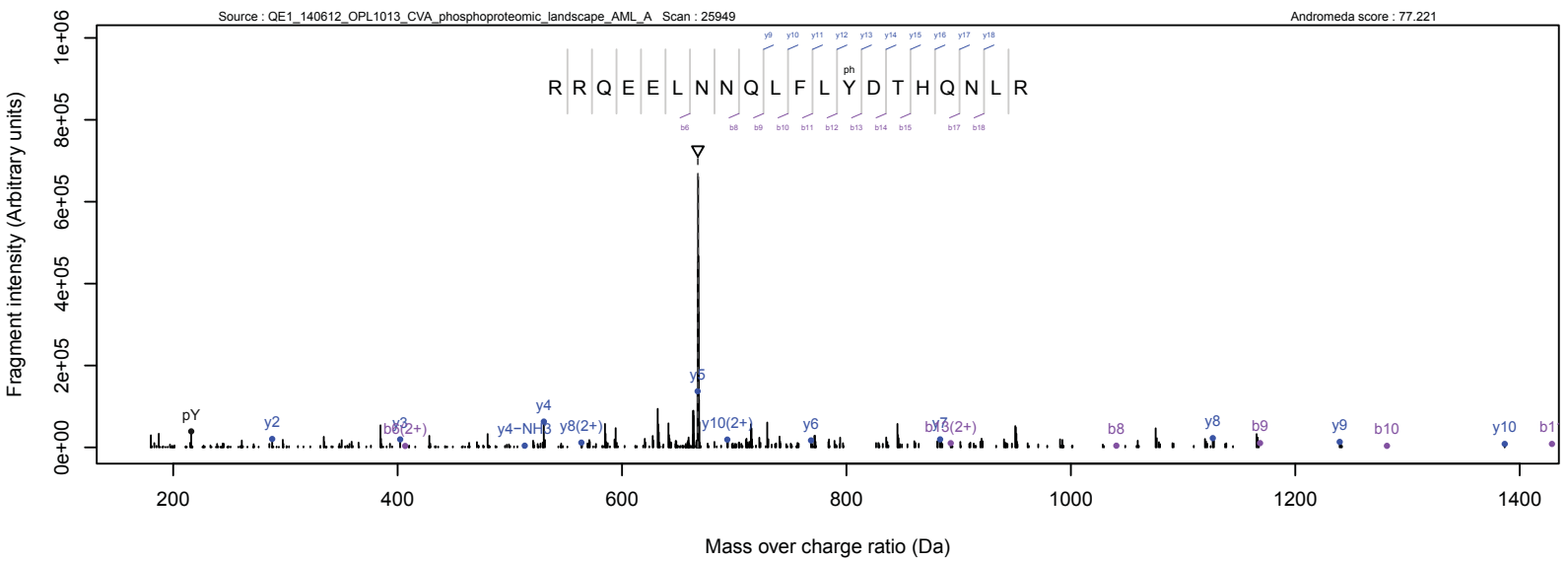
Andromeda score: 77.221



RRQEELNNQLFLYDTHQNLRL-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 25949

Andromeda score: 77.221

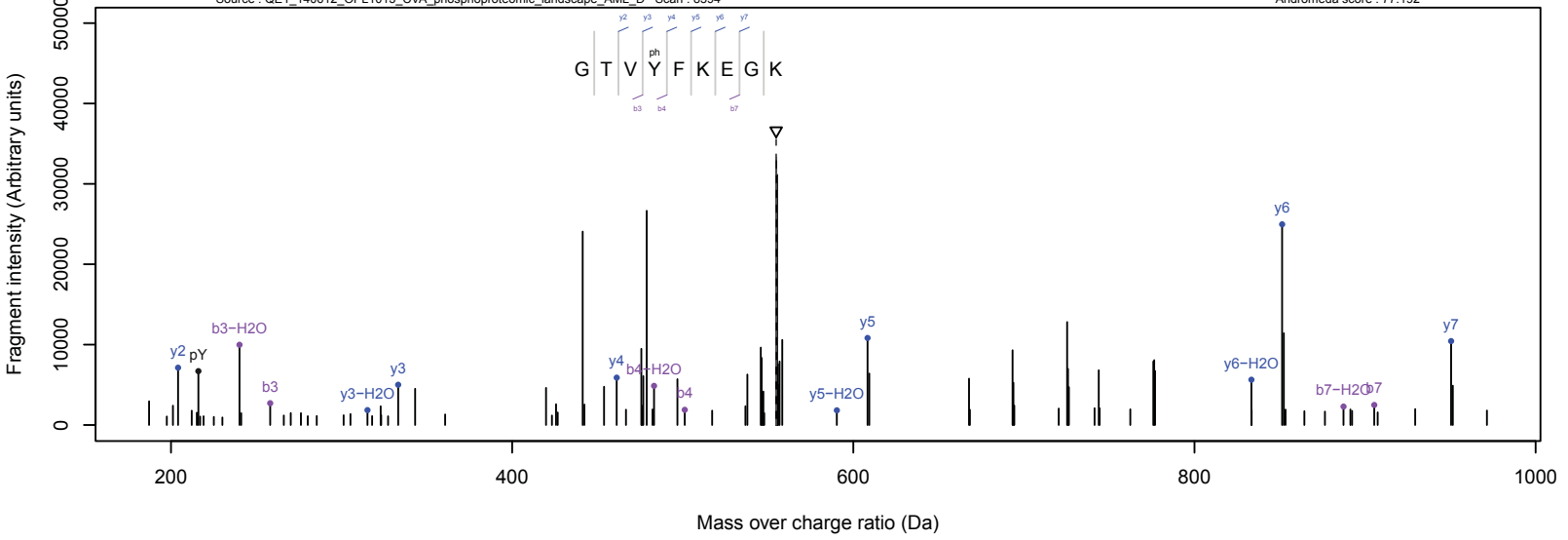




GTVY(ph)FKEGK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 8594

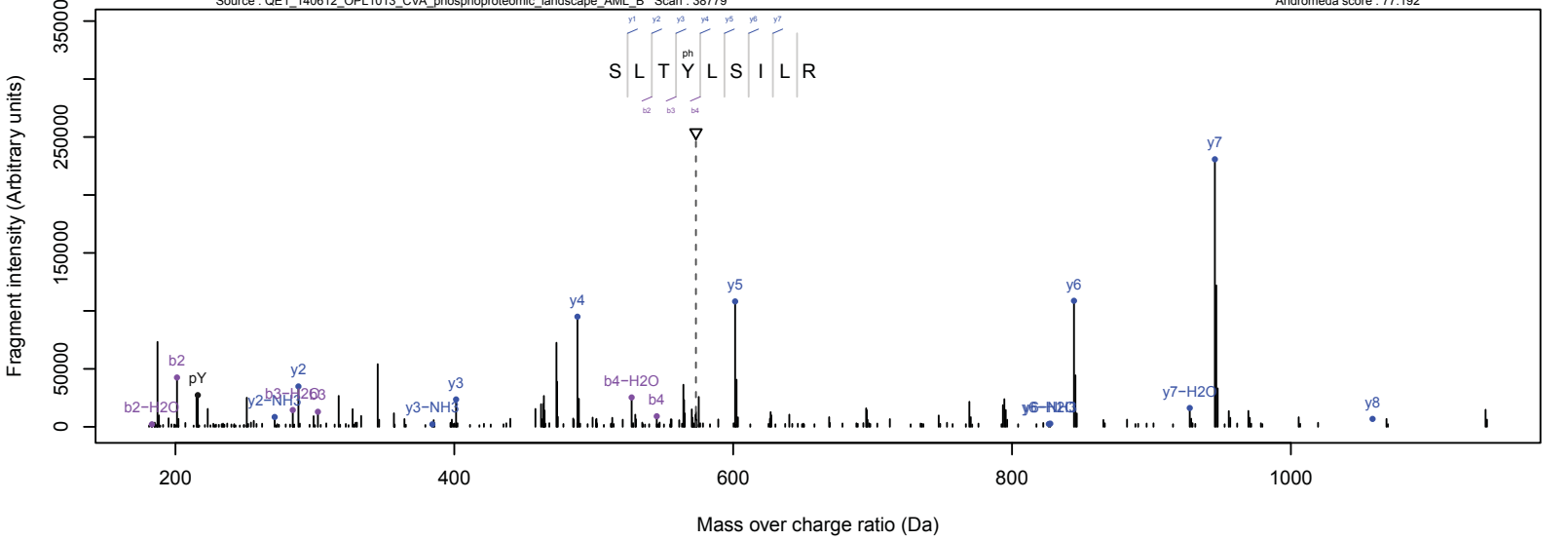
Andromeda score: 77.192



SLTY(ph)LSILR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 38779

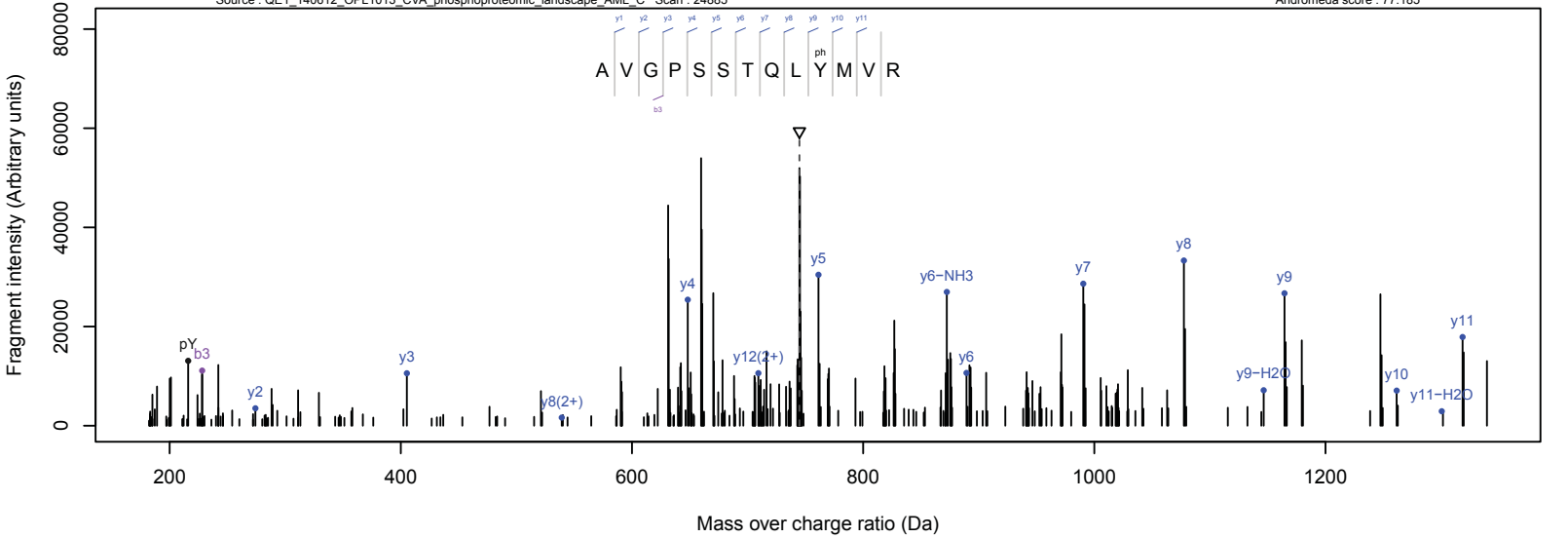
Andromeda score: 77.192



AVGSPSTQLY(ph)MVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24885

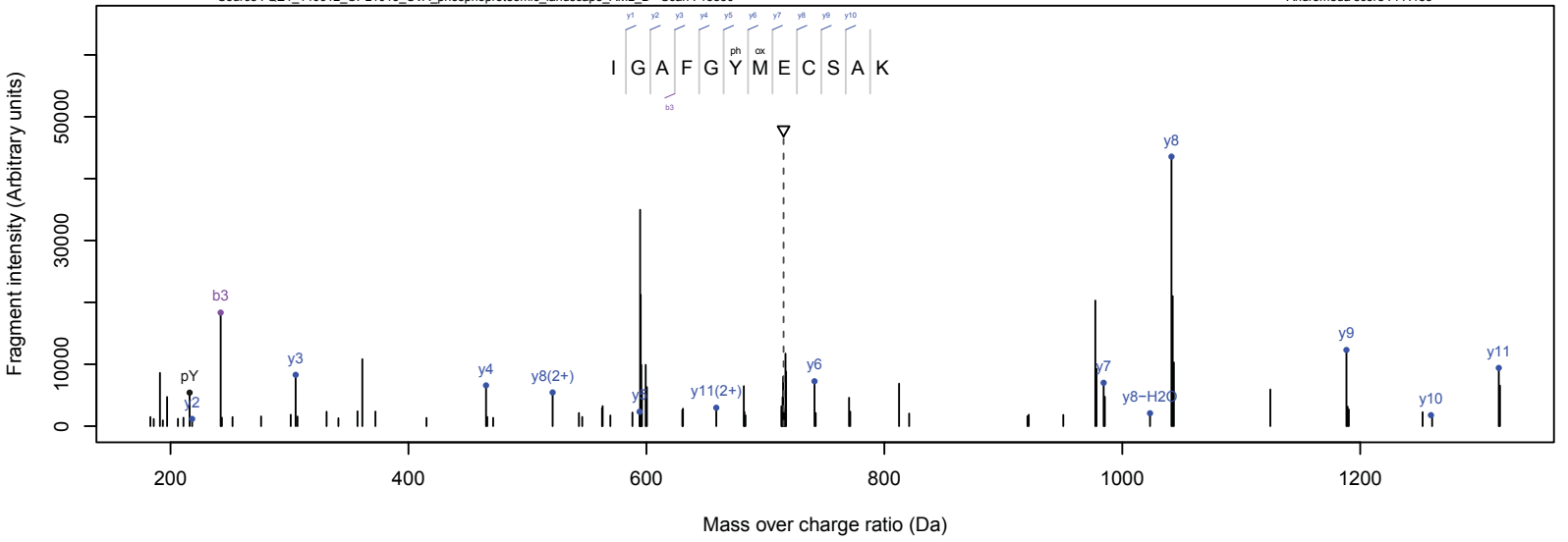
Andromeda score: 77.185



IGAFGY(ph)M(ox)ECSAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 15850

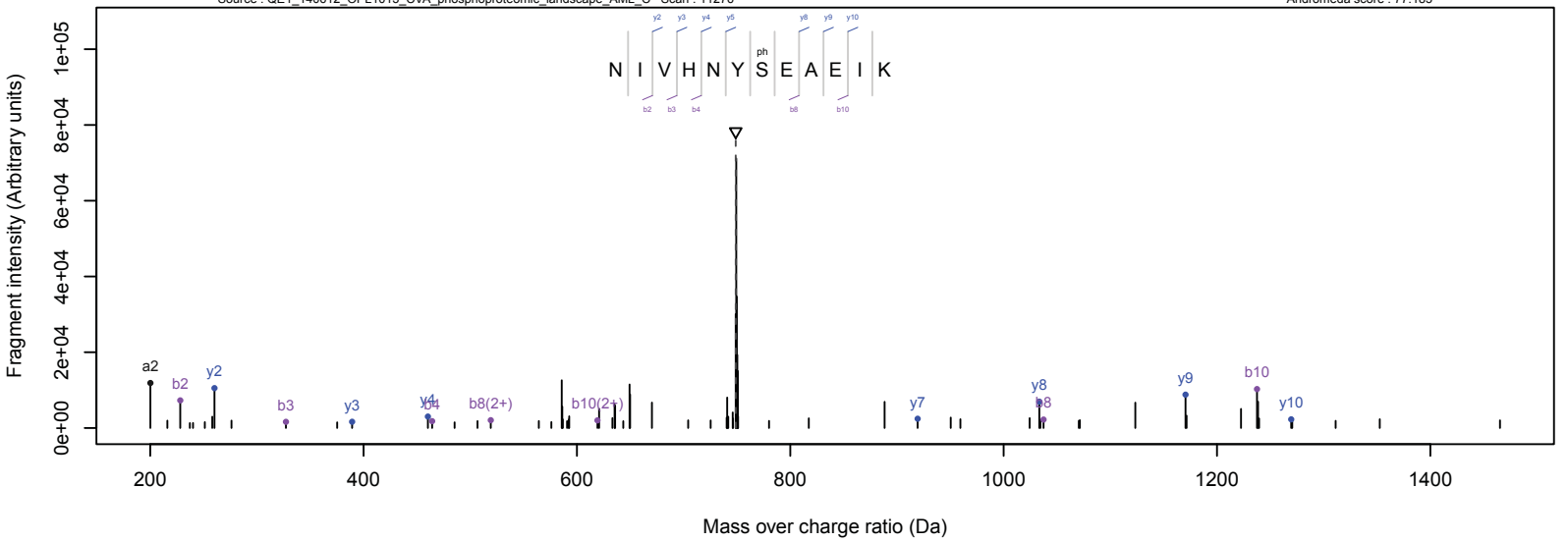
Andromeda score : 77.185



NIVHNYS(ph)EAEIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11276

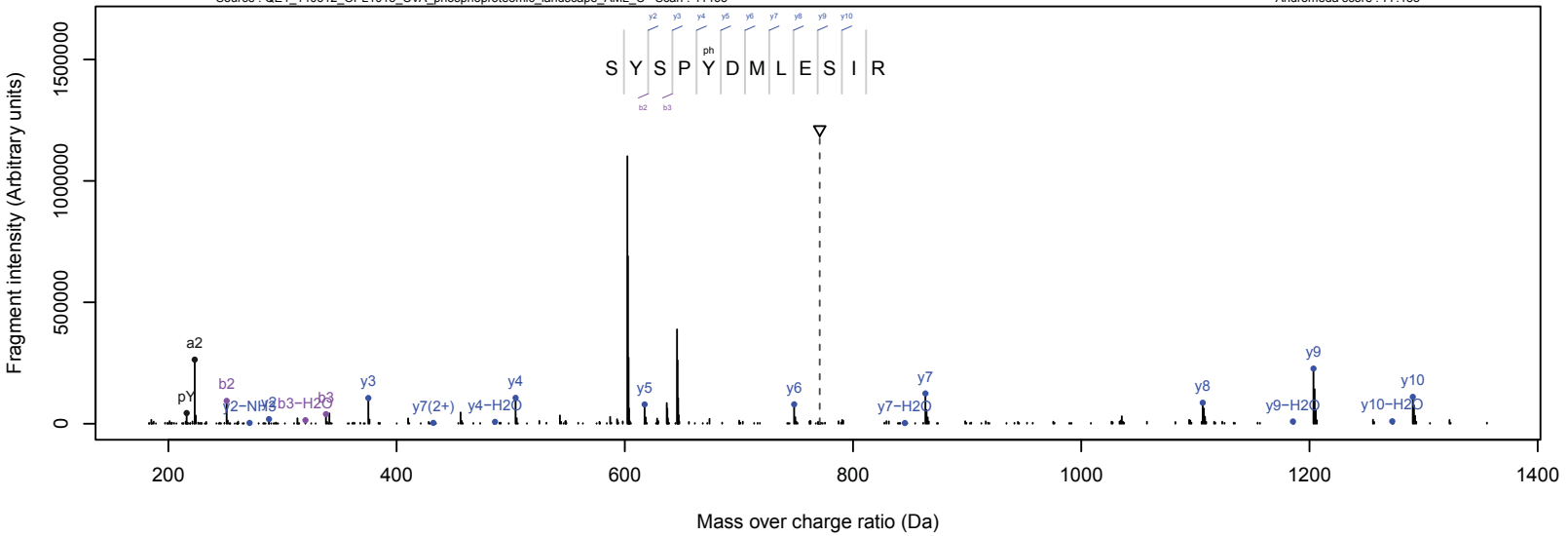
Andromeda score : 77.185



SYSPY(ph)DMLESIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41465

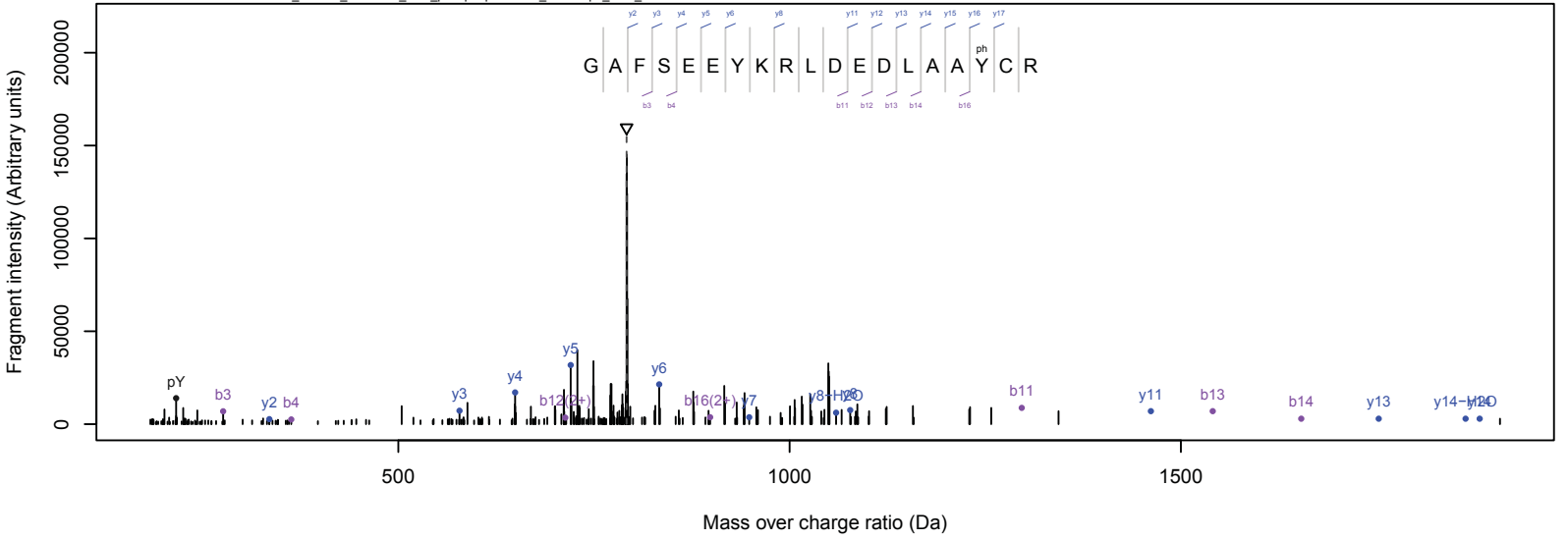
Andromeda score : 77.185



**GAFSEEYKRLDEDLAAY(ph)CR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32841

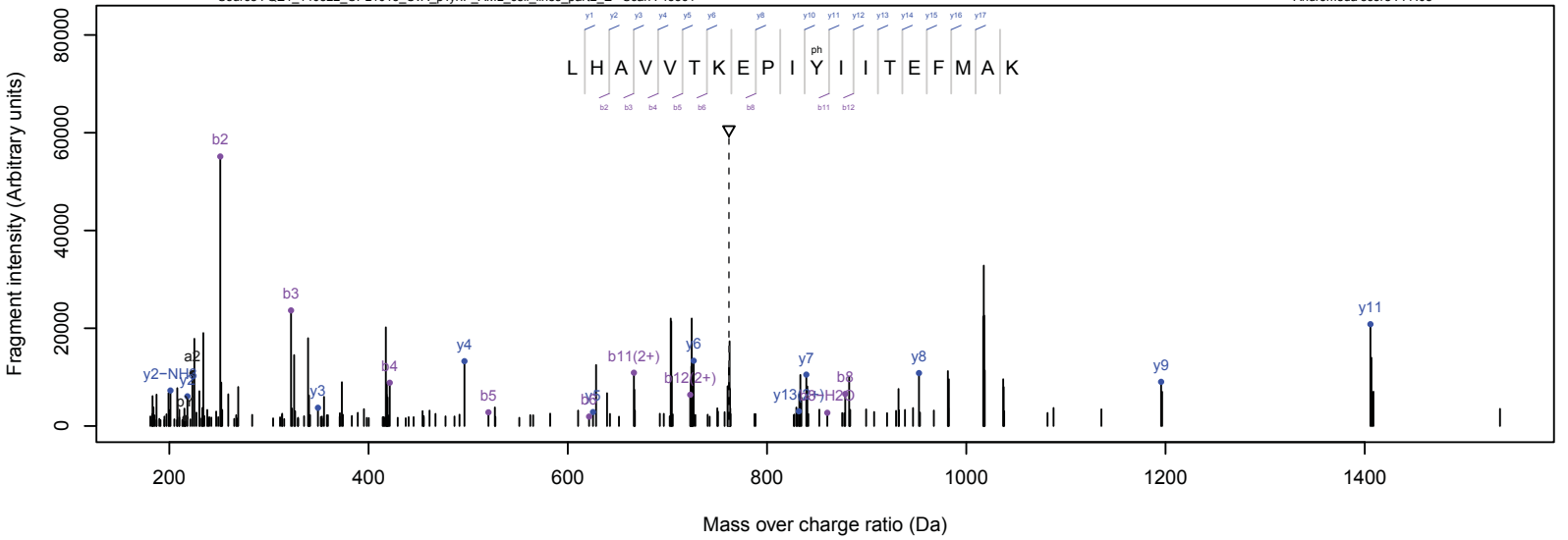
Andromeda score : 77.08



**LHAVVTKEPIY(ph)ITEFMAK-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 45561

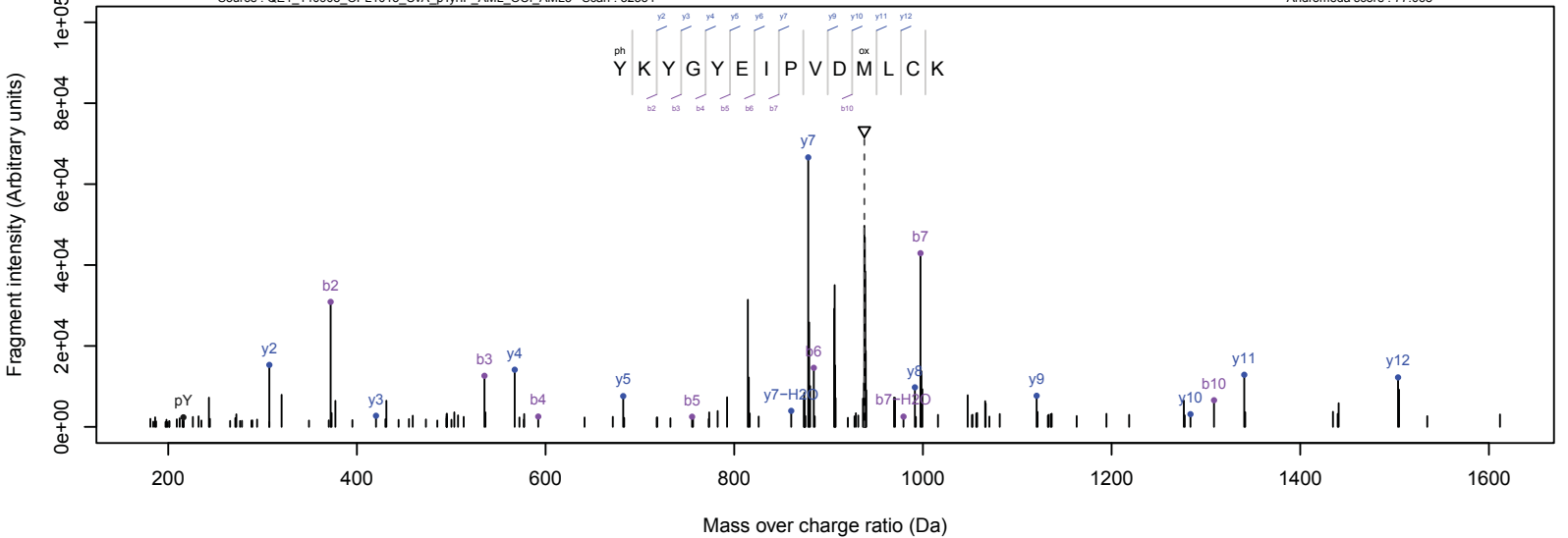
Andromeda score : 77.08



**Y(ph)KYGYEIPVDM(ox)LCK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 32354

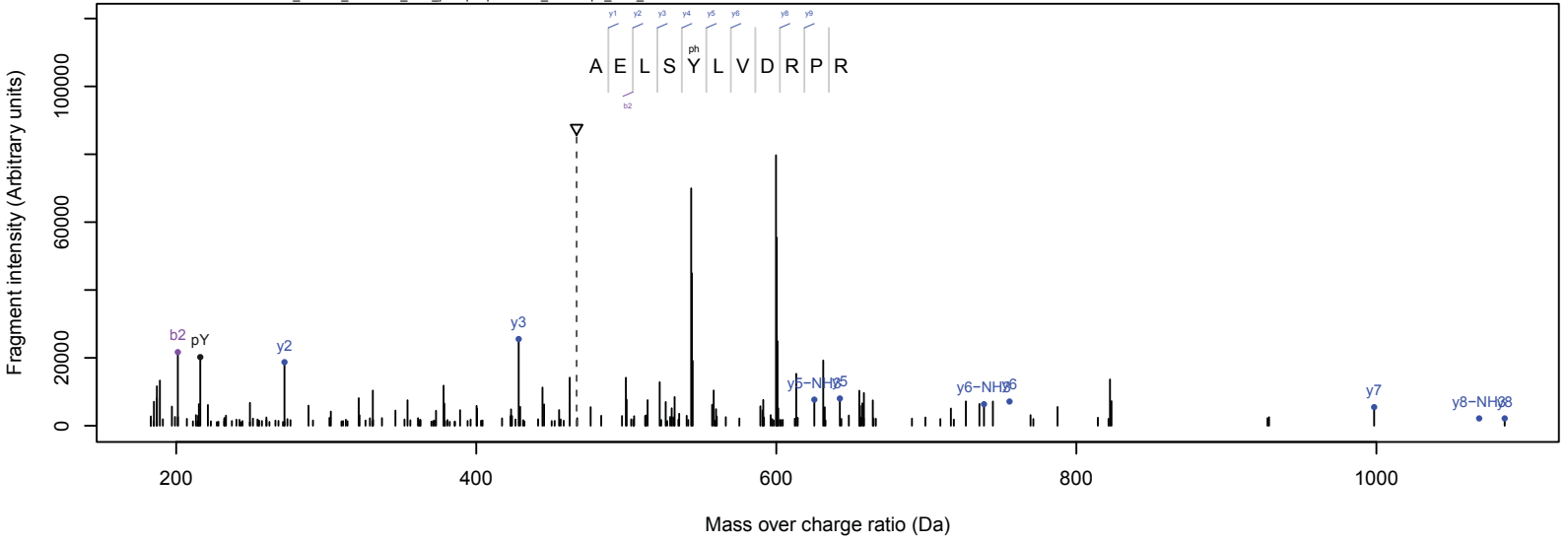
Andromeda score : 77.068



**AELSY(ph)LVDRPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23563

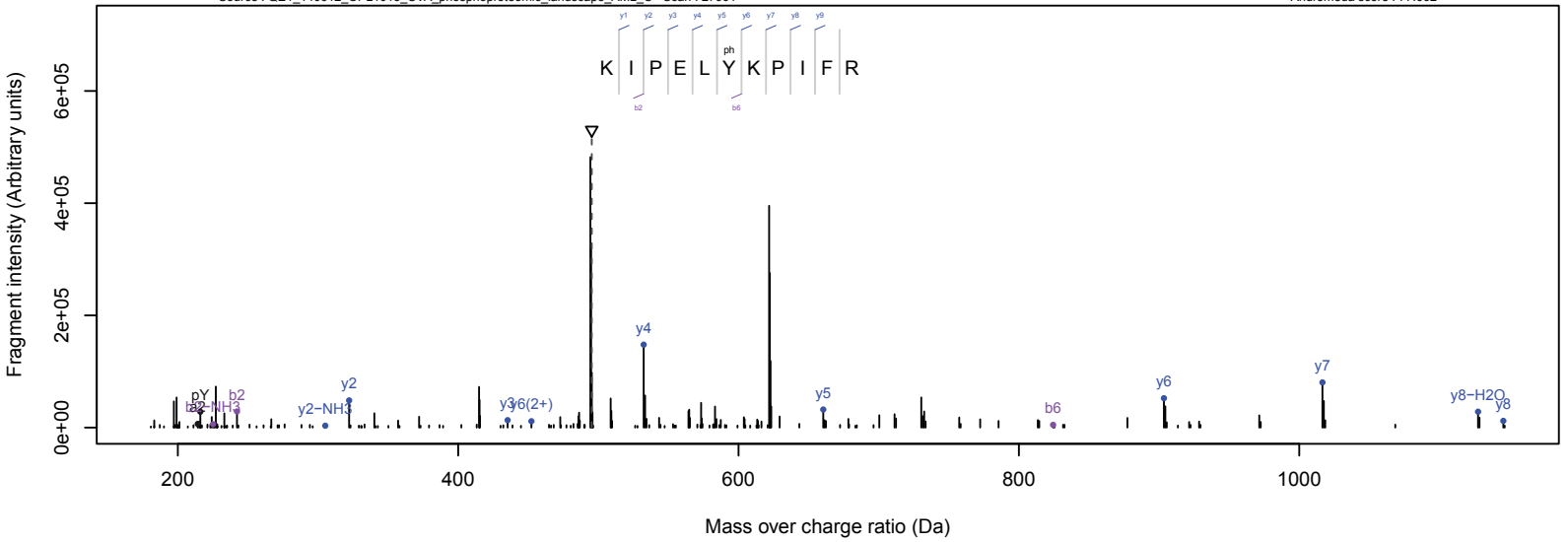
Andromeda score : 77.062



**KIPELY(ph)KPIFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27901

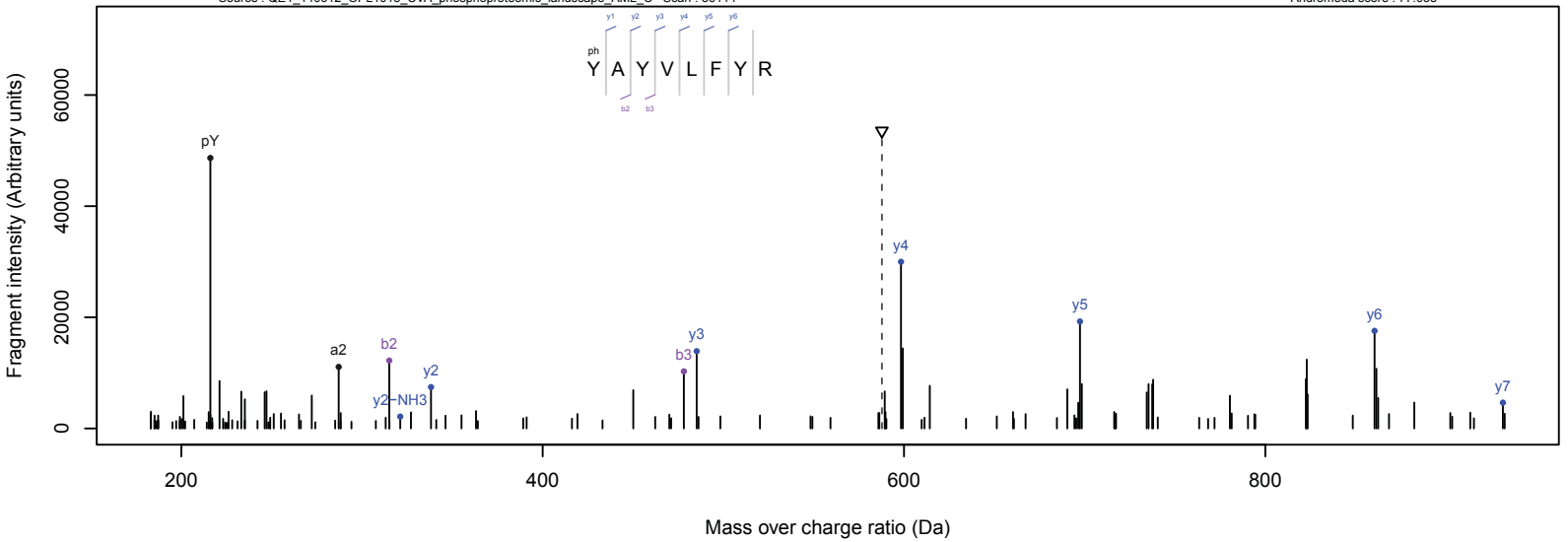
Andromeda score : 77.062



**Y(ph)AYLVFYR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36114

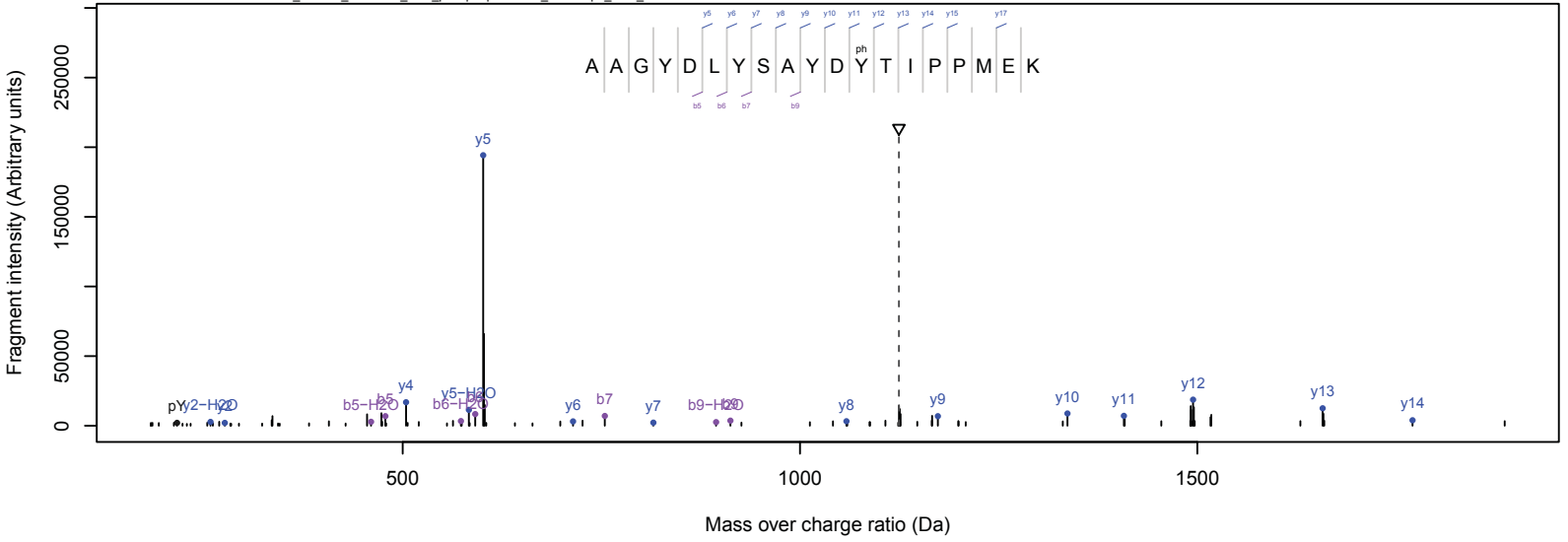
Andromeda score : 77.058



**AAGYDLYSAYDY(ph)TIPPMEK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41608

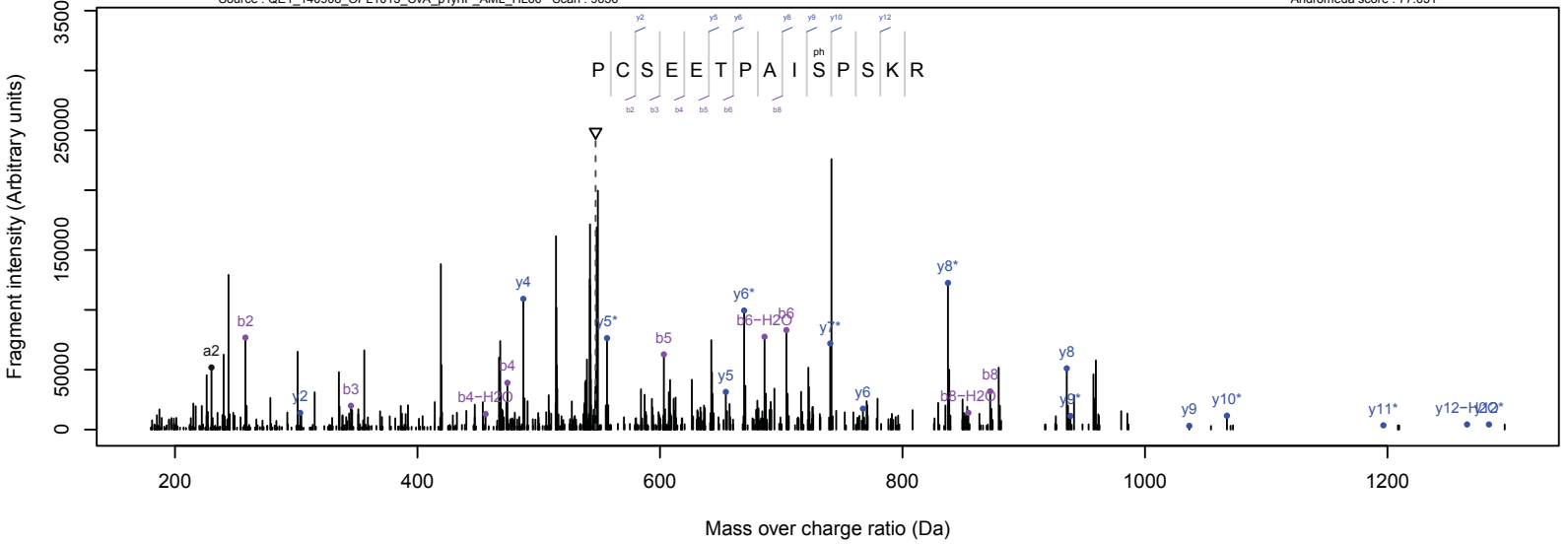
Andromeda score : 77.031



**PCSEETPAIS(ph)PSKR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 9636

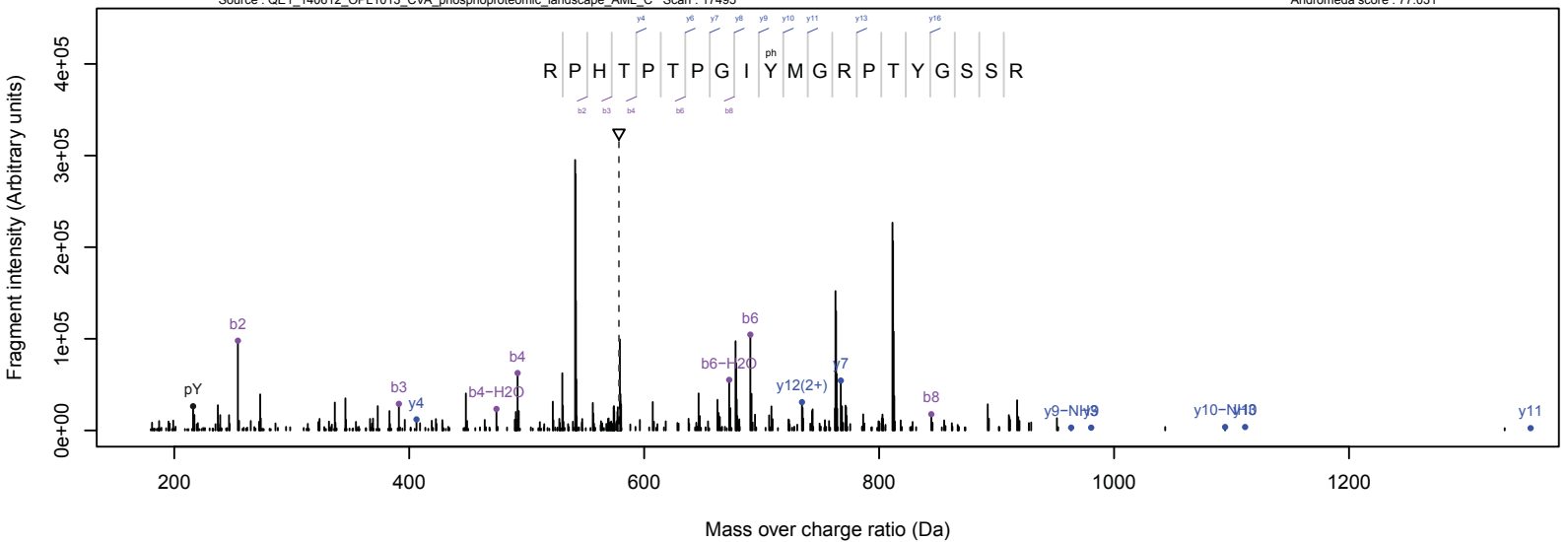
Andromeda score : 77.031



**RPHTPTGIY(ph)MGRPTYGSSR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17495

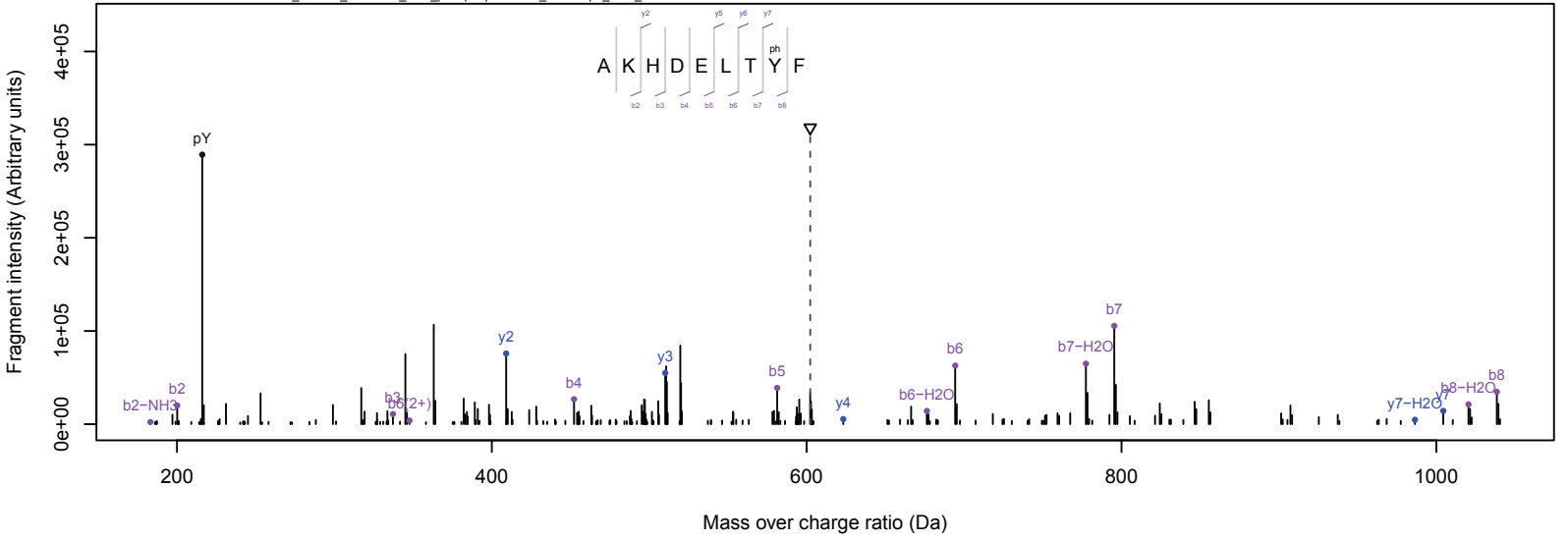
Andromeda score : 77.031



AKHDELTY(ph)F-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19971

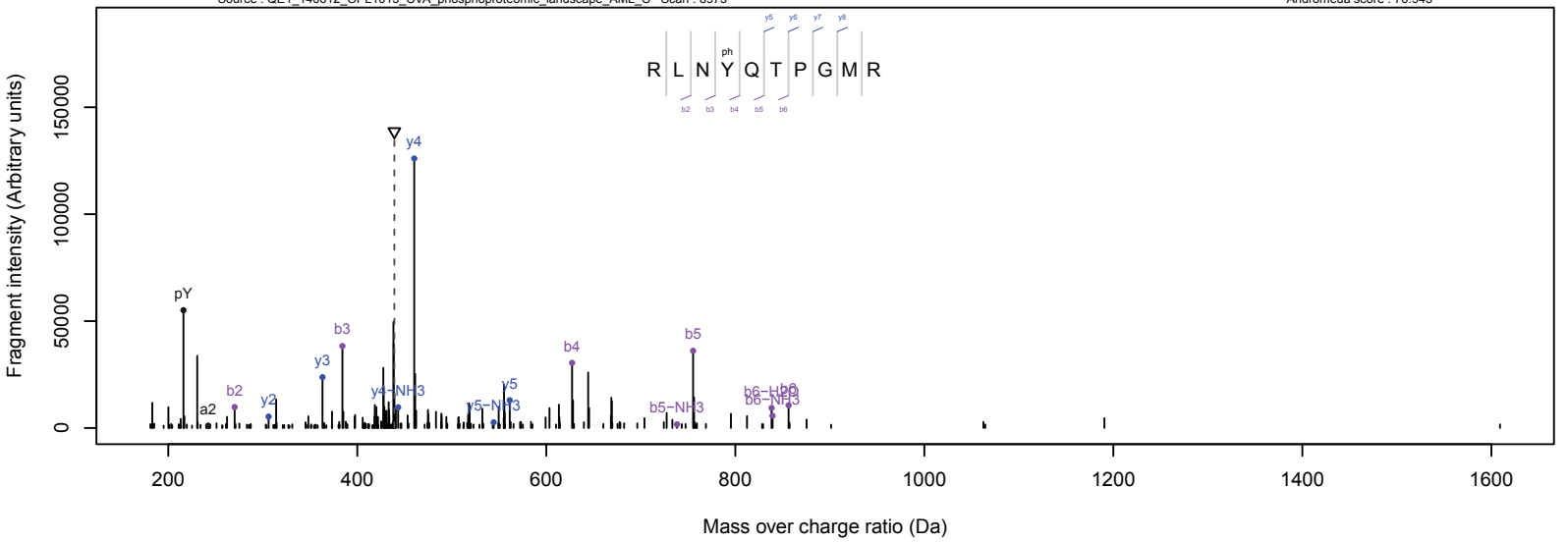
Andromeda score : 76.943



RLNY(ph)QTPGMR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8973

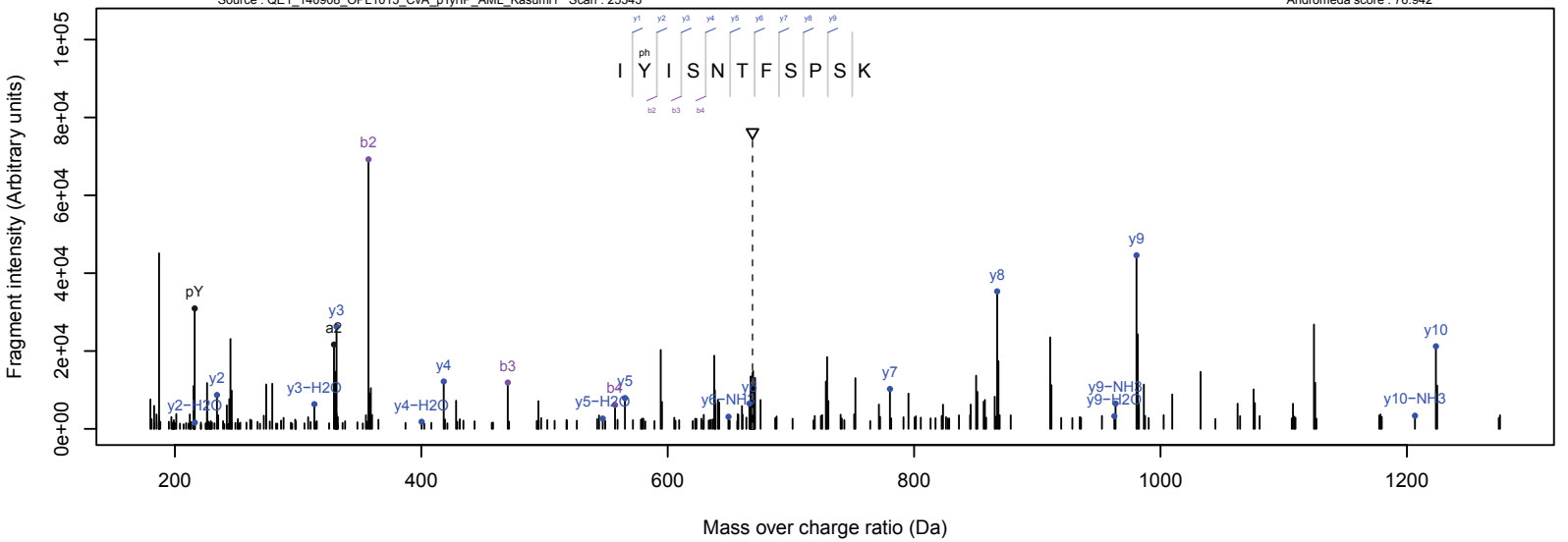
Andromeda score : 76.943



IY(ph)ISNTFSPSK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 25343

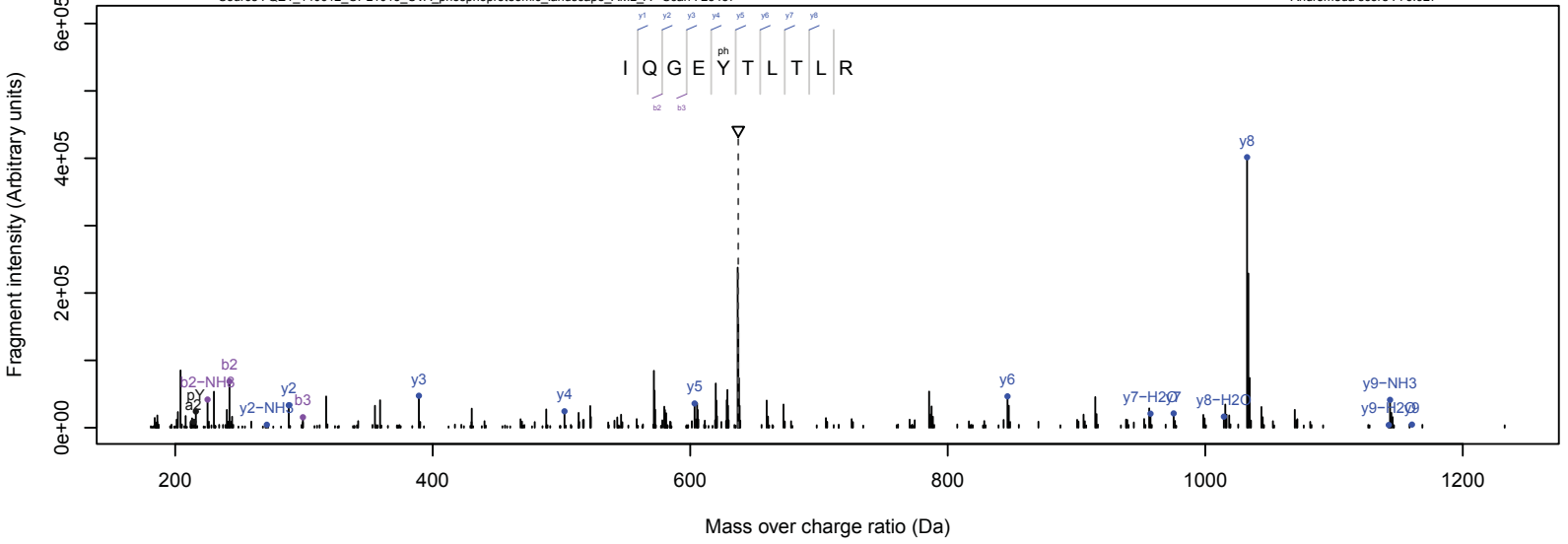
Andromeda score : 76.942



**IQGEY(ph)TLTLR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 23457

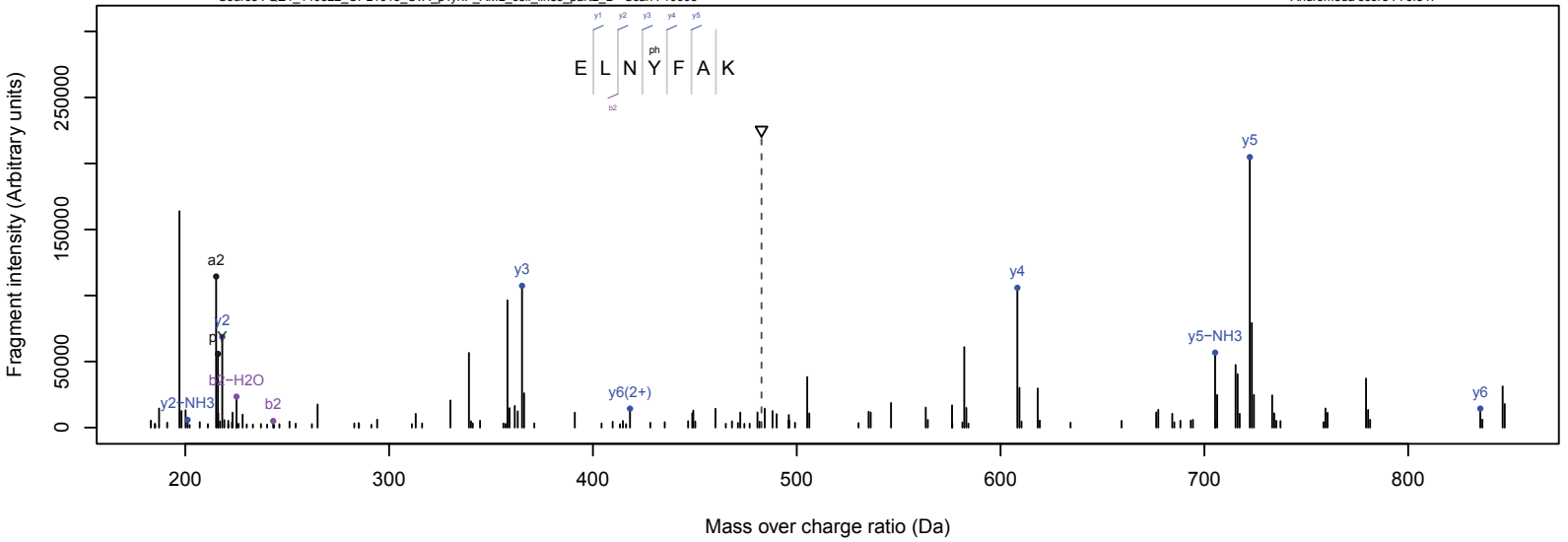
Andromeda score: 76.927



**ELNY(ph)FAK-2+**

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 10608

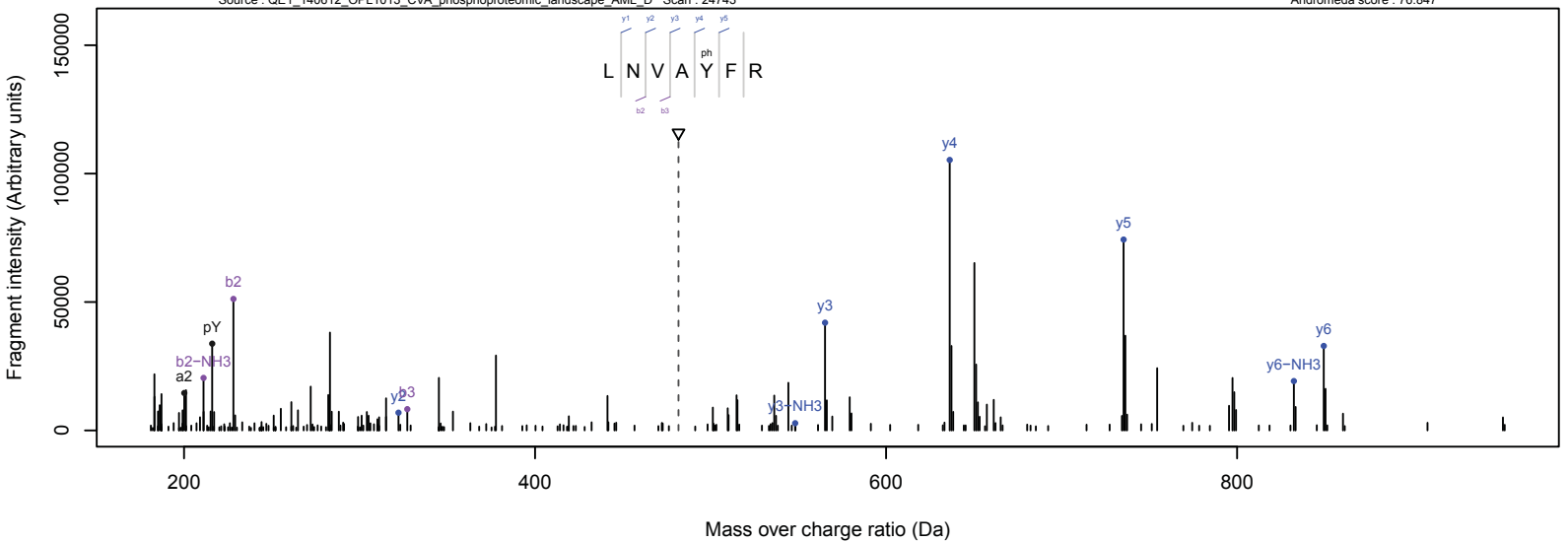
Andromeda score: 76.847



**LNVAY(ph)FR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 24743

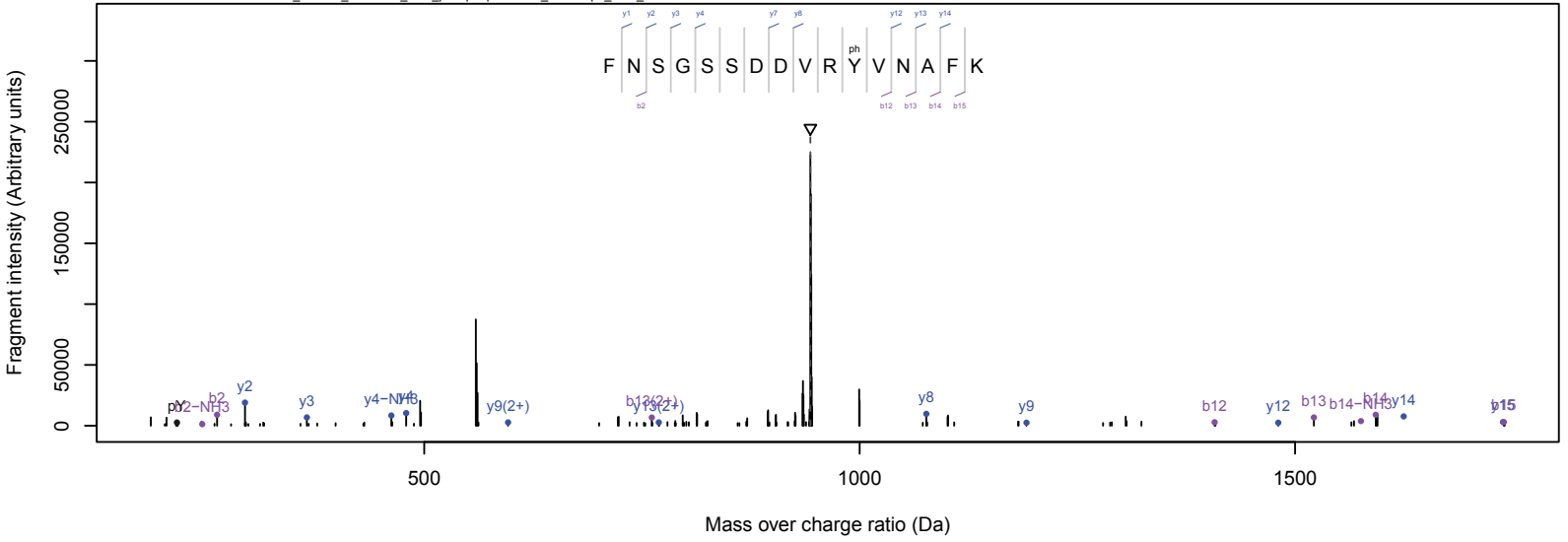
Andromeda score: 76.847



**FNSGSSDDVRY(ph)VNAFK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22620

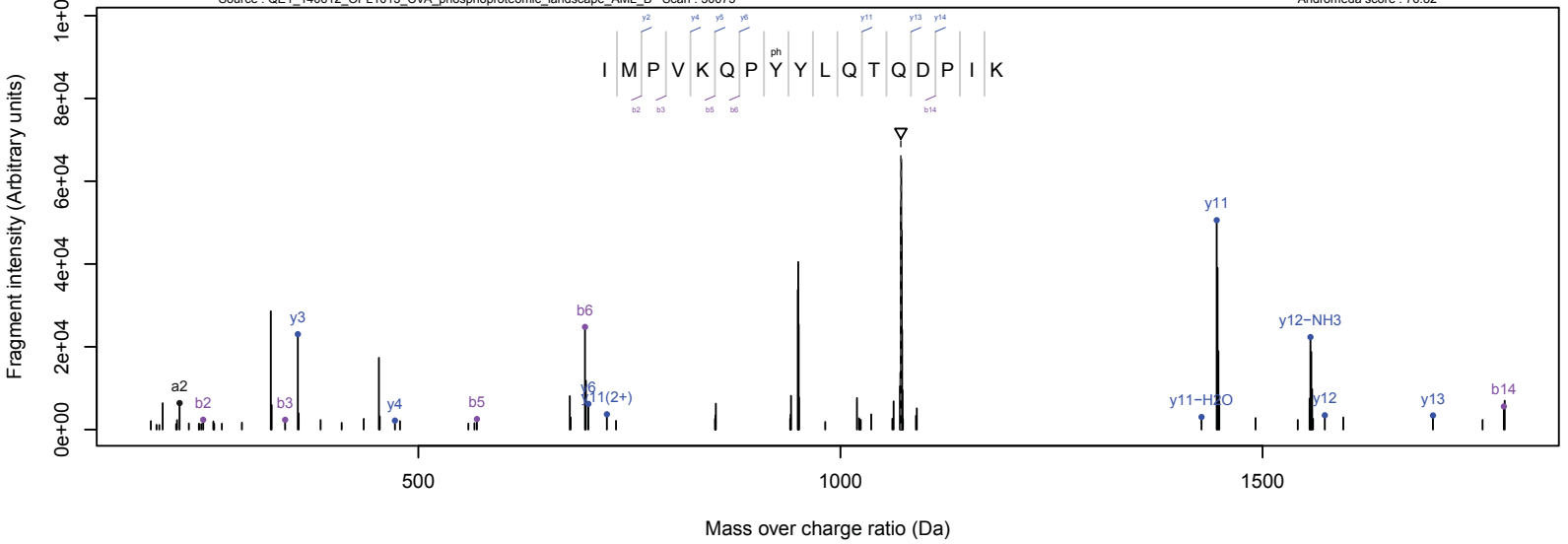
Andromeda score : 76.82



**IMPVKQPY(ph)YLQTQDPK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30073

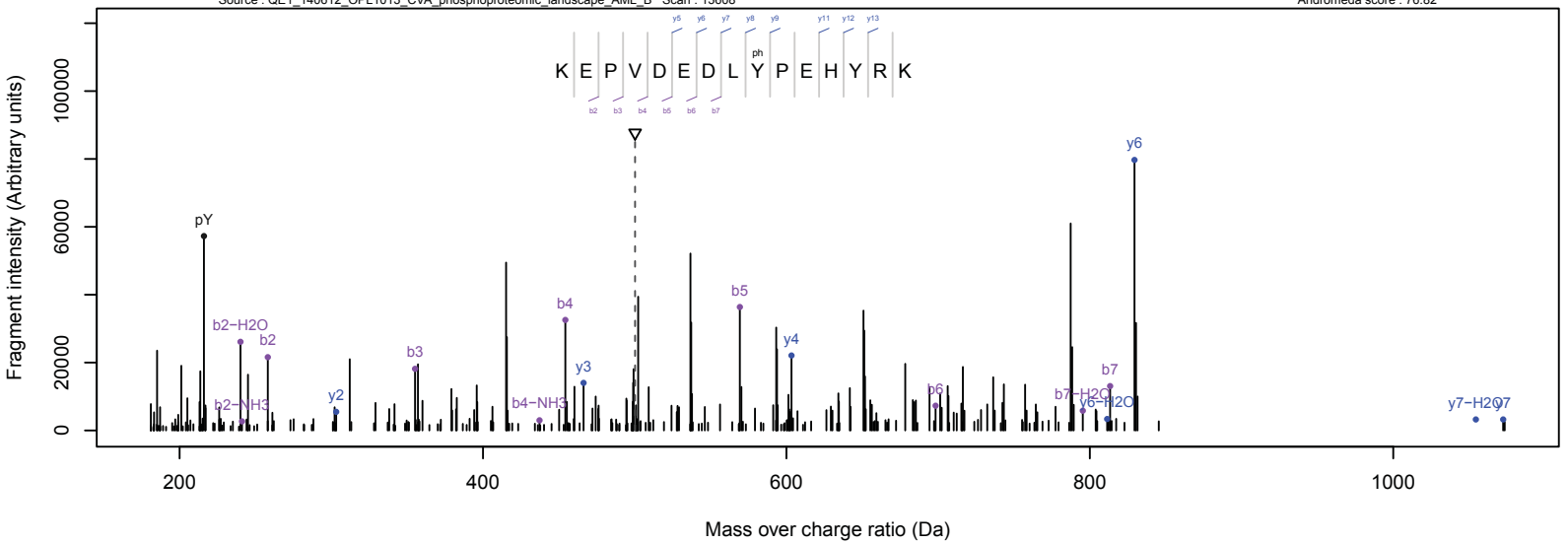
Andromeda score : 76.82



**KEPVDEDLY(ph)PEHYRK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13608

Andromeda score : 76.82

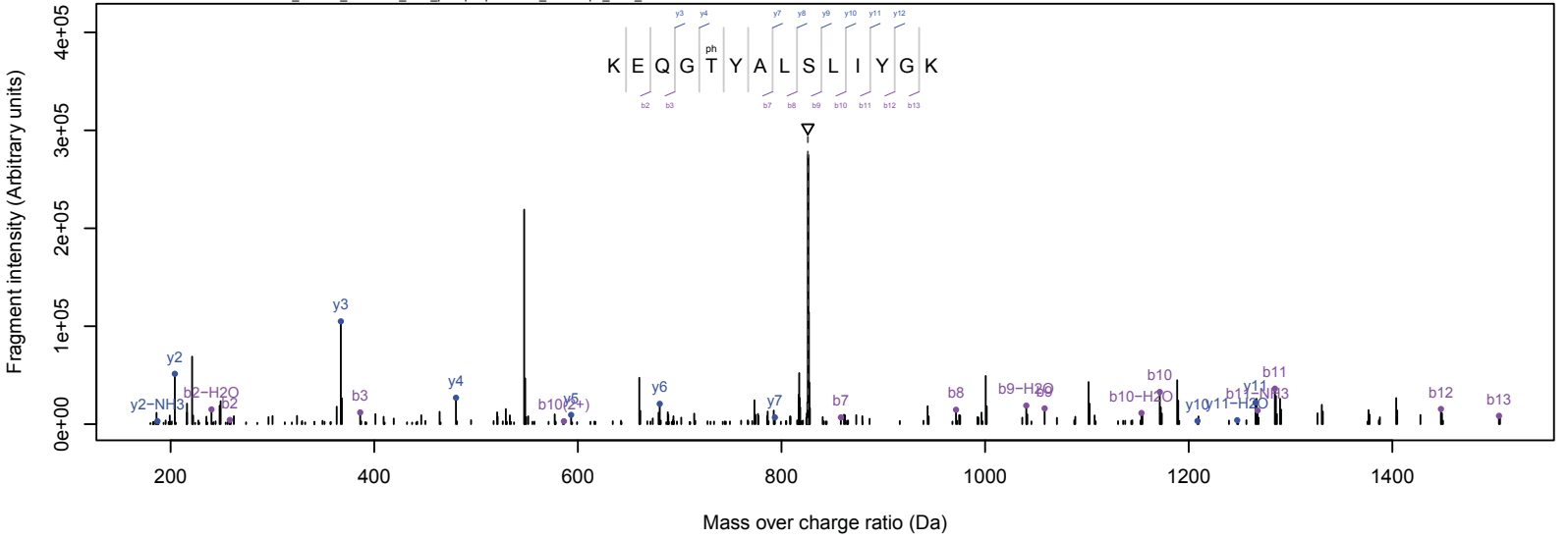




KEQGT(ph)YALSLIYGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31058

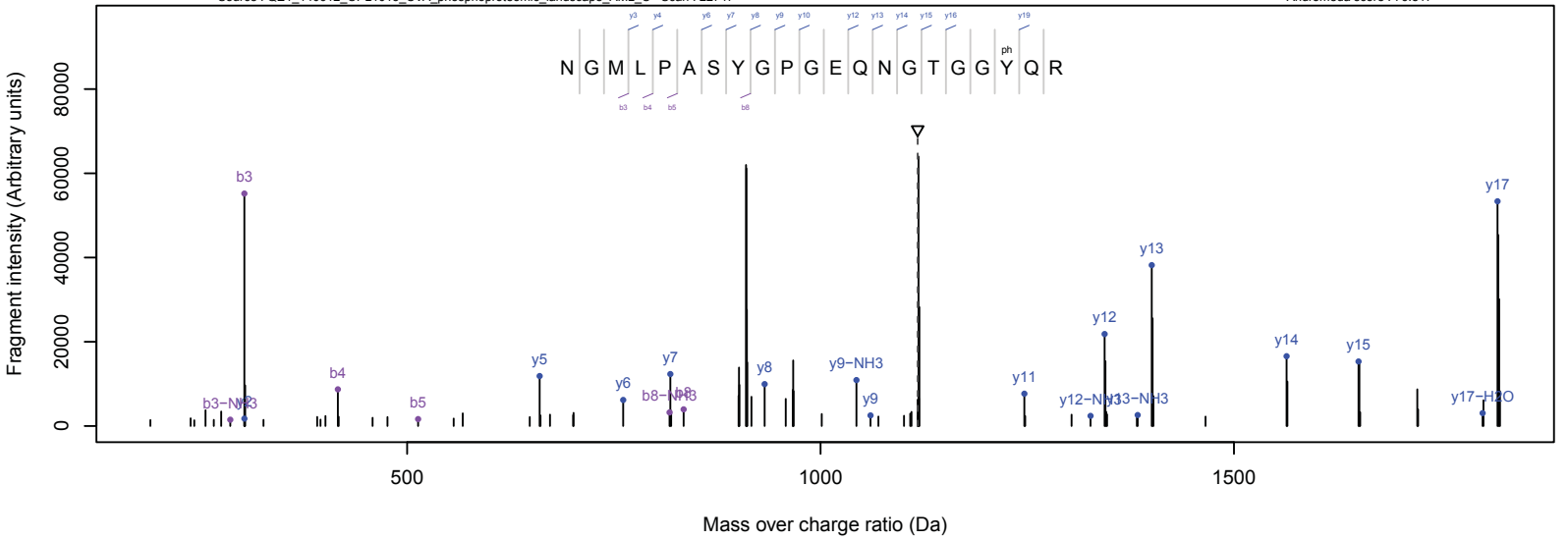
Andromeda score : 76.82



NGMLPASYGPGEQNGTGGY<sup>ph</sup>QR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22747

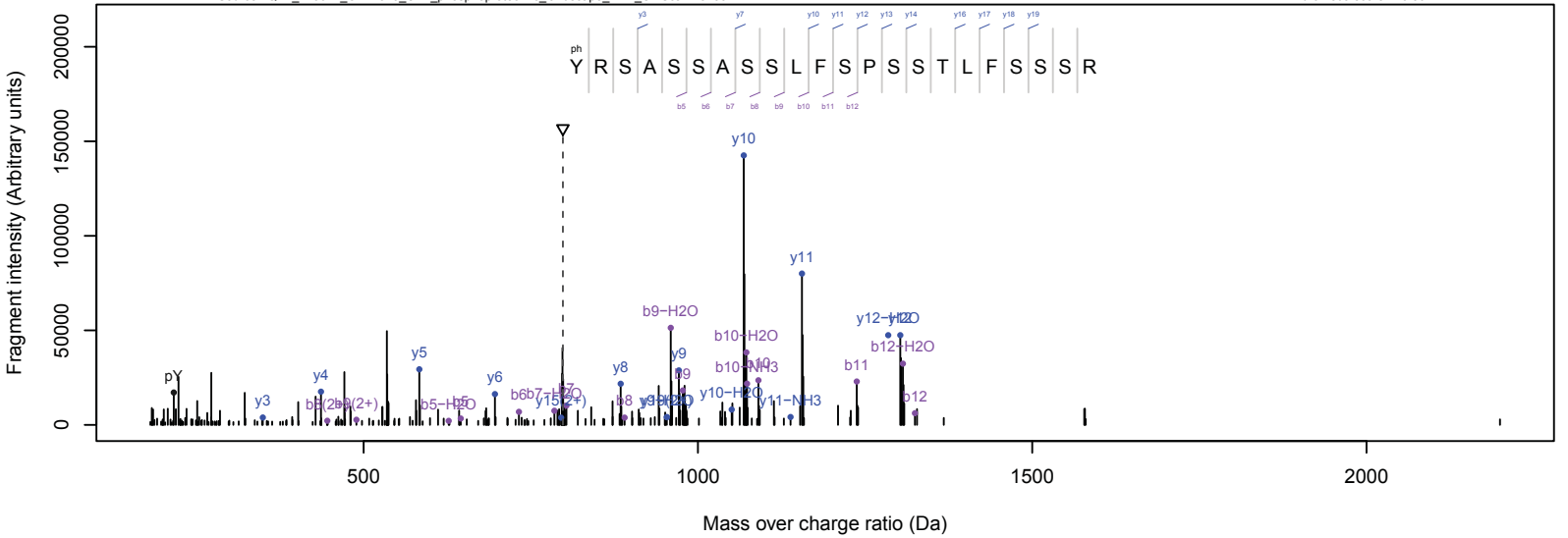
Andromeda score : 76.817



Y<sup>ph</sup>RSASSASSLFS<sup>ph</sup>SPSS<sup>ph</sup>T<sup>ph</sup>LFSSSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37537

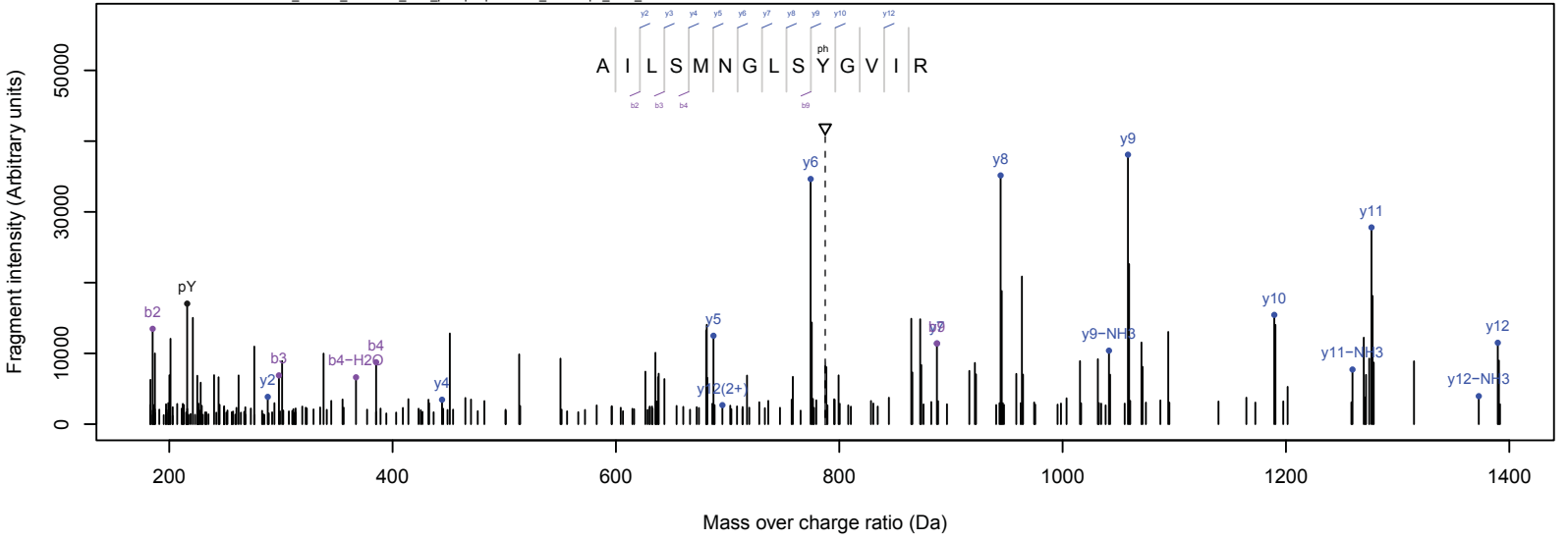
Andromeda score : 76.807



**AILS M N G L S Y G V I R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42319

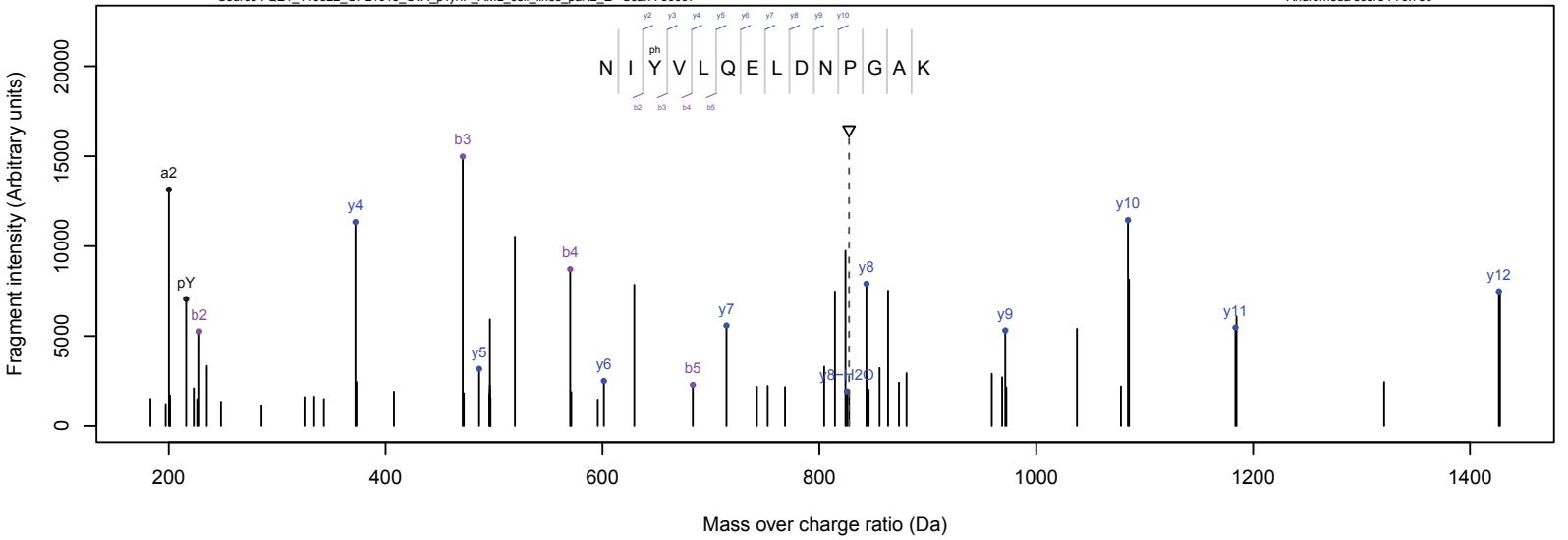
Andromeda score : 76.759



**NIY(ph)VLQELDNP GAK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 30097

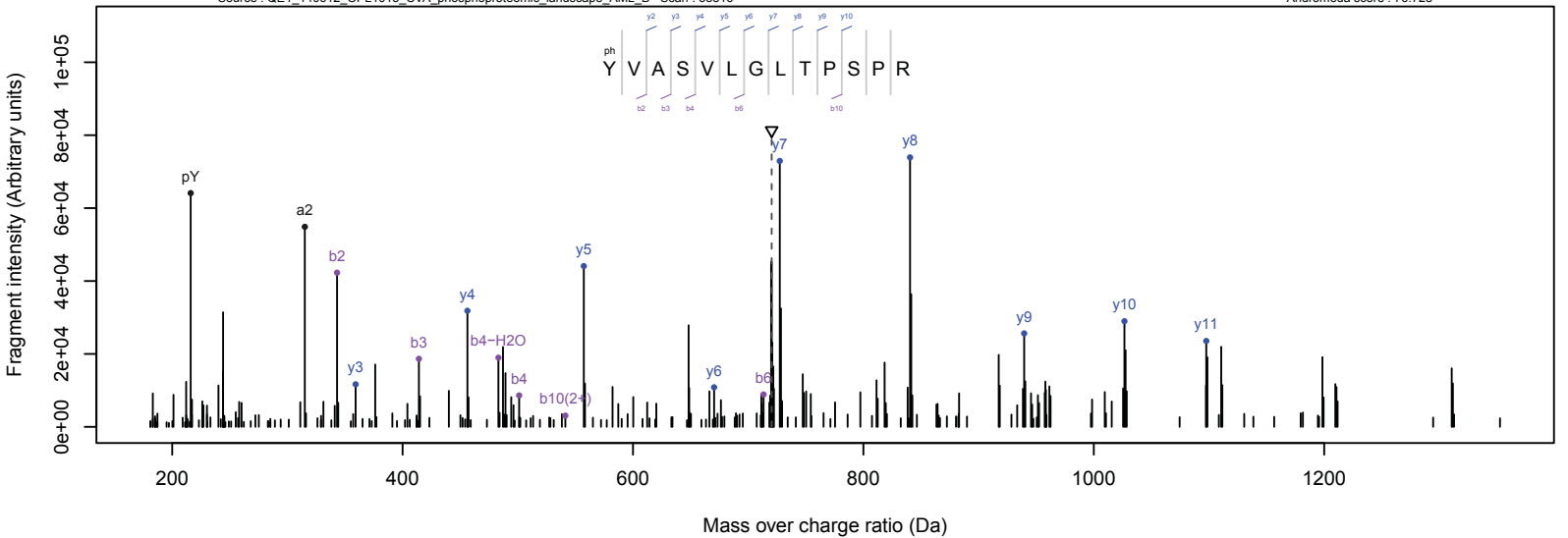
Andromeda score : 76.759



**Y(ph)VASV L G L T P S P R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33816

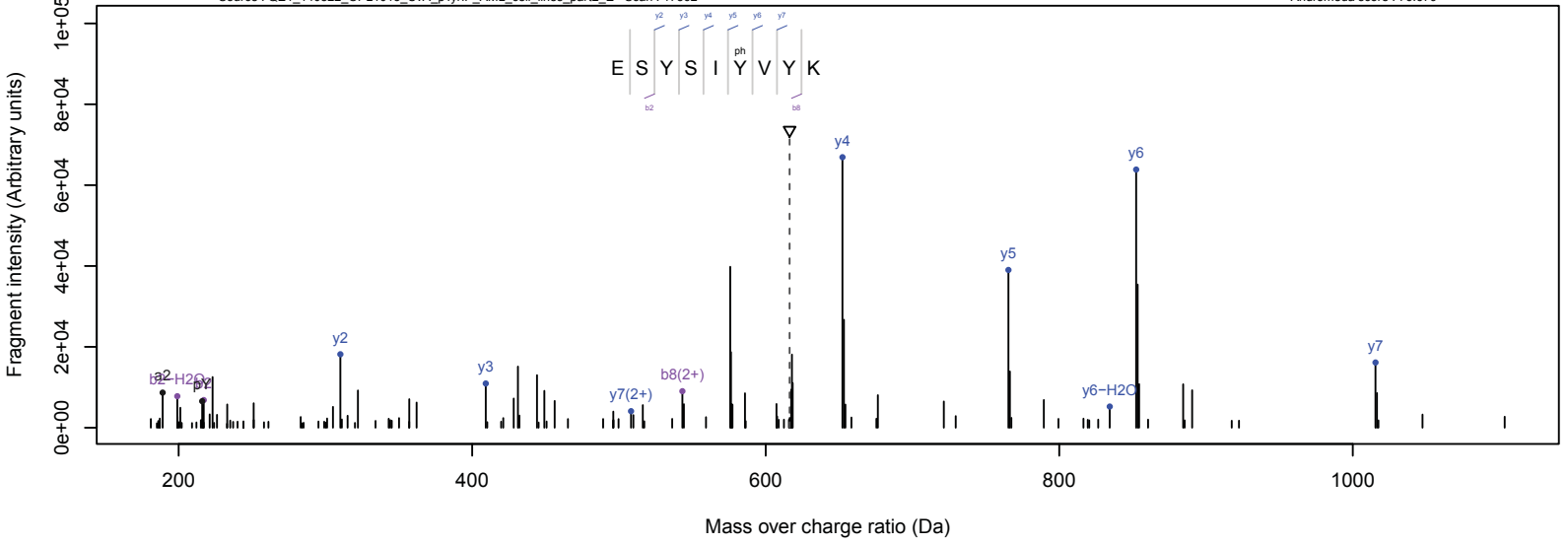
Andromeda score : 76.728



ESYSIY(ph)VYK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 17362

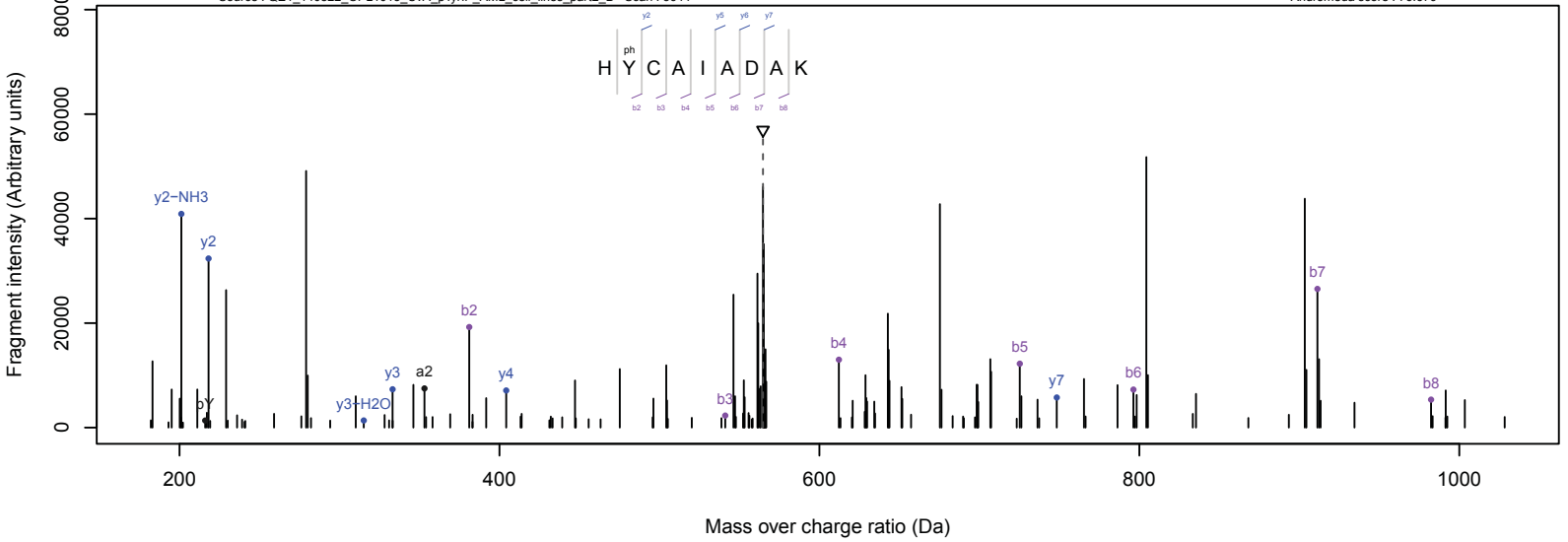
Andromeda score : 76.679



HY(ph)CAIADAK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3644

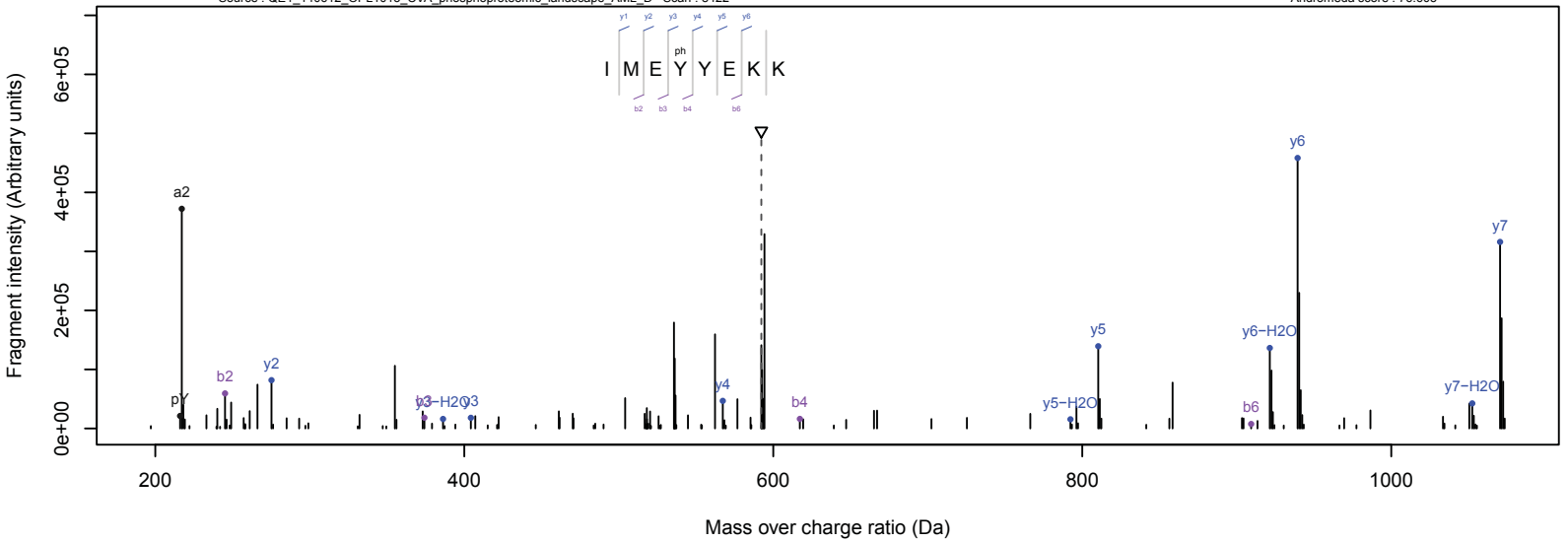
Andromeda score : 76.679



IMEY(ph)YEKK-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan : 8422

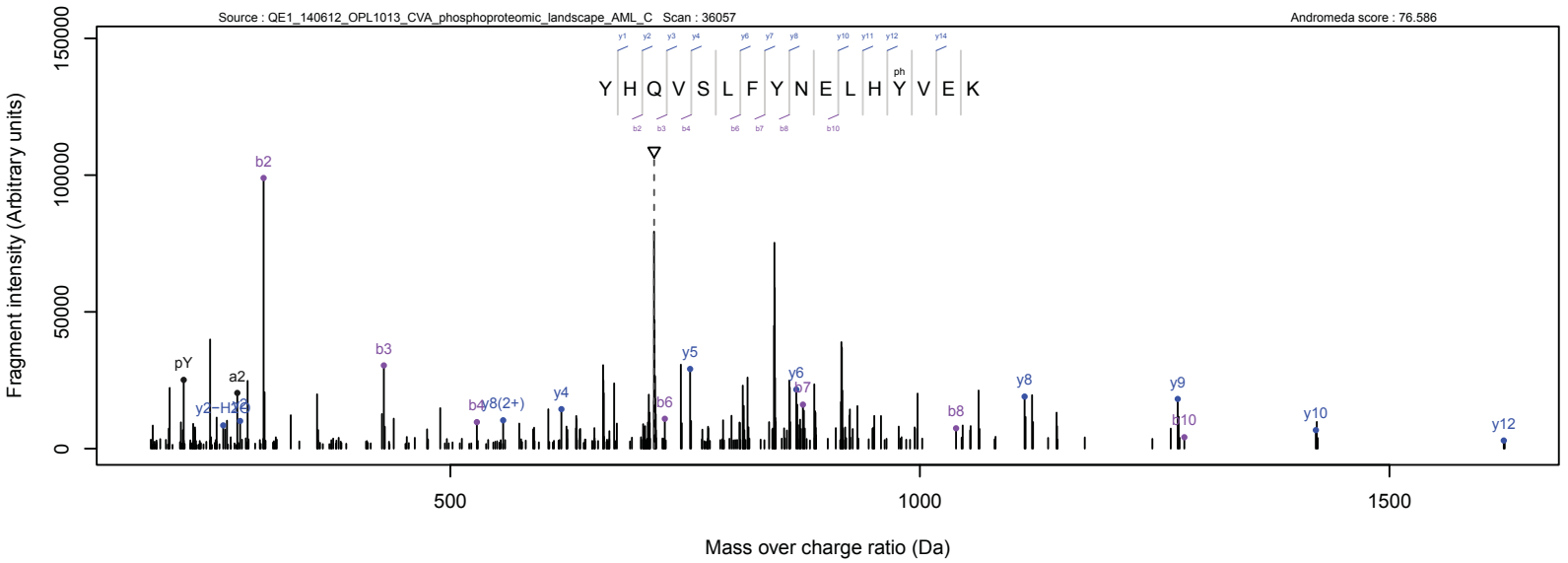
Andromeda score : 76.605



**YHQVSLFYNELHYVEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36057

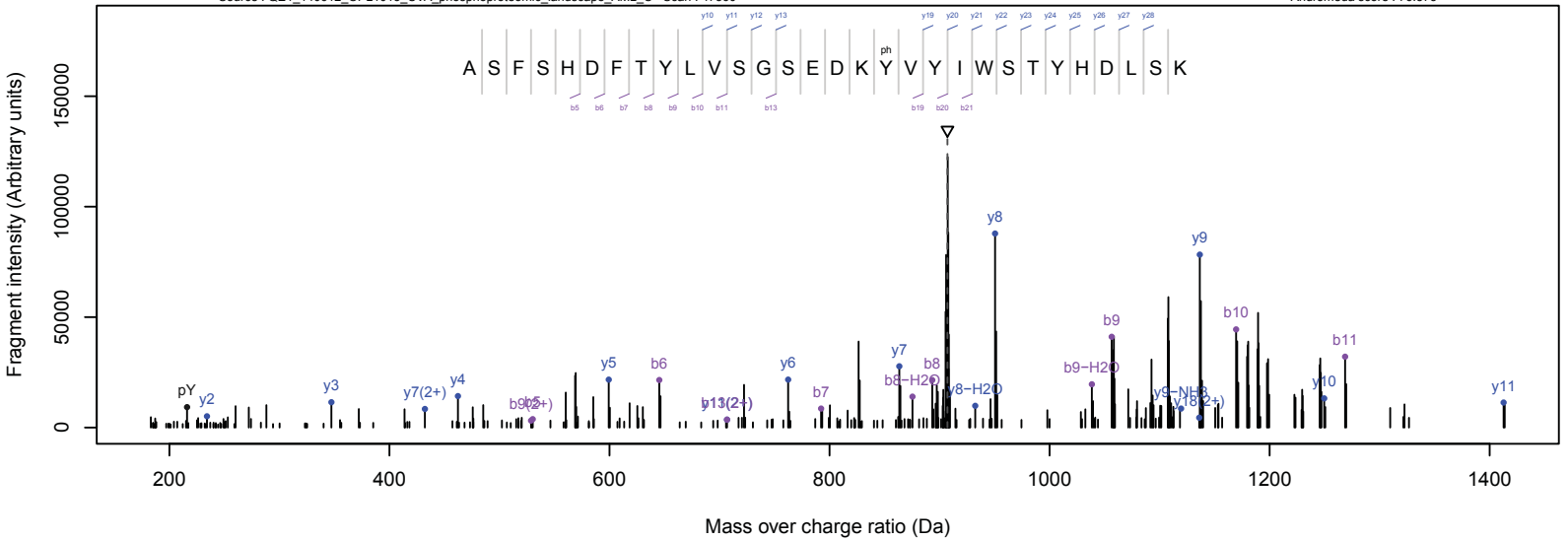
Andromeda score : 76.586



**ASFSHDFTYLVSGSEDKYVYIWSITYHDLISK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47360

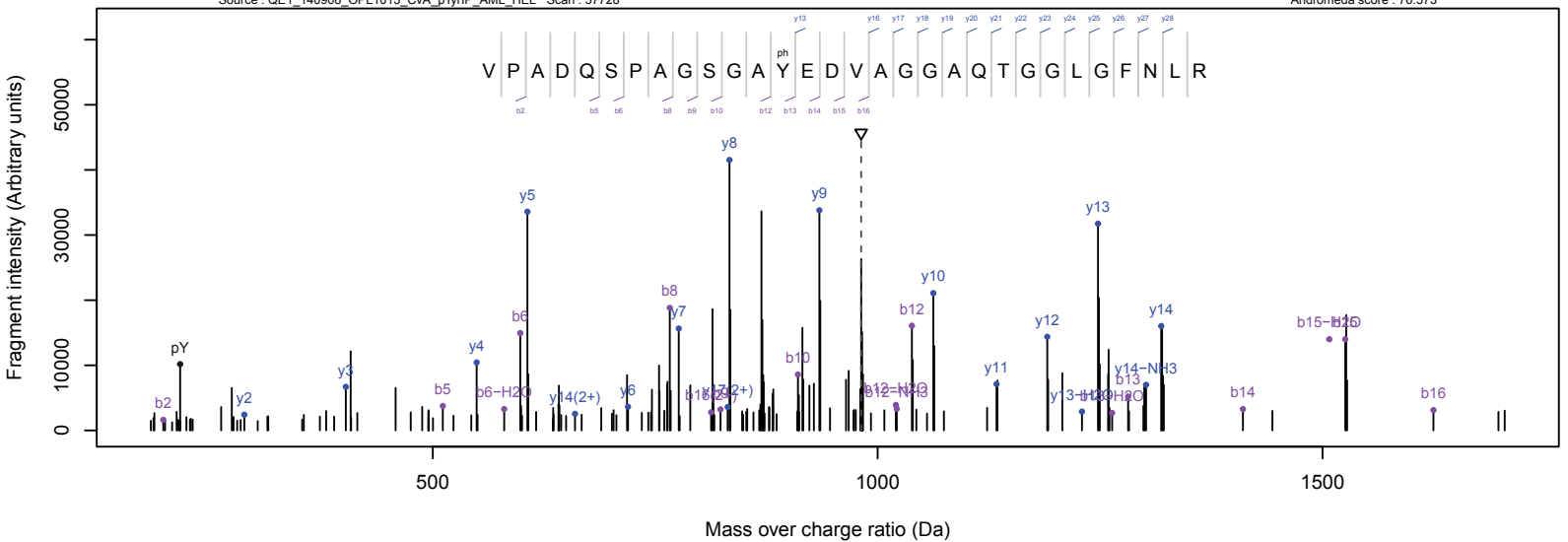
Andromeda score : 76.573



**VPADQSPAGSGAYEDVAGGAQTGGGLGFNLR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37728

Andromeda score : 76.573

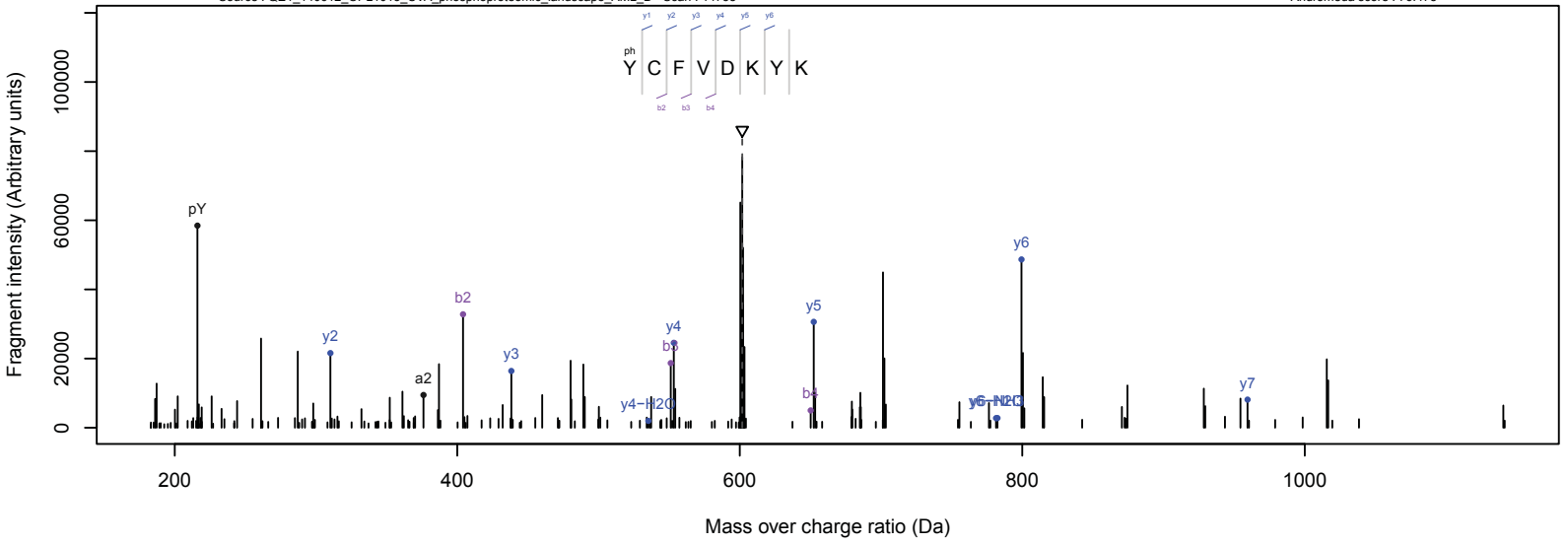




Y(ph)CFVDKYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14758

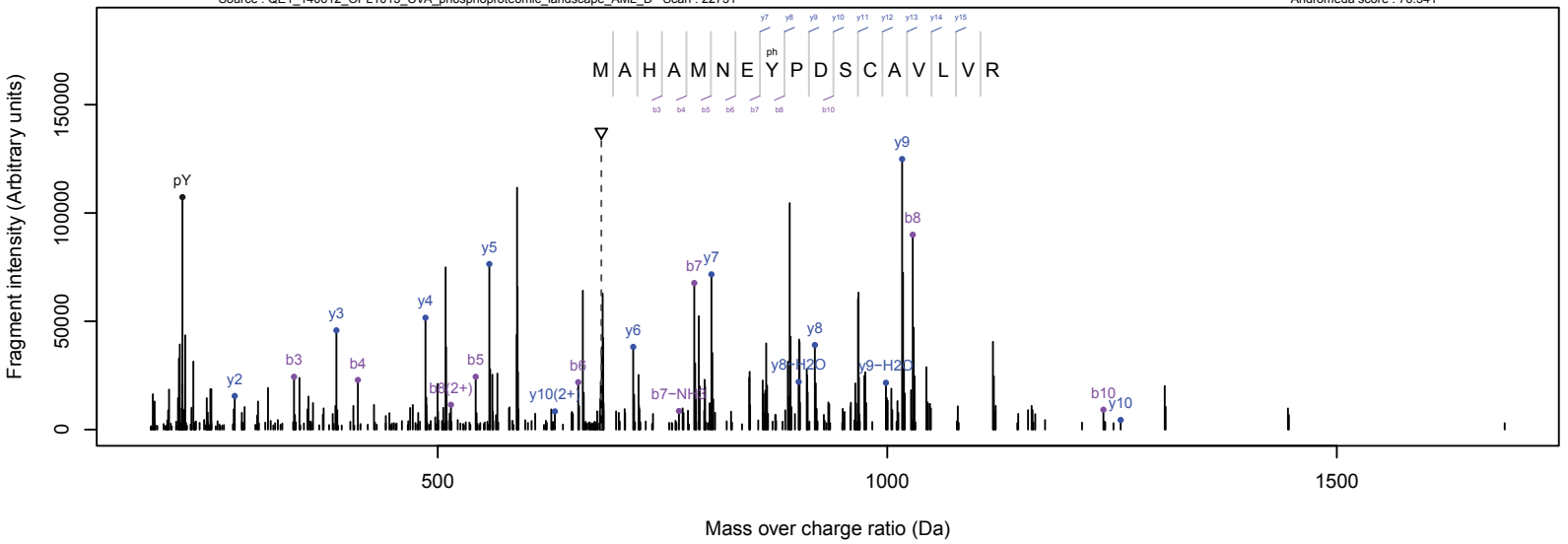
Andromeda score : 76.478



MAHAMNEY(ph)PDSCAVLVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22731

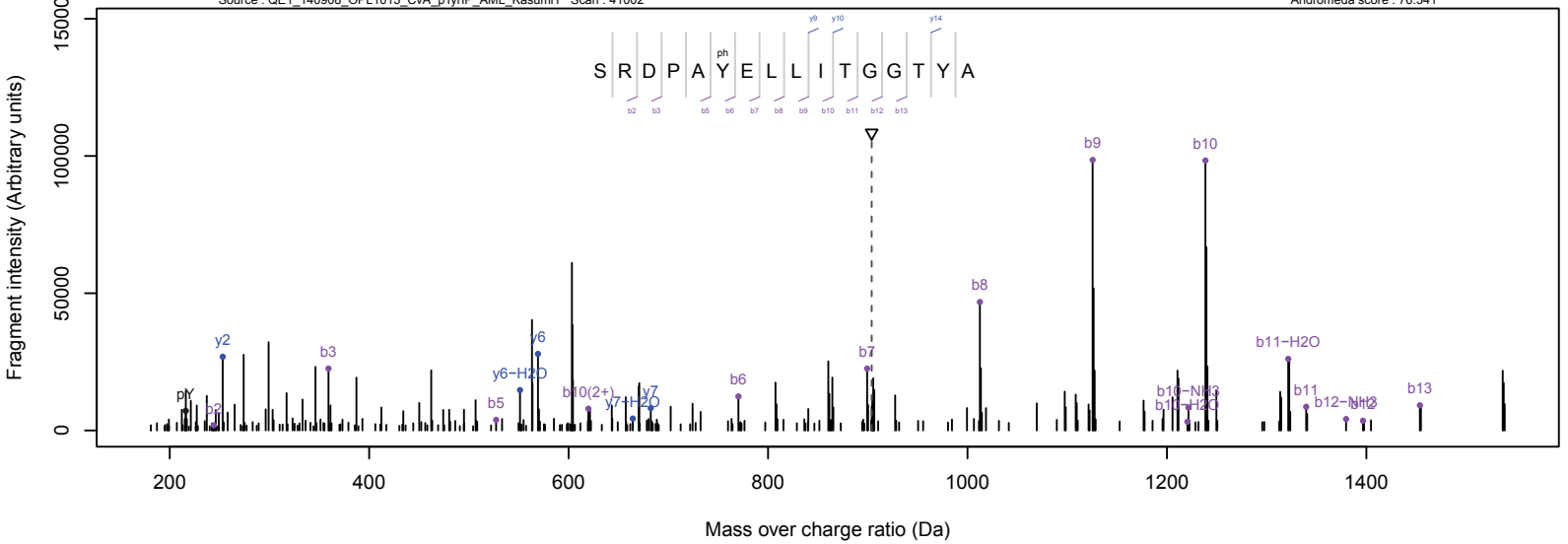
Andromeda score : 76.341



SRDPAY(ph)ELLITGGTYA-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 41002

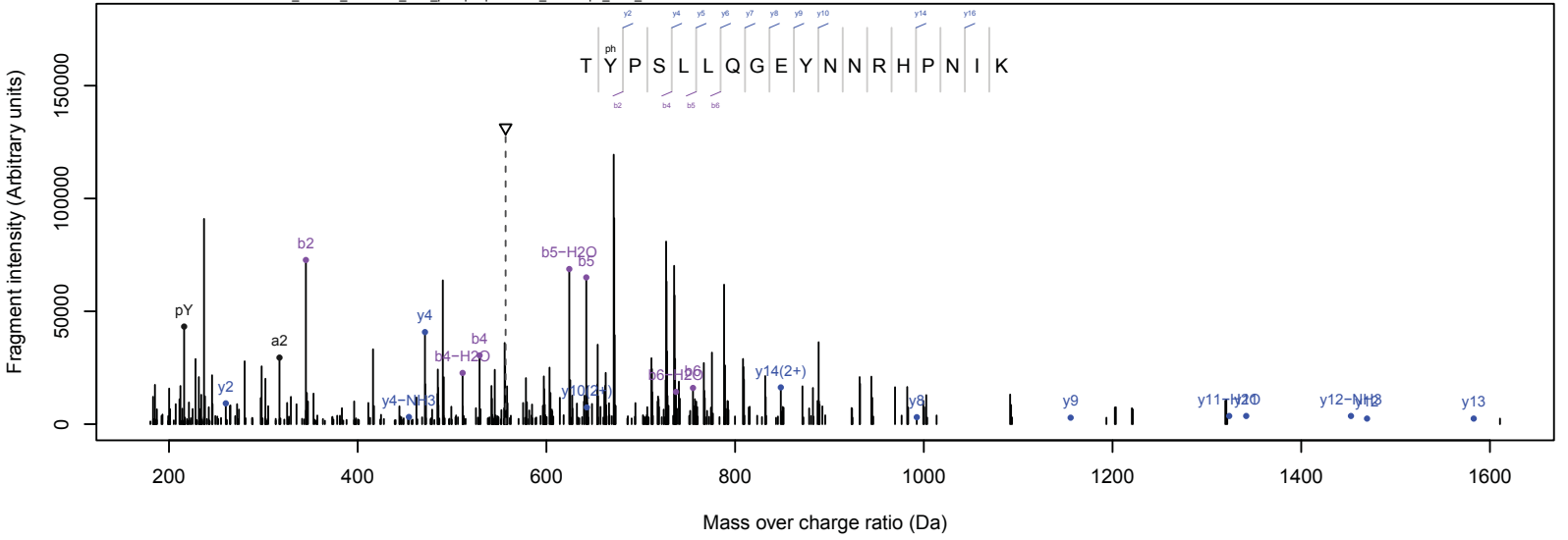
Andromeda score : 76.341



TY(ph)PSLLQGEYNNRHPNIK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21573

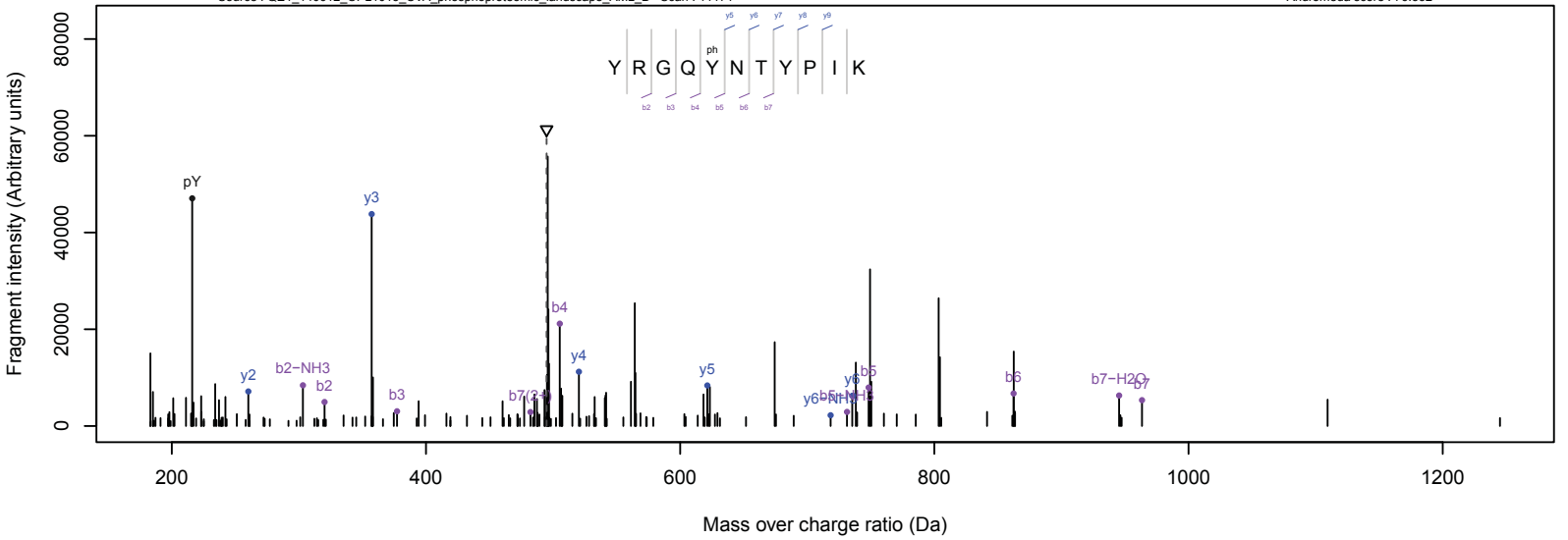
Andromeda score : 76.341



YRGQY(ph)NTYPIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11171

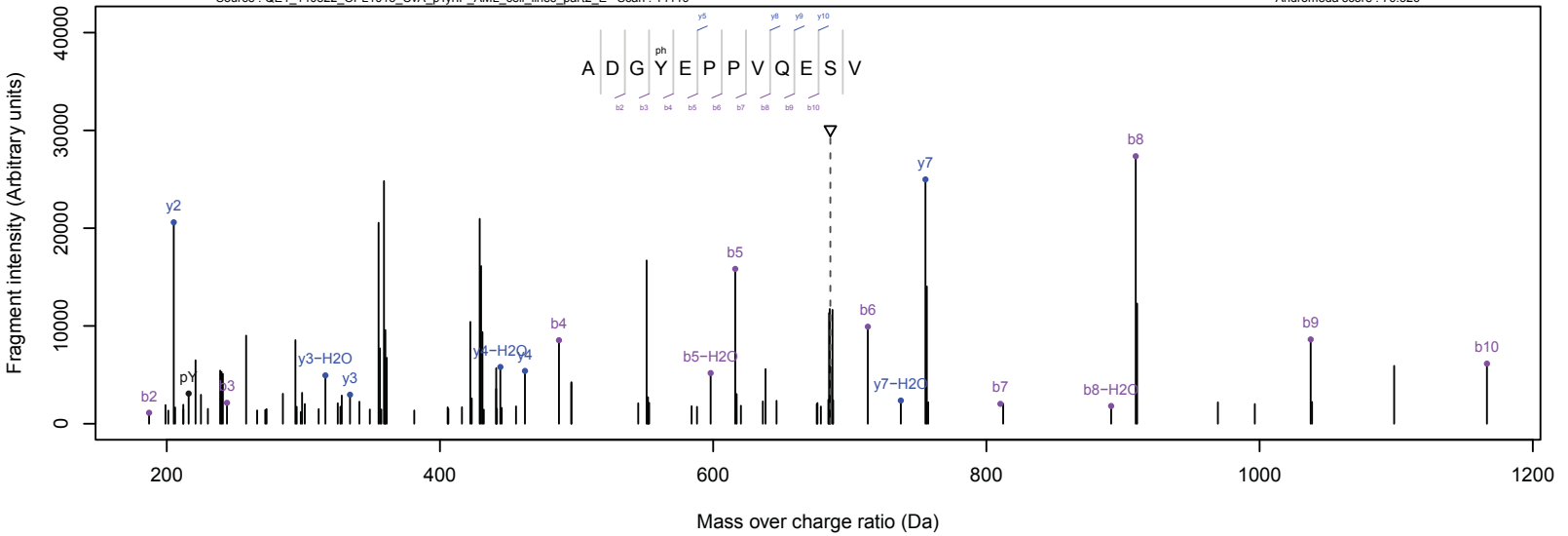
Andromeda score : 76.332



ADGY(ph)EPPVESV-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 14119

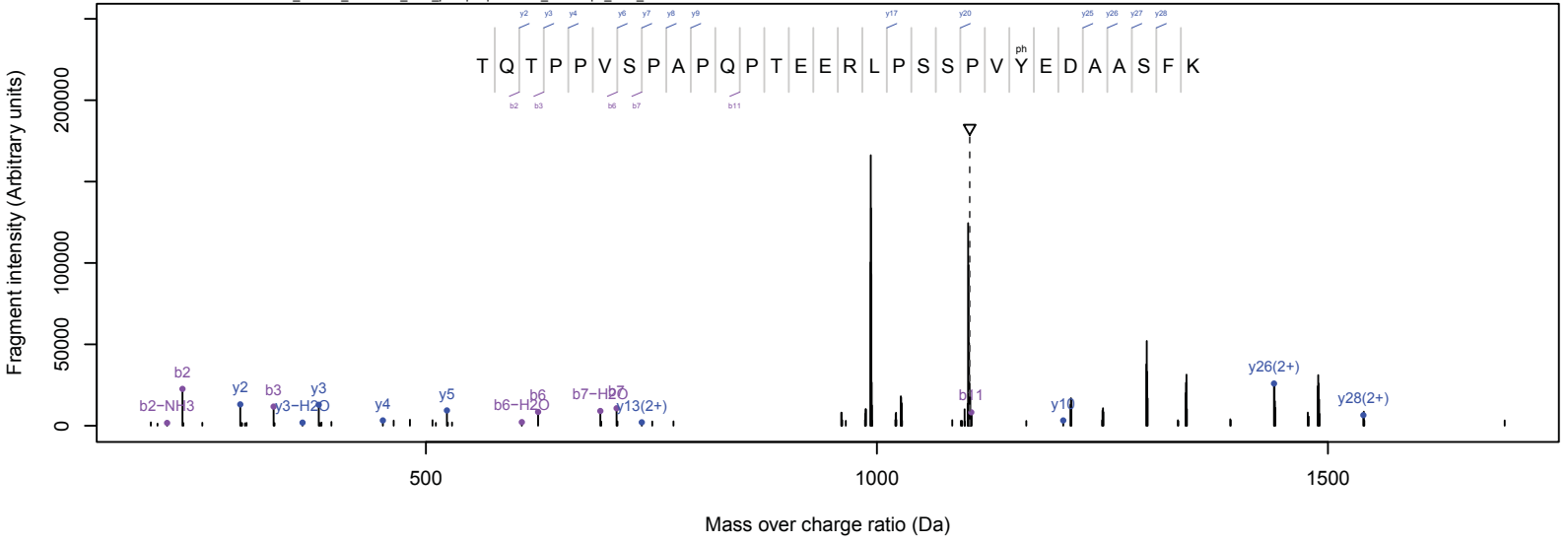
Andromeda score : 76.326



TQTTPVSPAPQTEERLPSSPV(ph)EDAASFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 32204

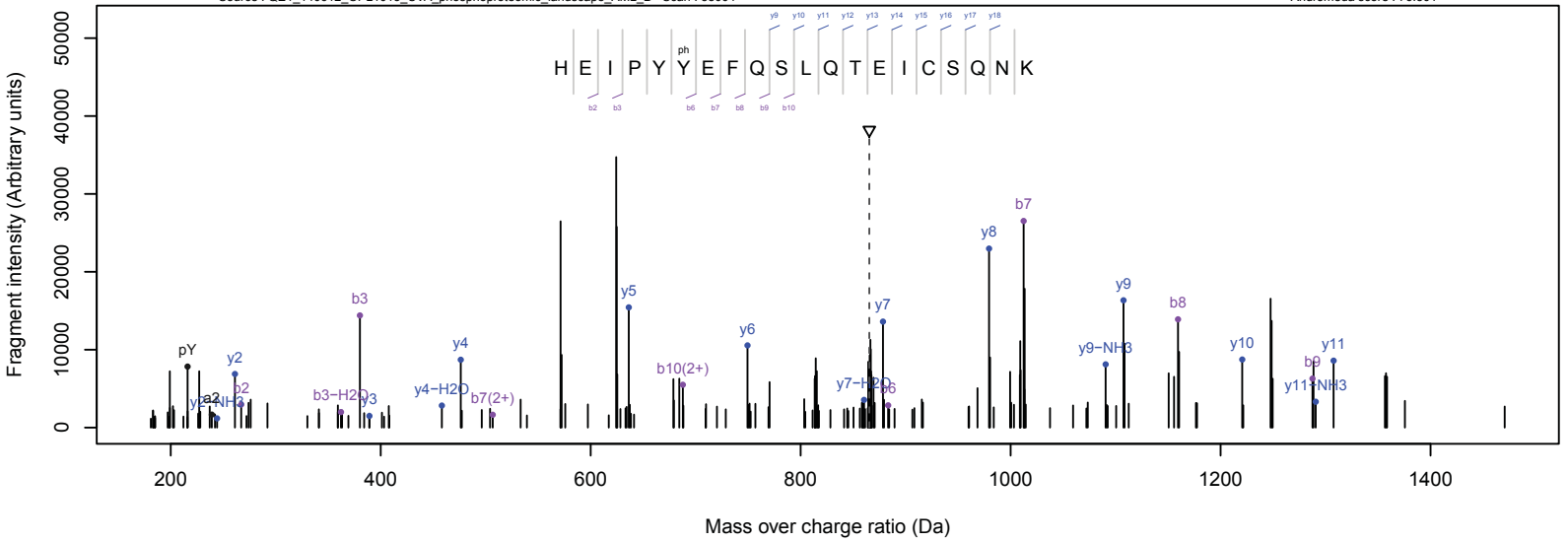
Andromeda score : 76.31



HEIPYY(ph)EFQSLQTEICSQNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38004

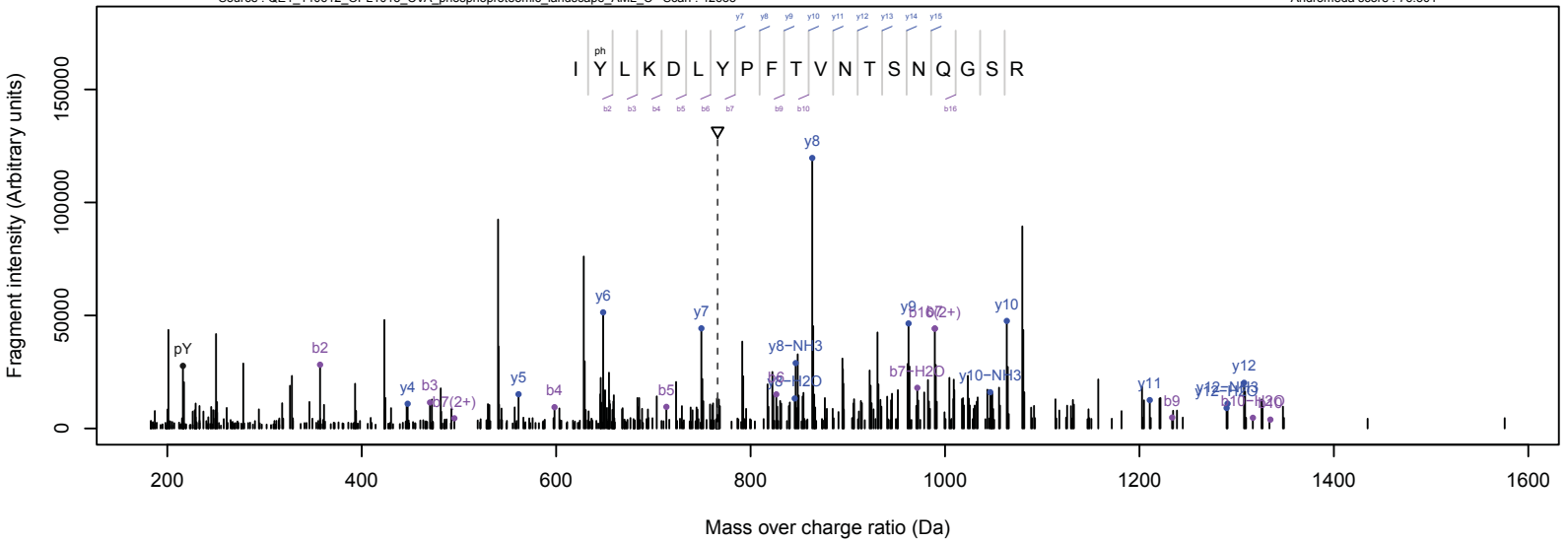
Andromeda score : 76.301



IY(ph)LKDLYPFTVNTSNQGSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42635

Andromeda score : 76.301

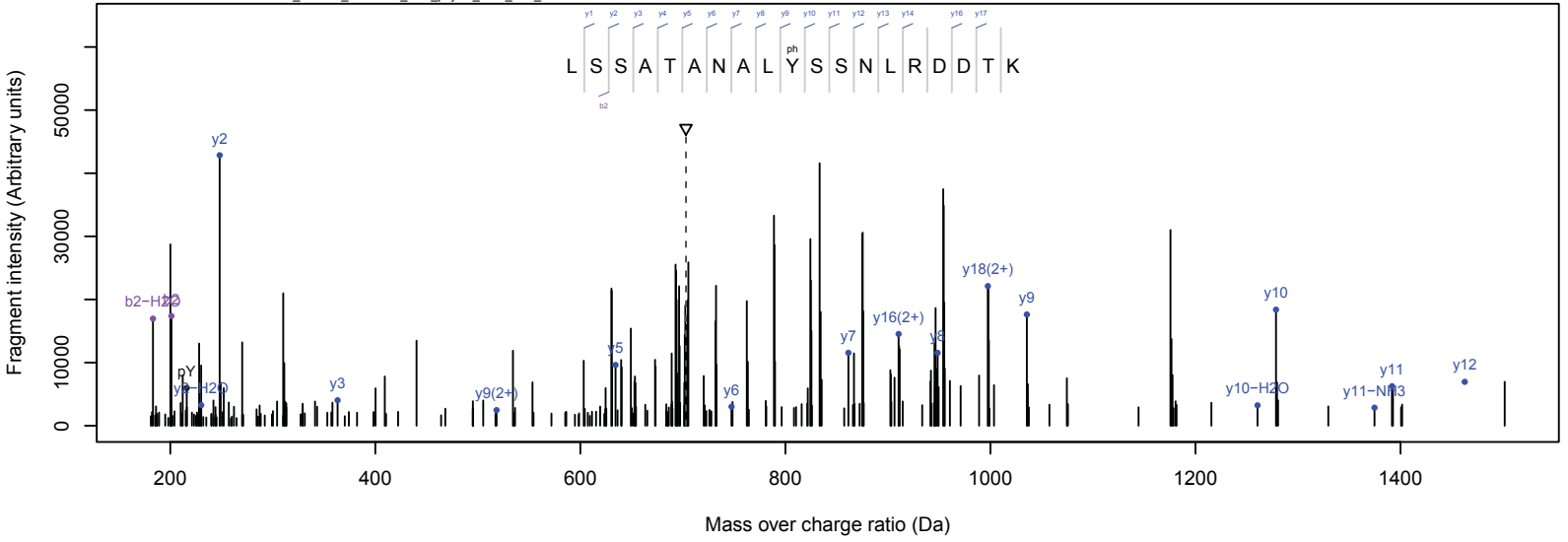




LSSATANALY(ph)SSNLRDDTK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 21139

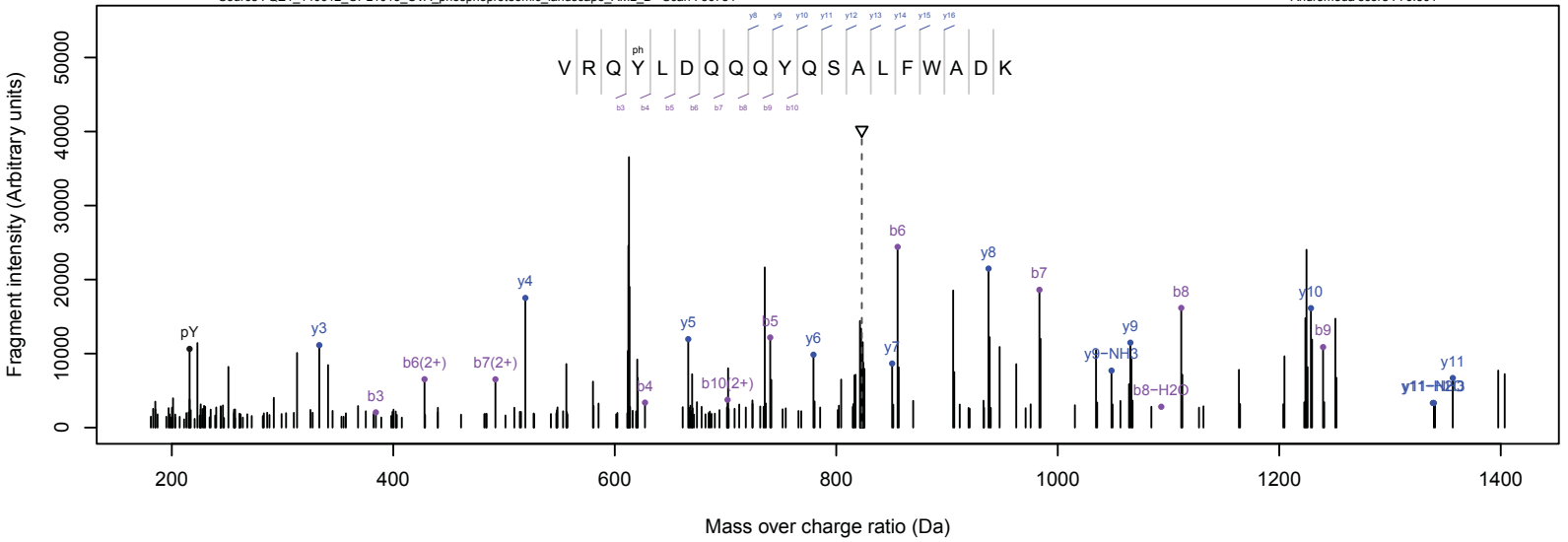
Andromeda score : 76.301



VRQY(ph)LDQQYQSALFWADK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan : 36784

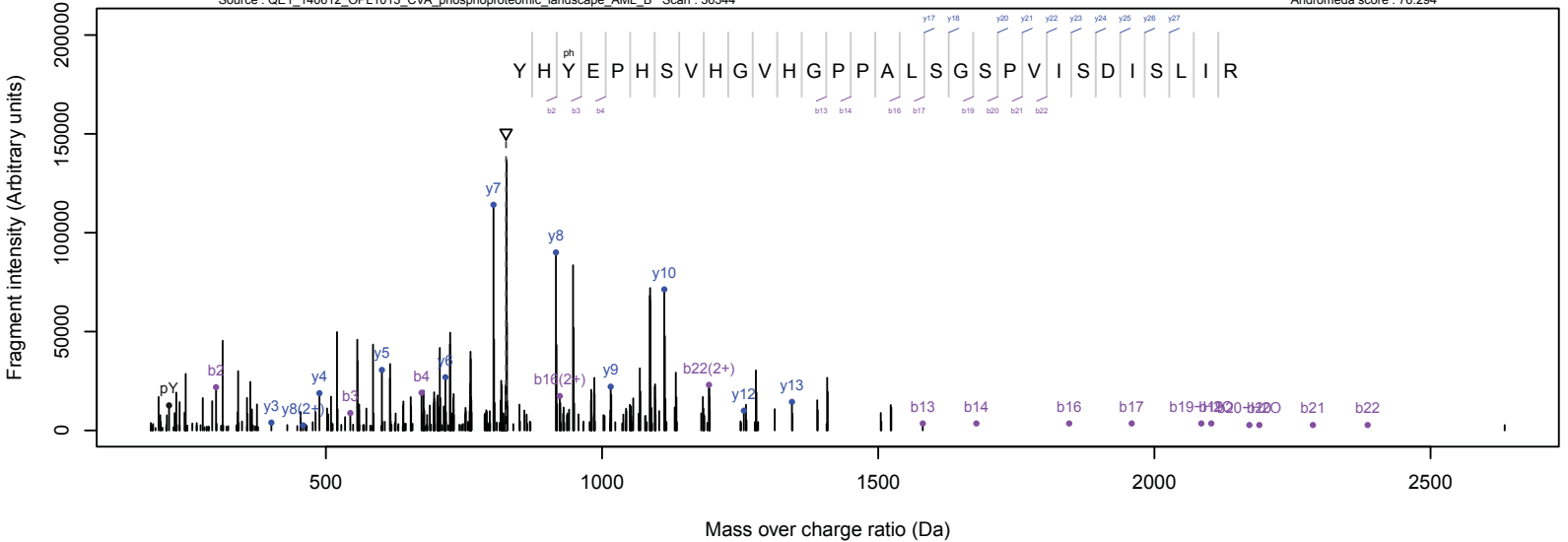
Andromeda score : 76.301



YHY(ph)EPHSVHG VHGPPALSGSPVISDISLIR-4+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan : 36344

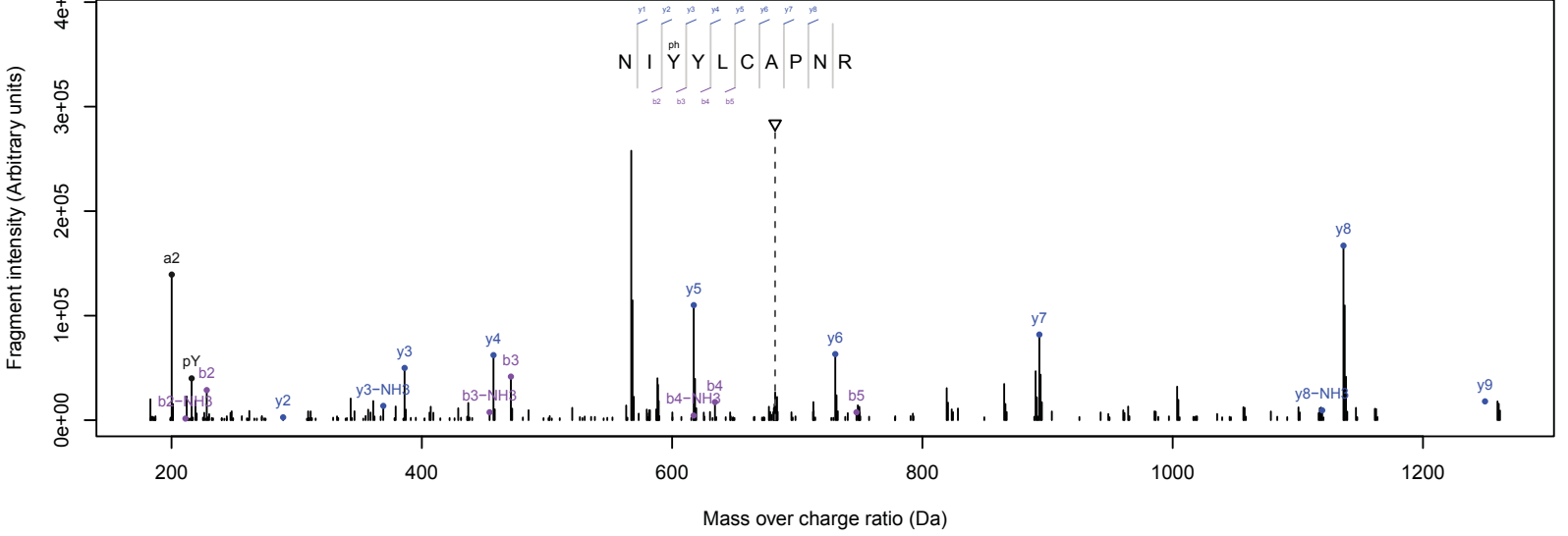
Andromeda score : 76.294



NIY(ph)YLCAPNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23992

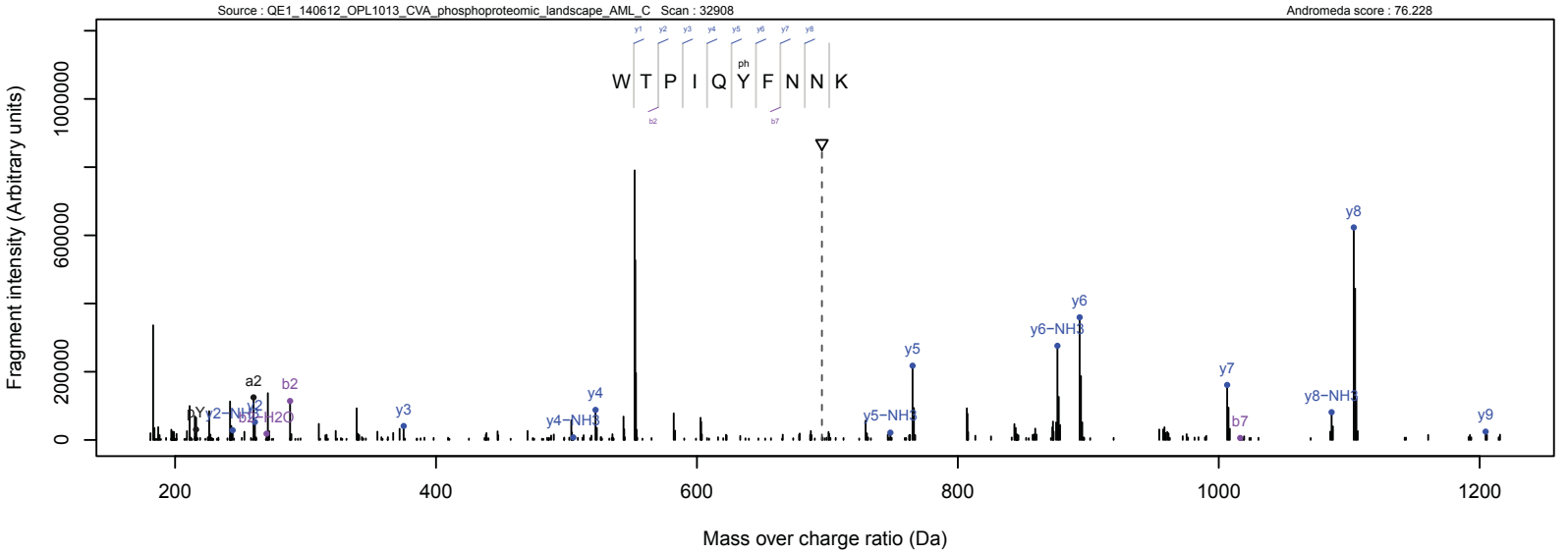
Andromeda score : 76.228



WTPIQY(ph)FNKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32908

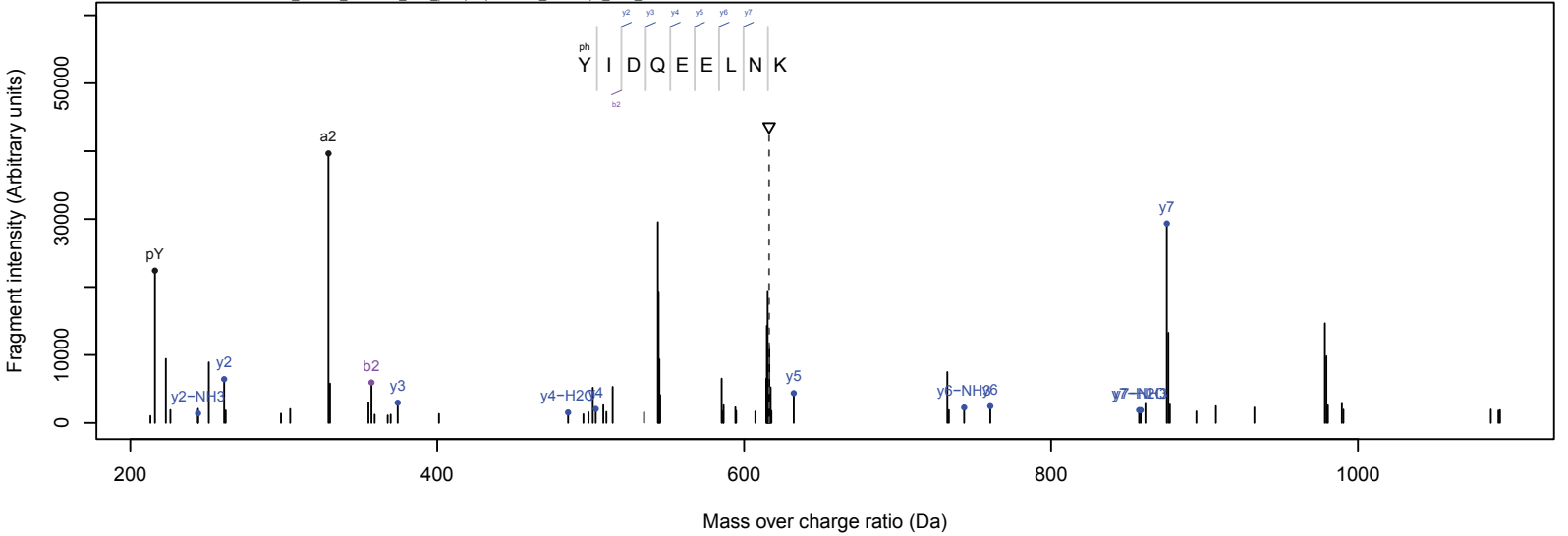
Andromeda score : 76.228



Y(ph)IDQEELNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9501

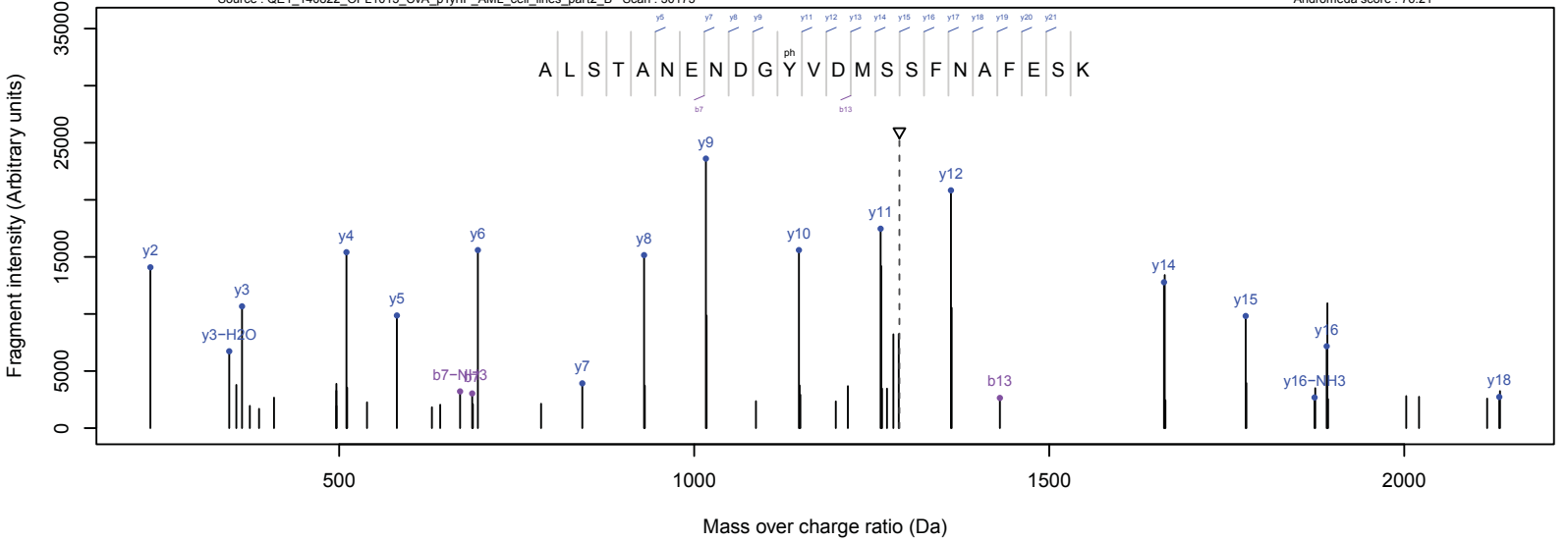
Andromeda score : 76.221



ALSTANENDGY(ph)VDMSSFNAFESK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 30173

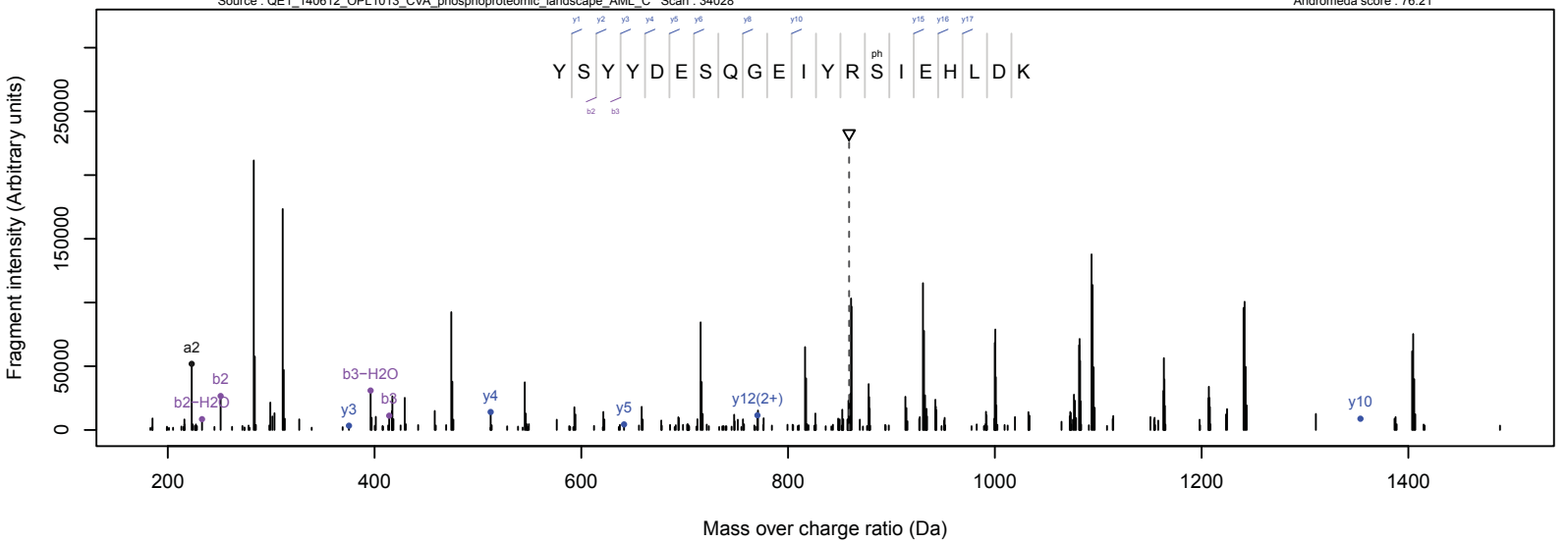
Andromeda score : 76.21



YSYDESQGEIYR(ph)IEHLDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34028

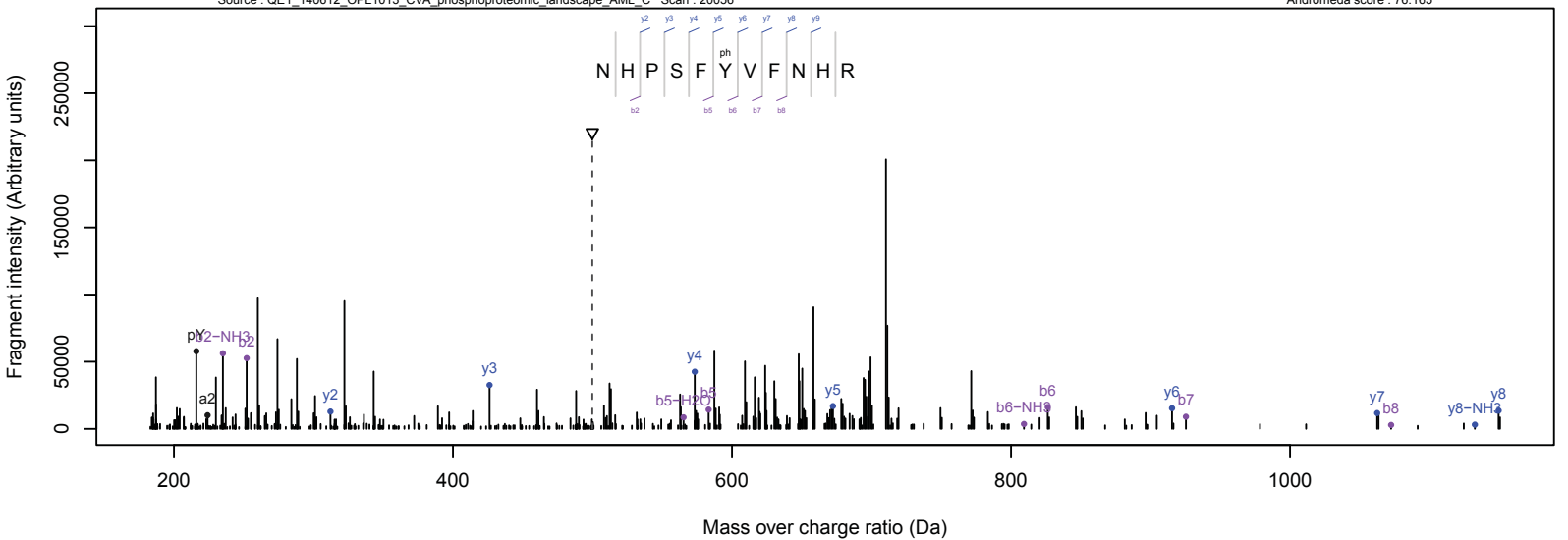
Andromeda score : 76.21



NHPSFY(ph)VFNHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20056

Andromeda score : 76.165

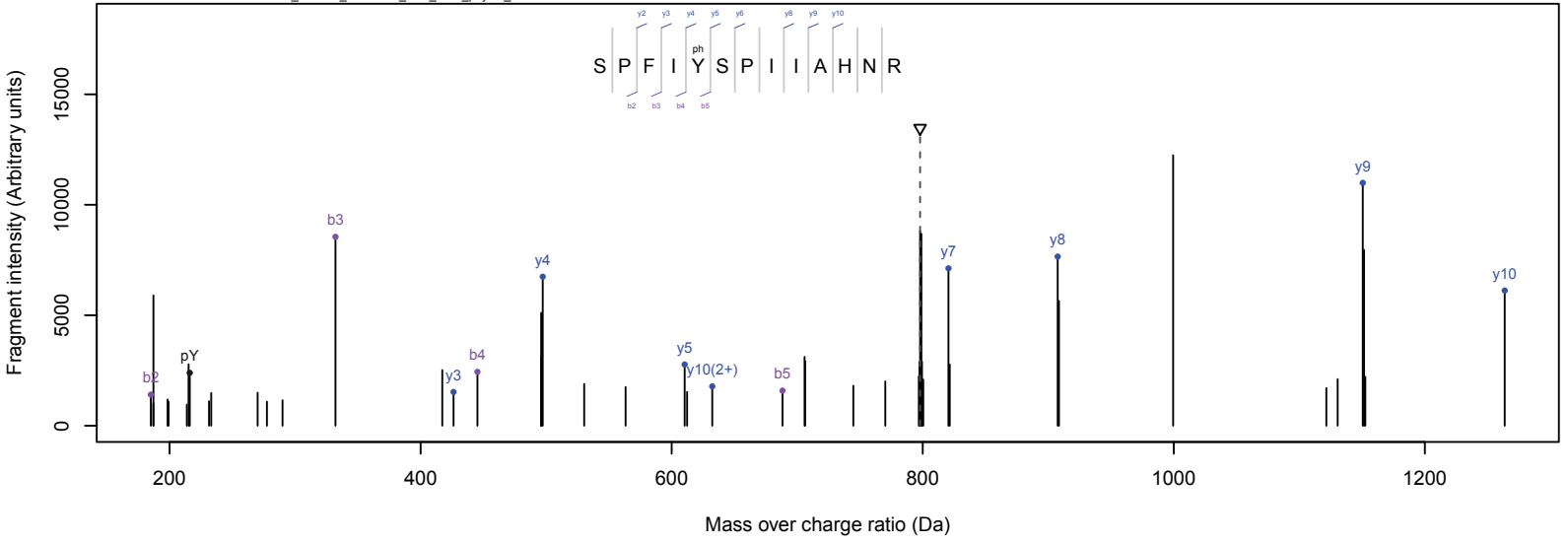




SPFIY(ph)SPIAHNR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 30646

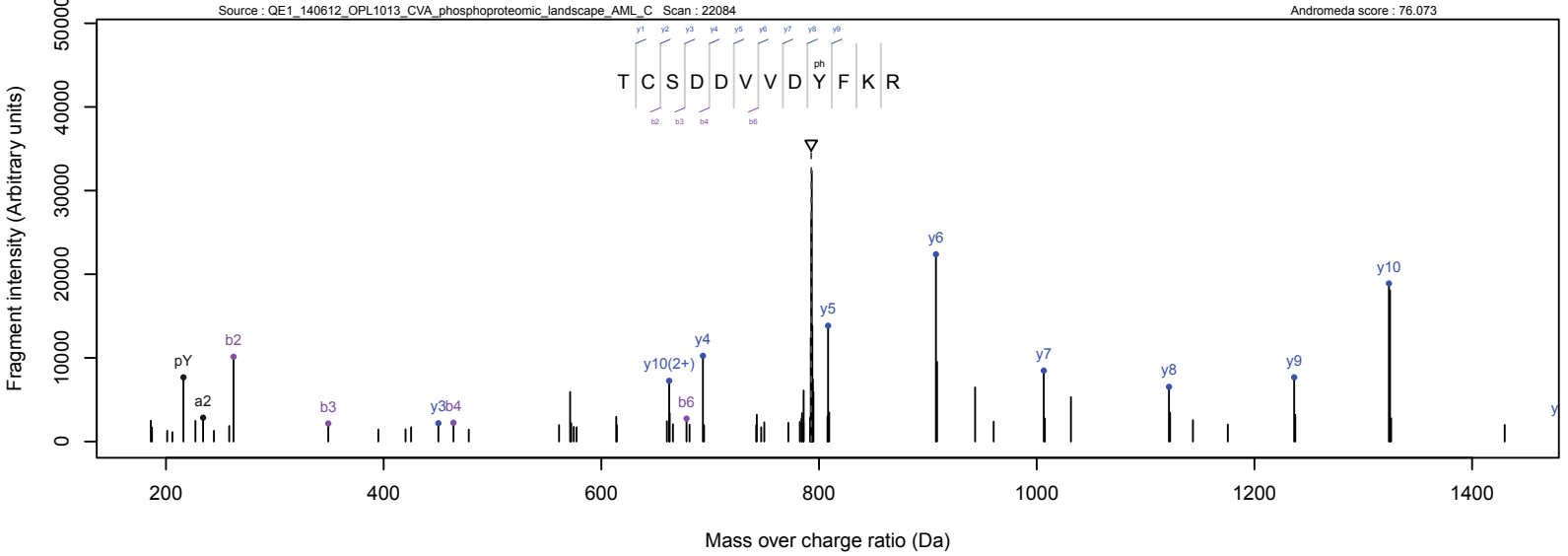
Andromeda score : 76.073



TCSDDVVDY(ph)FKR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22084

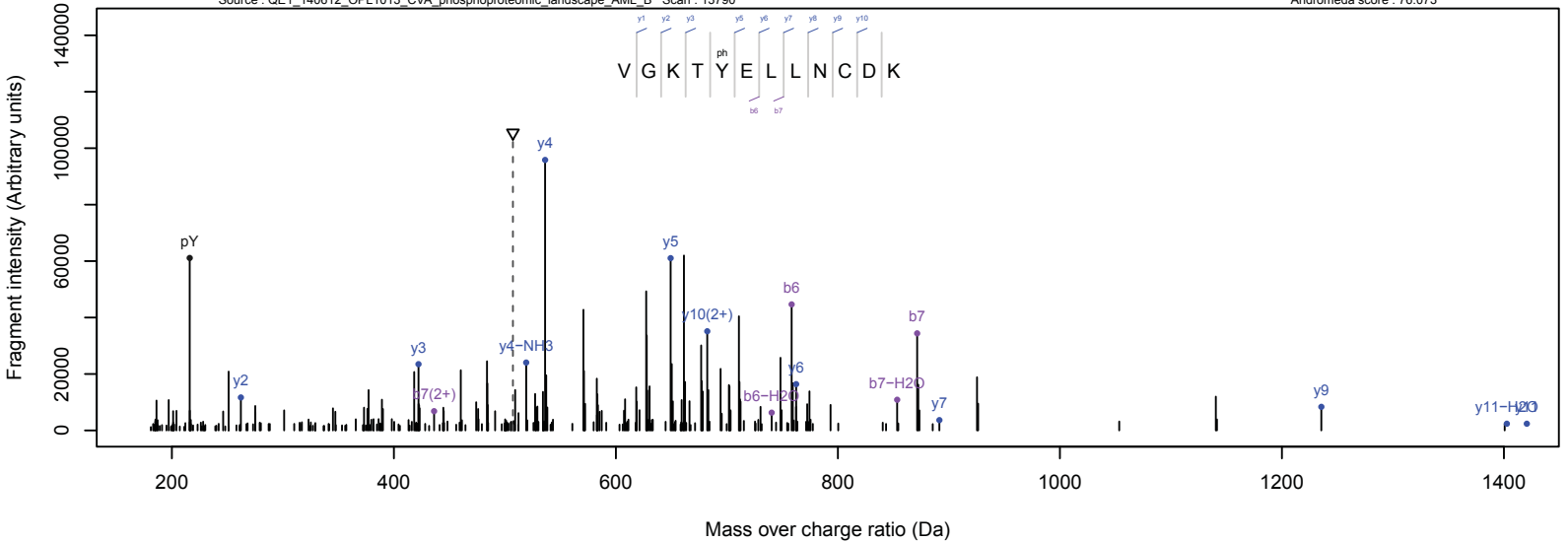
Andromeda score : 76.073



VGKTY(ph)ELLNCDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13790

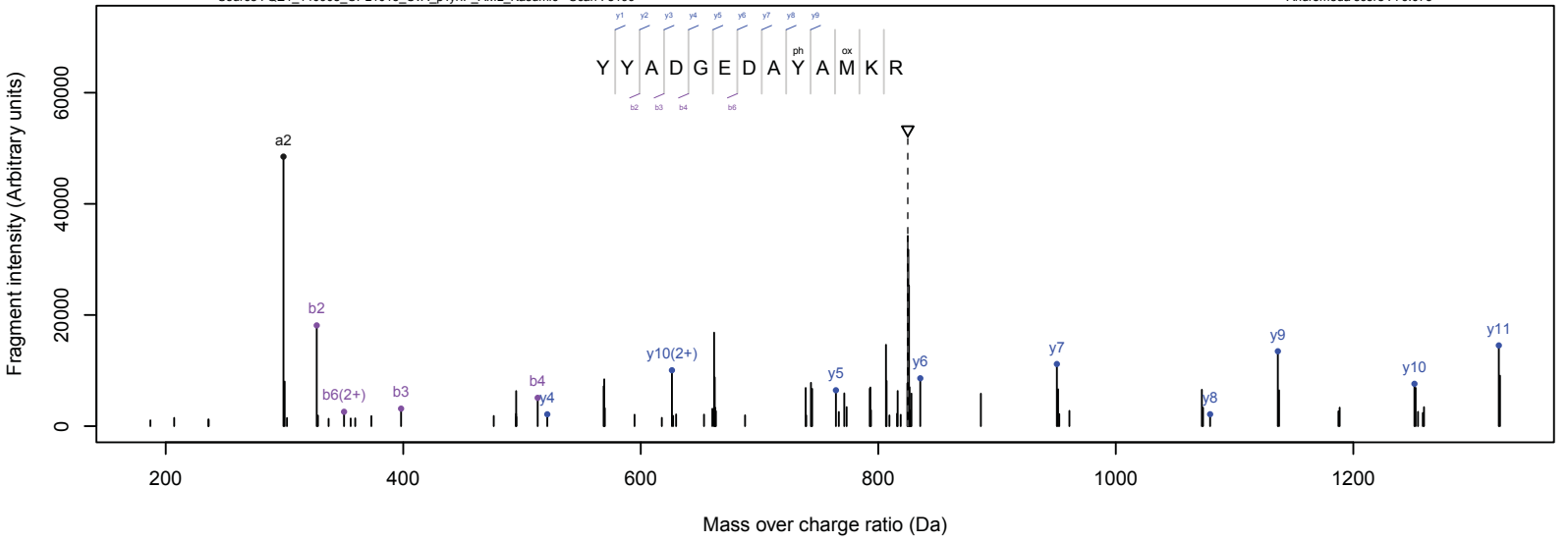
Andromeda score : 76.073



YYADGEDAY(ph)AM(ox)KR-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 8199

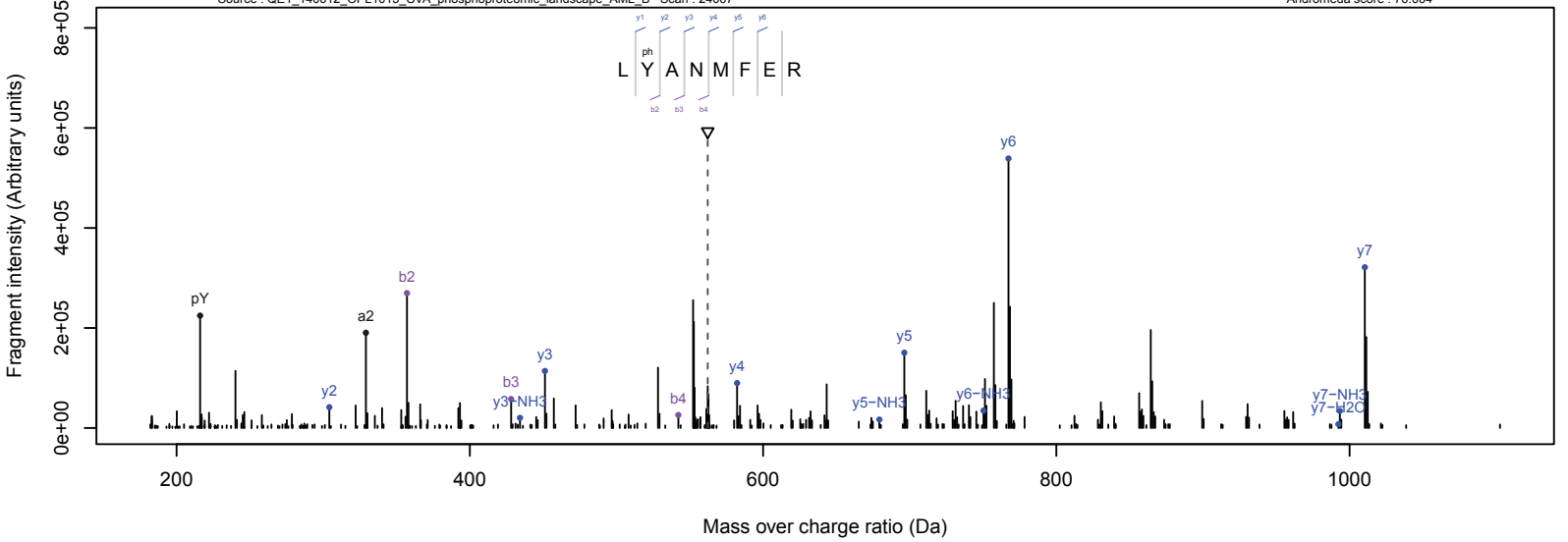
Andromeda score : 76.073



LY(ph)ANMFER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24067

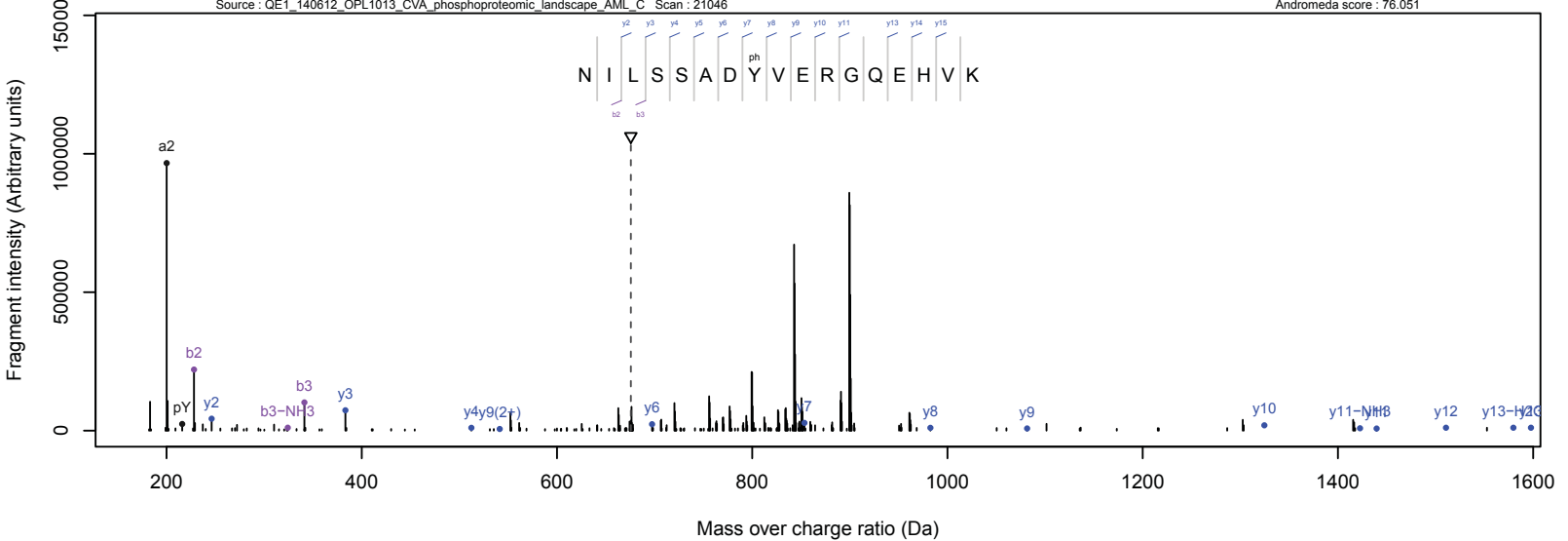
Andromeda score : 76.064



NILSSADY(ph)VERGQEHVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21046

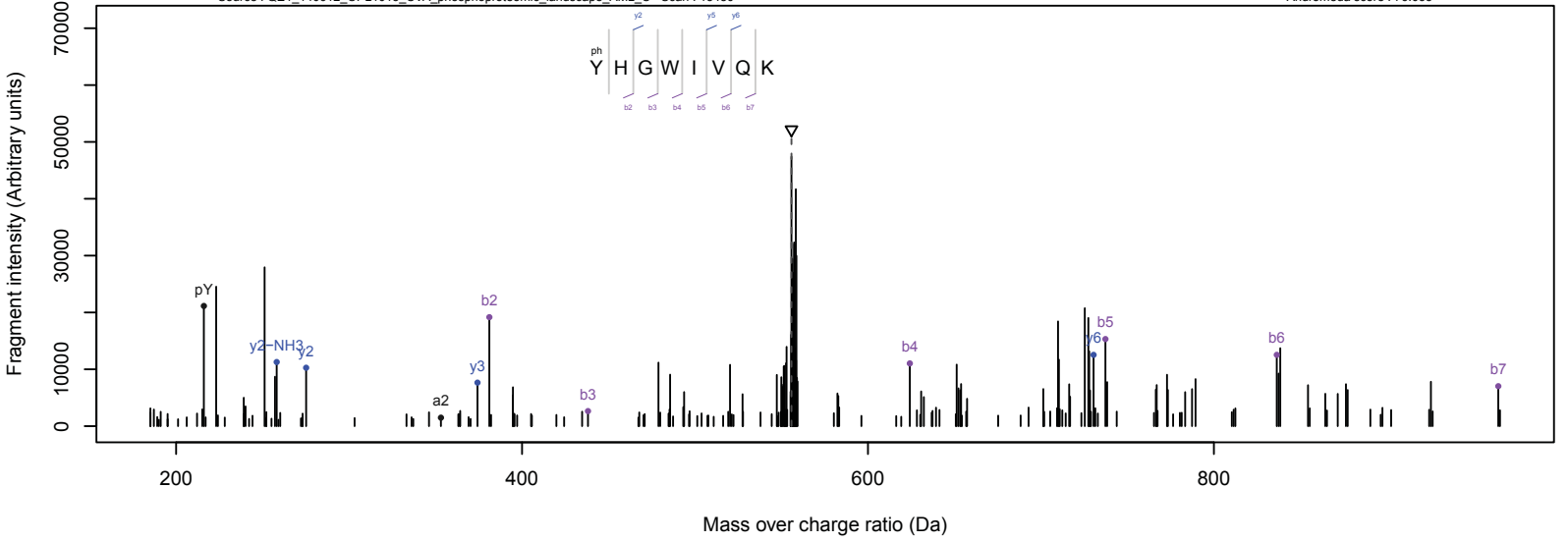
Andromeda score : 76.051



Y(ph)HGWIVQK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13480

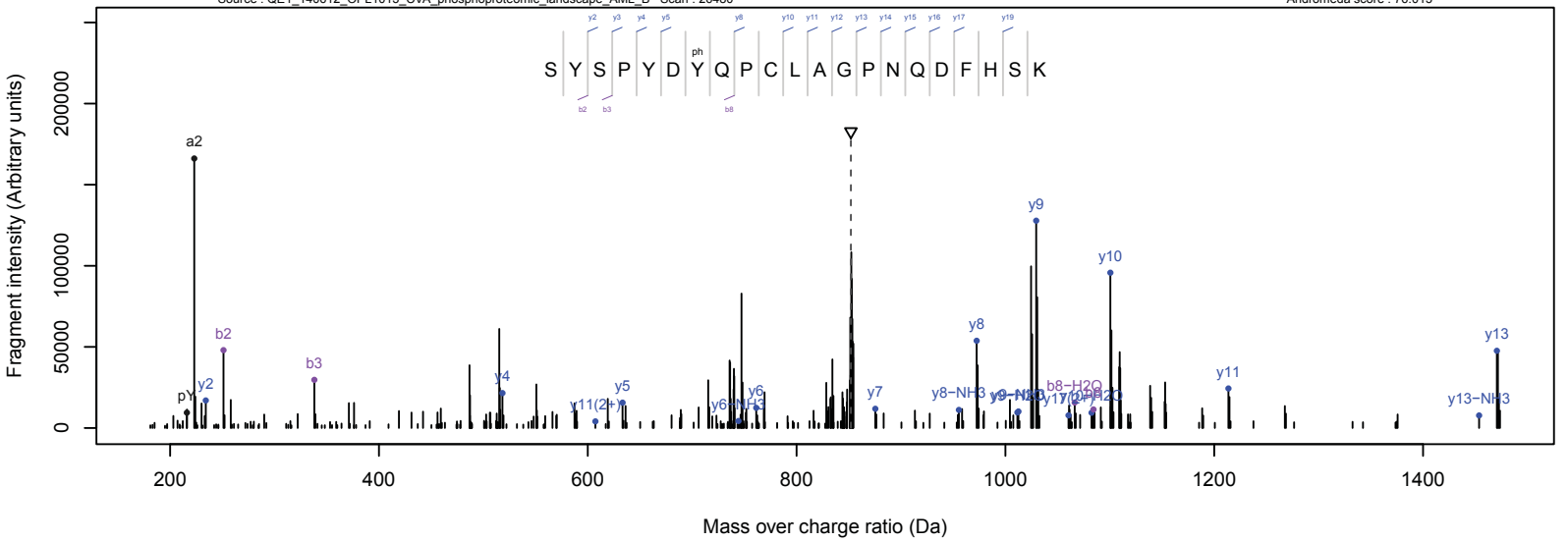
Andromeda score: 76.035



SYSPYDY(ph)QPCLAGPNQDFHSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 26480

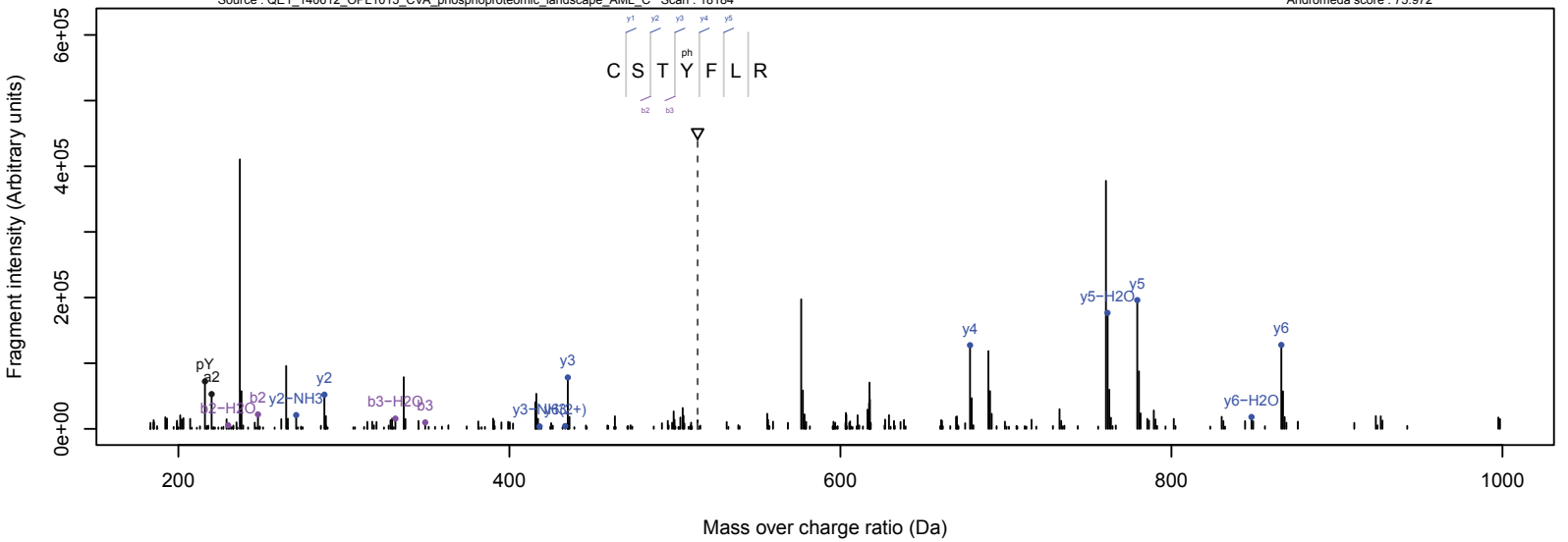
Andromeda score: 76.015



CSTY(ph)FLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18184

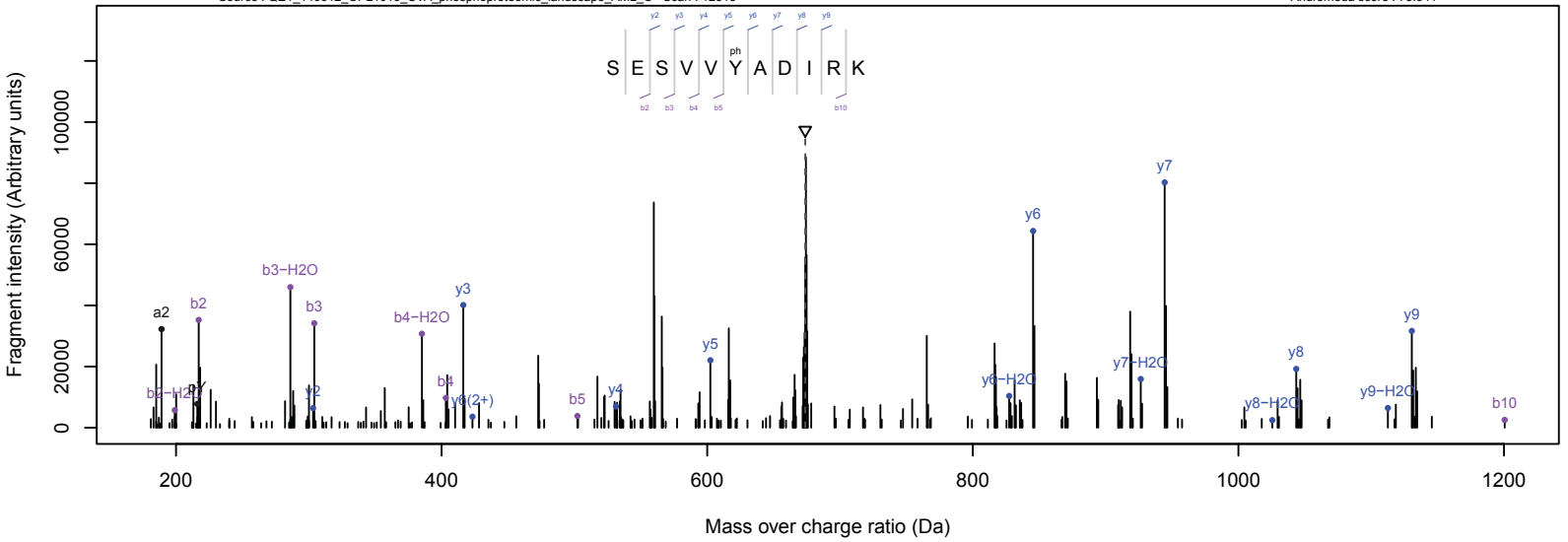
Andromeda score: 75.972



SESVVY(ph)ADIRK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12818

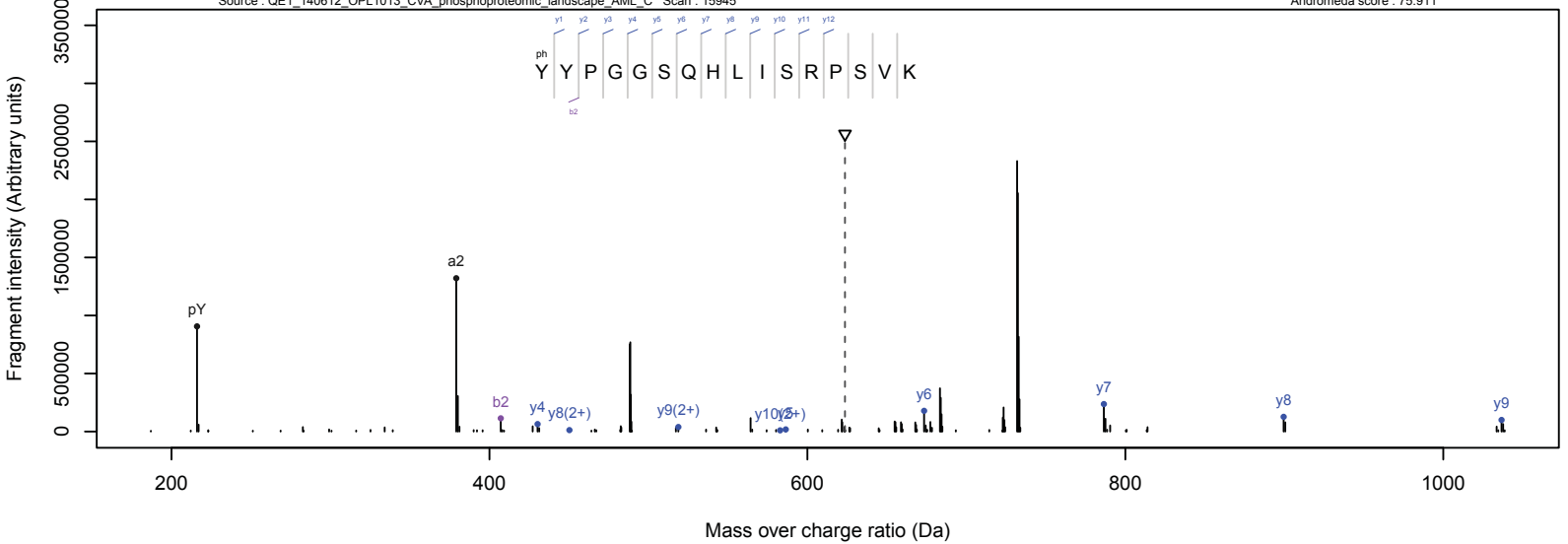
Andromeda score : 75.941



Y(ph)YPPGGSQHLISRPVSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15945

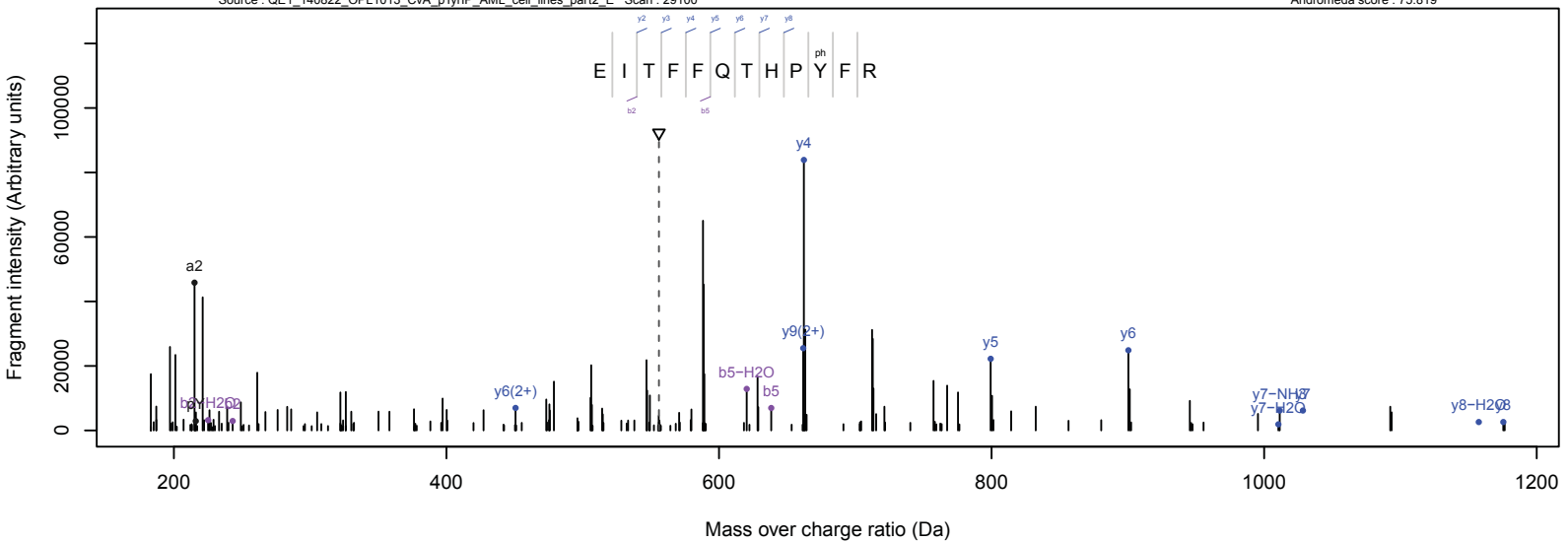
Andromeda score : 75.911



EITFFQTHPY(ph)FR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 29100

Andromeda score : 75.819

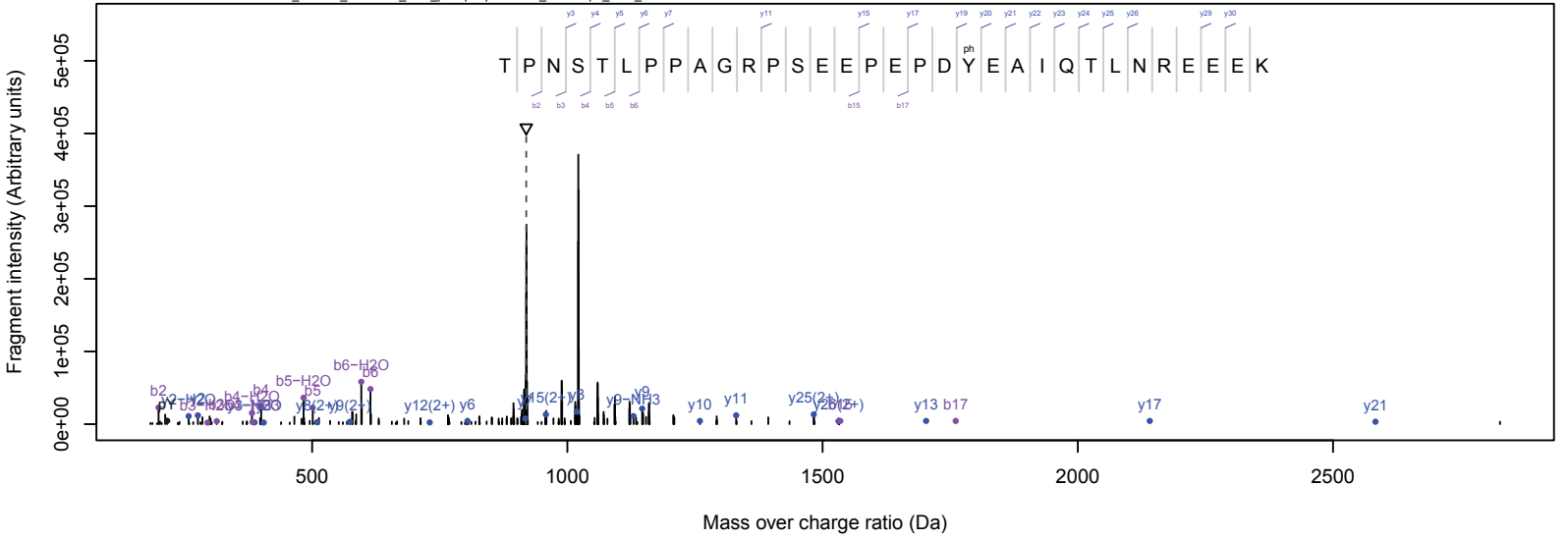




TPNSTLPPAGRPSEEPEDYEAIQTLNREEEK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28932

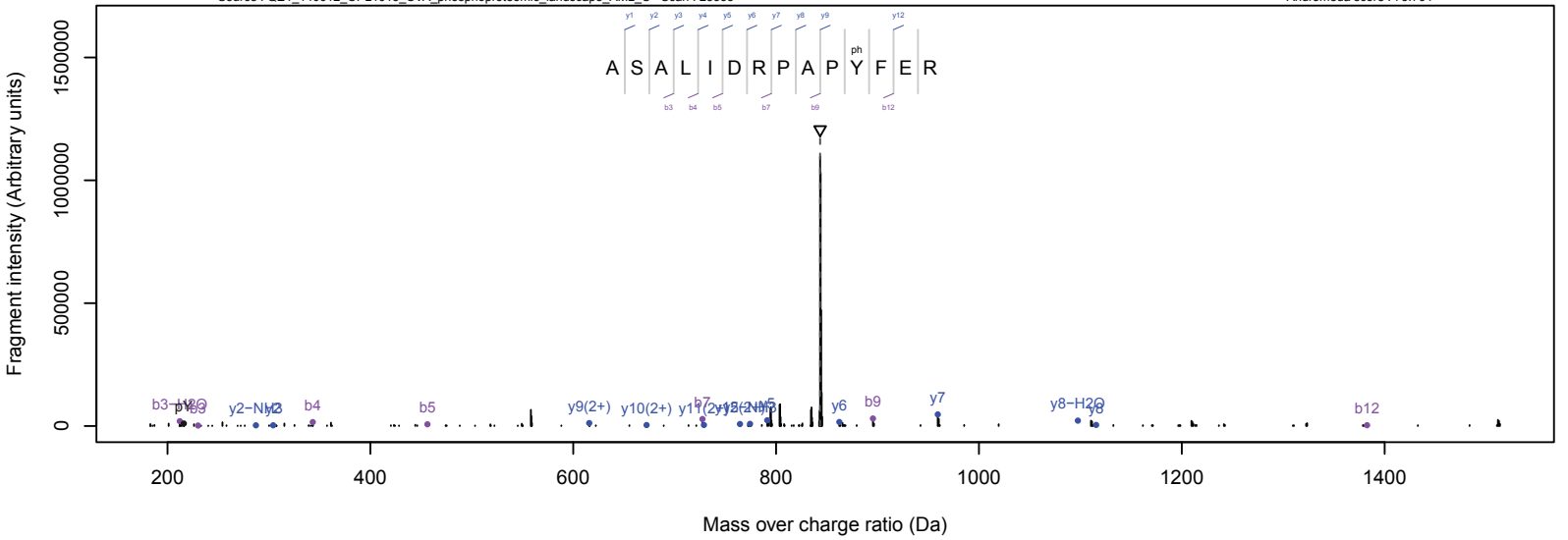
Andromeda score : 75.766



ASALIDRPAPYFER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25335

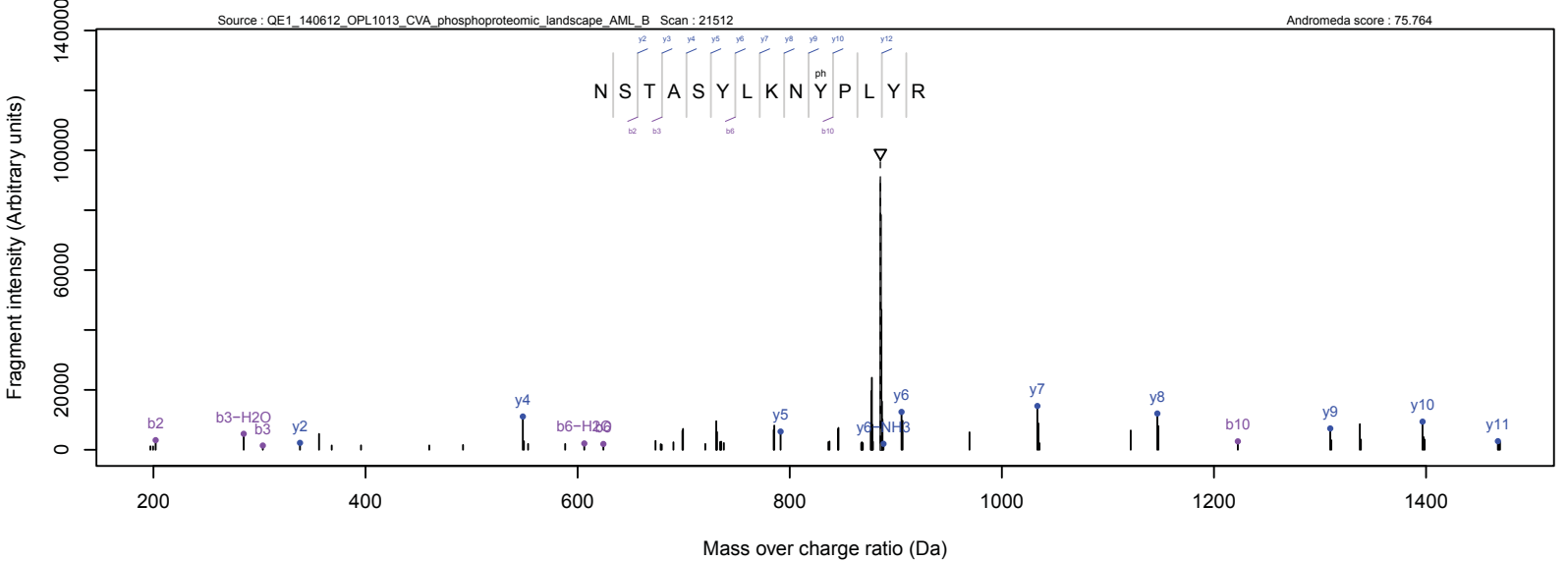
Andromeda score : 75.764



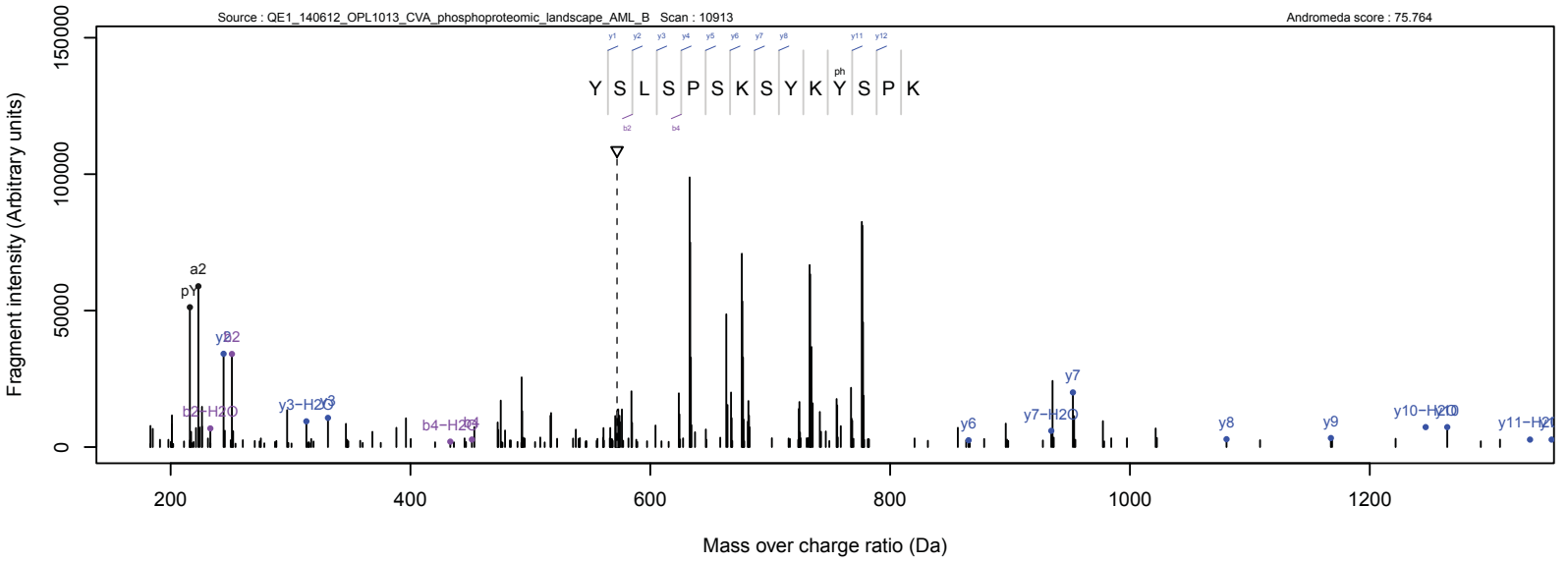
NSTASYLKNYPLYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21512

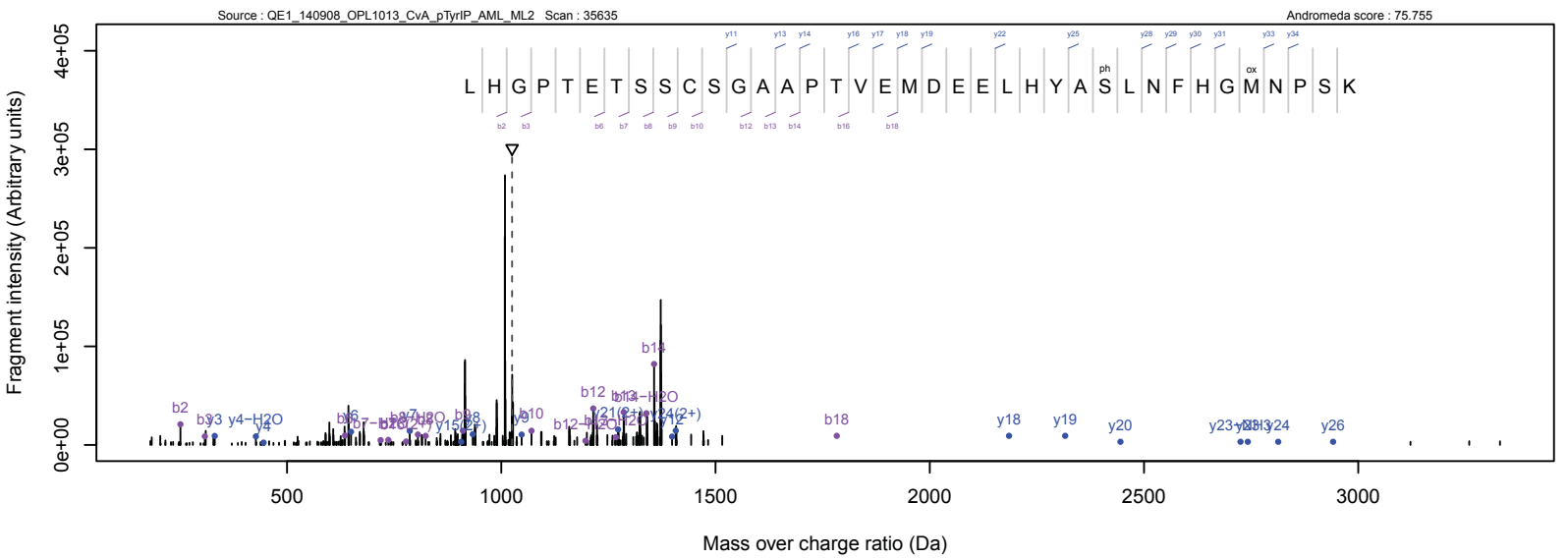
Andromeda score : 75.764



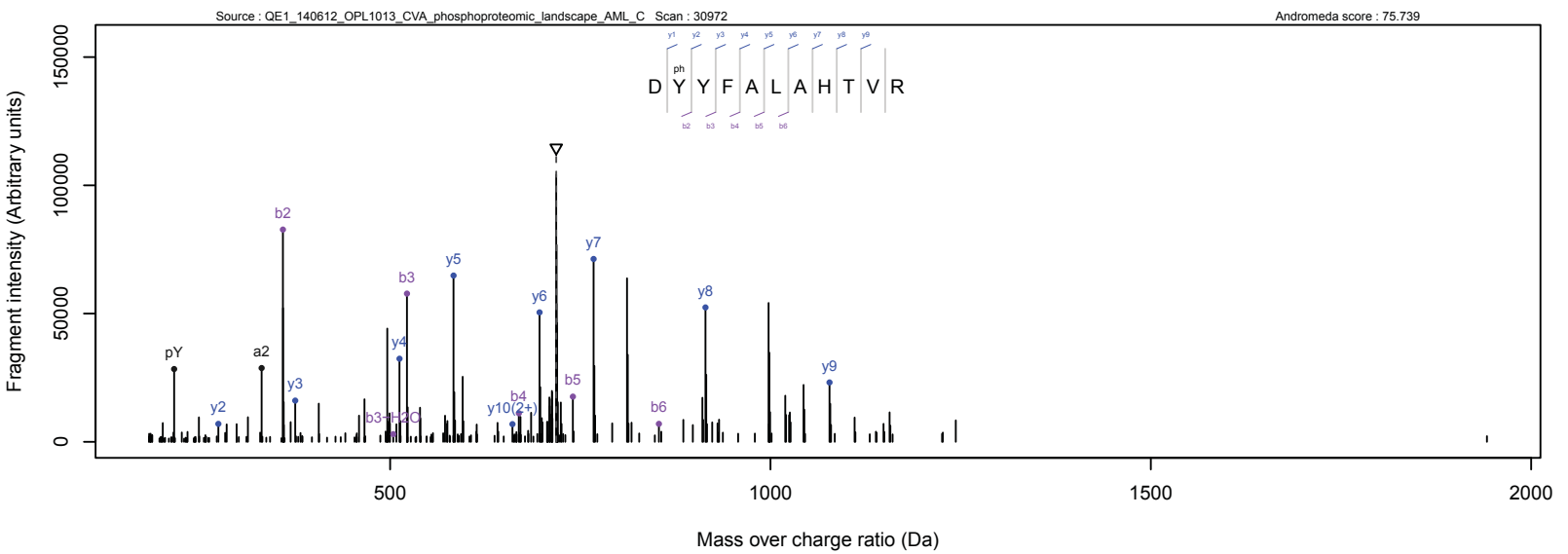
YLSLSPKSYKY(ph)SPK-3+



LHGPTETSSCSGAAPTVEEMDEELHYAS(ph)LNFHGM(ox)NPSK-4+



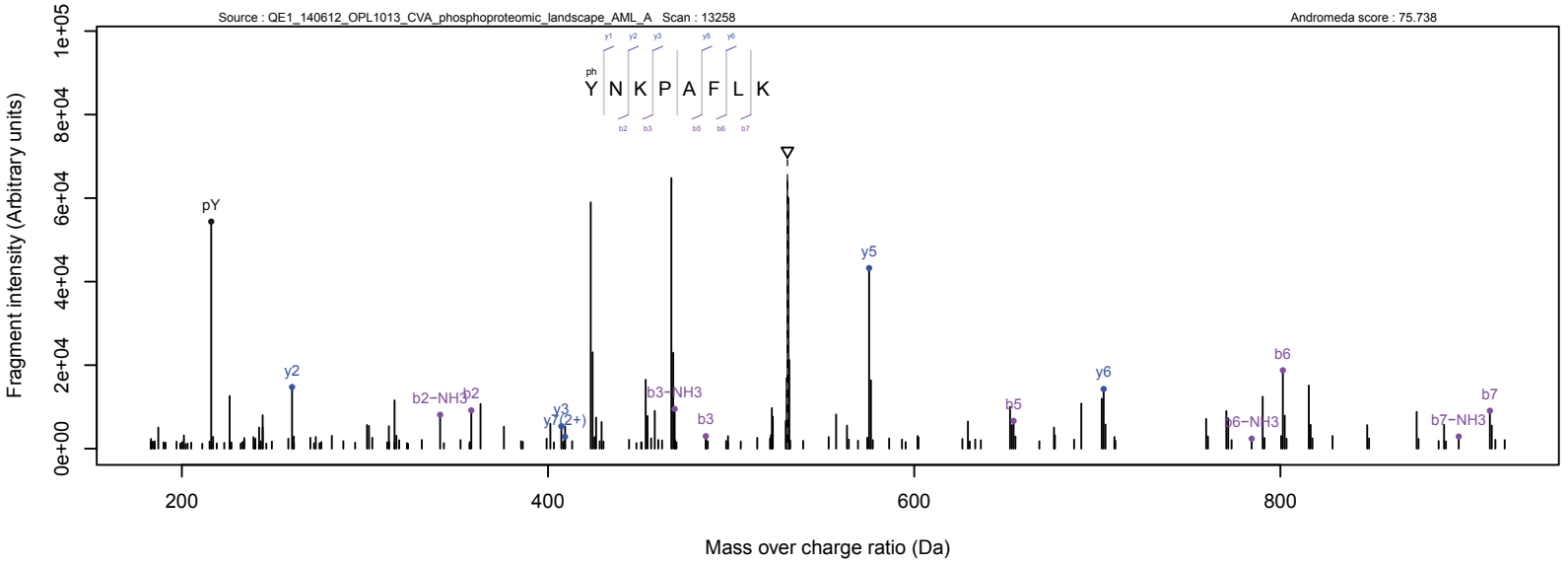
DY(ph)YFALAHTVR-2+



Y(ph)NKPAFLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 13258

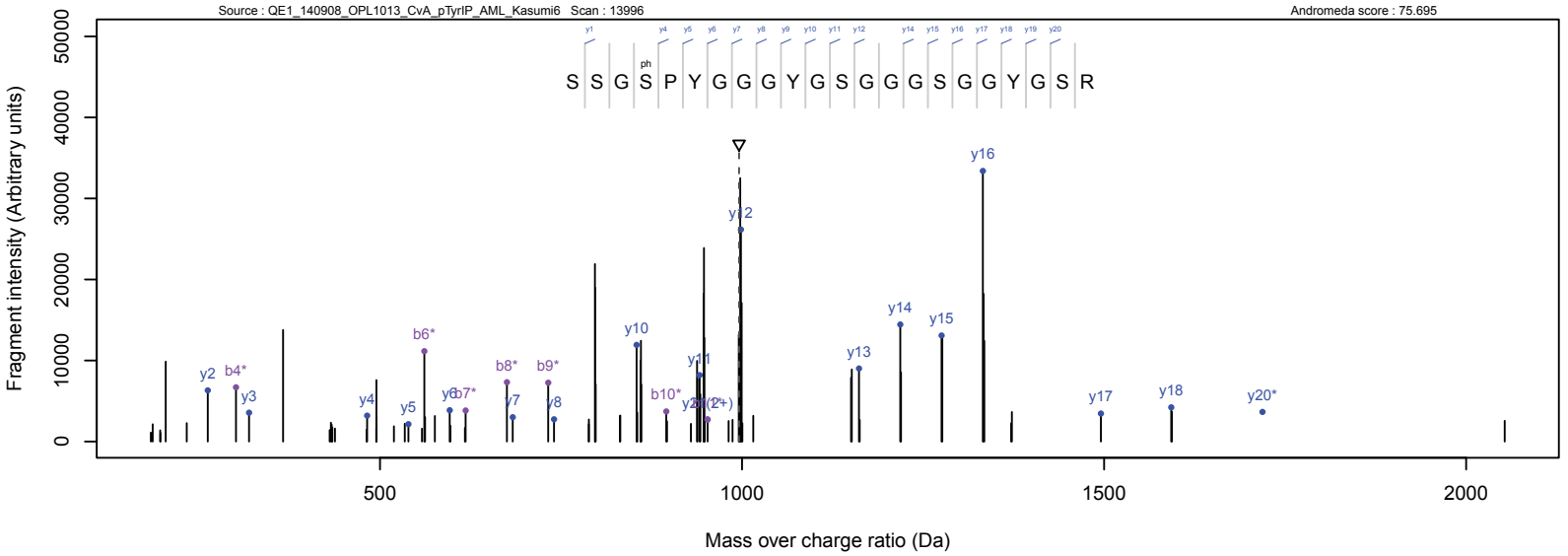
Andromeda score : 75.738



SSGS(ph)PYGGGYSGGGGYGSR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 13996

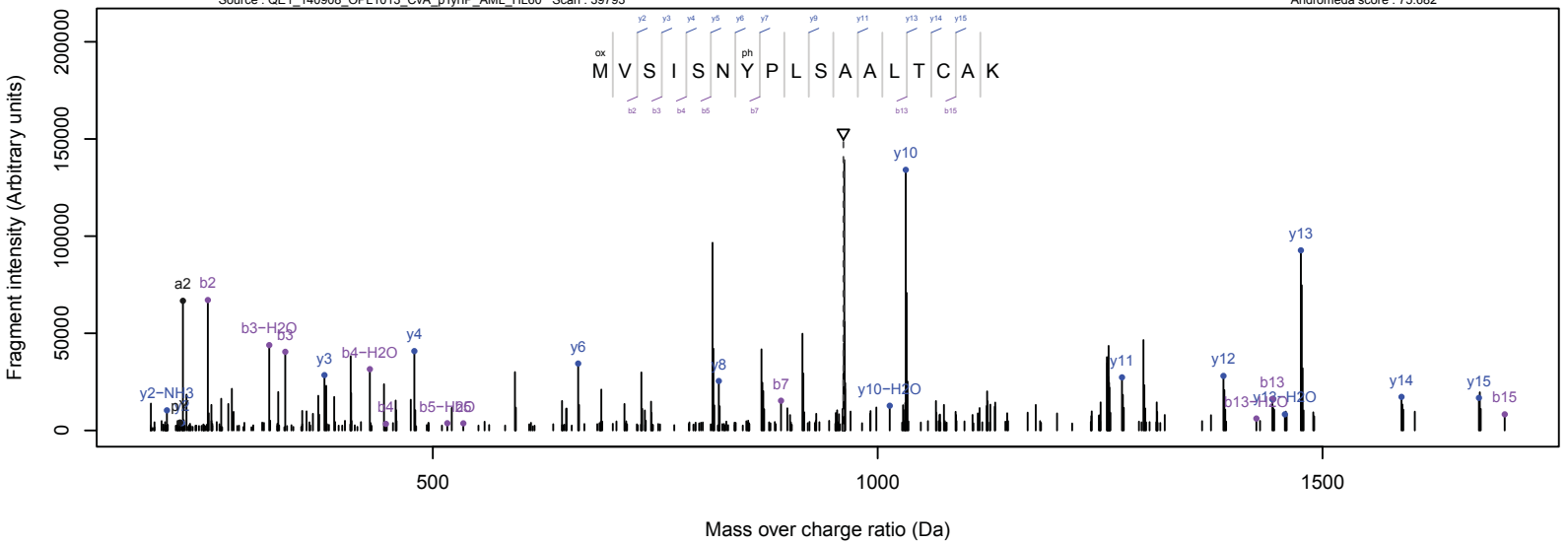
Andromeda score : 75.695



M(ox)VSISNY(ph)PLSAALTC AK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 39793

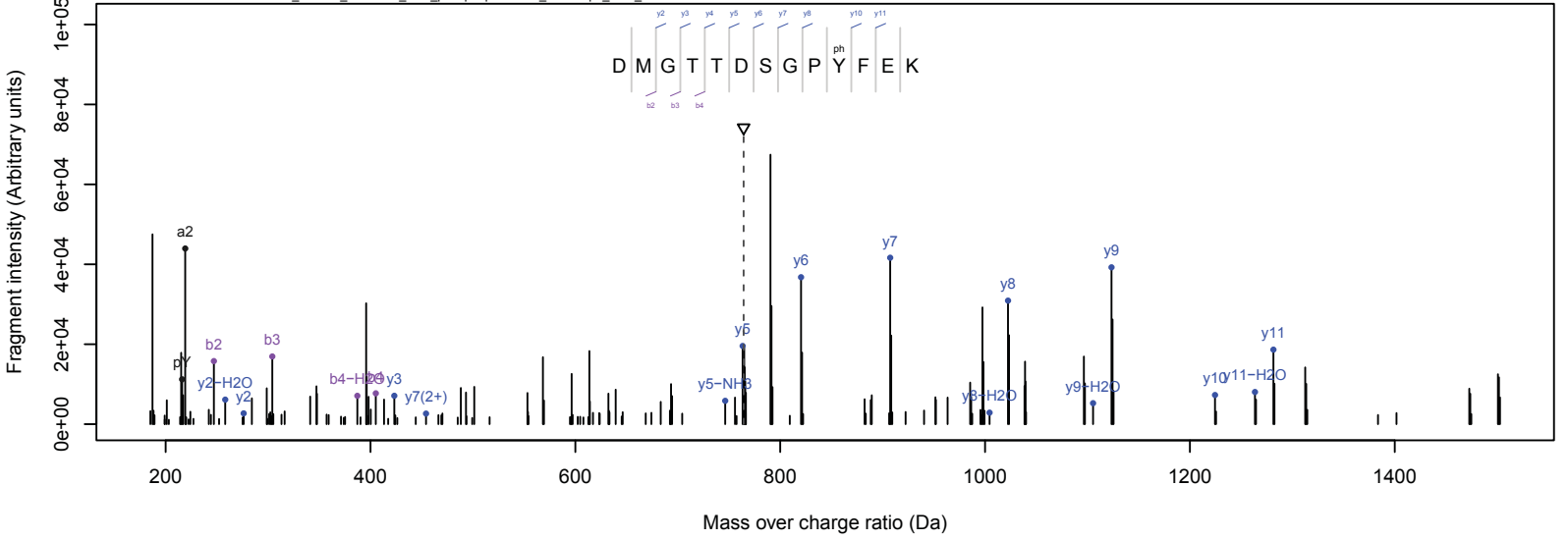
Andromeda score : 75.682



**DMGTTDGPY(ph)FEK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17945

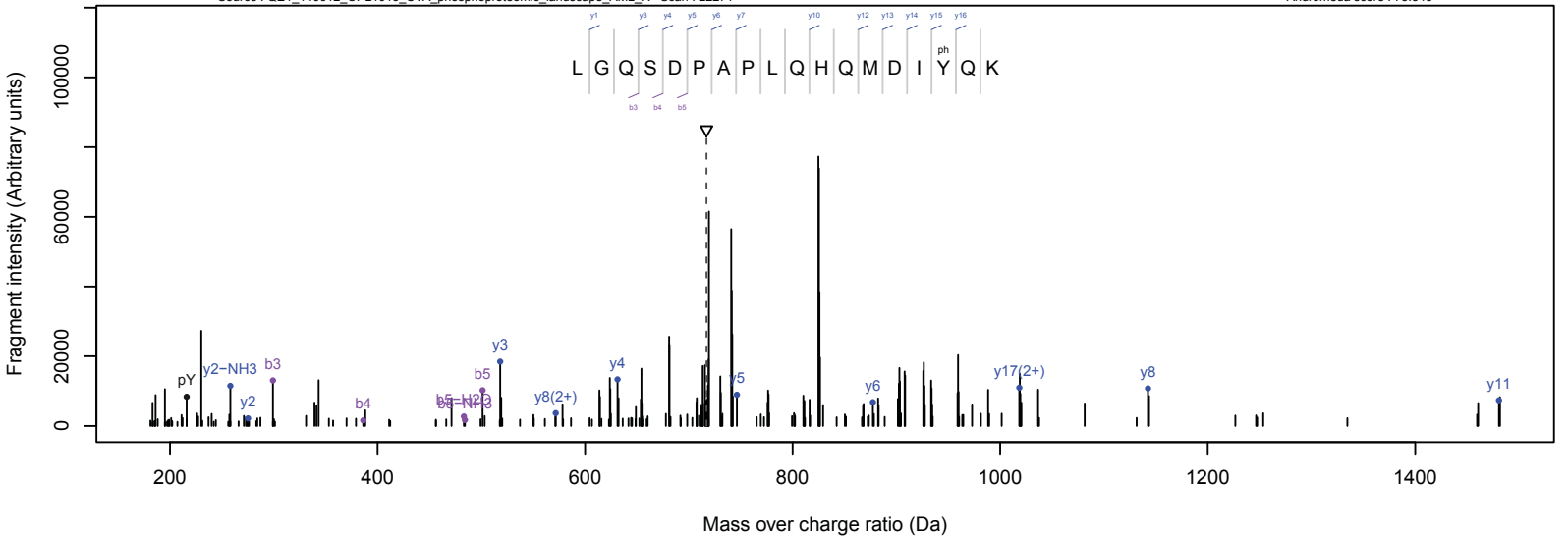
Andromeda score: 75.589



**LGQSDPAPLQHQMIDIYQK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 22271

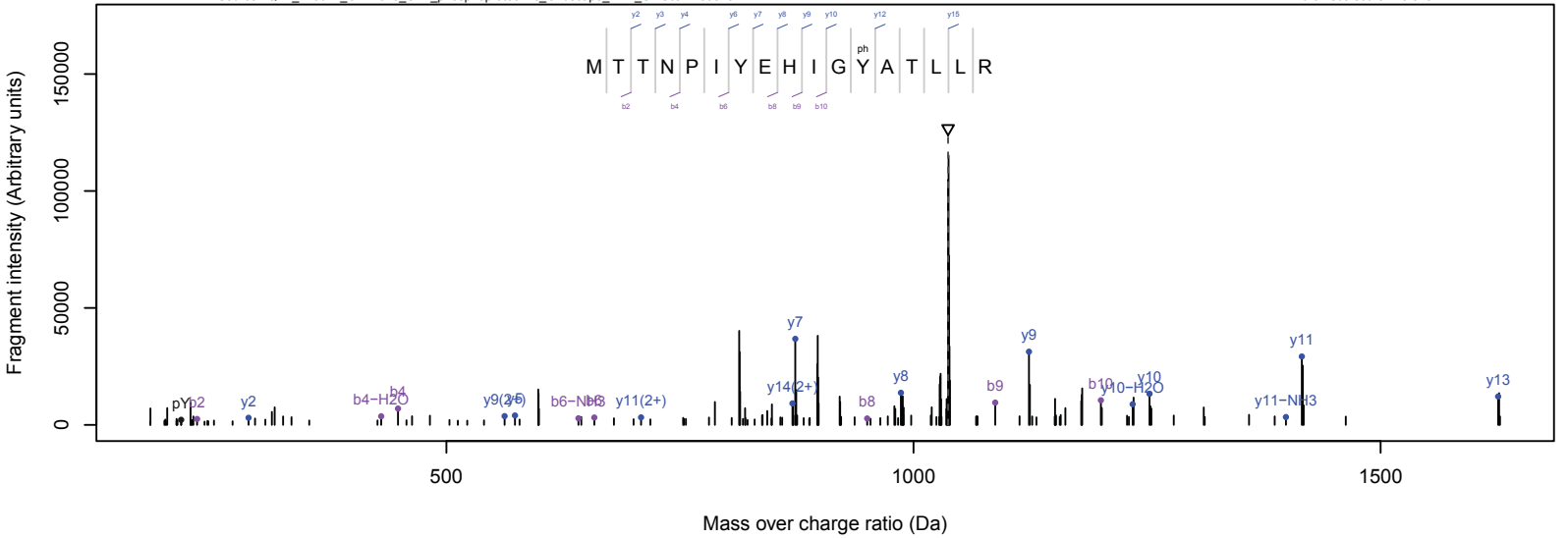
Andromeda score: 75.548



**MTTNIYEHIGYATLLR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 39528

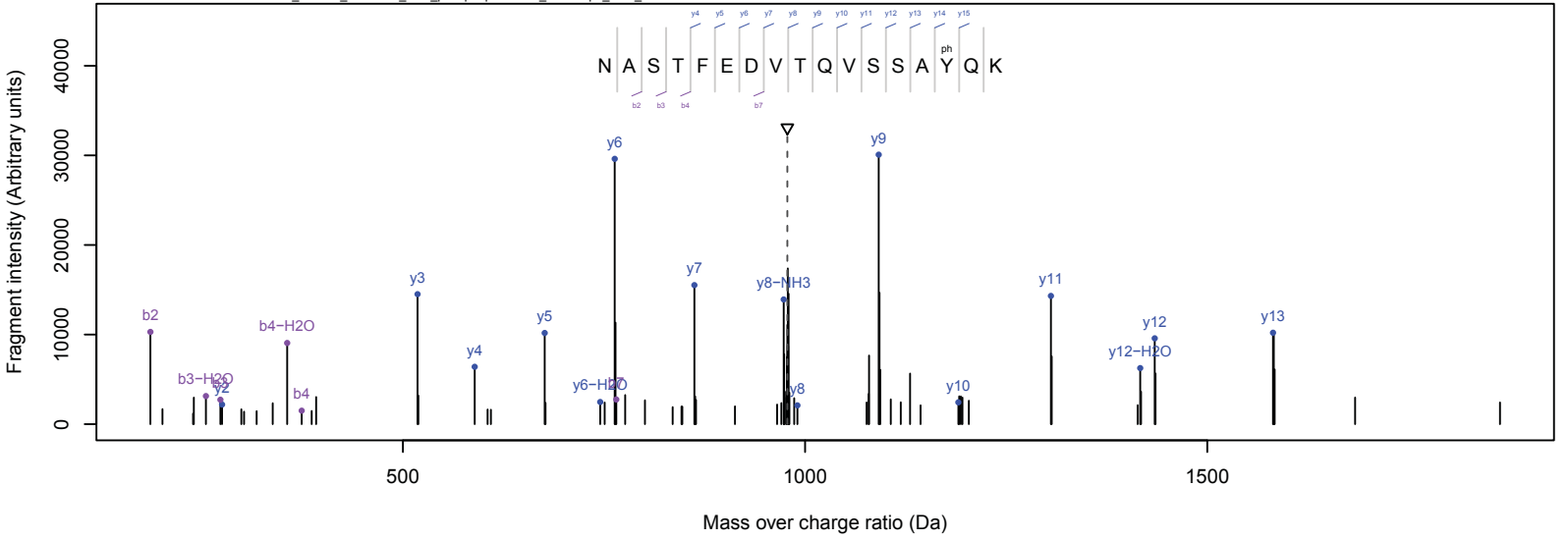
Andromeda score: 75.548



NASTFEDVTQVSSAYQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22546

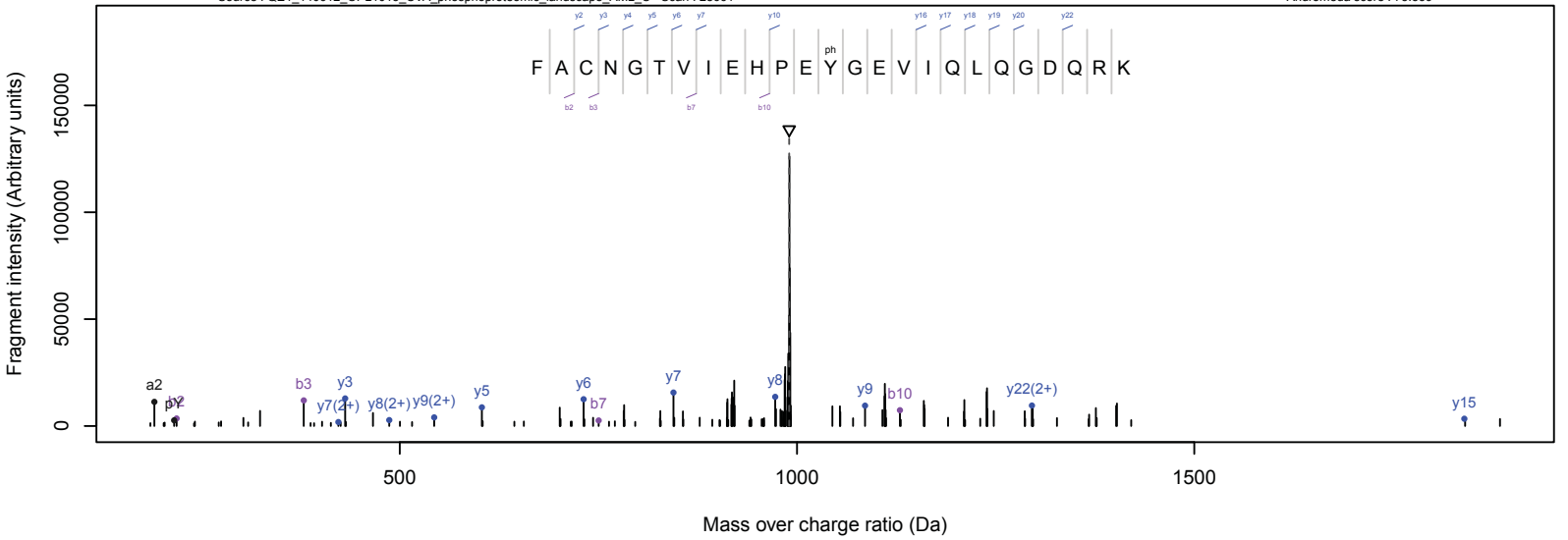
Andromeda score : 75.548



FACNGTVIEHPEYGEVILQLQGDRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28661

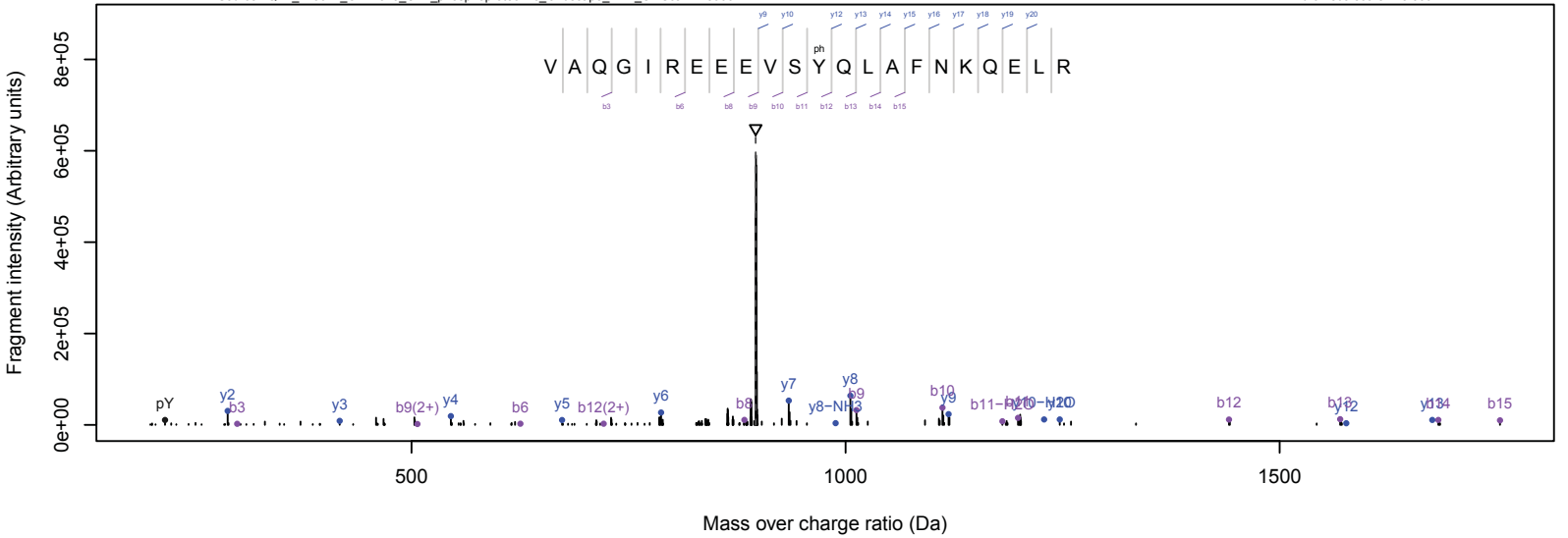
Andromeda score : 75.539



VAQGIREEEVS YQLAFNKQELR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28906

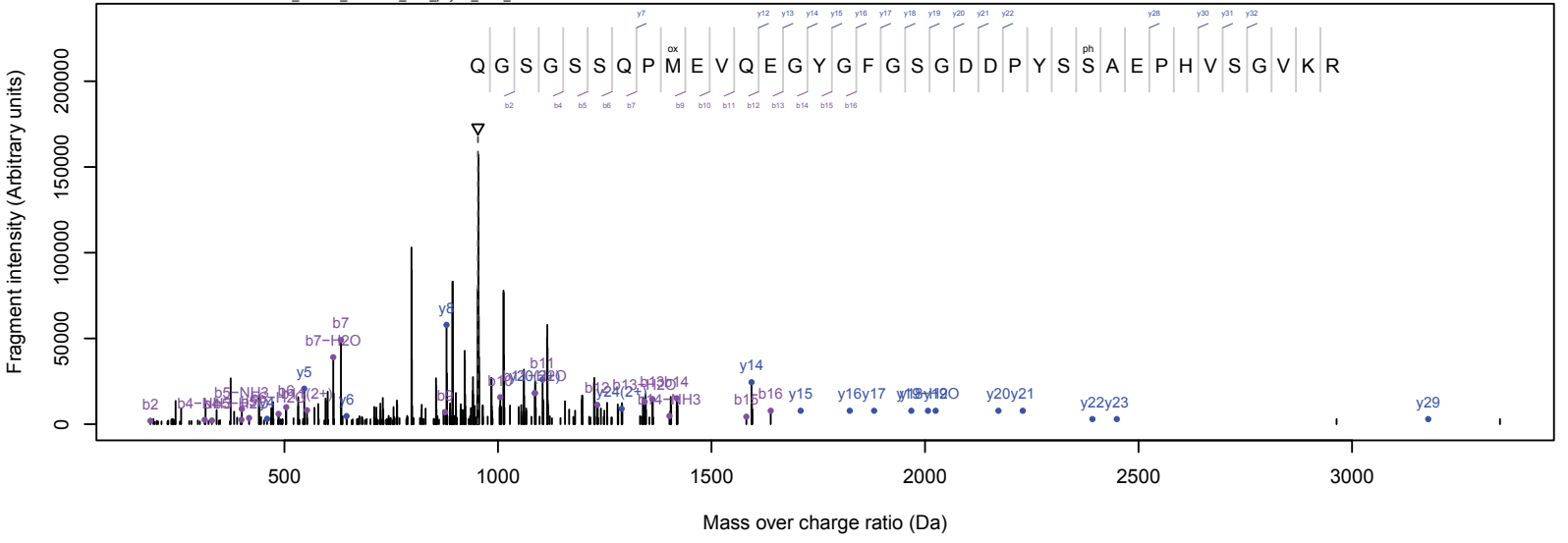
Andromeda score : 75.539



**QGSSSQPM(ox)EVQEGYFGSGDDPYSS(ph)AEPHVSQVKR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 24653

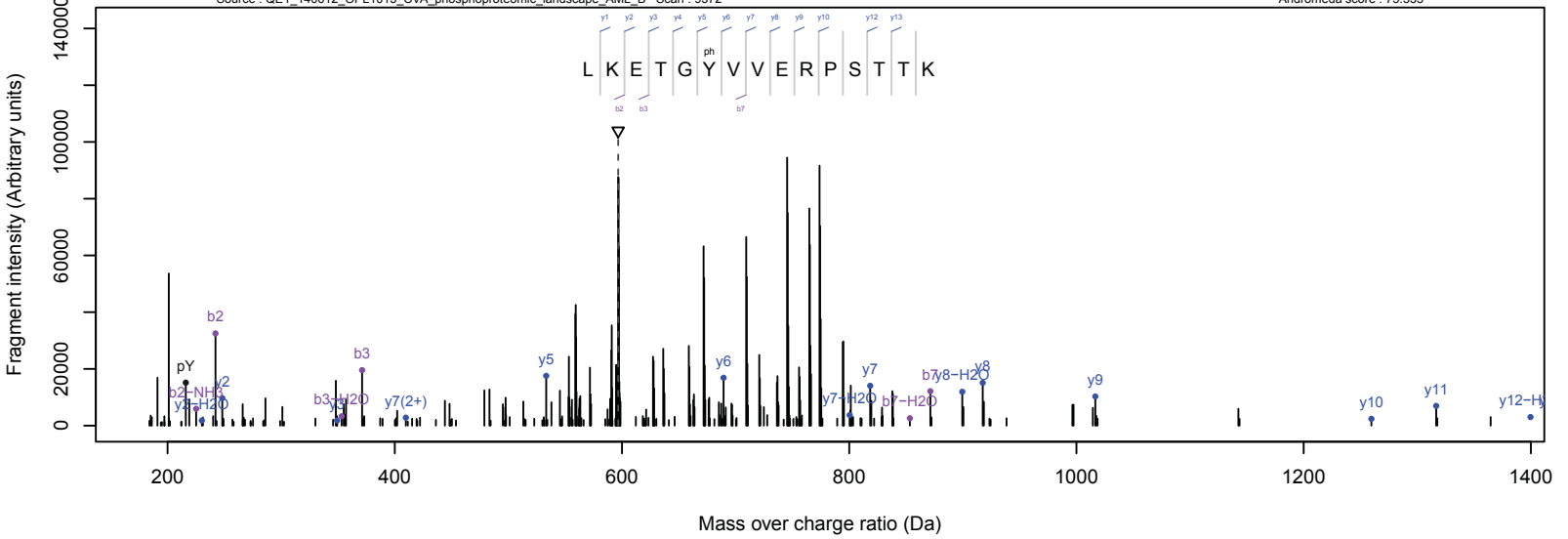
Andromeda score : 75.536



**LKETGY(ph)VVERPSTTK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9372

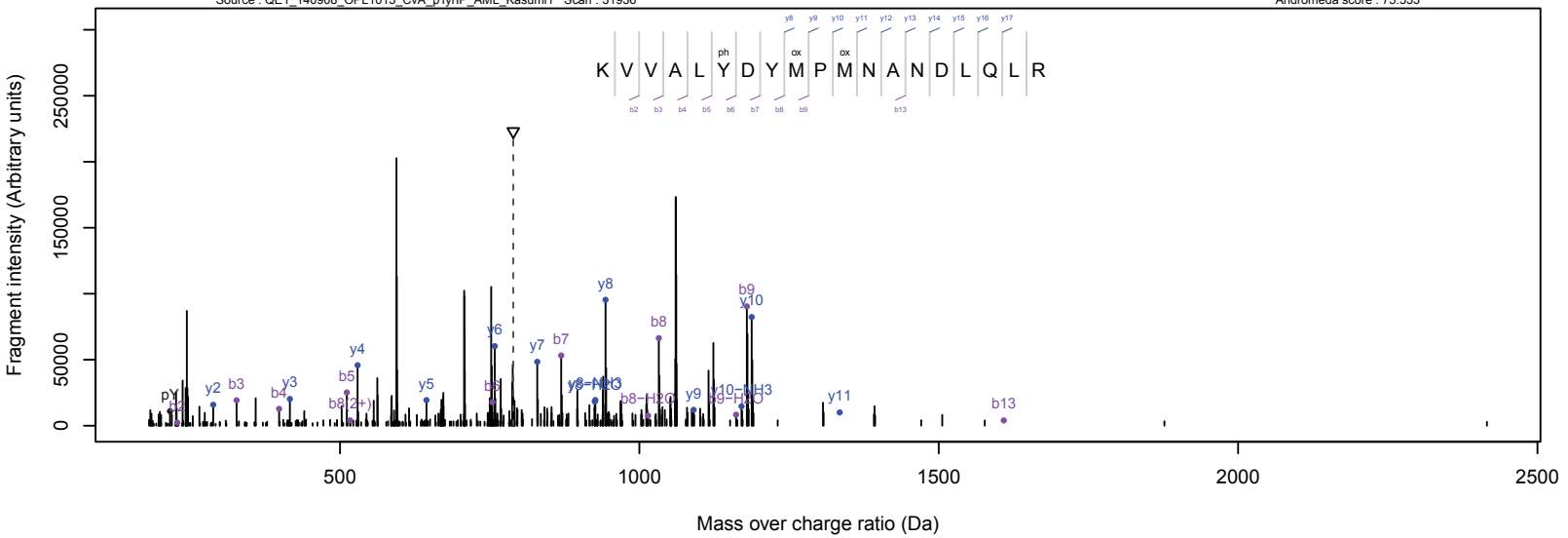
Andromeda score : 75.535



**KVVALYDYM(ox)PM(ox)NANDLQLR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 31936

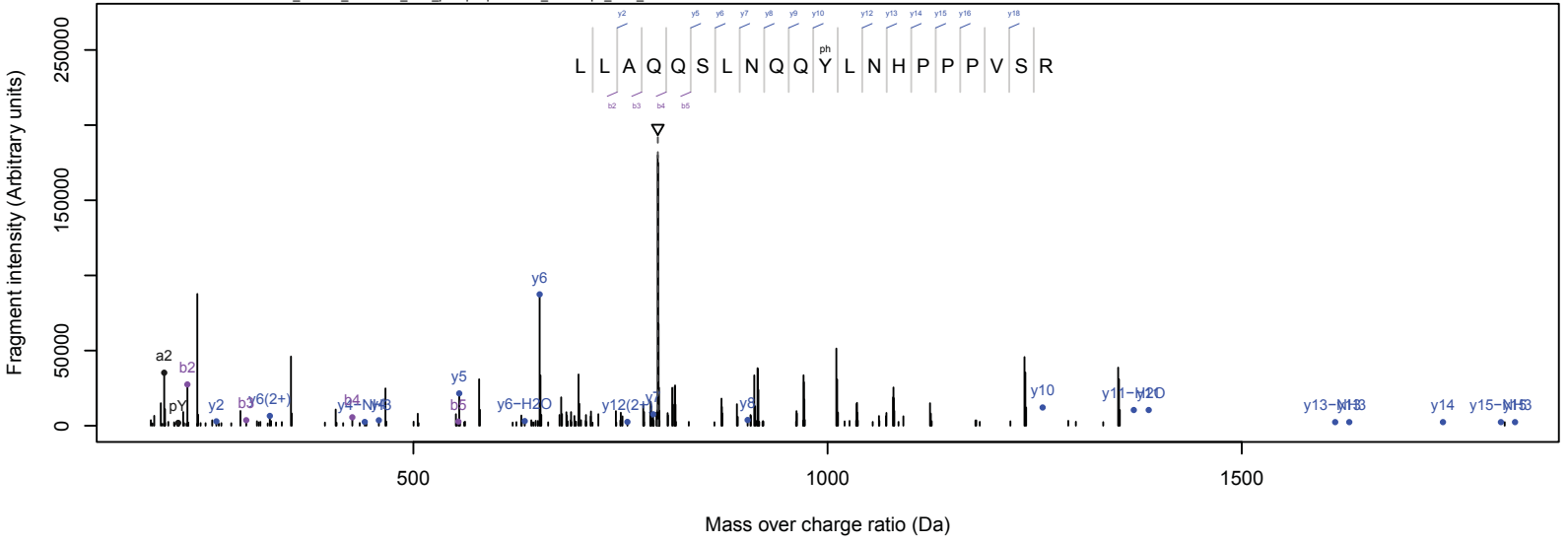
Andromeda score : 75.533



LLAQQSLNQQY(ph)LNHPPPVS<sup>R</sup>-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25351

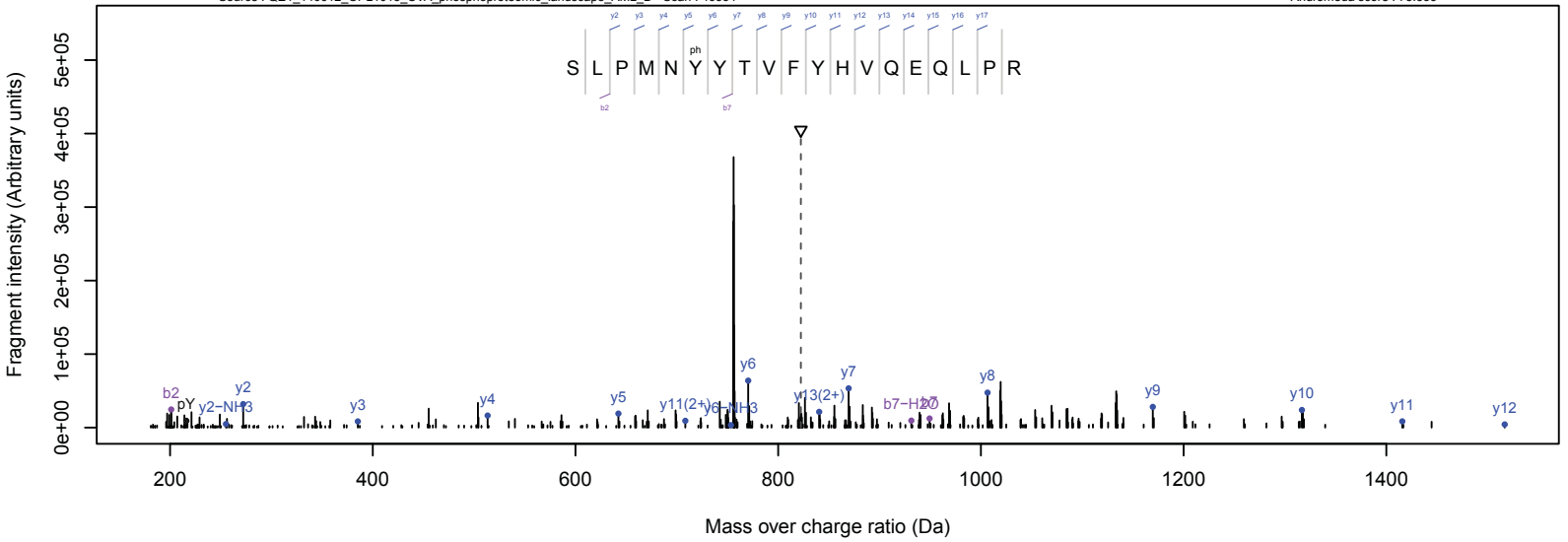
Andromeda score : 75.533



SLPMNY(ph)YTFYHVQEQLP<sup>R</sup>-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 46534

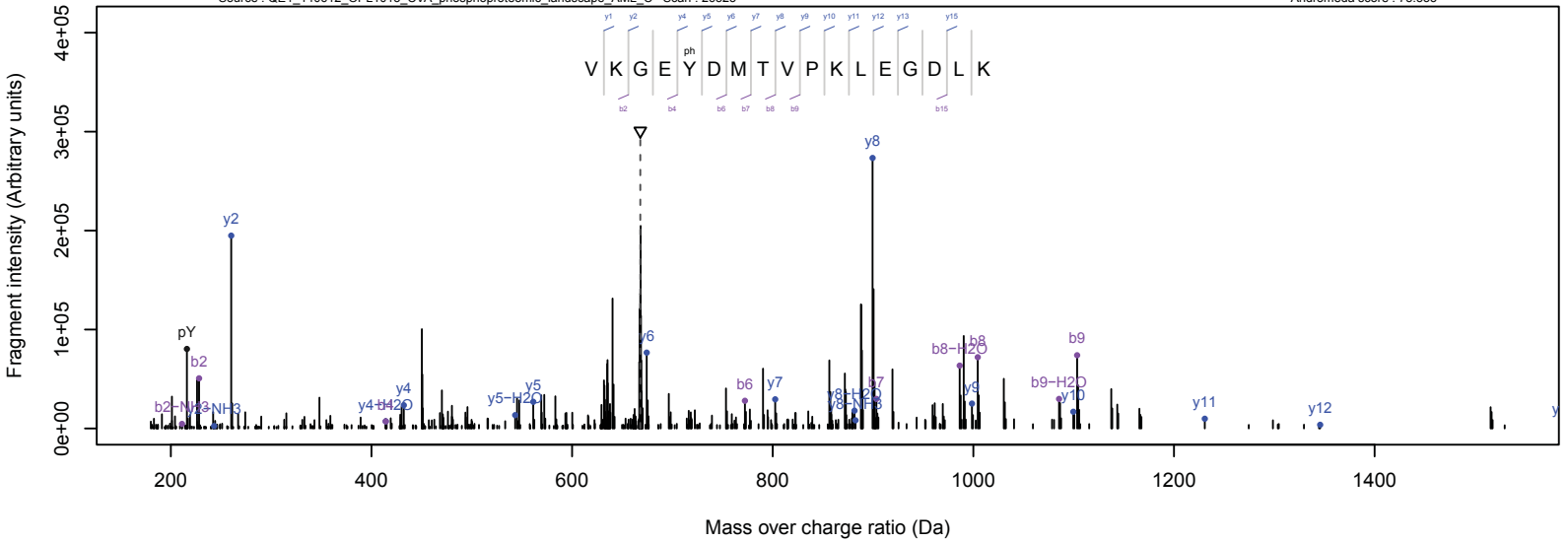
Andromeda score : 75.533



VKGEY(ph)DMTPKLEGLD<sup>LK</sup>-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26323

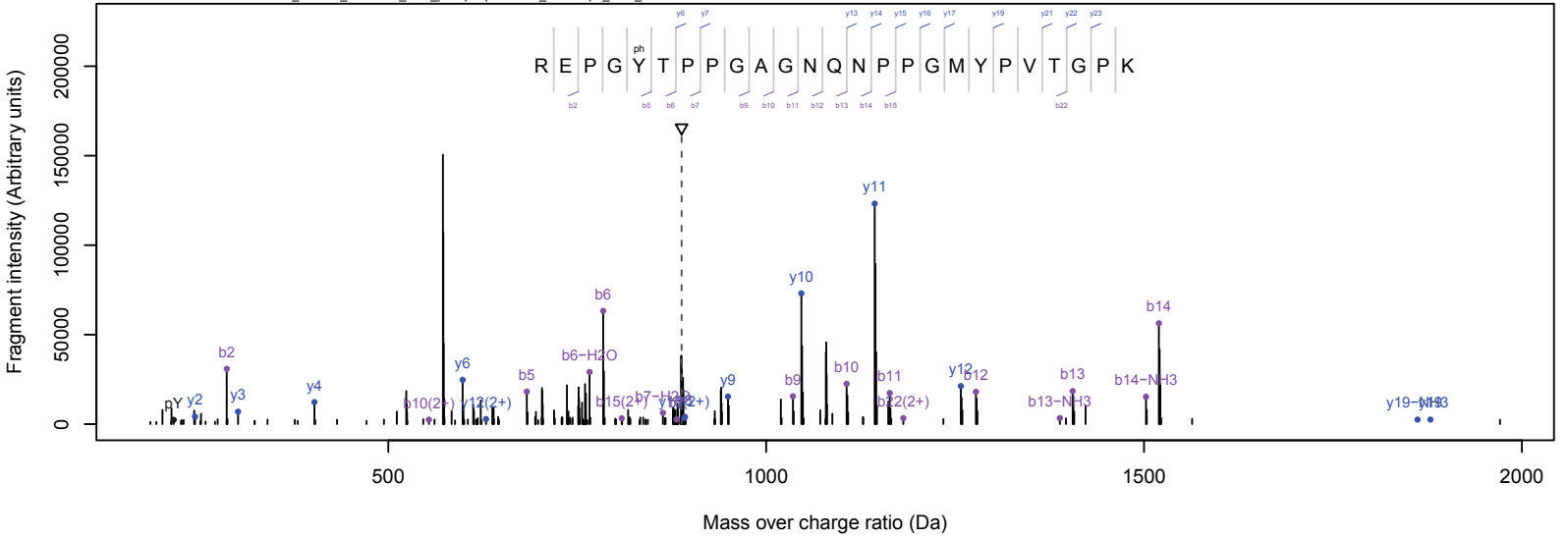
Andromeda score : 75.533



REPGY(ph)TPPGAGNQNP PGMY PVTGPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22537

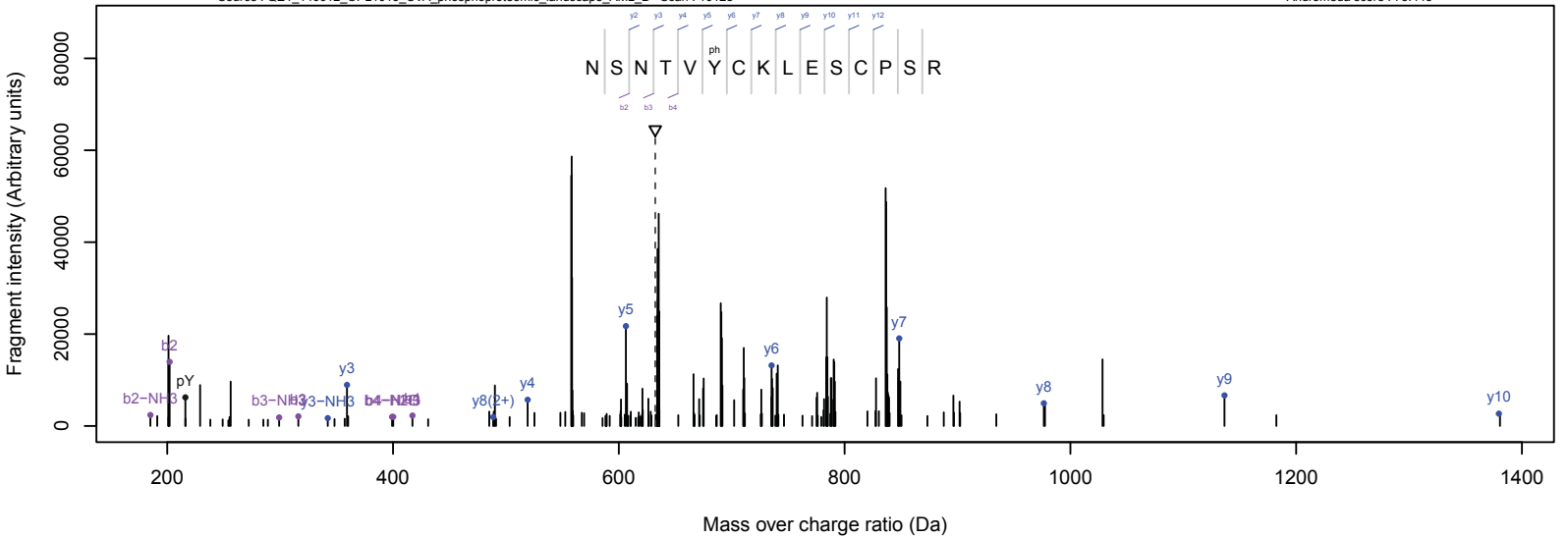
Andromeda score : 75.529



NSNTVY(ph)CKLESCPSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10128

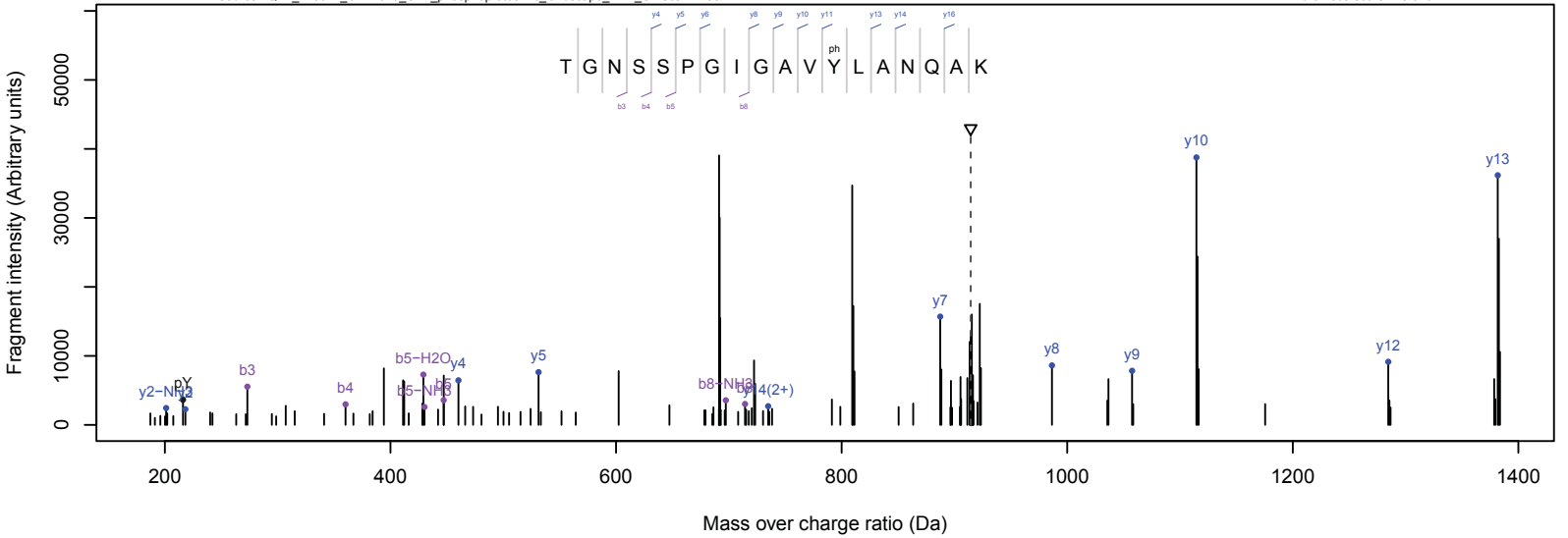
Andromeda score : 75.445



TGNSSPGAVY(ph)LANQAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25811

Andromeda score : 75.376

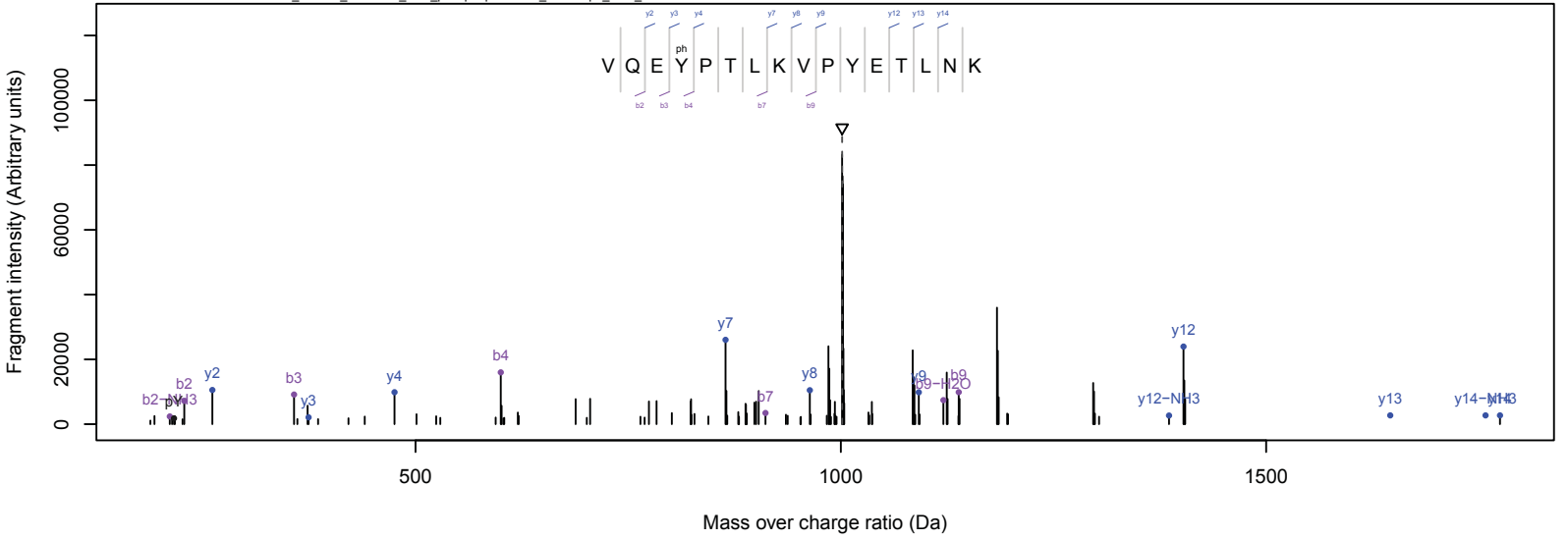




VQEY(ph)PTLKVPYETLNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29668

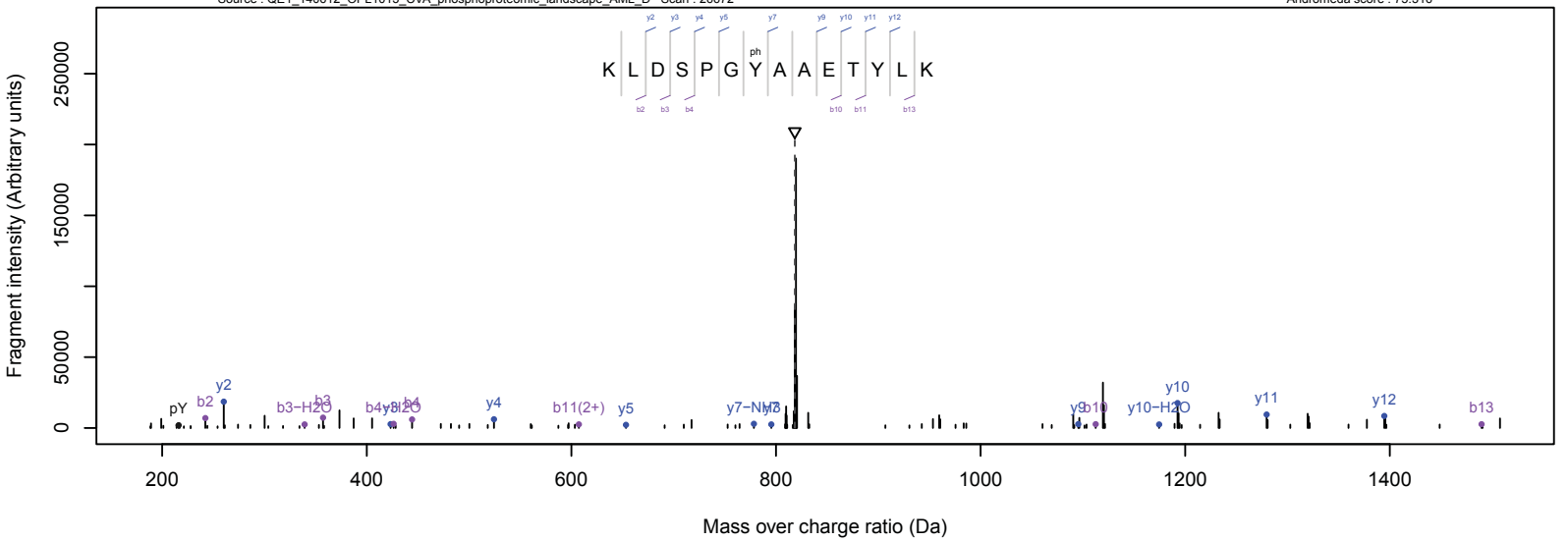
Andromeda score : 75.376



KLDSPG(ph)AAETYLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 26672

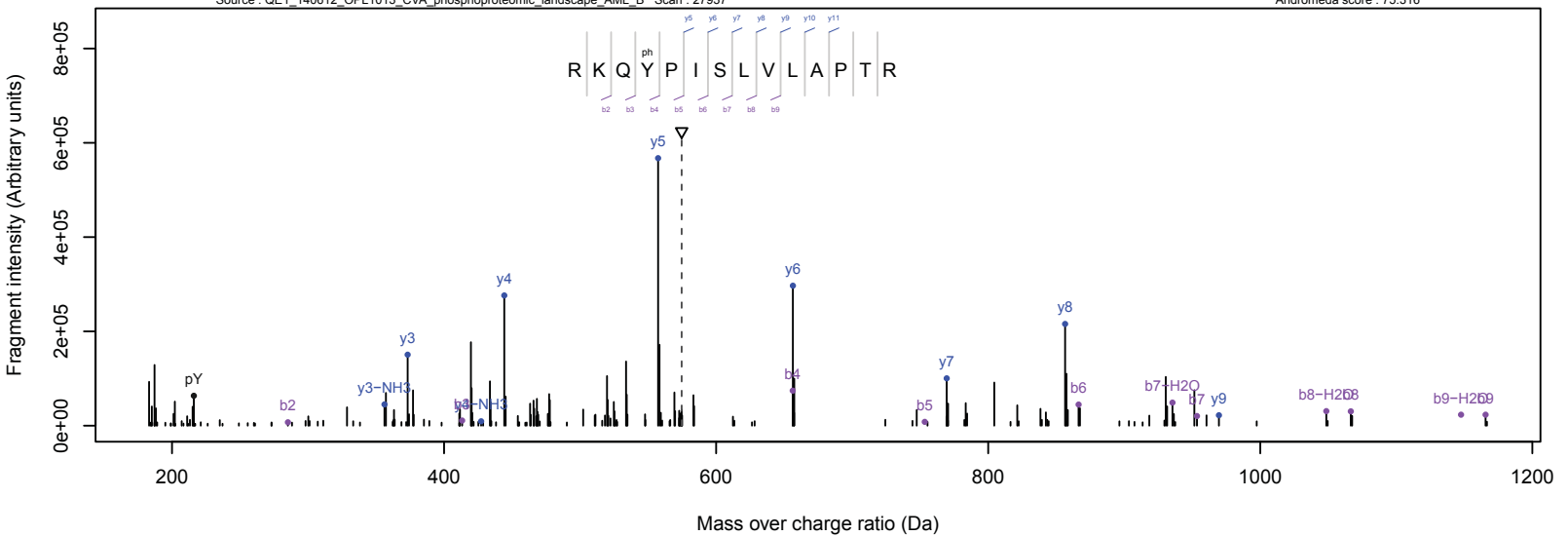
Andromeda score : 75.316



RKQY(ph)PISLV LAPTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27937

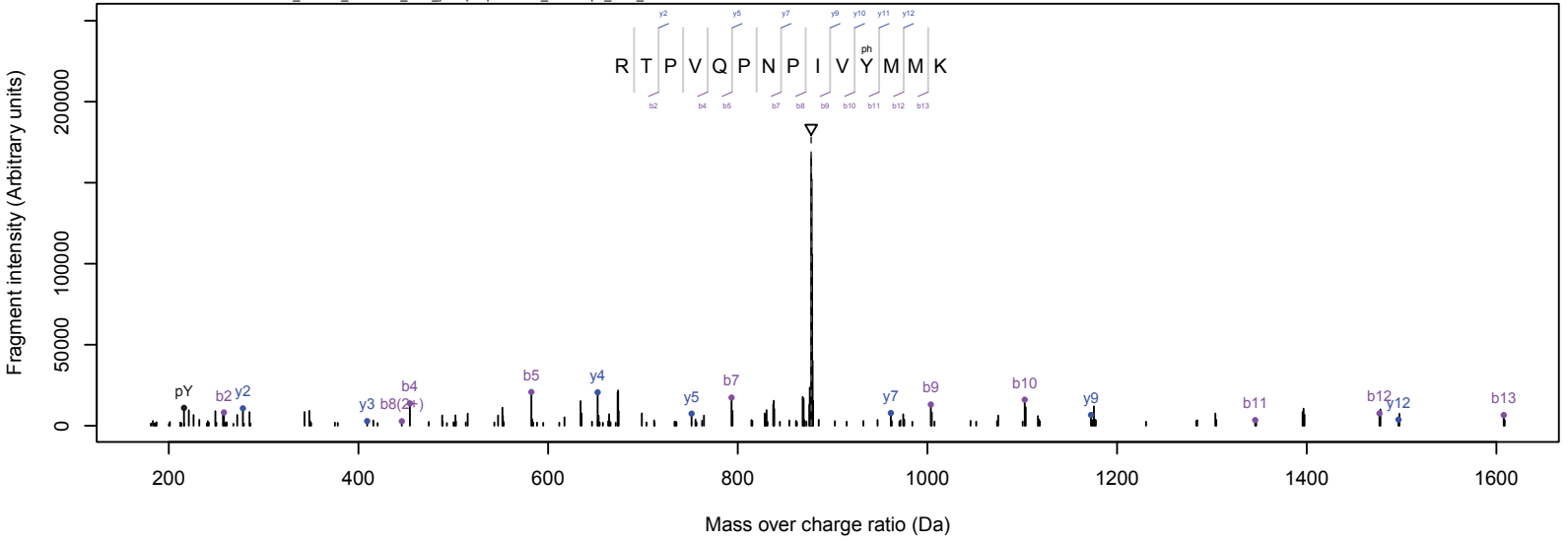
Andromeda score : 75.316



RTPVQPNPIVY(ph)MMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34904

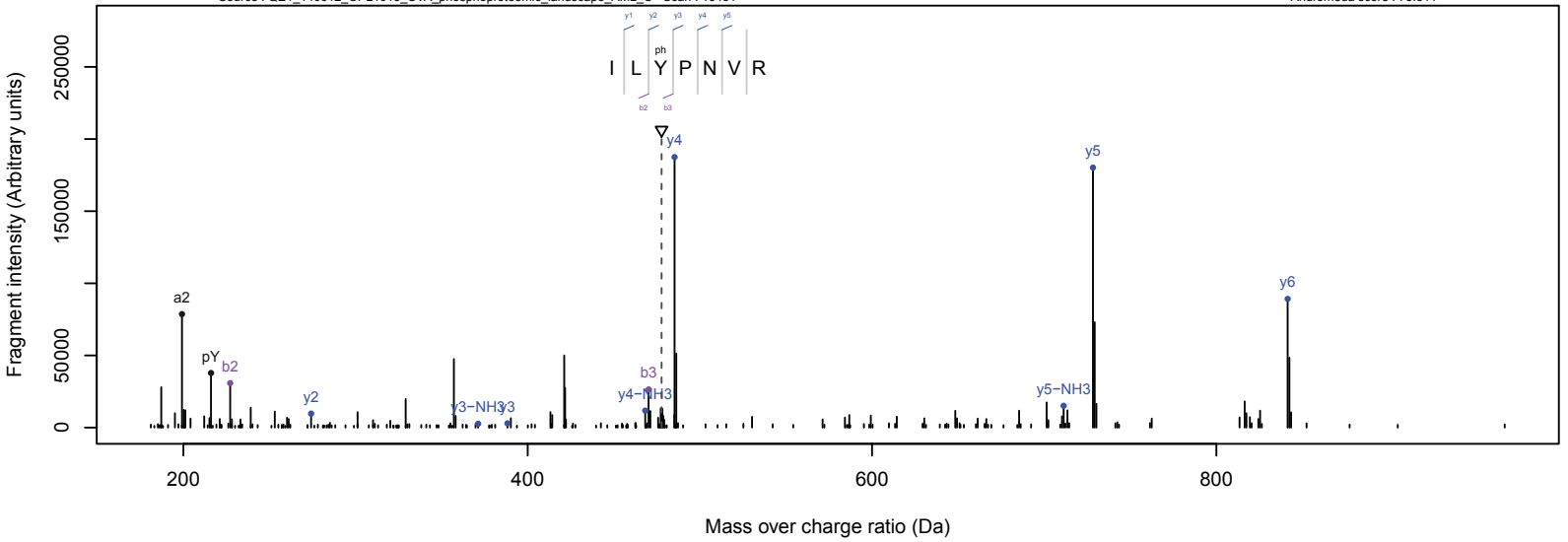
Andromeda score : 75.316



ILY(ph)PNVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16451

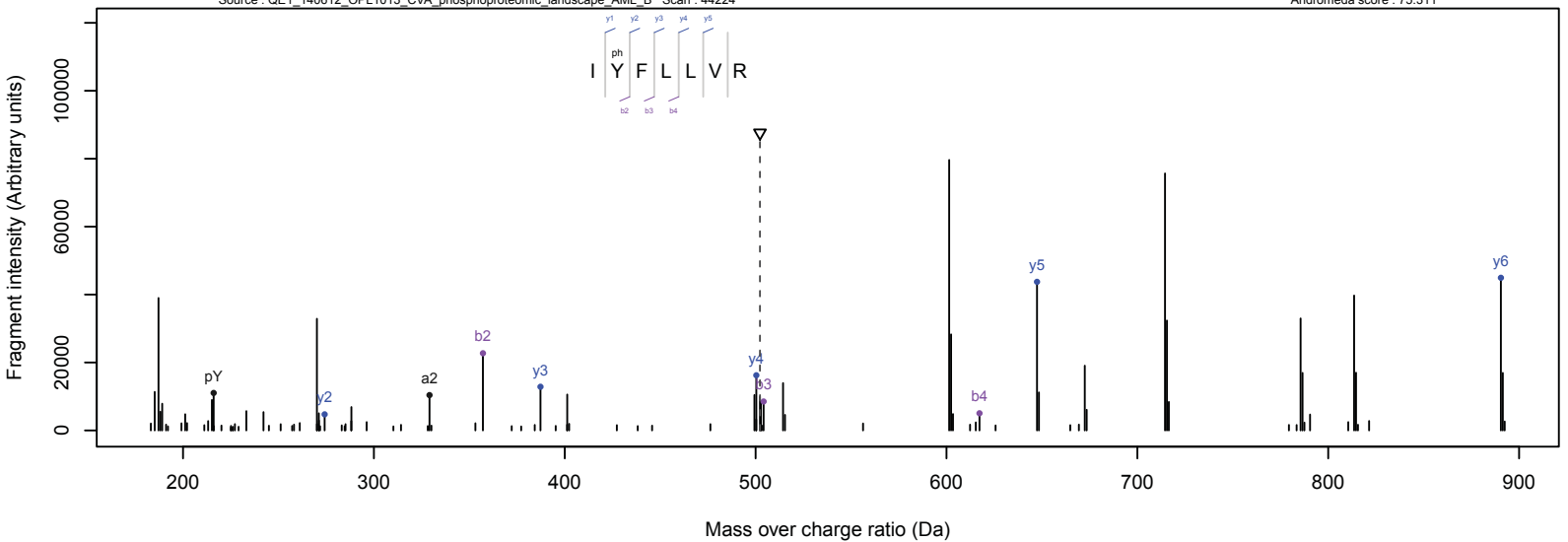
Andromeda score : 75.311



IY(ph)FLLVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44224

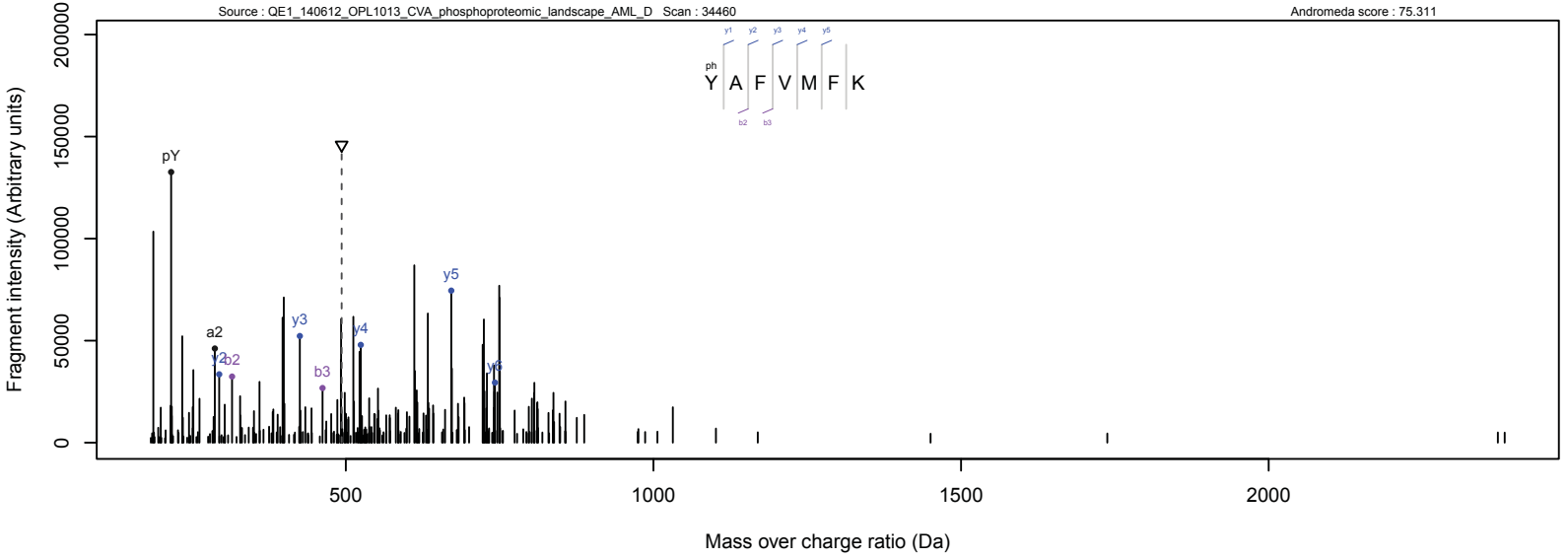
Andromeda score : 75.311



**Y(ph)AFVMFK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 34460

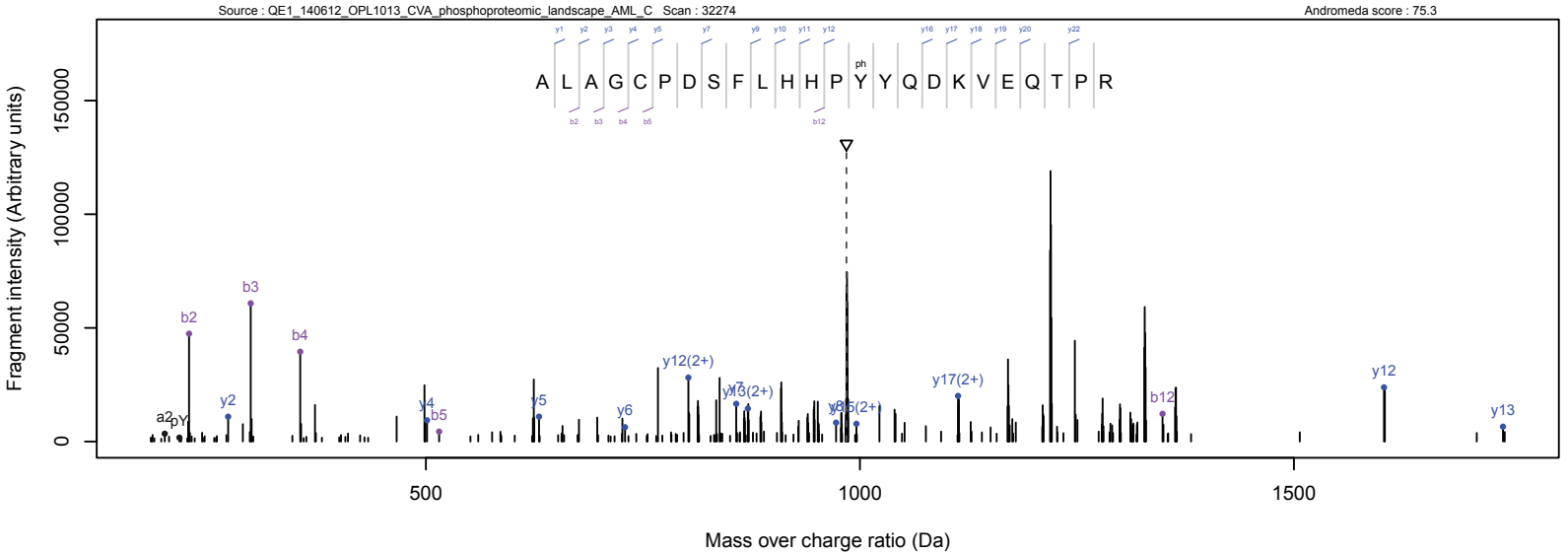
Andromeda score : 75.311



**(ac)ALAGCPDSFLHHPY(ph)YQDKVEQTPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32274

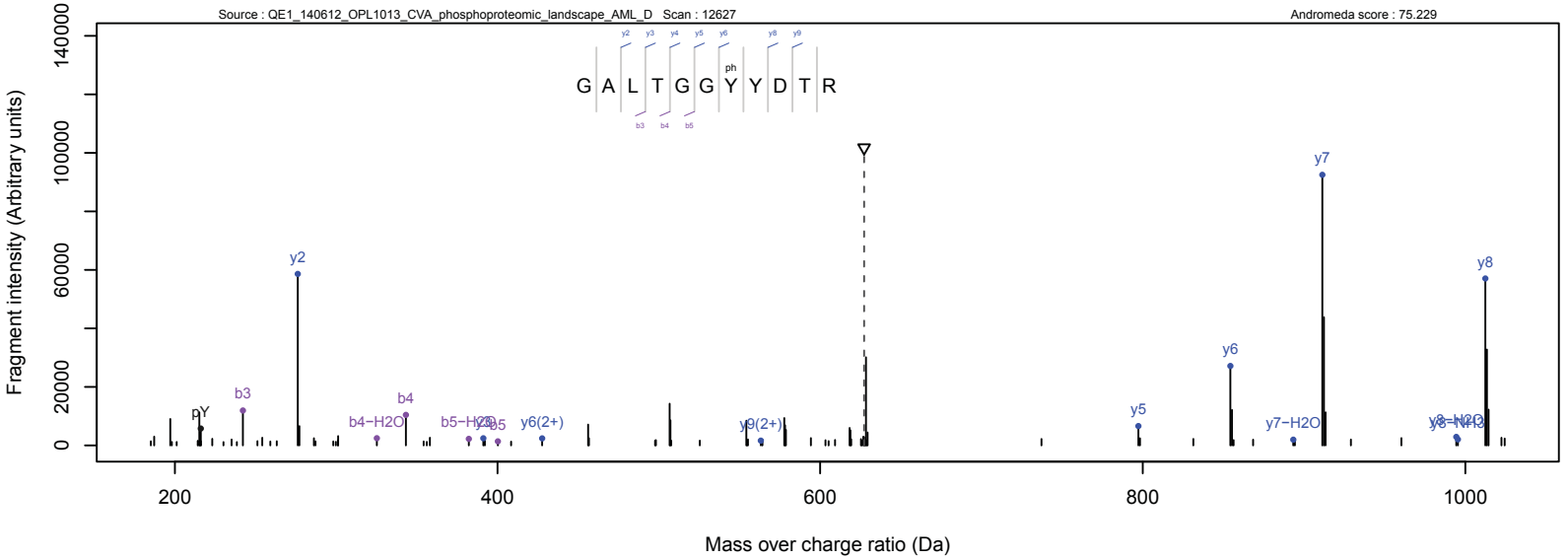
Andromeda score : 75.3



**GALGGY(ph)YDTR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12627

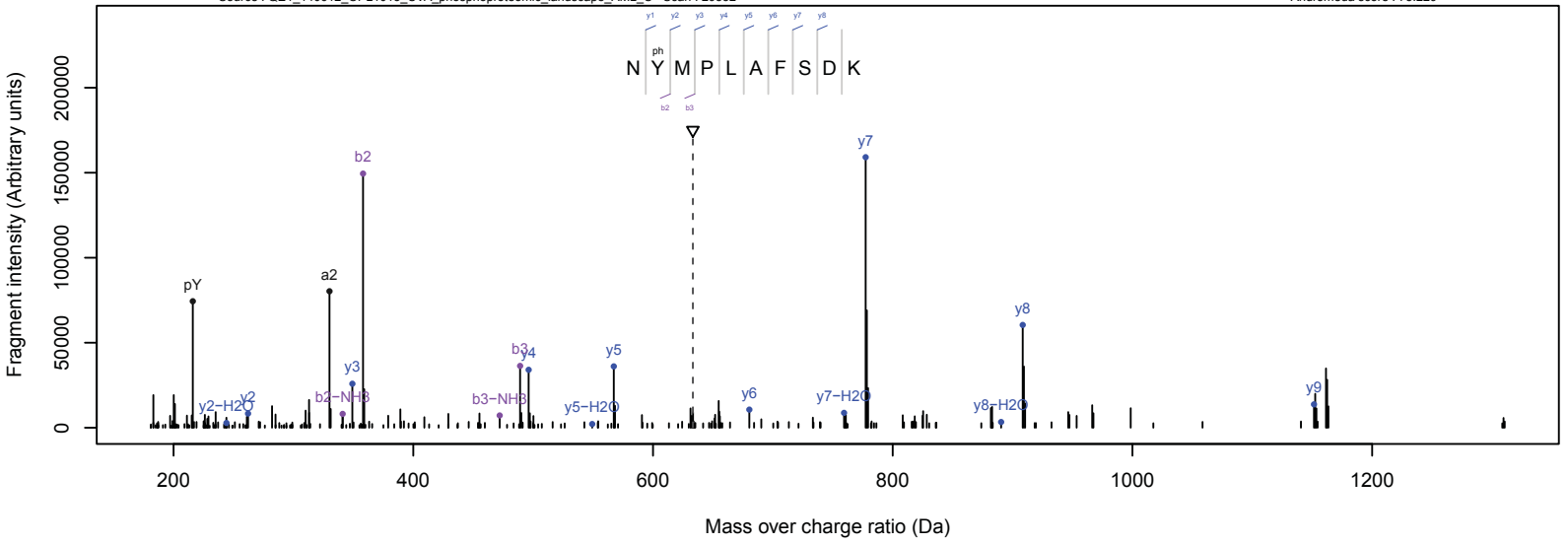
Andromeda score : 75.229



NY(ph)MPLAFSDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29532

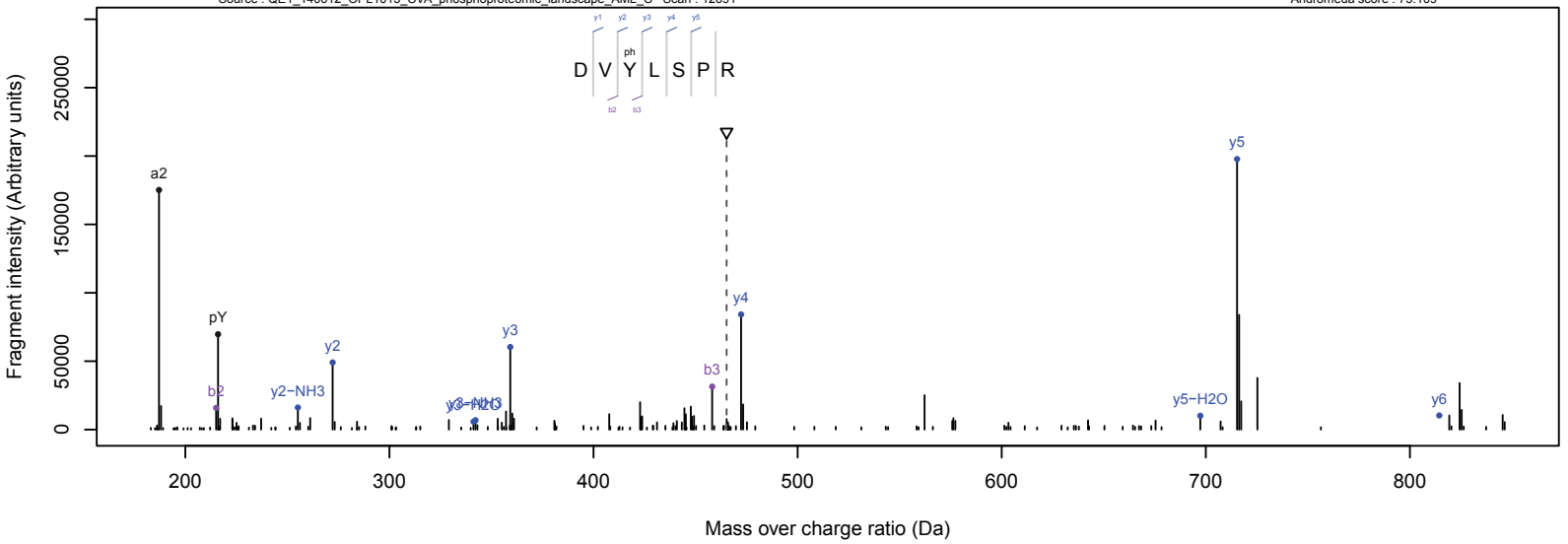
Andromeda score : 75.229



DVY(ph)LSPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12691

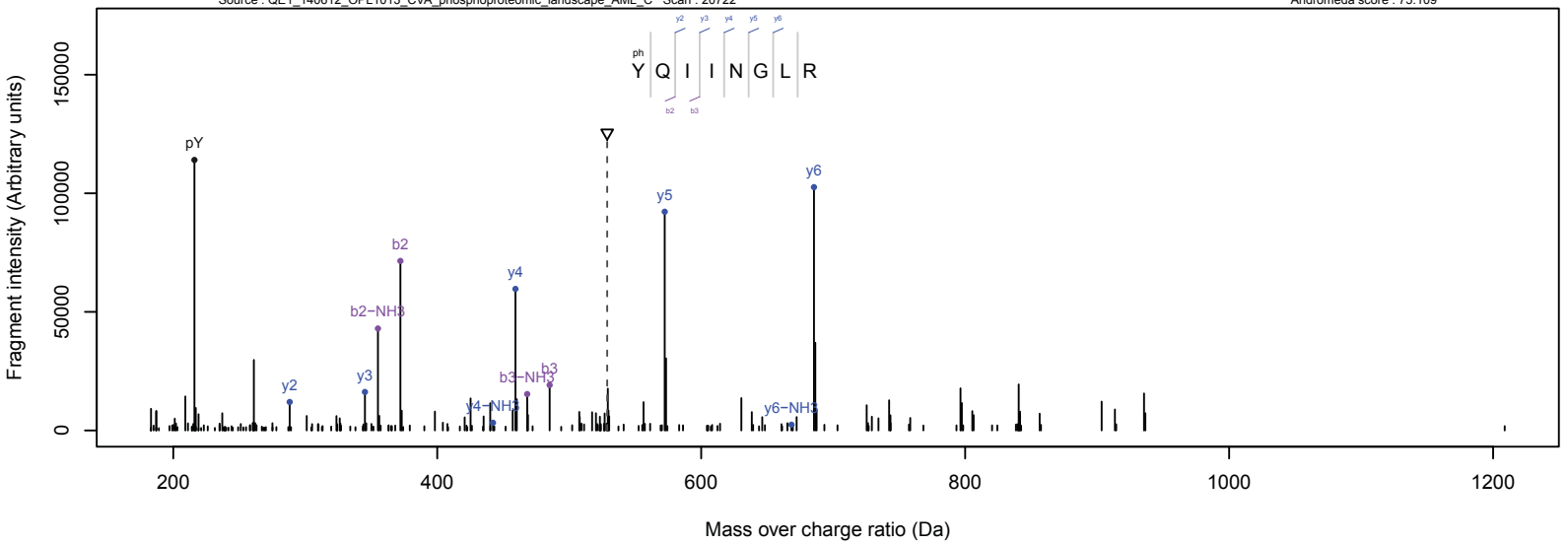
Andromeda score : 75.109



Y(ph)QIINGLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20722

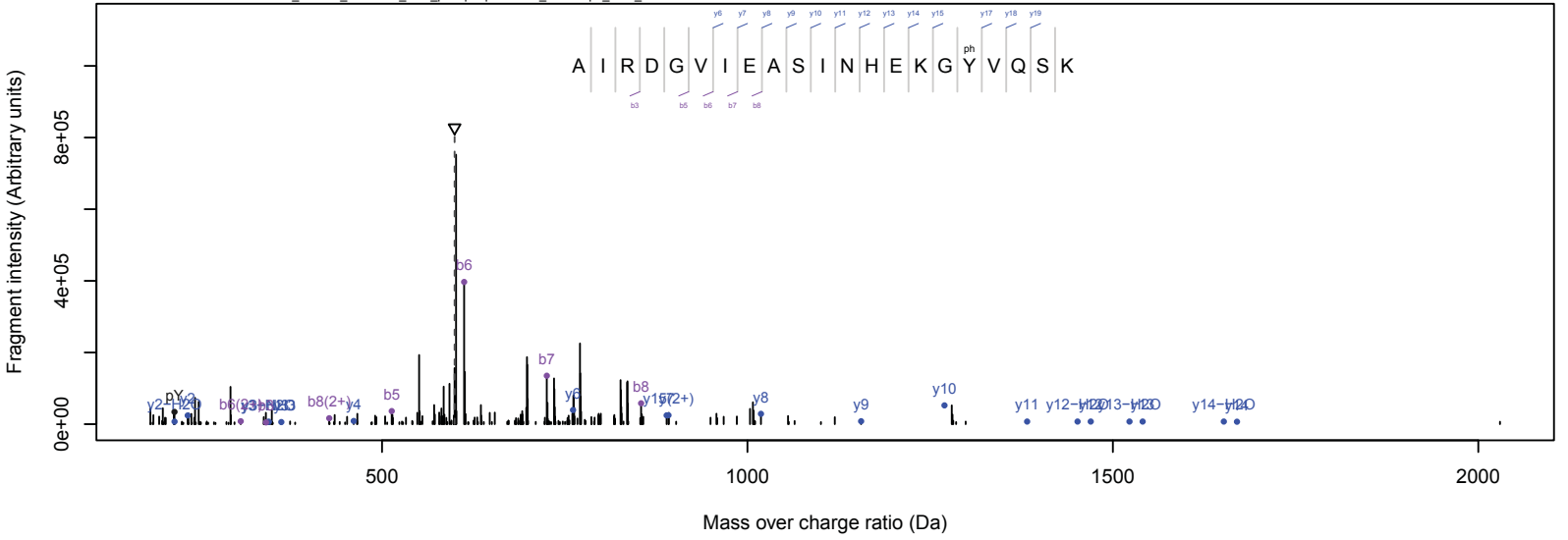
Andromeda score : 75.109



AIRDGVEASINHEKGYVQSK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21198

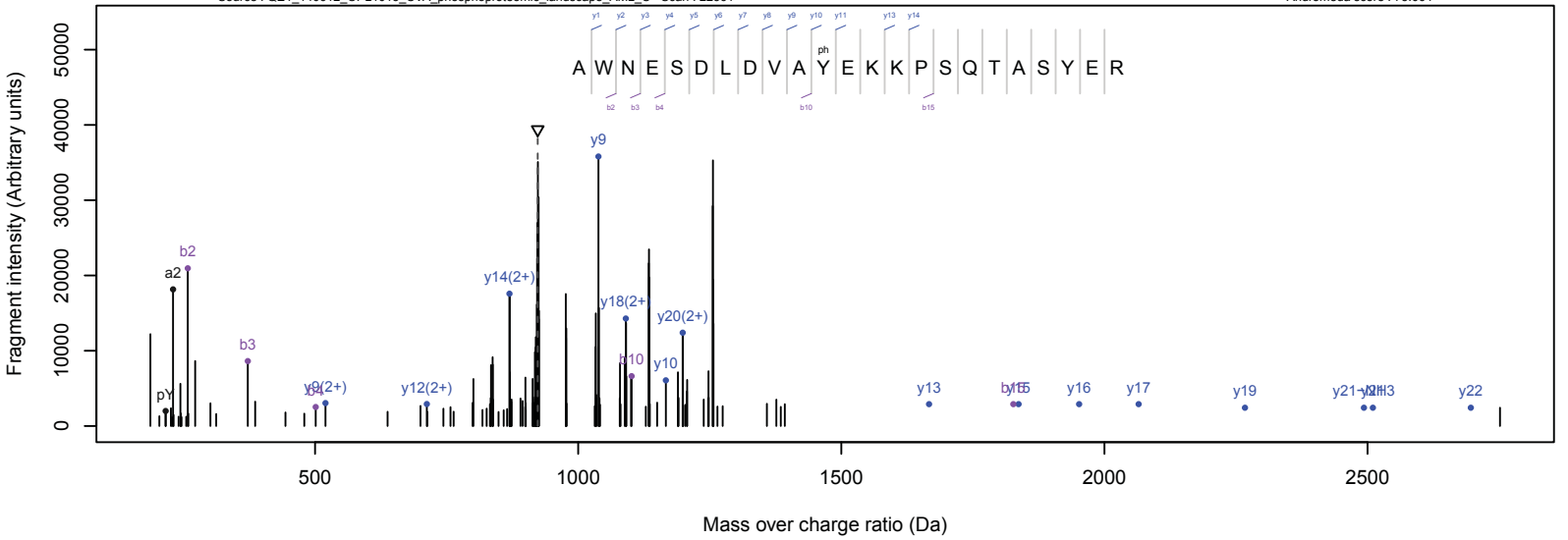
Andromeda score : 75.064



AWNESDLVAYEKKPSQTASYER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22561

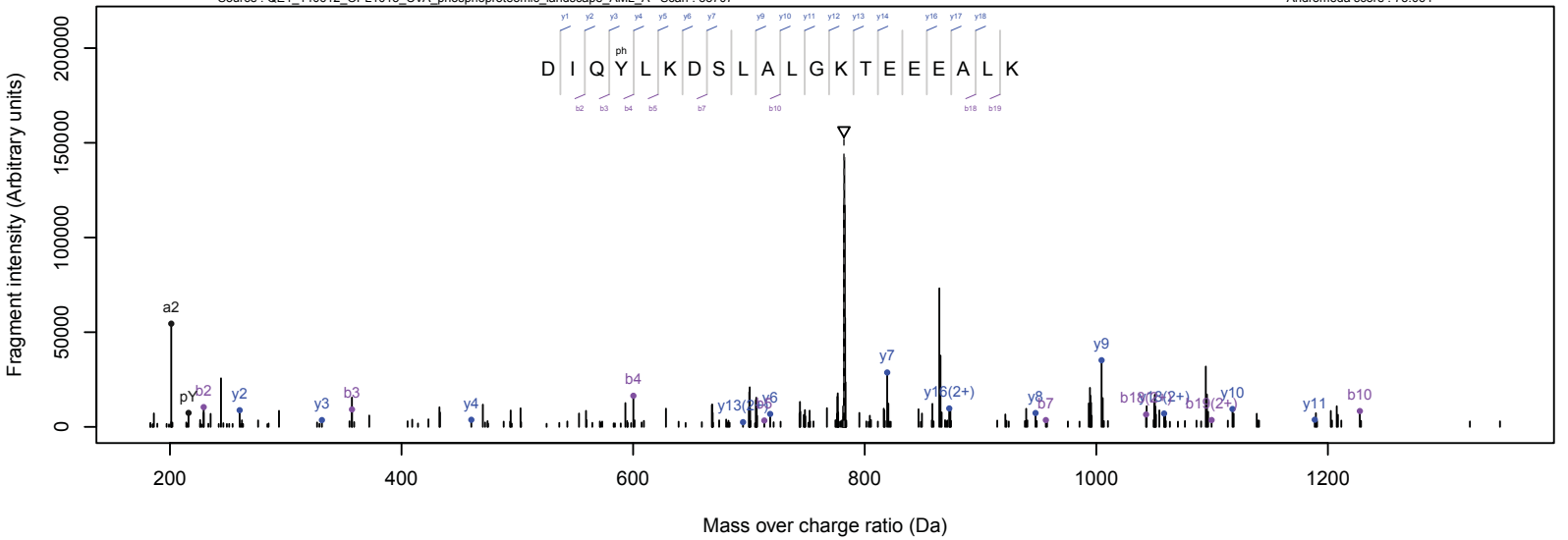
Andromeda score : 75.064



DIQY(ph)LKDSLALGKTEEEALK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 35707

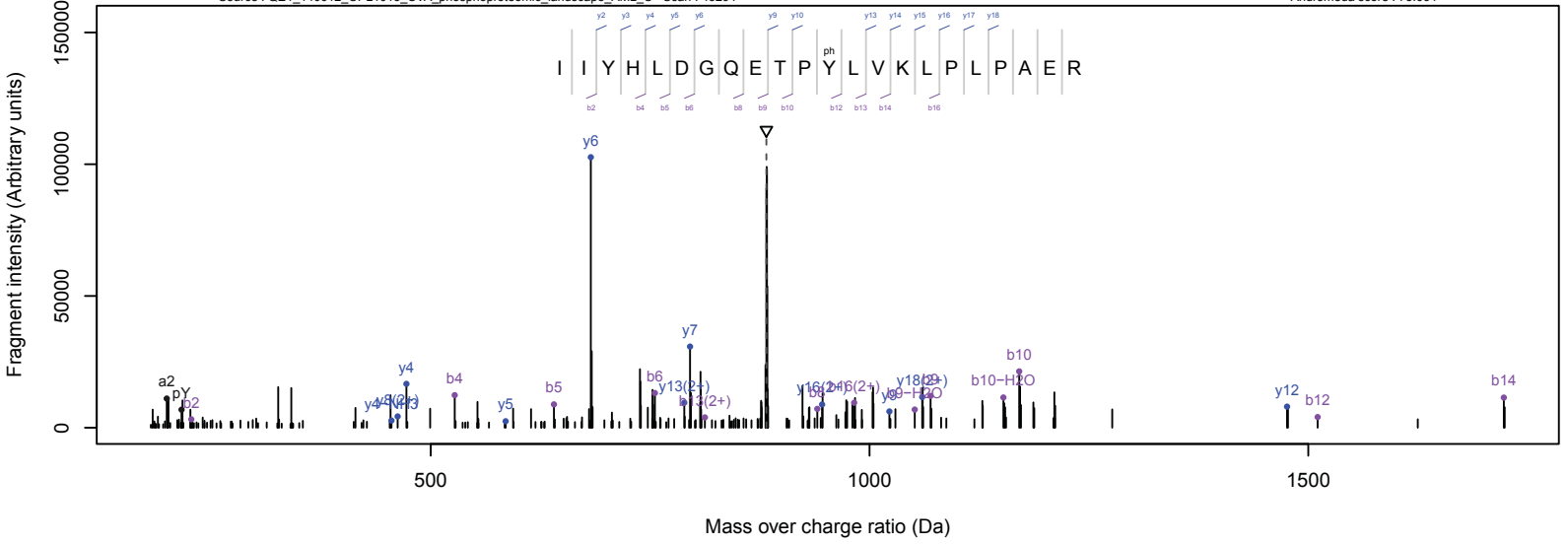
Andromeda score : 75.064



IYYHLDGQETPYLVKLP LPAER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43294

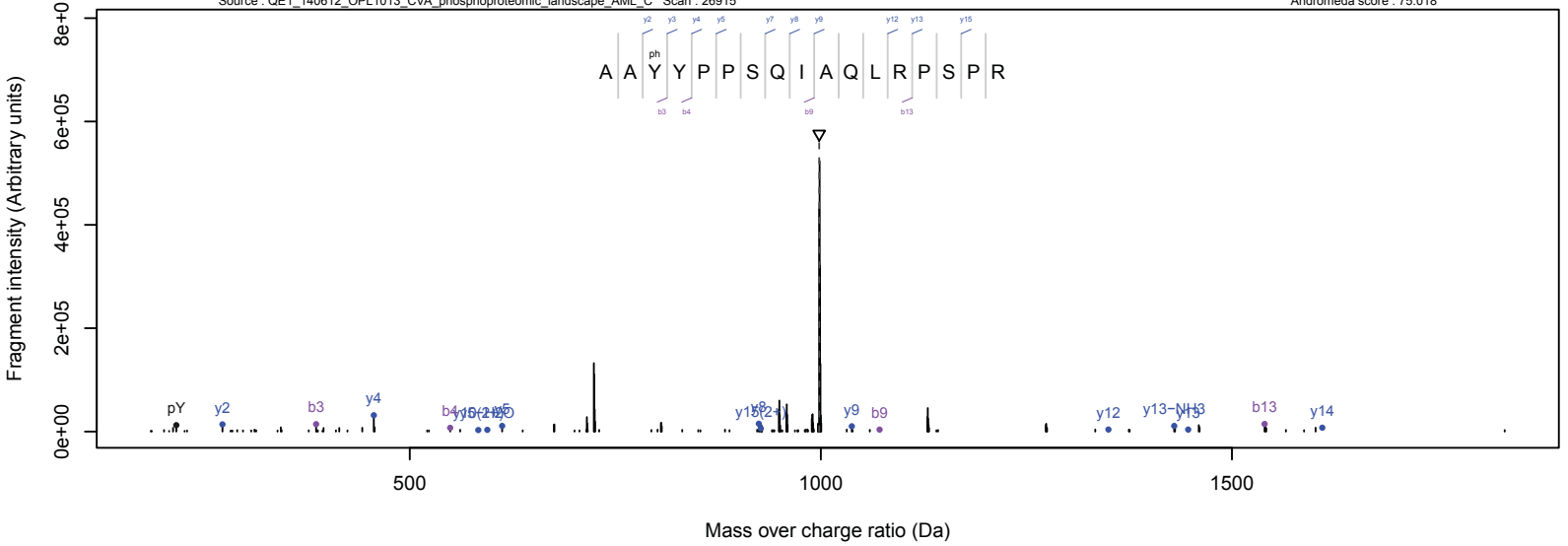
Andromeda score : 75.064



AA Y(ph)YPPSQI AQLRPSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26915

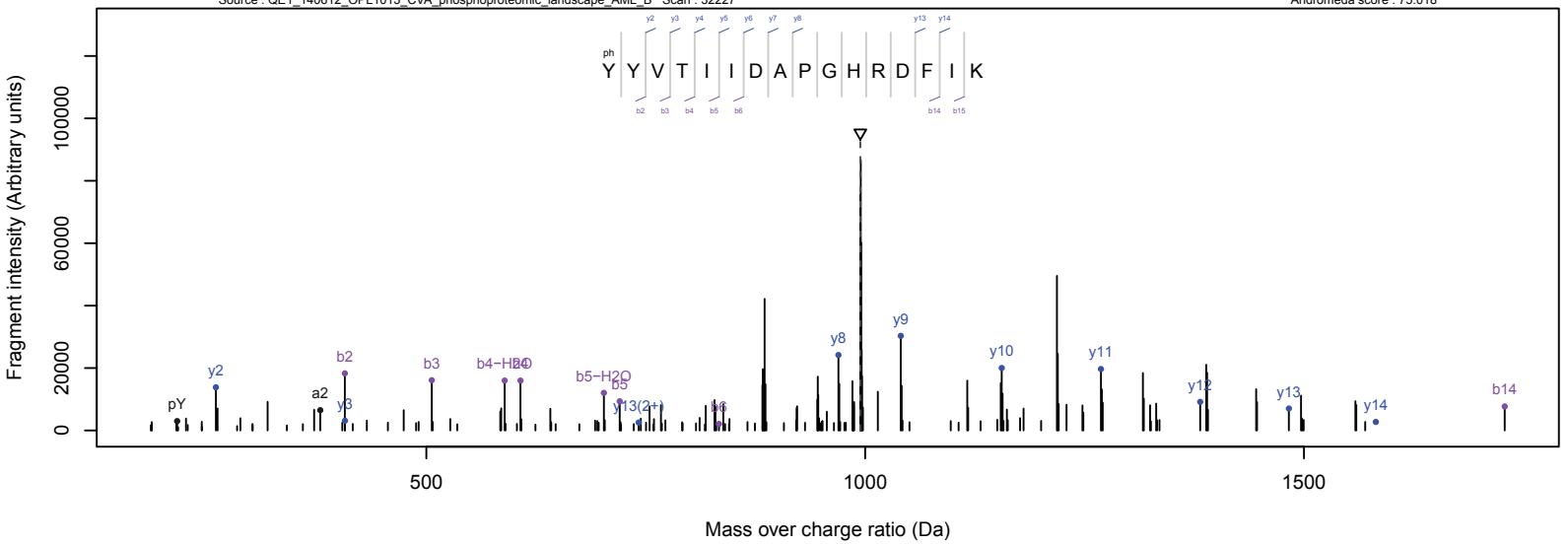
Andromeda score : 75.018



Y(ph)YVTIIDAPGHRDFIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32227

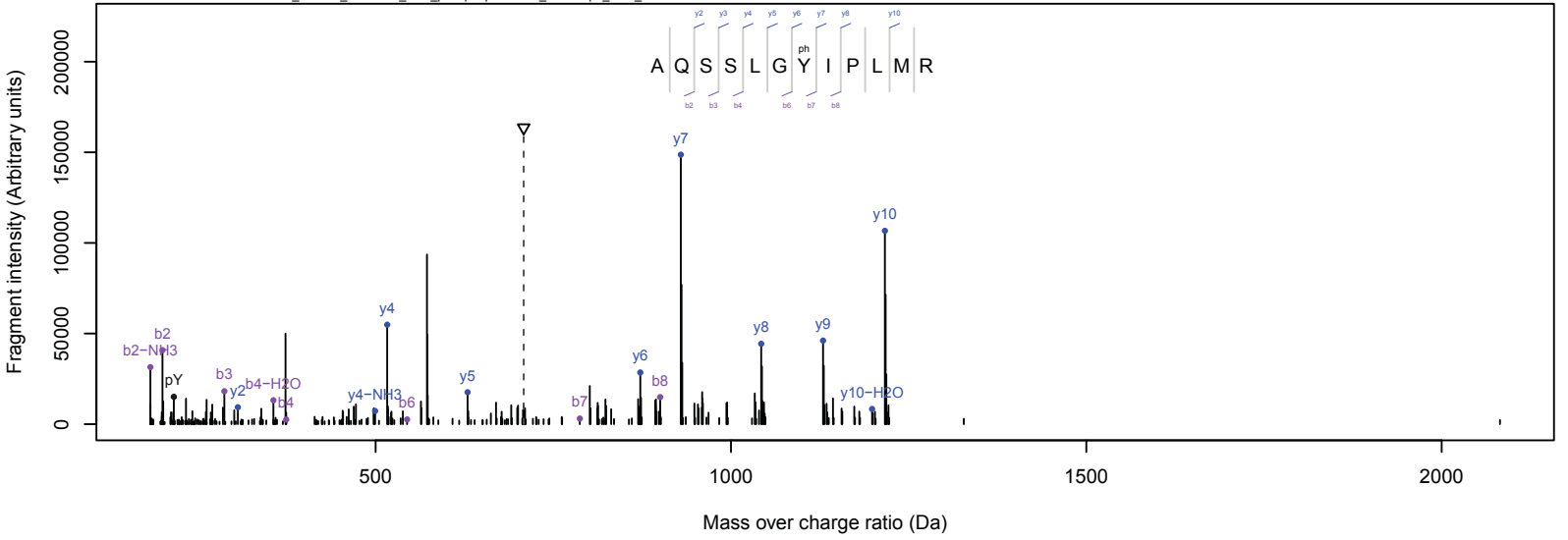
Andromeda score : 75.018



**AQSSLG(ph)PLMR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37783

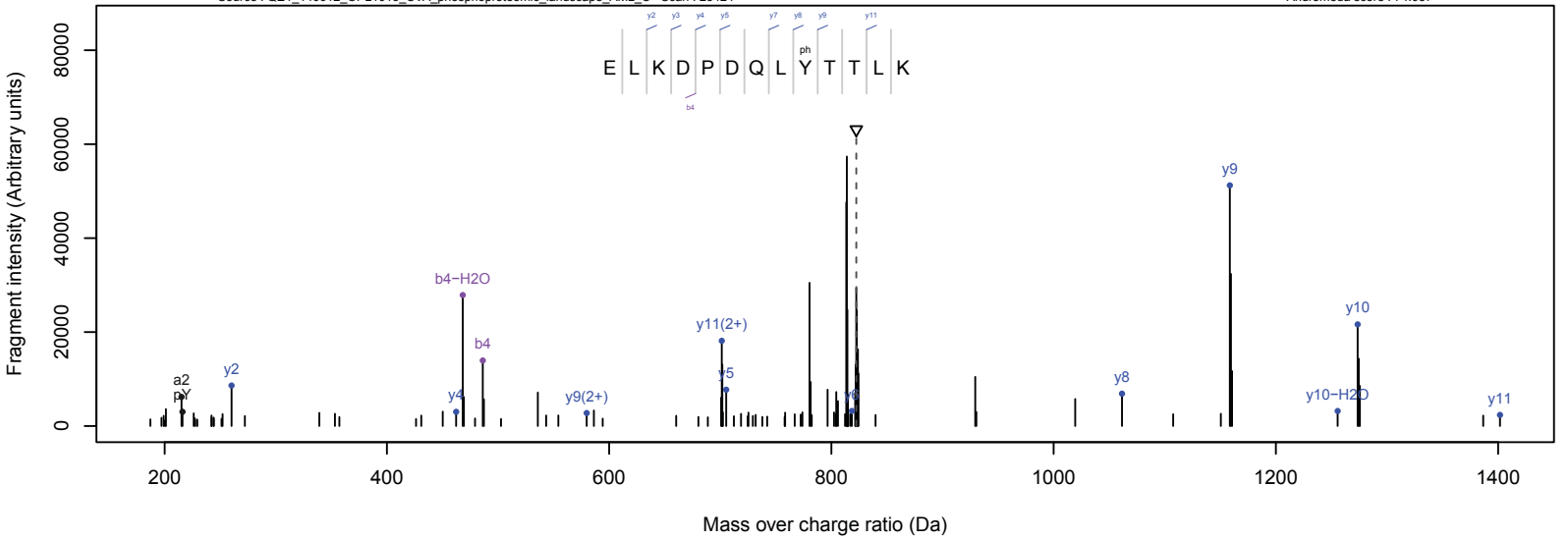
Andromeda score: 74.987



**ELKDPDQLY(ph)TTLK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23424

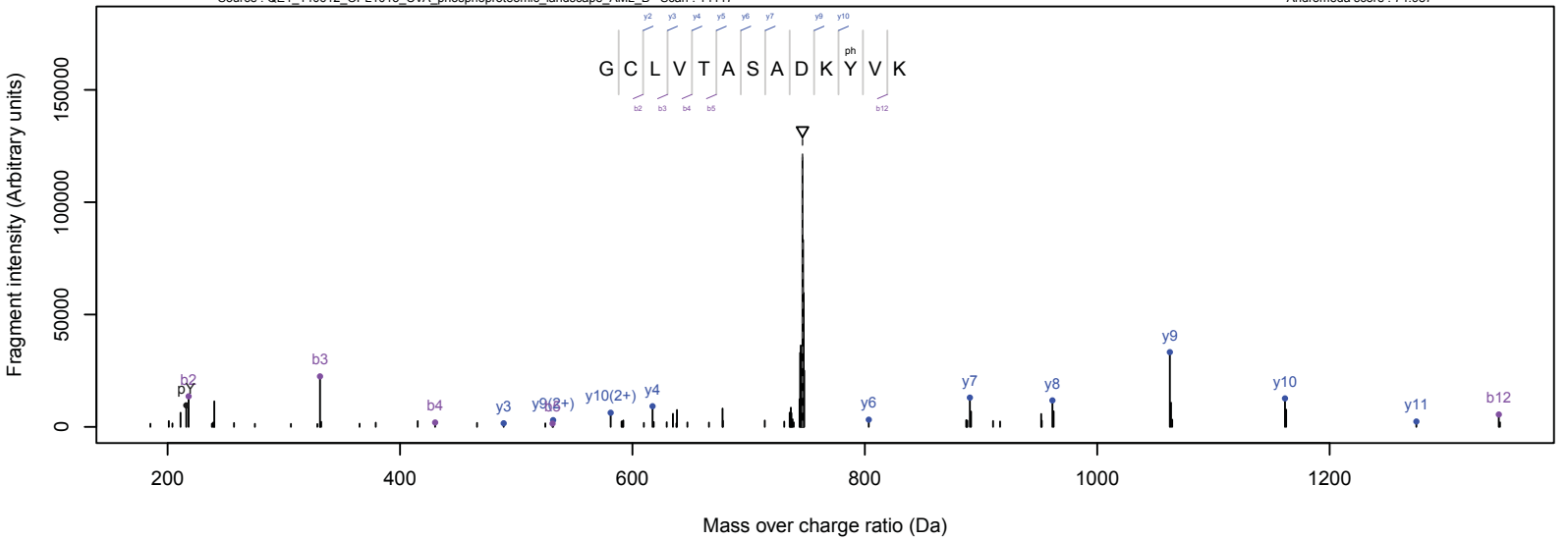
Andromeda score: 74.987



**GCLVTASADKY(ph)VK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14117

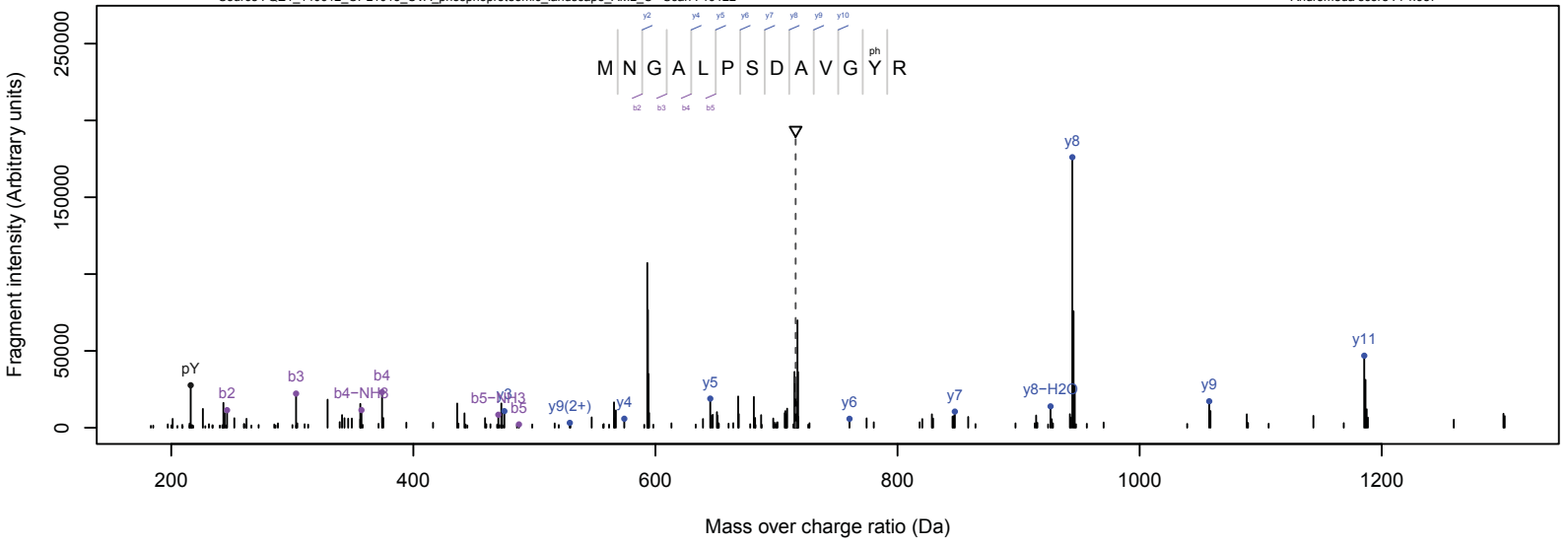
Andromeda score: 74.987



**MNGALPSDAVG(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19122

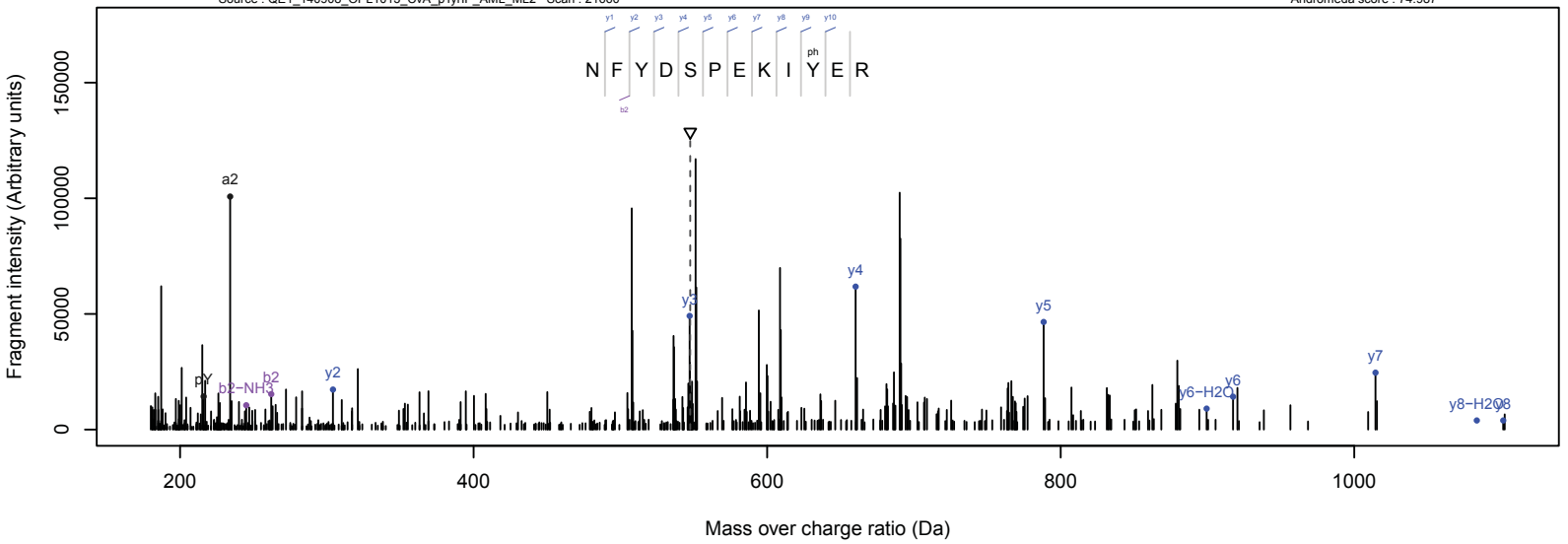
Andromeda score : 74.987



**NFYDSPEKIY(ph)ER-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 21660

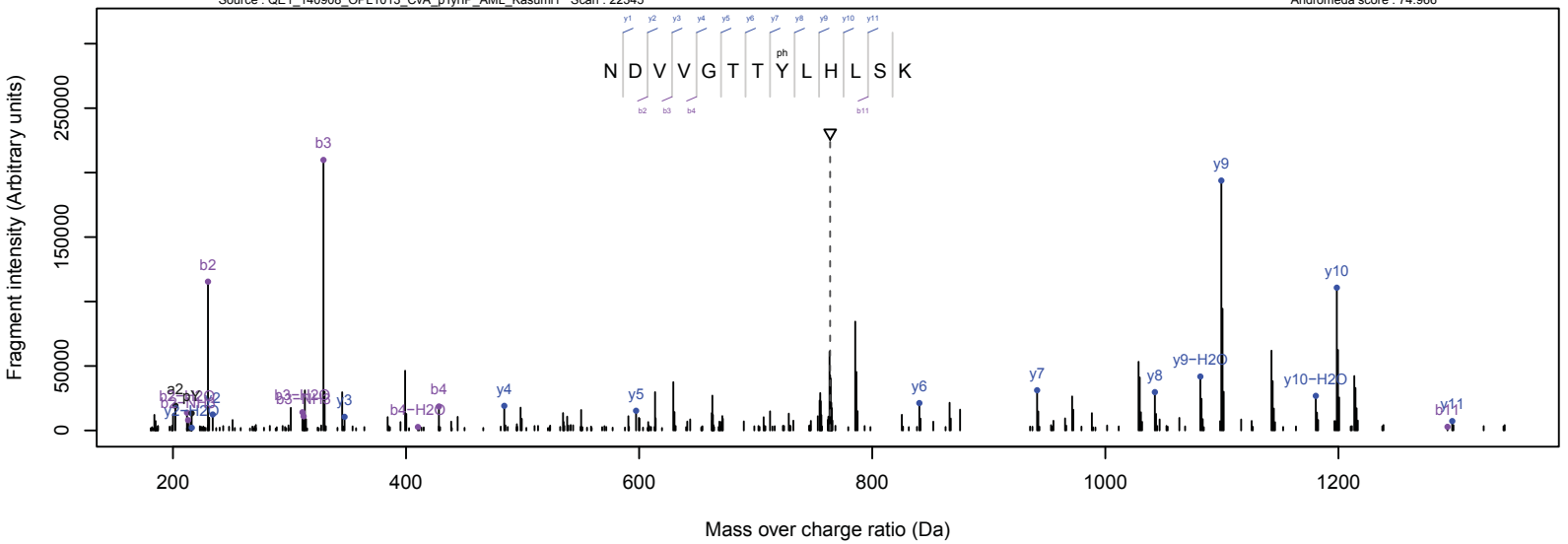
Andromeda score : 74.987



**NDVVGTTY(ph)LHLSK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 22345

Andromeda score : 74.966

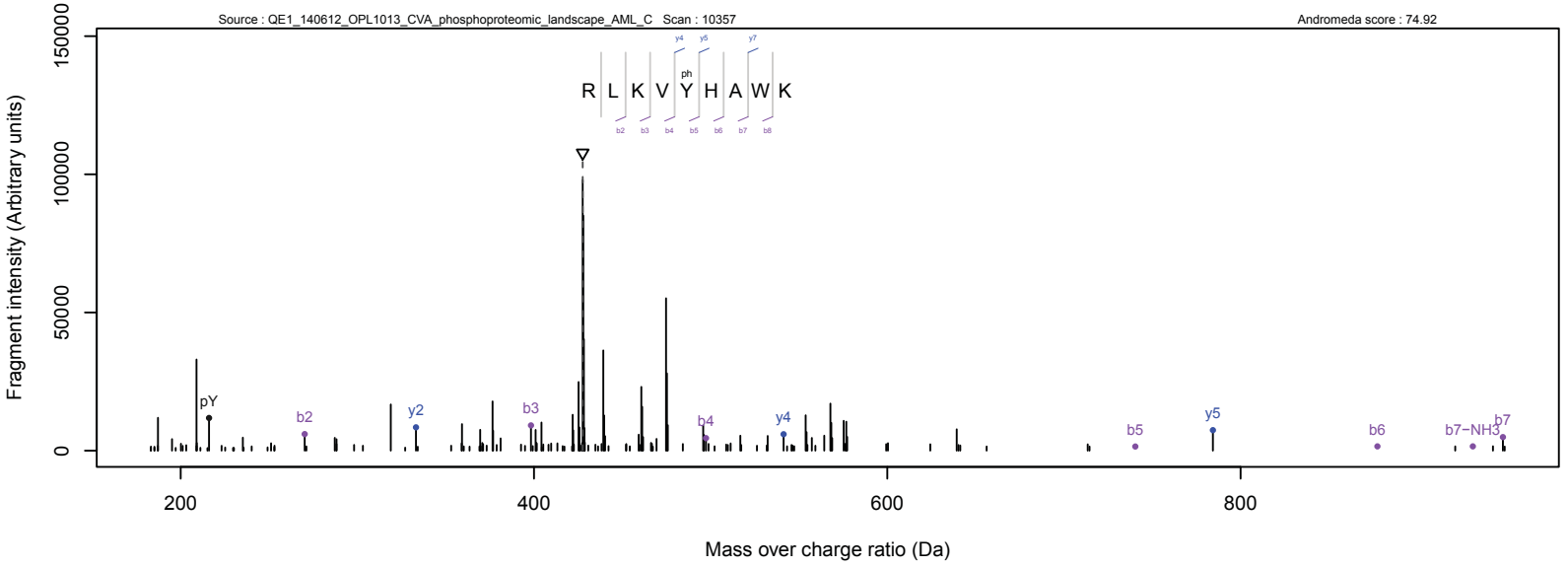




RLKVY(ph)HAWK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10357

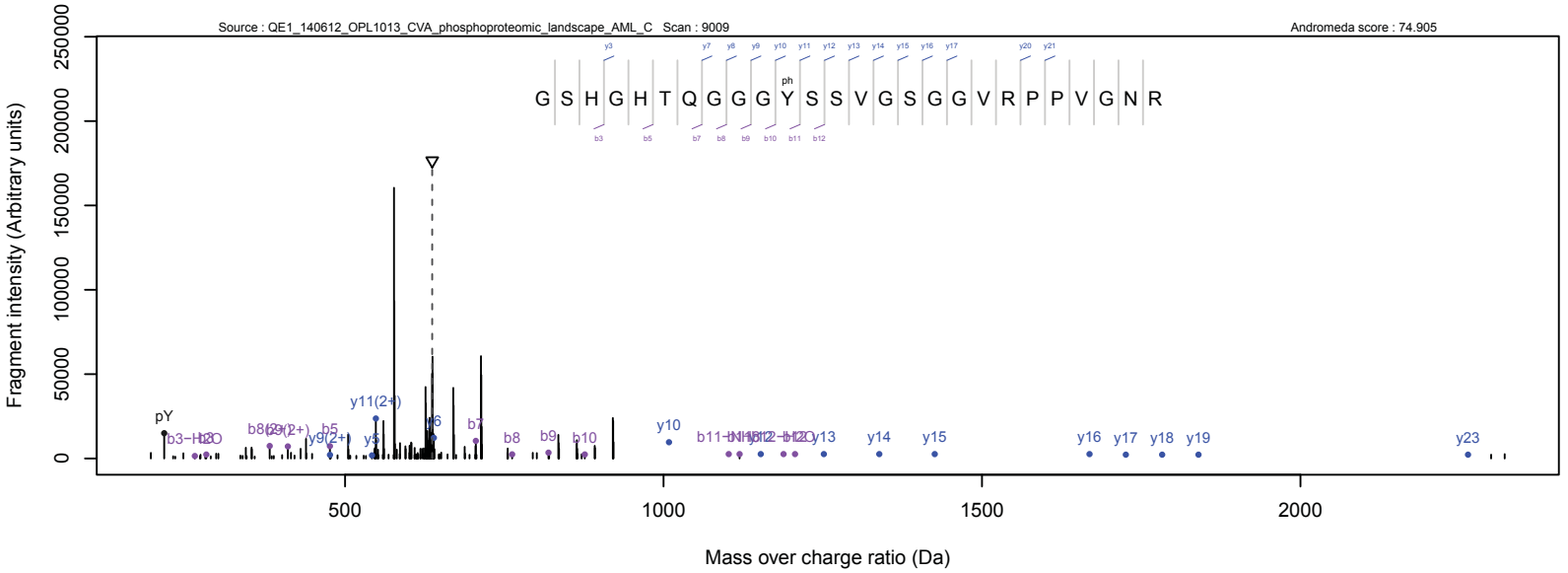
Andromeda score : 74.92



GSHGHTQGGY(ph)SSVSGGVRPPVGNR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9009

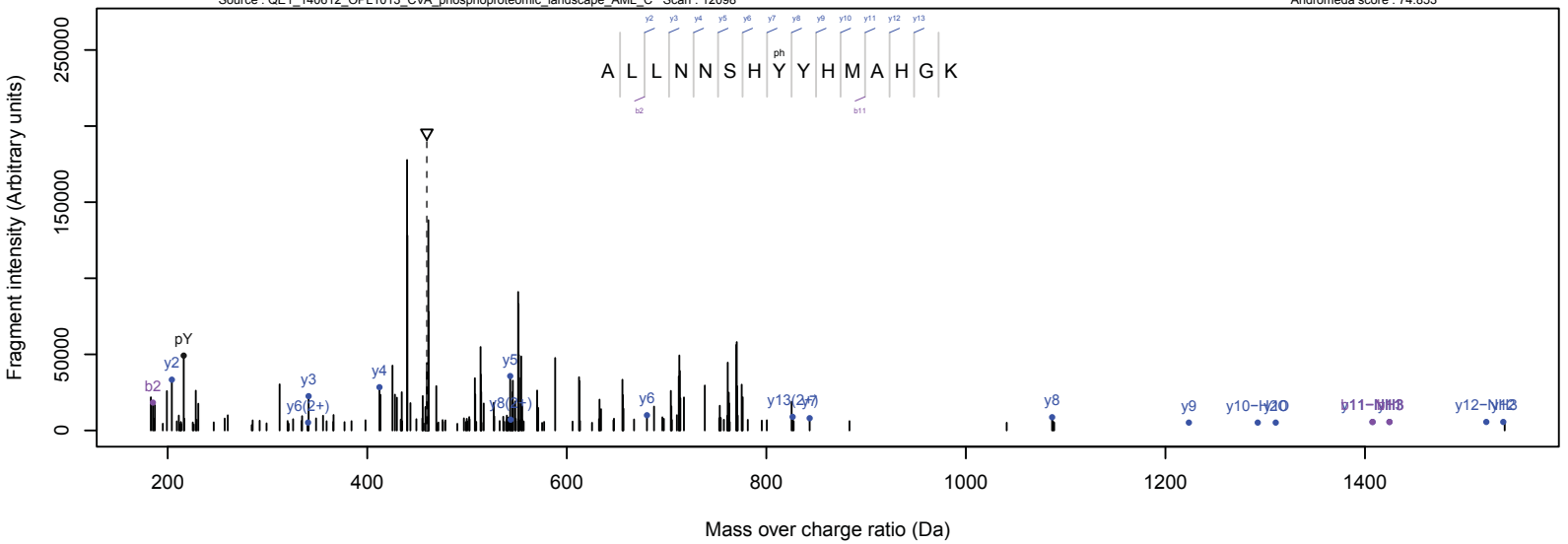
Andromeda score : 74.905



ALLNNSHY(ph)YHMAHGK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12098

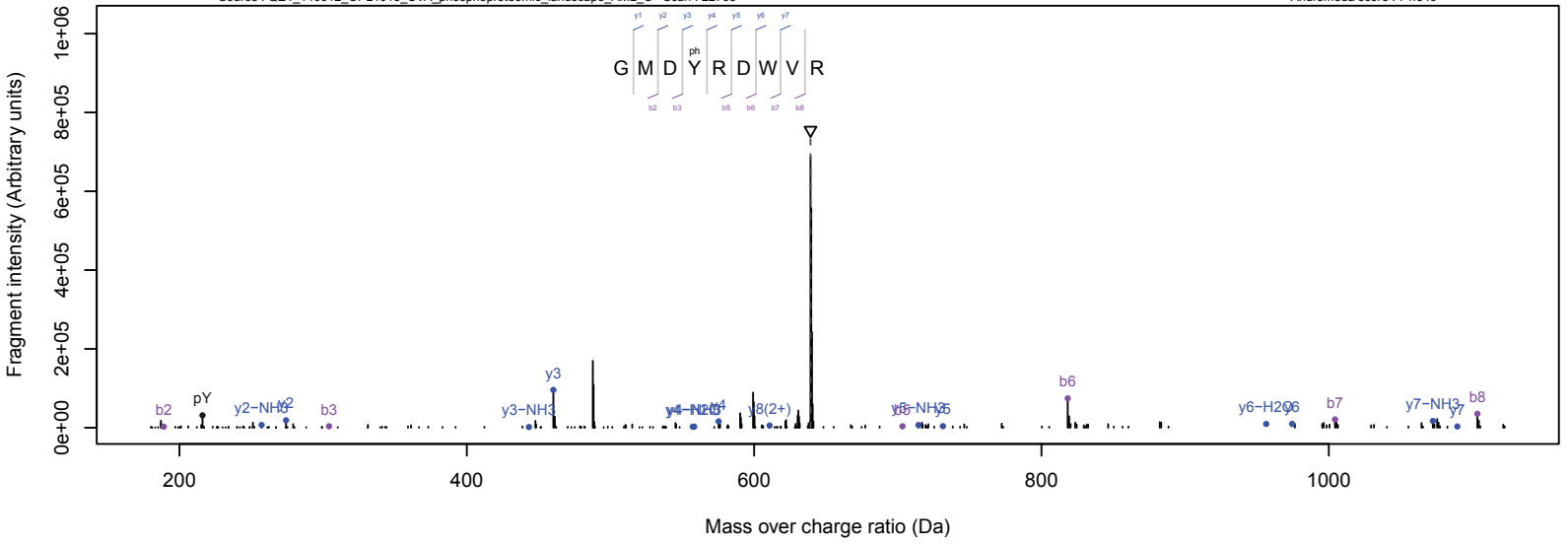
Andromeda score : 74.853



GMDY(ph)RDWVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 22785

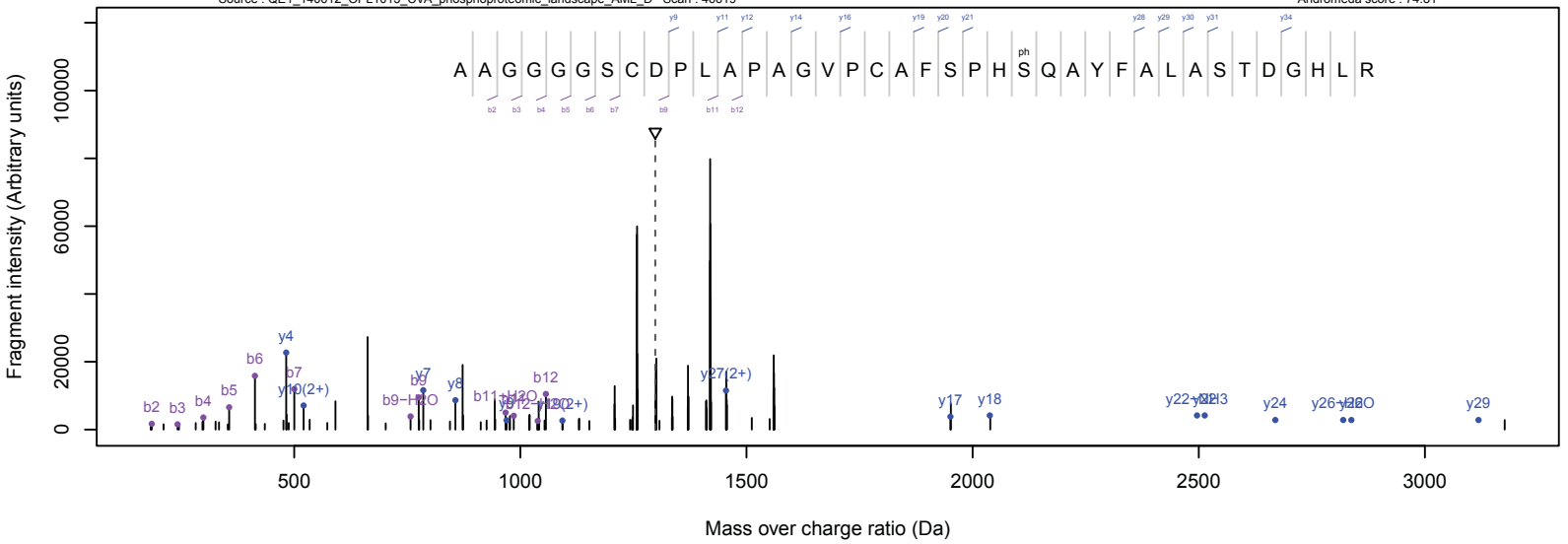
Andromeda score: 74.849



(ac)AAGGGGSCDPLAPAGVPCAFSPHS(ph)QAYFALASTDGHLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 46619

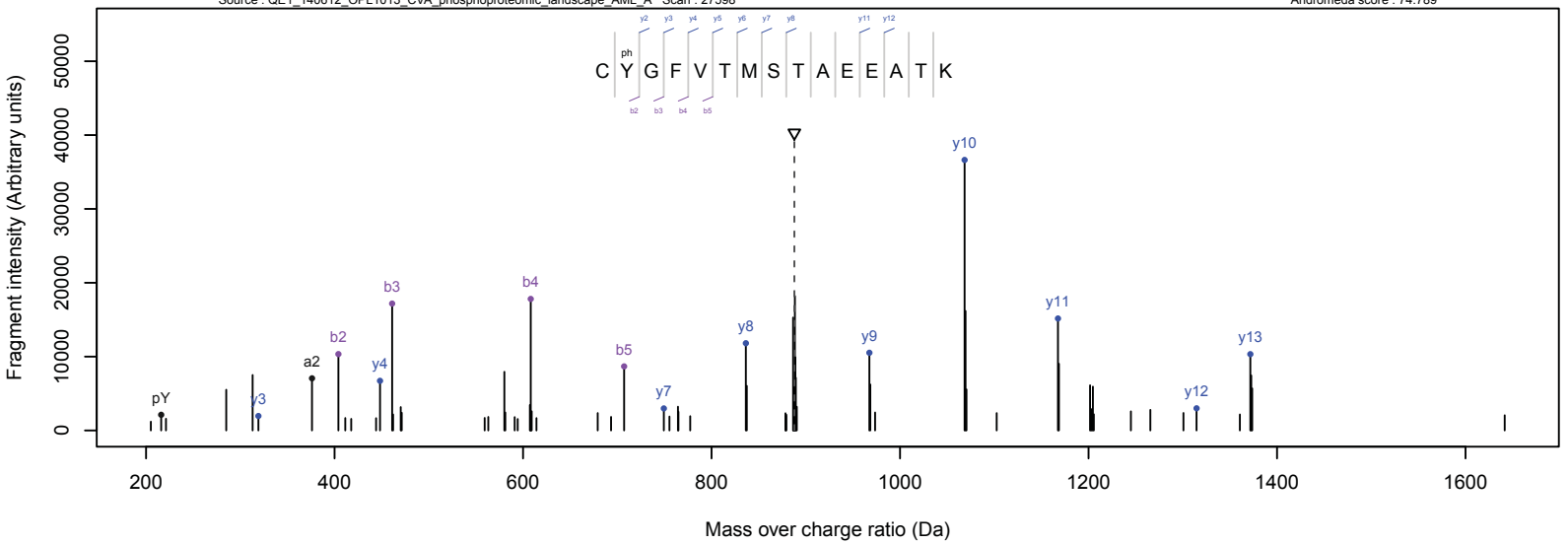
Andromeda score: 74.81



CY(ph)GFVTMSTAEATK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 27598

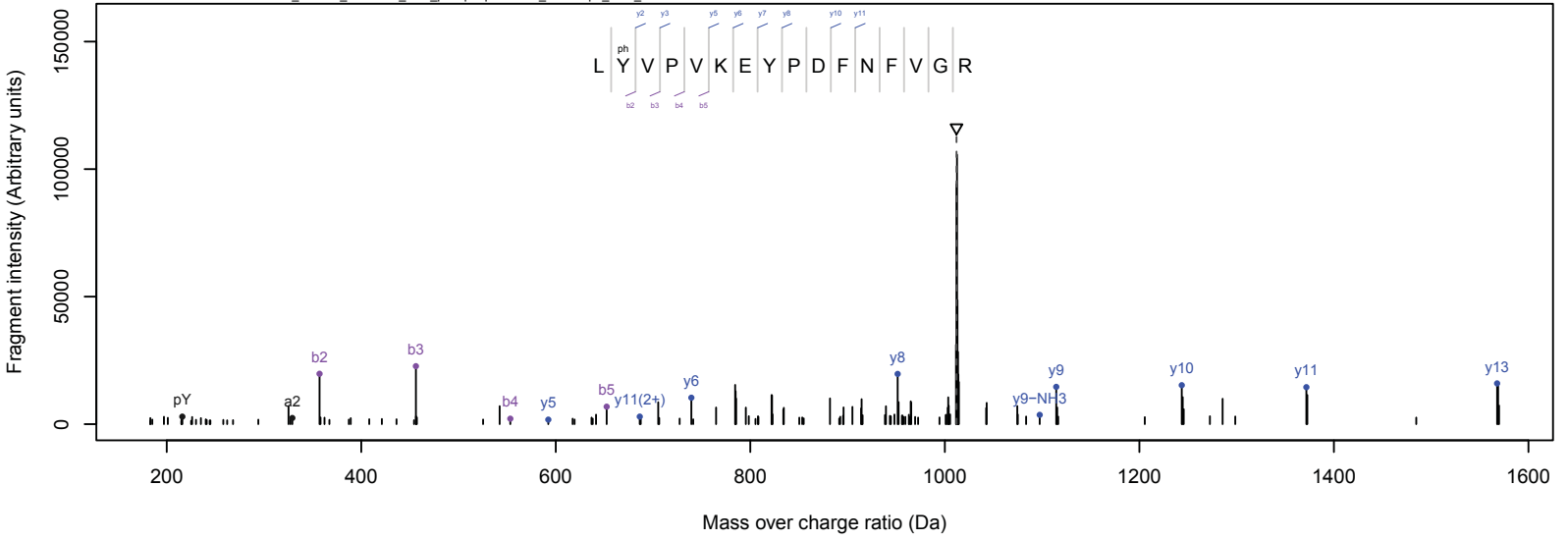
Andromeda score: 74.789



LY(ph)VPVKEYPDFNFVGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 42864

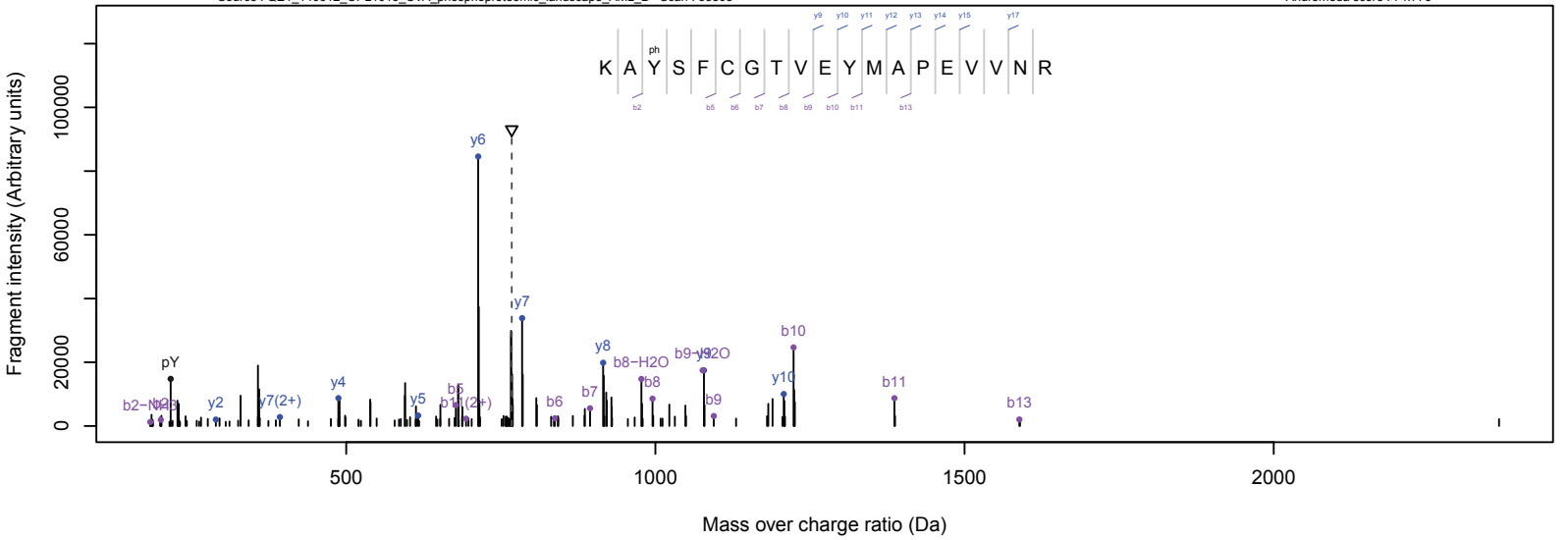
Andromeda score : 74.789



KAY(ph)SFCGTVEYMAPEVVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33808

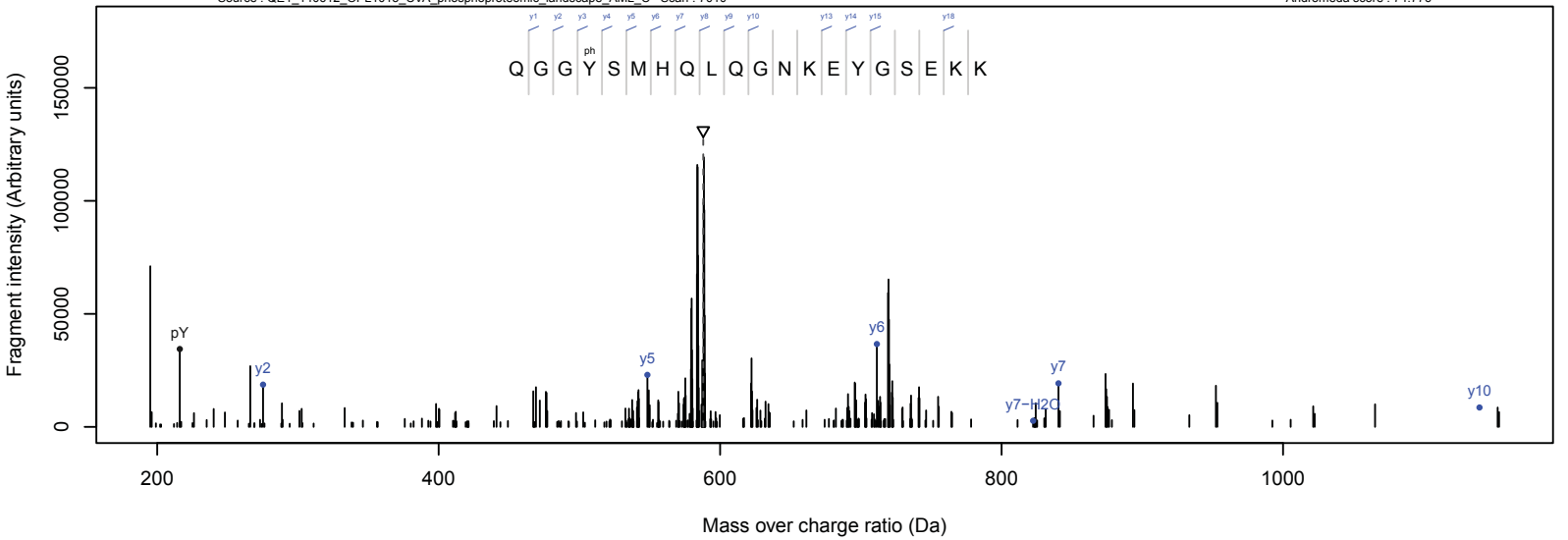
Andromeda score : 74.776



QGGY(ph)SMHQLQGNKEYGSEKK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7010

Andromeda score : 74.776

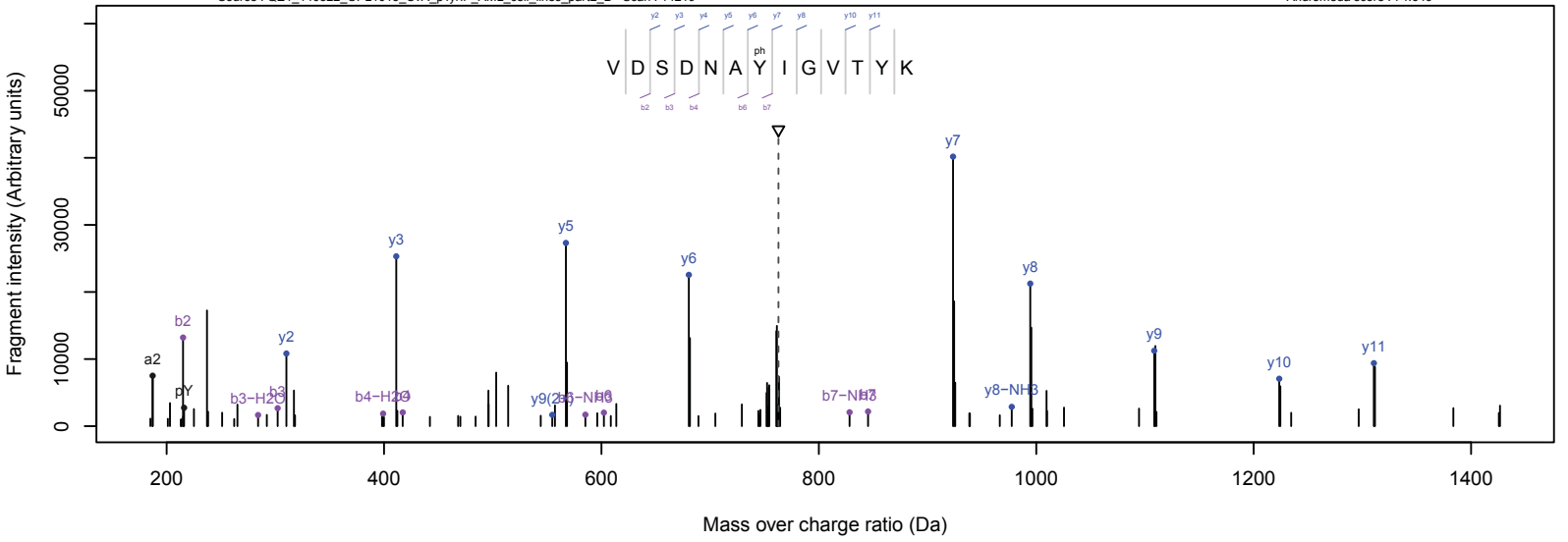




VSDSNAY(ph)IGVTYK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 14219

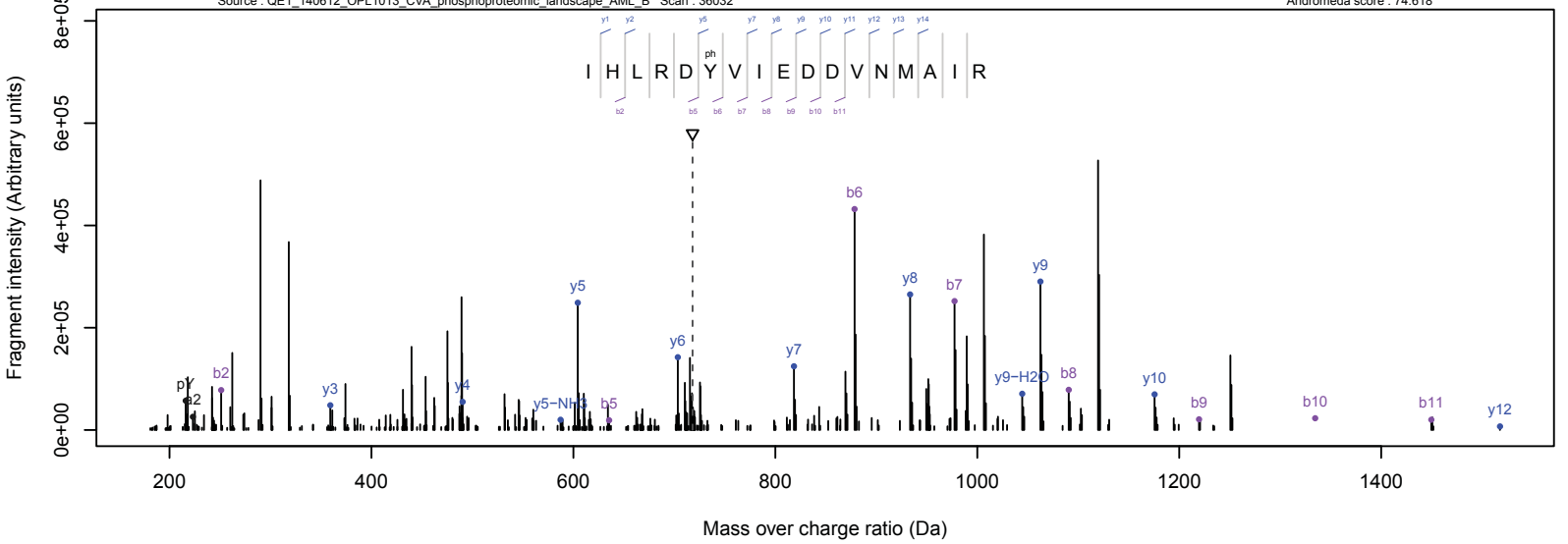
Andromeda score: 74.649



IHLRDY(ph)VEDDVNMAIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 36032

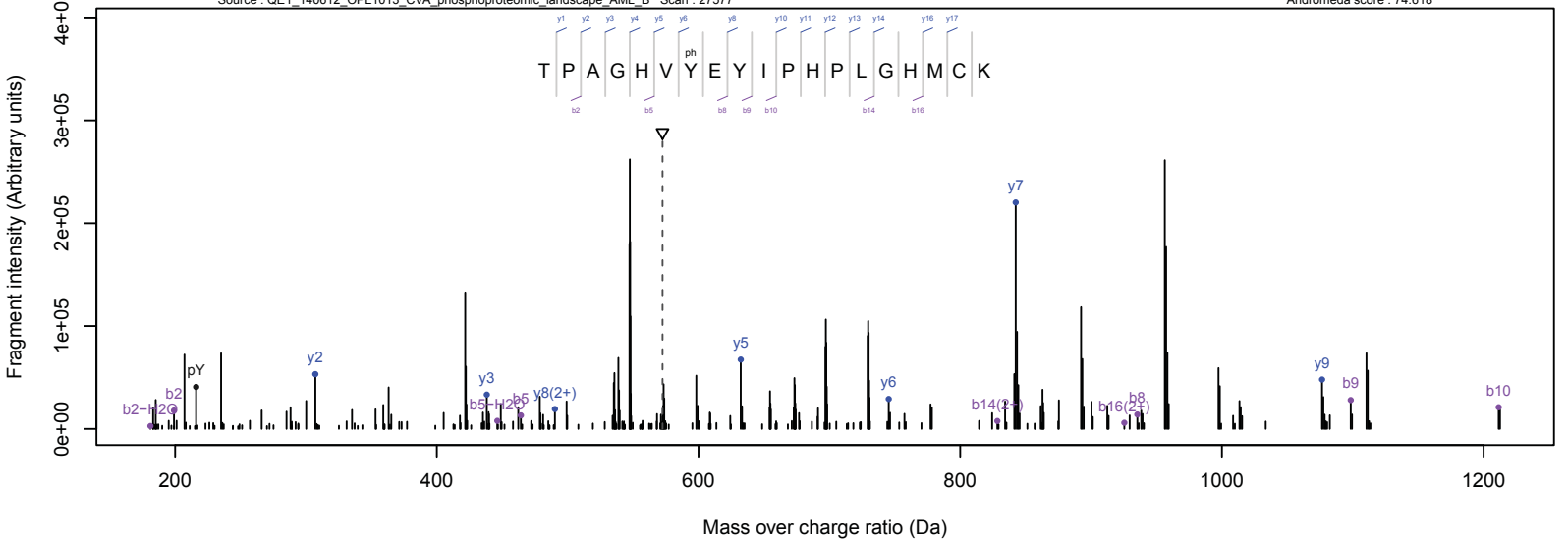
Andromeda score: 74.618



TPAGHVY(ph)EYIPHLGHMCK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27577

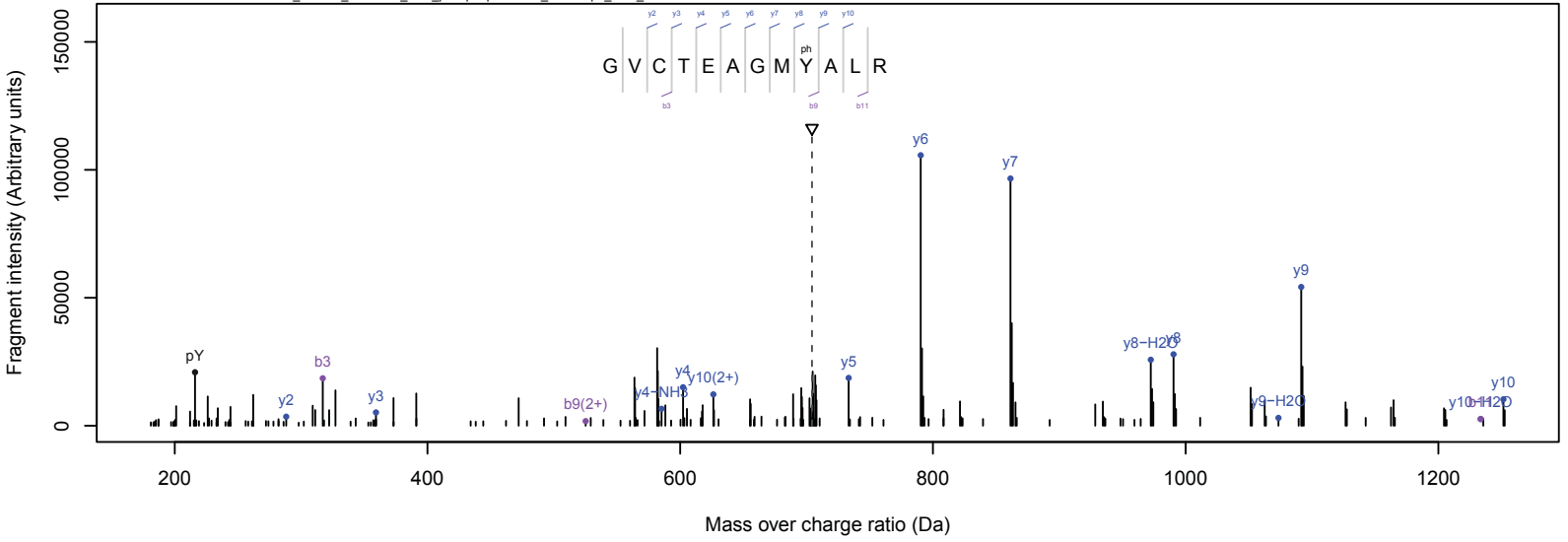
Andromeda score: 74.618



GVCTEAGMY(ph)ALR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21971

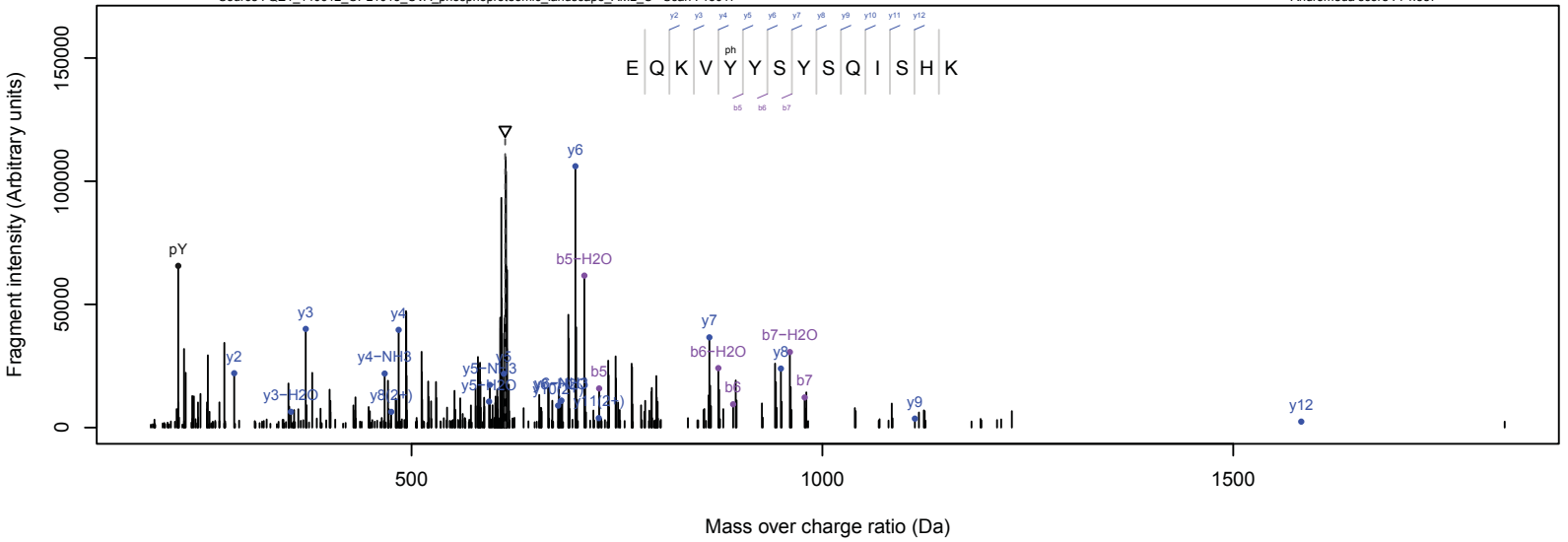
Andromeda score : 74.611



EQKVY(ph)YSYSQISHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13017

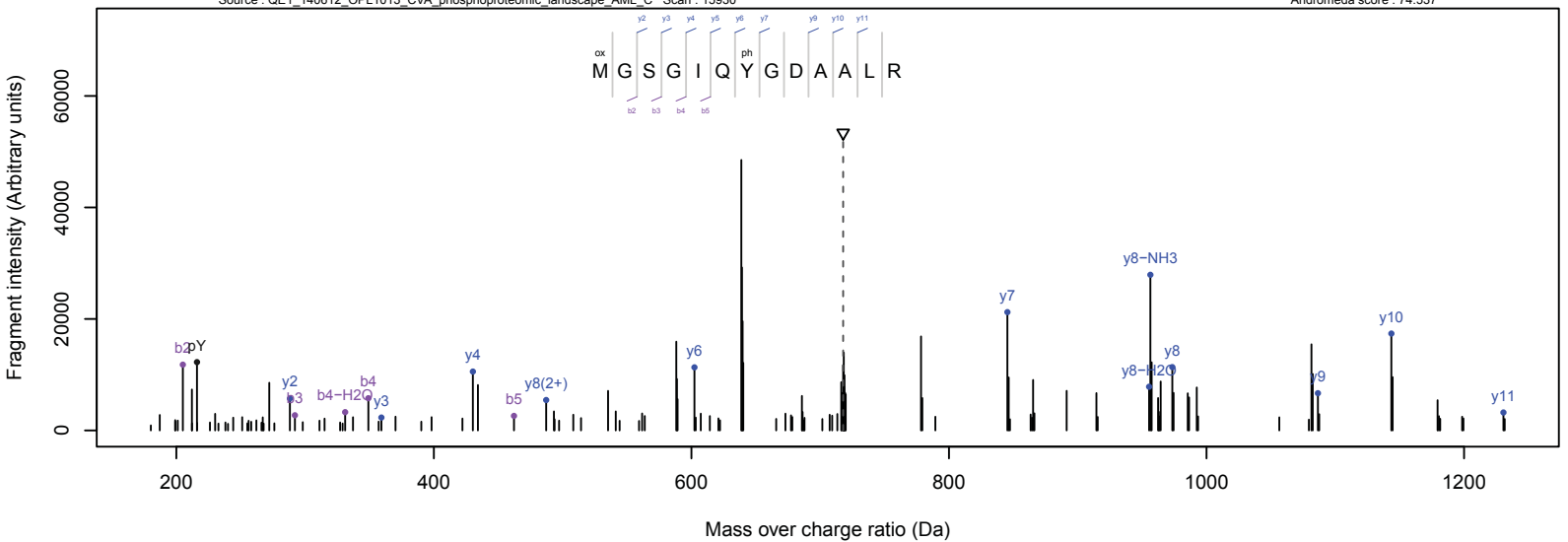
Andromeda score : 74.587



M(ox)GSGIQY(ph)GDAALR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15930

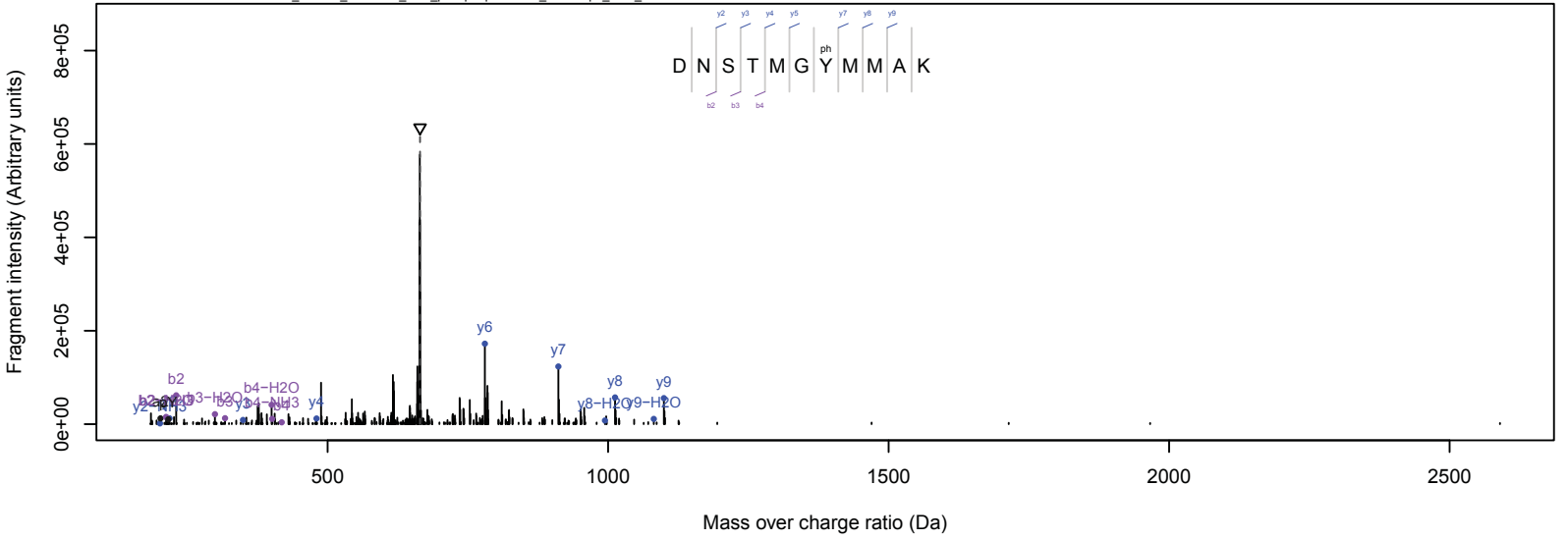
Andromeda score : 74.537



**DNSTMGY(ph)MMAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17224

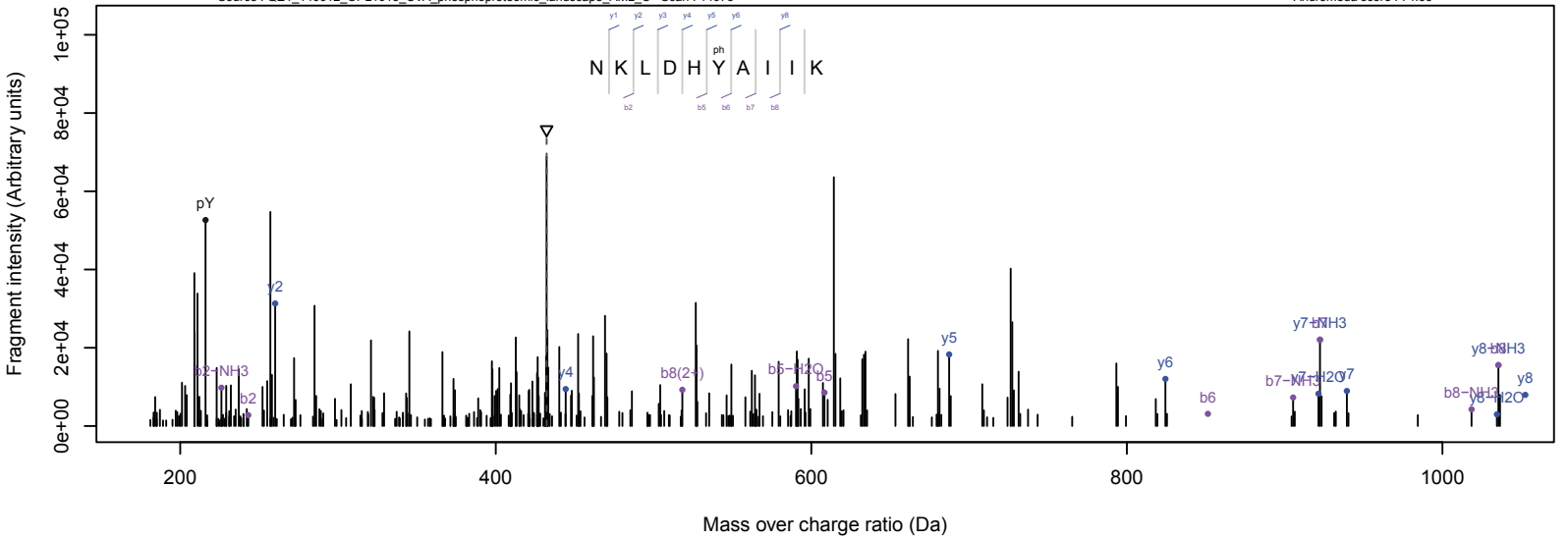
Andromeda score : 74.53



**NKLDHY(ph)AIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14973

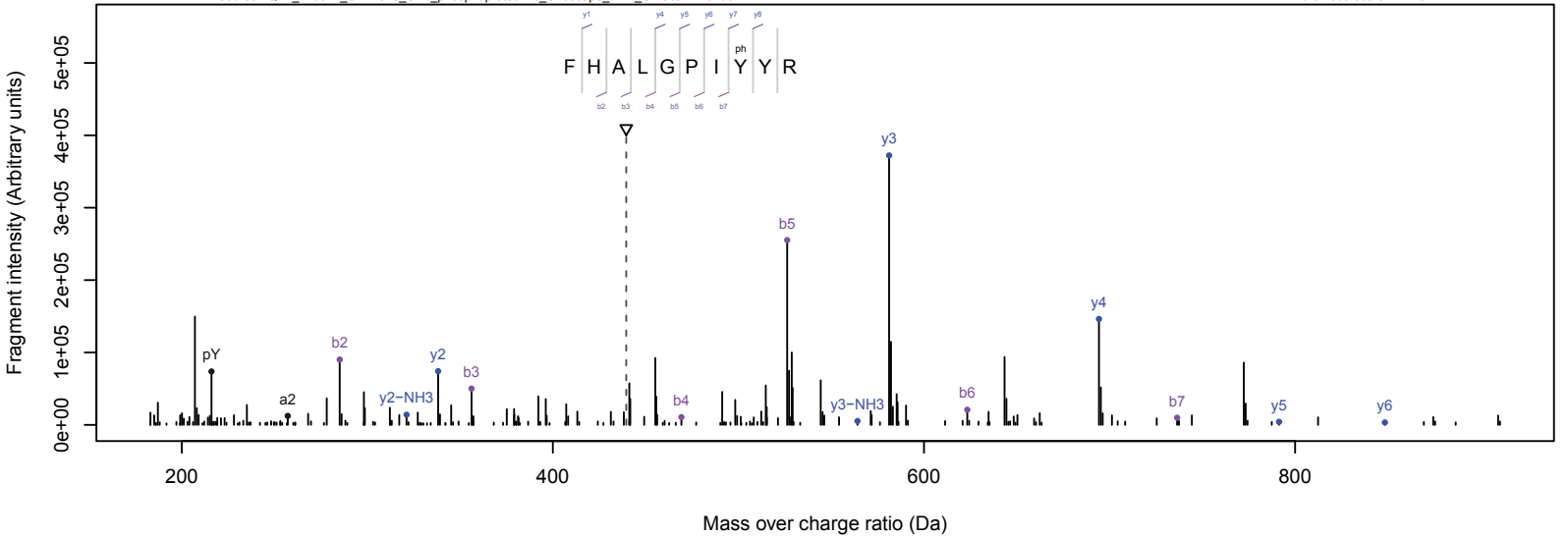
Andromeda score : 74.53



**FHALGPIY(ph)YR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29498

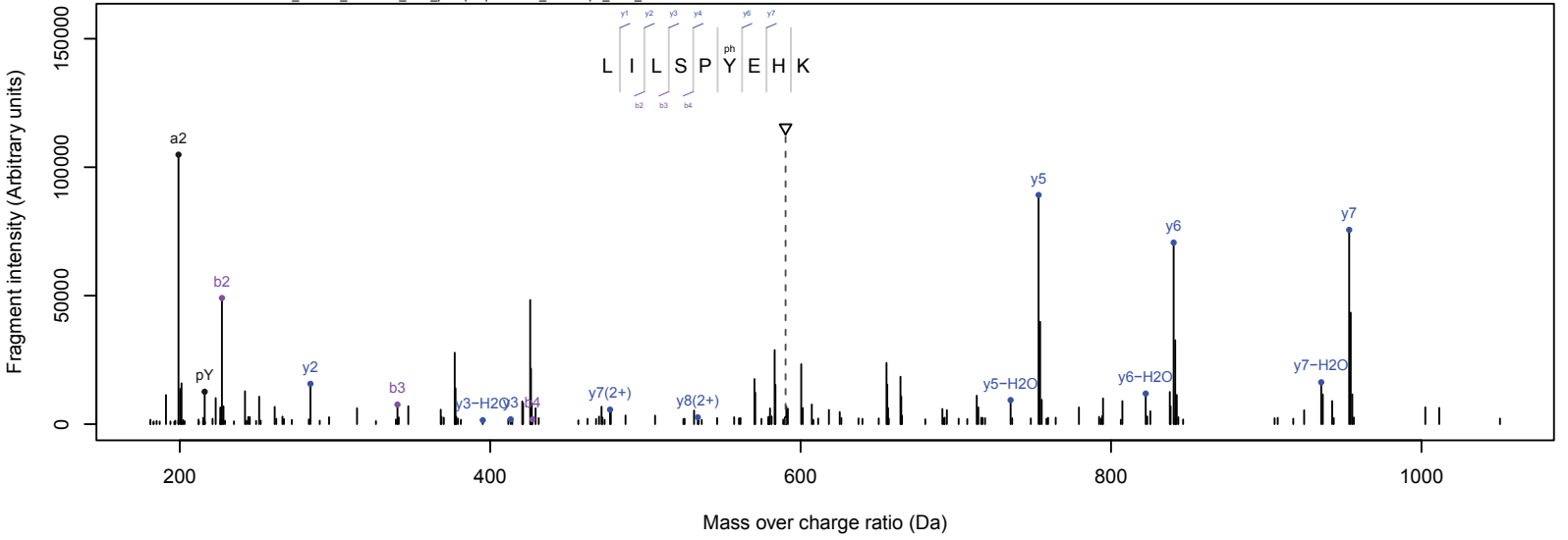
Andromeda score : 74.464



LILSPY(ph)EHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15748

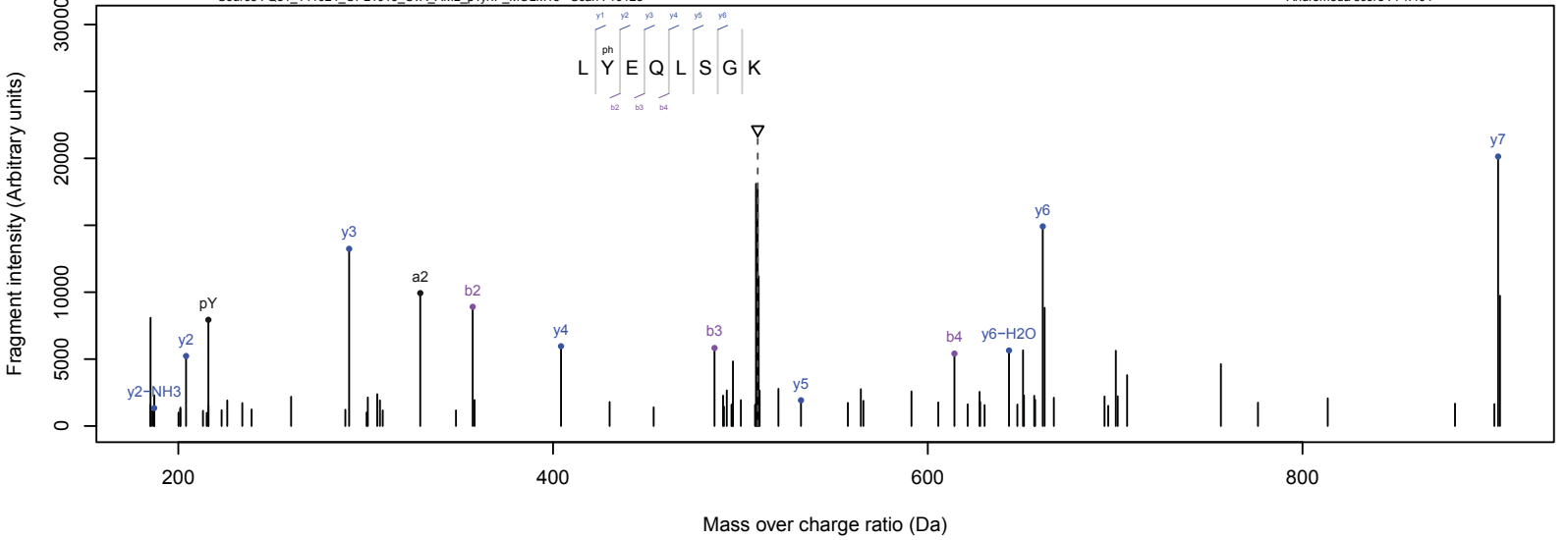
Andromeda score : 74.464



LY(ph)EQLSGK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyriP\_MOLM13 Scan : 10128

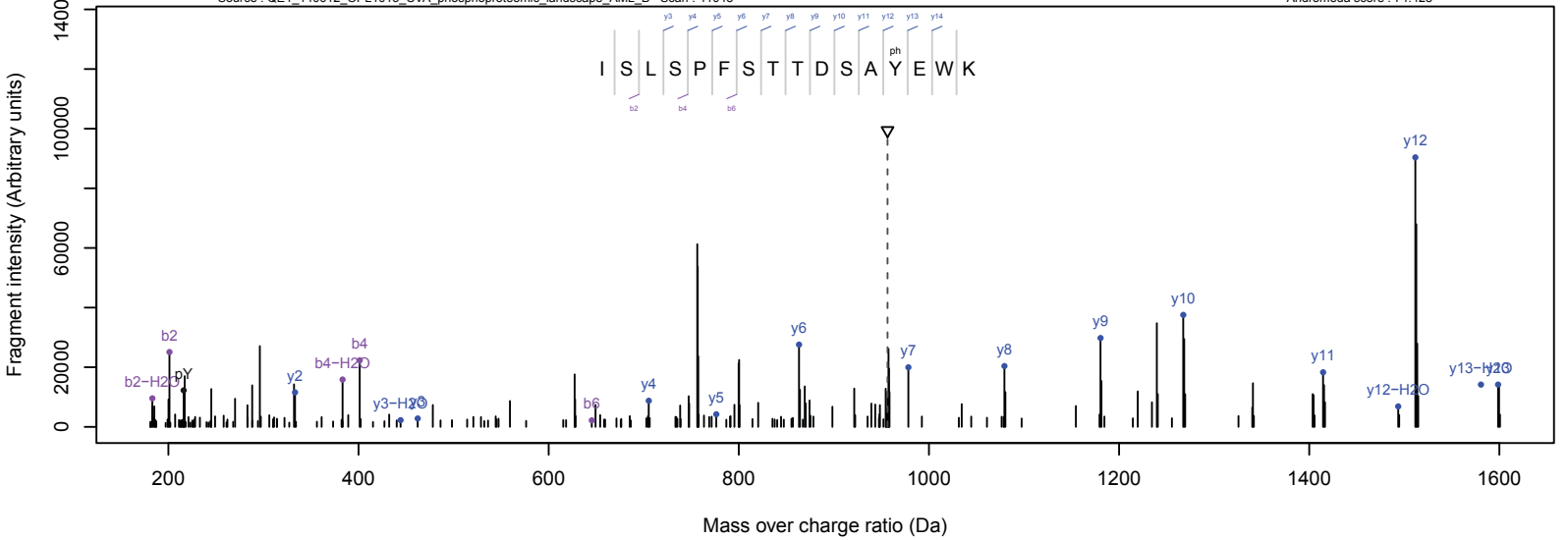
Andromeda score : 74.464



ISLSPFSTDSAY(ph)EWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 41048

Andromeda score : 74.428

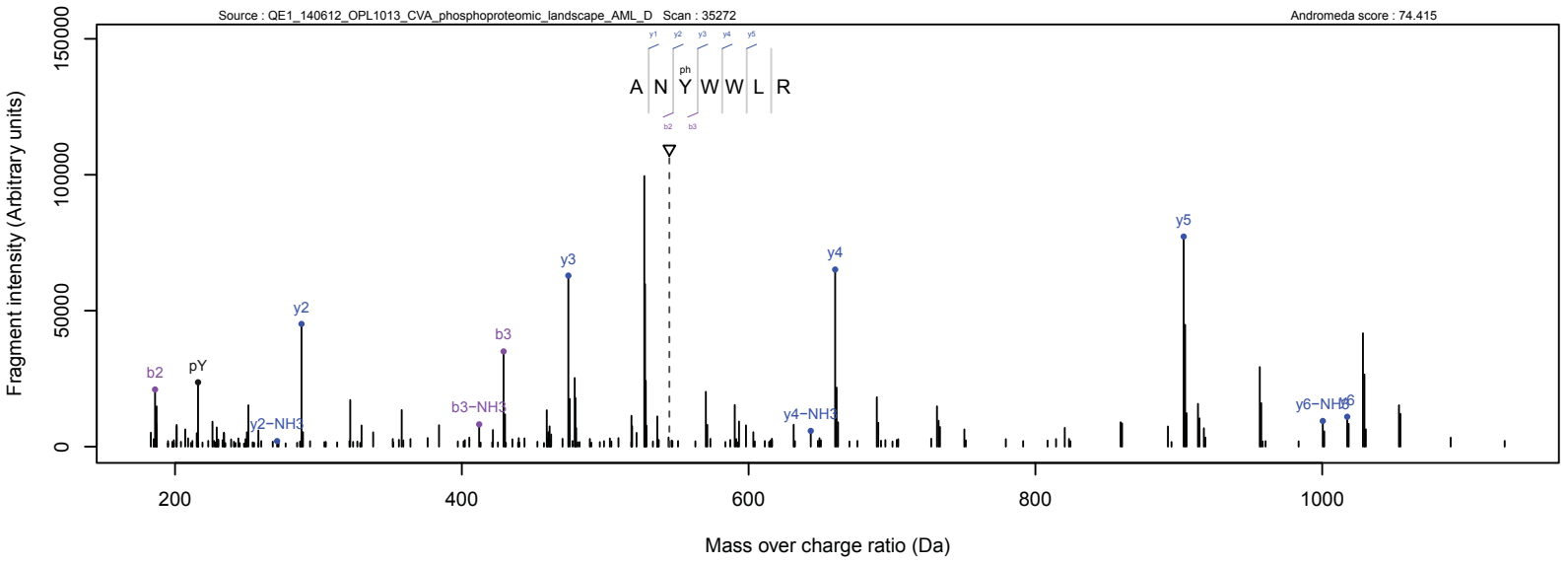




ANY(ph)WWLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 35272

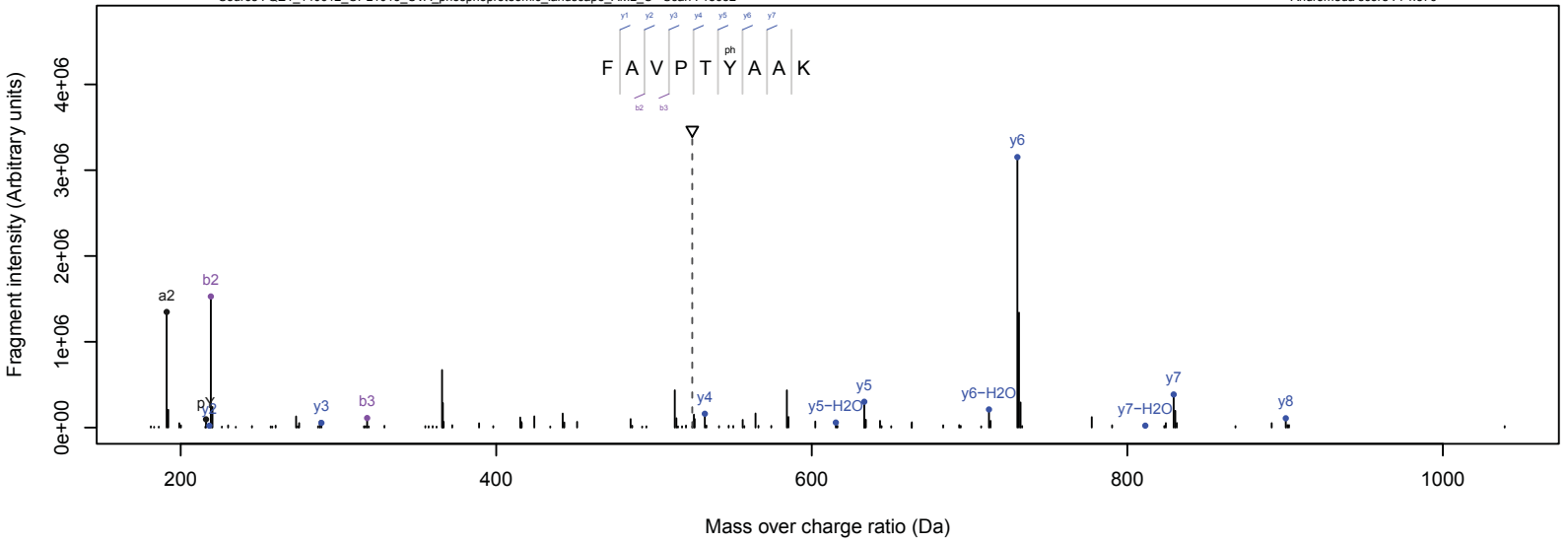
Andromeda score: 74.415



FAVPTY(ph)AAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18652

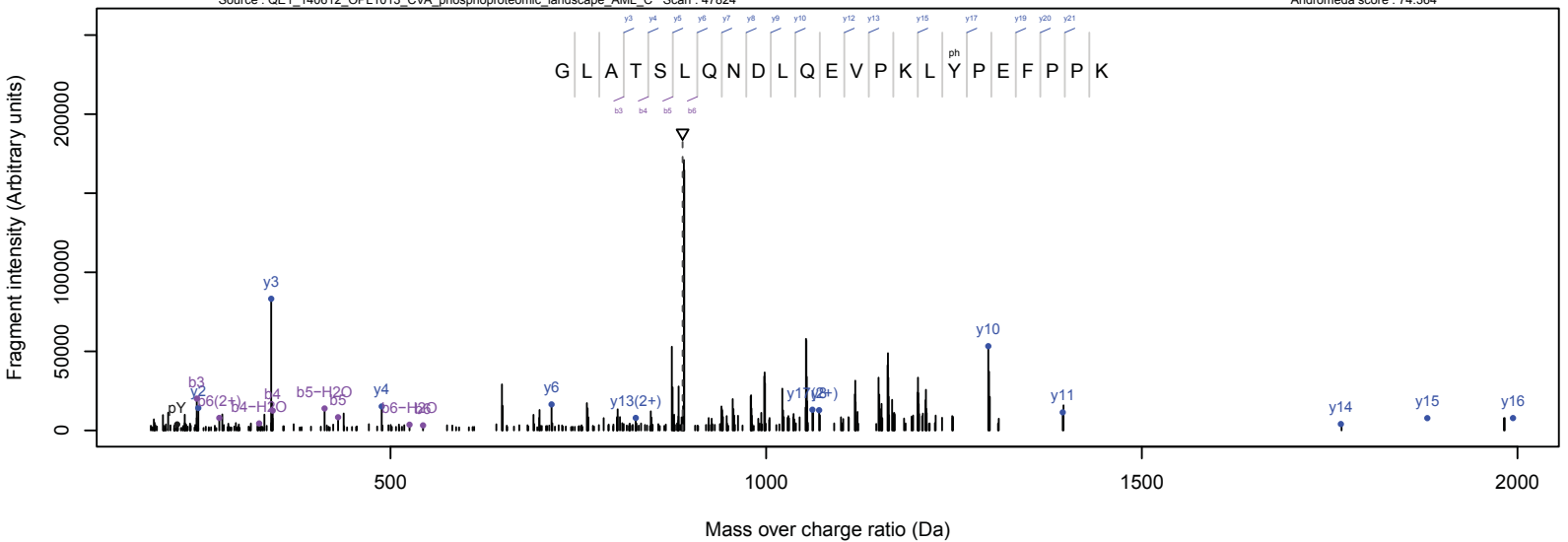
Andromeda score: 74.376



GLATSLQNDLQEVKLYPEFPK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 47824

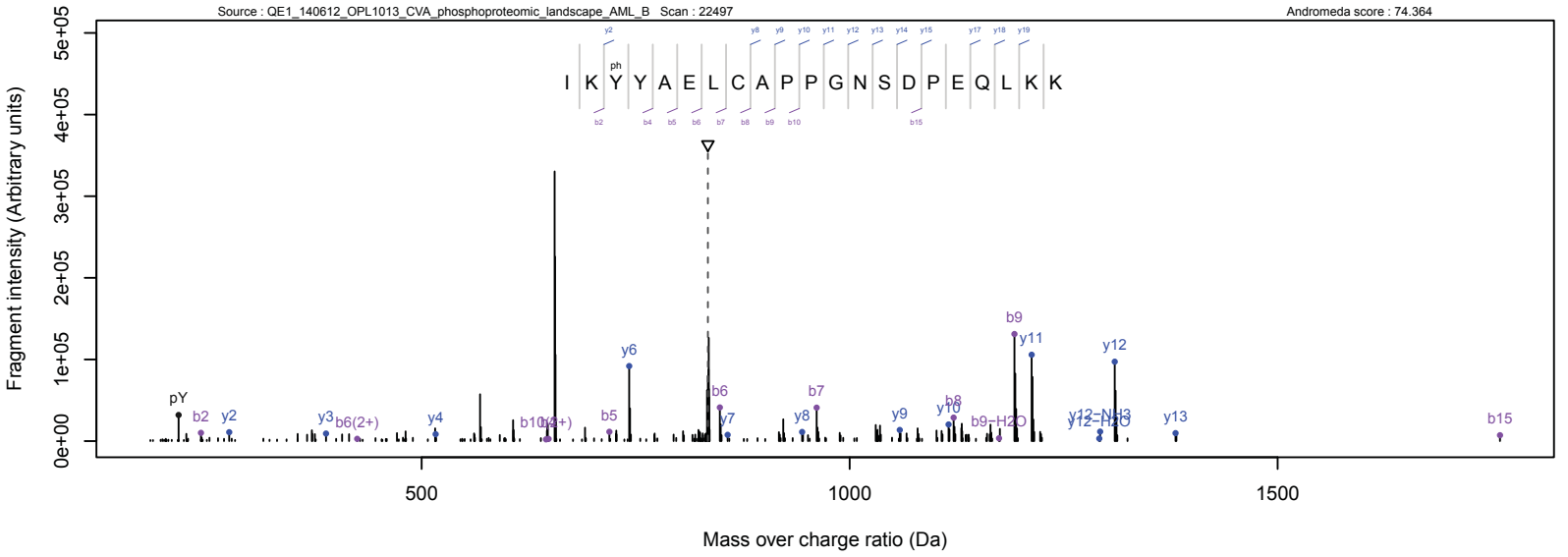
Andromeda score: 74.364



**IKY(ph)YALCAPPGNSDPEQLKK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22497

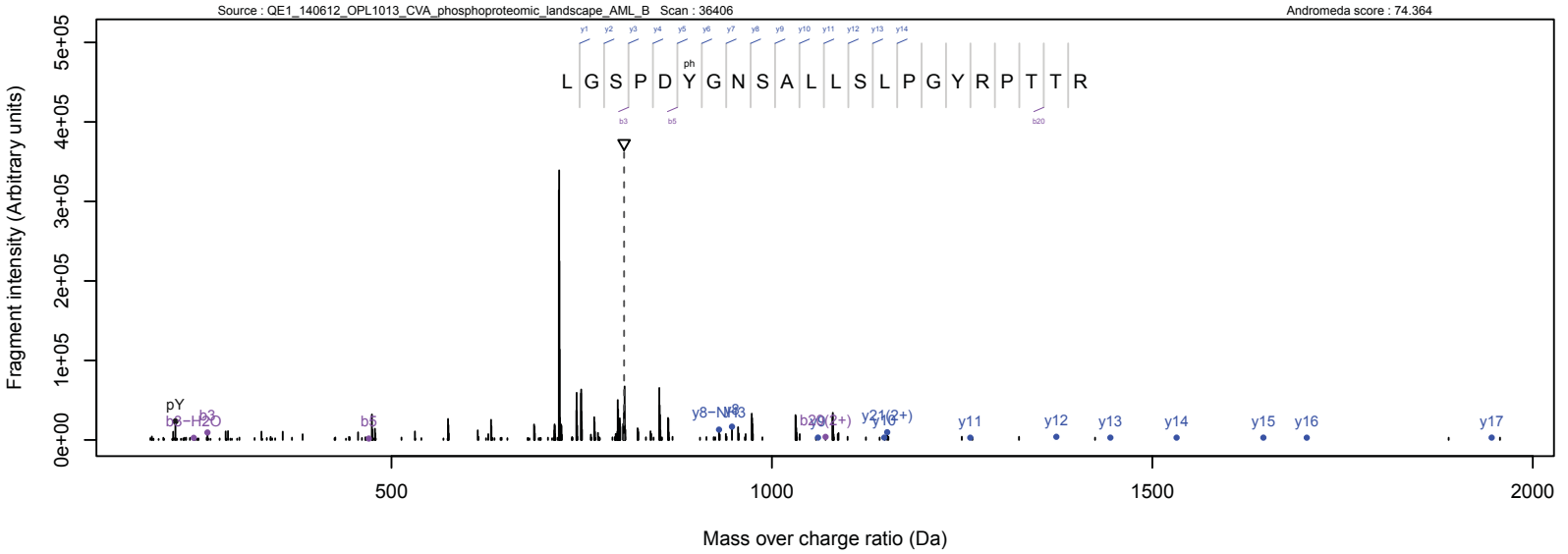
Andromeda score : 74.364



**LGSPDY(ph)GNSALLSLPGYRPTTR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36406

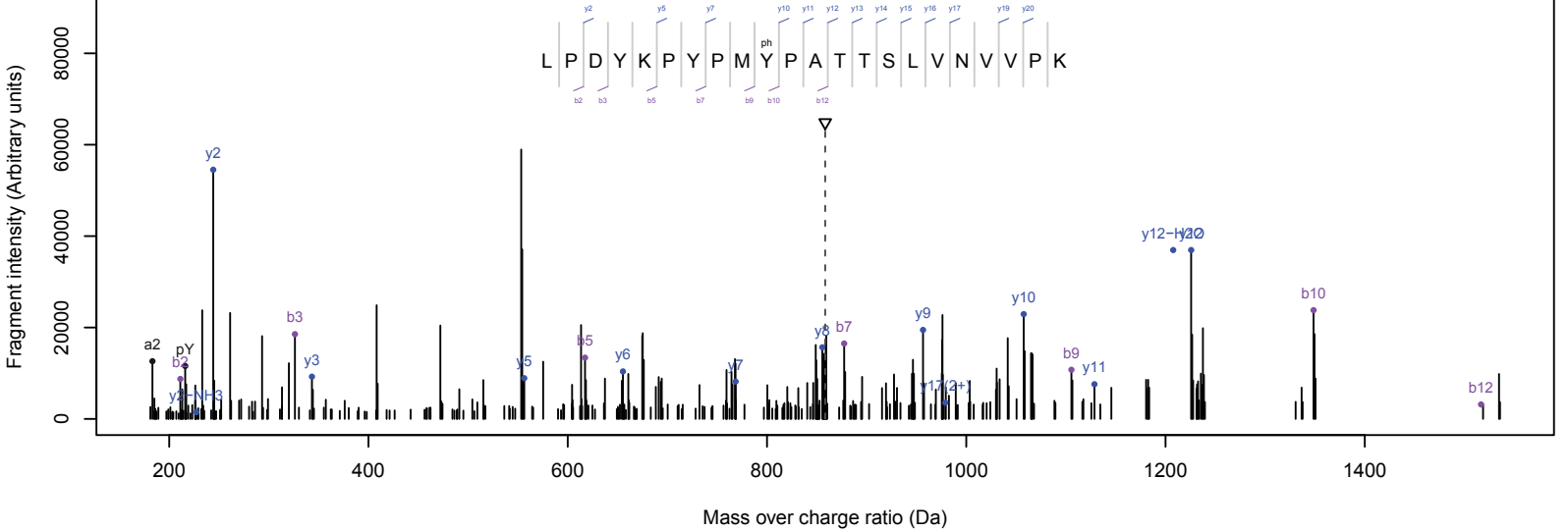
Andromeda score : 74.364



**LPDYKPYPMY(ph)PATTSLVNVVPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43618

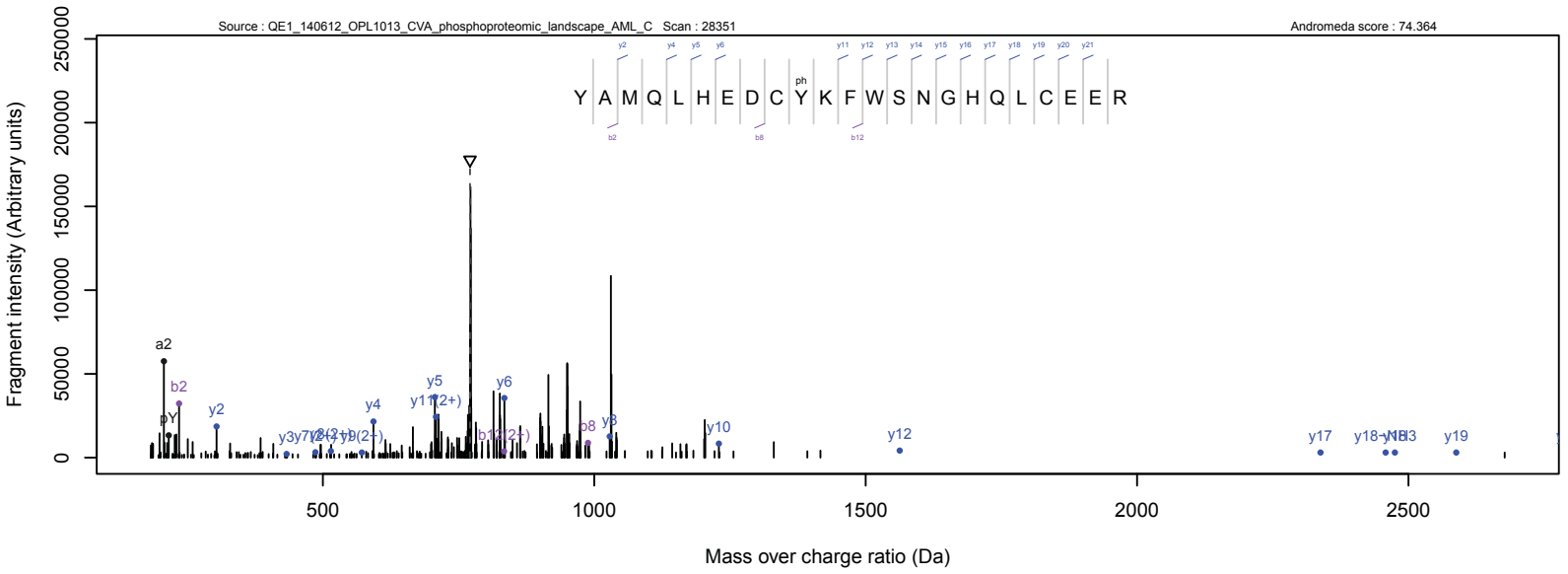
Andromeda score : 74.364



**YAMQLHEDCY(ph)KFWWSNGHQLCEER-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28351

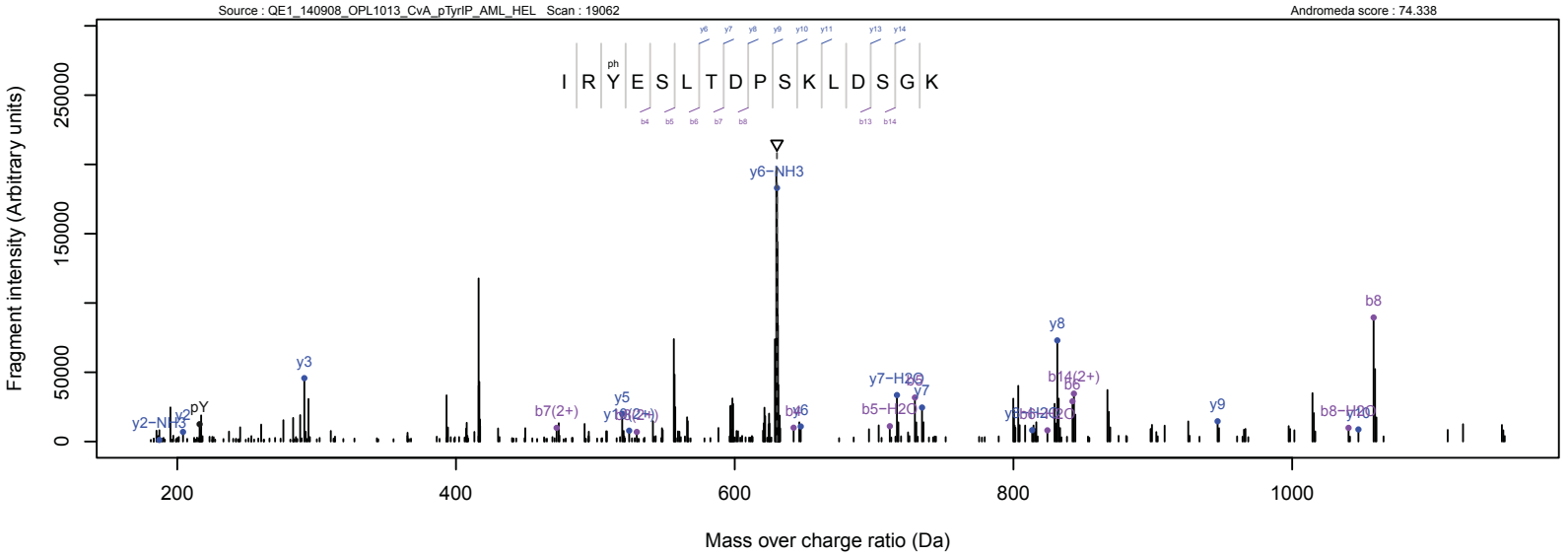
Andromeda score : 74.364



**IRY(ph)ESLTDP SKLD SGK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 19062

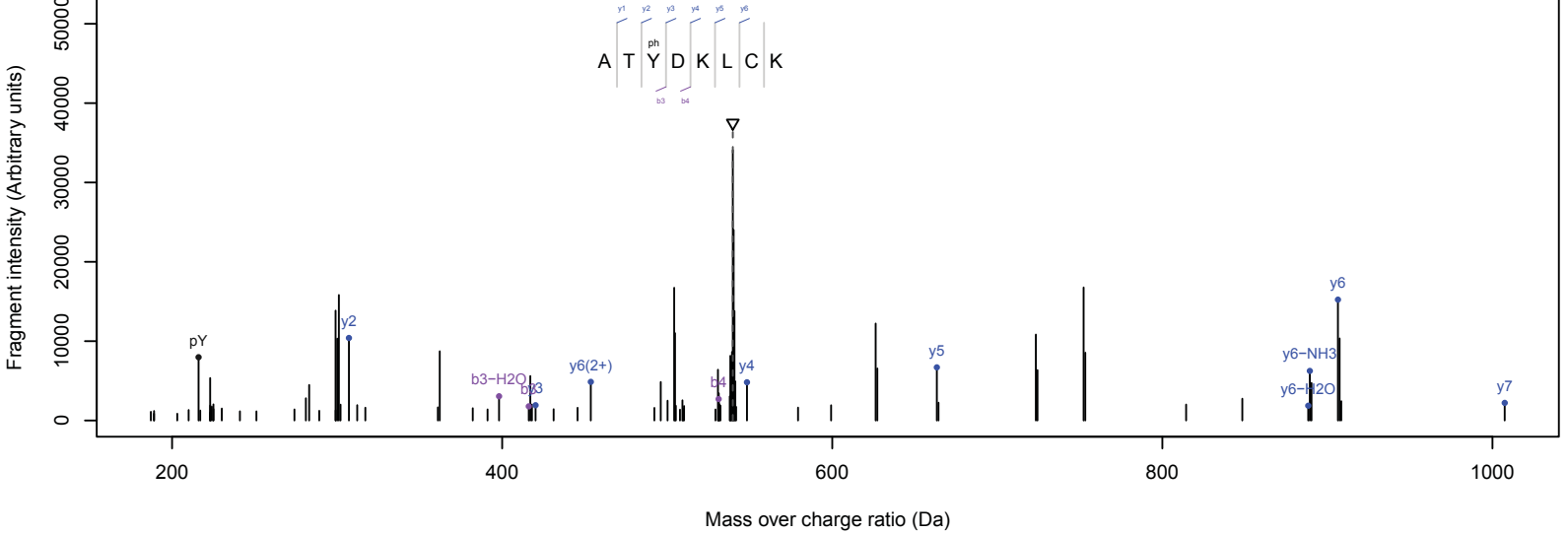
Andromeda score : 74.338



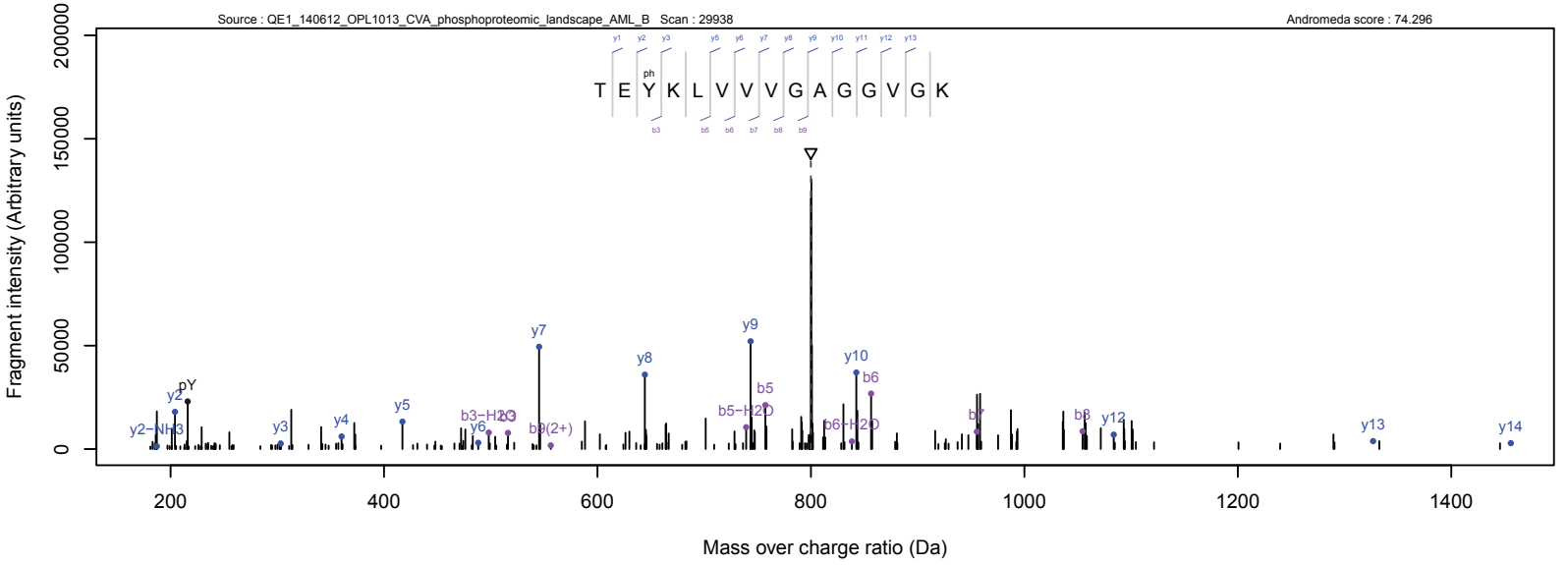
**ATY(ph)DKLCK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3046

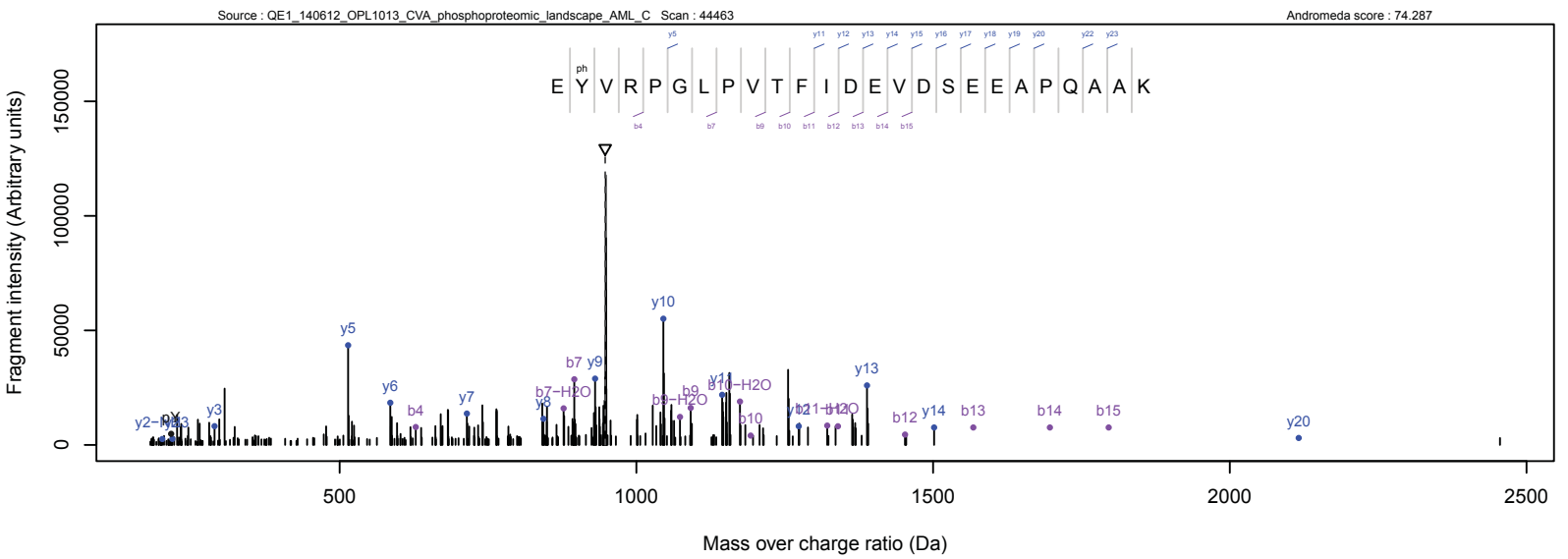
Andromeda score : 74.301



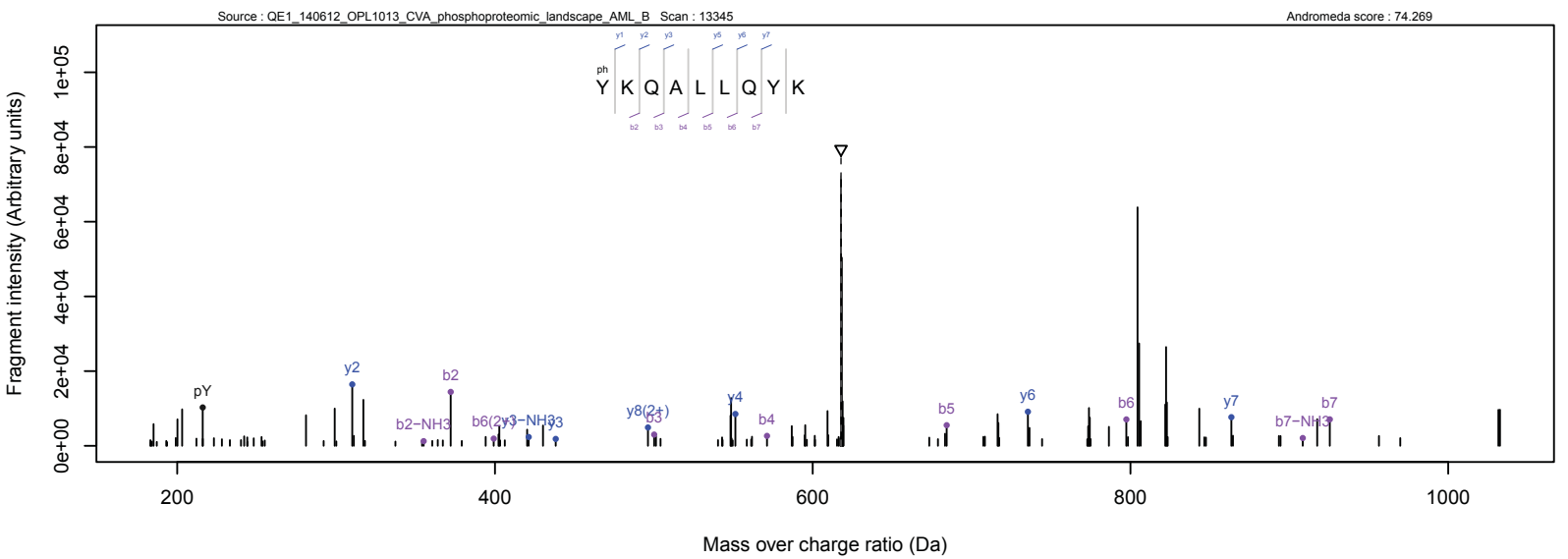
(ac)TEY(ph)KLVVVVGAGGVGK-2+



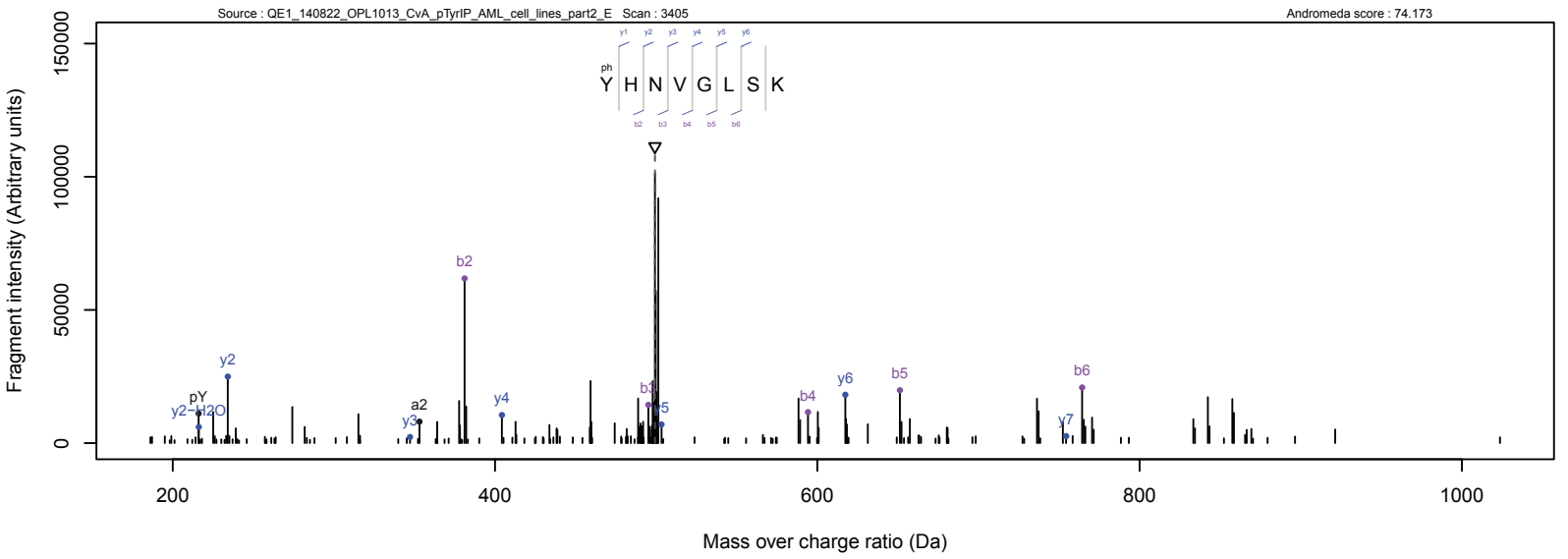
EY(ph)VRGPLVTFIDEVDSEEAPQAAK-3+



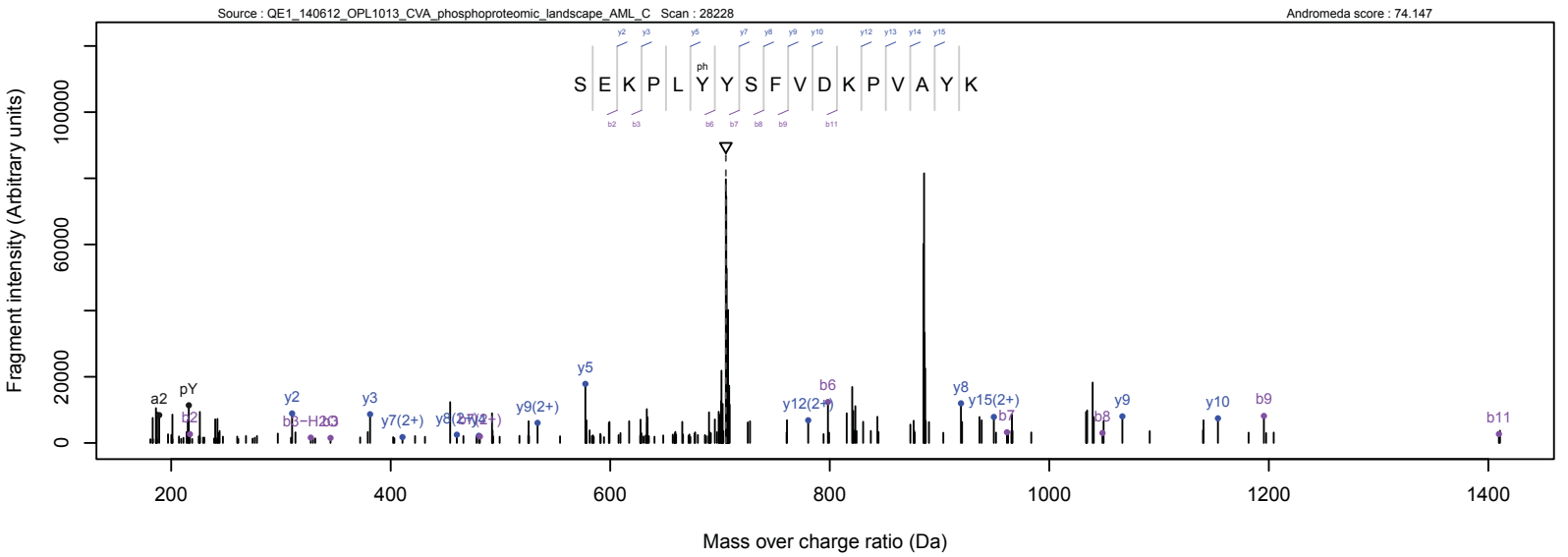
Y(ph)KQALLQYK-2+



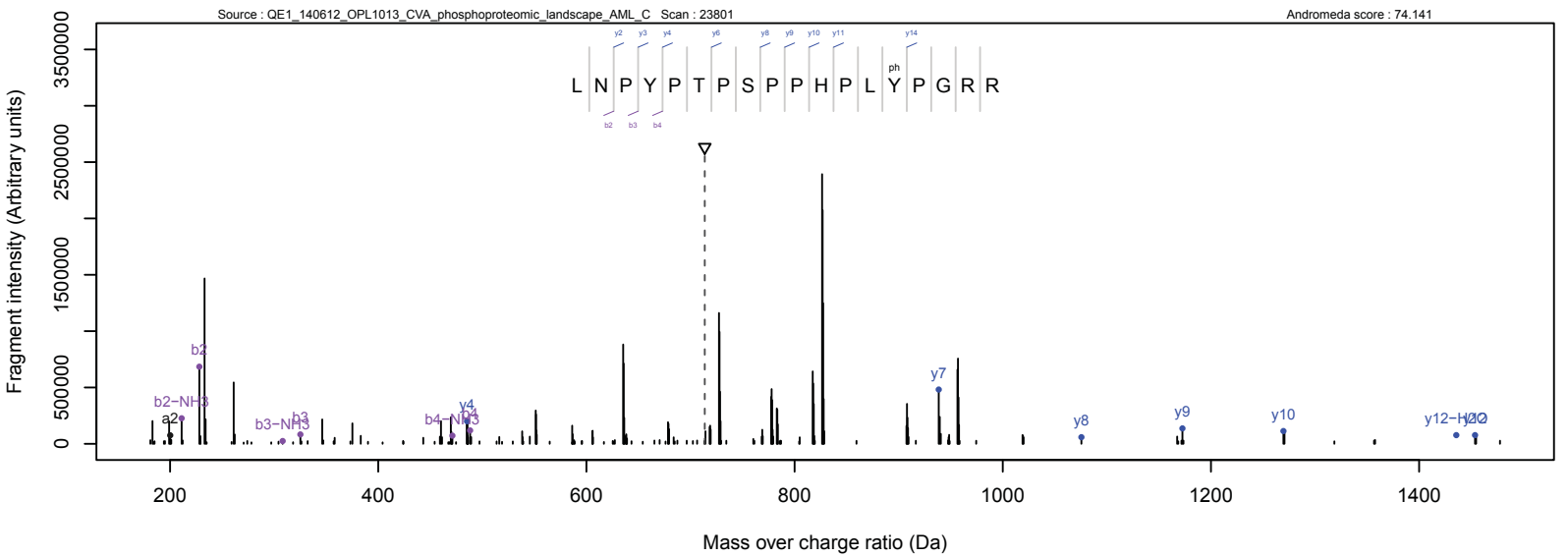
Y(ph)HNVGLSK-2+



SEKPLY(ph)YSFVDKPVAYK-3+



LNPYPTSPHPPLY(ph)PGRR-3+

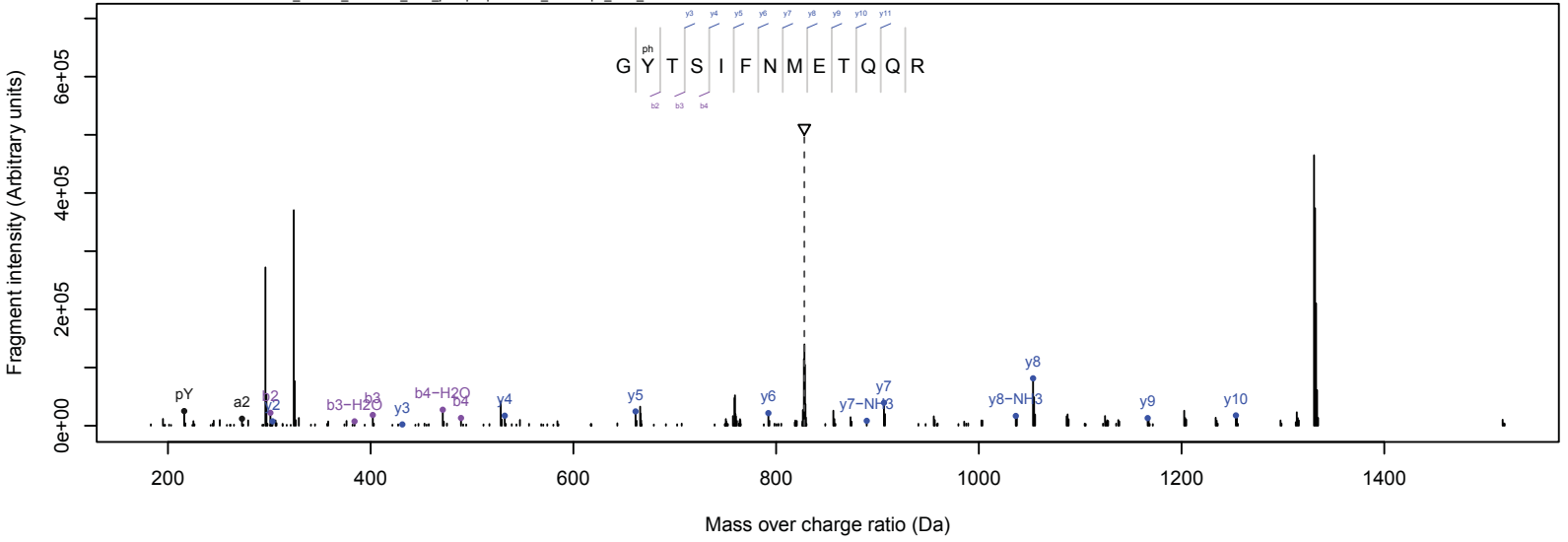




GY(ph)TSIFNMETQQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29595

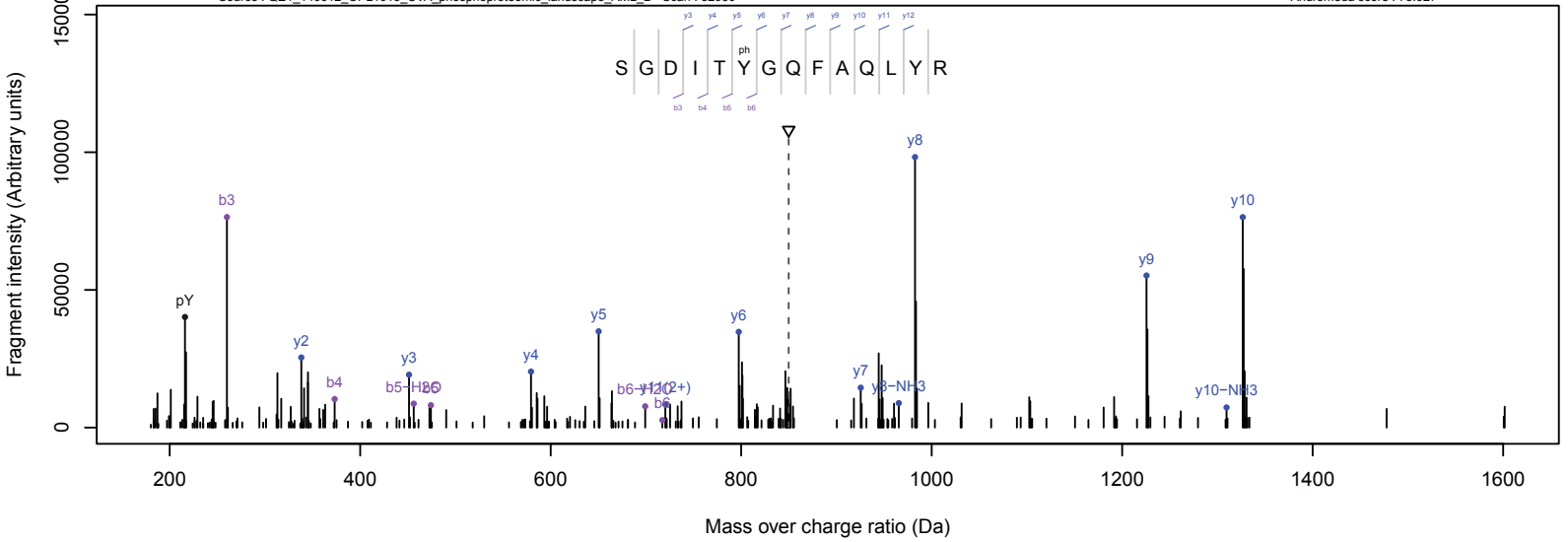
Andromeda score : 73.927



SGDITY(ph)GQFAQLYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32950

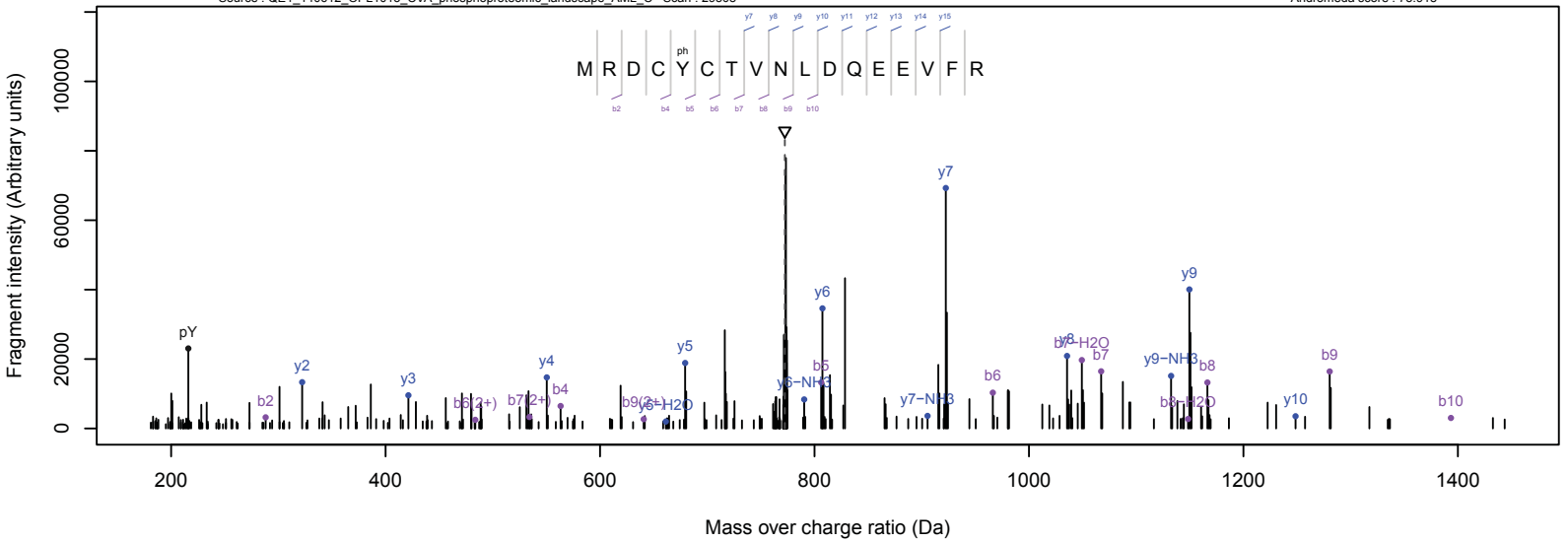
Andromeda score : 73.927



MRDCY(ph)CTVNLDEEVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29508

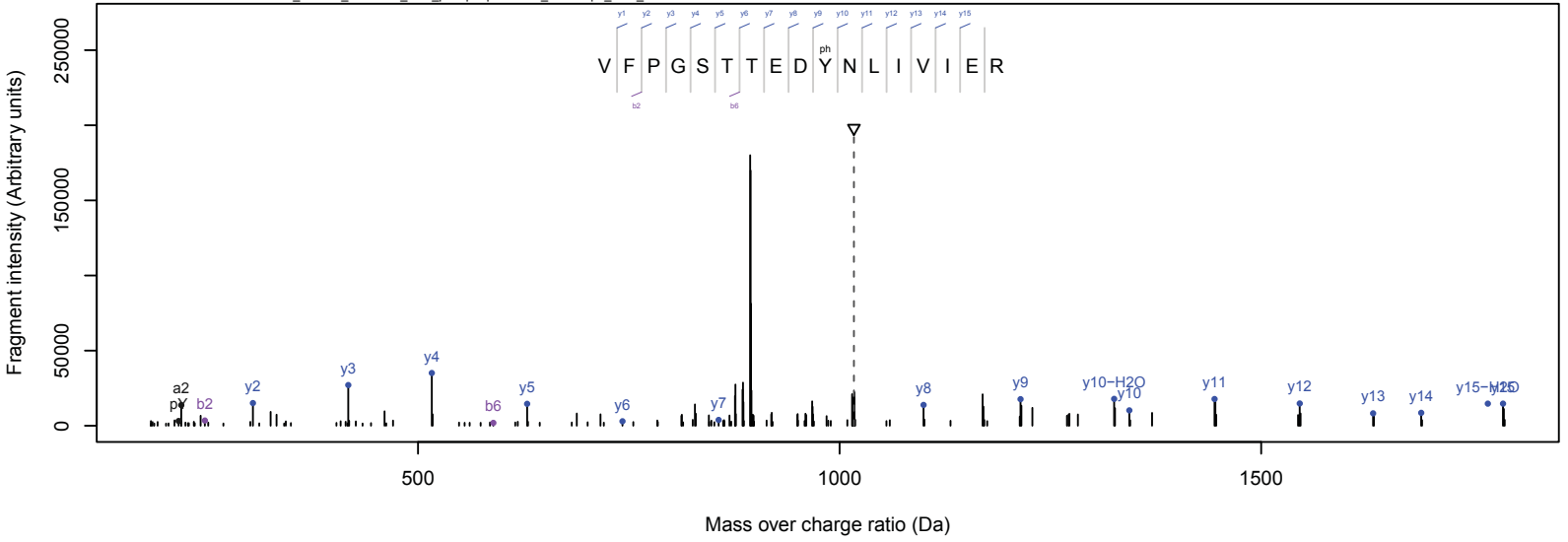
Andromeda score : 73.918



VFPGSTEDY(ph)NLIVIER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40509

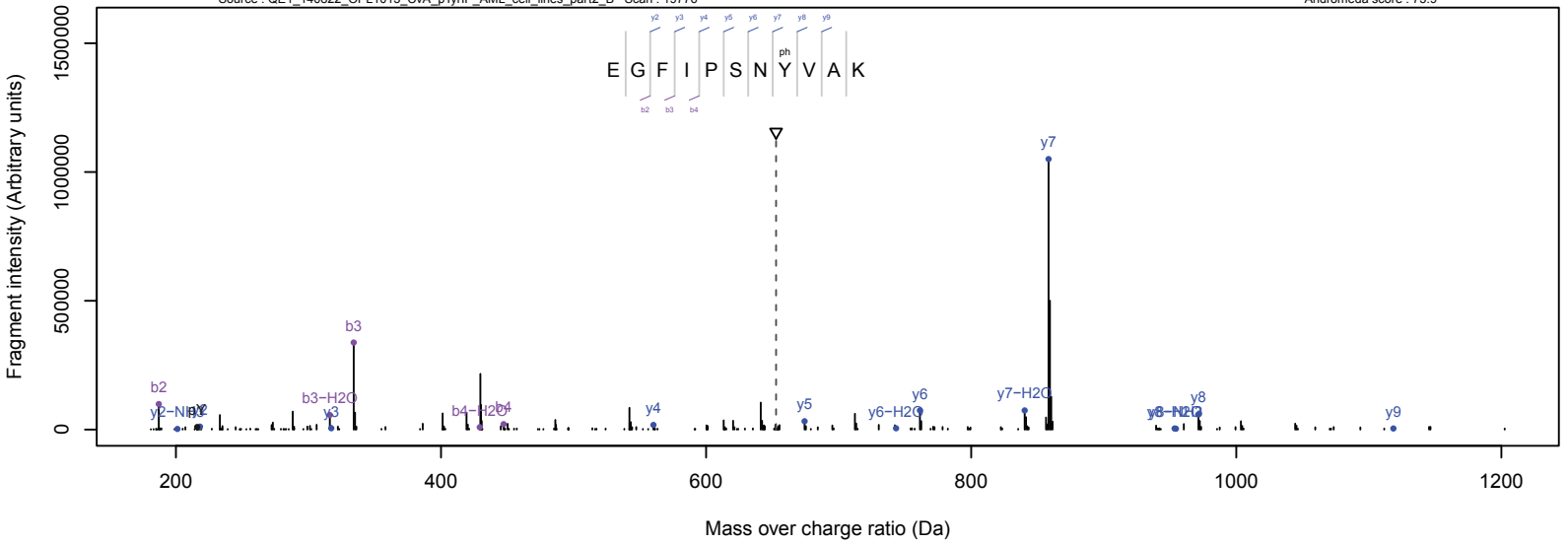
Andromeda score : 73.918



EGFIPSNY(ph)VAK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 19776

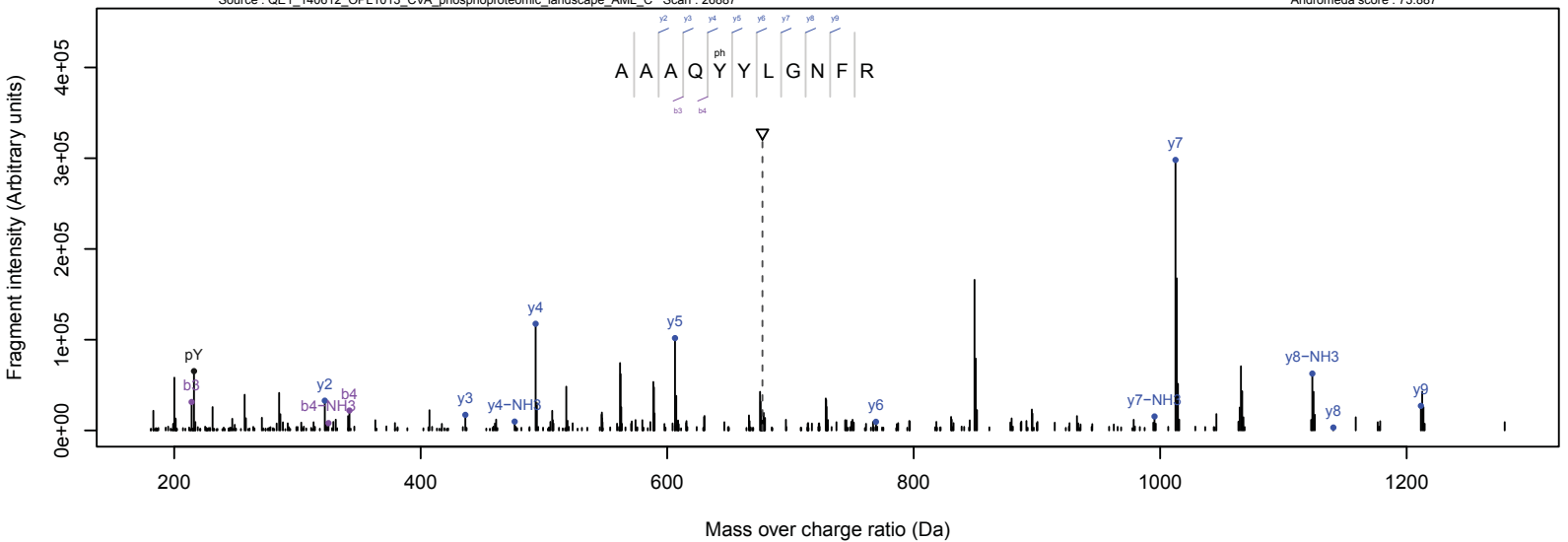
Andromeda score : 73.9



AAAQY(ph)YLGNFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26887

Andromeda score : 73.887

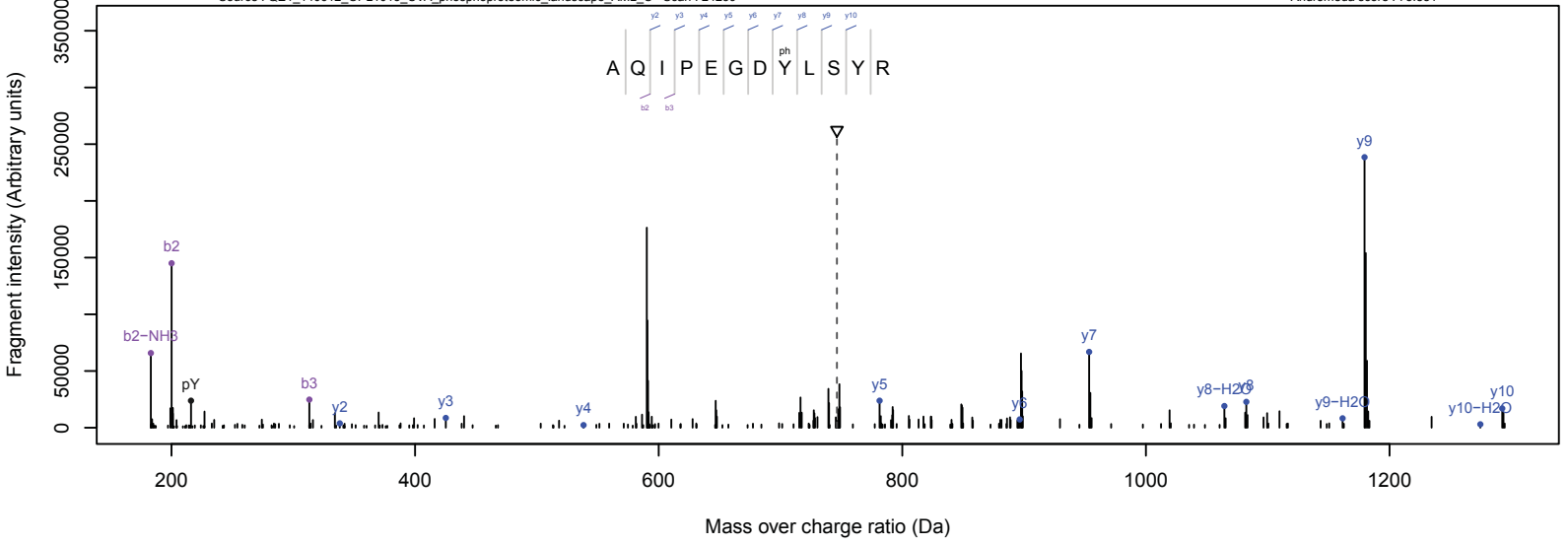




**AQIPEGDY(ph)LSYR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24259

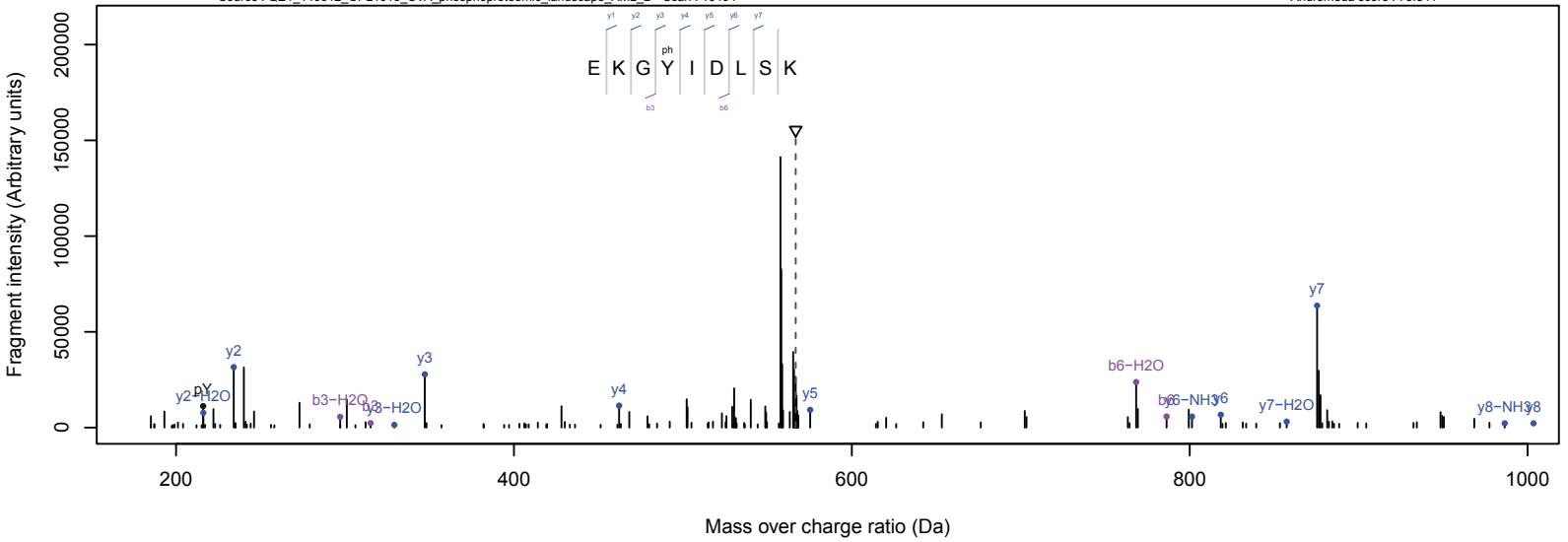
Andromeda score : 73.881



**EKGYP(h)IDLSK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 10494

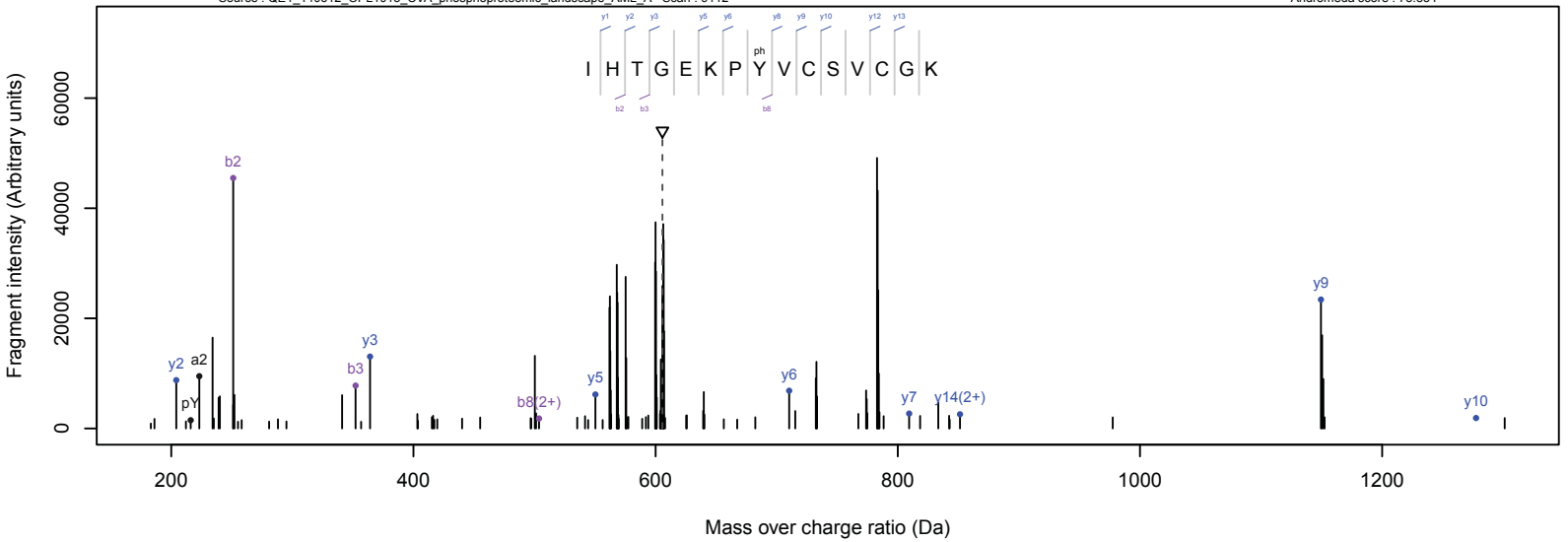
Andromeda score : 73.841



**IHTGKPY(ph)VCSVCGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 9112

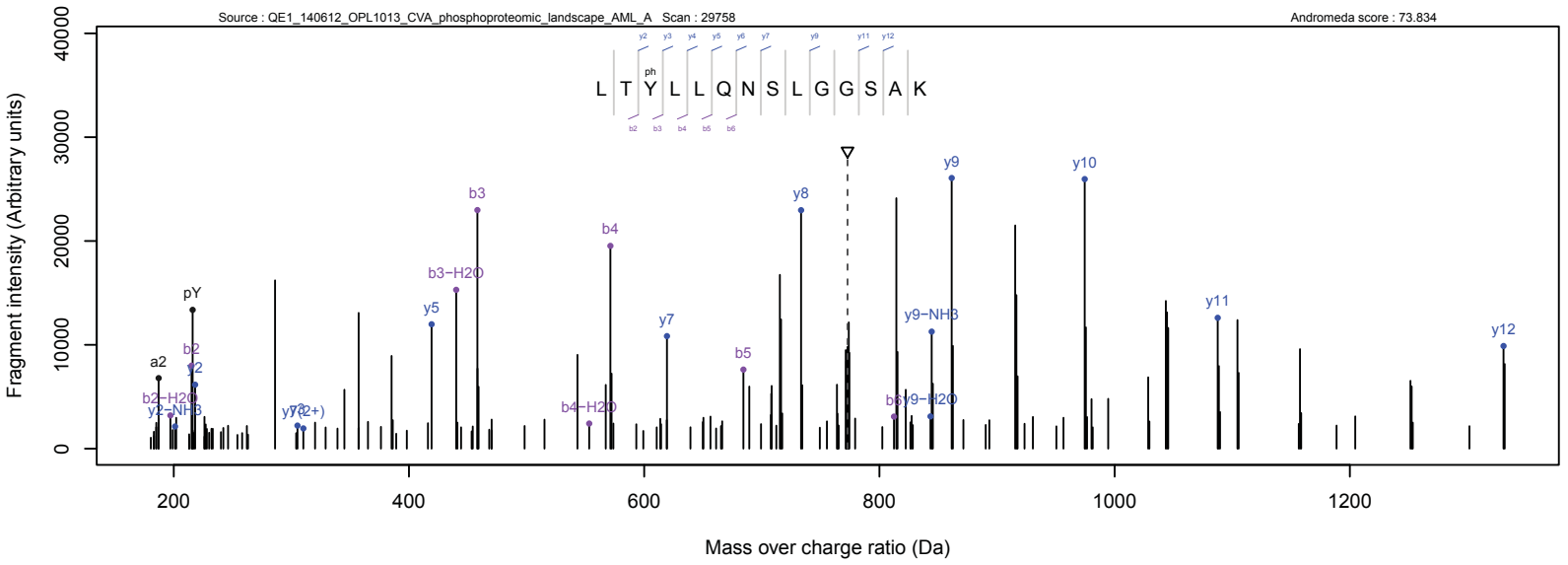
Andromeda score : 73.834



LTY(ph)LLQNSLGGSAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29758

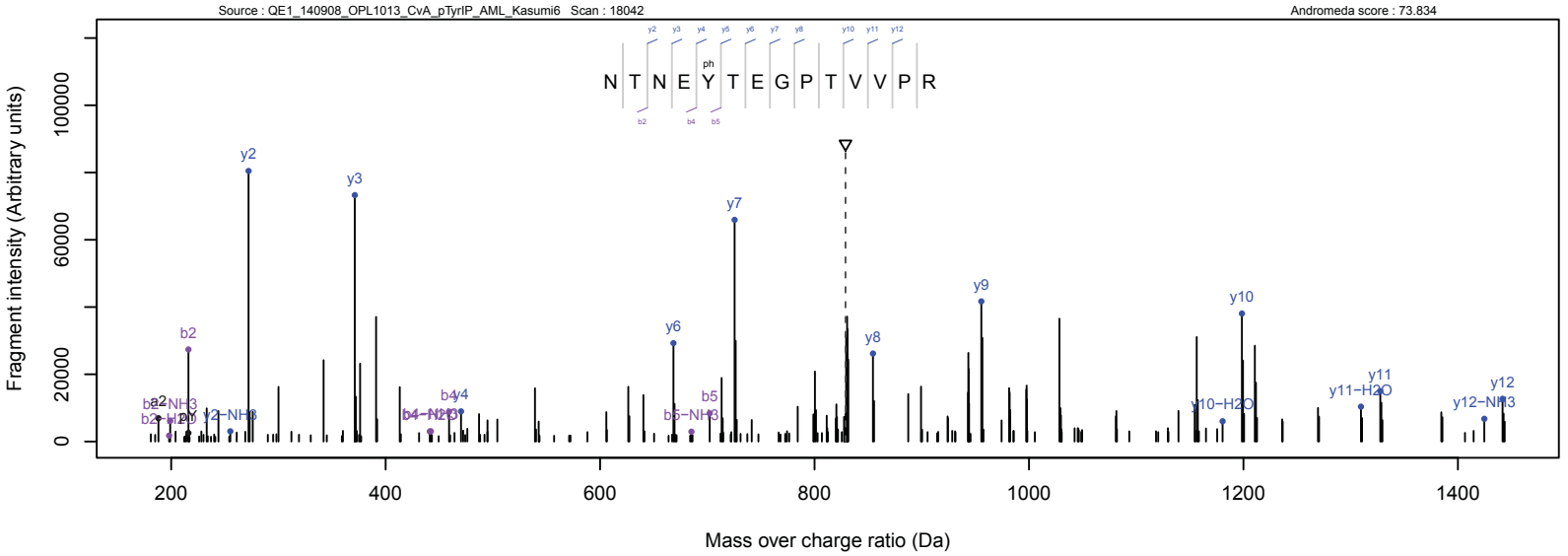
Andromeda score : 73.834



NTNEY(ph)TEGPTVPR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 18042

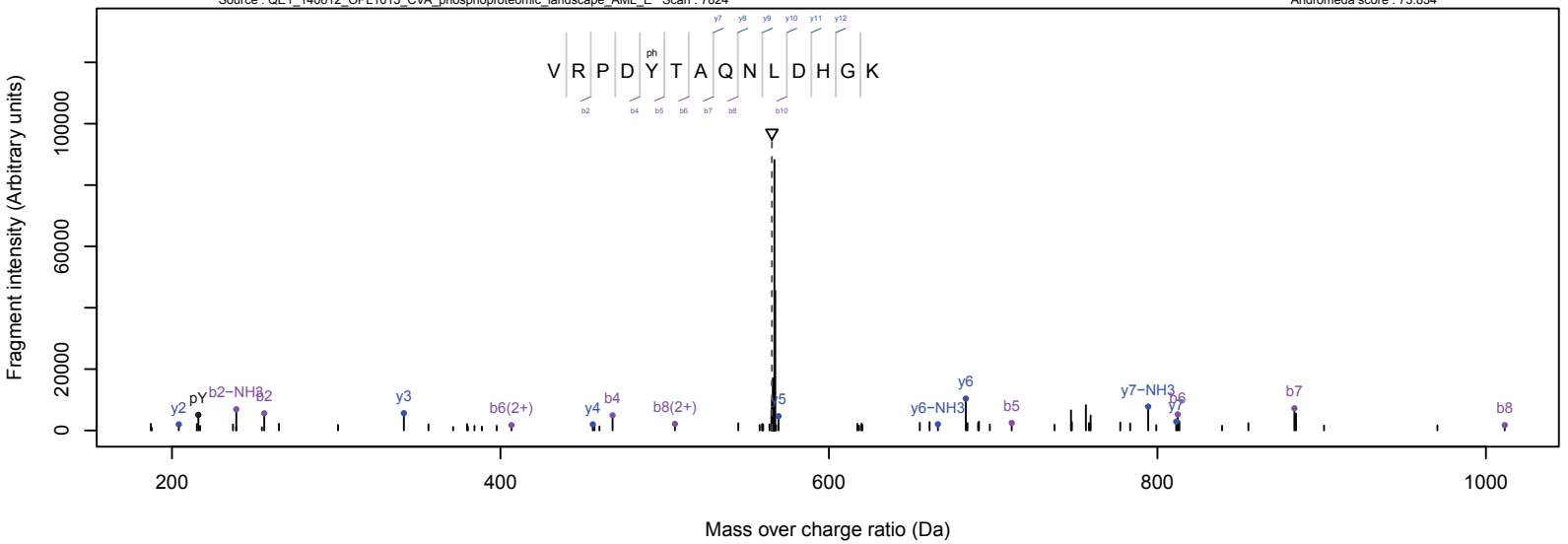
Andromeda score : 73.834



VRPDY(ph)TAQNLDHGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 7824

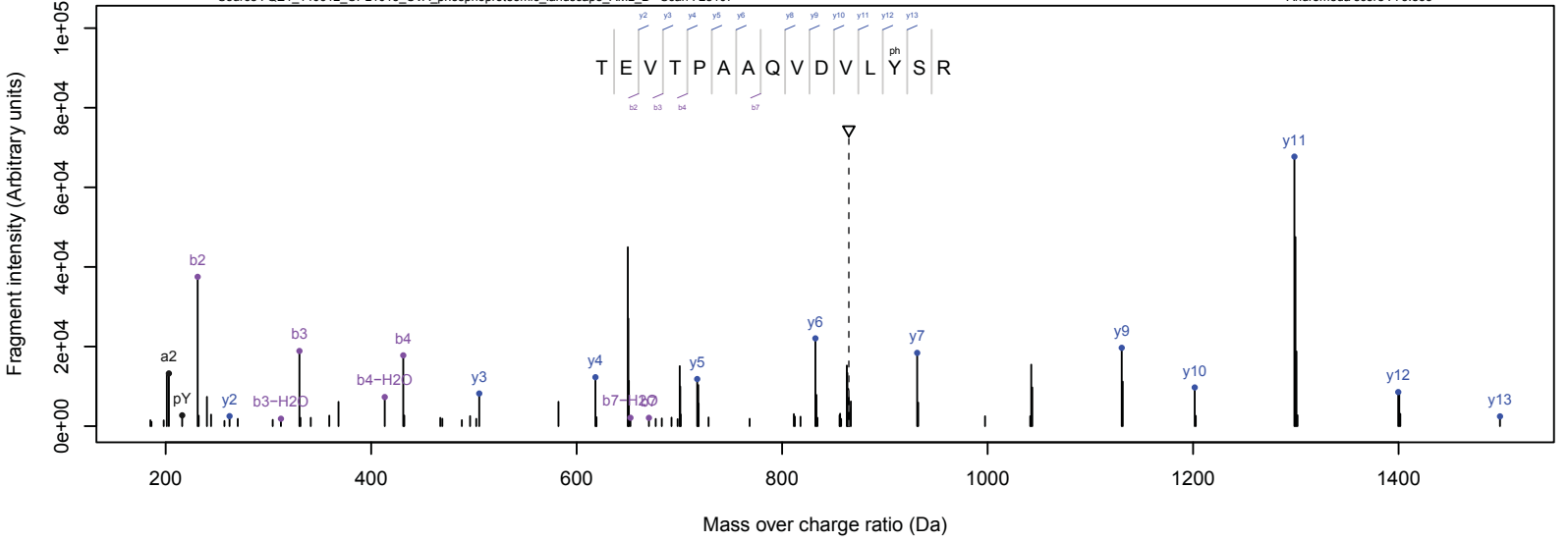
Andromeda score : 73.834



TEVTPAAQVDVLY(ph)SR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28107

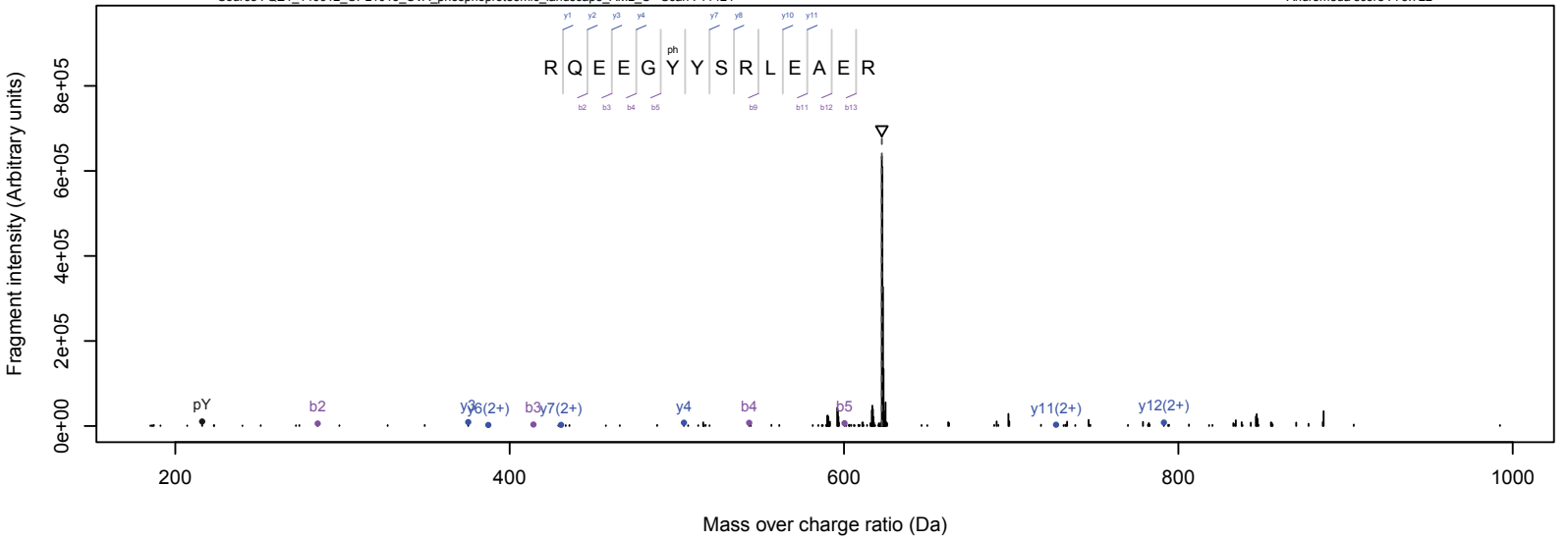
Andromeda score : 73.833



RQEEGY(ph)YSRLEAER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11424

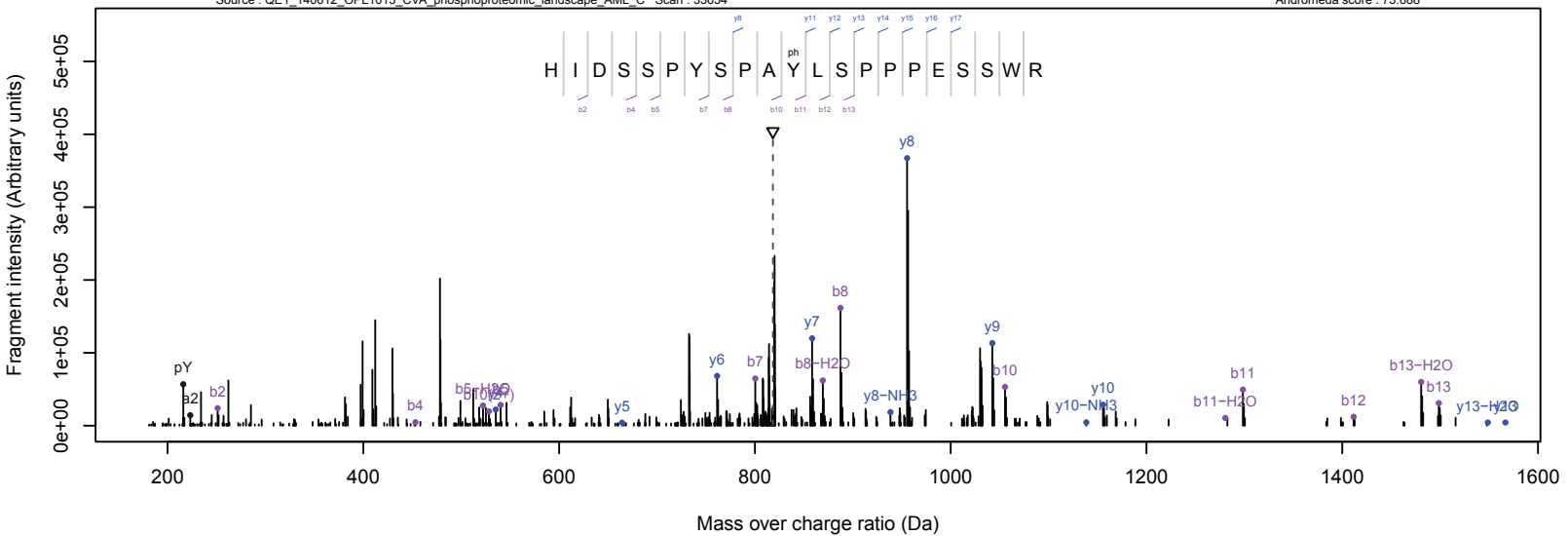
Andromeda score : 73.722



HIDSSPYSPAY(ph)LSPPPESSWR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33634

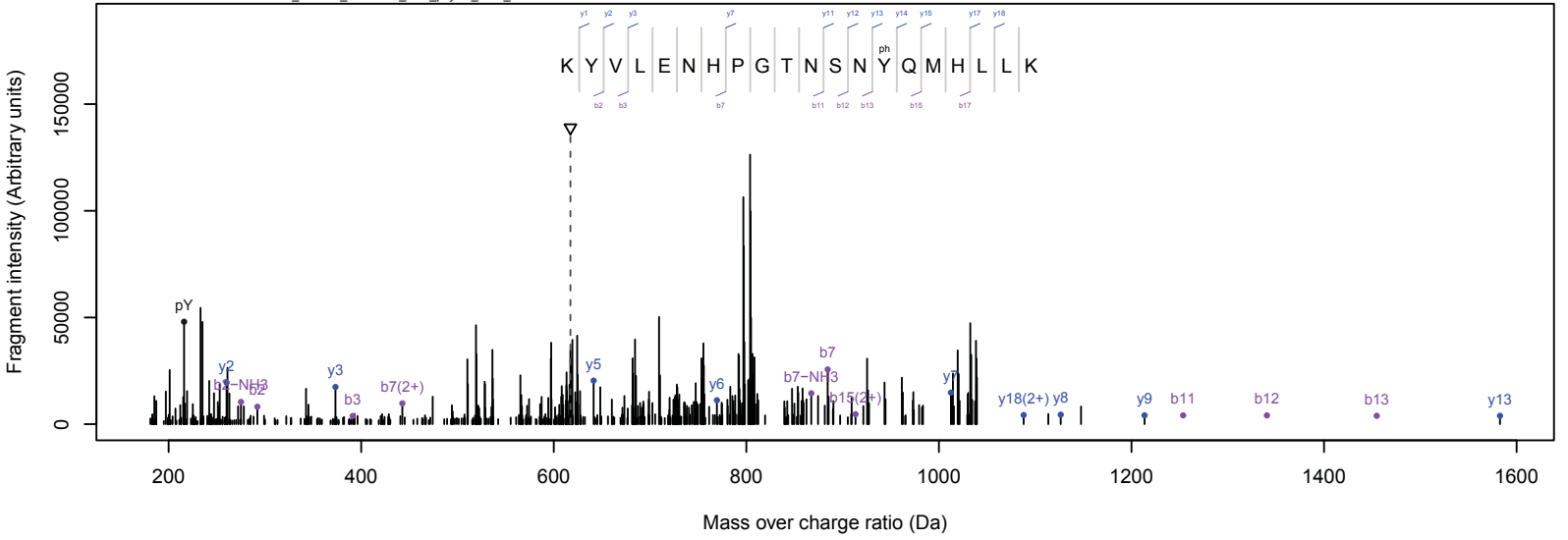
Andromeda score : 73.688



**KYVLENHPTNSNY(ph)QMHLK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 21771

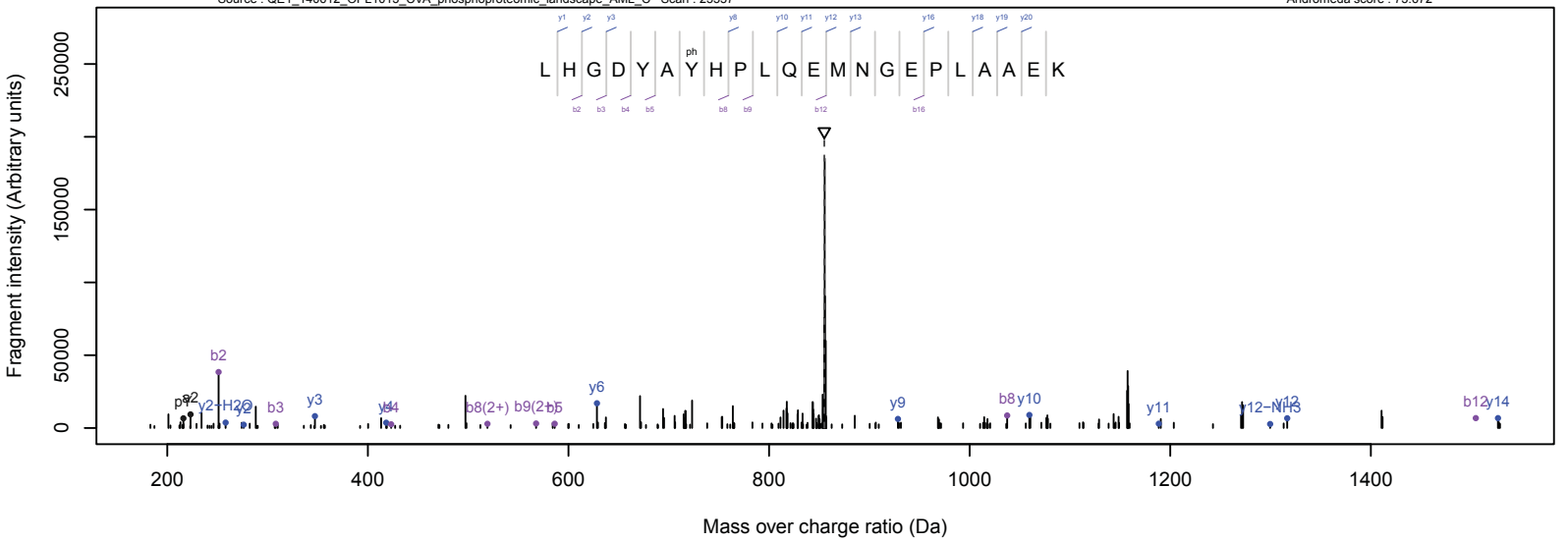
Andromeda score : 73.688



**LHG DYAY(ph)HPLQEMNGEPLAAEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25557

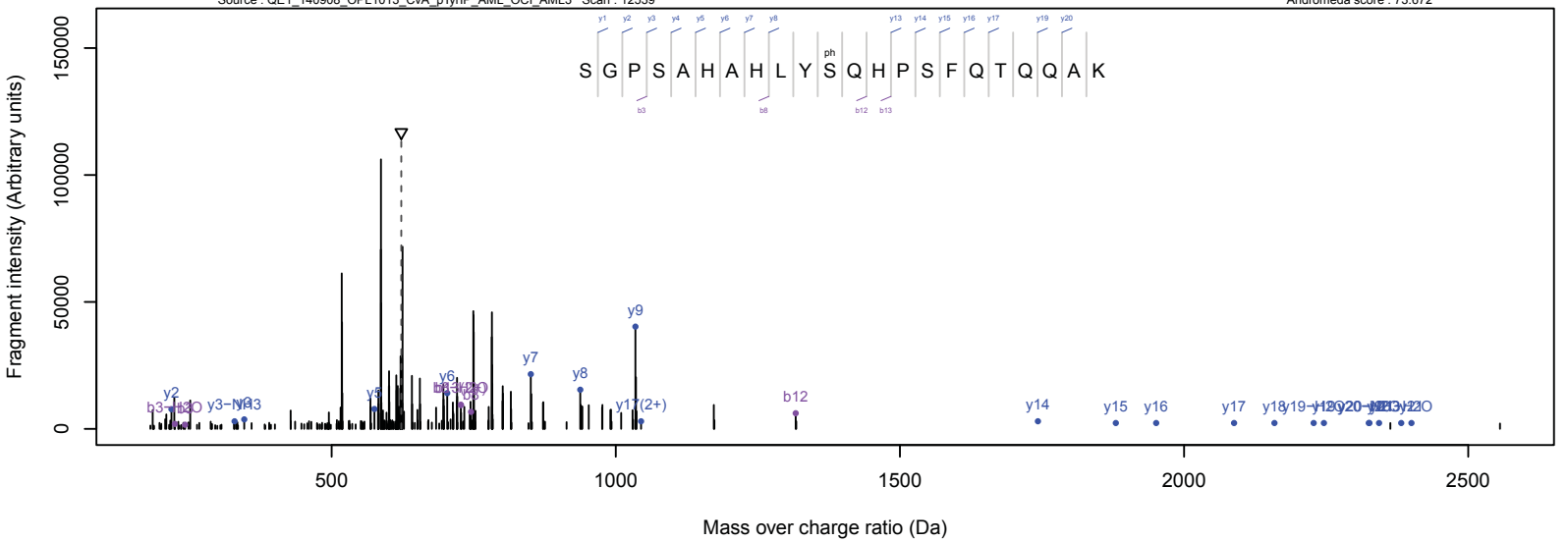
Andromeda score : 73.672



**SGPSAHALYS(ph)QHPSFQTQQA K-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 12539

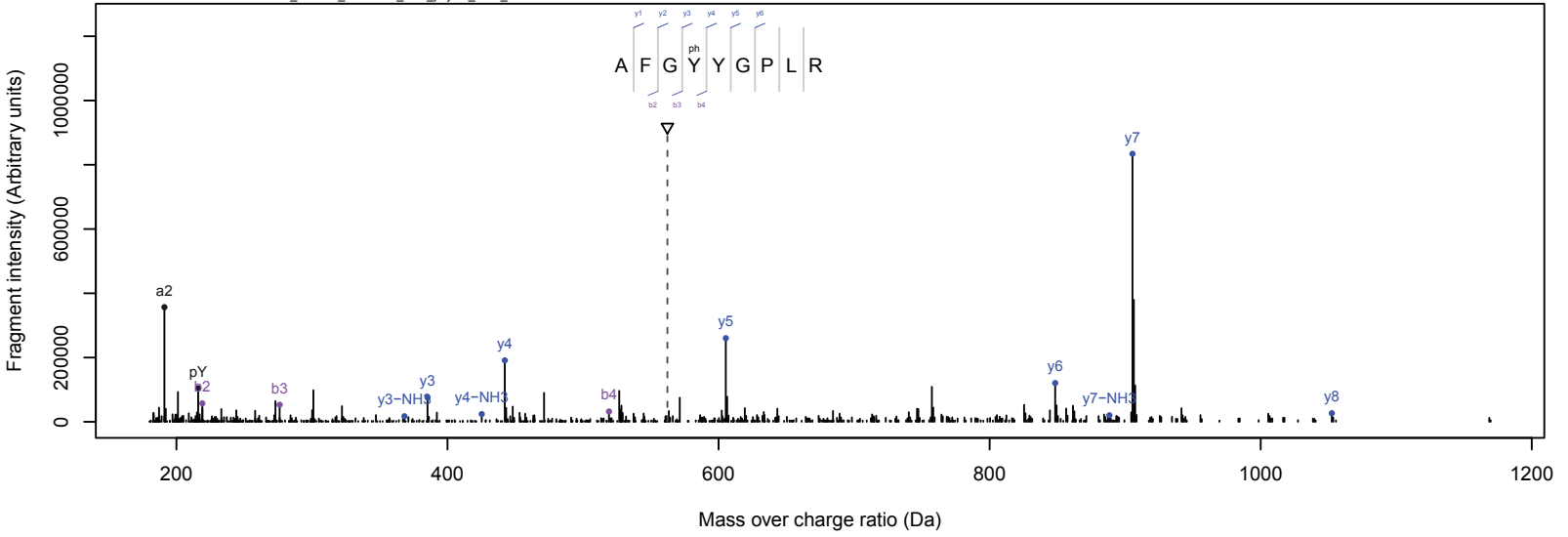
Andromeda score : 73.672



AFGY(ph)YGPLR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 30689

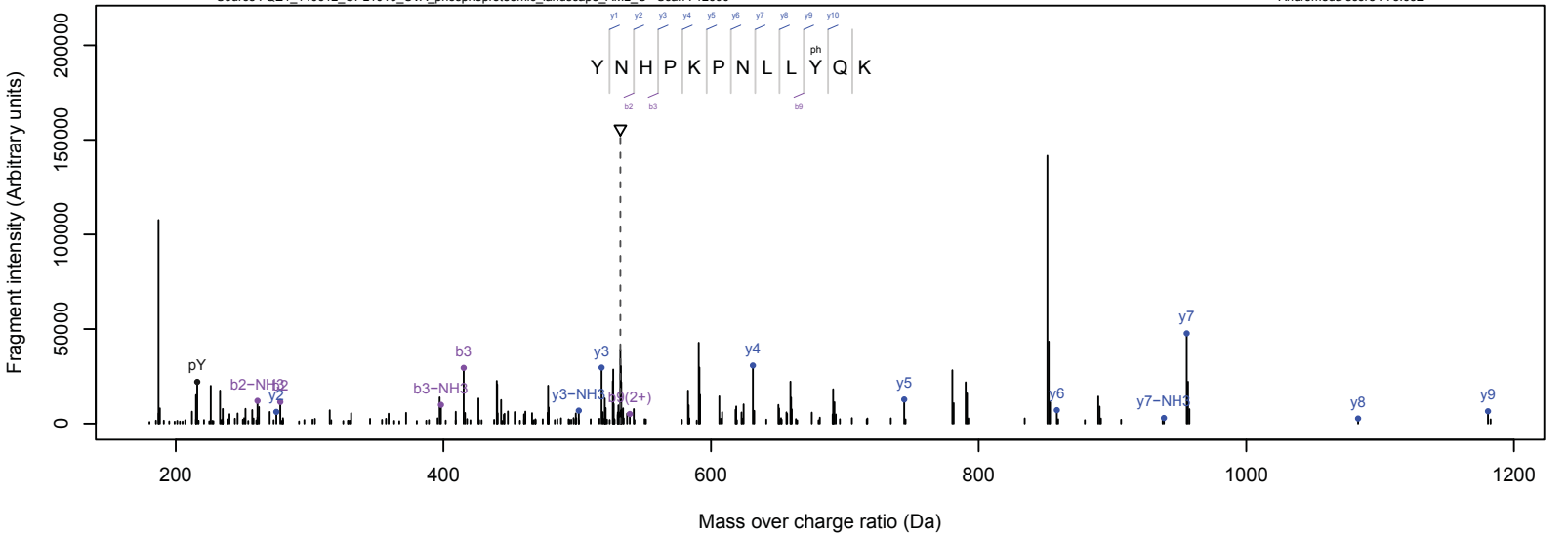
Andromeda score : 73.665



YNHPKNLLY(ph)QK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12690

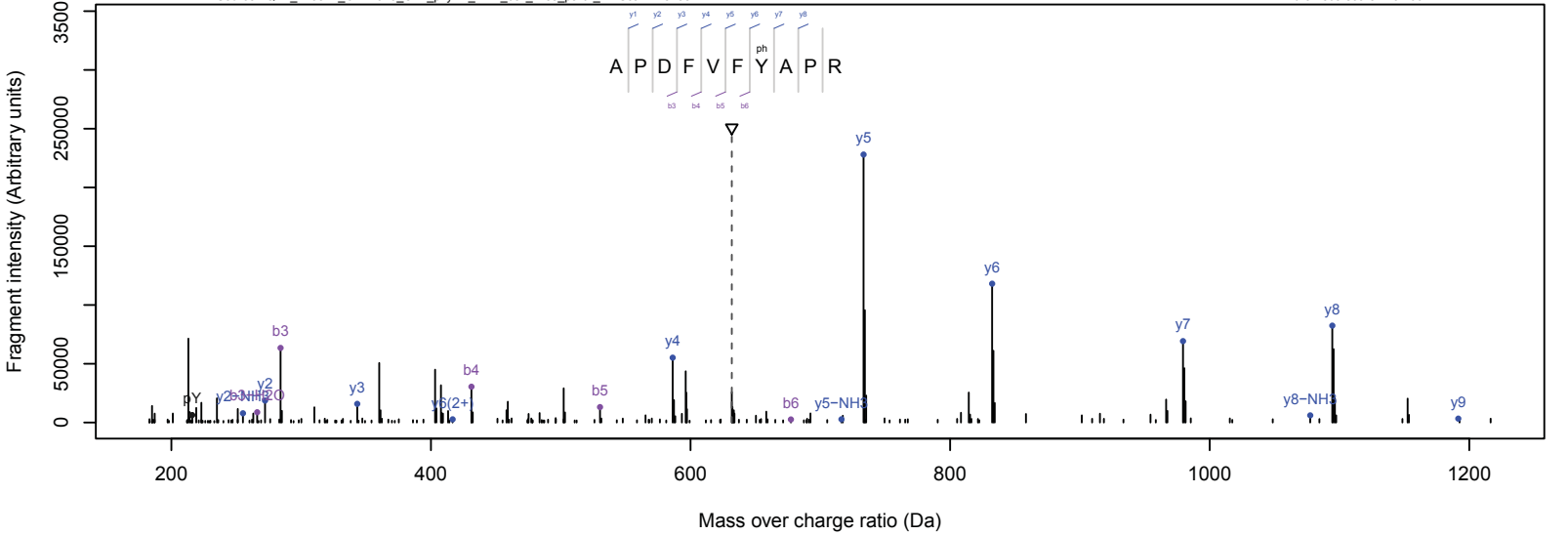
Andromeda score : 73.632



APDFVY(ph)APR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 28235

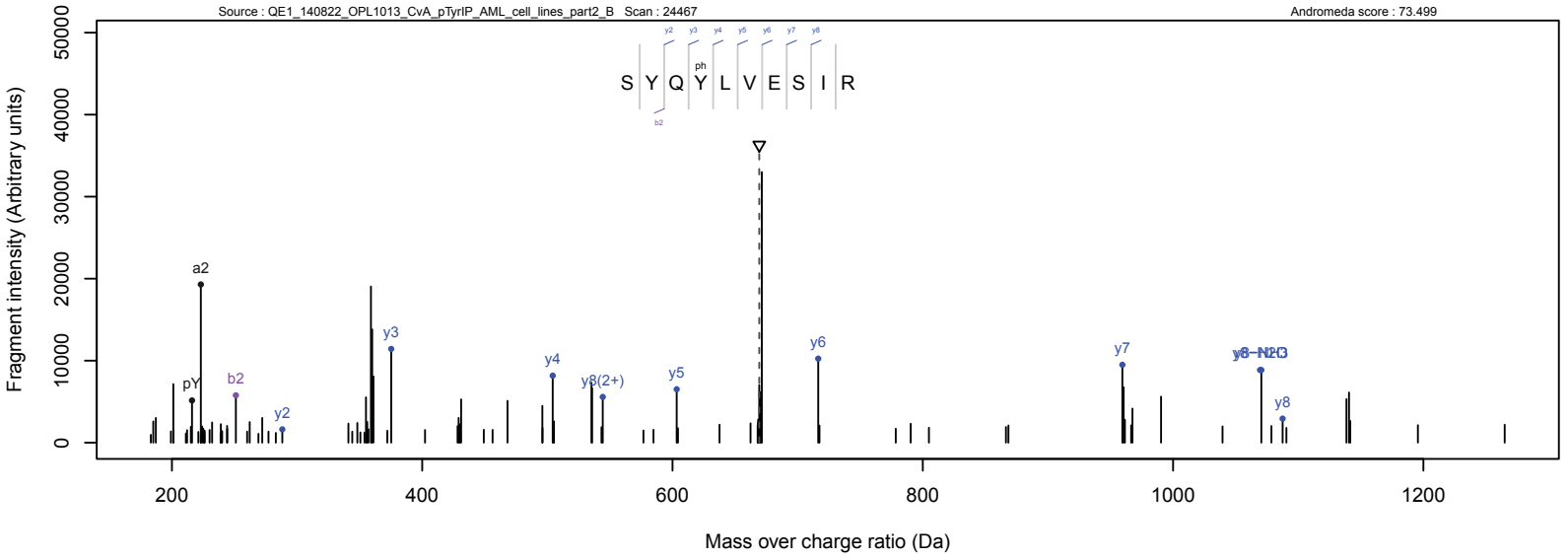
Andromeda score : 73.499



**SYQY(ph)LVESIR-2+**

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 24467

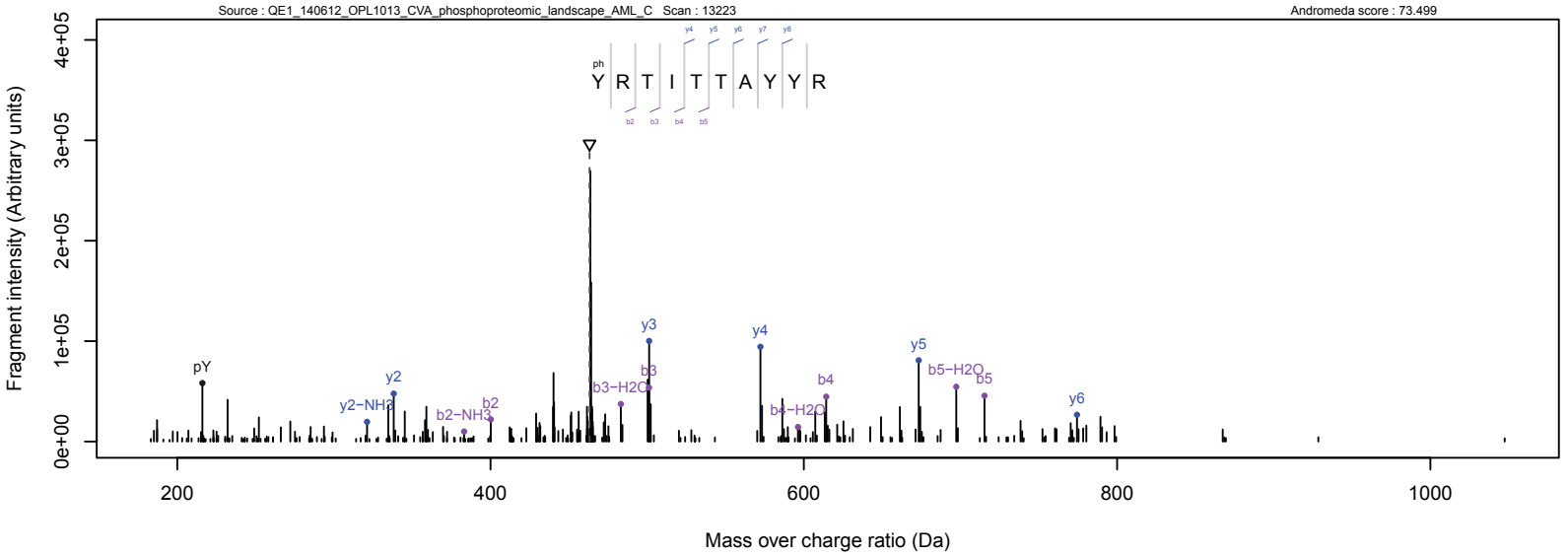
Andromeda score: 73.499



**Y(ph)RTITTAYR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13223

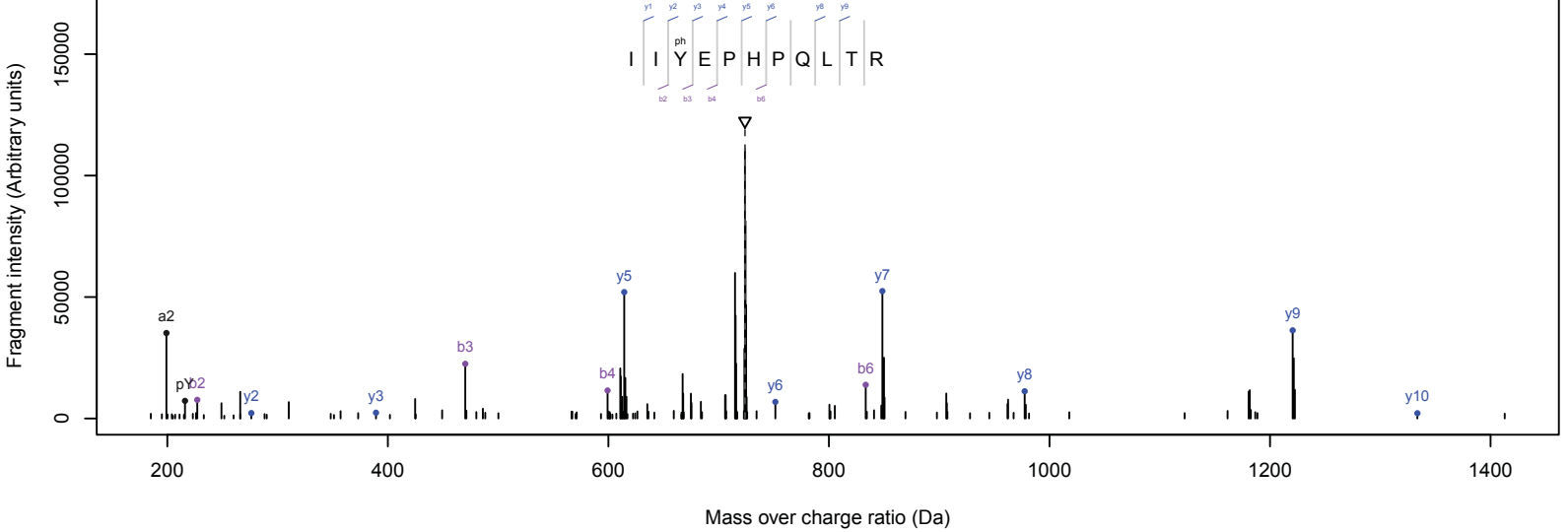
Andromeda score: 73.499



**IY(ph)EPHQLTR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18146

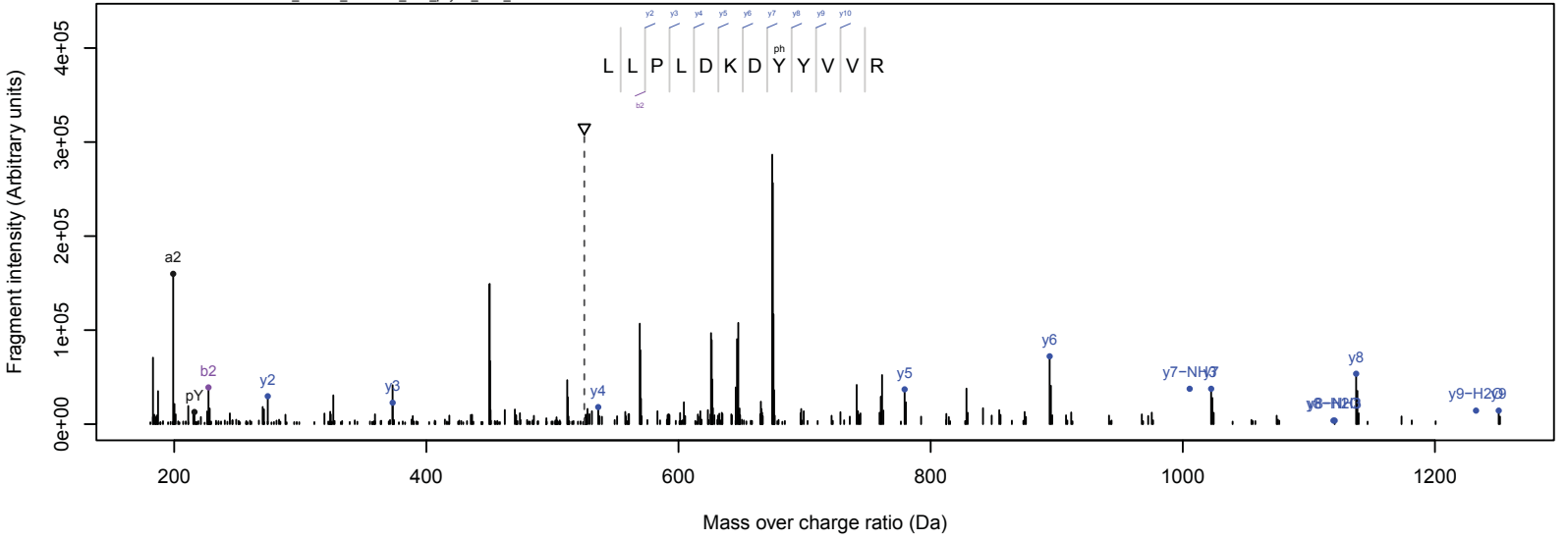
Andromeda score: 73.435



LLPLDKDY(ph)YVVR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 33501

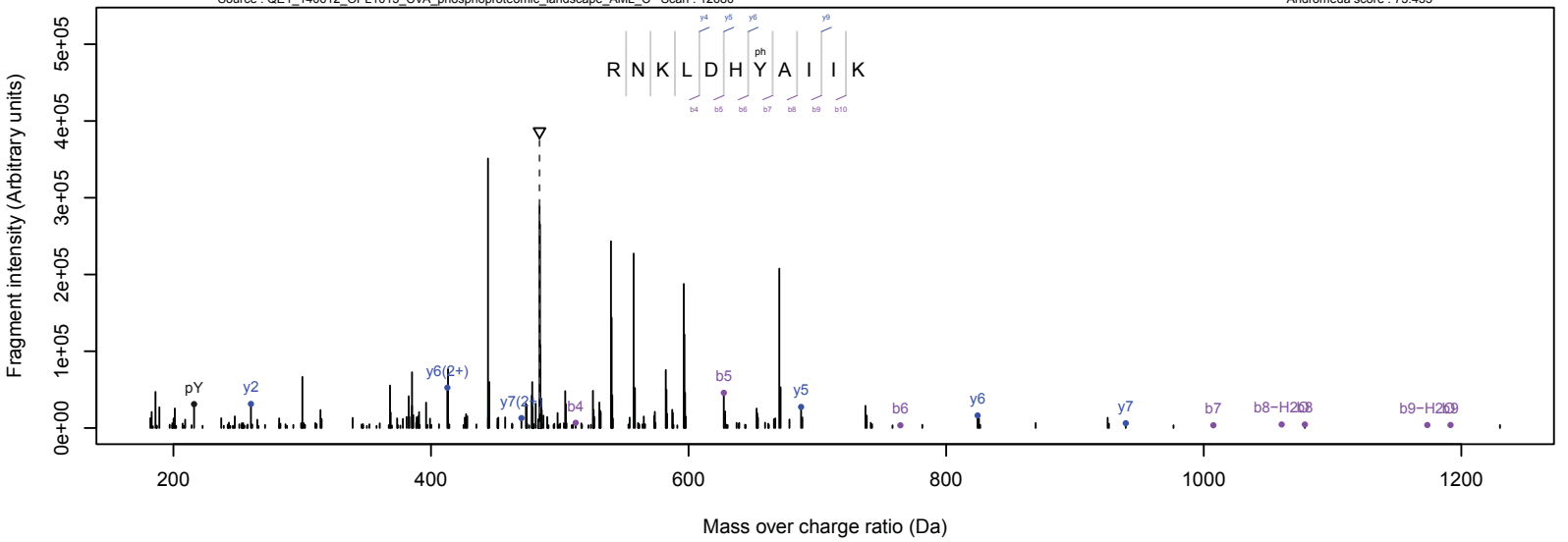
Andromeda score : 73.435



RNKLDHY(ph)AIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12680

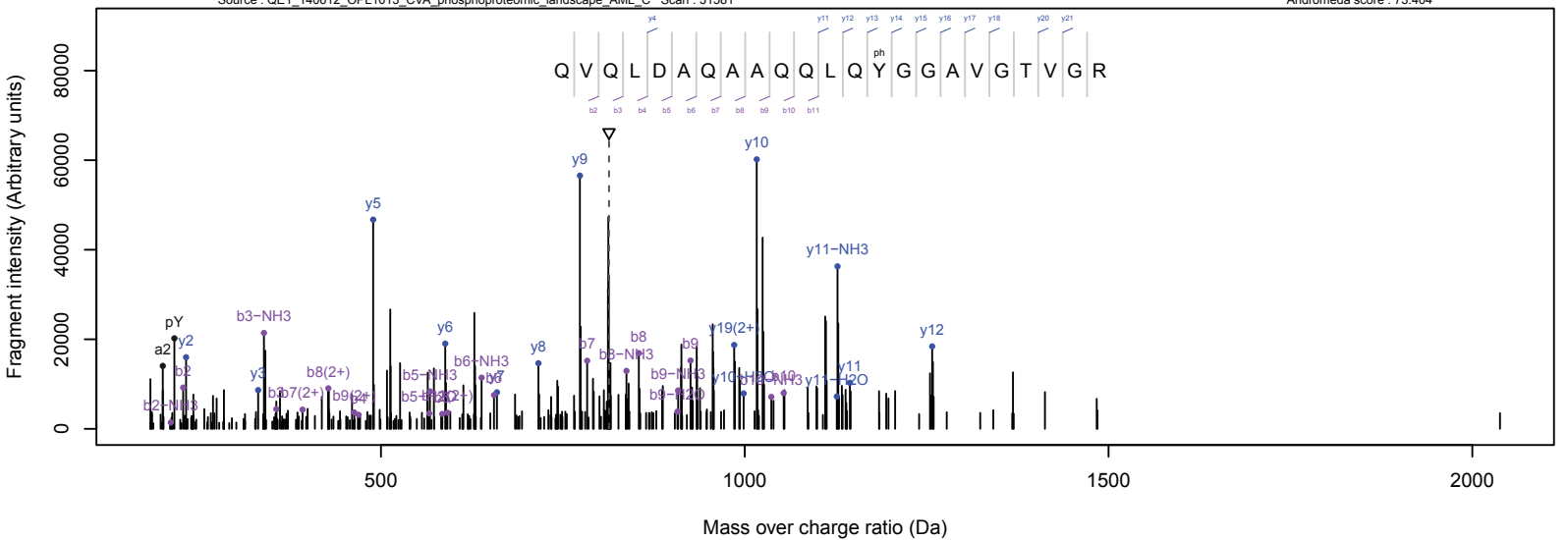
Andromeda score : 73.435



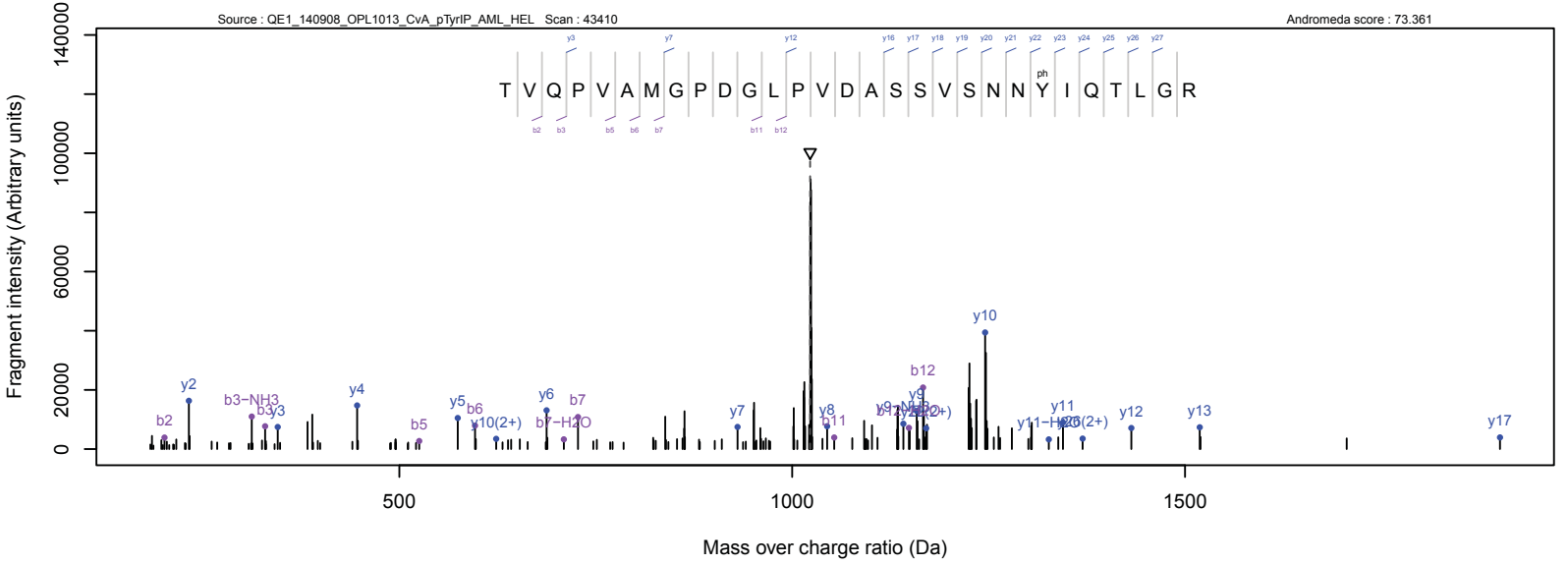
QVQLDAQAQQLQY(ph)GGAVGTVGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31581

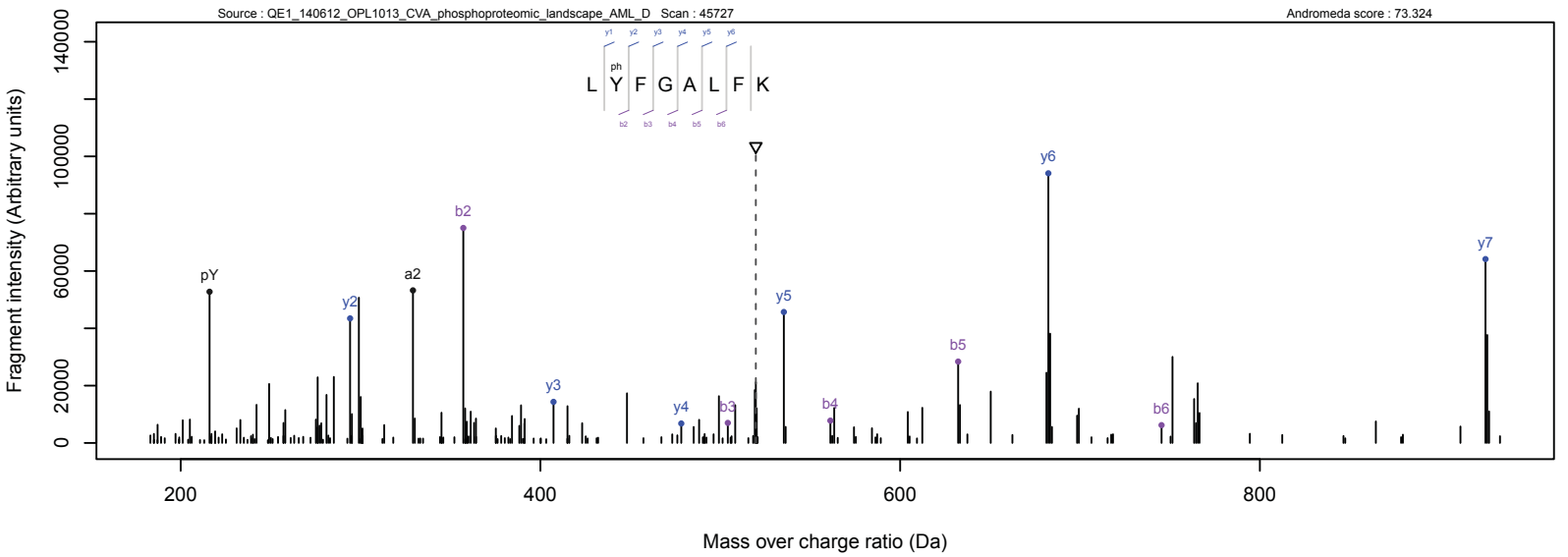
Andromeda score : 73.404



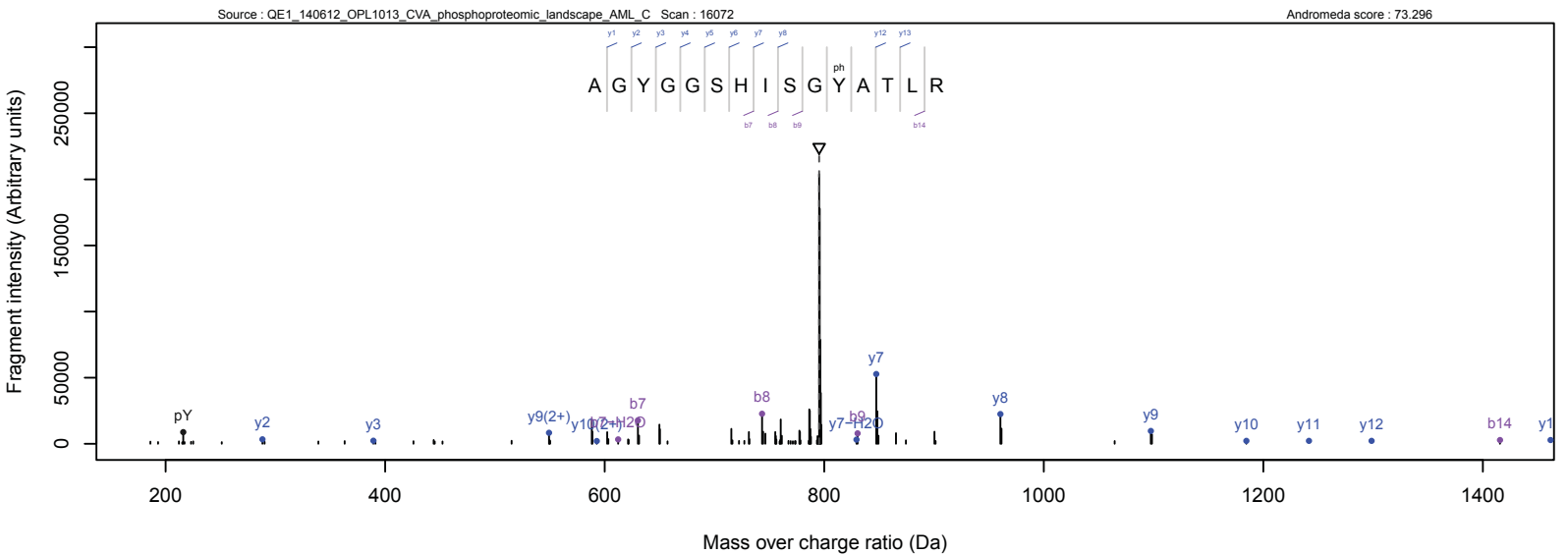
TVQPVAMGPDGLPVDASSVSNNY(ph)IQLGR-3+



LY(ph)FGALFK-2+



AGYGGSHISG(ph)ATLR-2+

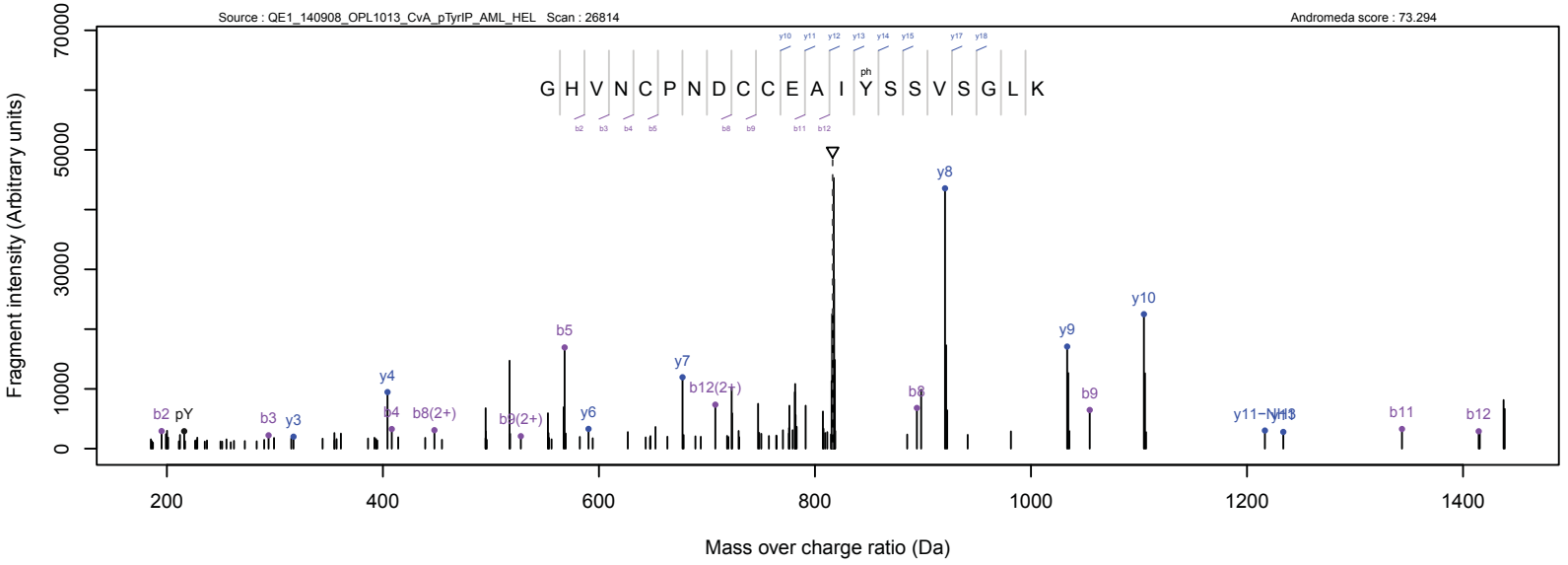




**GHVNCPCCEAIY(ph)SSVSLK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 26814

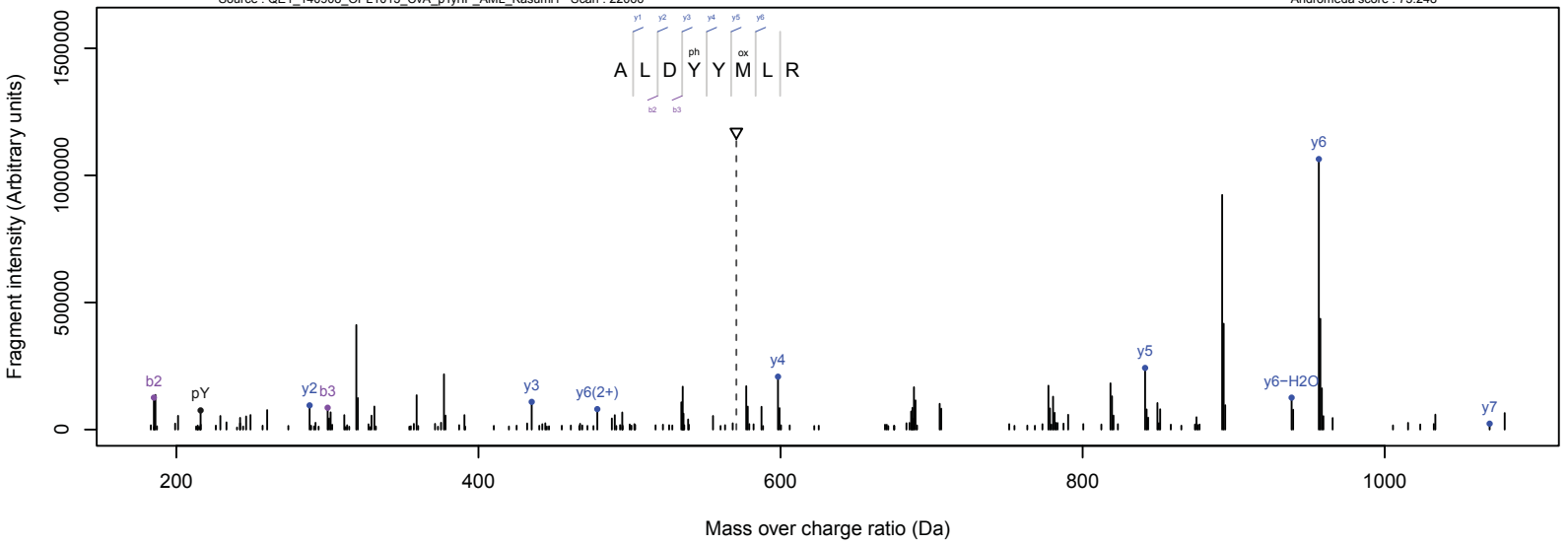
Andromeda score : 73.294



**ALDY(ph)YM(ox)LR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 22066

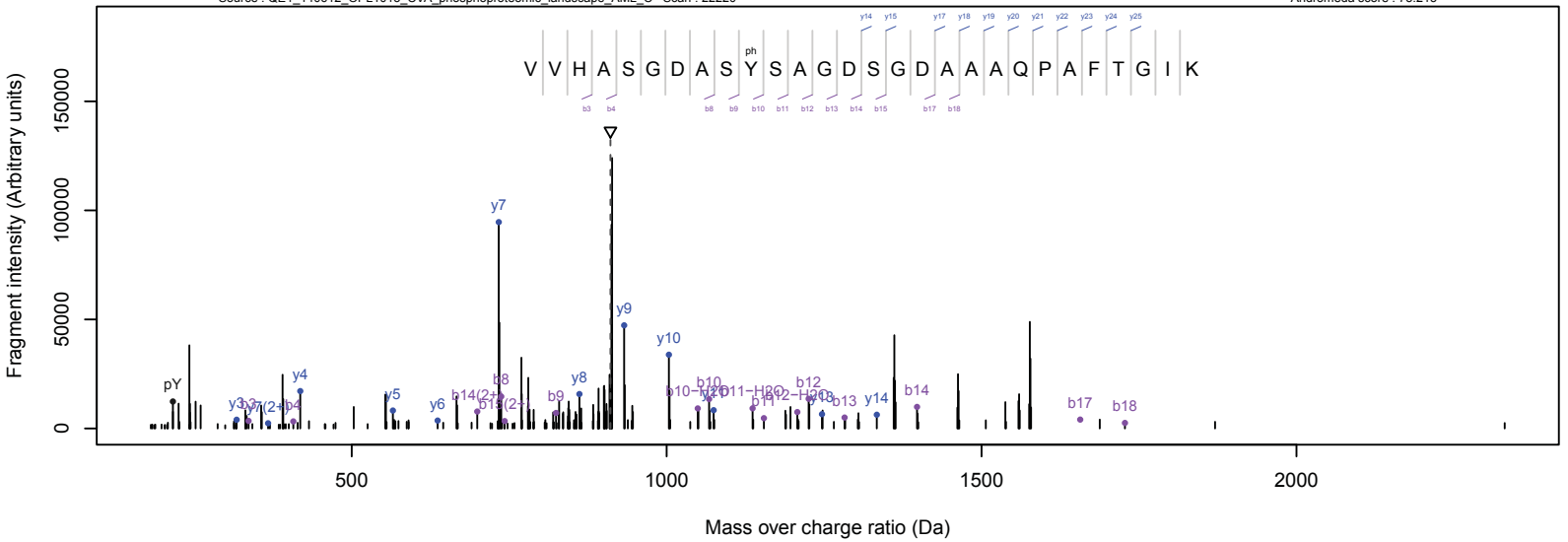
Andromeda score : 73.248



**VVHASGDASY(ph)SAGDSGDAQAQPAFTGIK-3+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 22220

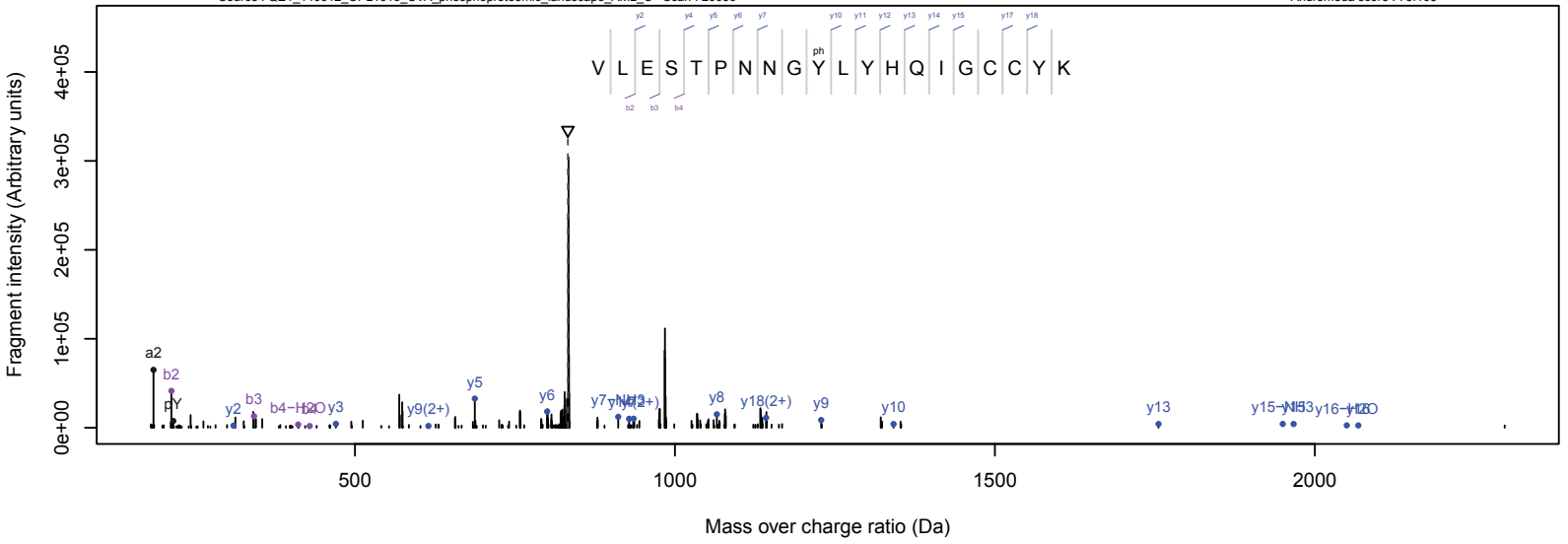
Andromeda score : 73.213



VLESTPNNGY(ph)LYHQIGCCYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26636

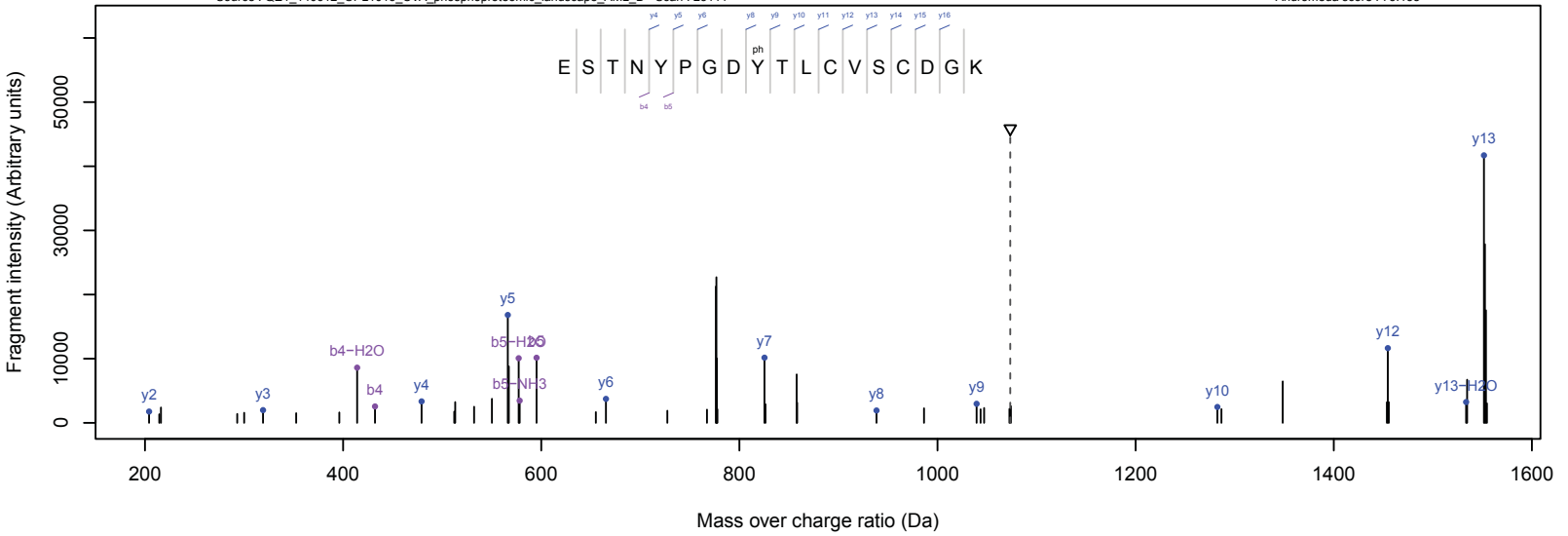
Andromeda score : 73.168



ESTNYPGDY(ph)TLCVSCDGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23111

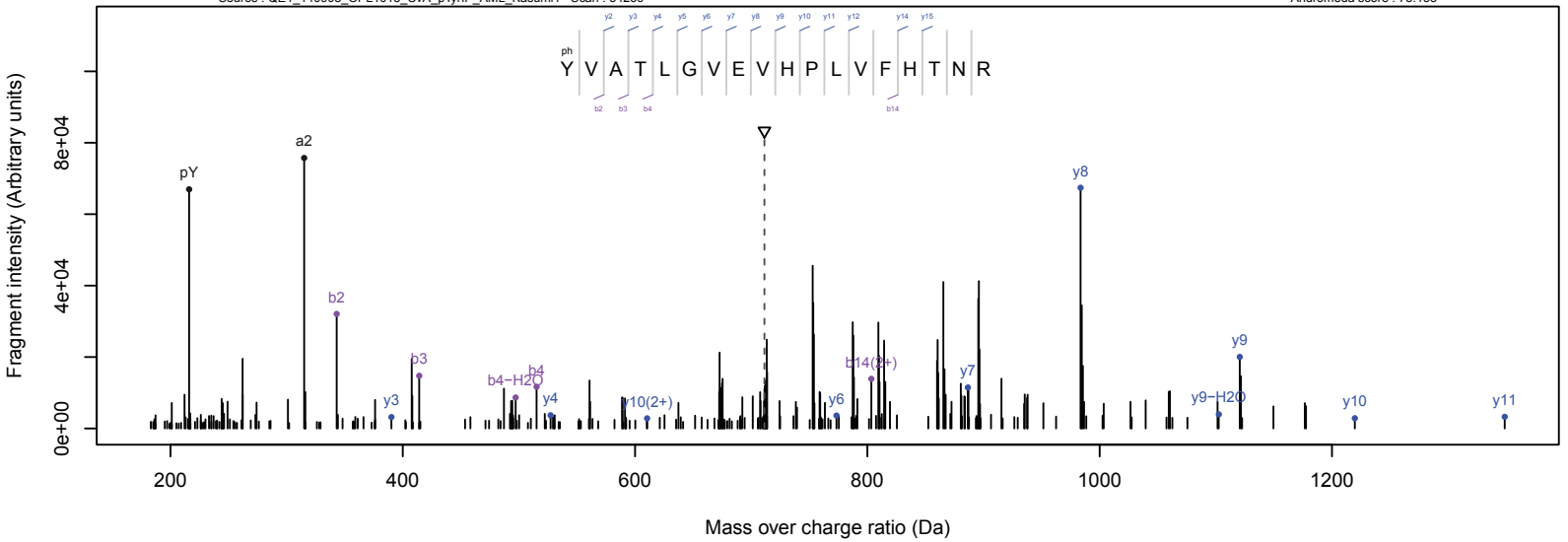
Andromeda score : 73.138



Y(ph)VATLGVEVHPLVFHTNR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 34239

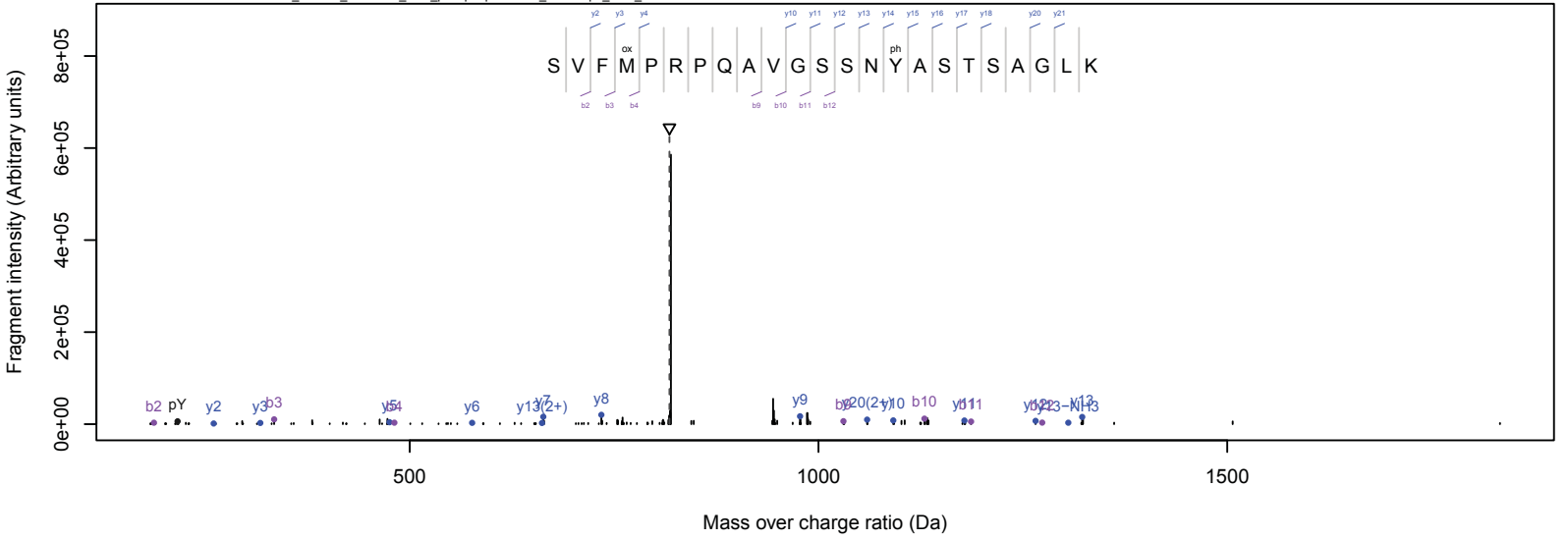
Andromeda score : 73.138



SVFM(ox)PRPQAVGSSNY(ph)ASTSAGLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26178

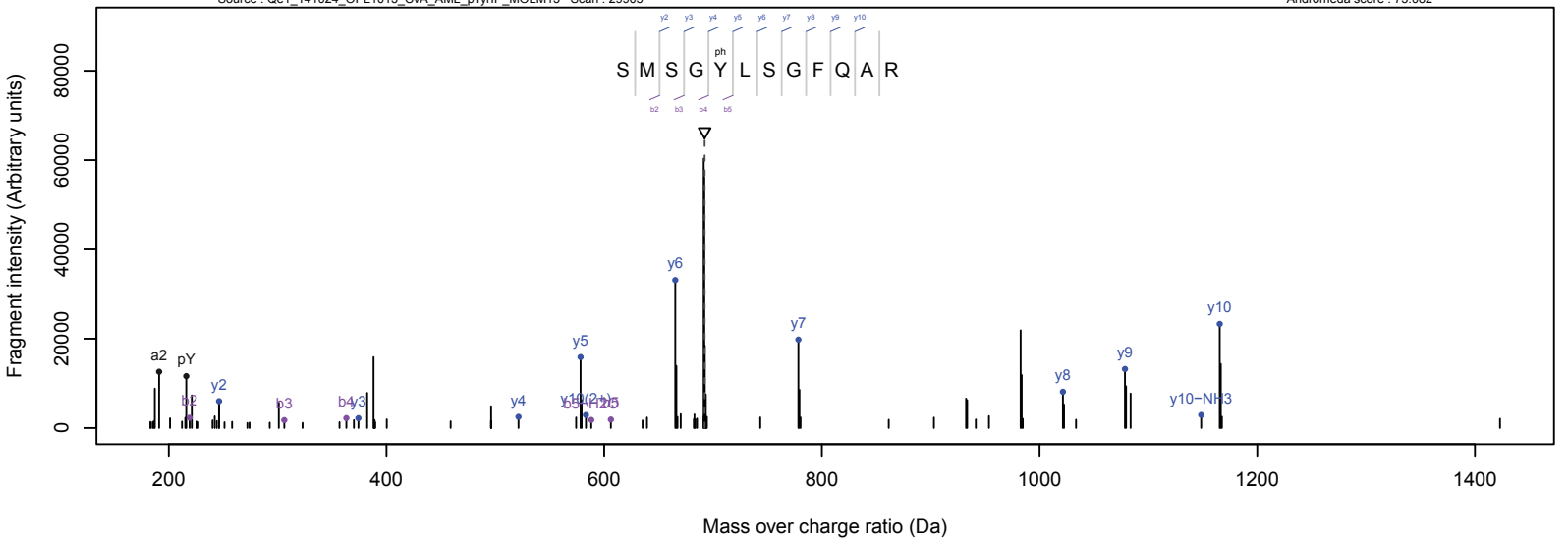
Andromeda score : 73.11



SMSGY(ph)LSGFQAR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 29905

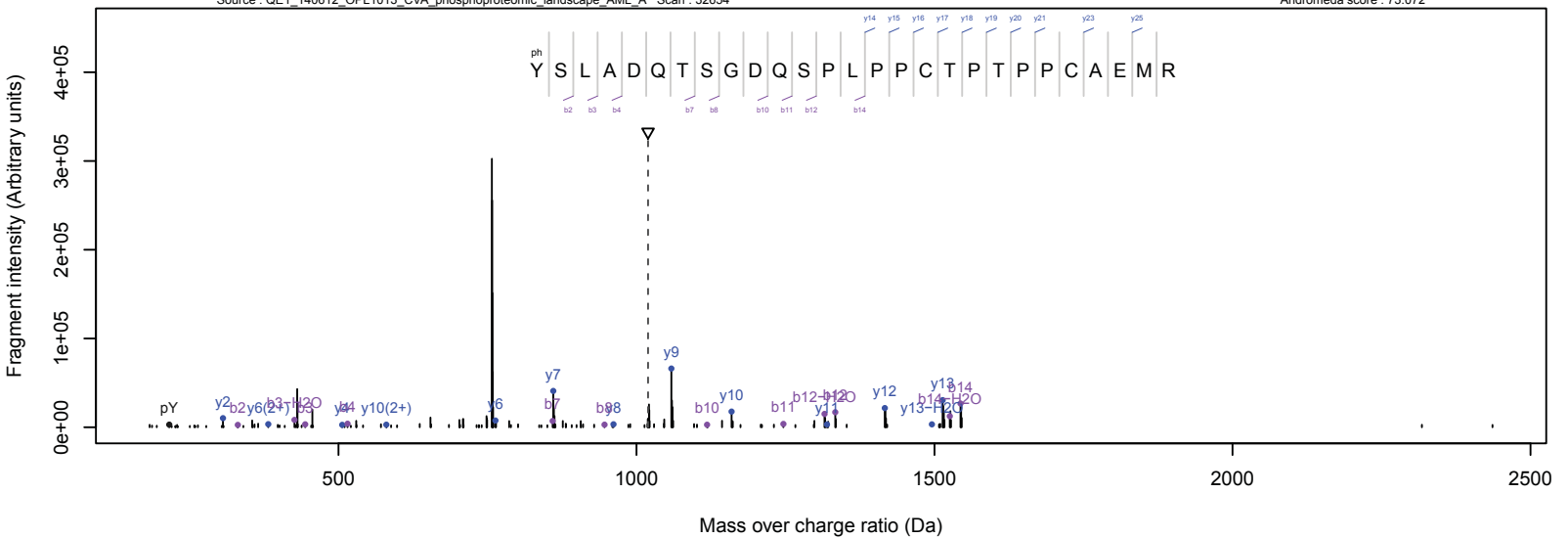
Andromeda score : 73.082



Y(ph)SLADQTS G D Q S P L P P C T P T P P C A E M R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32654

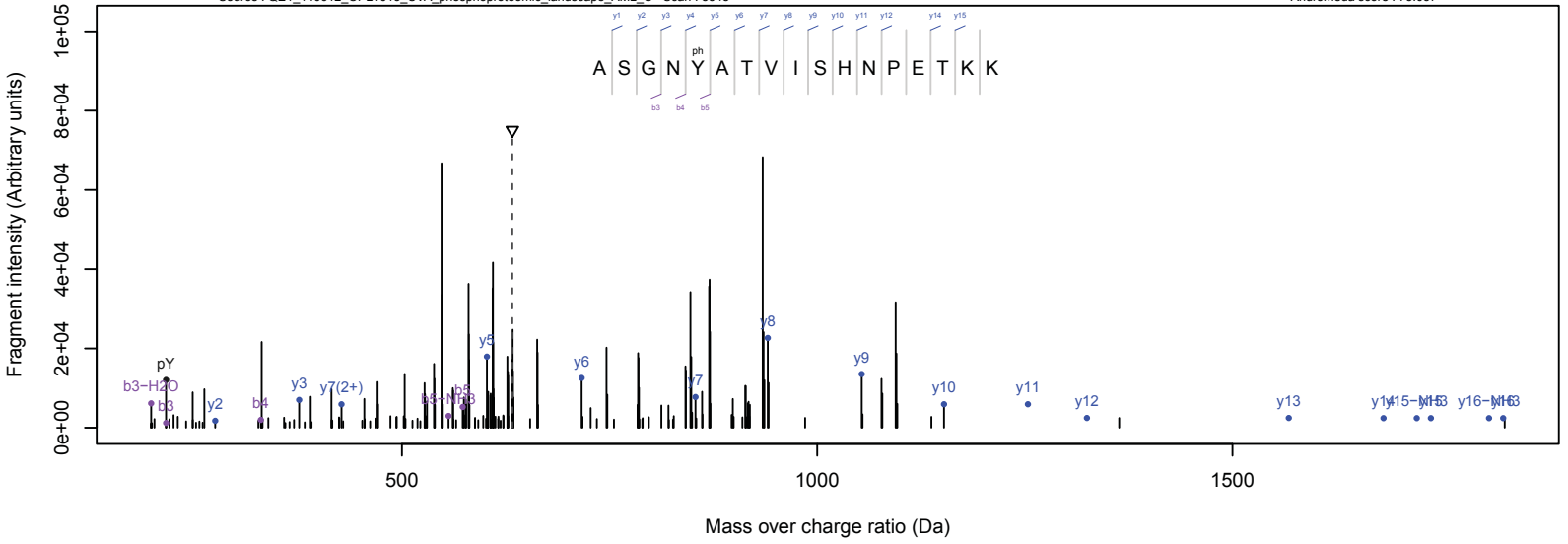
Andromeda score : 73.072



ASGNY(ph)ATVISHNPETKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8548

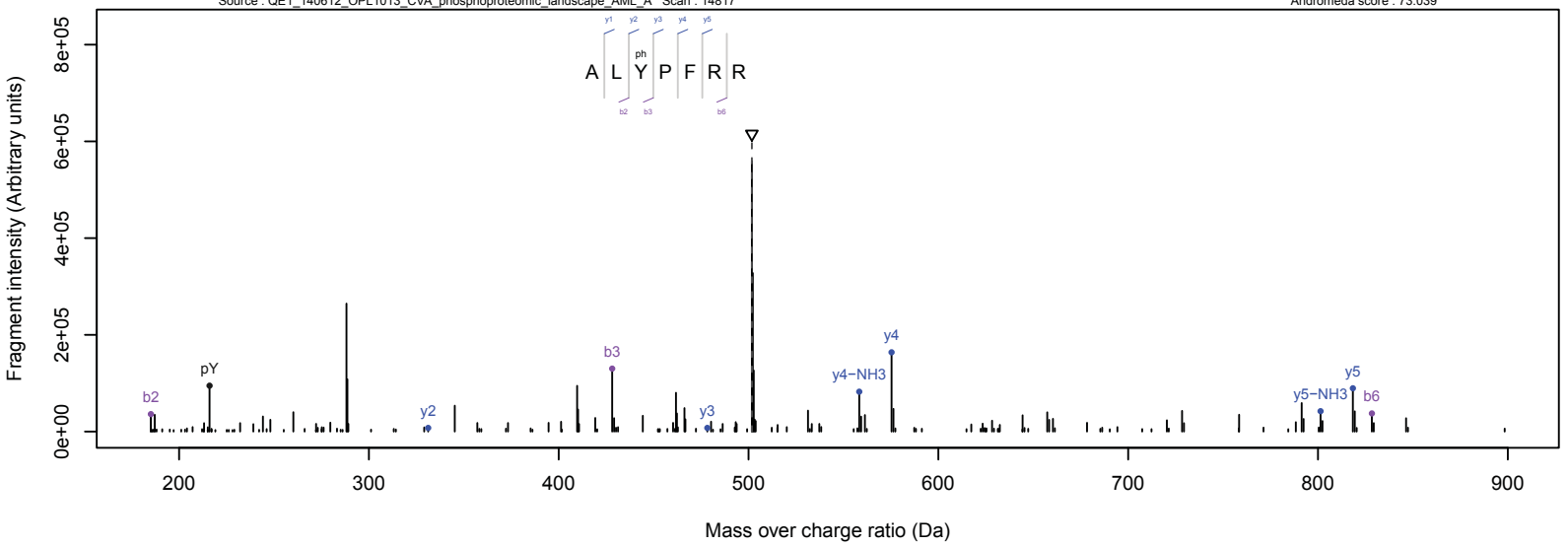
Andromeda score : 73.057



ALY(ph)PFRR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 14817

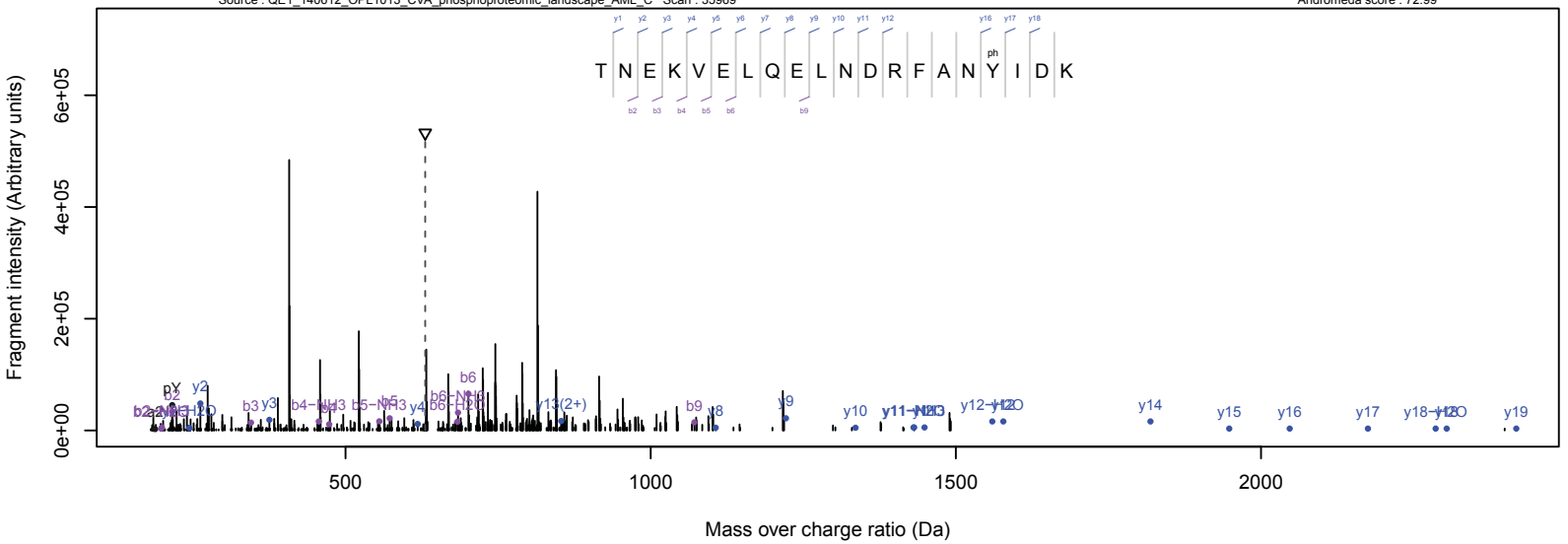
Andromeda score : 73.039



TNEKVELQELNDRFANY(ph)IDK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35969

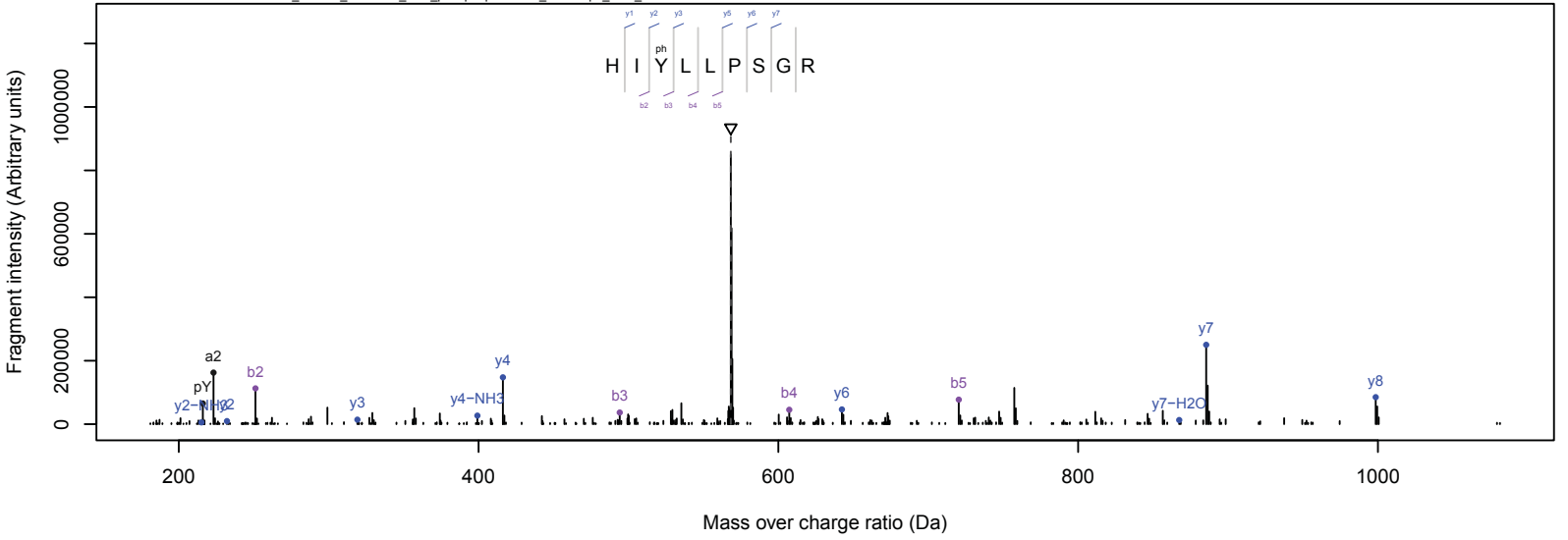
Andromeda score : 72.99



HIY(ph)LLPSGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23805

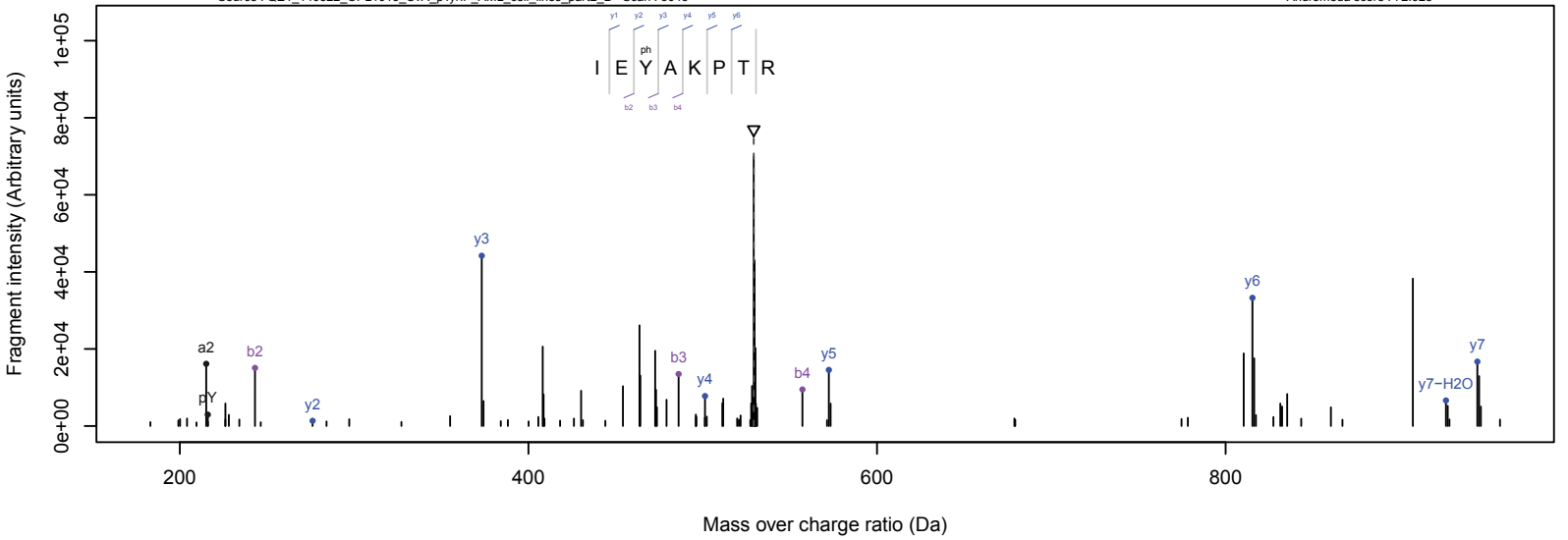
Andromeda score : 72.928



IEY(ph)AKPTR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3045

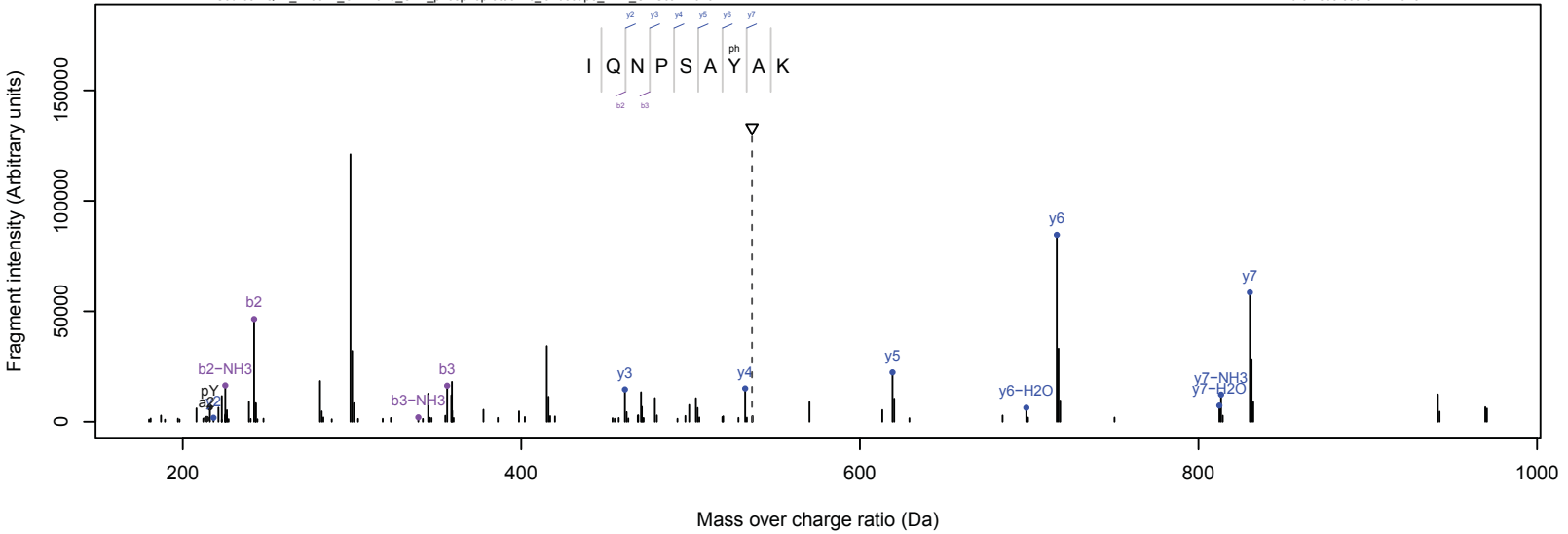
Andromeda score : 72.928



IQNPSAY(ph)AK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6231

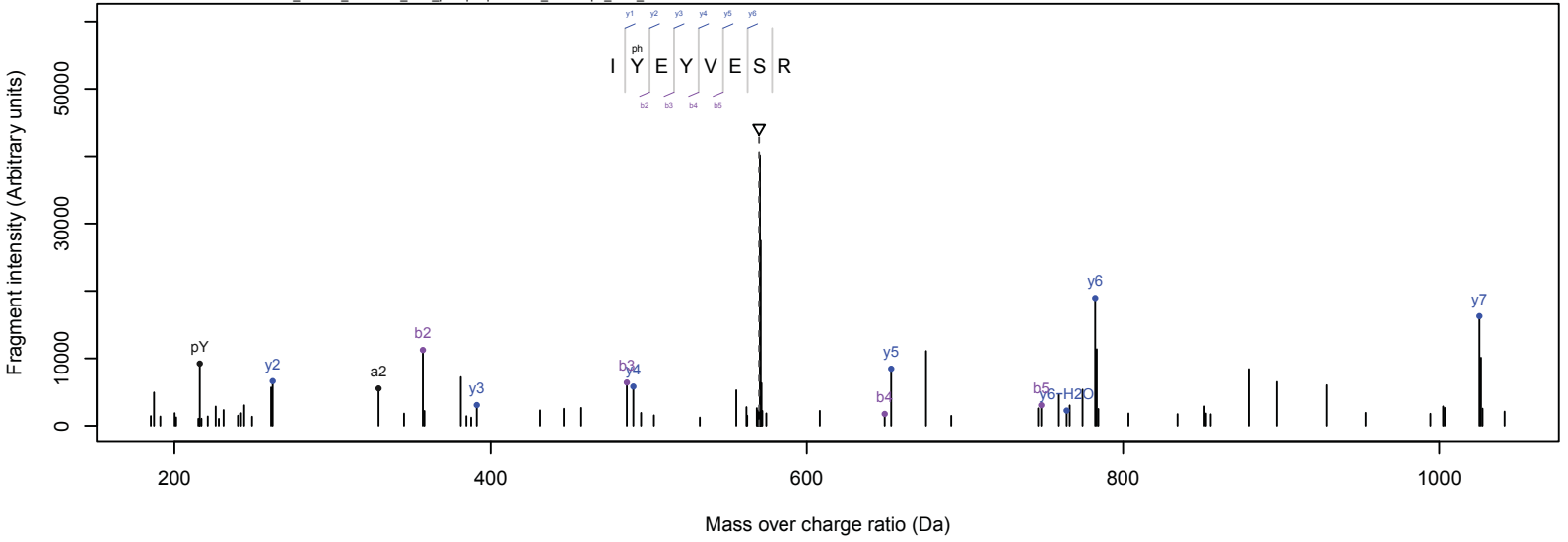
Andromeda score : 72.928



IY(ph)EYVESR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 11120

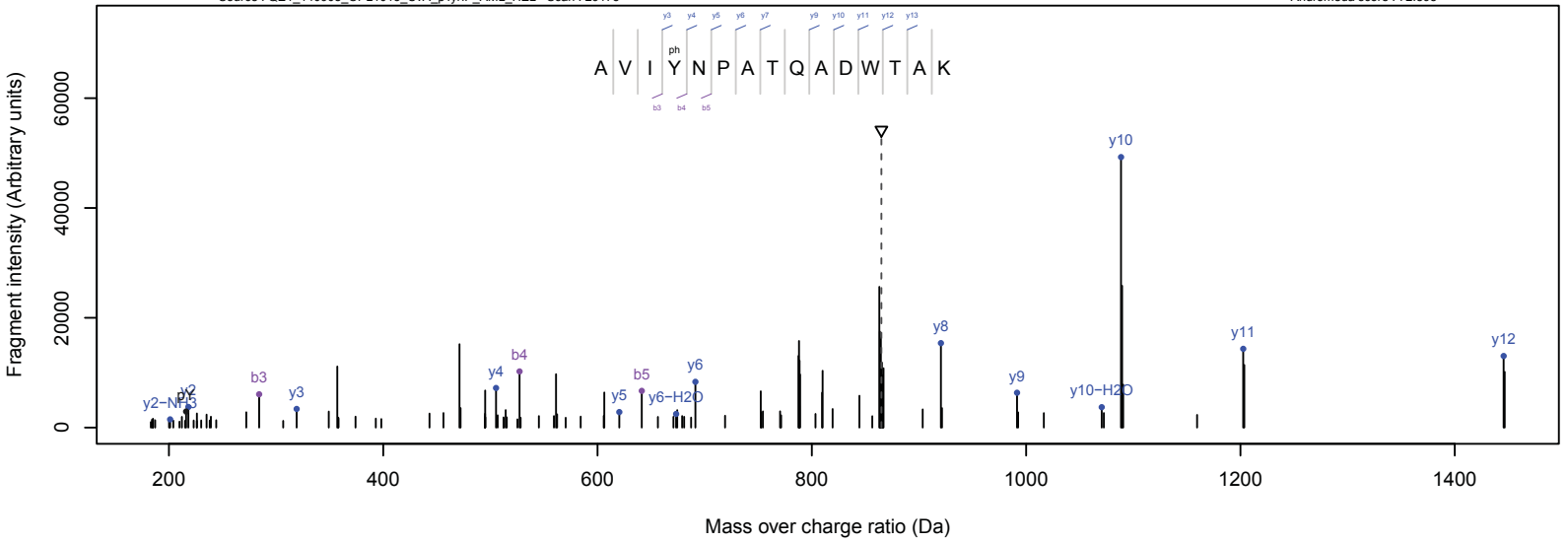
Andromeda score: 72.928



AVIY(ph)NPATQADWTAK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 29175

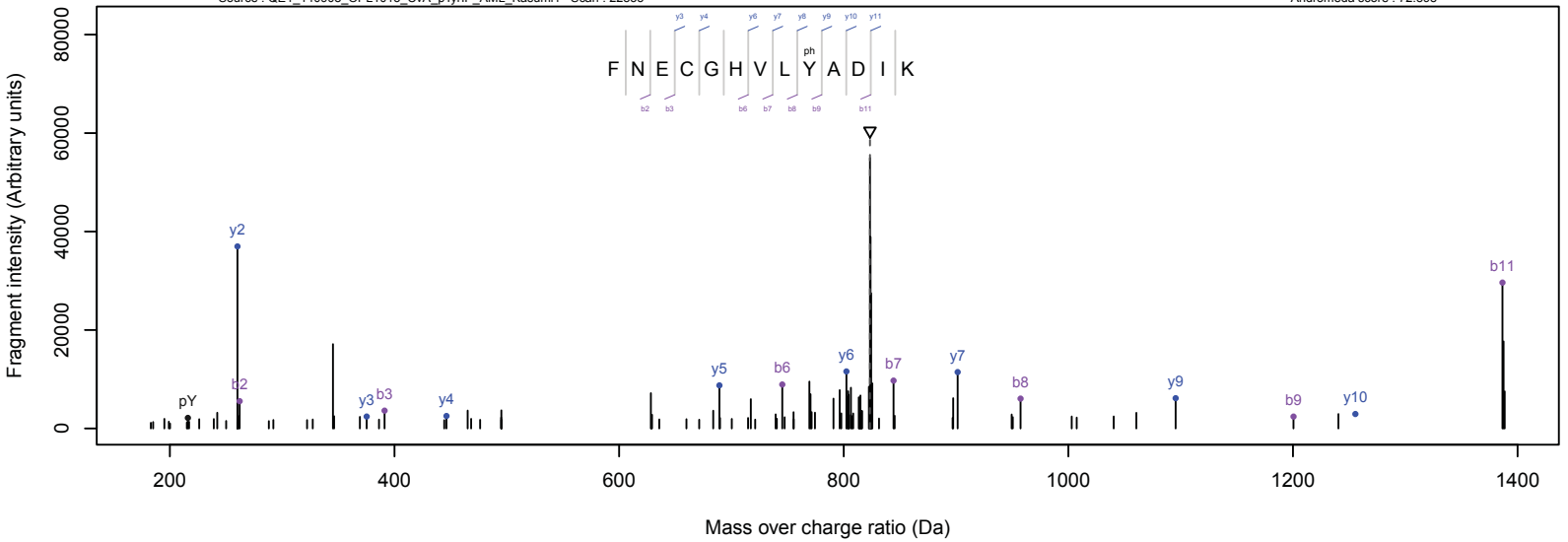
Andromeda score: 72.898



FNECGHVLY(ph)ADIK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 22835

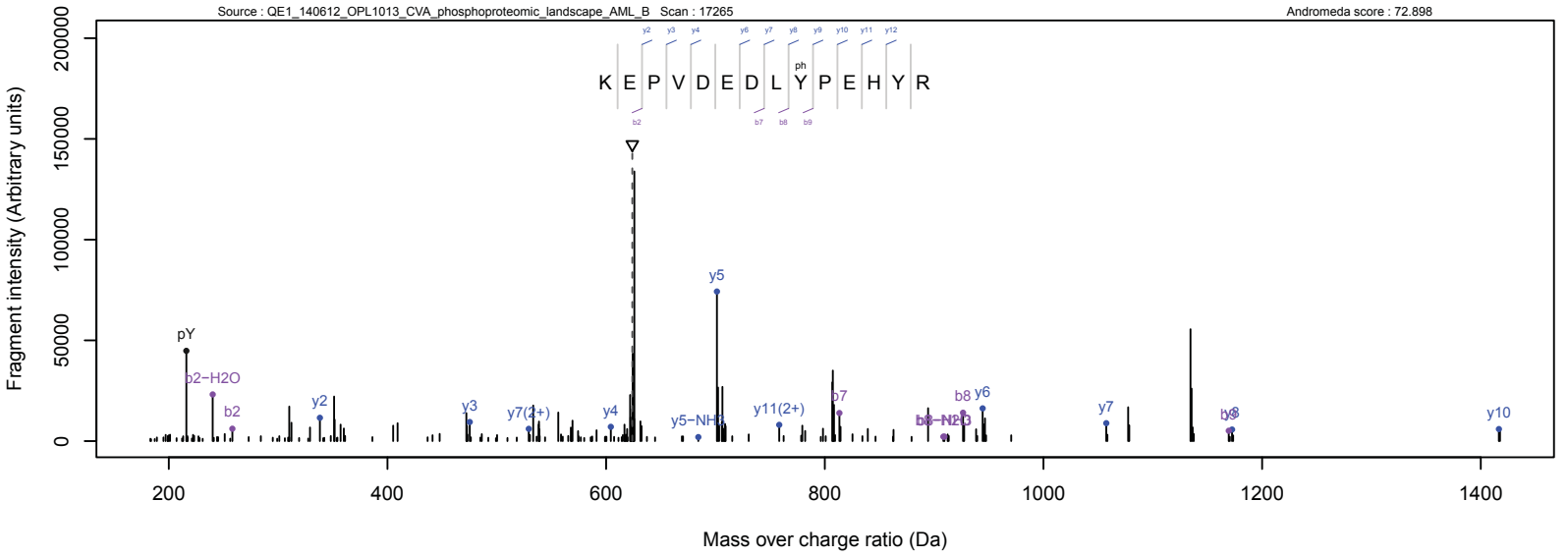
Andromeda score: 72.898



**KEPVDEDLY(ph)PEHYR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17285

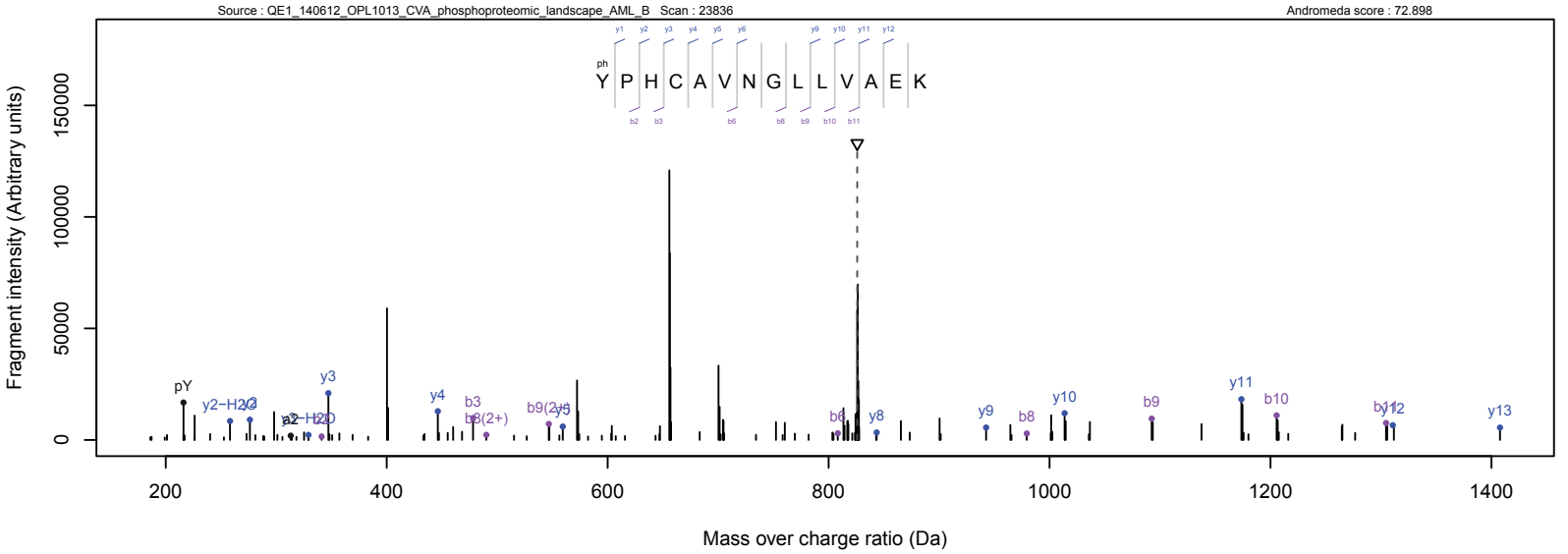
Andromeda score : 72.898



**Y(ph)PHCAVNGLLVAEK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23836

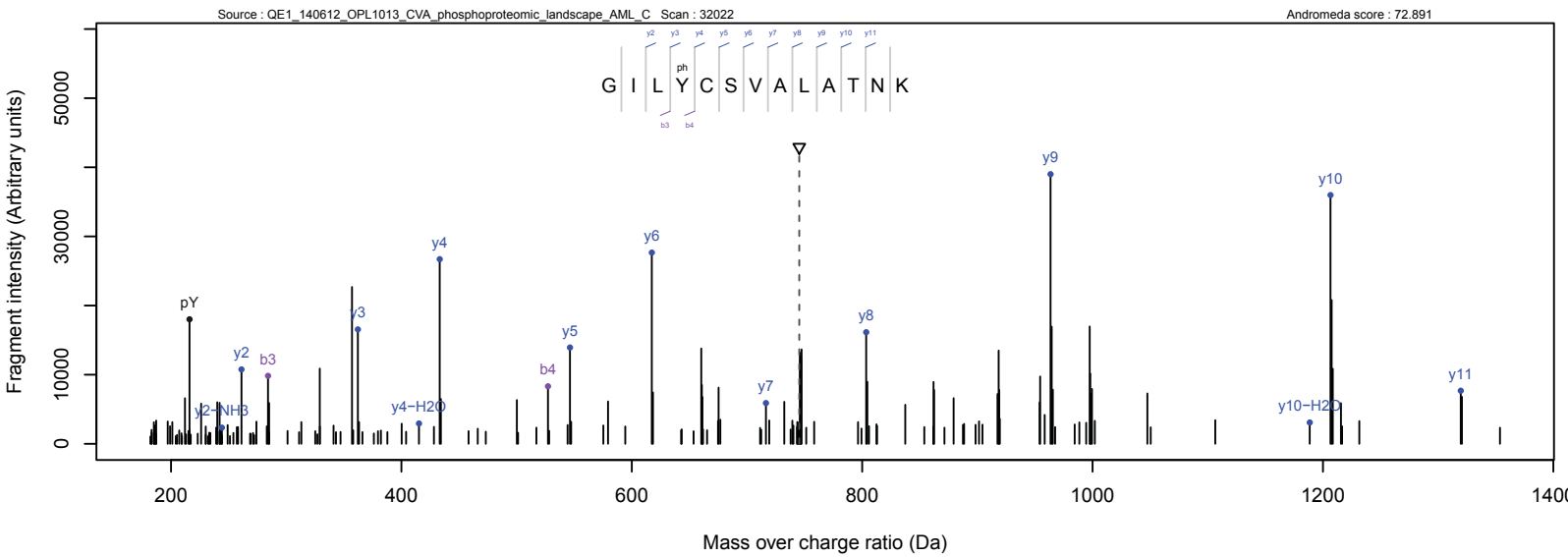
Andromeda score : 72.898



**GILY(ph)CSVALATNK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32022

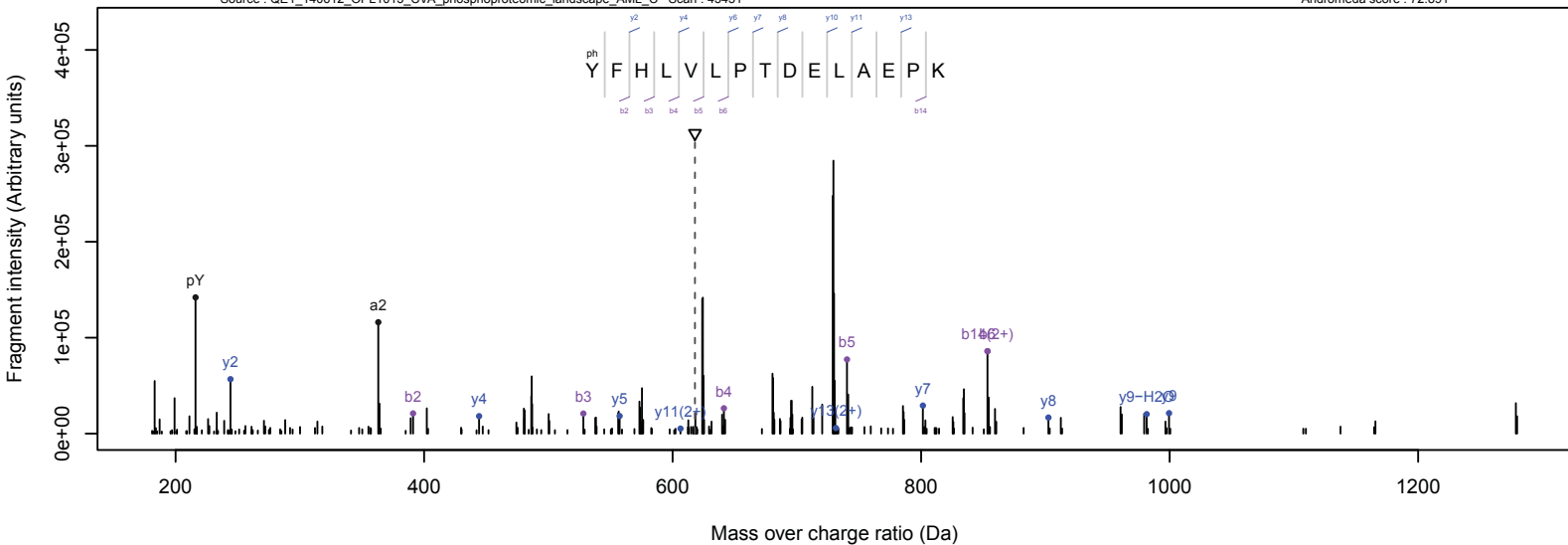
Andromeda score : 72.891



Y(ph)FHLVLPDELAEPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43451

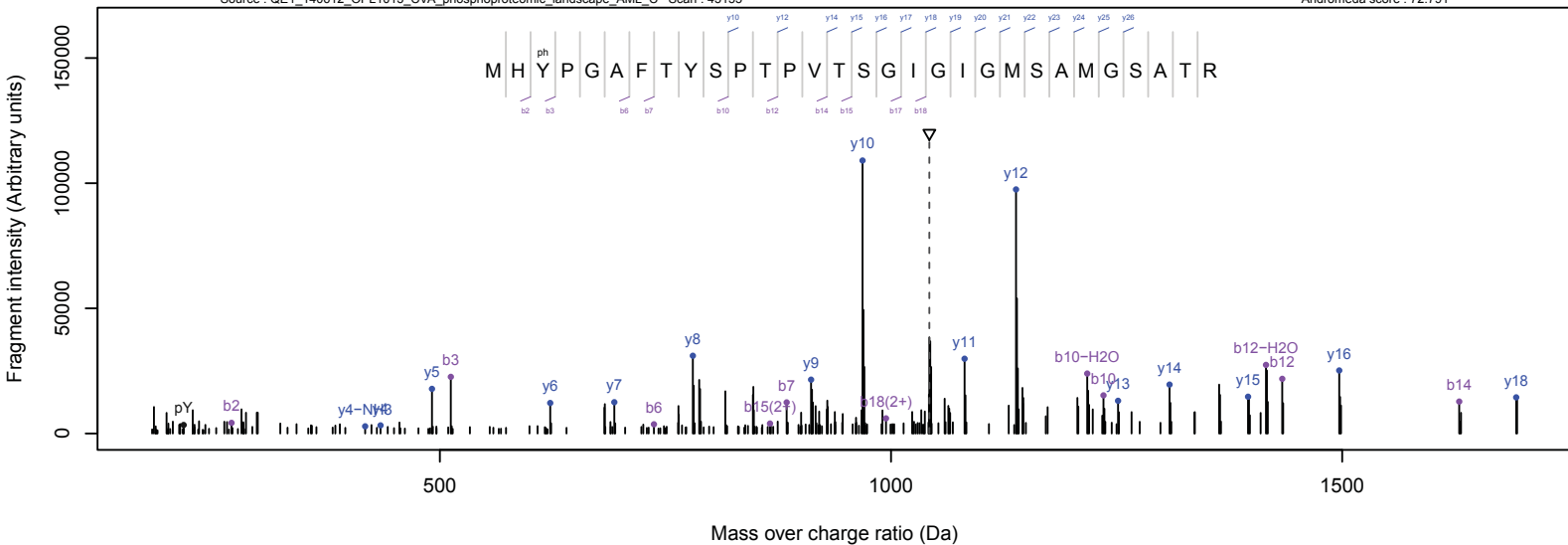
Andromeda score : 72.891



MHY(ph)PGAFTYSPTPTVTSIGIGIGMSAMGSATR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45155

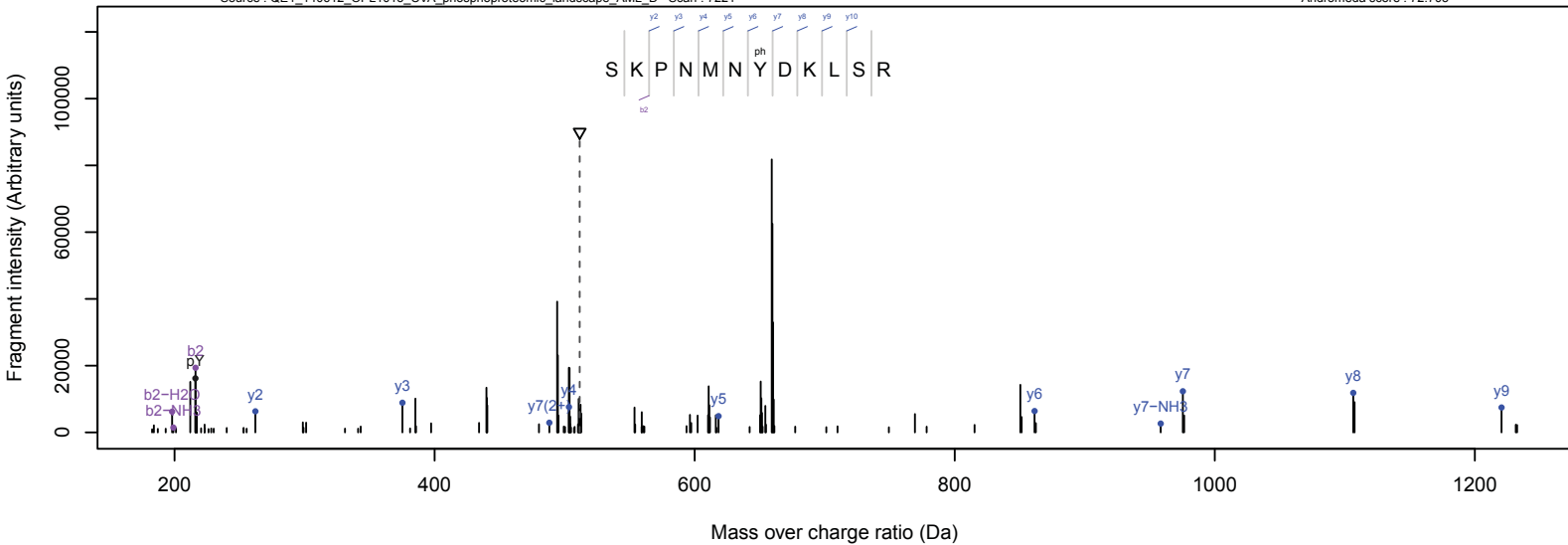
Andromeda score : 72.791



SKPNMNY(ph)DKLSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 7221

Andromeda score : 72.705

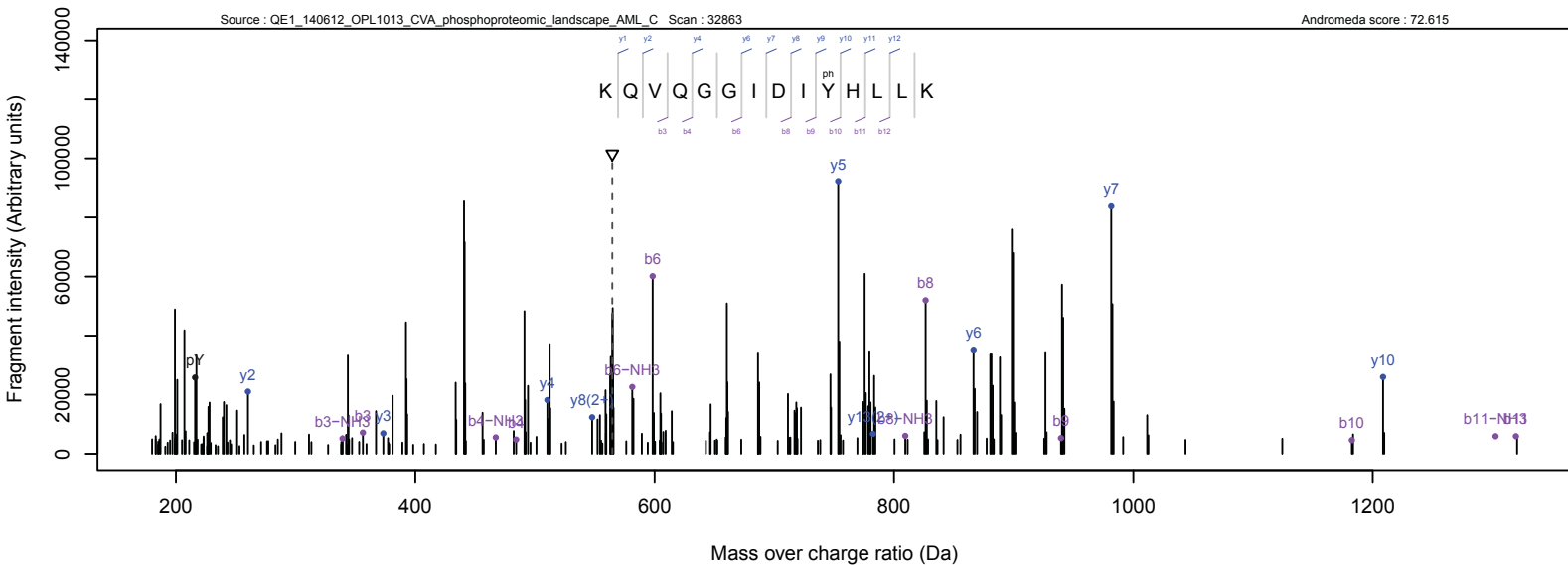




**KVQGGIDIYHLLK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32863

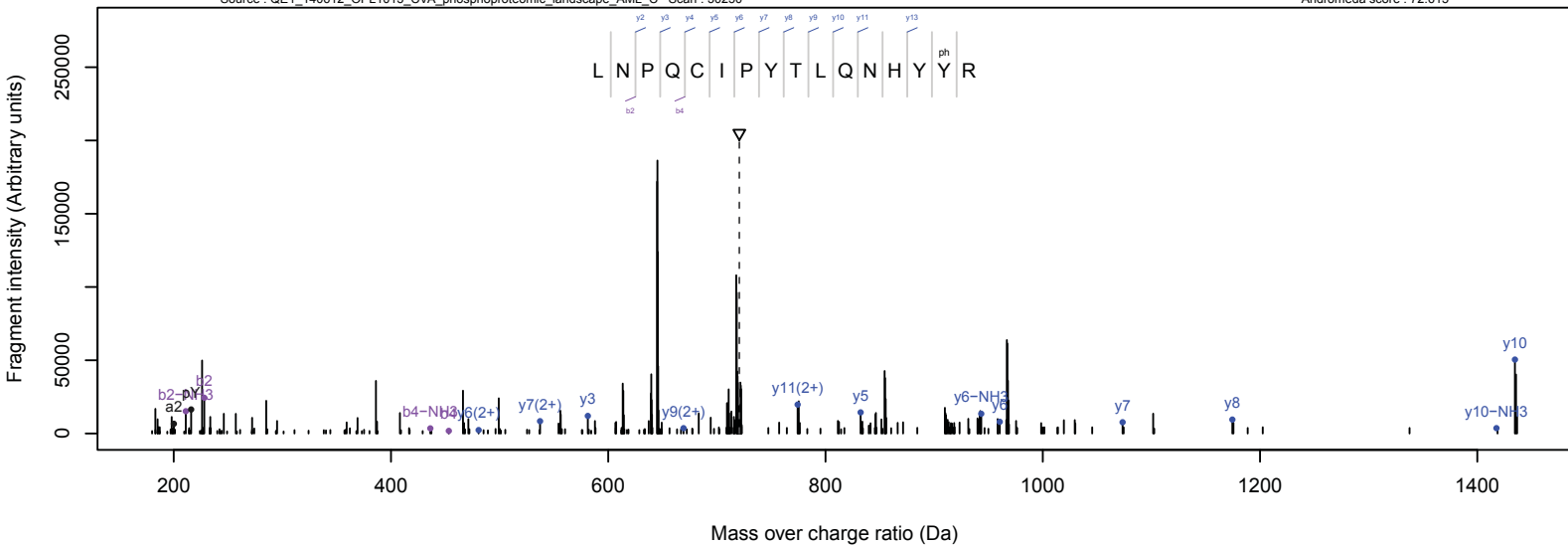
Andromeda score : 72.615



**LNPQCIPYTLQNHYYR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30250

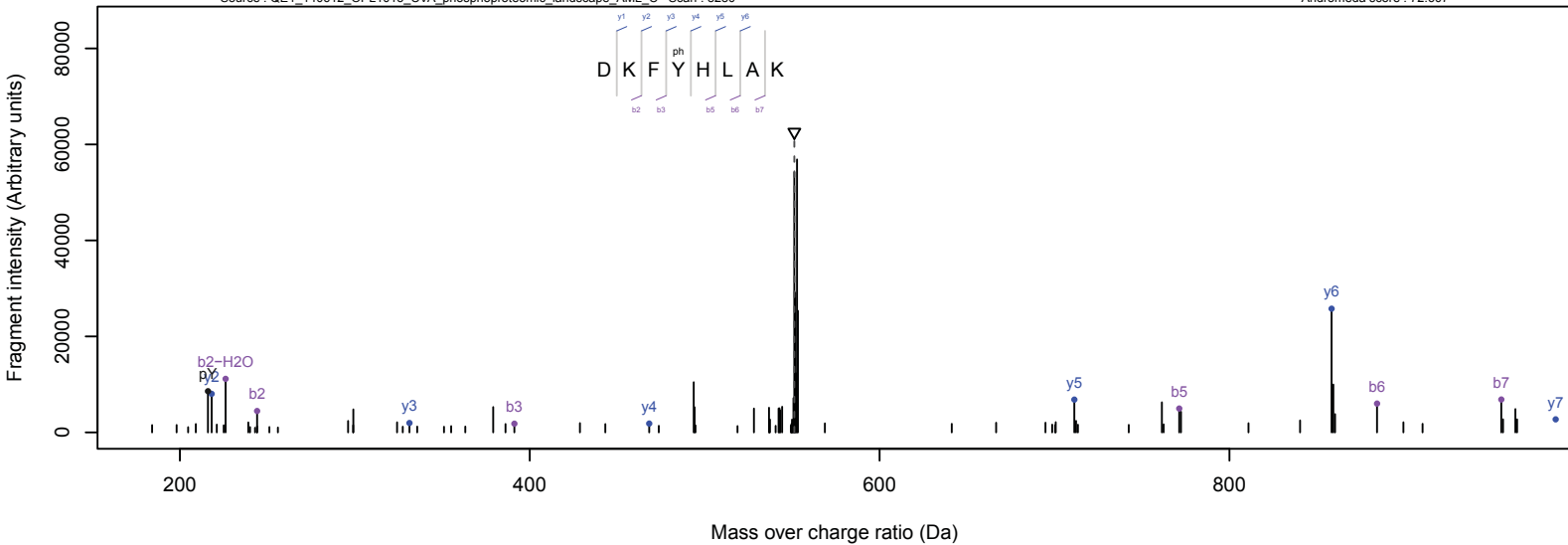
Andromeda score : 72.615



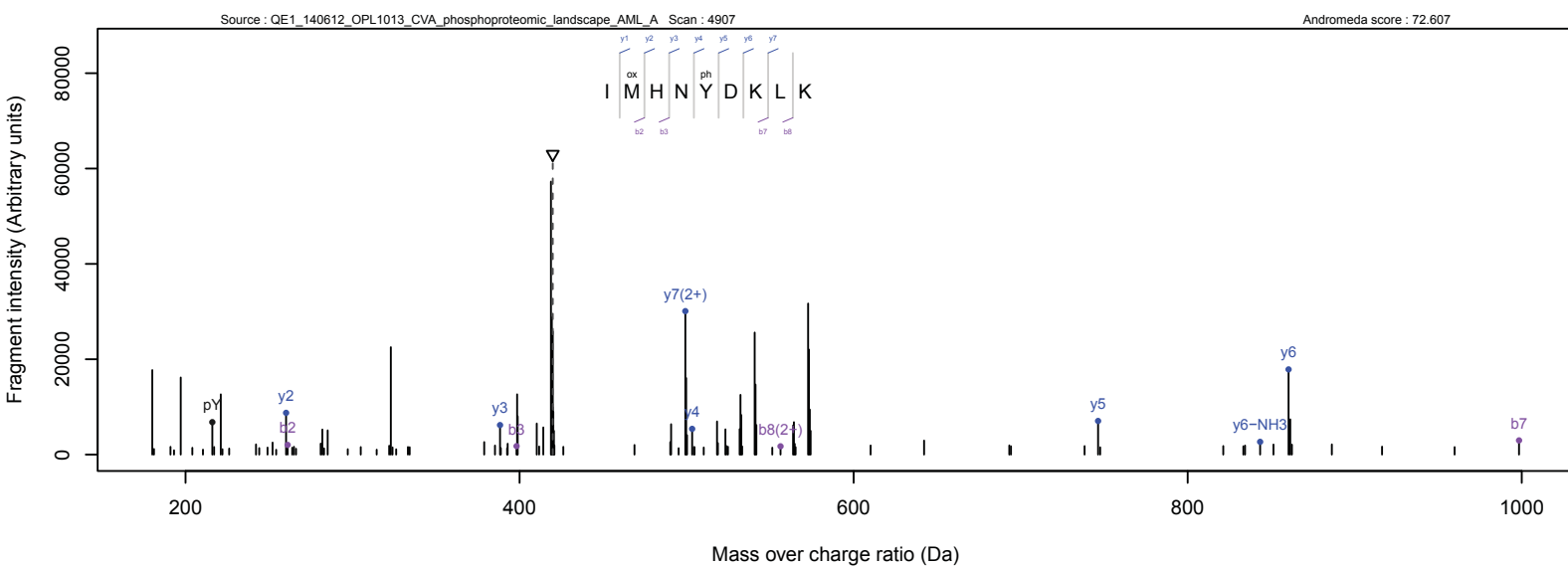
**DKFYHLLAK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8280

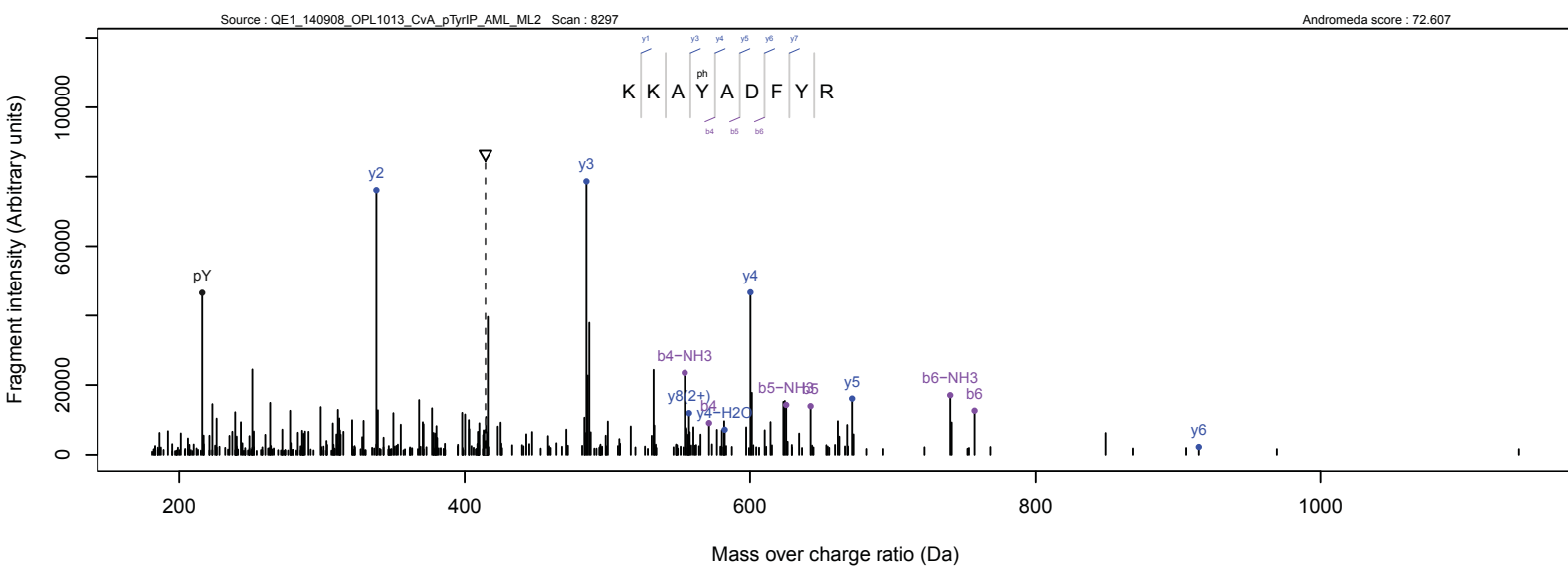
Andromeda score : 72.607



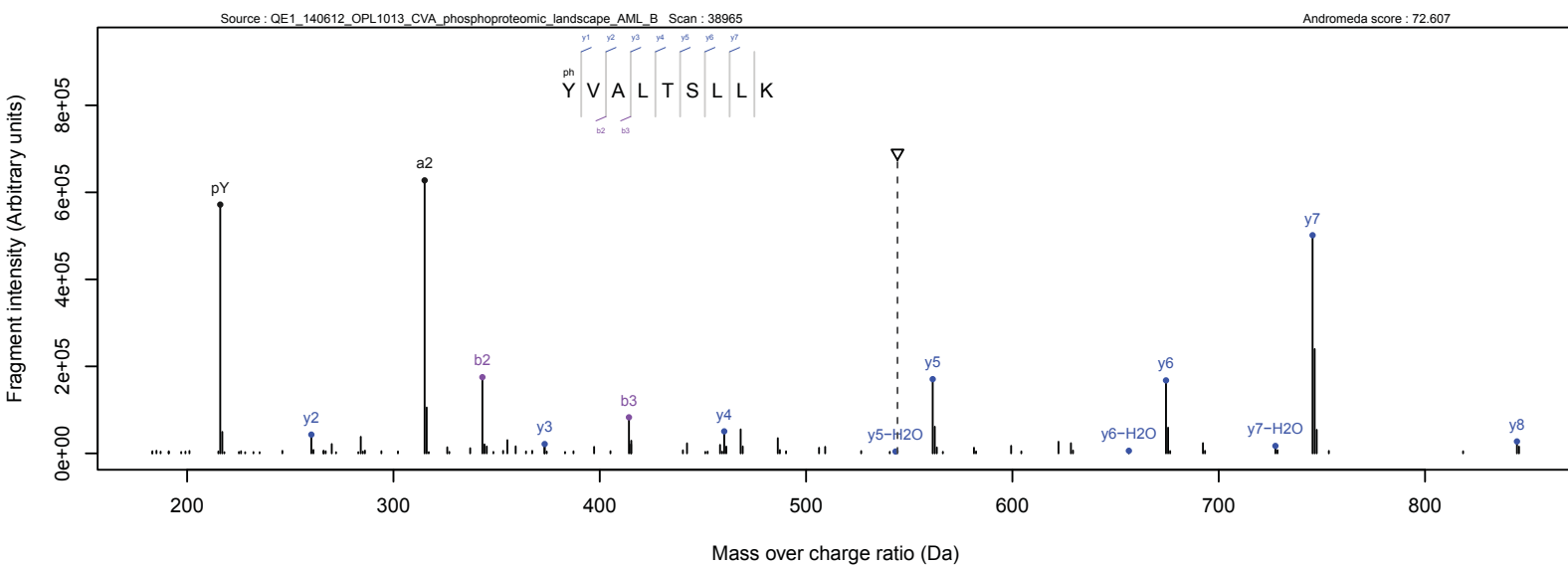
## IM(ox)HNY(ph)DKLK-3+



## KKAY(ph)ADFYR-3+

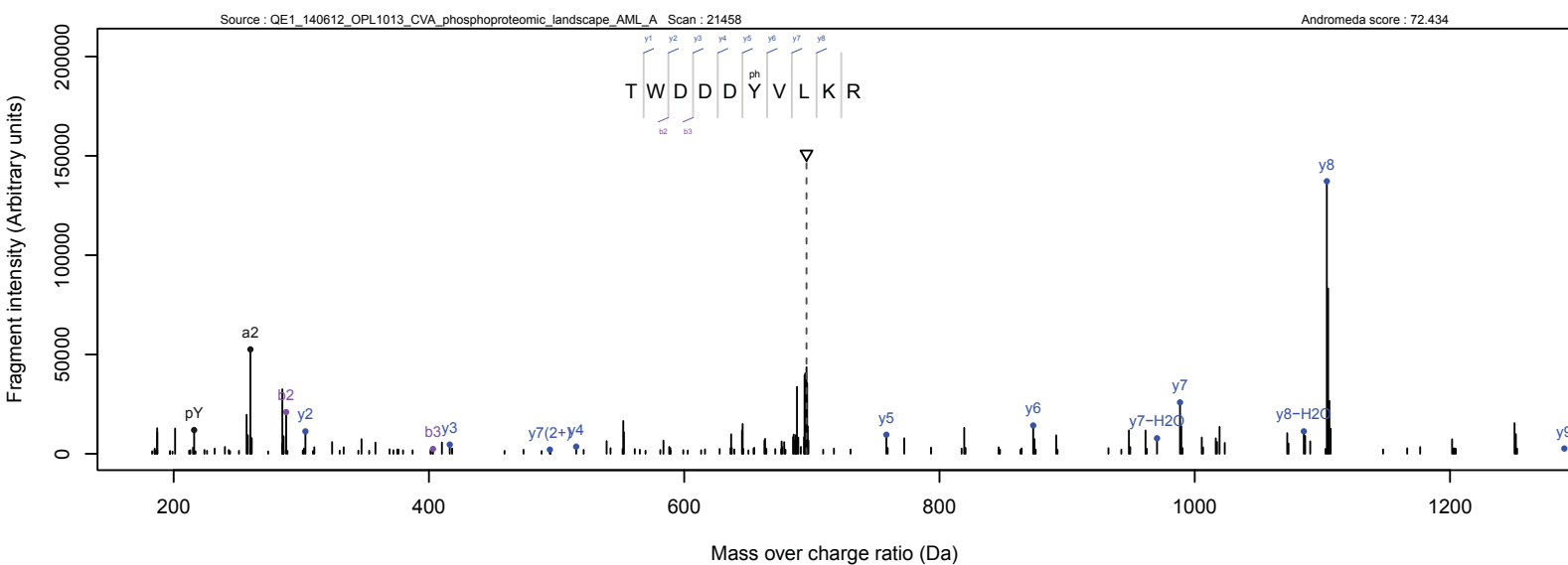


## Y(ph)VALTSLLK-2+

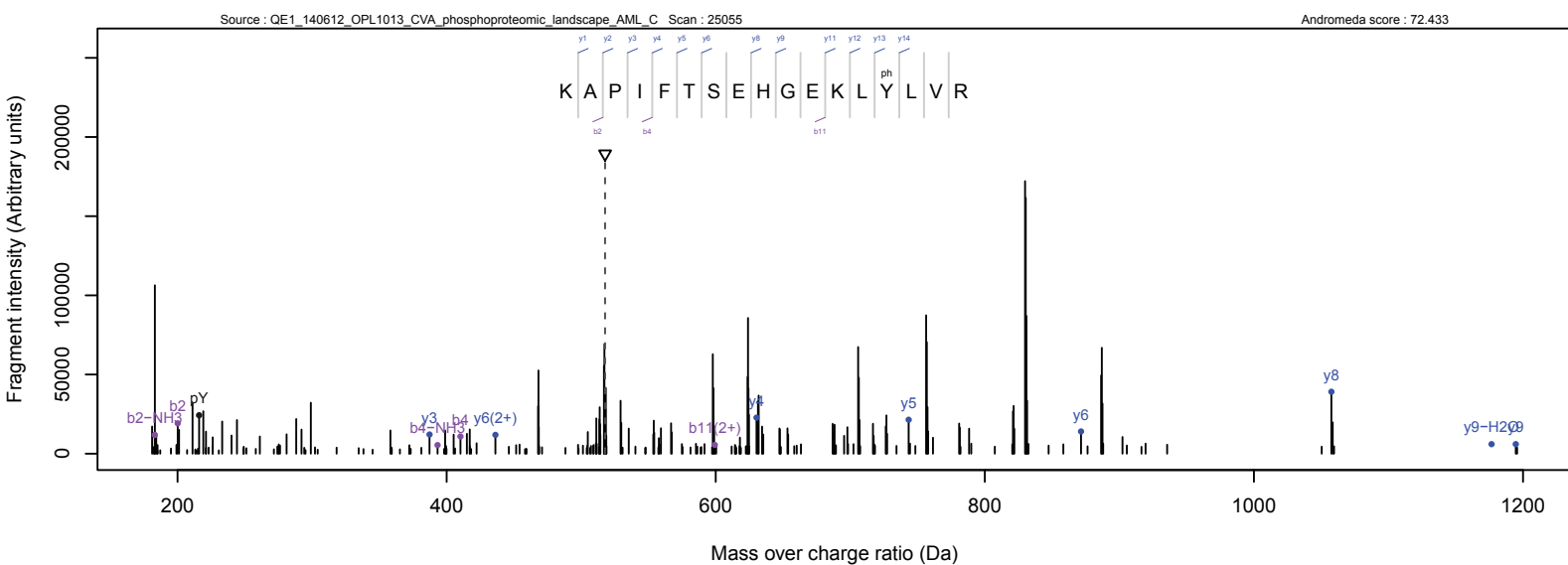




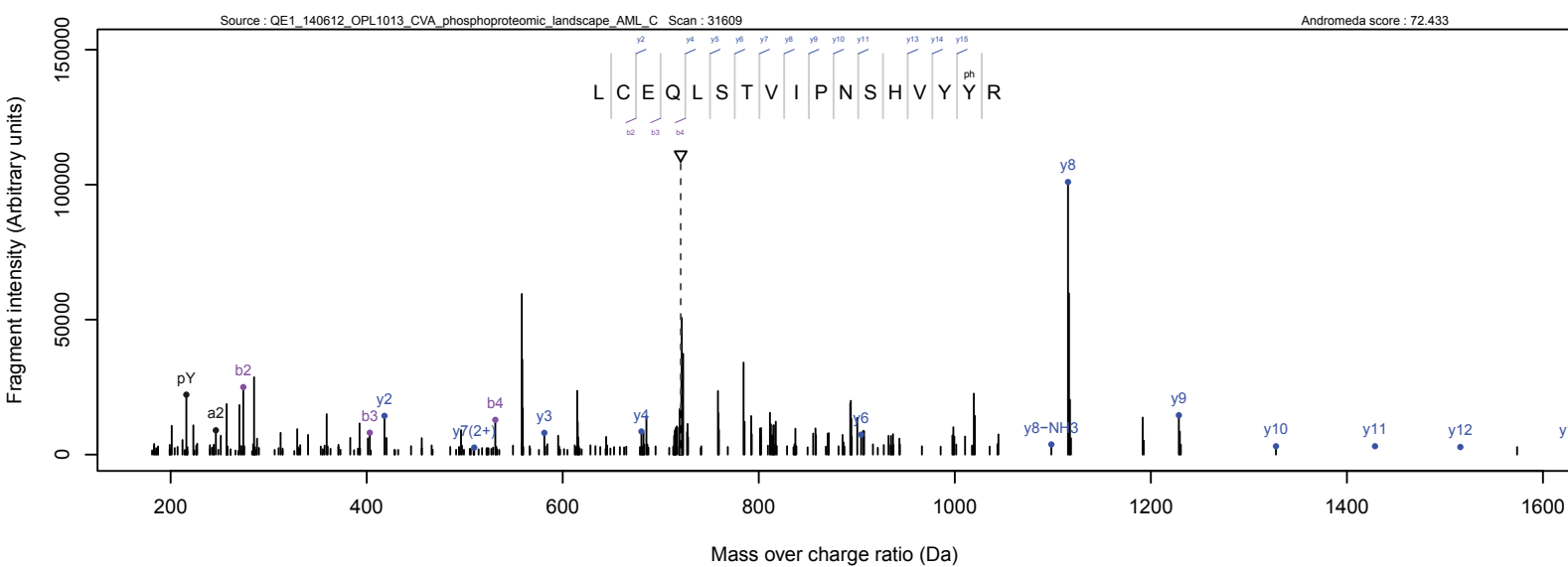
## TWDDDY(ph)V LKR-2+



## KAPIFTSEHGKLY LVR-4+



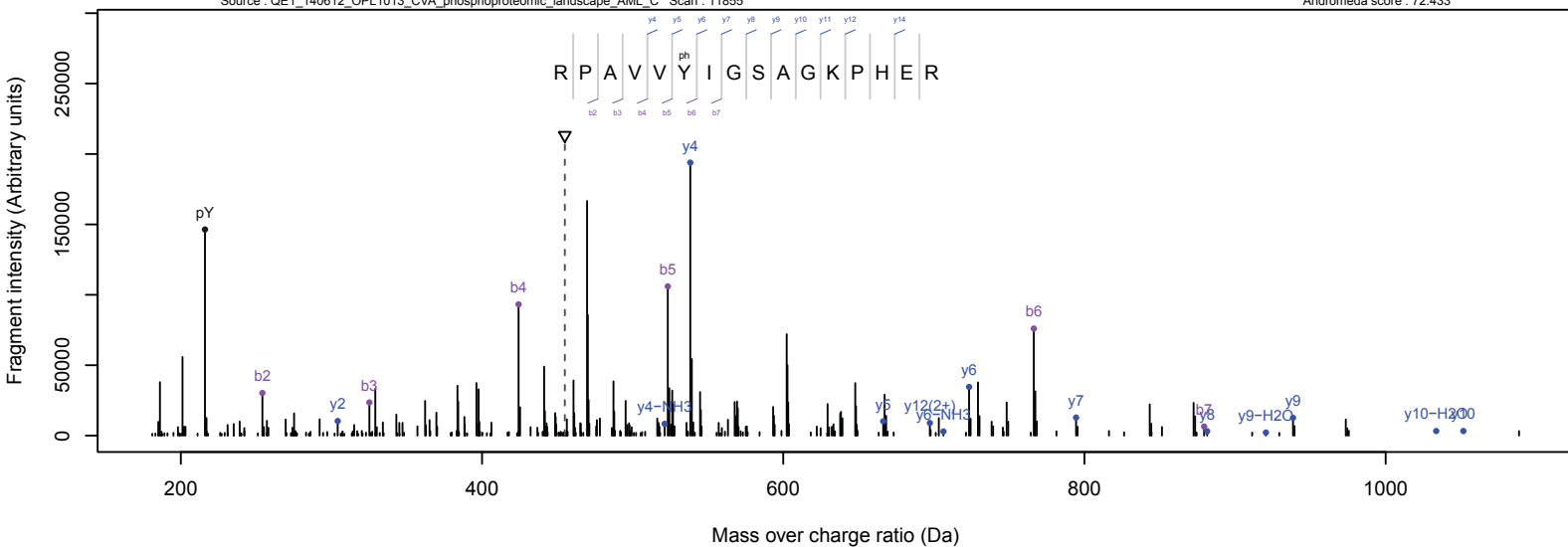
## LCEQLSTVIPNSHVYY R-3+



RPAVVY(ph)IGSAGK<sup>4+</sup>

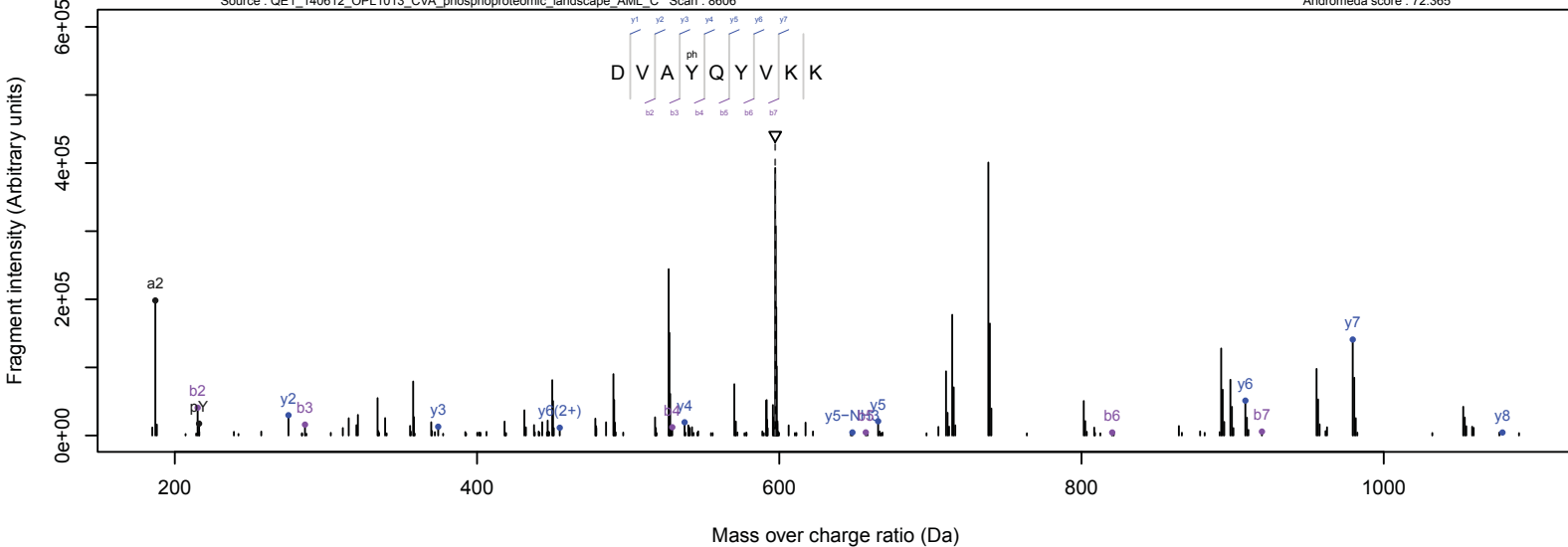
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11855

Andromeda score : 72.433

DVAY(ph)QYVKK<sup>2+</sup>

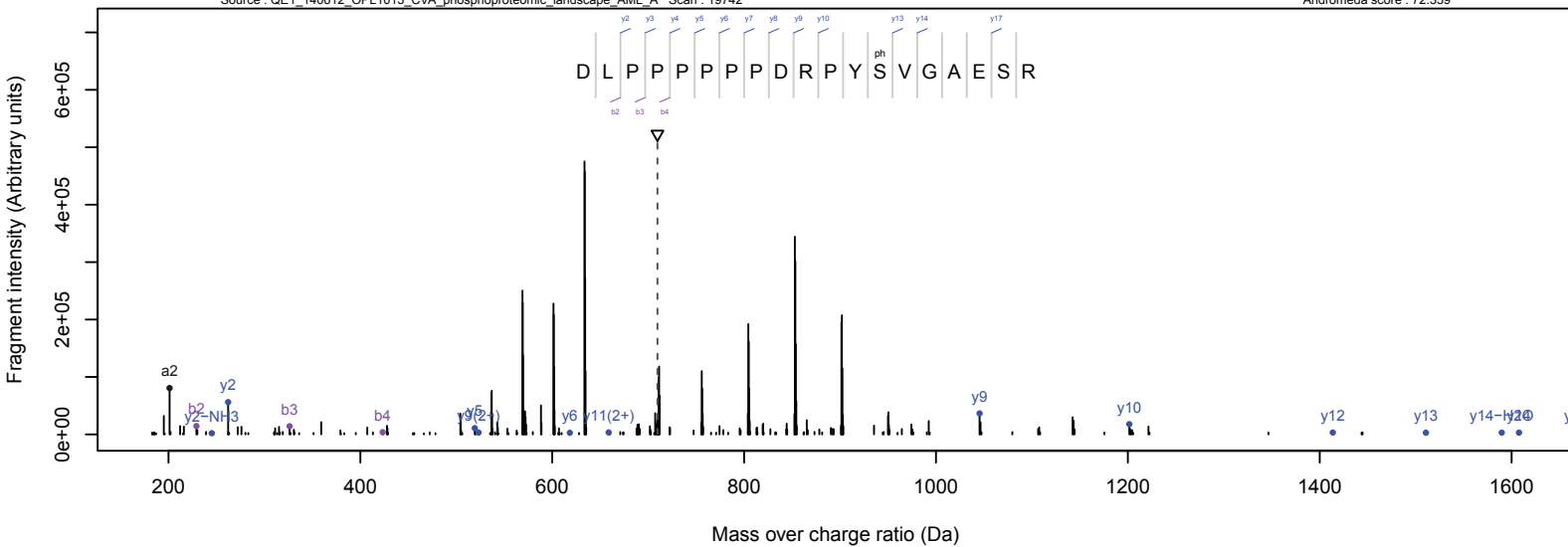
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8606

Andromeda score : 72.365

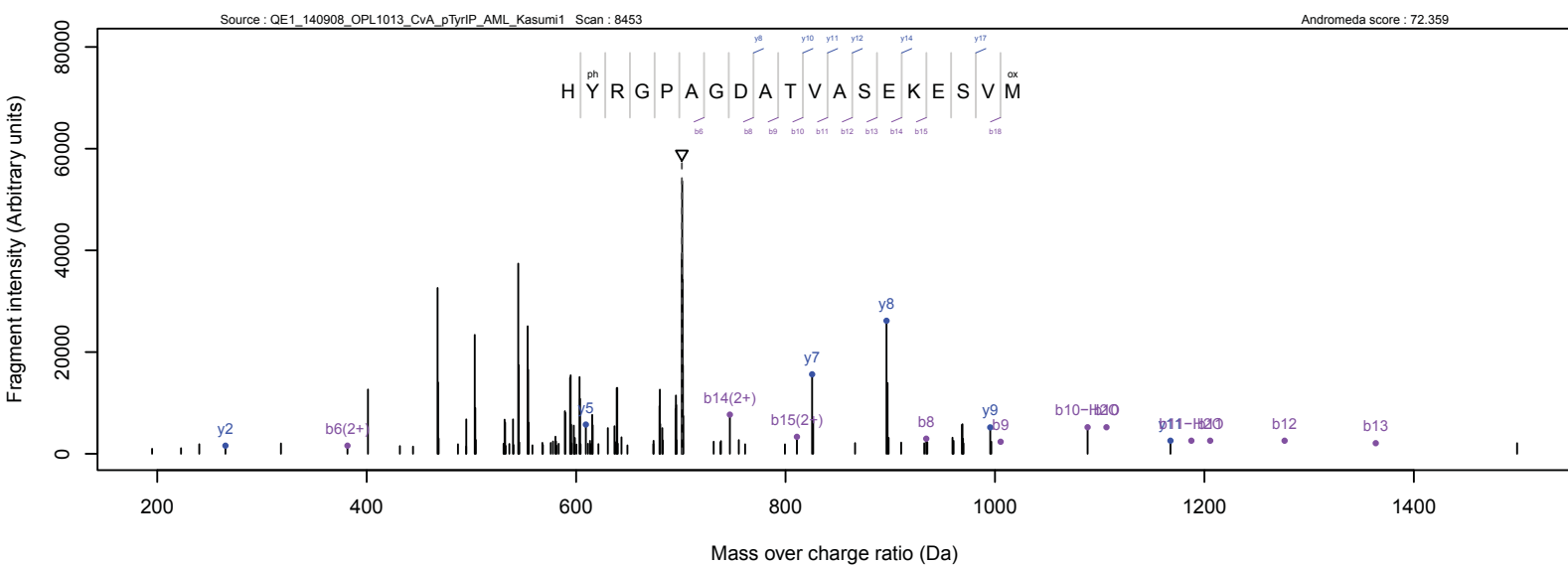
DLPPPPPPDRPYS(ph)VGAESR<sup>3+</sup>

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 19742

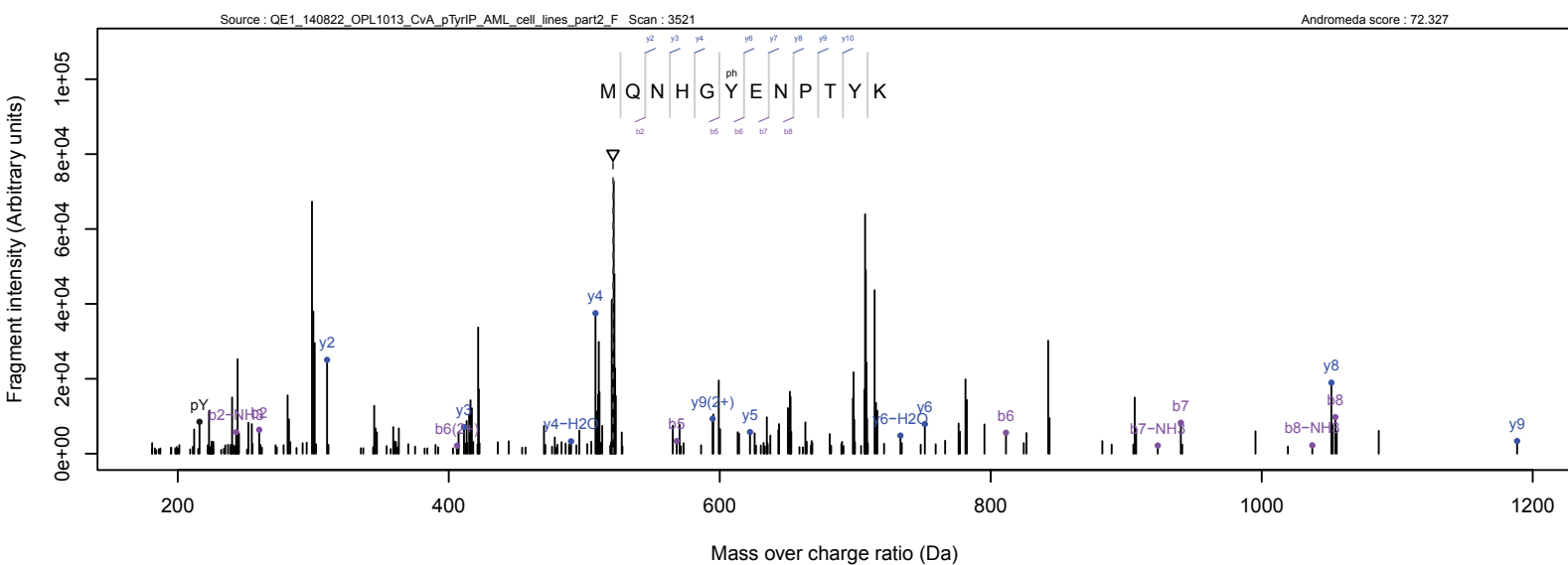
Andromeda score : 72.359



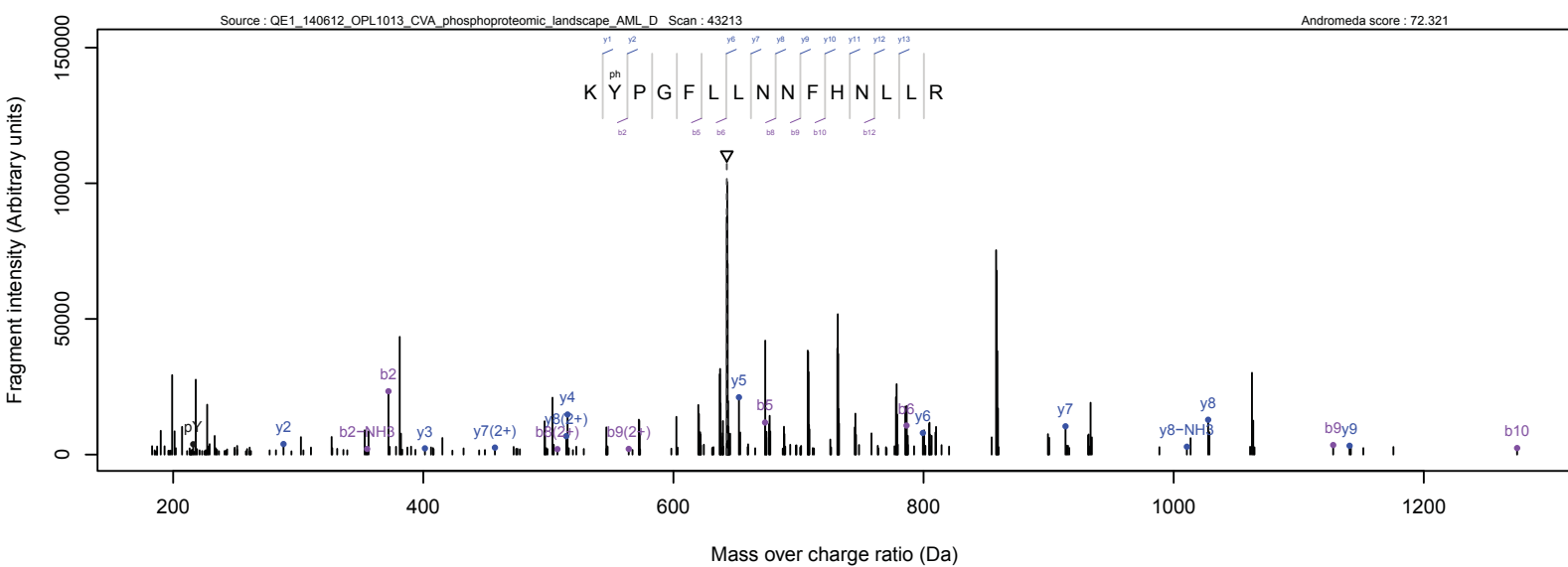
## HY(ph)RGPAGDATVASEKESVM(ox)-3+



## MQNHGY(ph)ENPTYK-3+



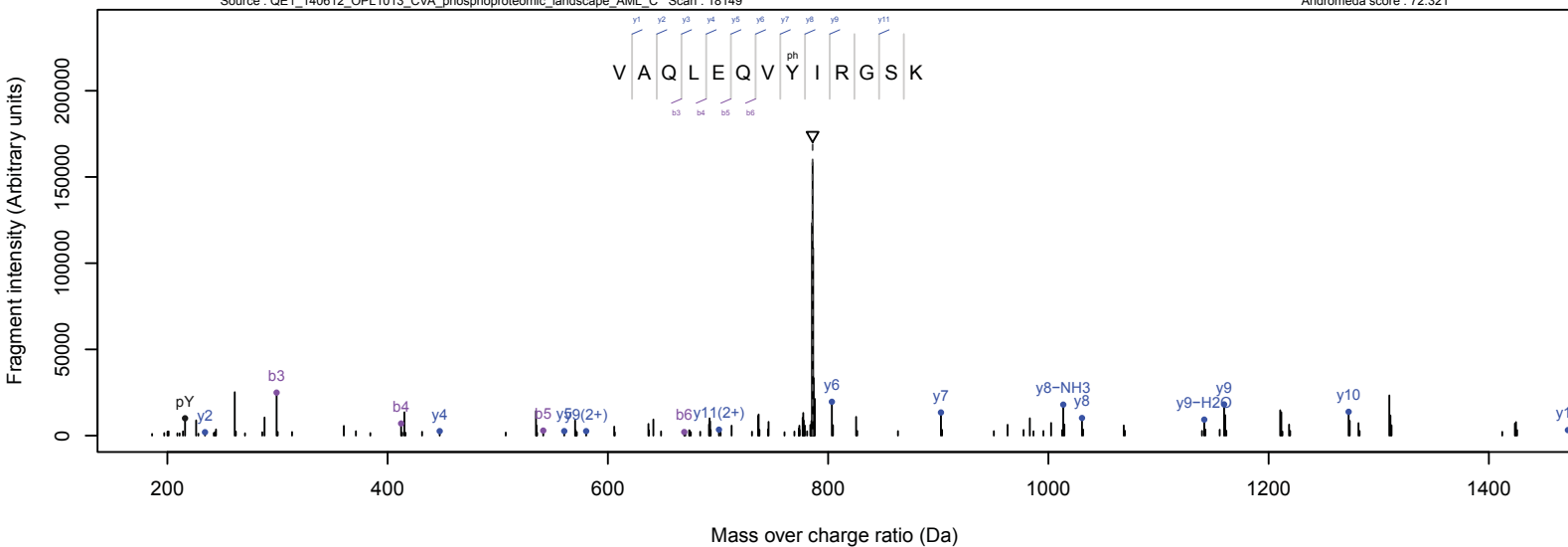
## KY(ph)PGFLLNFFHLLR-3+



VAQLEQVY(ph)IRGSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18149

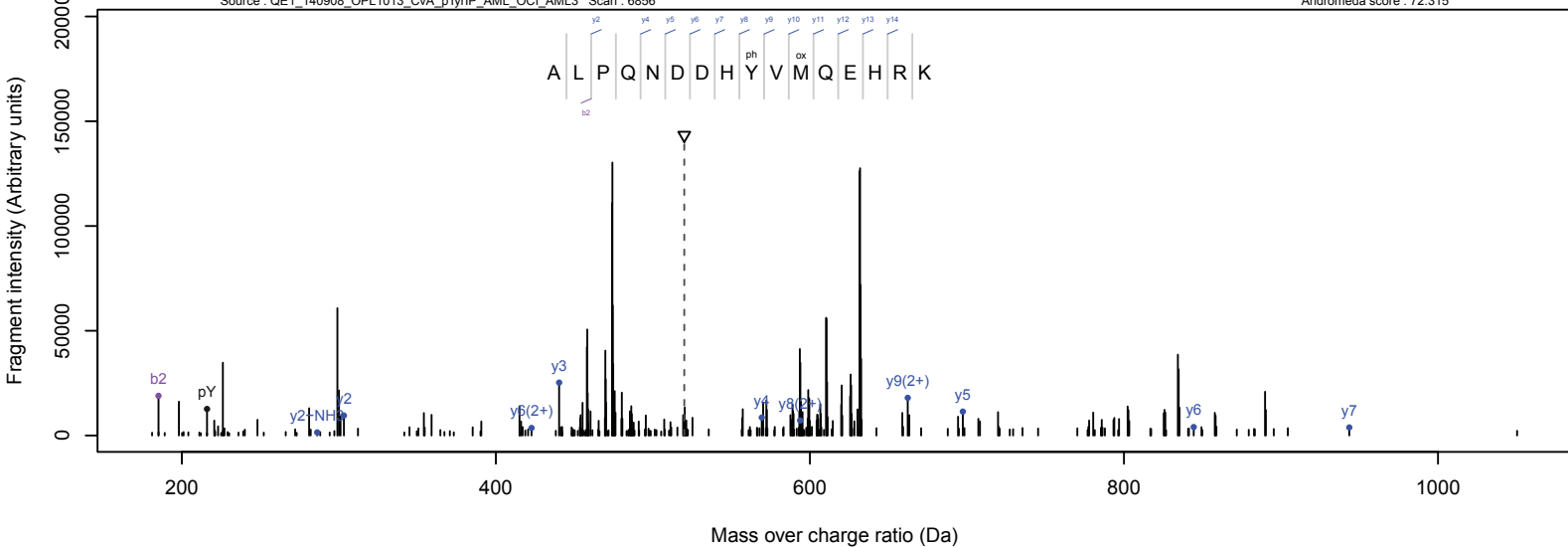
Andromeda score : 72.321



ALPQNDDHY(ph)VM(ox)QEHRK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 6856

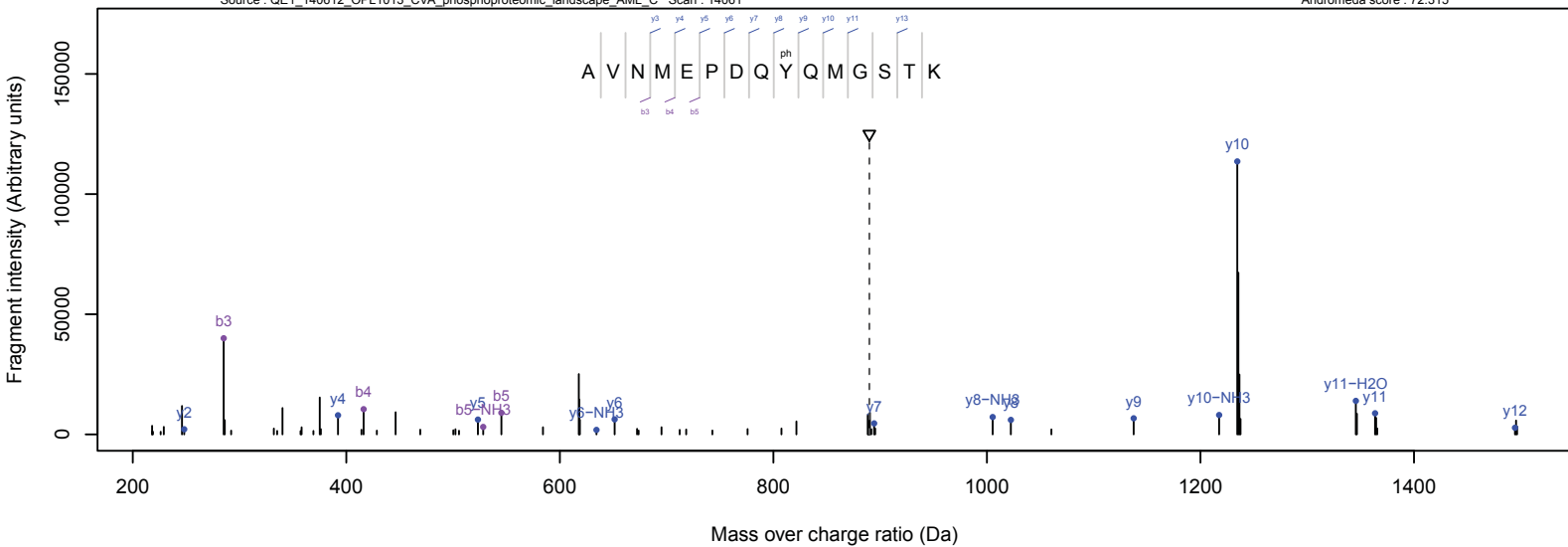
Andromeda score : 72.315



AVNMEPDQYQMGSTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14061

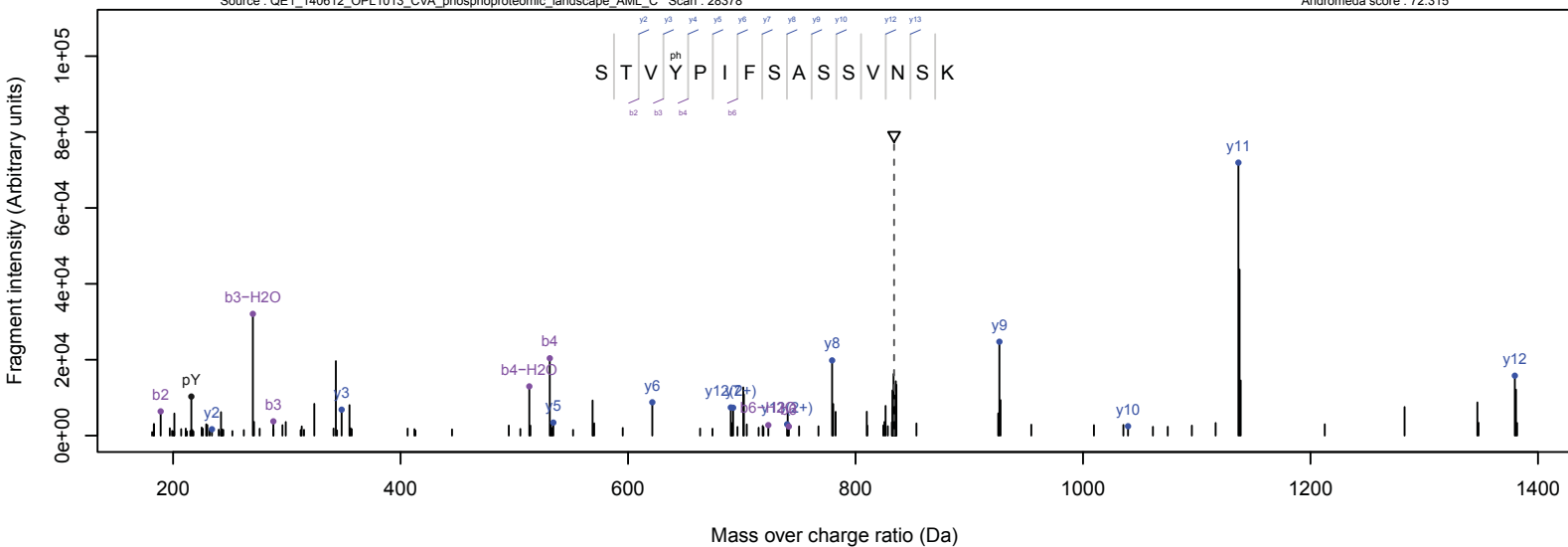
Andromeda score : 72.315



## STVY(ph)PIFSASSVNSK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 28378

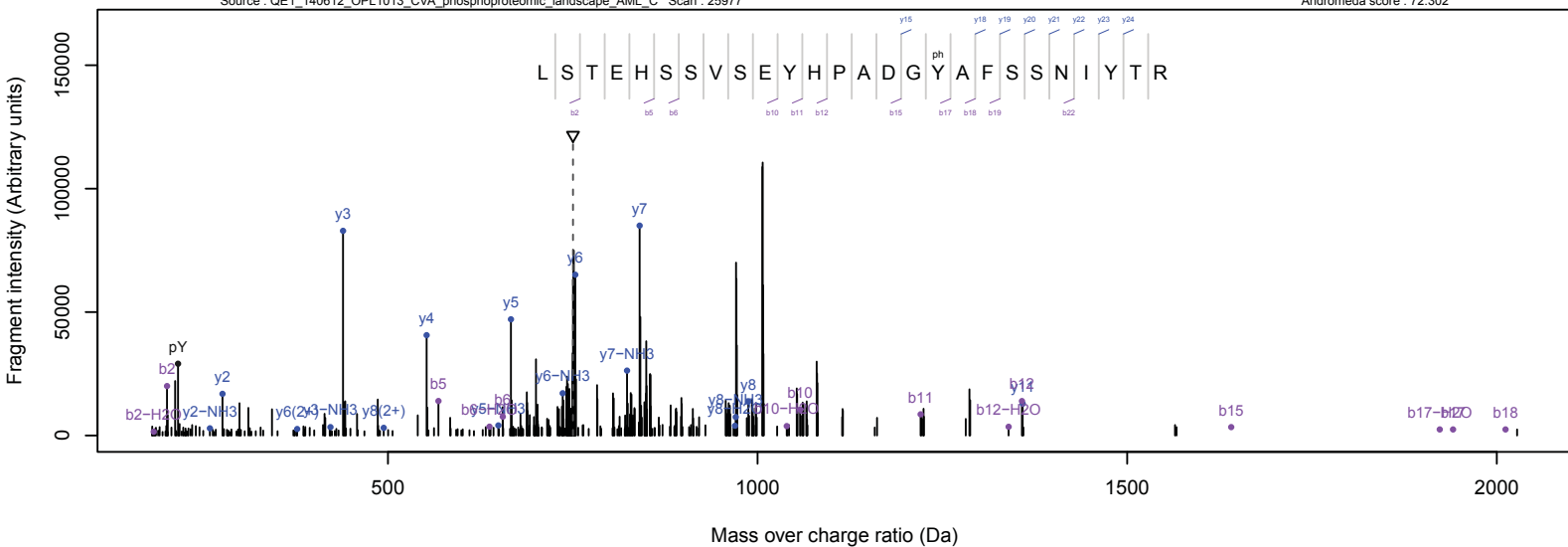
Andromeda score: 72.315



## LSTEHSVSEYHPADG(ph)AFSSNIYTR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25977

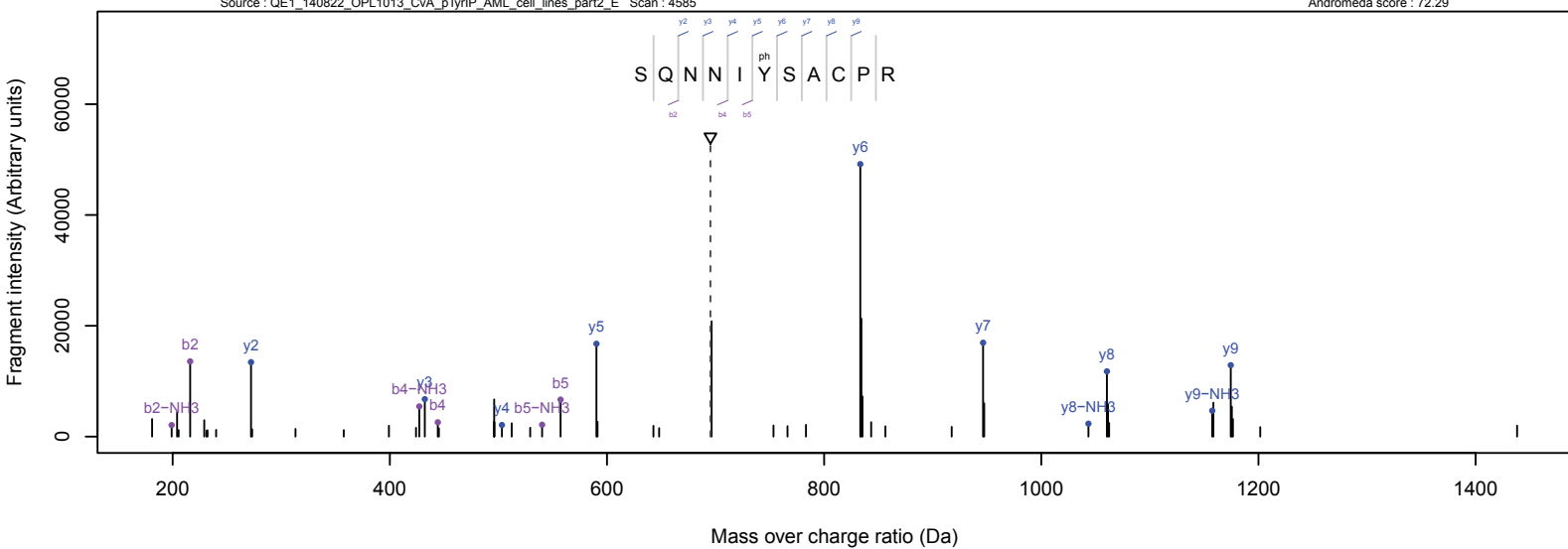
Andromeda score: 72.302



## SQNNIY(ph)SACPR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 4585

Andromeda score: 72.29

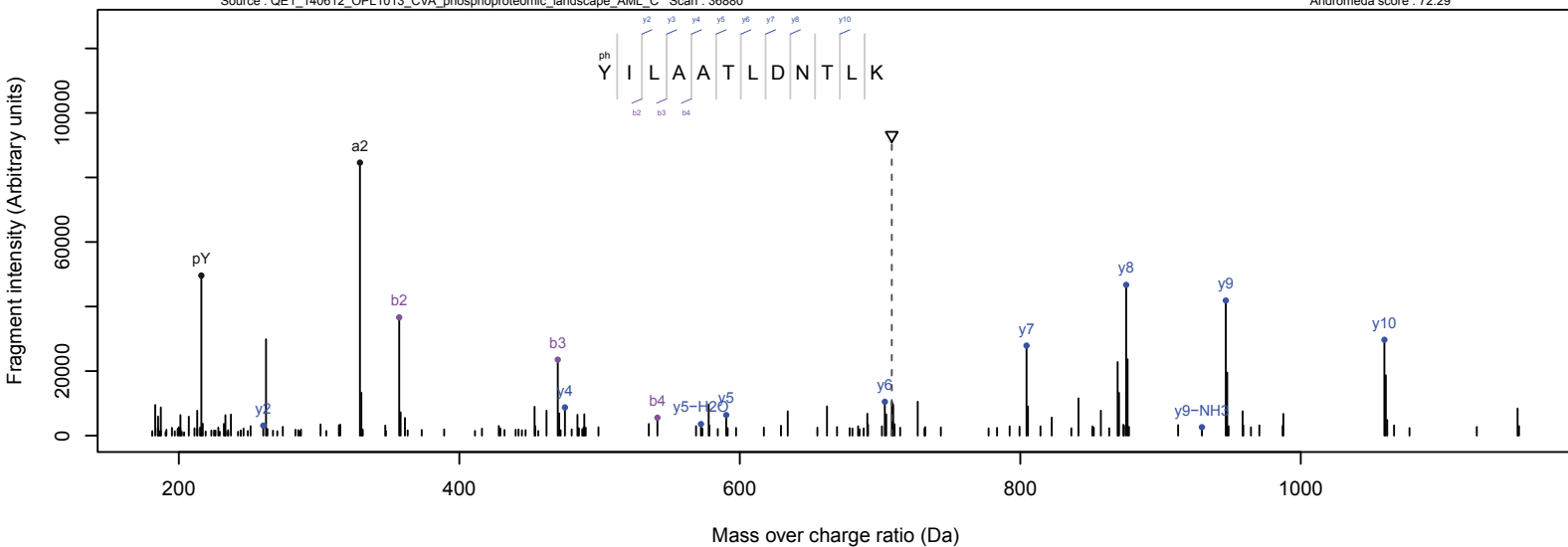




Y(ph)ILAATLDNTLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36880

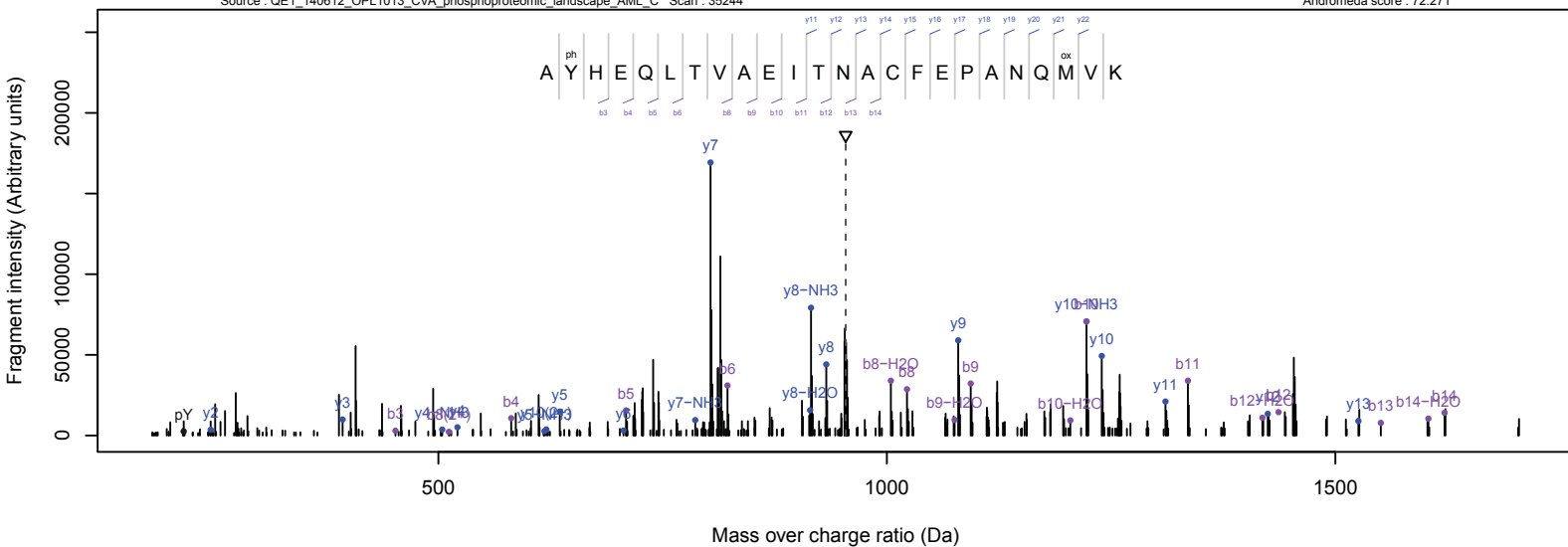
Andromeda score : 72.29



AY(ph)HEQLTVAEITNACFEPANQM(ox)VK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35244

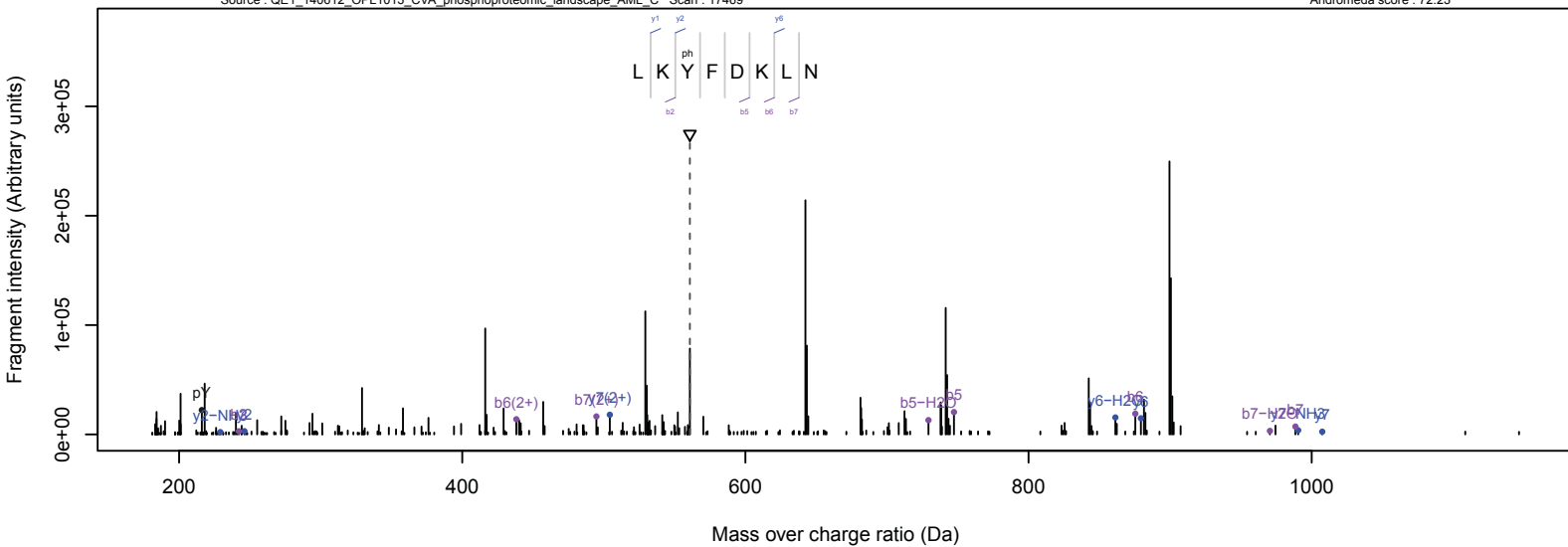
Andromeda score : 72.271



LKY(ph)FDKLN-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17469

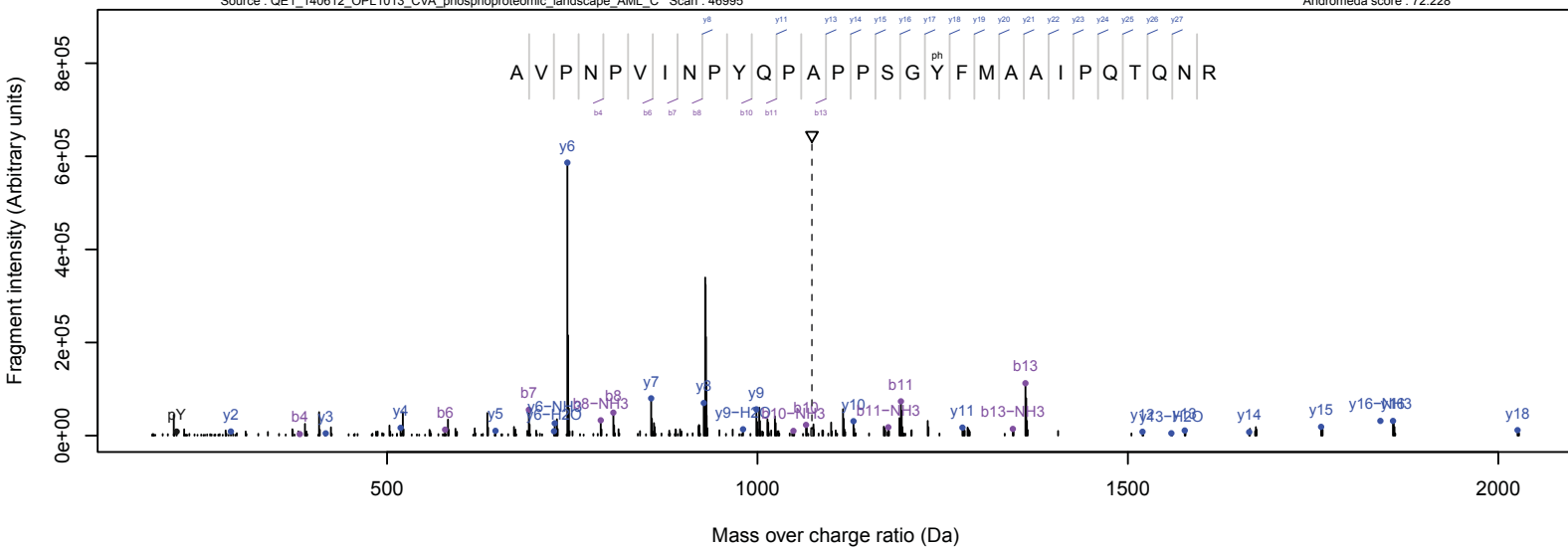
Andromeda score : 72.23



AVPNPVINPYQPAPPSPGYFMAAIPQTQNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46995

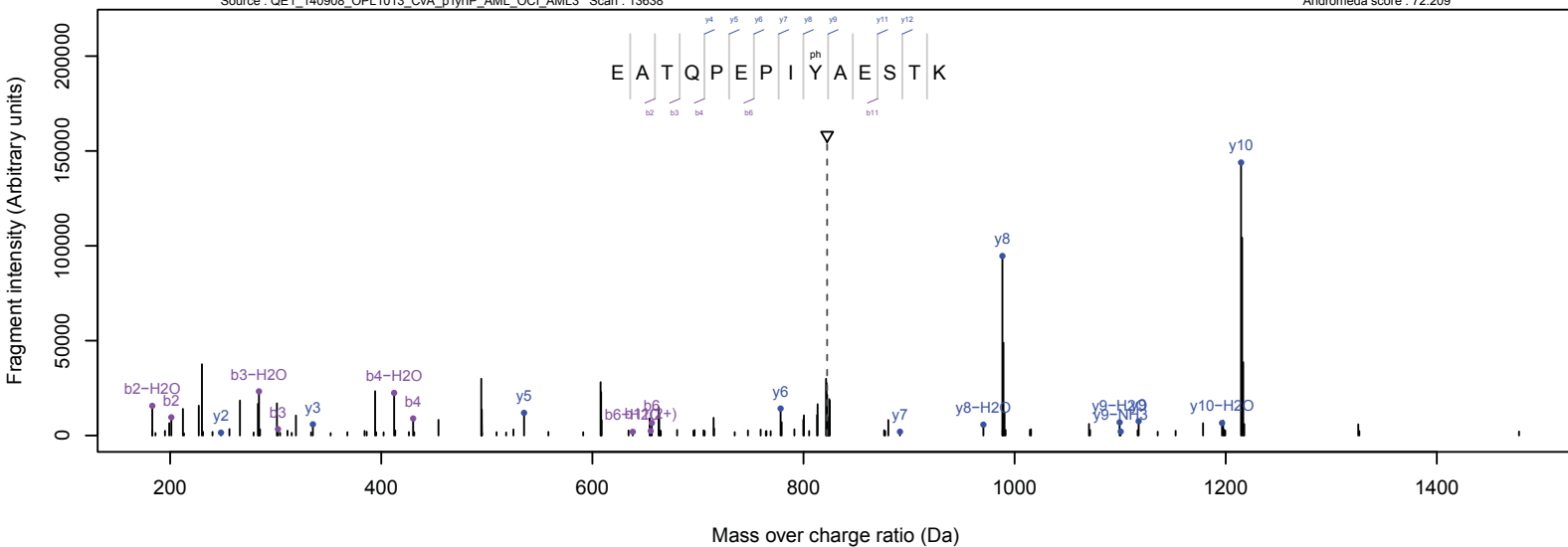
Andromeda score : 72.228



EATQPEPIY(ph)AESTK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 13638

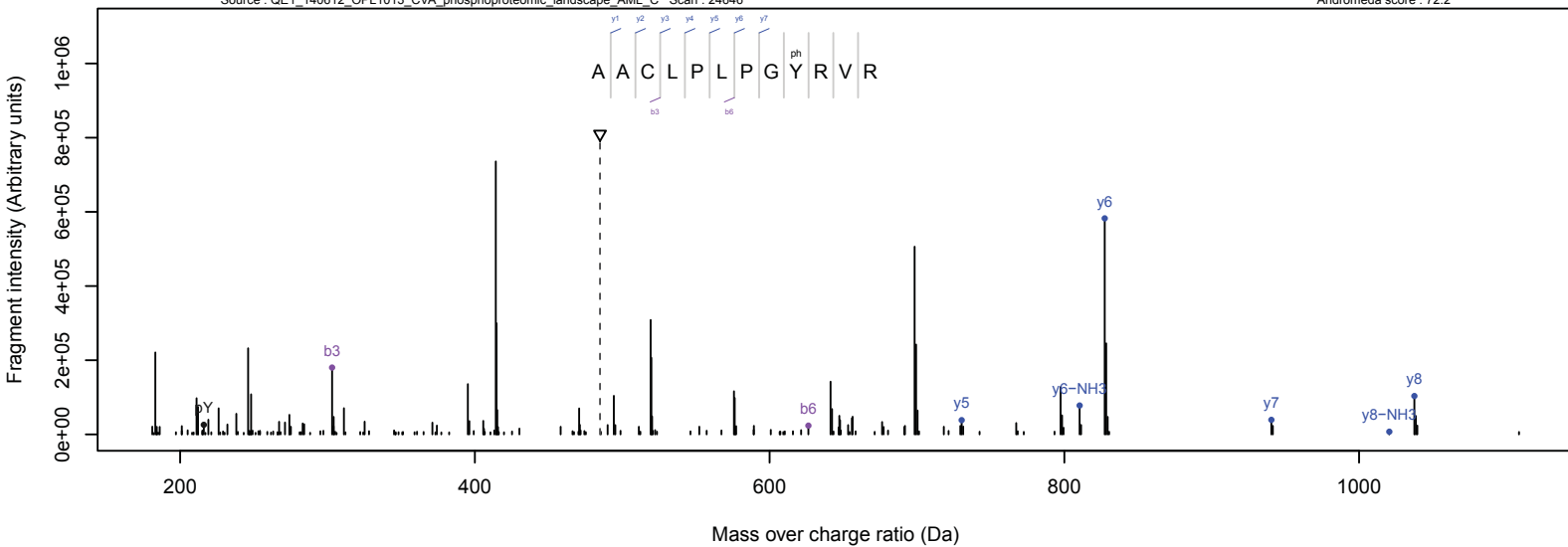
Andromeda score : 72.209



AACLPLPGY(ph)RVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24646

Andromeda score : 72.2

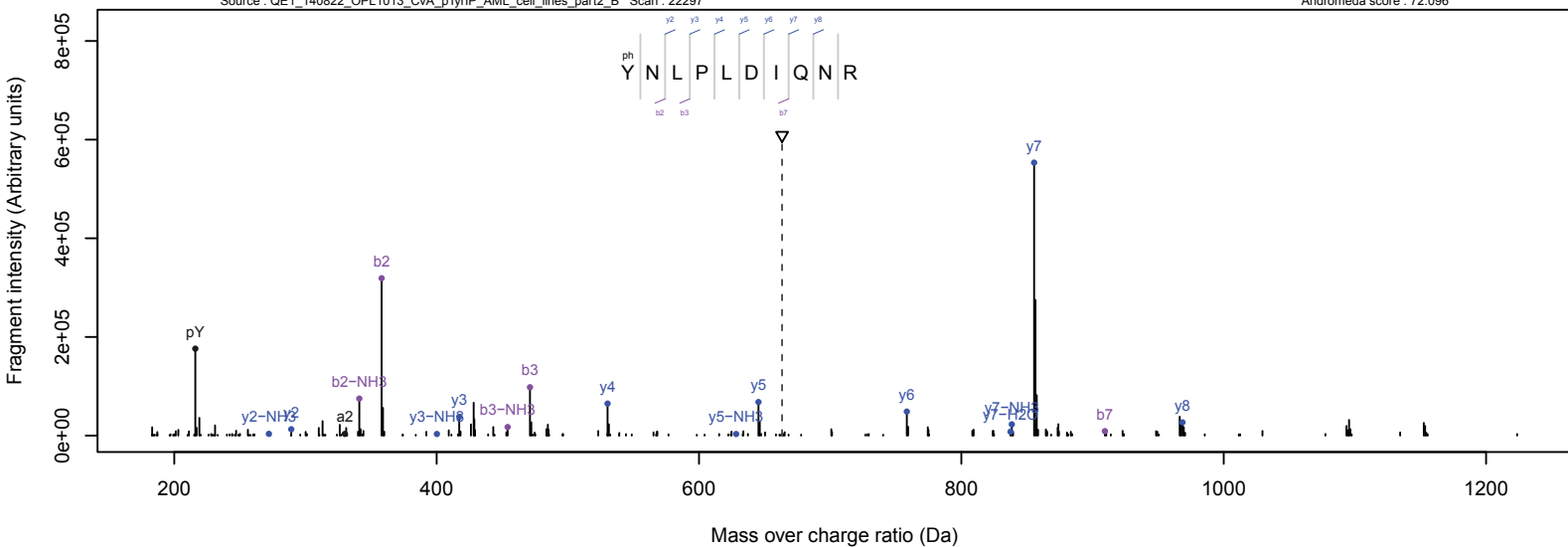




Y(ph)NLPLDIQNR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 22297

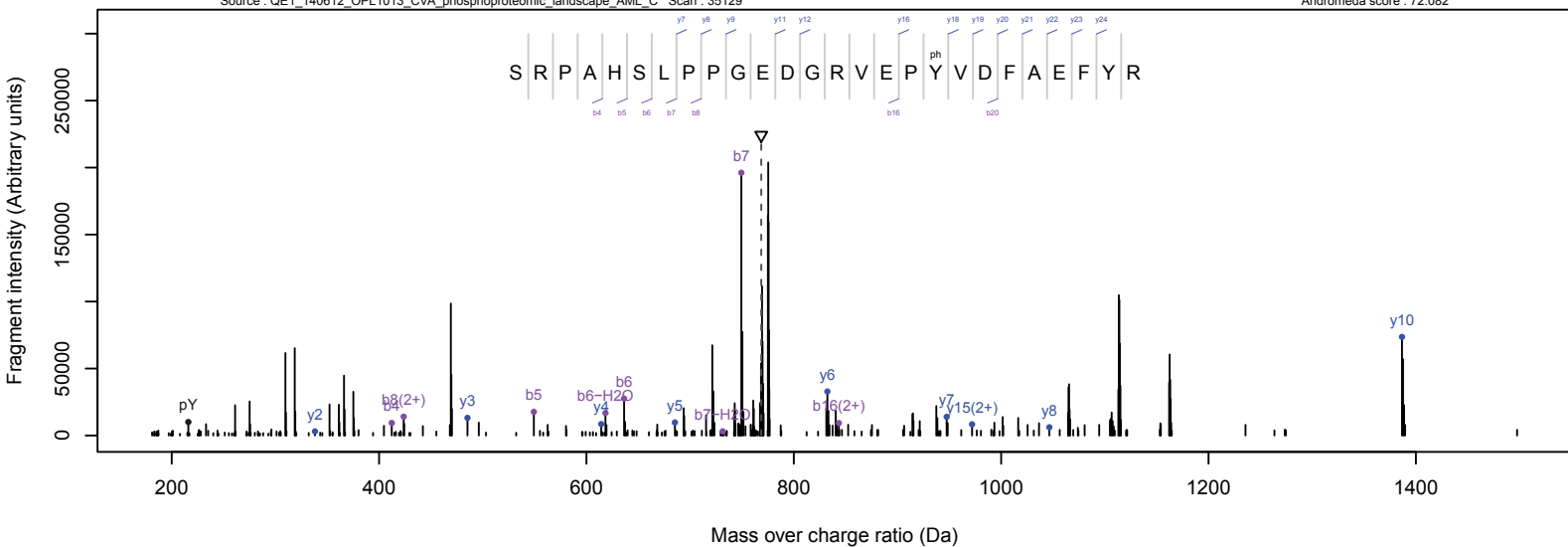
Andromeda score : 72.096



SRPAHSLPPGEDGRVEPY(ph)VDFAEFYR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35129

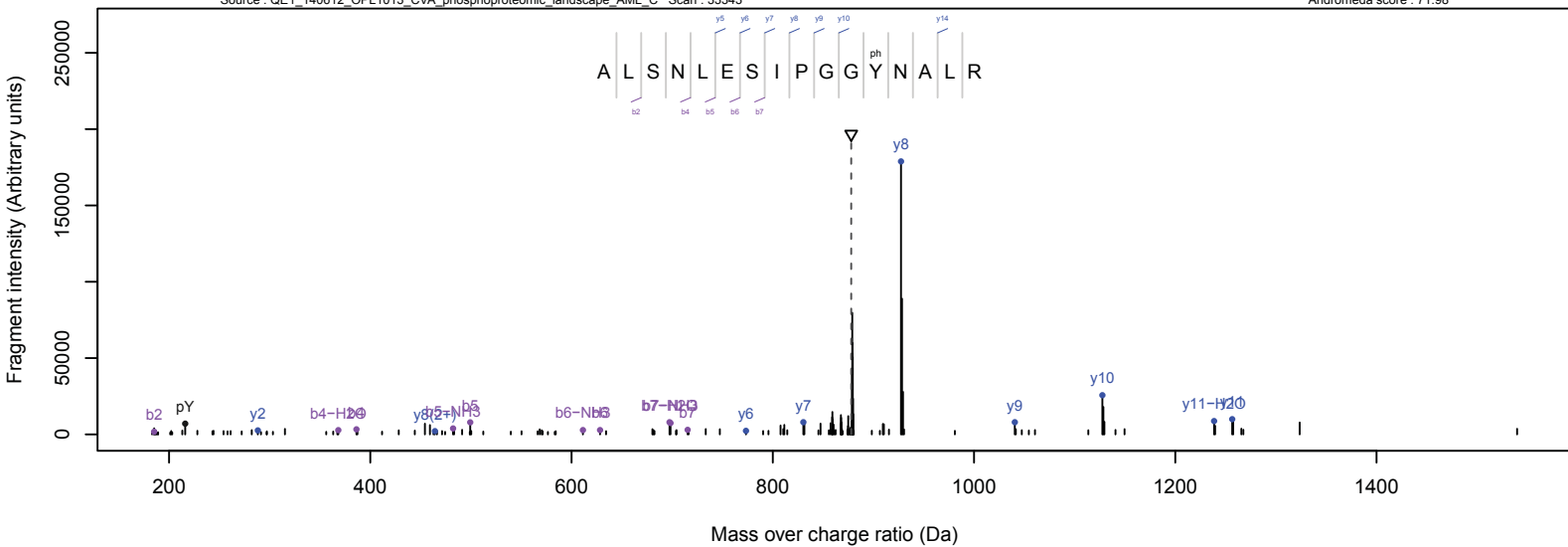
Andromeda score : 72.082



ALSNLEIPGGY(ph)NALR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33343

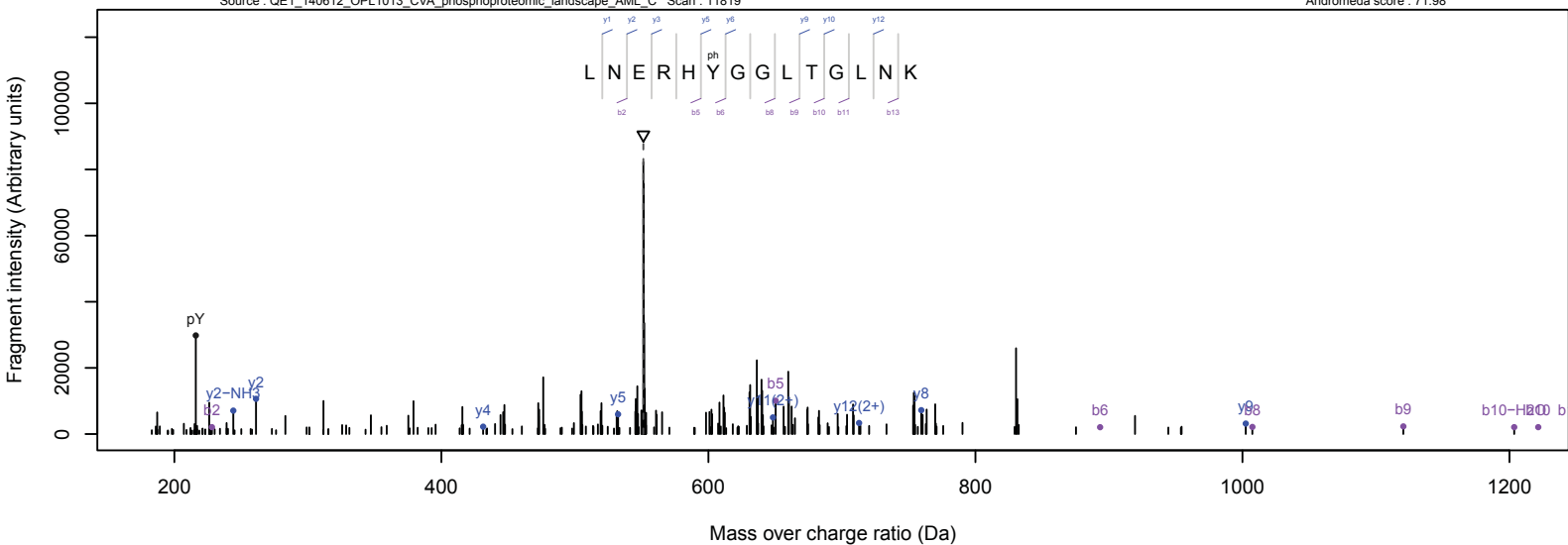
Andromeda score : 71.98



LNERHY(ph)GGLTGLNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11819

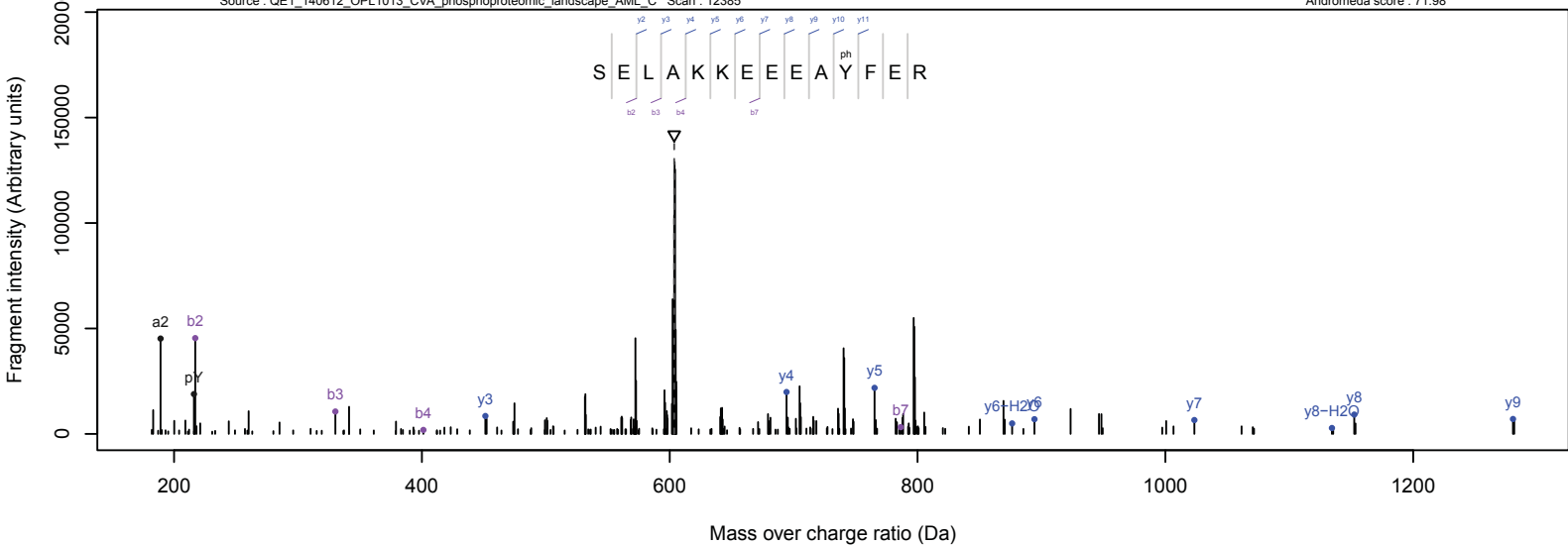
Andromeda score : 71.98



SELAKKEEEAY(ph)FER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12385

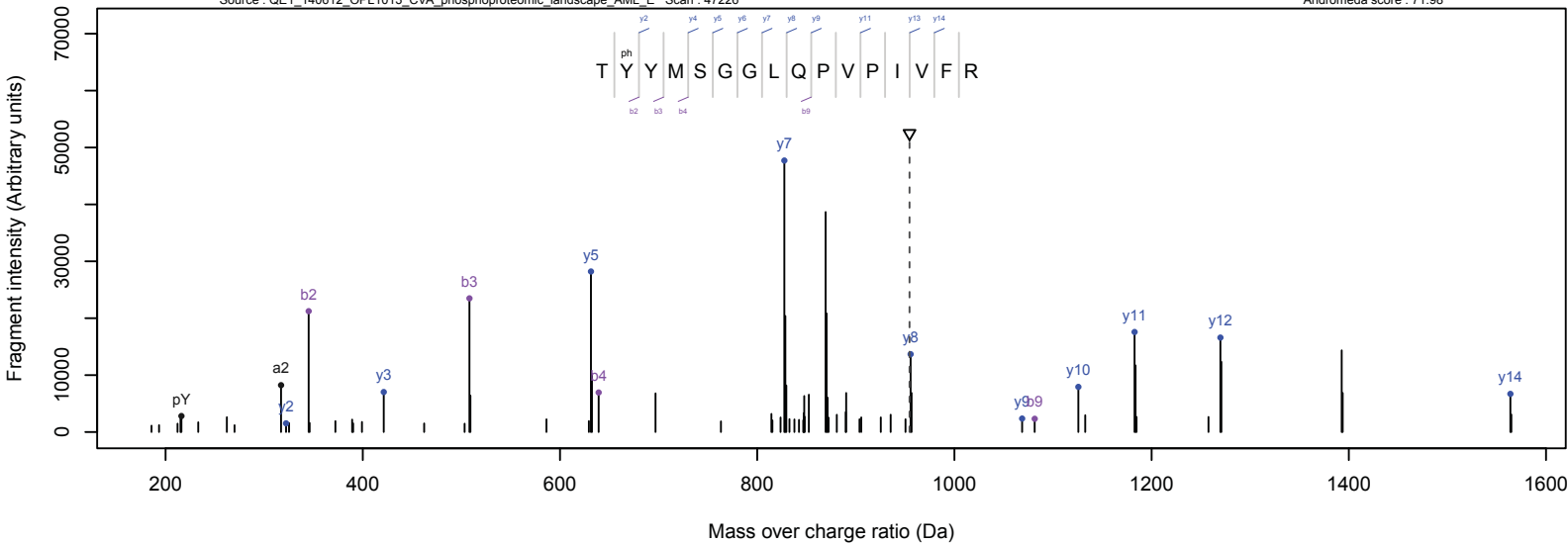
Andromeda score : 71.98



TY(ph)YMSGGLQPPIVFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 47226

Andromeda score : 71.98

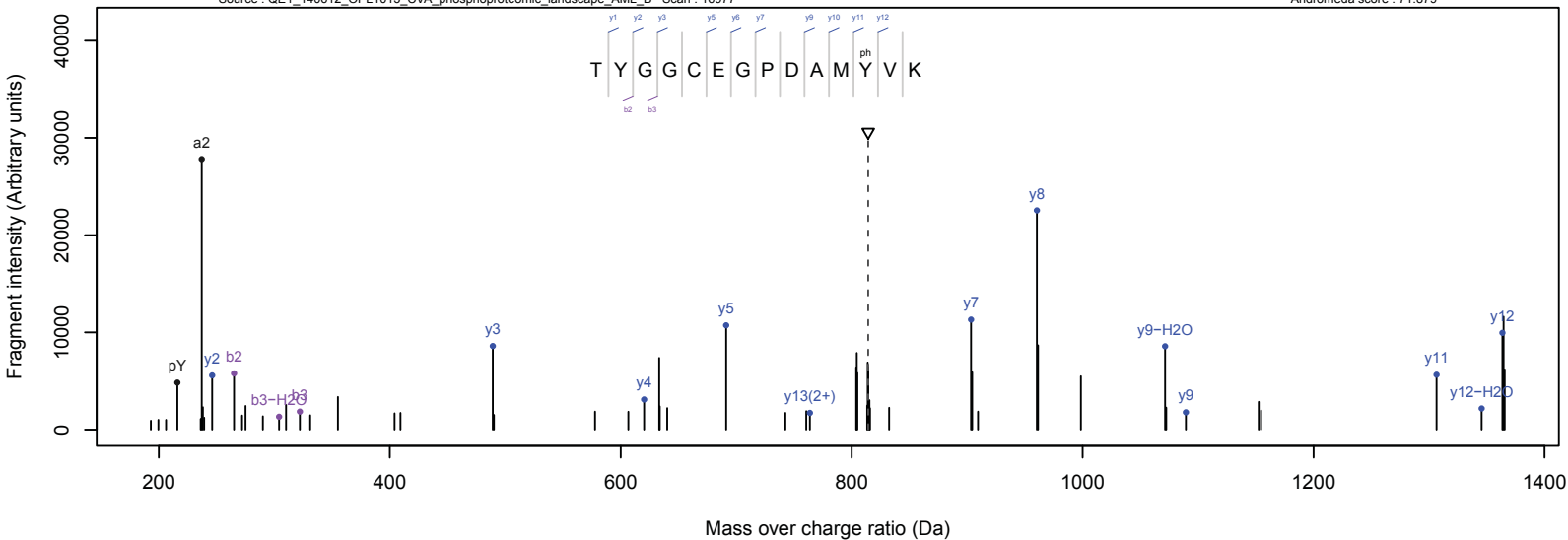




## TYGGCEGPDAMY(ph)VK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 16977

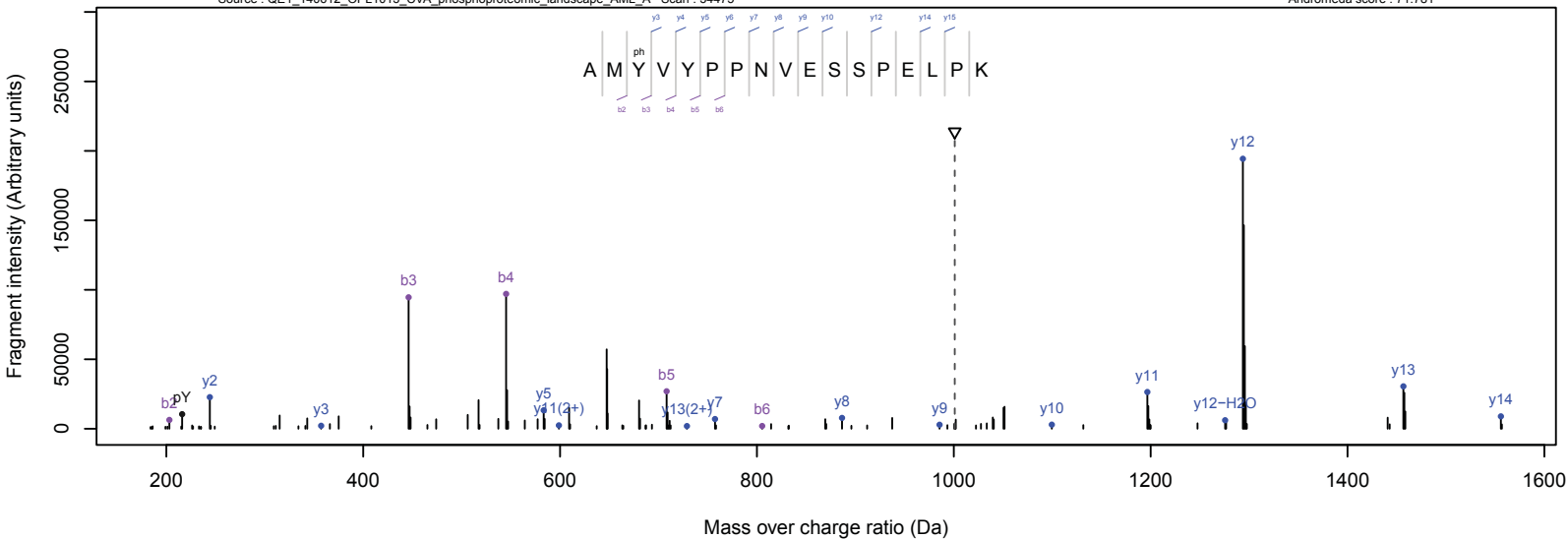
Andromeda score: 71.879



## AMY(ph)VYPNVESPELPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 34475

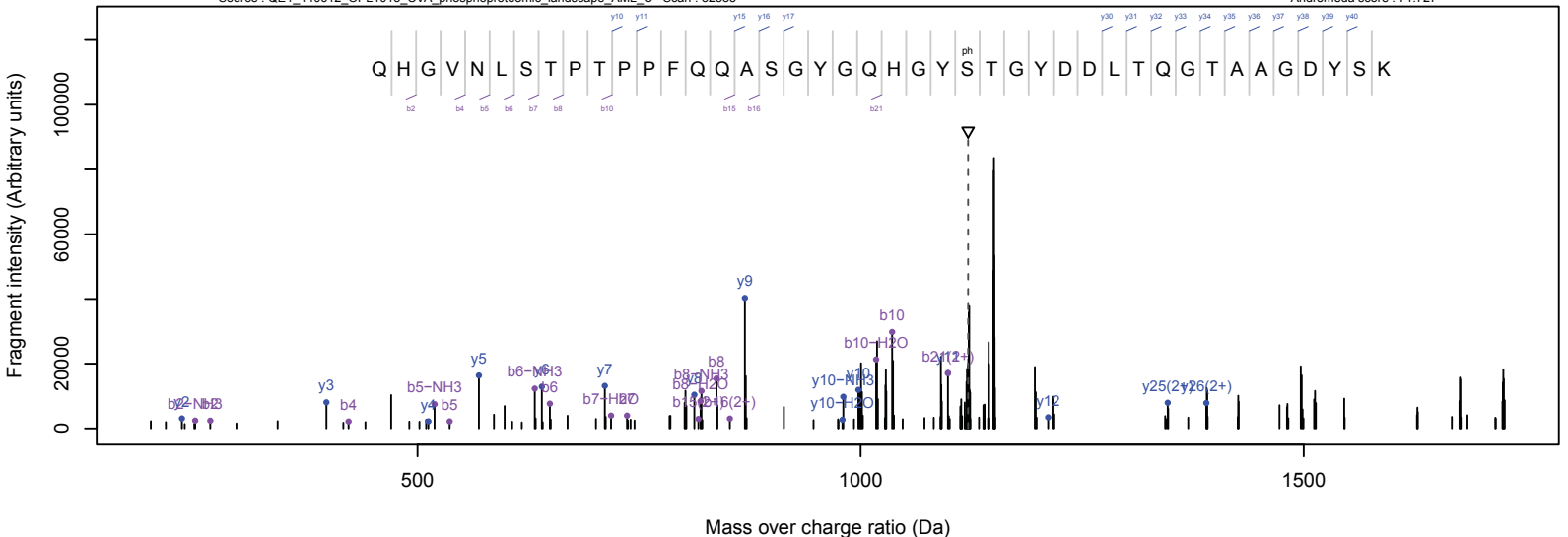
Andromeda score: 71.781



## QHGVNLSPTPPFQQASGYGQHGY(ph)TGYDDLQTGAAGDYSK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 32585

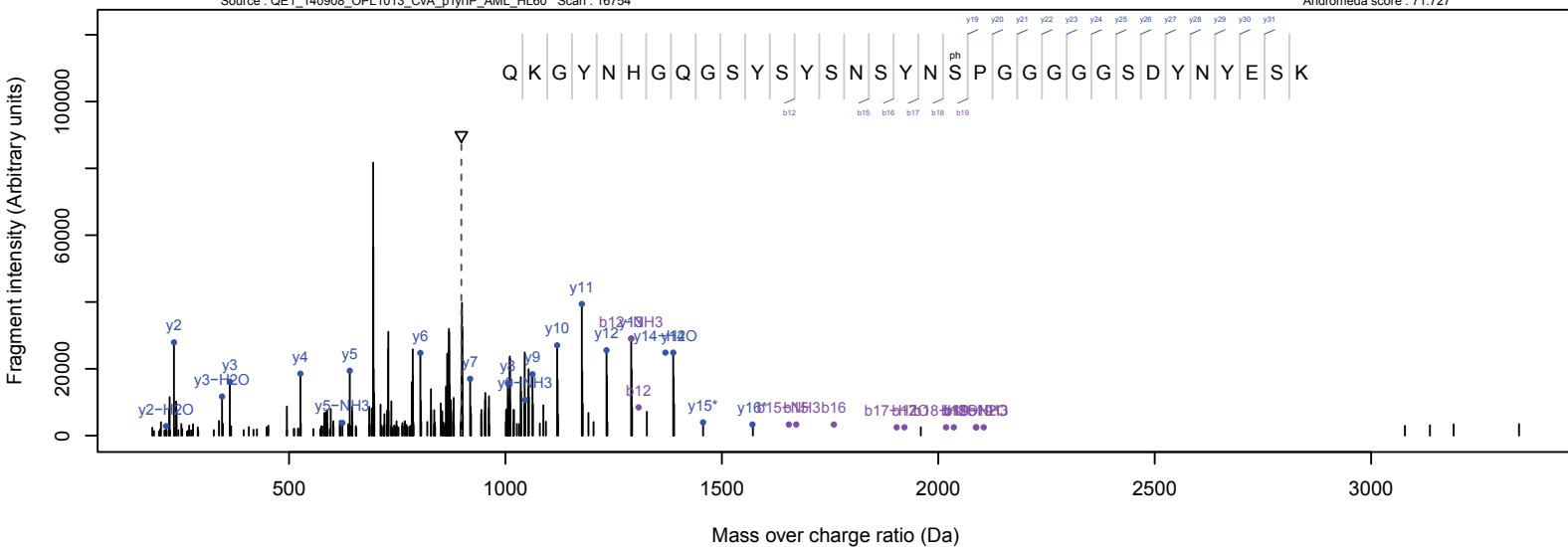
Andromeda score: 71.727



**KQGYNHGQGSYSYSNSYN<sup>ph</sup>PGGGGSDYNYESK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 16754

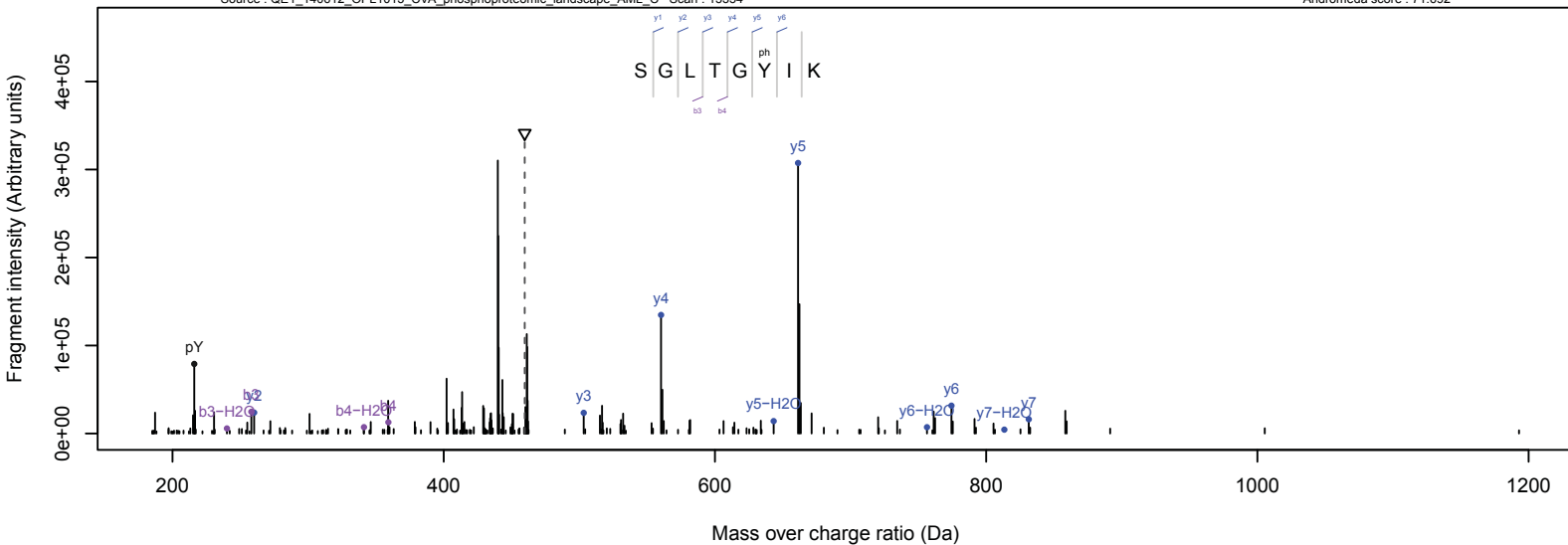
Andromeda score : 71.727



**SGLTGY<sup>ph</sup>IK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15354

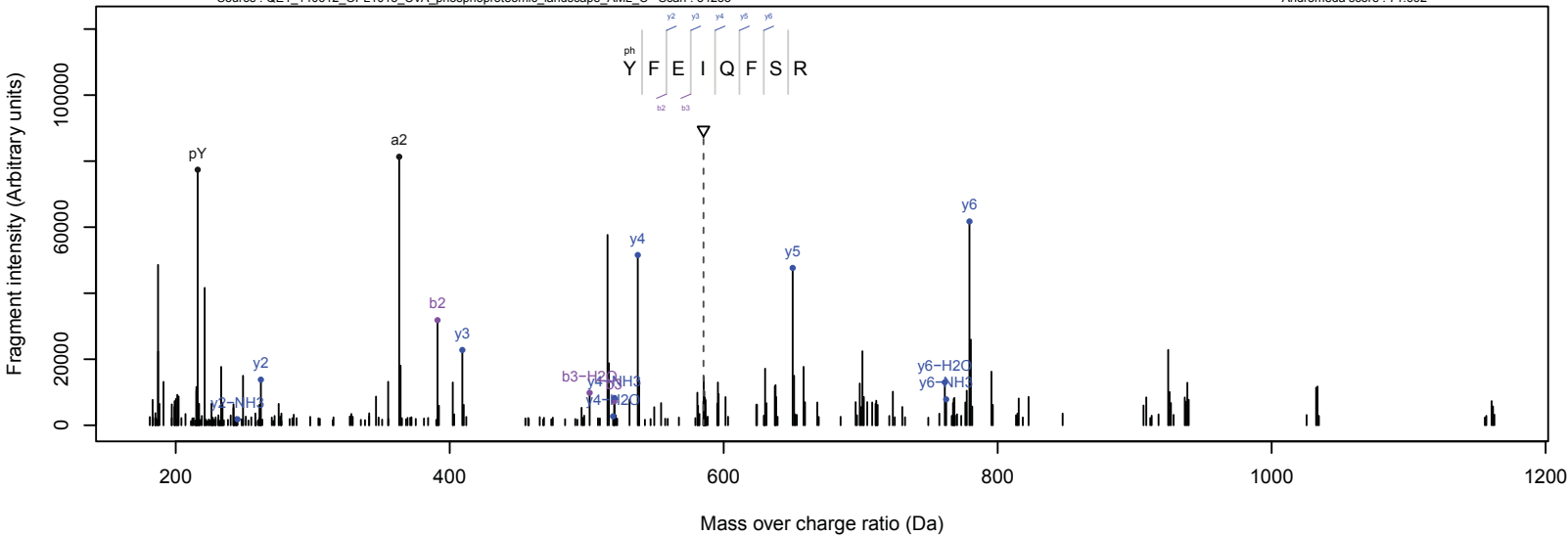
Andromeda score : 71.692



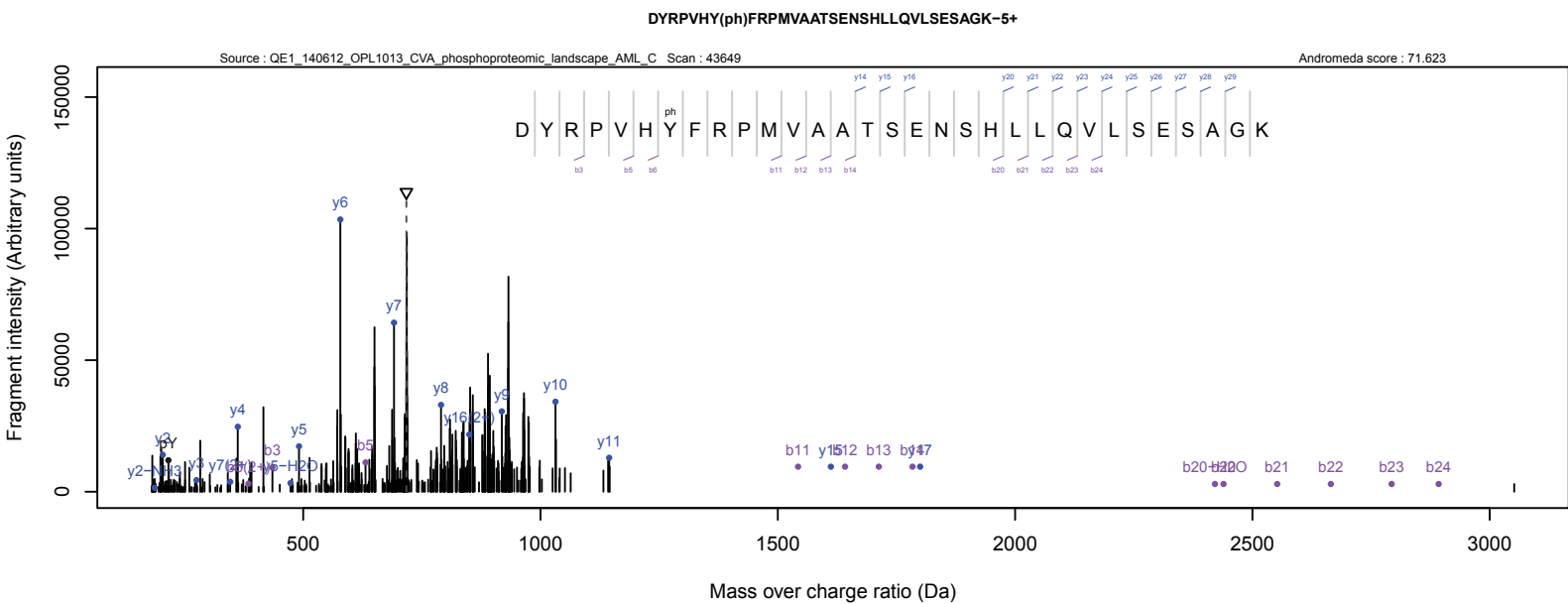
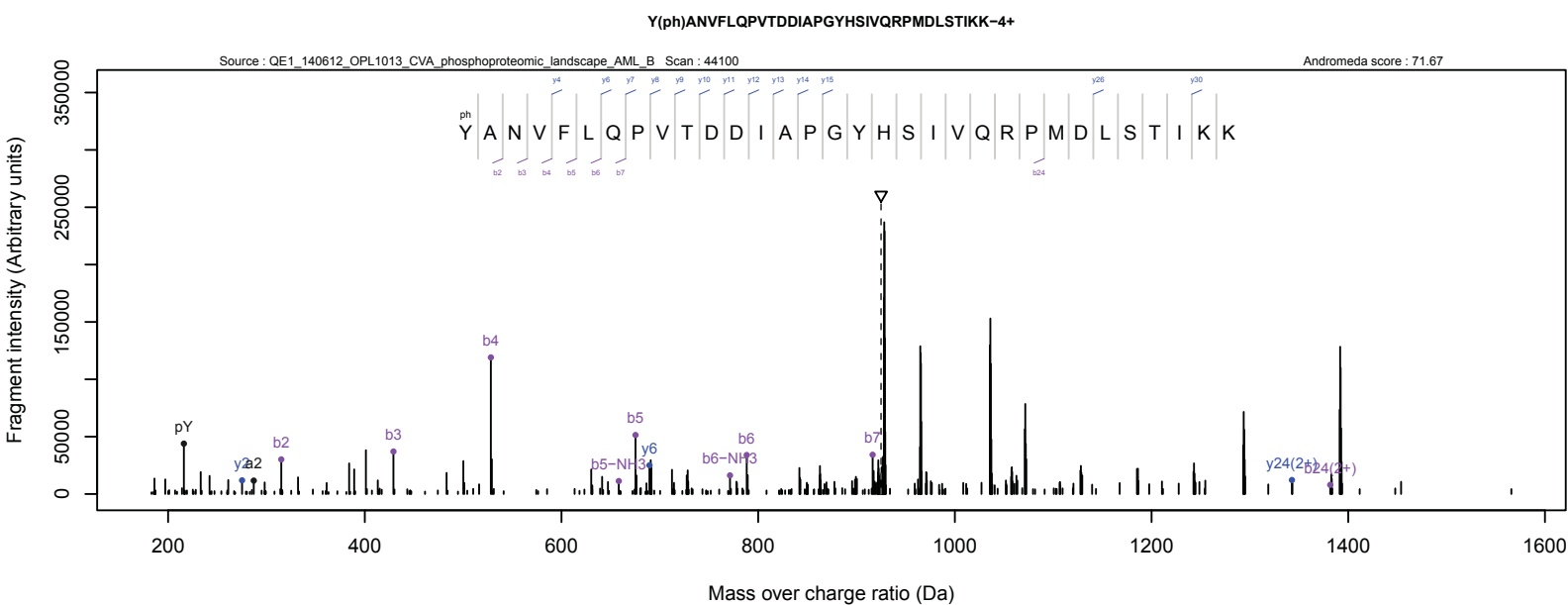
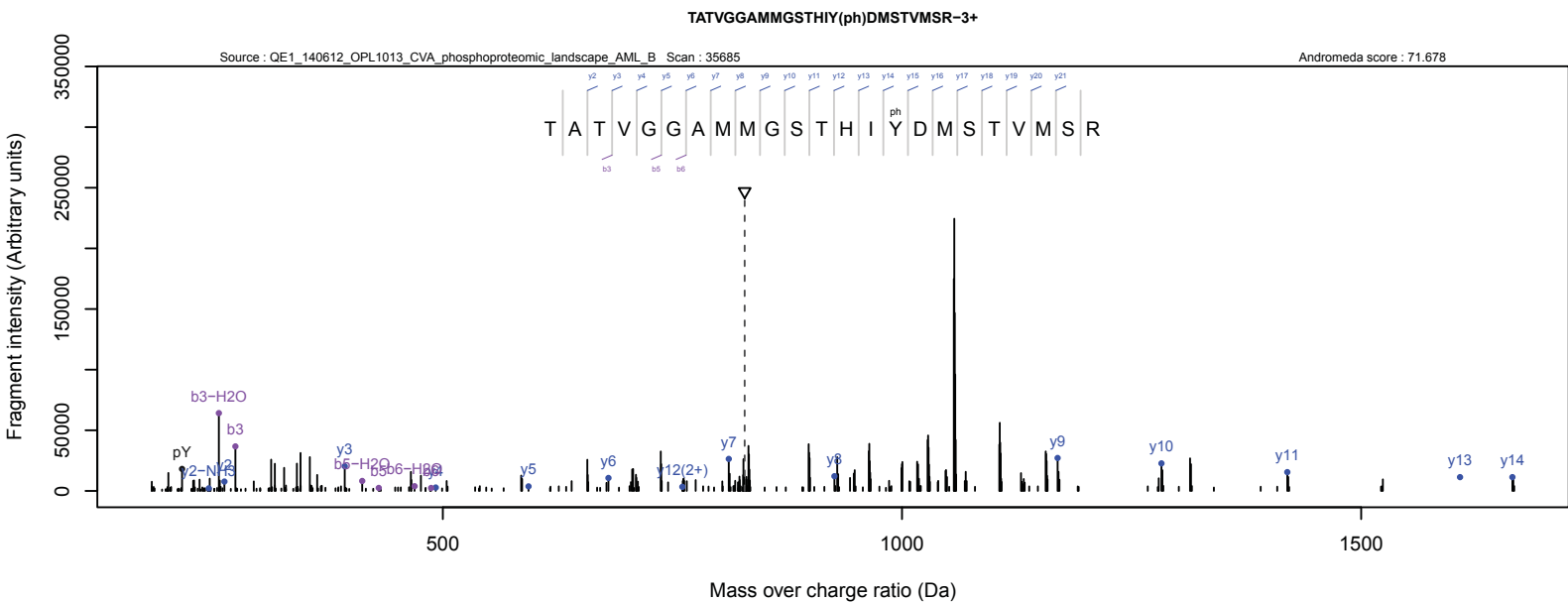
**Y<sup>ph</sup>FEIQFSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34255

Andromeda score : 71.692



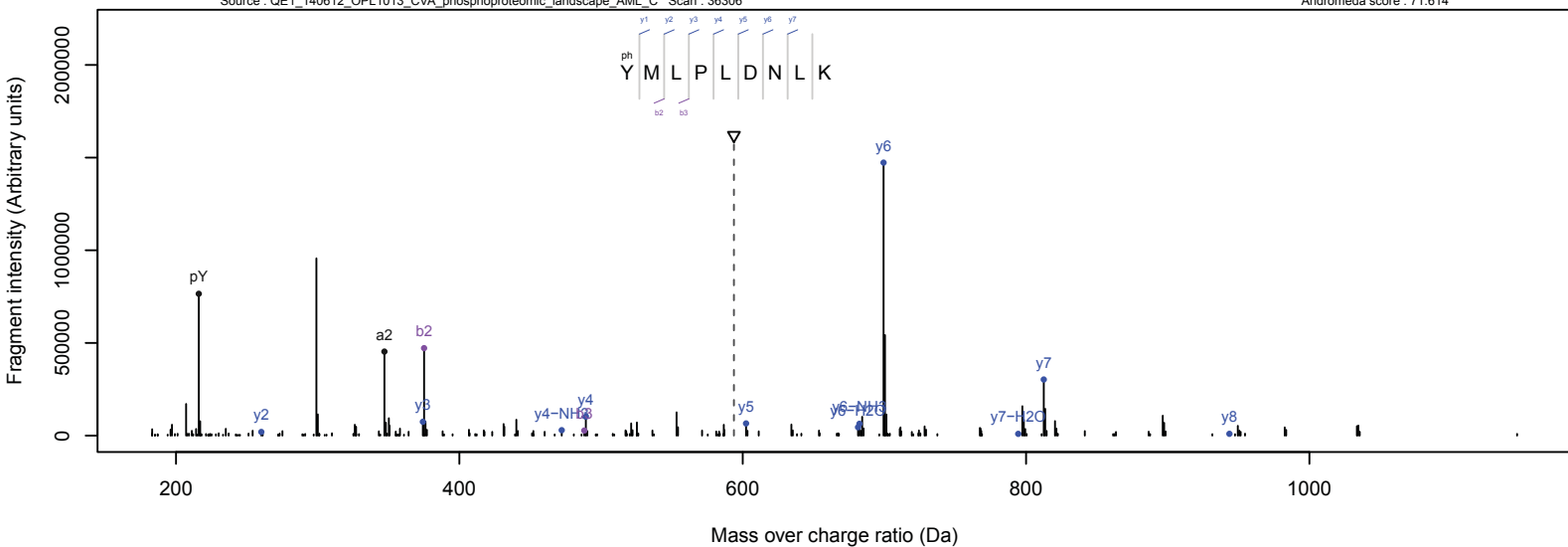




**Y(ph)MLPLDNLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36306

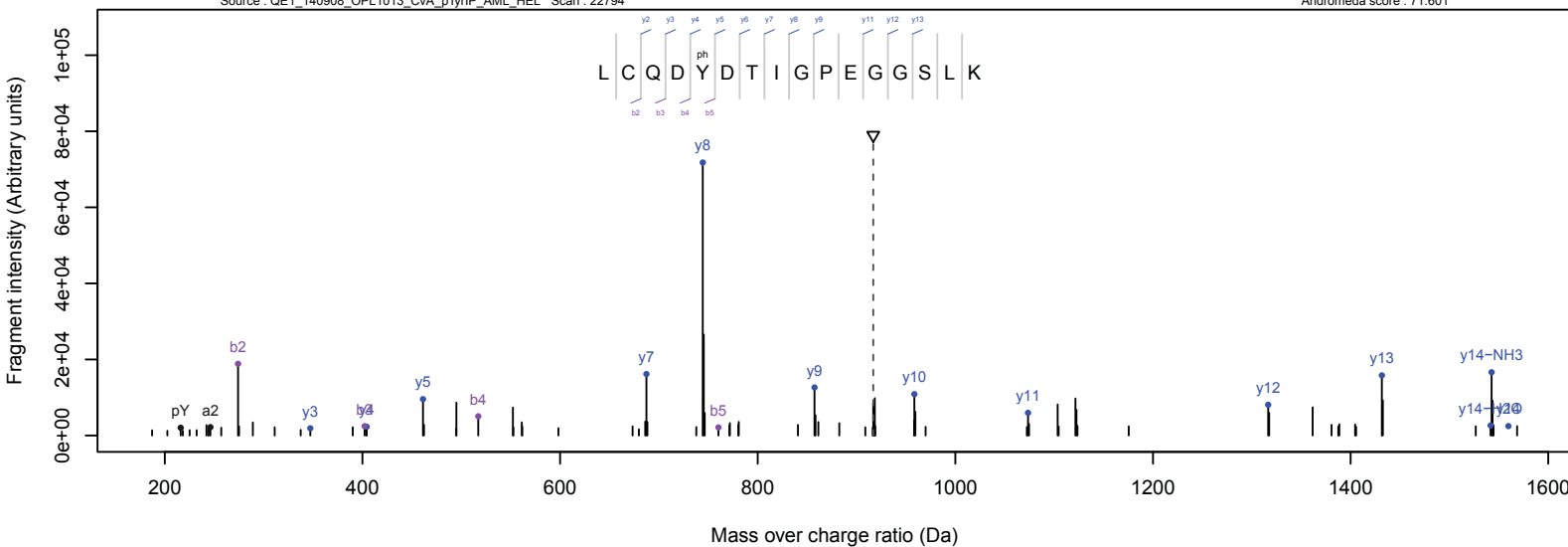
Andromeda score : 71.614



**LCQDY(ph)DTIGPEGSLK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 22794

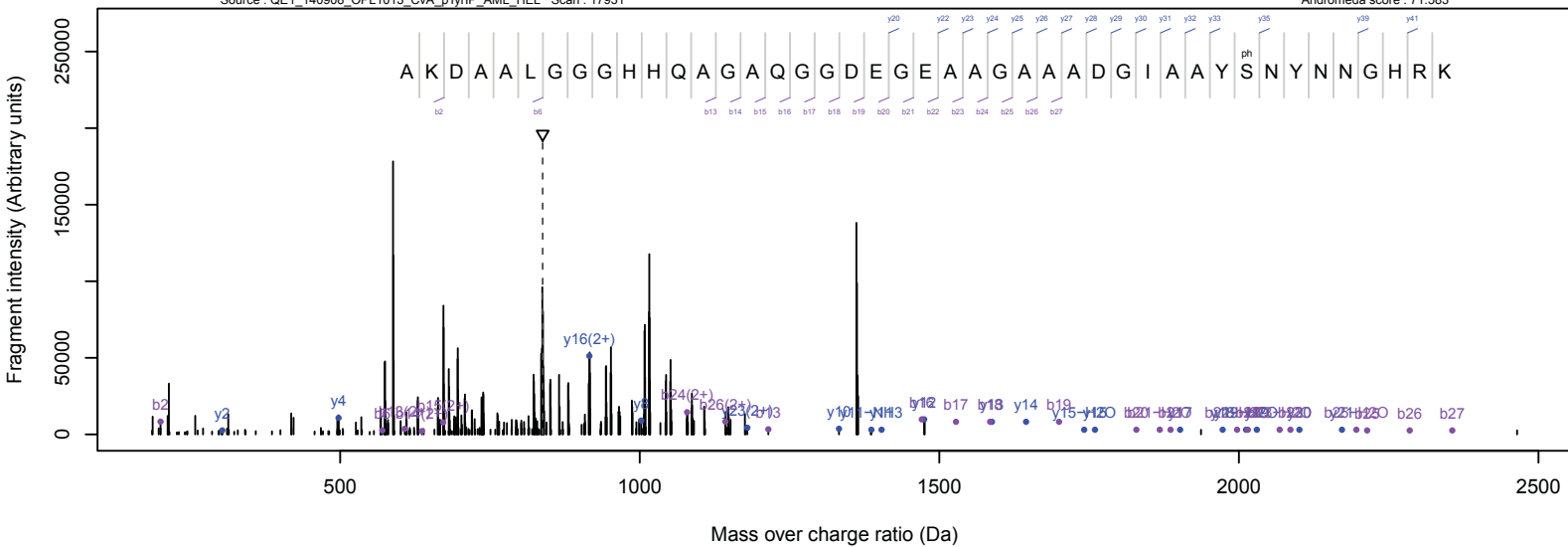
Andromeda score : 71.601



**AKDAALGGHHQAGAQQGDEGEAAGAAADGIAAYS(ph)NYNNGHRK-5+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 17931

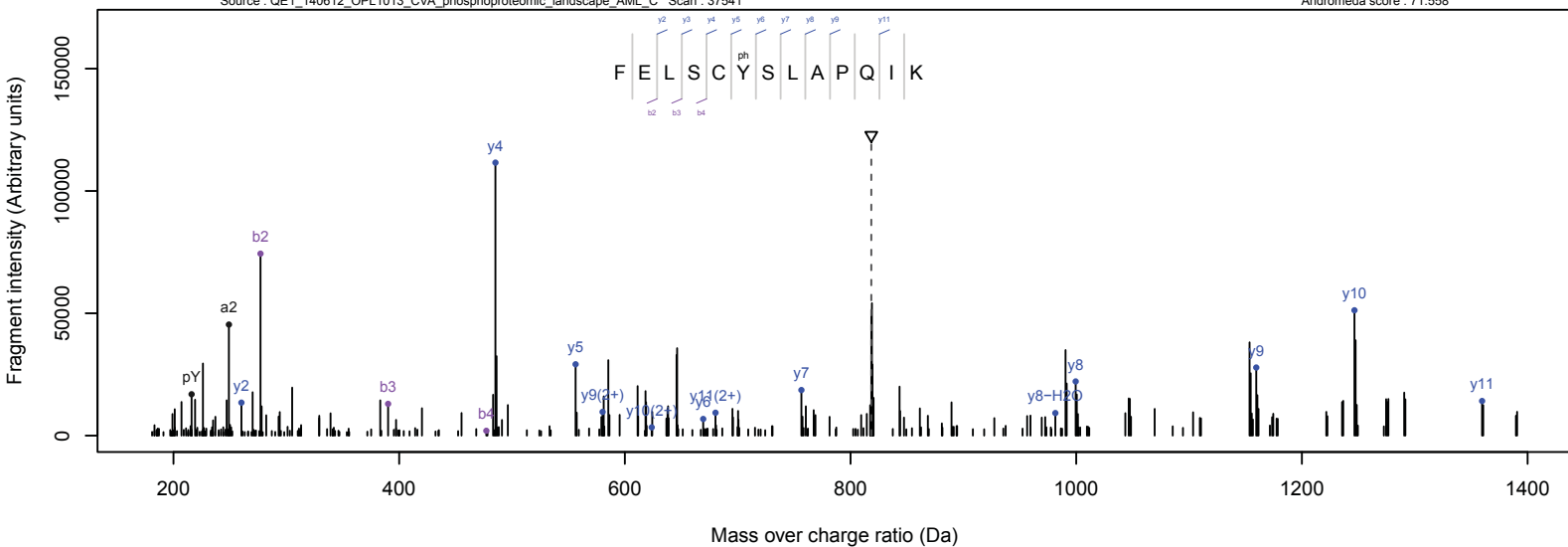
Andromeda score : 71.583



FELSCY(ph)SLAPQIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37541

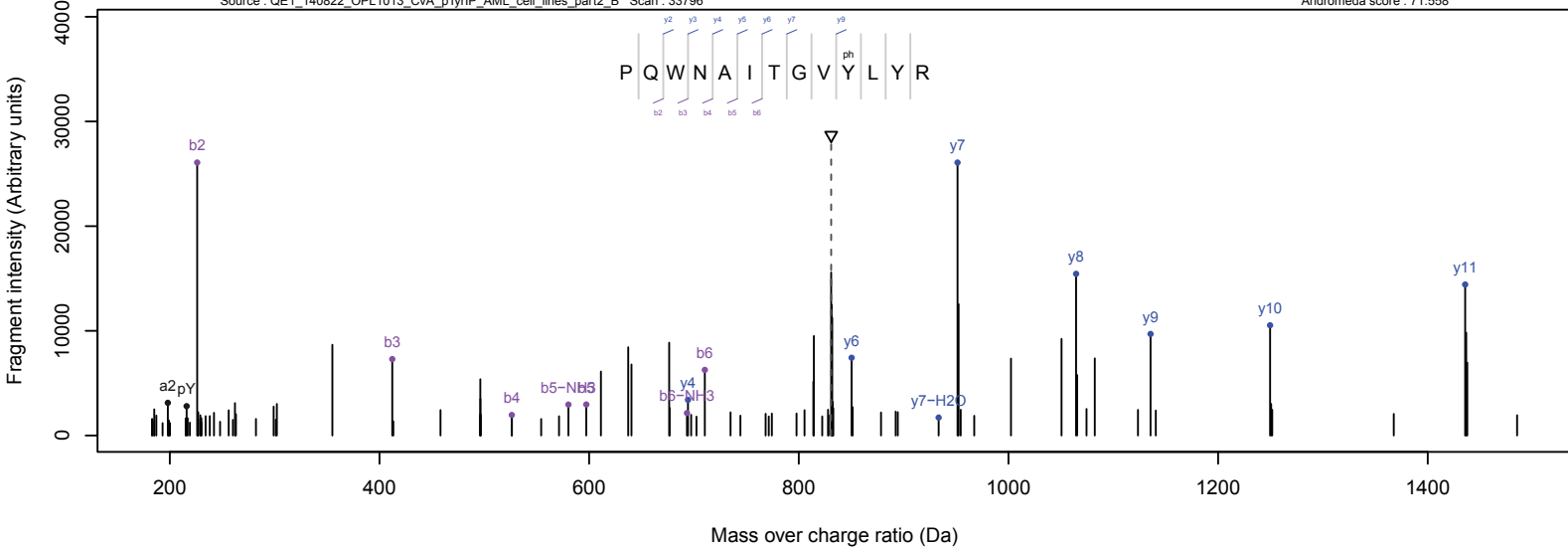
Andromeda score : 71.558



PQWNAITGVY(ph)LYR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 33796

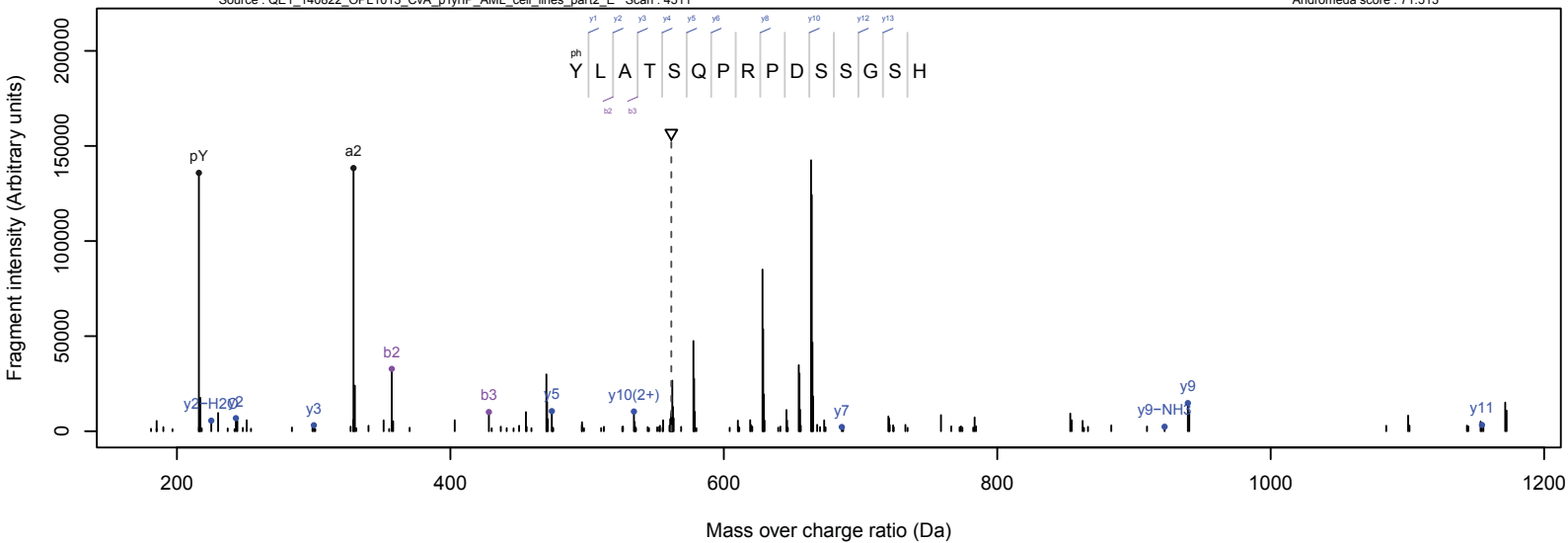
Andromeda score : 71.558



Y(ph)LATSQRPDSSGSH-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 4511

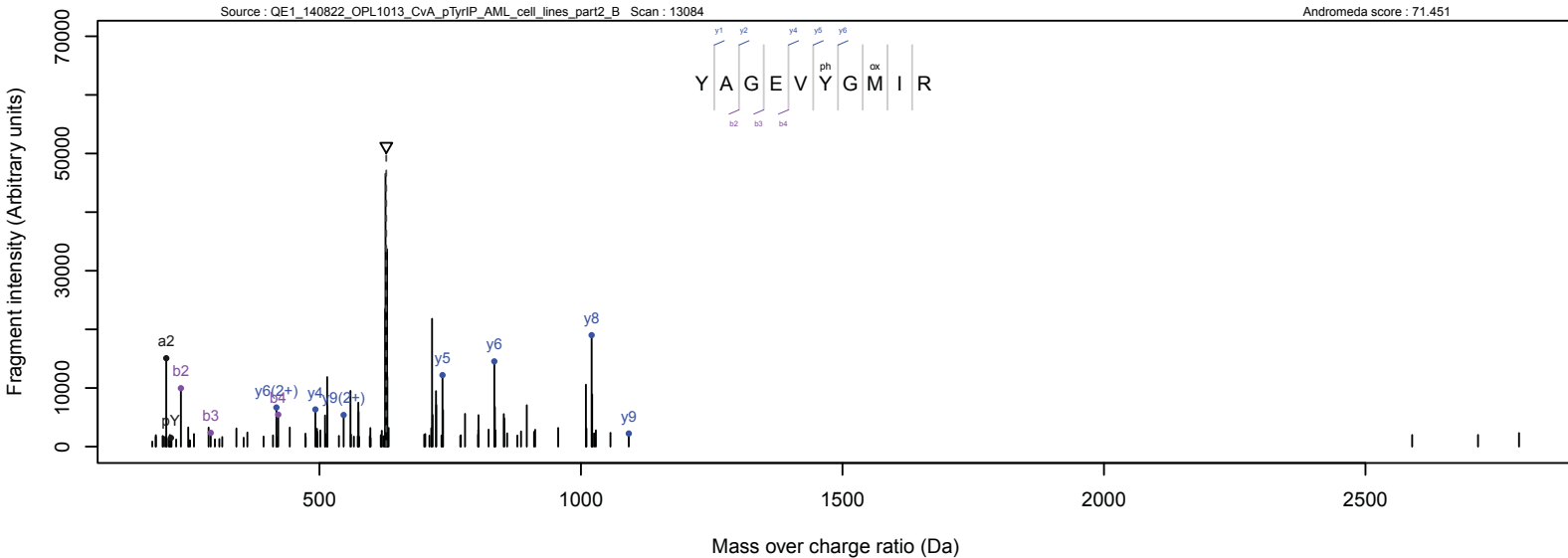
Andromeda score : 71.513



YAGEVY(ph)GM(ox)IR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 13084

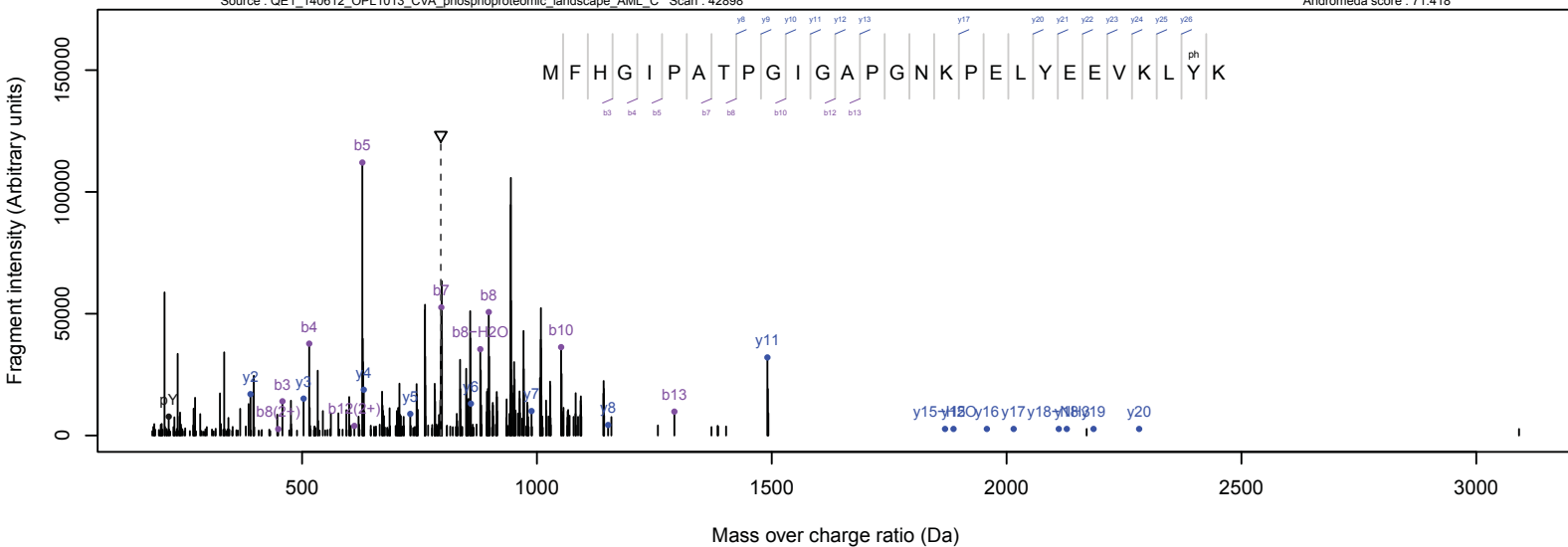
Andromeda score : 71.451



(ac)MFHGIPATPGIGAPGNKPELYEEVKLY(ph)K-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42898

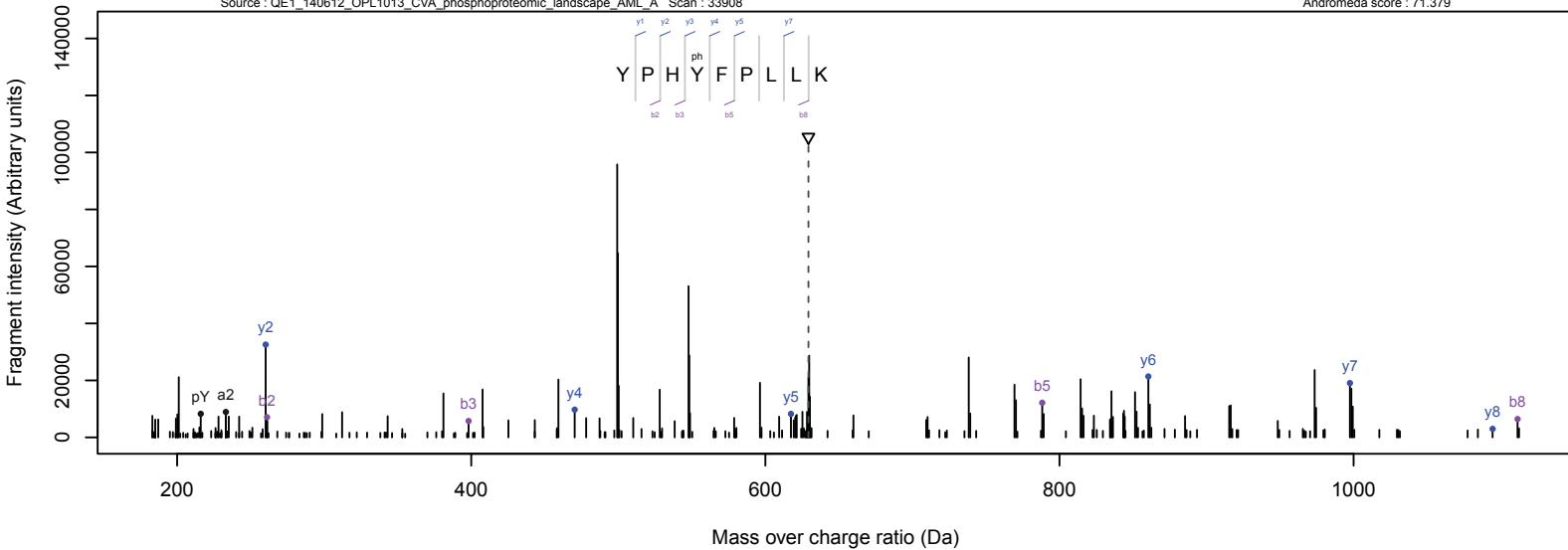
Andromeda score : 71.418



YPHY(ph)FLLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33908

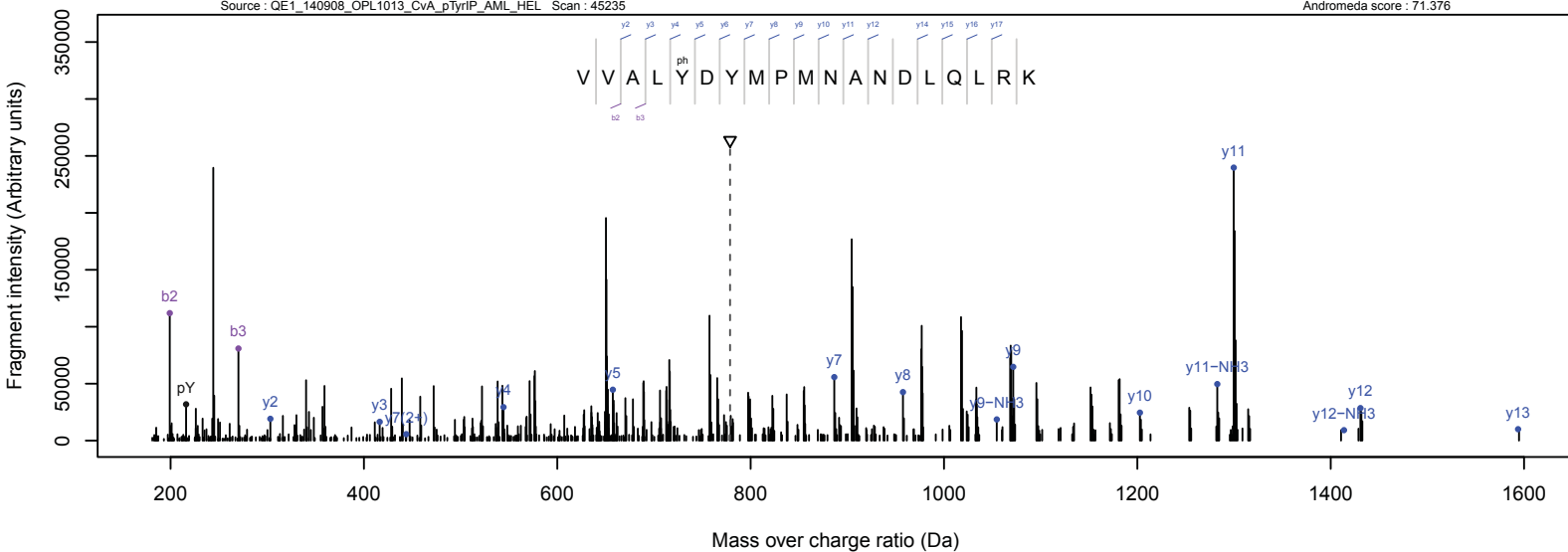
Andromeda score : 71.379



VVALY(ph)DYMPMNANDLQLRK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 45235

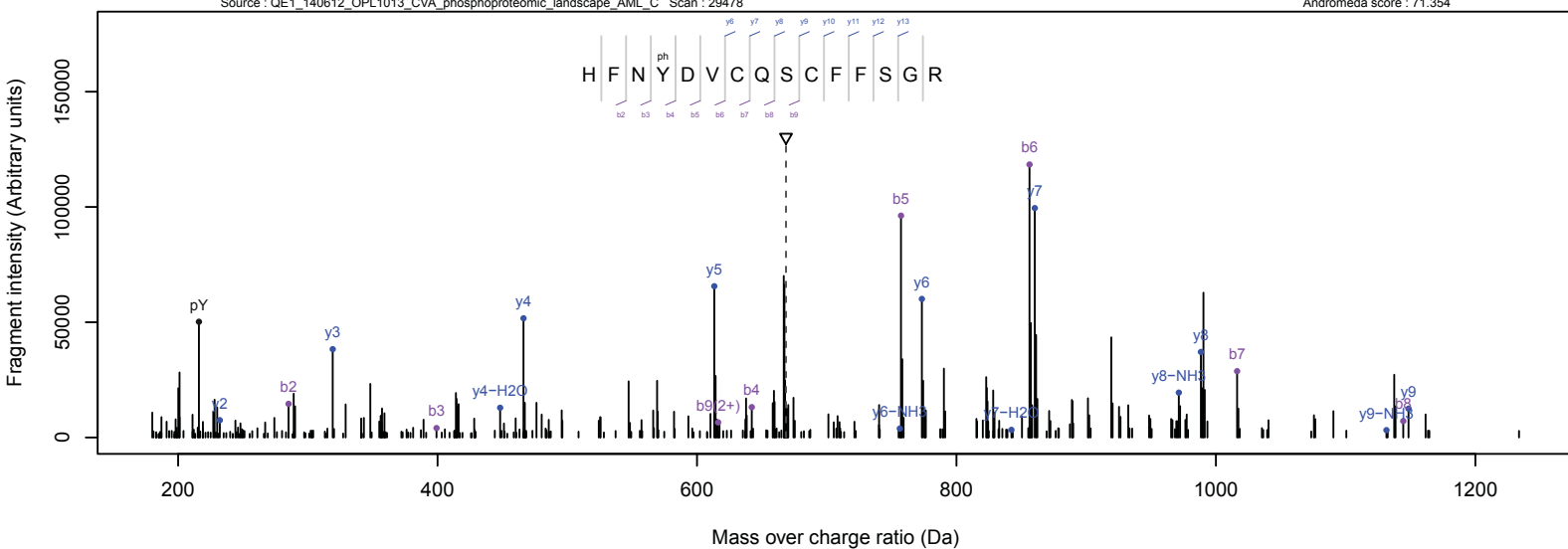
Andromeda score : 71.376



HFNY(ph)DVCQSCFFSGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29478

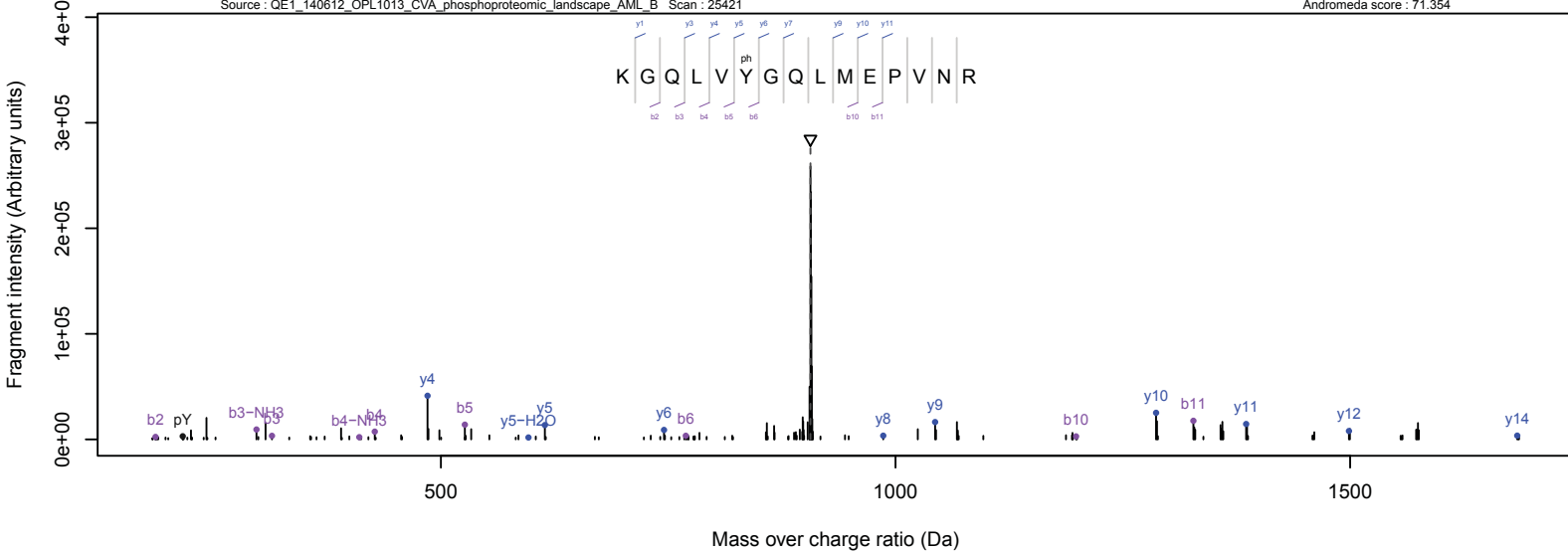
Andromeda score : 71.354



KGQLVY(ph)GQLMEPVNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25421

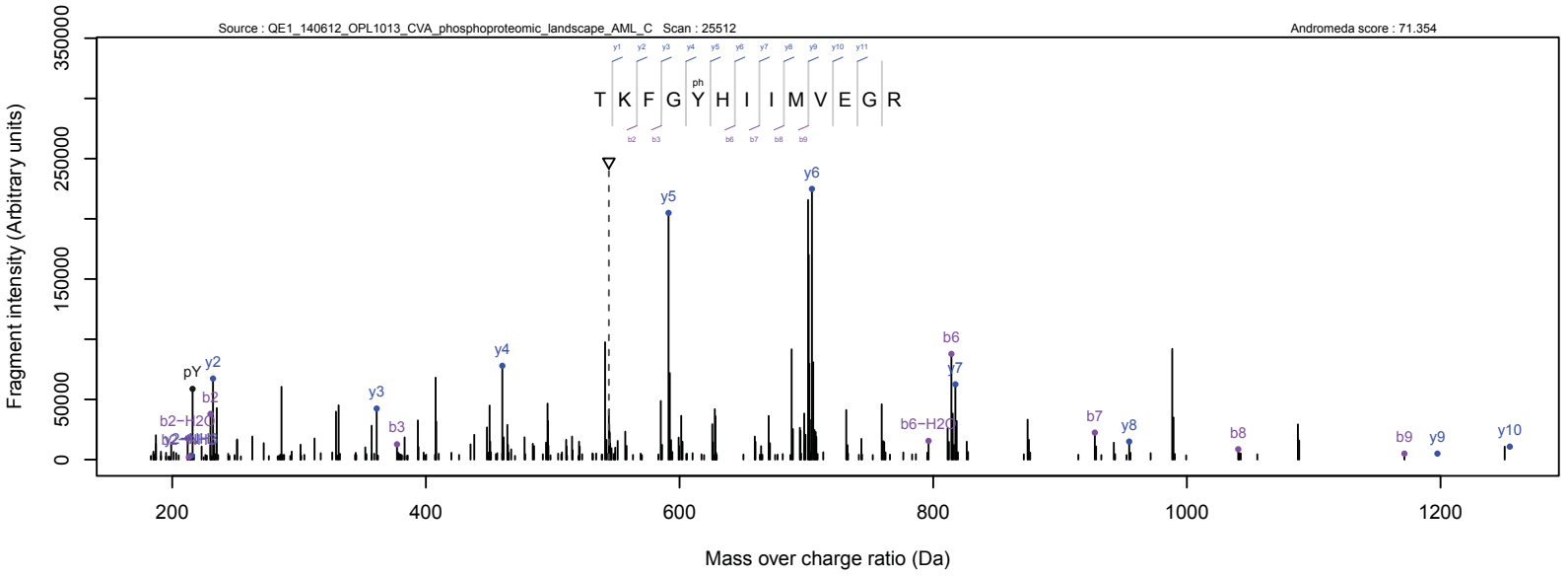
Andromeda score : 71.354



**TKFGY(ph)HIMVEGR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25512

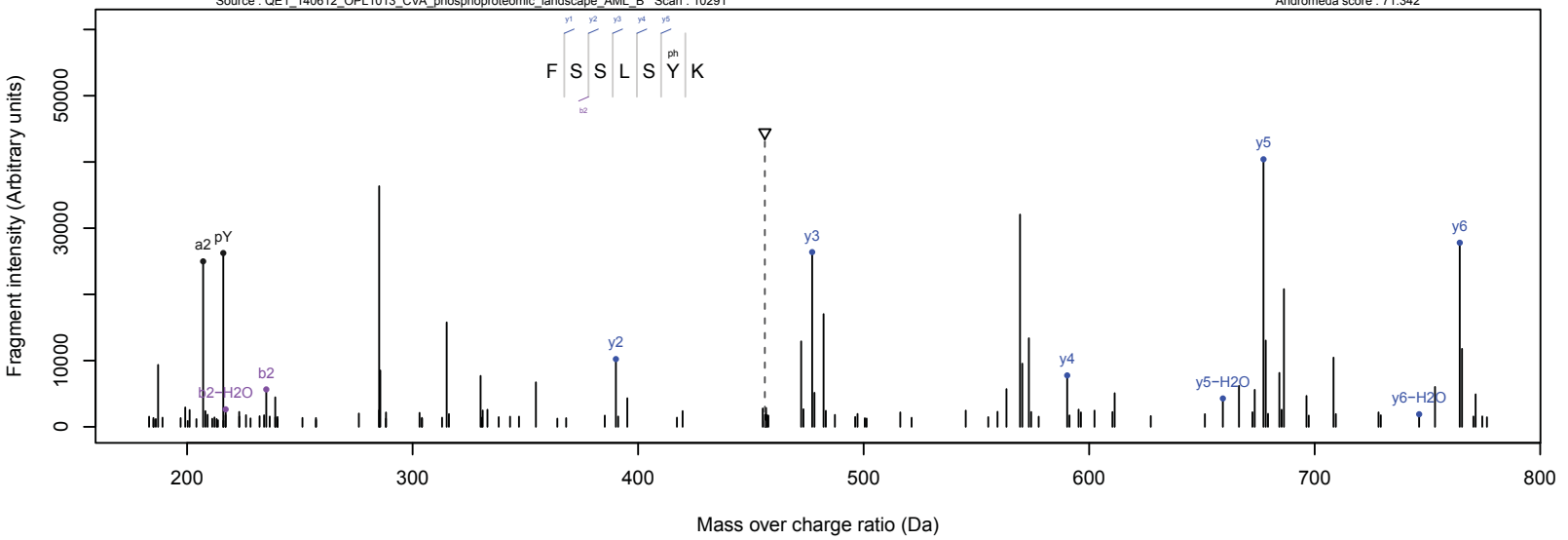
Andromeda score : 71.354



**FSSLSY(ph)K-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10291

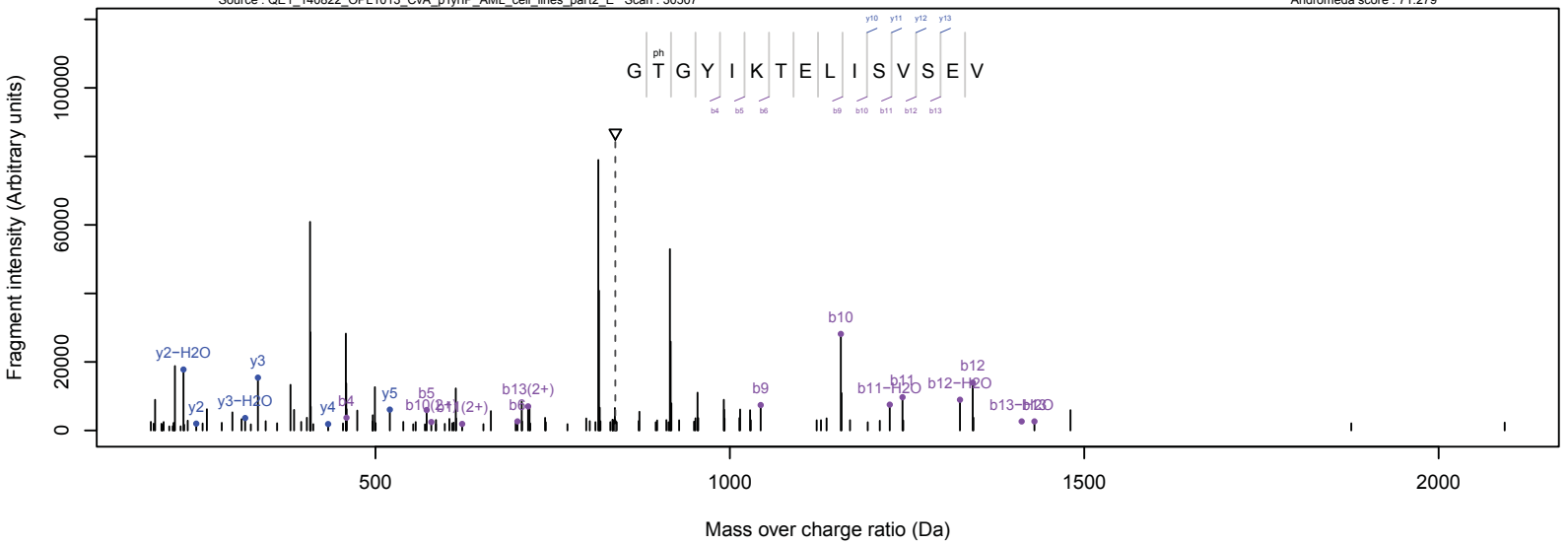
Andromeda score : 71.342



**GT(ph)GYIKTELISVSEV-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 30507

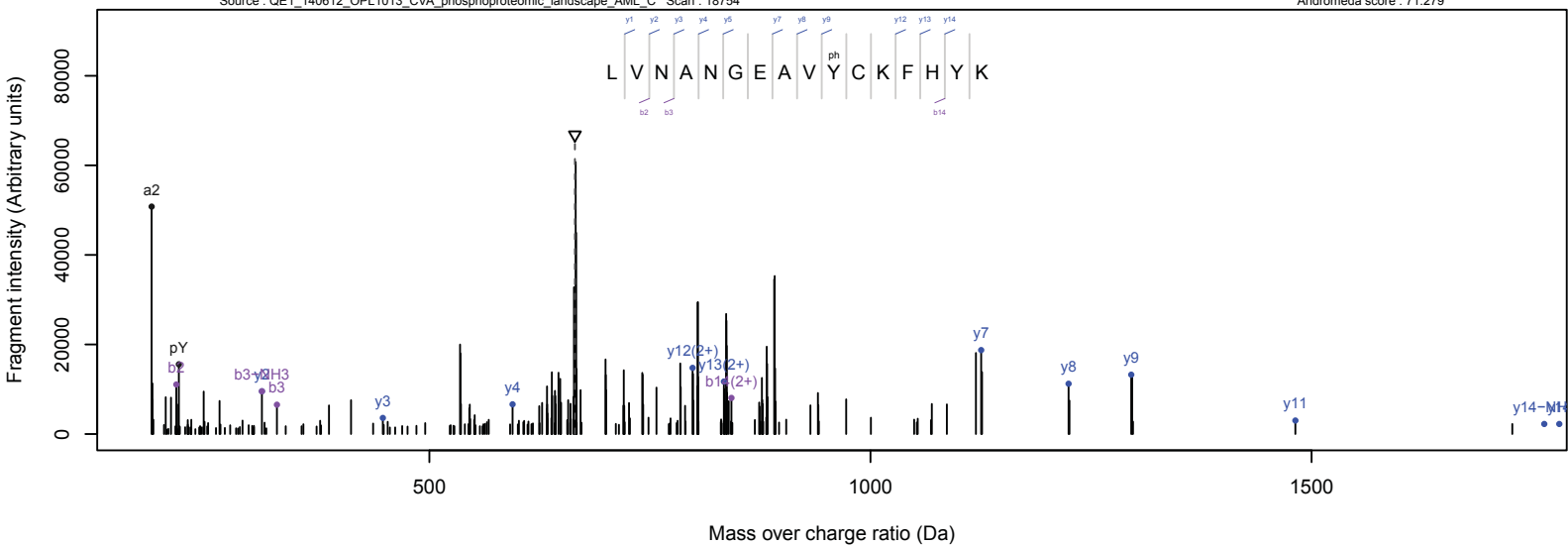
Andromeda score : 71.279



LVNANGEAVY(ph)CKFHVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18754

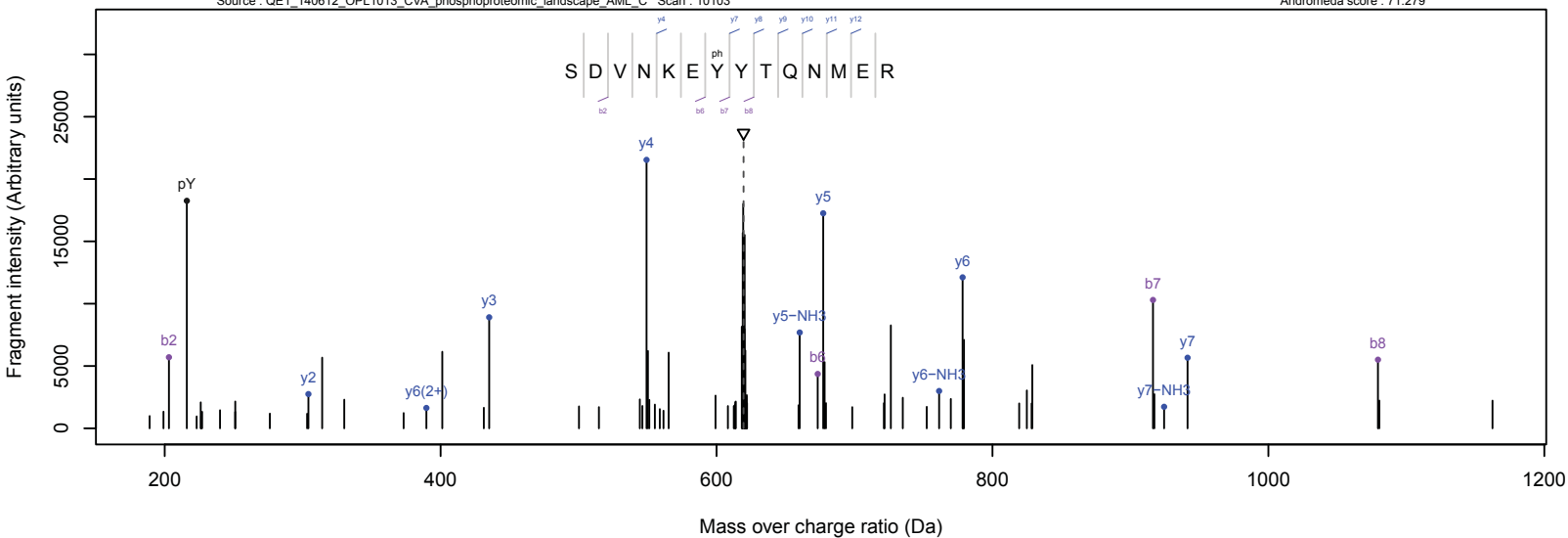
Andromeda score : 71.279



SDVNKEY(ph)YTQNMER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10103

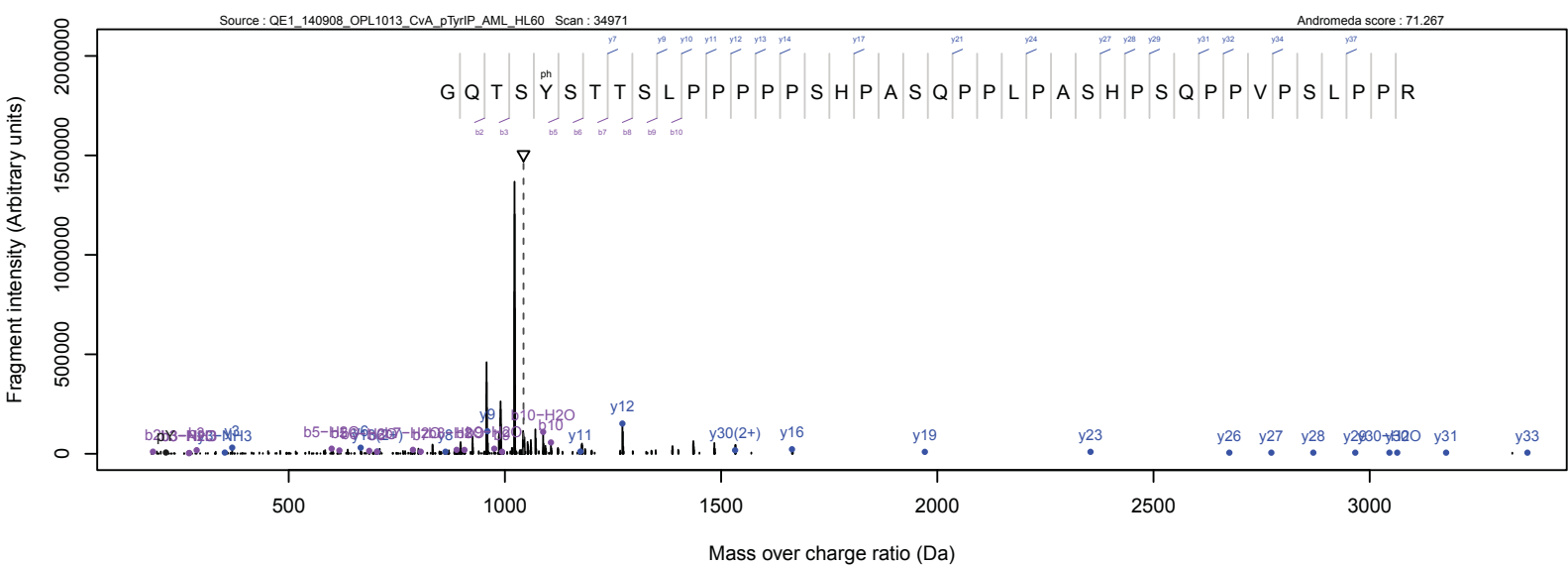
Andromeda score : 71.279



GQTSY(ph)STTSLPPPPSPASQPPLPASHPSQPPVPSLPPR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 34971

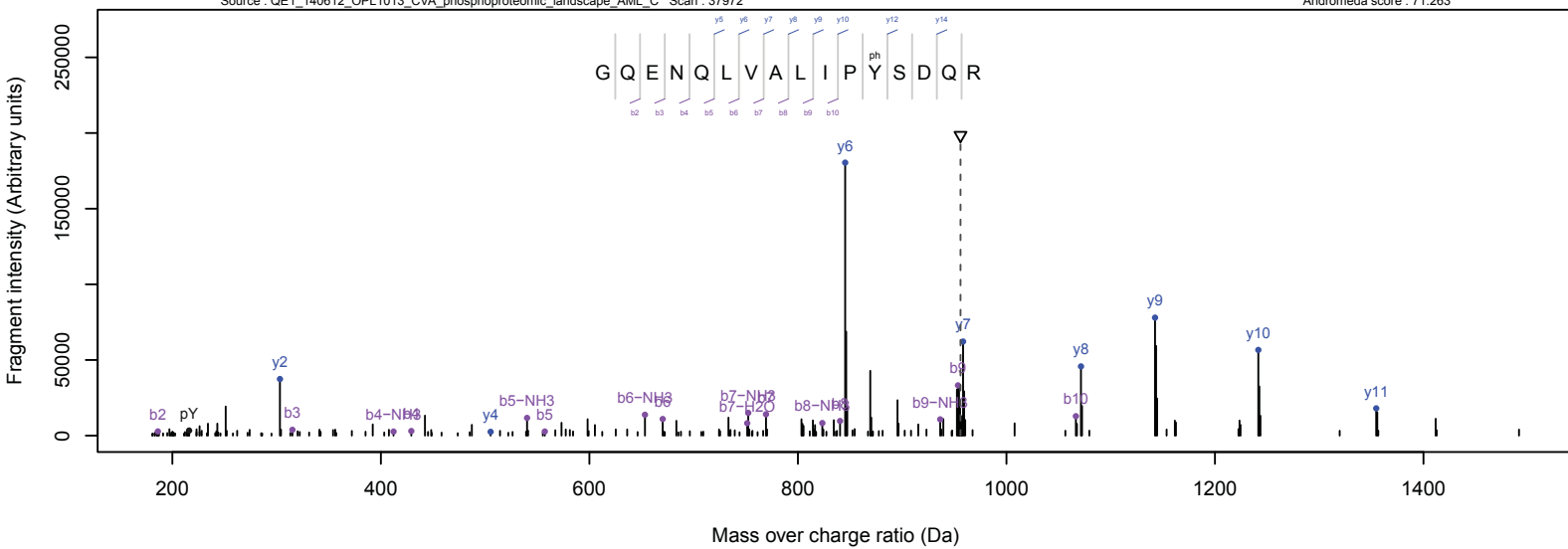
Andromeda score : 71.267



## GQENQLVIP(ph)SDQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37972

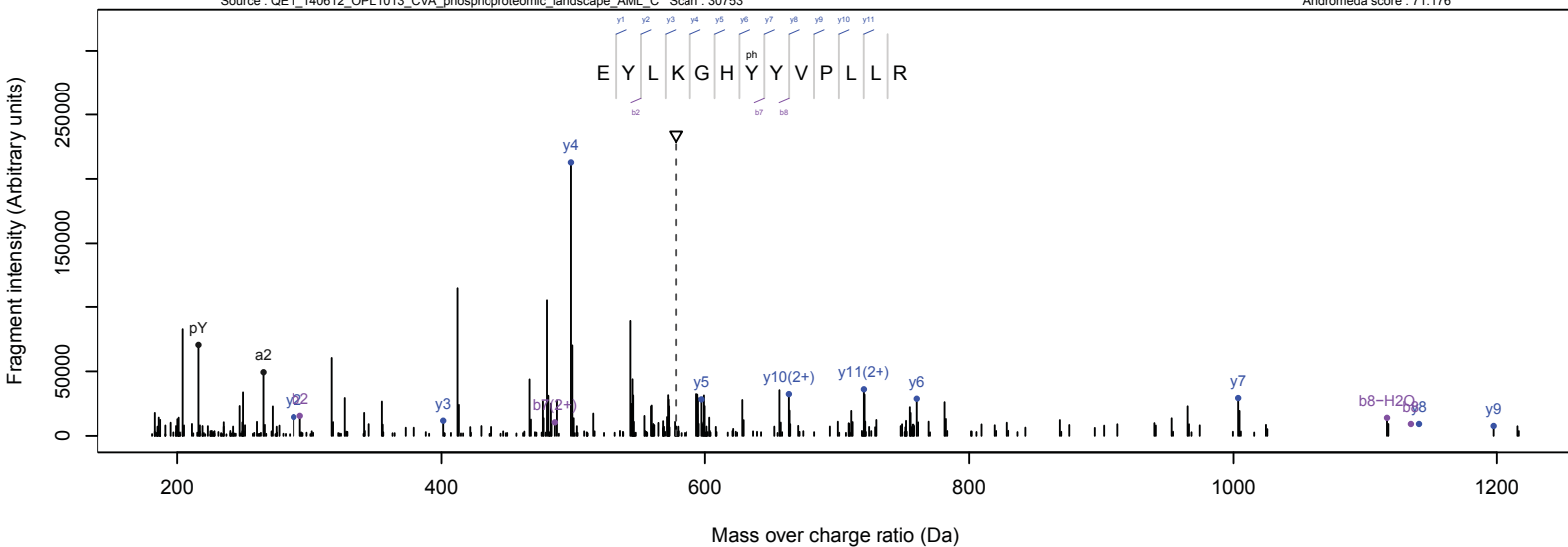
Andromeda score : 71.263



## EYLKGHY(ph)YVPLLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30753

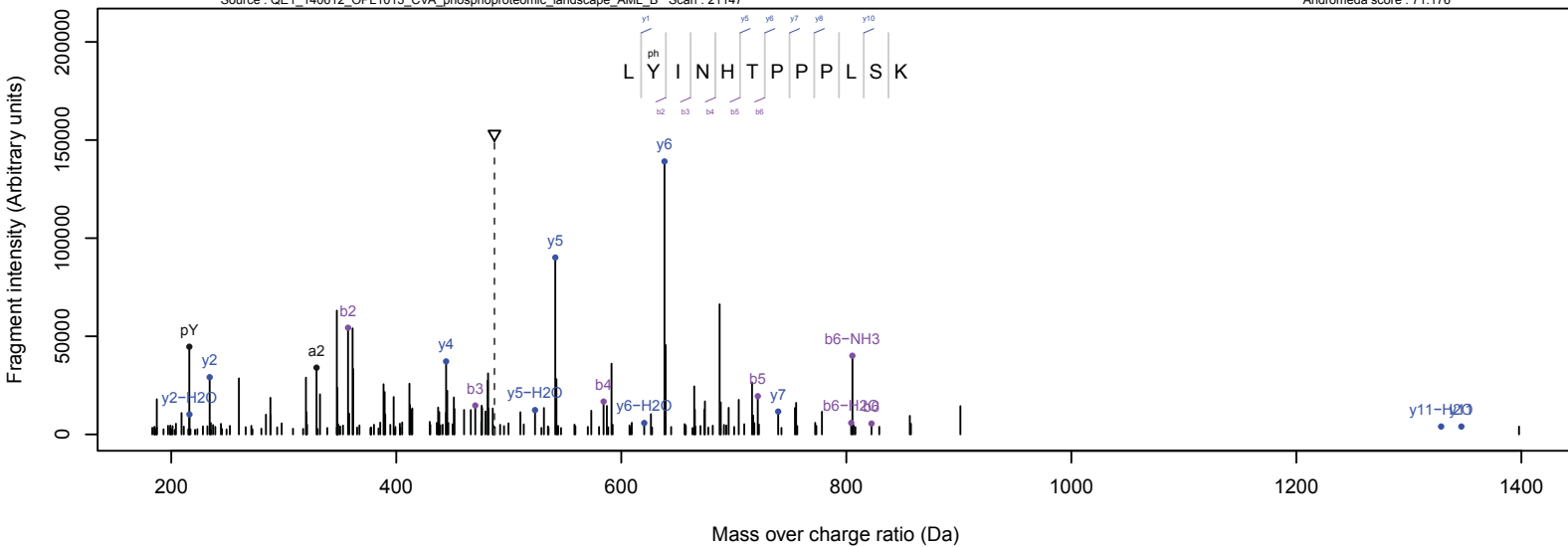
Andromeda score : 71.176



## LY(ph)INHTPPPLSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21147

Andromeda score : 71.176

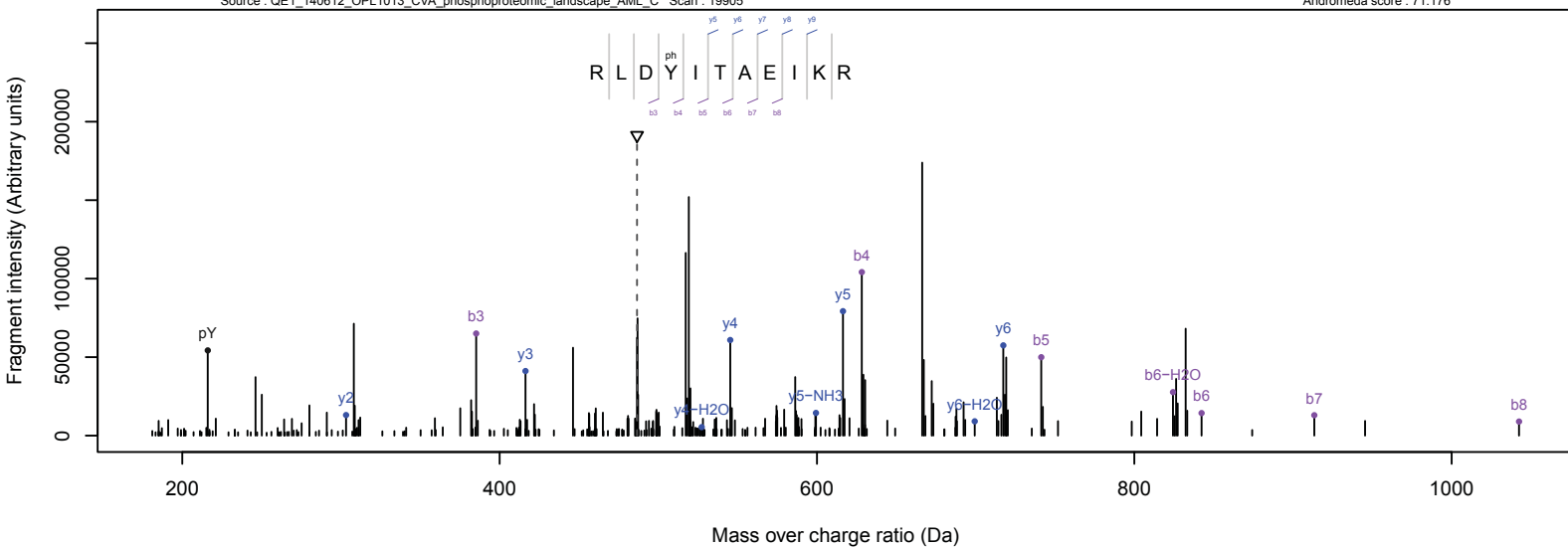




## RLDY(ph)ITAEIKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19905

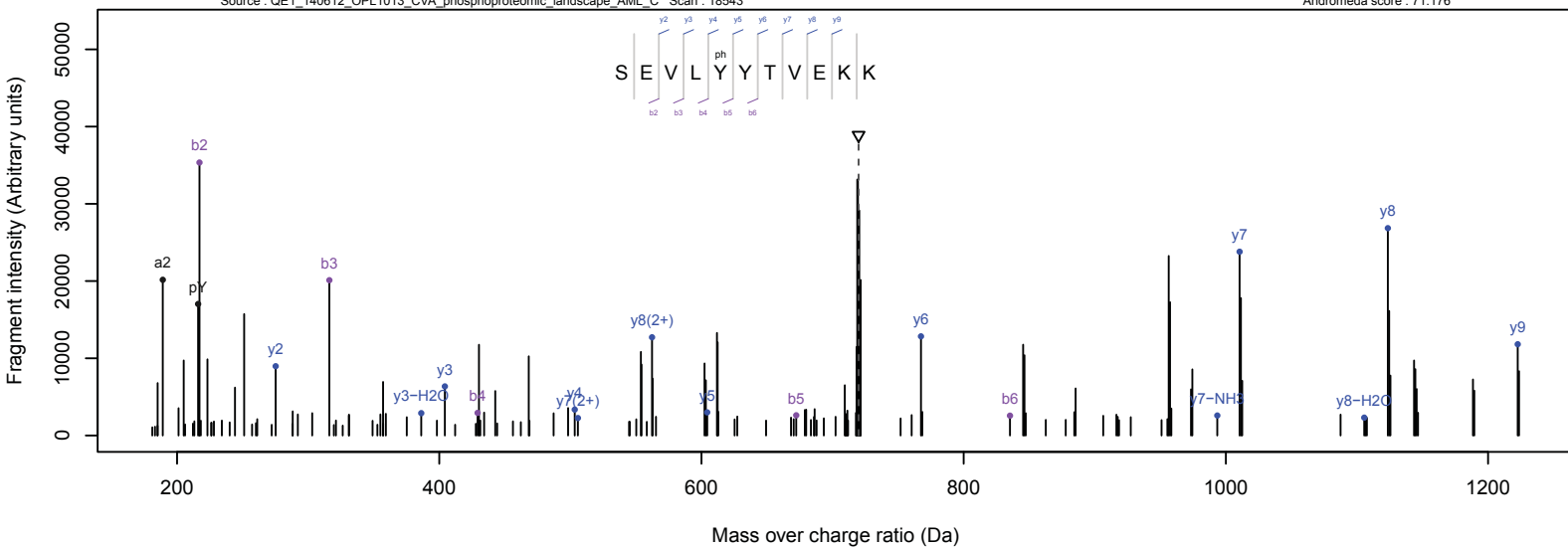
Andromeda score : 71.176



## SEVLY(ph)YTV EK K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18543

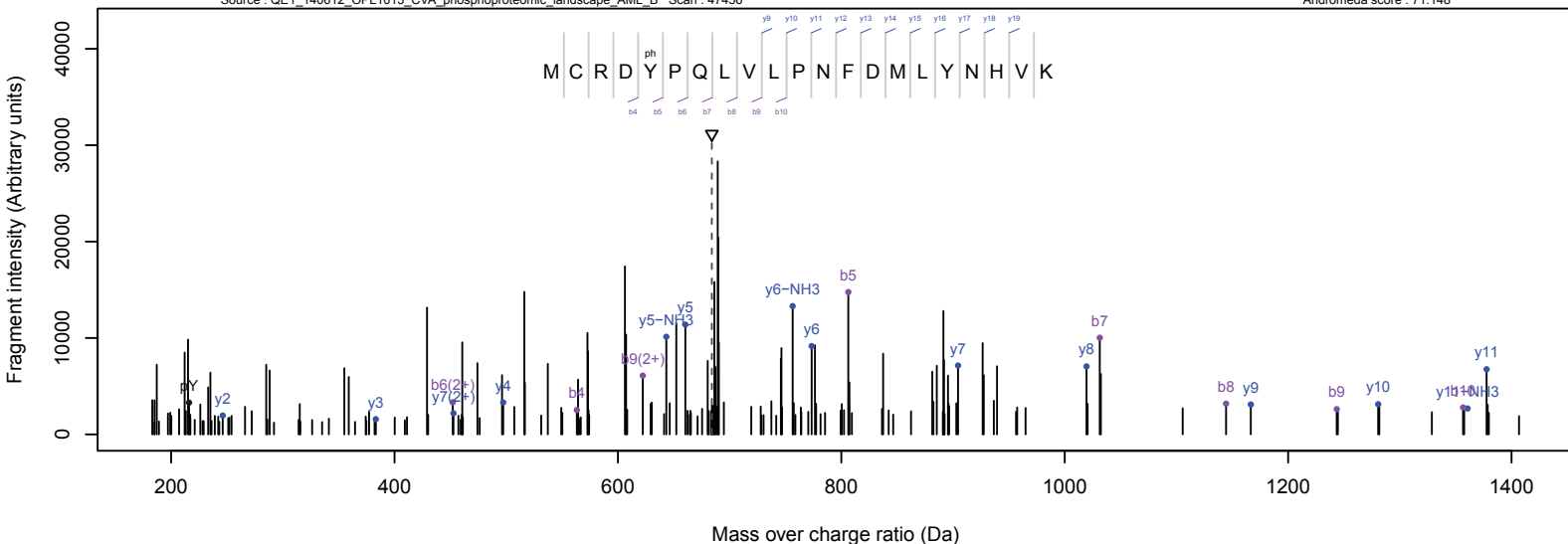
Andromeda score : 71.176



## MCRDY(ph)PQLVLPNF DMLYNHVK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47456

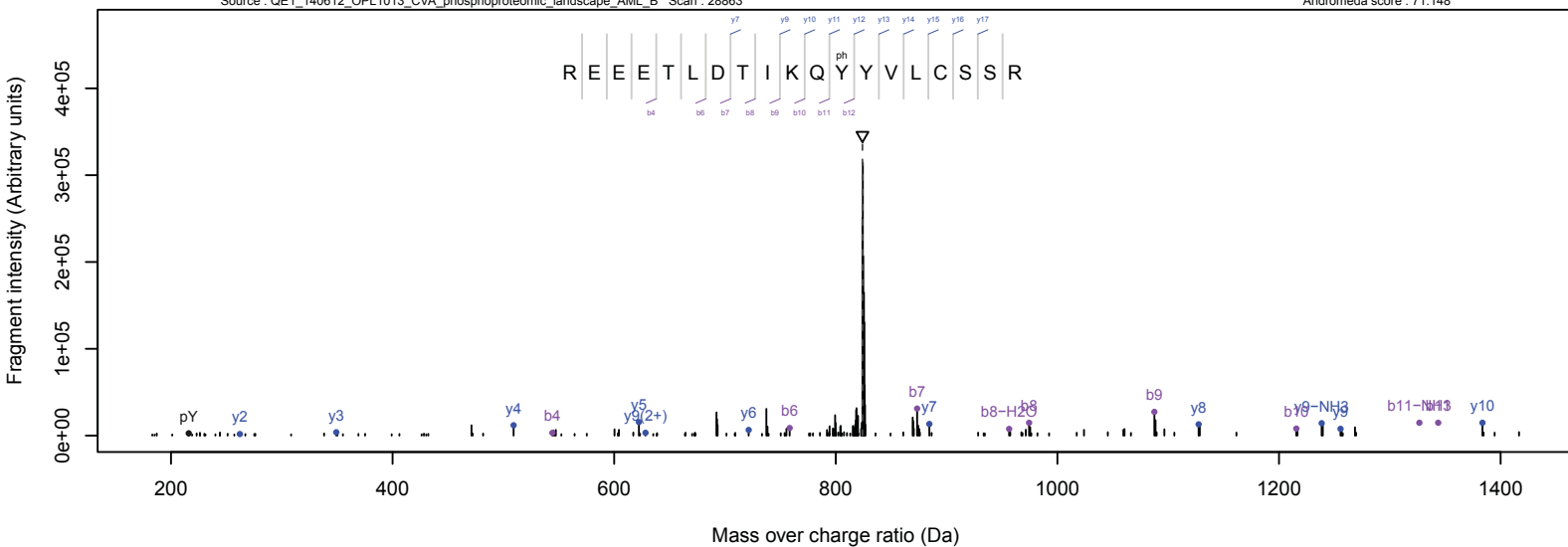
Andromeda score : 71.148



REEETLDTIKQ(ph)YVLCSSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28863

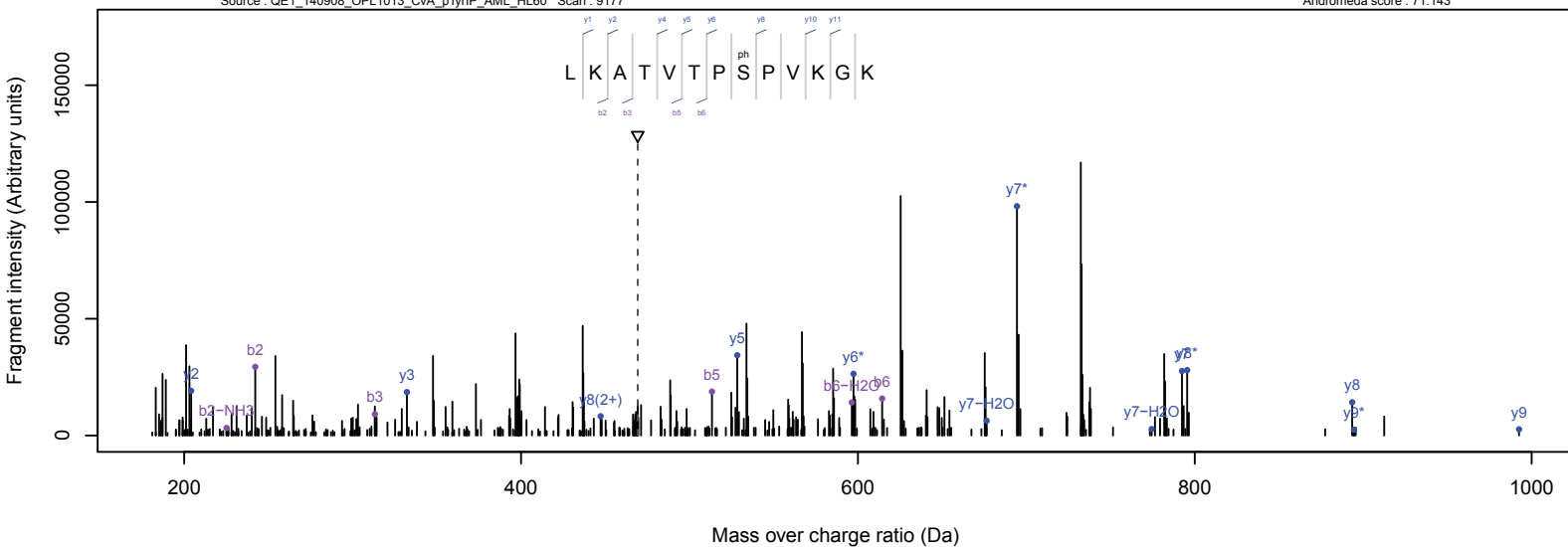
Andromeda score : 71.148



LKATVTPS(ph)PVK GK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 9177

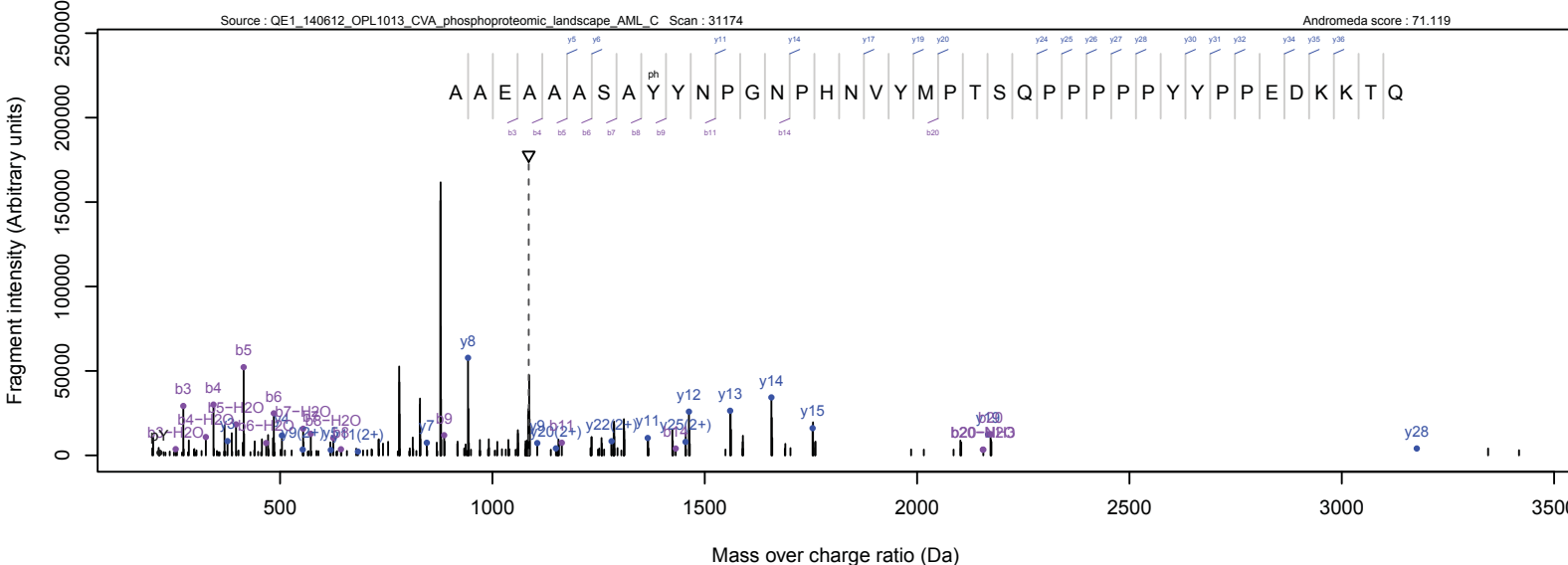
Andromeda score : 71.143



AAEAAASAY(ph)YNGNPHNVYMPYPPPPYPPEDKKTQ-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31174

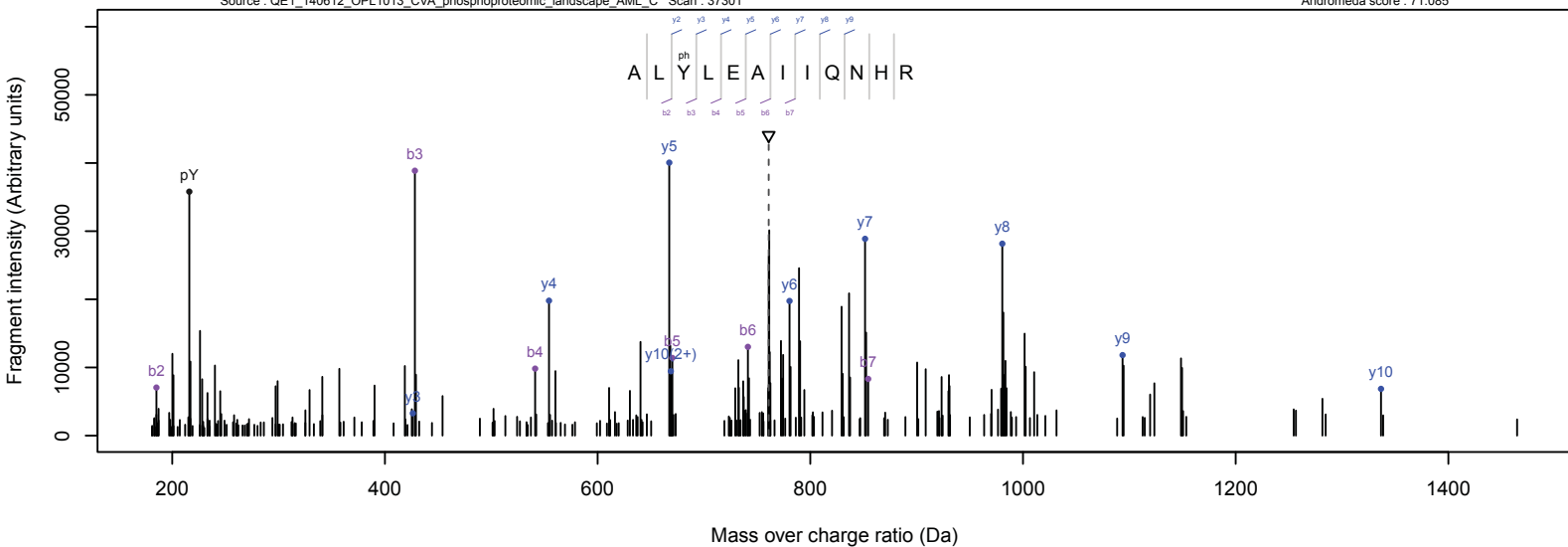
Andromeda score : 71.119



ALY(ph)LEAIQNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37301

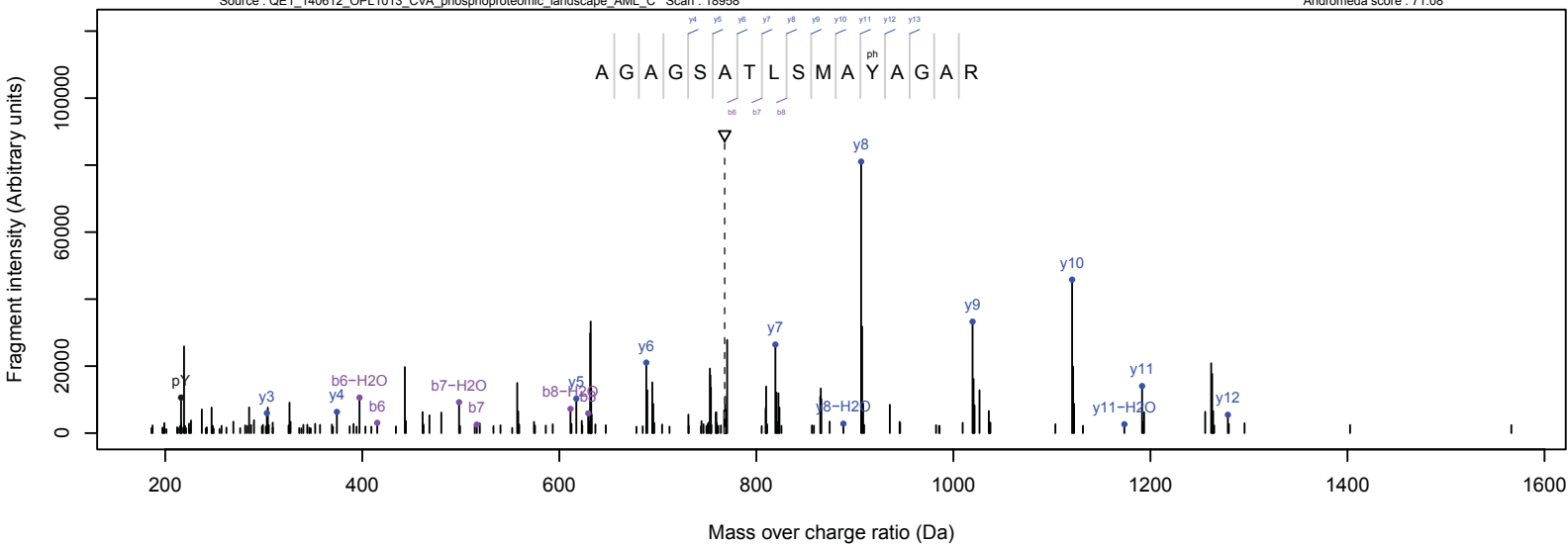
Andromeda score : 71.085



AGAGSATLSMAY(ph)AGAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18958

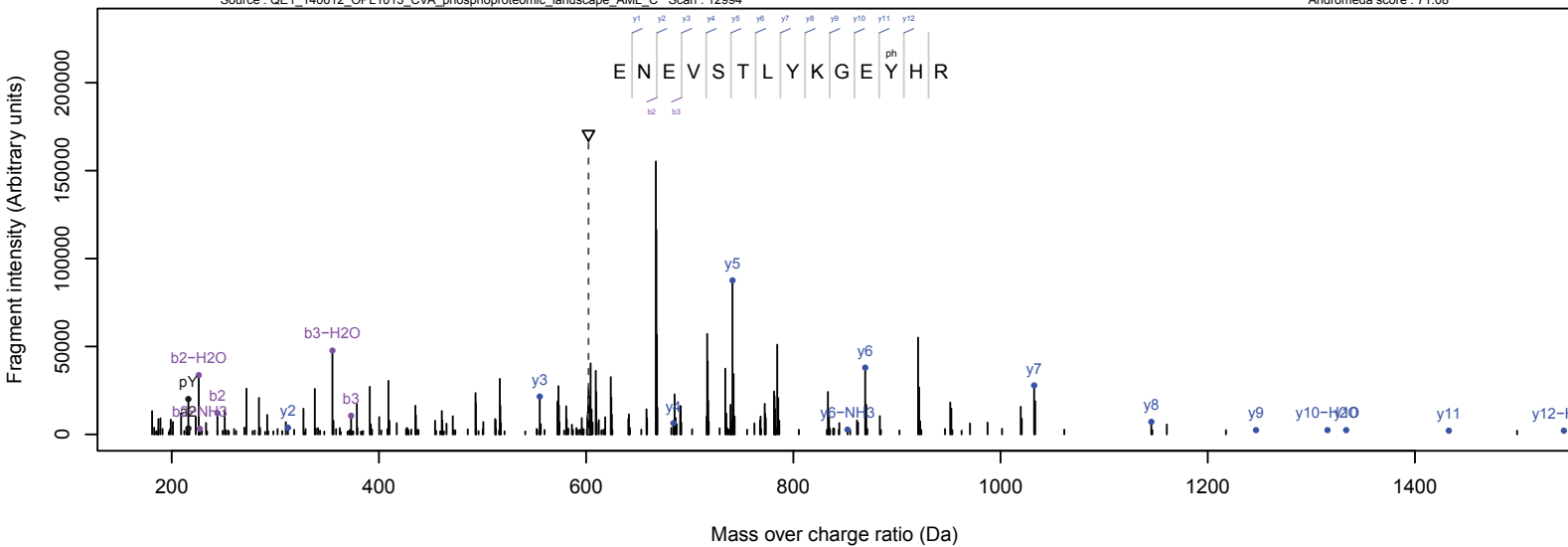
Andromeda score : 71.08



ENEVSTLYKGEY(ph)HR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12994

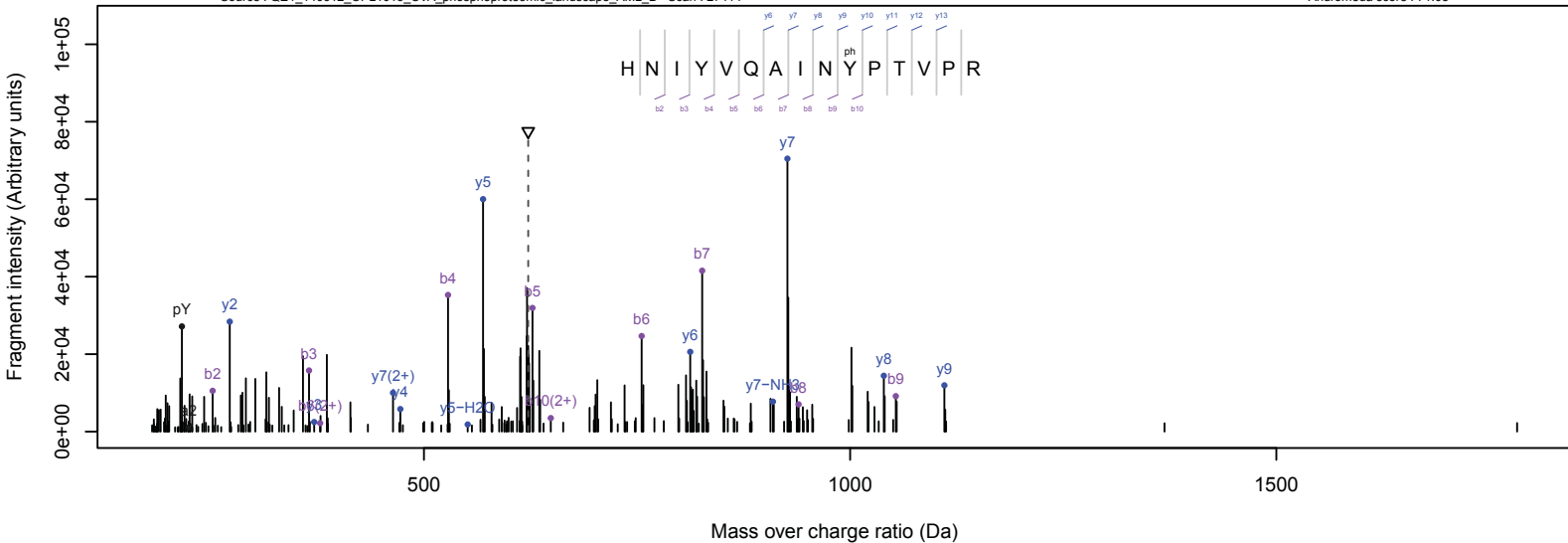
Andromeda score : 71.08



## HNIYVQAINY(ph)PTVPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27414

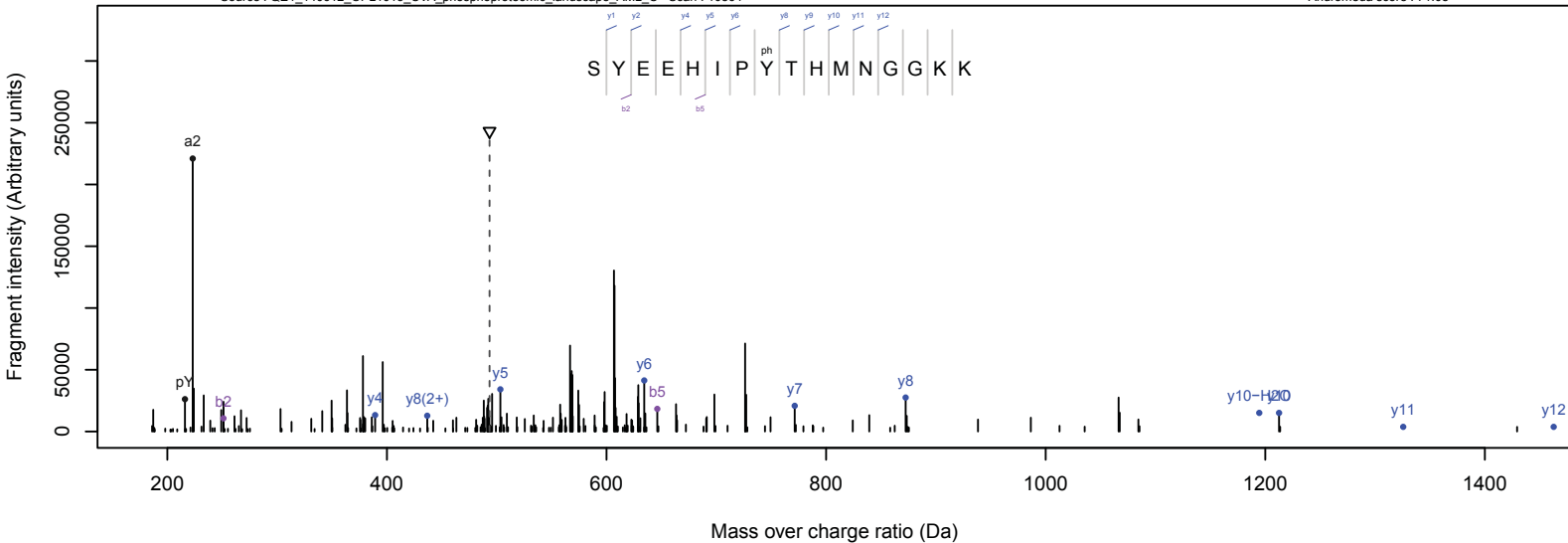
Andromeda score : 71.08



## SYEEHIPY(ph)THMNGGKK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10864

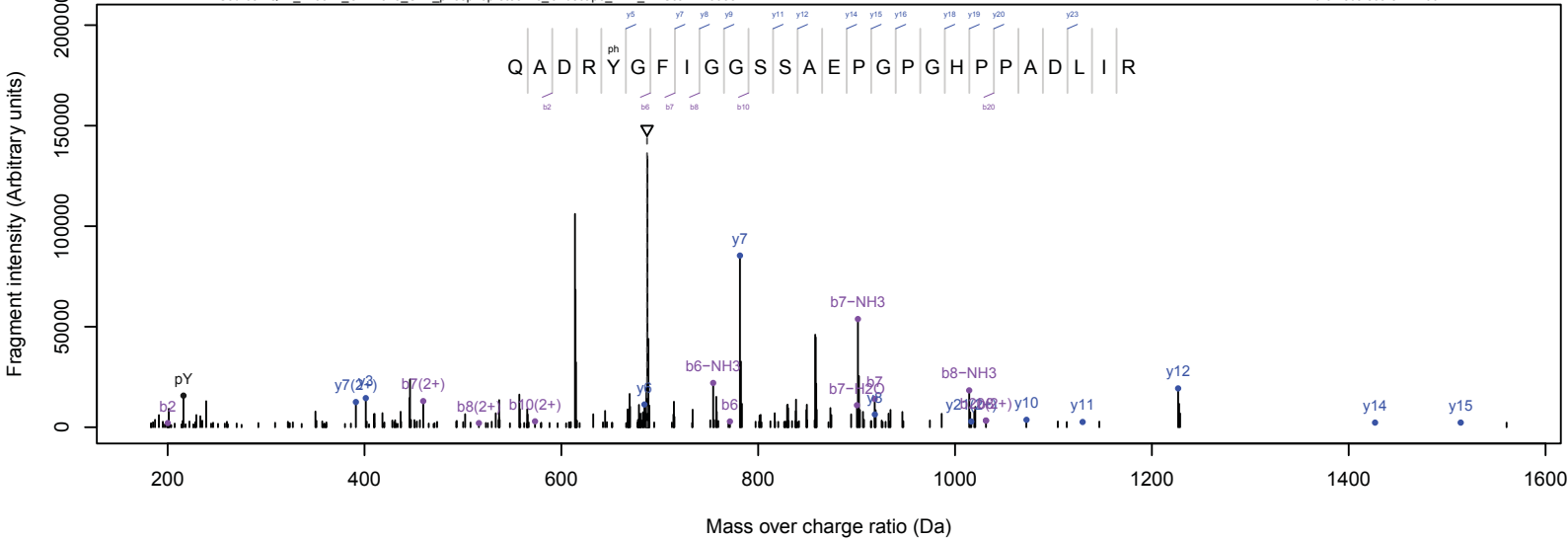
Andromeda score : 71.08



## QADRY(ph)GFIGGSSAEPGPGHPADLIR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29958

Andromeda score : 71.06

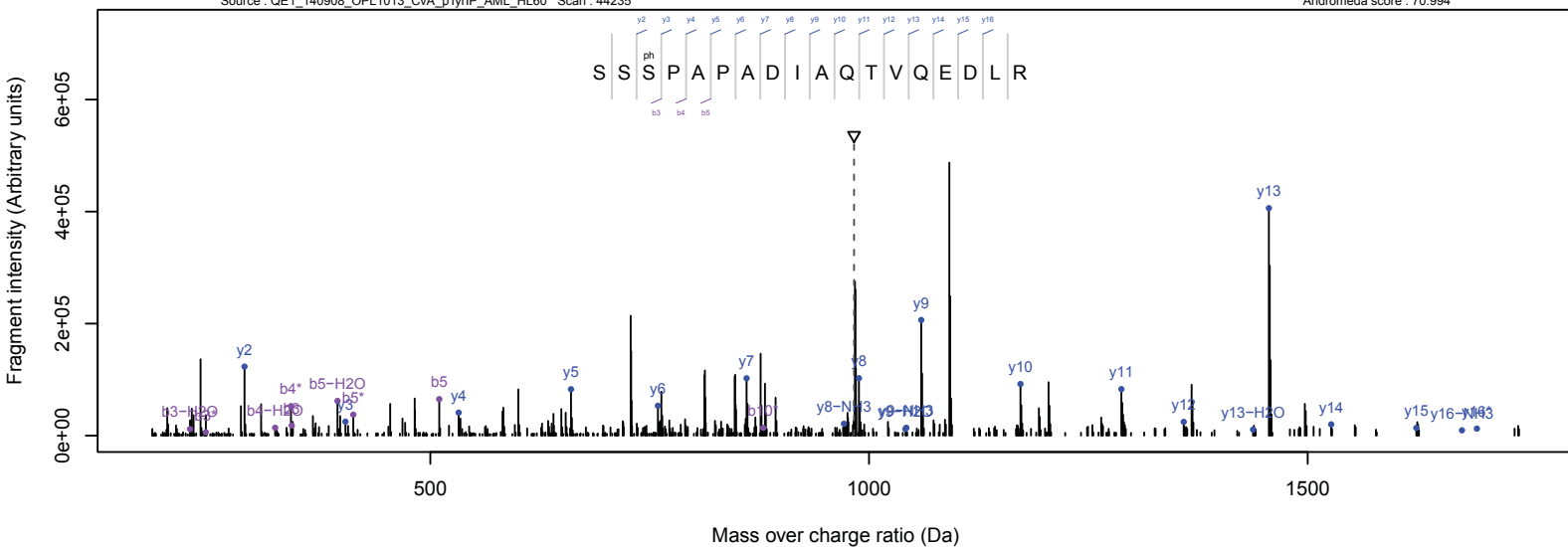




## SSS(ph)PAPADIAQTVQEDLR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 44235

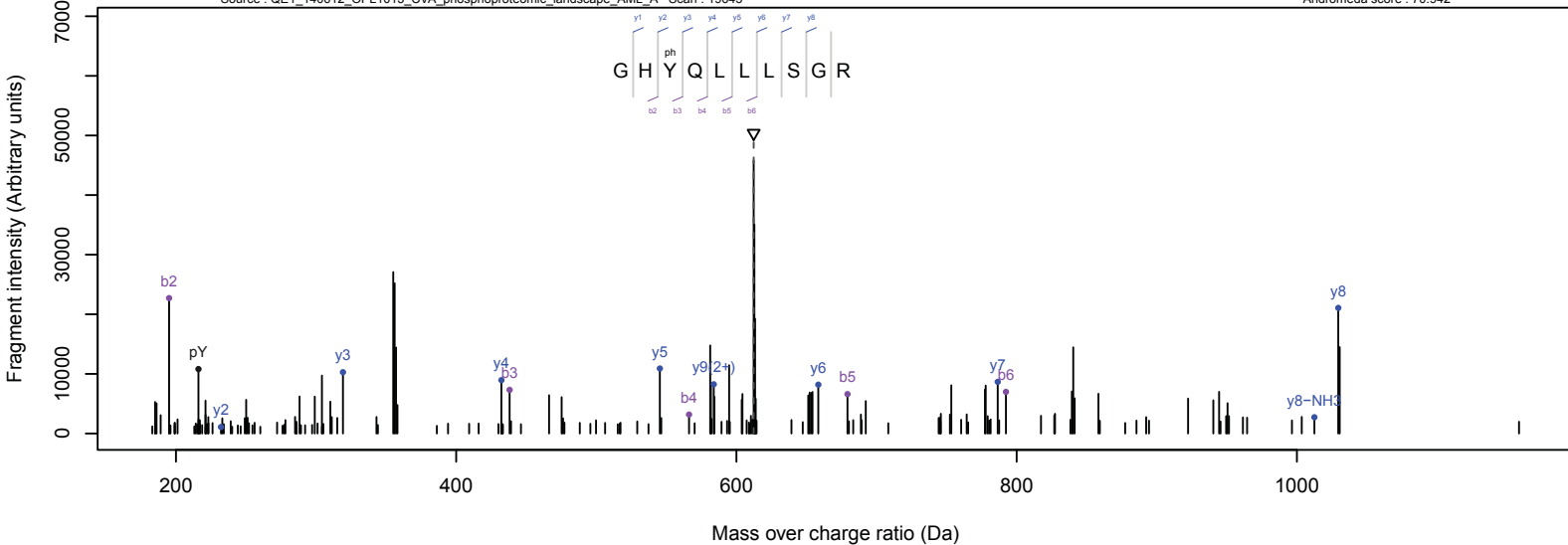
Andromeda score : 70.994



## GHY(ph)QLLLSGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 19645

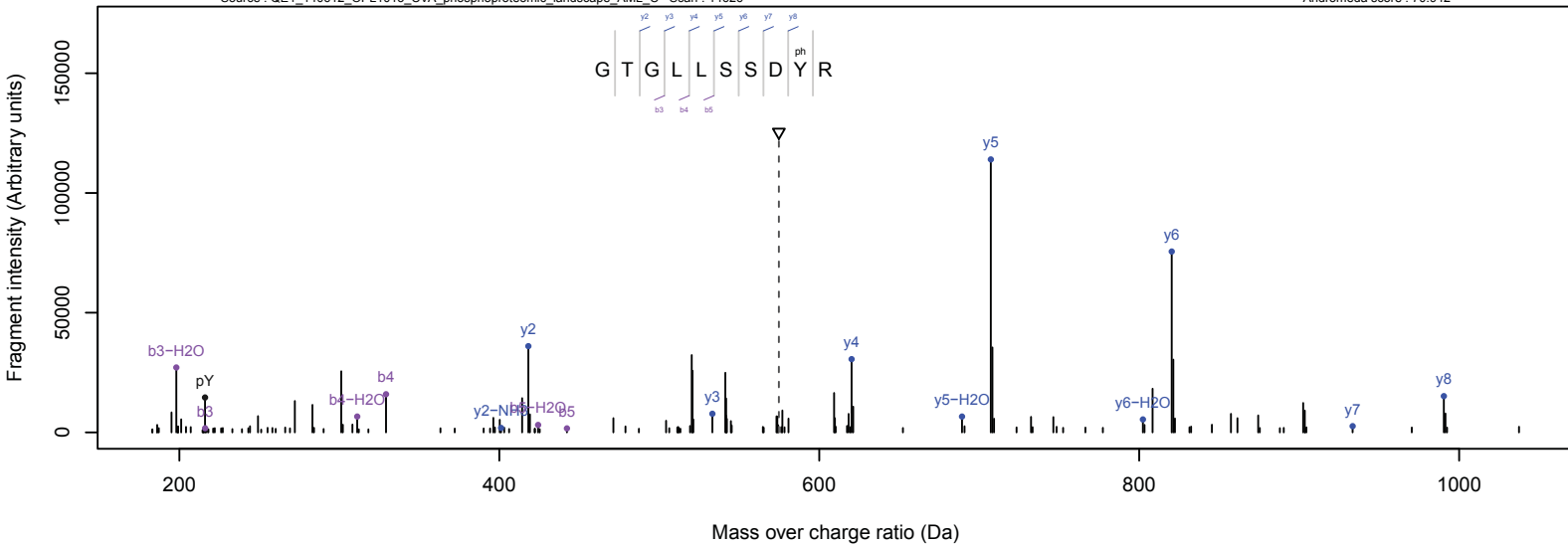
Andromeda score : 70.942



## GTGLSSDY(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14626

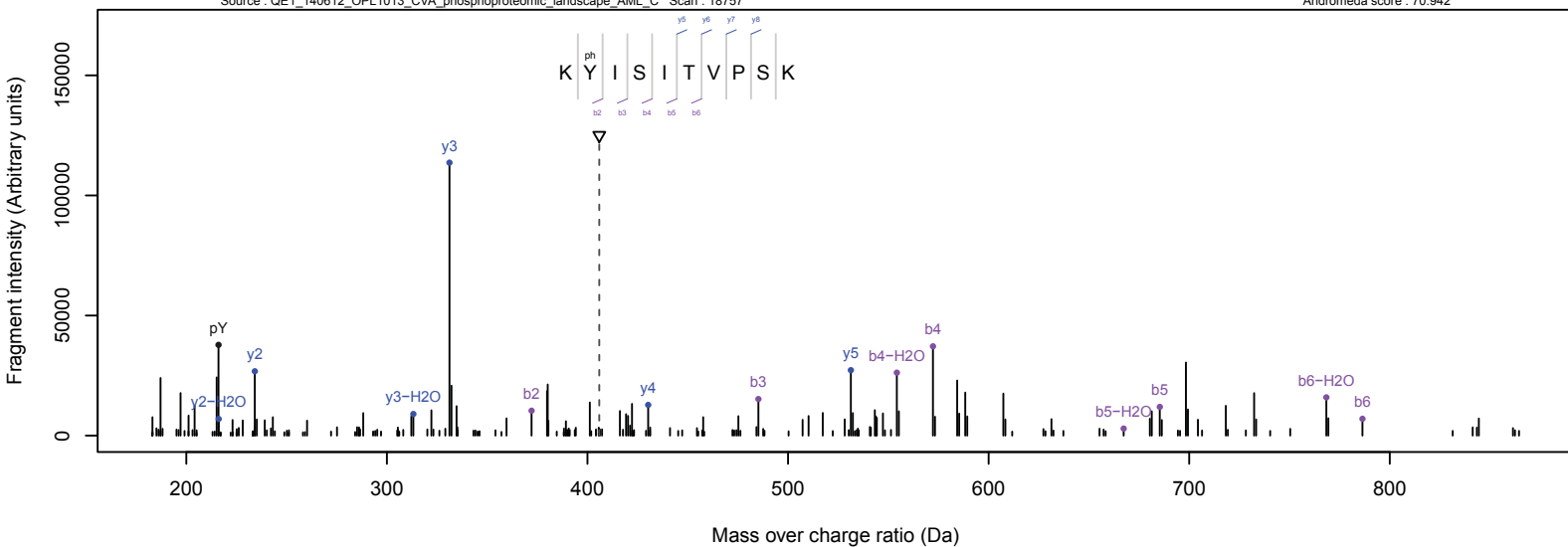
Andromeda score : 70.942



KY(ph)ISITVPSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18757

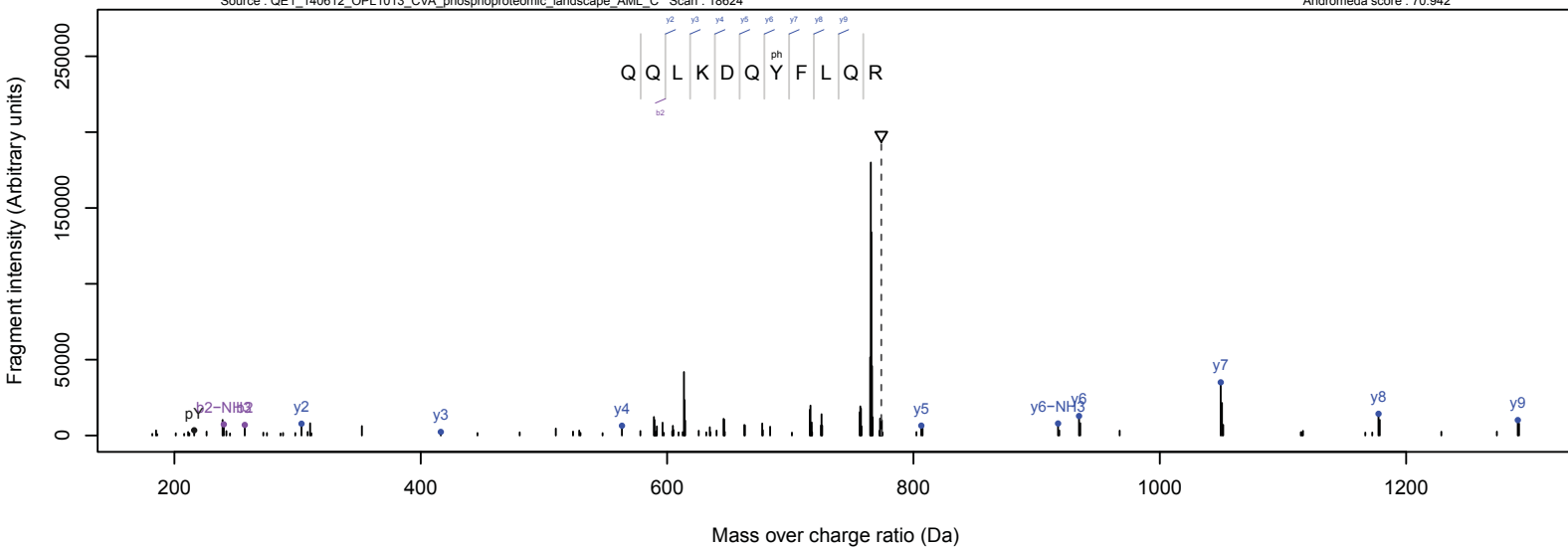
Andromeda score : 70.942



QLKDKQY(ph)FLQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18624

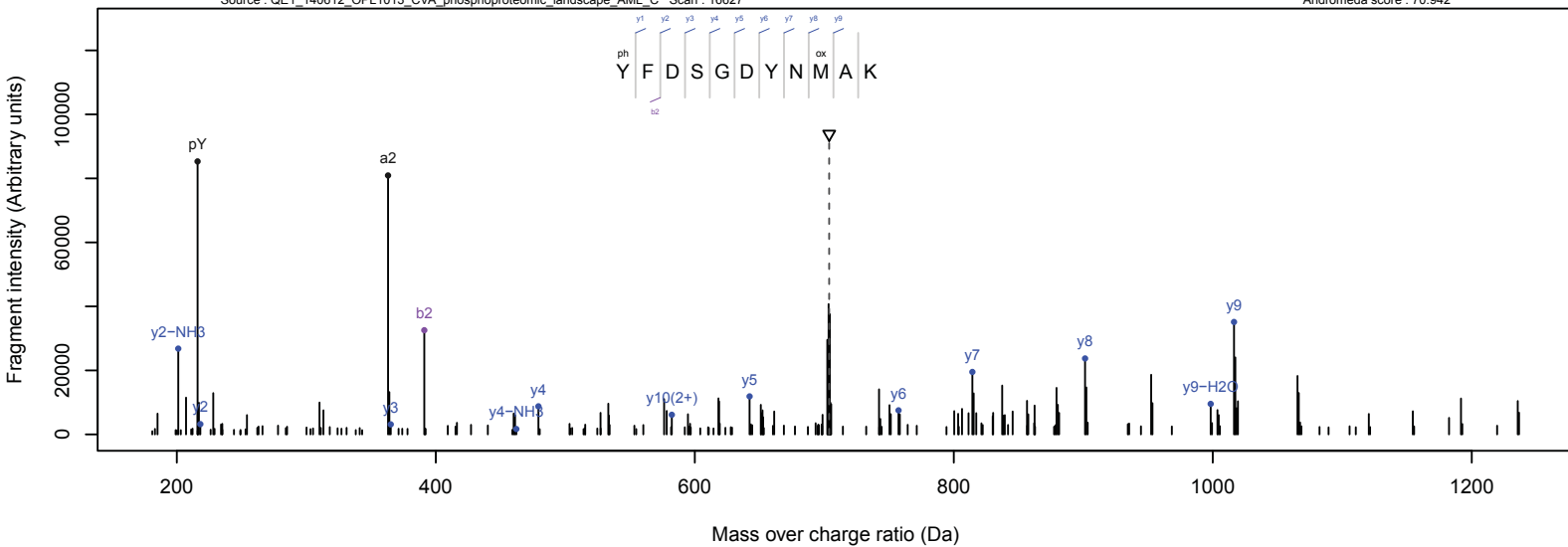
Andromeda score : 70.942

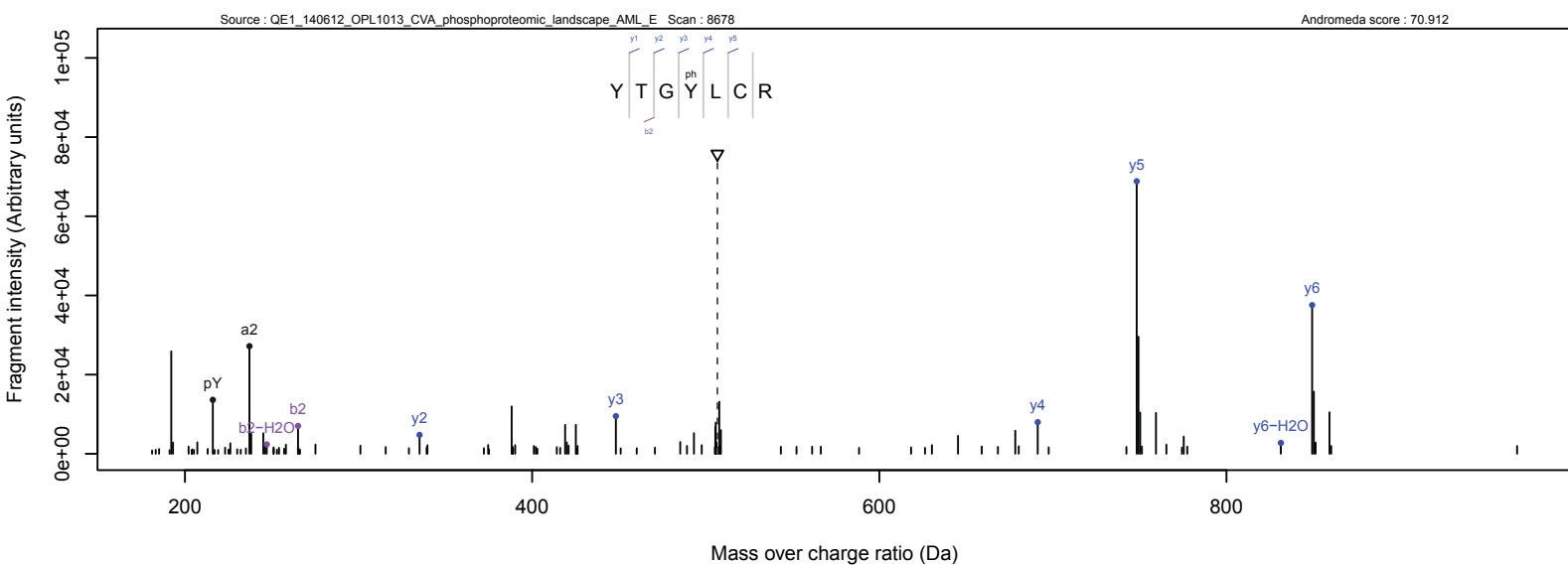
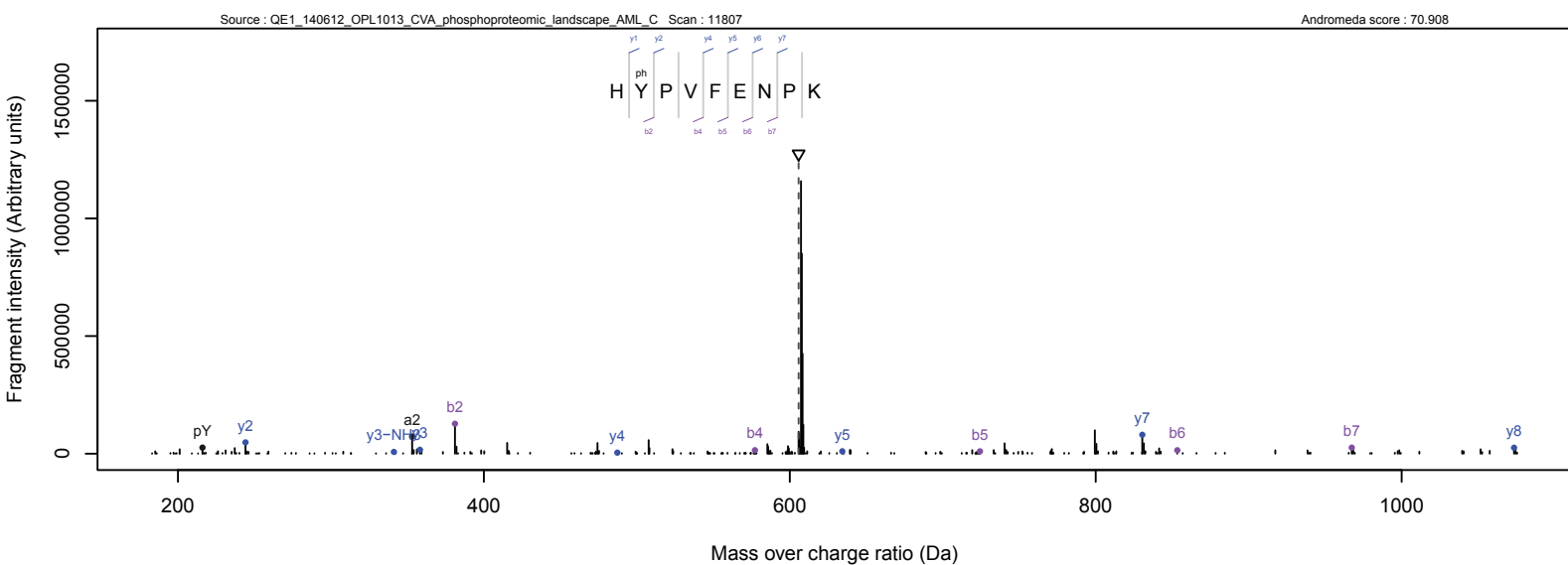
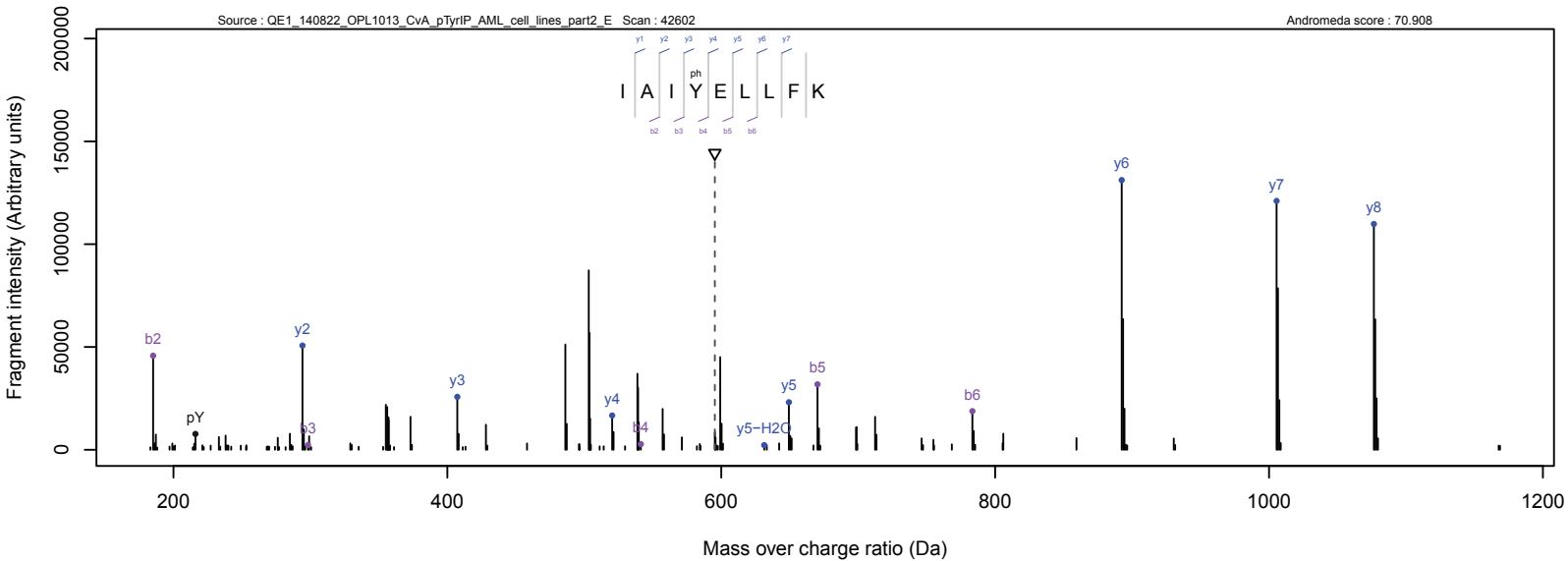


Y(ph)FDSGDYNM(ox)AK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16627

Andromeda score : 70.942



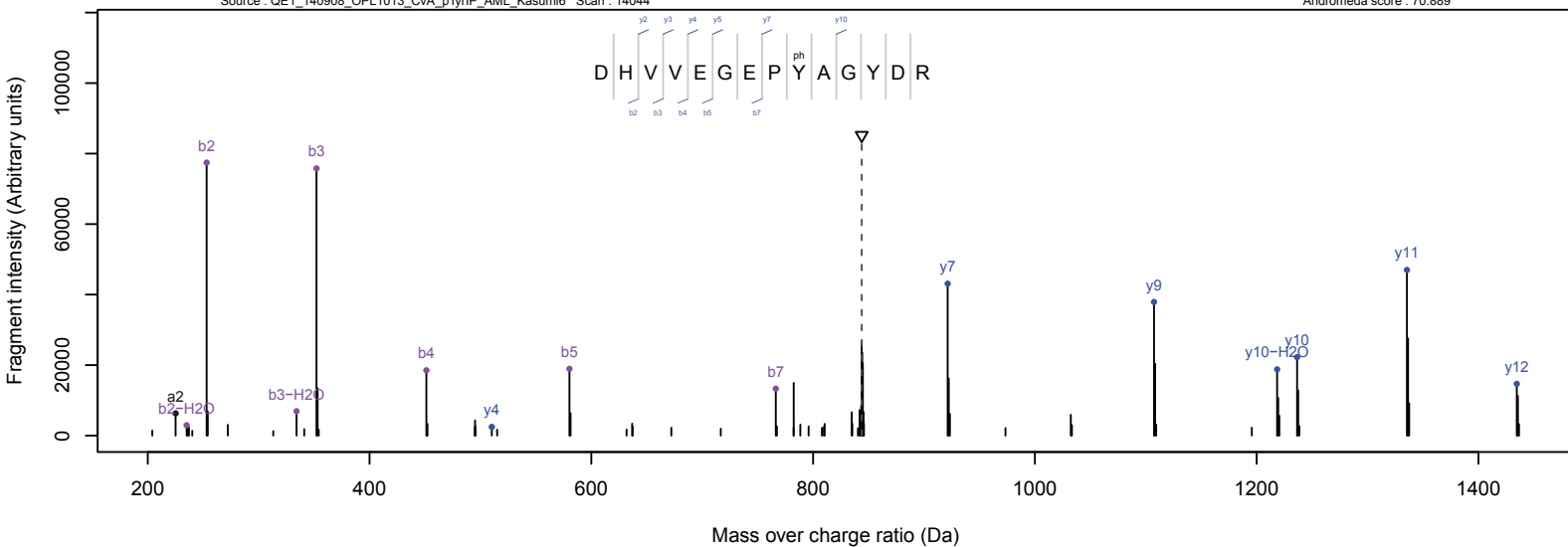
YTG<sup>(ph)</sup>LCR-2+HY<sup>(ph)</sup>PVFENPK-2+IAIY<sup>(ph)</sup>ELLFK-2+



## DHVVEGEPY(ph)AGYDR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 14044

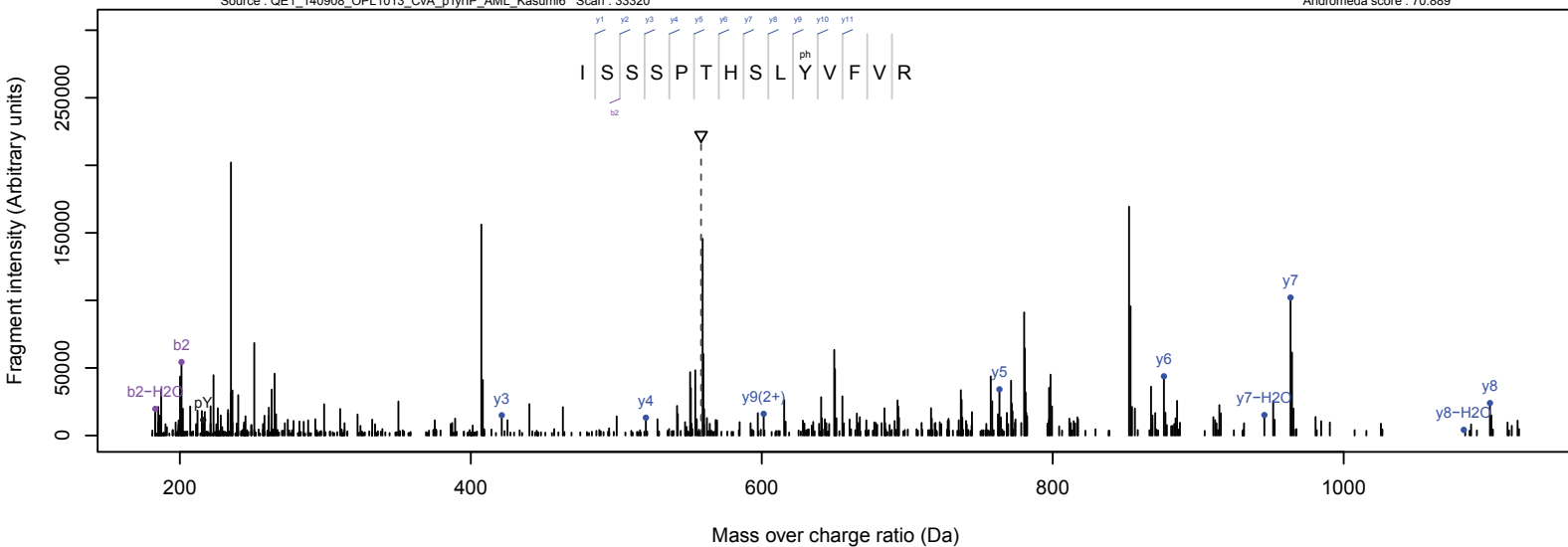
Andromeda score : 70.889



## ISSPHTSLY(ph)VFVR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 33320

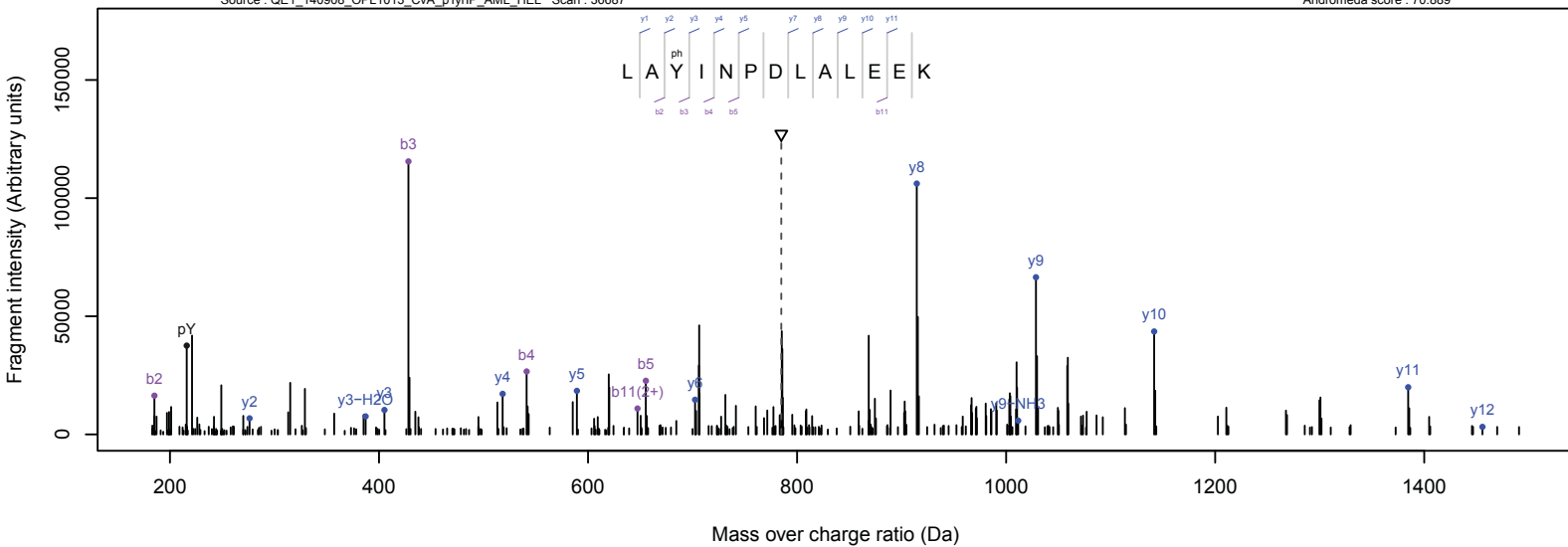
Andromeda score : 70.889



## LAY(ph)INPDLALEEK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 36687

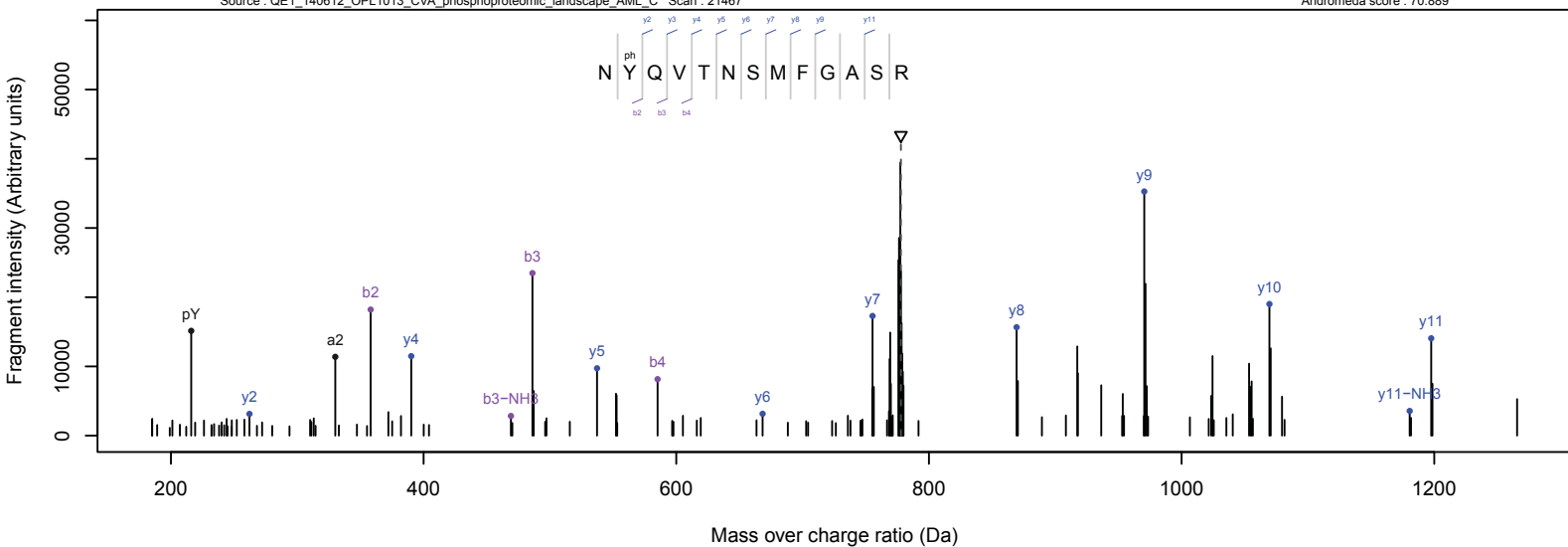
Andromeda score : 70.889



## NY(ph)QVTNSMFGASR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21467

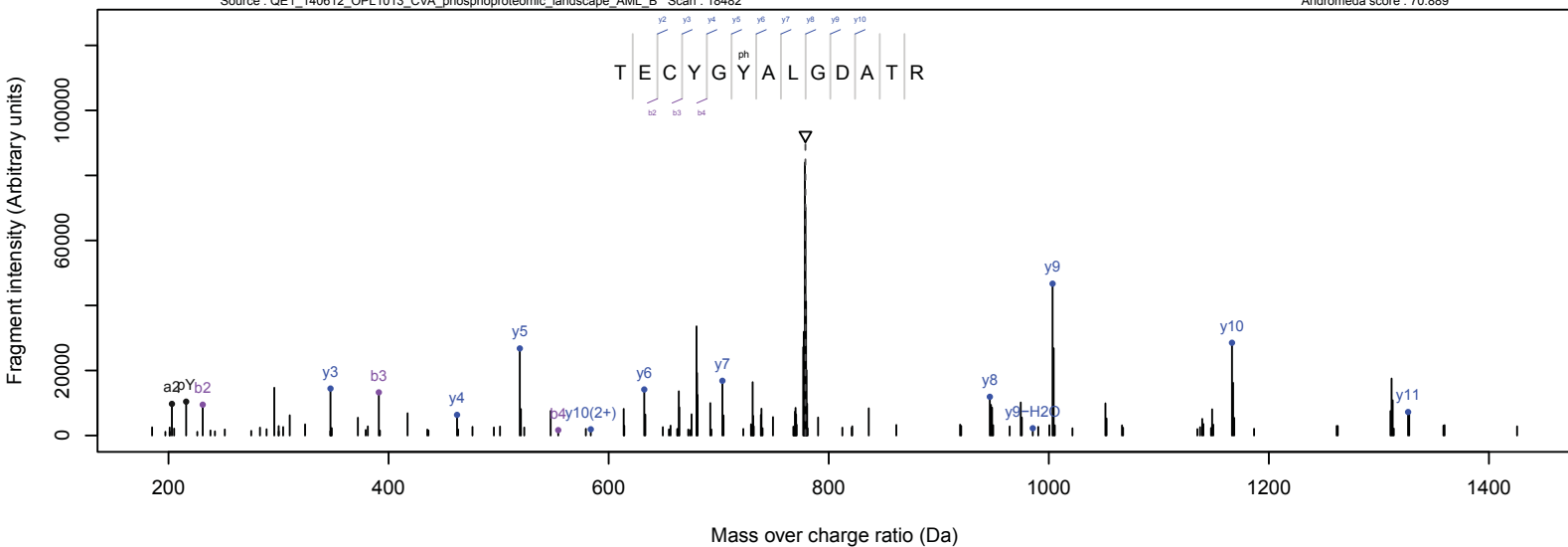
Andromeda score : 70.889



## TECYG(ph)ALGDATR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18482

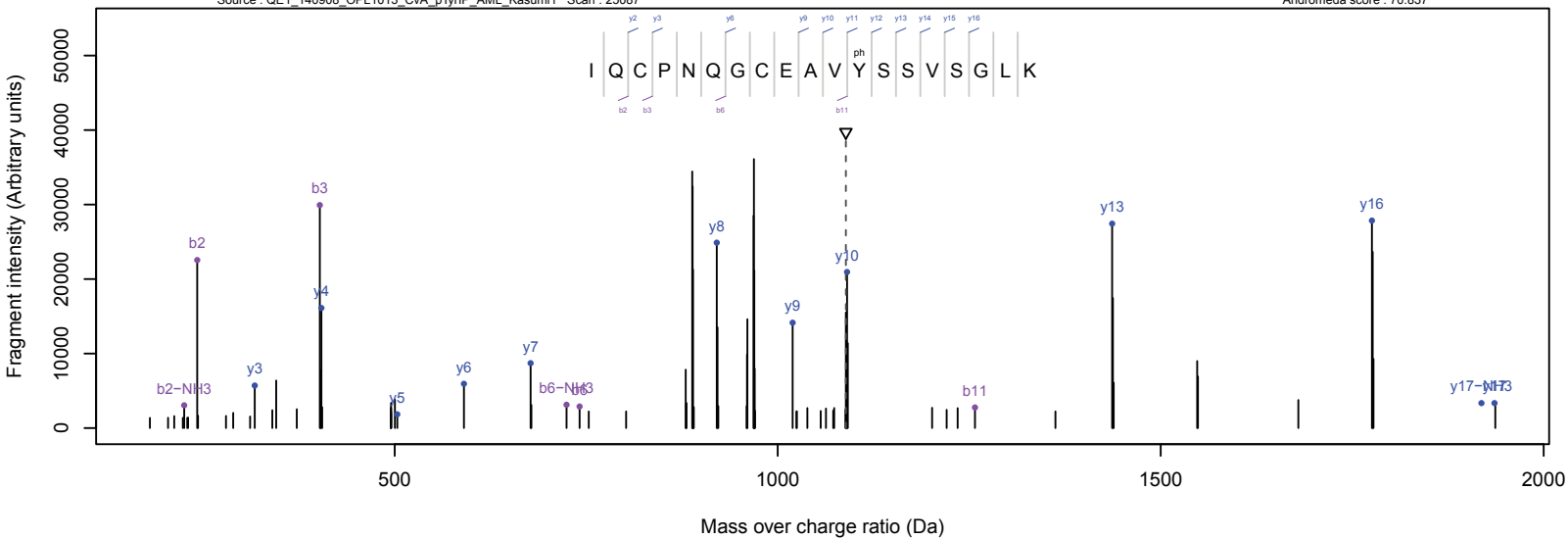
Andromeda score : 70.889



## IQCPNQGEAVY(ph)SSVSLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 25087

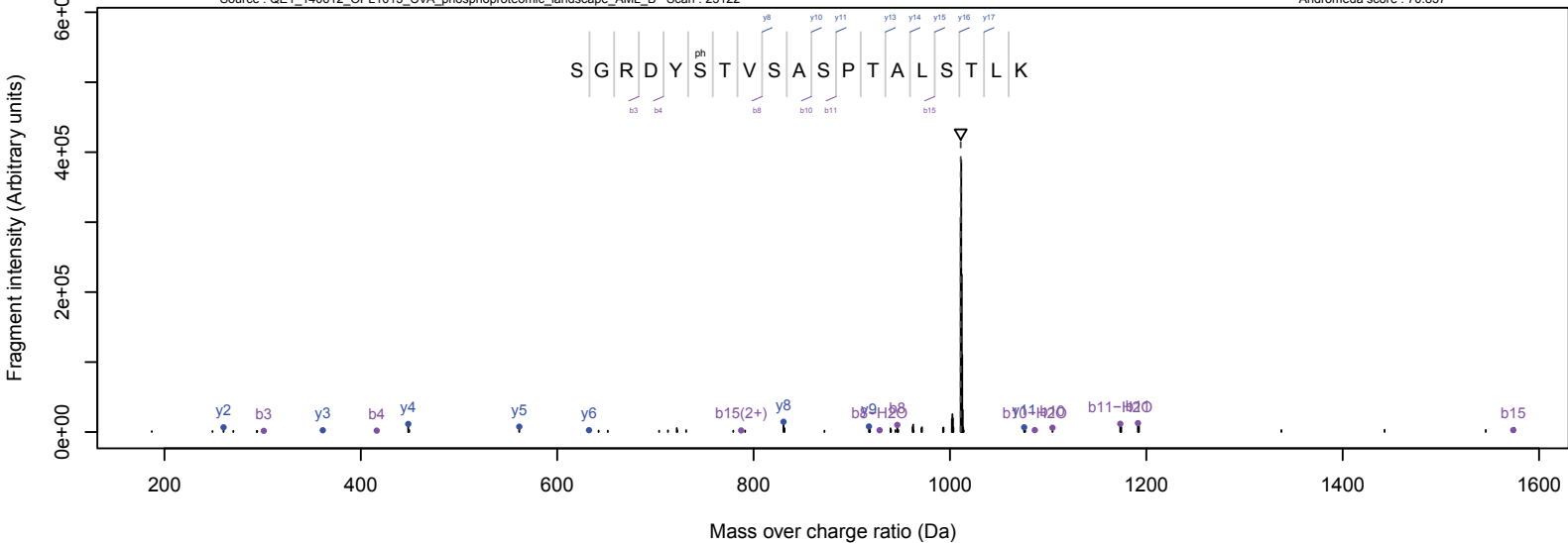
Andromeda score : 70.837



SGRDYS(ph)TVSASPTALSTLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23122

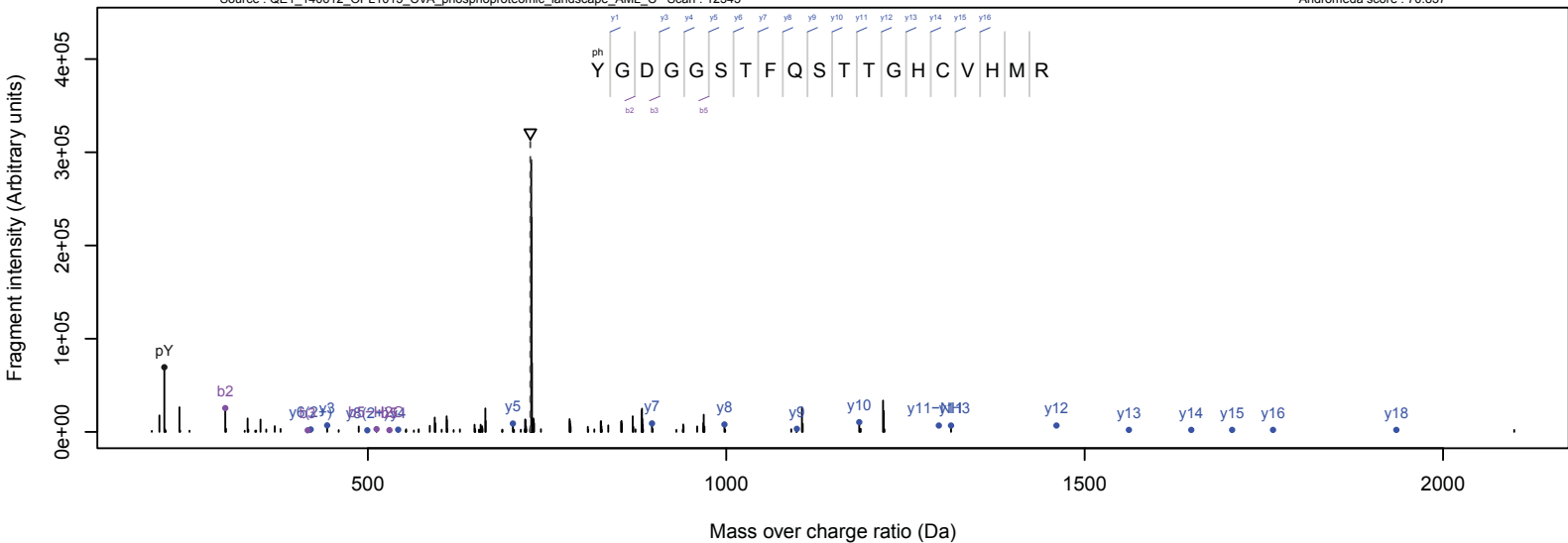
Andromeda score : 70.837



Y(ph)GDGGSTFQSTTGHCVHMR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12345

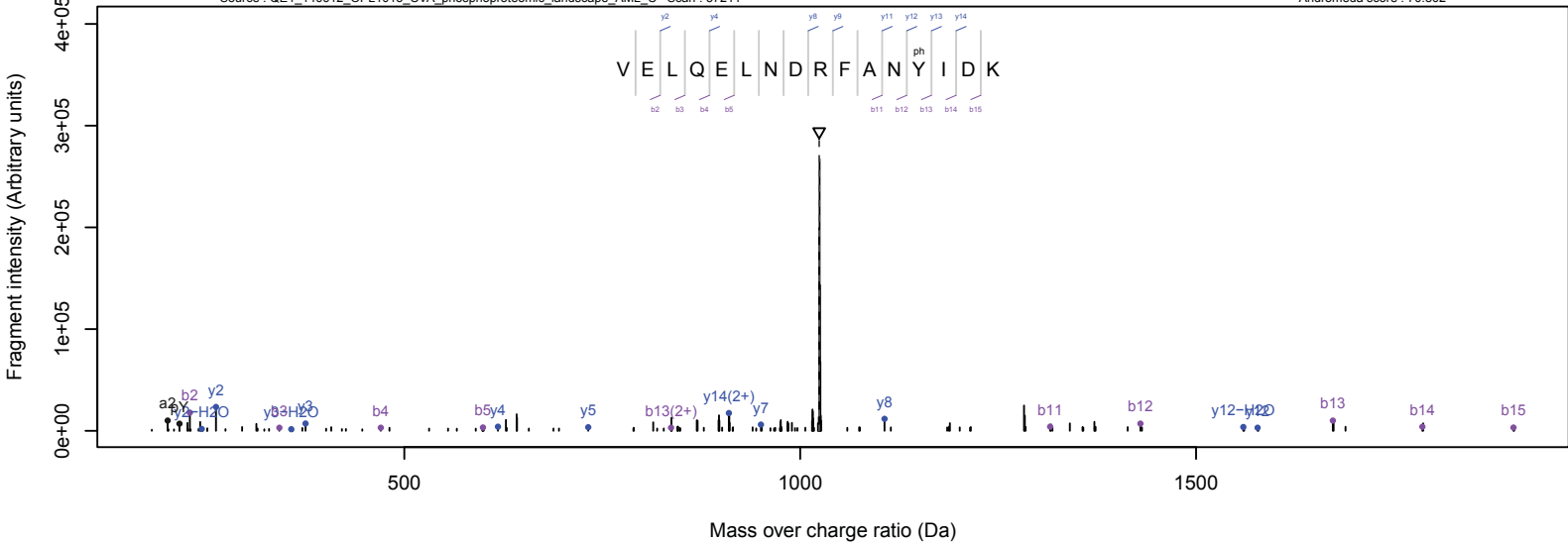
Andromeda score : 70.837



VELQELNDRFANYIDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37211

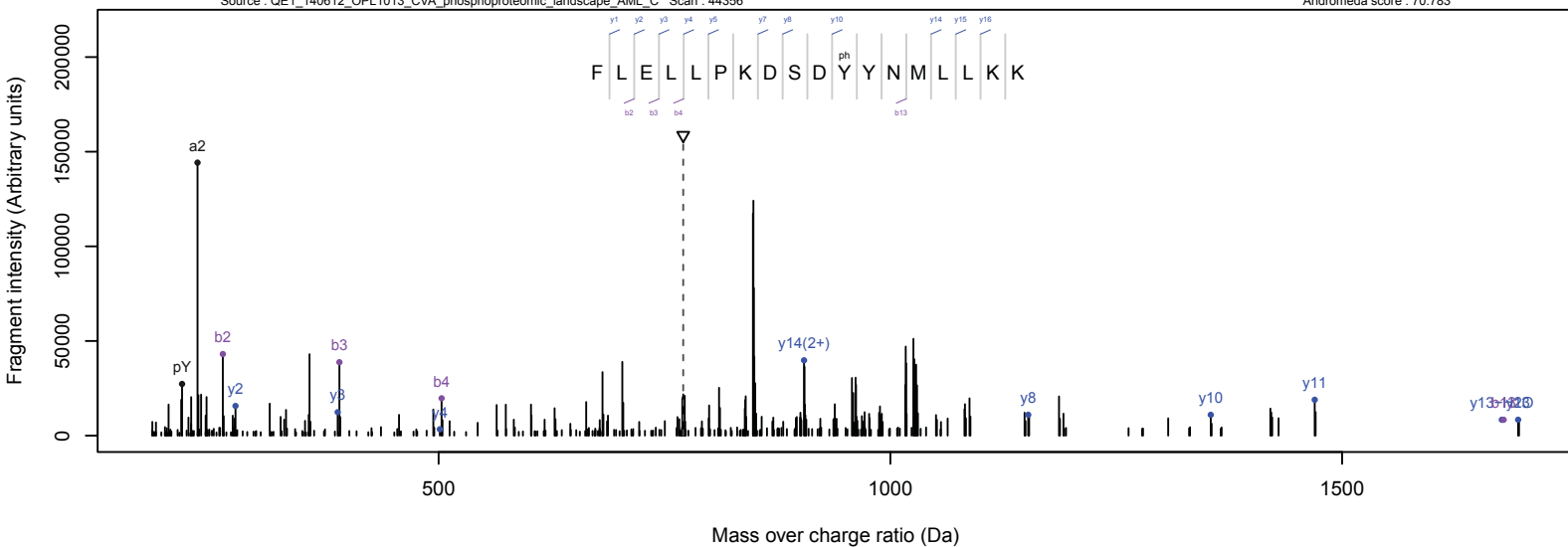
Andromeda score : 70.802



## FLELLPKDSY(ph)YNMLLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44356

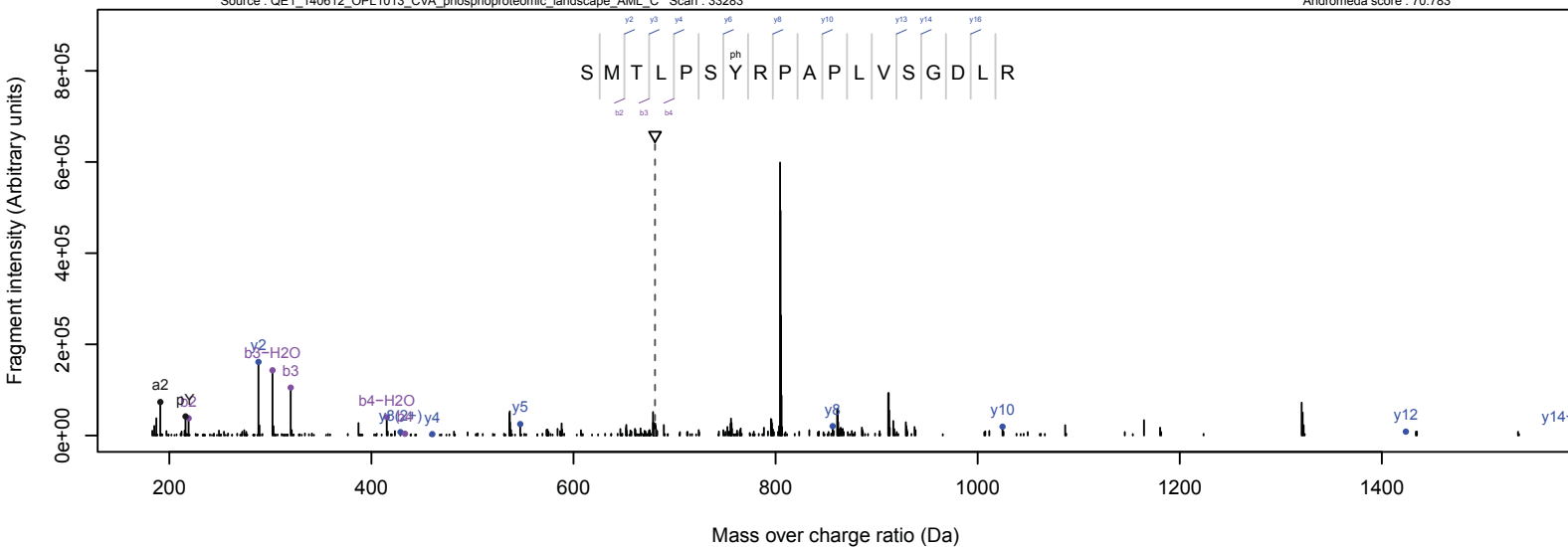
Andromeda score : 70.783



## SMTLPSY(ph)RPAPLVSGDLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33283

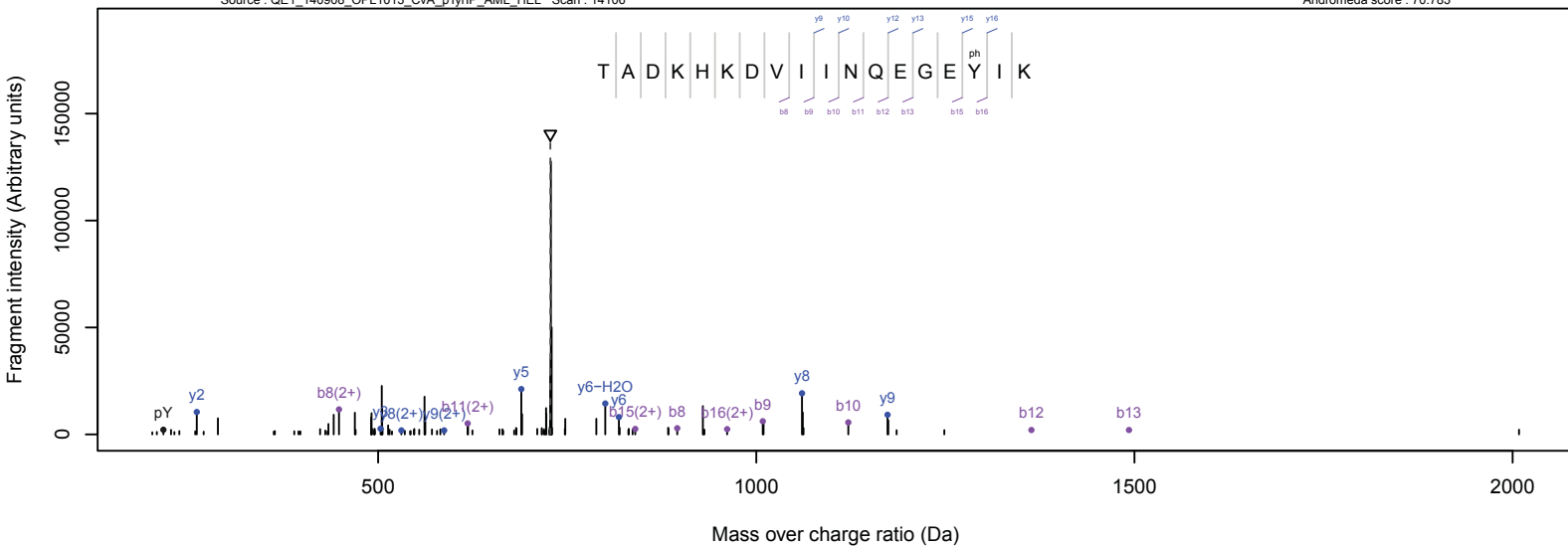
Andromeda score : 70.783



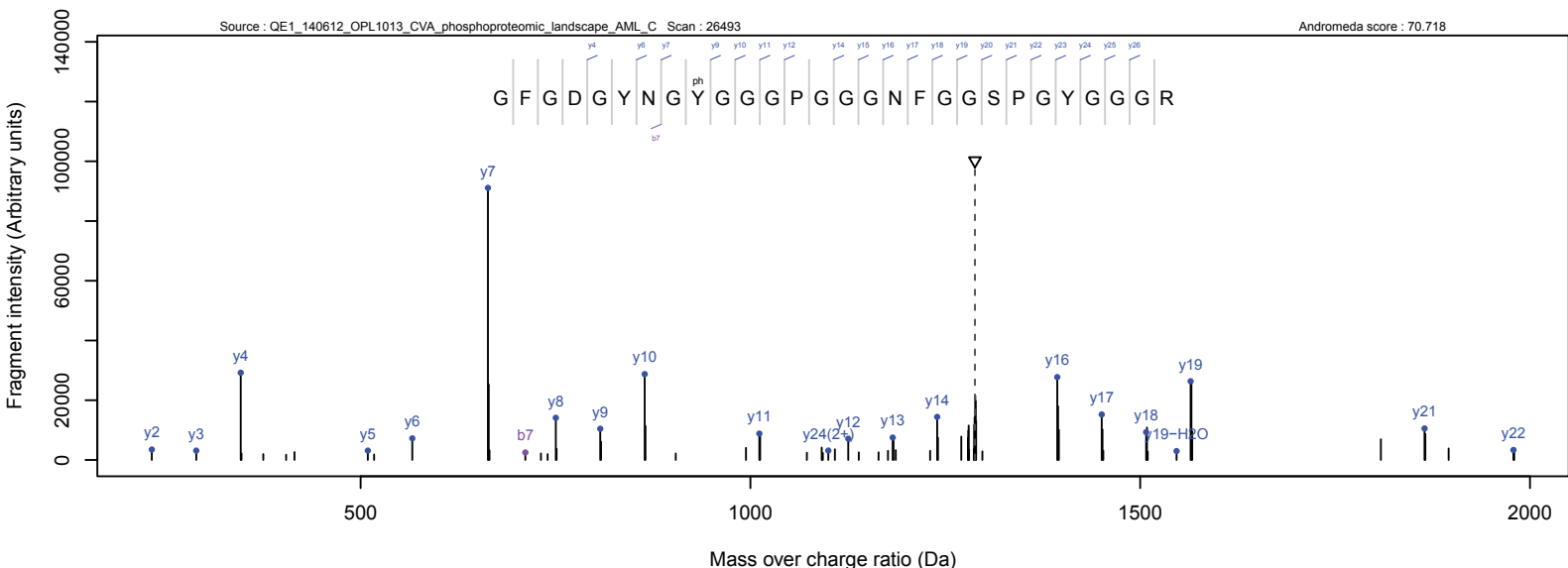
## TADKHKDVIINQEGEY(ph)IK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 14106

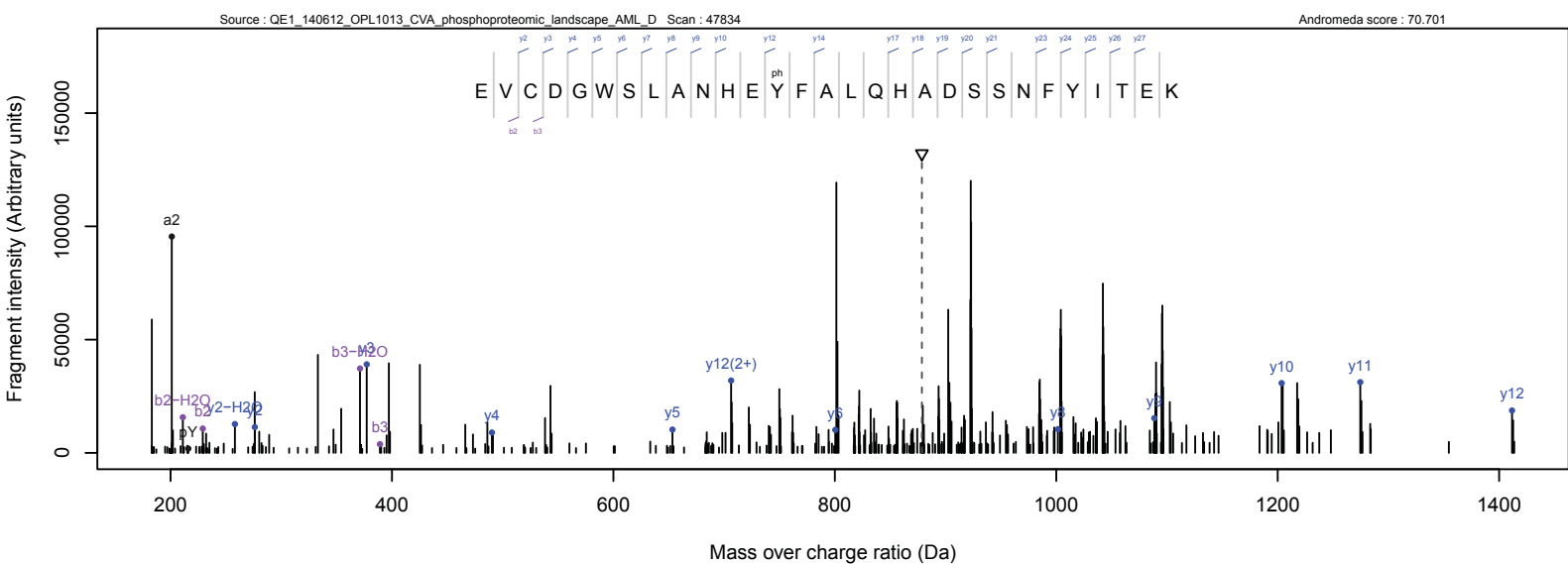
Andromeda score : 70.783



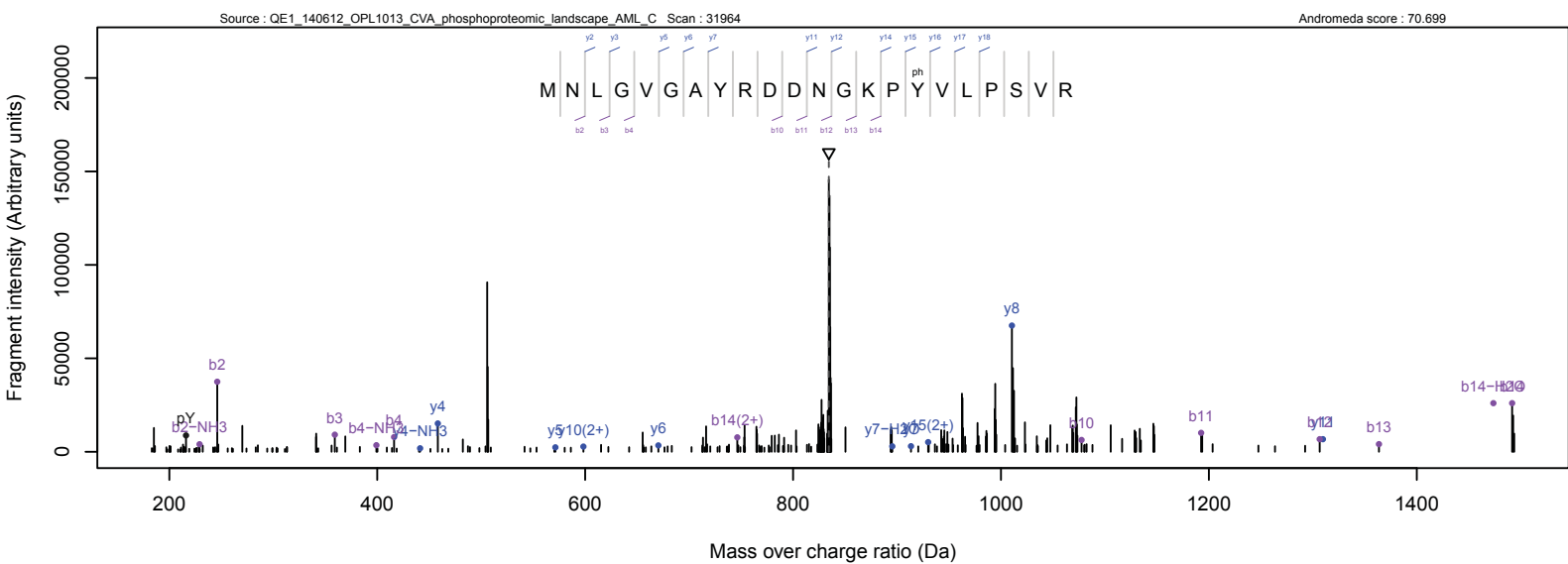
**GFGDGYNGY(ph)GGPGGGNFGGSPGYGGGR-2+**



**EVCDGWSLANHEY(ph)FALQHADSSNFYITEK-4+**



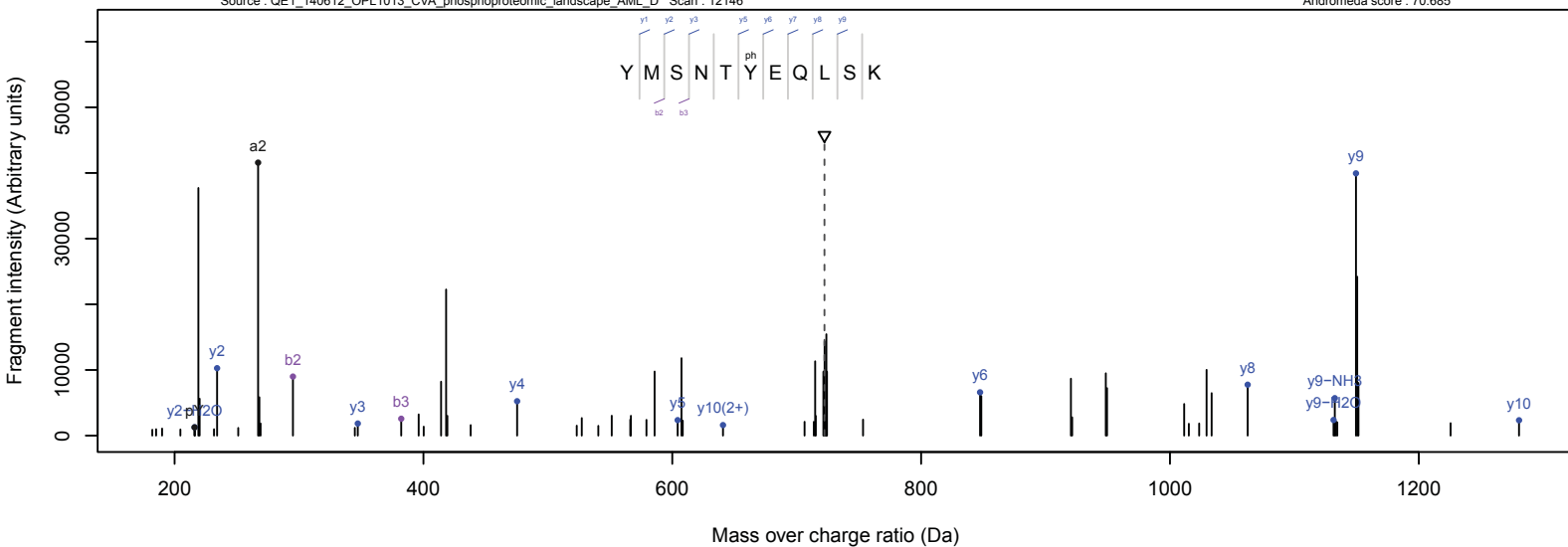
**MNLGVGAYRDDNGKPYVLPVSR-3+**



## YMSNTY(ph)EQLSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12146

Andromeda score : 70.685

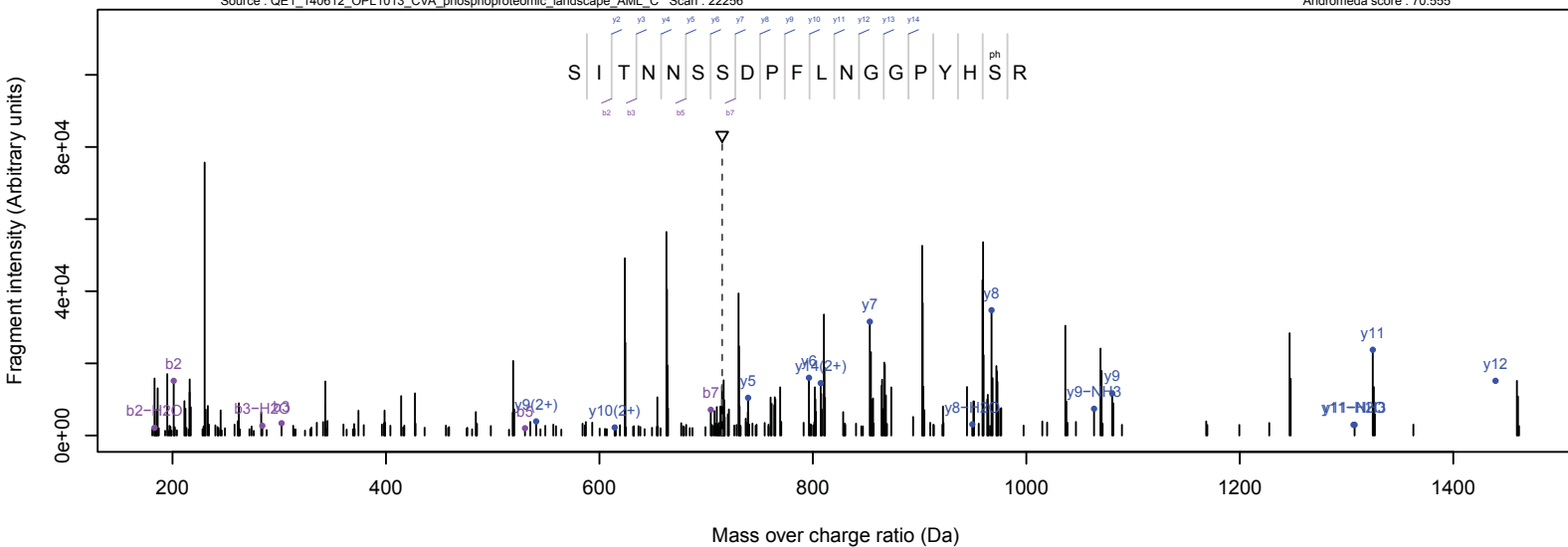


Mass over charge ratio (Da)

## SITNNSDPFLNGGPYHSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22256

Andromeda score : 70.555

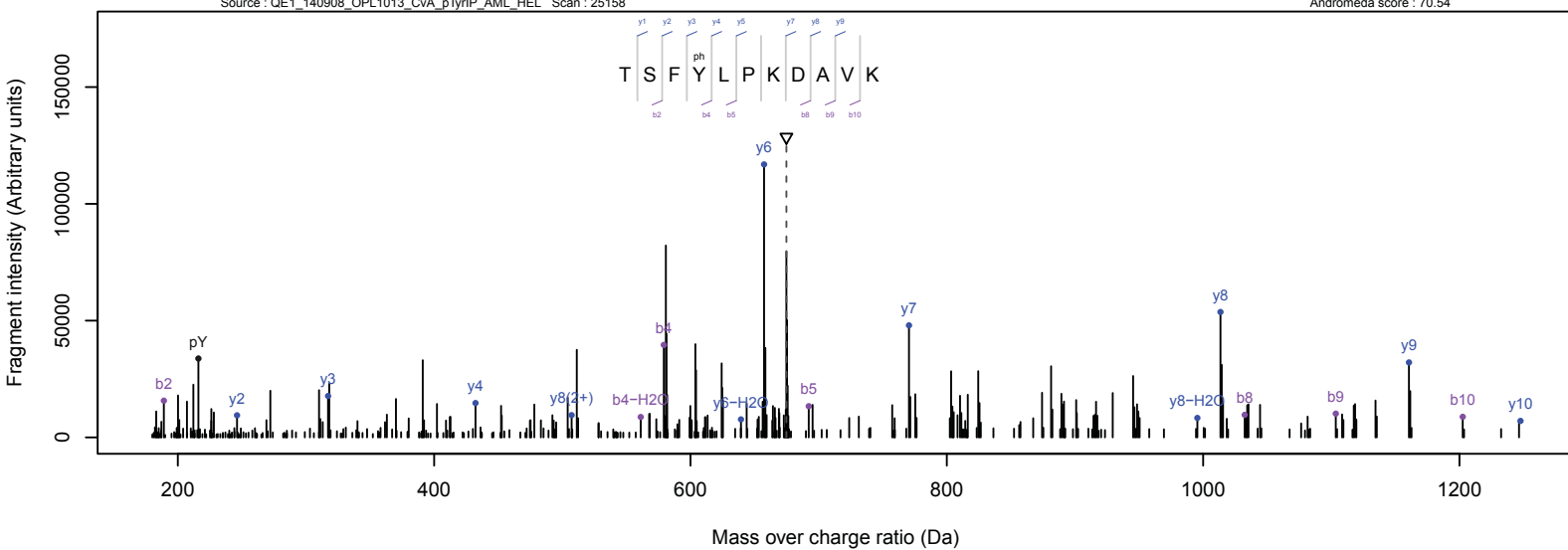


Mass over charge ratio (Da)

## TSFY(ph)LPKDAVK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 25158

Andromeda score : 70.54

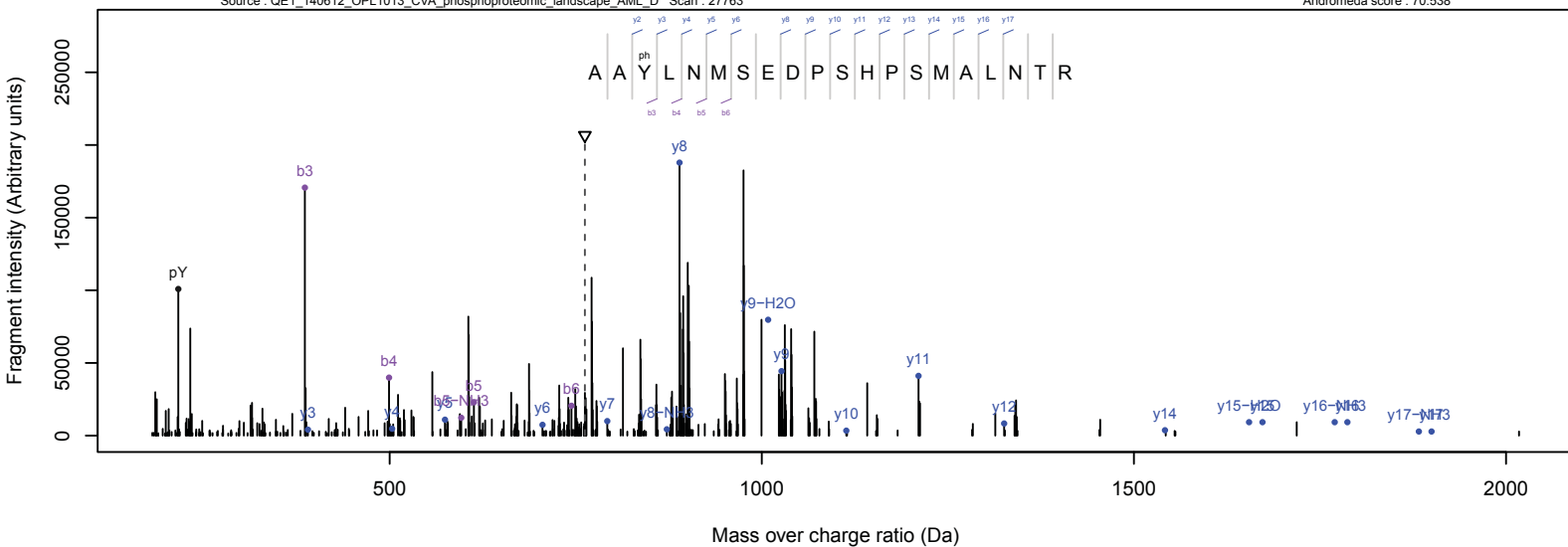


Mass over charge ratio (Da)

**AA Y(ph)LNMS EDP SHP S M A L N T R**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 27763

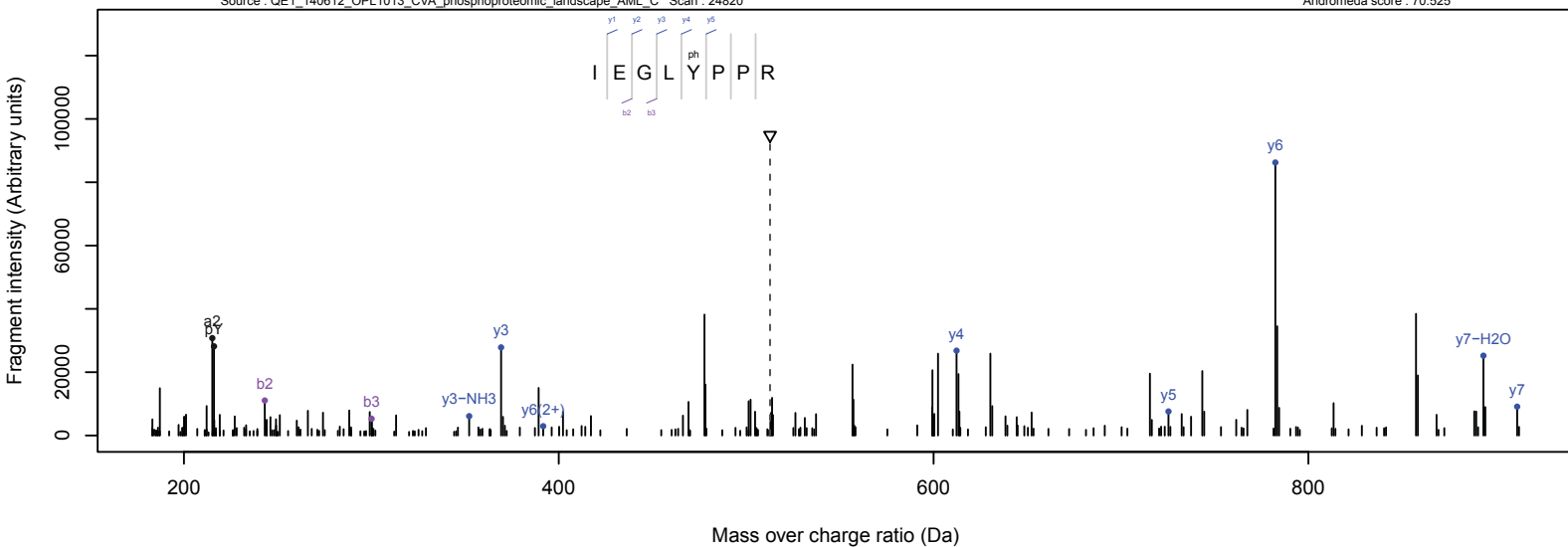
Andromeda score : 70.538



**IEGL Y(ph) P P R**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24820

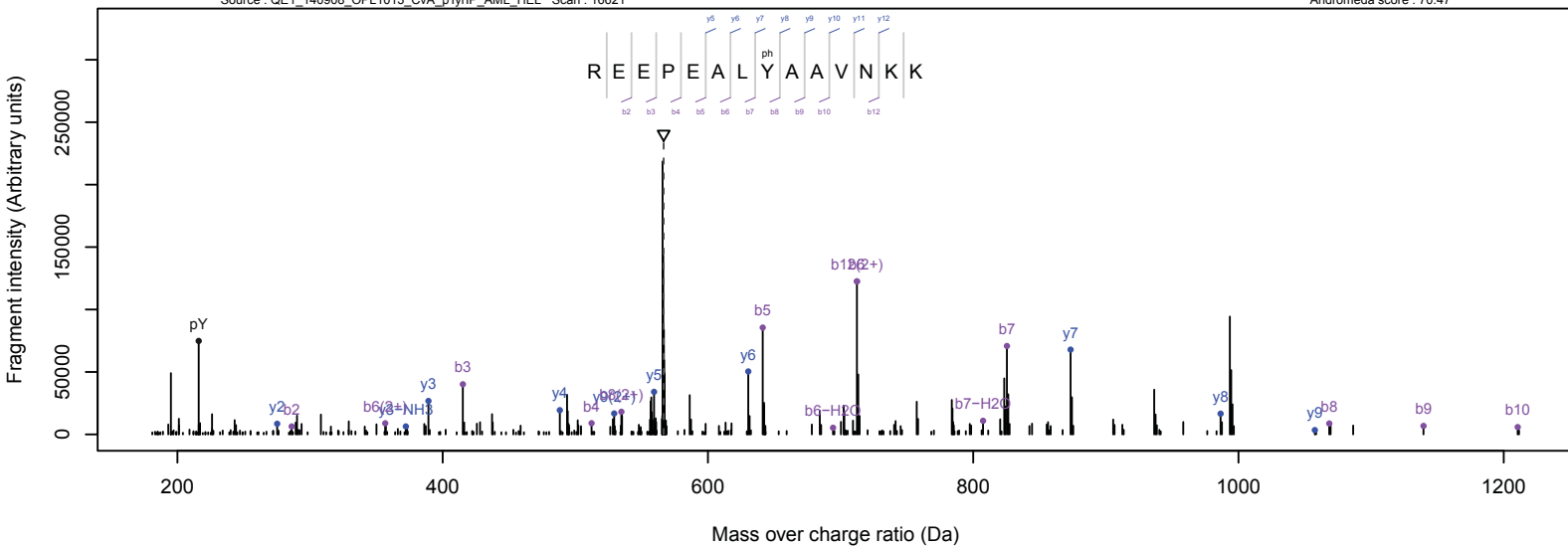
Andromeda score : 70.525



**REEPEAL Y(ph) AAVNKK**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 16621

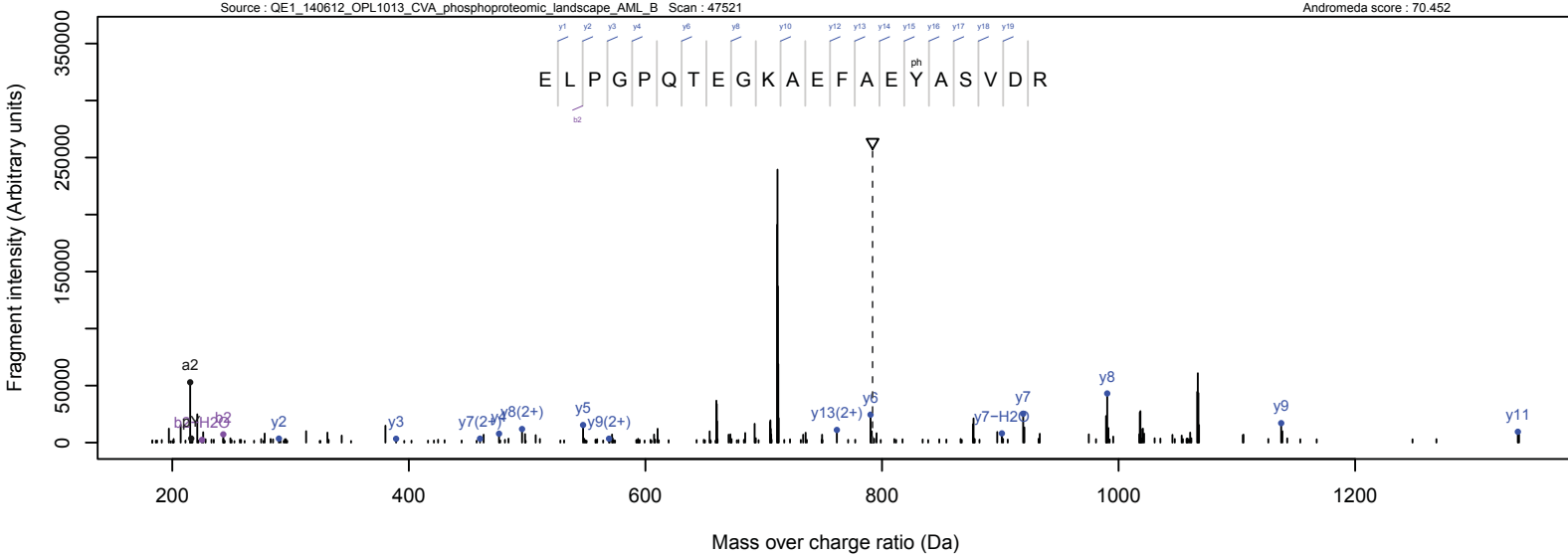
Andromeda score : 70.47



## ELPGPQTEGKAEYASVDR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 47521

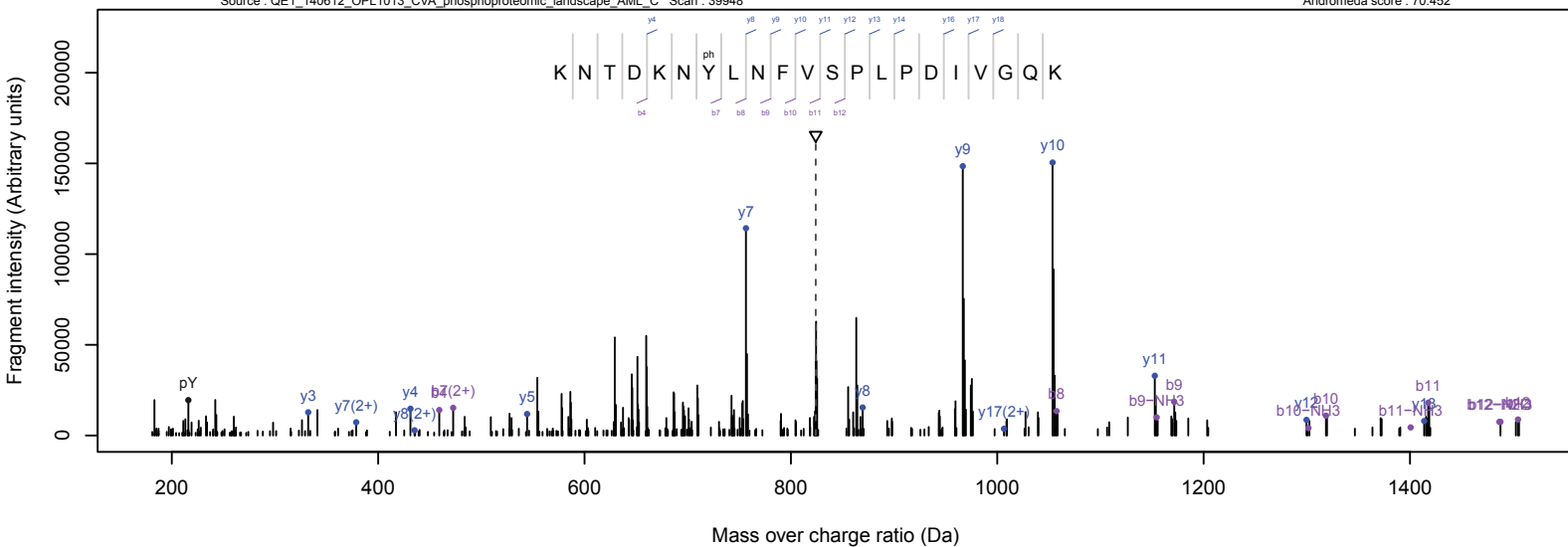
Andromeda score: 70.452



## KNTDKNYLNFVSPDPDVGQK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 39948

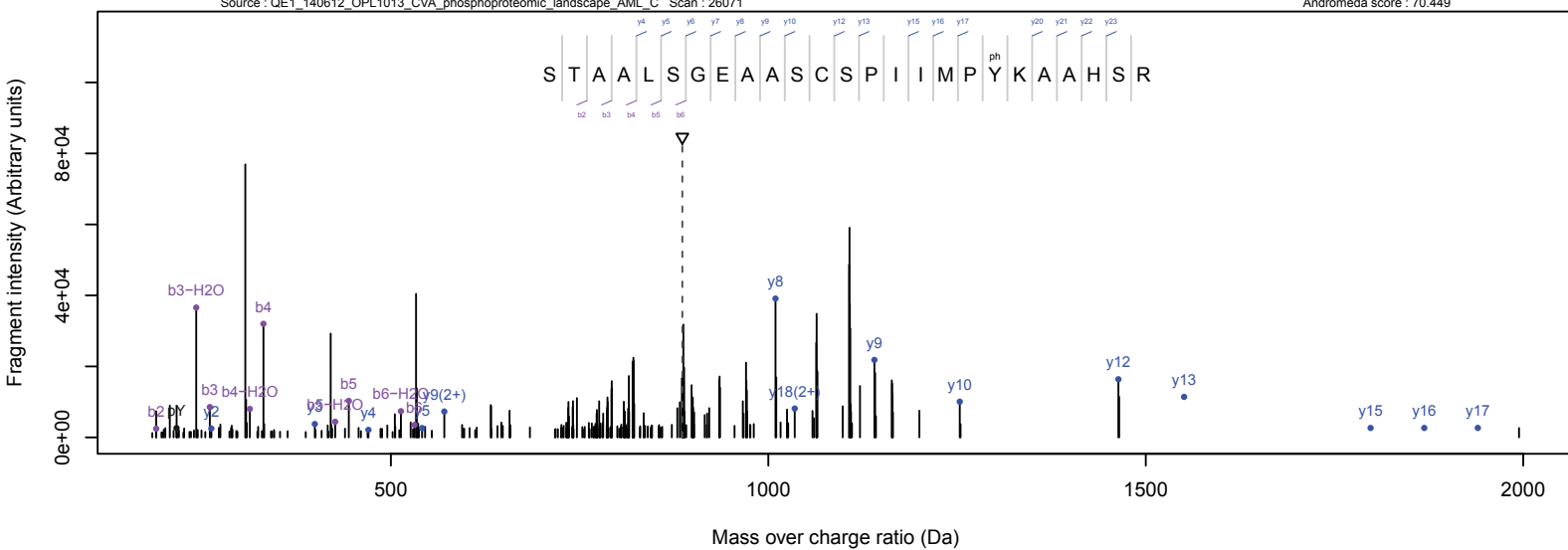
Andromeda score: 70.452



## STAALSGEAASCSPYKAAHSR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26071

Andromeda score: 70.449

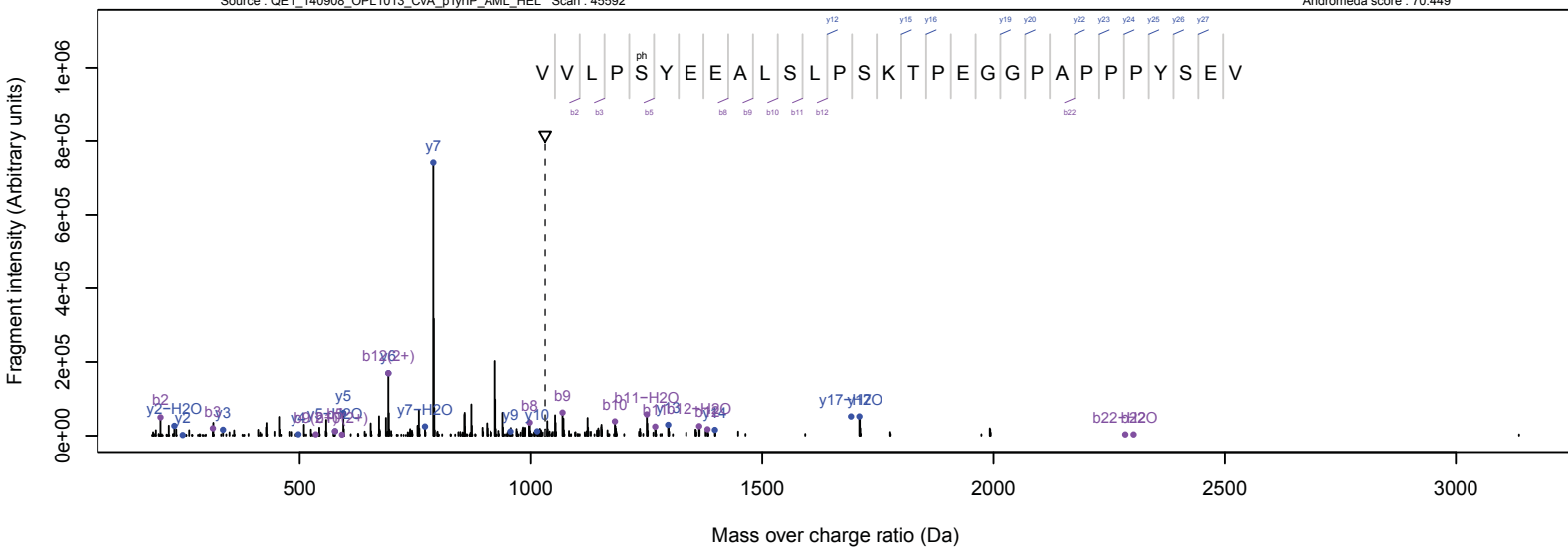




VVLPS(ph)YEEALS LPSK TPEG GPAPPYSEV-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 45592

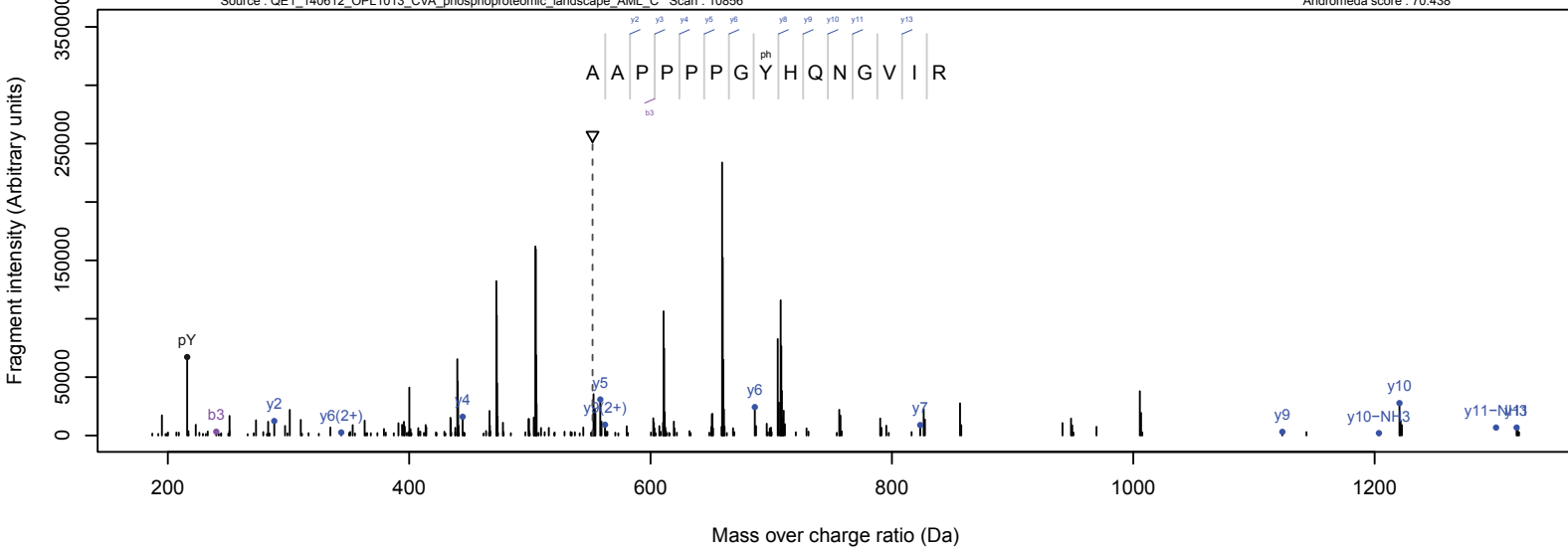
Andromeda score : 70.449



AAPPPGY(ph)HQNGVIR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 10856

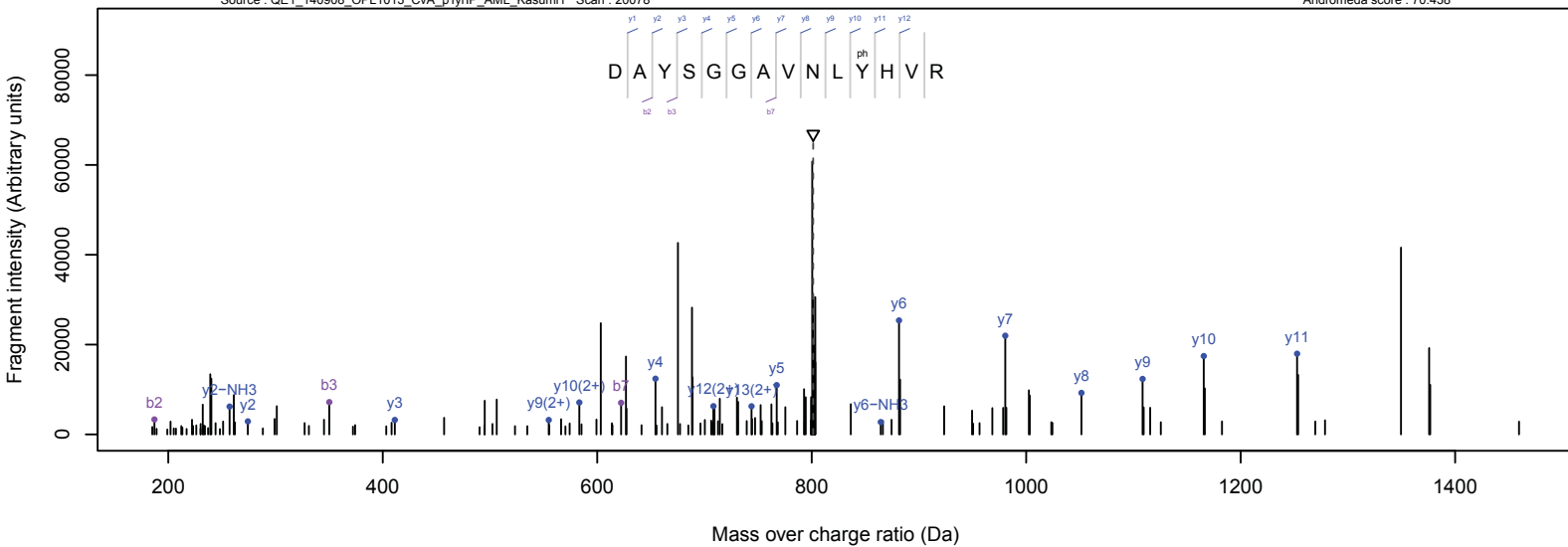
Andromeda score : 70.438



DAYSGAVNLY(ph)HVR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 20078

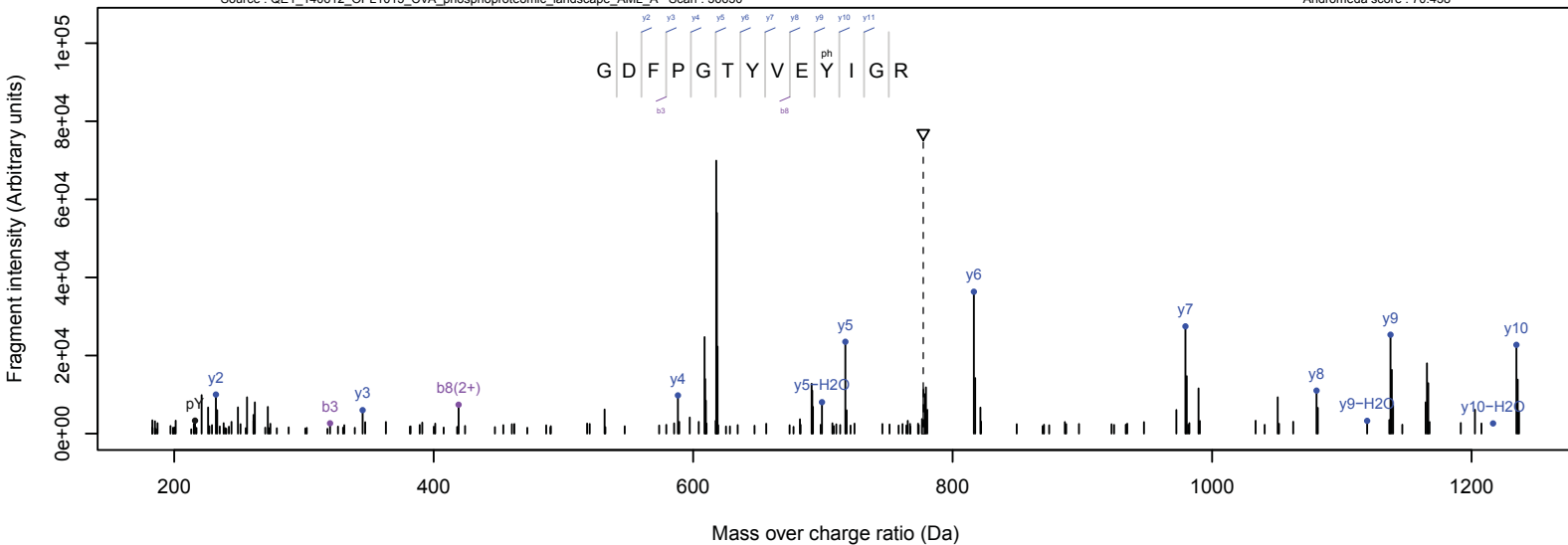
Andromeda score : 70.438



## GDFPGTYVEY(ph)IGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36630

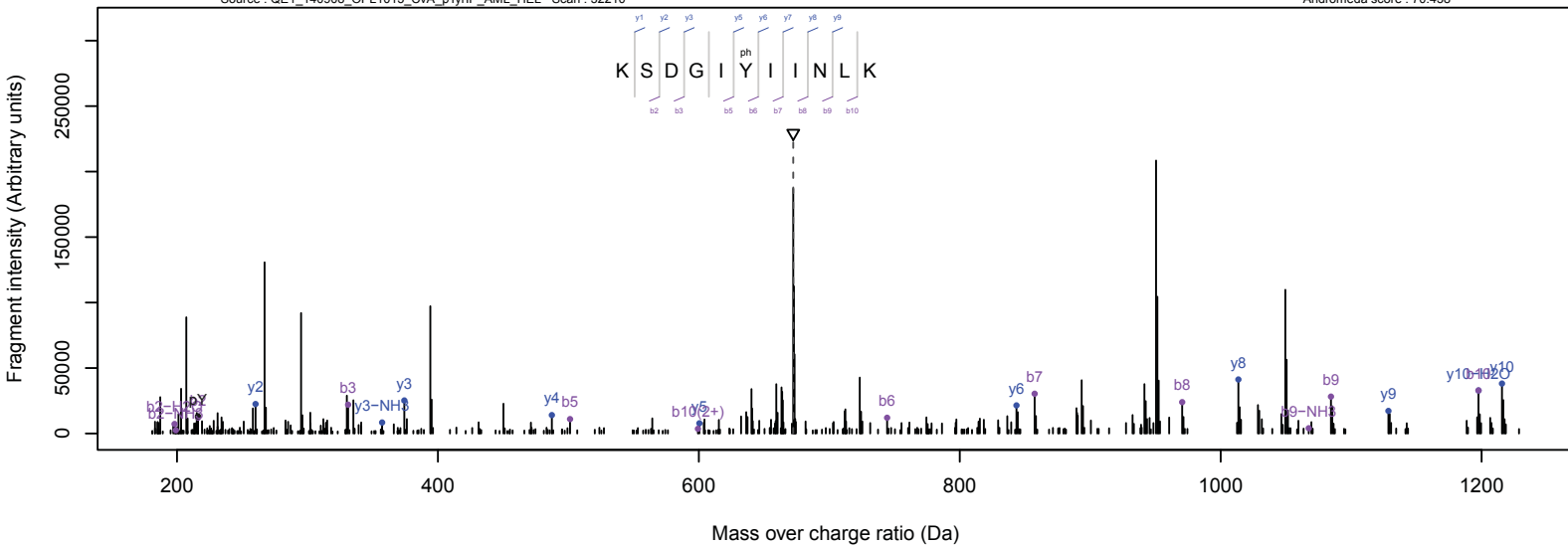
Andromeda score : 70.438



## KSDGIY(ph)IINLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32210

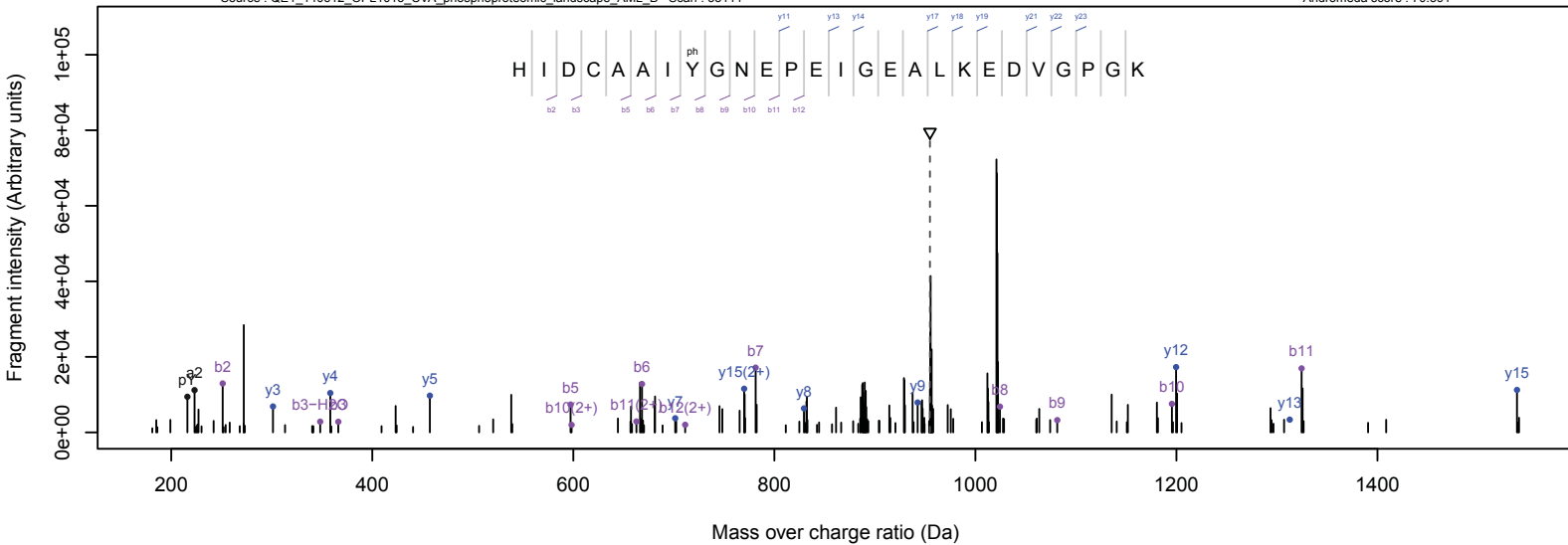
Andromeda score : 70.438



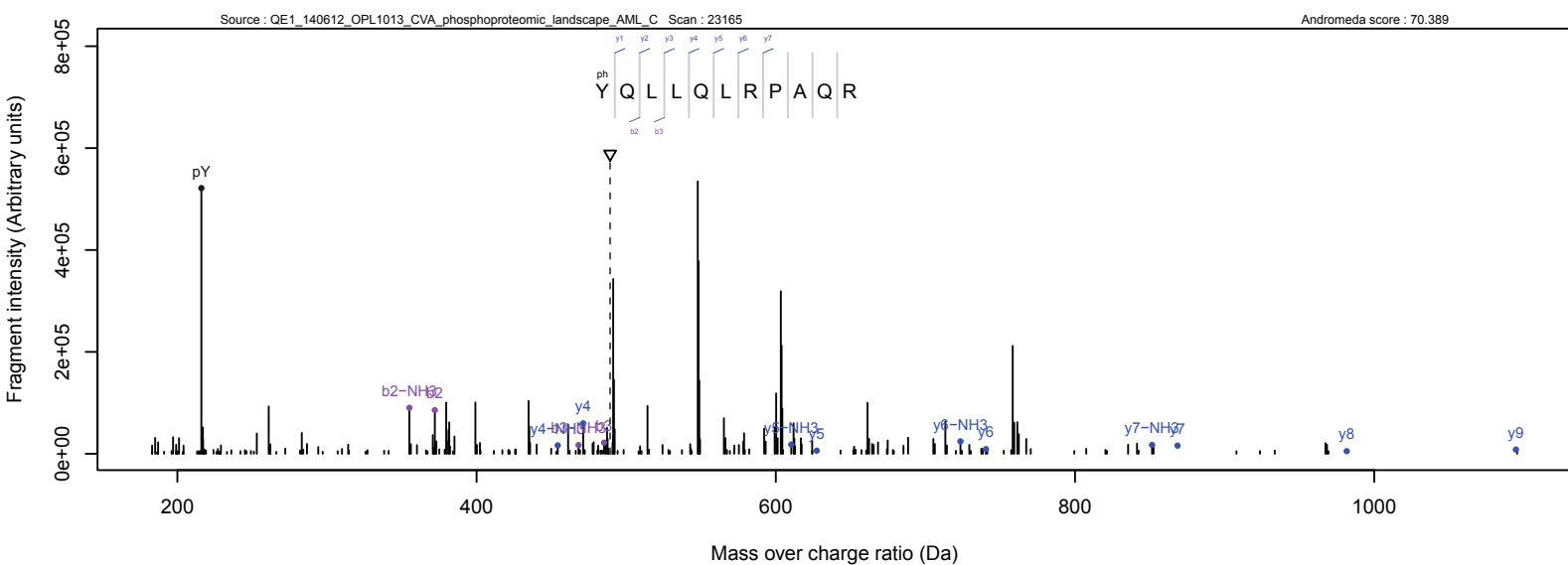
## HIDCAAIY(ph)GNEPEIGEALKEVDVGP GK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33111

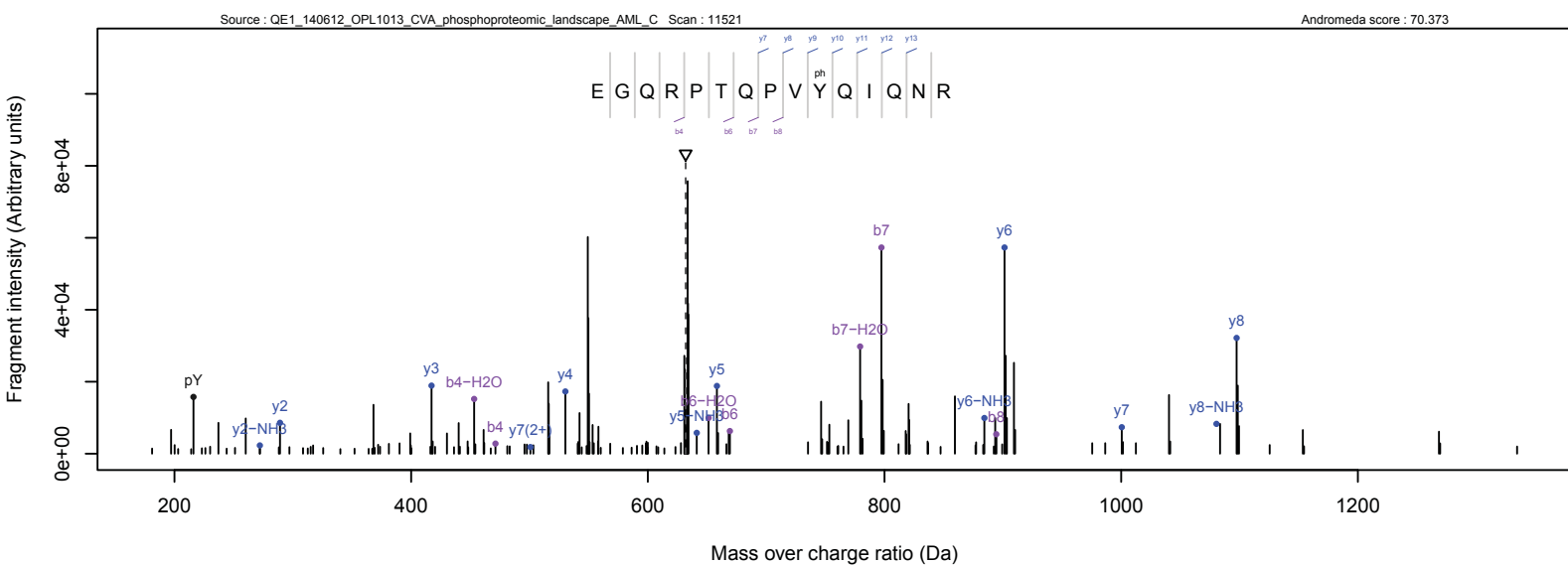
Andromeda score : 70.391



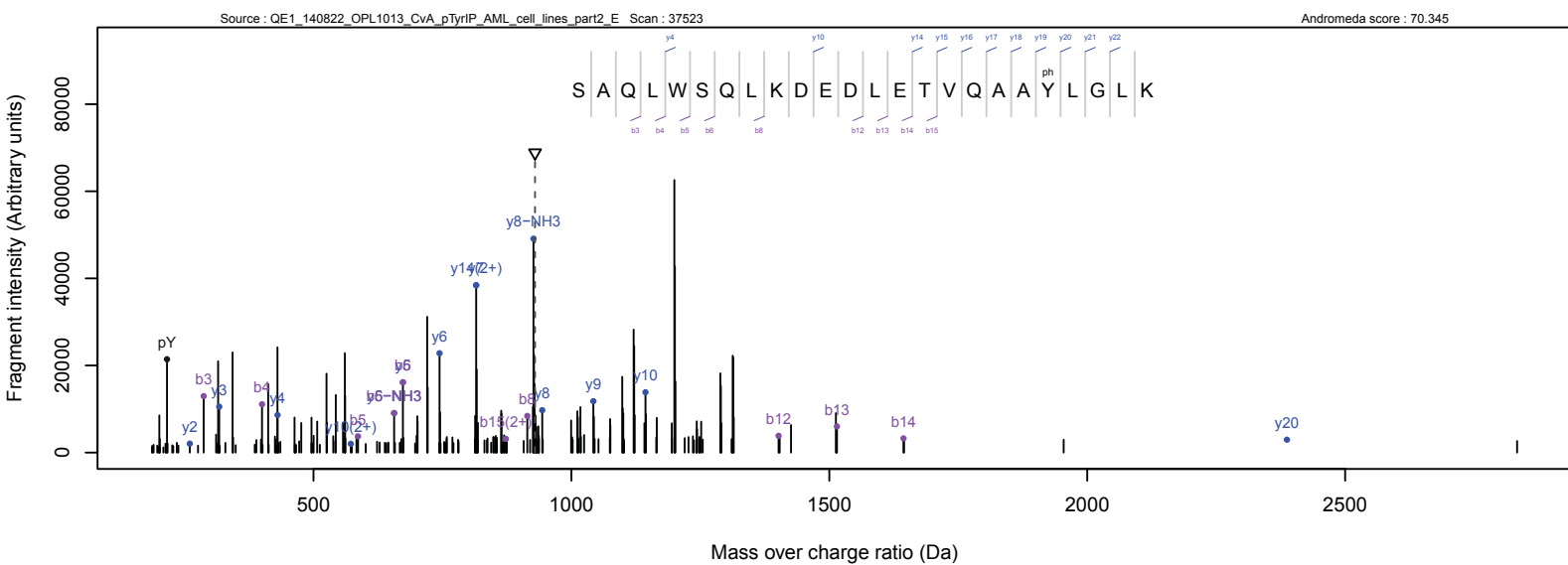
## Y(ph)QLQLRPAQR-3+



## EGQRPTQPVYQIQNR-3+



## SAQLWSQLKDEDLETVQAAYLGLK-3+

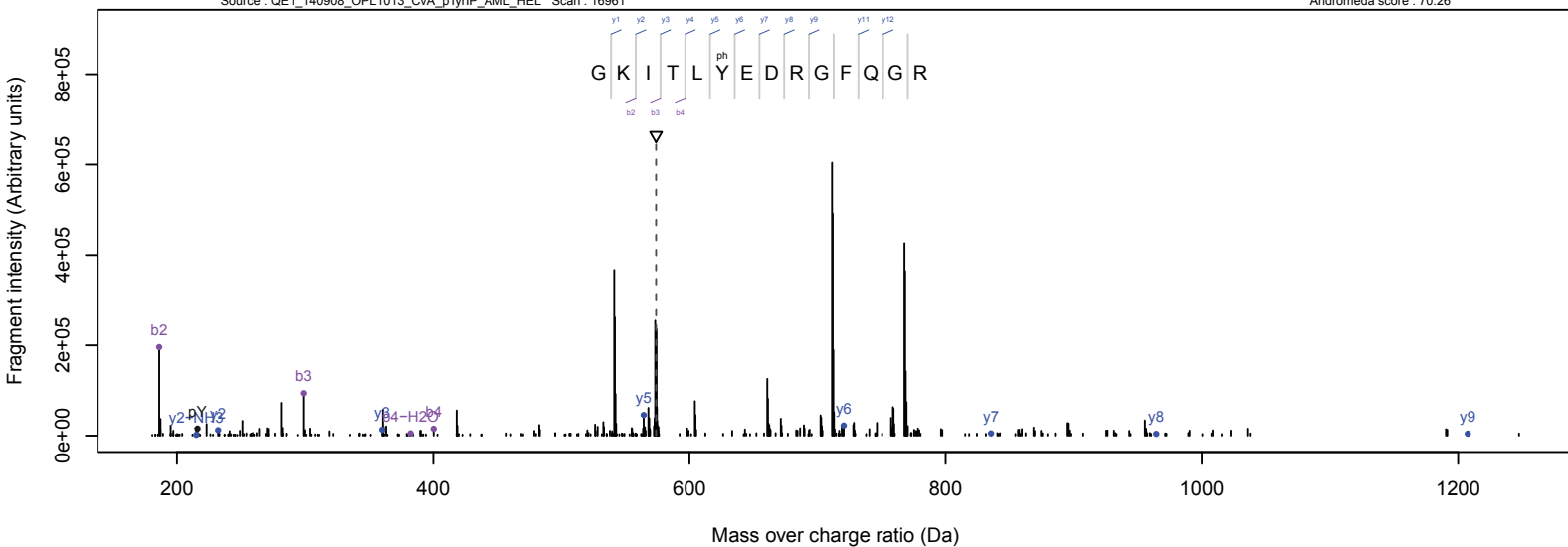




## GKITLY(ph)EDRGFQGR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 16961

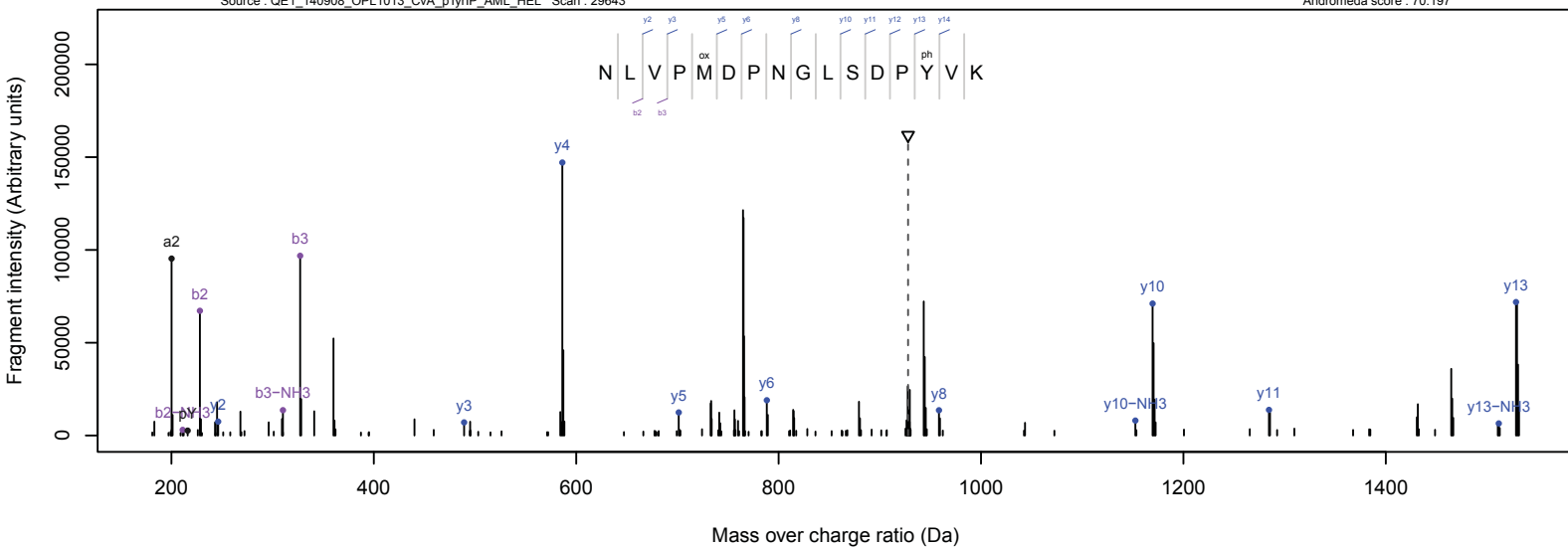
Andromeda score : 70.26



## NLVPM(ox)DPNGLSDPY(ph)VK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29643

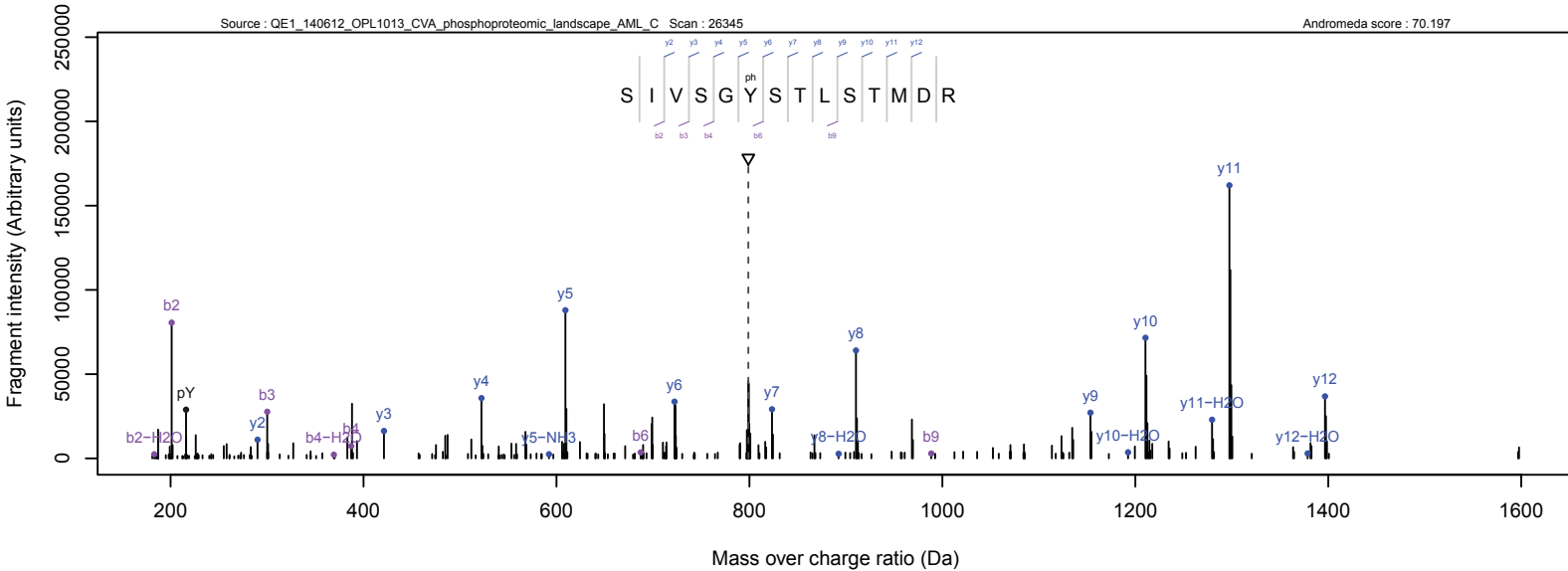
Andromeda score : 70.197



## SIVSGY(ph)STLSTMDR-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 26345

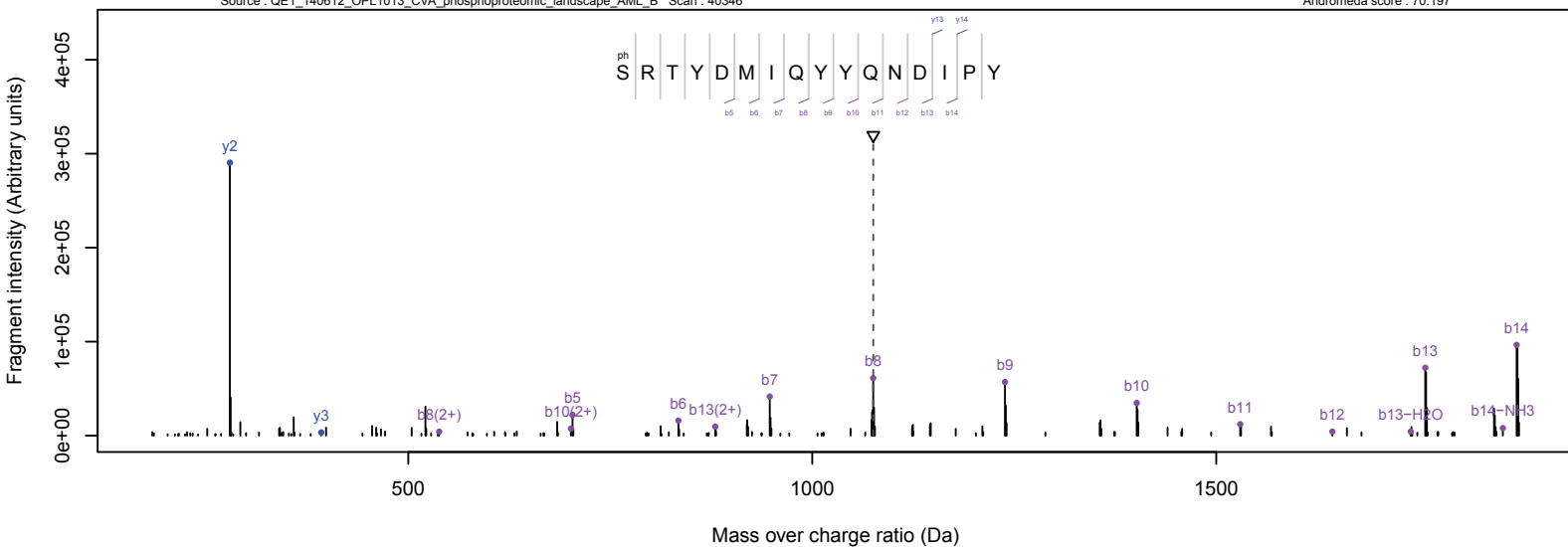
Andromeda score : 70.197



**S(ph)RTYDMIQYYQNDIPY-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40346

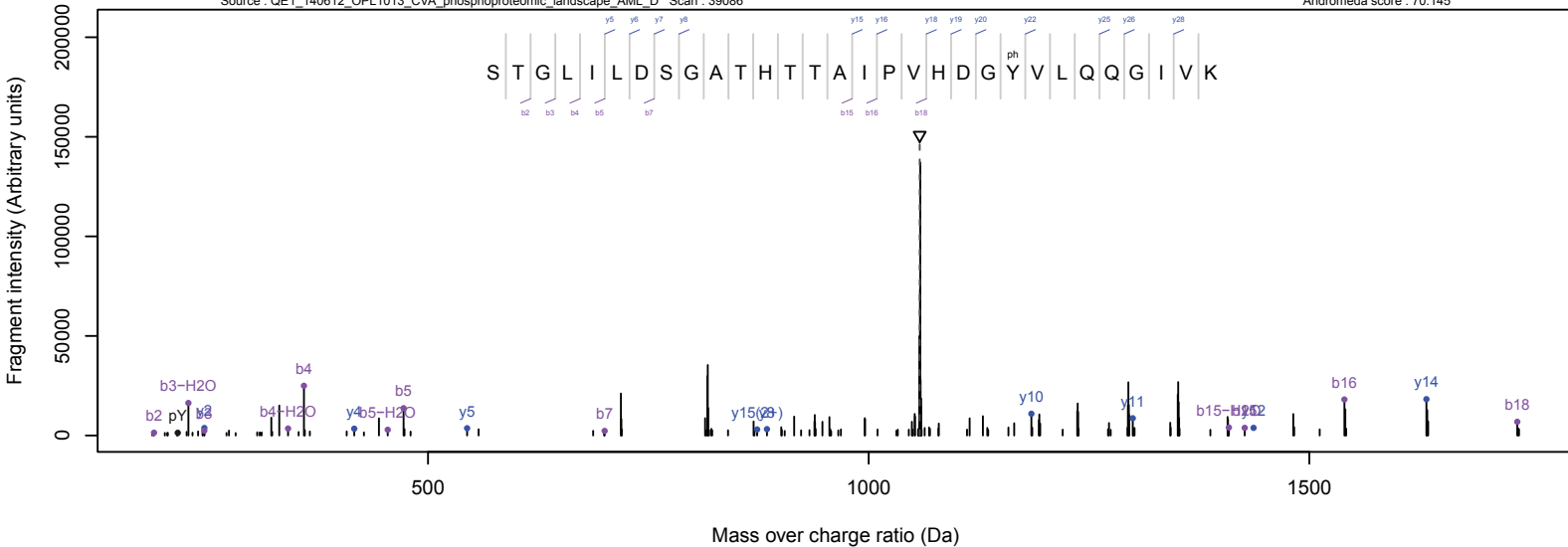
Andromeda score : 70.197



**STGLILDGATHHTTAIPVHDG(ph)VLQQGIVK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 39086

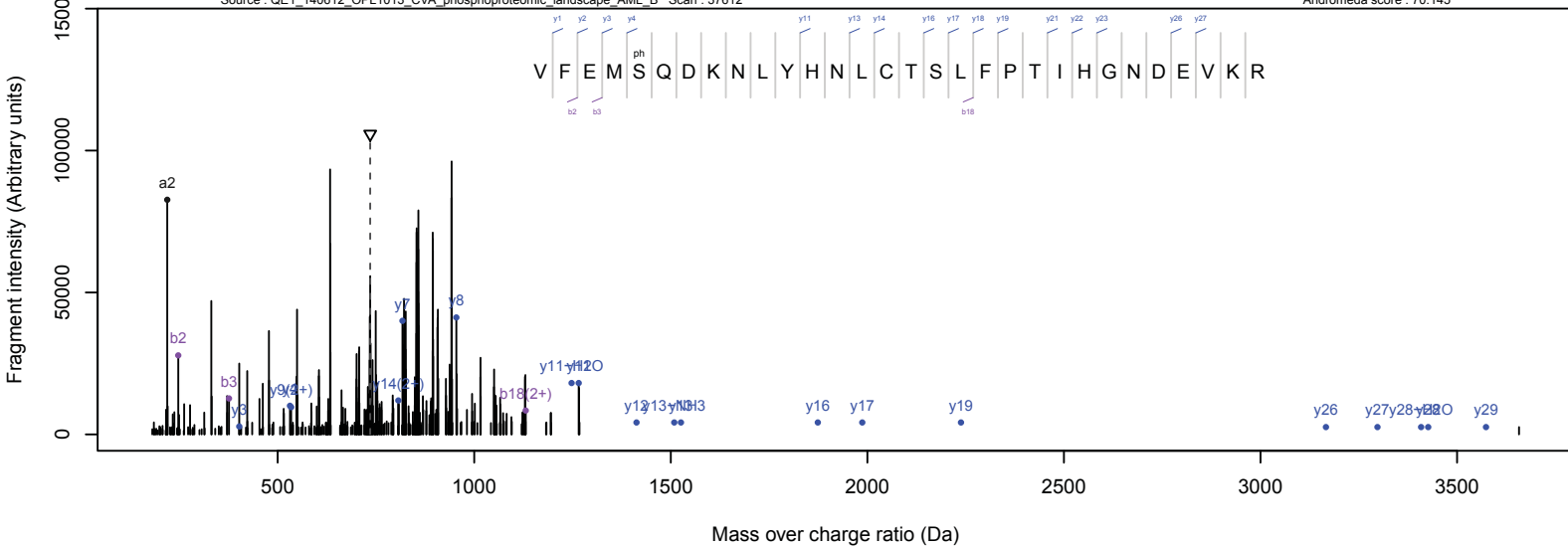
Andromeda score : 70.145



**VFEMS(ph)QDKNLYHNLCTSLFPTIHGNDVVK-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37612

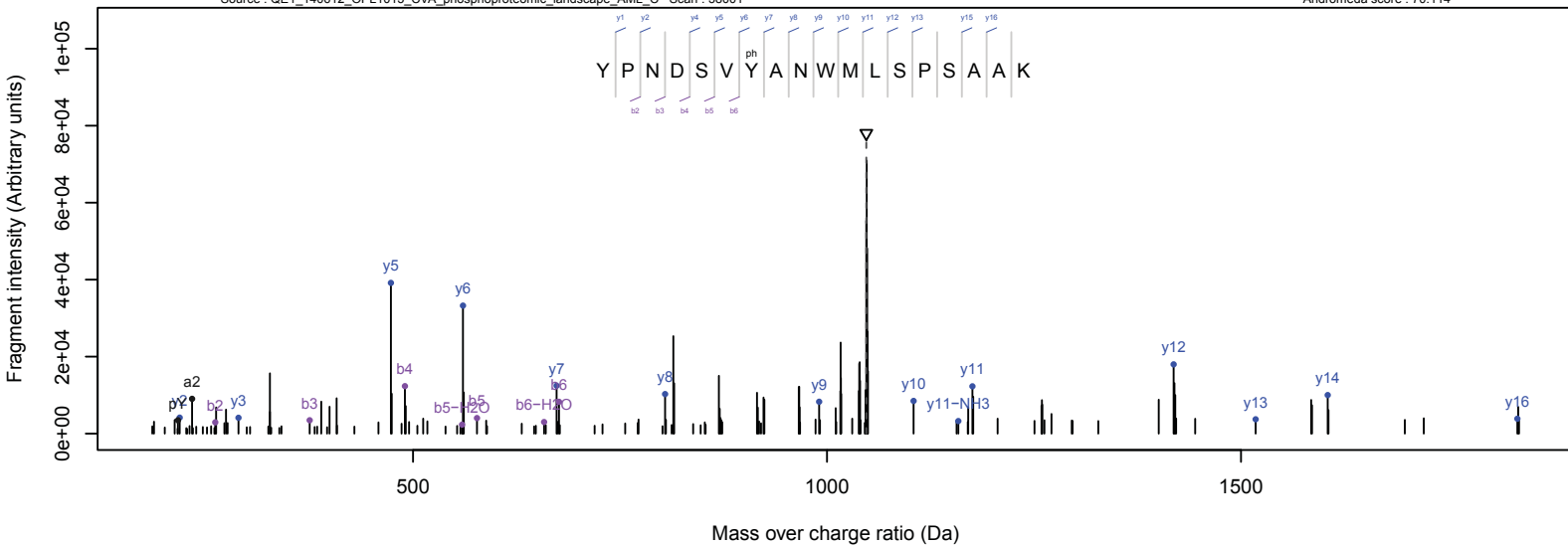
Andromeda score : 70.145



YPNDSVY(ph)ANWMLSPSAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38001

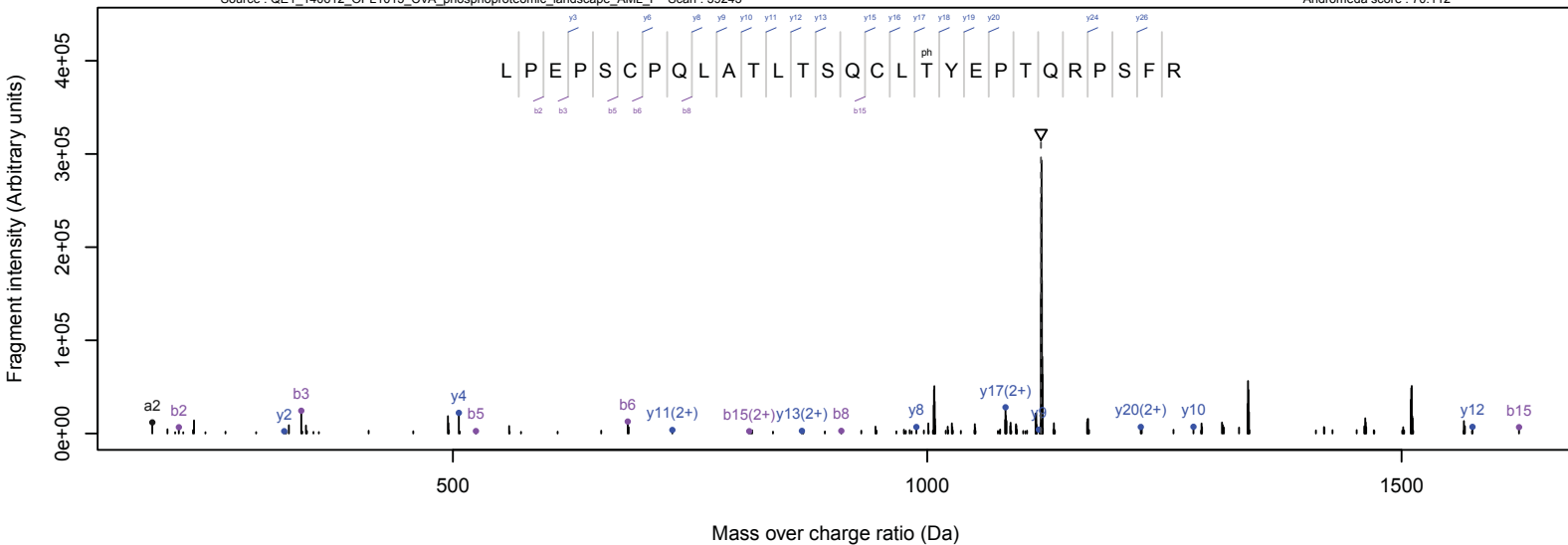
Andromeda score : 70.114



LPEPSCQLATLTSQCLT(ph)YEPTQRPSFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 39243

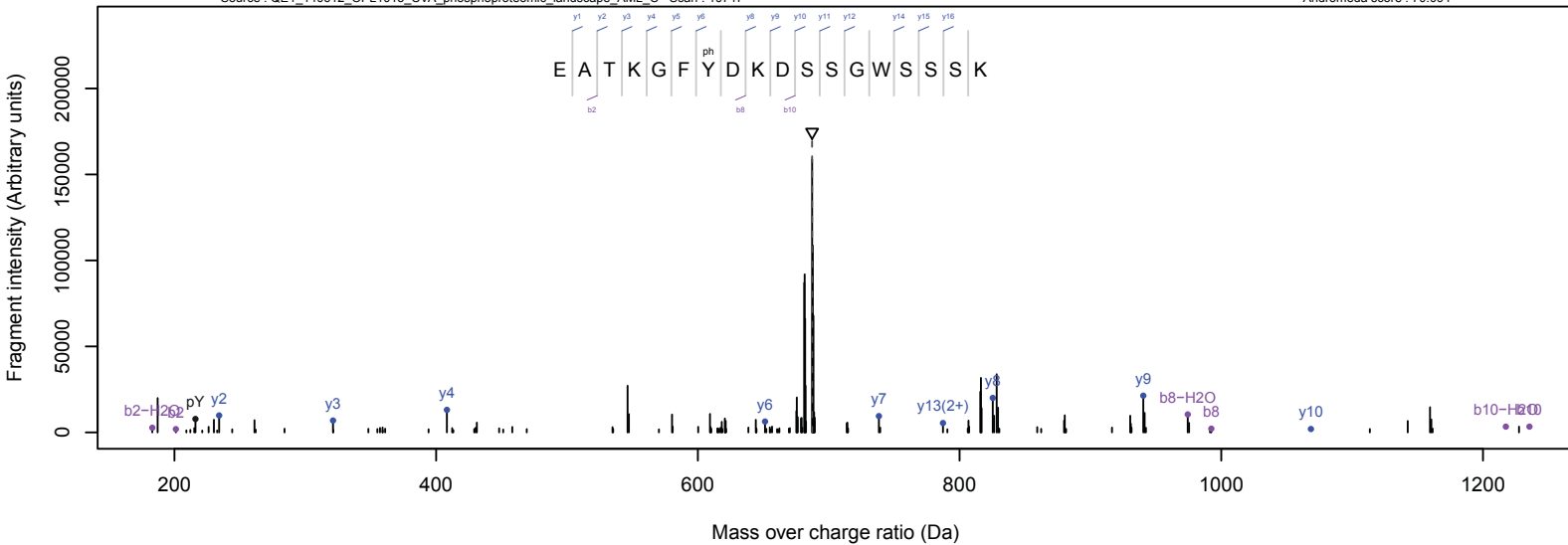
Andromeda score : 70.112

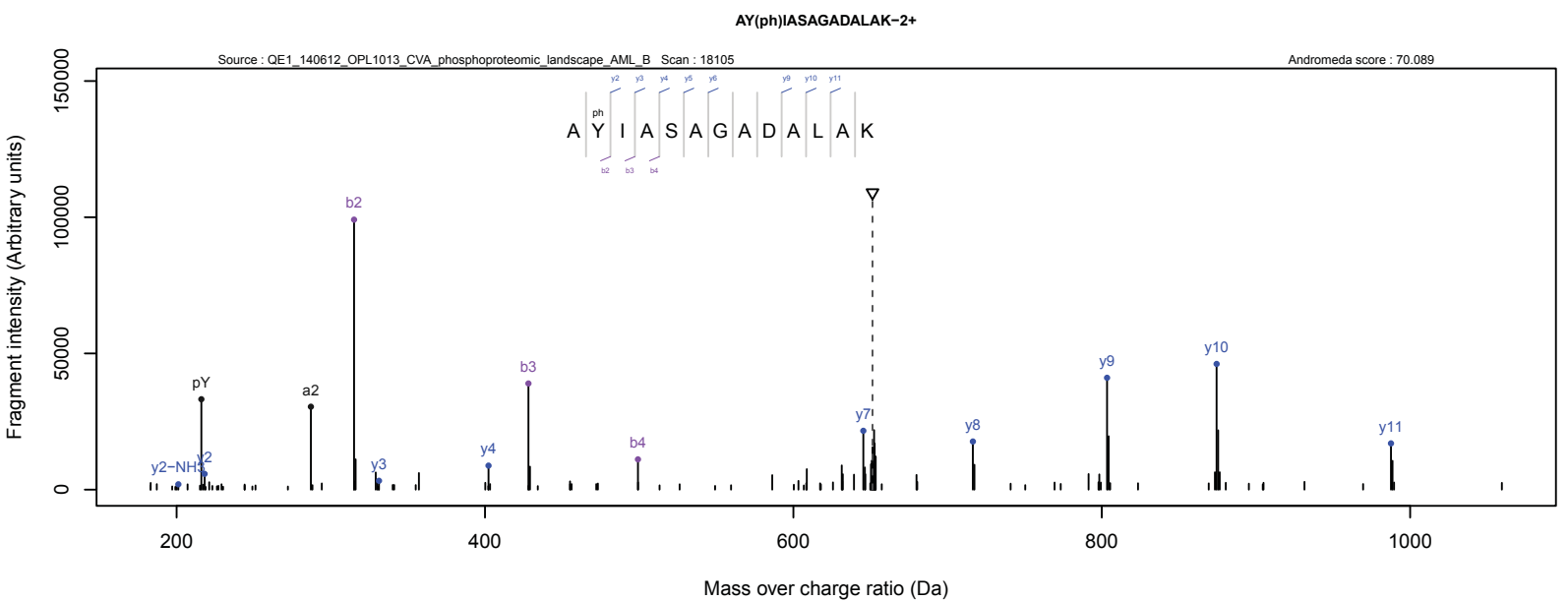
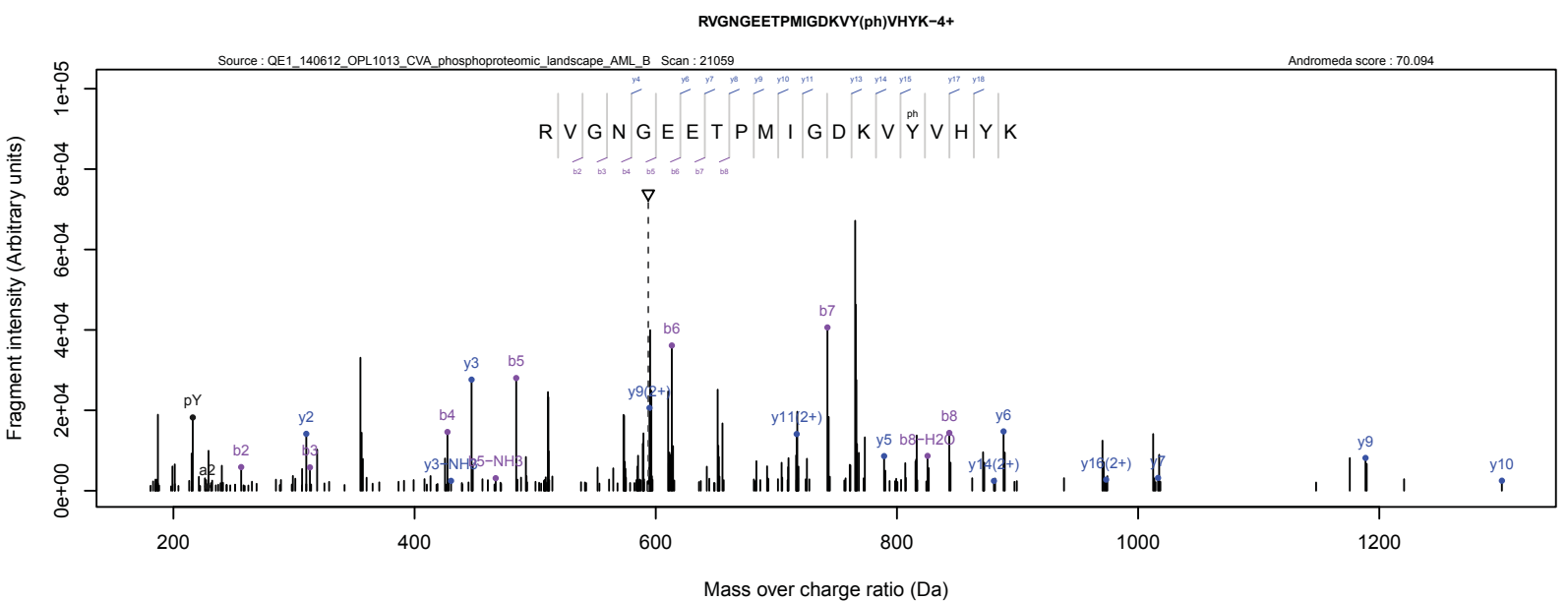
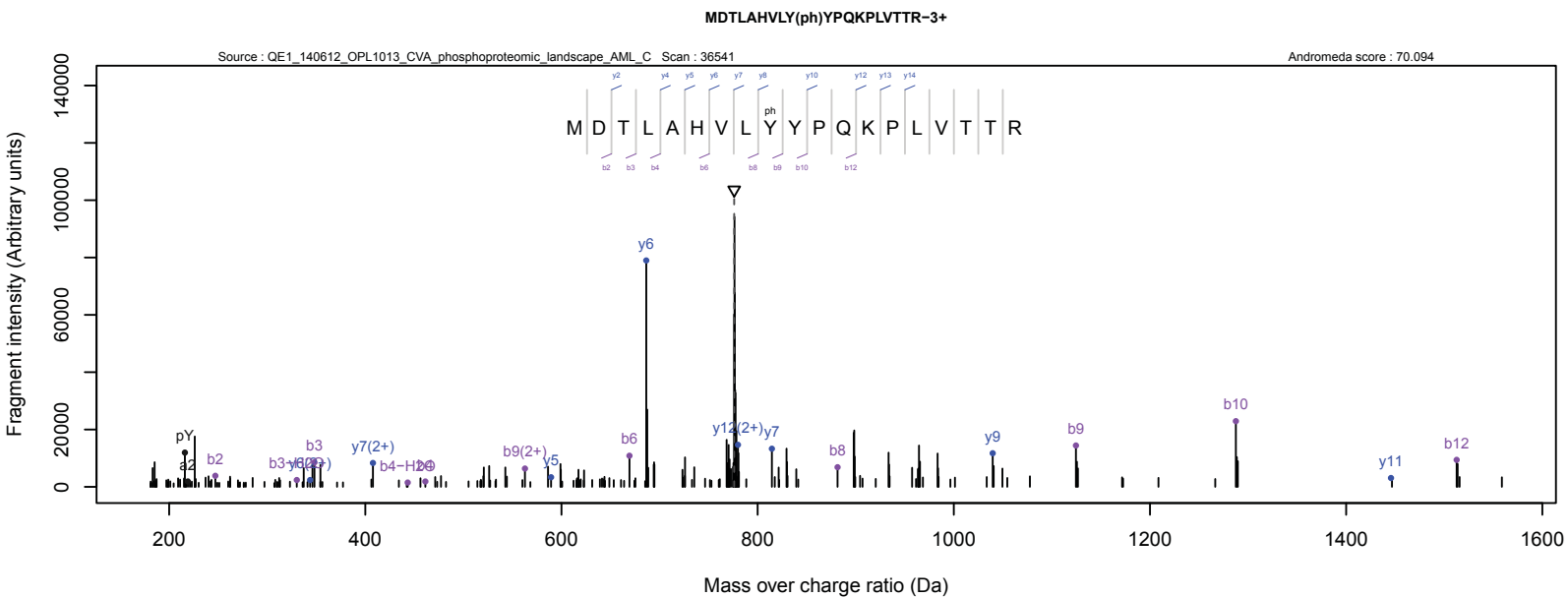


EATKGFYD(K)SSGSWSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10747

Andromeda score : 70.094



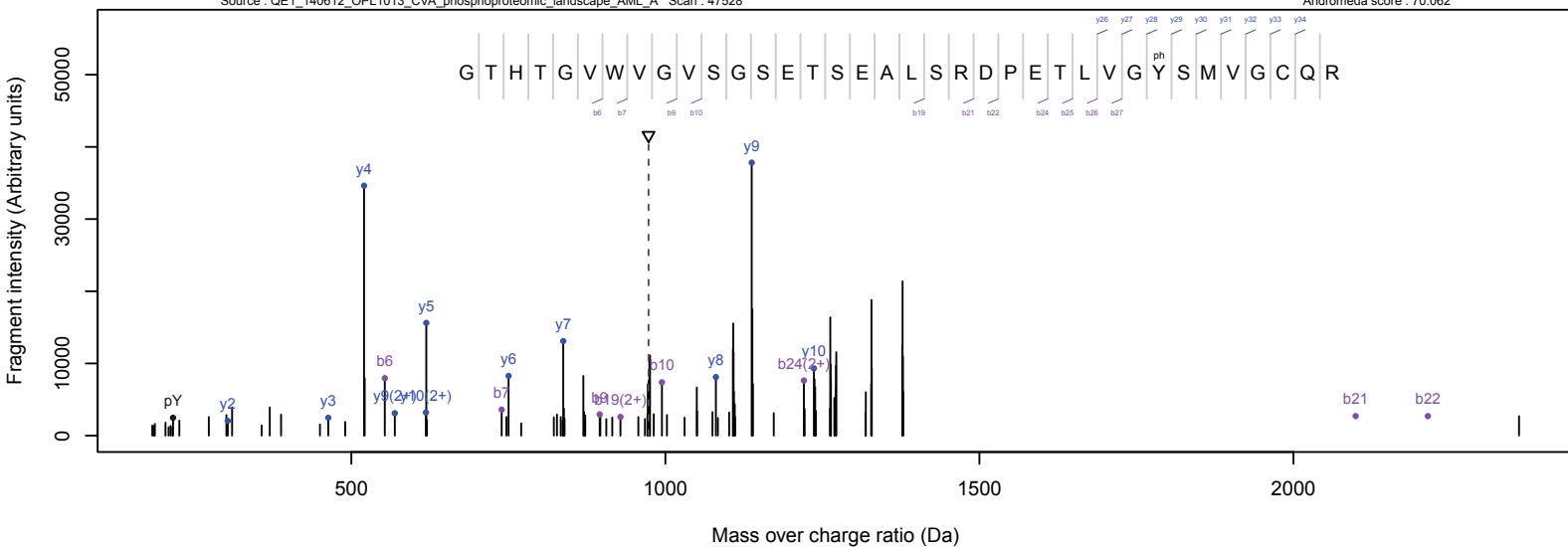




GTHTGVWVGSGETSEALSRDPETLVGYPHSMVGCQR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 47528

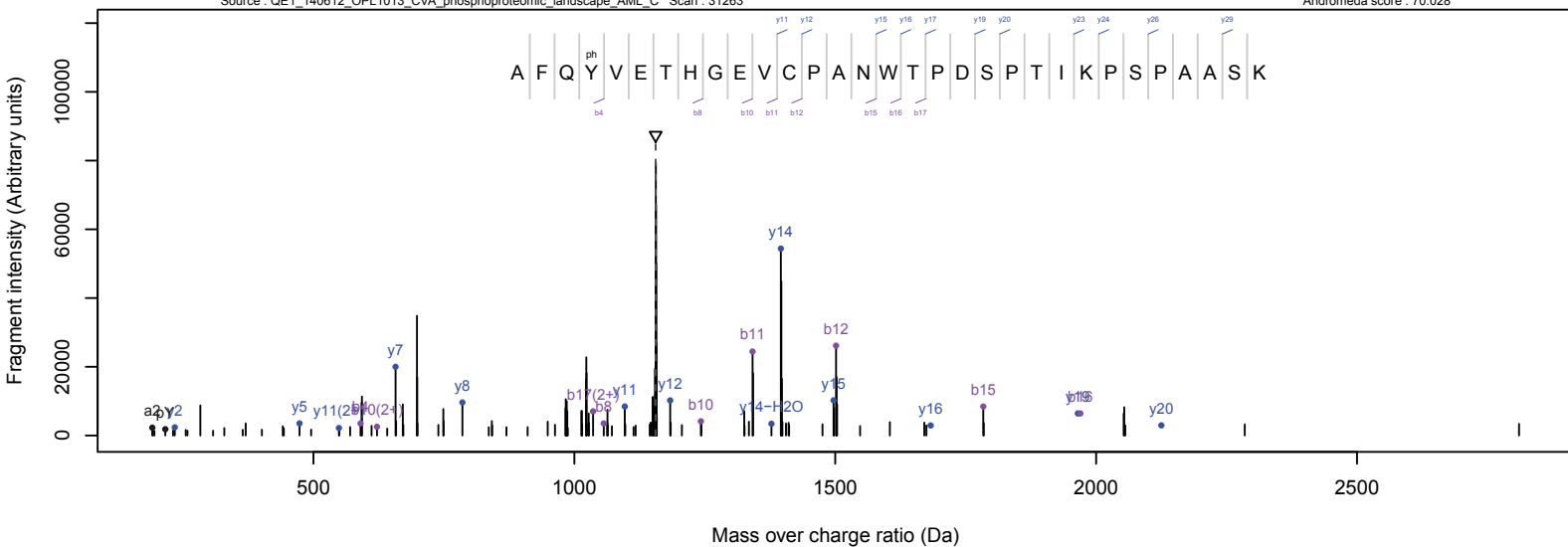
Andromeda score : 70.062



AFQY(V)ETHGEVCPANWTPDPTIKPSPAASK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31263

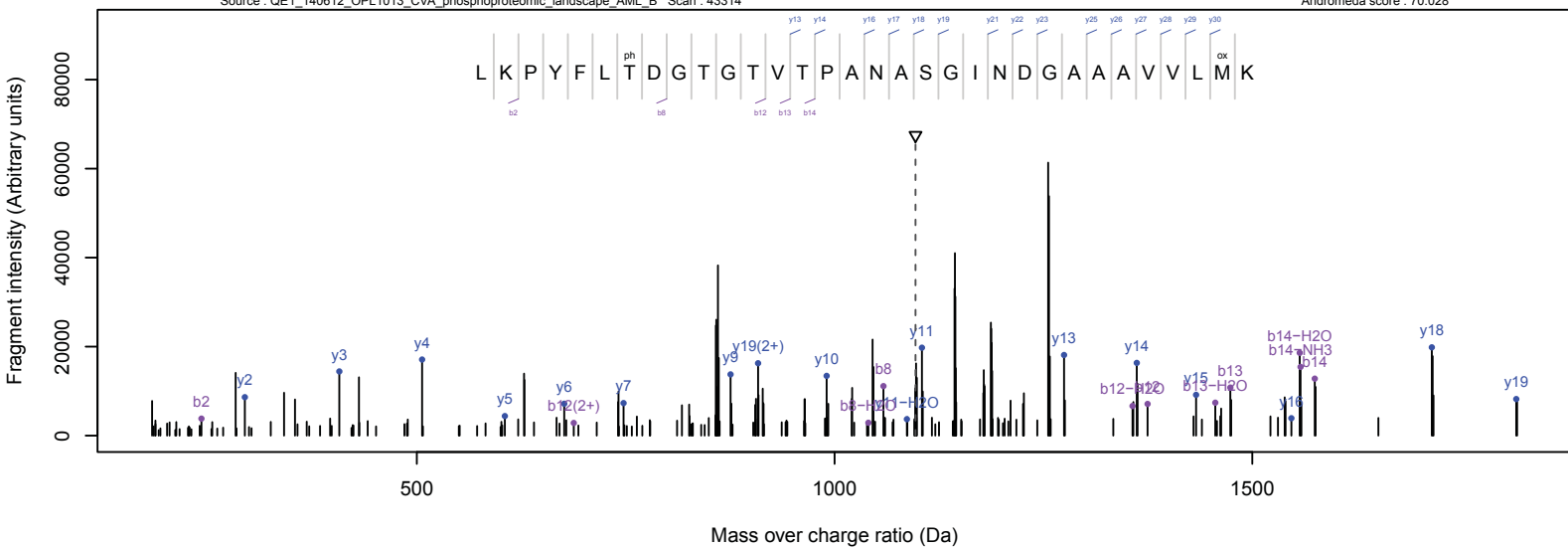
Andromeda score : 70.028



## LKP YFL(ph)DGTGTVTPANASGINDGAAAVVLM(ox)K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43314

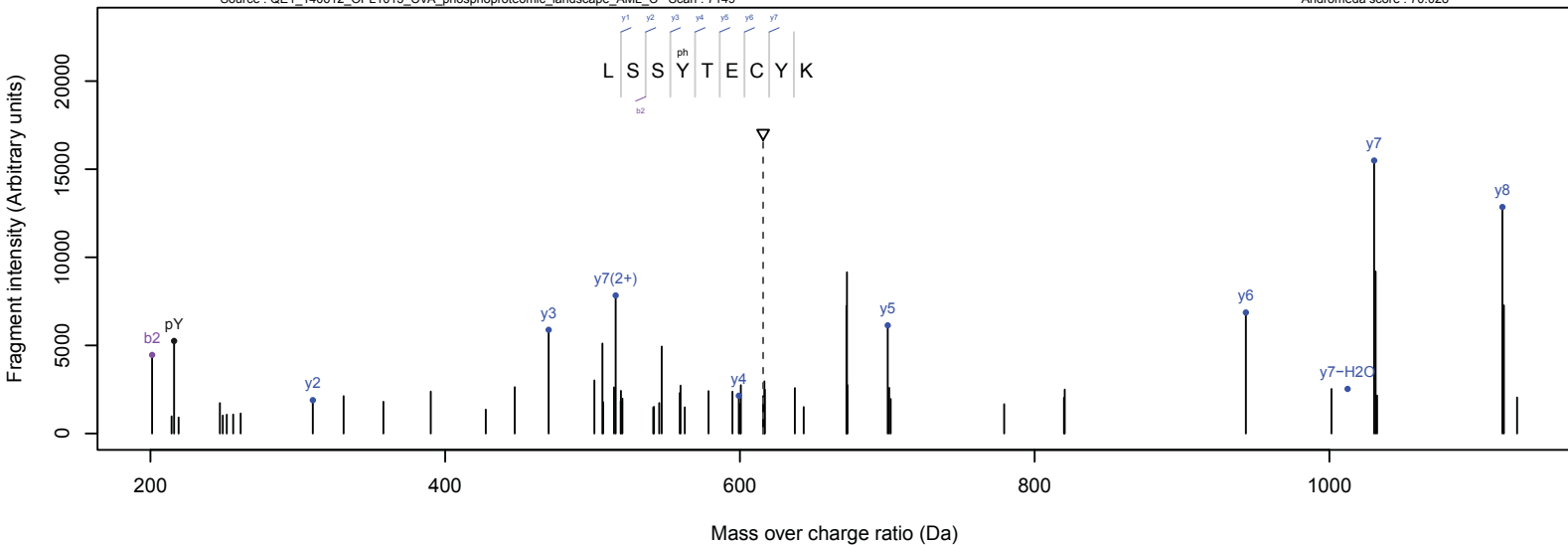
Andromeda score : 70.028



## LSS Y(ph)TECYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7149

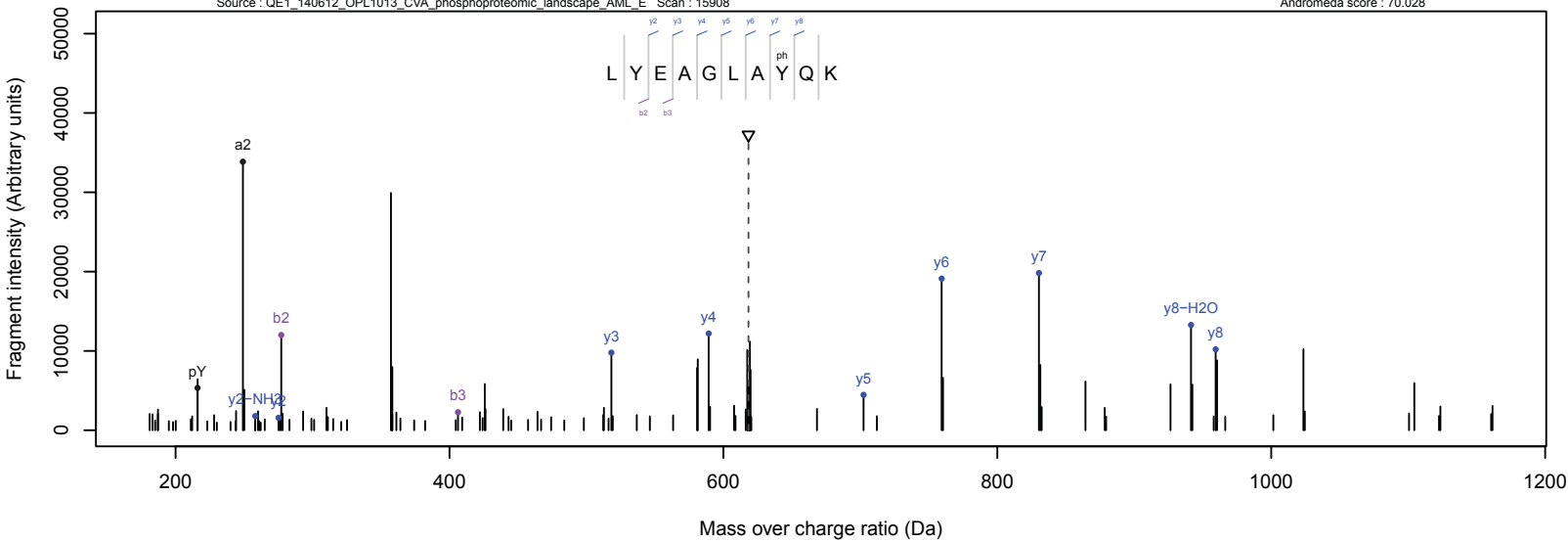
Andromeda score : 70.028



## LYEAGL(ph)YQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 15908

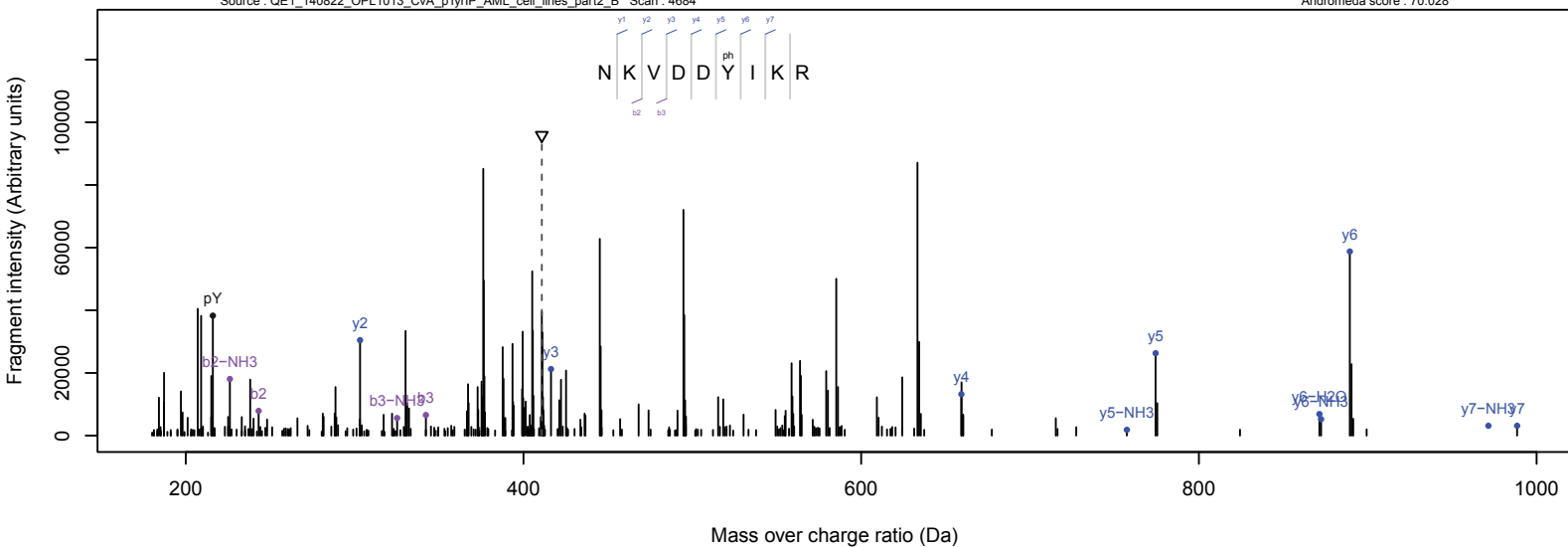
Andromeda score : 70.028



## NKVDDY(ph)IKR-3+

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 4684

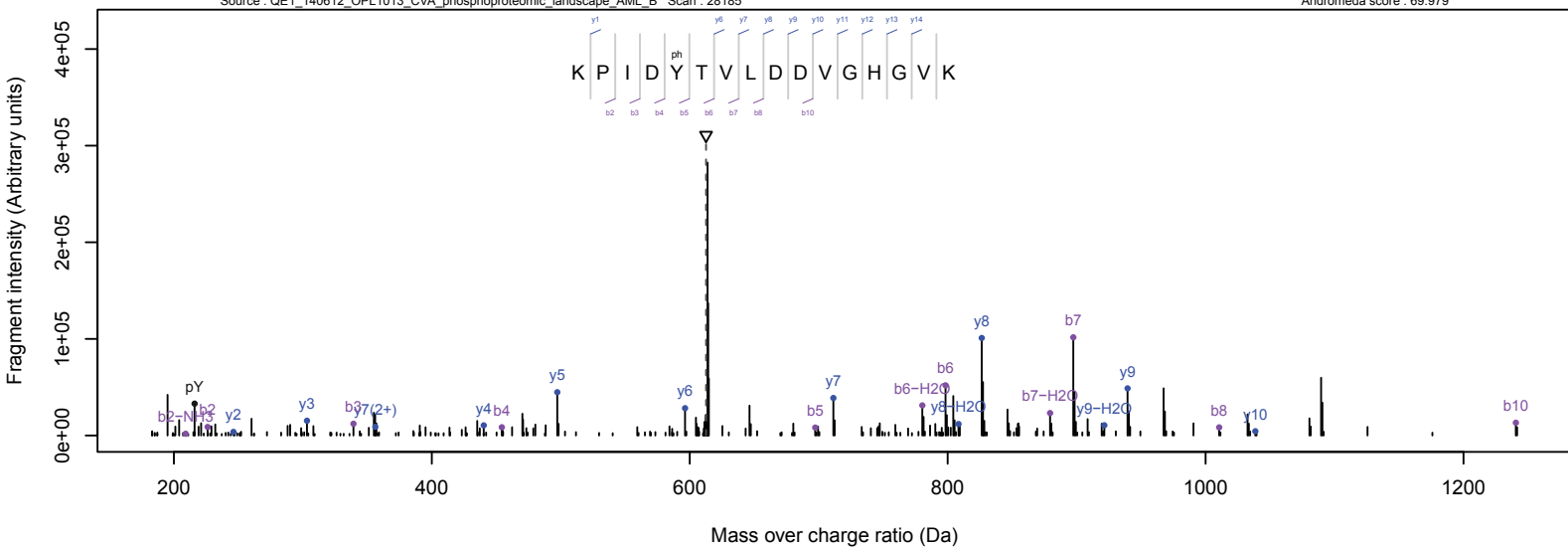
Andromeda score: 70.028



## KPIDY(ph)TVLDDVGHGVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 28185

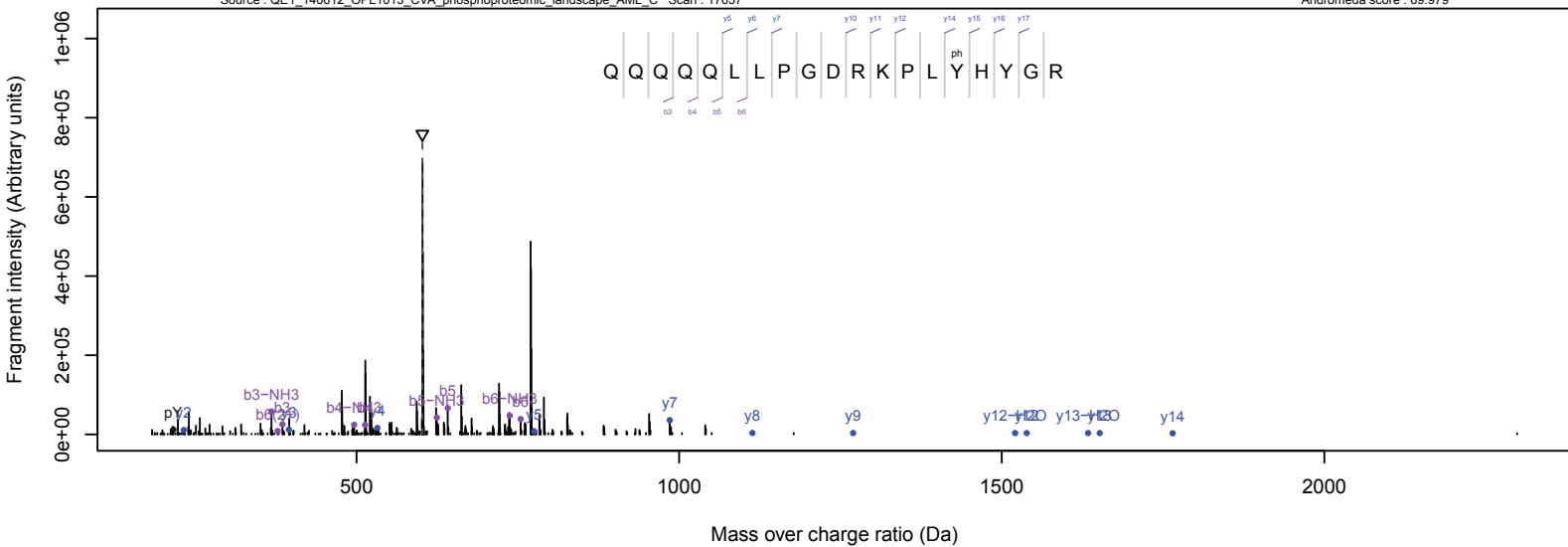
Andromeda score: 69.979



## QQQQLLPGDRKPLYHYGR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17637

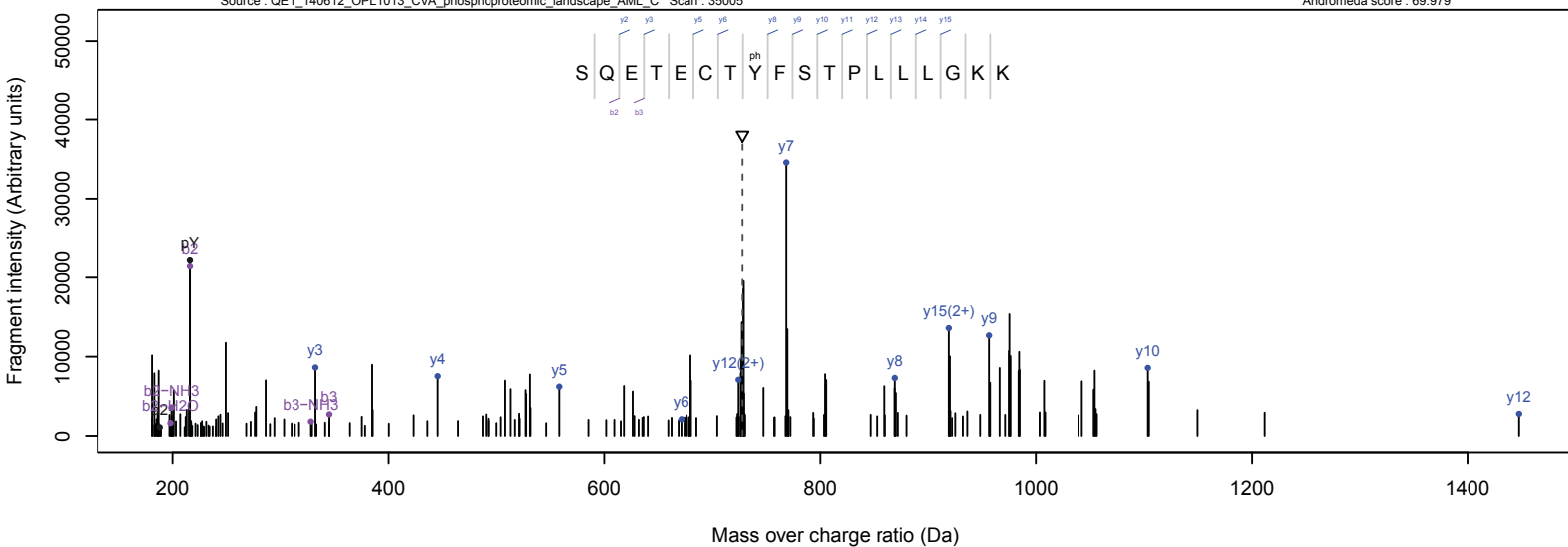
Andromeda score: 69.979



## SQETECTY(ph)FSTPLLLGKK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 35005

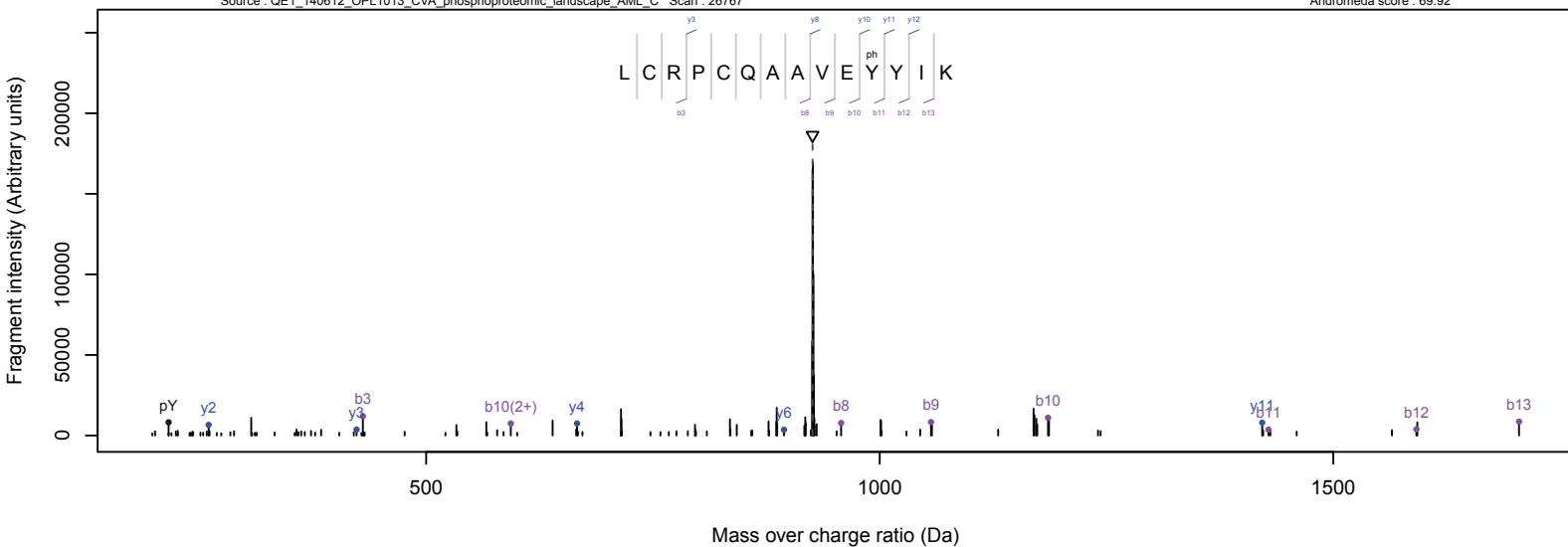
Andromeda score: 69.979



## LCRPCQAAVEY(ph)YIK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26767

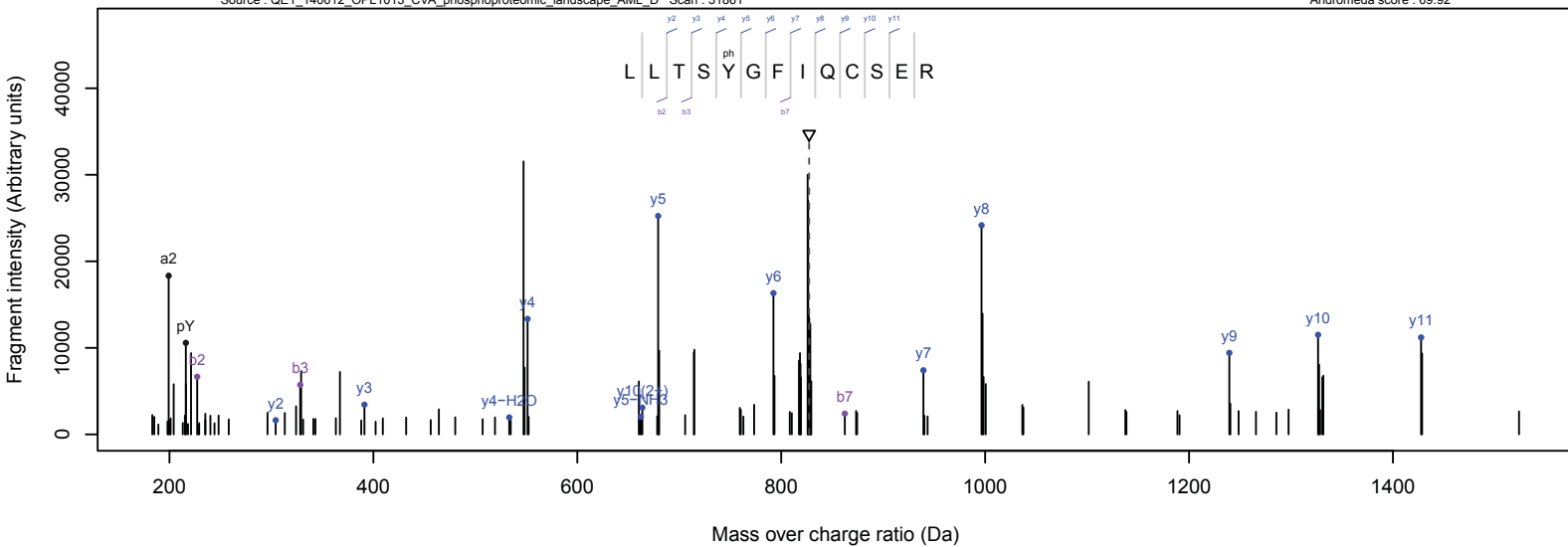
Andromeda score: 69.92



## LLTSY(ph)GFIQCSEK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 31861

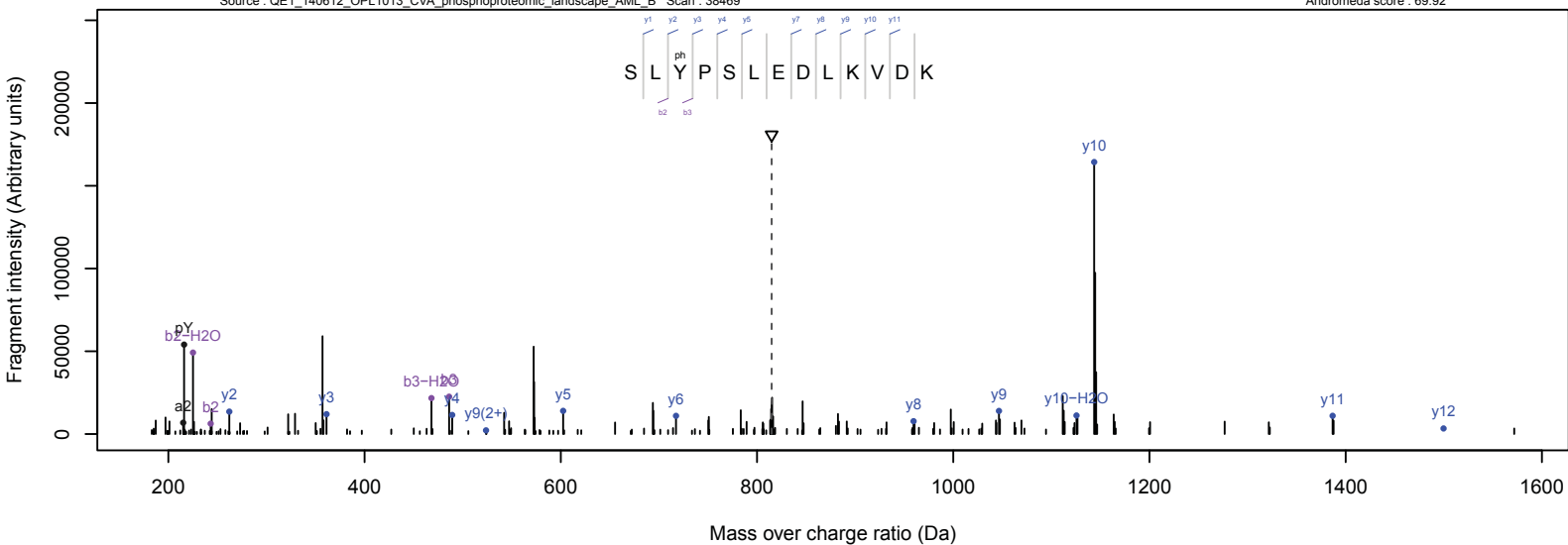
Andromeda score: 69.92



## (ac)SLY(ph)PSLEDLKVDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38469

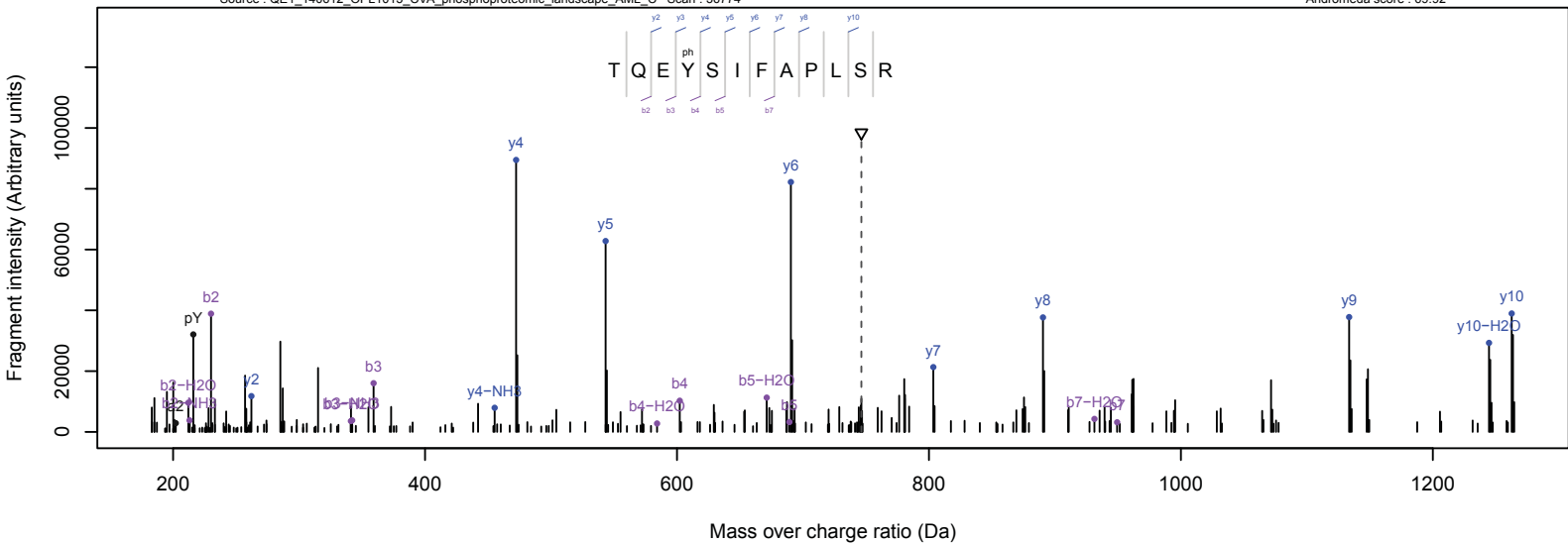
Andromeda score : 69.92



## TQEY(ph)SIFAPLSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36774

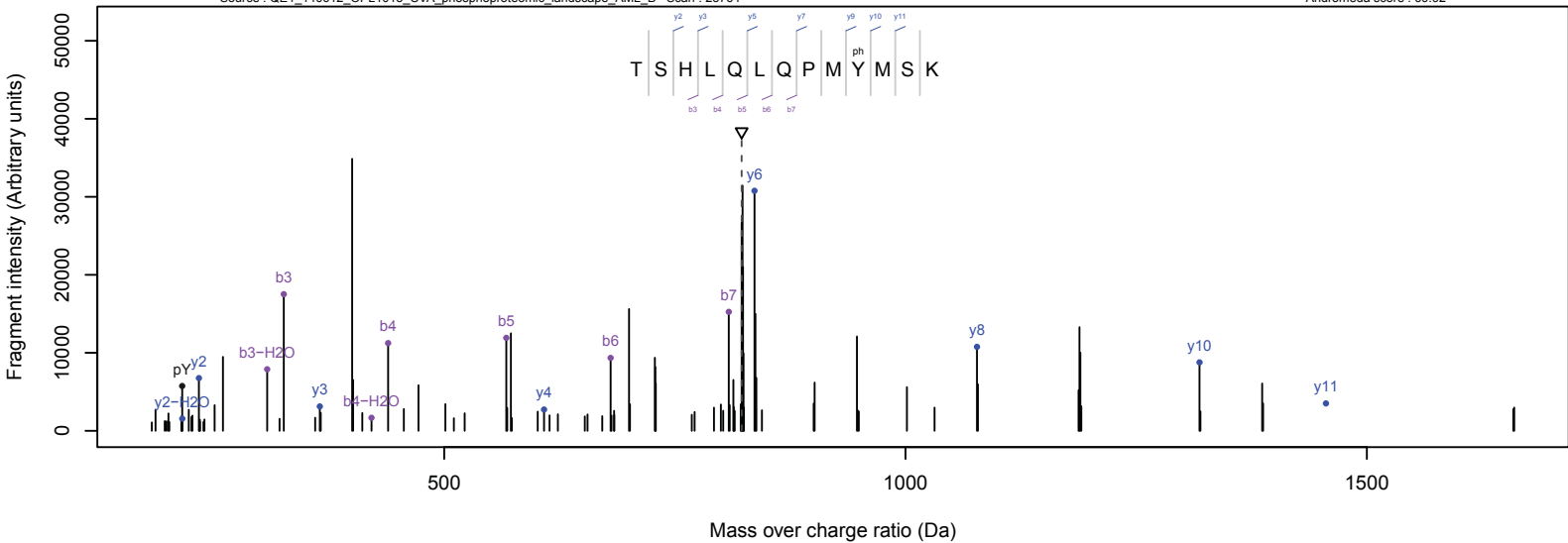
Andromeda score : 69.92



## TSHLQLQPMY(ph)MSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23764

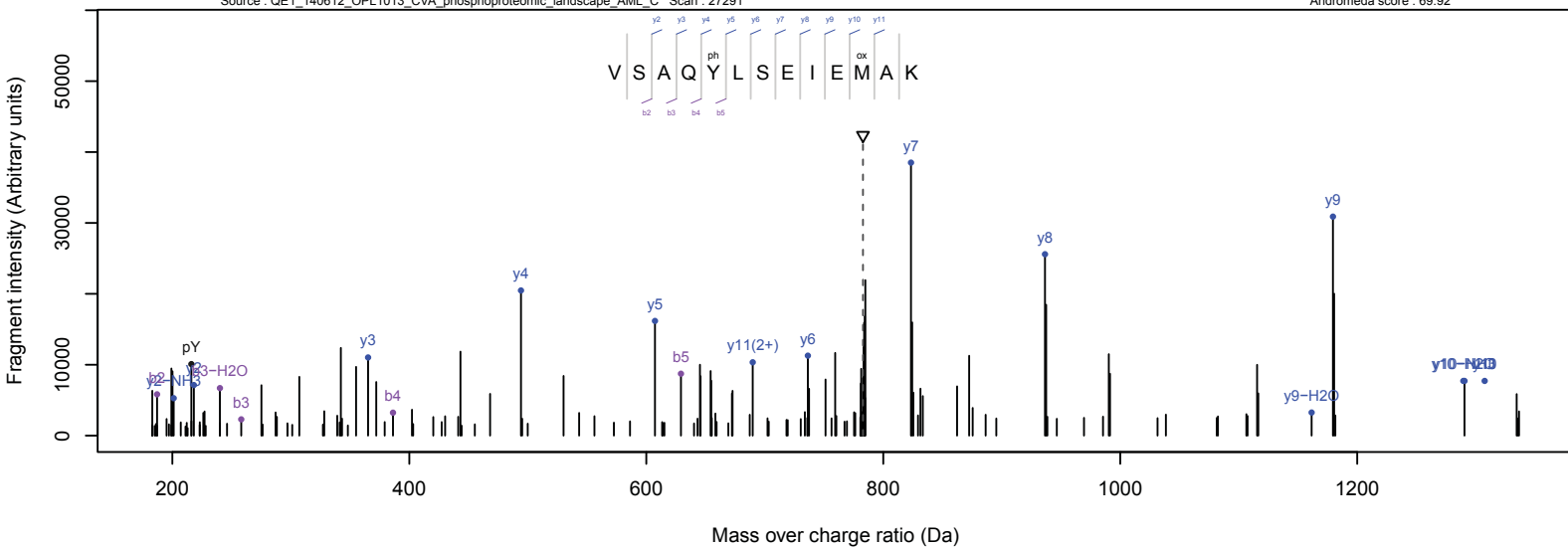
Andromeda score : 69.92



## VSAQY(ph)LSEIEM(ox)AK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27291

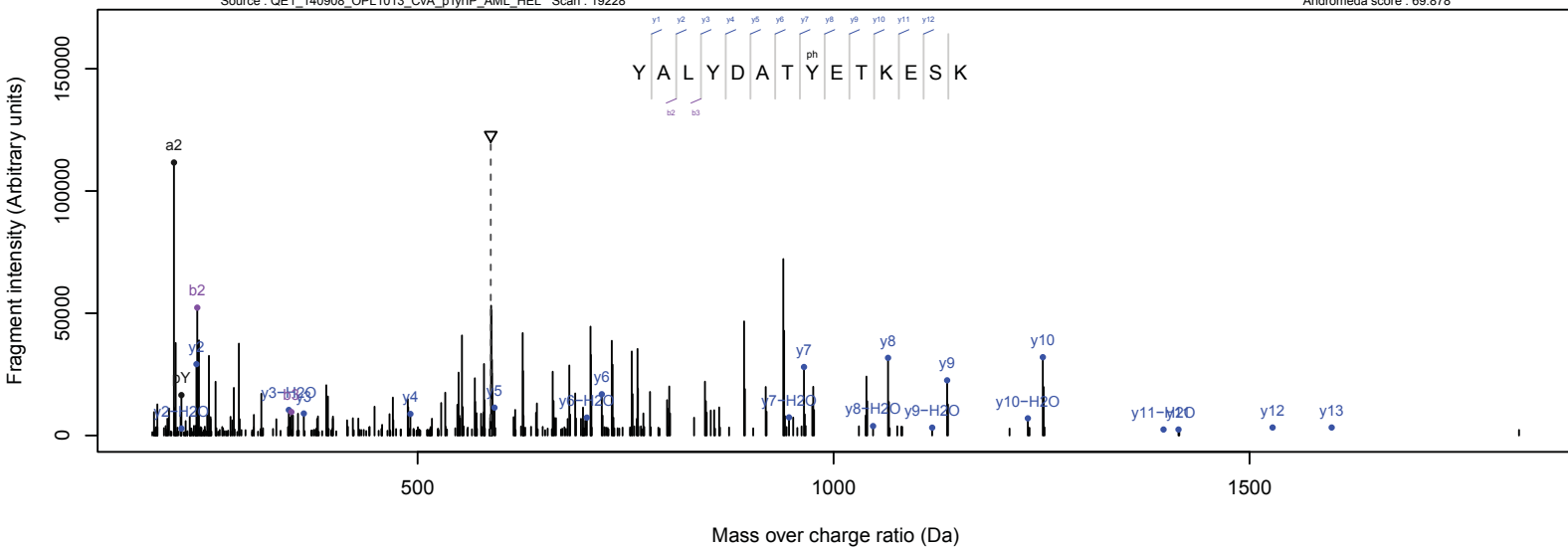
Andromeda score : 69.92



## YALYDATY(ph)ETKESK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 19228

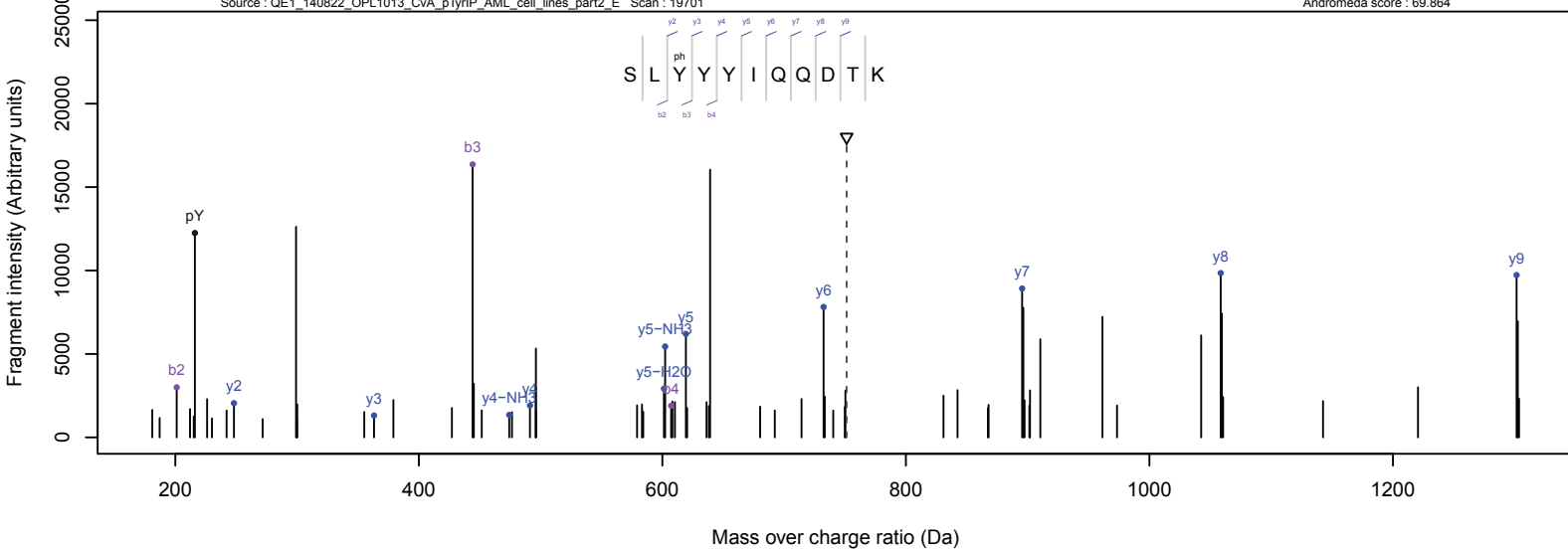
Andromeda score : 69.878



## SLY(ph)YIQQDTK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 19701

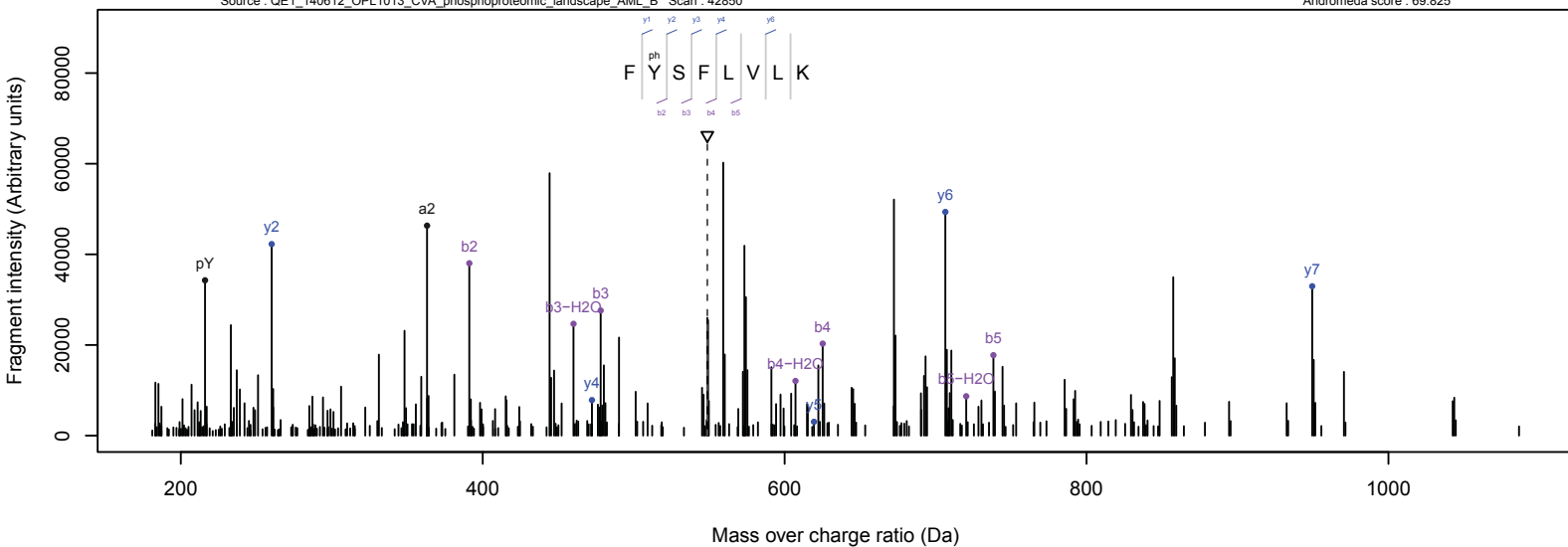
Andromeda score : 69.864



## FY(ph)SFLVLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42850

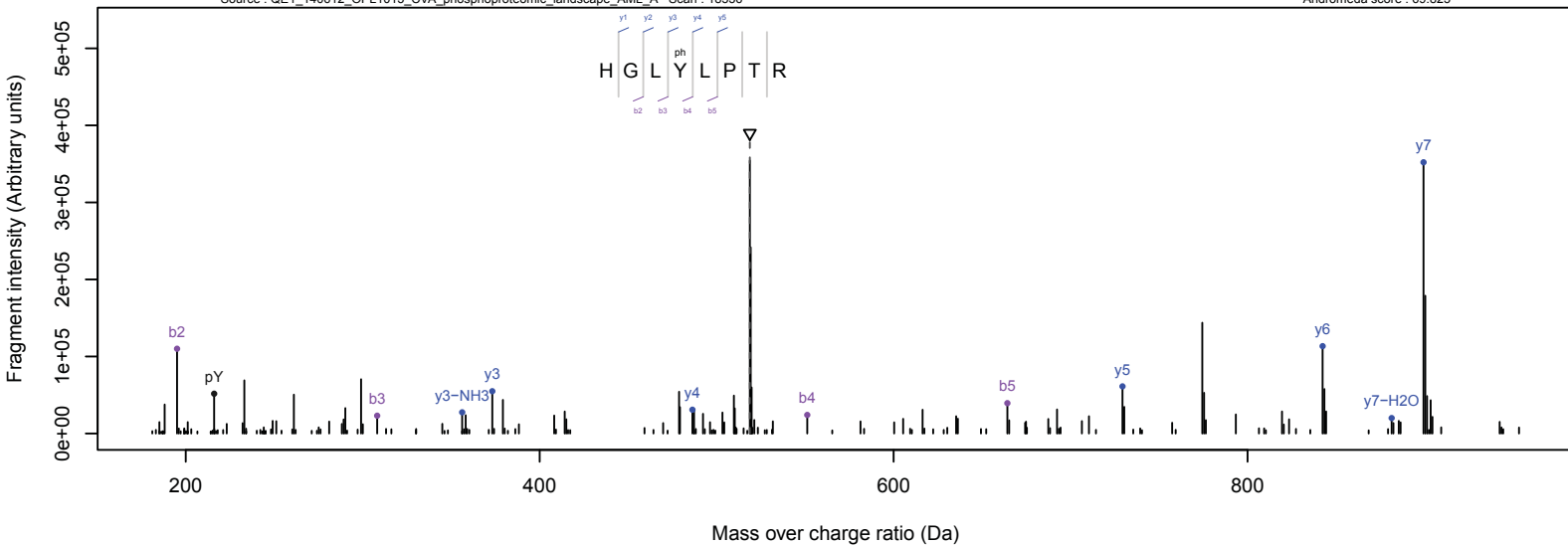
Andromeda score : 69.825



## HGLY(ph)LPTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 18536

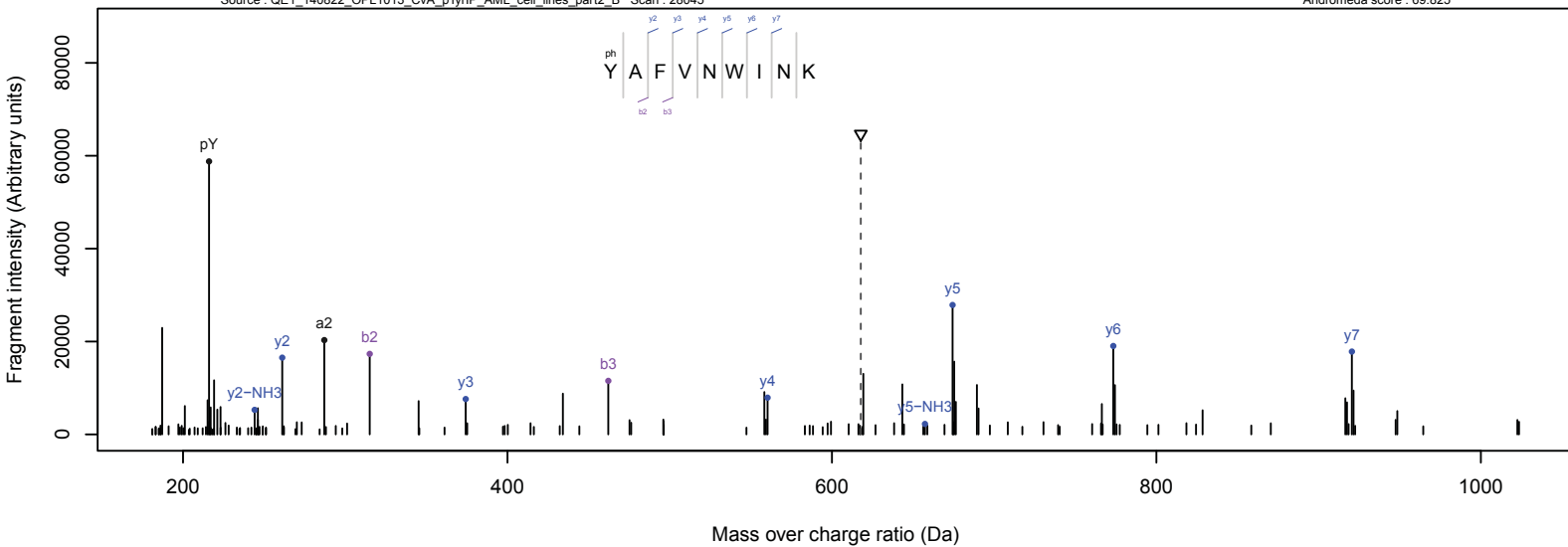
Andromeda score : 69.825



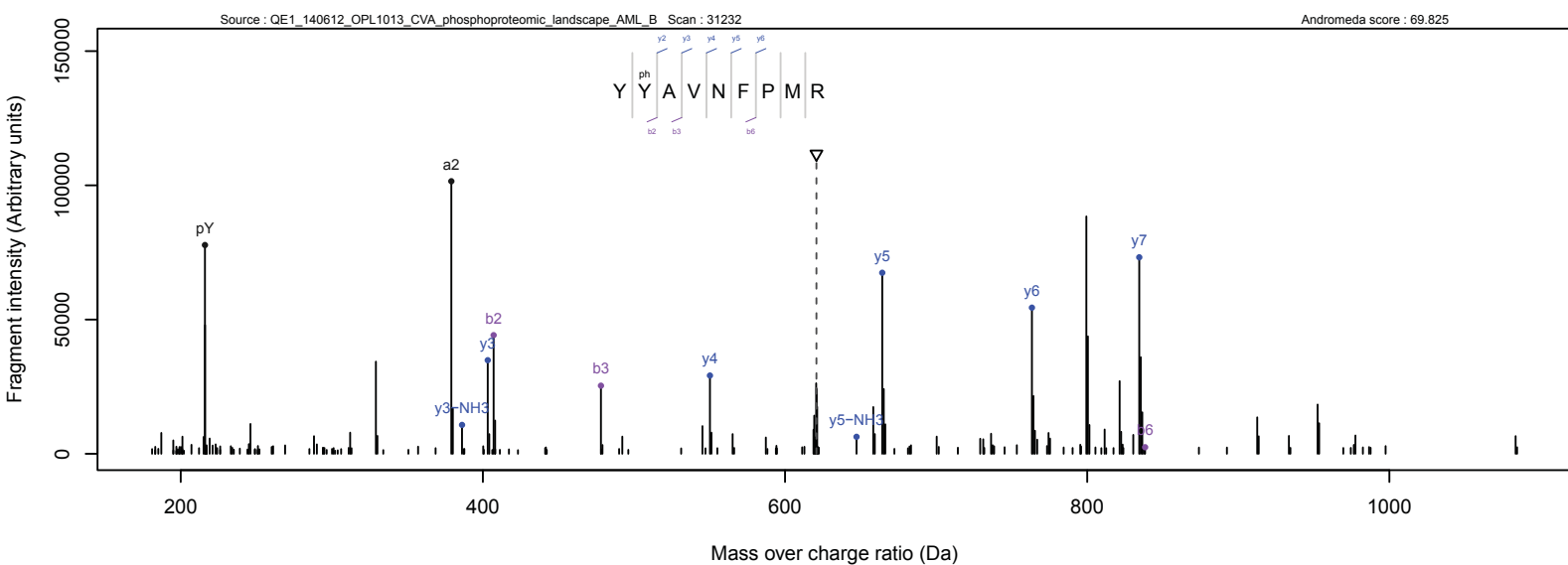
## Y(ph)AFVNWINK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 28045

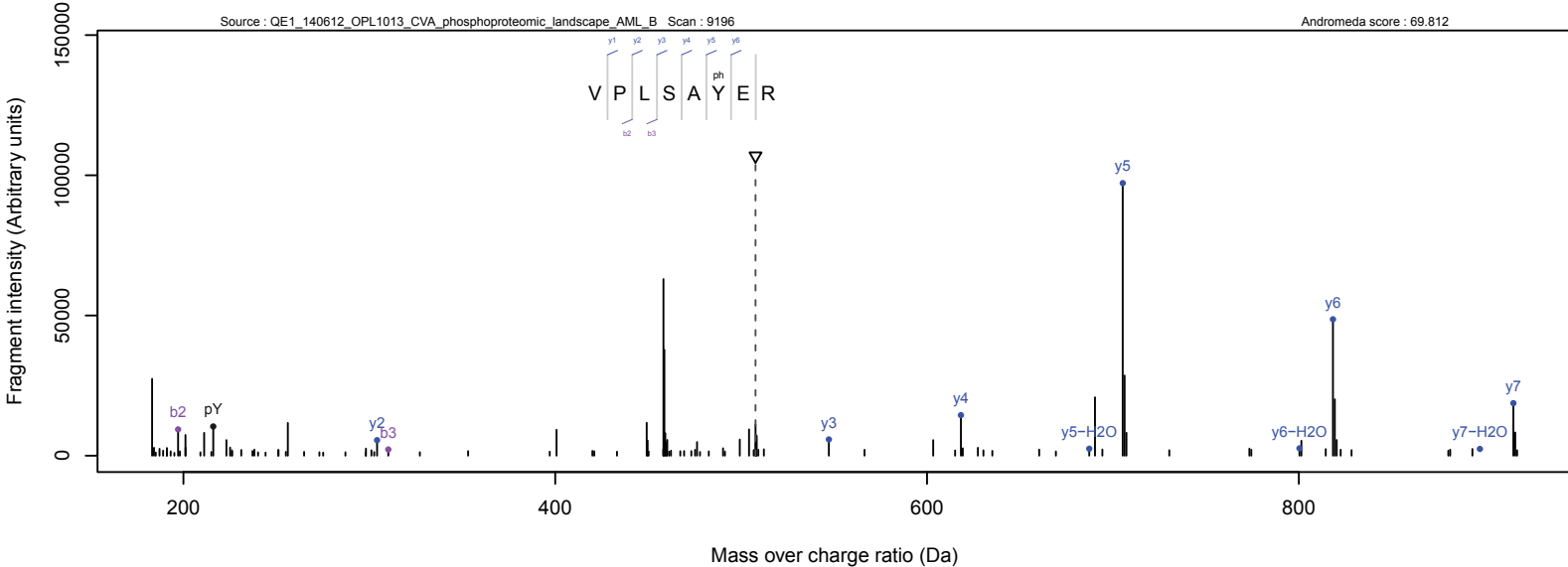
Andromeda score : 69.825



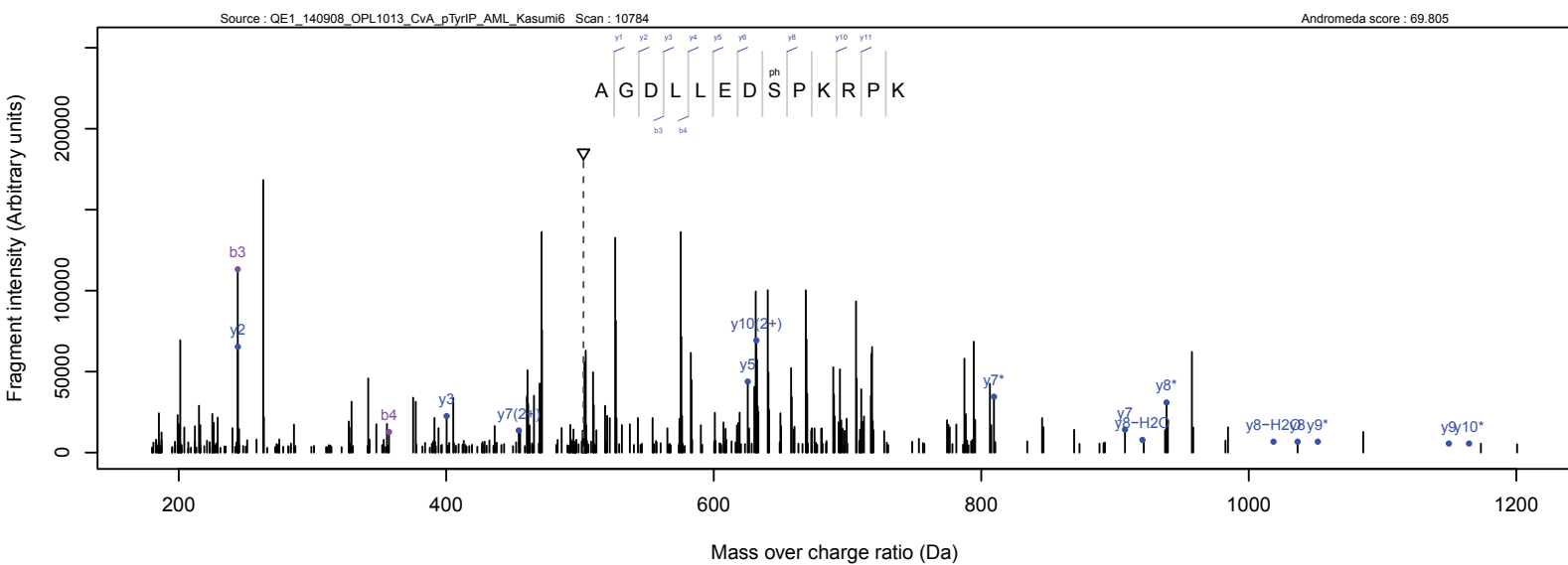
## YY(ph)AVNFPMR-2+



## VPLSAY(ph)ER-2+



## AGDLEDS(ph)PKRPK-3+

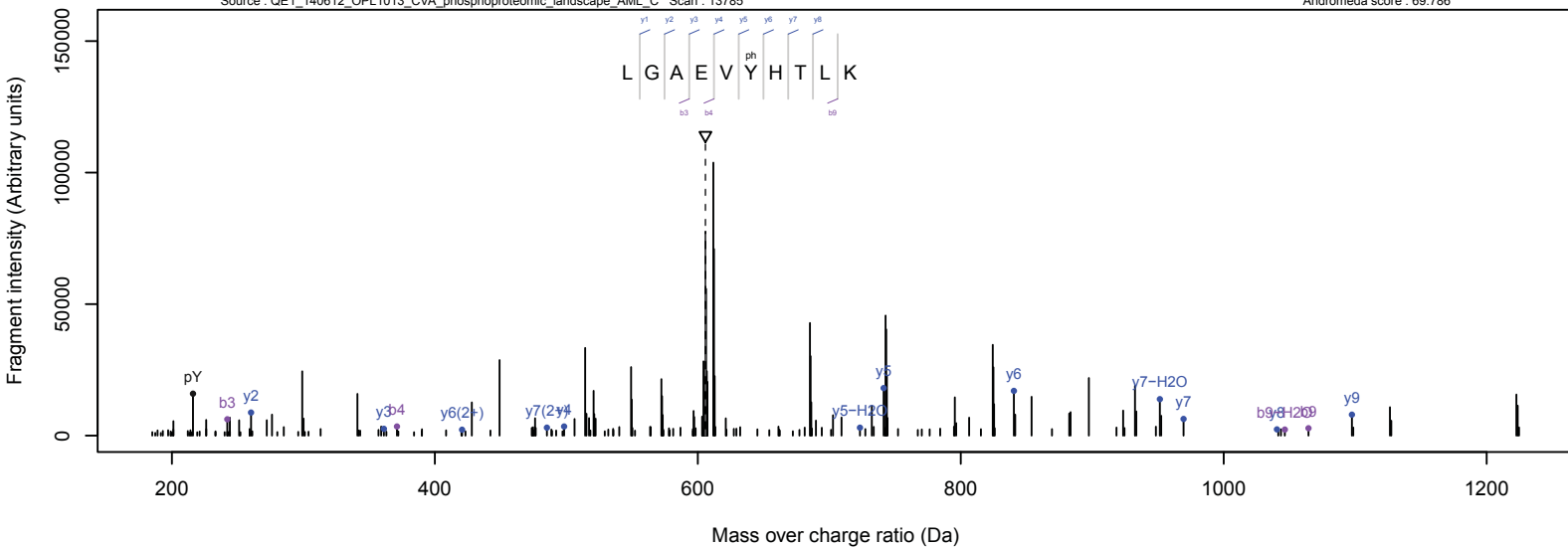




**LGAEVY(ph)HTLK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13785

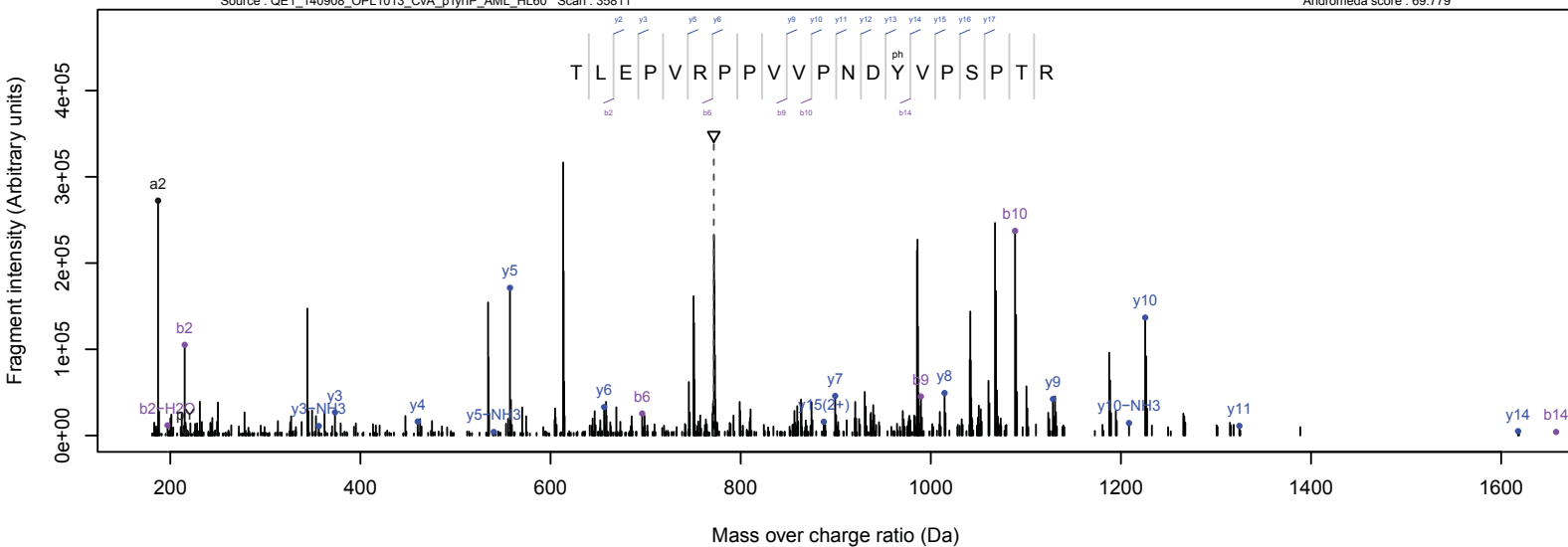
Andromeda score : 69.786



**TLEPVRPPVNDY(ph)VPSPTR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 35811

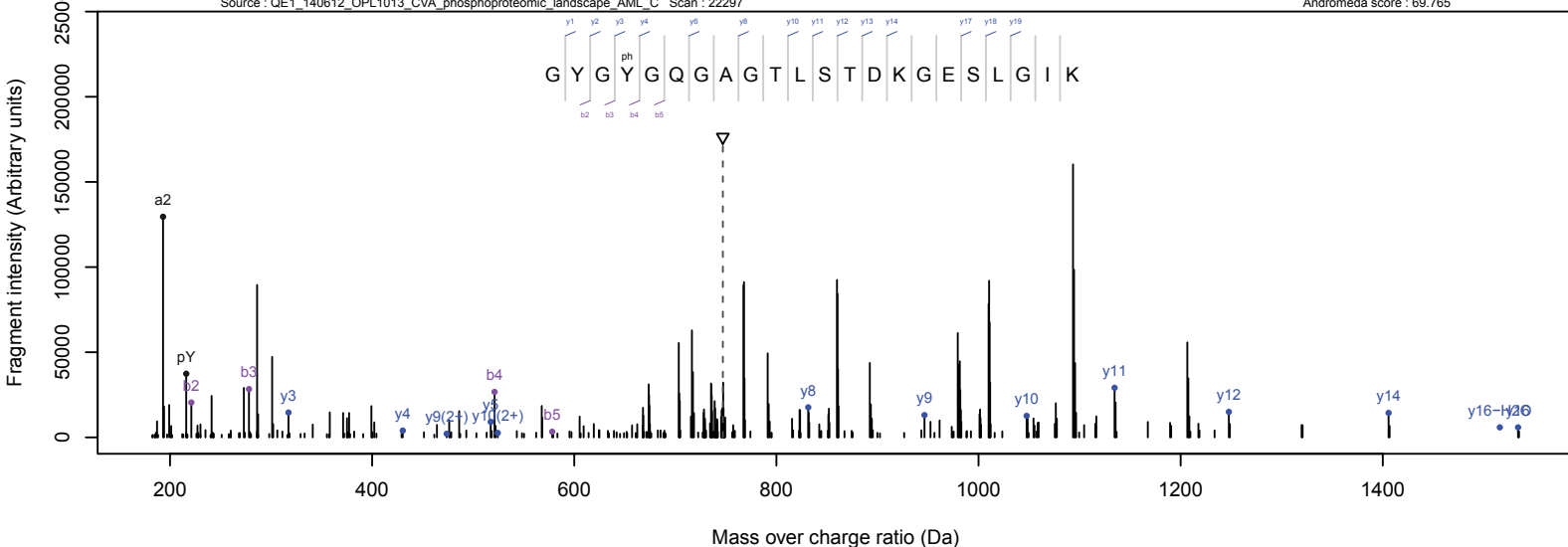
Andromeda score : 69.779



**GYGY(ph)GQGAGTLSTDKGESLGIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22297

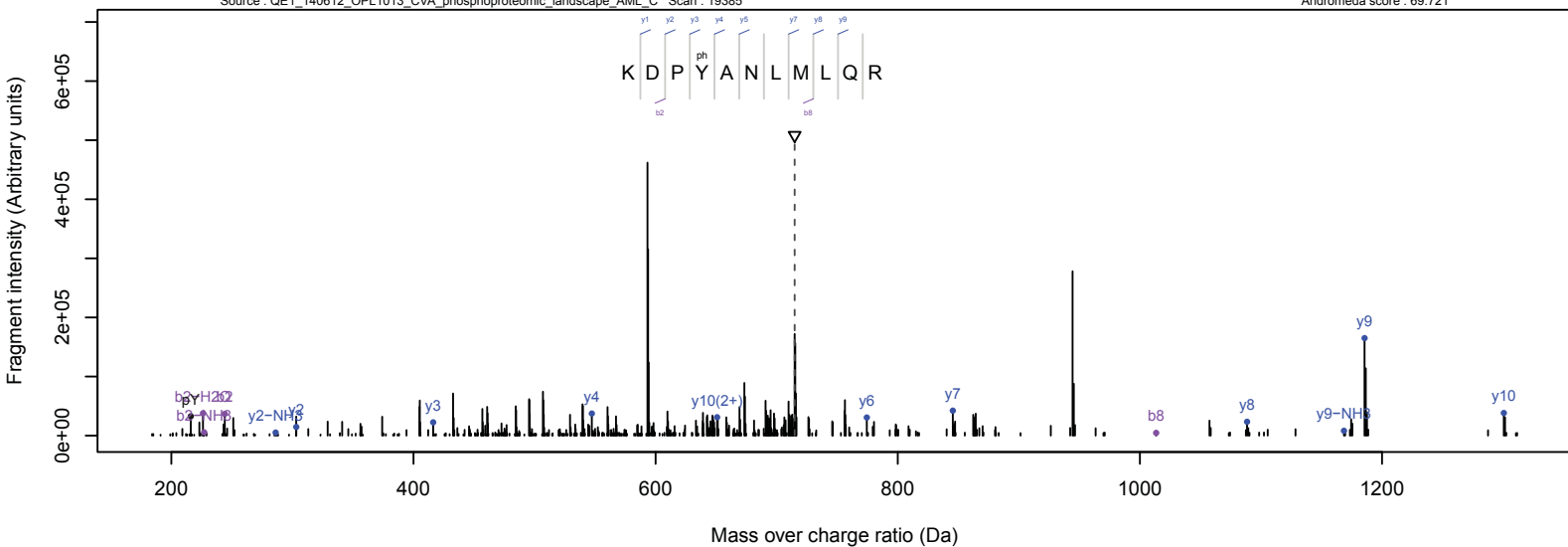
Andromeda score : 69.765



## KDPY(ph)ANMLQR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19385

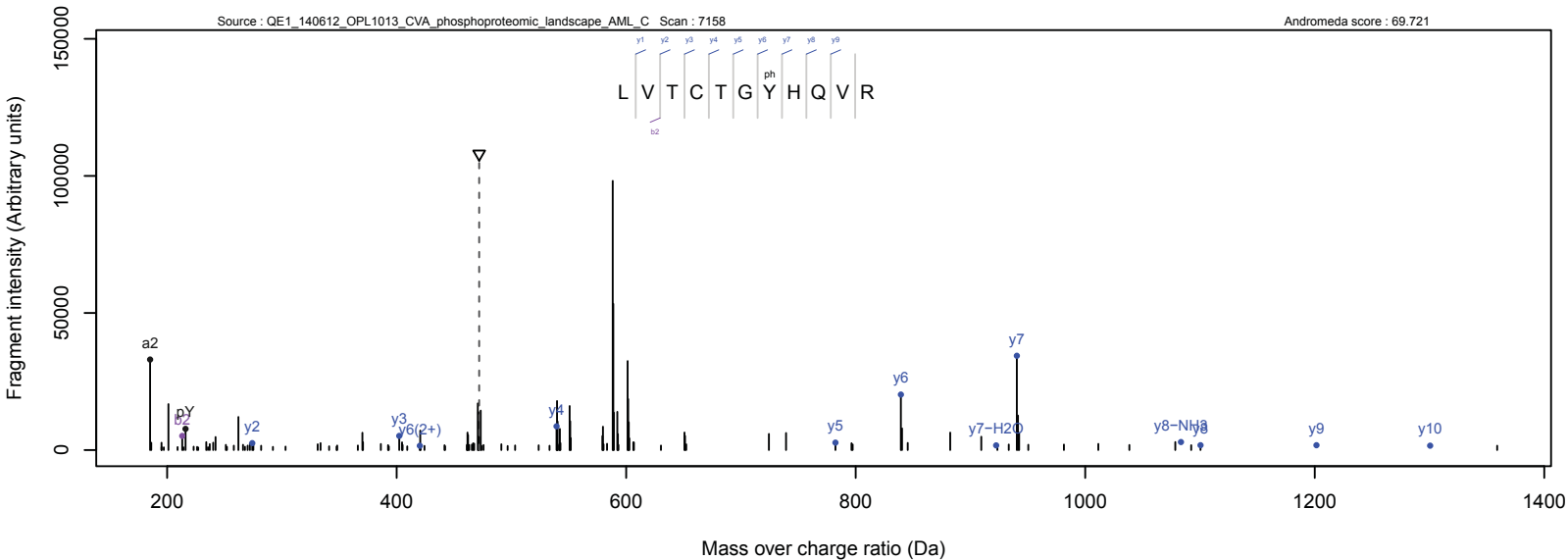
Andromeda score: 69.721



## LVTCTGY(ph)HQVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 7158

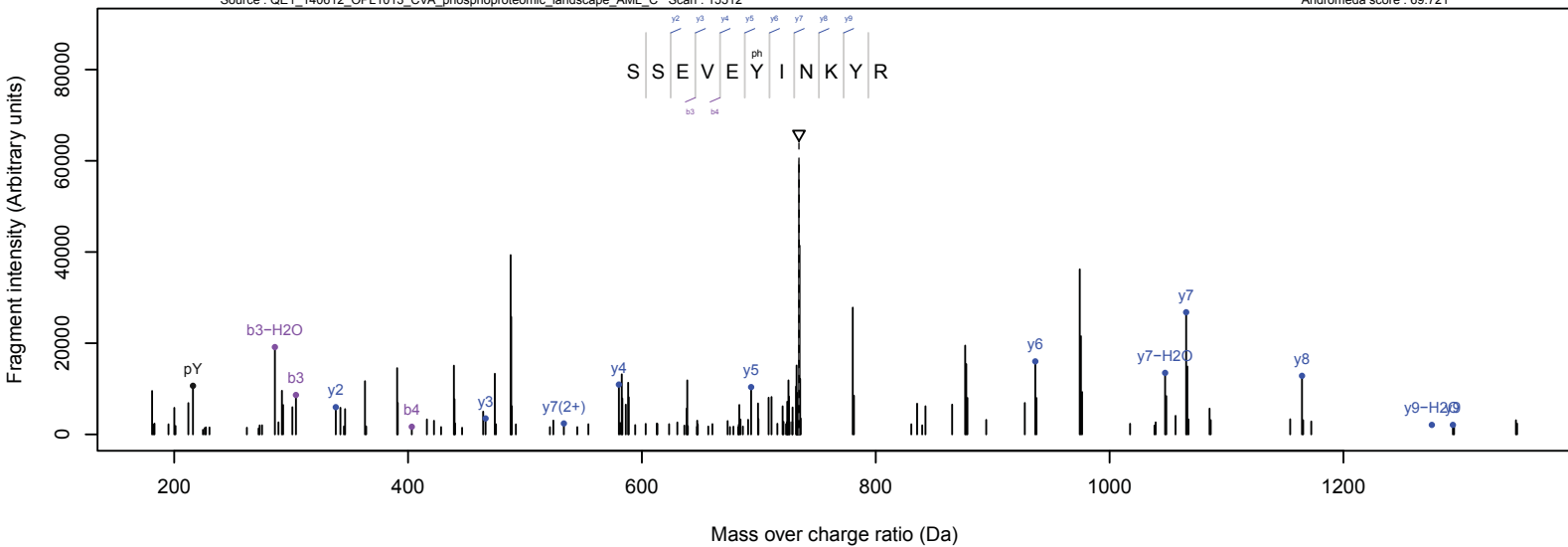
Andromeda score: 69.721



## SSEVEY(ph)INKYR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15512

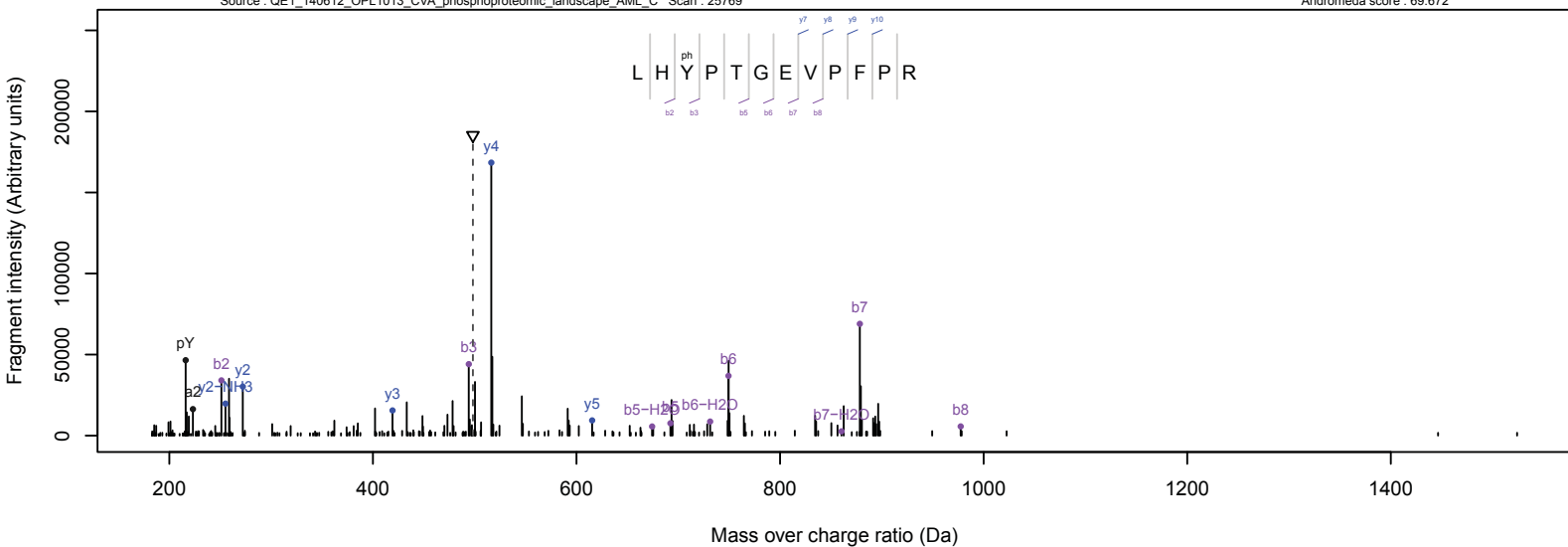
Andromeda score: 69.721



LHY(ph)PTGEVFPFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25769

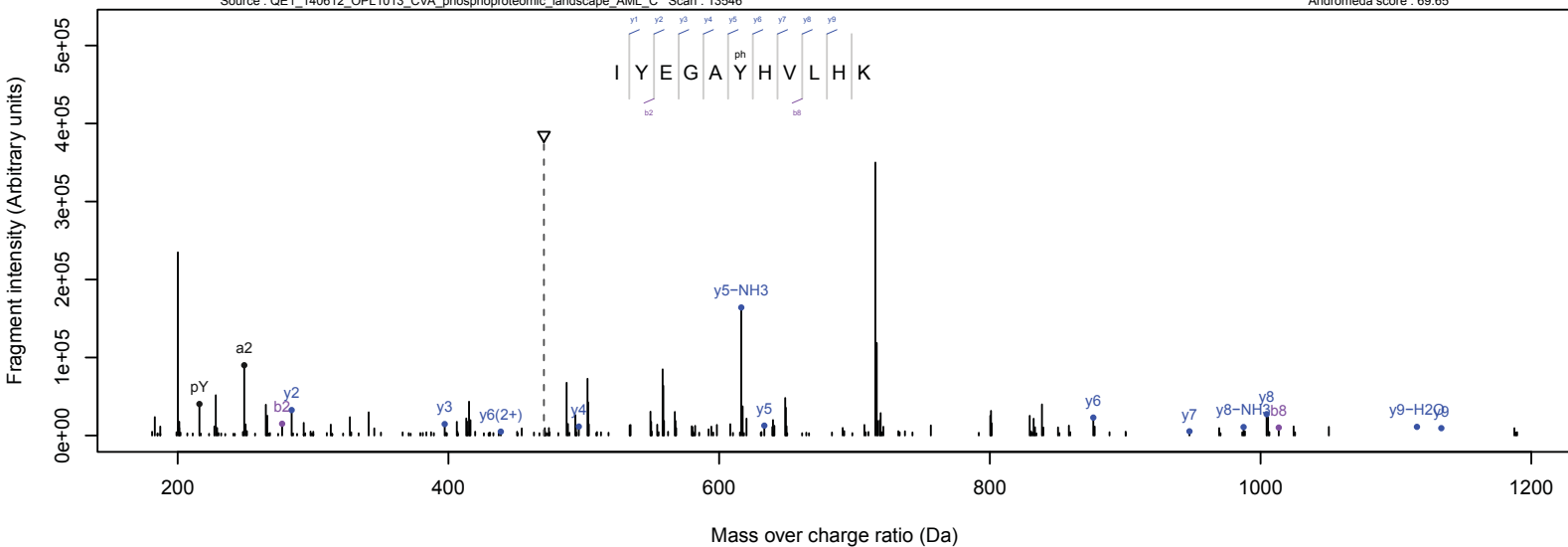
Andromeda score : 69.672



IYEGAY(ph)HVLHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13546

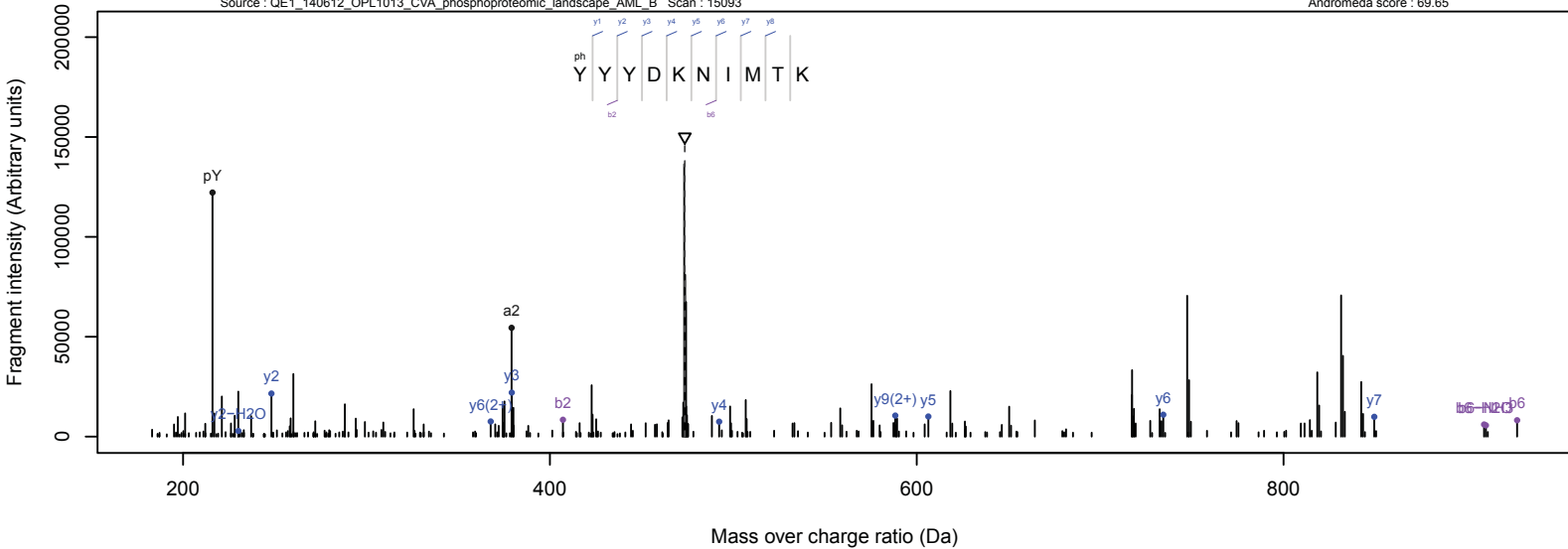
Andromeda score : 69.65



Y(ph)YYDKNIMTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15093

Andromeda score : 69.65

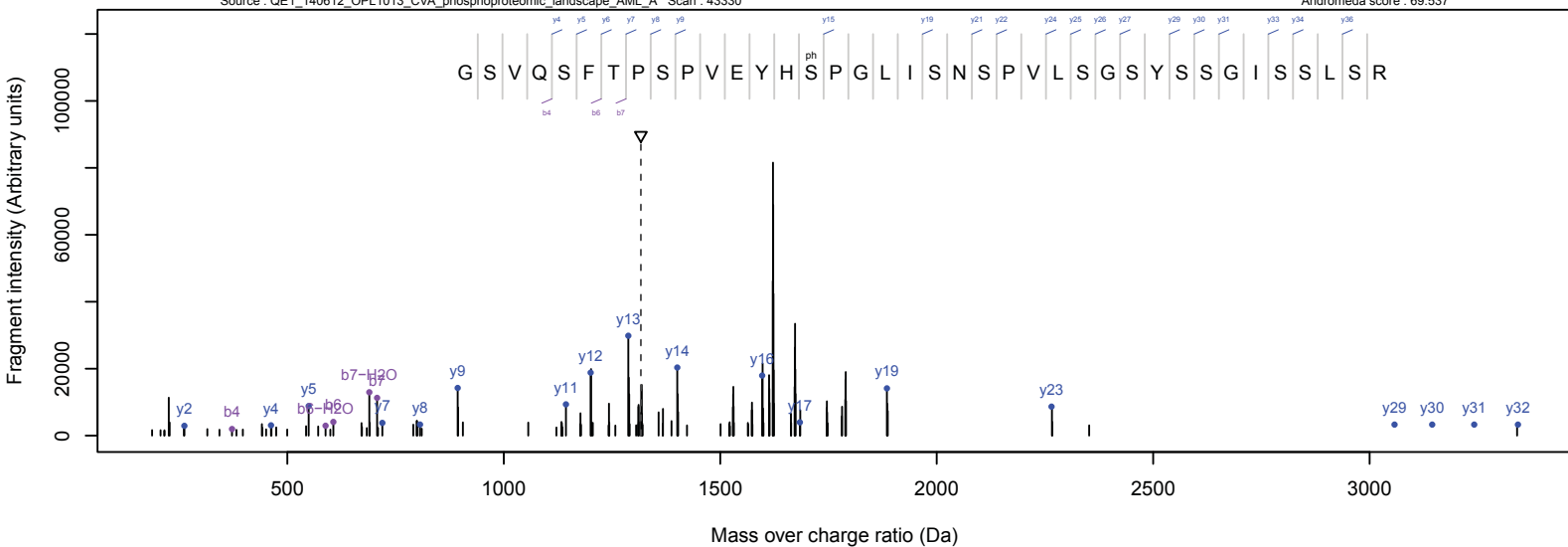




GSVQSFPTSPVEYHSPGLISNSPVLGSGYSYSSGISLSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 43330

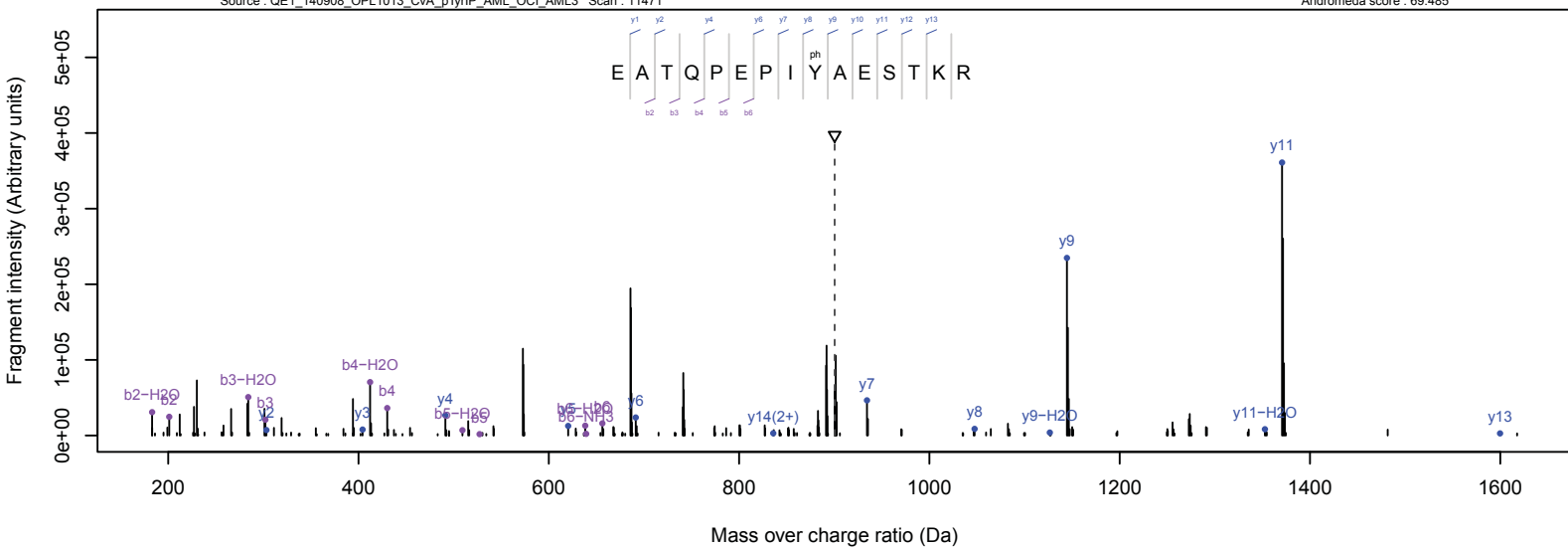
Andromeda score : 69.537



EATQPEPIY(ph)AESTKR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 11471

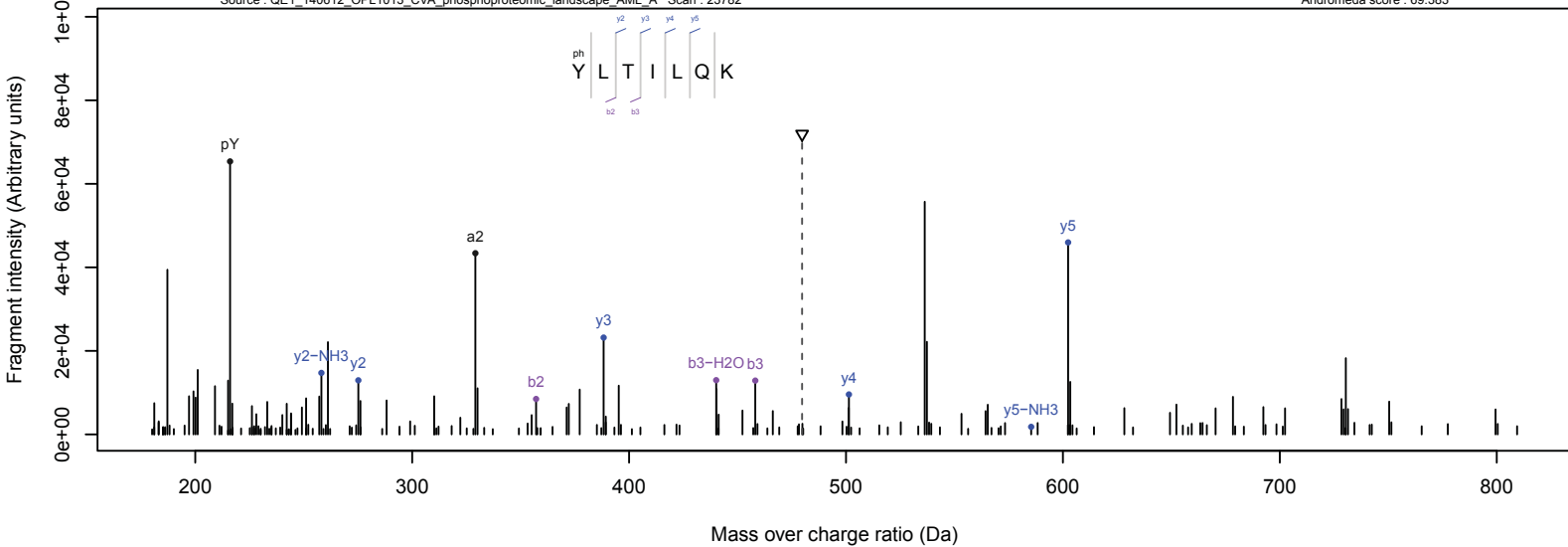
Andromeda score : 69.485



Y(ph)LTILQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 23782

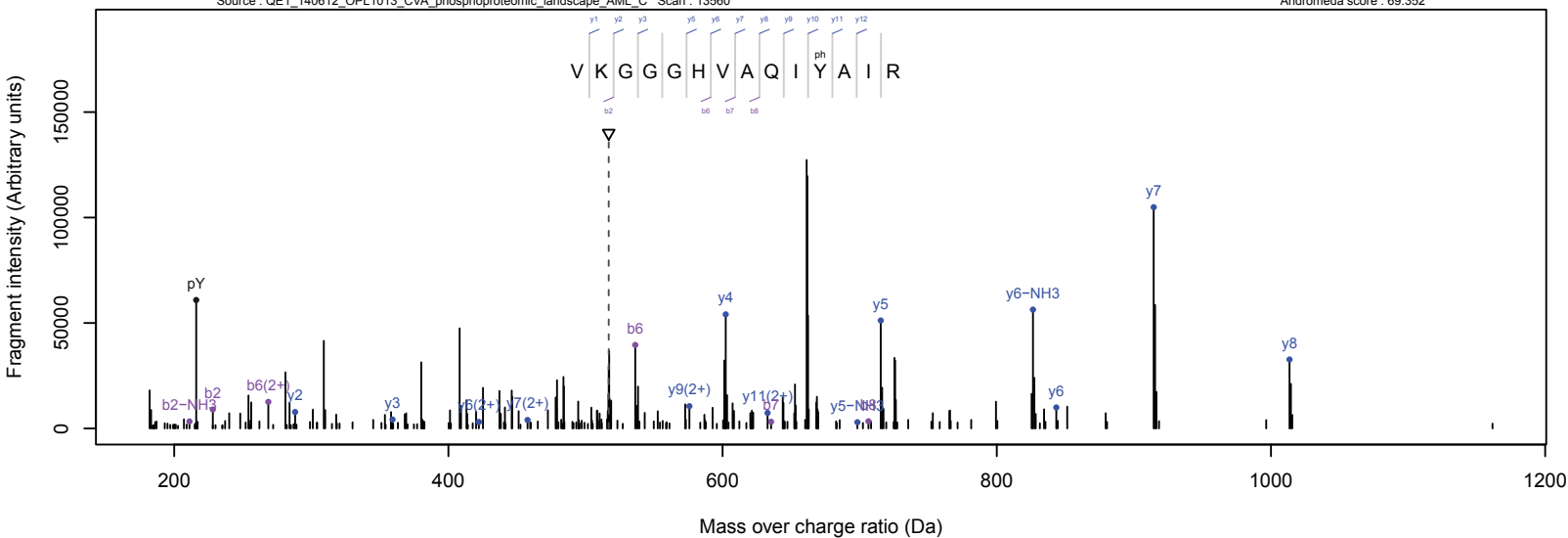
Andromeda score : 69.383



## VKGGHVAQIYAIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13560

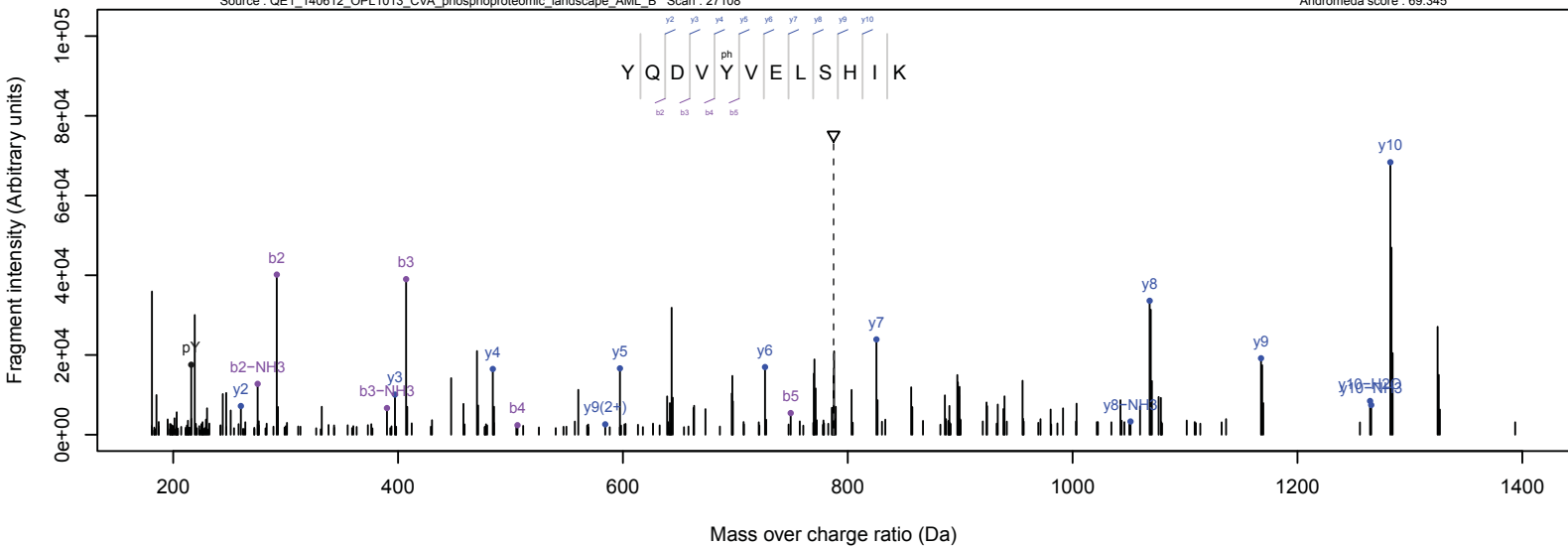
Andromeda score: 69.352



## YQDVYVELSHIK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27108

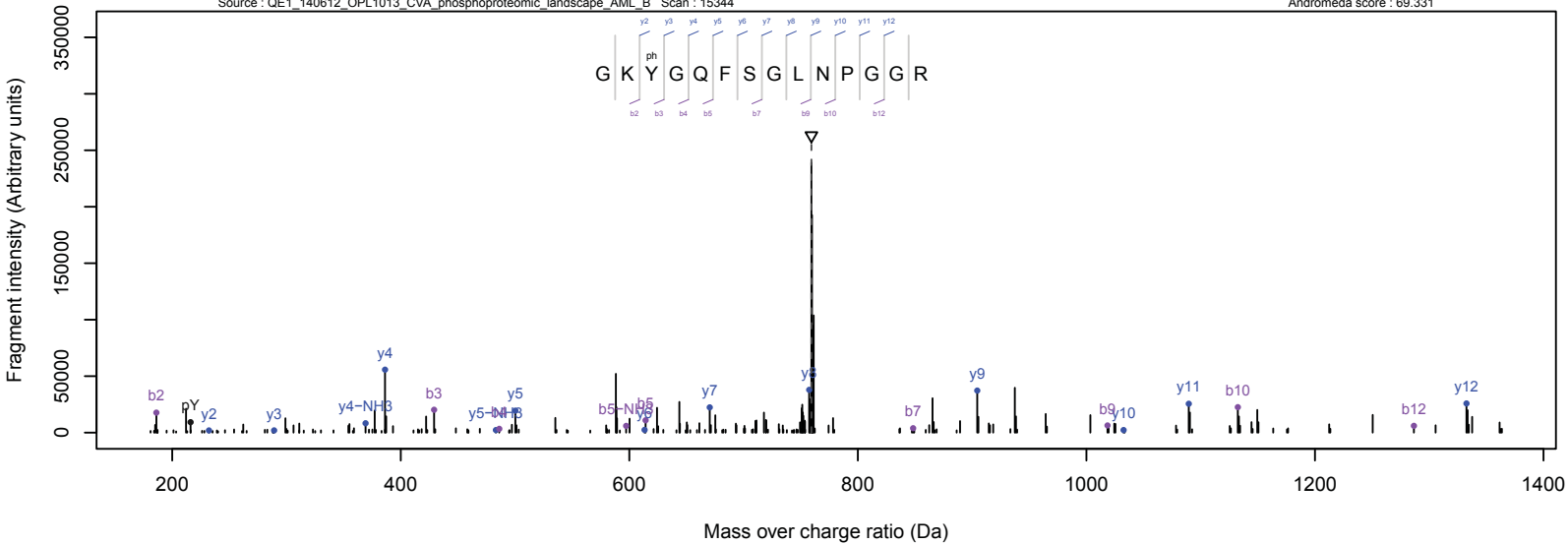
Andromeda score: 69.345



## GKY(ph)GQFSLNPGGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15344

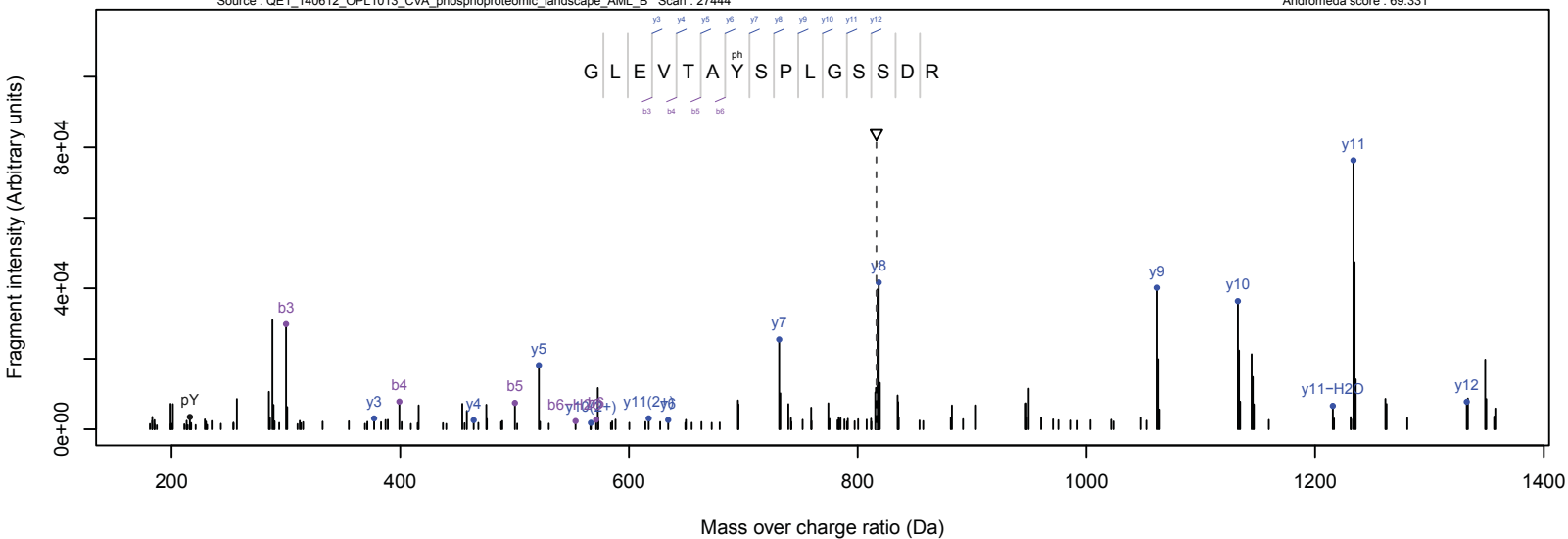
Andromeda score: 69.331



## GLEVTAY(ph)SPLGSSDR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27444

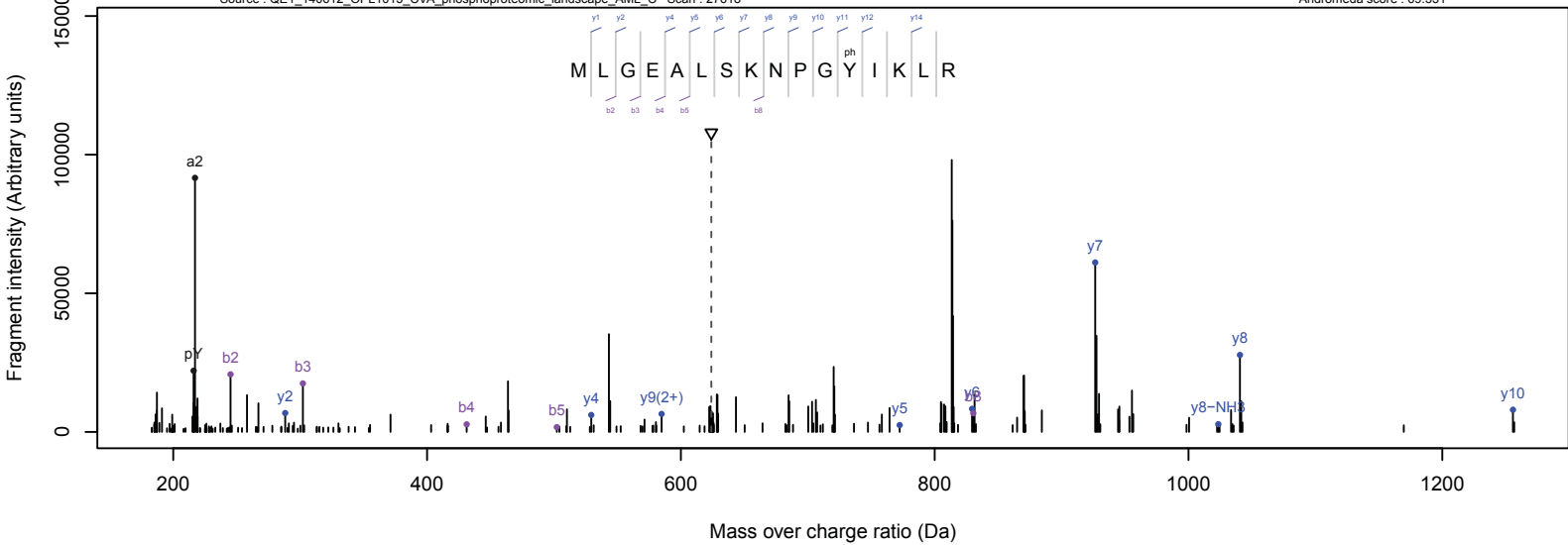
Andromeda score: 69.331



## MLGEALSKNPGY(ph)IKLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 27618

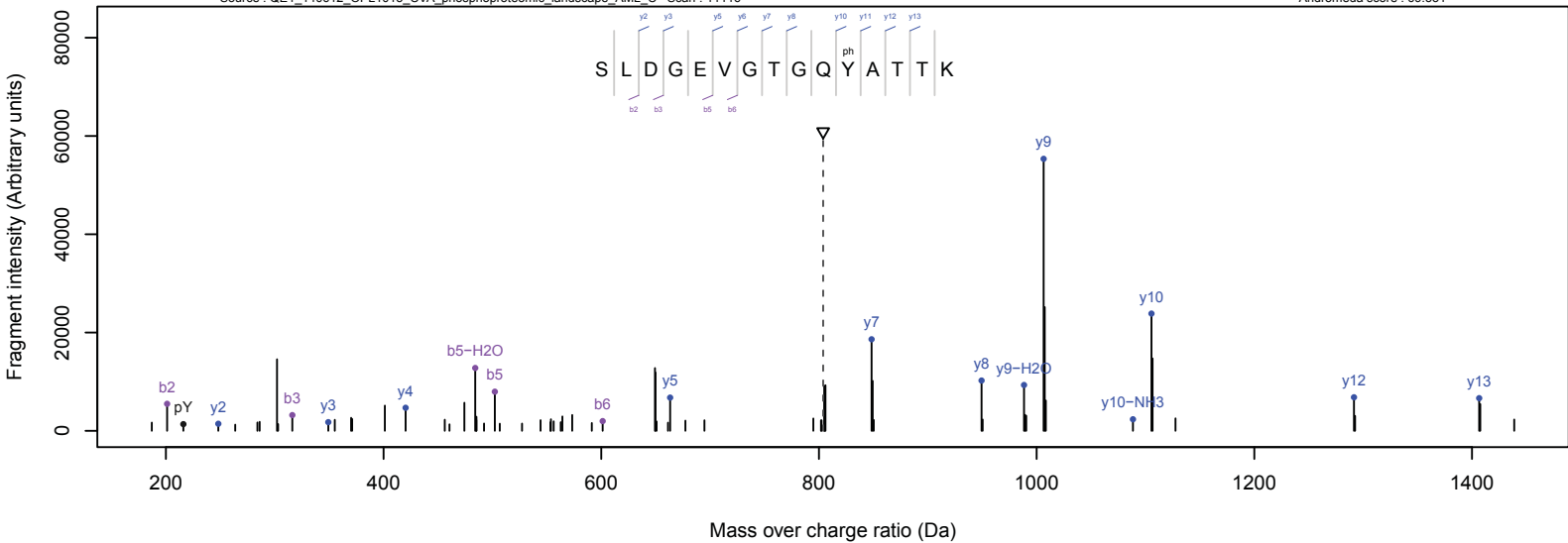
Andromeda score: 69.331



## SLDGEVGTGQY(ph)ATTK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11116

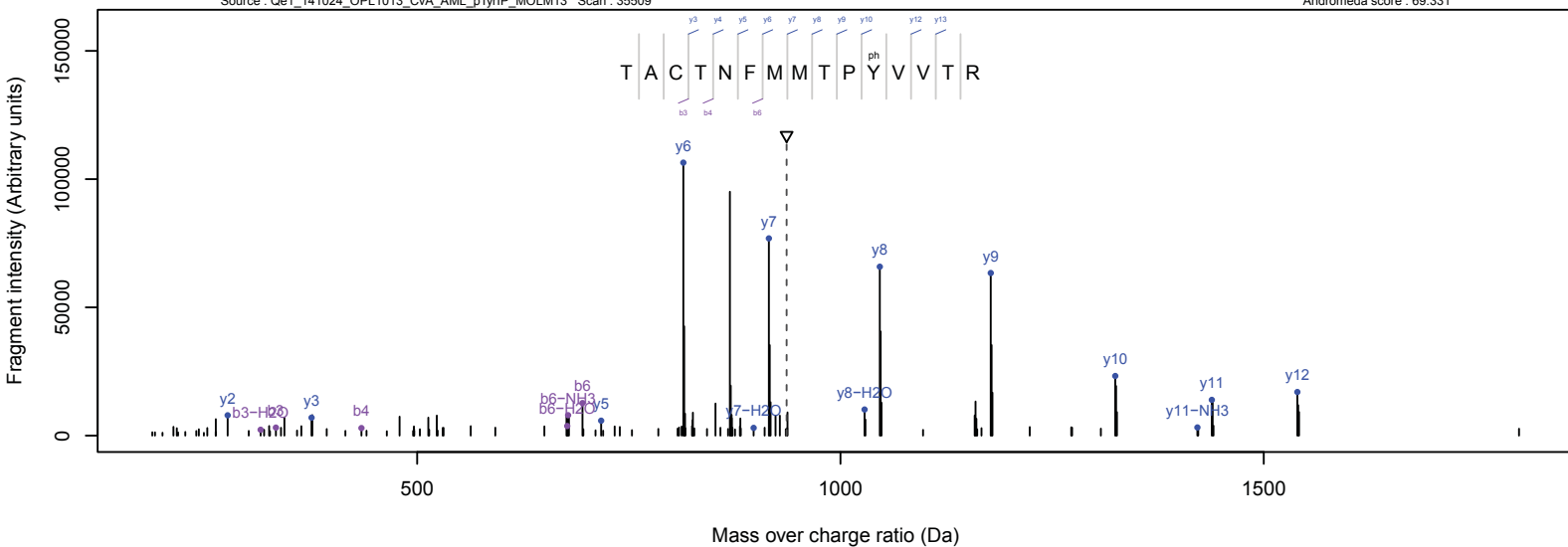
Andromeda score: 69.331



## TACTNFMMPY(ph)VVTR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 35509

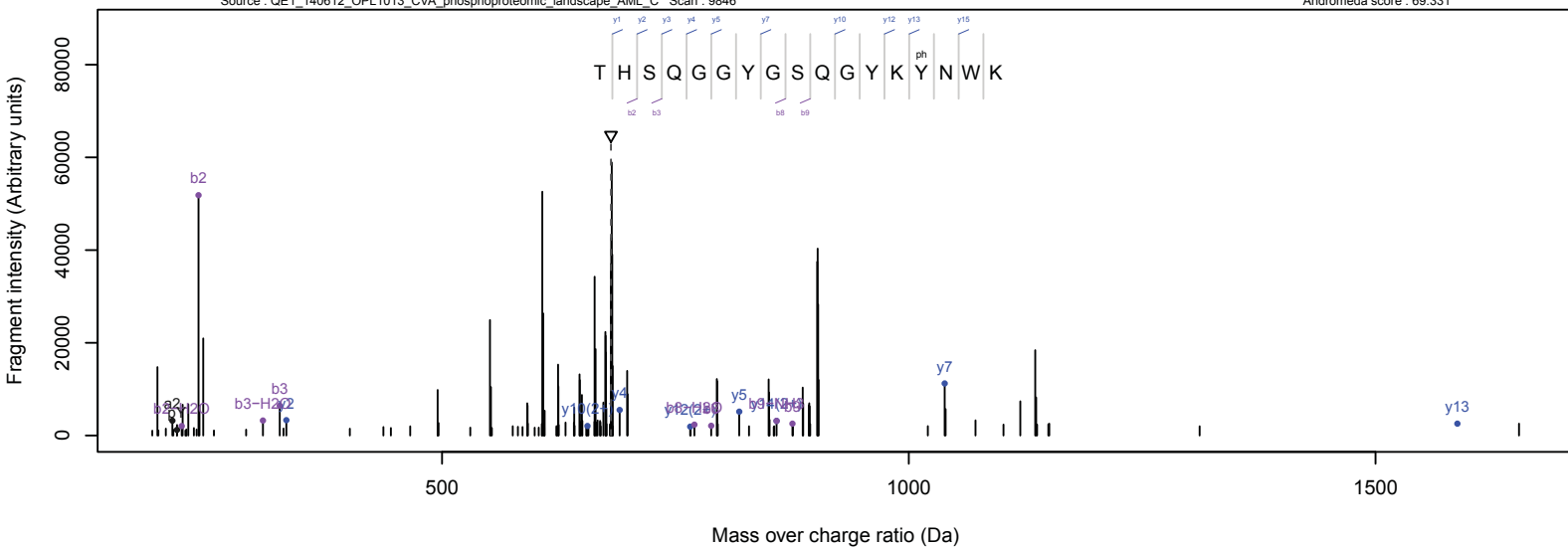
Andromeda score : 69.331



## THSQGGYGSQGYK(ph)NWK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9846

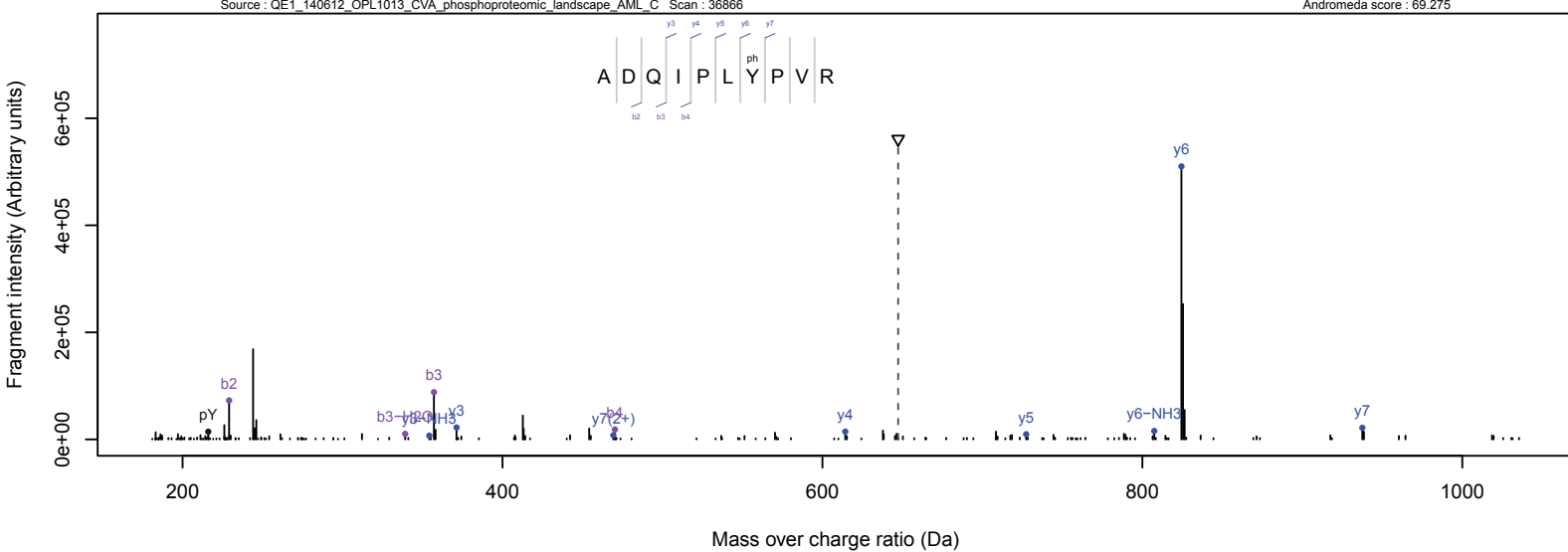
Andromeda score : 69.331



## (ac)ADQIPLY(ph)PVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36866

Andromeda score : 69.275

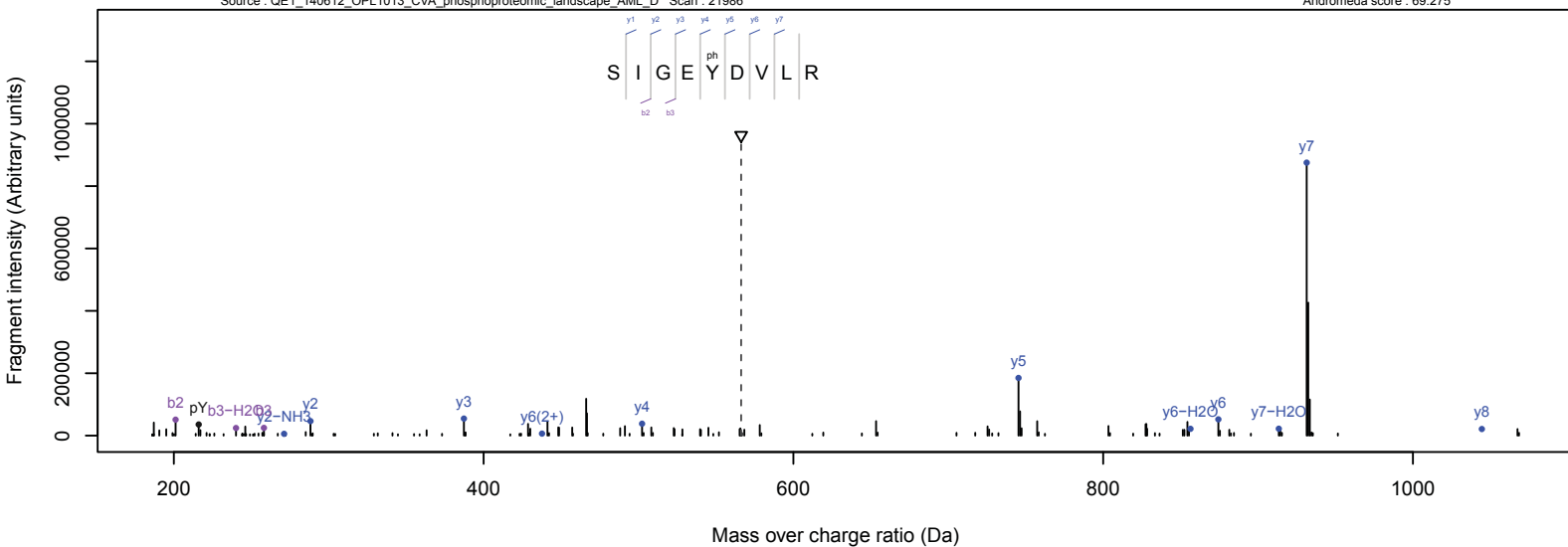




## SIGEY(ph)DVLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 21986

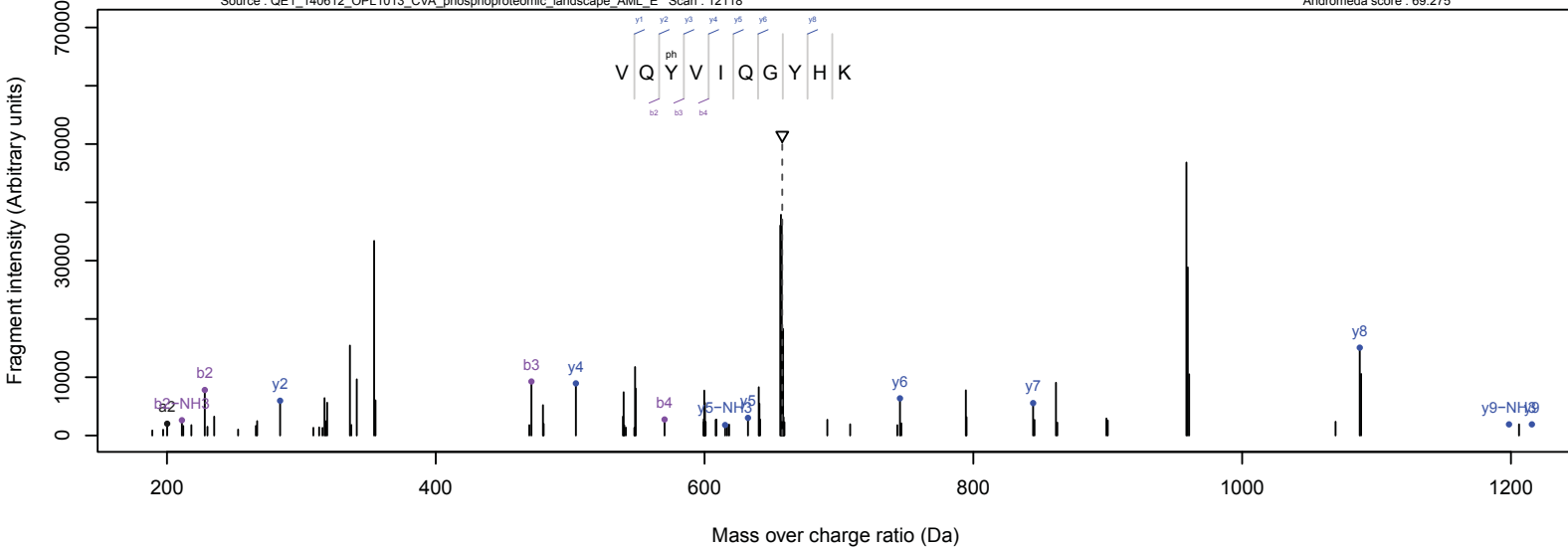
Andromeda score: 69.275



## VQY(ph)VIQGYHK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 12118

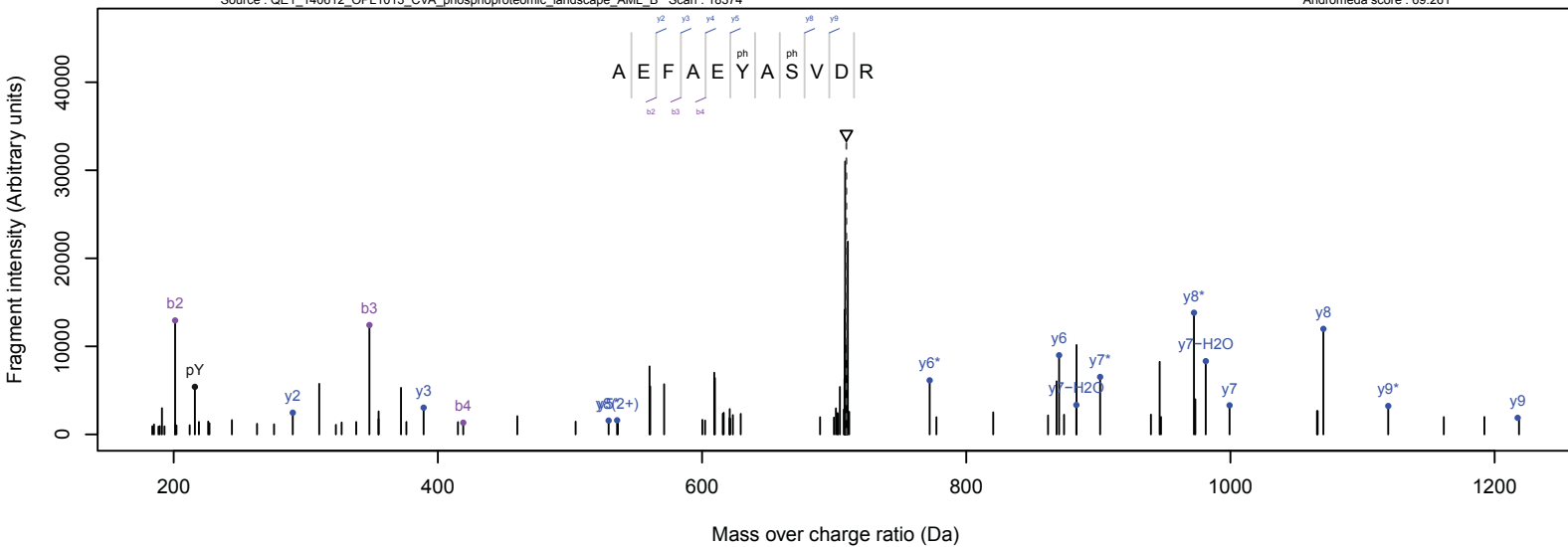
Andromeda score: 69.275



## AEFAEY(ph)AS(ph)VDR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18374

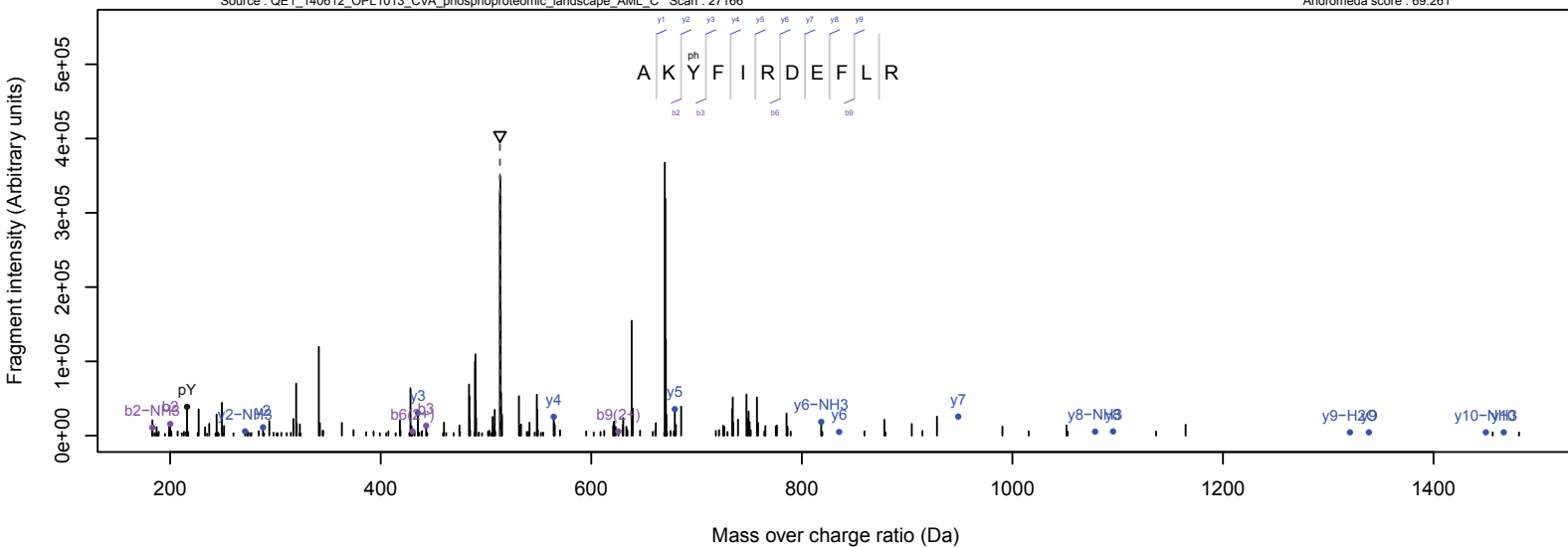
Andromeda score: 69.261



## AKY(ph)FIRDEFLLR

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27166

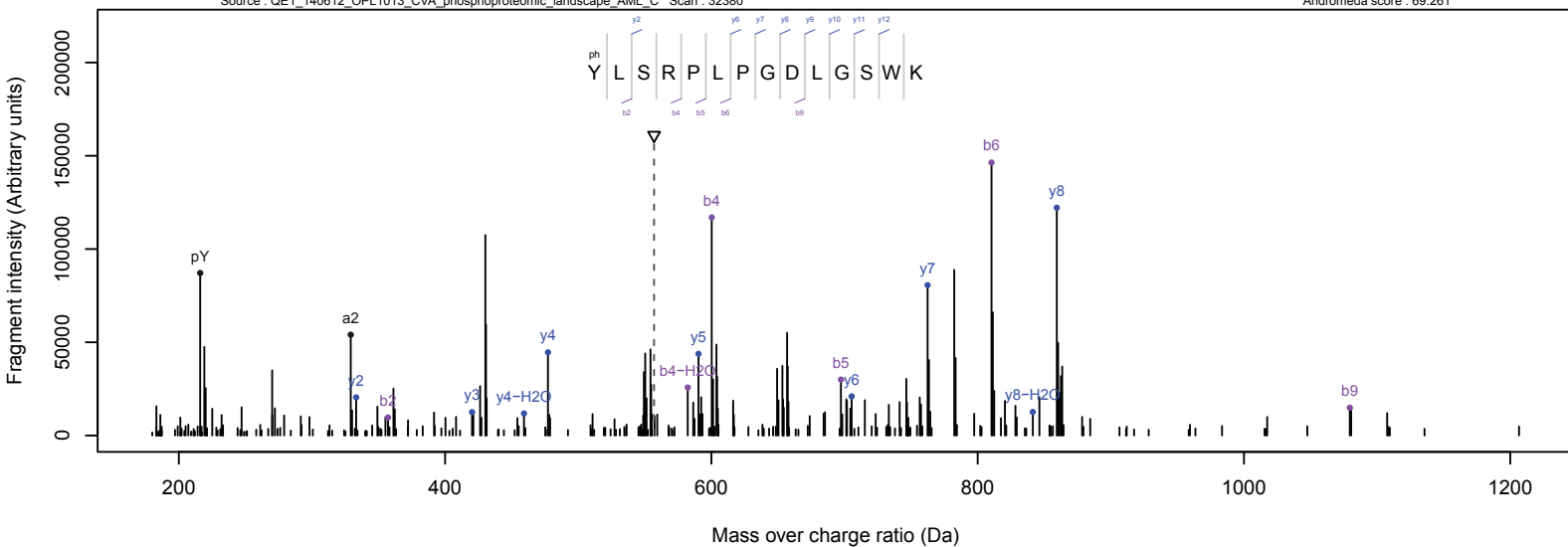
Andromeda score : 69.261



## Y(ph)LSRPLPGDLGSWK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32380

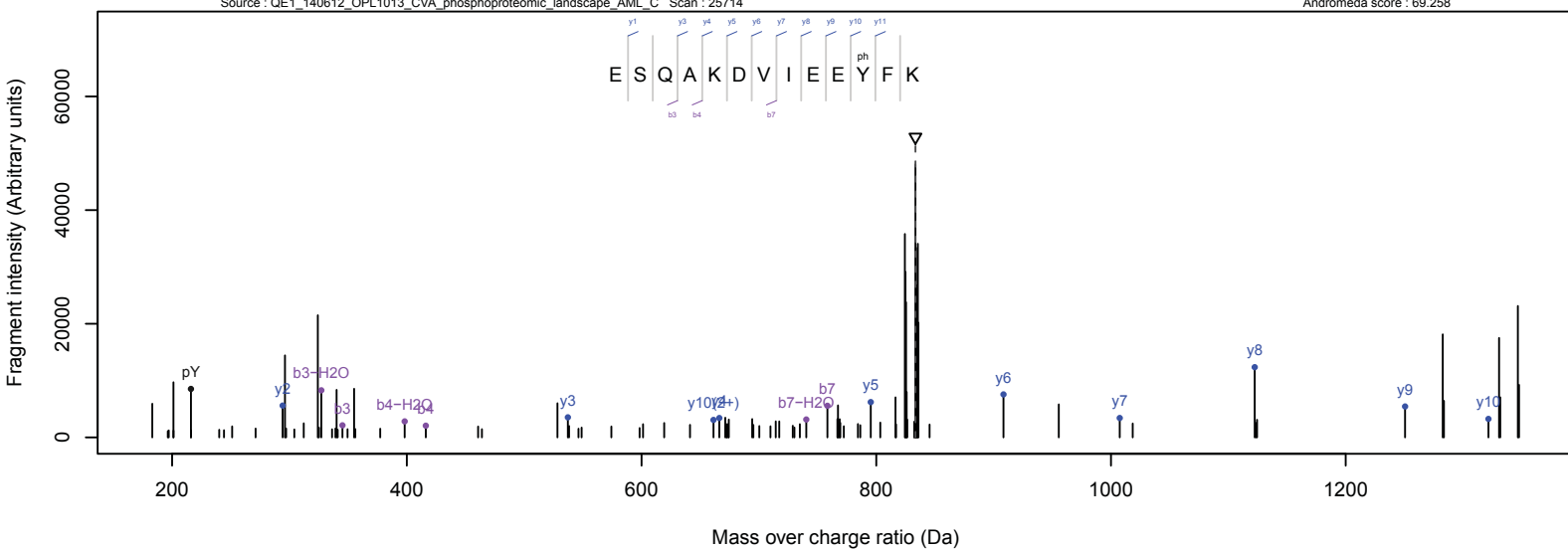
Andromeda score : 69.261



## ESQAKDVEEY(ph)FK

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25714

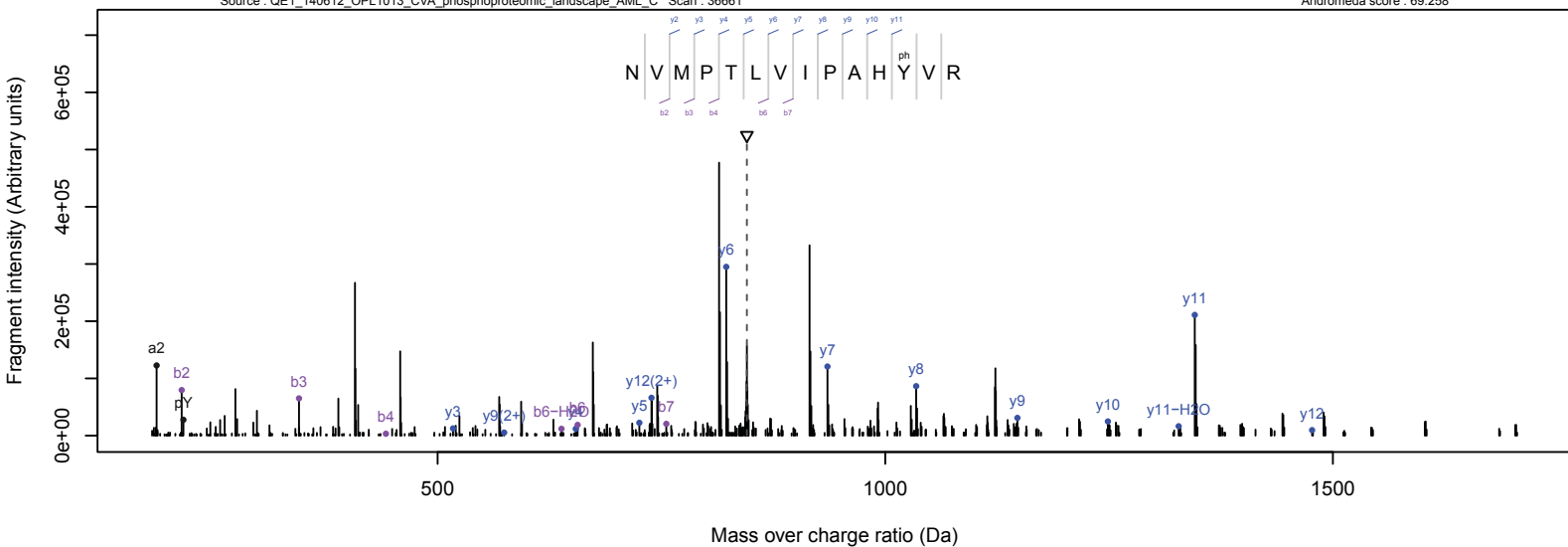
Andromeda score : 69.258



NVMPTLVIPAHYVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36661

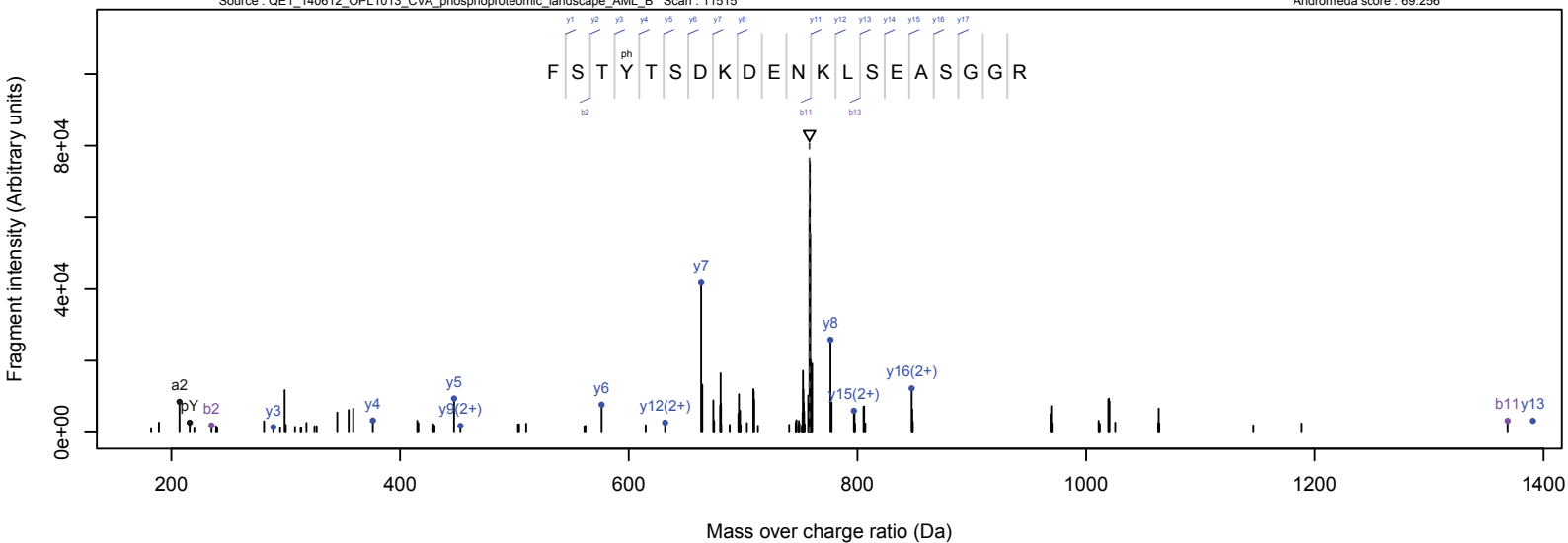
Andromeda score: 69.258



FSTY(ph)TSDKDENKLSAEGGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 11515

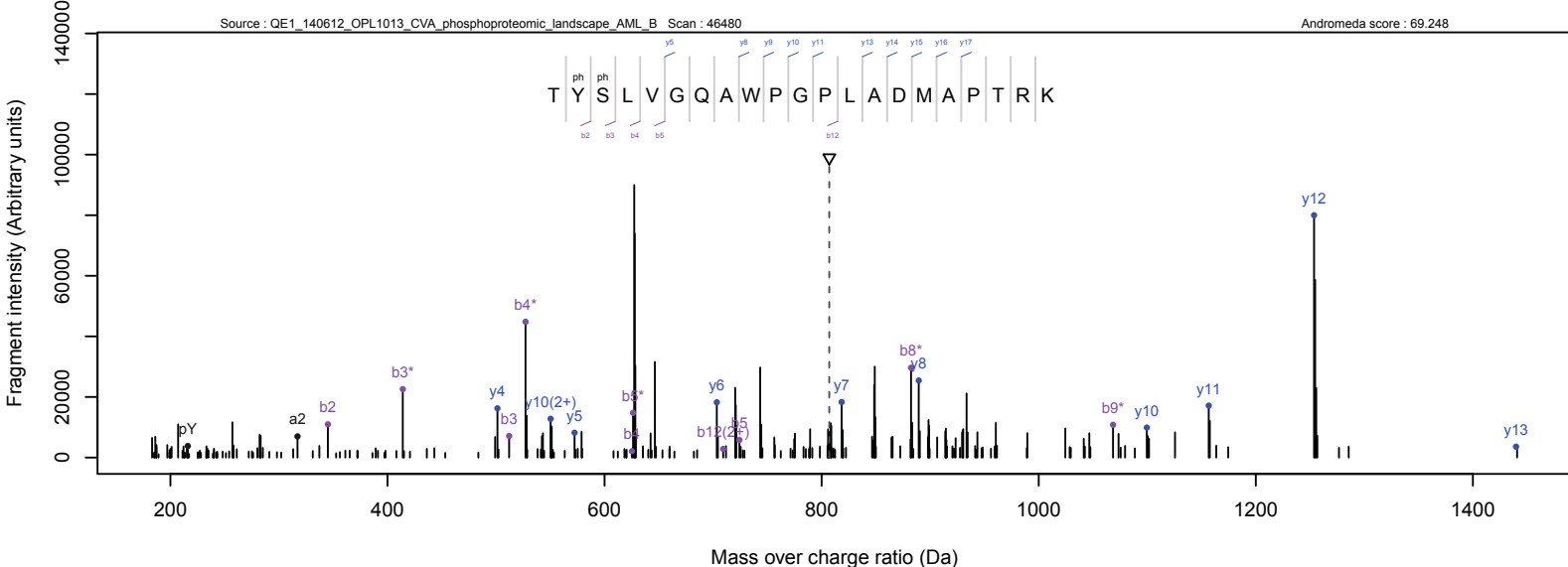
Andromeda score: 69.256



TY(ph)S(ph)LVGQAWPGPLADMAPTRK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 46480

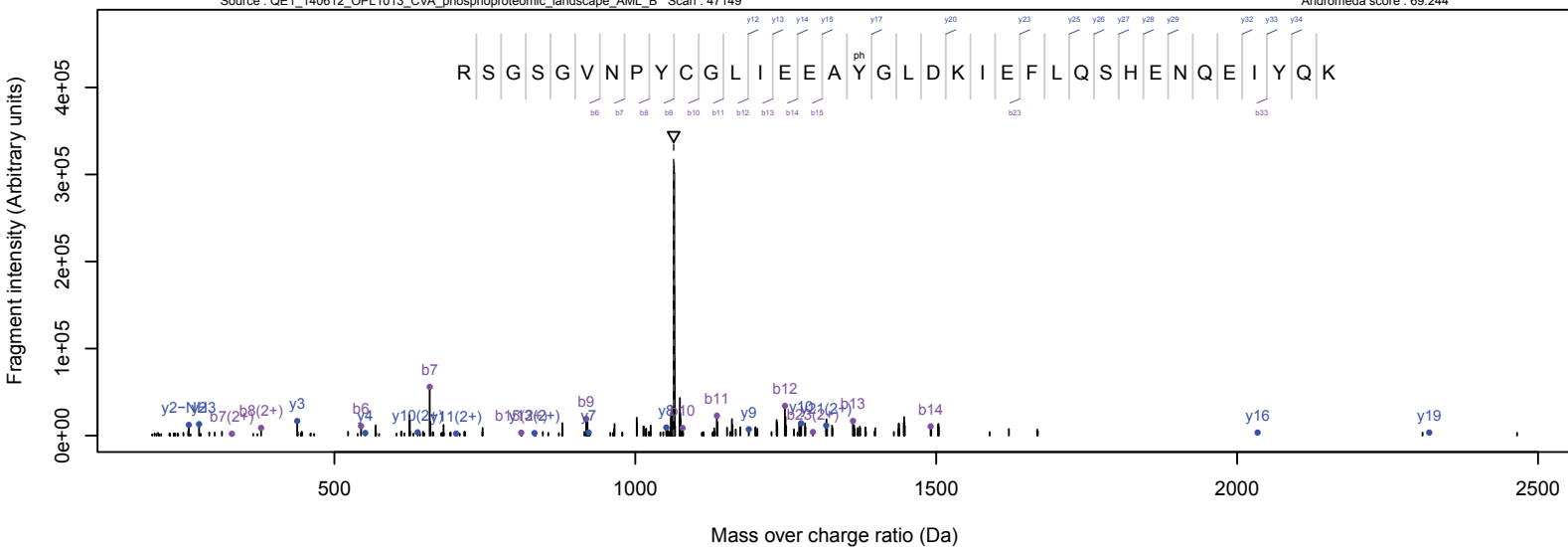
Andromeda score: 69.248



## RSGSGVNPYCGLIIEEAY(ph)GLDKIEFLQSHENQEIYQK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47149

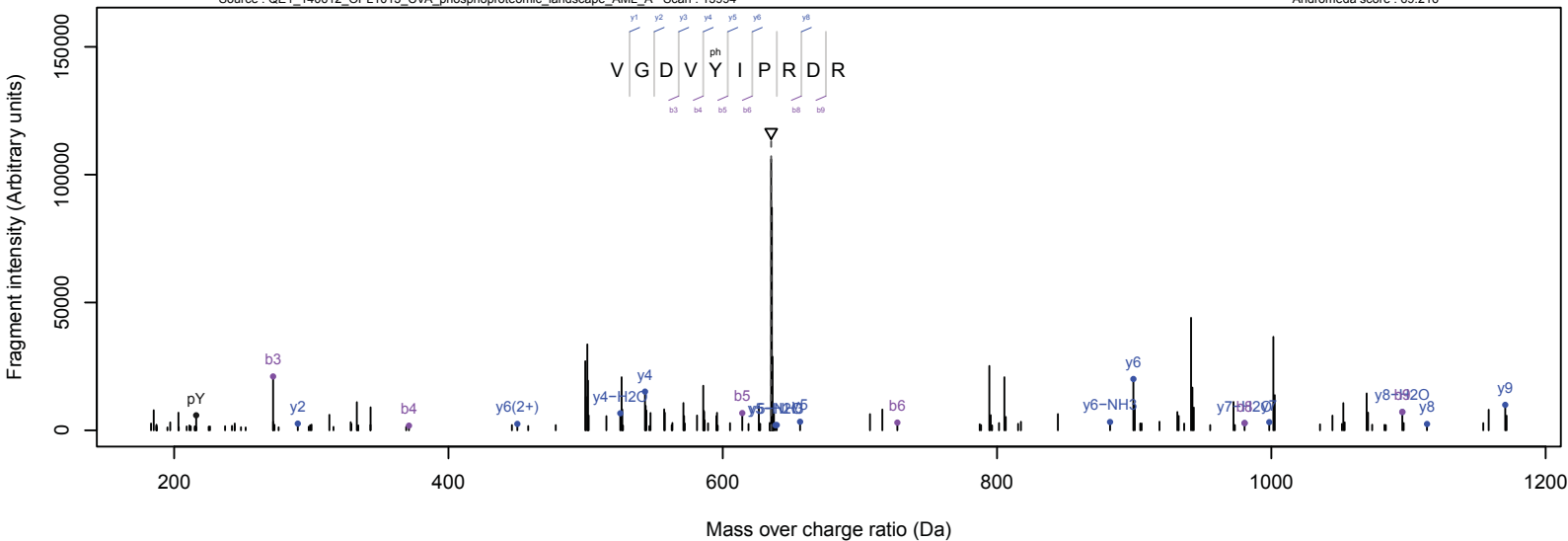
Andromeda score : 69.244



## VGDVY(ph)IPRDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 13994

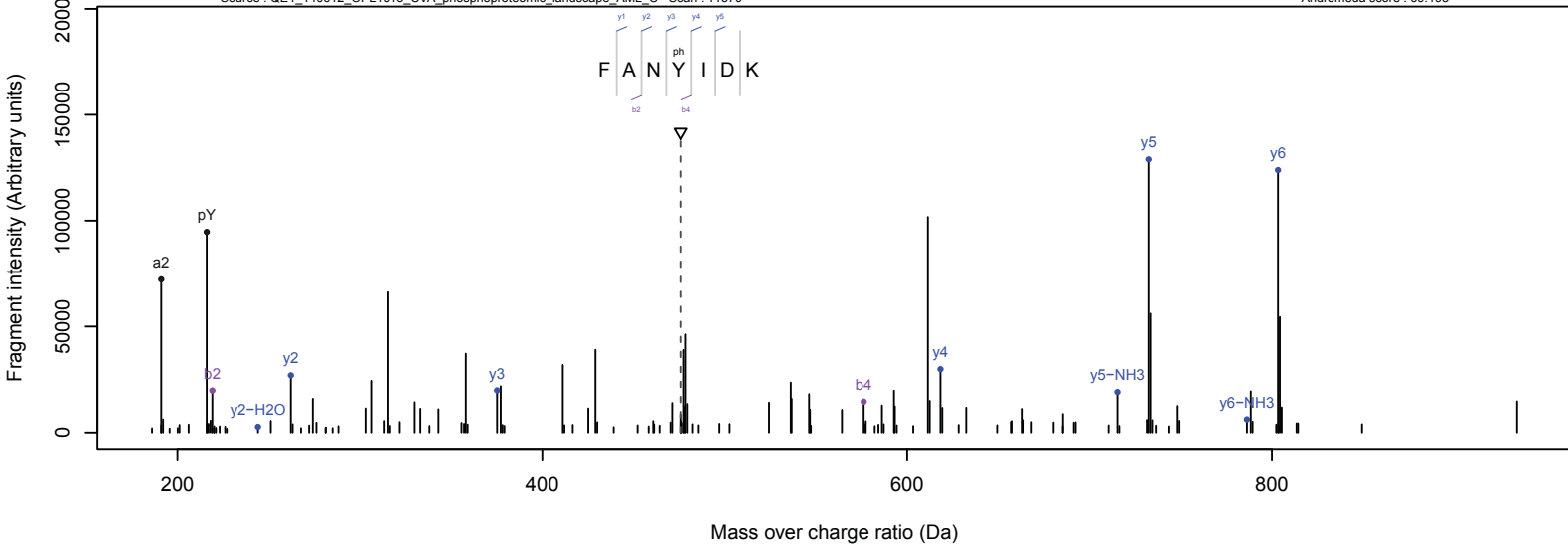
Andromeda score : 69.216



## FANYIDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11370

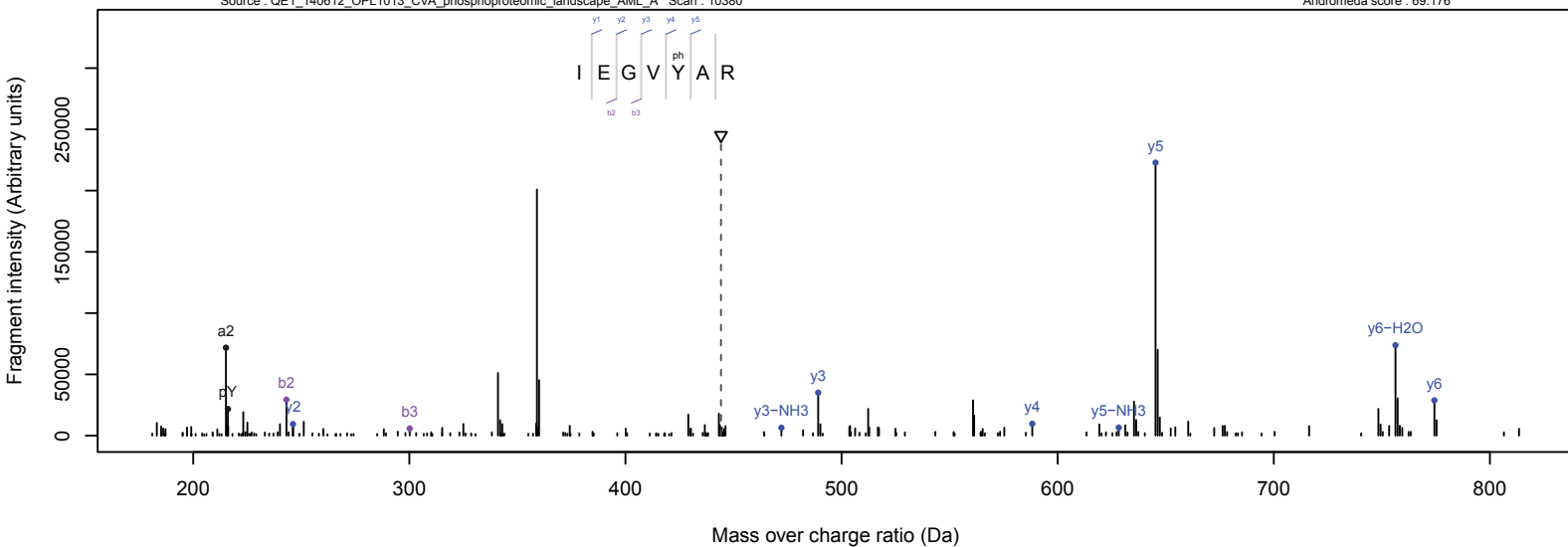
Andromeda score : 69.198



## IEGVY(ph)AR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 10380

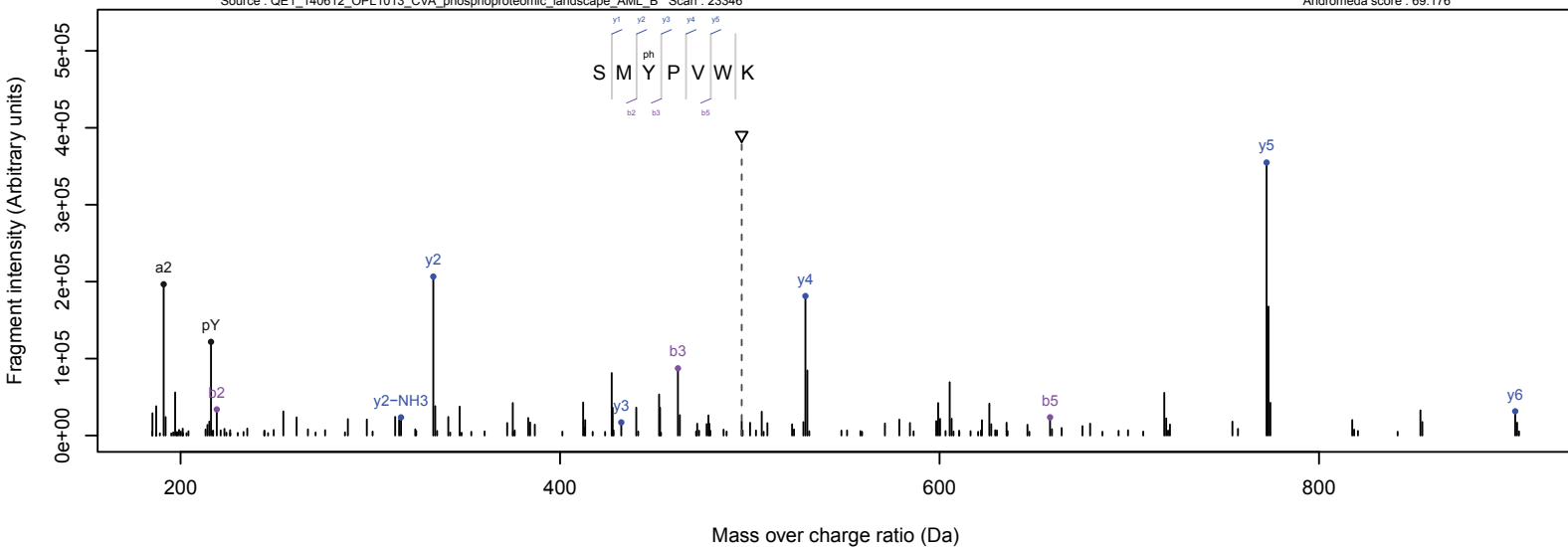
Andromeda score: 69.176



## SMY(ph)PVWK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 23346

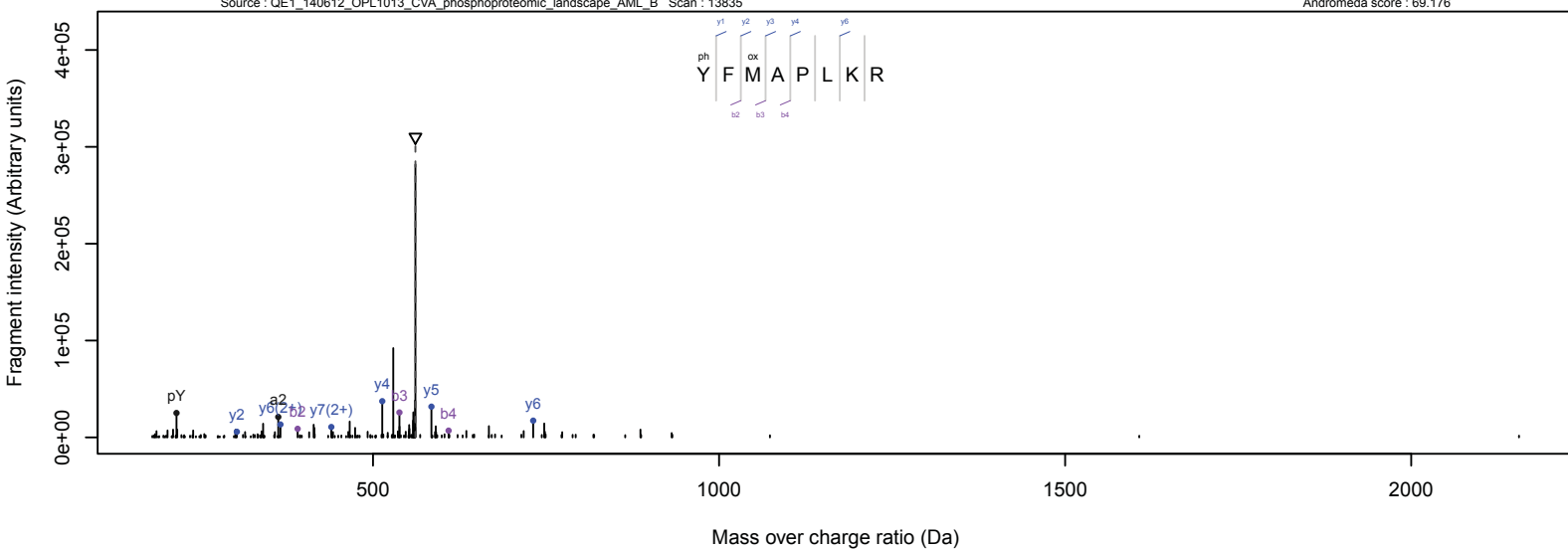
Andromeda score: 69.176



## Y(ph)FM(ox)APLKR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 13835

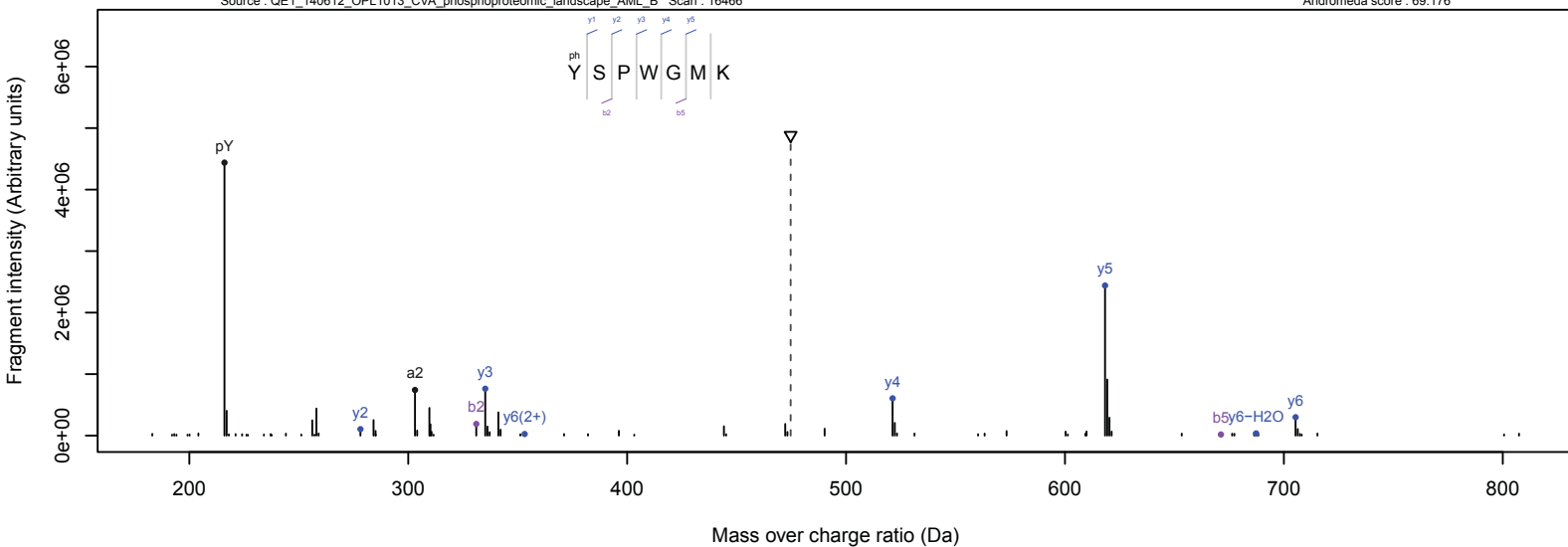
Andromeda score: 69.176



## Y(ph)SPWGMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16466

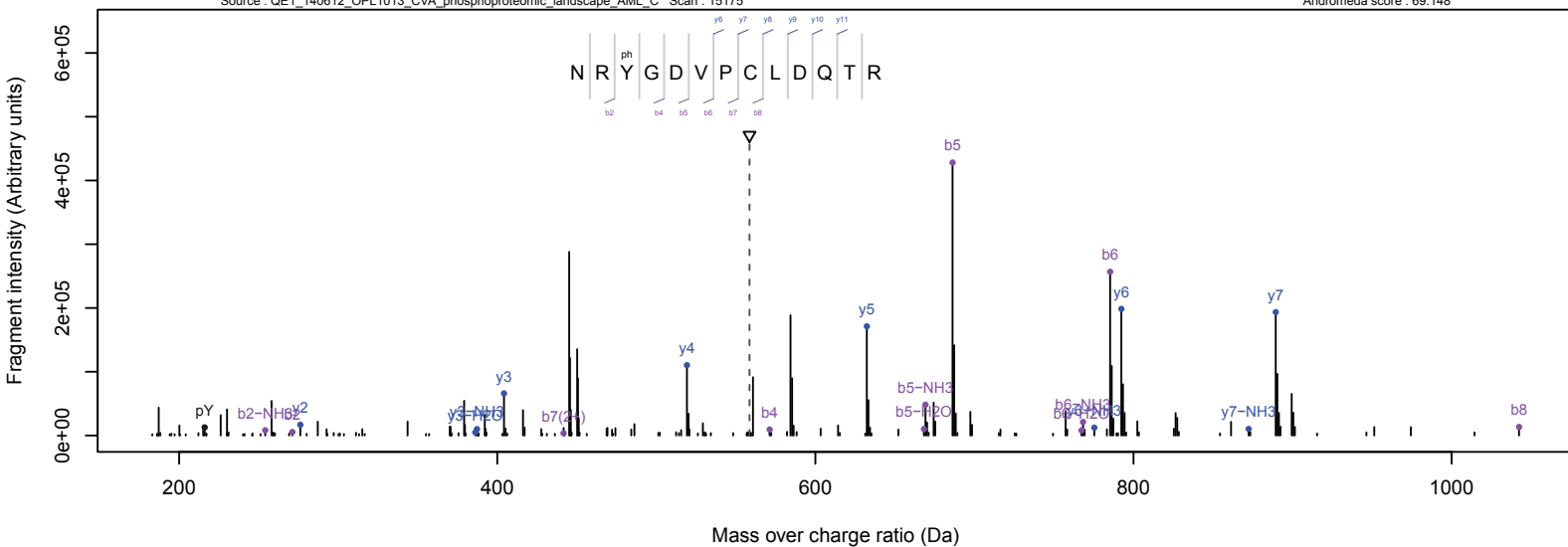
Andromeda score : 69.176



## NRY(ph)GDVPCLDQTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15175

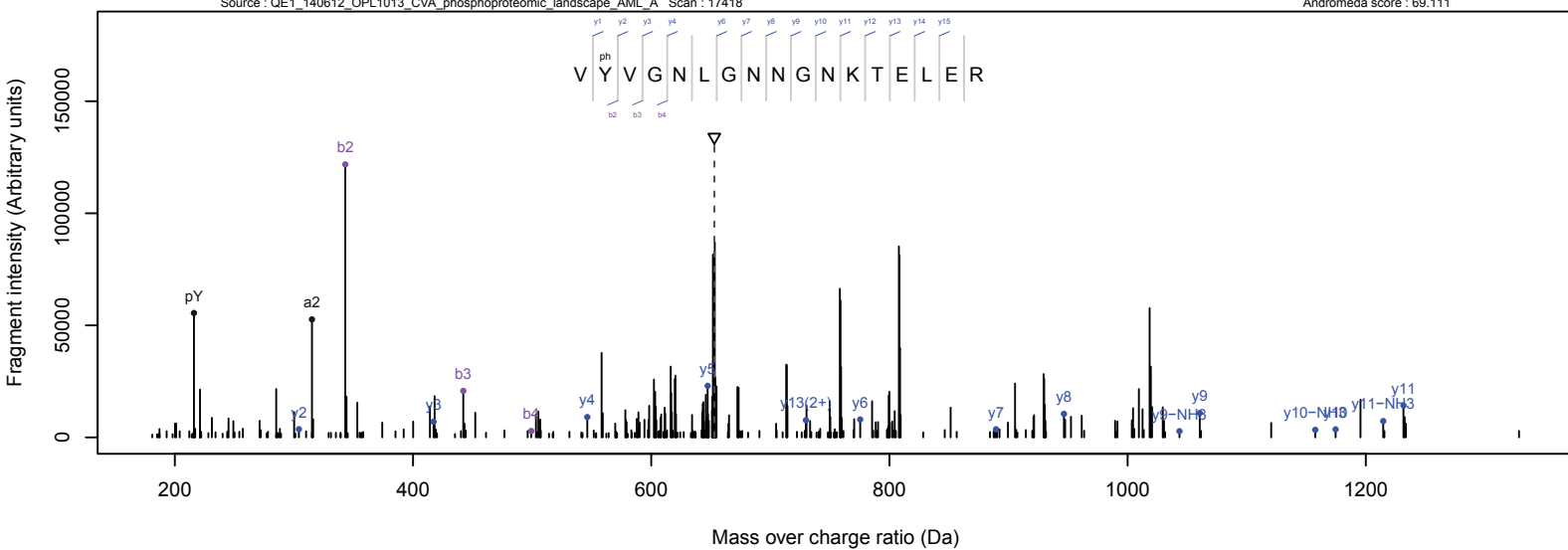
Andromeda score : 69.148



## VY(ph)VGNLGNNGKTELER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17418

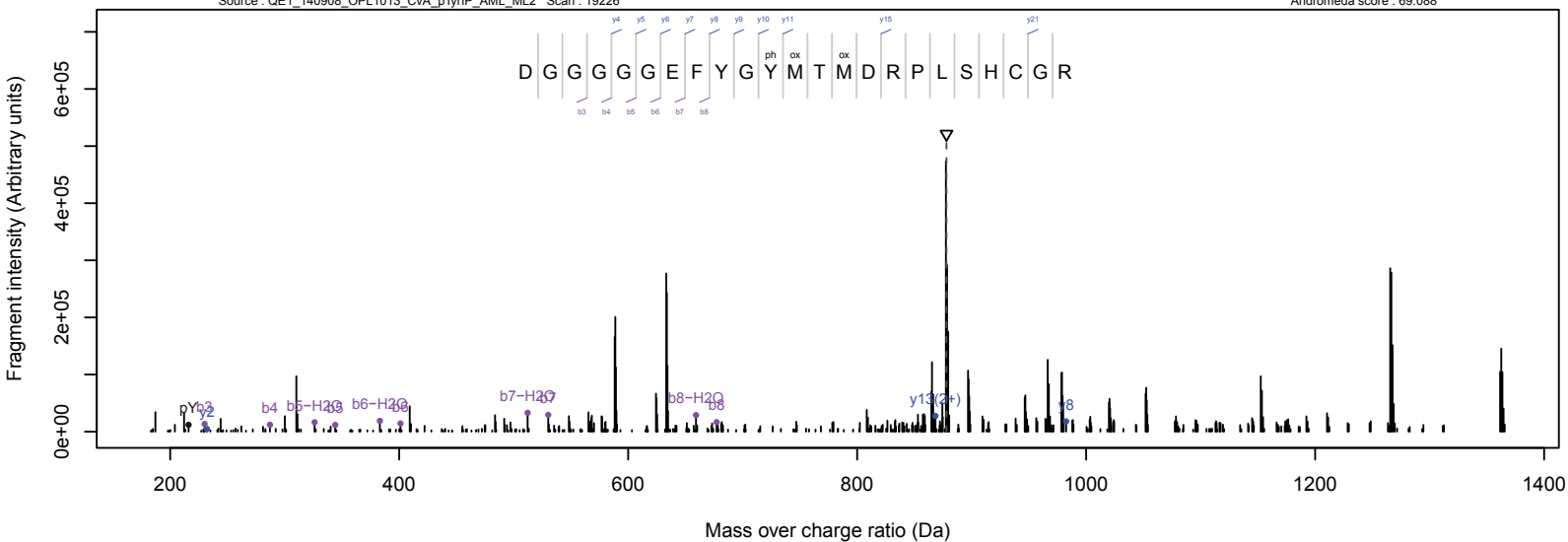
Andromeda score : 69.111



DGGGGGEFYGYM(ox)TM(ox)DRPLSHCGR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 19226

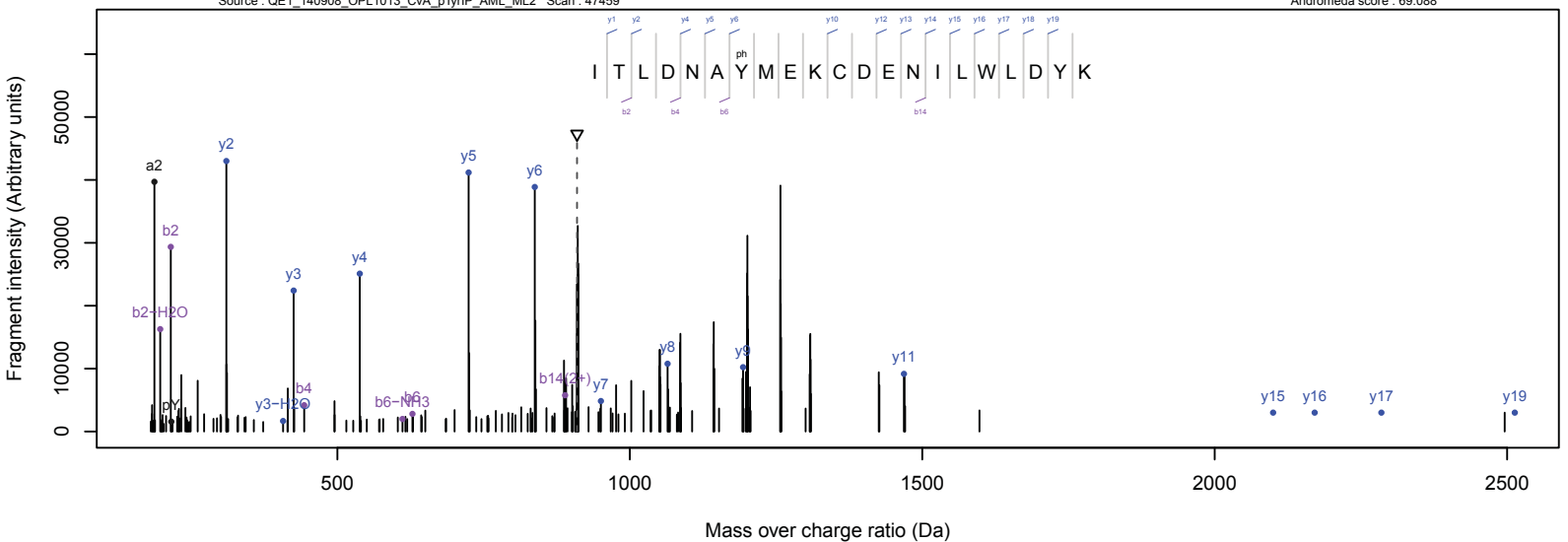
Andromeda score : 69.088



ITLDNAY(ph)MEKCDENILWLDYK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 47459

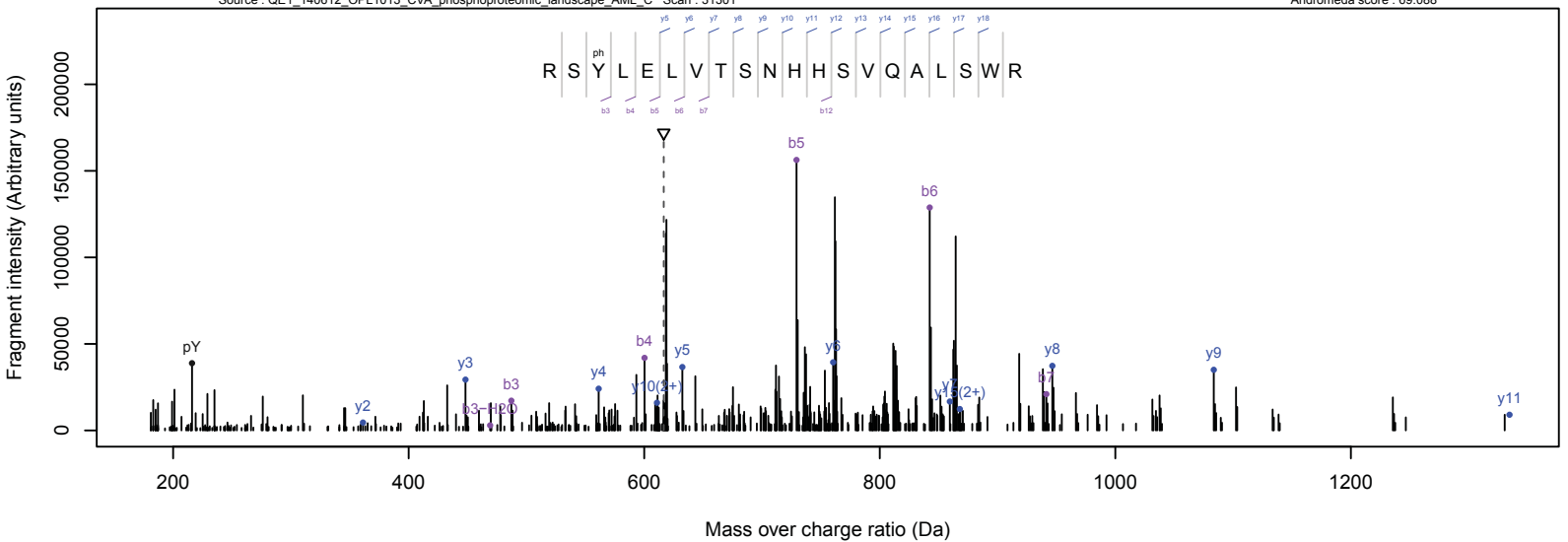
Andromeda score : 69.088



RSY(ph)LELVTSNHHSVQALSWR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31301

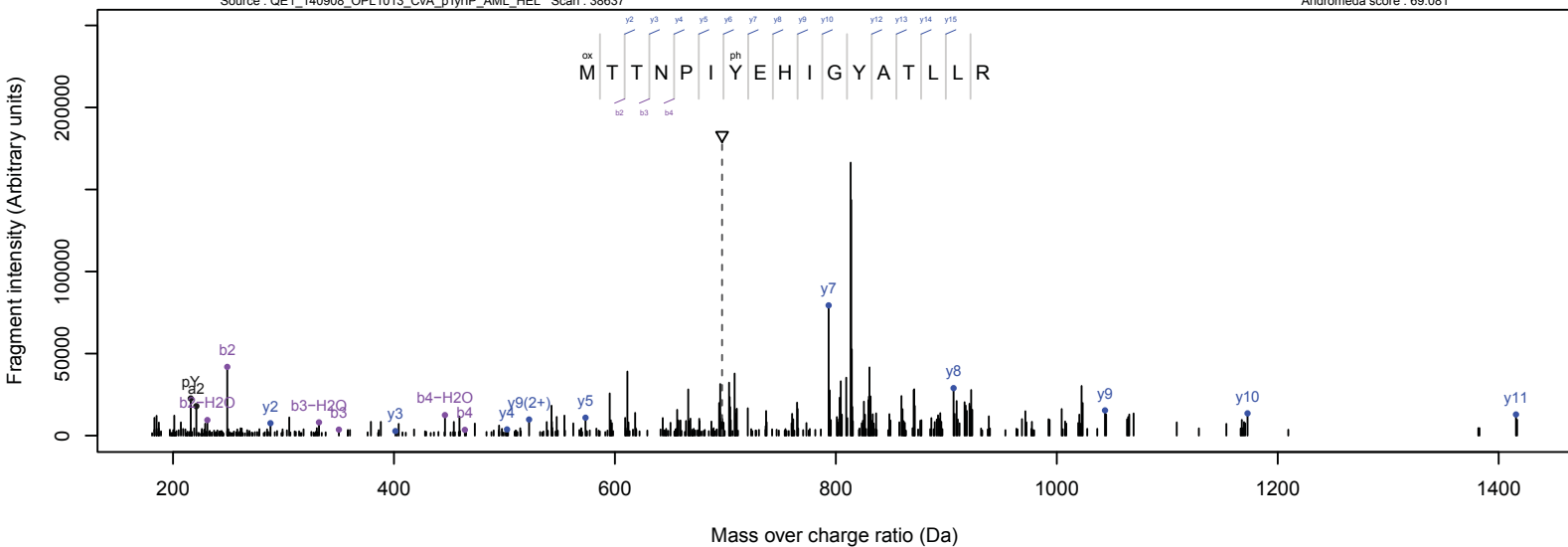
Andromeda score : 69.088



## M(ox)TTNPIY(ph)EHIGYATLLR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 38637

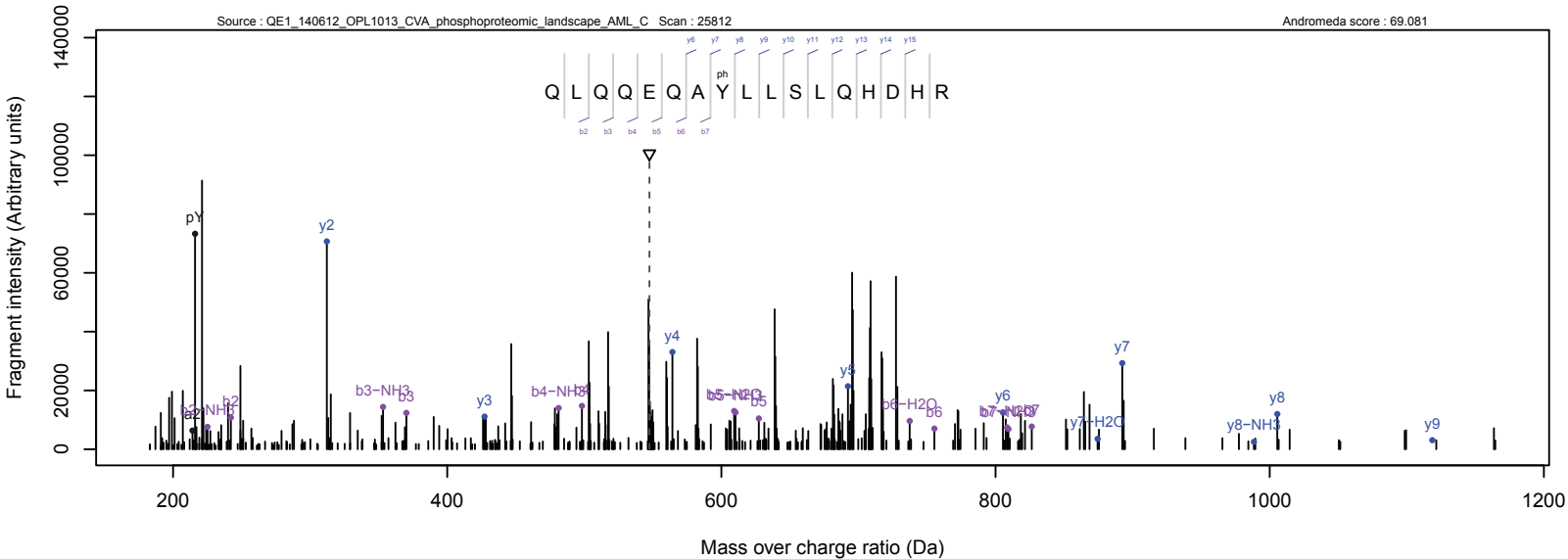
Andromeda score : 69.081



## QLQQEQAY(ph)LLSLQHDHR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25812

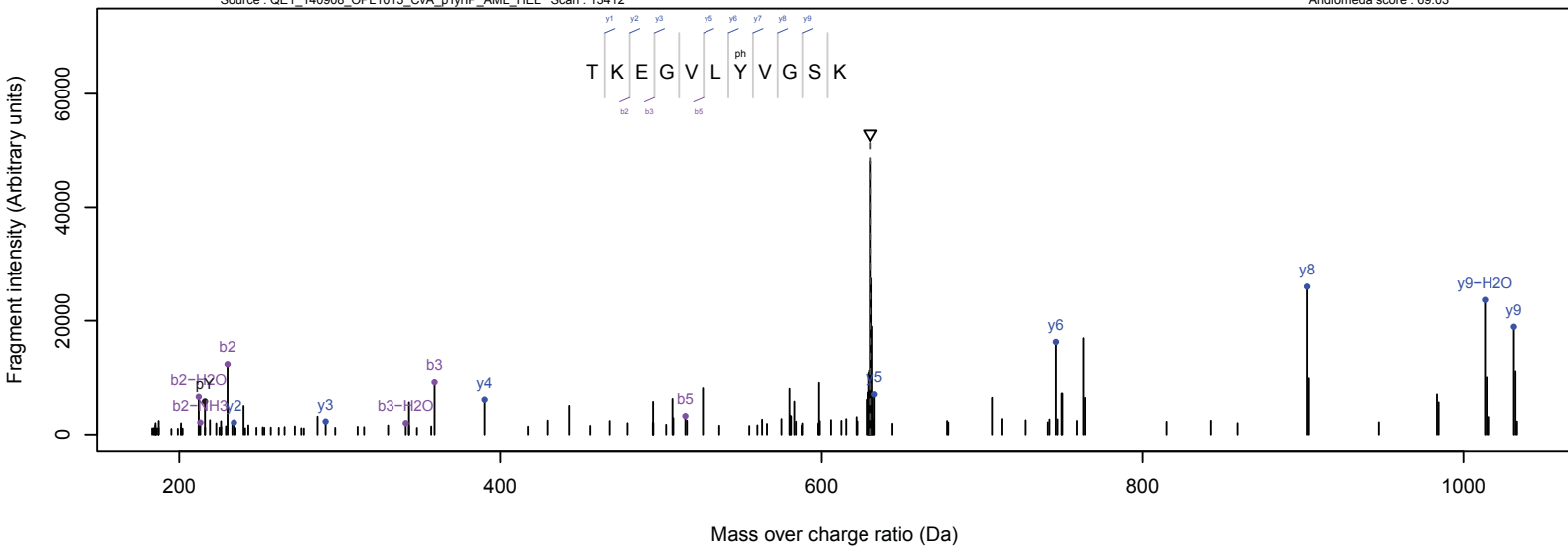
Andromeda score : 69.081



## TKEGVLY(ph)VGSK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 13412

Andromeda score : 69.03

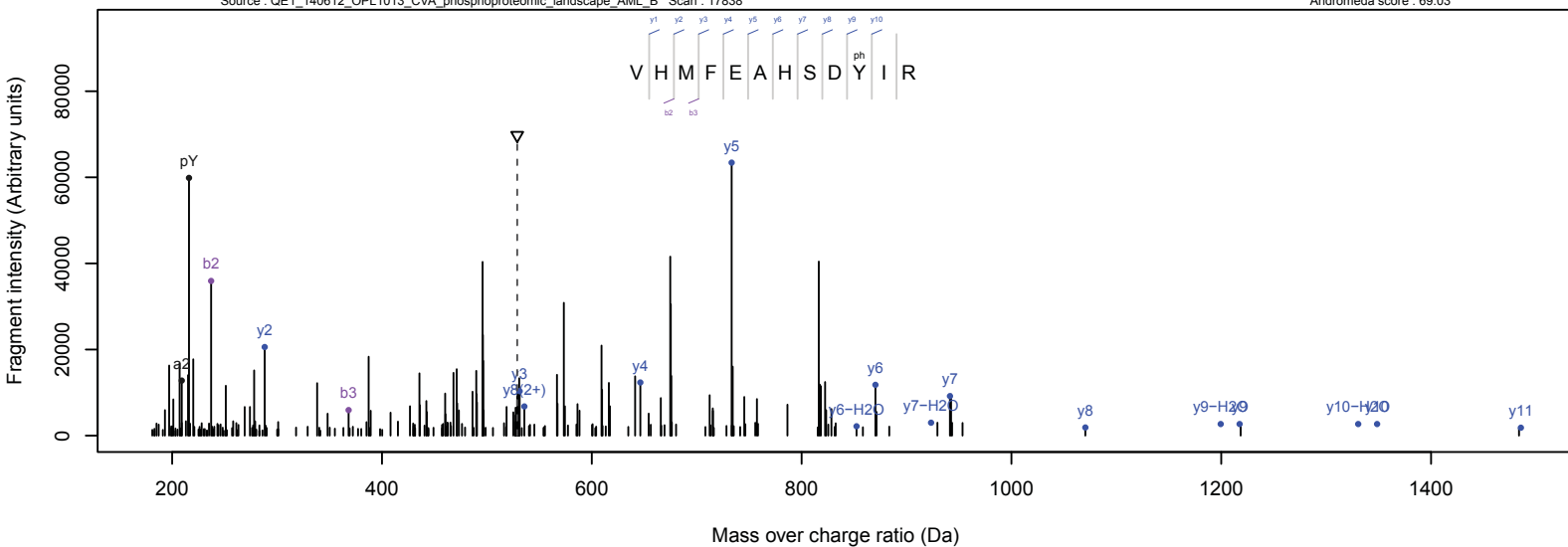




**VHMFEAHSY(ph)IR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17838

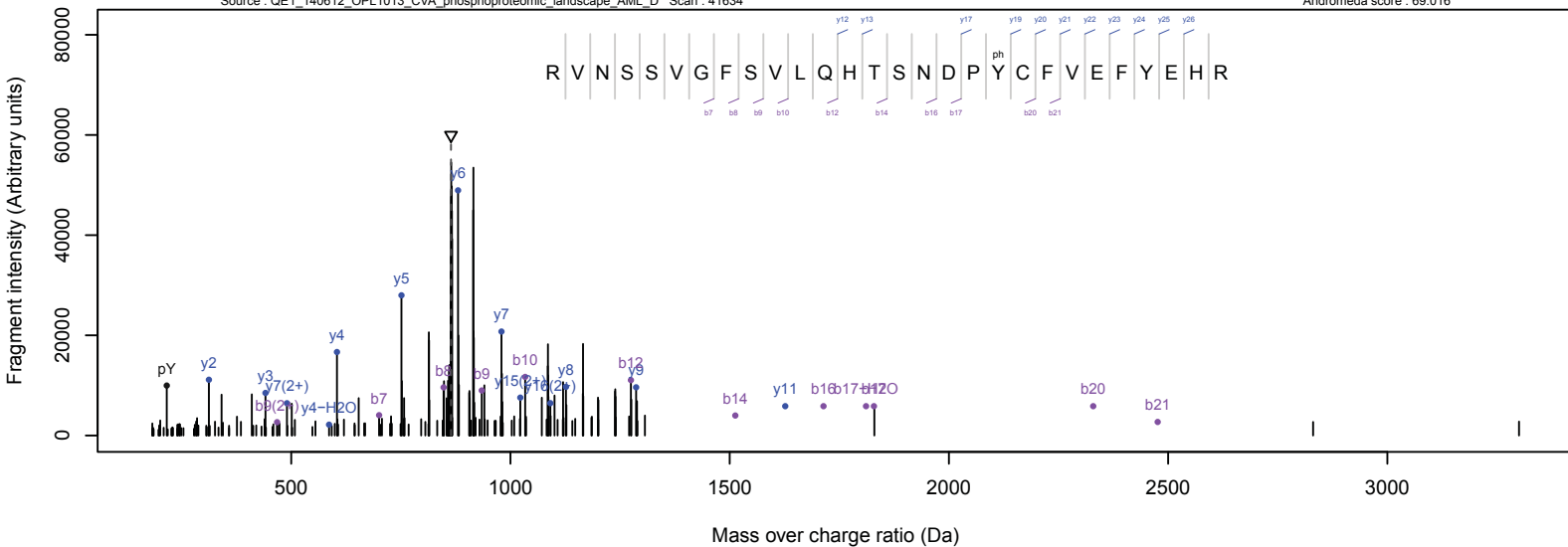
Andromeda score : 69.03



**RVNSSVGFSLQHTSNDP(ph)CFVEFYEHR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41634

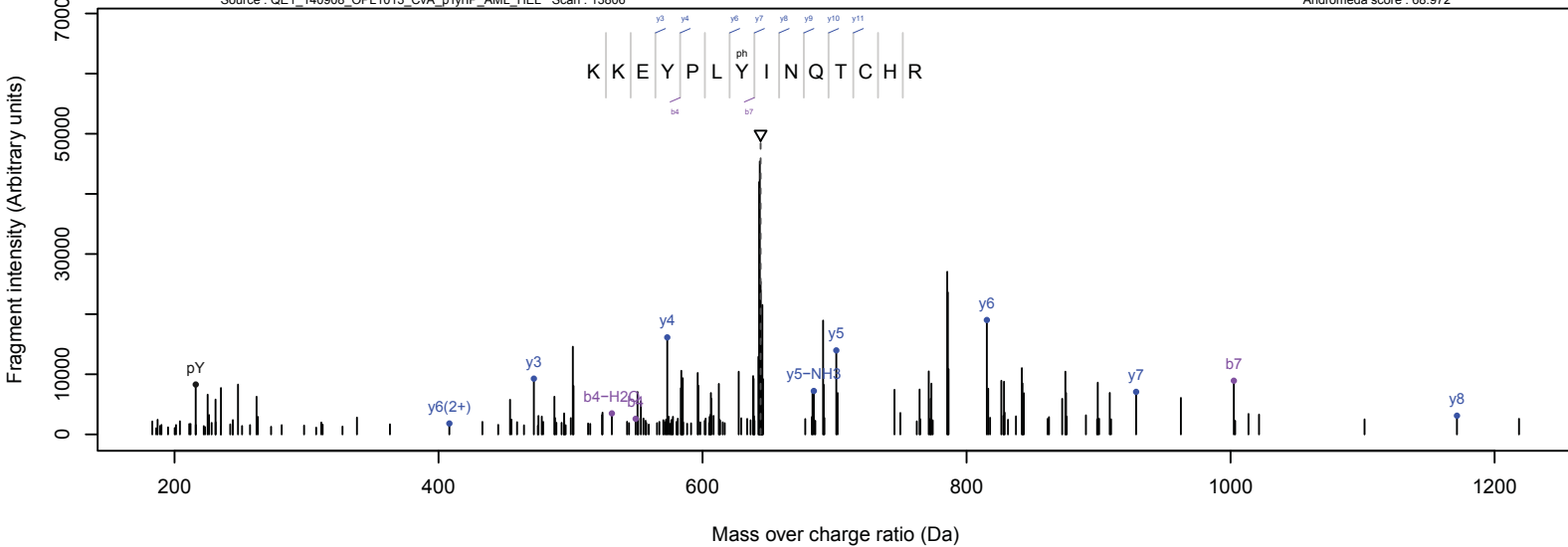
Andromeda score : 69.016



**KKEYPLY(ph)INQTCR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 13806

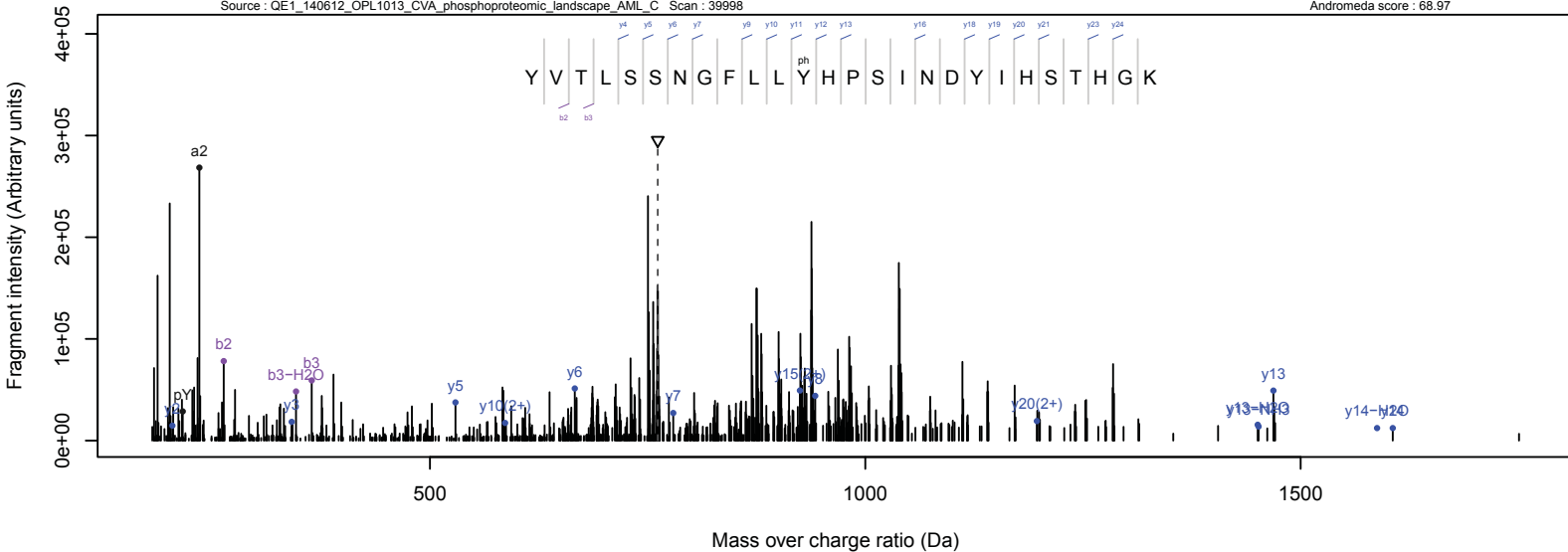
Andromeda score : 68.972



## YVTLSSNGFLLY(ph)HPSINDYIHSTHGK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39998

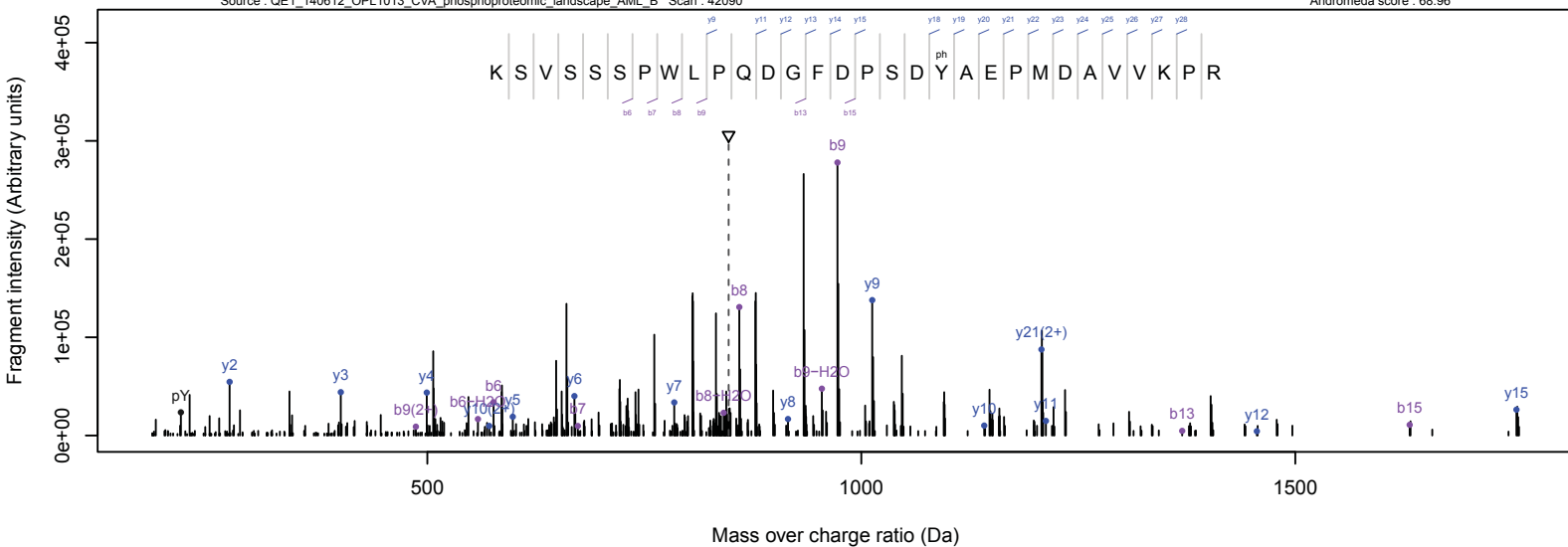
Andromeda score : 68.97



## KSVSSSPWLPQDGFDPSPDY(ph)AEPMDAVVKPR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42090

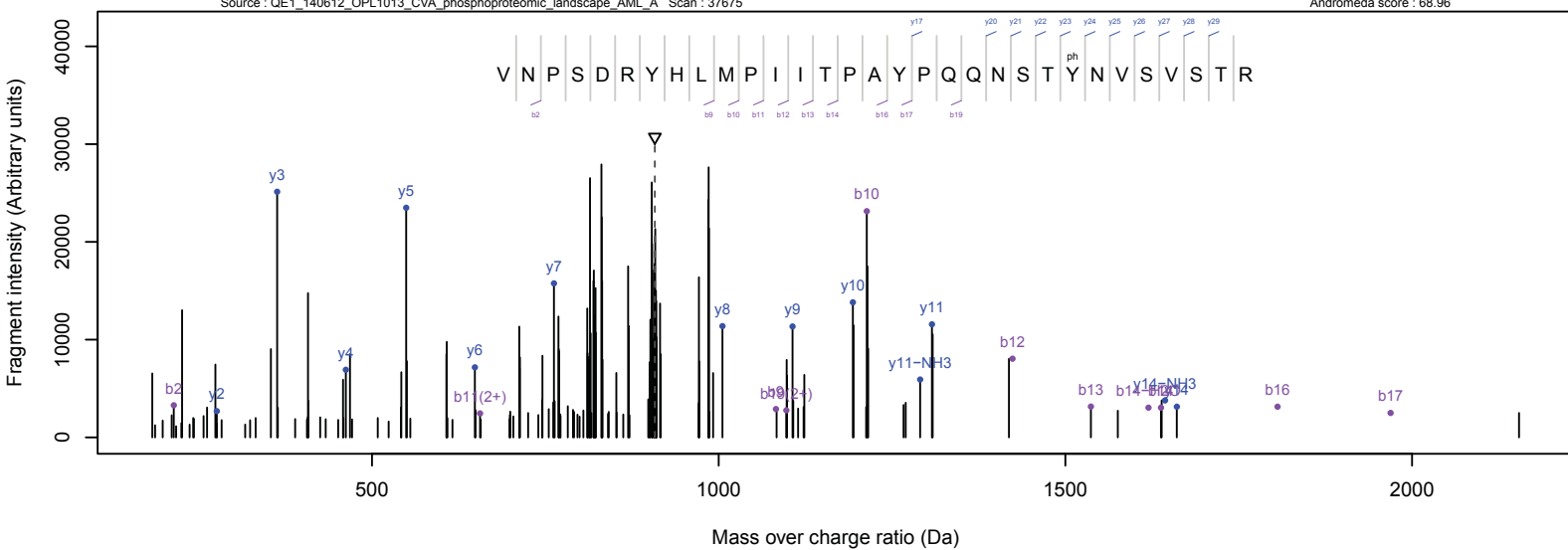
Andromeda score : 68.96



## VNPSDRYHLMPIITPAYPQQNSTY(ph)NVSSTR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 37675

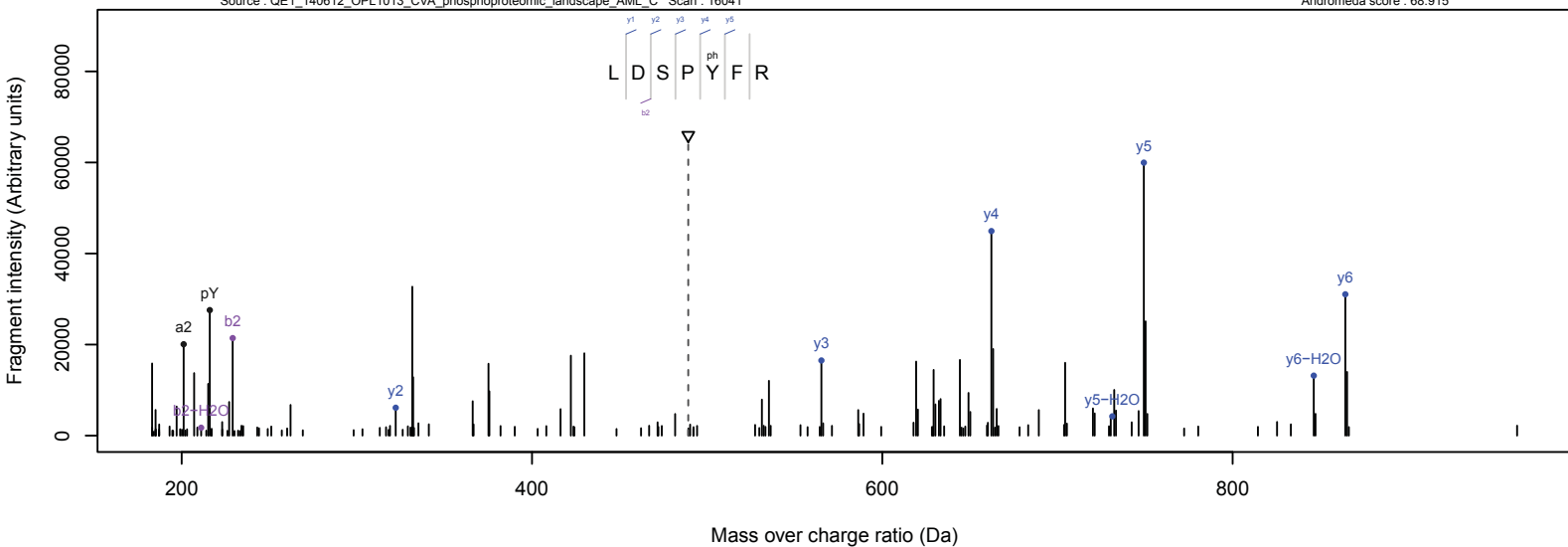
Andromeda score : 68.96



**LDSPY(ph)FR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16041

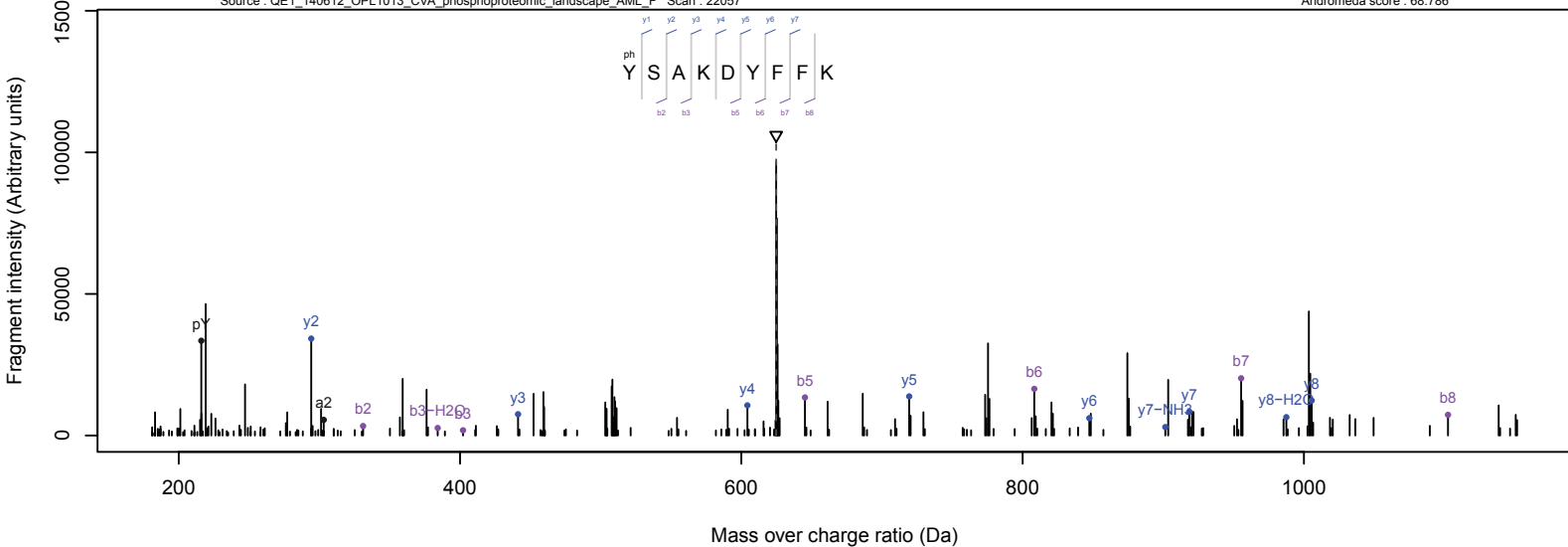
Andromeda score : 68.915



**Y(ph)SAKDYFFK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 22057

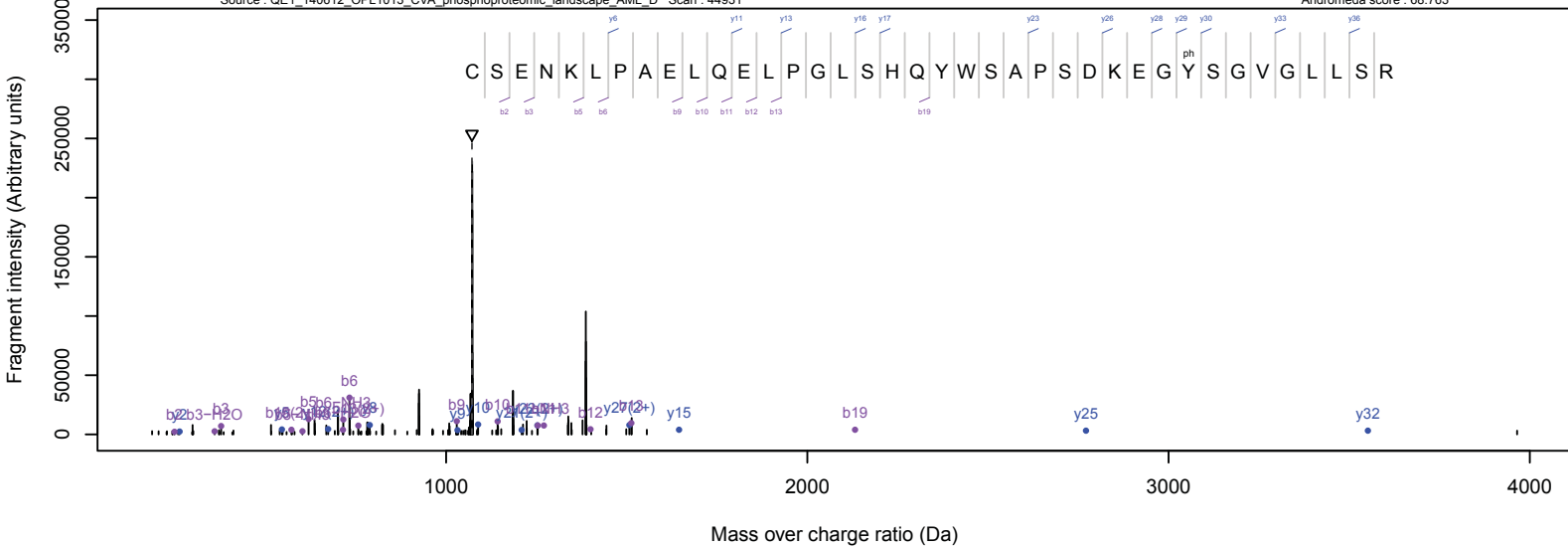
Andromeda score : 68.786



**CSENKLP AELQELPGLSHQYWSAPS DKEGY(ph)SGVGLLSR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 44931

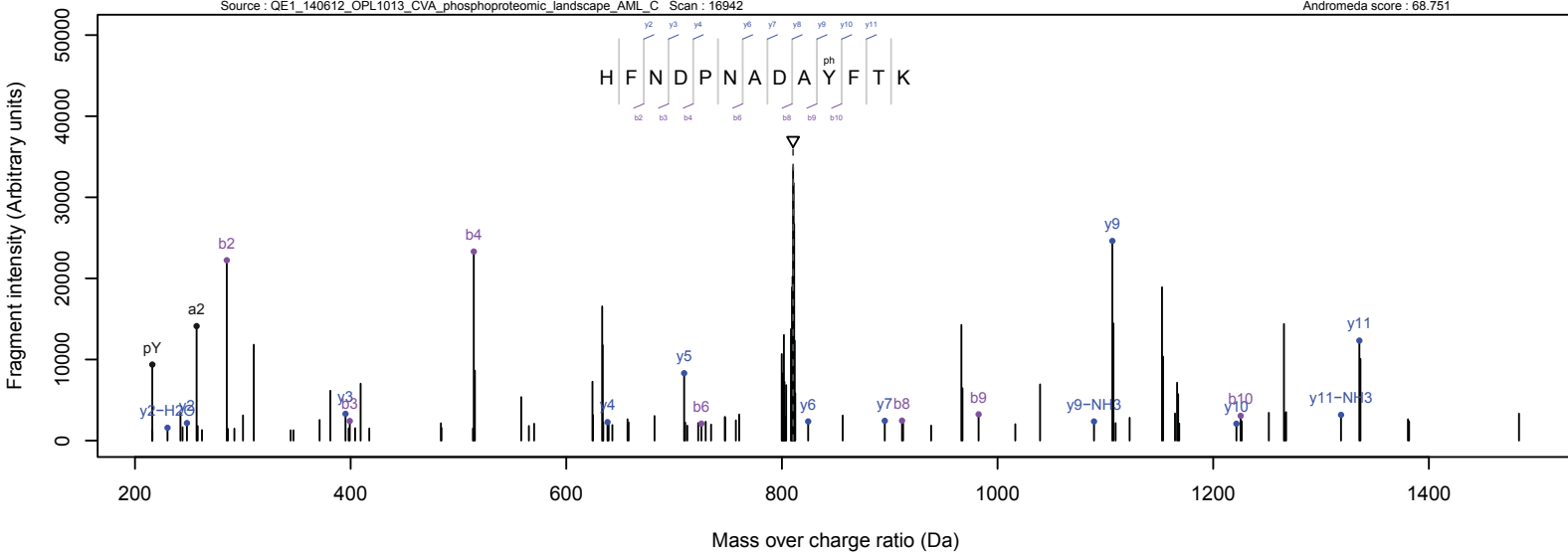
Andromeda score : 68.763



**HFNDPNADAYFTK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16942

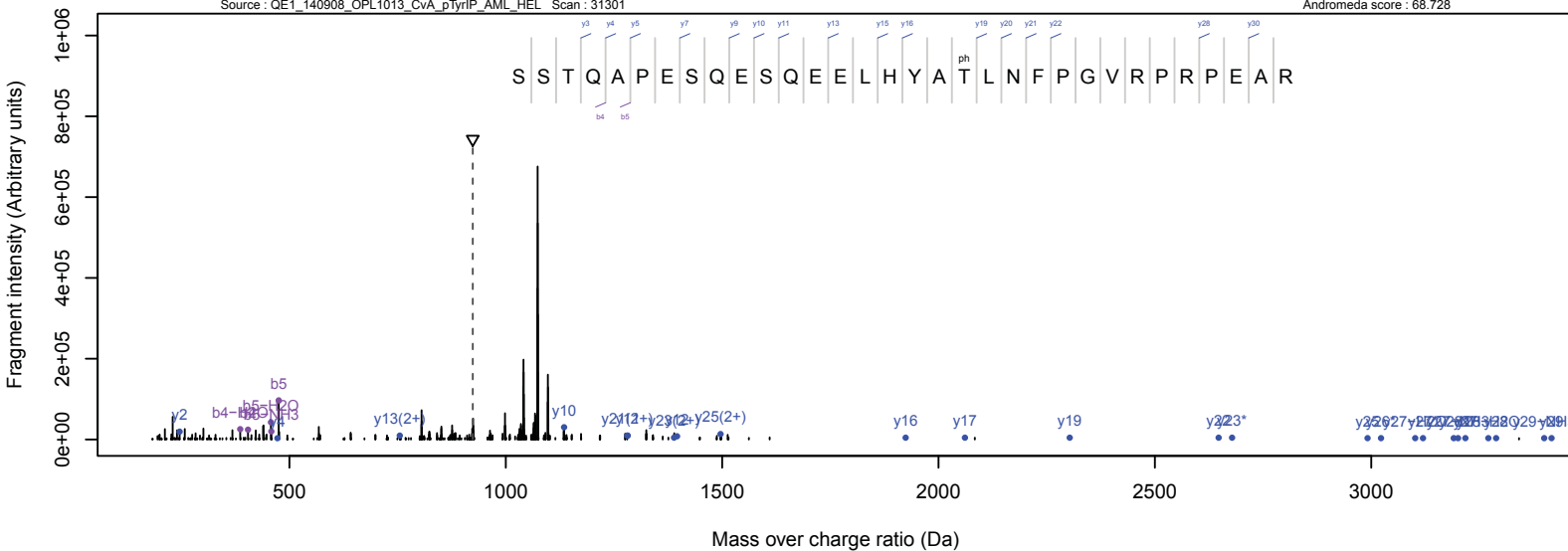
Andromeda score : 68.751



**SSTQAPESQESQEEELHYATLNFGVGRPRPEAR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31301

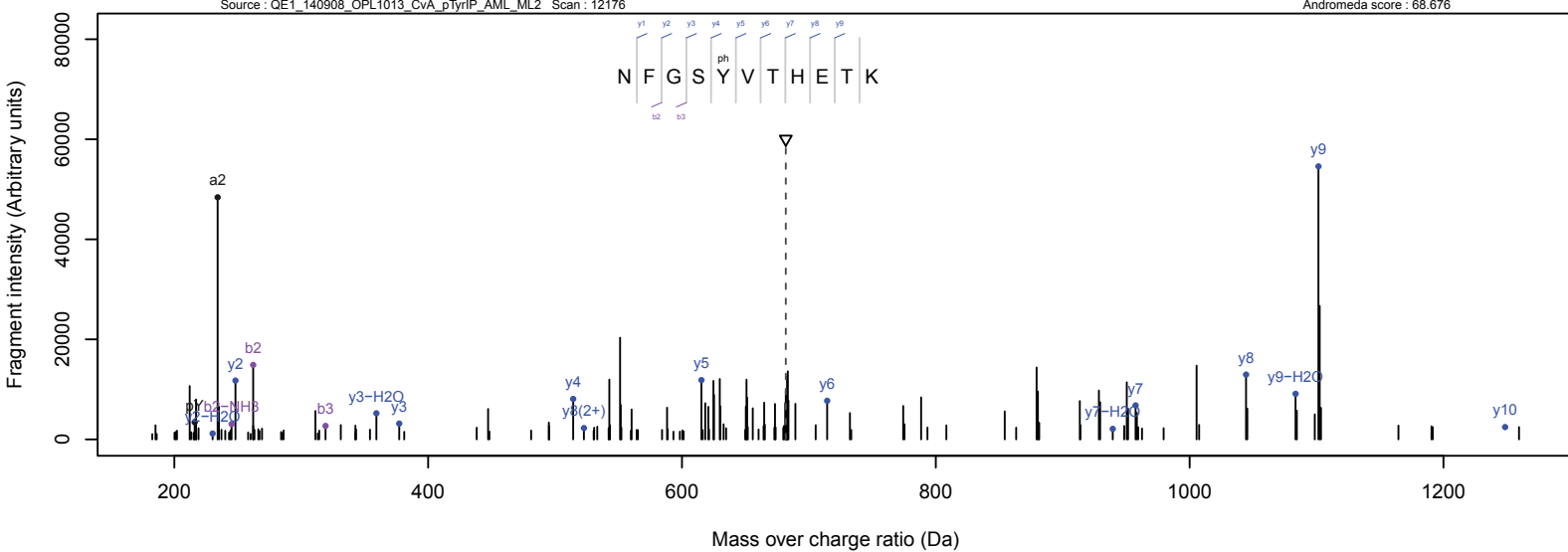
Andromeda score : 68.728



**NFGSYVTHETK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 12176

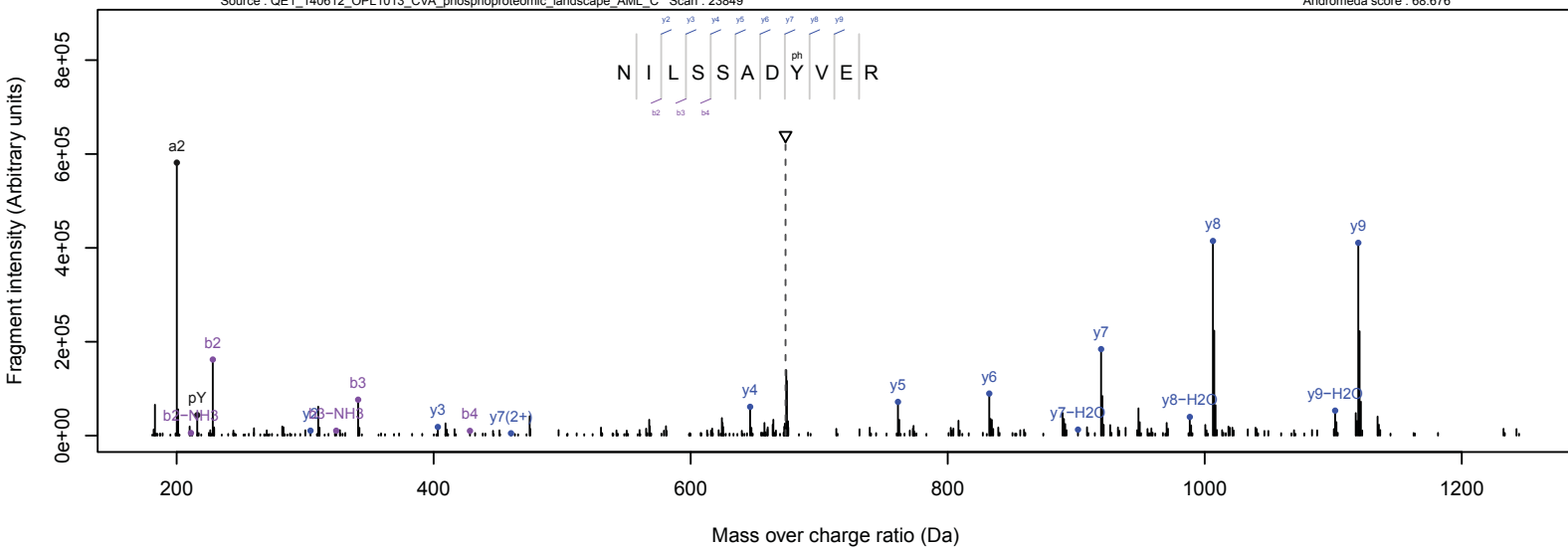
Andromeda score : 68.676



NILSSADY(ph)VER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23849

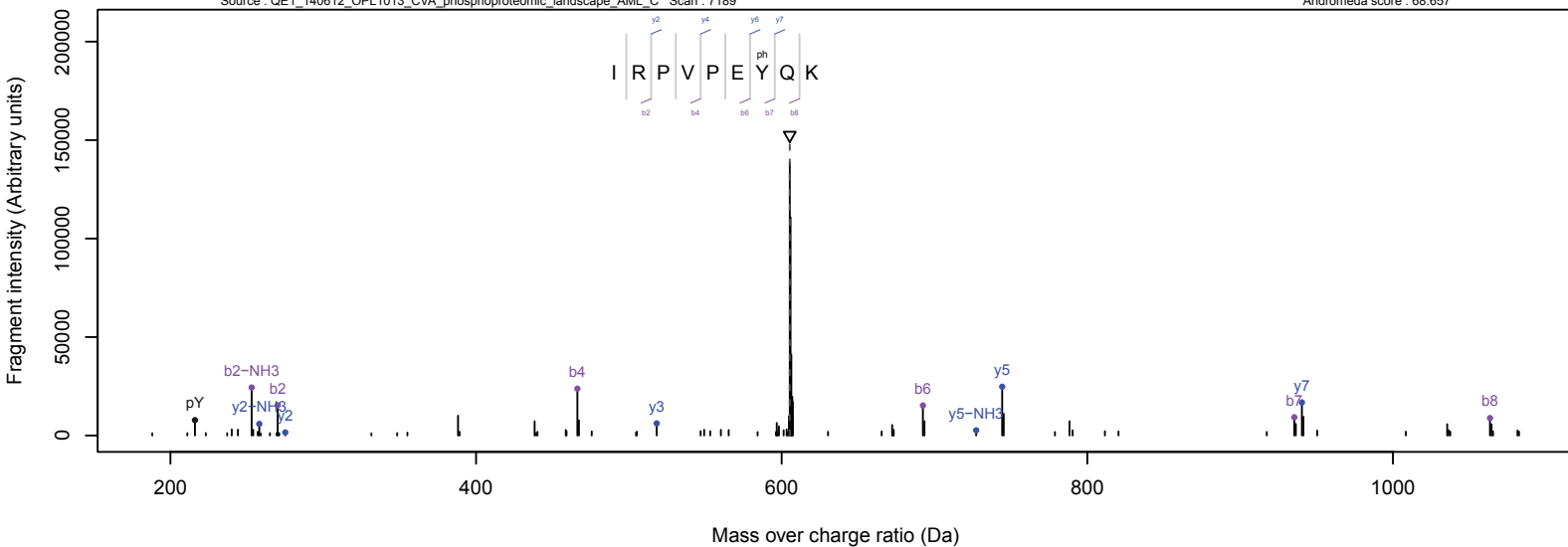
Andromeda score : 68.676



IRPVPEY(ph)QK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7189

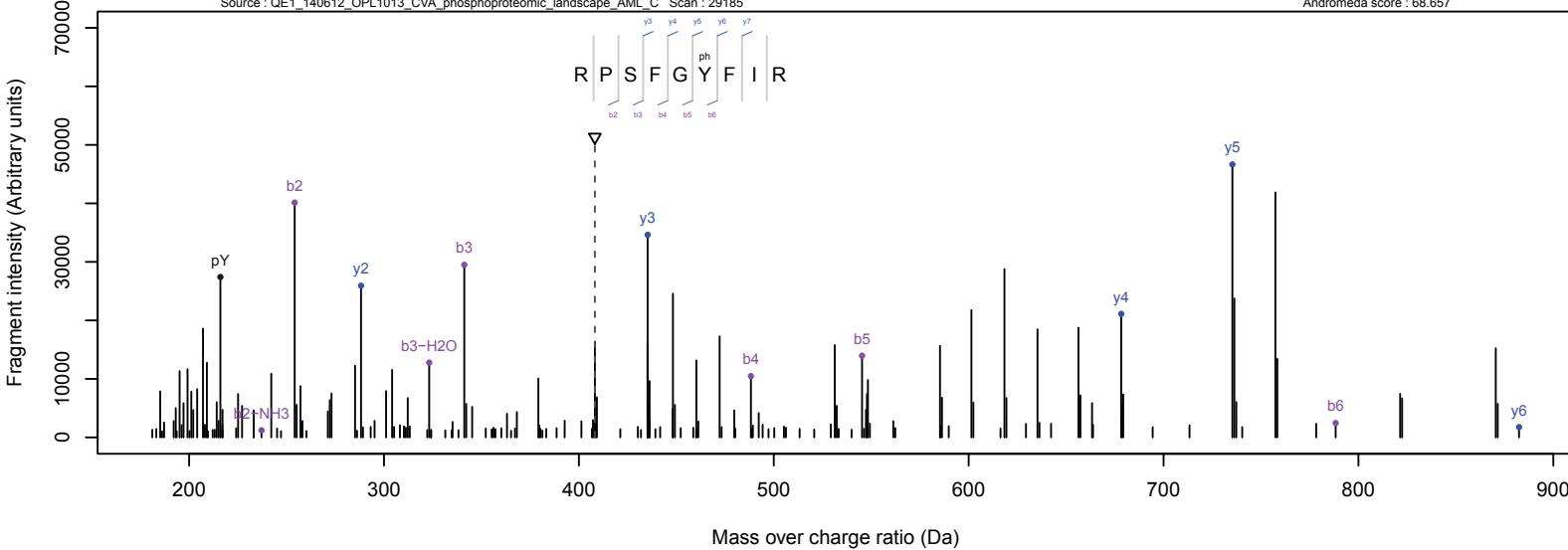
Andromeda score : 68.657



RPSFGY(ph)FIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29185

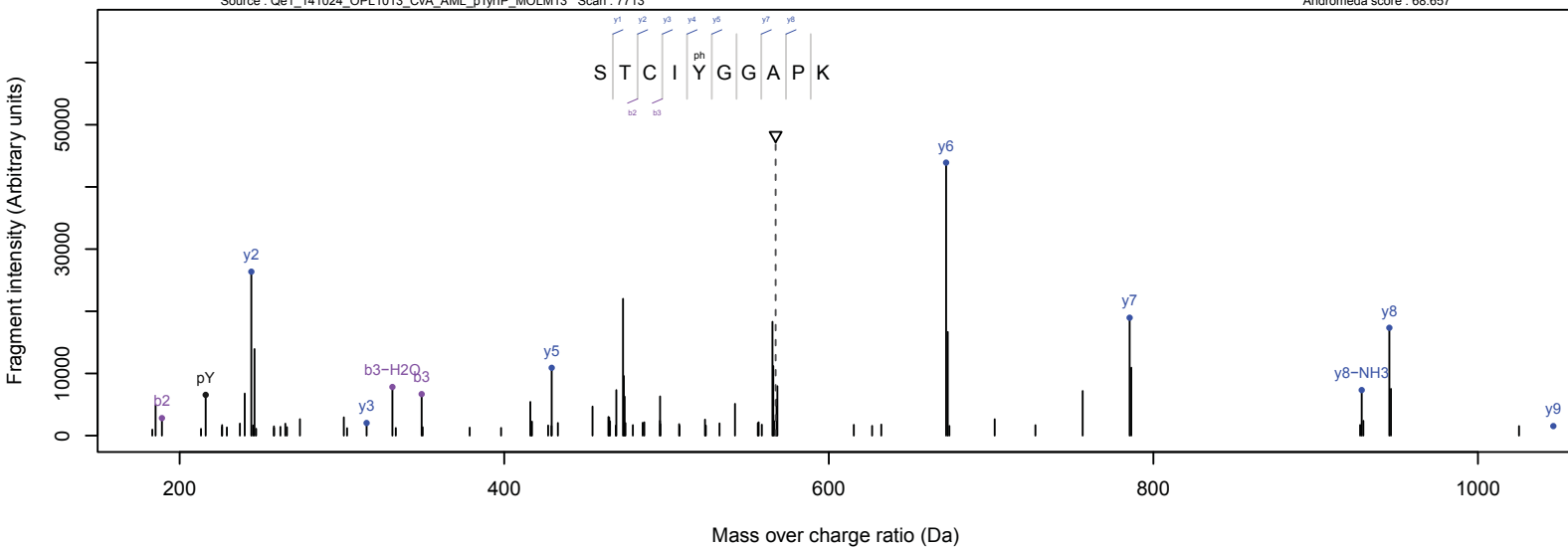
Andromeda score : 68.657



## STCIY(ph)GGAPK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 7713

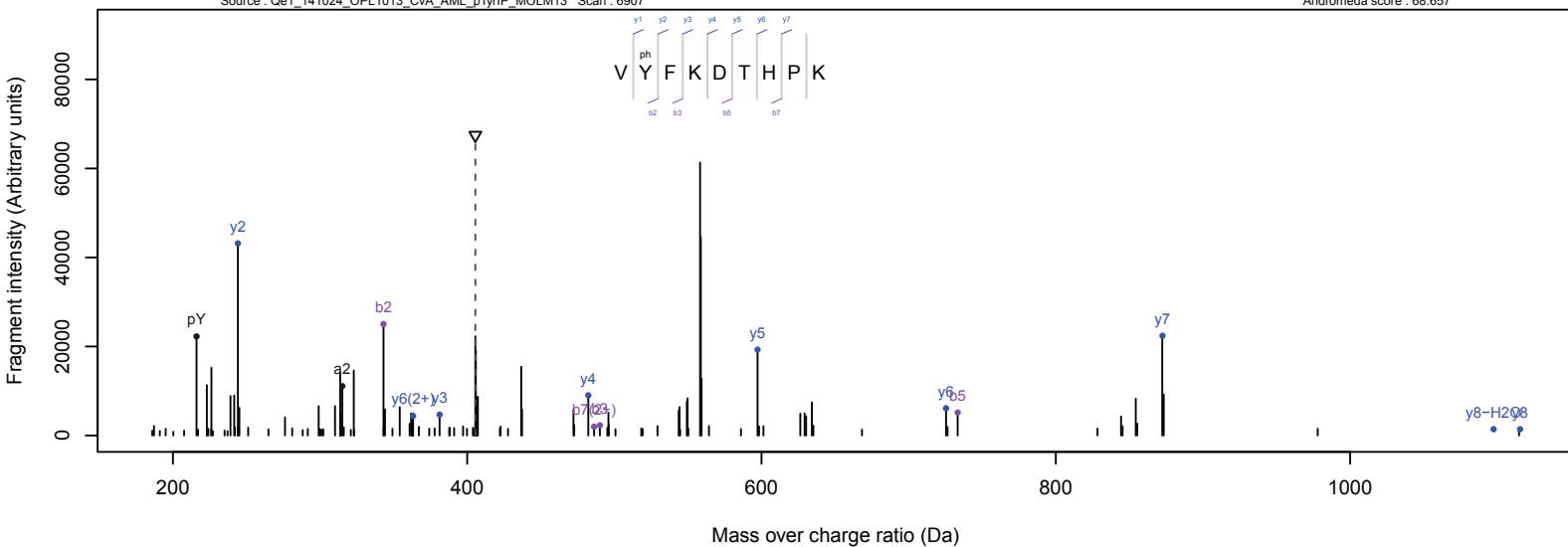
Andromeda score : 68.657



## VY(ph)FKDTHPK-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 6907

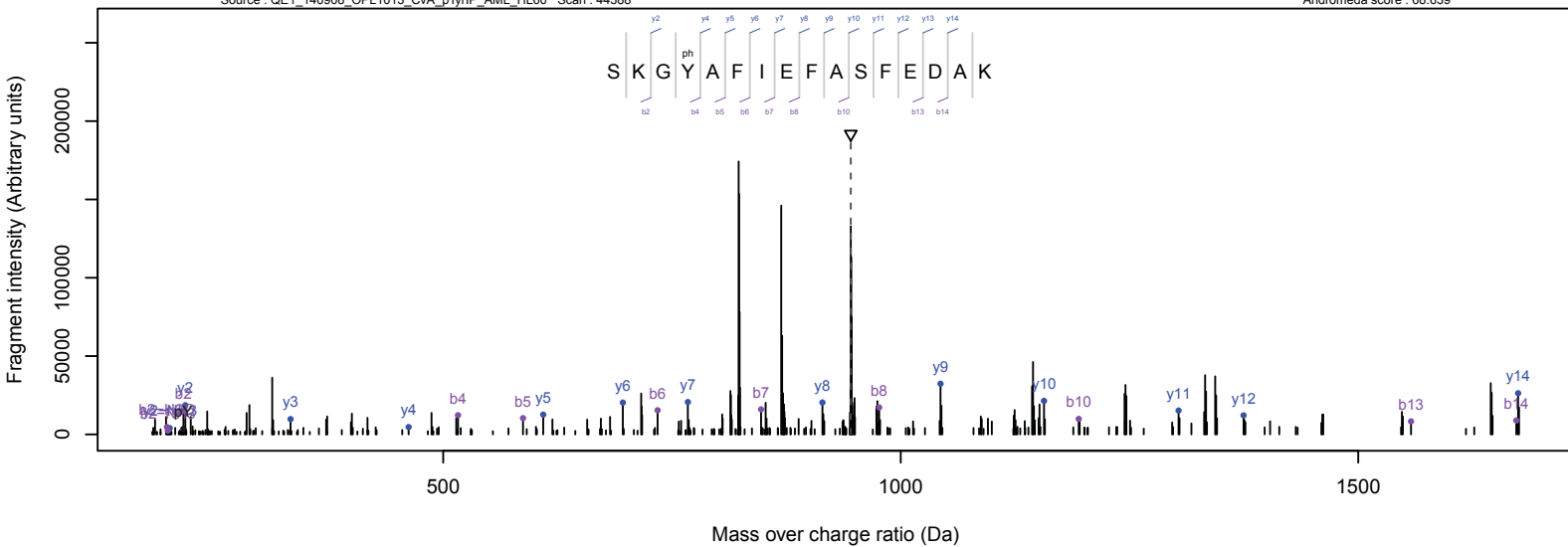
Andromeda score : 68.657



## SKGY(ph)AFIEFASFE DAK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 44388

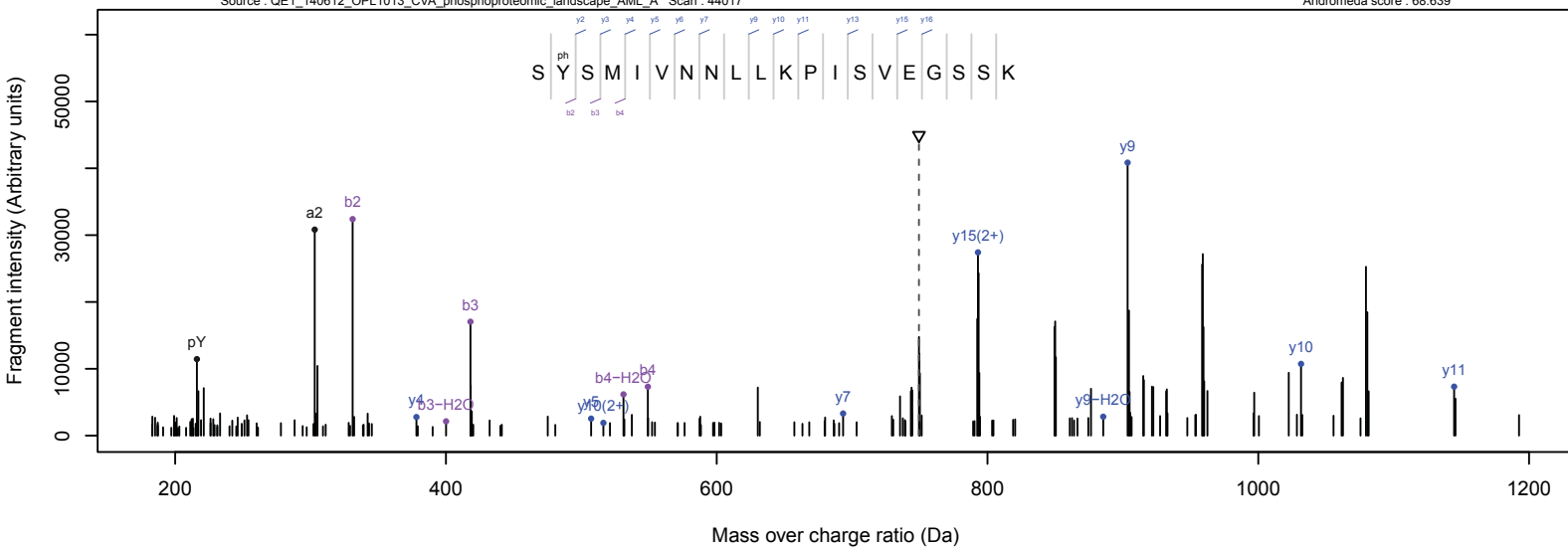
Andromeda score : 68.639



## SY(ph)SMIVNNLLKPISEVGSSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 44017

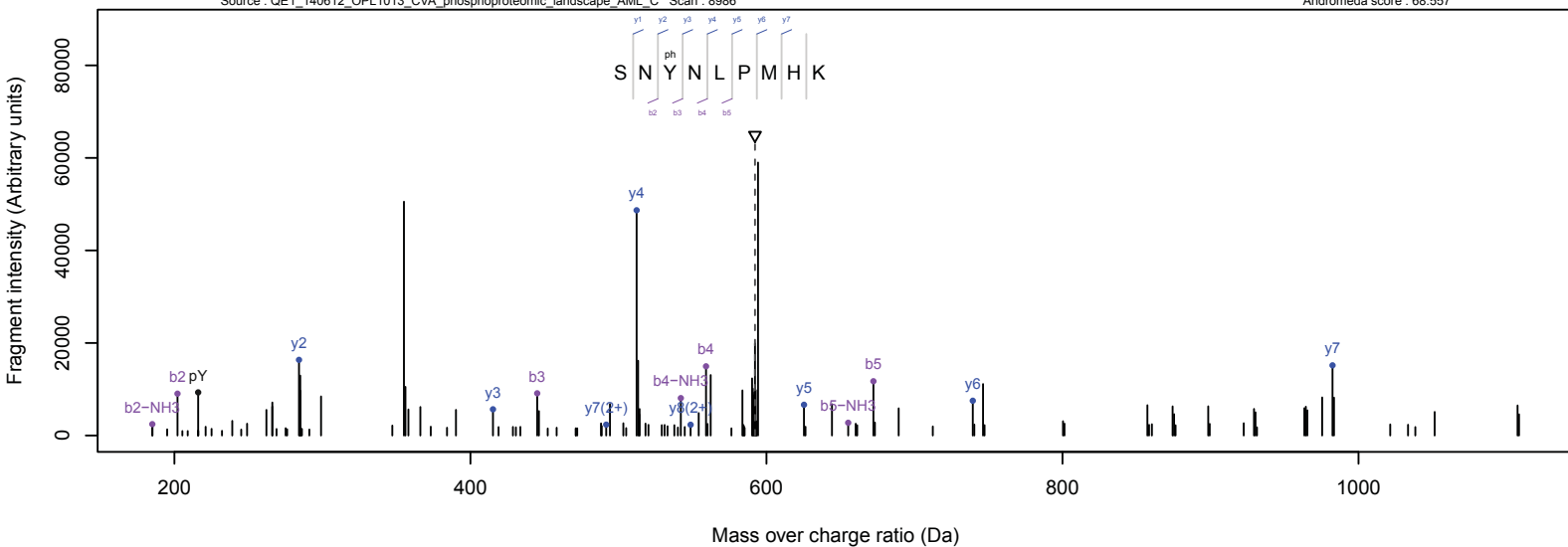
Andromeda score : 68.639



## SNY(ph)NLPMHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8986

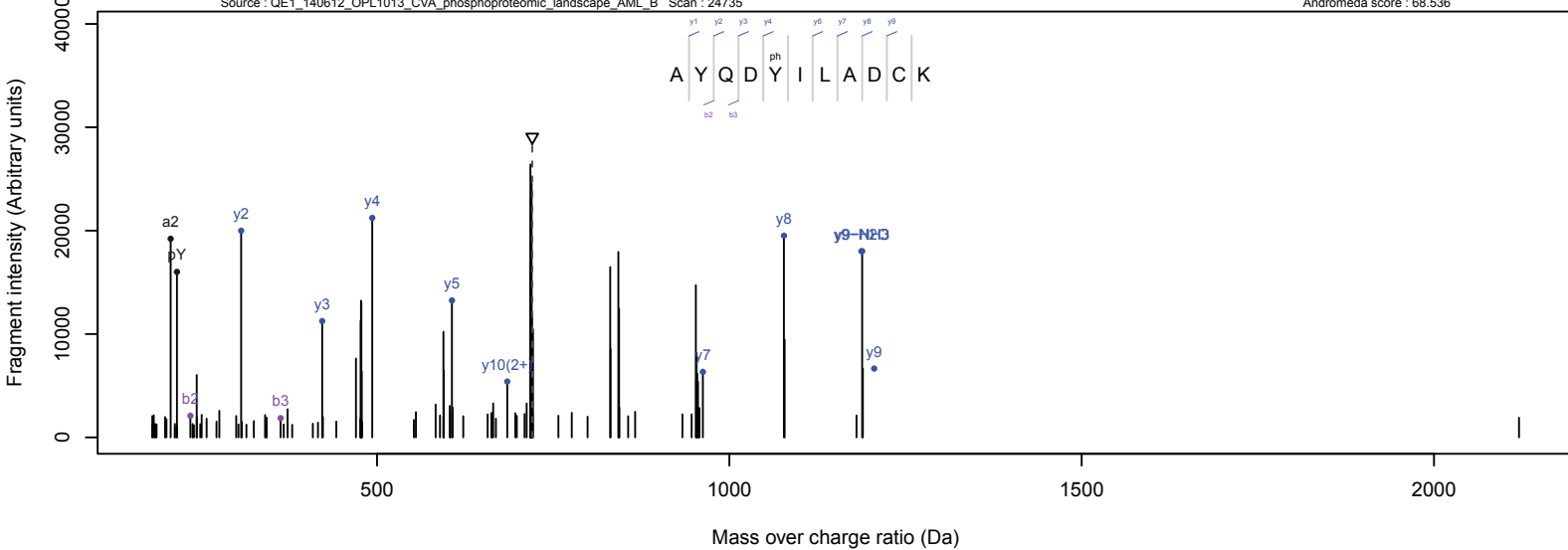
Andromeda score : 68.557



## AYQDY(ph)ILADCK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24735

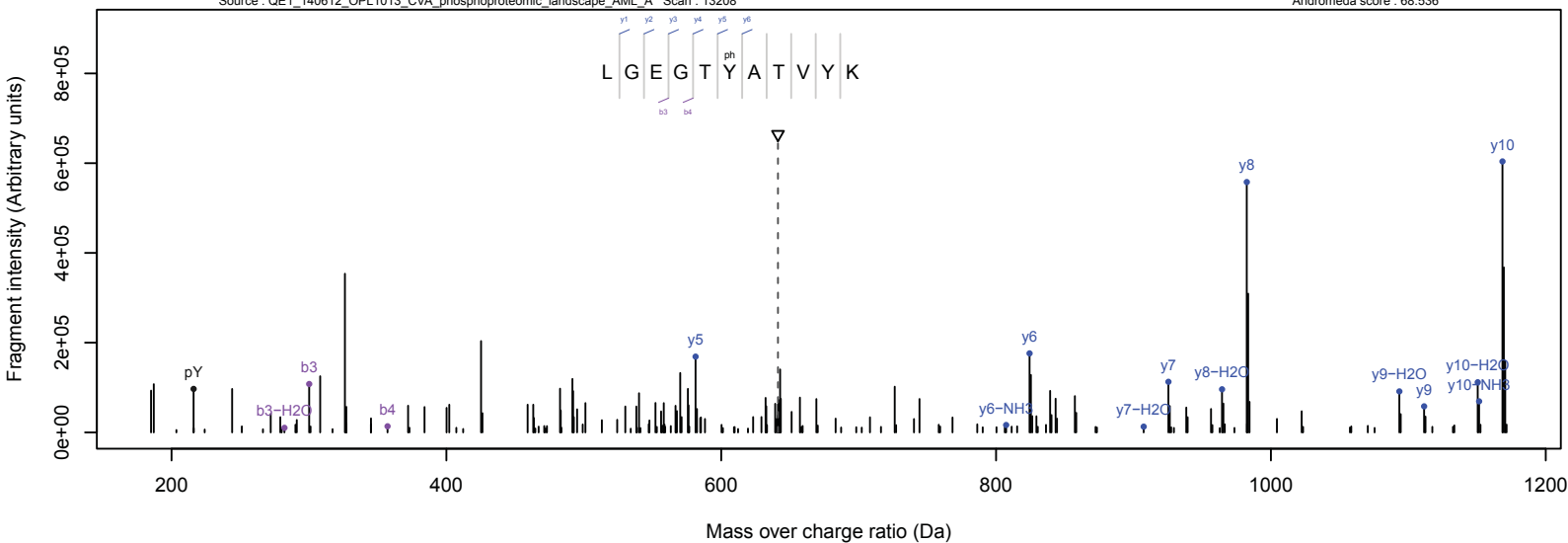
Andromeda score : 68.536



## LGEGTY(ph)ATVYK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 13208

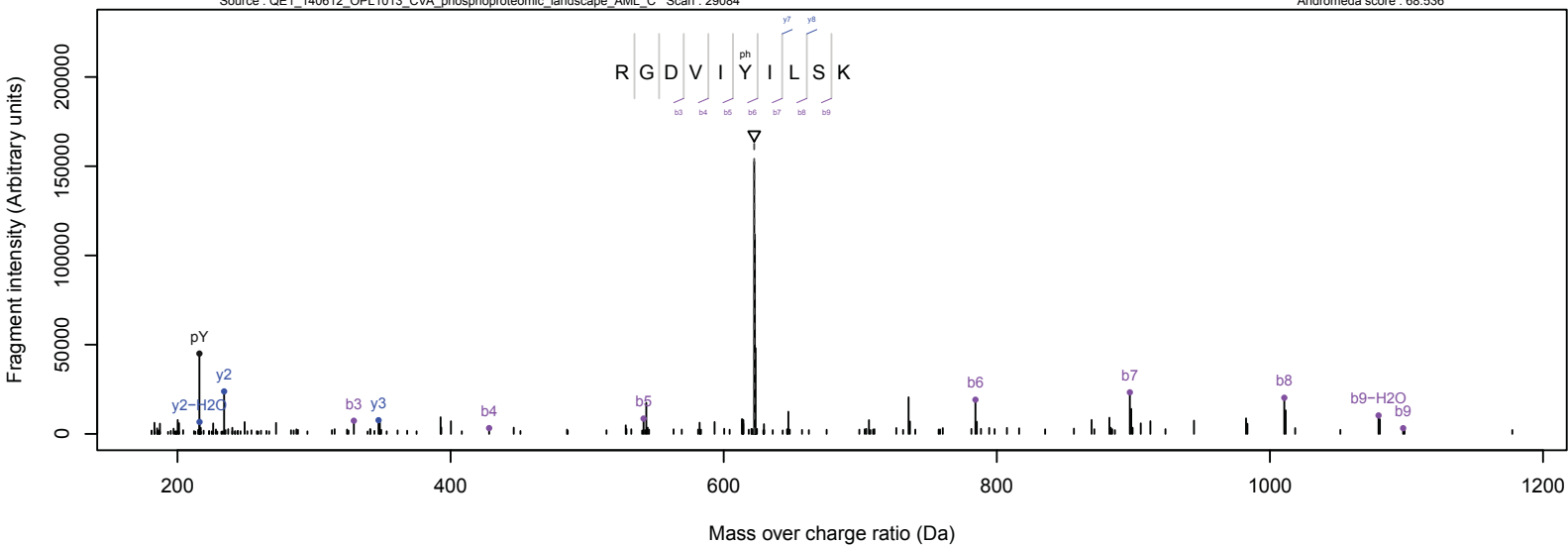
Andromeda score: 68.536



## RGDVIY(ph)ILSK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 29084

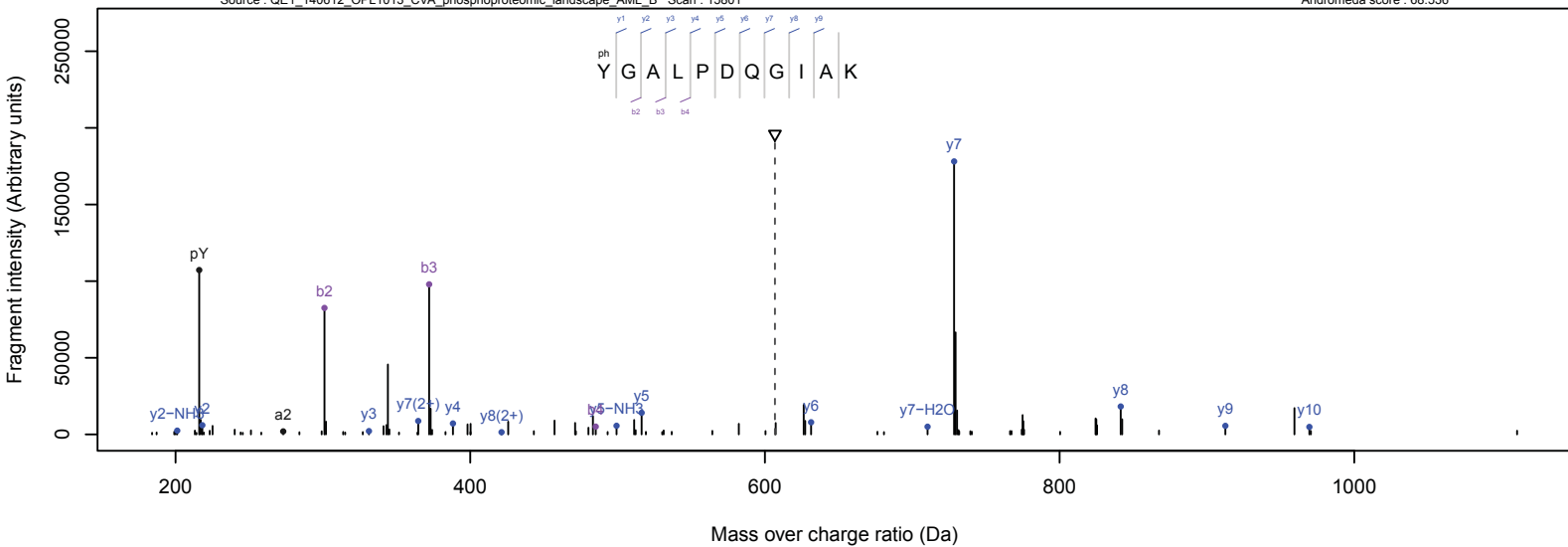
Andromeda score: 68.536



## Y(ph)GALPDQGIK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15801

Andromeda score: 68.536

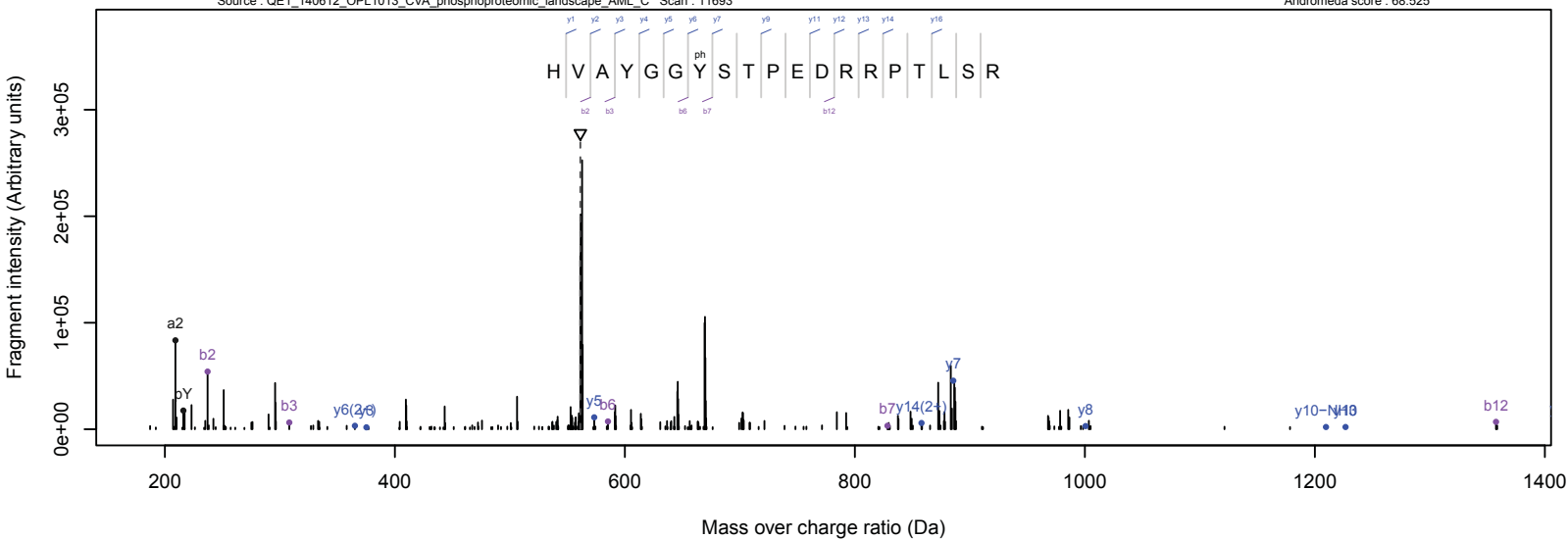




## HVAYGGY(ph)STPEDRRPTLSR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11693

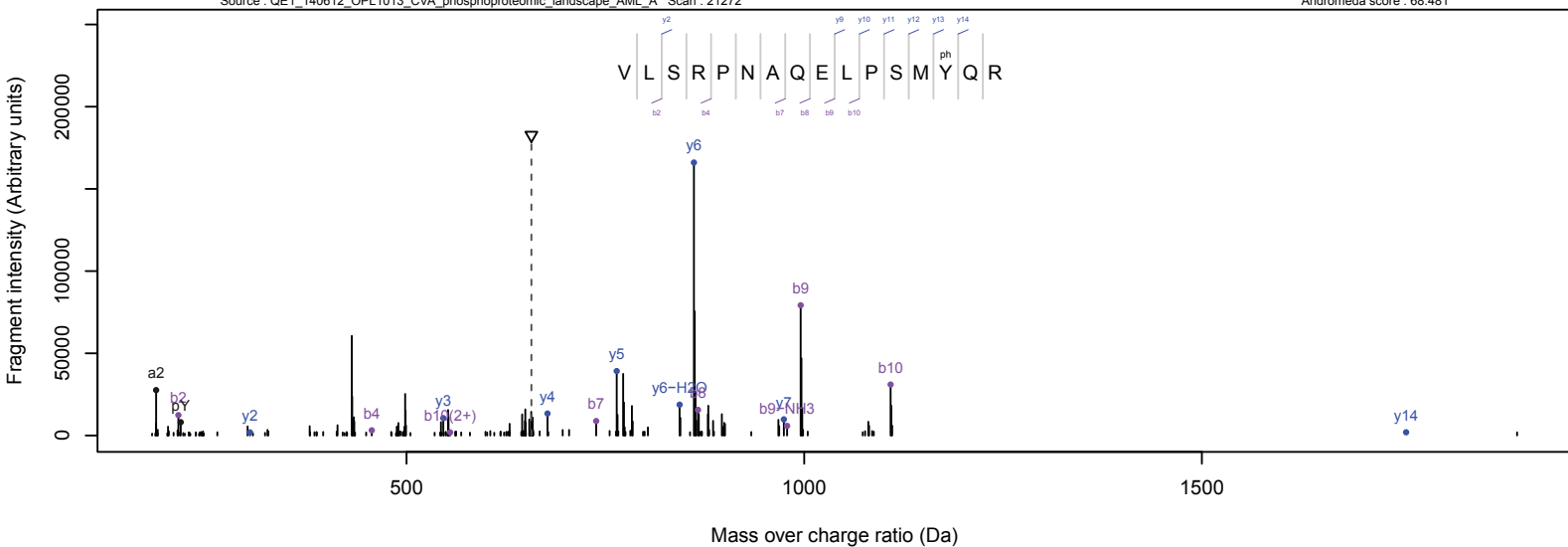
Andromeda score: 68.525



## VLSRPNAQELPSMY(ph)QR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 21272

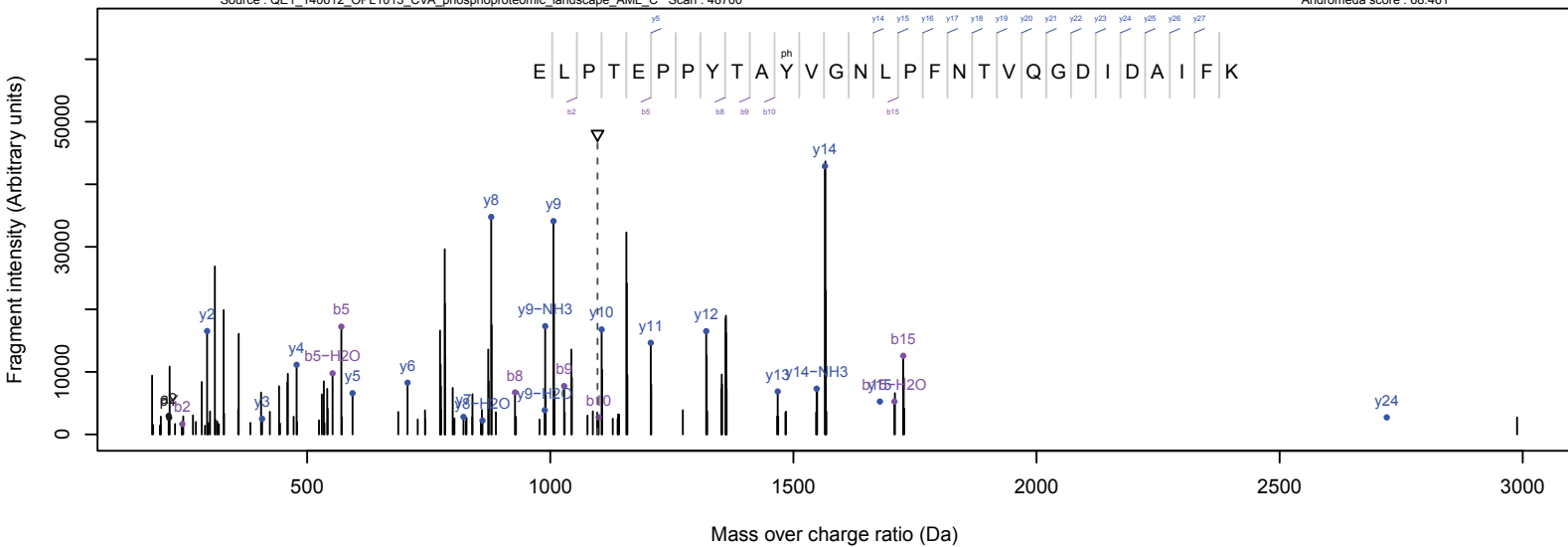
Andromeda score: 68.481



## ELPTEPPYTA(ph)YVGNLPFNTVQGDIDAIFK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 48700

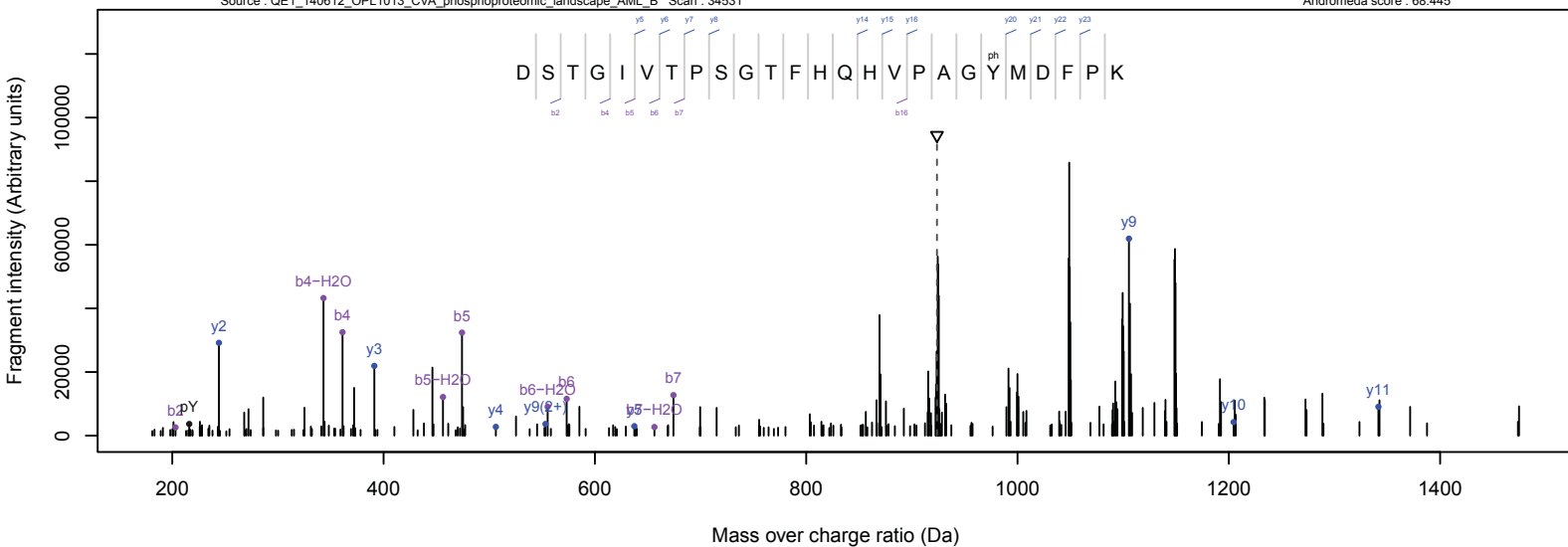
Andromeda score: 68.461



## DSTGIVTSPGTFHQHVPAGY(ph)MDFPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34531

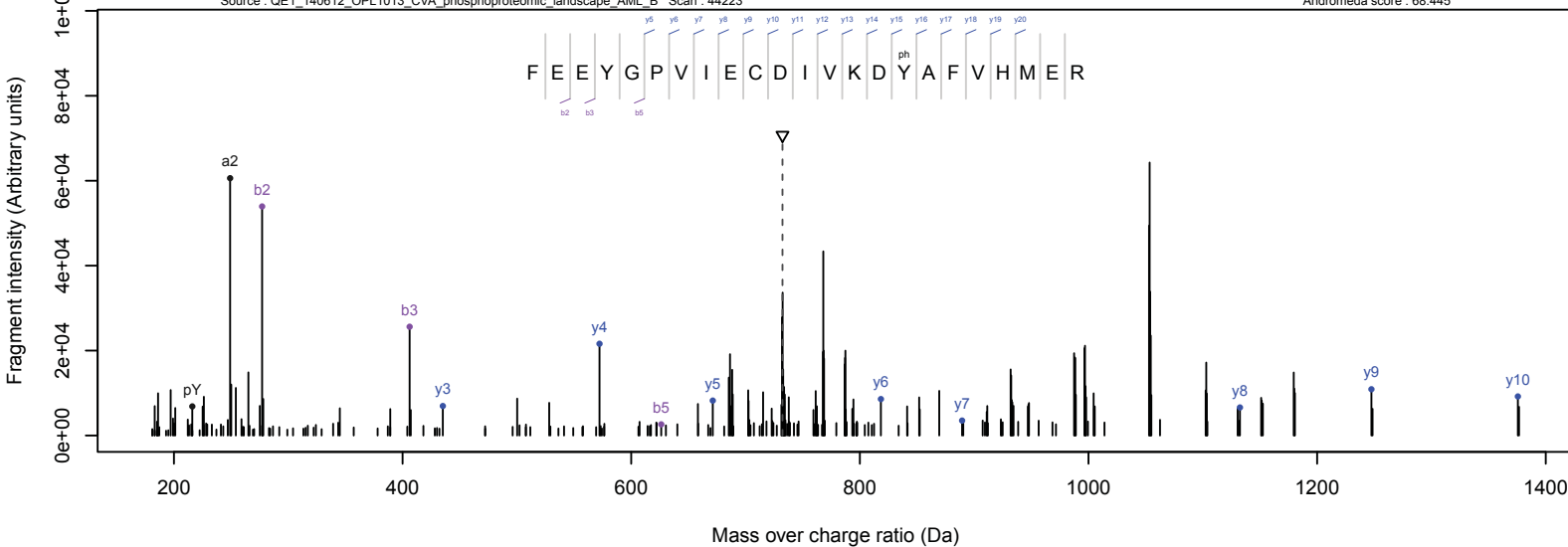
Andromeda score : 68.445



## FEEYGPVIECDIVKDY(ph)AFVHMER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44223

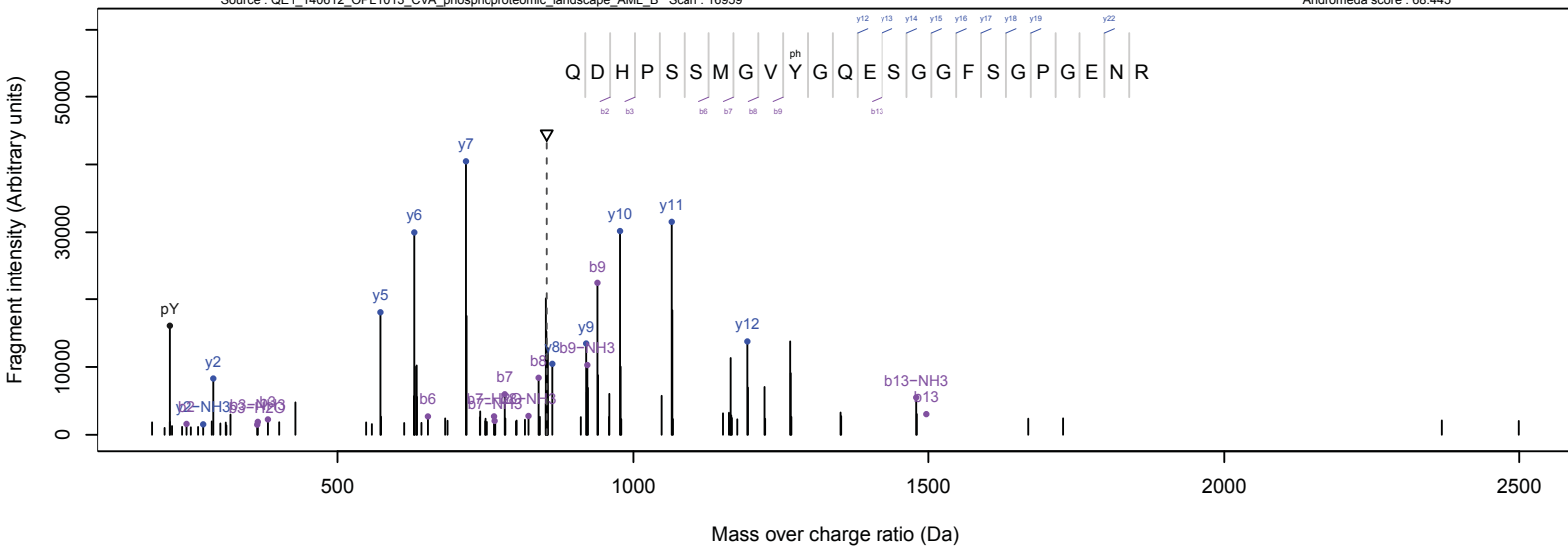
Andromeda score : 68.445



## QDHPSSMGVY(ph)QESGGFSGPGENR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16959

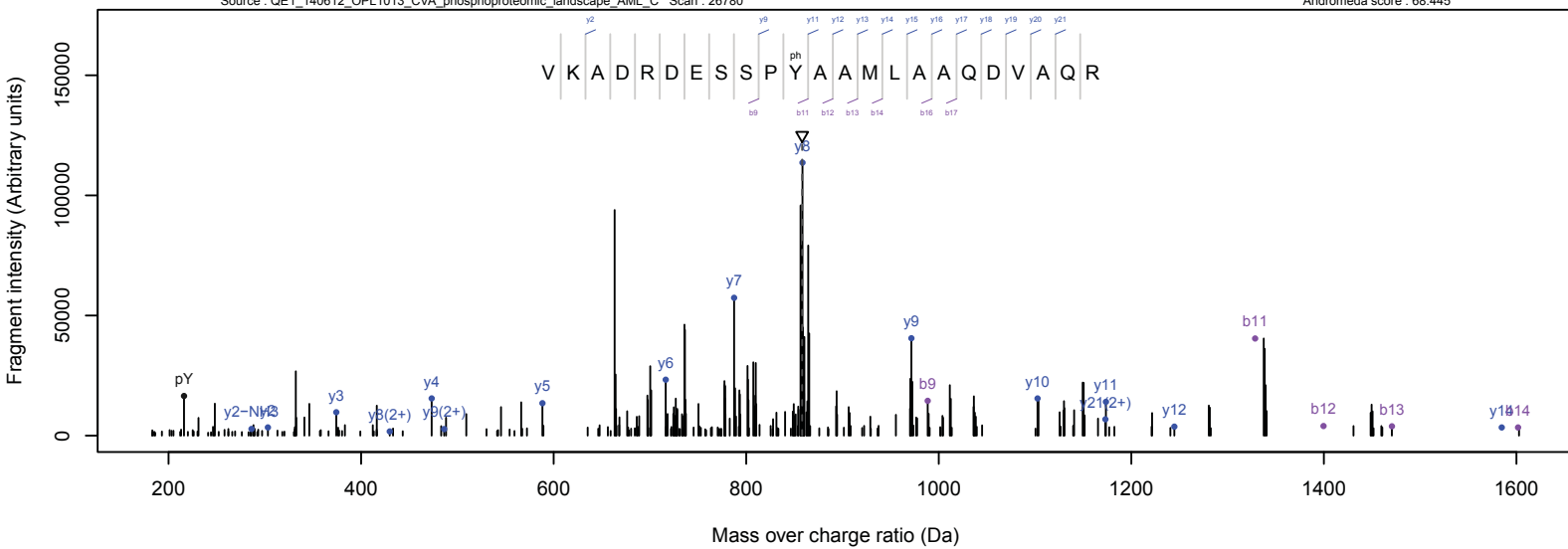
Andromeda score : 68.445



## VKADRDESSPY(ph)AAMLAQDVAQR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26780

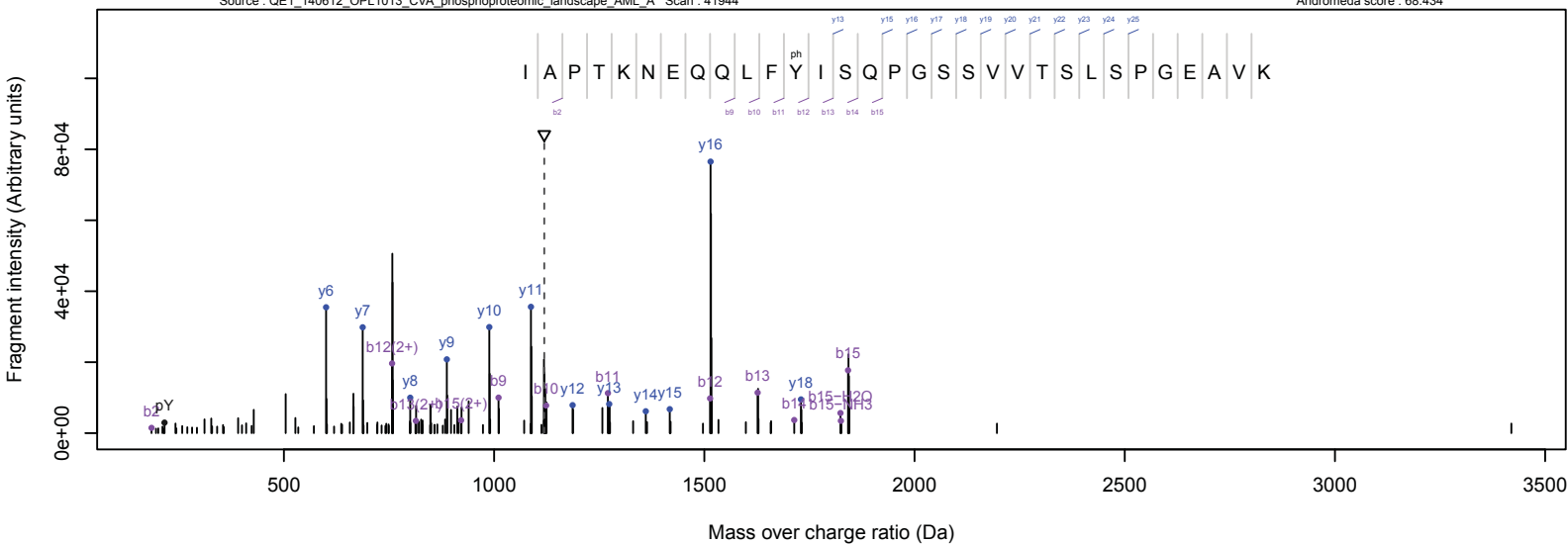
Andromeda score: 68.445



## IAPTKEQLFY(ph)ISQPGSSVVTSLSPGEAVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 41944

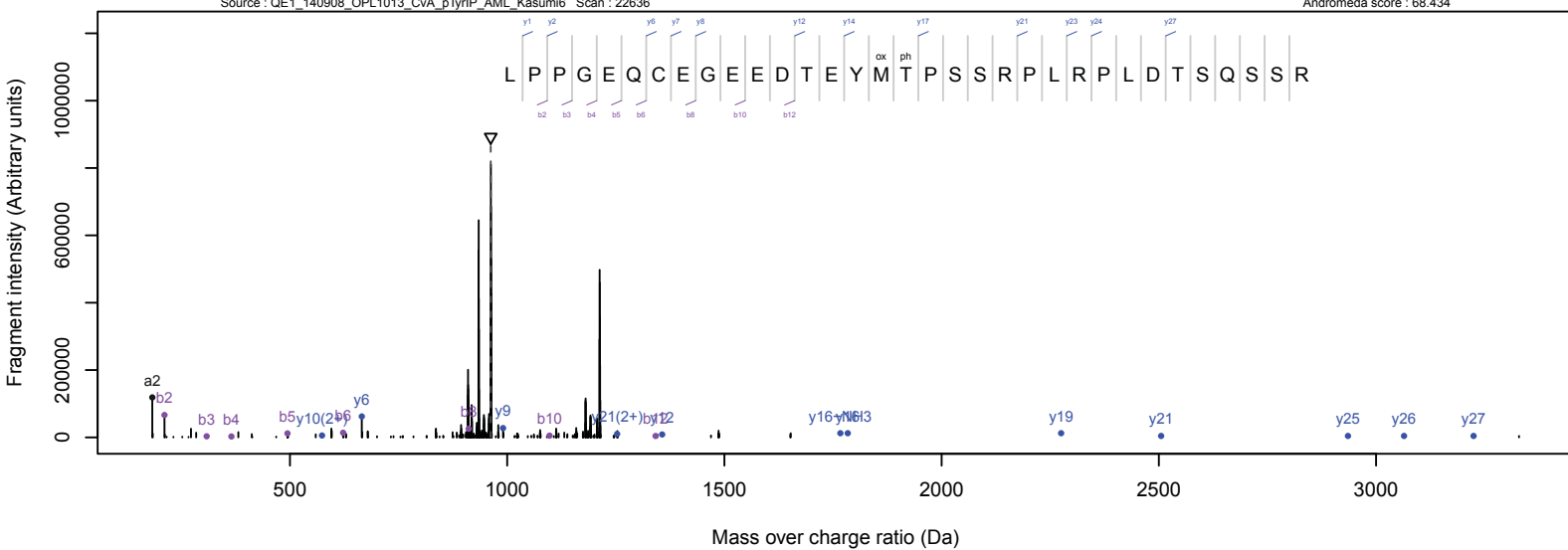
Andromeda score: 68.434



## LPPGEQCEGEDTEYM(ox)T(ph)PSSRPLRLDTSQSSR-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 22636

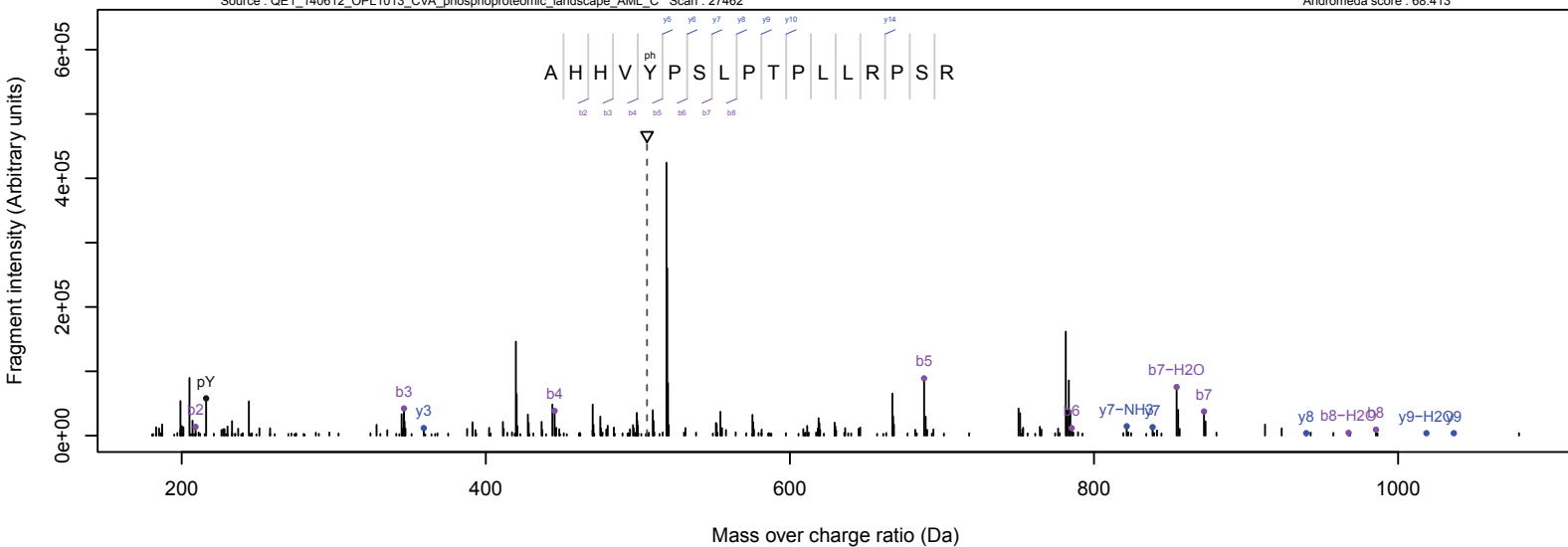
Andromeda score: 68.434



## AHHVY(ph)PSLPTLLRPSR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27462

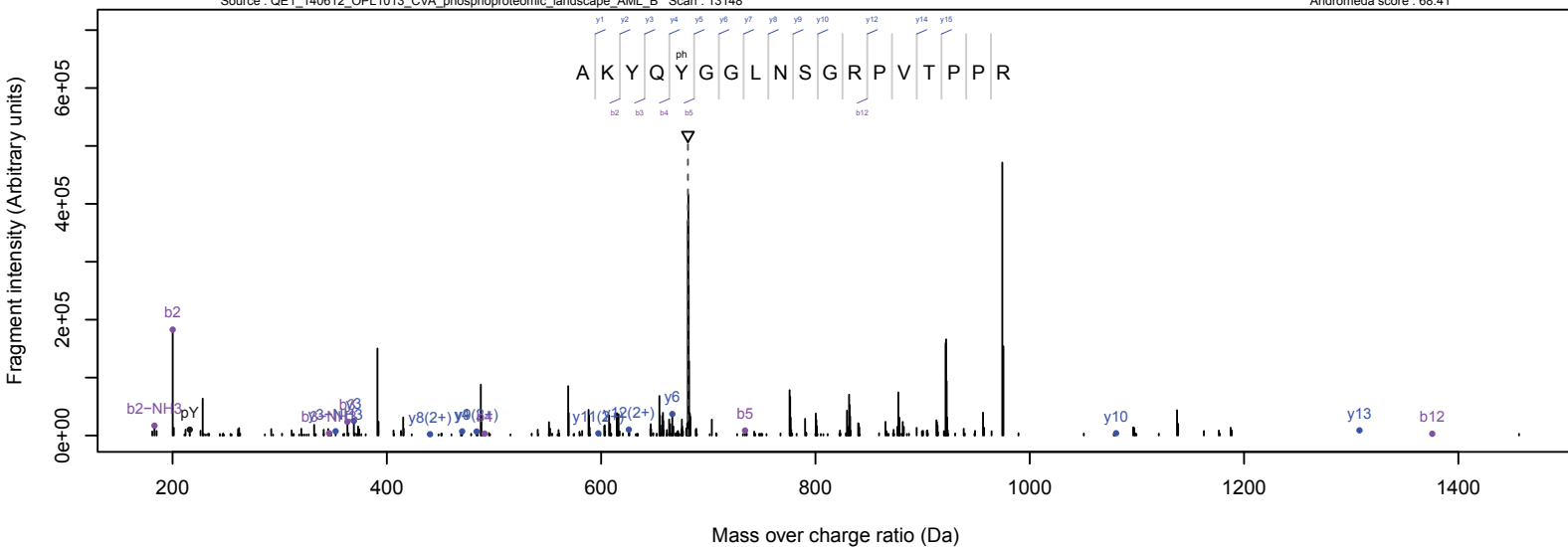
Andromeda score : 68.413



## AKYQY(ph)GGLNSGRPVTTPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13148

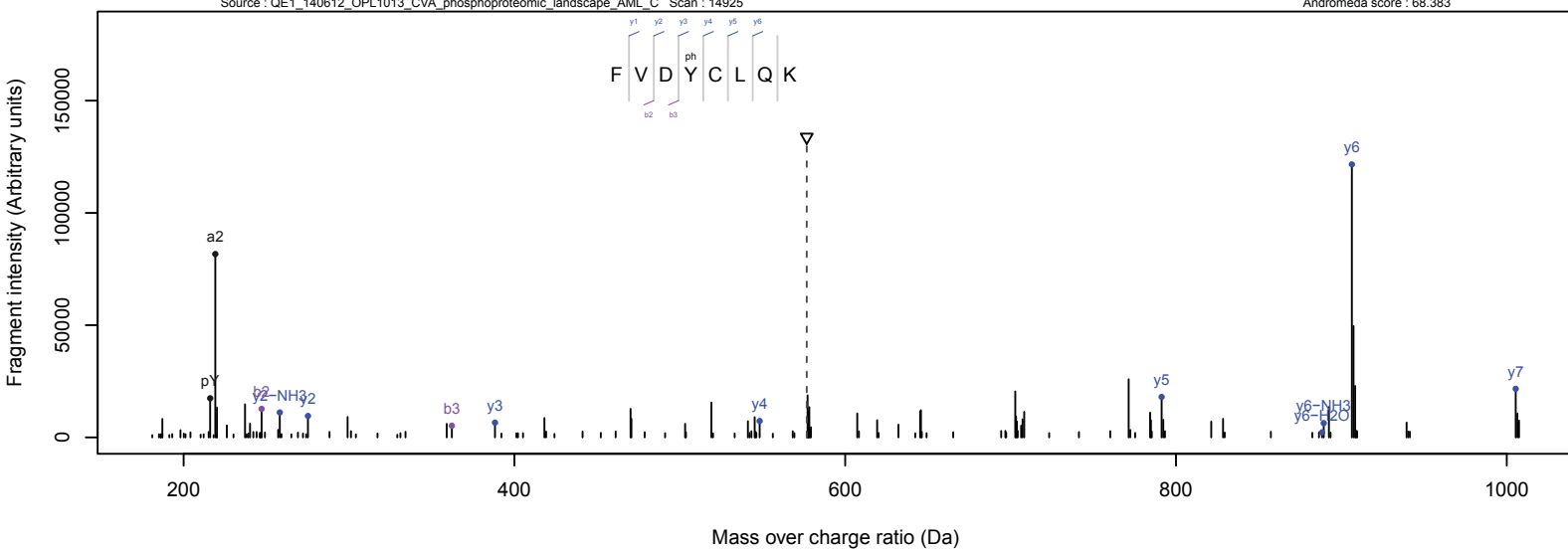
Andromeda score : 68.41



## FVDY(ph)CLQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14925

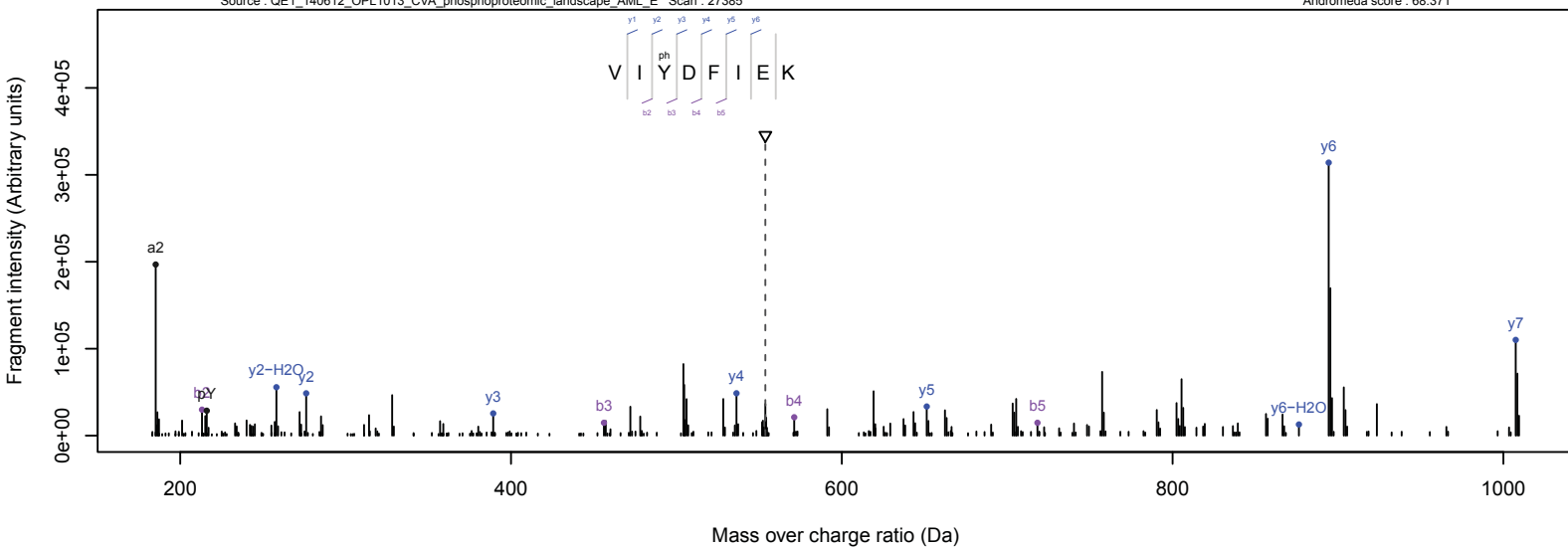
Andromeda score : 68.383



## VIY(ph)DFIEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 27385

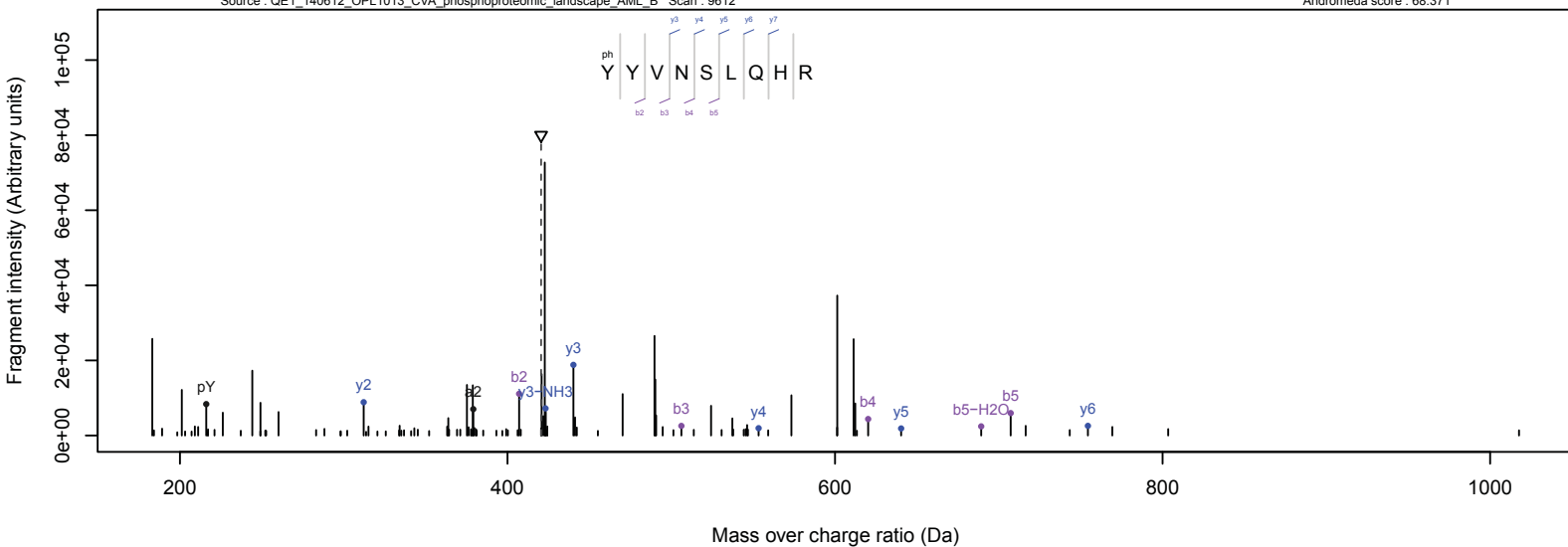
Andromeda score : 68.371



## Y(ph)YVNSLQHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9612

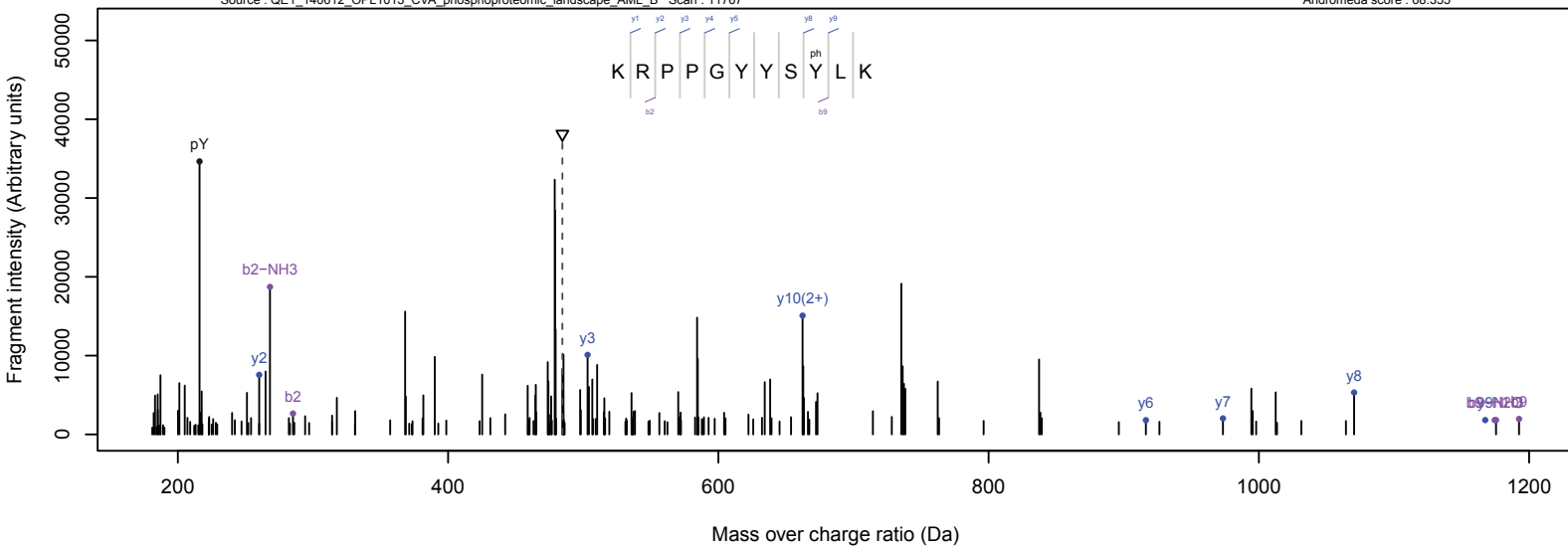
Andromeda score : 68.371

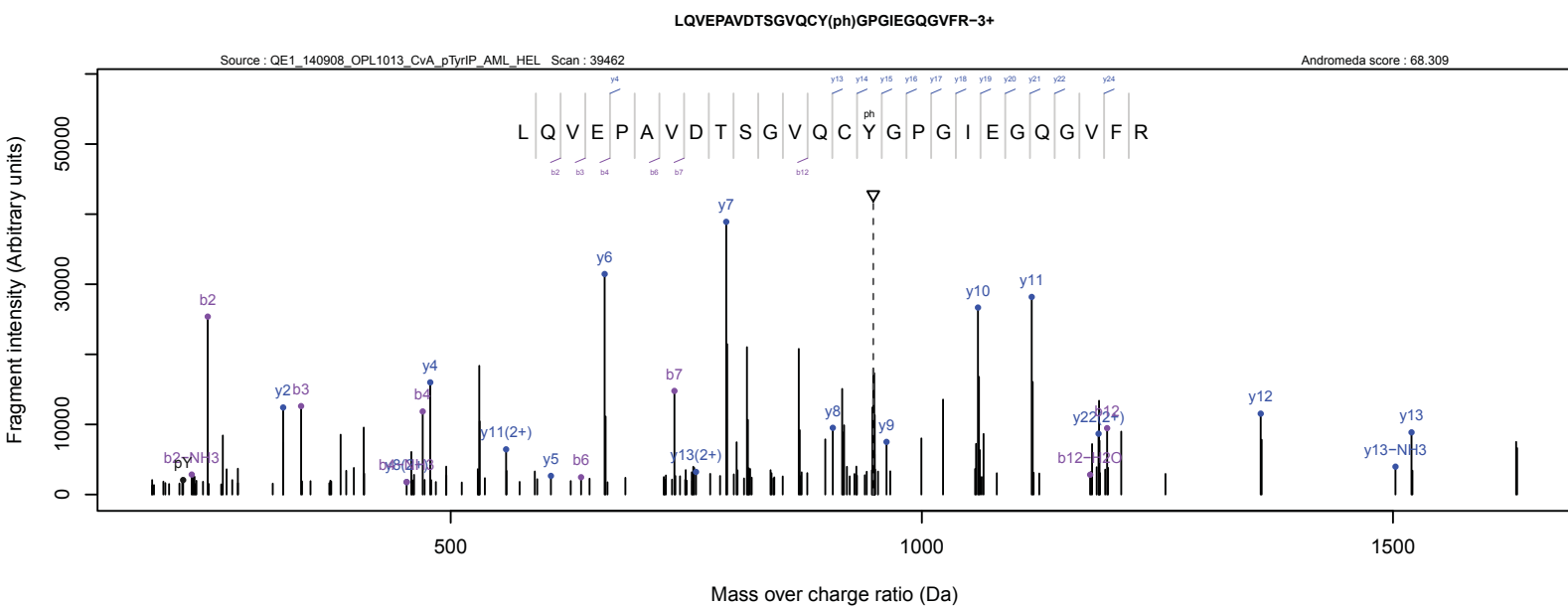
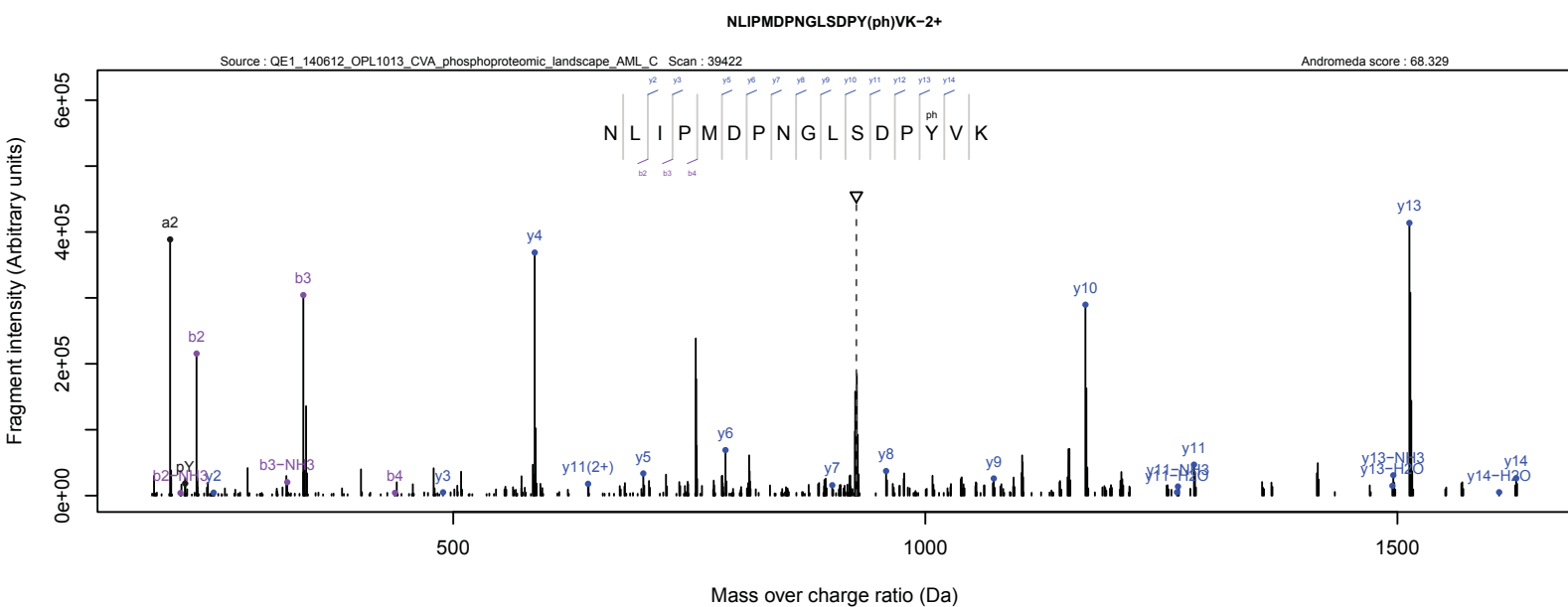
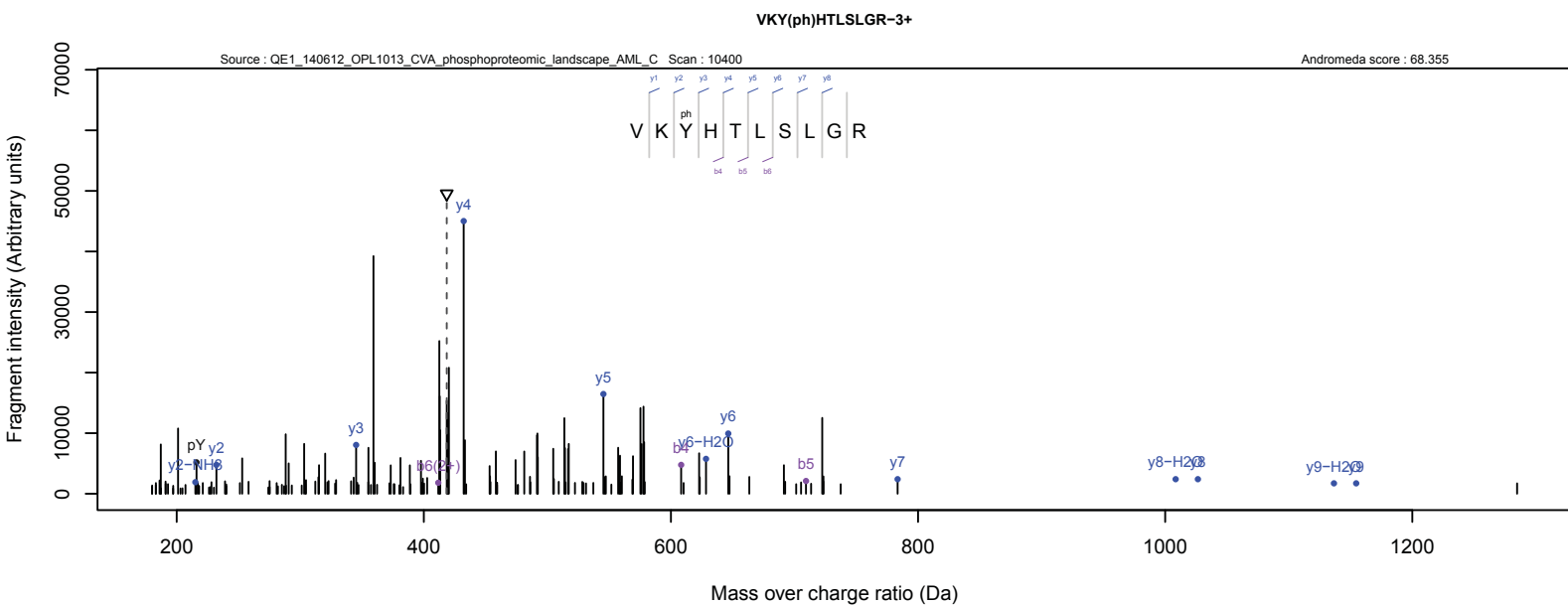


## KRPPGYYSY(ph)LK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11767

Andromeda score : 68.355

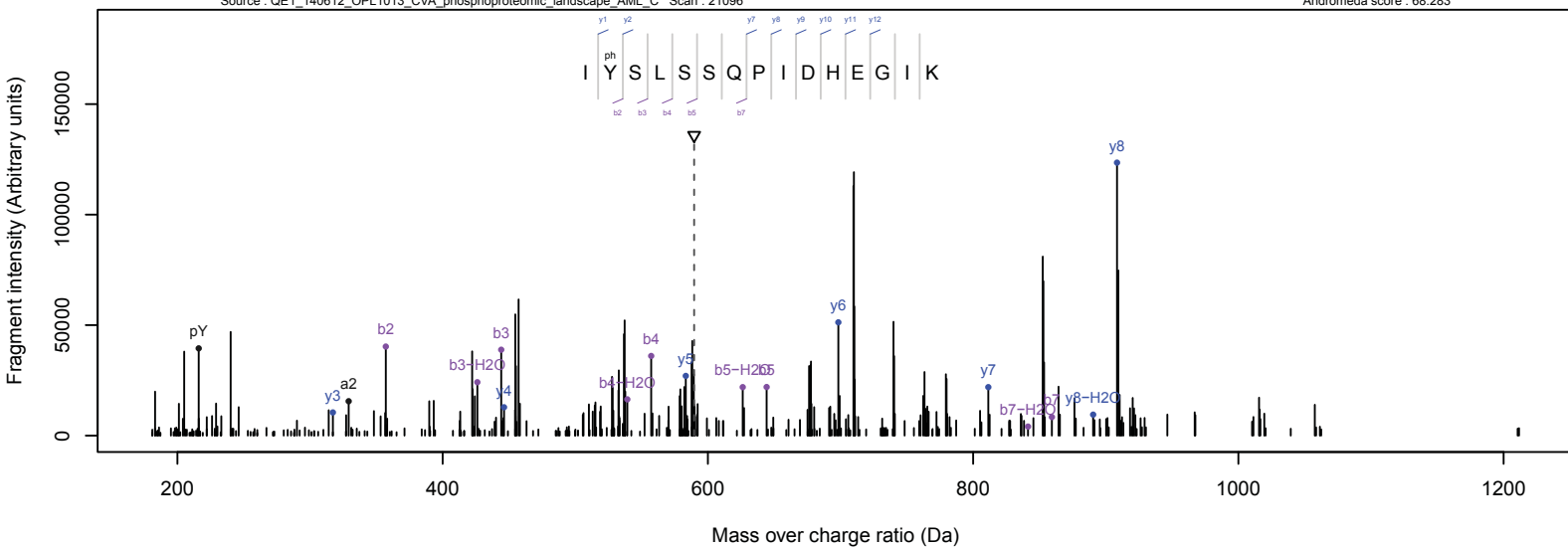




IY(ph)SLSSQPIDHEGIK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21096

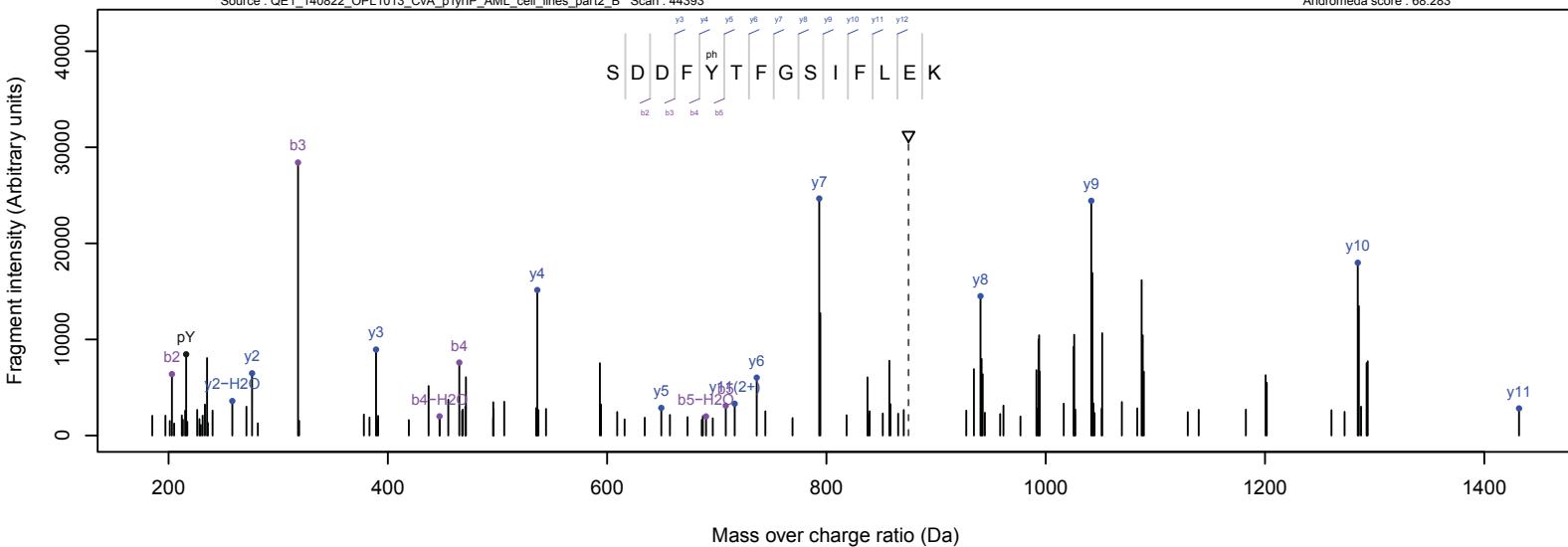
Andromeda score : 68.283



SDDF(ph)TFGSIFLEK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 44393

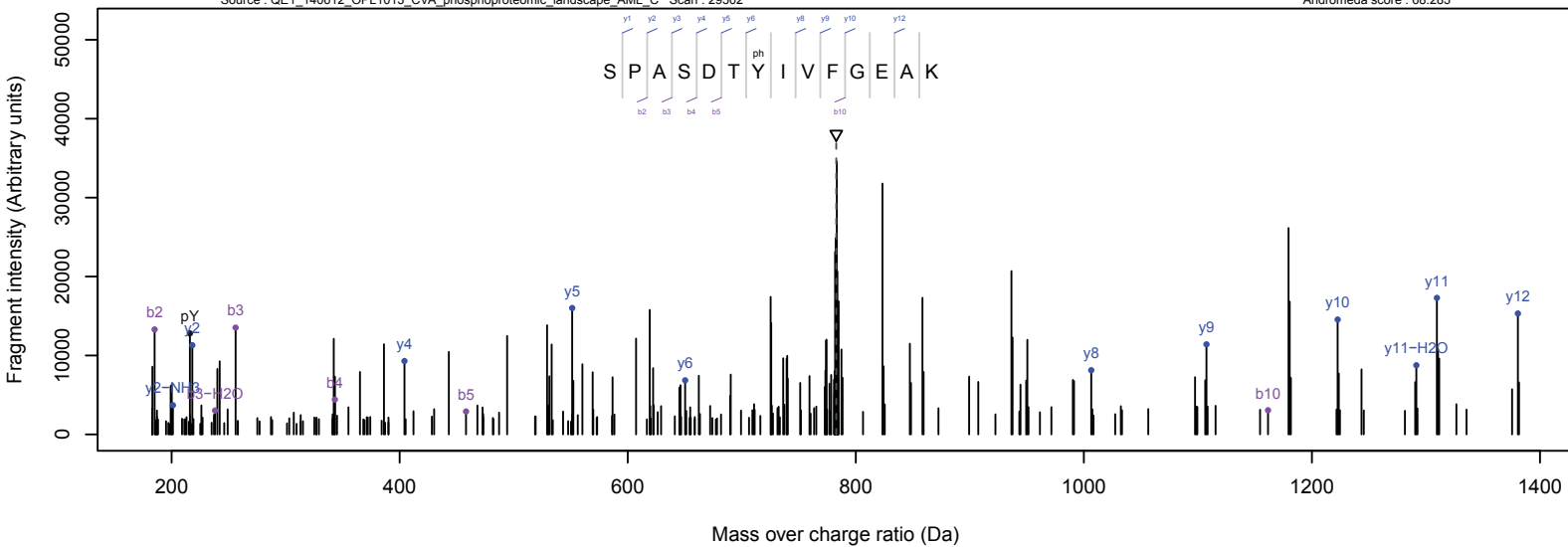
Andromeda score : 68.283



SPASDTY(ph)IVFGEAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29502

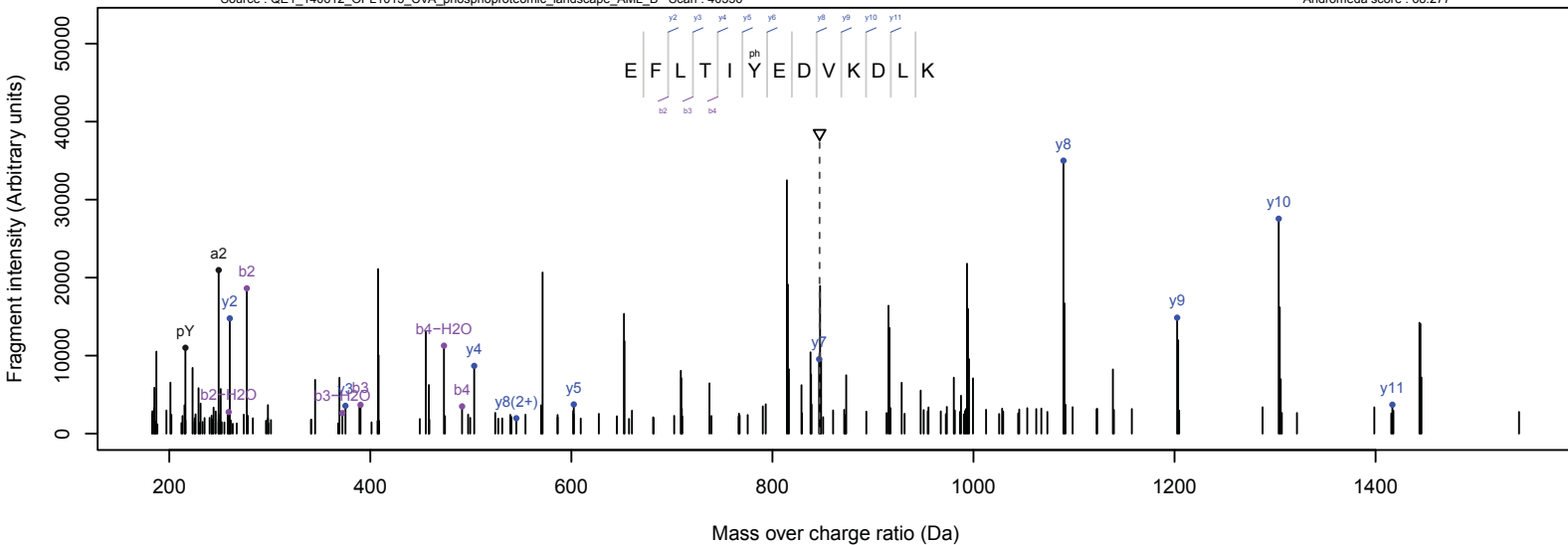
Andromeda score : 68.283



## EFLTIY(ph)EDVKDLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40356

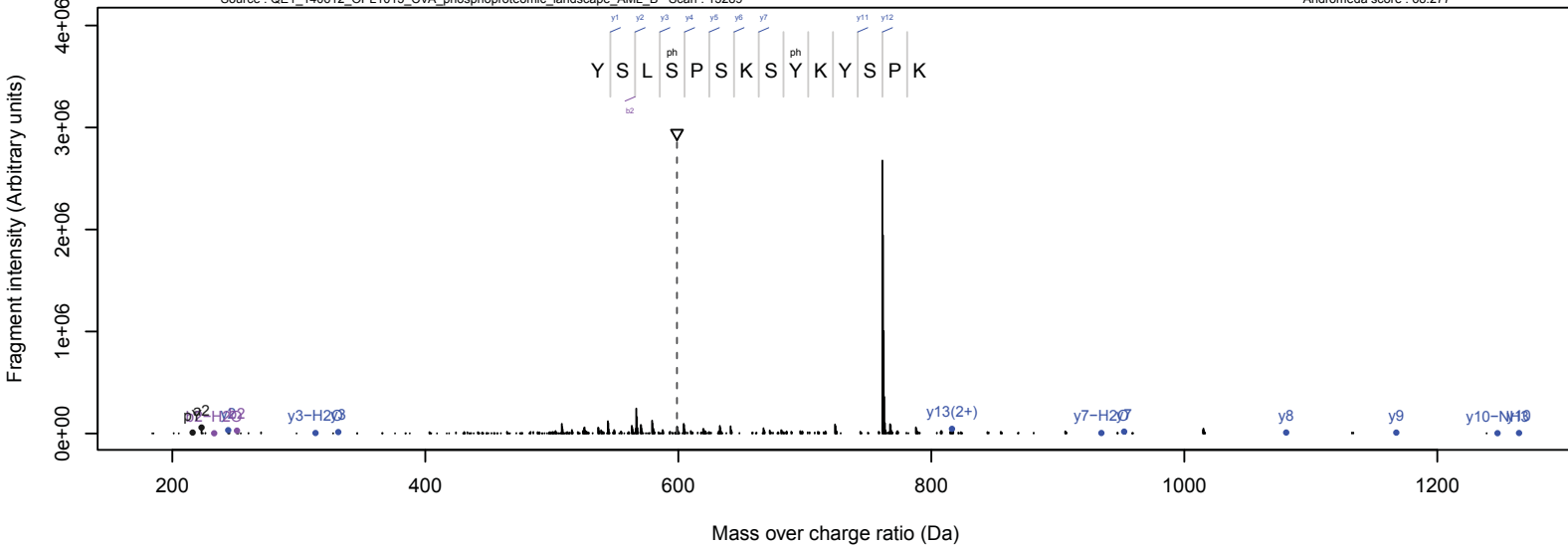
Andromeda score : 68.277



## YSLS(ph)PSKSY(ph)KYSYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13289

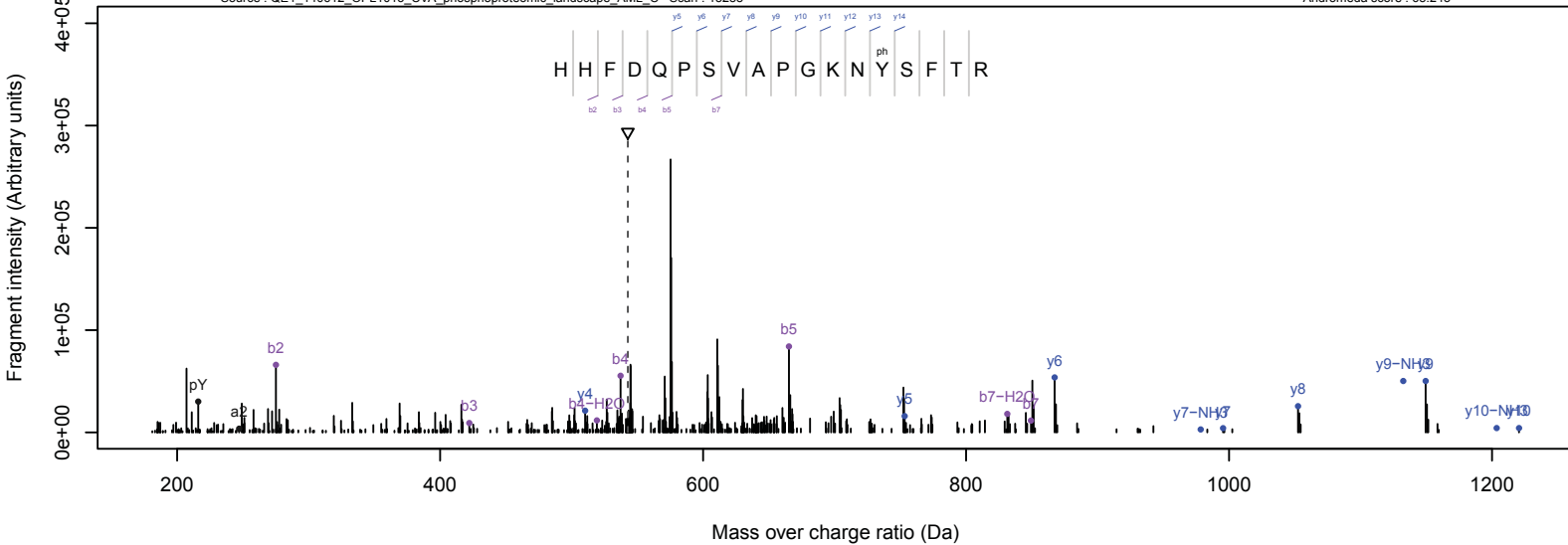
Andromeda score : 68.277



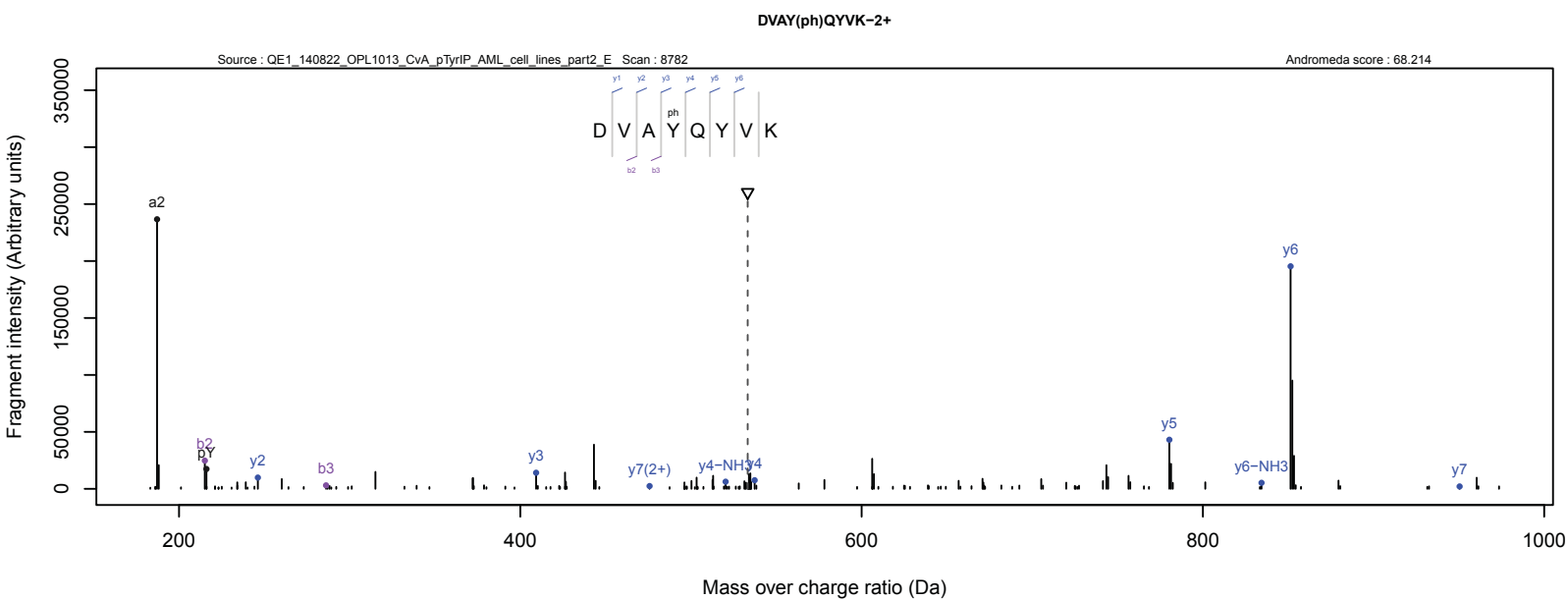
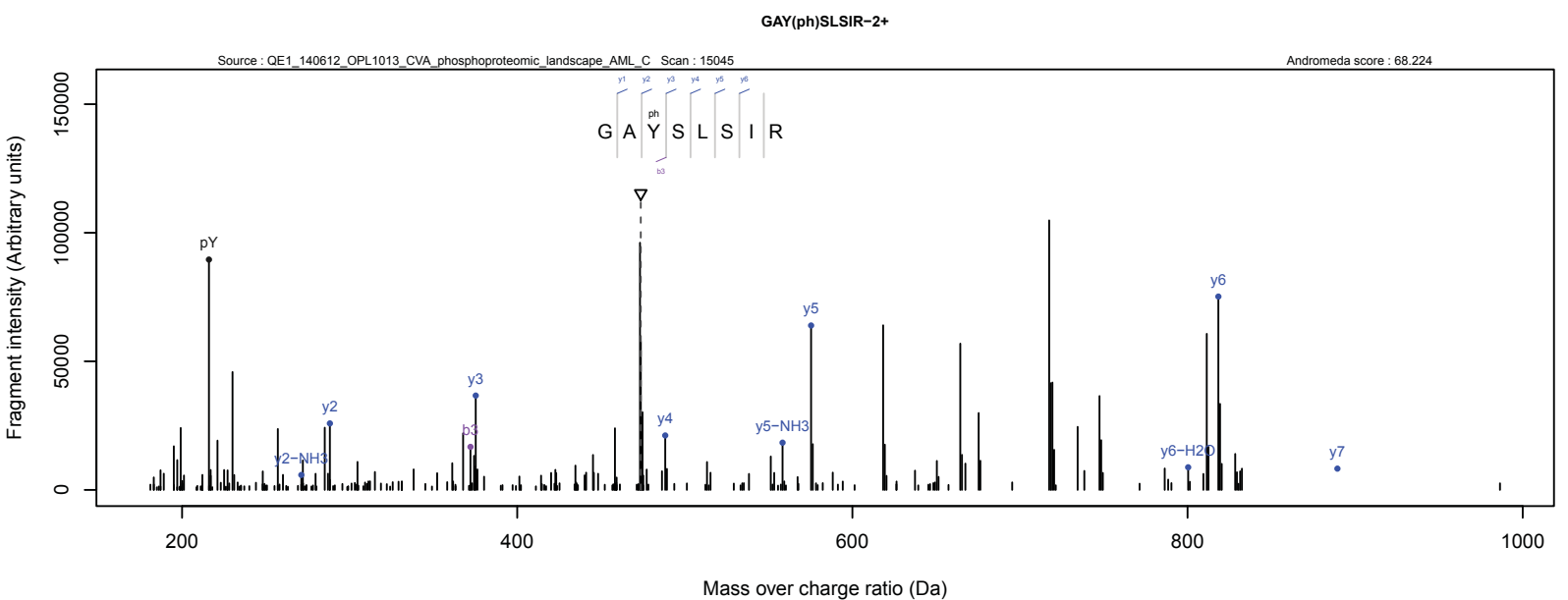
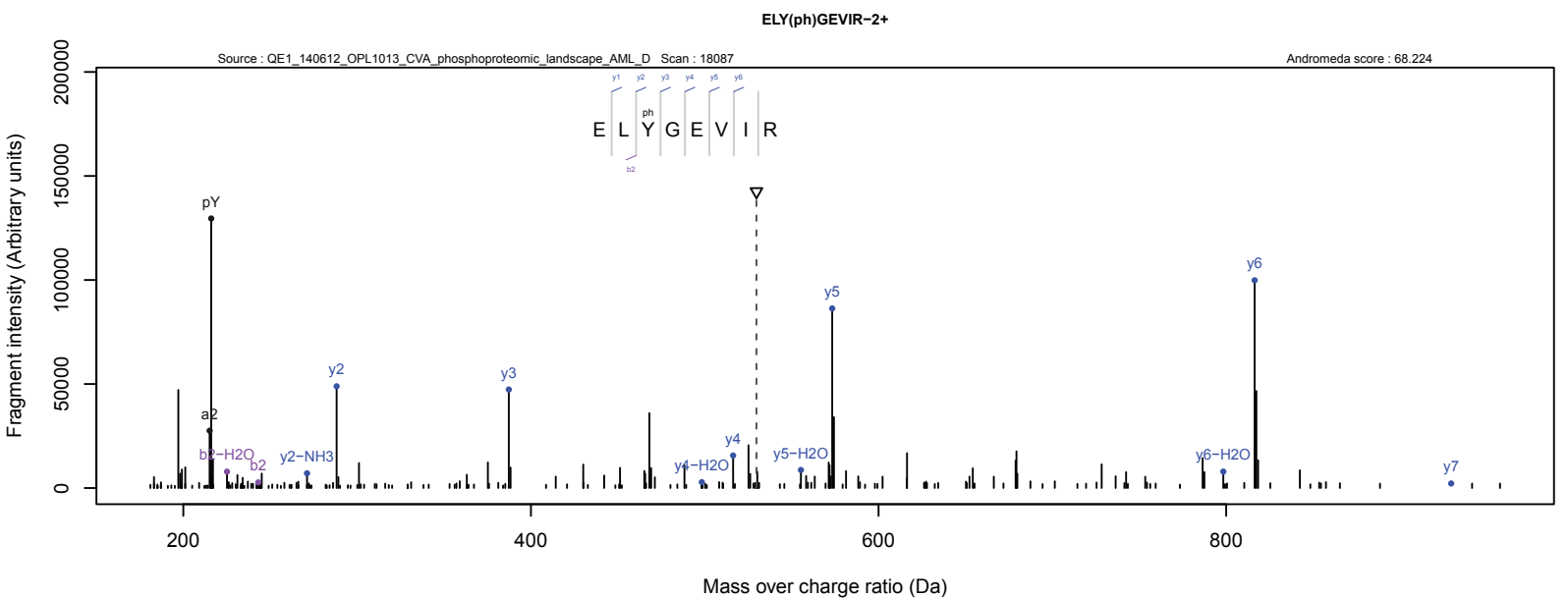
## HHFDQPSVAPGKNYSFTR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15288

Andromeda score : 68.243



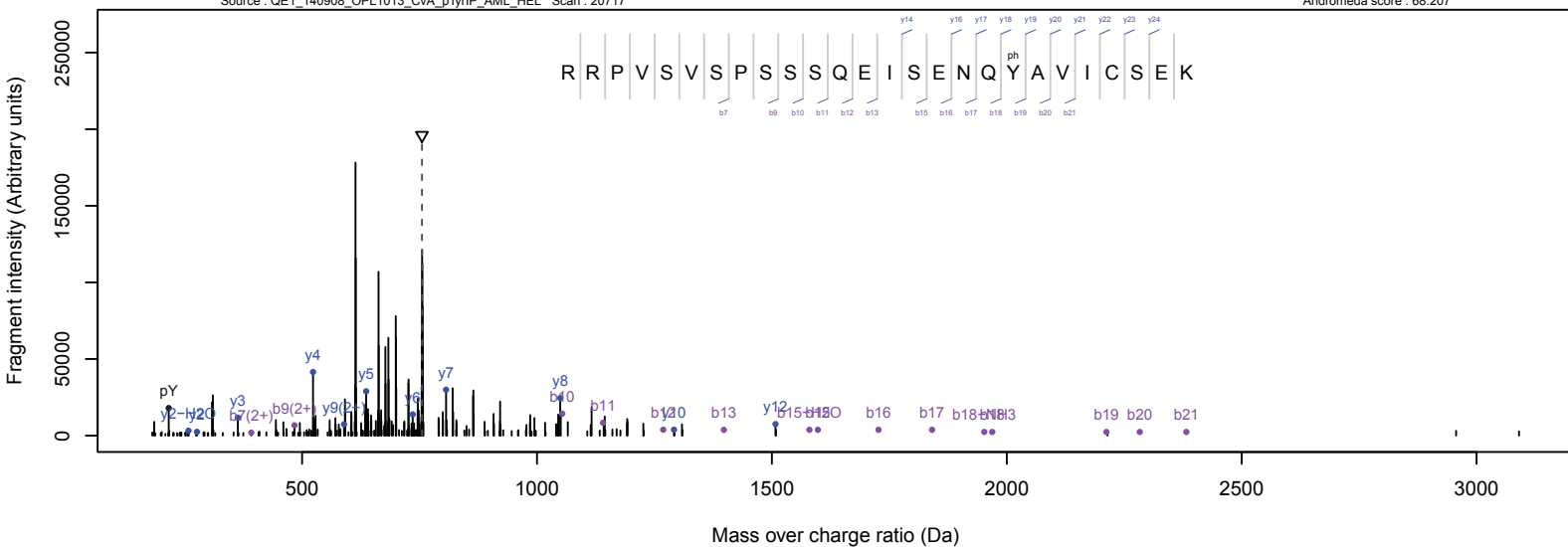




RRPVSVSPSSSQEISENQY(ph)AVICSEK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 20717

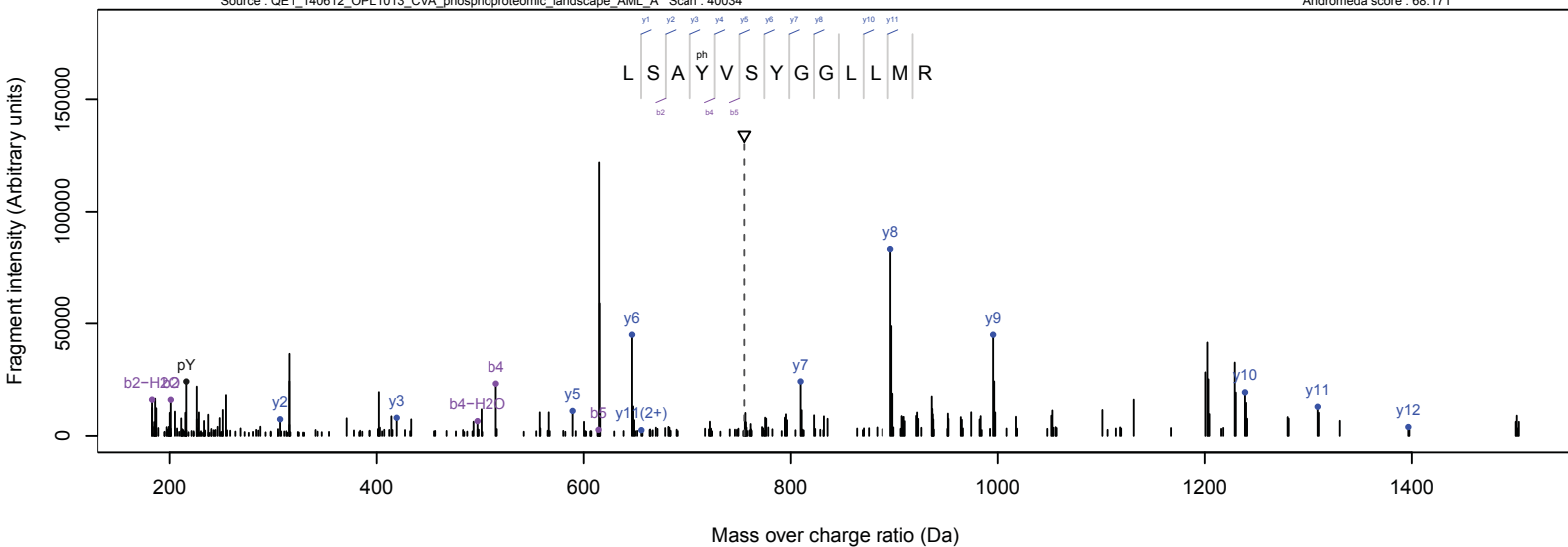
Andromeda score : 68.207



LSAY(ph)VSYGGLLMR-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_A Scan : 40034

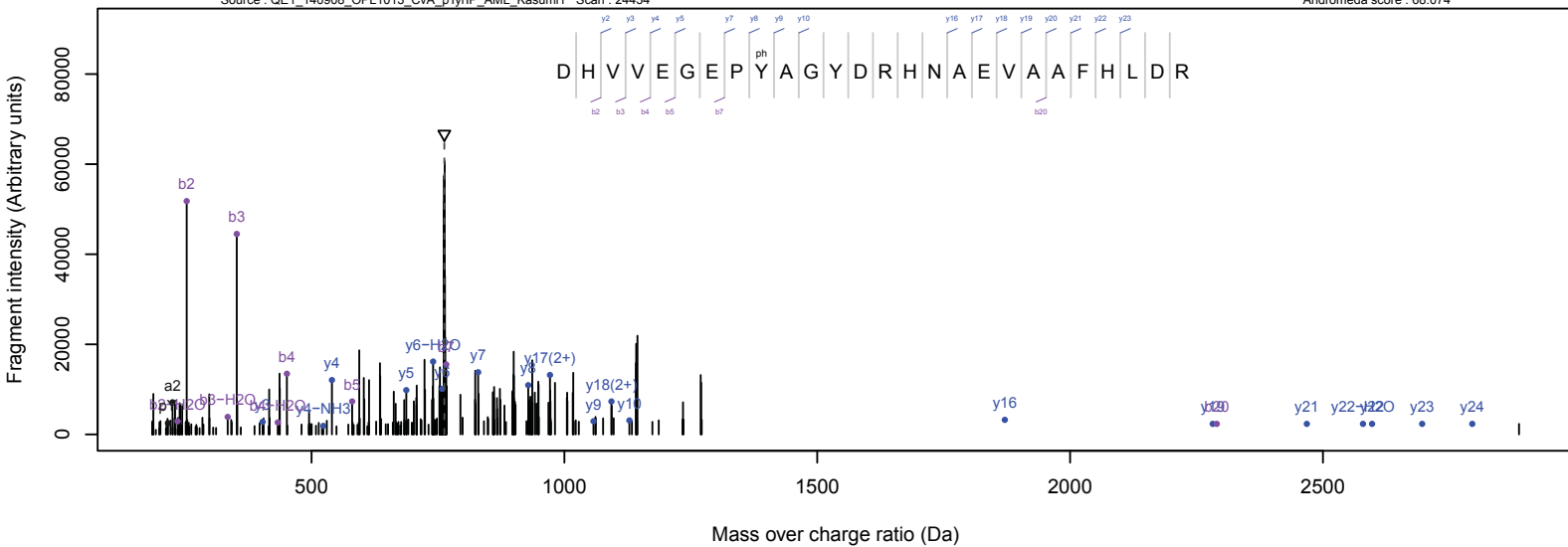
Andromeda score : 68.171



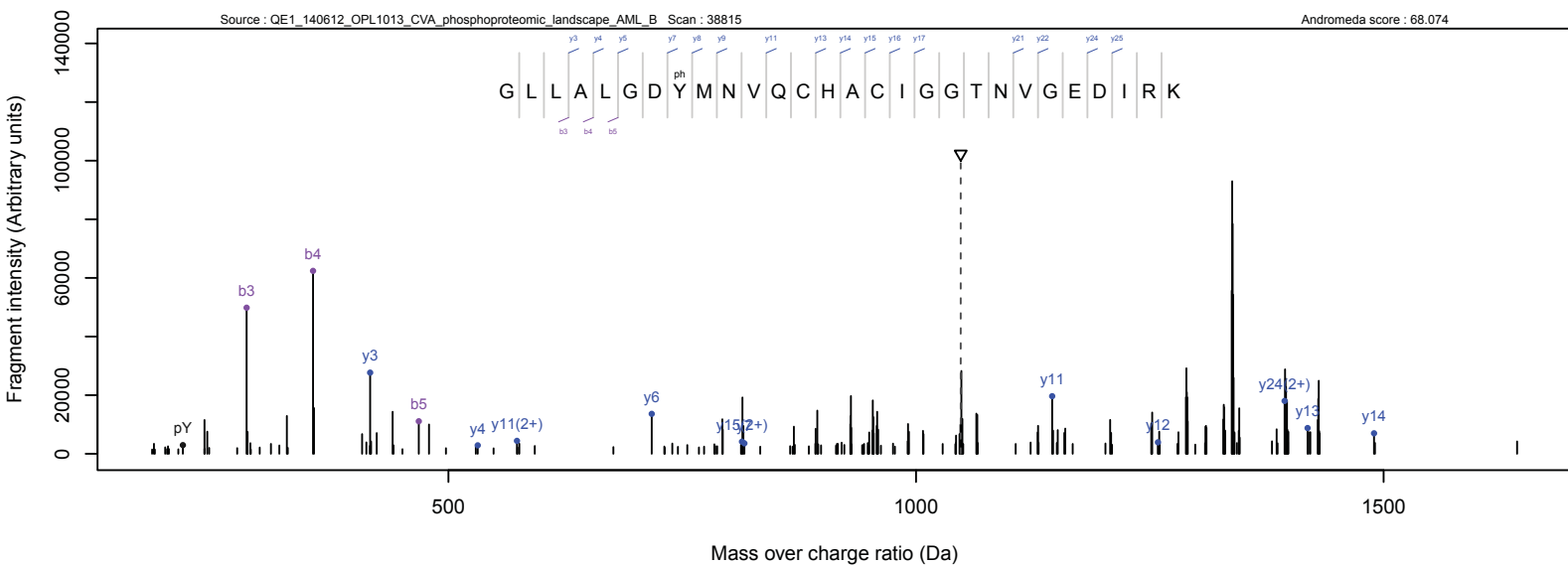
DHVVEGEPYAGYDRHNAEVAAFHLDR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 24434

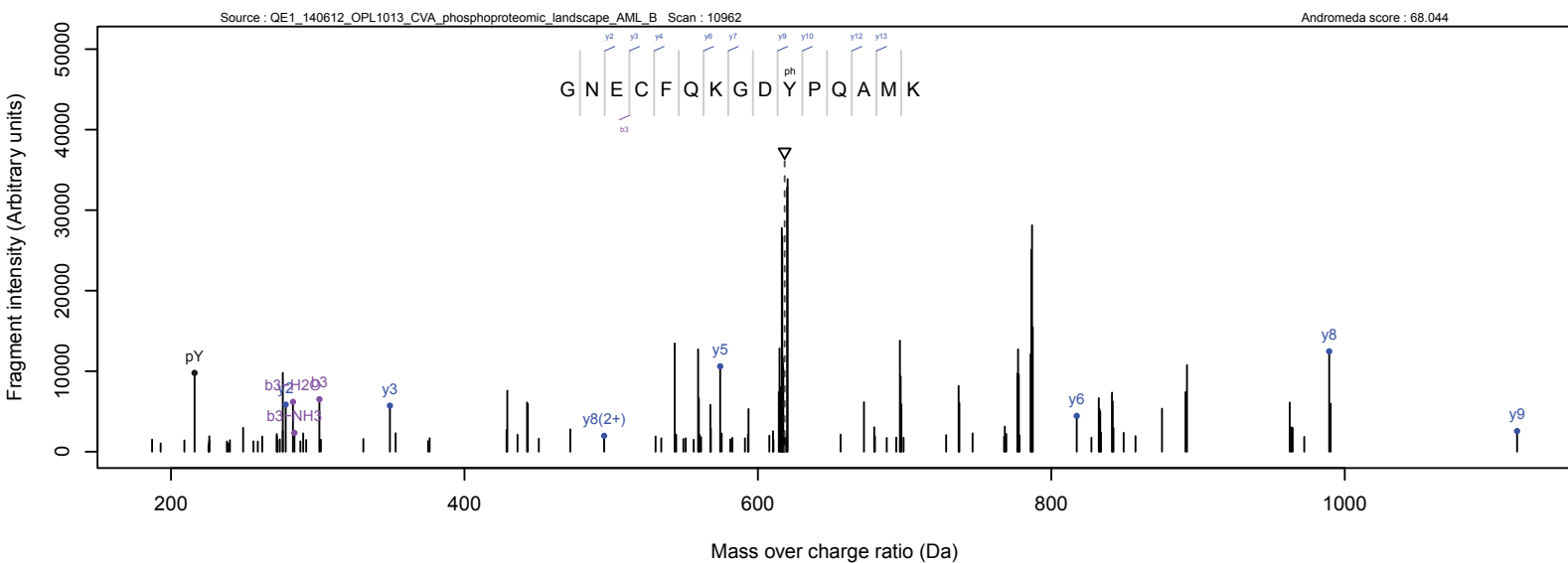
Andromeda score : 68.074



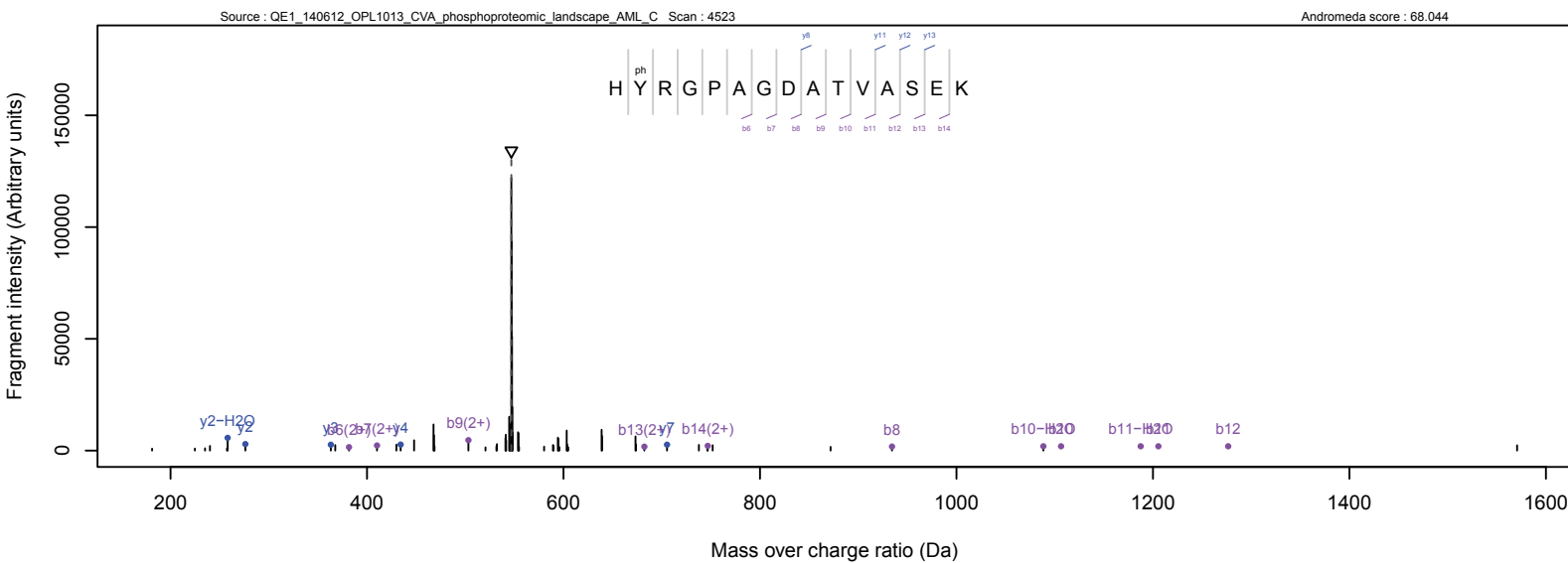
GLLALGDY(ph)MNVQCHACIGGTNVGEDIRK-3+

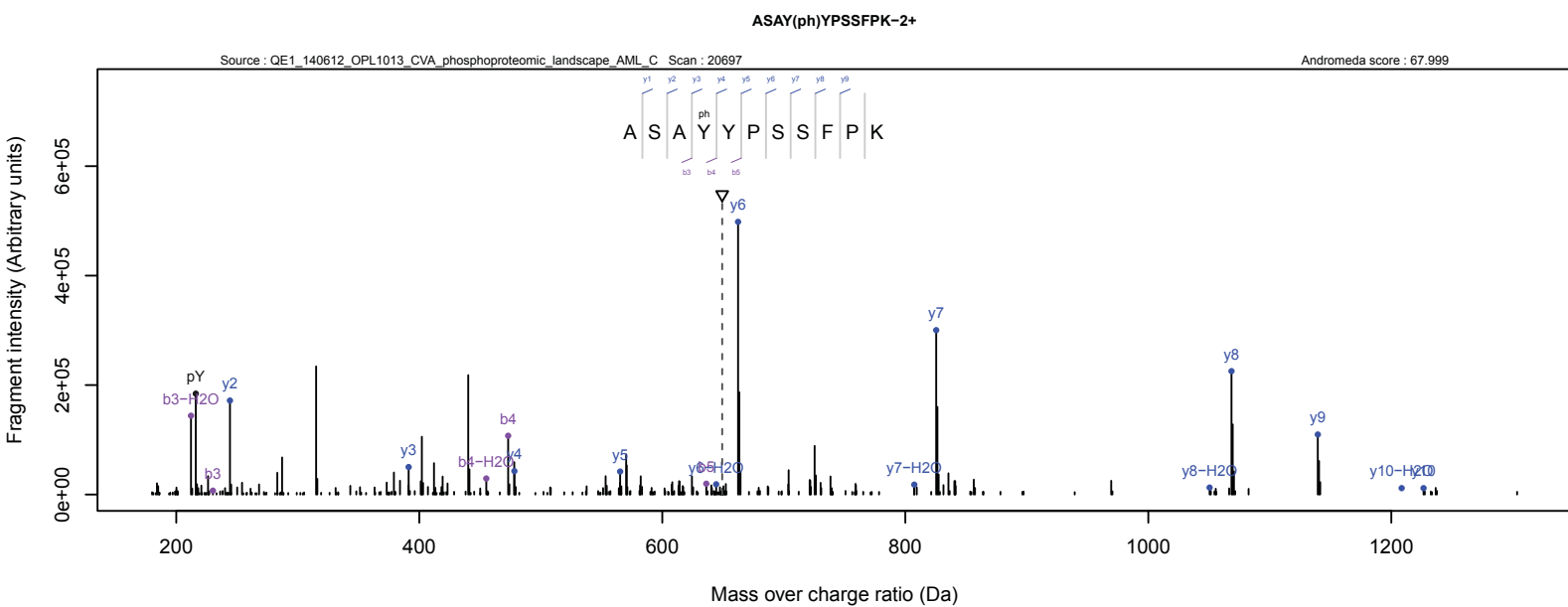
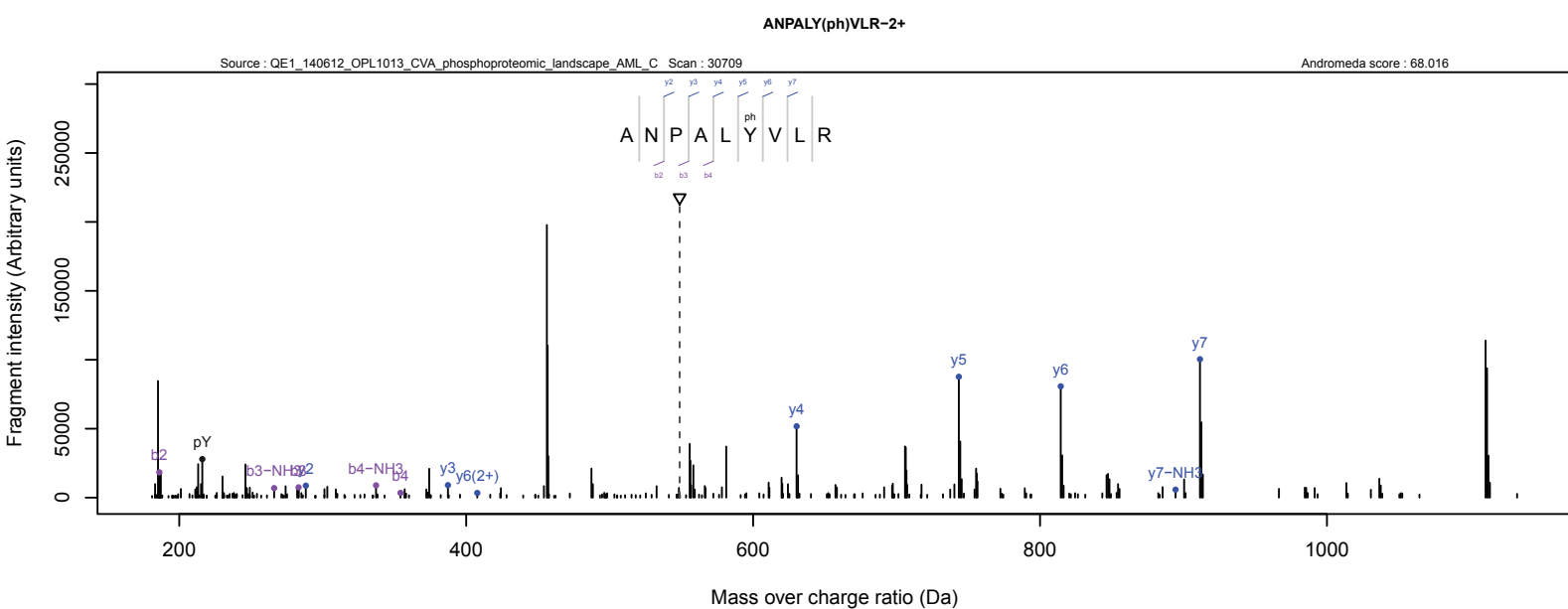
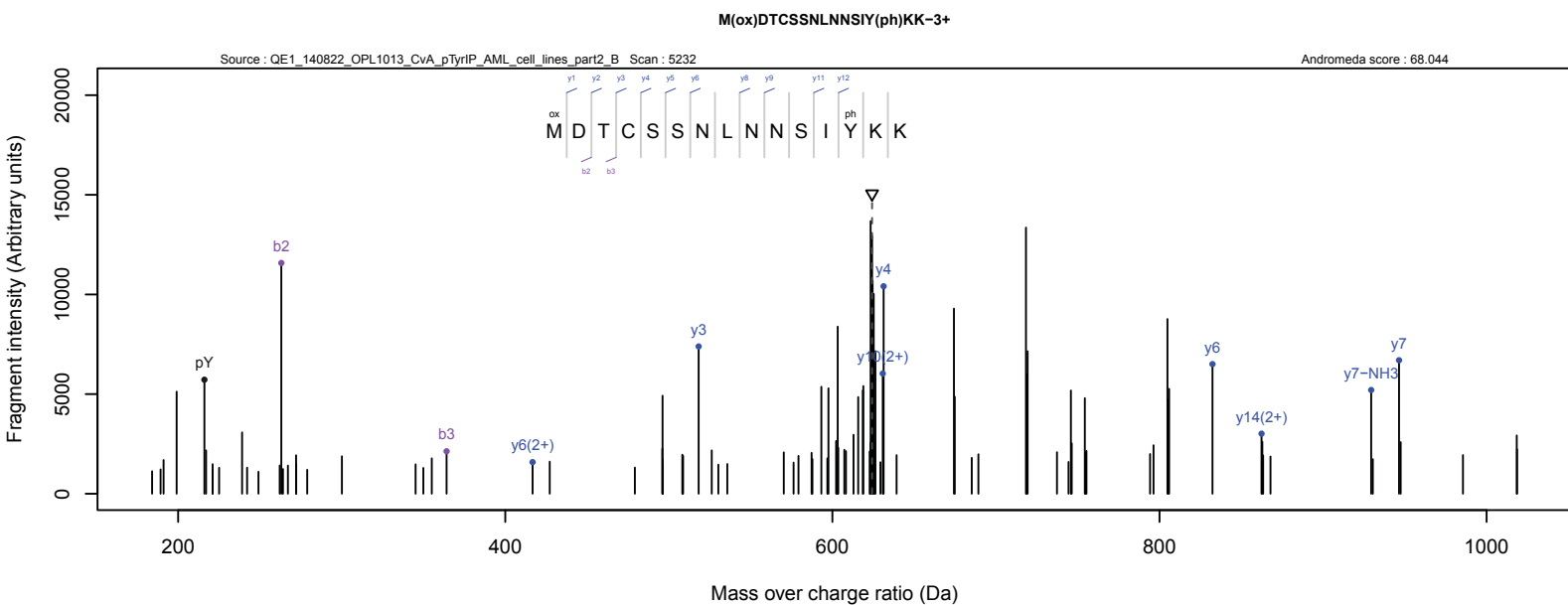


GNECFQKGDY(ph)PQAMK-3+



HY(ph)RGPAGDATVASEK-3+

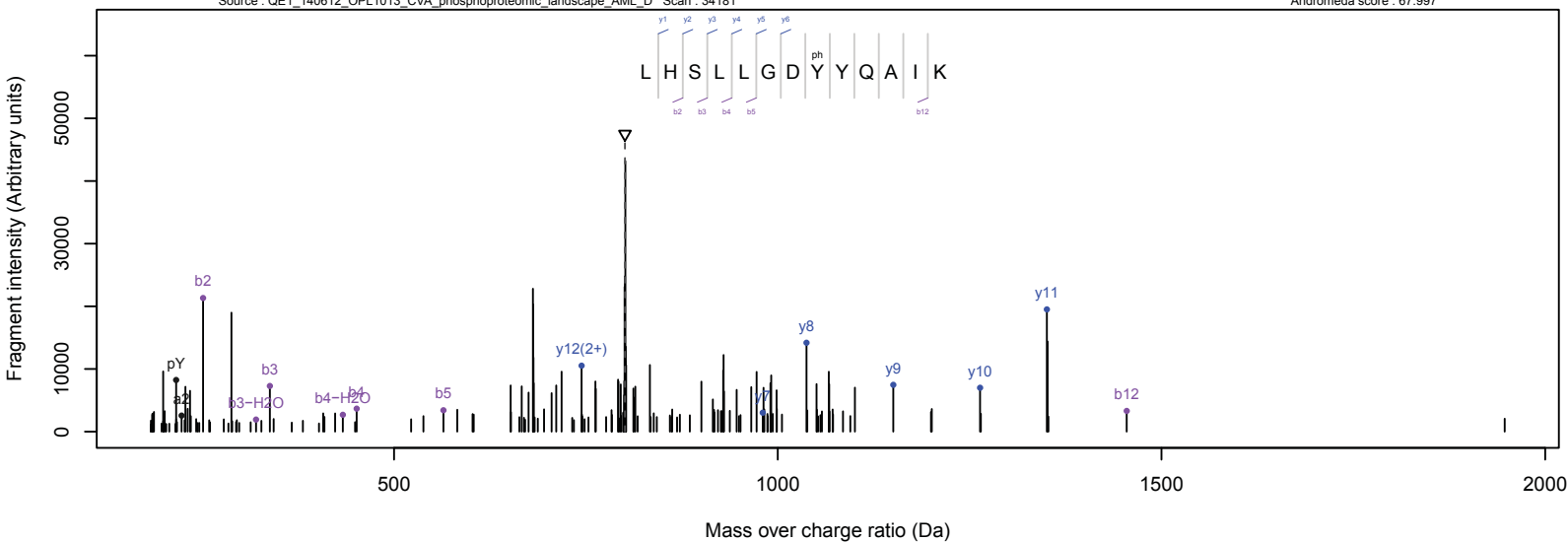




## LHSLLDY(ph)YQAIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 34181

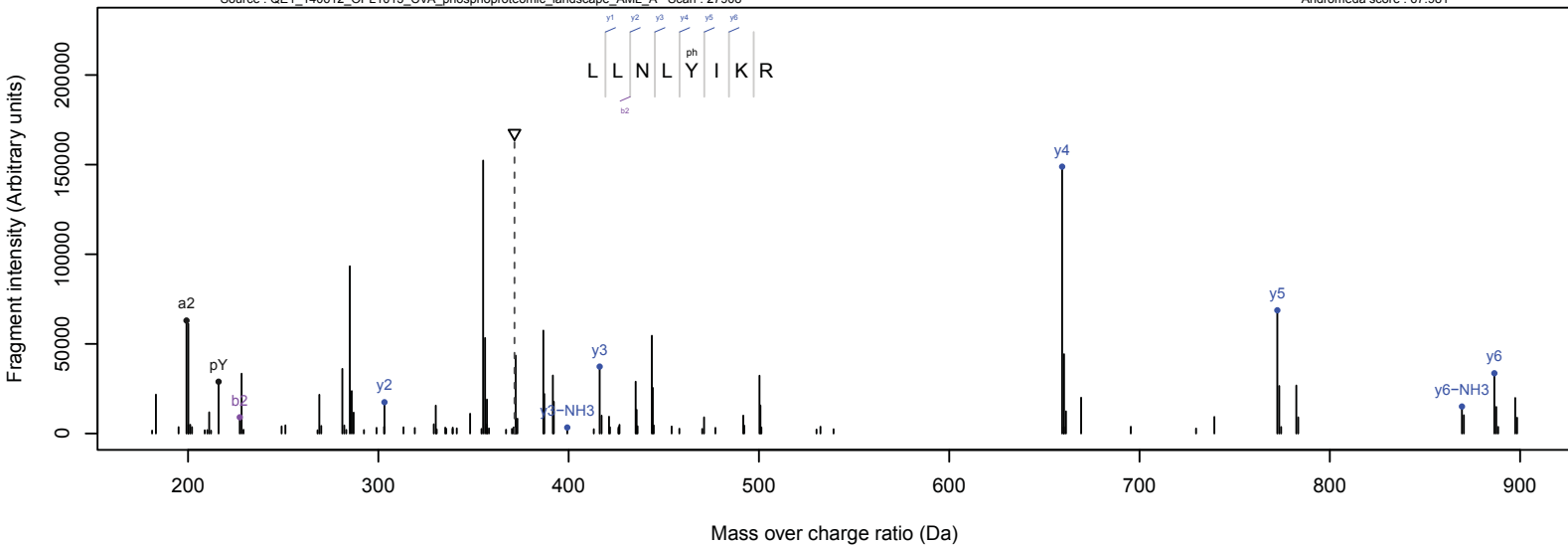
Andromeda score : 67.997



## LLNLY(ph)IKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27908

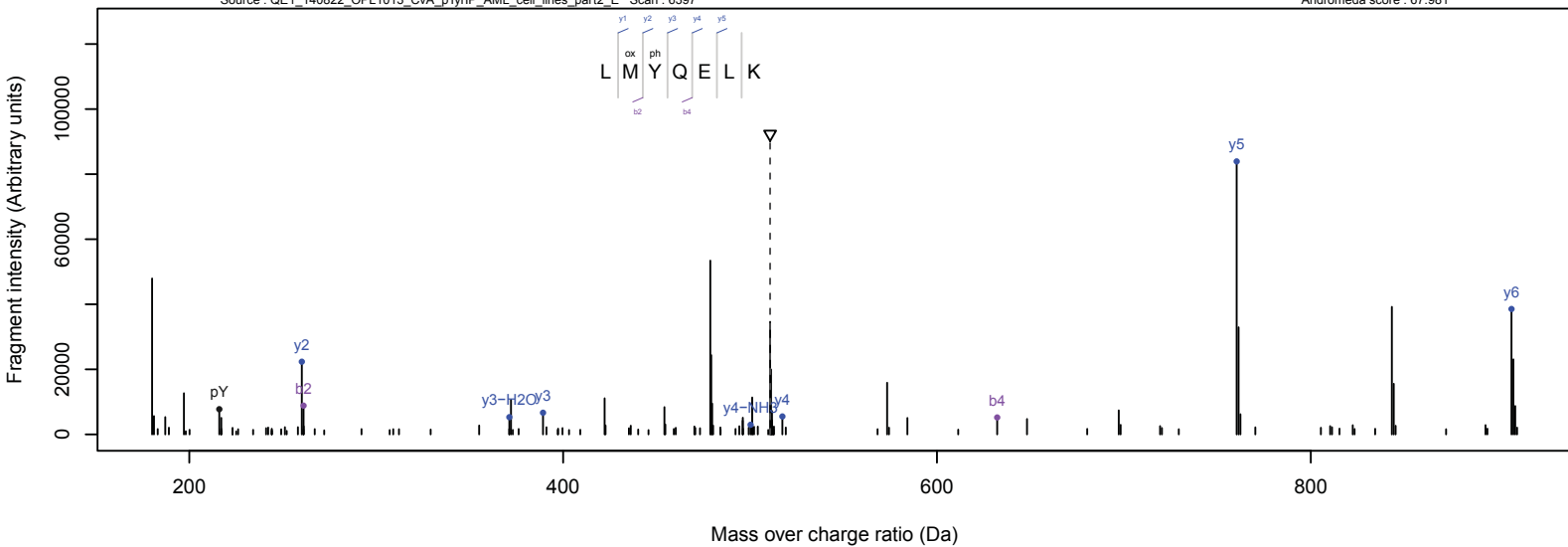
Andromeda score : 67.981



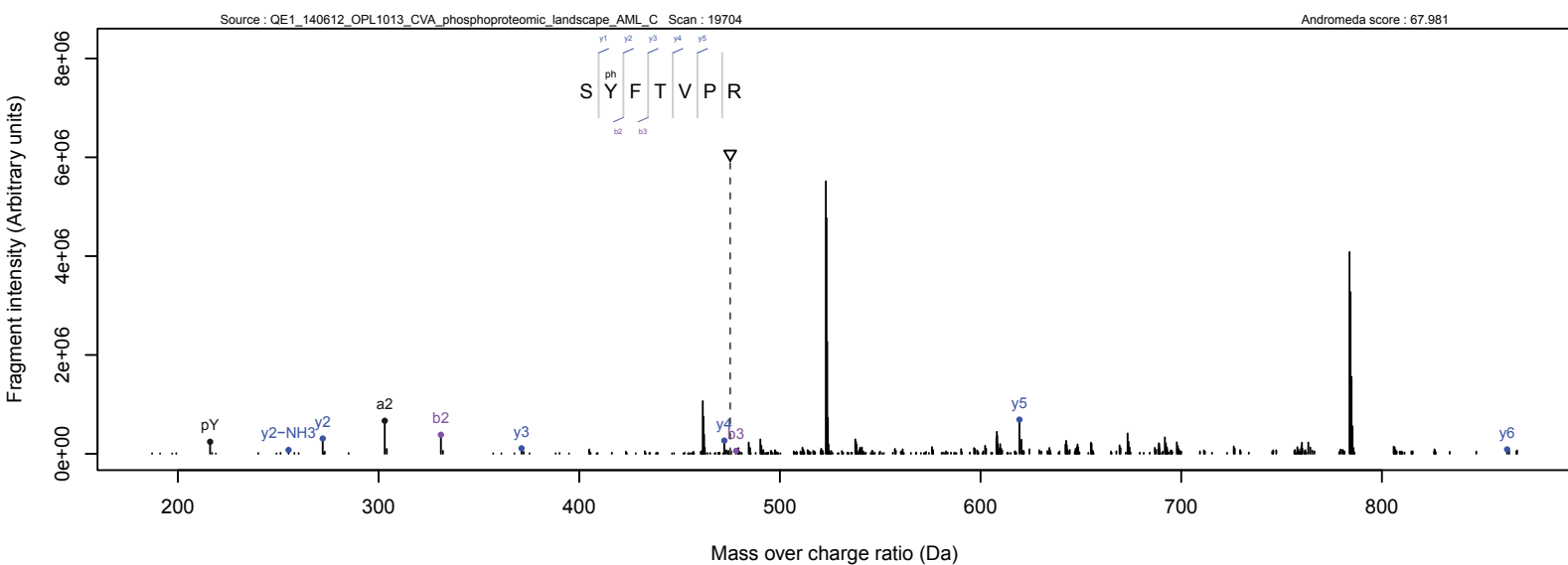
## LM(ox)Y(ph)QELK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 6397

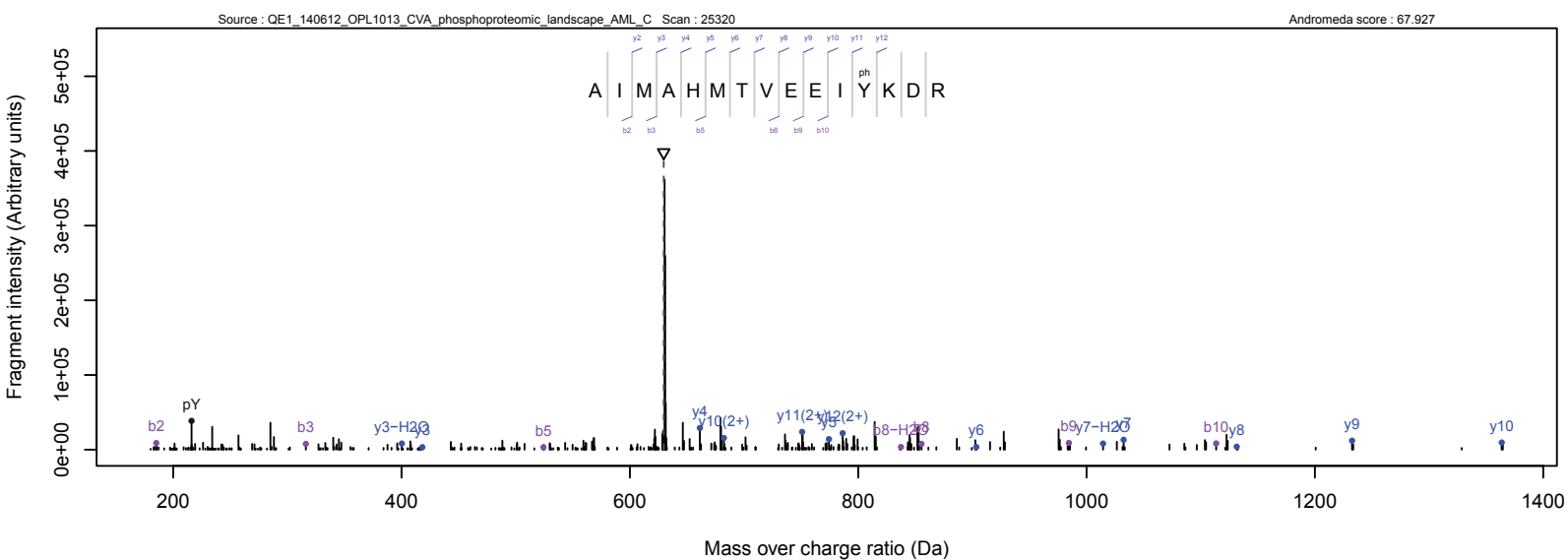
Andromeda score : 67.981



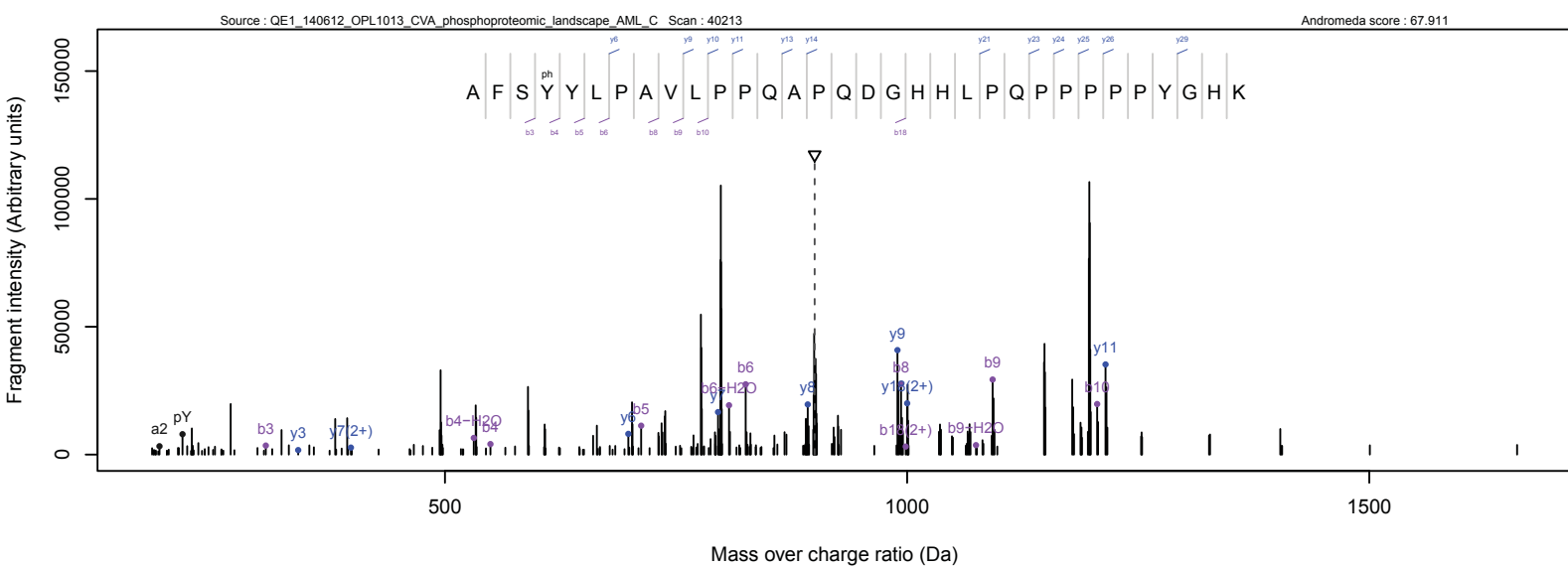
## SY(ph)FTVPR-2+



## AIMAHMTVEEIY(ph)KDR-3+



## AFSY(ph)YLPVLPQAPQDGHLLPQPPPPYGHK-4+

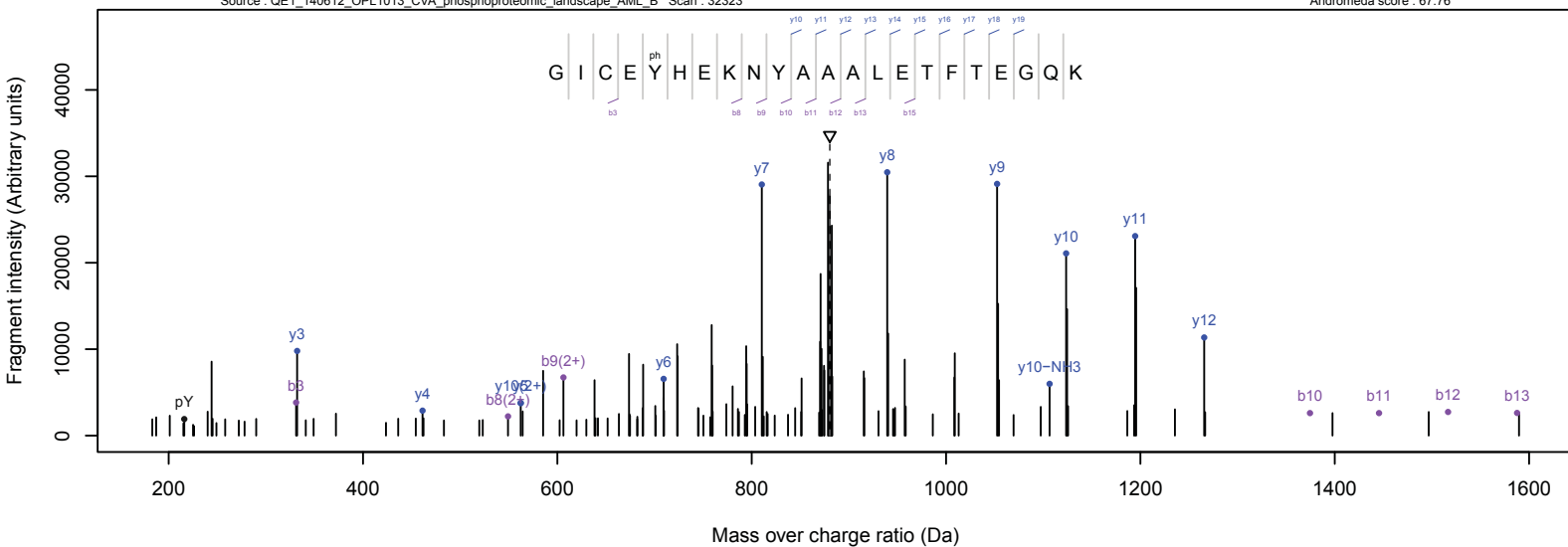




## GICEY(ph)HEKNYAAALETFTEGQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32323

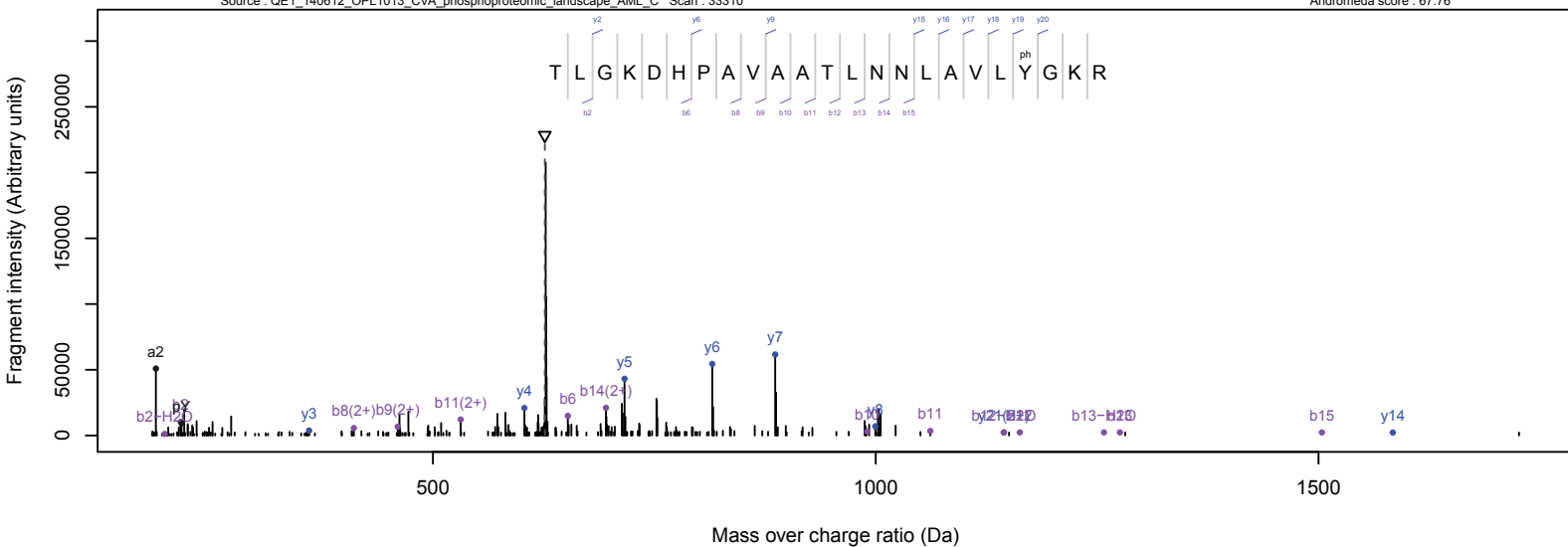
Andromeda score : 67.76



## TLGKDHPAVALNLAVALY(ph)GKR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33310

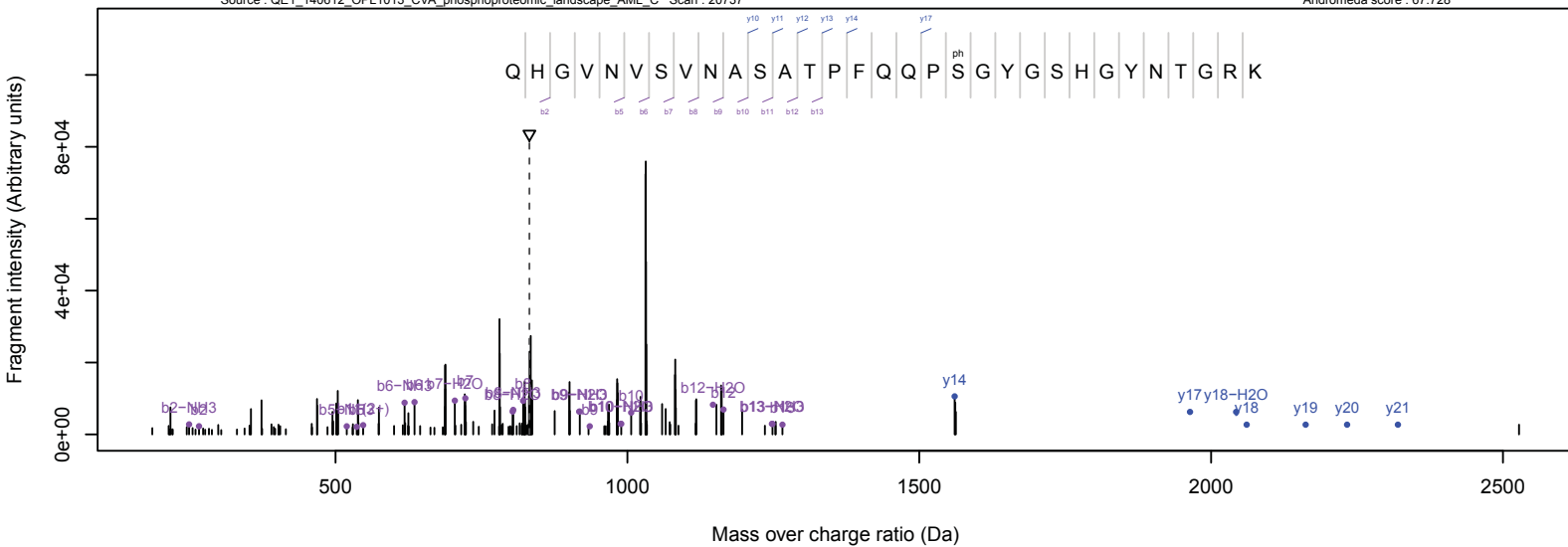
Andromeda score : 67.76



## QHGVNVSVAATPFQQPS(ph)YGSHGYNTGRK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20737

Andromeda score : 67.728

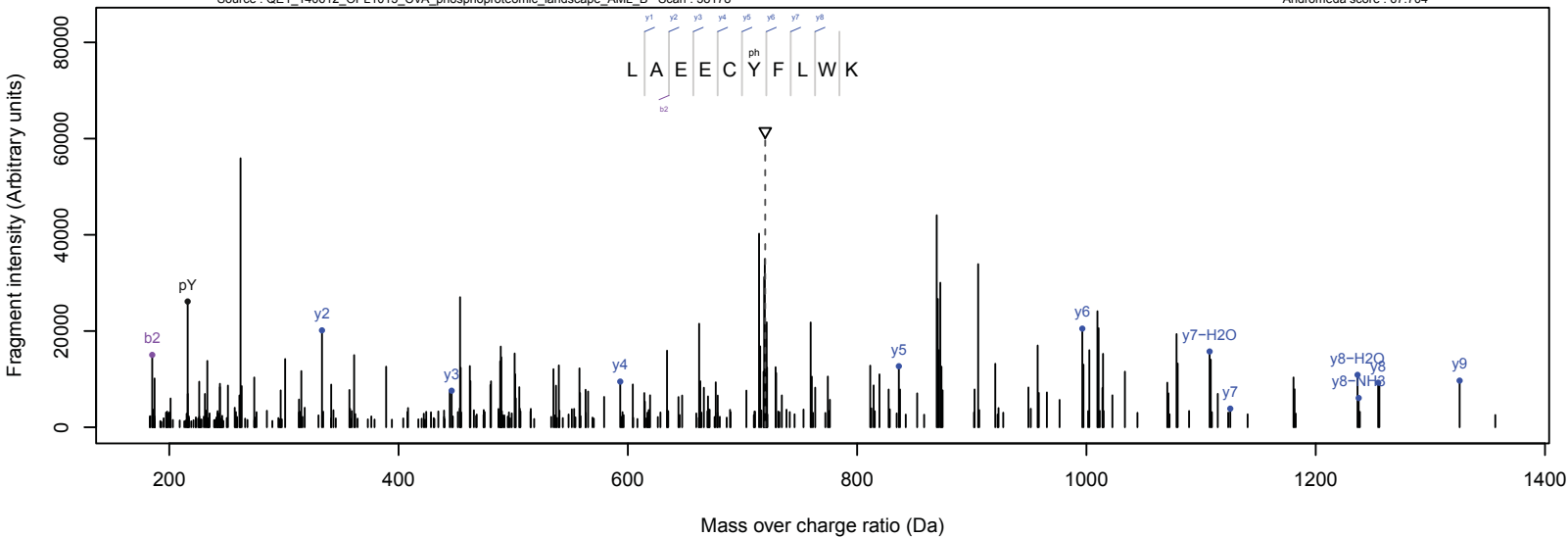




## LAEECY(ph)FLWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 38178

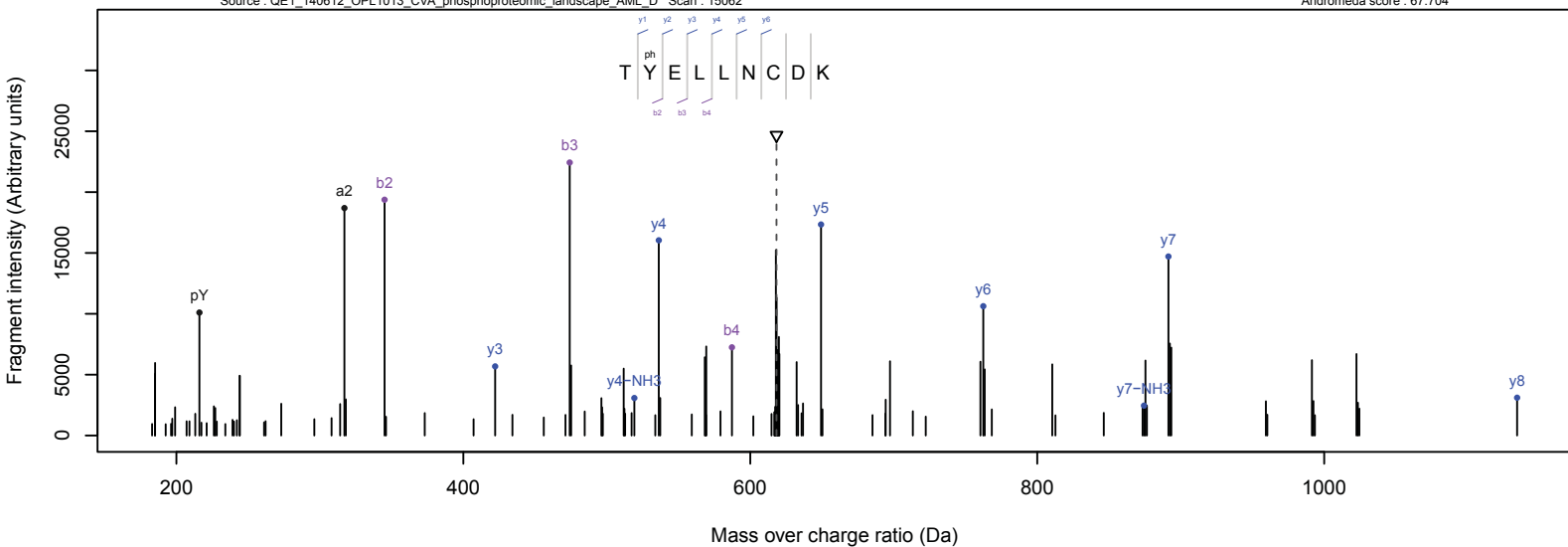
Andromeda score : 67.704



## TY(ph)ELLNCDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 15062

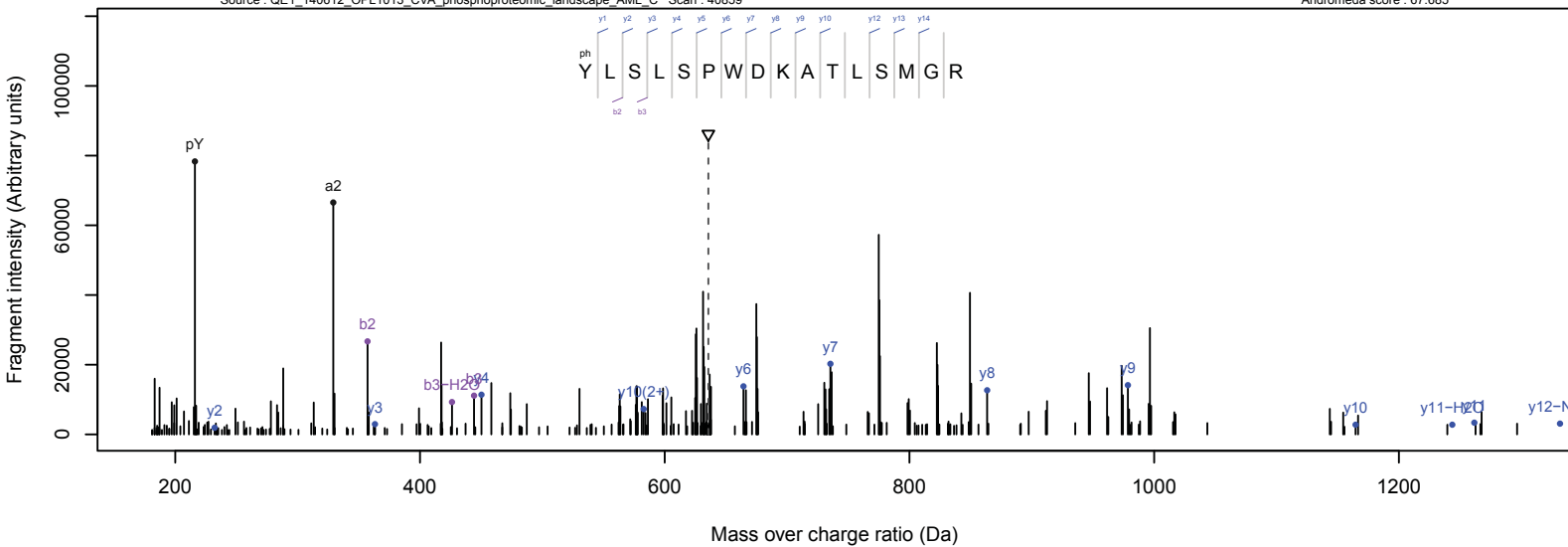
Andromeda score : 67.704



## Y(ph)LSLSPWDKATLSMGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40859

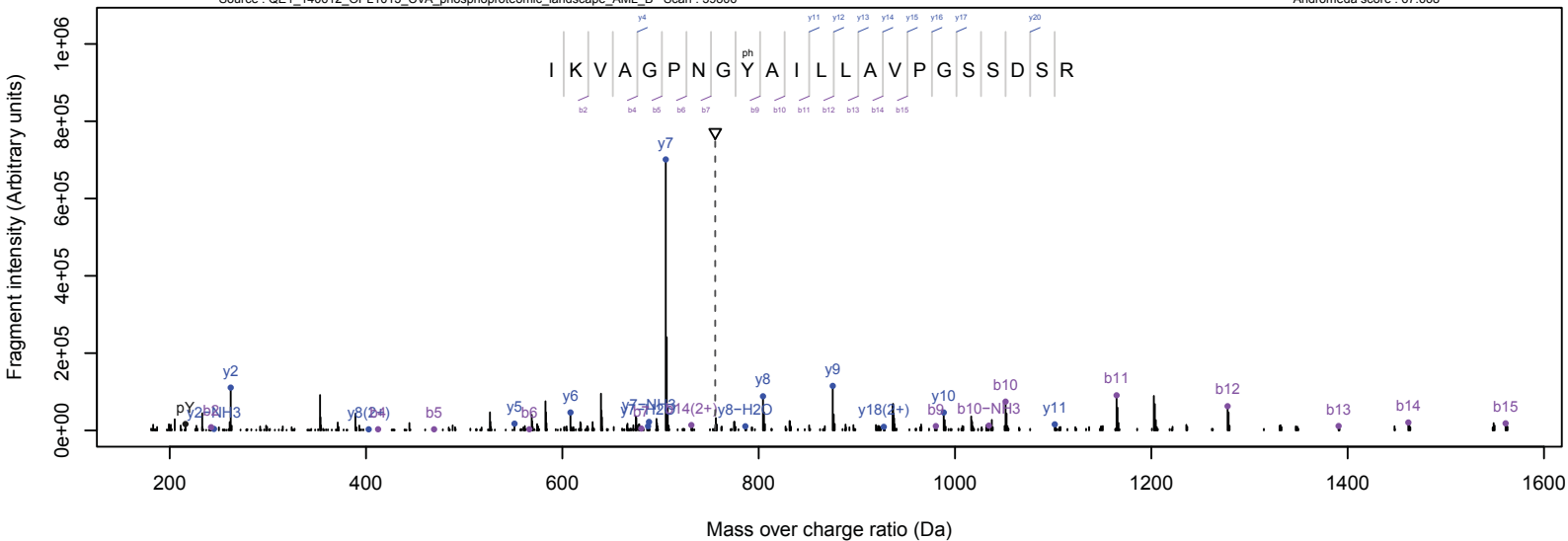
Andromeda score : 67.685



## IKVAGPNGY(ph)AILLAVPGSSDSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39806

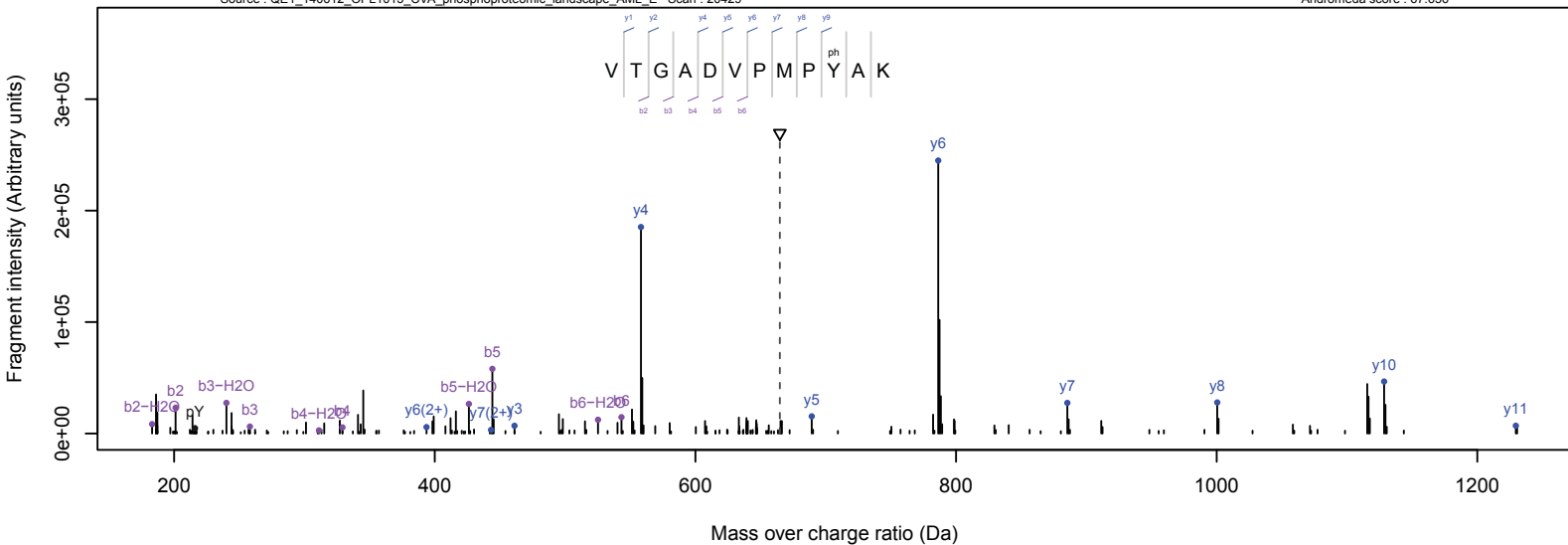
Andromeda score : 67.668



## VTGADVMPY(ph)AK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 20429

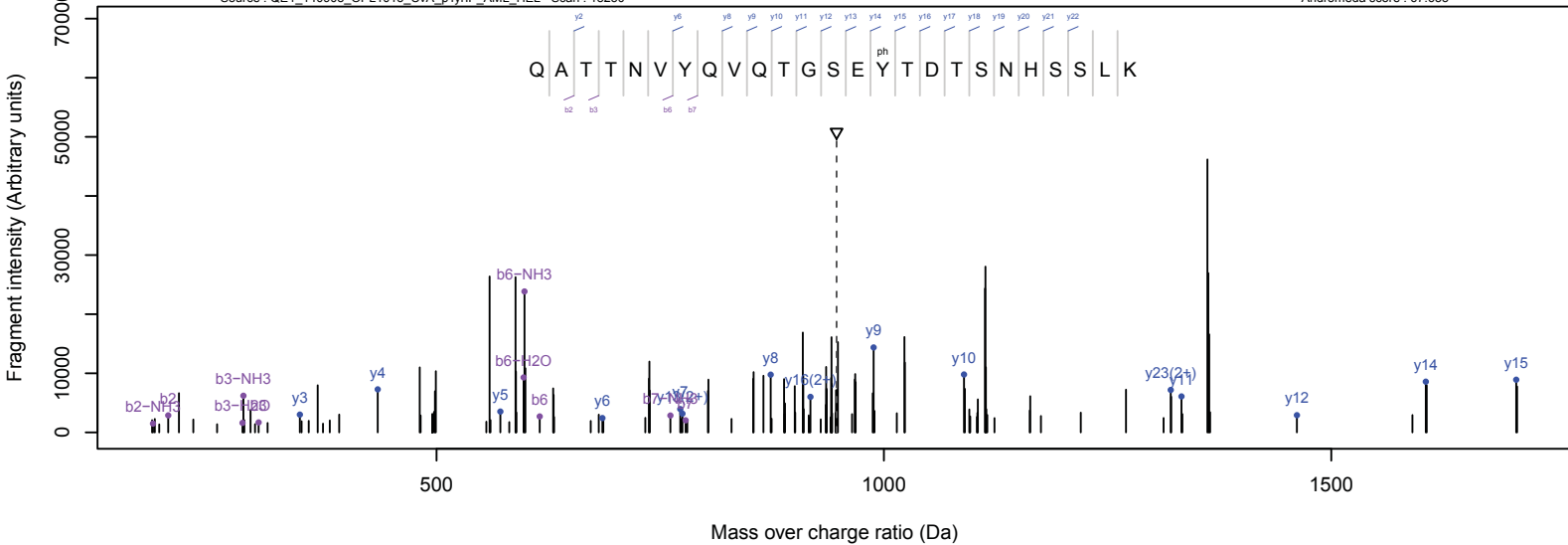
Andromeda score : 67.656



## QATTNVYQVQTGSEY(ph)TDSNHSSLK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 18250

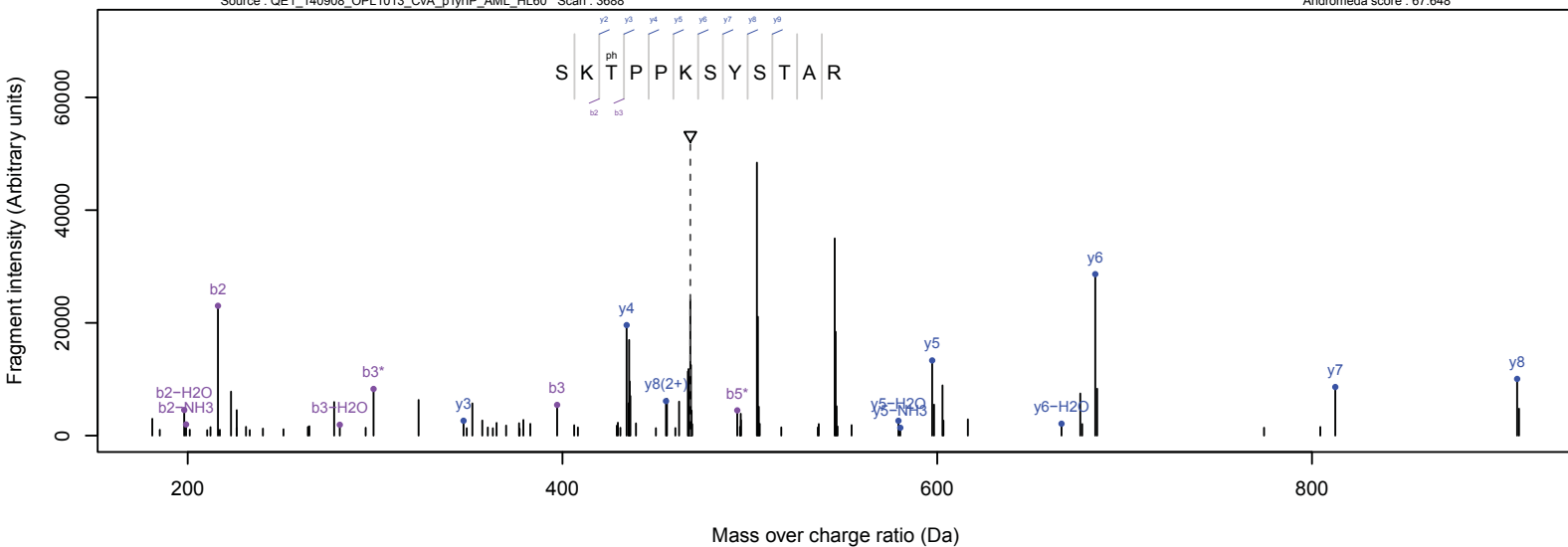
Andromeda score : 67.655



## SKT(ph)PPKSYSTAR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 3688

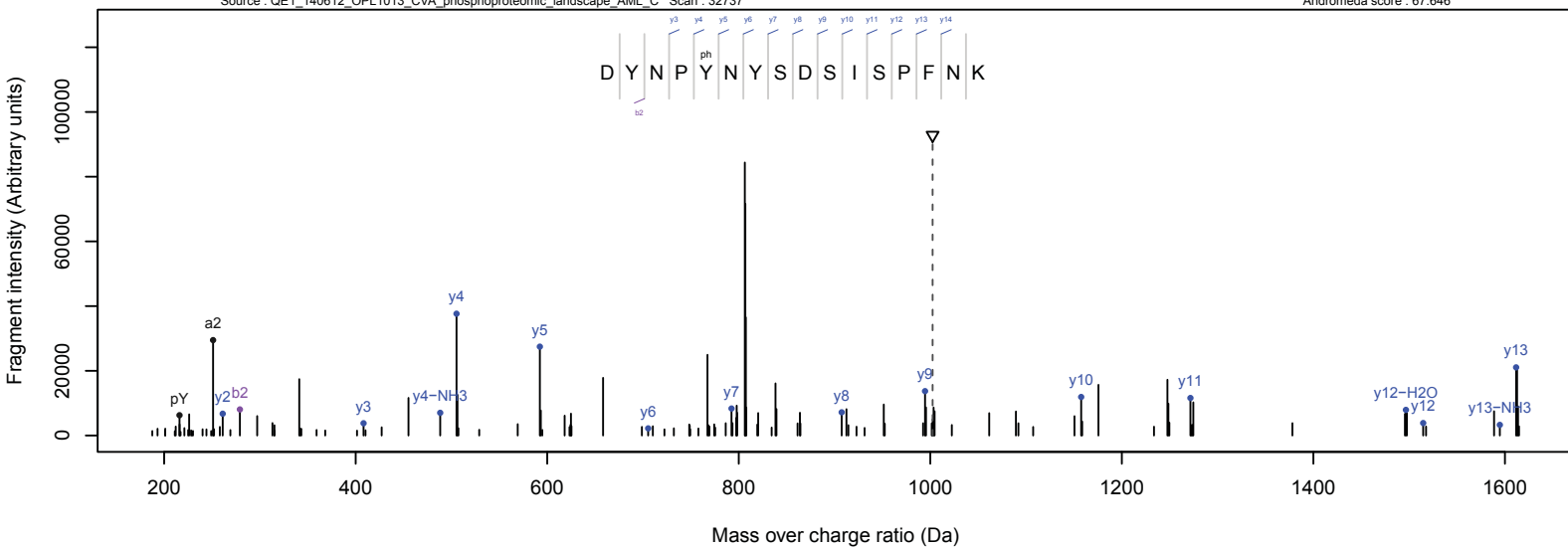
Andromeda score : 67.648



## DYNPY(ph)NYSDSISPFNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32737

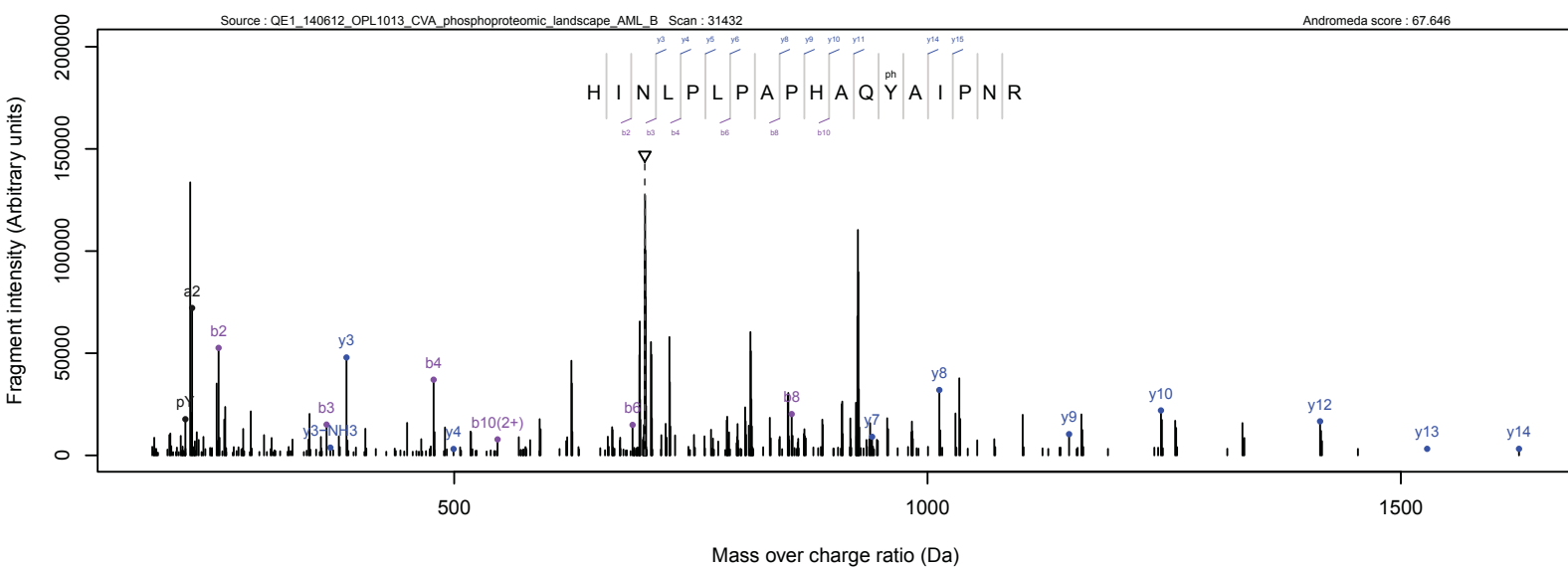
Andromeda score : 67.646



## HINLPLPAPHAQY(ph)AIPNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31432

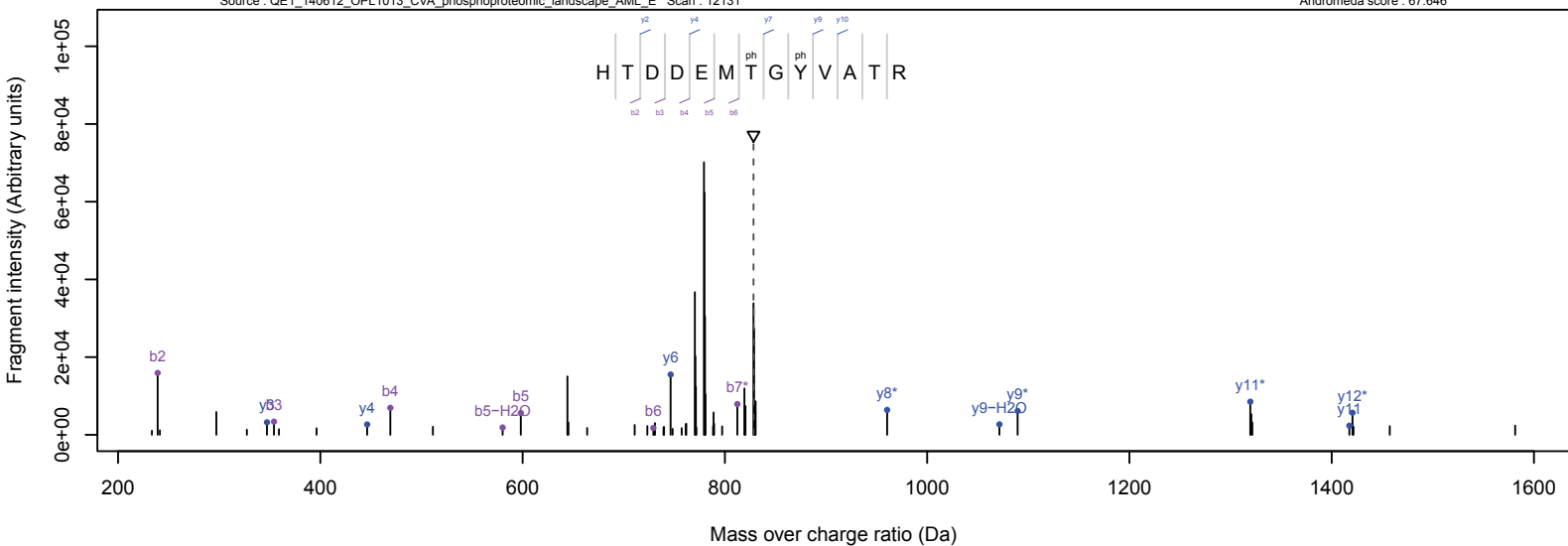
Andromeda score : 67.646



## HTDDEMT(ph)GYVATR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 12131

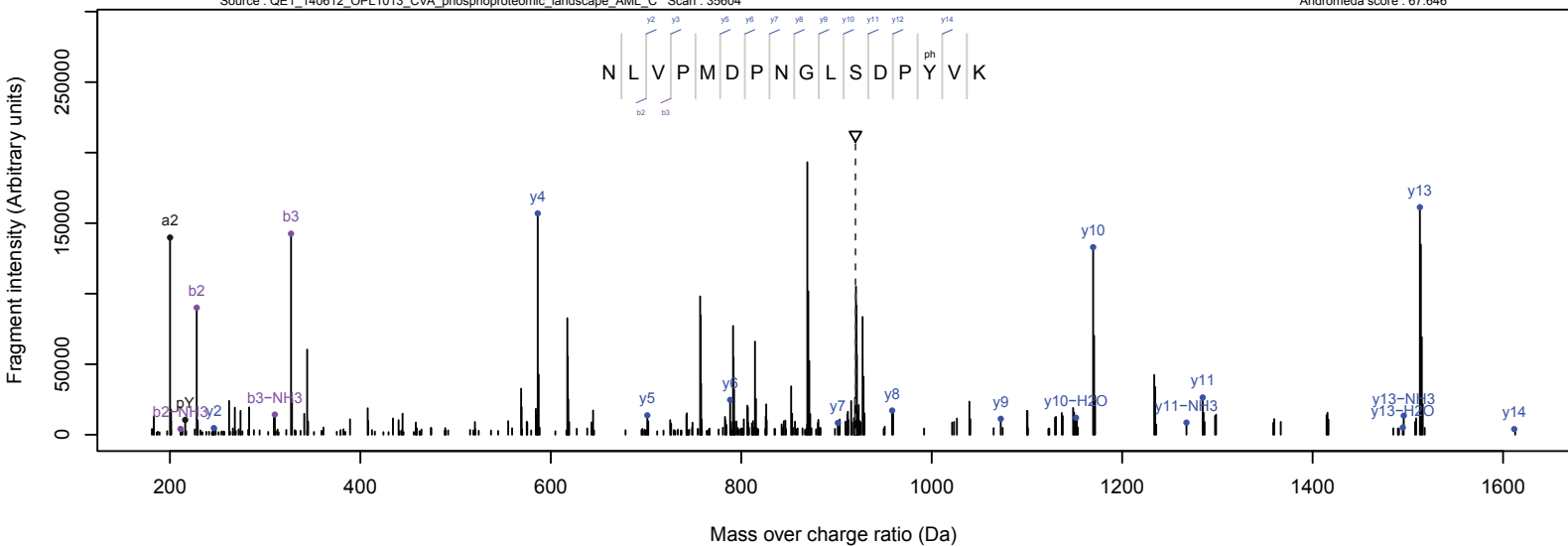
Andromeda score: 67.646



## NLVMPDPNGLSDPY(ph)VK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 35604

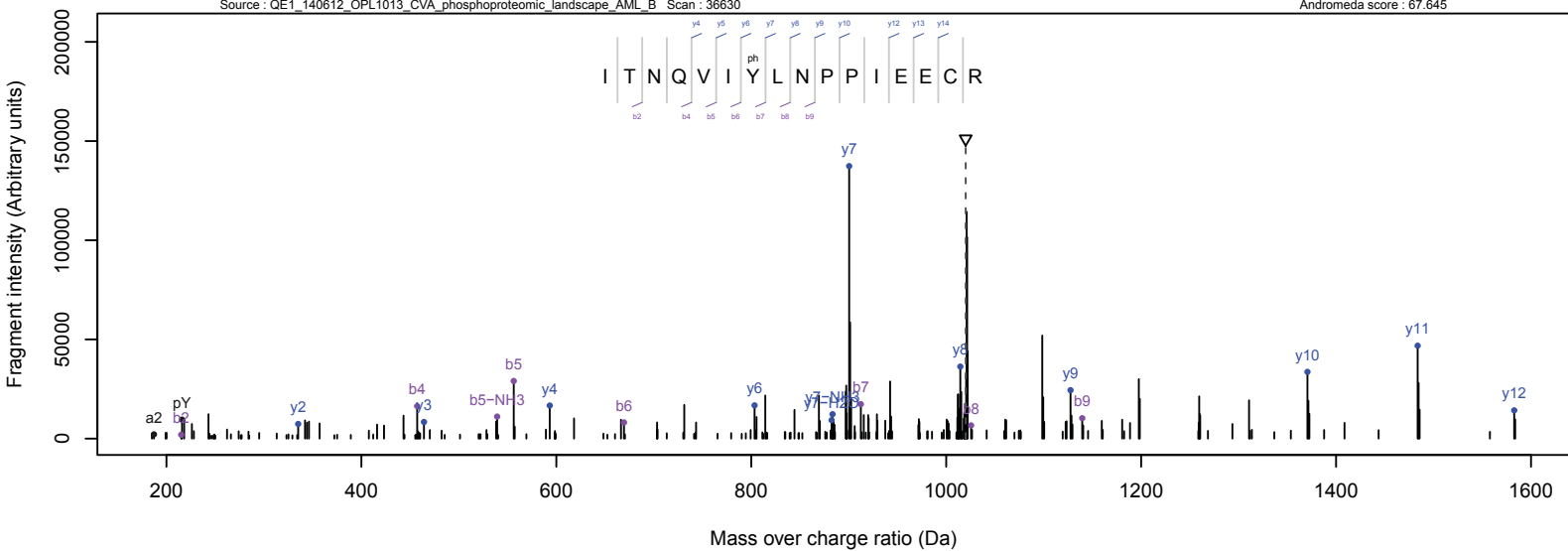
Andromeda score: 67.646



## ITNQVIY(L)NPPIEECR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 36630

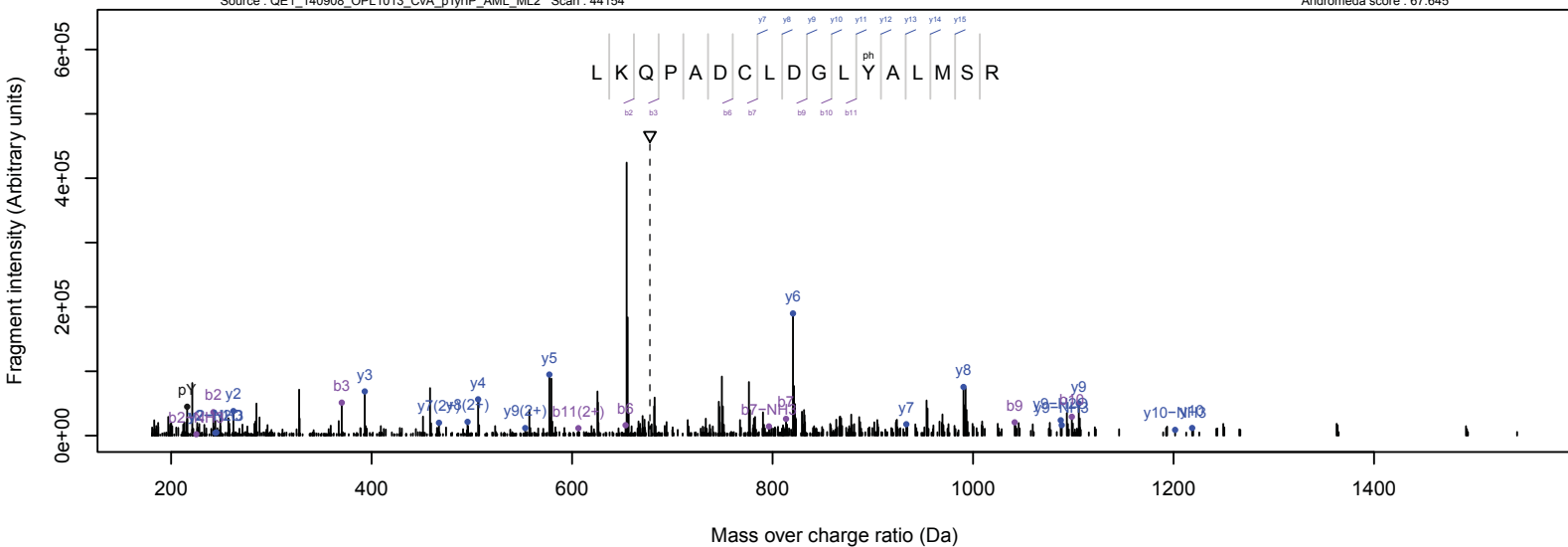
Andromeda score: 67.645



LKQPADCLDGLY(ph)ALMSR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan: 44154

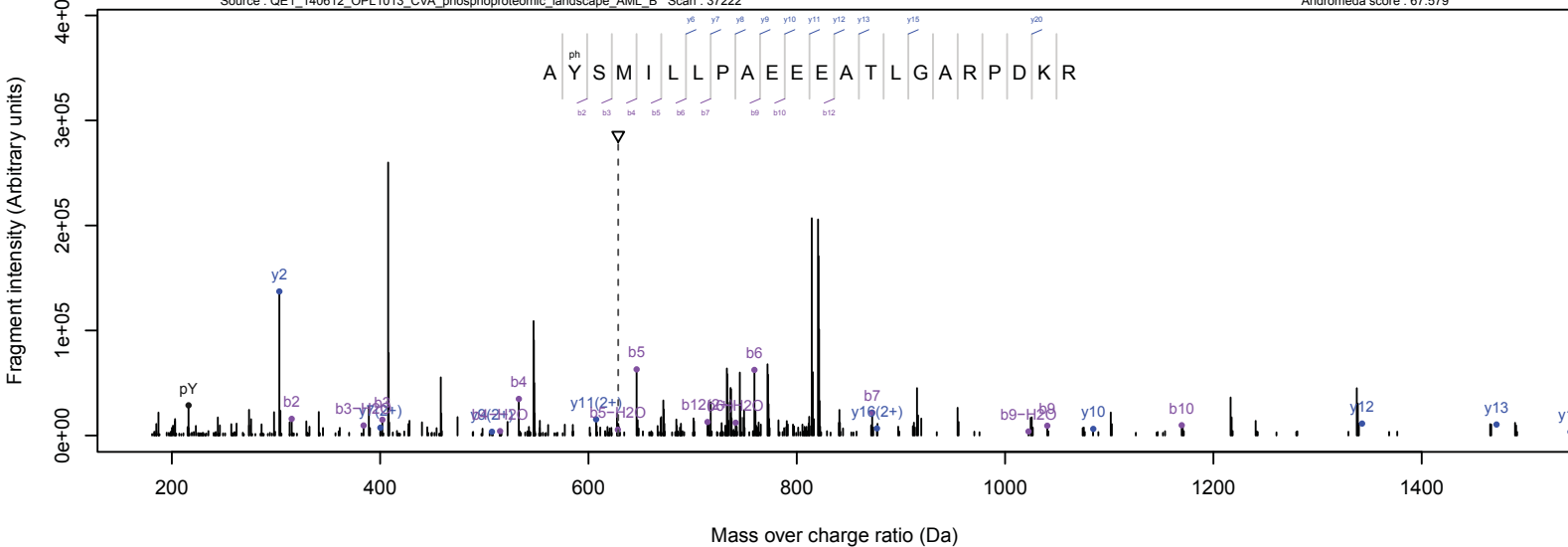
Andromeda score: 67.645



AY(ph)SMILLPAEEEEATLGARPKR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 37222

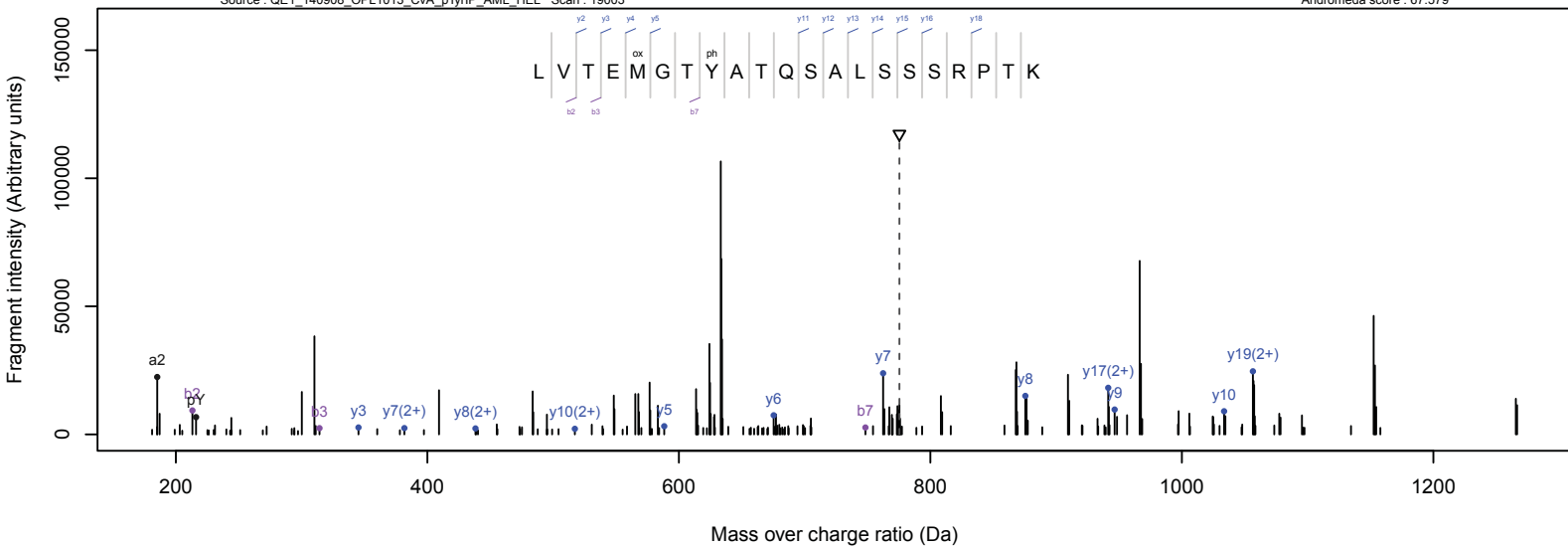
Andromeda score: 67.579

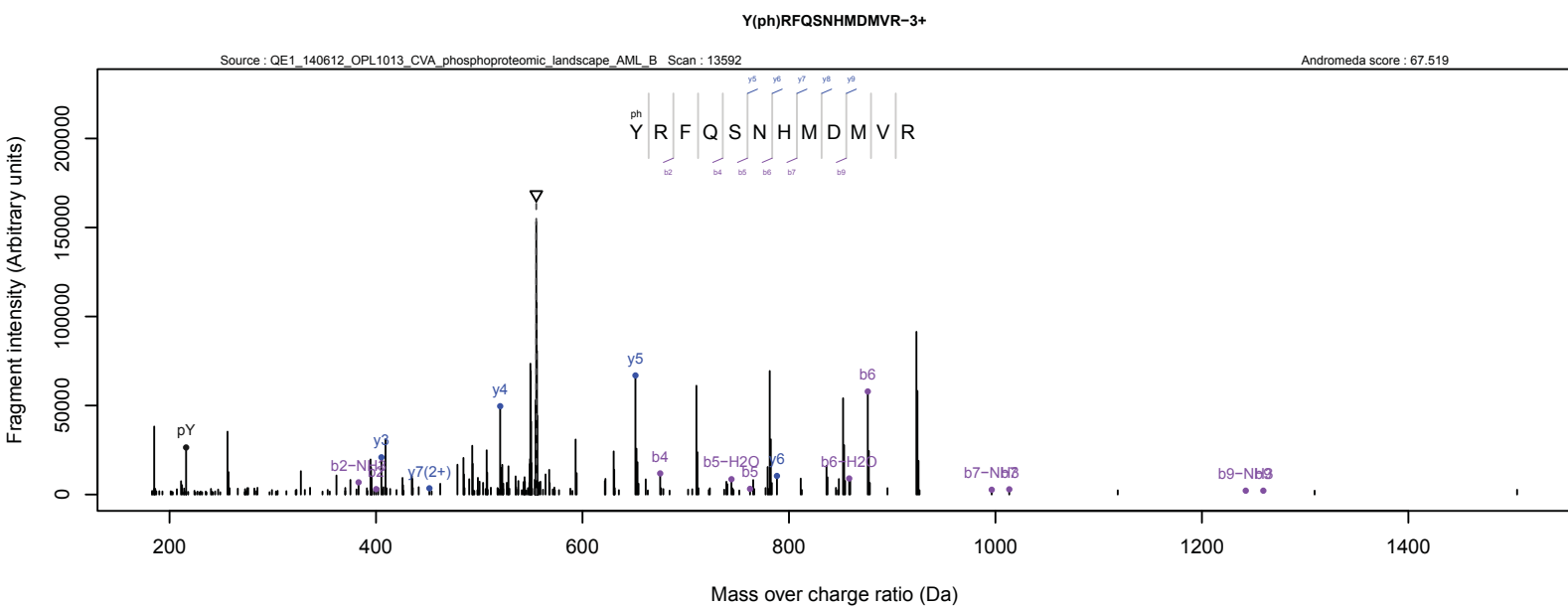
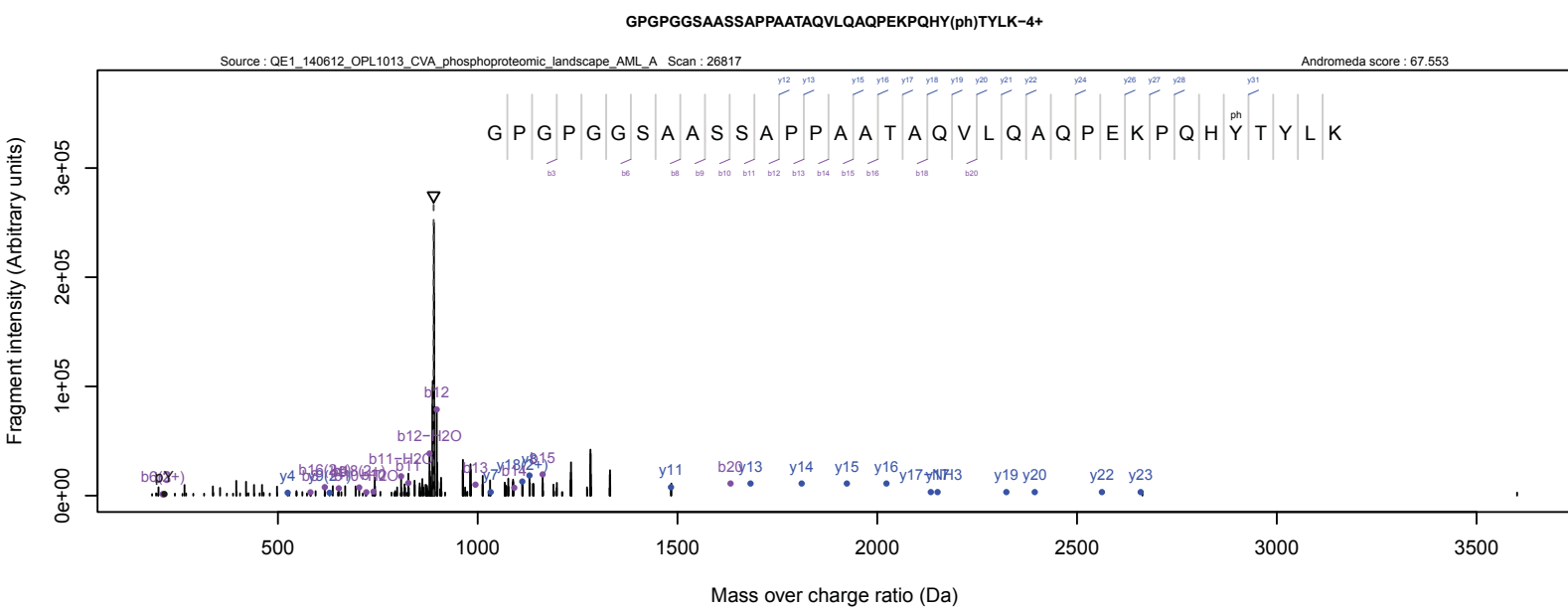
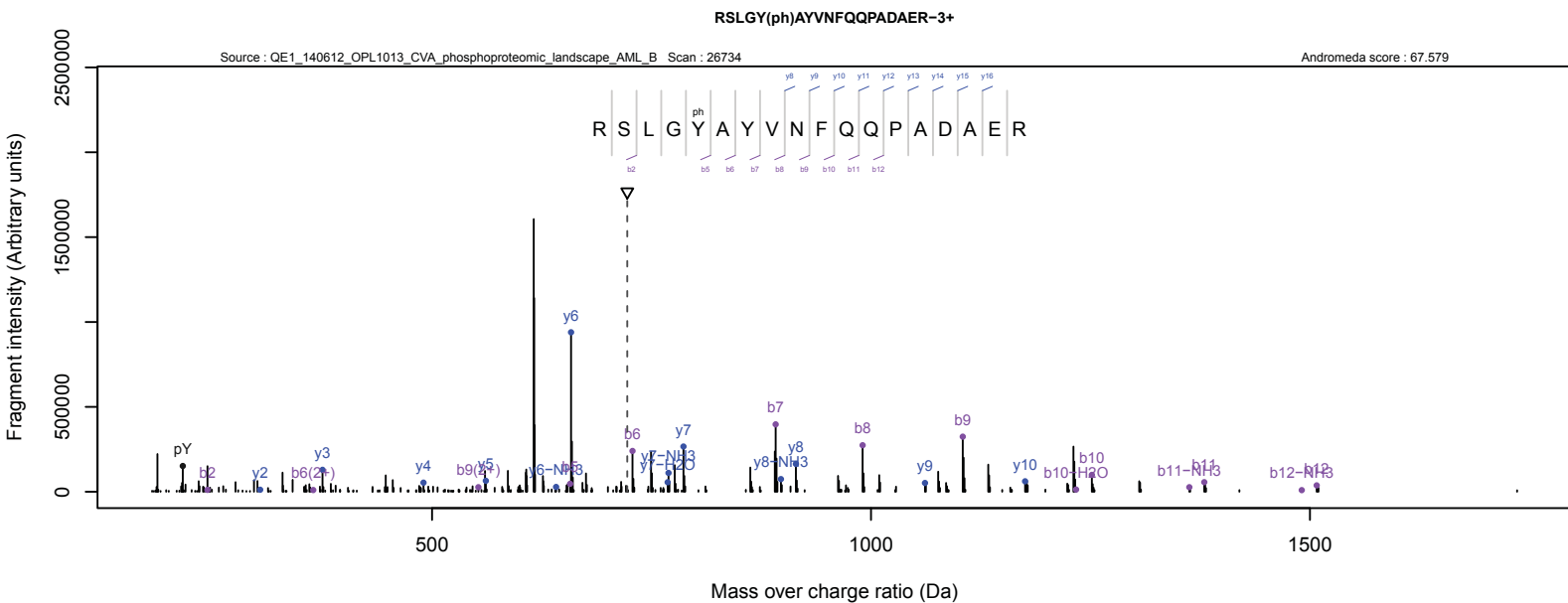


LVTEM(ox)GTY(ph)ATQSALSSSRPTK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 19003

Andromeda score: 67.579

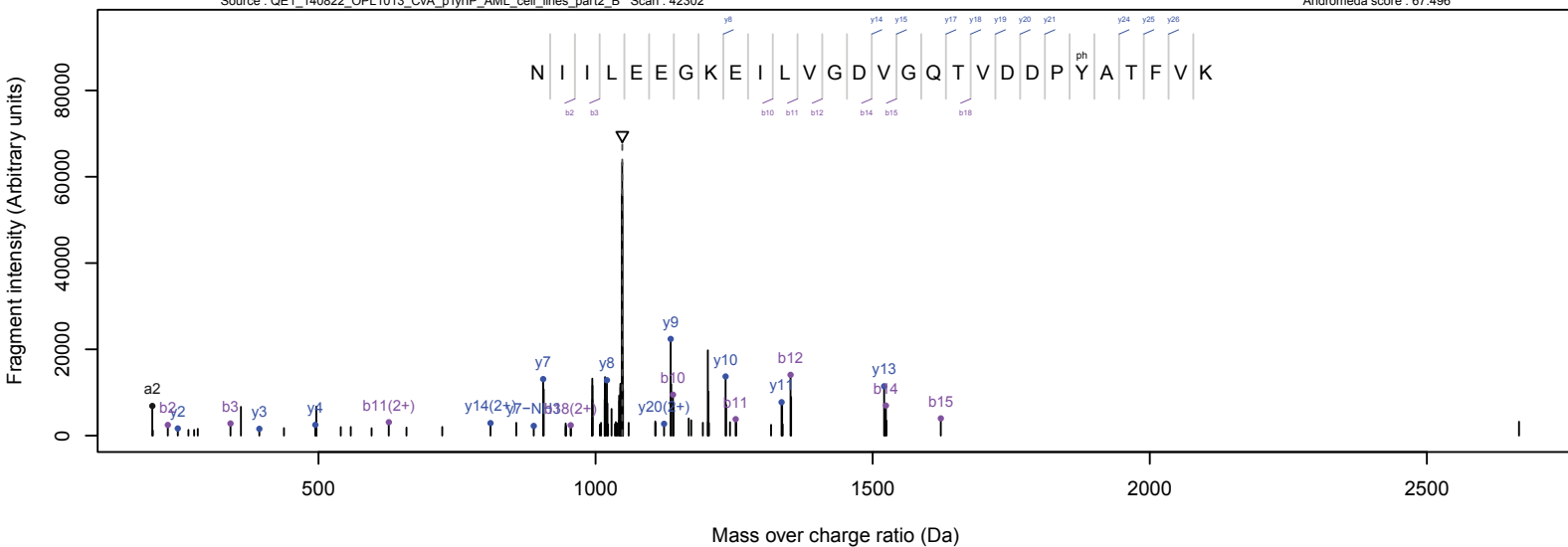




## NILEEGKEILVGDVGGQTVDDPY(ph)ATFVK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 42302

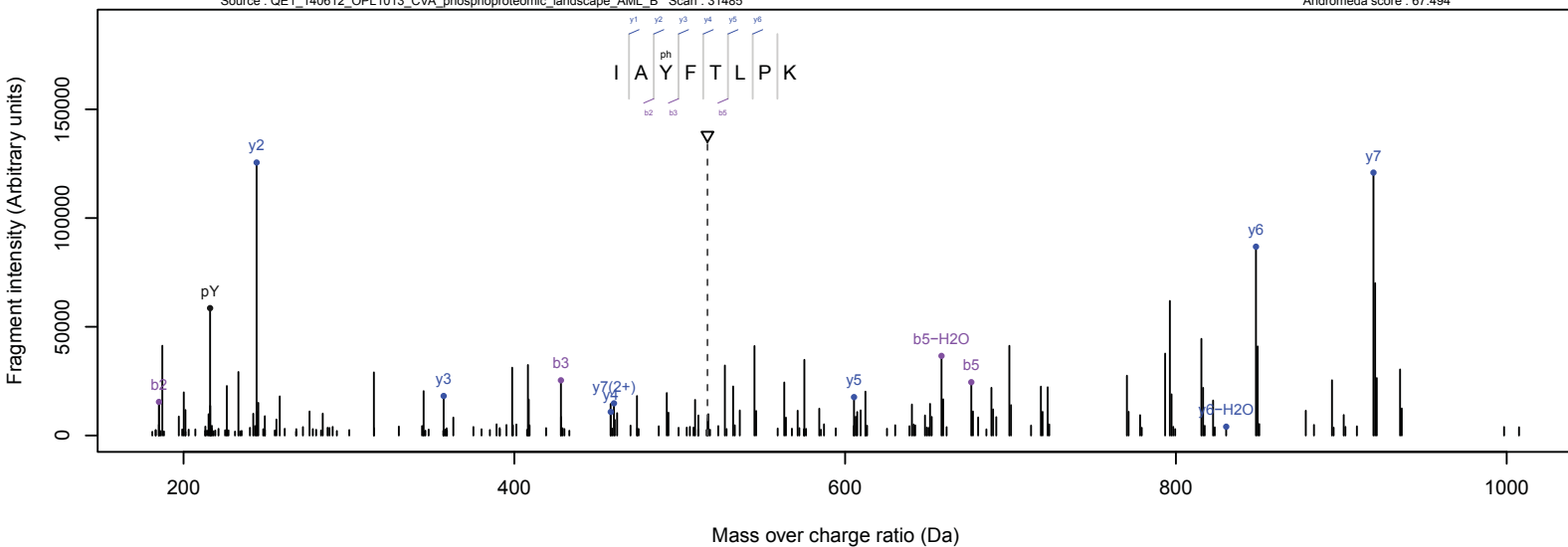
Andromeda score : 67.496



## IAY(ph)FTLPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31485

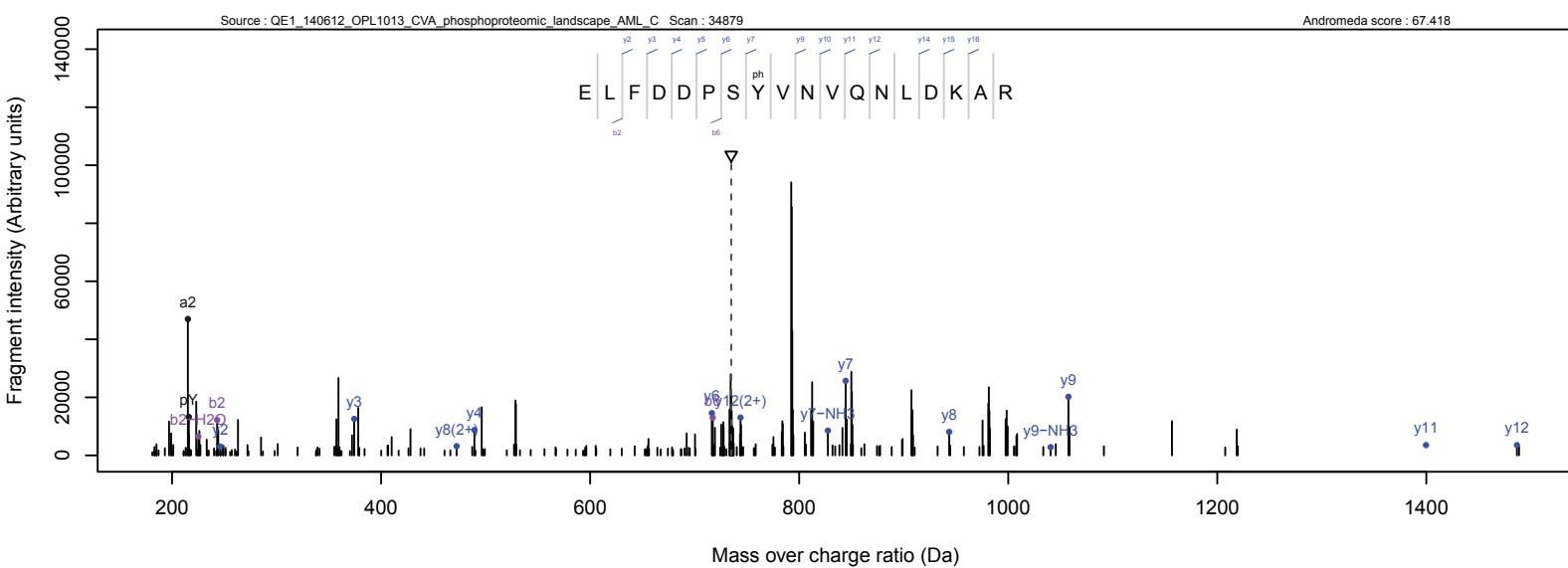
Andromeda score : 67.494



## ELFDDPSY(ph)VNVQNLDKAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34879

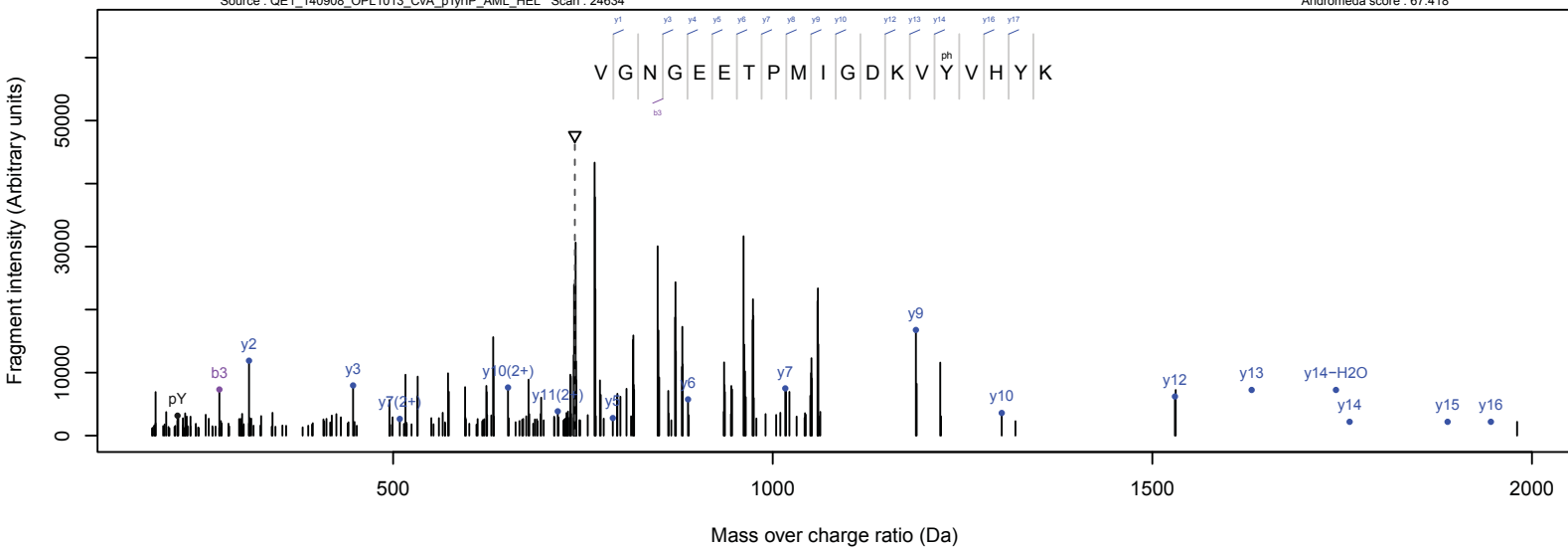
Andromeda score : 67.418



**VGNGEETPMIGDKVY(ph)VHYK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24634

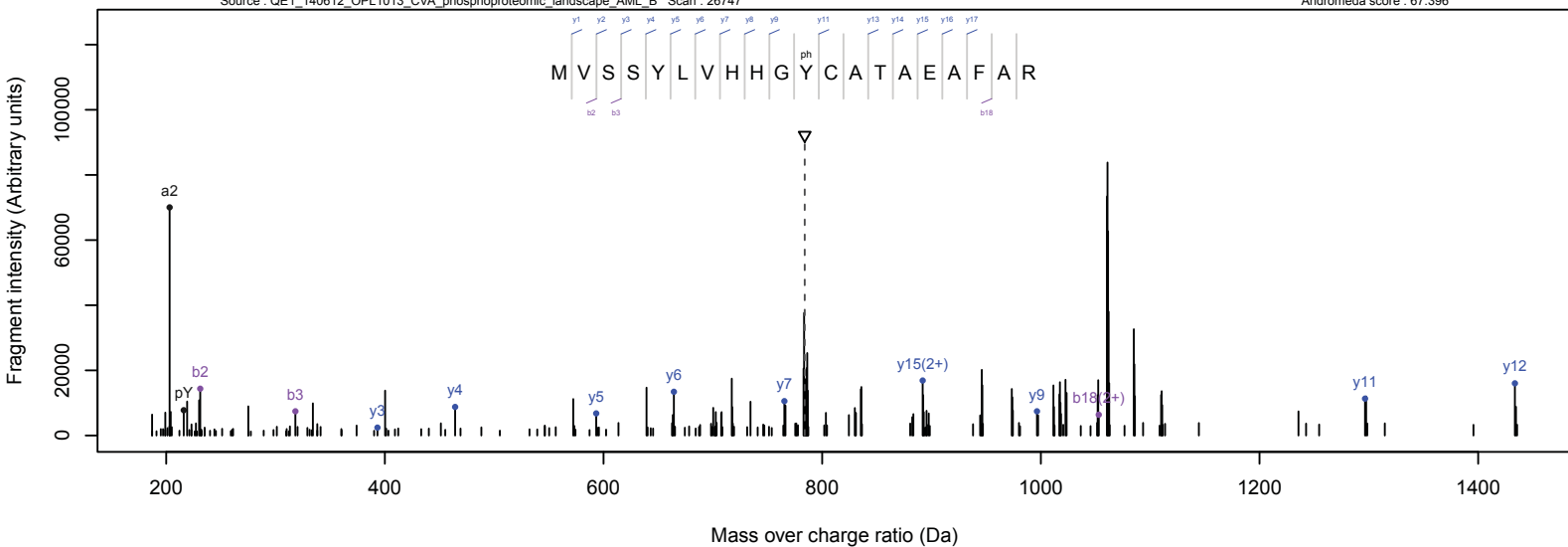
Andromeda score : 67.418



**MVSSYLHHGY(ph)CATAEFAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26747

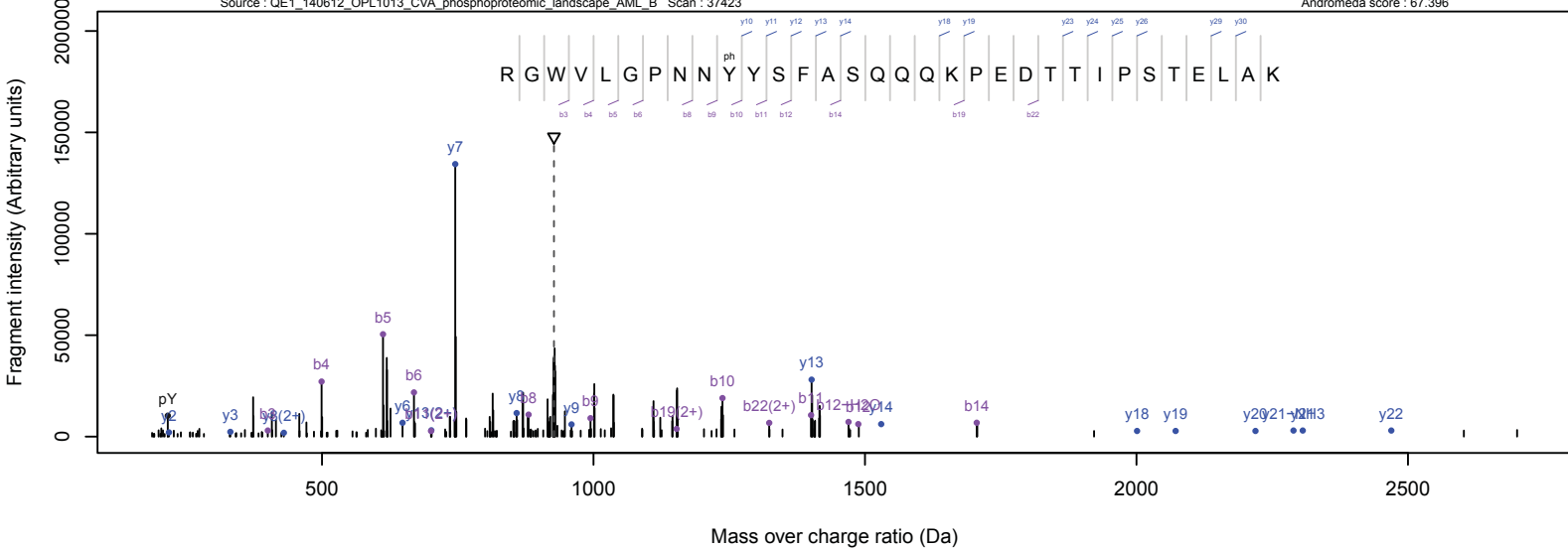
Andromeda score : 67.396



**RGWVLGPNNY(ph)YFASQQQKPEDTTIPSTELAK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37423

Andromeda score : 67.396

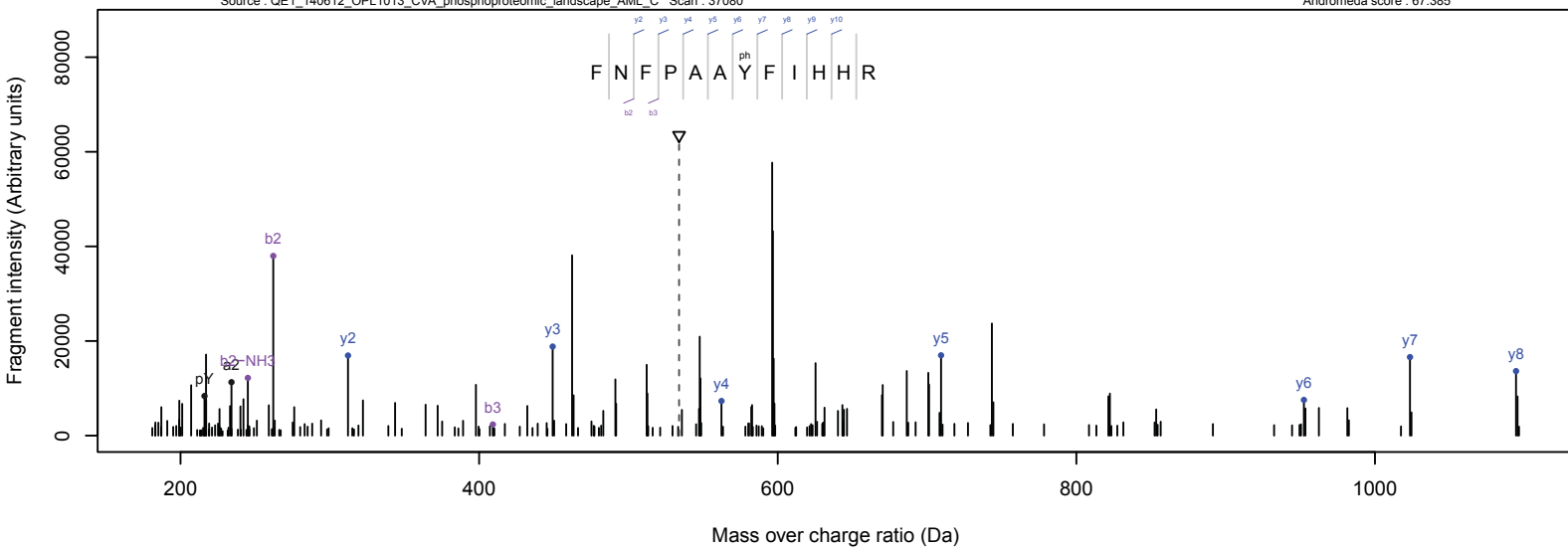




## FNFPAA(ph)FIHHR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37080

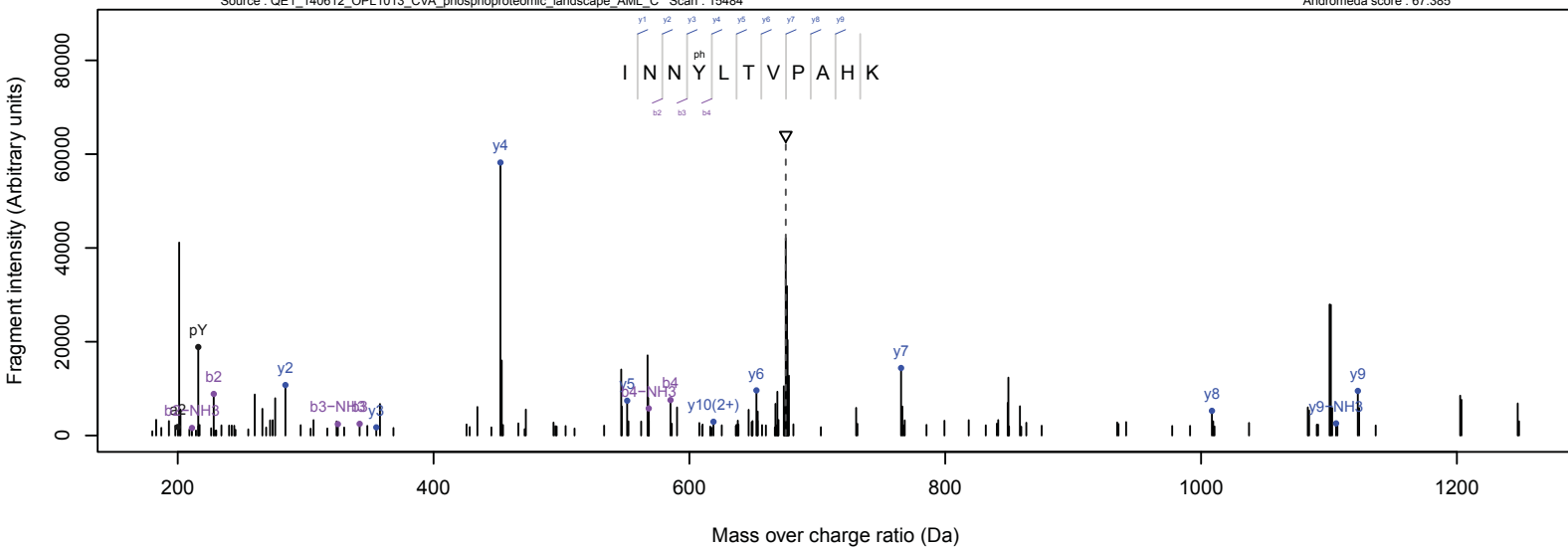
Andromeda score: 67.385



## INNY(ph)LTVPAHK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15484

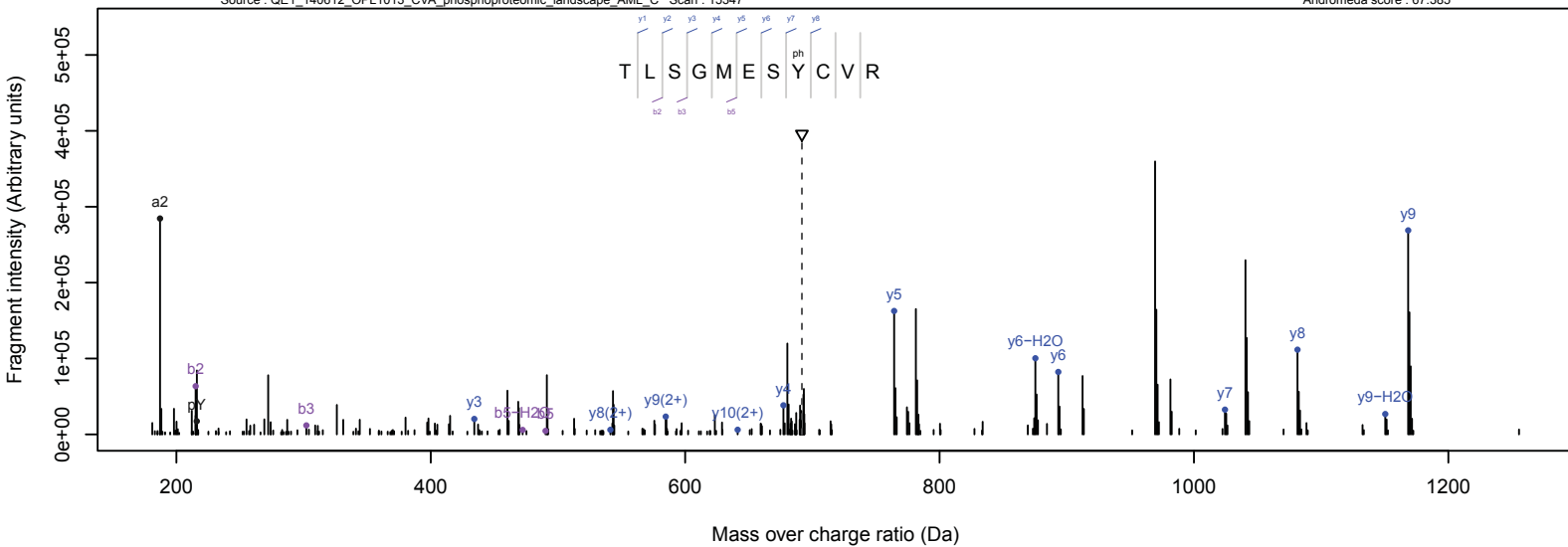
Andromeda score: 67.385



## TLSGMESY(ph)CVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15347

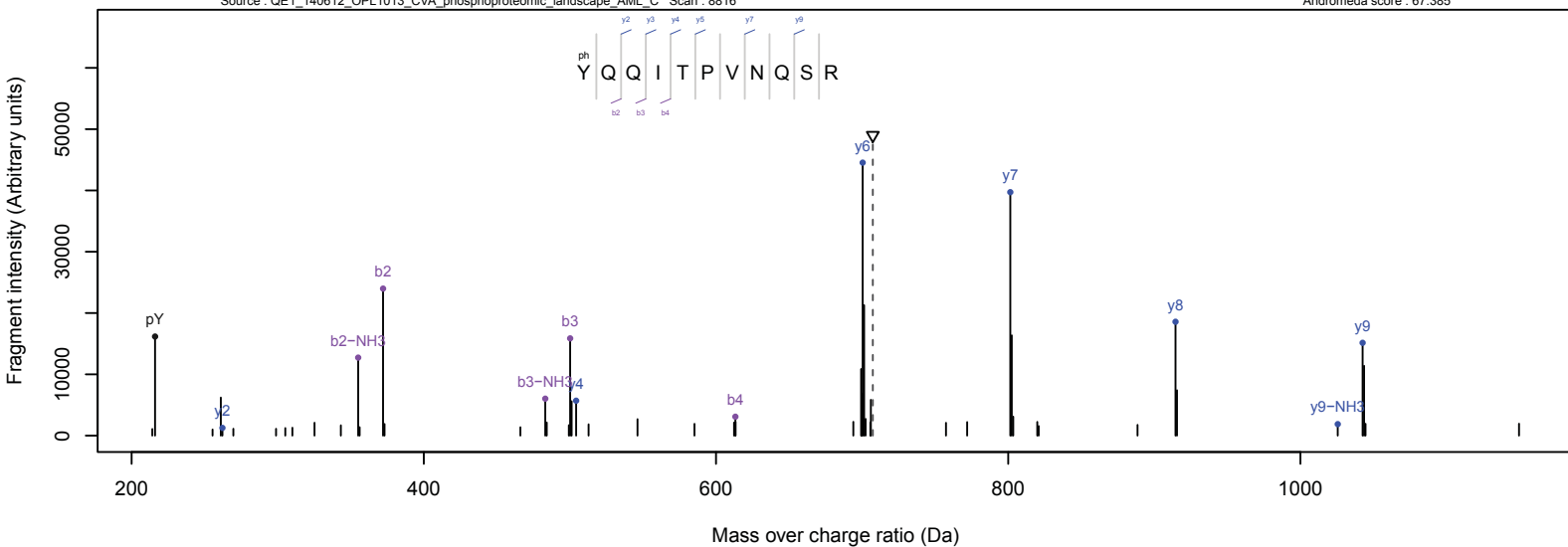
Andromeda score: 67.385



## Y(ph)QQITPVNQRS-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8816

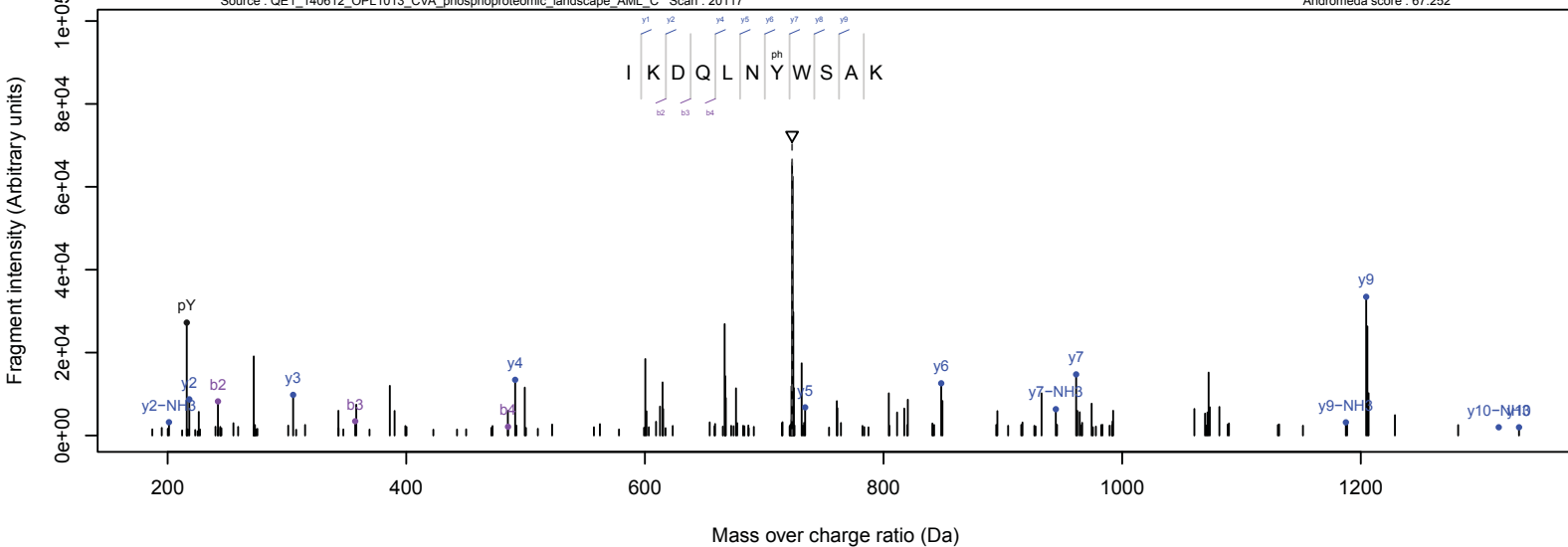
Andromeda score : 67.385



## IKDQLNY(ph)WSAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20117

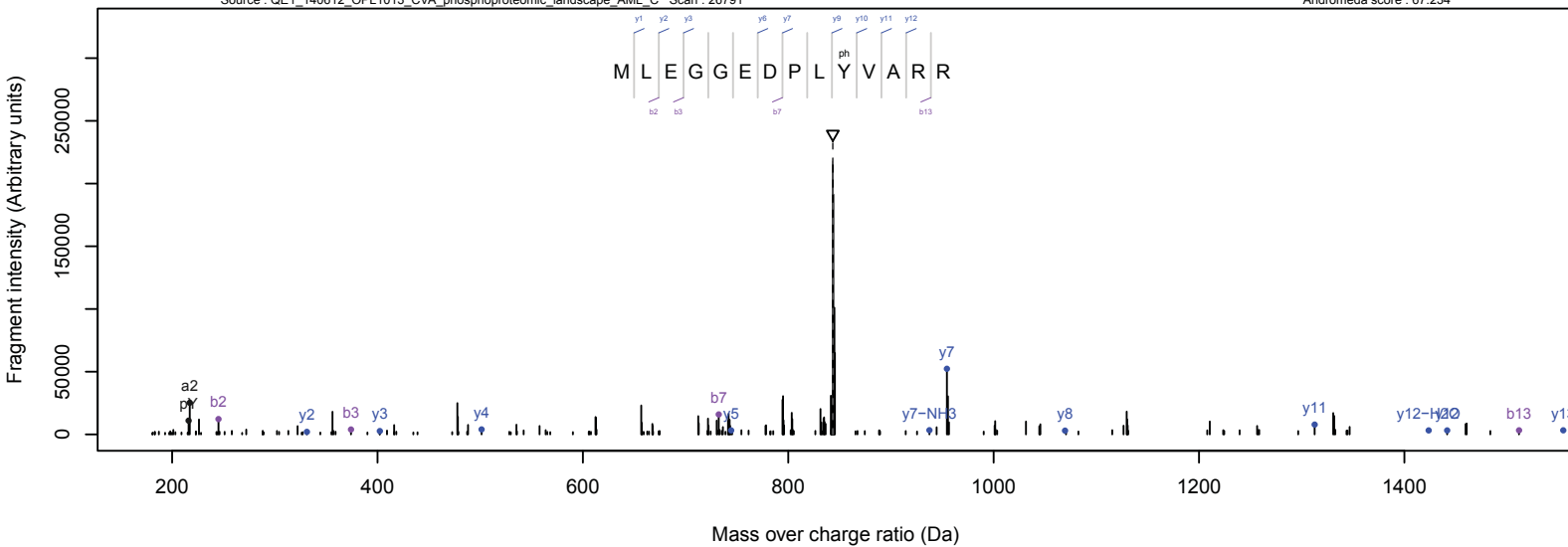
Andromeda score : 67.252



## MLEGGEDPLY(ph)VARR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26791

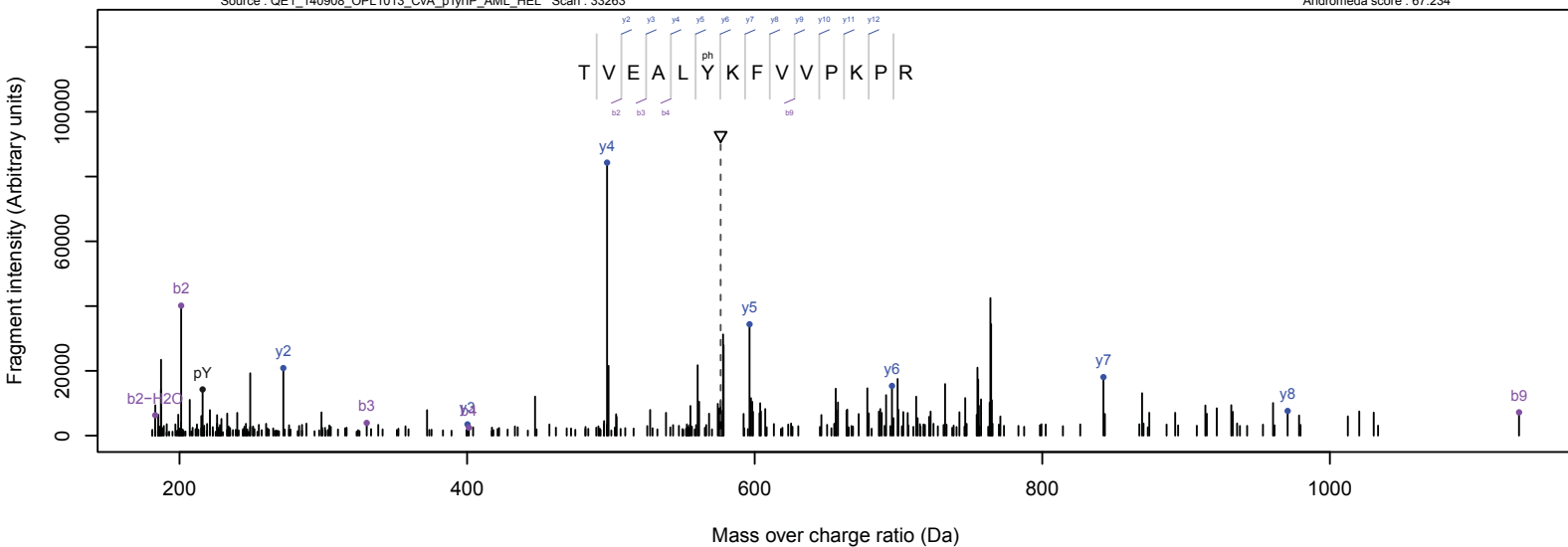
Andromeda score : 67.234



## TVEALY(ph)KFVVPKPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 33263

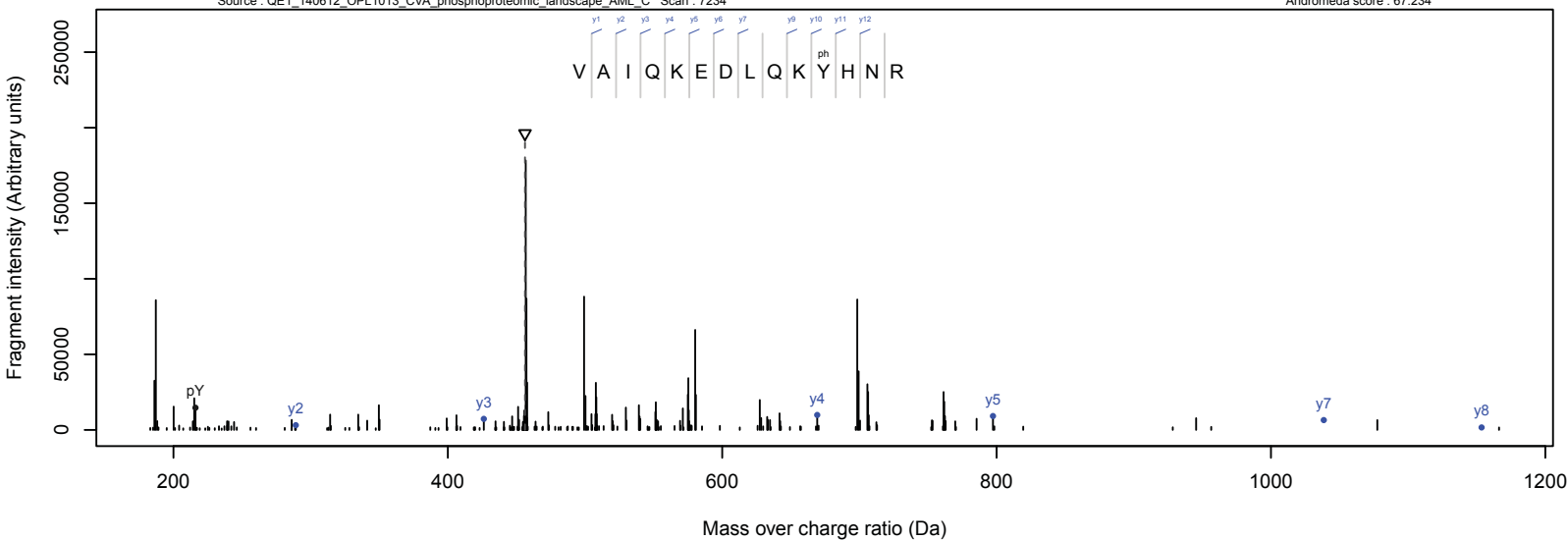
Andromeda score : 67.234



## VAIQKEDLQY(ph)HNR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7234

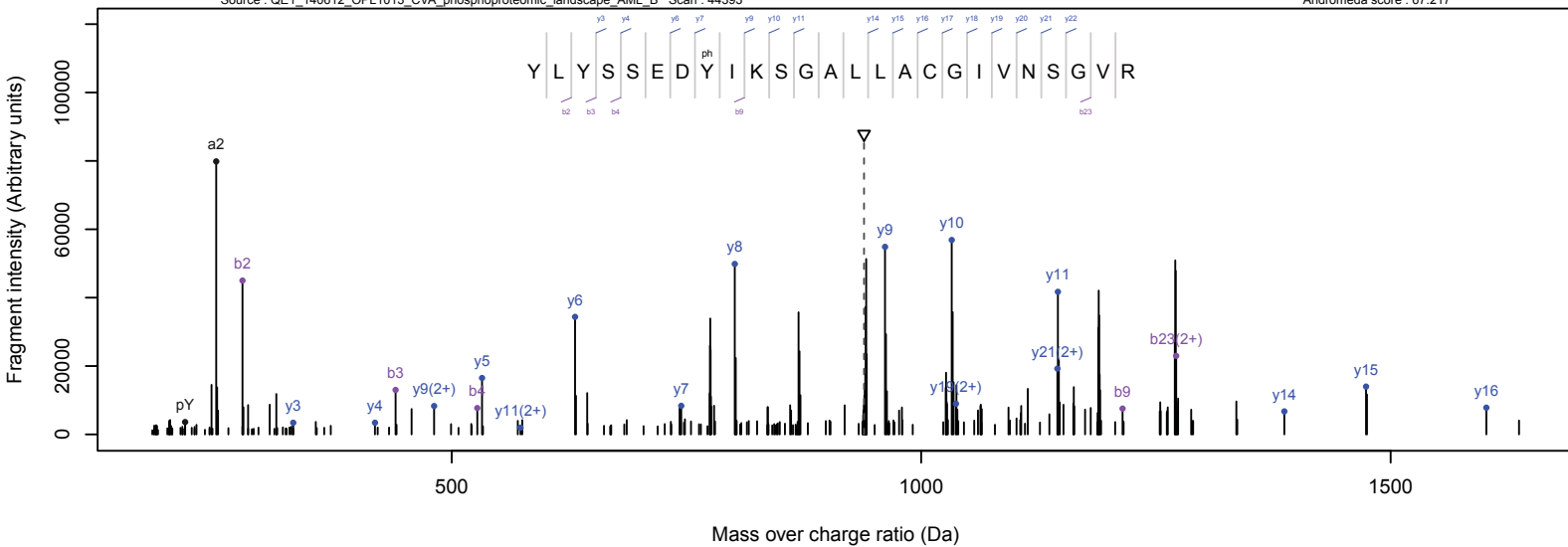
Andromeda score : 67.234



## YLYSSEDY(ph)IKSGALLACGIVNSGVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44393

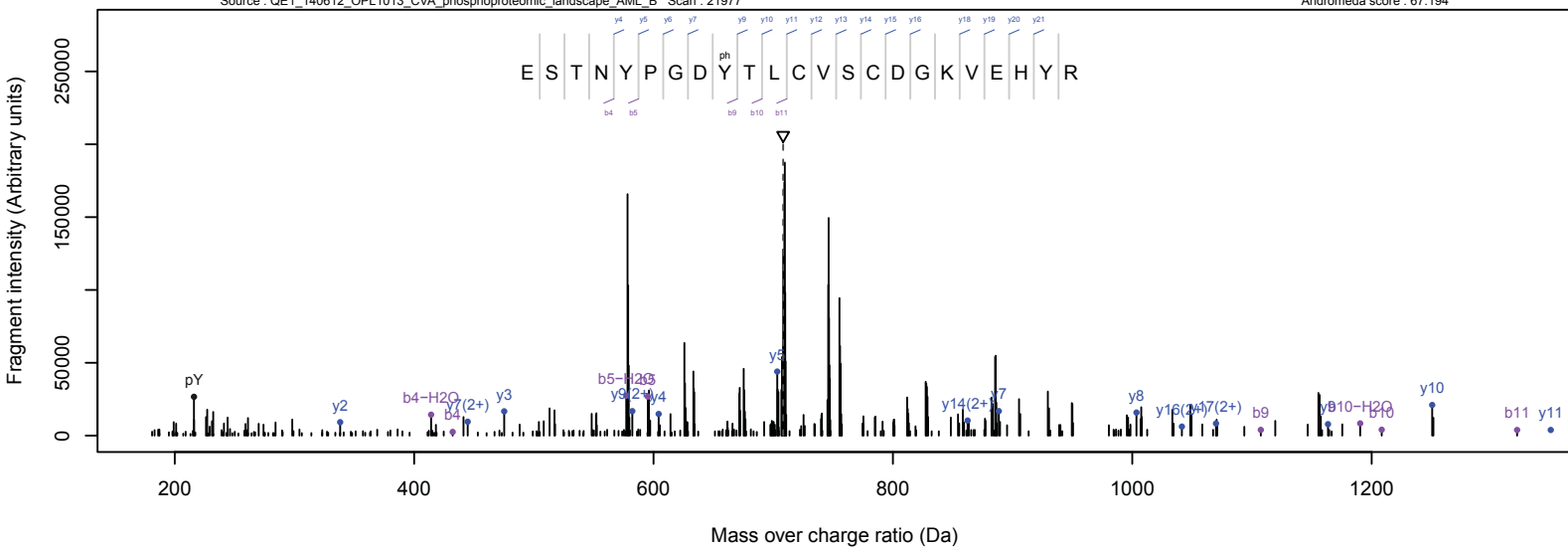
Andromeda score : 67.217



ESTNYPGDY(ph)TLCVSCDGKVEHYR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21977

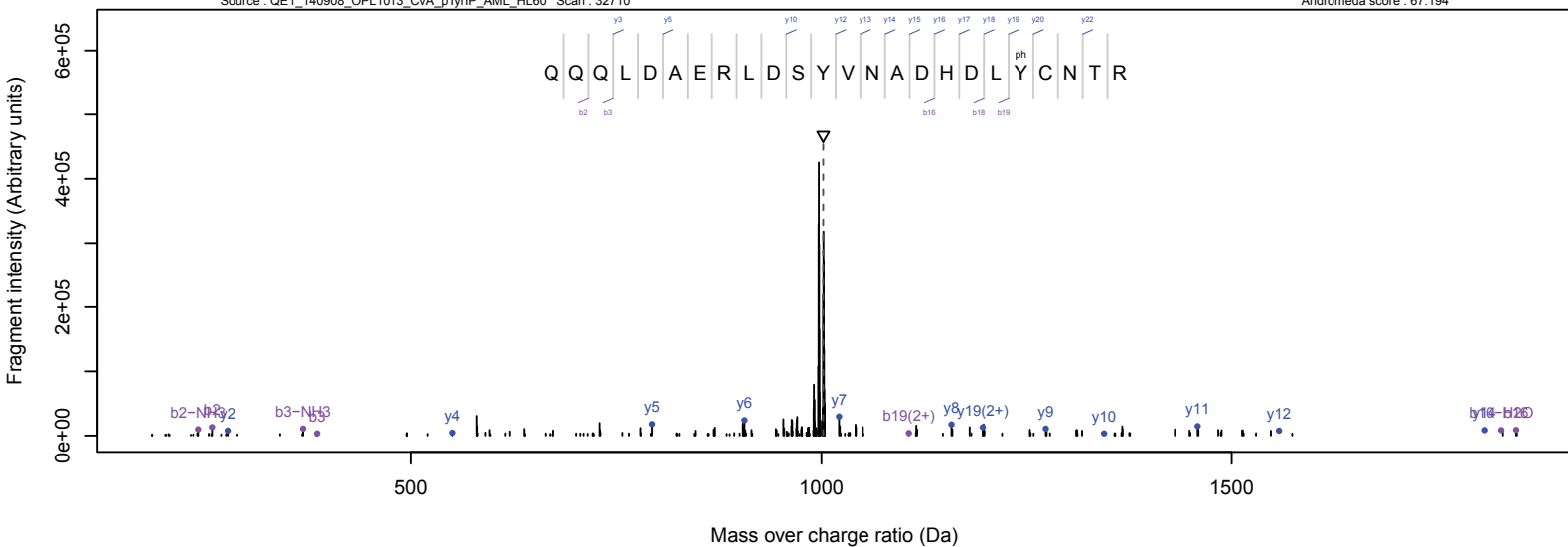
Andromeda score : 67.194



QQQLDAERLDSYVNADHDLY(ph)CNR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 32710

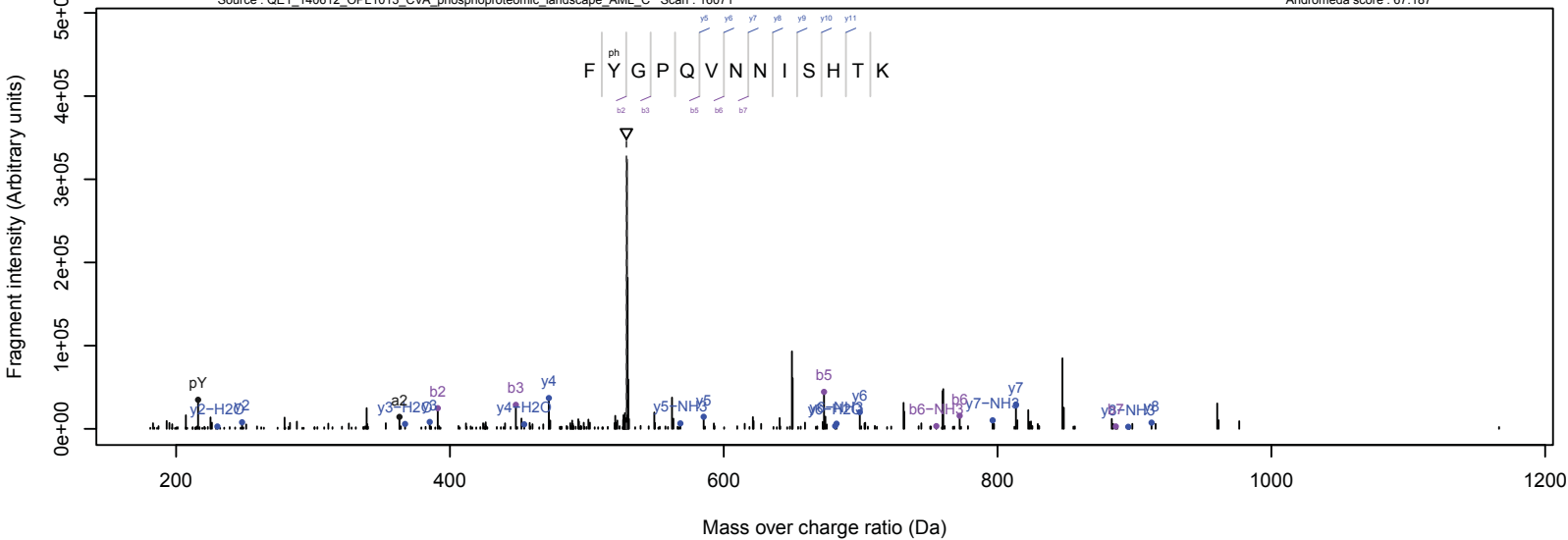
Andromeda score : 67.194



FY(ph)GPQVNNISHTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16071

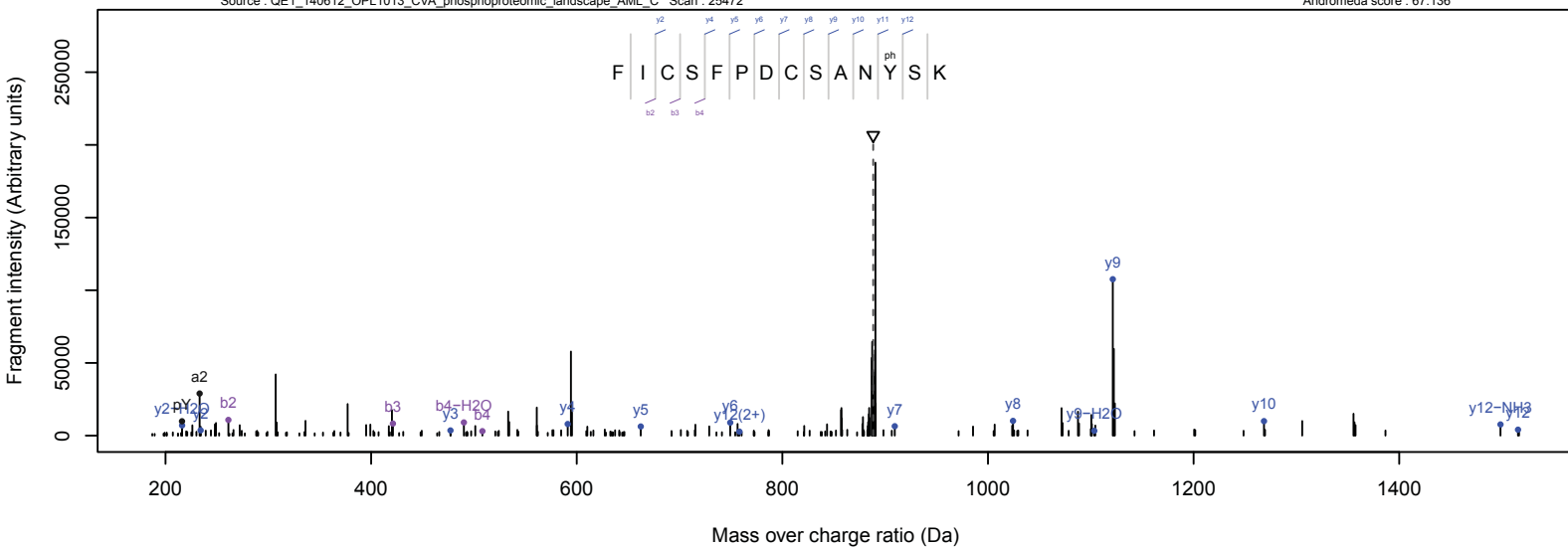
Andromeda score : 67.187



## FICSPDCSANY(ph)SK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25472

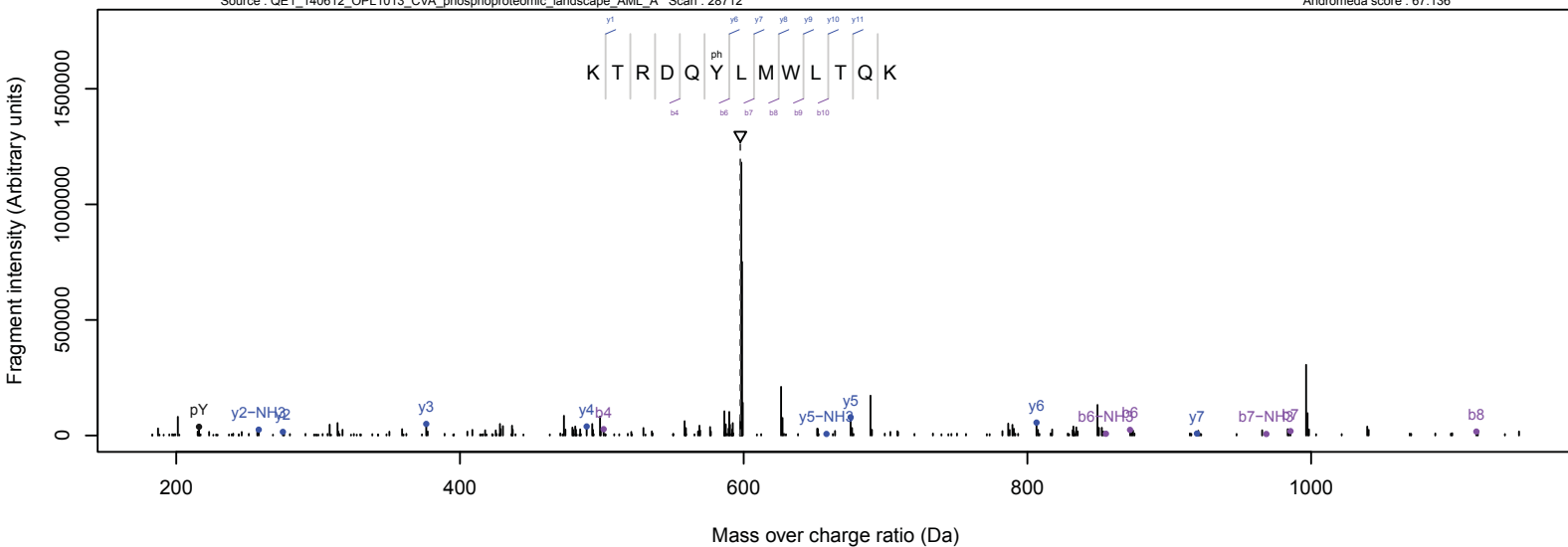
Andromeda score : 67.136



## KTRDQY(ph)LMWLTQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 28712

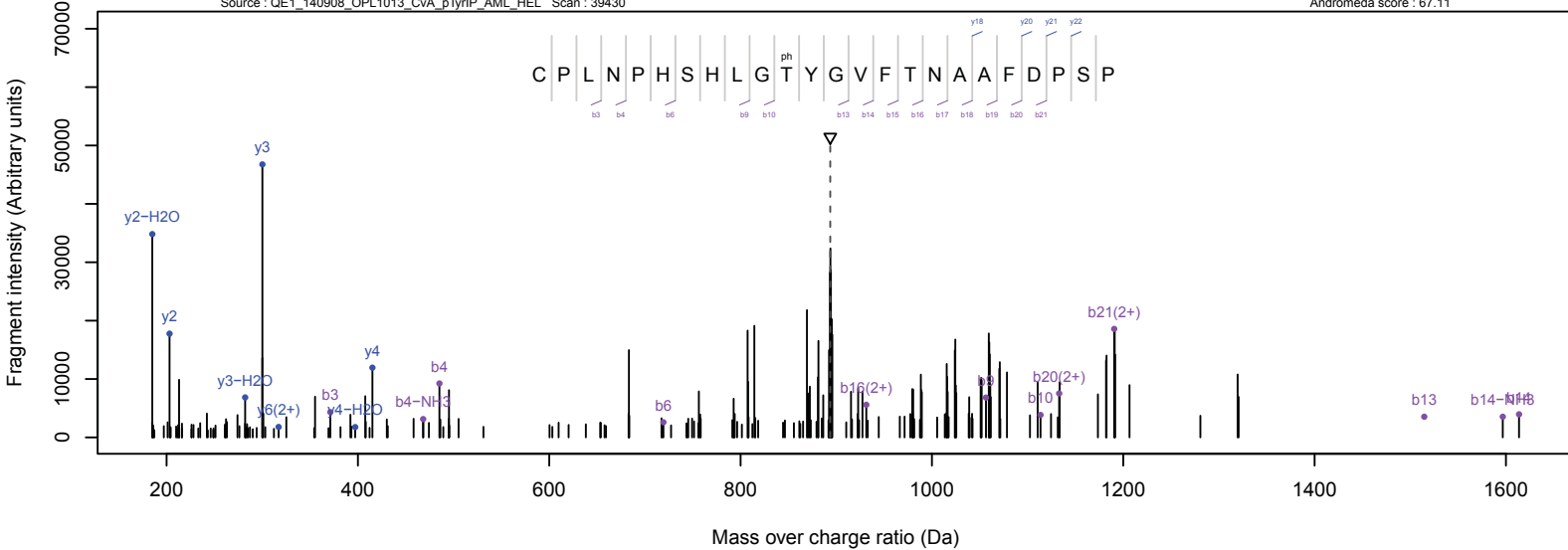
Andromeda score : 67.136



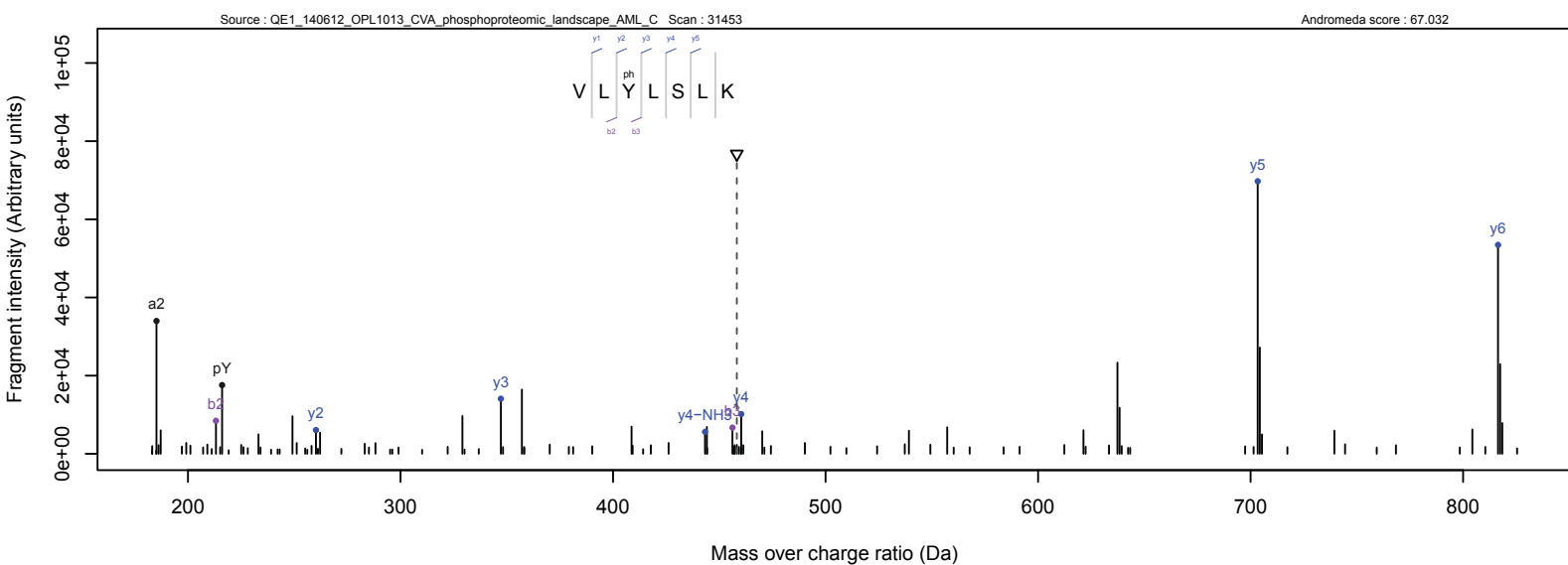
## CPLNPHSHLGT(ph)YGVFTNAAFDPS-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 39430

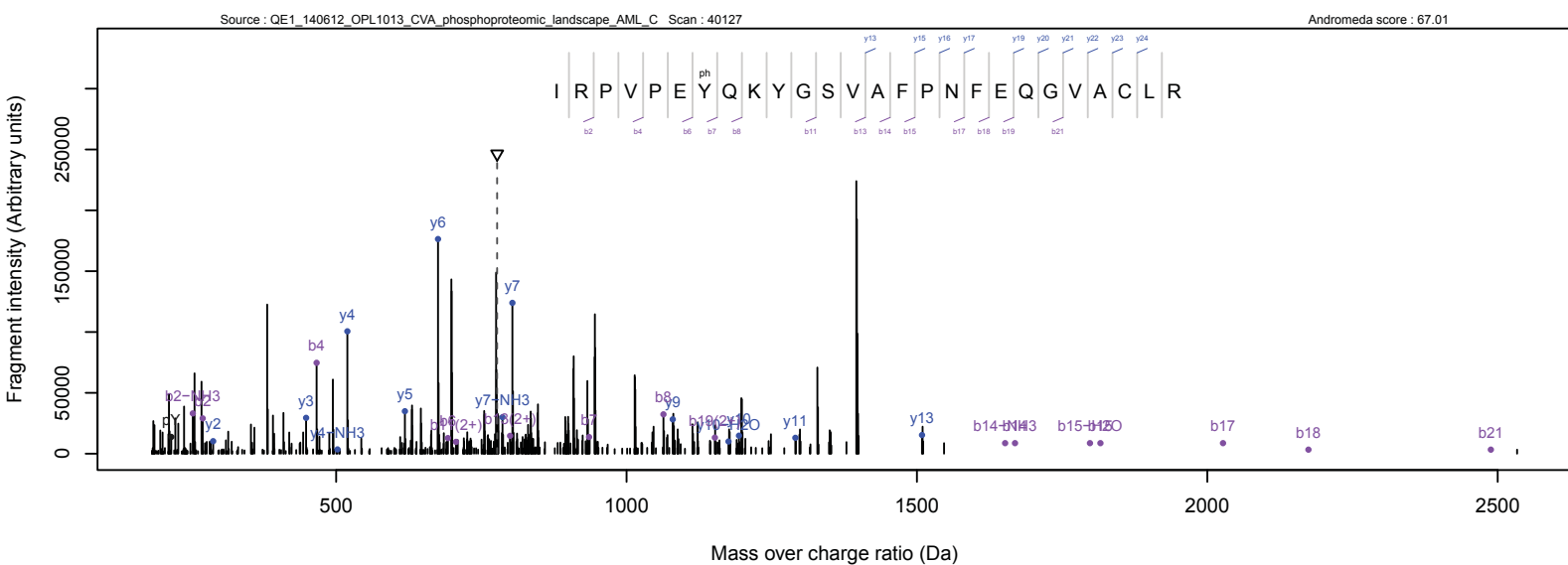
Andromeda score : 67.11



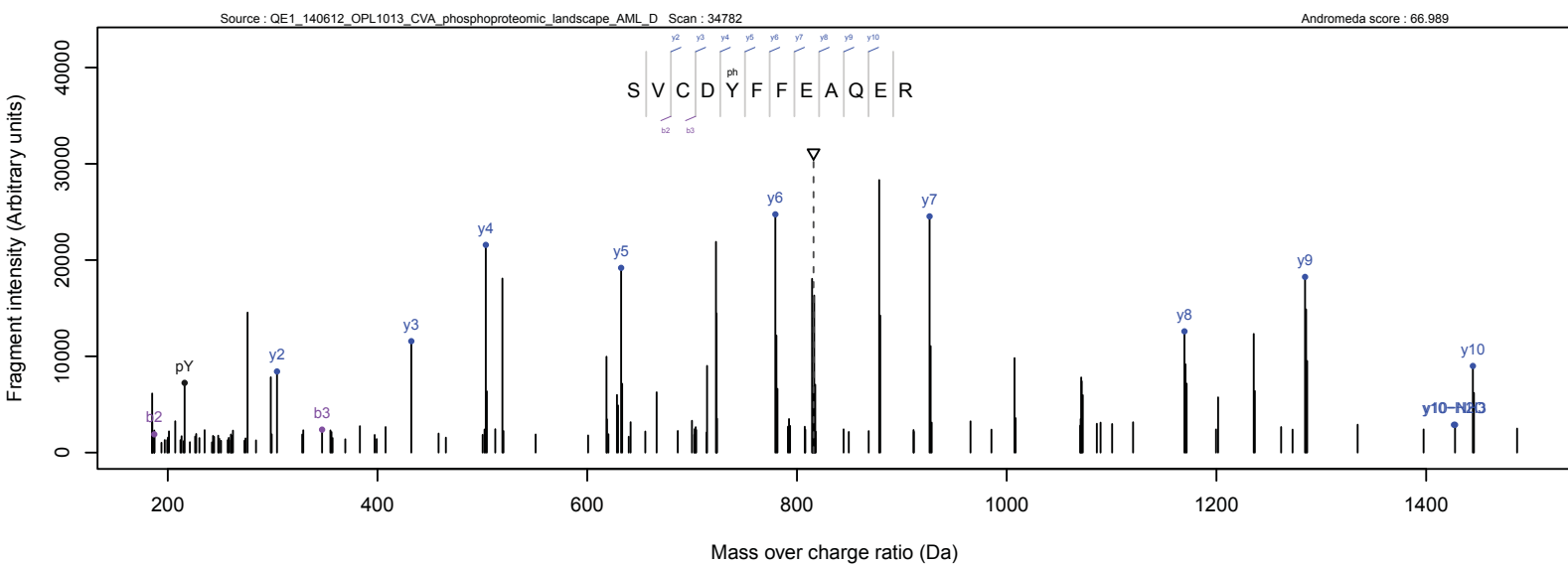
## VLY(ph)LSLK-2+



## IRPVPEY(ph)QKYGSVAFPNFEQGVACL R-4+



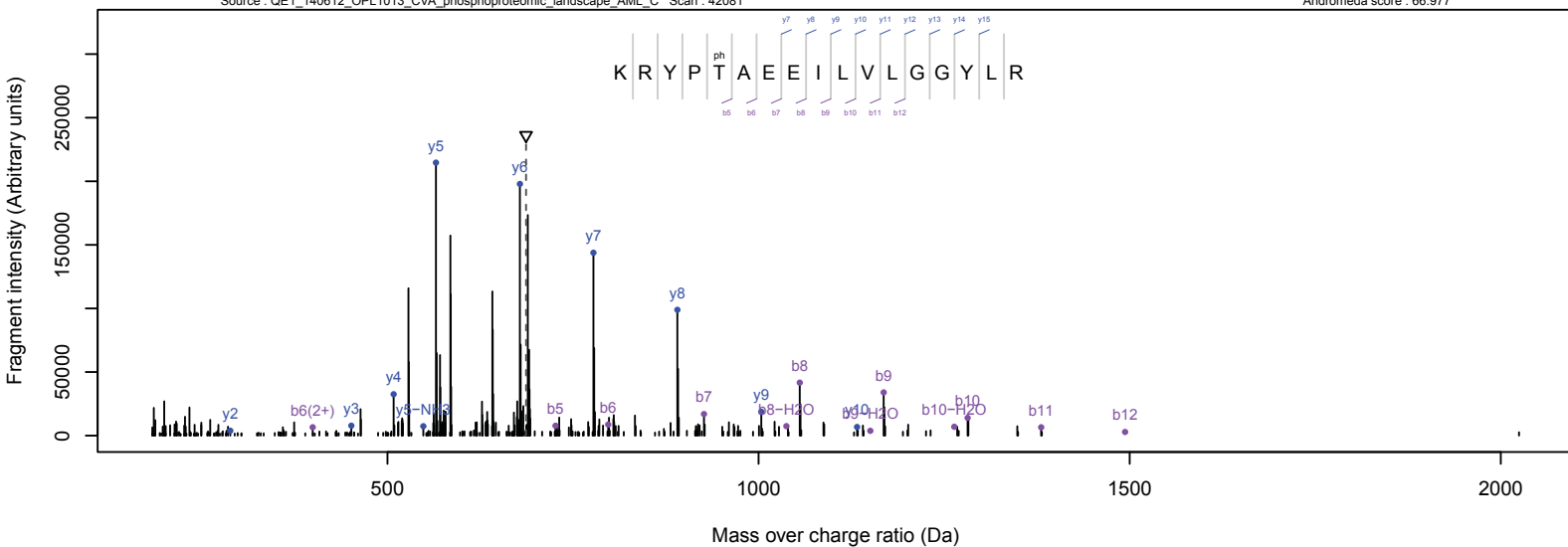
## SVC DY(ph)FFEAQER-2+



**KRYPT(ph)AEEILVGGYLR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 42081

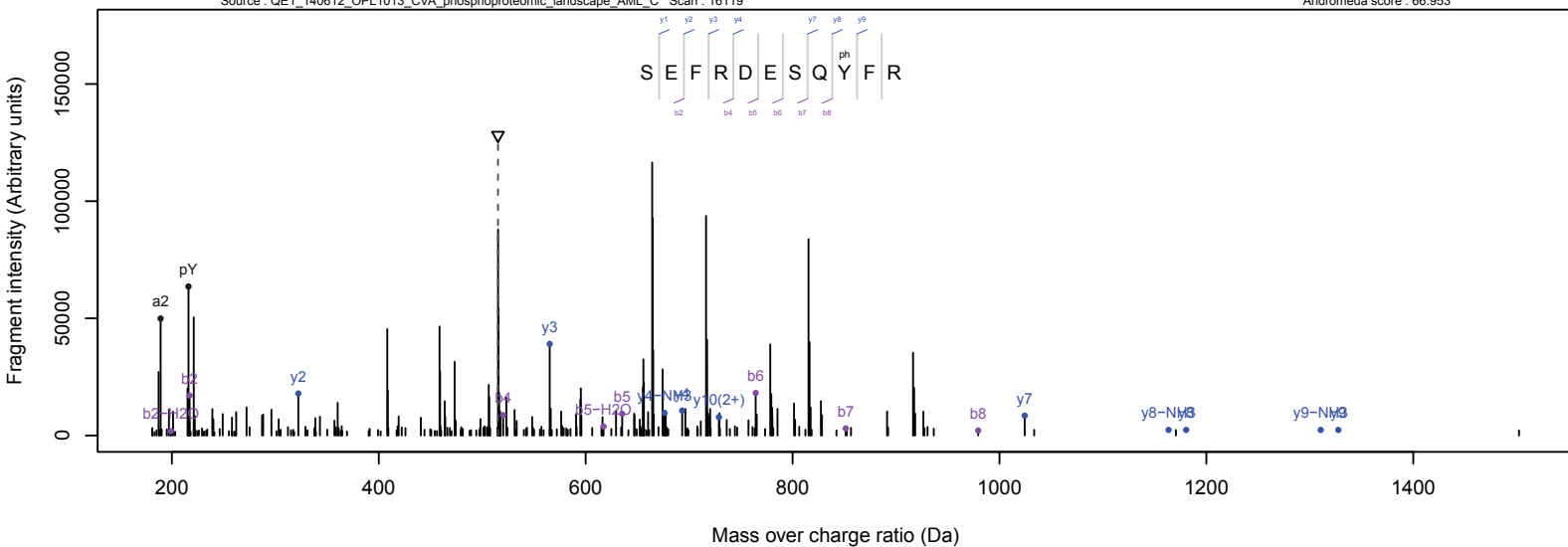
Andromeda score: 66.977



**SEFRDESQ(ph)FR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16119

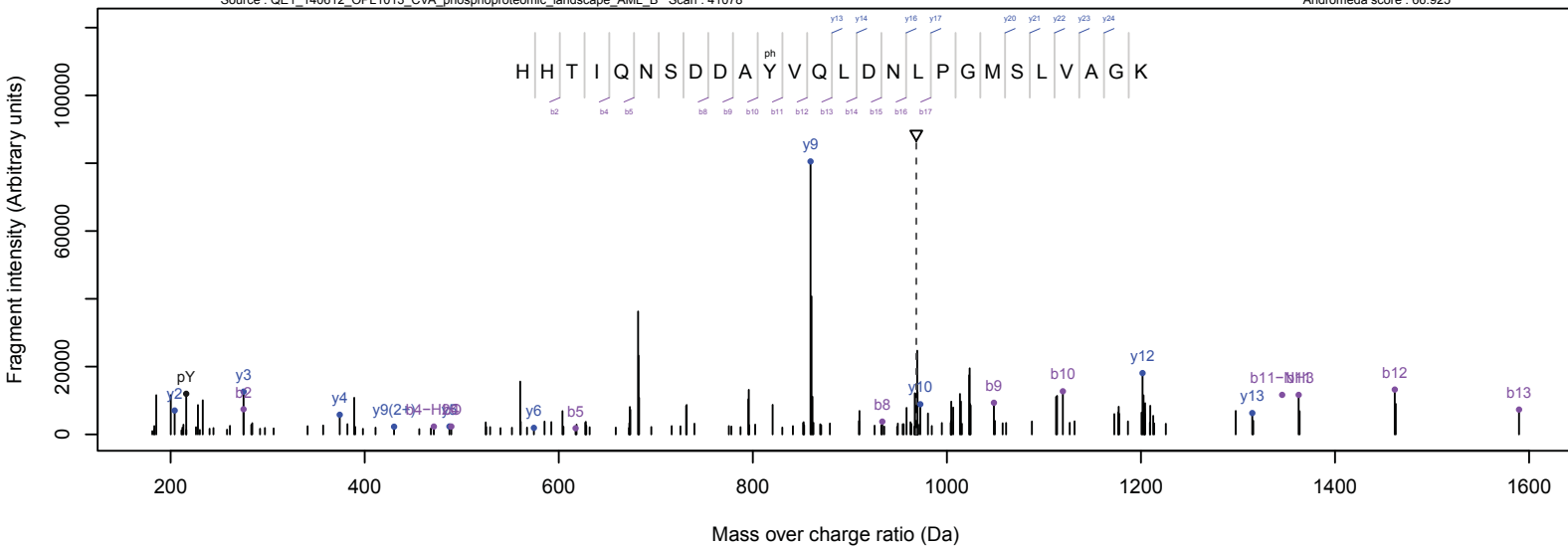
Andromeda score: 66.953



**HHTIQNSDDAY(ph)VQLDNLPGMSLVAGK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 41078

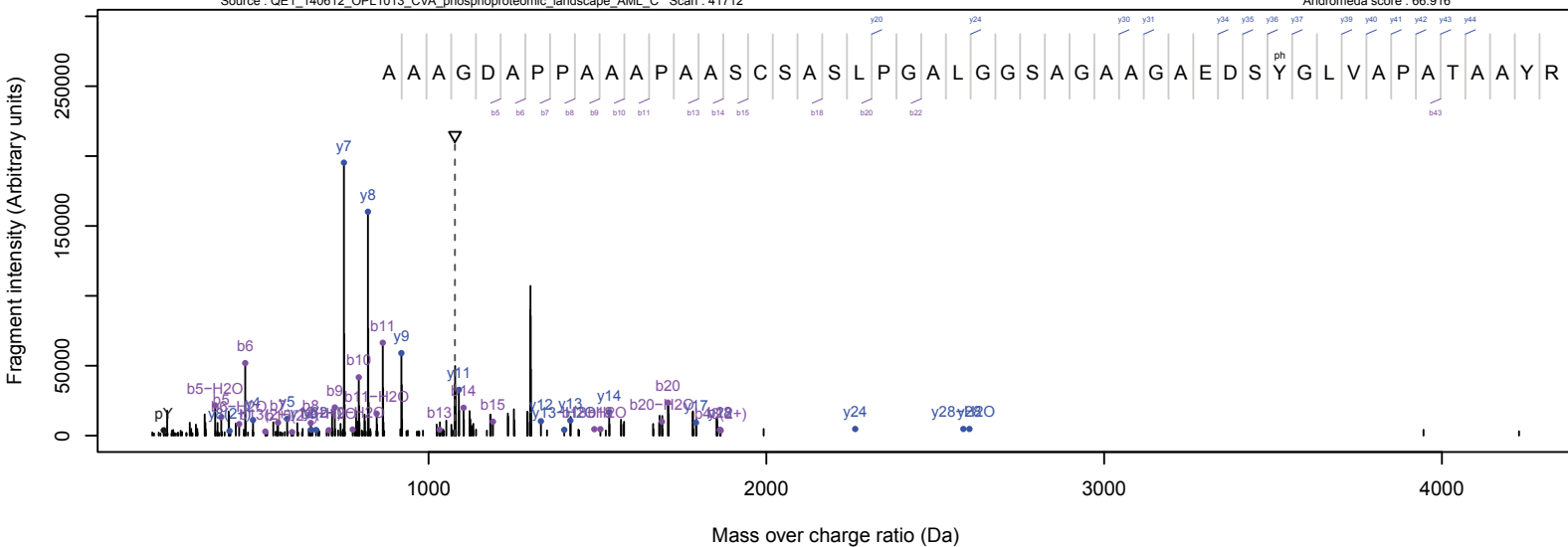
Andromeda score: 66.925



AAAGDAPPAAPAAAPAAASCASLPGALGGSAAGAEDSY(ph)GLVAPATAAYR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41712

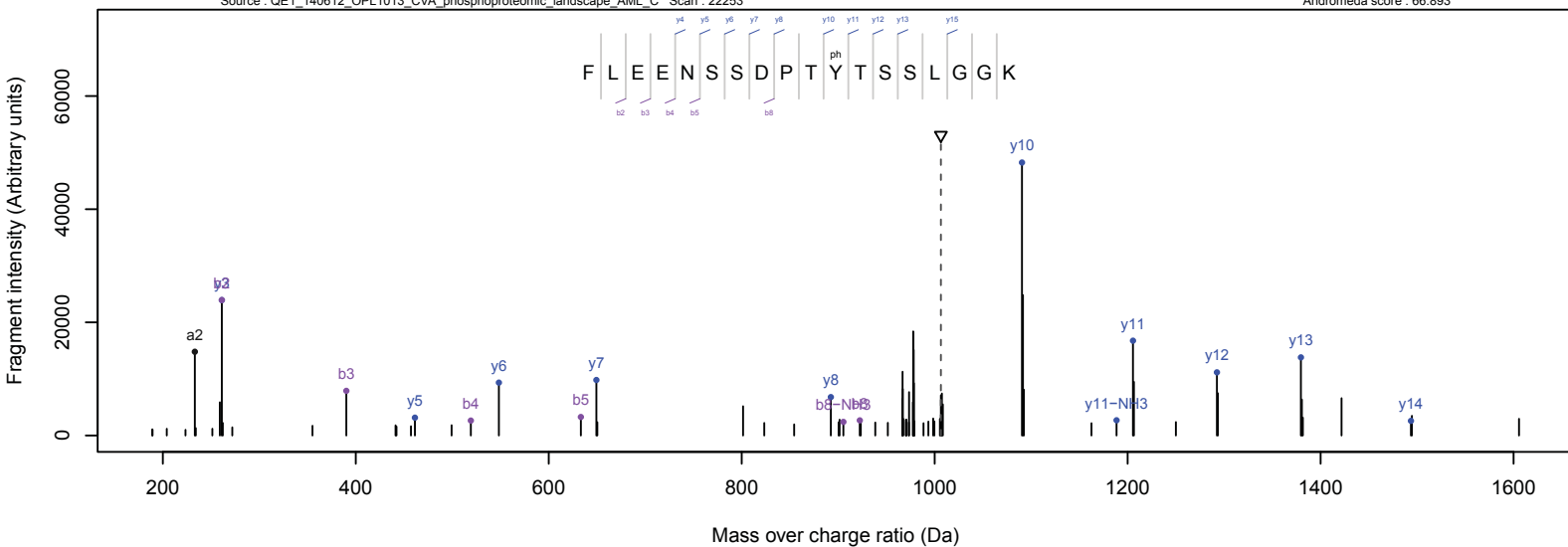
Andromeda score: 66.916



FLEENSSDPTY(ph)TSSLGGK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 22253

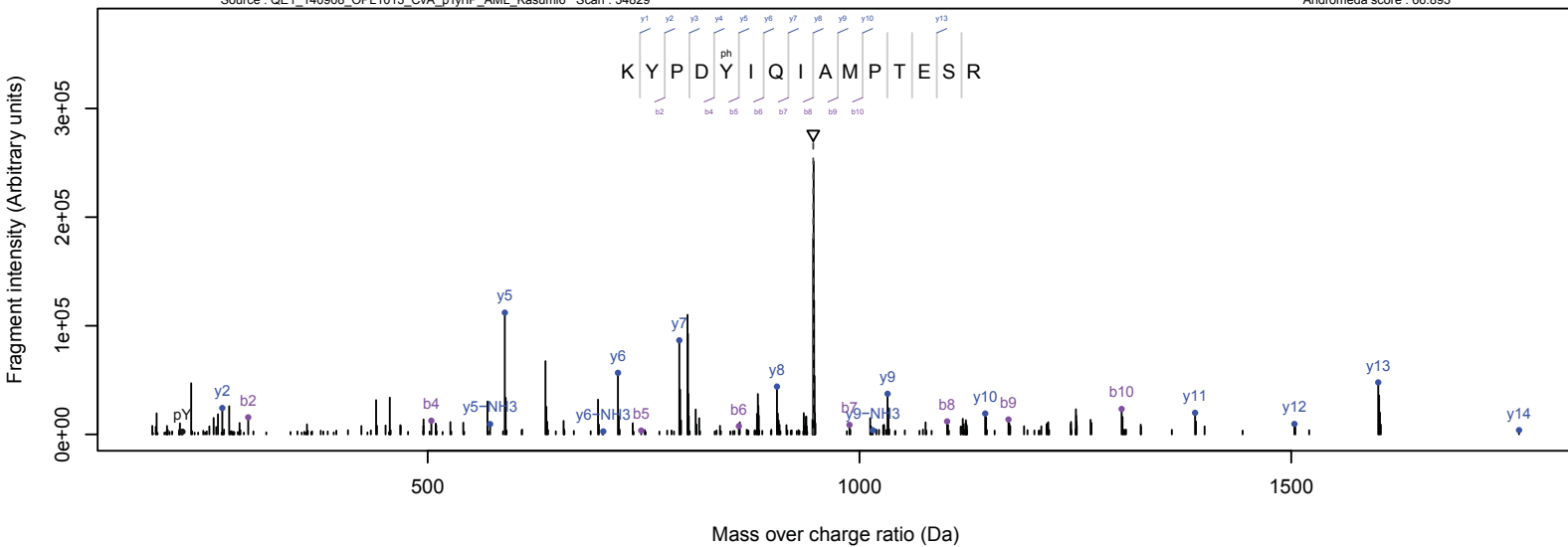
Andromeda score: 66.893



KYPDYIQIAMPTESR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 34829

Andromeda score: 66.893

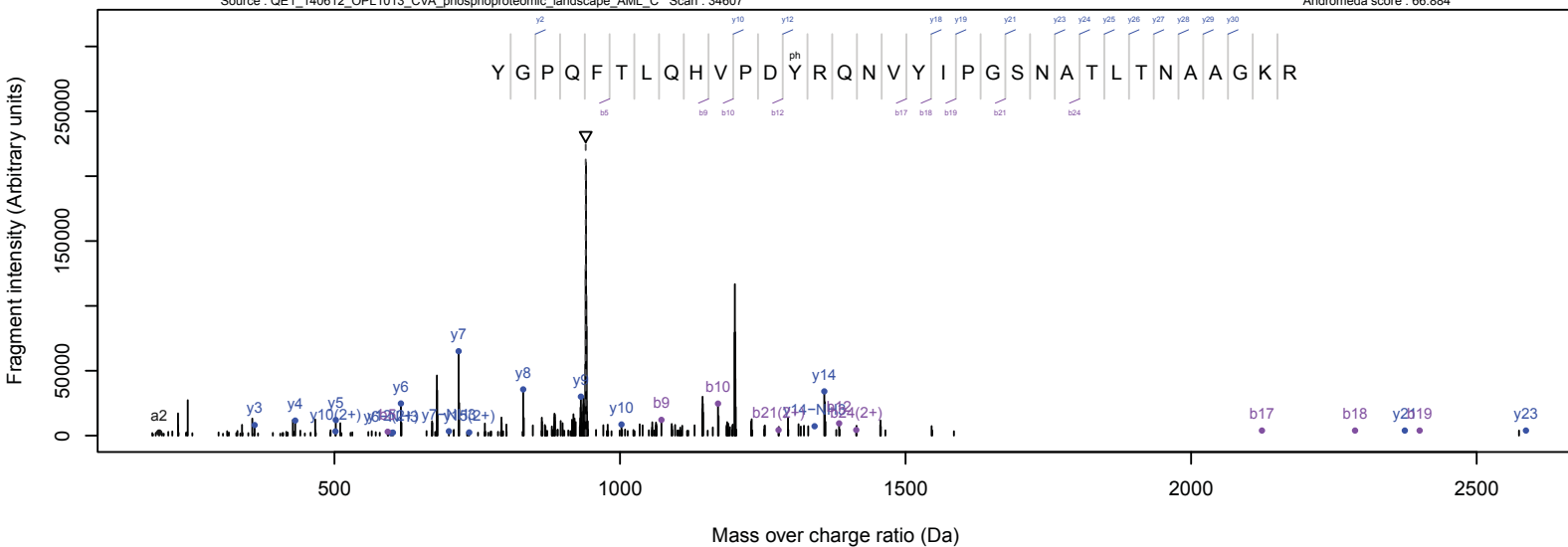




## YGPQFTLQHVPDY(ph)RQNVYIPGSNATLTNAAGKR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34607

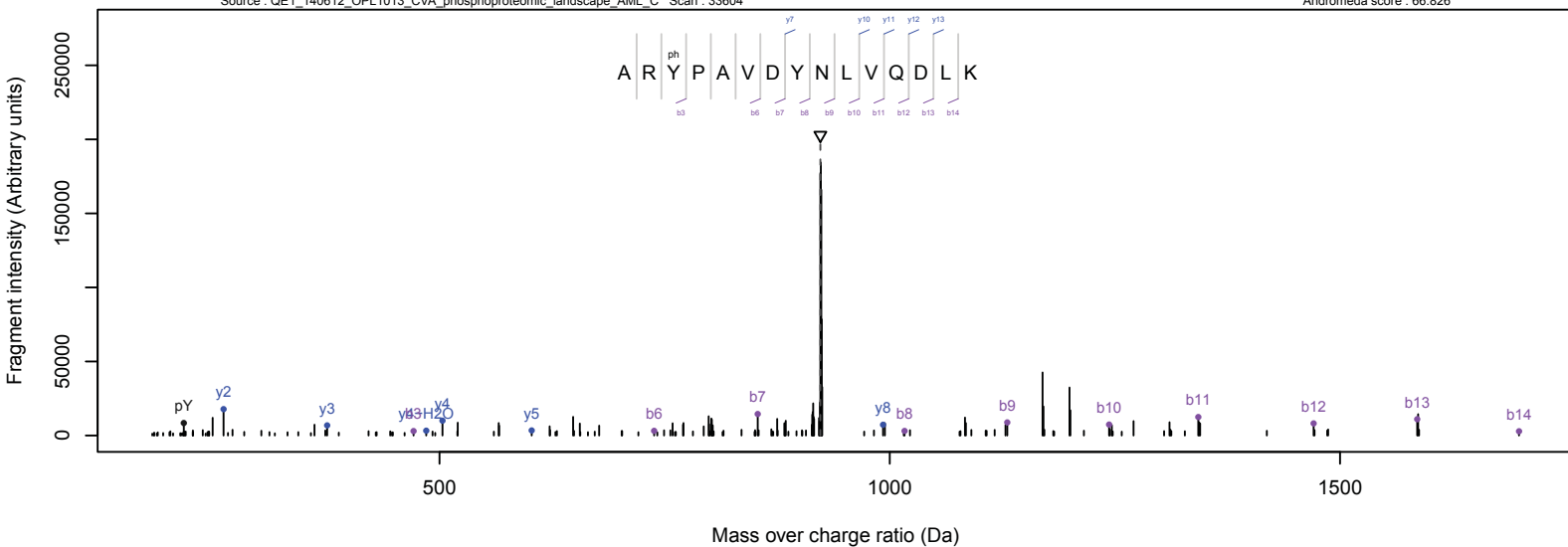
Andromeda score: 66.884



## ARY(ph)PAVDYNLVQDLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 33604

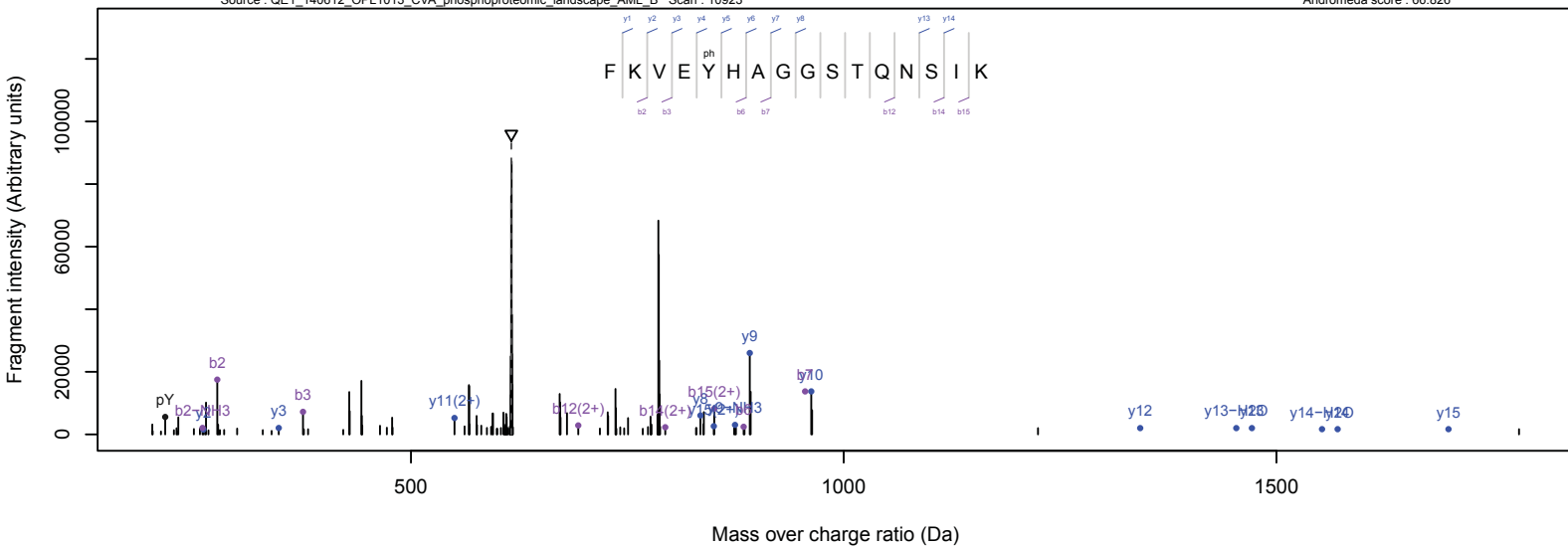
Andromeda score: 66.826



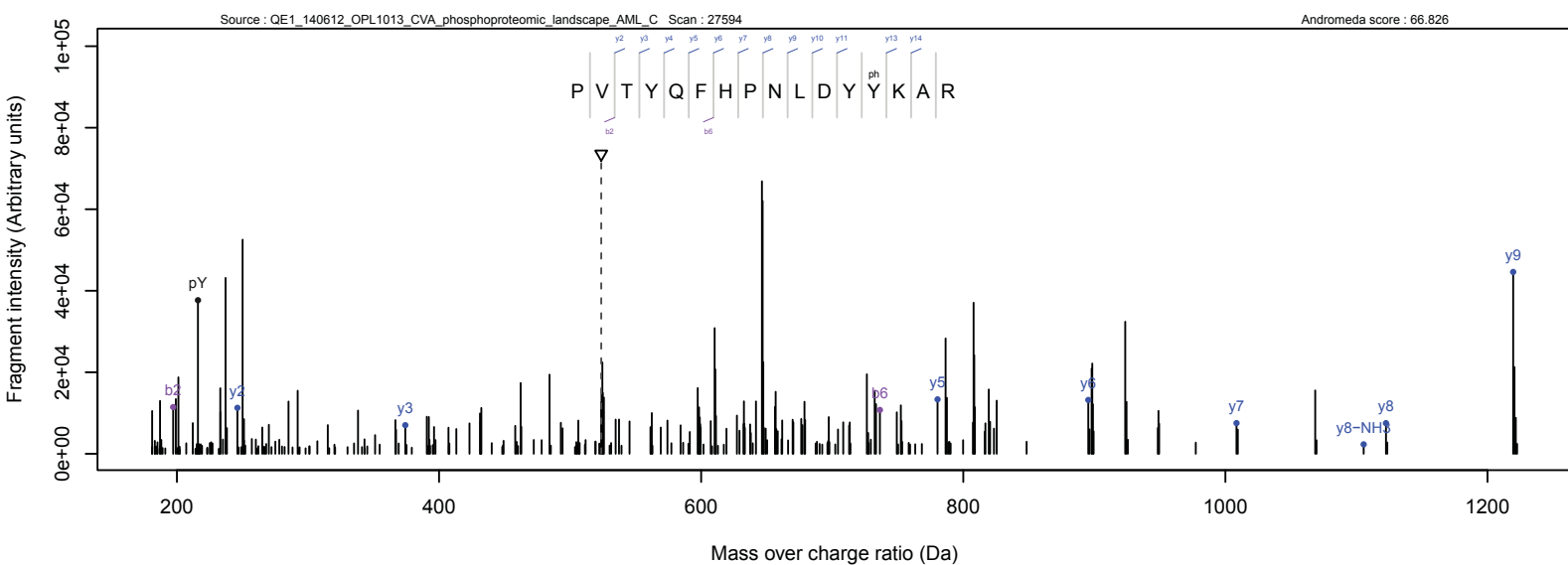
## FKVEY(ph)HAGGSTQNSIK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 10923

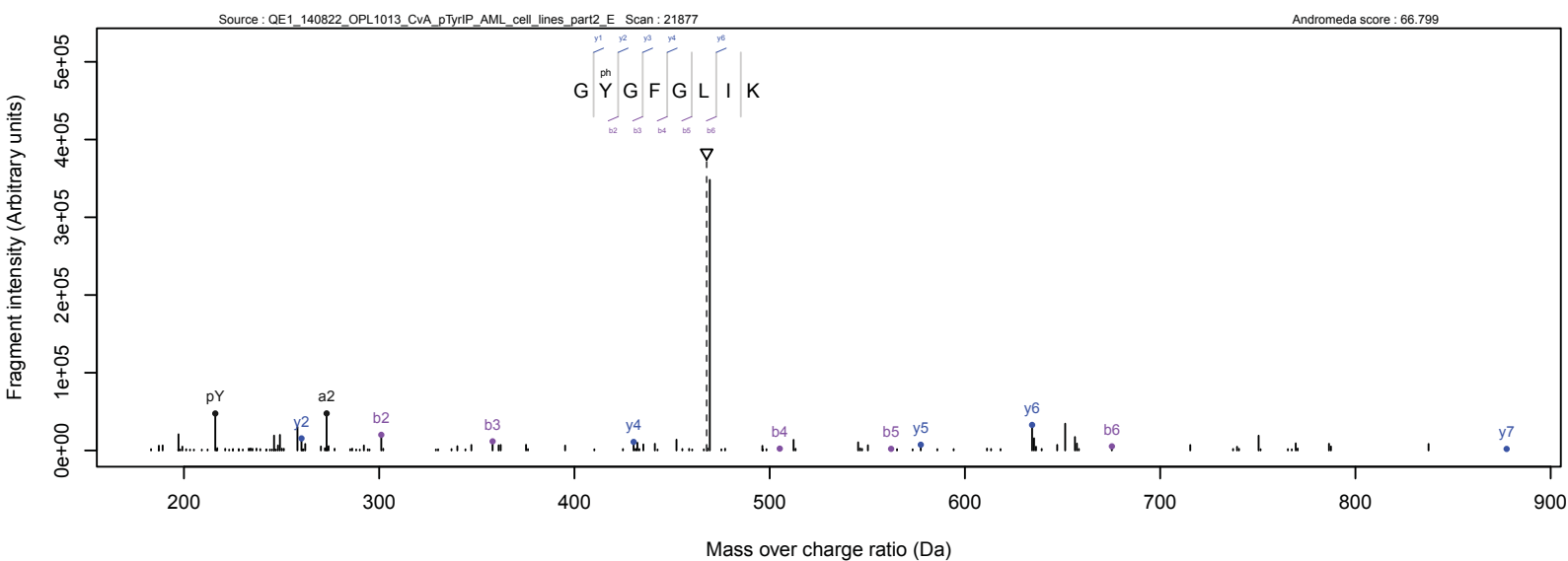
Andromeda score: 66.826



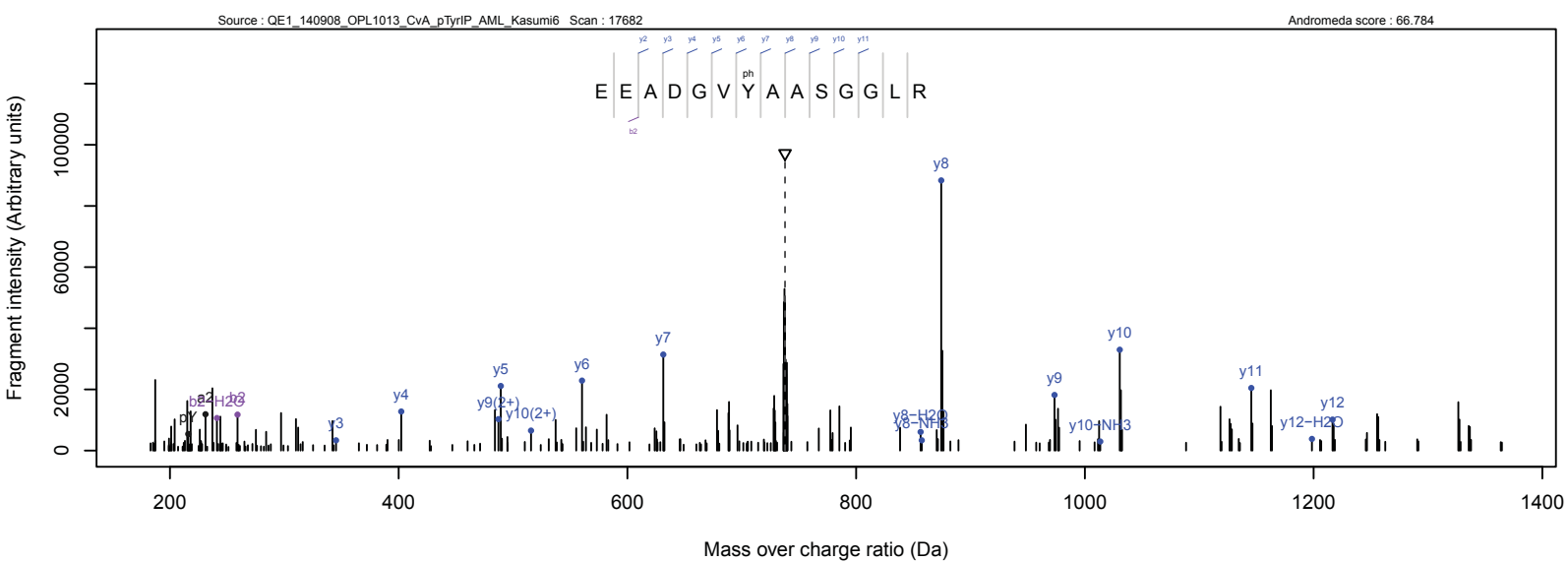
## PVTYQFHPNLDYY(ph)KAR-4+



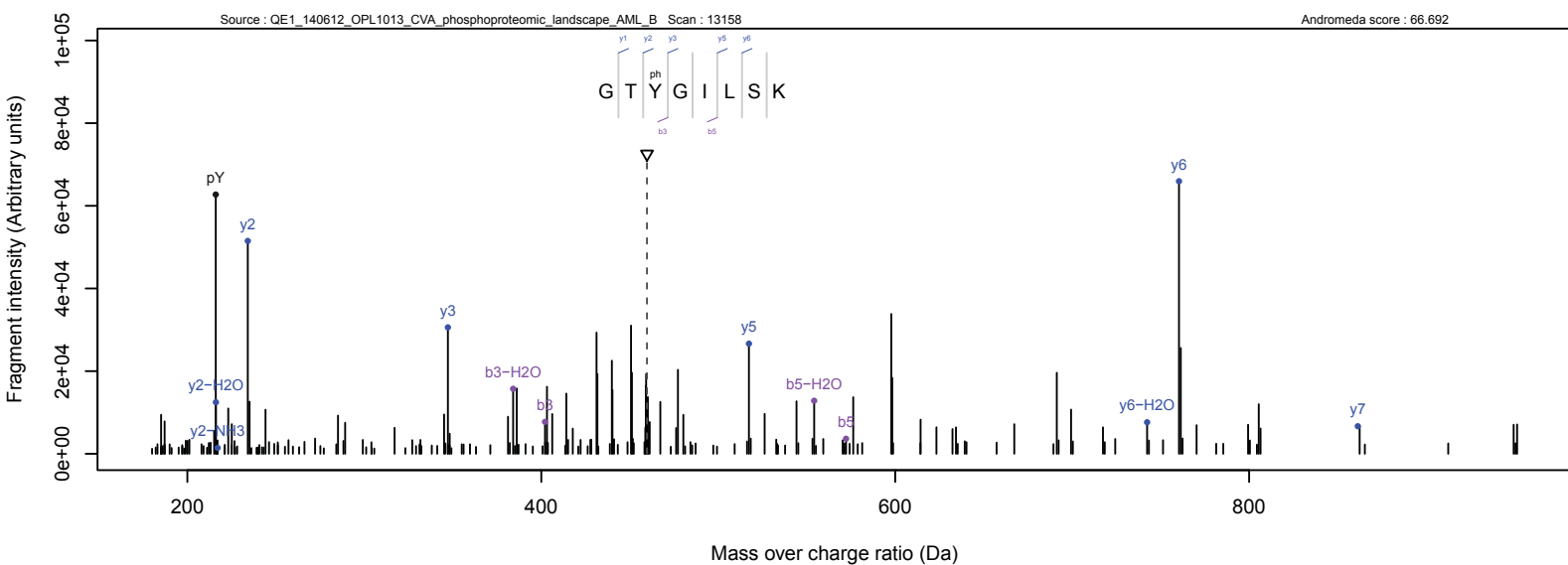
## GY(ph)GFGLIK-2+



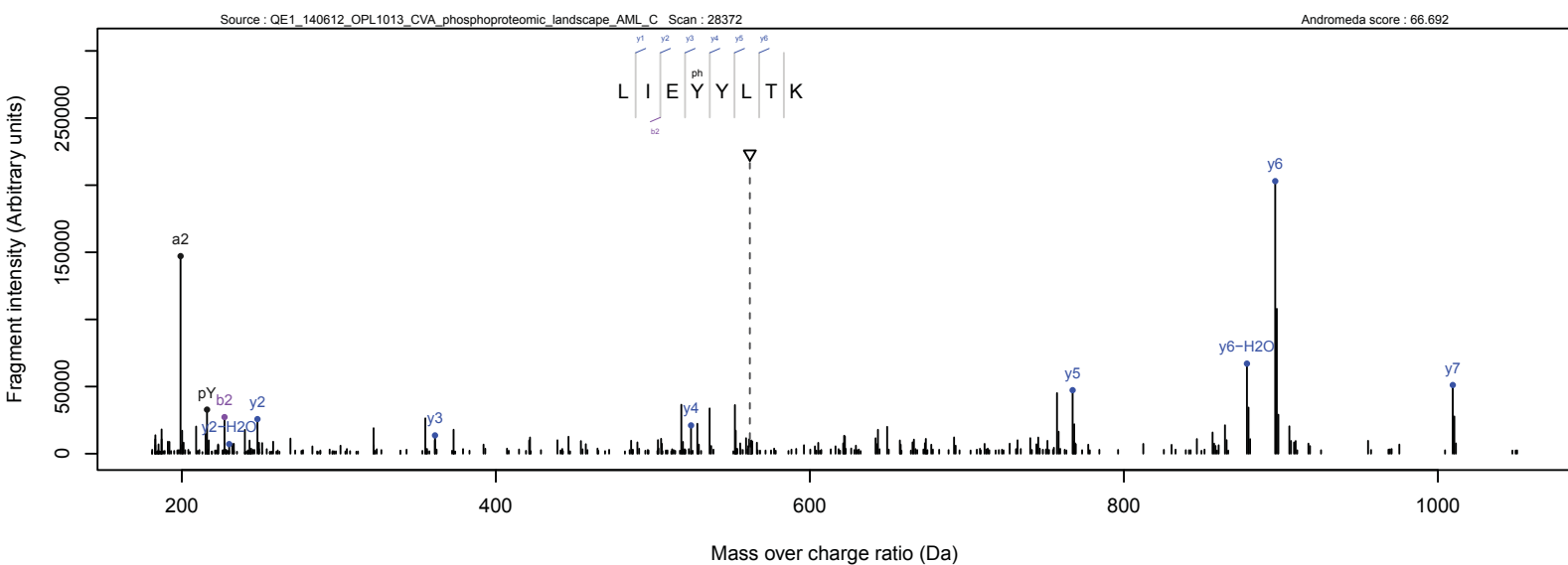
## EEADGVY(ph)AASGLR-2+



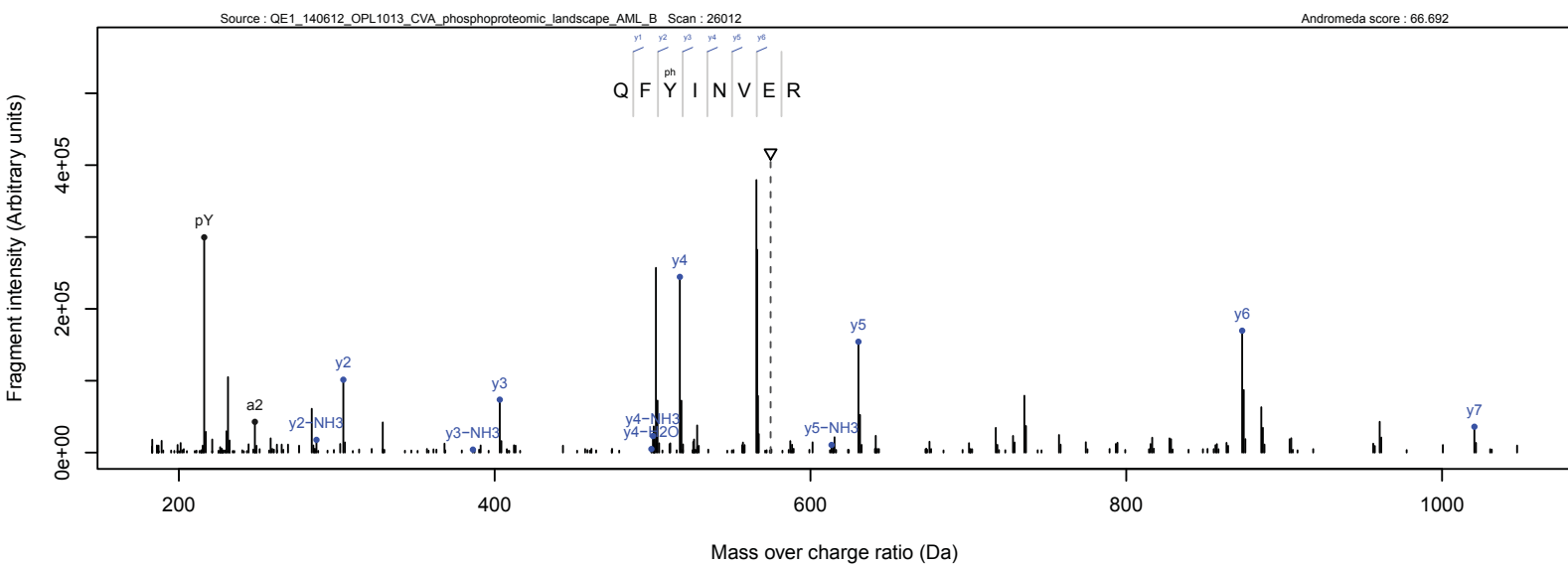
## GTY(ph)GILSK-2+



## LIEY(ph)YLTK-2+



## QFY(ph)INVER-2+

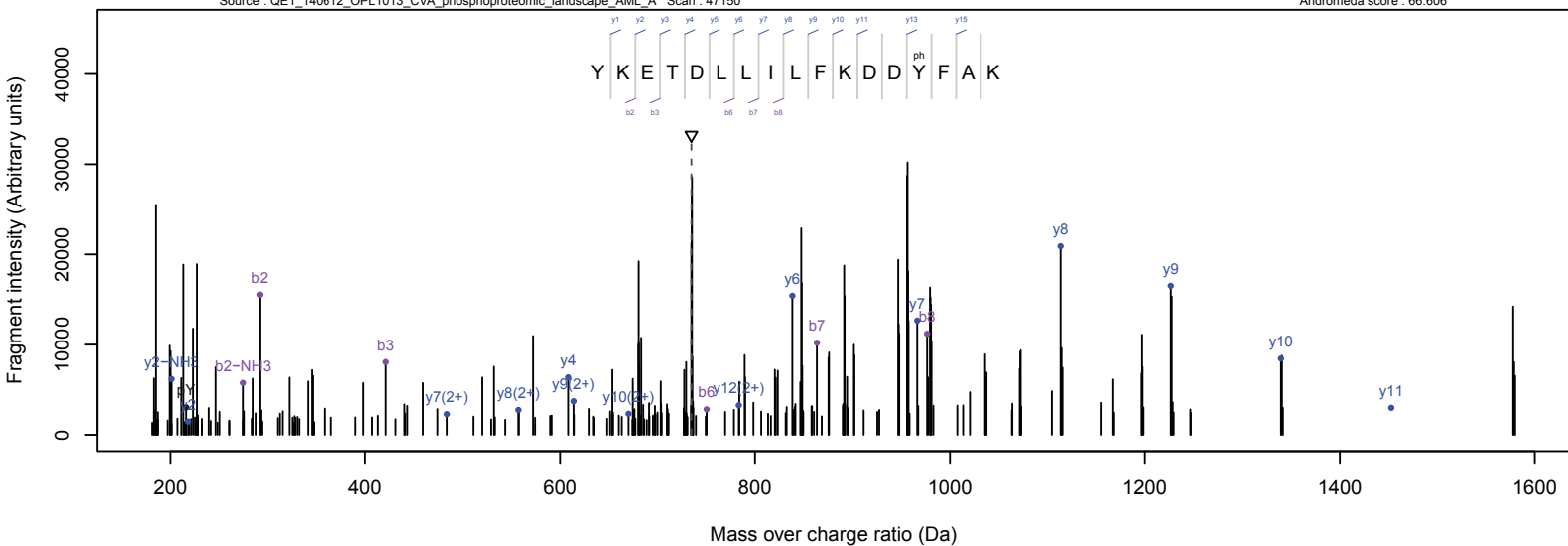




## YKETDLLILFKDDY(ph)FAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 47150

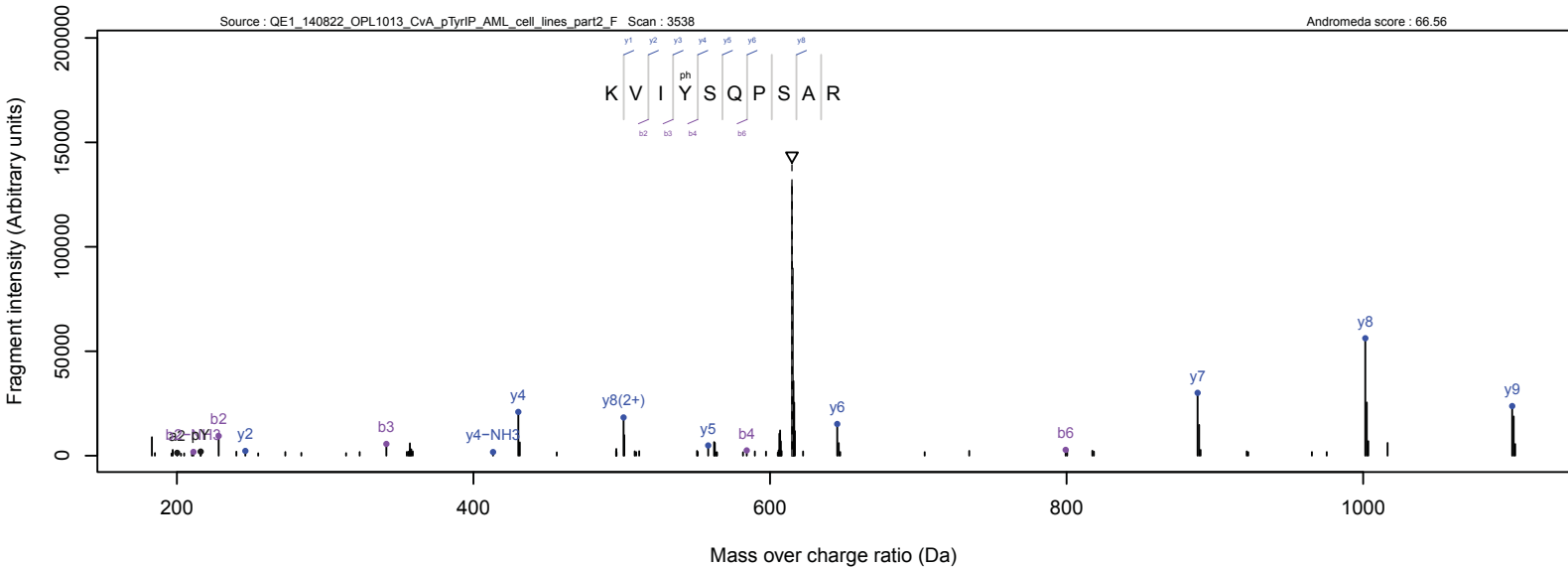
Andromeda score : 66.606



## KVIY(ph)SQPSAR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 3538

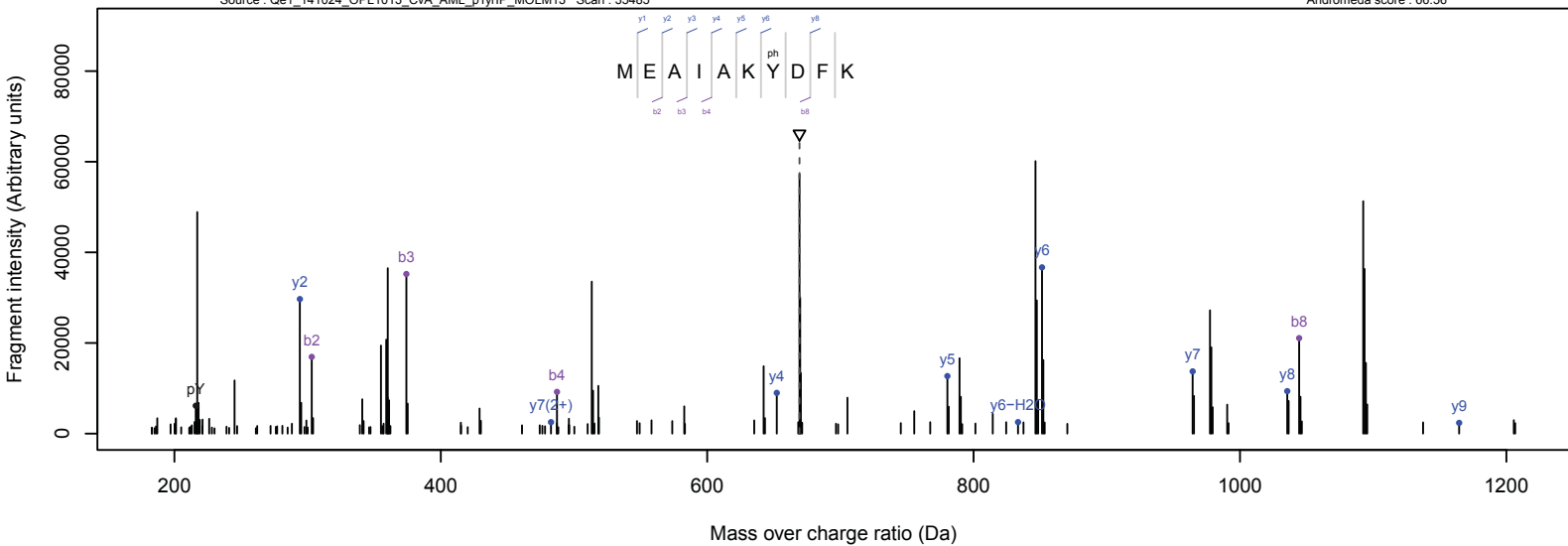
Andromeda score : 66.56



## (ac)MEAIKY(ph)DFK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 35485

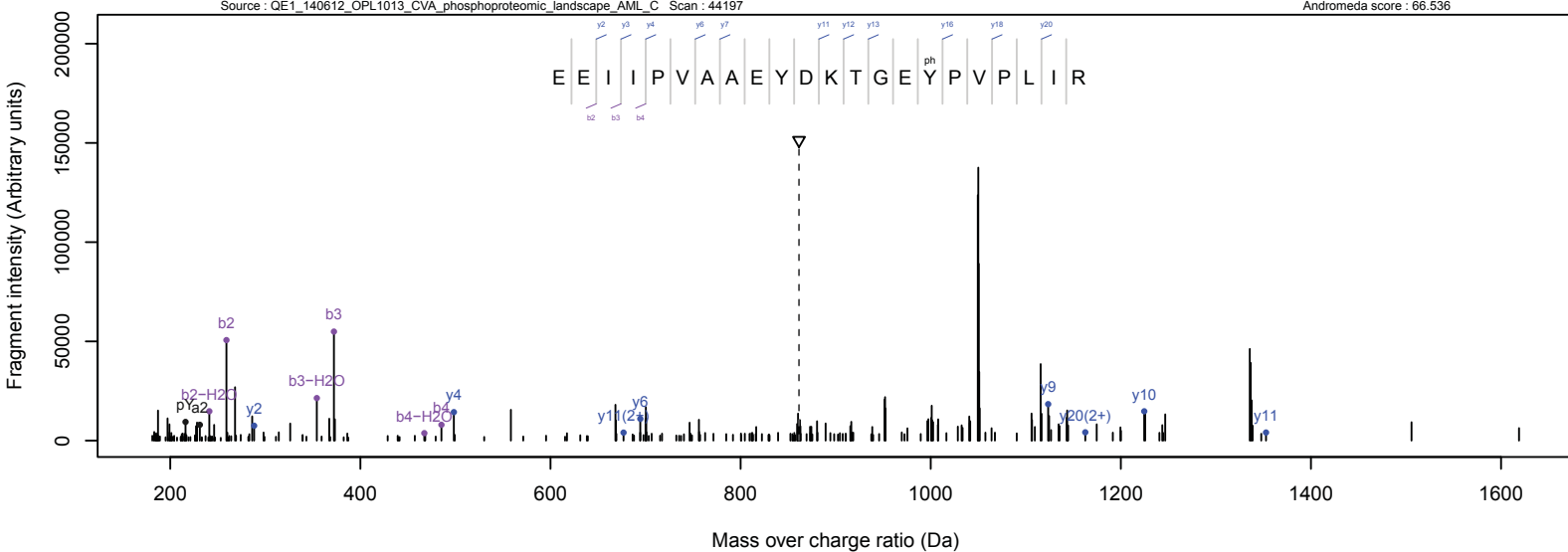
Andromeda score : 66.56



## EEIIPVAAEYDKTGEY(ph)PVPLIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44197

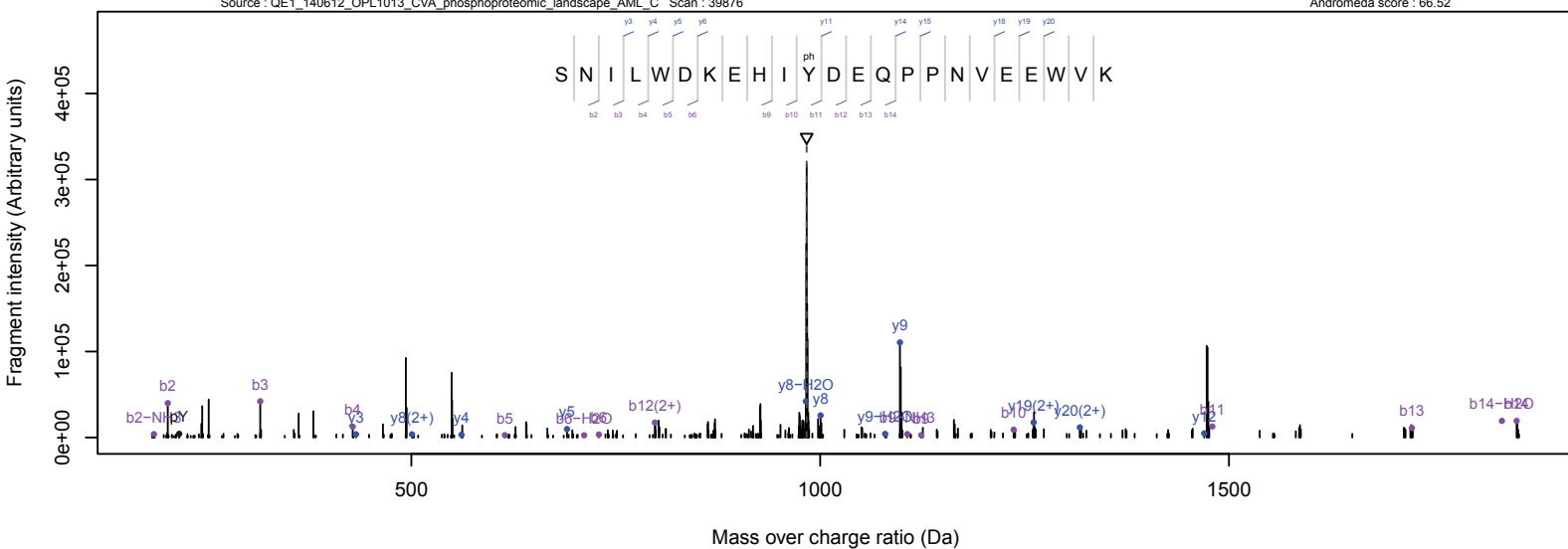
Andromeda score : 66.536



## SNILWDKEHIY(ph)DEQPPNVEEWVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39876

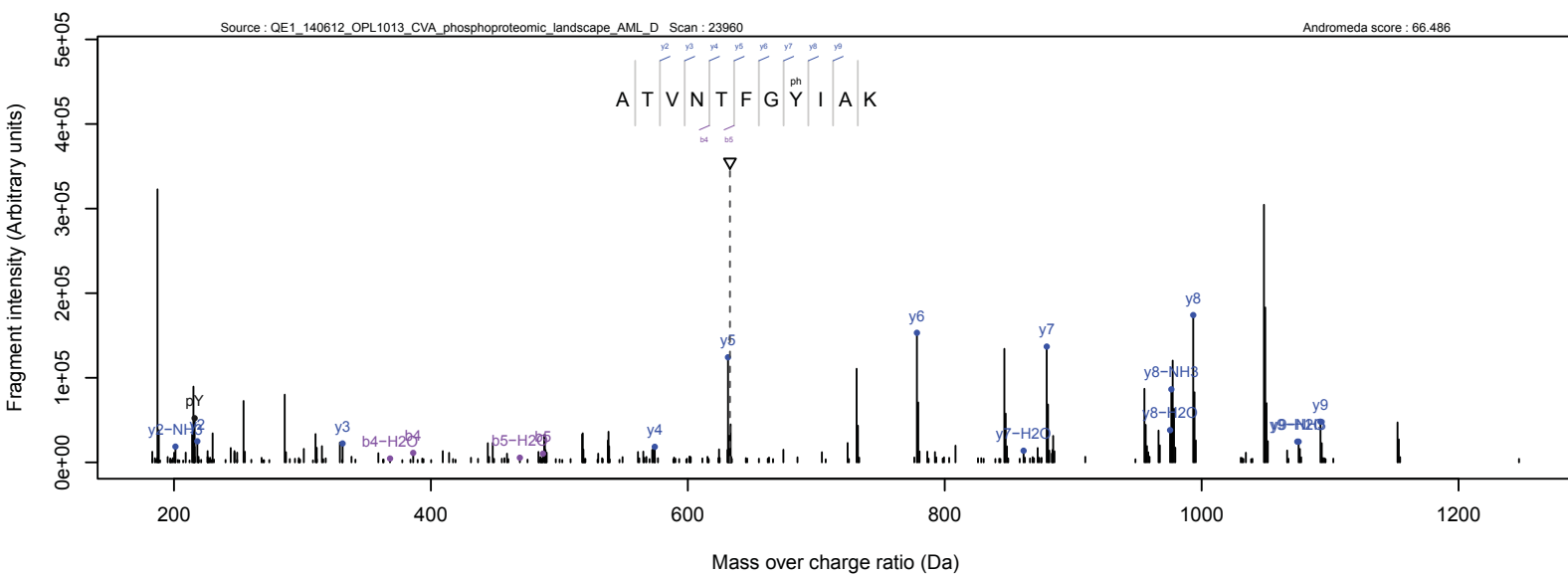
Andromeda score : 66.52



## ATVNTFGY(ph)IAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23960

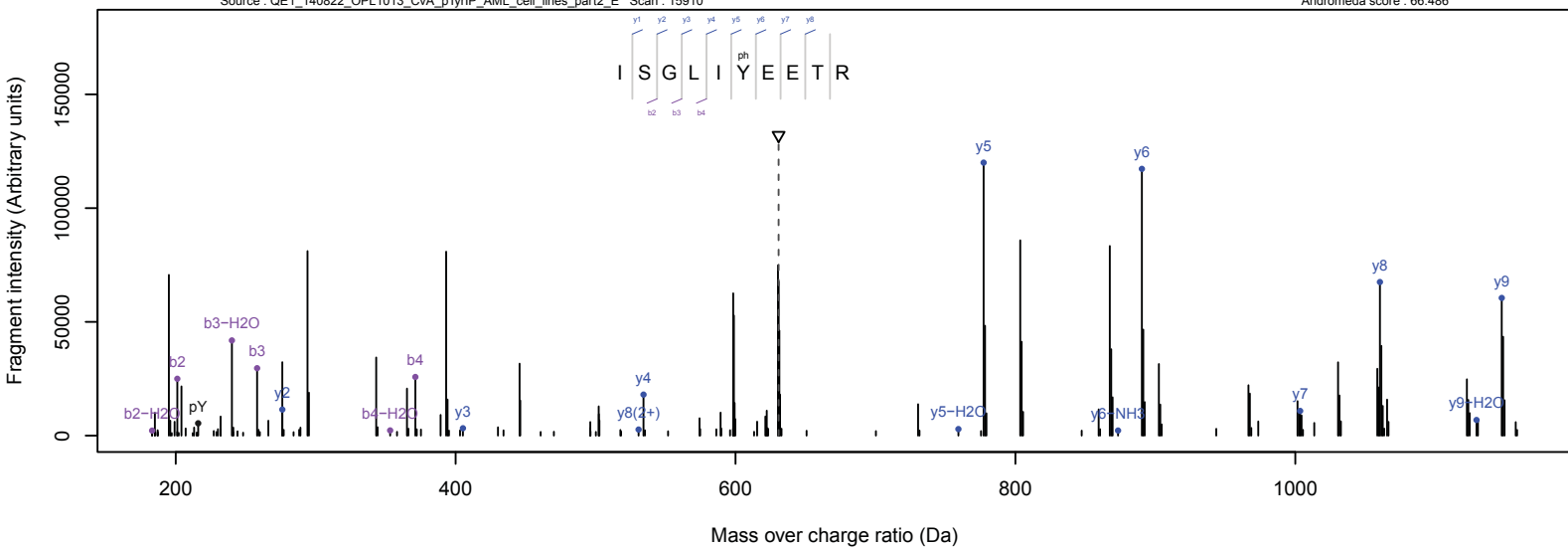
Andromeda score : 66.486



ISGLIY(ph)EETR-2+

Source: QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 15910

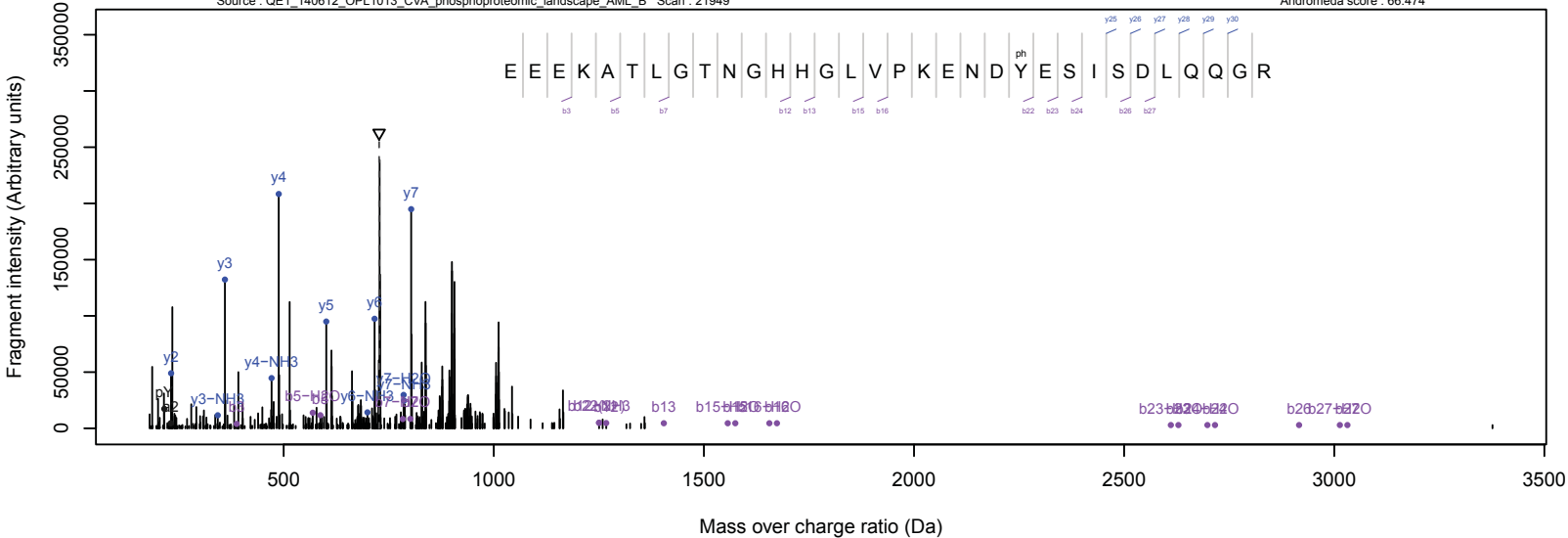
Andromeda score: 66.486



EEEKATLGTNGHHGLVLPKENDY(ph)ESIDLQQGR-5+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 21949

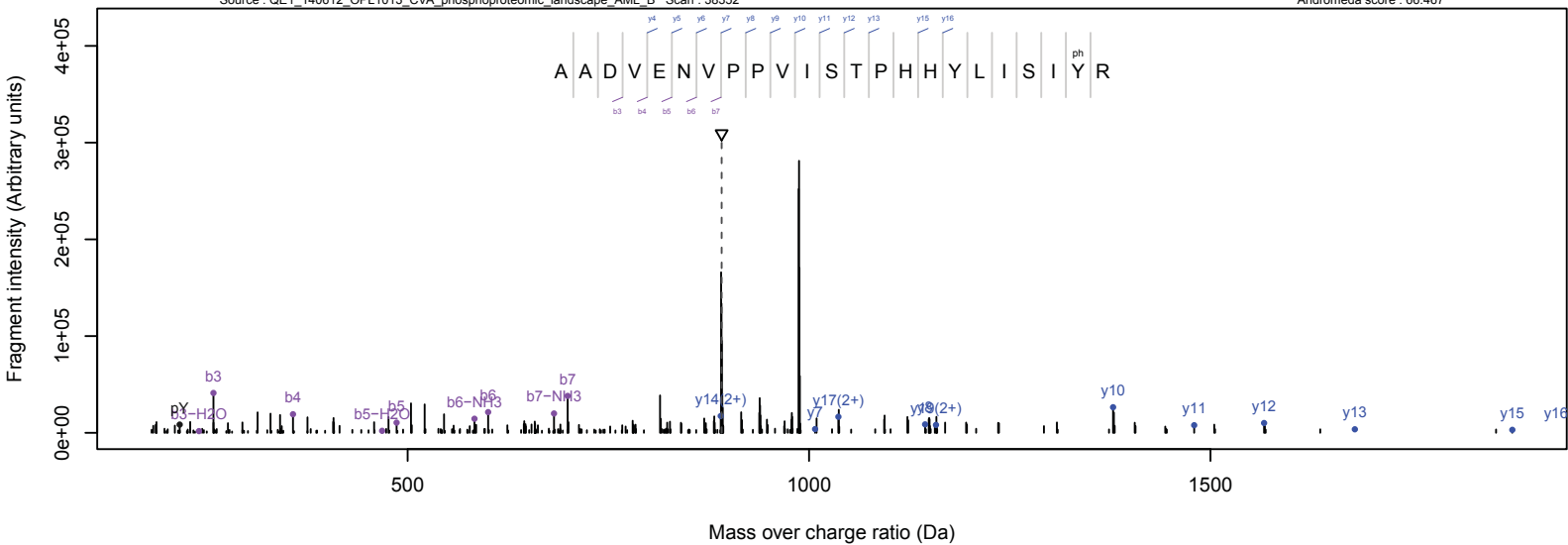
Andromeda score: 66.474



AADVENVPPVISTPHHYLISYIYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 38352

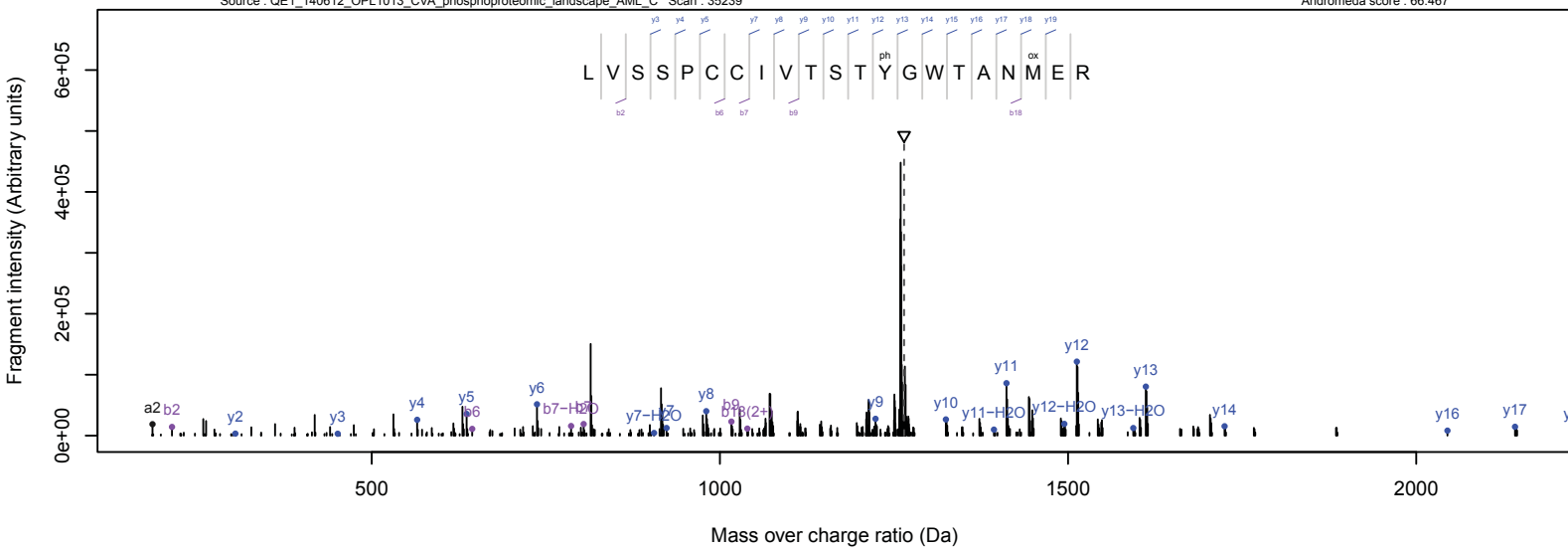
Andromeda score: 66.467



LVSSPCCIVTSTY<sup>ph</sup>GWTANM<sup>ox</sup>ER-2+

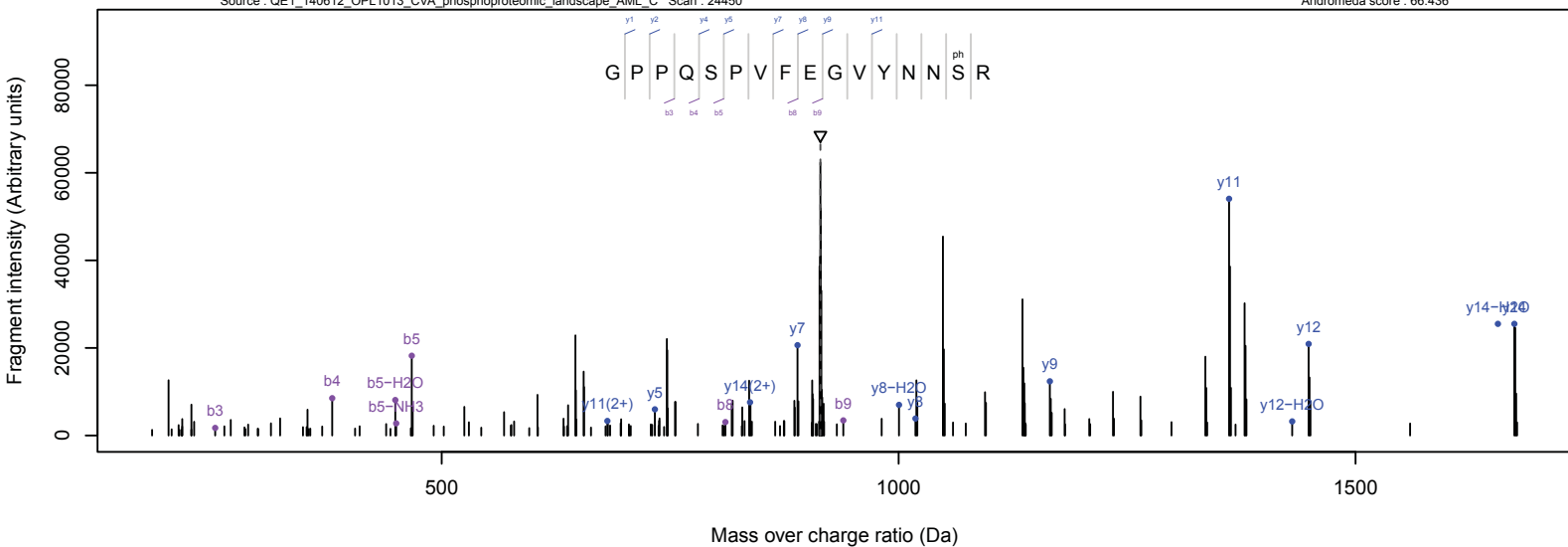
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35239

Andromeda score : 66.467

GPPQSPVFE<sup>ph</sup>GVYNN<sup>ph</sup>SR

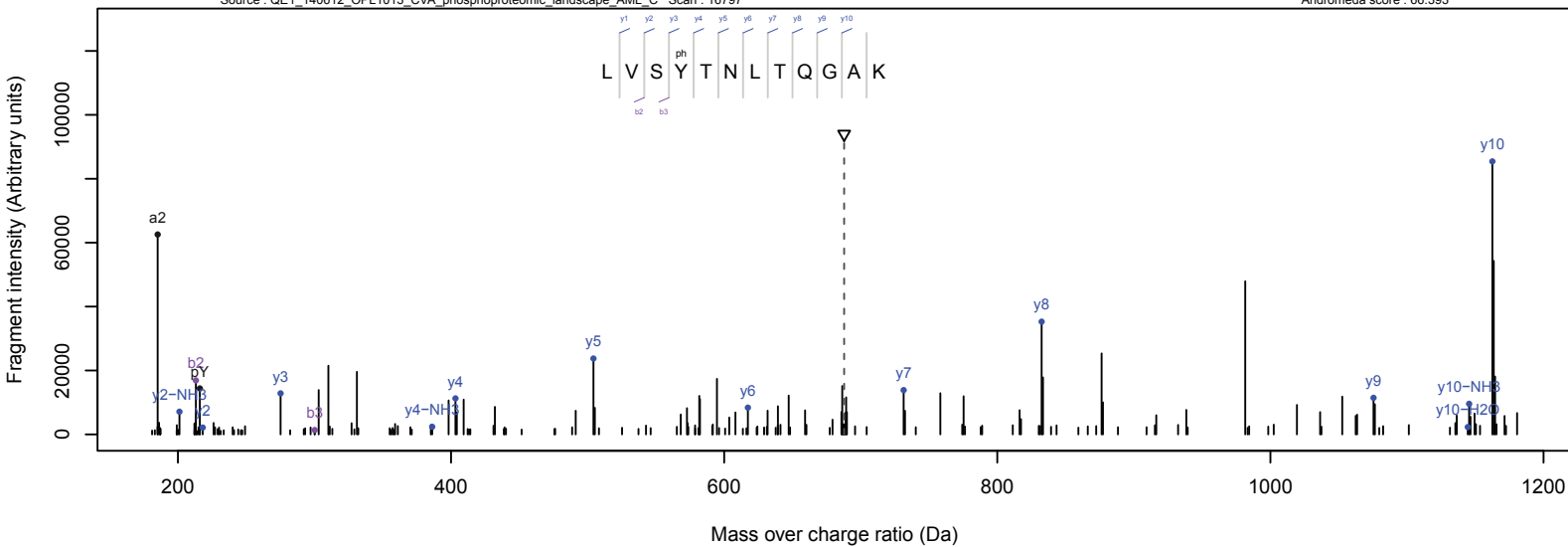
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24450

Andromeda score : 66.436

LVSY<sup>ph</sup>TNLTQGA<sup>ph</sup>K

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16797

Andromeda score : 66.393

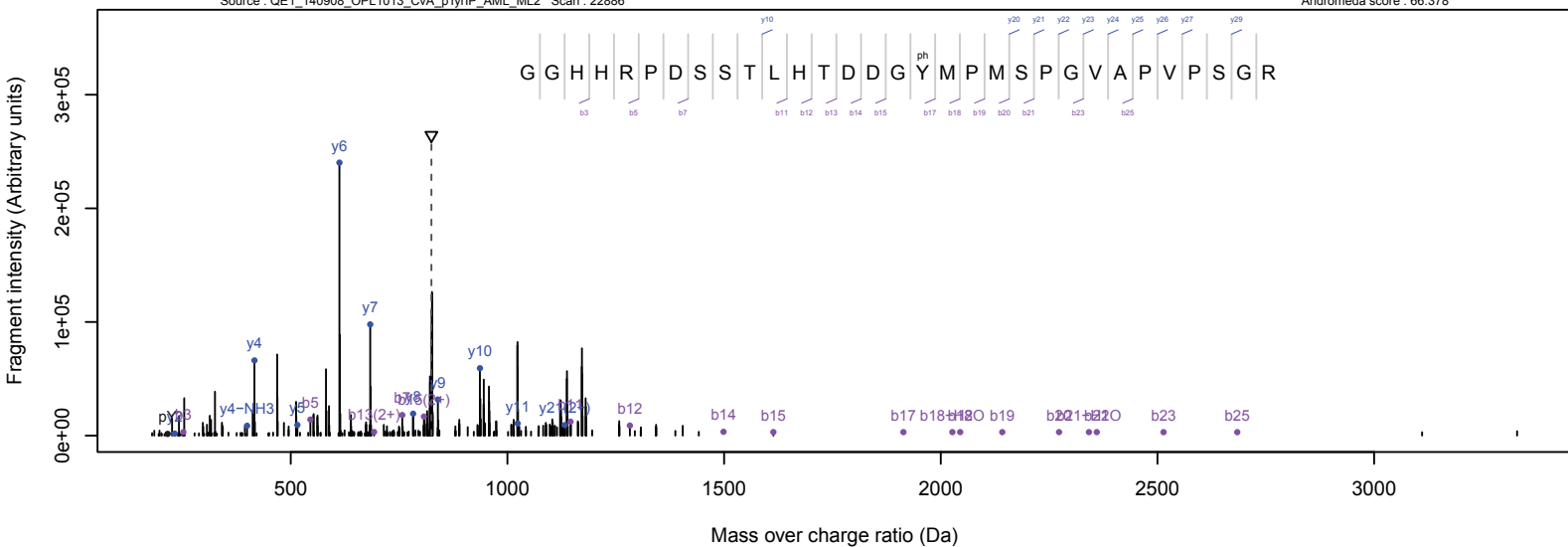




GGHHRPDSSTLHTDDGY(ph)MPMPGVPVPSGR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 22886

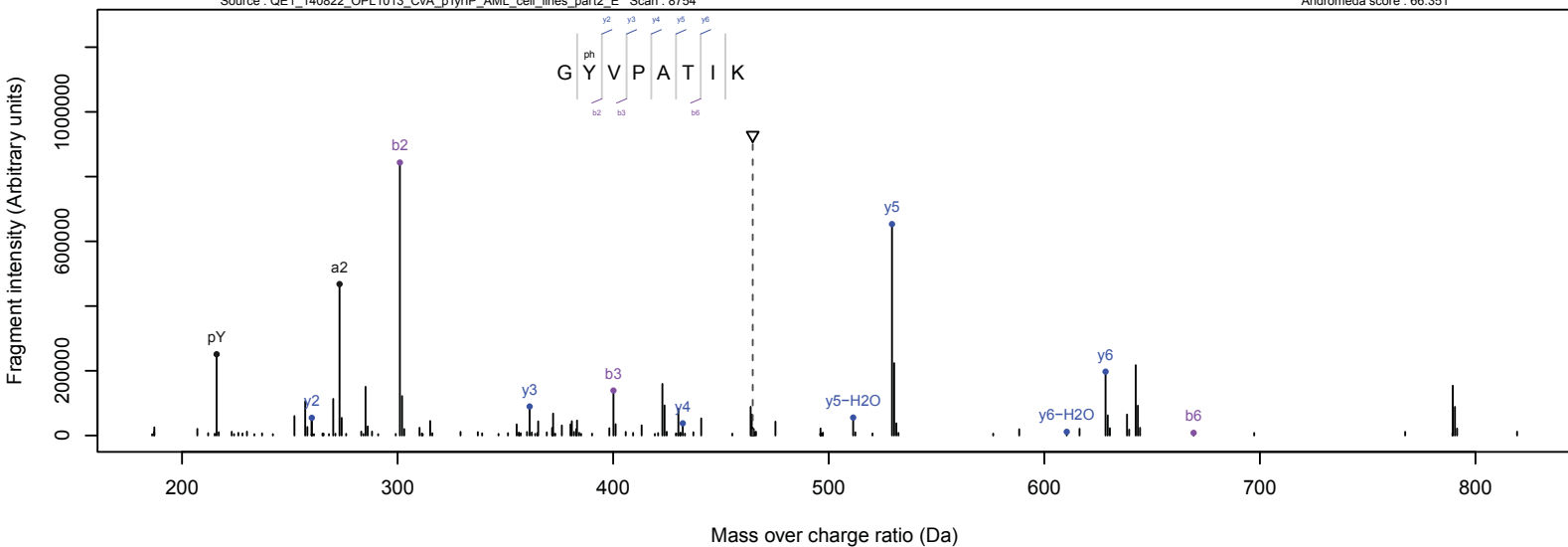
Andromeda score : 66.378



GY(ph)VPATIK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 8754

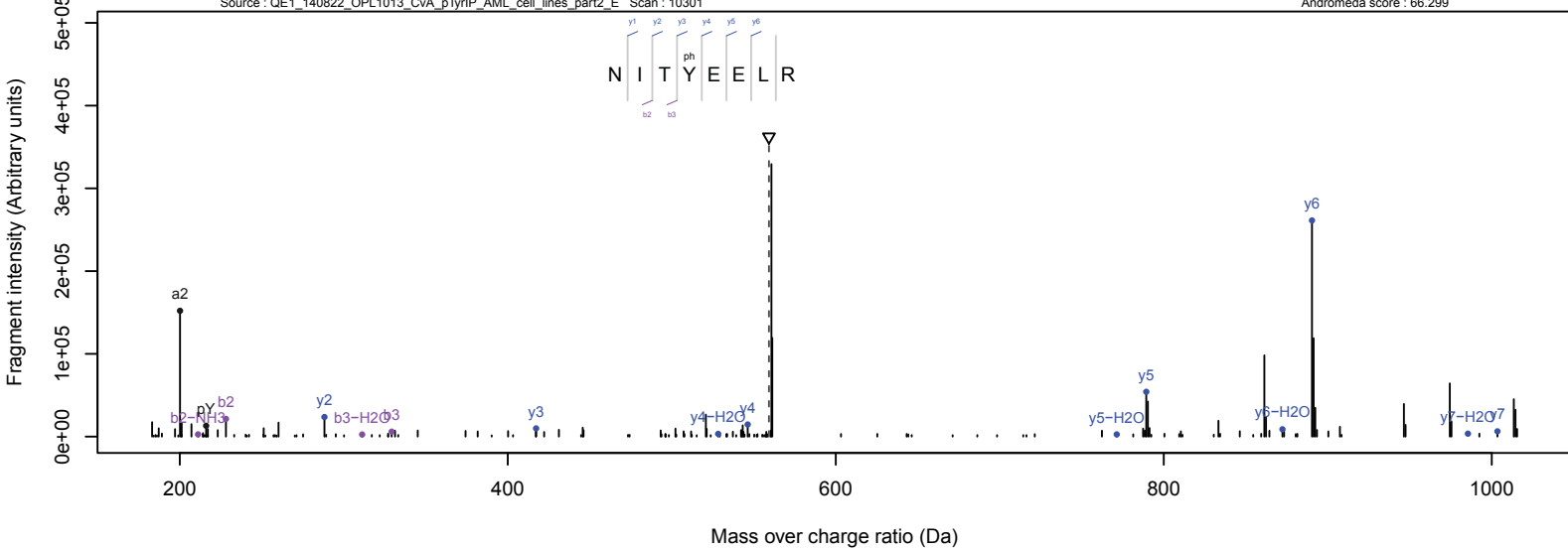
Andromeda score : 66.351



NITY(ph)EELR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 10301

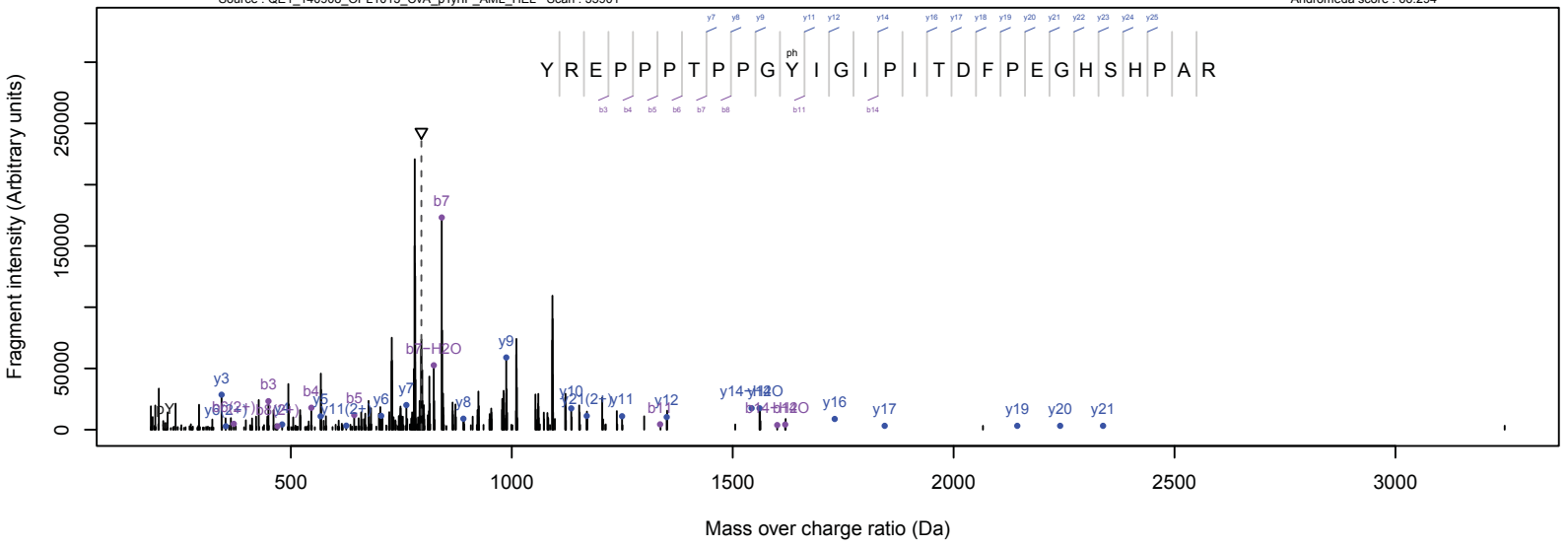
Andromeda score : 66.299



YREPPPTPPG(ph)IGIPITDFEGHSHPAR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35901

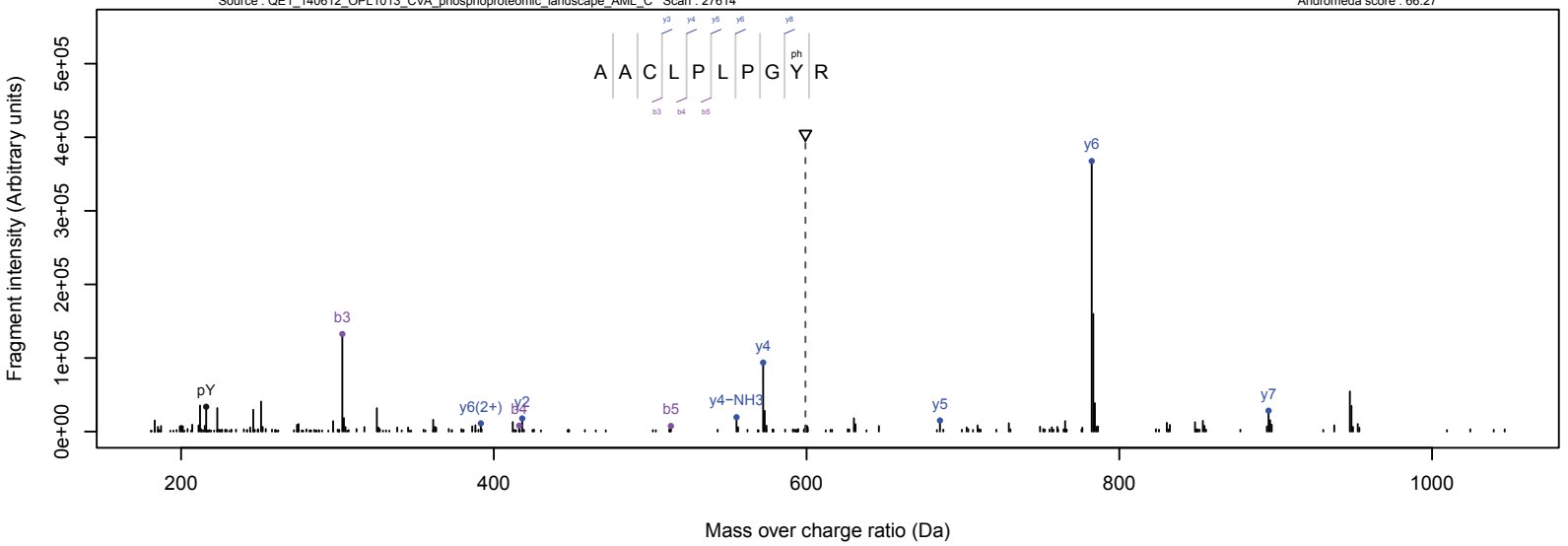
Andromeda score : 66.294



AACLPLPGY(ph)R-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 27614

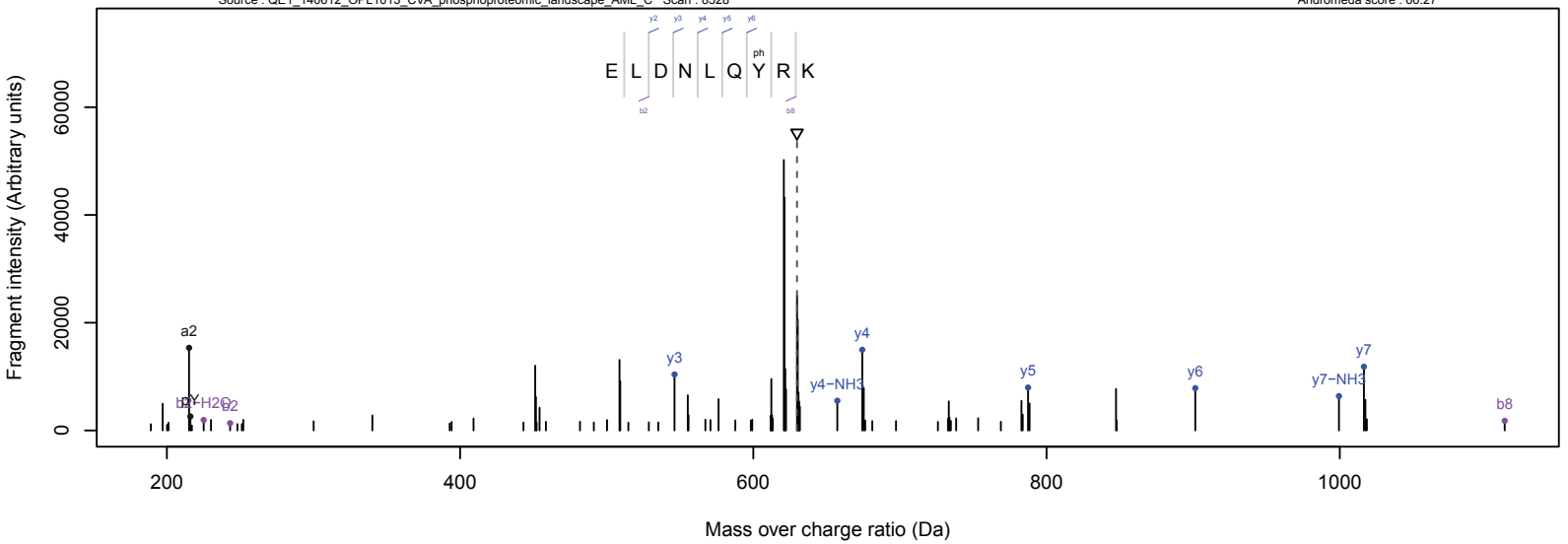
Andromeda score : 66.27



ELDNLQY(ph)RK-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 8528

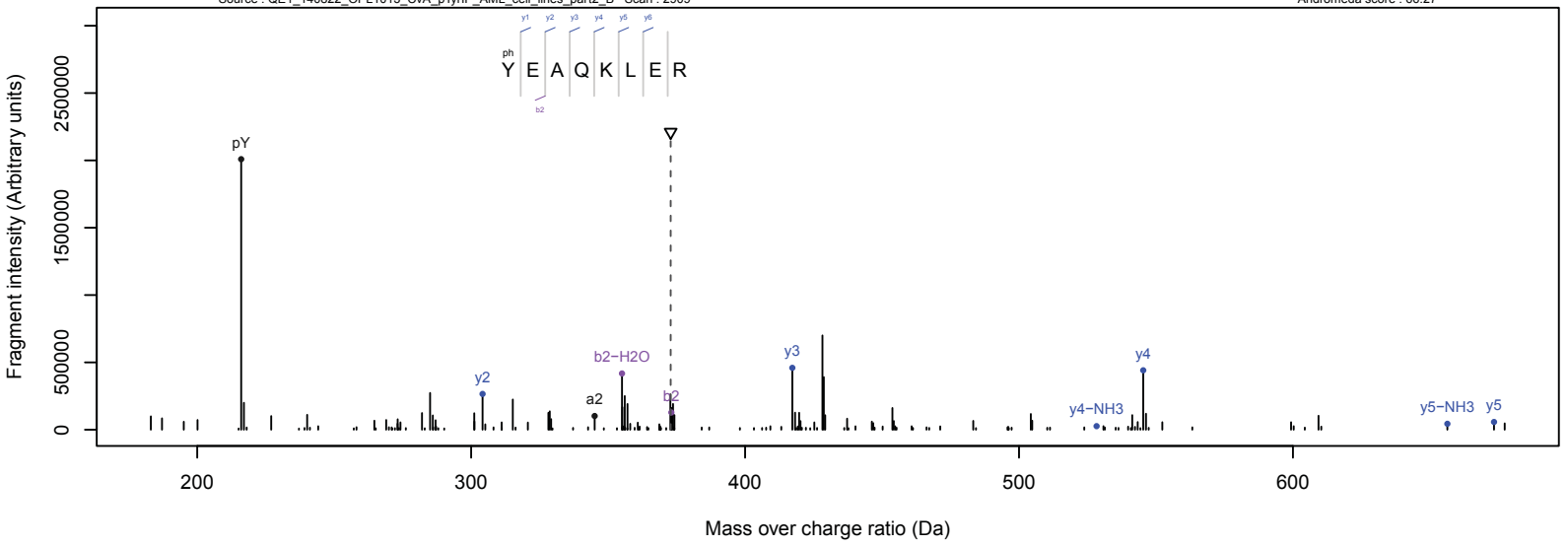
Andromeda score : 66.27



Y(ph)EAQKLER-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 2909

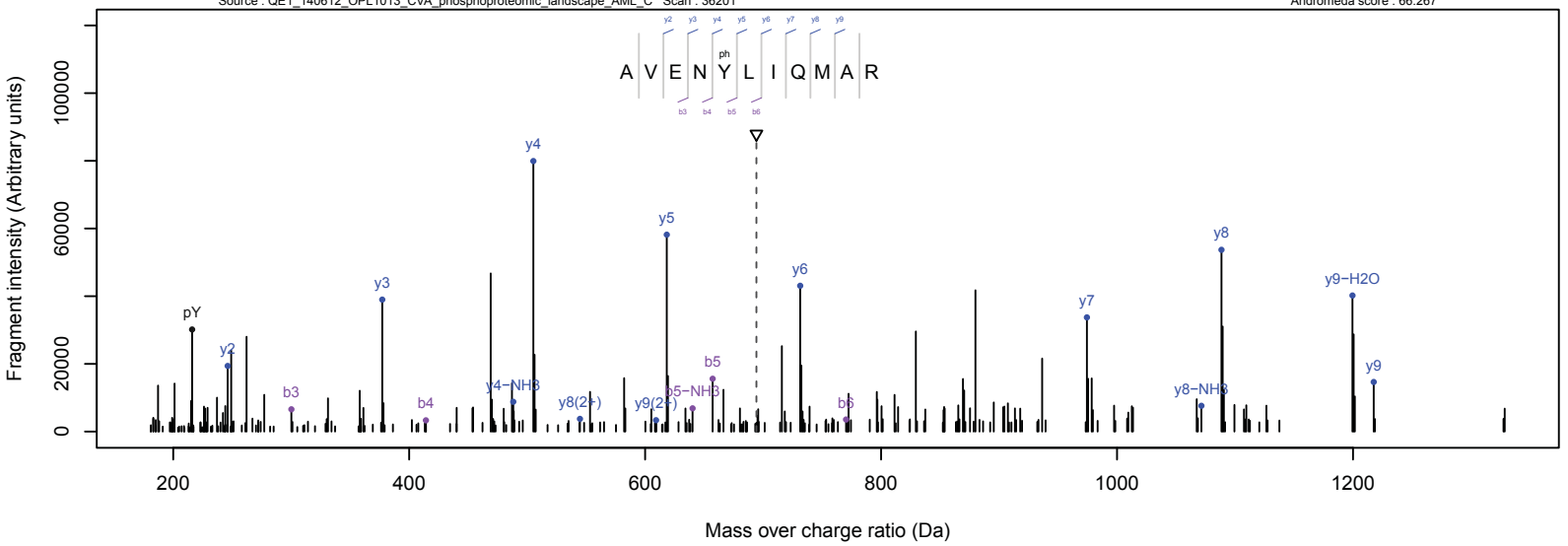
Andromeda score : 66.27



AVENY(ph)LIQMAR-2+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 36201

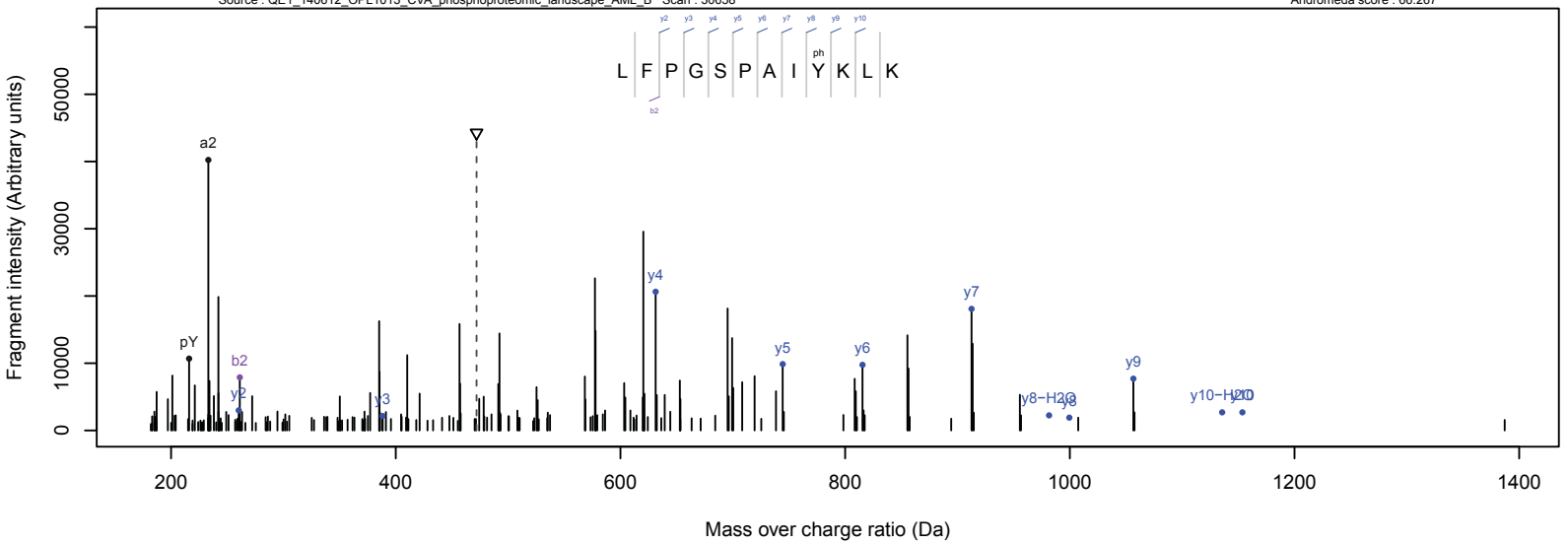
Andromeda score : 66.267

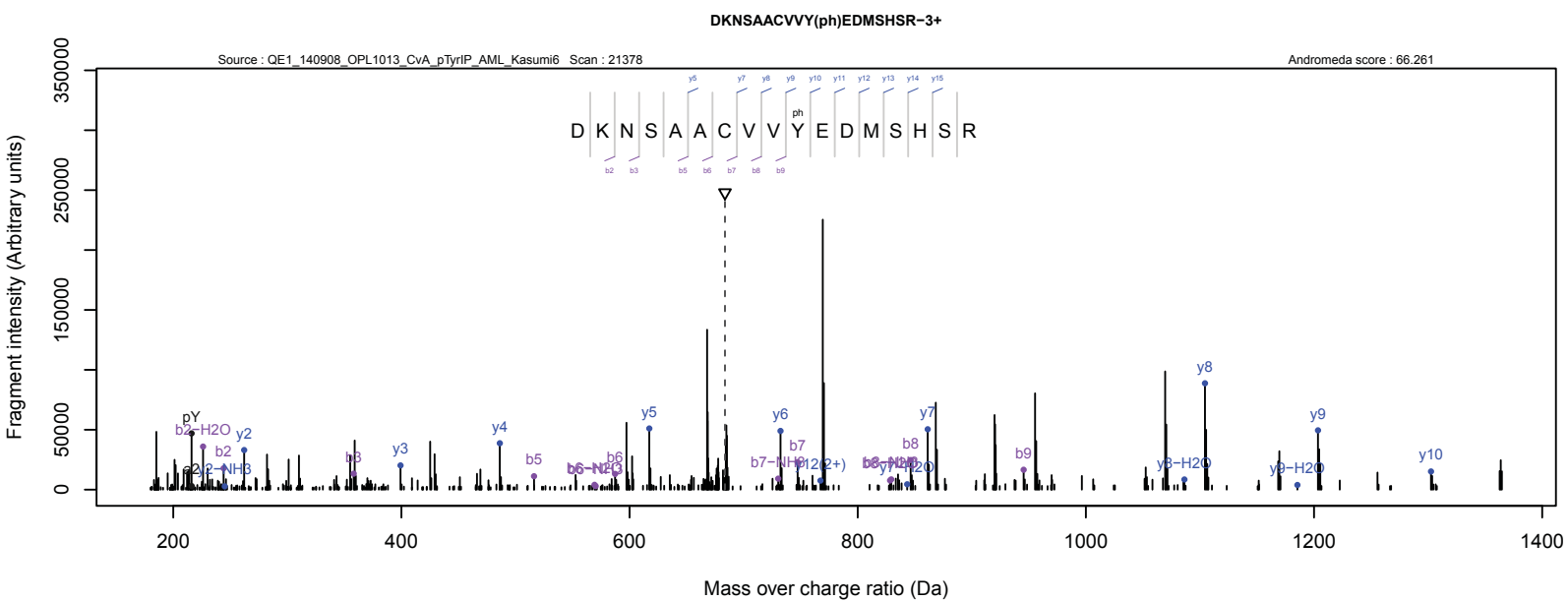
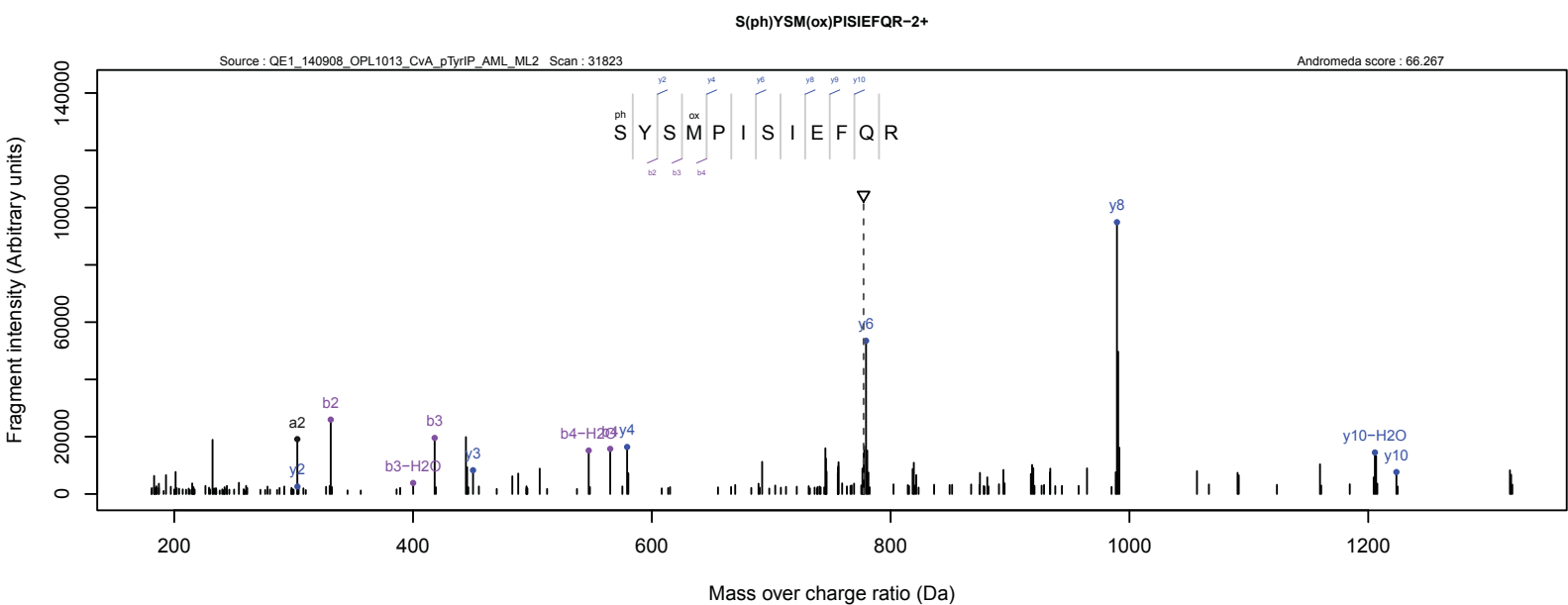
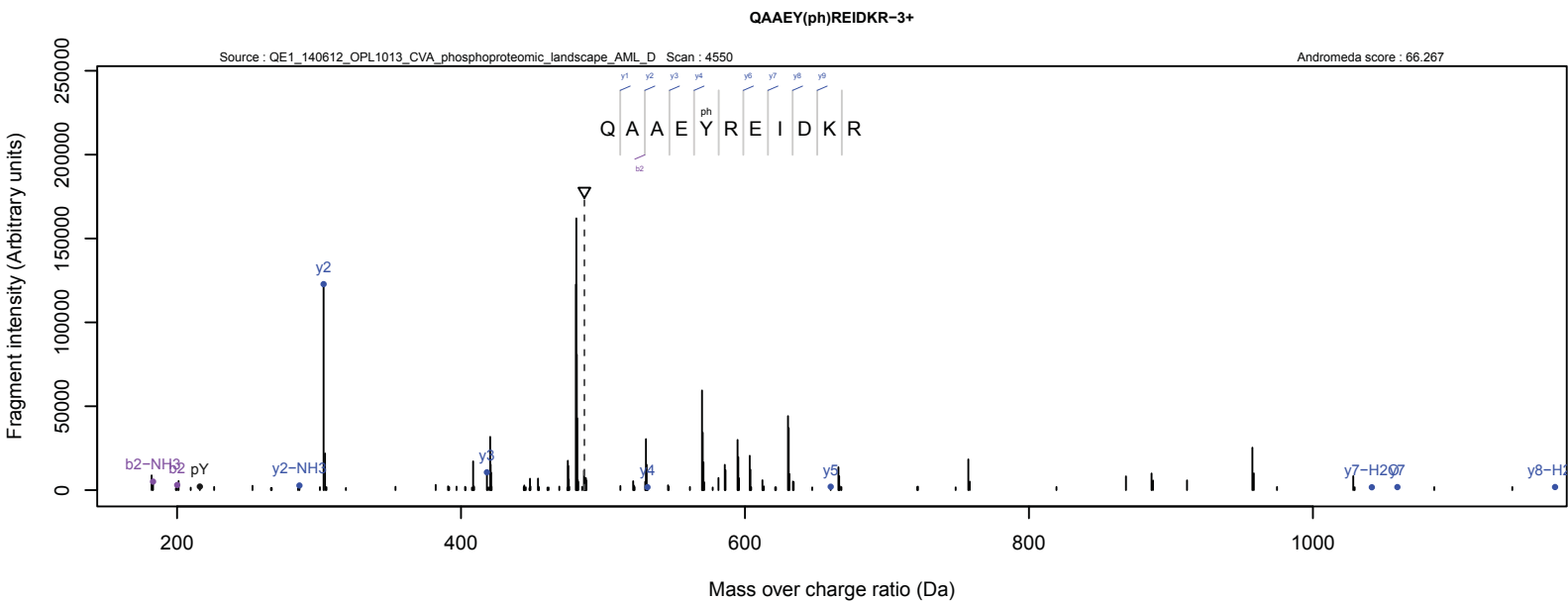


LFGSPAIY(ph)KLK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan : 30658

Andromeda score : 66.267

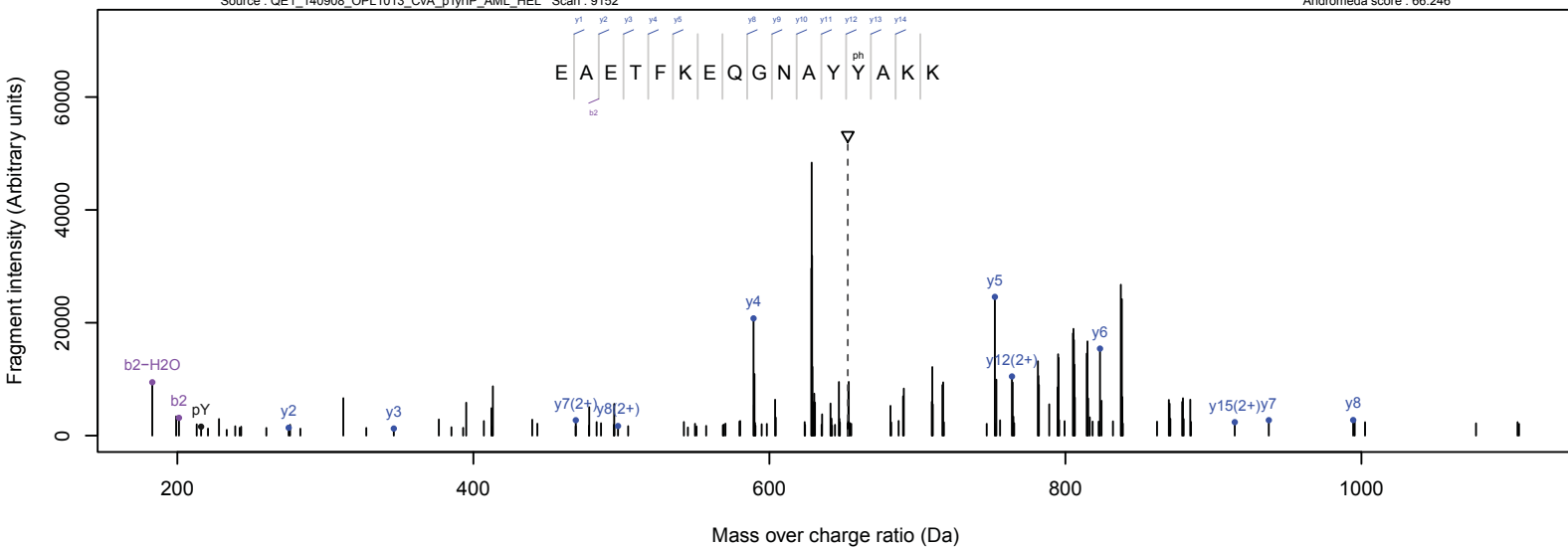




EAETFKEQGNAY(ph)AKK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 9152

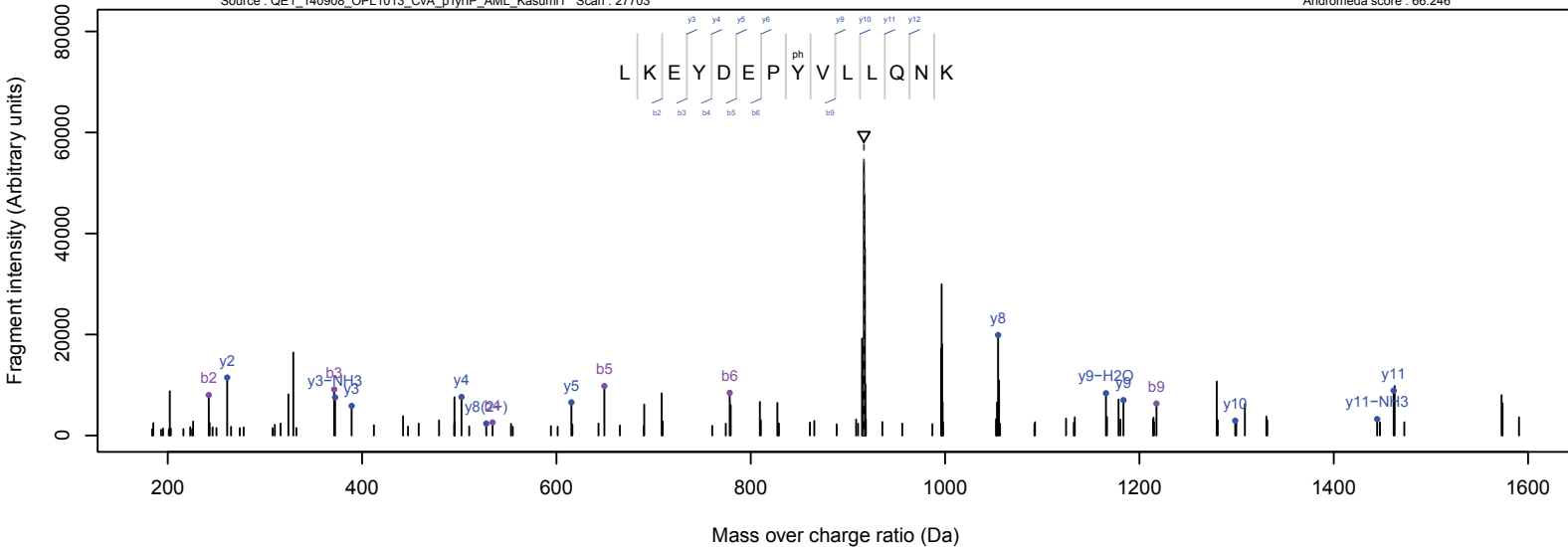
Andromeda score : 66.246



LKEYDEPY(ph)VLLQNK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27703

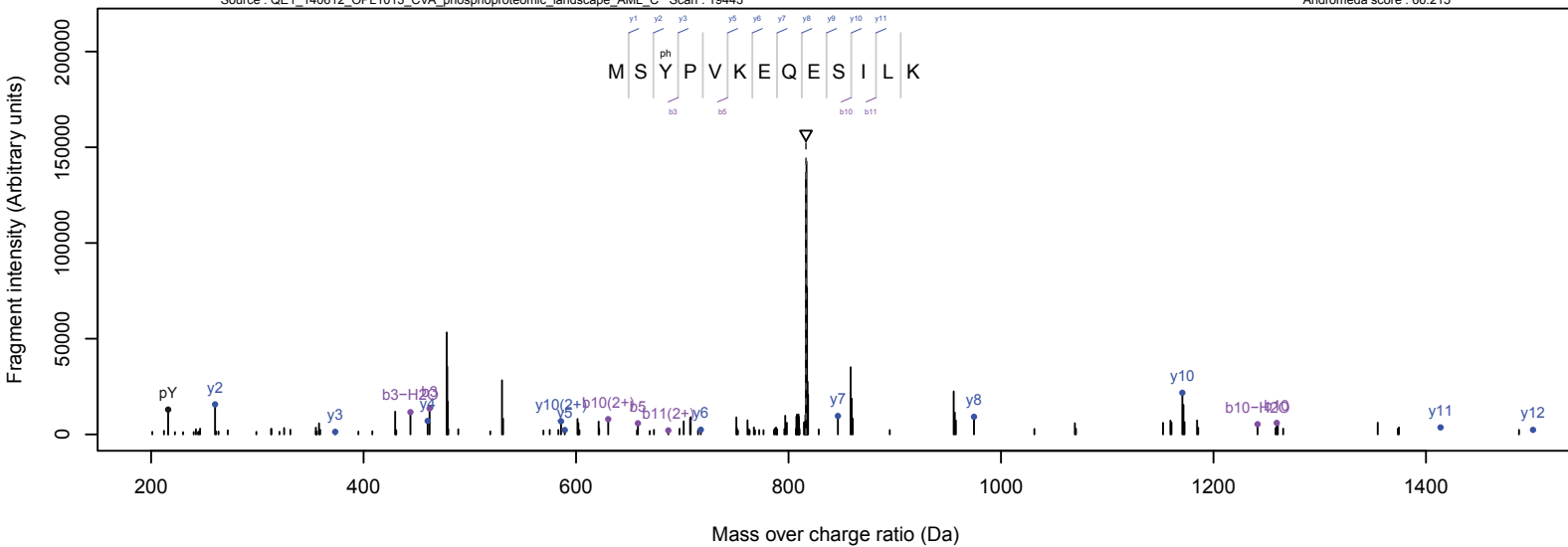
Andromeda score : 66.246



MSY(ph)PVKEQESILK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19443

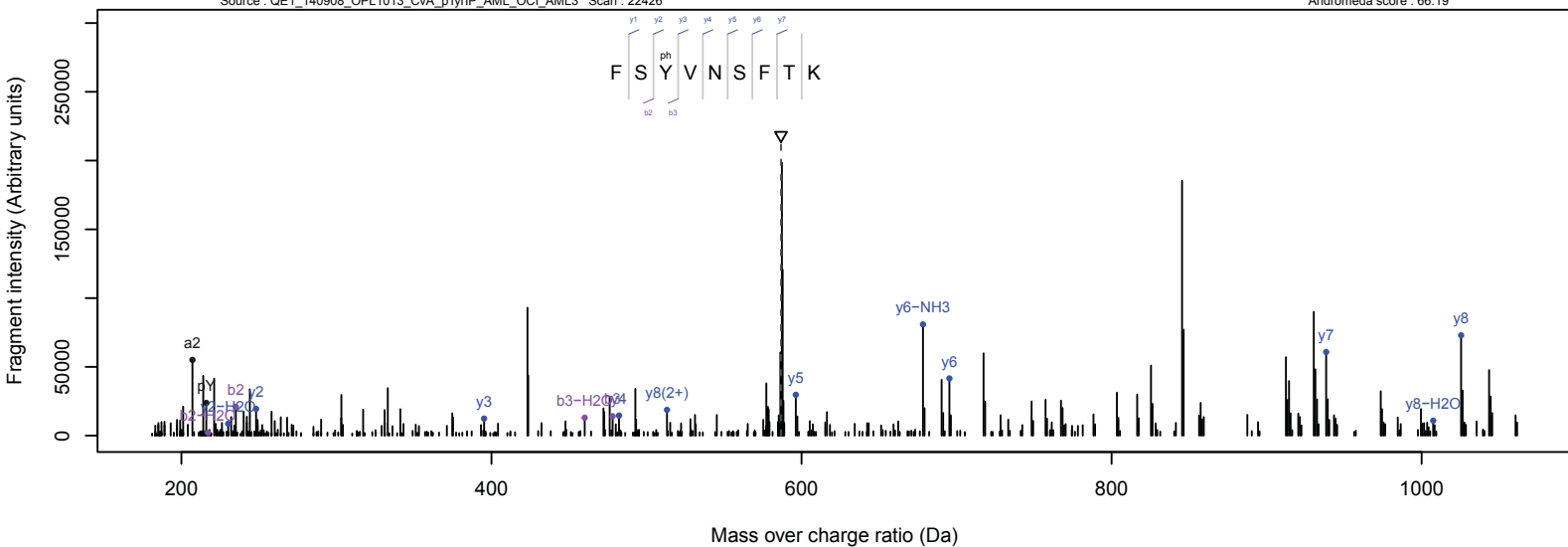
Andromeda score : 66.215



## FSY(ph)VNSFTK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 22426

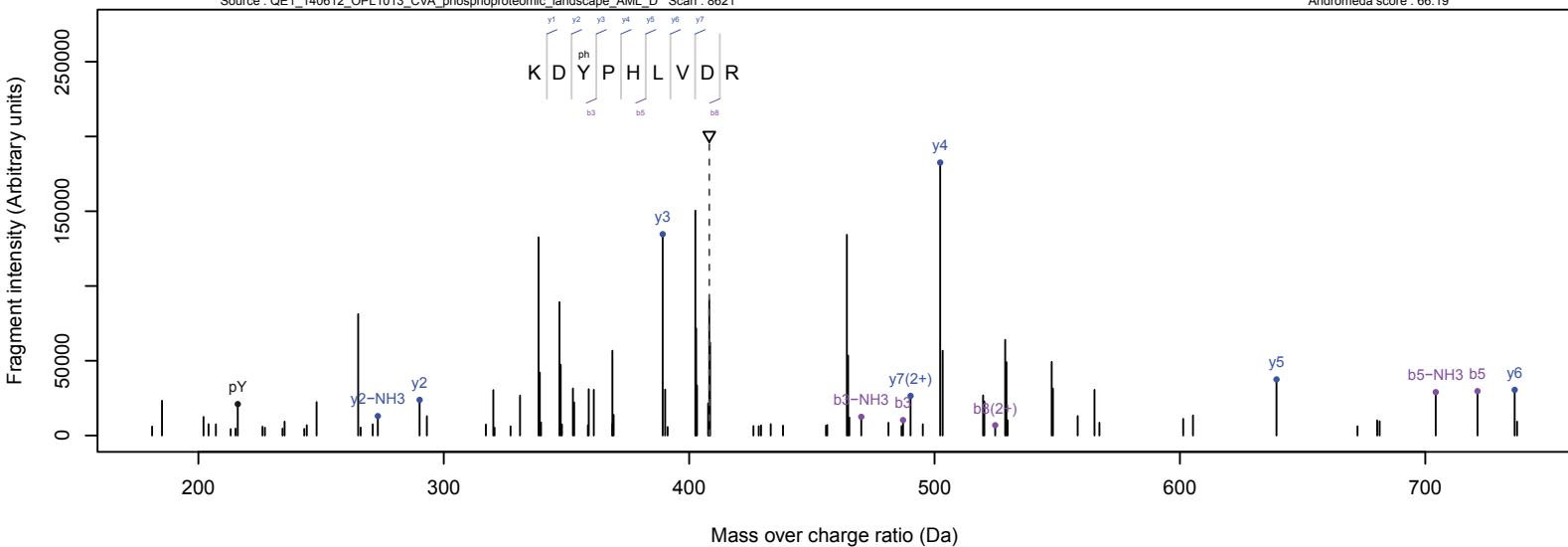
Andromeda score : 66.19



## KDY(ph)PHLVDR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan : 8621

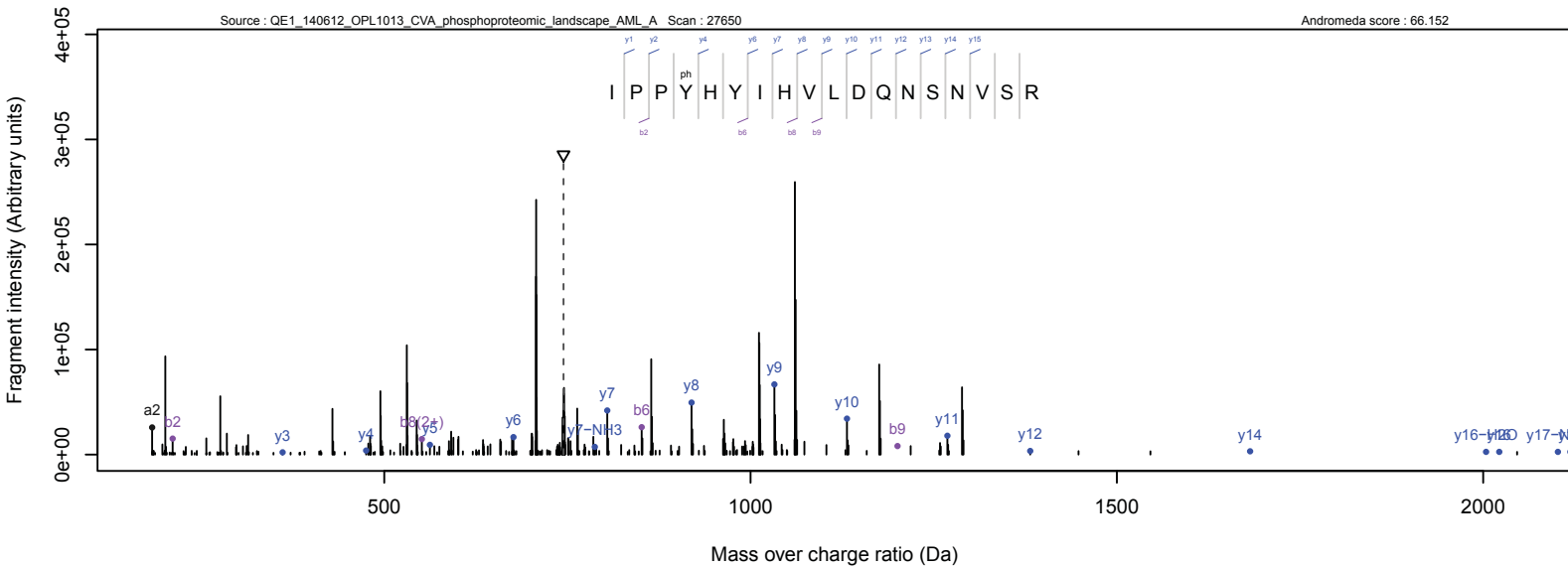
Andromeda score : 66.19



## IPPY(ph)HYIHVLDQNSVSR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_A Scan : 27650

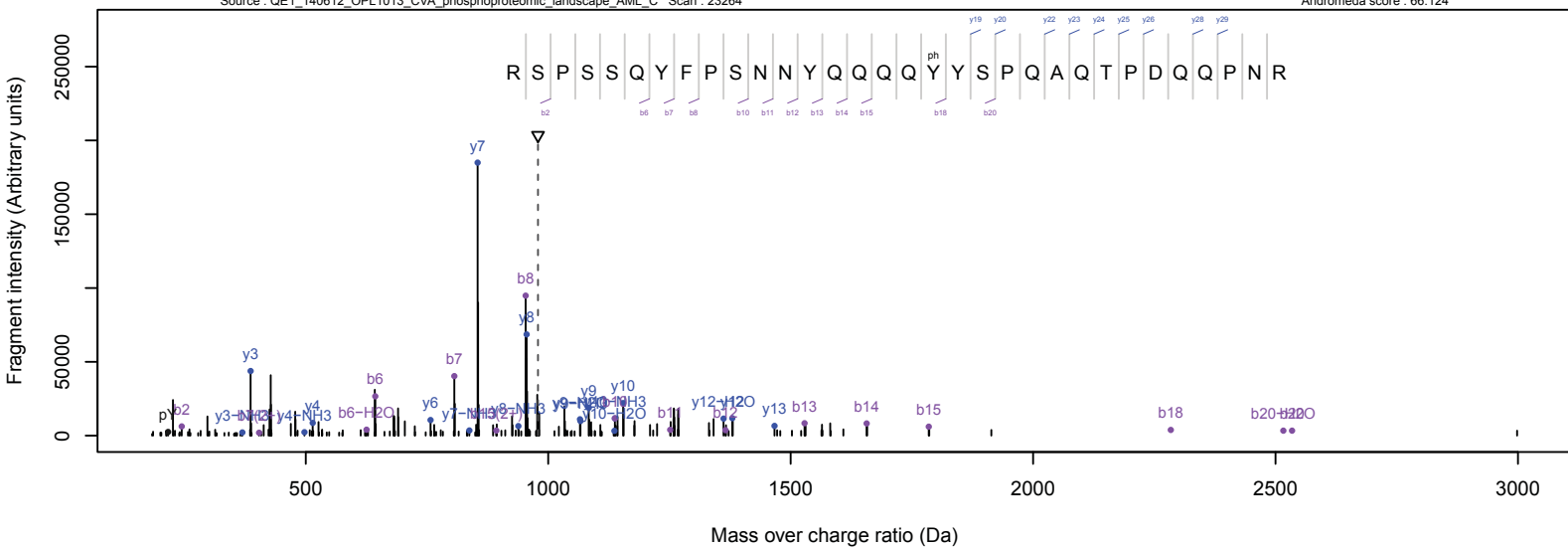
Andromeda score : 66.152



RSPSSQYFSPSNNYQQQY(ph)YSPQAQTPDQQPNR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23264

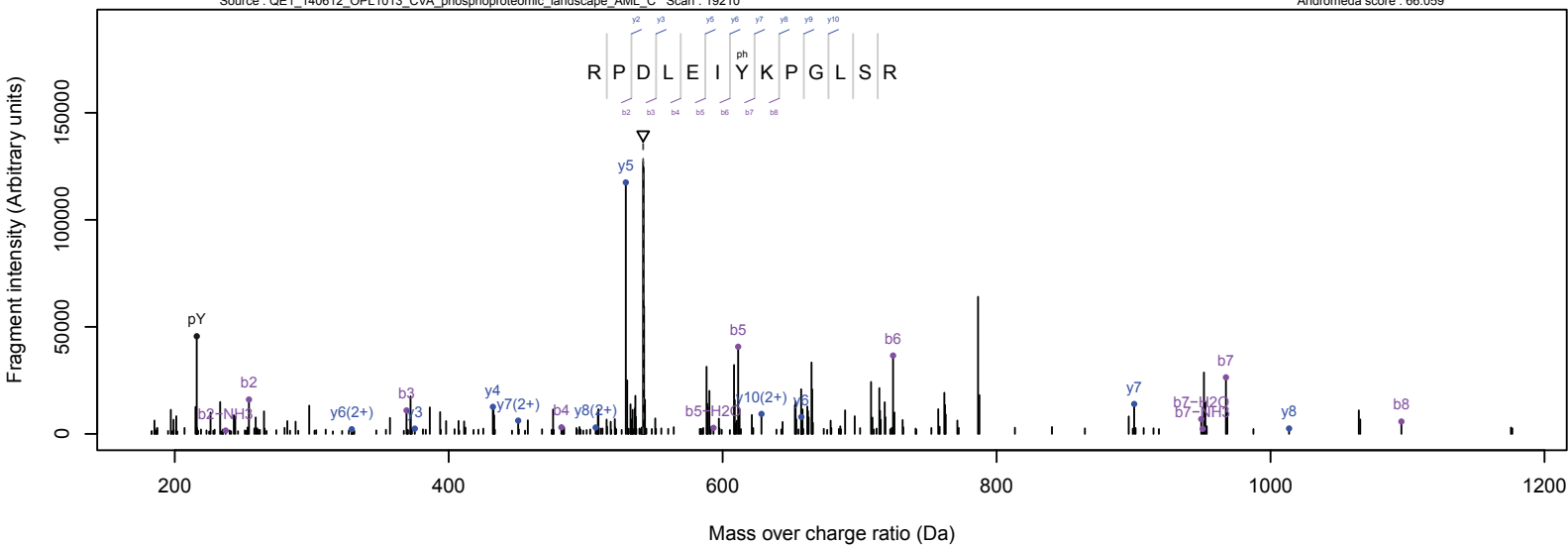
Andromeda score : 66.124



RPDLEIY(ph)KPGLSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19210

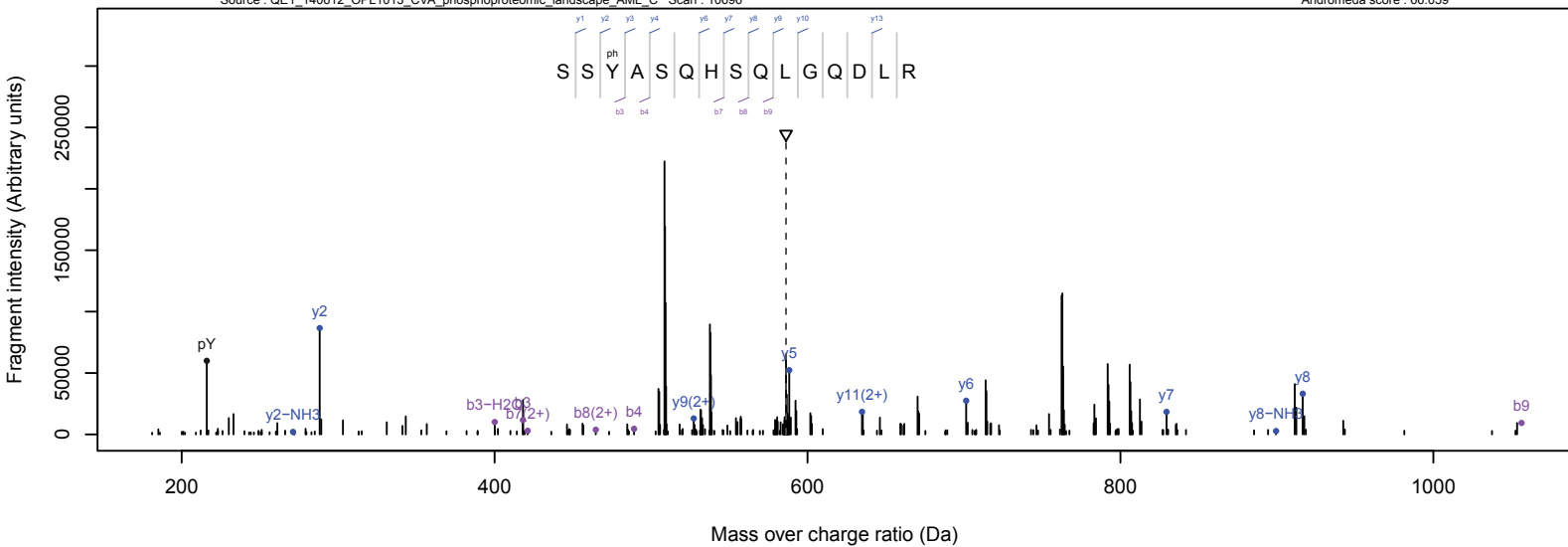
Andromeda score : 66.059



SSY(ph)ASQHSQLGQDLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10696

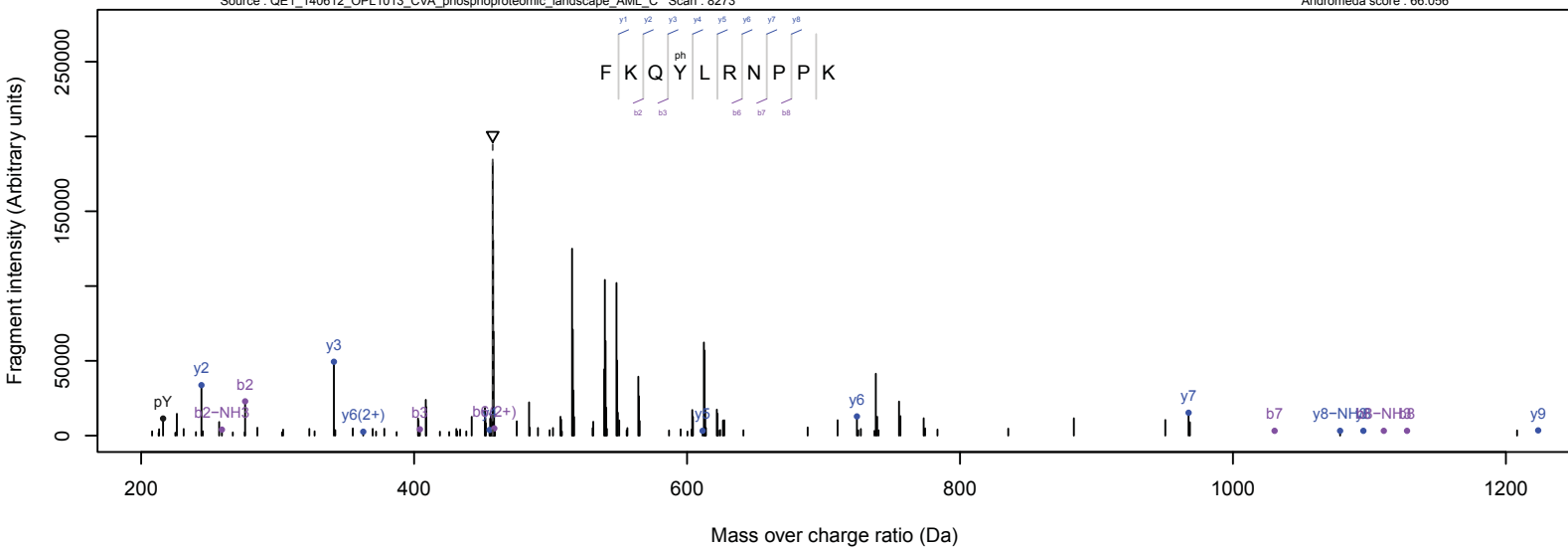
Andromeda score : 66.059



**FKQY(ph)LRNPPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8273

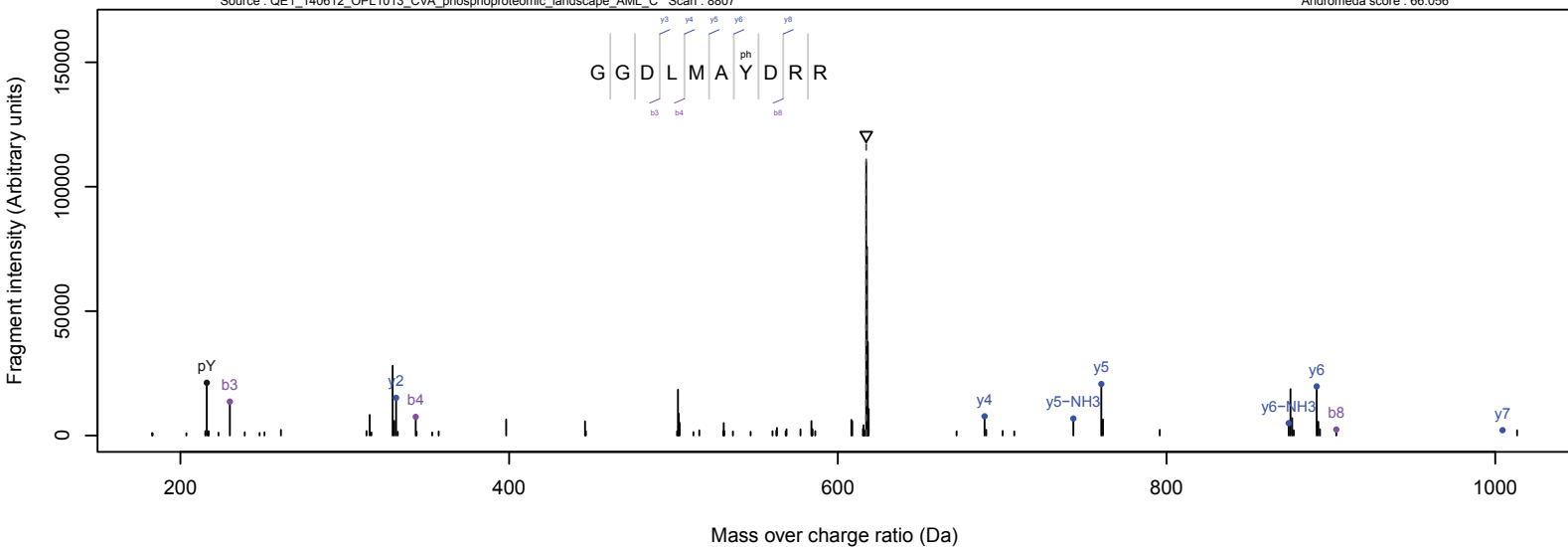
Andromeda score : 66.056



**GGDLMAY(ph)DRR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8807

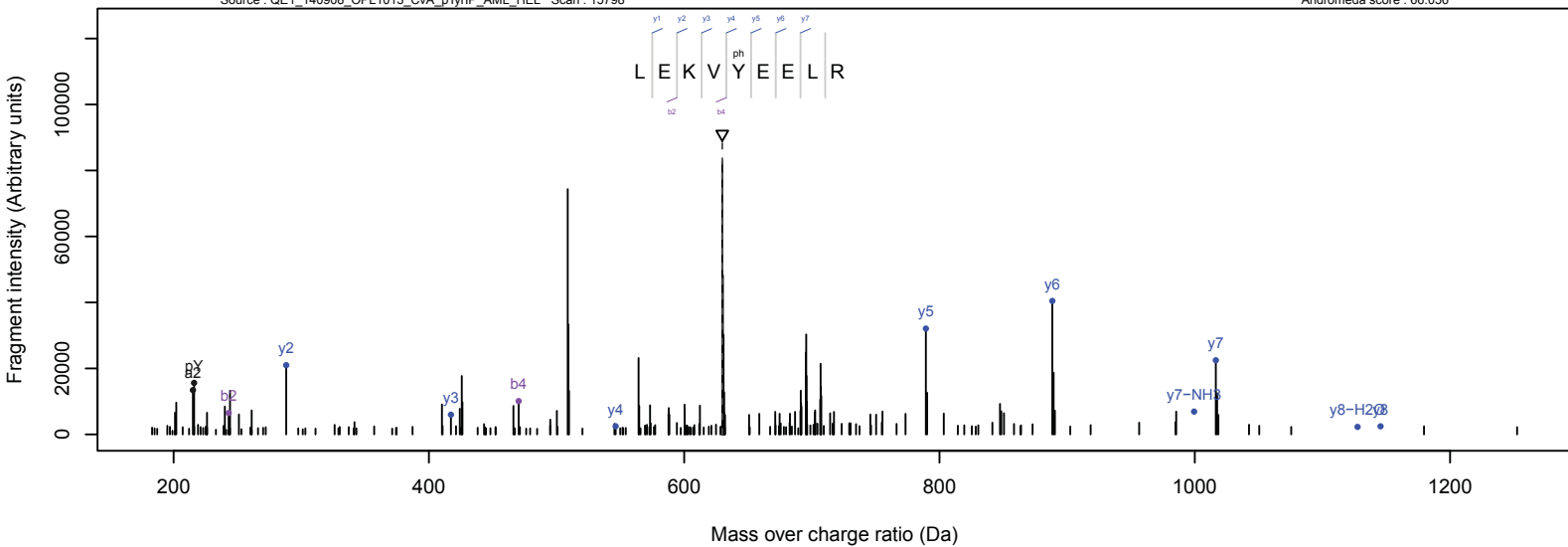
Andromeda score : 66.056



**LEKVV(ph)EELR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 15798

Andromeda score : 66.056

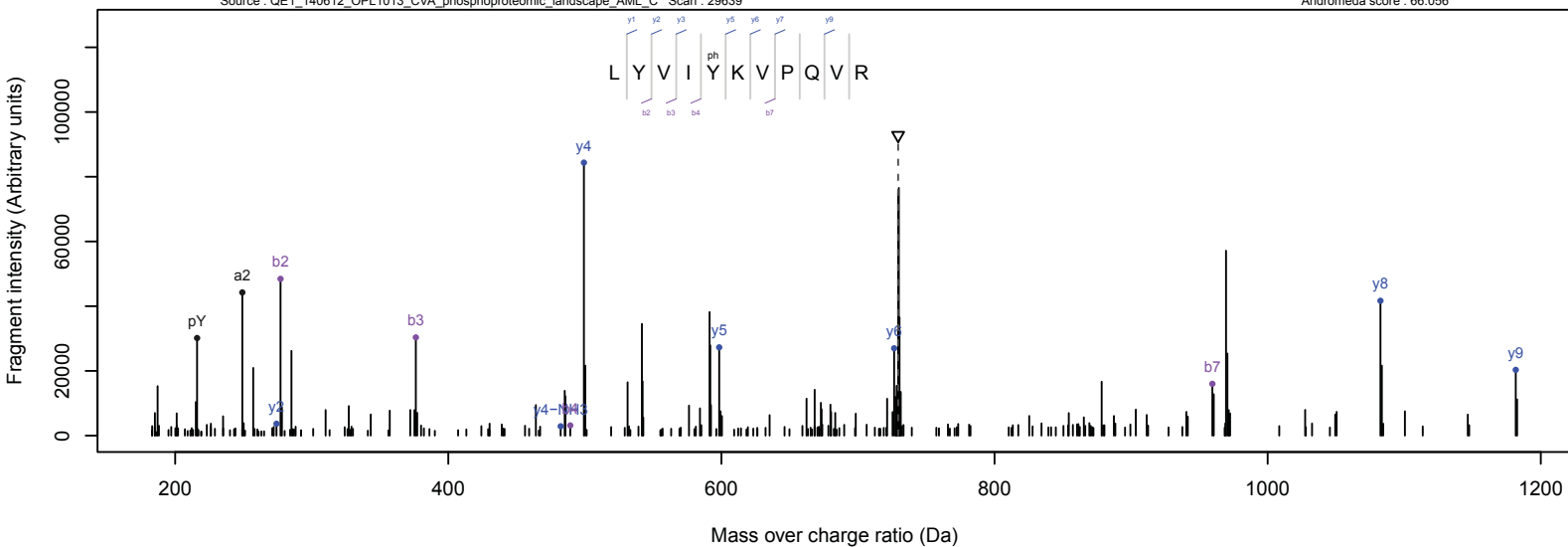




## LYVIY(ph)KVPQR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 29639

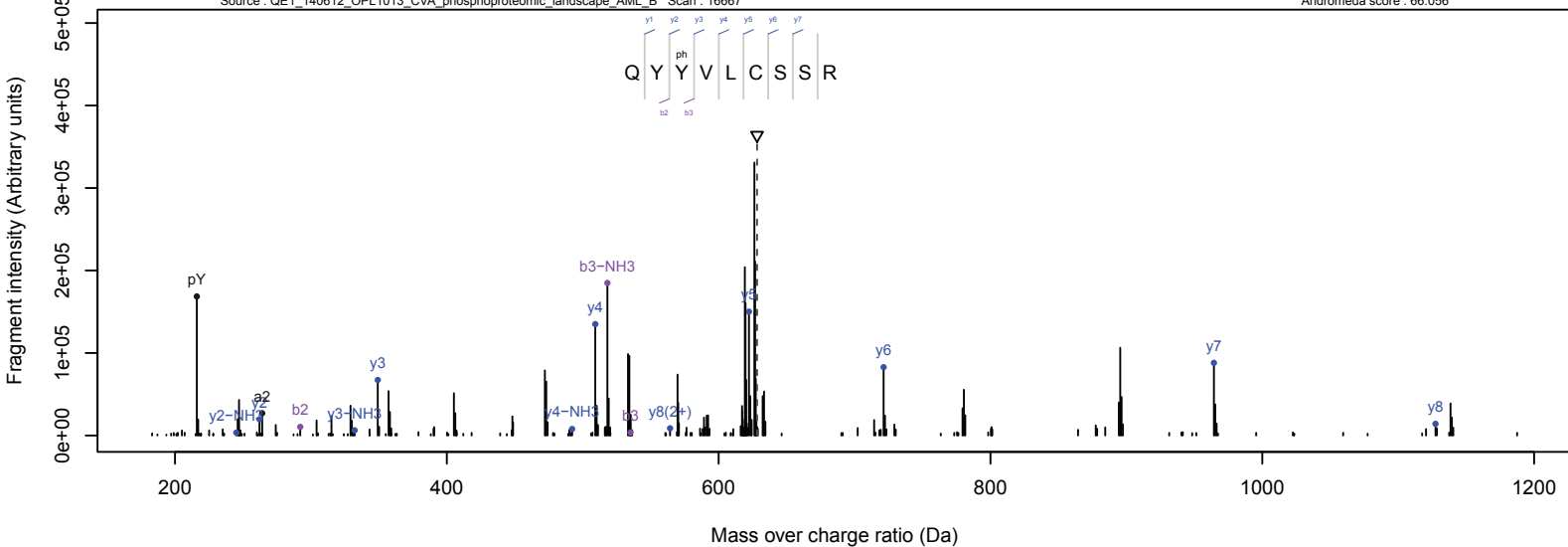
Andromeda score: 66.056



## QYY(ph)VLCSSR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 16667

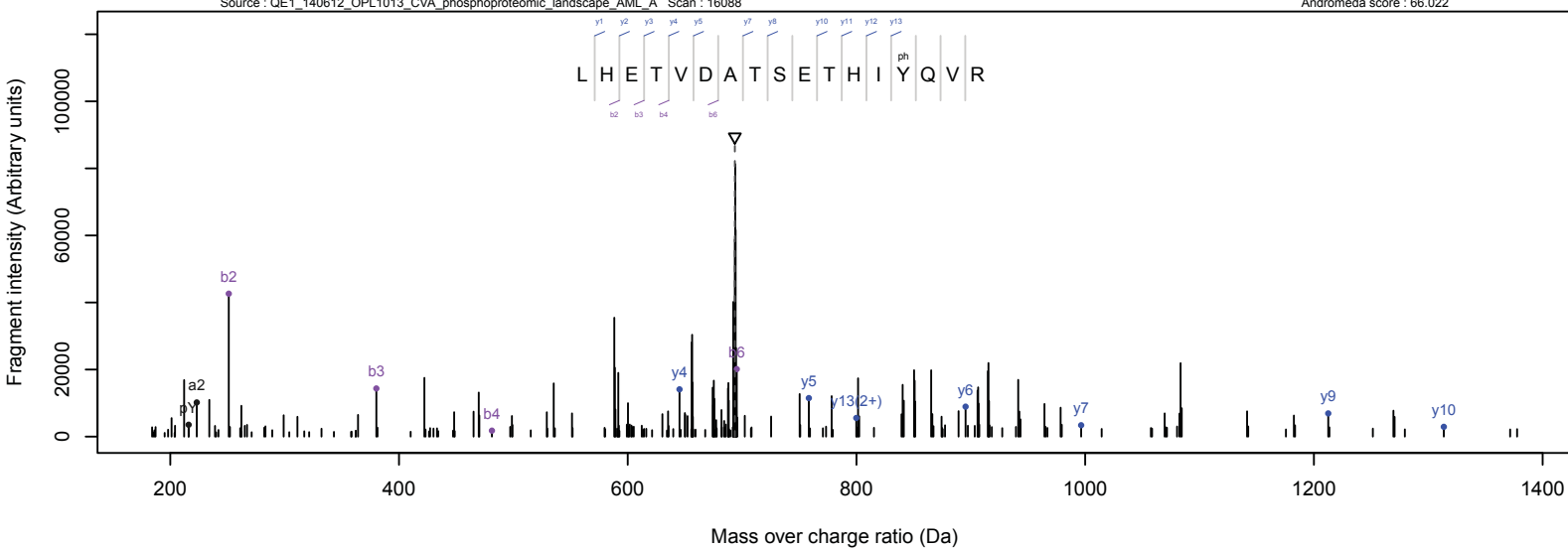
Andromeda score: 66.056



## LHETVDATSETHIY(ph)QVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 16088

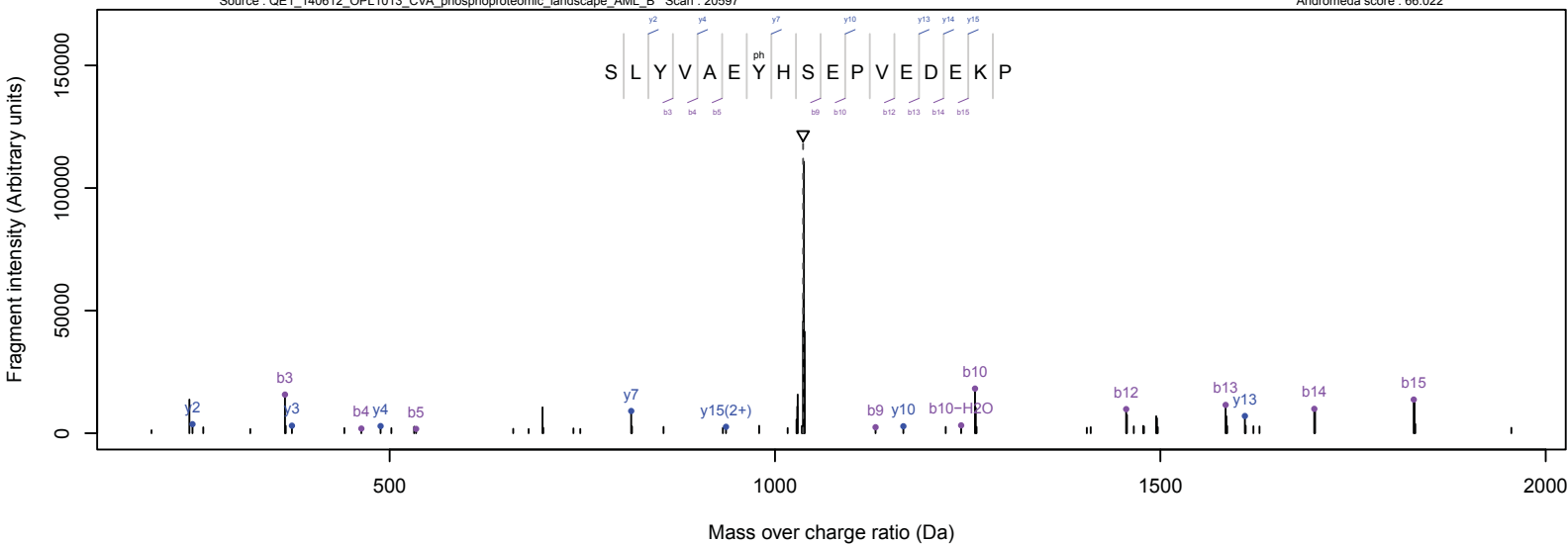
Andromeda score: 66.022



**SLYVAEY(ph)HSEPVDEDEK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 20597

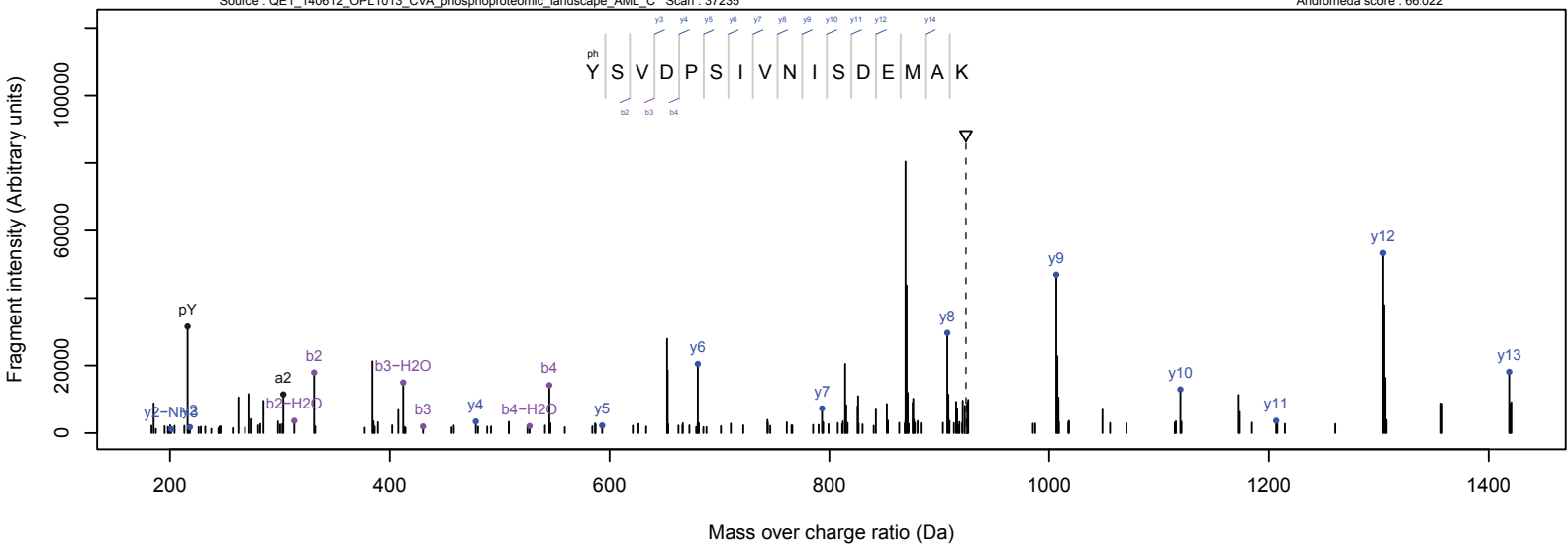
Andromeda score: 66.022



**Y(ph)SVDPSIVNISDEMAK-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37235

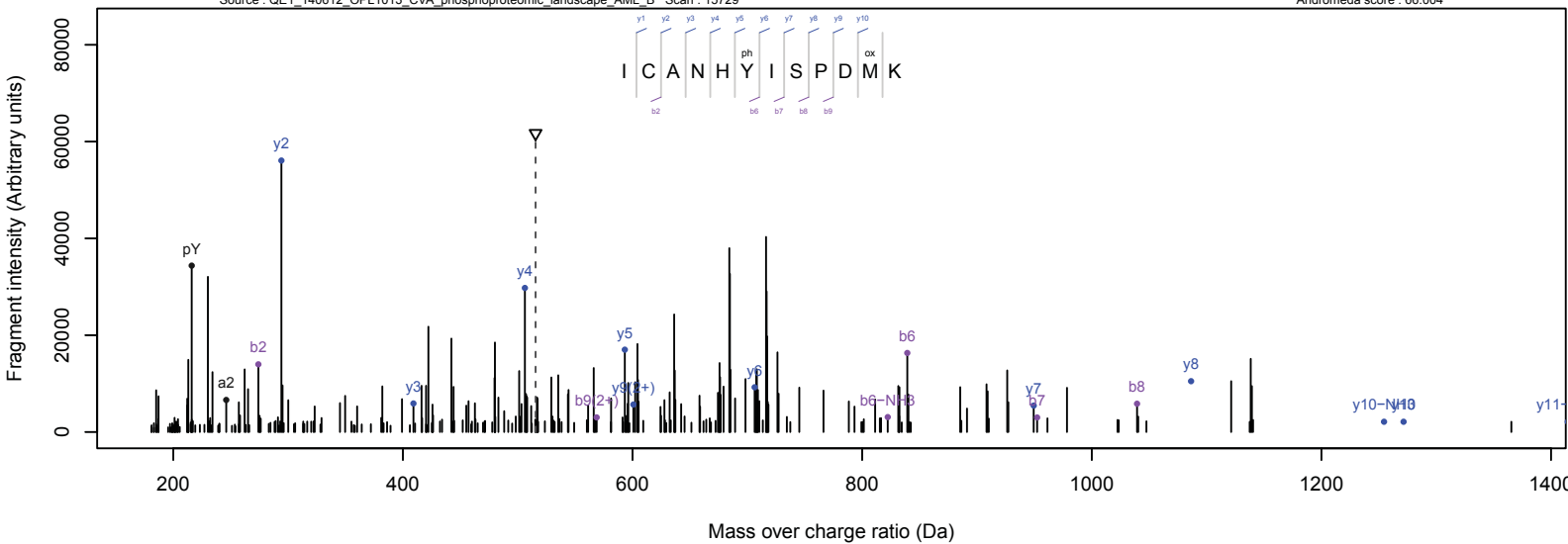
Andromeda score: 66.022



**ICANHYPH)ISPDM(ox)K-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15729

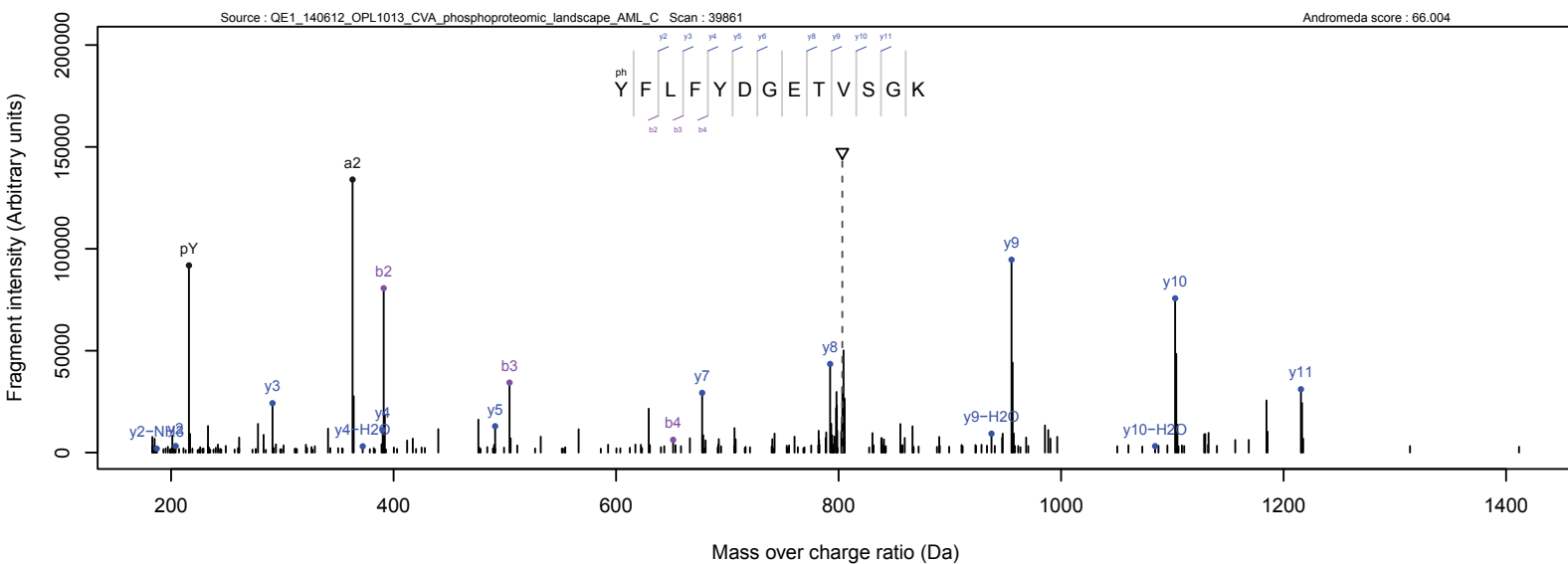
Andromeda score: 66.004



**Y(ph)FLFYDGETVSGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39861

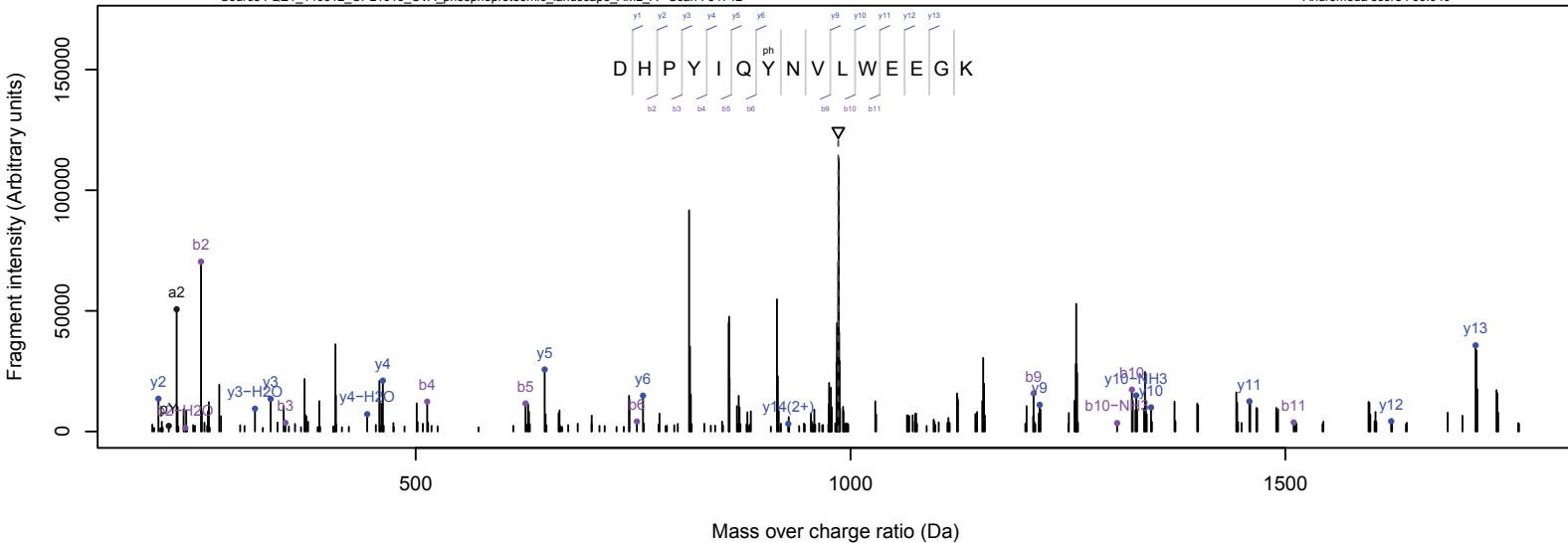
Andromeda score : 66.004



**DHPYIQY(ph)NVLWEEGK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34742

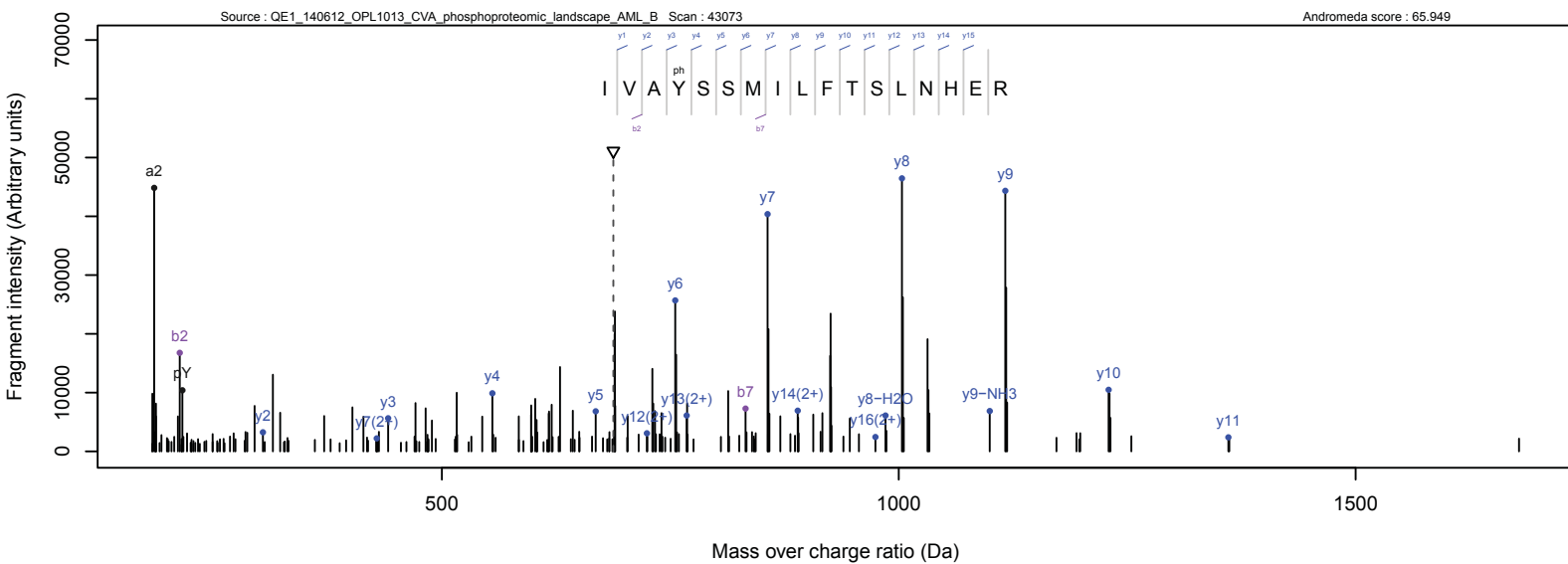
Andromeda score : 65.949



**IVAY(ph)SSMILFSLNHER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43073

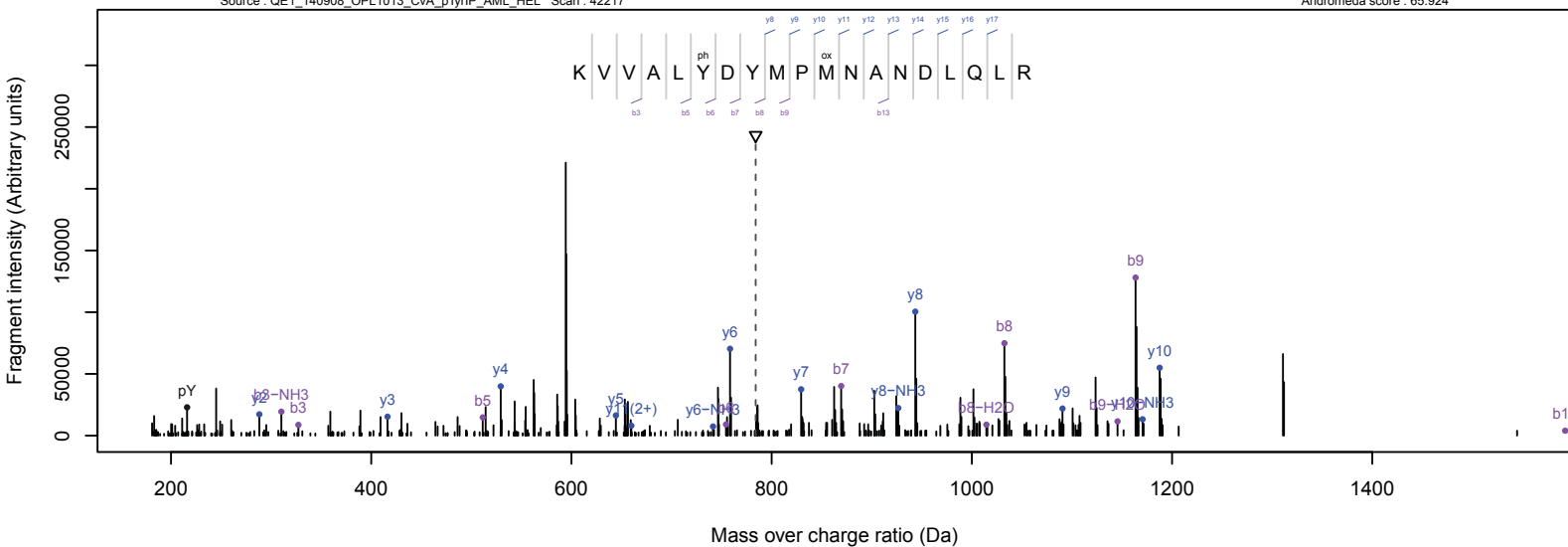
Andromeda score : 65.949



**KVVALY(ph)DYMPM(ox)NANDLQLR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 42217

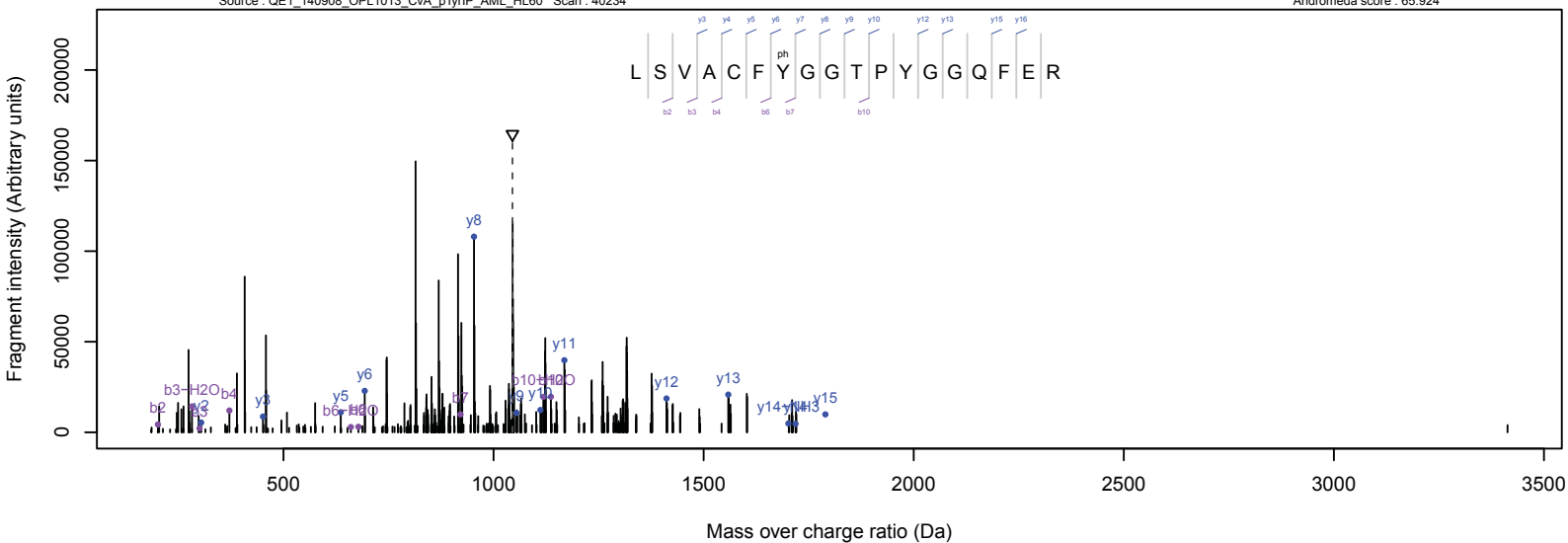
Andromeda score : 65.924



**LSVACFY(ph)GGTPYGGQFER-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 40234

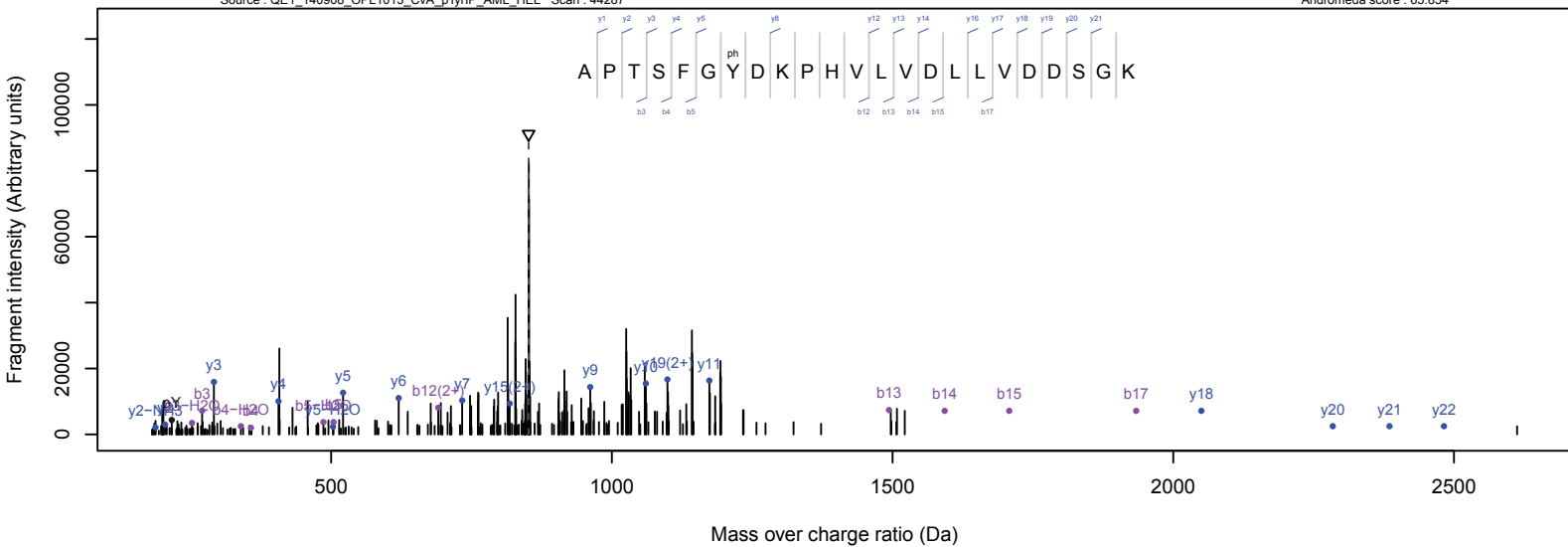
Andromeda score : 65.924



**APTSFGY(ph)DKPHVLVDLLVDDSGK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 44287

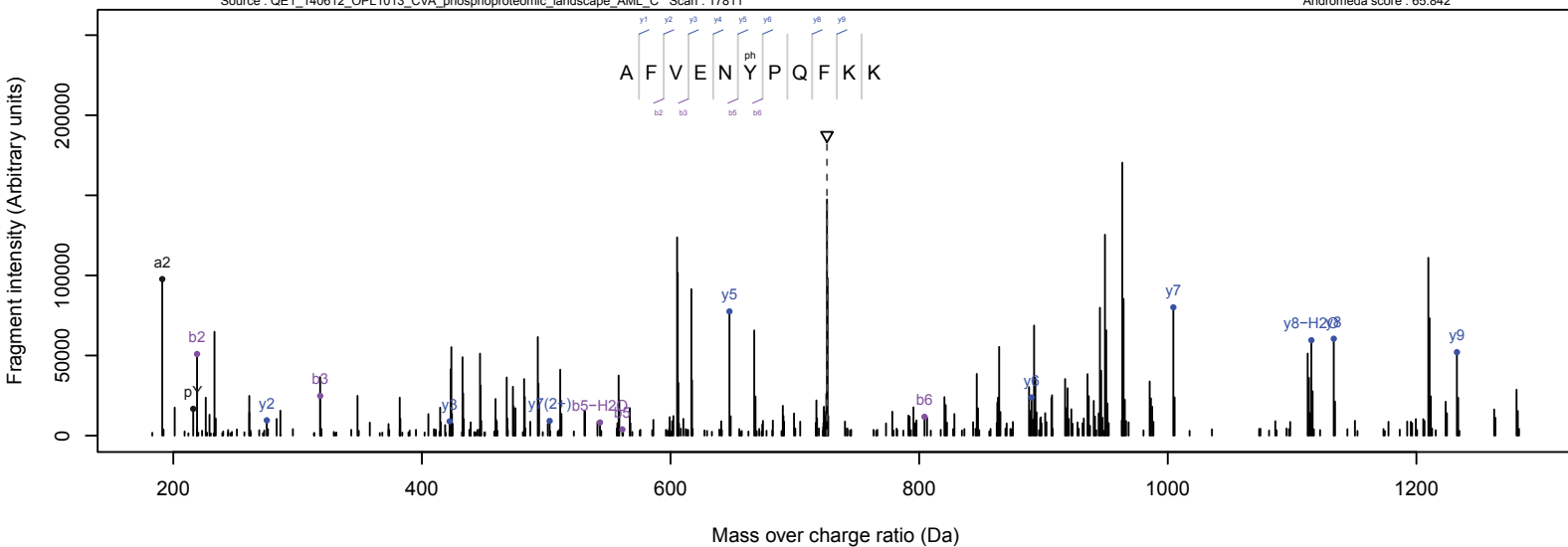
Andromeda score : 65.854



## AFVENY(ph)PQFKK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17811

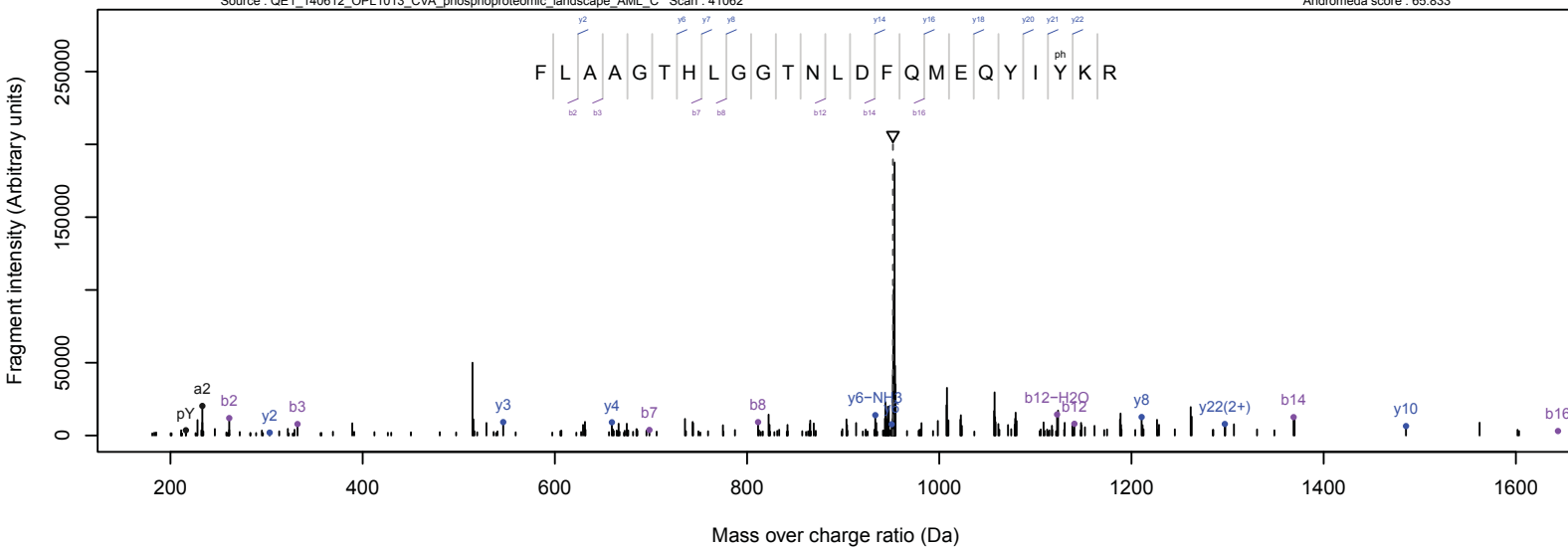
Andromeda score: 65.842



## FLAAGTHLGGTDLDFQMEQYIYKR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41062

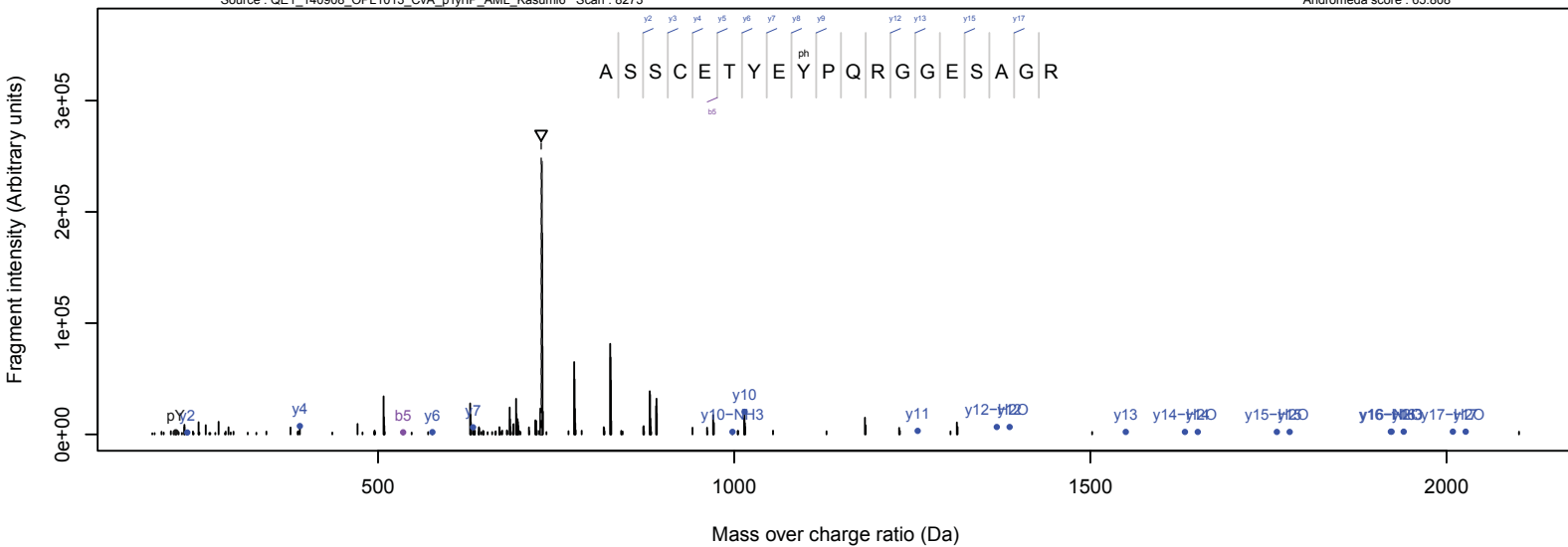
Andromeda score: 65.833



## ASSCETYEY(ph)PQRGGESAGR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 8273

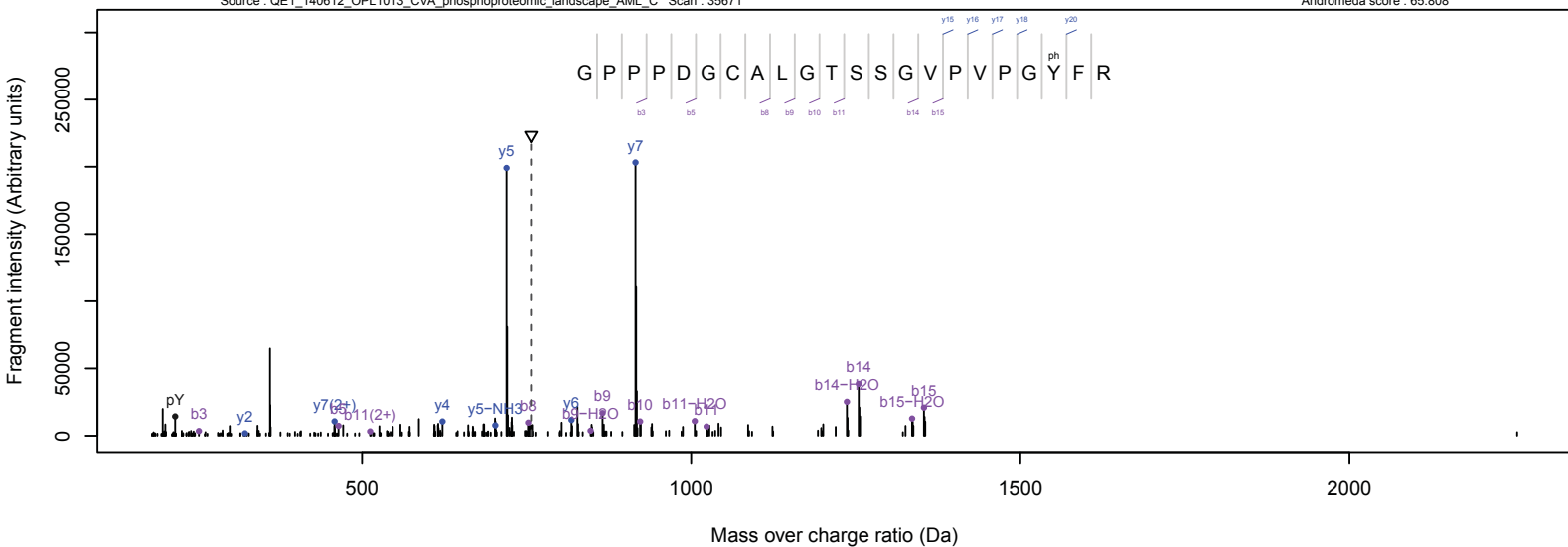
Andromeda score: 65.808



GPPPDGCALGTSVVPVPGYFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35671

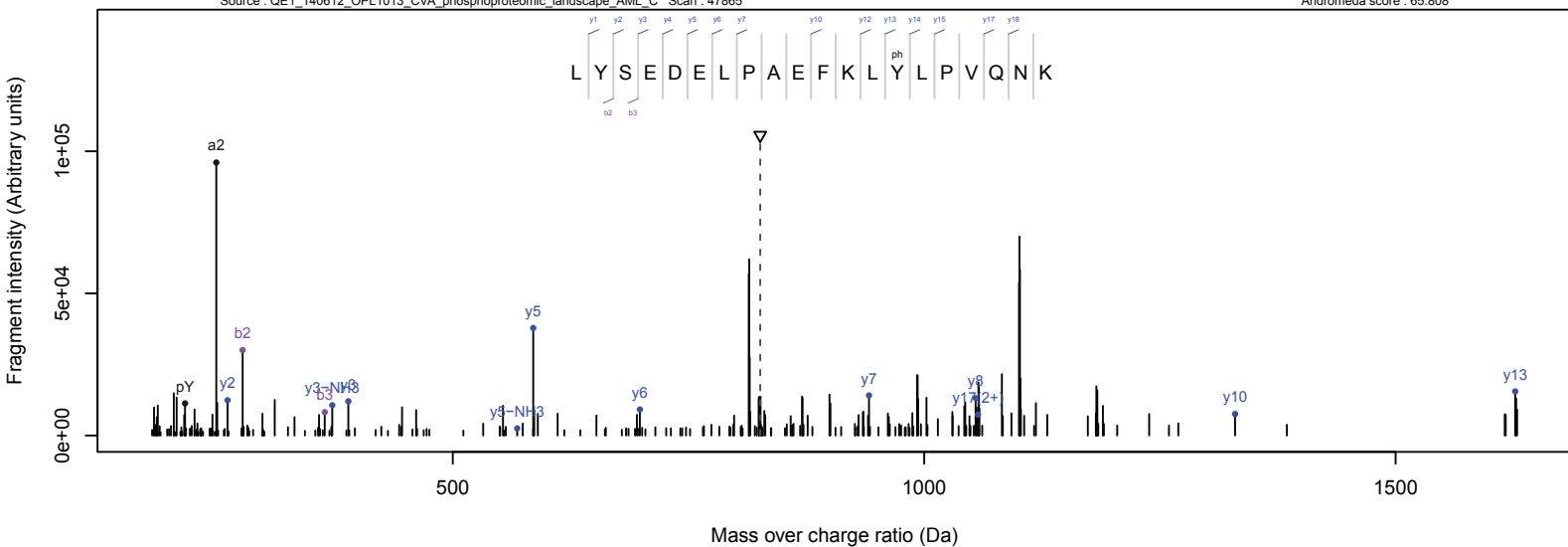
Andromeda score : 65.808



LYSEDELPAEFKLYLPVQNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47865

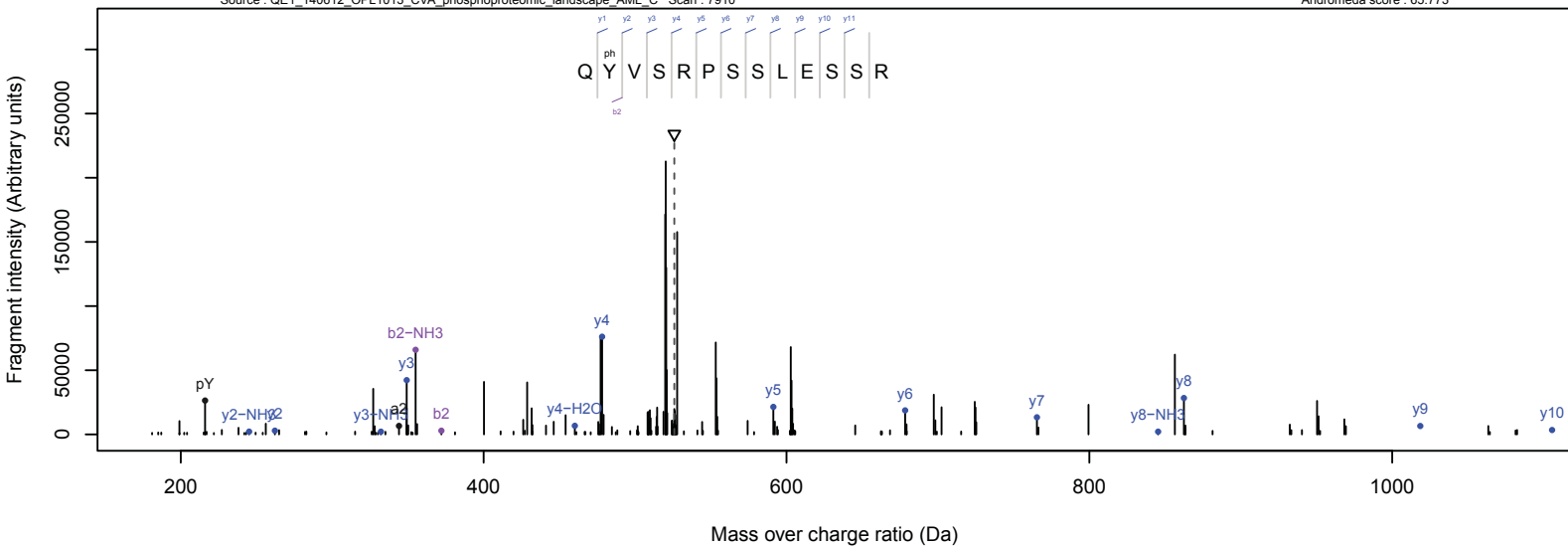
Andromeda score : 65.808



QYVSRPSSLESR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7910

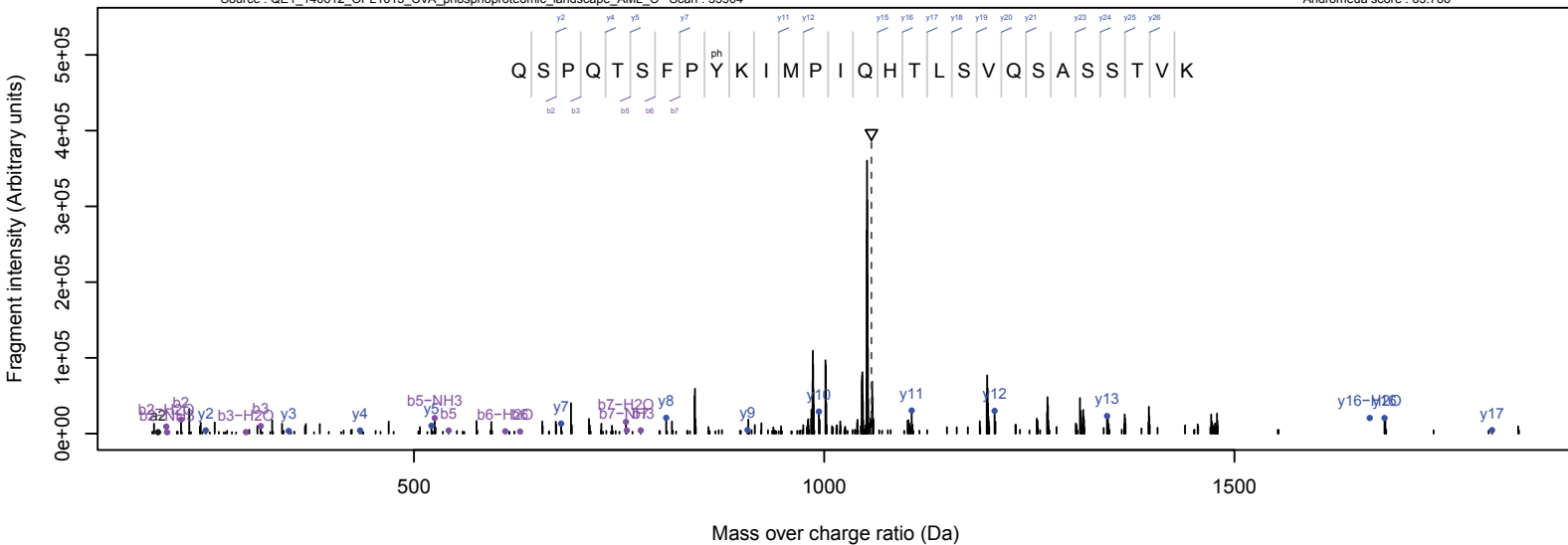
Andromeda score : 65.773



**QSPQTSFPY(ph)KIMPIIQHTLSVQSASSTVK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35904

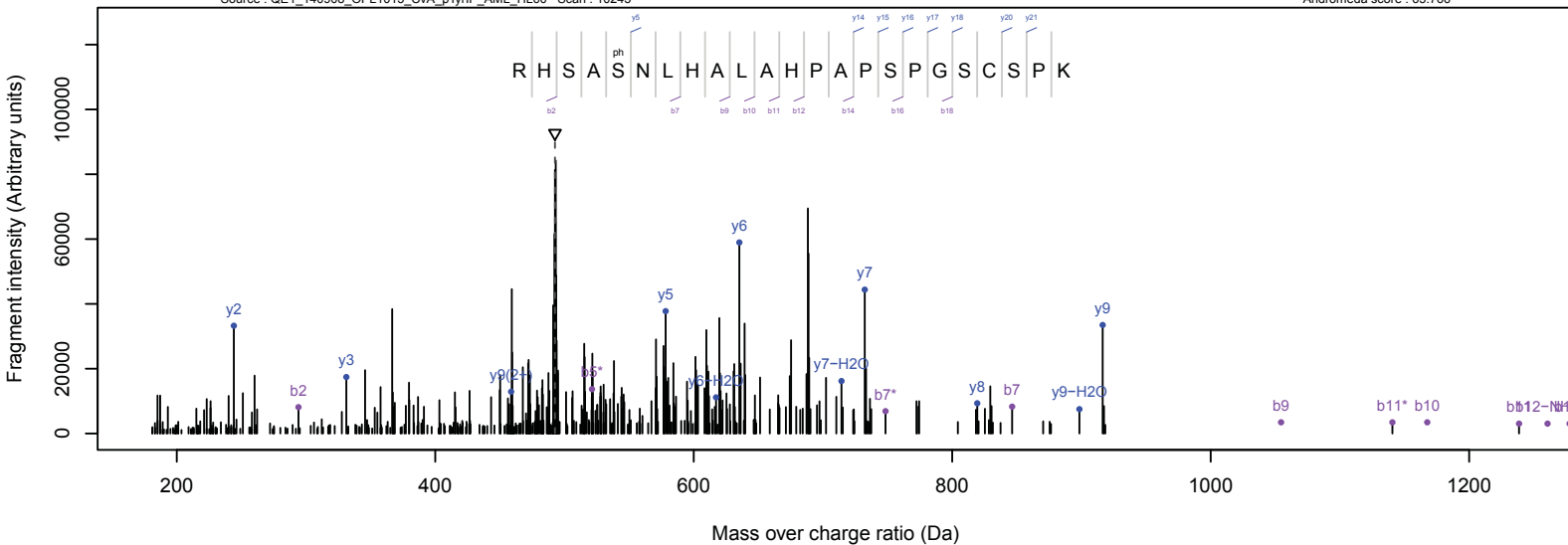
Andromeda score : 65.766



**RHSAS(ph)NLHALAHPAPSPGSCSPK-5+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 10243

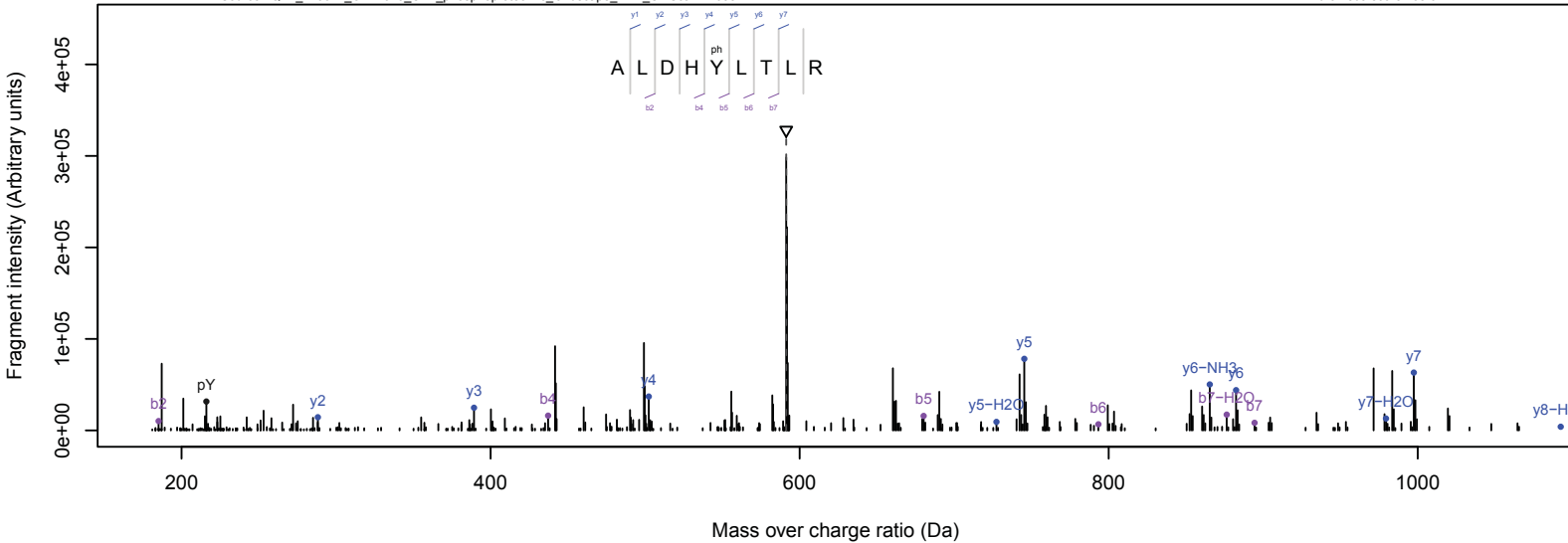
Andromeda score : 65.766



**ALDHY(ph)LTLR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26581

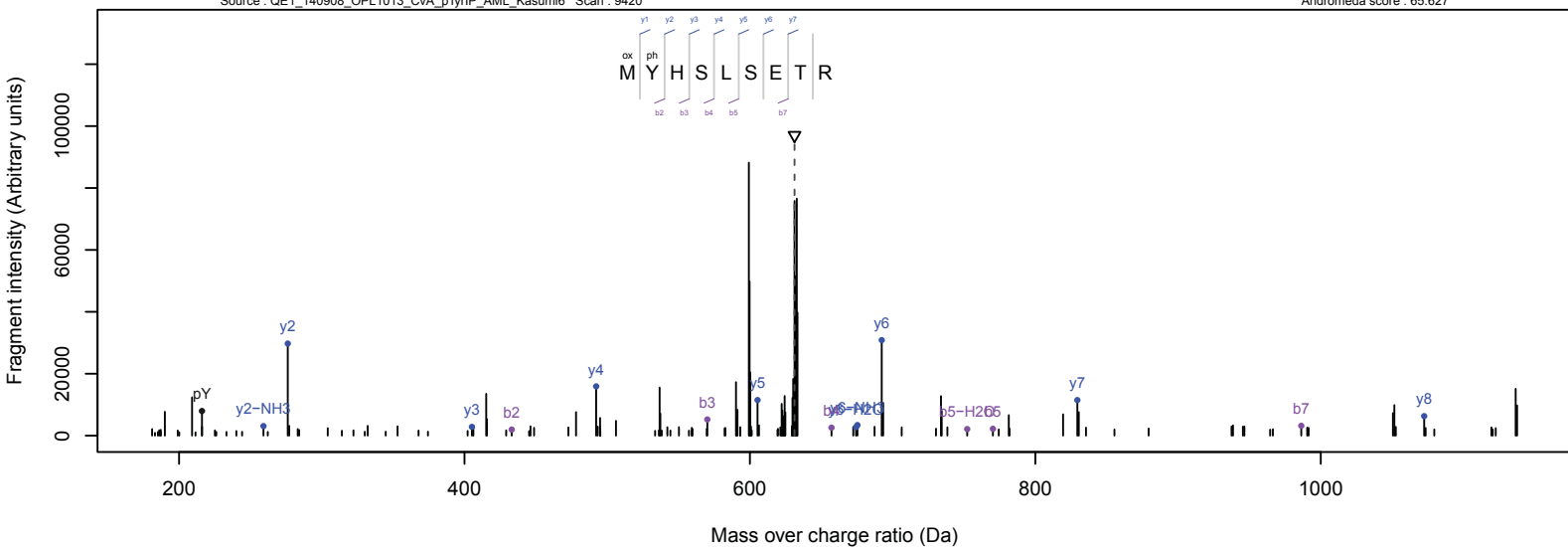
Andromeda score : 65.627



(ac)M(ox)Y(ph)HSLSETR-2+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan: 9420

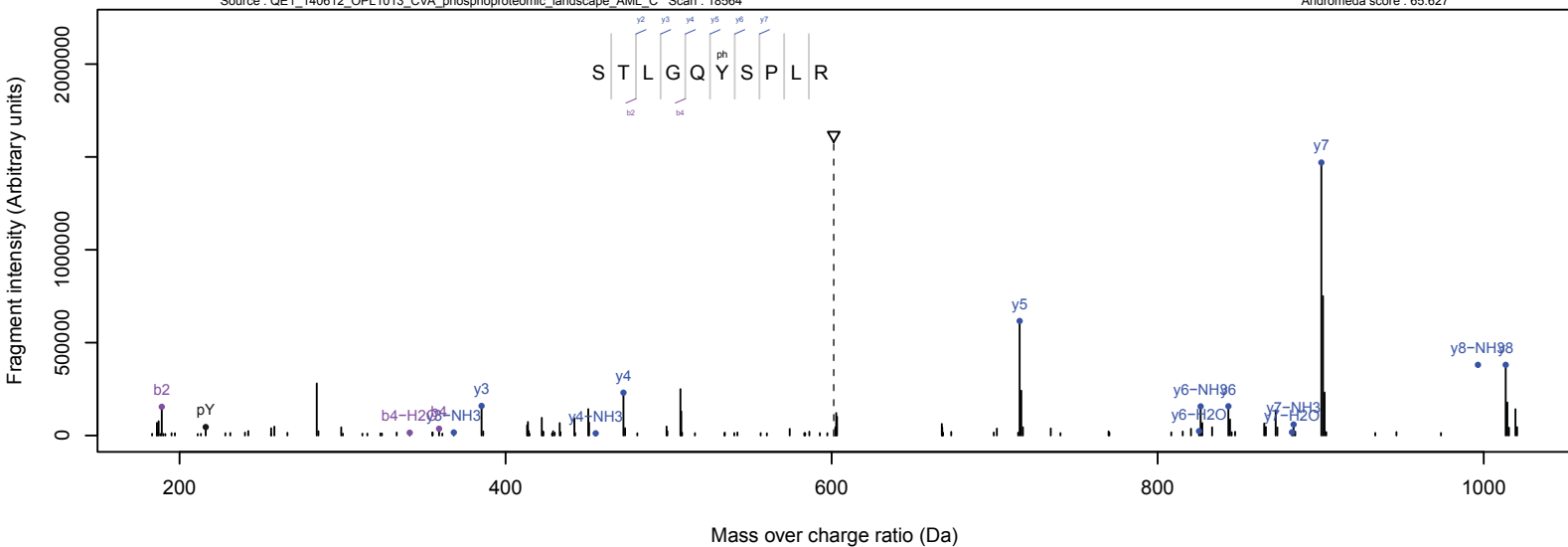
Andromeda score: 65.627



STLGQY(ph)SPLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18564

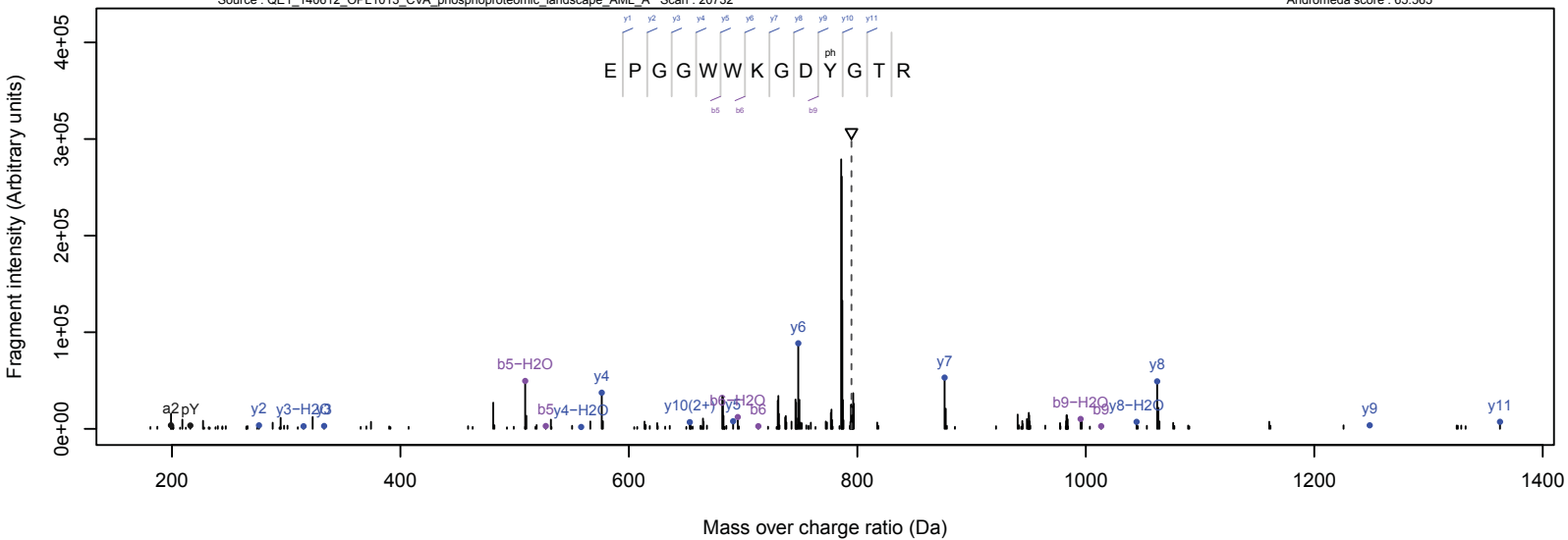
Andromeda score: 65.627



EPGGWVKDY(ph)GTR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 20732

Andromeda score: 65.565

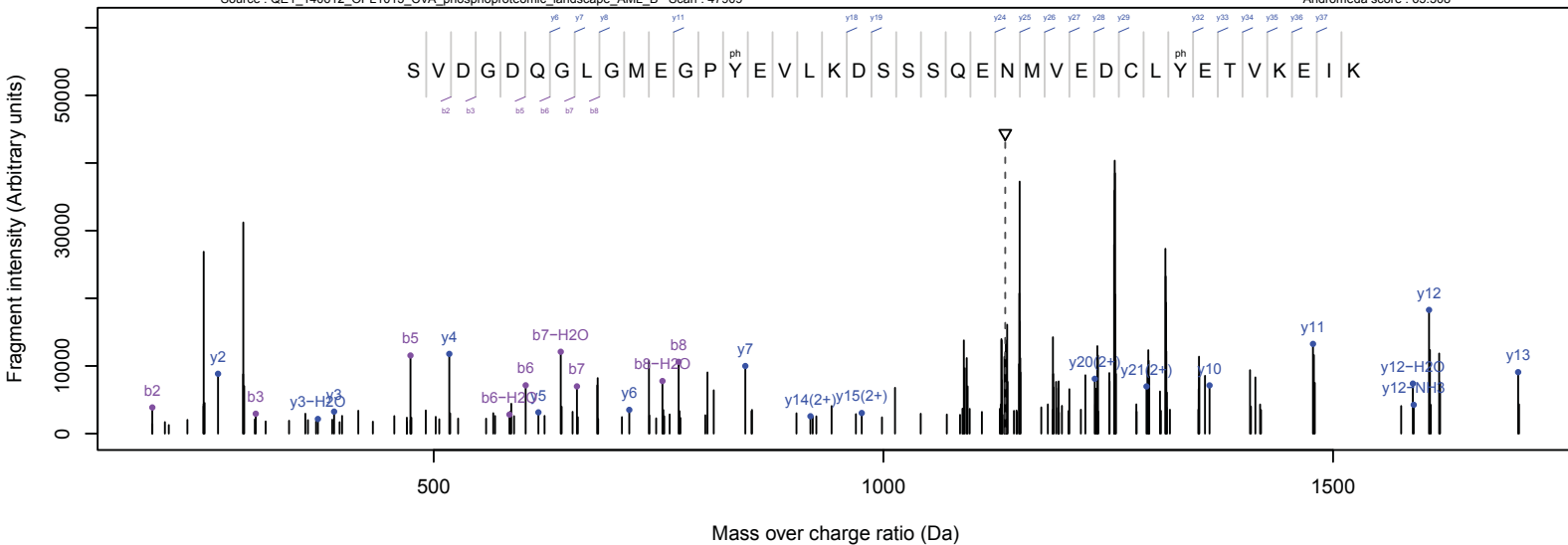




SVDGDQLGMEGPY(ph)EVLKDSSSQENMVEDCLY(ph)ETVKEIK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47909

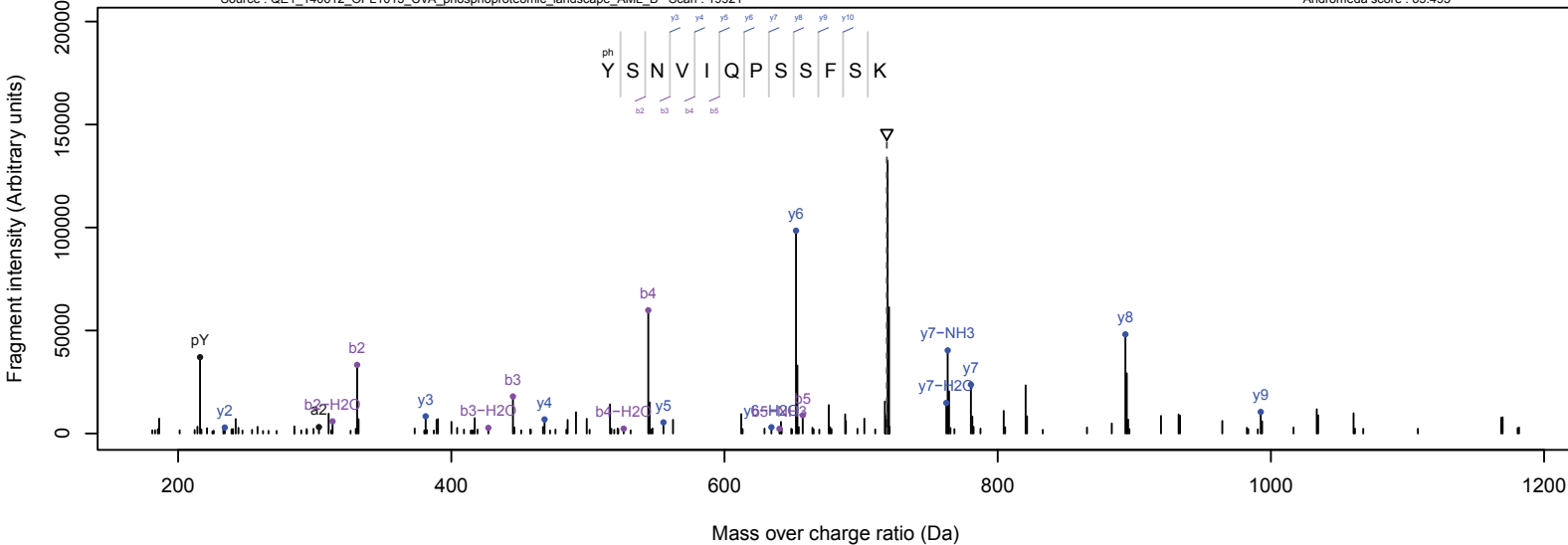
Andromeda score : 65.508



Y(ph)SNVIQSSFSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19921

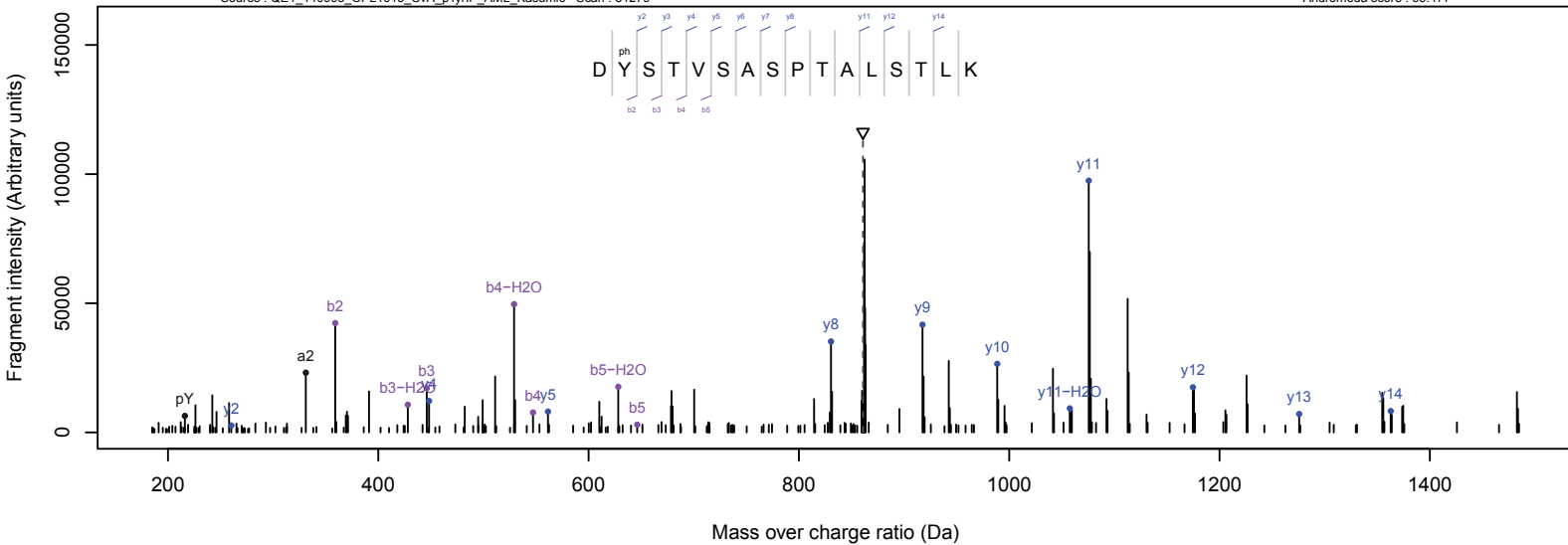
Andromeda score : 65.495



DY(ph)STVSASPTALSTLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 31279

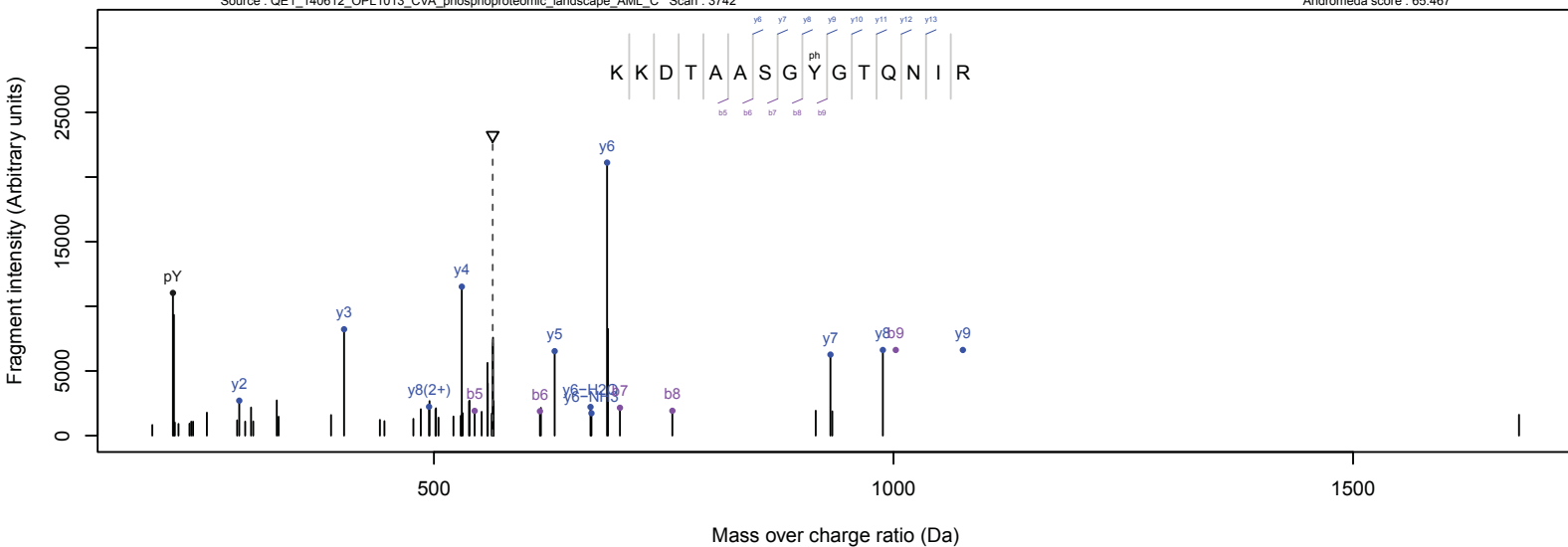
Andromeda score : 65.477



## KKDTAASGY(ph)GTQNIIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3742

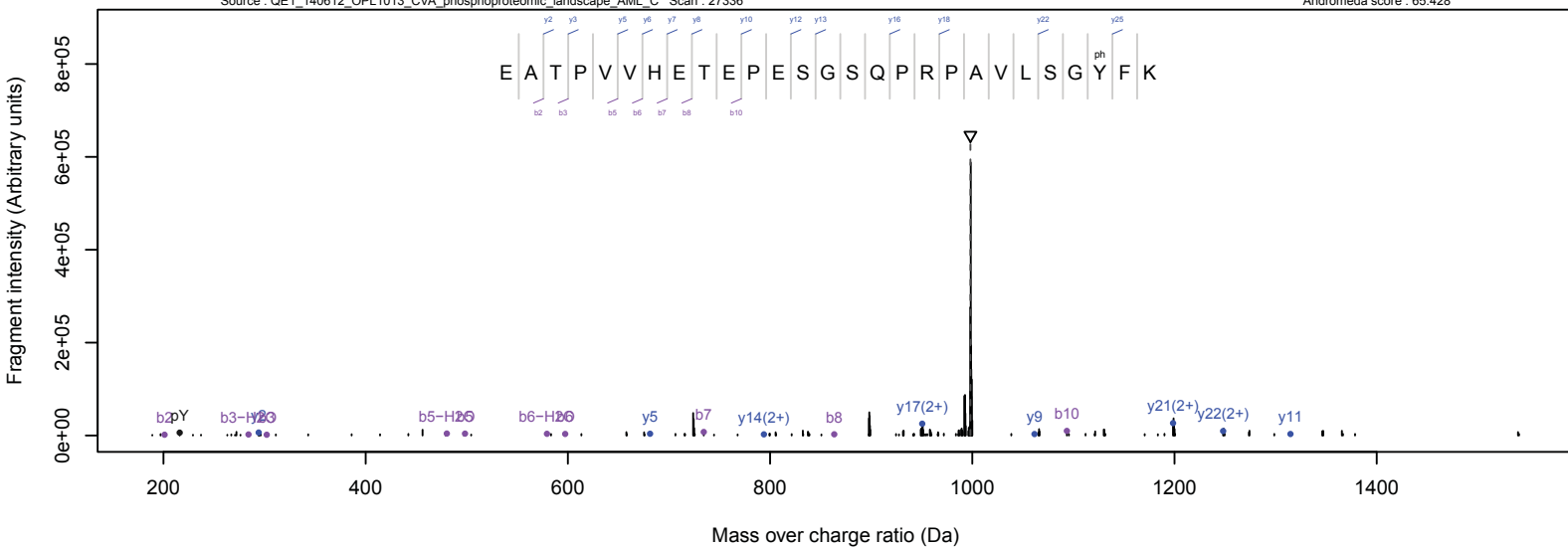
Andromeda score : 65.467



## EATPVVHETEPESGSRPAVLSGY(ph)FK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27336

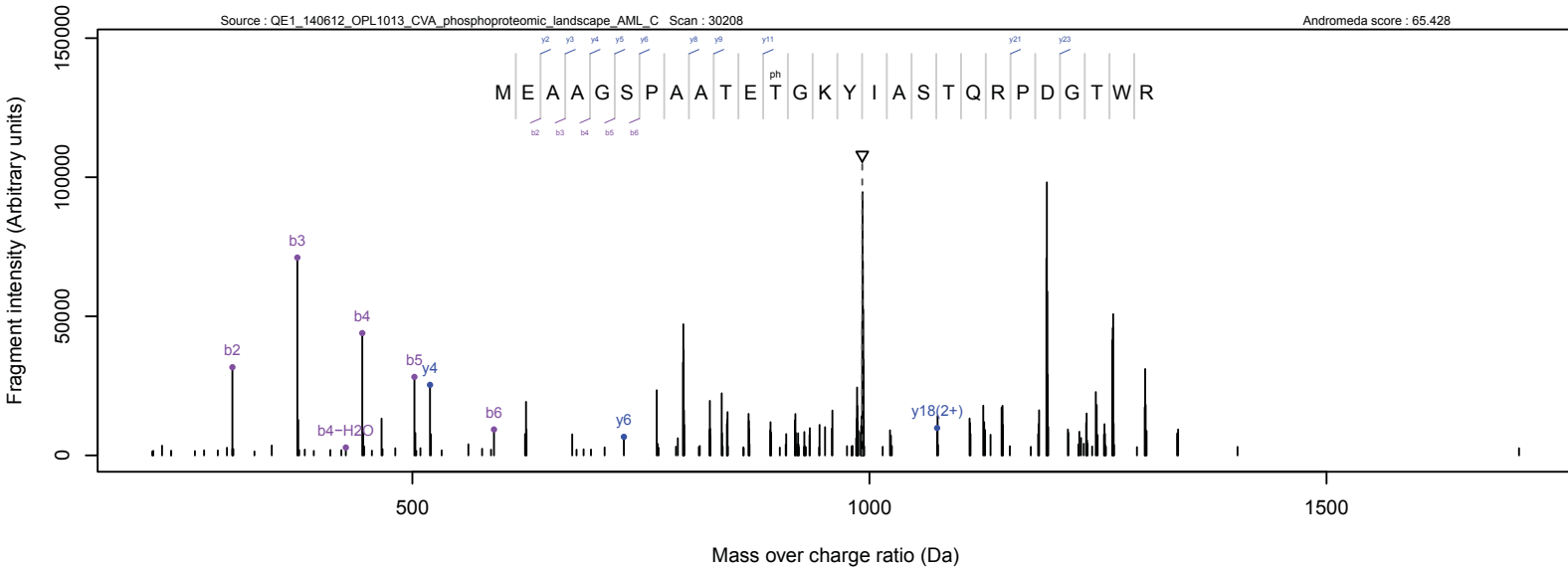
Andromeda score : 65.428



## (ac)MEAAGSPAATET(ph)GKYIASTQRPDGTWR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30208

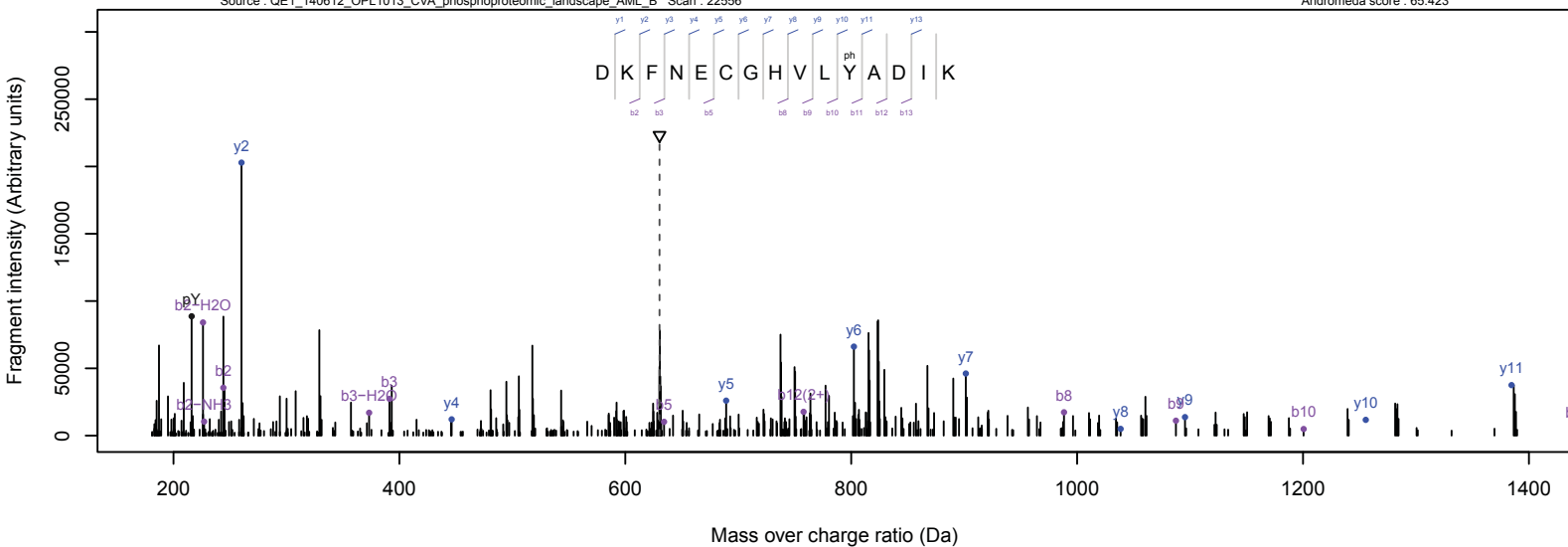
Andromeda score : 65.428



## DKFNECGHVL(ph)ADIK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 22556

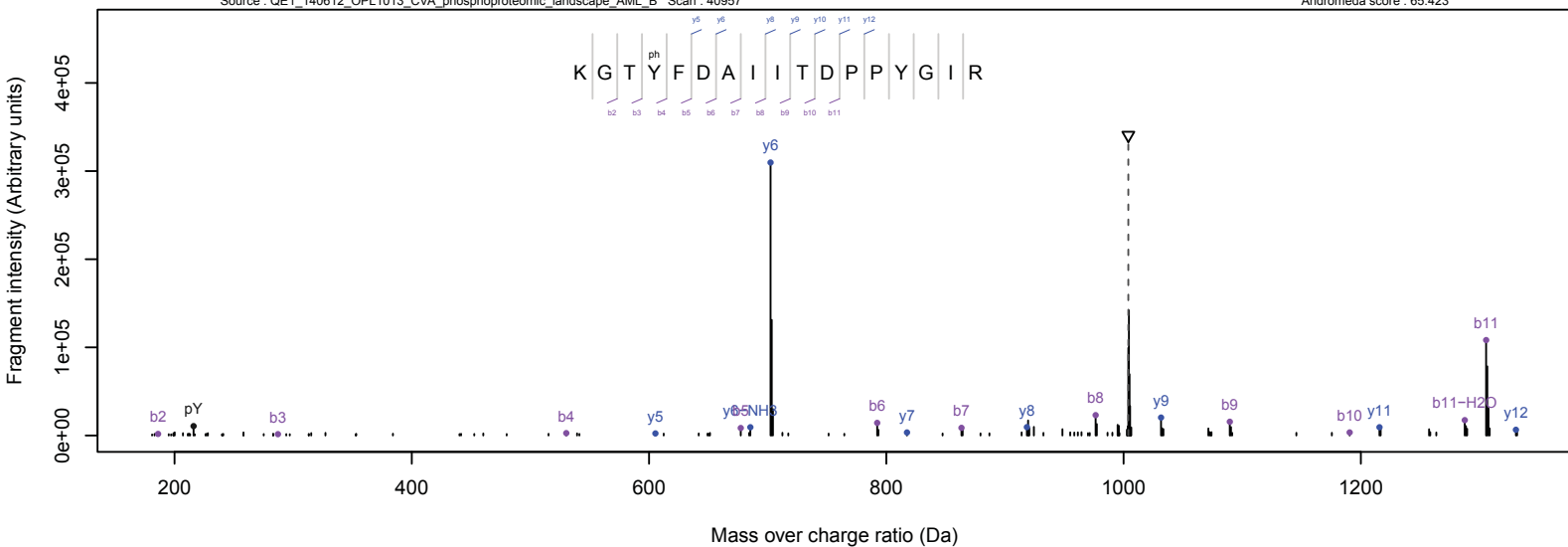
Andromeda score: 65.423



## KGTYPDAIITDPPYGI R-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 40957

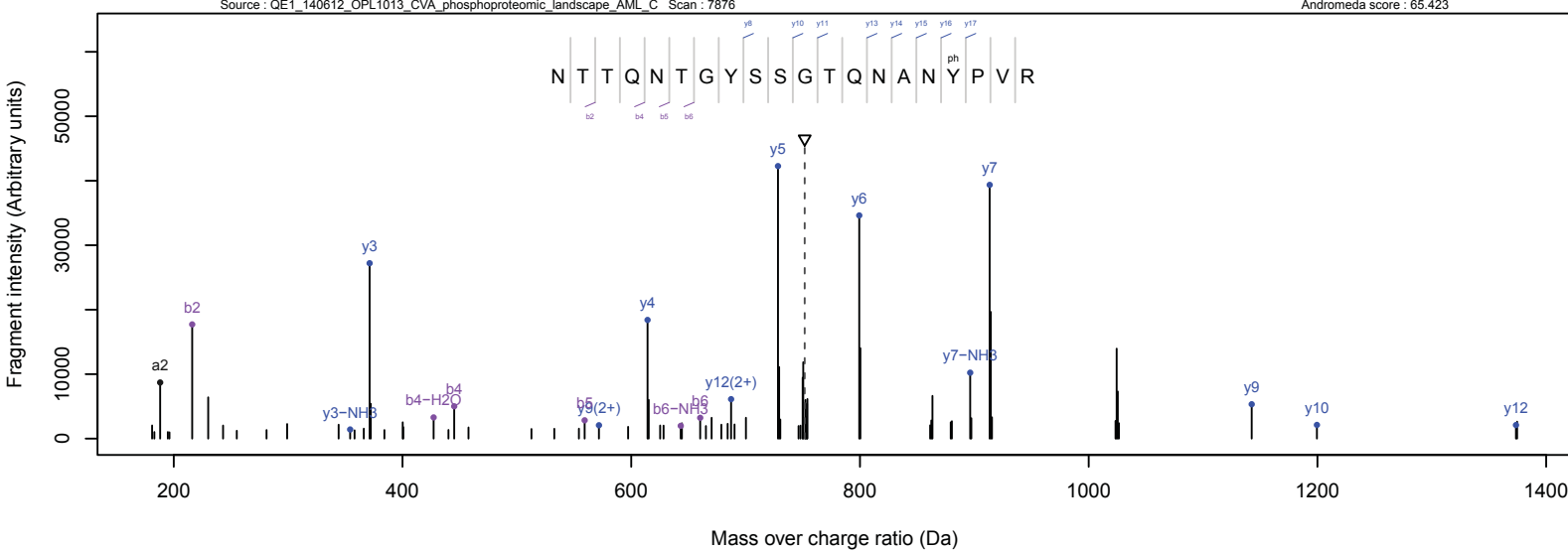
Andromeda score: 65.423



## NTTQNTGYSSGTQ NANY PVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 7876

Andromeda score: 65.423

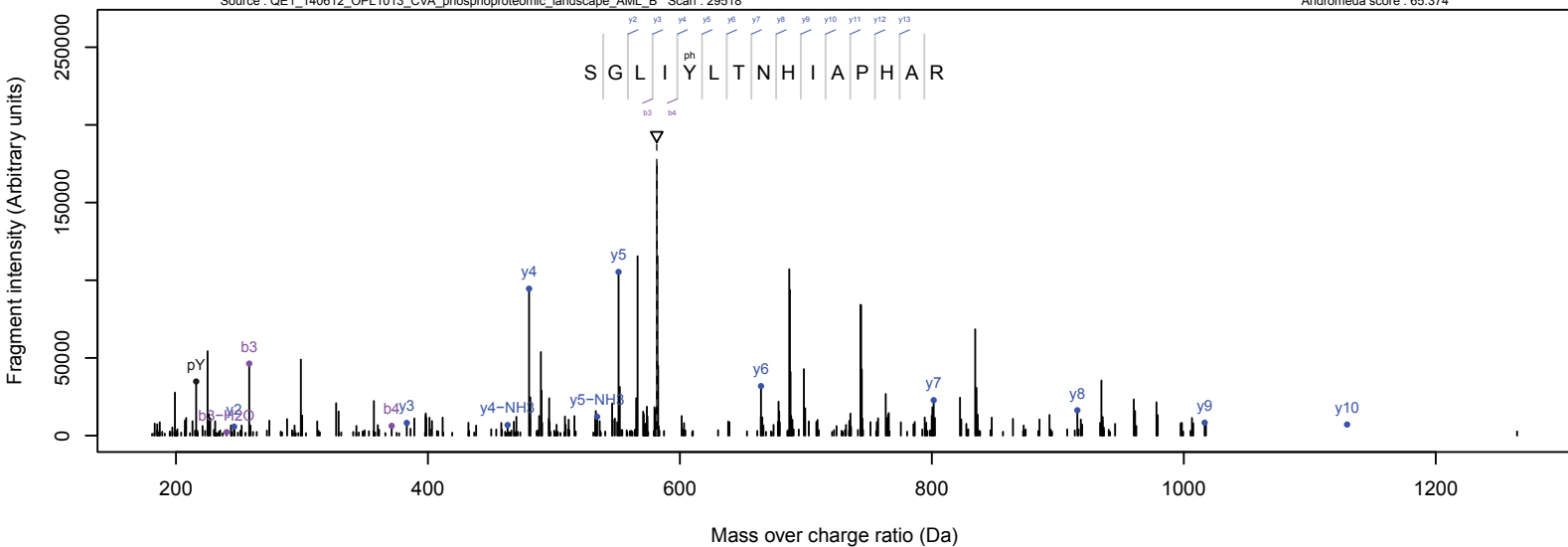




**SGLIY(ph)LTNHIAPHAR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29518

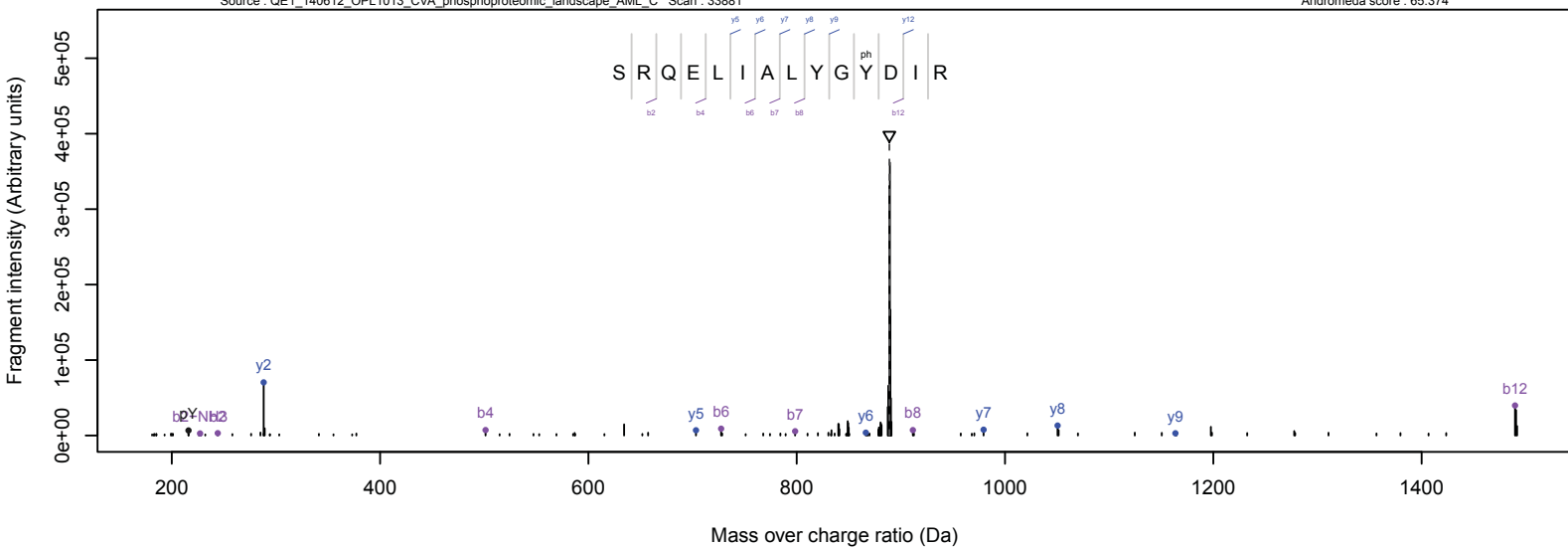
Andromeda score : 65.374



**SRQELIALYGY(ph)DIR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33881

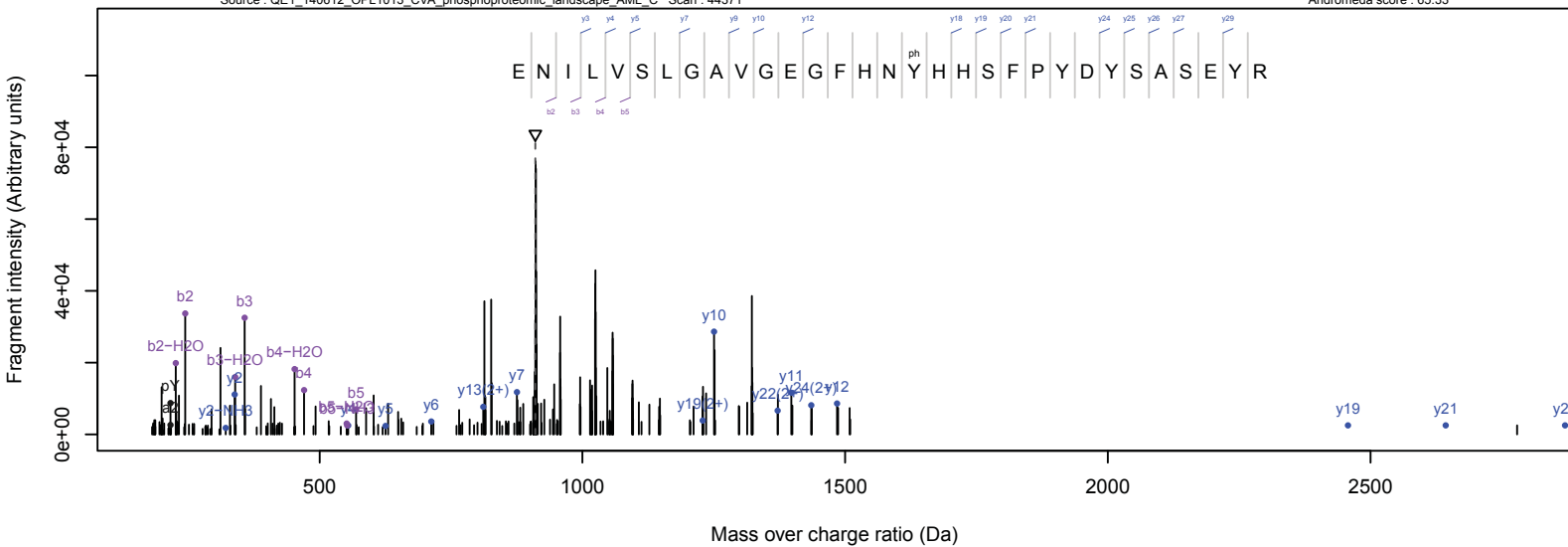
Andromeda score : 65.374



**ENILVSLGAVGEGFHNY(ph)HHSFPYDYSASEYR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44371

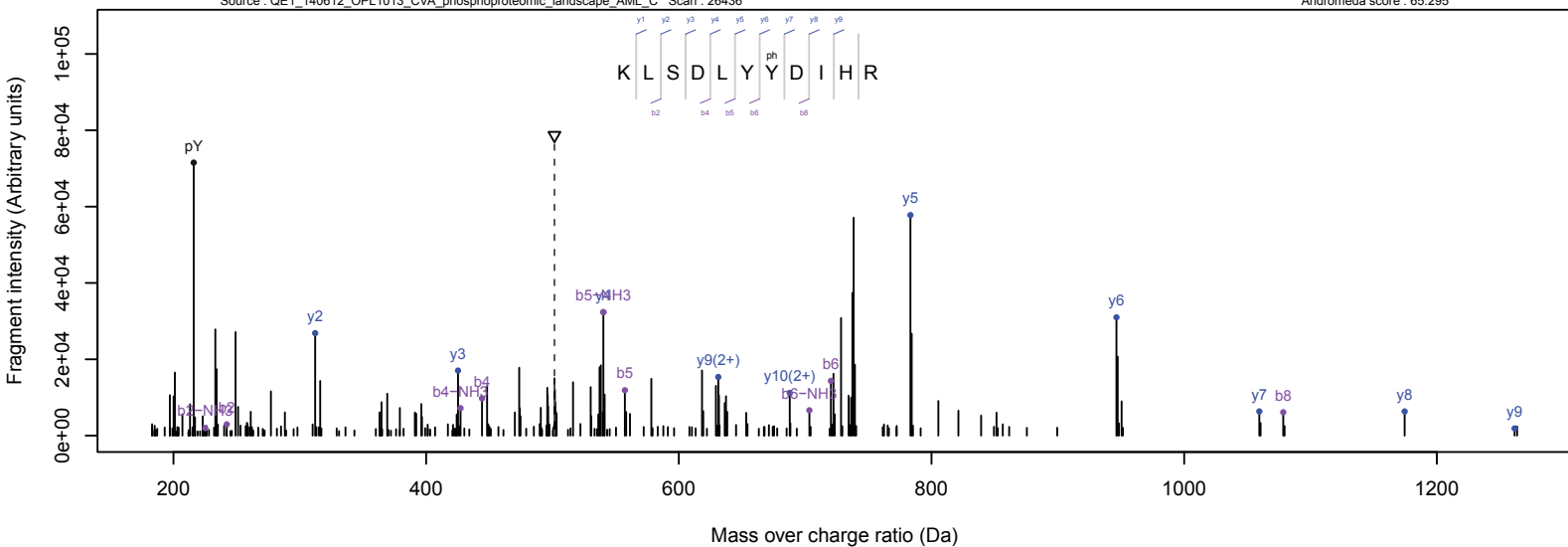
Andromeda score : 65.33



**KLSDLYY(ph)DIHR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26436

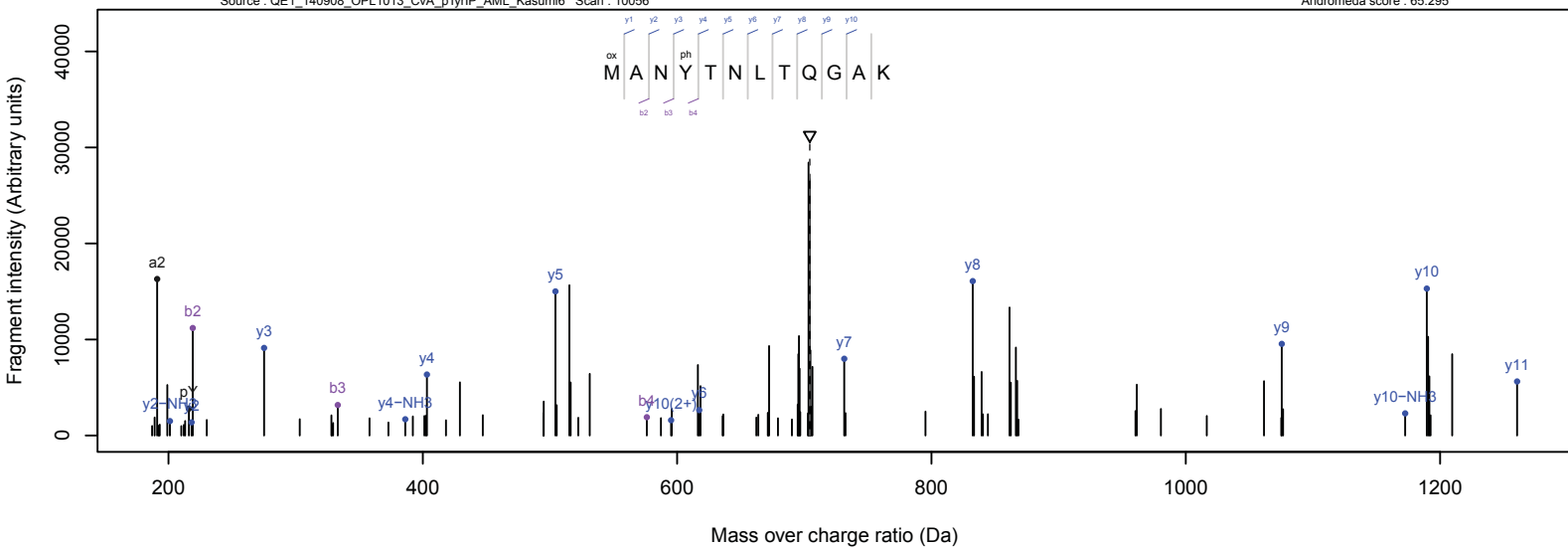
Andromeda score : 65.295



**M(ox)ANY(ph)TNLTQGAK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 10056

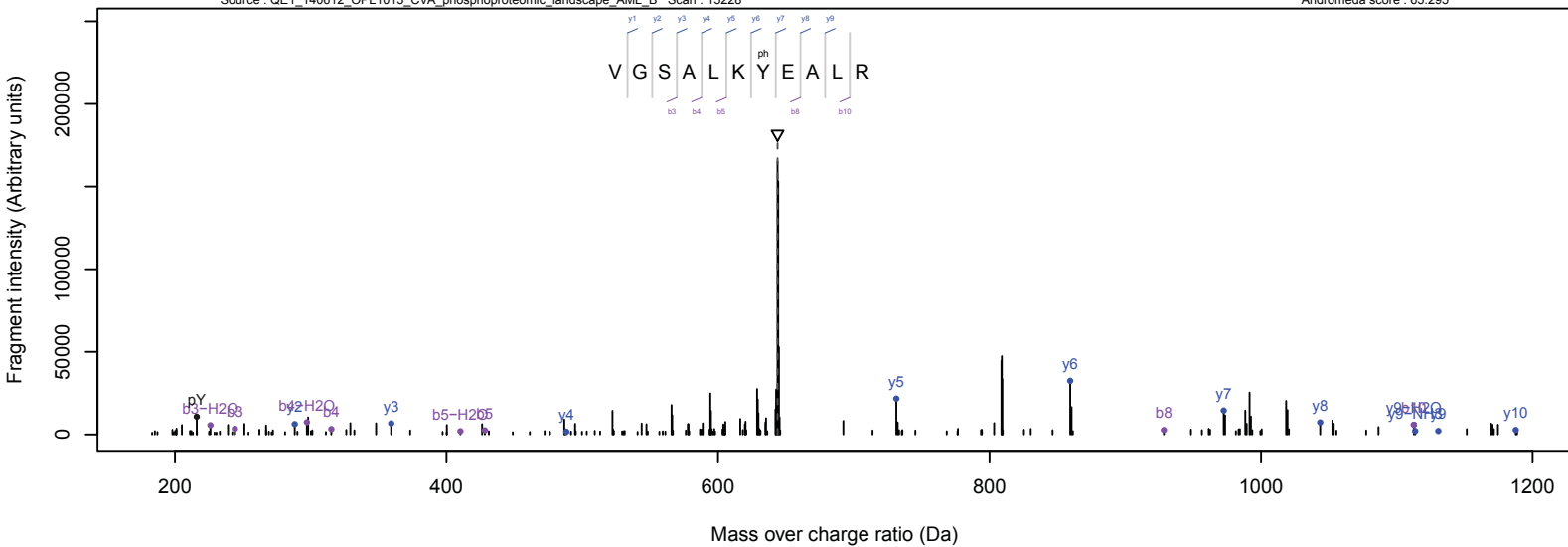
Andromeda score : 65.295

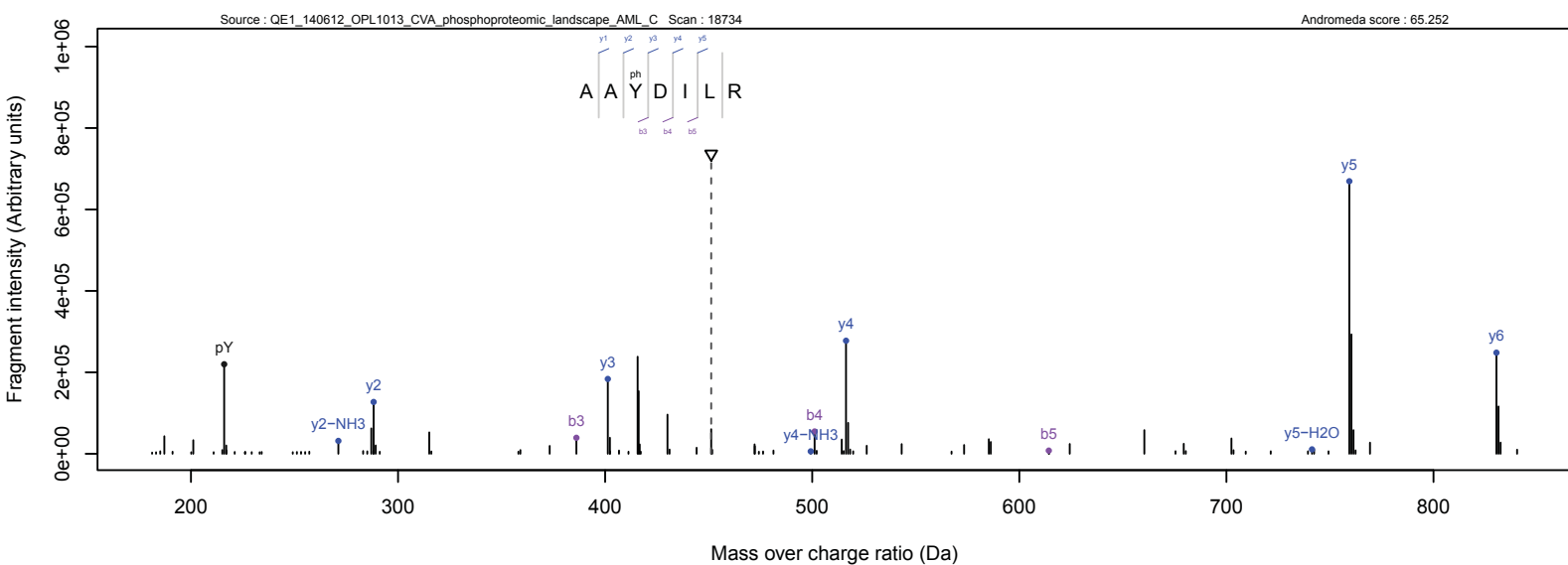


**VGSALKY(ph)EALR-2+**

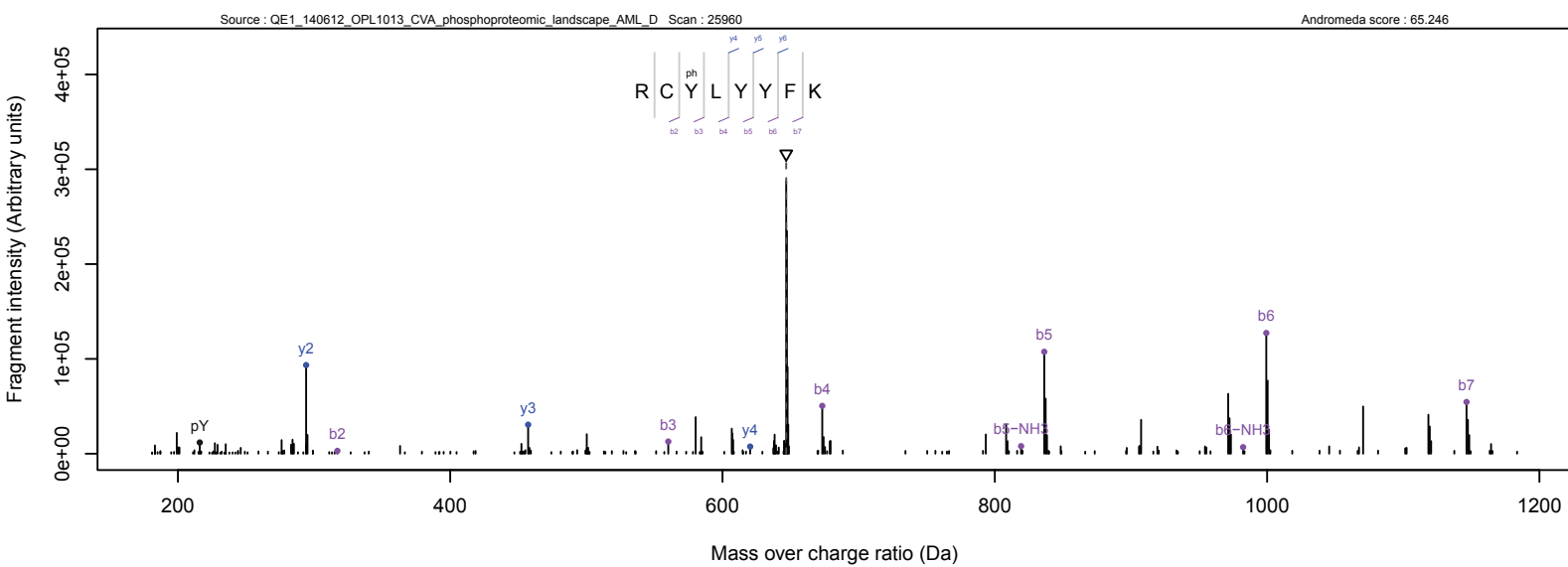
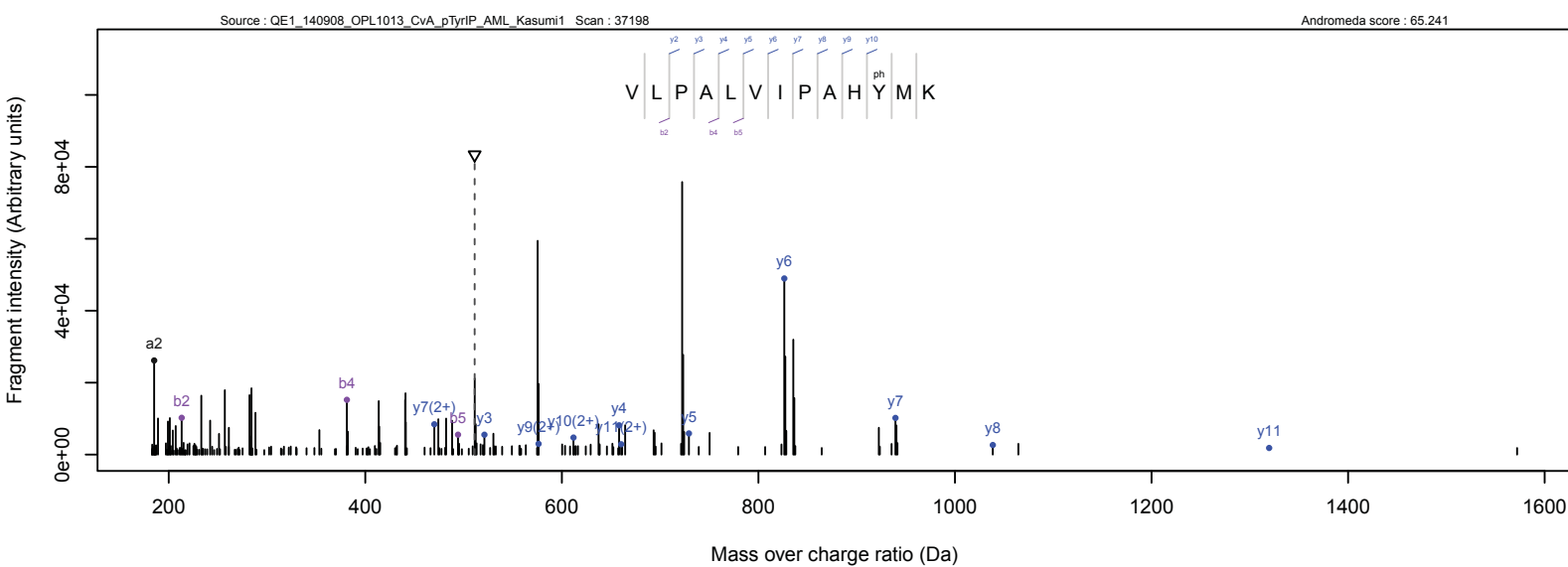
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15228

Andromeda score : 65.295



AA<sub>Y</sub>(ph)DILR-2+

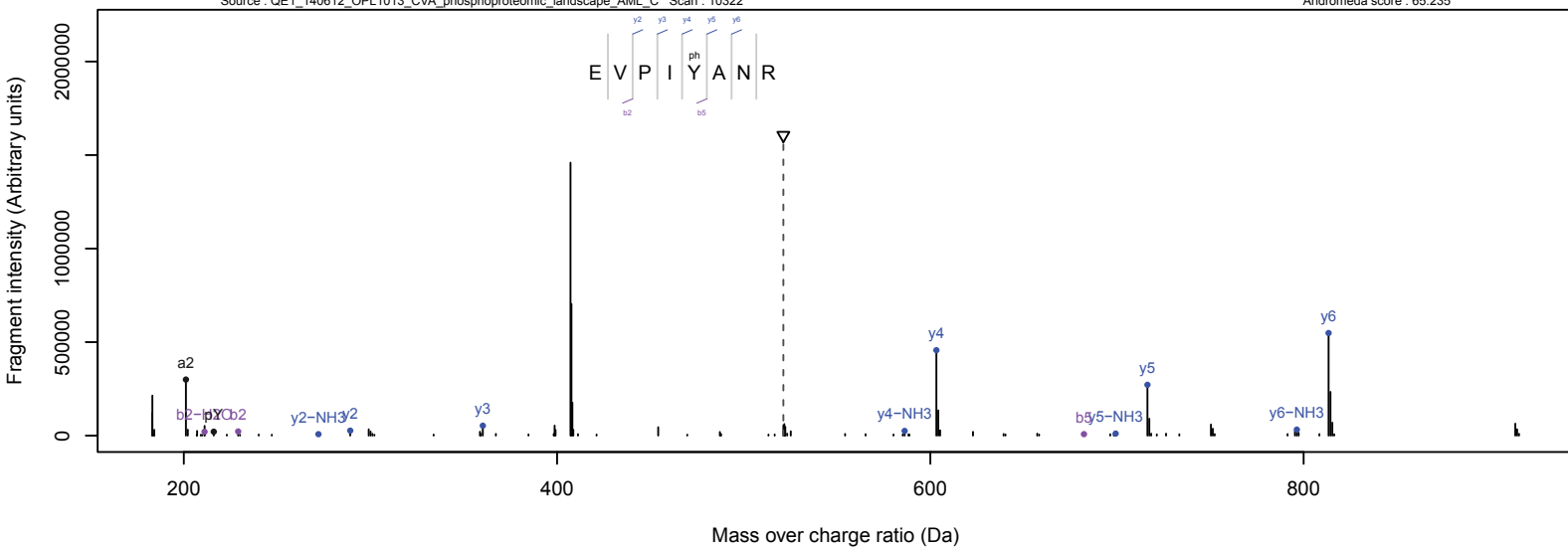
## RCY(ph)LYYFK-2+

VLPALVIPA<sub>HY</sub>(ph)MK-3+

## EVPIY(ph)ANR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10322

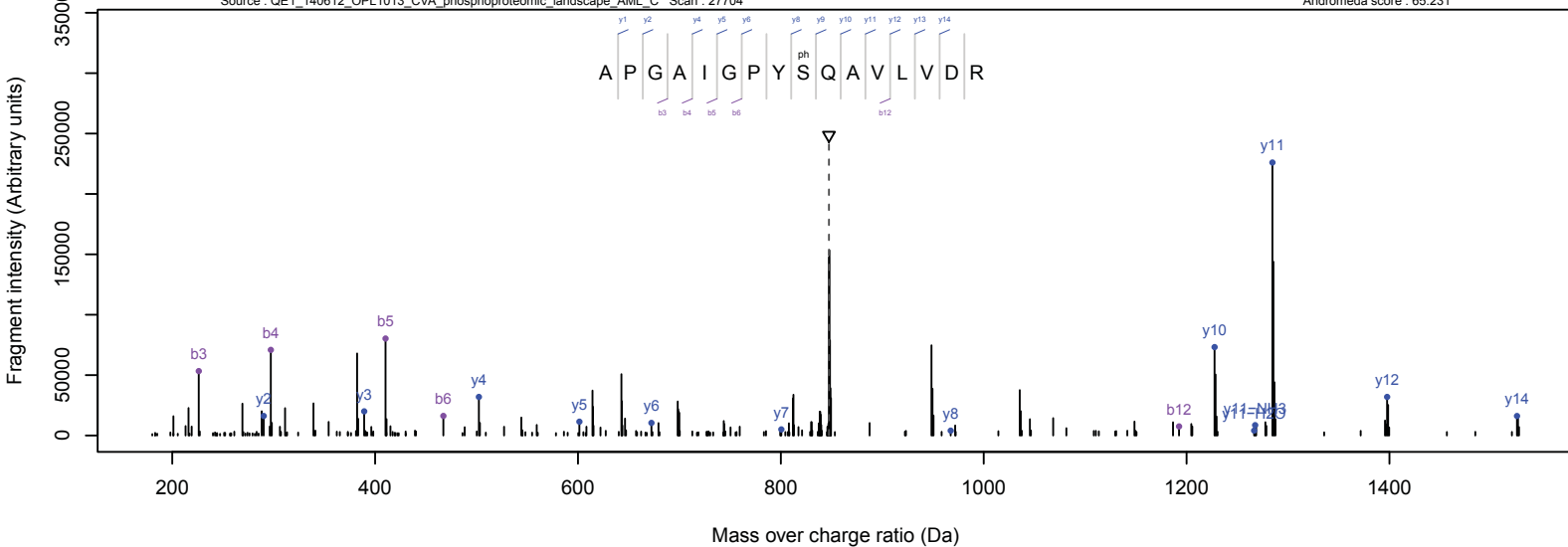
Andromeda score : 65.235



## APGAIGPYS(ph)QAVLVDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27704

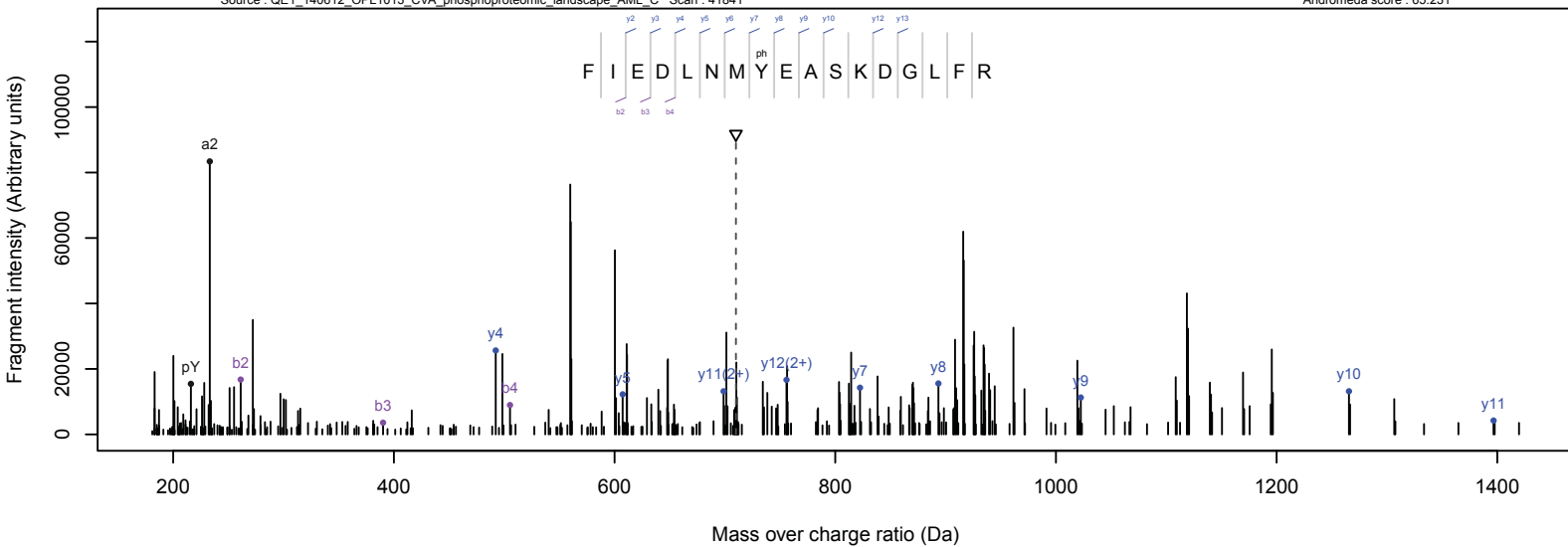
Andromeda score : 65.231



## FIEDLNMY(ph)EASKDGLFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41841

Andromeda score : 65.231

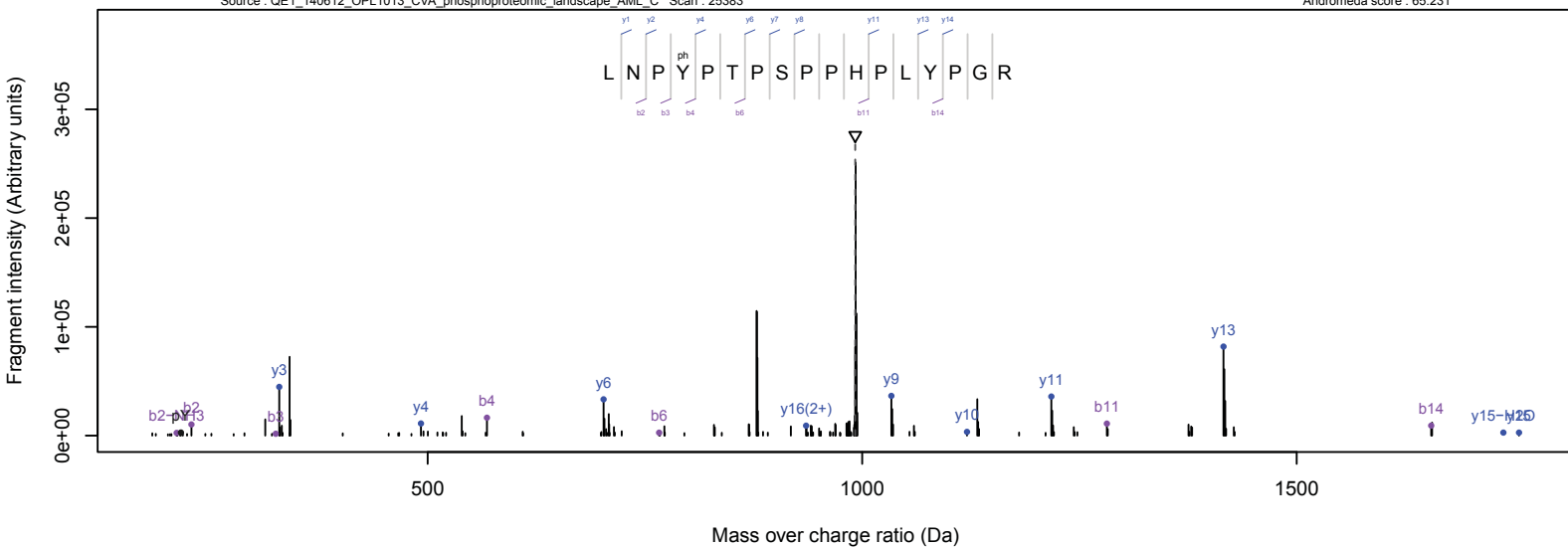




## LNPY(ph)PTSPHPLYPGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25383

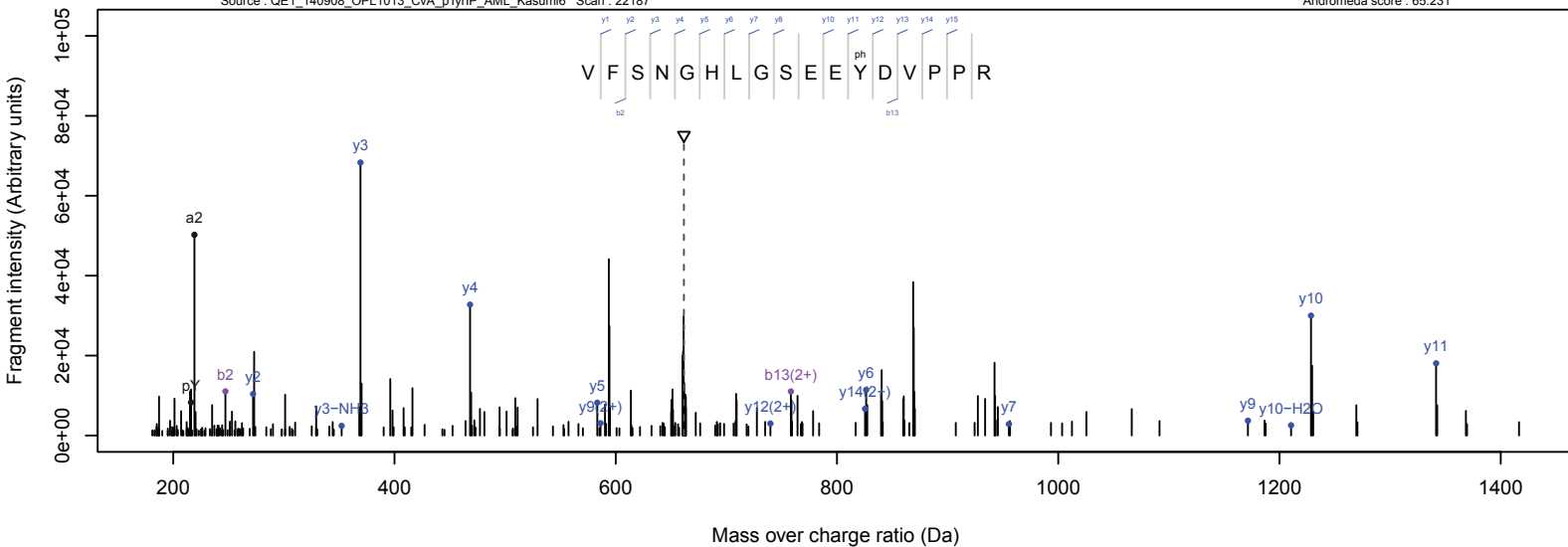
Andromeda score: 65.231



## VFSNGHLGSEE(ph)DVPPR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 22187

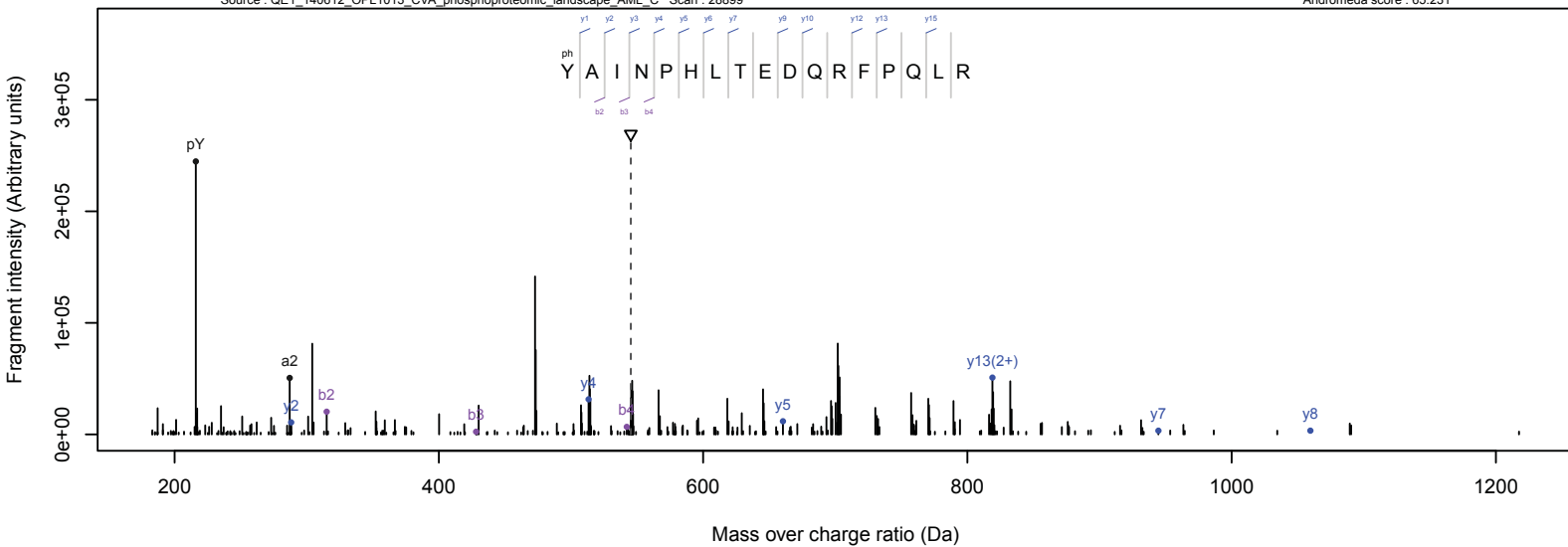
Andromeda score: 65.231



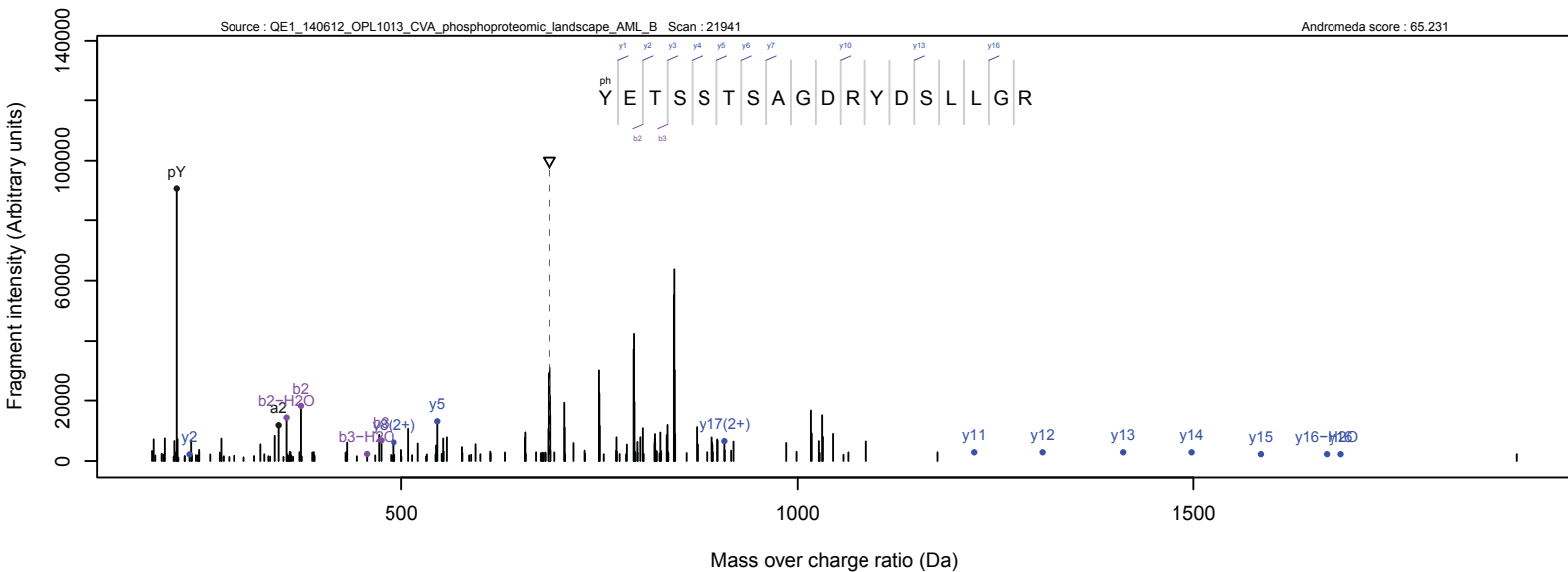
## Y(ph)AINPHLTDQRFPQLR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 28899

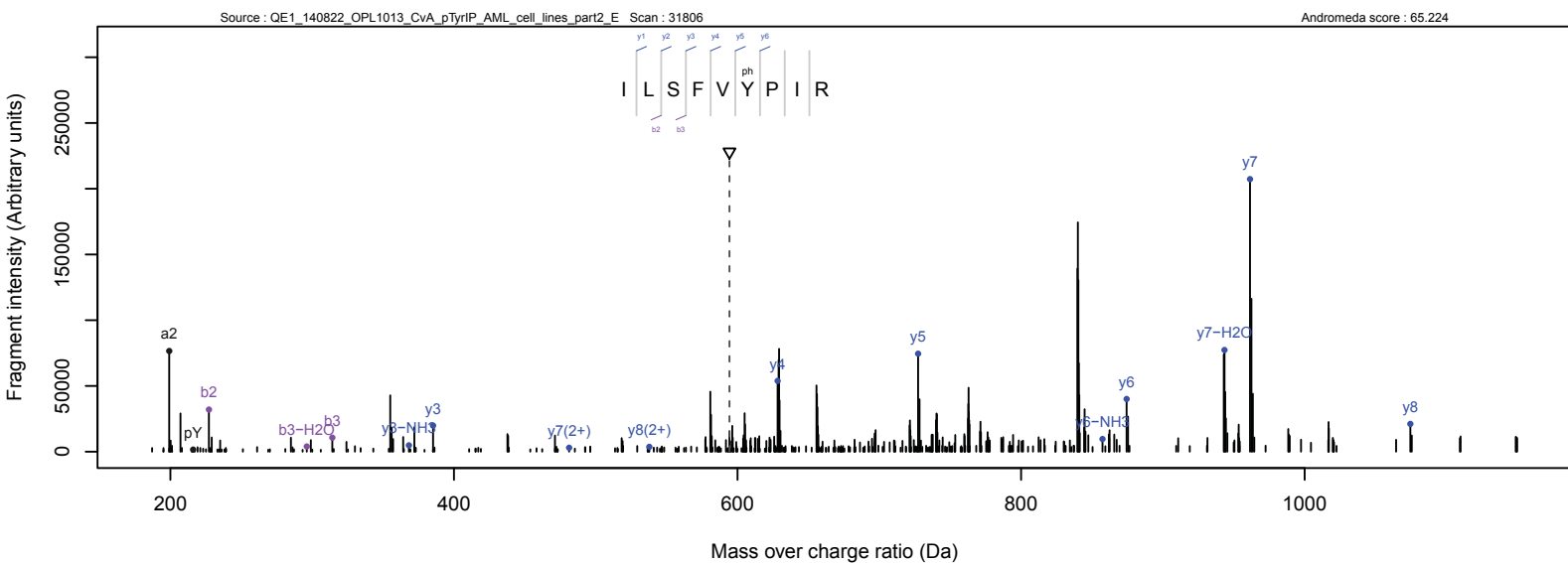
Andromeda score: 65.231



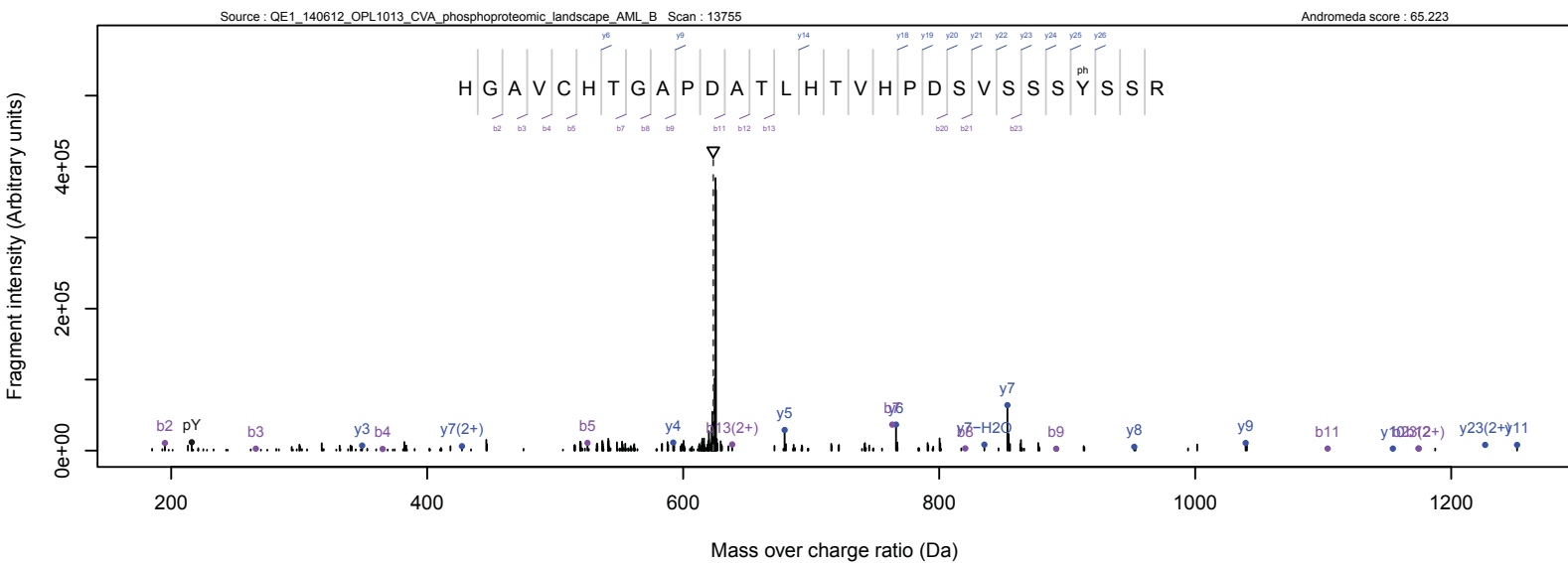
Y(ph)ETSSTSAGDRYDSLGR-3+



ILSFVY(ph)PIR-2+



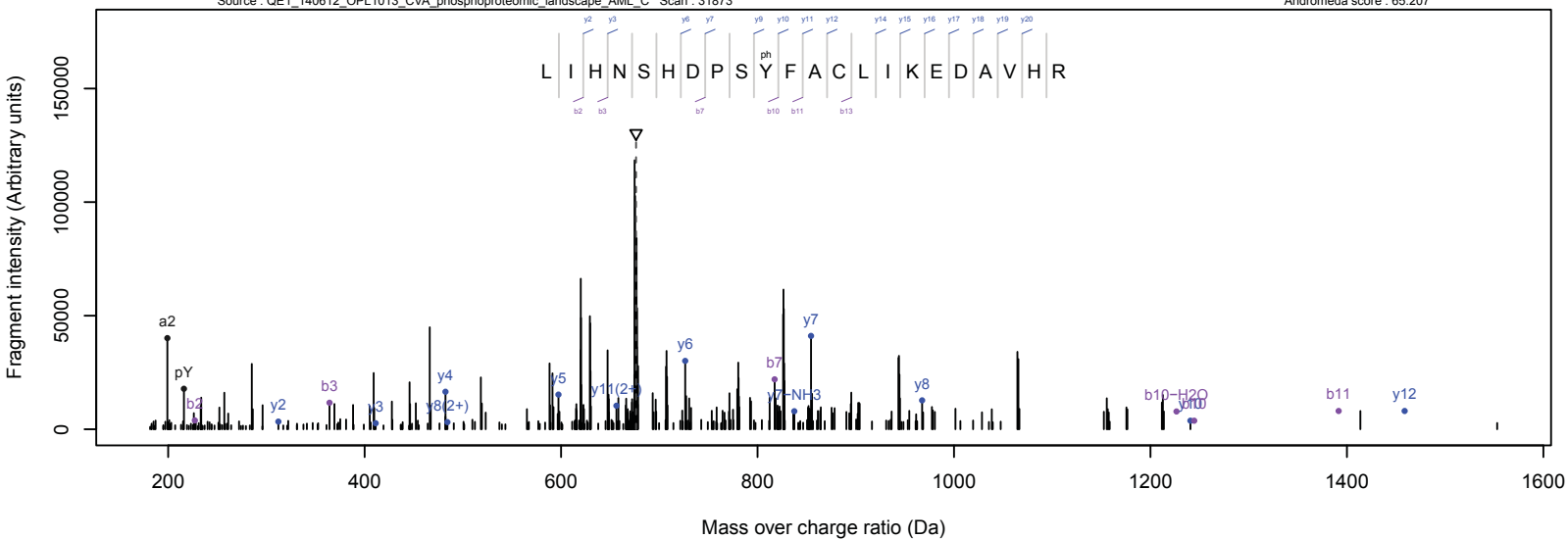
HGAVCHTGAPDATHVHPDVSSSSY(ph)SSR-5+



LIHNSHDPSY(ph)FACLIKEDAVHR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31873

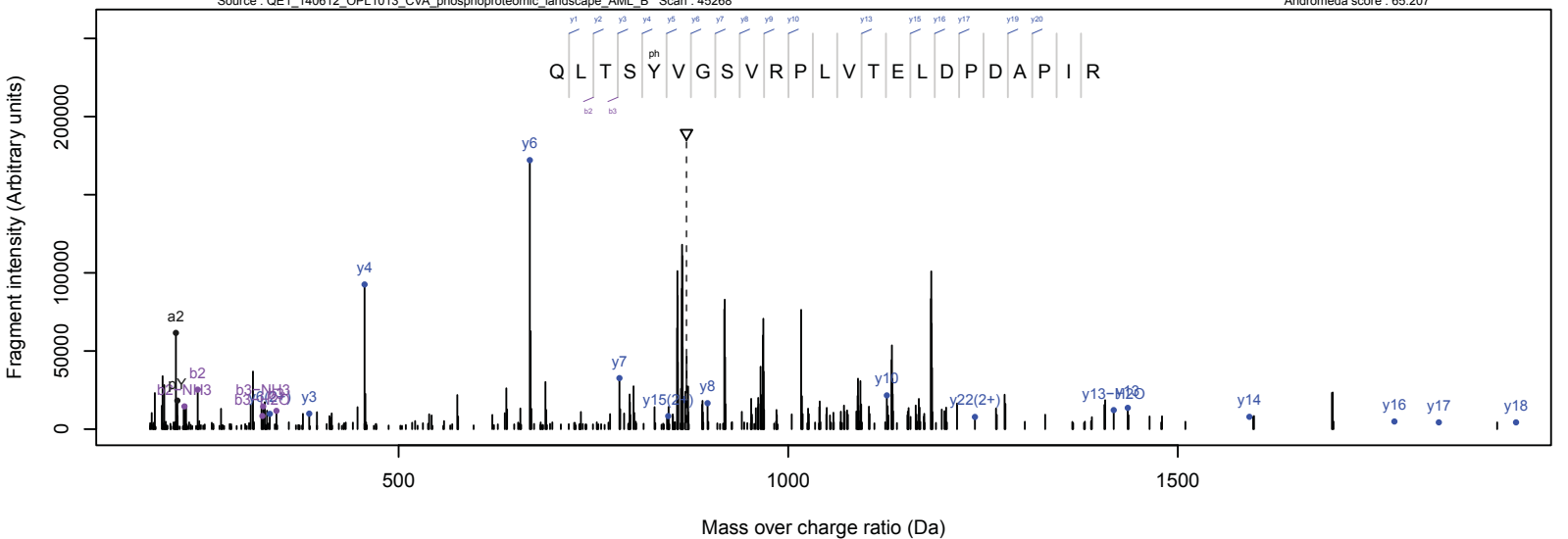
Andromeda score: 65.207



QLTSY(ph)VGSVRPLVTELPDAPIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 45268

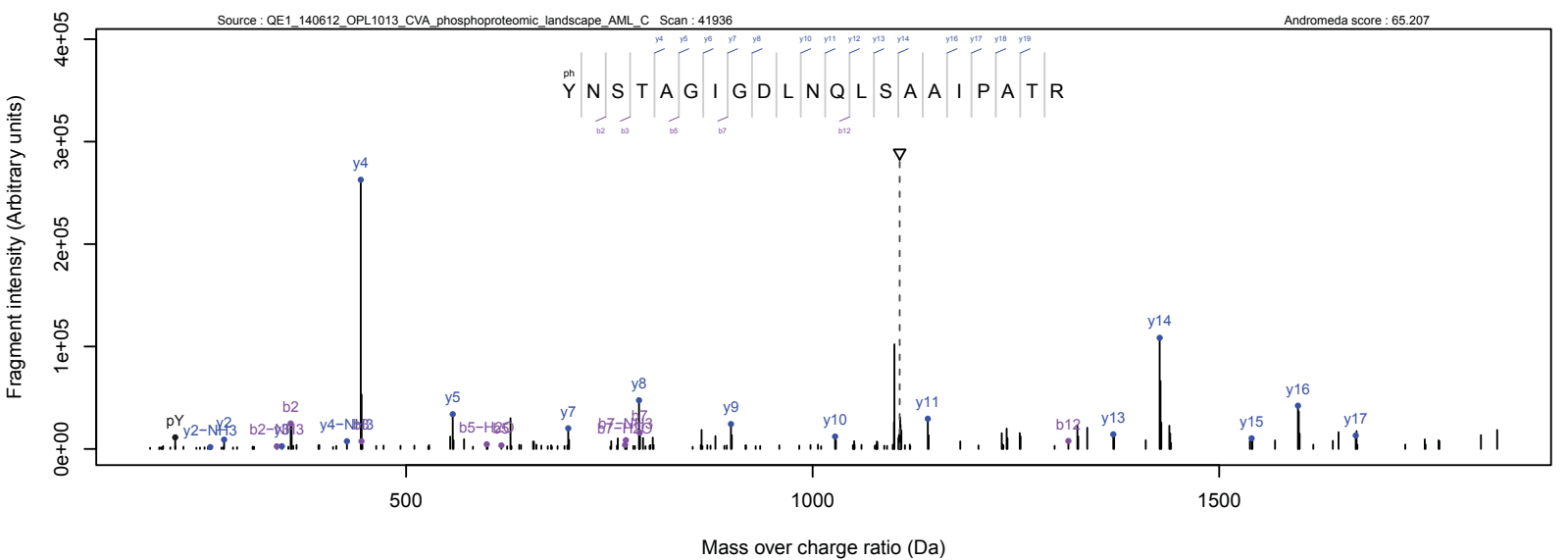
Andromeda score: 65.207



Y(ph)NSTAGIDLNQLSAAIPATR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41936

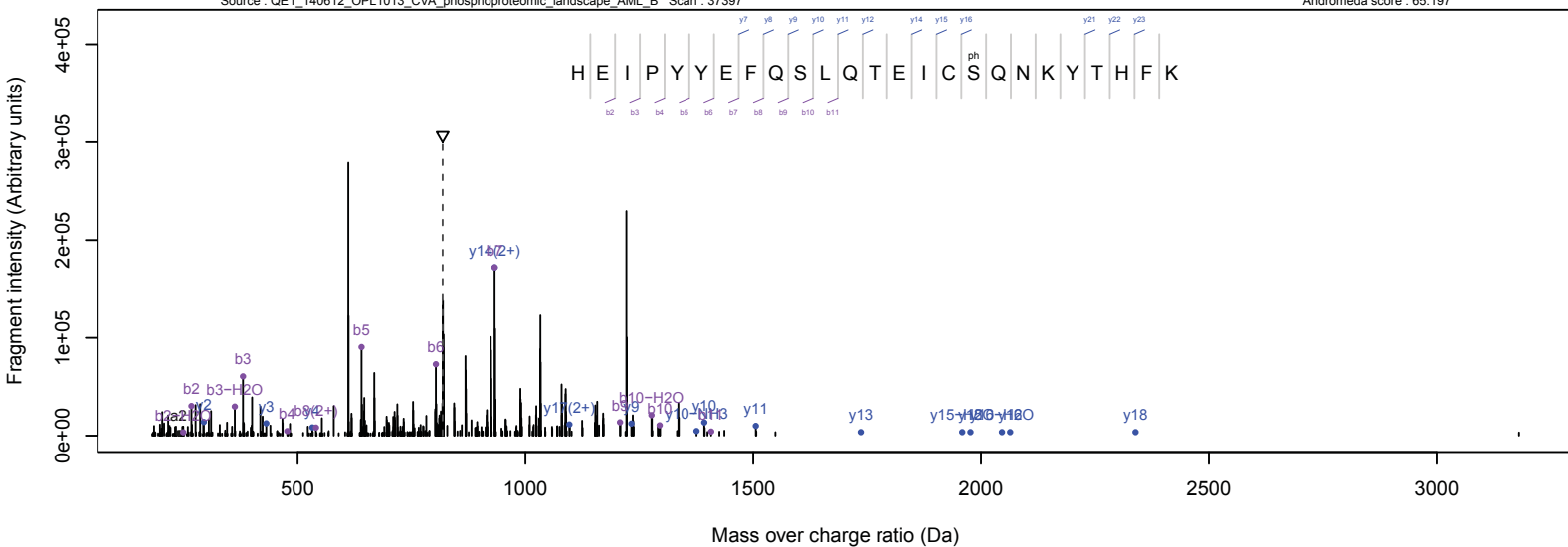
Andromeda score: 65.207



HEIPYYEFQSLQTEIC<sup>ph</sup>QNKYTHFK-4+

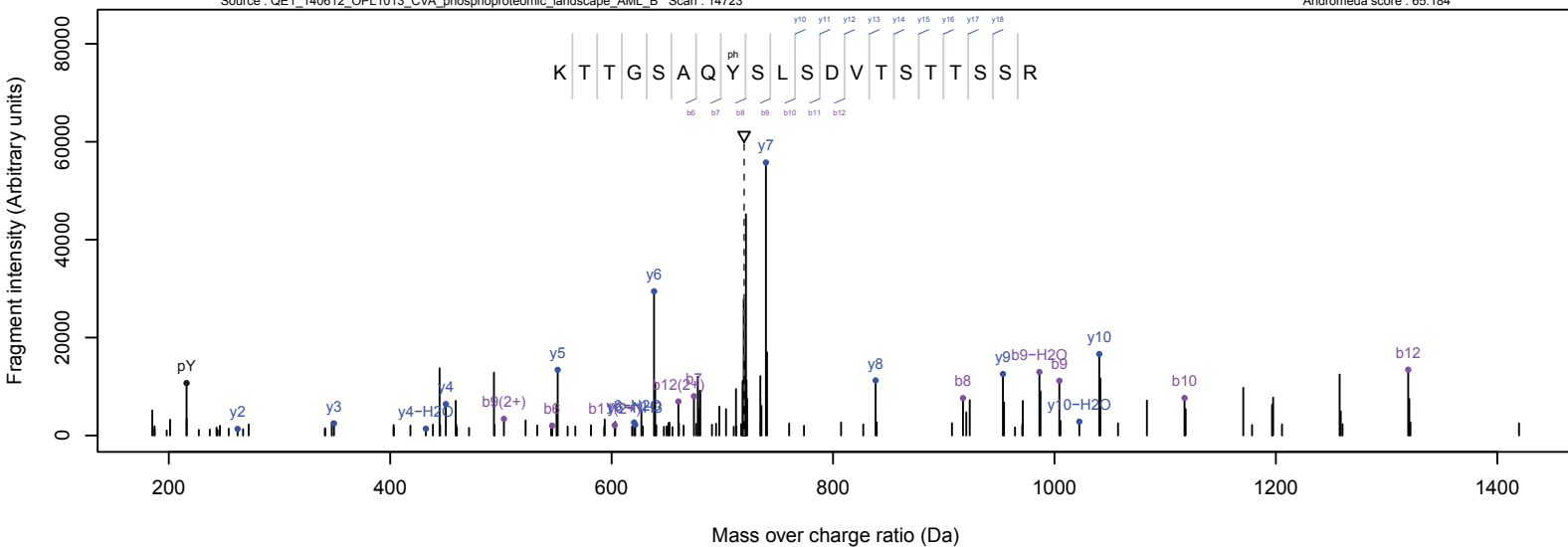
Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 37397

Andromeda score: 65.197

KTTGSAQY<sup>ph</sup>SLSDVTSTSSR-3+

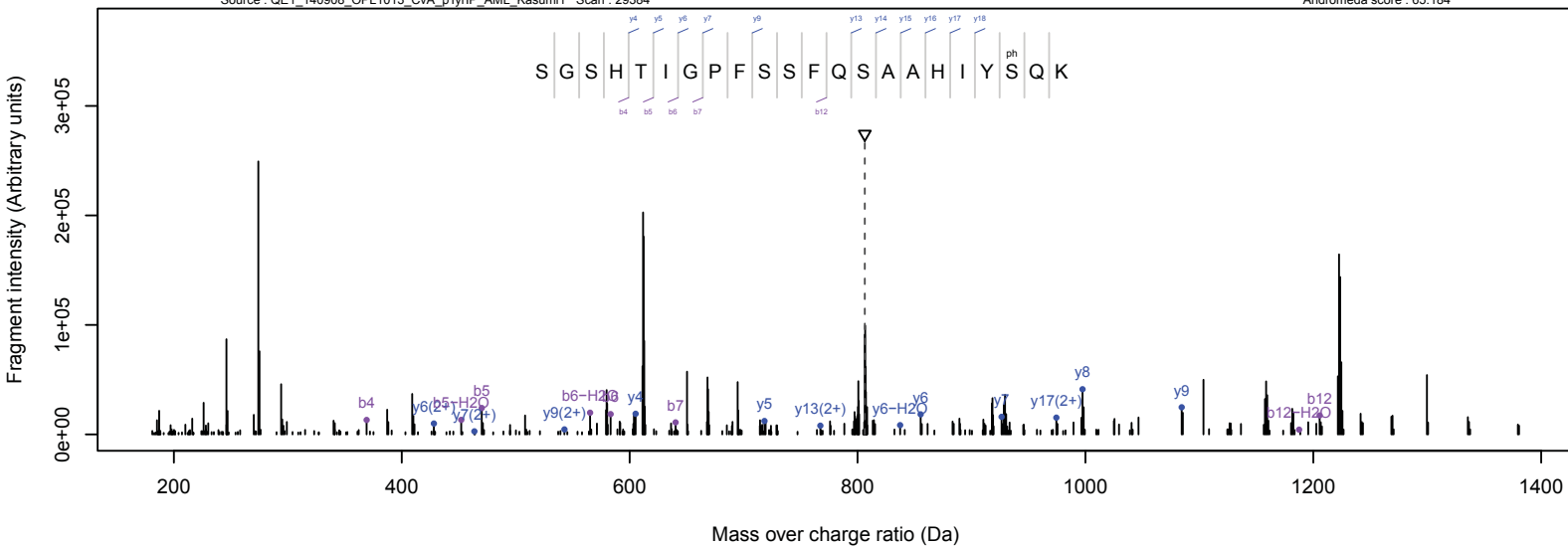
Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 14723

Andromeda score: 65.184

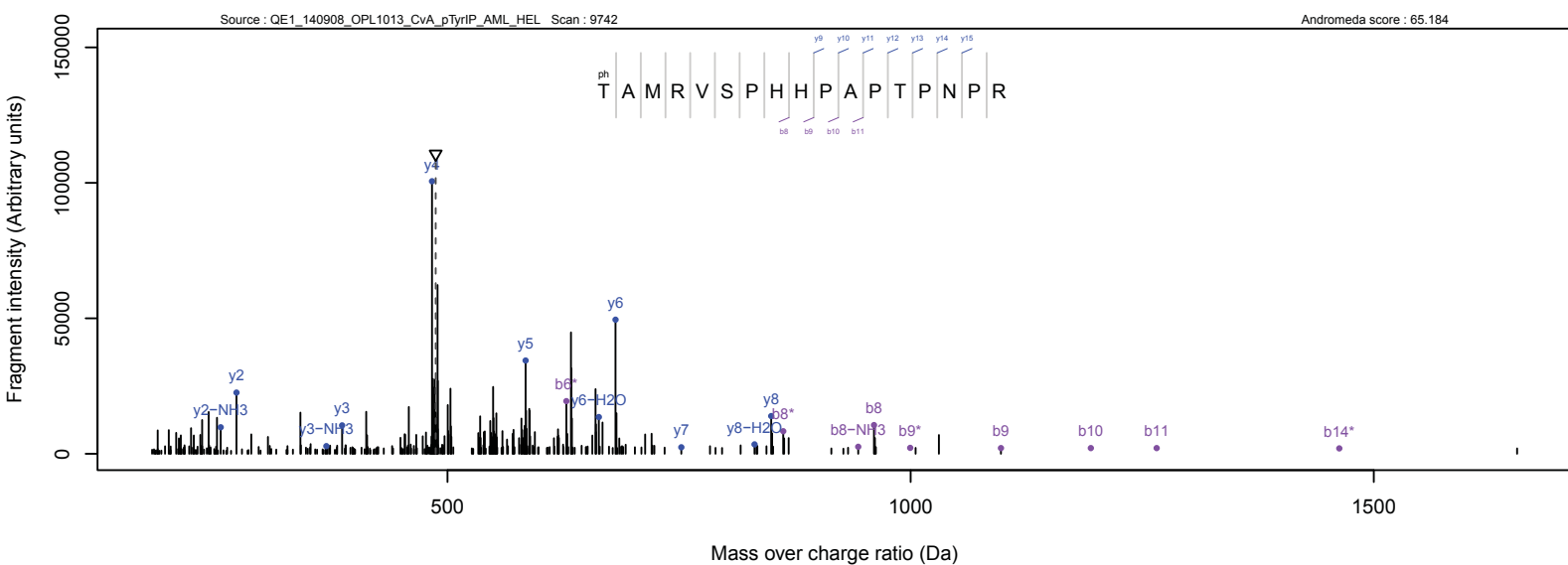
SGSHTIGPFSS<sup>ph</sup>QSAAHYISQK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 29384

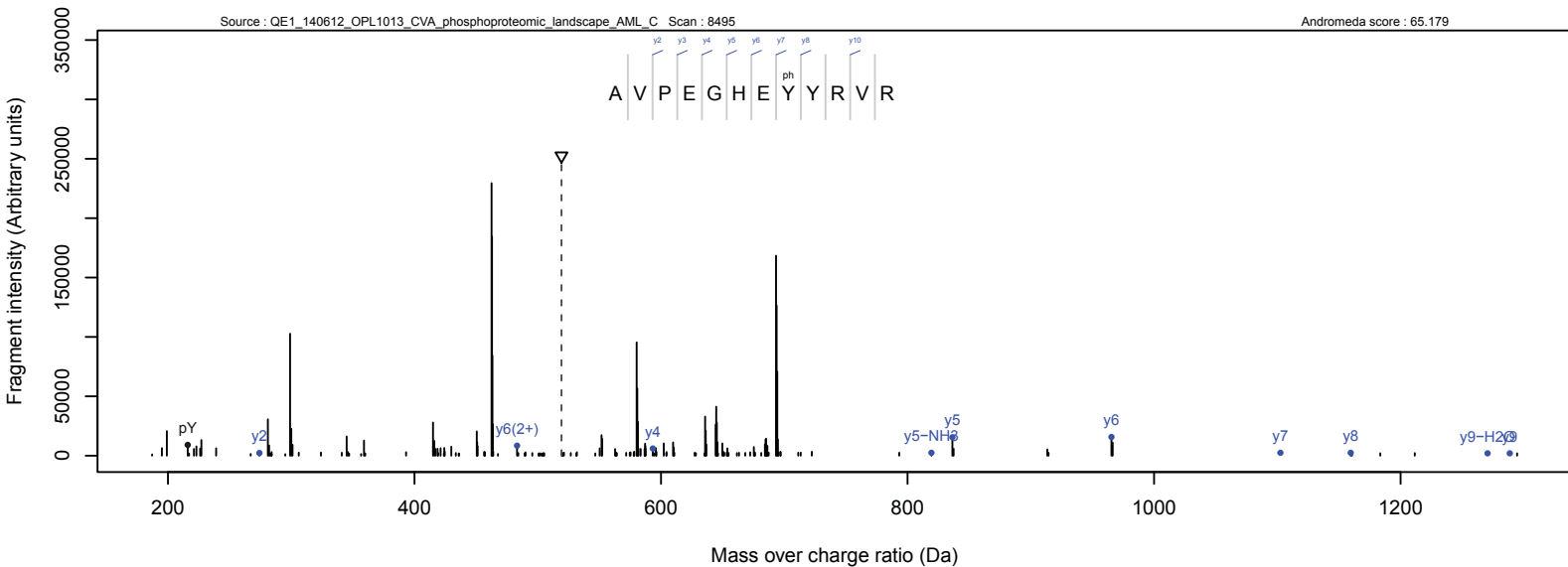
Andromeda score: 65.184



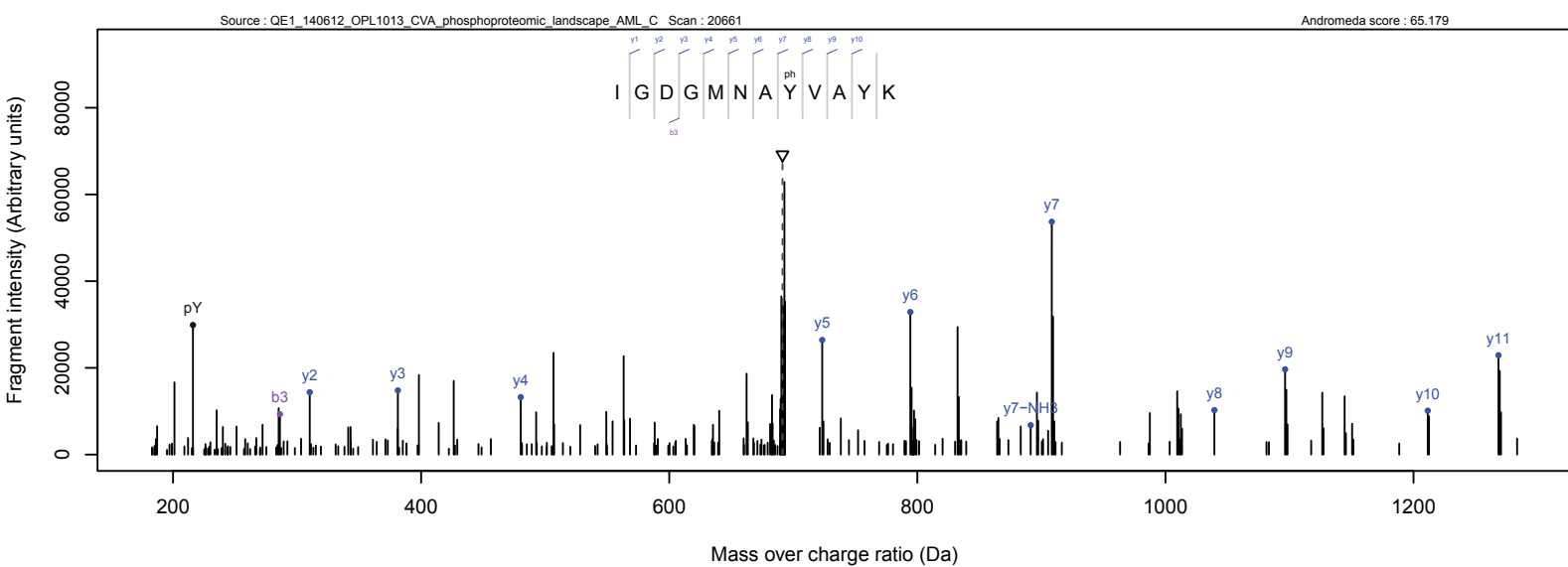
## T(ph)AMRVSPHHPAPTNP-4+



## AVPEGHEY(ph)YRVR-3+



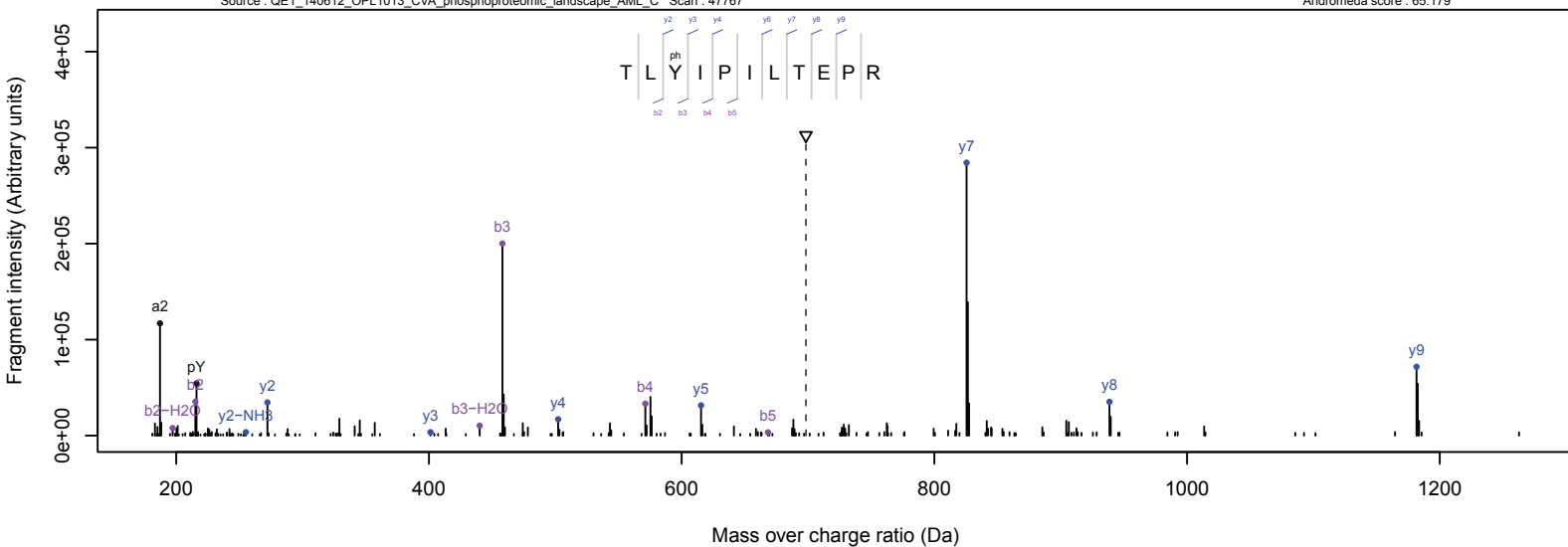
## IGDGMNAY(ph)VAYK-2+



**TLY(ph)IPILTEPR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 47767

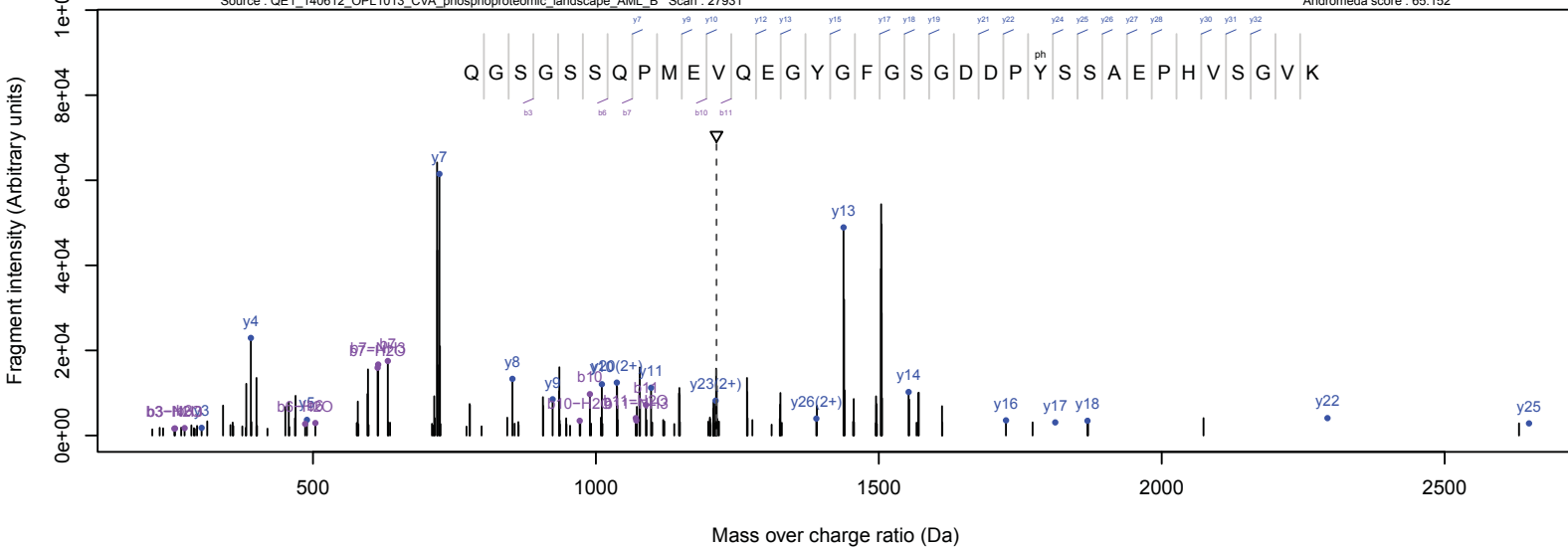
Andromeda score: 65.179



**QGSQSSQPMEVQEGYFGSGDDPY(ph)SSAEPHVSGVK-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 27931

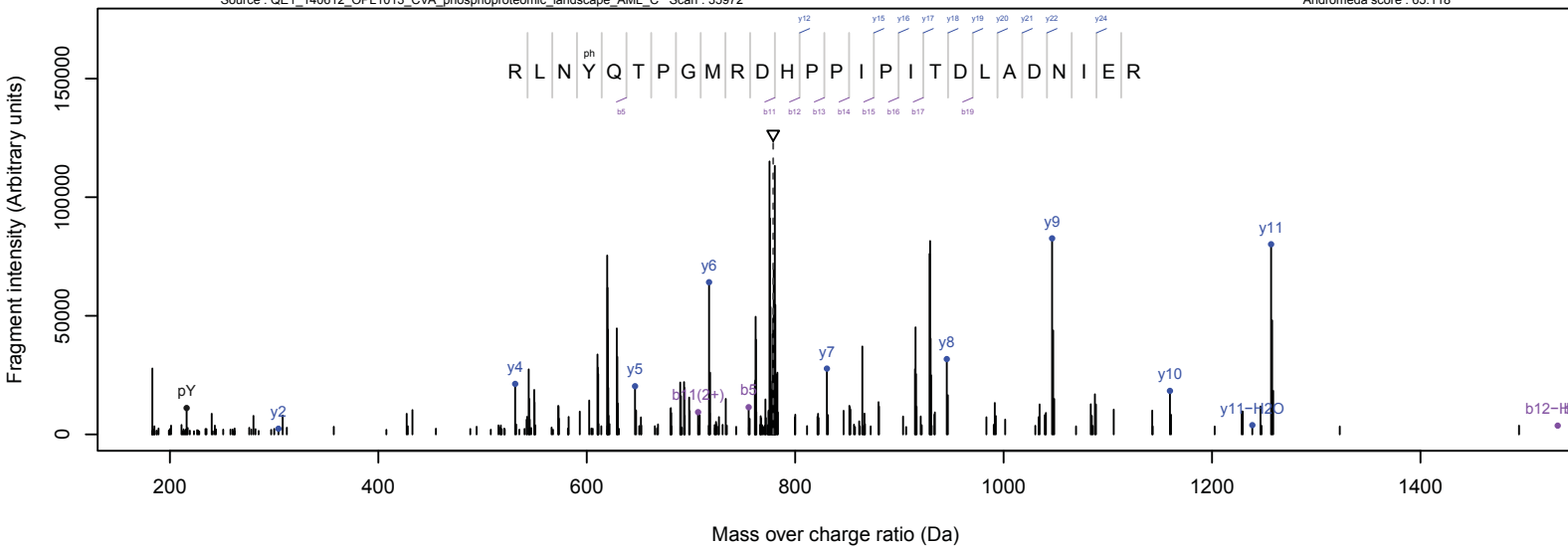
Andromeda score: 65.152



**RLNY(ph)QTPGMRDHPPIIPITDLADNIER-4+**

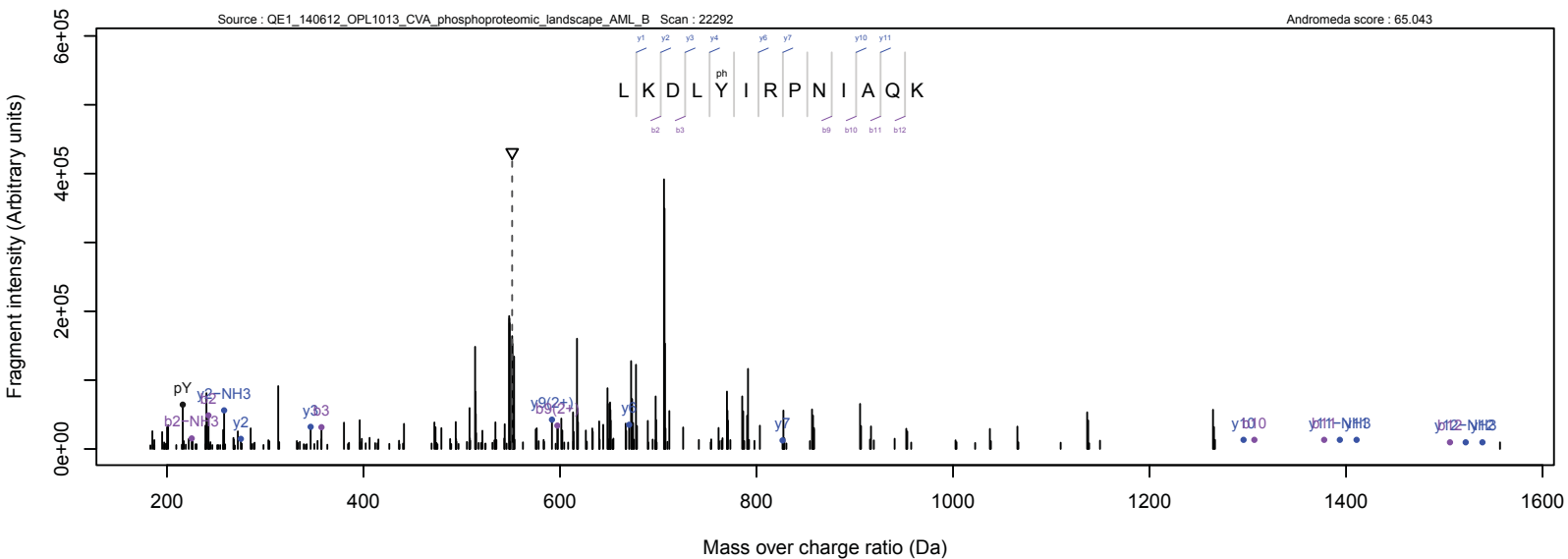
Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 35972

Andromeda score: 65.118

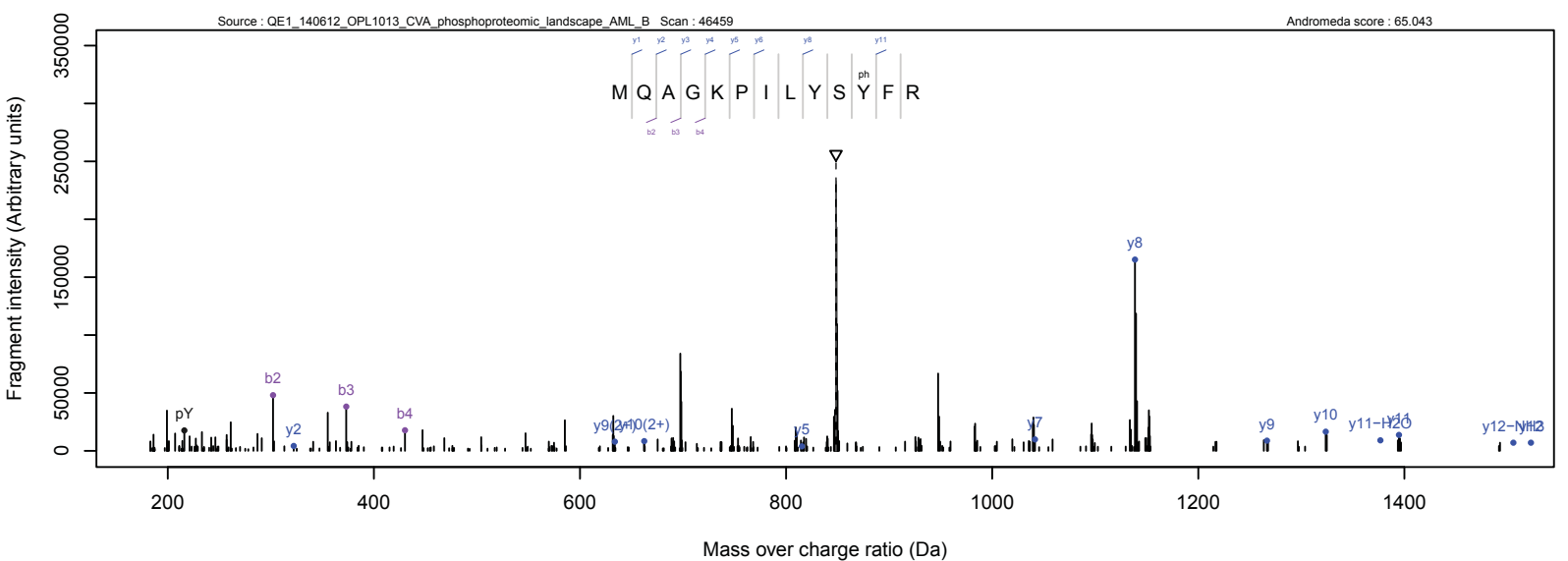




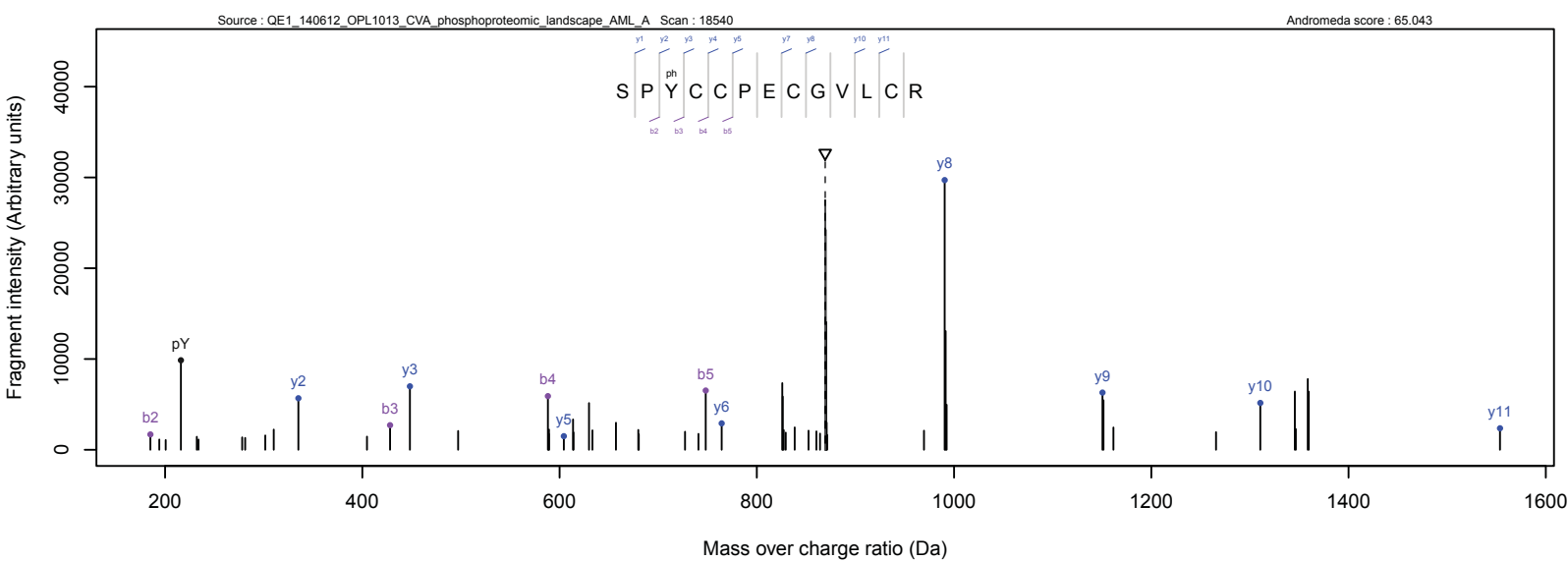
LKDLY(ph)IRPNIAQK-3+



(ac)MQAGKPILYS(ph)YFR-2+



SPY(ph)CCPEGVLCR-2+



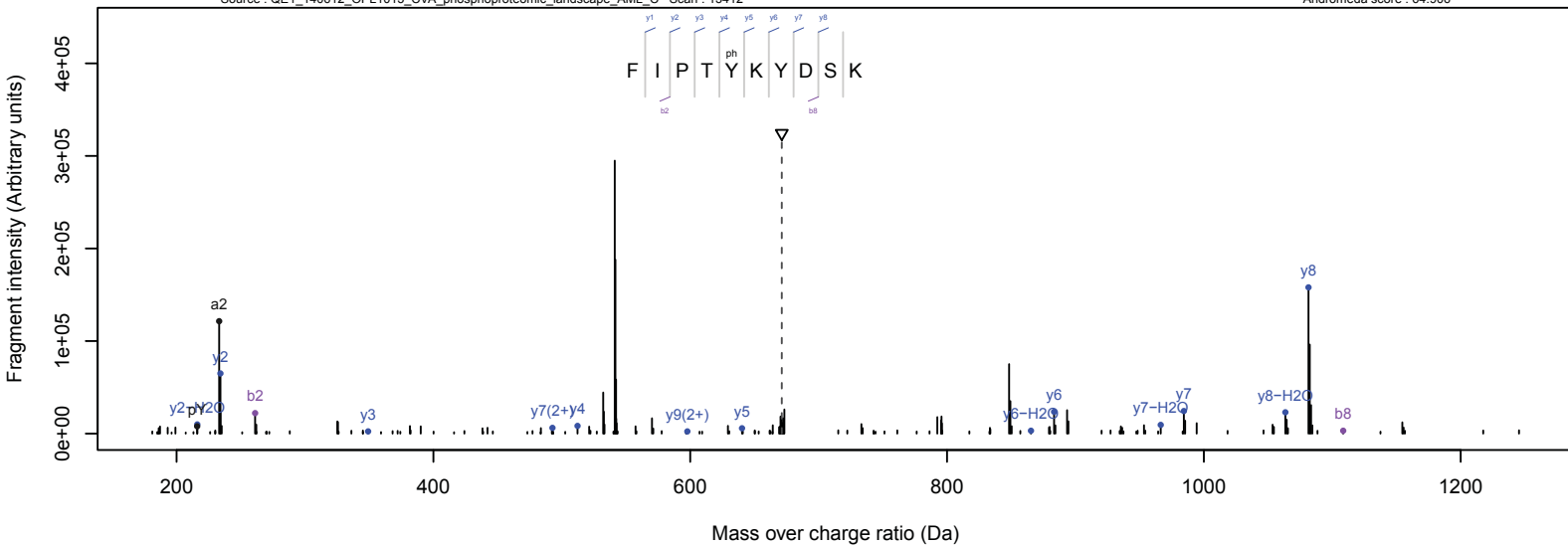




## FIPTY(ph)KYDSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13412

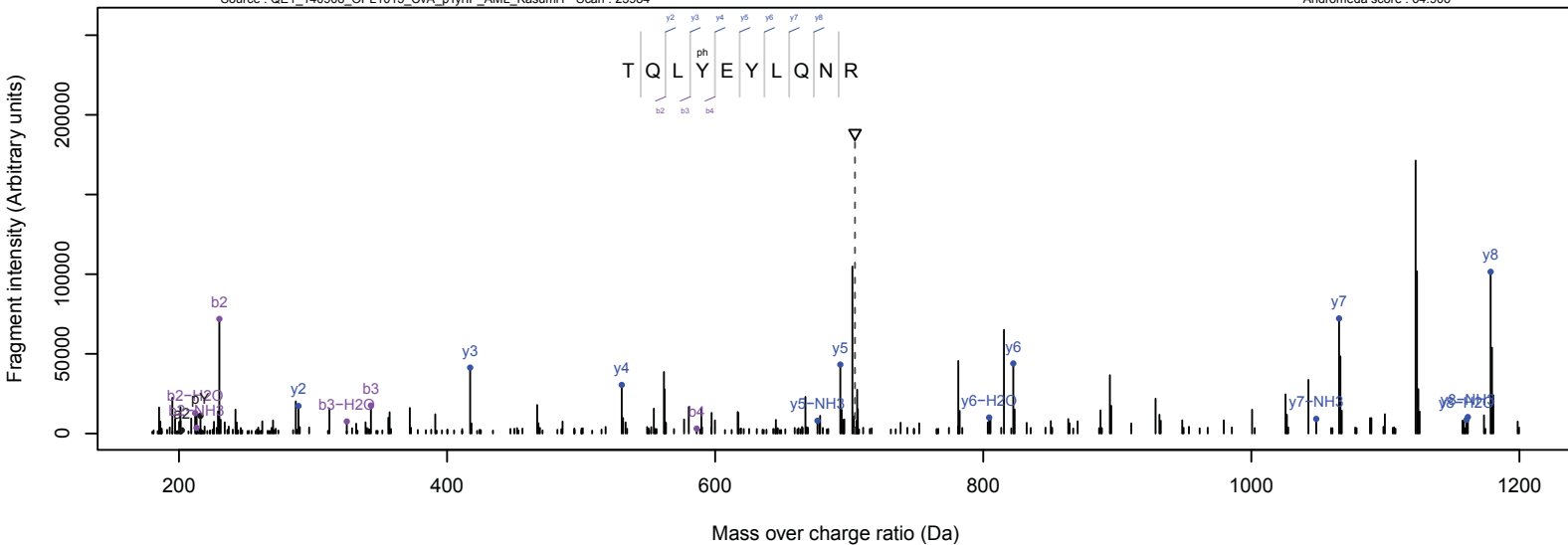
Andromeda score : 64.906



## TQLY(ph)EYLQNR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 25984

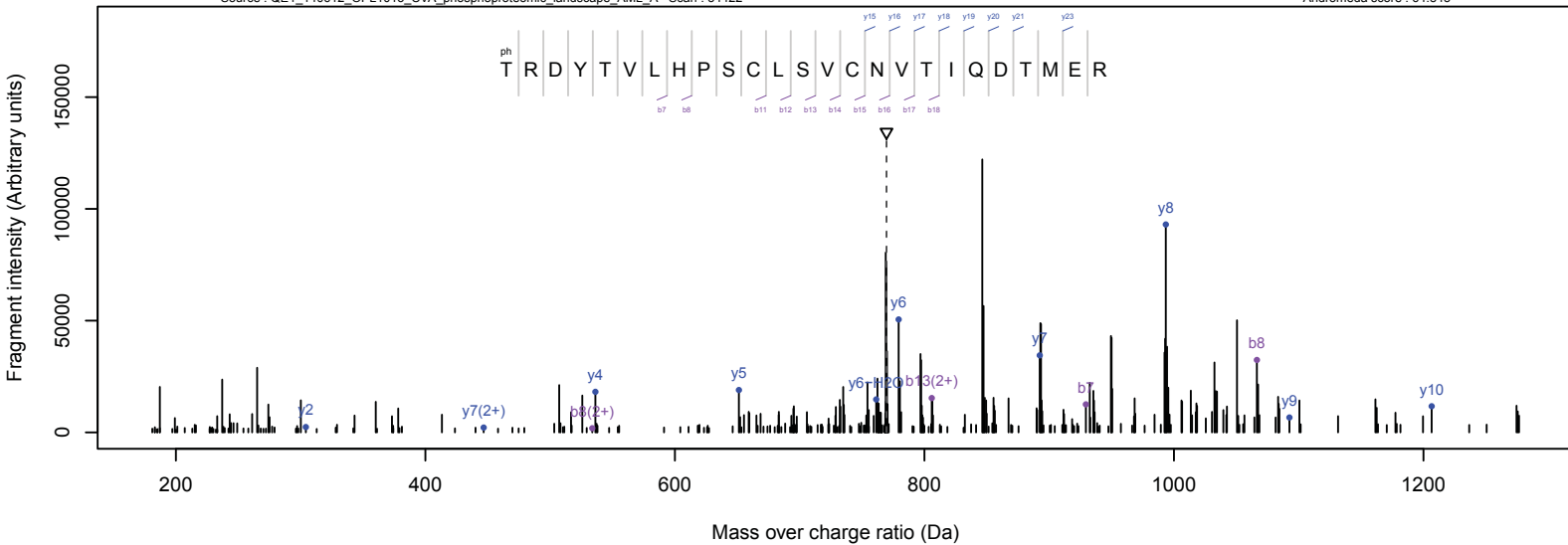
Andromeda score : 64.906



## T(ph)RDYTVLHPSCLSVCNVTIQDTMER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34422

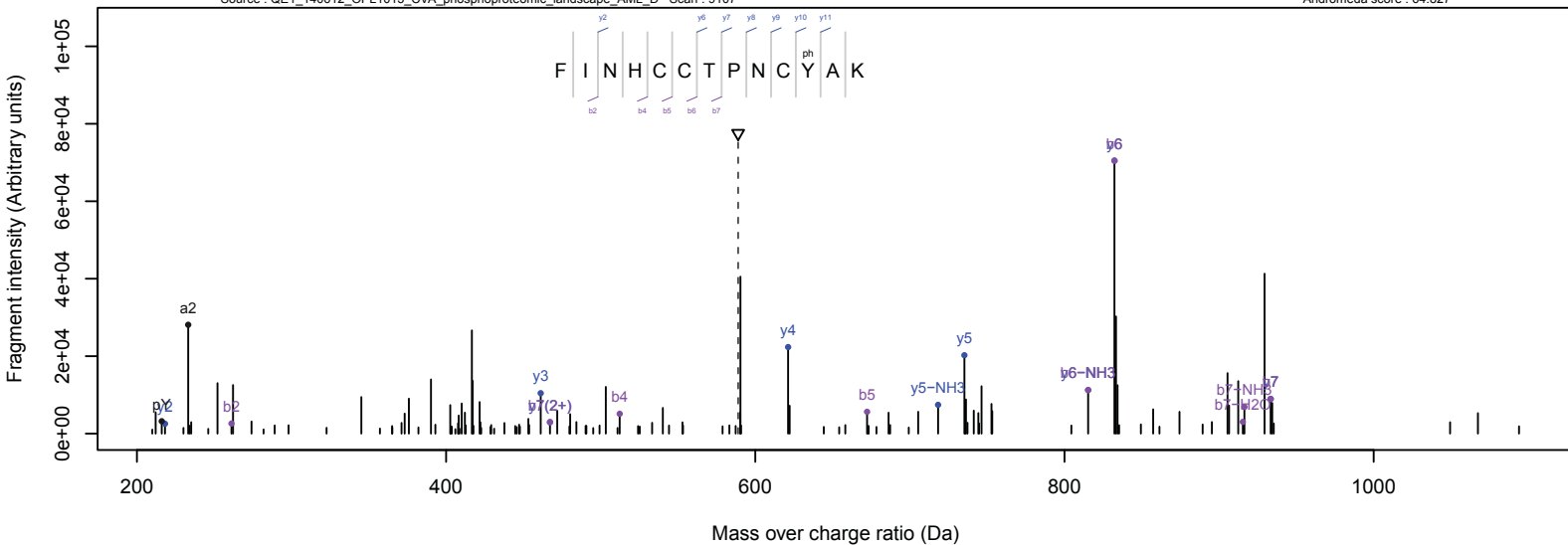
Andromeda score : 64.845



## FINHCCTPNCYAK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 9107

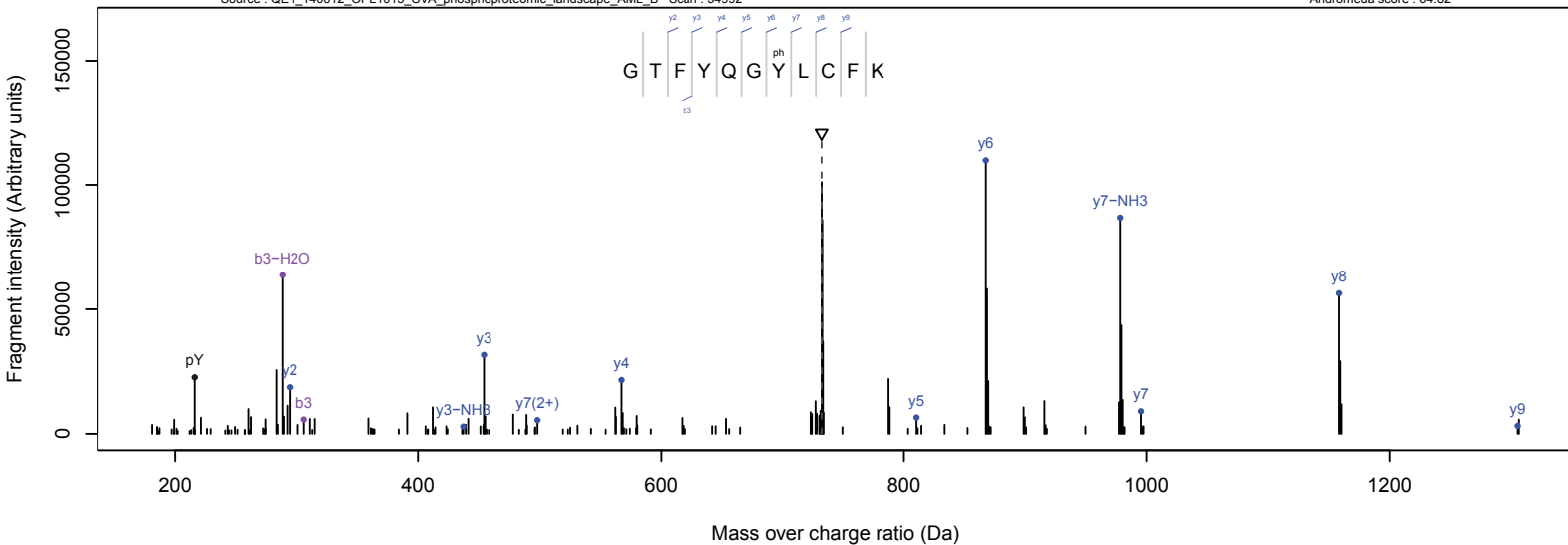
Andromeda score: 64.827



## GTFYQGYLFCFK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34992

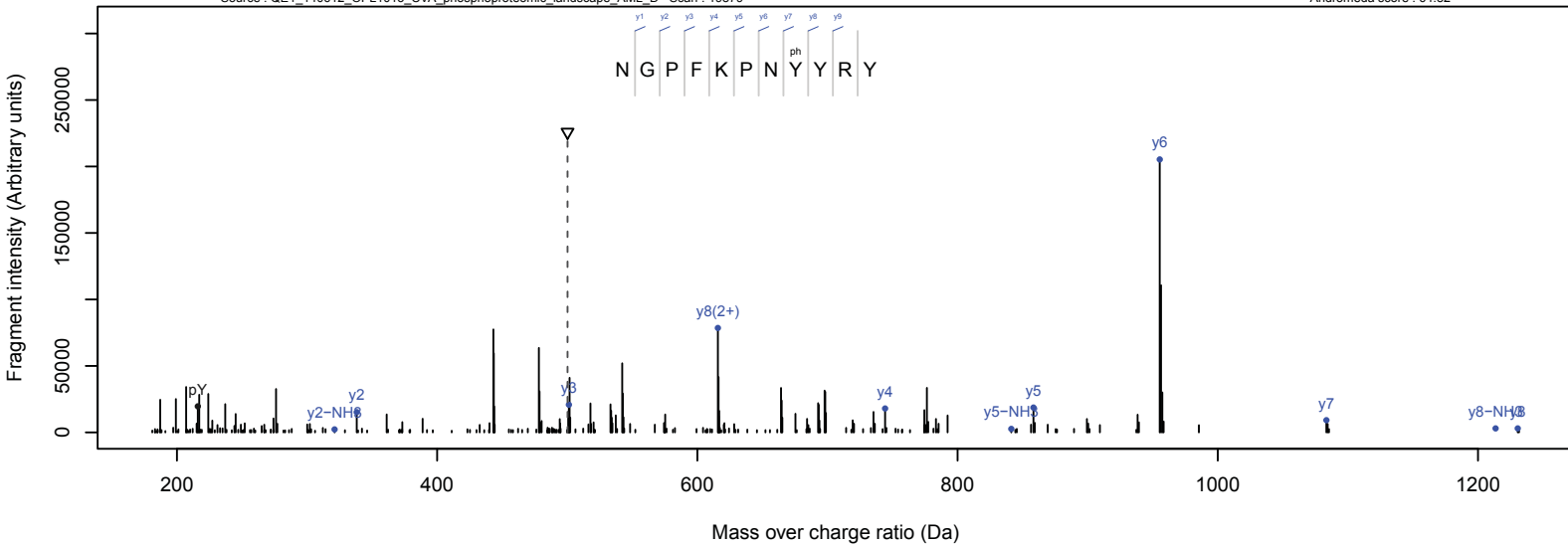
Andromeda score: 64.82



## NGPFKPNYYRY-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 19879

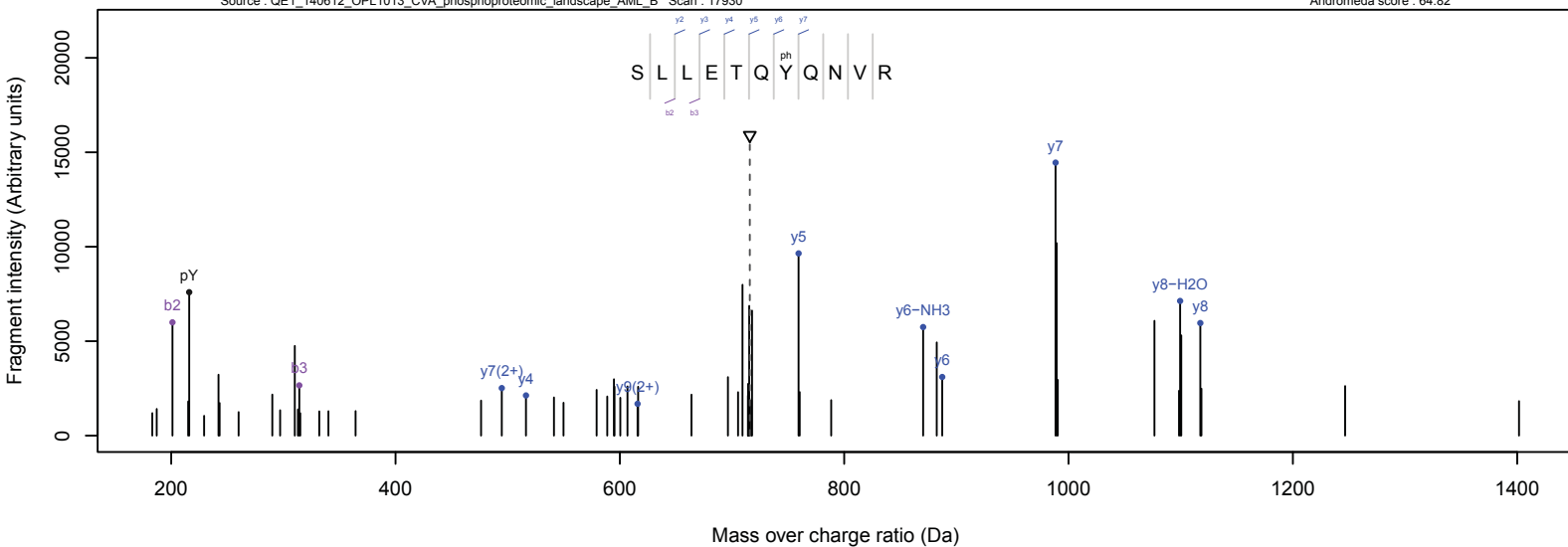
Andromeda score: 64.82



## SLLETQ(ph)QNVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 17930

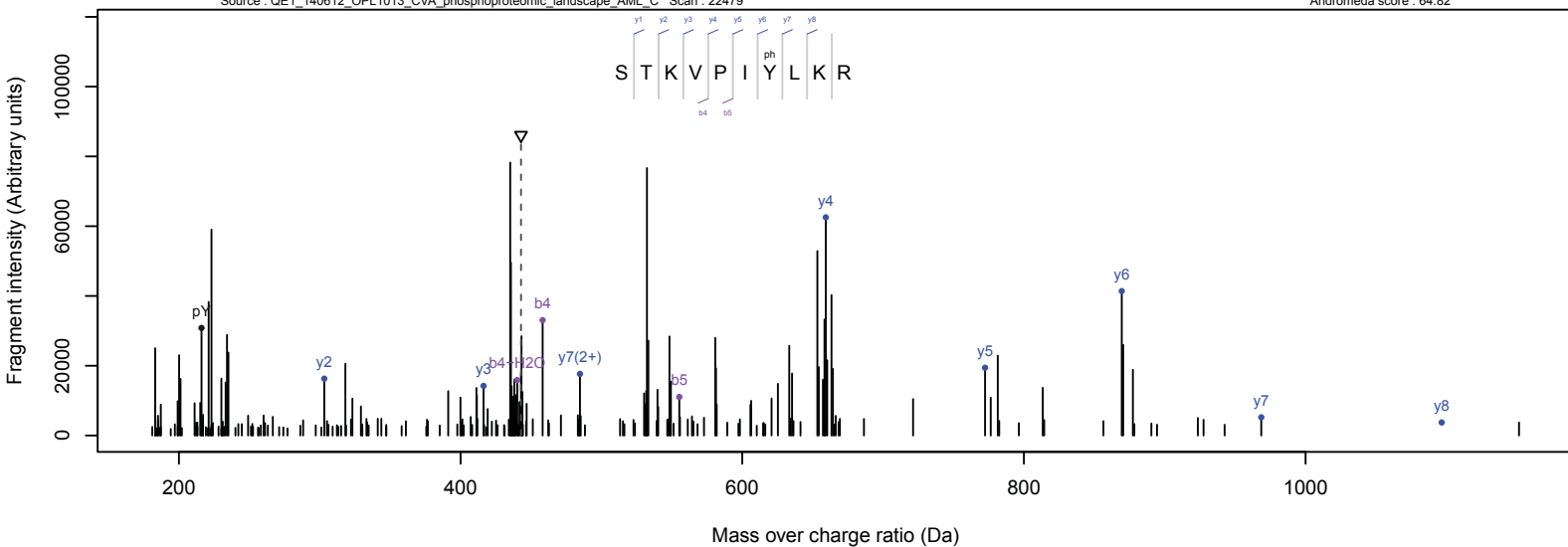
Andromeda score: 64.82



## (ac)STKVPYI(ph)LKR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 22479

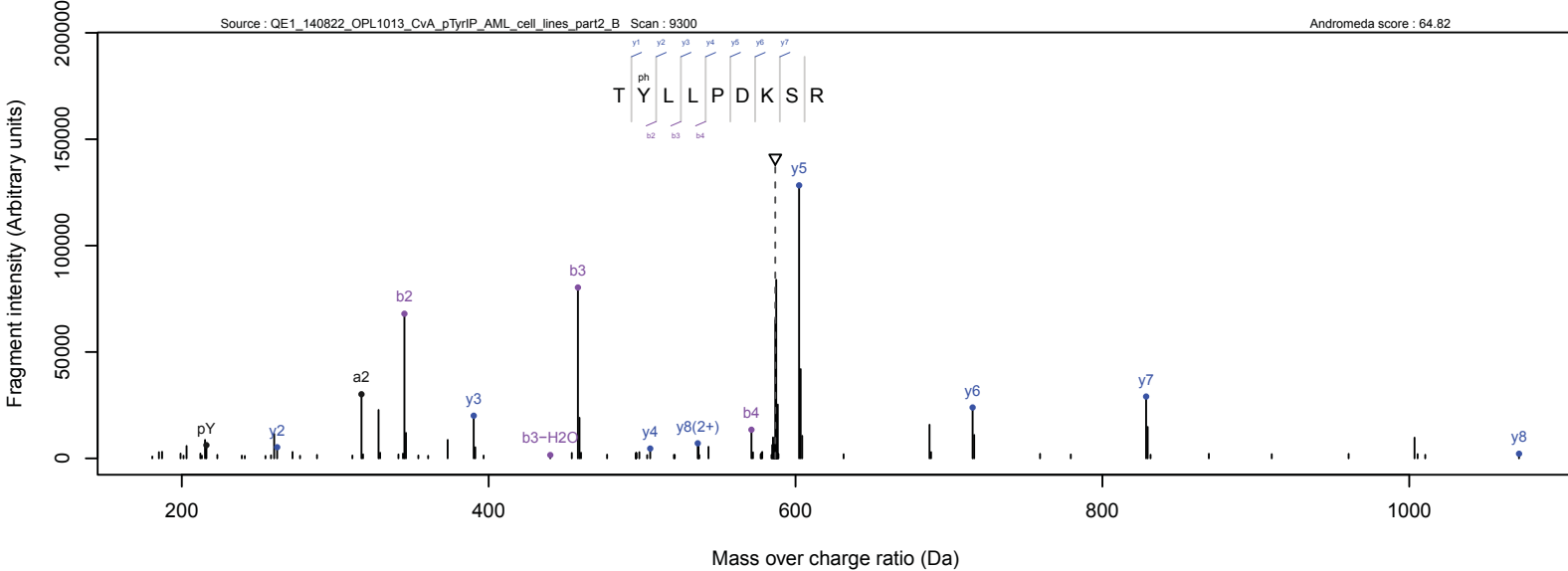
Andromeda score: 64.82



## TY(ph)LLPKSR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 9300

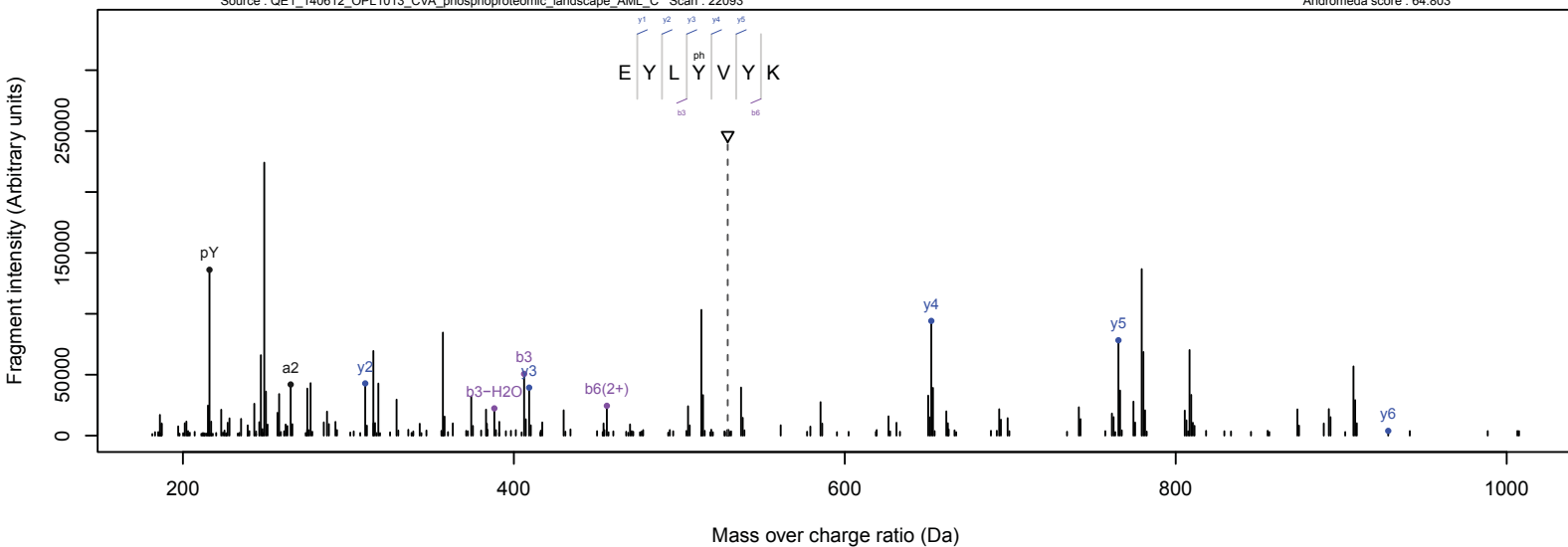
Andromeda score: 64.82



EYLY(ph)VYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22093

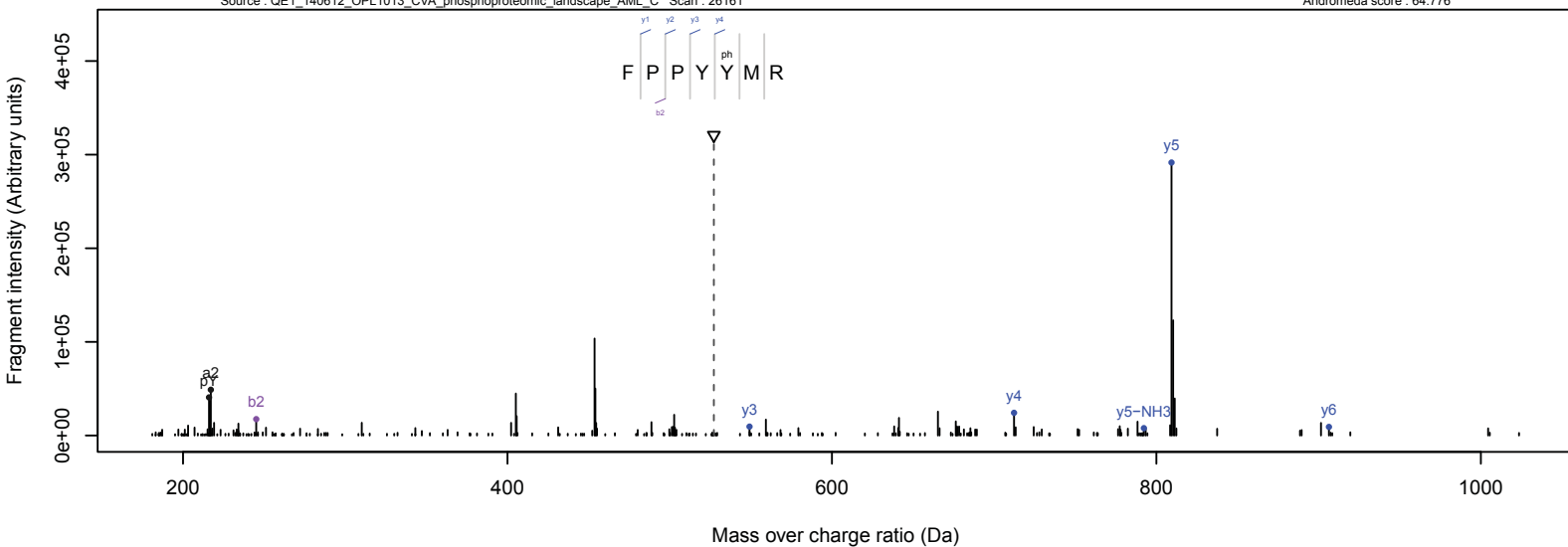
Andromeda score : 64.803



FPYYP(ph)MR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26161

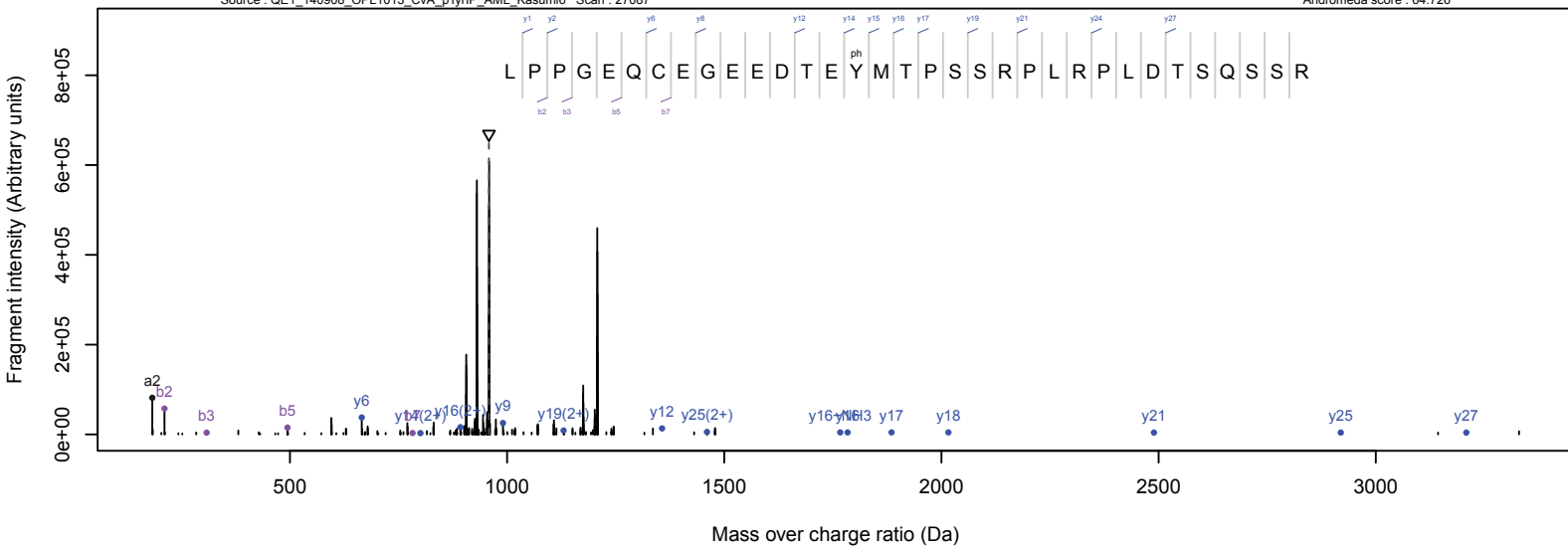
Andromeda score : 64.776



LPPGEQCEGEDTEY(ph)MTPSSRPLRLDTSQSSR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 27087

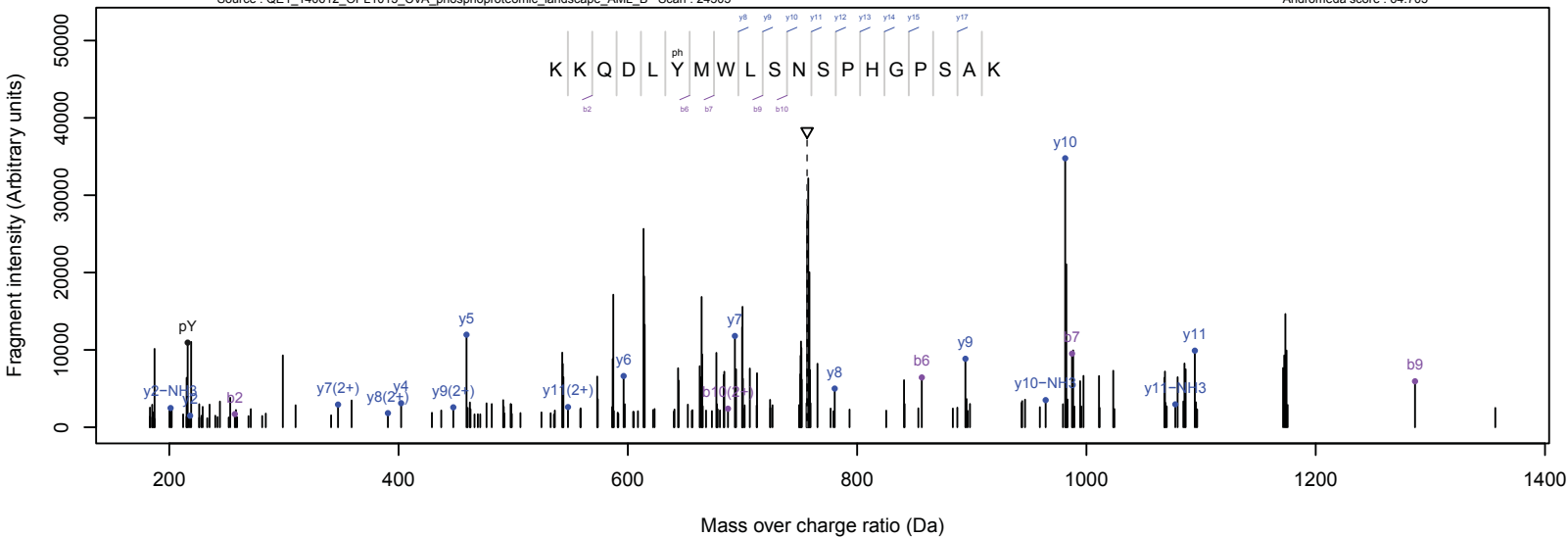
Andromeda score : 64.726



## KKQDLY(ph)MWLSNSPHGPSAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24505

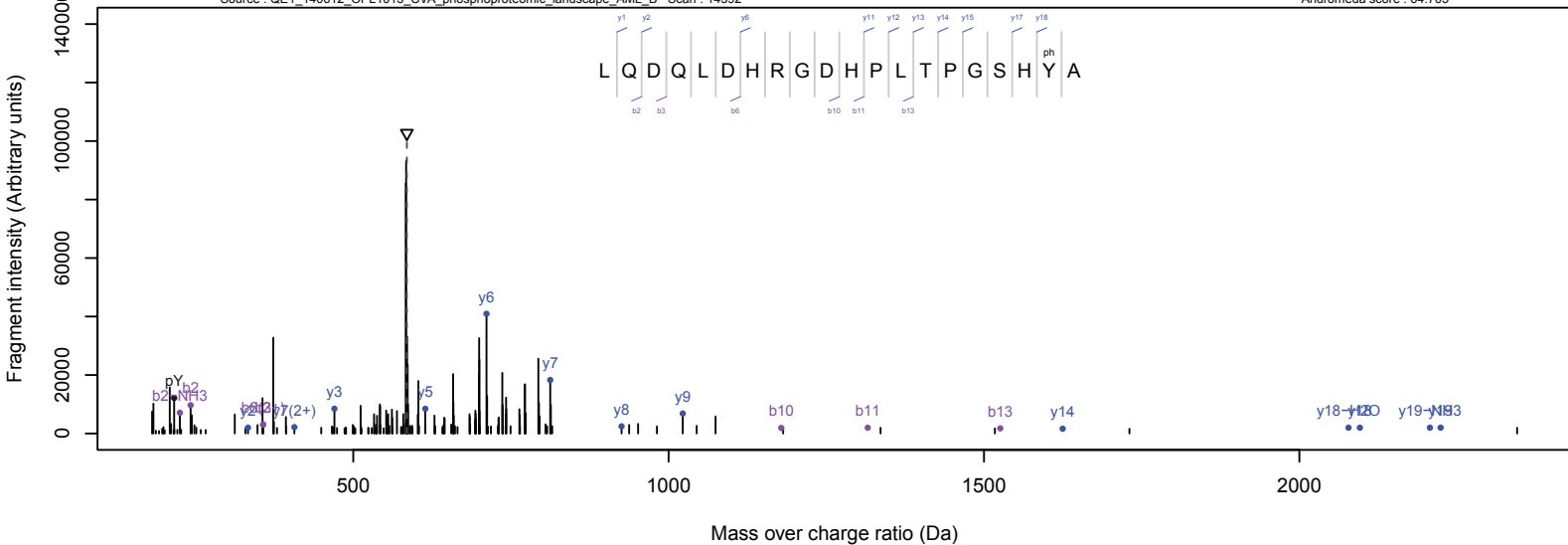
Andromeda score : 64.705



## LQDQLDHRGDHPLTPGSHY(ph)A-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14392

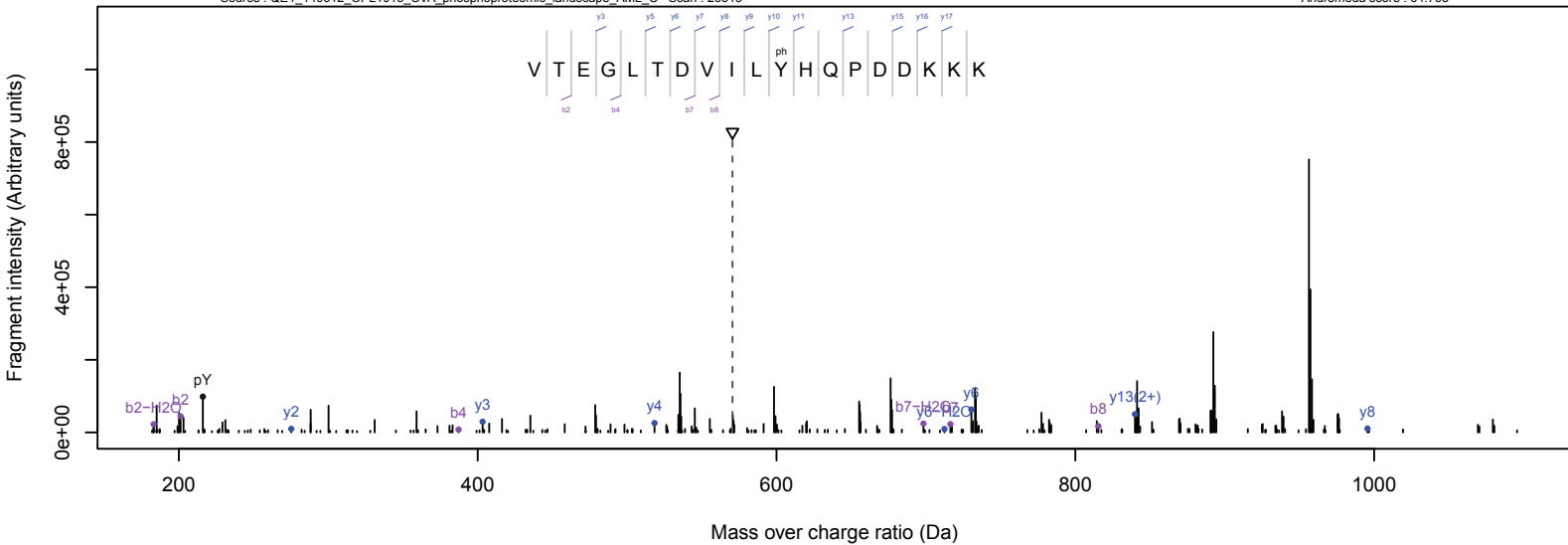
Andromeda score : 64.705



## VTEGLTDVILYHQPD DKKK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26313

Andromeda score : 64.705

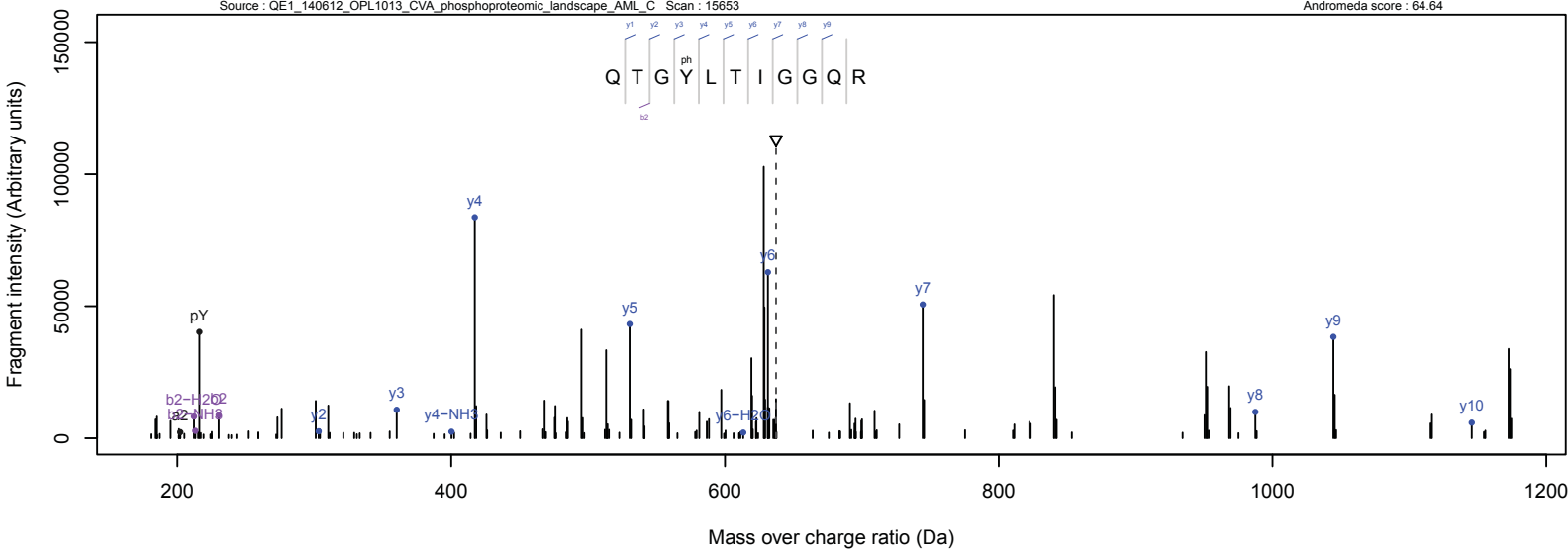




**QTGY(ph)LTIGGQR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15653

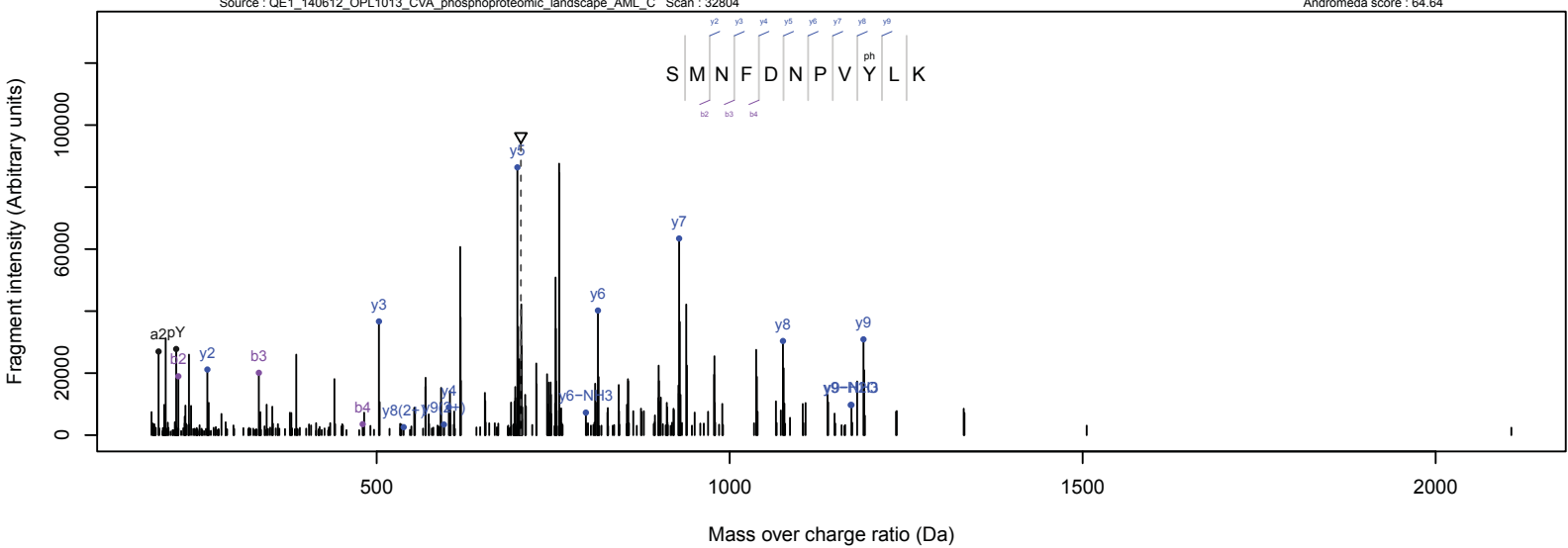
Andromeda score : 64.64



**SMNFDNPVY(ph)LK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32804

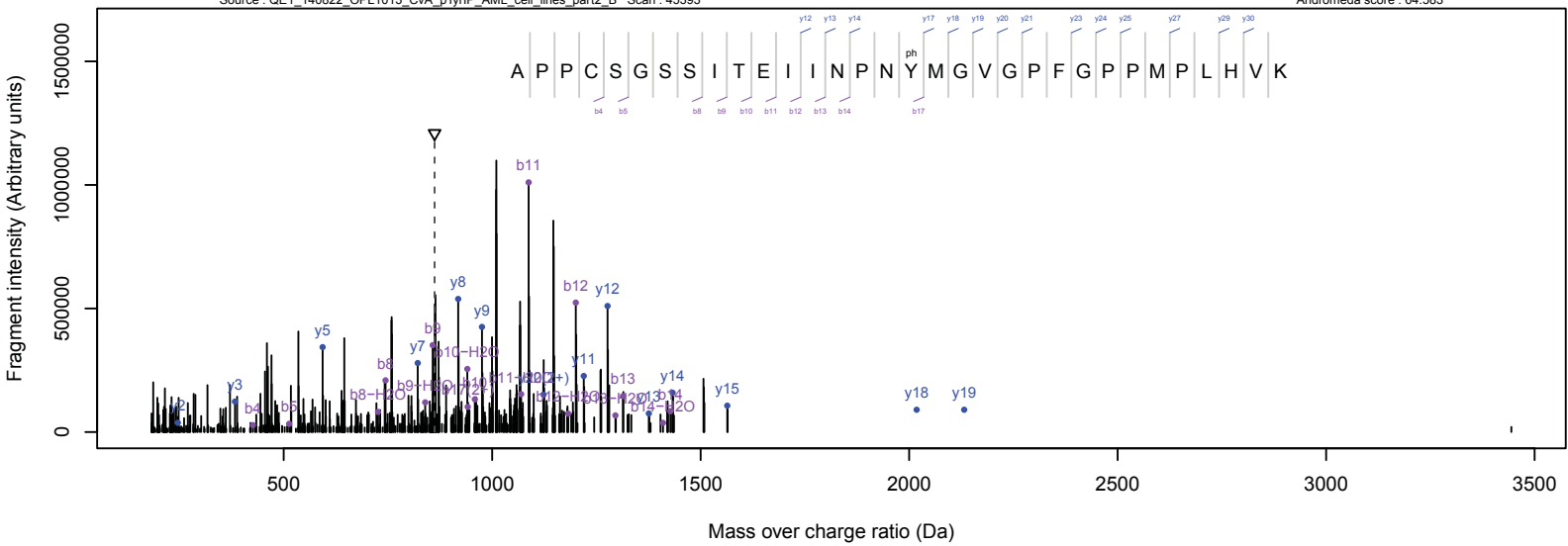
Andromeda score : 64.64



**APPCSGSSITEIINPNYMGVGFPPMPLHVK-4+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 45393

Andromeda score : 64.583

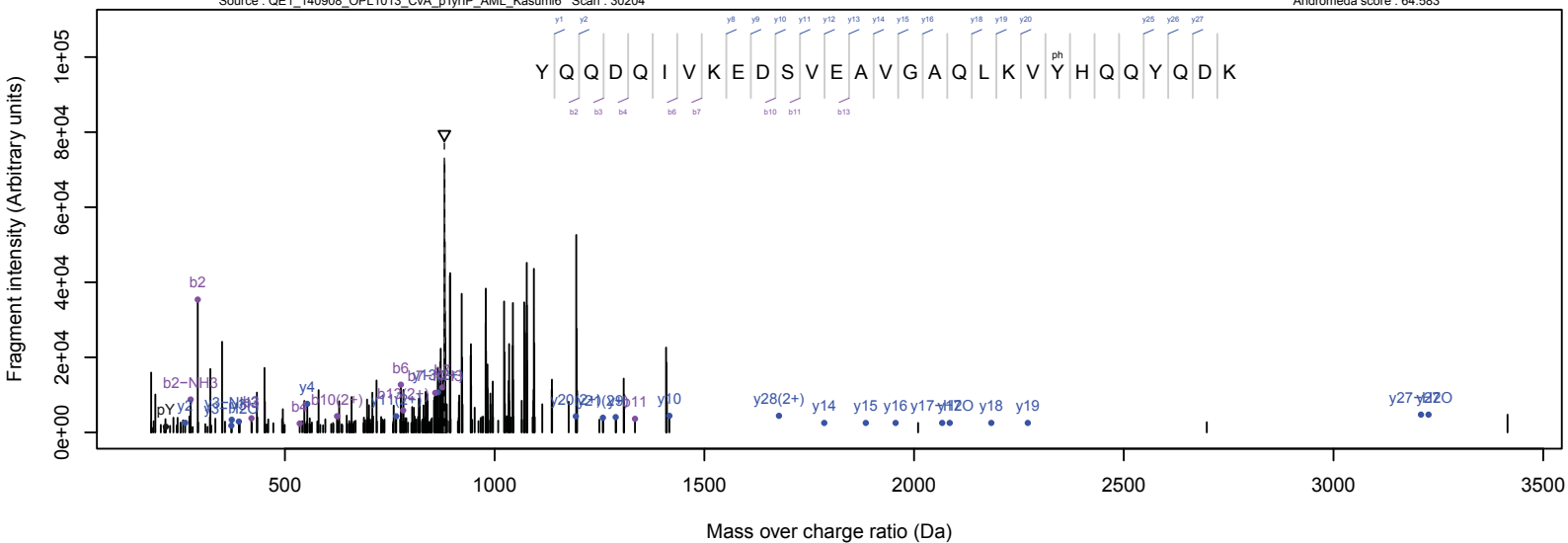




YQQDQIVKEDSVEAVGAQLK<sup>ph</sup>HQQYQDK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 30204

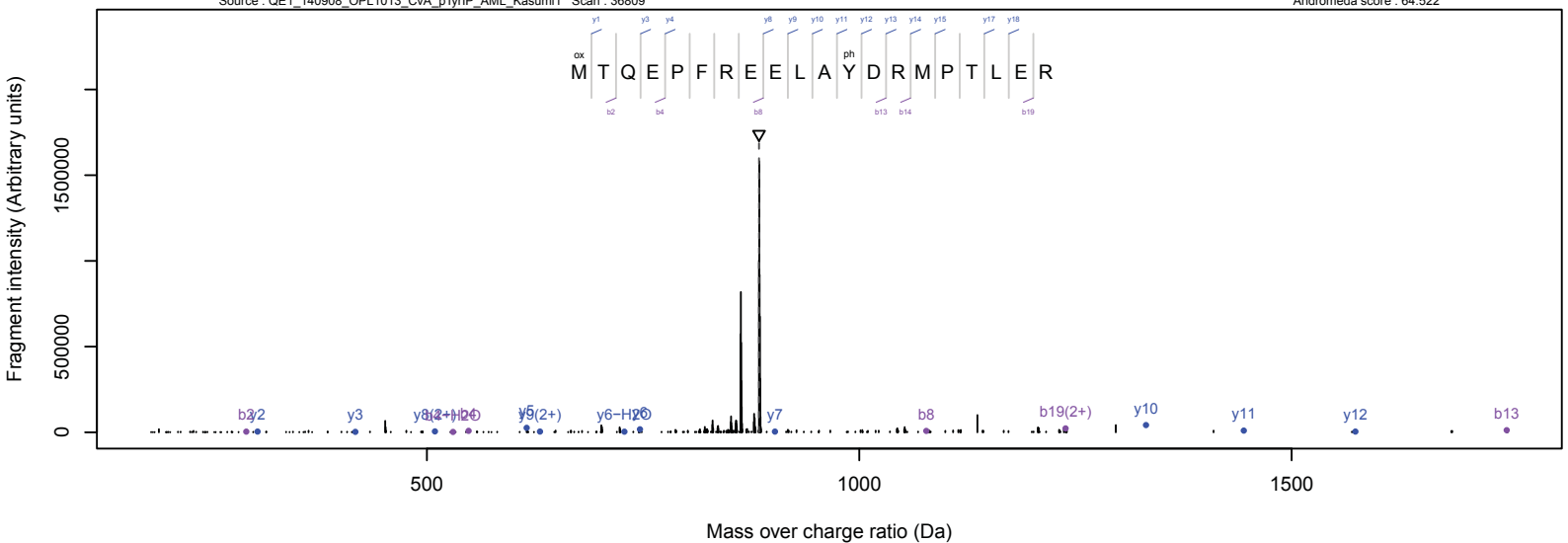
Andromeda score : 64.583



(ac)M(ox)TQEPFREELAY<sup>ph</sup>DRMPTLER-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 36809

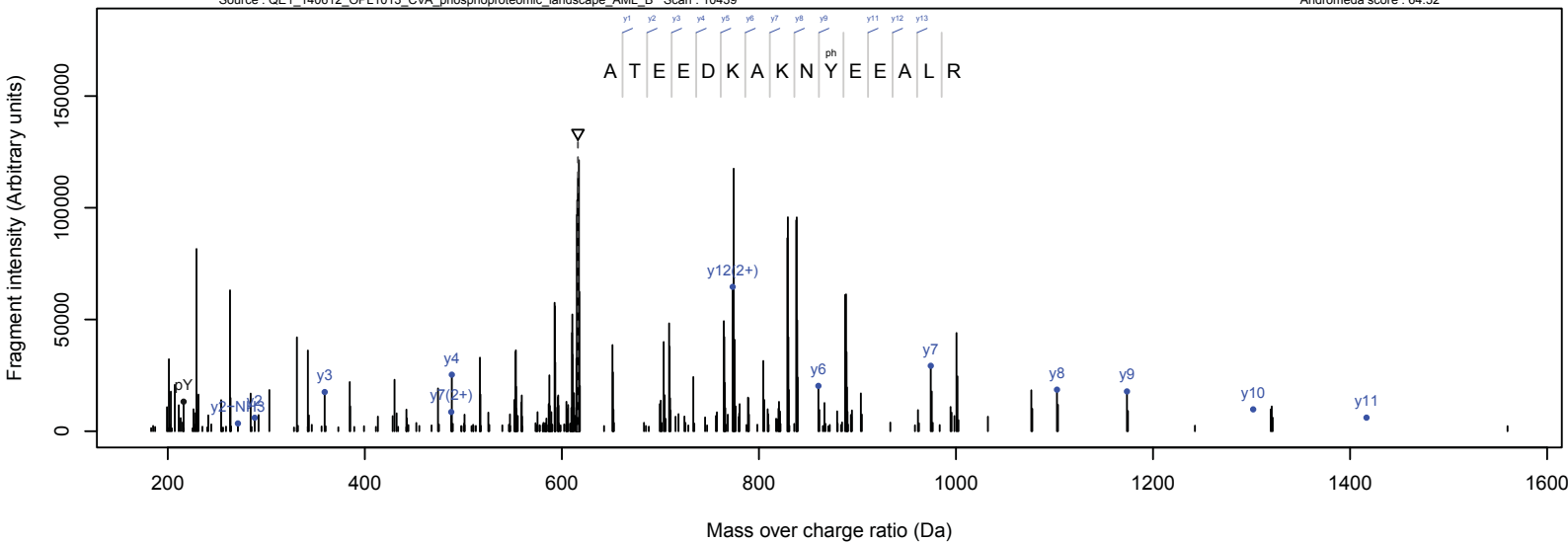
Andromeda score : 64.522



ATEEDKAKNY<sup>ph</sup>EELR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_B Scan : 10439

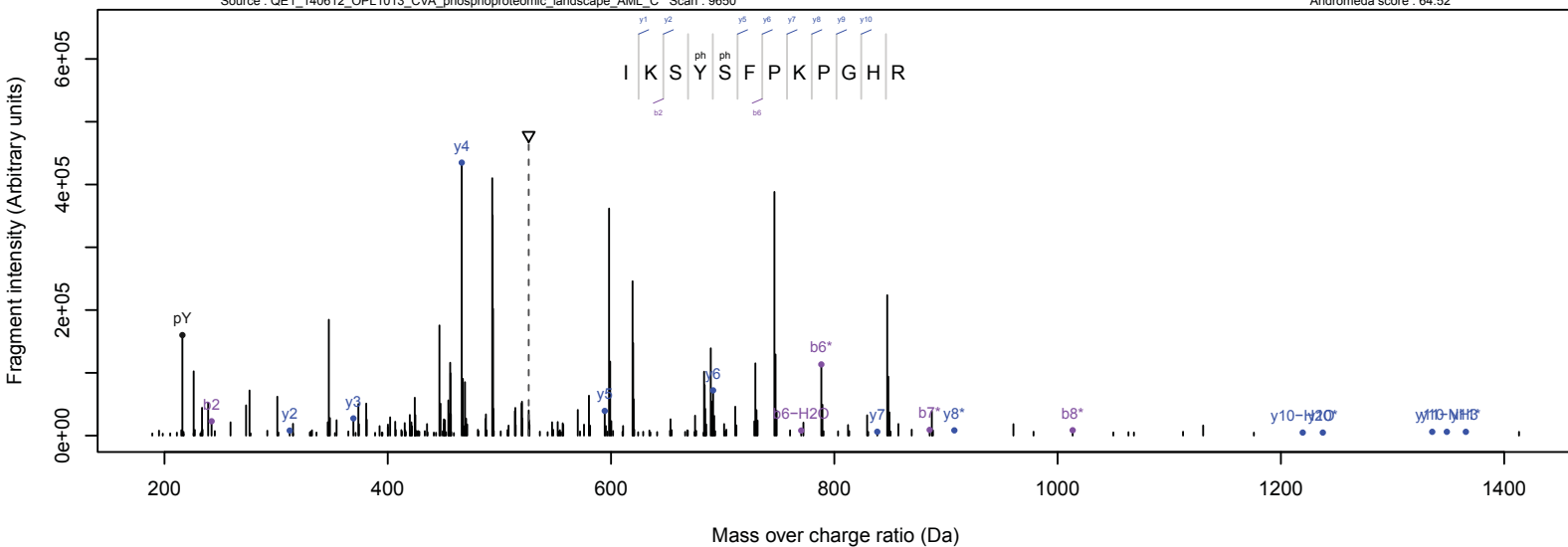
Andromeda score : 64.52



## IKSY(ph)S(ph)FPKPGHR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 9650

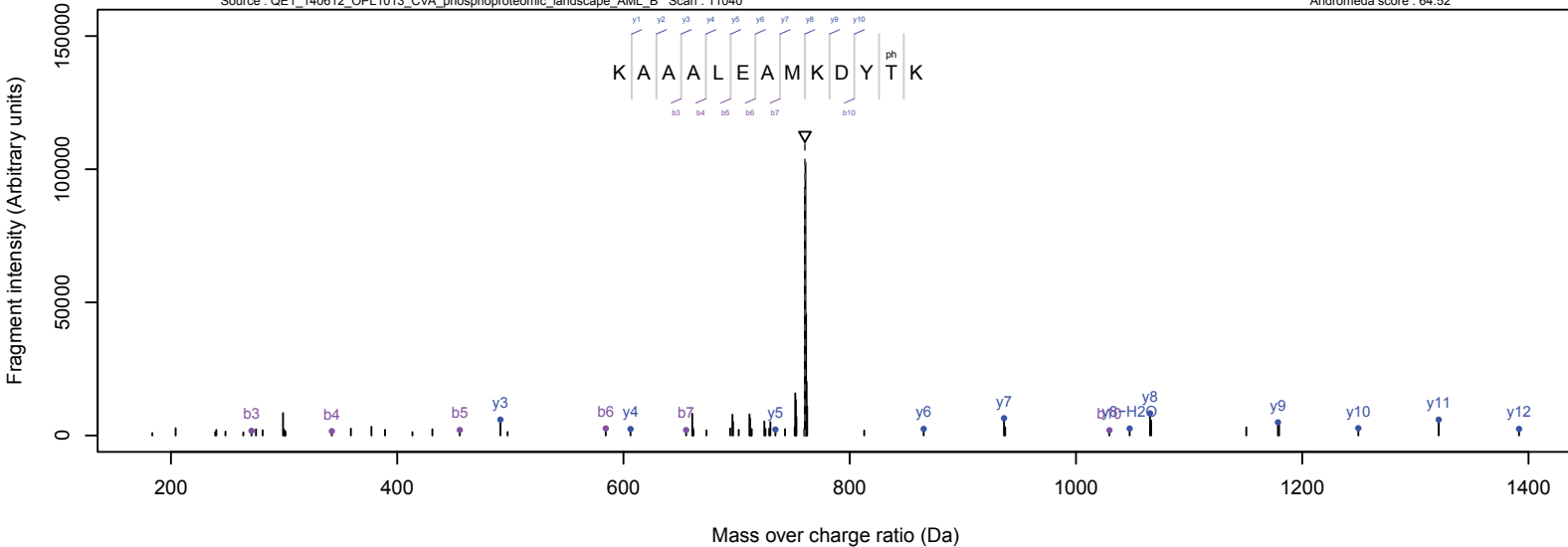
Andromeda score: 64.52



## KAAALEAMKDYT(ph)K-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 11040

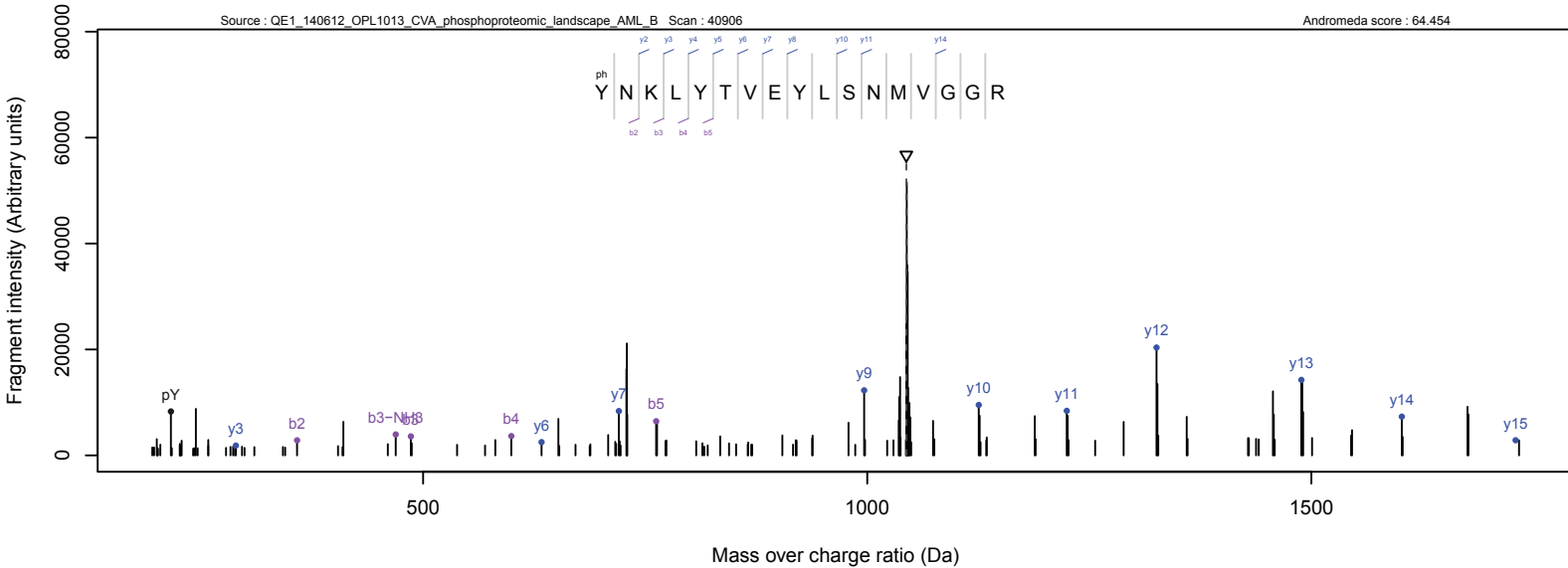
Andromeda score: 64.52



## Y(ph)NKLYTVEYLSNMVGGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 40906

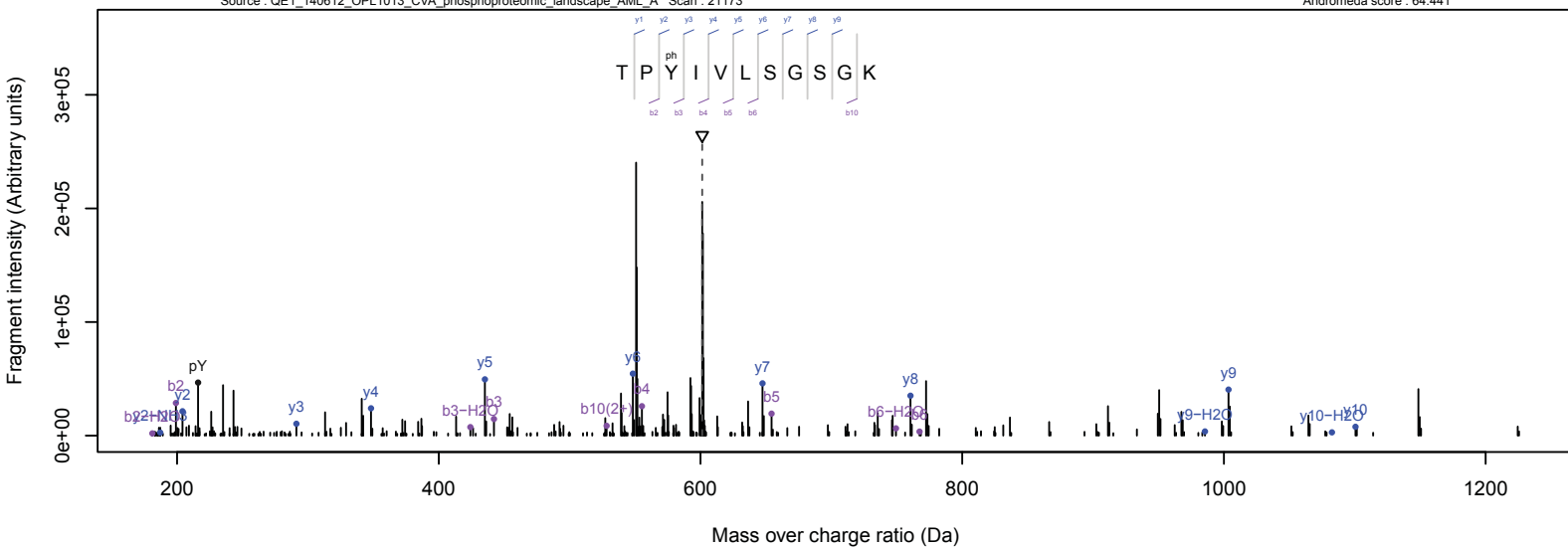
Andromeda score: 64.454



## TPY(ph)IVLSGSGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 21173

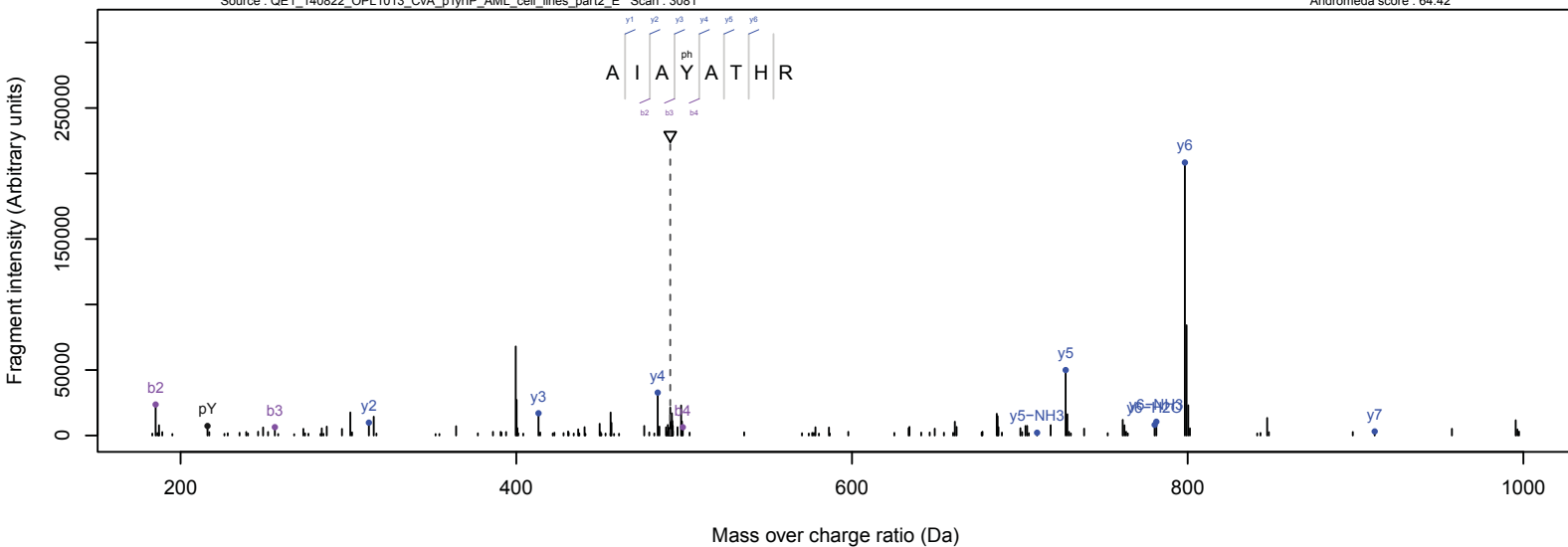
Andromeda score : 64.441



## AIAY(ph)ATHR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 3081

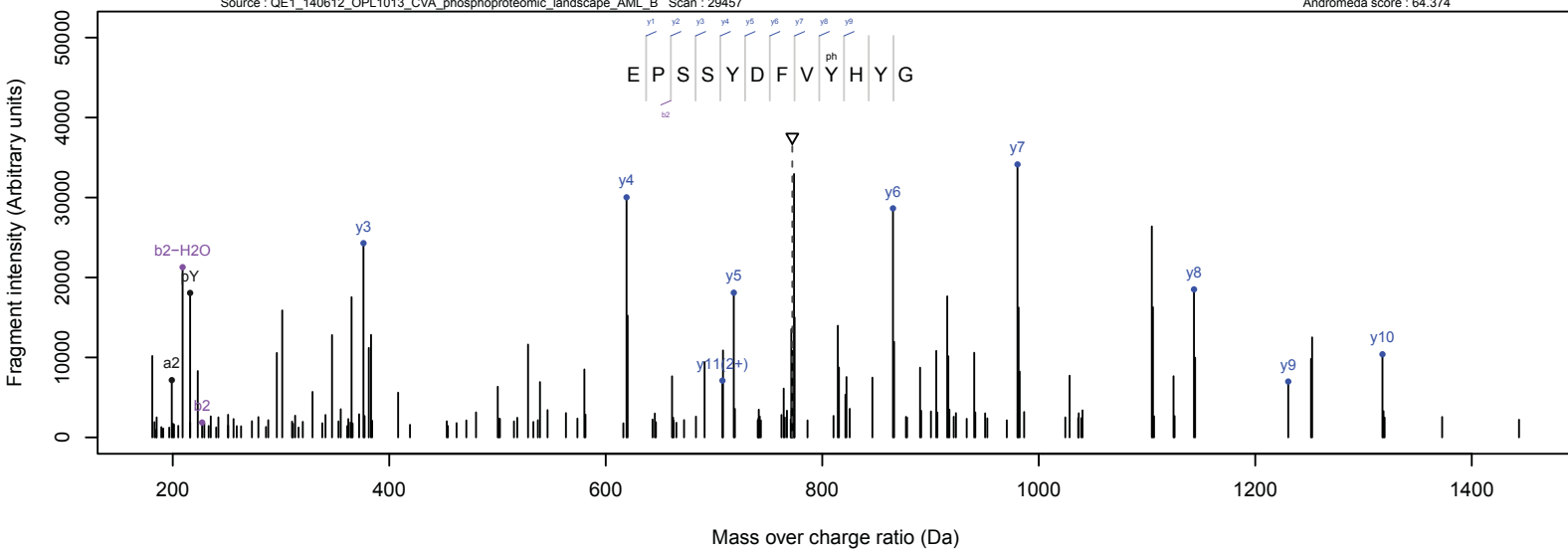
Andromeda score : 64.42



## EPSSYDFVY(ph)HYG-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29457

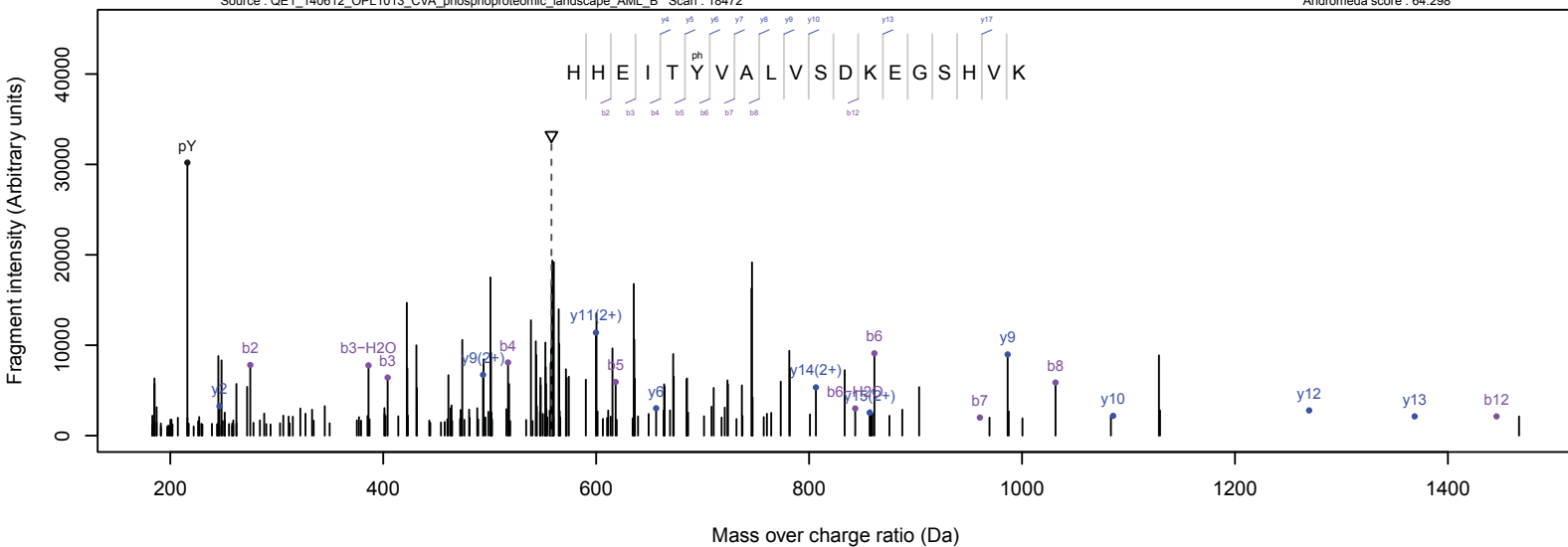
Andromeda score : 64.374



## HHEITY(ph)VALVSDKEGSHVK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18472

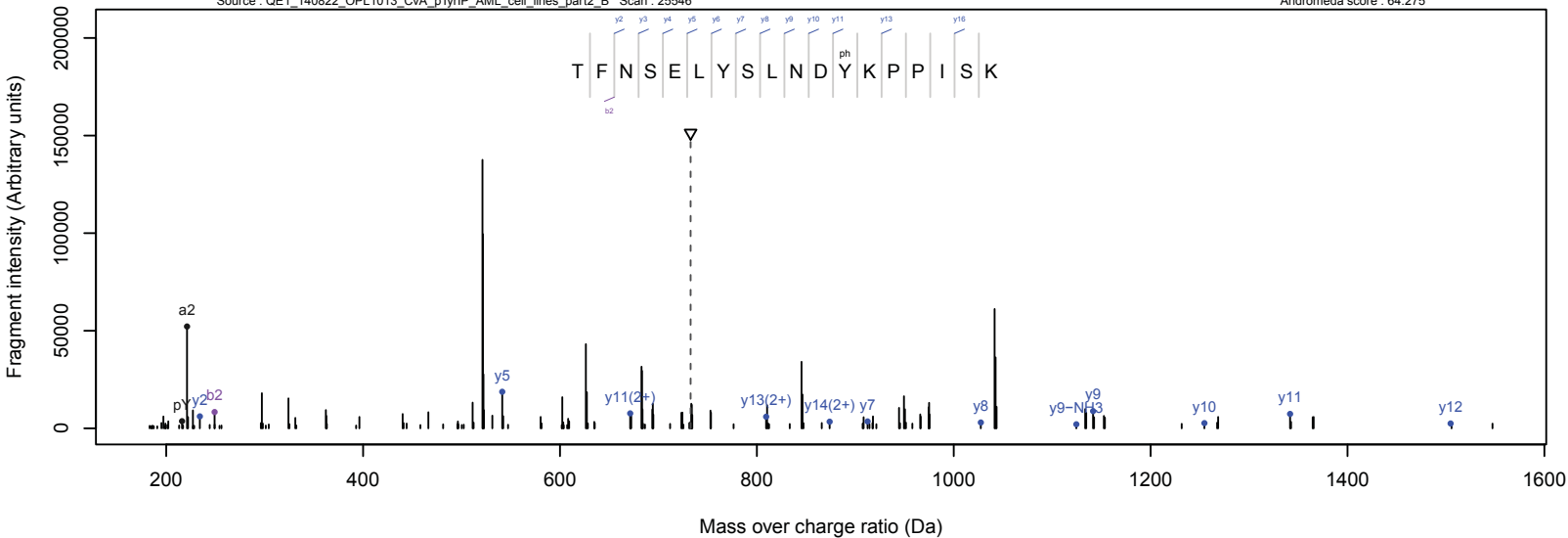
Andromeda score : 64.298



## TFNSELYSLNDY(ph)KPPISK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 25546

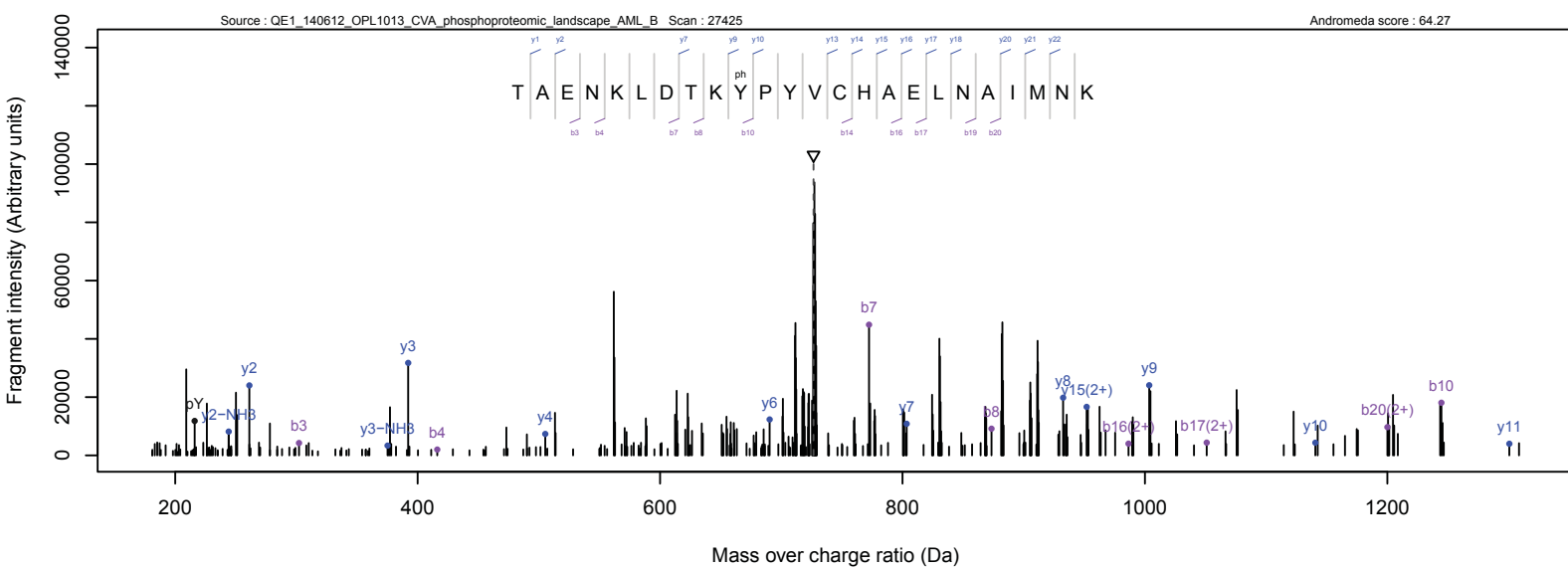
Andromeda score : 64.275



## TAENKLDTKY(ph)PYVCHAE LN A I M N K-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27425

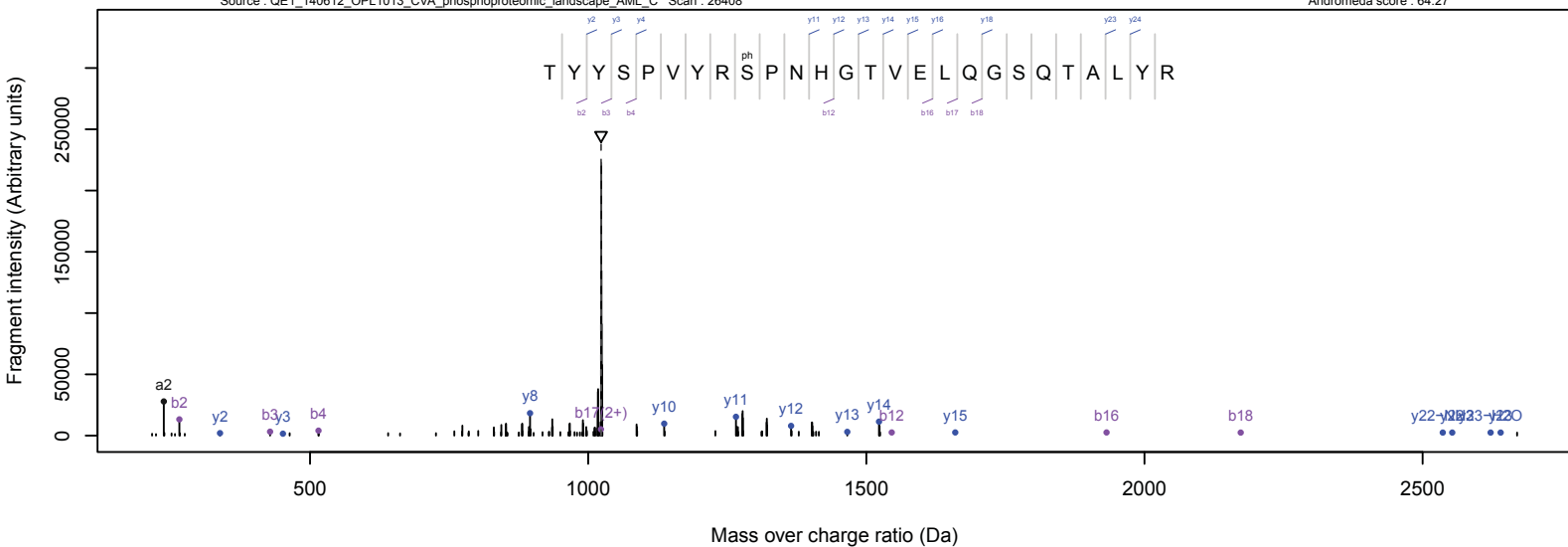
Andromeda score : 64.27



TYYSPVYRS(ph)PNHGTVELQGSQTALYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26408

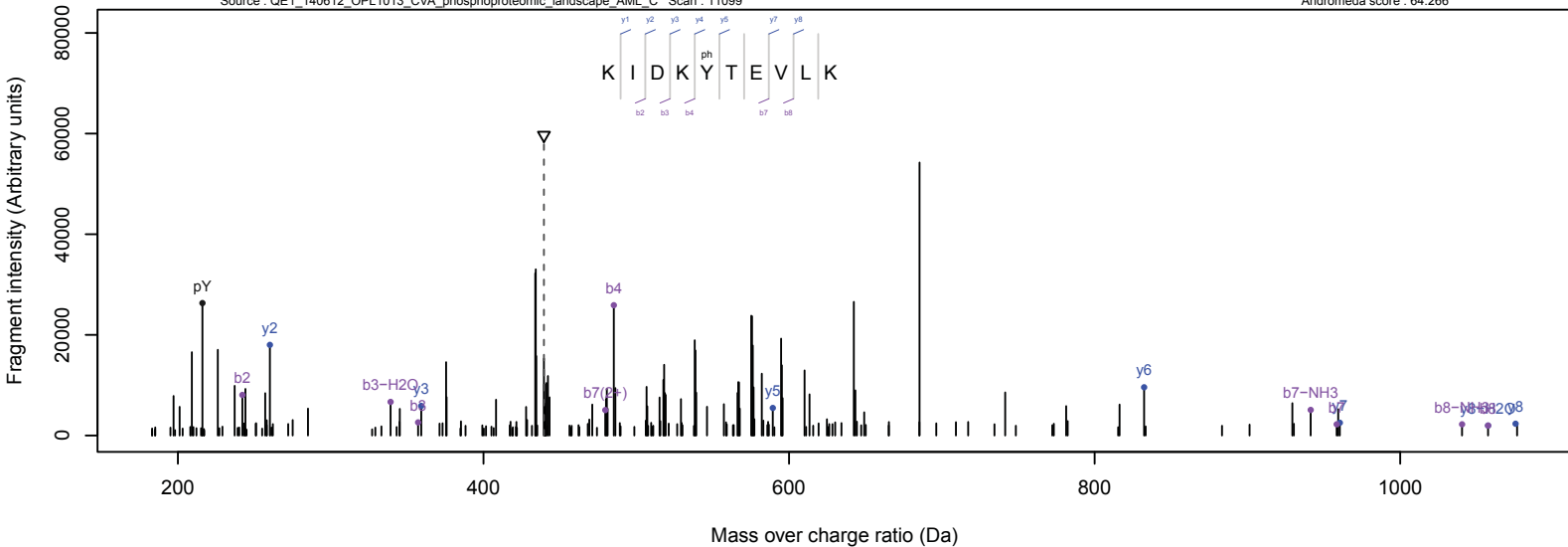
Andromeda score : 64.27



KIDKY(ph)TEVLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11099

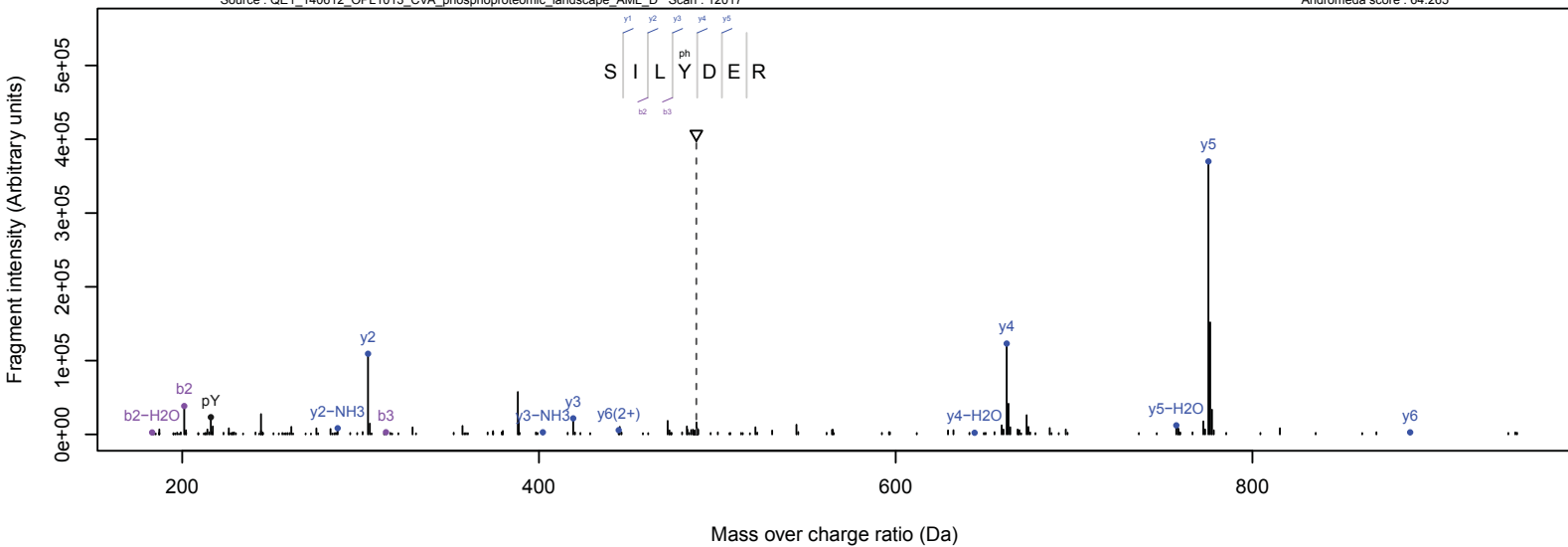
Andromeda score : 64.266

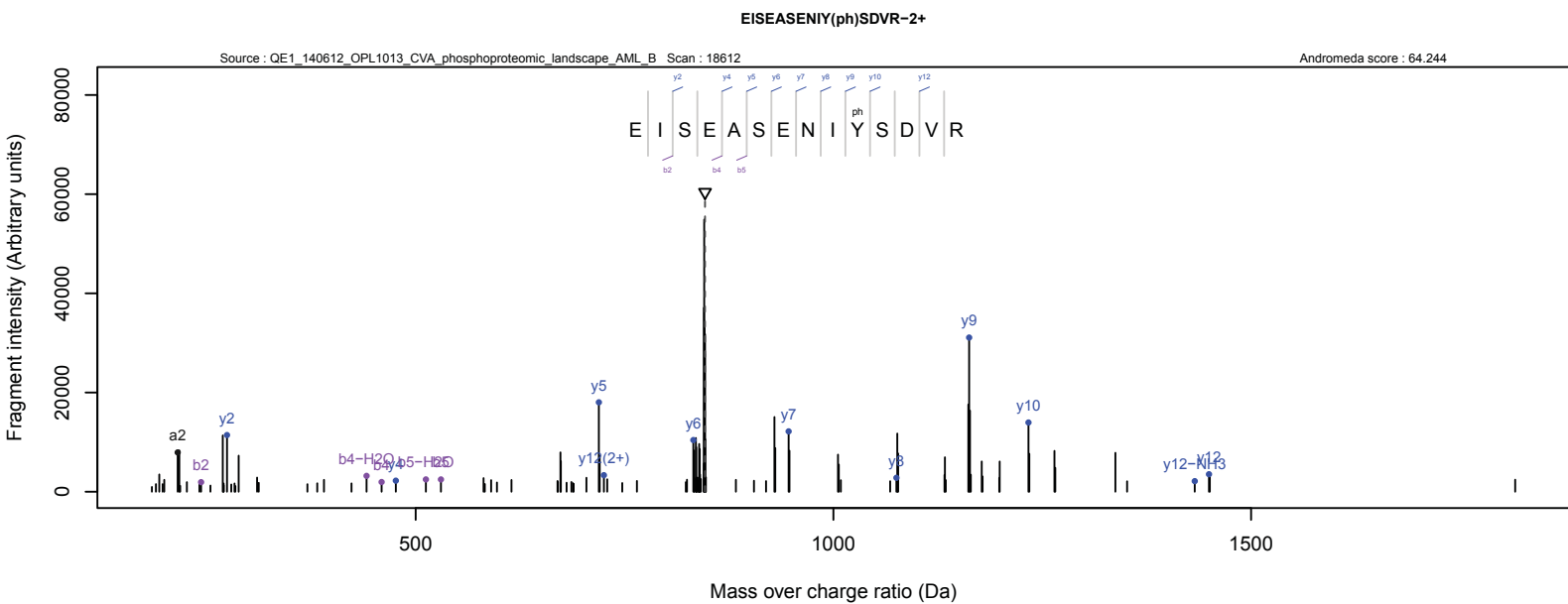
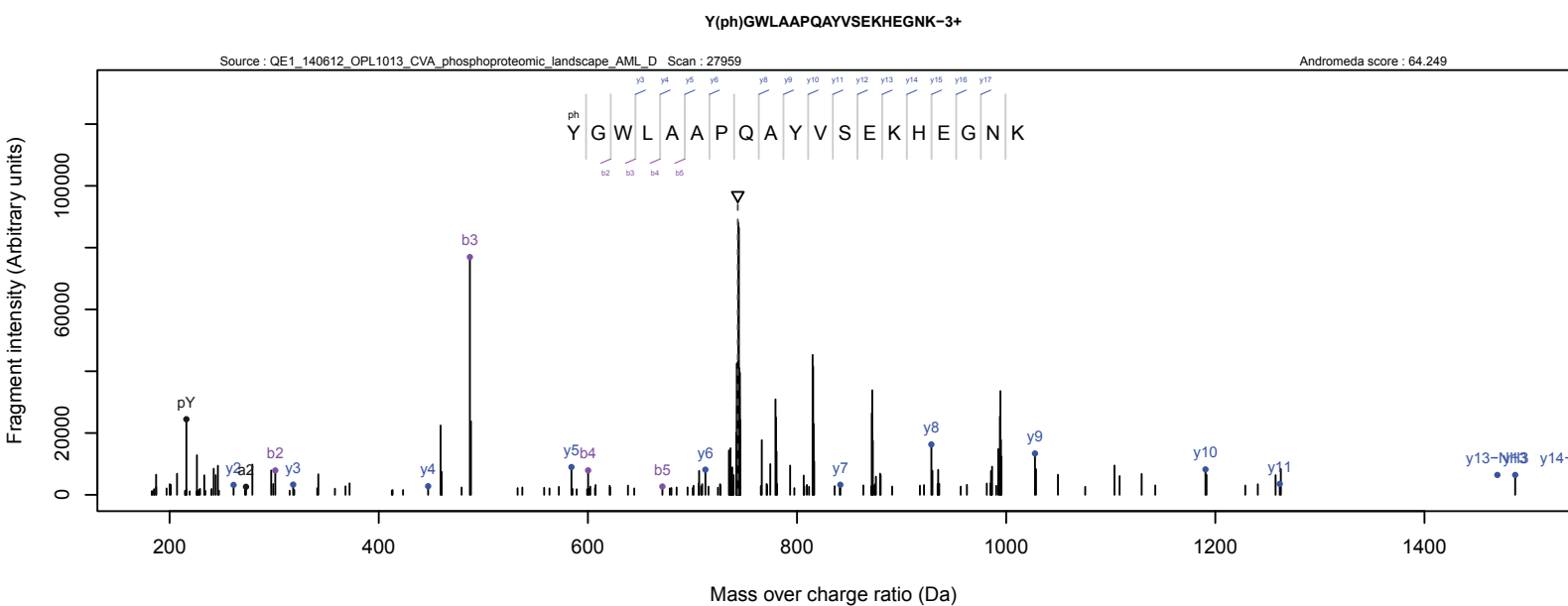
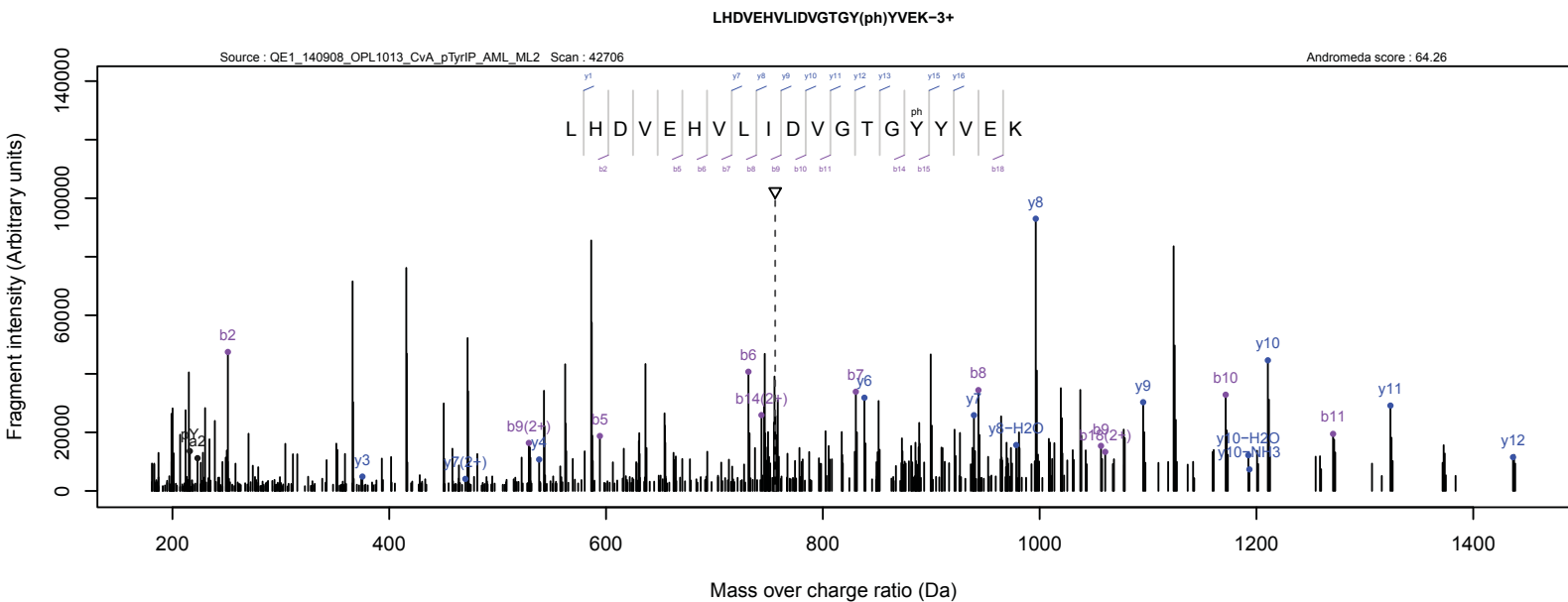


SILY(ph)DER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12017

Andromeda score : 64.265

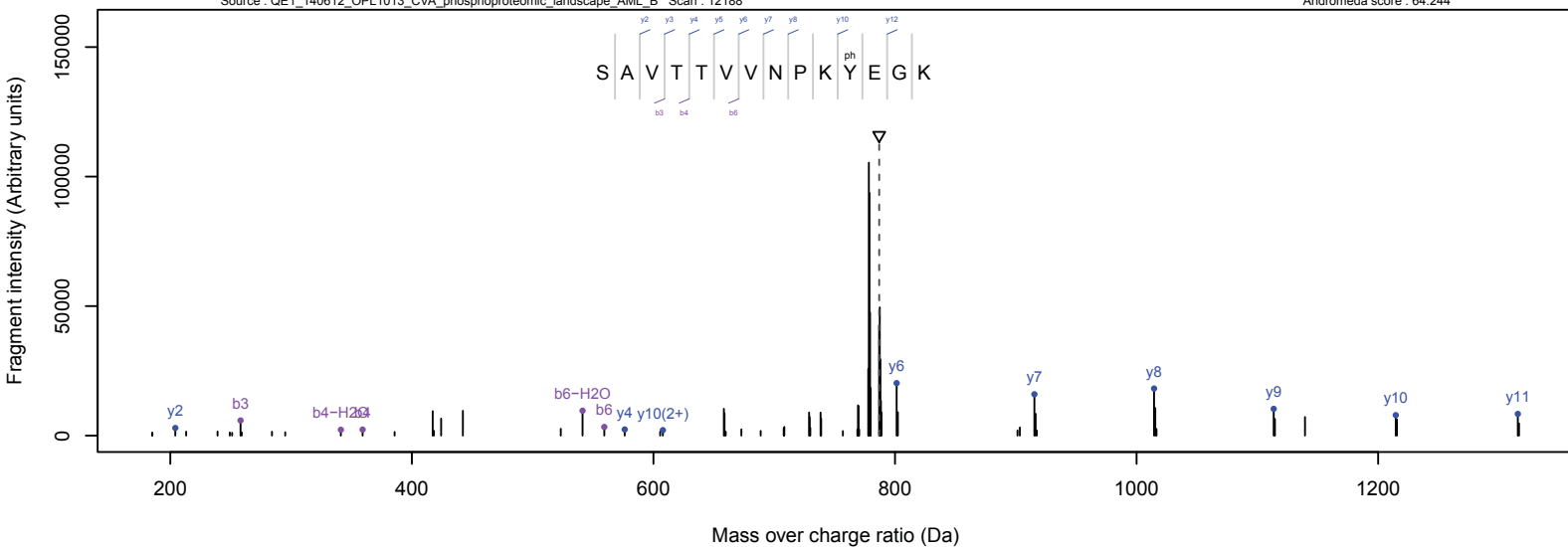




SAVTTVVNPK(ph)EGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12188

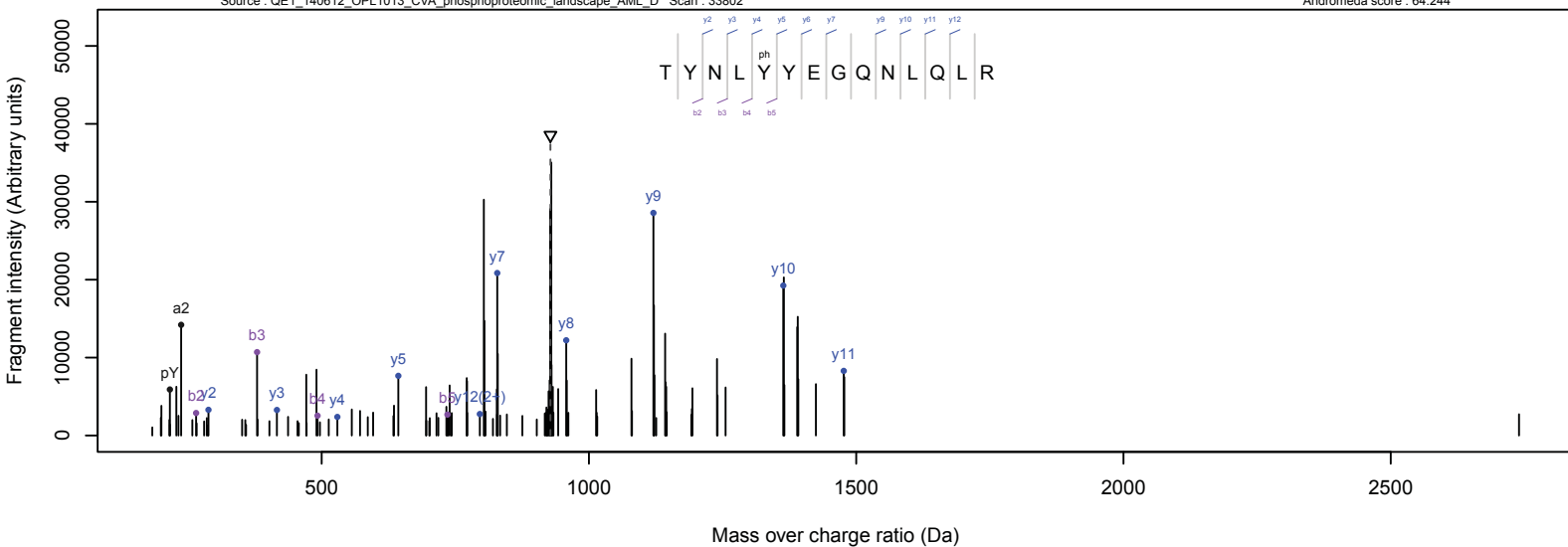
Andromeda score : 64.244



TYNLY(ph)YEGNQLQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33802

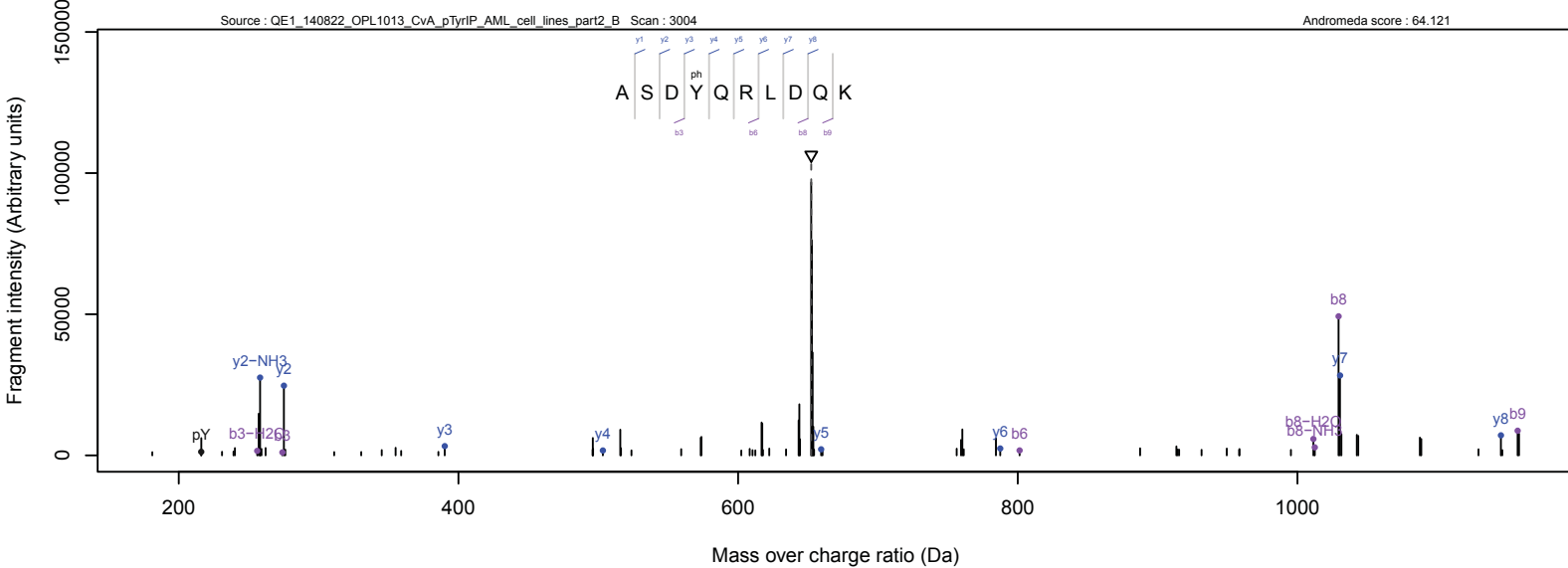
Andromeda score : 64.244



ASDY(ph)QLDQK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3004

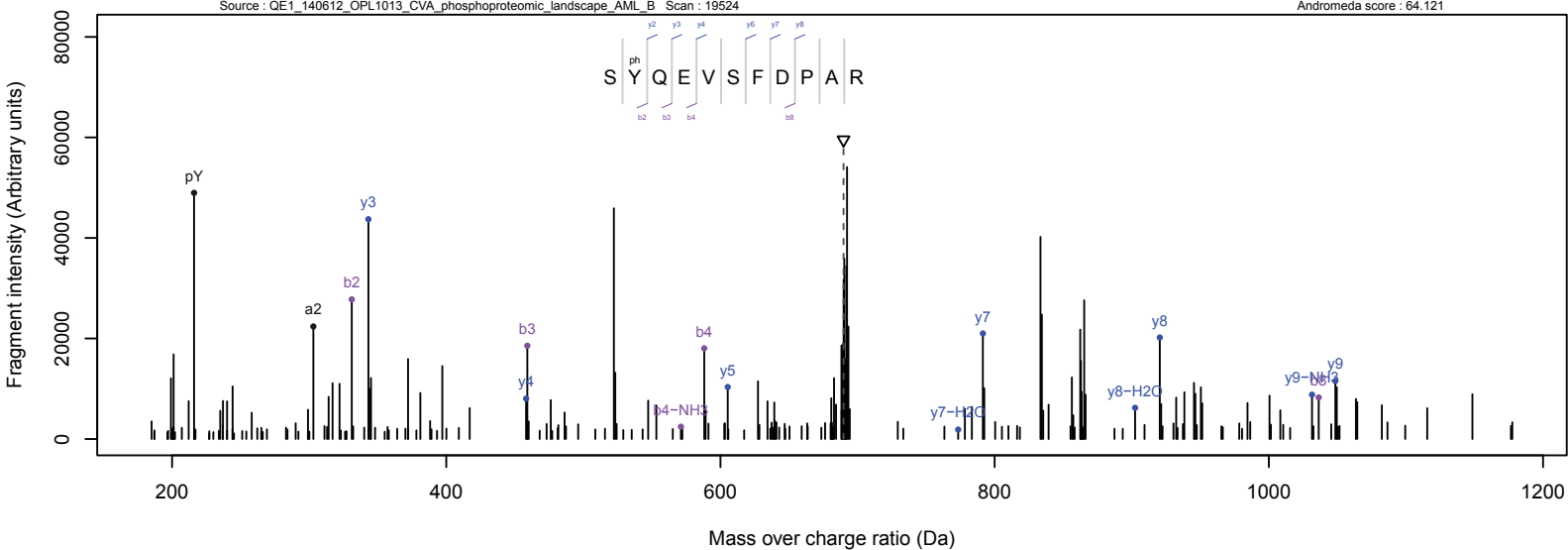
Andromeda score : 64.121



## SY(ph)QEVSFDPAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 19524

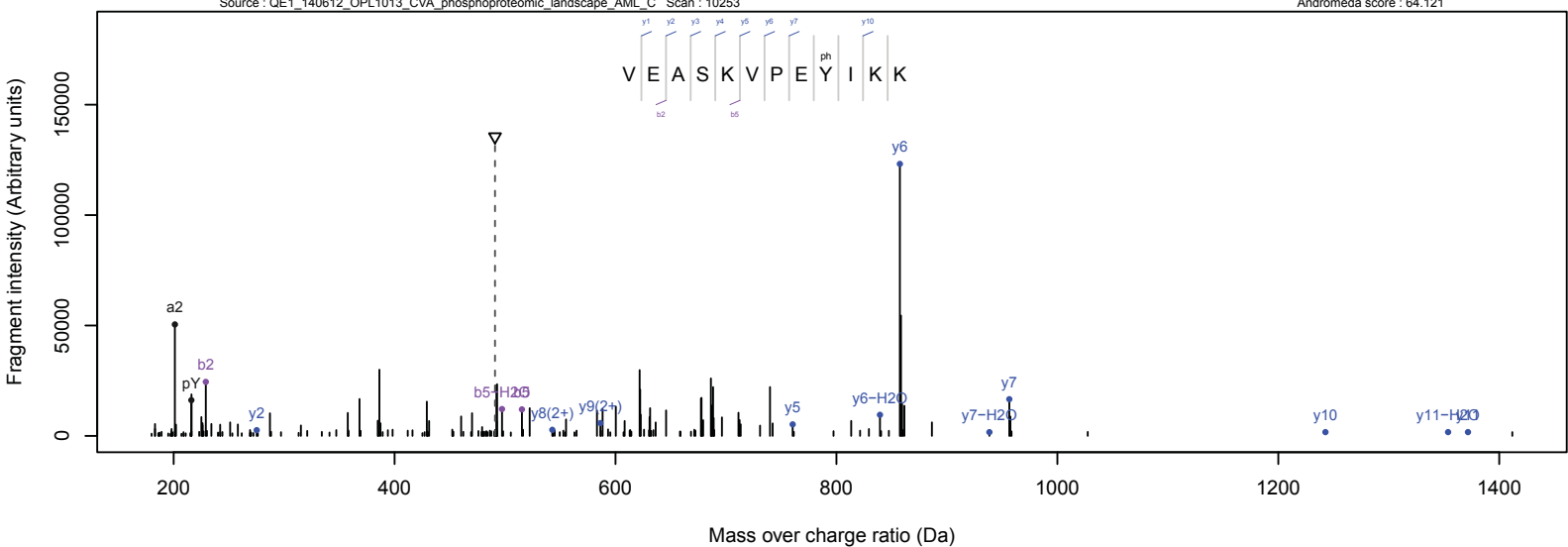
Andromeda score : 64.121



## VEASKVPEY(ph)IKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10253

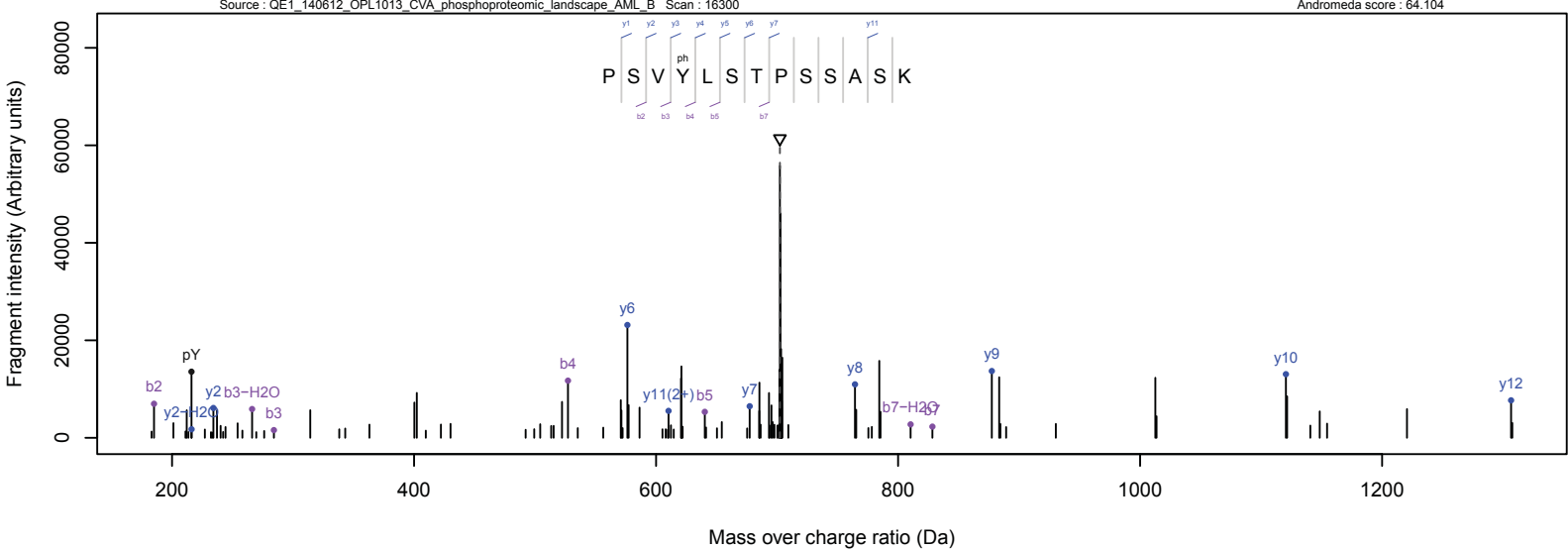
Andromeda score : 64.121



## PSVY(ph)LSTPSSASK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16300

Andromeda score : 64.104

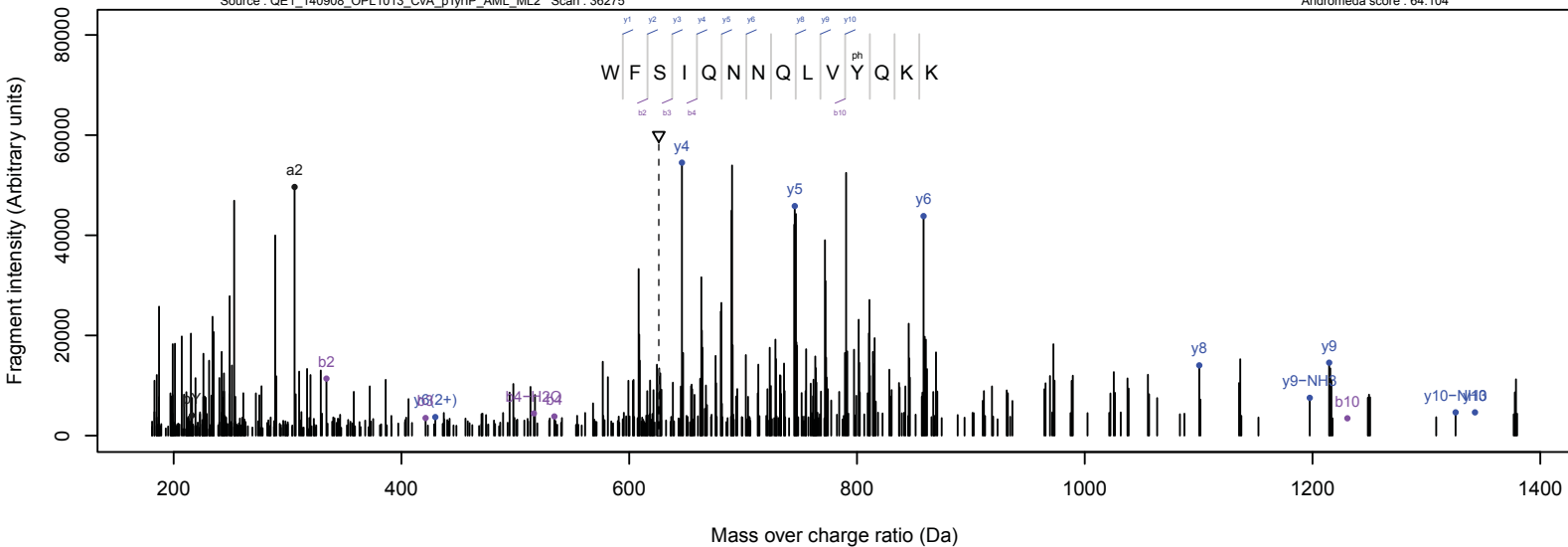




## WFSIQNNQLVY(ph)QKK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_ML2 Scan : 36275

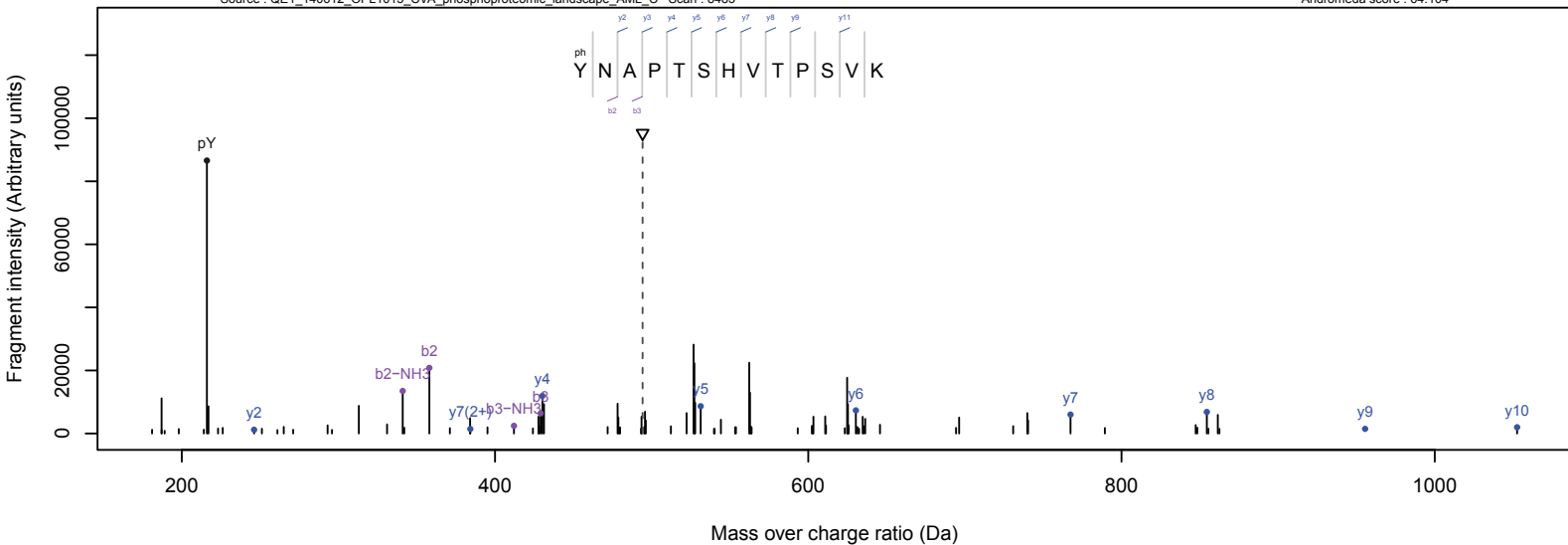
Andromeda score : 64.104



## Y(ph)NAPTSHVTPSVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8483

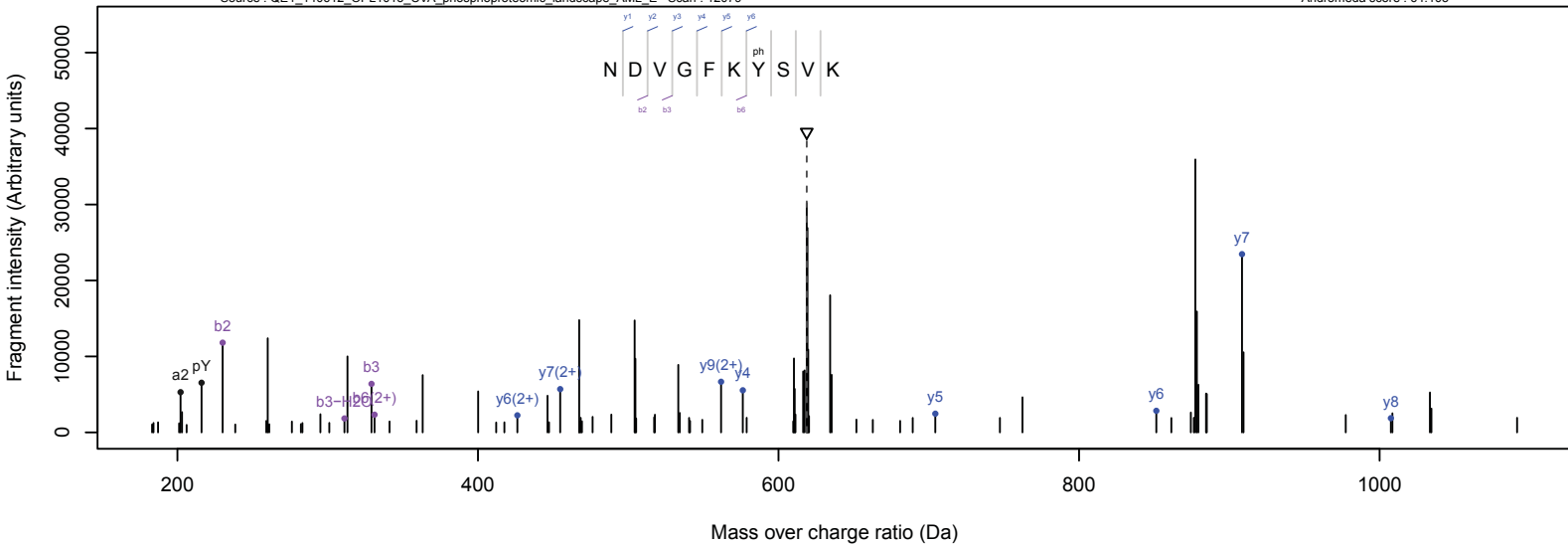
Andromeda score : 64.104



## NDVGFKY(ph)SVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 12079

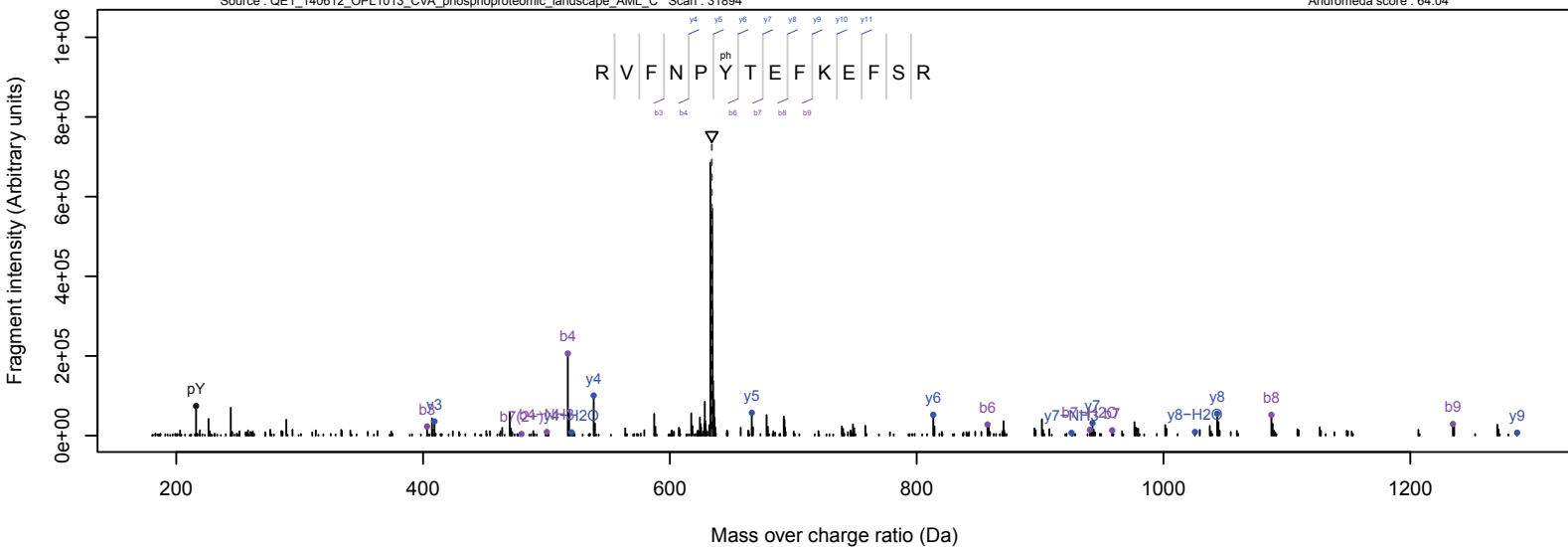
Andromeda score : 64.103



## RVFNPY(ph)TEFKEFSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31894

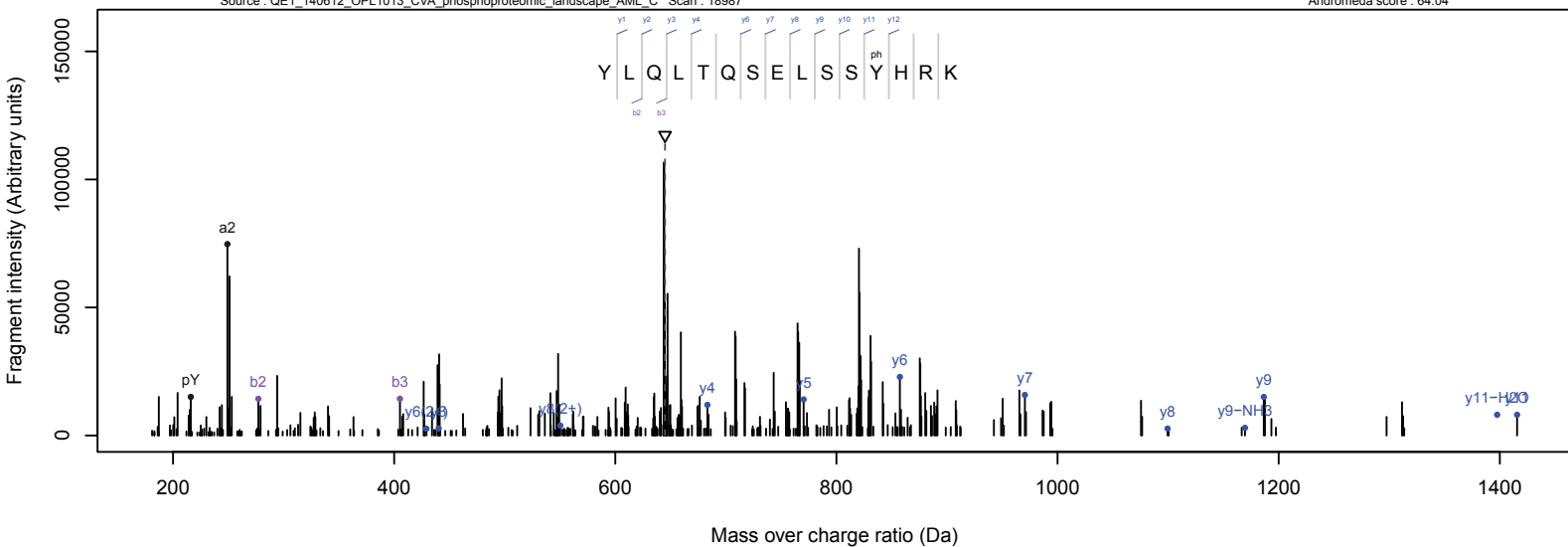
Andromeda score : 64.04



## YLQLTQSELSSY(ph)HRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18987

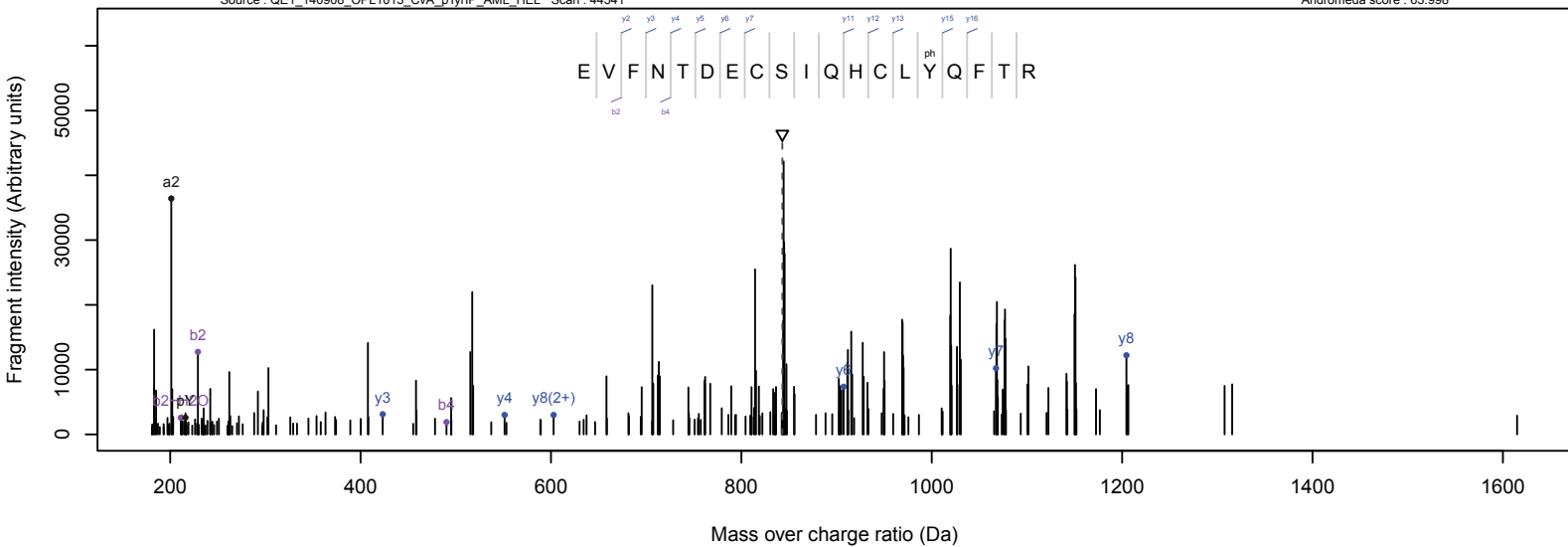
Andromeda score : 64.04



## EVFNTDECSIQHCLY(ph)QFTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 44541

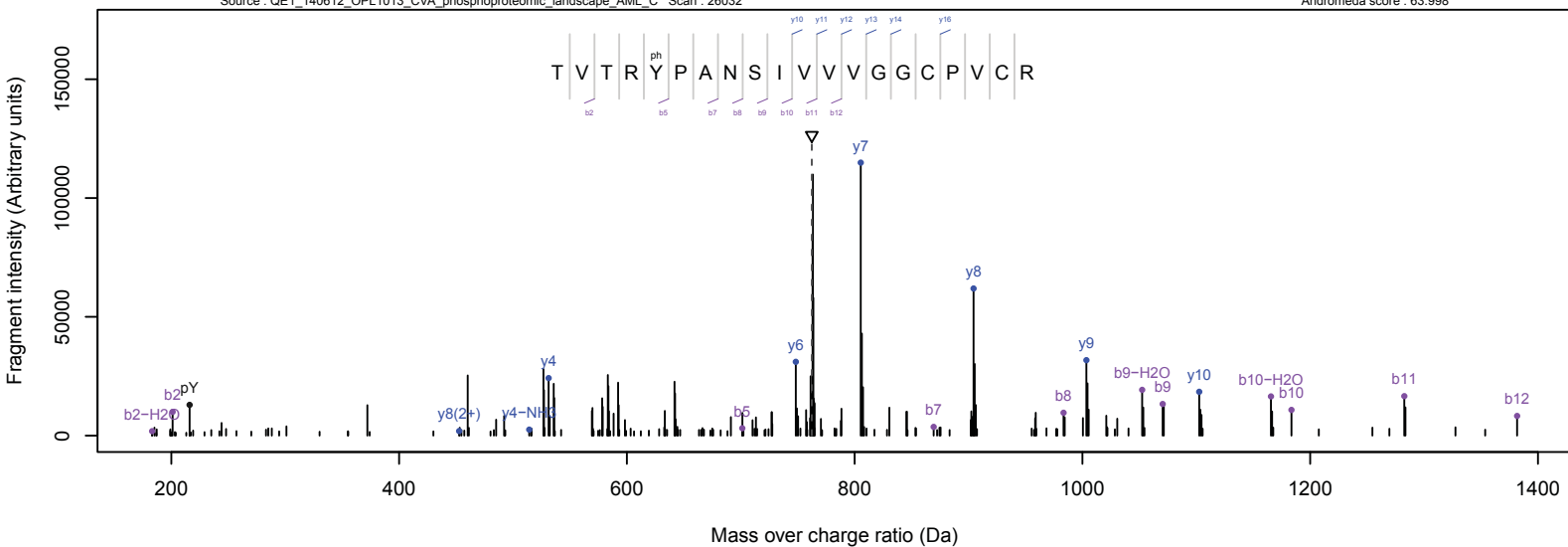
Andromeda score : 63.998



TVTRY(ph)PANSIVVGGCPVCR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26032

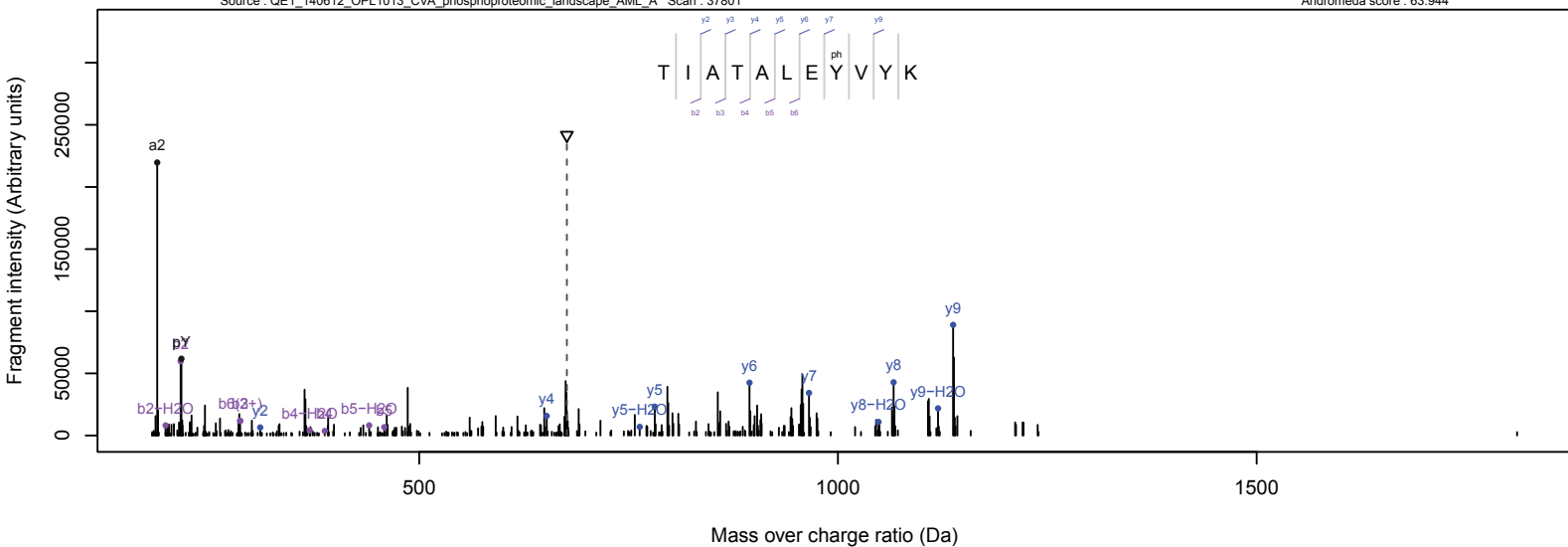
Andromeda score : 63.998



TIATALEY(ph)VYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 37801

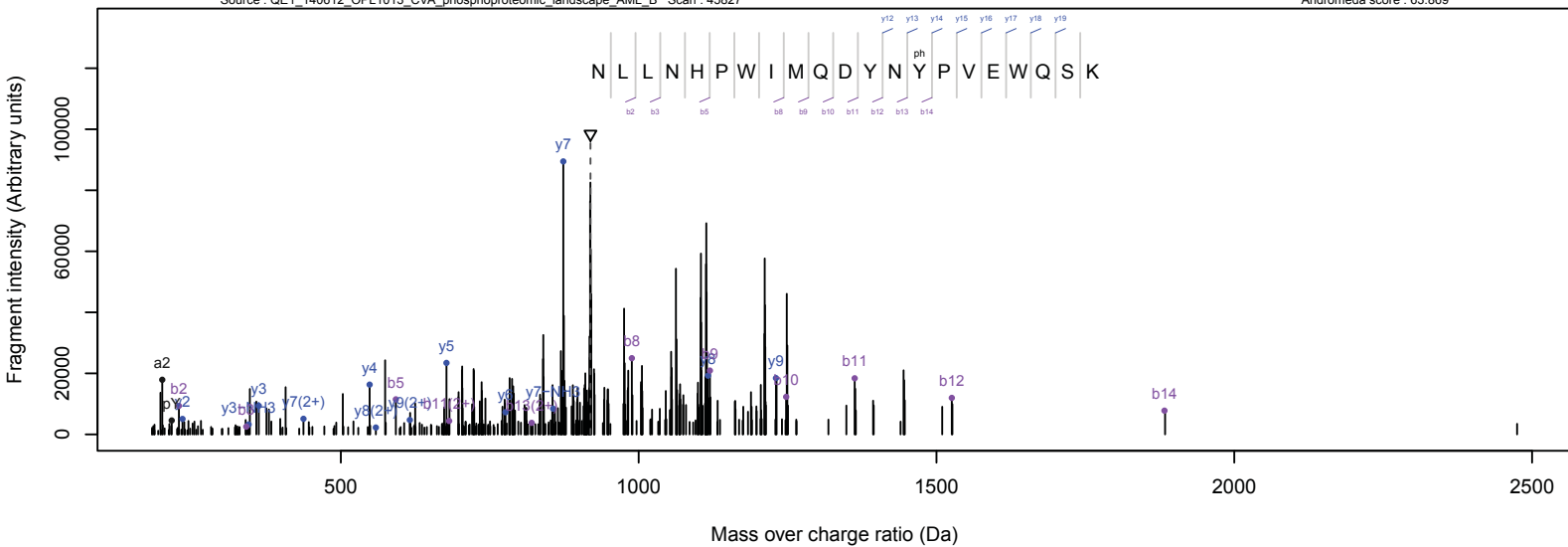
Andromeda score : 63.944



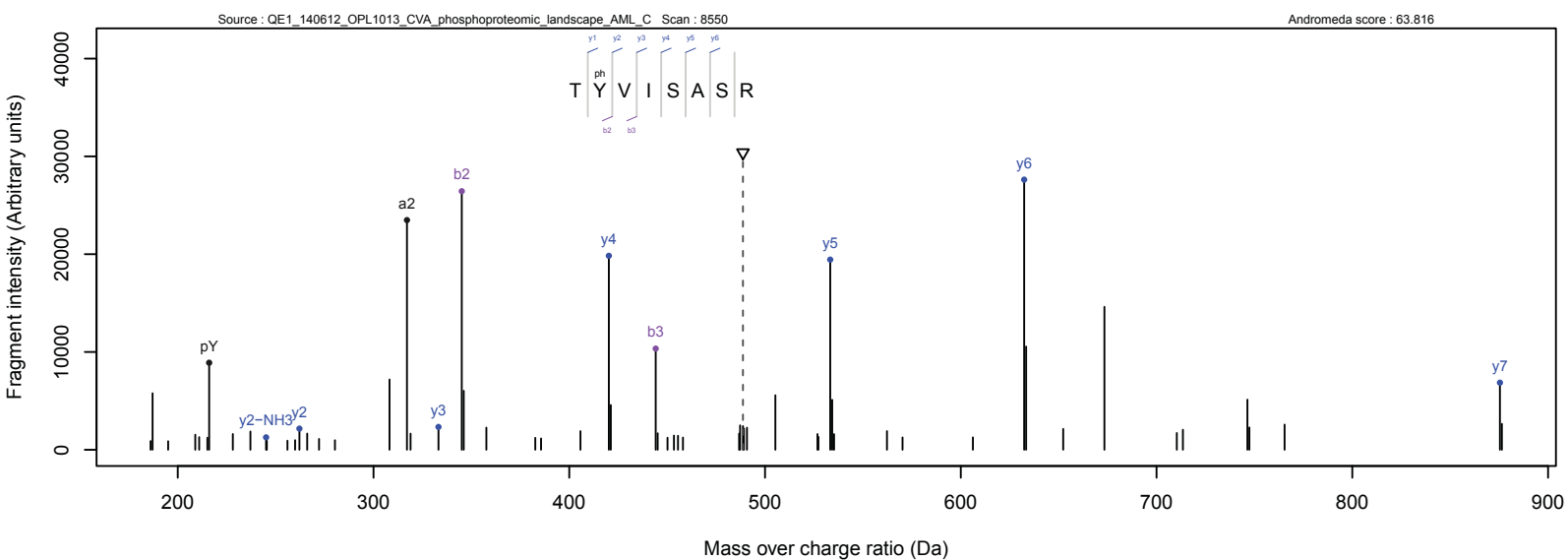
NLLNHPWIMQDYNYPVEWQSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45827

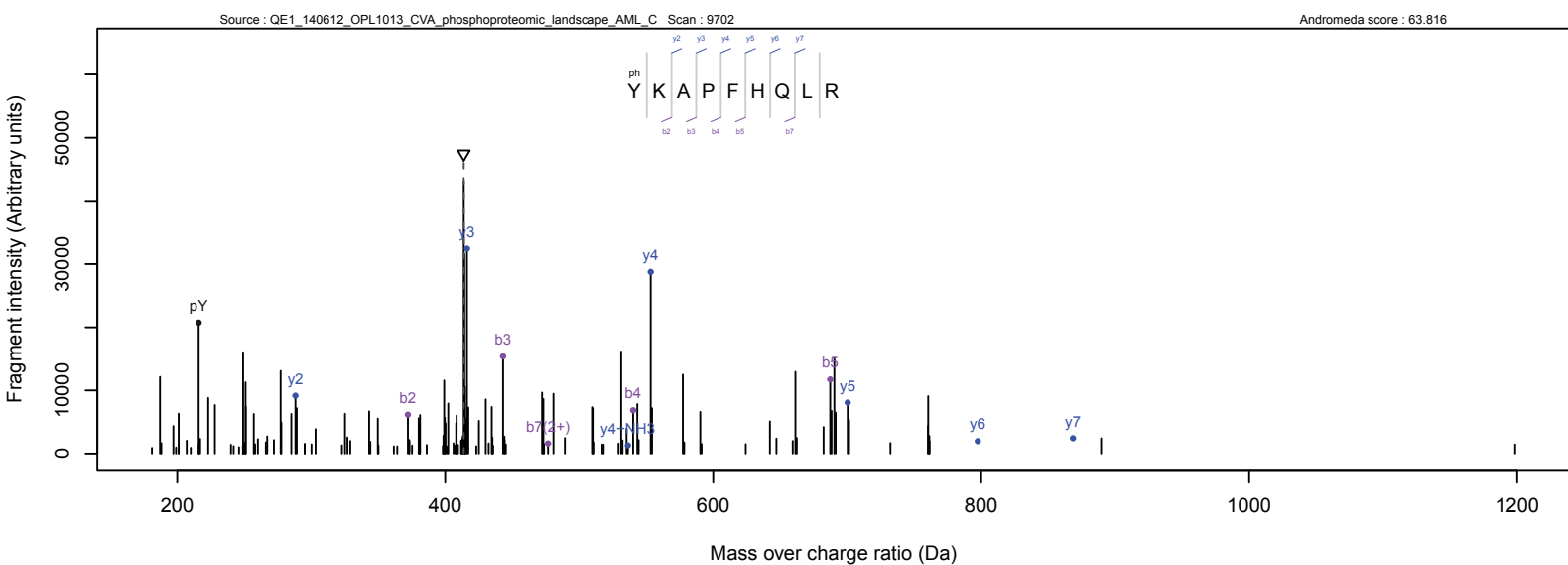
Andromeda score : 63.869



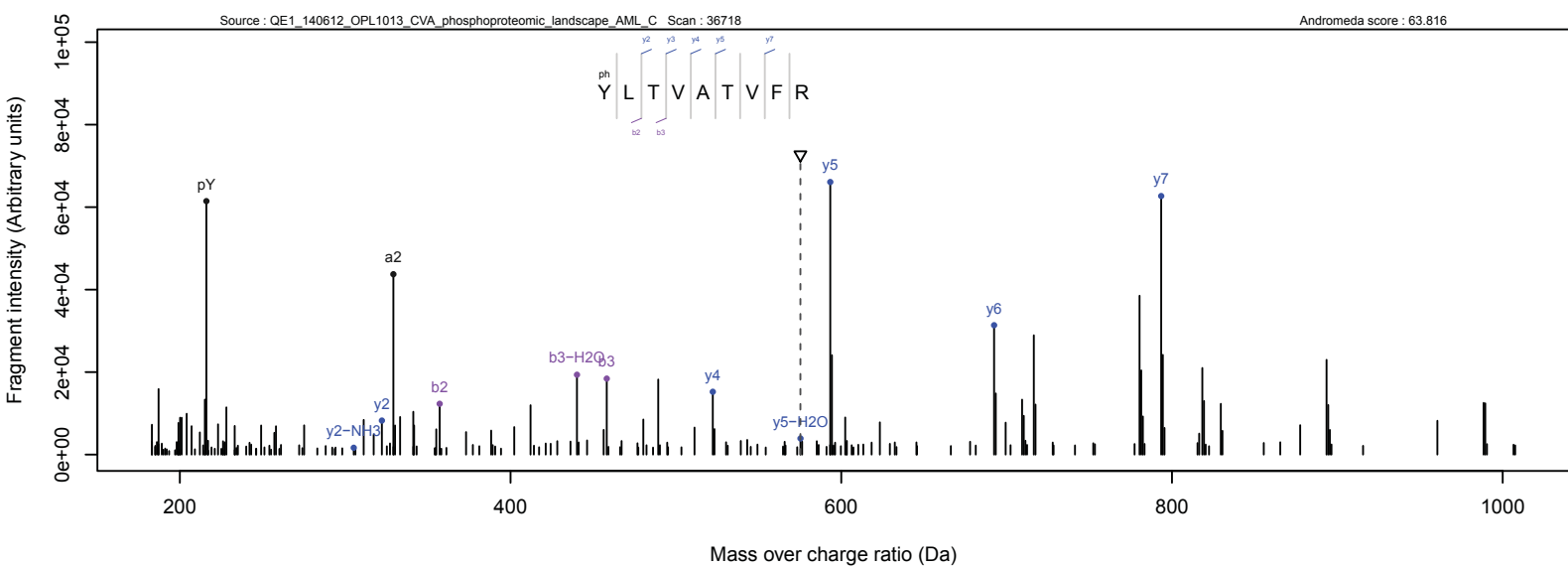
## TY(ph)VISASR-2+



## Y(ph)KAPFHQLR-3+



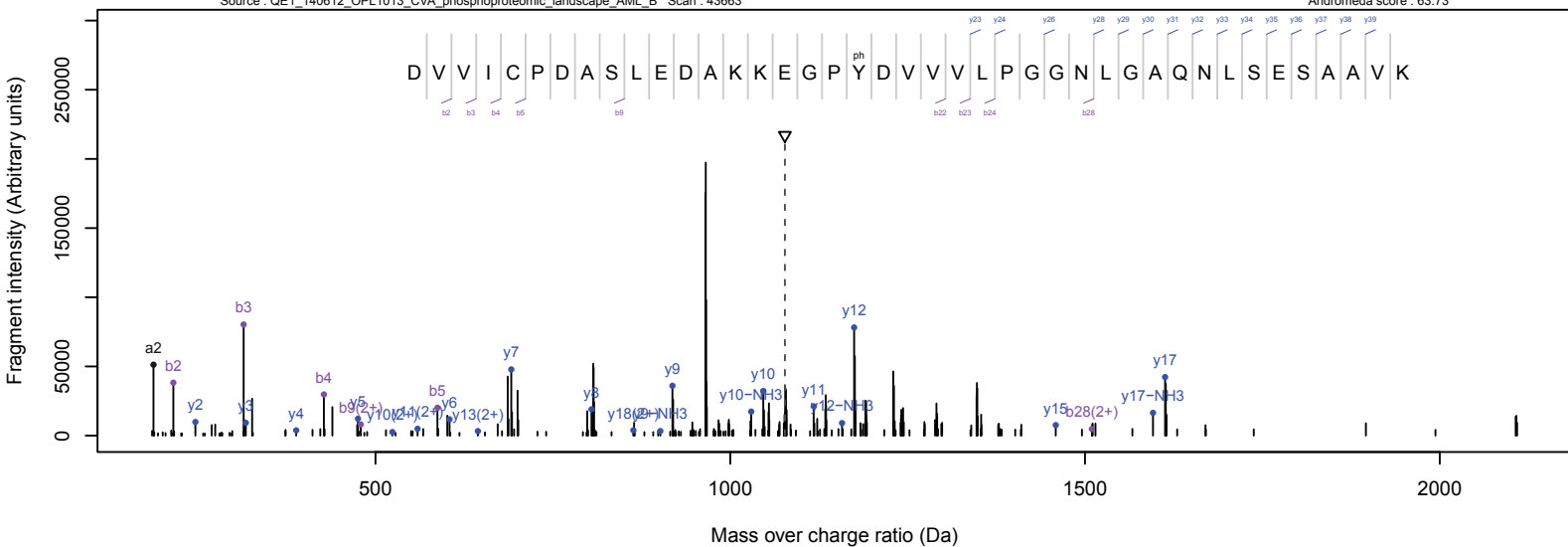
## Y(ph)LTVATVFR-2+



DVVICPDASLEDAKKEGY(ph)DVVVLPGGNLGAQNLSESAVK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43663

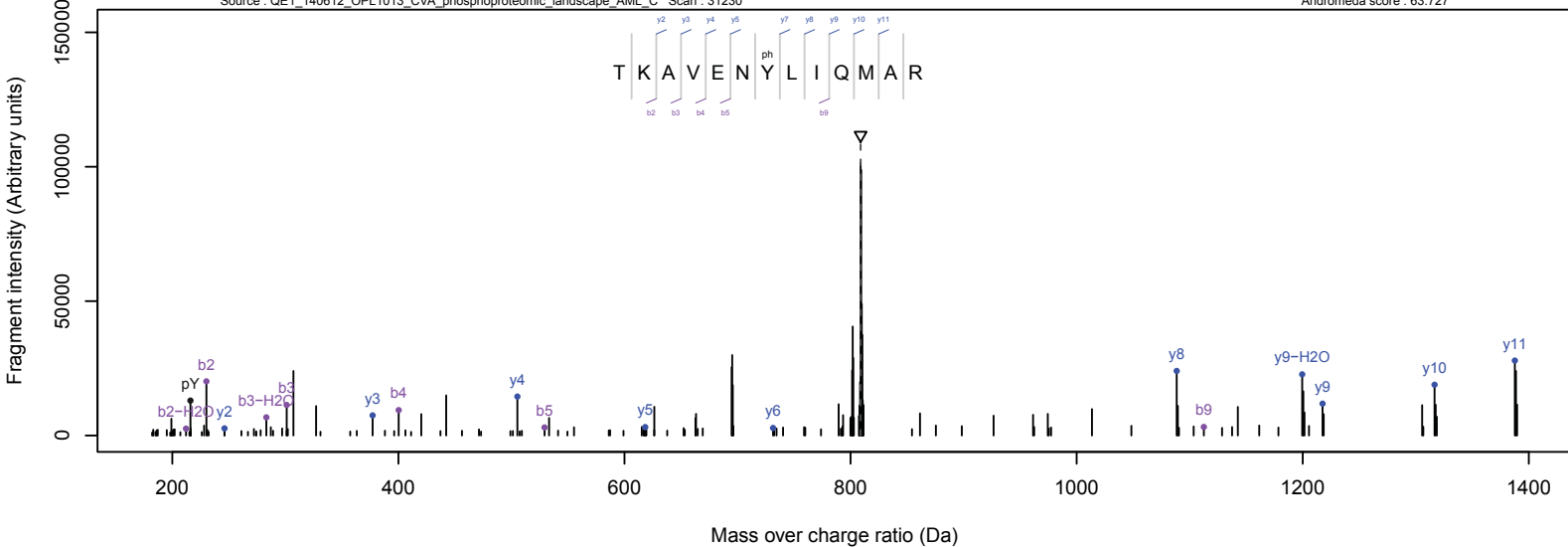
Andromeda score : 63.73



TKAVENY(ph)LIQMAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31230

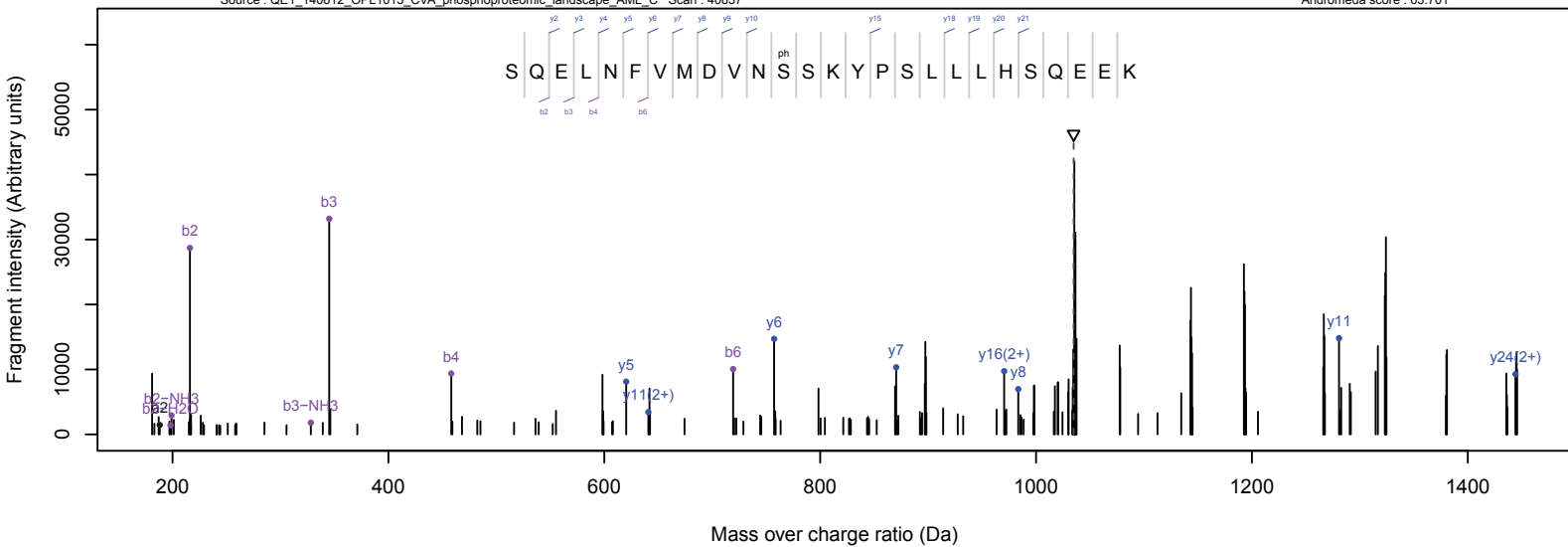
Andromeda score : 63.727



SQELNFVMDVNS(ph)SKYPSLLLSQEEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40837

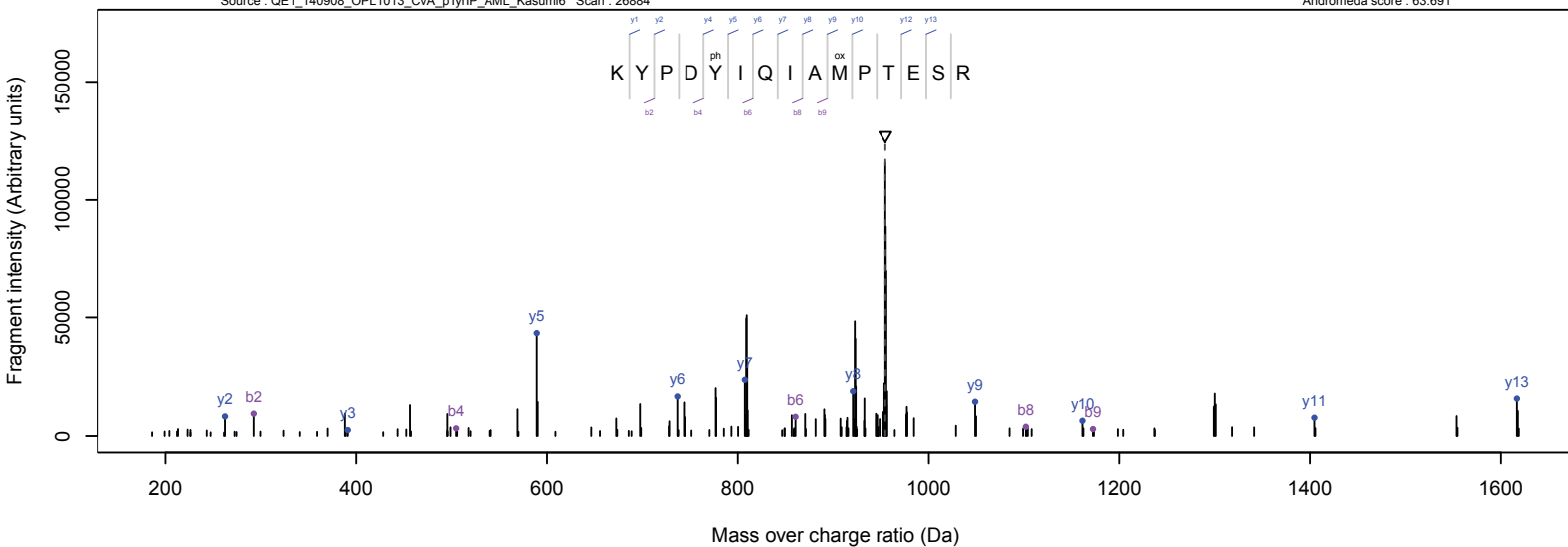
Andromeda score : 63.701



## KYPDY(ph)IQIAMPTESR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 26884

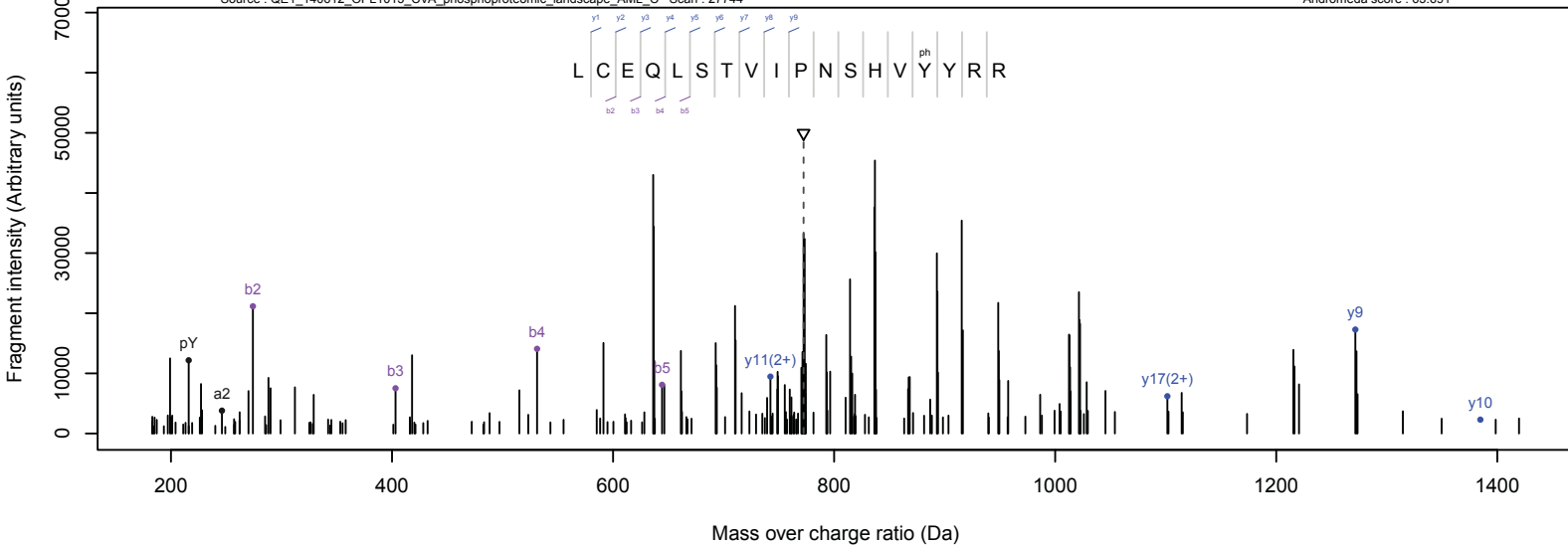
Andromeda score : 63.691



## LCEQLSTVIPNSHVY(ph)YRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27744

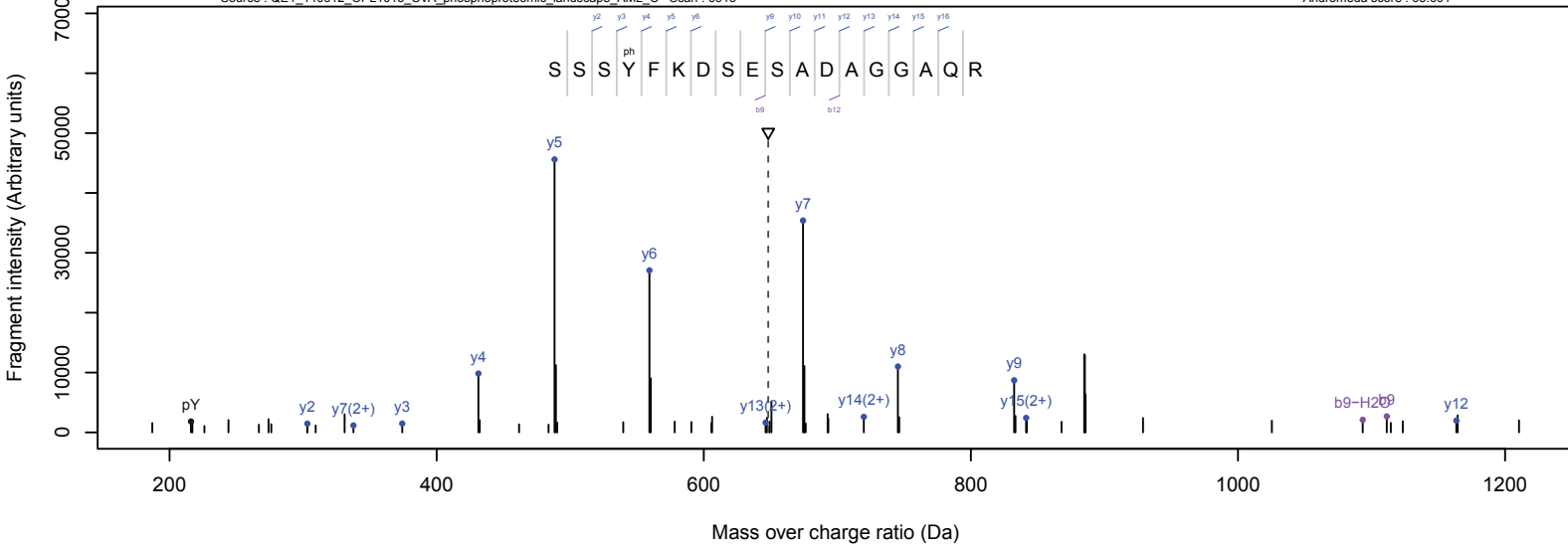
Andromeda score : 63.691



## SSSY(ph)FKDSESADAGGAQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6915

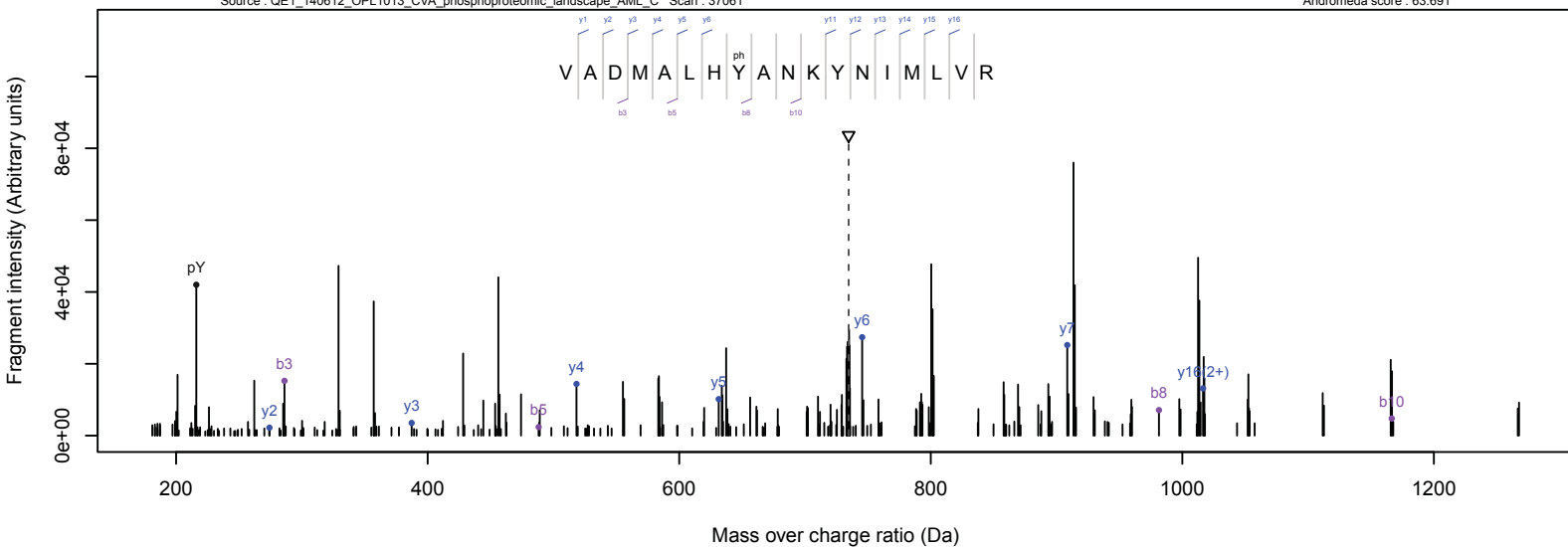
Andromeda score : 63.691



## VADMALHY(ph)ANKYNILVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37061

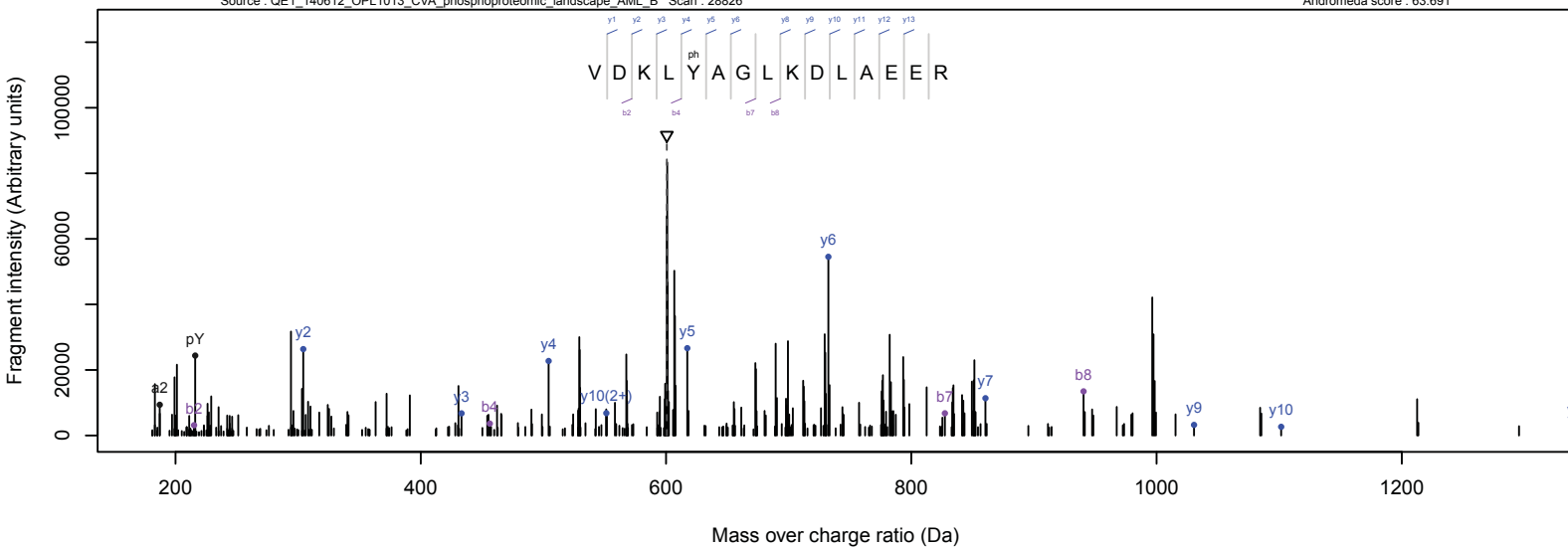
Andromeda score: 63.691



## VDKLY(ph)AGLKDLAER-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 28826

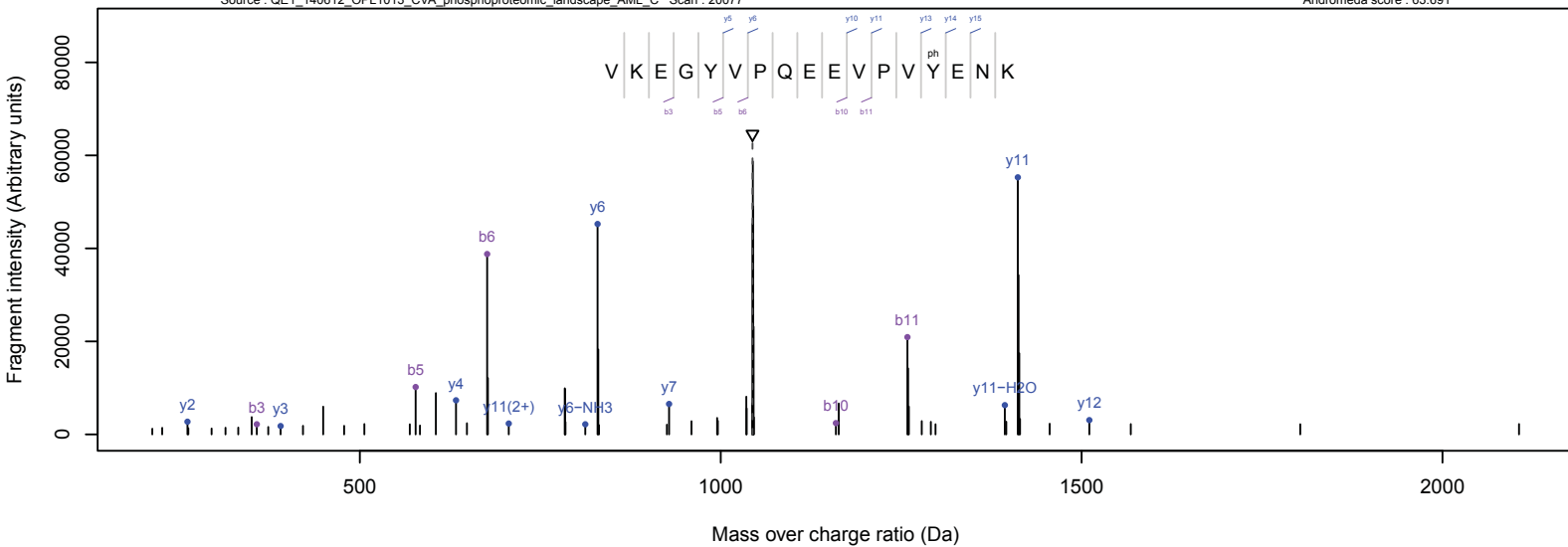
Andromeda score: 63.691



## VKEGYVPQEEV(ph)ENK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 20077

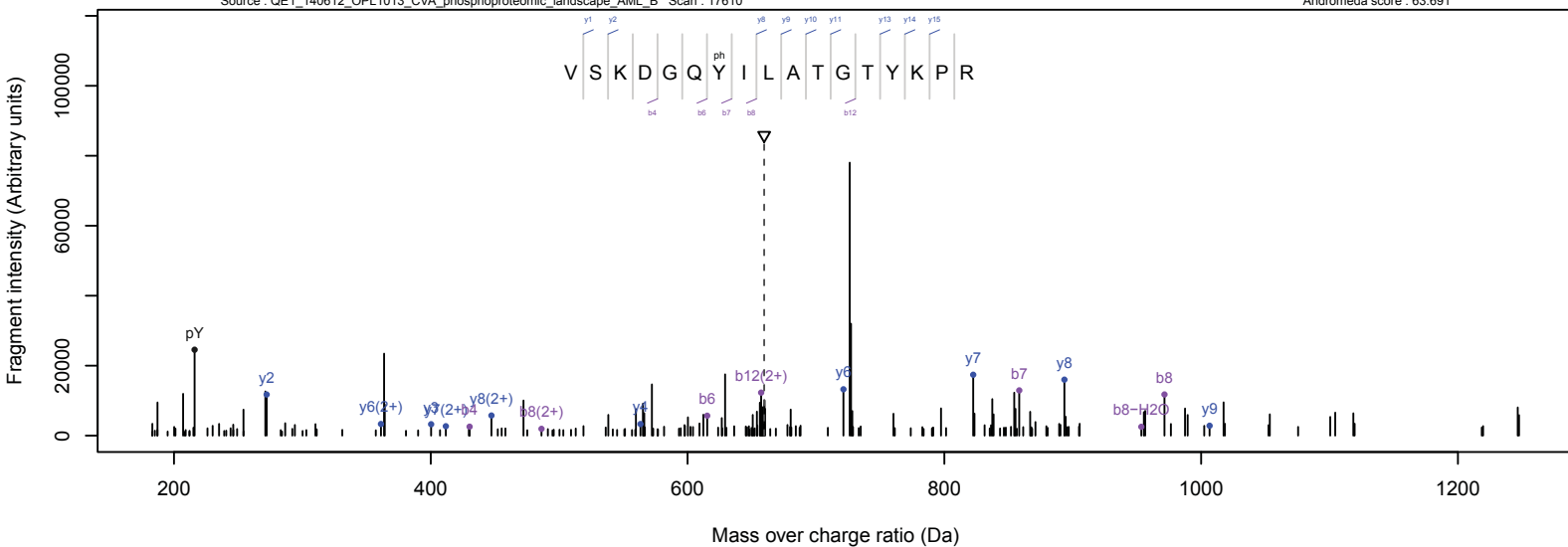
Andromeda score: 63.691



## VSKDGQY(ph)ILATGTYKPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17610

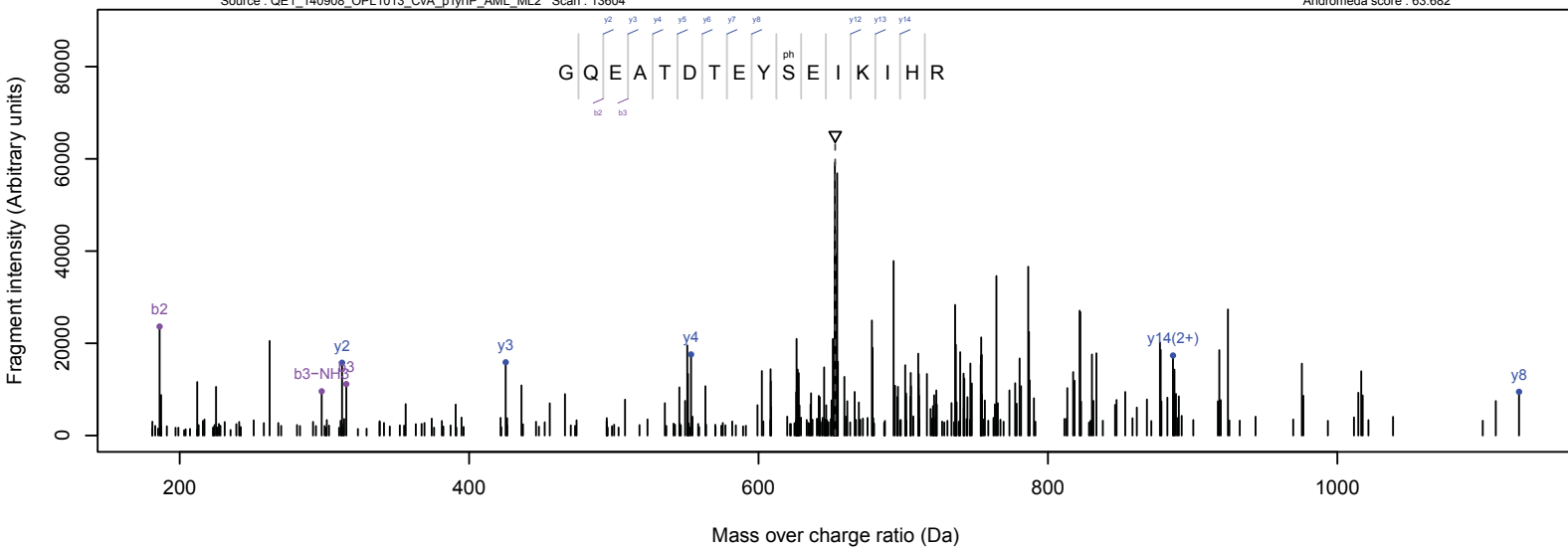
Andromeda score : 63.691



## GQEATDTEYS(ph)EIKIHR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 13604

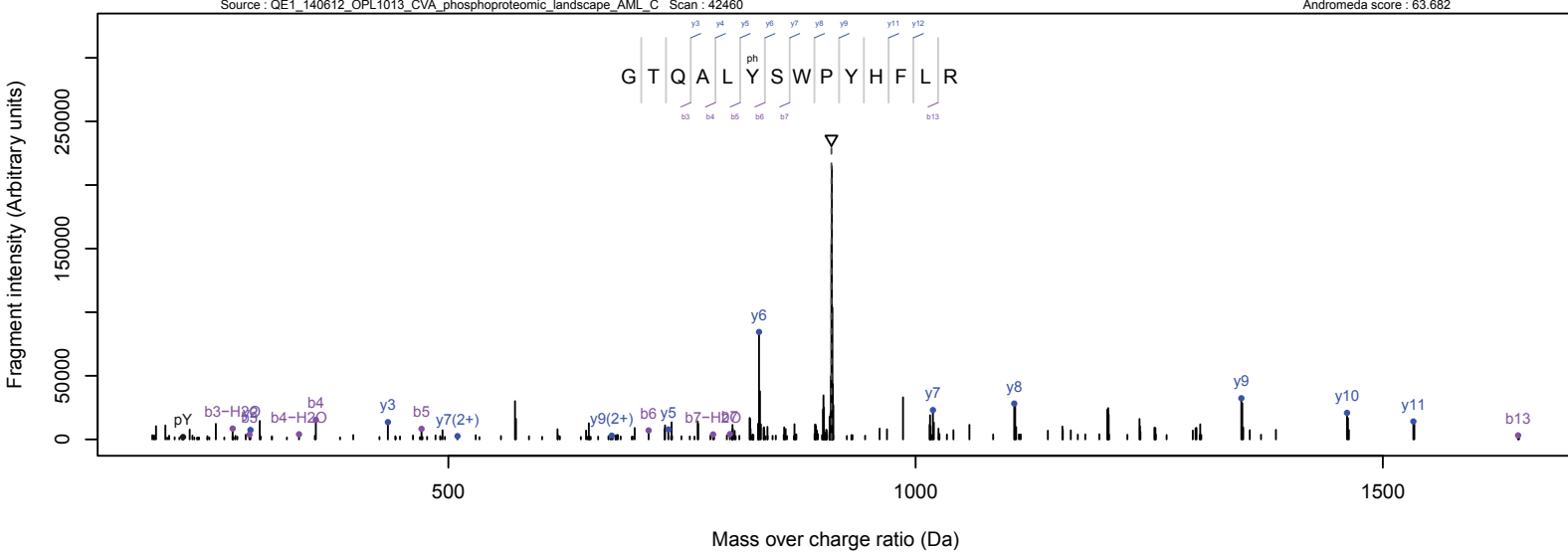
Andromeda score : 63.682



## GTQALY(ph)SWPYHFLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42460

Andromeda score : 63.682

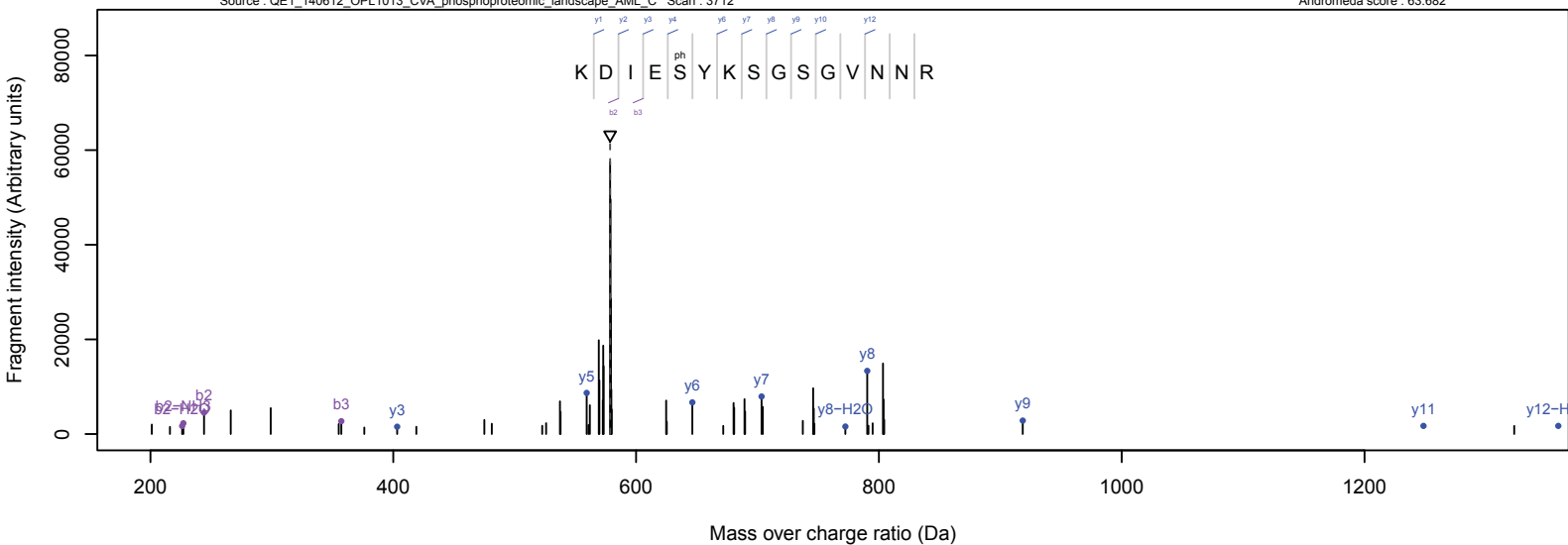




**KDIES(ph)YKSGSGVNNR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3712

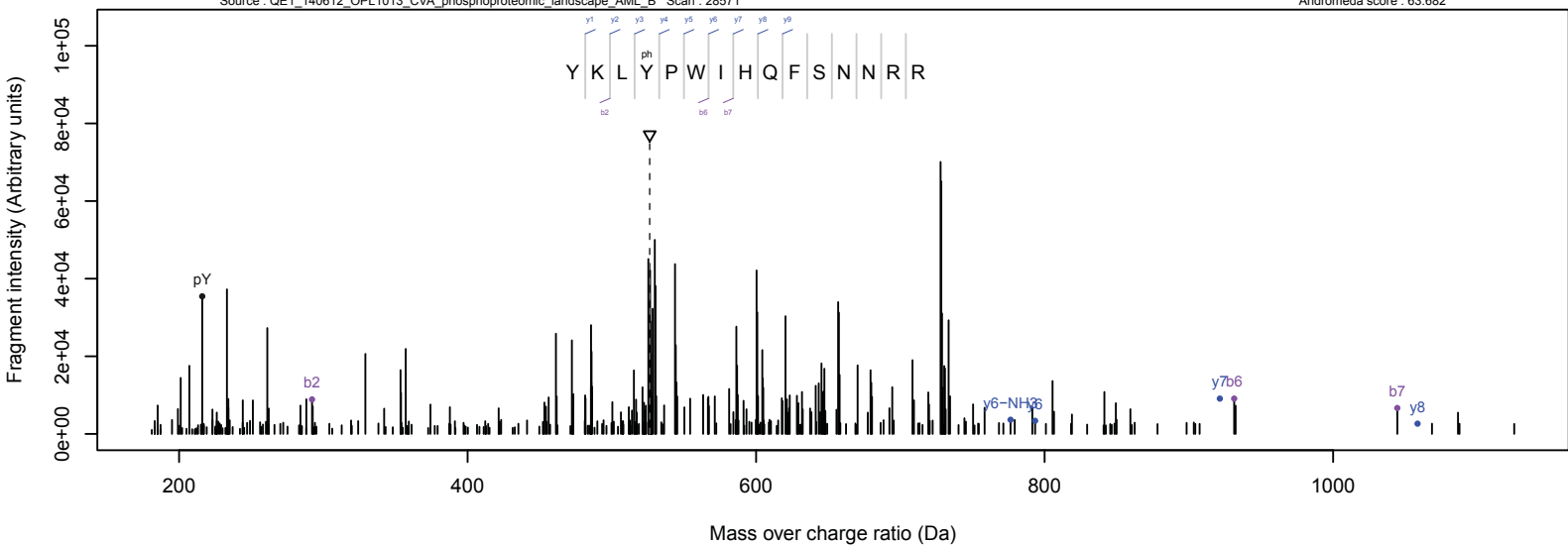
Andromeda score : 63.682



**YKLY(ph)PWIHQFSNNRR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28571

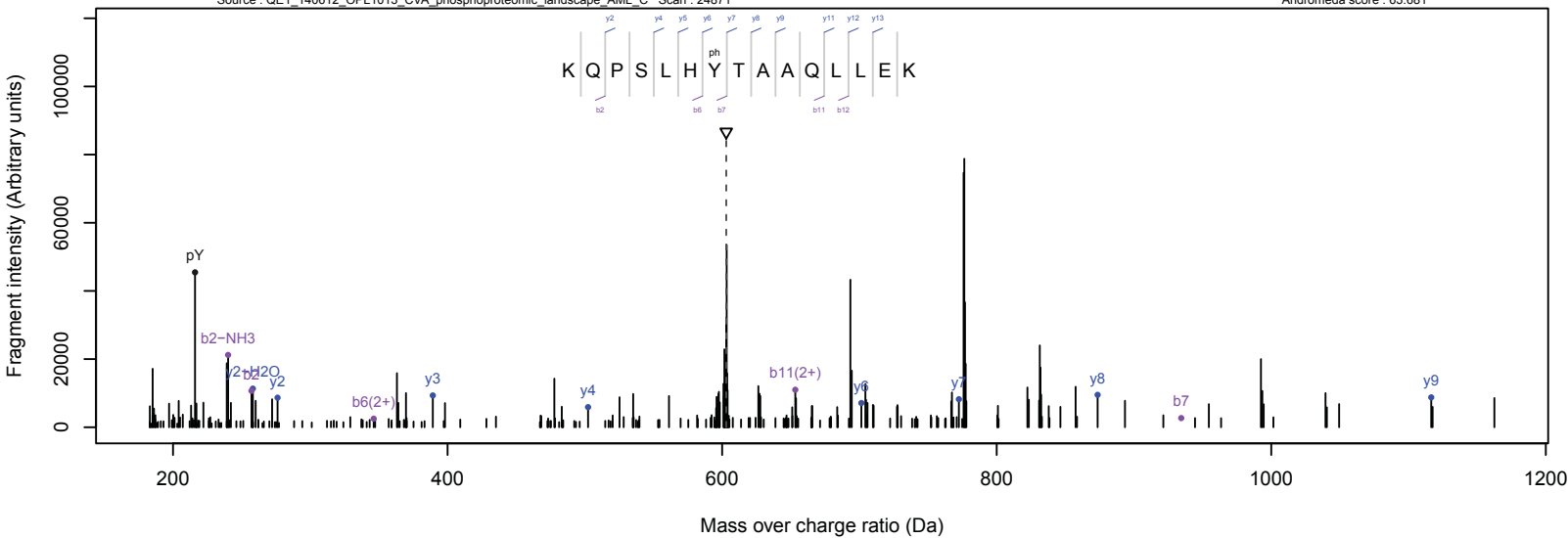
Andromeda score : 63.682

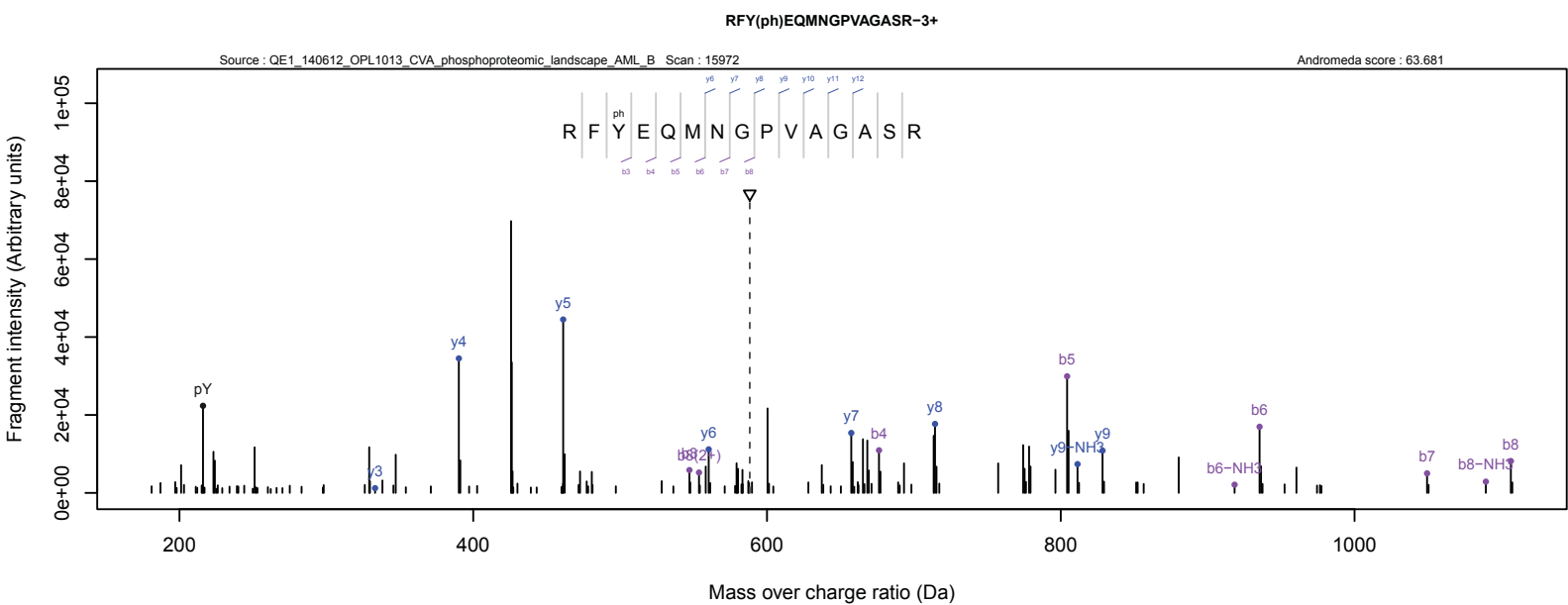
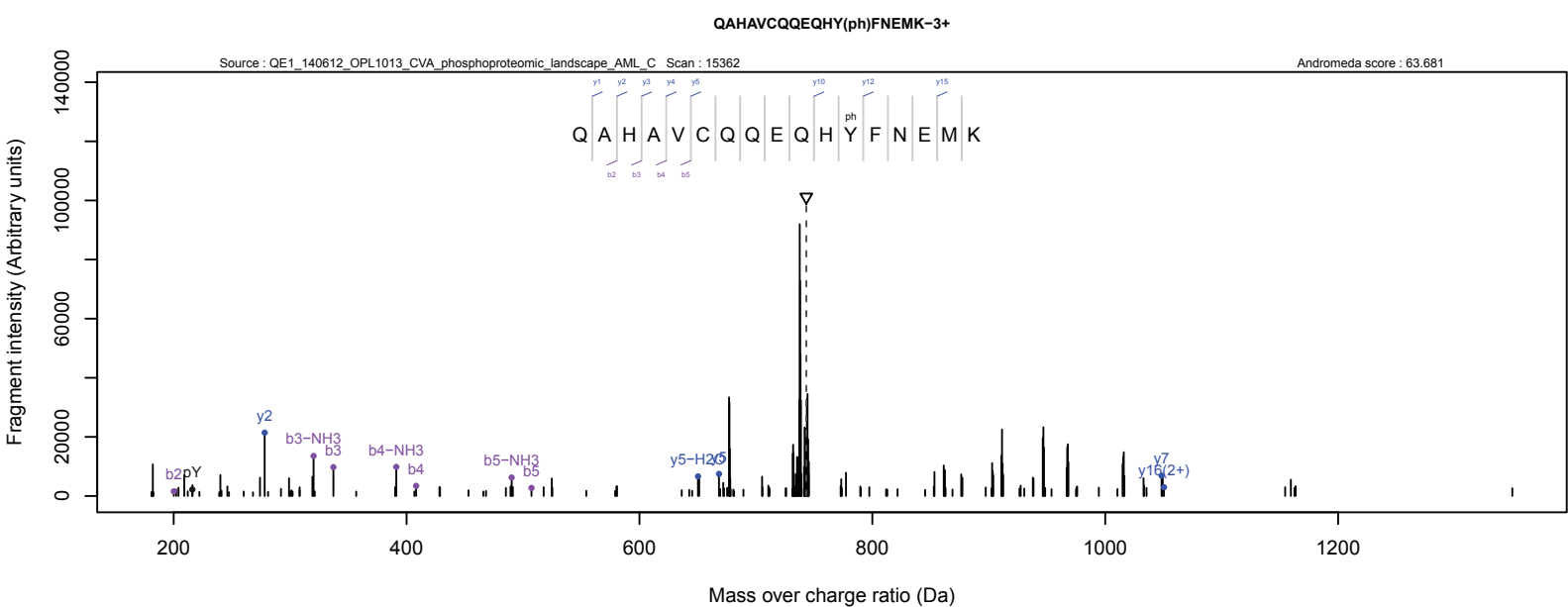
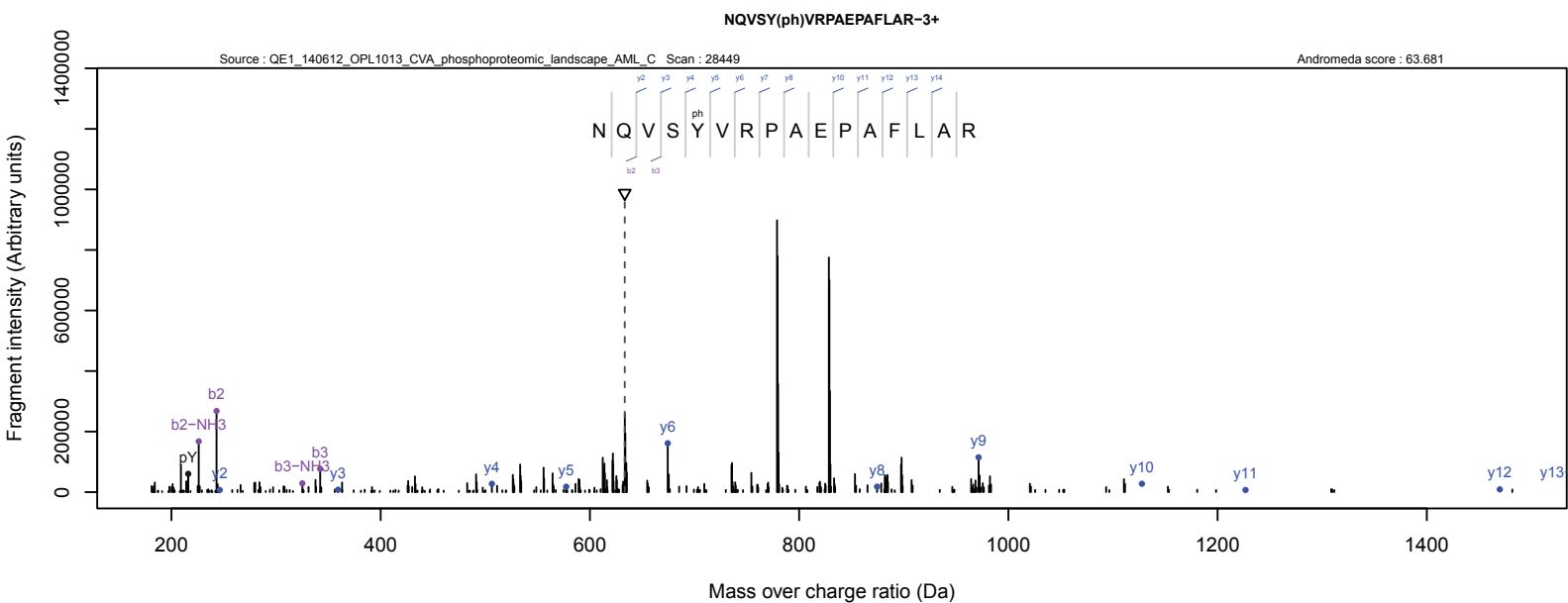


**KQPSLHY(ph)TAAQLLEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24871

Andromeda score : 63.681

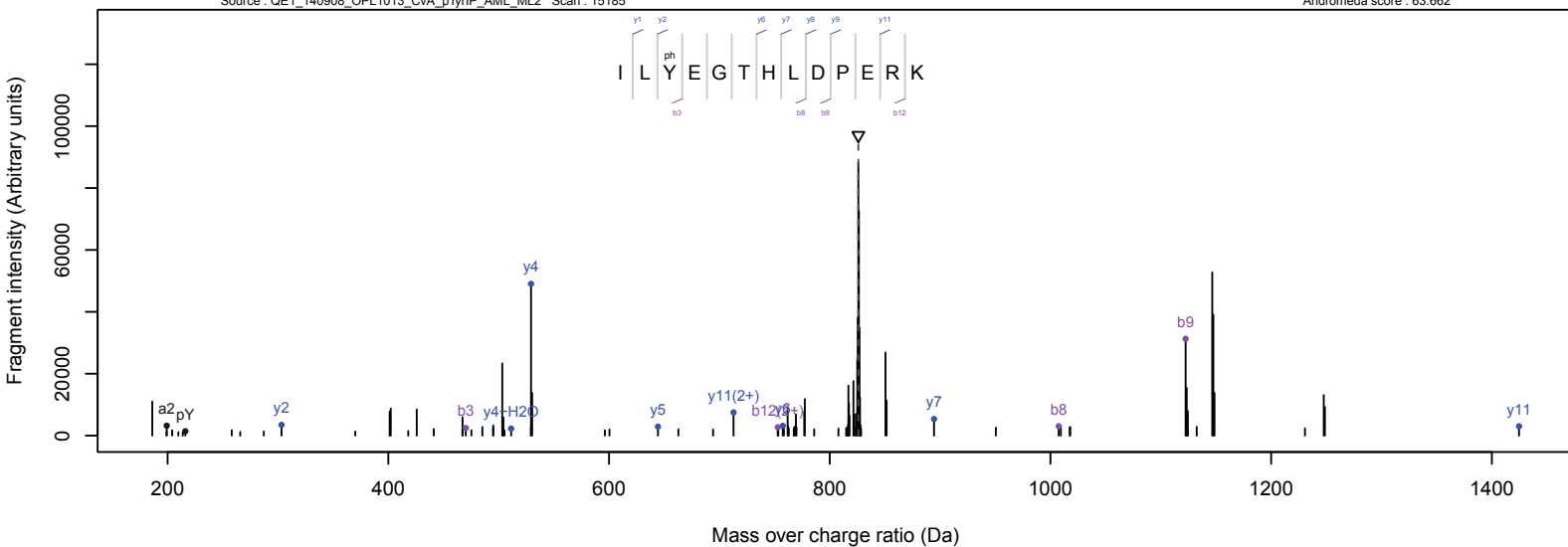




## ILY(ph)EGTHLDPERK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 15185

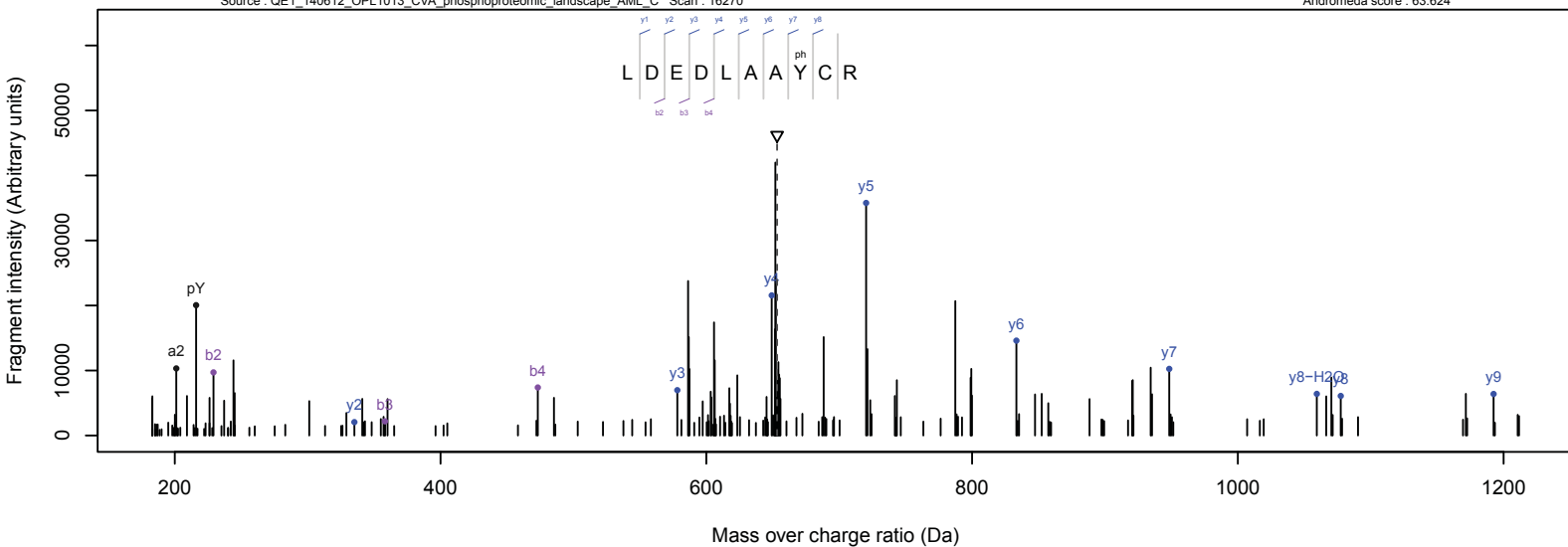
Andromeda score : 63.662



## LDEDLAAY(ph)CR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16270

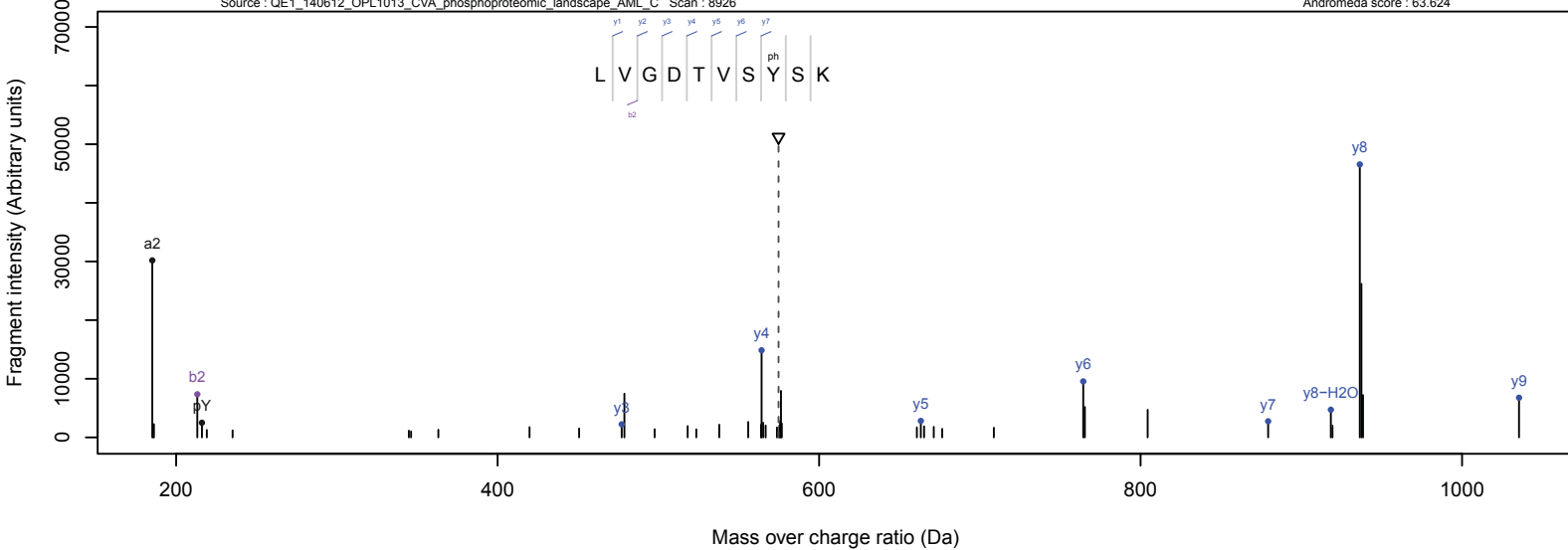
Andromeda score : 63.624



## LVGDTVSY(ph)SK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8926

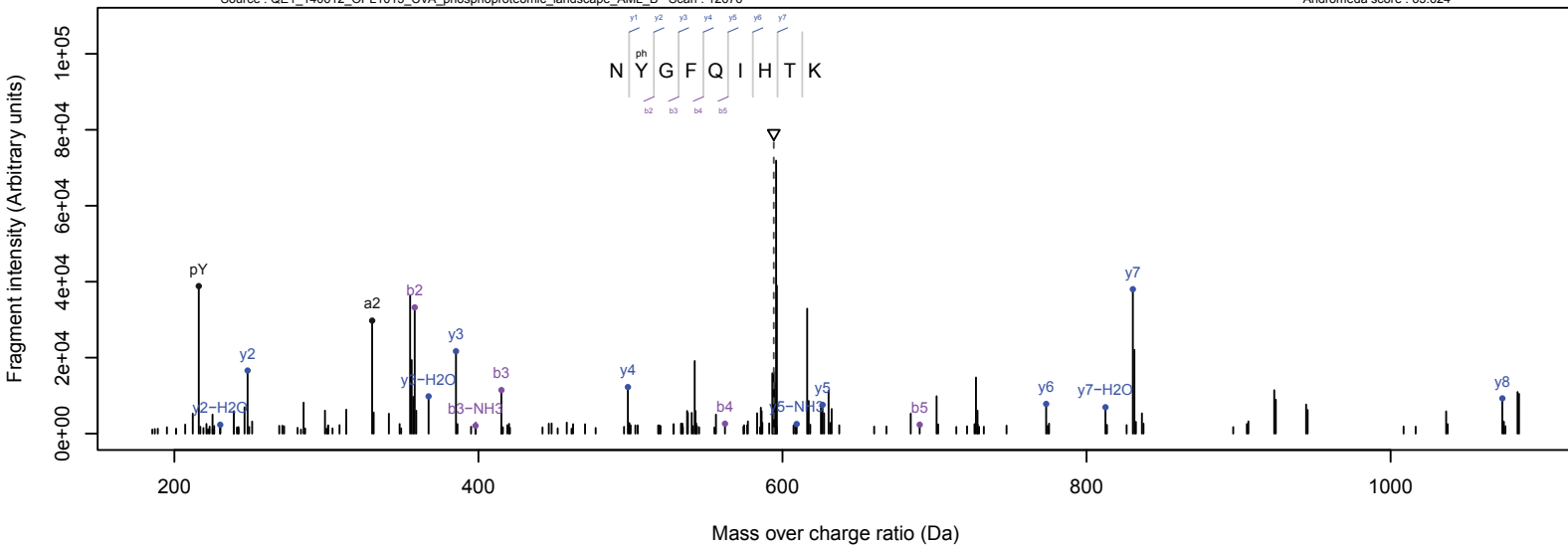
Andromeda score : 63.624



## NY(ph)GFQIHTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12676

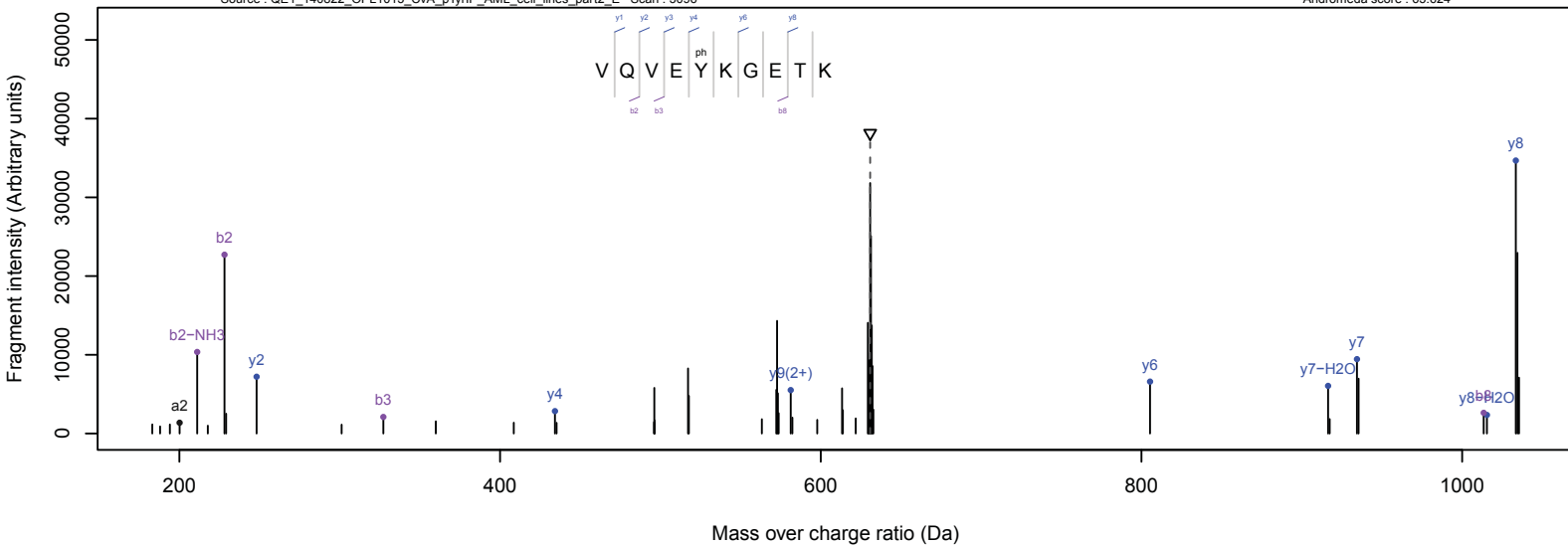
Andromeda score : 63.624



## VQVEY(ph)KGETK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 3096

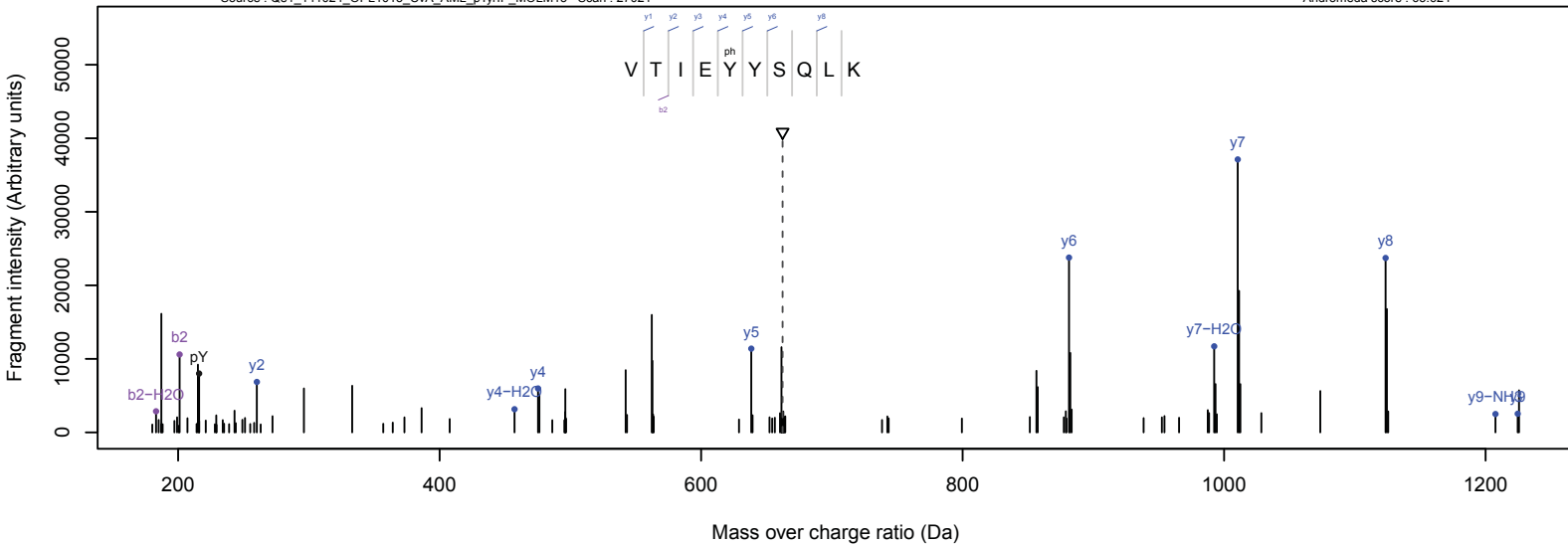
Andromeda score : 63.624



## VTIEY(ph)YSQLK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 27021

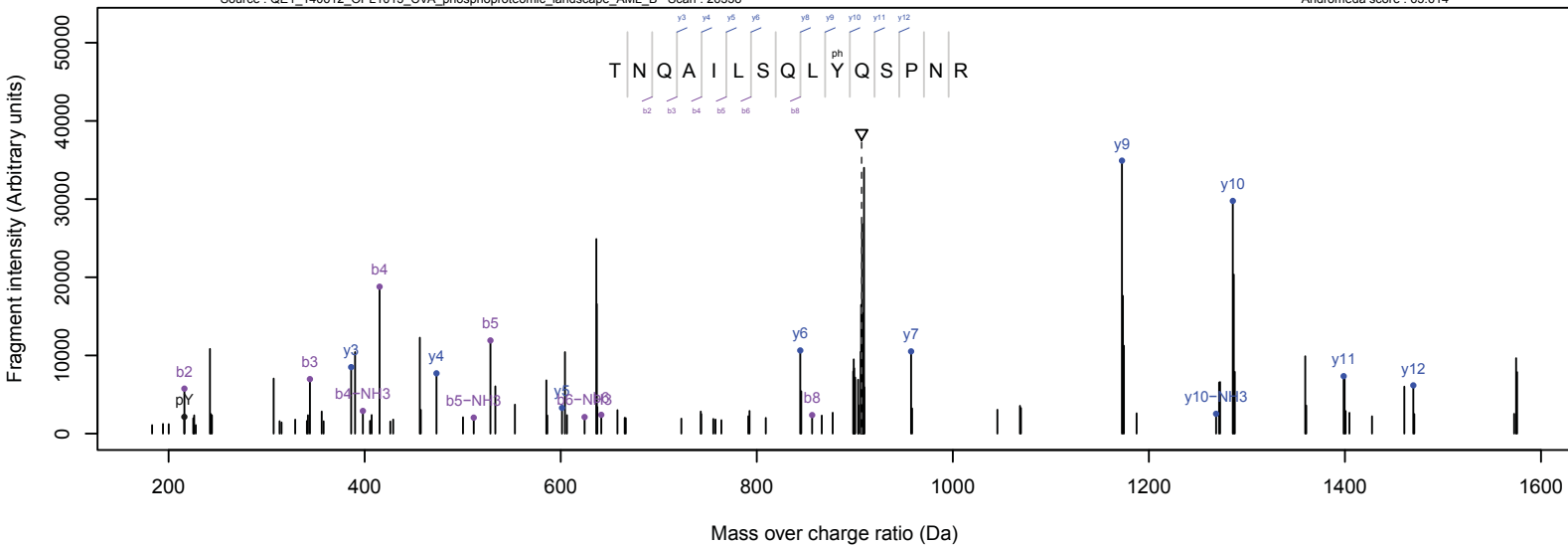
Andromeda score : 63.624



TNQAILSQLY(ph)QSPNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26338

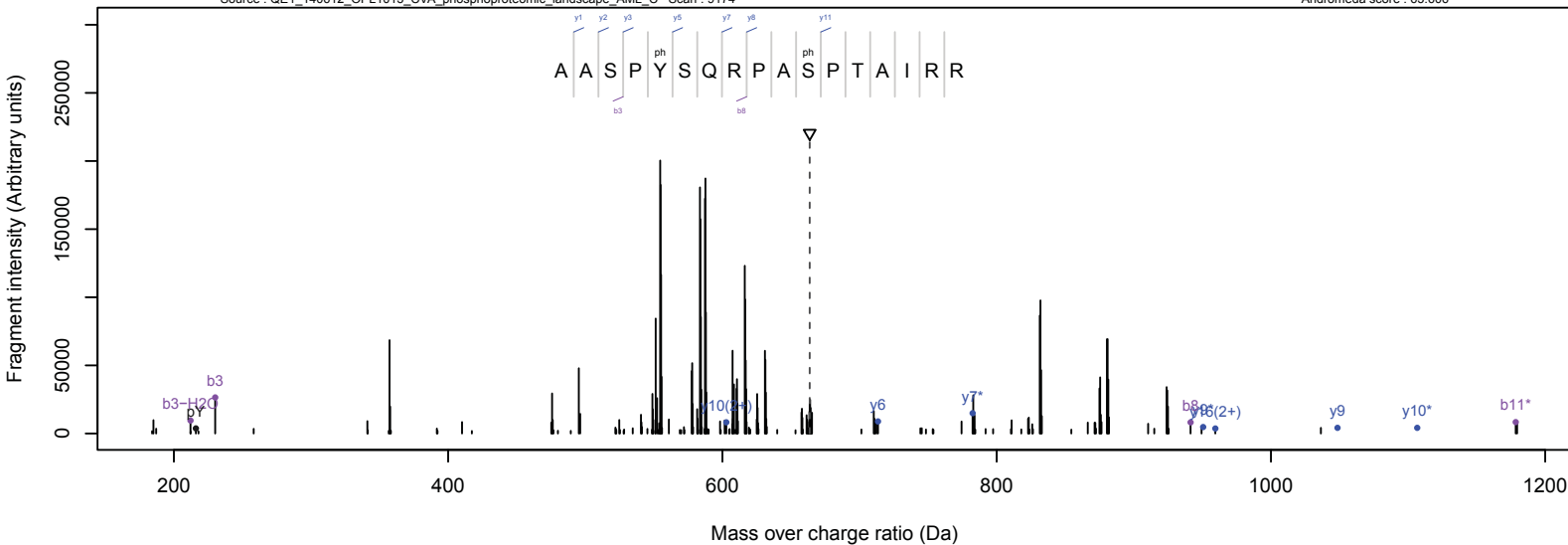
Andromeda score : 63.614



AASPY(ph)SQRPAS(ph)PTAIRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9174

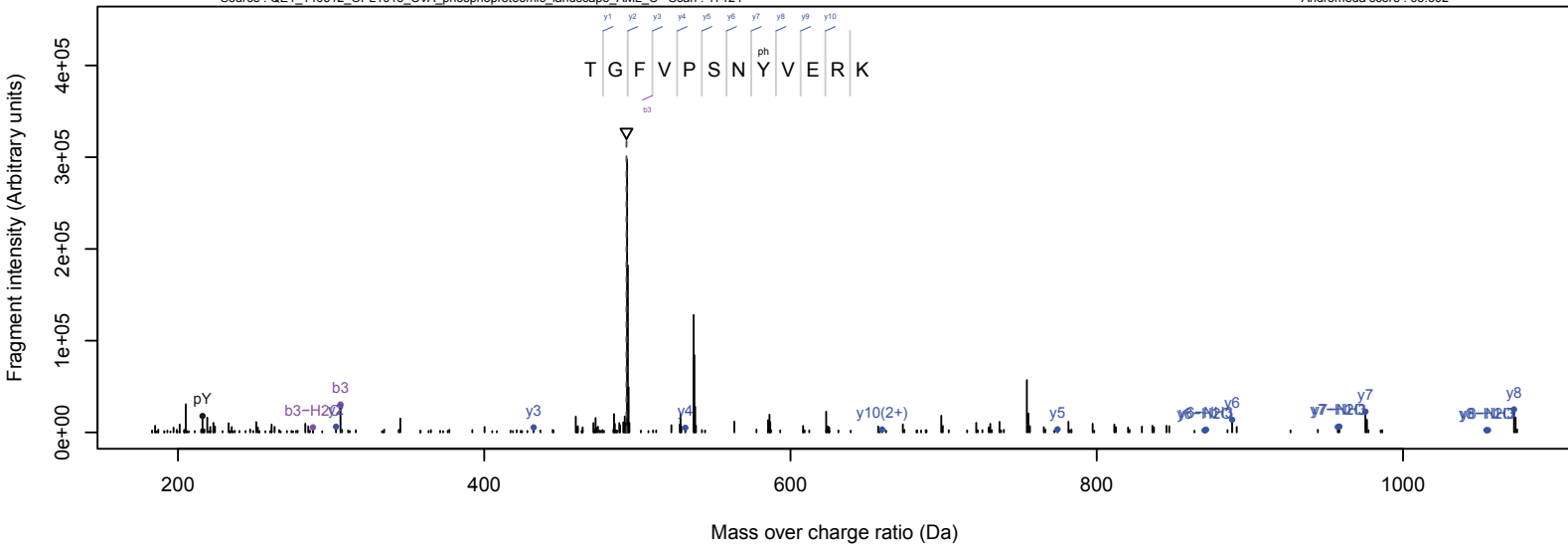
Andromeda score : 63.606



TGFVPSNY(ph)VERK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17124

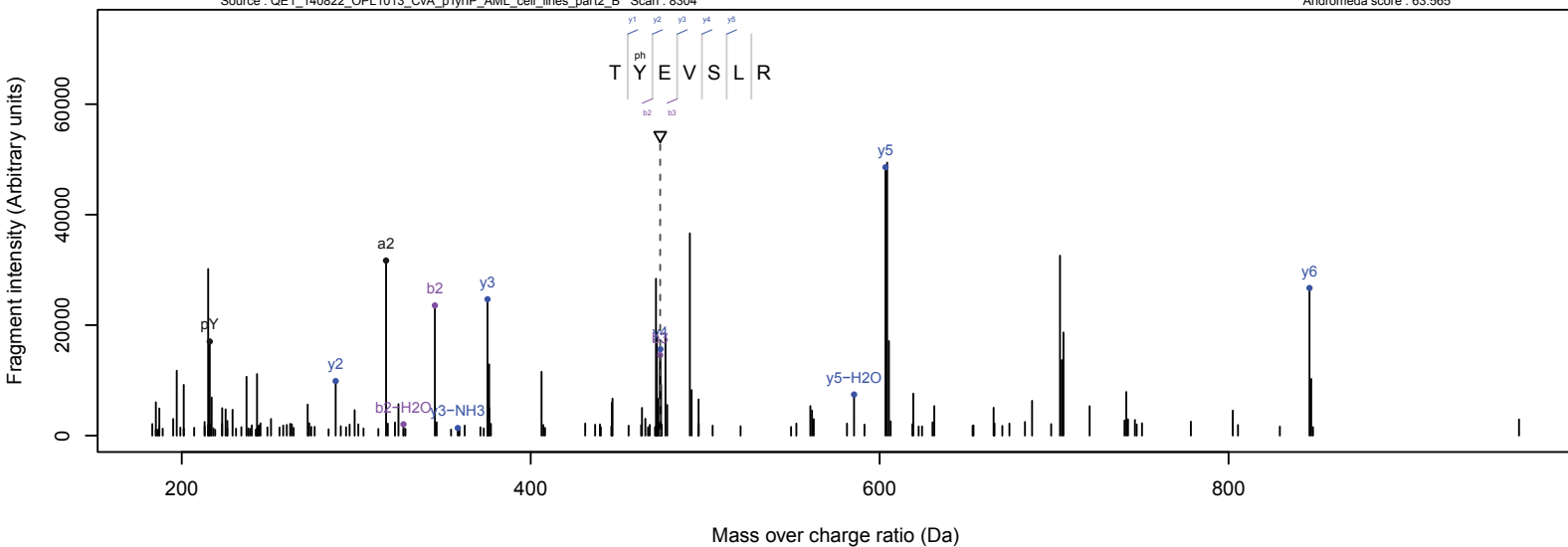
Andromeda score : 63.602



## TY(ph)EVSLR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 8304

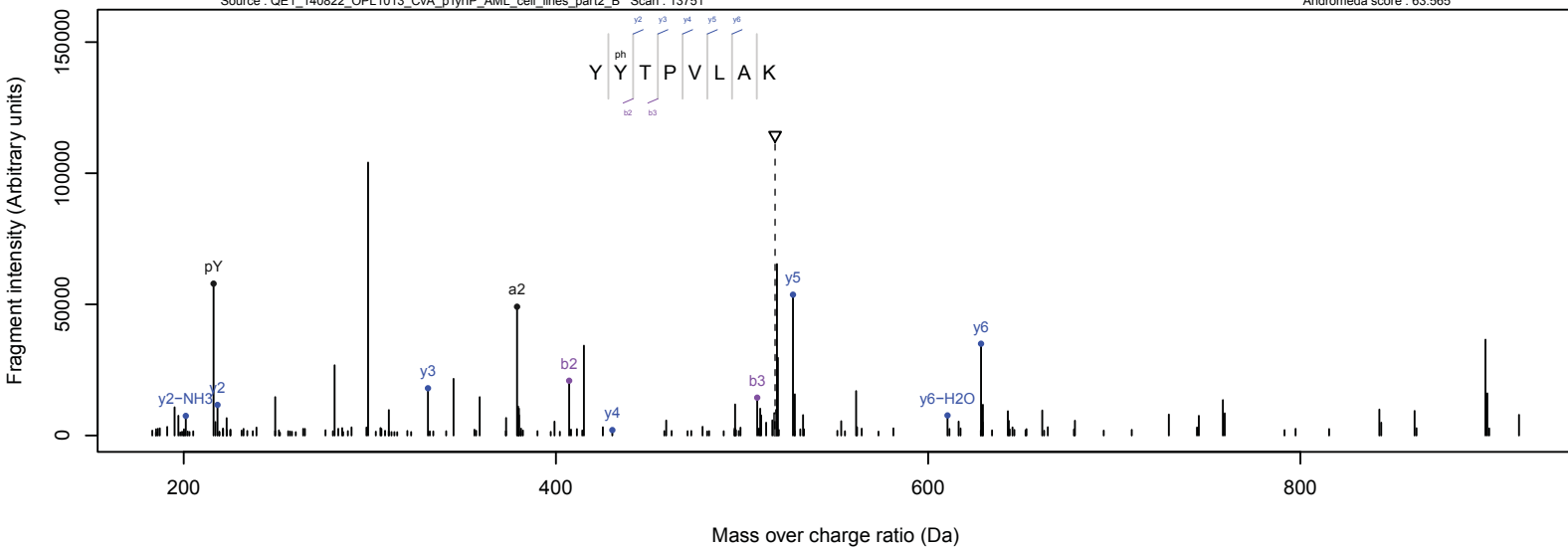
Andromeda score : 63.565



## YY(ph)TPVLAK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 13751

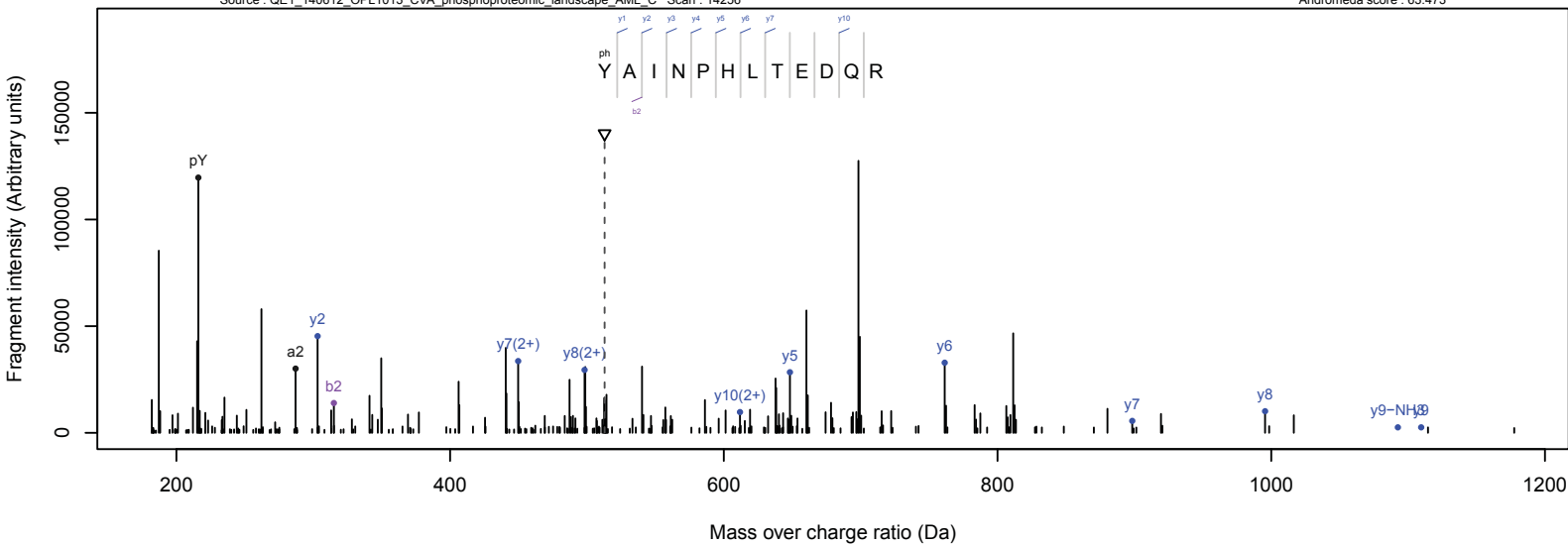
Andromeda score : 63.565



## Y(ph)AINPHLTEDQR-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 14256

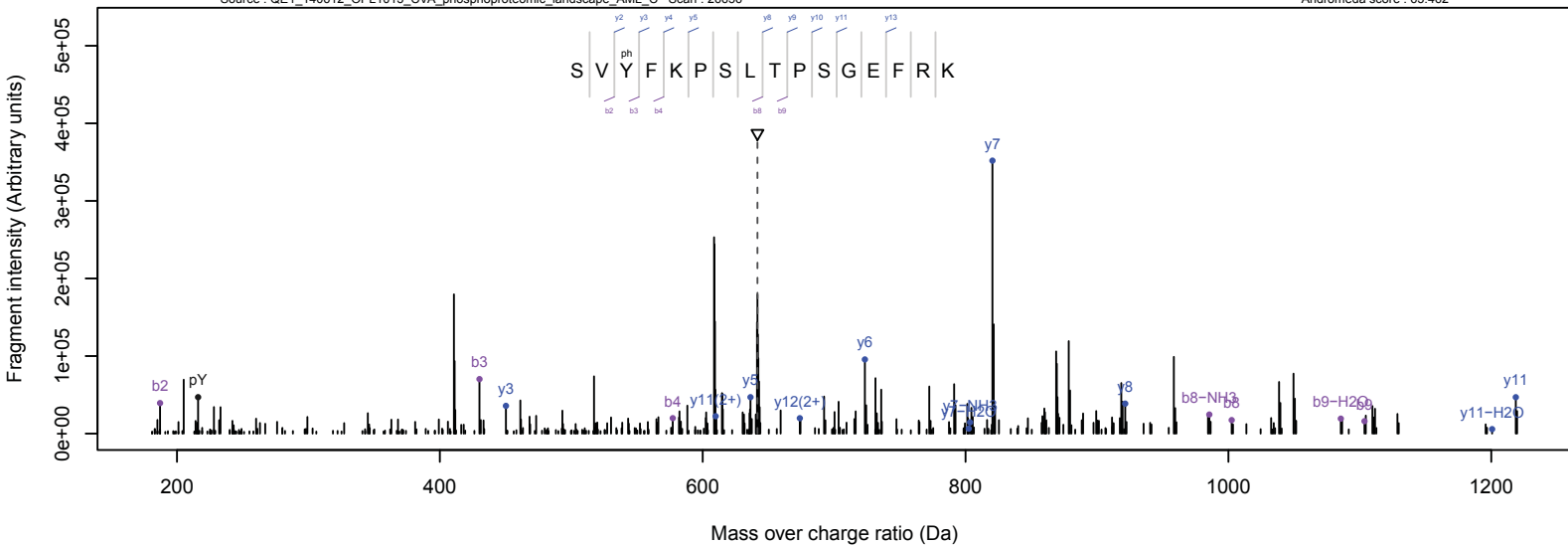
Andromeda score : 63.473



## SVY(ph)FKPSLTSPSGEFRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26656

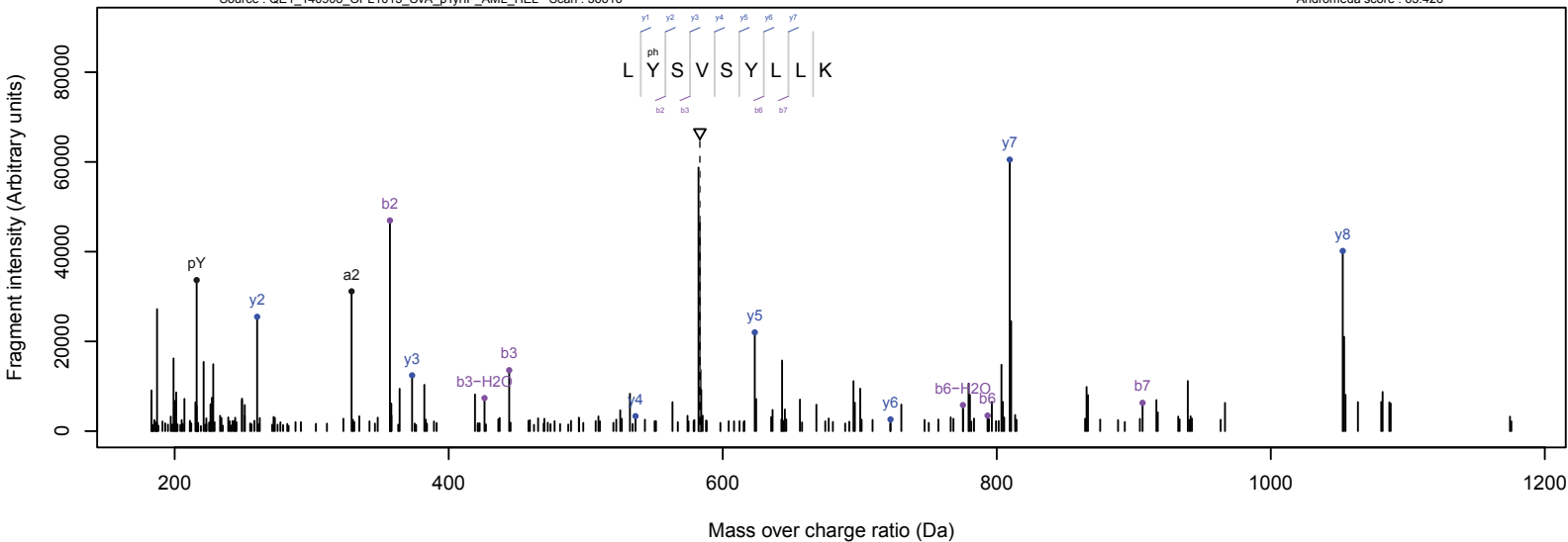
Andromeda score : 63.462



## LY(ph)SVSYLLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 36616

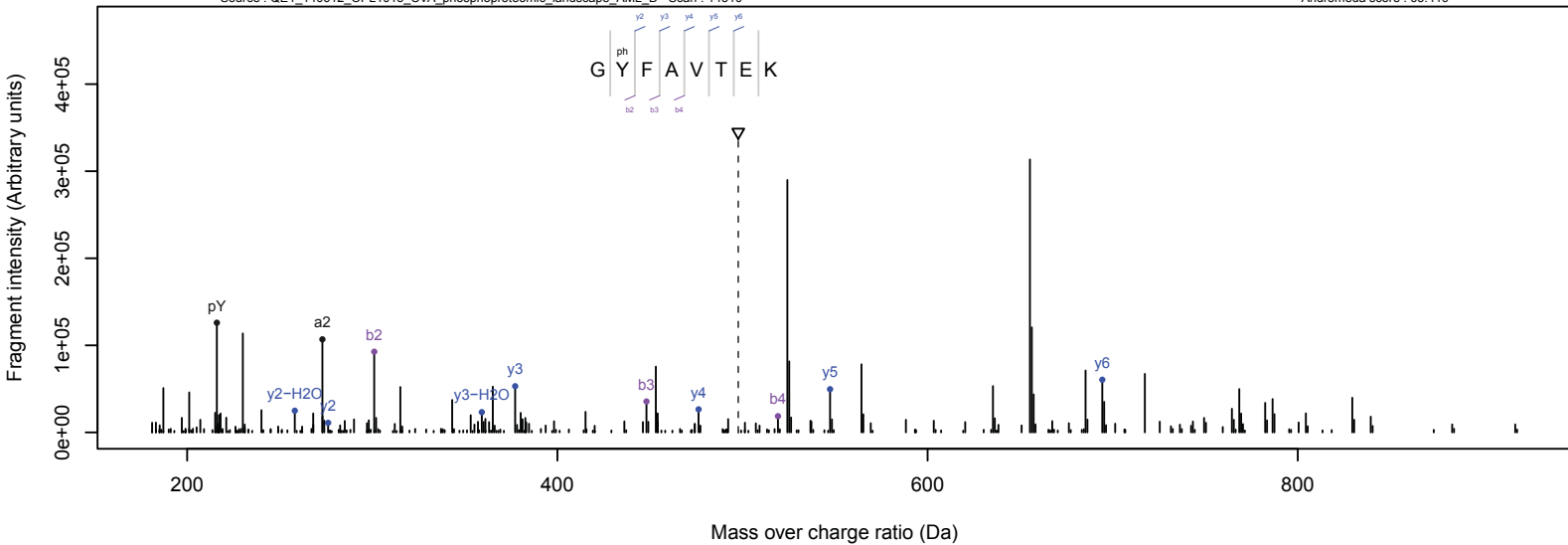
Andromeda score : 63.426



## GY(ph)FAVTEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14310

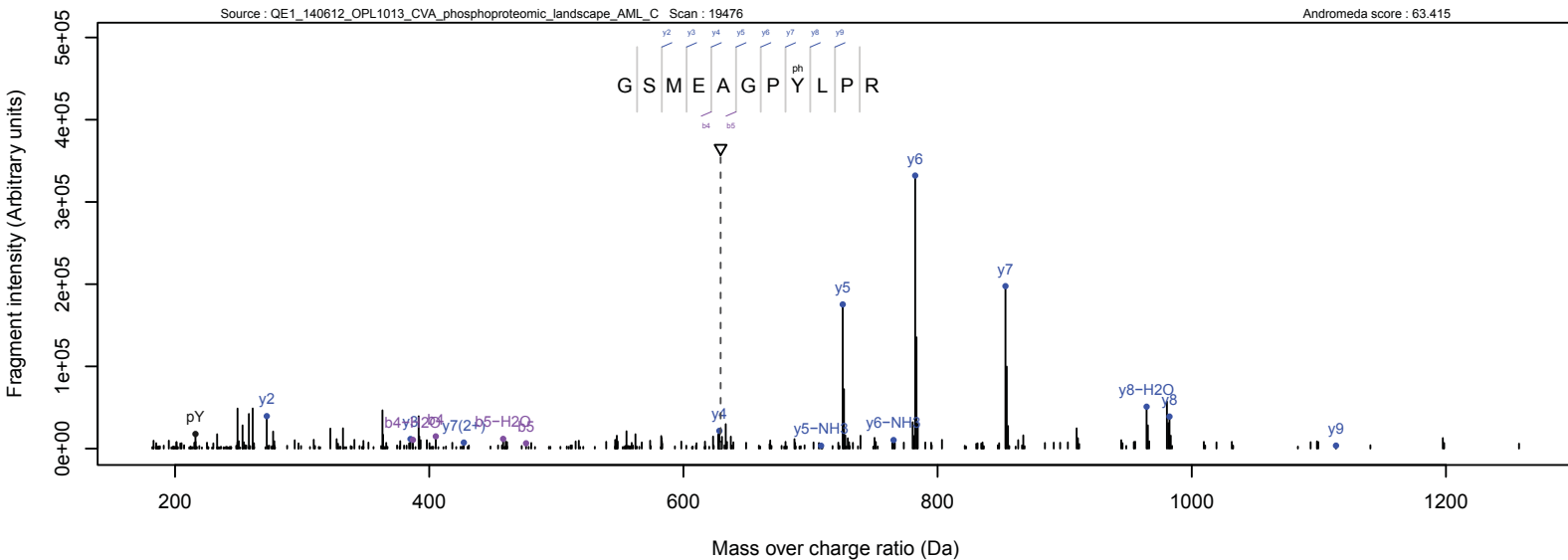
Andromeda score : 63.419



**GSMEAGPY(ph)LPR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19476

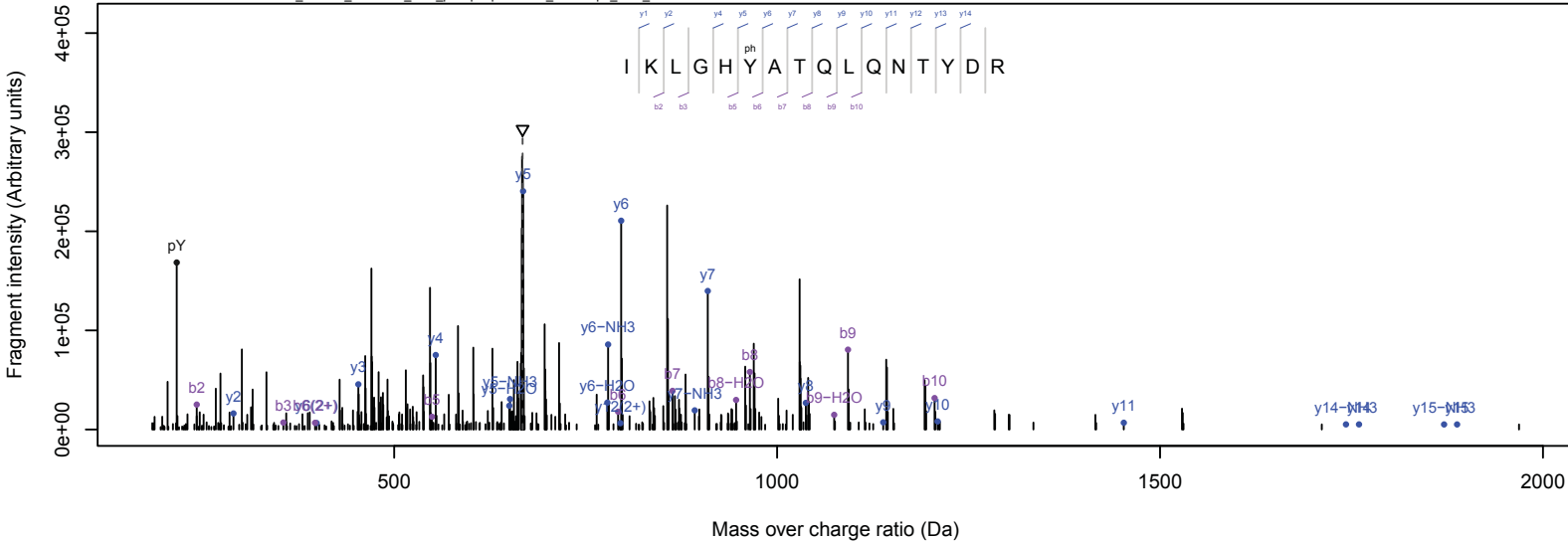
Andromeda score: 63.415



**IKLGHY(ph)ATQLQNTYDR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 23093

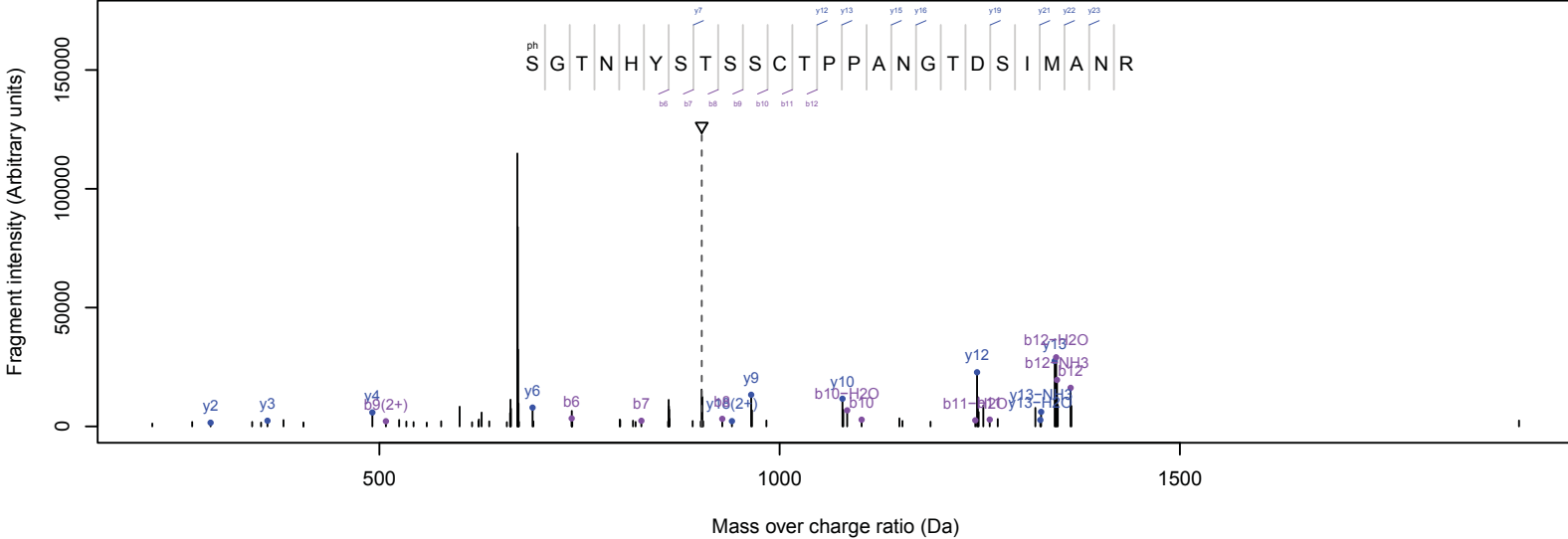
Andromeda score: 63.385



**S(ph)GTNHYSTSSCTPPANGTDSIMANR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15033

Andromeda score: 63.376

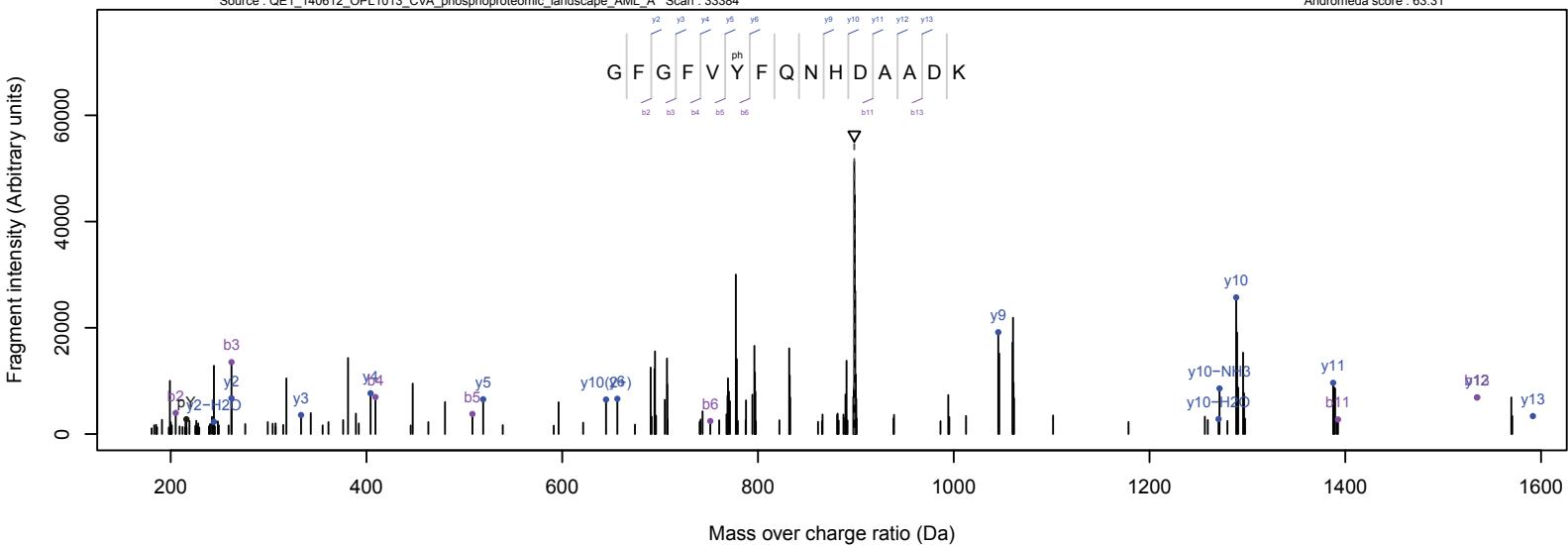




## GFGFVY(ph)FQNHDAADK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33384

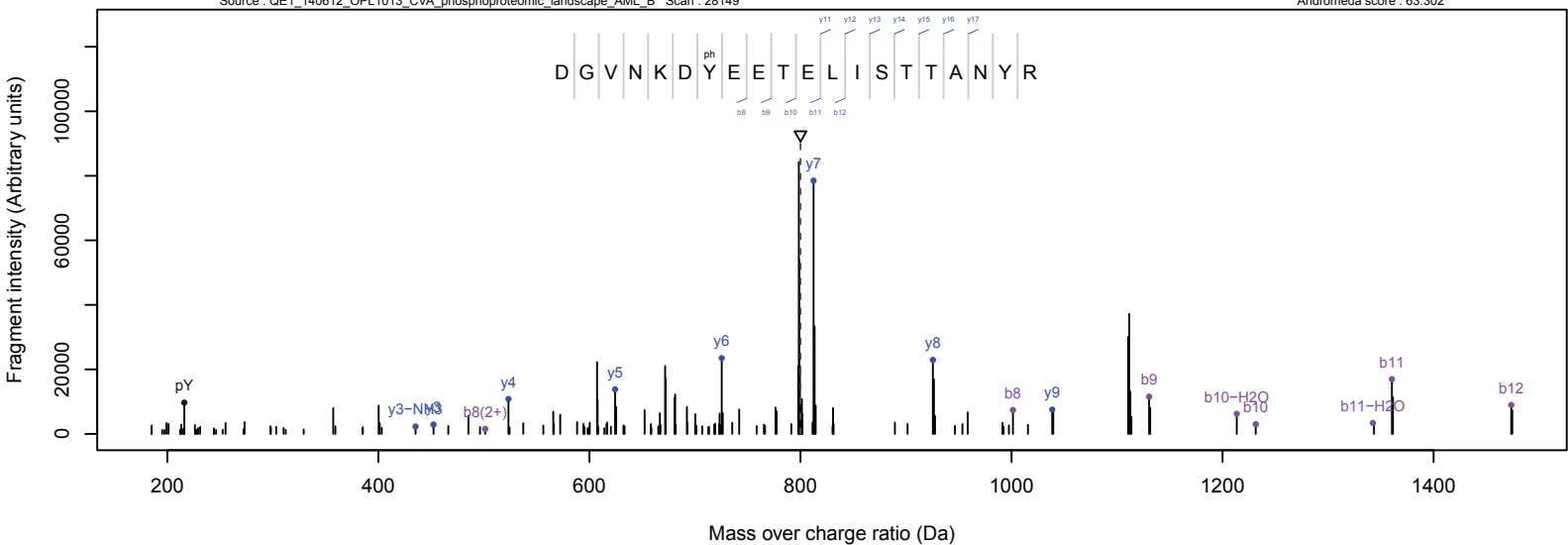
Andromeda score : 63.31



## DGVNKDY(ph)EETELISTANYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28149

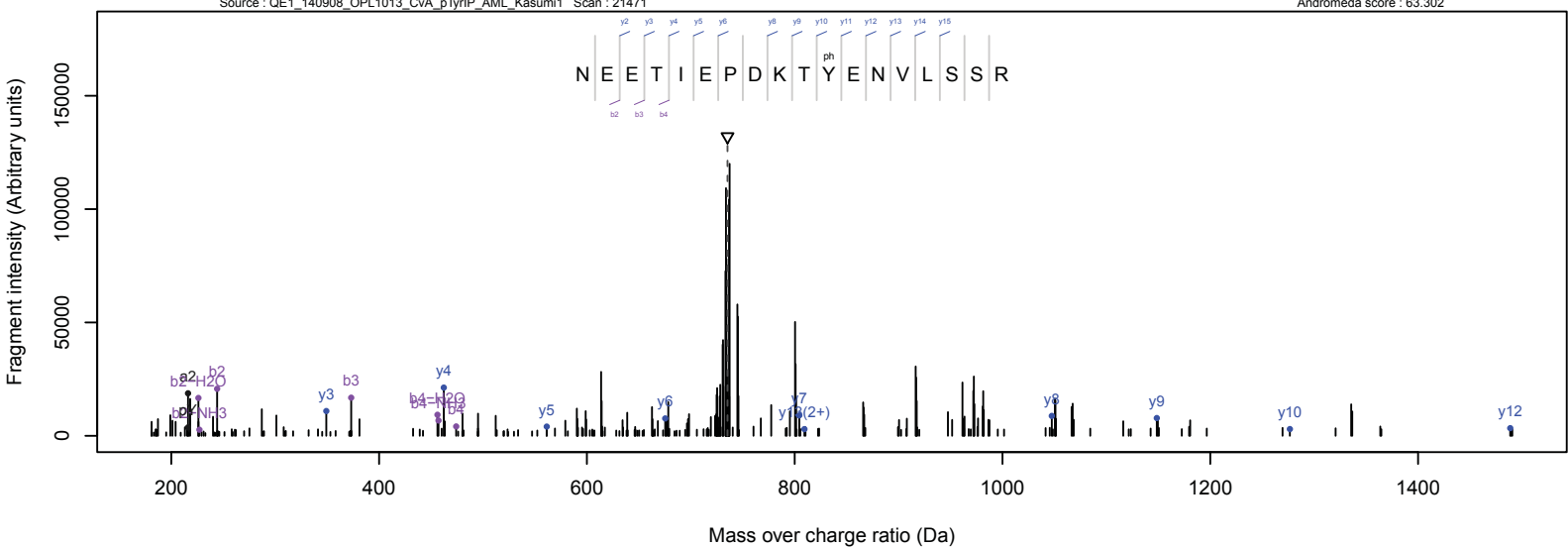
Andromeda score : 63.302



## NEETIEPDKTY(ph)ENVLSSR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 21471

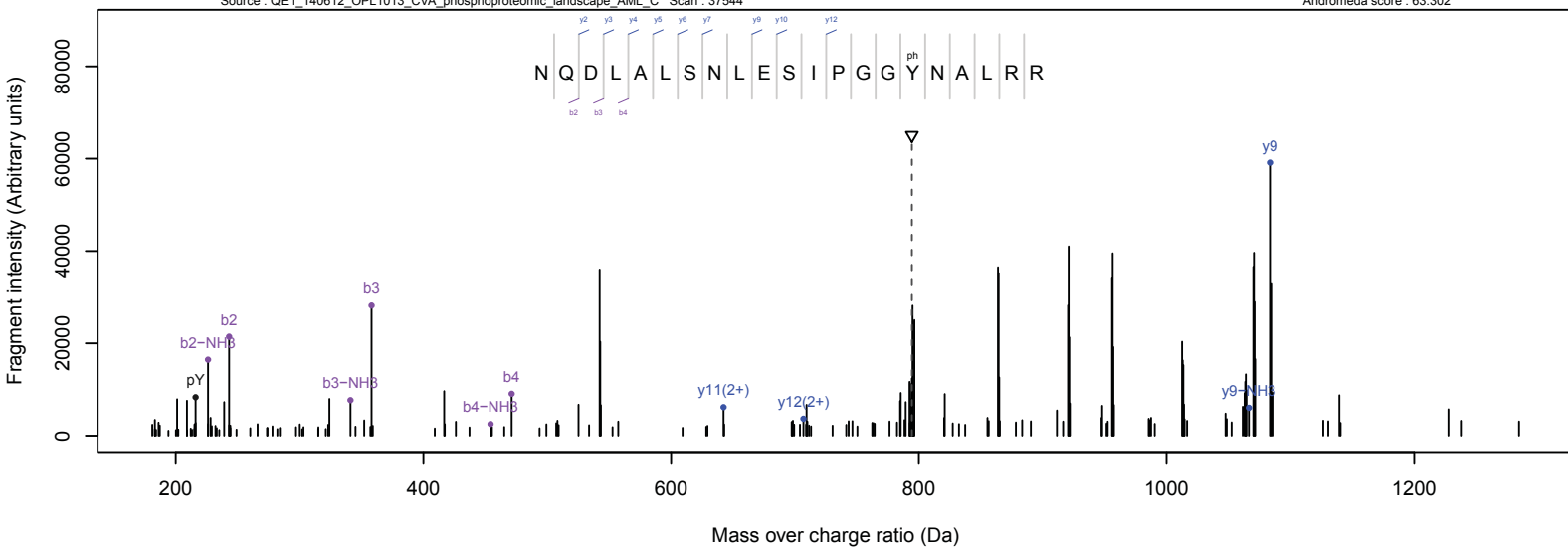
Andromeda score : 63.302



NQDLALSLESIPGGY<sup>ph</sup>NALRR-3+

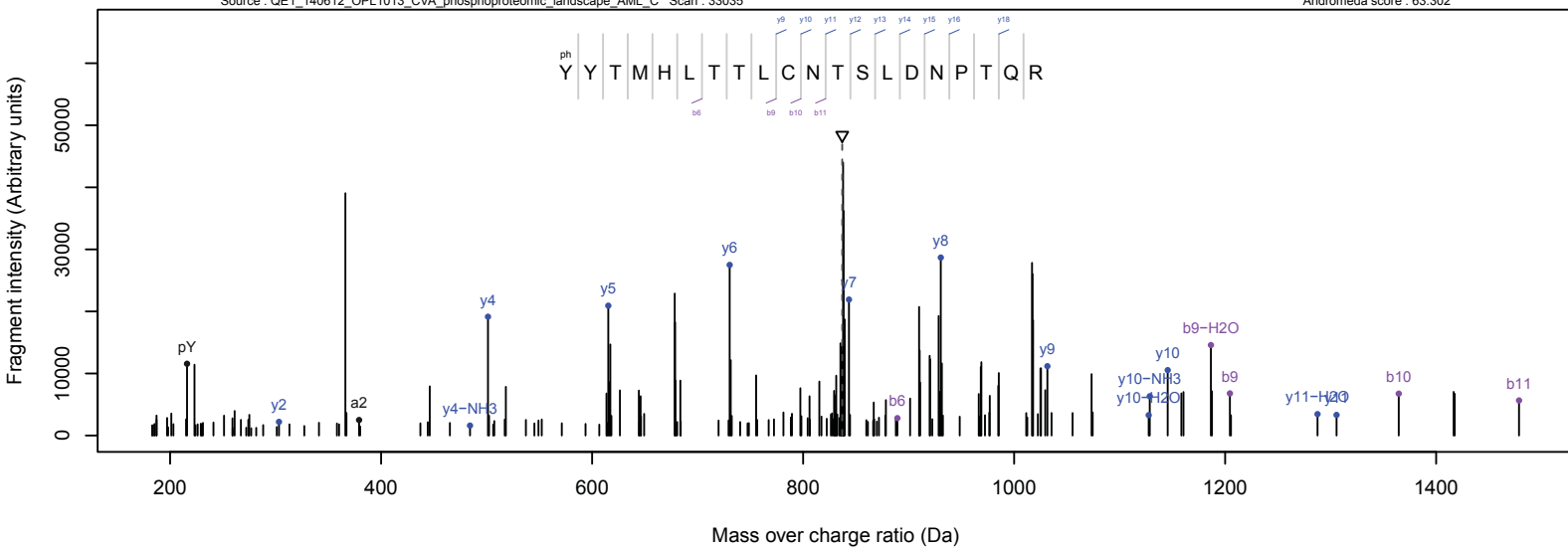
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37544

Andromeda score : 63.302

Y<sup>ph</sup>YTMHLTTLCLNTSLDNPTR-3+

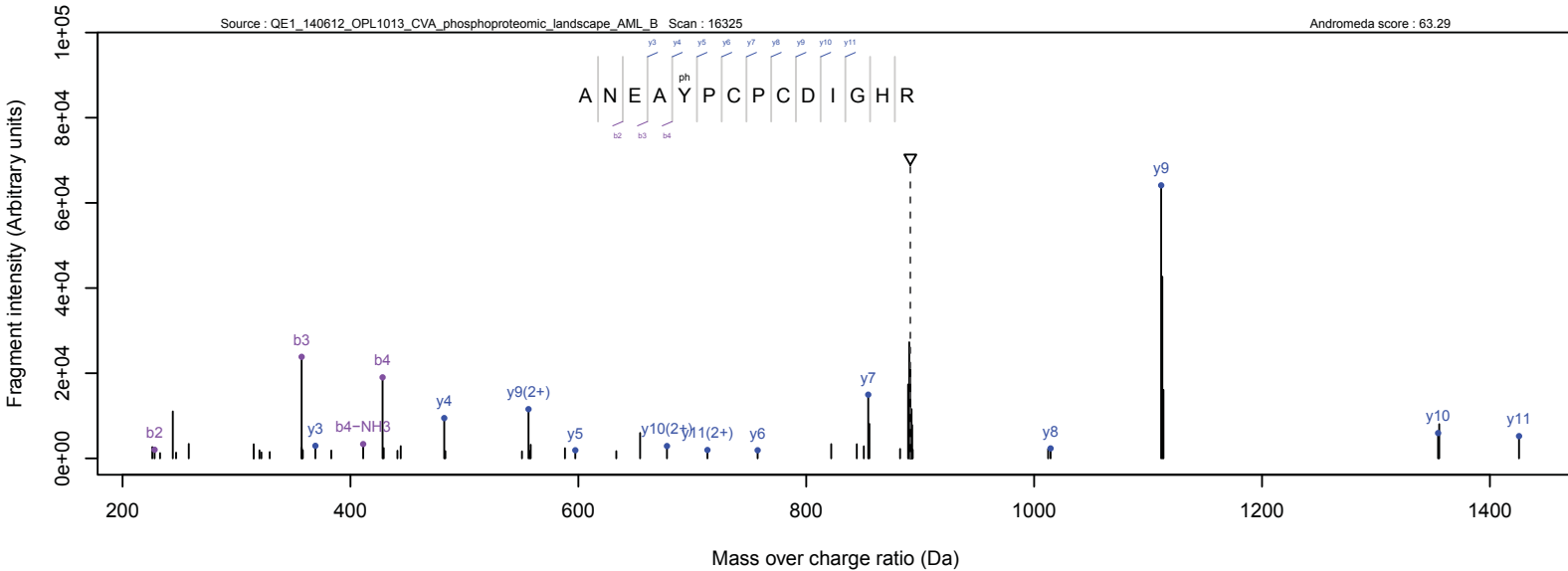
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33035

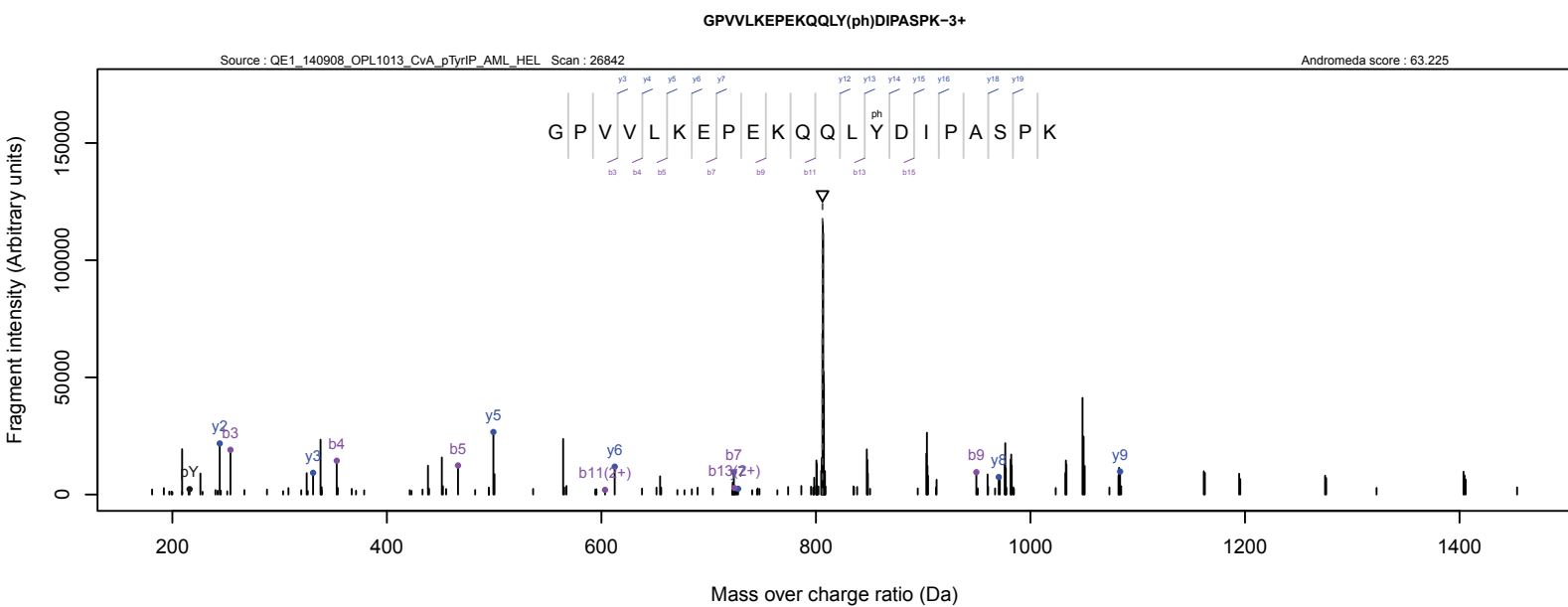
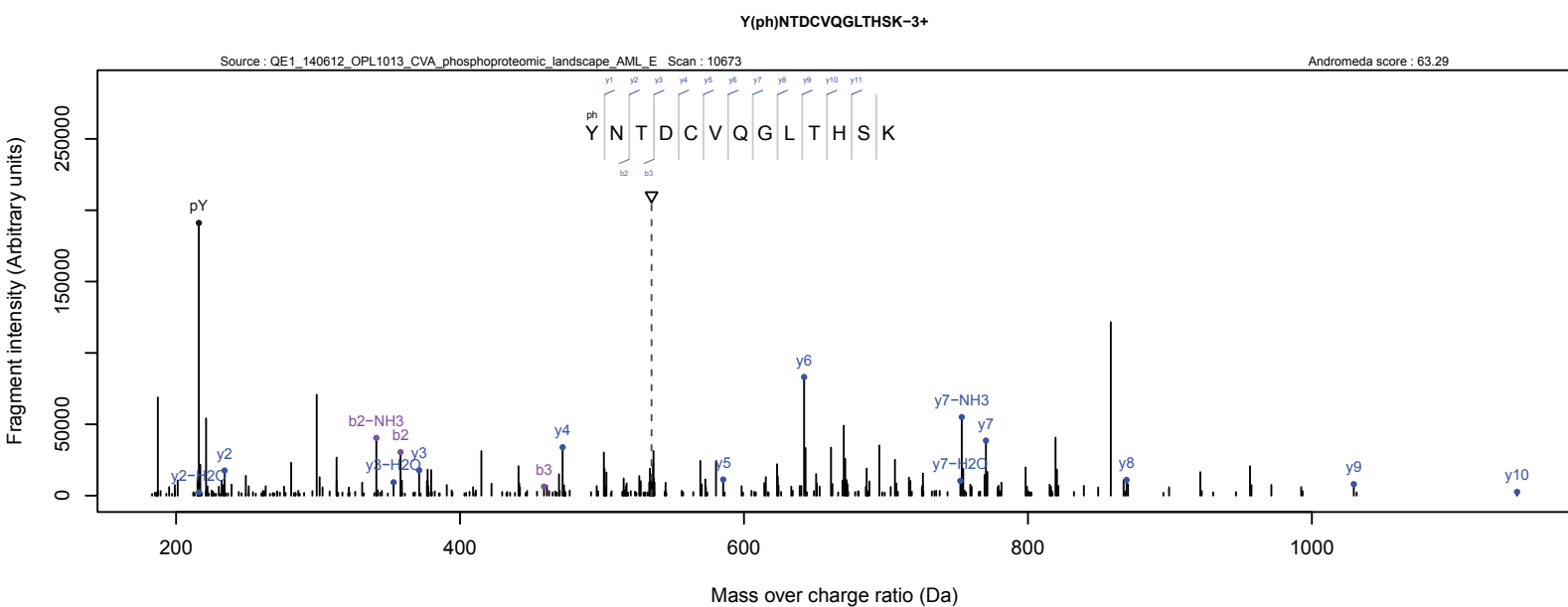
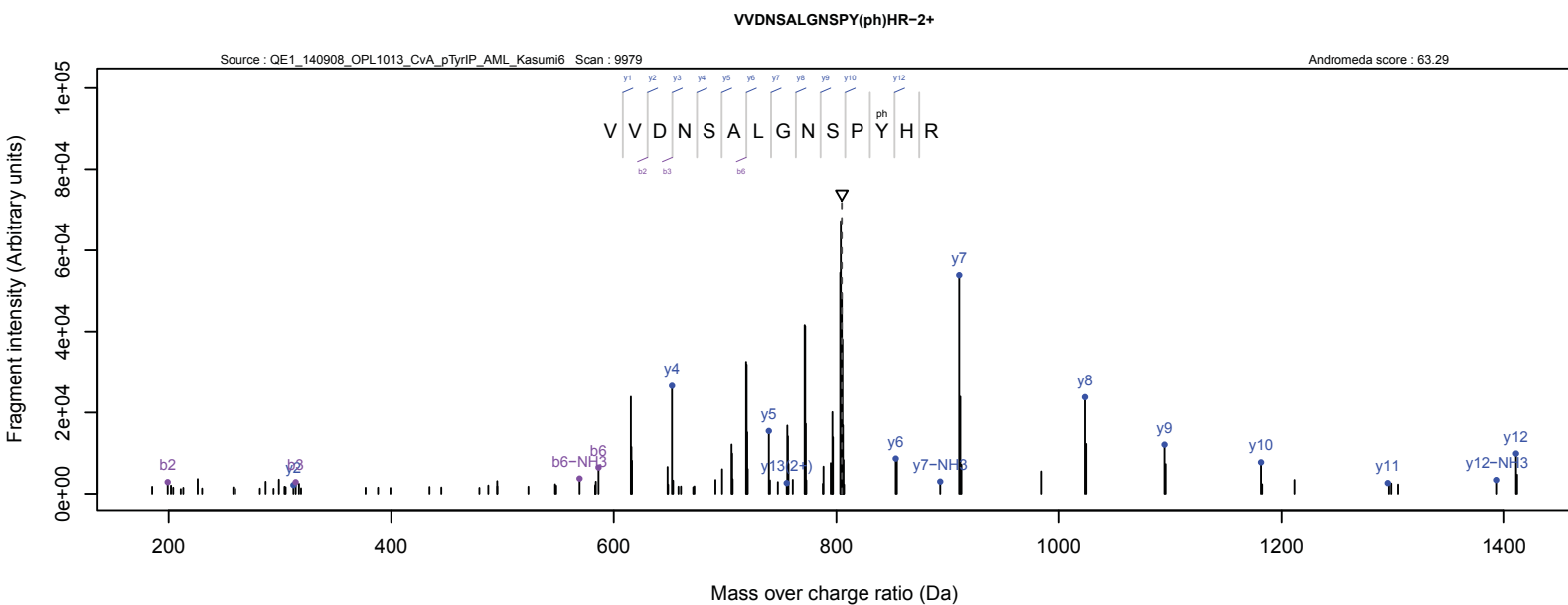
Andromeda score : 63.302

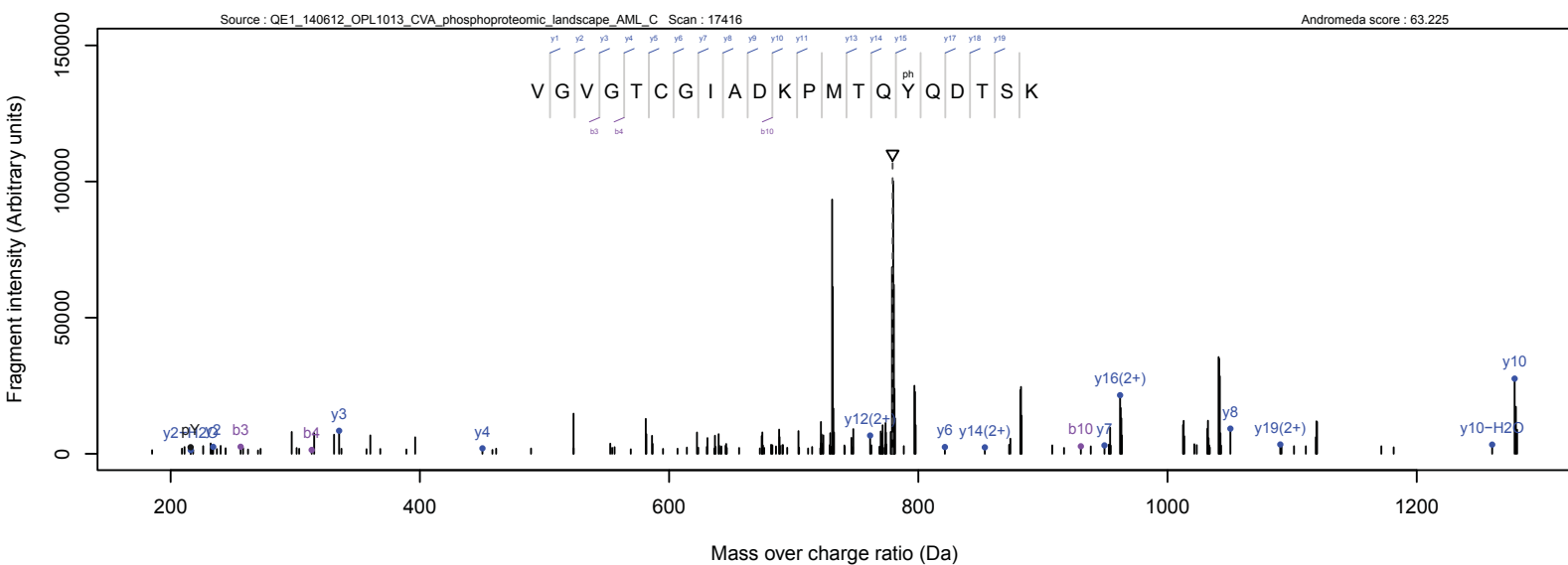
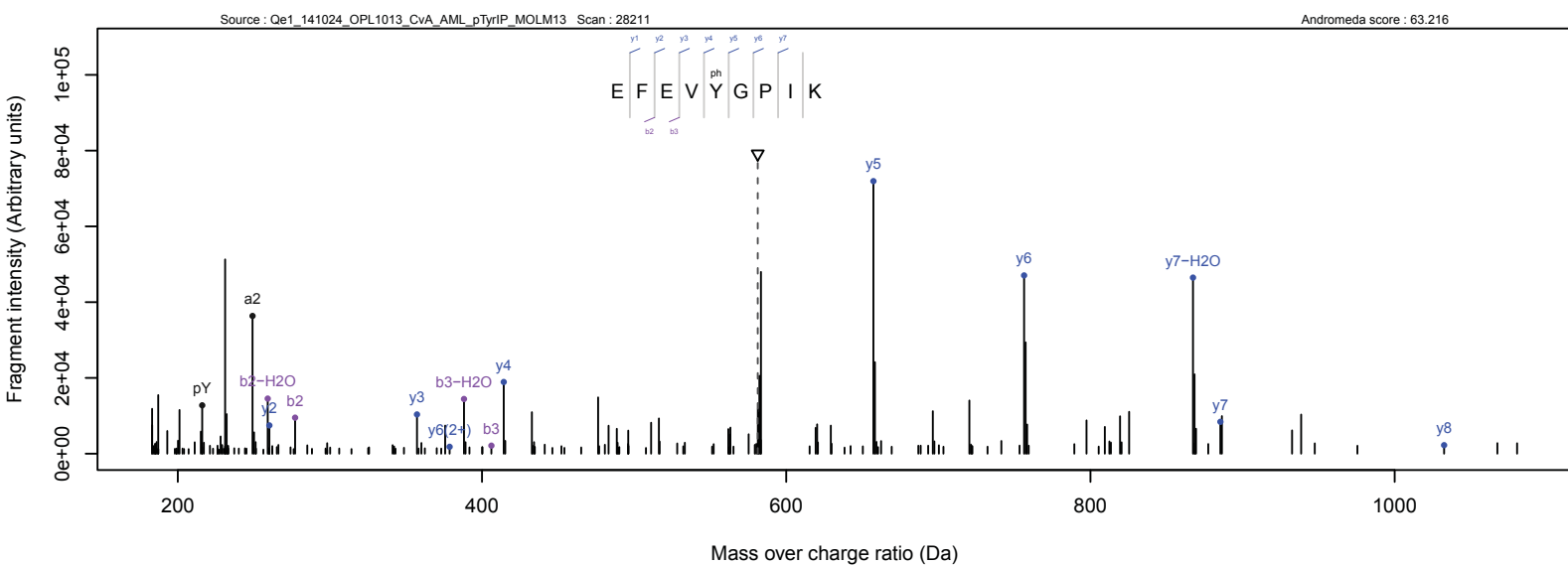
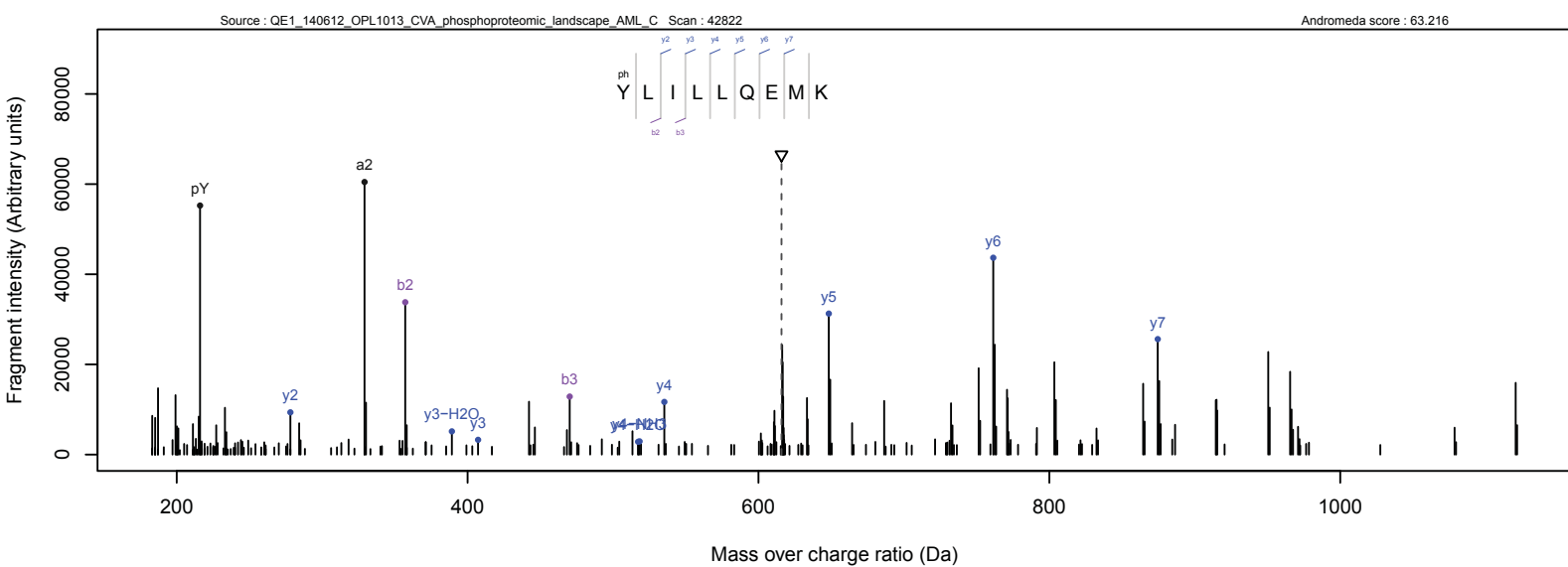
(ac)ANEAY<sup>ph</sup>PCPCDIGHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16325

Andromeda score : 63.29





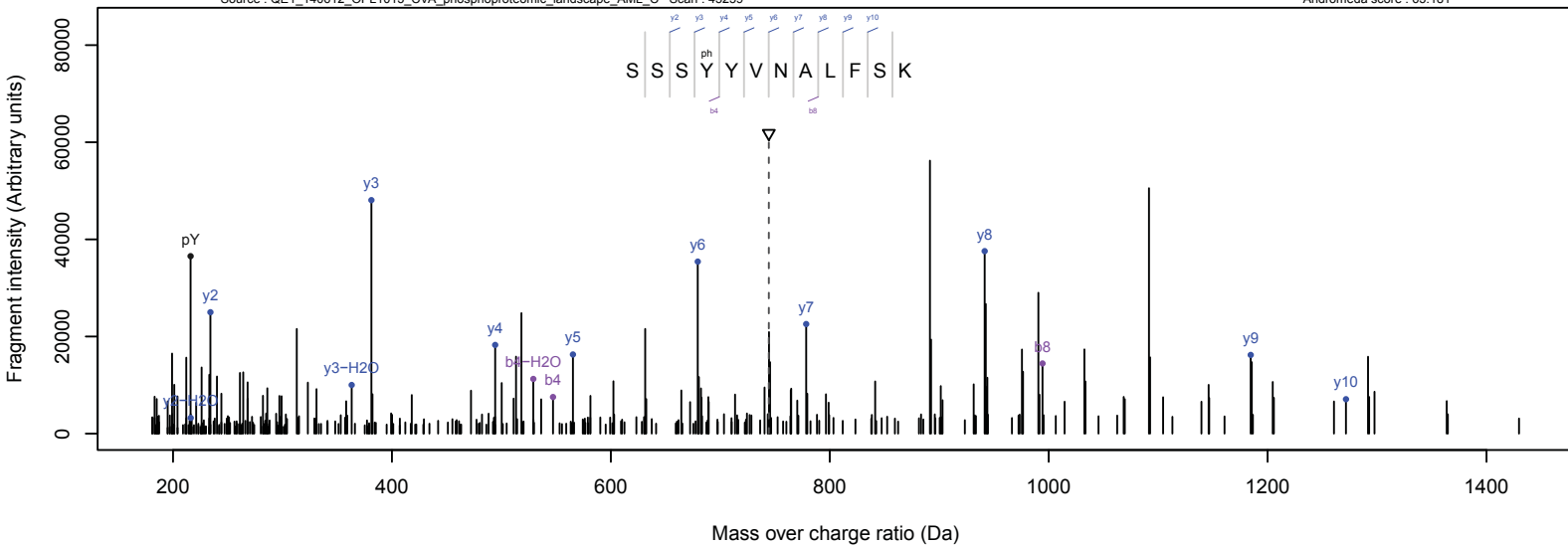
VGVGTCGIADKPM<sup>ph</sup>TQYQDTSK-3+EFEV<sup>ph</sup>GPIK-2+Y<sup>ph</sup>LILLQEMK-2+



## (ac)SSSY(ph)YVNALFSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45259

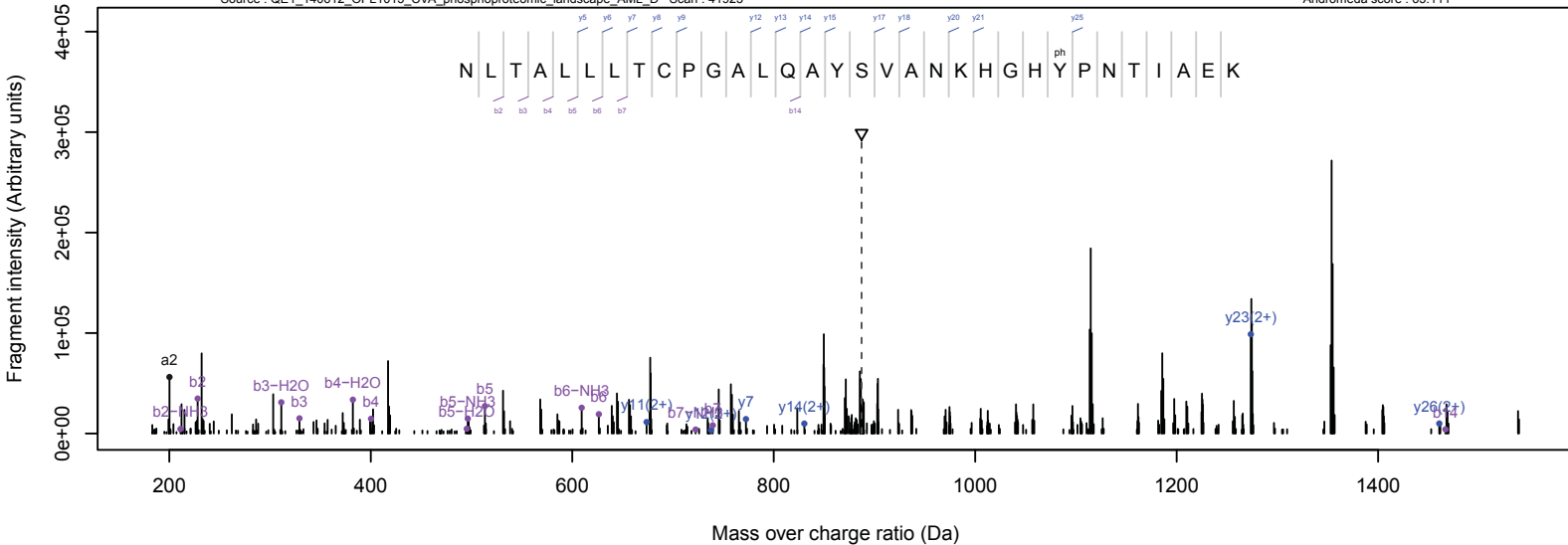
Andromeda score : 63.181



## NLTALLTCPGALQAYSVANKHGHY(ph)PNTIAEK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 41923

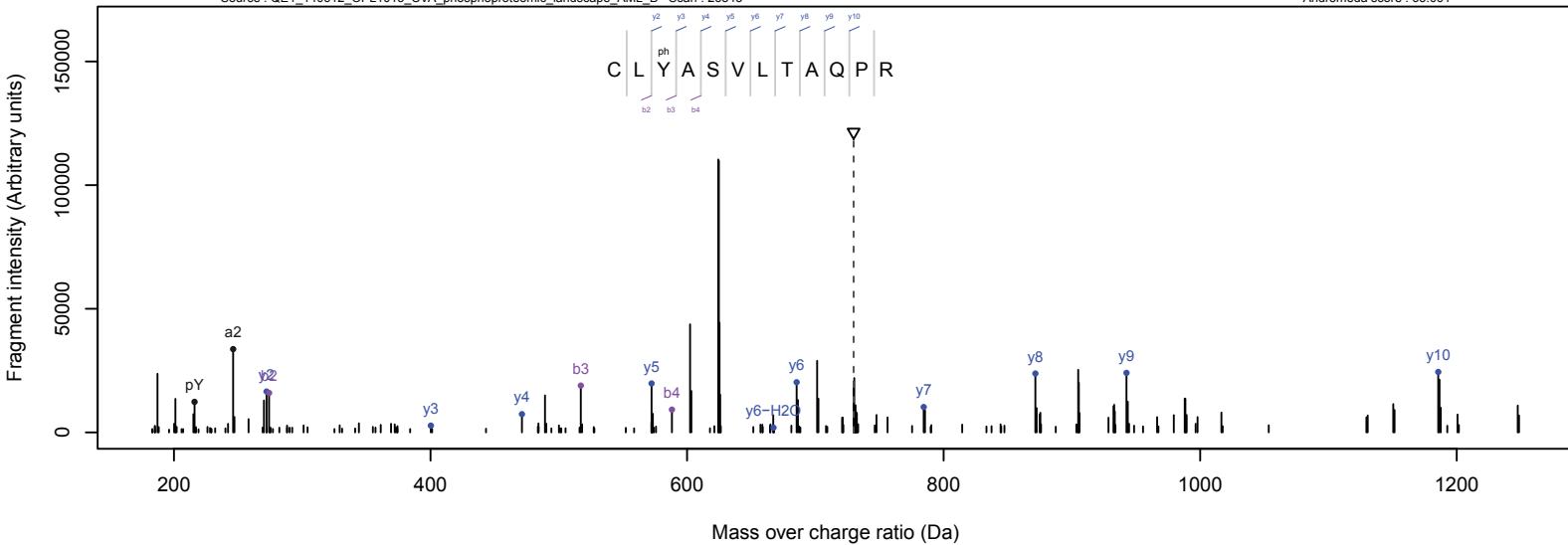
Andromeda score : 63.111



## CLY(ph)ASVLTAQPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26343

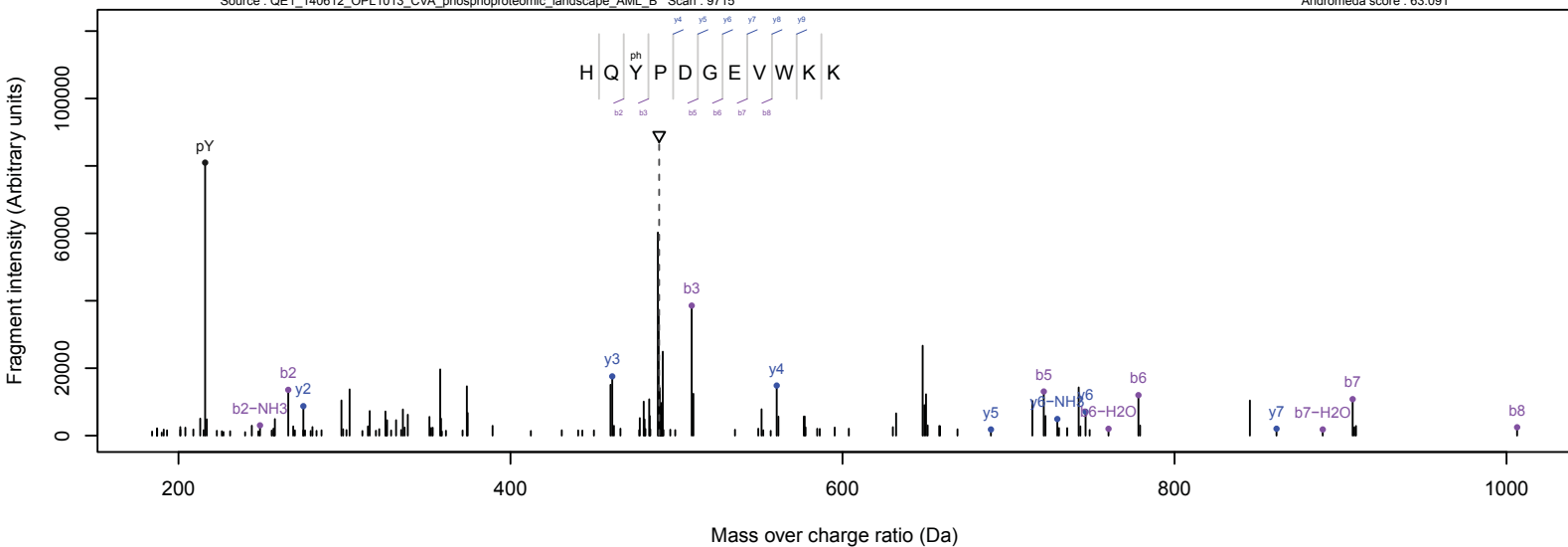
Andromeda score : 63.091



HQY(ph)PDGEVWKK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 9715

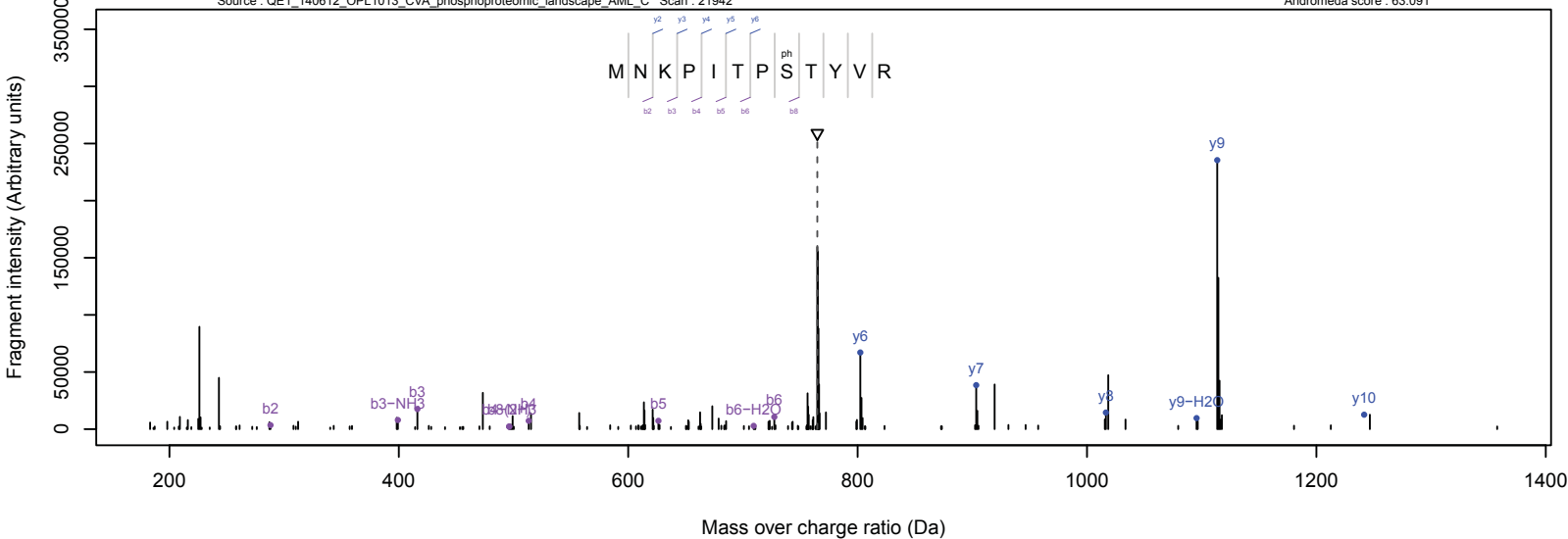
Andromeda score: 63.091



(ac)MNKPITPS(ph)TYVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21942

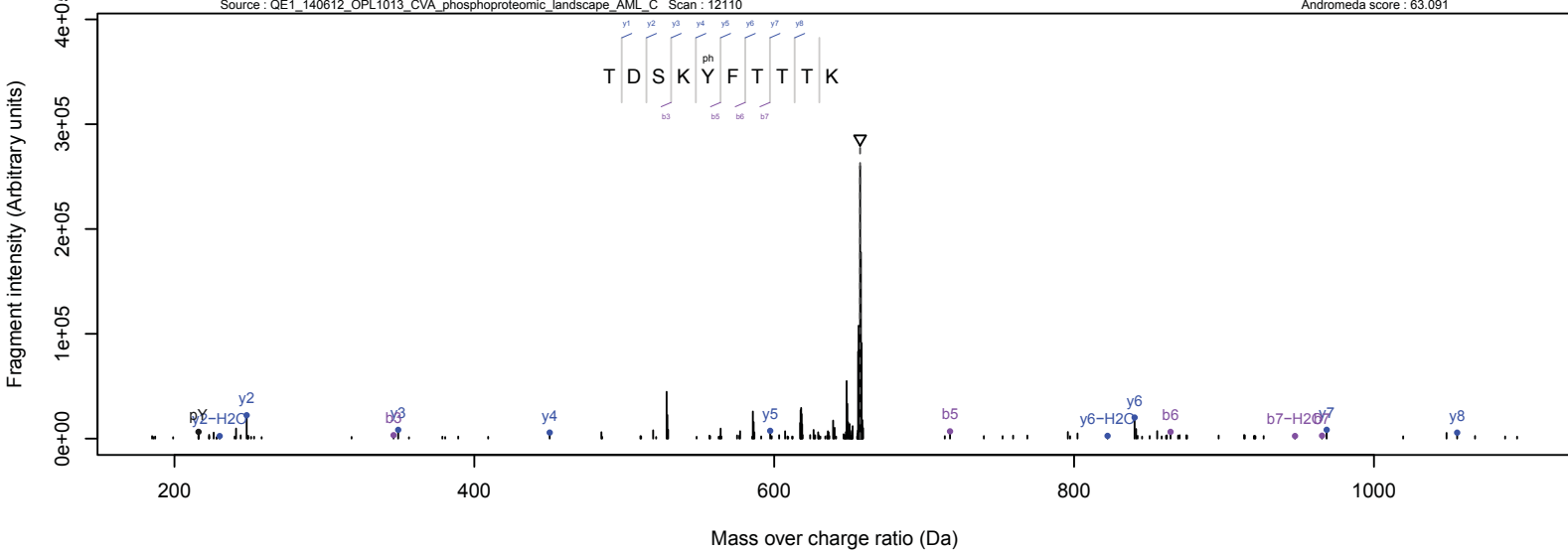
Andromeda score: 63.091

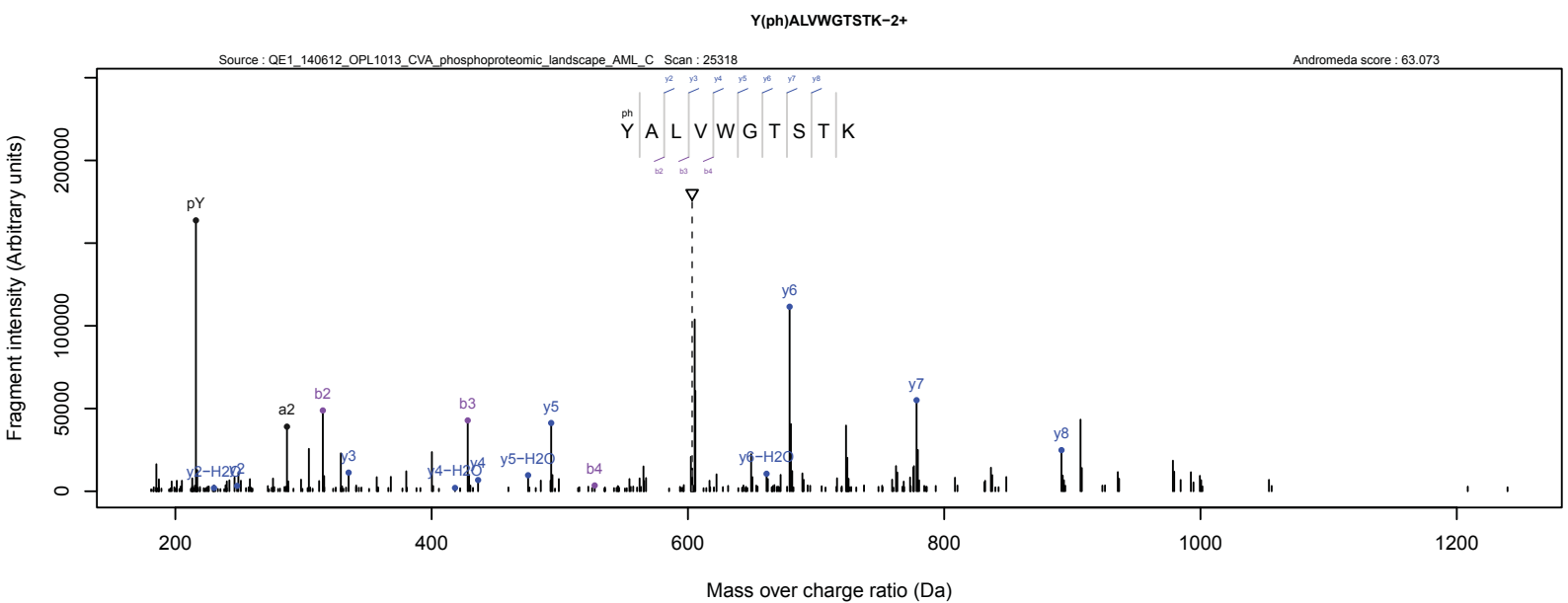
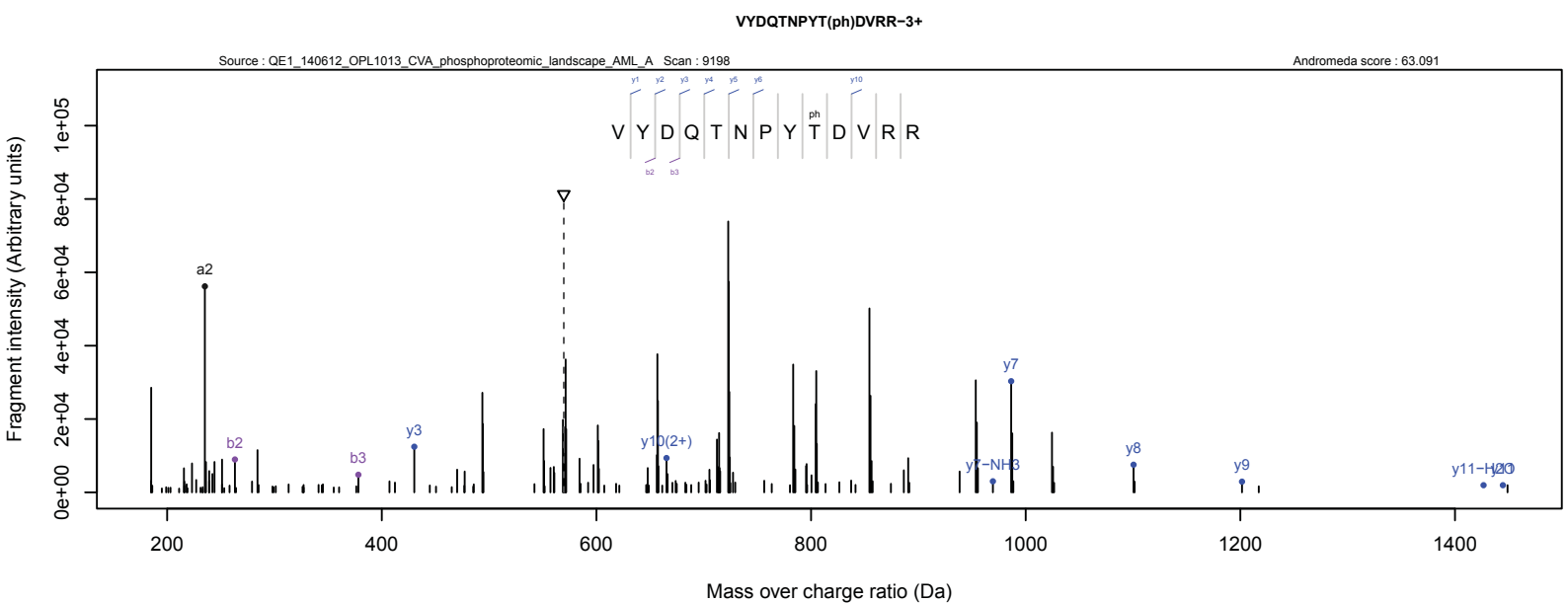
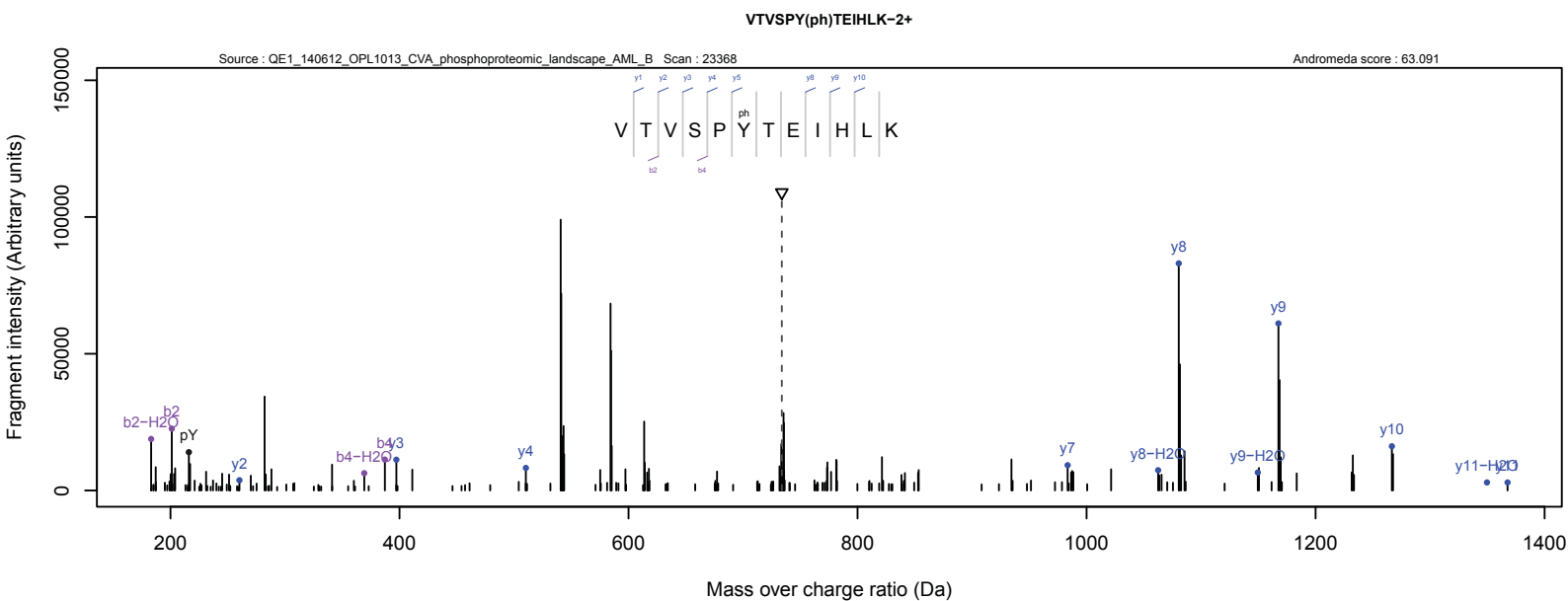


(ac)TDSKY(ph)FTTTK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12110

Andromeda score: 63.091



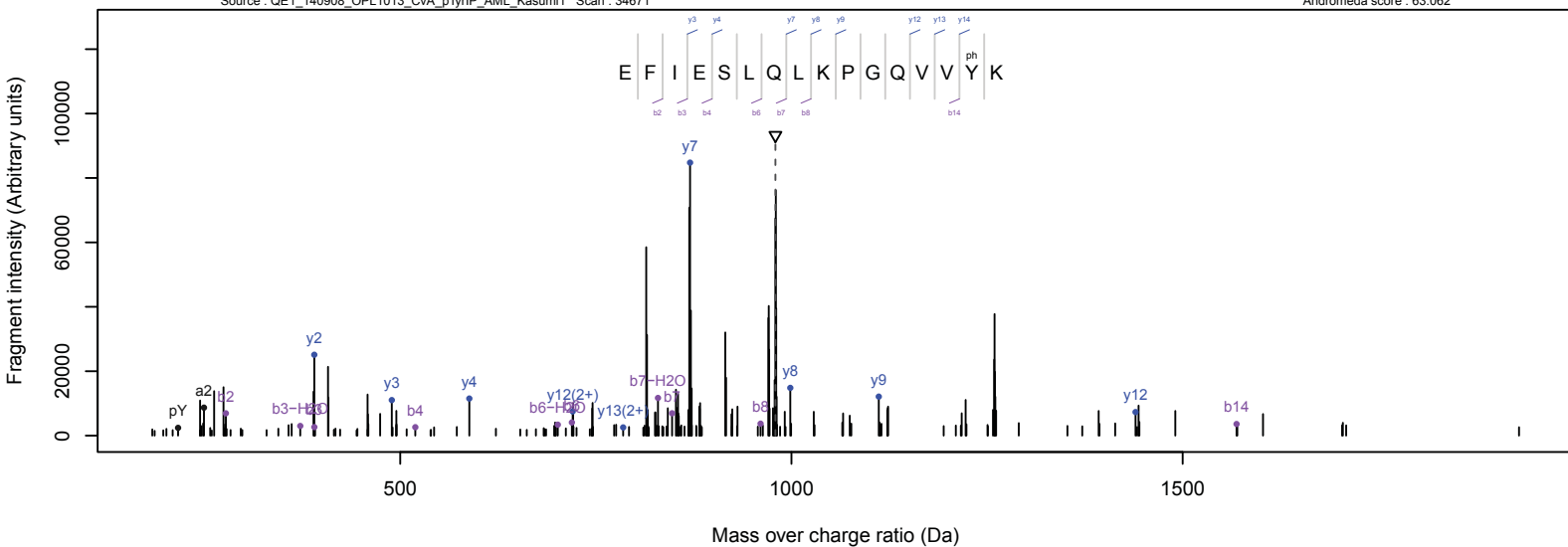




## EFIESLQKPGQVVY(ph)K-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 34671

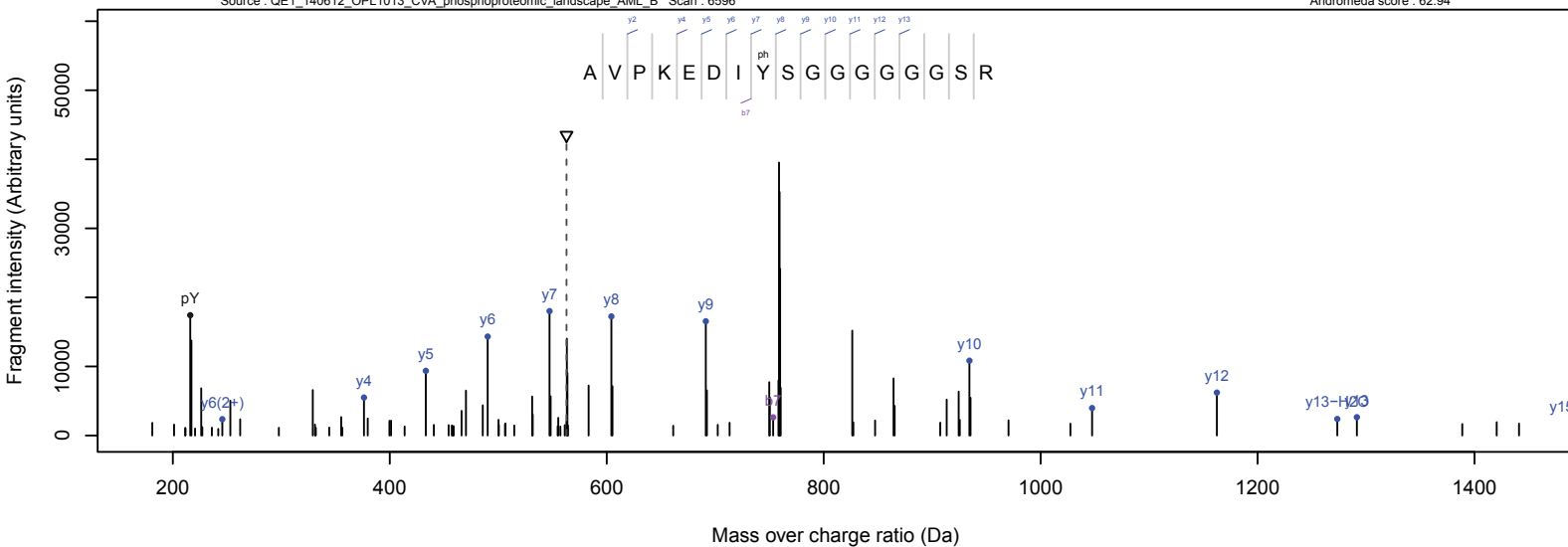
Andromeda score : 63.062



## AVPKEDIYSGGGGGSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 6596

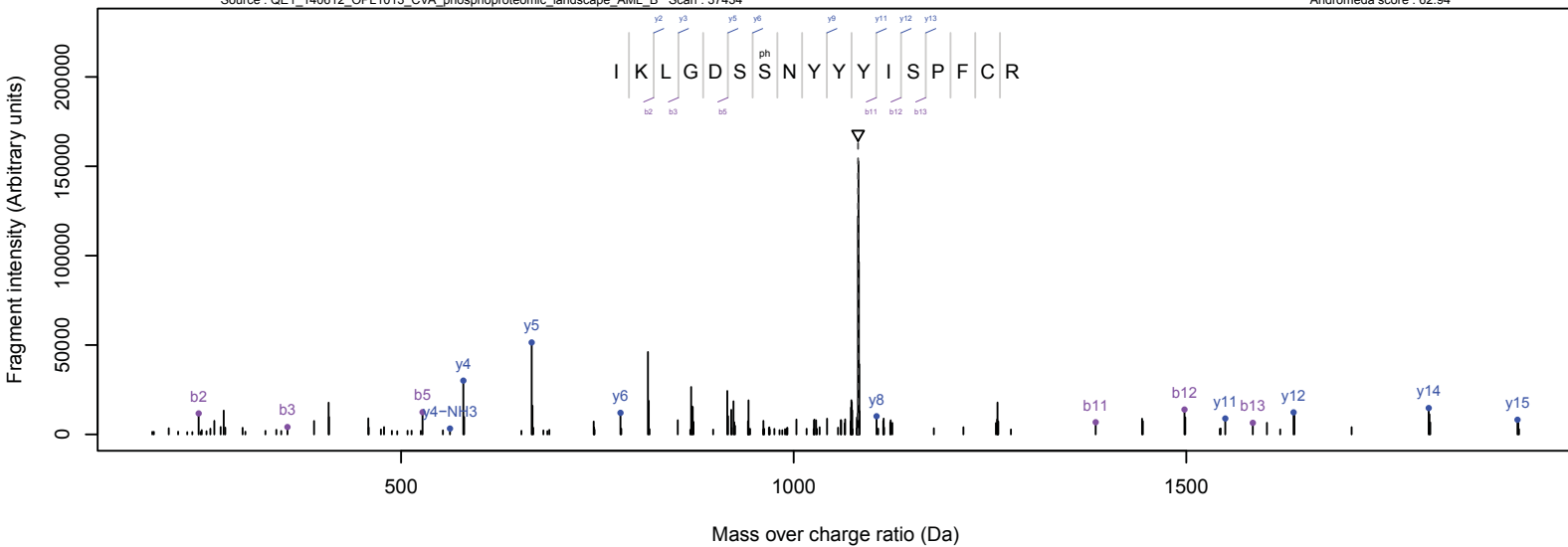
Andromeda score : 62.94

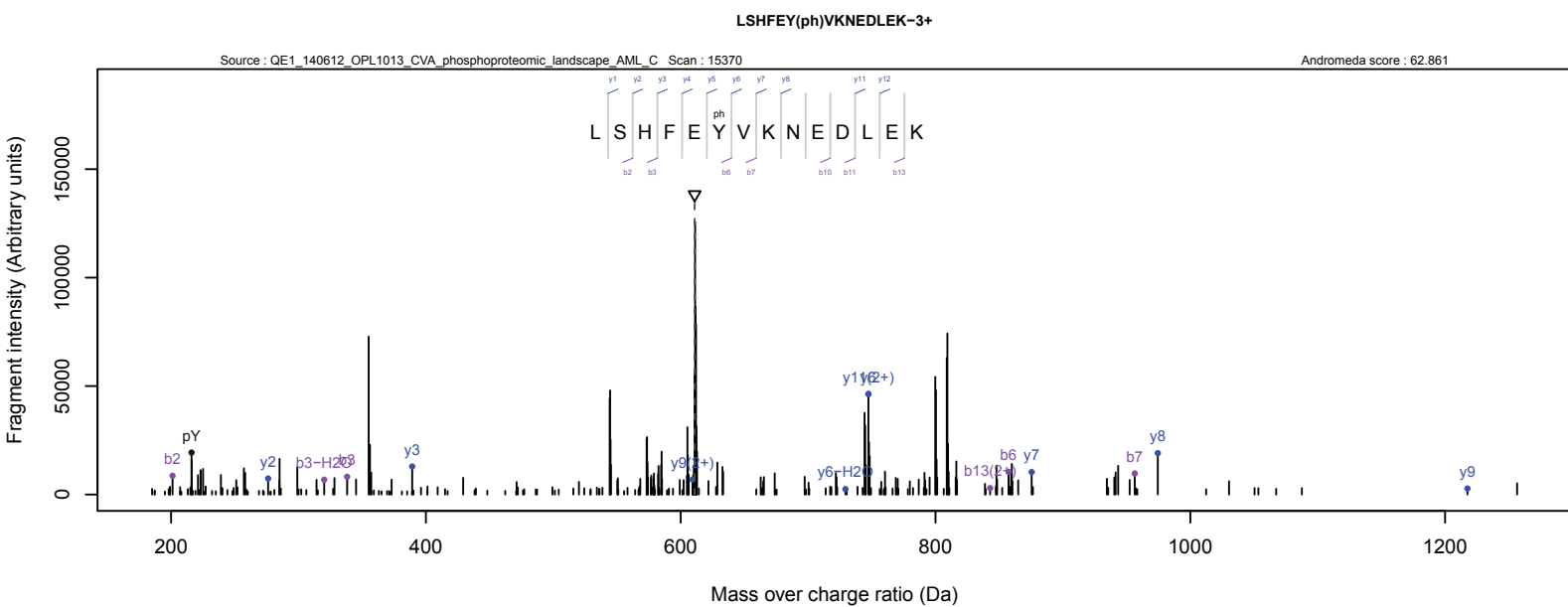
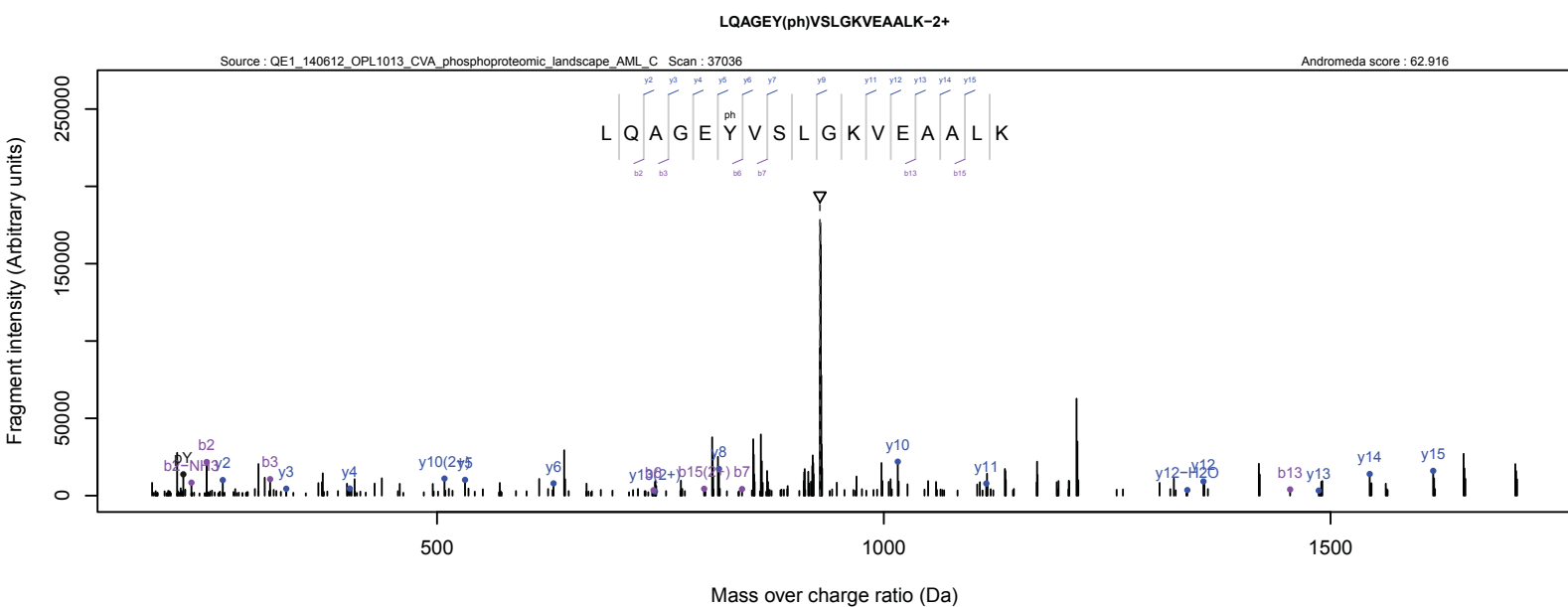
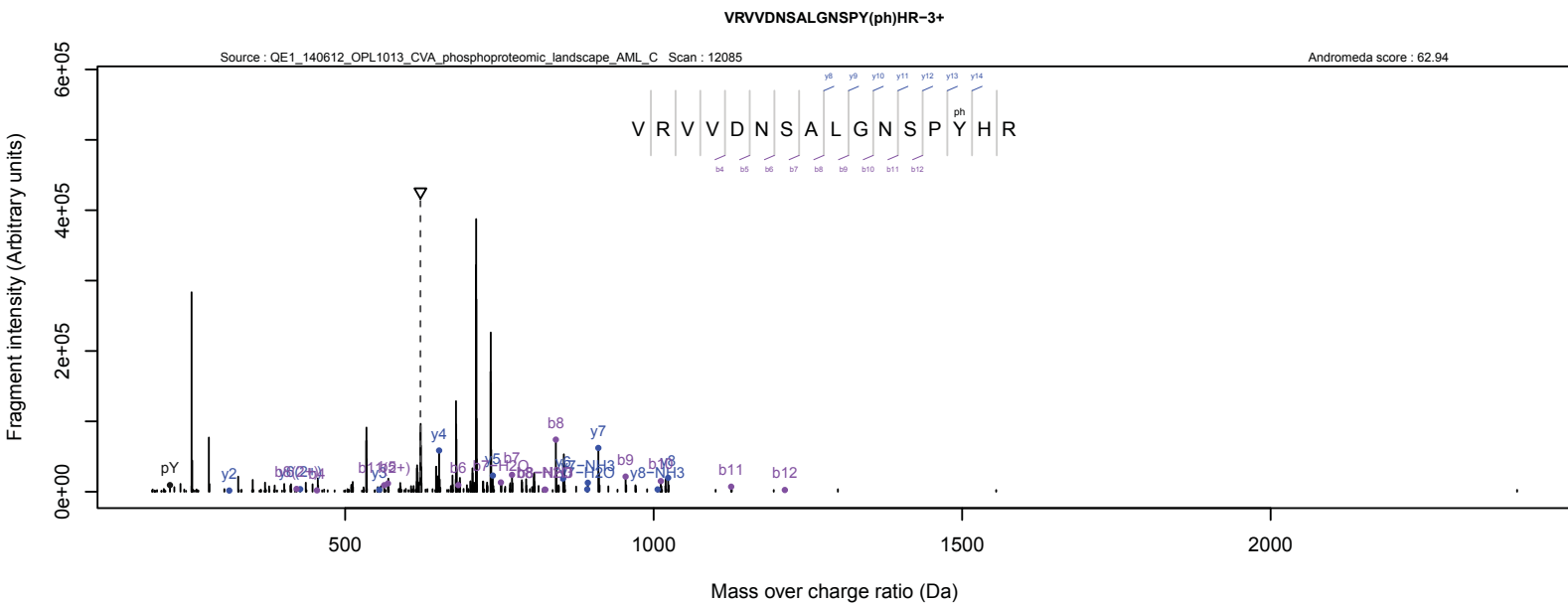


## IKLGDSS(ph)NYIYISPFGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37434

Andromeda score : 62.94

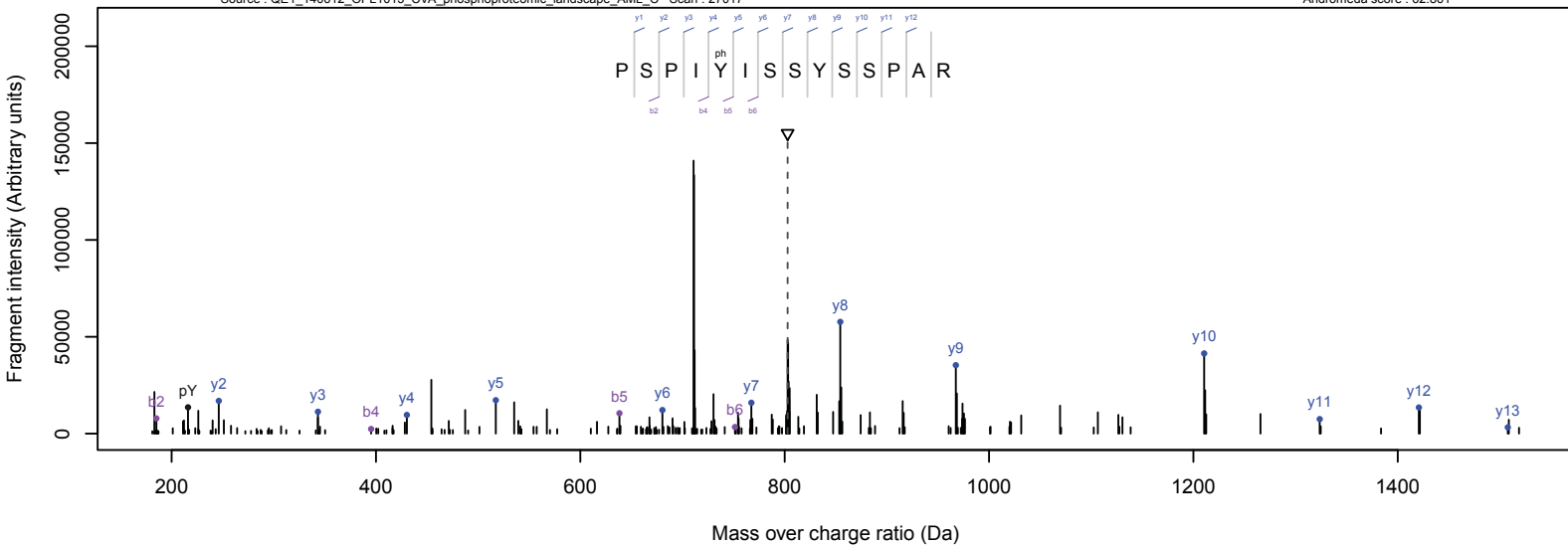




PSPIY(ph)ISSYSSPAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27017

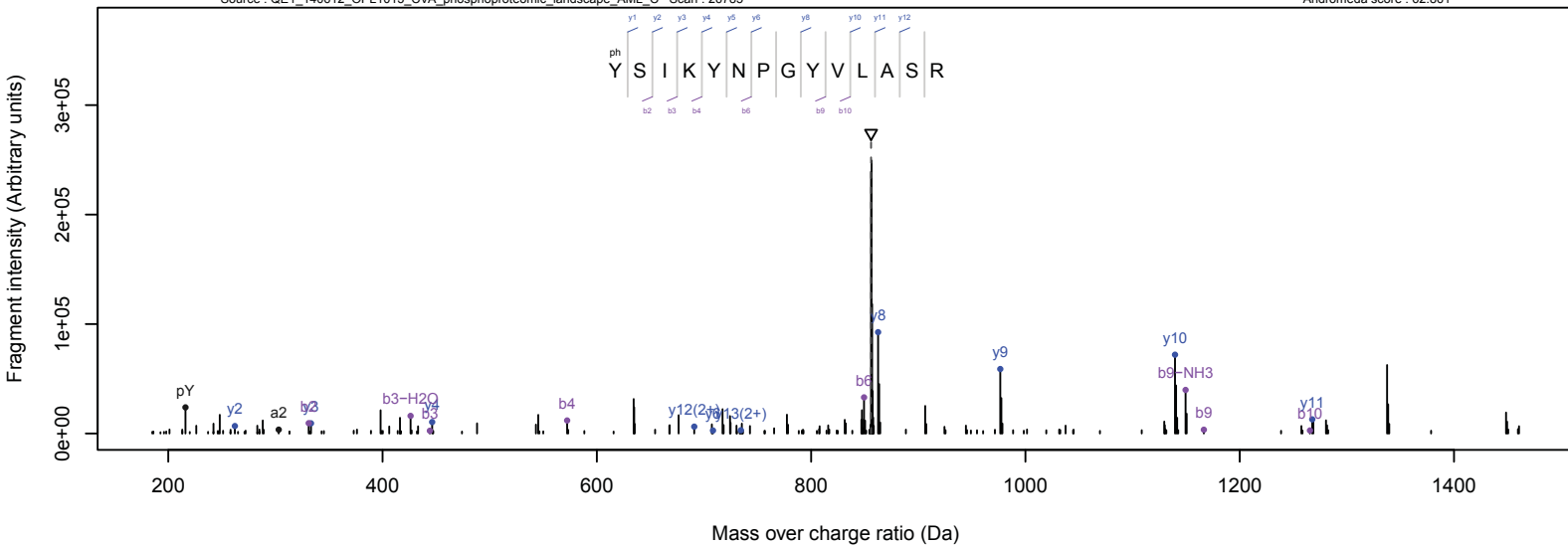
Andromeda score : 62.861



Y(ph)SIKYNPGYVLASR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26785

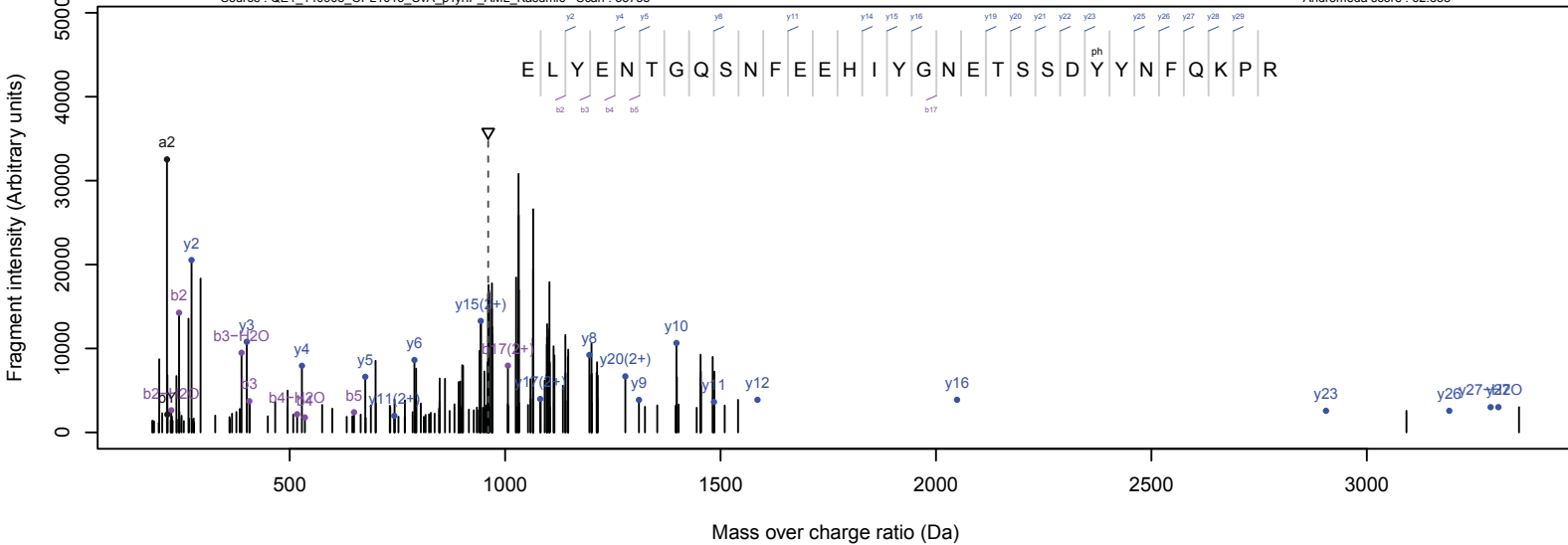
Andromeda score : 62.861



ELYENTGQSNFEEHIYGNETS(ph)YNFQKPR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 33738

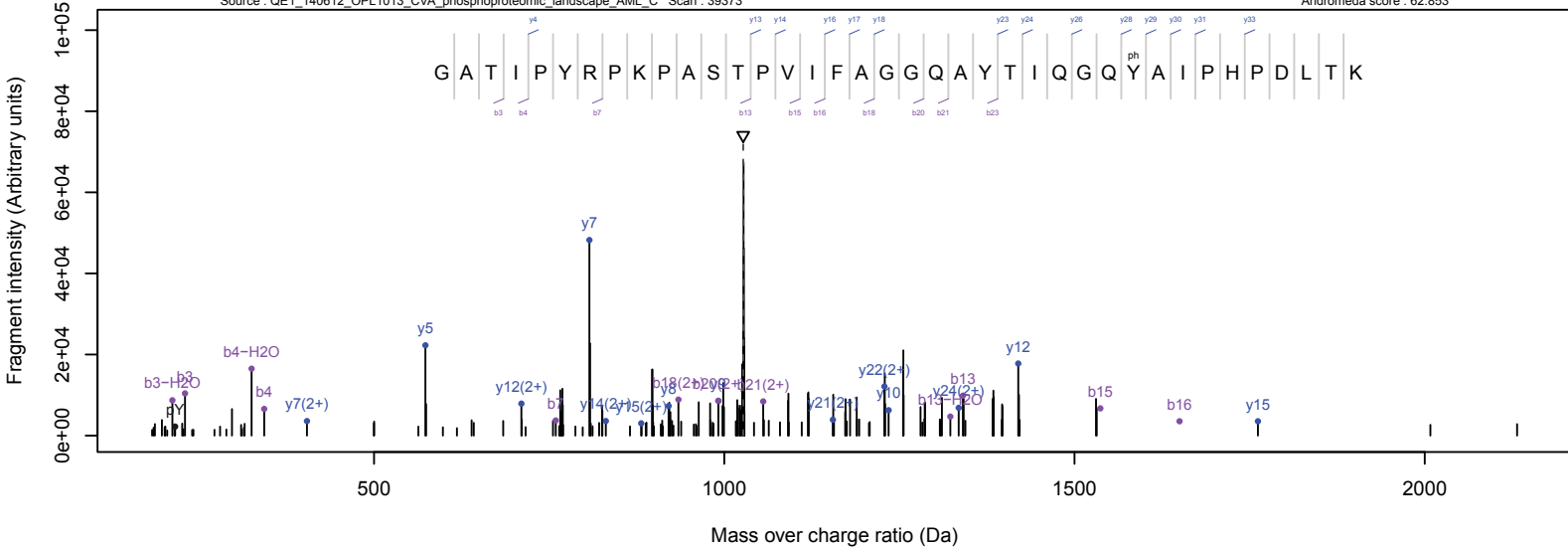
Andromeda score : 62.858



GATIPYRPKPASTPVI FAGGQAYTIQQQY(ph)AIPHPDLTK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39373

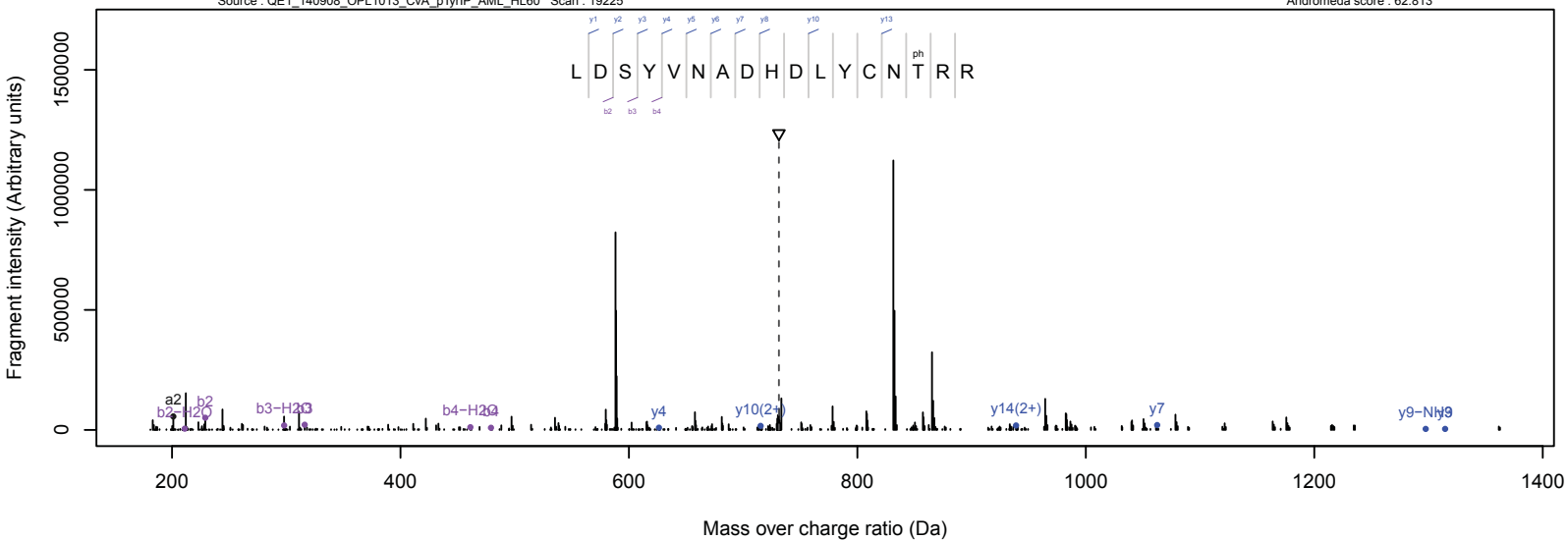
Andromeda score : 62.853



LDSYVNADHDLYCNT(ph)RR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 19225

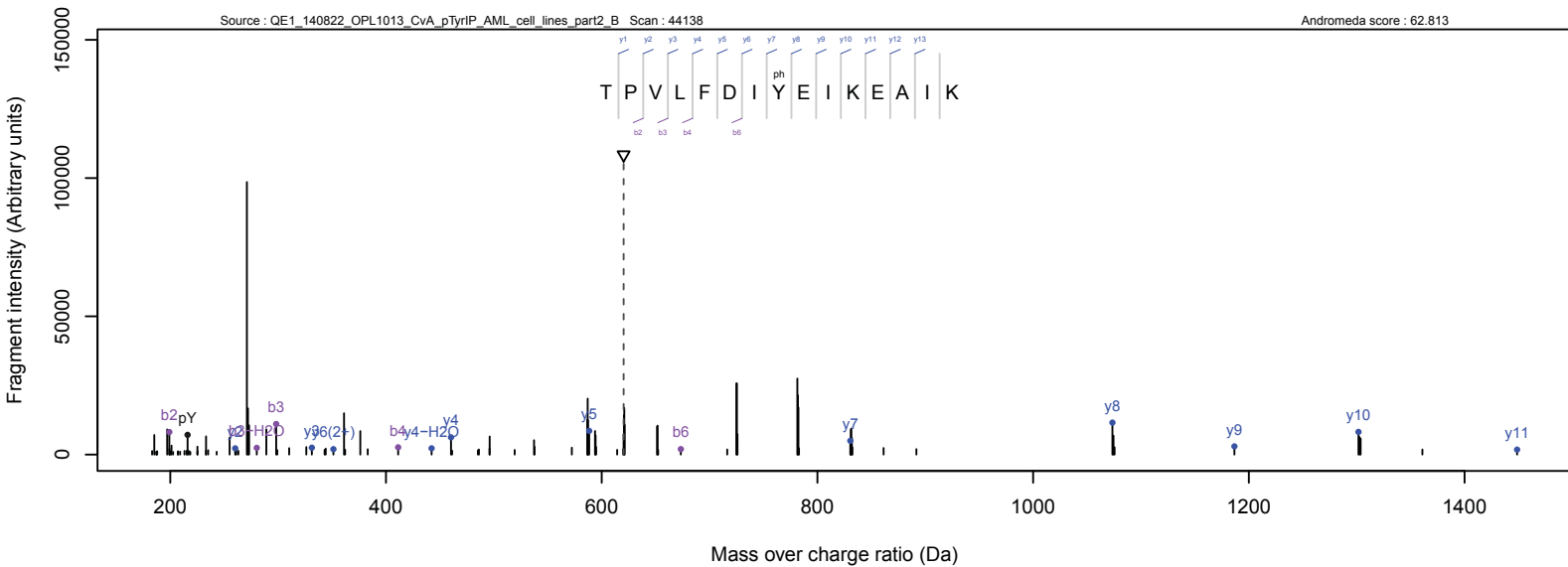
Andromeda score : 62.813



TPVLFDIY(ph)EIKEAIK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 44138

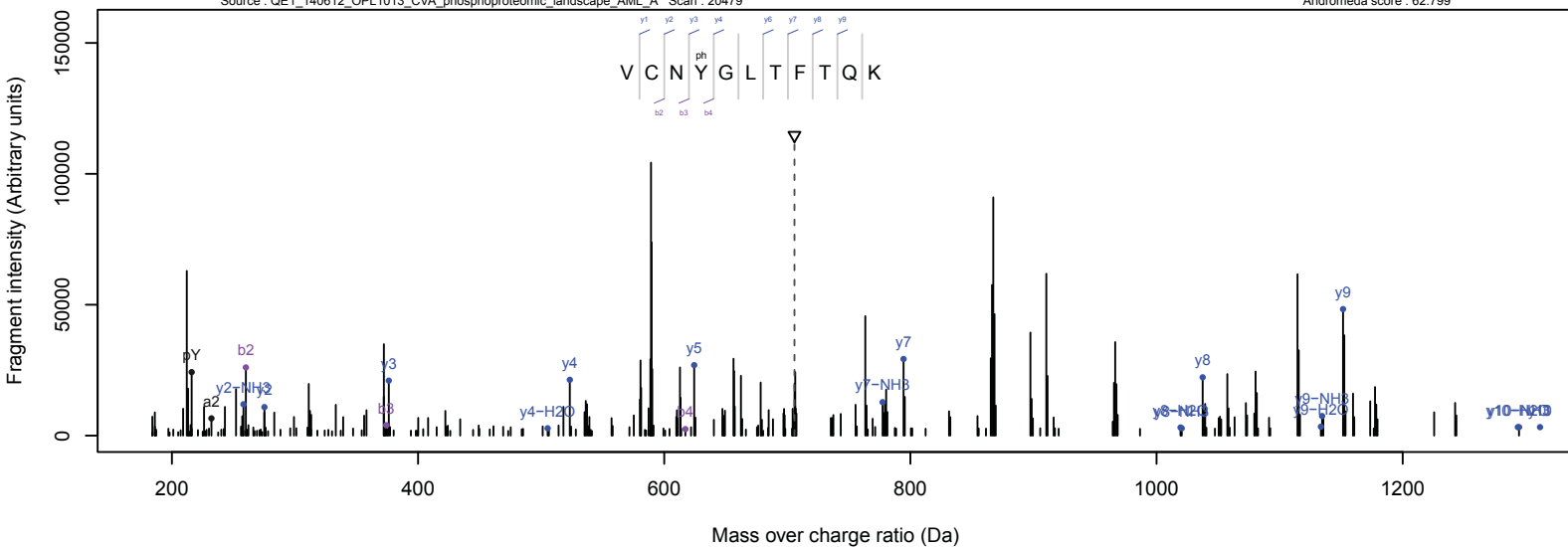
Andromeda score : 62.813



VCNY(ph)GLTFTQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 20479

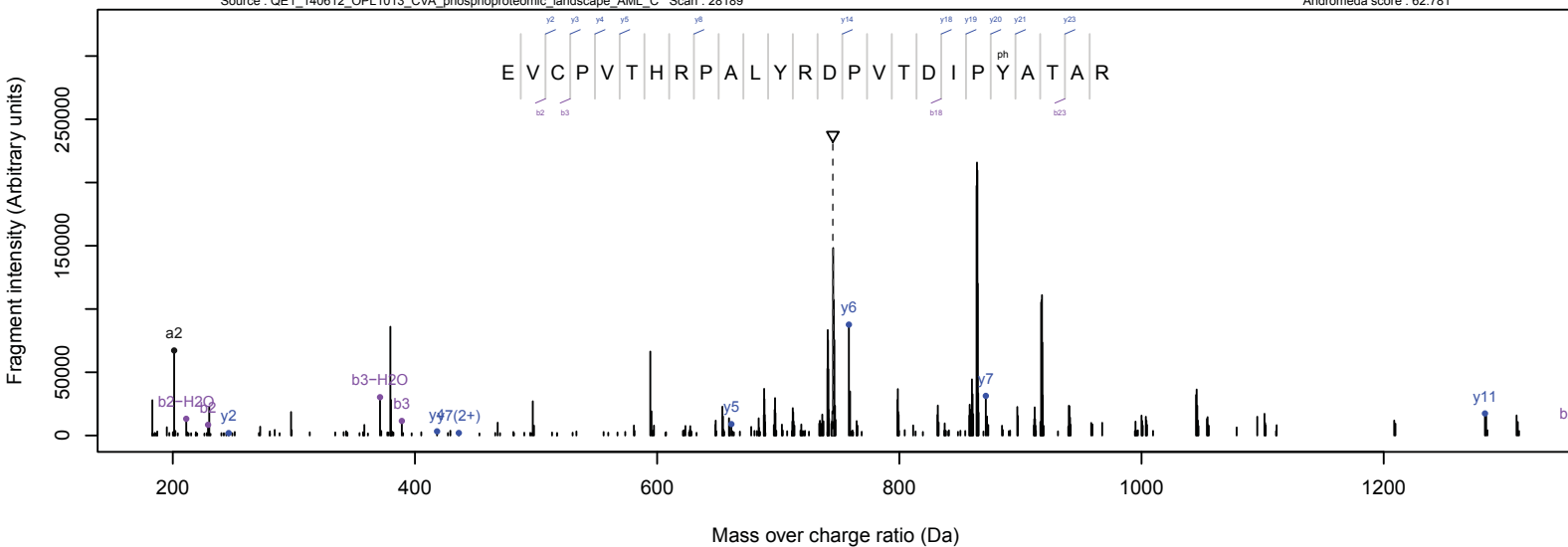
Andromeda score : 62.799



EVCPTVTHRPAlyRDPVTDIPY(ph)ATAR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28189

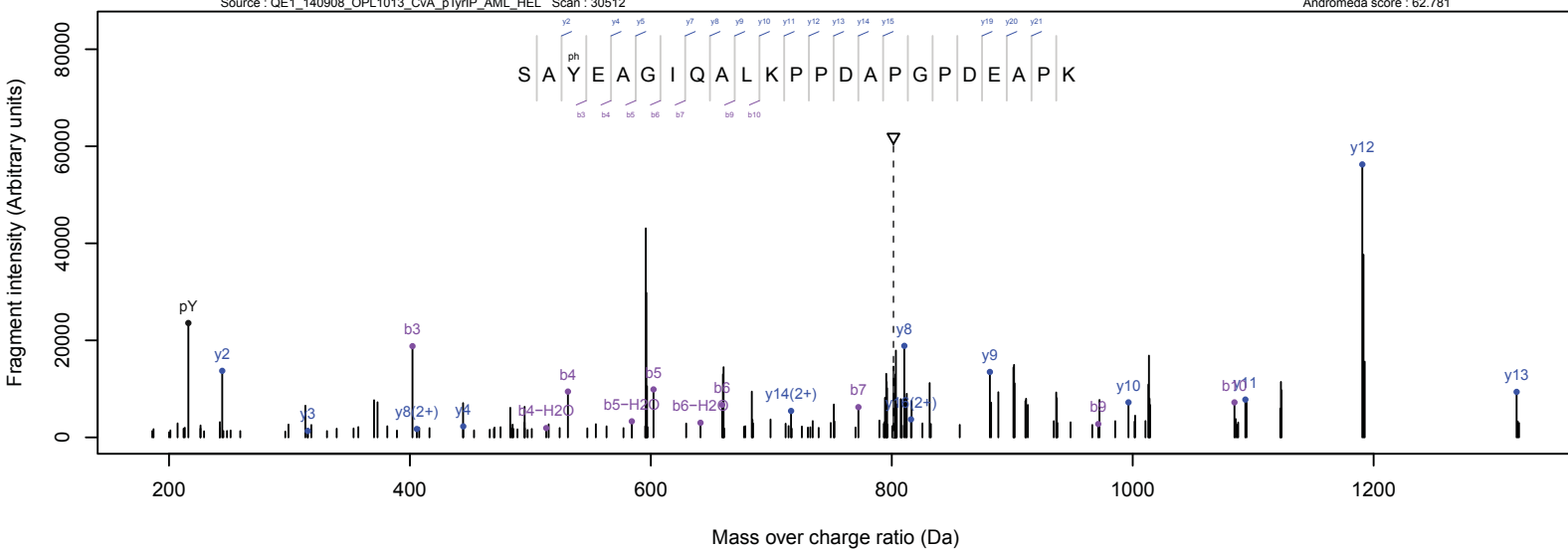
Andromeda score : 62.781



SAY(ph)EAGIQALKPPDAPGDEAPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 30512

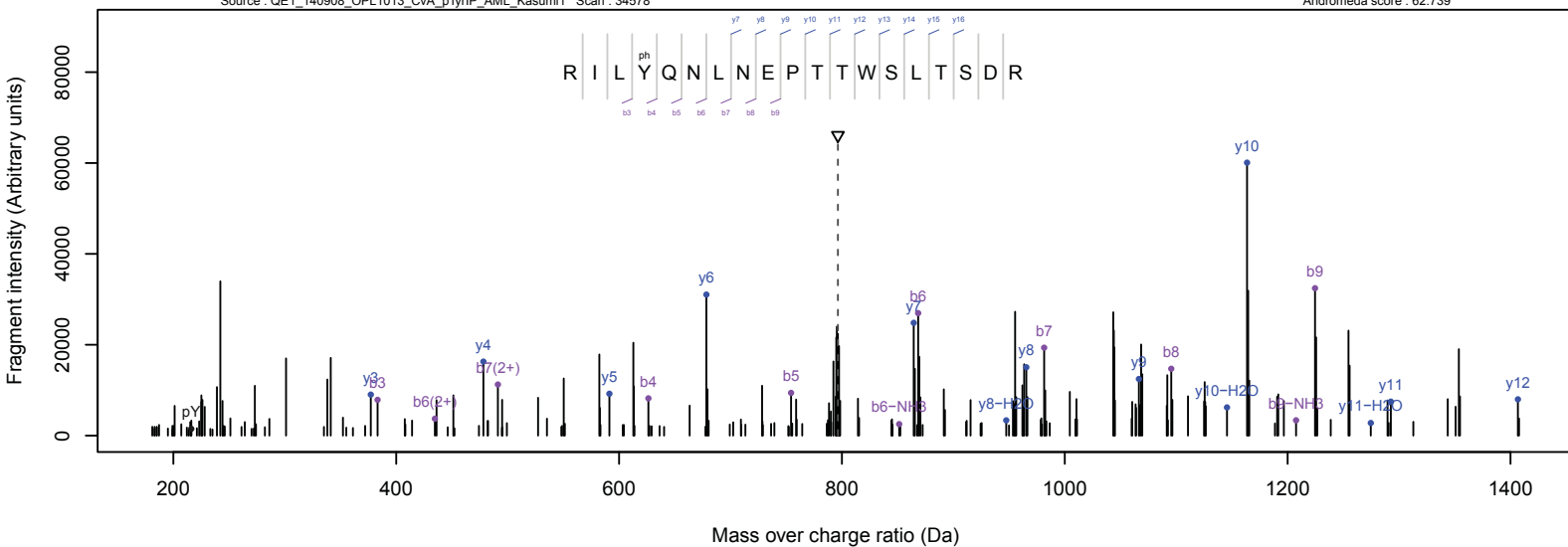
Andromeda score : 62.781



RILY(ph)QLNEPTTWSLTSR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 34578

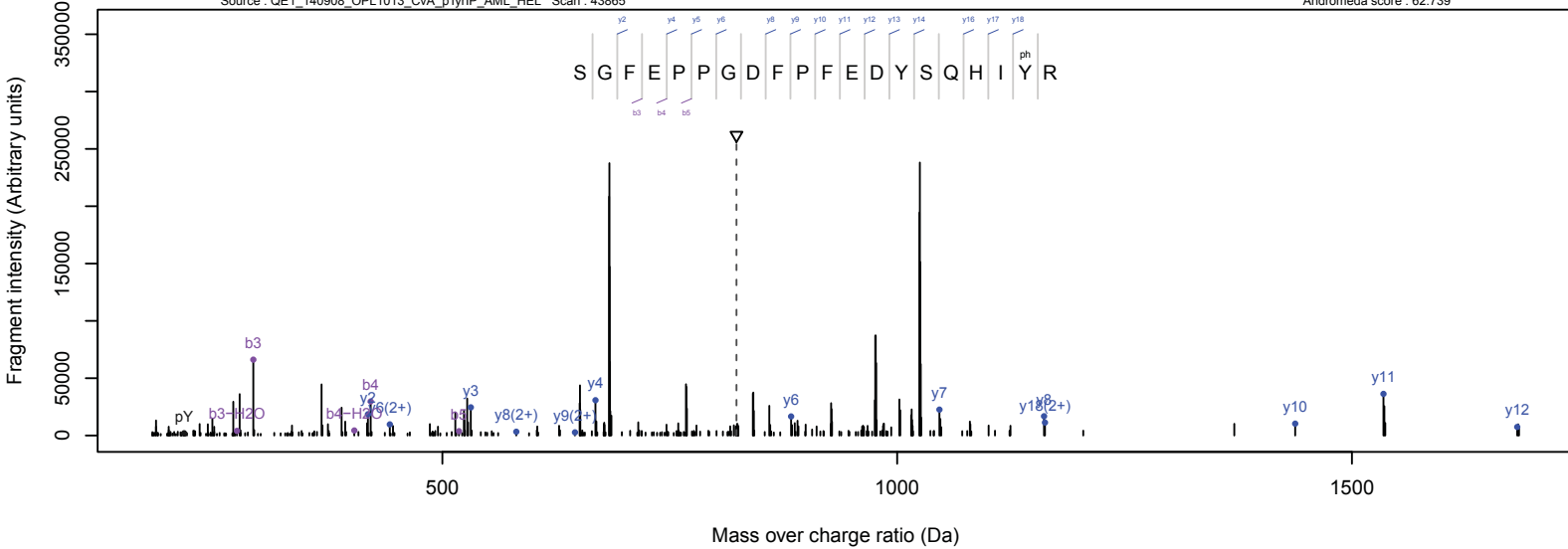
Andromeda score : 62.739



SGFEPPGDFPFEDYSQH(Y)R-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 43865

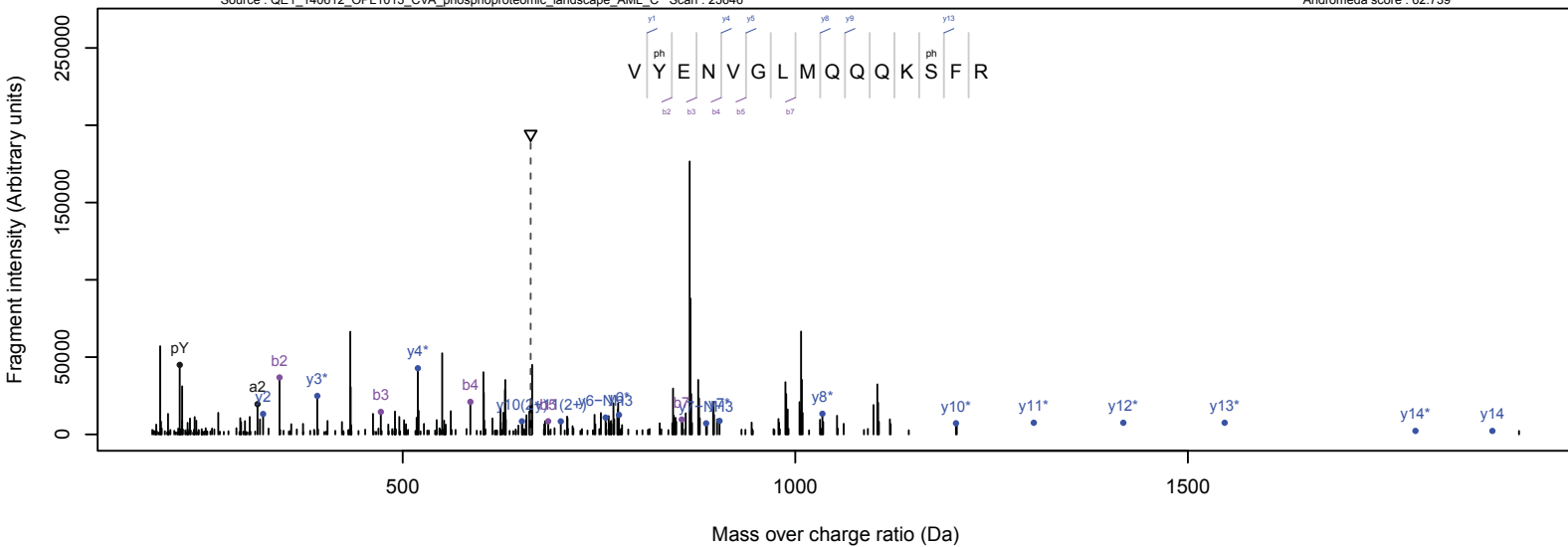
Andromeda score : 62.739



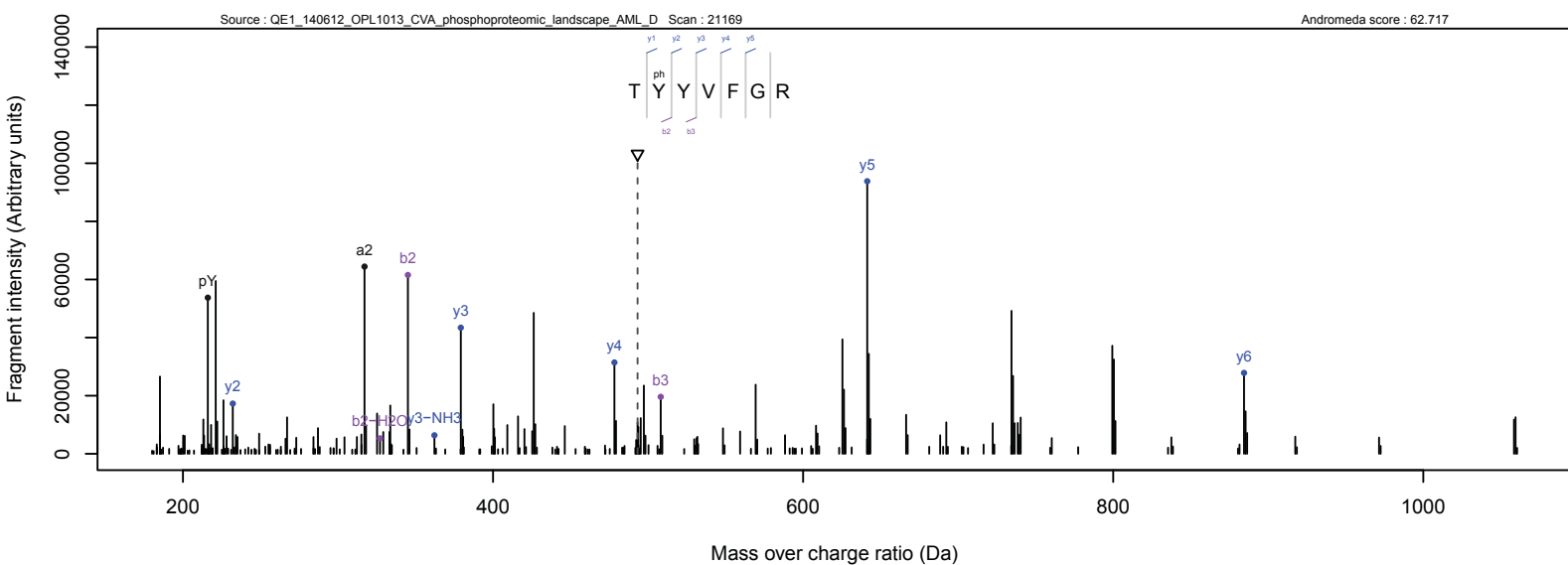
VY(ph)ENVGLMQQKS(ph)FR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23646

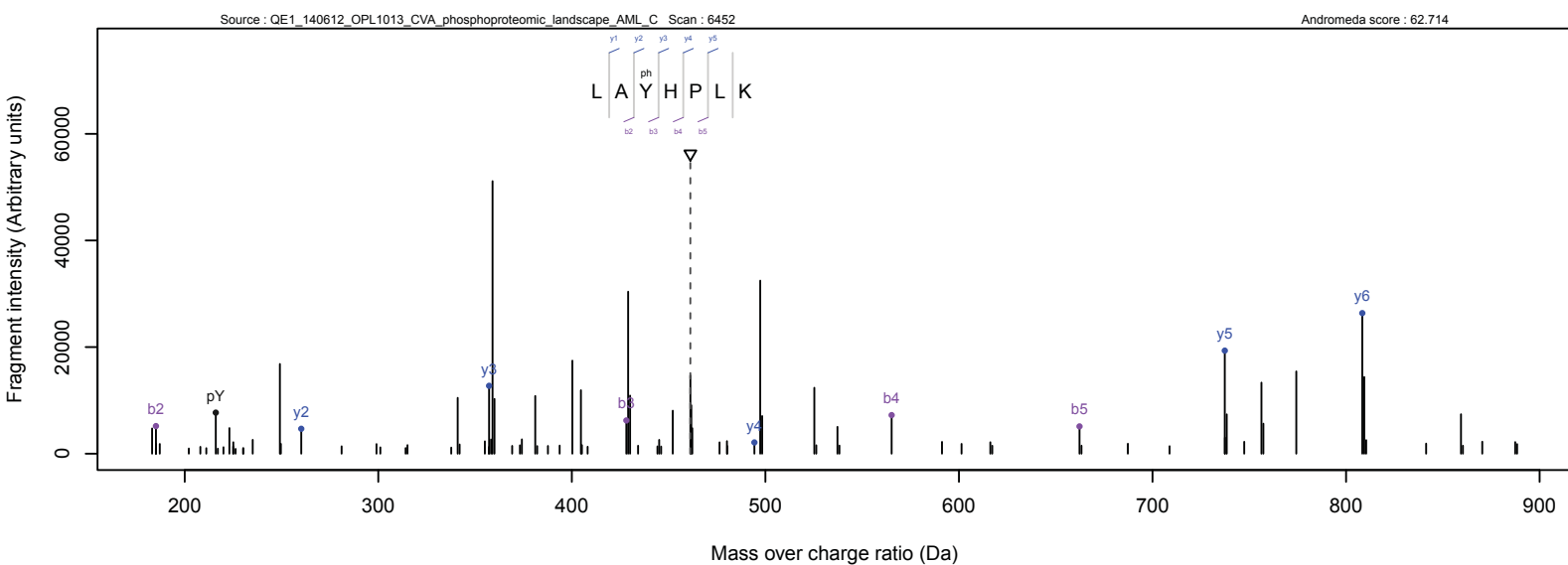
Andromeda score : 62.739



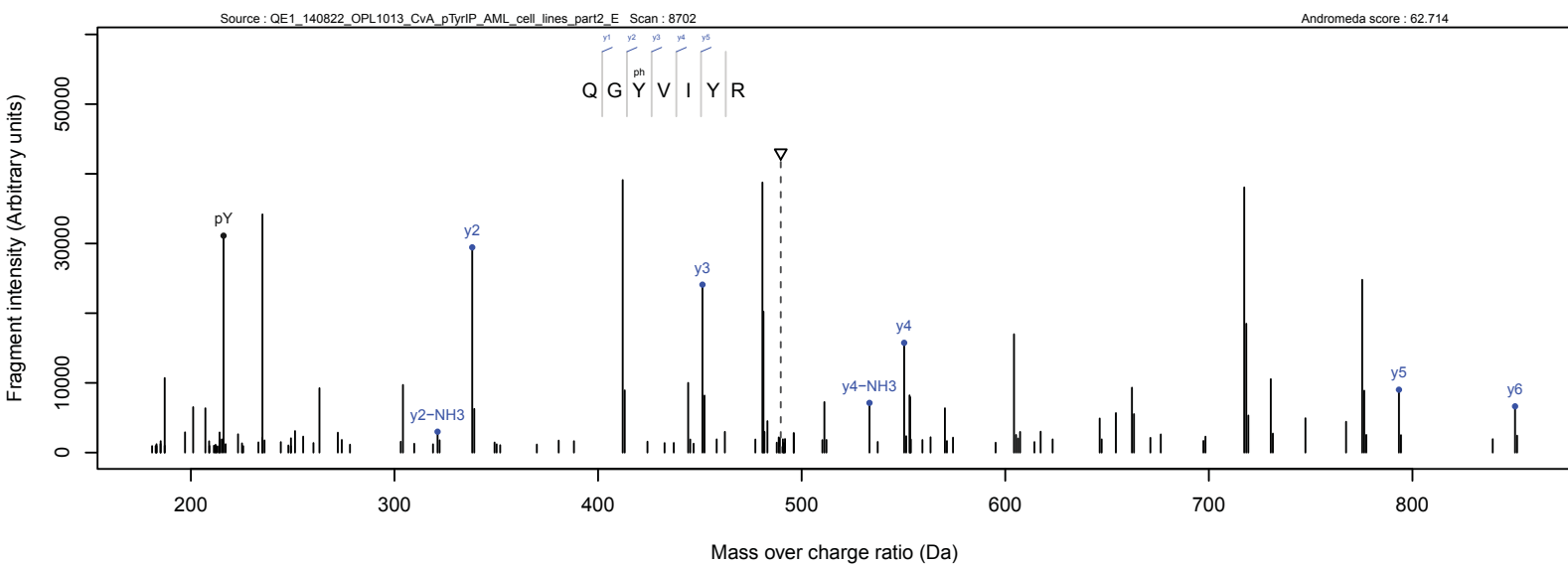
## TY(ph)YVFGR-2+



## LAY(ph)HPLK-2+



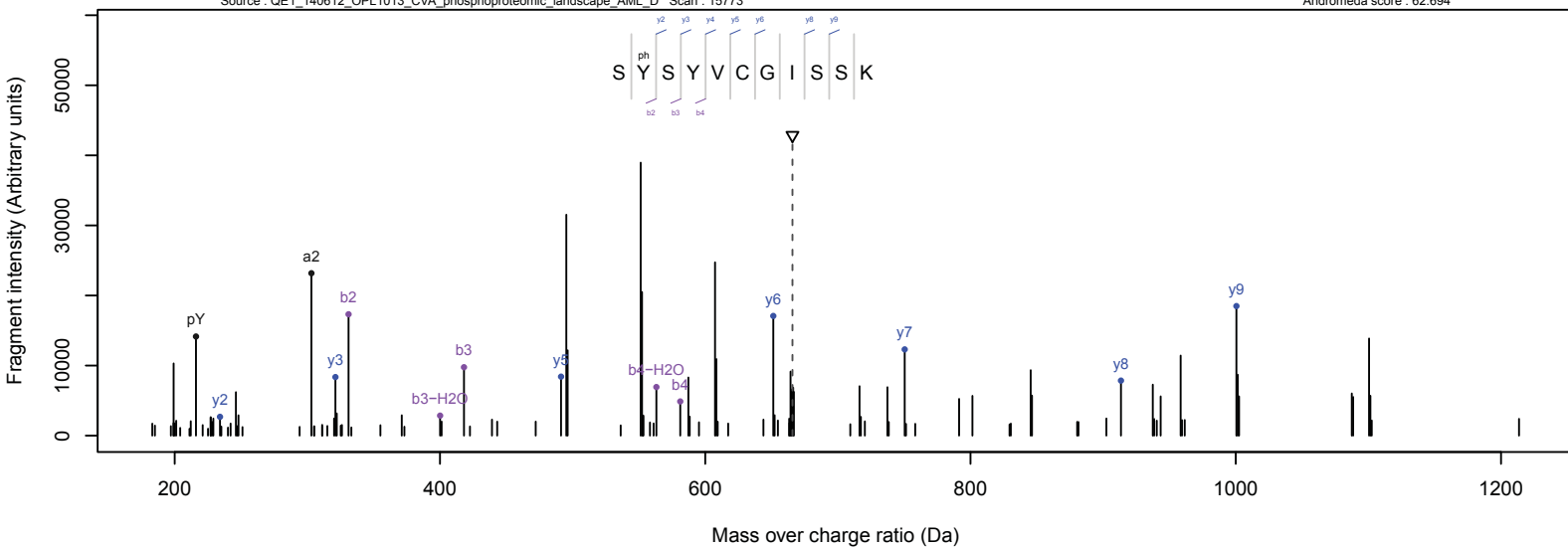
## QGY(ph)VIYR-2+



## SY(ph)SYVCGISSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 15773

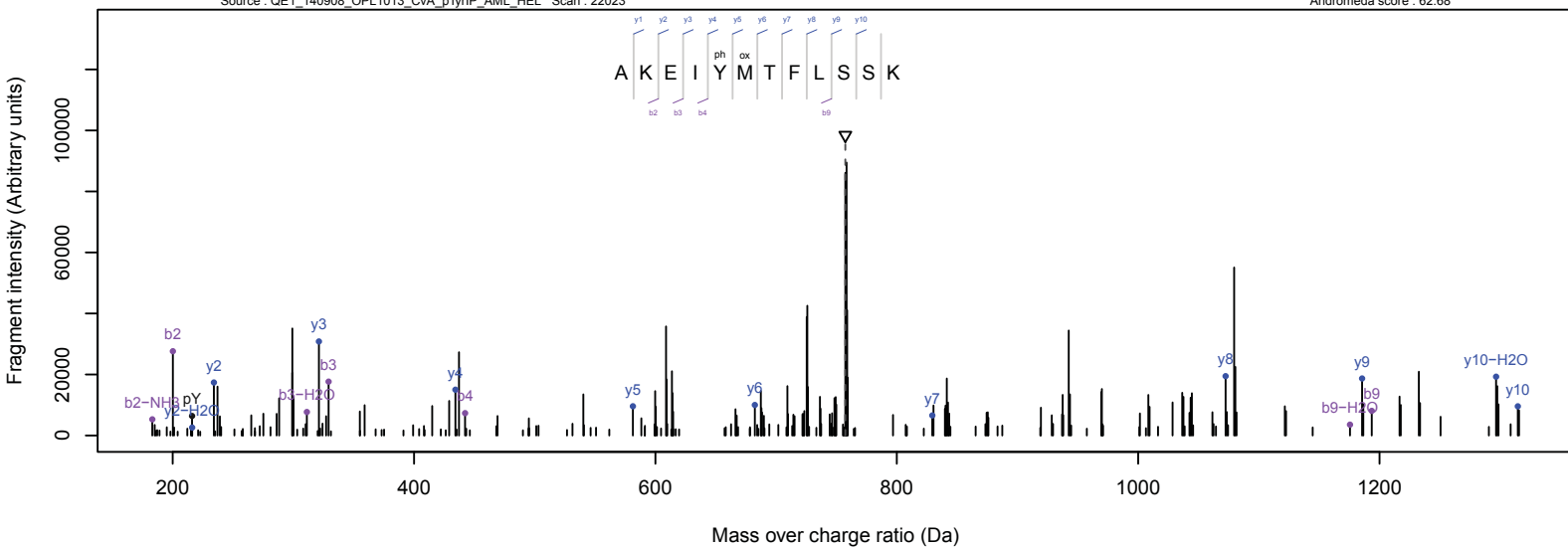
Andromeda score : 62.694



## AKEIY(ph)M(ox)TFLSSK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 22023

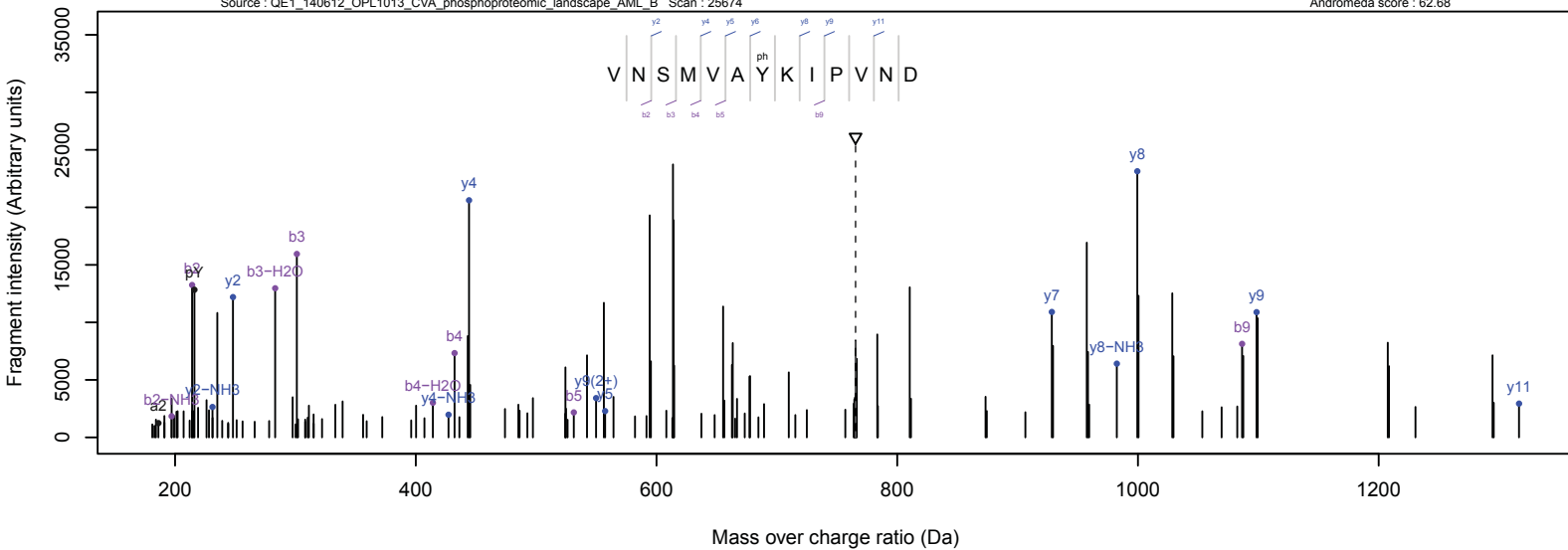
Andromeda score : 62.68



## VNSMVAY(ph)KIPVND-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25674

Andromeda score : 62.68

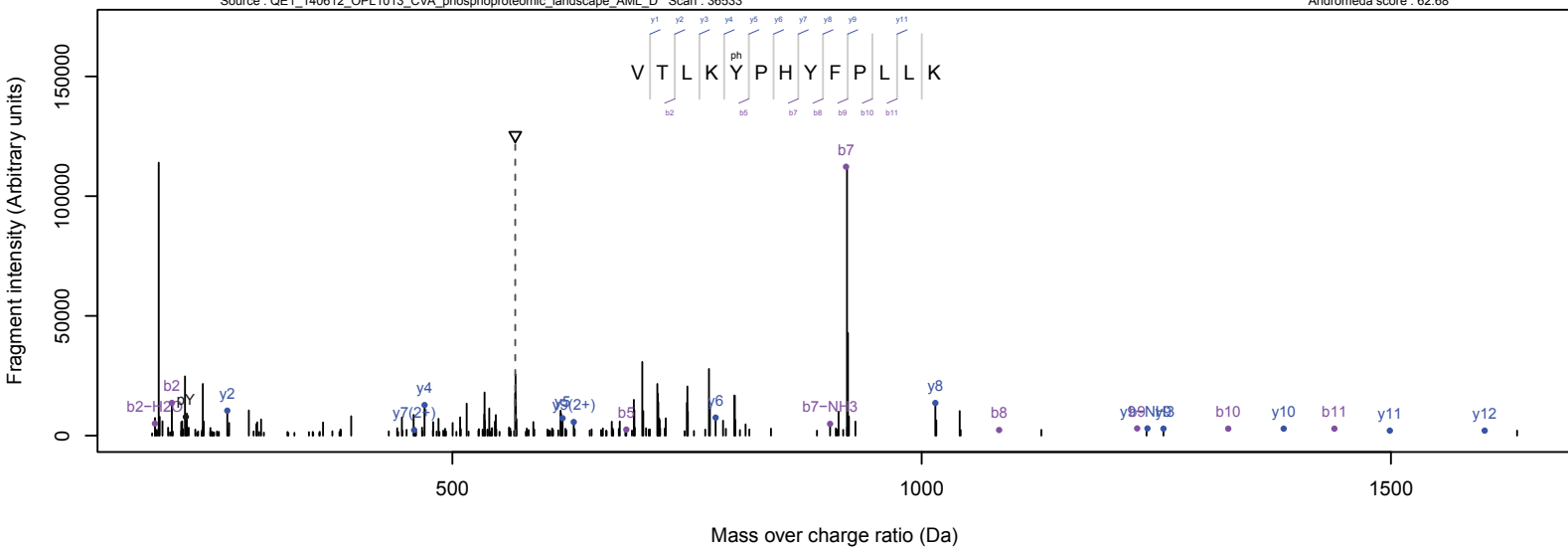




VTLKY(ph)PHYFLLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 36533

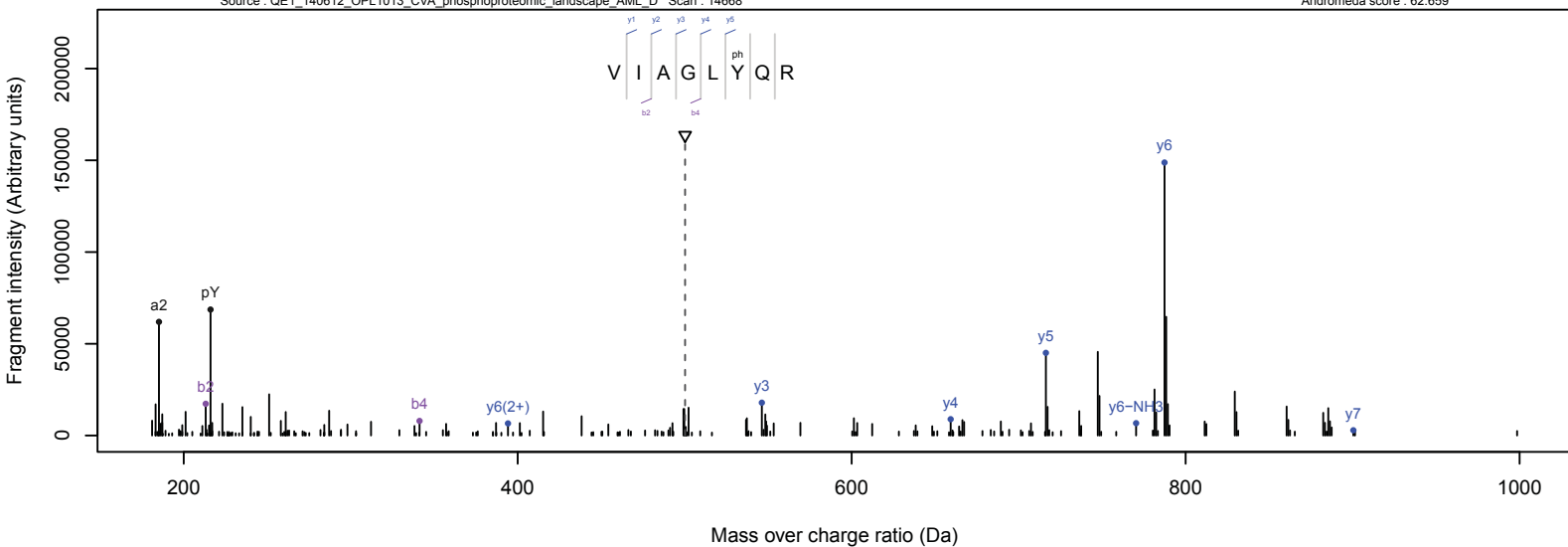
Andromeda score : 62.68



VIAGLY(ph)QR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14668

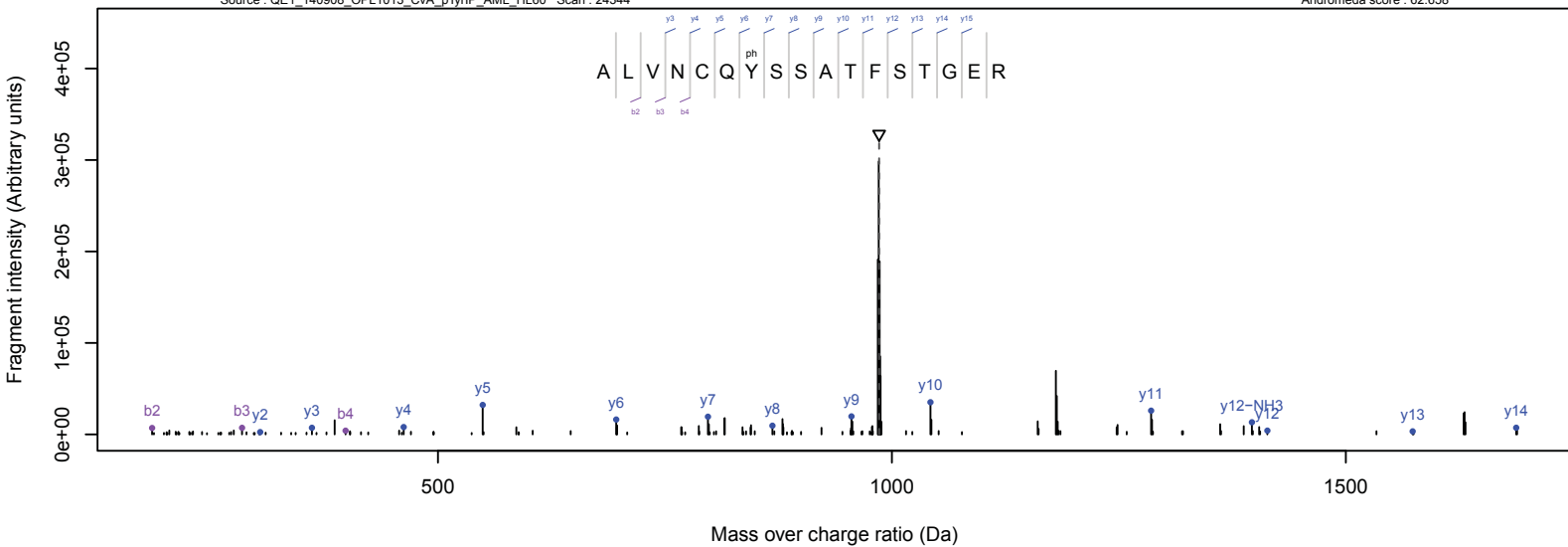
Andromeda score : 62.659



ALVNCQY(ph)SSATFSTGER-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 24344

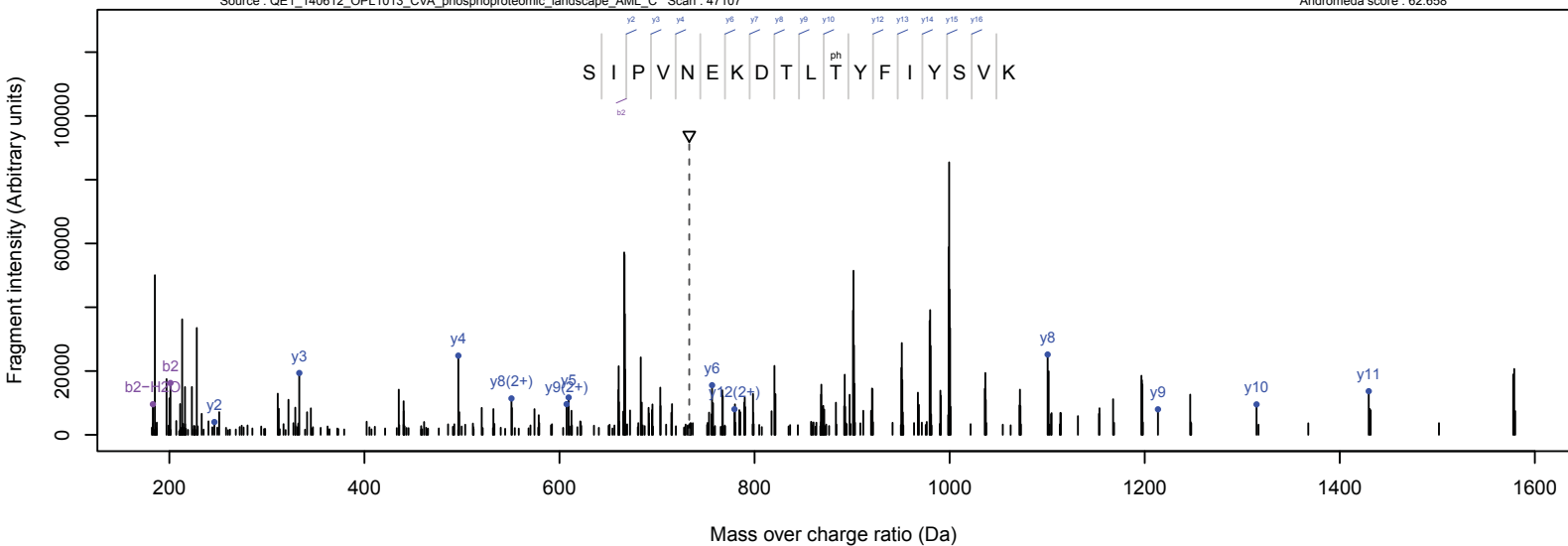
Andromeda score : 62.658



## SIPVNEKDTLT(ph)YFIYSVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 47107

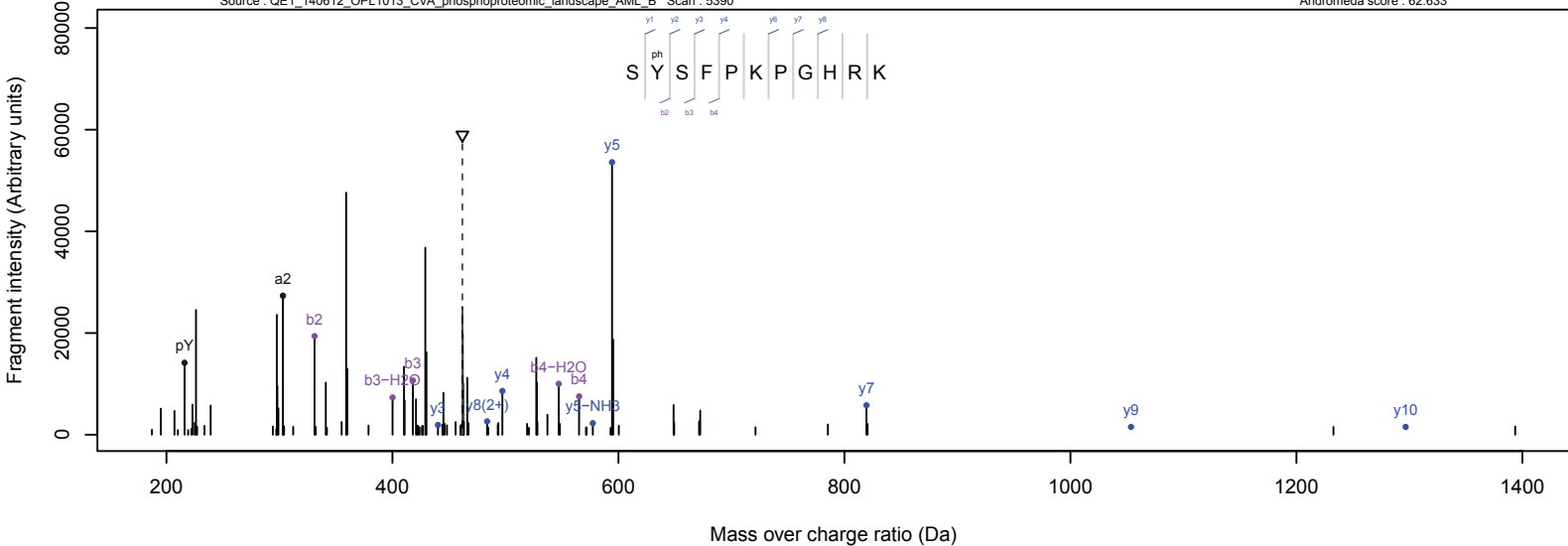
Andromeda score: 62.658



## SY(ph)SFPKPGHRK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 5390

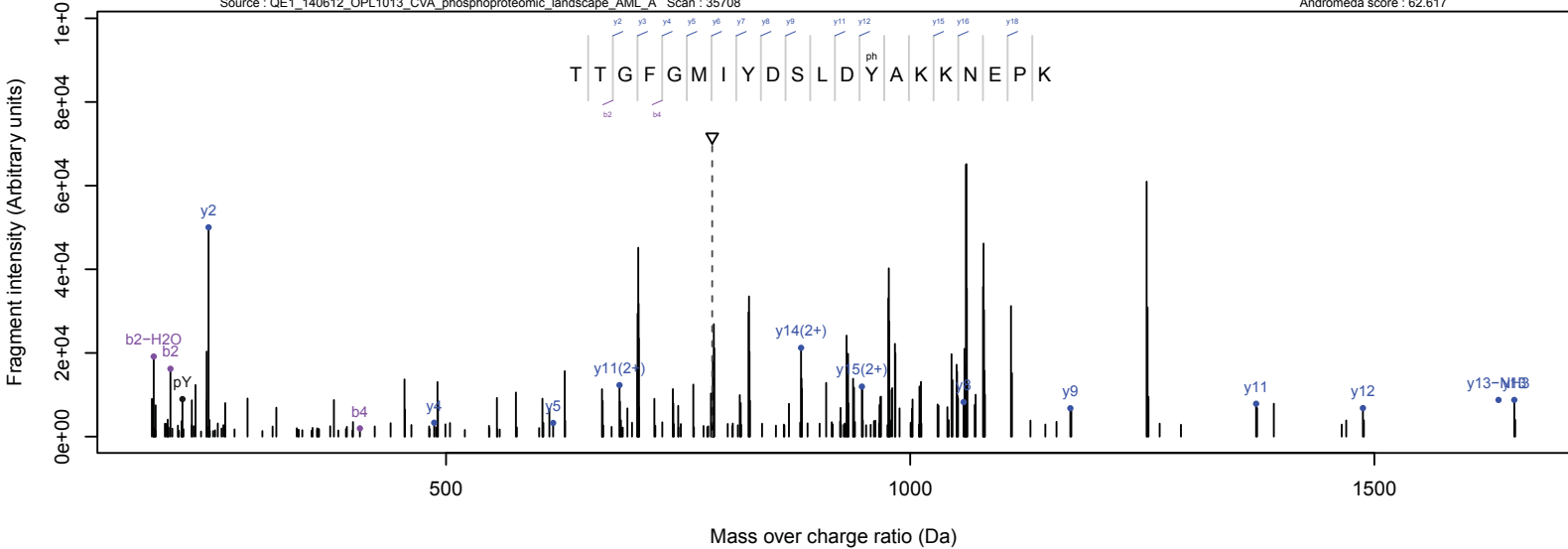
Andromeda score: 62.633



## TTGFGMIYDSL DY(ph)AKKNEPK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 35708

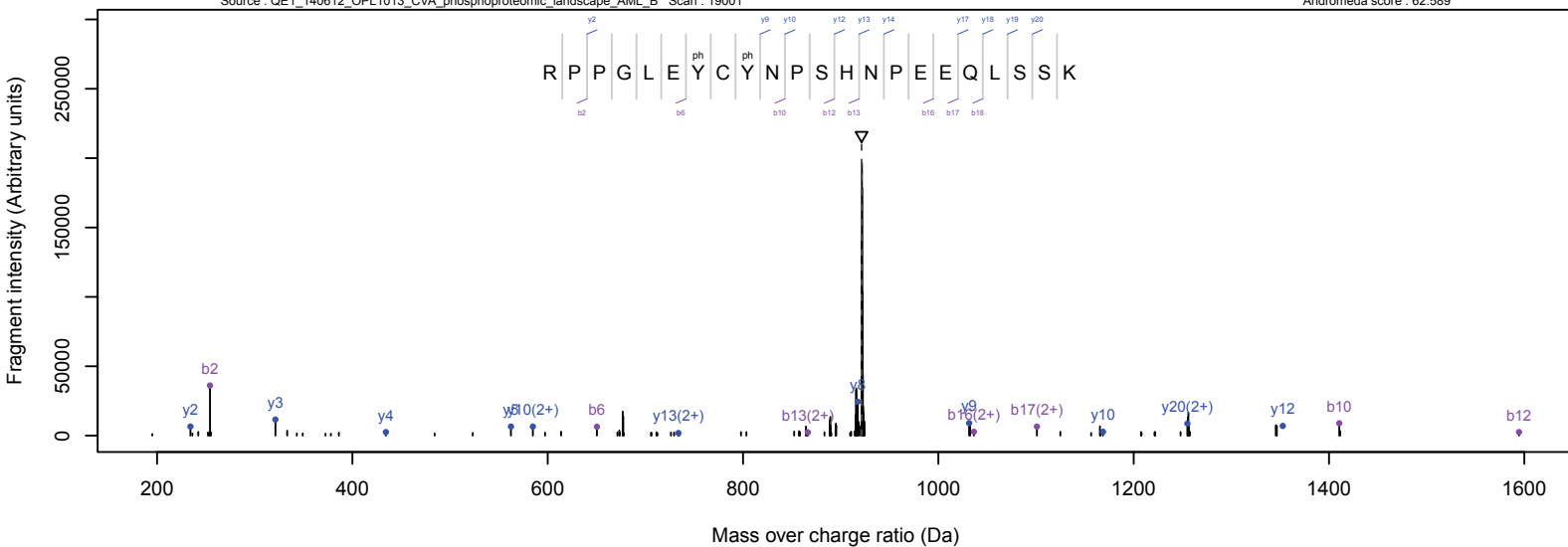
Andromeda score: 62.617



## RPPGLEYP(h)CYNPESHNP EEQLSSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 19001

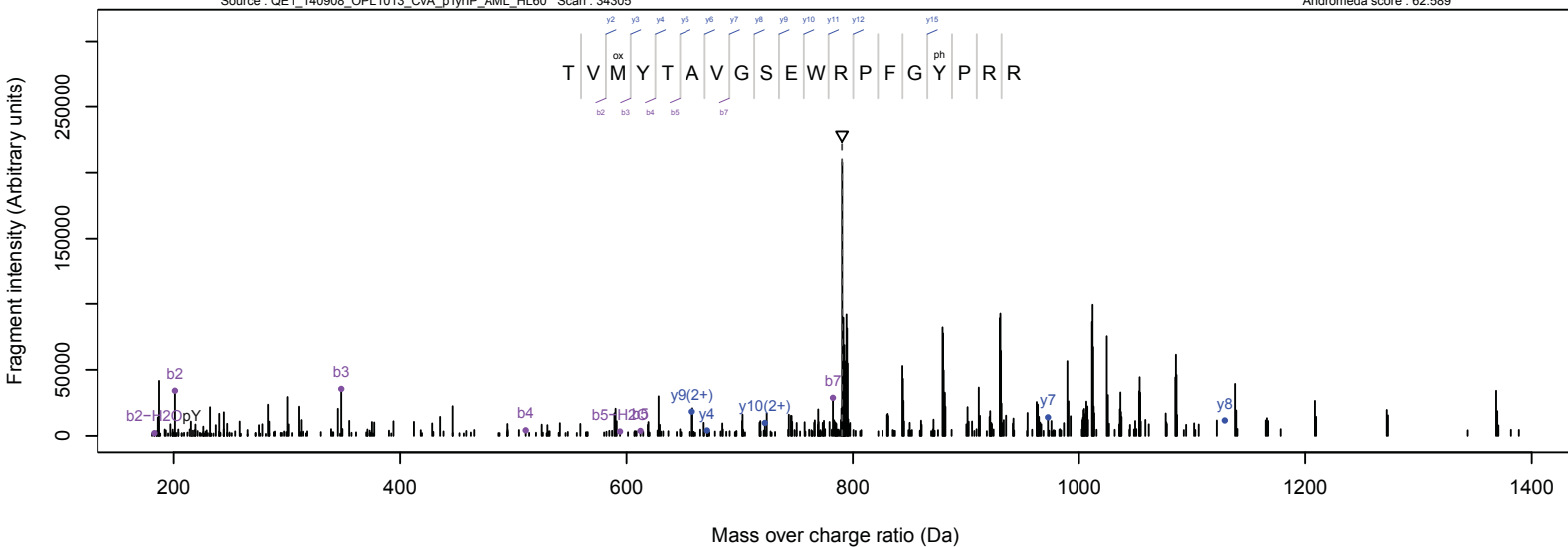
Andromeda score: 62.589



## TVM(ox)YTAGVSEWRPFY(h)PRR-3+

Source: QE1\_140908\_OPL1013\_Cva\_pTyrIP\_AML\_HL60 Scan: 34305

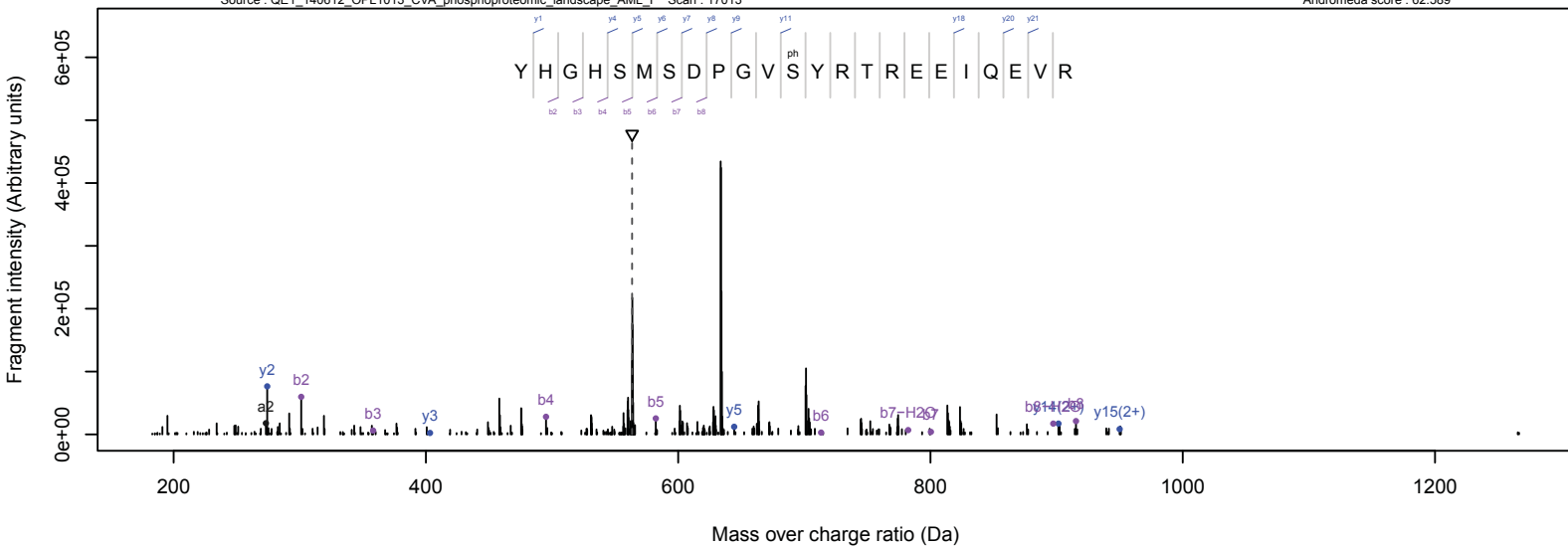
Andromeda score: 62.589



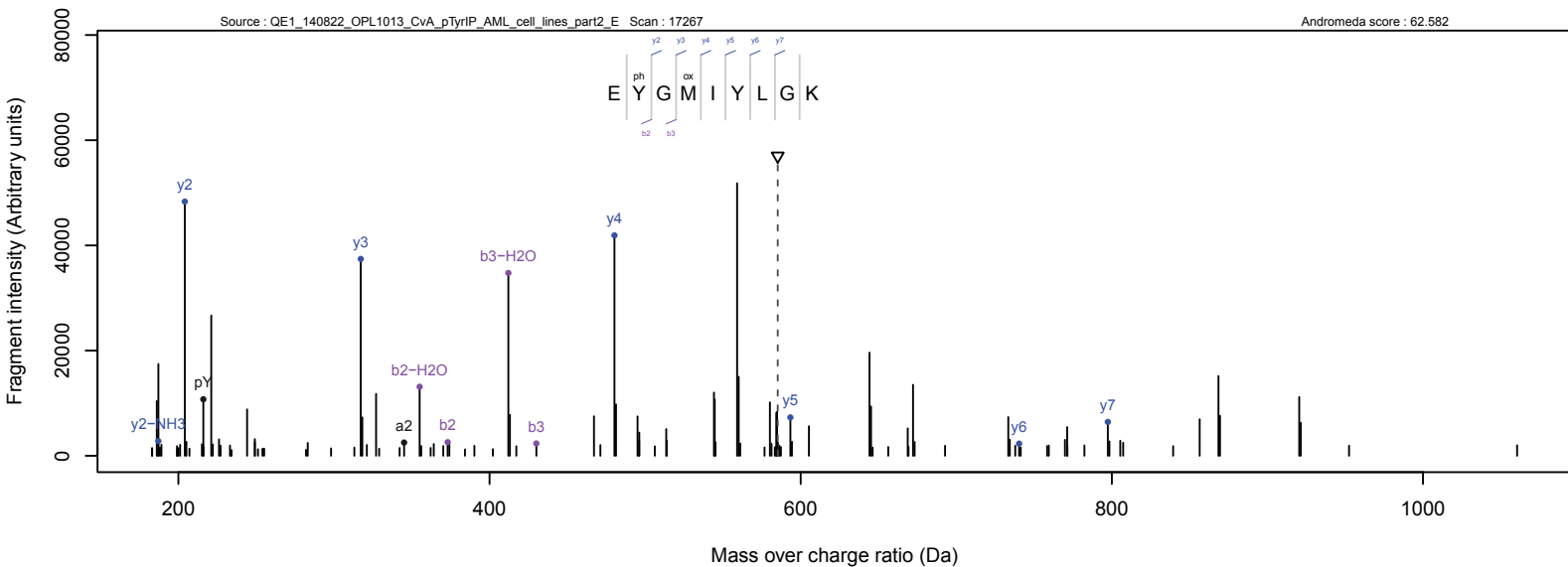
## YHGHSMDPGVS(h)YRTREEIQEVR-5+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan: 17013

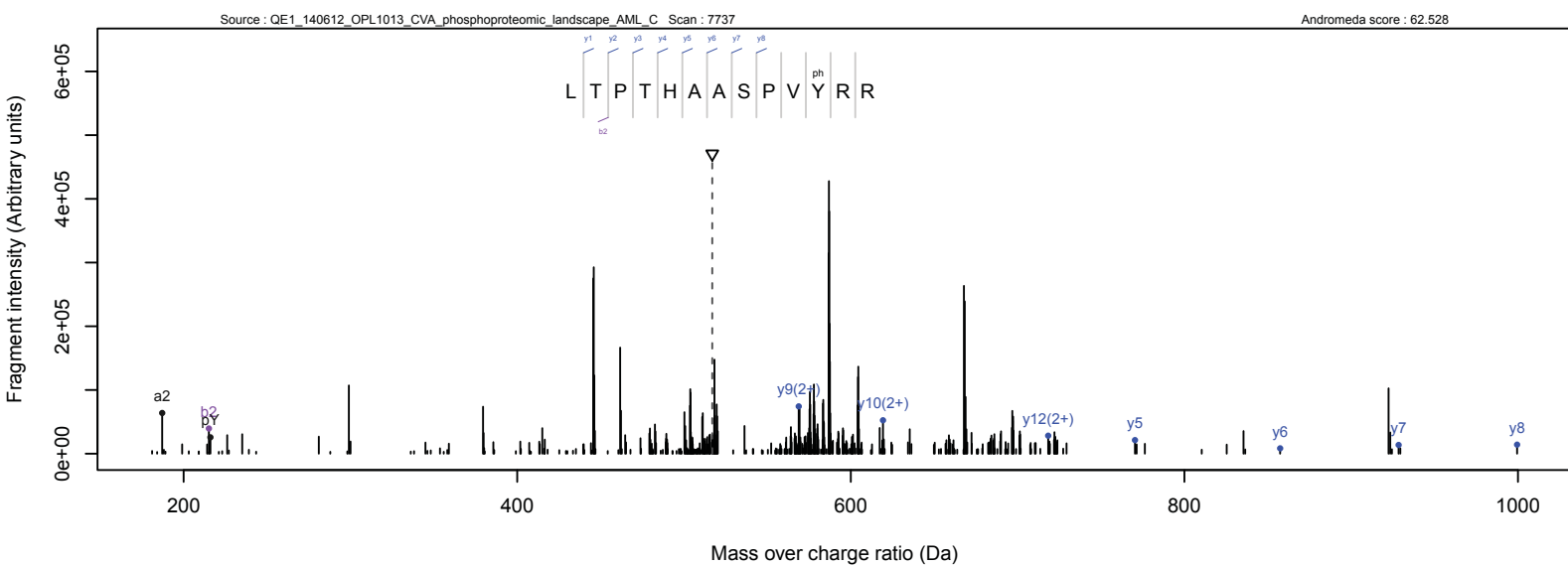
Andromeda score: 62.589



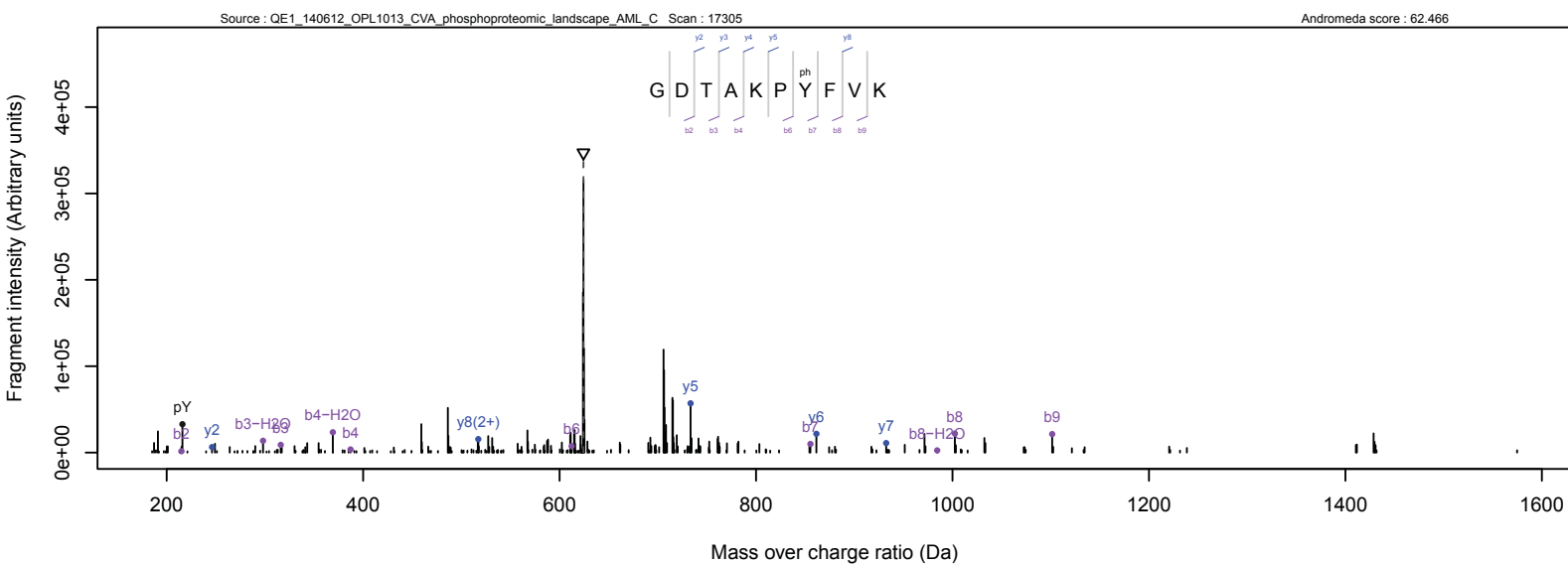
## EY(ph)GM(ox)IYLGK-2+



## LTPTHAASPVY(ph)RR-3+



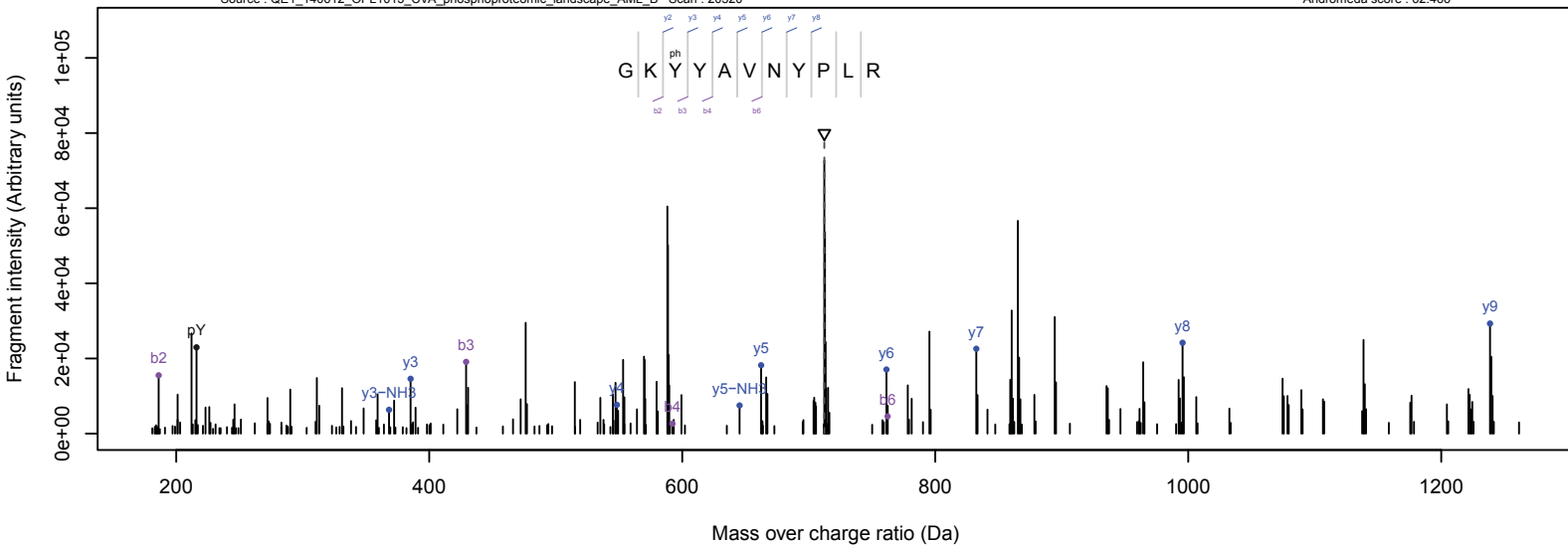
## (ac)GDTAKPY(ph)FVK-2+



## GKY(ph)YAVNYPLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20520

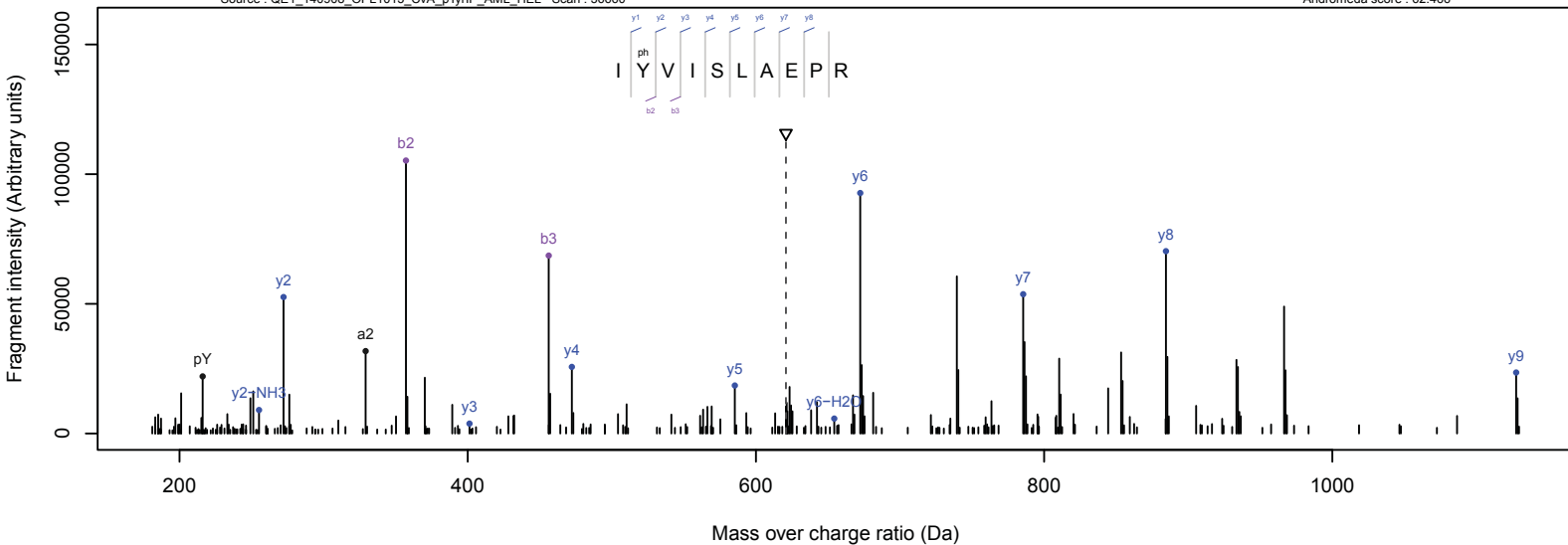
Andromeda score : 62.466



## IY(ph)VISLAEPR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 36660

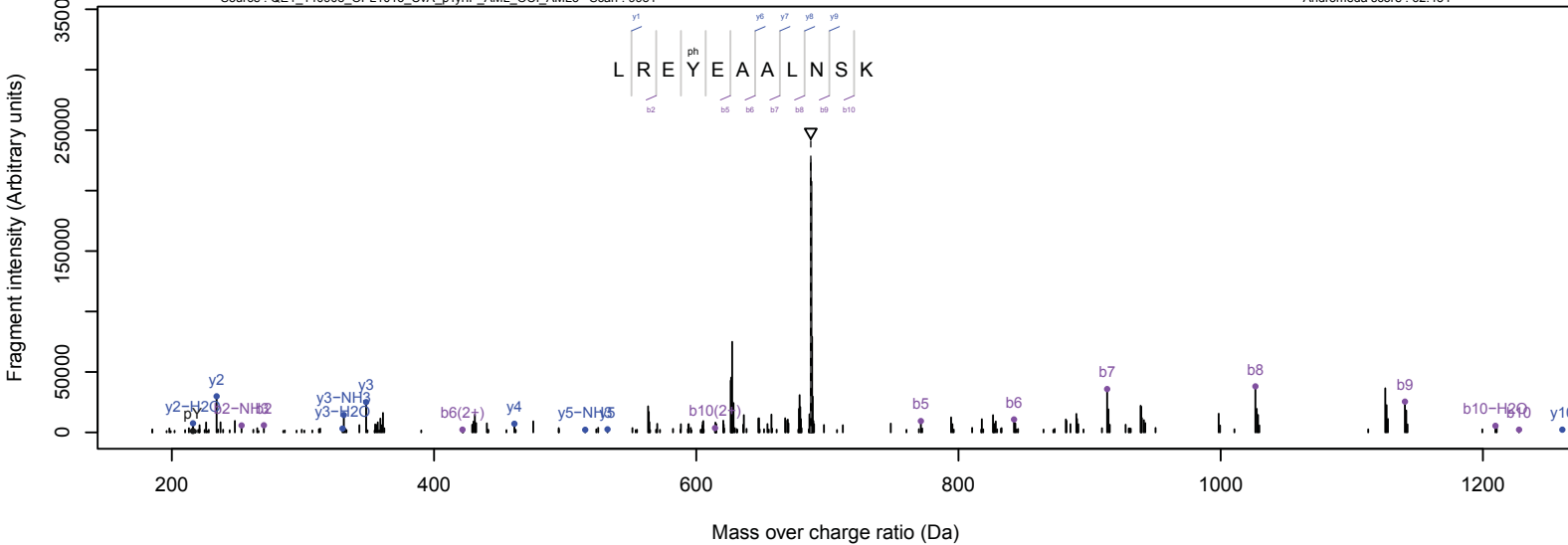
Andromeda score : 62.466



## LREY(ph)EAALNSK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 9951

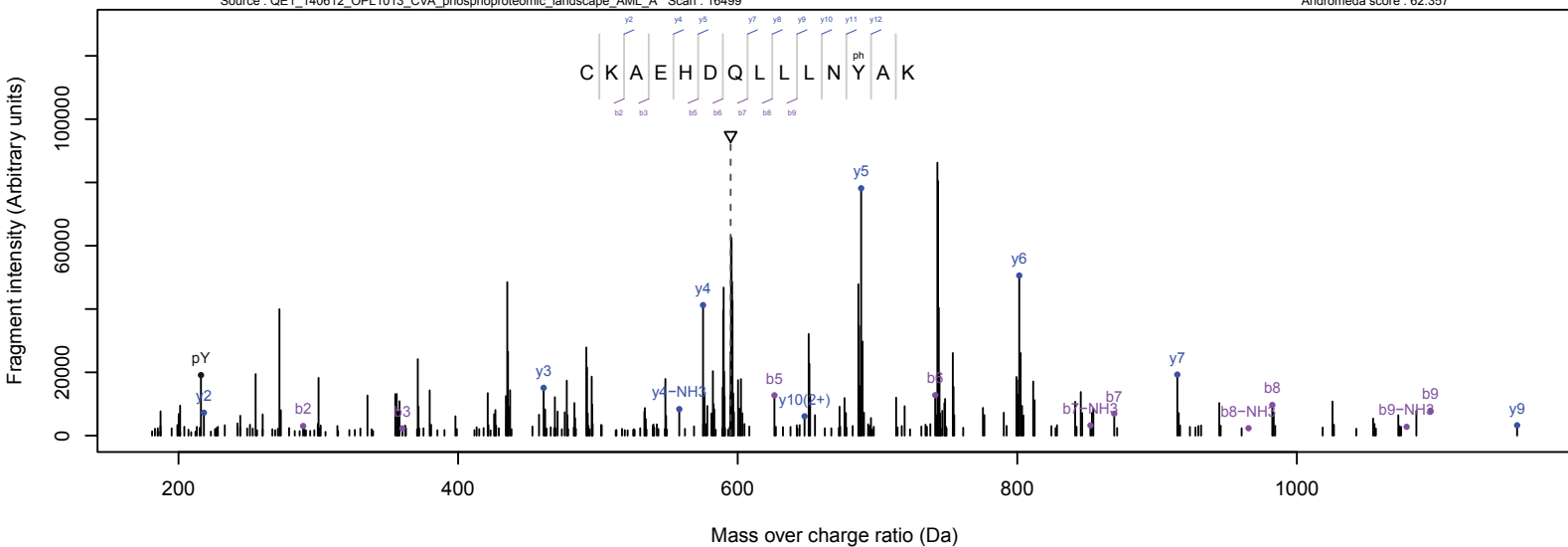
Andromeda score : 62.454



## CKAEHDQLLLNYAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16499

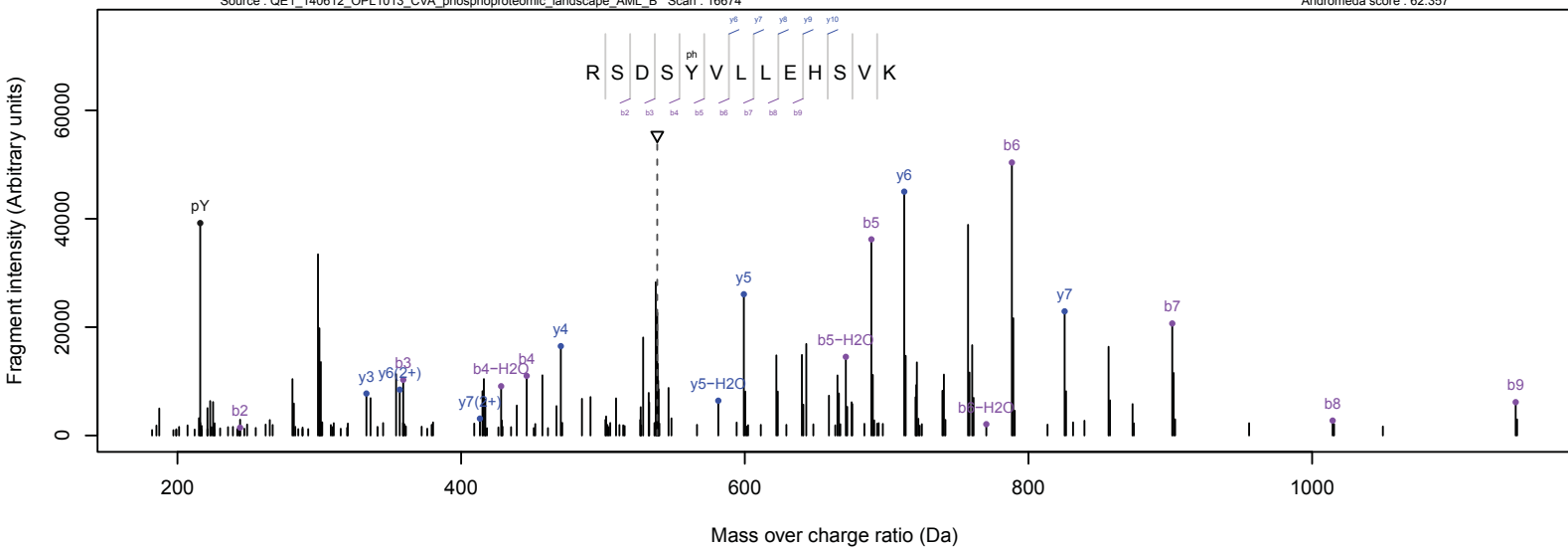
Andromeda score : 62.357



## RSDSY(ph)VLEHSVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16674

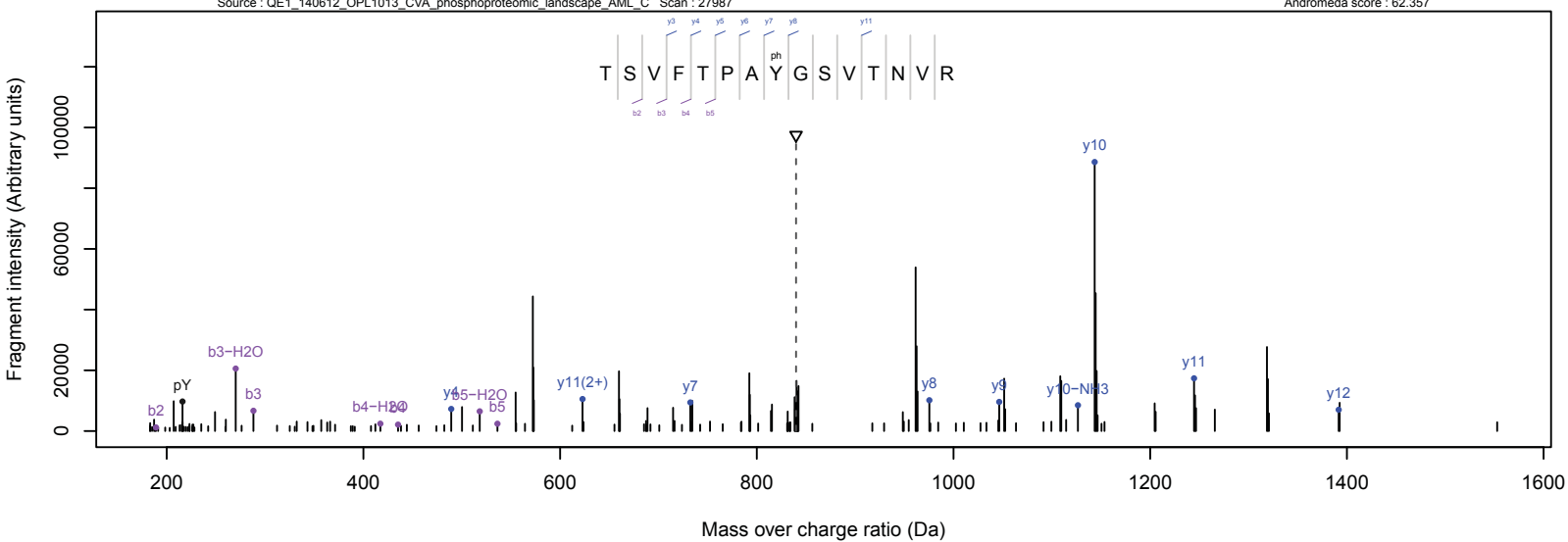
Andromeda score : 62.357



## TSVFTPAY(ph)GSVTNVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27987

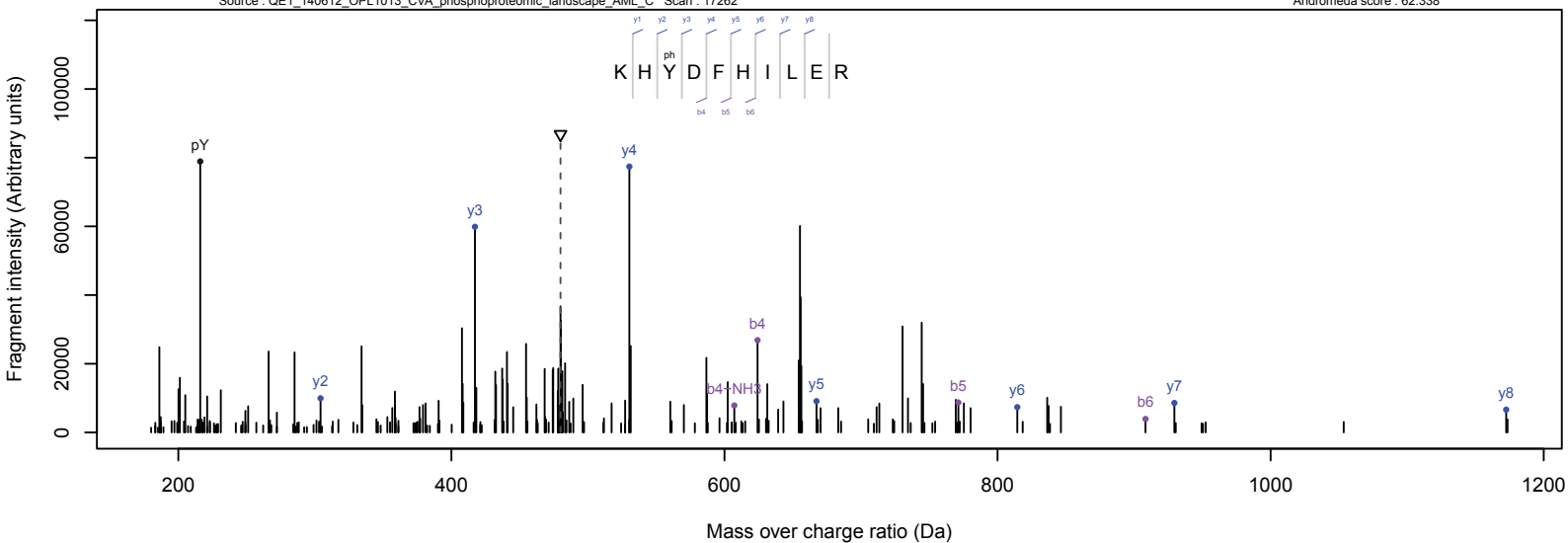
Andromeda score : 62.357



**KHY(ph)DFHILER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17262

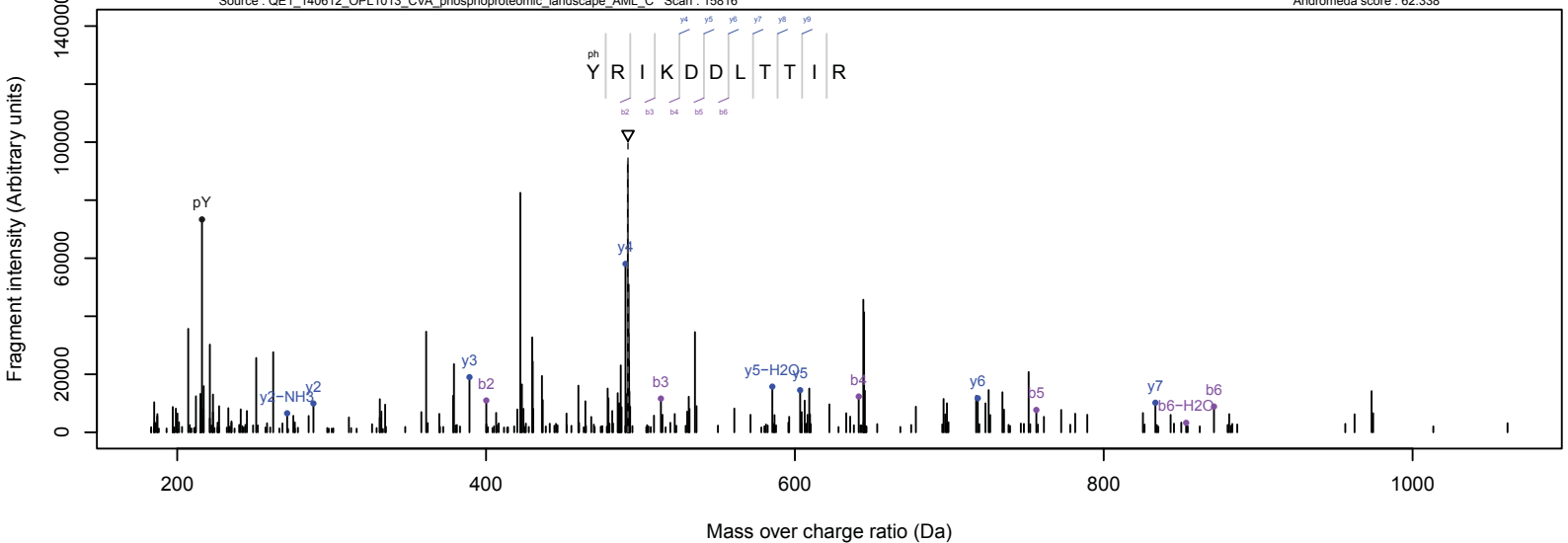
Andromeda score : 62.338



**Y(ph)RIKDDLTTIR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15816

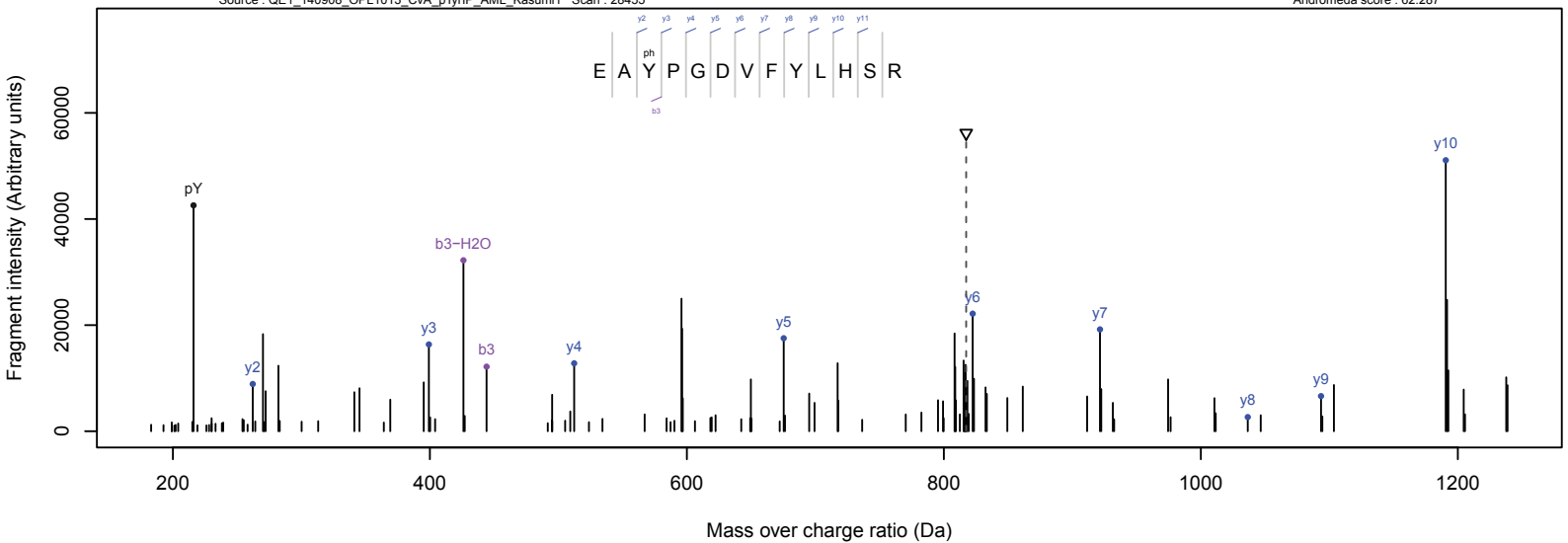
Andromeda score : 62.338



**EAY(ph)PGDVFYLHSR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 28455

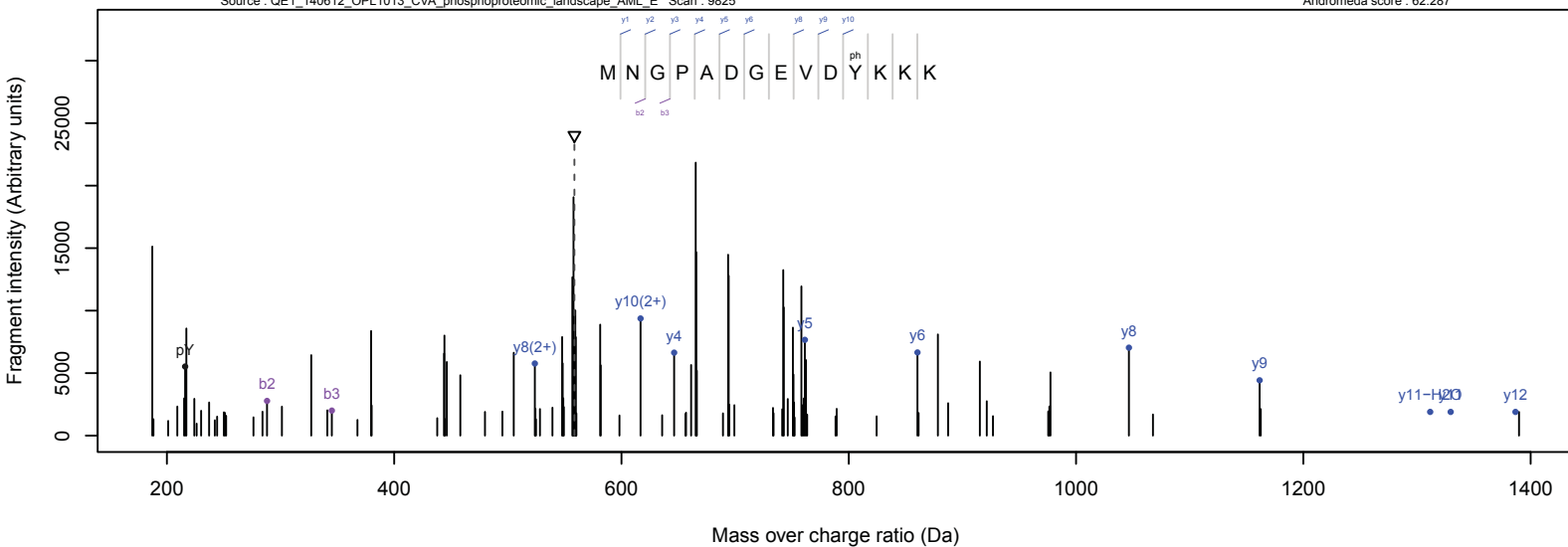
Andromeda score : 62.287



**(ac)MNGPADGEVDY(ph)KKK-3+**

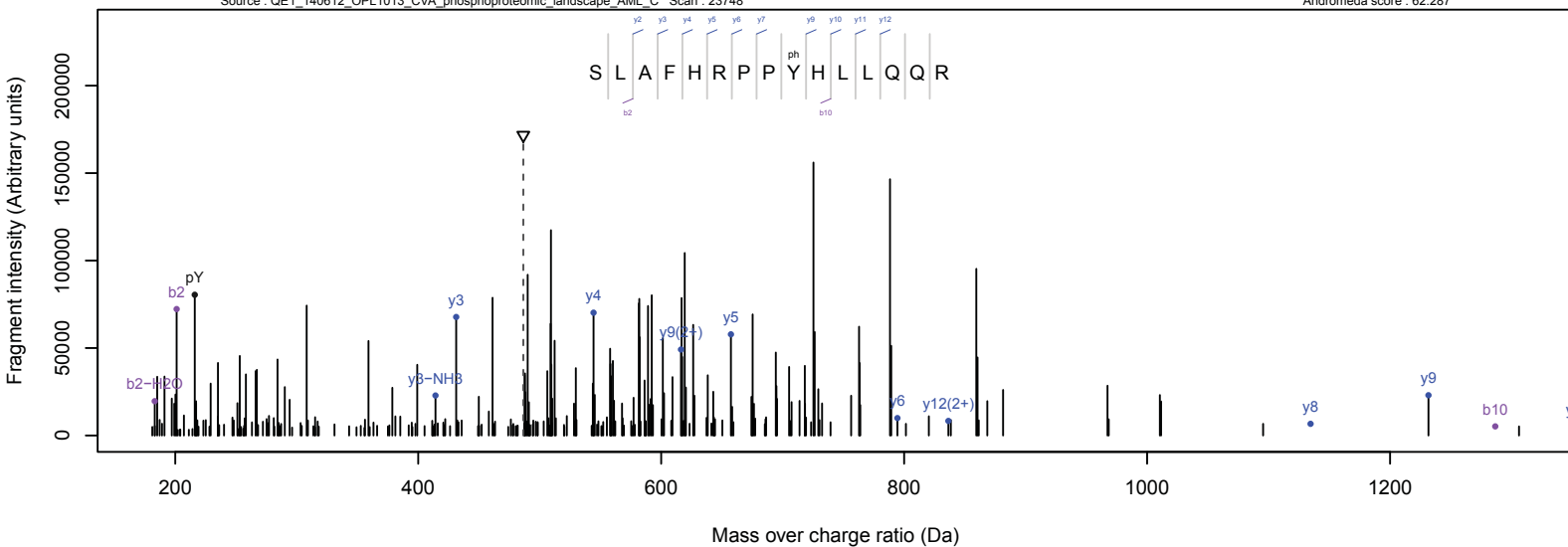
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 9825

Andromeda score : 62.287

**SLAFHRPPY(ph)HLLQQR-4+**

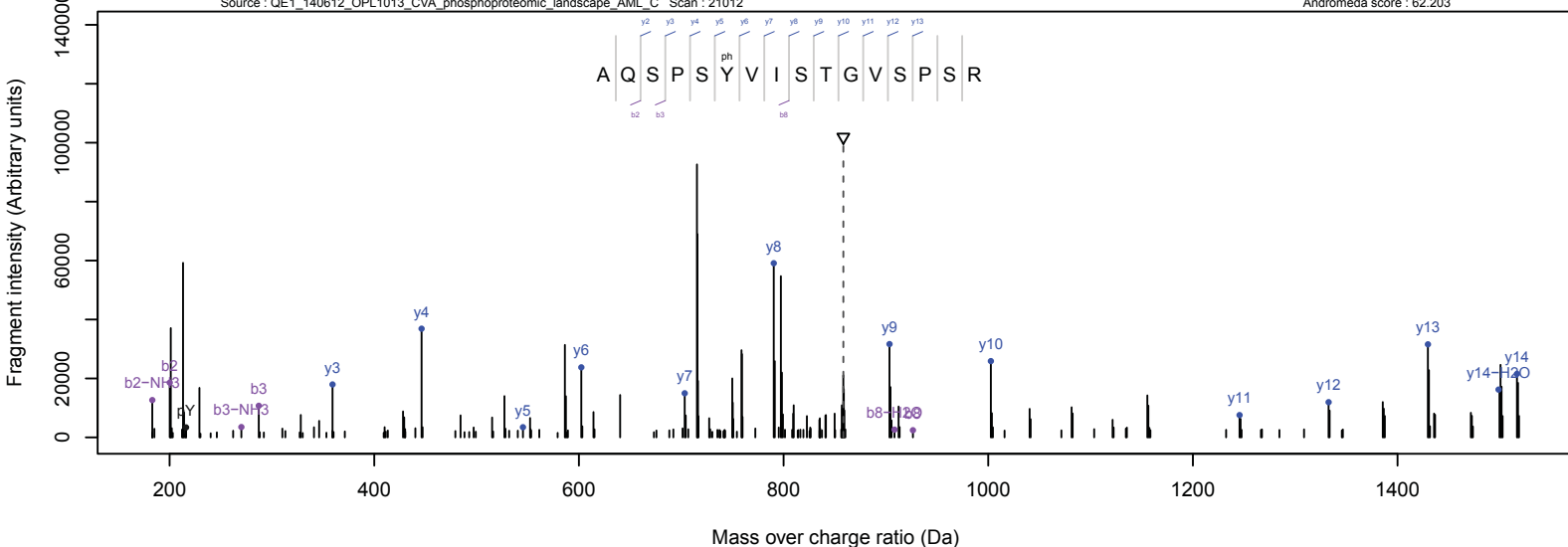
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23748

Andromeda score : 62.287

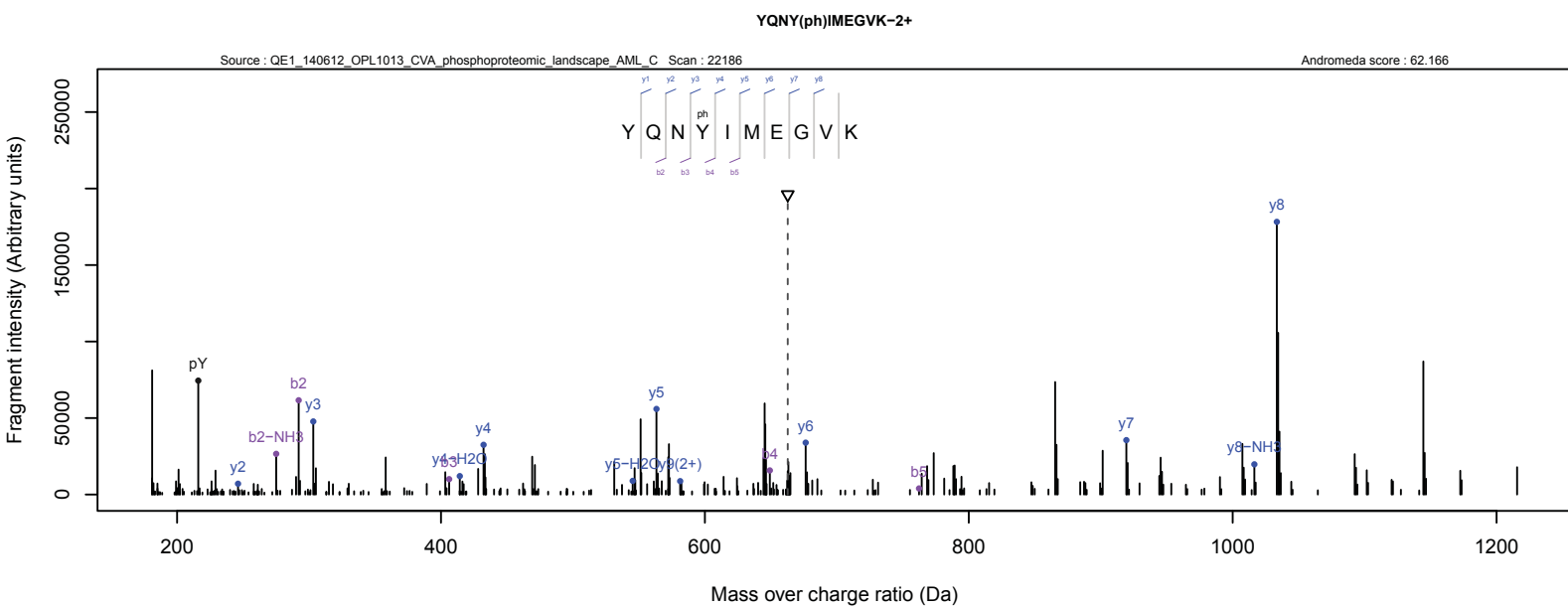
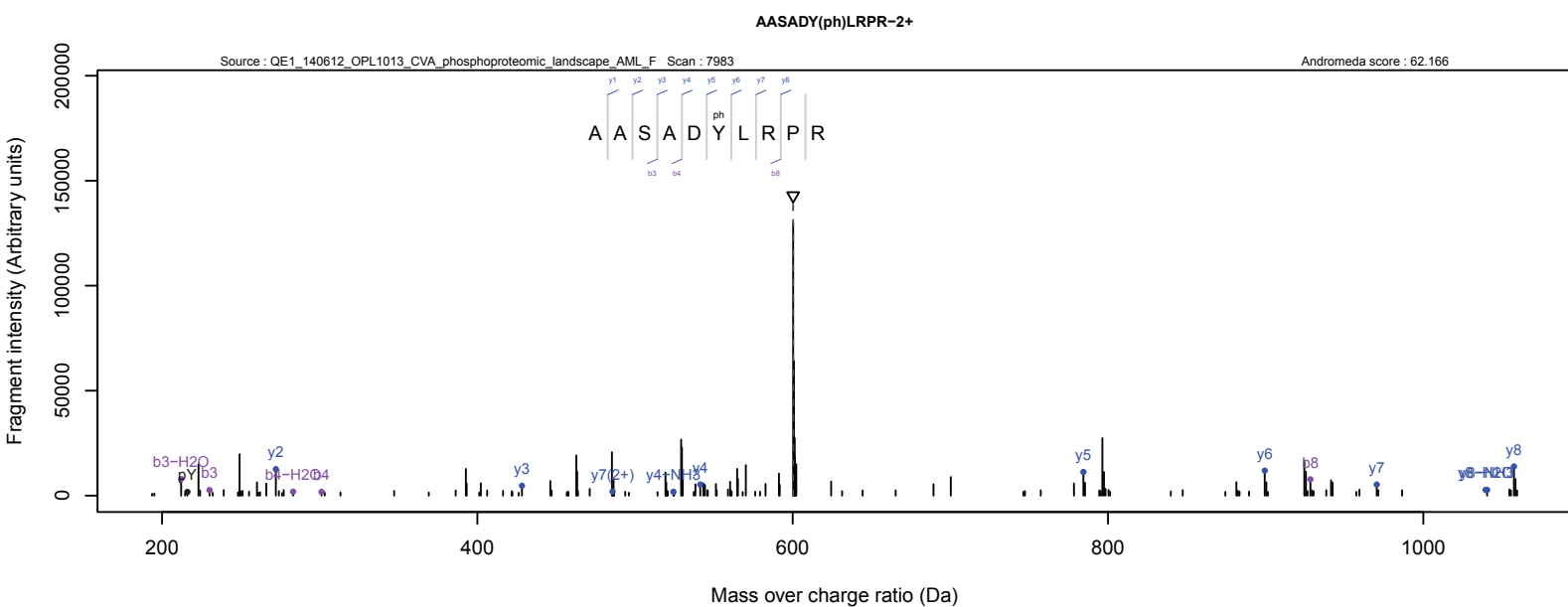
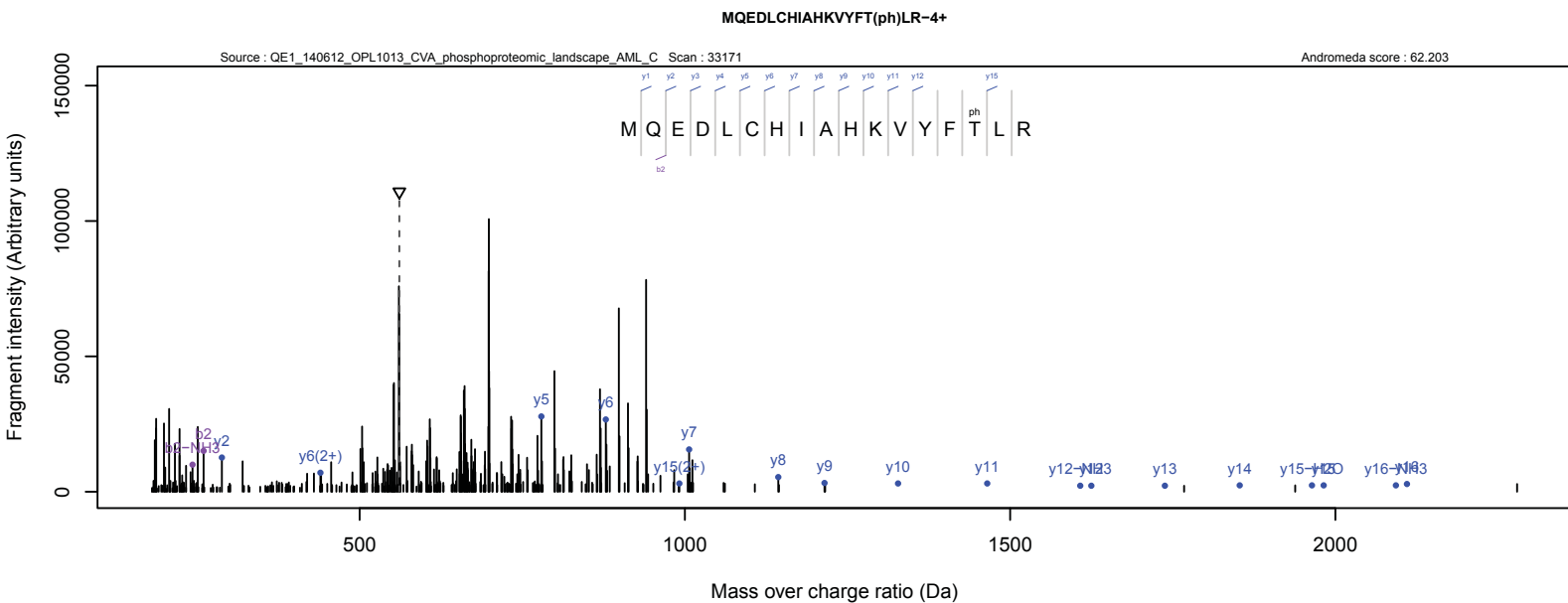
**AQSPSY(ph)VISTGVSPSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21012

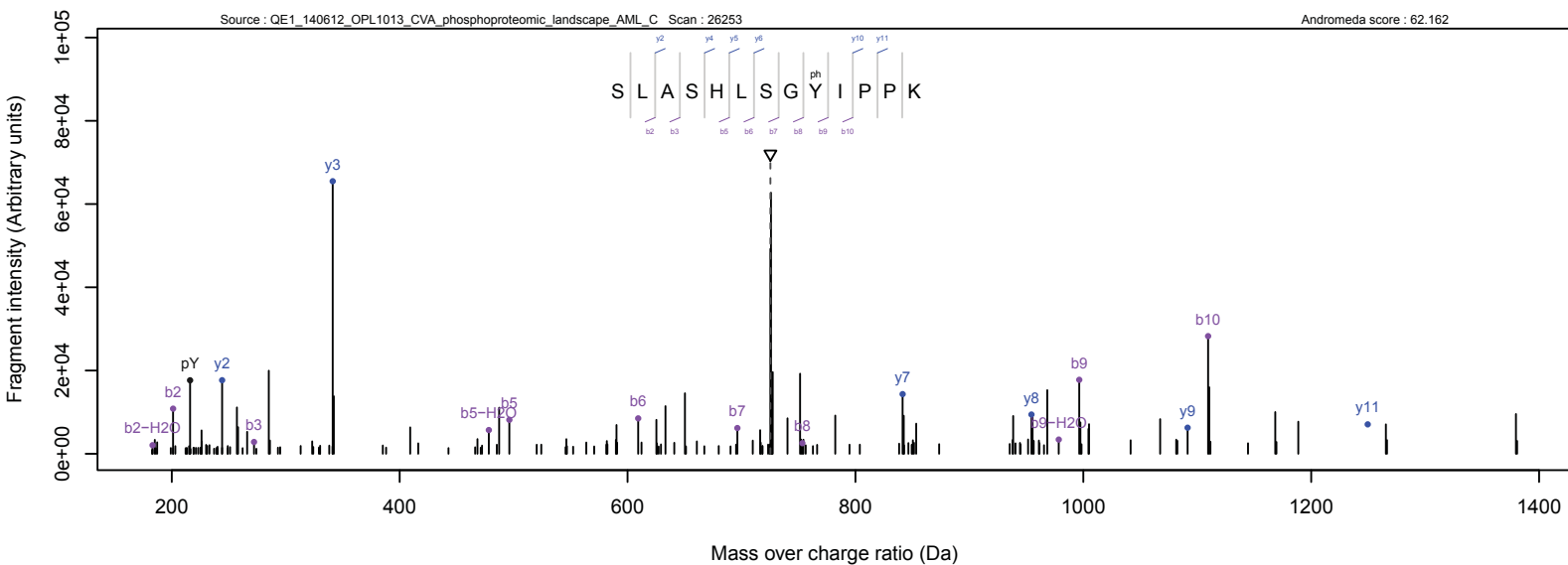
Andromeda score : 62.203



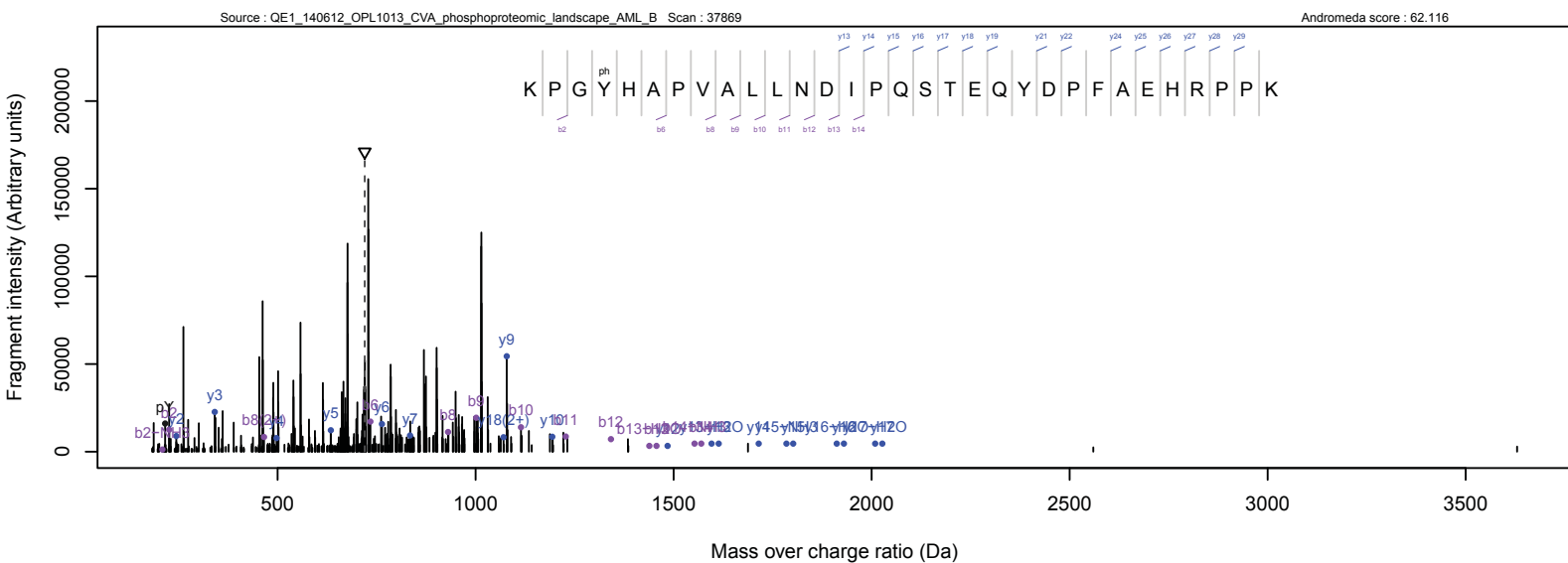




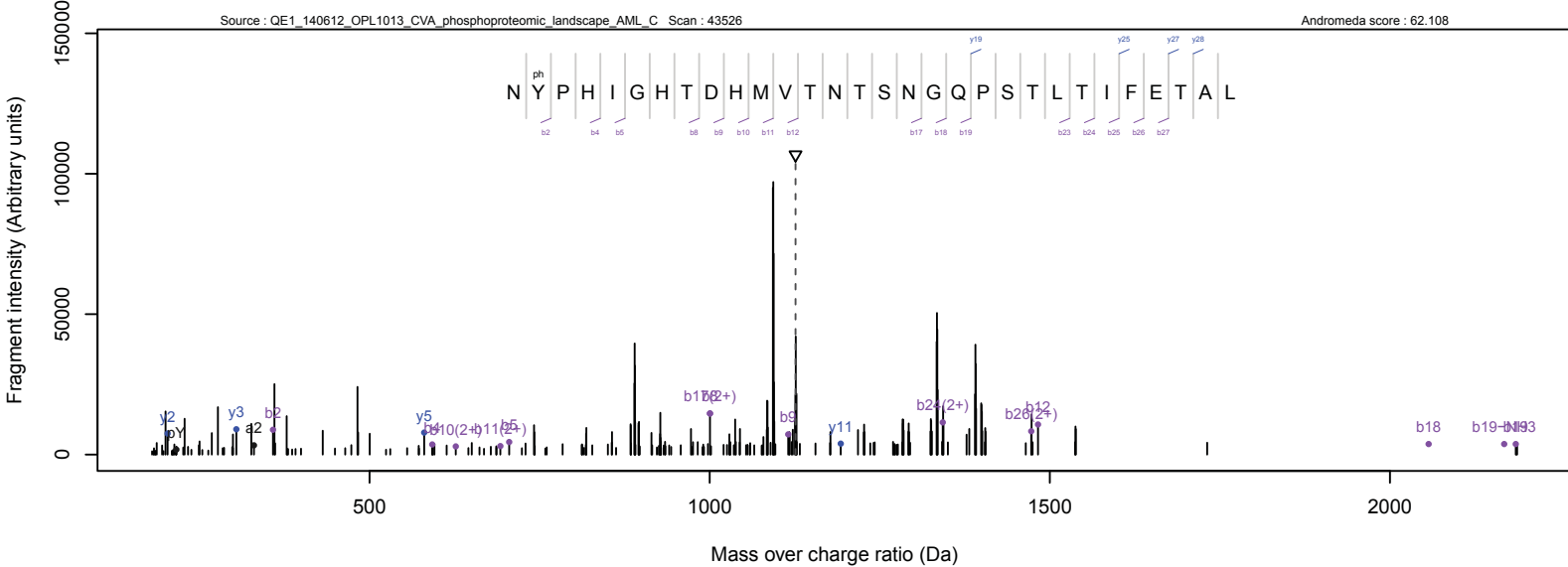
SLASHLSGY(ph)IPPK-2+



KPGY(ph)HAPVALLNDIPQSTEQYDPPFAEHRPPK-5+



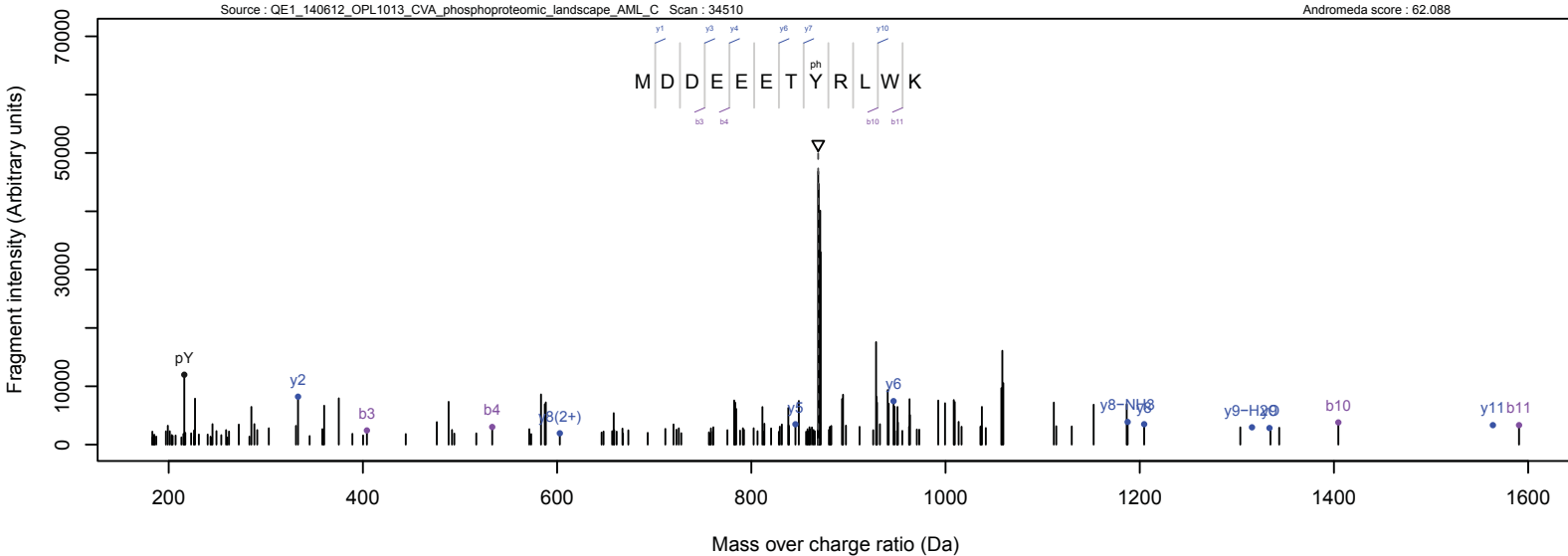
NY(ph)PHIGHTDHMTNTSNGQPSTLTIFETAL-3+



(ac)MDDEEETY(ph)RLWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34510

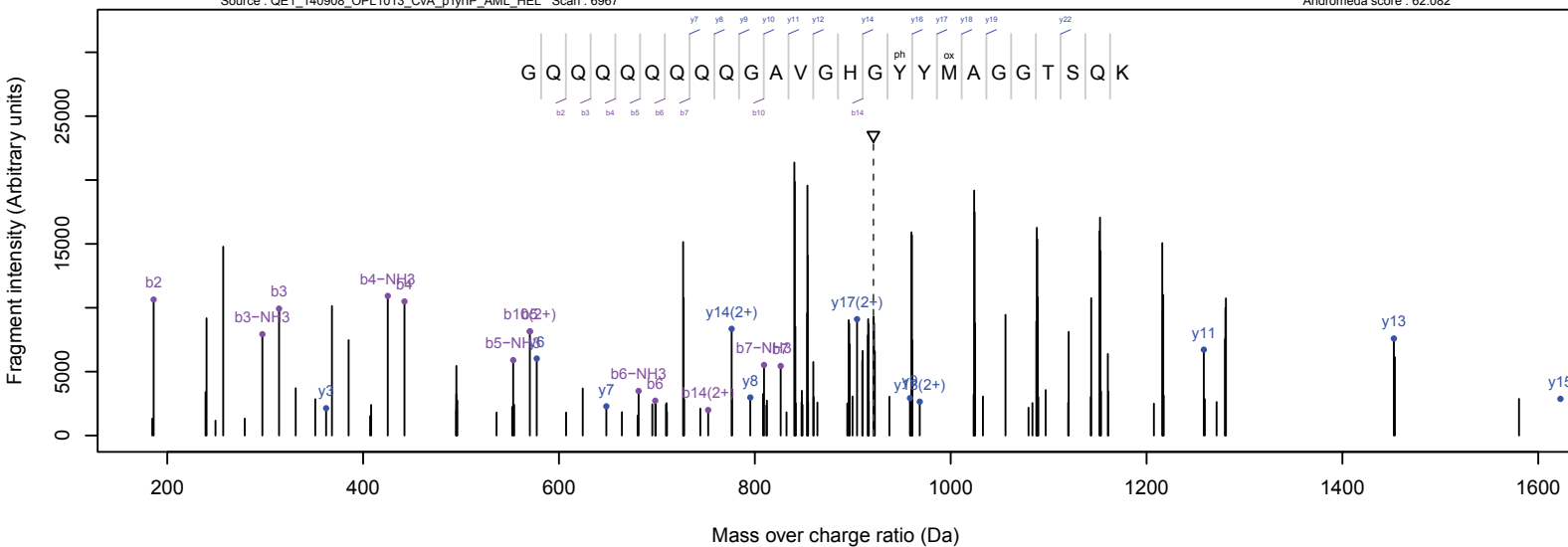
Andromeda score : 62.088



GQQQQQQQGVGHGY(ph)YM(ox)AGGTSQK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 6967

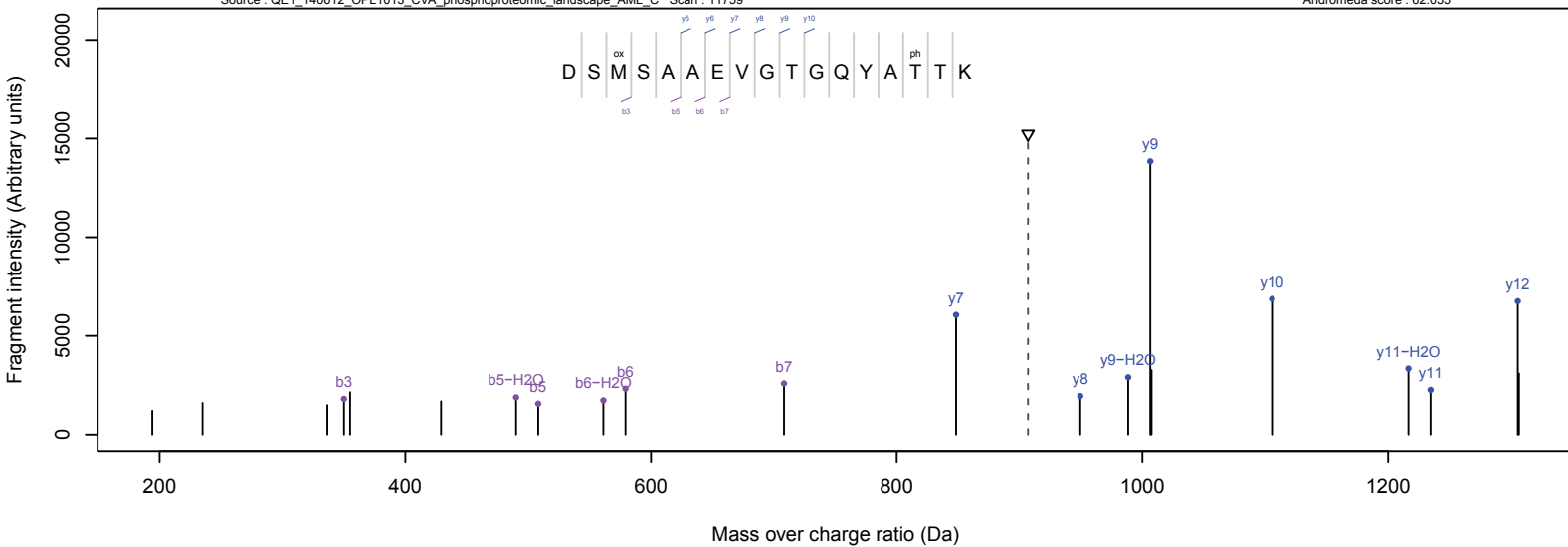
Andromeda score : 62.082



DSM(ox)SAAEVGTGQYAT(ph)TK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11759

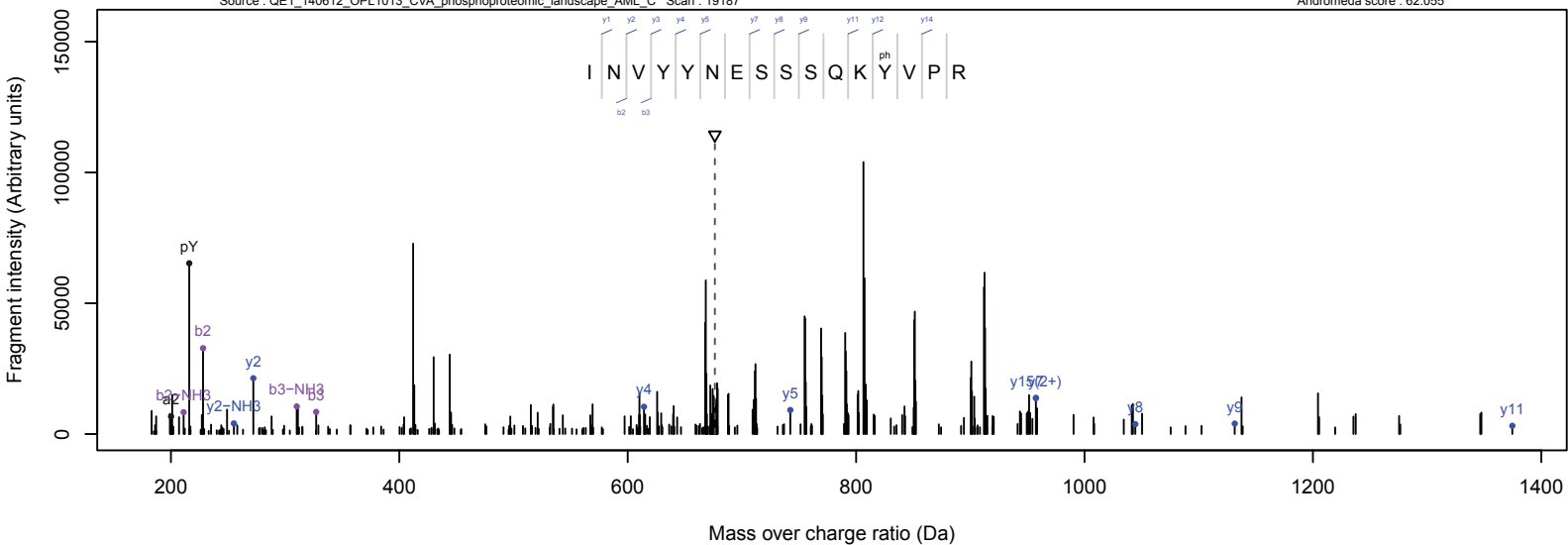
Andromeda score : 62.055



## INVYNESSQKY(ph)VPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19187

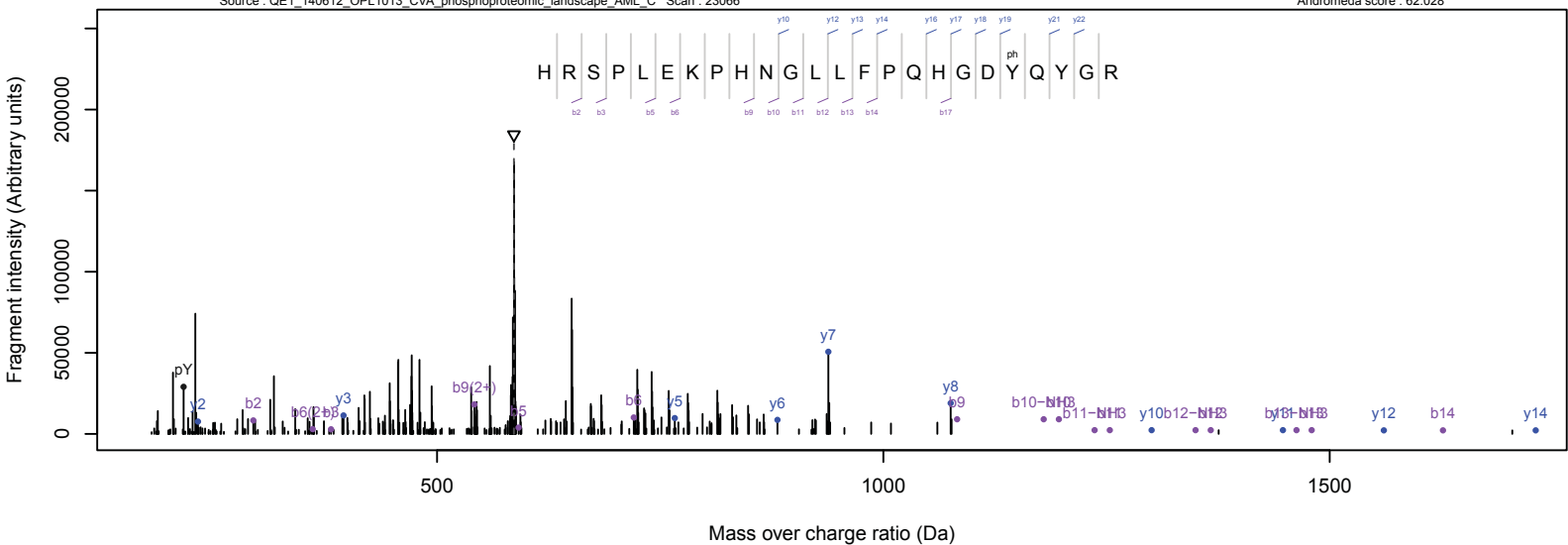
Andromeda score: 62.055



## HRSPLEKPHNGLLPQHGDY(ph)QYGR-5+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23066

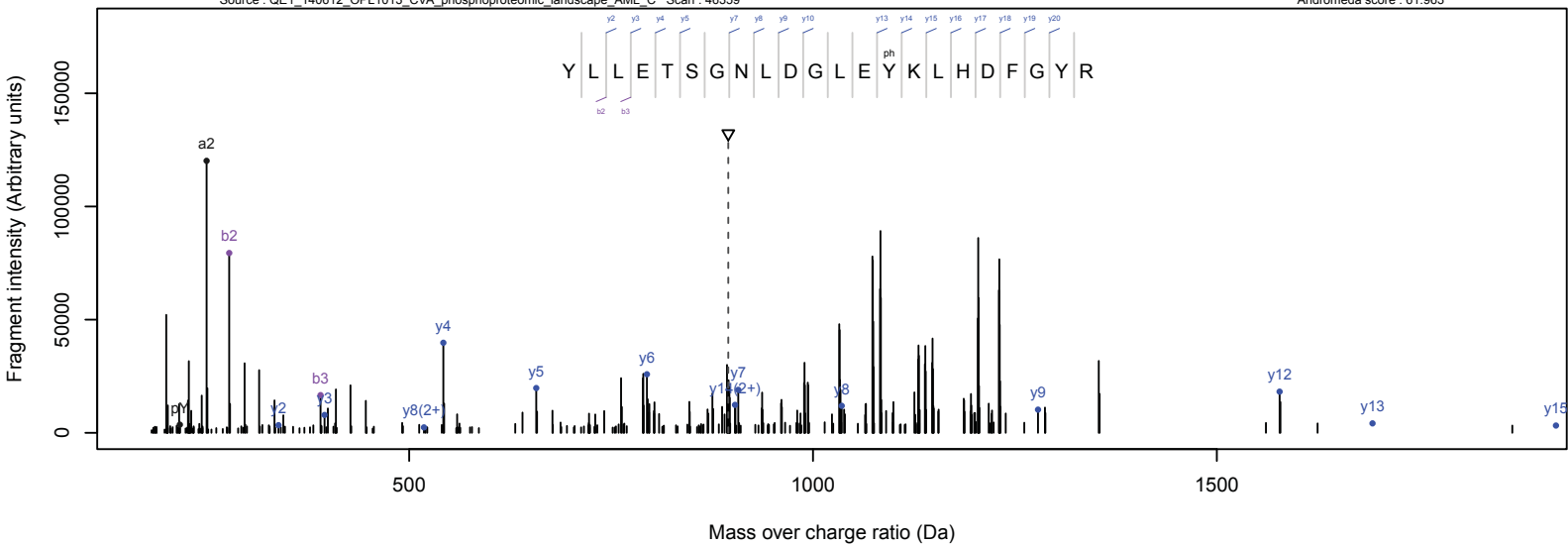
Andromeda score: 62.028



## YLLETSGNLDGLEYPK(LHDFGYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 46359

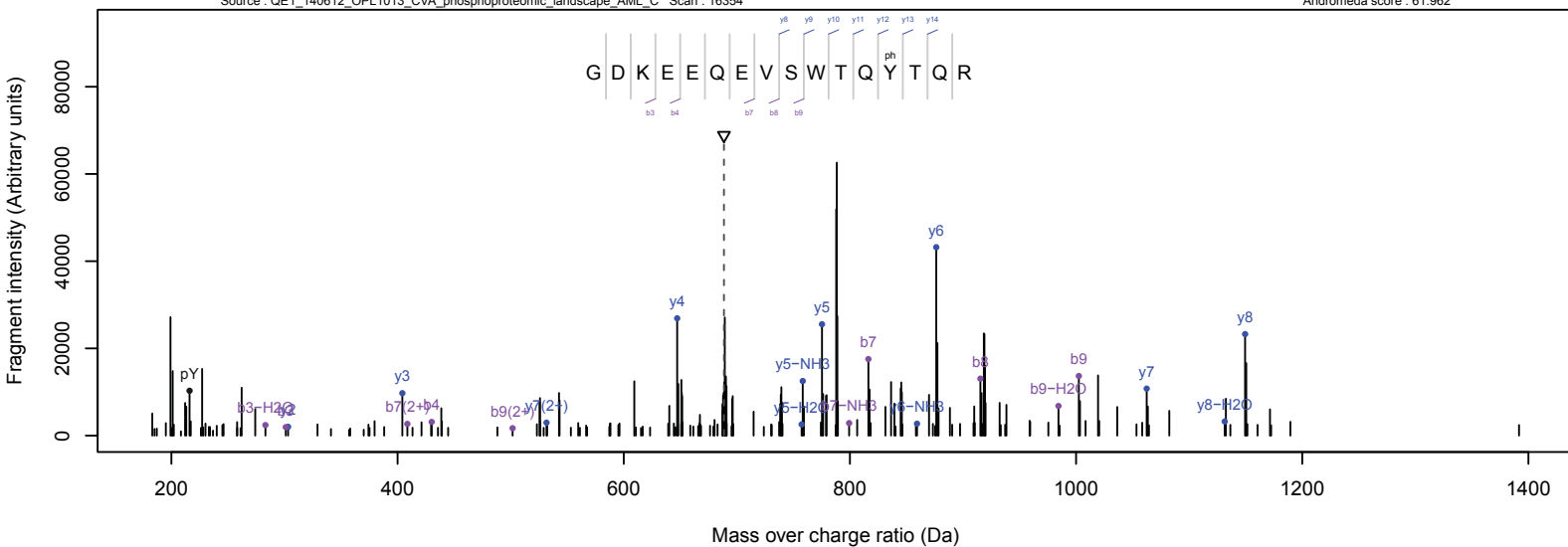
Andromeda score: 61.963



## GDKEEQVSWTQY(ph)TQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16354

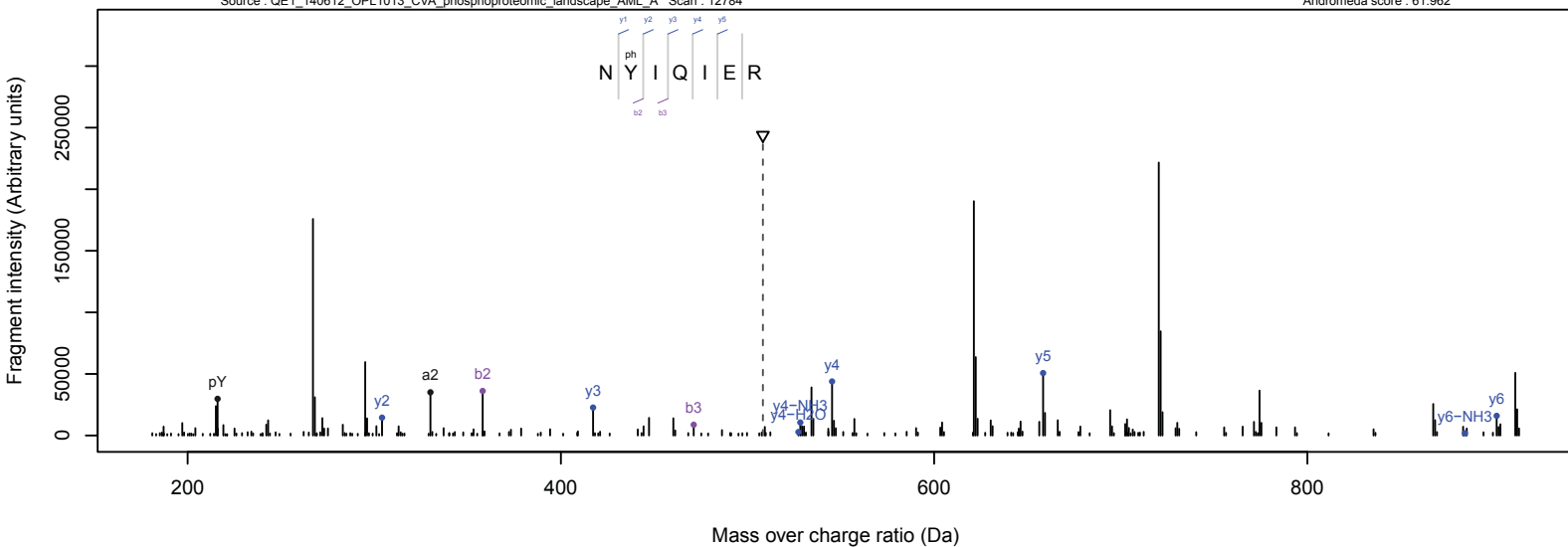
Andromeda score : 61.962



## NY(ph)IQIER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12784

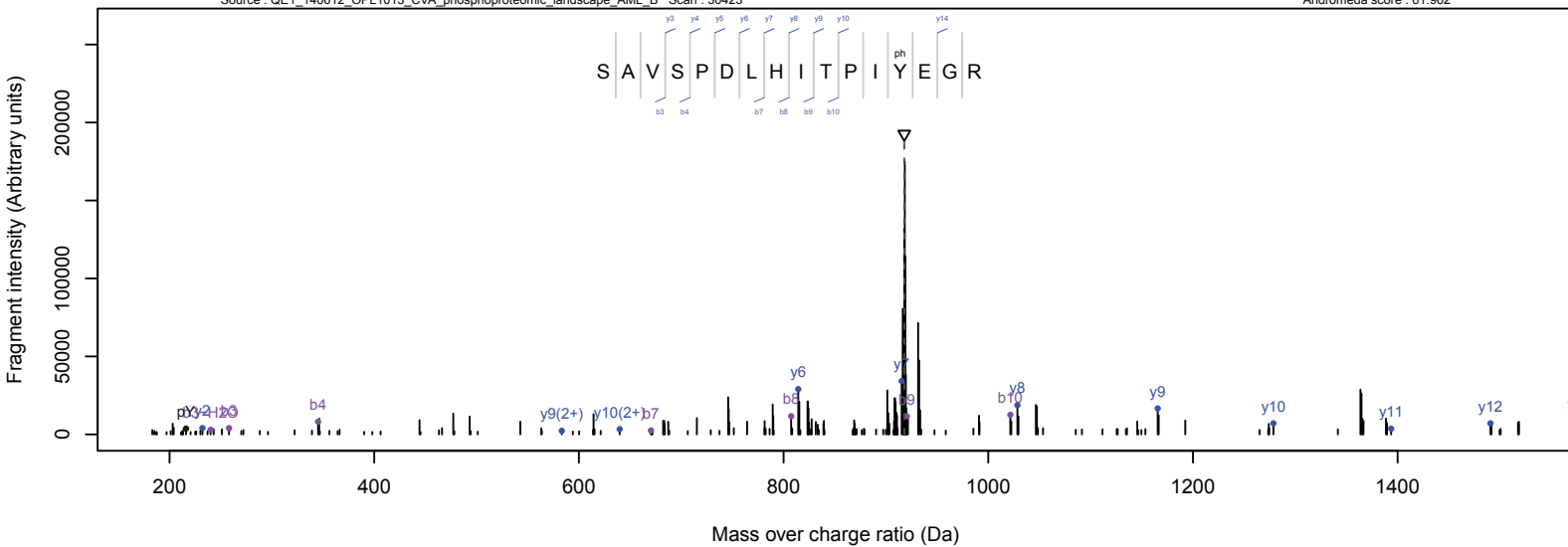
Andromeda score : 61.962



## SAVSPDLHITPIY(ph)EGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30423

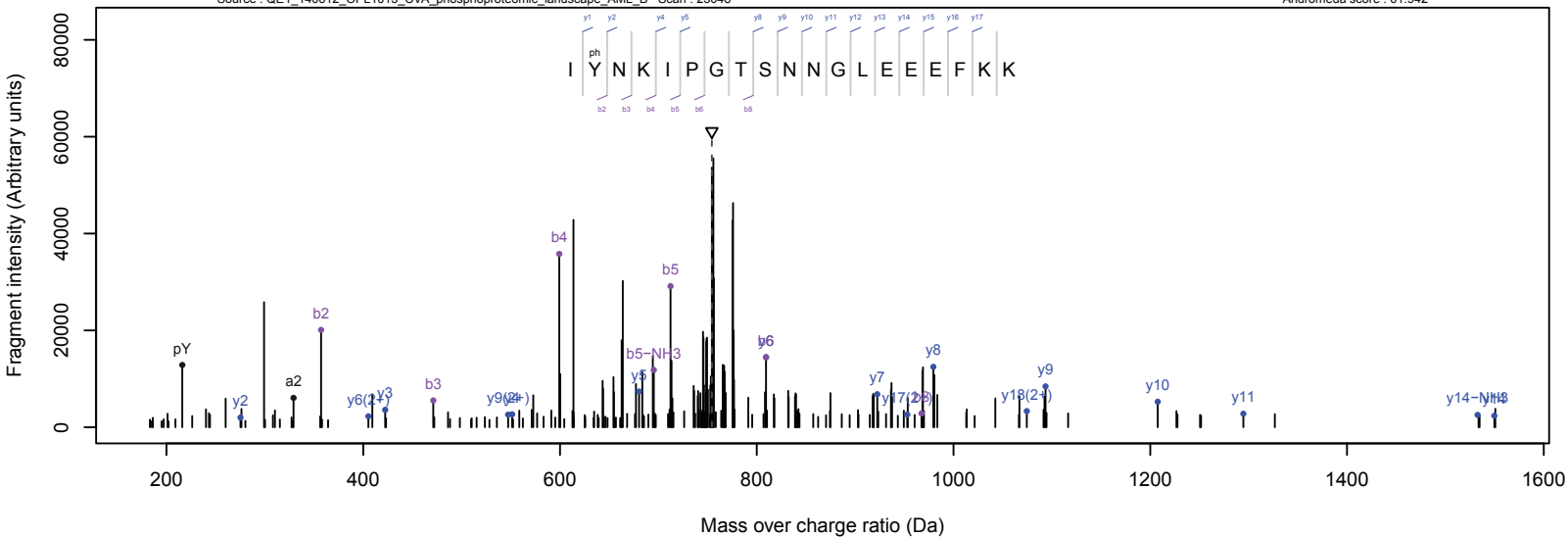
Andromeda score : 61.962



## IY(ph)NKIPGTSNNGLEEFKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23048

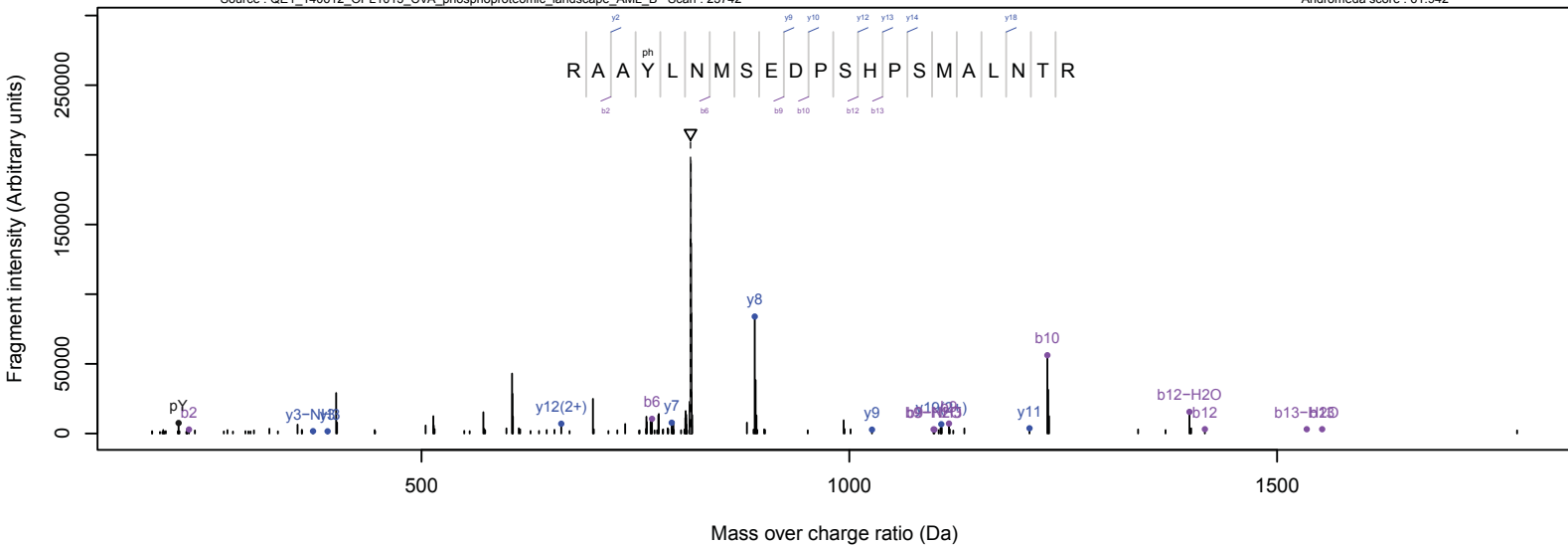
Andromeda score : 61.942



## RAAY(ph)LNMSSEDPSPHPSMALNTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23742

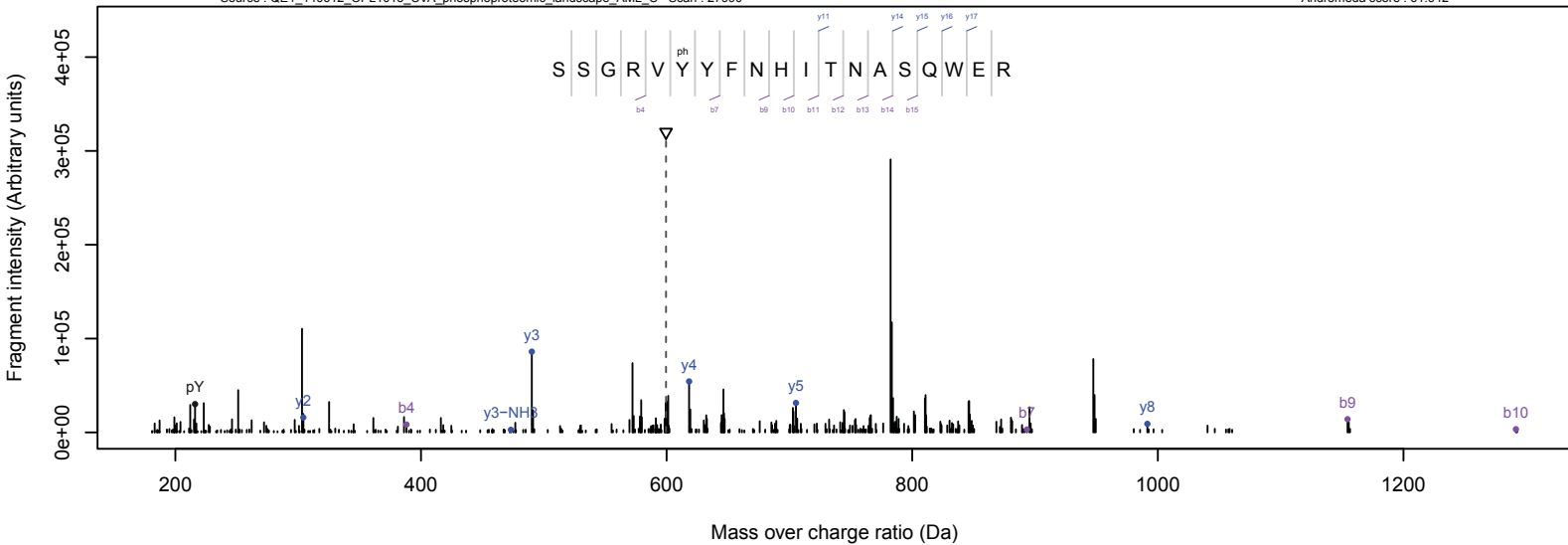
Andromeda score : 61.942



## SSGRVY(ph)YFNHITNASQWER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27990

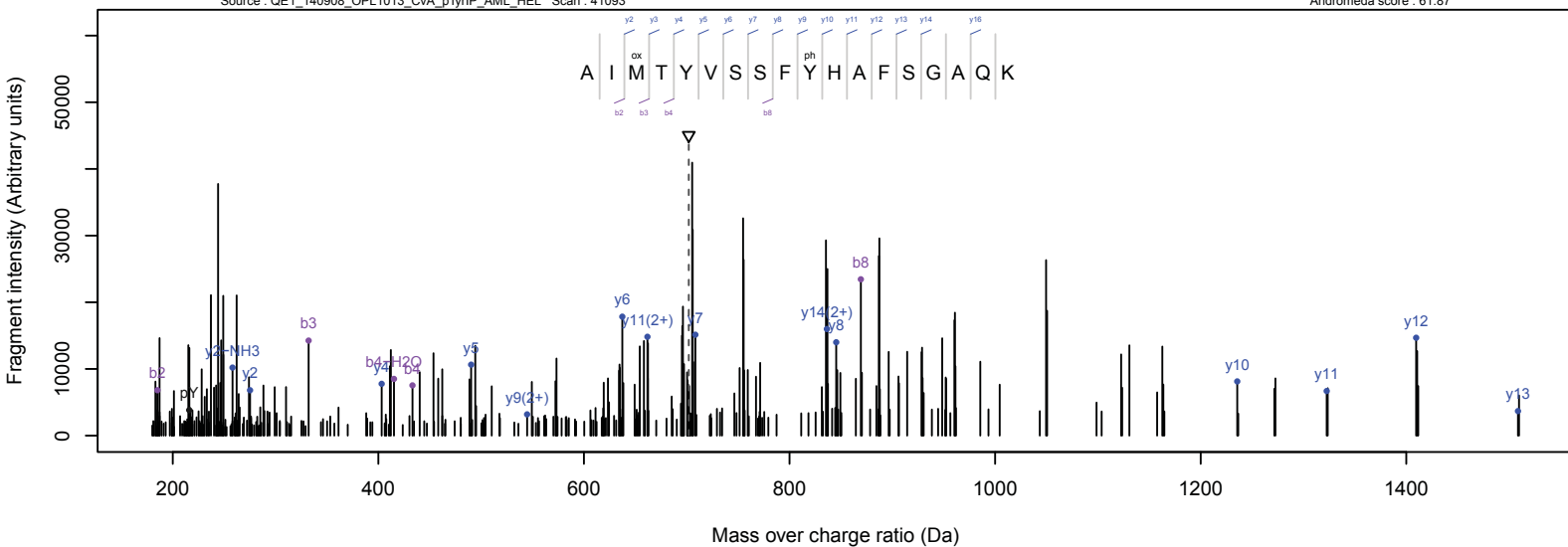
Andromeda score : 61.942



## AIM(ox)TYVSSFY(ph)HAFSGAQK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 41093

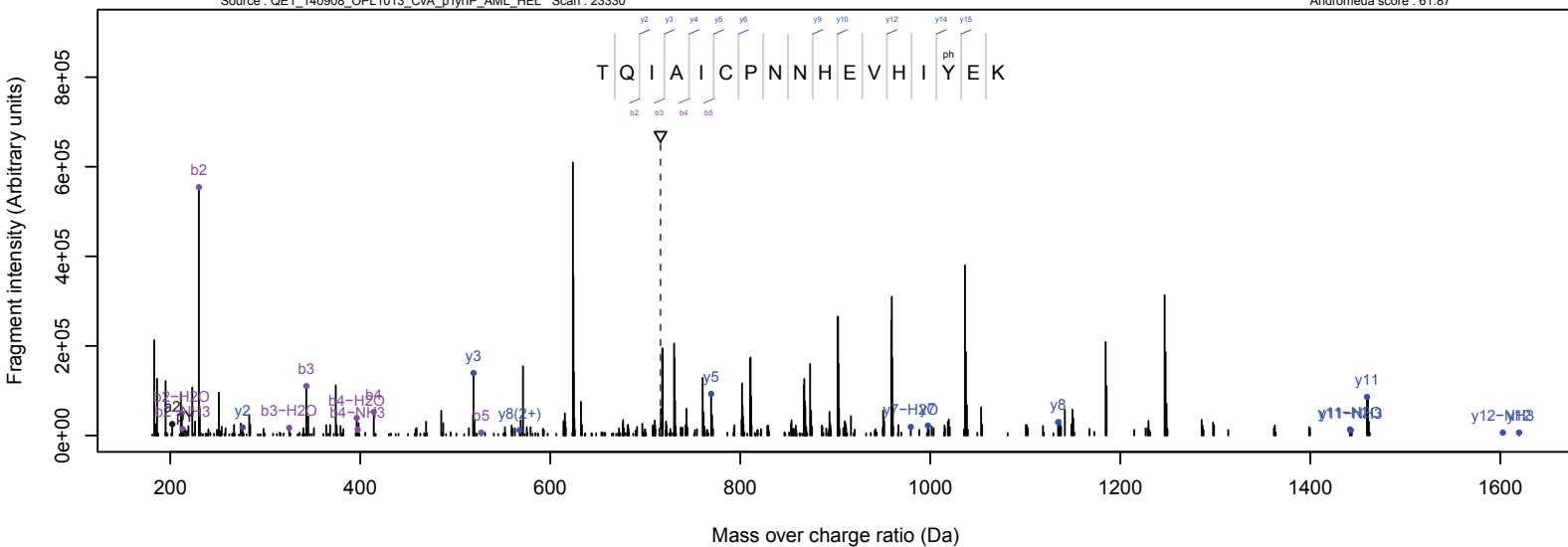
Andromeda score : 61.87



## TQIAICPNNHEVHIYEK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23330

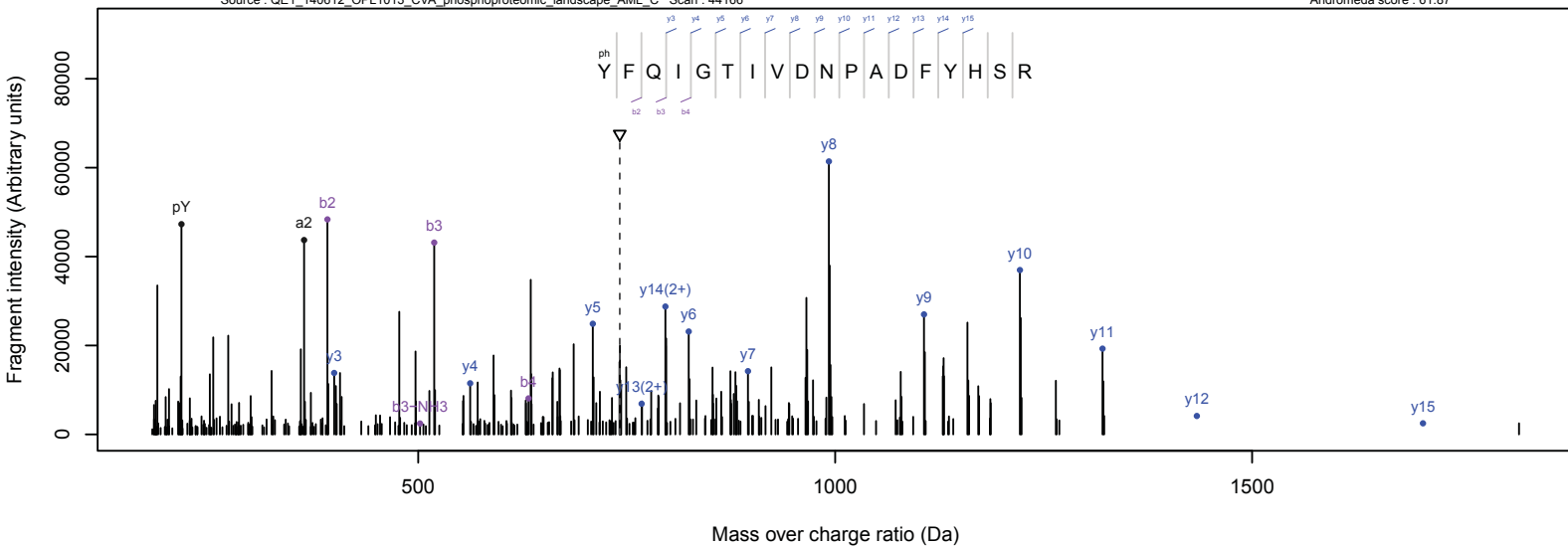
Andromeda score : 61.87



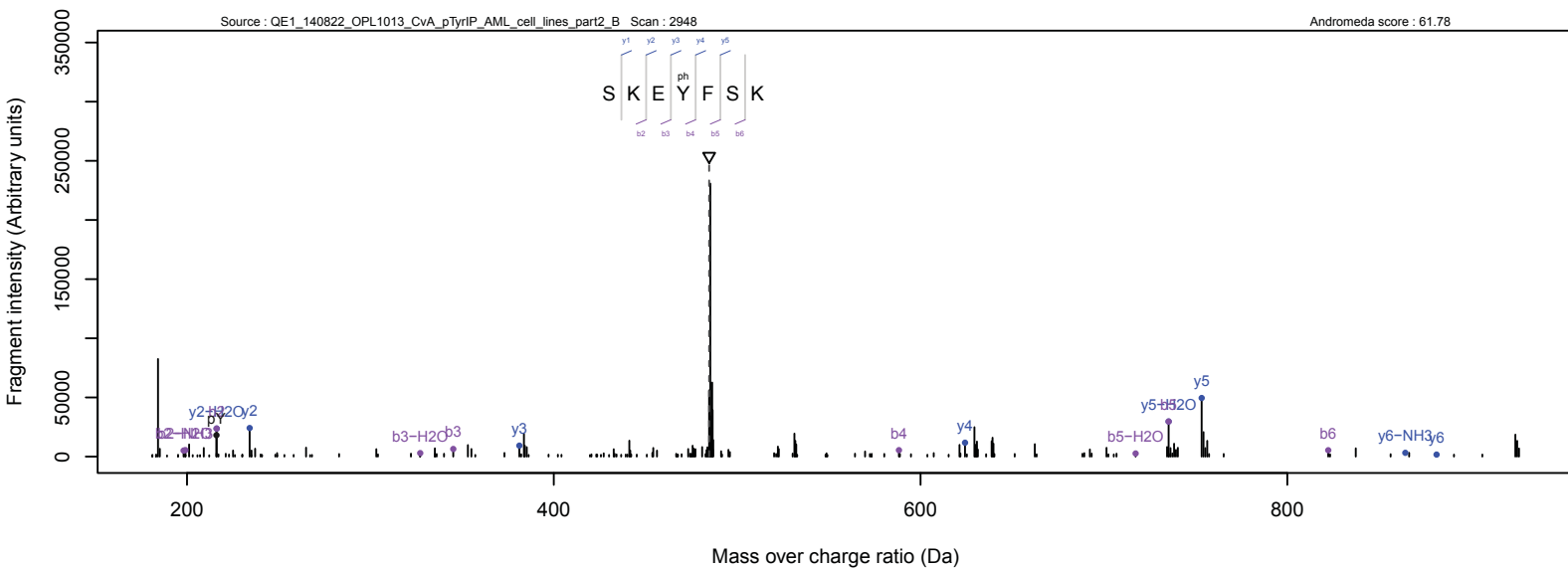
## Y(ph)FQIGTIVDNPADFYHSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44166

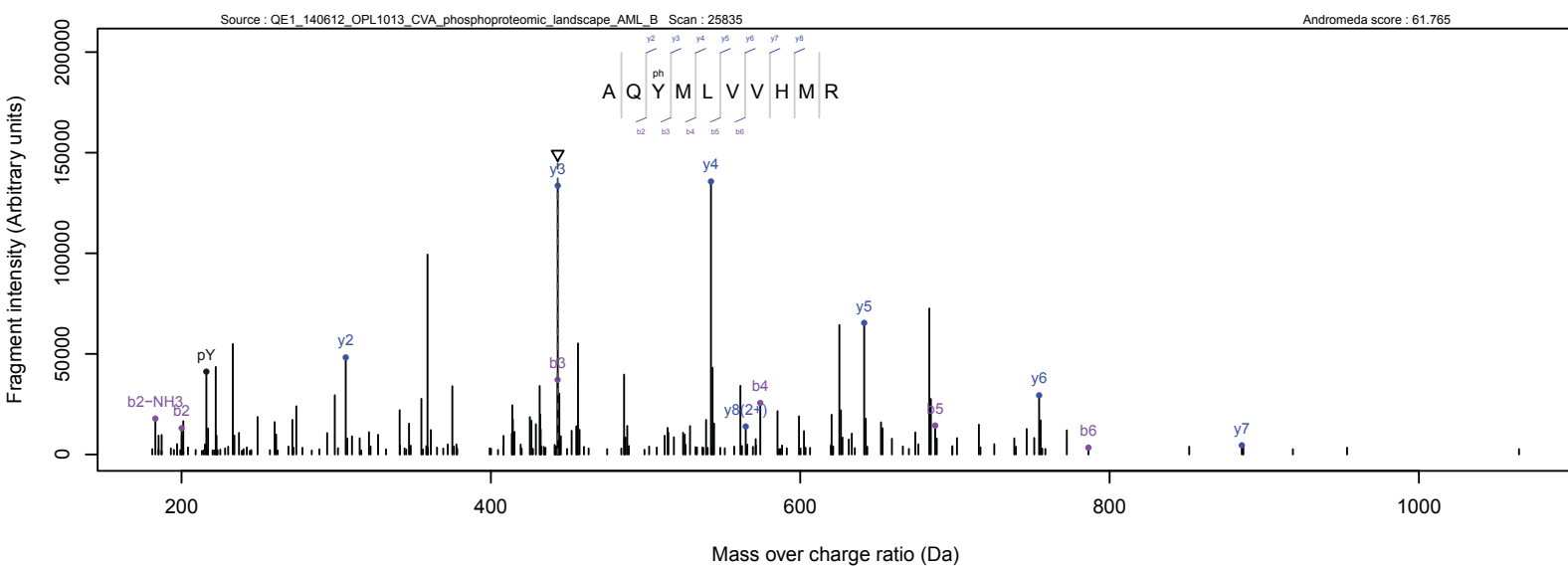
Andromeda score : 61.87



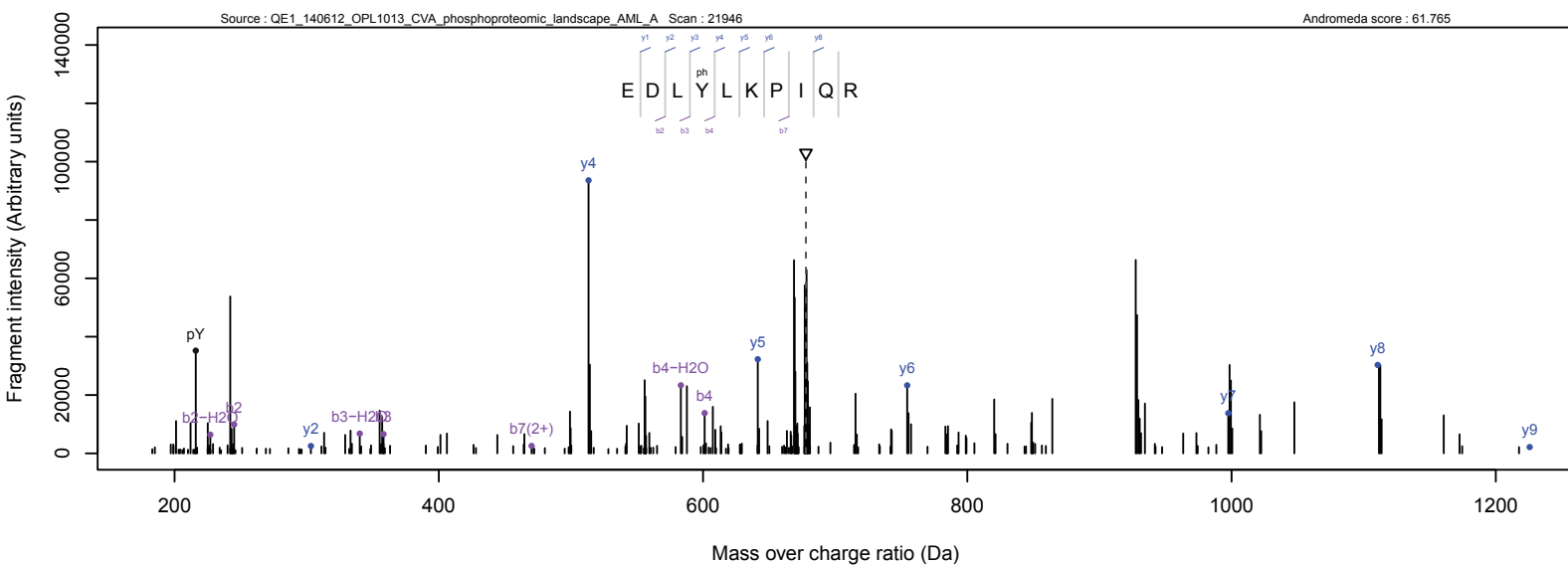
## SKEY(ph)FSK-2+



## AQY(ph)MLVVHMR-3+



## EDLY(ph)LKPIQR-2+

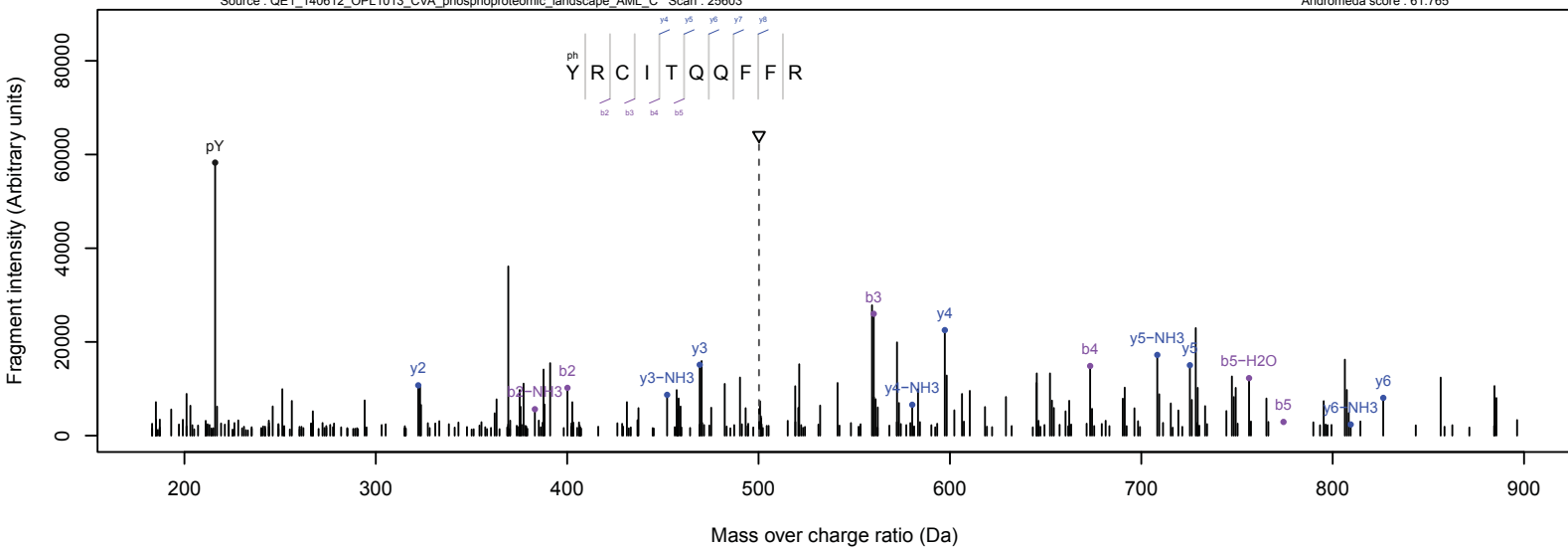




Y(ph)RCITQQFFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25603

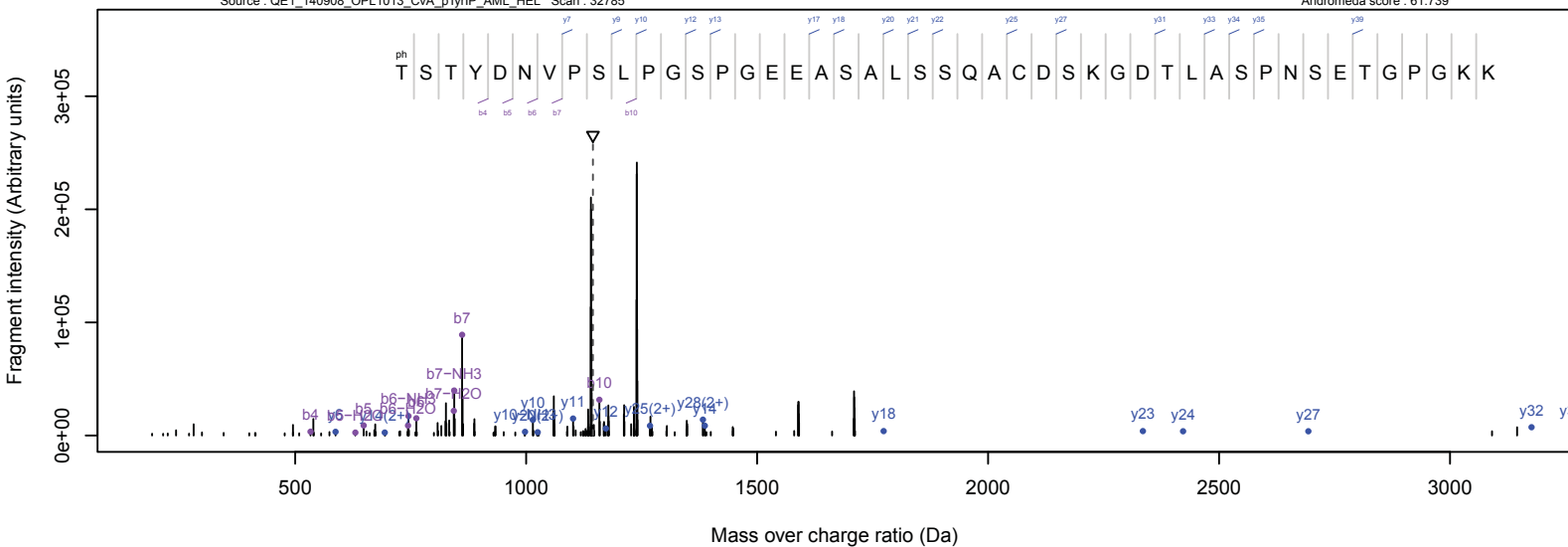
Andromeda score : 61.765



T(ph)STYDNPVSLPGSPGEEASALSSQACDSKGDTLASPNSSETGPGKK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32785

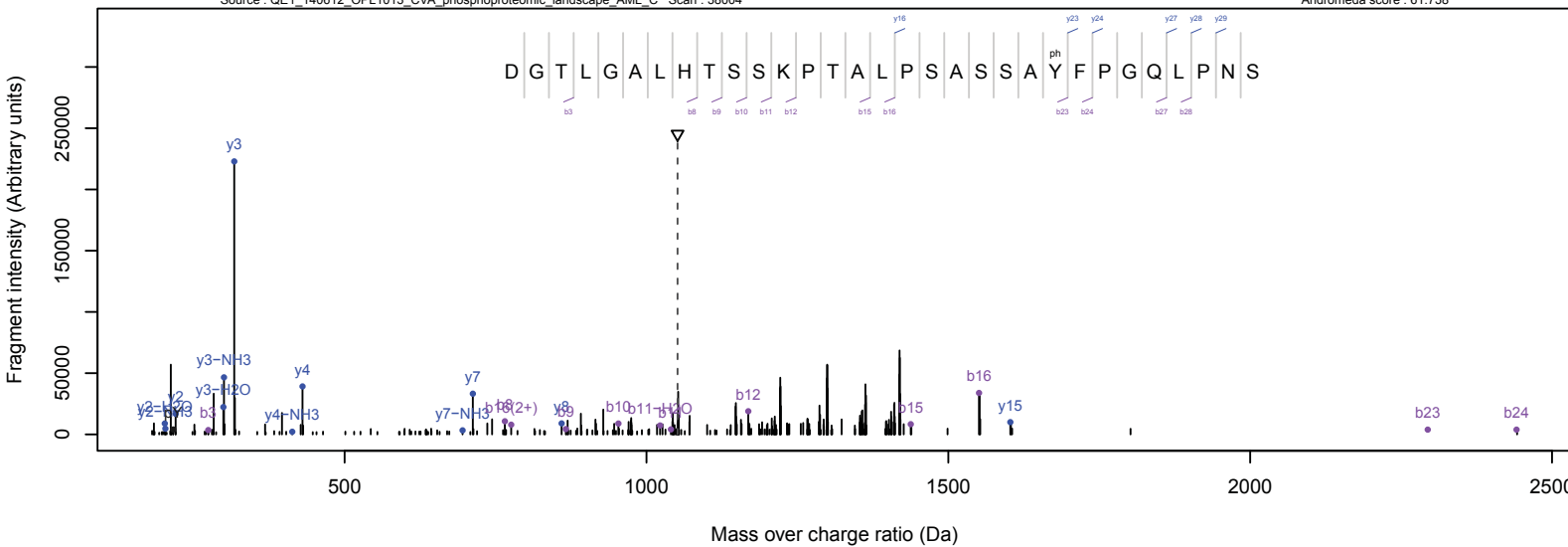
Andromeda score : 61.739



DGTLGALHTSSKPTALPSASSAY(ph)FPGQLPNS-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38004

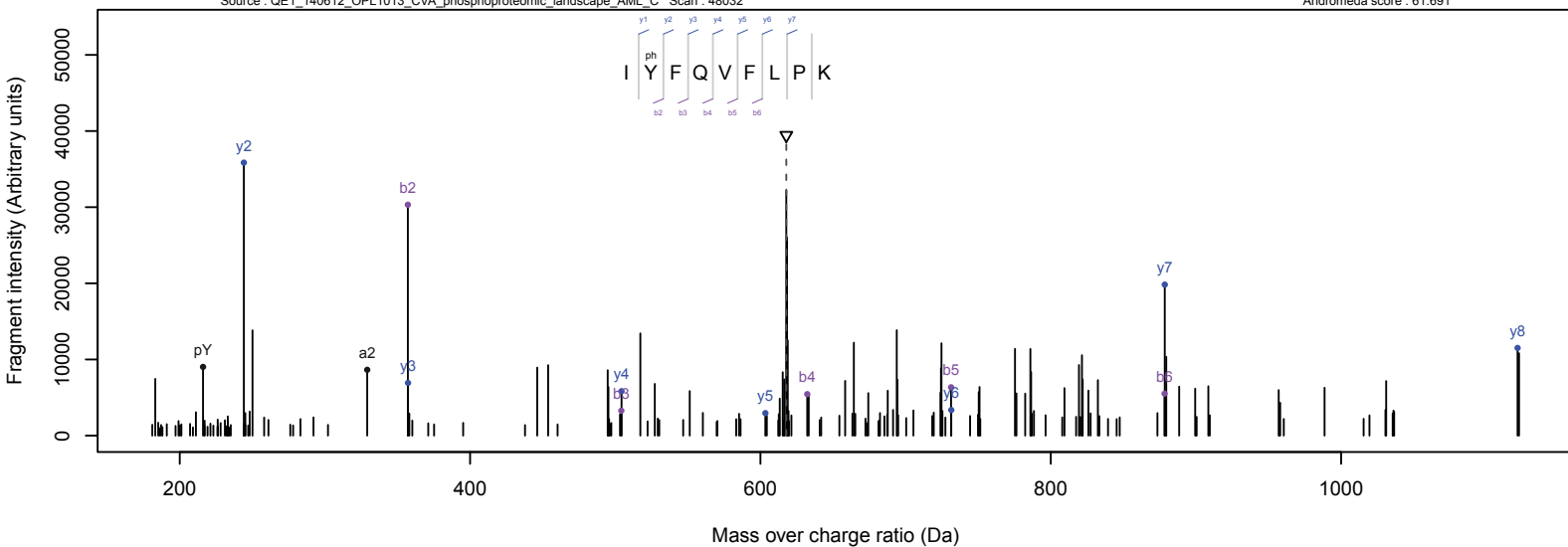
Andromeda score : 61.738



## IY(ph)FQVFLPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 48032

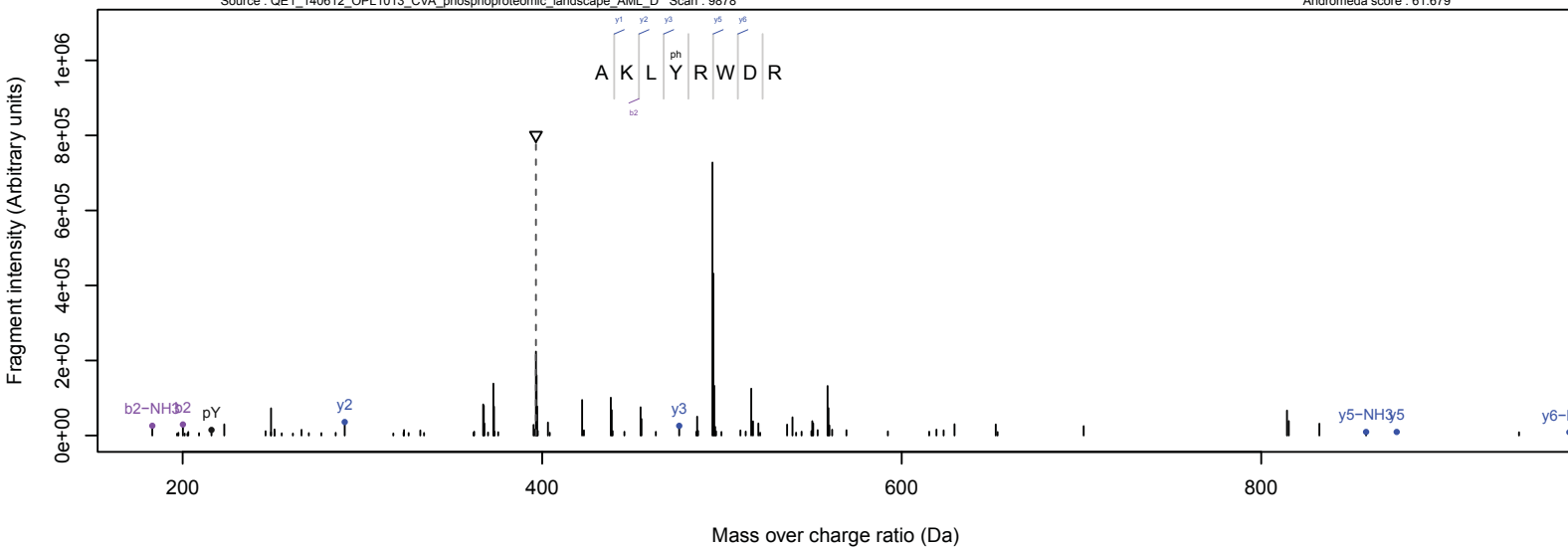
Andromeda score : 61.691



## AKLY(ph)RWDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9878

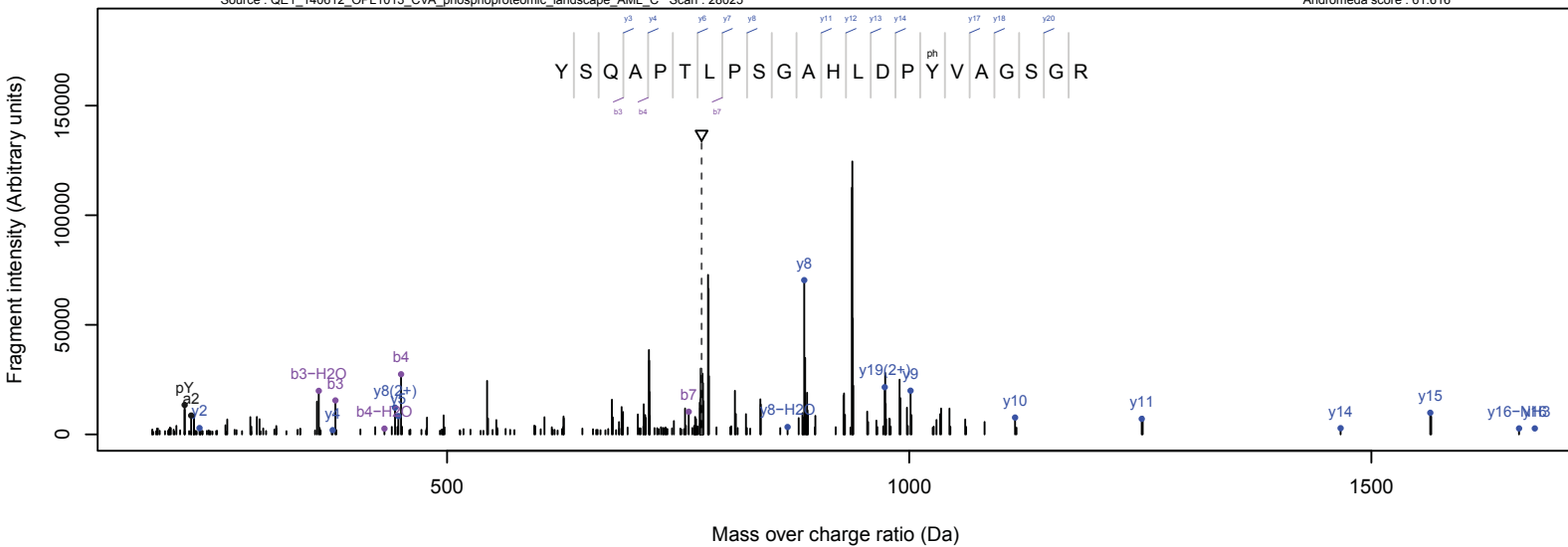
Andromeda score : 61.679



## YSQAPTLPSGAHLDPY(ph)VAGSGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28025

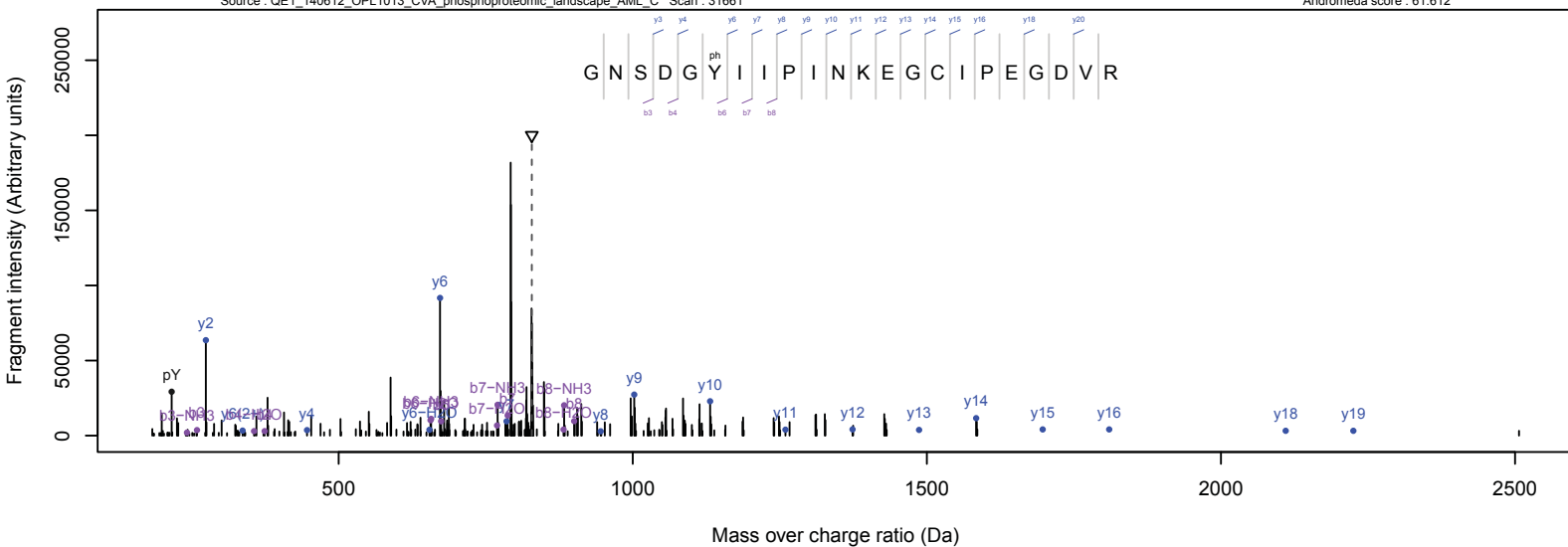
Andromeda score : 61.616



## GNSDGY(ph)IIPINKEGCIPEGDVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31661

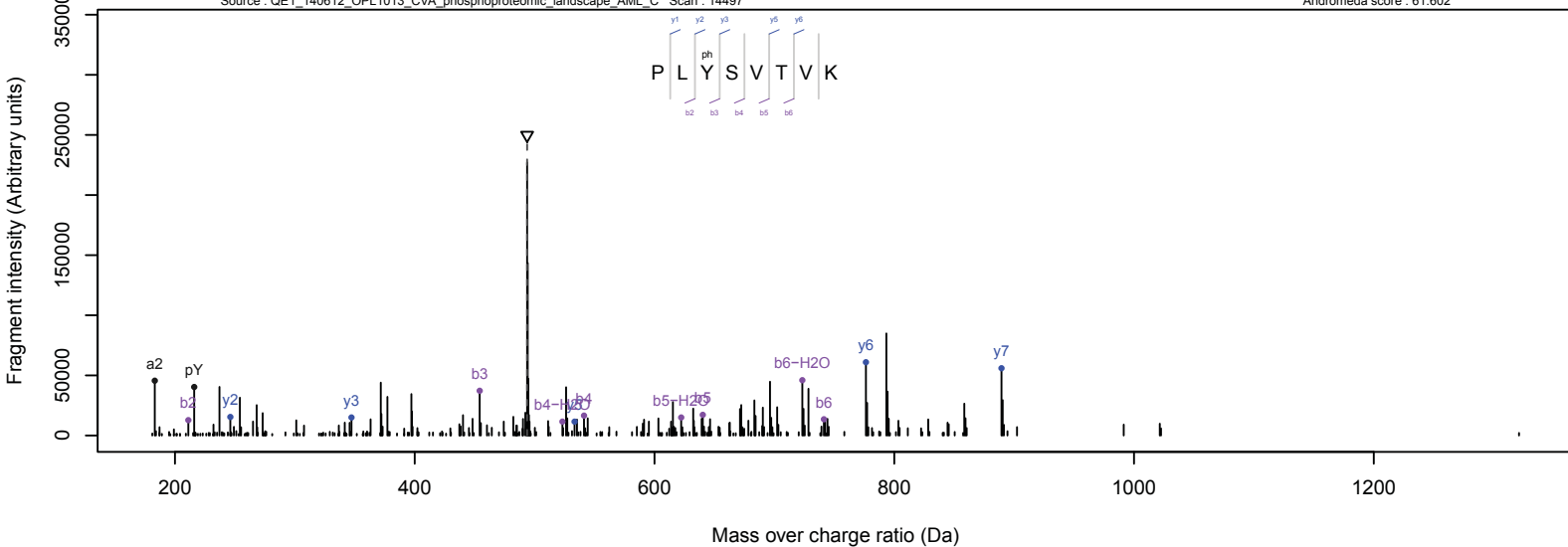
Andromeda score: 61.612



## PLY(ph)SVTVK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14497

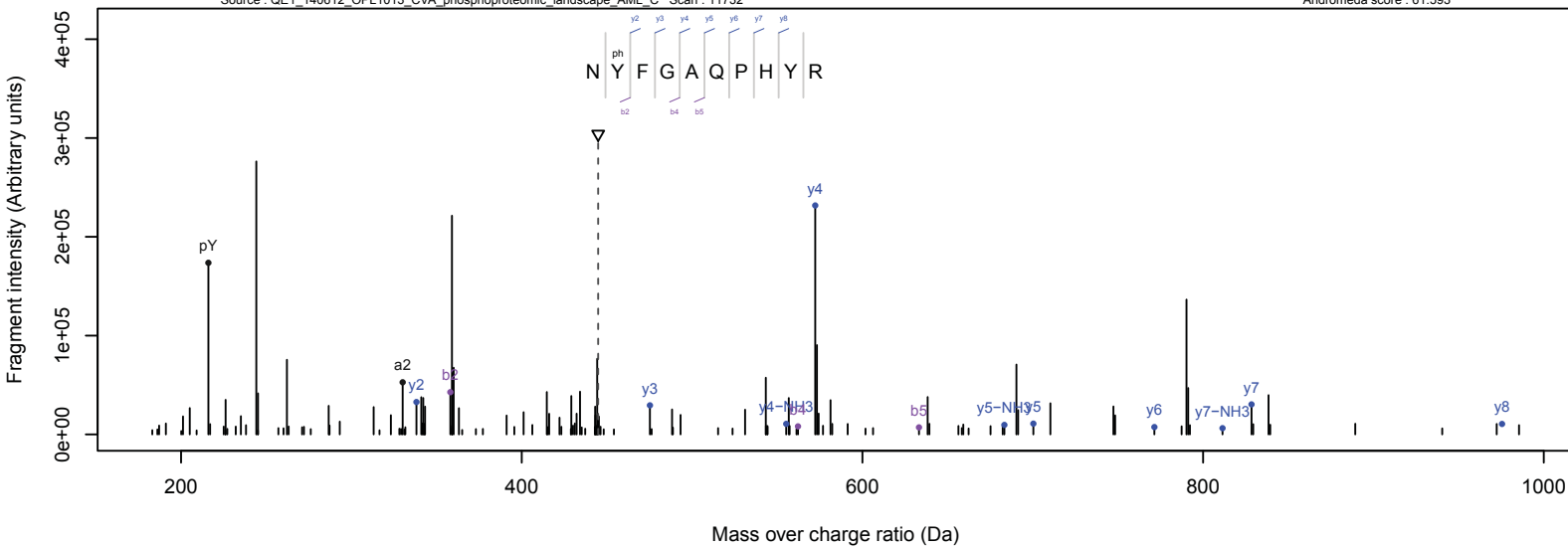
Andromeda score: 61.602



## NY(ph)FGAQPHYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11732

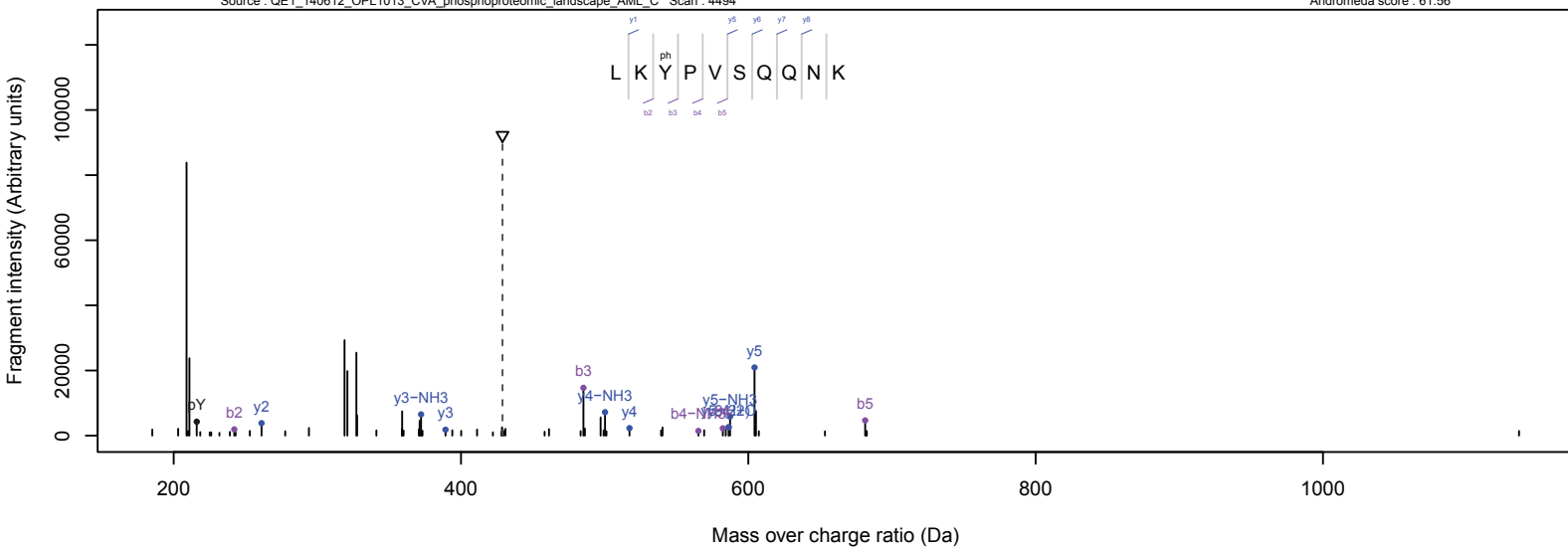
Andromeda score: 61.593



## LKY(ph)PVSQQNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 4494

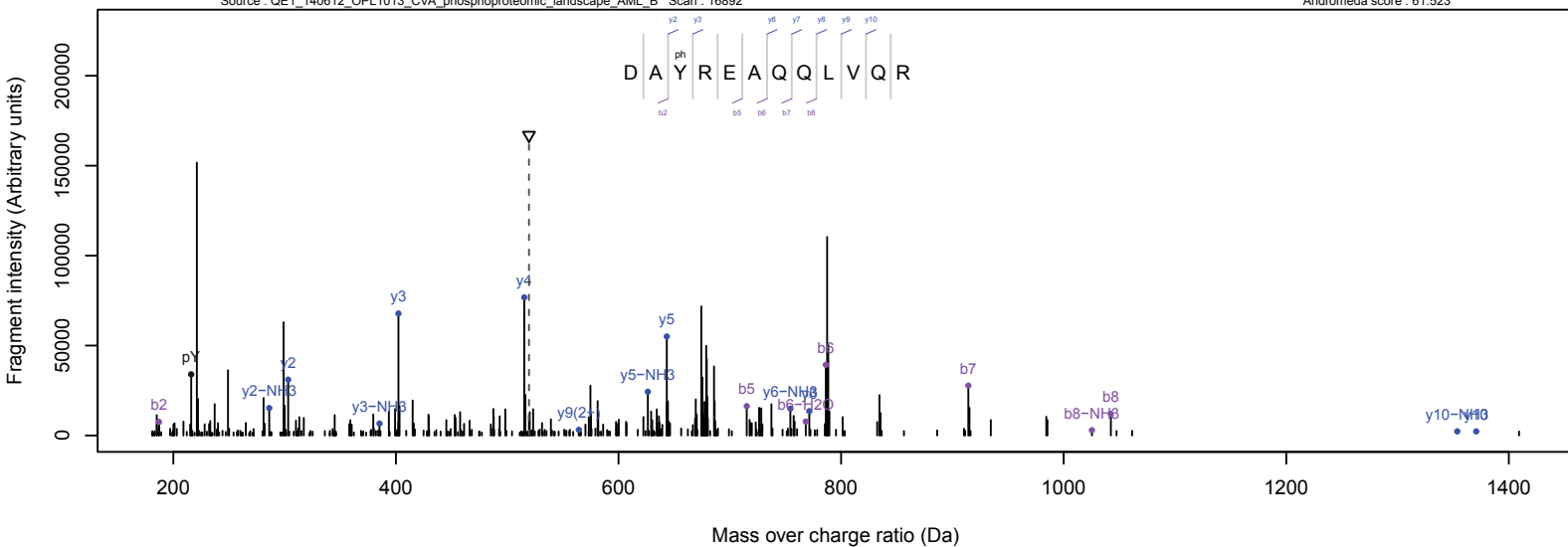
Andromeda score : 61.56



## DAY(ph)REAQLVQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16892

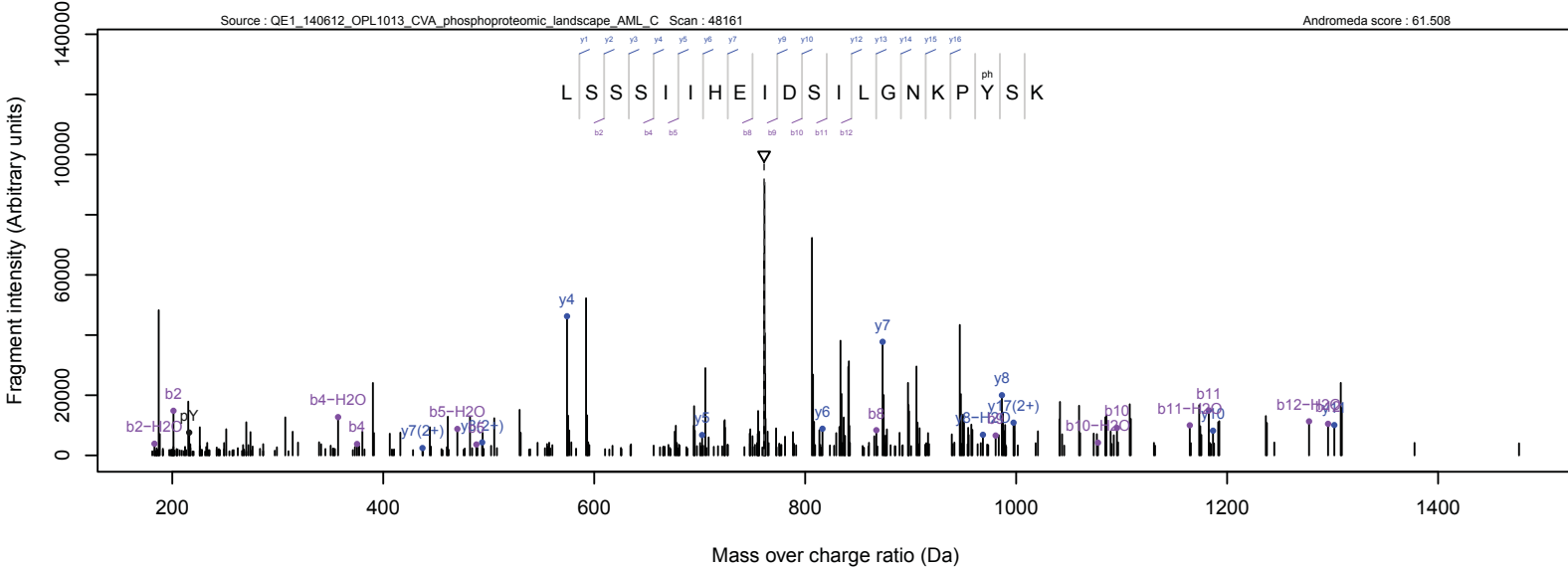
Andromeda score : 61.523



## LSSSIHEIDSLGKPY(ph)SK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 48161

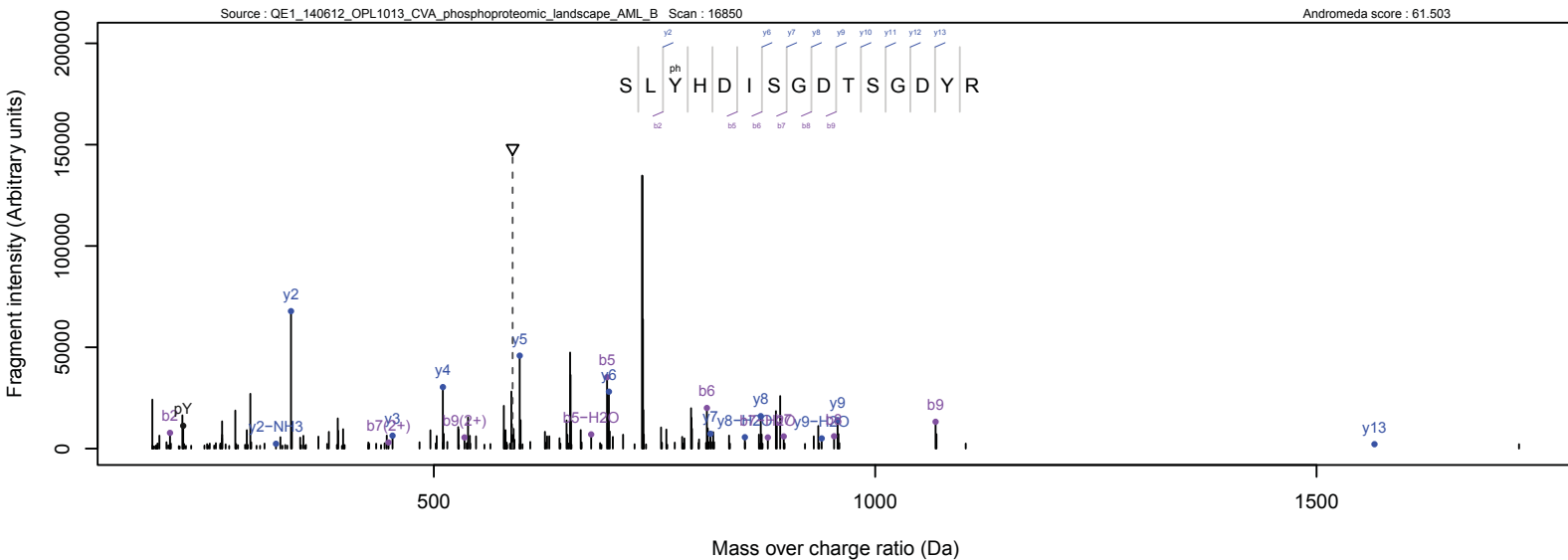
Andromeda score : 61.508



**SLY(ph)HDISGDTSGDYR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16850

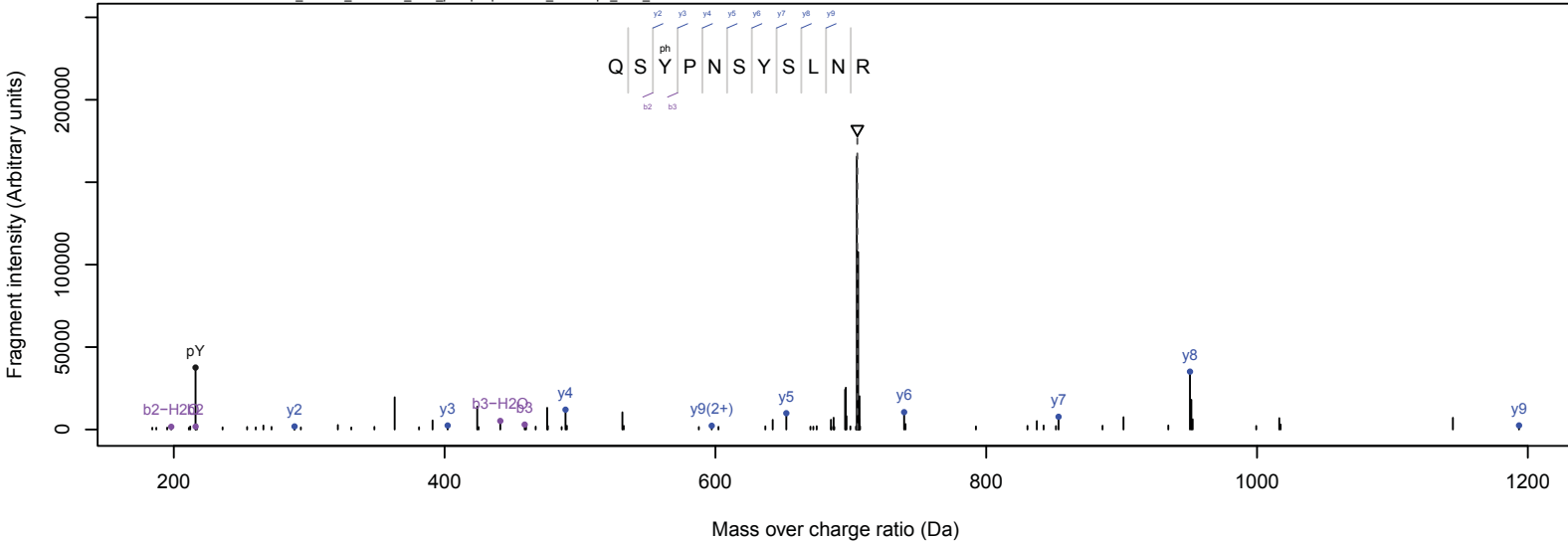
Andromeda score : 61.503



**QSY(ph)PNSYSLNR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11367

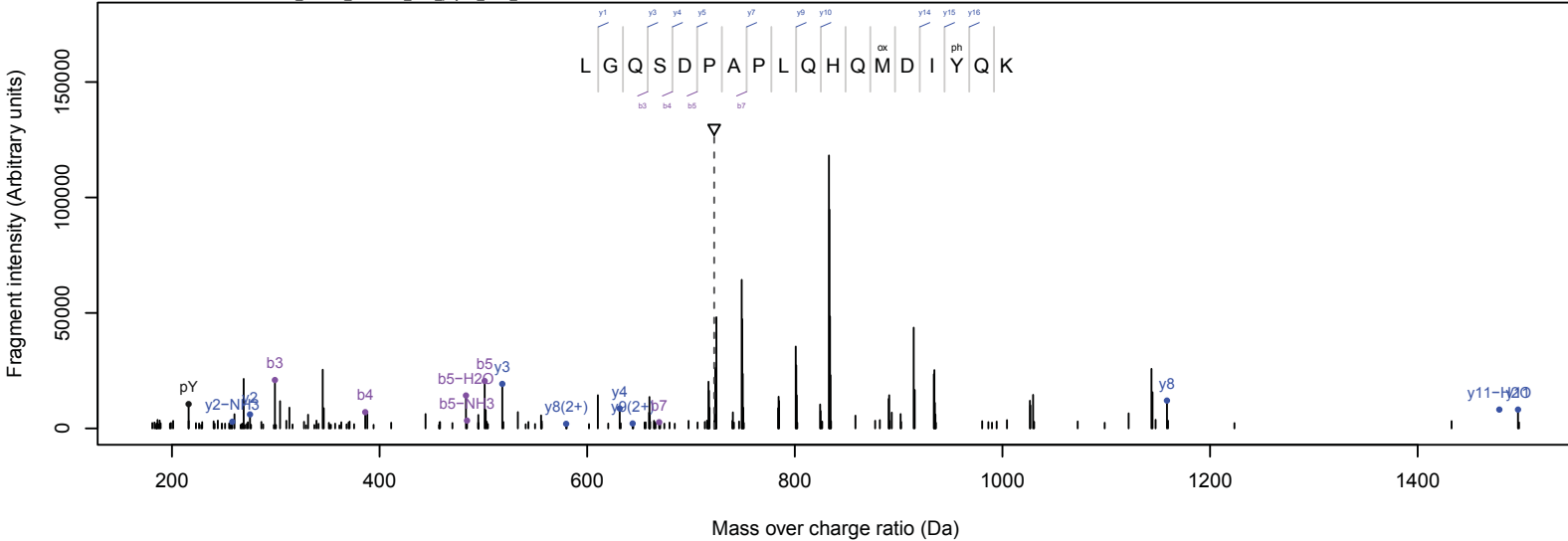
Andromeda score : 61.48



**LGQSDPAPLQHQM(ox)DIY(ph)QK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 15256

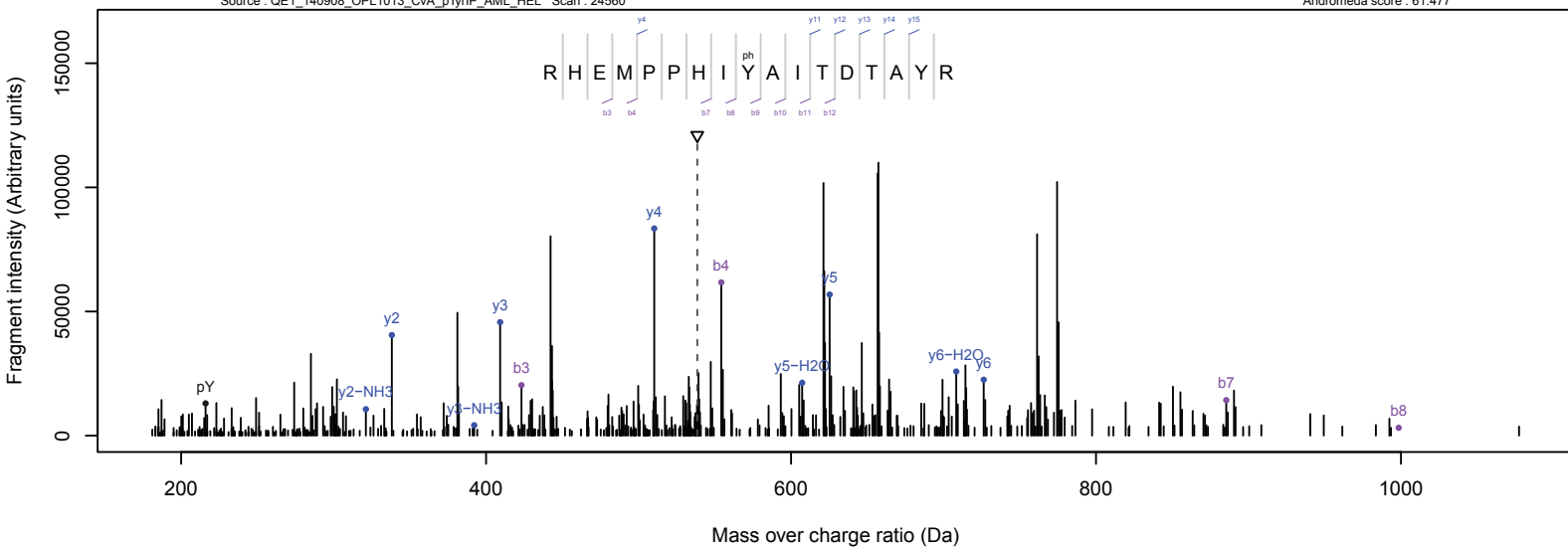
Andromeda score : 61.477



RHEMPPHIY(ph)AITDTAYR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24560

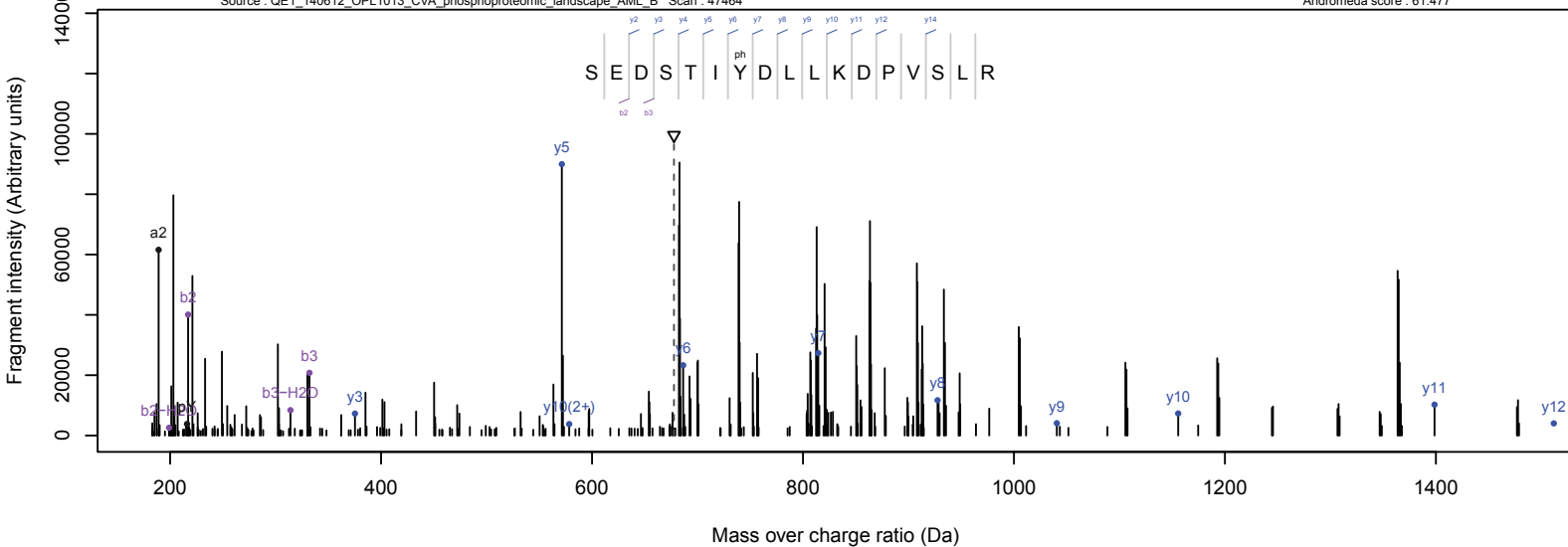
Andromeda score : 61.477



SEDSTIY(ph)DLLKDPVSLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 47464

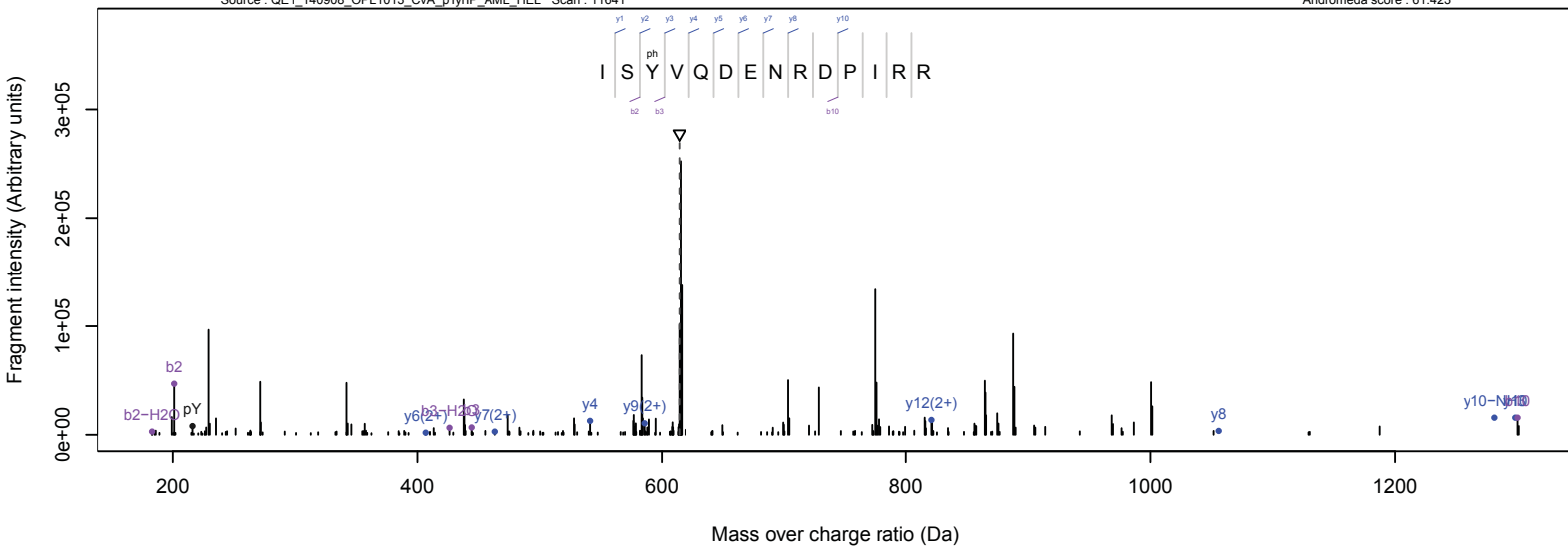
Andromeda score : 61.477



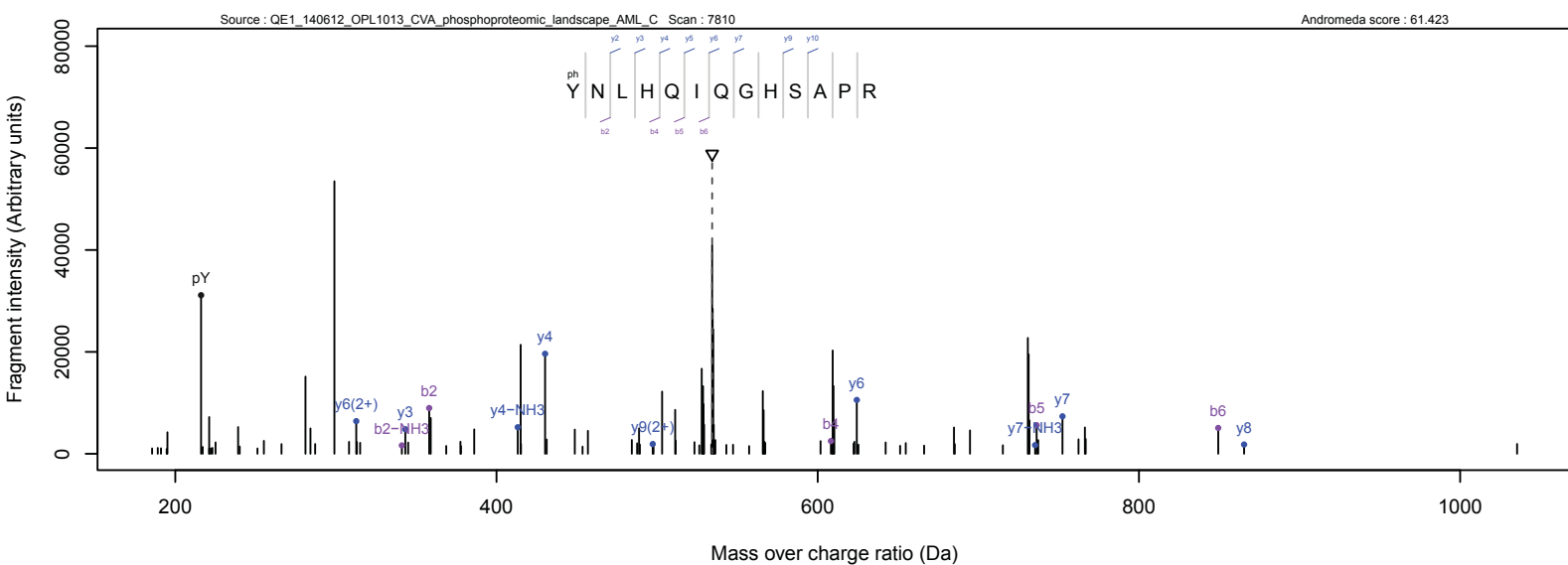
ISY(ph)VQDENRDPIRR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 11641

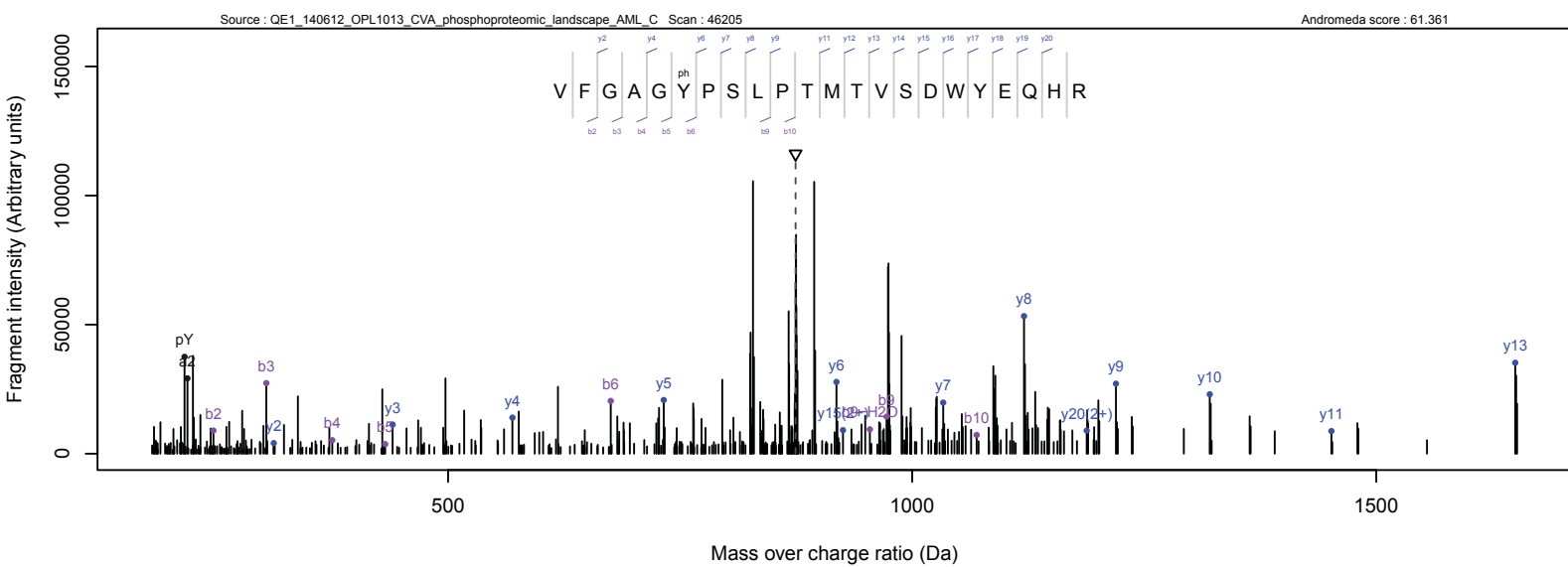
Andromeda score : 61.423



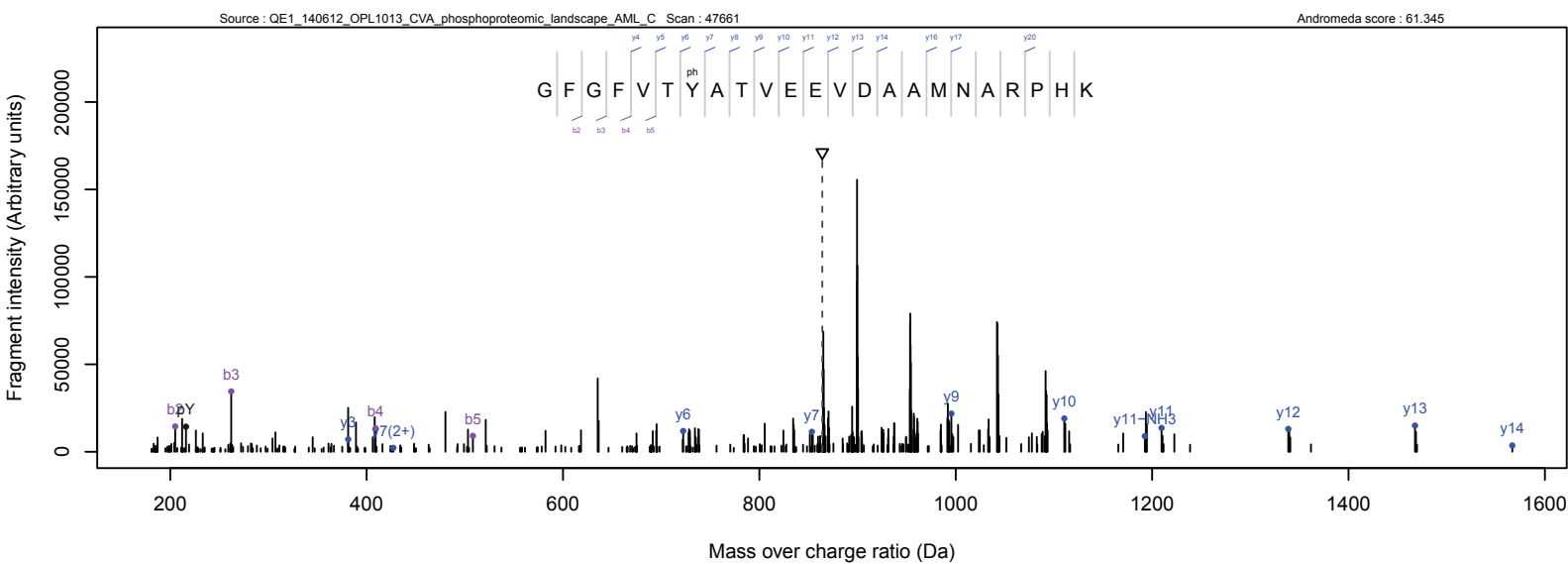
## Y(ph)NLHQIQGHSAPR-3+



## VFGAGY(ph)PSLPTMTVSDWYEQHR-3+



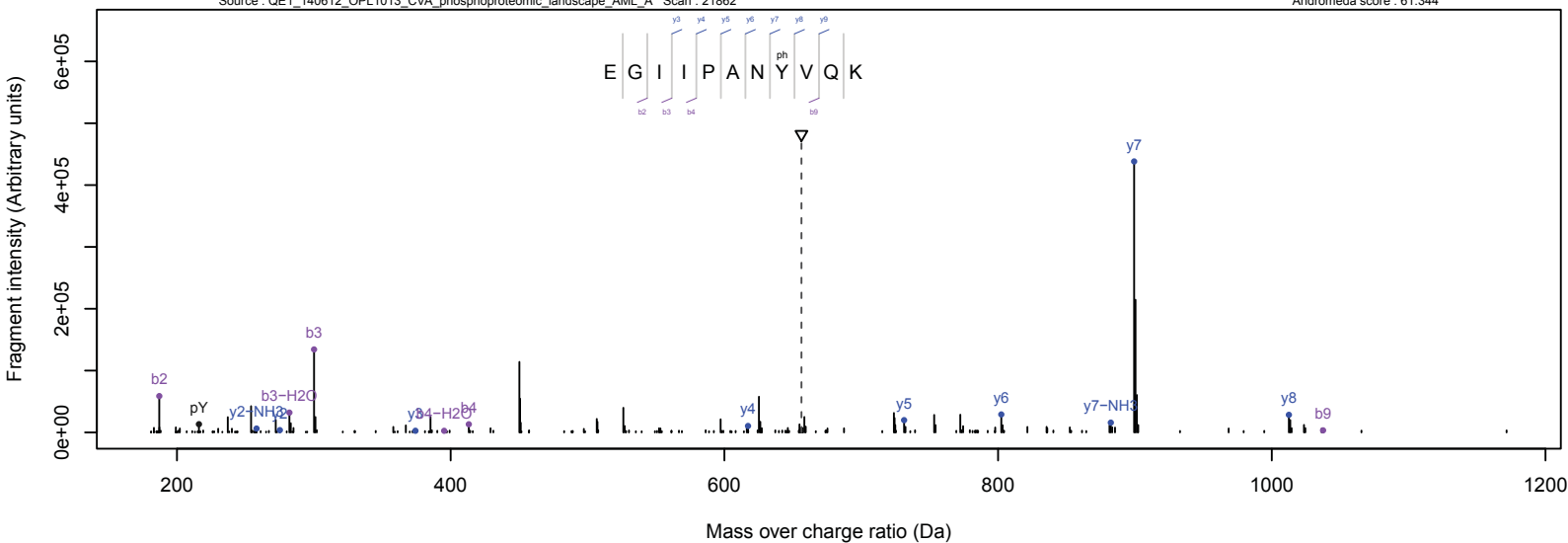
## GFGFVY(ph)ATVEEVDAAAMNARPHK-3+



## EGIIPANY(ph)VQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 21862

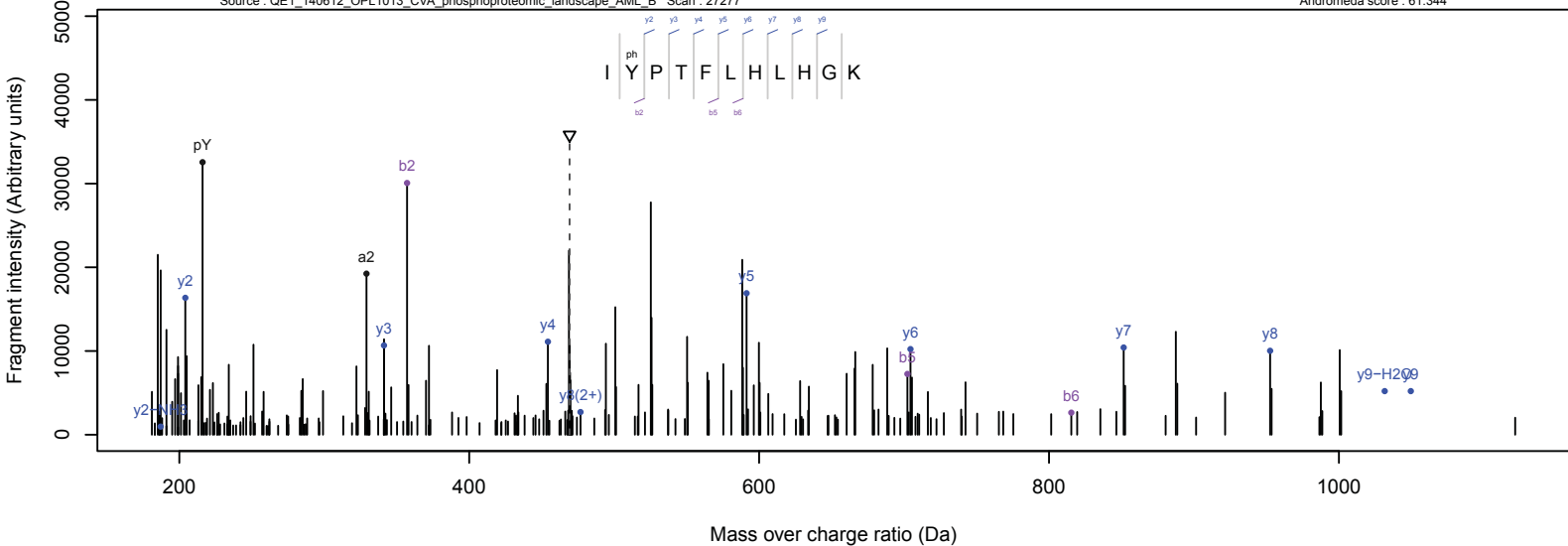
Andromeda score : 61.344



## IY(ph)PTFLHLHGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27277

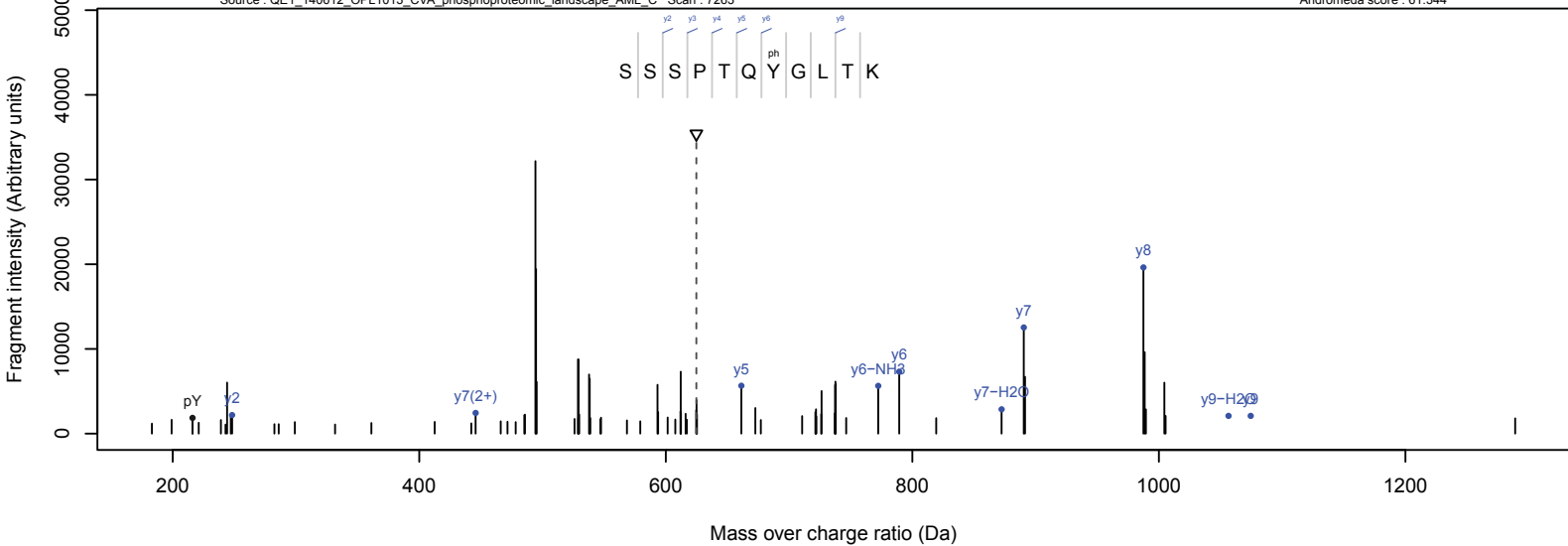
Andromeda score : 61.344



## SSSPQY(ph)GLTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7263

Andromeda score : 61.344

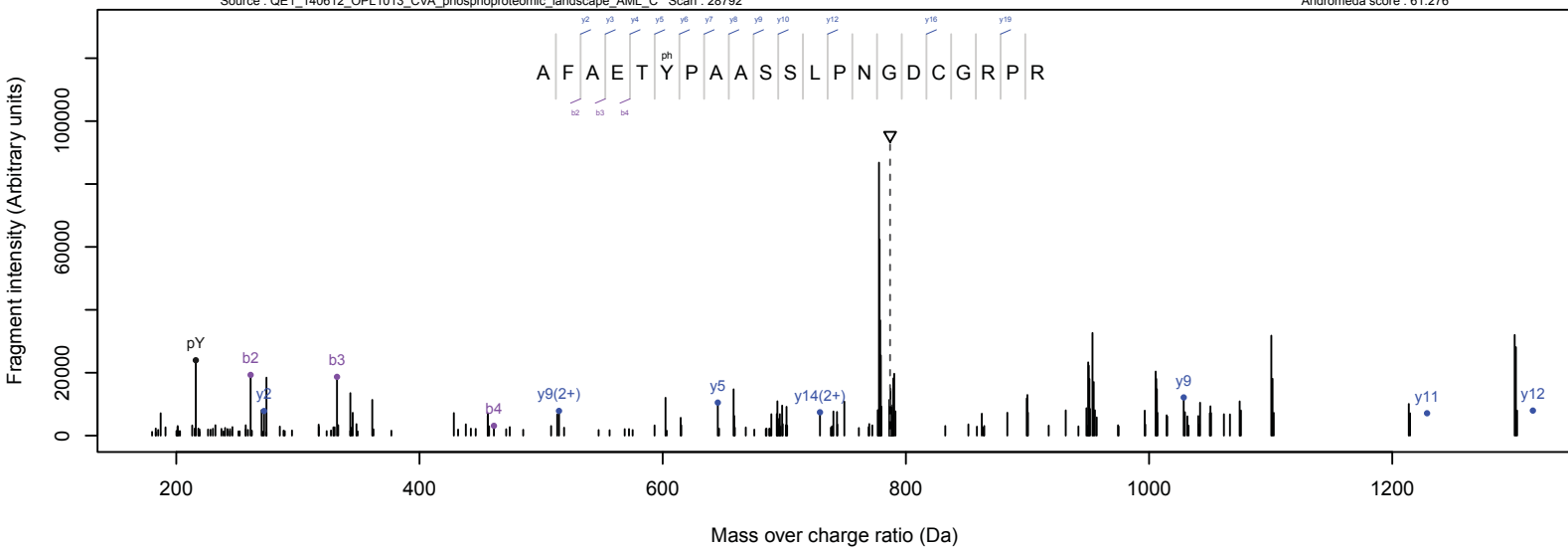




## (ac)AFAETY(ph)PAASSLPNGDCGRPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28792

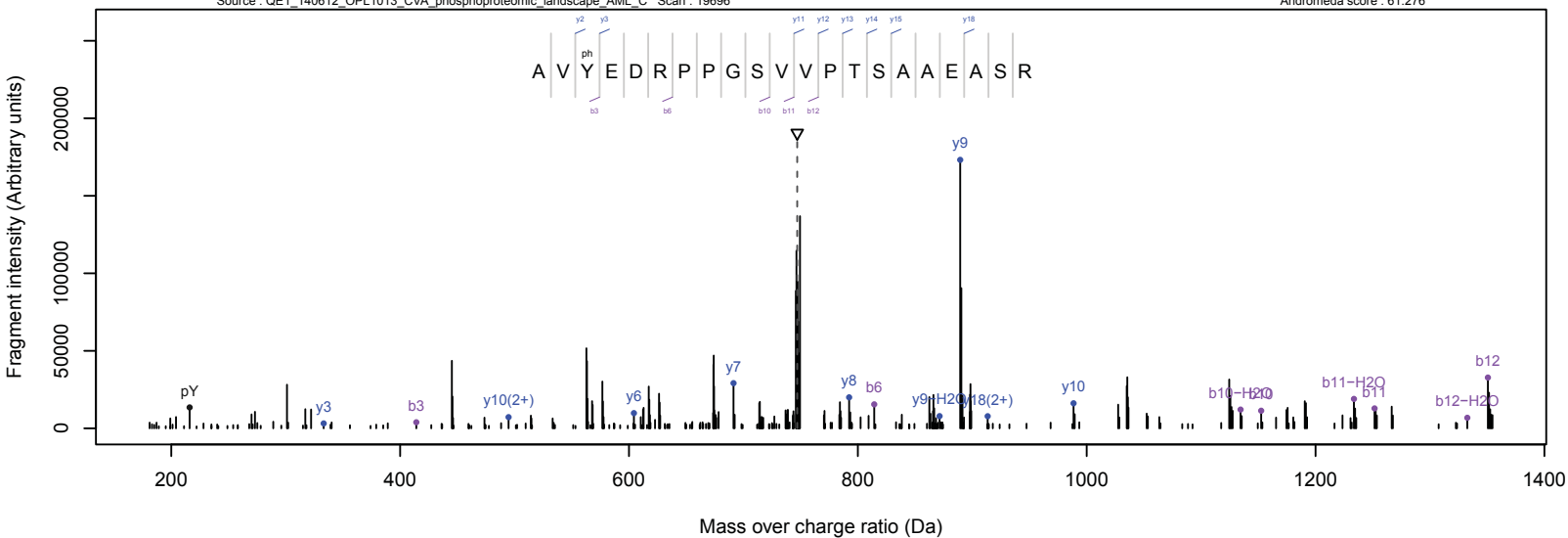
Andromeda score : 61.276



## AVY(ph)EDRPPGSVVPTSAAEASR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19696

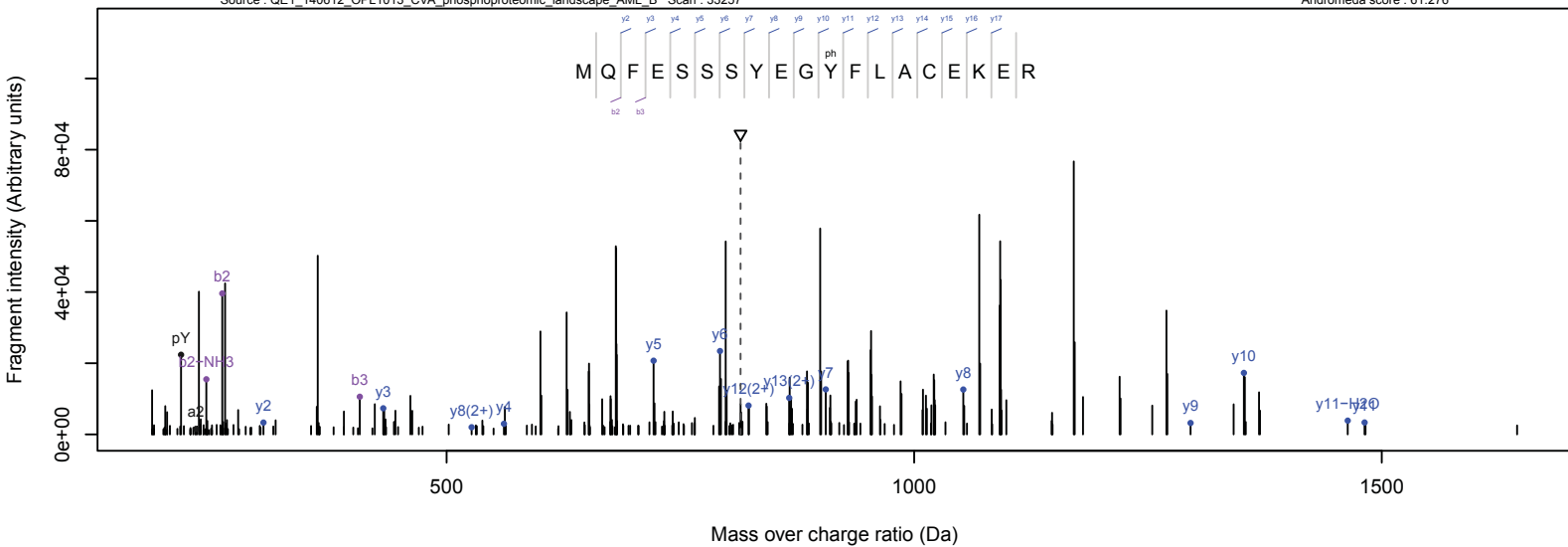
Andromeda score : 61.276



## MQFESSYEGY(ph)FLACEKER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33257

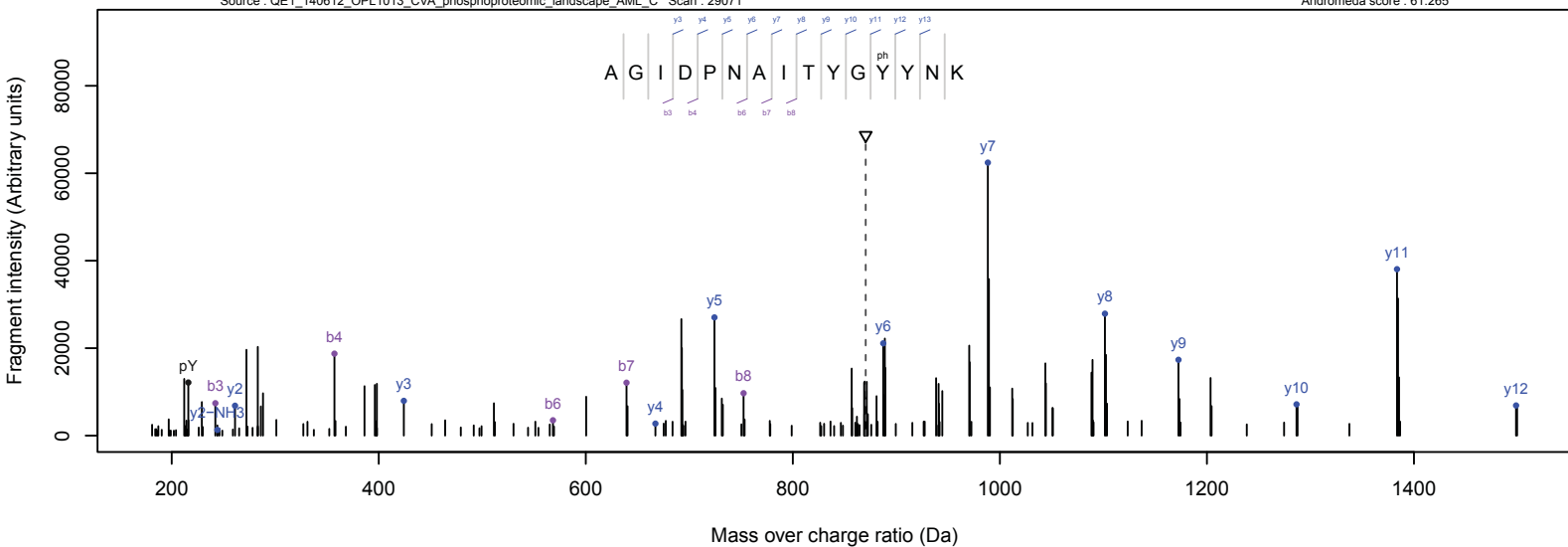
Andromeda score : 61.276



## AGIDPNAITYG(ph)YNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29071

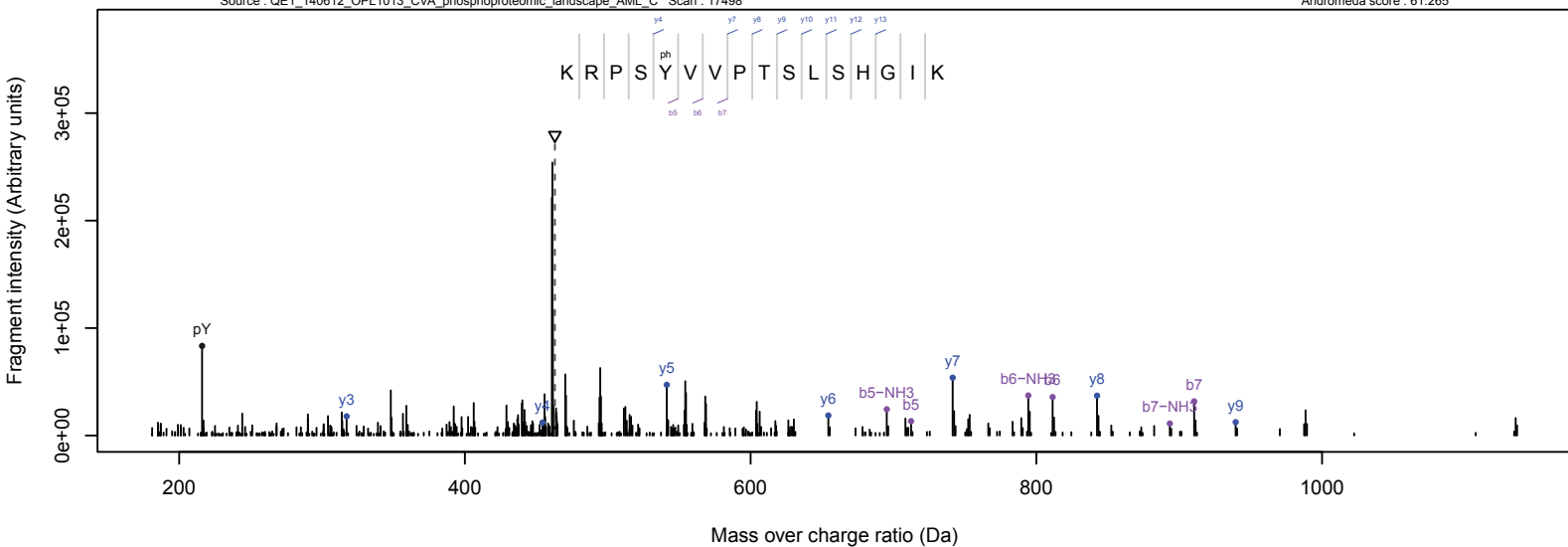
Andromeda score : 61.265



## KRPSY(ph)VVPTSLSHGIK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17498

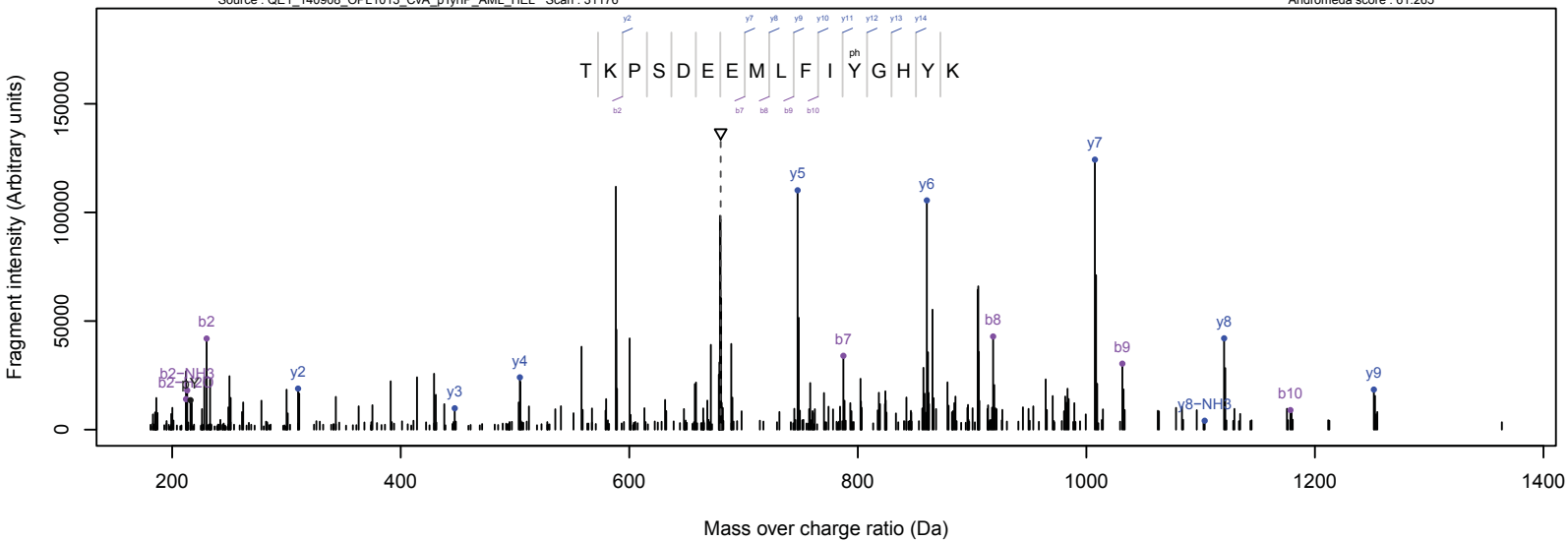
Andromeda score : 61.265



## TKPSDEEMLFY(ph)GHYK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31176

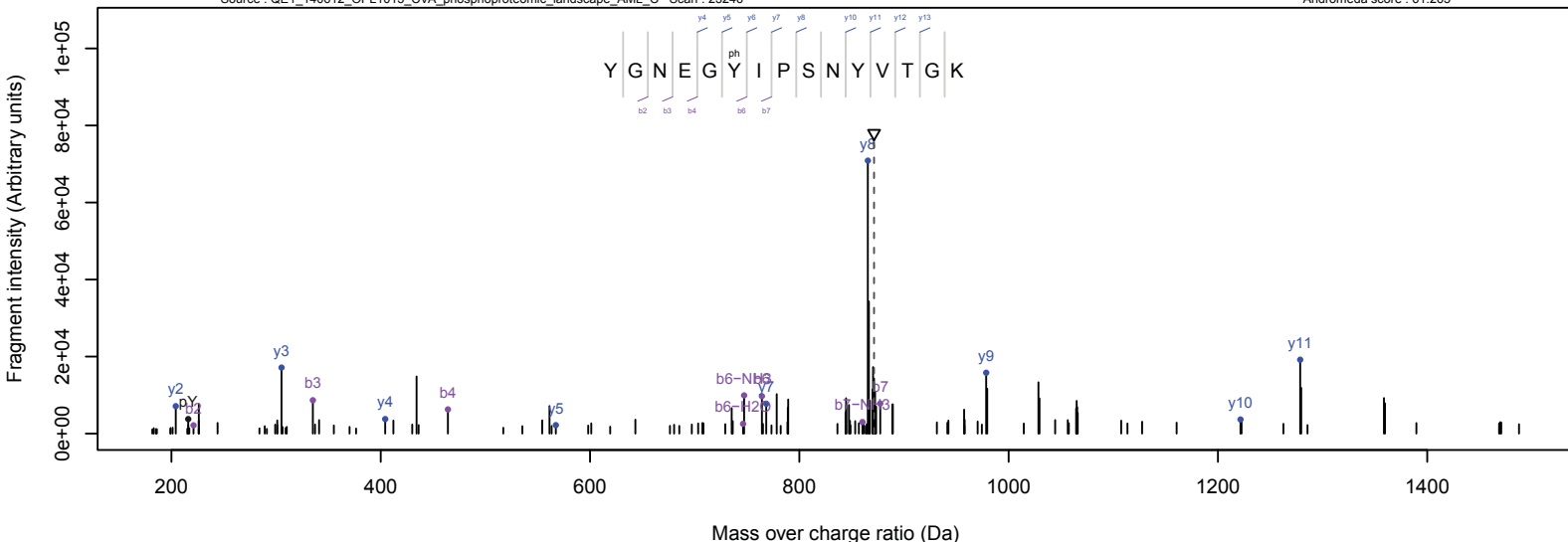
Andromeda score : 61.265



YGNEGY(ph)IPSNYVTGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23246

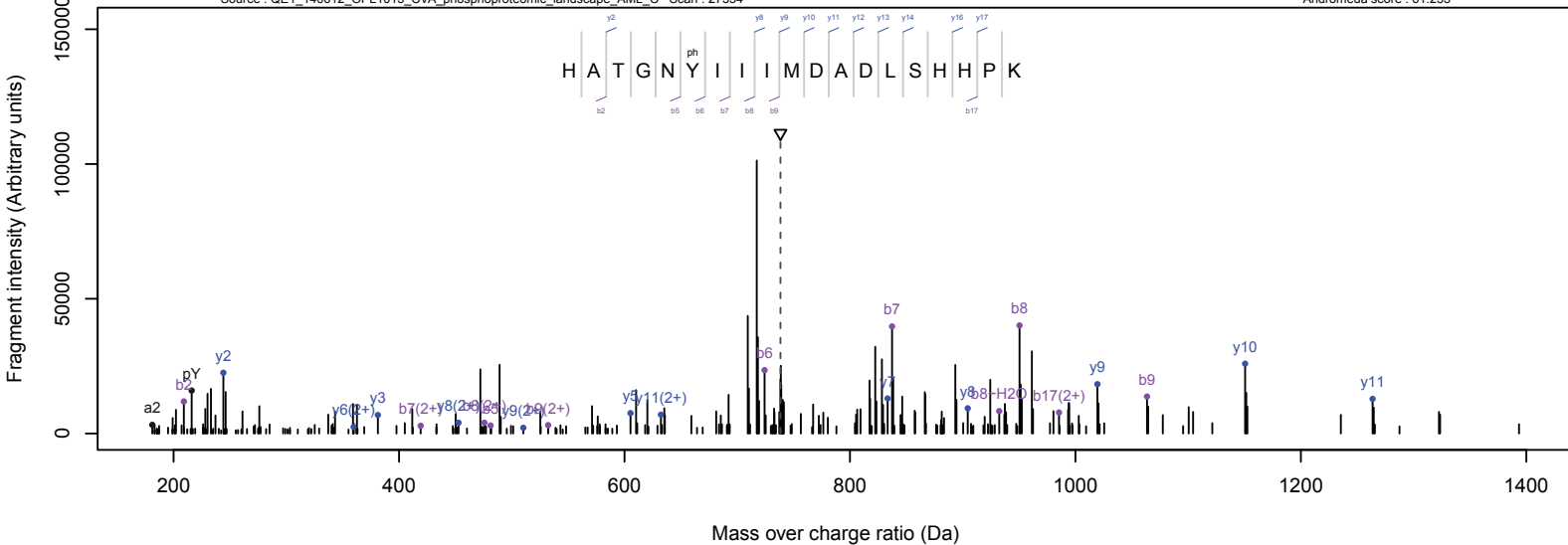
Andromeda score : 61.265



HATGNY(ph)IIMDADLSHHPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27554

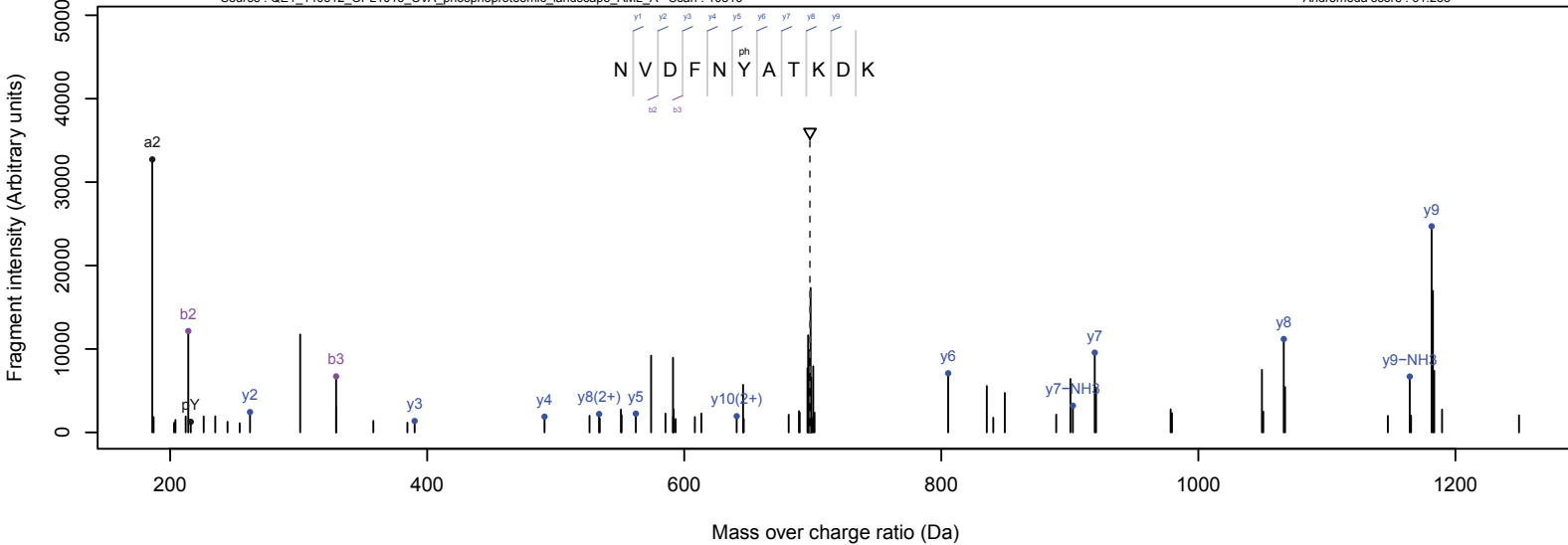
Andromeda score : 61.253



NVDFNY(ph)ATKDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 10816

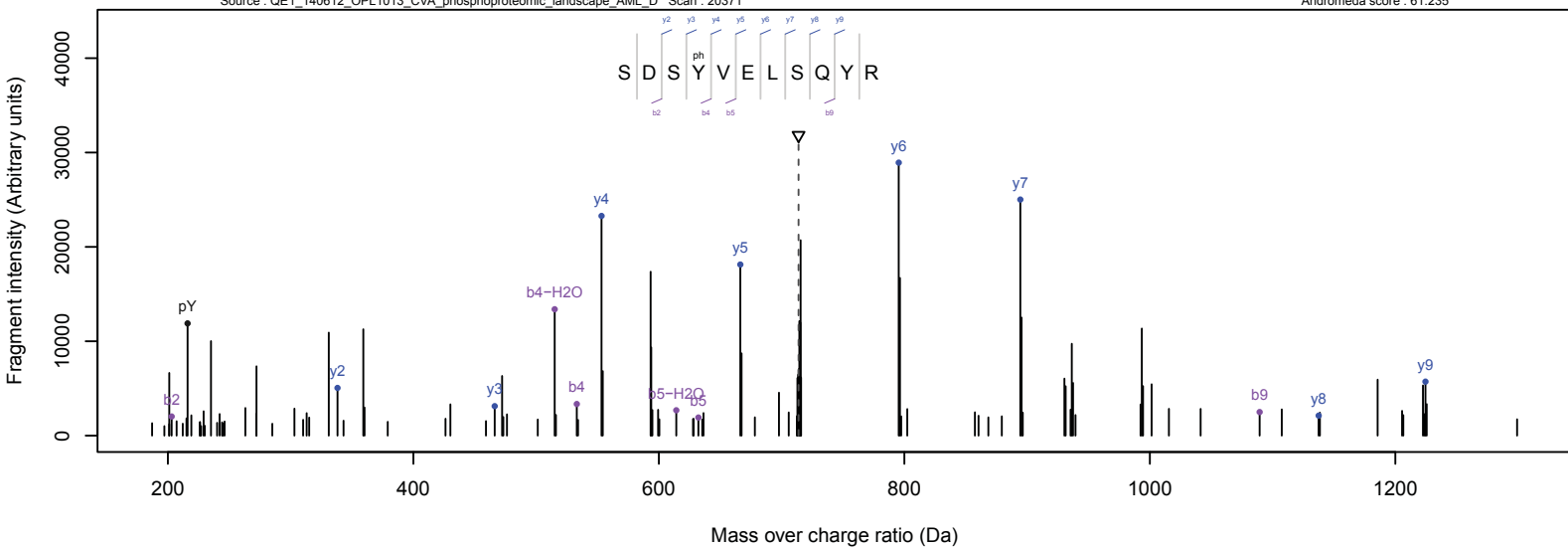
Andromeda score : 61.235



SDSY(ph)VELSQYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 20371

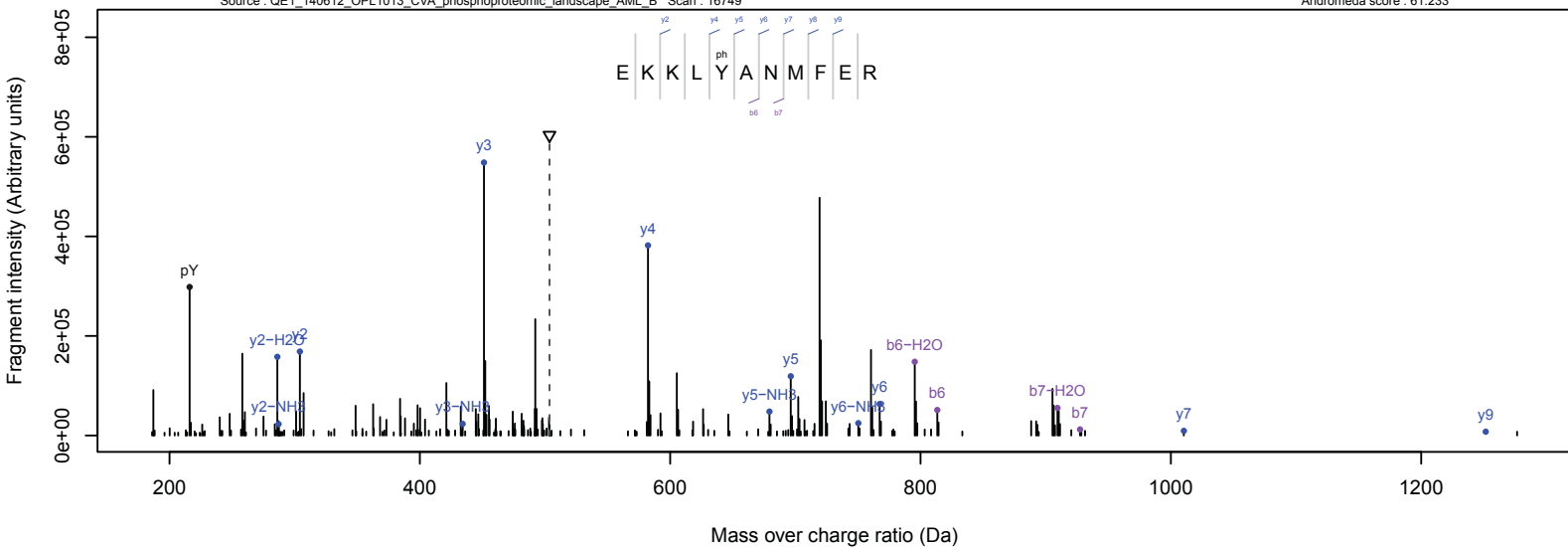
Andromeda score : 61.235



EKKLY(ph)ANMFER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16749

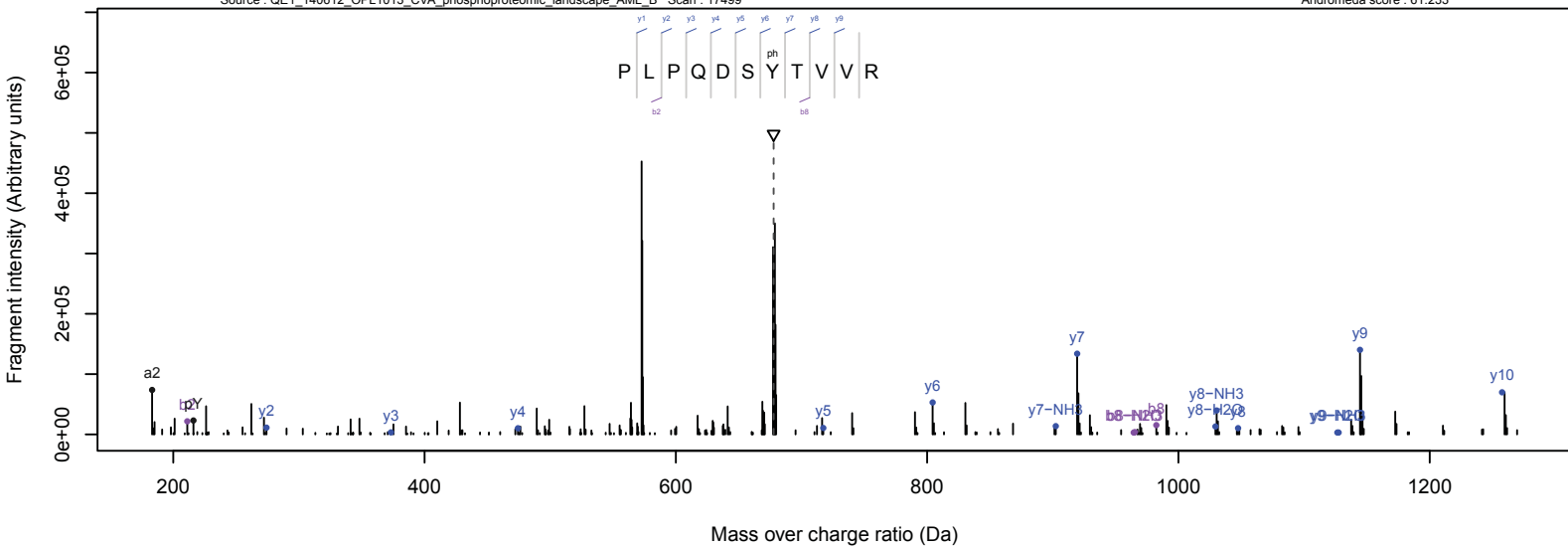
Andromeda score : 61.233



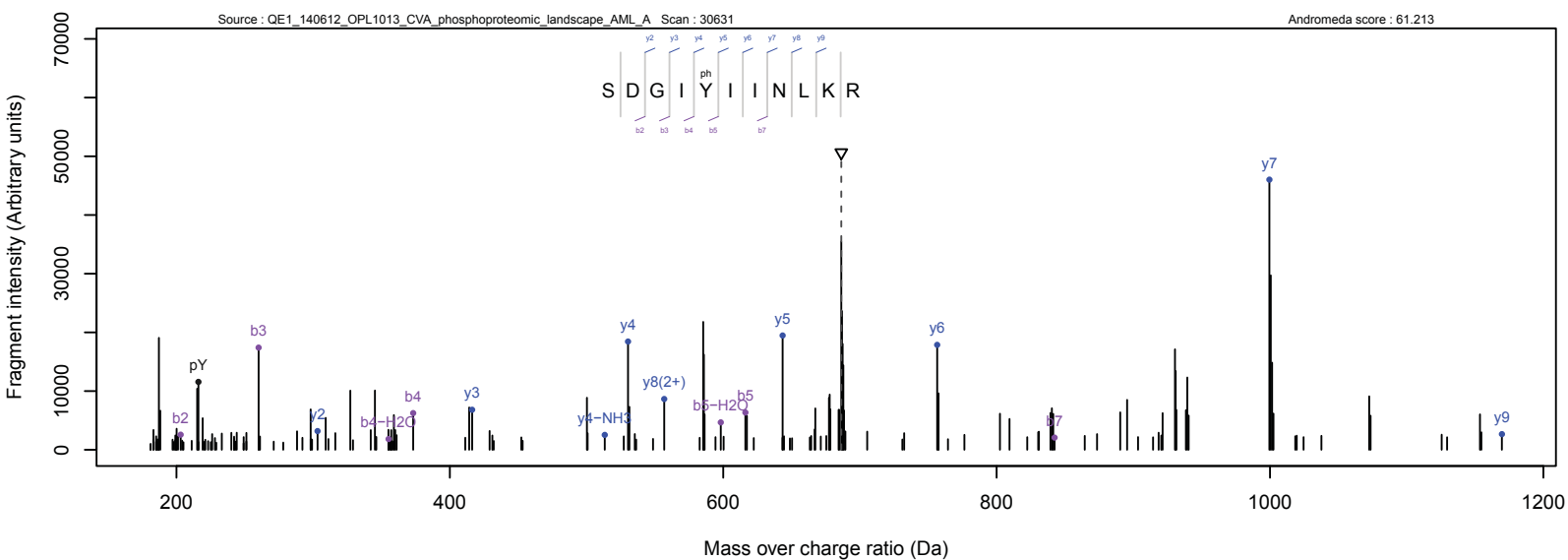
PLPQDSY(ph)TVVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17499

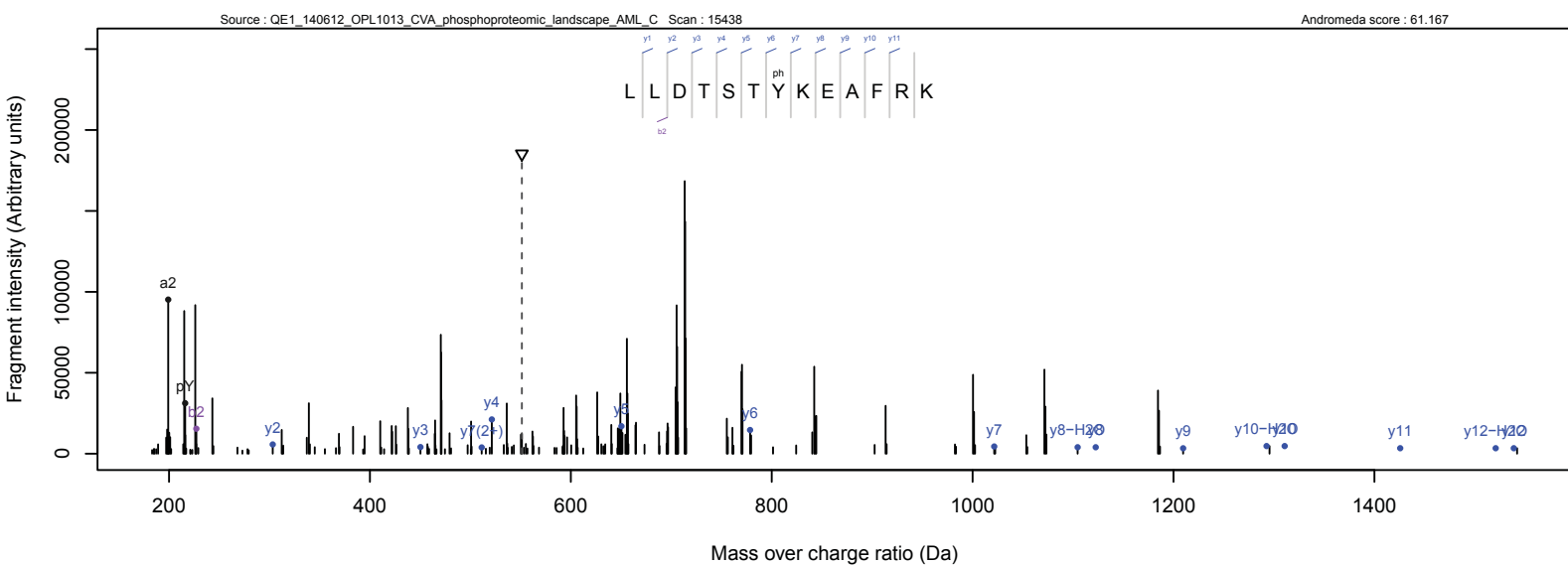
Andromeda score : 61.233



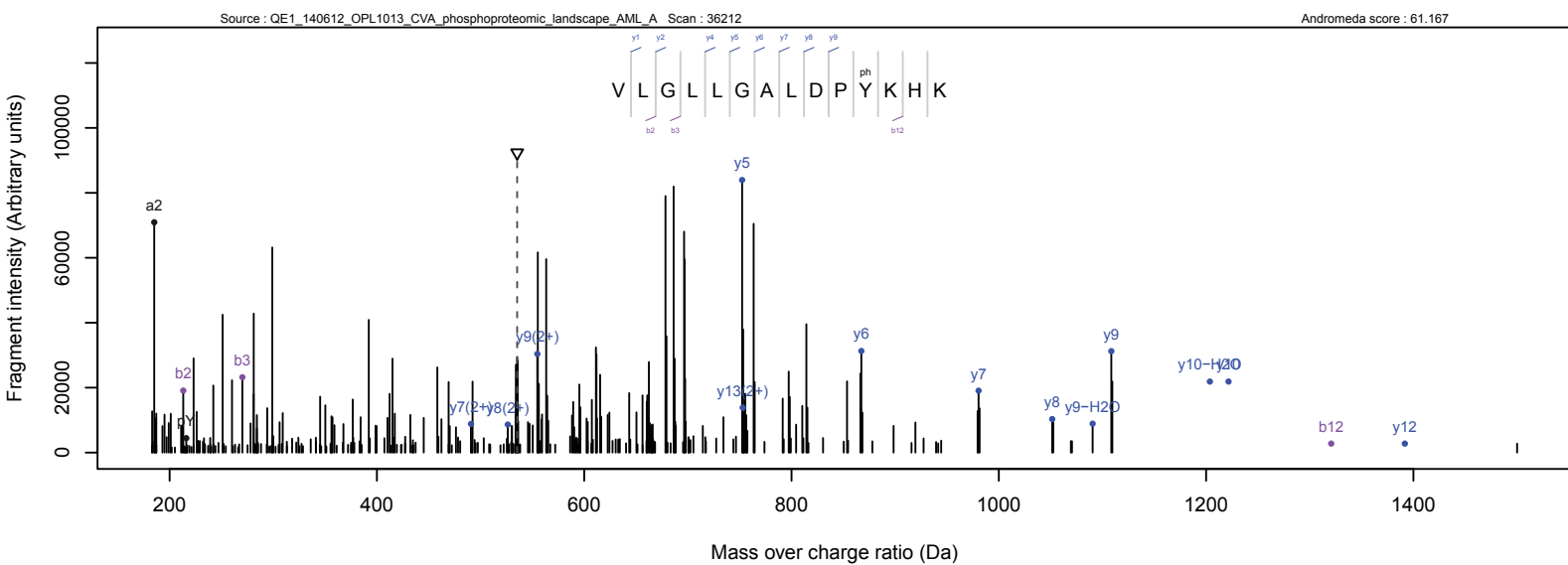
## SDGIY(ph)IINLKR-2+

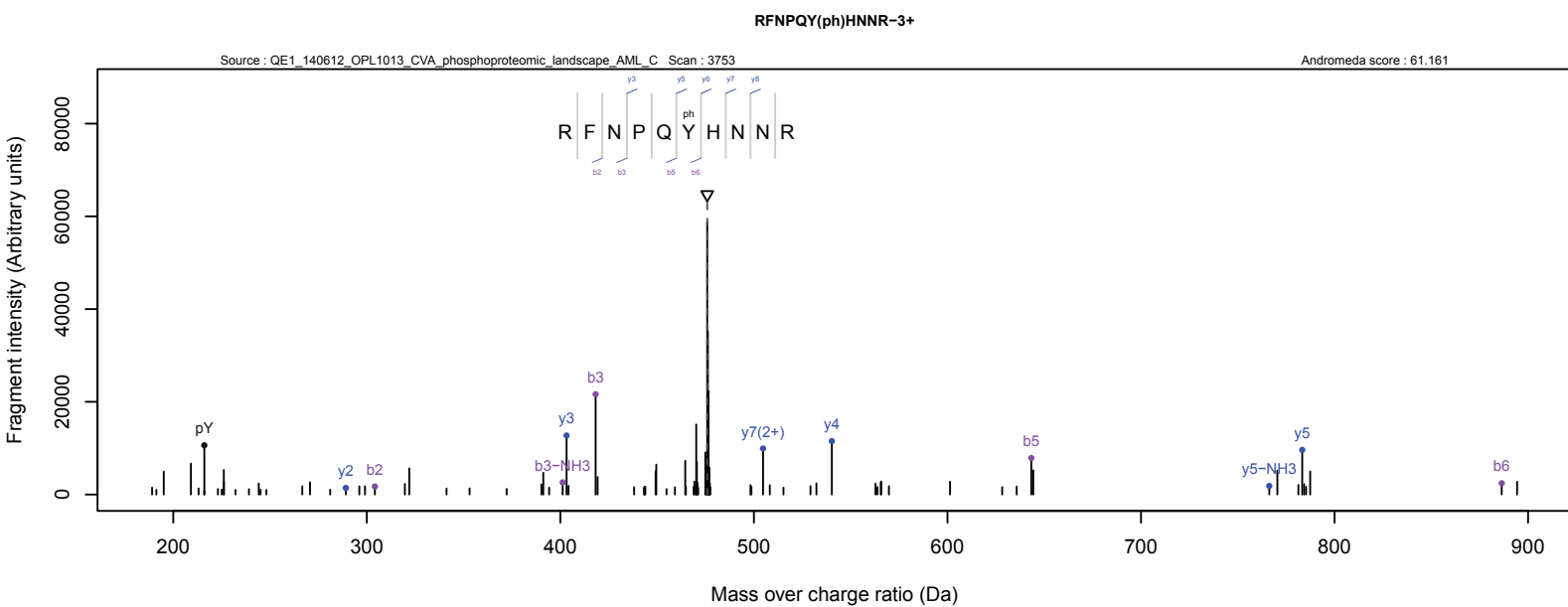
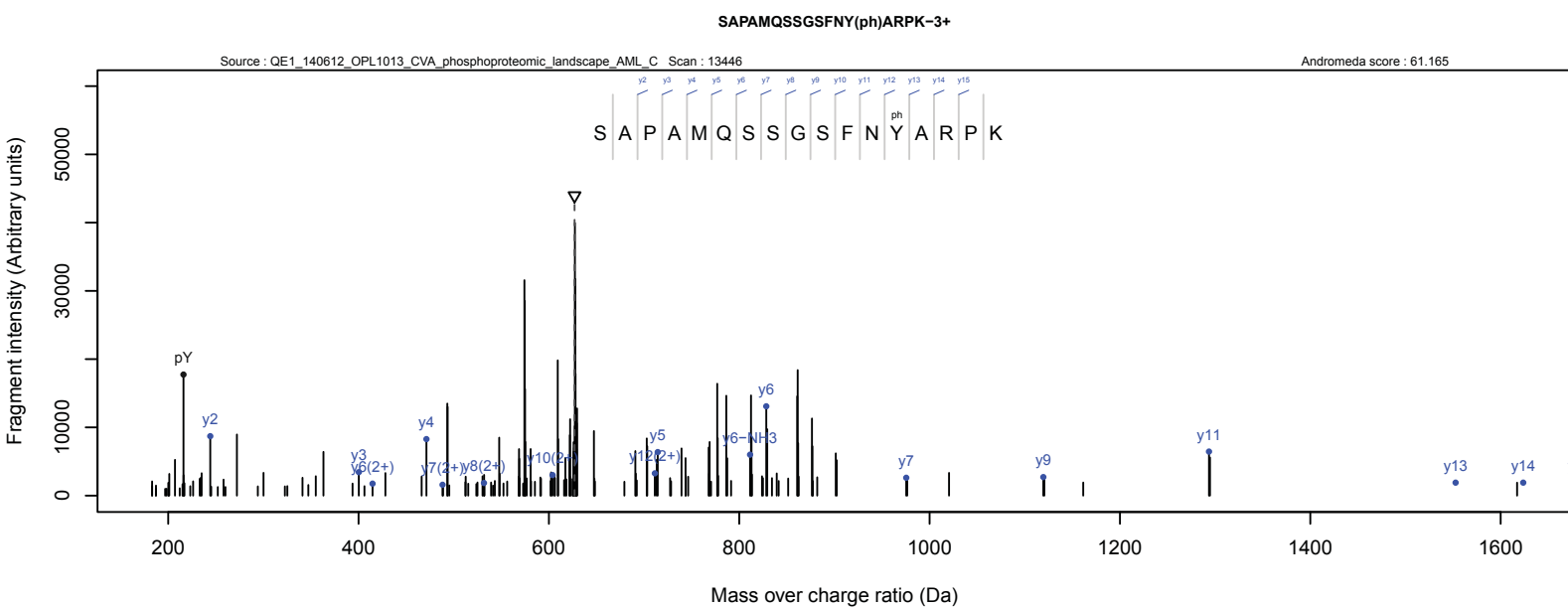
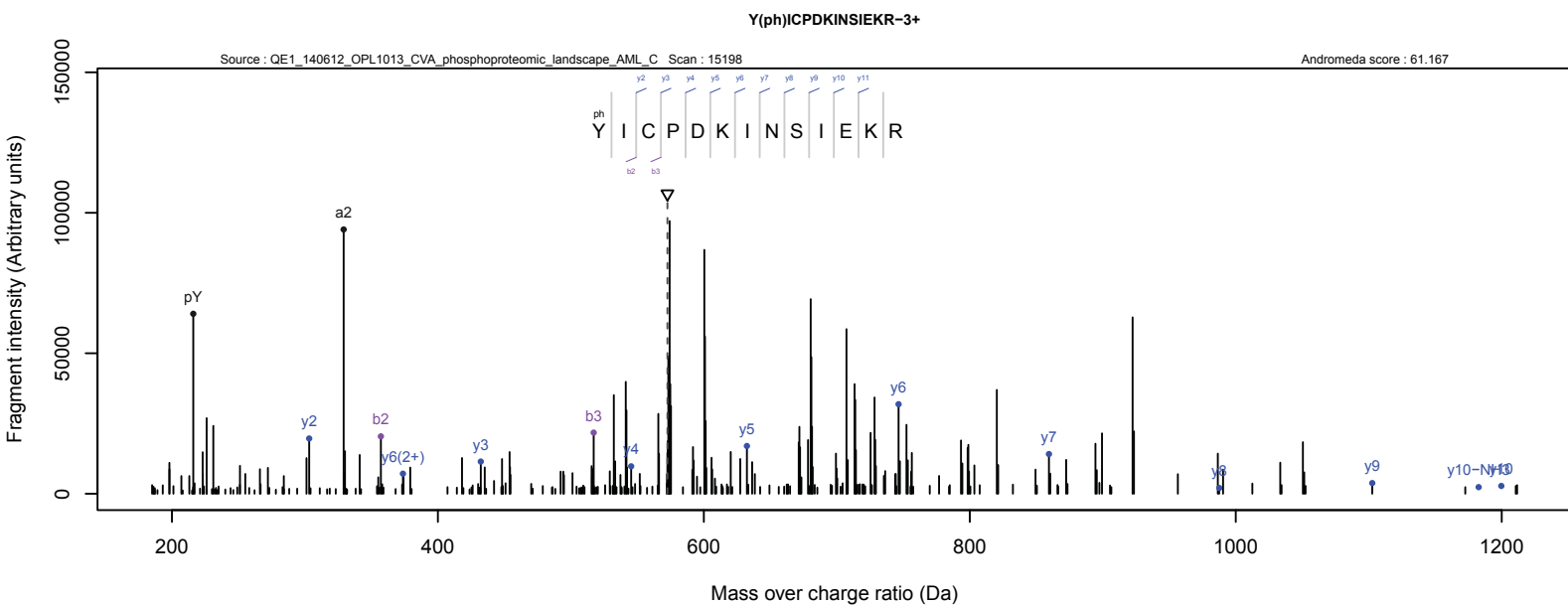


## LLDTSTY(ph)KEAFRK-3+



## VLGLLALDYPY(ph)KHK-3+



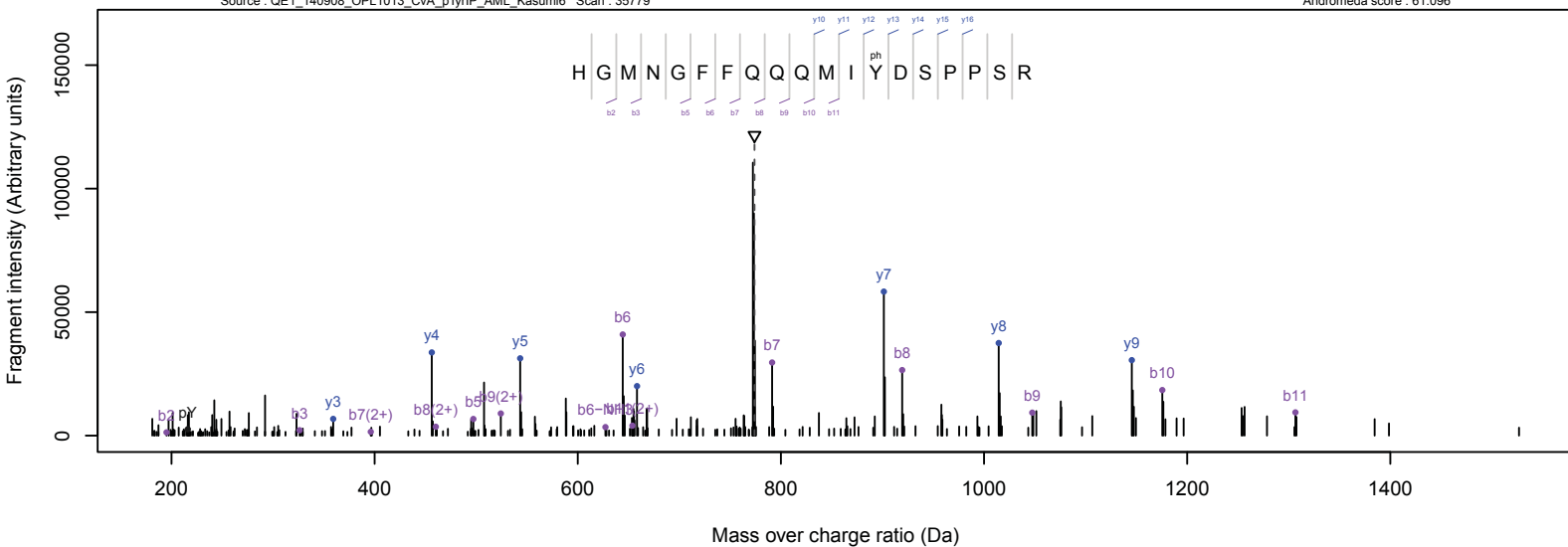




HGMNGFFQQMIY(ph)DSPPSR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 35779

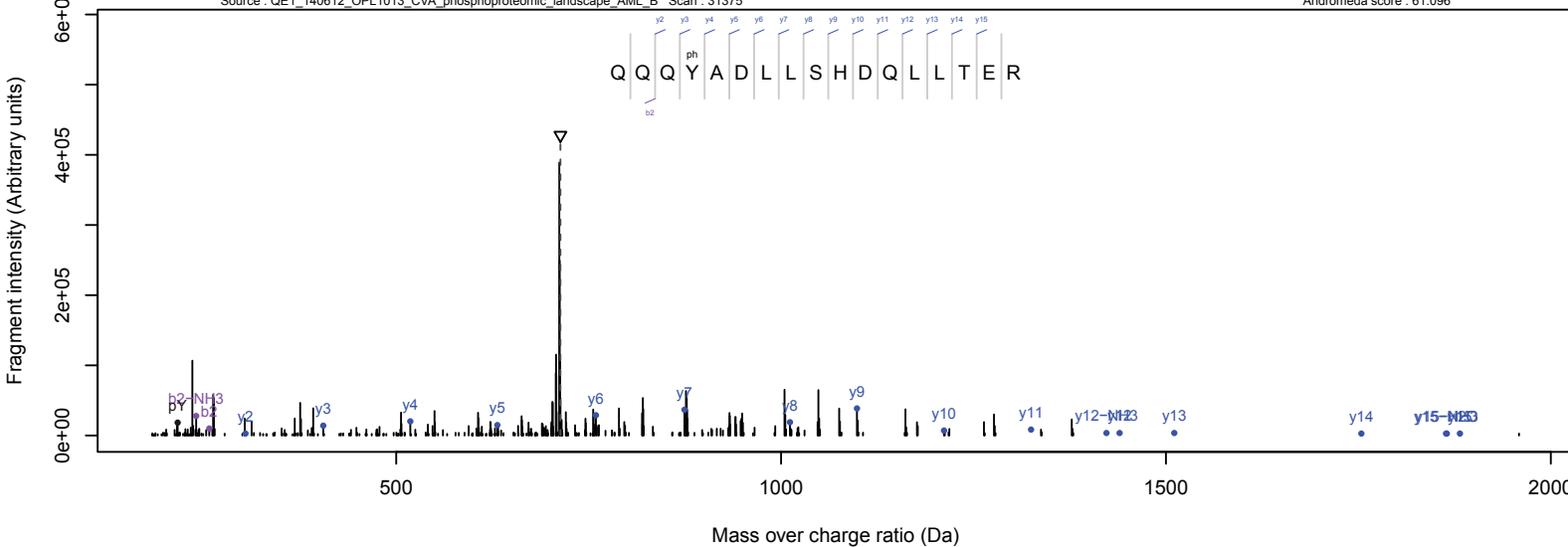
Andromeda score : 61.096



QQQY(ph)ADLLSHDQLLTER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31375

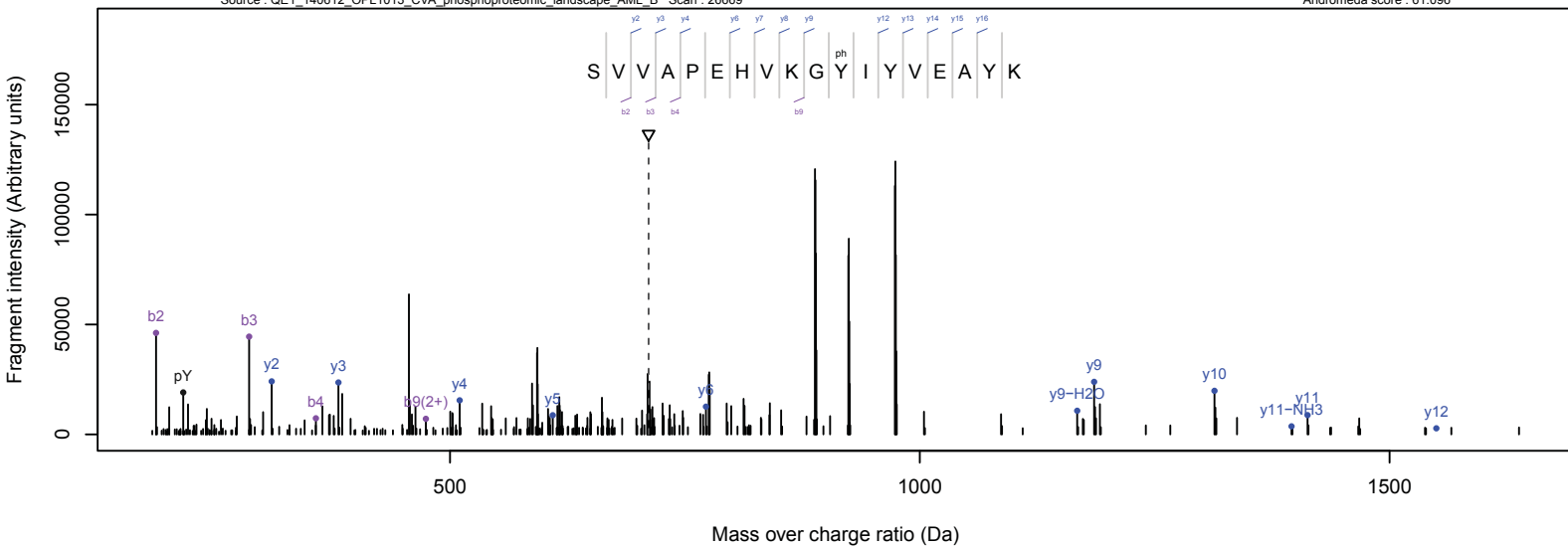
Andromeda score : 61.096



SVVAPEHVKG(ph)YVEAYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26669

Andromeda score : 61.096

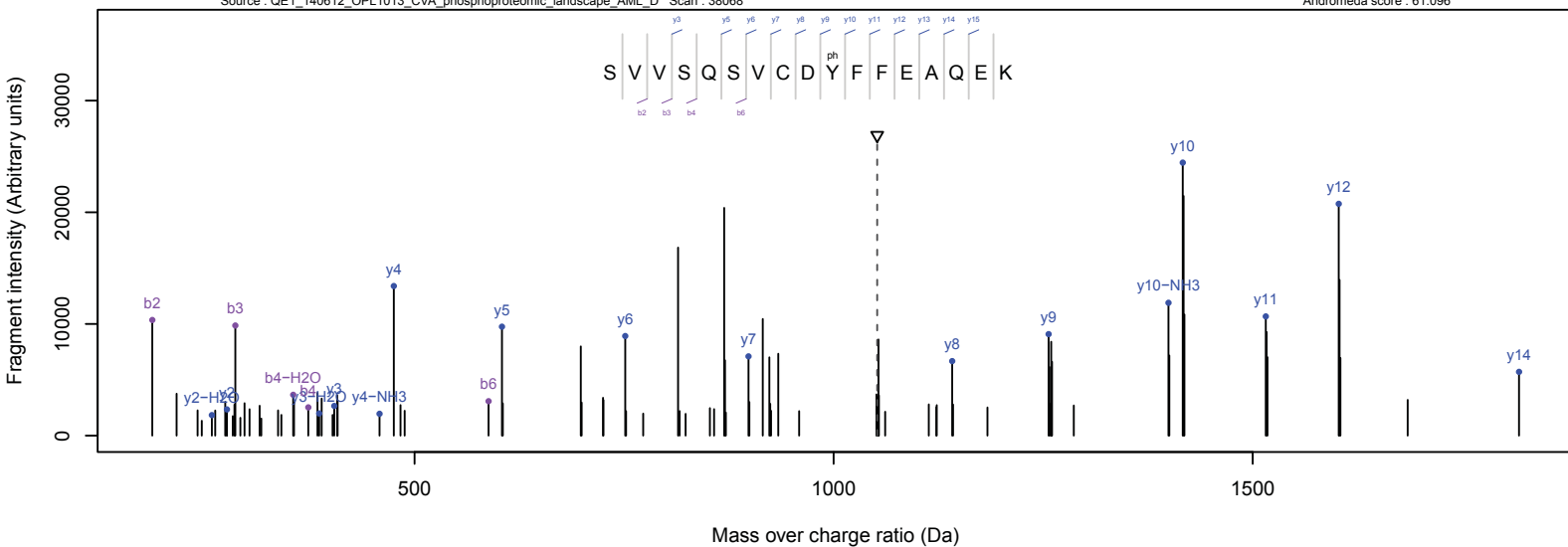




## SVVSQSVCDY(ph)FFEAQEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 38068

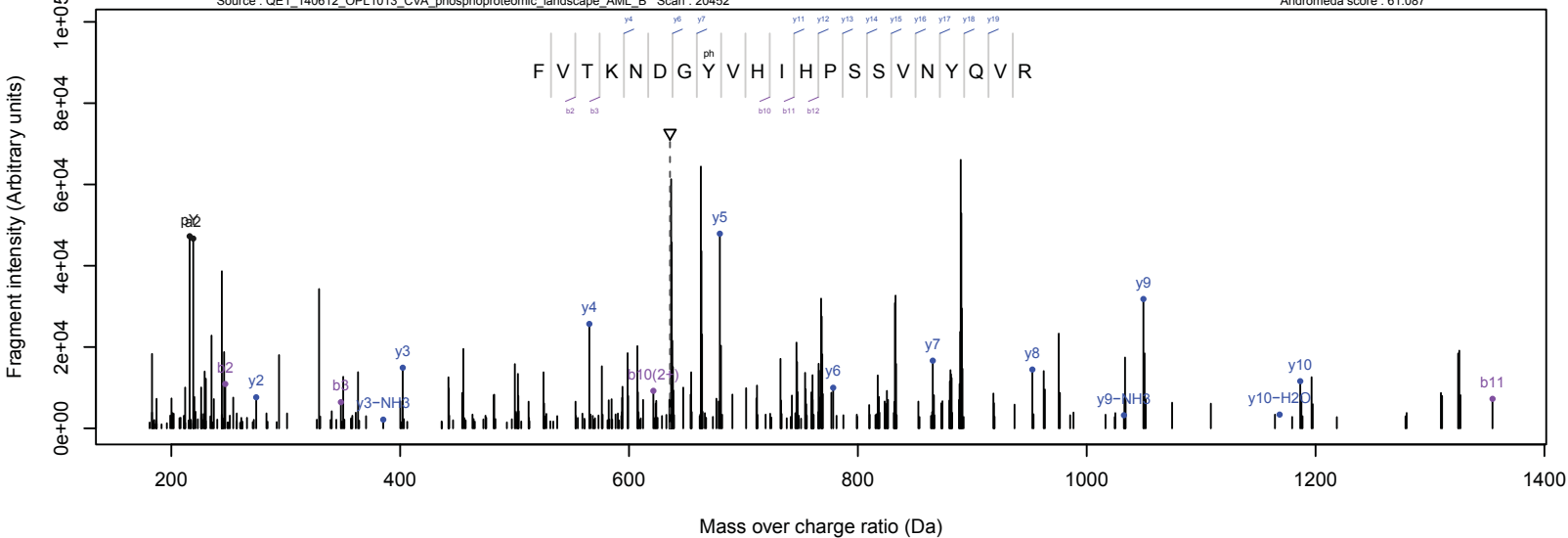
Andromeda score : 61.096



## FVTKNDGY(ph)VHIHPSSVNYQVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20452

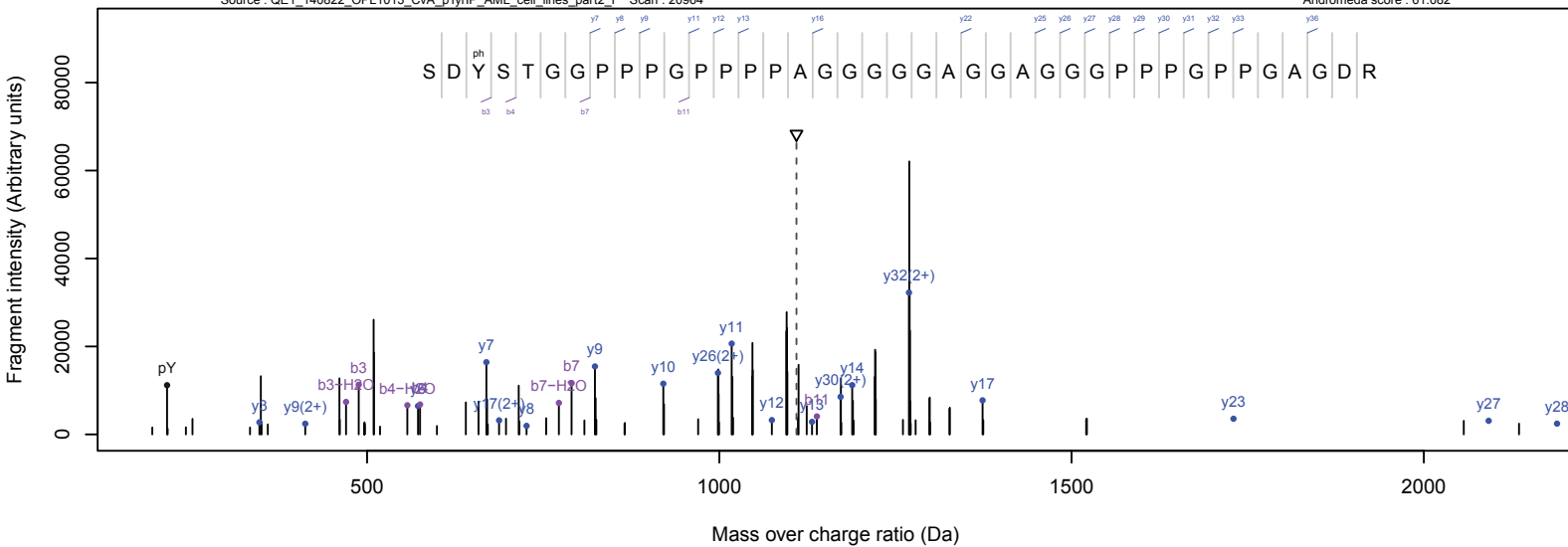
Andromeda score : 61.087



## (ac)SDY(ph)STGGPPPGPPPAGGGGGAGGAGGPPPPPPGAGDR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 20964

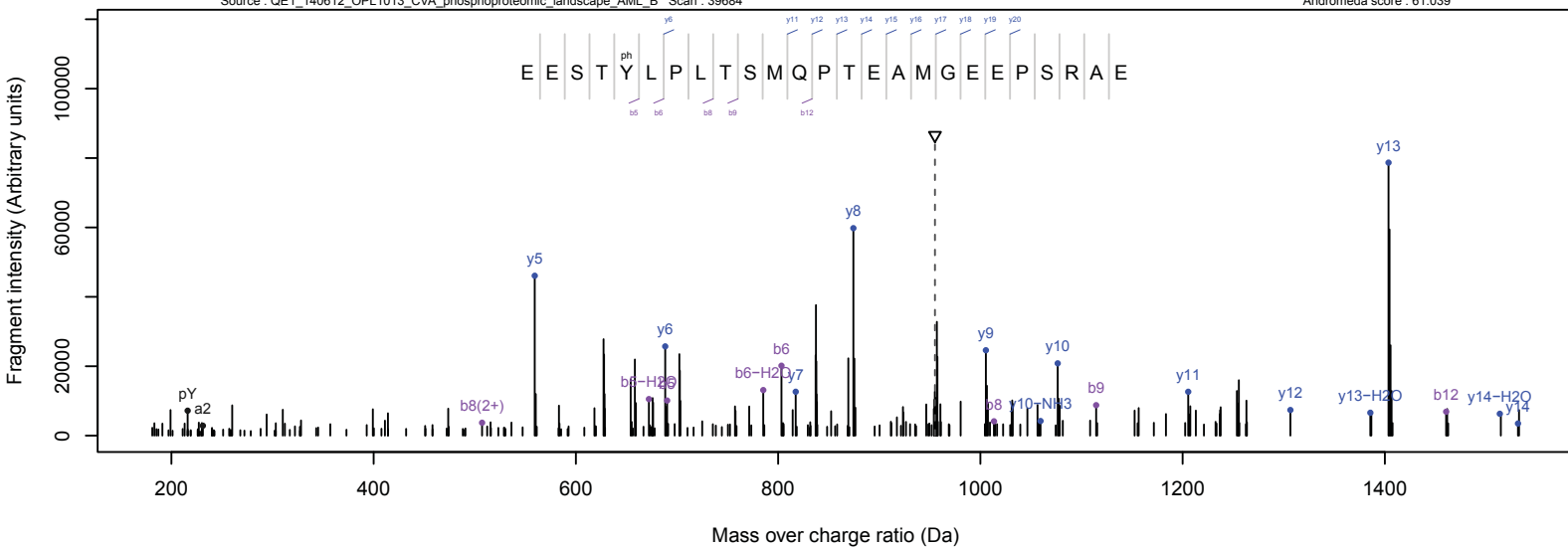
Andromeda score : 61.082



**EESTY(ph)LPLTSMQPTEAMGEEPSRAE-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39684

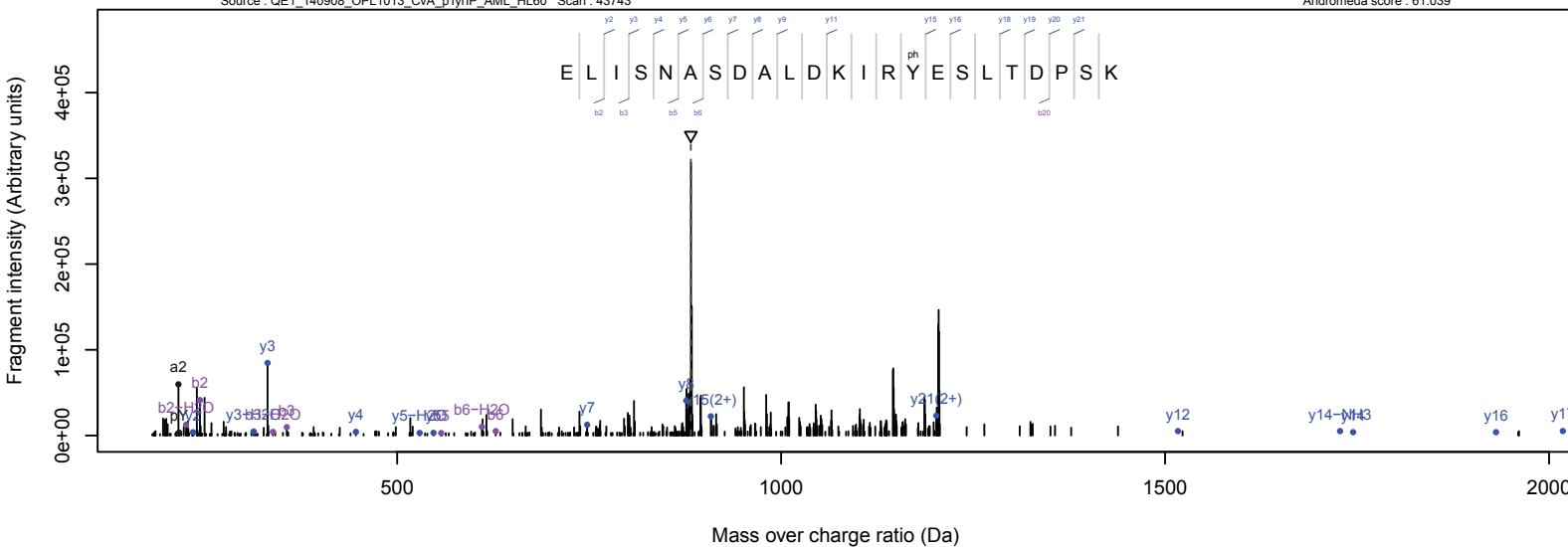
Andromeda score : 61.039



**ELISNASDALDKIRY(ph)ESLTDPK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 43743

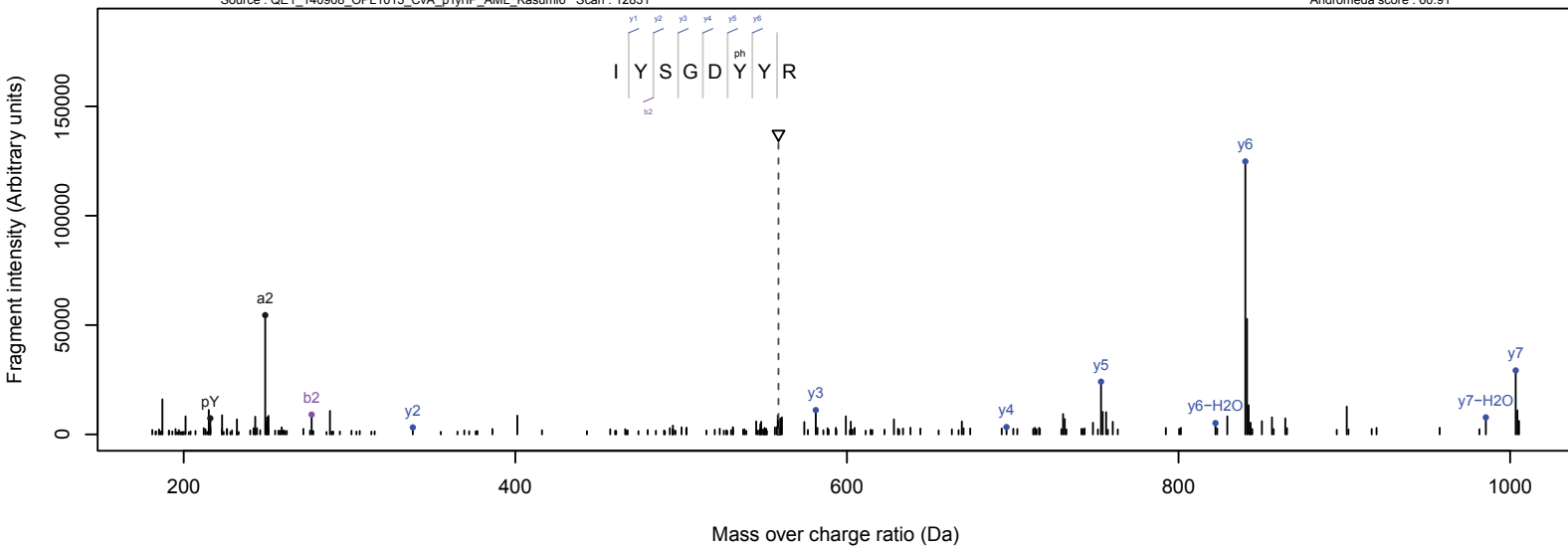
Andromeda score : 61.039



**IYSGDY(ph)YR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 12831

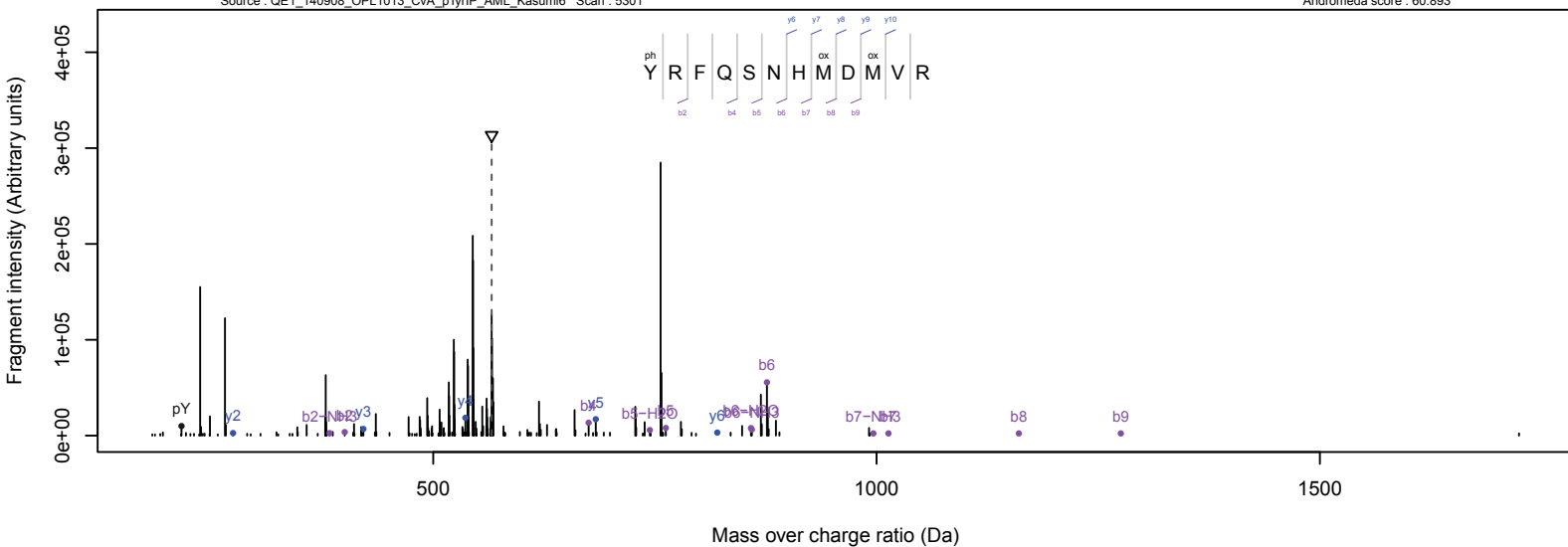
Andromeda score : 60.91



## Y(ph)RFQSNHM(ox)DM(ox)VR-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan: 5301

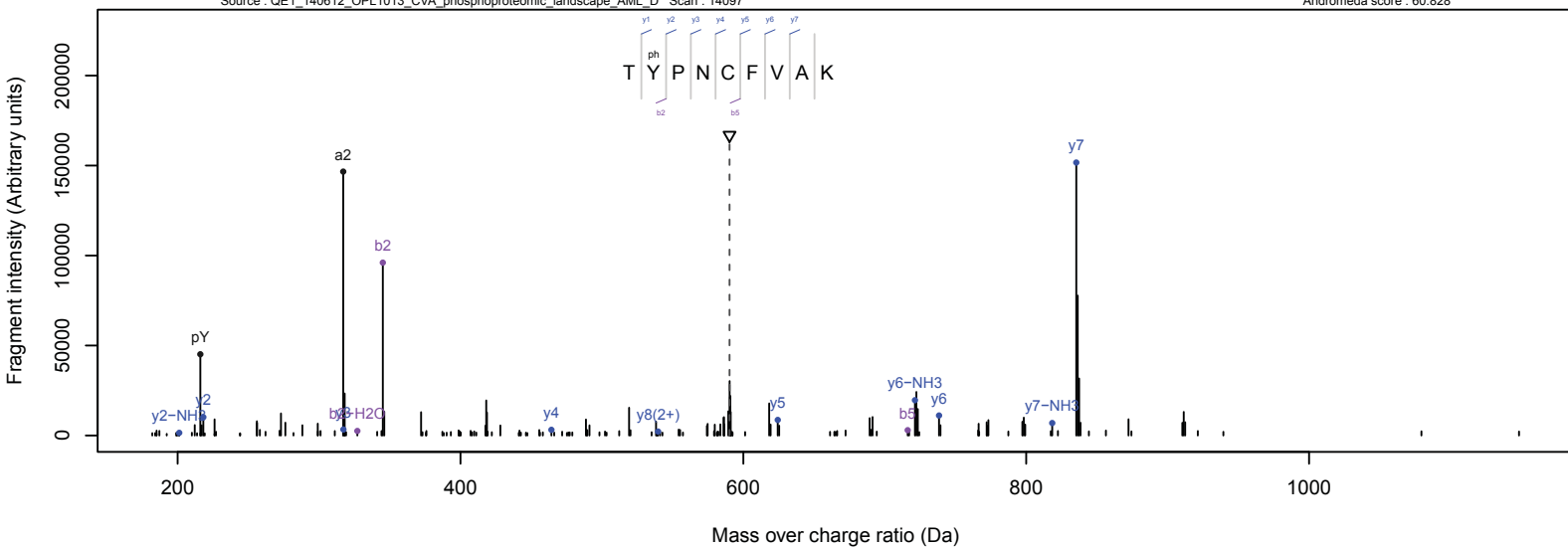
Andromeda score: 60.893



## TY(ph)PNCFVAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 14097

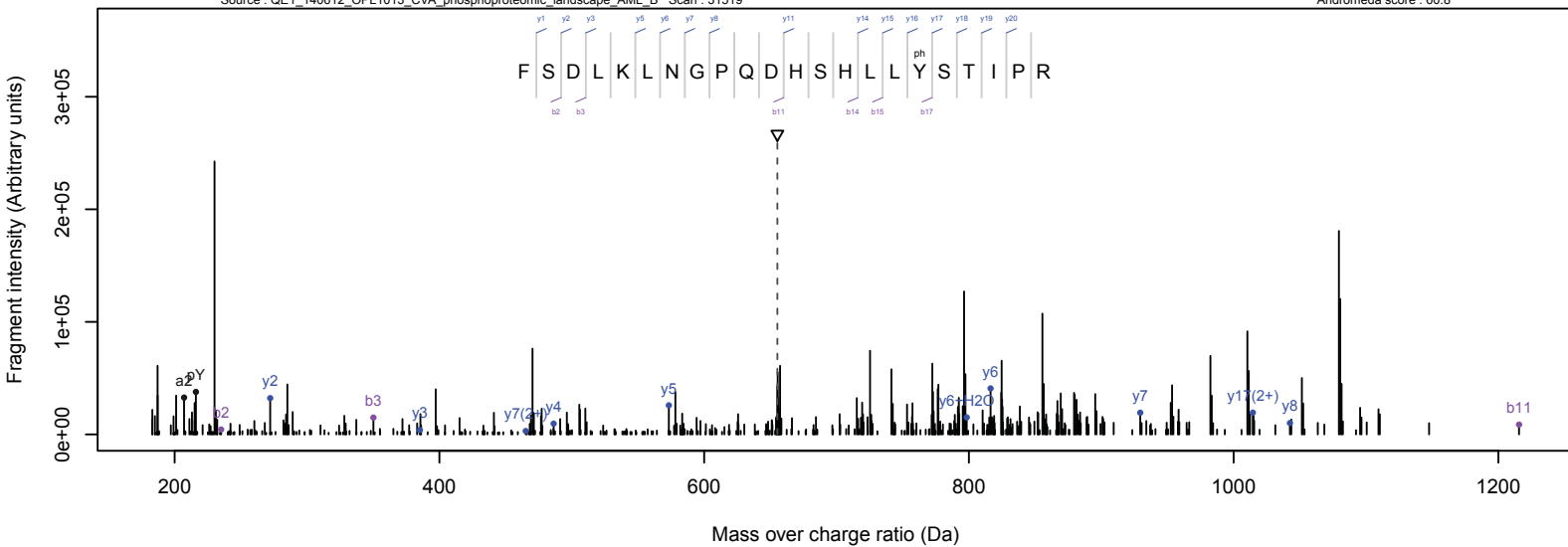
Andromeda score: 60.828



## FSDLKLNPGQDHSLLY(ph)STIPR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 31519

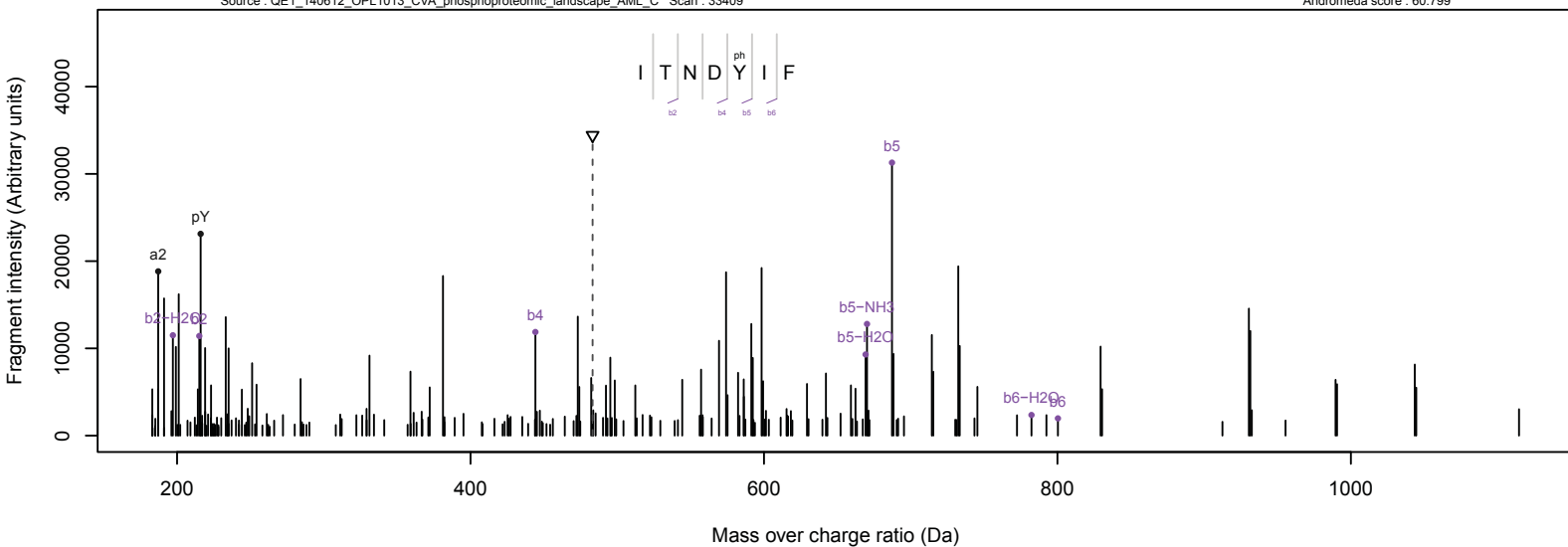
Andromeda score: 60.8



ITNDY(ph)F-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33409

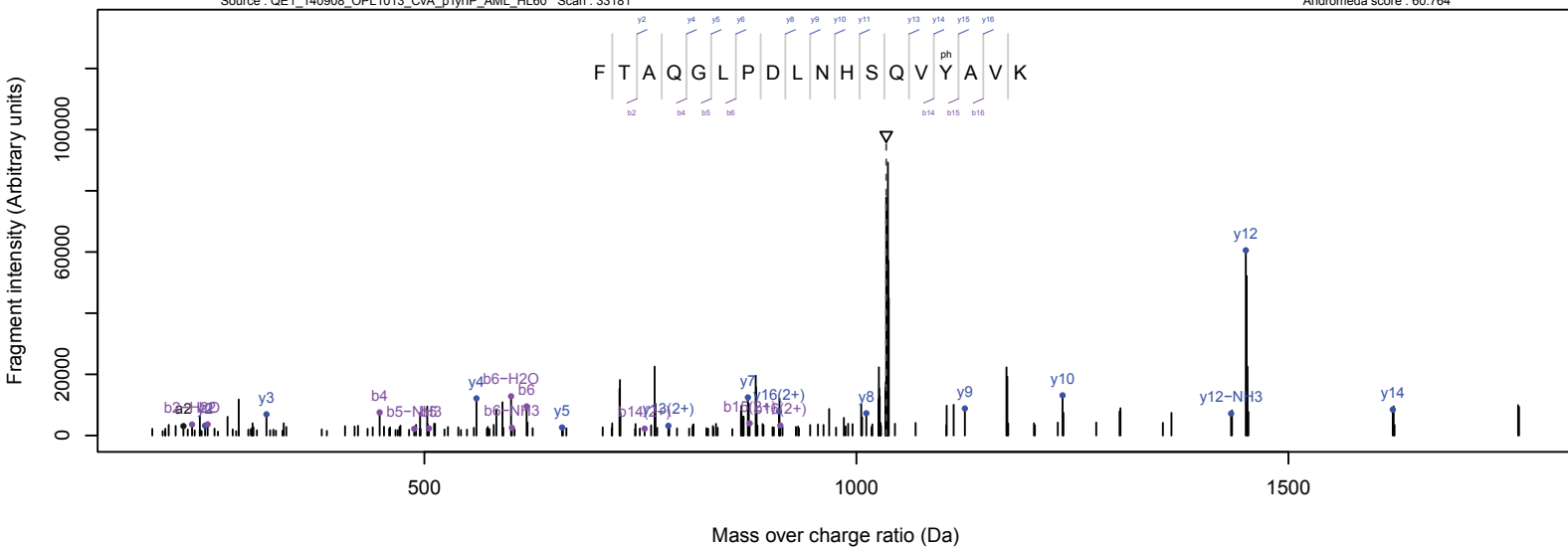
Andromeda score : 60.799



FTAQGLPDLNHSQV(ph)AVK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 33181

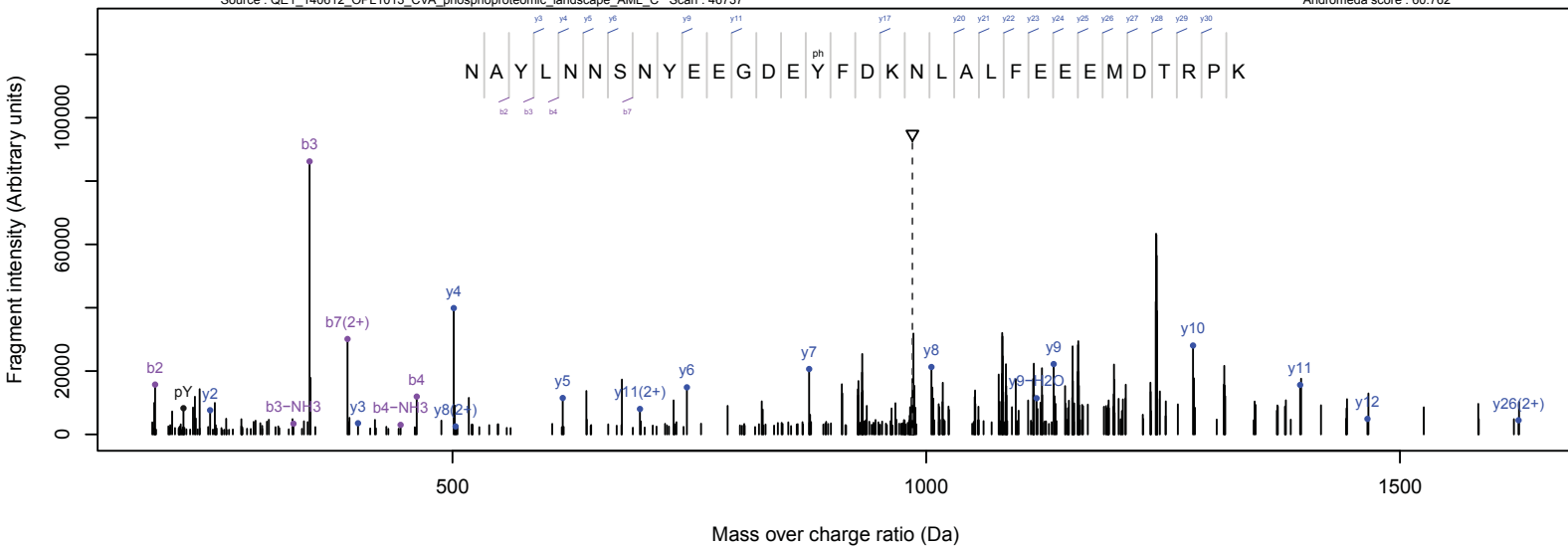
Andromeda score : 60.764



NAYLNNSEEGDEY(ph)FDKNLALFEEEMDTRPK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46737

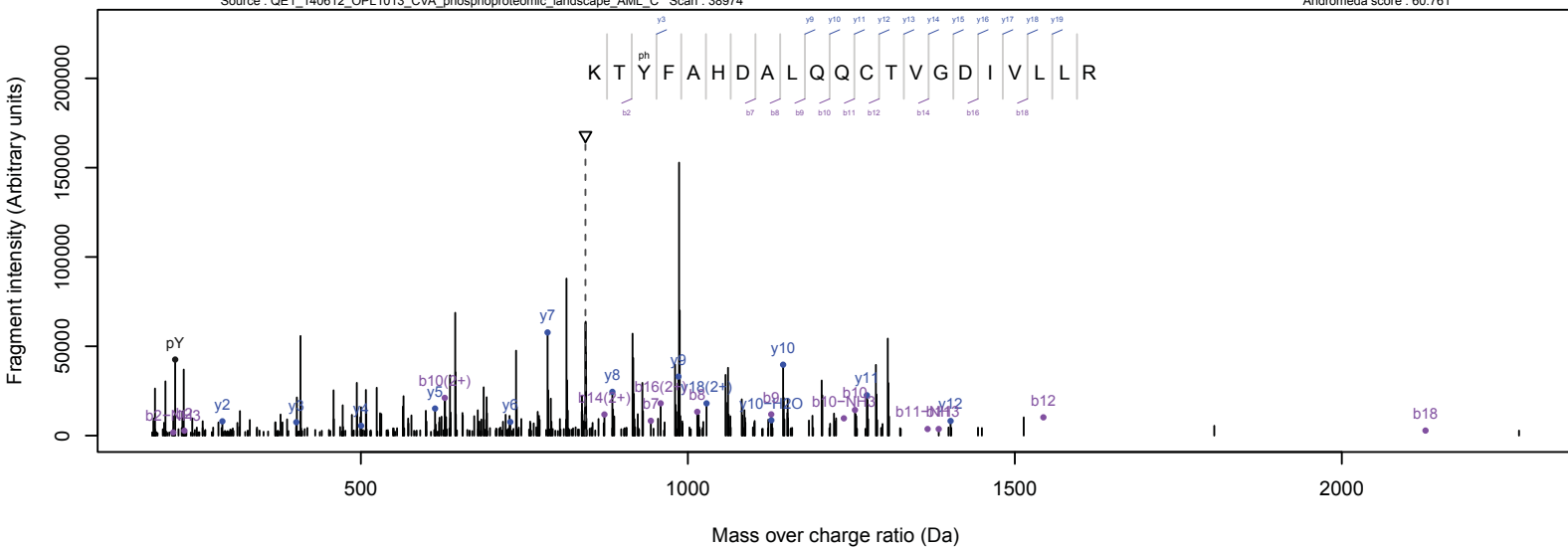
Andromeda score : 60.762



KTY(ph)FAHDALQQCTVGDIVLLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38974

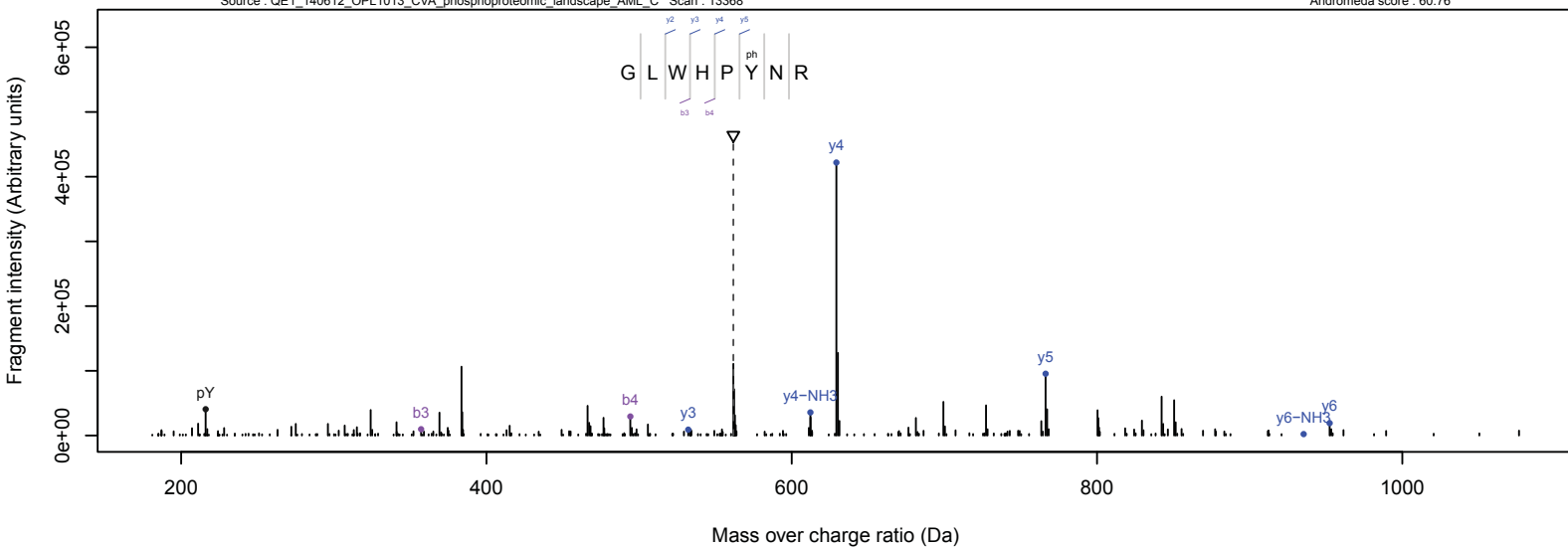
Andromeda score : 60.761



GLWHPY(ph)NR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13368

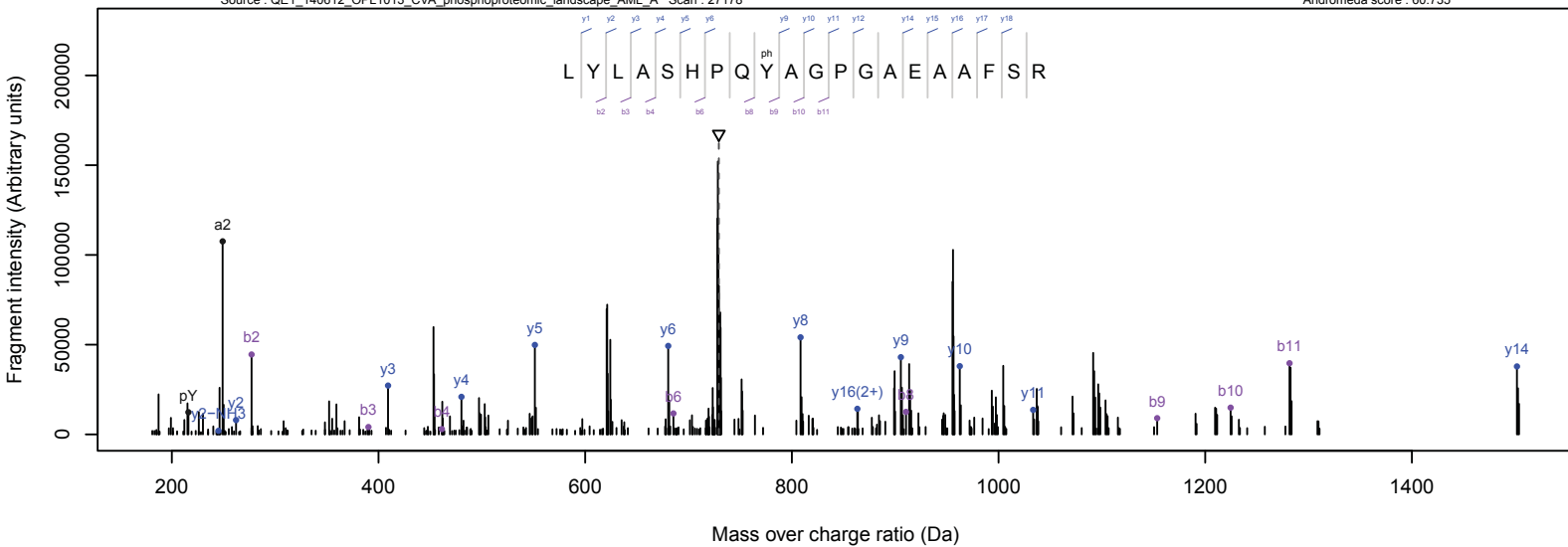
Andromeda score : 60.76



LYLASHPQ(ph)AGPGAEEAFSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27178

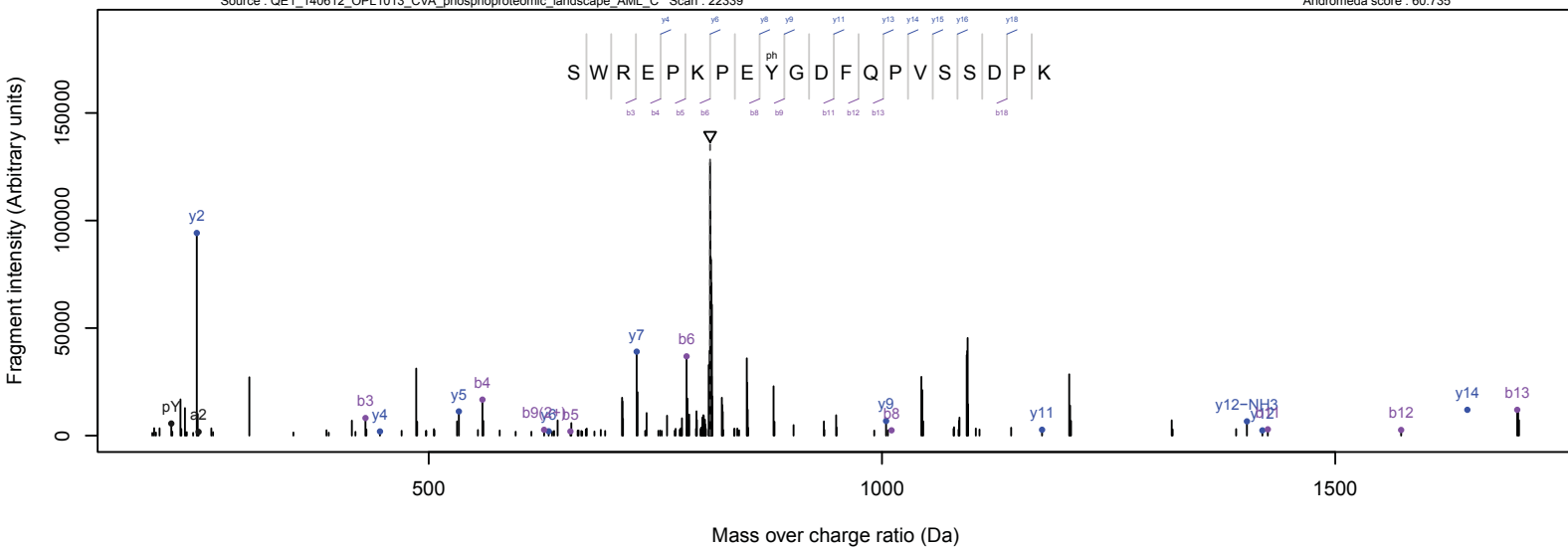
Andromeda score : 60.735



## SWREPKPEY(ph)GDFQPVSSDPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22339

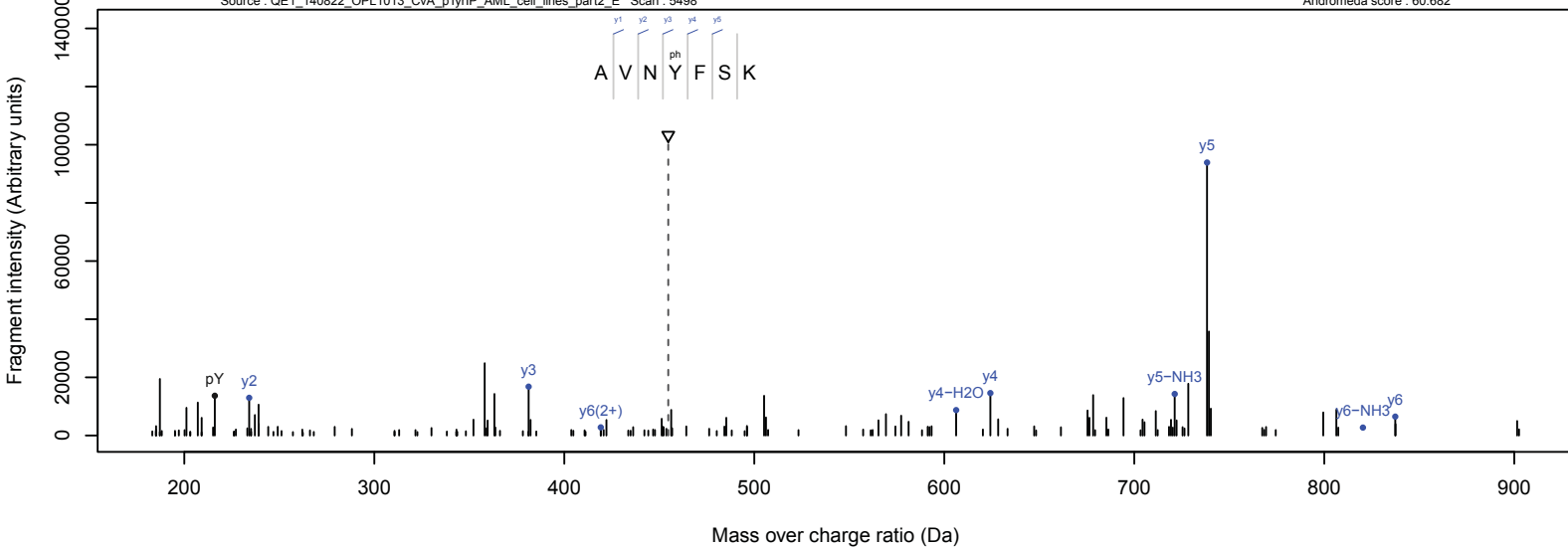
Andromeda score : 60.735



## AVNY(ph)FSK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 5498

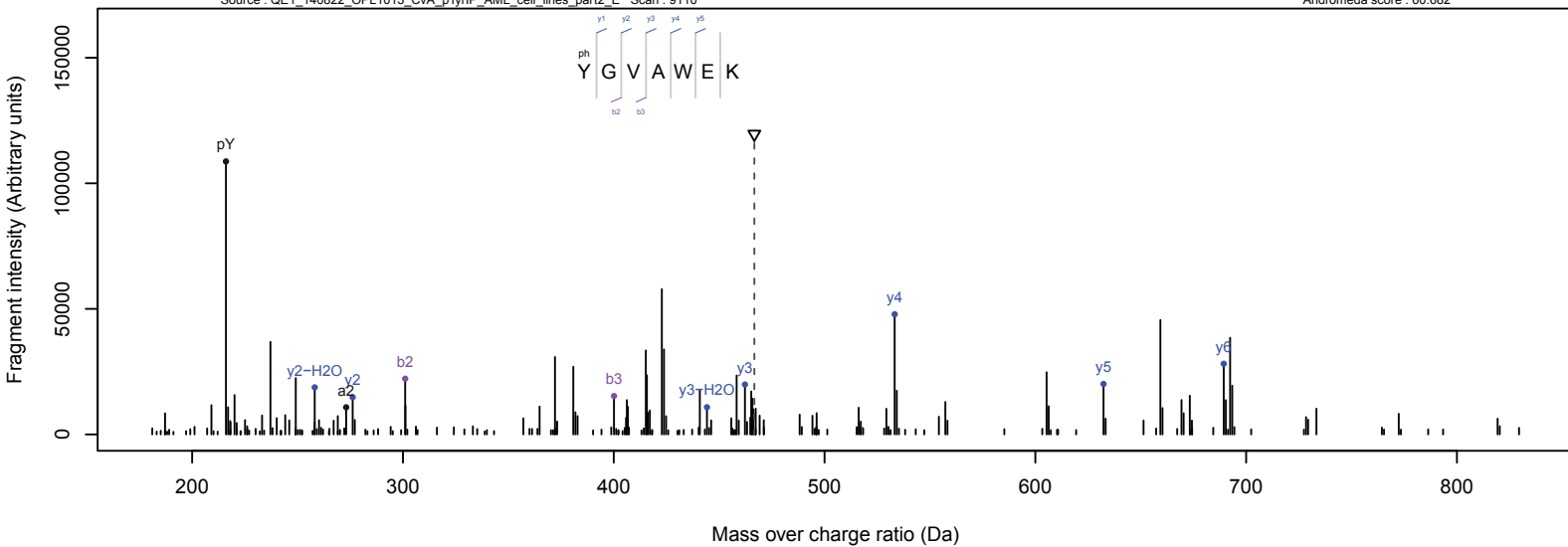
Andromeda score : 60.682



## Y(ph)GVAWEK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 9110

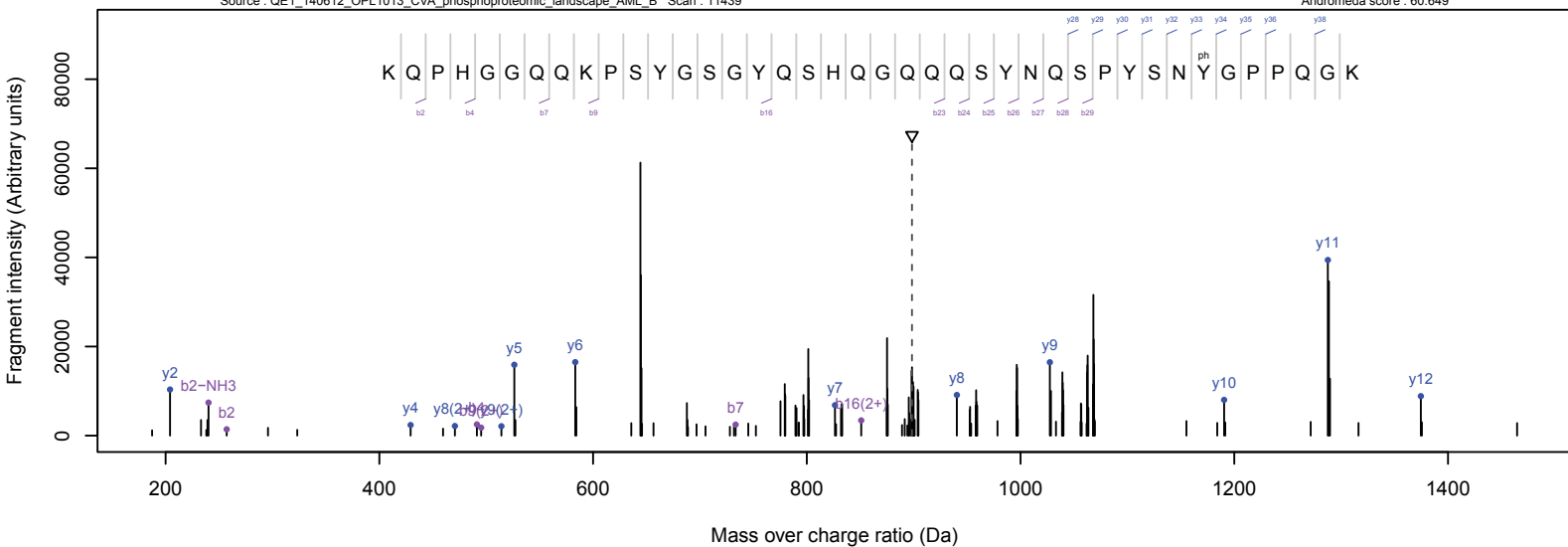
Andromeda score : 60.682



**KQPHGGQKPSYSGYQSHQGQQSYNQSPYSNY(ph)GPPQGK-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11439

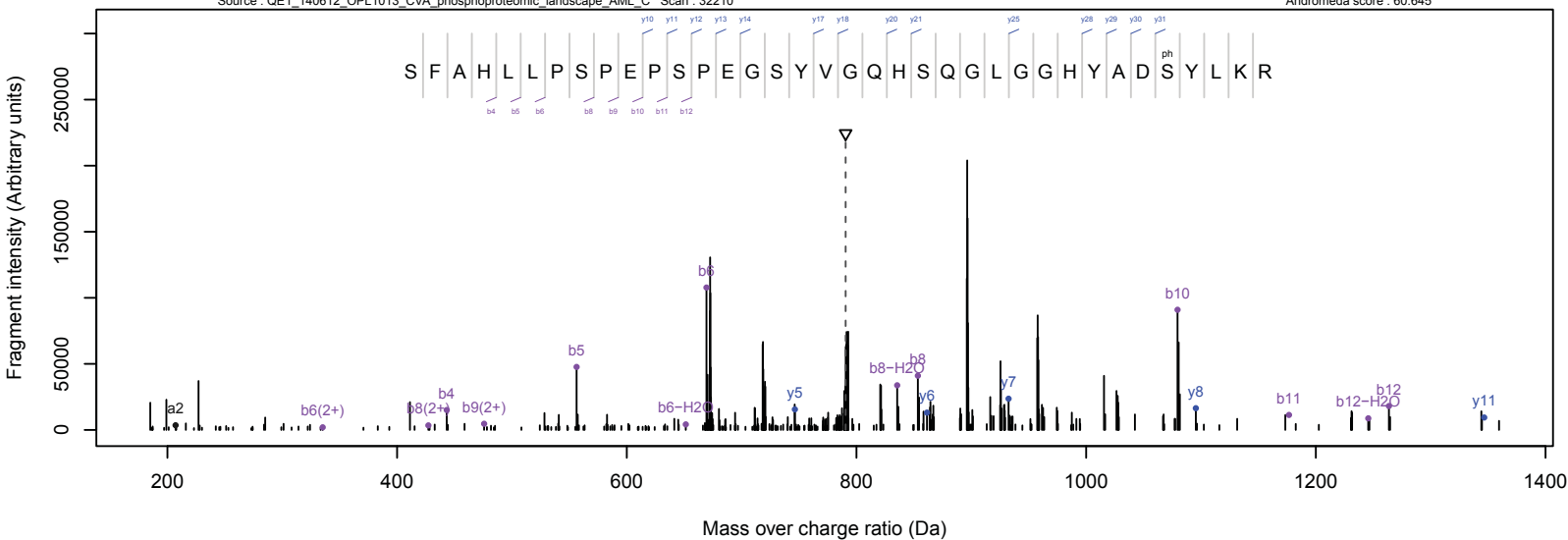
Andromeda score : 60.649



**SFAHLLPSEPSPEGSYVGQHSQGLGGHYADS(ph)YLKR-5+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32210

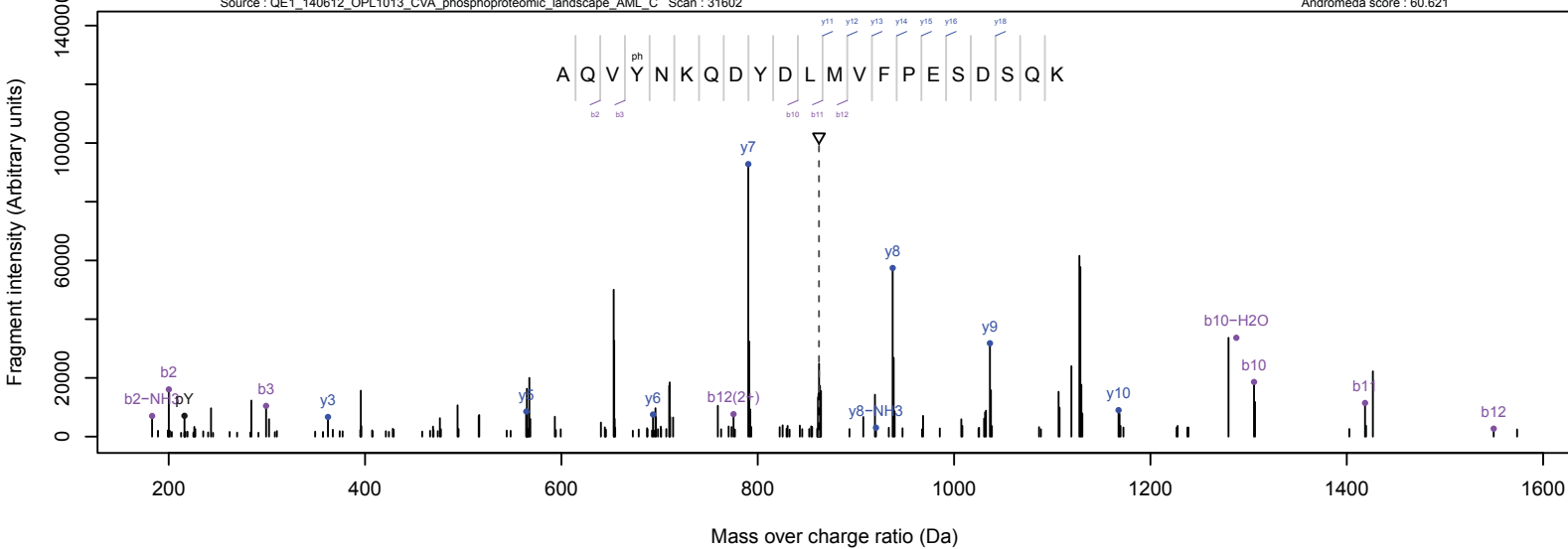
Andromeda score : 60.645

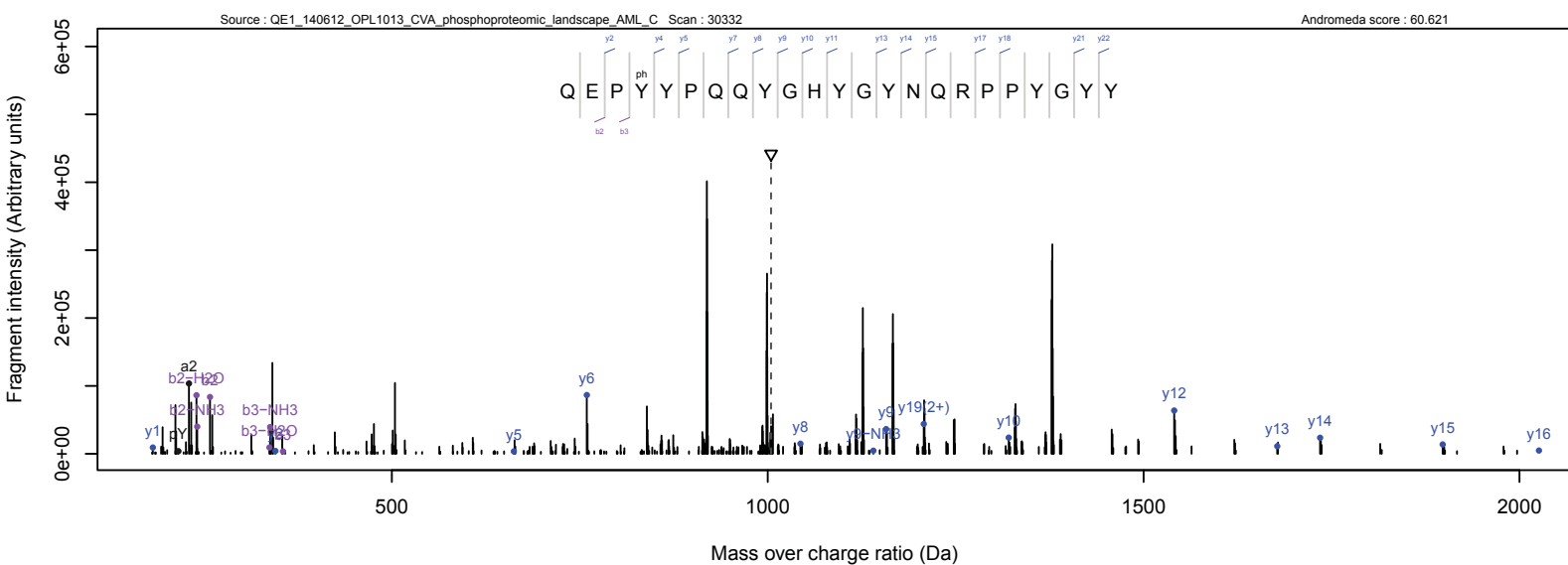
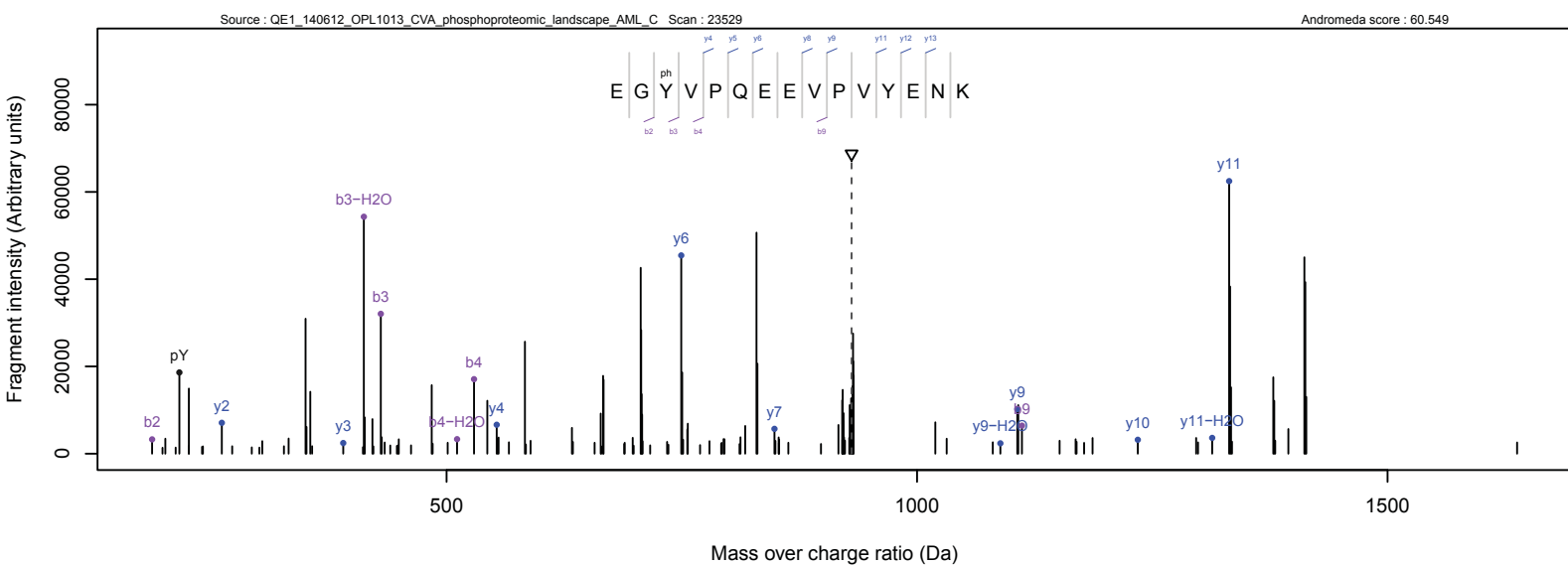
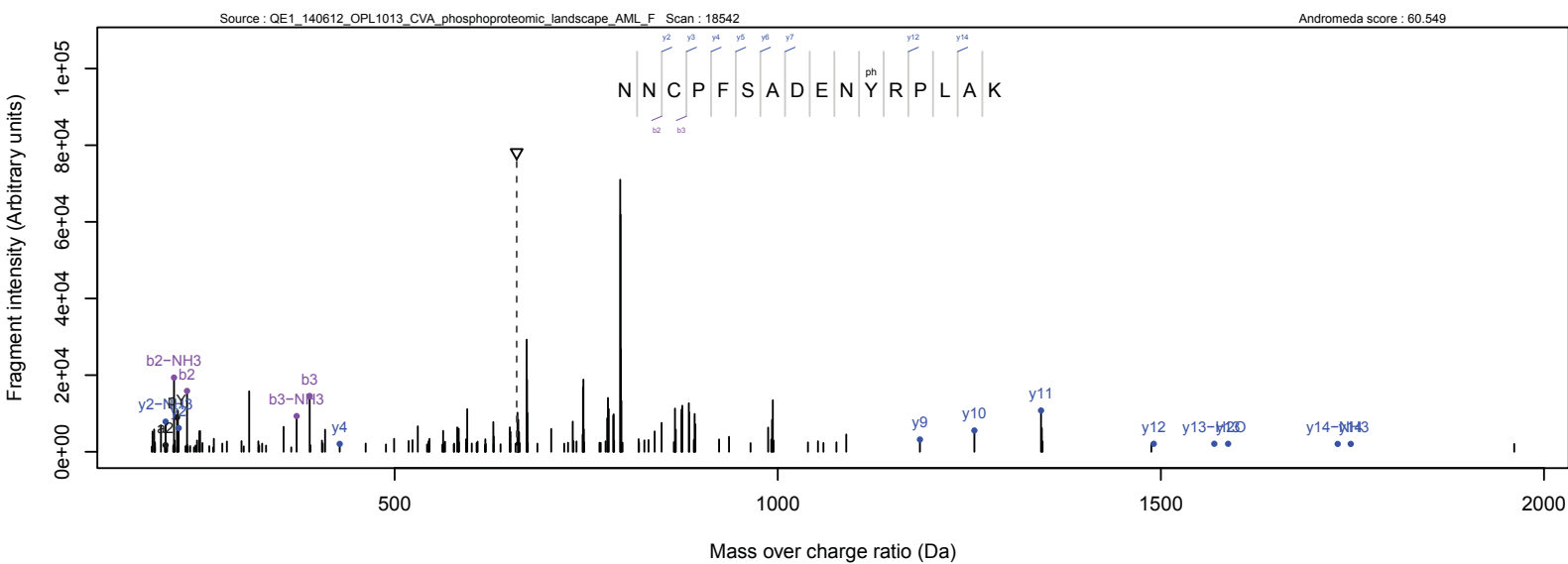


**AQVY(ph)NKQDYDLMVFPESDSQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31602

Andromeda score : 60.621



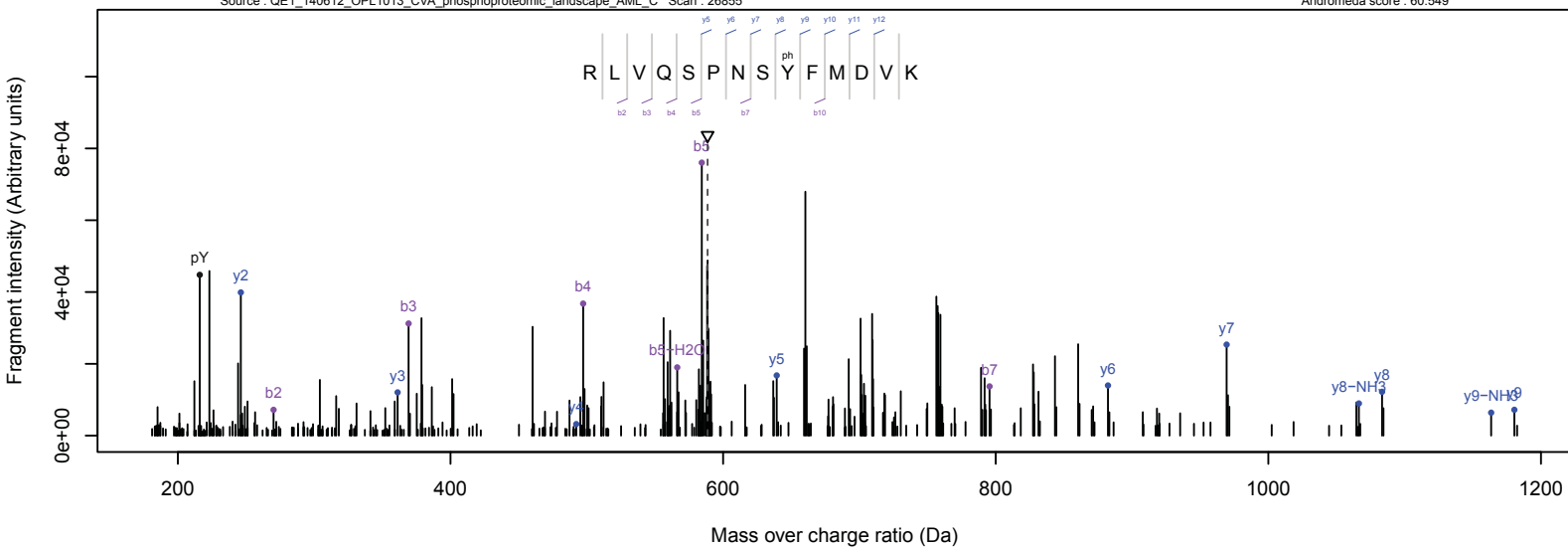
QE<sup>ph</sup>YPQYGHYGNQRPPYGY-3+EGY<sup>ph</sup>VPQEEVPVYENK-2+NNCPFSADENY<sup>ph</sup>RPLAK-3+



RLVQSPNS(ph)FMDVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26855

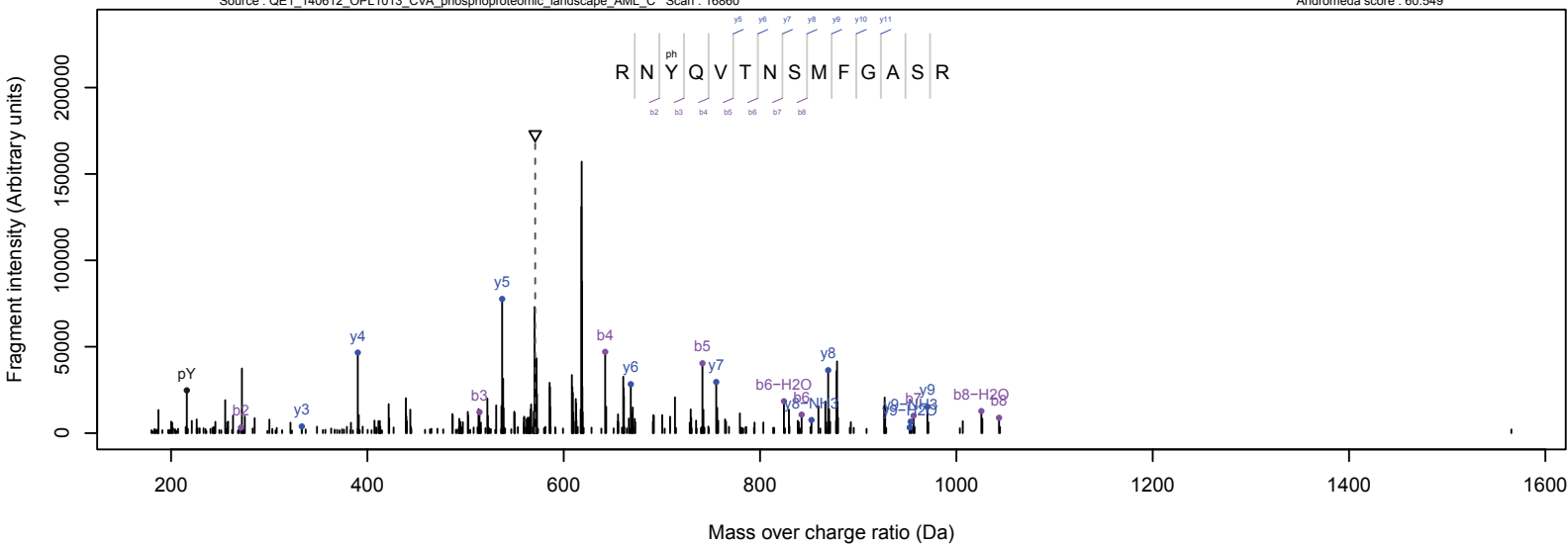
Andromeda score : 60.549



RNY(ph)QVTNSMFGASR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16860

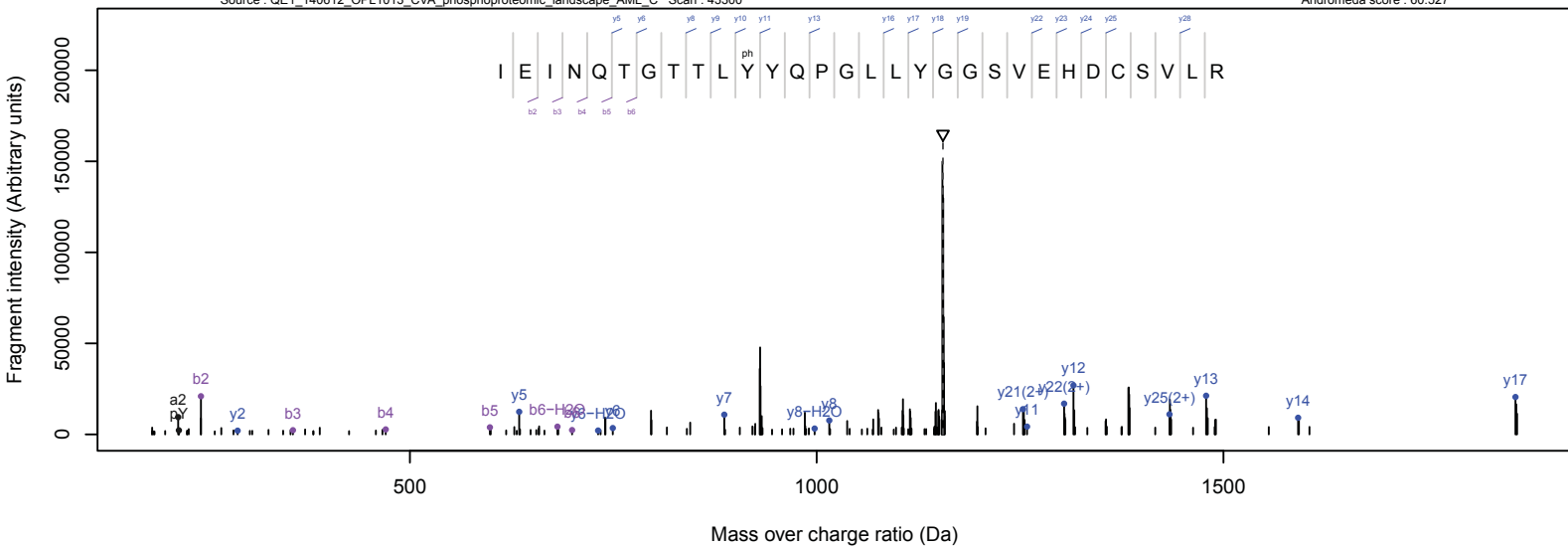
Andromeda score : 60.549



IEINQTTLY(ph)YQPLLGGVSEHDCSVLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43300

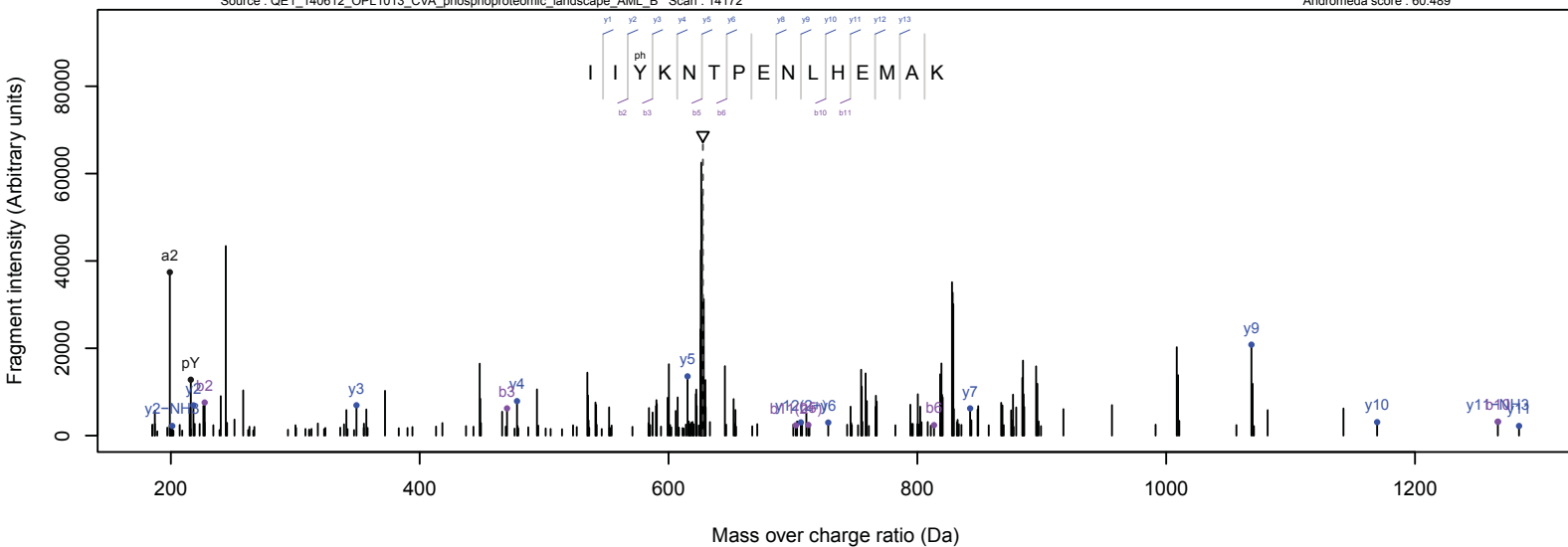
Andromeda score : 60.527



## IIY(ph)KNTPENLHEMAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14172

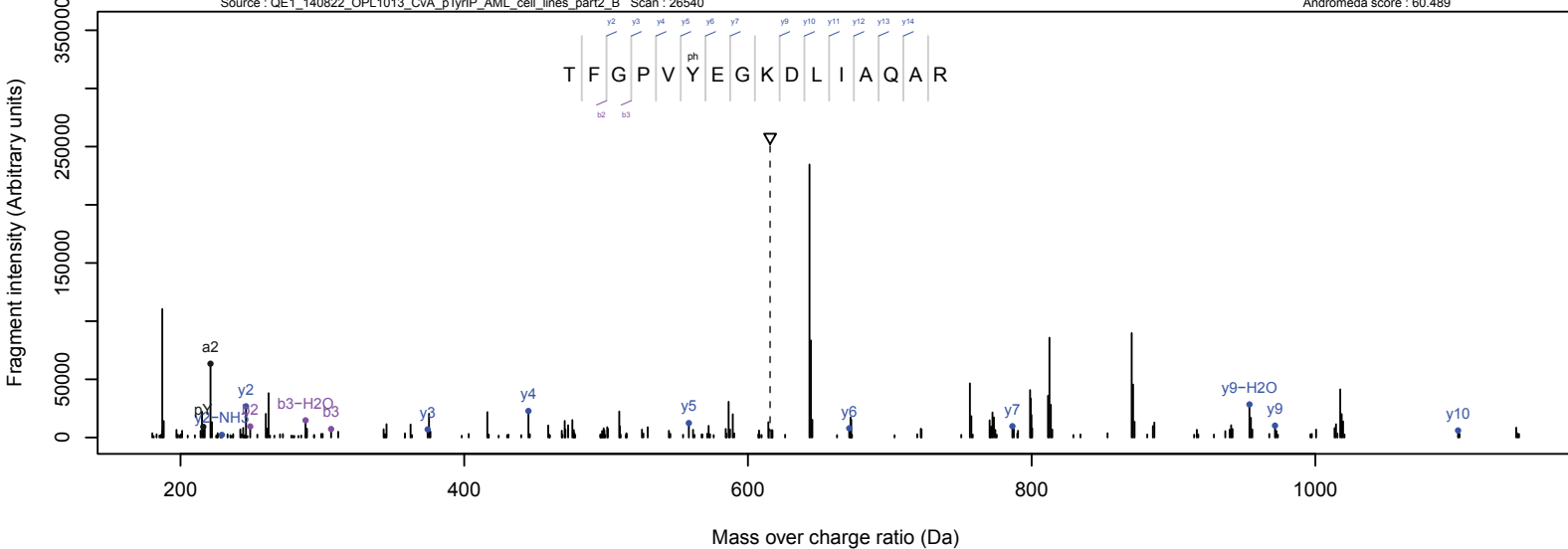
Andromeda score : 60.489



## TFGPVY(ph)EGKDLIAQAR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 26540

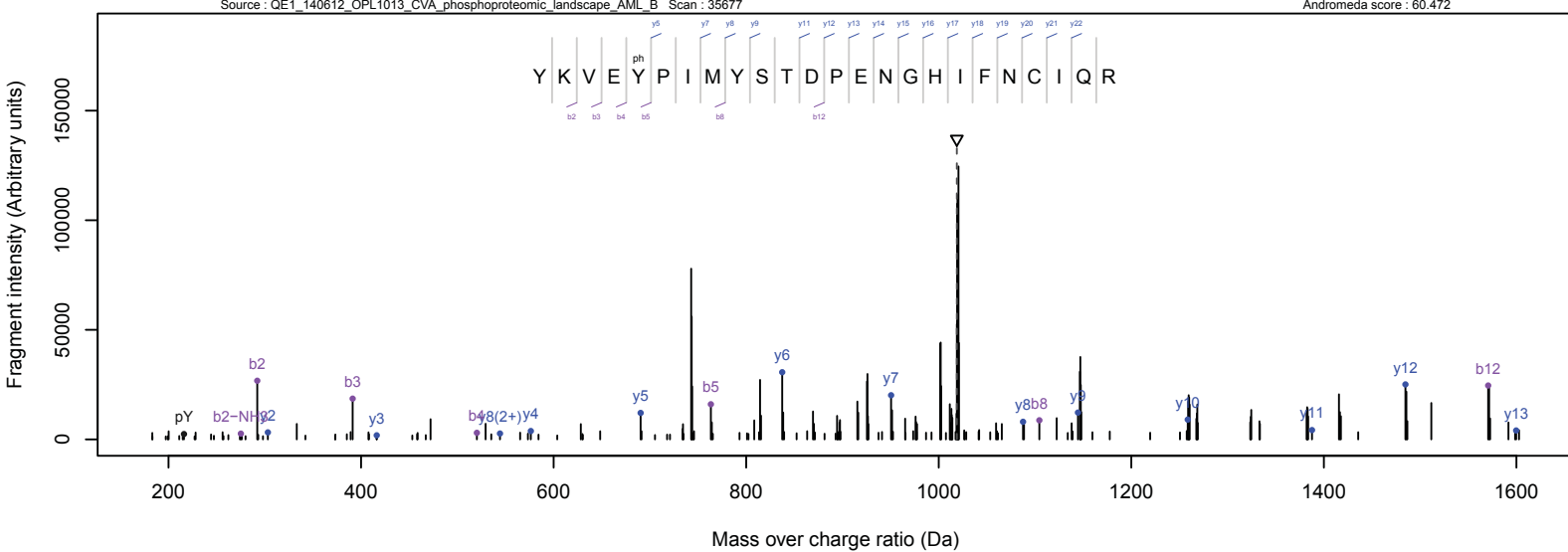
Andromeda score : 60.489



## YKVEY(ph)PIMYSTDPENGHIFNCIQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35677

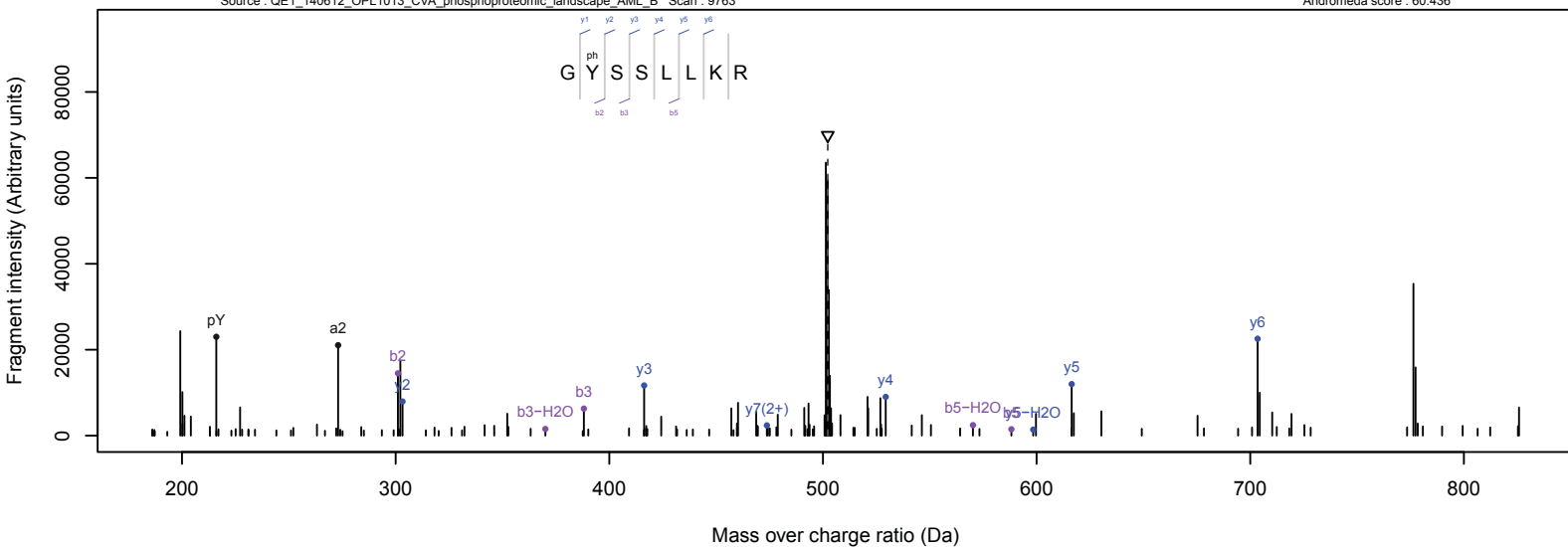
Andromeda score : 60.472



## GY(ph)SSLLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9763

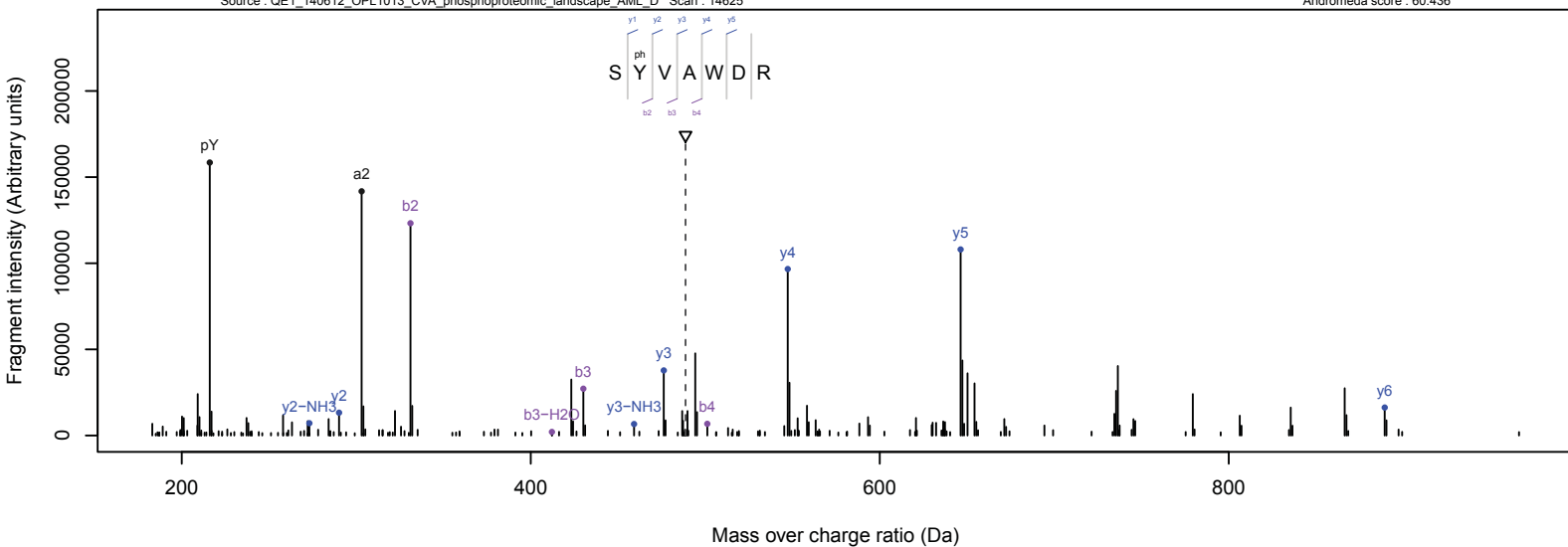
Andromeda score : 60.436



## SY(ph)VAWDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14625

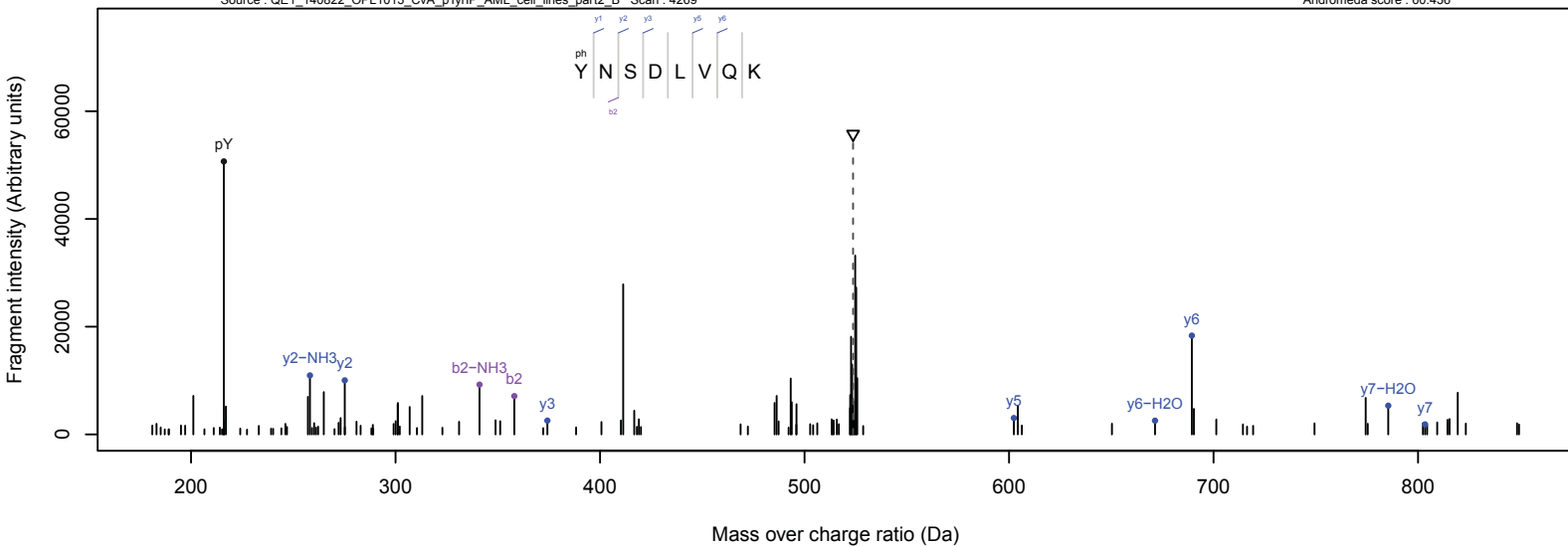
Andromeda score : 60.436



## Y(ph)NSDLVQK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 4269

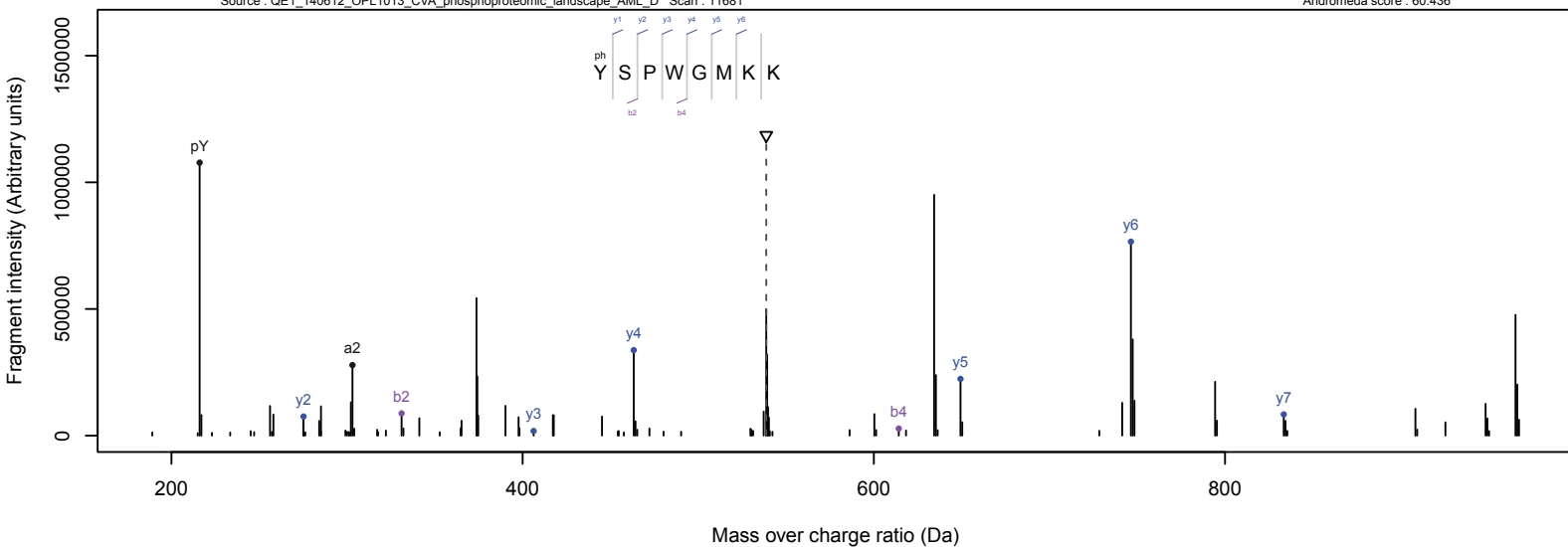
Andromeda score : 60.436



Y(ph)SPWGMKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 11681

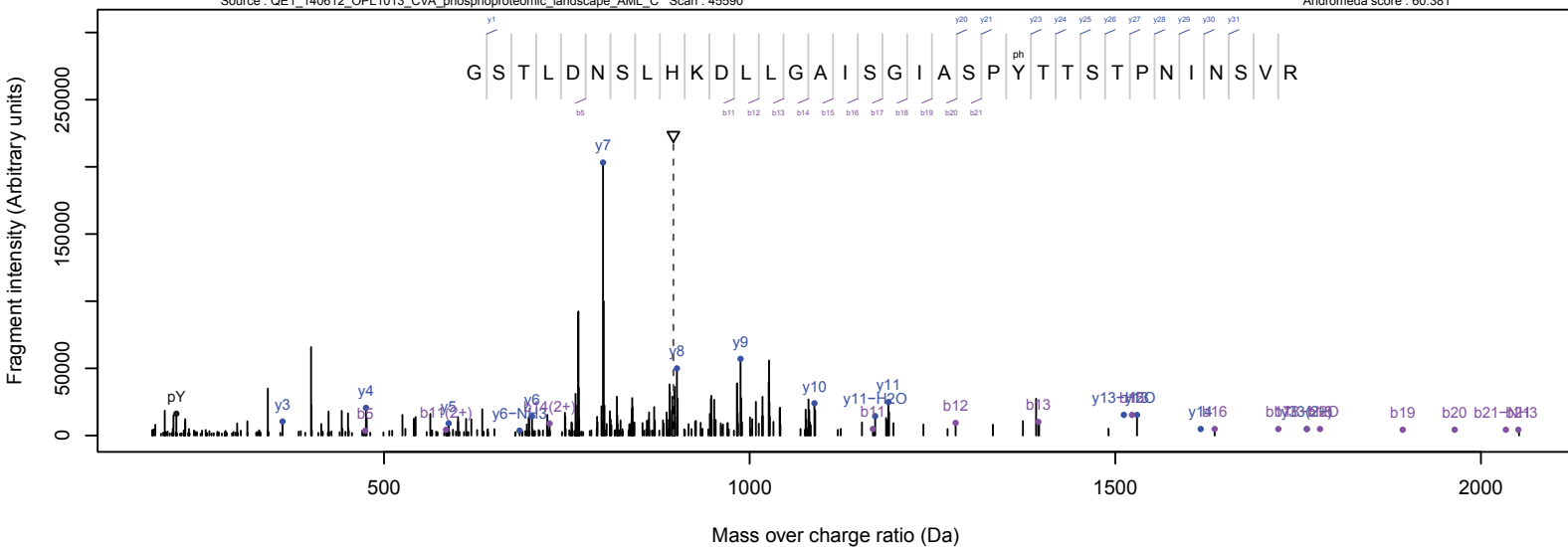
Andromeda score : 60.436



GSTLDNSLHKDLLGAISGIASPY(ph)TTSTPNINSVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45590

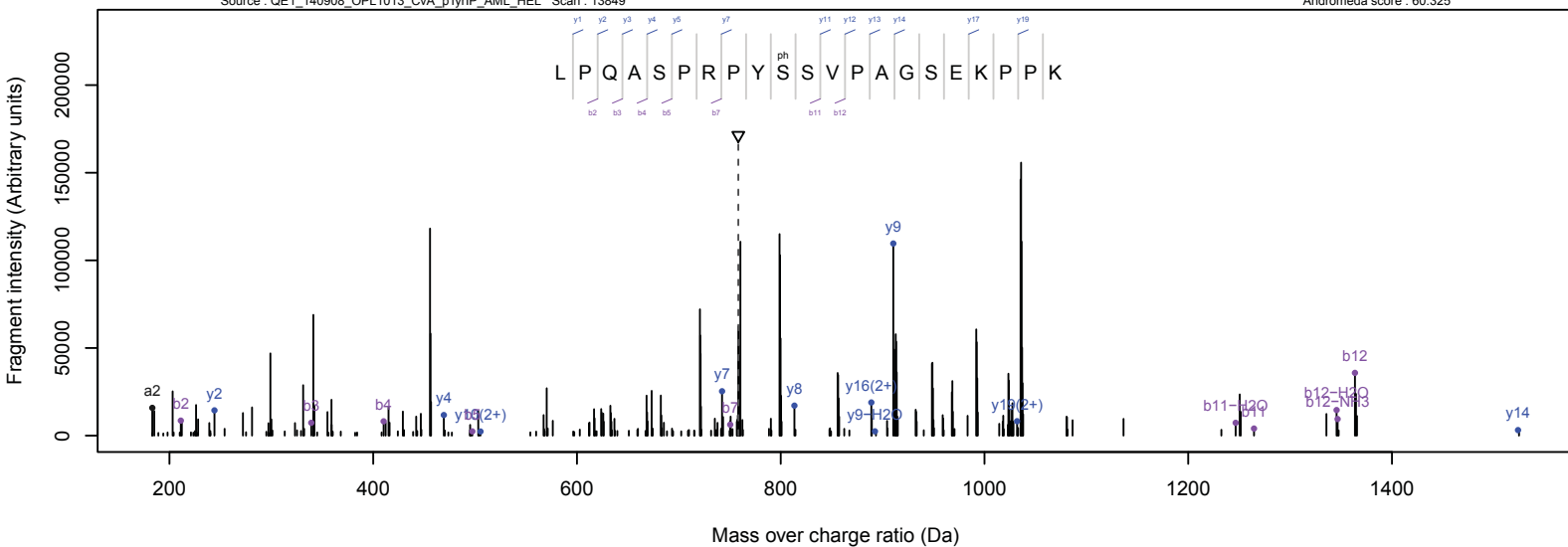
Andromeda score : 60.381



LPQASRPYS(ph)SVPAGSEKPPK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 13849

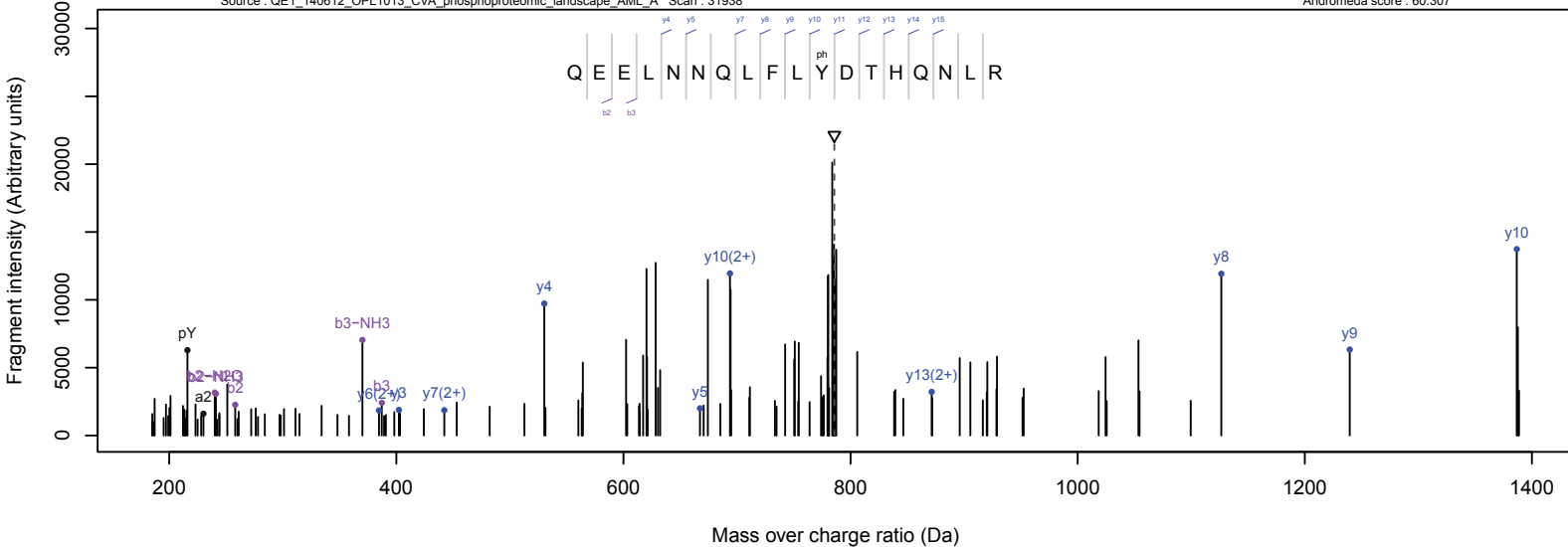
Andromeda score : 60.325



QEELNNQLFLY(ph)DTHQNLRL-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 31938

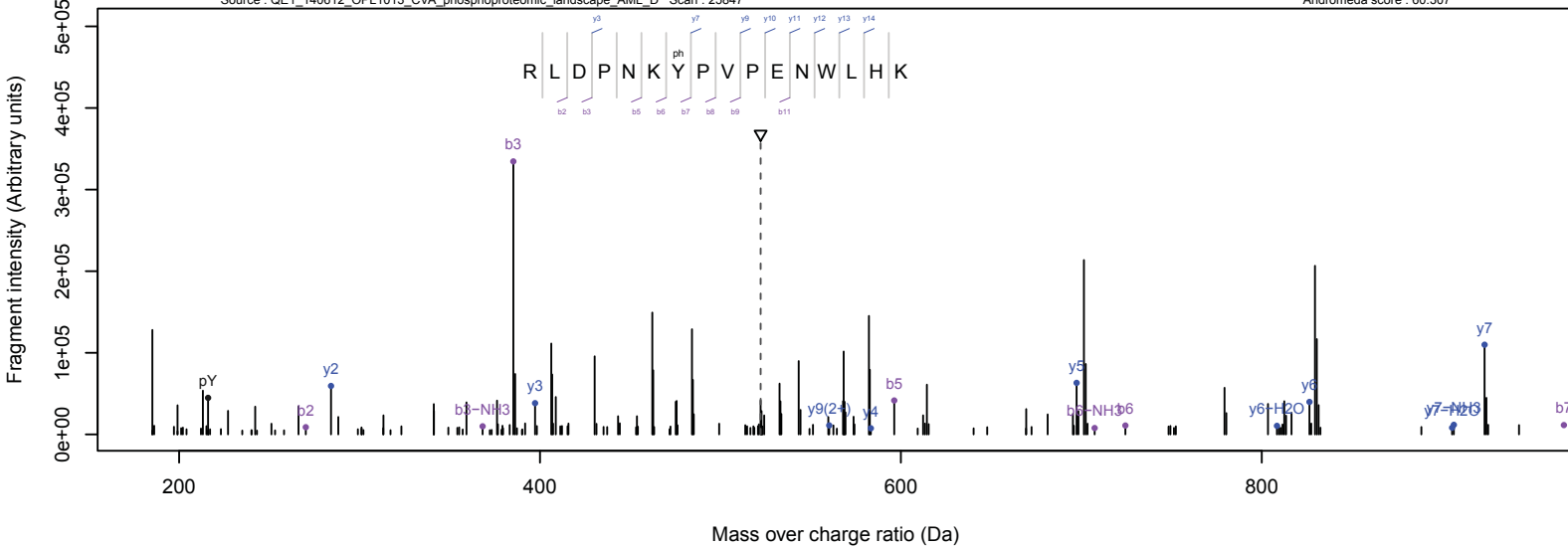
Andromeda score : 60.307



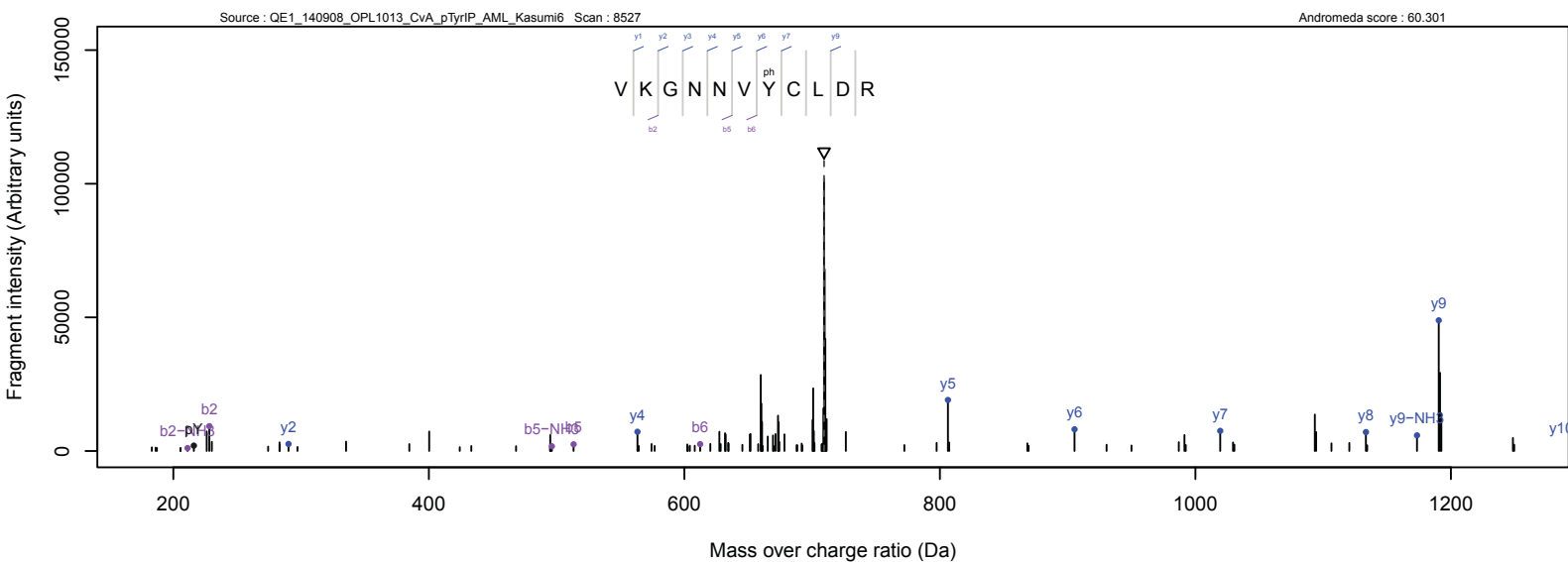
RLDPNKY(ph)PVPENWLHK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 25847

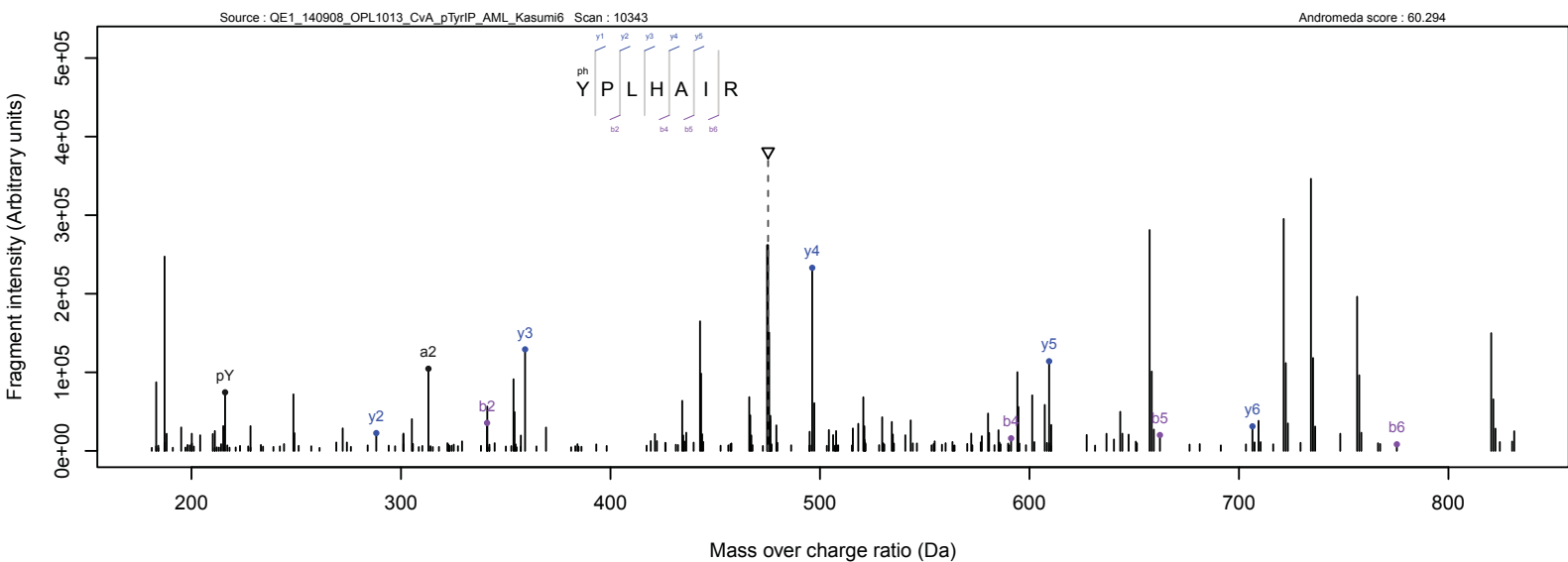
Andromeda score : 60.307



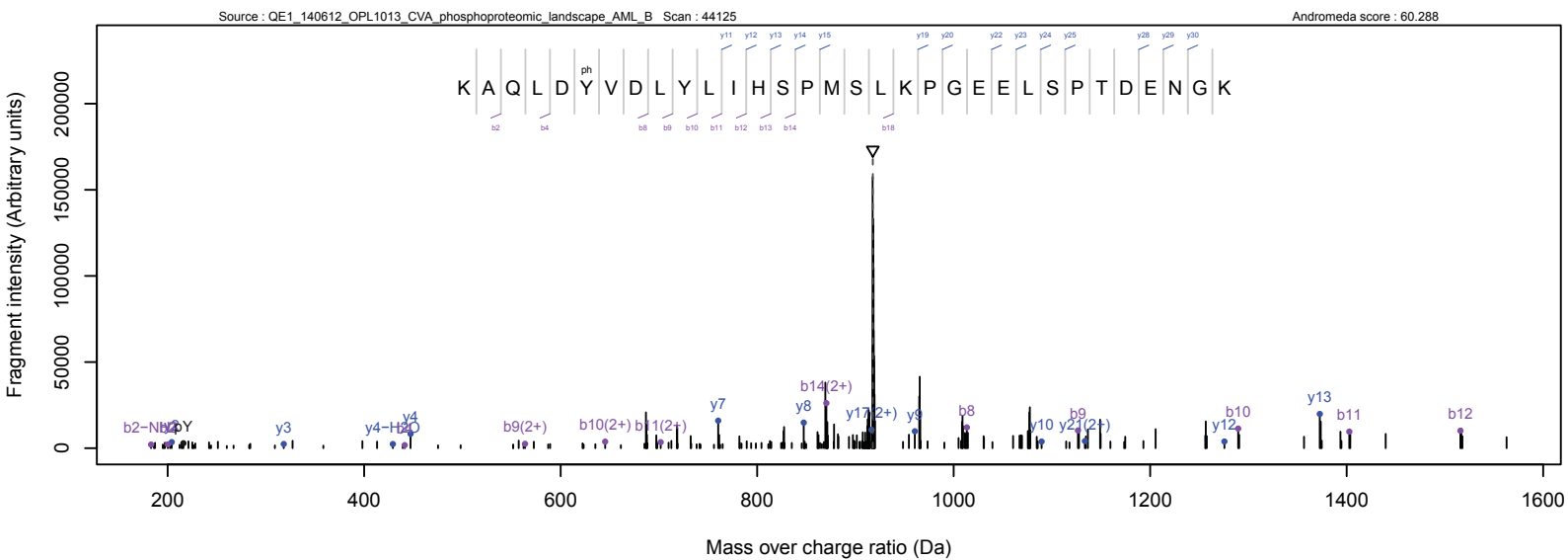
VKGNNVY(ph)CLDR-2+



Y(ph)PLHAIR-2+



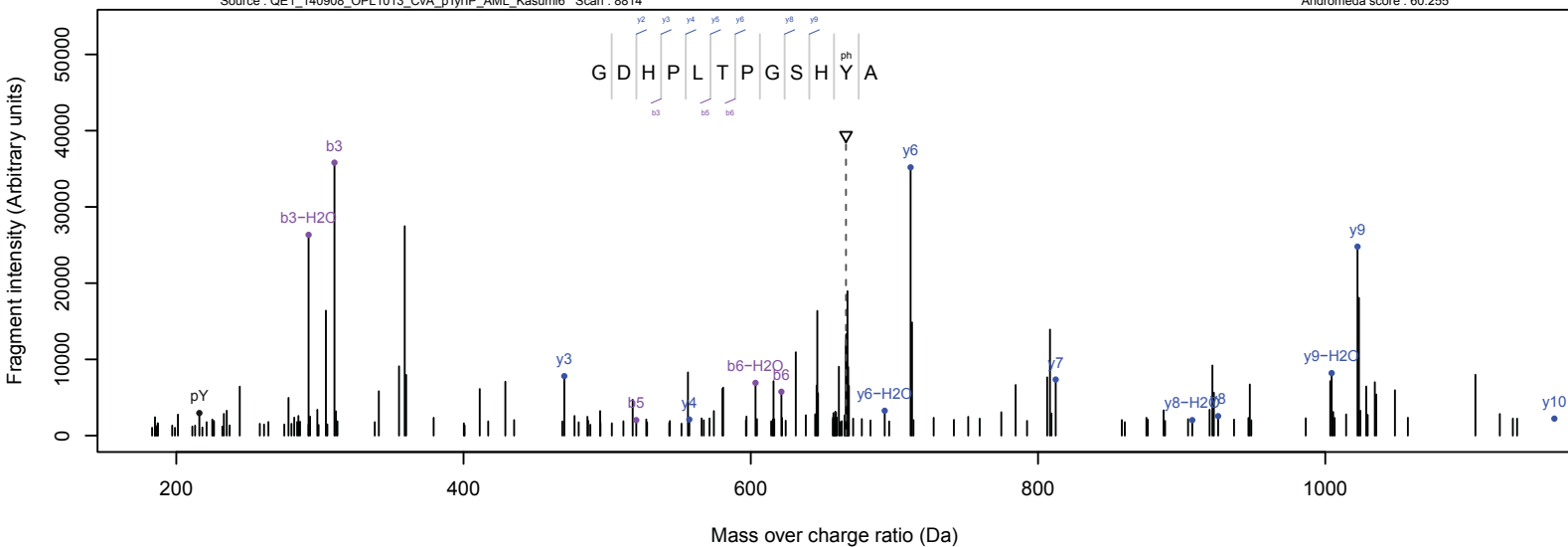
KAQLDY(ph)VDLYLIHSPMSLKPGEEELSPDENGK-4+



## GDHPLTPGSHY(ph)A-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 8814

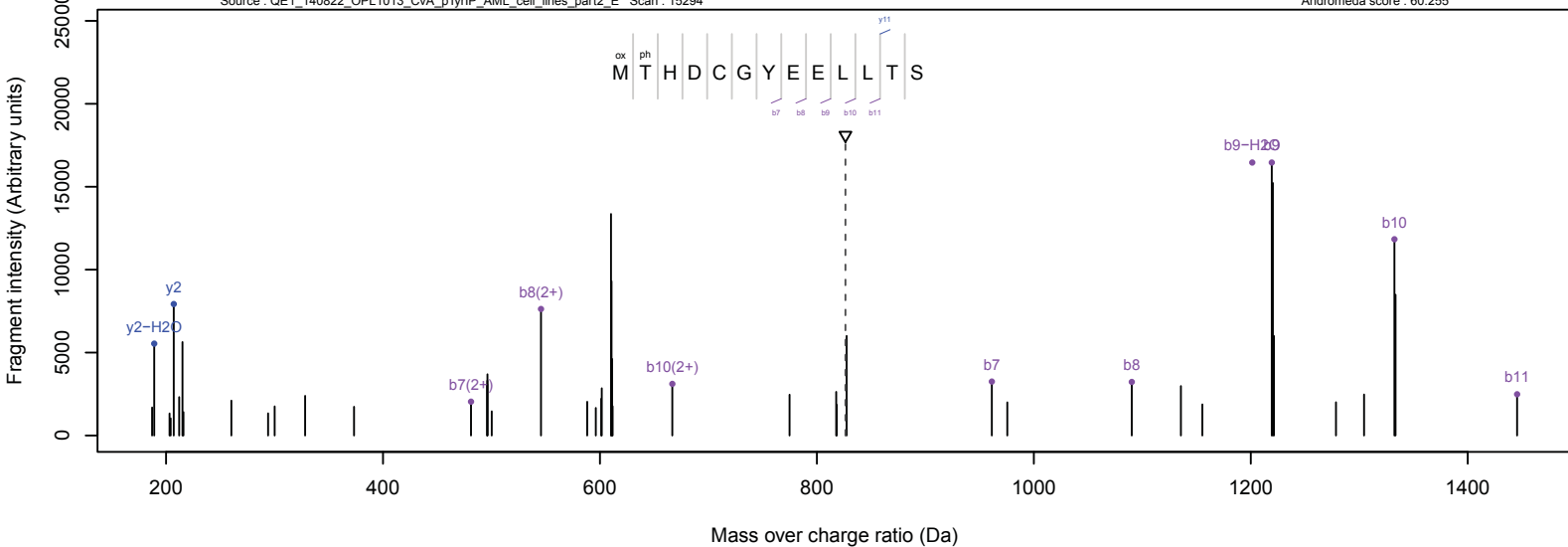
Andromeda score: 60.255



## M(ox)T(ph)HDCGYEELLTS-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 15294

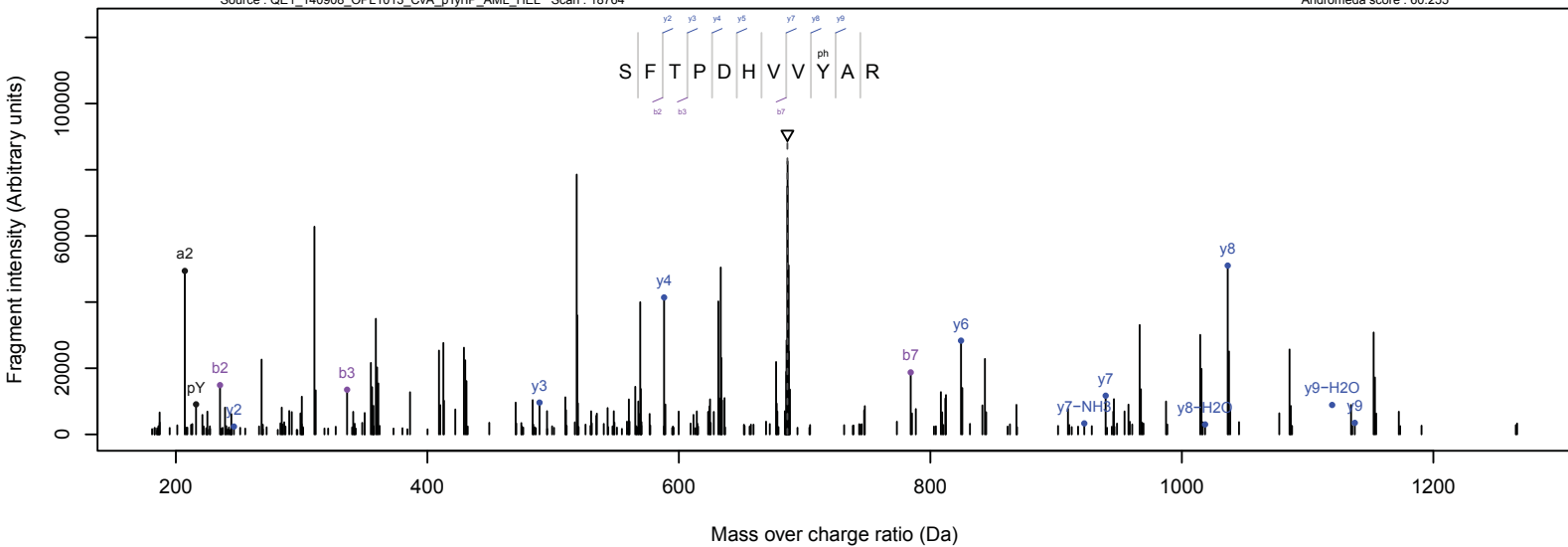
Andromeda score: 60.255



## SFTPDHVY(ph)AR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 18764

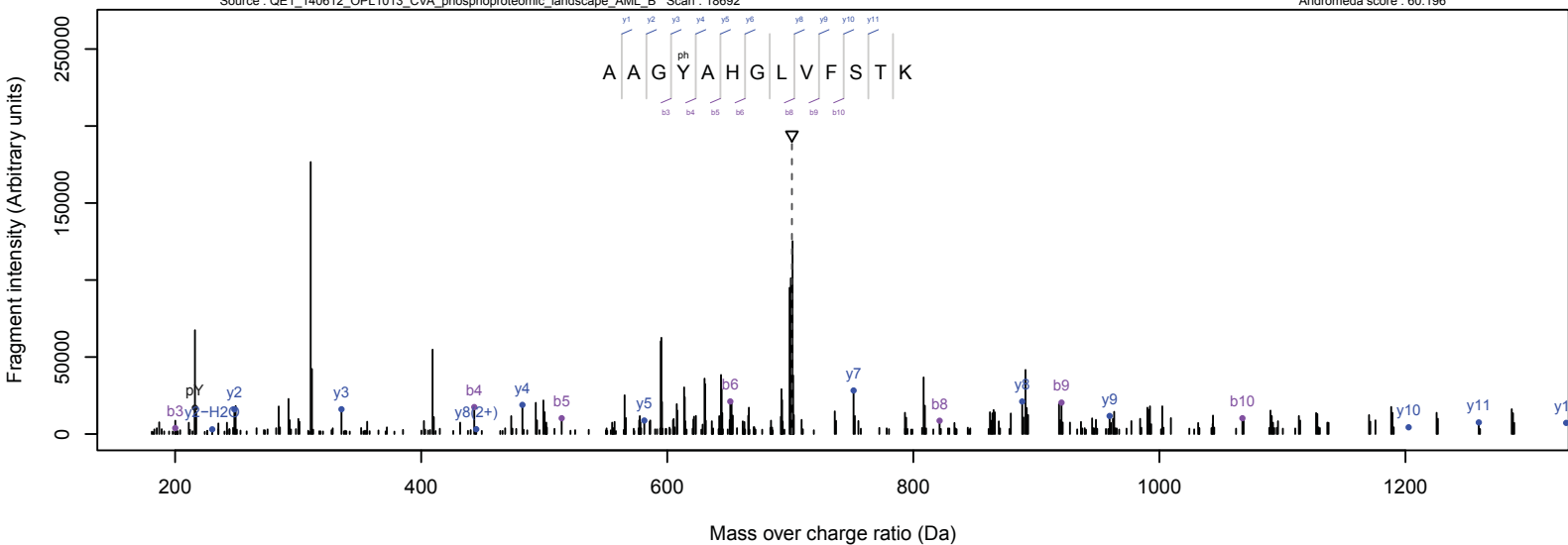
Andromeda score: 60.255



## AAGY(ph)AHGLVFSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18692

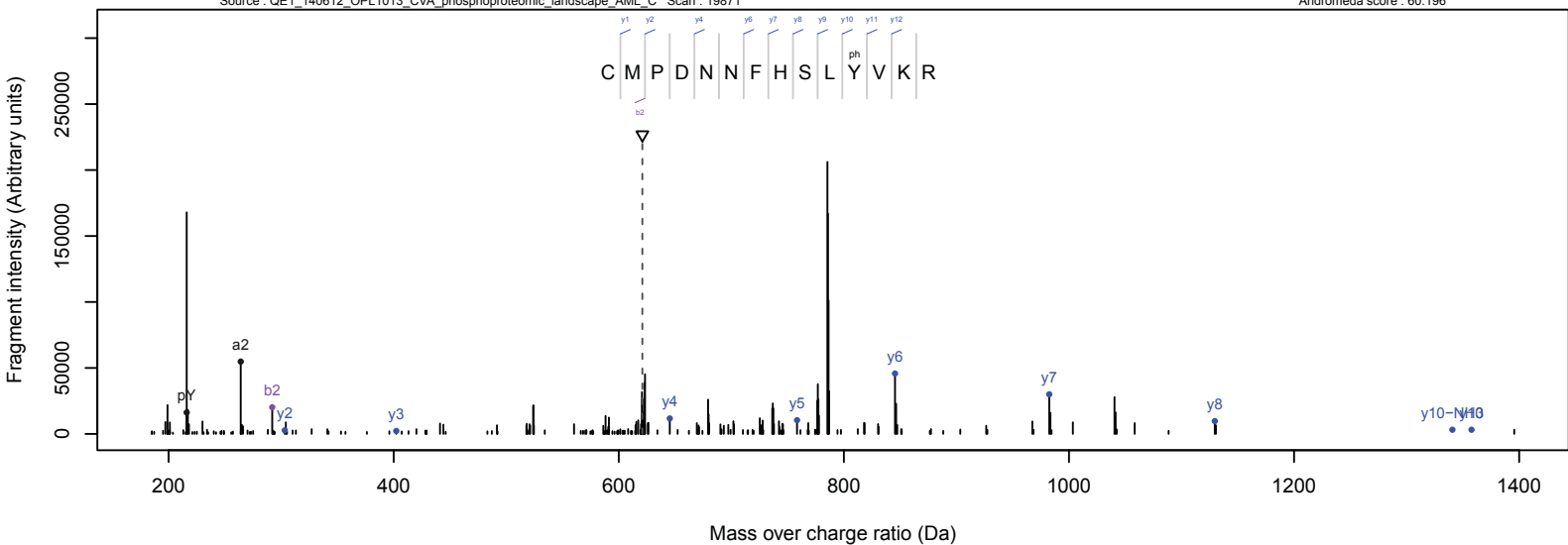
Andromeda score : 60.196



## CMPDNNFHSLY(ph)VKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19871

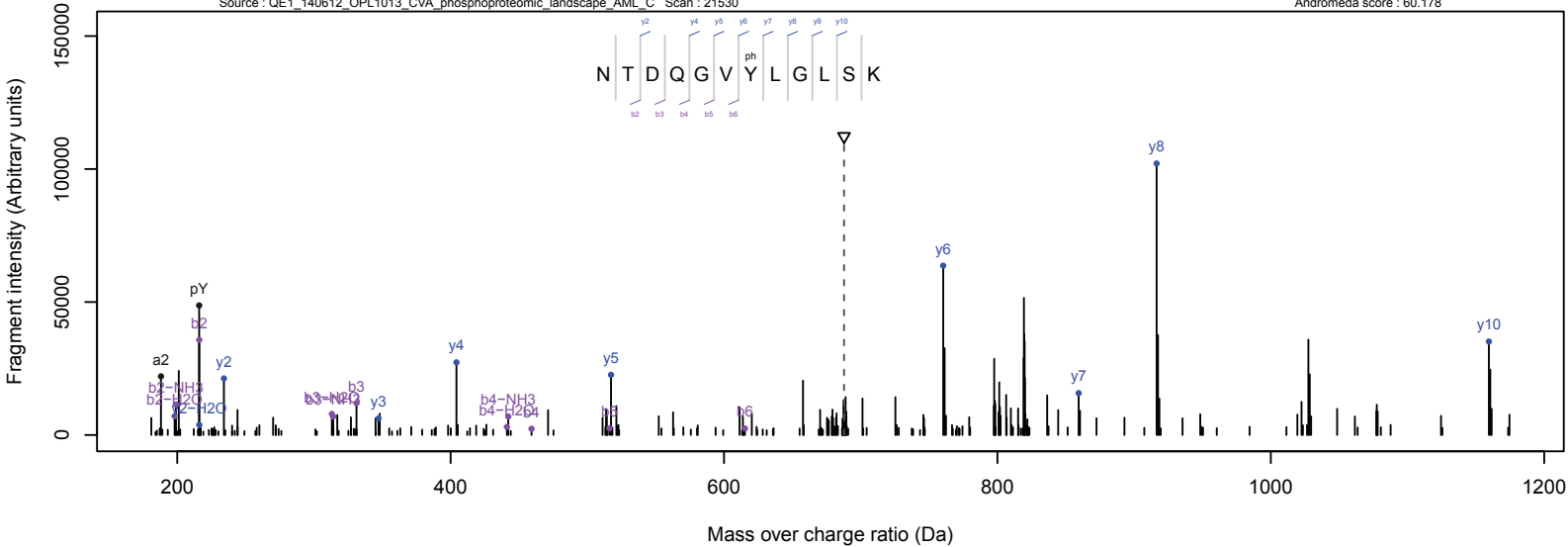
Andromeda score : 60.196



## NTDQGVY(ph)LGLSK-2+

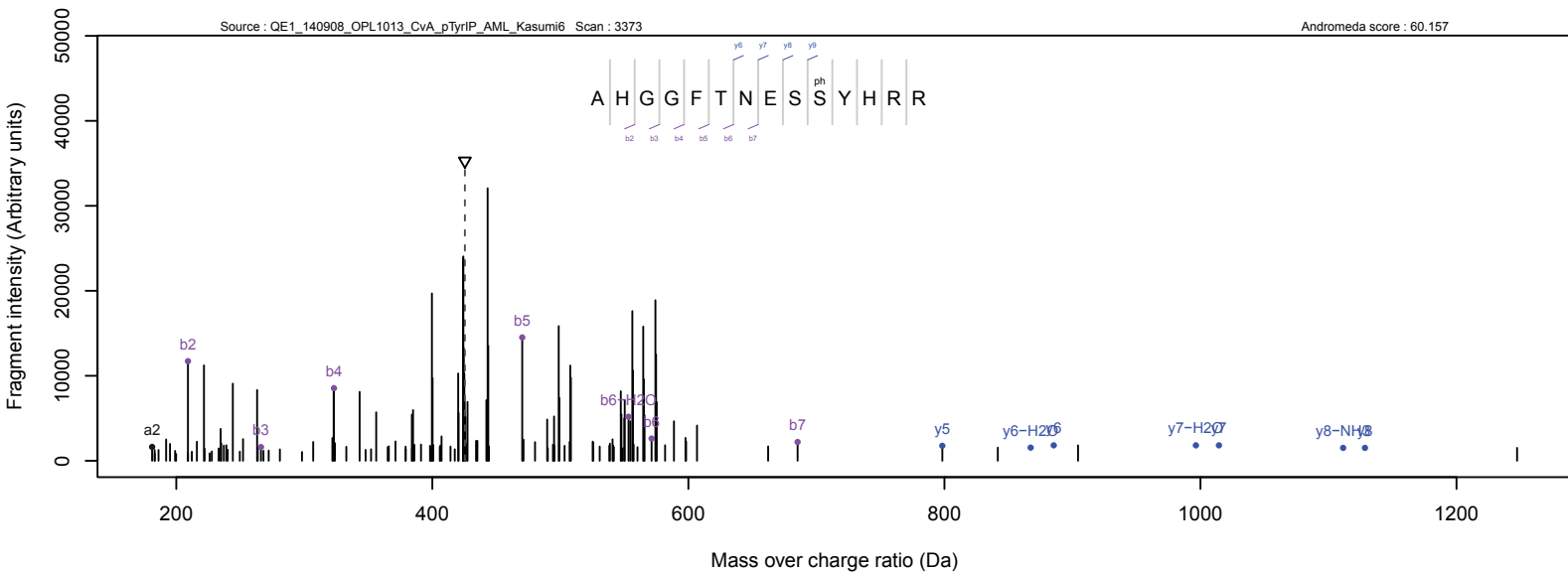
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21530

Andromeda score : 60.178

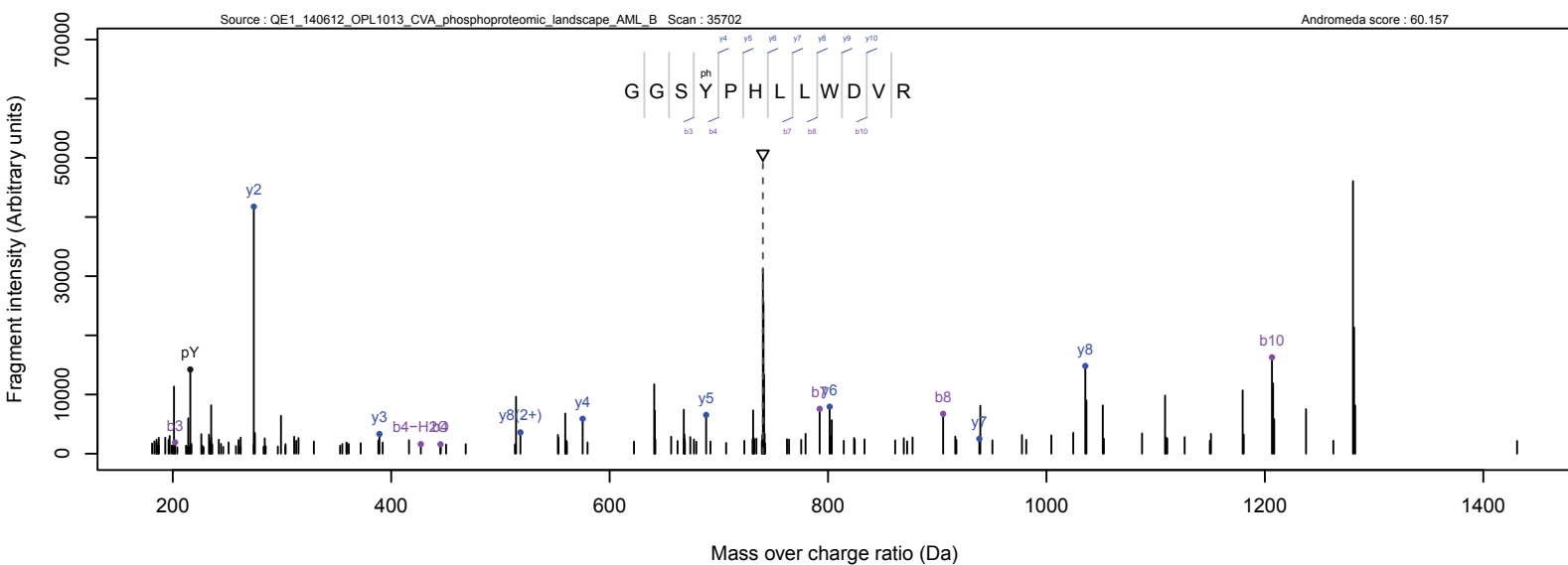




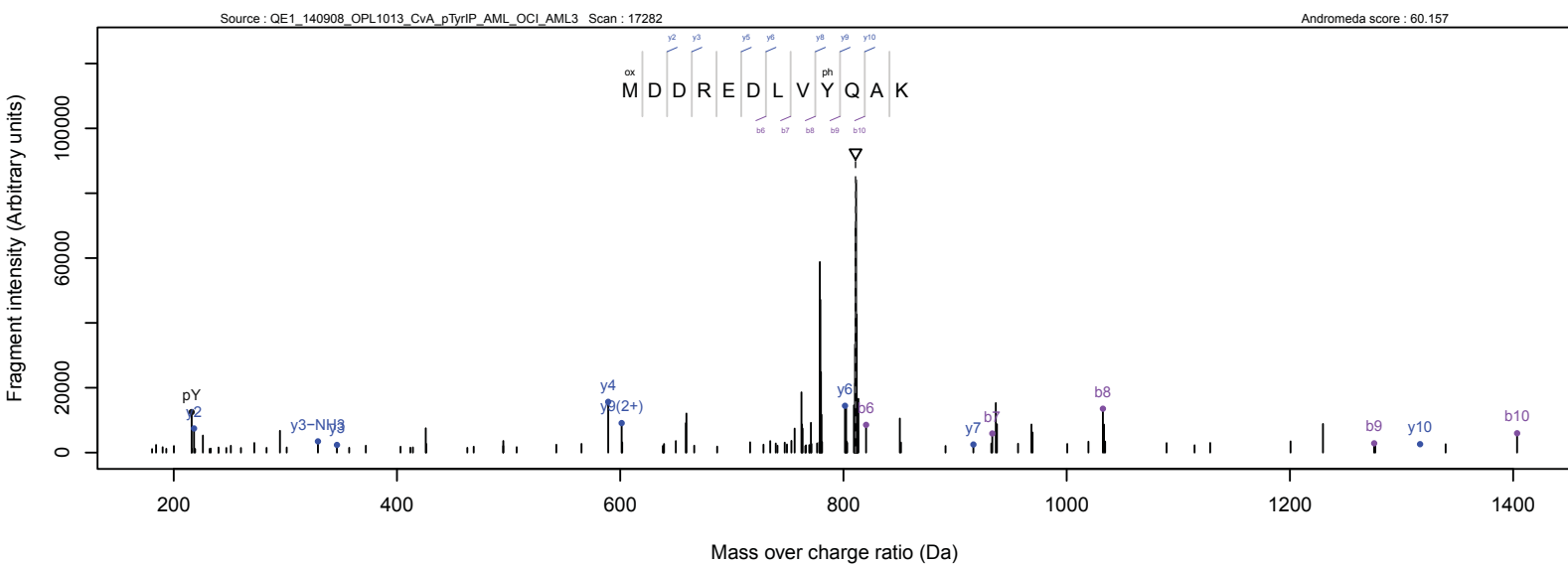
AHGGFTNESS(ph)YHRR-4+



GGSY(ph)PHLLWDVR-2+



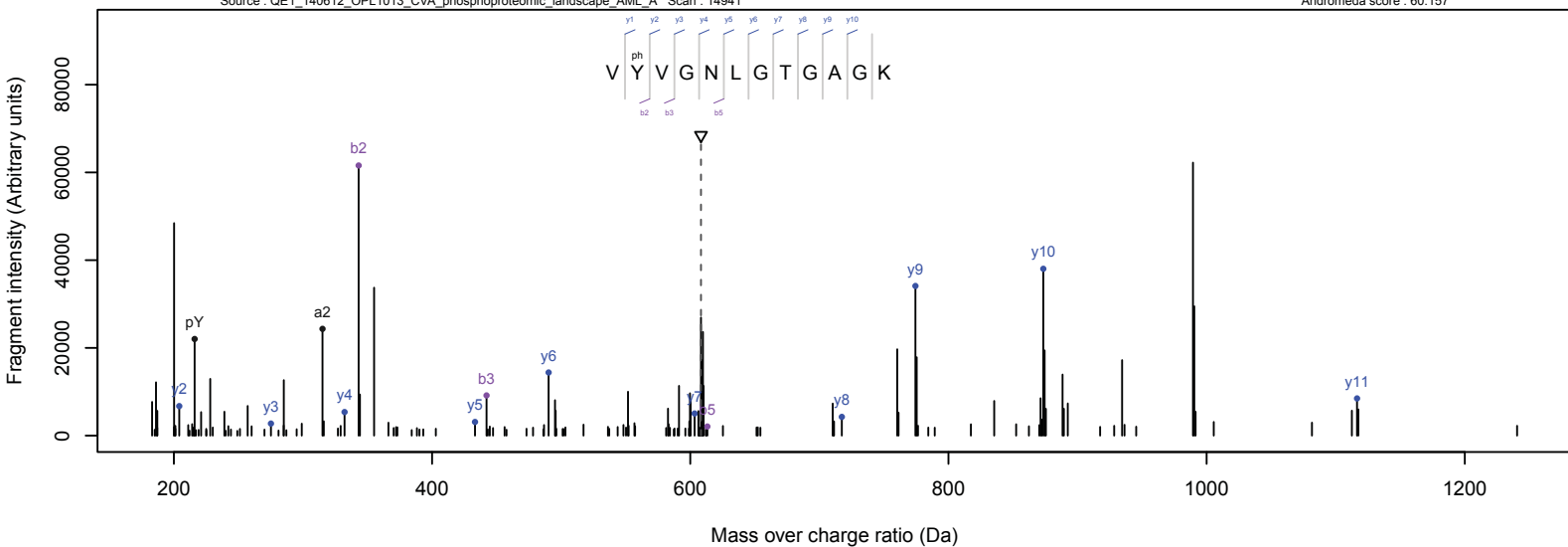
(ac)M(ox)DDREDLVY(ph)QAK-2+



## VY(ph)VGNLTGAGK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 14941

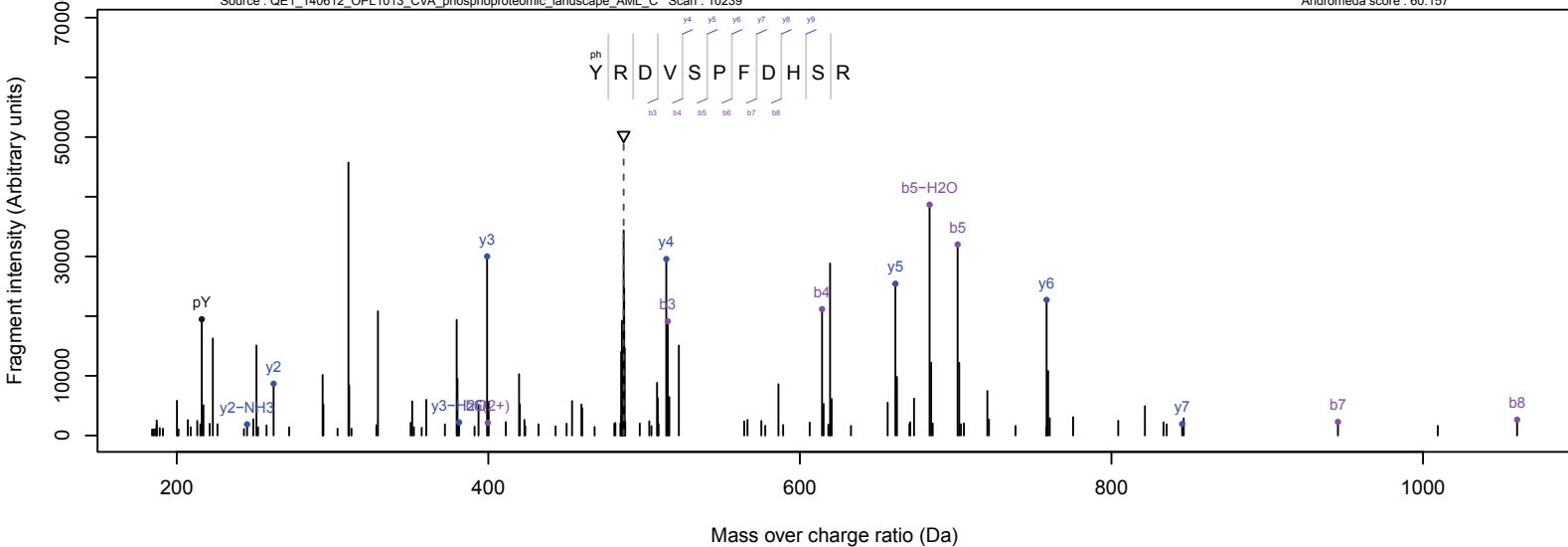
Andromeda score: 60.157



## Y(ph)RDVSPFDHSR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10239

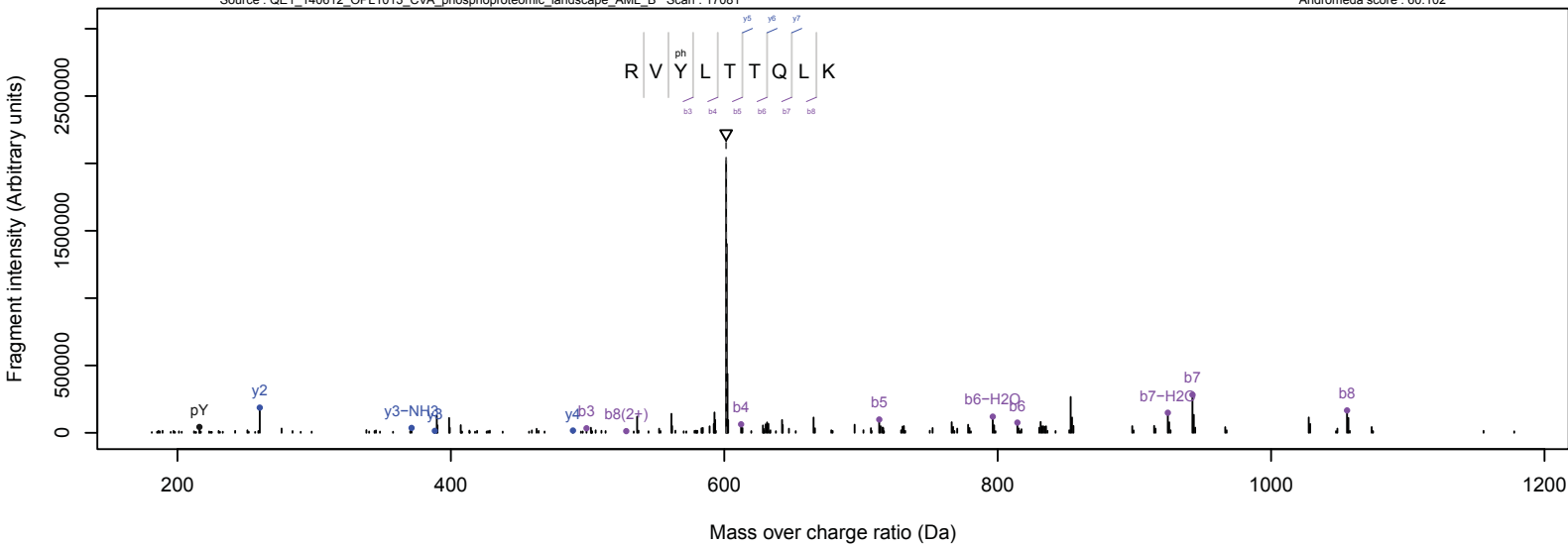
Andromeda score: 60.157



## RVY(ph)LTQLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 17081

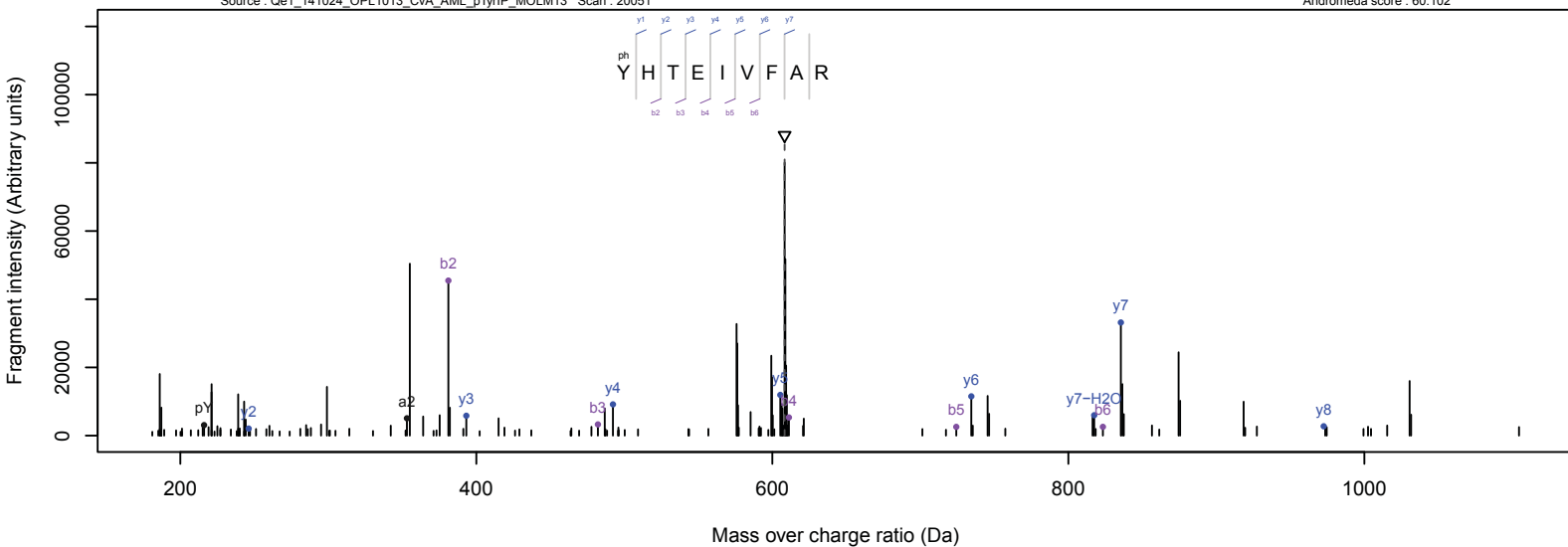
Andromeda score: 60.102



Y(ph)HTEIVFAR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 20051

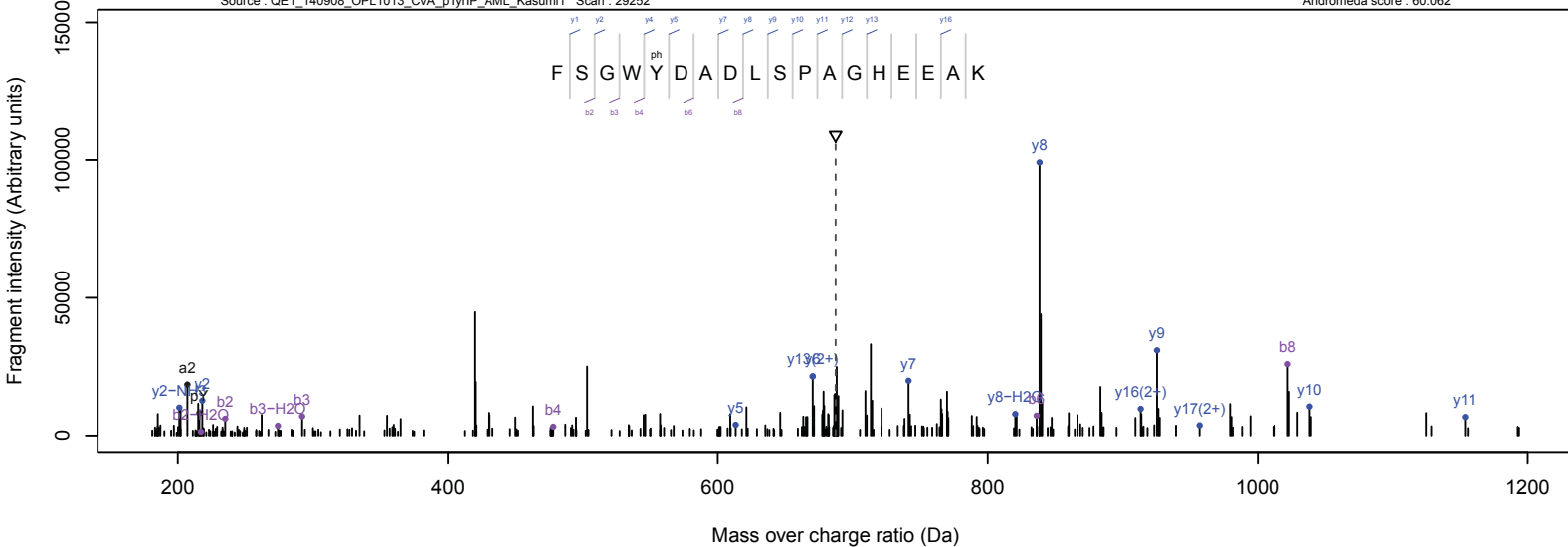
Andromeda score : 60.102



FSGWY(ph)DADLSPAGHEEAK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 29252

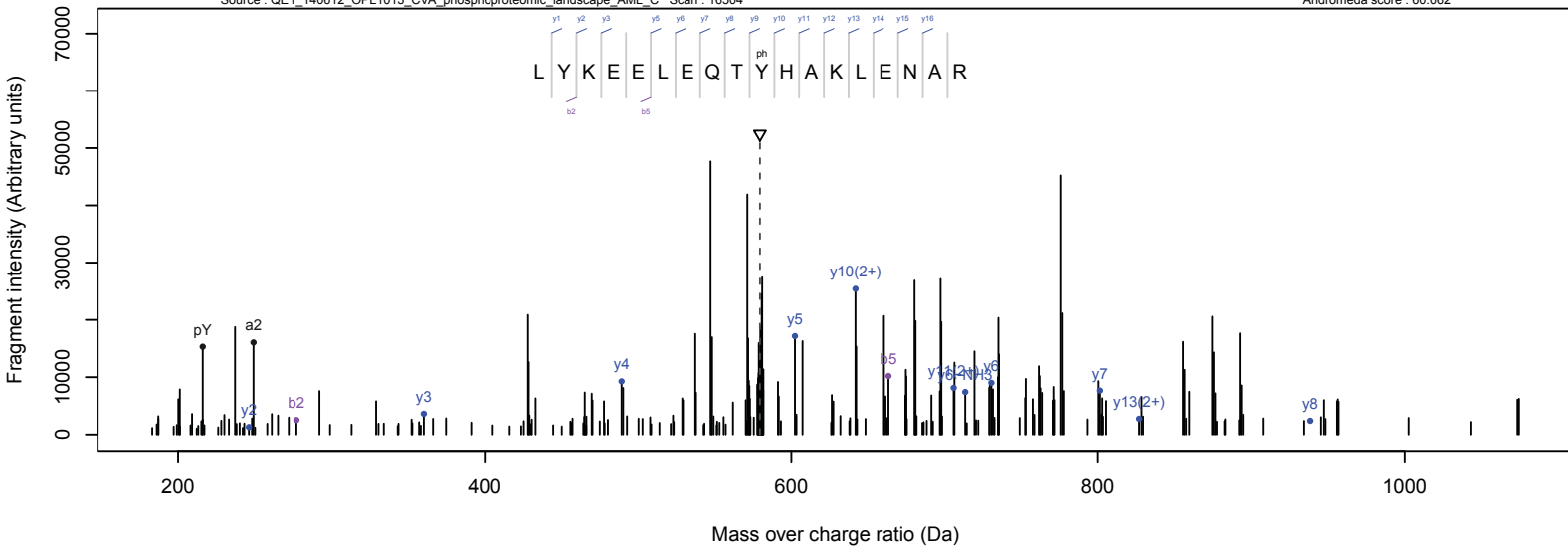
Andromeda score : 60.062



LYKEELEQTY(ph)HAKLENAR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16504

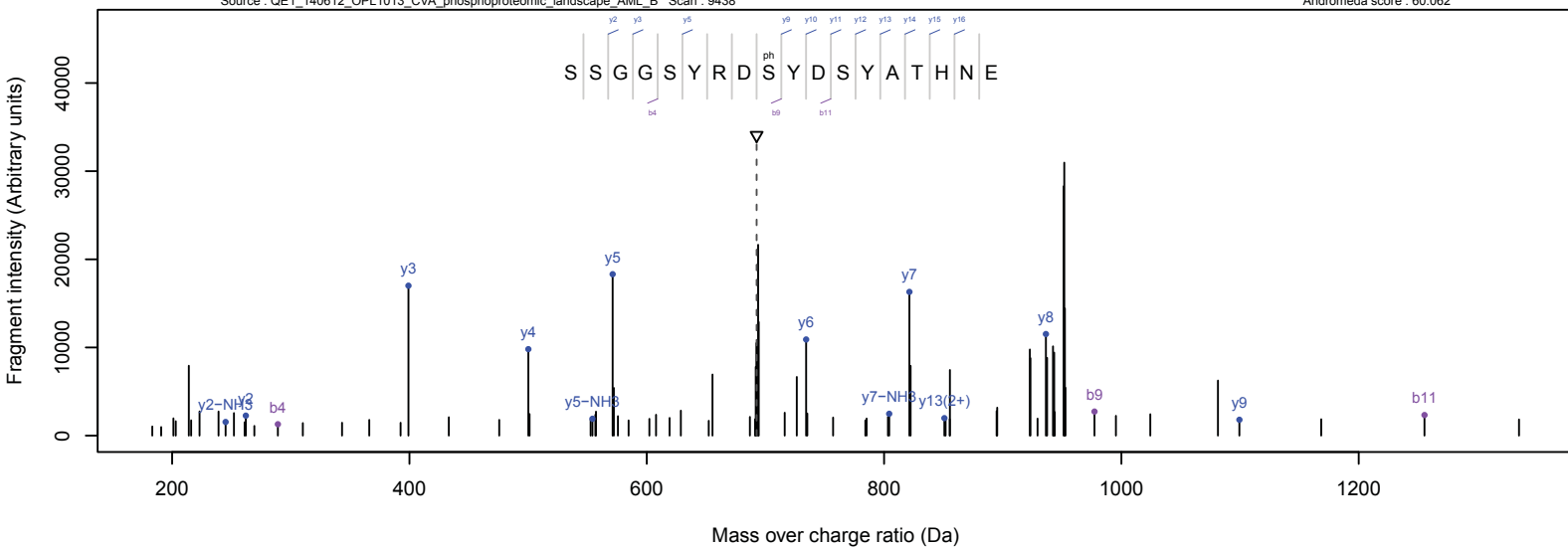
Andromeda score : 60.062



## SSGGSYRDS(ph)YDSYATHNE-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 9438

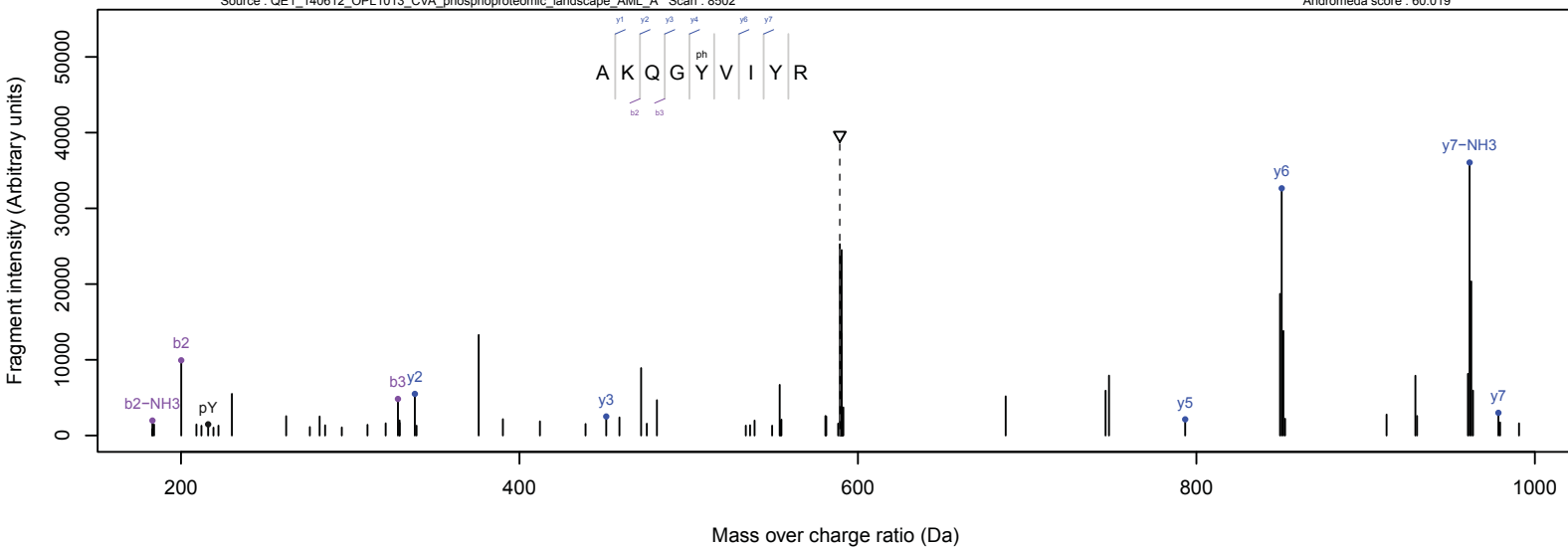
Andromeda score: 60.062



## AKQGY(ph)VIYR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 8502

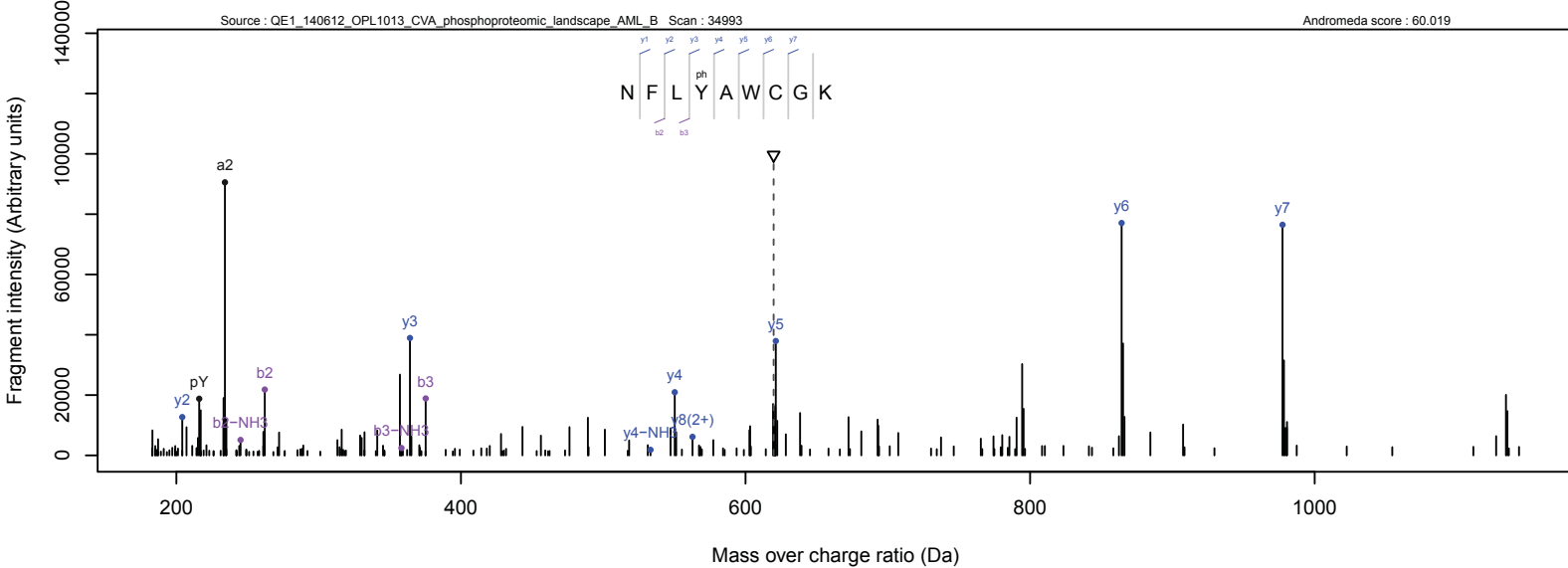
Andromeda score: 60.019



## NFLY(ph)AWCGK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 34993

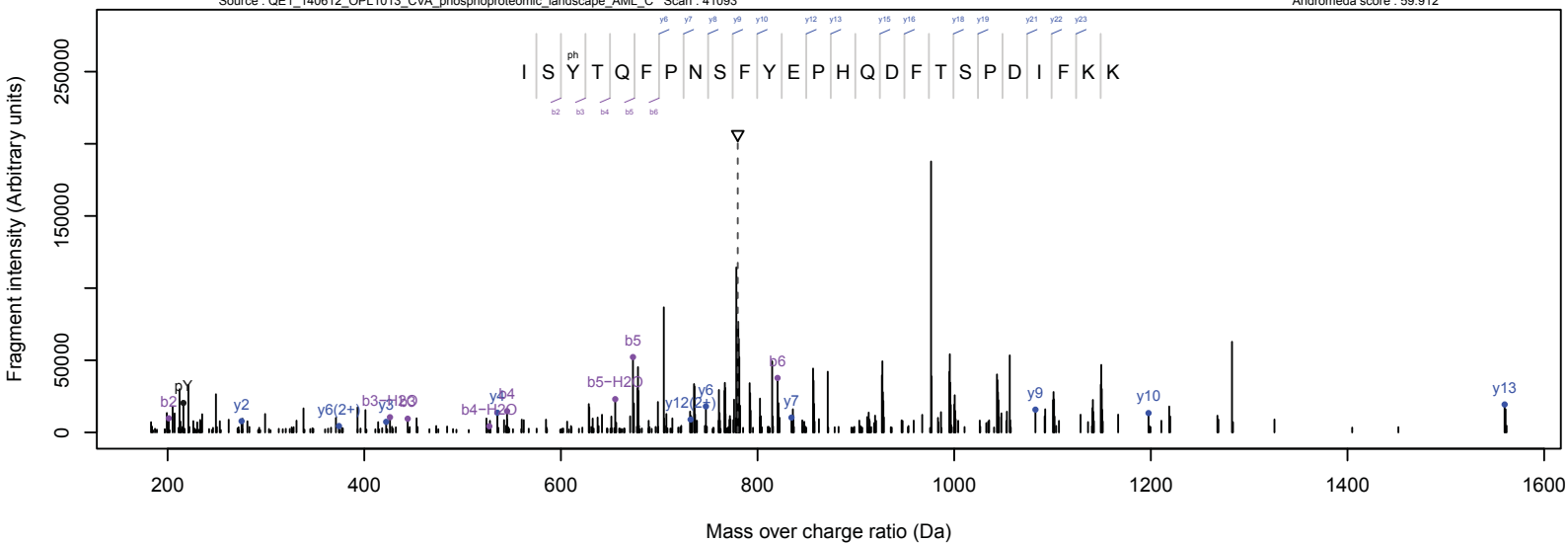
Andromeda score: 60.019



ISY(ph)TQFPNSFYEPHQDFTSPDIFKK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41093

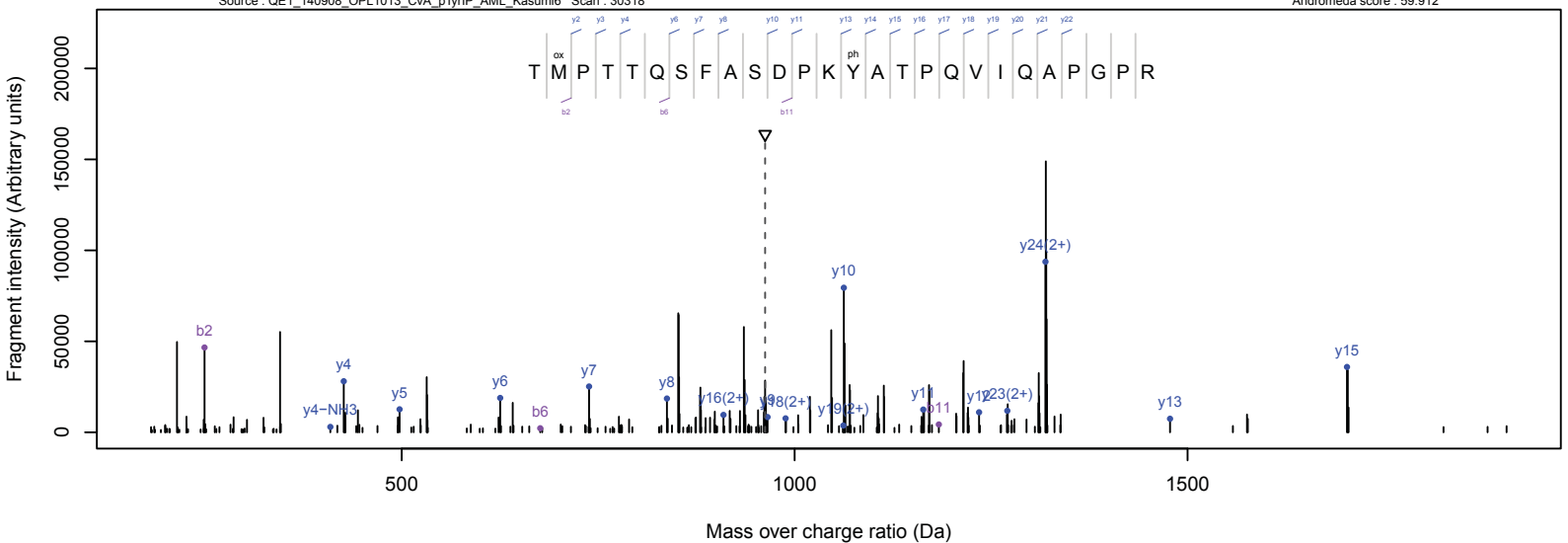
Andromeda score : 59.912



TM(ox)PTTQSFASDPK(ph)ATPQVIQAPGPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 30318

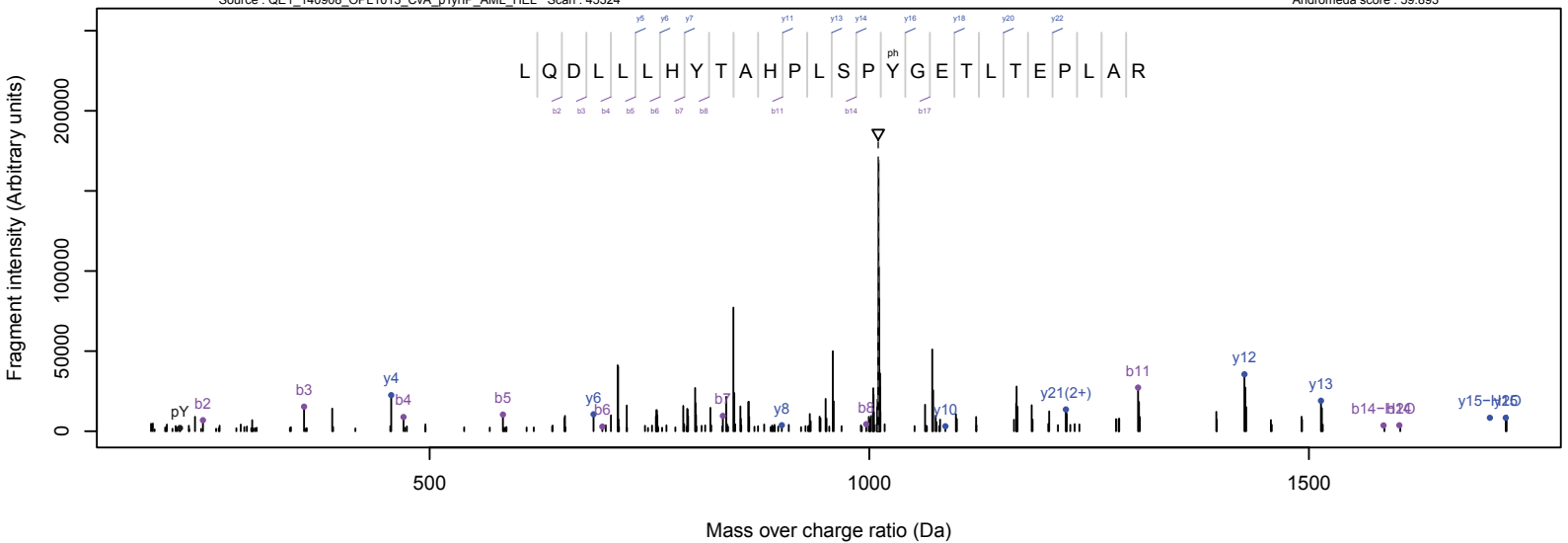
Andromeda score : 59.912

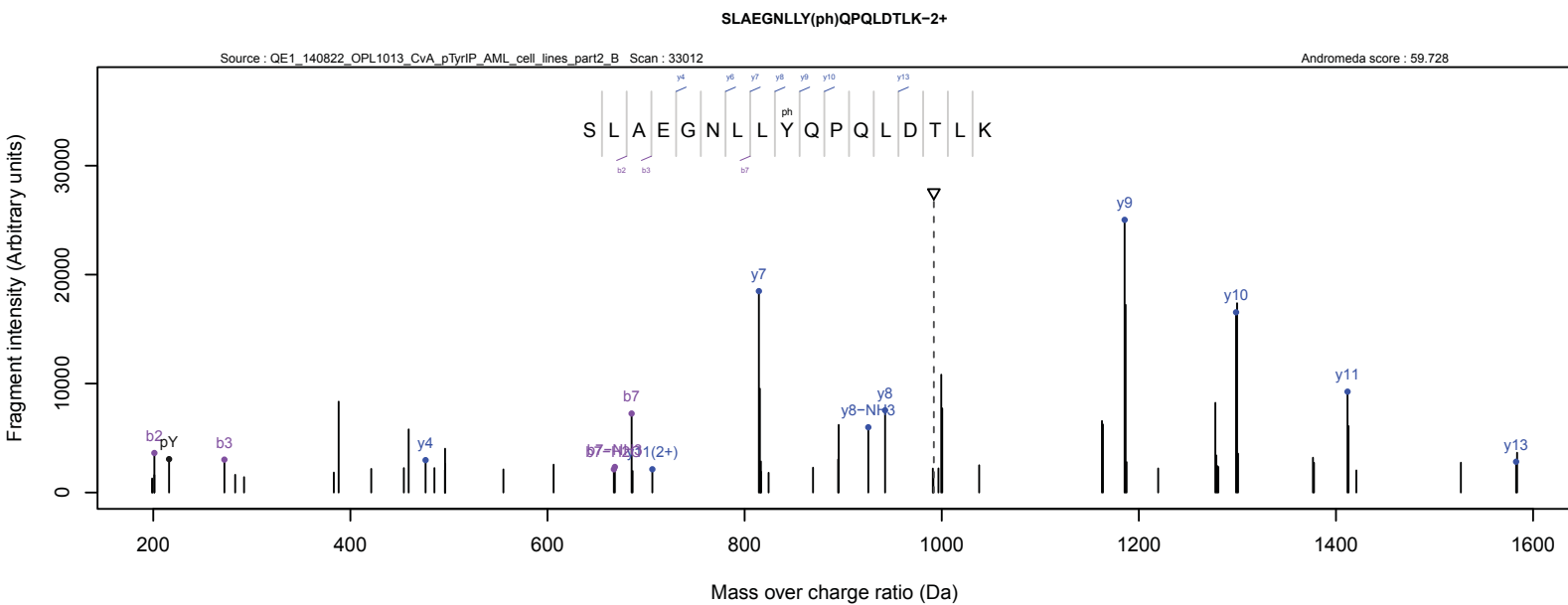
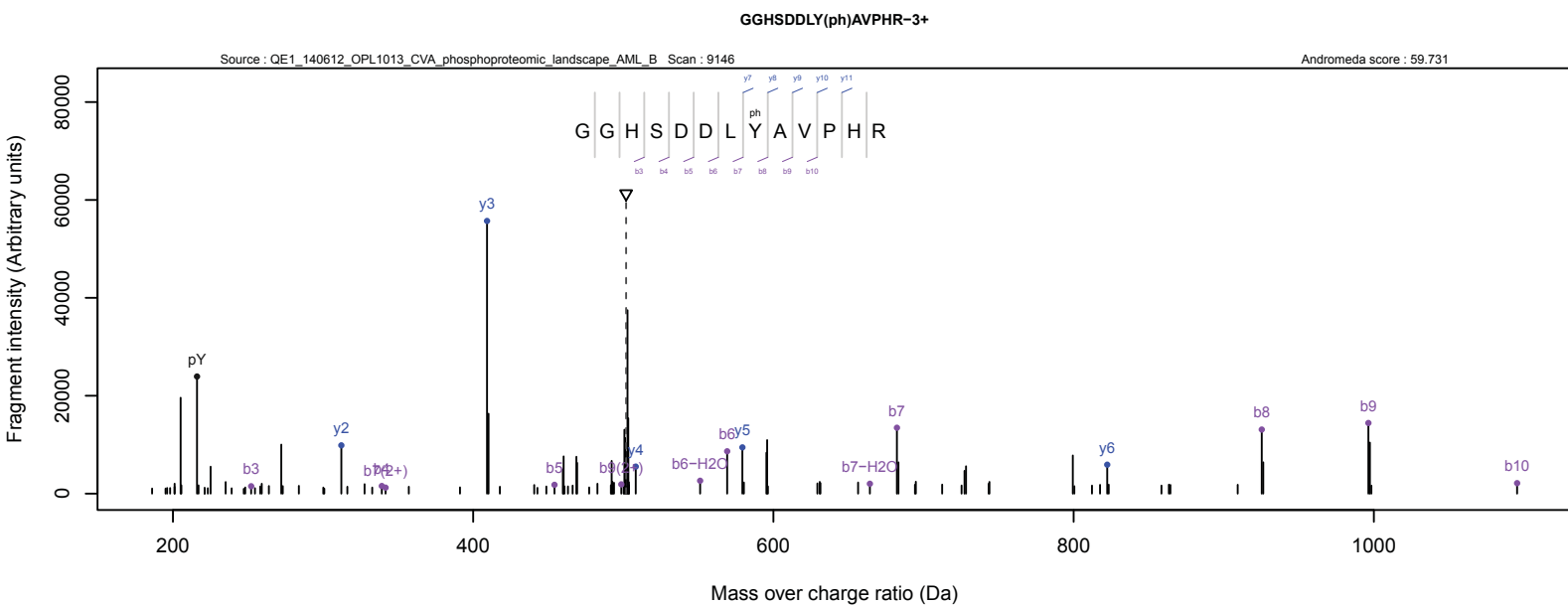
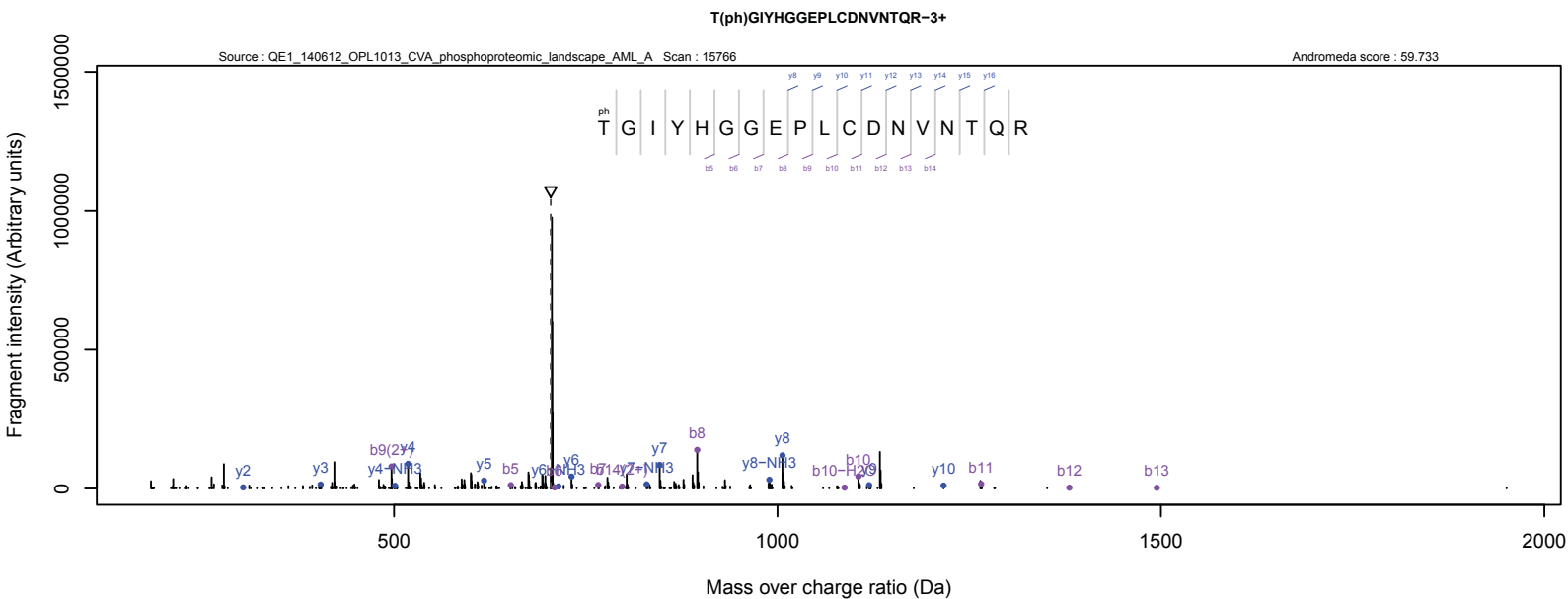


LQDLLLHYTAHPLSPYGETLTEPLAR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 45324

Andromeda score : 59.893

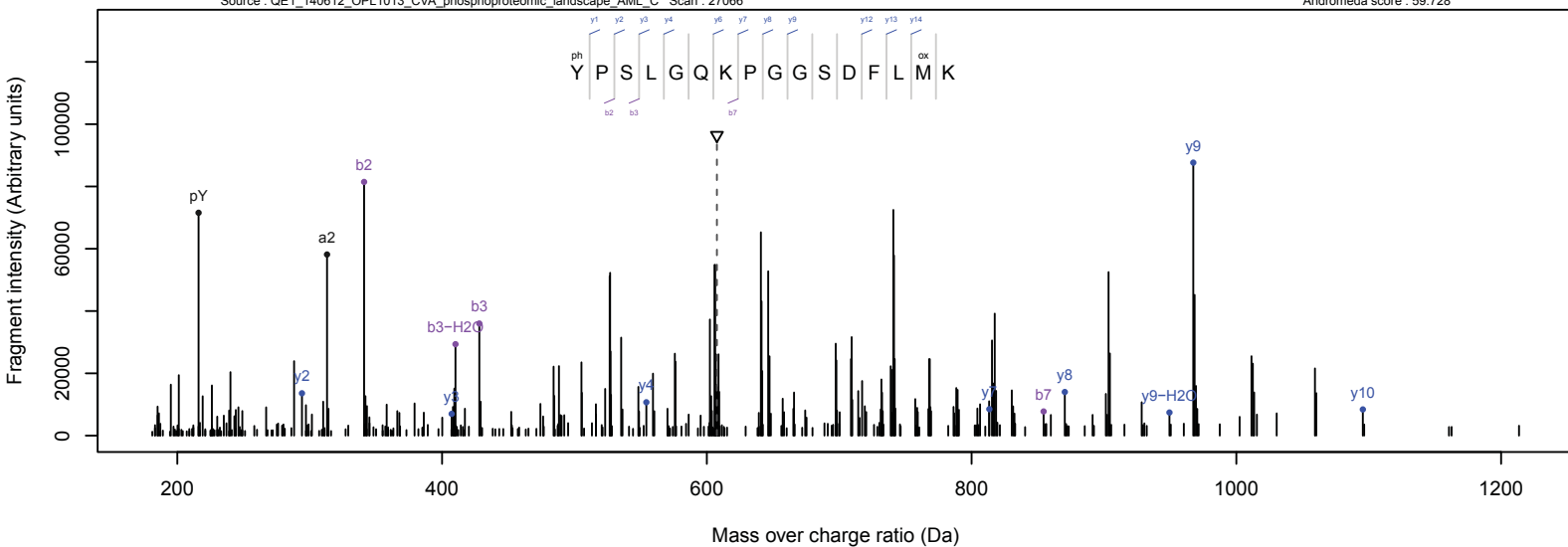




## Y(ph)PSLGQKPGGSDFLM(ox)K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27066

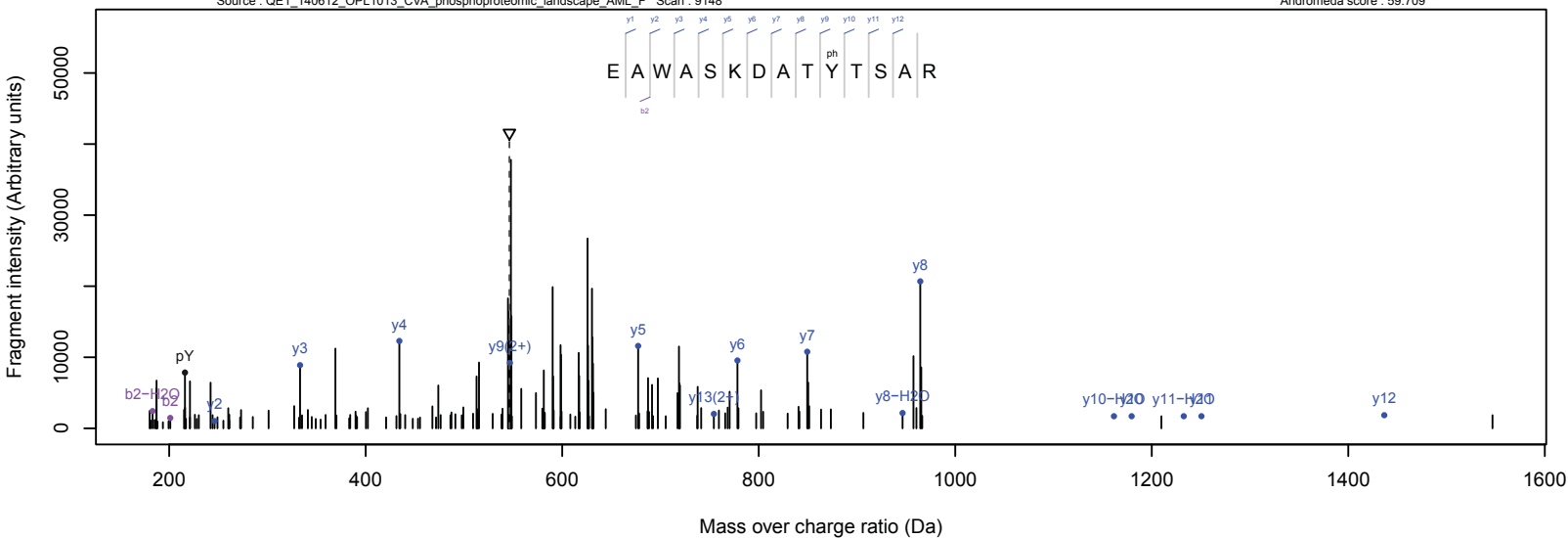
Andromeda score : 59.728



## EAWASKDATY(ph)TSAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 9148

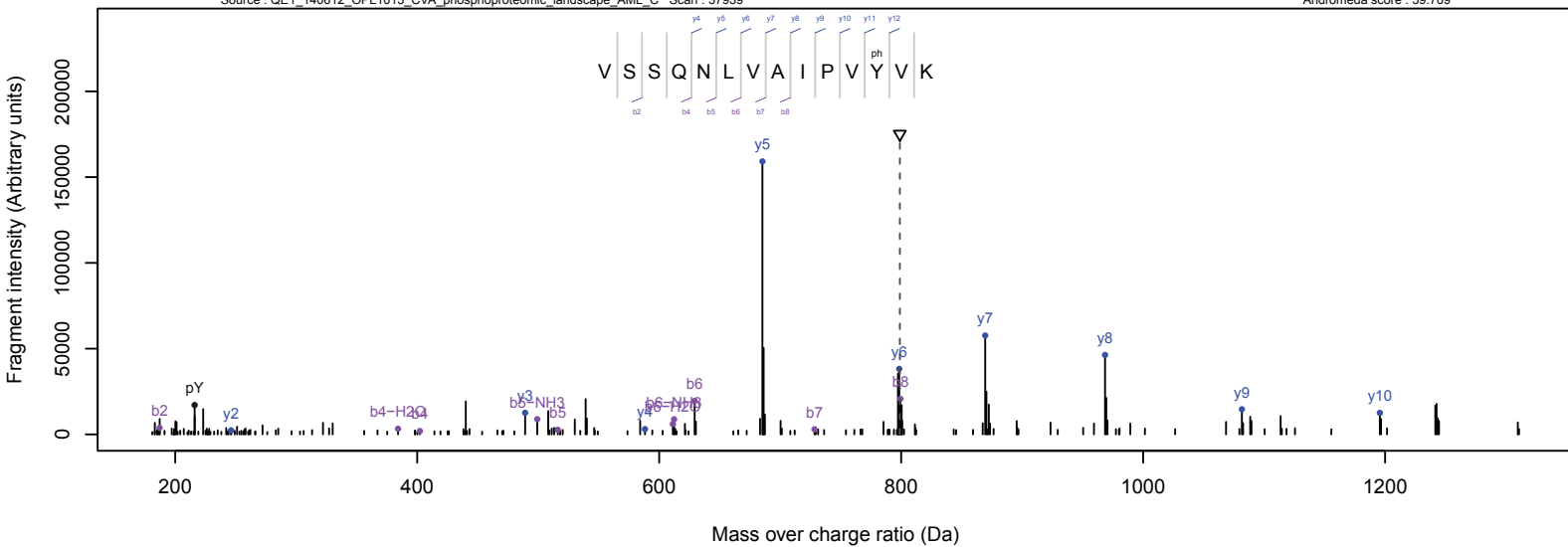
Andromeda score : 59.709



## VSSQLVAIPVY(ph)VK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37939

Andromeda score : 59.709



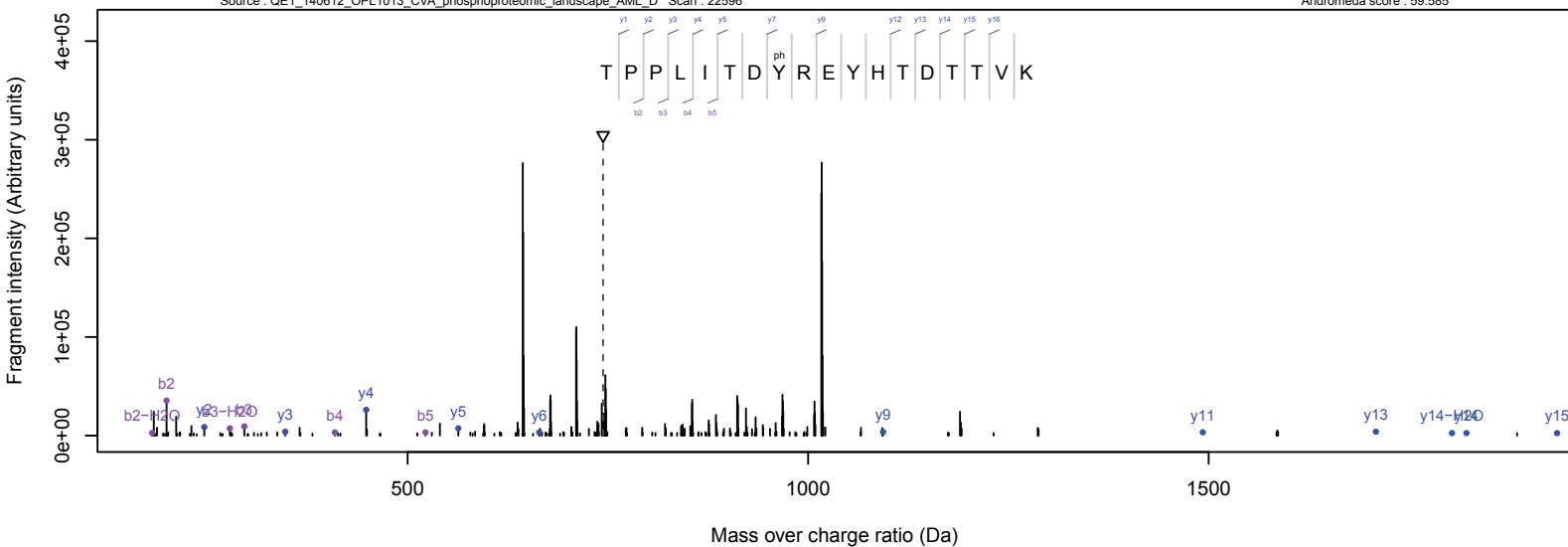




TPPLITDY(ph)REYHTDTTVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 22596

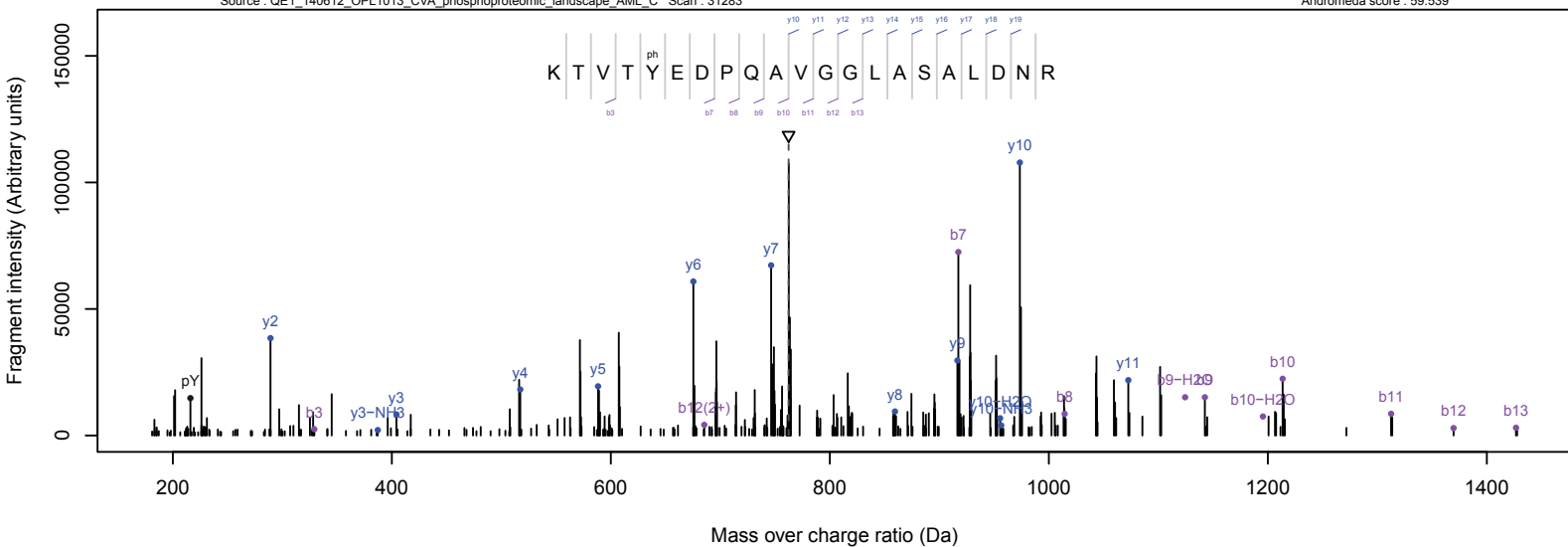
Andromeda score: 59.585



KTVTY(ph)EDPQAVGGLASALDNR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 31283

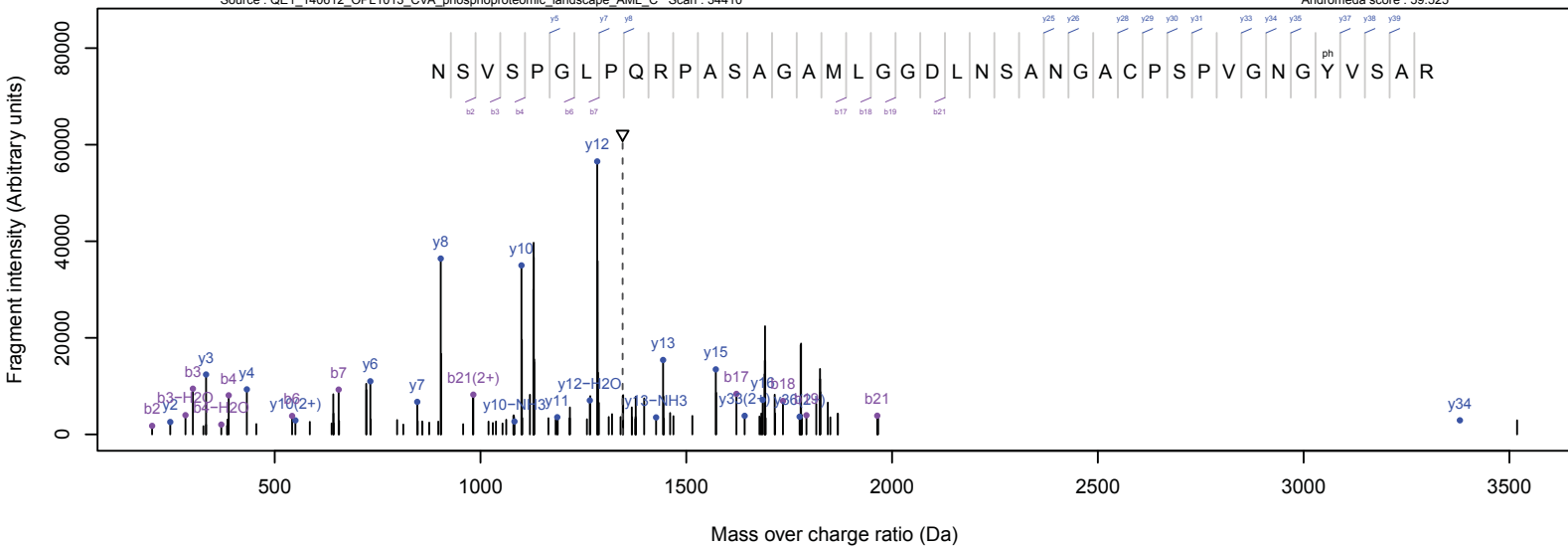
Andromeda score: 59.539



NSVSPGLPQRPASAGAMLGGLNSANGACPSVGNNGY(ph)VSAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34410

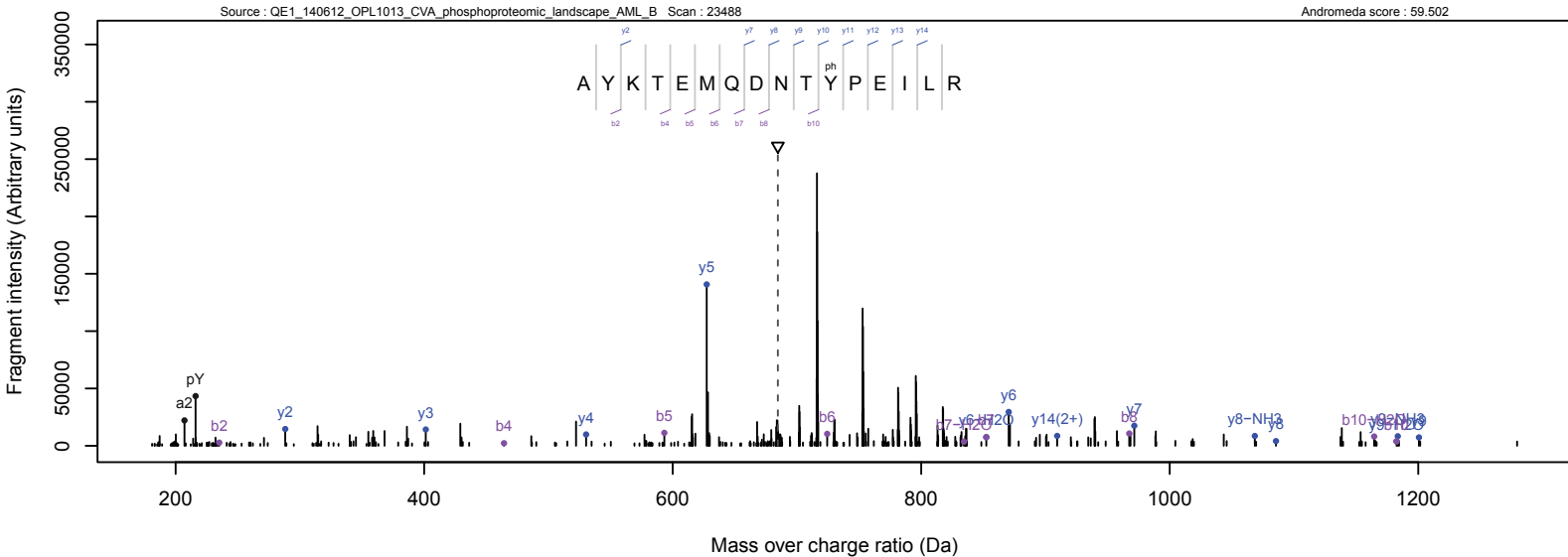
Andromeda score: 59.525



AYKTEMQDNTY(ph)PEILR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23488

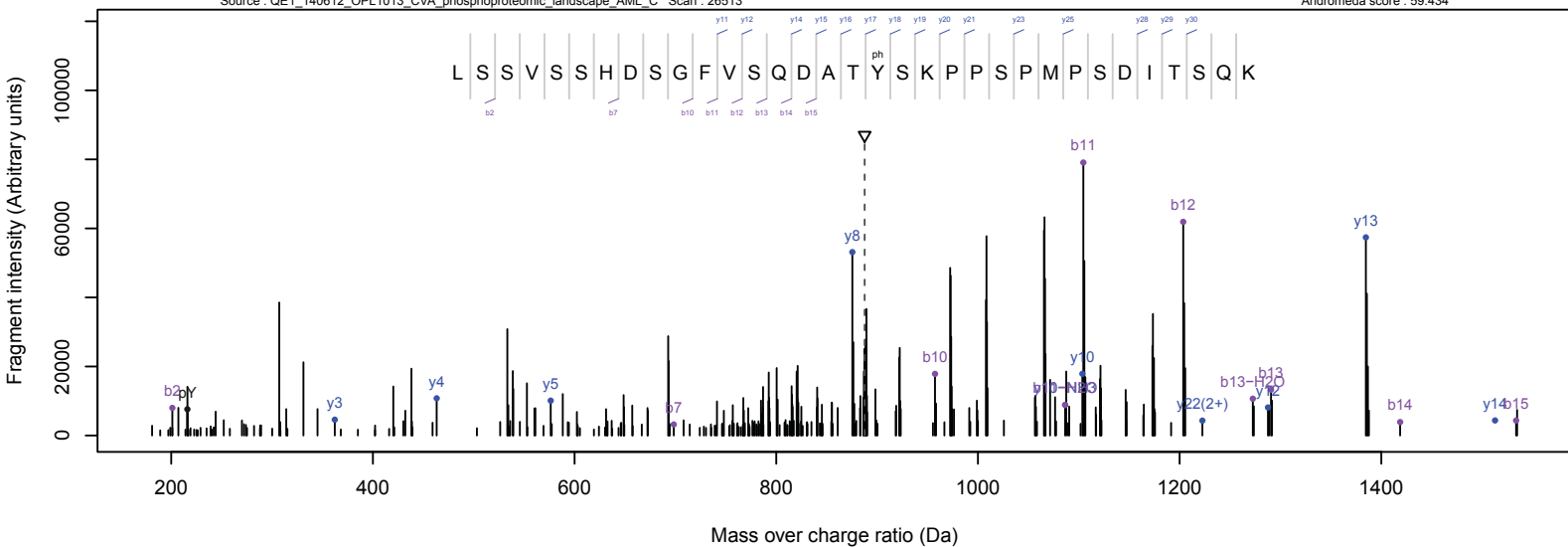
Andromeda score : 59.502



LSSVSSHDSGFVSDATY(ph)SKPPSPMPDITSQK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26513

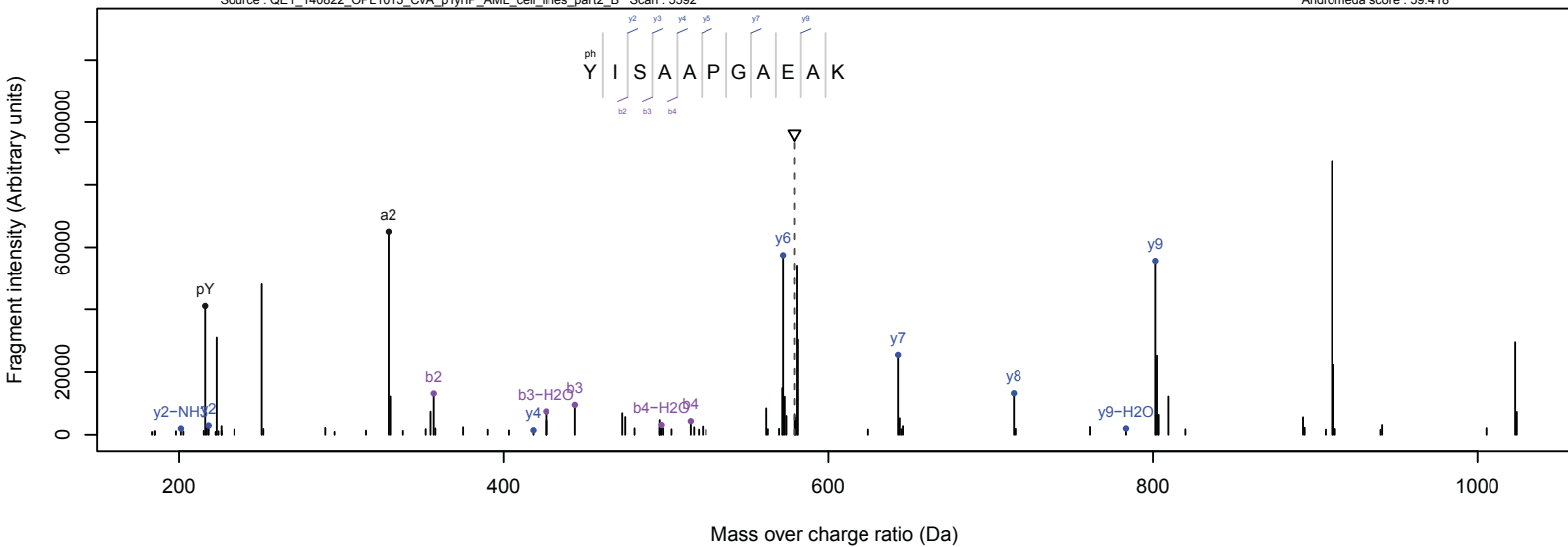
Andromeda score : 59.434



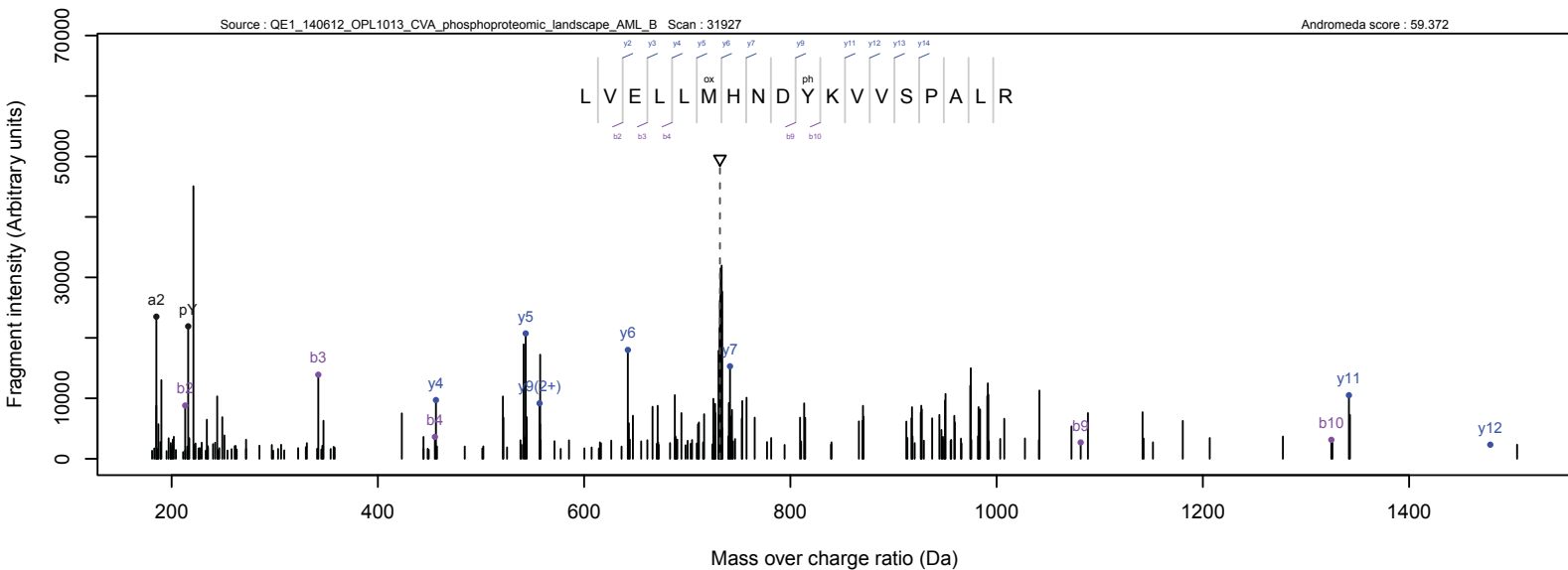
Y(ph)ISAAPGAEAK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 5592

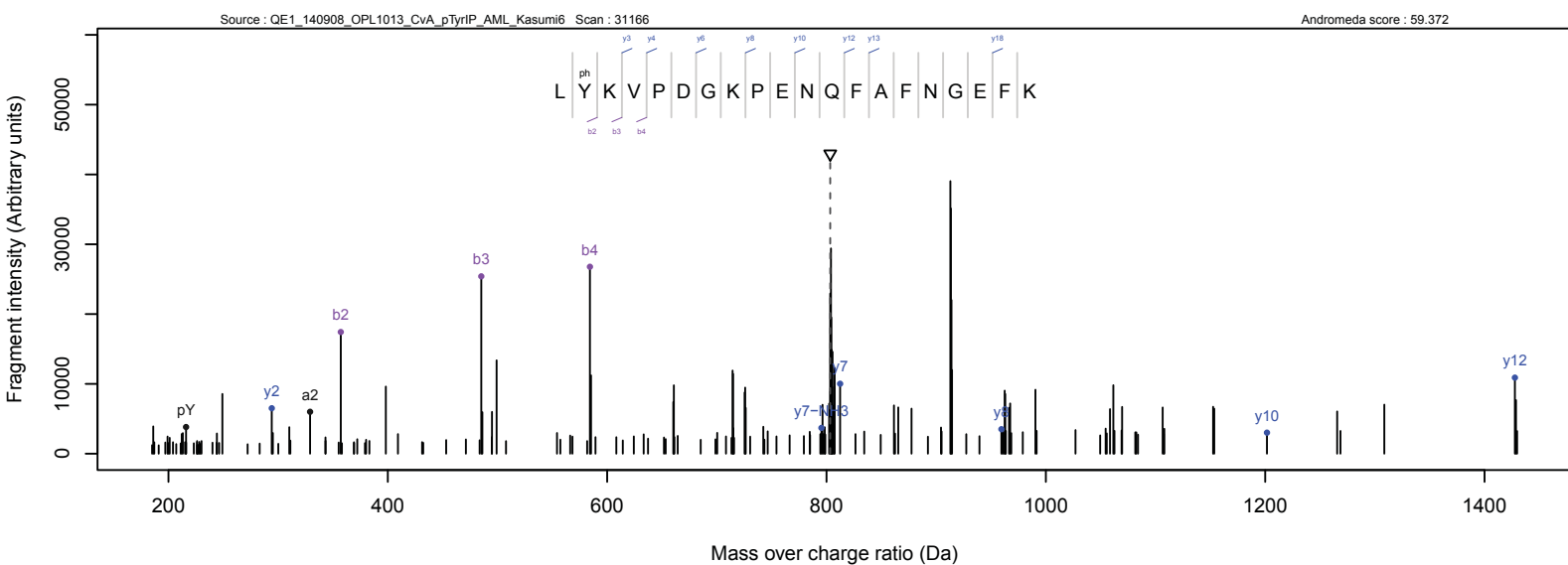
Andromeda score : 59.418



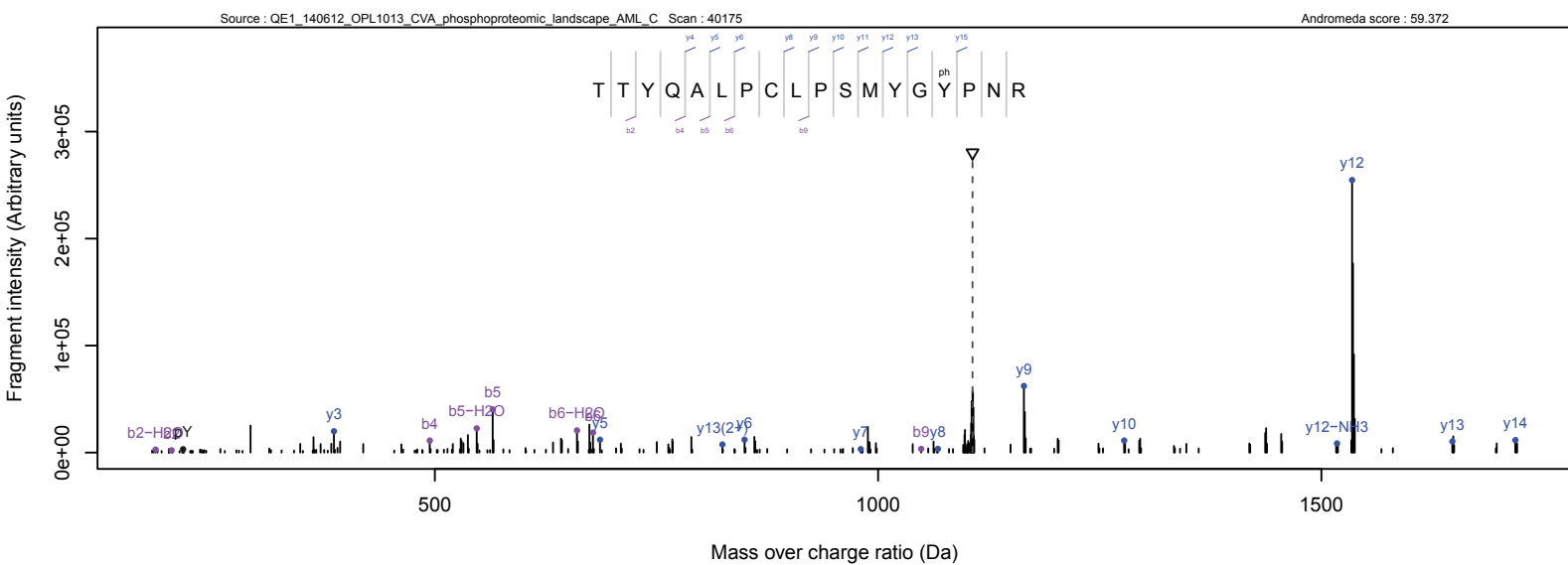
LVELLM(ox)HNDY(ph)KVVPALR-3+



LY(ph)KVPDQKPEQFAFNGEFK-3+



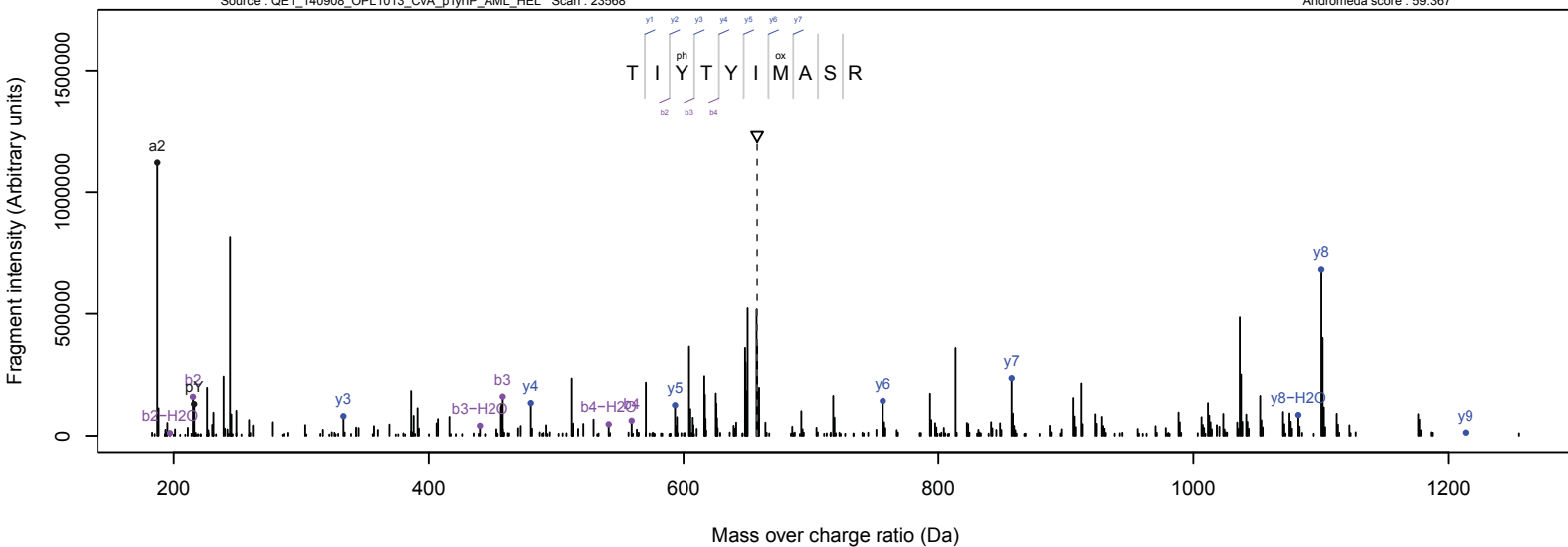
TTYQALPCLPSMYG(ph)PNR-2+



## TIY(ph)TYIM(ox)ASR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23568

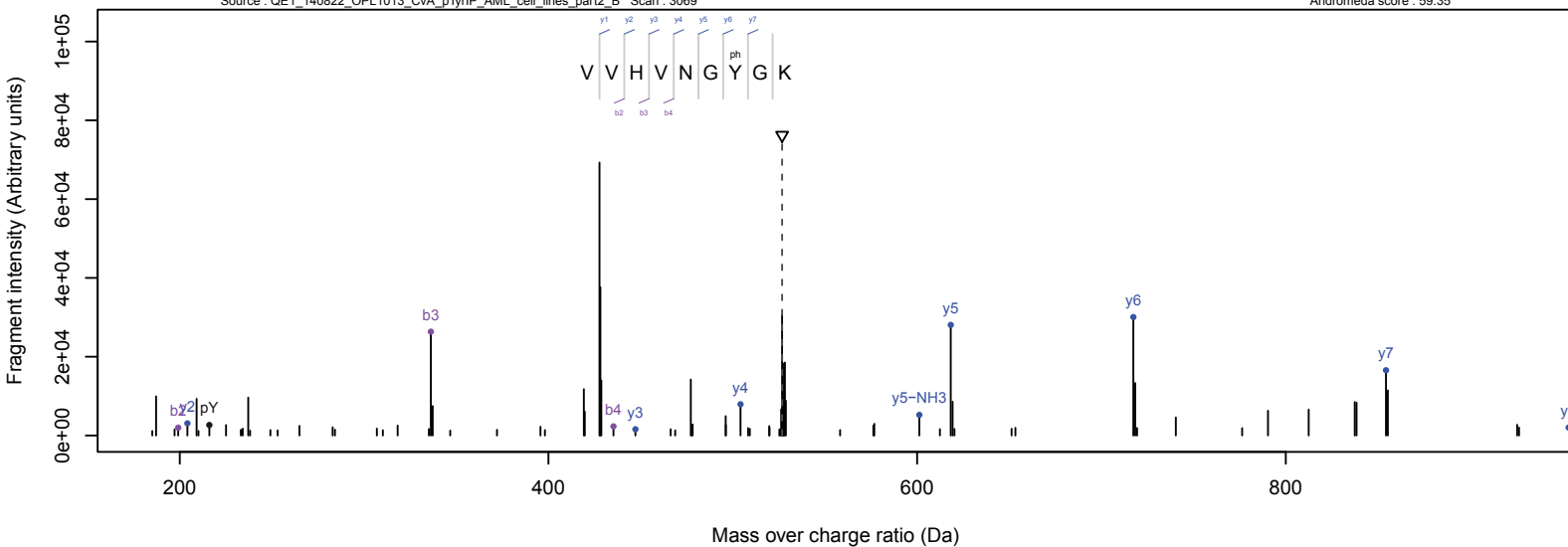
Andromeda score : 59.367



## VVHVNGY(ph)GK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3069

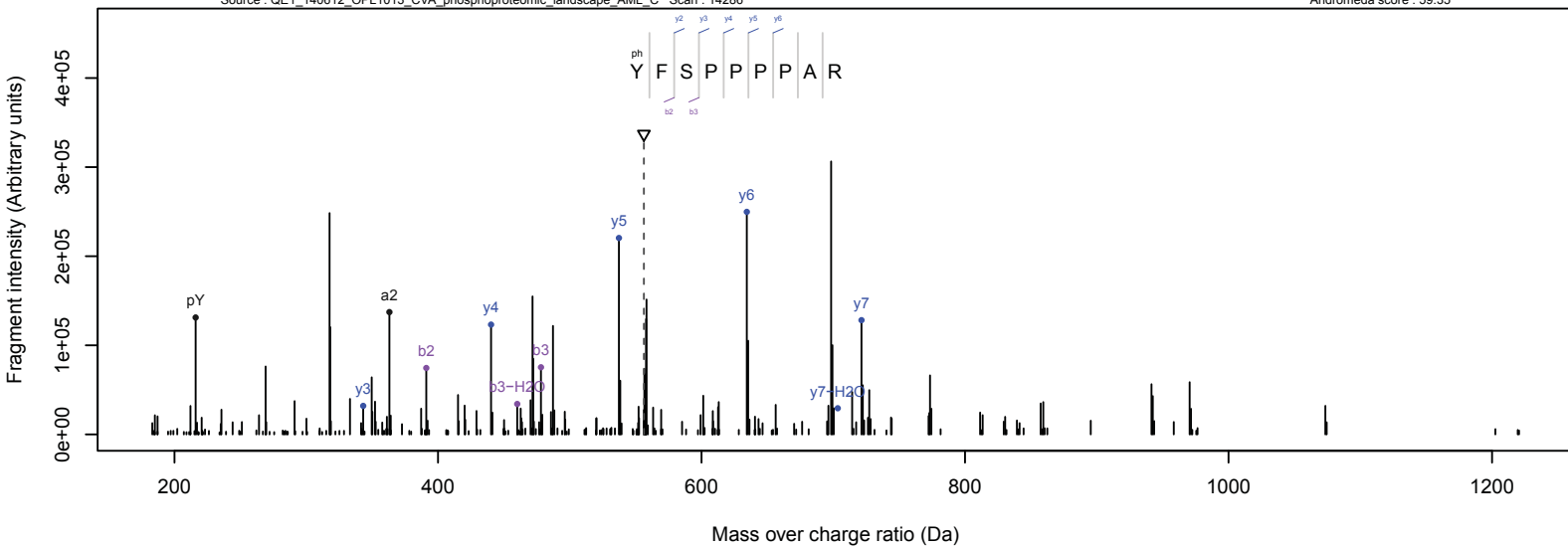
Andromeda score : 59.35



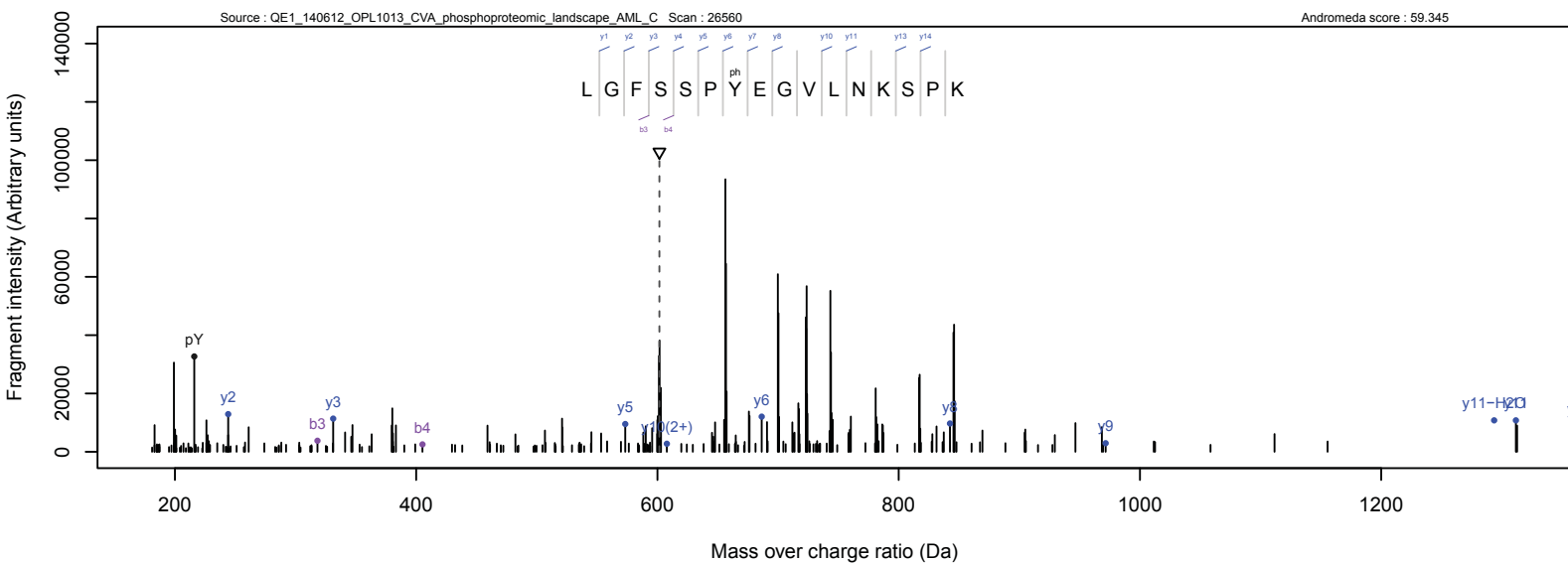
## Y(ph)FSPPPAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14286

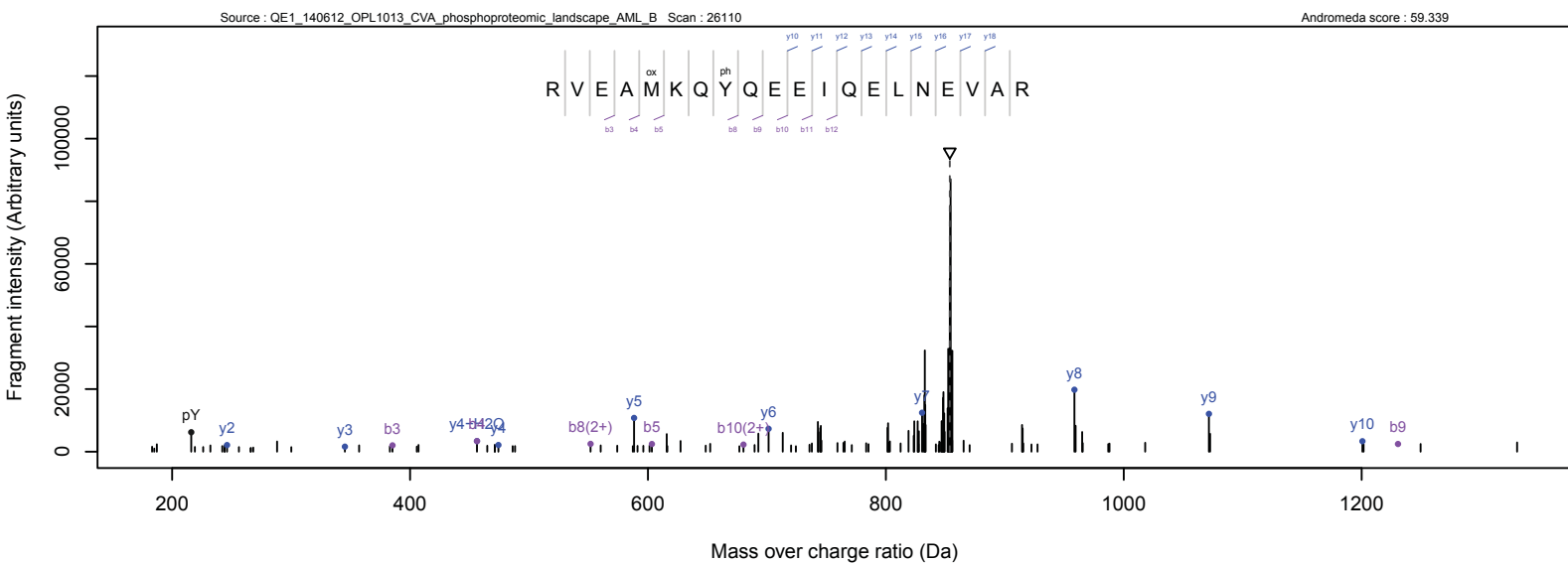
Andromeda score : 59.35



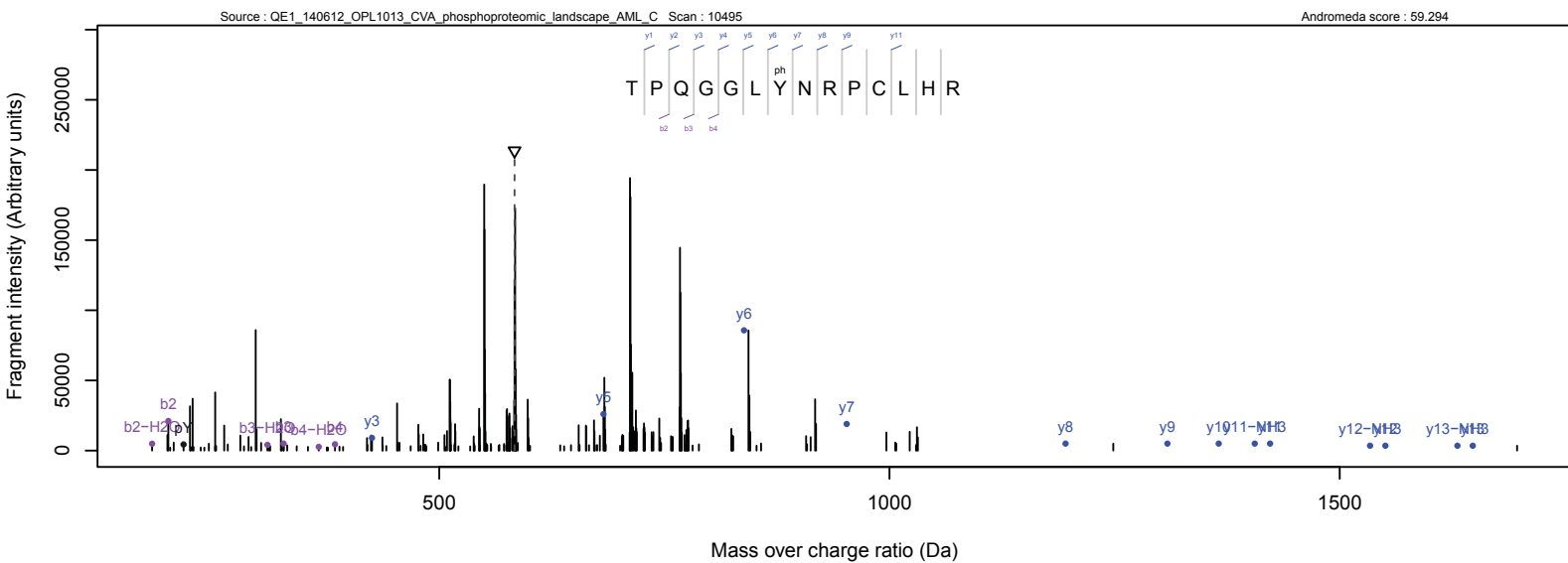
**LGFSSPY(ph)EGLNKSPK-3+**



**RVEAM(ox)KQY(ph)QEEIQELNEVAR-3+**



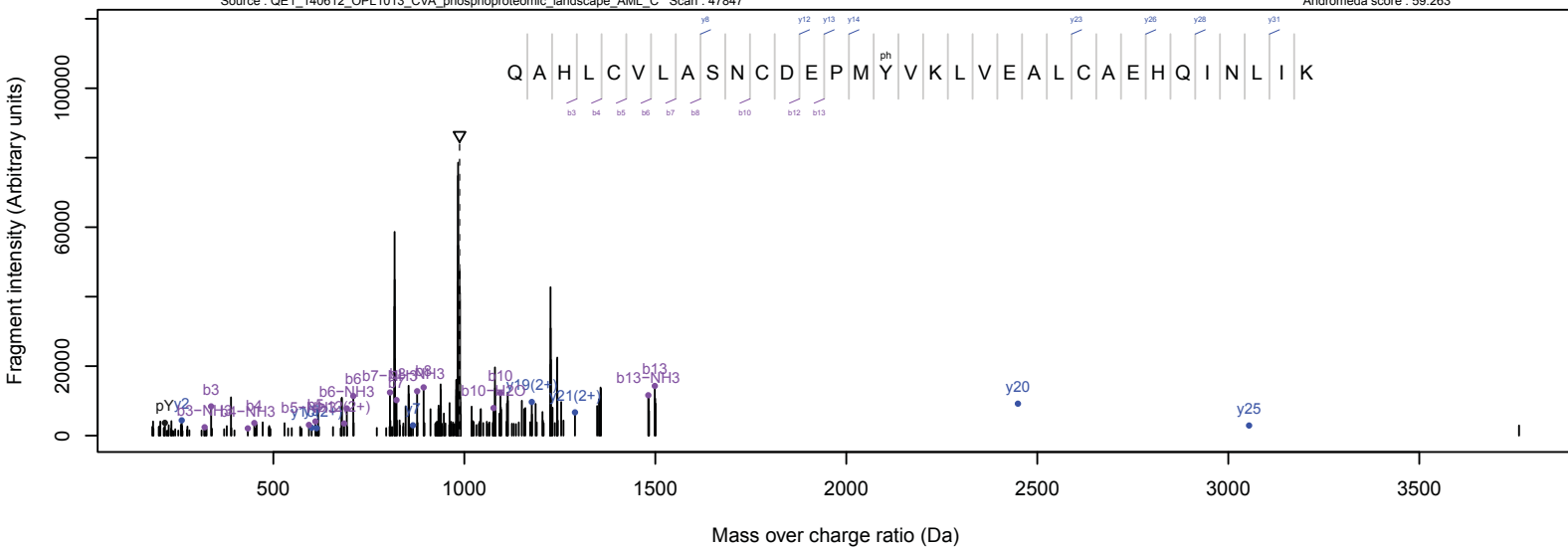
**TPQGGLY(ph)NRPCLHR-3+**



QAHLCVLASNCDEPMY(ph)VKLVEALCAEHQINLIK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47847

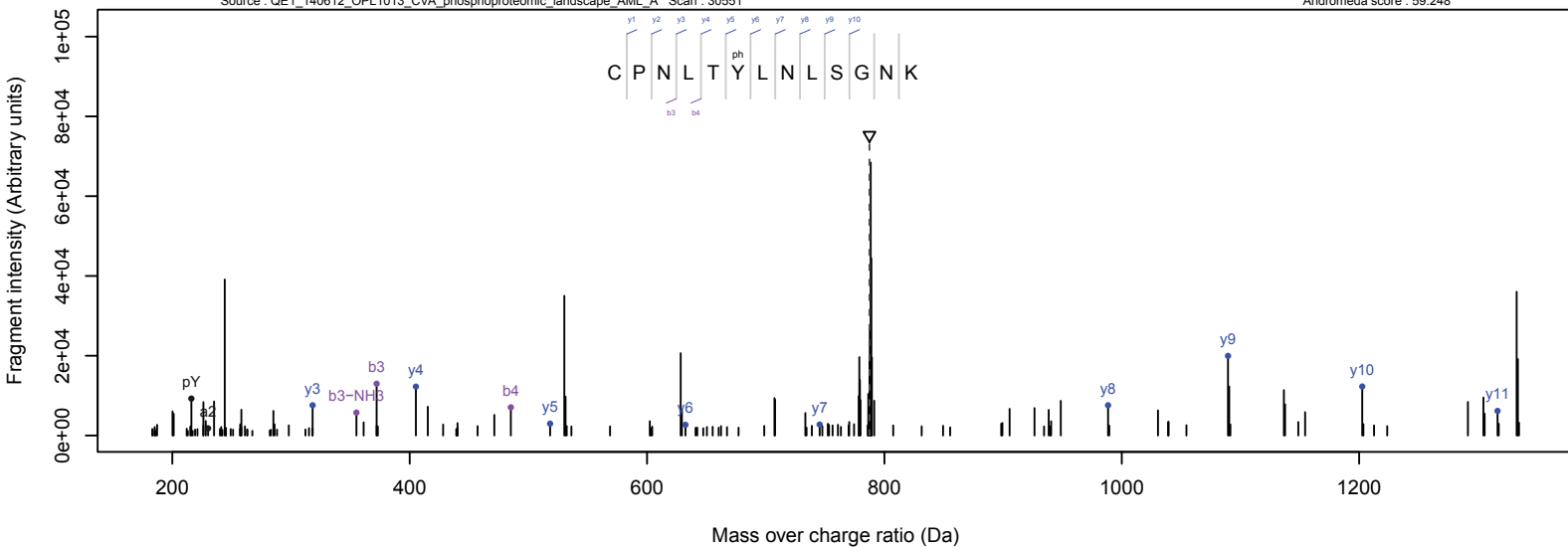
Andromeda score : 59.263



CPNLTY(ph)LNLSGNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 30551

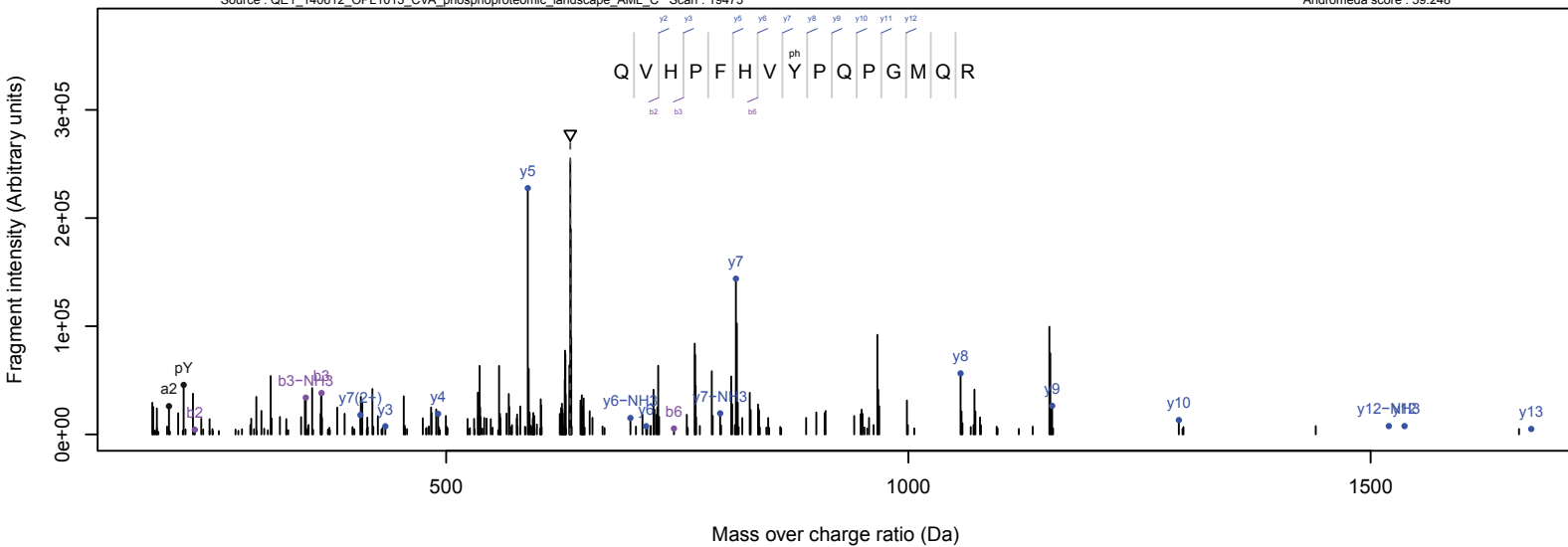
Andromeda score : 59.248



QVHPFHVY(ph)PQPGMQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19475

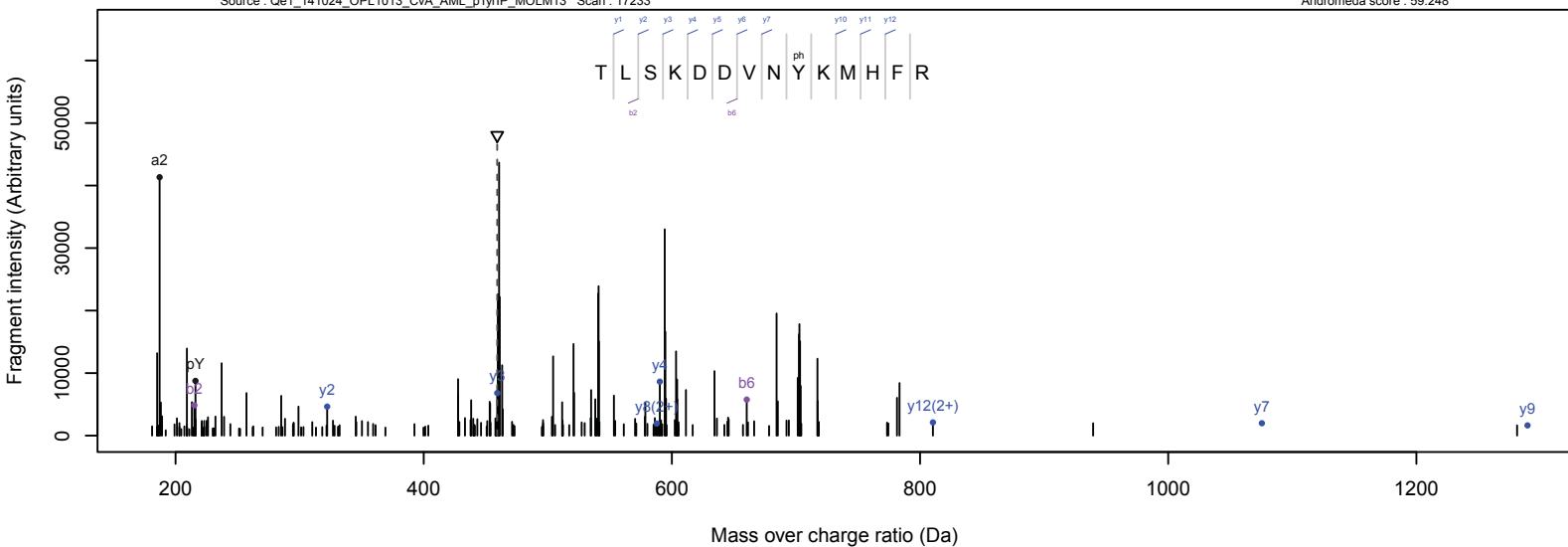
Andromeda score : 59.248



## TLSKDDVNY(ph)KMHFR-4+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 17233

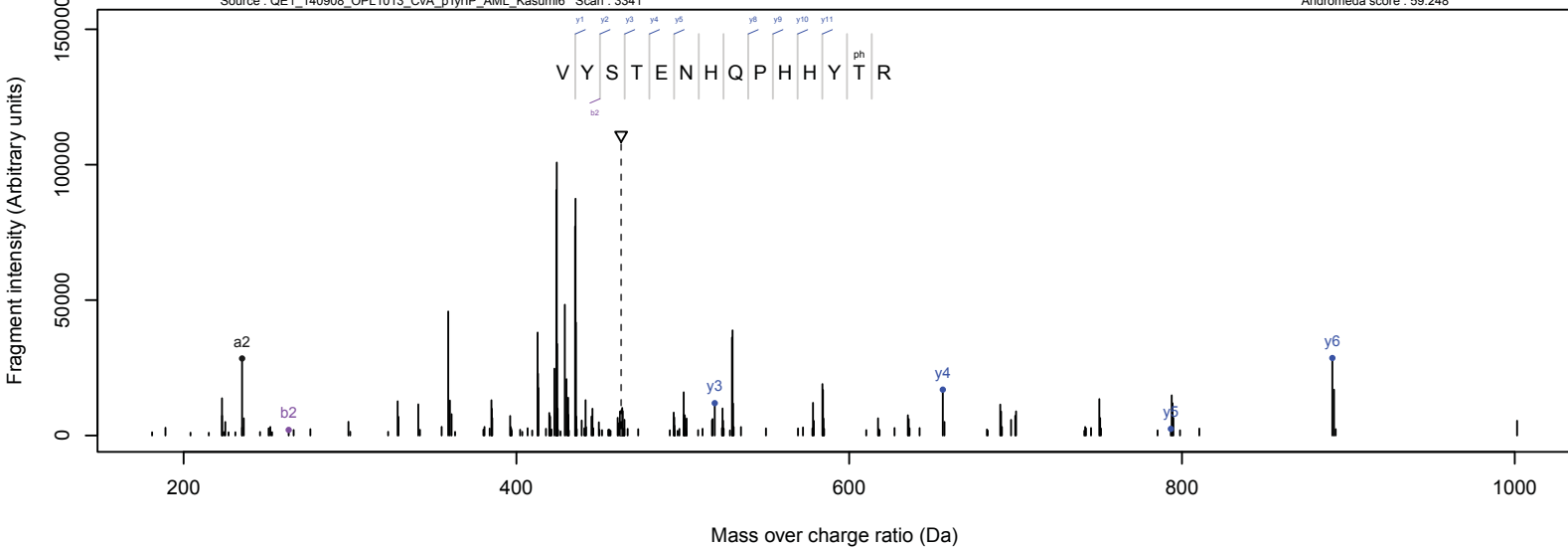
Andromeda score : 59.248



## VYSTENHQPHHYT R-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 3341

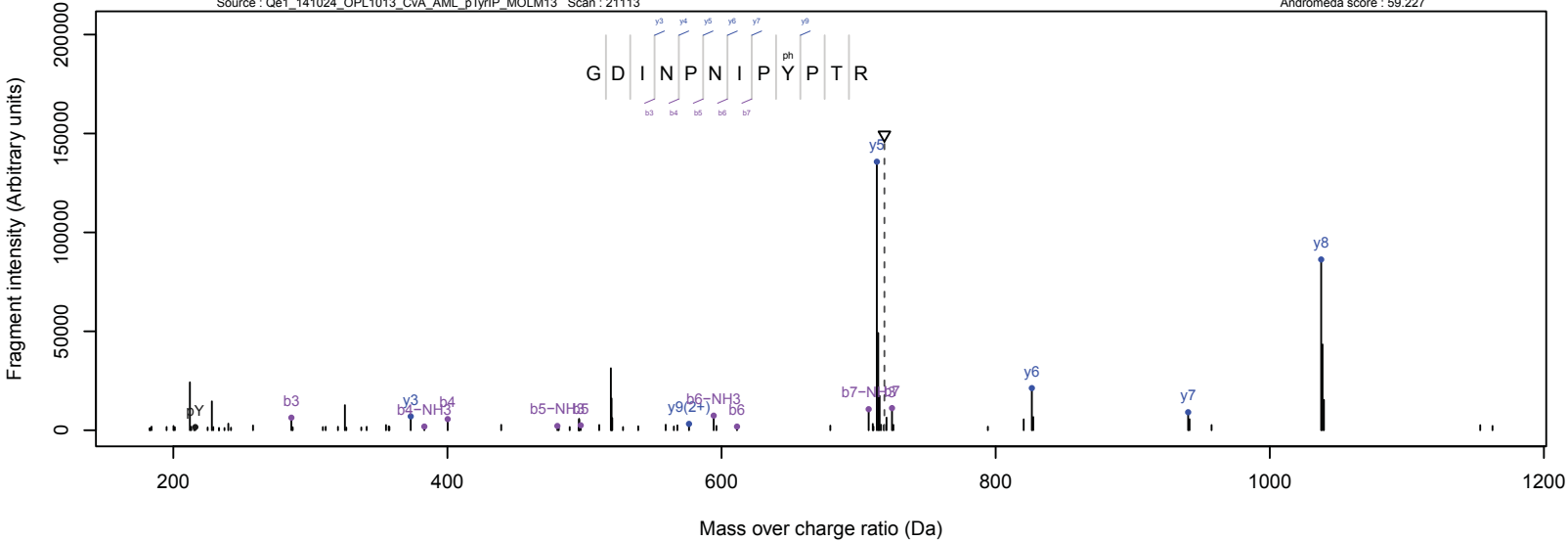
Andromeda score : 59.248

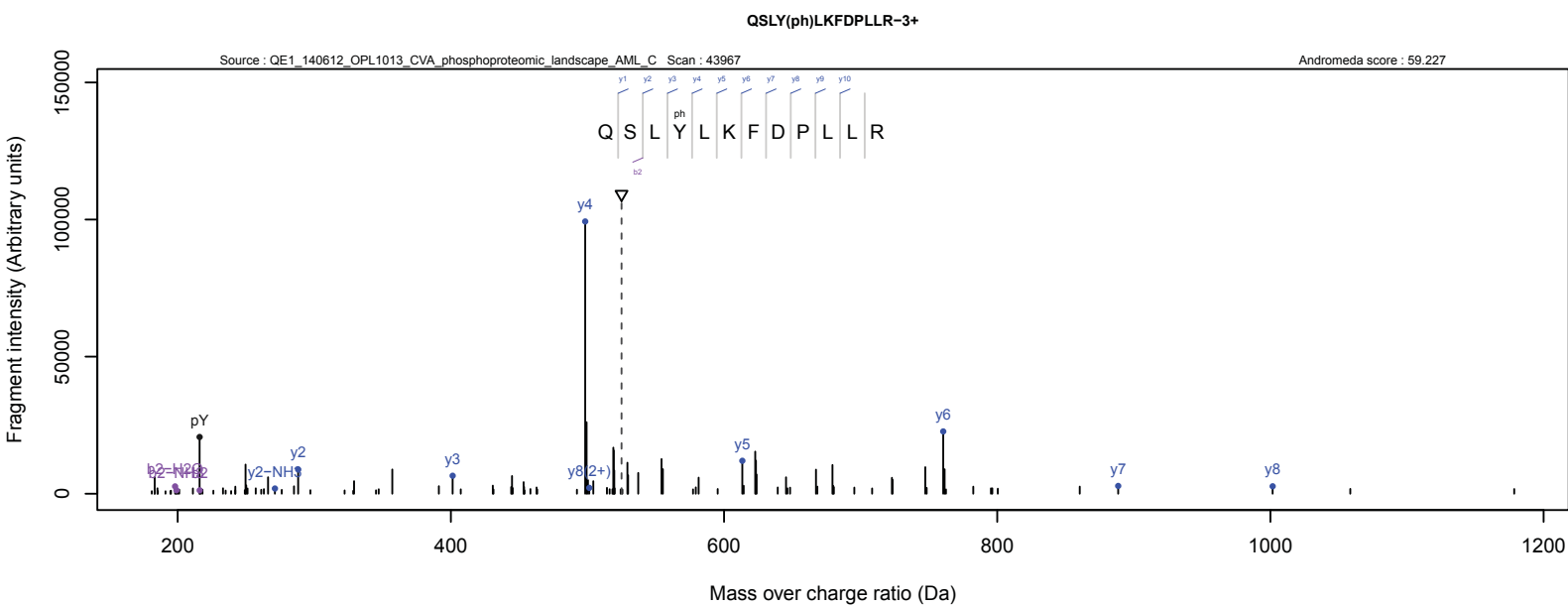
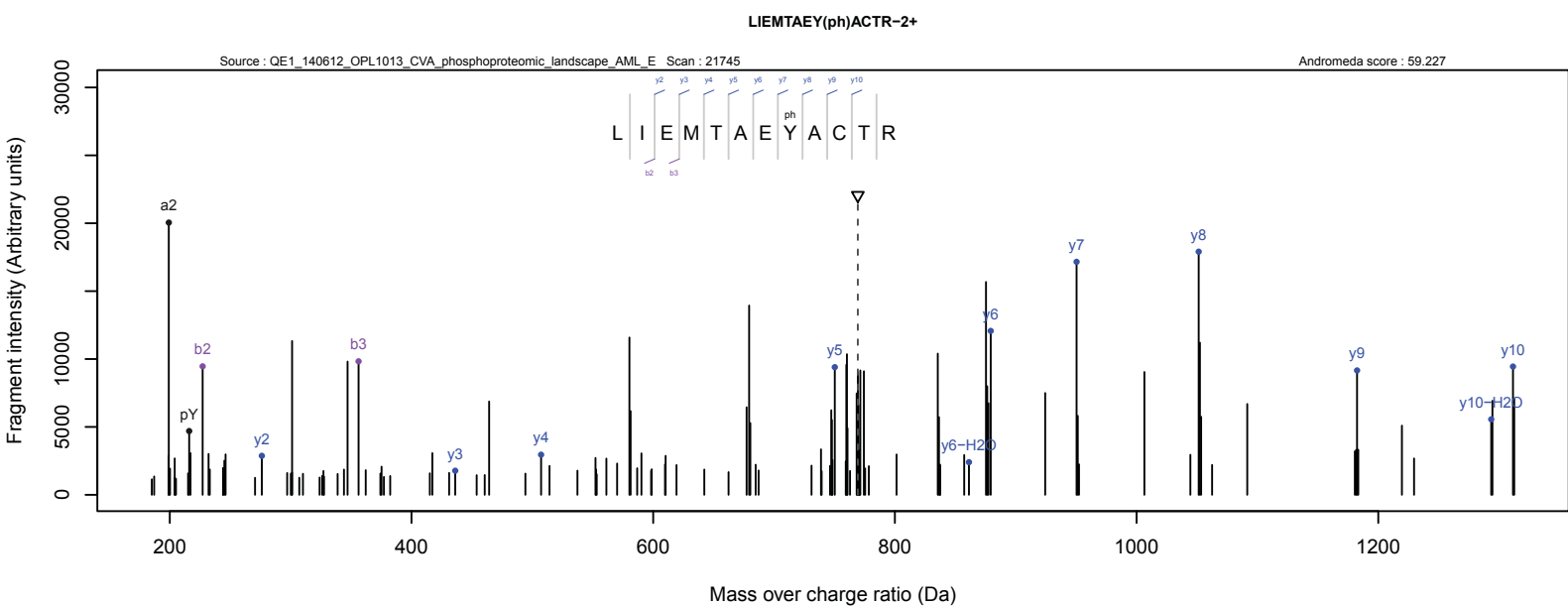
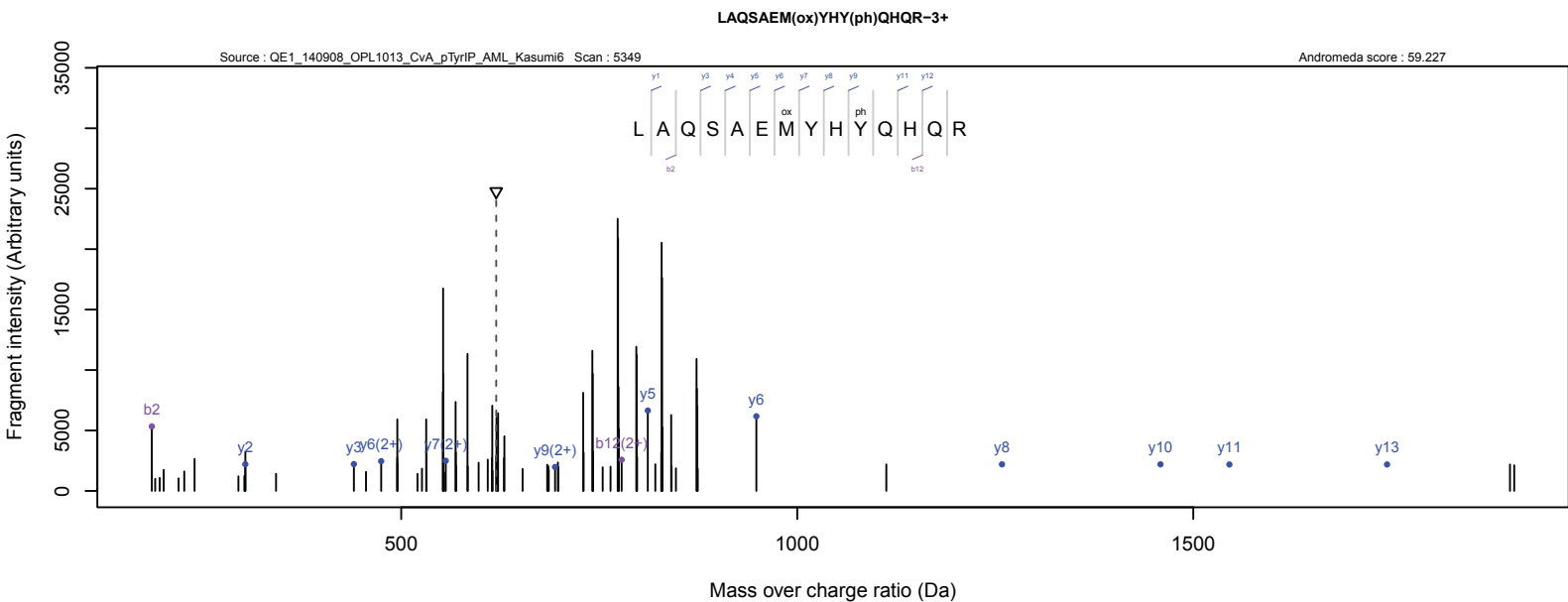


## GDINPNIPY P T R-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 21113

Andromeda score : 59.227



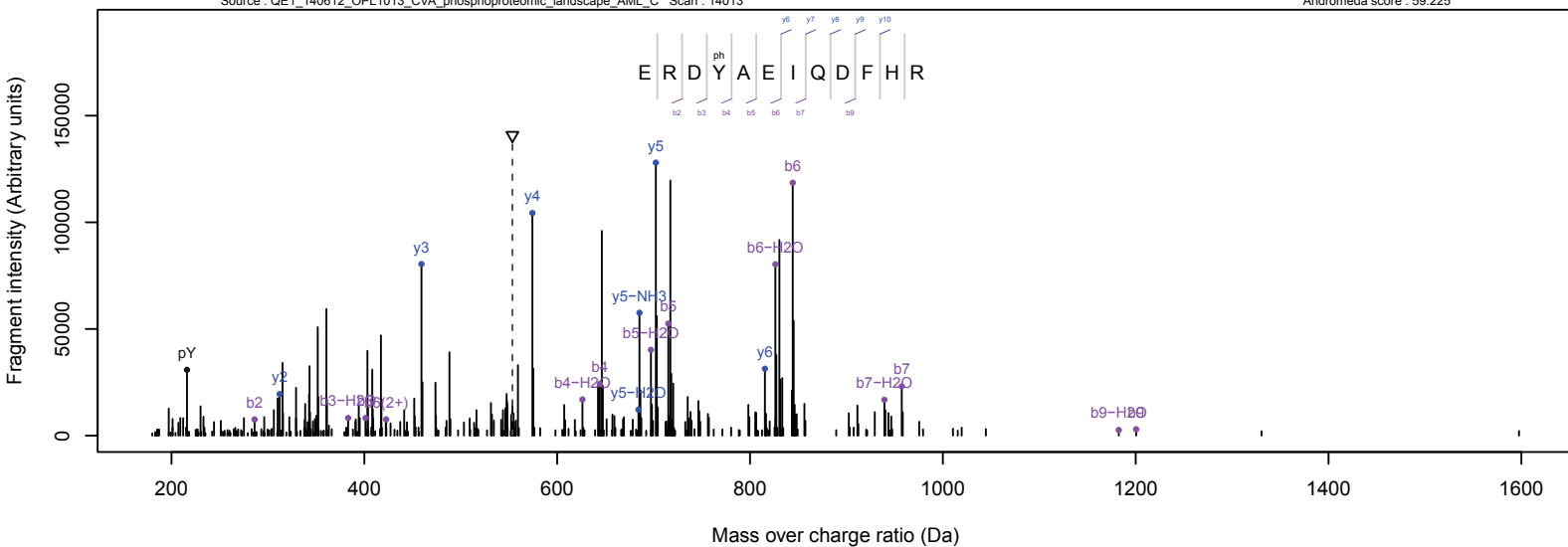




## ERDY(ph)AEIQDFHR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 14013

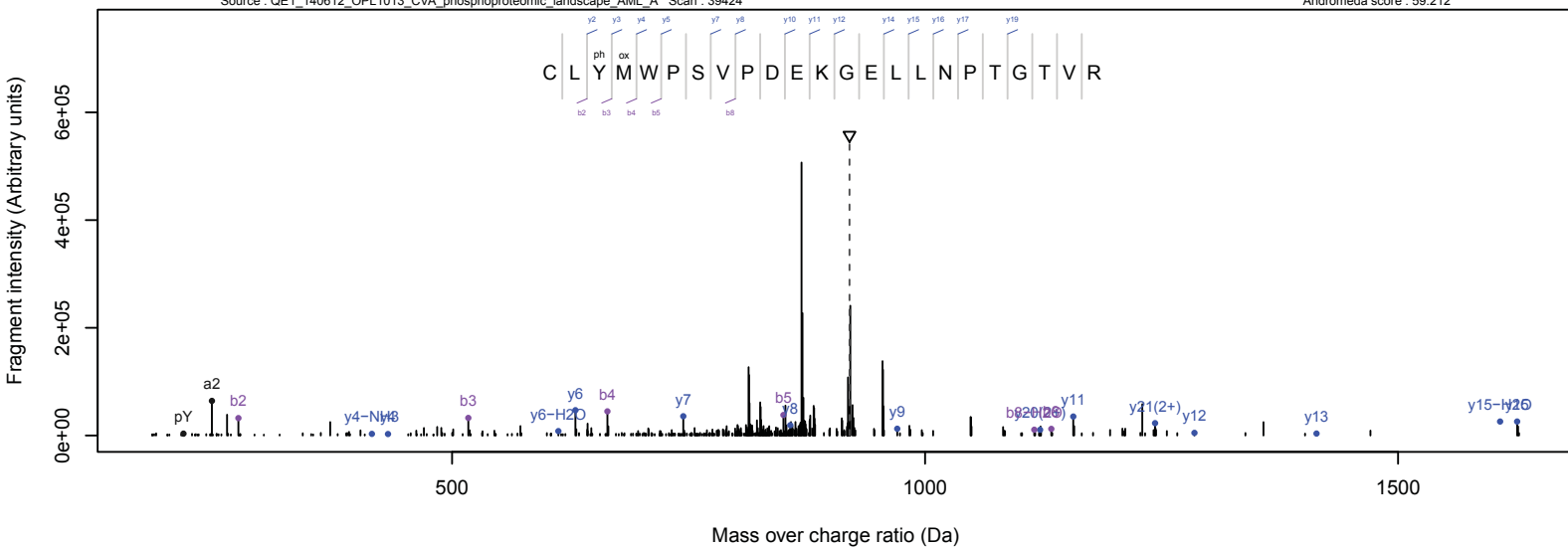
Andromeda score: 59.225



## CLY(ph)M(ox)WPSVPDEKGE LLNPTGTVR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 39424

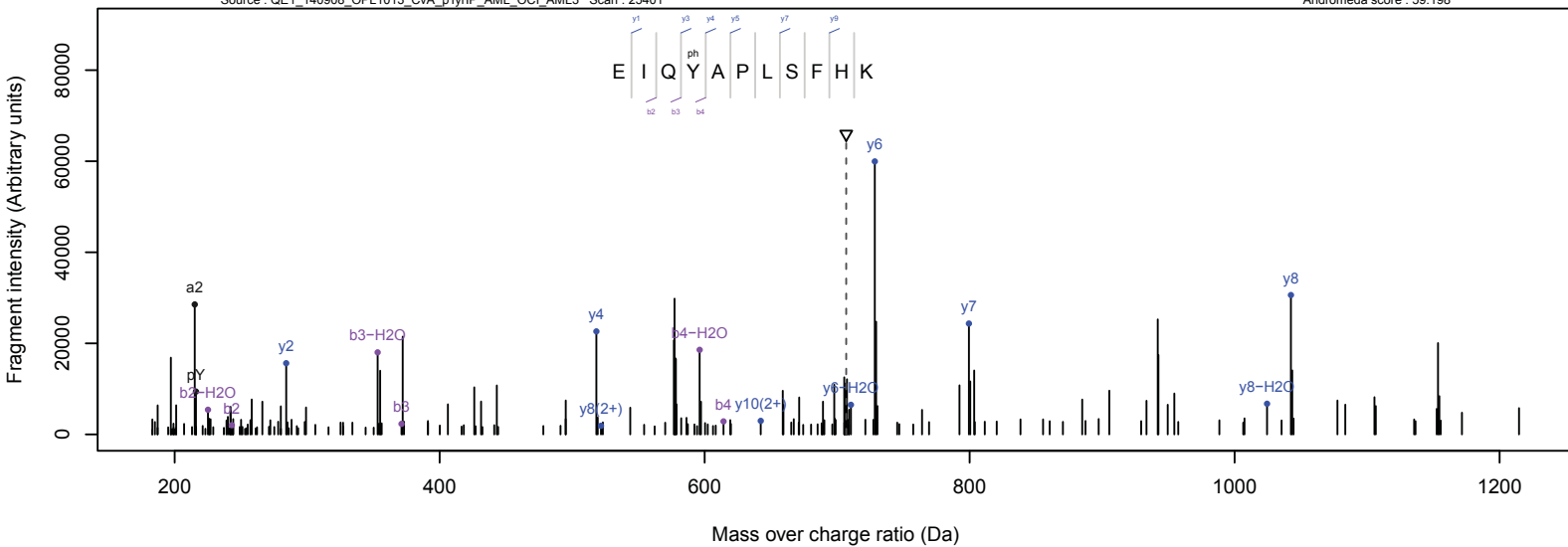
Andromeda score: 59.212



## EIQY(ph)APLSFHK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 25401

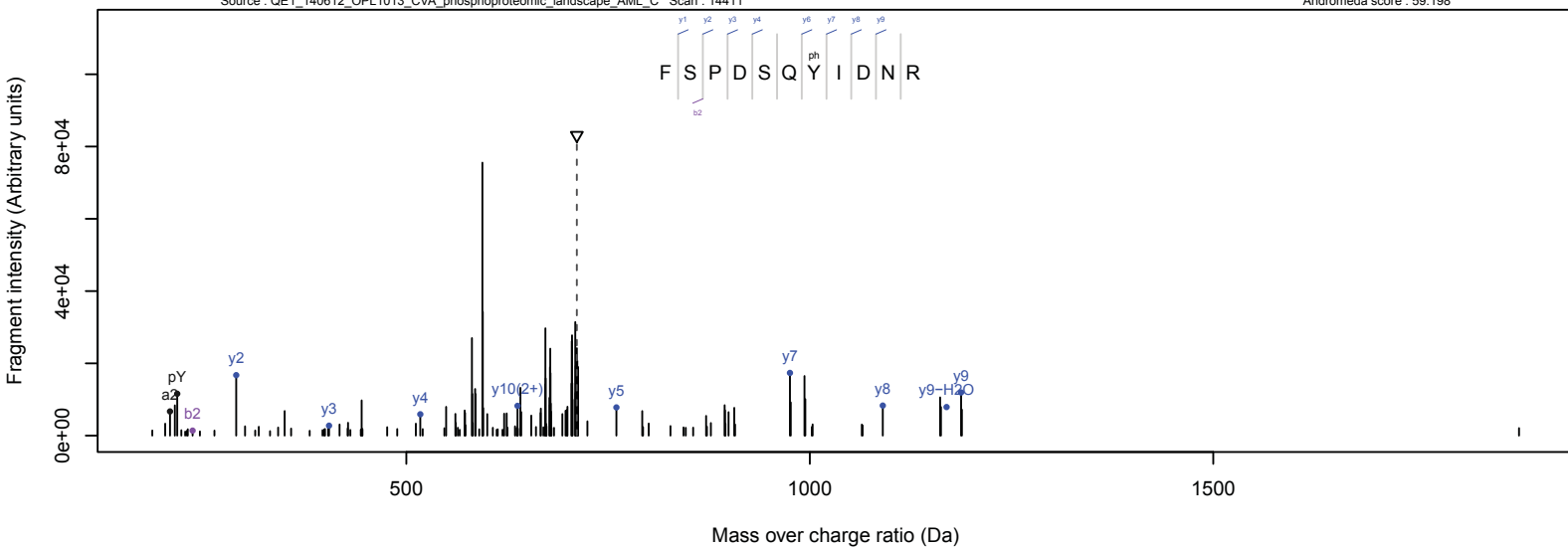
Andromeda score: 59.198



## FSPDSQ(ph)IDNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14411

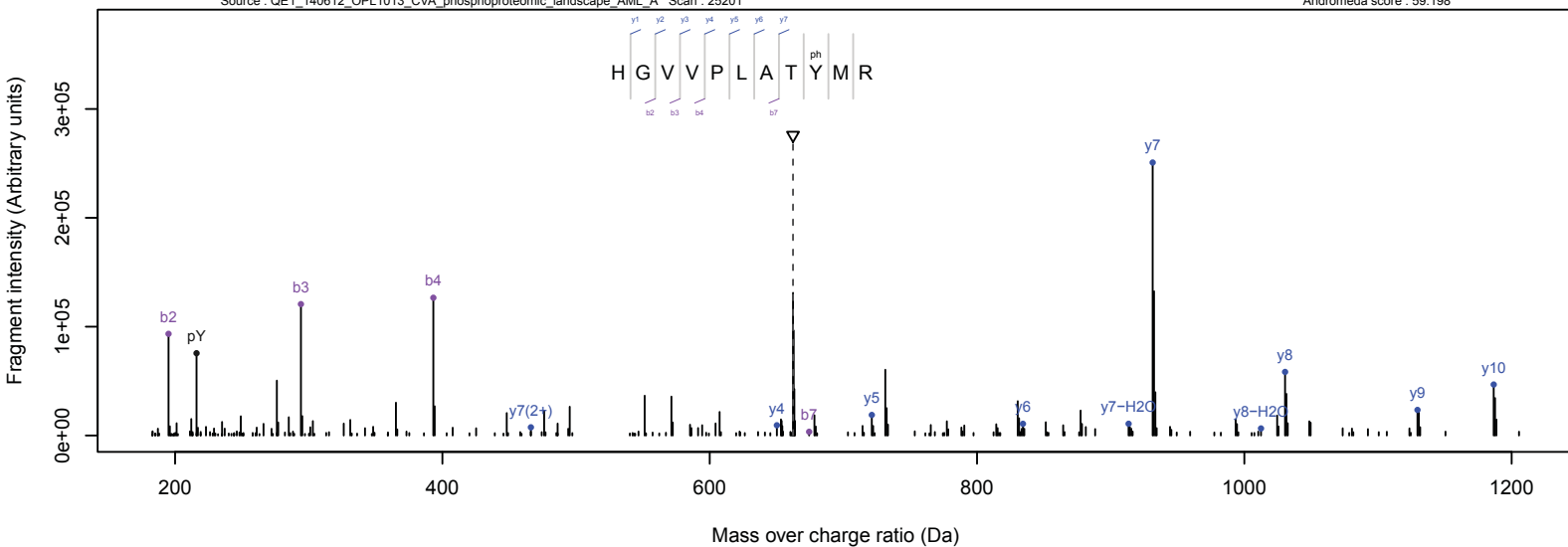
Andromeda score : 59.198



## HGVVPLATY(ph)MR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25201

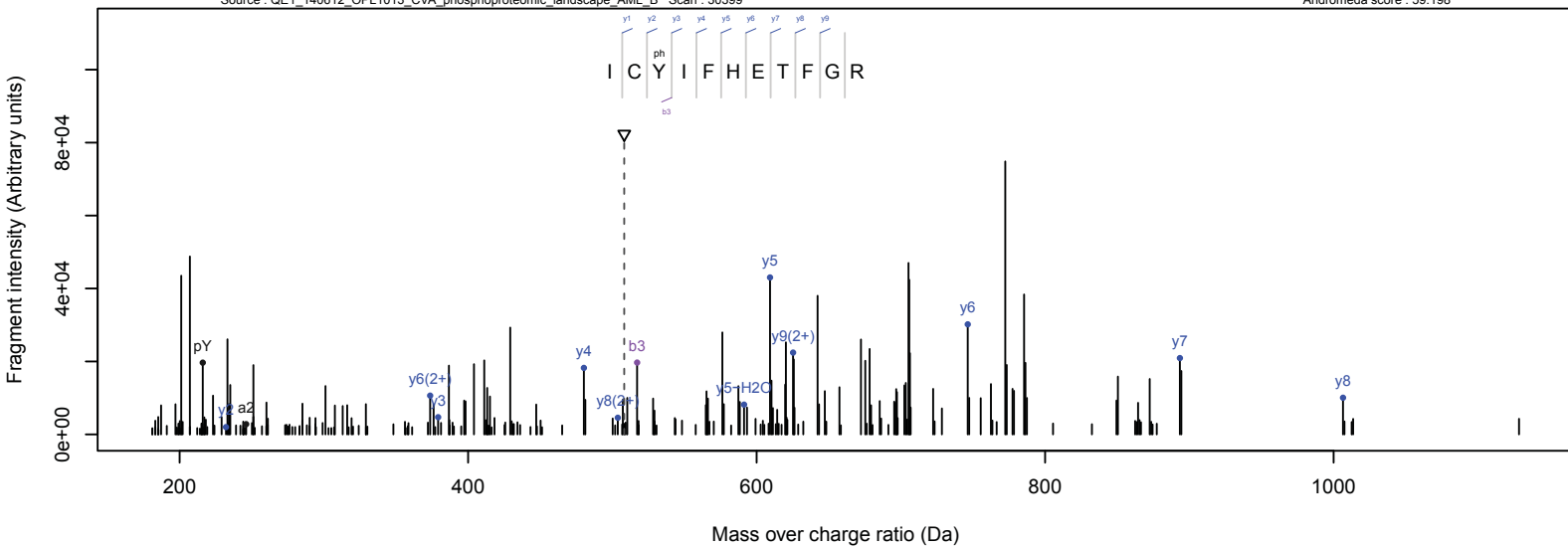
Andromeda score : 59.198



## ICY(ph)IFHETFGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30399

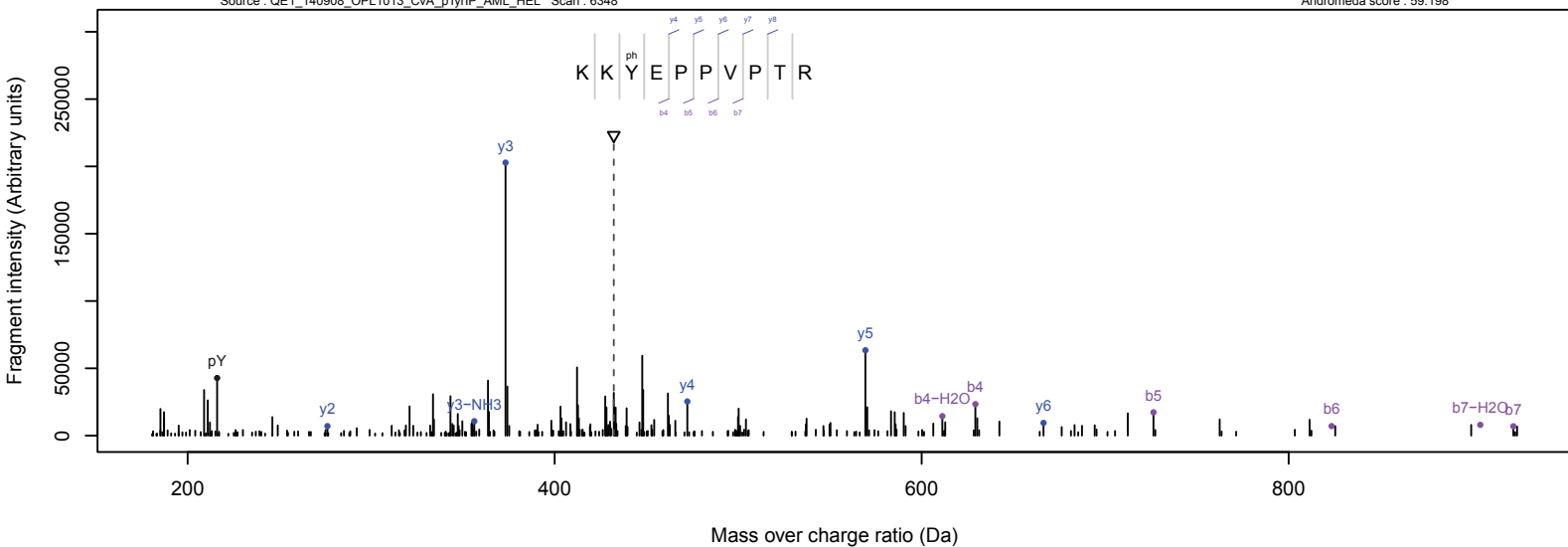
Andromeda score : 59.198



## KKY(ph)EPPVPTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 6348

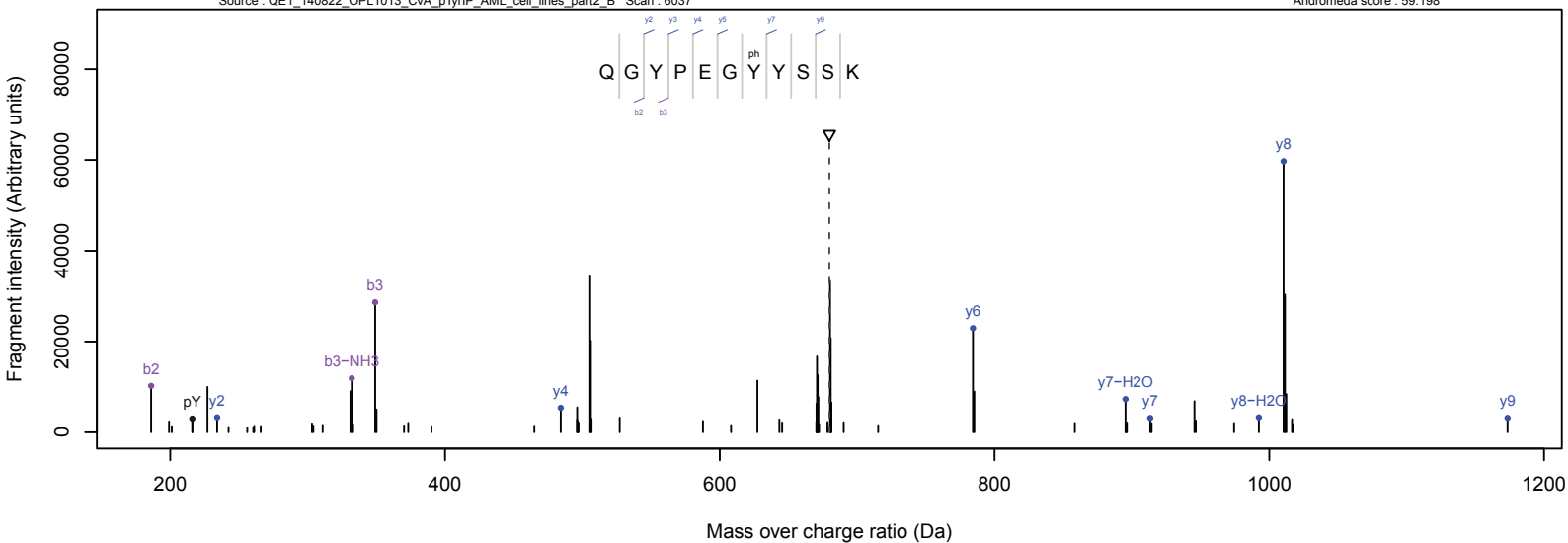
Andromeda score : 59.198



## QGYPEGY(ph)YSSK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 6037

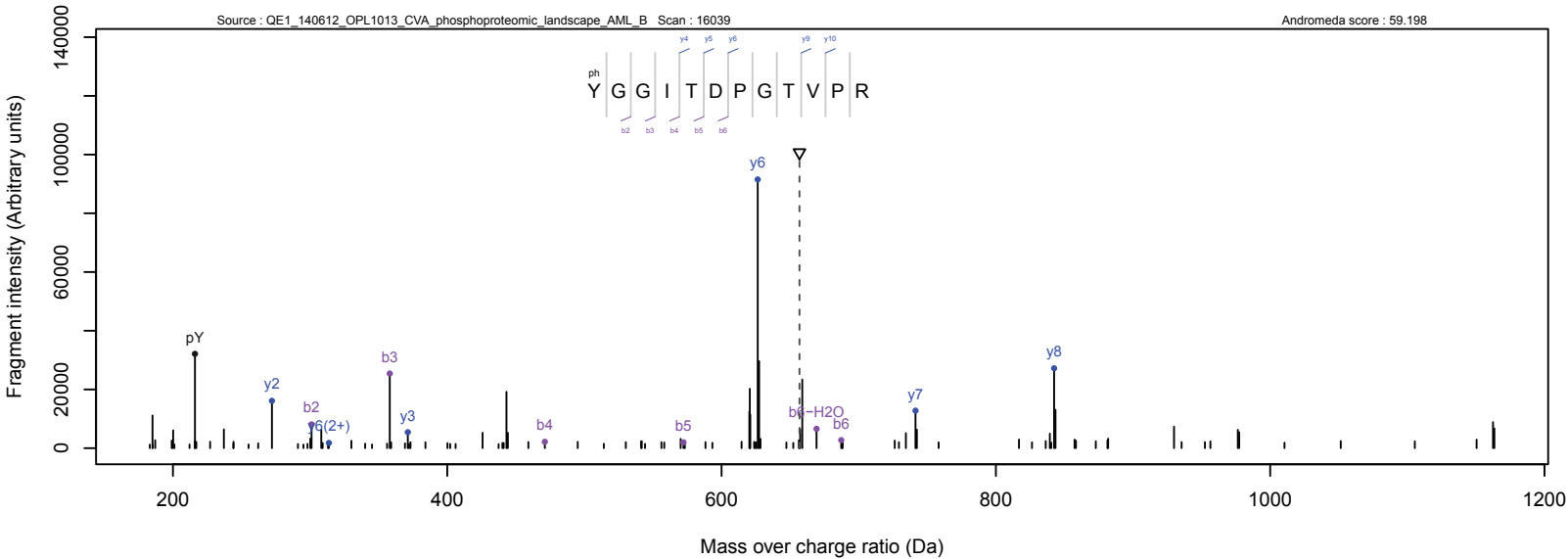
Andromeda score : 59.198



## Y(ph)GGITDPGTVPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16039

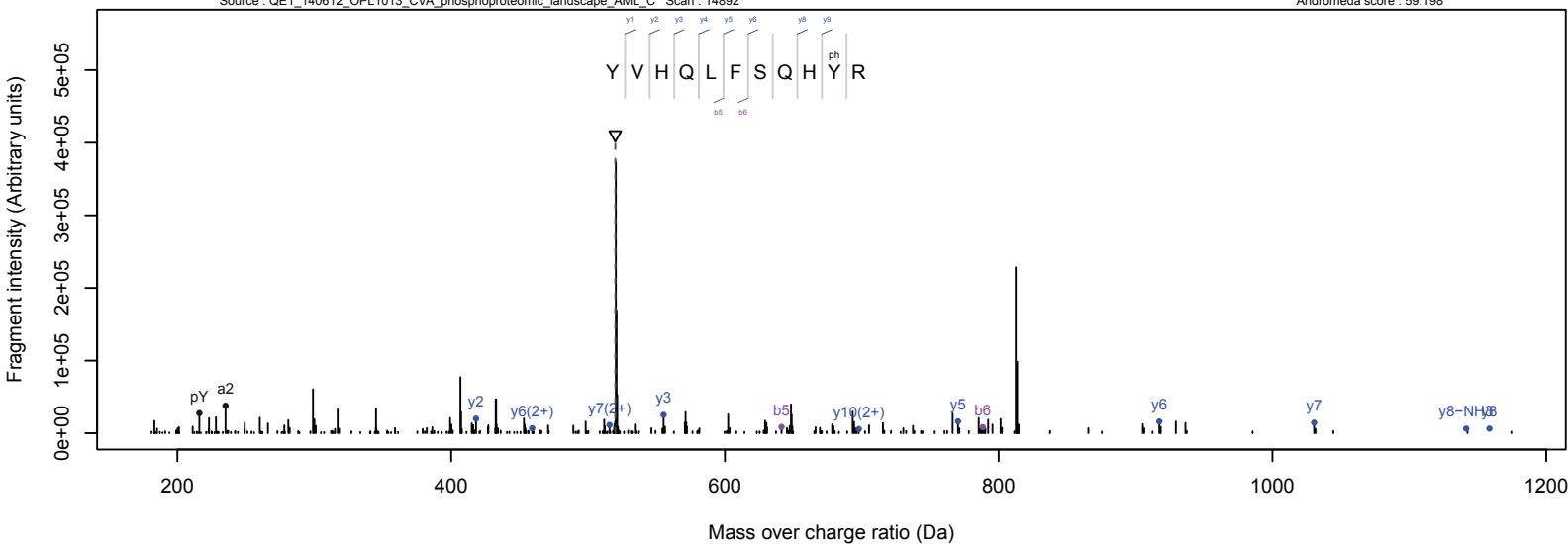
Andromeda score : 59.198



## YVHQLFSQHY(ph)R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14892

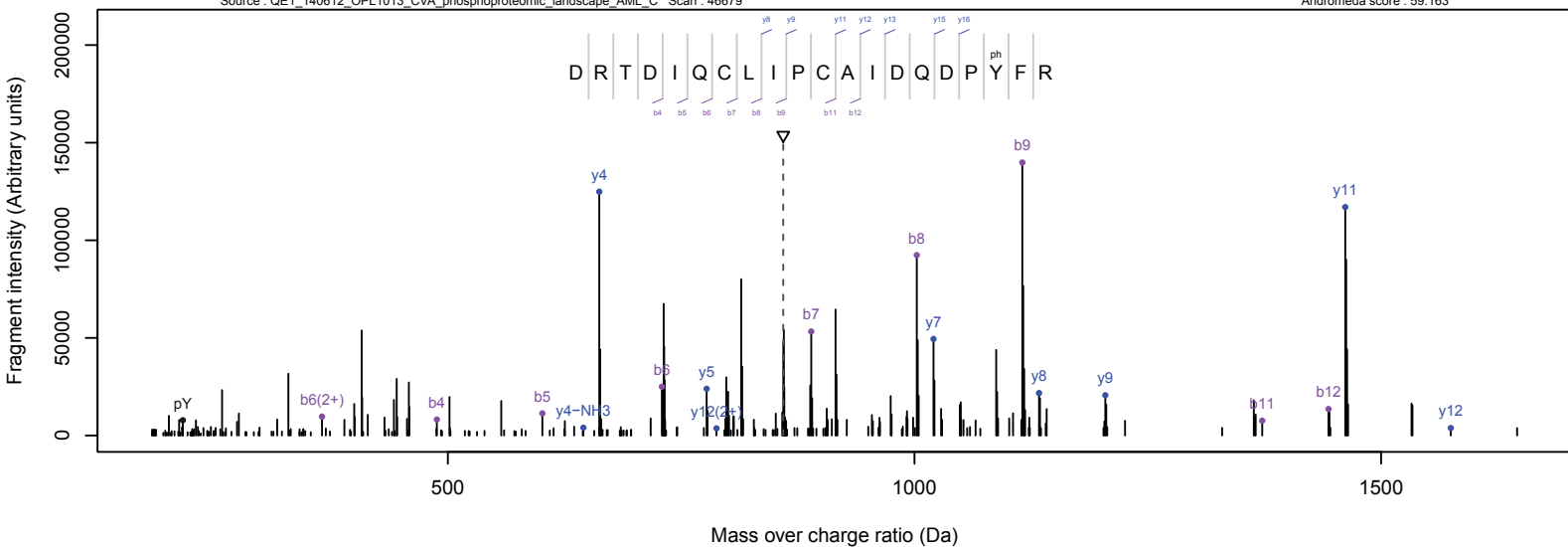
Andromeda score : 59.198



## DRTDIQLIPCAIDQDPY(ph)FR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46679

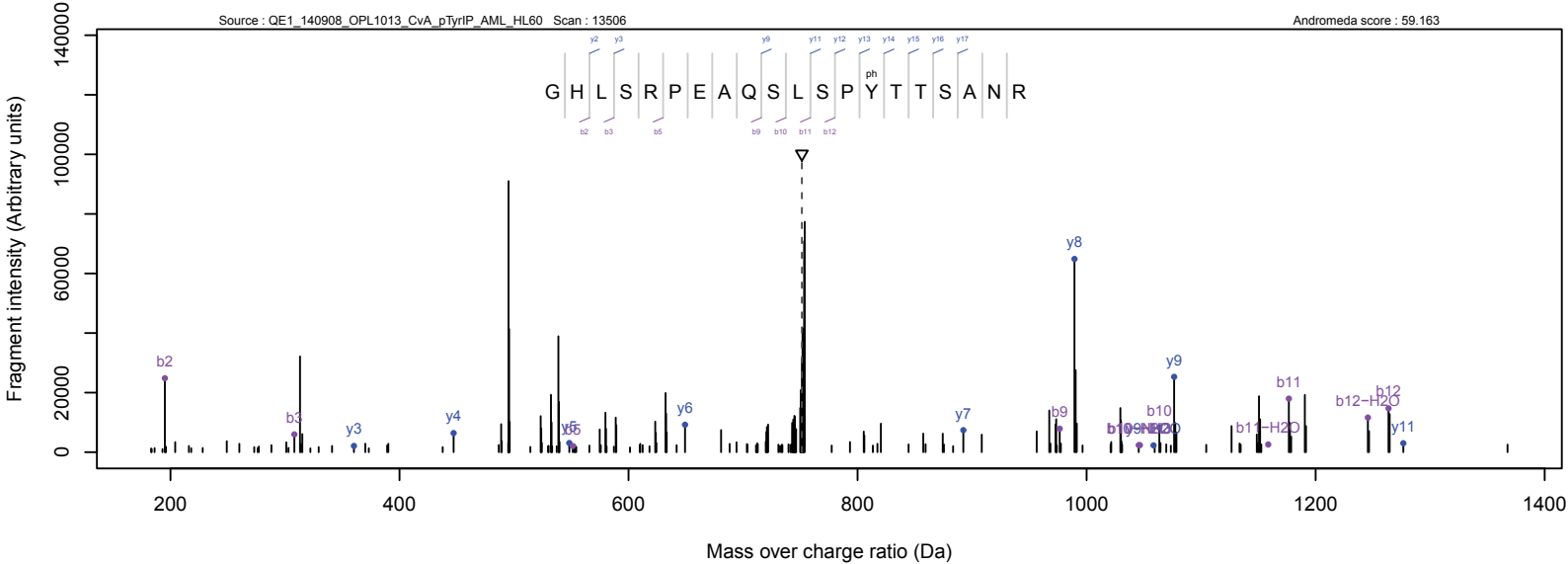
Andromeda score : 59.163



## GHLSRPEAQSLSPYTTSANR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 13506

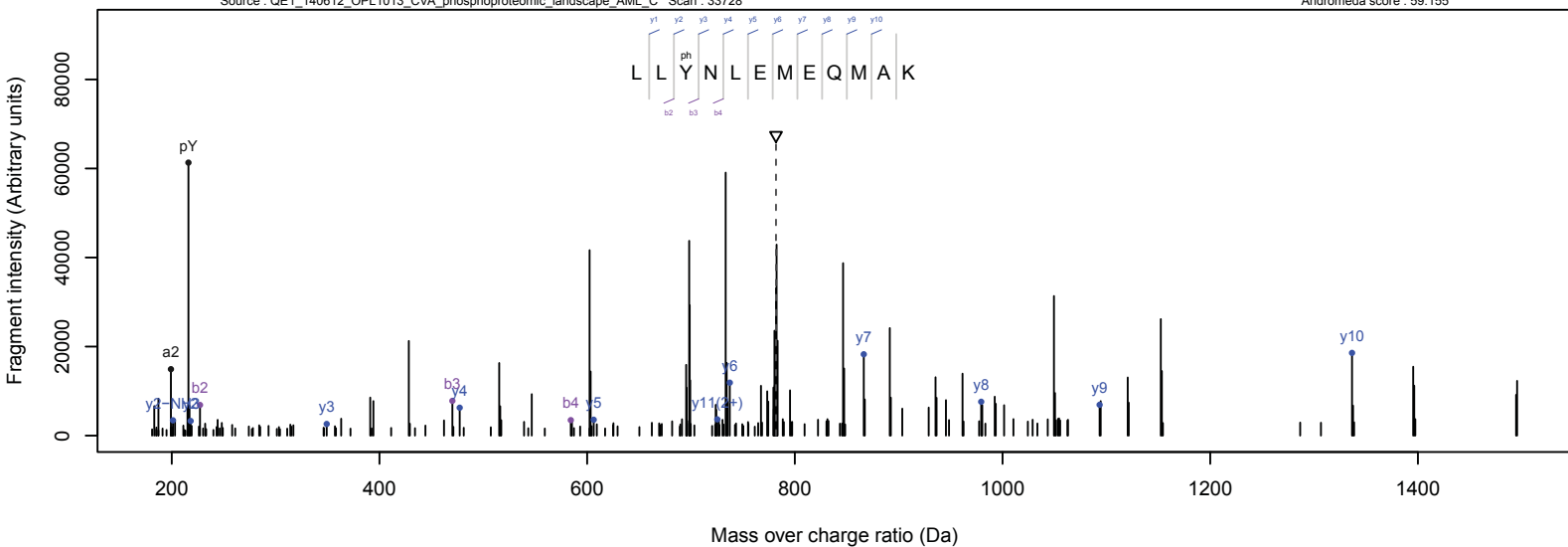
Andromeda score : 59.163



LLY(ph)NLEMEQMAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33728

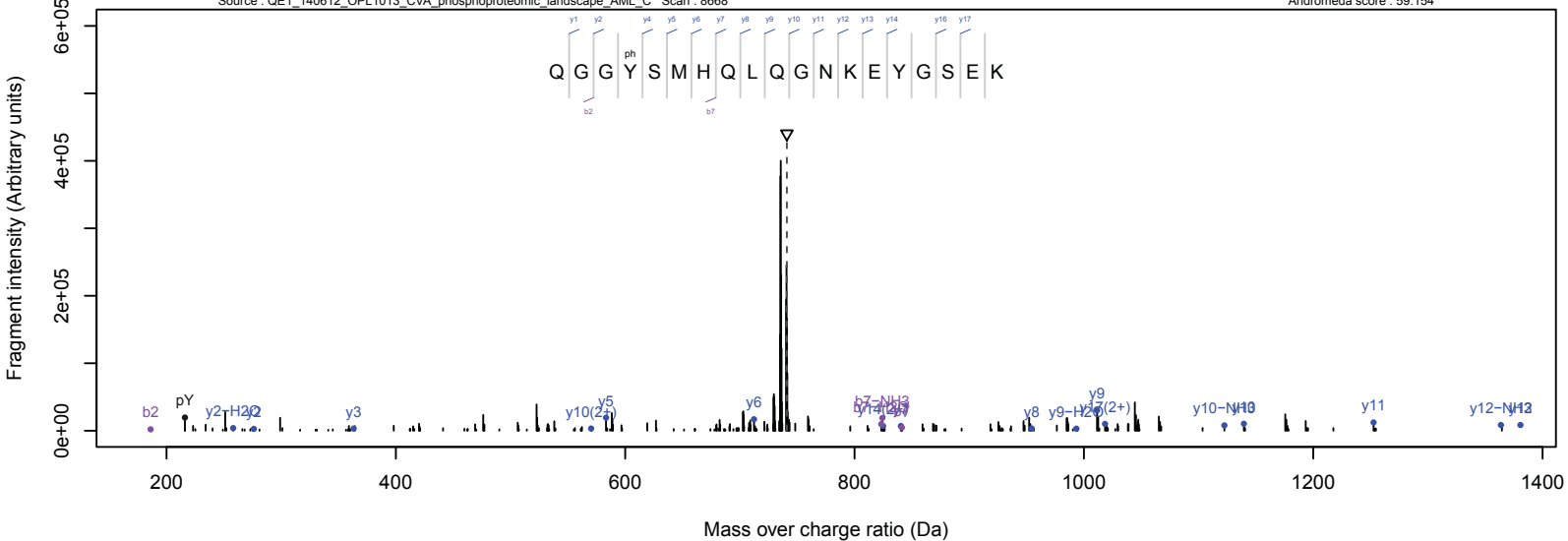
Andromeda score : 59.155



QGGY(ph)SMHQLQGNKEYGSEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8668

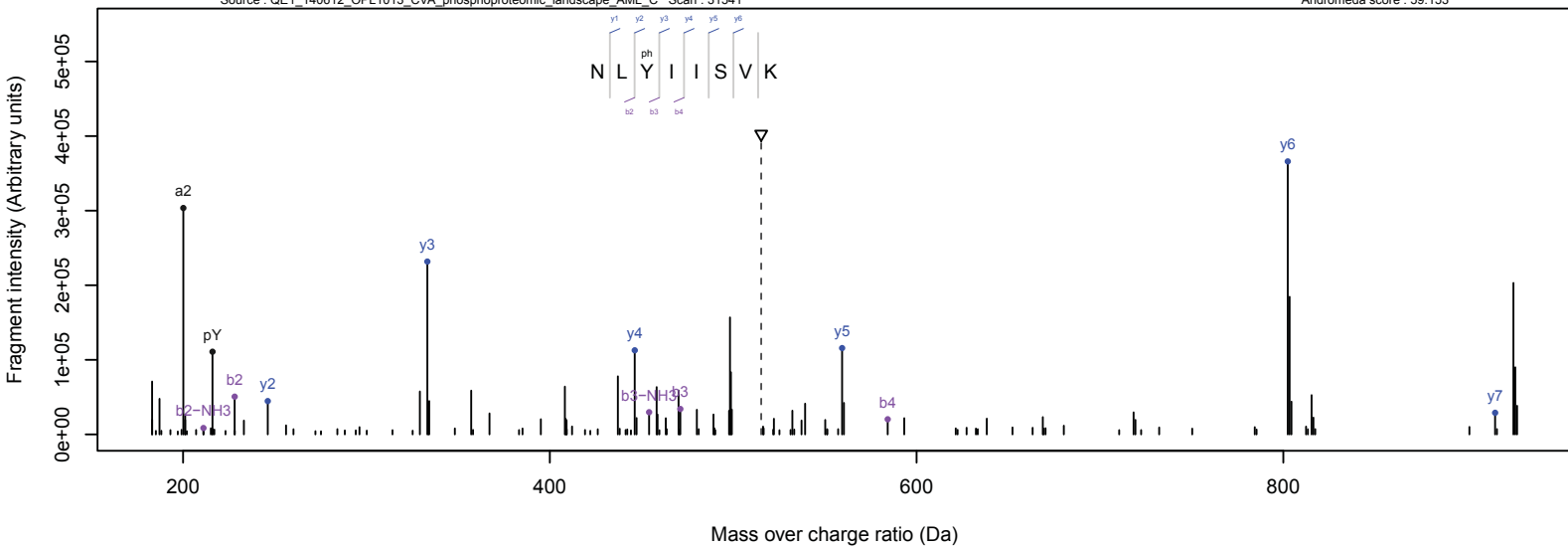
Andromeda score : 59.154



NLY(ph)IISVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31541

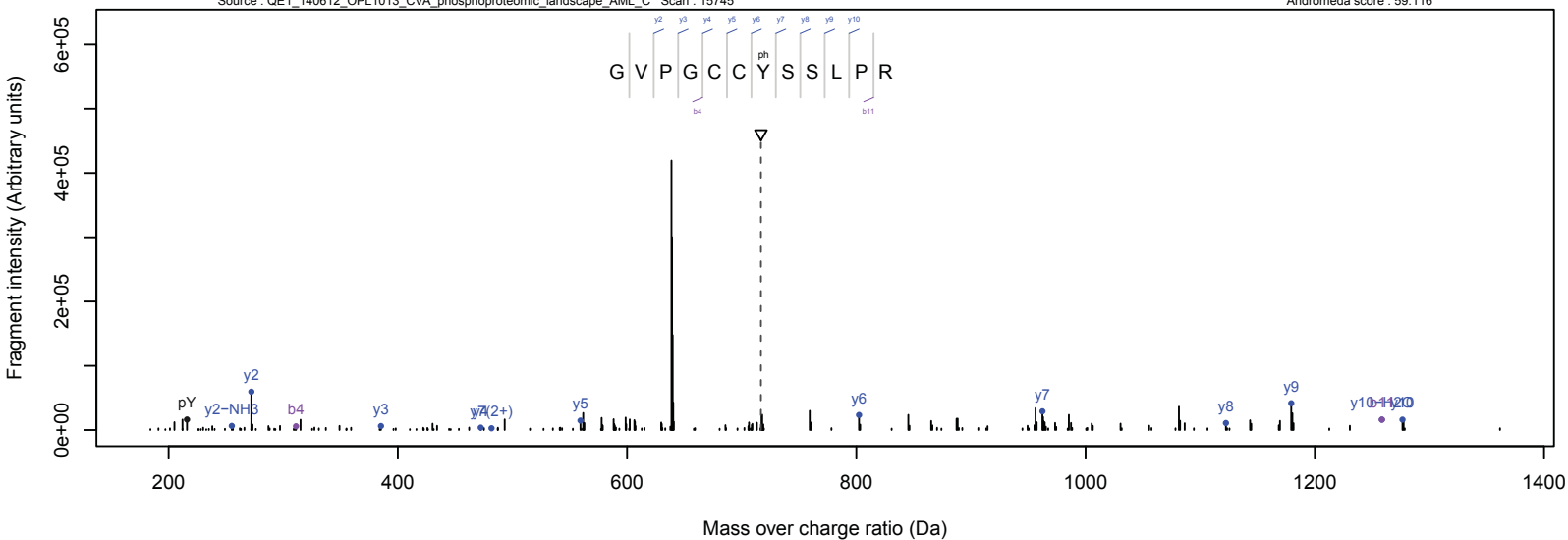
Andromeda score : 59.153



## GVPGCCY(ph)SSLPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15745

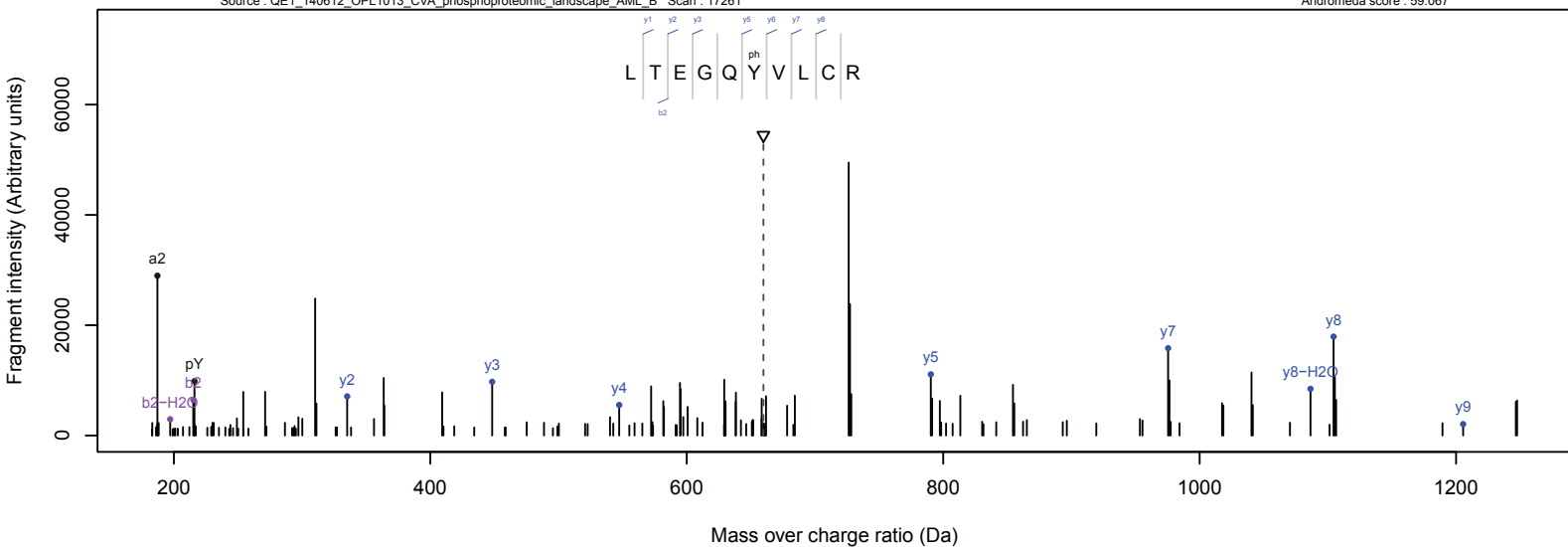
Andromeda score: 59.116



## LTEGQY(ph)VLCR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 17261

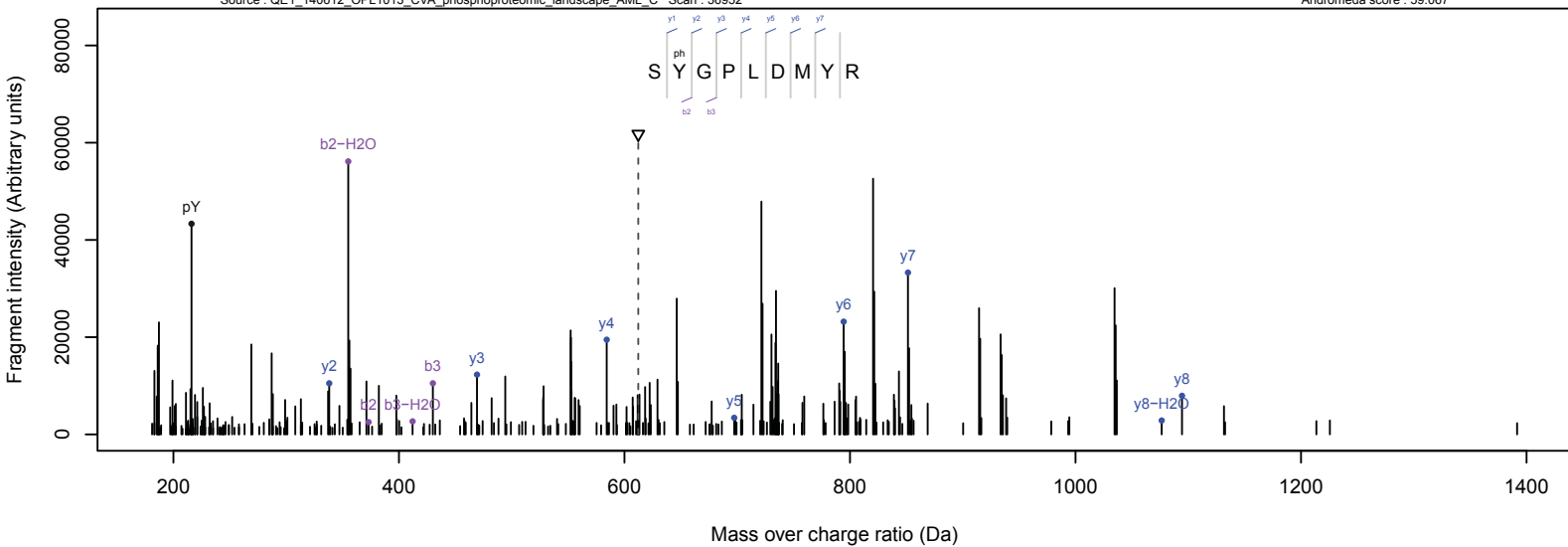
Andromeda score: 59.067



## (ac)SY(ph)GPLDMYR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36952

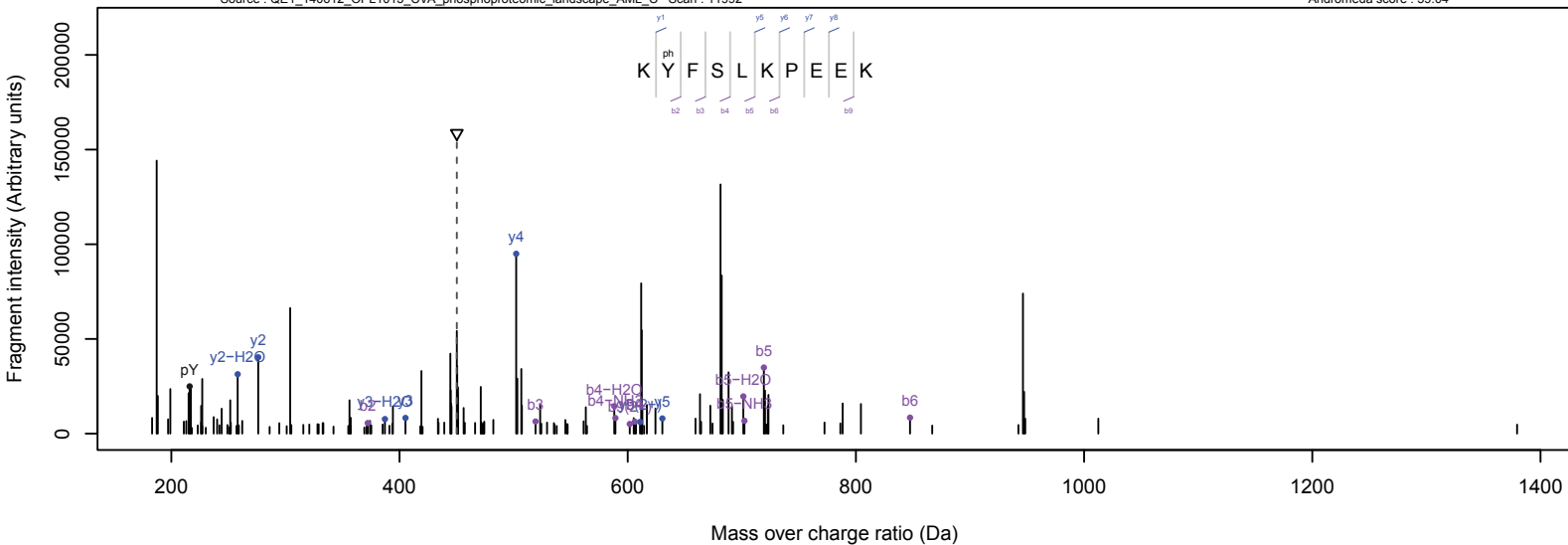
Andromeda score: 59.067



**KY(ph)FSLKPEEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11992

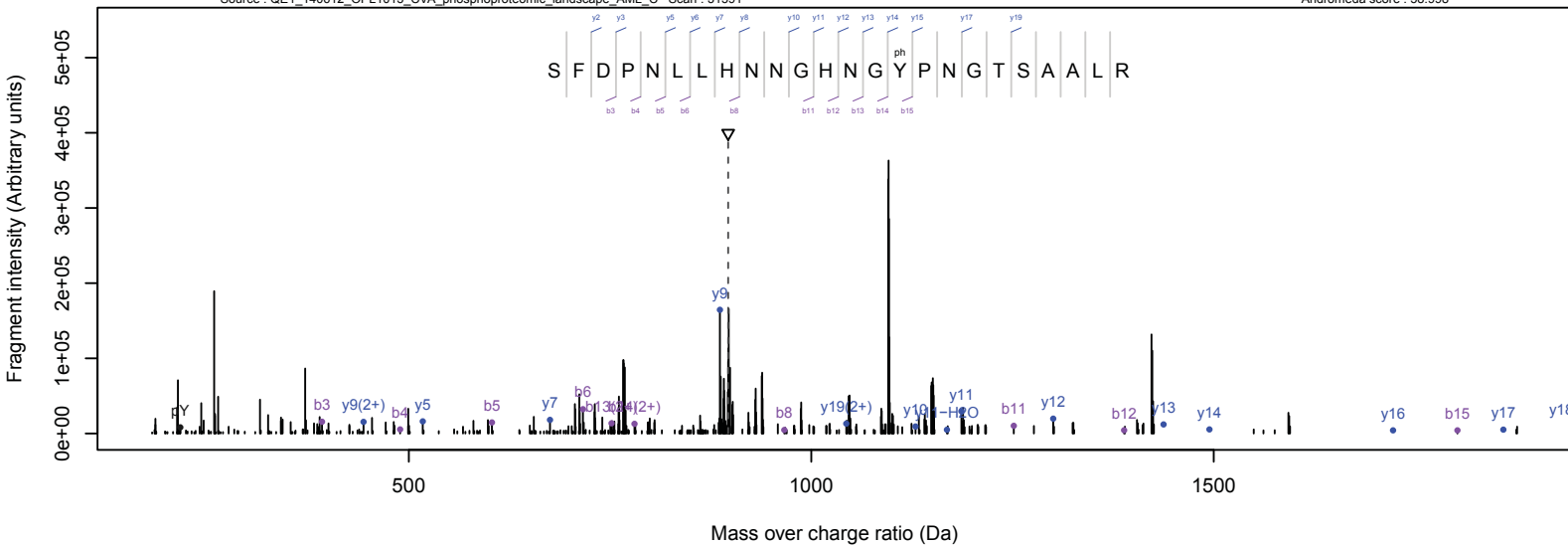
Andromeda score : 59.04



**(ac)SFDPNLLHNNHNGY(ph)PNGTSAALR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31591

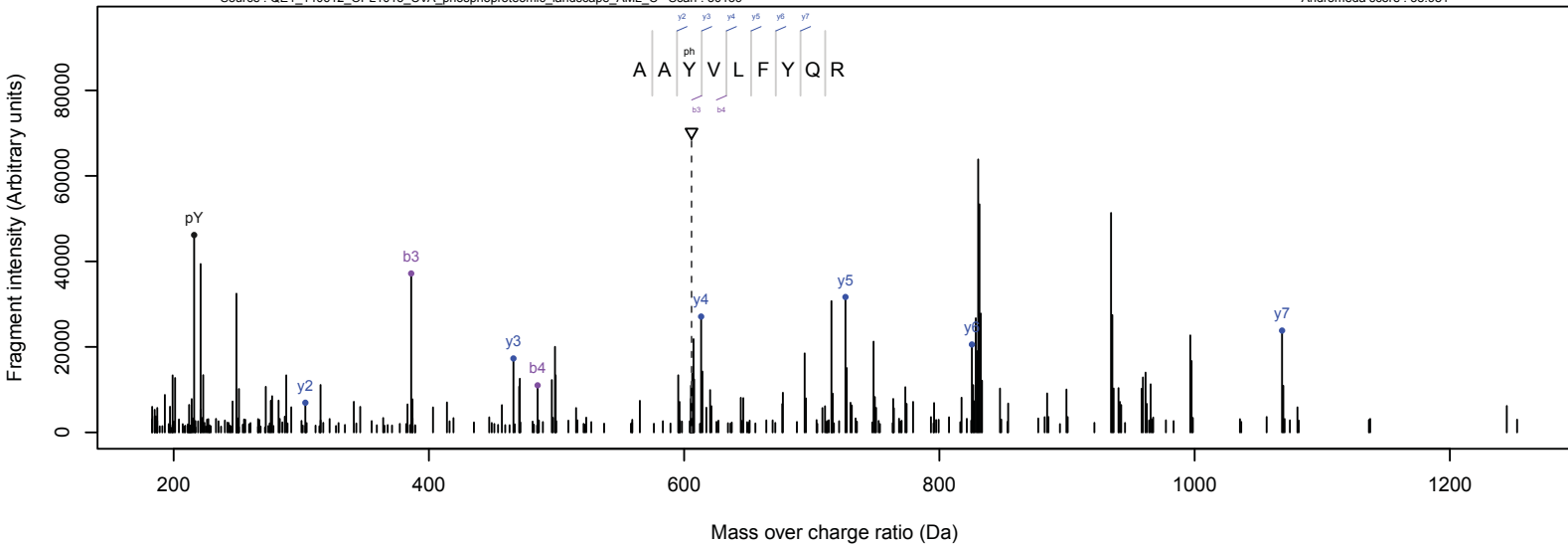
Andromeda score : 58.998



**AA Y(ph)VLFYQR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30139

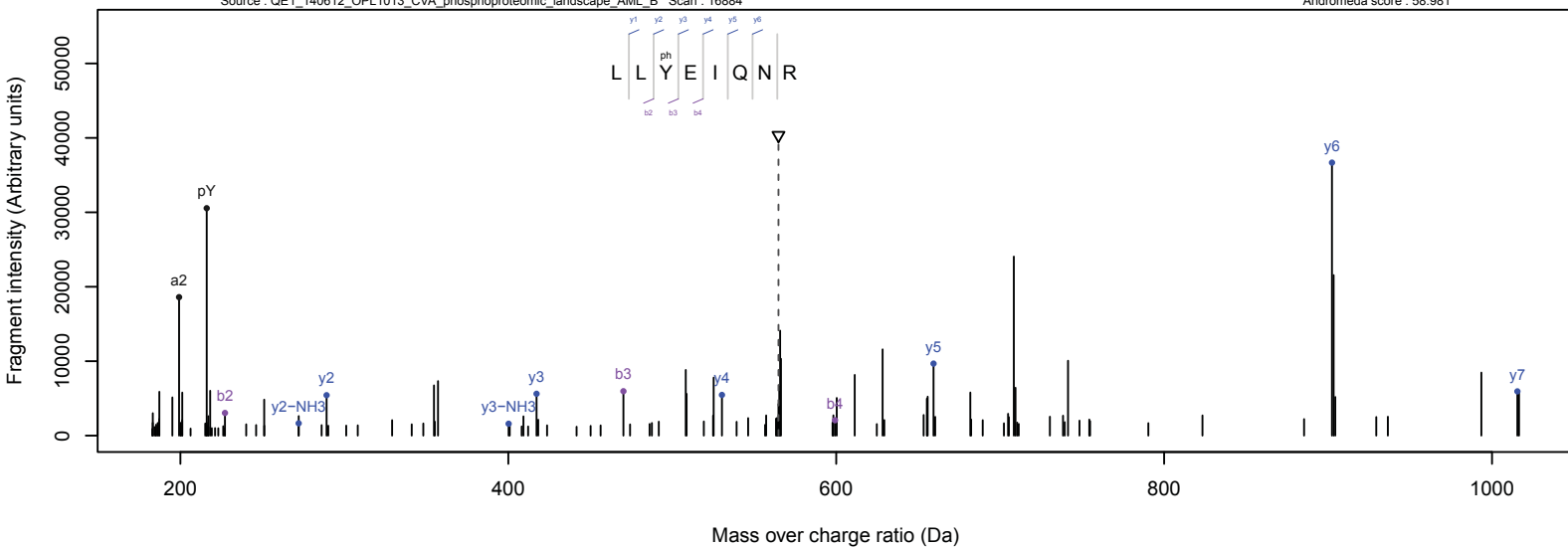
Andromeda score : 58.981



## LLY(ph)EIQNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16884

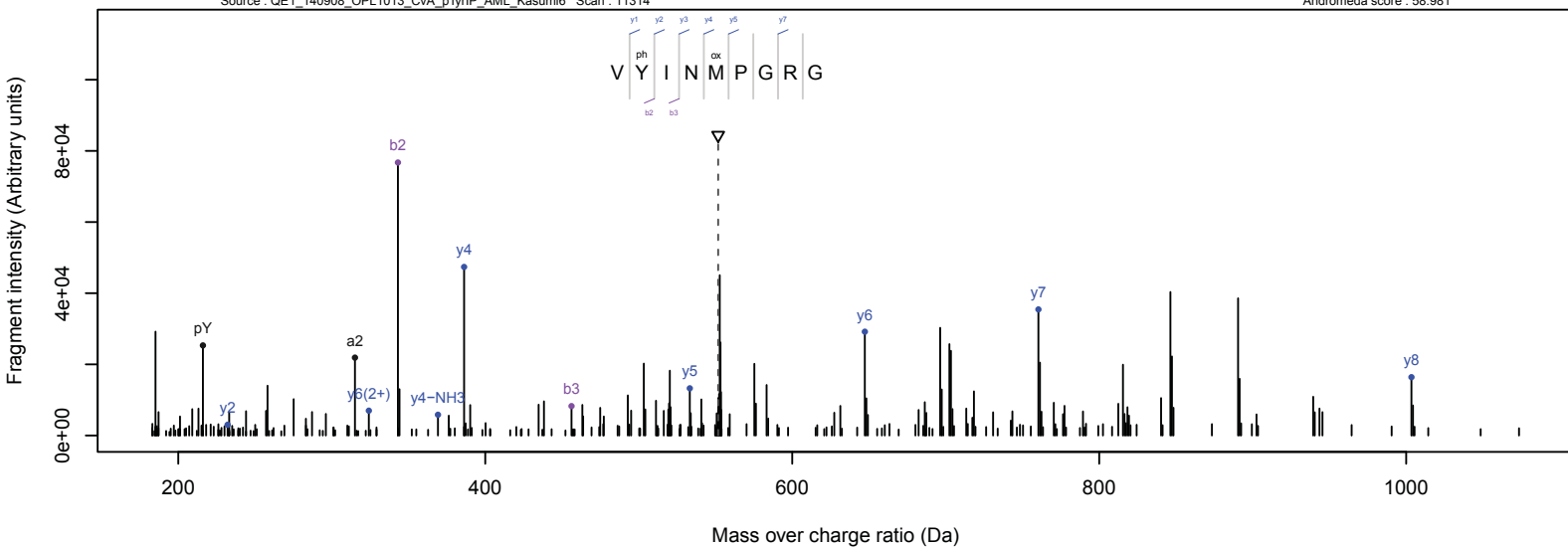
Andromeda score : 58.981



## VY(ph)INM(ox)PGRG-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 11314

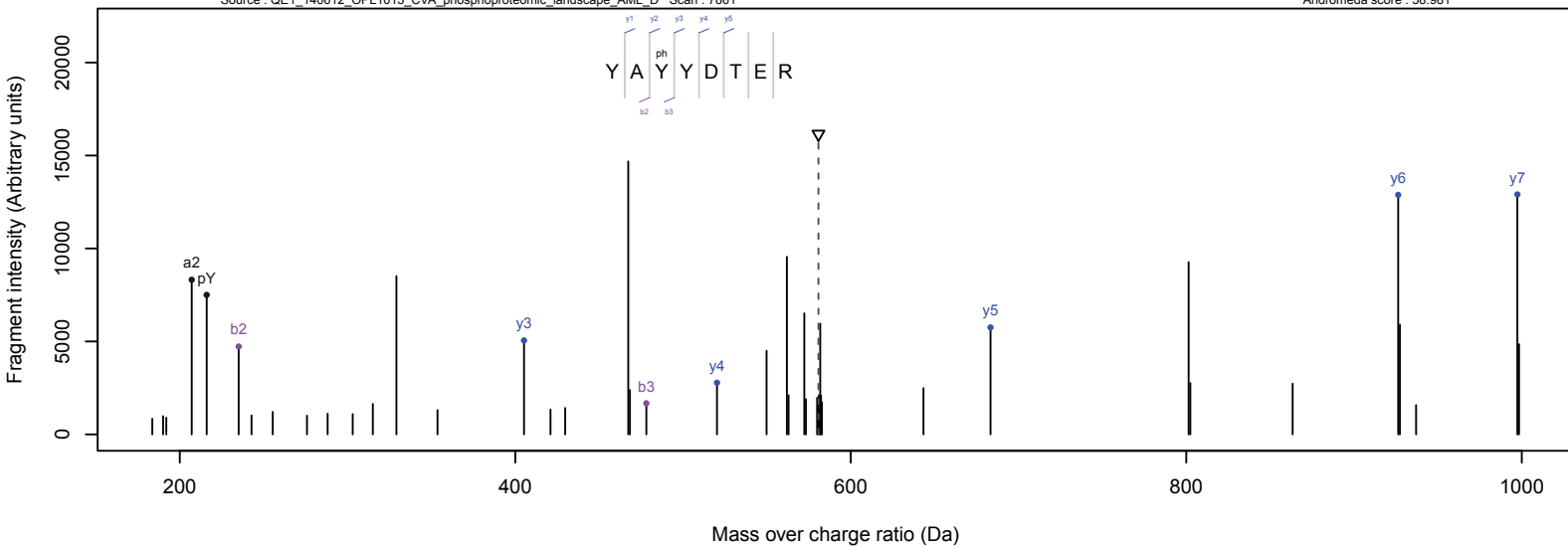
Andromeda score : 58.981



## YAY(ph)YDTER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 7861

Andromeda score : 58.981

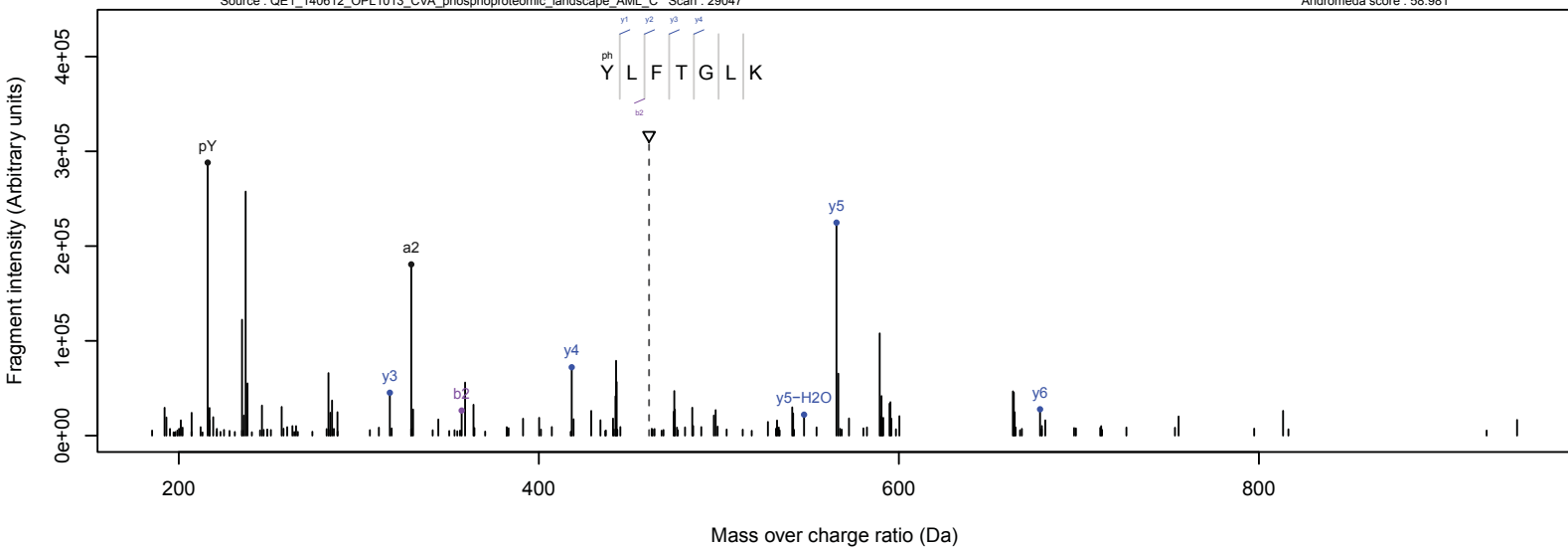




## Y(ph)LFTGLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29047

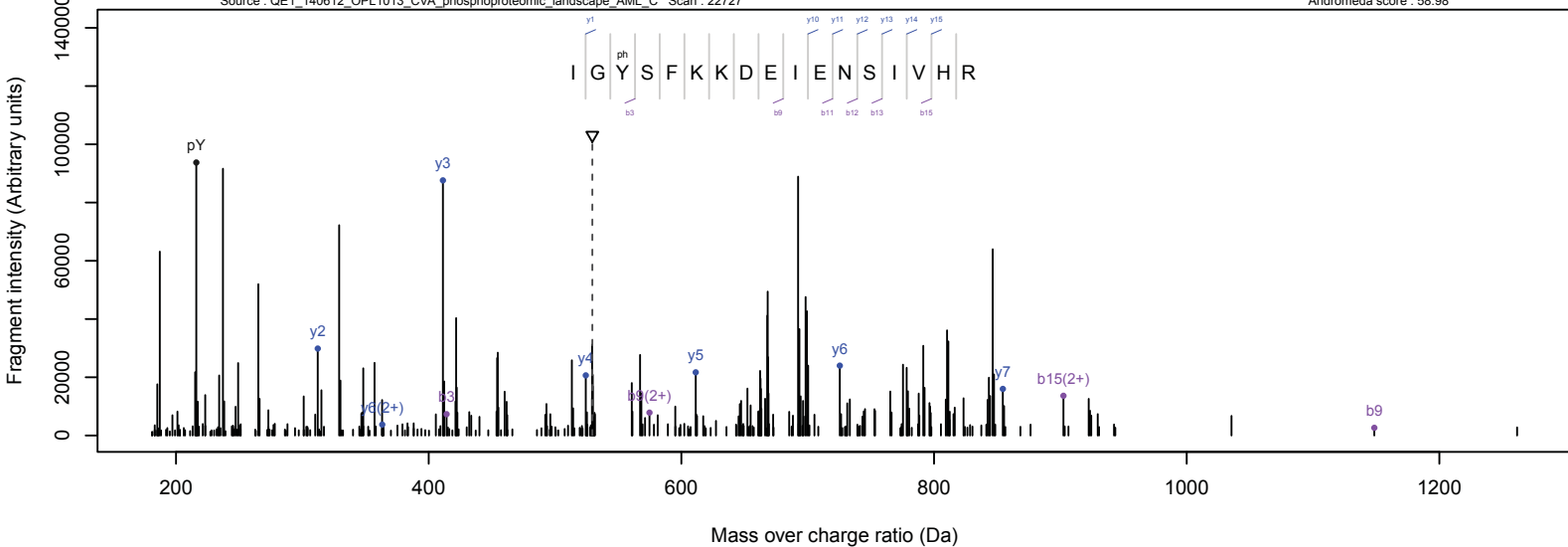
Andromeda score : 58.981



## IGY(ph)SFKKDEIENSIVHR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22727

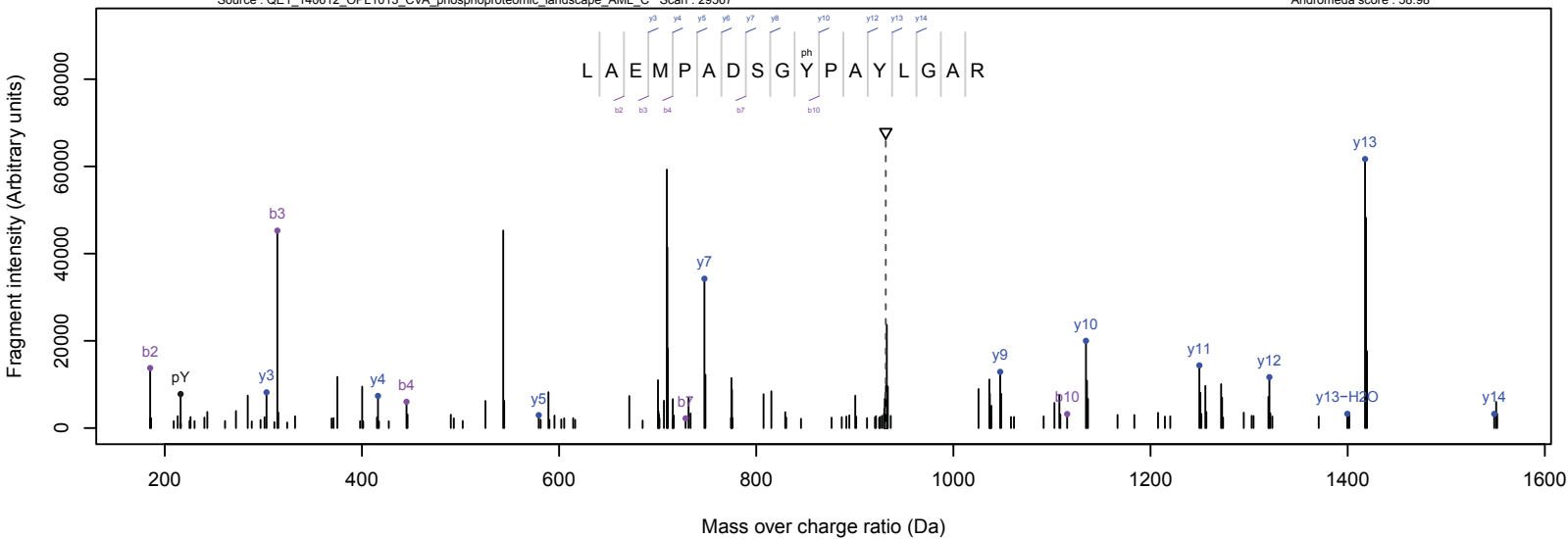
Andromeda score : 58.98



## LAEMPADSGY(ph)PAYLGAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29567

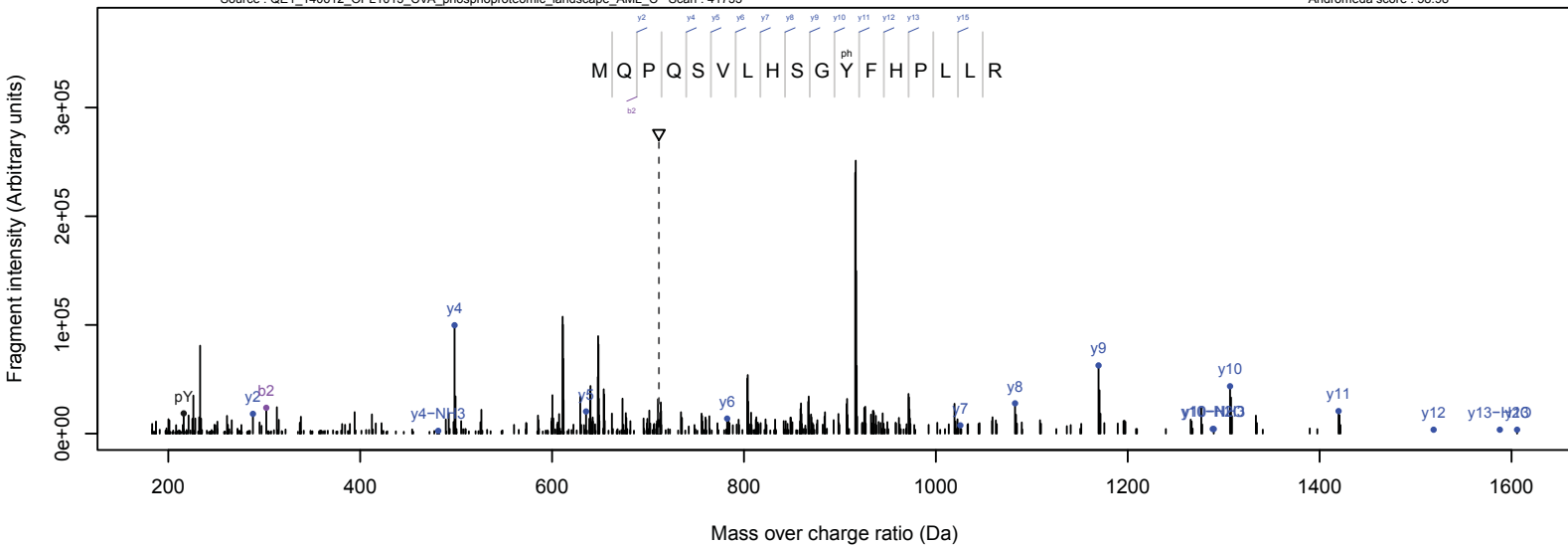
Andromeda score : 58.98



(ac)MQPQSVLHSGY(ph)FHPLLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 41733

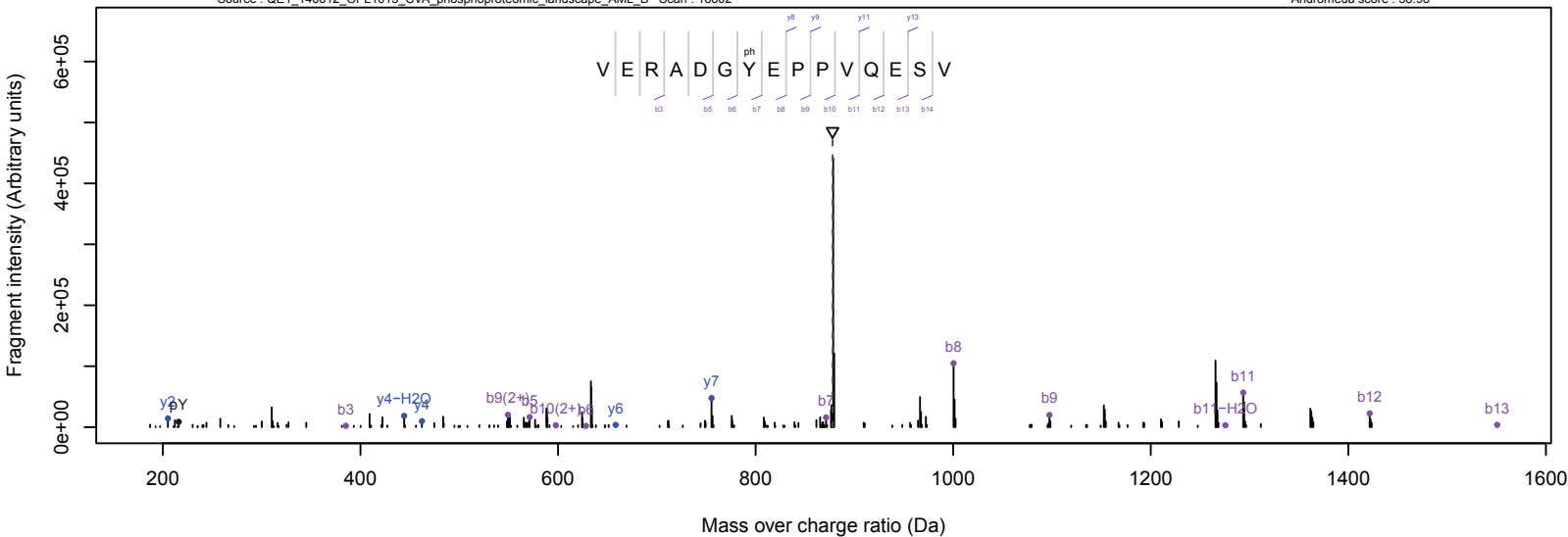
Andromeda score: 58.98



VERADGY(ph)EPPVQESV-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 16802

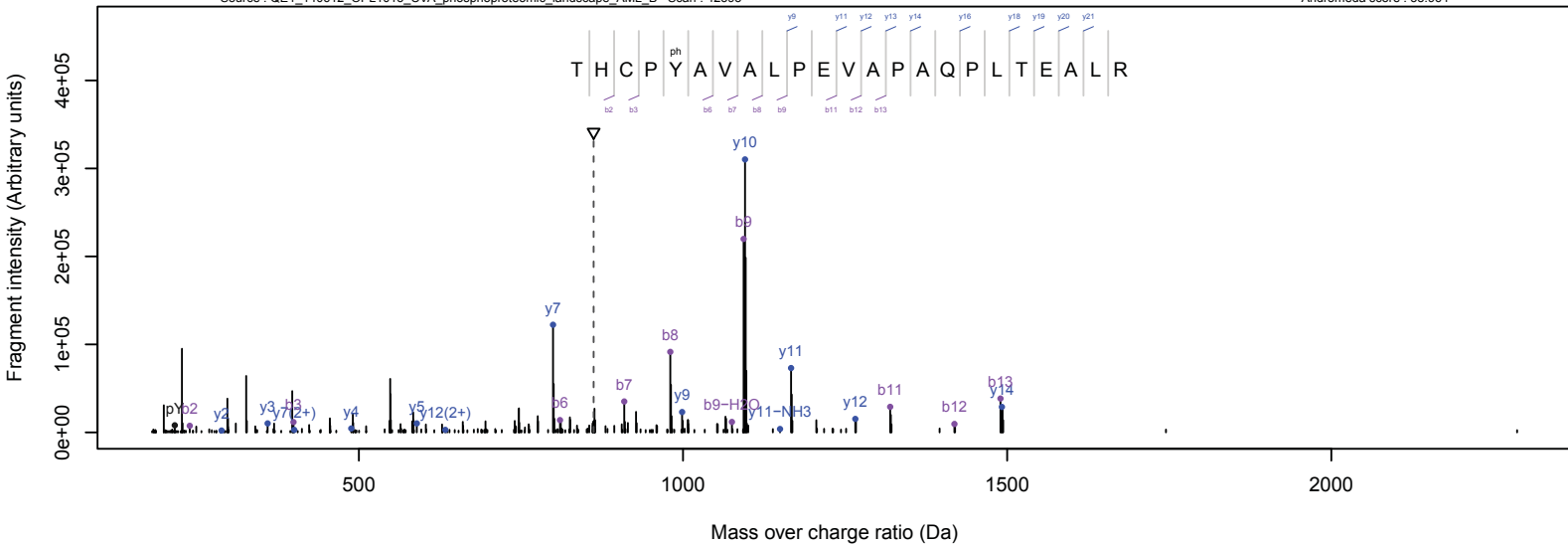
Andromeda score: 58.98



THCPY(ph)AVALPEVAPAQLTEALR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 42395

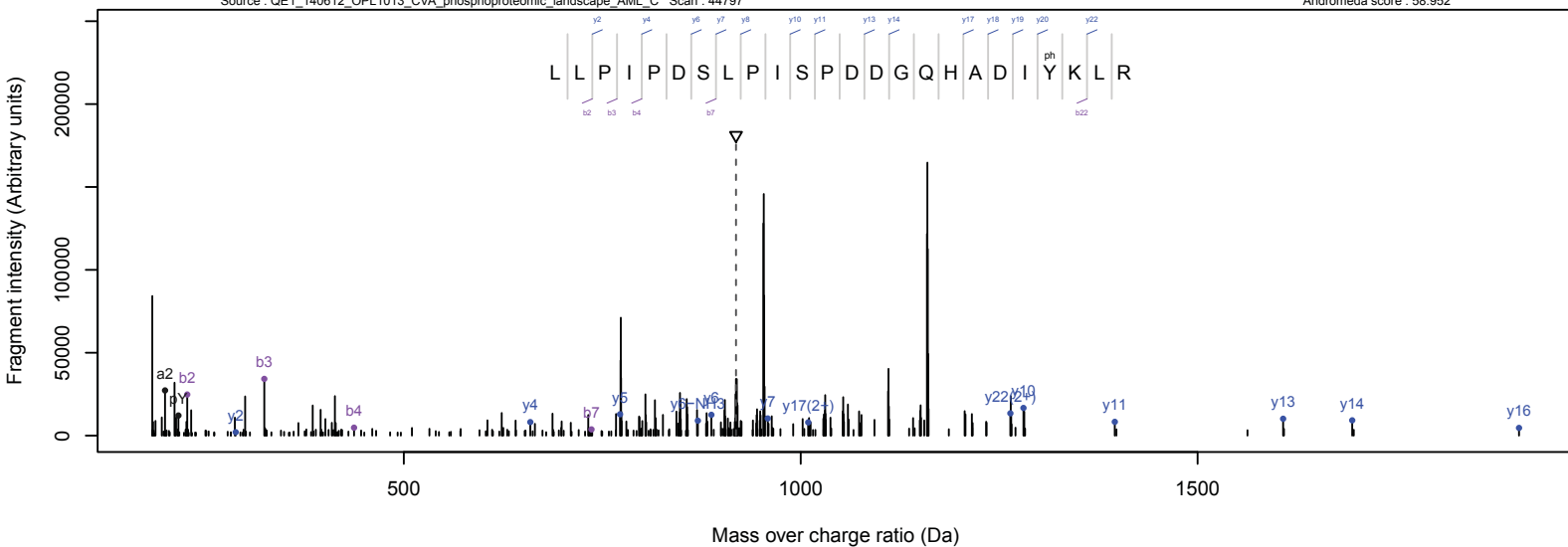
Andromeda score: 58.964



LLPIPDSLPISPDGQHADIYKLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44797

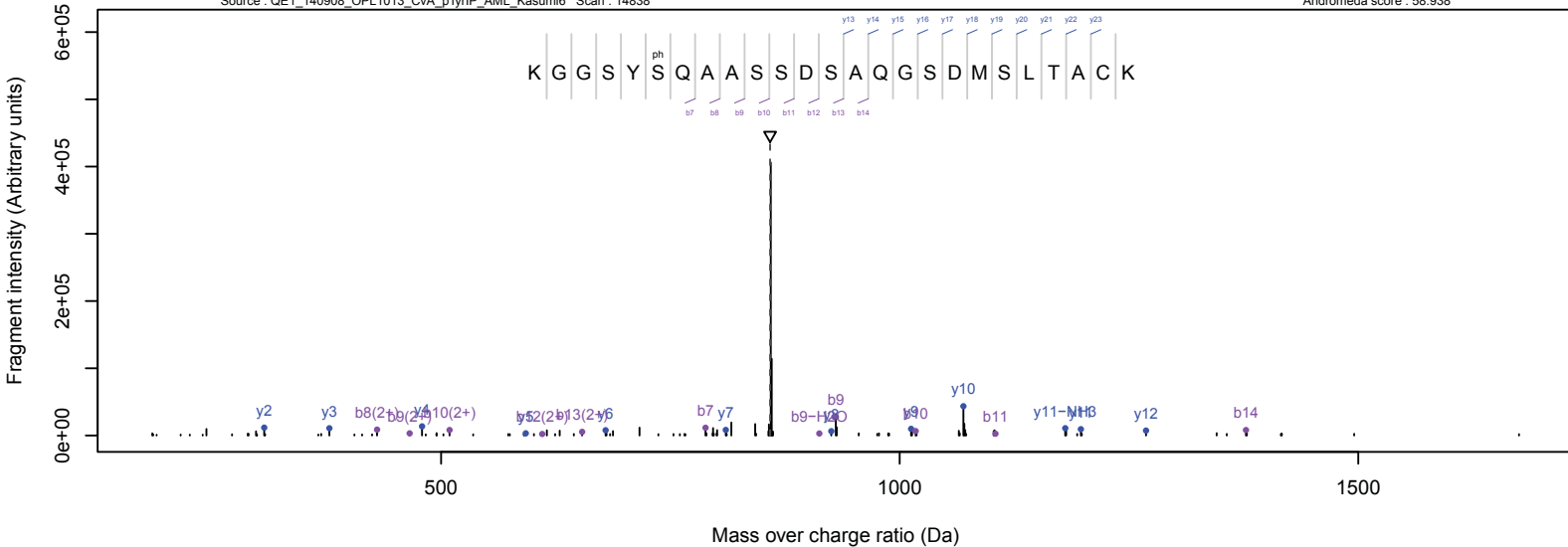
Andromeda score : 58.952



KGGSYS(ph)QAASSDQAQSDMSLTACK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 14838

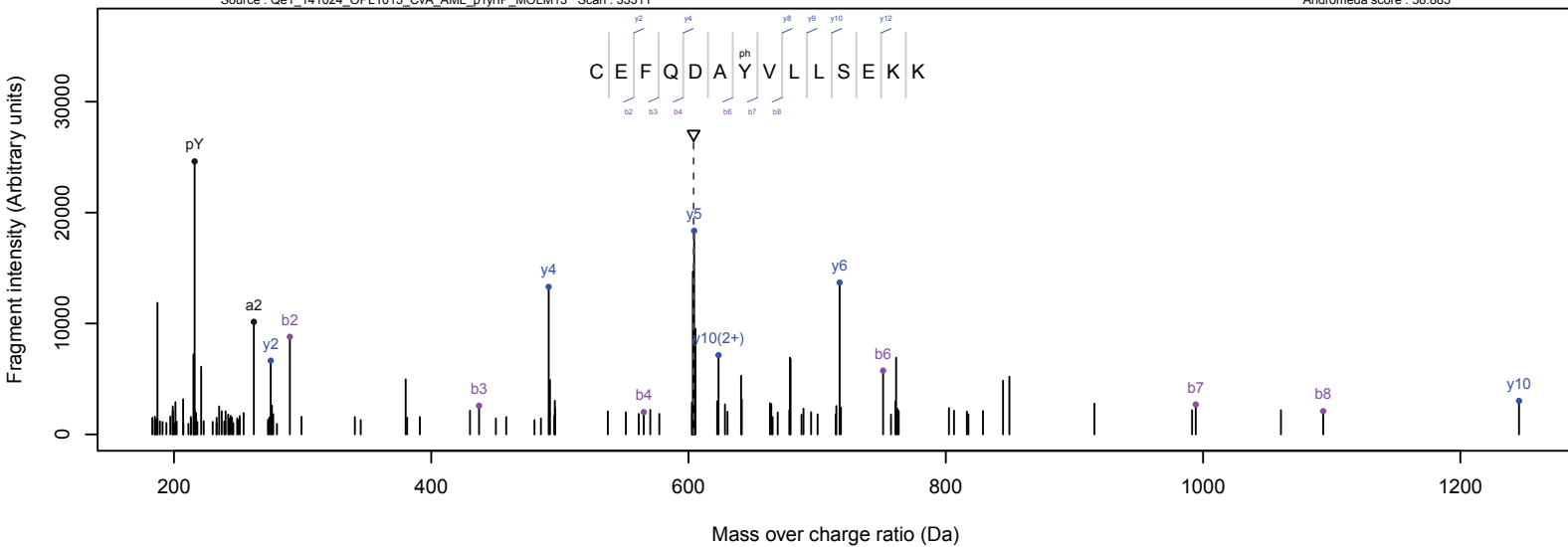
Andromeda score : 58.938



CEFQDAYVLLSEKK-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 33311

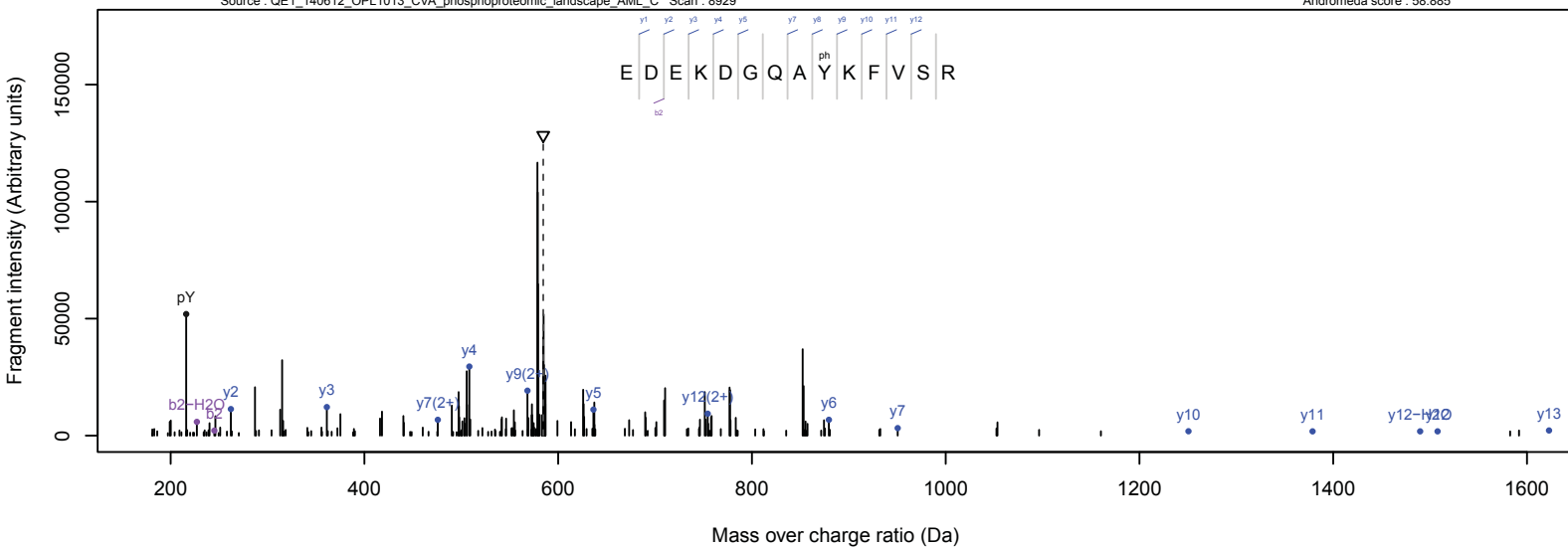
Andromeda score : 58.885



EDEKDGQAY(ph)KFVSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8929

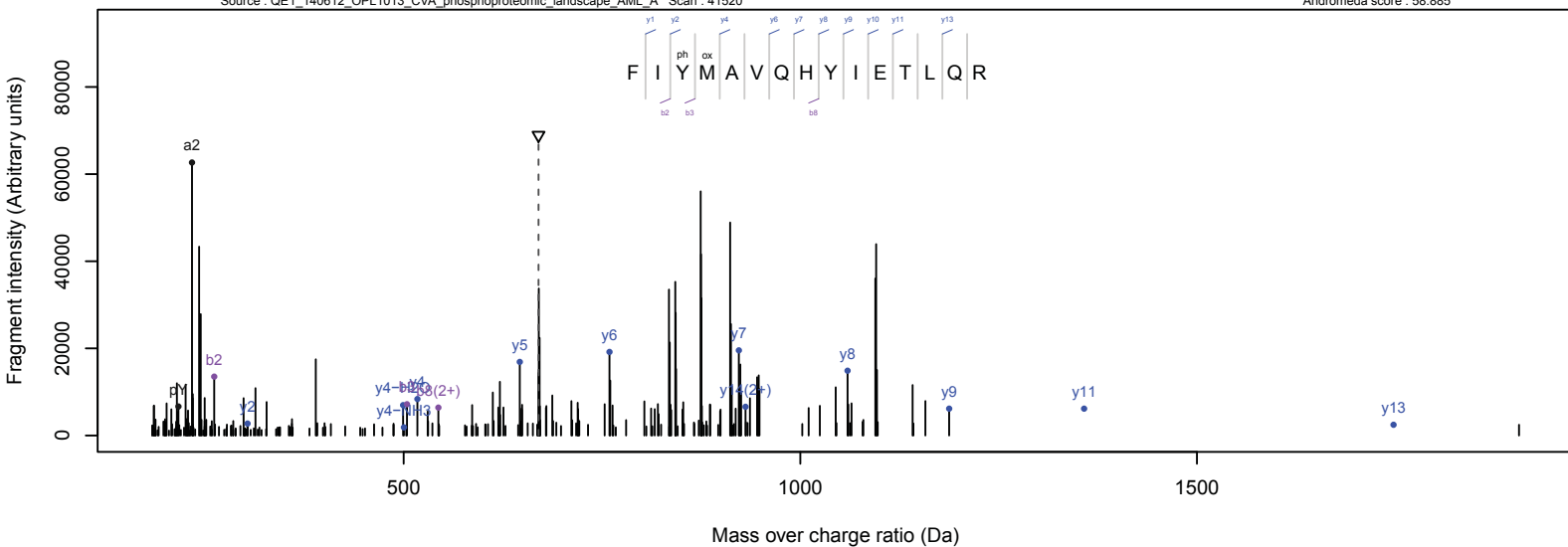
Andromeda score : 58.885



FIY(ph)M(ox)AVQHYIETLQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 41520

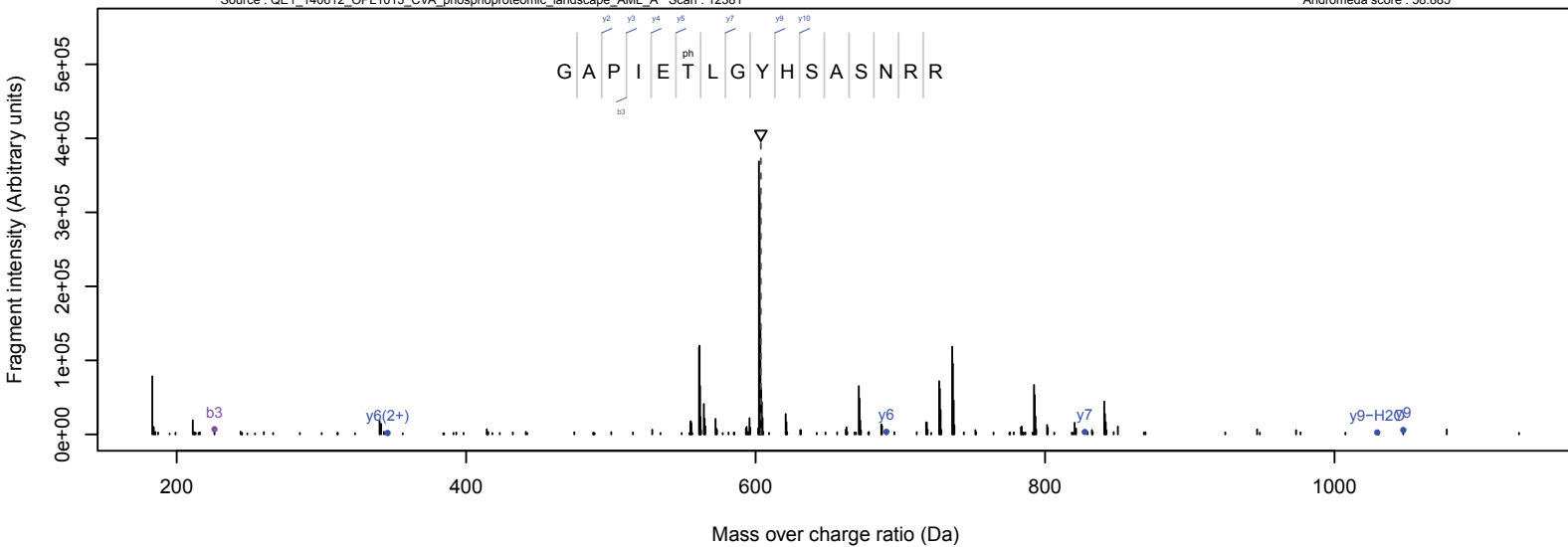
Andromeda score : 58.885



GAPIET(ph)LGYHSASNRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12381

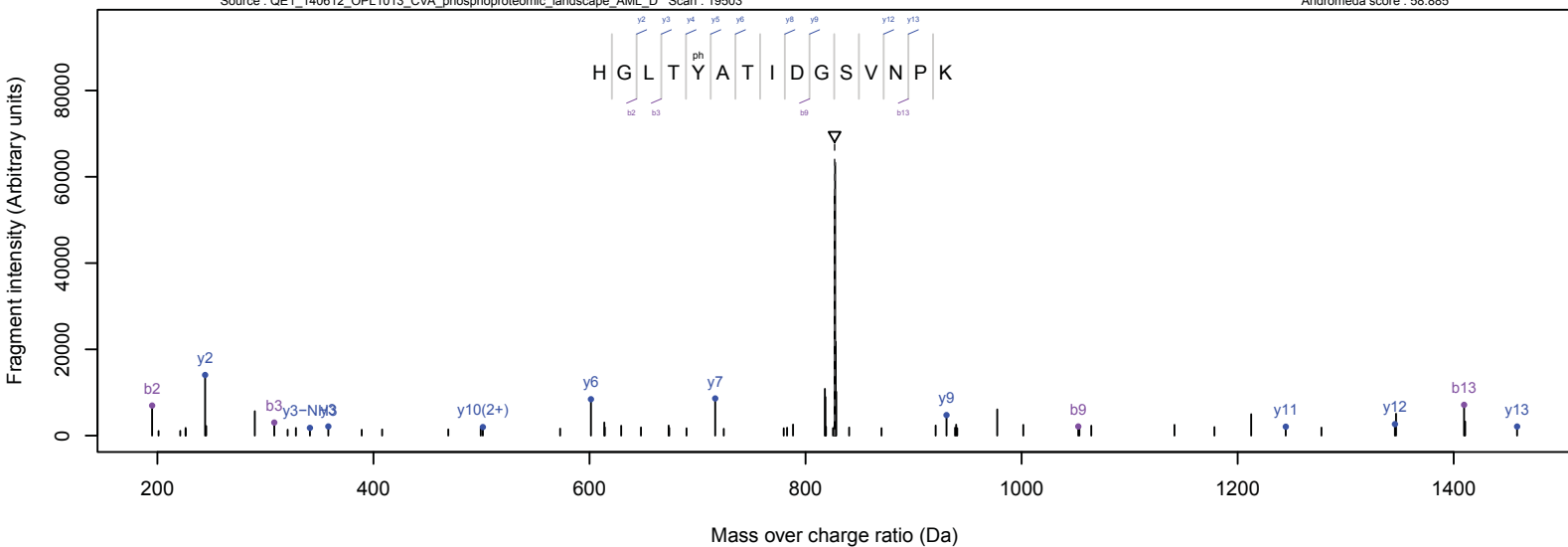
Andromeda score : 58.885



HGLTY(ph)ATIDGSVNPk-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 19503

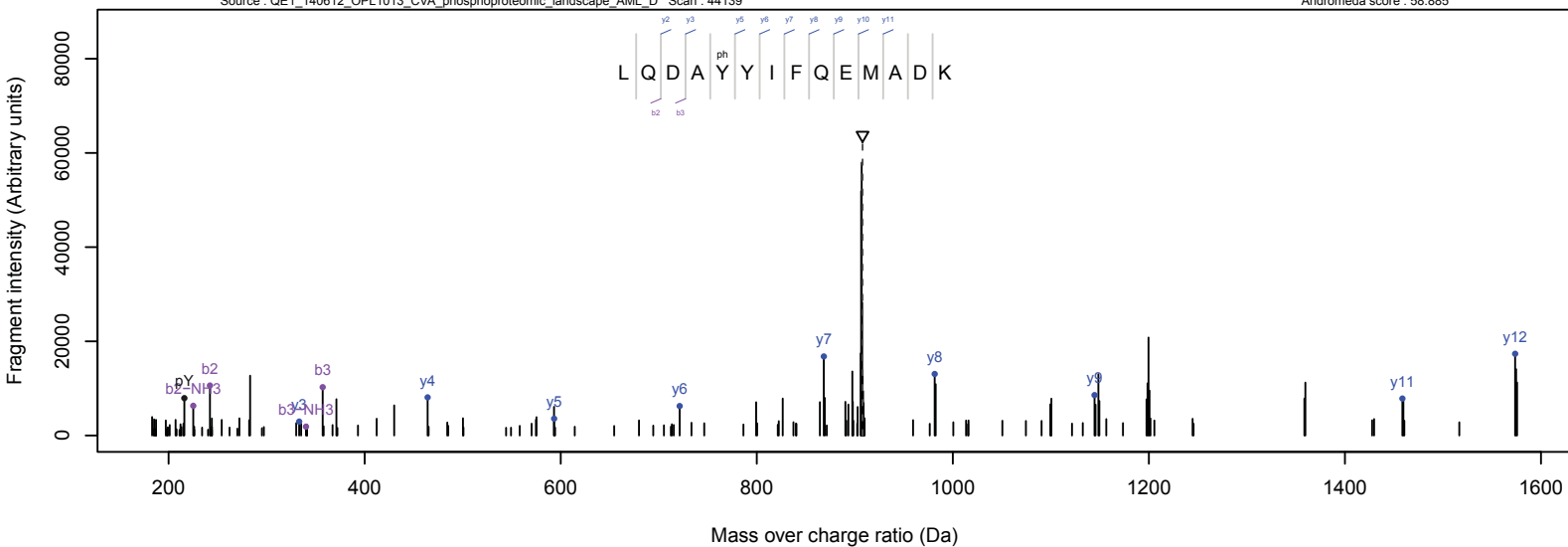
Andromeda score: 58.885



LQDAY(ph)YIFQEMADK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 44139

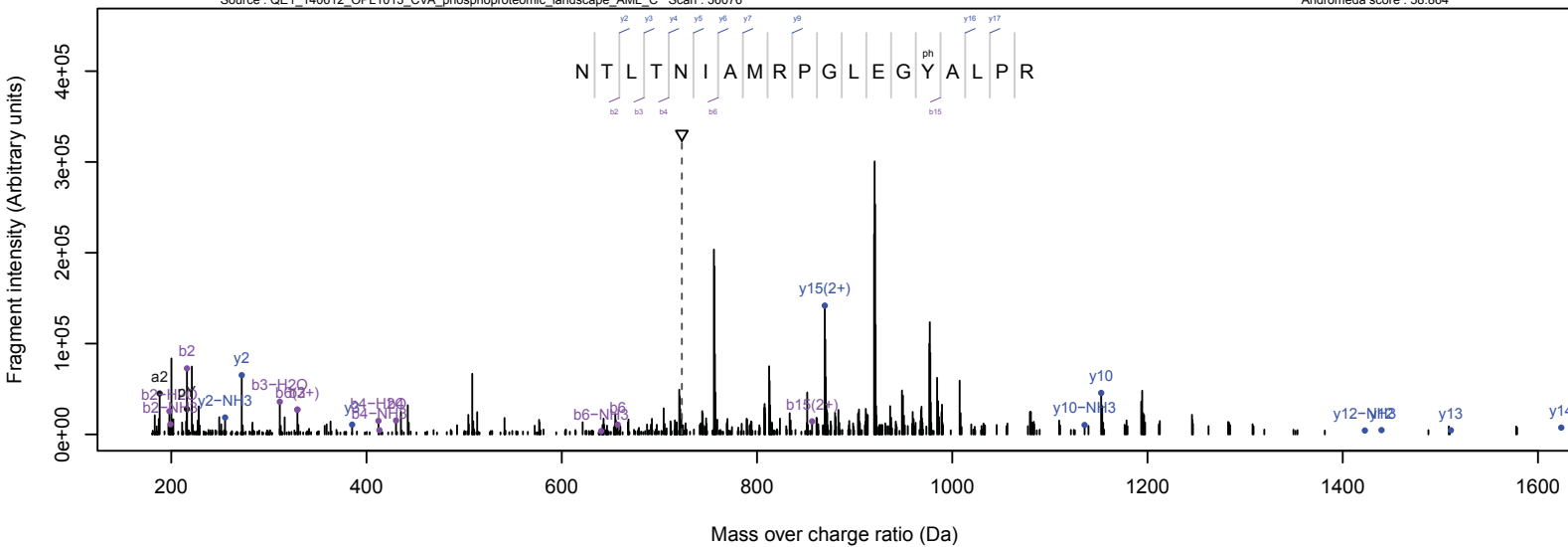
Andromeda score: 58.885



NLTNIAMRPGLEG(ph)YALPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36076

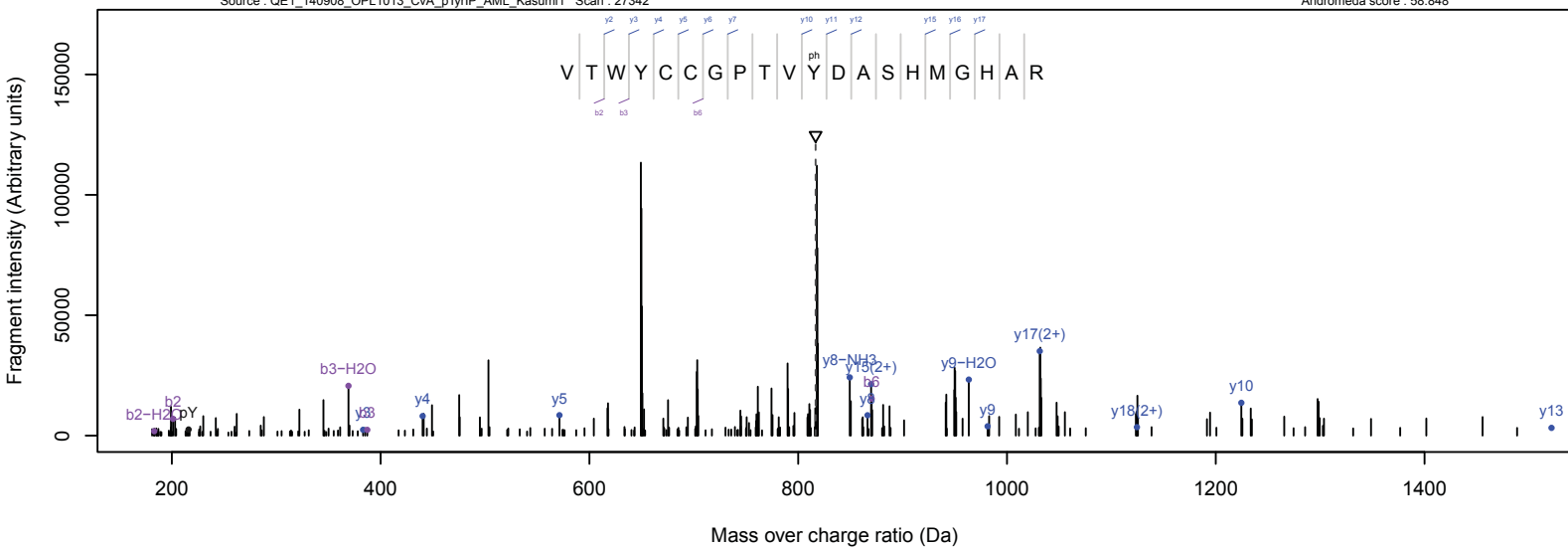
Andromeda score: 58.864



## VTWYCCGPTVY(ph)DASHMGHAR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27342

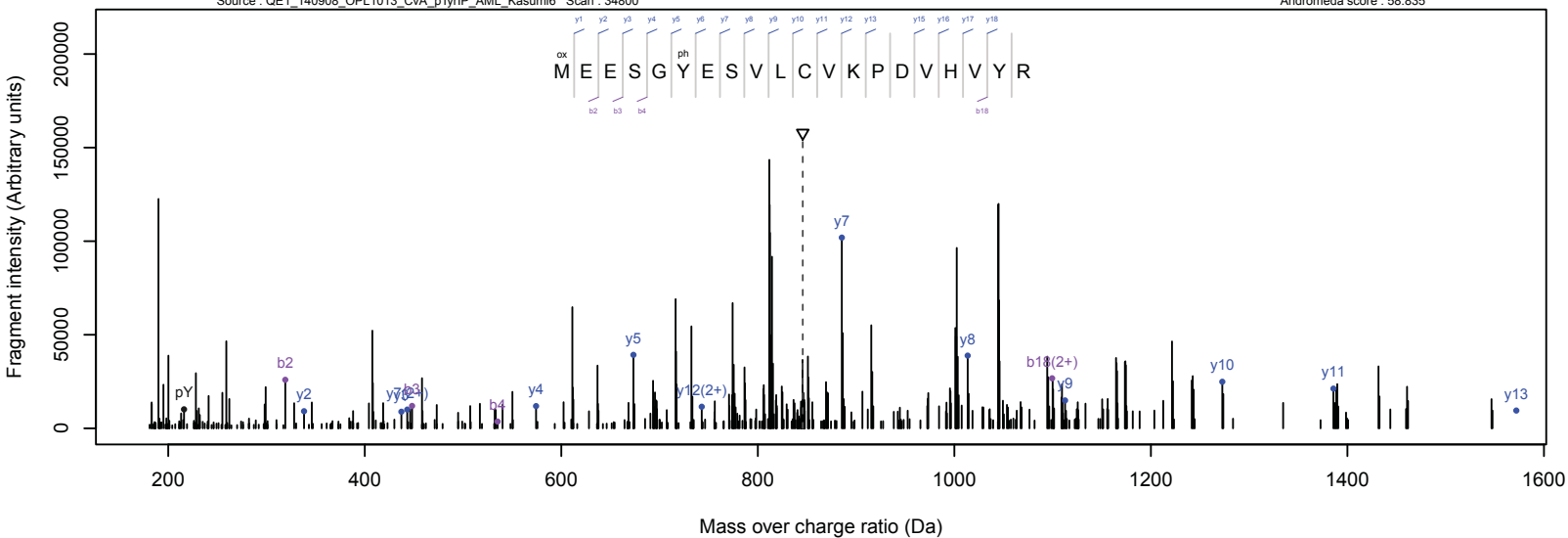
Andromeda score : 58.848



## (ac)M(ox)EESGY(ph)ESVLCVKPDVHVYR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 34800

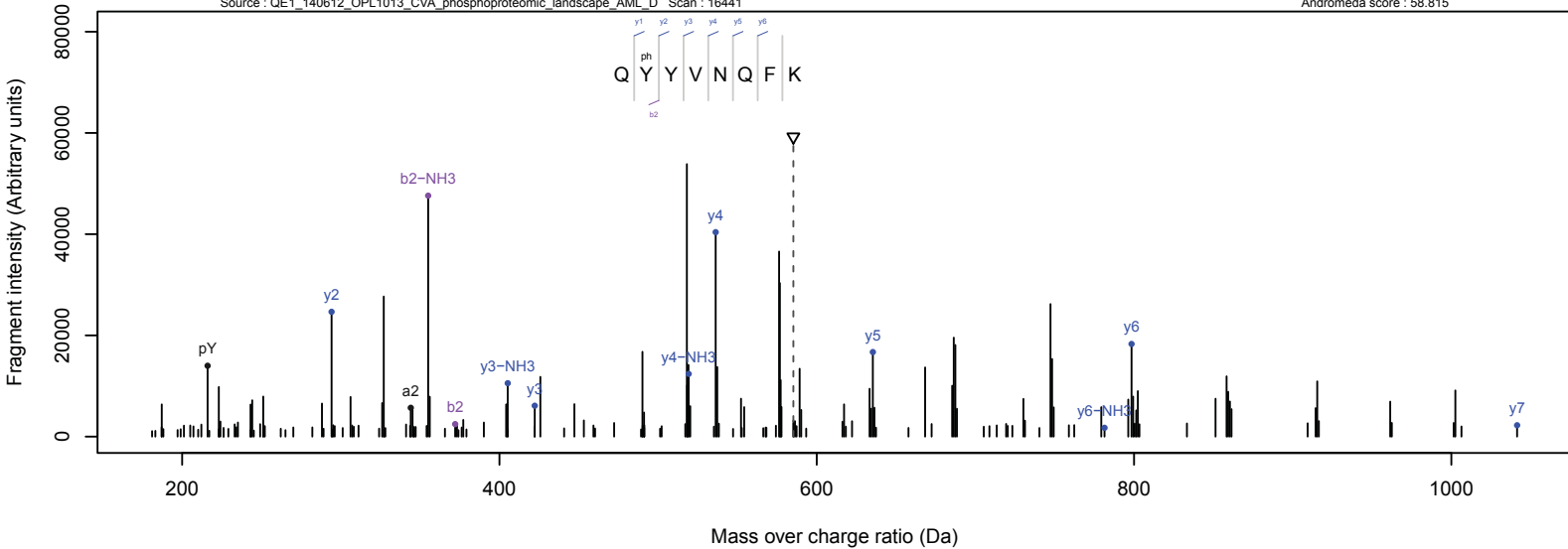
Andromeda score : 58.835



## QY(ph)YVNQFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 16441

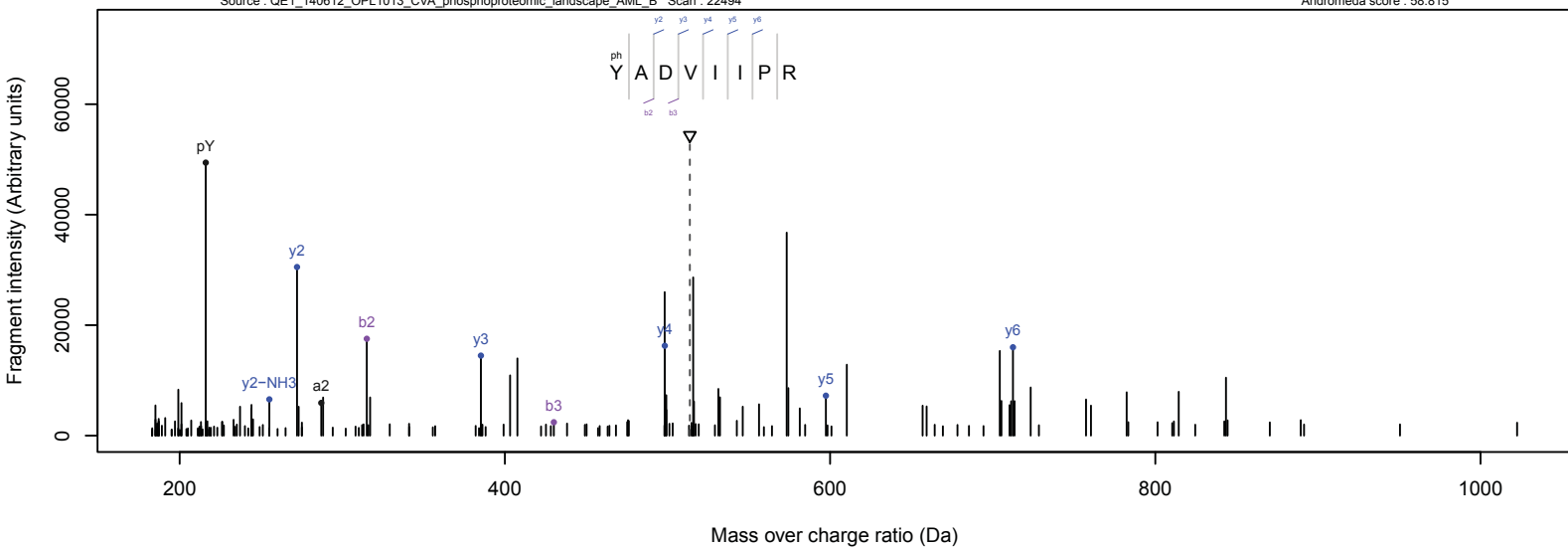
Andromeda score : 58.815



## Y(ph)ADVIIPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22494

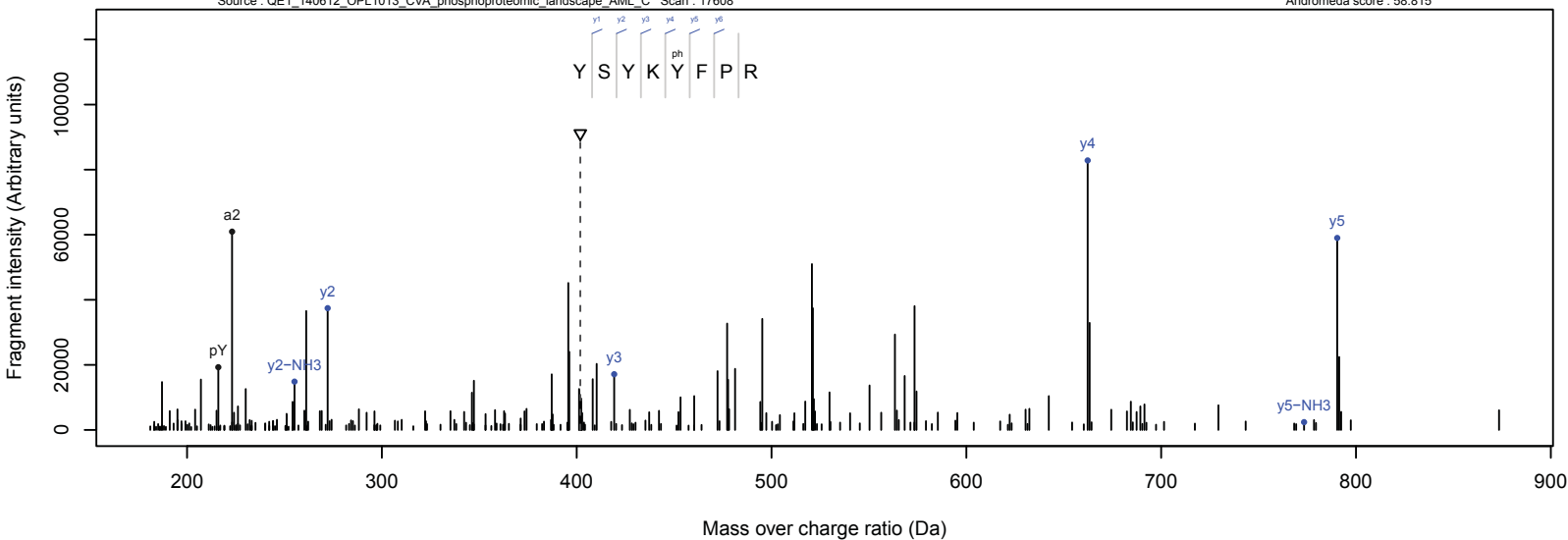
Andromeda score : 58.815



## YSYKY(ph)FPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17608

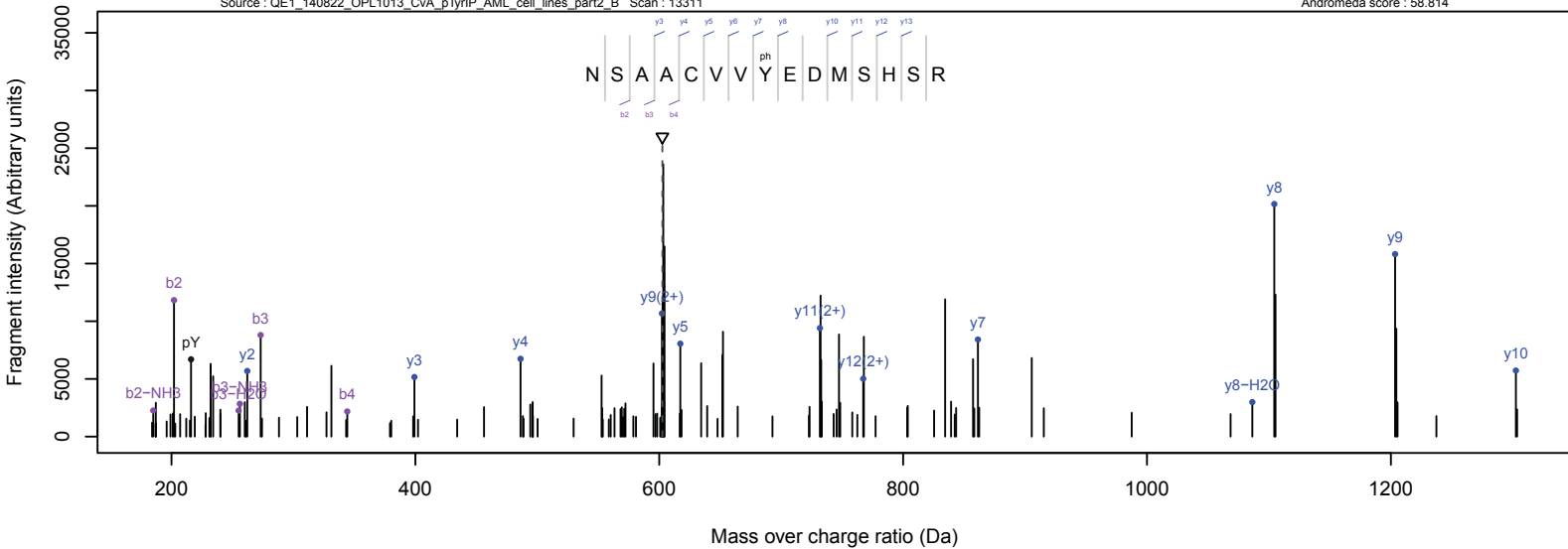
Andromeda score : 58.815



## NSAACVVY(ph)EDMSHSR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 13311

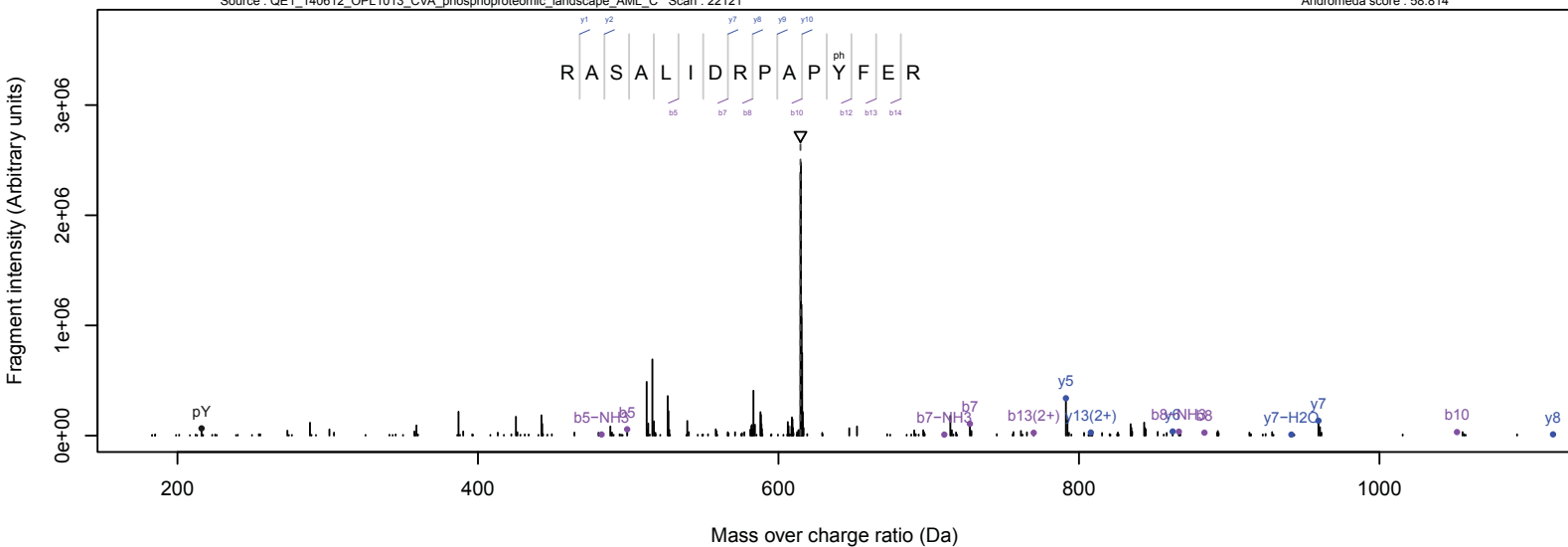
Andromeda score : 58.814



## RASALIDRPAPY(ph)FER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22121

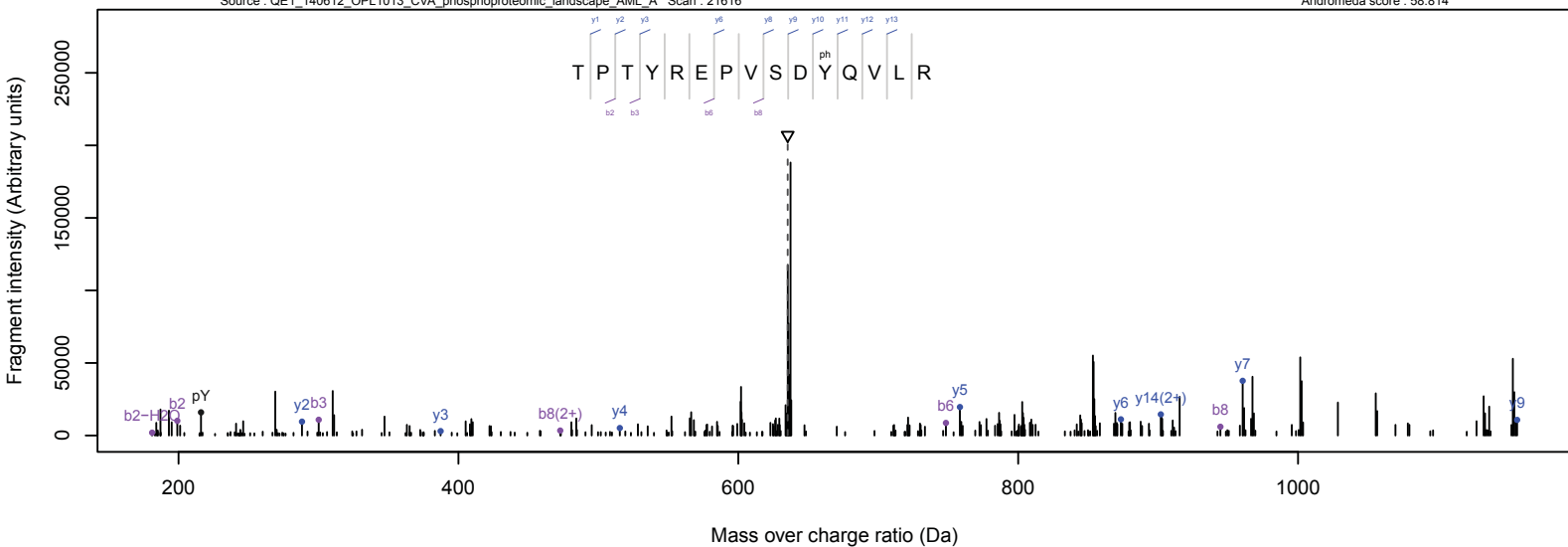
Andromeda score : 58.814



## TPTYREPVS DY(ph)QVLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 21616

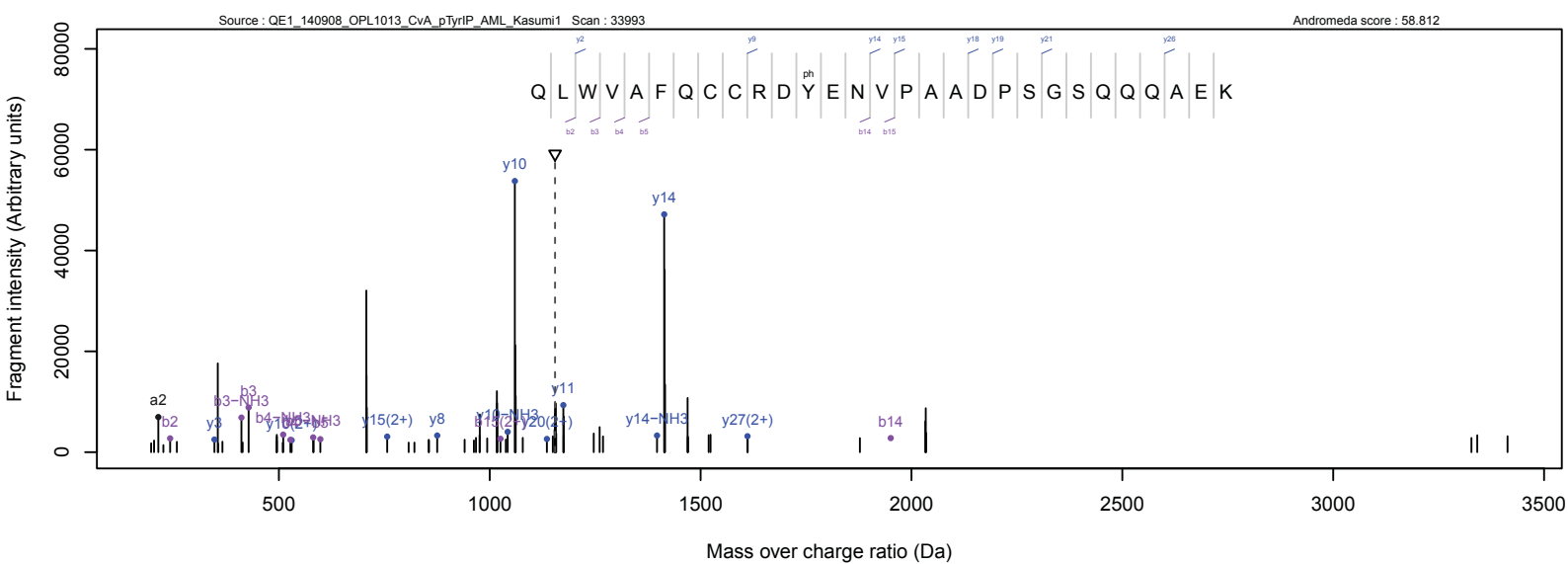
Andromeda score : 58.814



## QLWVAFQCCR DY(ph)ENVPAAADPSGSQQQAEK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 33993

Andromeda score : 58.812

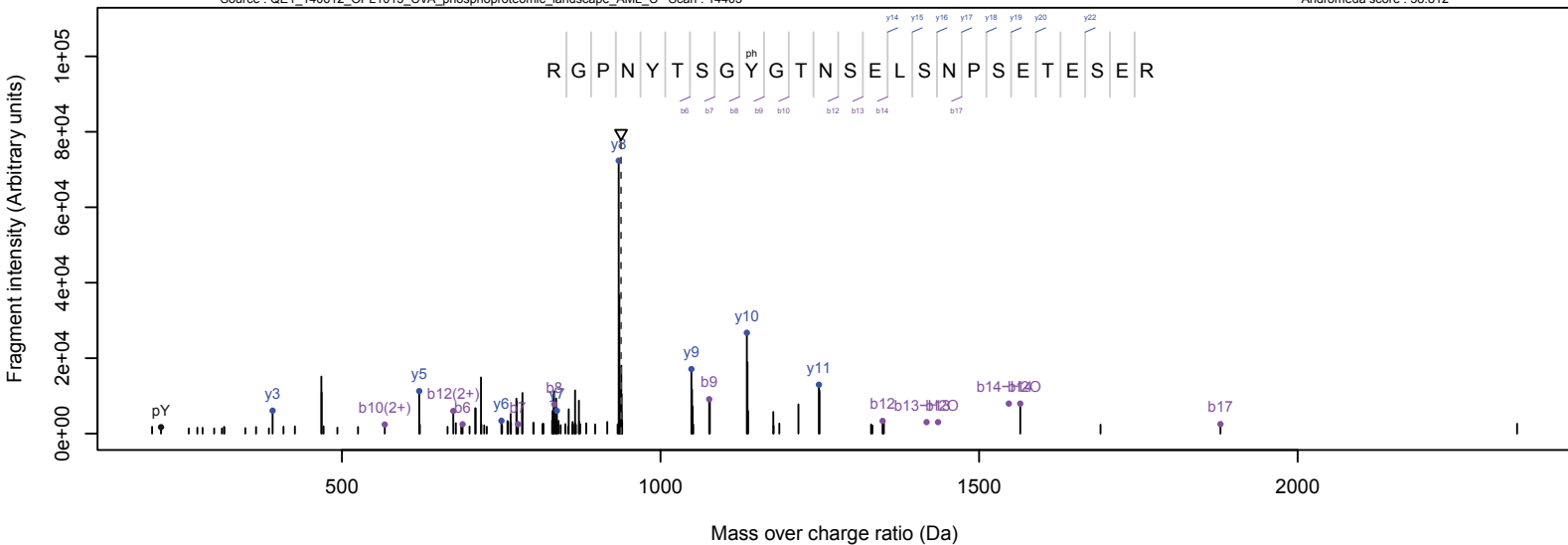




**RGPNYTSGY(ph)GTNSELSNPSETESER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14403

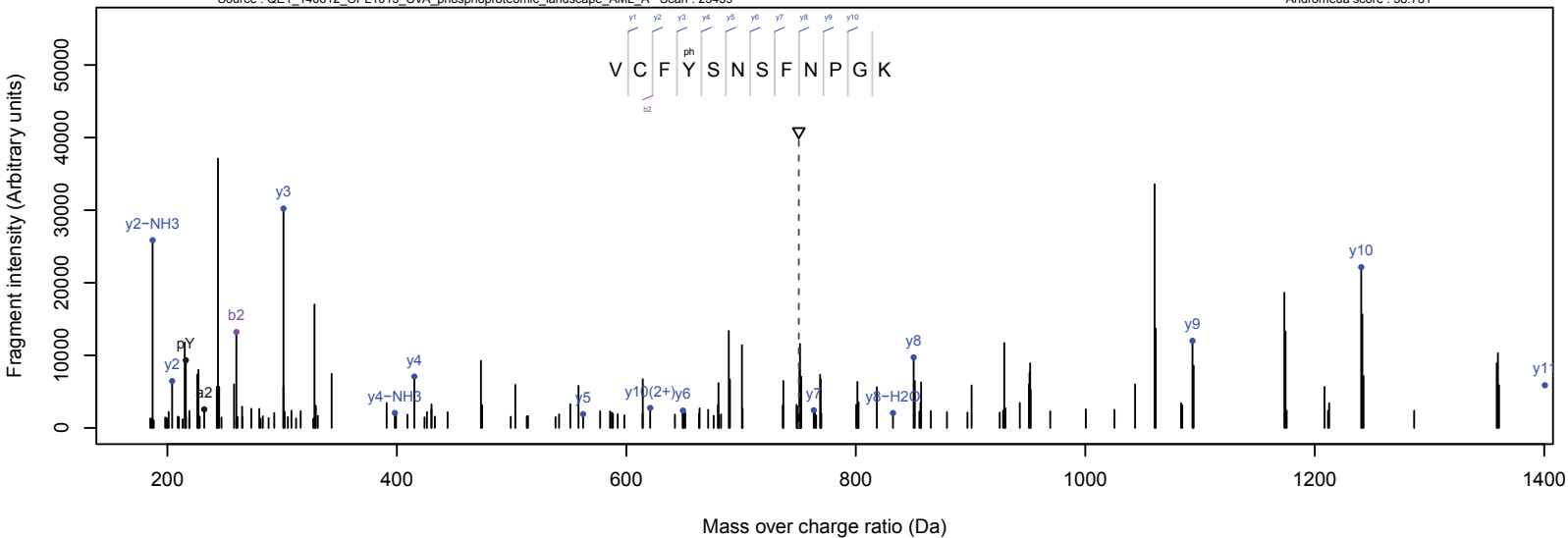
Andromeda score : 58.812



**VCFY(ph)SNSFNGPK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 23439

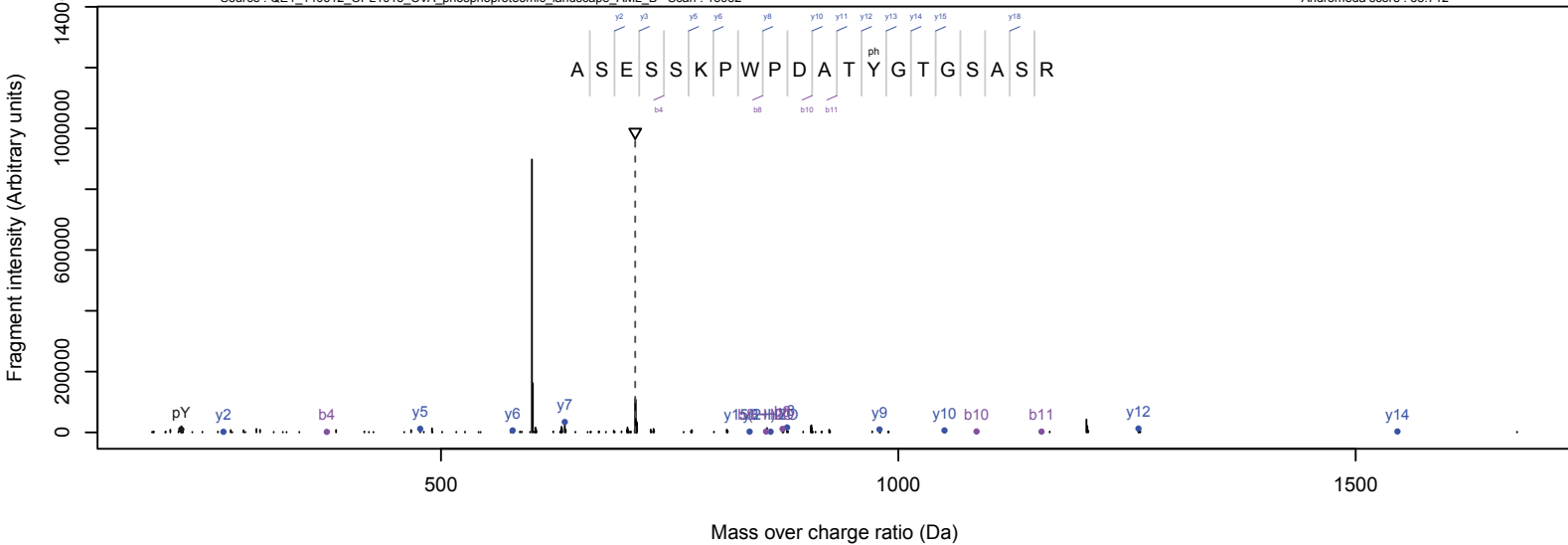
Andromeda score : 58.781



**ASESSKPWPDATY(ph)GTGSASR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13982

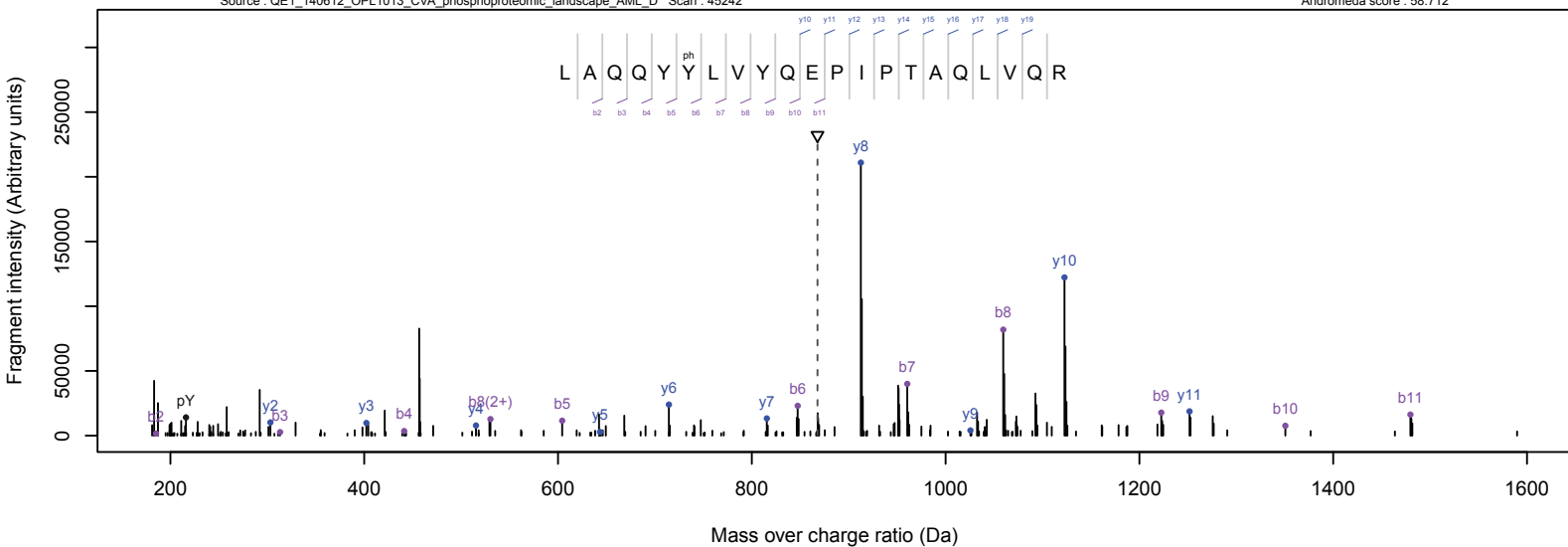
Andromeda score : 58.712



LAQQYY(ph)LVYQEPIPTAQLVQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 45242

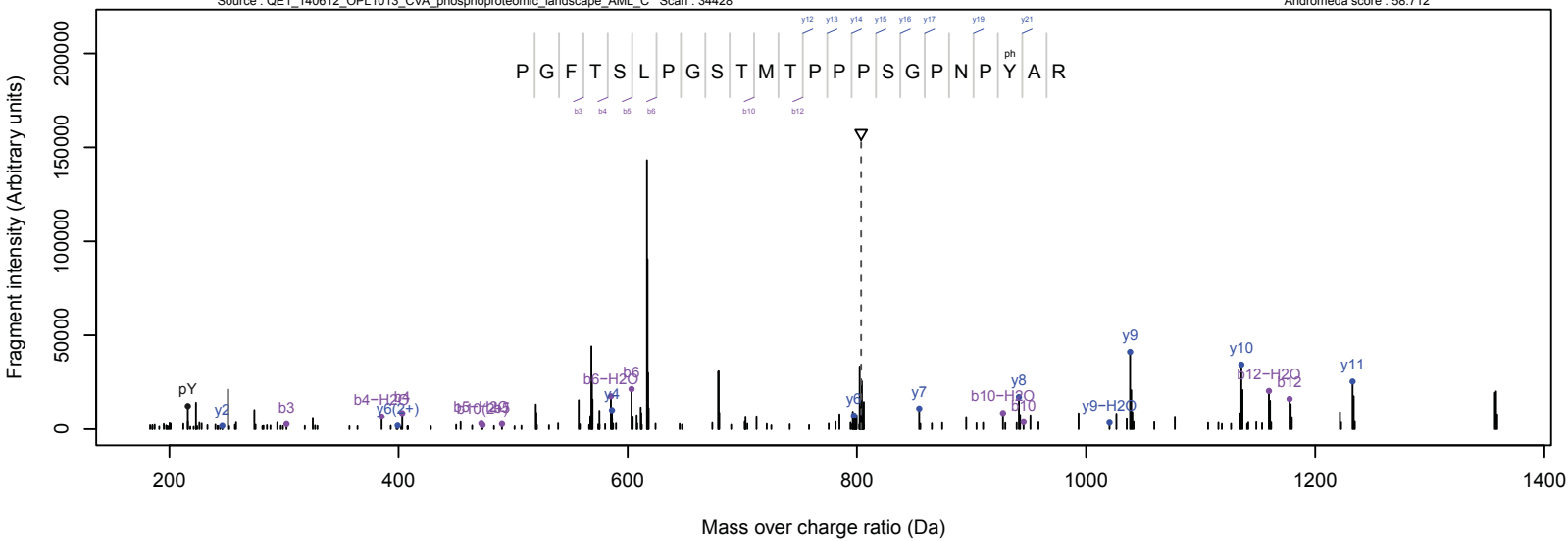
Andromeda score : 58.712



PGFTSLPGSTMTPPSPGNPY(ph)AR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34428

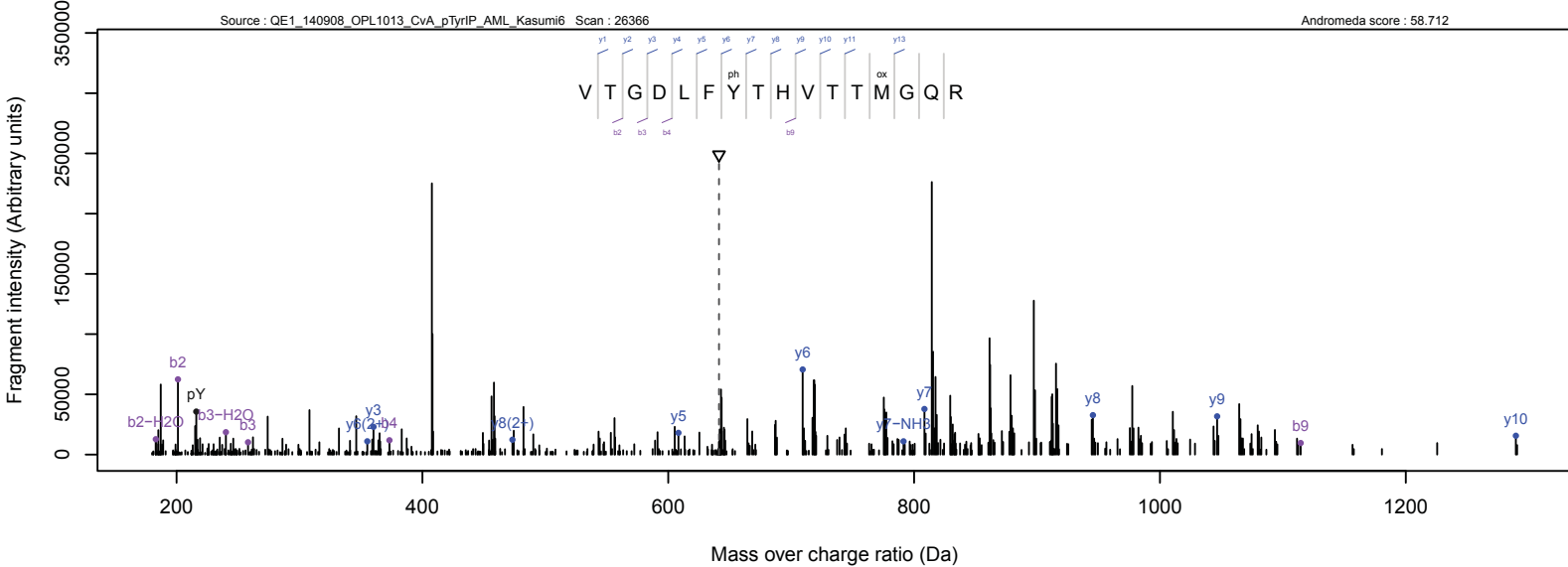
Andromeda score : 58.712



VTGDLFY(ph)THVTM(ox)GQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 26366

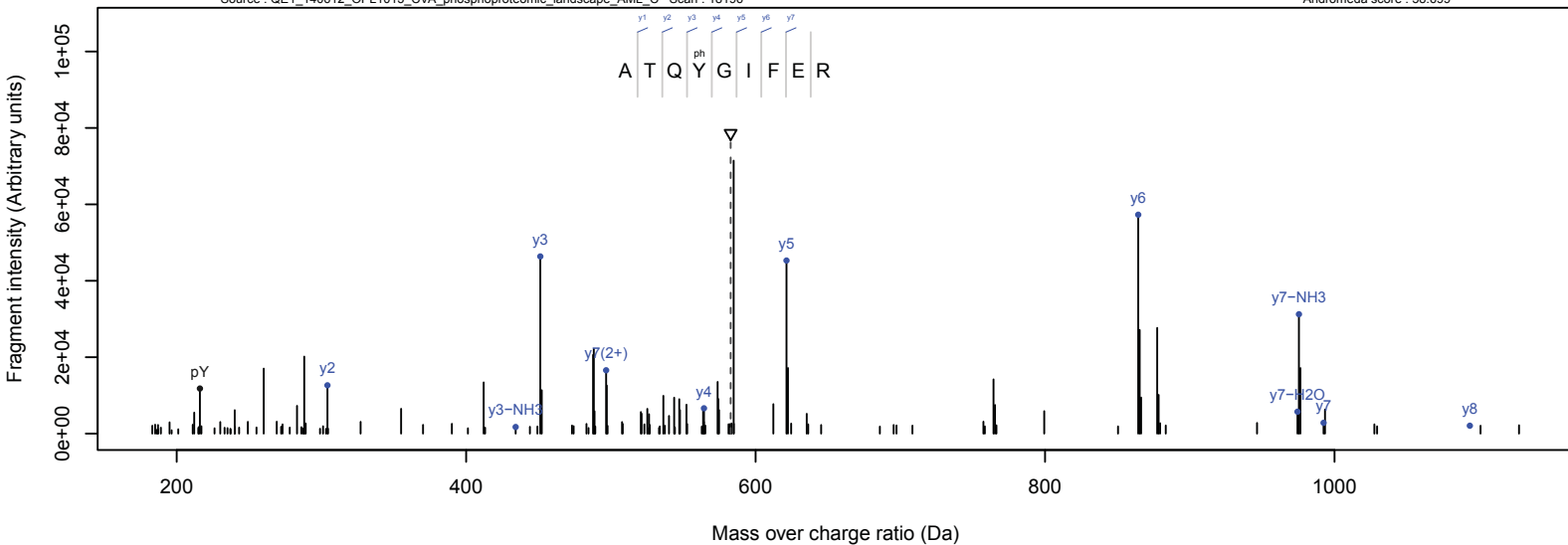
Andromeda score : 58.712



## ATQY(ph)GIFER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18196

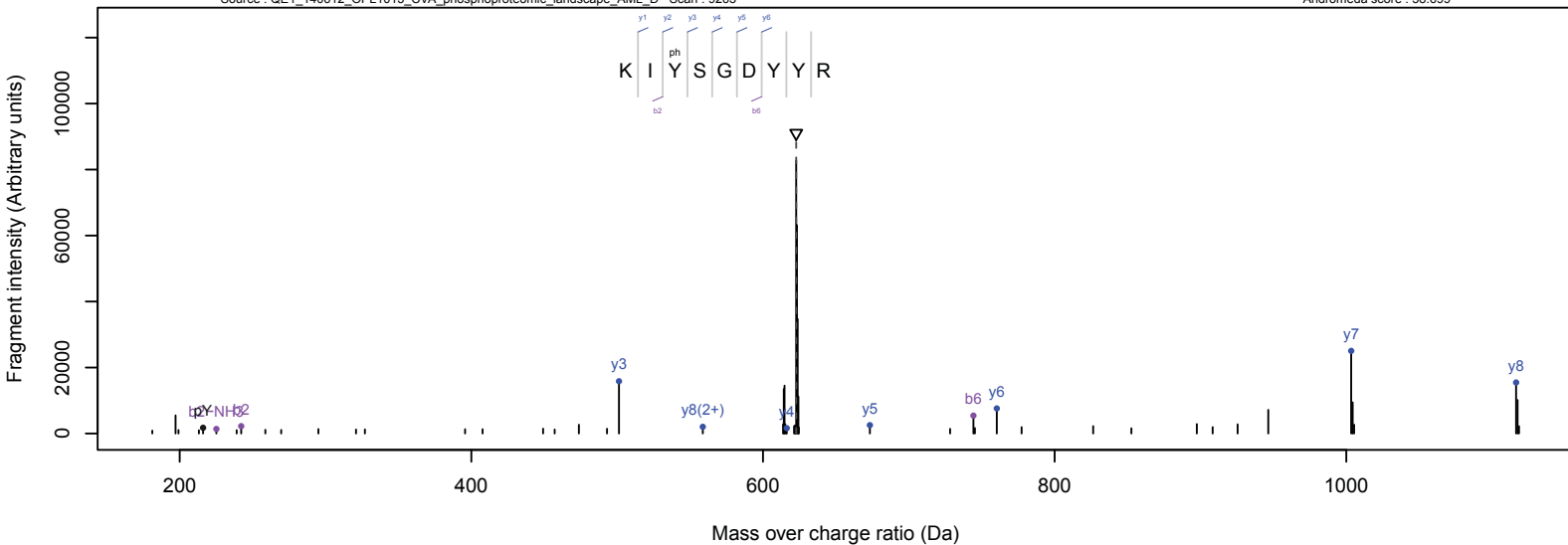
Andromeda score : 58.699



## KIY(ph)SGDYR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9263

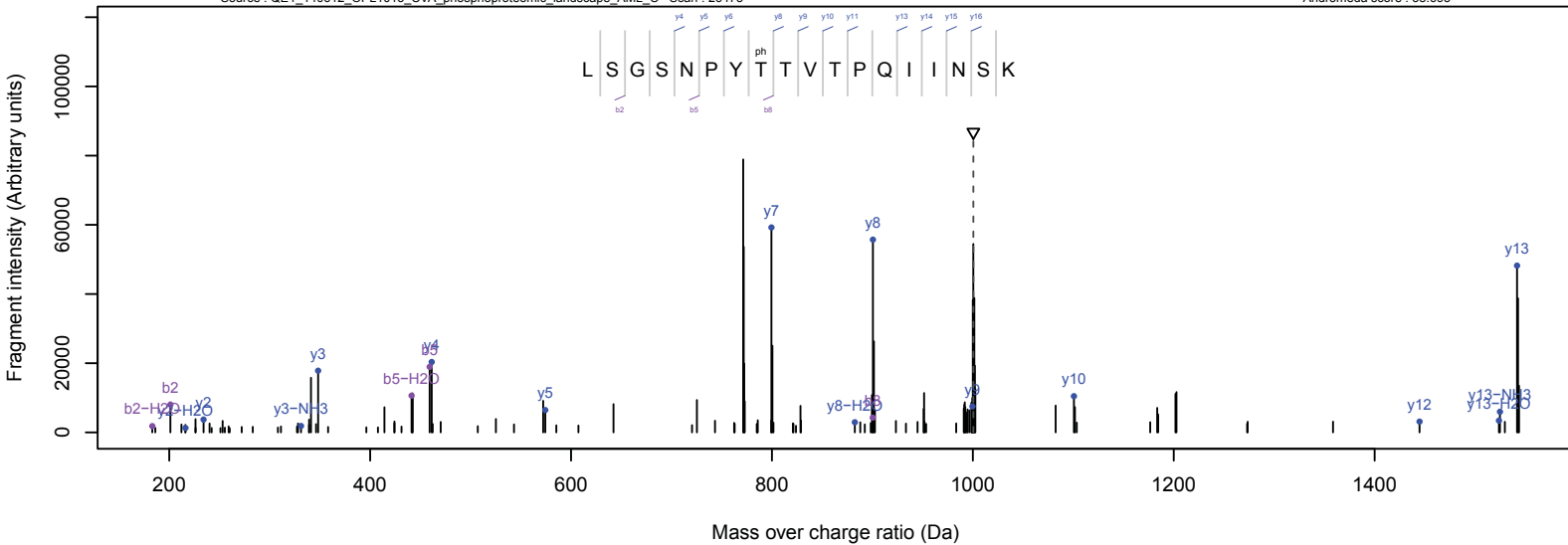
Andromeda score : 58.699



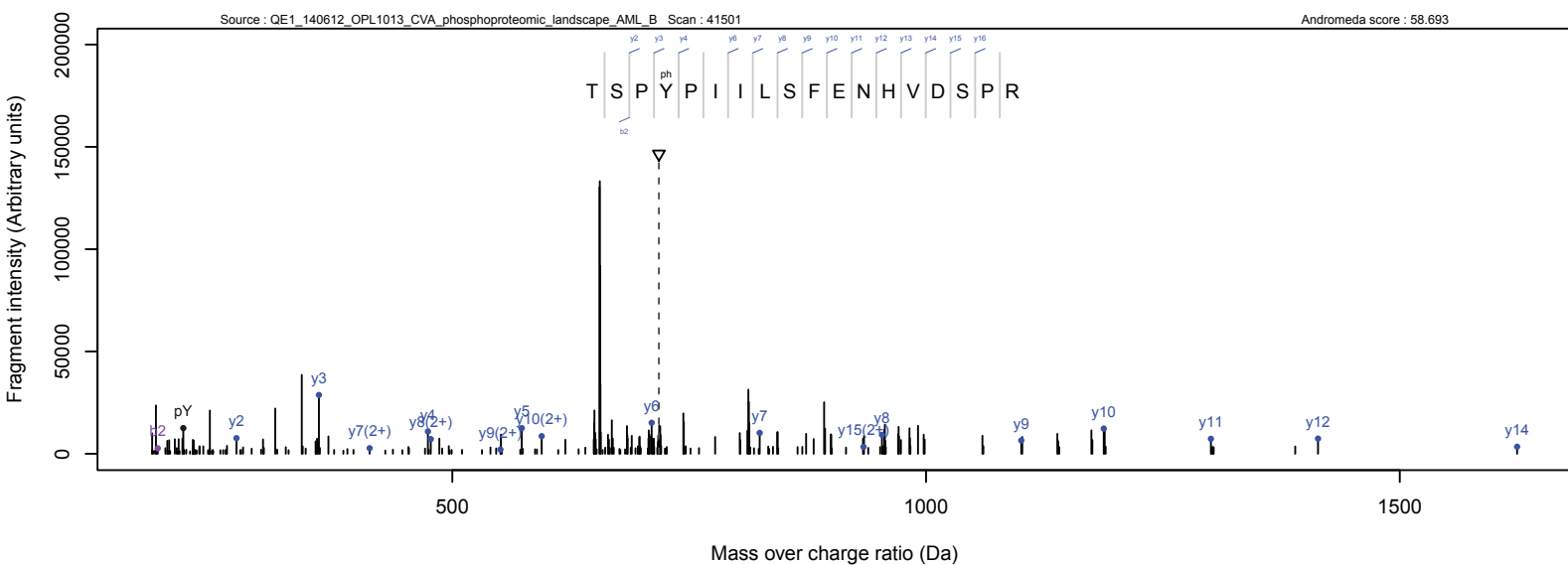
## LSGSNPY(ph)TVTPQIINSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26473

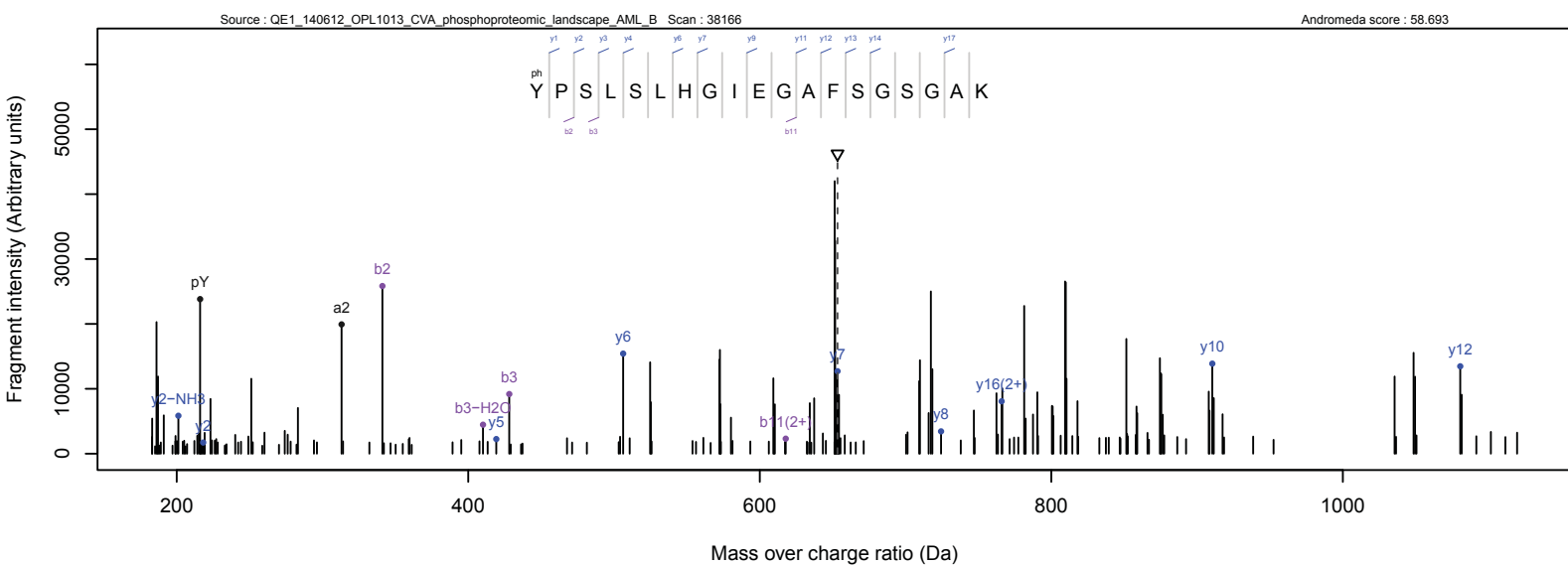
Andromeda score : 58.693



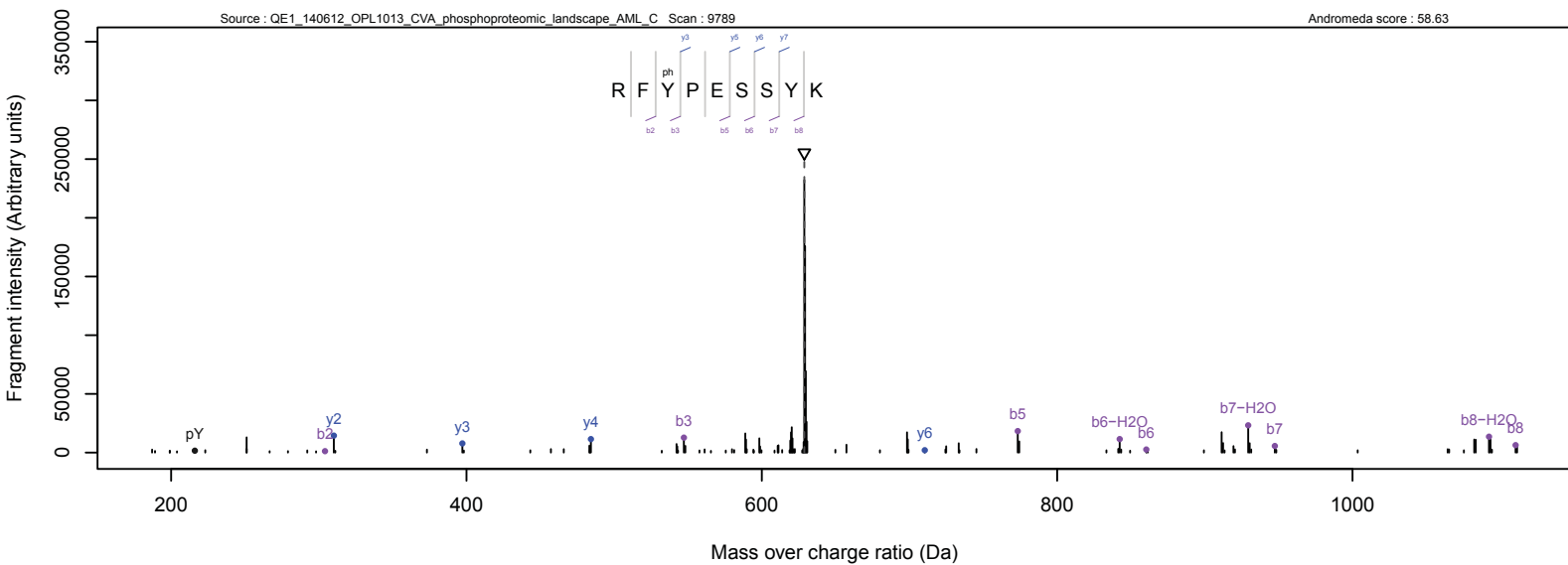
## TSPY(ph)PIILSFENHVDSPR-3+



## Y(ph)PSLSLHGIEGAFSGSGAK-3+



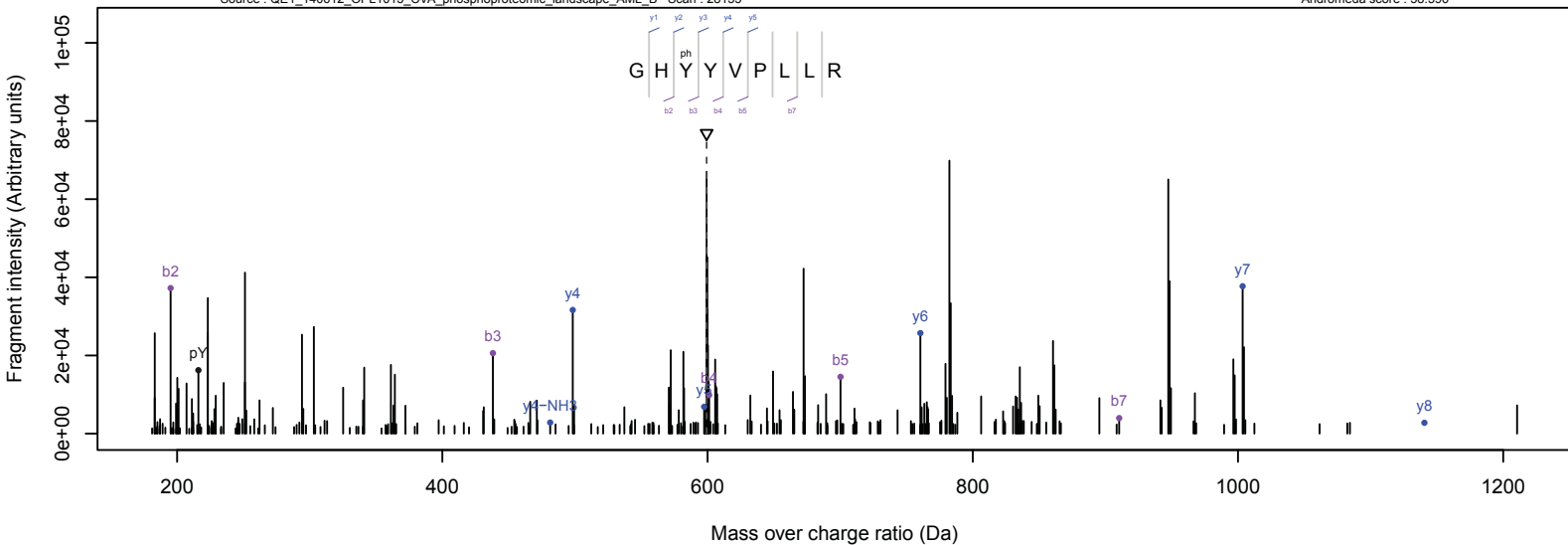
## RFY(ph)PESSYK-2+



## GHY(ph)YVPLLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28135

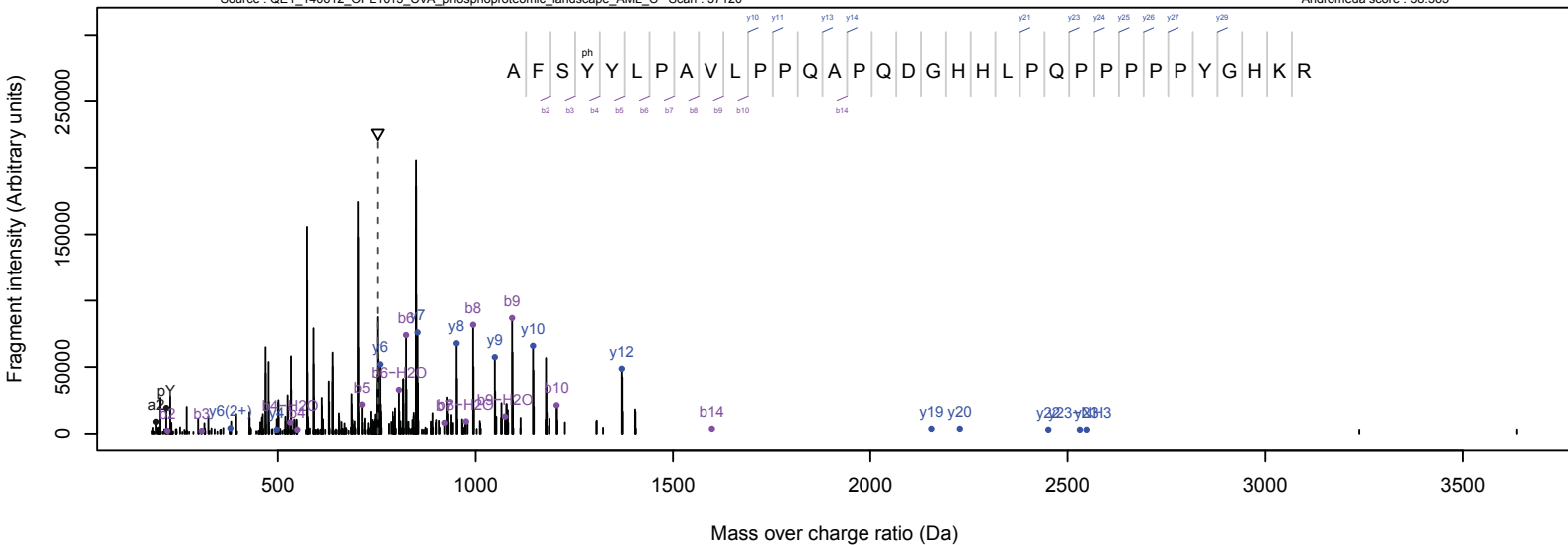
Andromeda score : 58.596



## AFSY(ph)YLPVLPQAPQDGHHLQPPPPYPYGHKR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37120

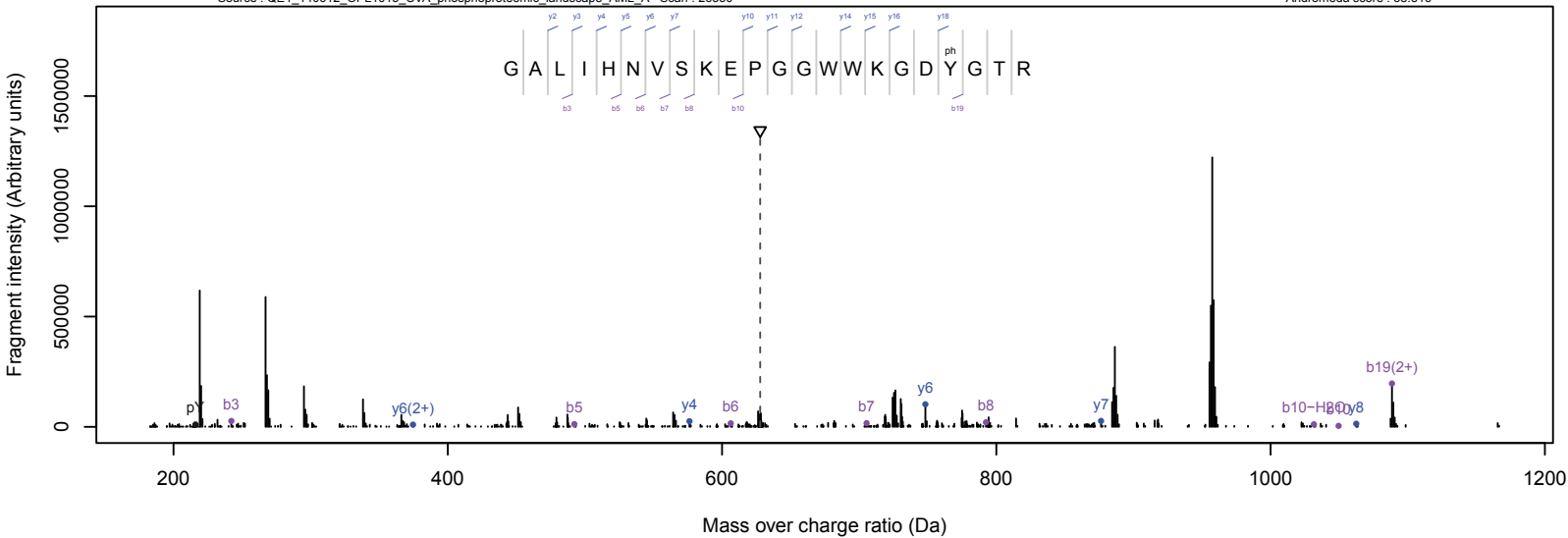
Andromeda score : 58.565



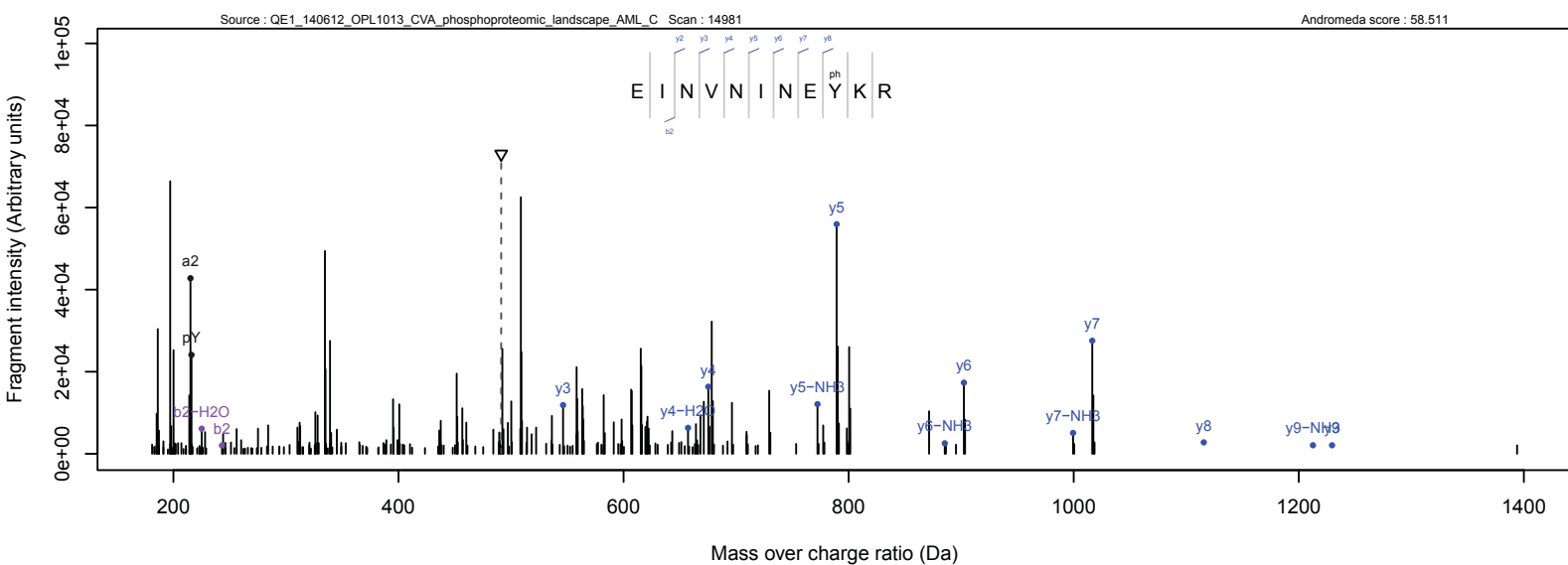
## GALIHNSKEPGGWWKGDY(ph)GTR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26330

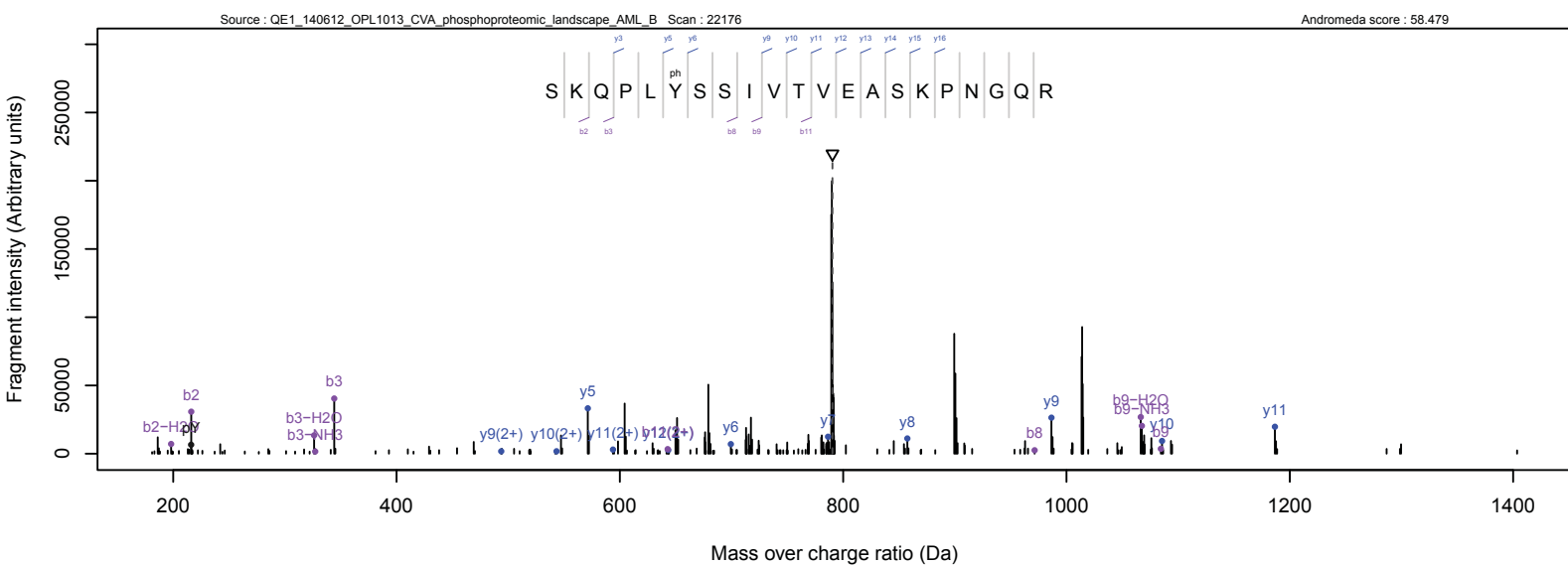
Andromeda score : 58.516



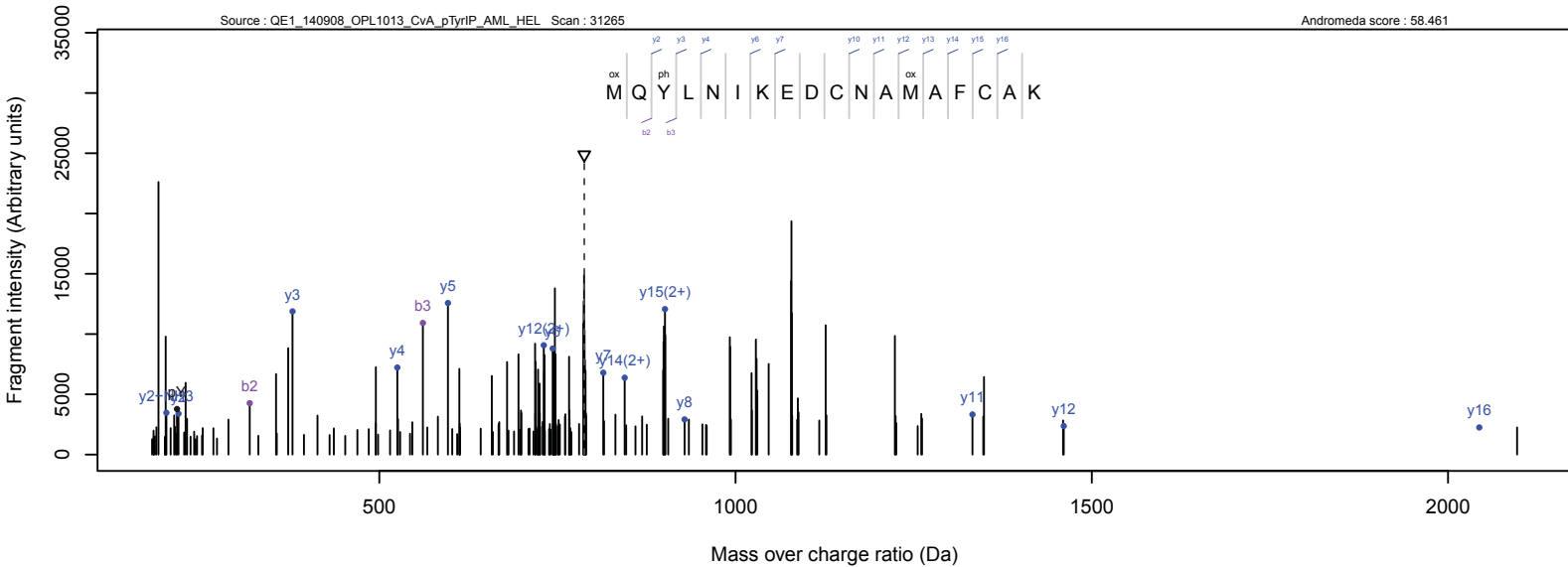
## EINVNINEY(ph)KR-3+



## SKQPLY(ph)SSIVTVEASKPNGQR-3+



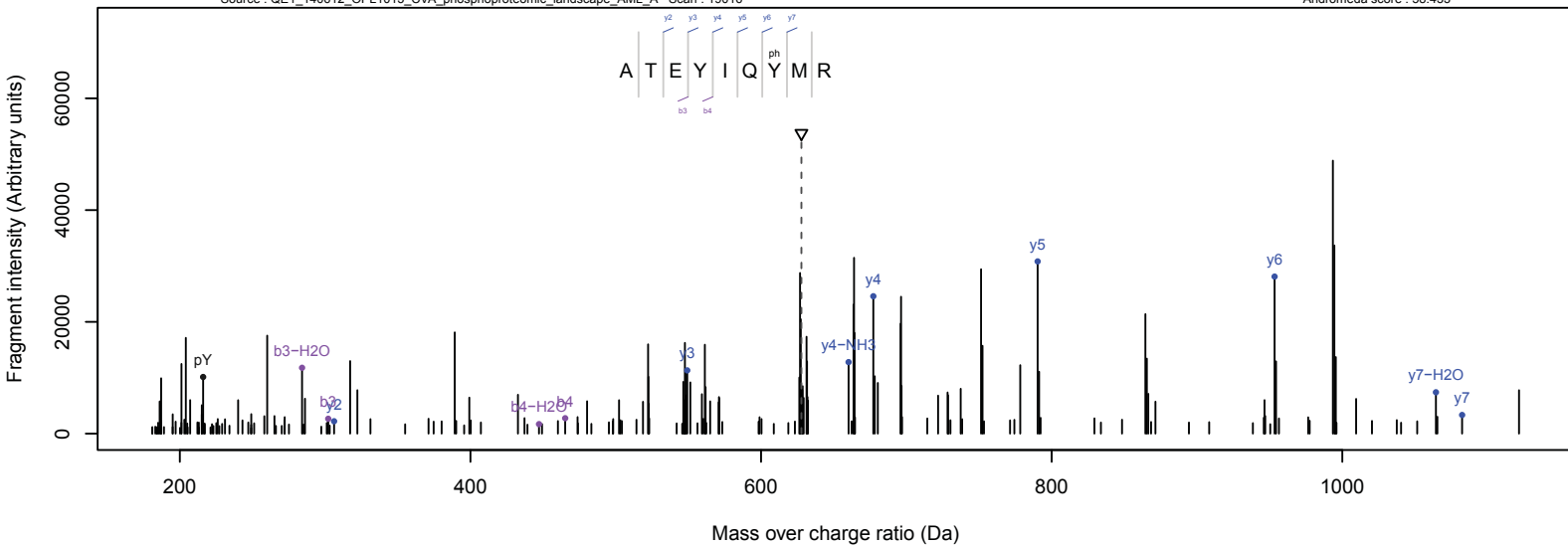
## (ac)M(ox)QY(ph)LNIKEDCNAM(ox)AFCAK-3+



ATEYIQ(ph)MR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 19016

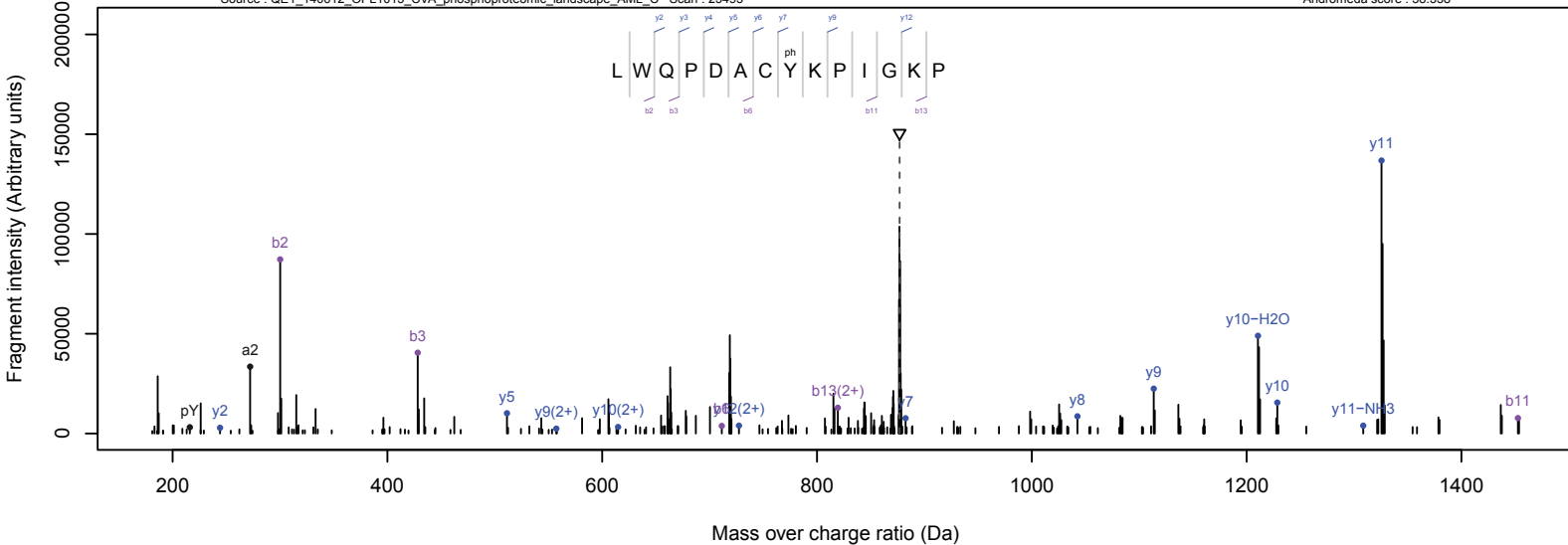
Andromeda score : 58.433



LWQPDACY(ph)KPIGKP-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23493

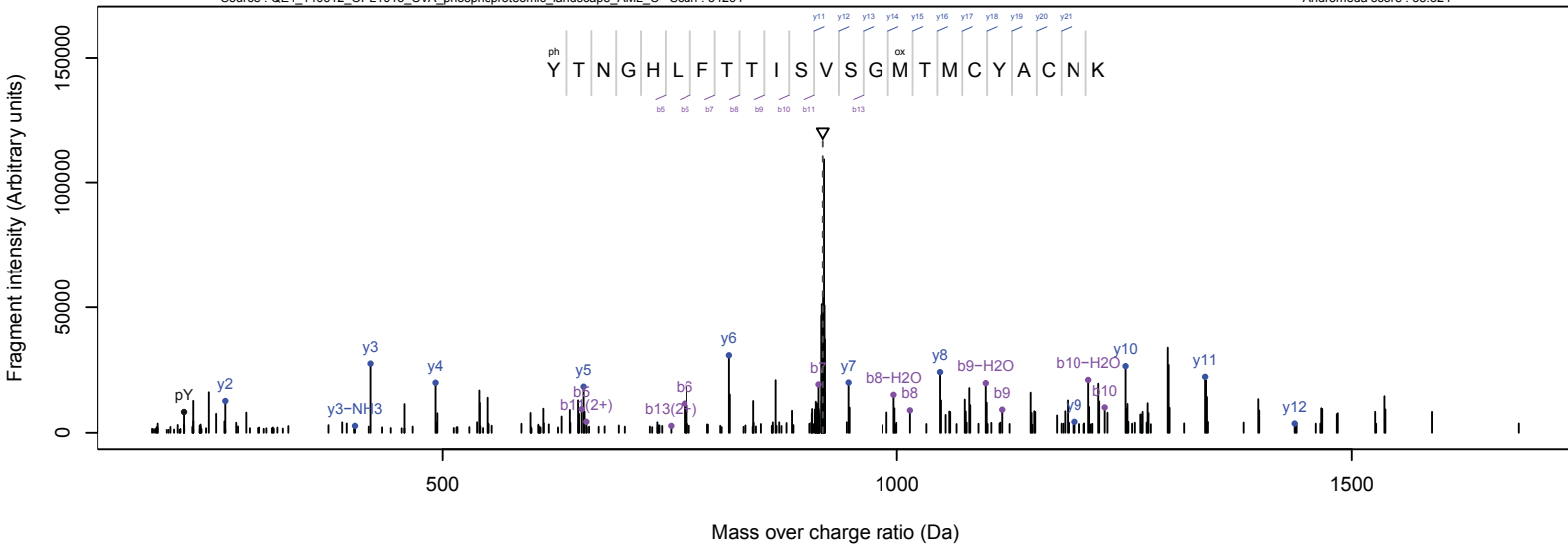
Andromeda score : 58.338



Y(ph)TNGHLFTTISVSGM(ox)TMCYACNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34264

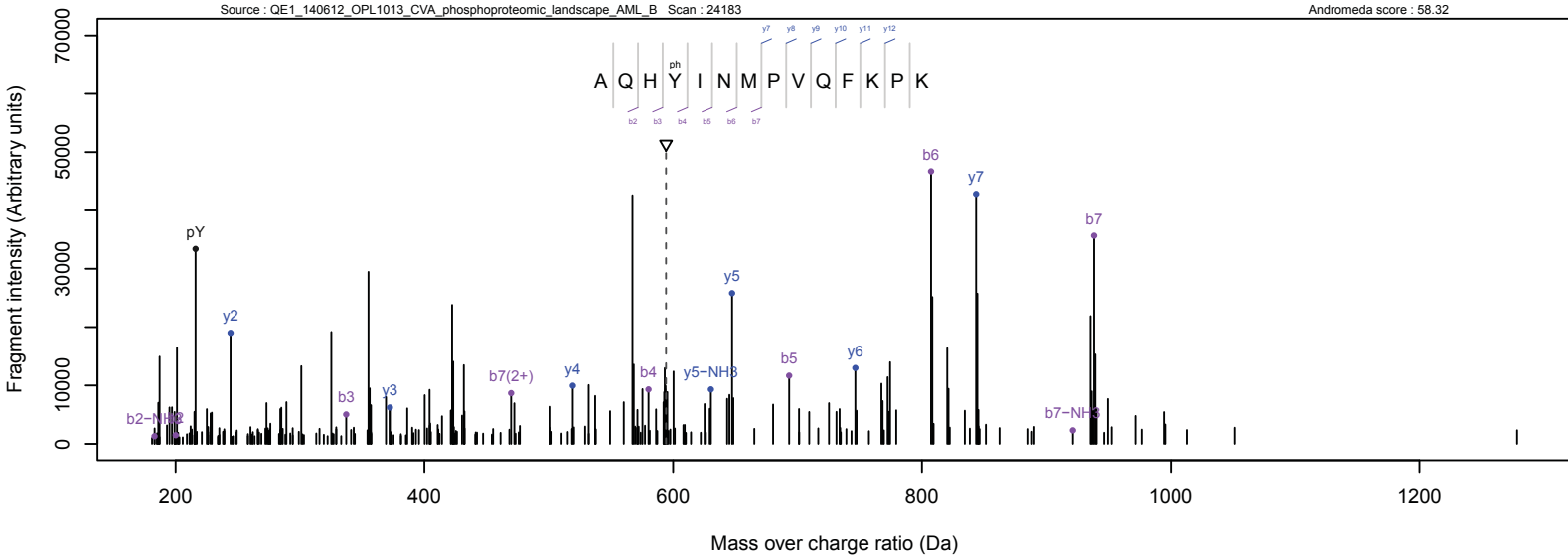
Andromeda score : 58.324



## AQHY(ph)INMPVQFKPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24183

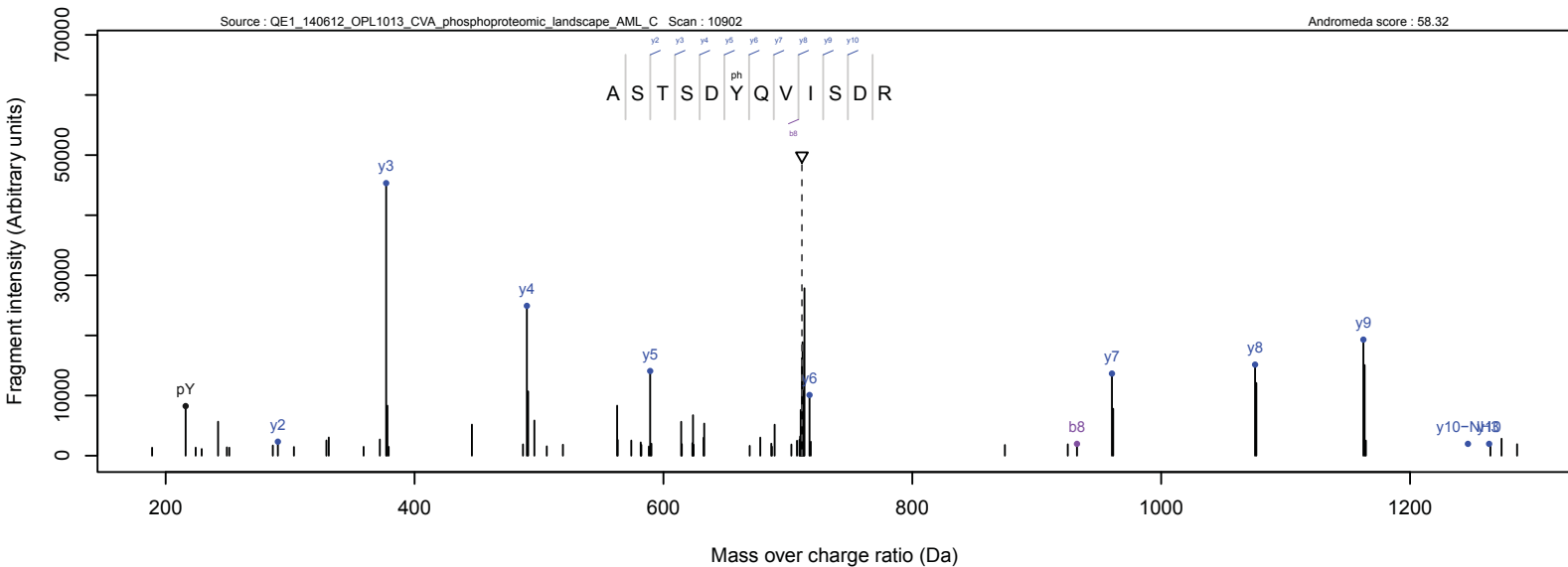
Andromeda score : 58.32



## ASTSDY(ph)QVISDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10902

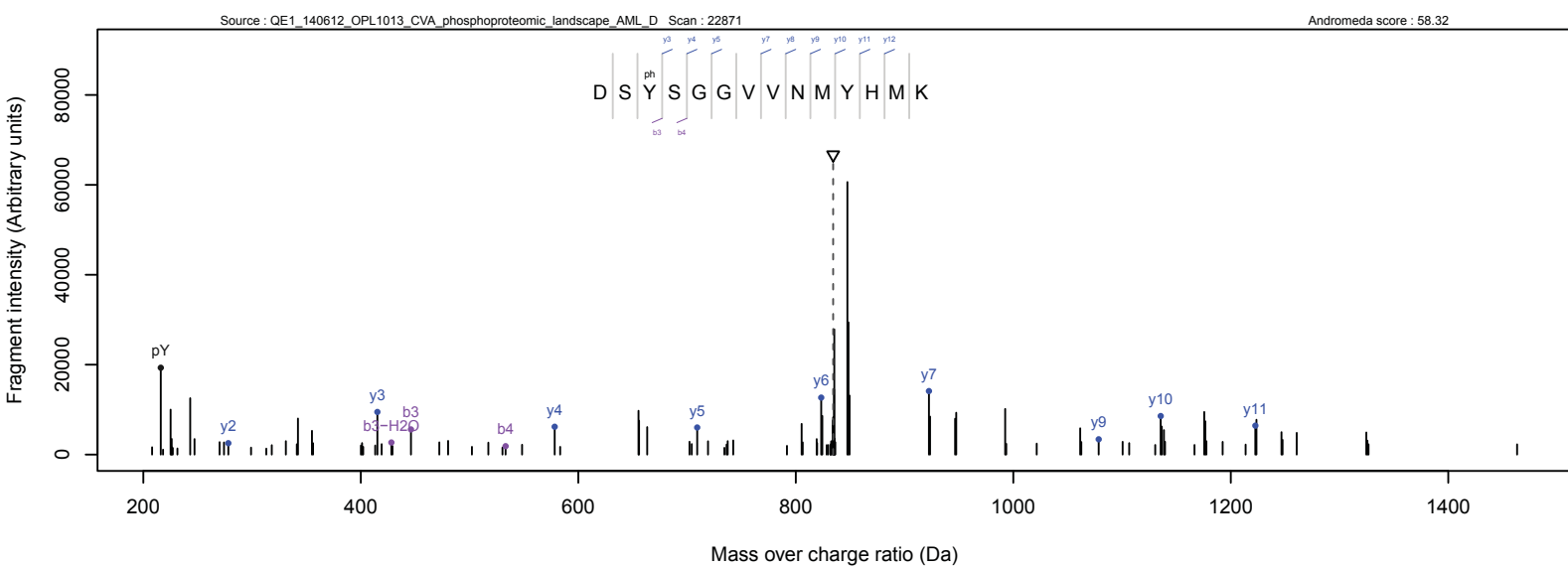
Andromeda score : 58.32



## DSY(ph)SGGVNMYHMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22871

Andromeda score : 58.32

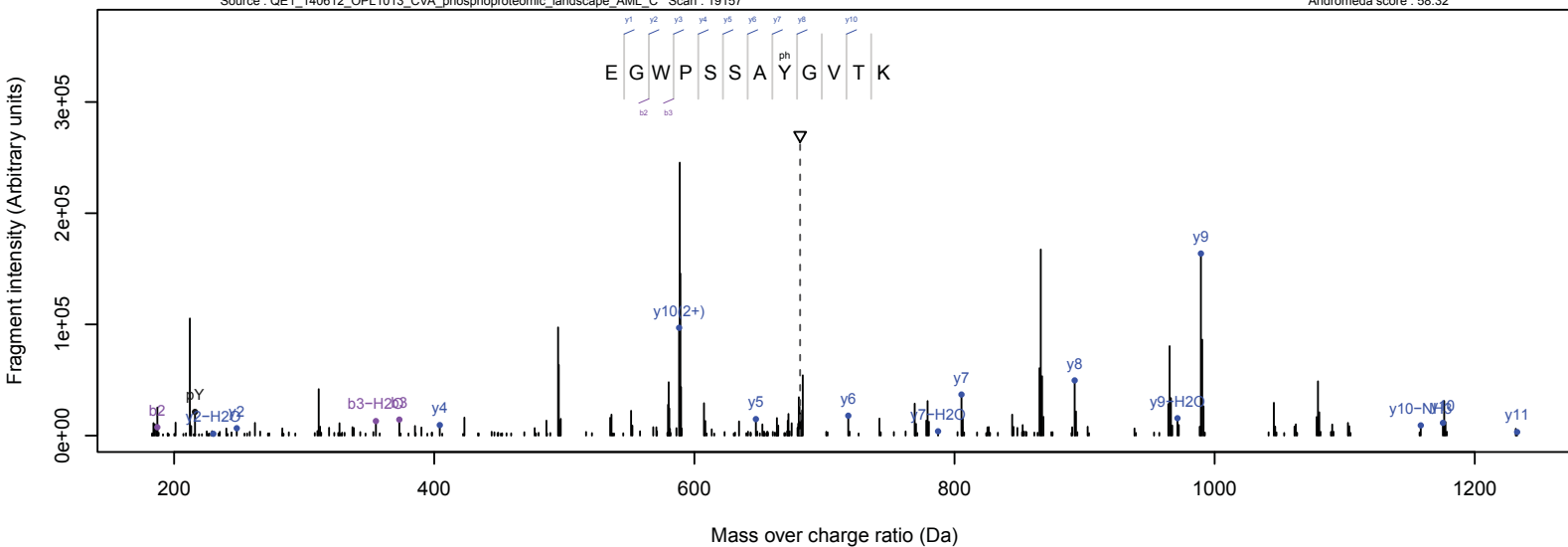




## EGWPSSAY(ph)GVTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19157

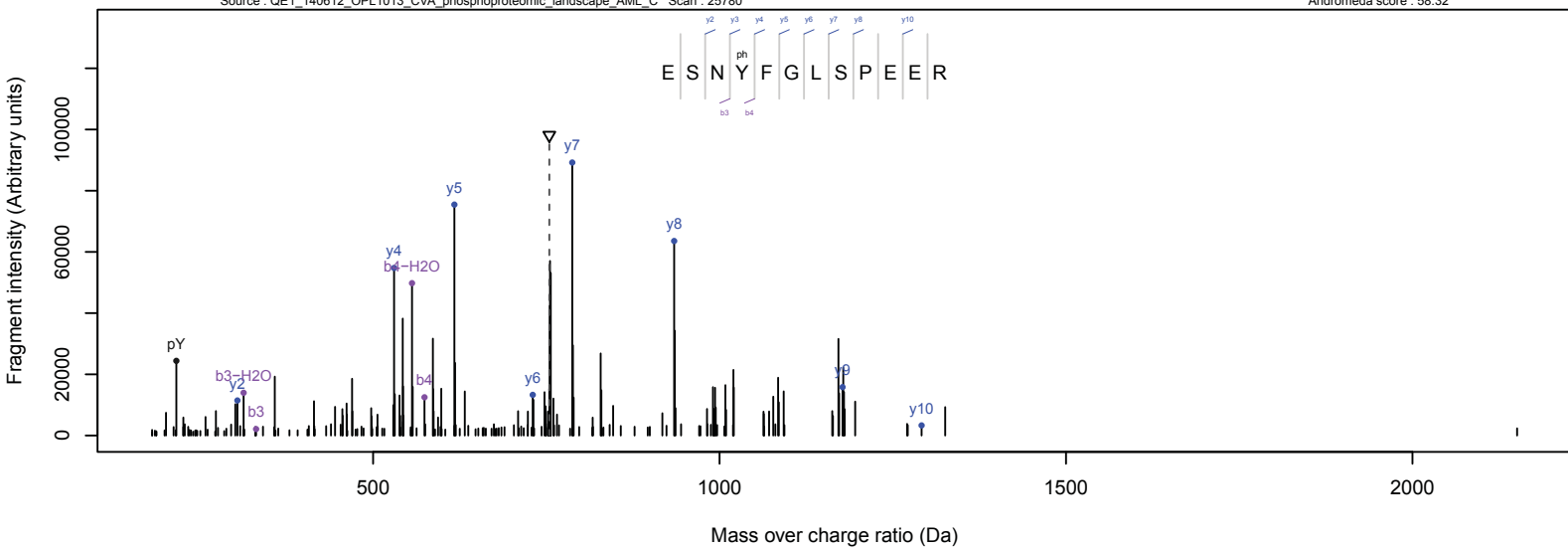
Andromeda score : 58.32



## ESNY(ph)FGLSPEER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25780

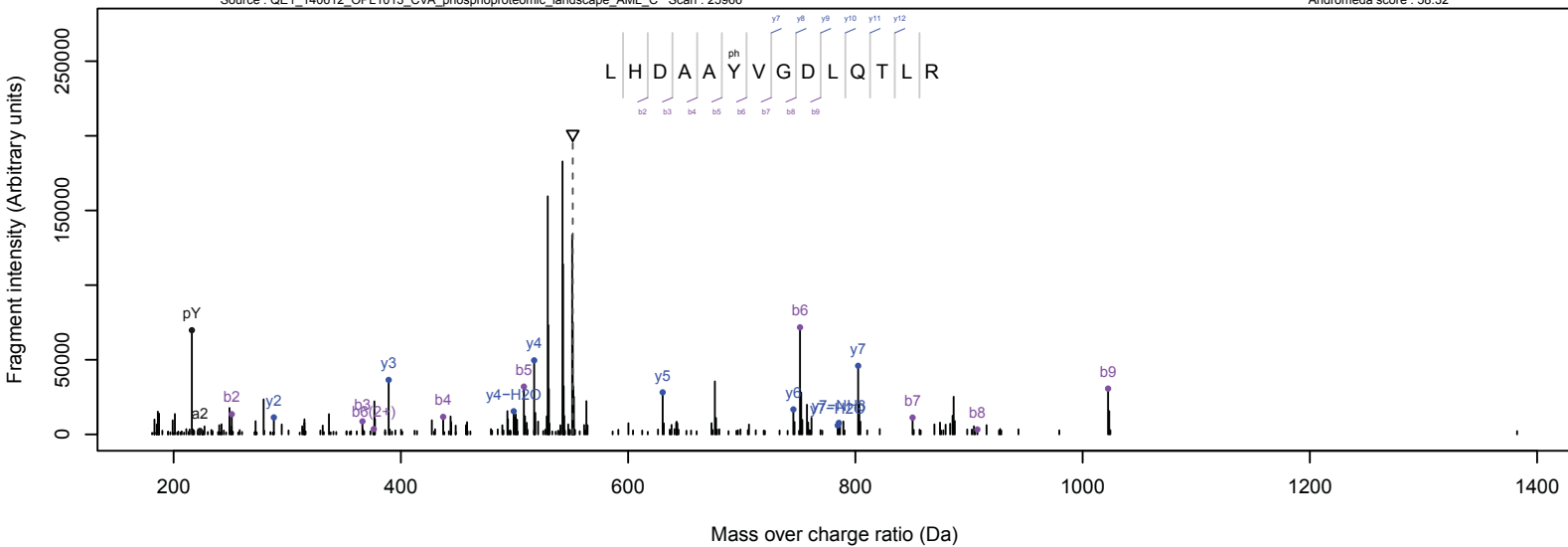
Andromeda score : 58.32



## LHDAAY(ph)VGDLQTLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25966

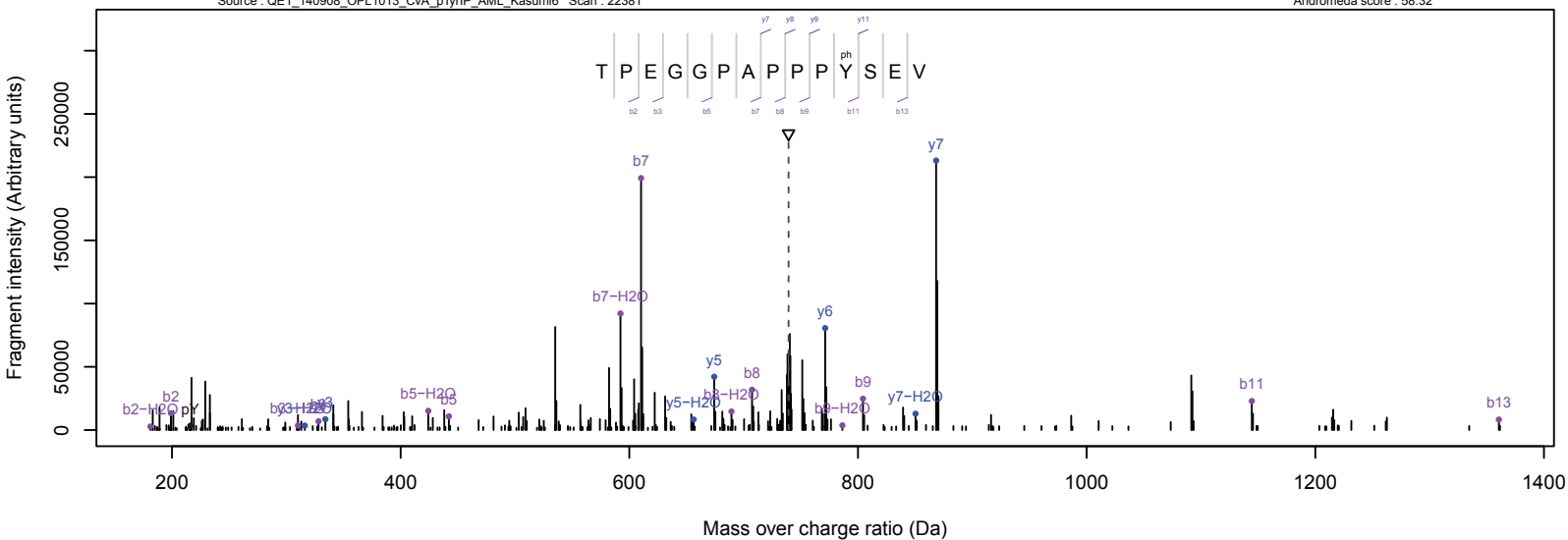
Andromeda score : 58.32



## TPEGGPAPPYSEV-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 22381

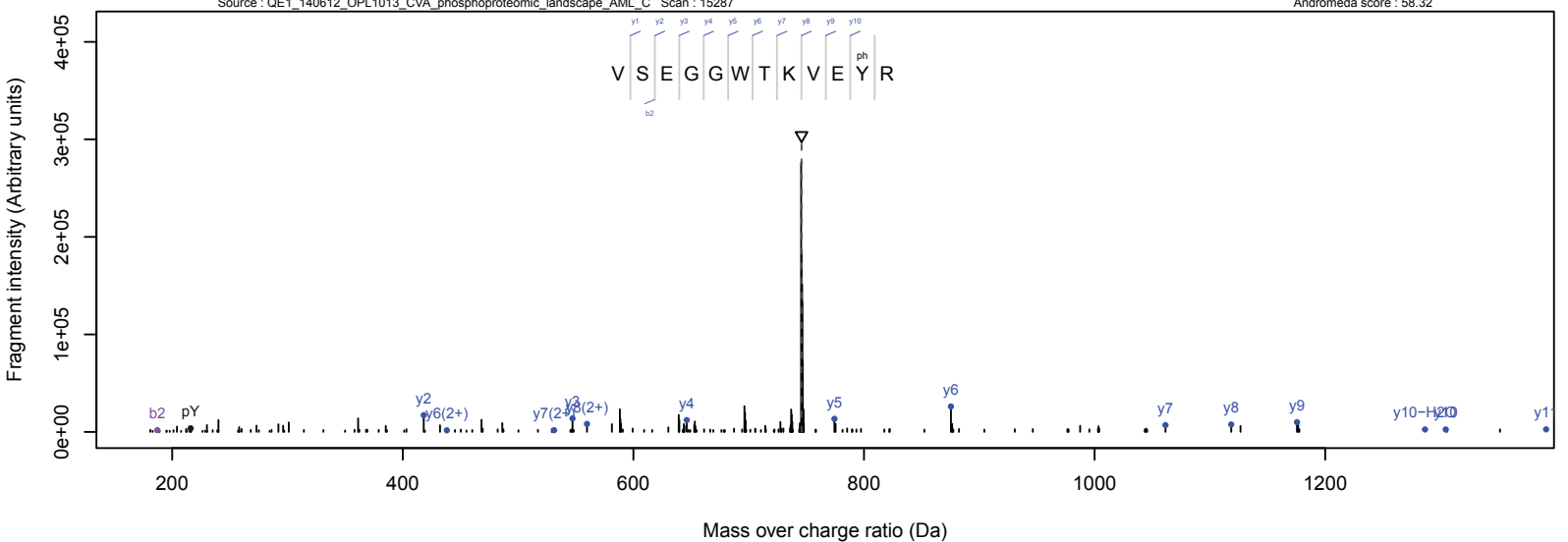
Andromeda score: 58.32



## VSEGGWTKVEYR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15287

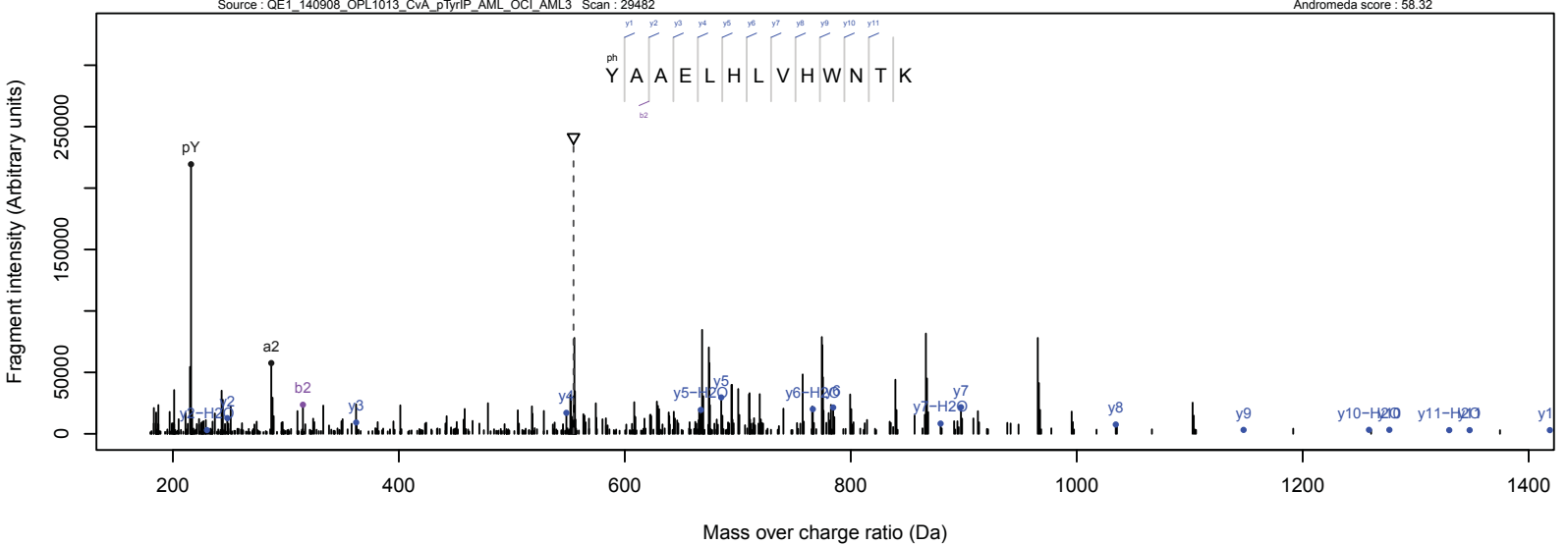
Andromeda score: 58.32



## Y(ph)AAELHLVHWNTK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 29482

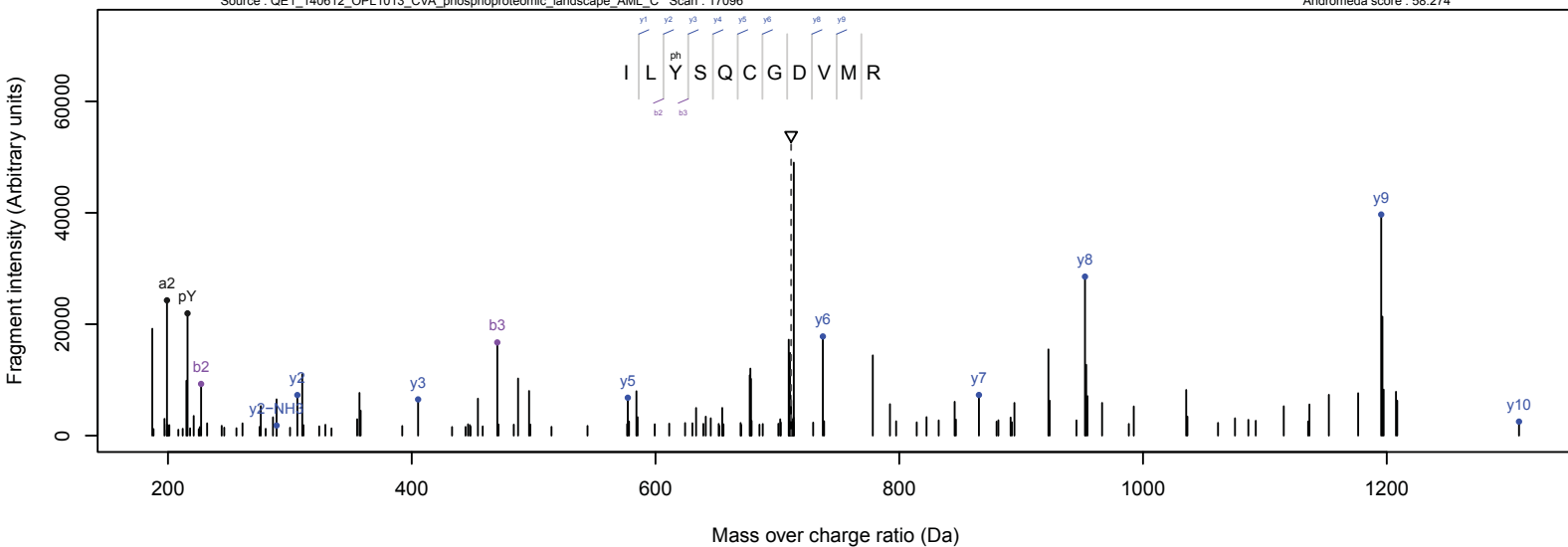
Andromeda score: 58.32



ILY(ph)SQCGDVMR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17096

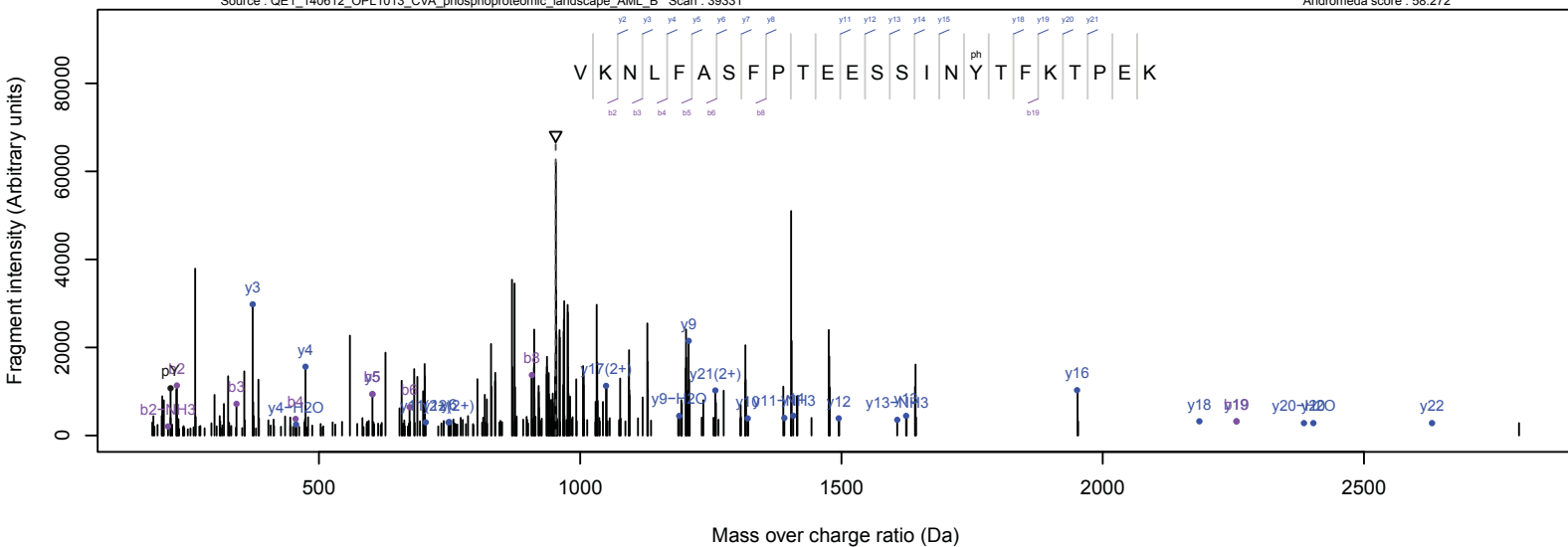
Andromeda score: 58.274



VKNLFASFTEESSINY(ph)TFKTPEK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 39331

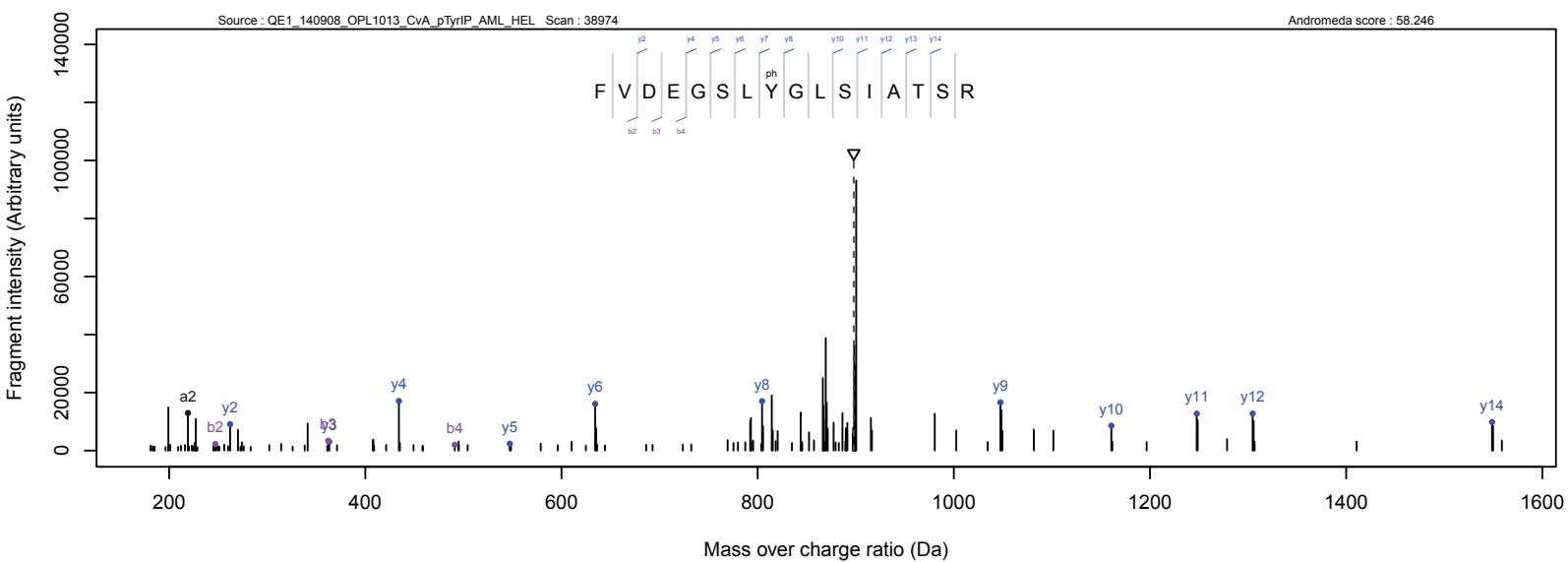
Andromeda score: 58.272



FVDEGSLYGLSIATSR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 38974

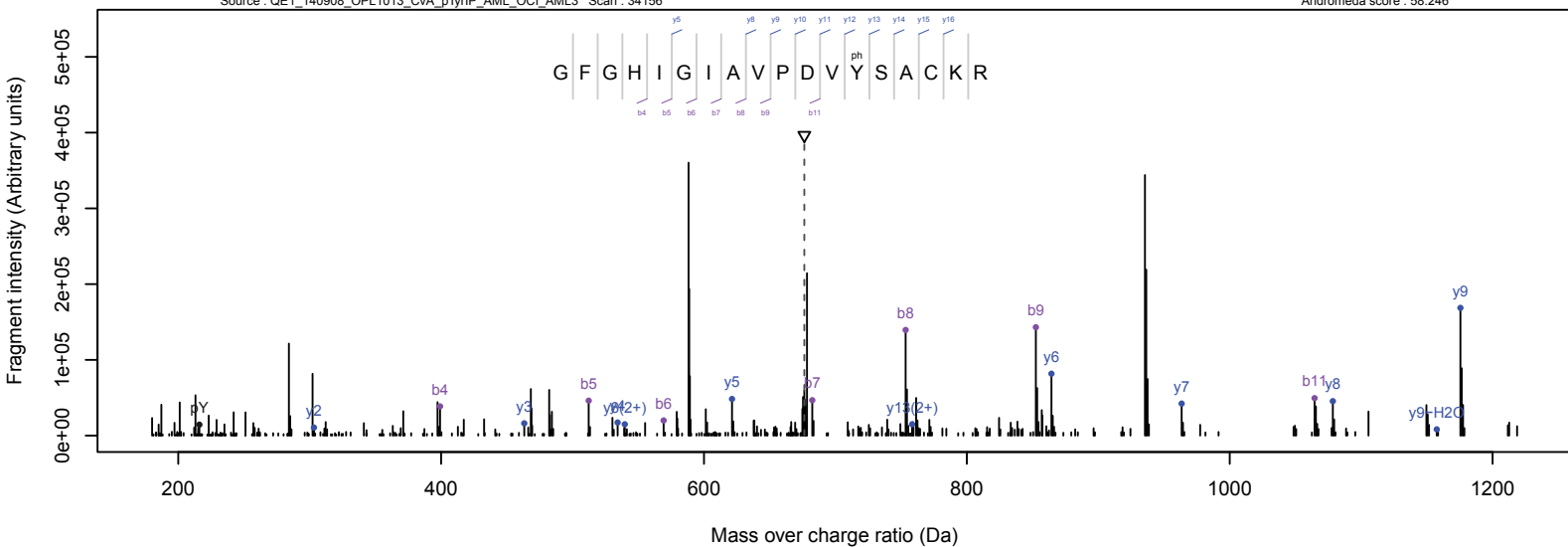
Andromeda score: 58.246



GFGHIGIAPDVY(ph)SACKR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 34156

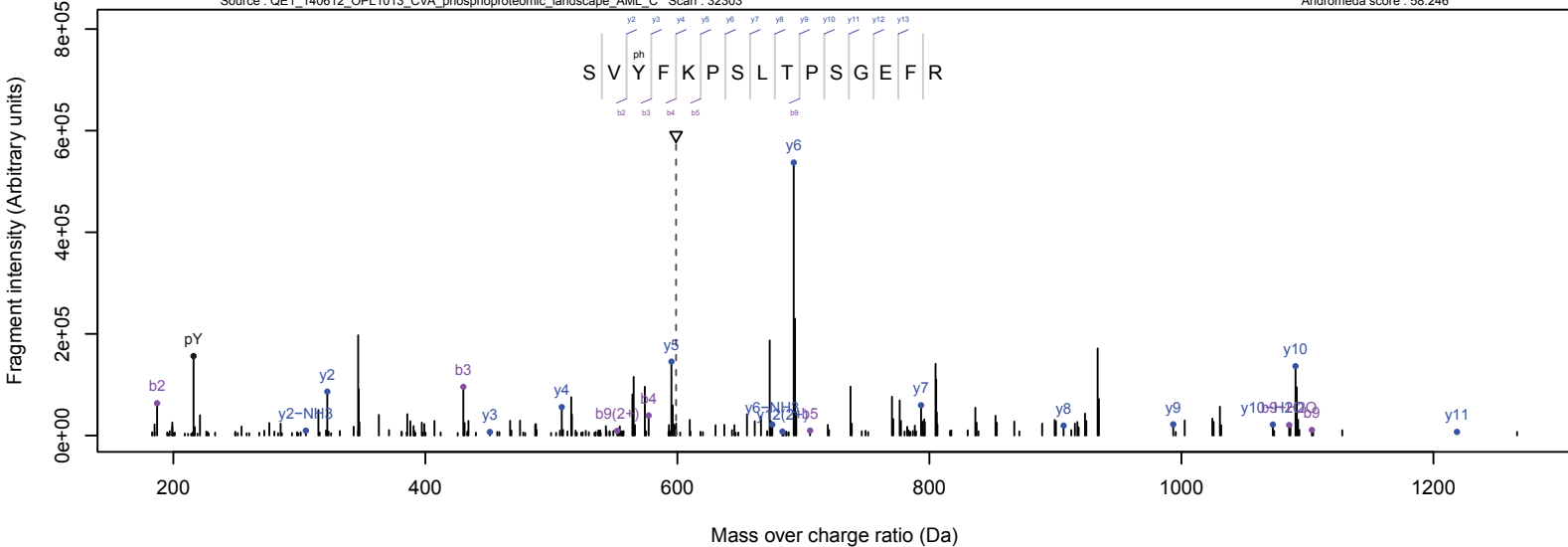
Andromeda score : 58.246



SVY(ph)FKPSLTSPGSEFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32303

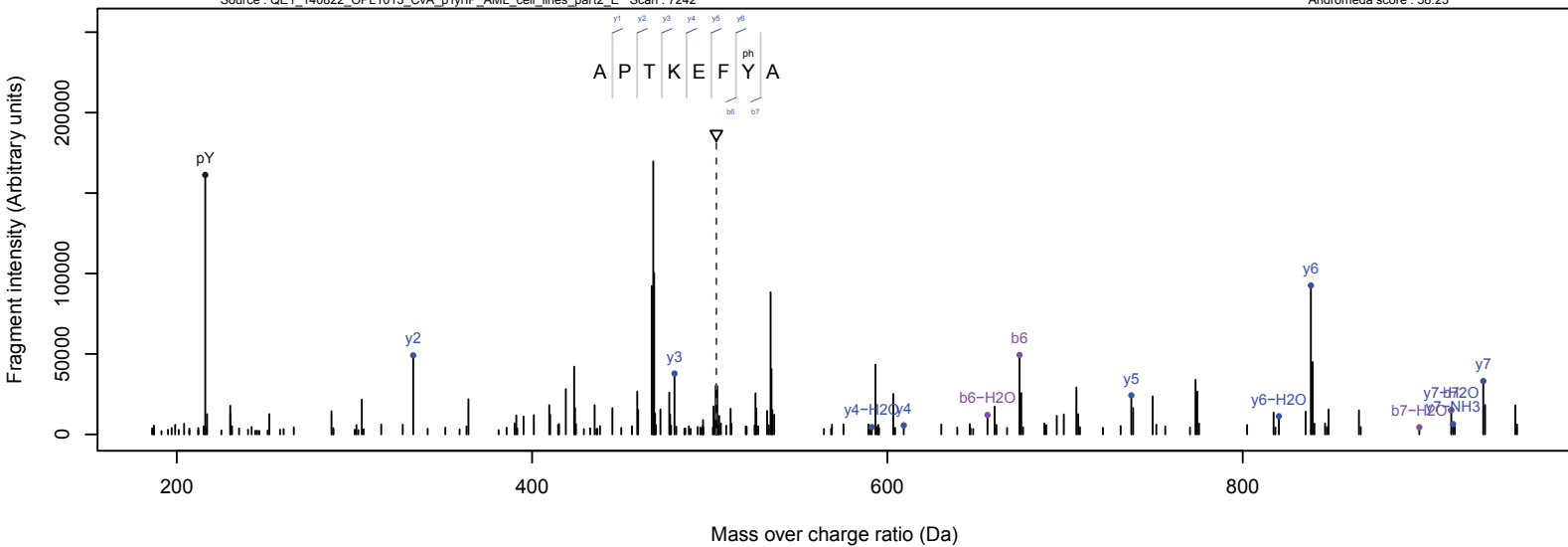
Andromeda score : 58.246



APTKEFY(ph)A-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 7242

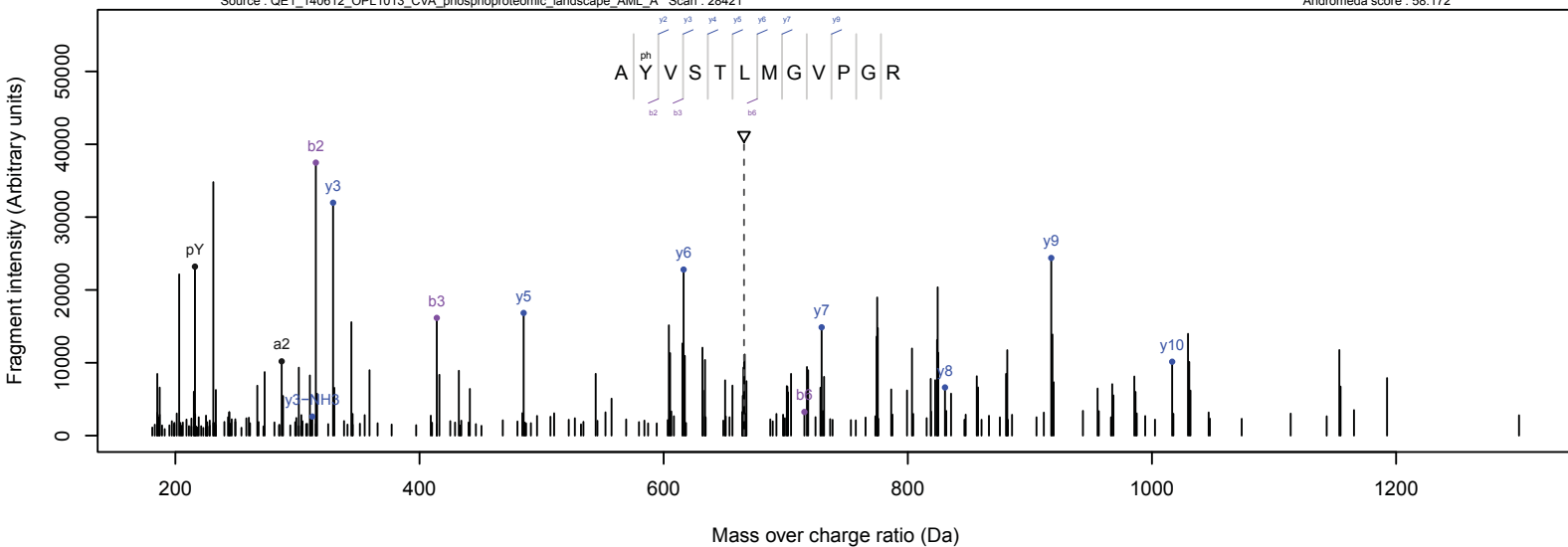
Andromeda score : 58.23



## AY(ph)VSTLMGVPGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 28421

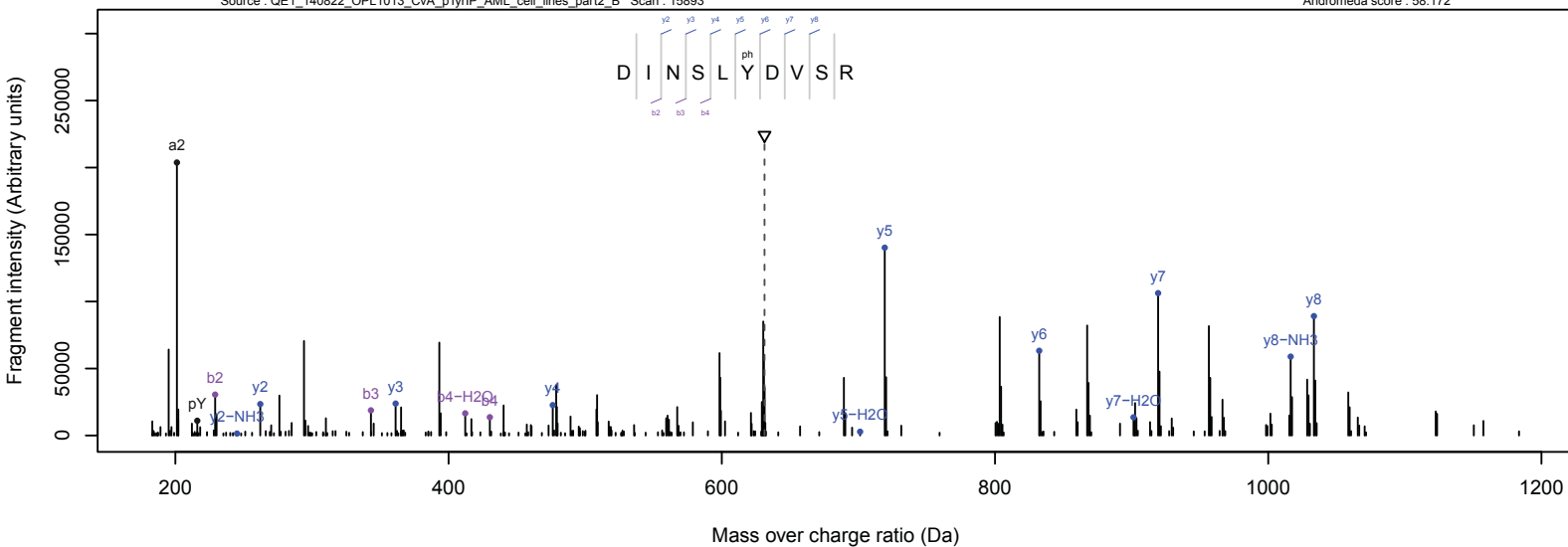
Andromeda score: 58.172



## DINSLY(ph)DVSR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 15893

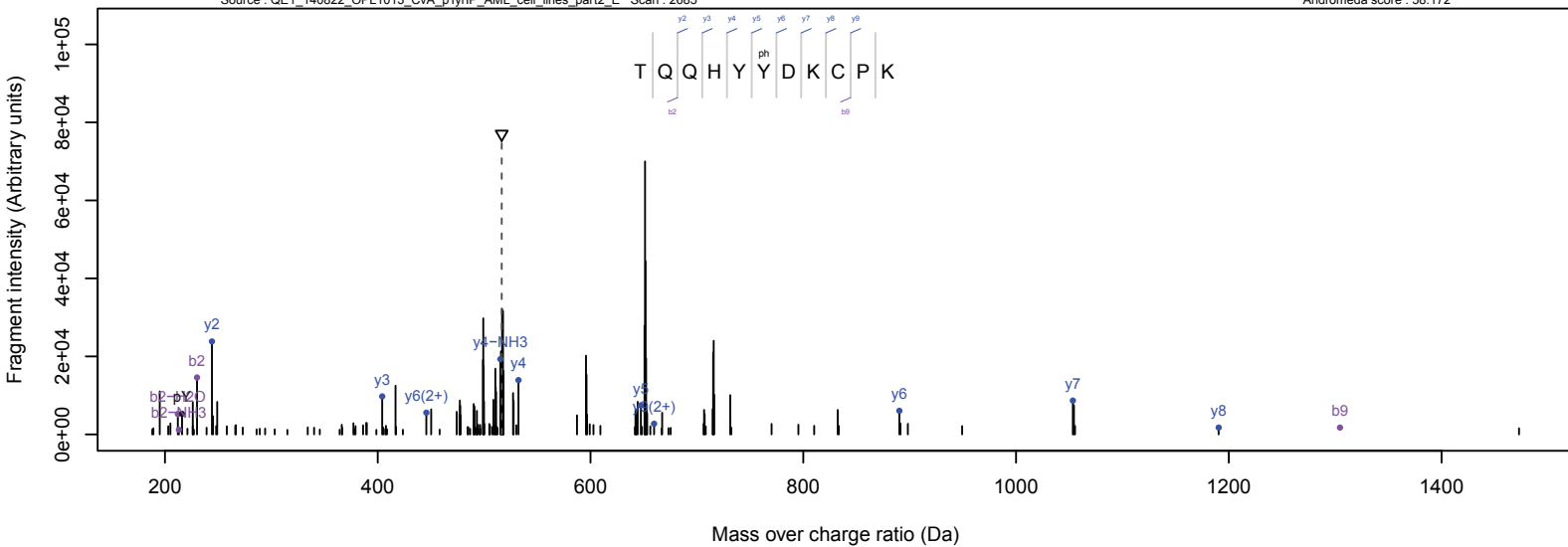
Andromeda score: 58.172



## TQQHY(ph)DKCPK-3+

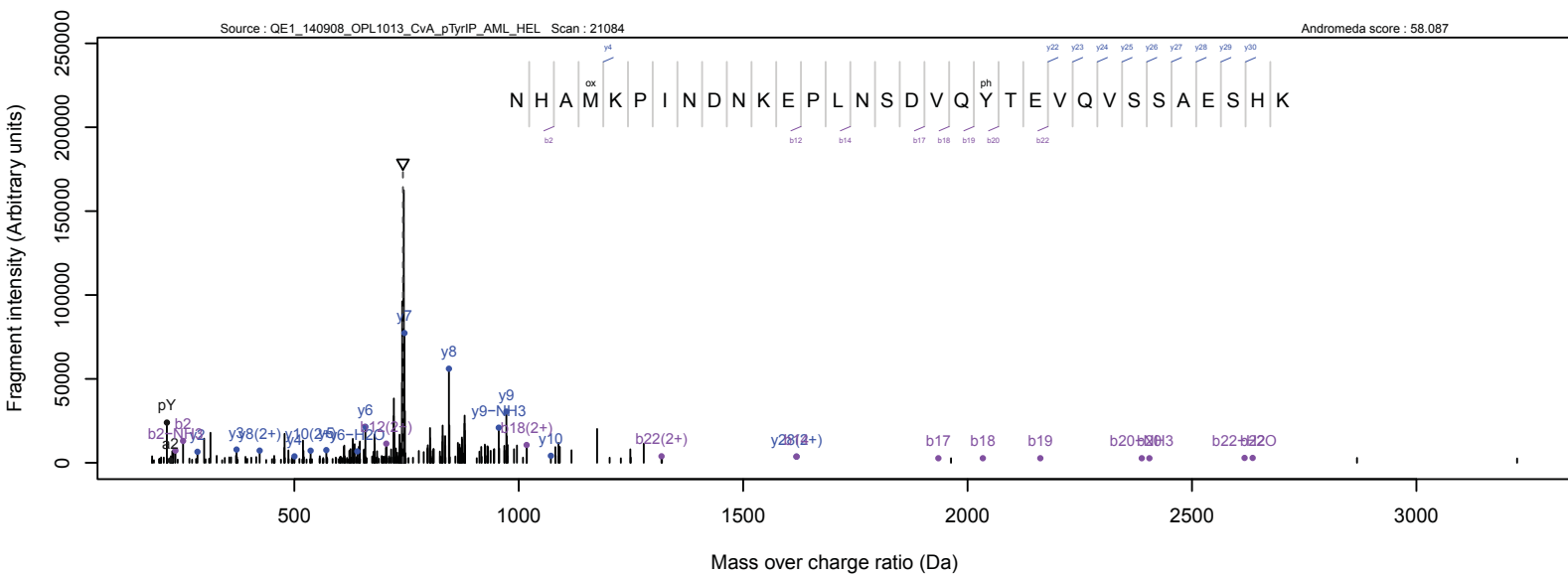
Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 2685

Andromeda score: 58.172

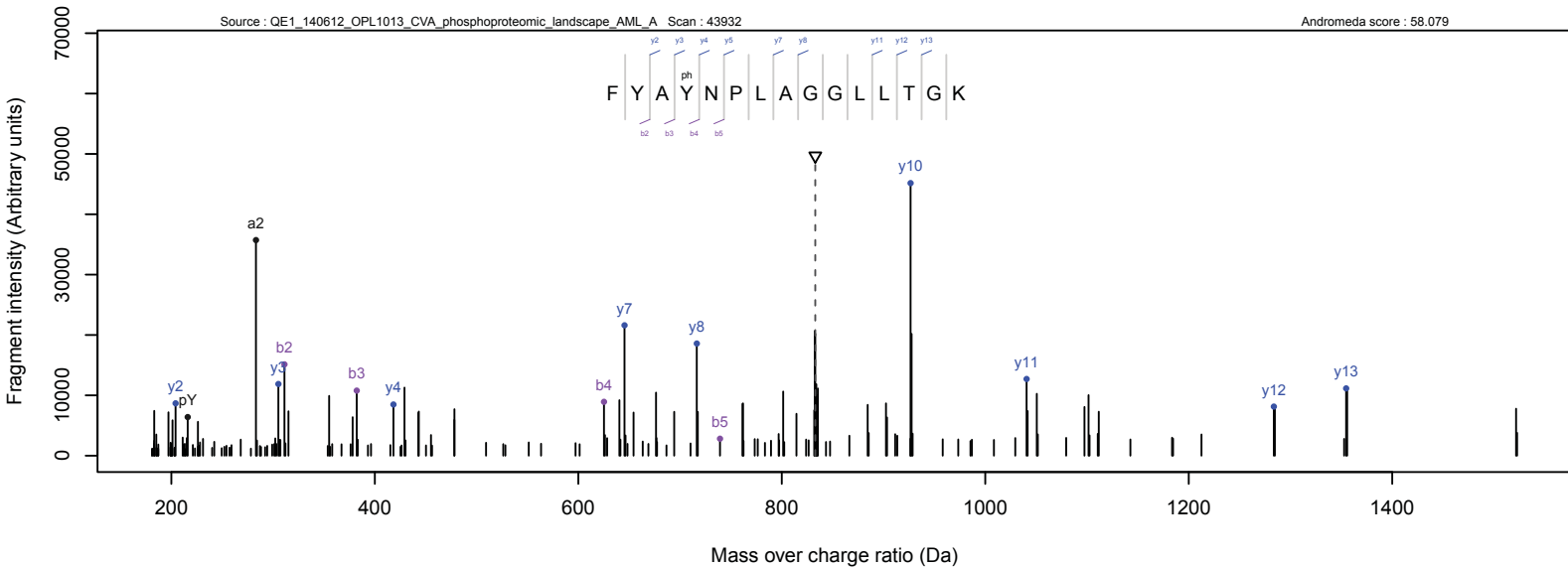




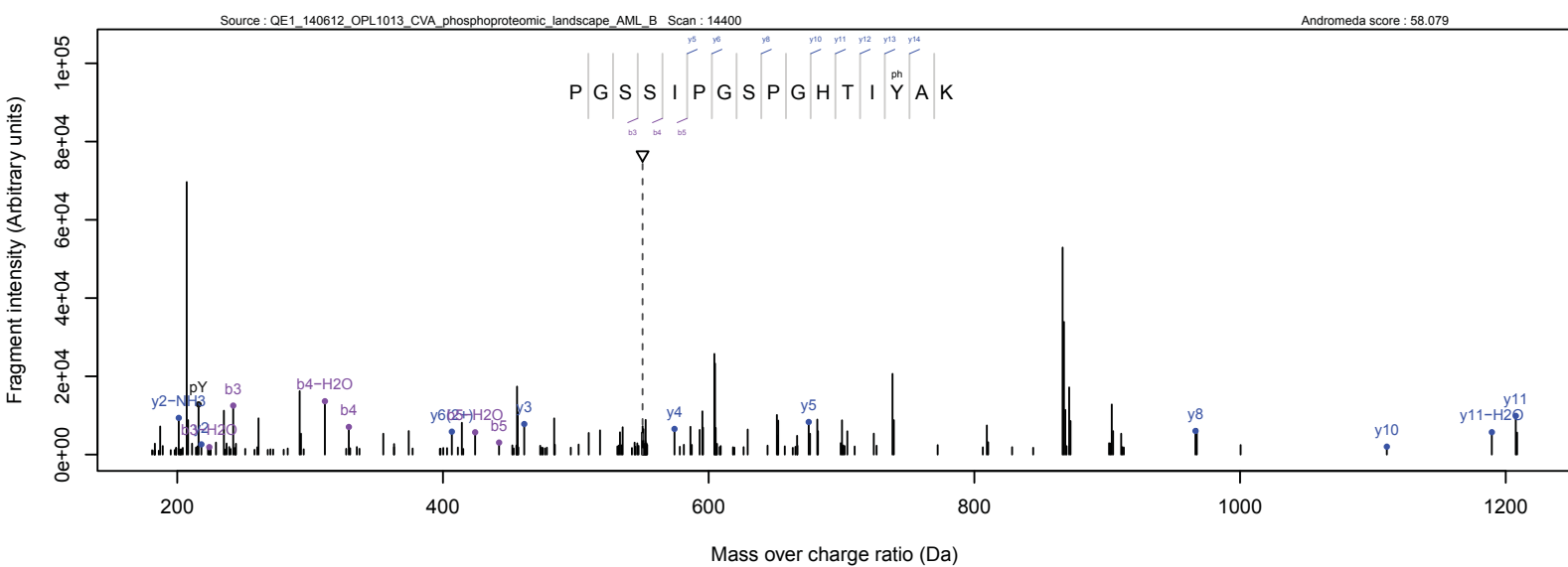
NHAM(ox)KPINDNKEPLNSDVQY(ph)TEVQVSSAESHK-5+



FYAY(ph)NPLAGLLTGK-2+



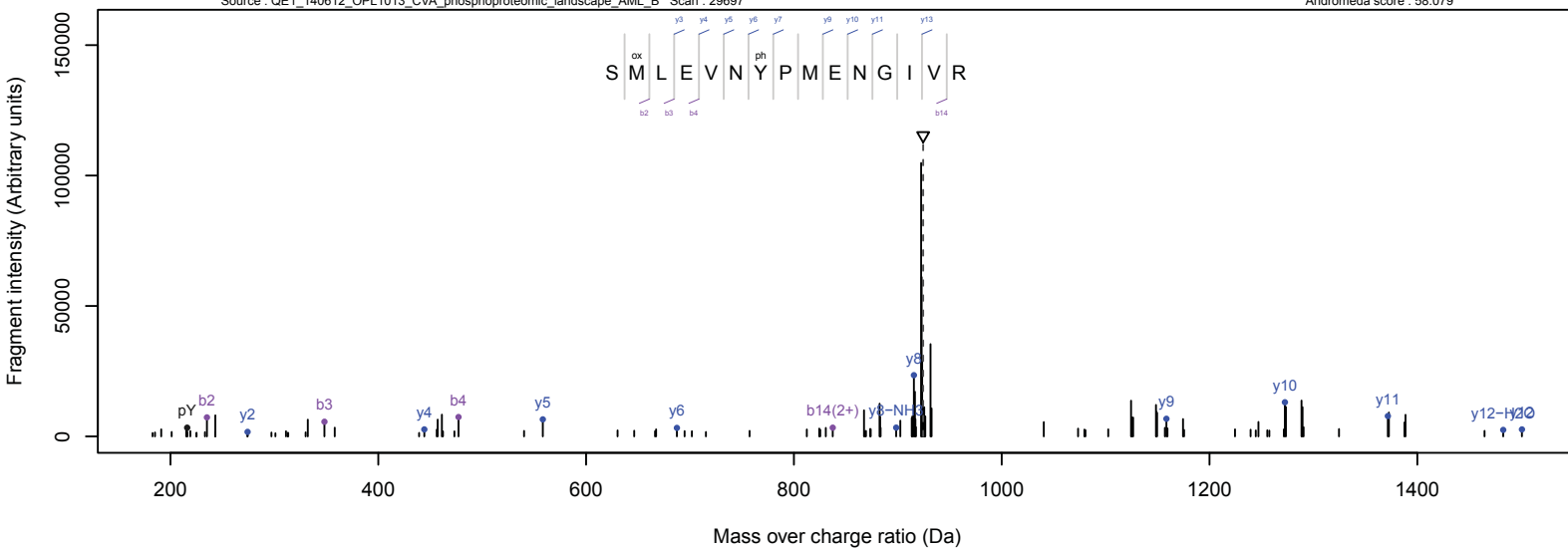
PGSSIPGSPGHTIY(ph)AK-3+



## SM(ox)LEVNY(ph)PMENGIVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29697

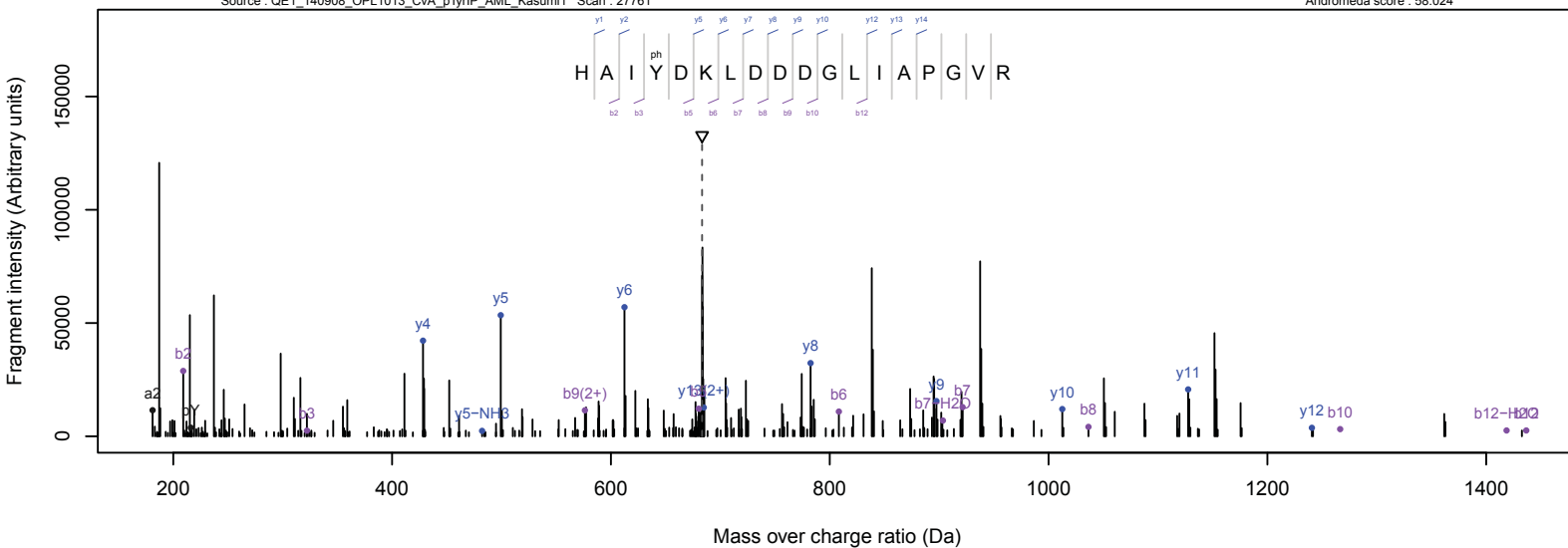
Andromeda score : 58.079



## HAIY(ph)DKLDDGLIAPGVR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 27761

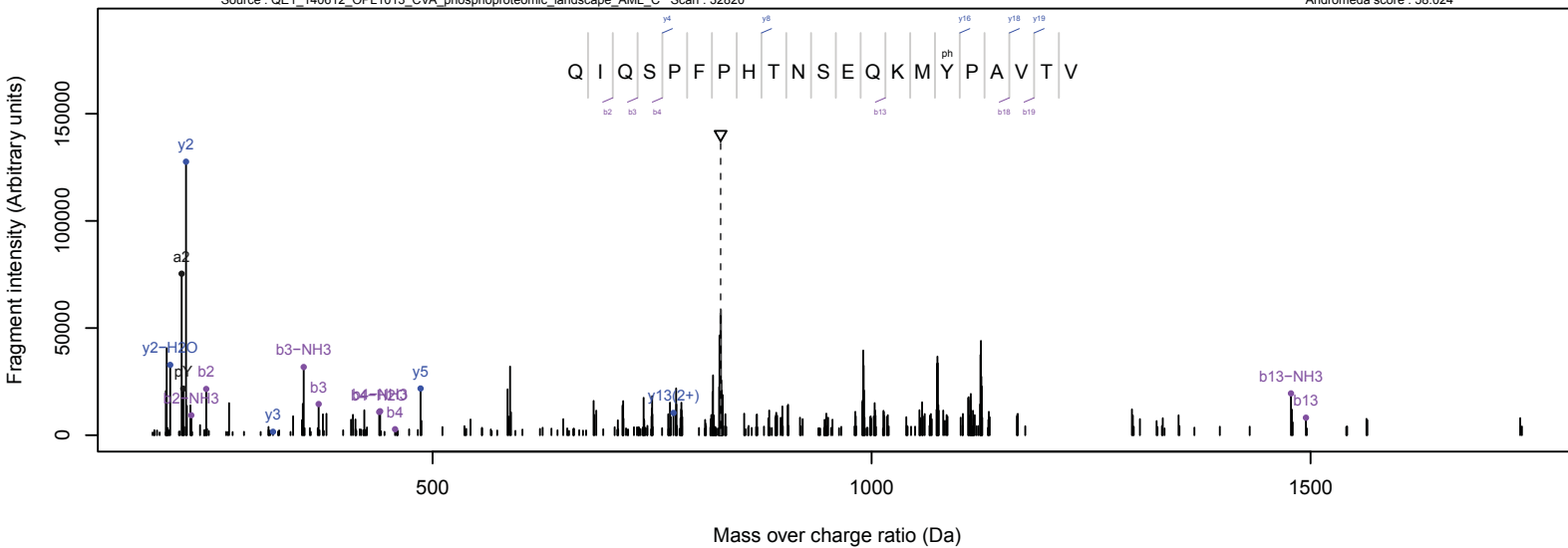
Andromeda score : 58.024



## QIQSPFPHTNSEQKMY(ph)PAVTV-3+

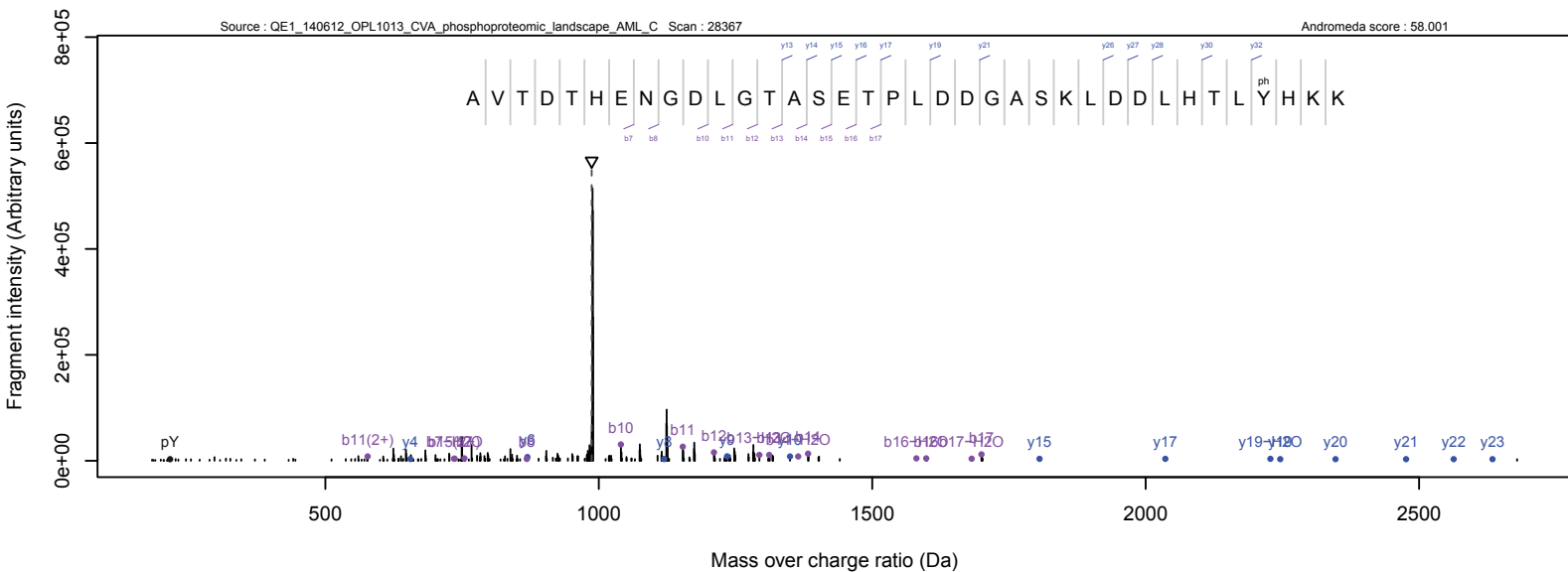
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32820

Andromeda score : 58.024

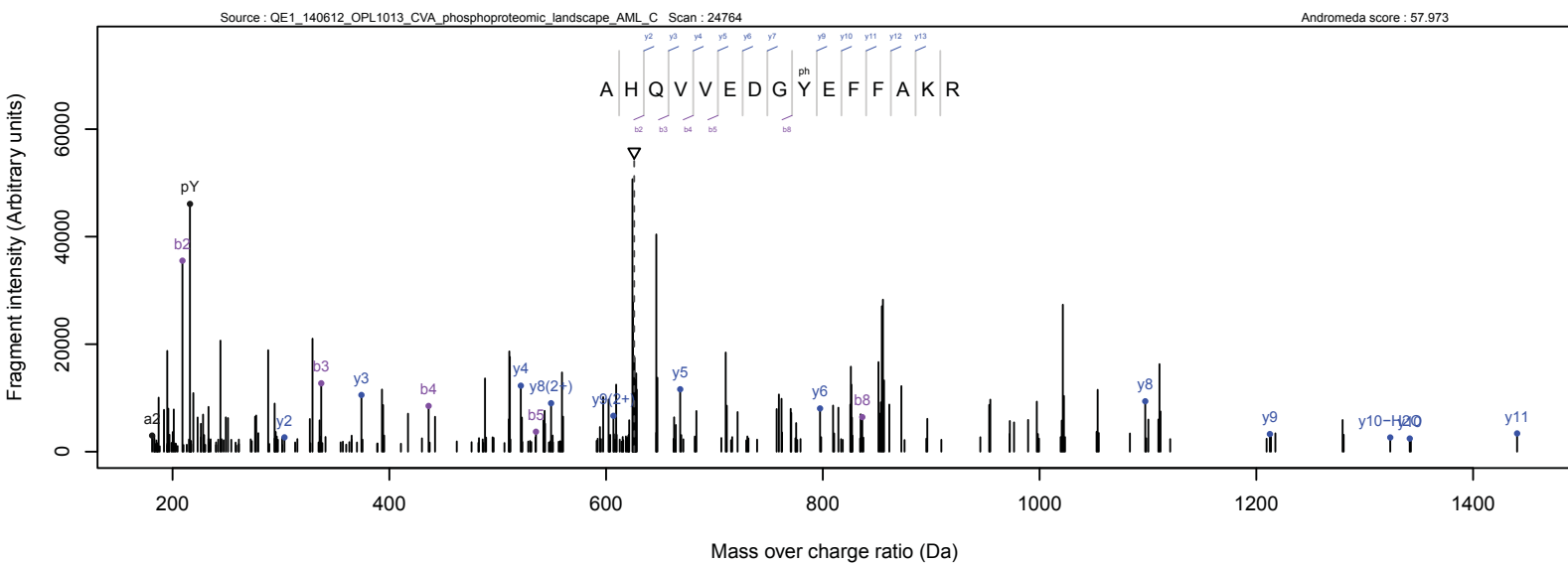




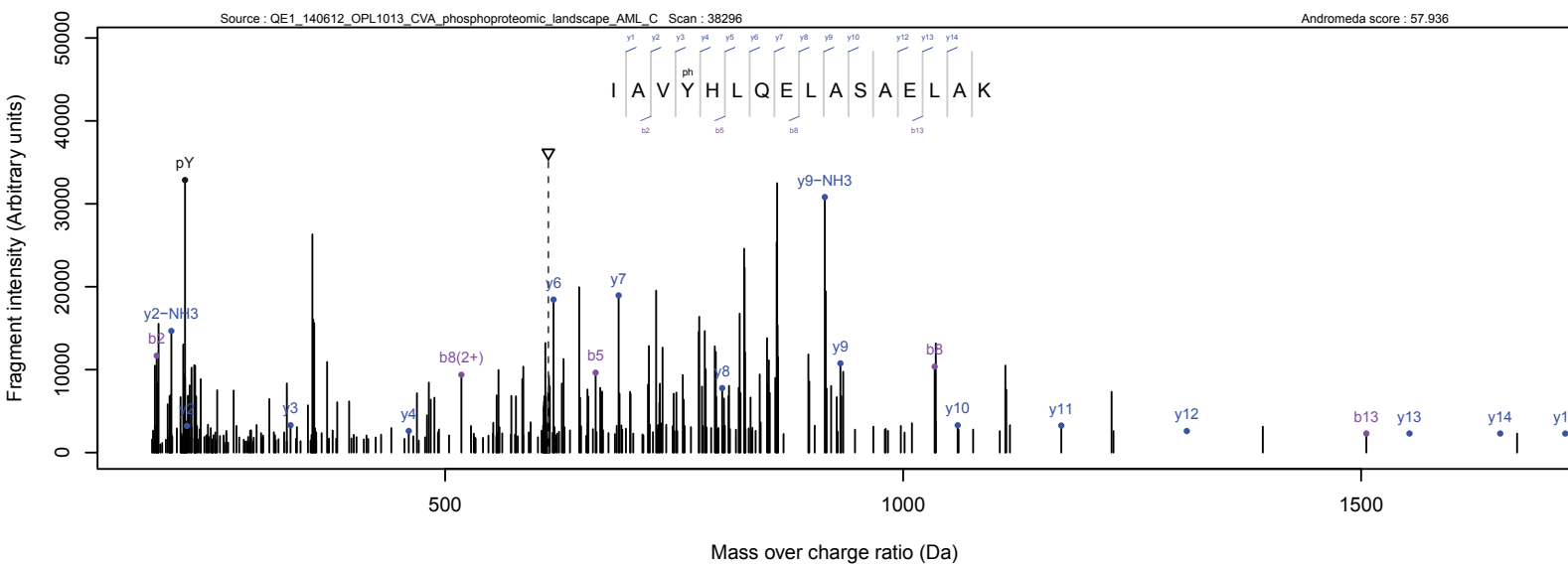
AVTDTHE<sup>ph</sup>NGDLGTASE<sup>ph</sup>TPLDDGASKLDDLHTLY<sup>ph</sup>HKK-4+



AHQVVEDGY<sup>ph</sup>EFFAKR-3+



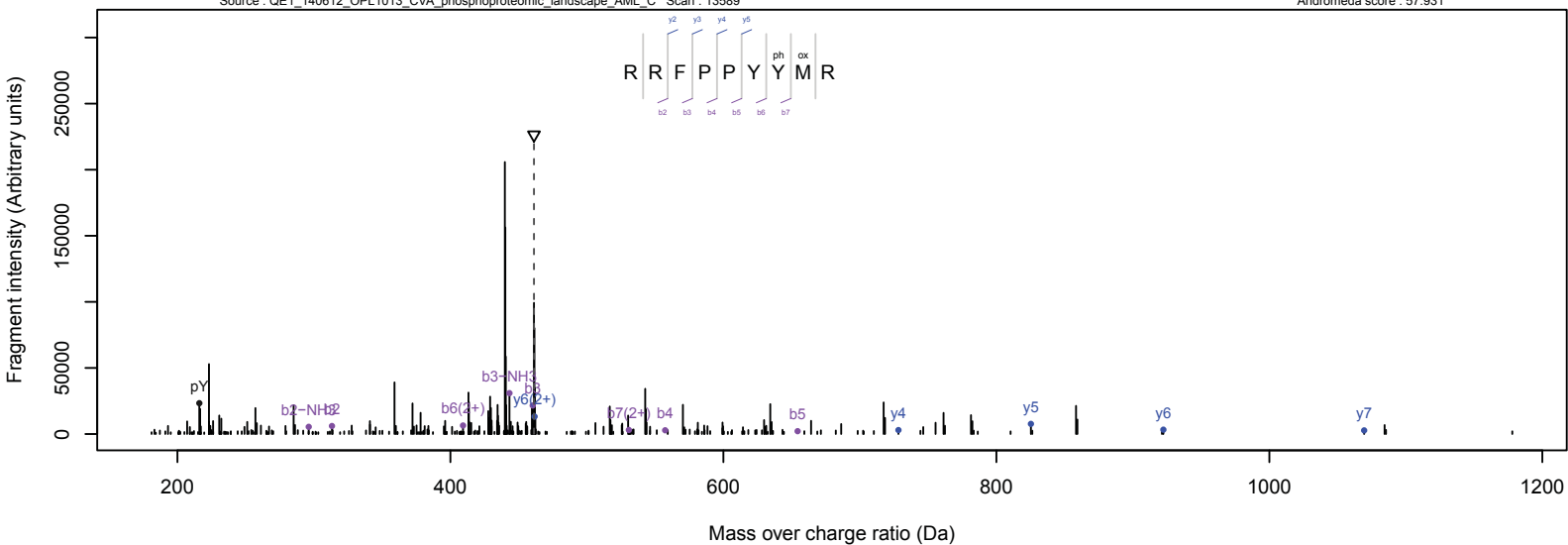
IAVY<sup>ph</sup>HLQELASAE<sup>ph</sup>LAK-3+



## RRFPYY(ph)M(ox)R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13589

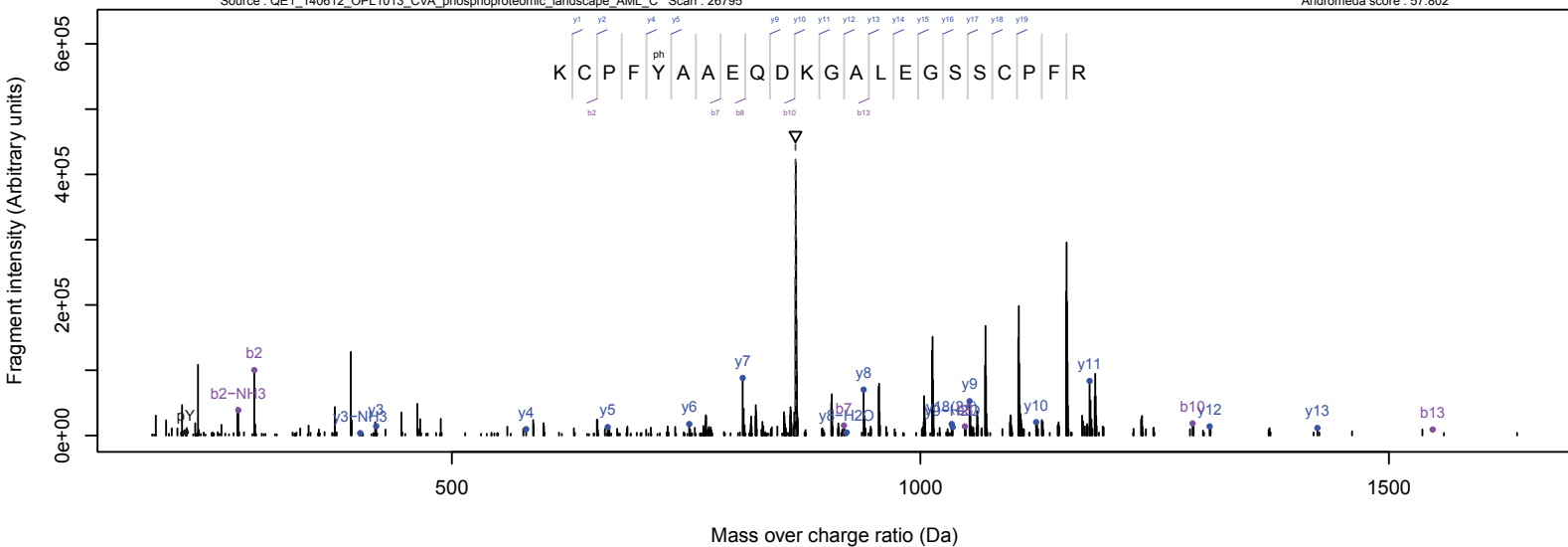
Andromeda score : 57.931



## KCPFY(ph)AAEQDKGALGSSCPFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26795

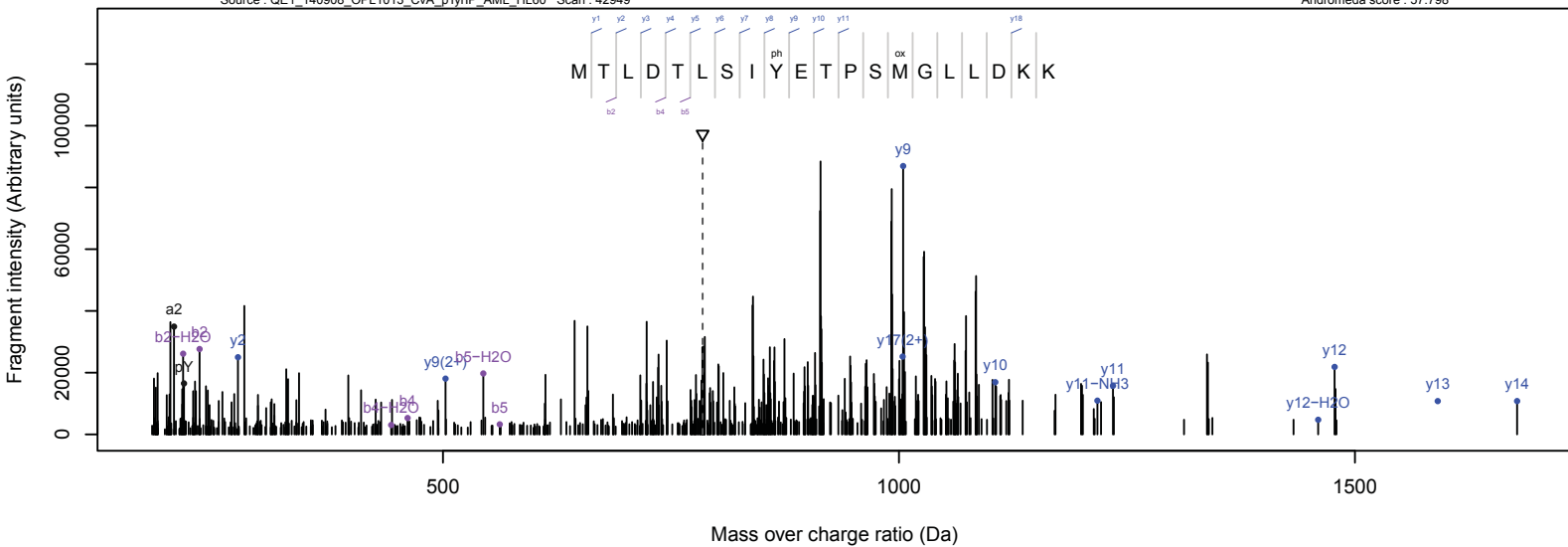
Andromeda score : 57.802

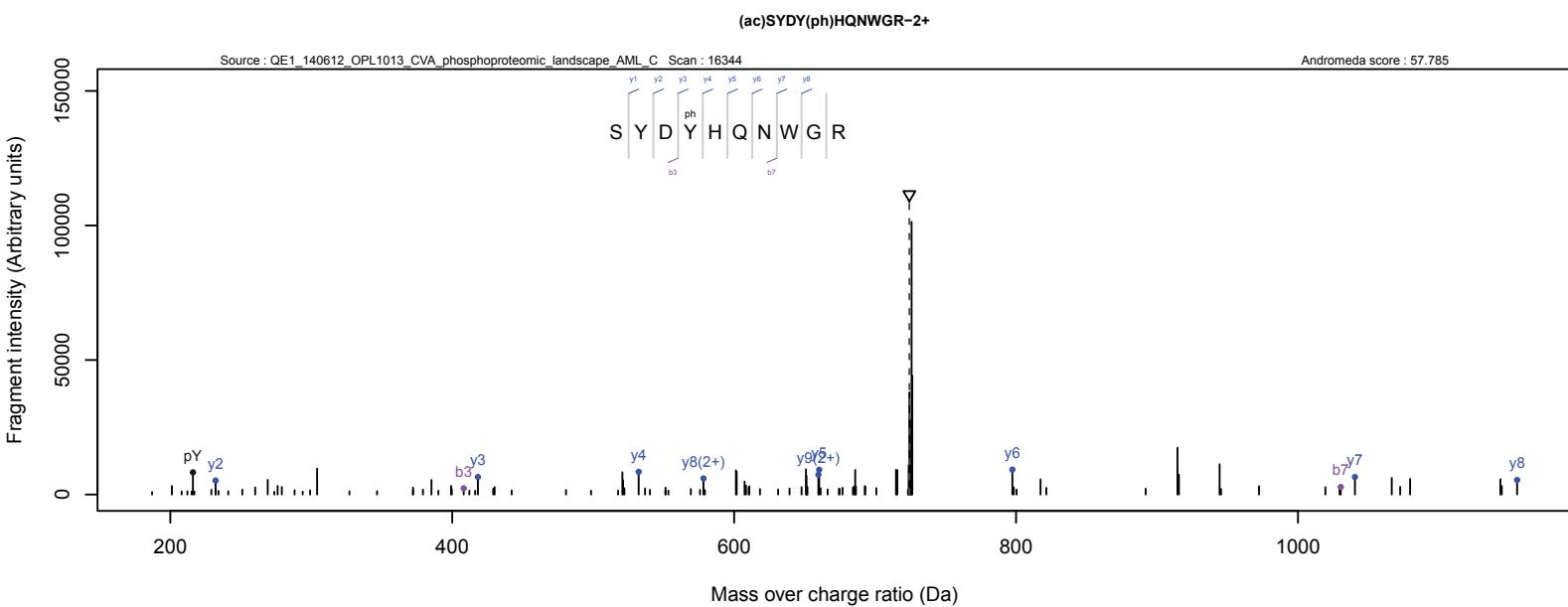
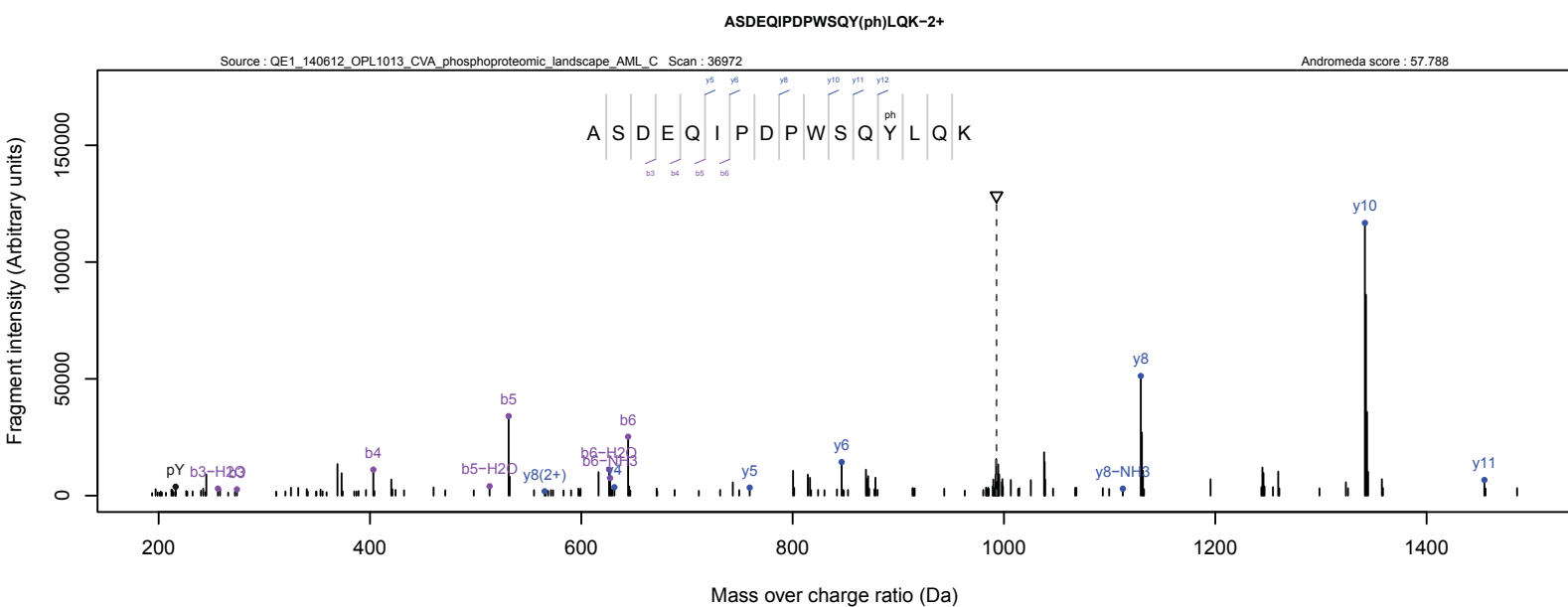
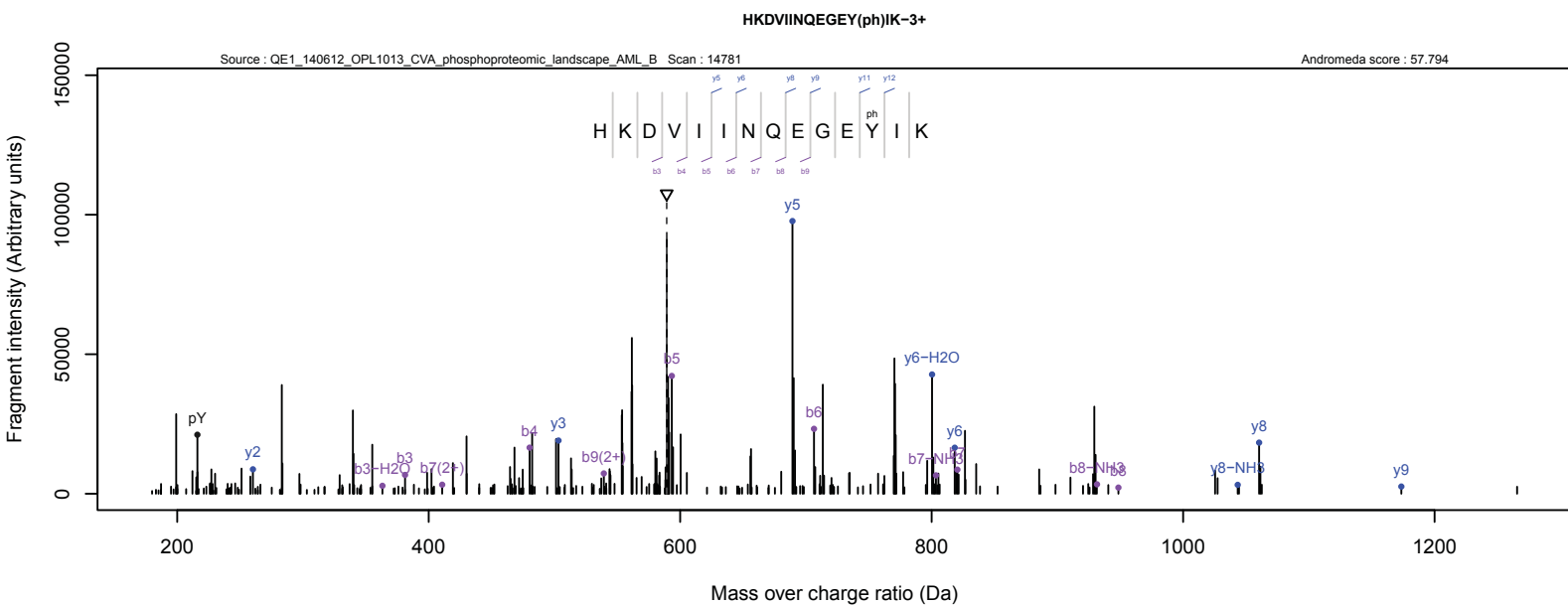


## MTLDTLSIY(ph)ETPSM(ox)GLLDKK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 42949

Andromeda score : 57.798

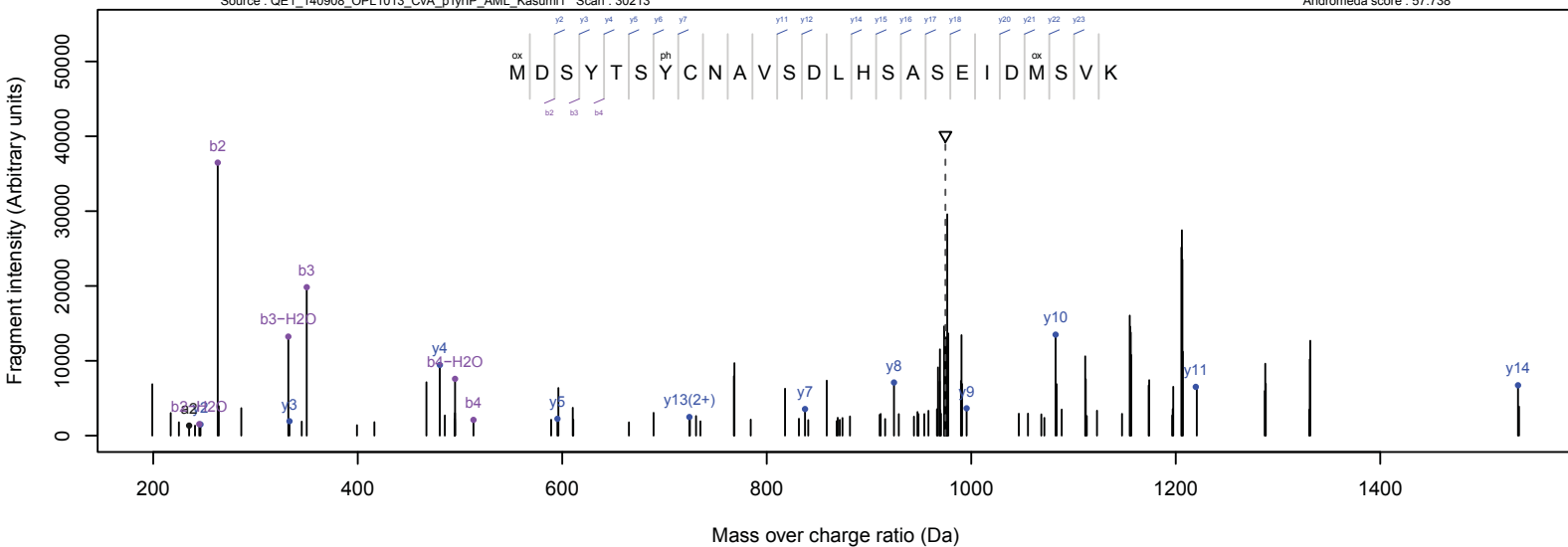




## M(ox)DSYTSY(ph)CNAVSDLHSASEIDMSVK-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan: 30213

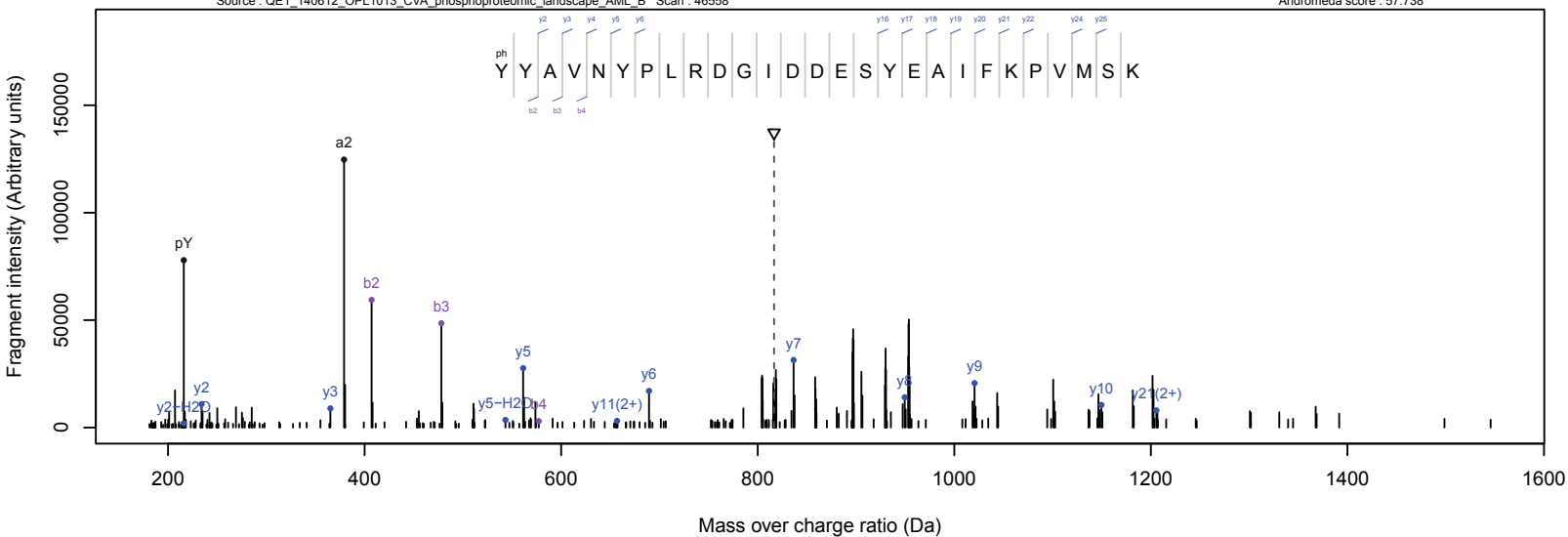
Andromeda score: 57.738



## Y(ph)YAVNYPLRDGIDDES YEAI FKP VMSK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 46558

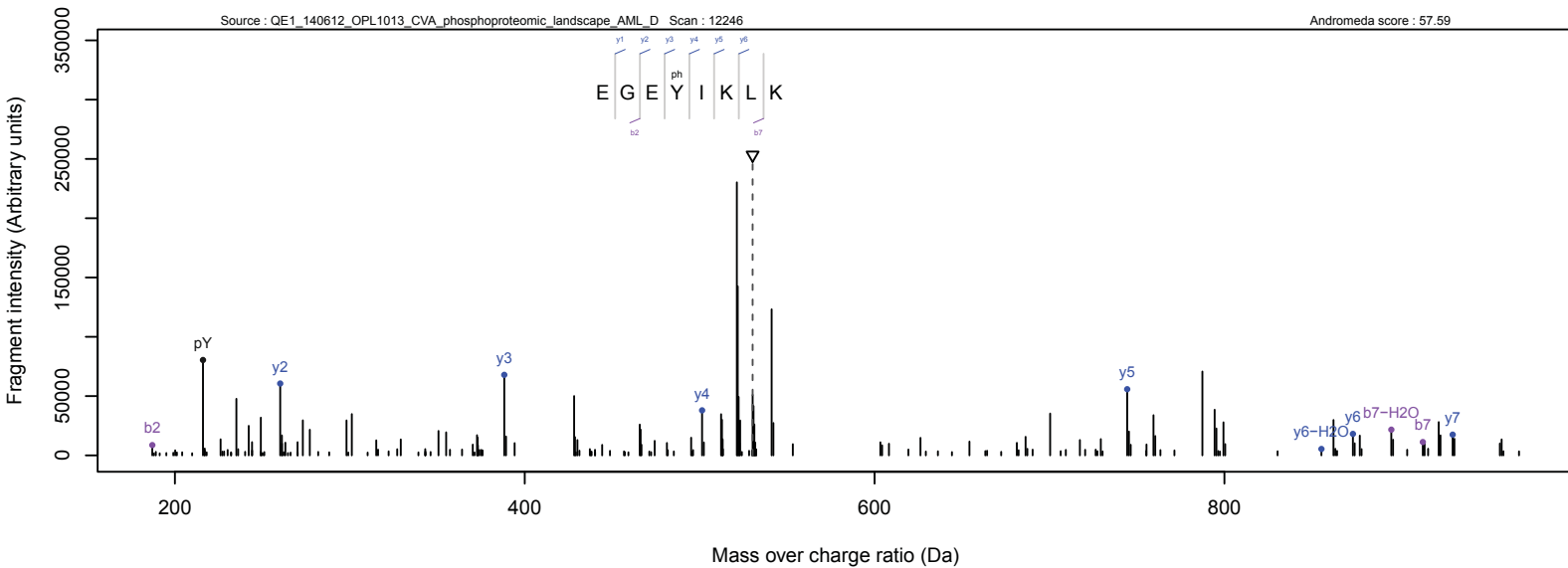
Andromeda score: 57.738



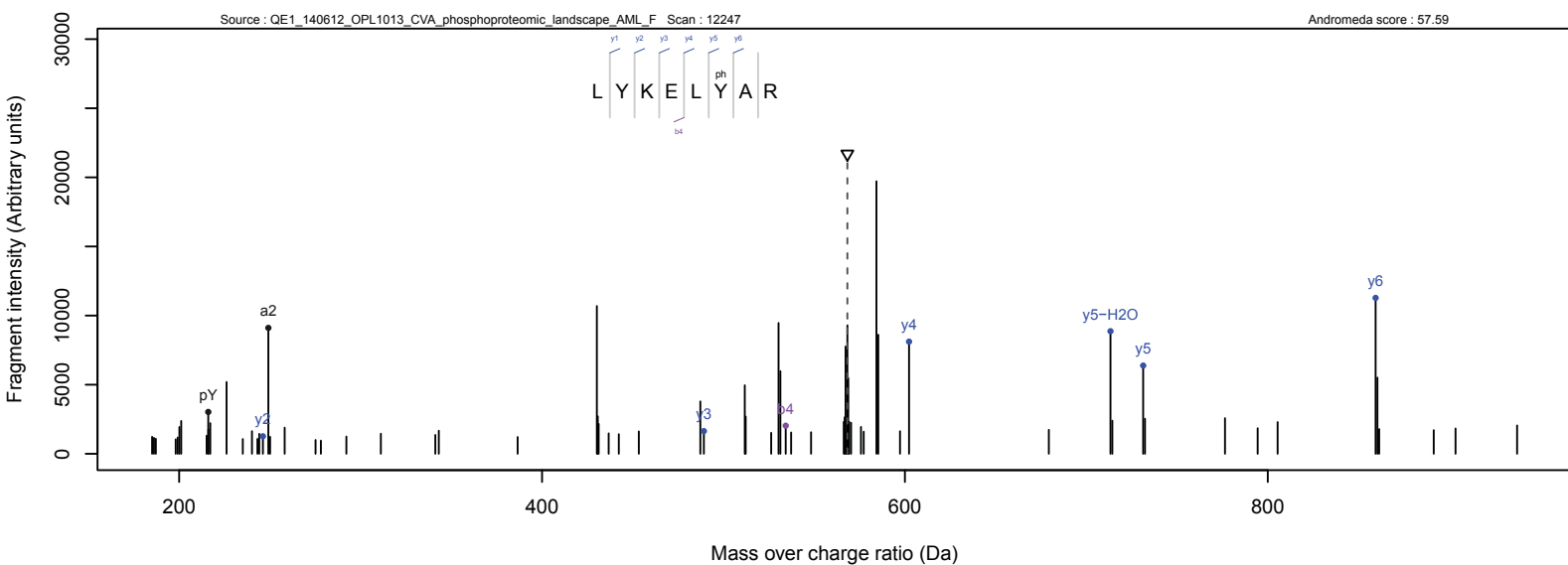
## EGEY(ph)IKLK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 12246

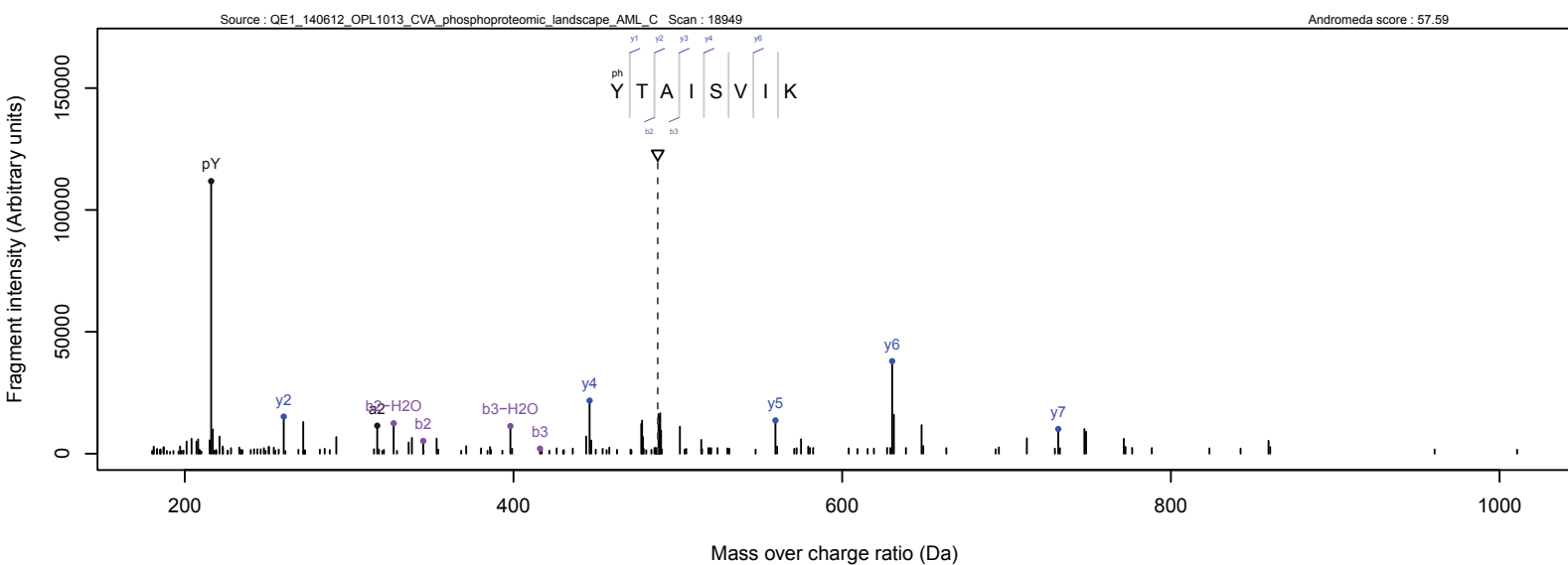
Andromeda score: 57.59



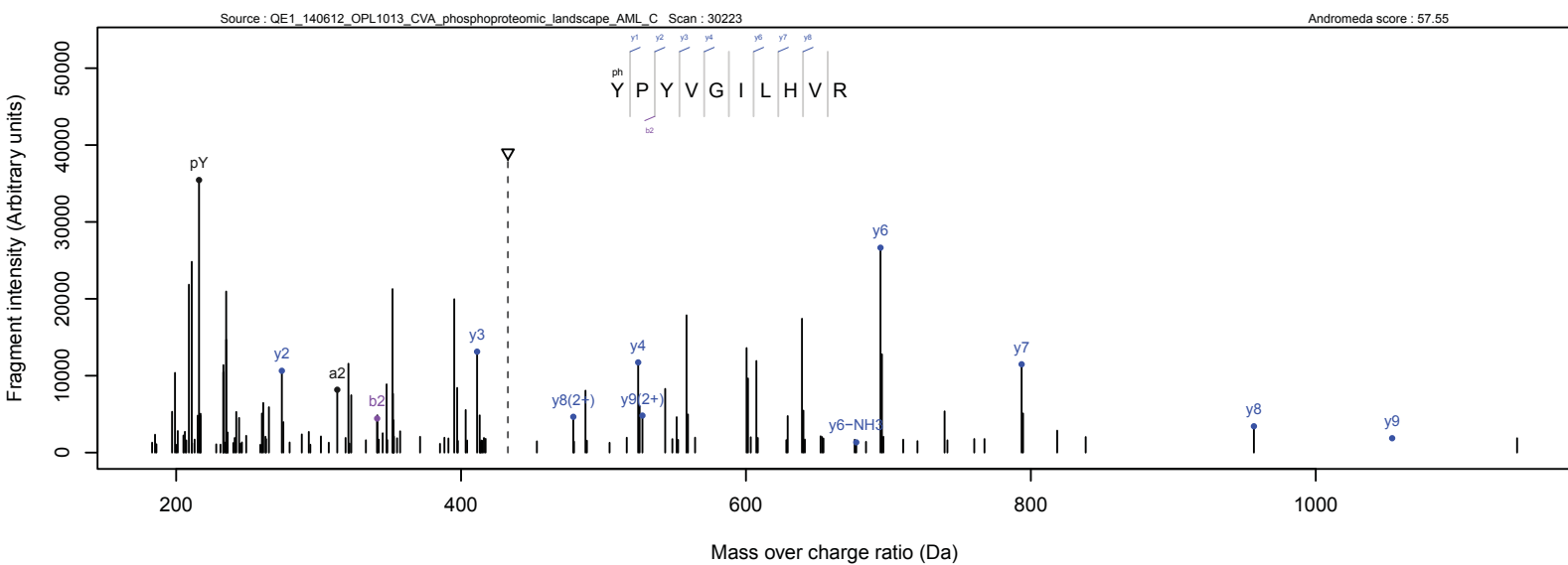
## LYKELY(ph)AR-2+



## Y(ph)TAISVIK-2+



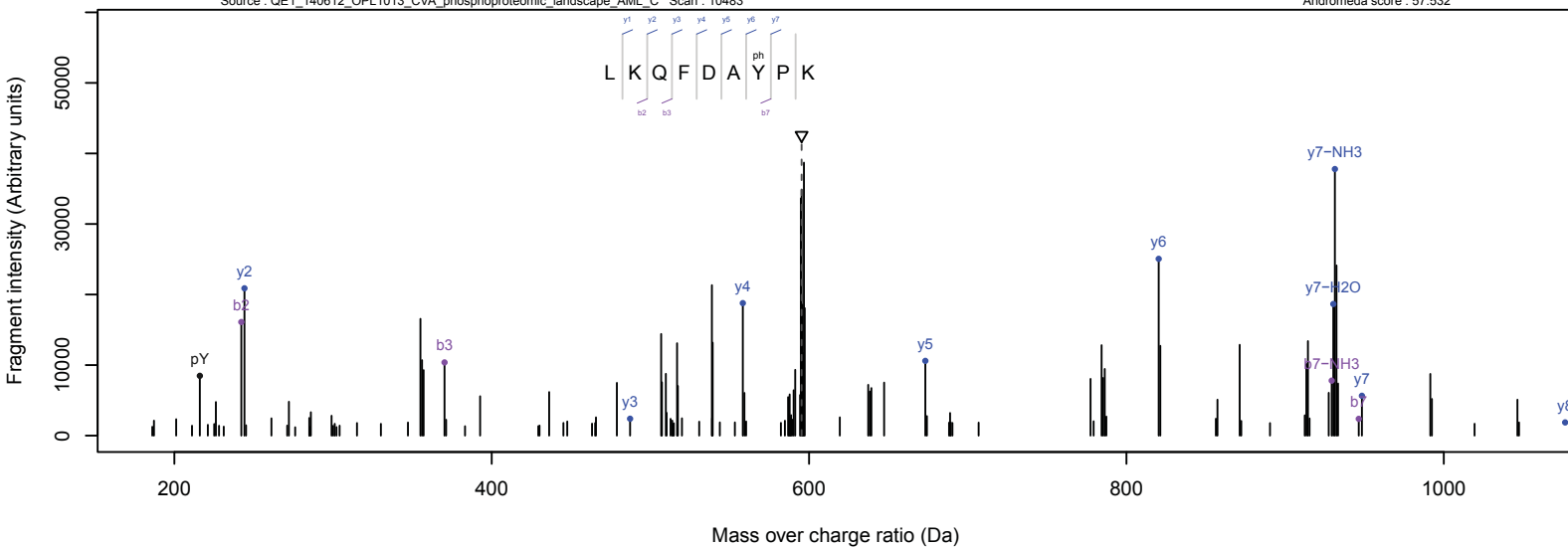
## Y(ph)PYVGILHVR-3+



## LKQFDAY(ph)PK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10483

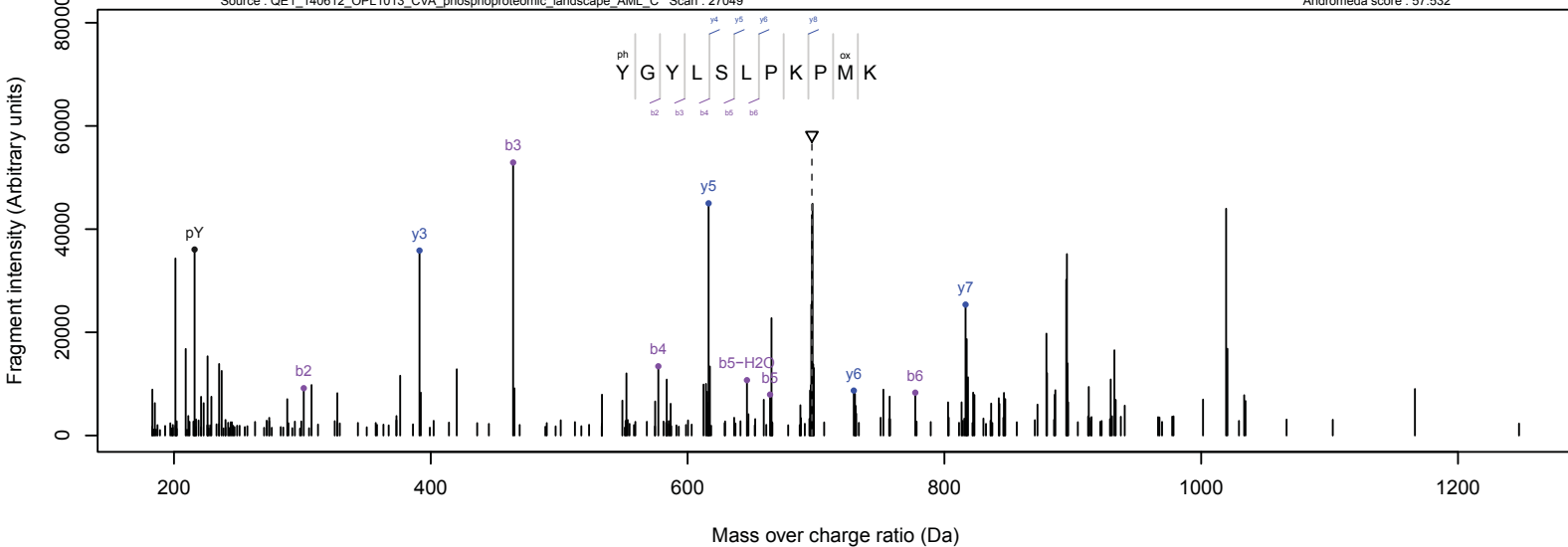
Andromeda score : 57.532



## Y(ph)GYLSLPKM(ox)K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27049

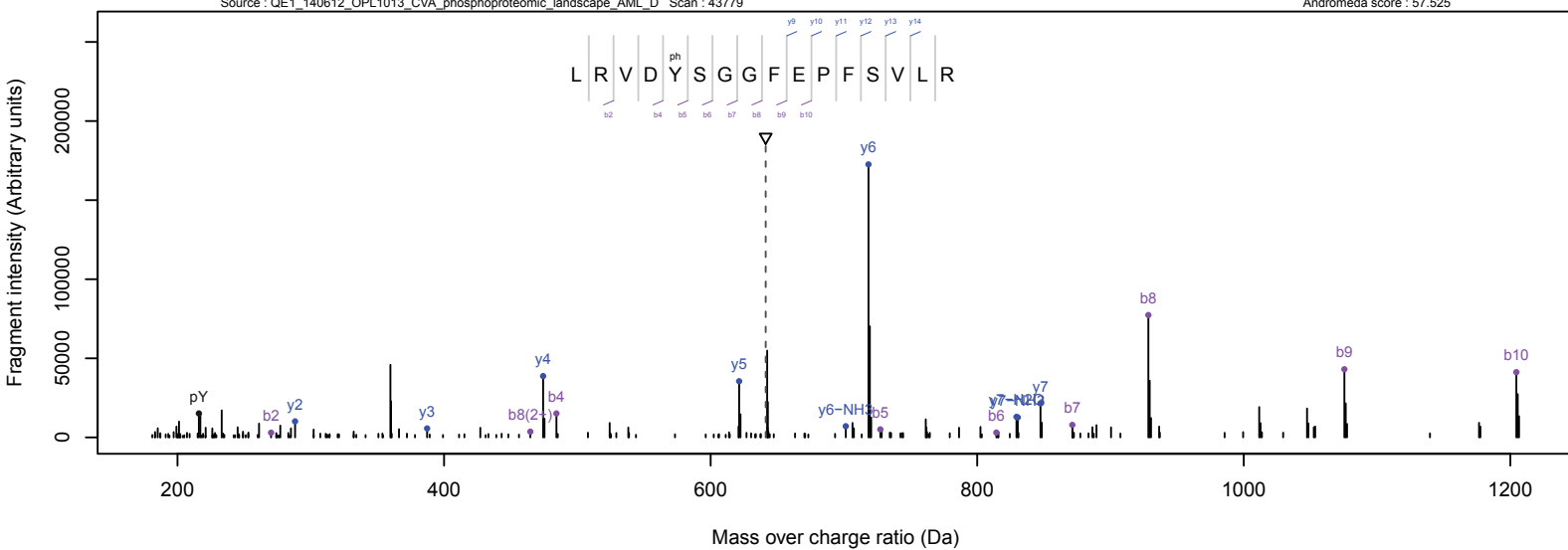
Andromeda score : 57.532



## LRVDY(ph)SGGFEPFSLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 43779

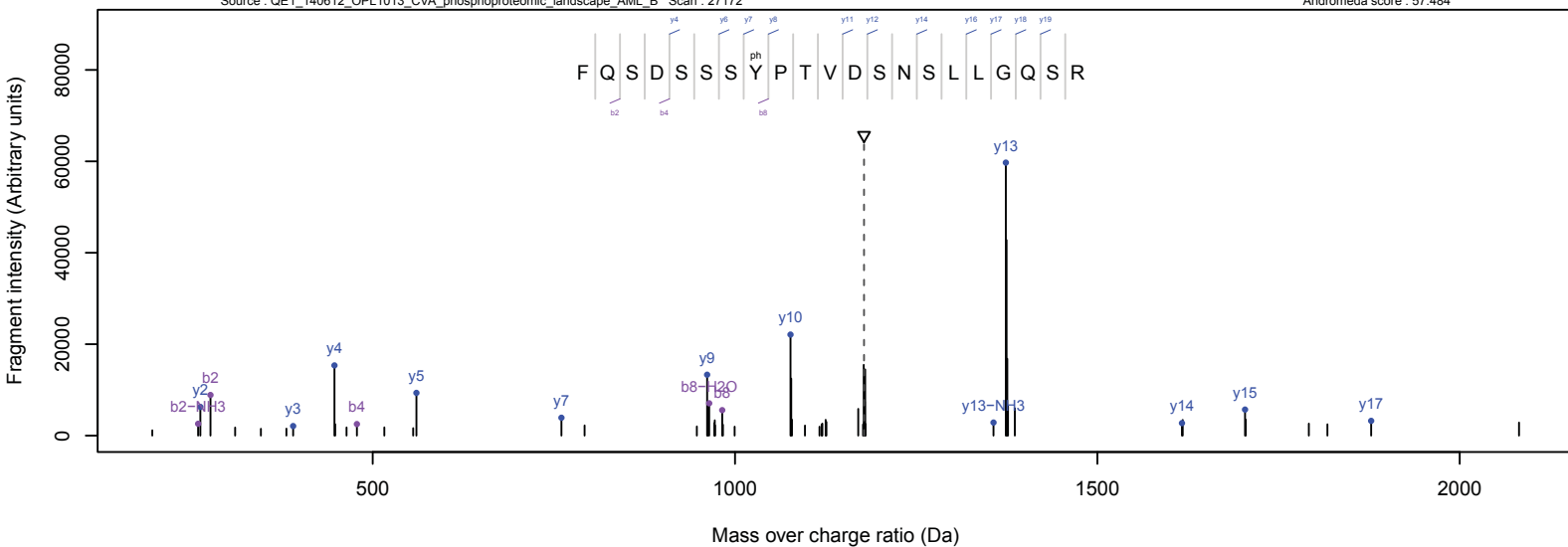
Andromeda score : 57.525



## FQDSSSY(ph)PTVDSNLLGQSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27172

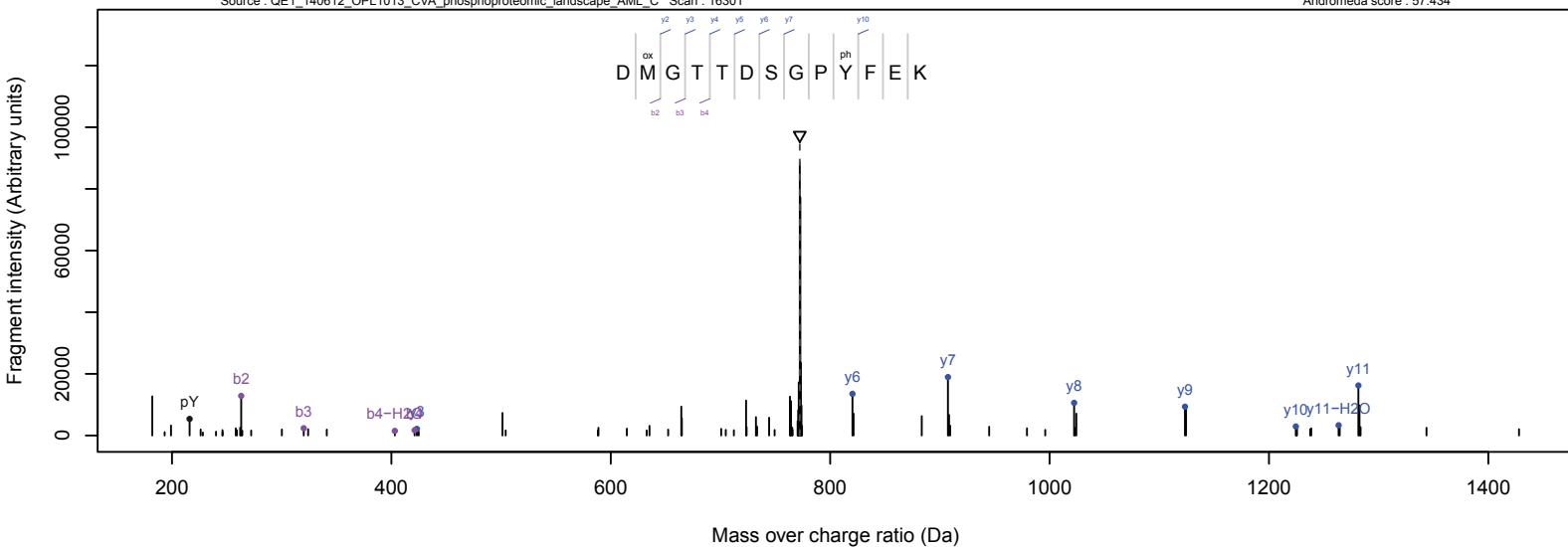
Andromeda score : 57.484



## DM(ox)GTTDSGPY(ph)FEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16301

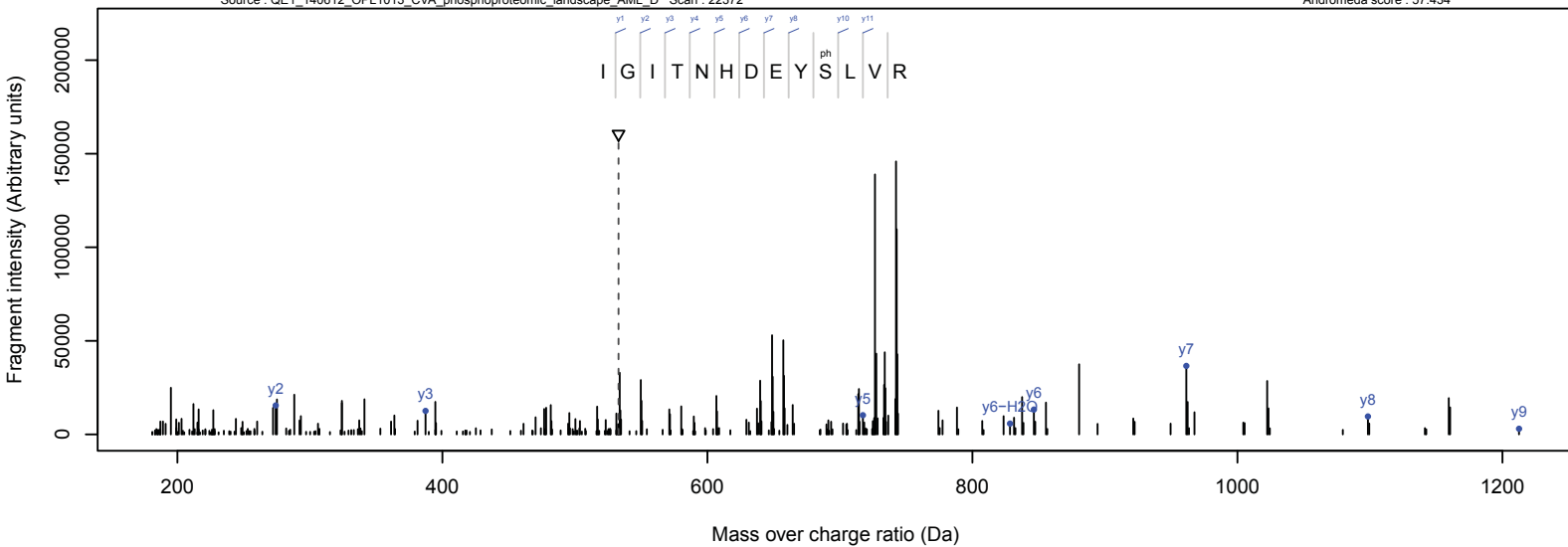
Andromeda score : 57.434



## IGITNHDEYS(ph)LVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22372

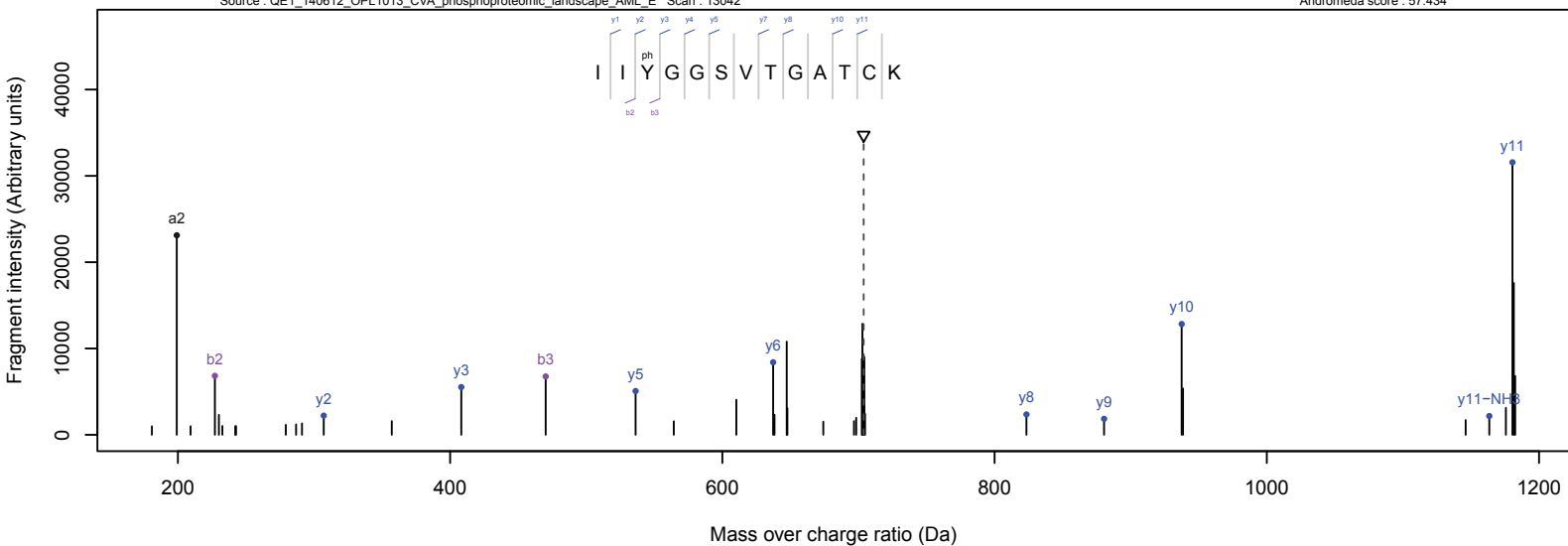
Andromeda score : 57.434



## IIY(ph)GGSVTGATCK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 13042

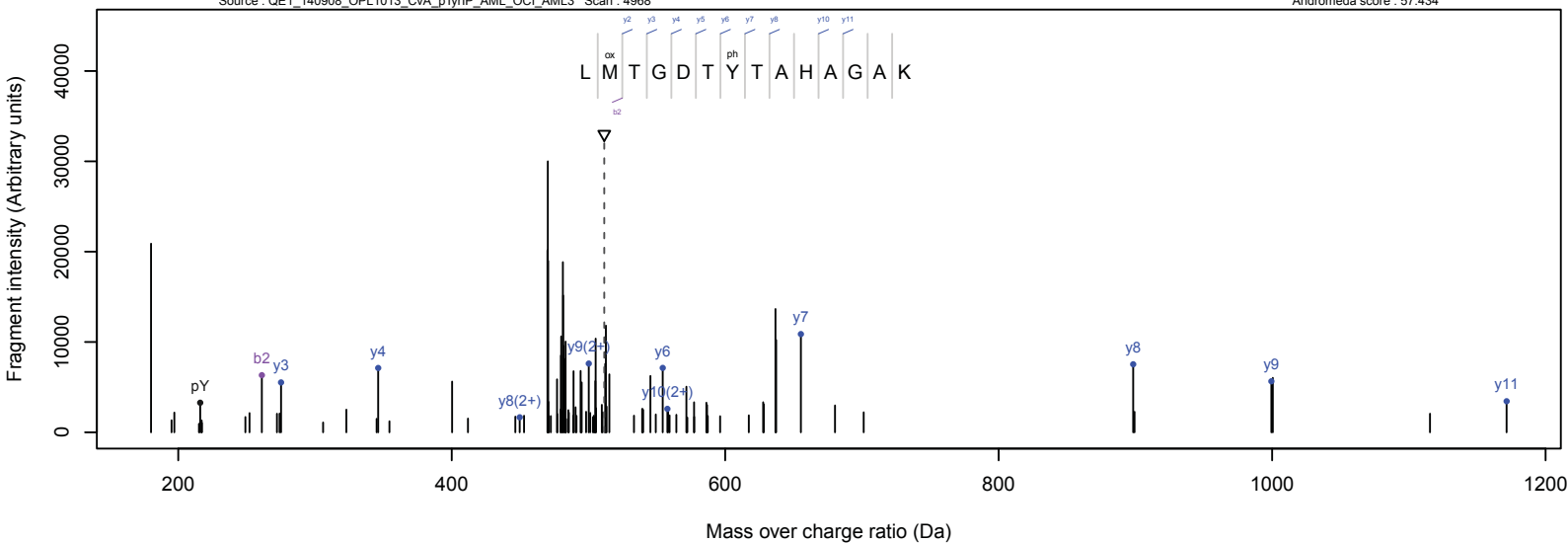
Andromeda score : 57.434



## LM(ox)TGDTY(ph)TAHAGAK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 4968

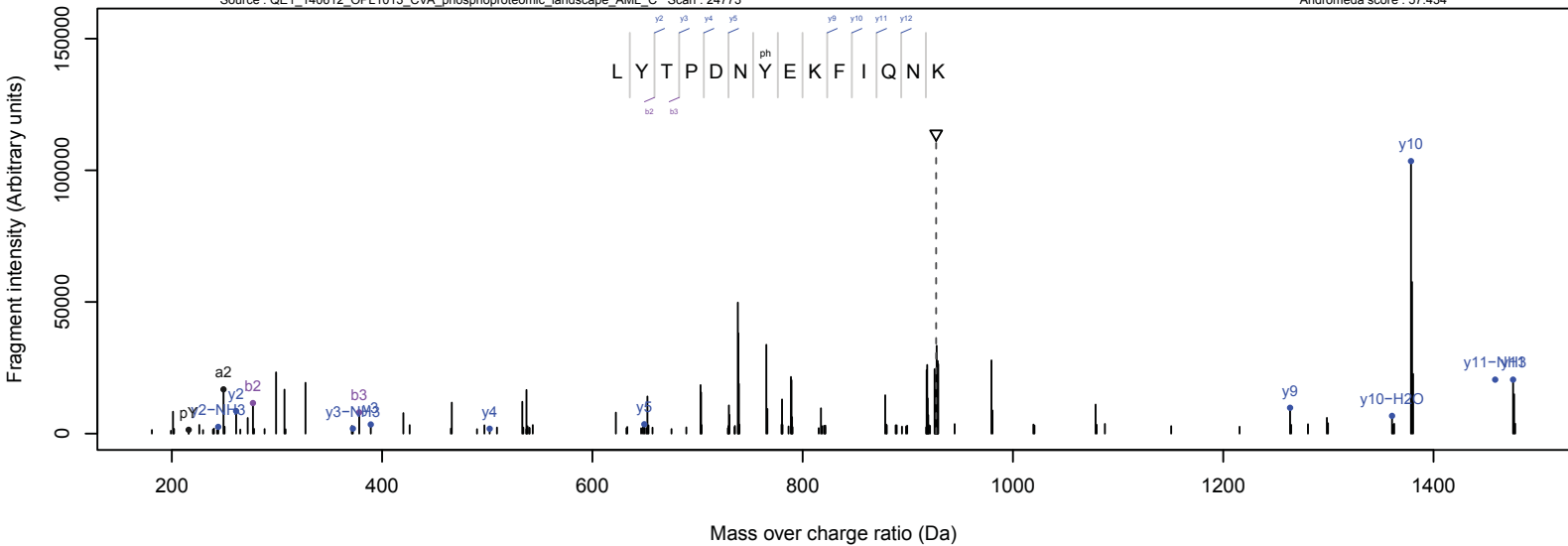
Andromeda score : 57.434



## LYTPDNY(ph)EKFIQNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24773

Andromeda score : 57.434

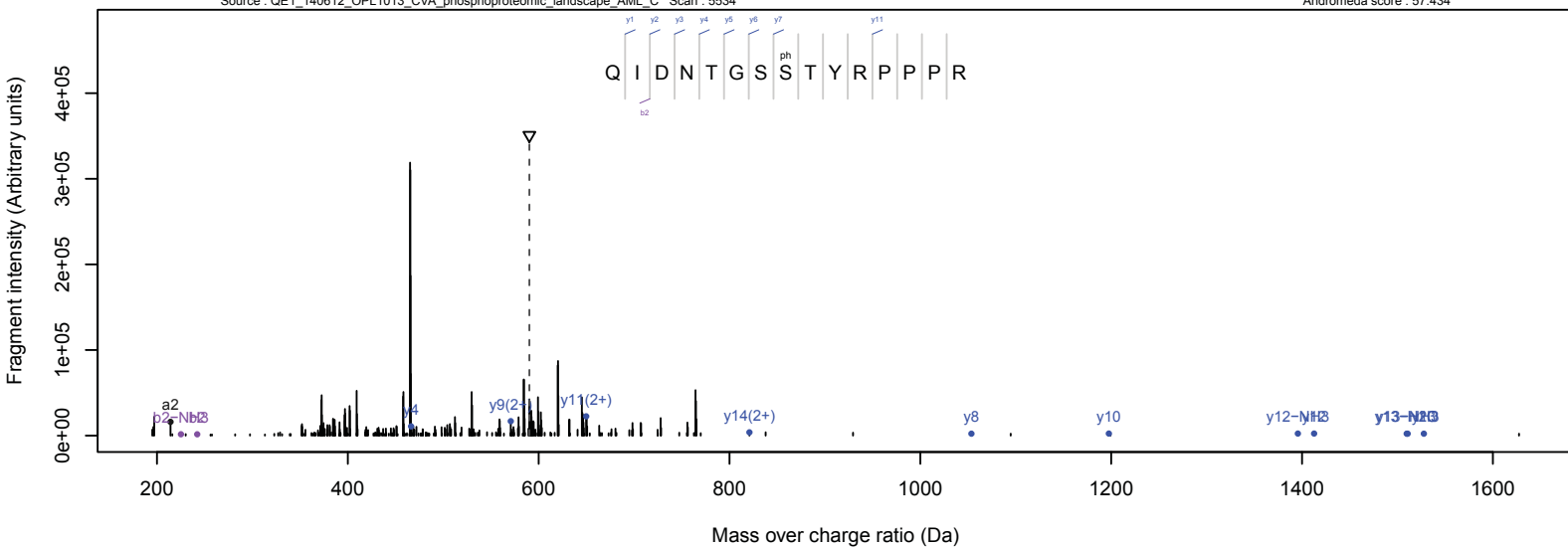




## QIDNTGSS(ph)TYRPPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 5534

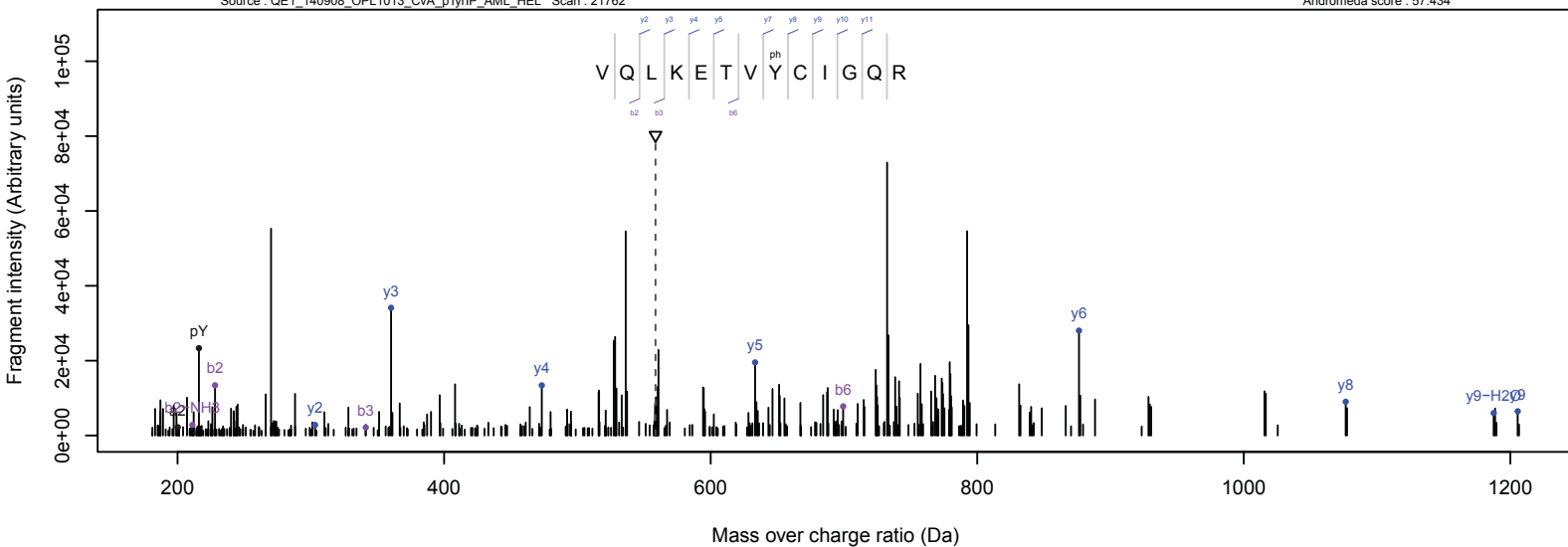
Andromeda score : 57.434



## VLKETVY(ph)CIGQR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 21762

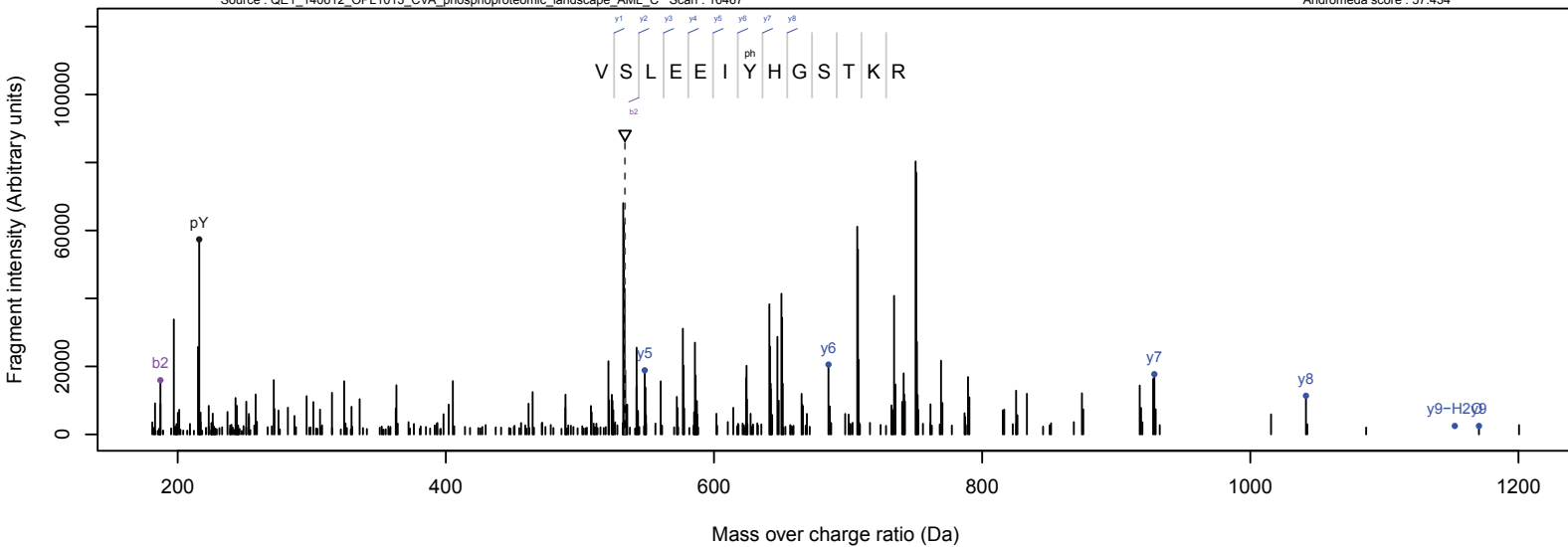
Andromeda score : 57.434



## VSLEEIY(ph)HGSTKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16467

Andromeda score : 57.434

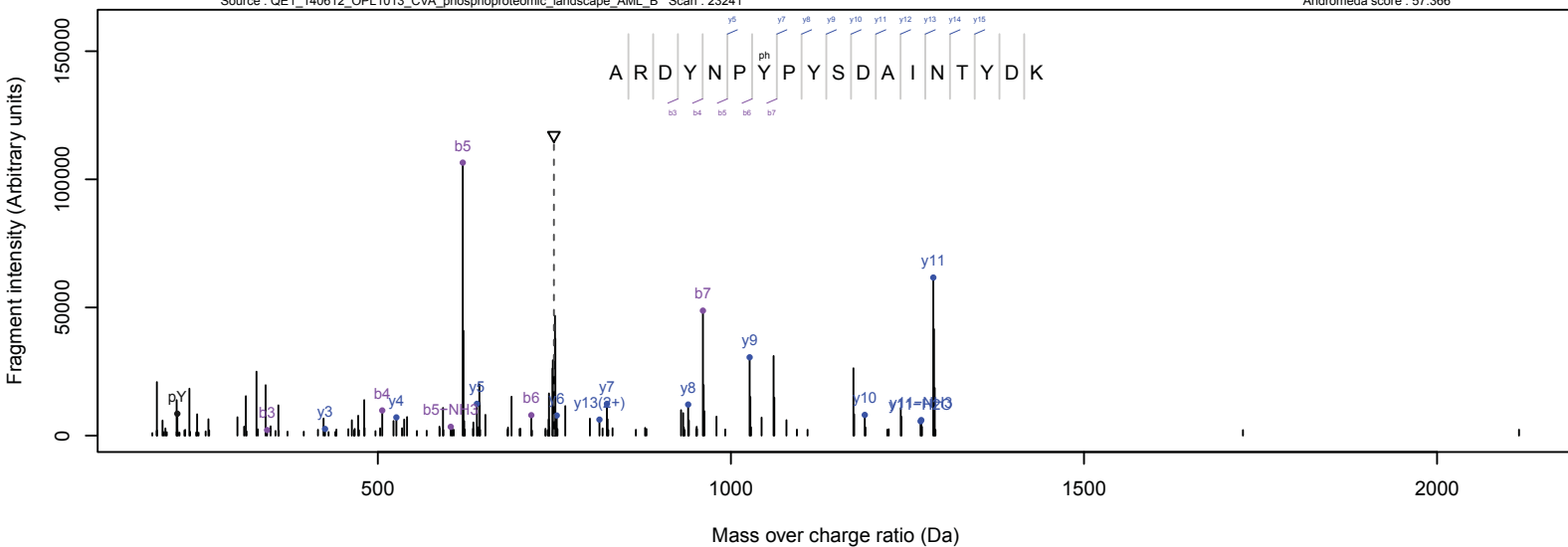




## ARDYNPY(ph)PYSDAINTYDK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23241

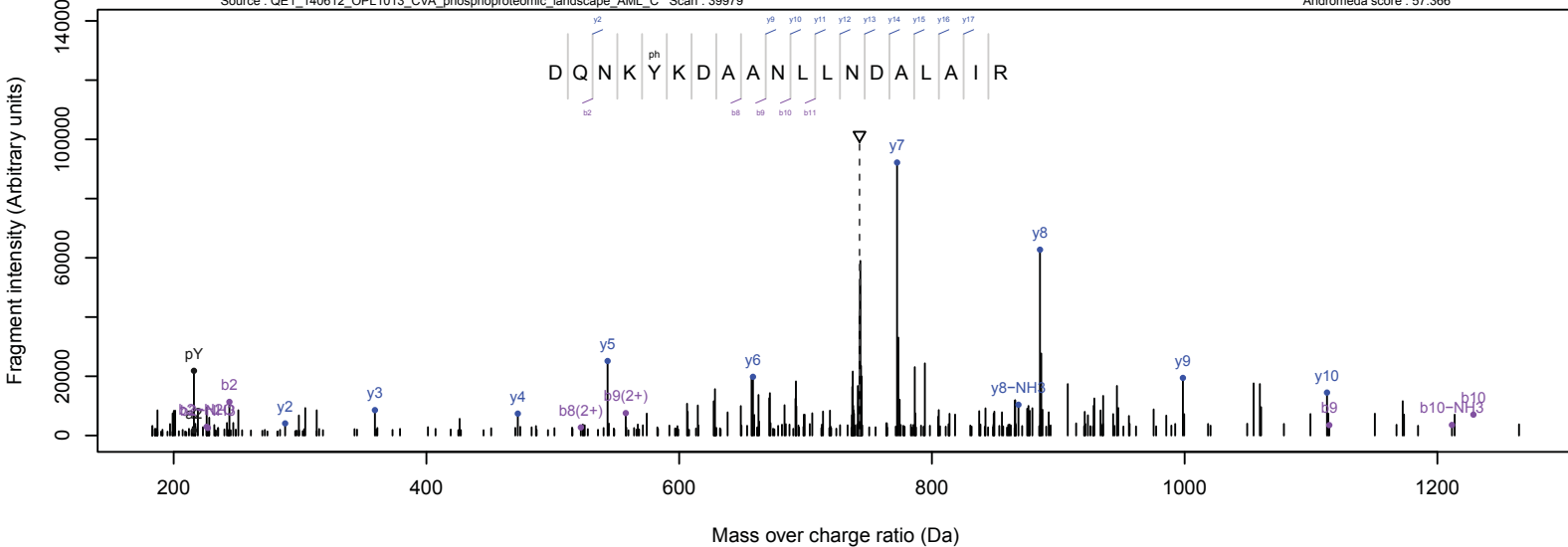
Andromeda score : 57.366



## DQNKY(ph)KDAANLLNDALAIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39979

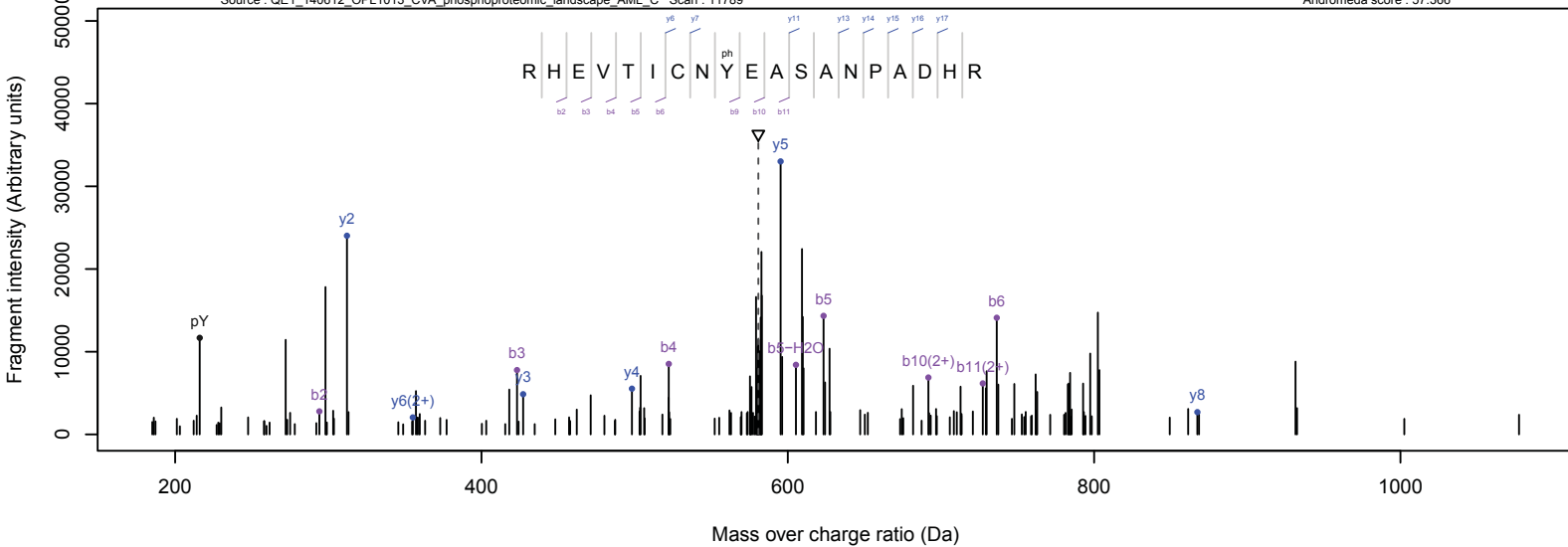
Andromeda score : 57.366



## RHEVTICNY(ph)EASNPADHR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11789

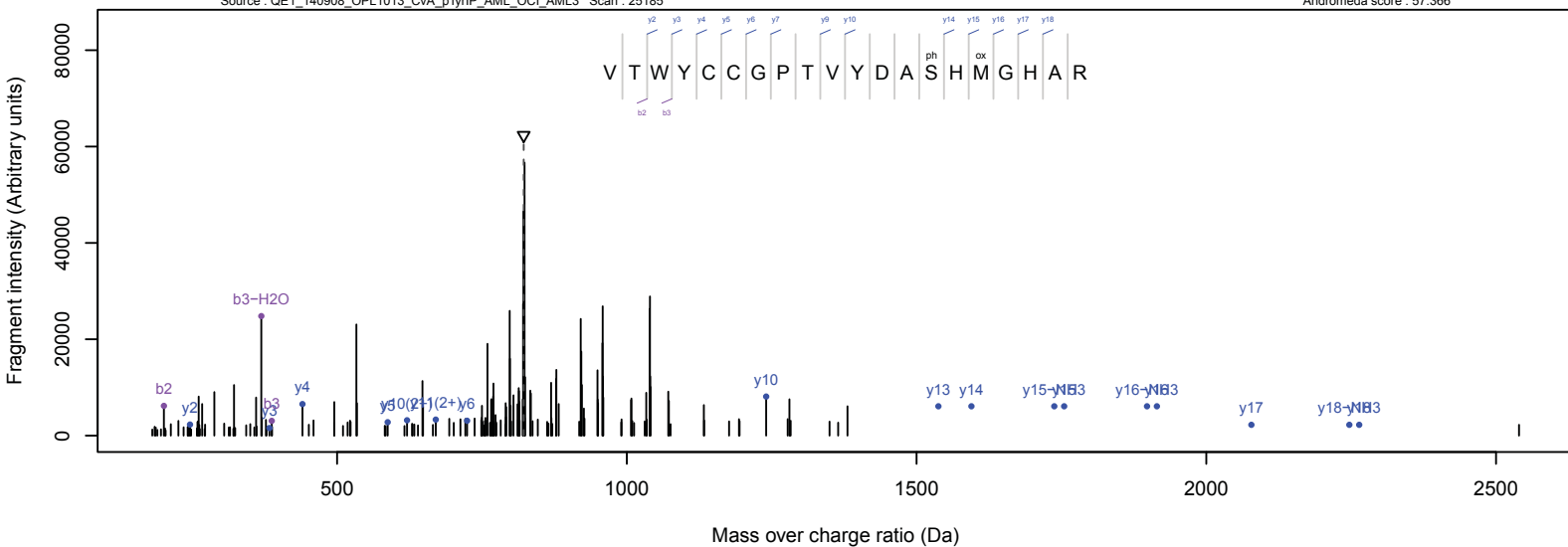
Andromeda score : 57.366



VTWYCCGPTVYDAS(ph)HM(ox)GHAR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 25185

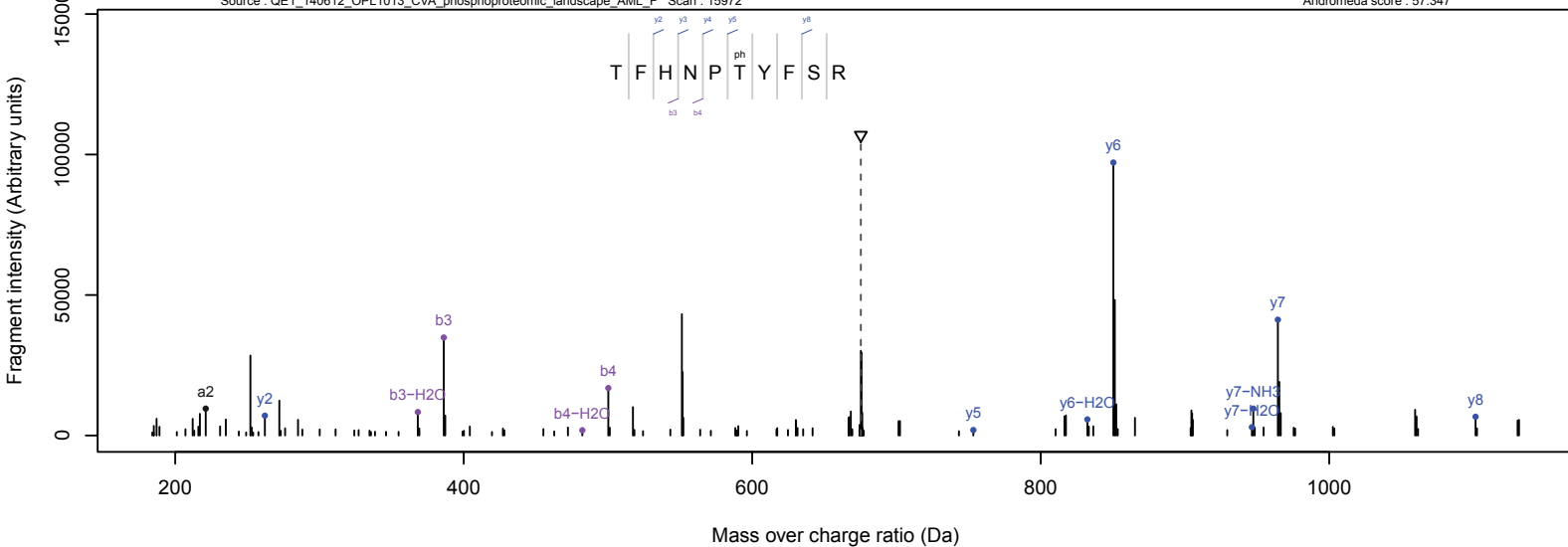
Andromeda score : 57.366



TFHNPT(ph)YFSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 15972

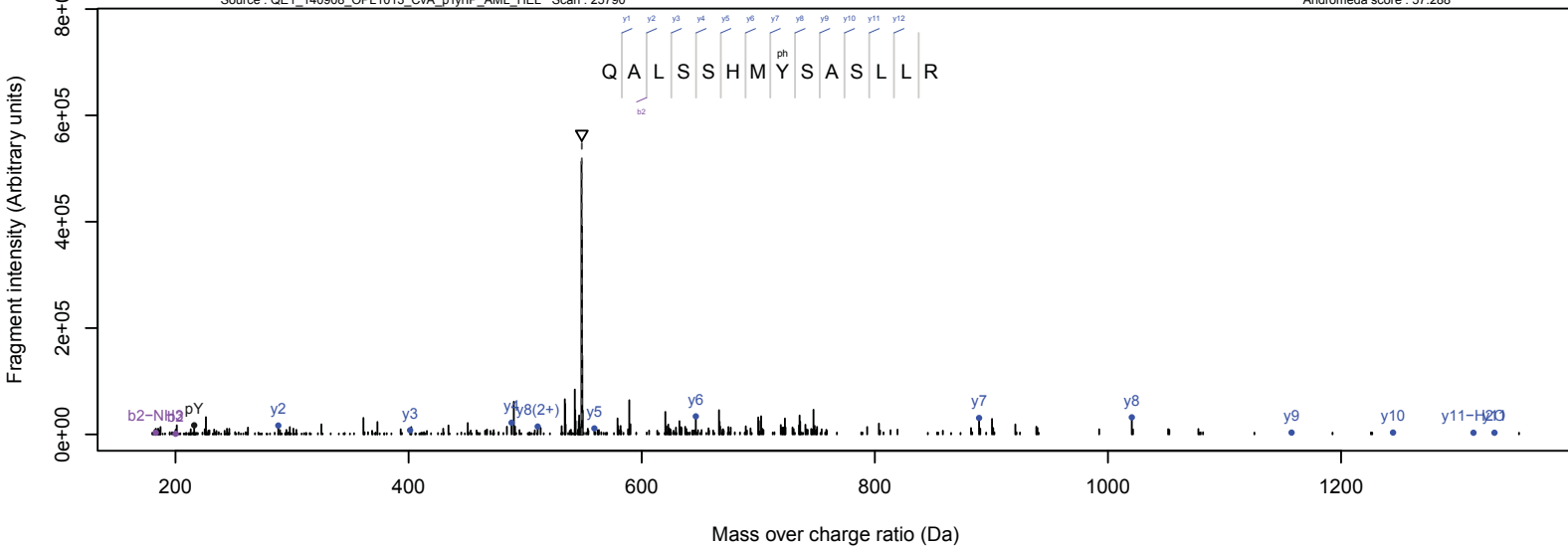
Andromeda score : 57.347



QALSSHMY(ph)SASLLR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 25790

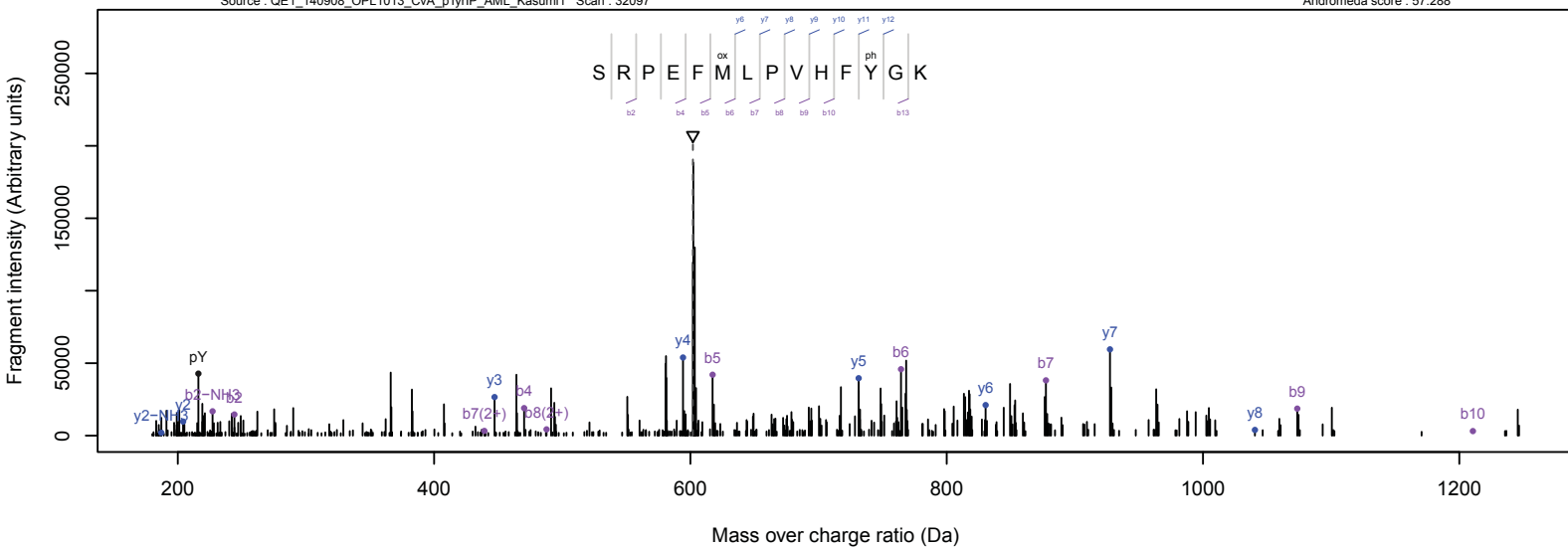
Andromeda score : 57.288



SRPEFM(ox)LPVHFY(ph)GK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 32097

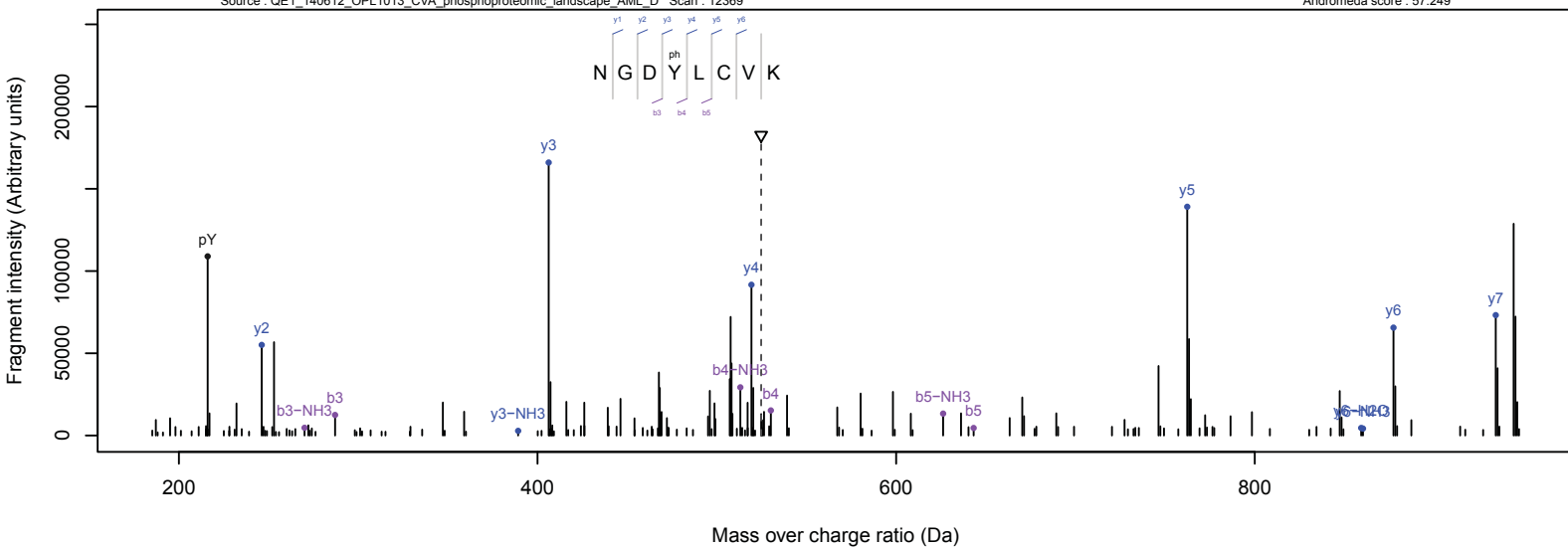
Andromeda score : 57.288



NGDY(ph)LCVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12369

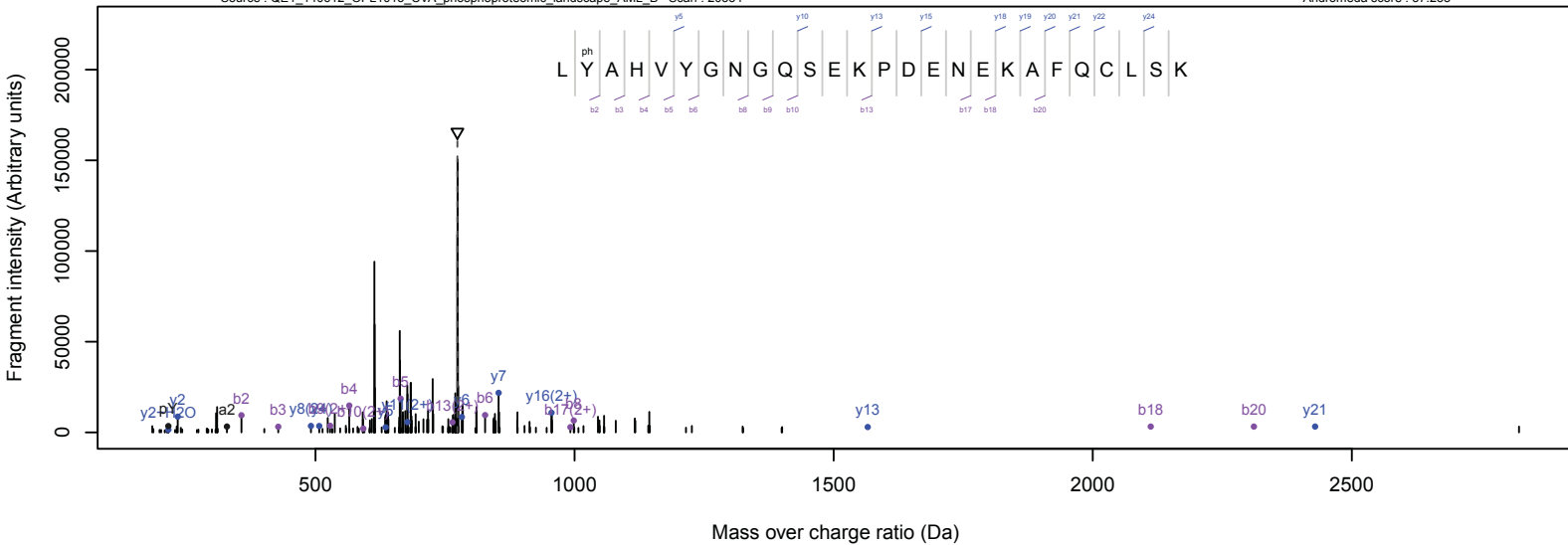
Andromeda score : 57.249



LY(ph)AHVYGNQSEKPDENEKAFQCLSK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20334

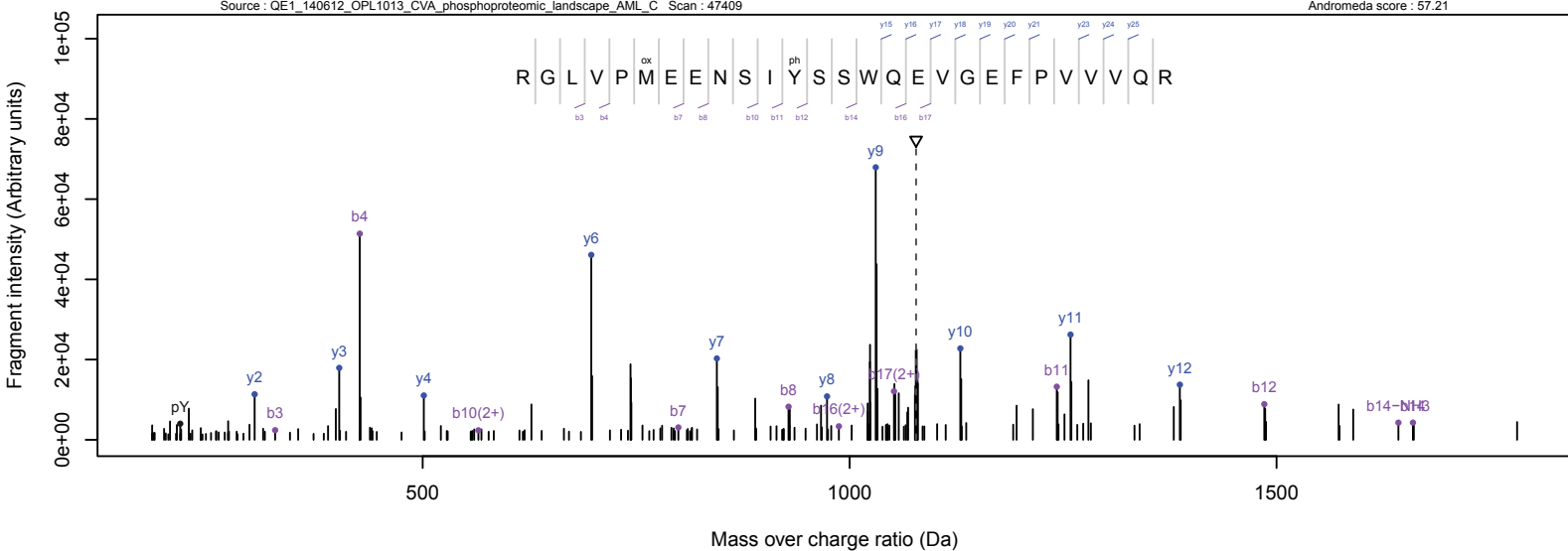
Andromeda score : 57.238



RGLVPM(ox)EENSIY(ph)SSWQEVGEFVVVQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47409

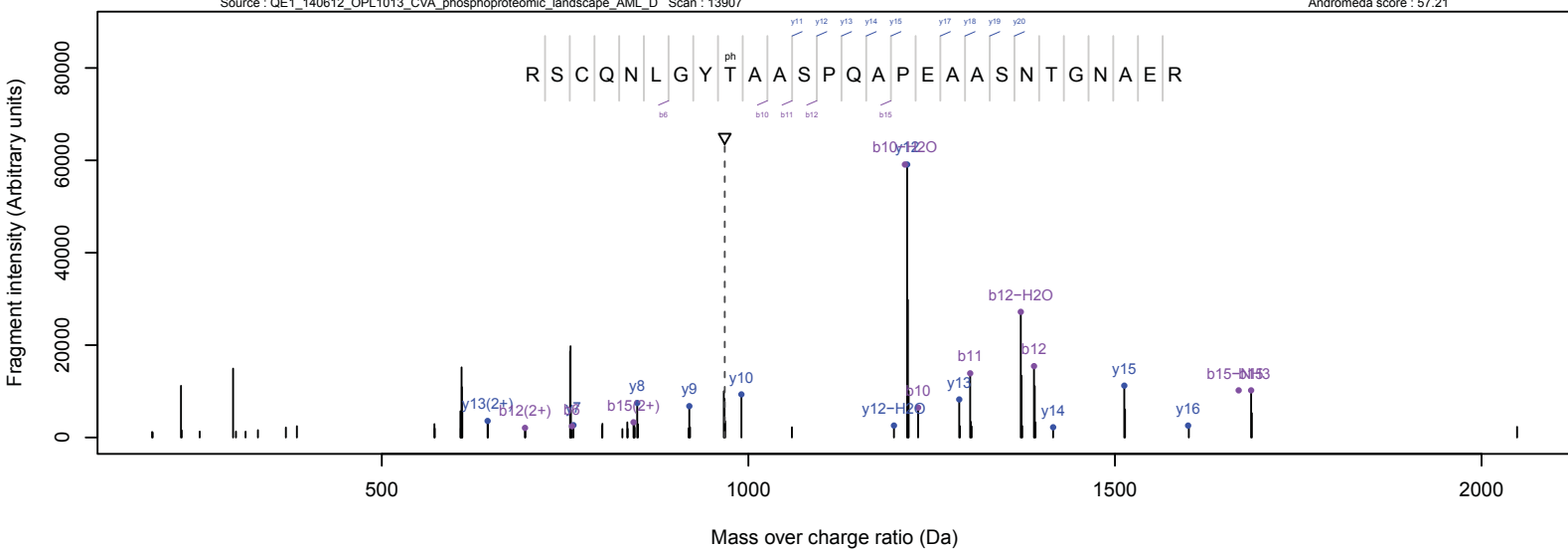
Andromeda score : 57.21



RSCQNLGYT(ph)AASPQAPEAASNTGNAER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 13907

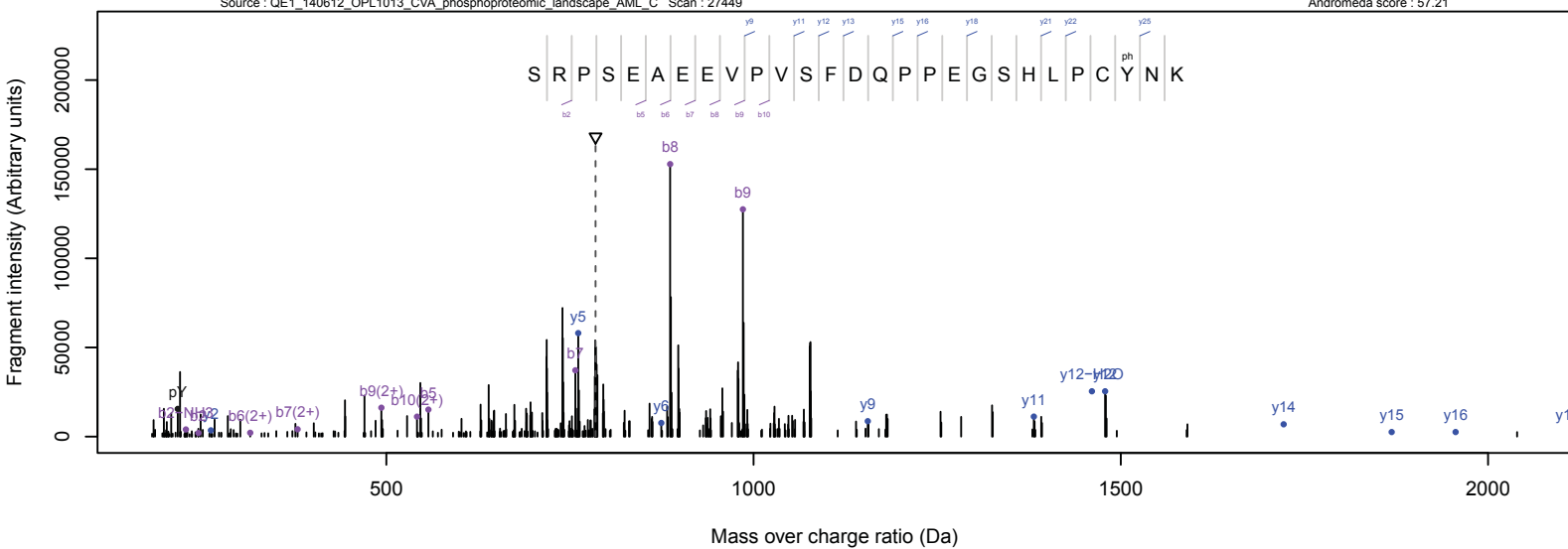
Andromeda score : 57.21



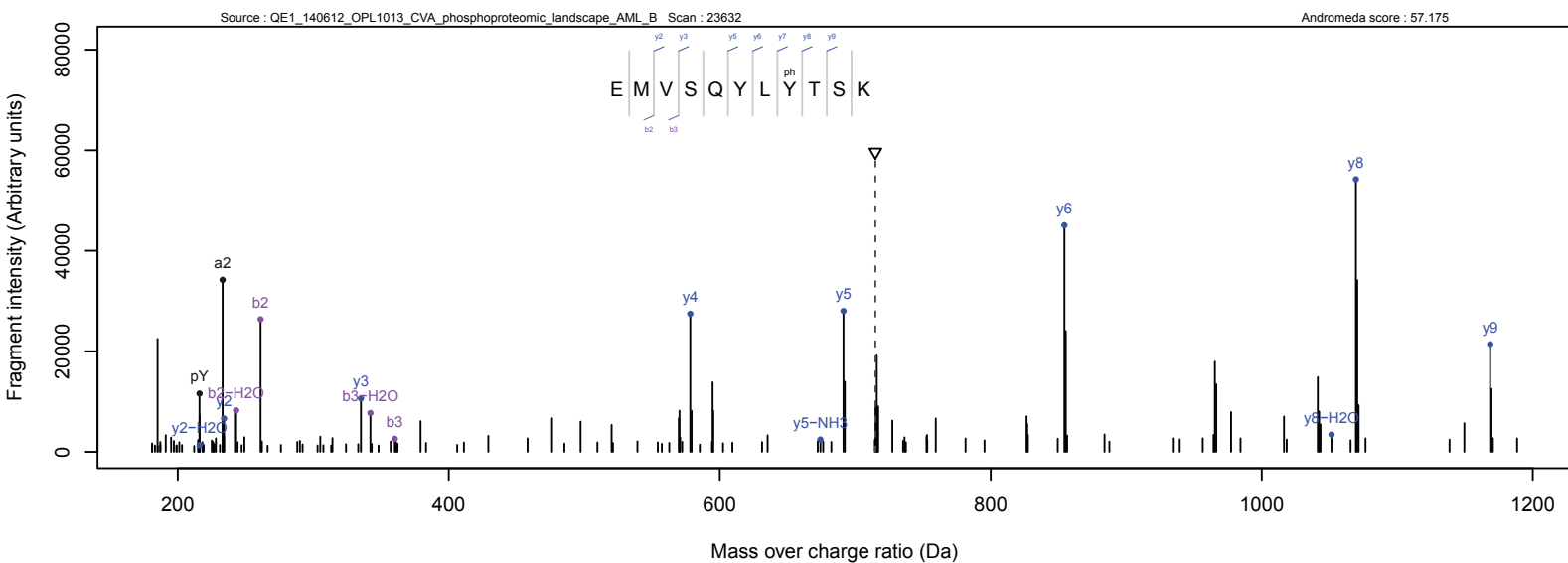
SRPSEAEVPSFDQPPEGSHLPCY(ph)NK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27449

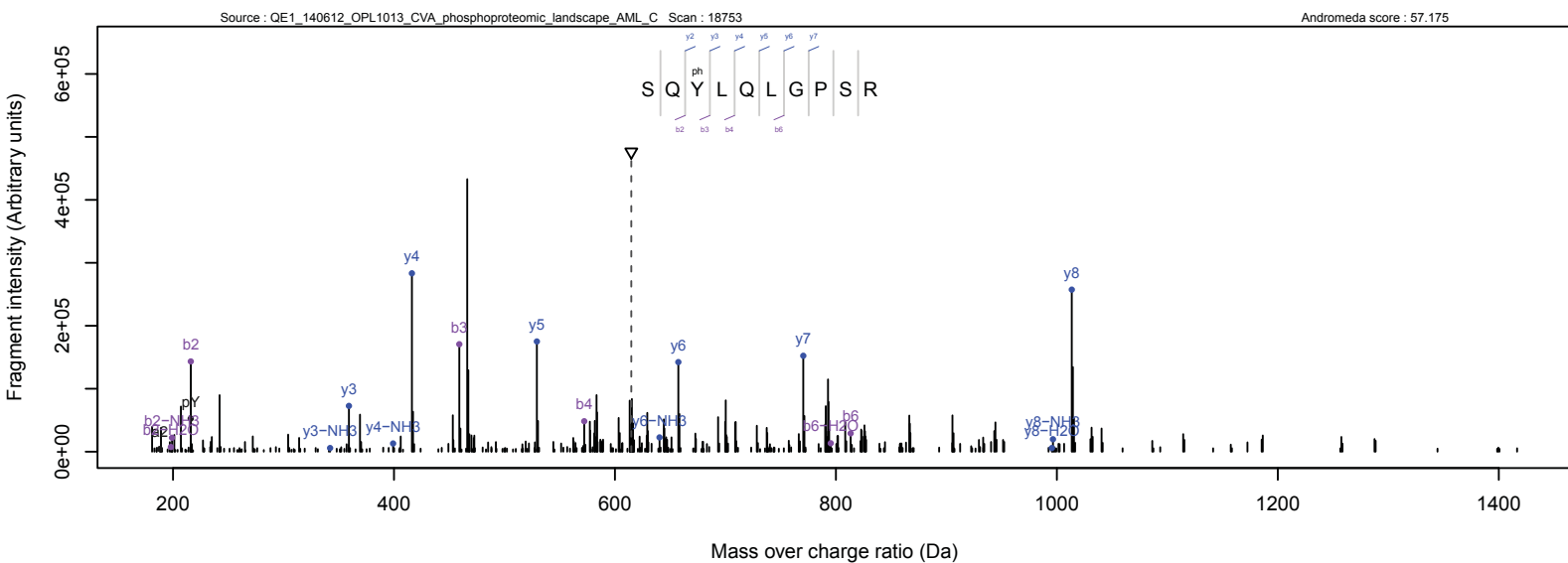
Andromeda score : 57.21



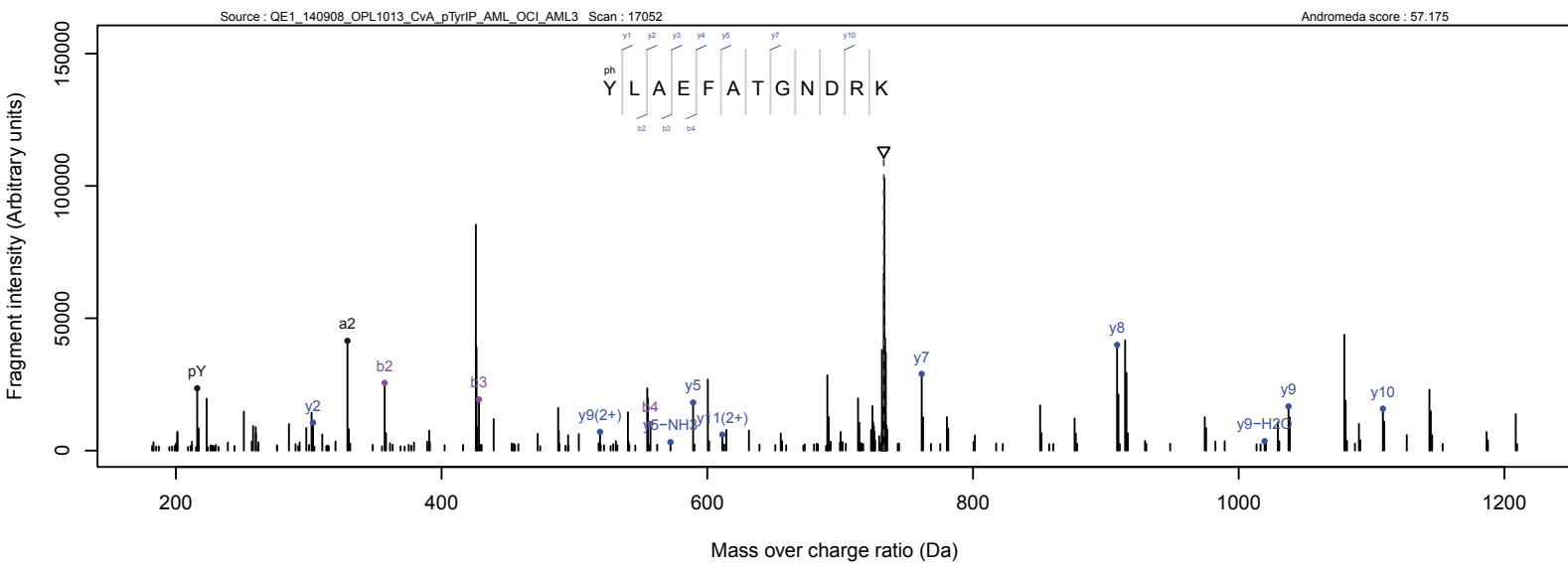
EMVSQYLY(ph)TSK-2+



SQY(ph)LQLGPSR-2+



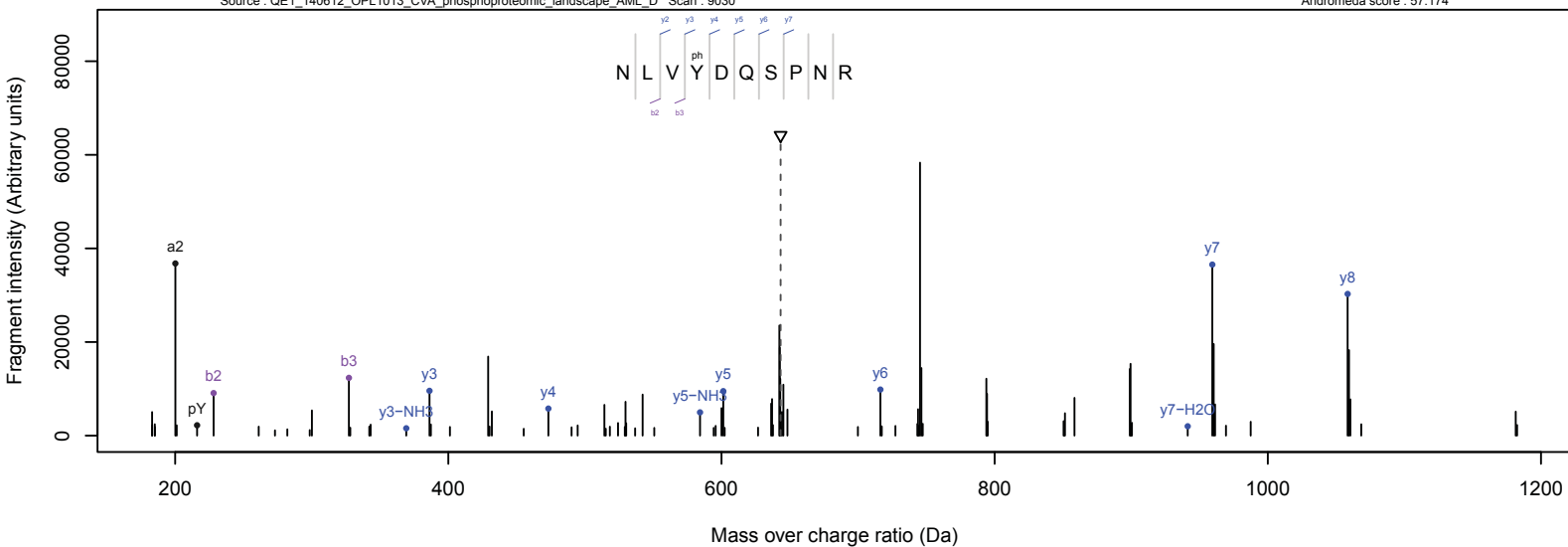
Y(ph)LAEFATGNDRK-2+



## NLVY(ph)DQSPNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9030

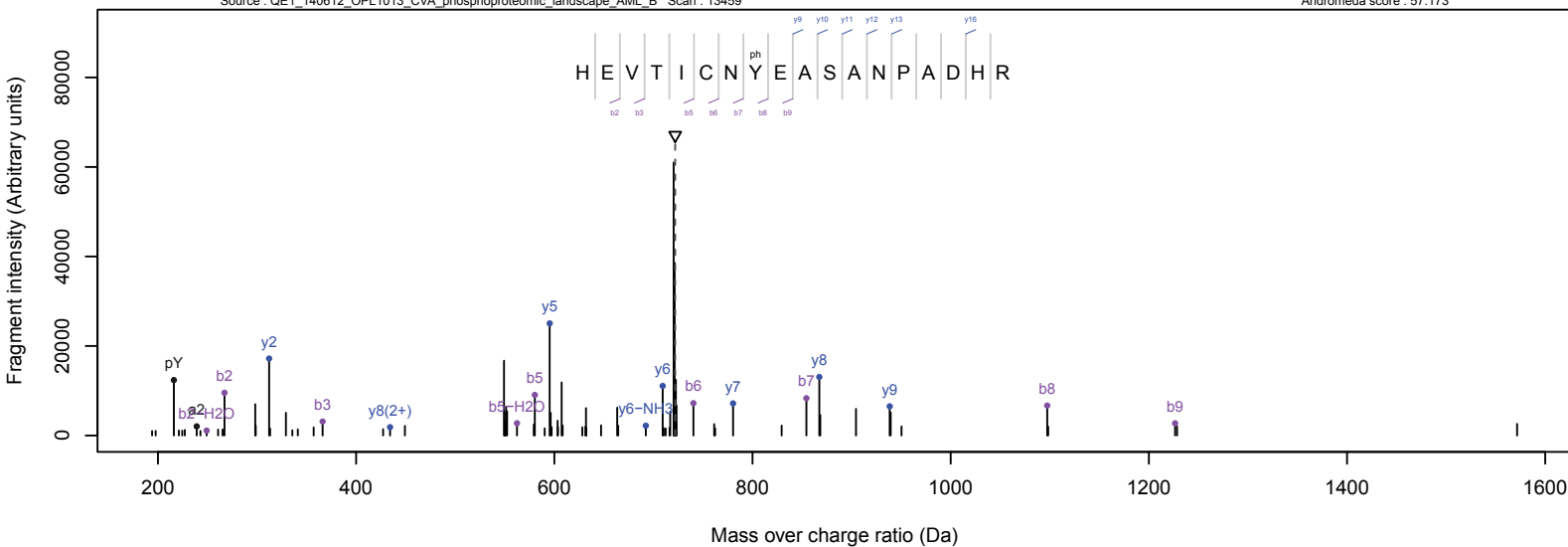
Andromeda score : 57.174



## HEVTICNY(ph)EASANPADHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13459

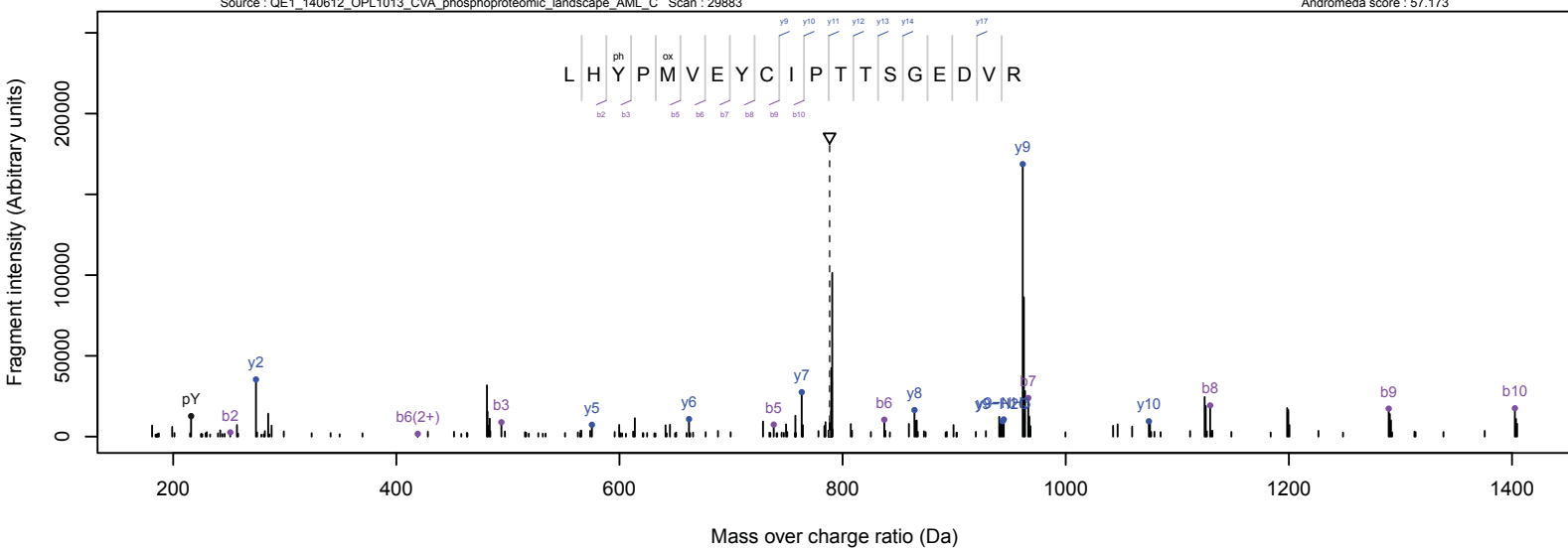
Andromeda score : 57.173



## LHY(ph)PM(ox)VEYCIPTTSGEDVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29883

Andromeda score : 57.173

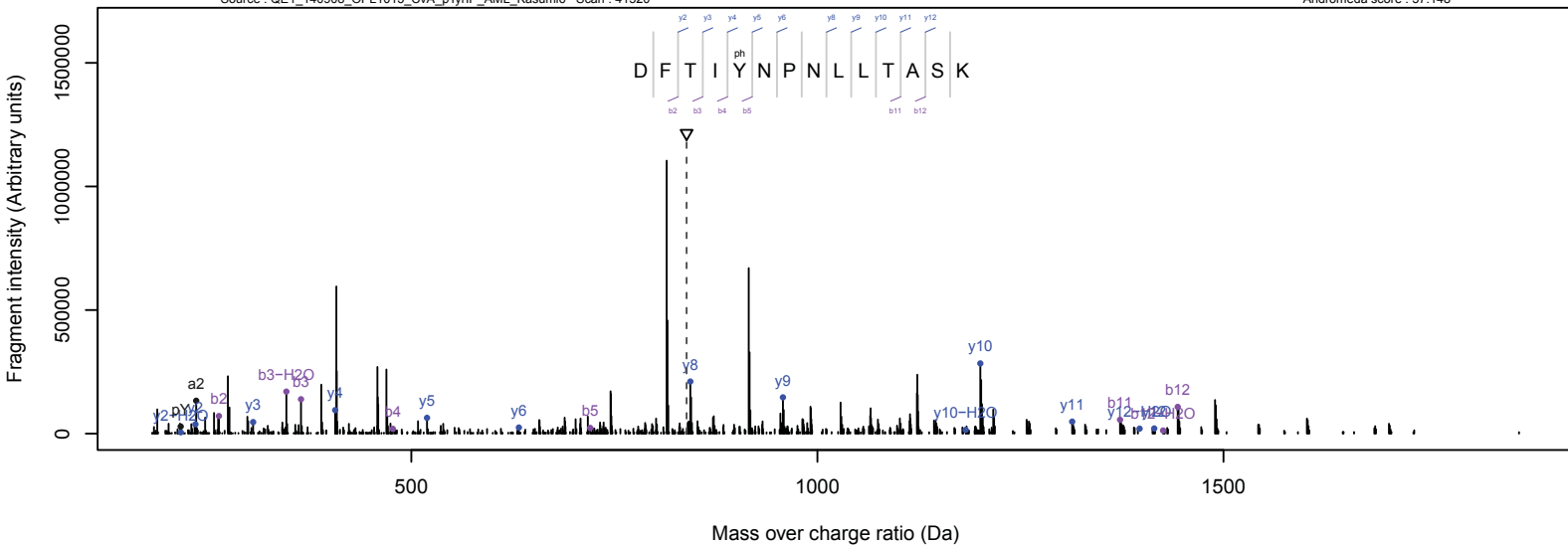




DFTIY(ph)NPLLTASK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 41520

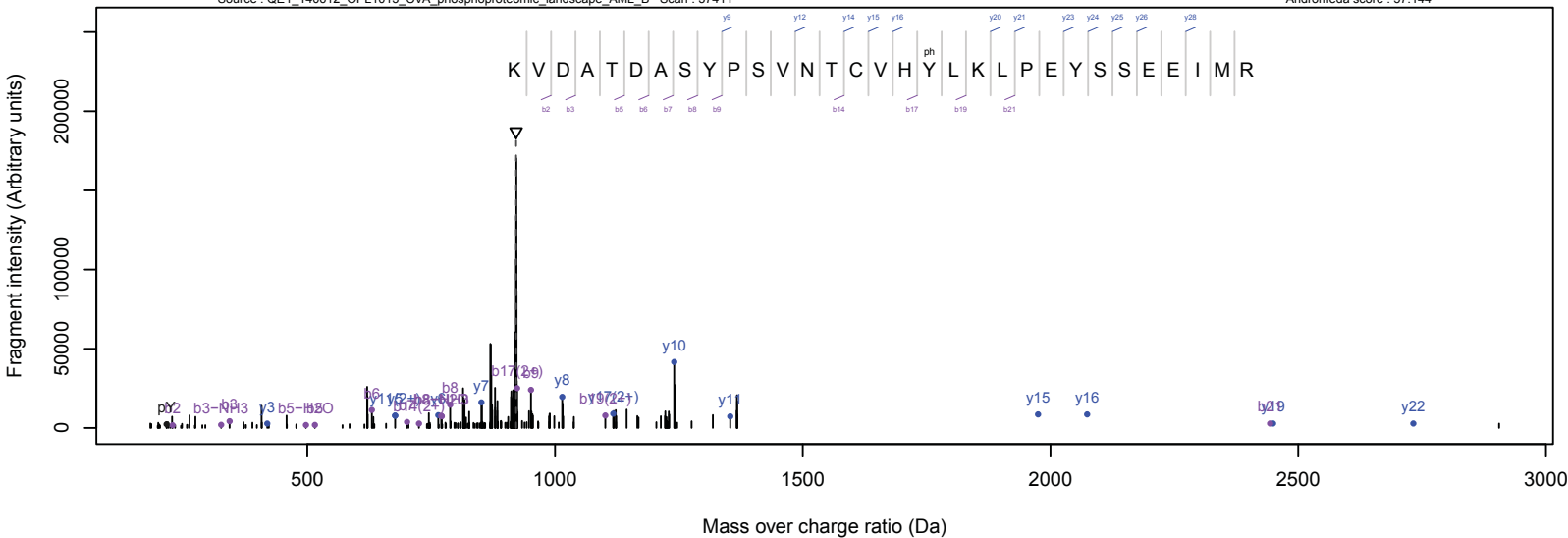
Andromeda score: 57.148



KVVDATDASYPVNTCVHY(ph)LKLPESSEEIMR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 37411

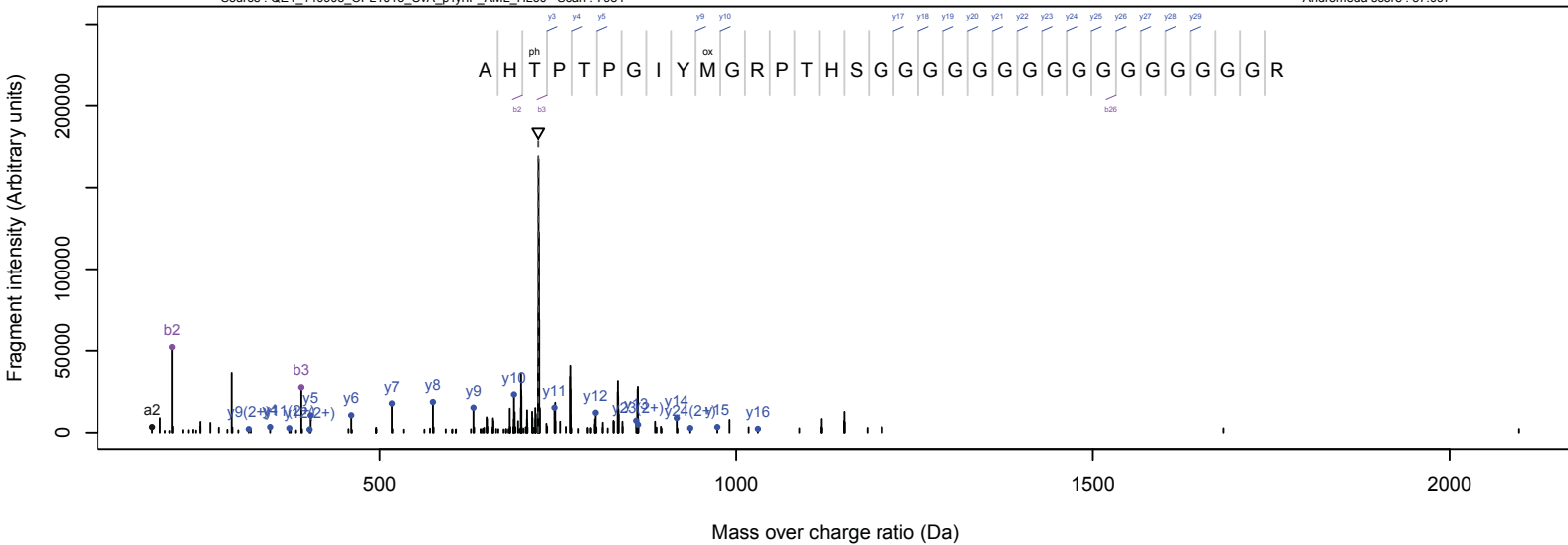
Andromeda score: 57.144



AHT(ph)PTPGIYMGRPTHSGGGGGGGGGGGGGGR-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan: 7984

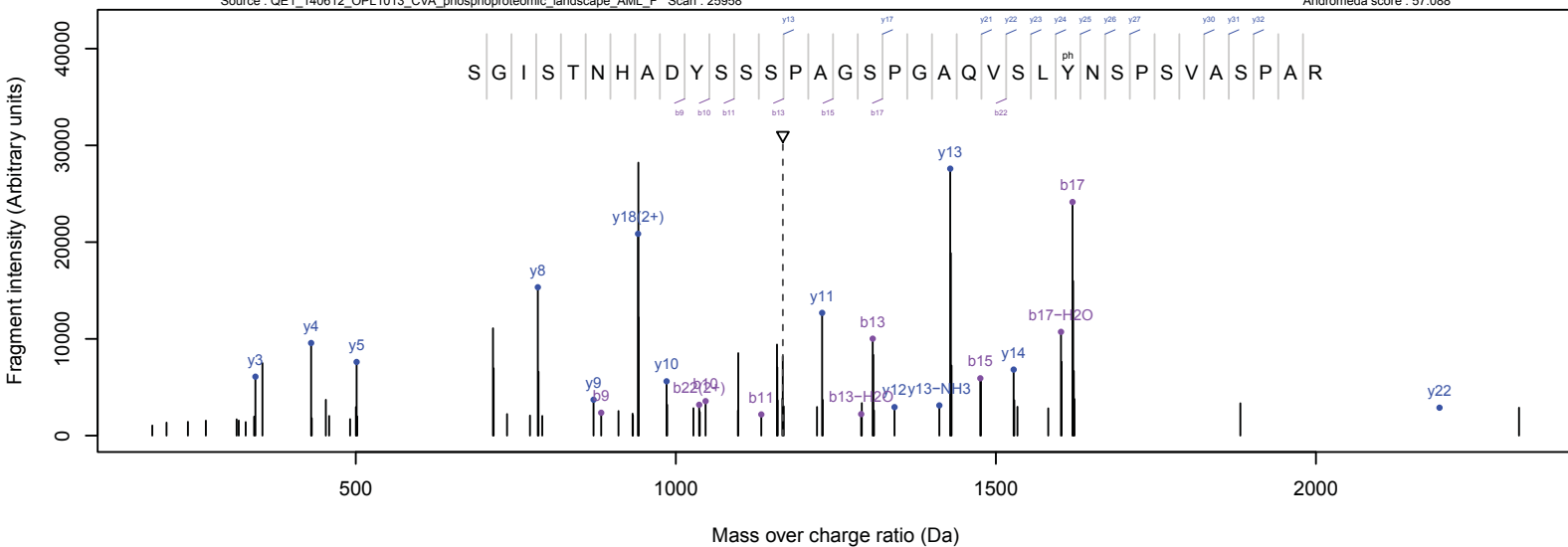
Andromeda score: 57.097



## SGISTNHADYSSSPAGSPGAQVSLY(ph)NSPSVASPAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 25958

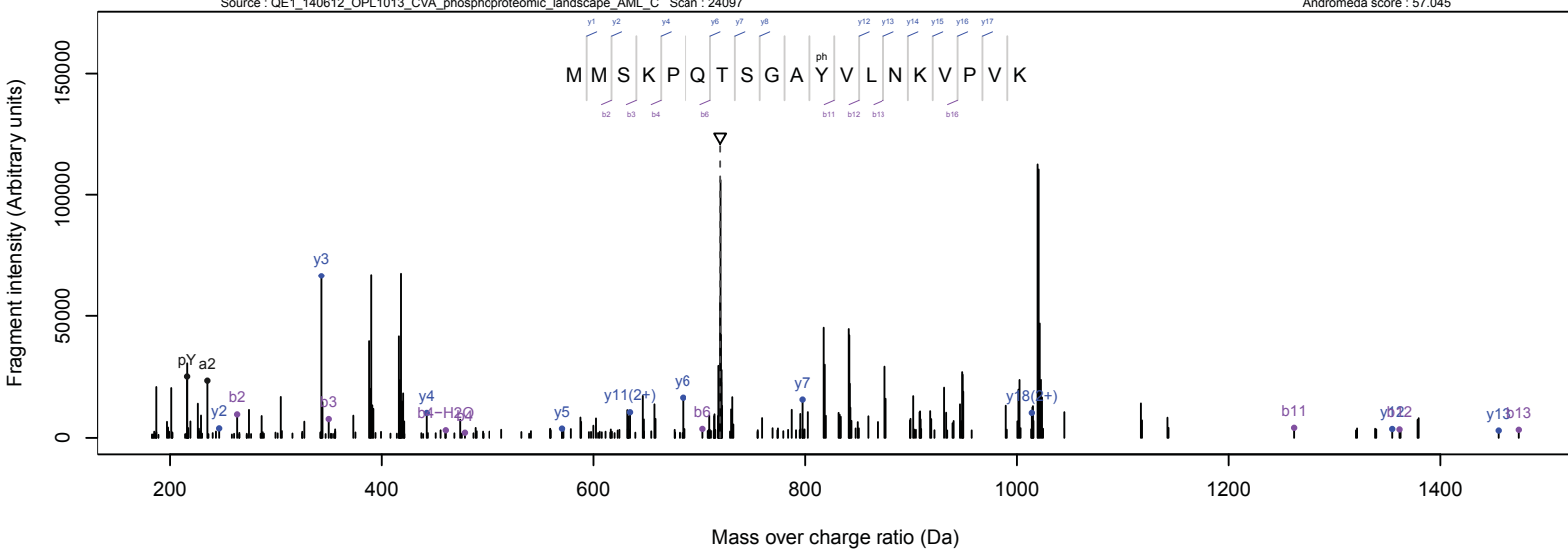
Andromeda score : 57.088



## MMSKPQTSGAYVLNKVPVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24097

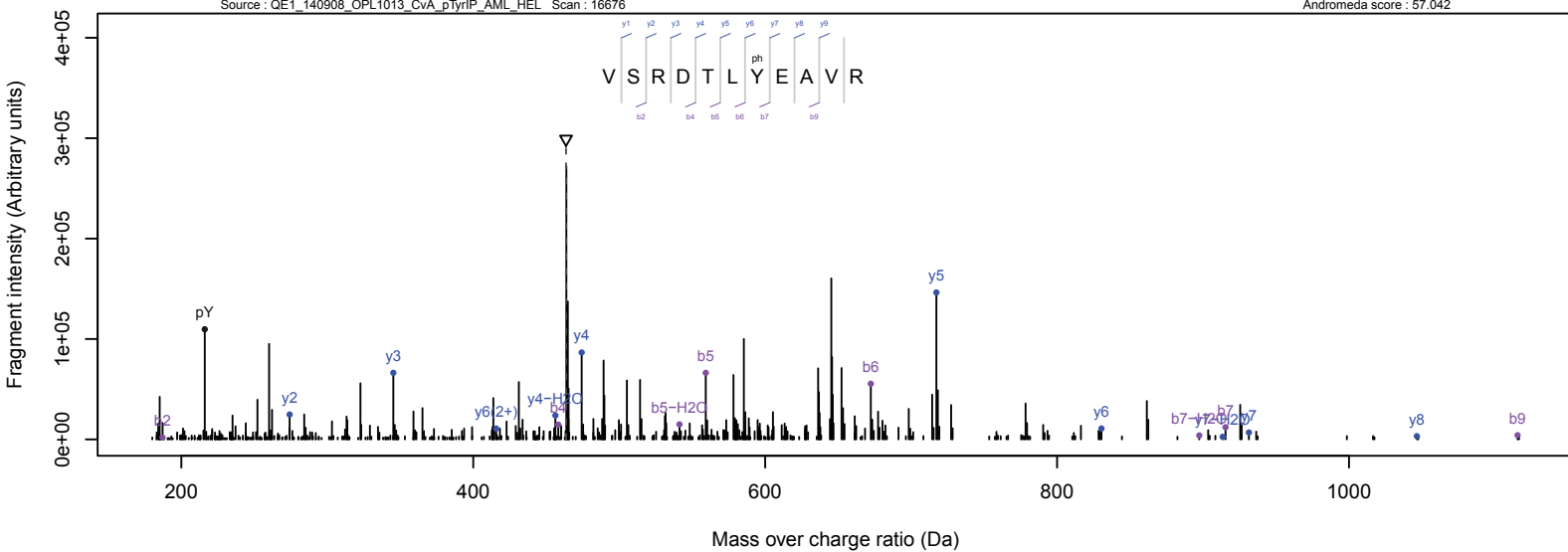
Andromeda score : 57.045



## VSRDTLYEAVR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 16676

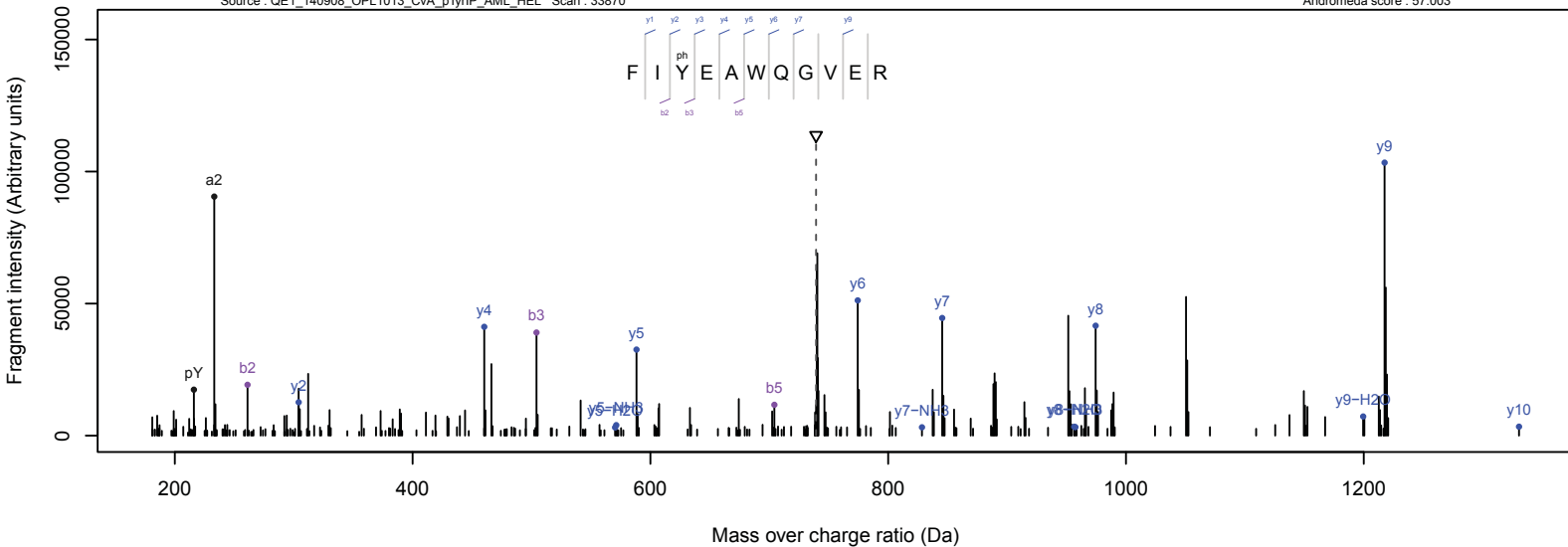
Andromeda score : 57.042



FIY(ph)EAWQGV<sup>ER</sup>-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 33870

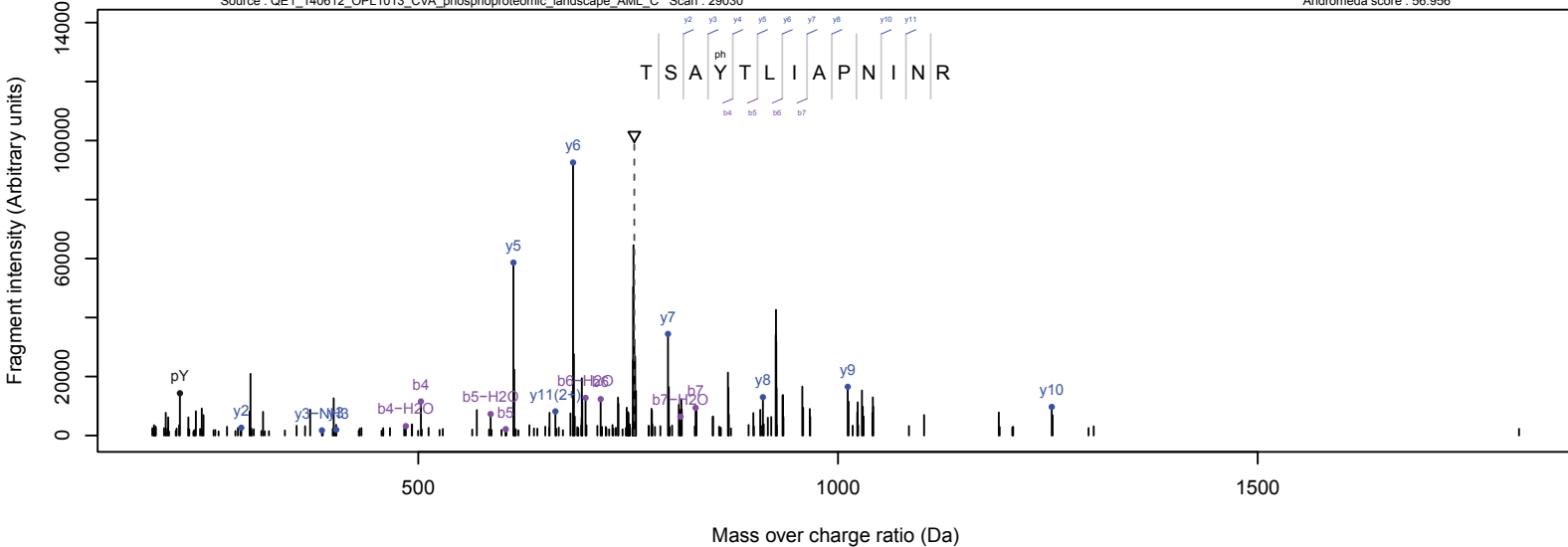
Andromeda score: 57.003



## TSAY(ph)TLIAPNIR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 29030

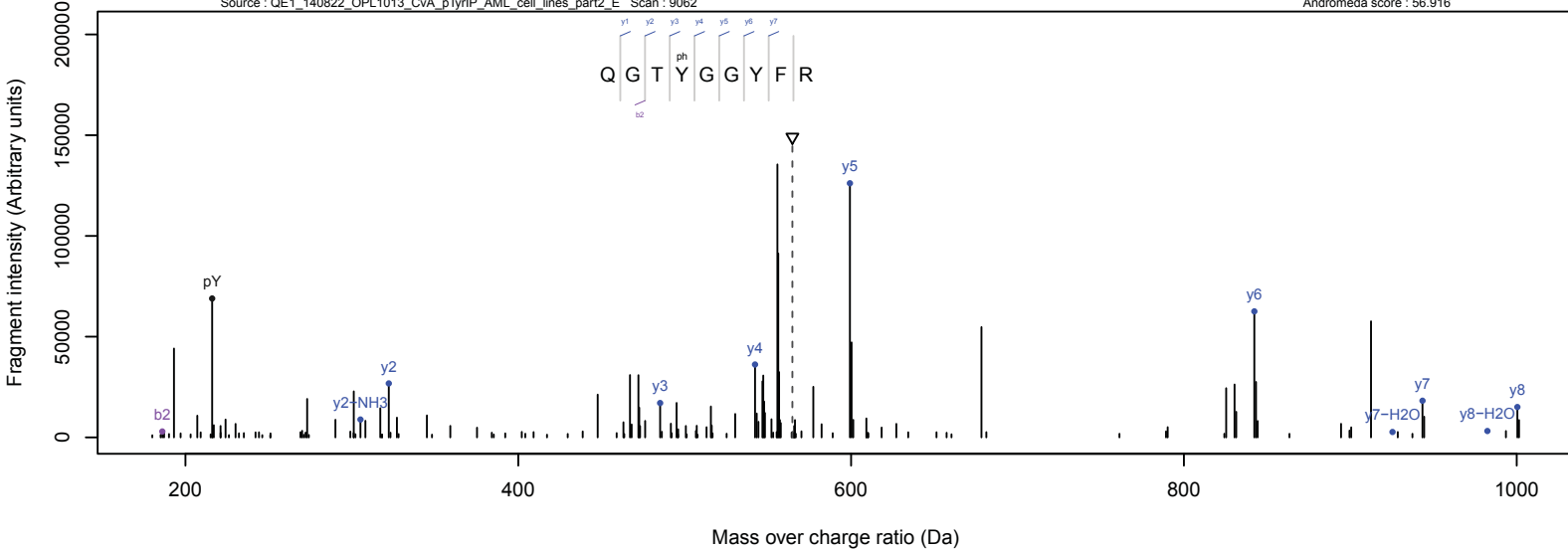
Andromeda score: 56.956



## QGTY(ph)GGYFR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 9062

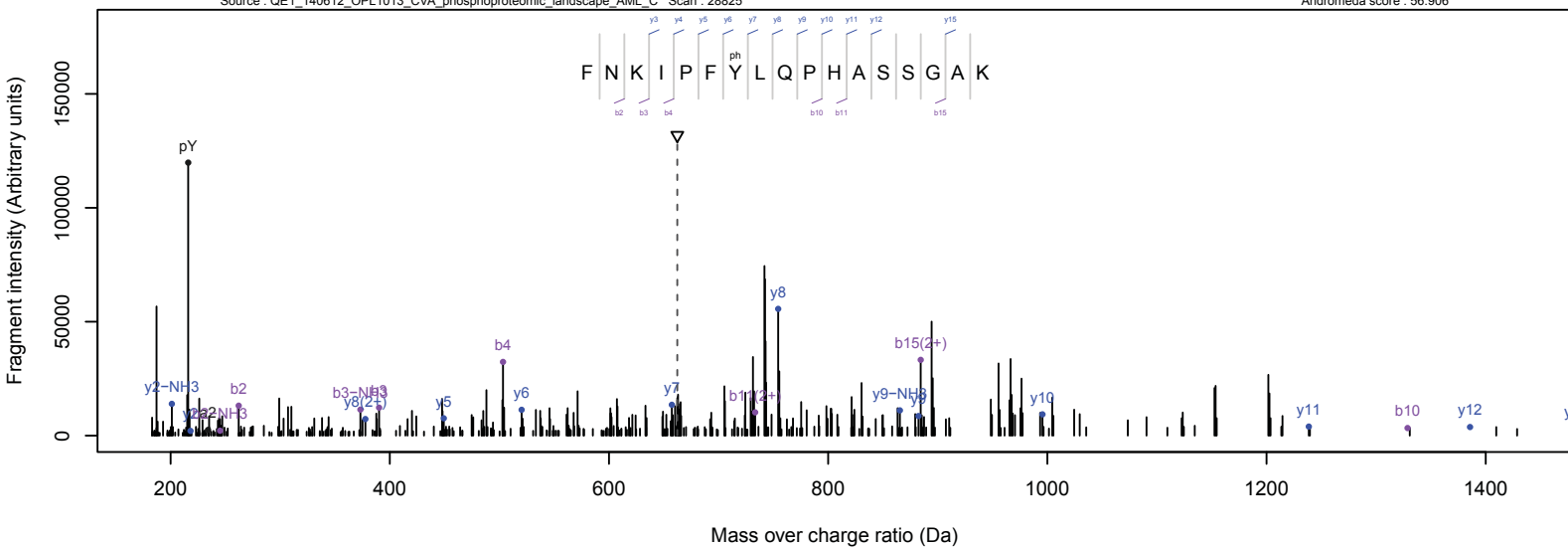
Andromeda score: 56.916



**FNKIPFY(ph)LQPHASSGAK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28825

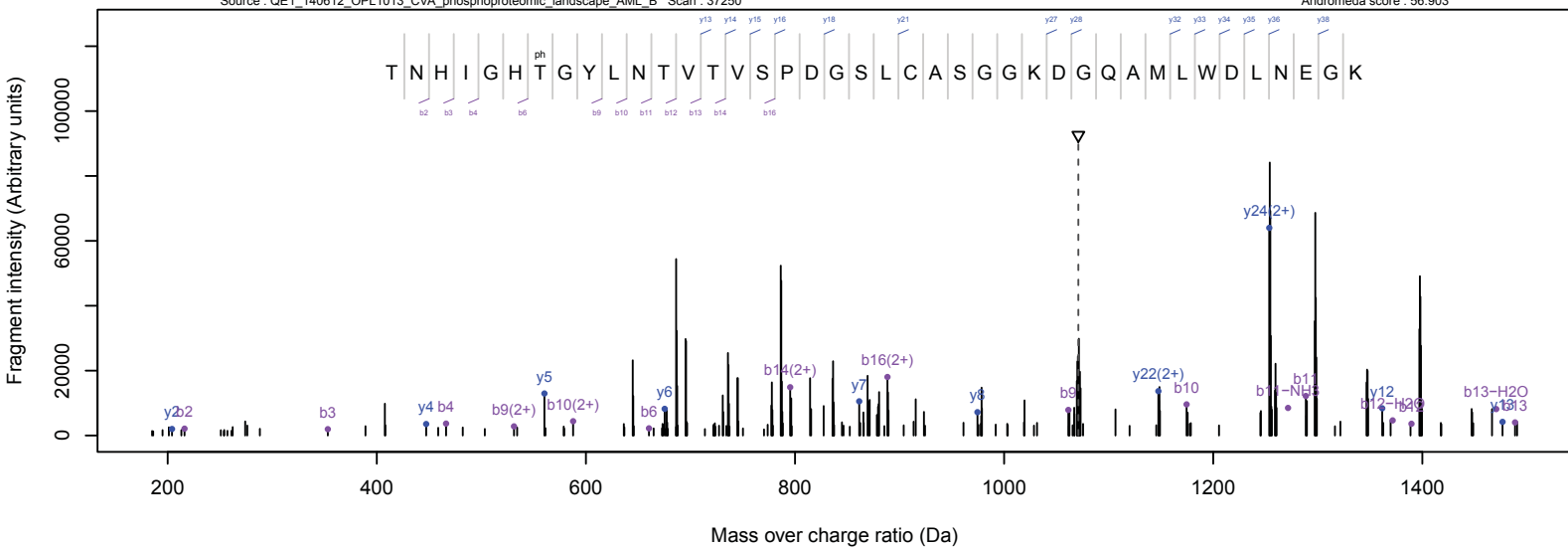
Andromeda score : 56.906



**TNHIGHT(ph)GYLNTVTVSPDGLSCASGGKDGQAMLWDLNEGK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37250

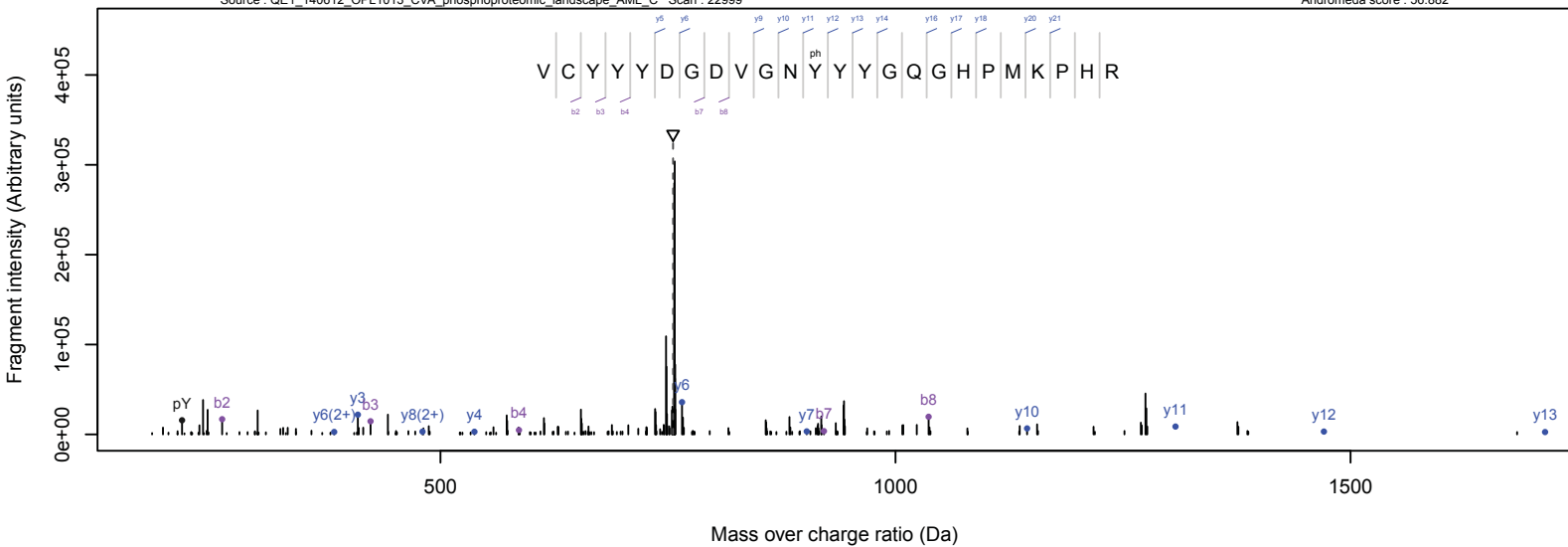
Andromeda score : 56.903



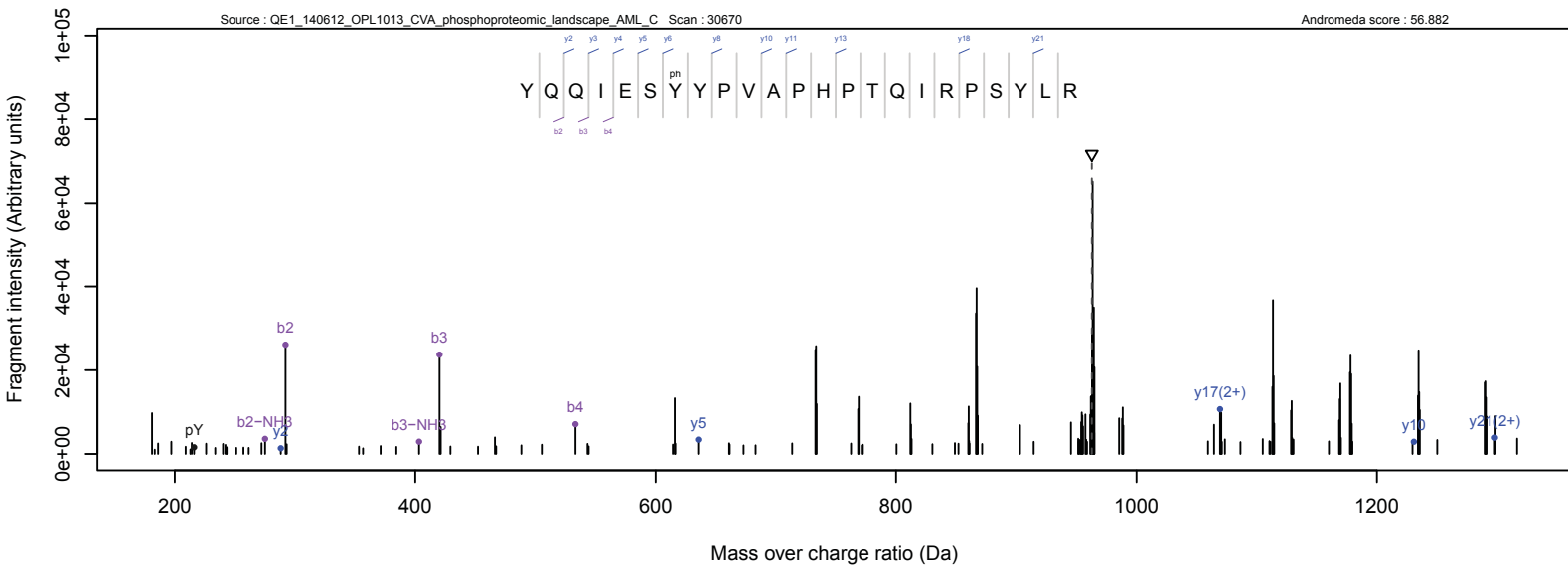
**VCYYDGDVGNYPYGGQHPMKPHR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22999

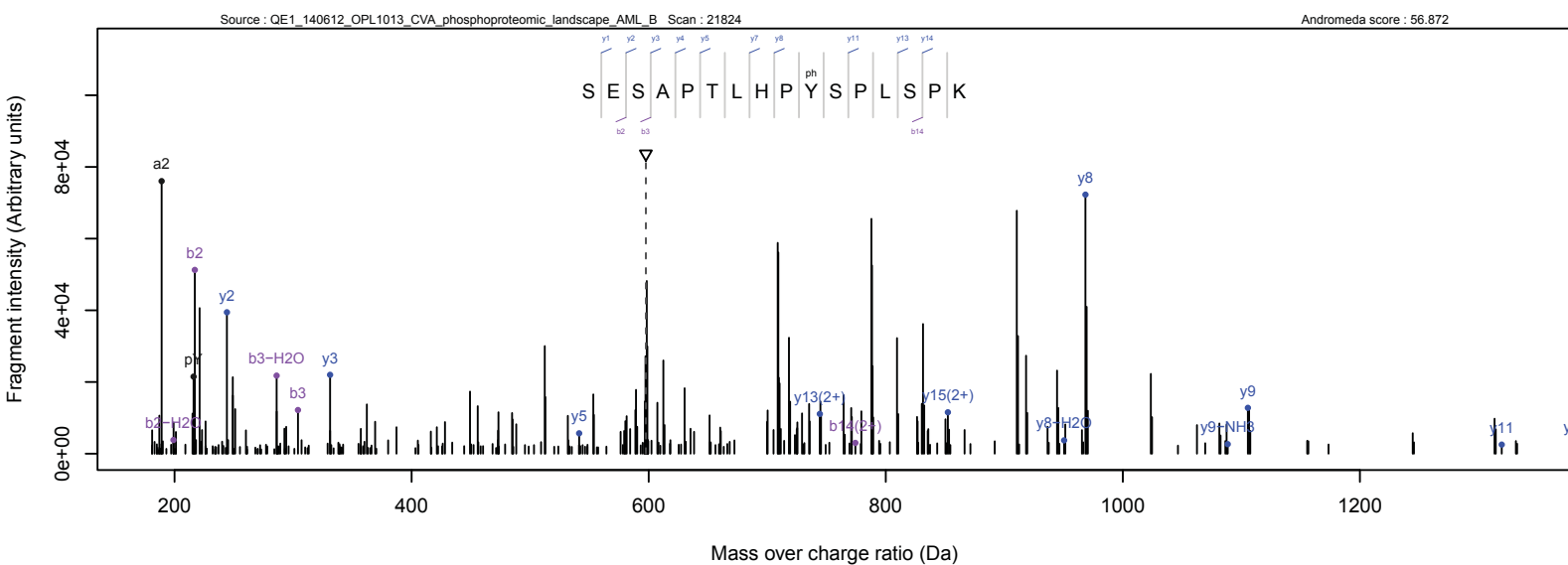
Andromeda score : 56.882



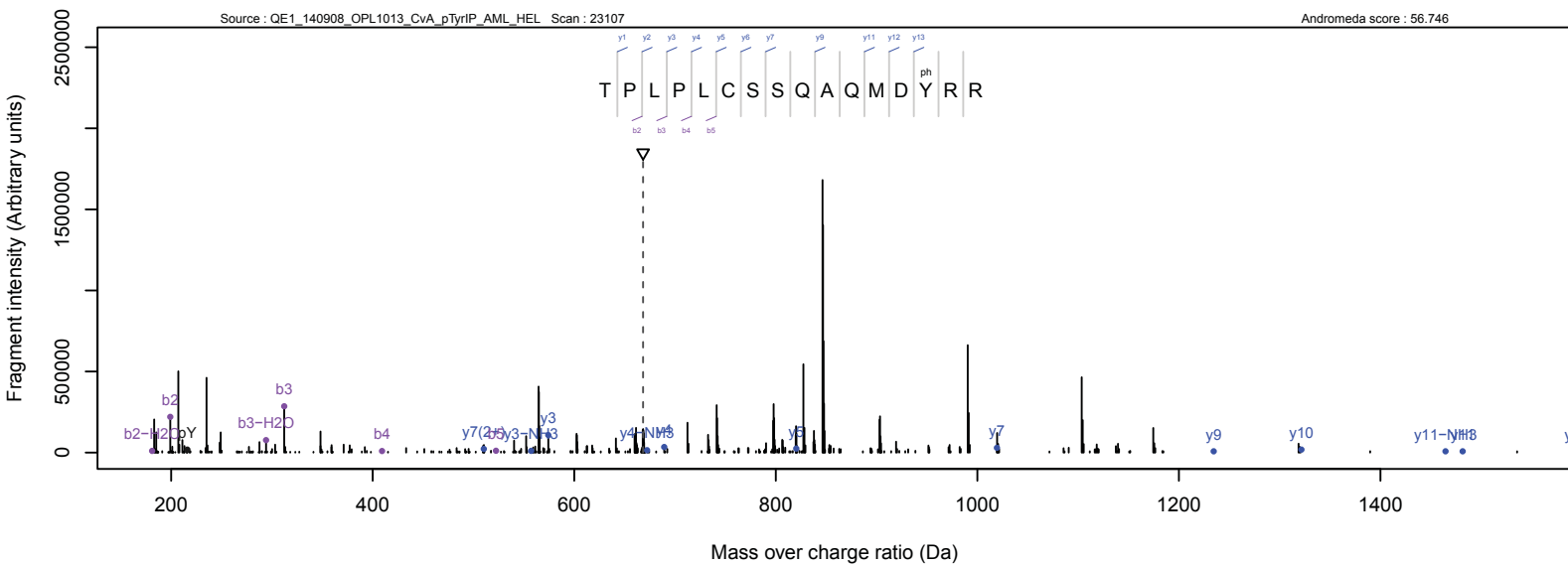
YQQIESY(ph)YPVAPHPTQIRPSYL<sup>R</sup>-3+



SESAPTLHPY(ph)SPLSPK<sup>-3+</sup>



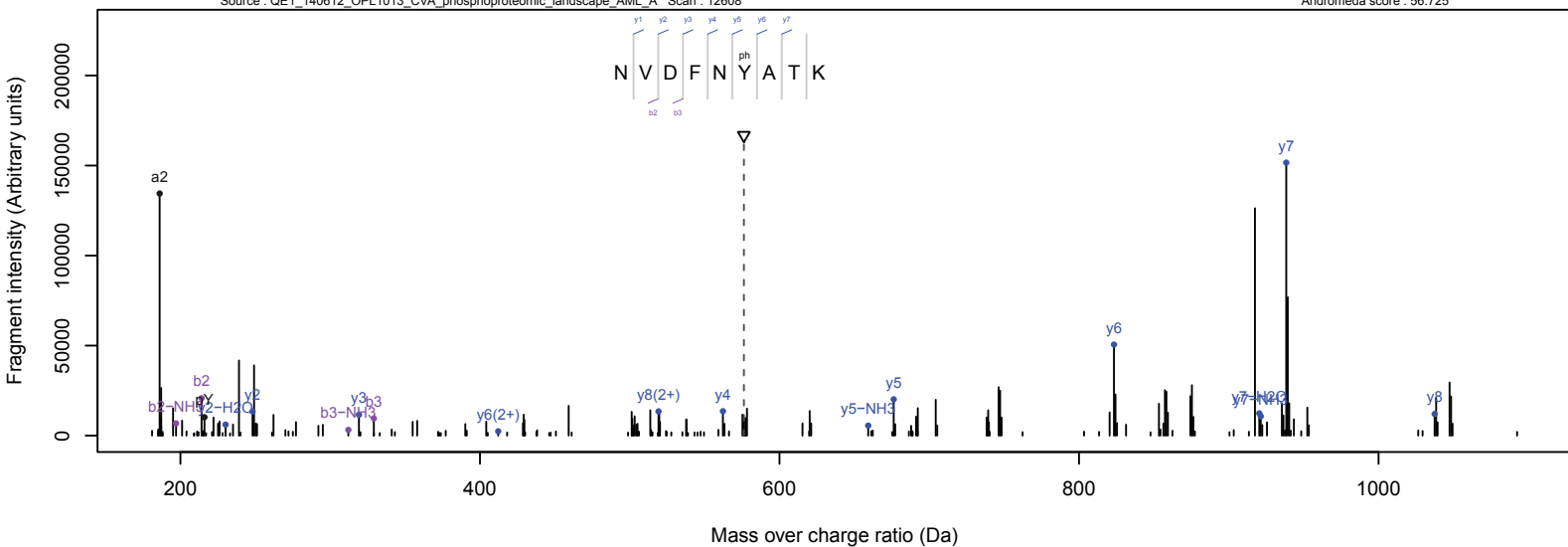
TPLPLCSSQAQMDY(ph)RR<sup>-3+</sup>



## NVDFNY(ph)ATK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12608

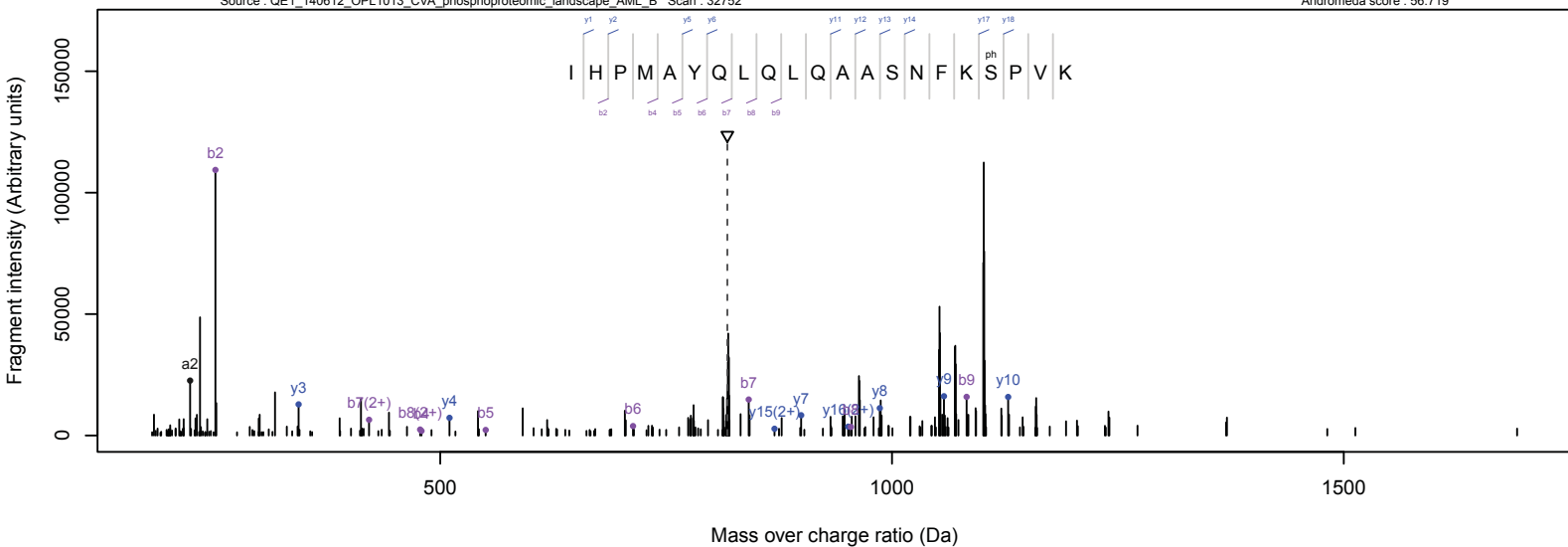
Andromeda score : 56.725



## IHPMAYQLQLQAASNFKS(ph)PVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32752

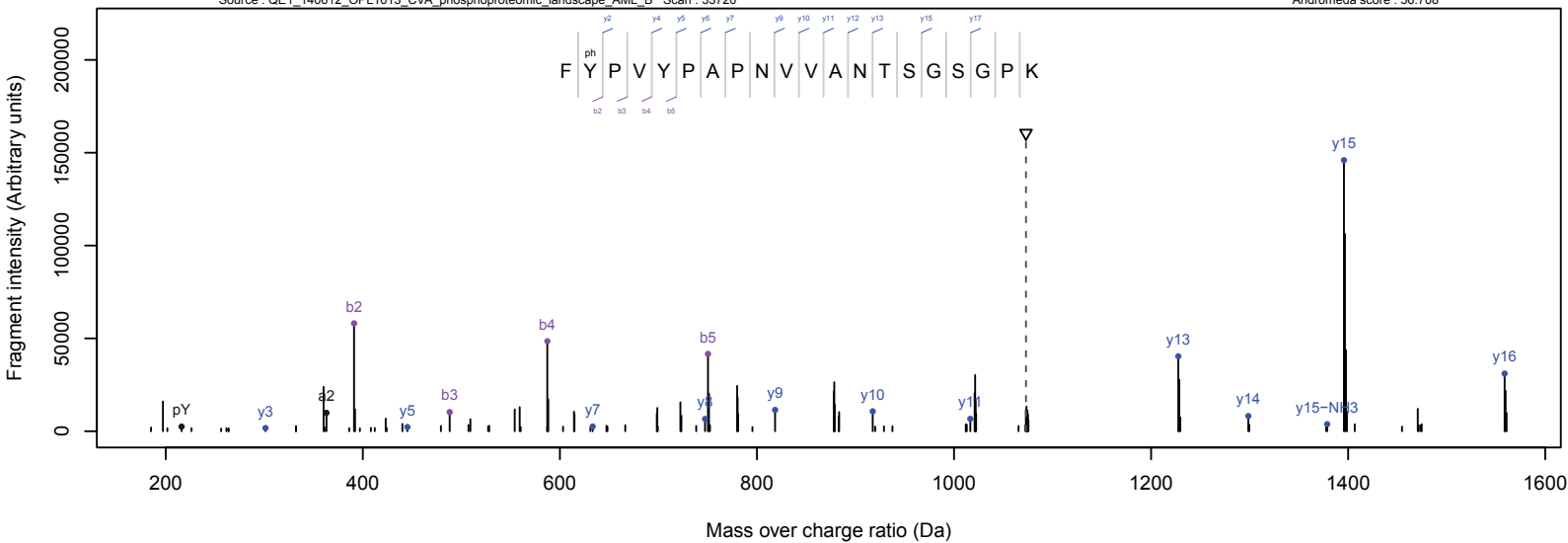
Andromeda score : 56.719



## FY(ph)PVYPAPNVVANTSGSGPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33720

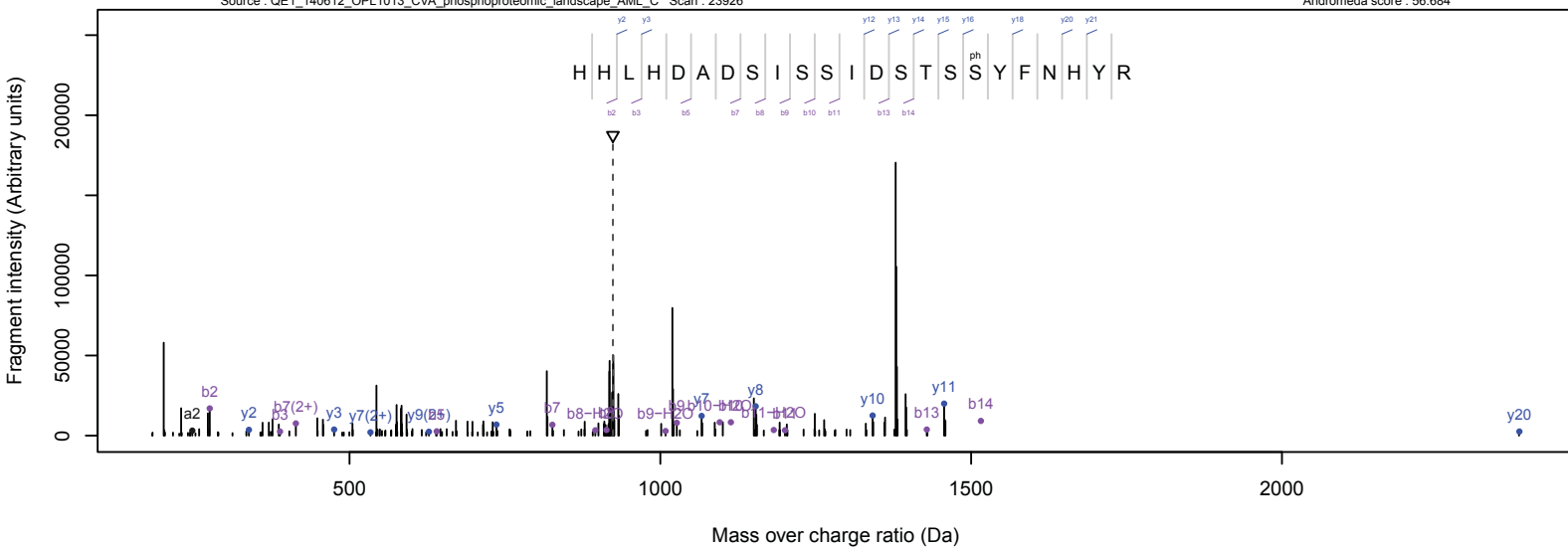
Andromeda score : 56.708



HHLHDADSISSIDSTSS<sup>ph</sup>YFNHYR-3+

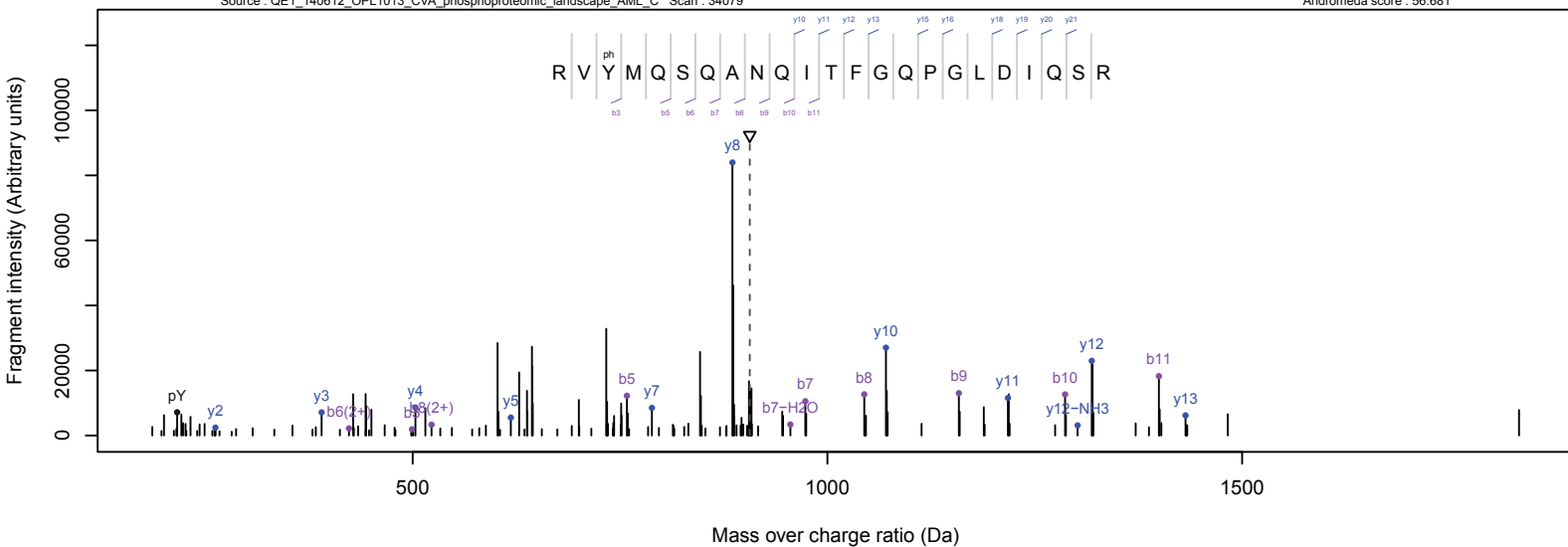
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23926

Andromeda score : 56.684

RVY<sup>ph</sup>MQSQANQITFGQPLDIQSR-3+

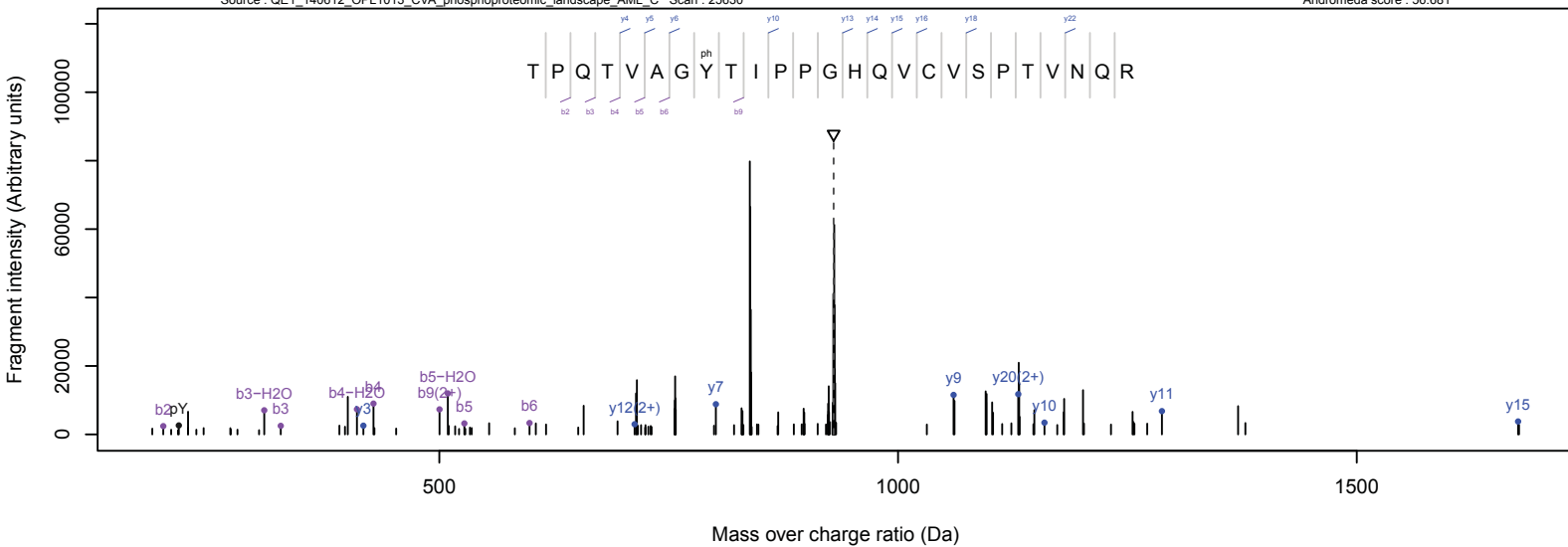
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34079

Andromeda score : 56.681

TPQTVAGY<sup>ph</sup>TIPPGHQVCVSP TVNQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25630

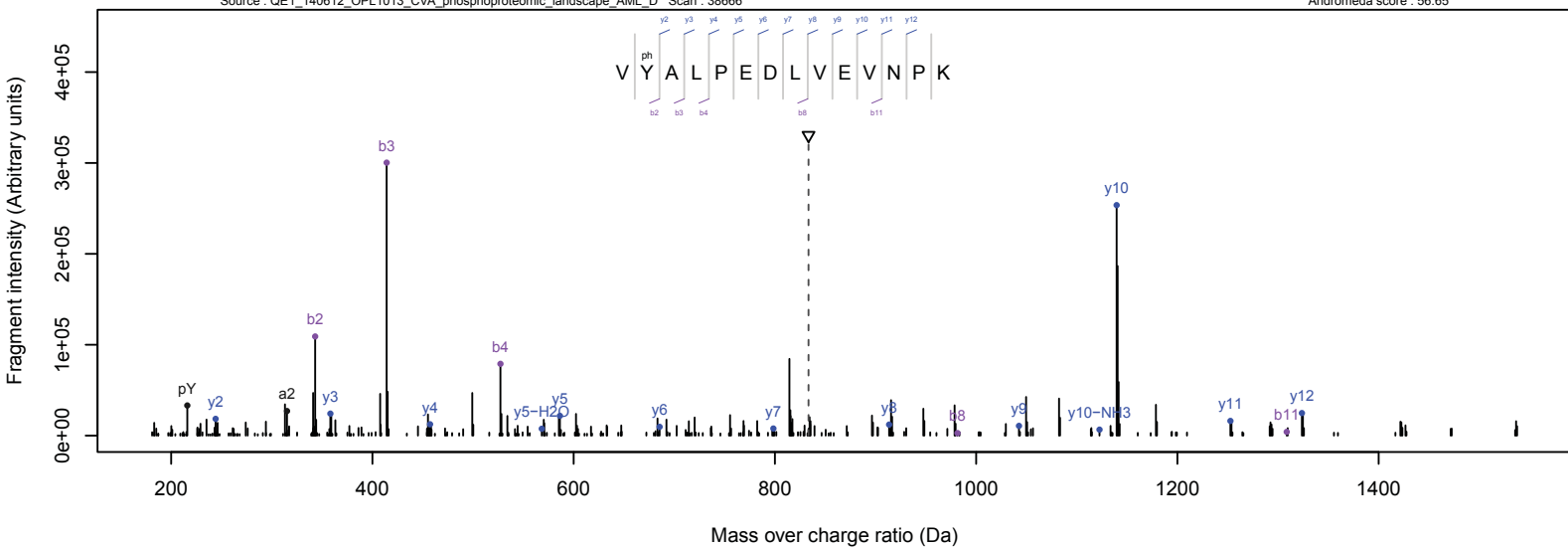
Andromeda score : 56.681



## VY(ph)ALPEDLVEVNPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 38666

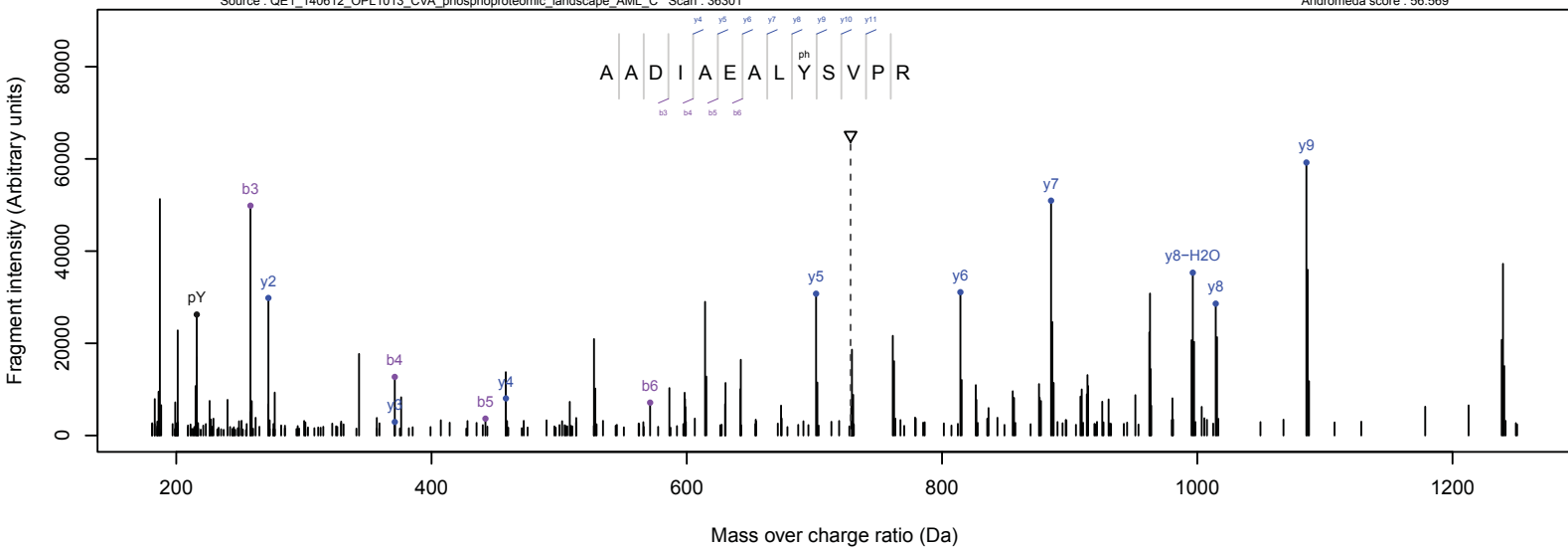
Andromeda score: 56.65



## AADIAEALY(ph)SVPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 36301

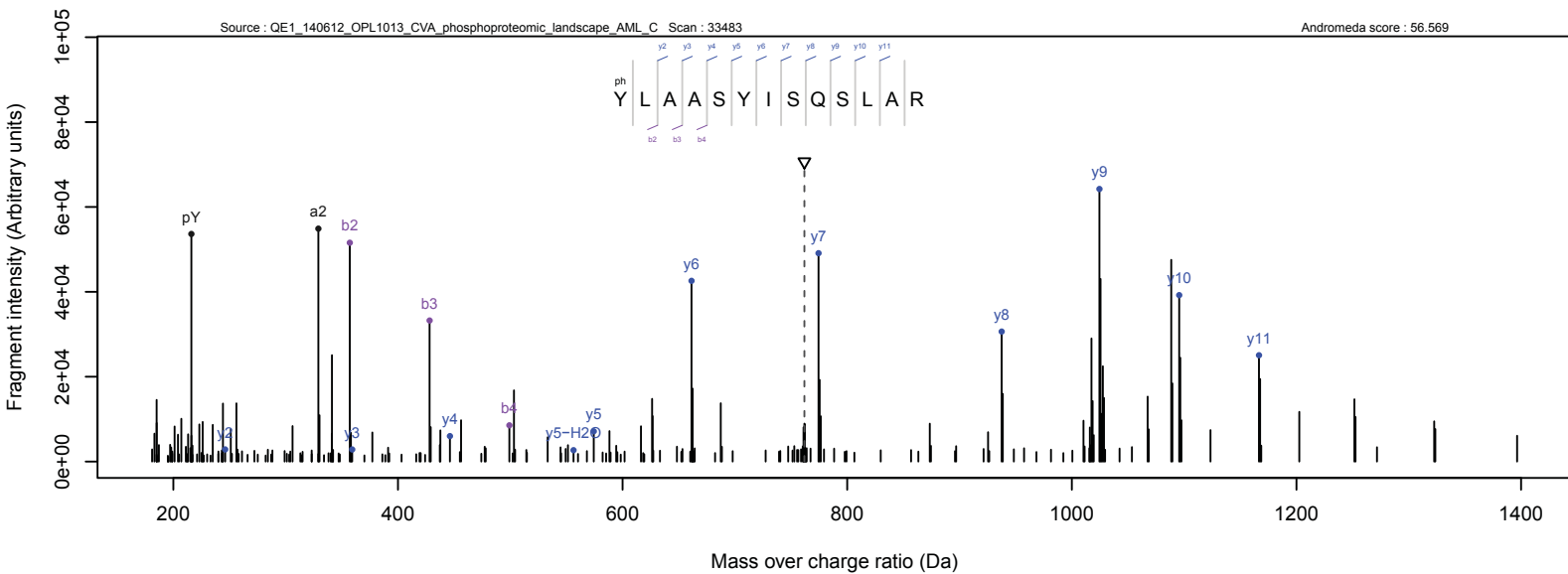
Andromeda score: 56.569



## Y(ph)LAASYISQSLAR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 33483

Andromeda score: 56.569

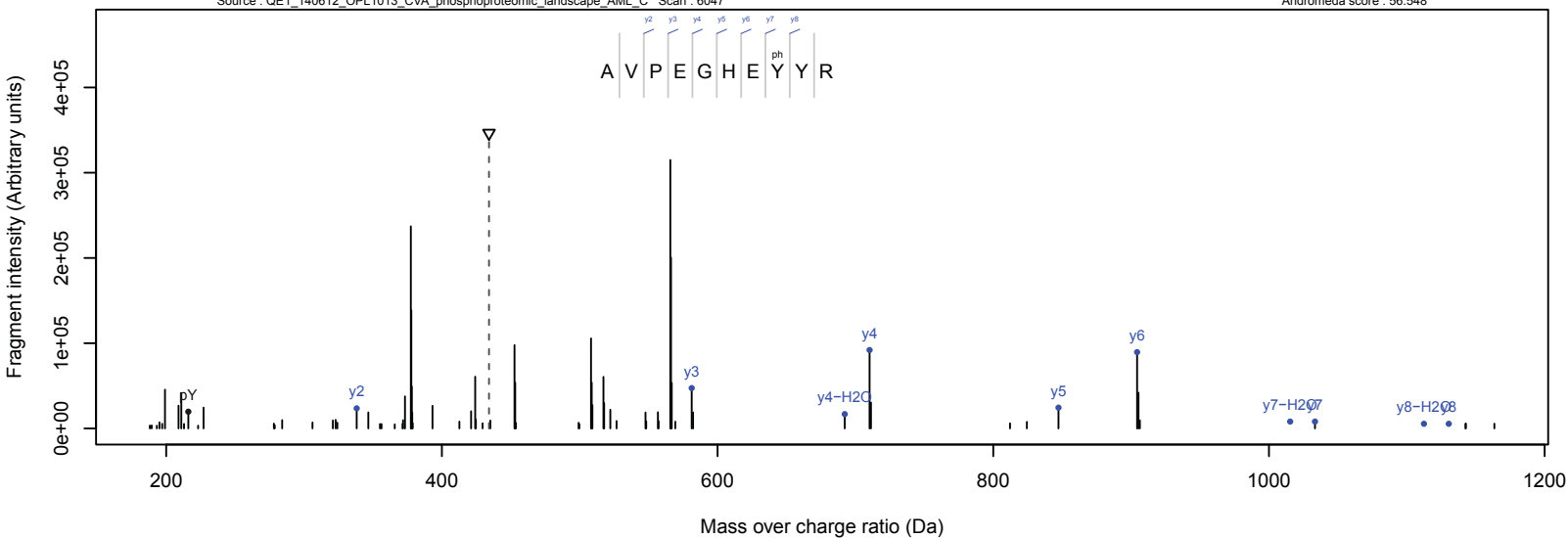




AVPEGHEY<sup>(ph)</sup>YR-3+

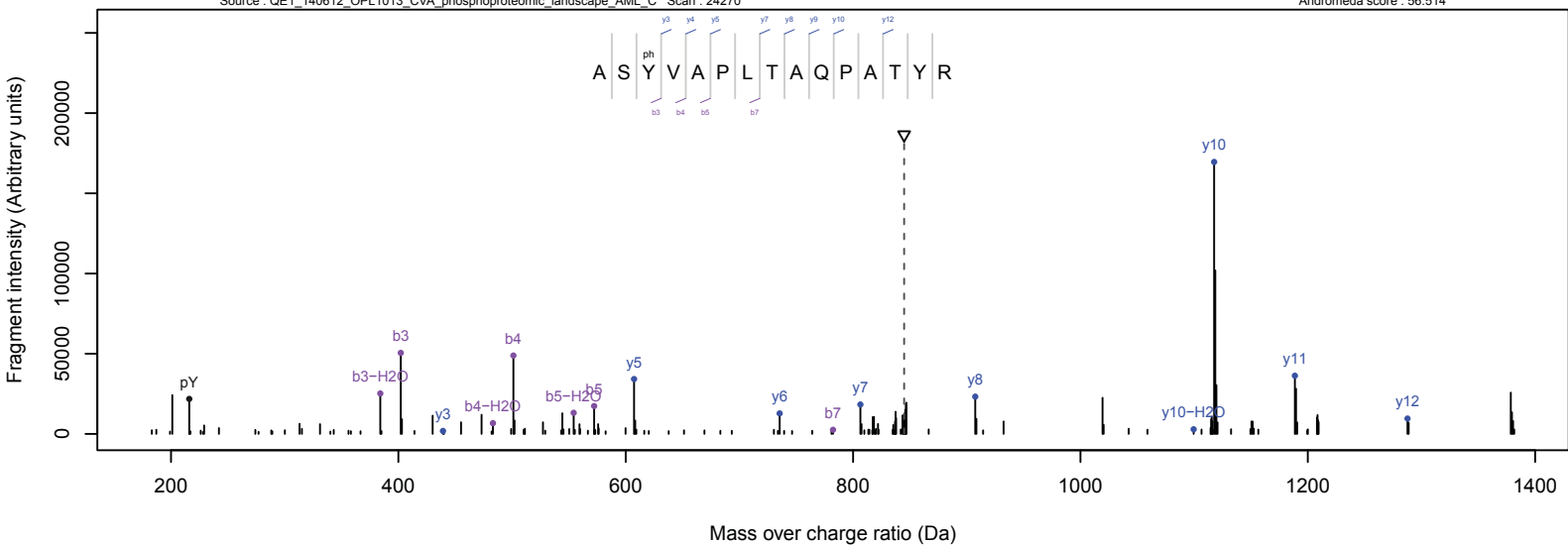
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6047

Andromeda score : 56.548

ASY<sup>(ph)</sup>VAPLTAQPATYR-2+

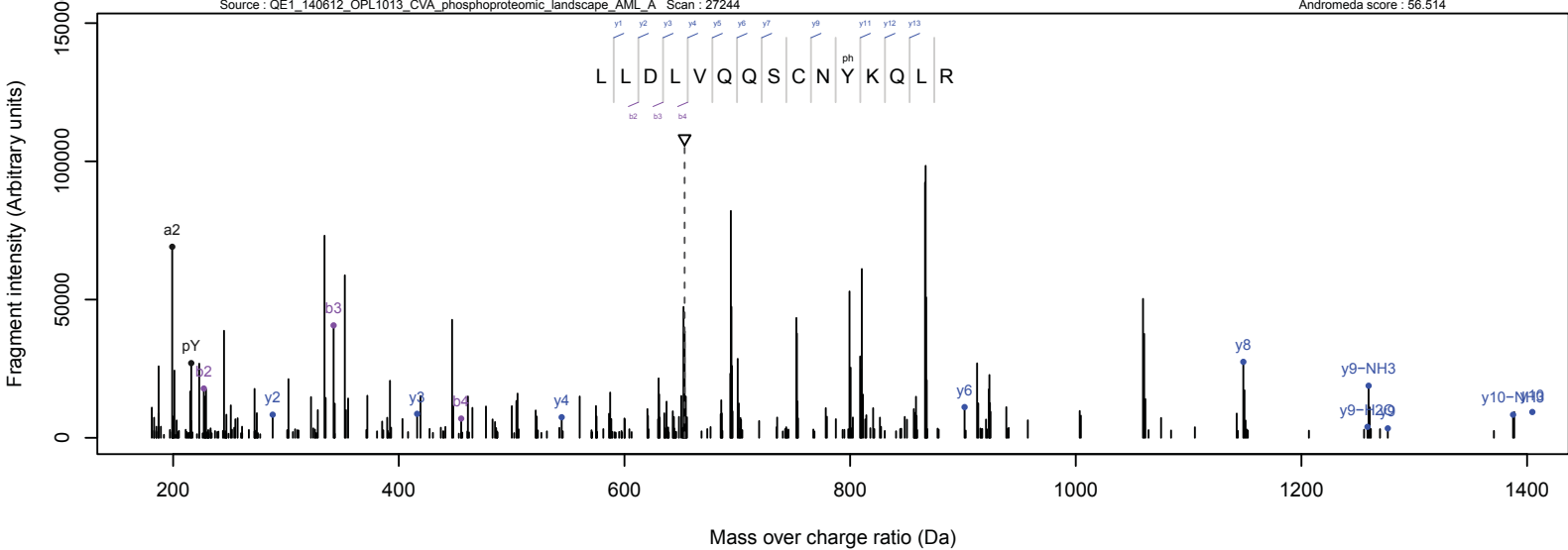
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24270

Andromeda score : 56.514

LLDLVQQSCNY<sup>(ph)</sup>KQLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27244

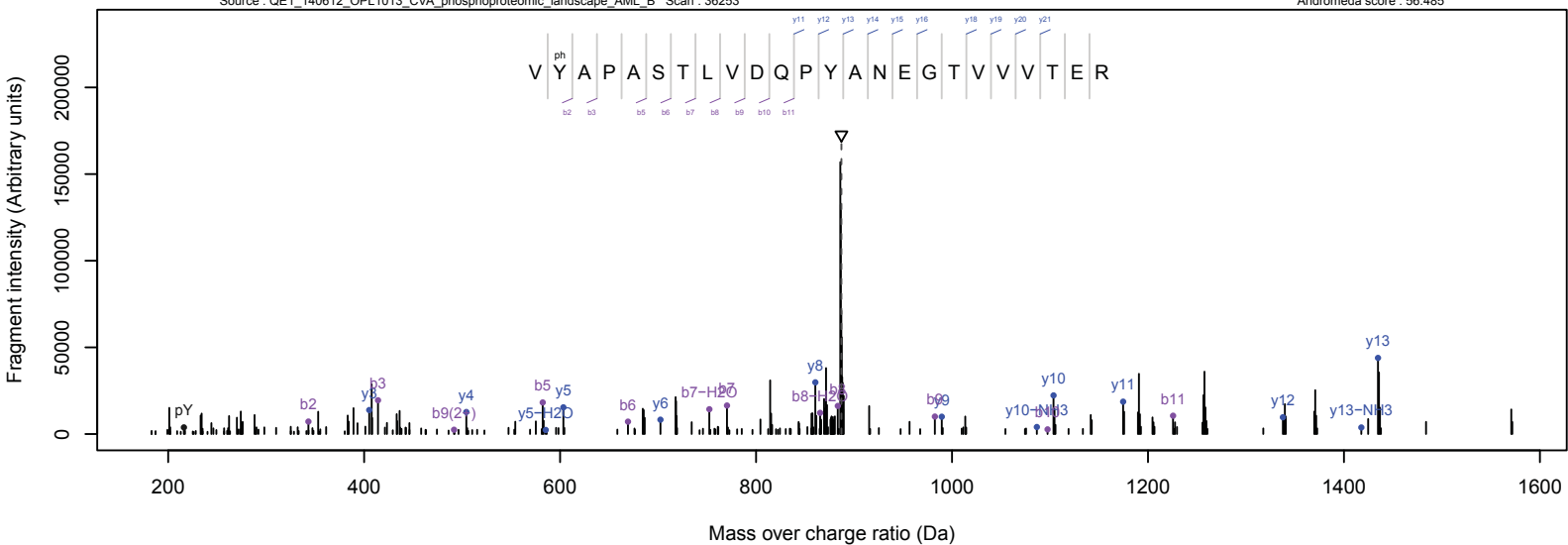
Andromeda score : 56.514



VY(ph)APASTLVLDQPYANEGTVVVTTER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 36253

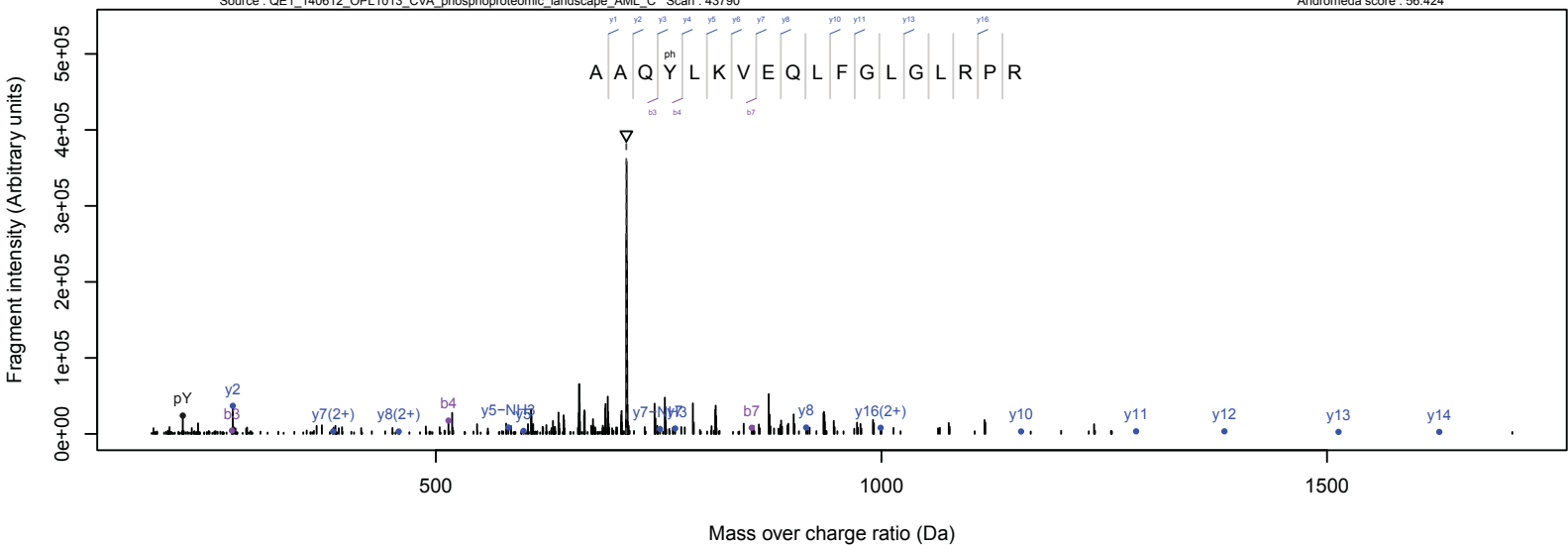
Andromeda score : 56.485



AAQY(ph)LKVEQLFGLGLRPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43790

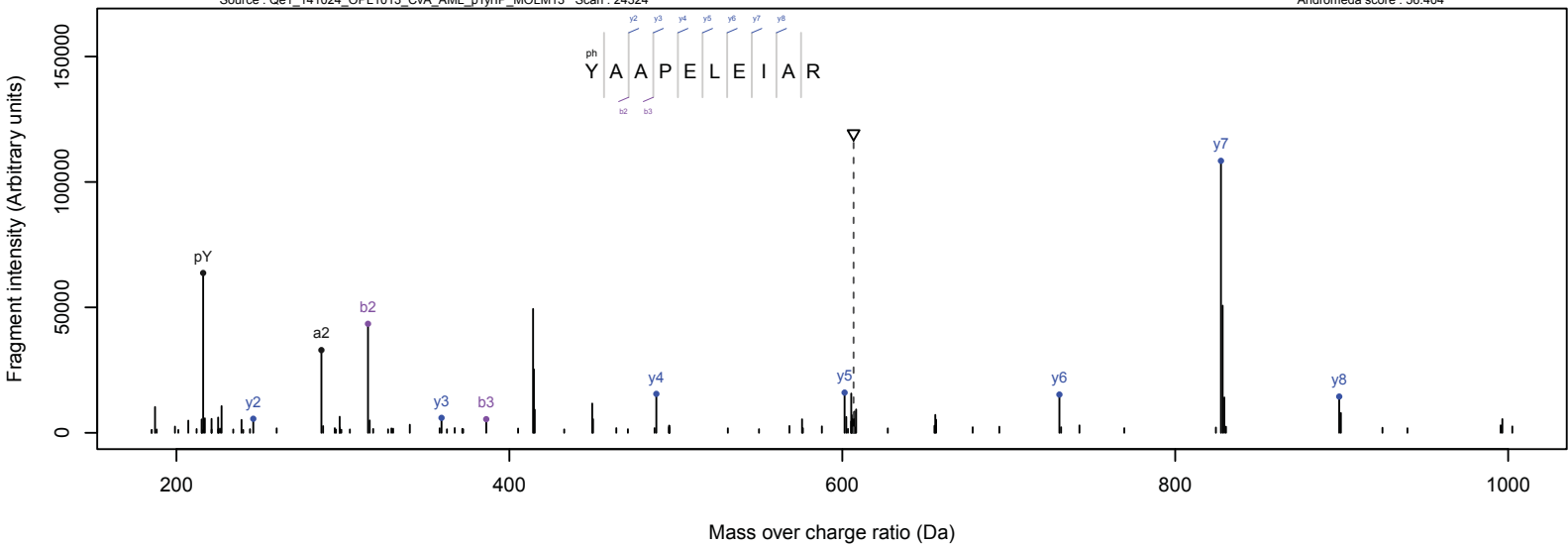
Andromeda score : 56.424



Y(ph)AAPELEIAR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 24324

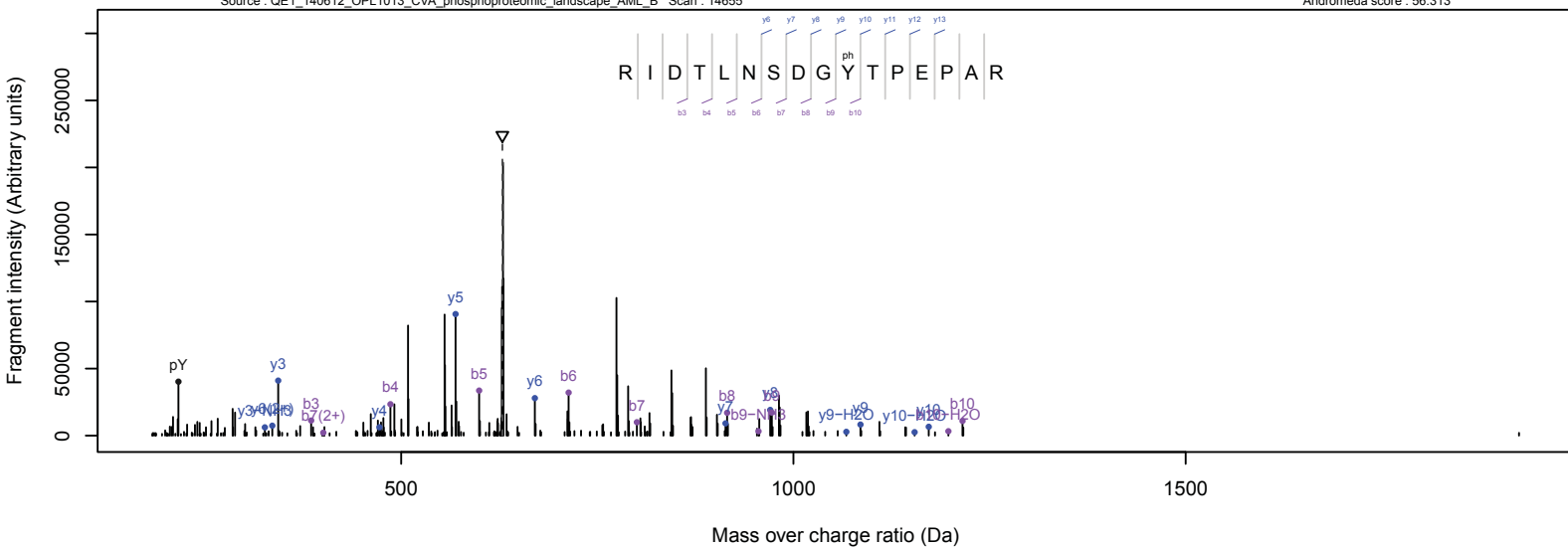
Andromeda score : 56.404



RIDTLNSDGY(ph)TPEPAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14655

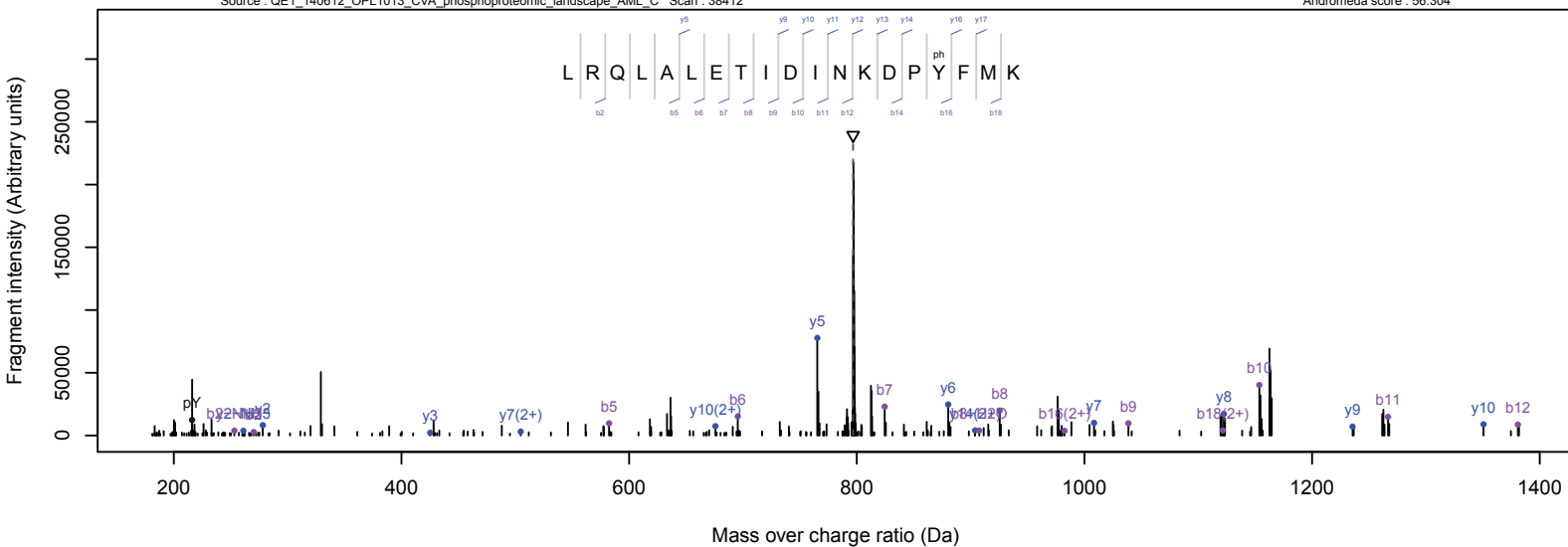
Andromeda score : 56.313



LRQLALETIDINKDYPYFMK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38412

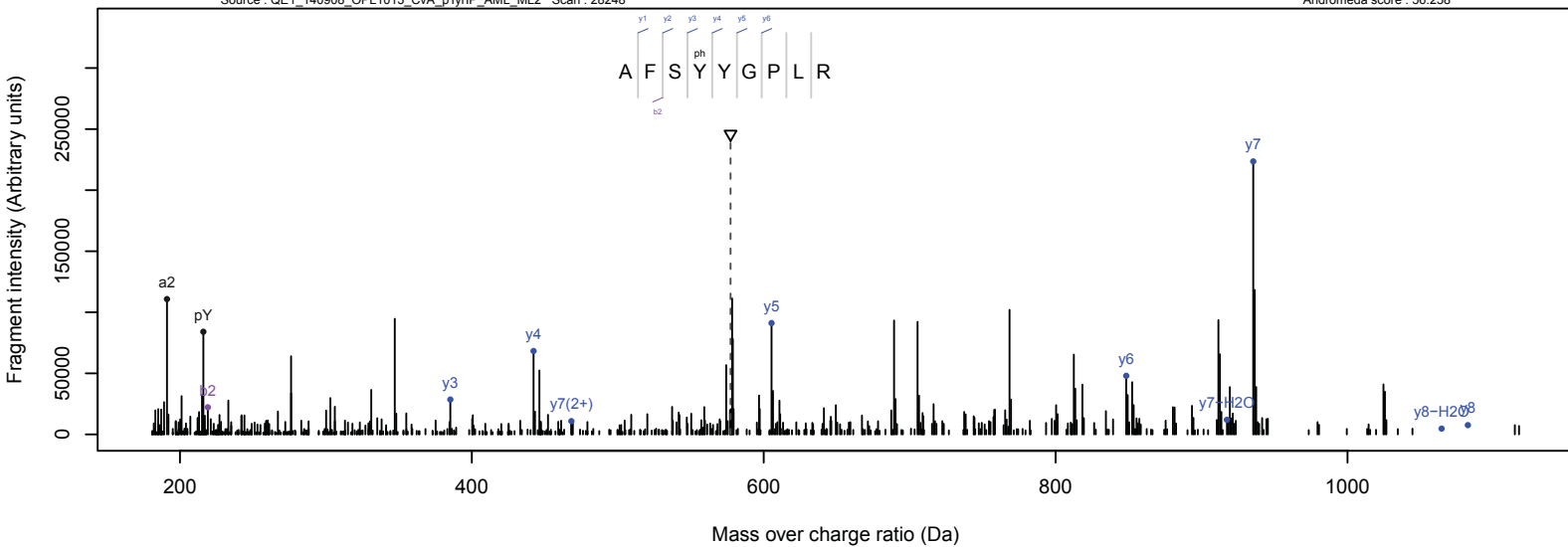
Andromeda score : 56.304



AFSY(ph)YGPLR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 28248

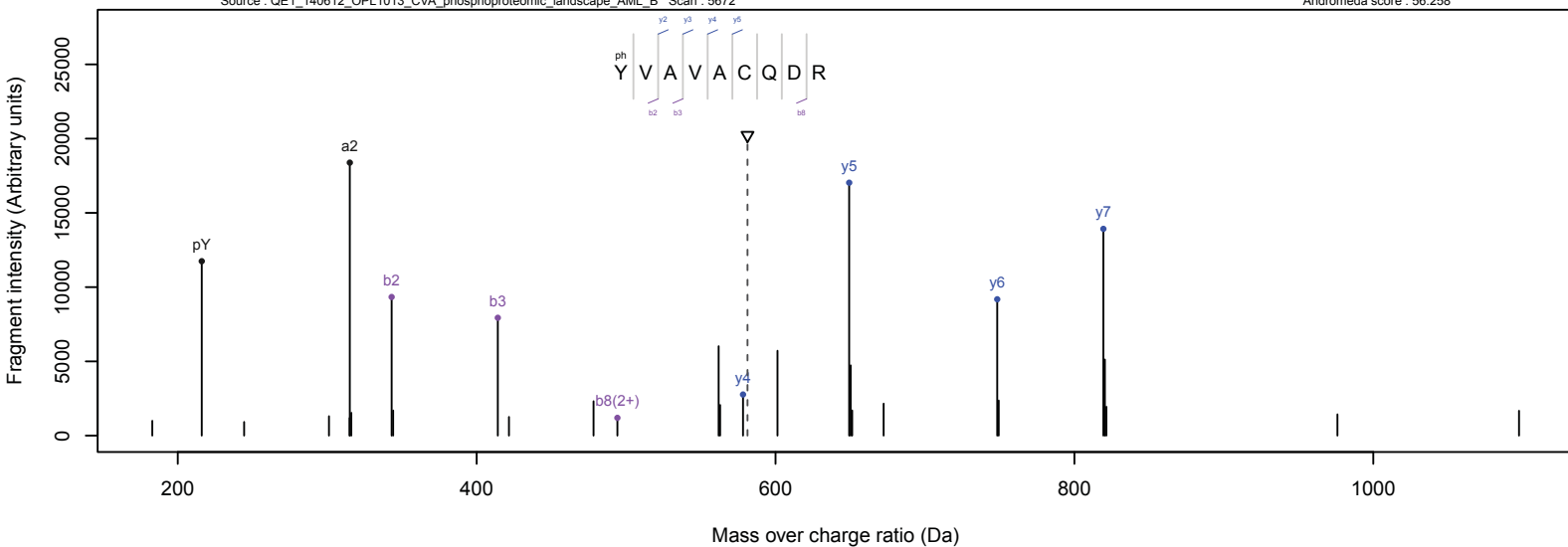
Andromeda score : 56.258



## Y(ph)VAVACQDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 5672

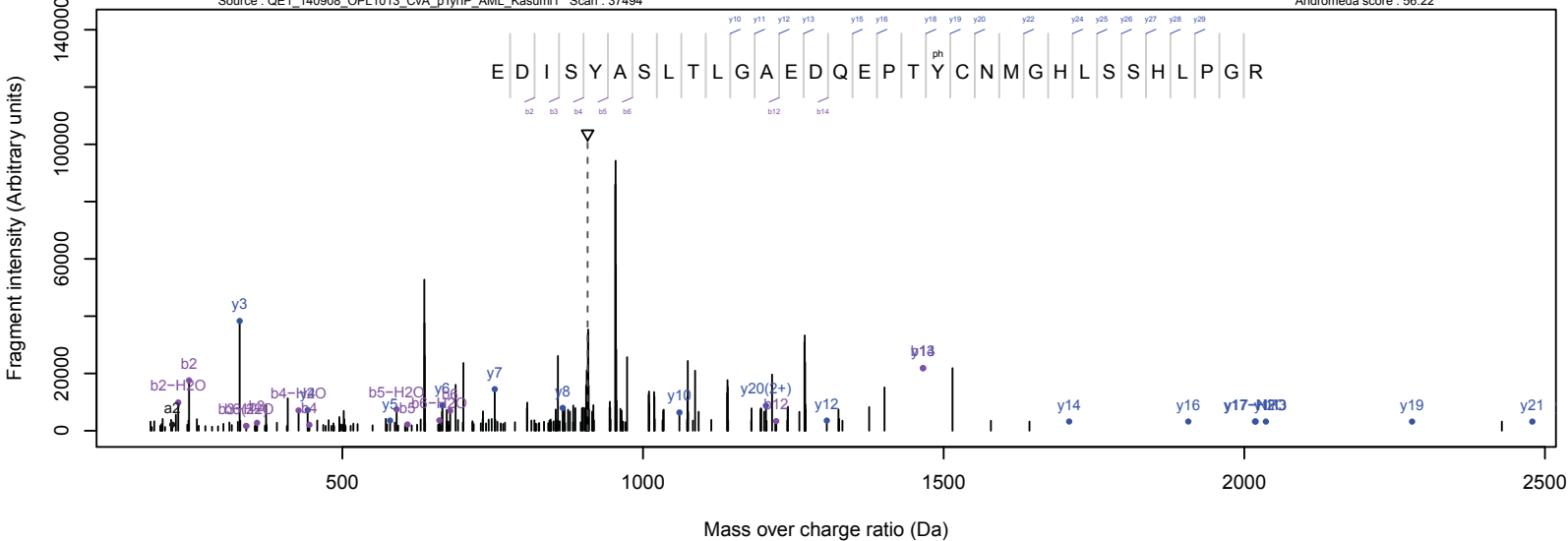
Andromeda score : 56.258



## EDISYASLTTLGAEDQEPTY(ph)CNMGLSSHLPLGR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 37494

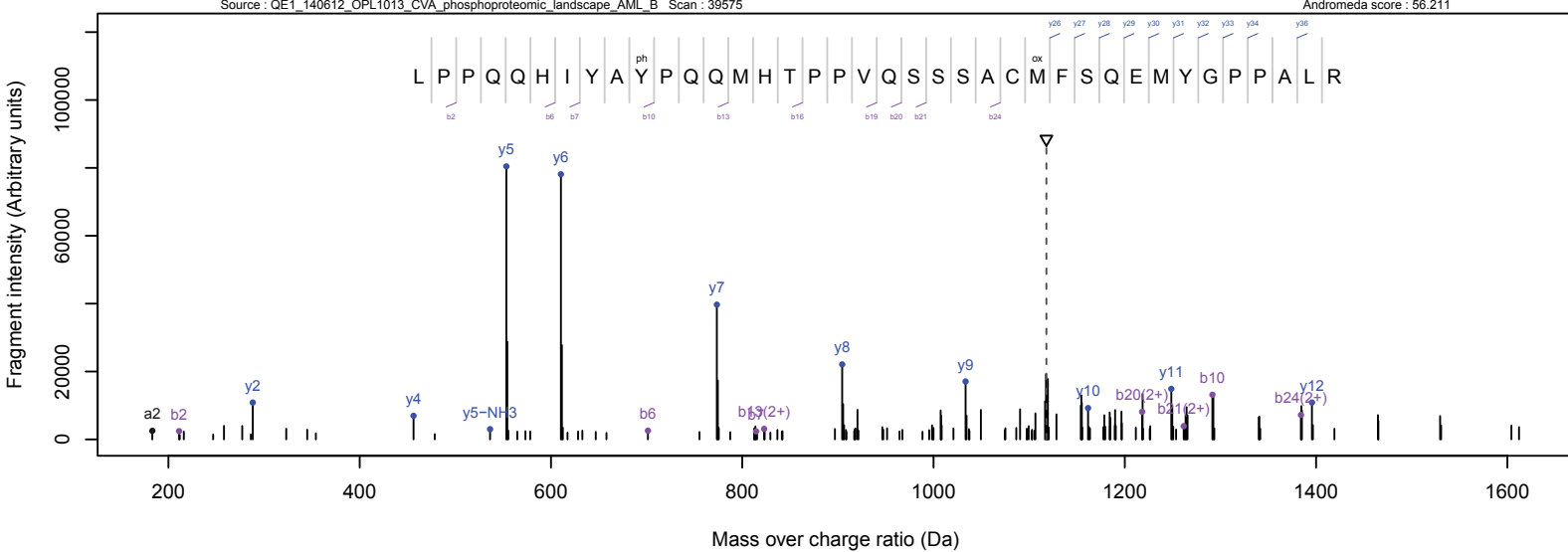
Andromeda score : 56.22



## LPPQQHIYAY(ph)PQQMHTPPVQSSSACM(ox)FSQEMYGPPALR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39575

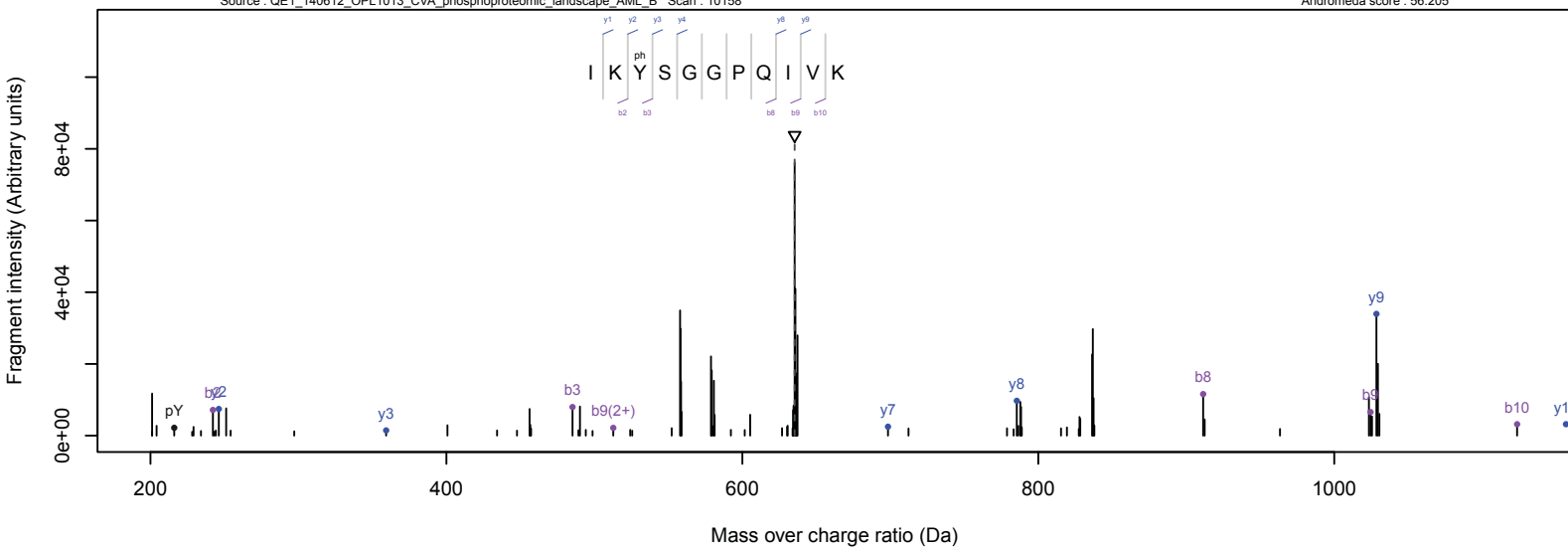
Andromeda score : 56.211



## IKY(ph)SGGPQIVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10158

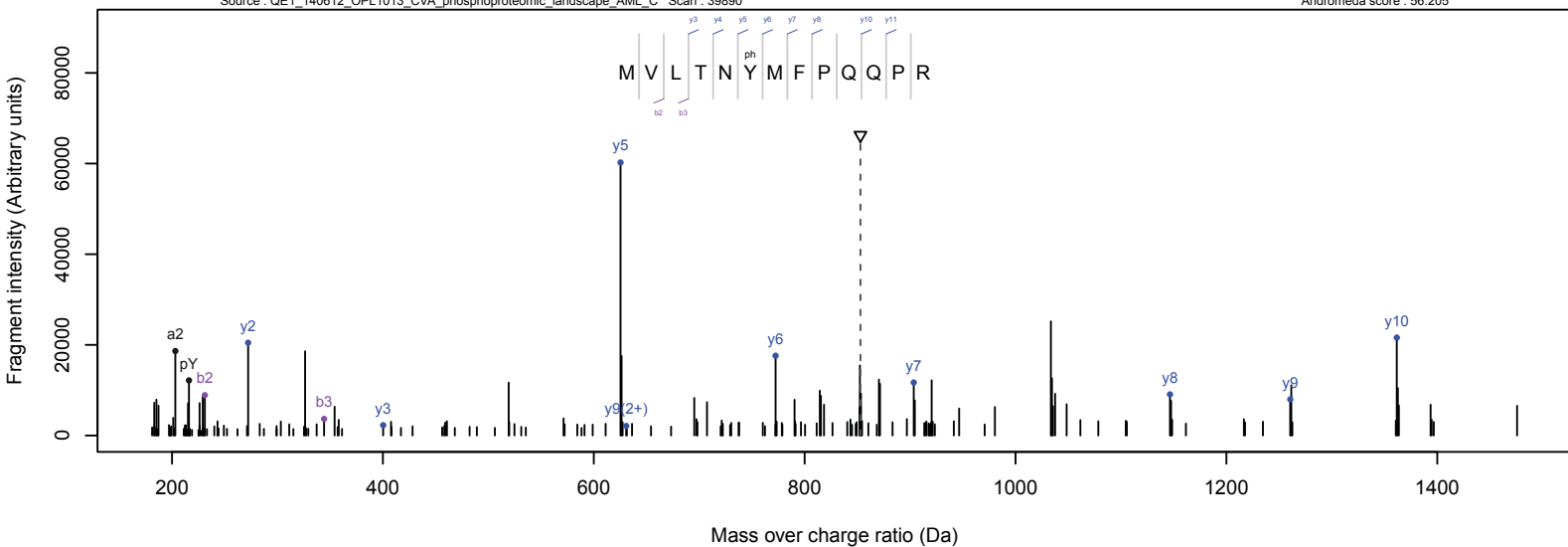
Andromeda score : 56.205



## MVLNTY(ph)MFPQQPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39890

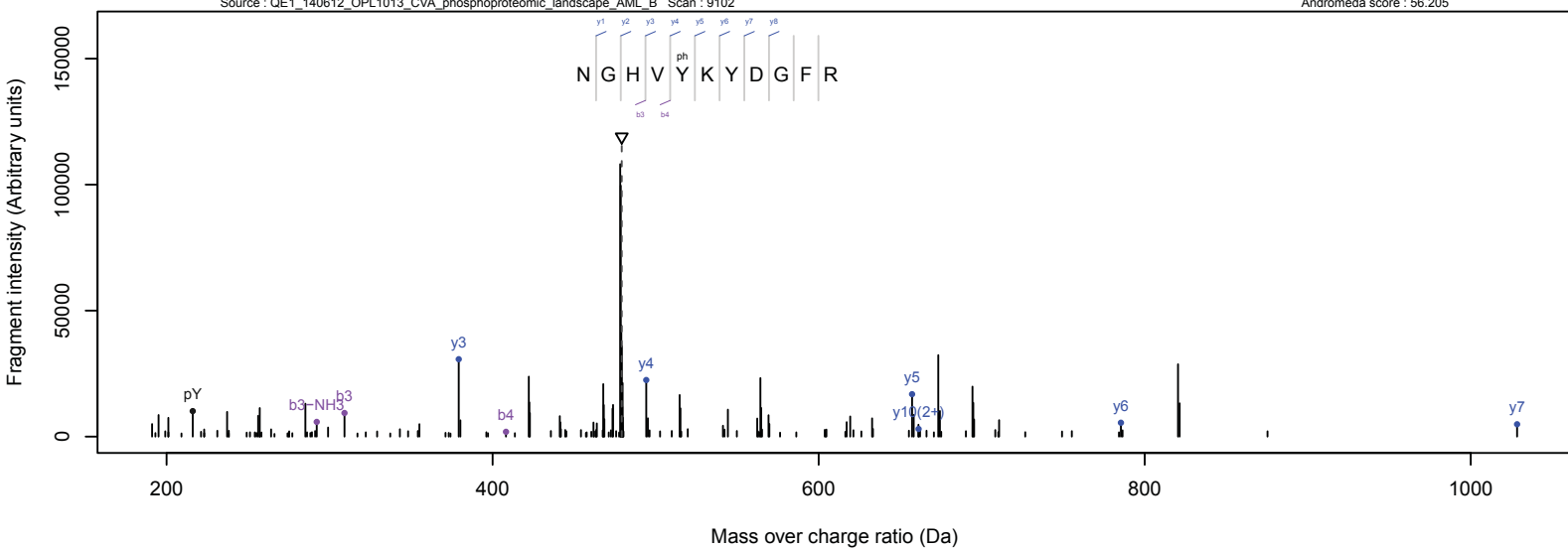
Andromeda score : 56.205



## NGHVY(ph)KYDGF-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9102

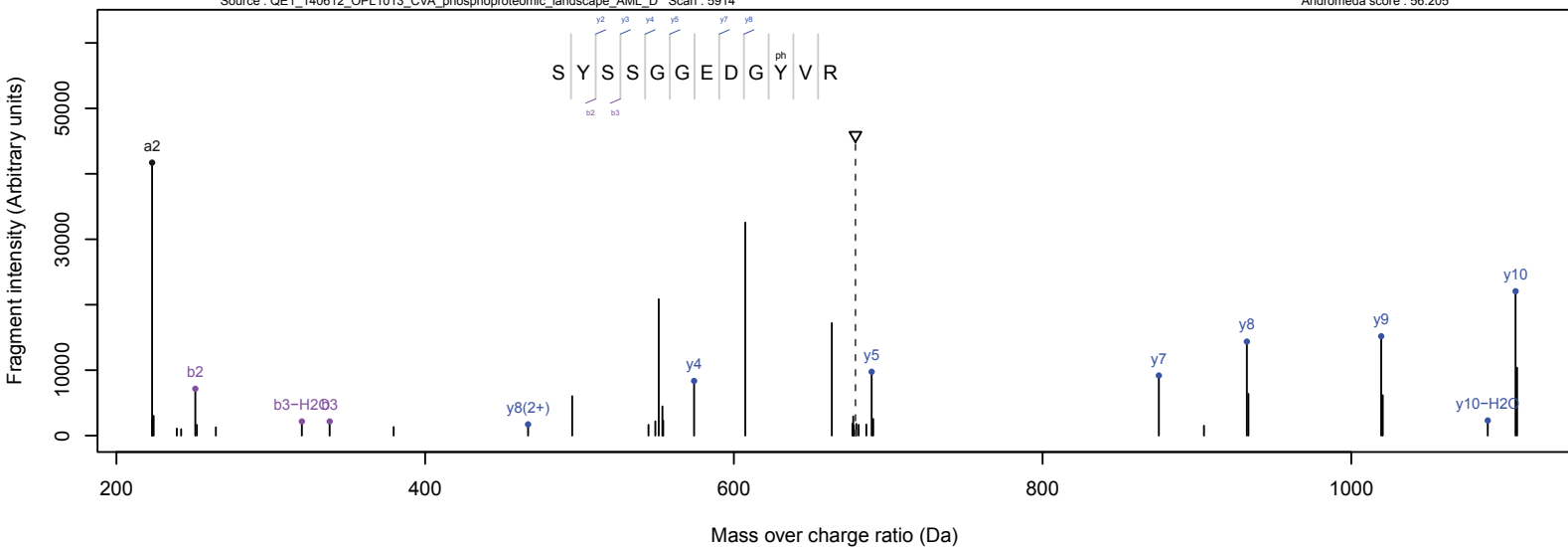
Andromeda score : 56.205



**SYSSGGEDGY(ph)VR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 5914

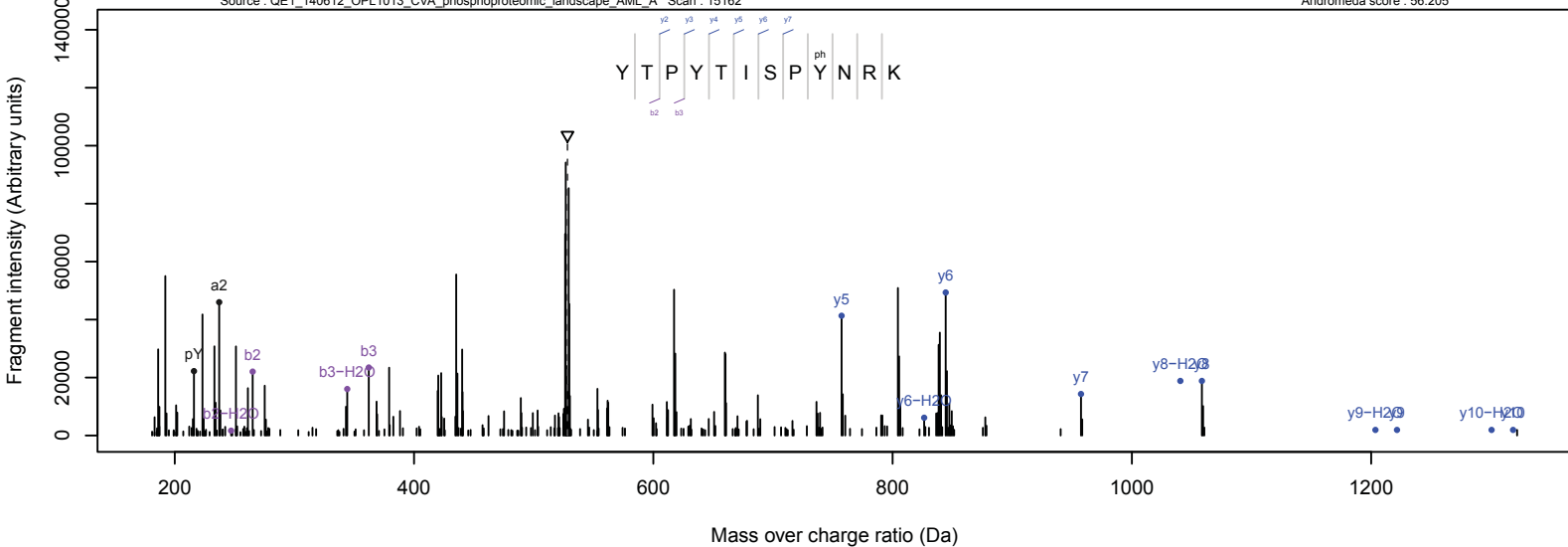
Andromeda score : 56.205



**YTPYTISPY(ph)NRK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 15162

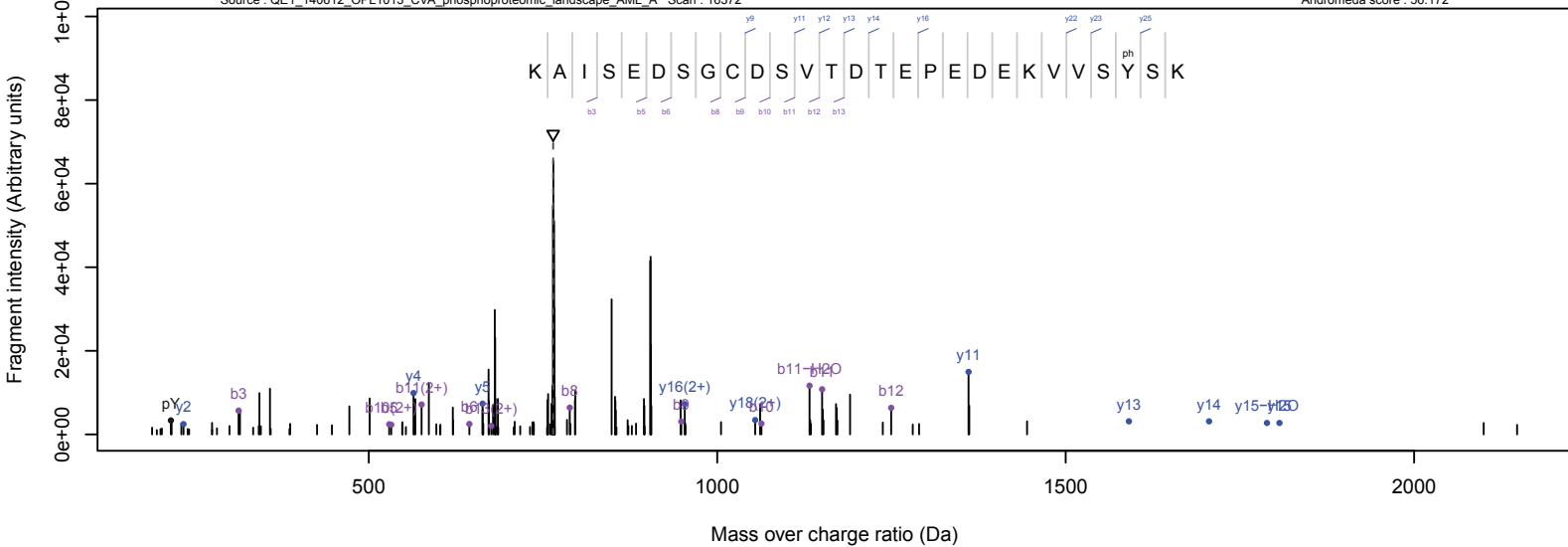
Andromeda score : 56.205

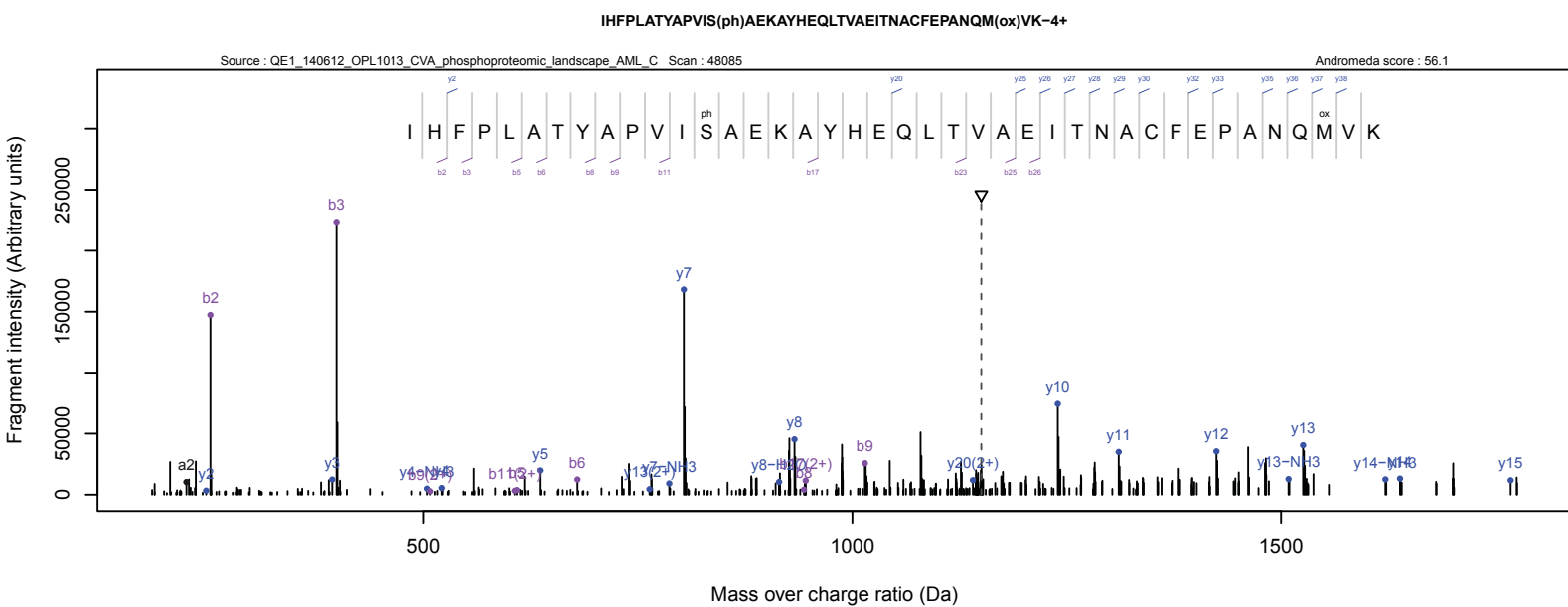
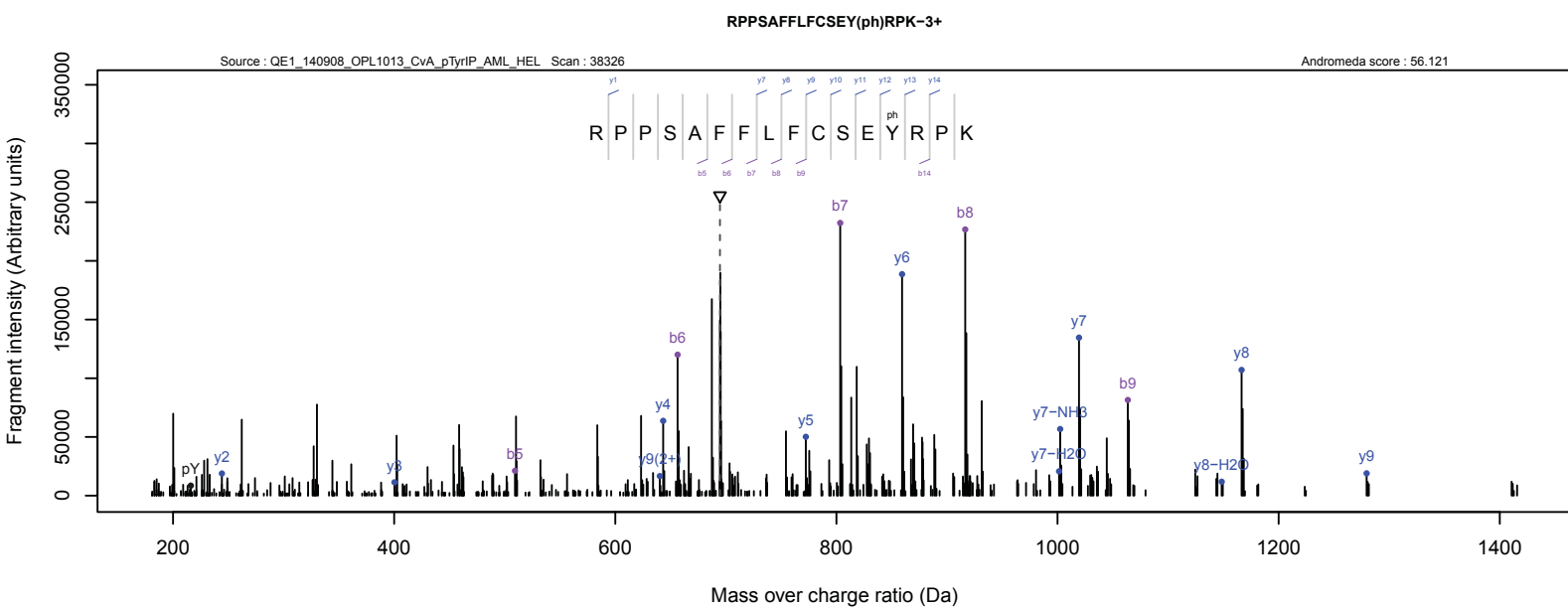
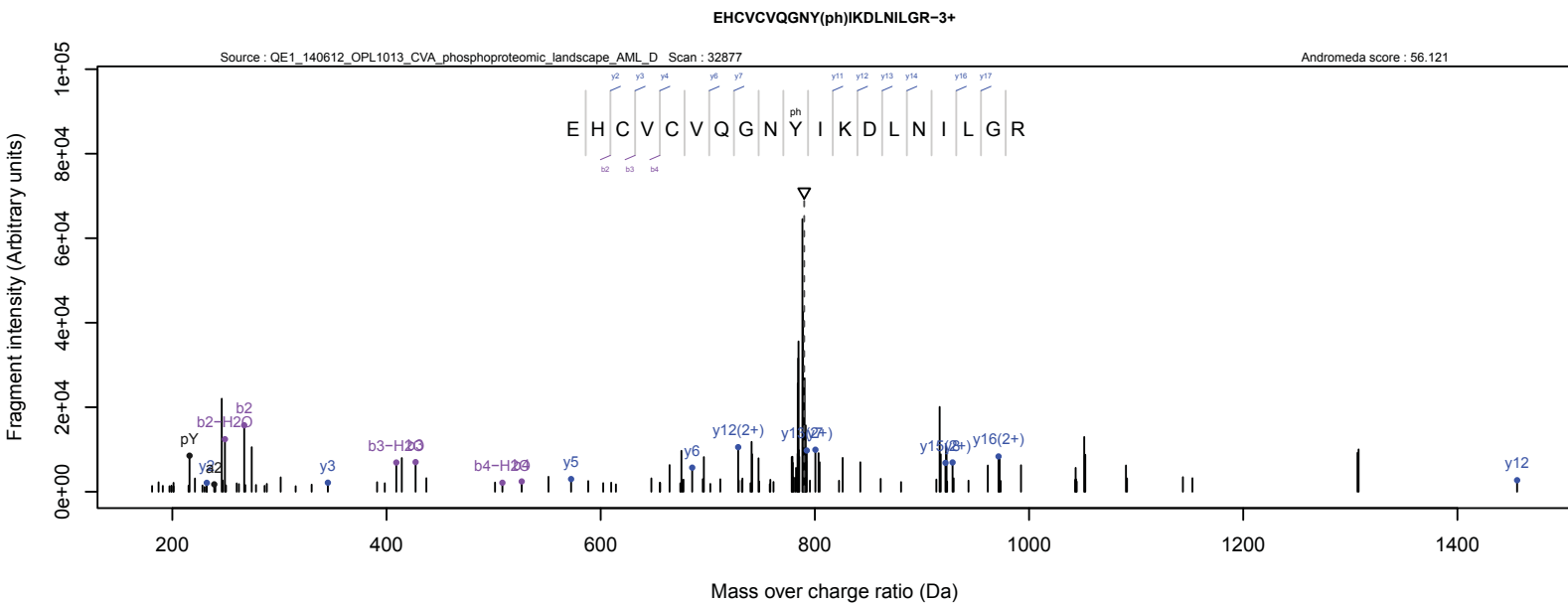


**KAISEDGCVTDTEPEDEKVVSY(ph)SK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16372

Andromeda score : 56.172

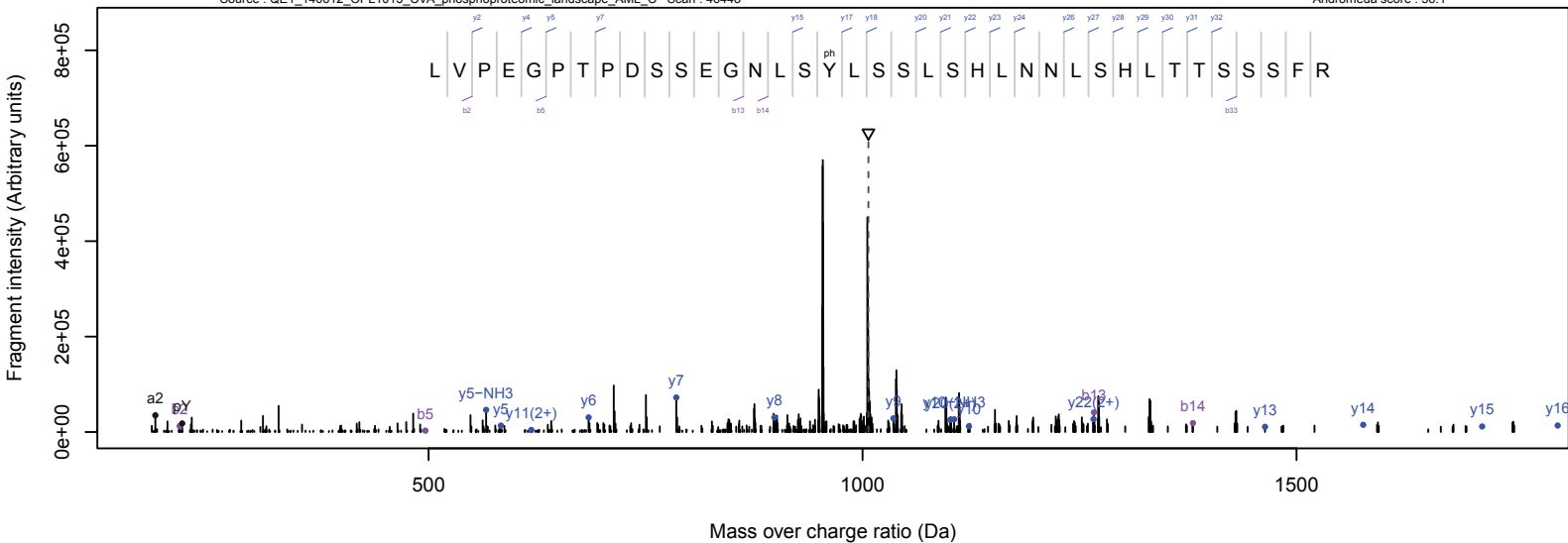




LVPEGPTDSSEGNLS(ph)LSSLNLSLHLLTSSSFR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46448

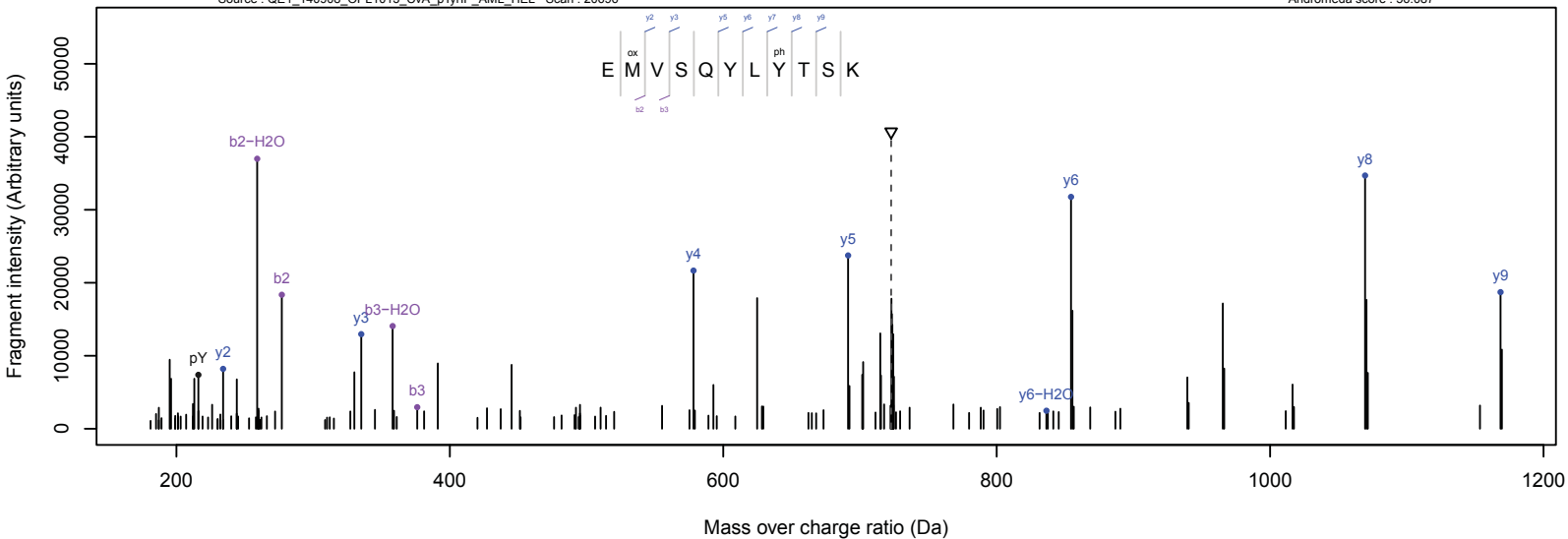
Andromeda score : 56.1



EM(ox)VSQYLY(ph)TSK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 20696

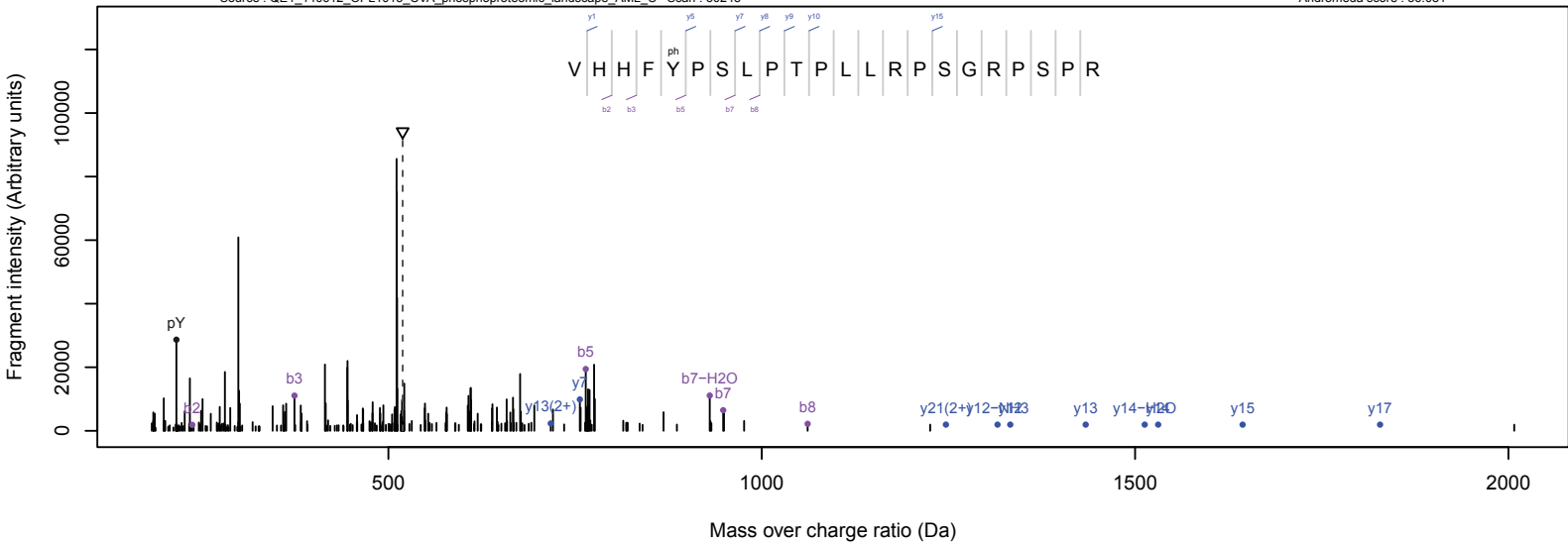
Andromeda score : 56.087



VHHFY(ph)PSLPTLLRPSGRPSR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30248

Andromeda score : 56.081

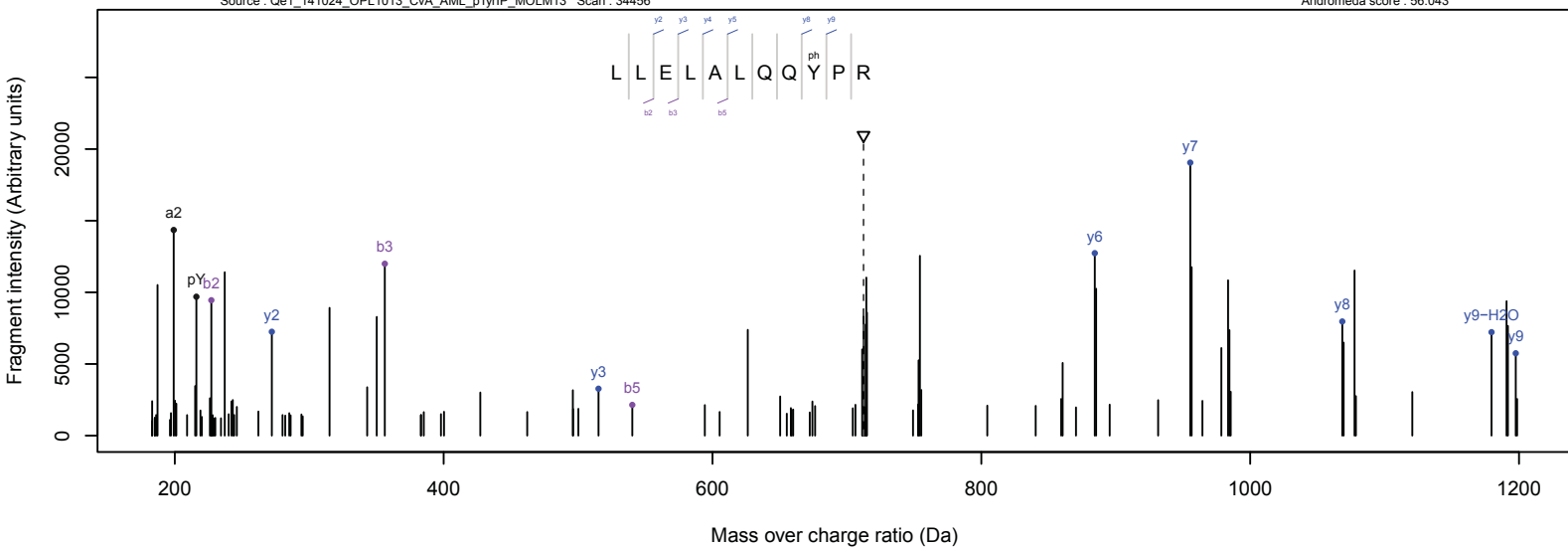




## LLELALQY(ph)PR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 34456

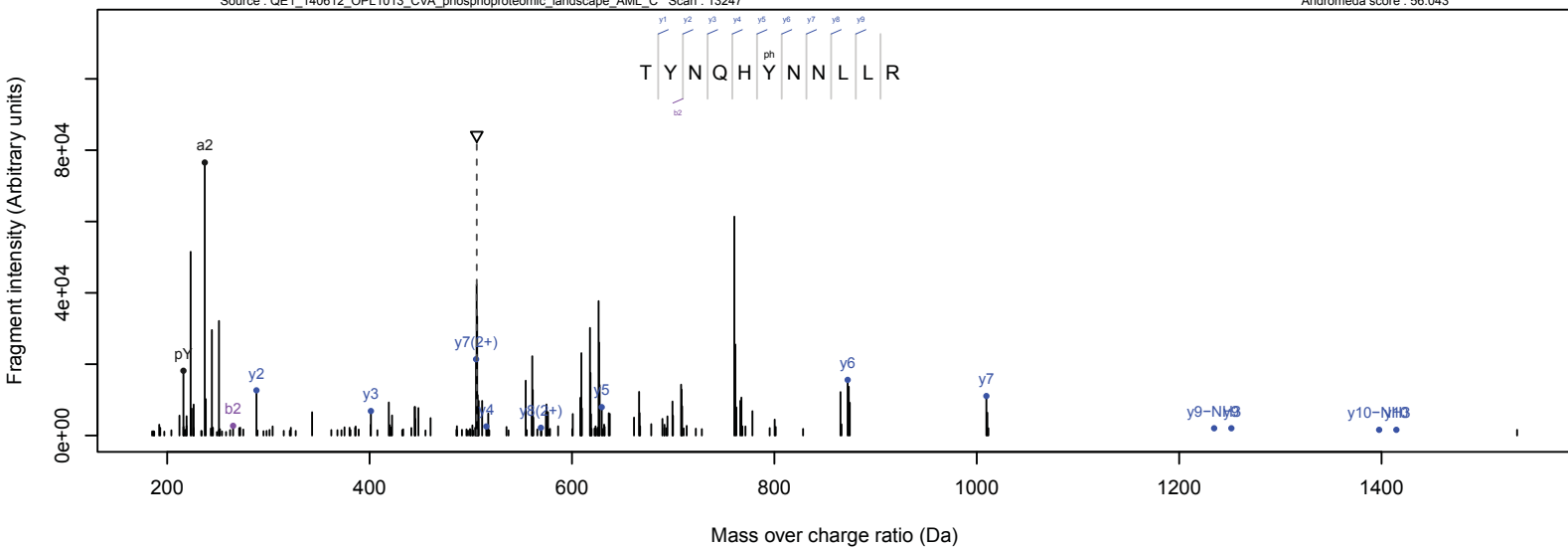
Andromeda score : 56.043



## TYNQHY(ph)NNLLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13247

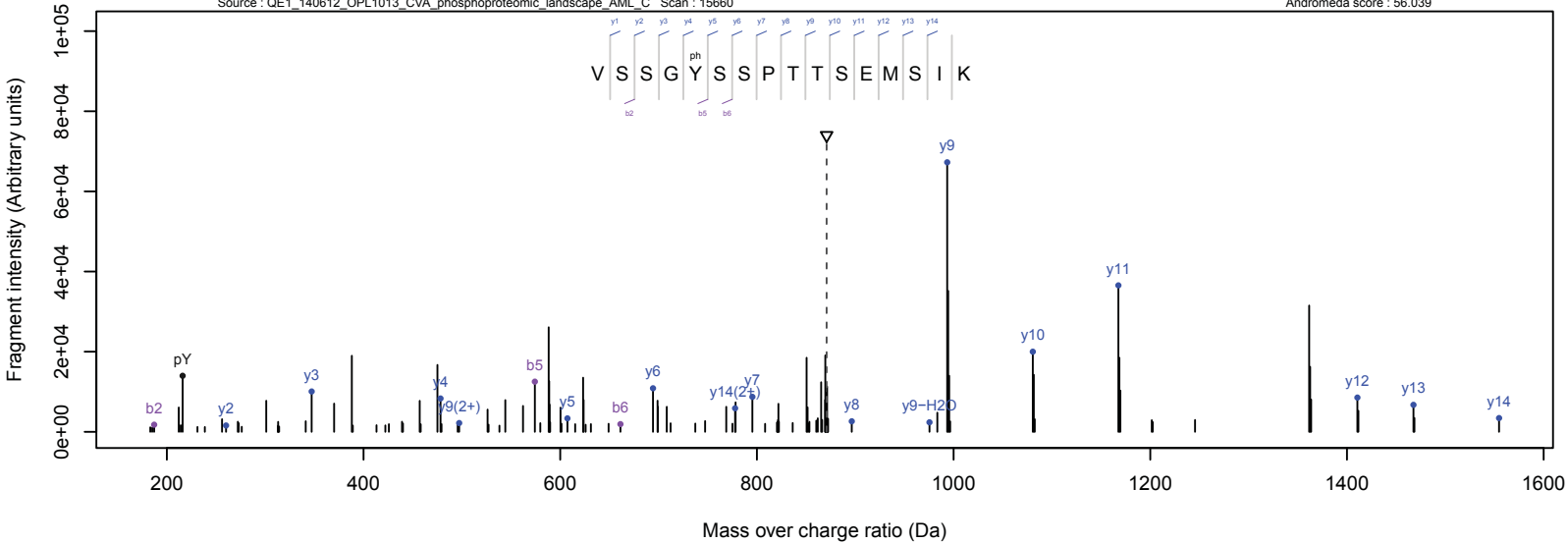
Andromeda score : 56.043



## VSSGY(ph)SSPTTSEMSIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15660

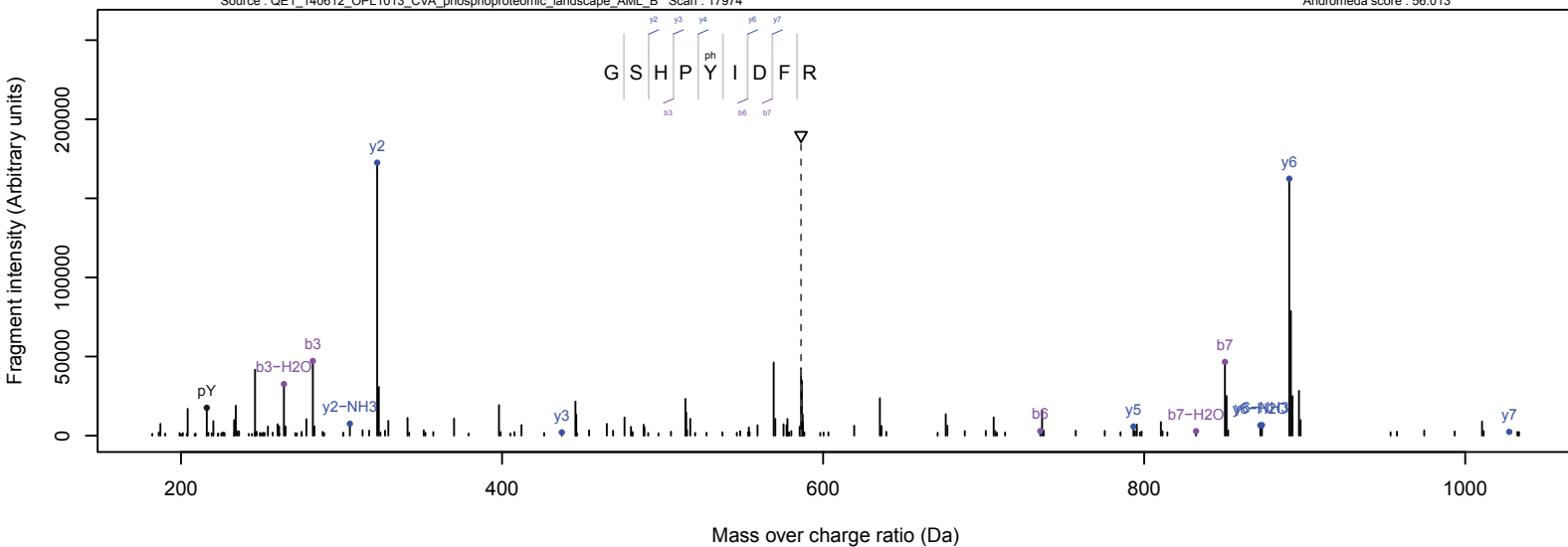
Andromeda score : 56.039



## GSHPY(ph)IDFR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 17974

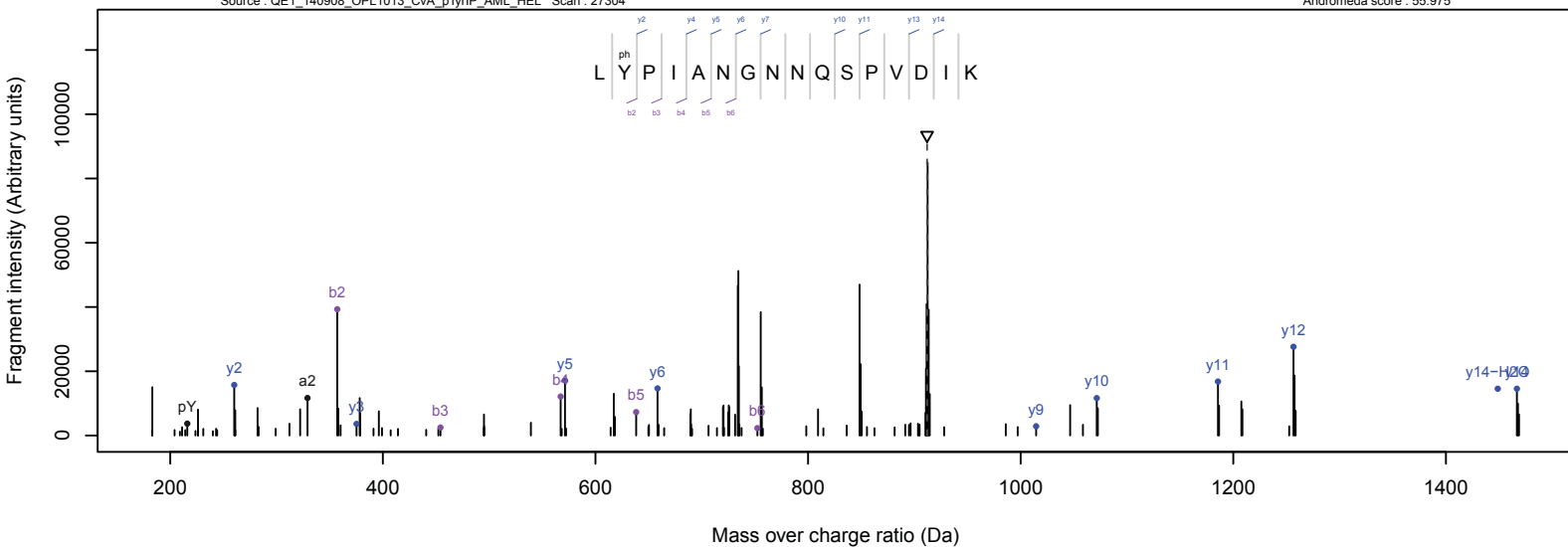
Andromeda score: 56.013



## LY(ph)PIANGNNQSPVDIK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 27304

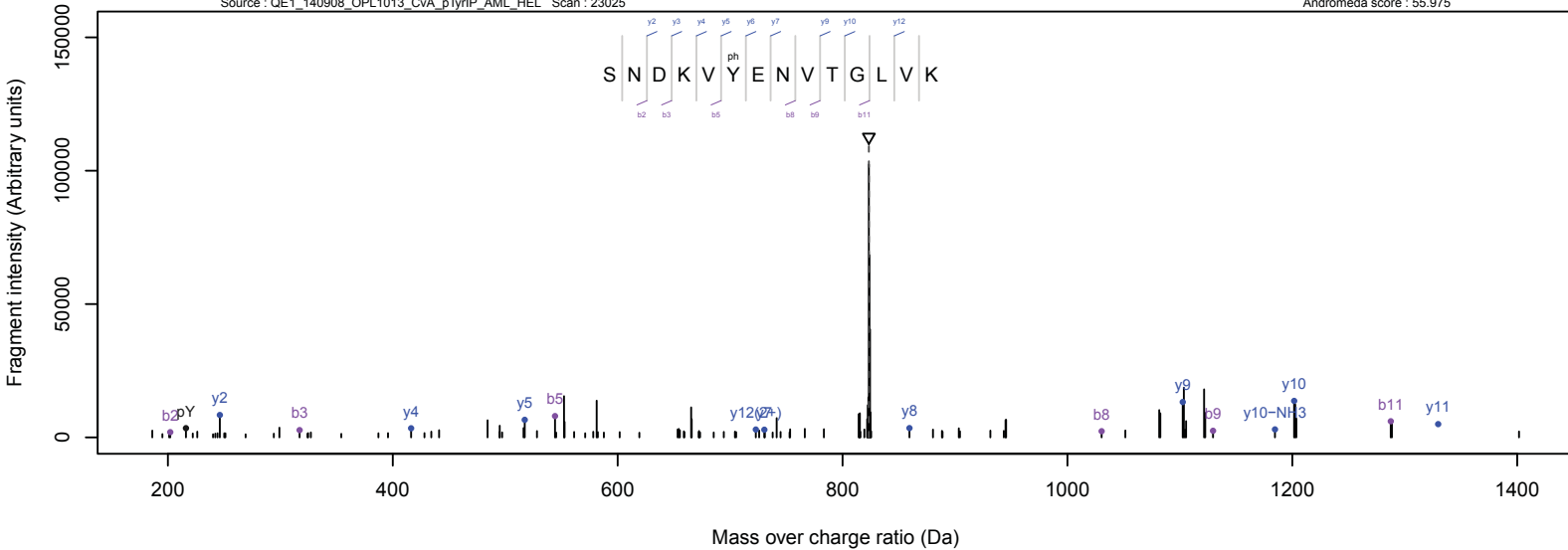
Andromeda score: 55.975



## SNDKVY(ph)ENVTLVK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 23025

Andromeda score: 55.975

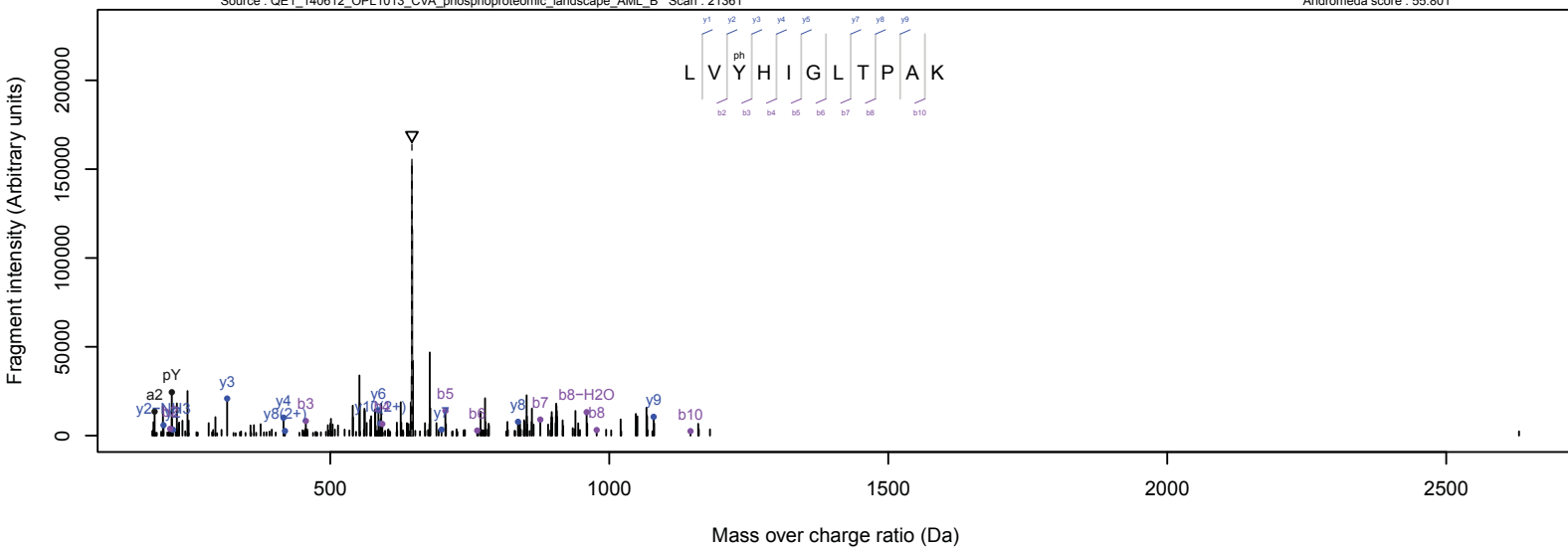




LVY(ph)HIGLTPAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21361

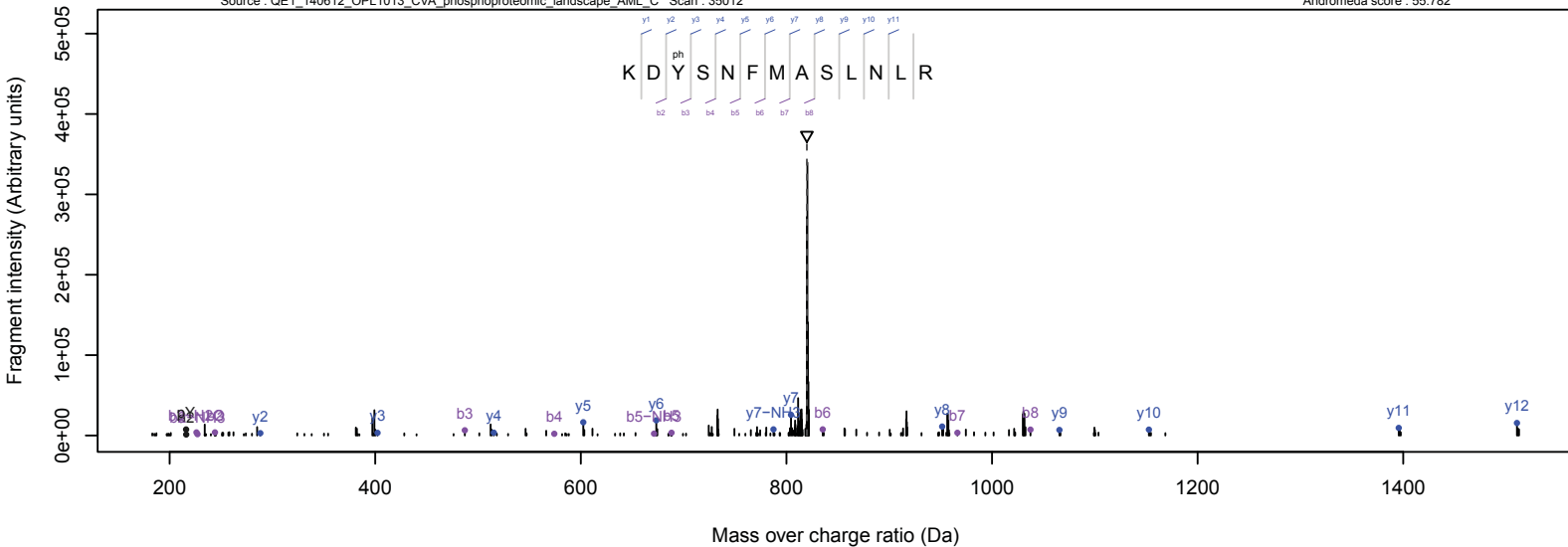
Andromeda score : 55.801



KDY(ph)SNFMASLNLRL-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35012

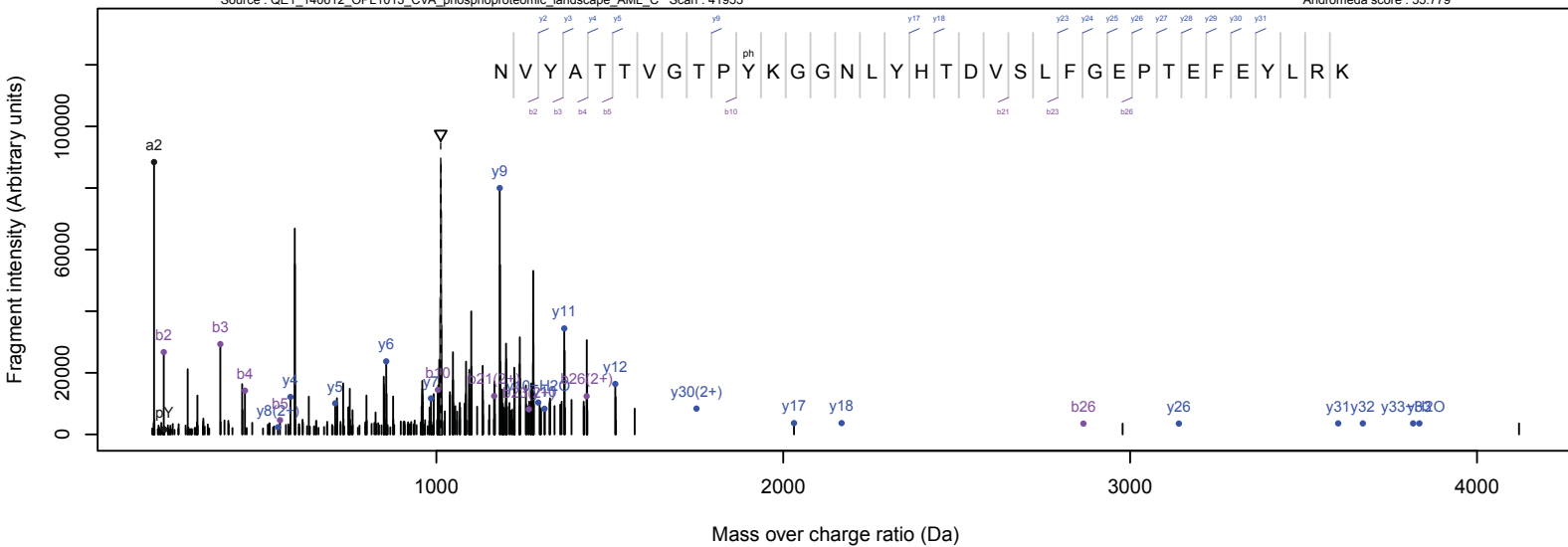
Andromeda score : 55.782



NVYATTVGTPY(ph)KGGNLYHTDVSLEFGEPTFEYLRK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41955

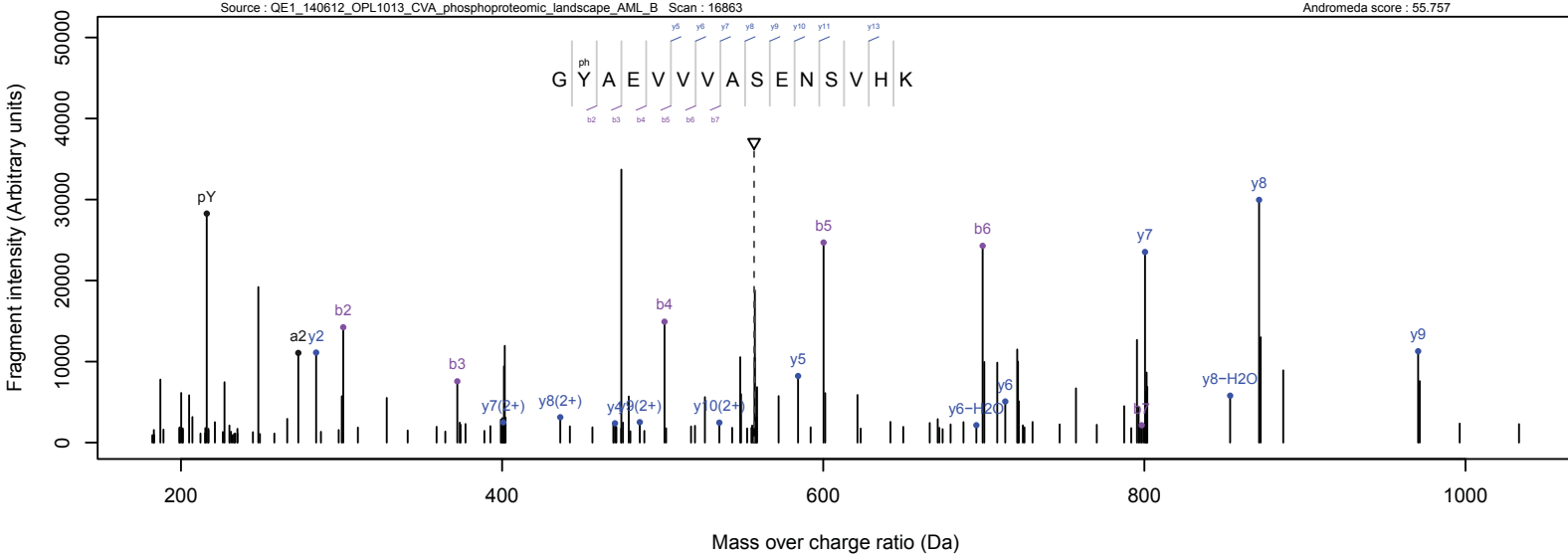
Andromeda score : 55.779



## GY(ph)AEVVVASENSVHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16863

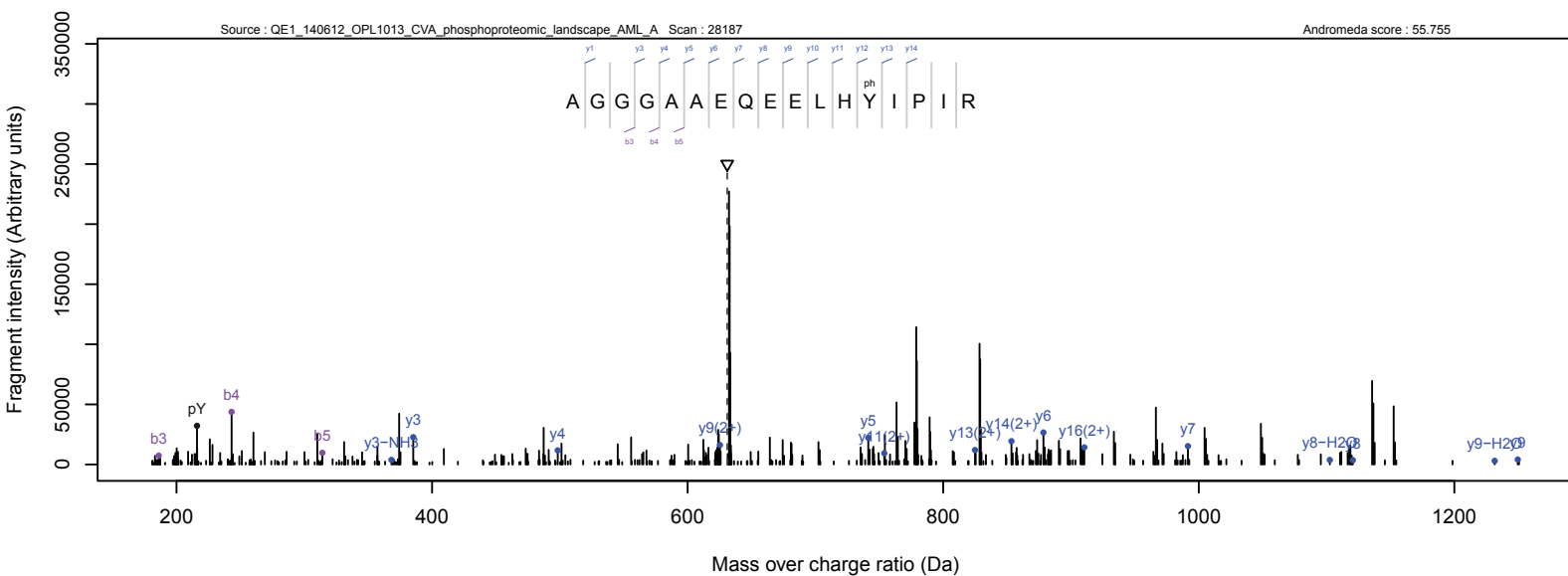
Andromeda score : 55.757



## AGGGAAEQEELHY(ph)IPIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 28187

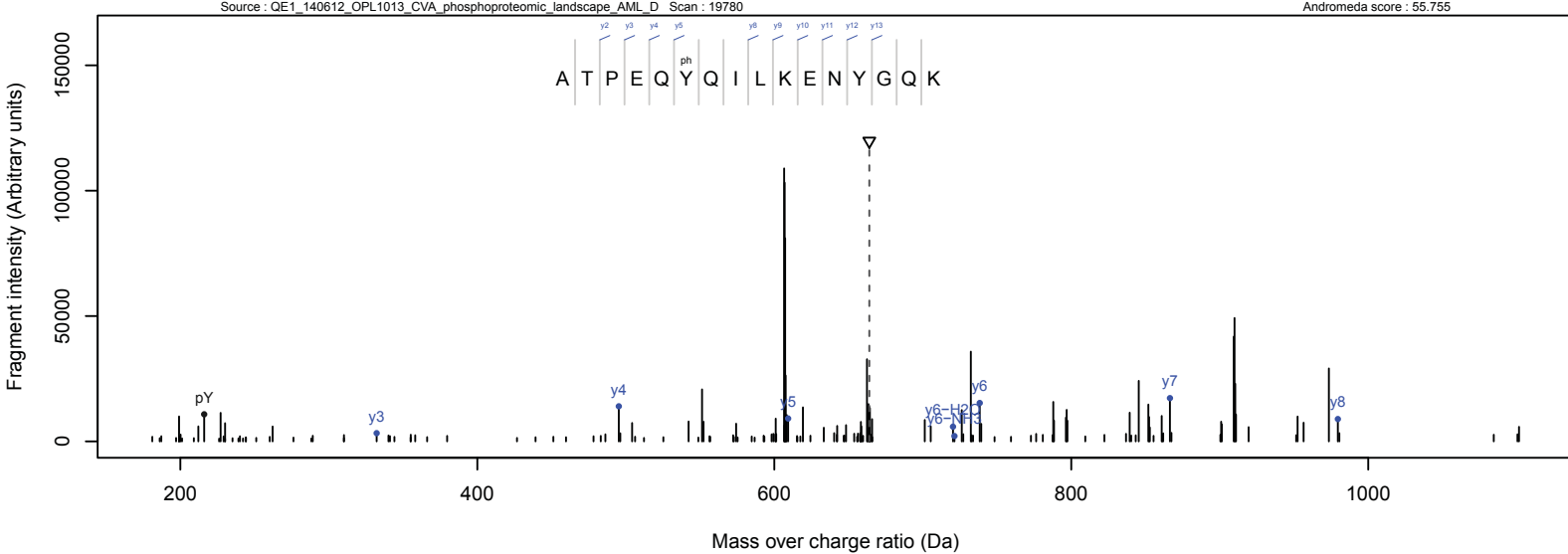
Andromeda score : 55.755



## ATPEQY(ph)ILKENYGQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 19780

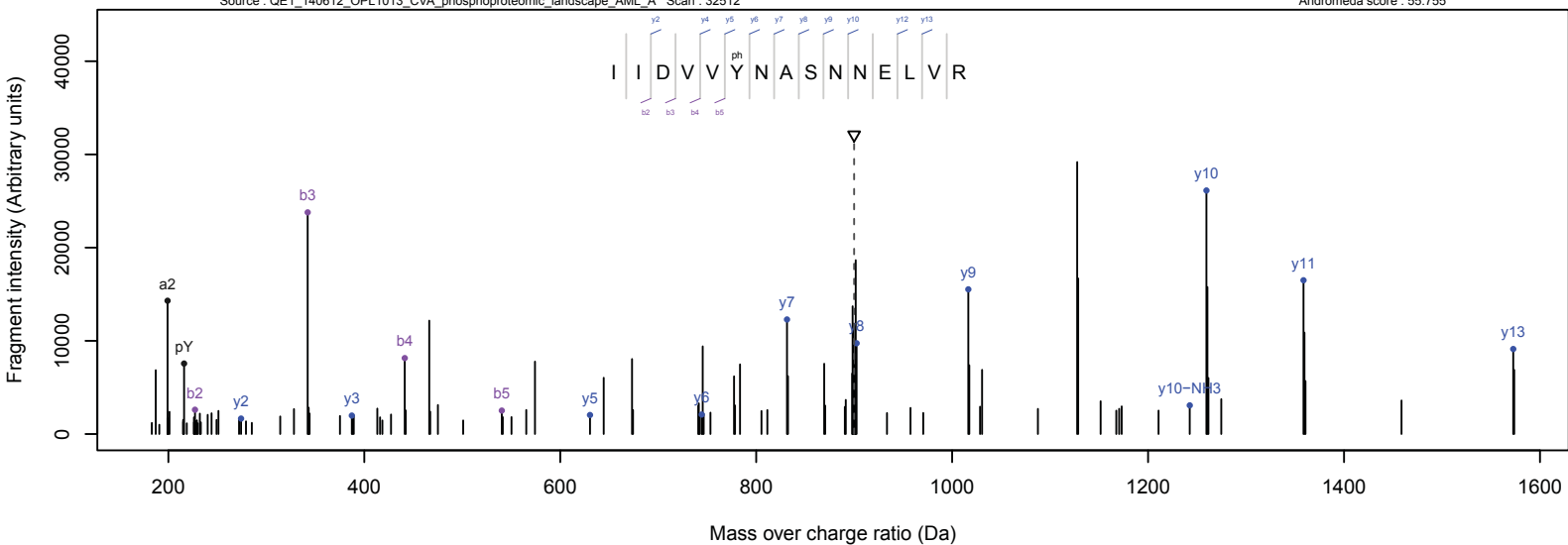
Andromeda score : 55.755



## IIDVVY(ph)NASNNELVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32512

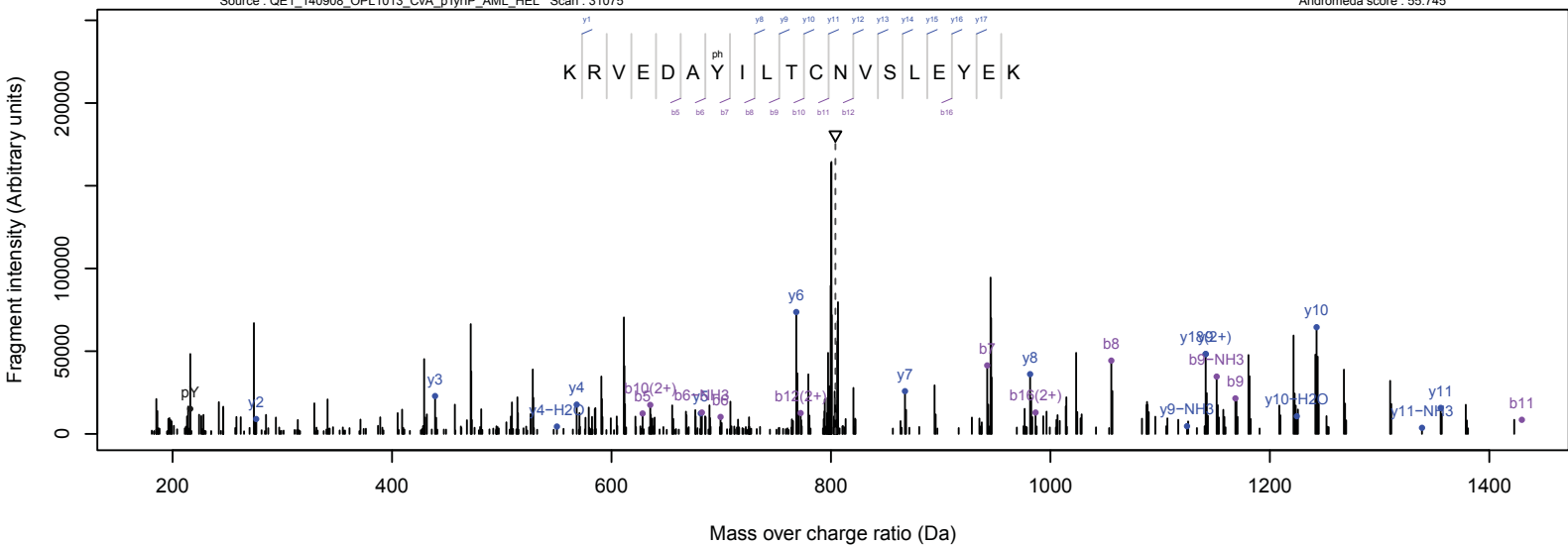
Andromeda score : 55.755



## KRVEDAY(ph)ILTCNVSLEYEK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31075

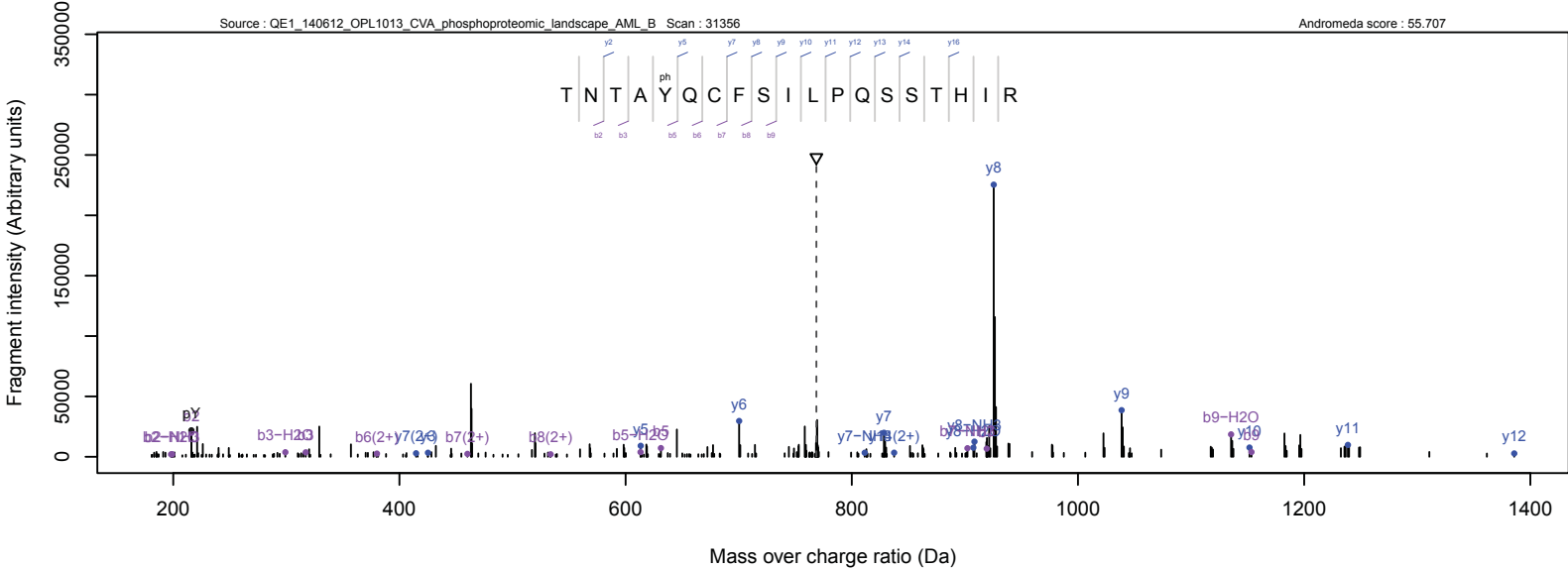
Andromeda score : 55.745



## TNTAY(ph)QCFSLPQSSTHIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31356

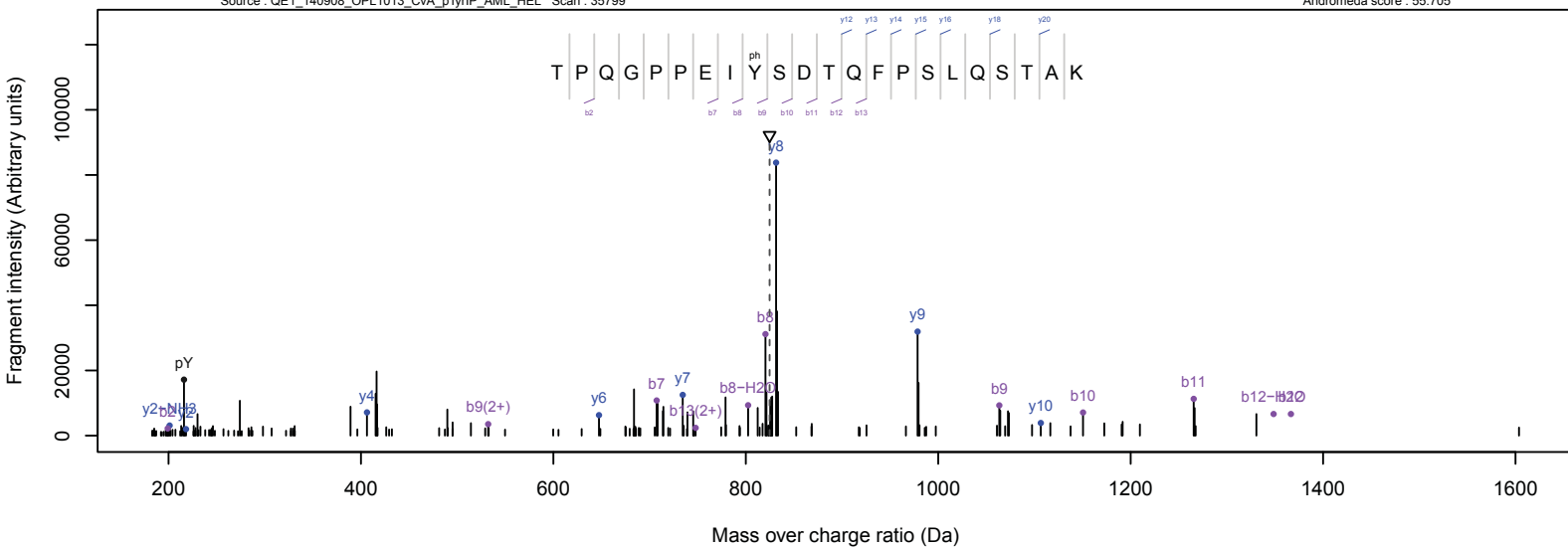
Andromeda score : 55.707



TPQGPPEIY(ph)SDTQFSLQSTAK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 35799

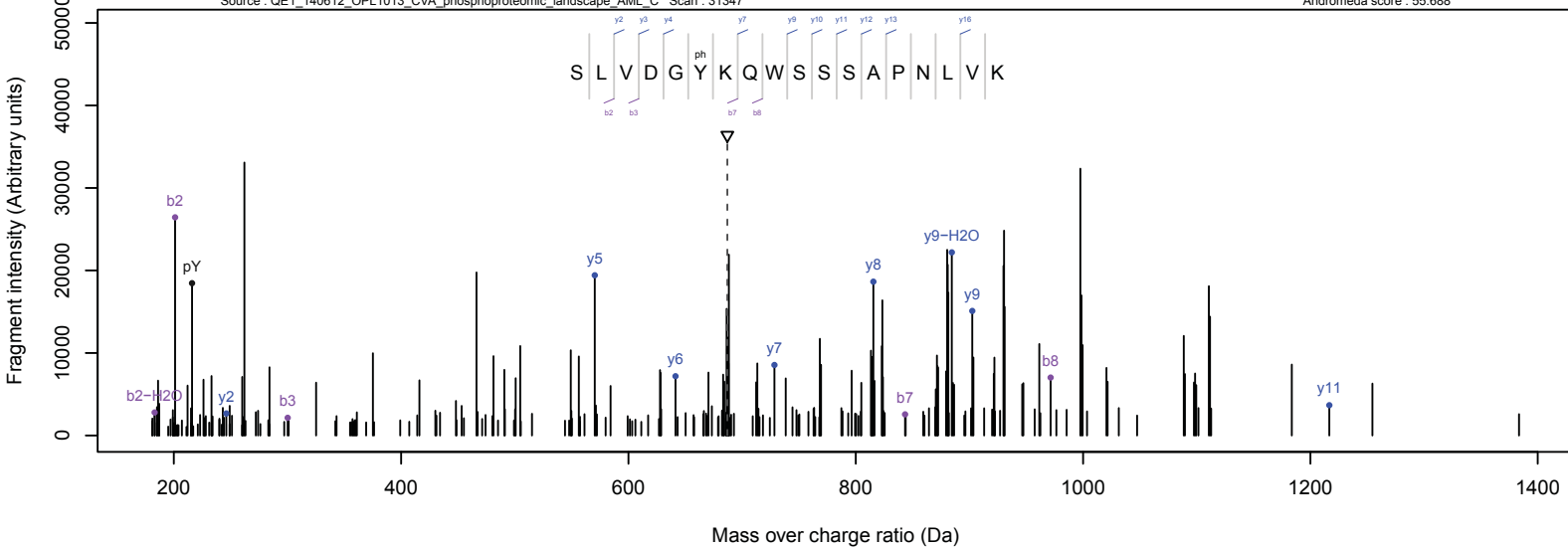
Andromeda score : 55.705



SLVDGY(ph)KQWSSAPNLVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31347

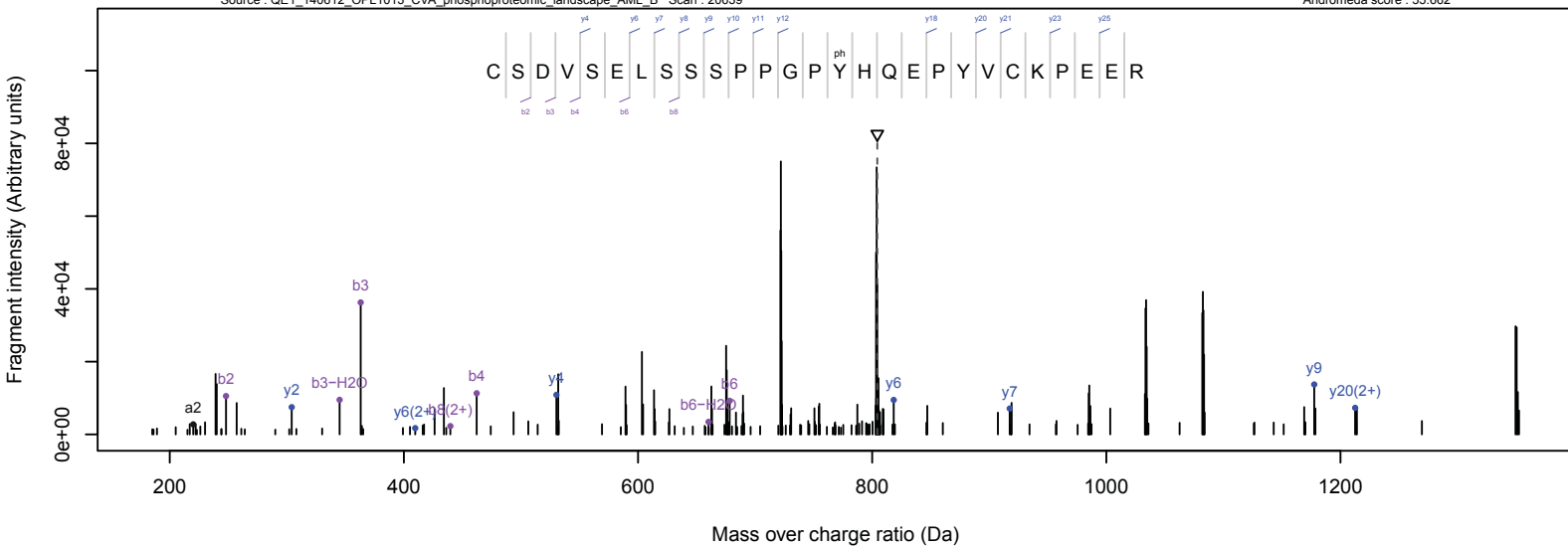
Andromeda score : 55.688



CSDVSELSSSPGYPYHQEPYVCKPEER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20639

Andromeda score : 55.662



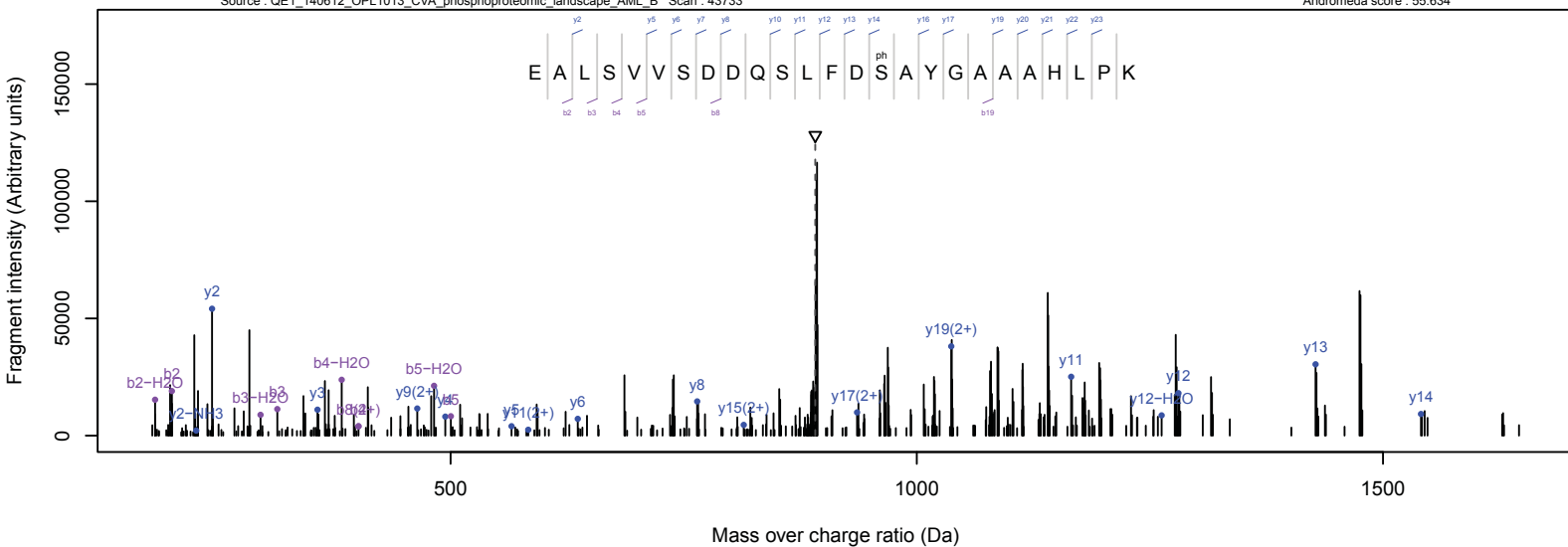




EALSVVSD DQSLFDS(ph)AYGAAHLPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 43733

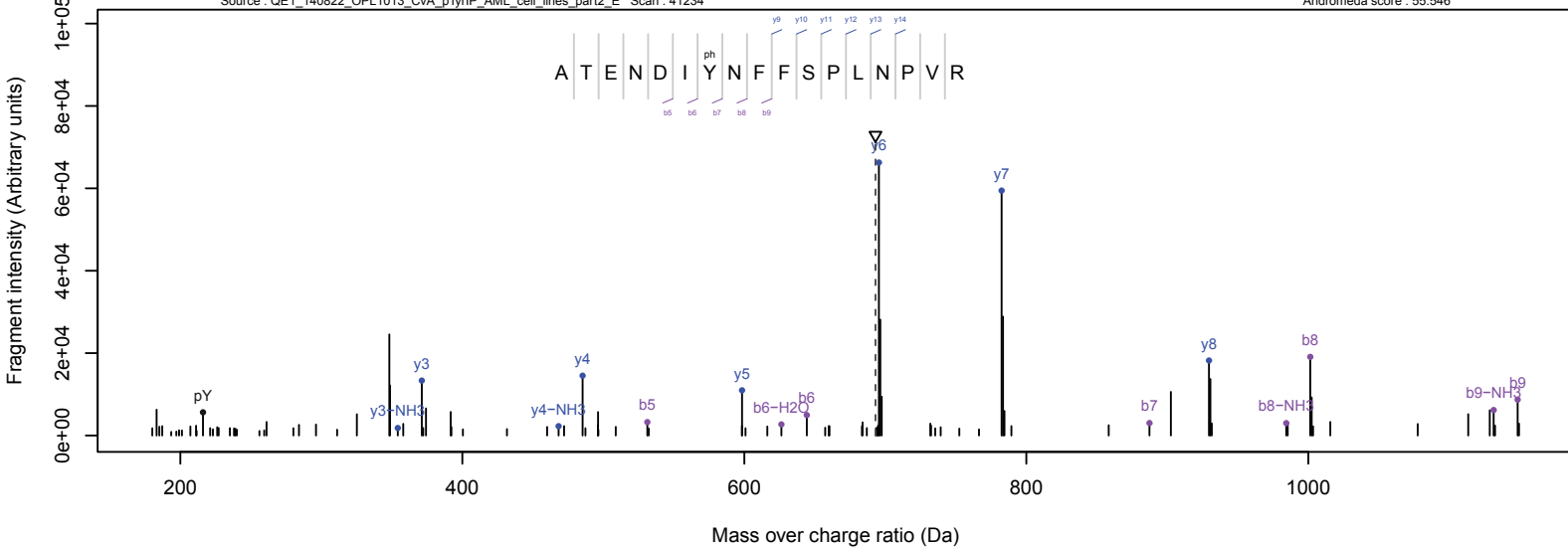
Andromeda score : 55.634



ATENDIY(ph)NFFSPLNPVR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 41234

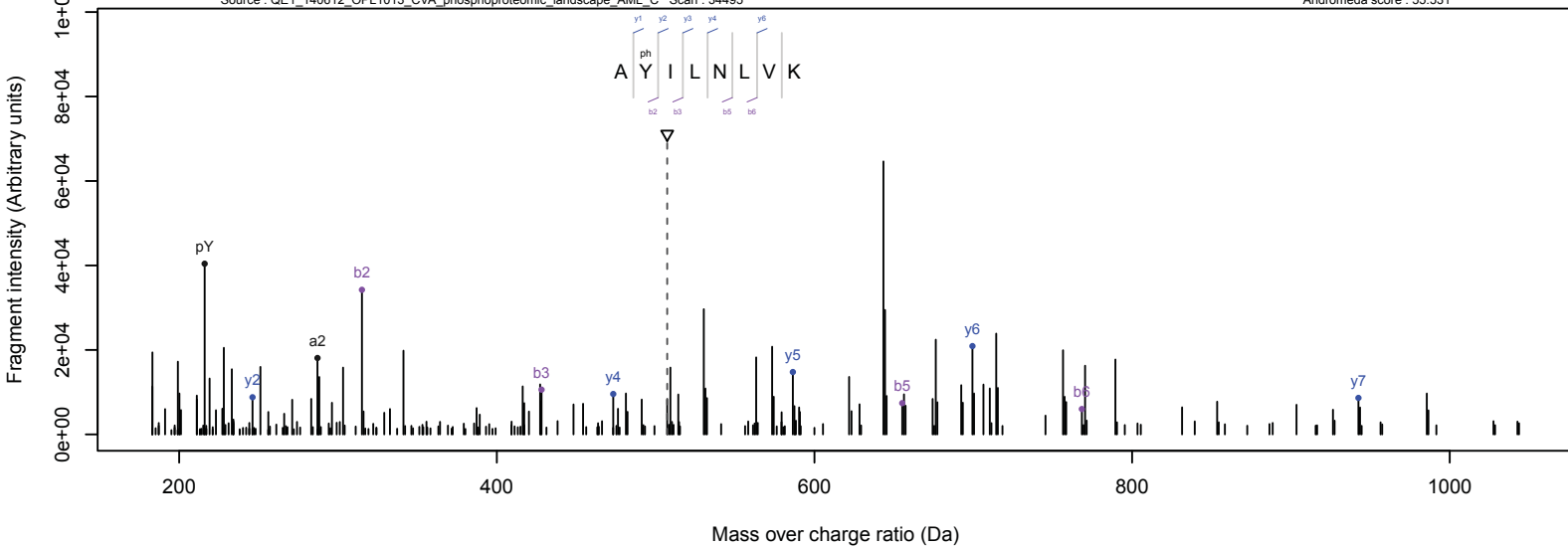
Andromeda score : 55.546



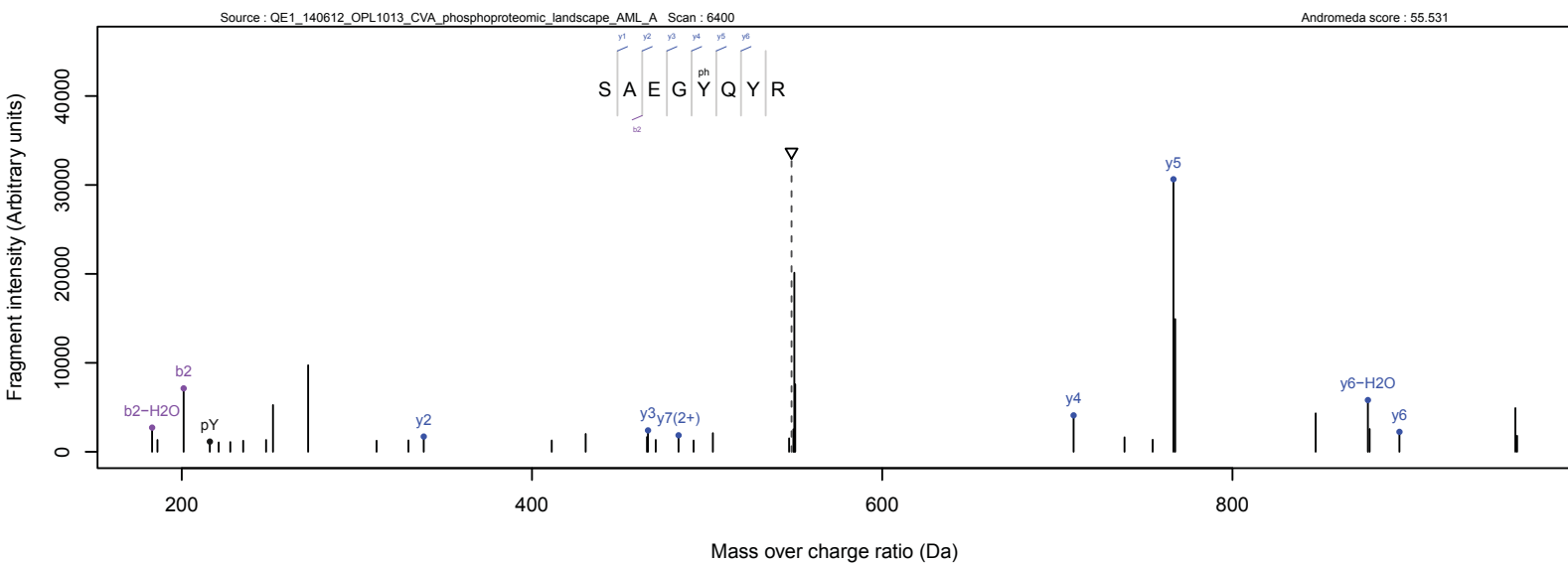
AY(ph)ILNLVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34495

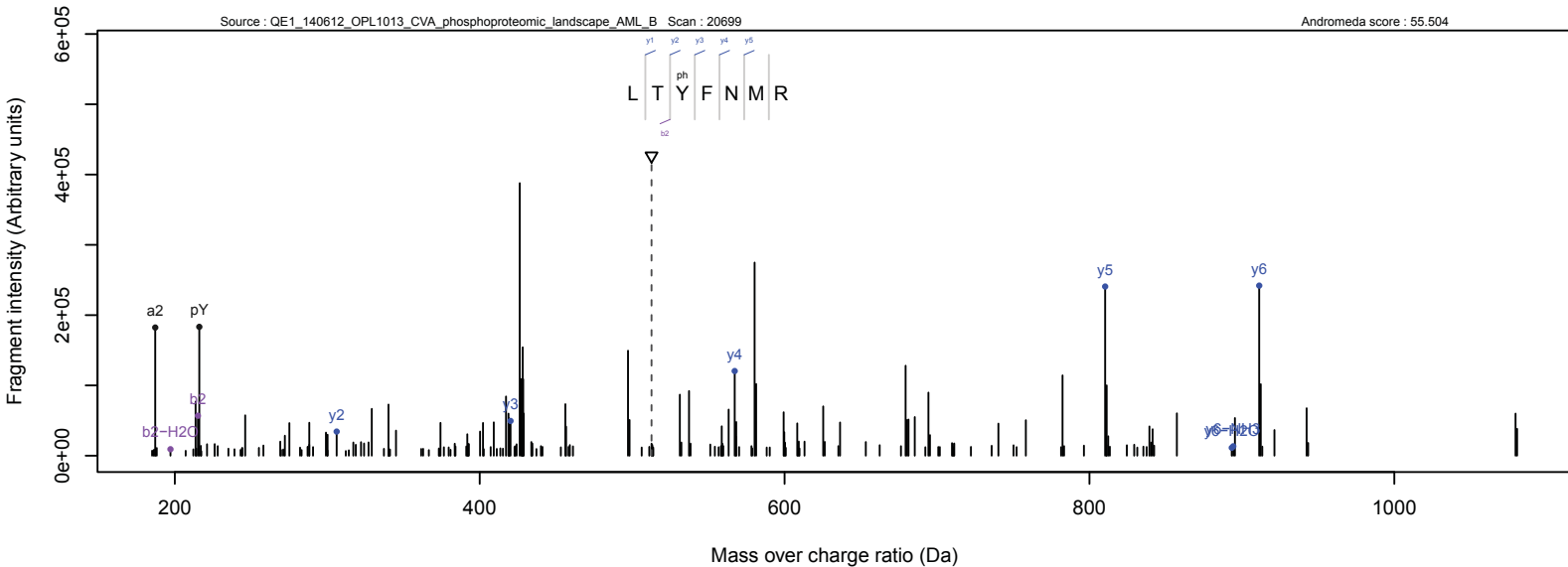
Andromeda score : 55.531



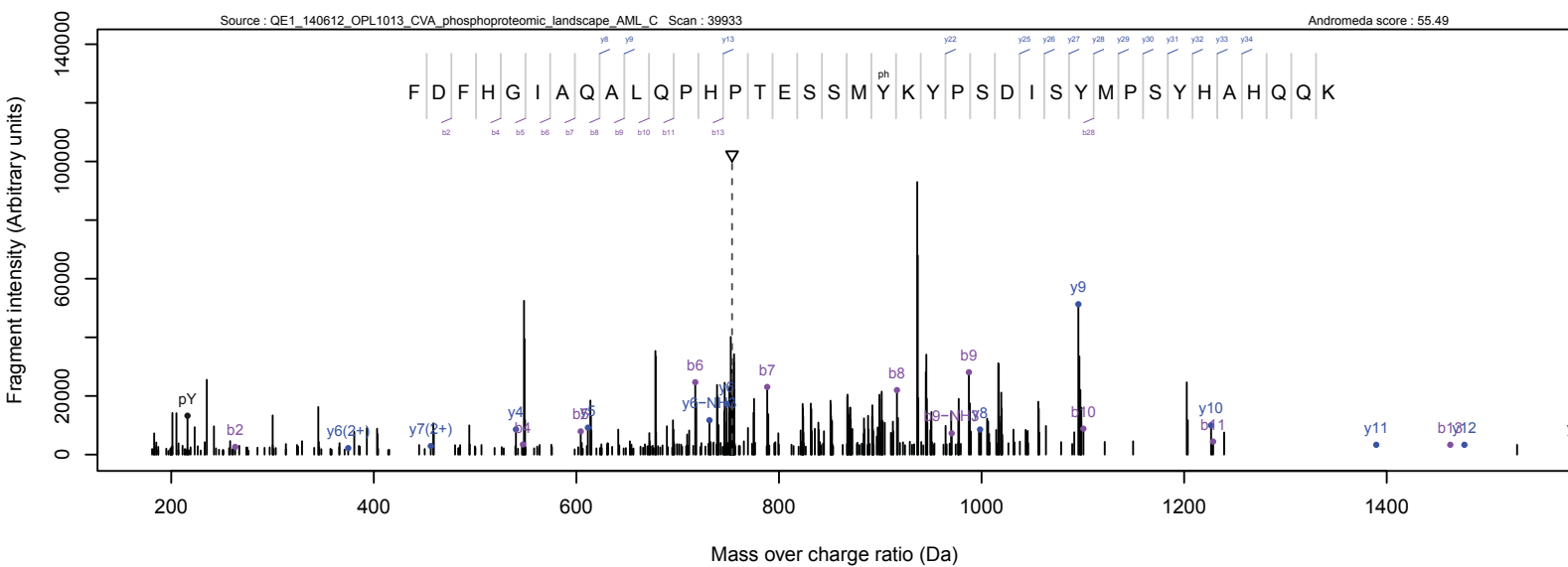
(ac)SAEGY(ph)QYR-2+



LTY(ph)FNMR-2+



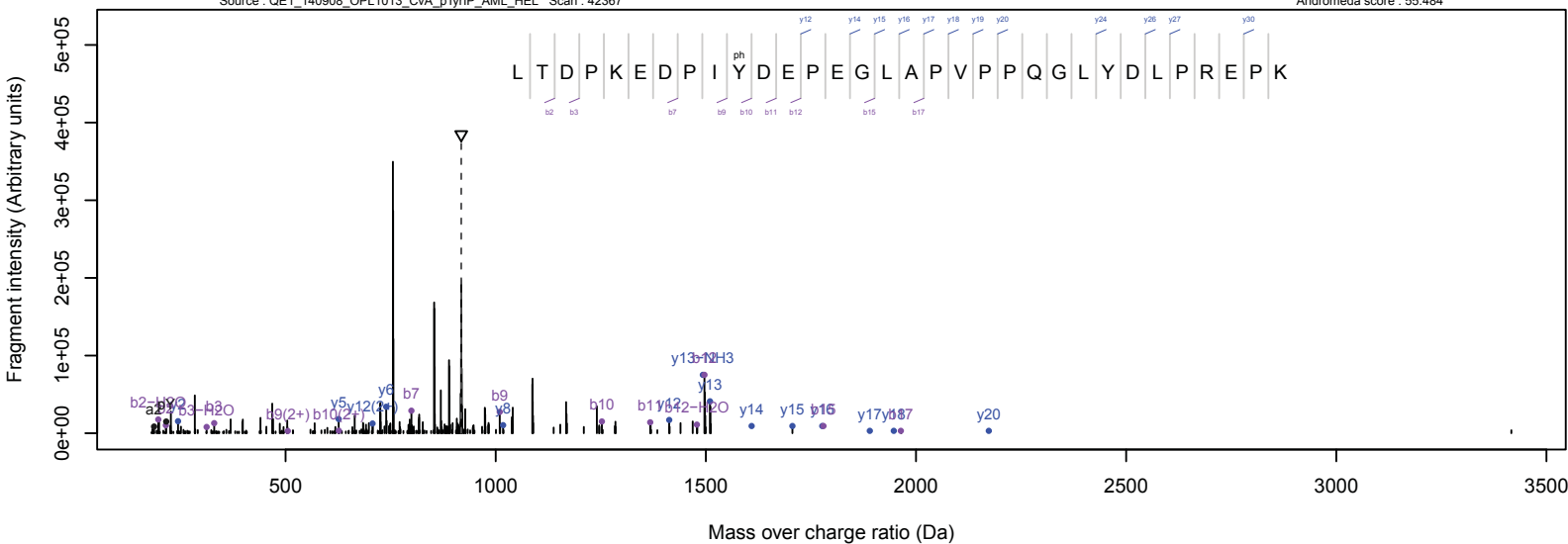
FD FHGIAQLQPHPTESMYKYPSPDISYMP SYH AHQK-6+



**LTDPKEDPIYDEPEGLAPVPPQGLYDLPREP-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 42367

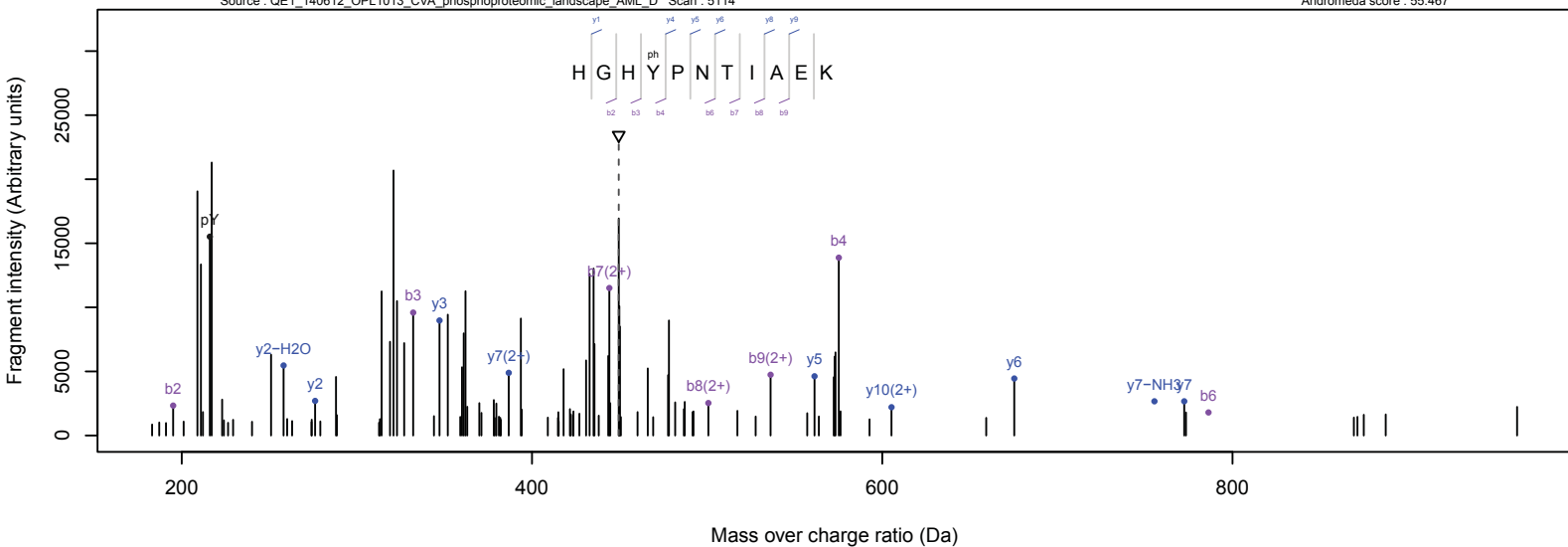
Andromeda score : 55.484



**HGHY(ph)PNTIAEK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 5114

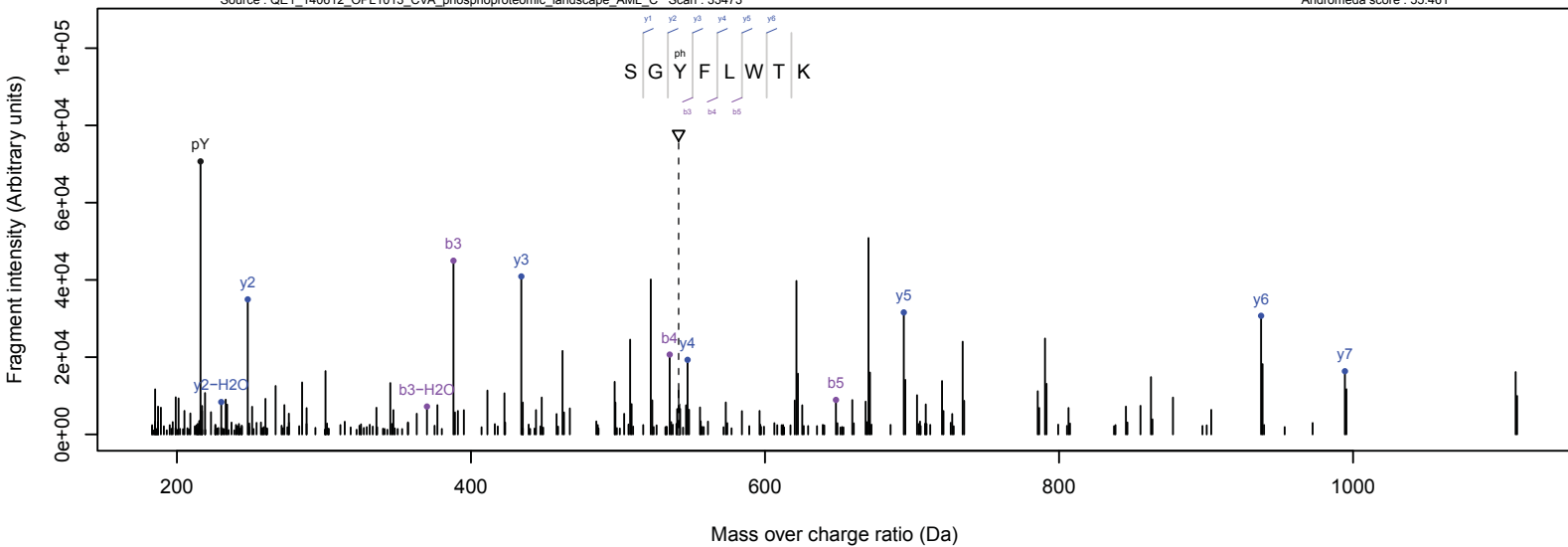
Andromeda score : 55.467



**SGY(ph)FLWTK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33473

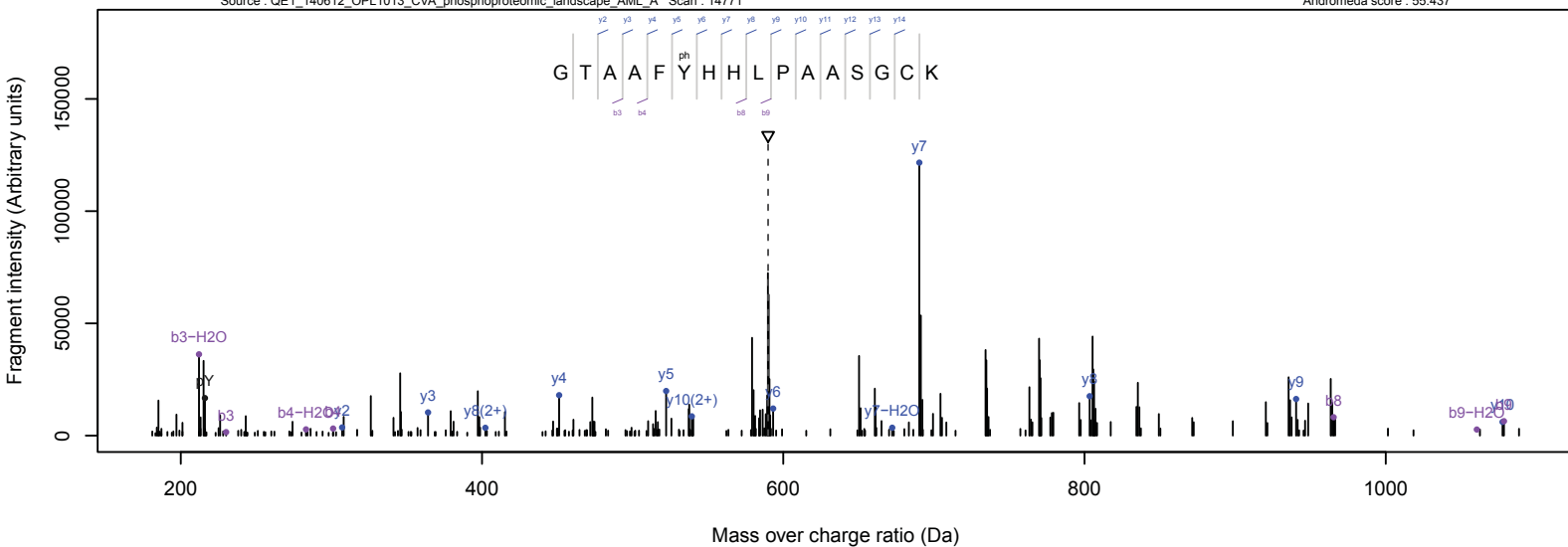
Andromeda score : 55.461



## GTAAFY(ph)HHLPAASGCK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 14771

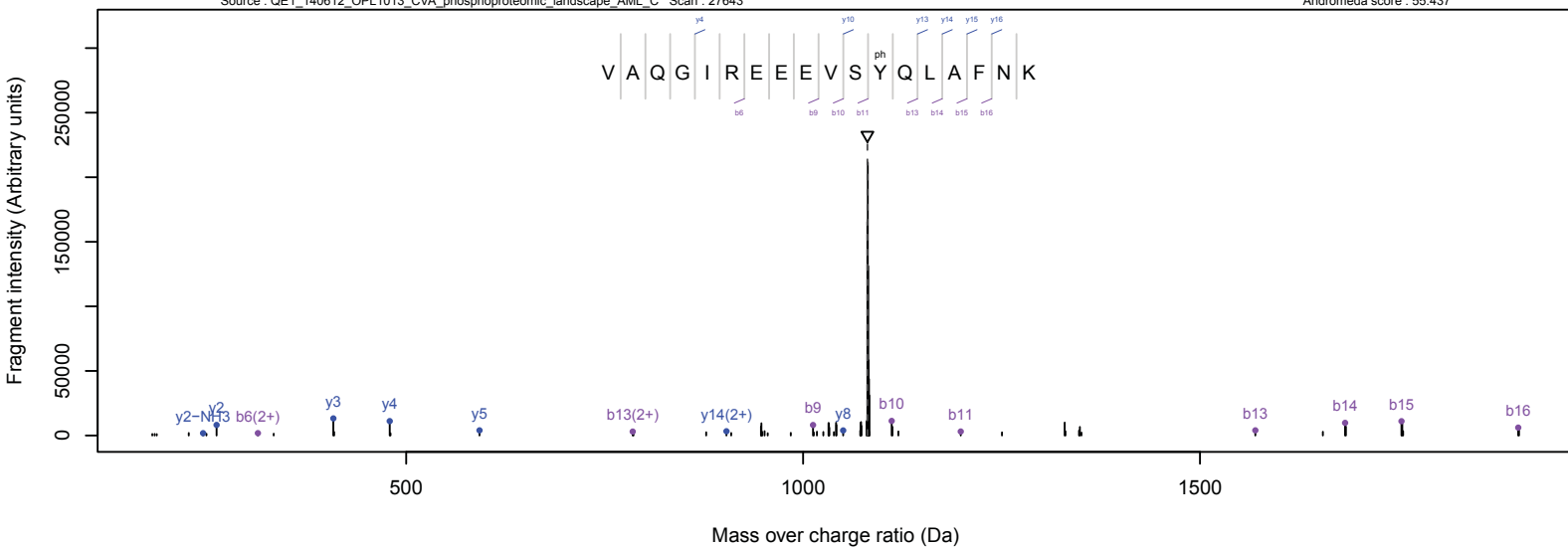
Andromeda score: 55.437



## VAQGIRREEVSY(ph)QLAFNK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 27643

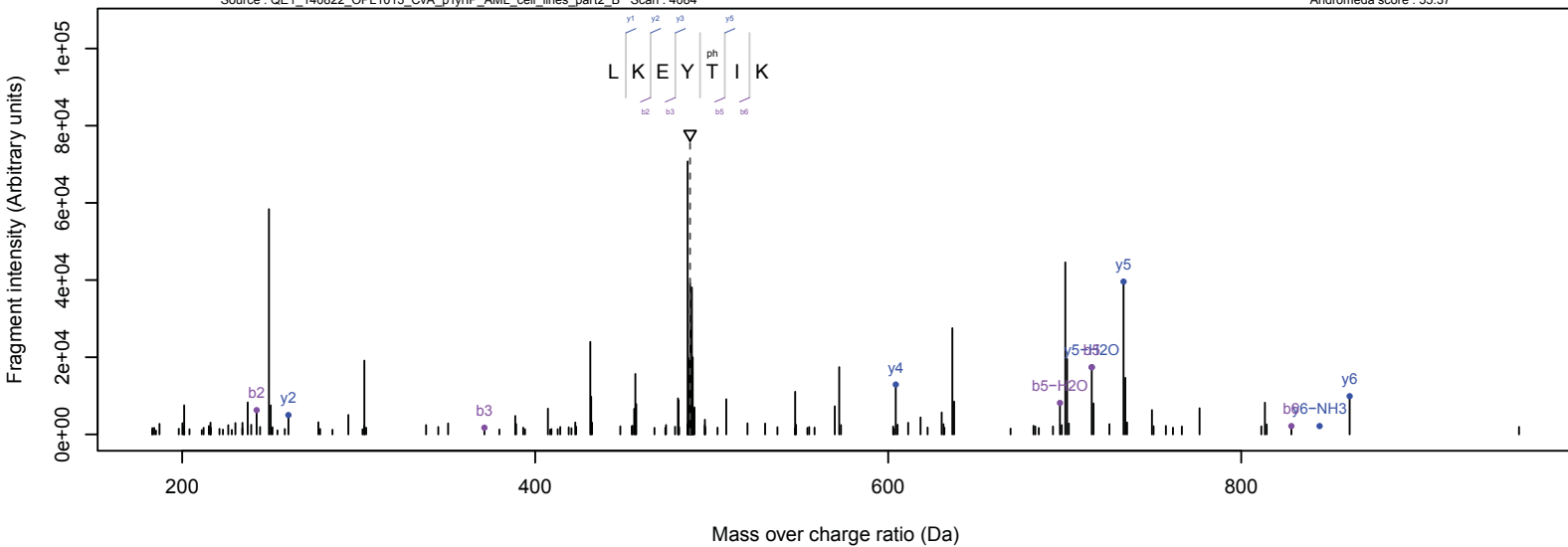
Andromeda score: 55.437



## LKEYT(ph)IK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 4084

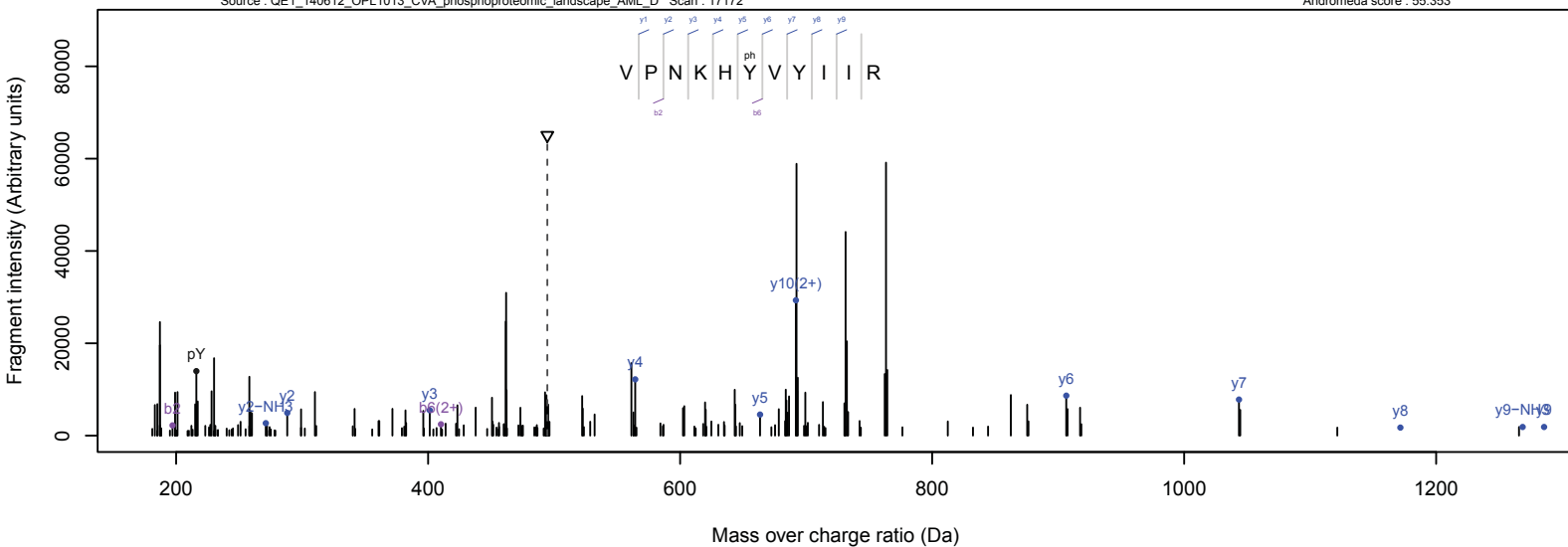
Andromeda score: 55.37



VPNKHY(ph)VYIIR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 17172

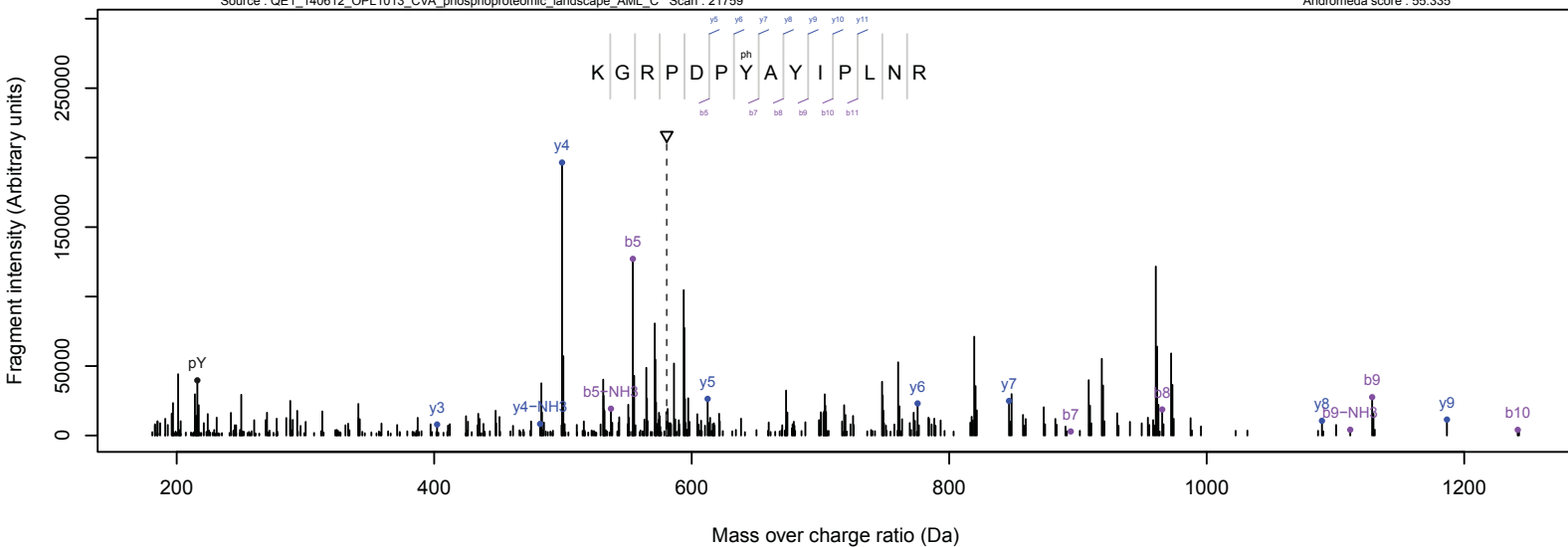
Andromeda score : 55.353



KGRDPY(ph)AYIPLNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21759

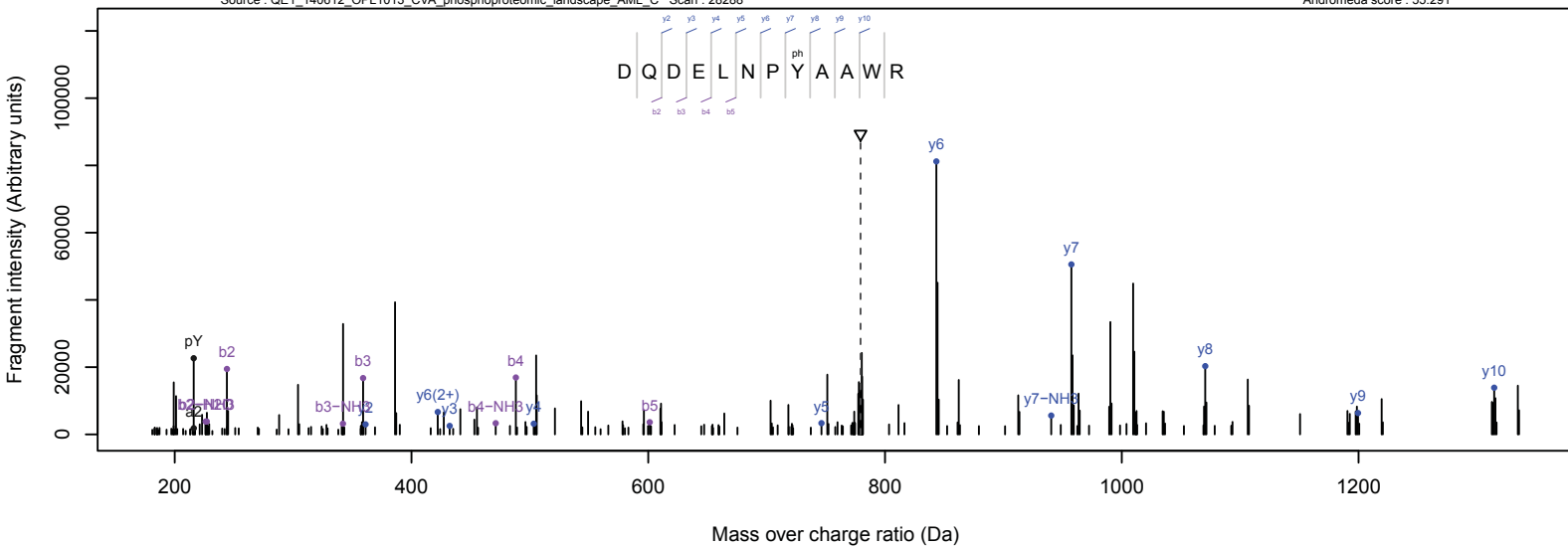
Andromeda score : 55.335



DQDELNPY(ph)AAWR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28288

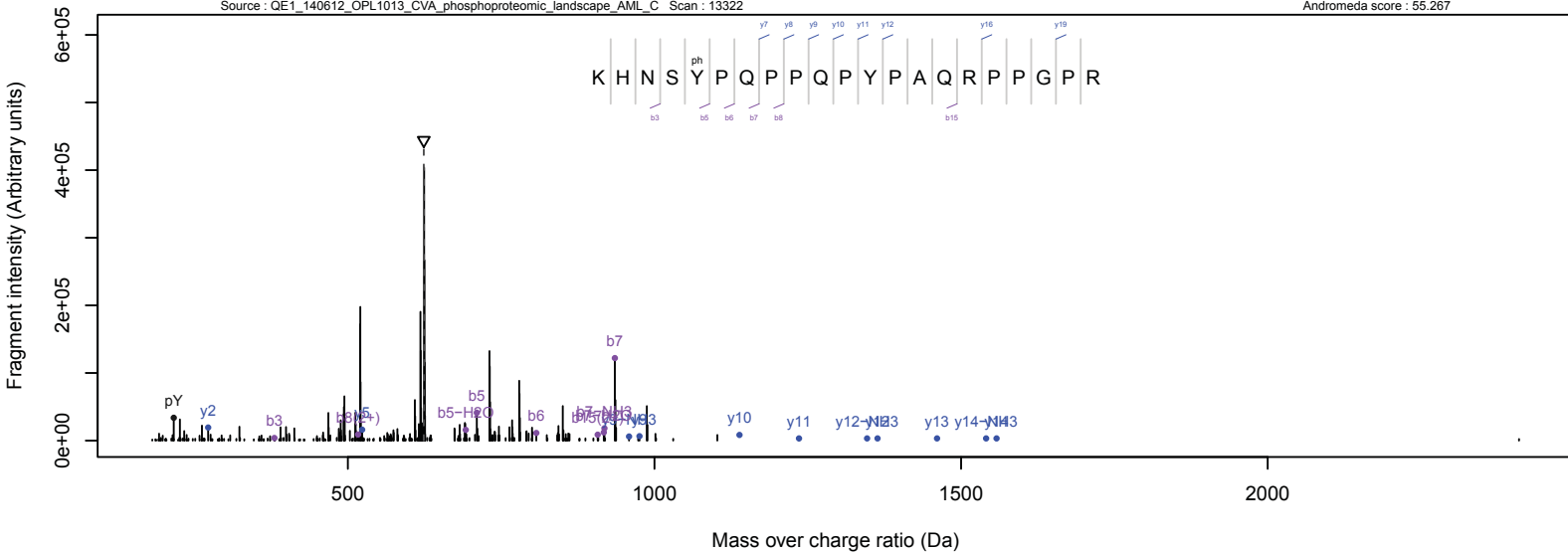
Andromeda score : 55.291



**KHNSY(ph)PQPYPYPAQRPPGPR-4+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13322

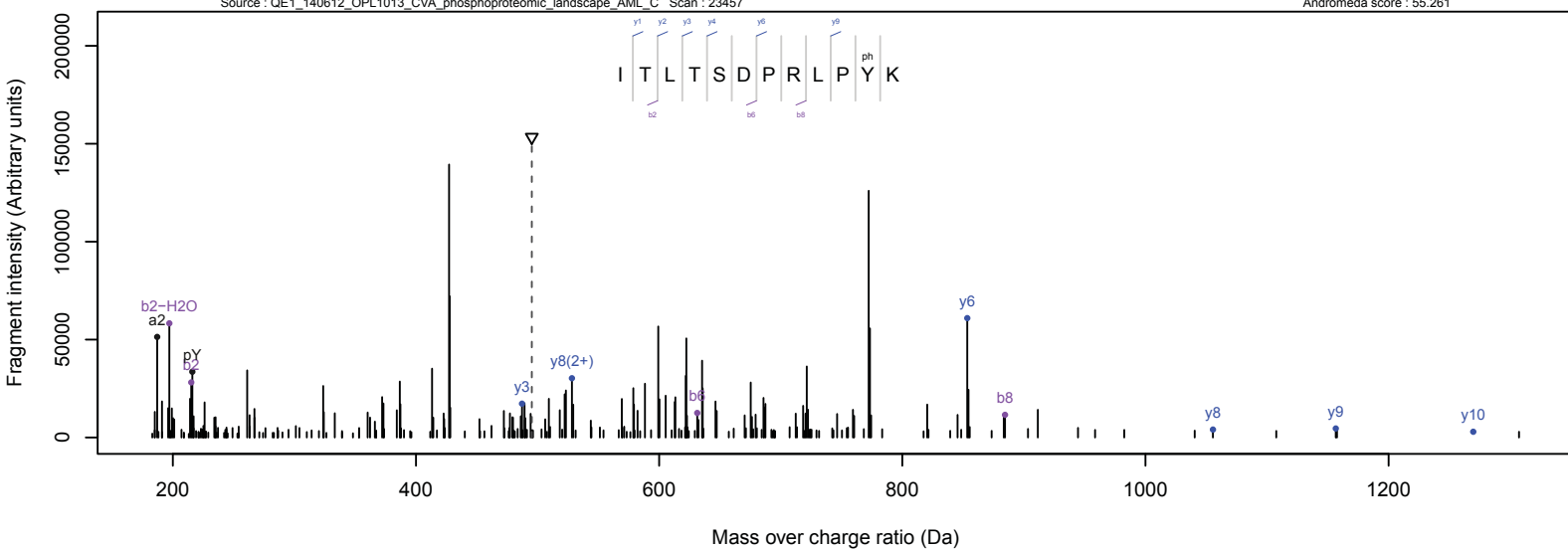
Andromeda score: 55.267



**ITLTSDPRLPY(ph)K-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23457

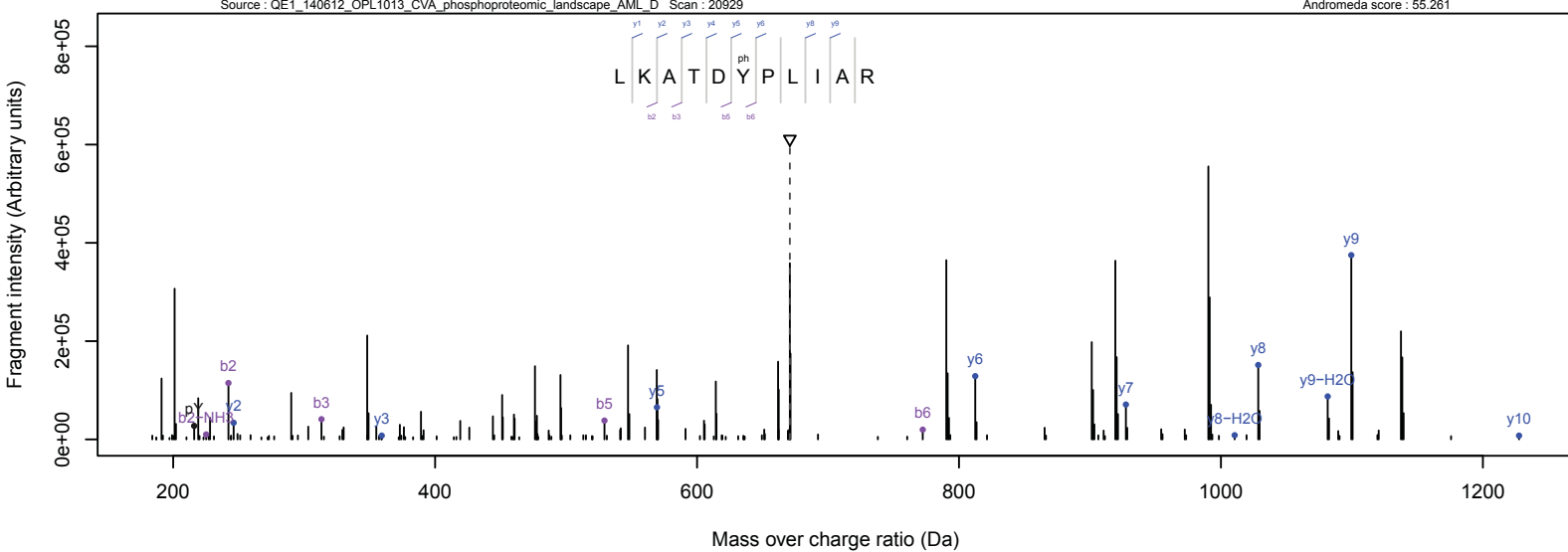
Andromeda score: 55.261



**LKATDY(ph)PLIAR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 20929

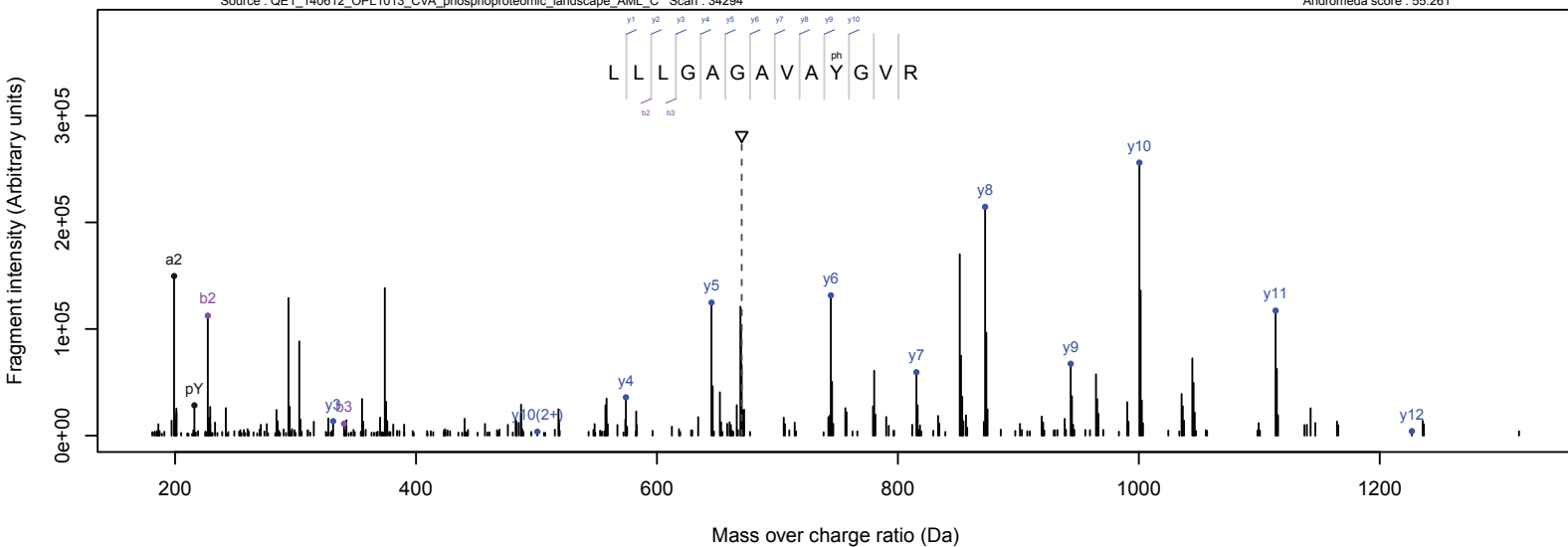
Andromeda score: 55.261



## LLLGAVAY(ph)GVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34294

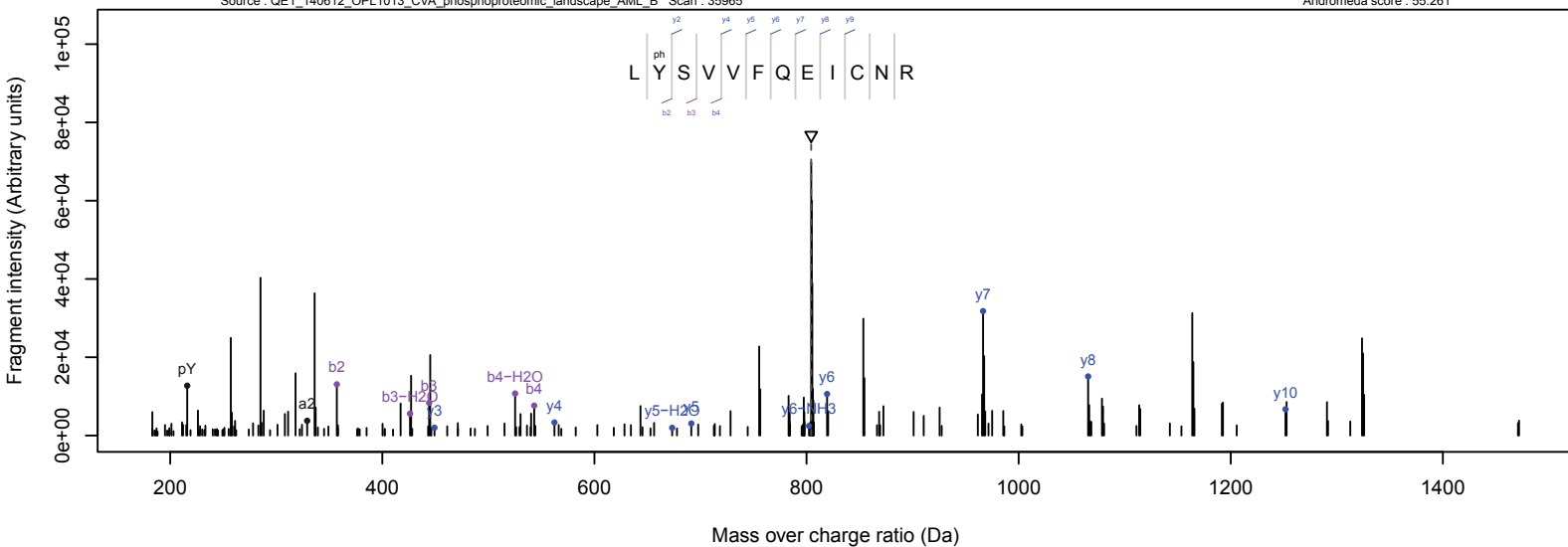
Andromeda score : 55.261



## LY(ph)SVVFQEICNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35965

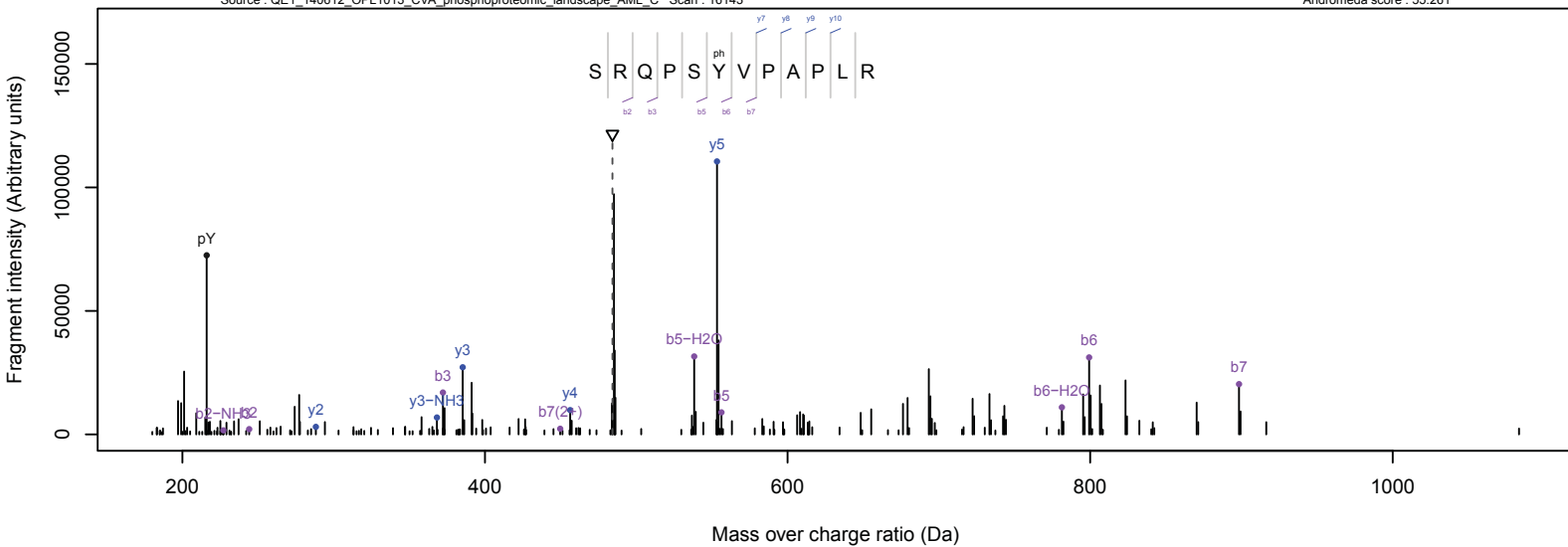
Andromeda score : 55.261



## SRQPSY(ph)VPAPLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16143

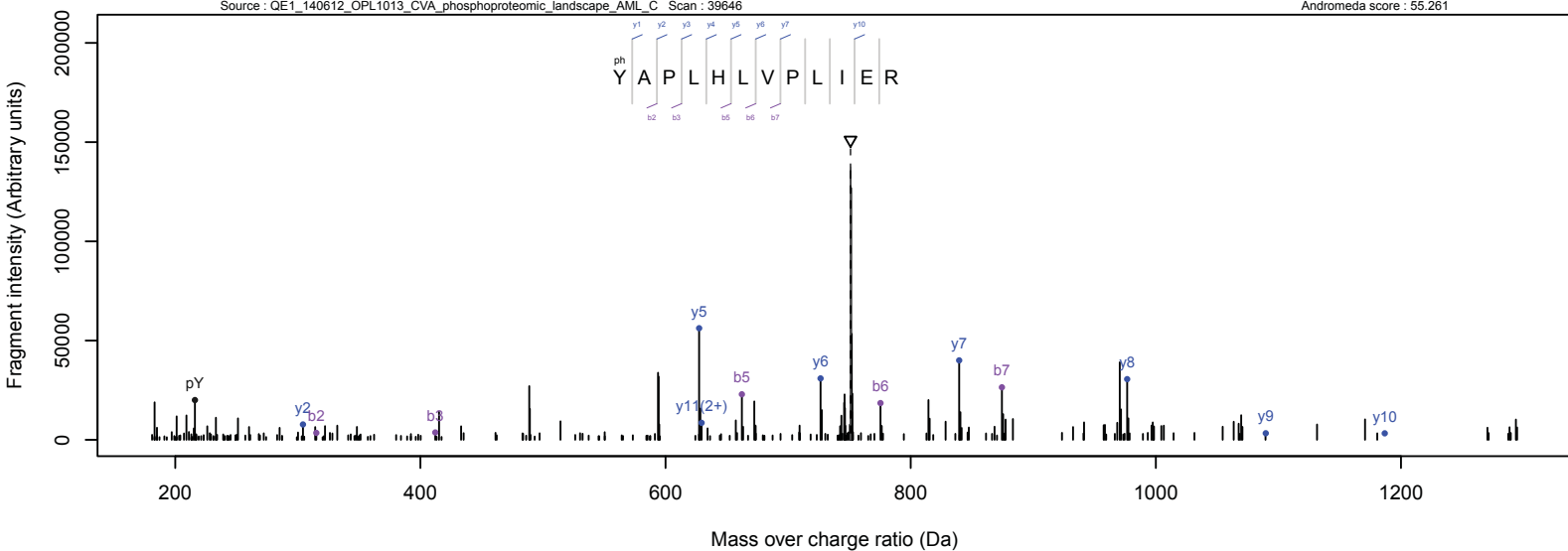
Andromeda score : 55.261



Y(ph)APLHLVPLIER-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 39646

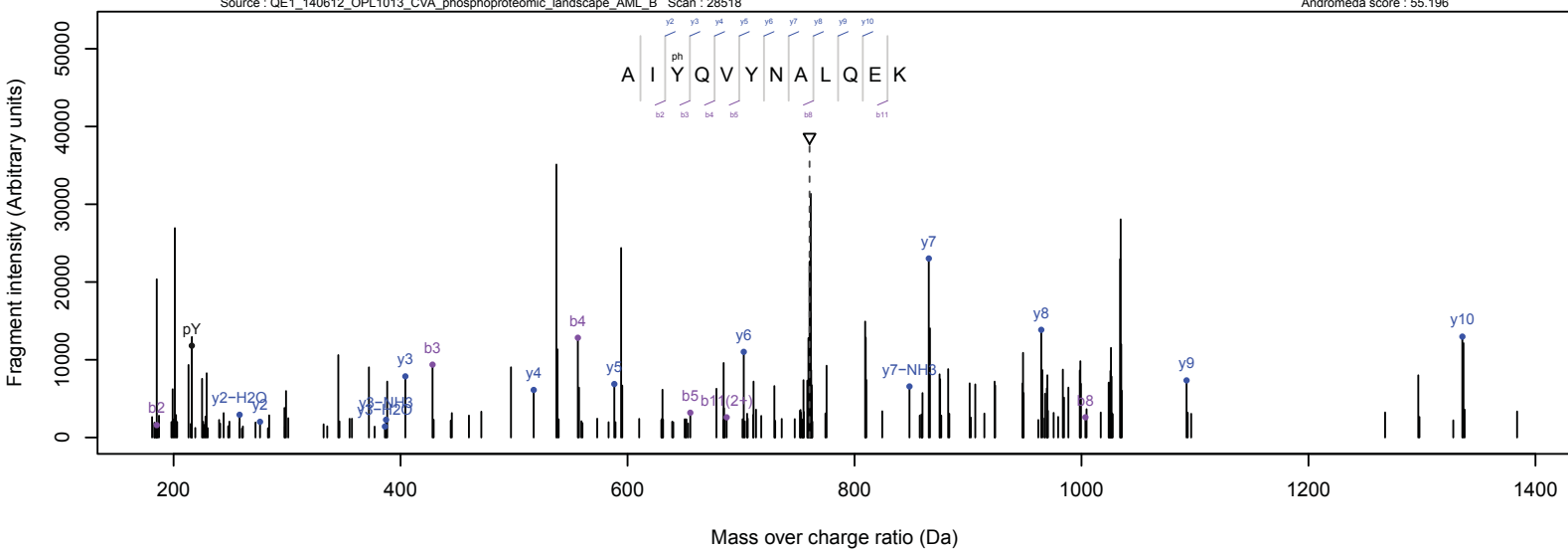
Andromeda score: 55.261



AIY(ph)QVYNALQEK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 28518

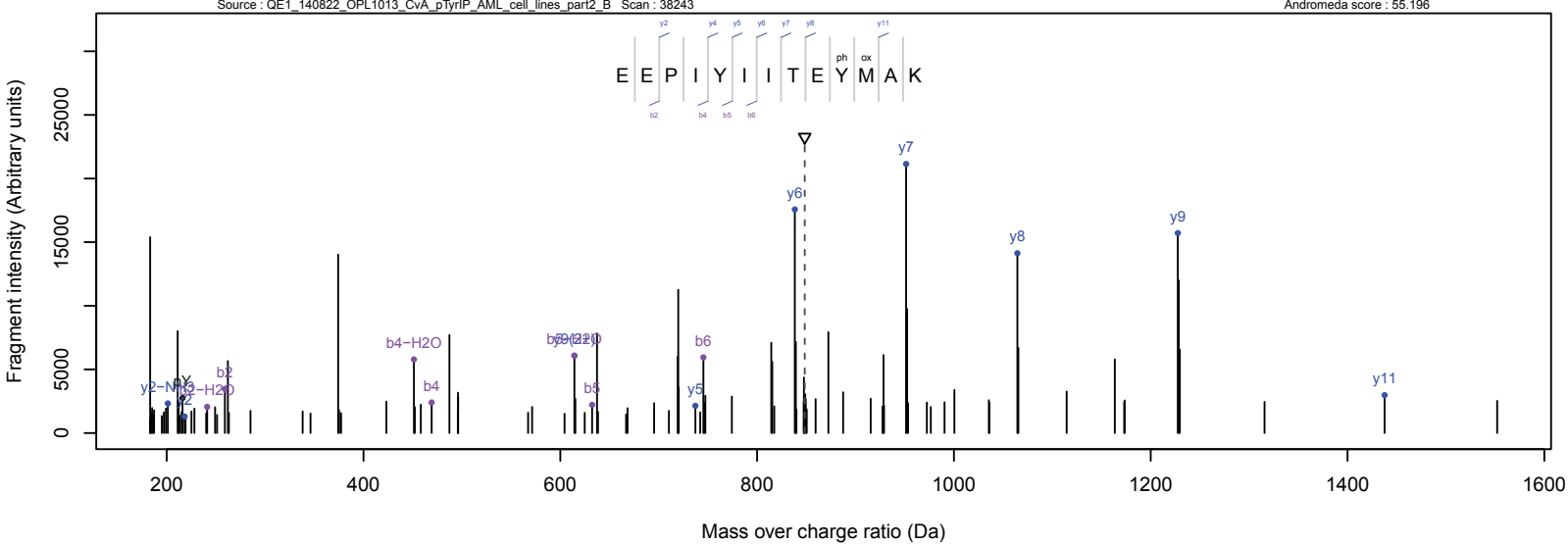
Andromeda score: 55.196



EEPIIITEY(ph)M(ox)AK-2+

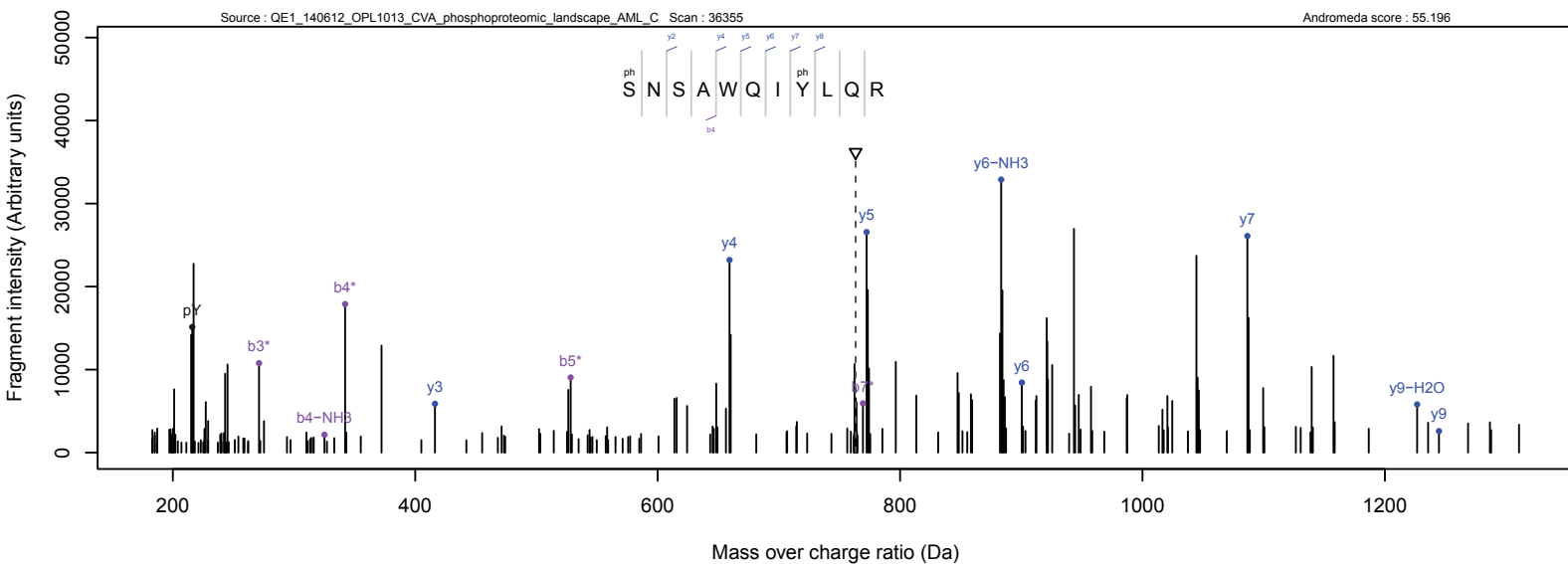
Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 38243

Andromeda score: 55.196

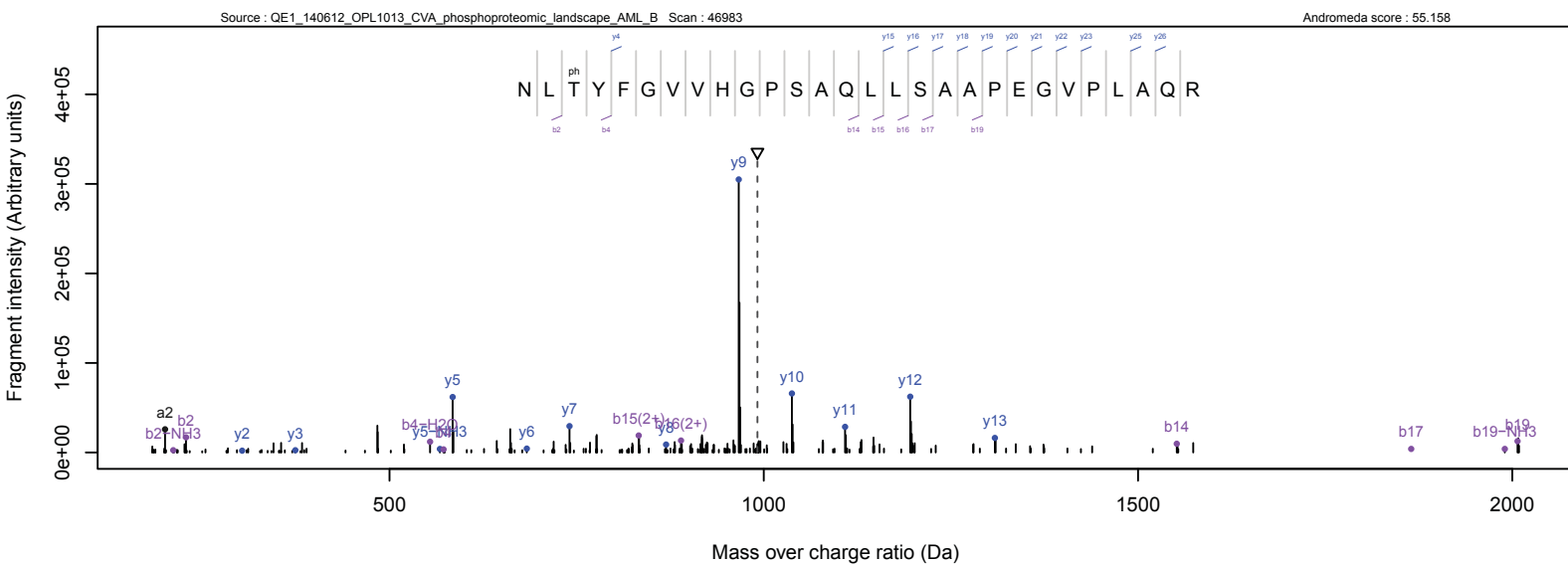




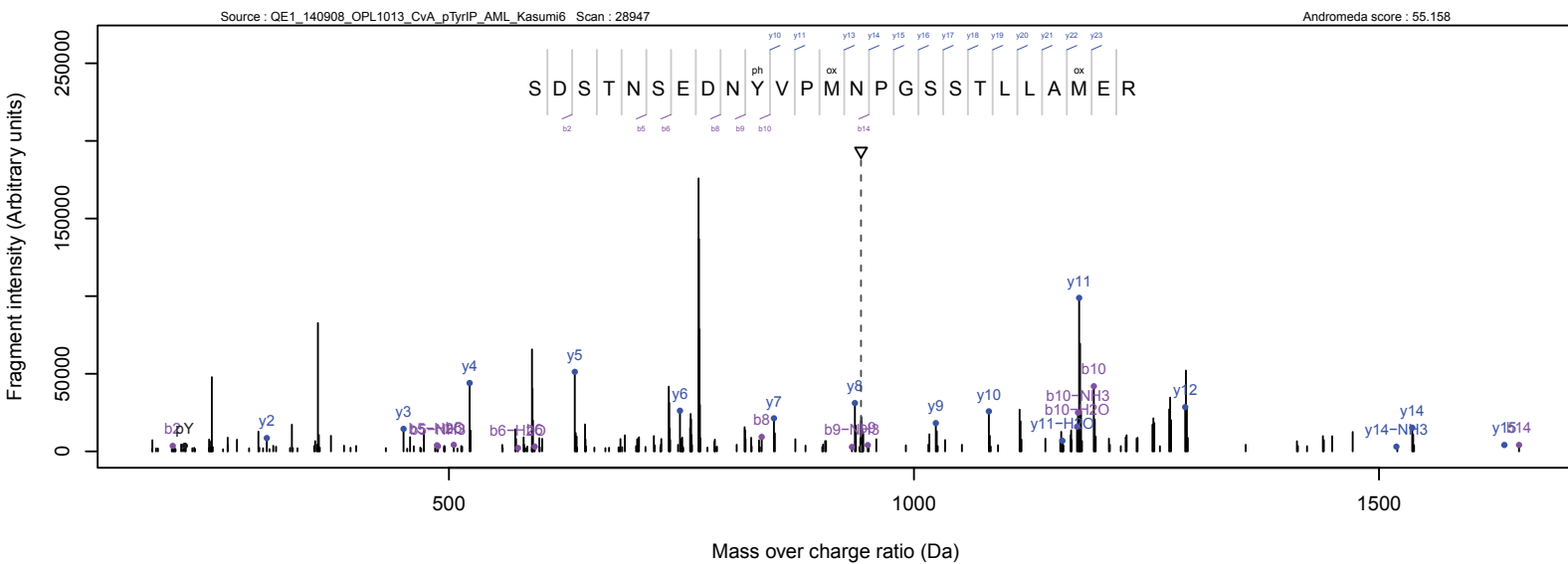
S(ph)NSAWQIY(ph)LQR-2+



NLT(ph)YFGVVHGPSAQLLSAAPEGVPLAQR-3+



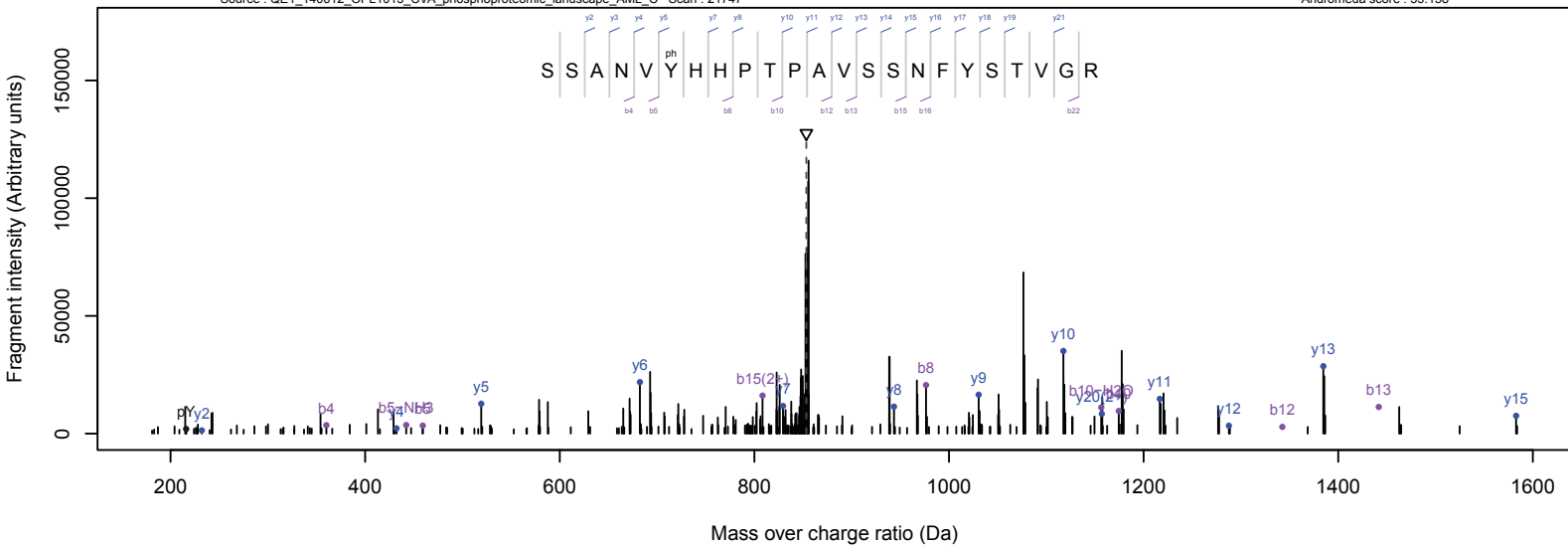
SDSTNSEDN(ph)VPMNPGSSTLLAMER-3+



## SSANVY(ph)HHPTPAVSSNFYSTVGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21747

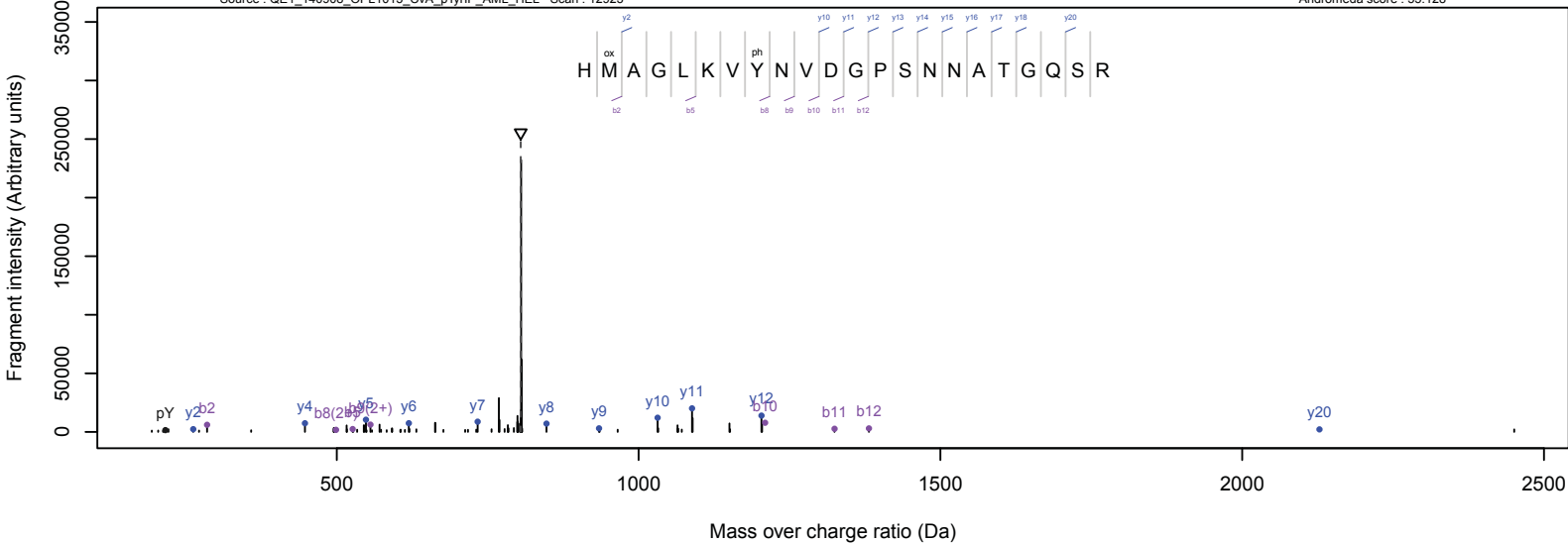
Andromeda score : 55.158



## HM(ox)AGLKVY(ph)NVDGSPNNATGQSR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 12923

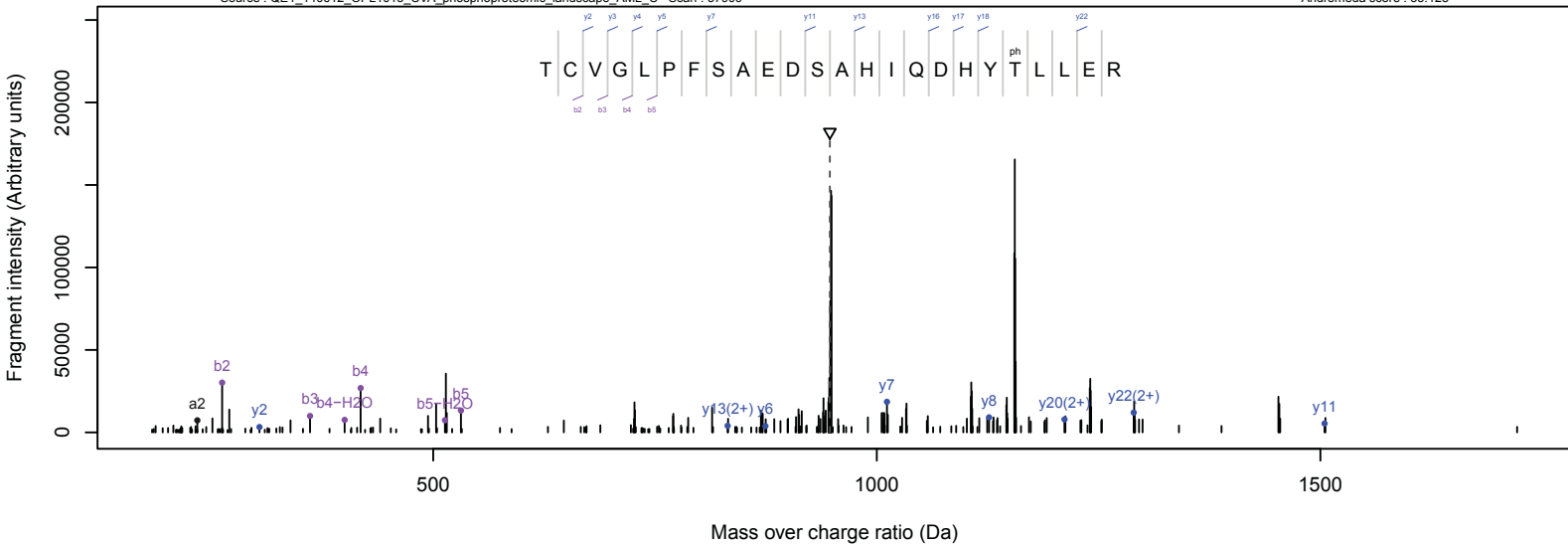
Andromeda score : 55.128



## TCVGLPFSAEHSAHQDHYT(ph)LLER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37909

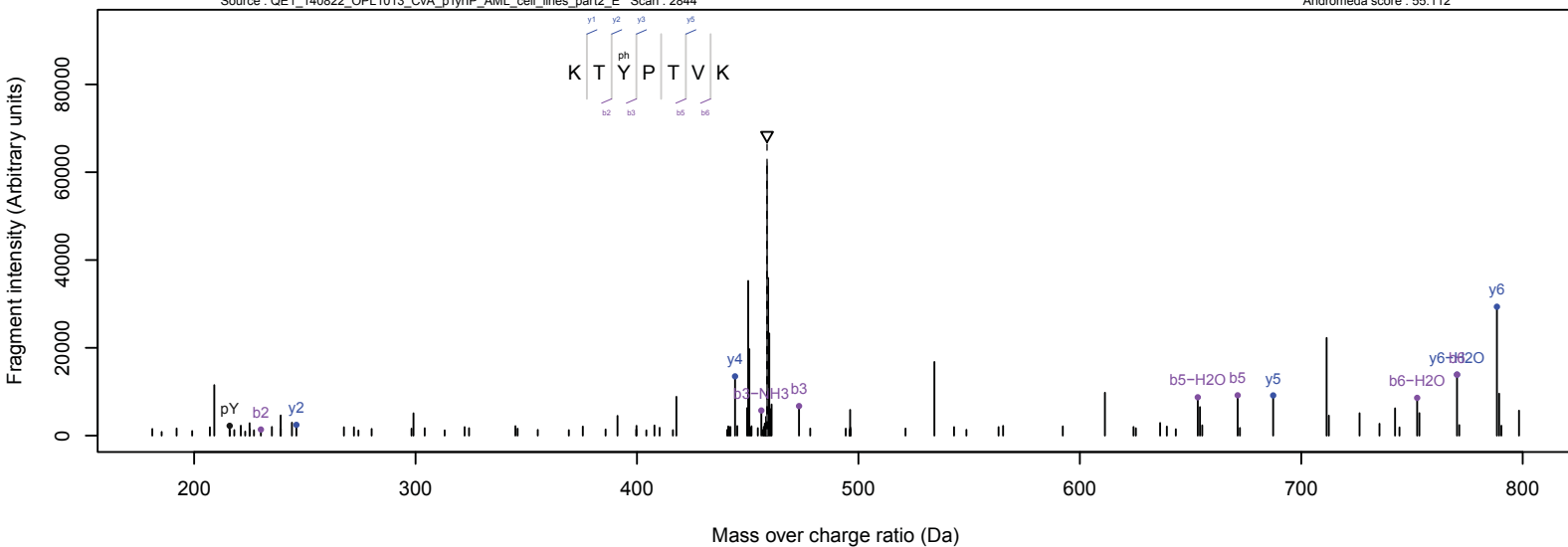
Andromeda score : 55.128



**KTY(ph)PTVK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 2844

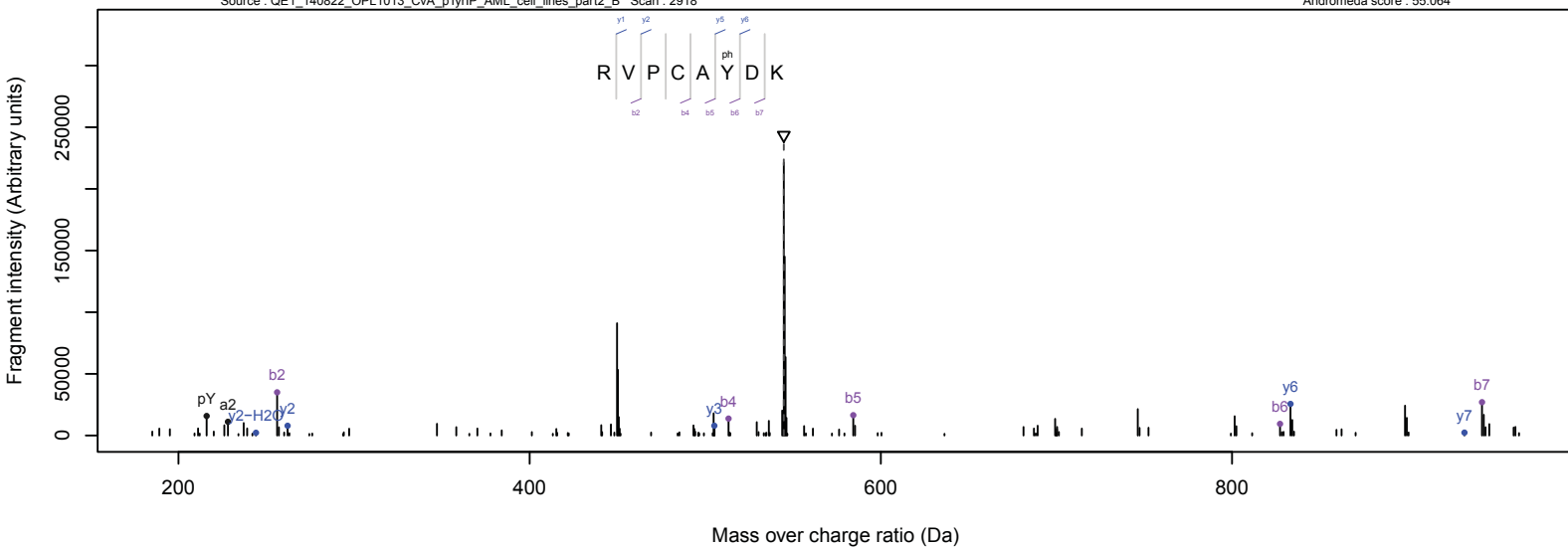
Andromeda score : 55.112



**RVPCAY(ph)DK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 2918

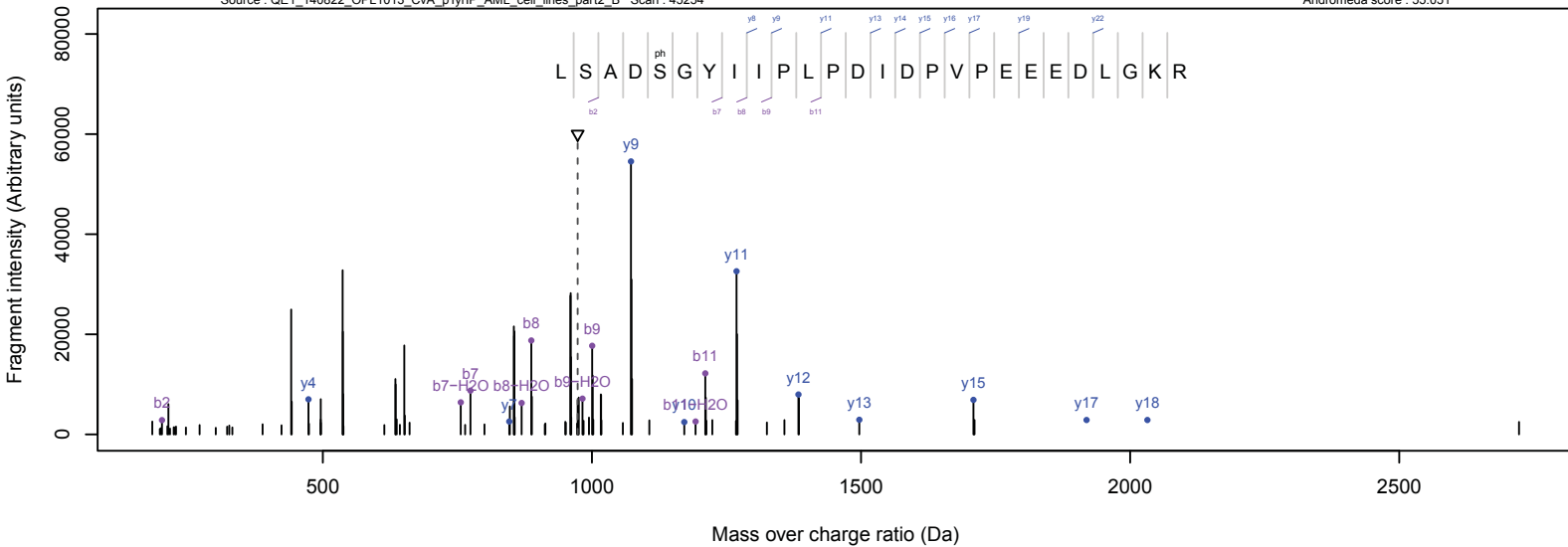
Andromeda score : 55.064



**LSADS(ph)GYIILPDIDPVPEEEDLGKR-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 45254

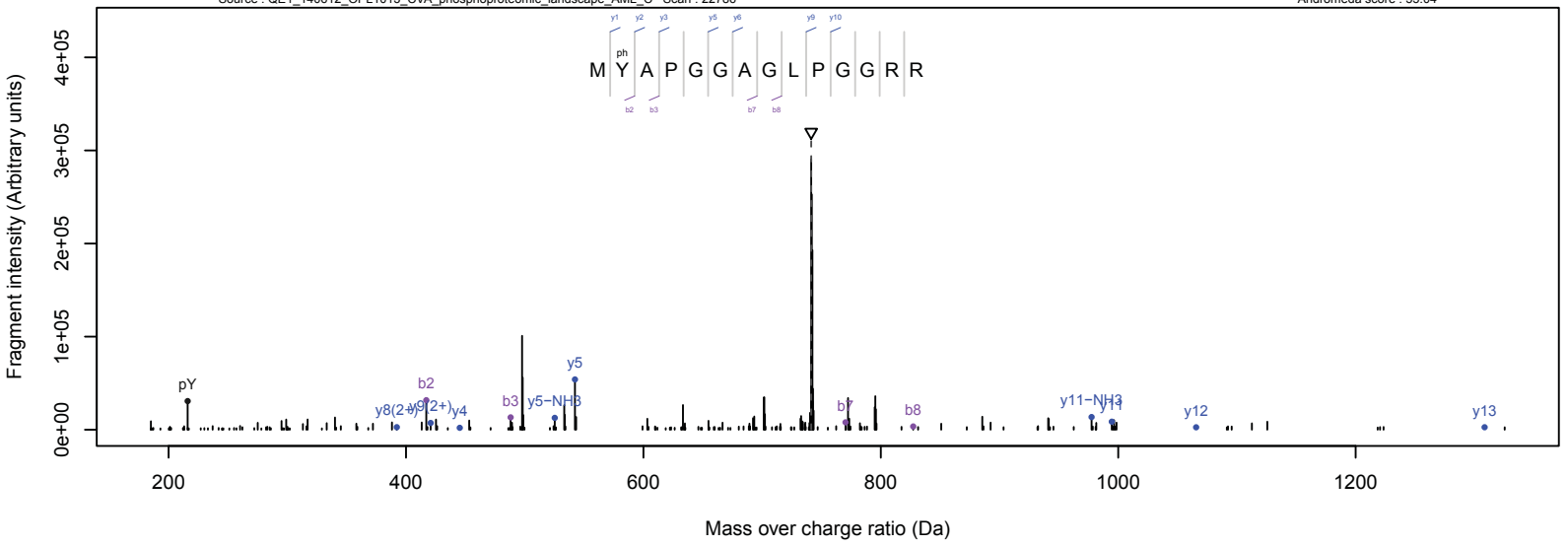
Andromeda score : 55.051



(ac)MY(ph)APGGAGLPGRR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22786

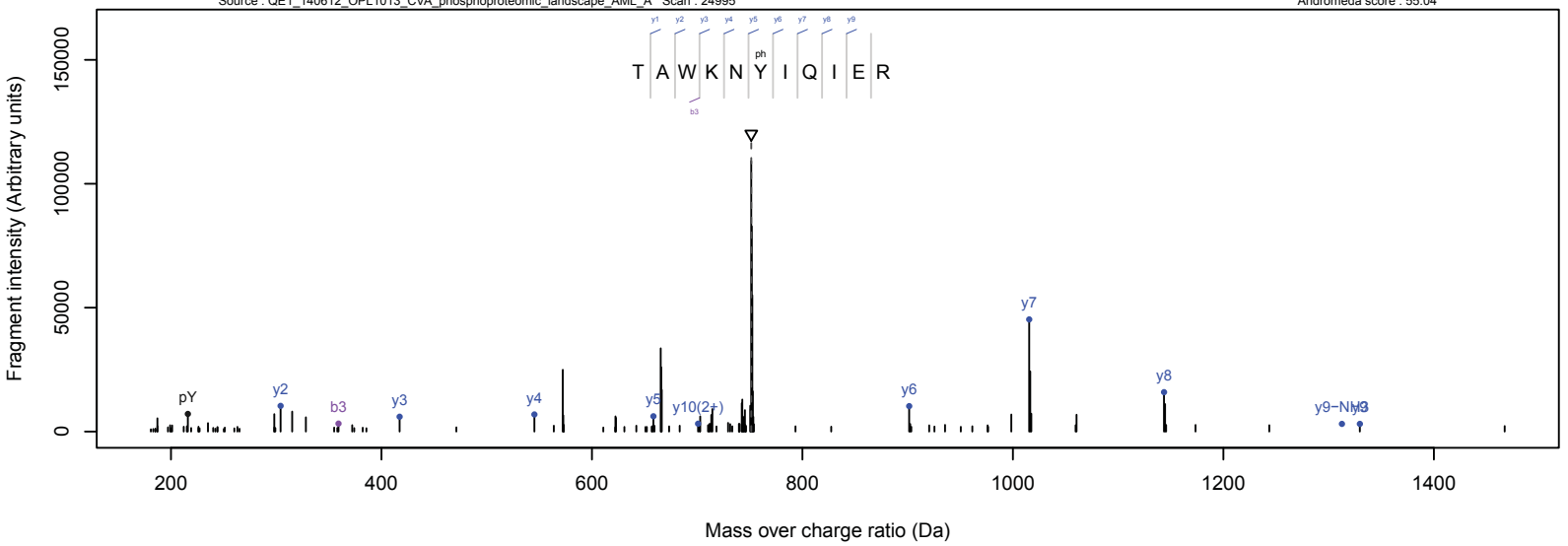
Andromeda score : 55.04



TAWKNY(ph)IQIER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 24995

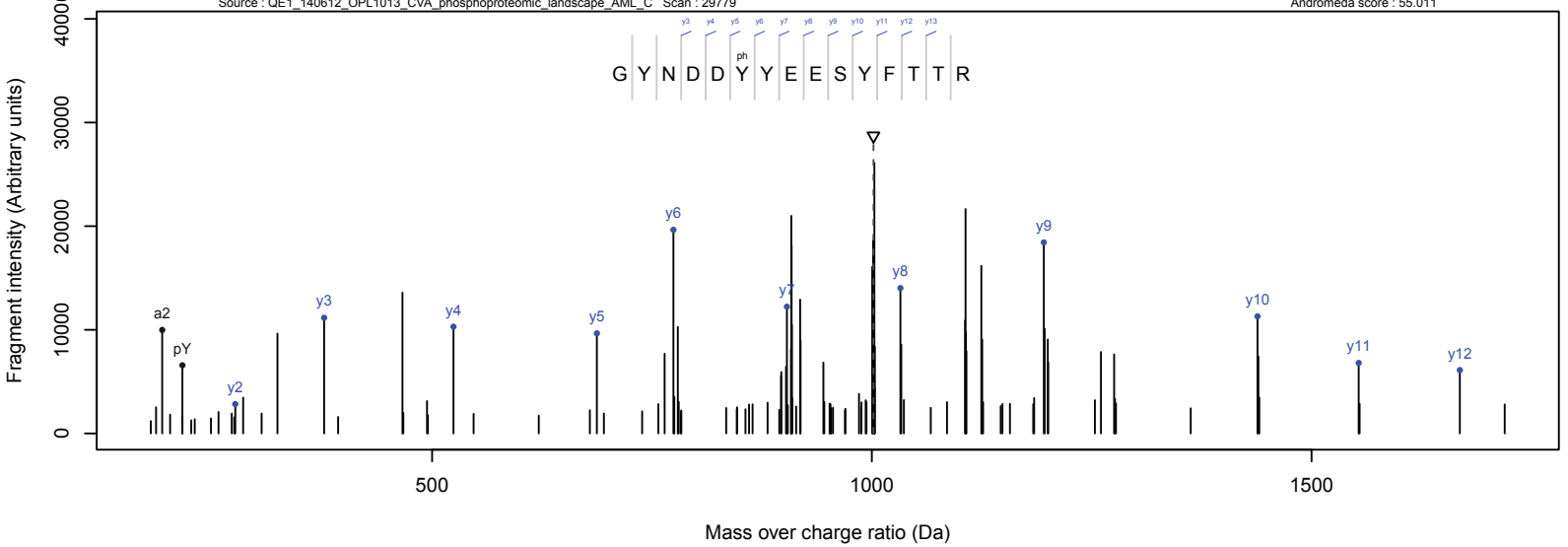
Andromeda score : 55.04



GYNDY(ph)YEESYFTTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29779

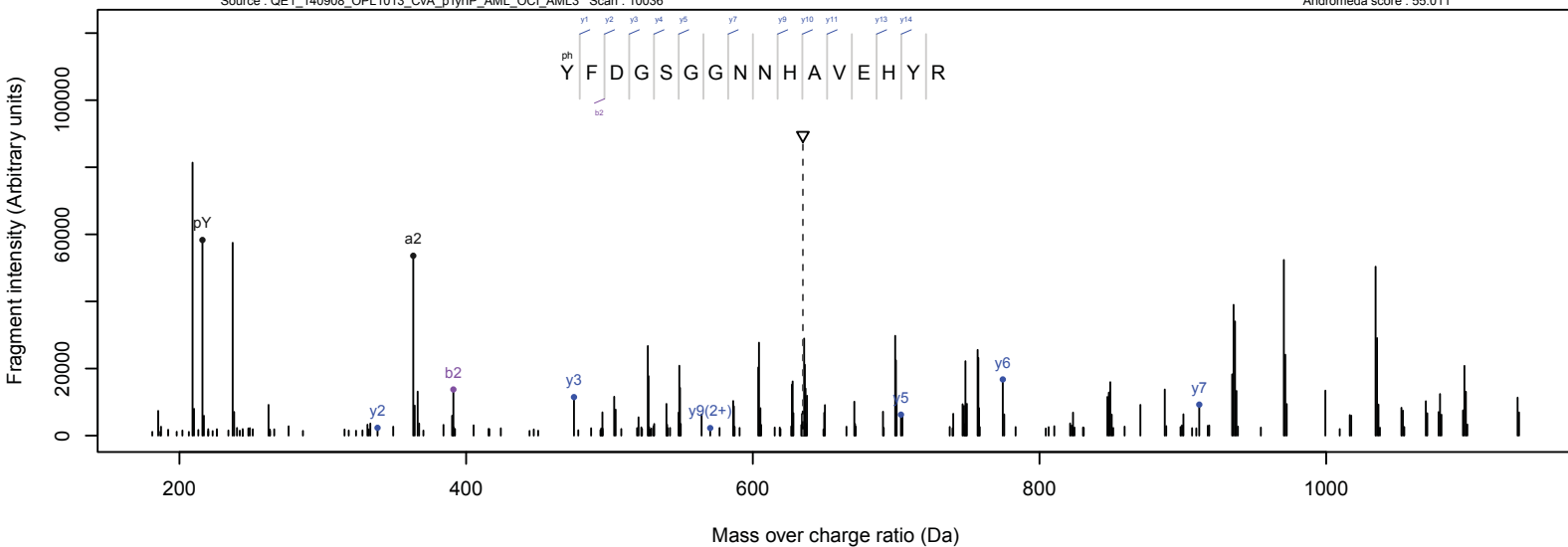
Andromeda score : 55.011



## Y(ph)FDGSGGNHVEHYR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 10036

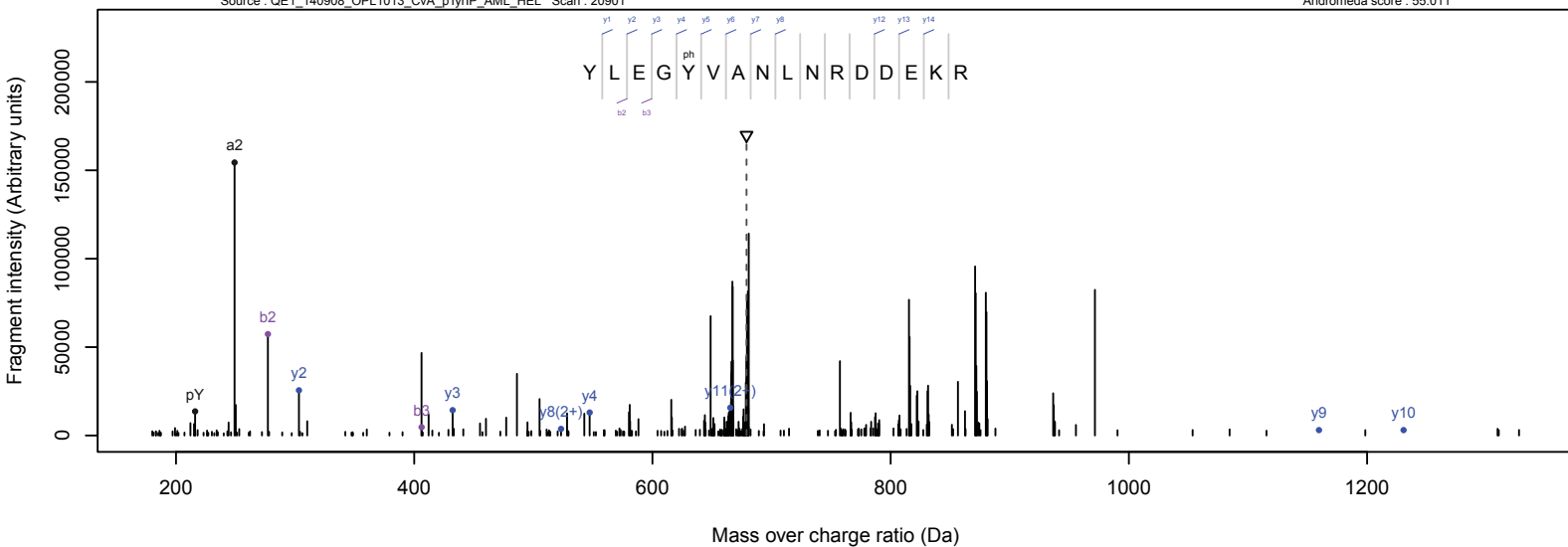
Andromeda score : 55.011



## YLEGY(ph)VANLRDDEKR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 20901

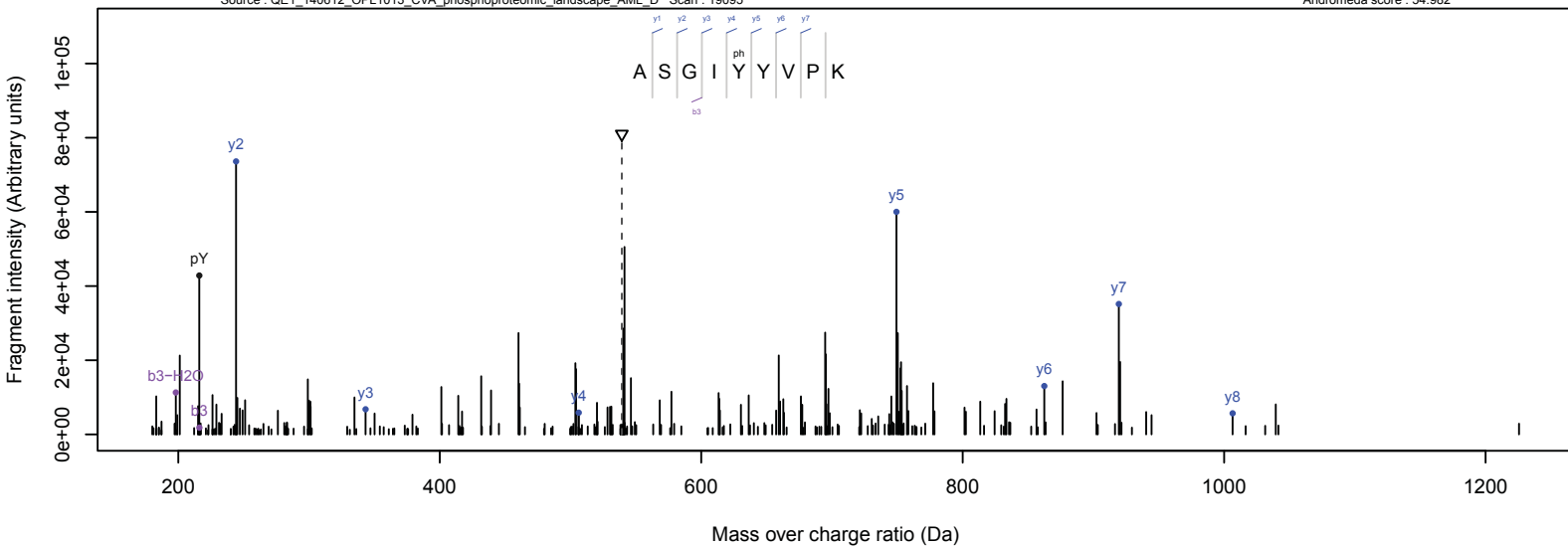
Andromeda score : 55.011



## ASGIY(ph)YVPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 19095

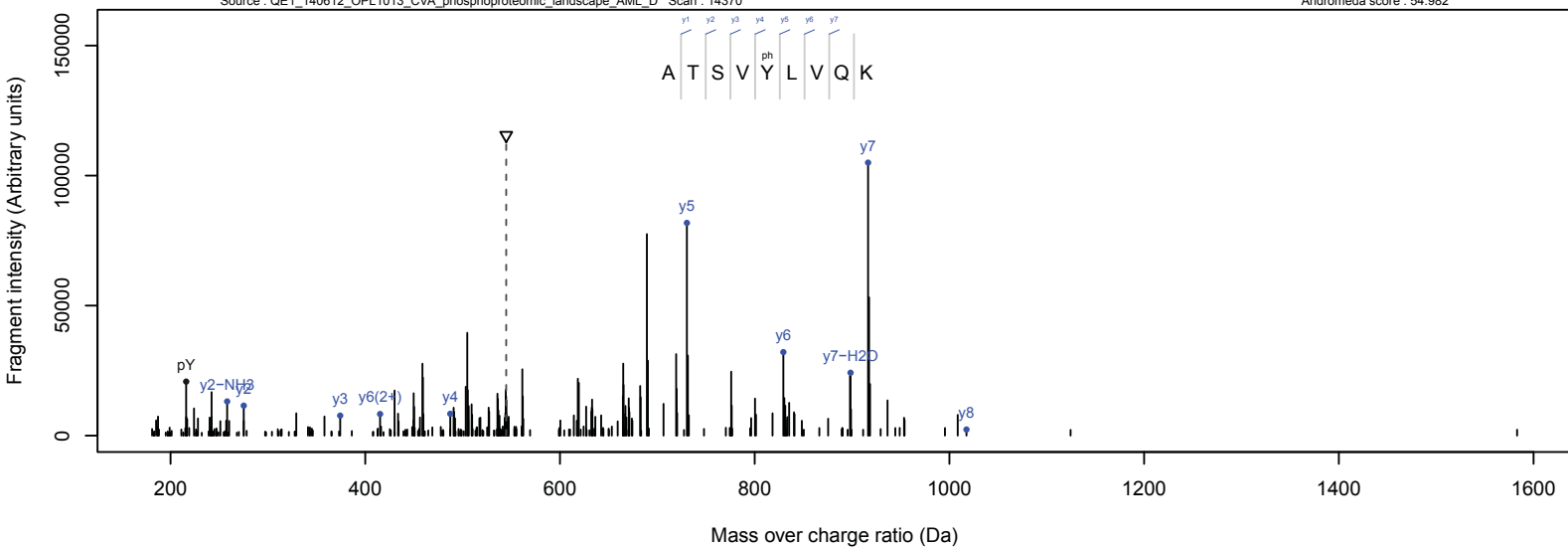
Andromeda score : 54.982



## ATSVY(ph)LVQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 14370

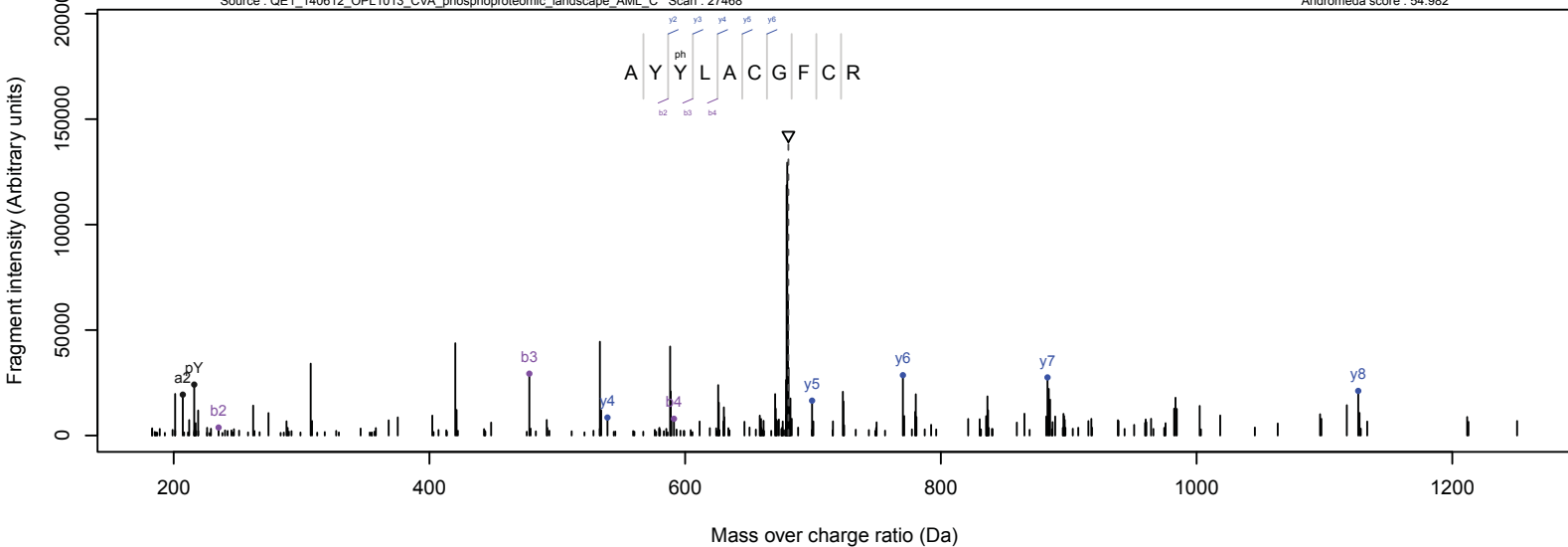
Andromeda score : 54.982



## AYY(ph)LACGFCR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27468

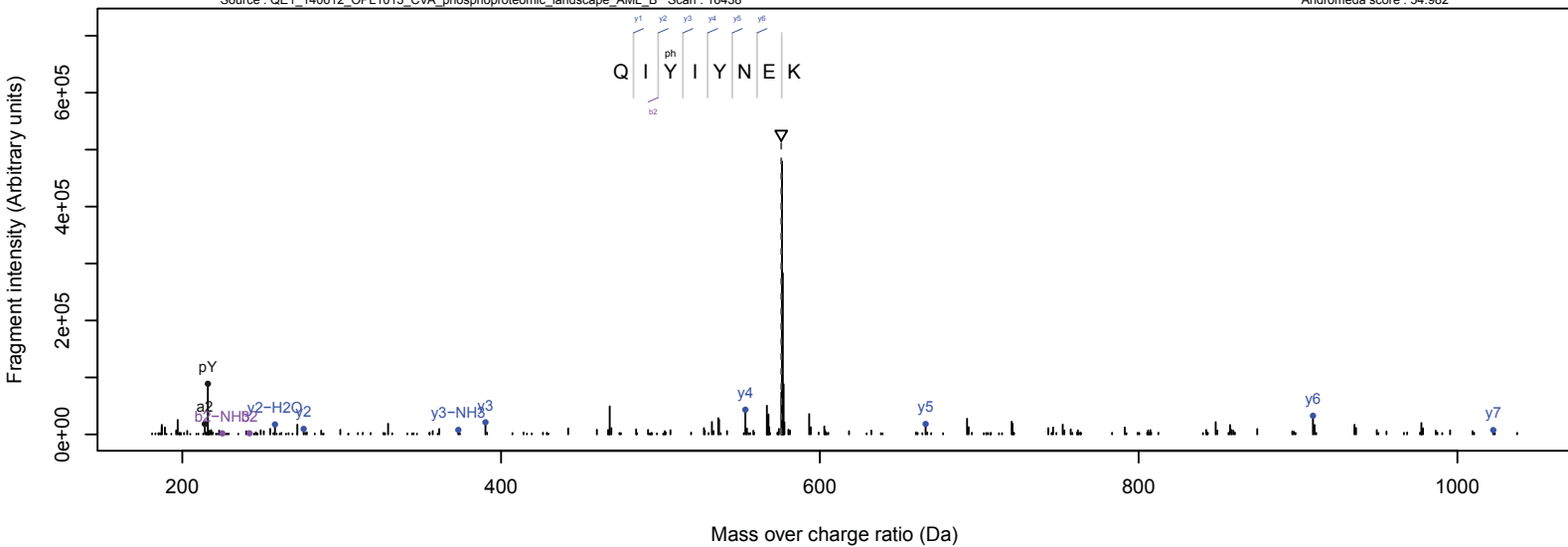
Andromeda score : 54.982



## QIY(ph)IYNEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16438

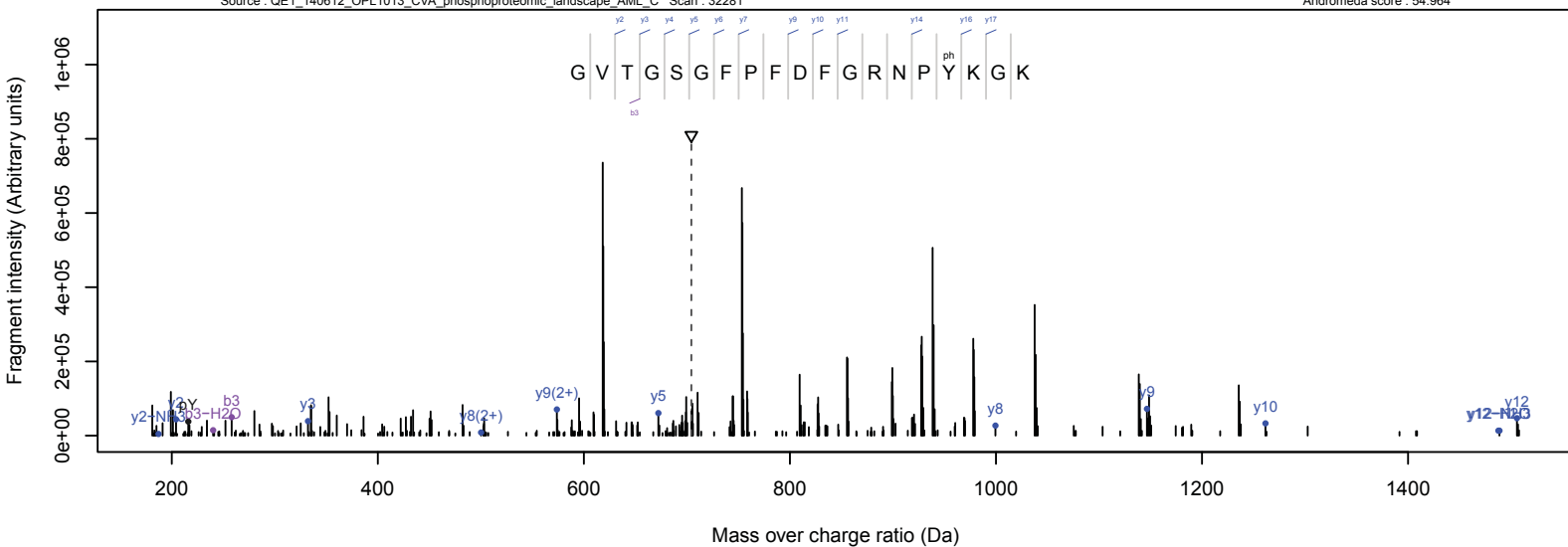
Andromeda score : 54.982



GVTGSGFFDFGRNPYKKGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32281

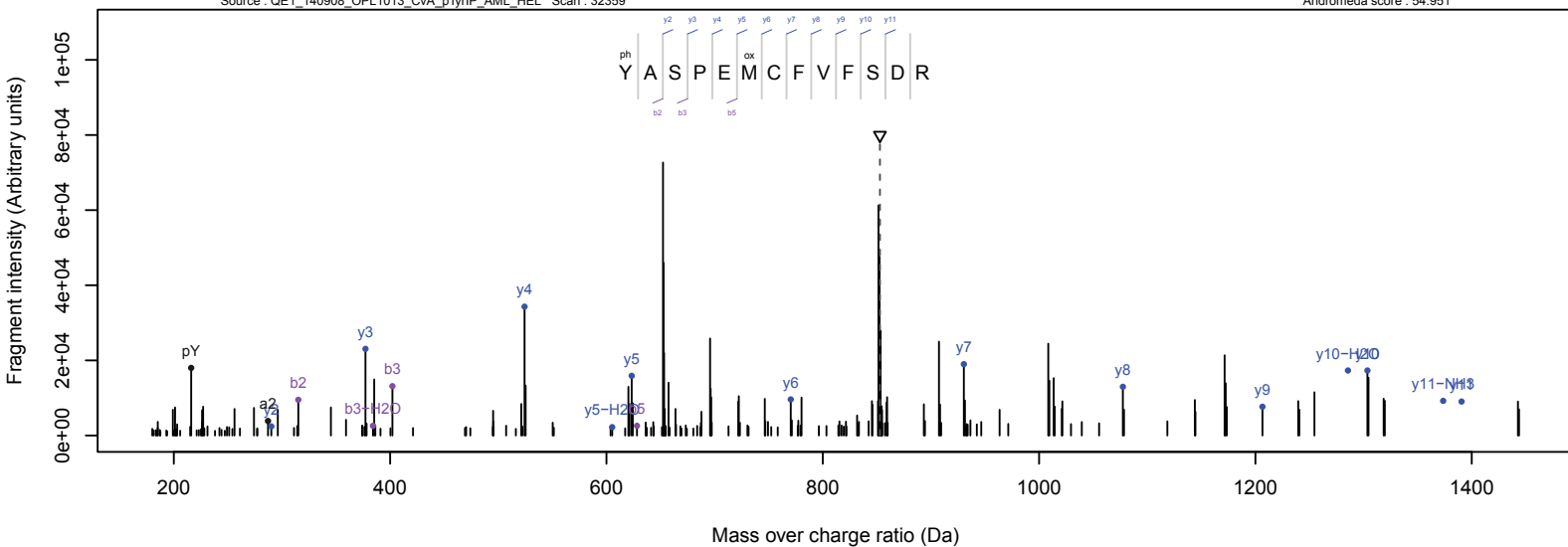
Andromeda score : 54.964



Y(ph)ASPEM(ox)CFVFSDR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32359

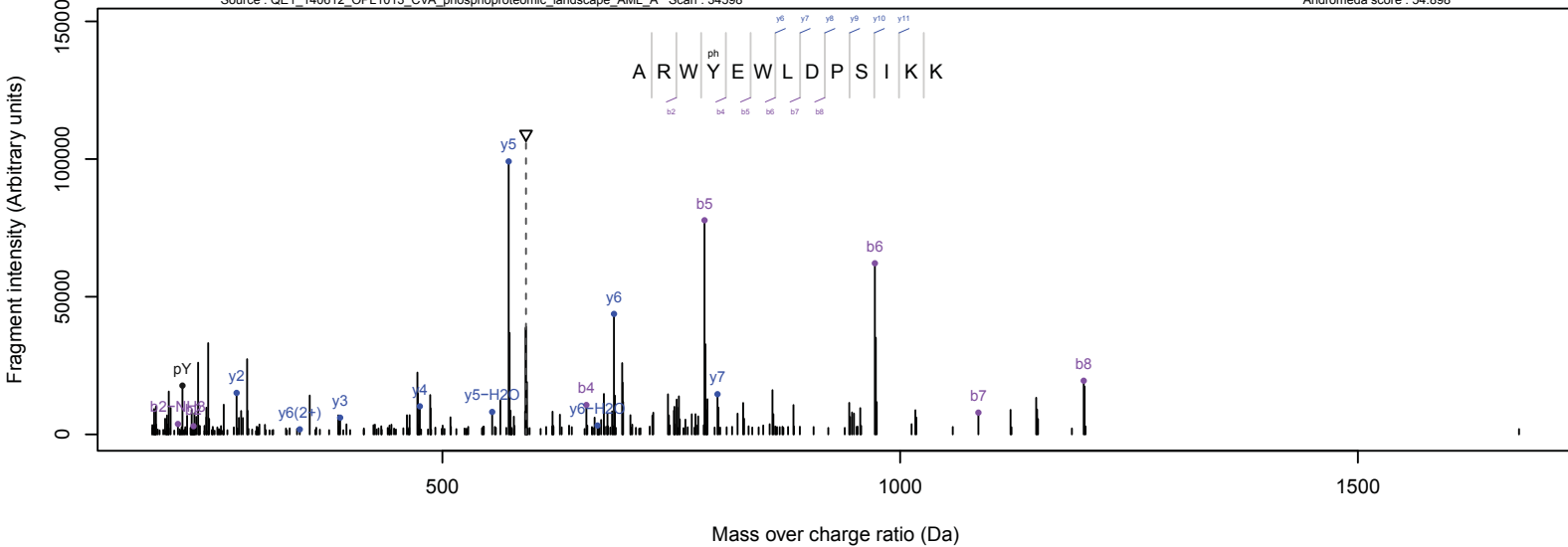
Andromeda score : 54.951

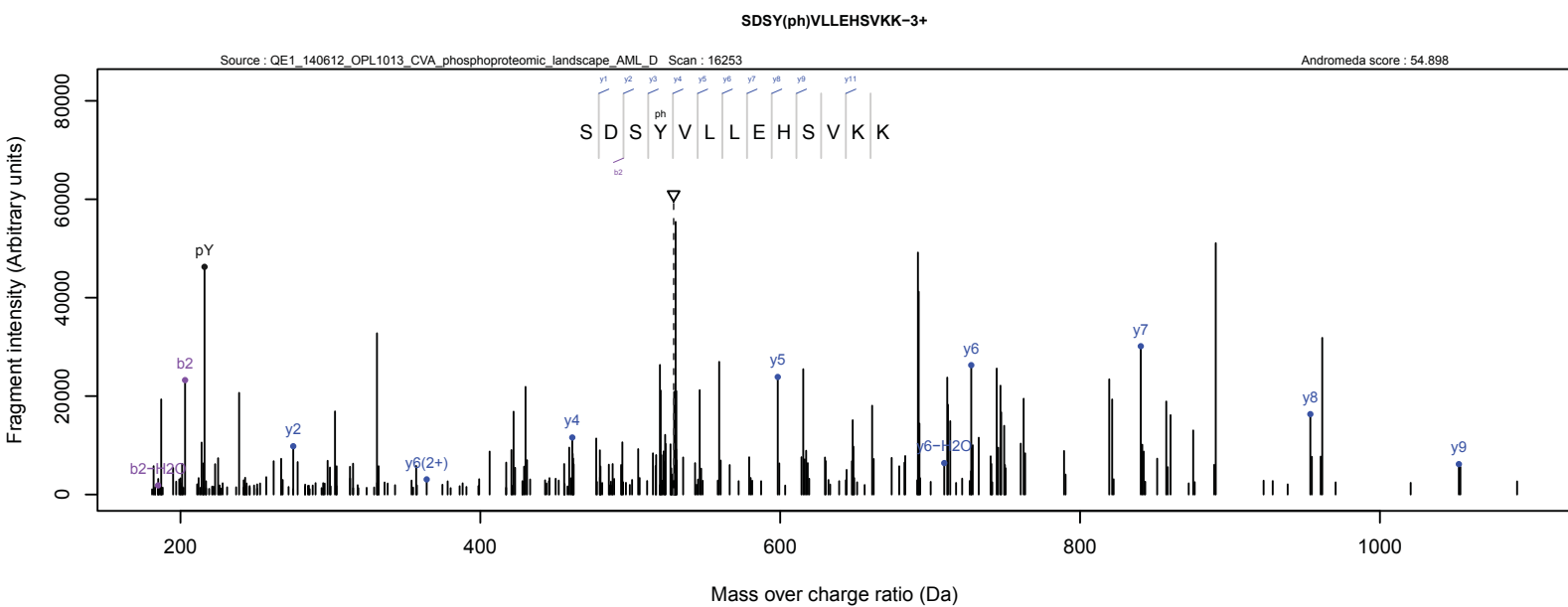
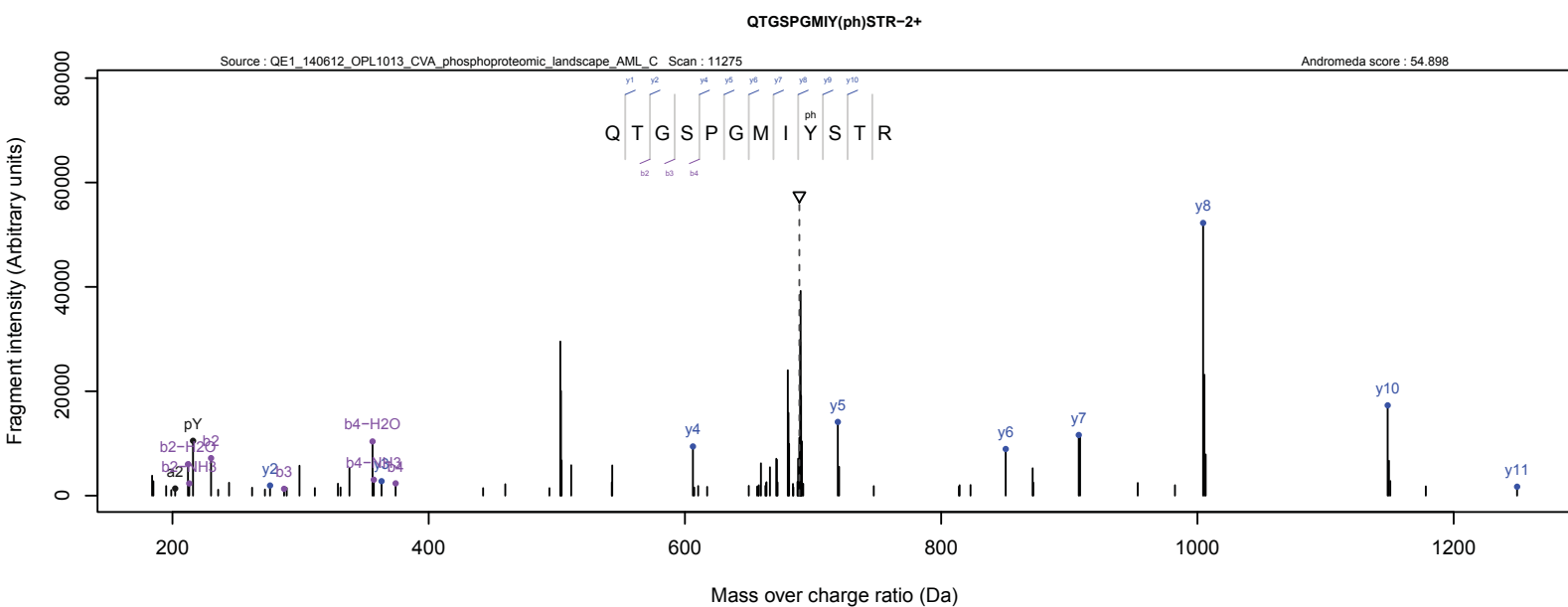
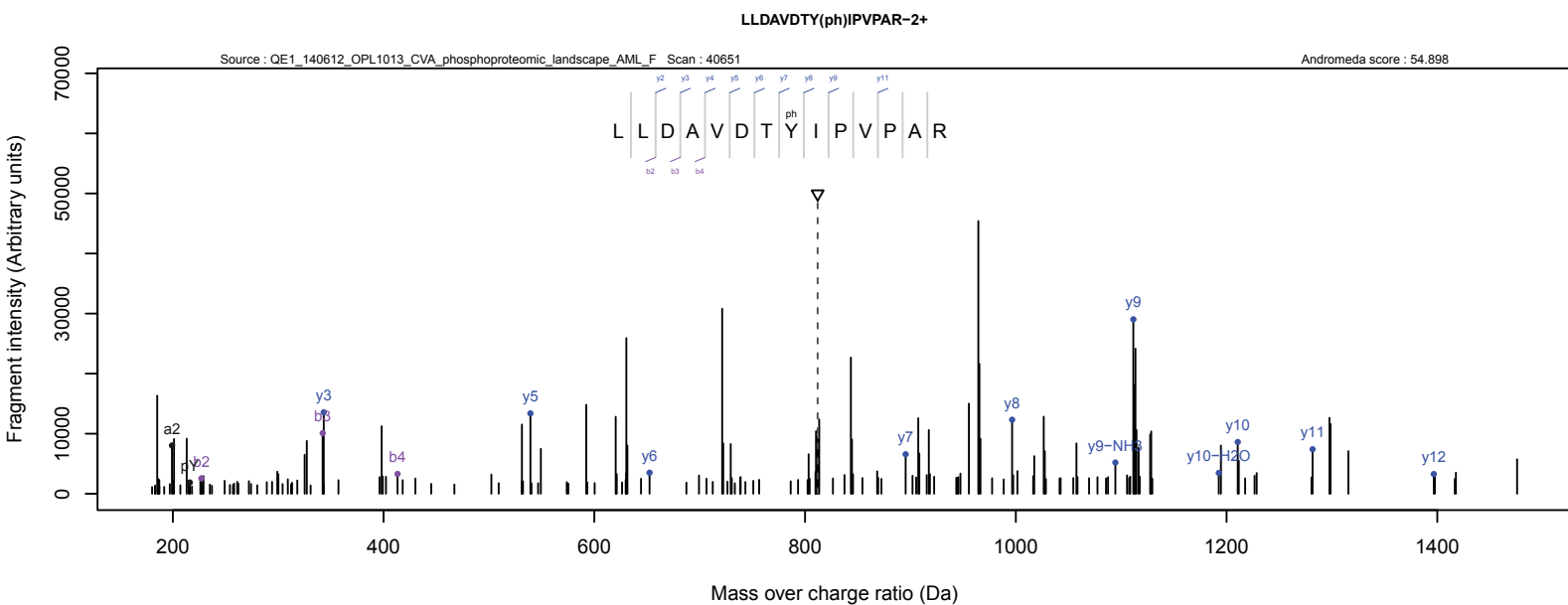


ARWY(ph)EWLDPSIKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34598

Andromeda score : 54.898



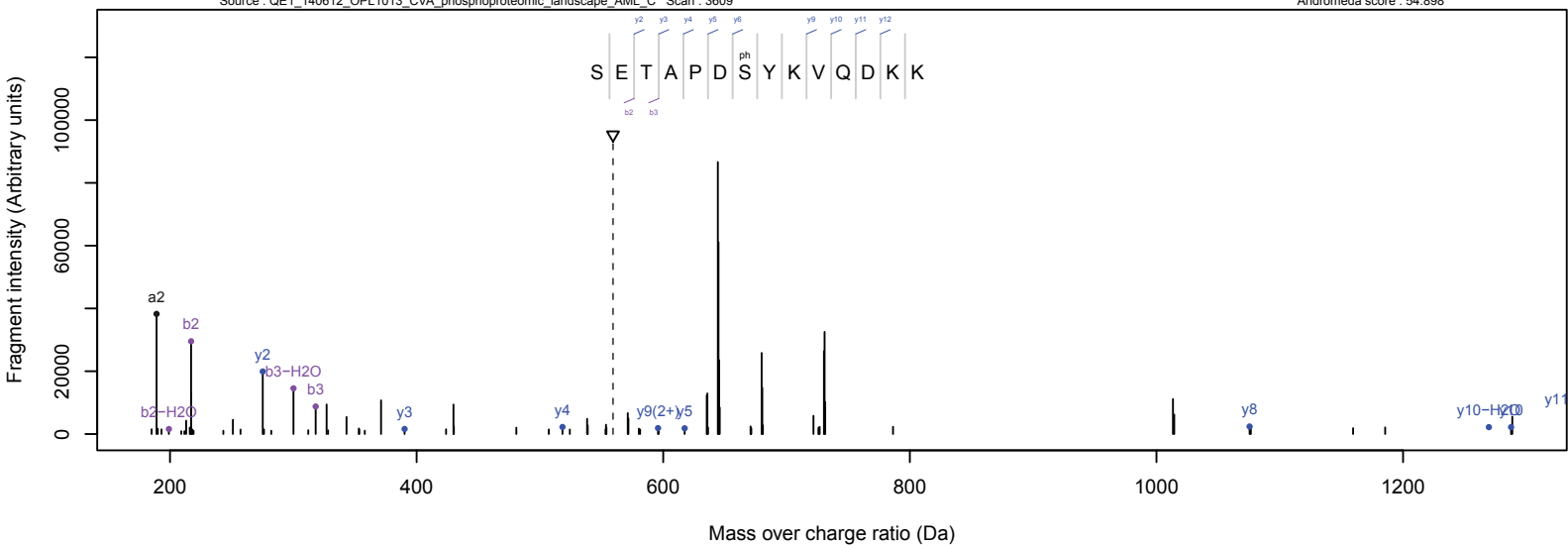




## SETAPDS(ph)YKVQDKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3609

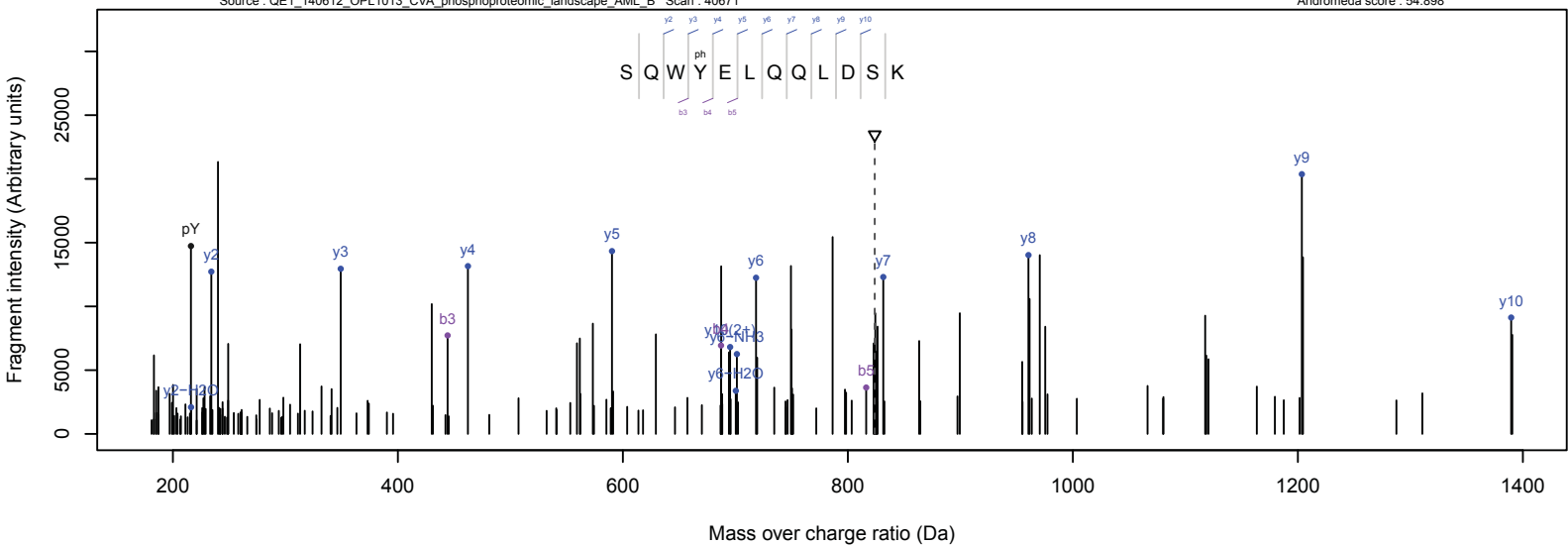
Andromeda score : 54.898



## (ac)SQWY(ph)ELQLDSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40671

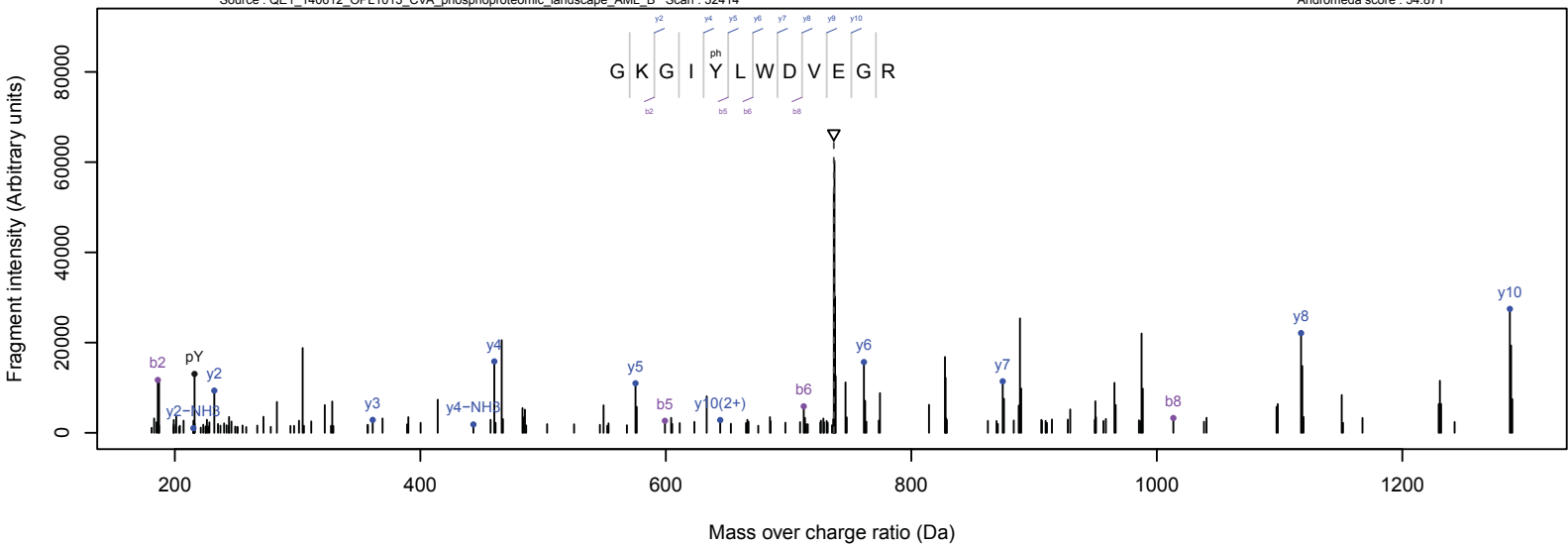
Andromeda score : 54.898



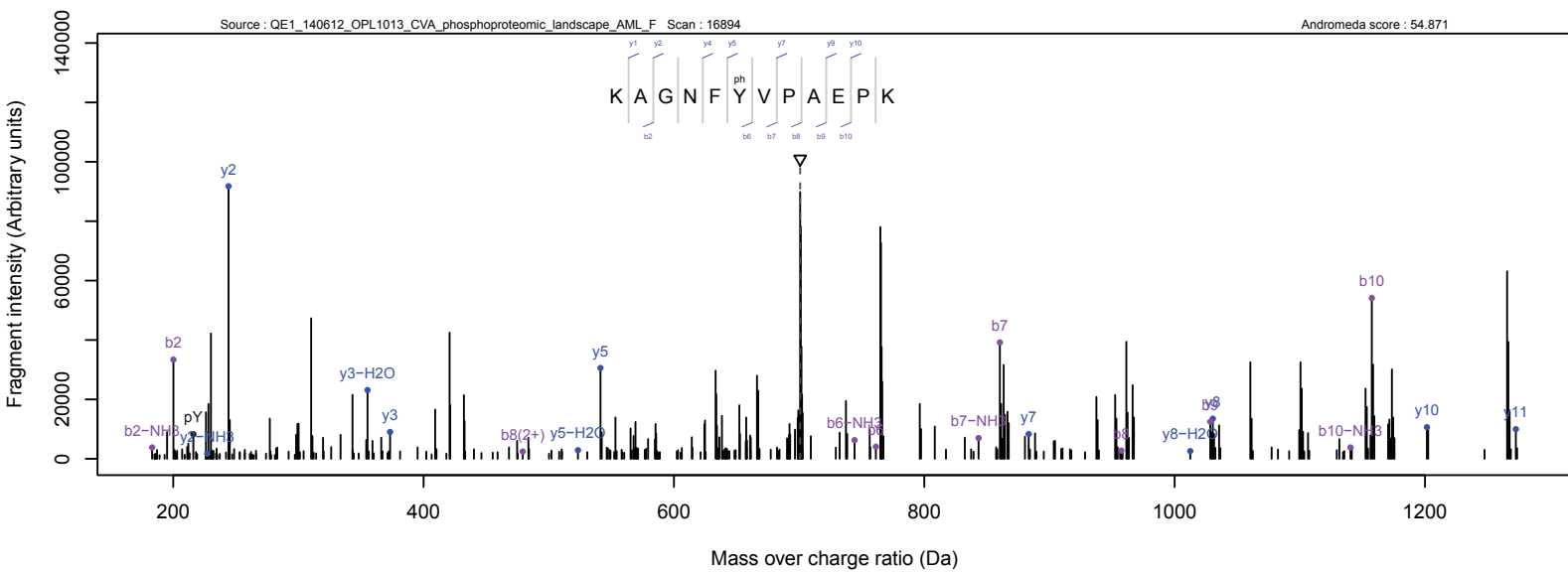
## GKGIY(ph)LWDVEGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32414

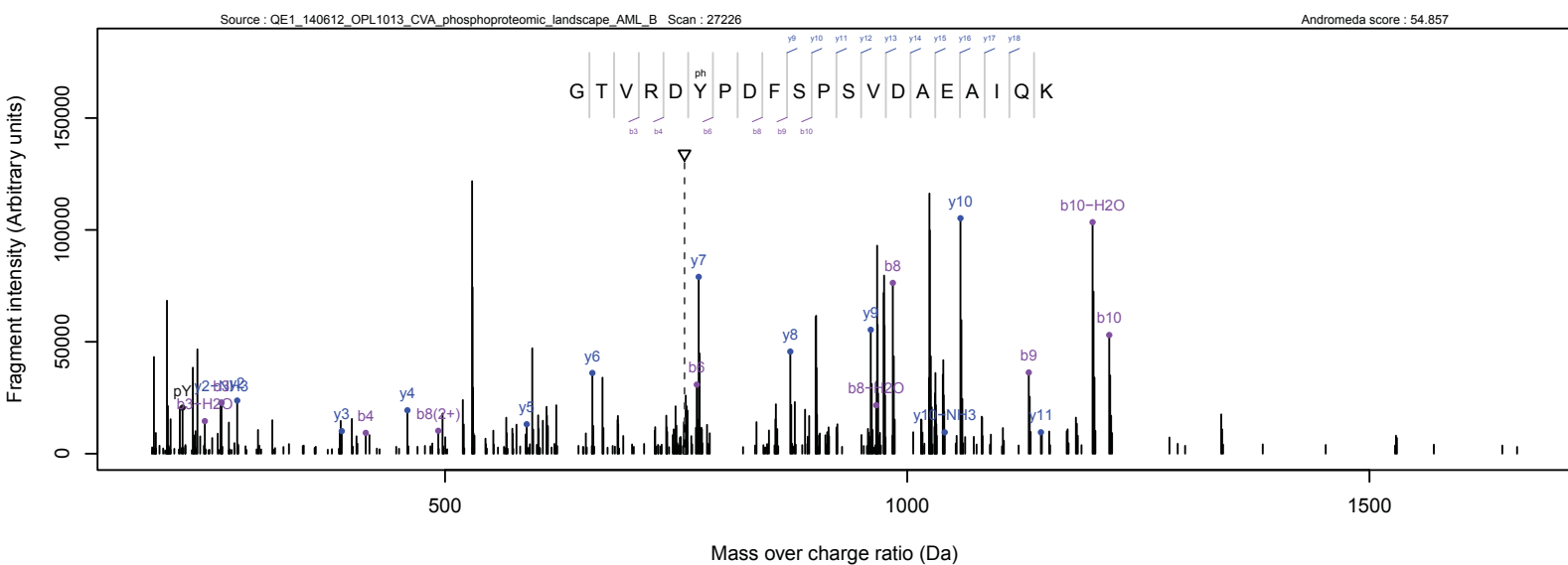
Andromeda score : 54.871



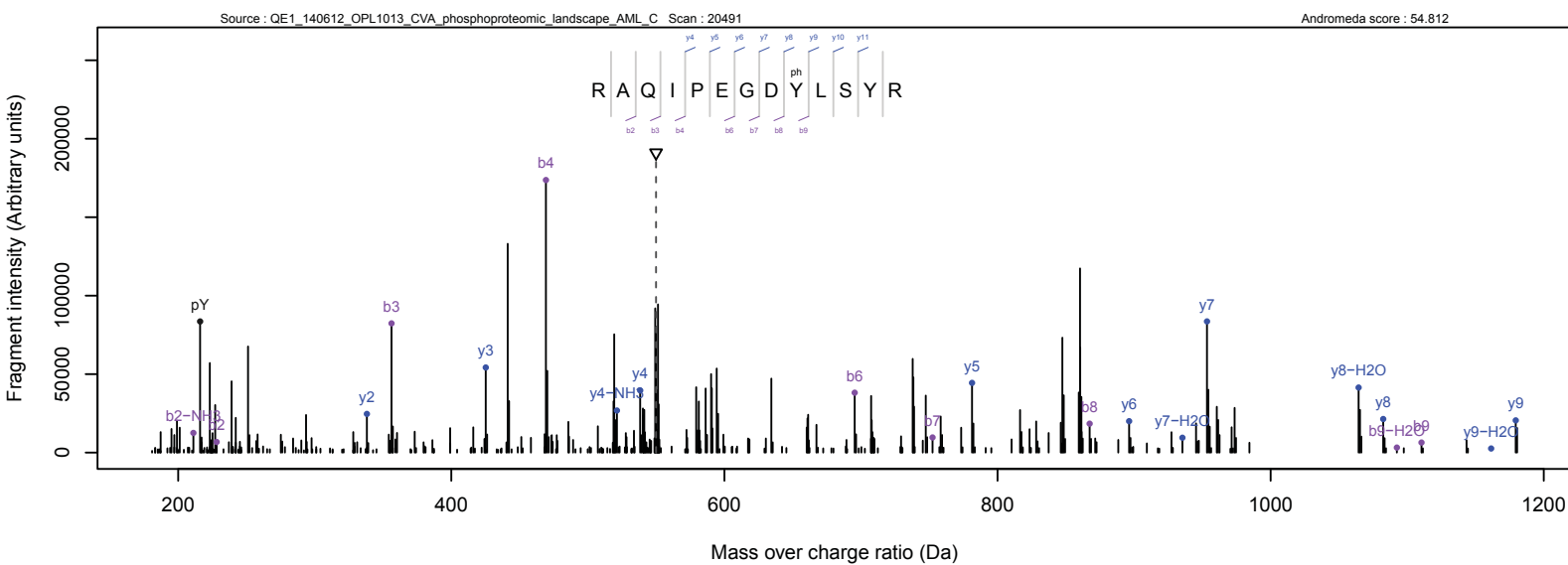
## KAGNFY(ph)VPAEPK



## GTVRDY(ph)PDFSPSVDAAEIQK



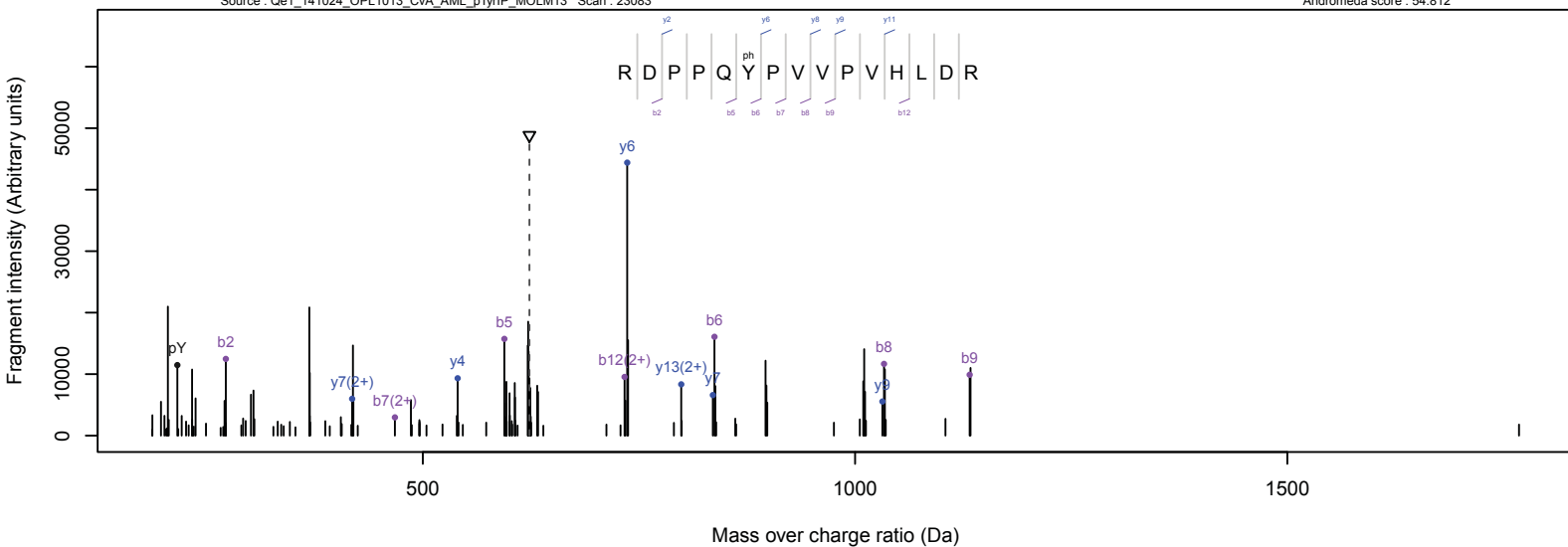
## RAQIEGDY(ph)LSYR



## RDPPQY(ph)PVVPHLDR-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 23083

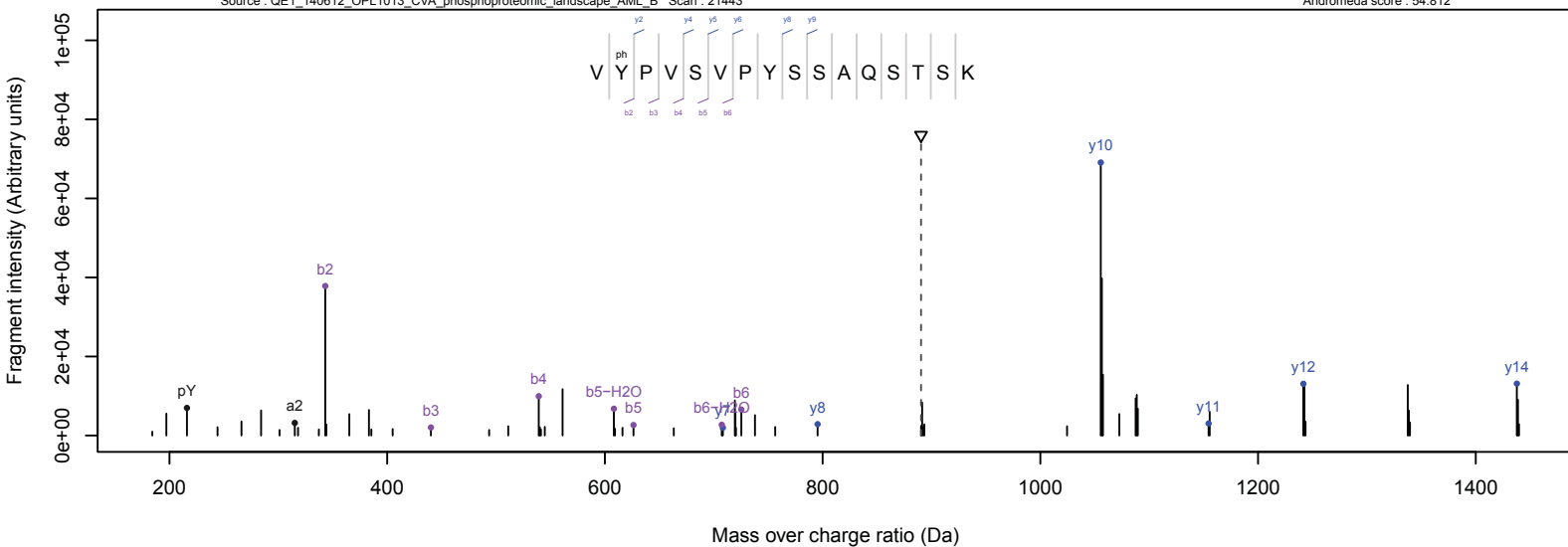
Andromeda score : 54.812



## VY(ph)PVSVPYSSAQSTSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21443

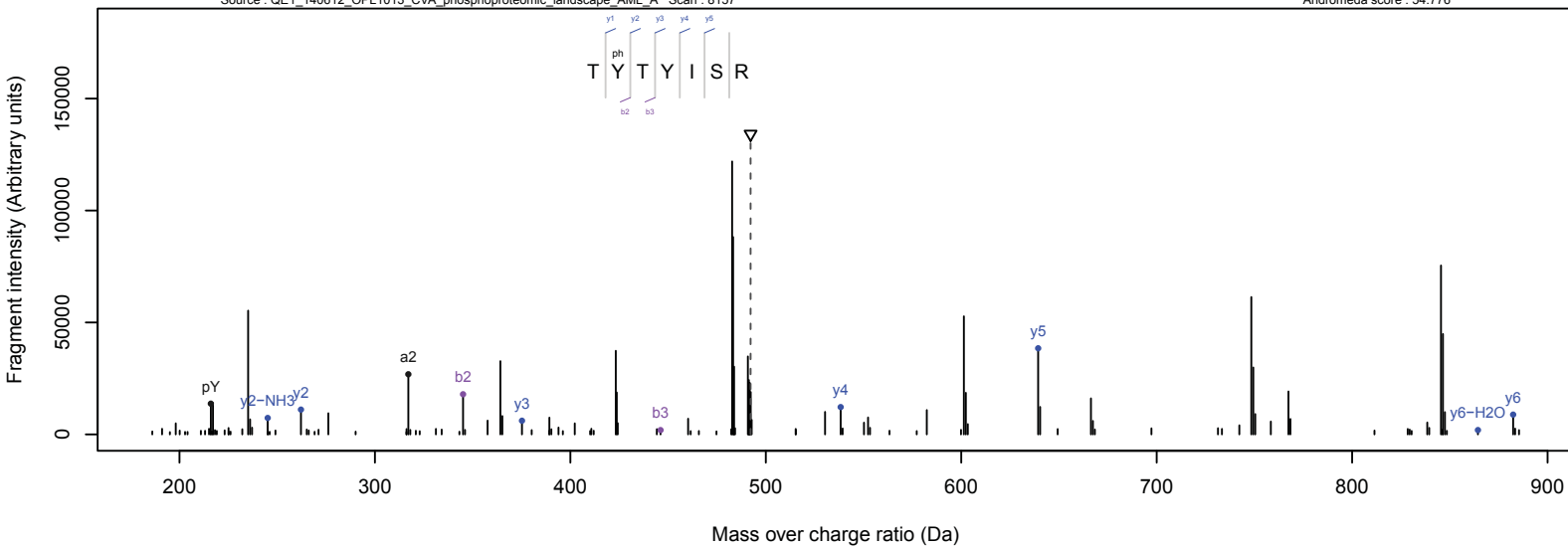
Andromeda score : 54.812



## TY(ph)TYISR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 8157

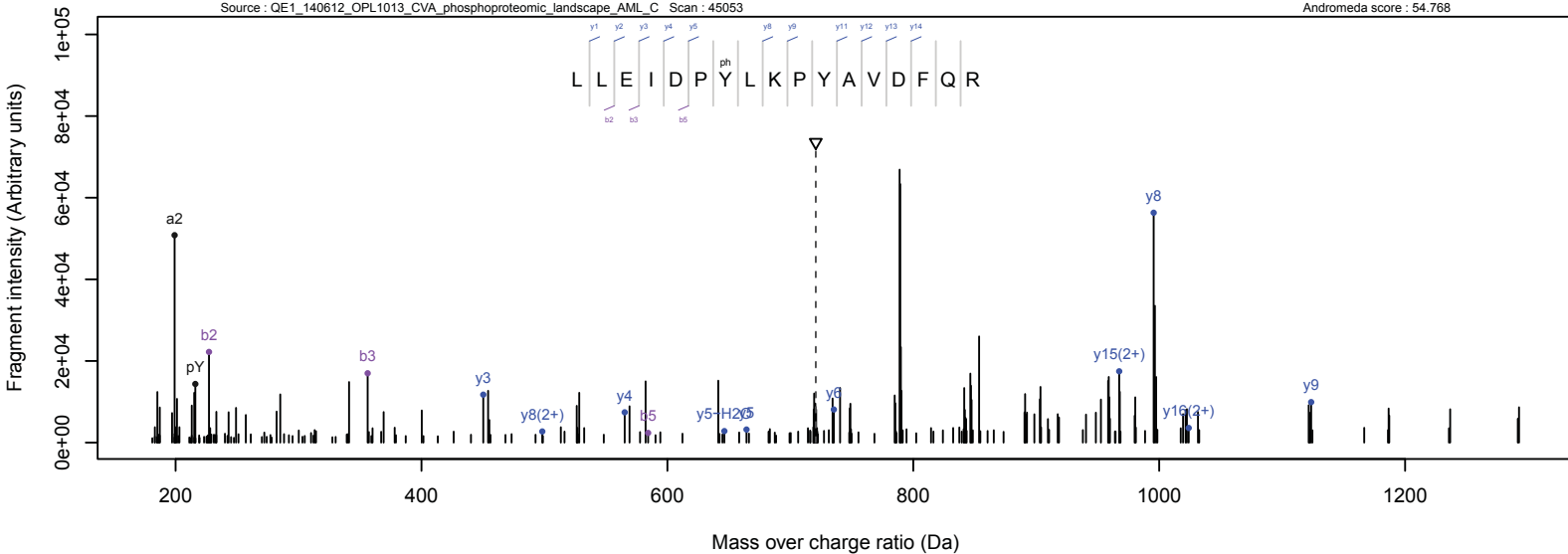
Andromeda score : 54.776



## LLEIDPY(ph)LKPYAVDFQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45053

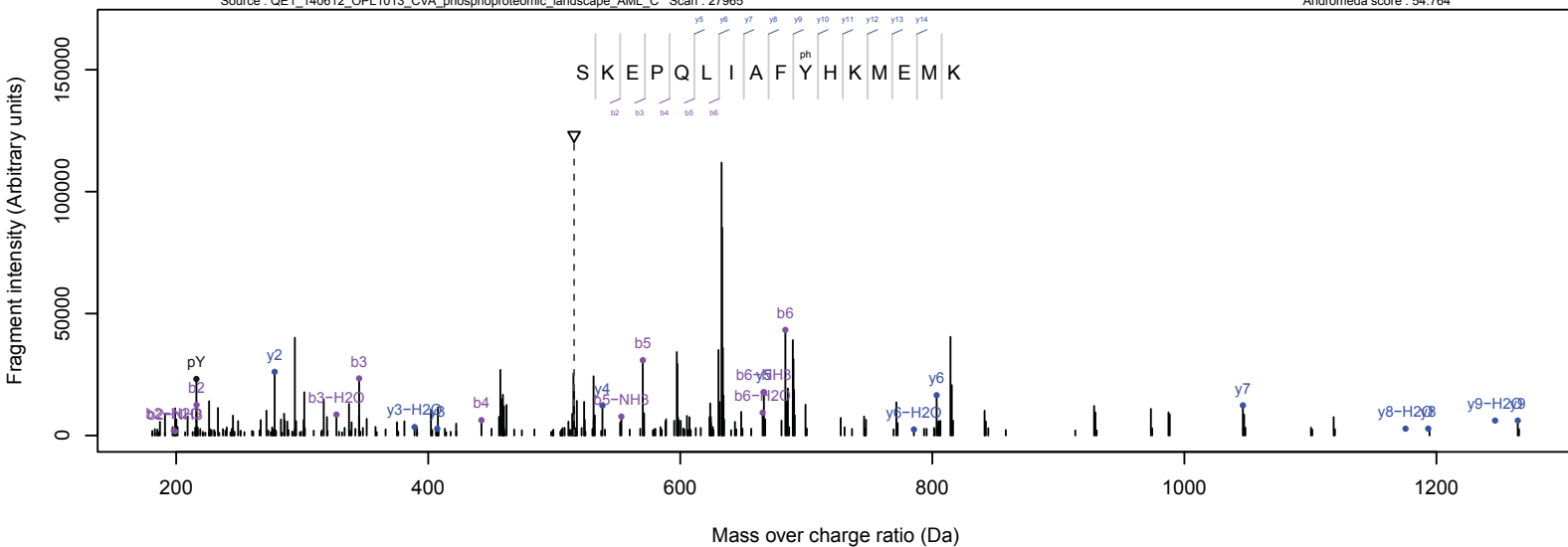
Andromeda score : 54.768



## SKEPQLIAFY(ph)HKMEMK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27965

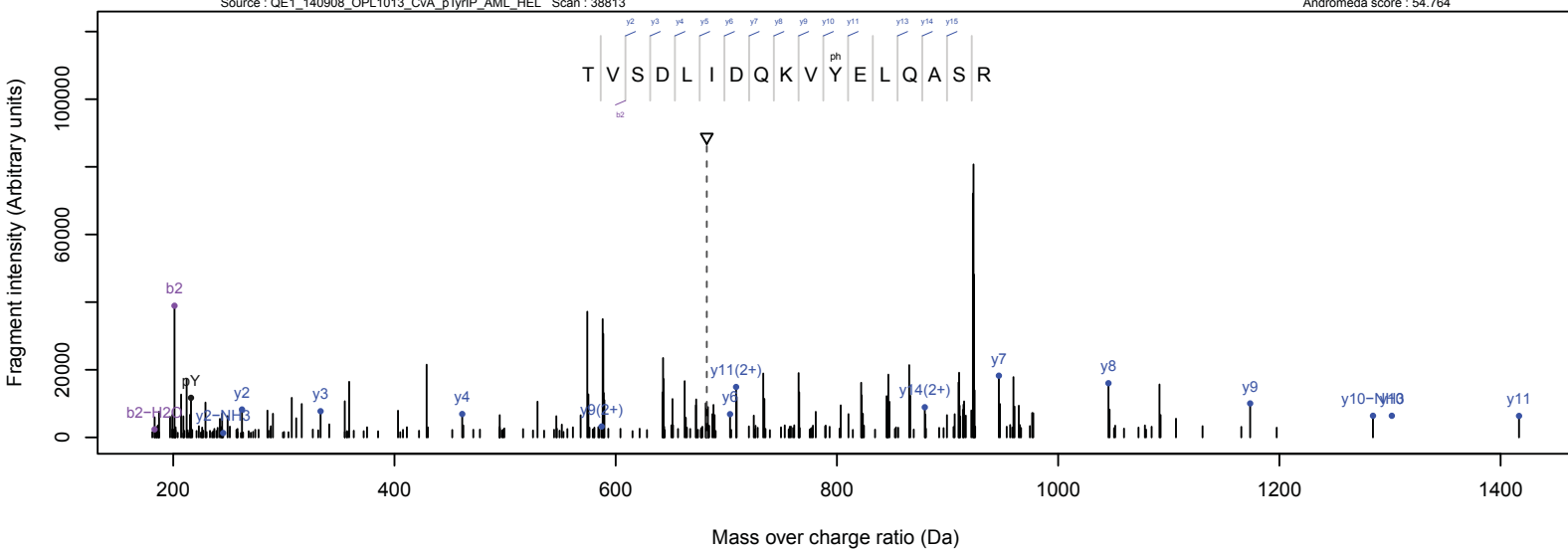
Andromeda score : 54.764



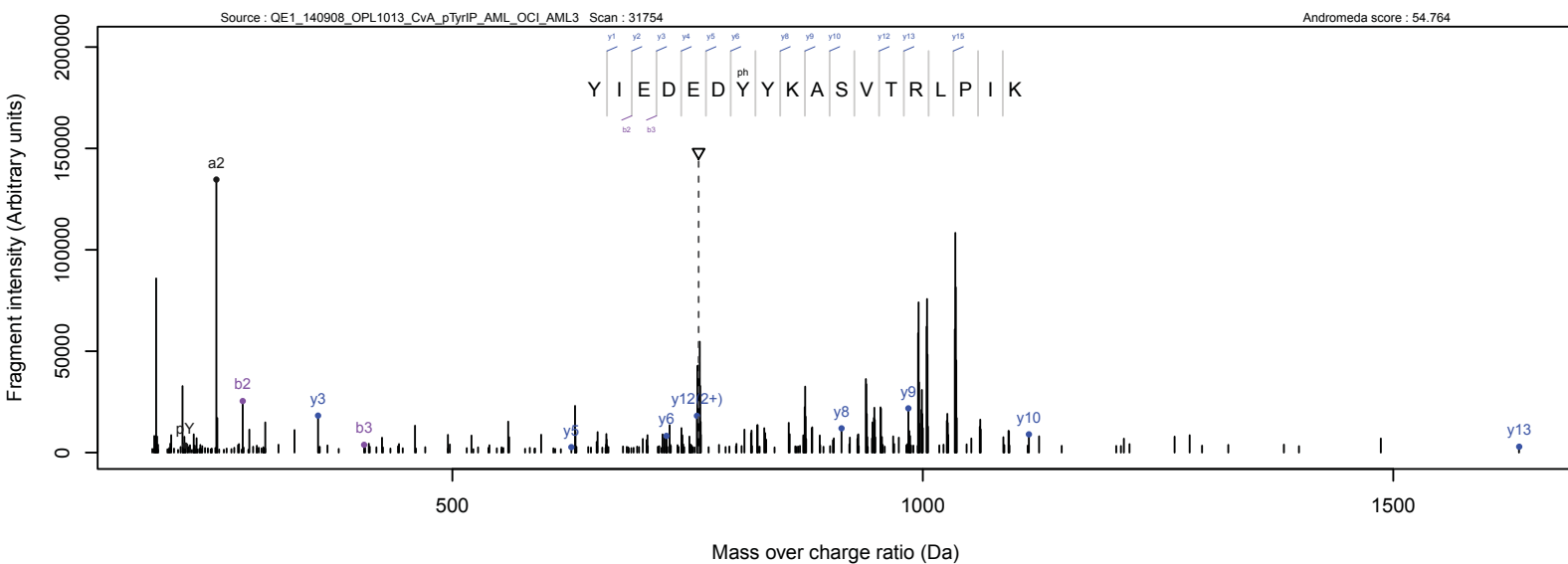
## TVSDLIQKVV(ph)ELQASR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 38813

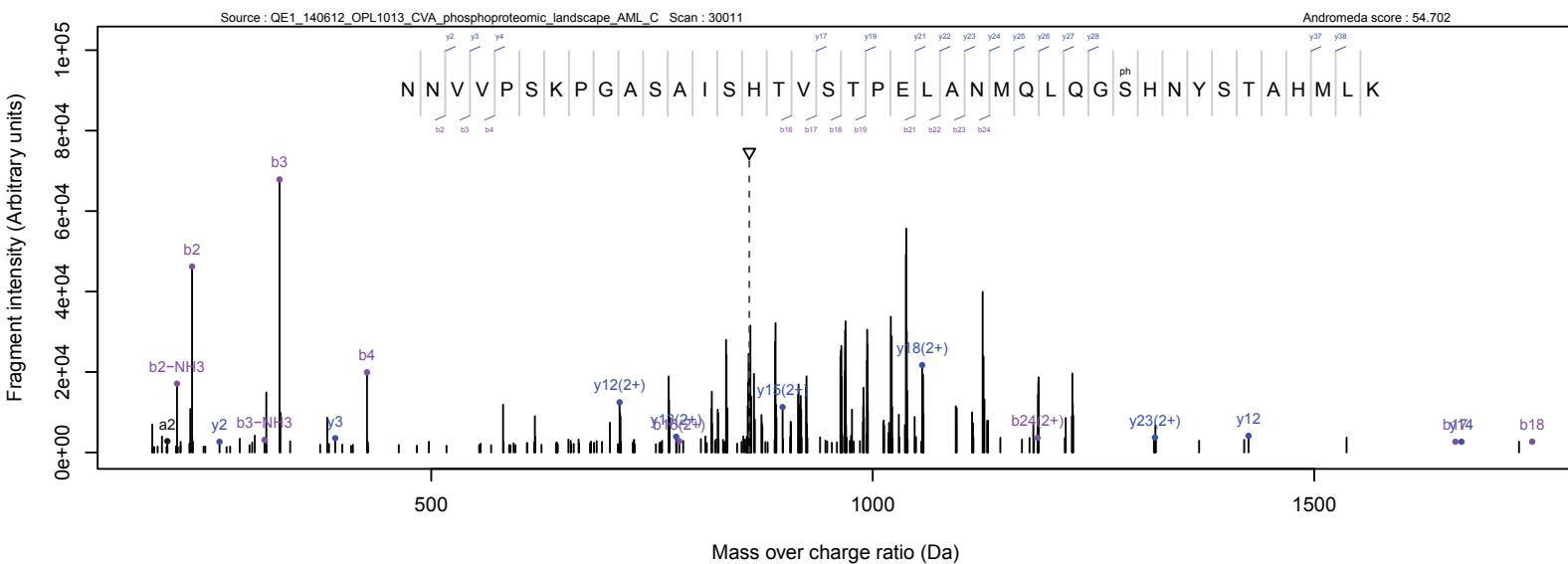
Andromeda score : 54.764



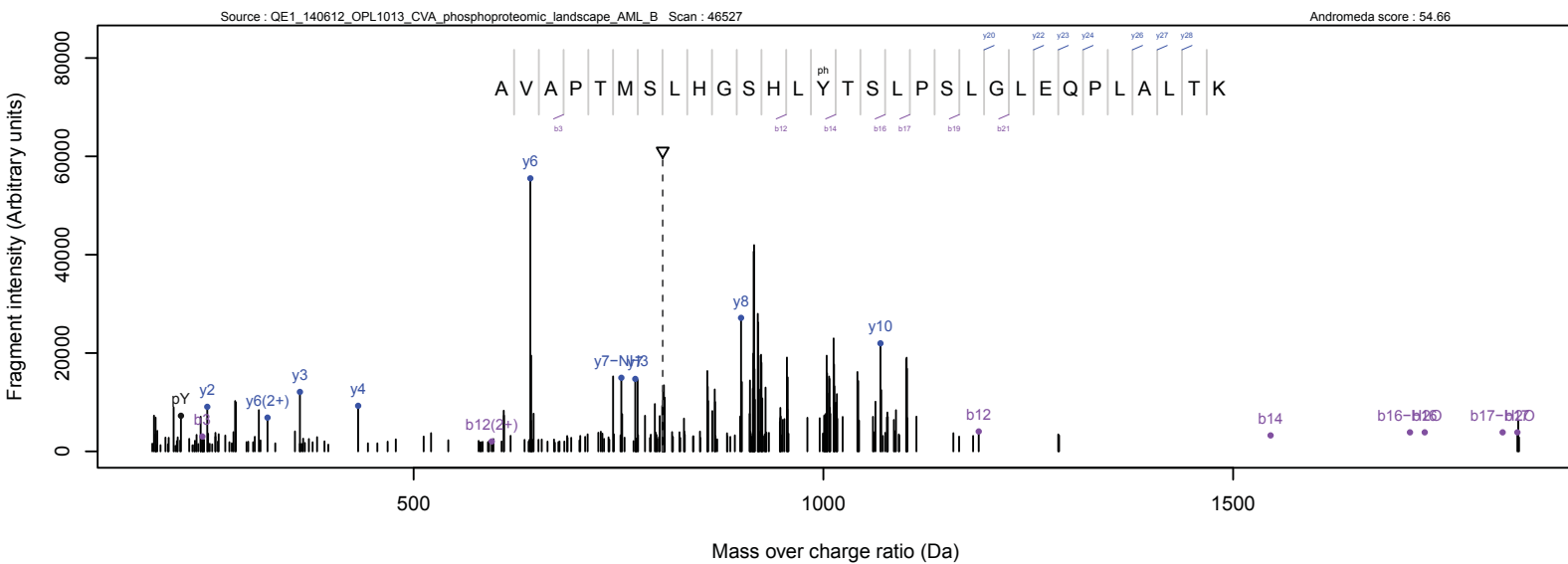
**YIEDEDY(ph)YKASVTRLPIK-3+**



**NNVPSKPGASAIHTVSTPELANMQLQGS(ph)HNYSTAHMLK-5+**



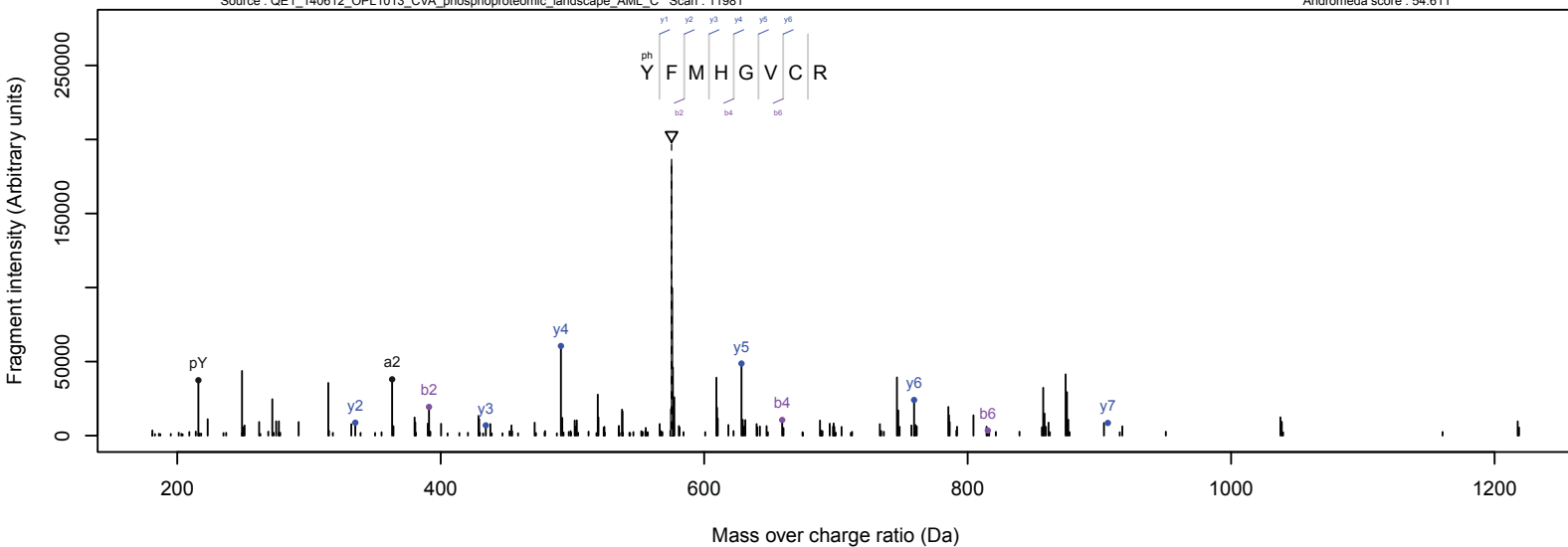
**AVAP TMSLHGSHLY(ph)TSLPSLGLQPLALTK-4+**



## Y(ph)FMHGVCR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11981

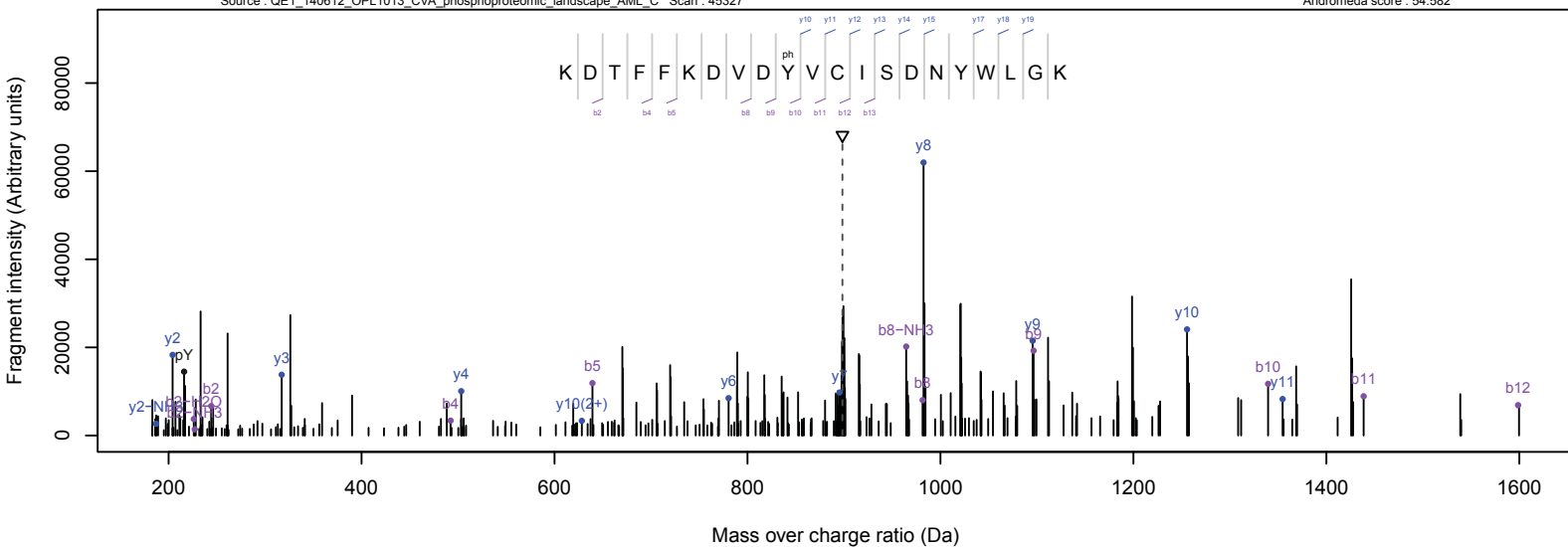
Andromeda score: 54.611



## KDTFFKDVDY(ph)VCISDNYWLGK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 45327

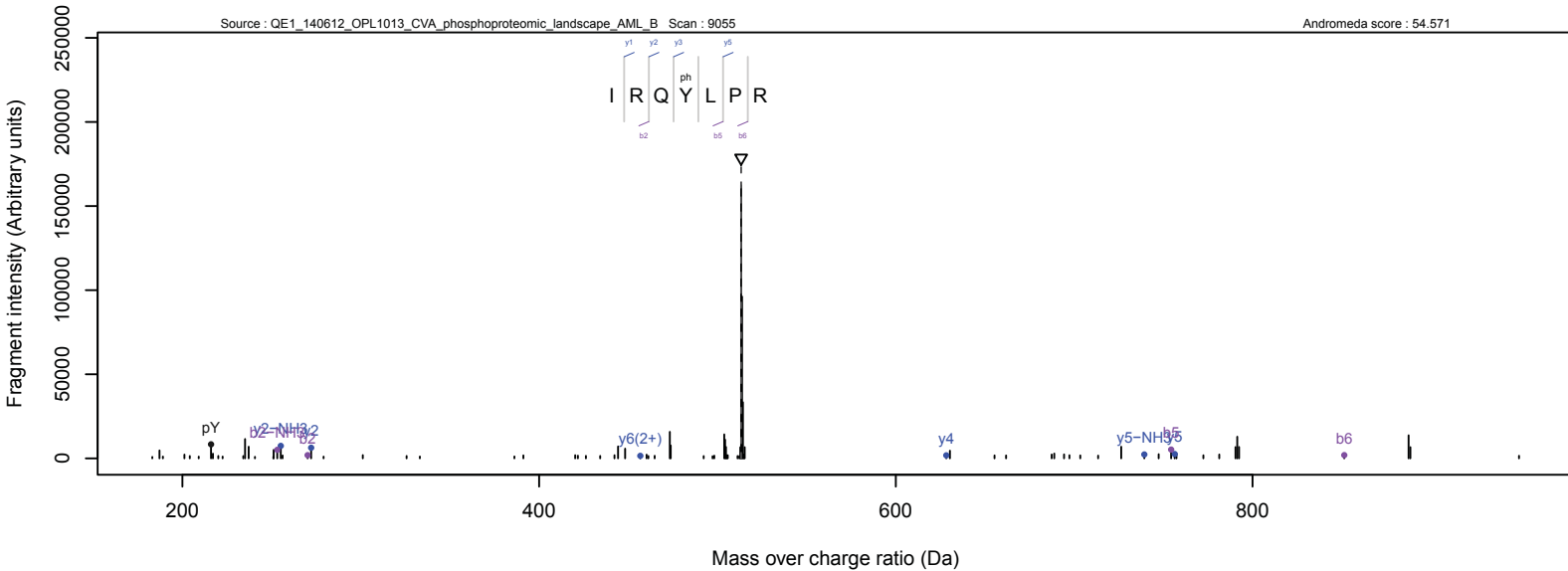
Andromeda score: 54.582



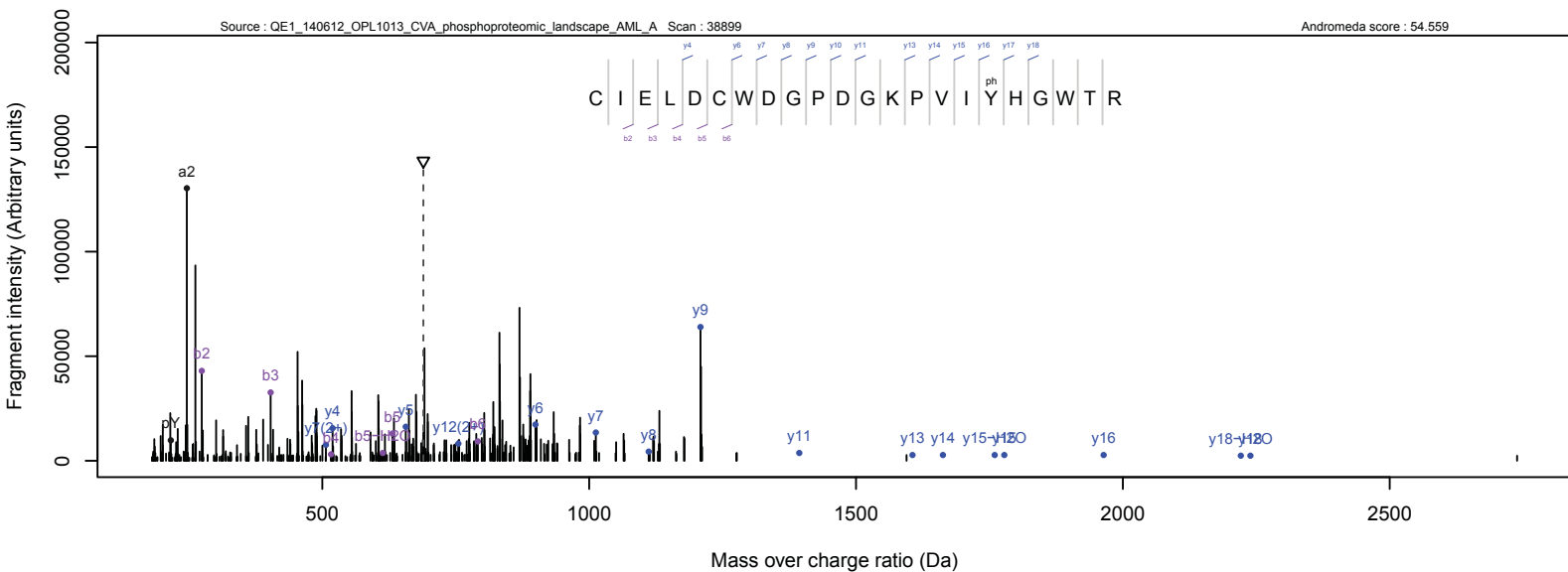
## IRQY(ph)LPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 9055

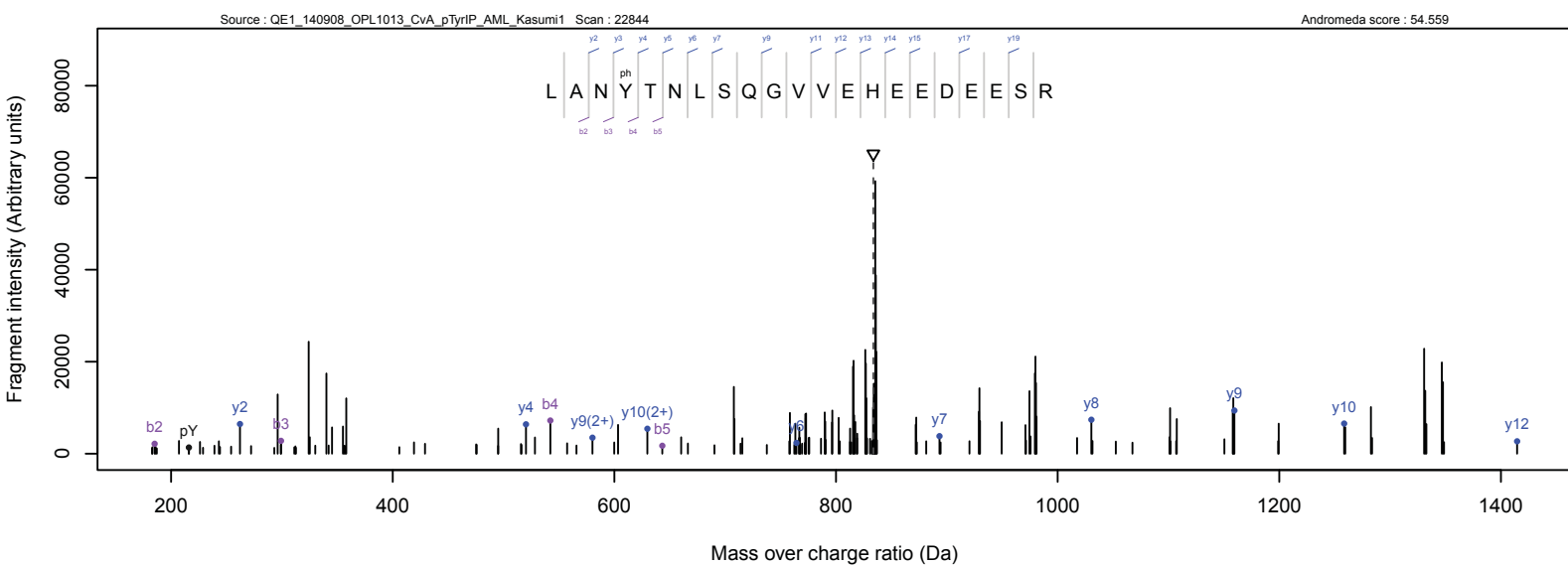
Andromeda score: 54.571



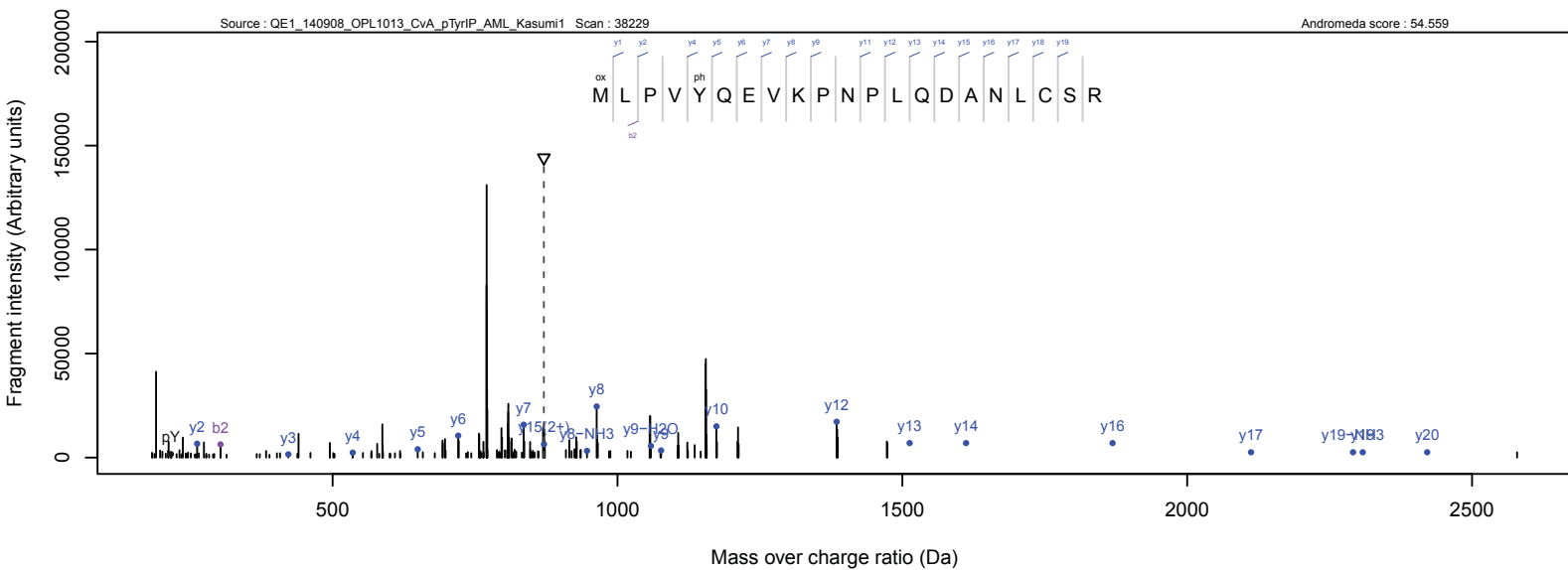
**CIELDCWDGPKVIY(ph)HGWTR-4+**



**LANY(ph)TNLSQGVVEHEEESR-3+**



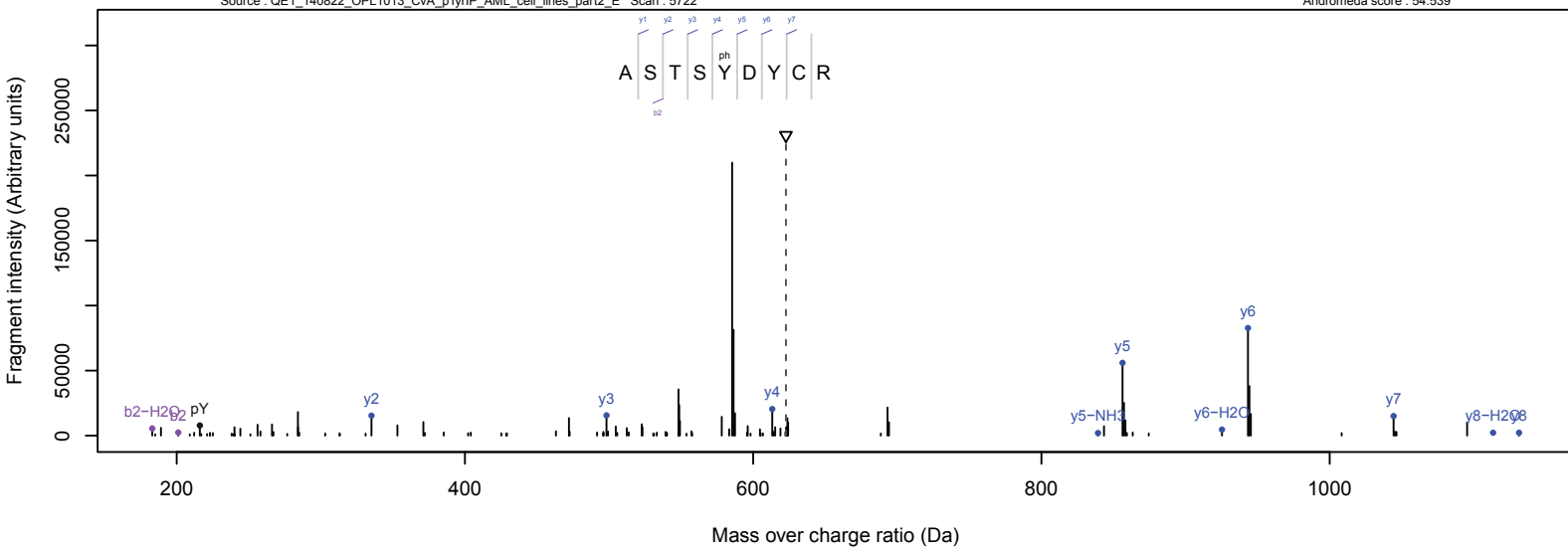
**(ac)M(ox)LPVY(ph)QEVKPNPLQDANLCSR-3+**



## (ac)ASTSY(ph)DYCR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 5722

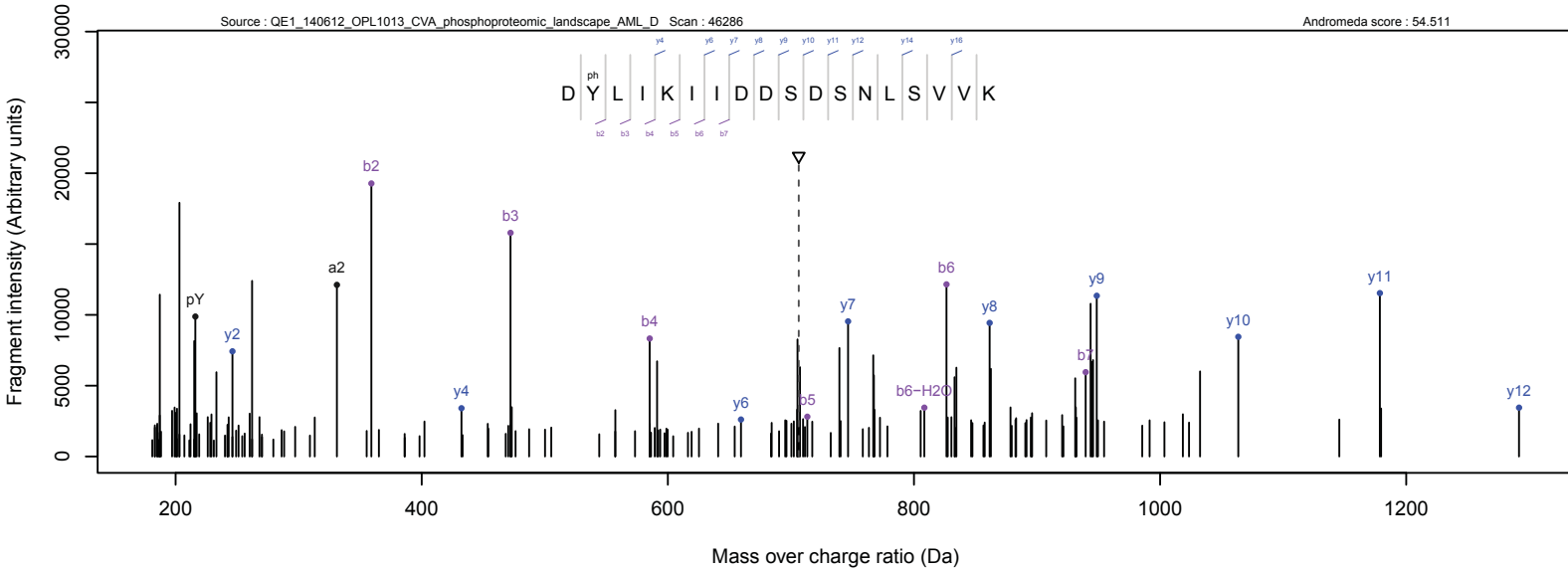
Andromeda score: 54.539



## DY(ph)LIKIIDSDSNLSVVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 46286

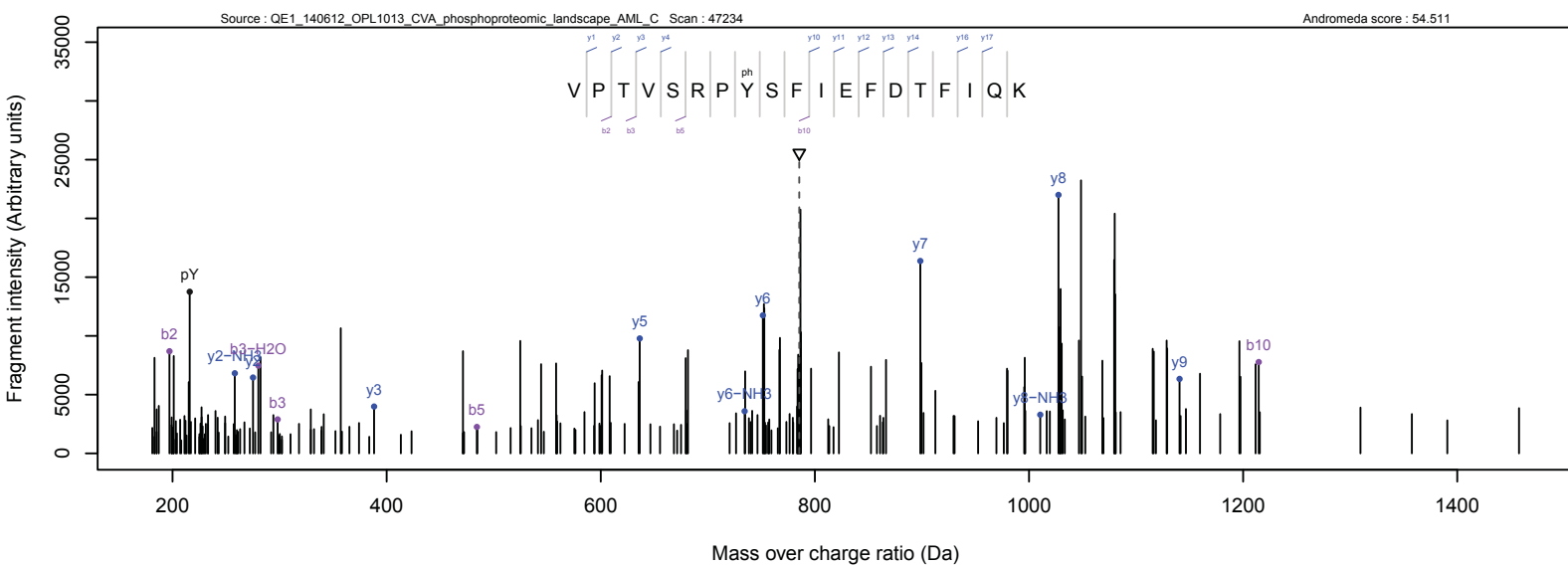
Andromeda score: 54.511



## VPTVSRPY(ph)SFIEFDTFIQK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 47234

Andromeda score: 54.511

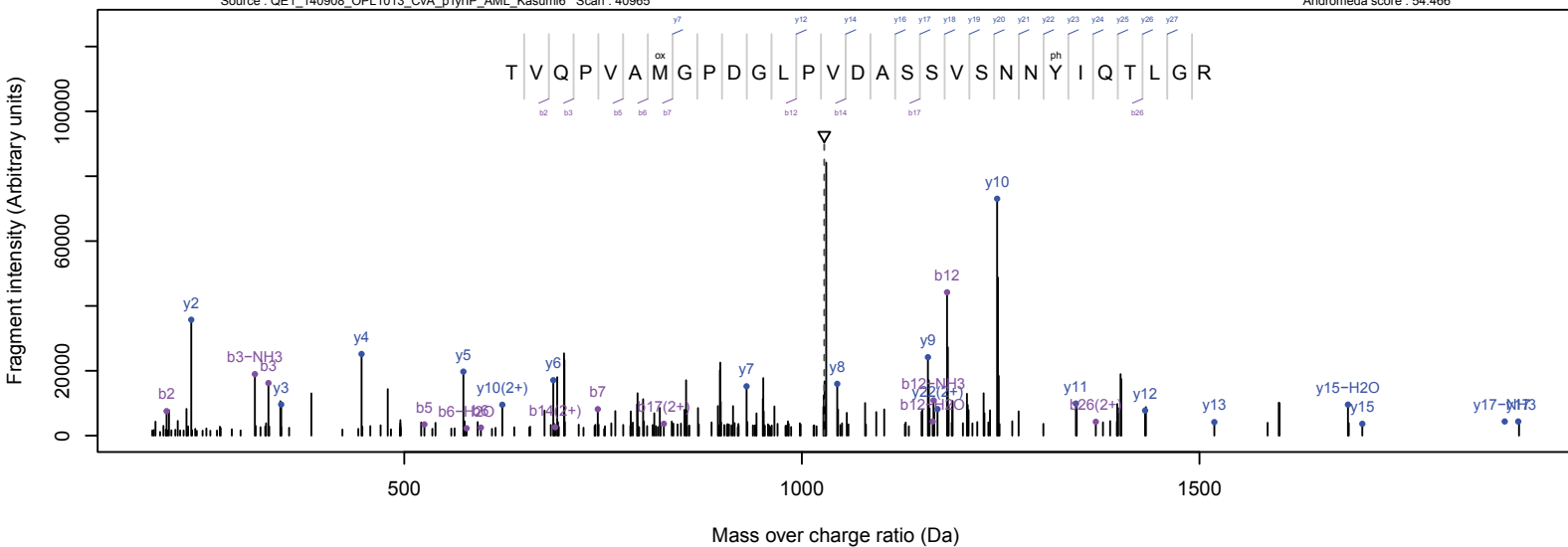




TVQPVAM(ox)GPDGLPVDASSVSNY(ph)IQLTGR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 40965

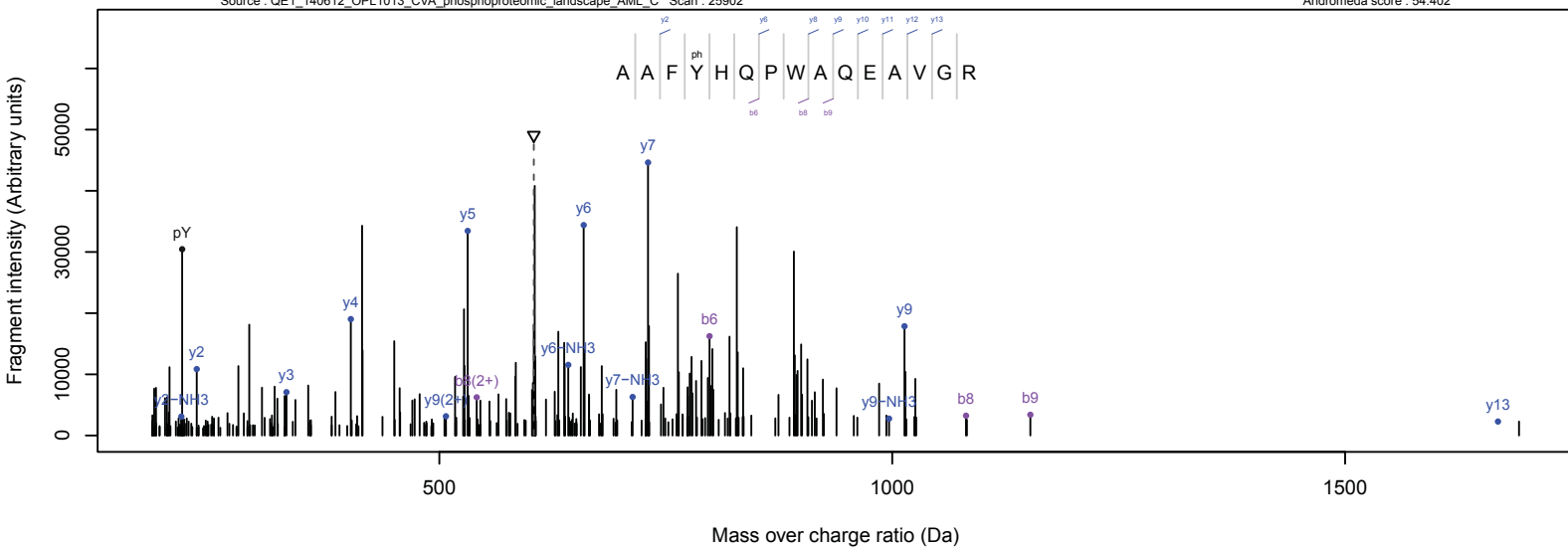
Andromeda score: 54.466



AAFY(ph)HQPWAQEA VGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25902

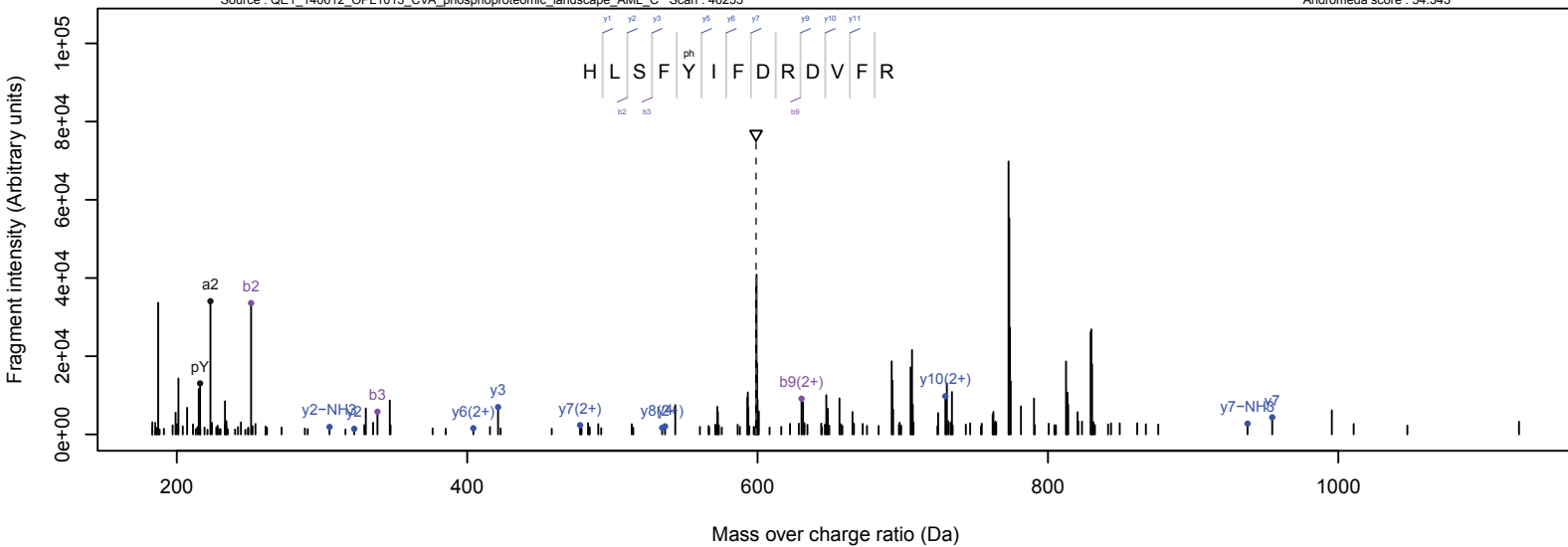
Andromeda score: 54.402



HLSFY(ph)IFDRD VFR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 46255

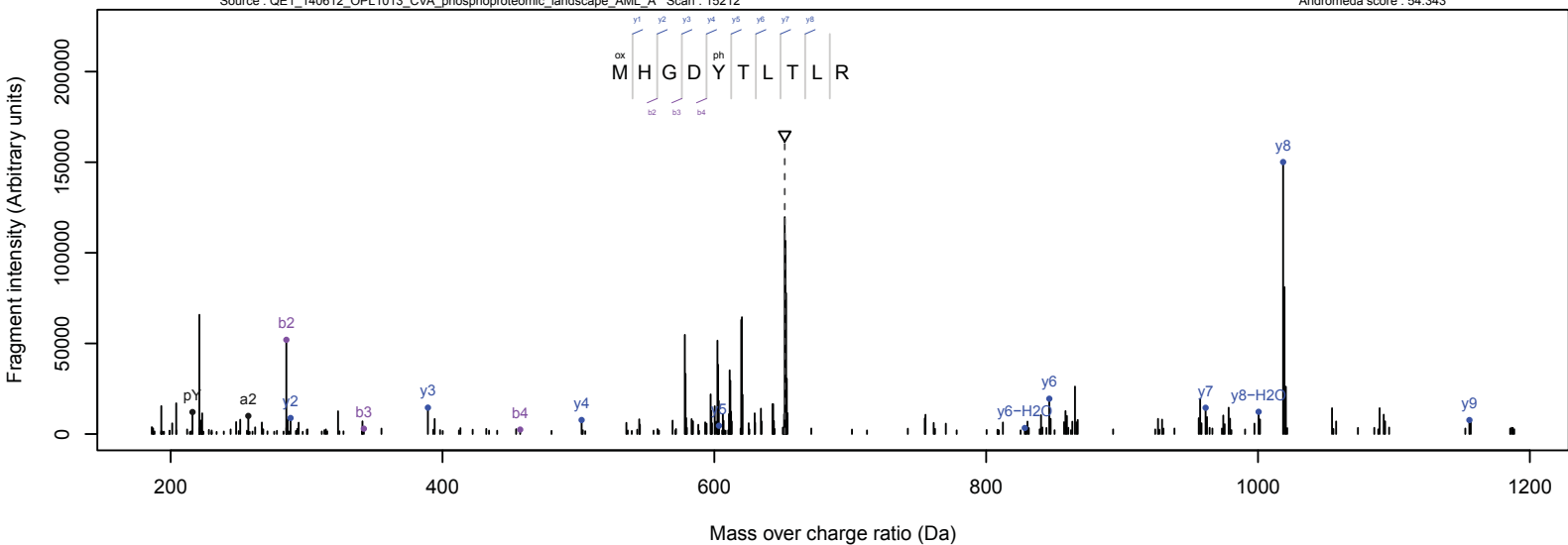
Andromeda score: 54.343



M(ox)HG DY(ph)TLTLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 15212

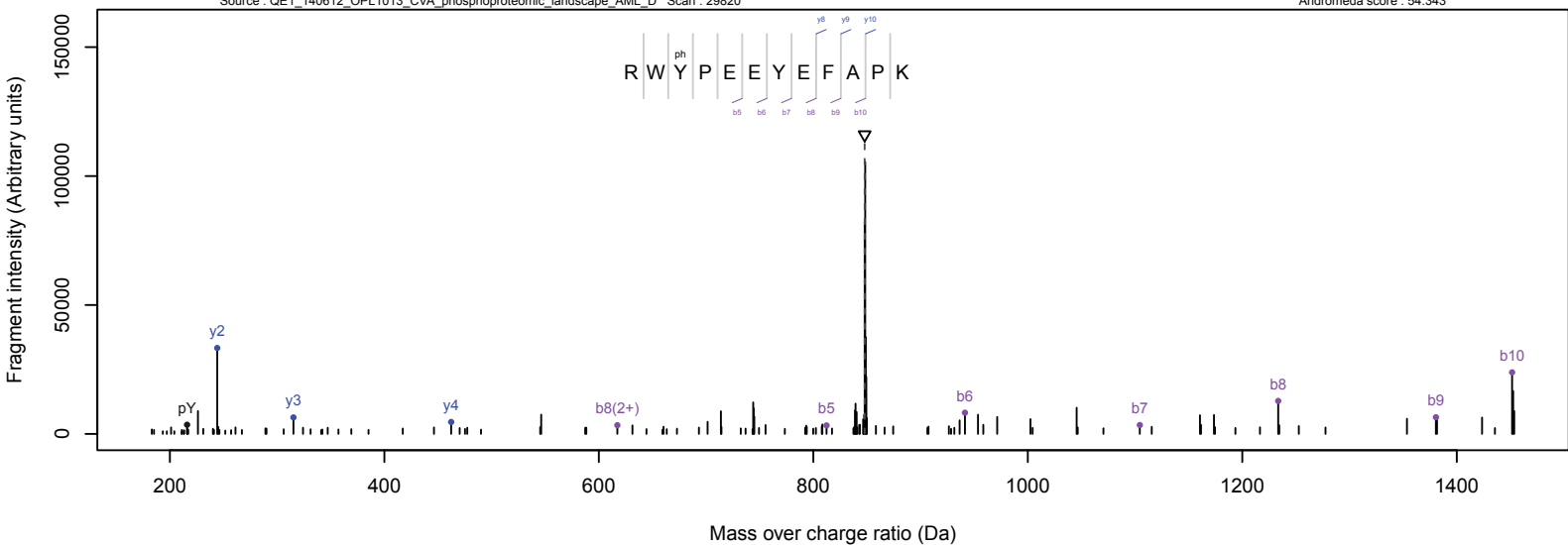
Andromeda score: 54.343



RWY(ph)PEEYEFAPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 29820

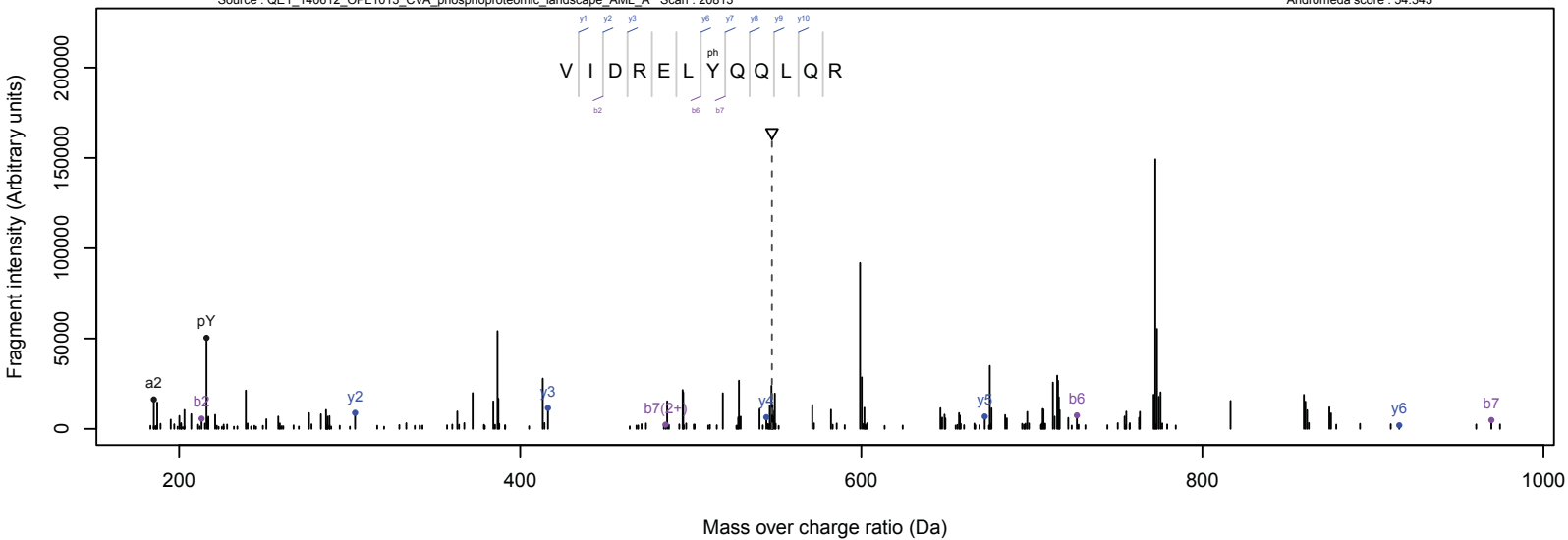
Andromeda score: 54.343



VIDRELY(ph)QLQR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 20813

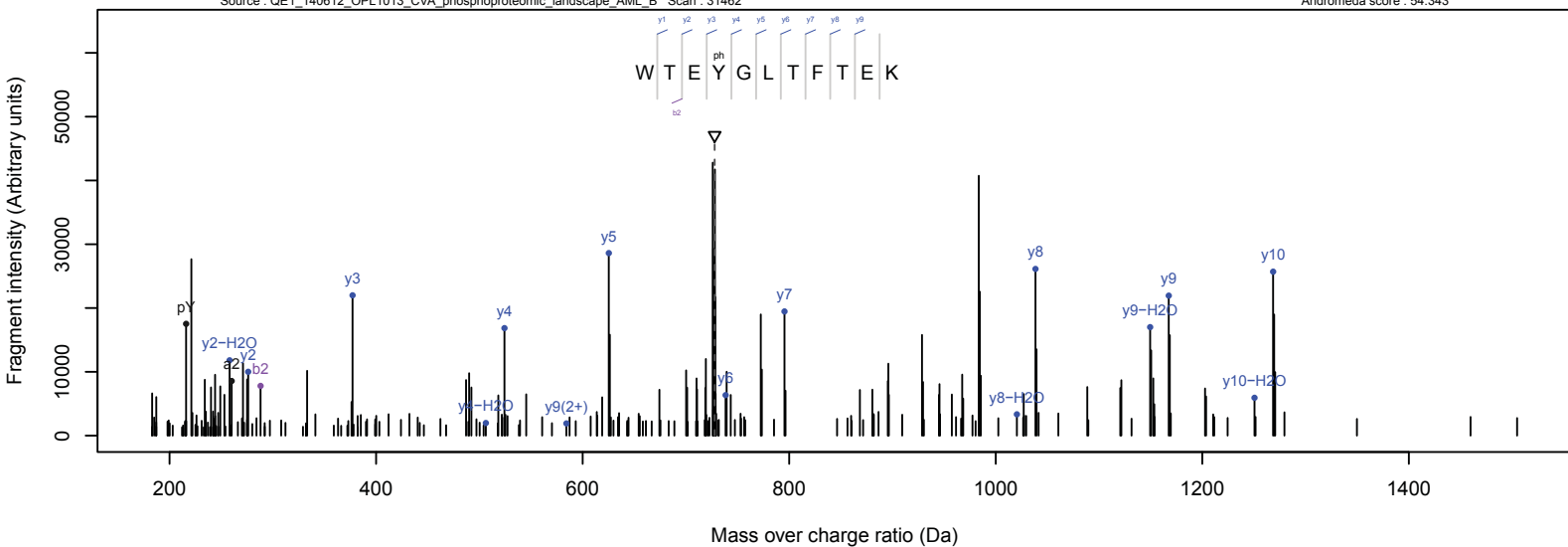
Andromeda score: 54.343



WTEY(ph)GLTFTEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31462

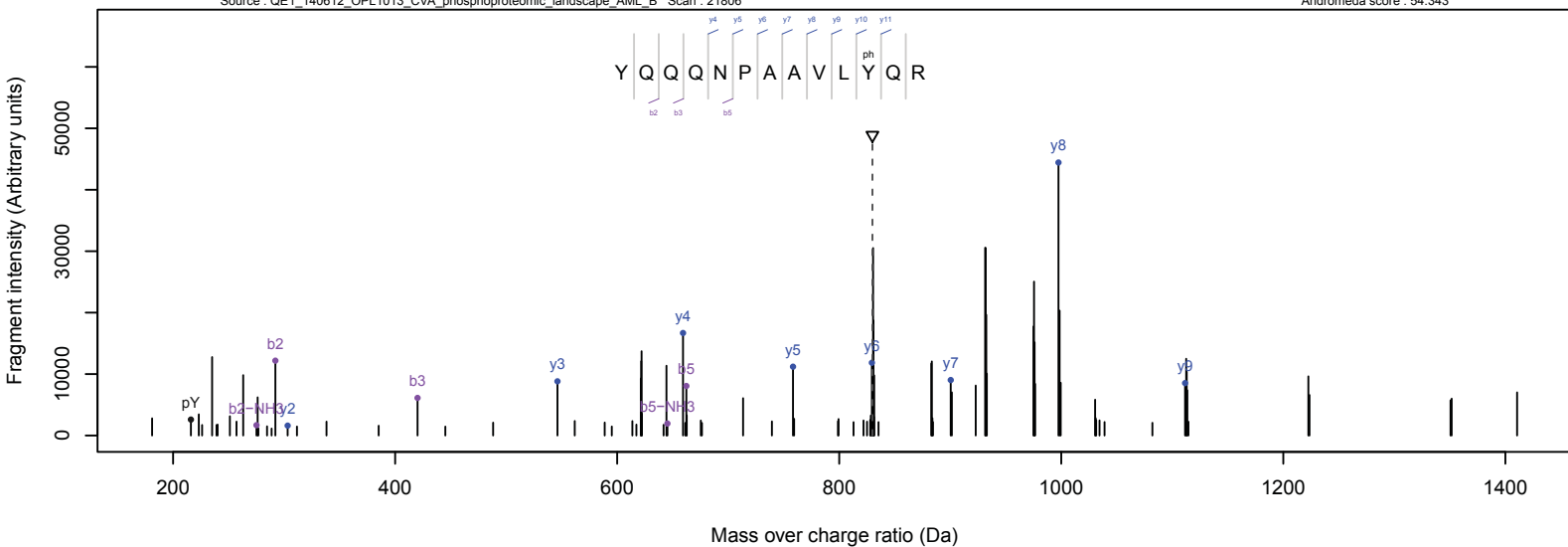
Andromeda score : 54.343



YQQNPAAVLY(ph)QR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21806

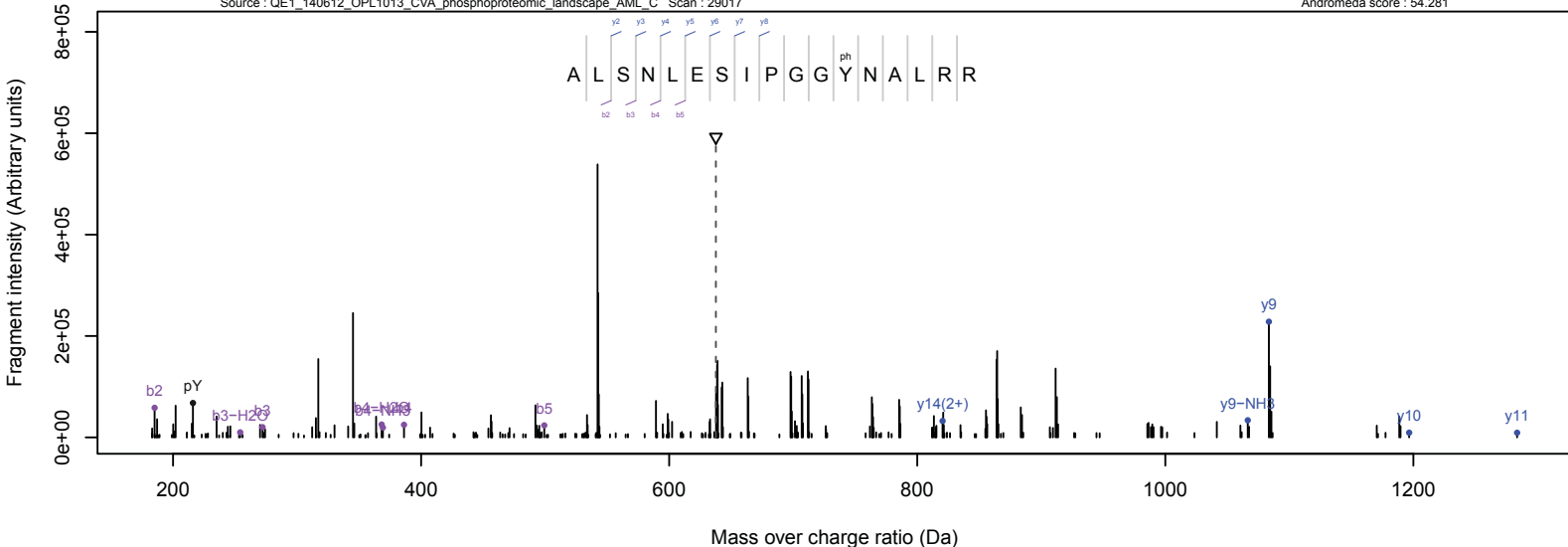
Andromeda score : 54.343



ALSNLEISIPGGY(ph)NALRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29017

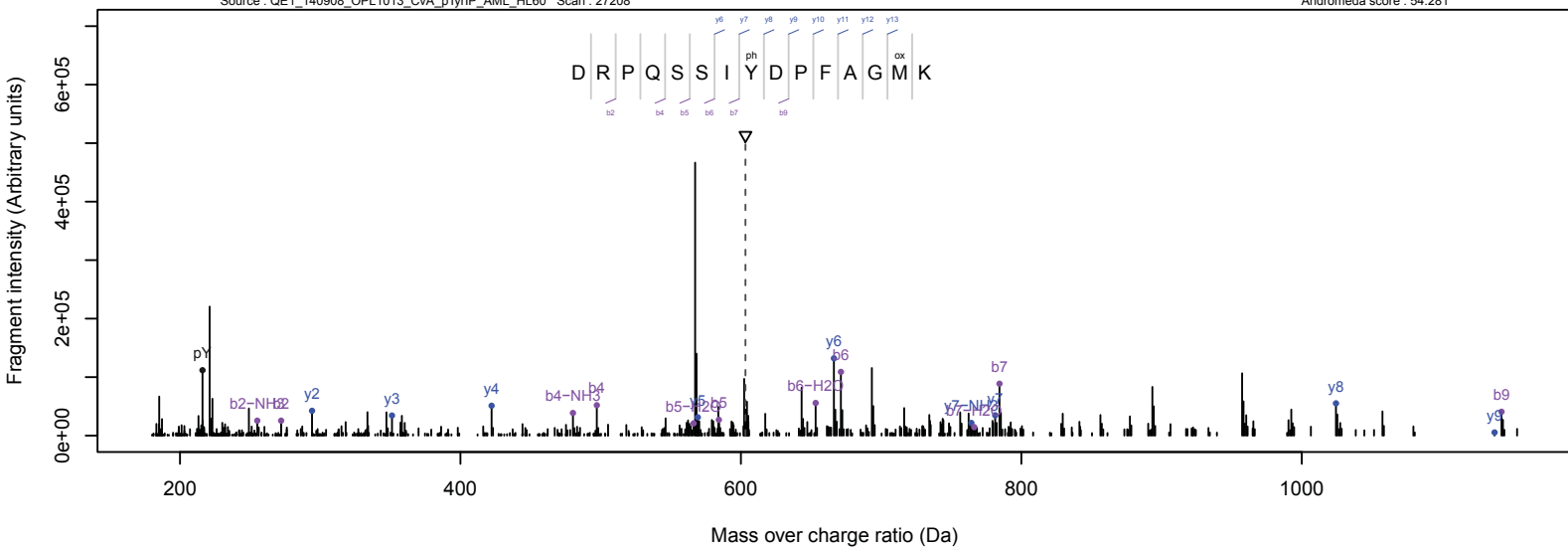
Andromeda score : 54.281



DRPQSSY(ph)DPFAGM(ox)K-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 27208

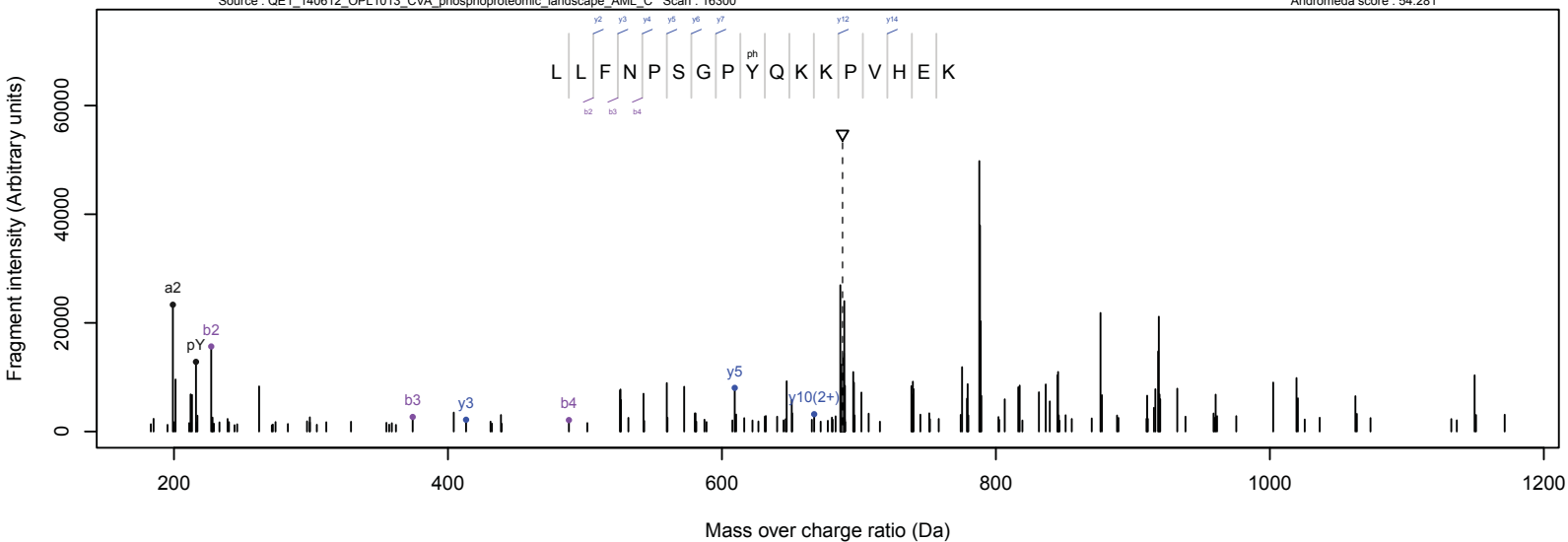
Andromeda score : 54.281



LLFNPSGPY(ph)QKPVHEK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 16300

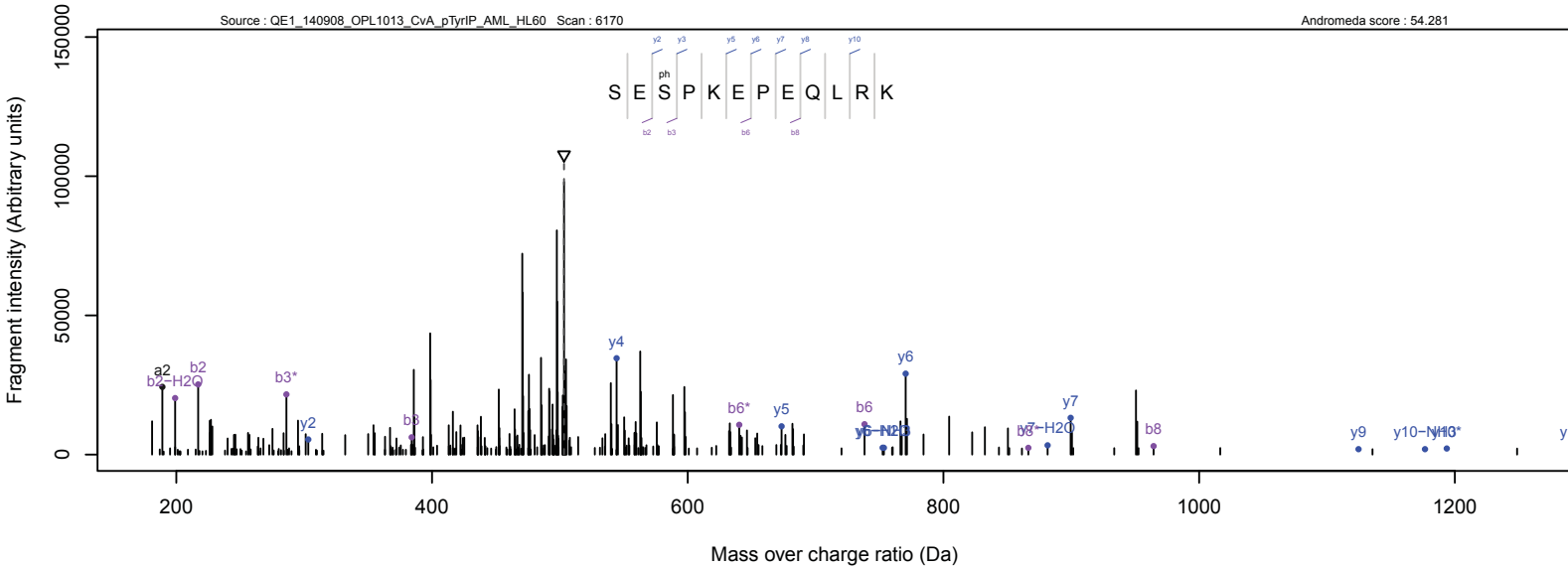
Andromeda score : 54.281



SES(ph)PKEPEQLRK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 6170

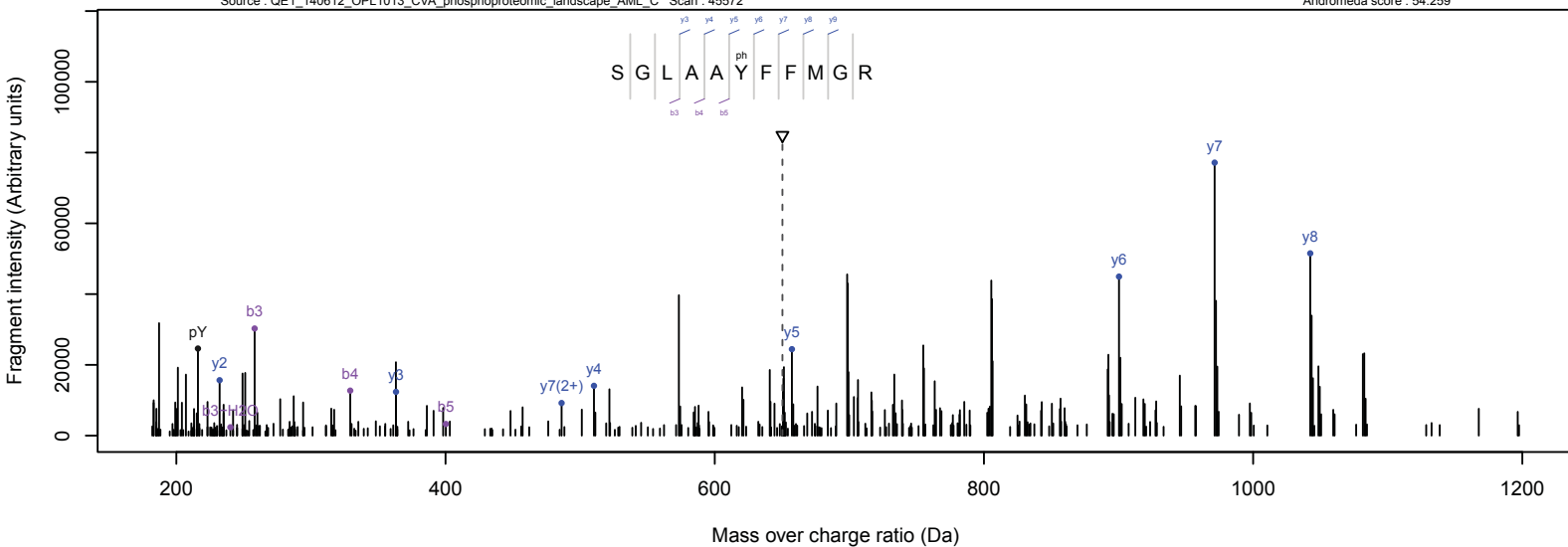
Andromeda score : 54.281



## SGLAAY(ph)FFMGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45572

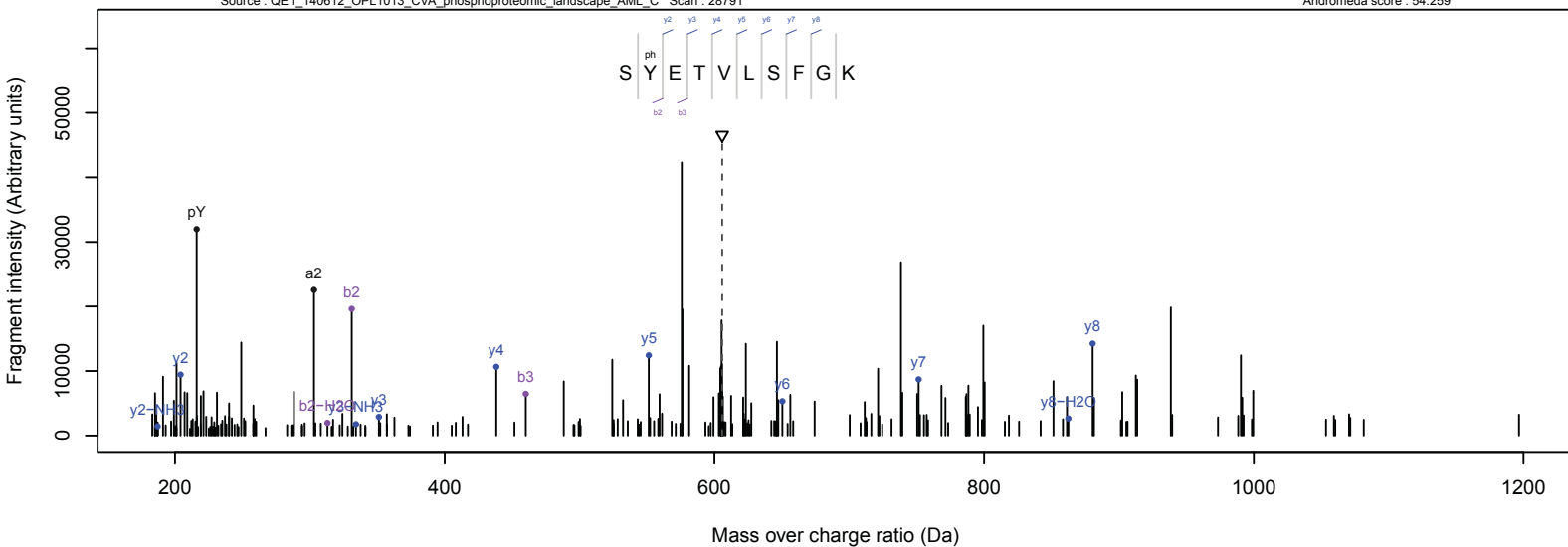
Andromeda score : 54.259



## SY(ph)ETVLSFGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28791

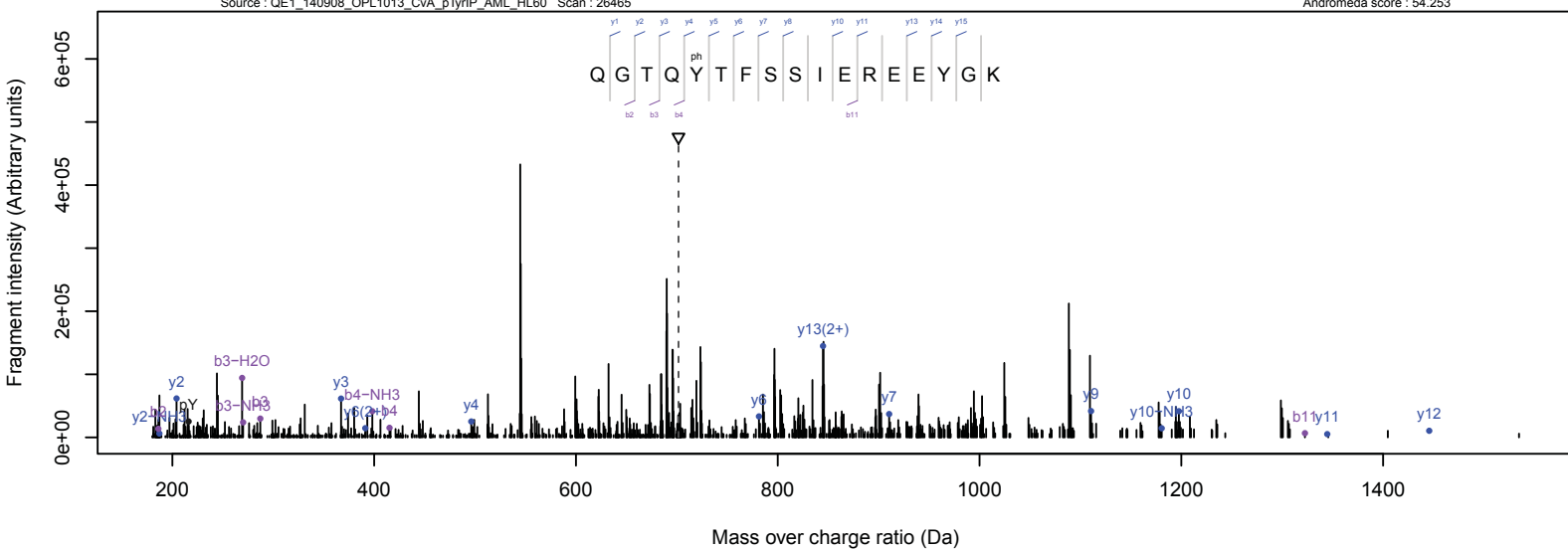
Andromeda score : 54.259



## QGTQY(ph)TFSSIEREEYGK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 26465

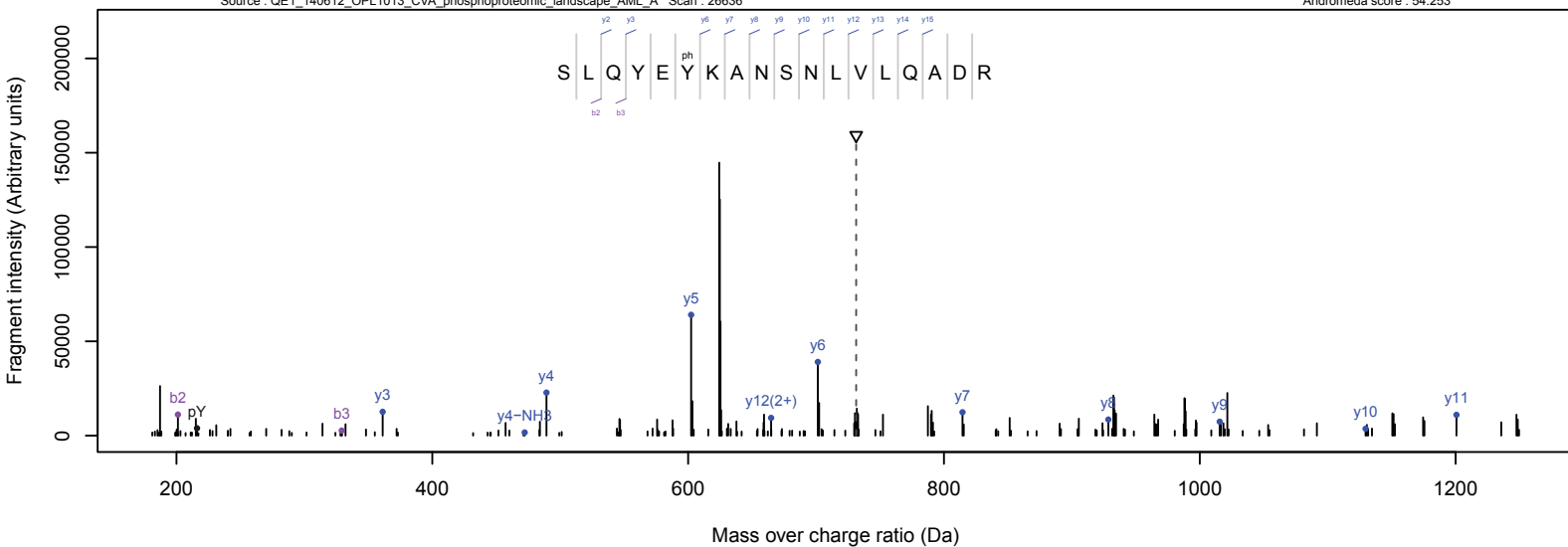
Andromeda score : 54.253



SLQY<sup>EY</sup>(ph)KANSNLVLQADR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26636

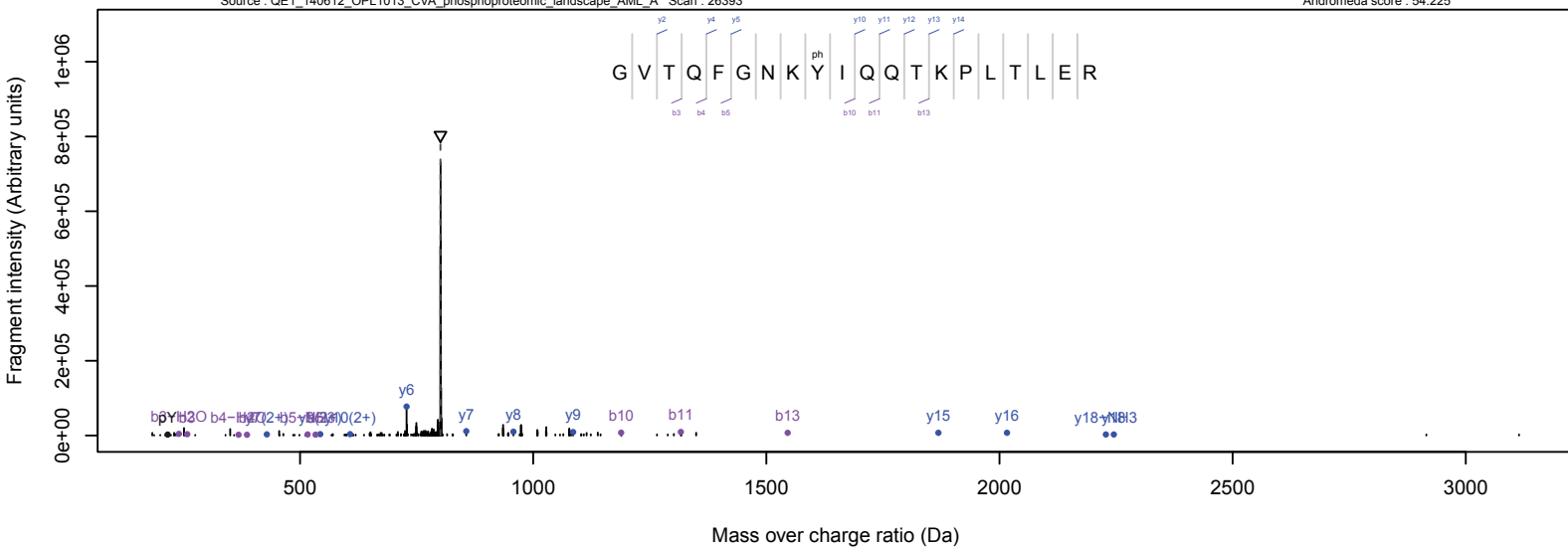
Andromeda score : 54.253



GVTQFGNK<sup>Y</sup>(ph)IQQT<sup>K</sup>PLTLER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26393

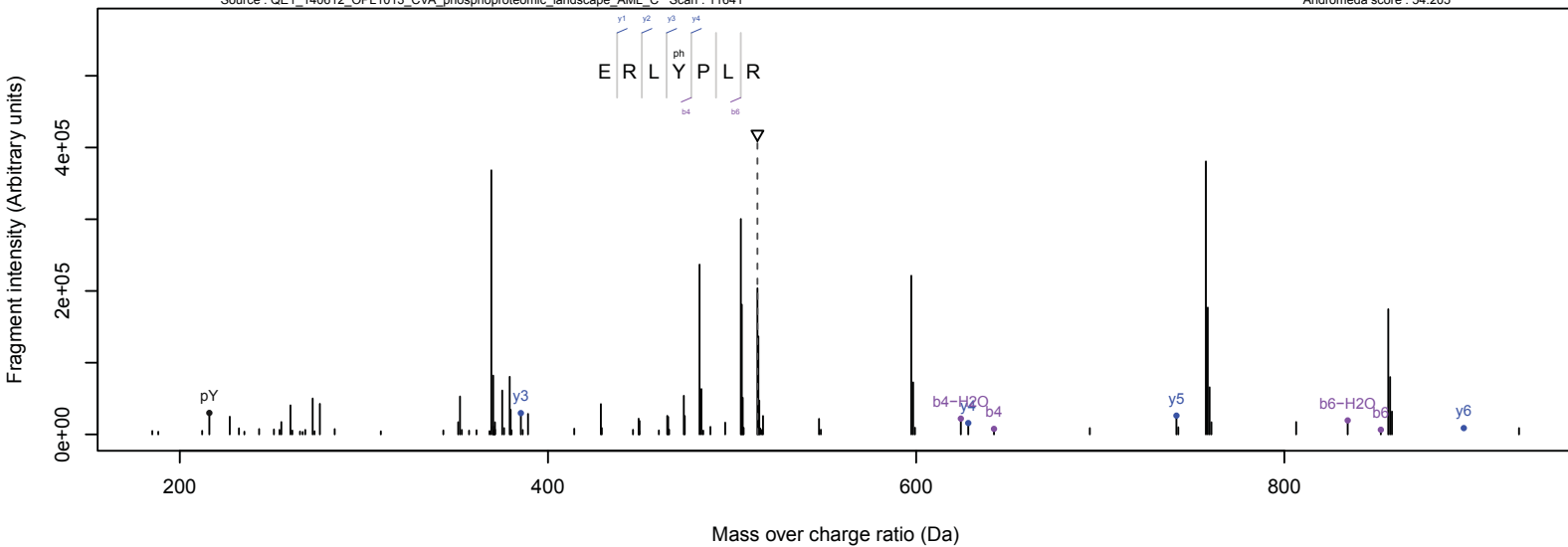
Andromeda score : 54.225



ERLY<sup>Y</sup>(ph)PLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11641

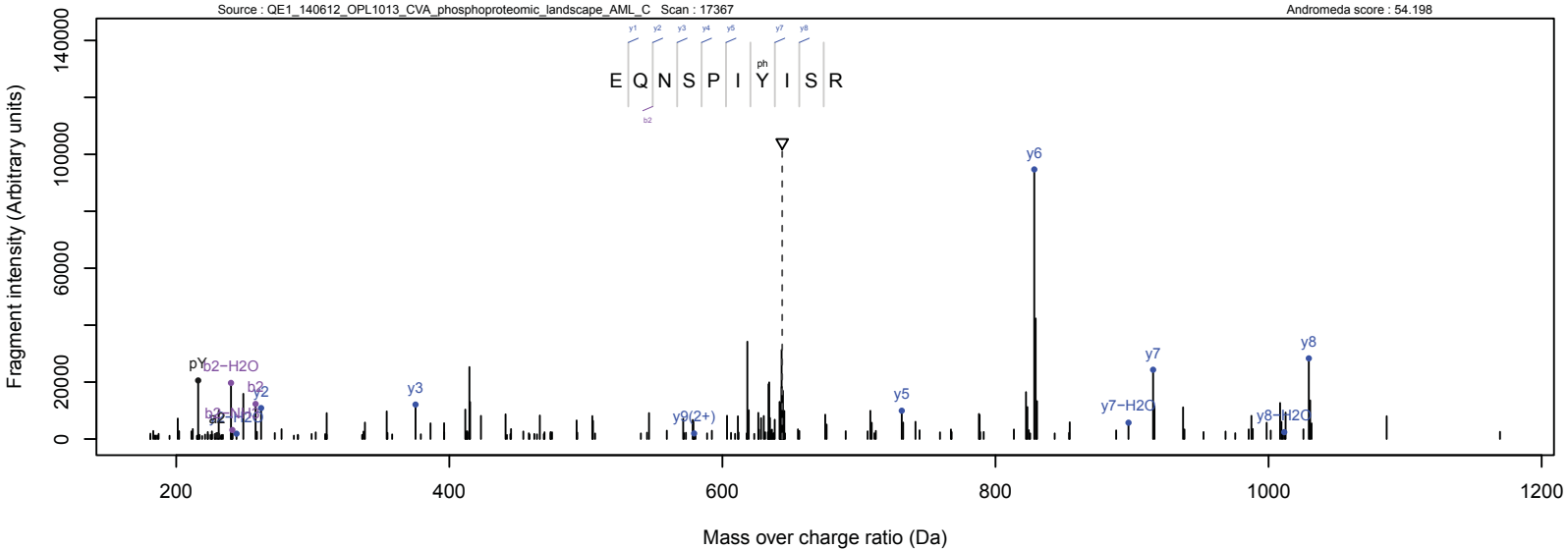
Andromeda score : 54.205



**EQNSPIY(ph)ISR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17367

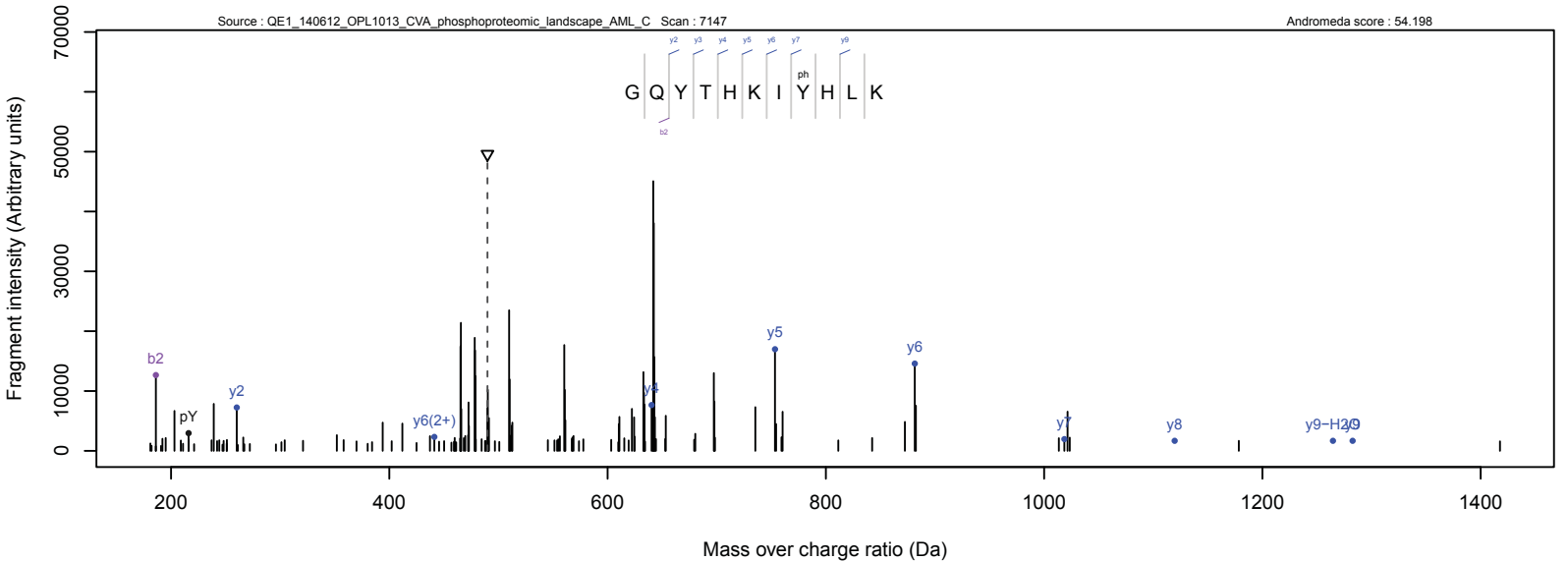
Andromeda score : 54.198



**GQYTHKIY(ph)HLK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7147

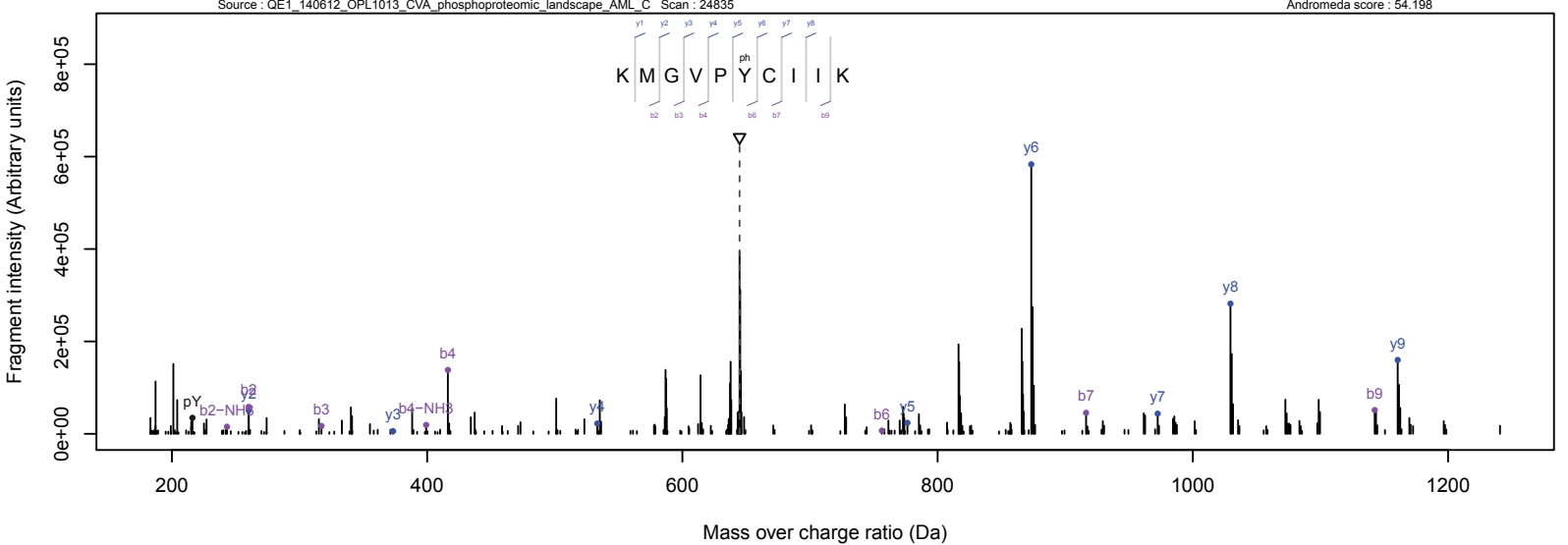
Andromeda score : 54.198



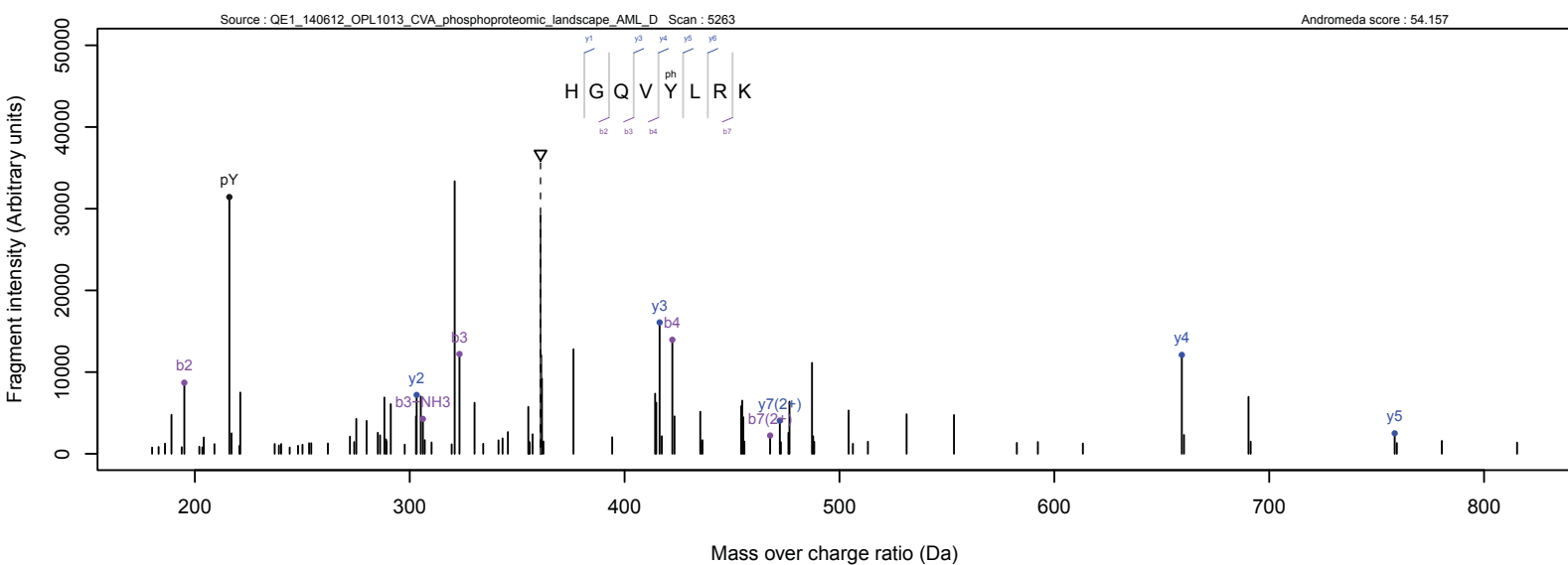
**KMGVPY(ph)CIK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24835

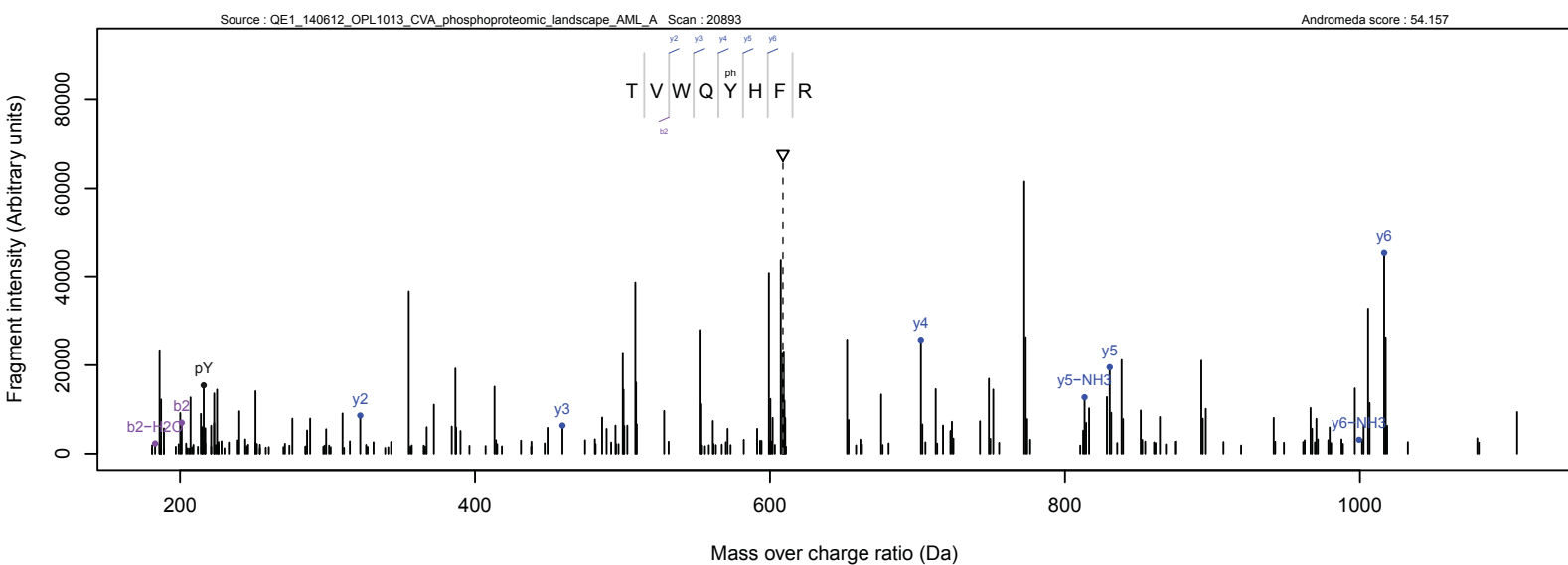
Andromeda score : 54.198



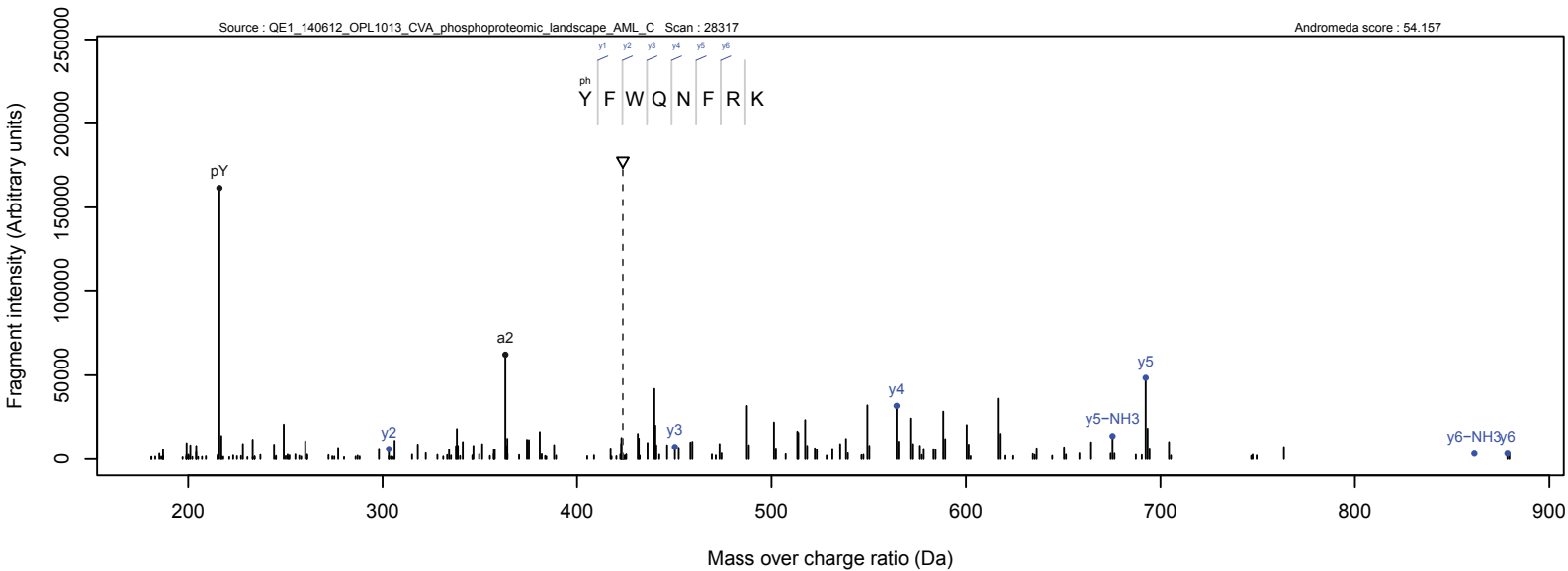
## HGQVY(ph)LRK-3+



## TVWQY(ph)HFR-2+

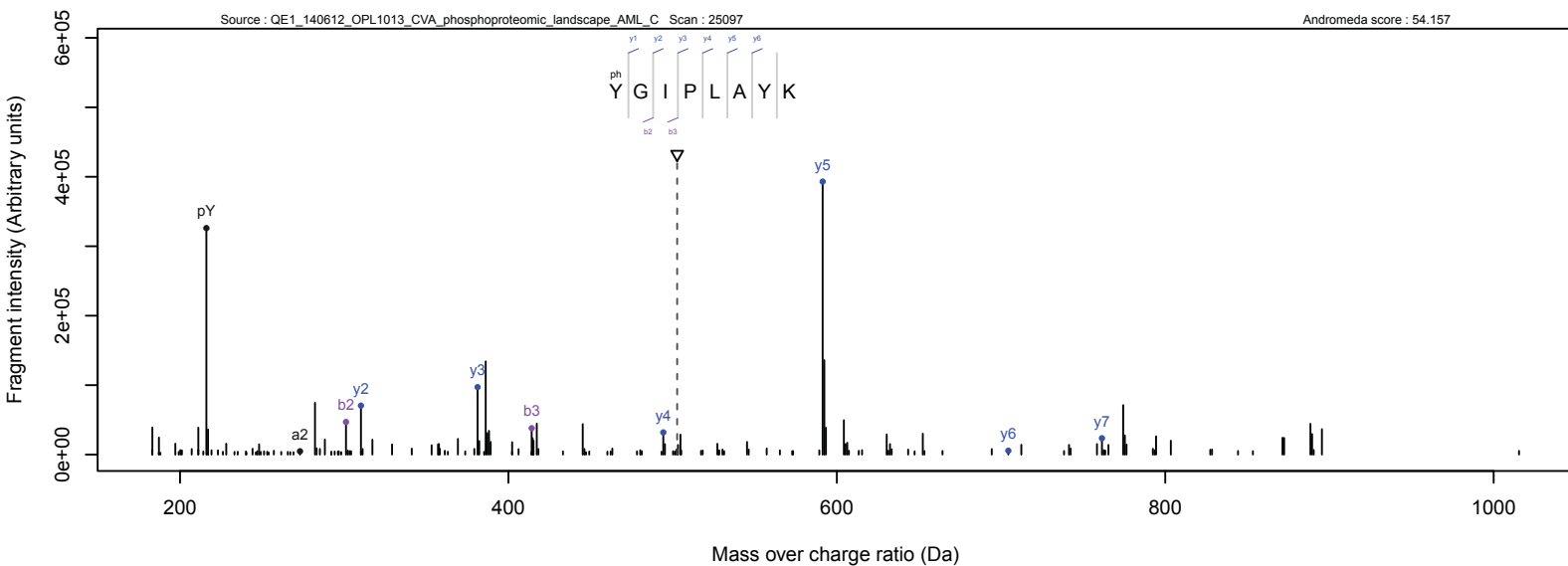


## Y(ph)FWQNF RK-3+

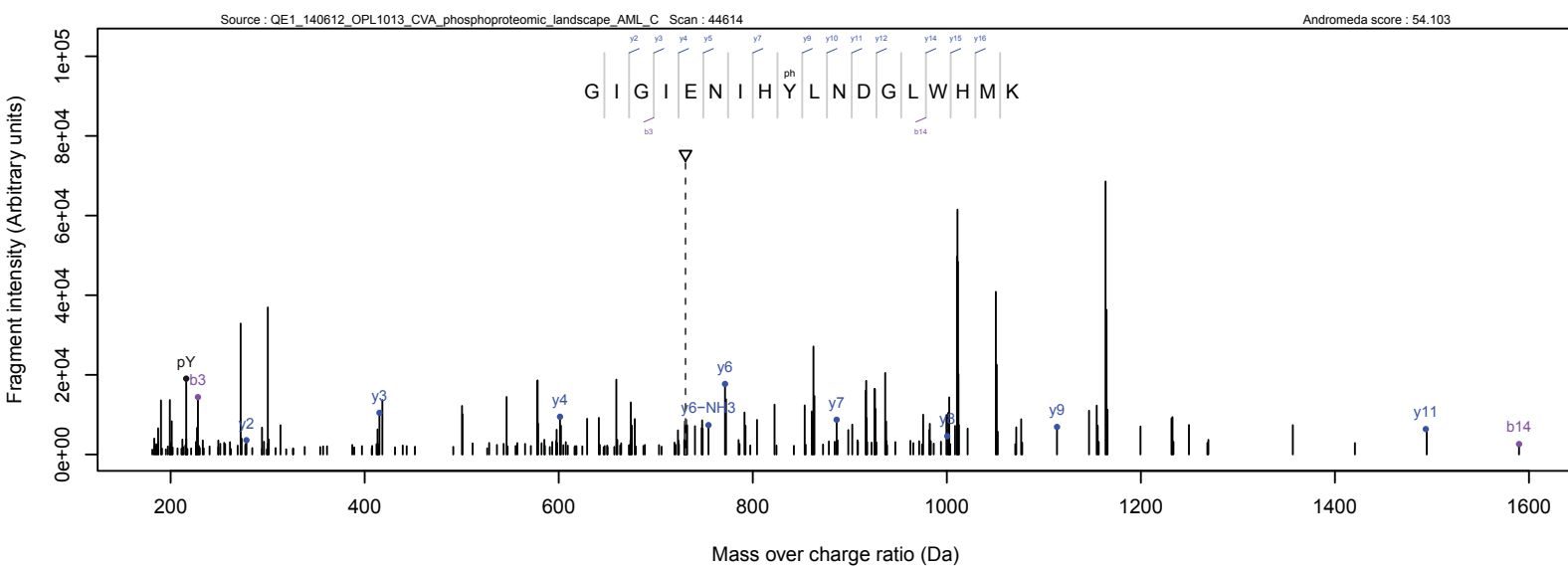




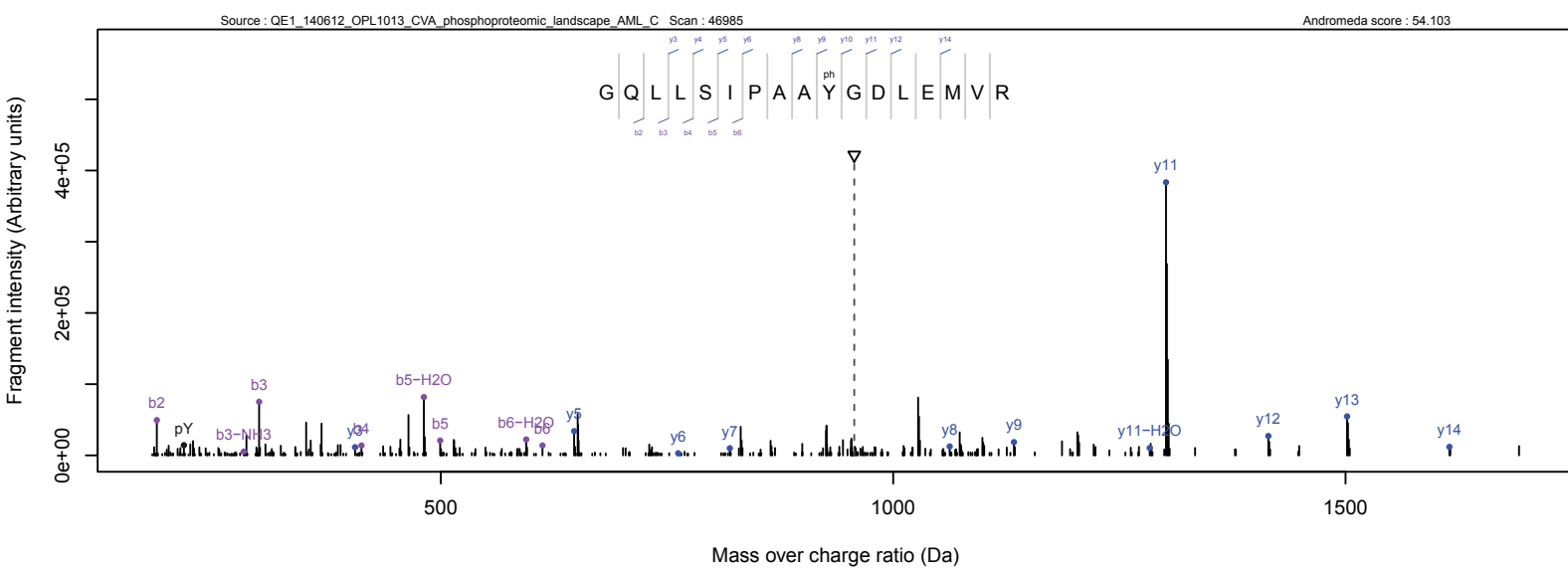
## Y(ph)GIPLAYK-2+



## GIGIENHY(ph)LNDGLWHMK-3+



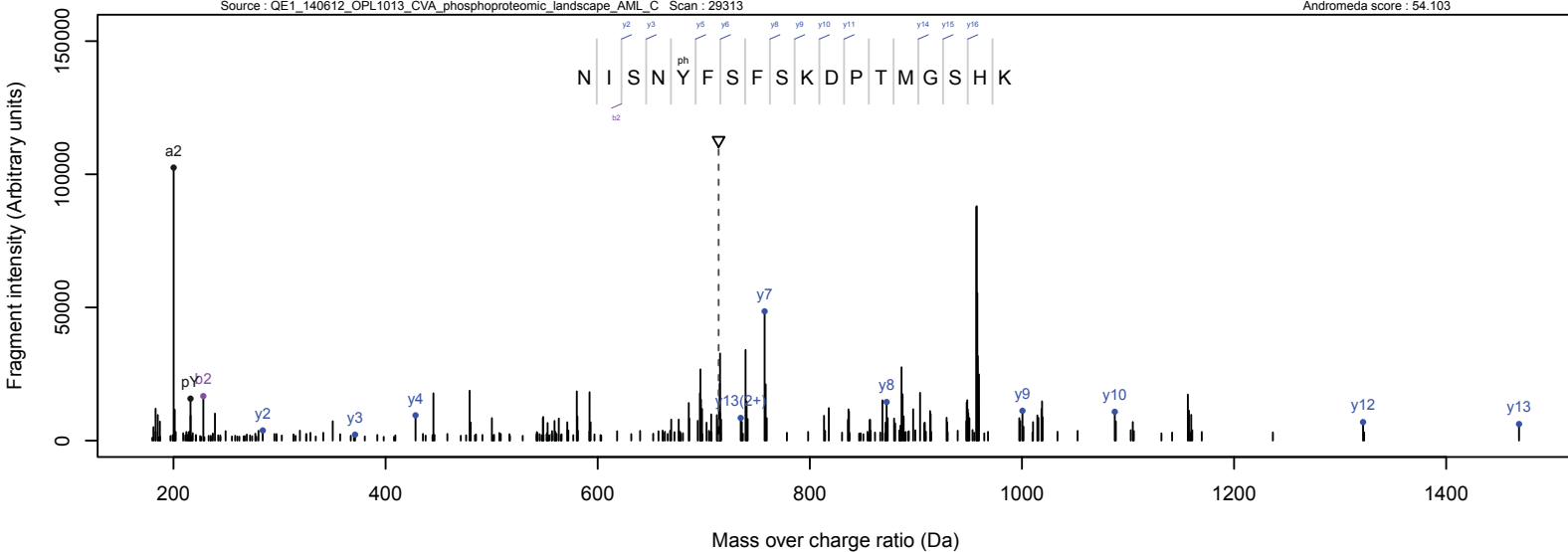
## GQLLSIPAAYGDLEMVR-2+



## NISNY(ph)FFSKDPTMGSHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29313

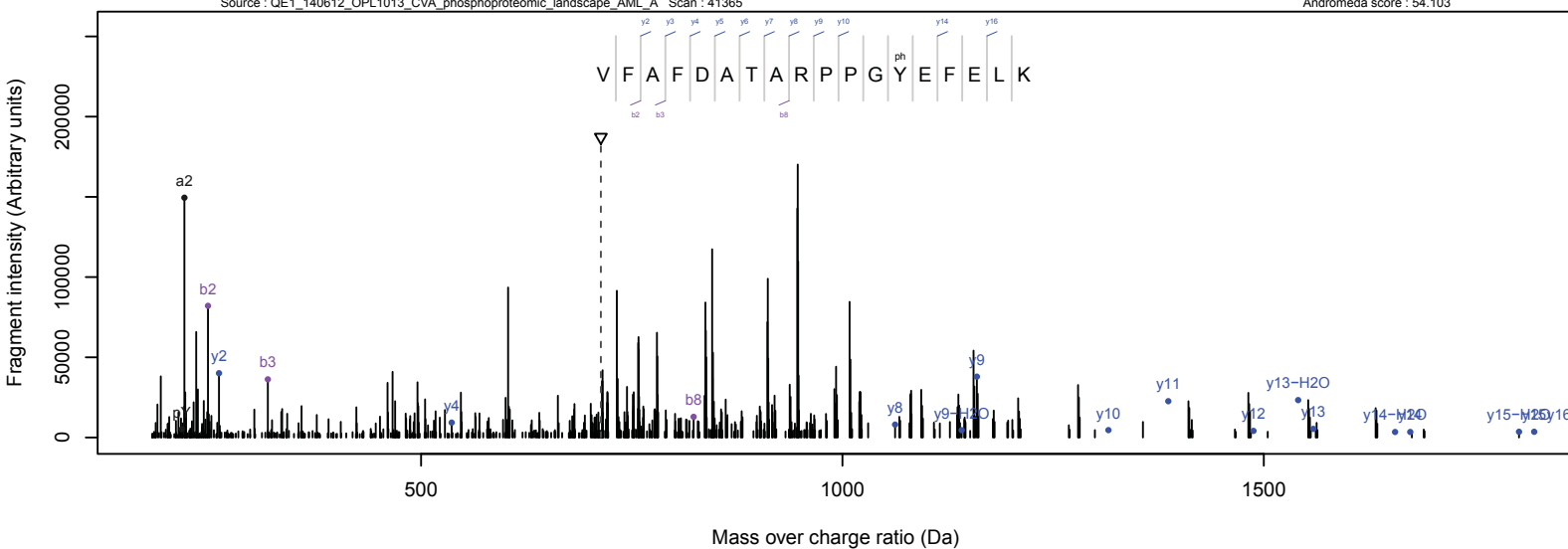
Andromeda score : 54.103



## VFADFATARPPGY(ph)EFELK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 41365

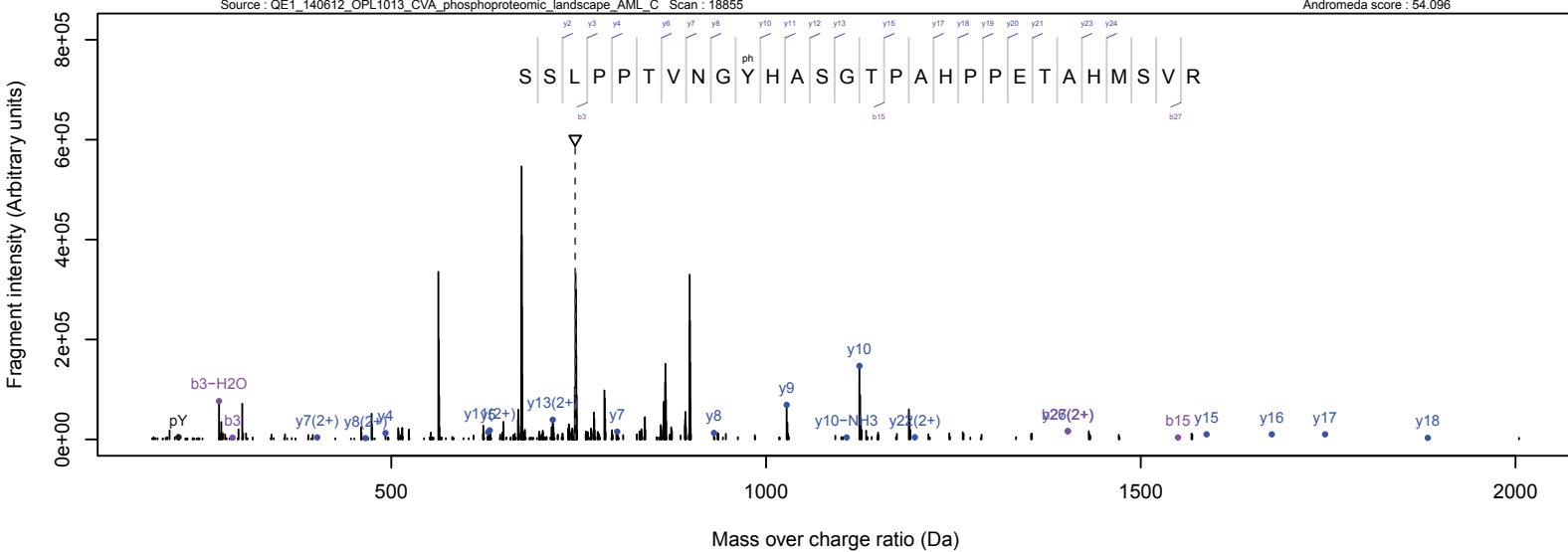
Andromeda score : 54.103



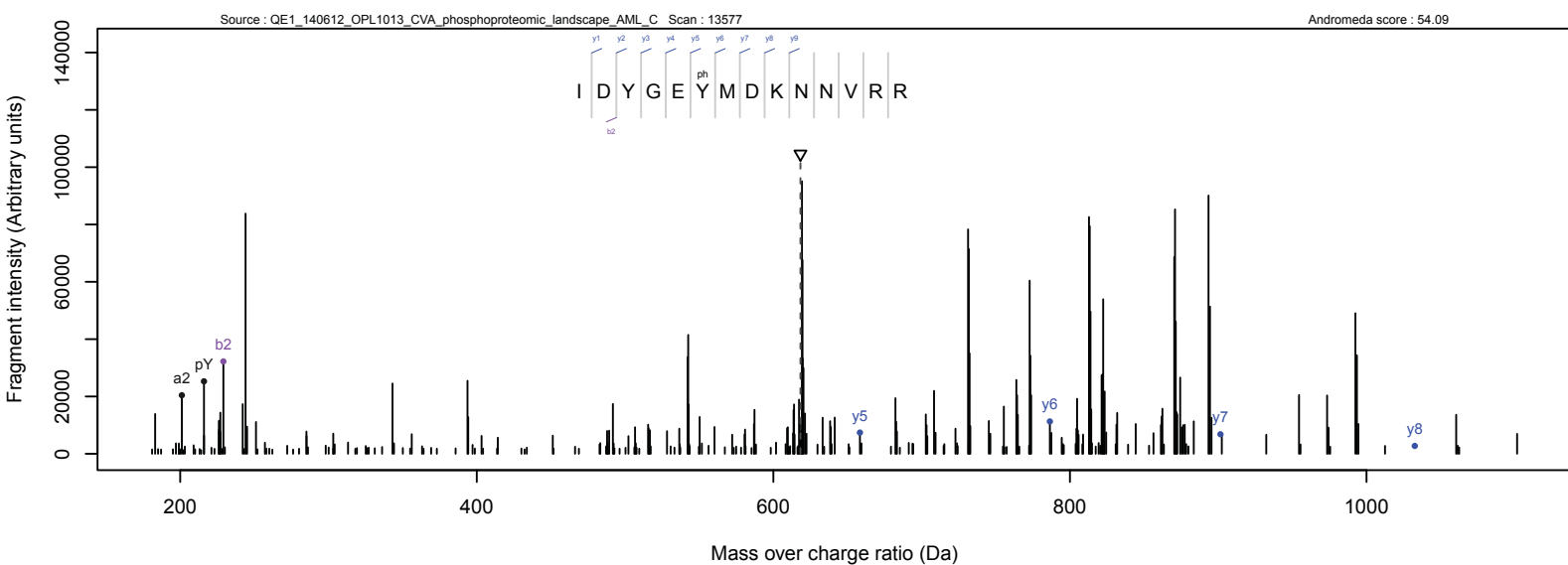
## SSLPPTVNGY(ph)HASGTPAHPPEAHMSVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18855

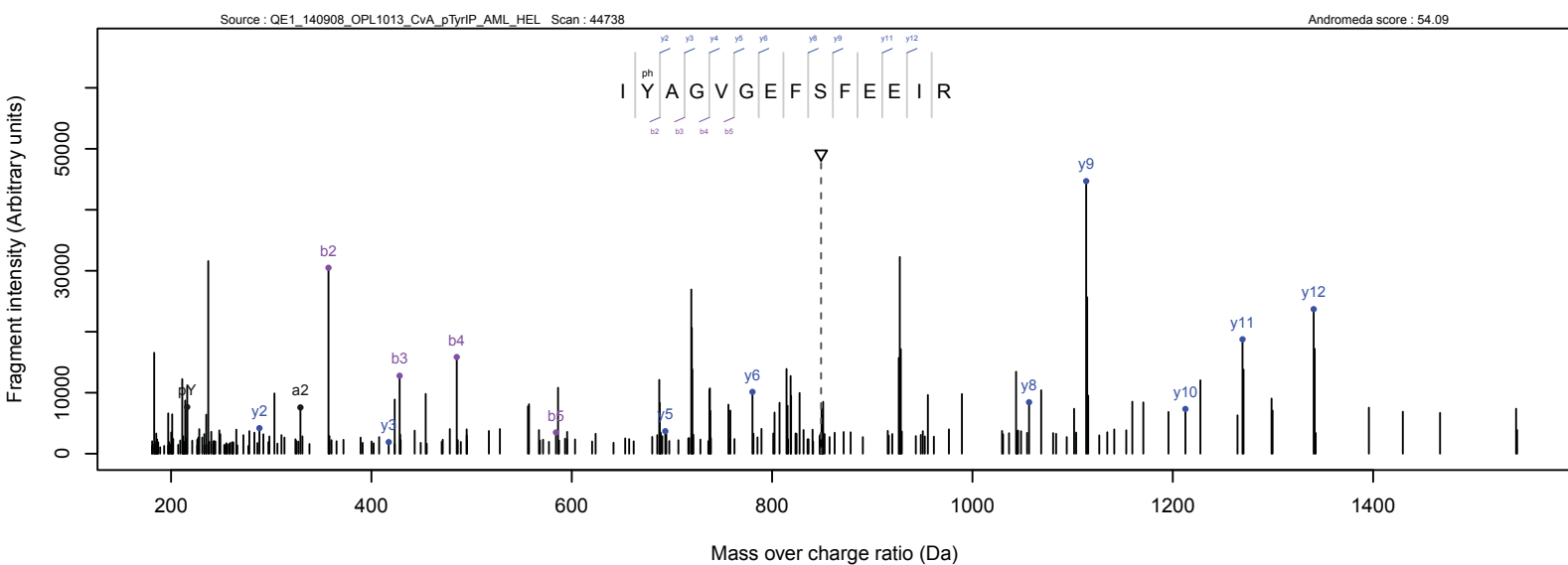
Andromeda score : 54.096



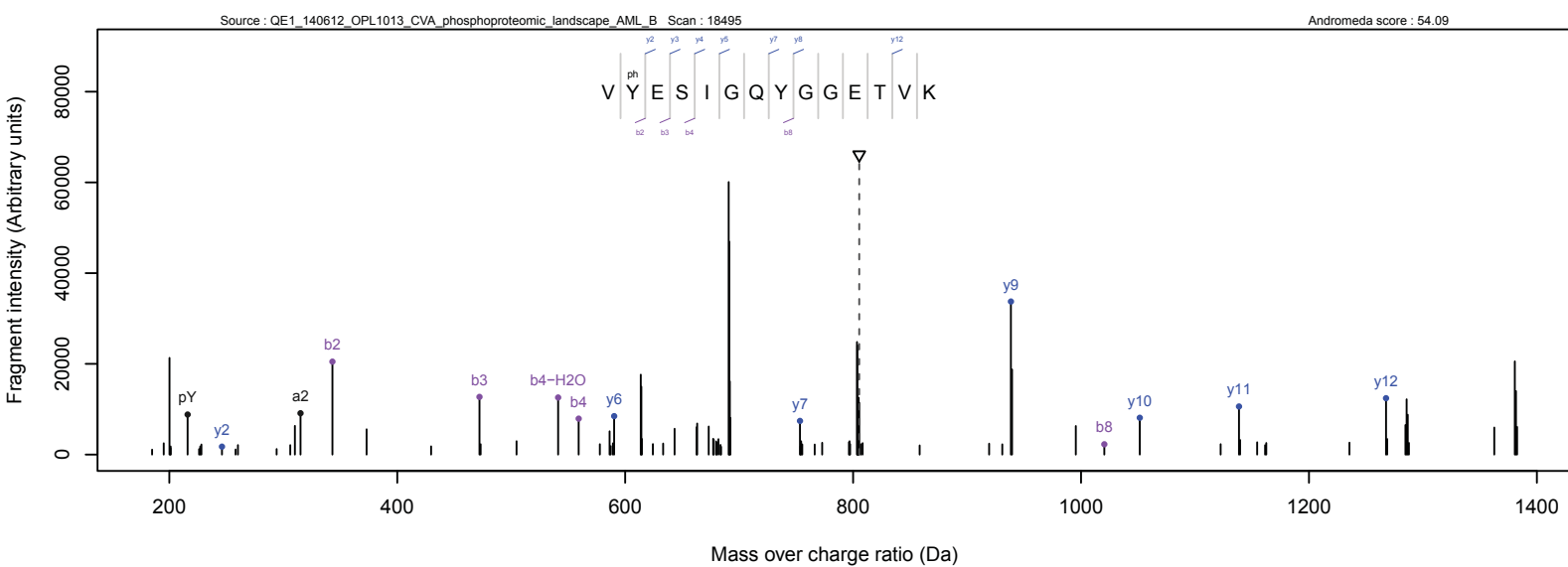
## IDYGEY(ph)MDKNNVRR-3+



## IY(ph)AGVGEFSFEEIR-2+



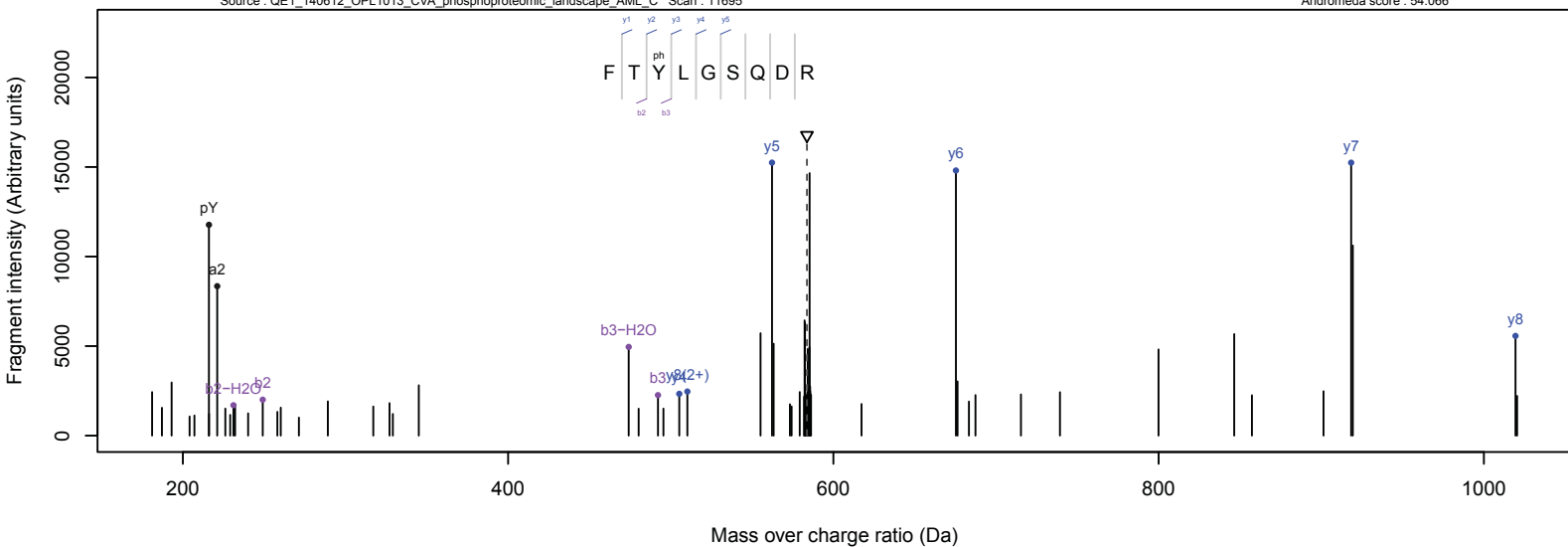
## VY(ph)ESIGQYGGETVK-2+



## FTY(ph)LGSQDR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11695

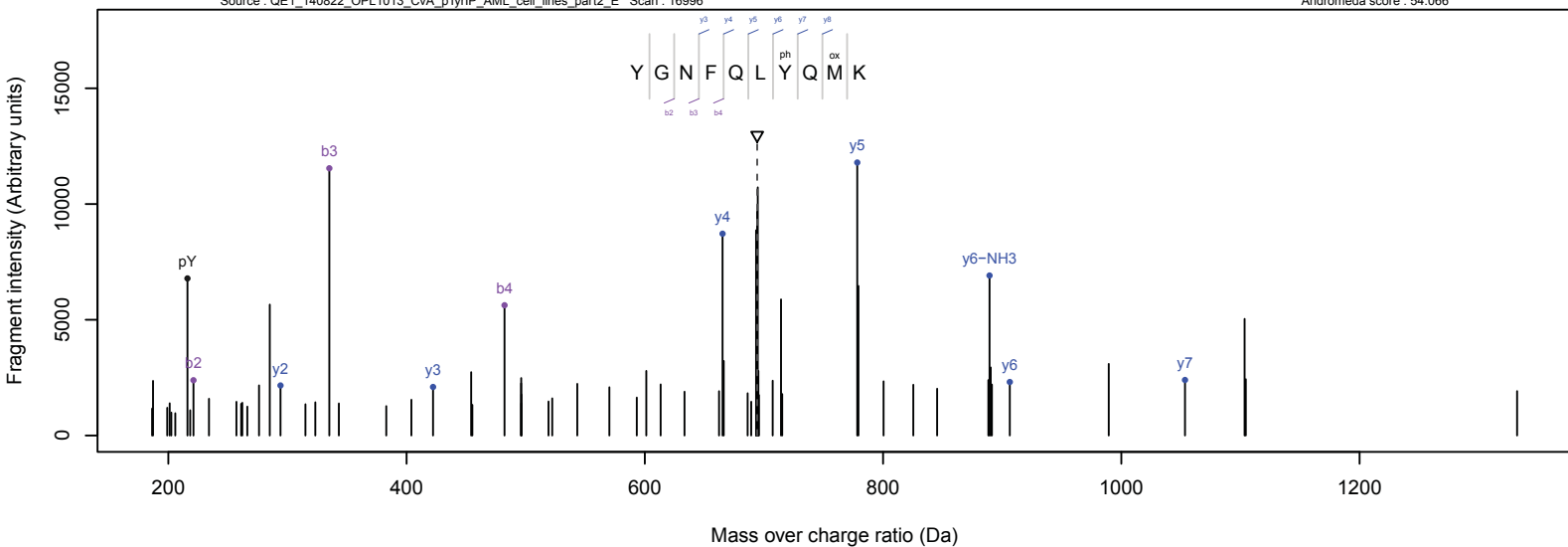
Andromeda score: 54.066



## YGNFQLYQM(ox)K-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 16996

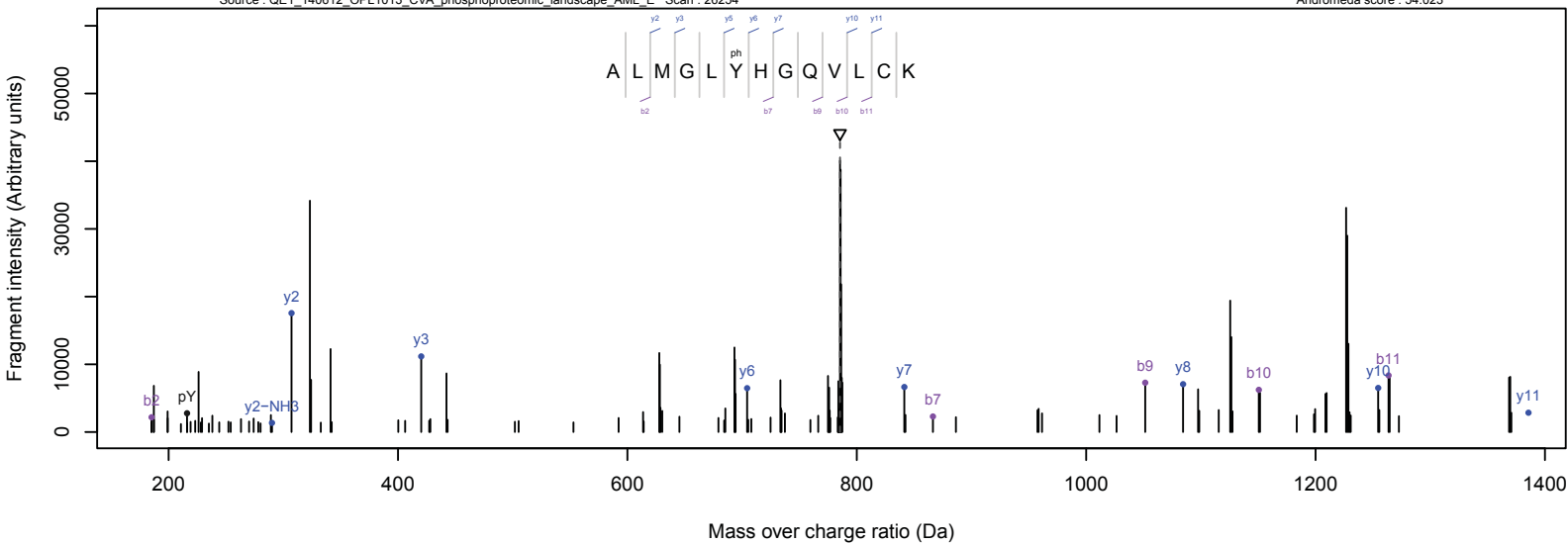
Andromeda score: 54.066



## ALMGLY(pH)HGQVLCK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 26254

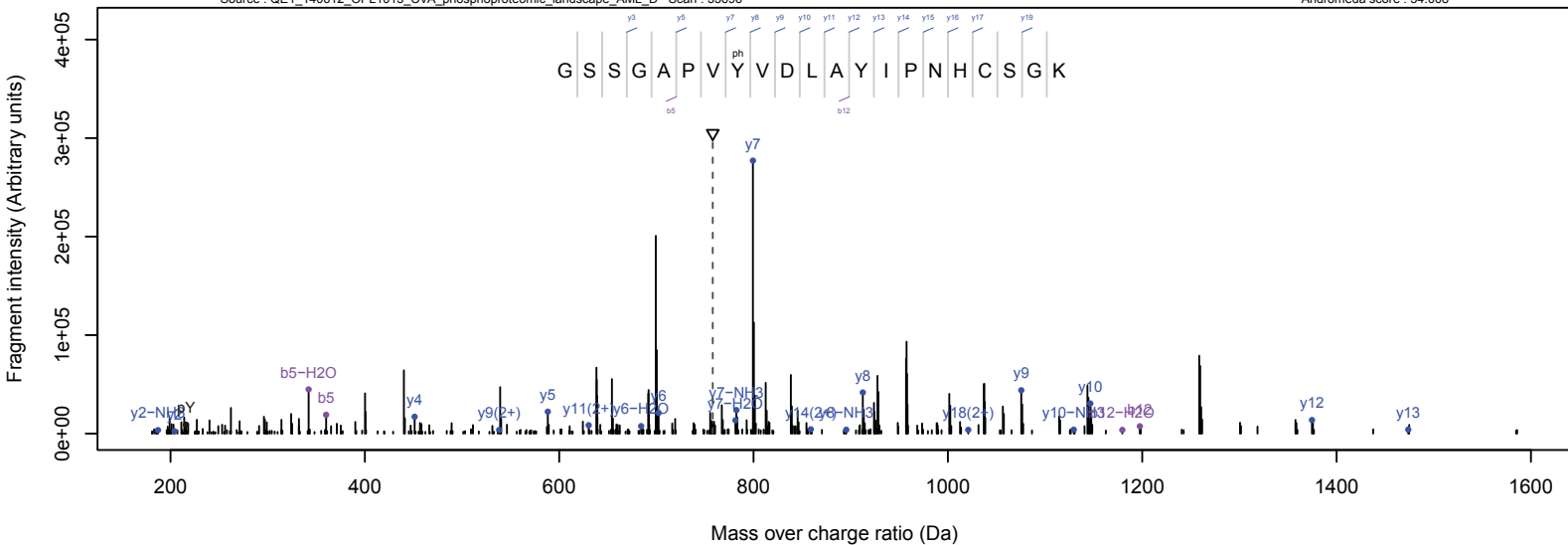
Andromeda score: 54.023



## GSSGAPVY(ph)VDLAYIPNHCSGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33696

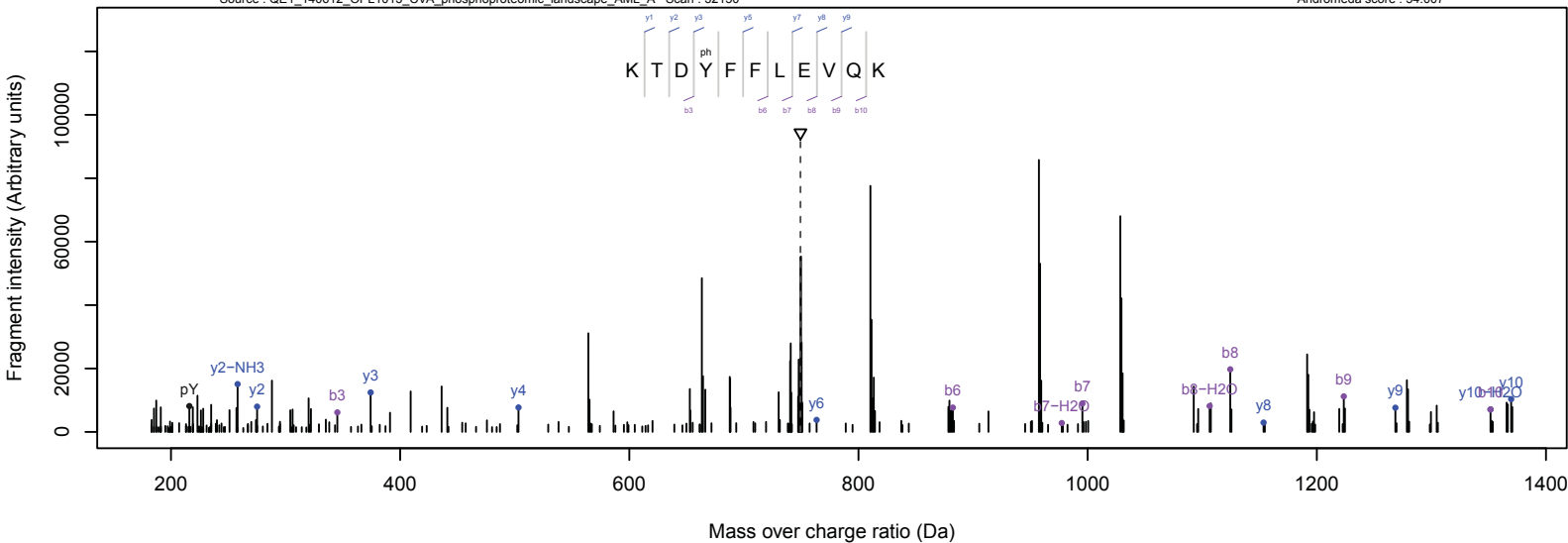
Andromeda score : 54.008



## KTDY(ph)FFLEVQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32150

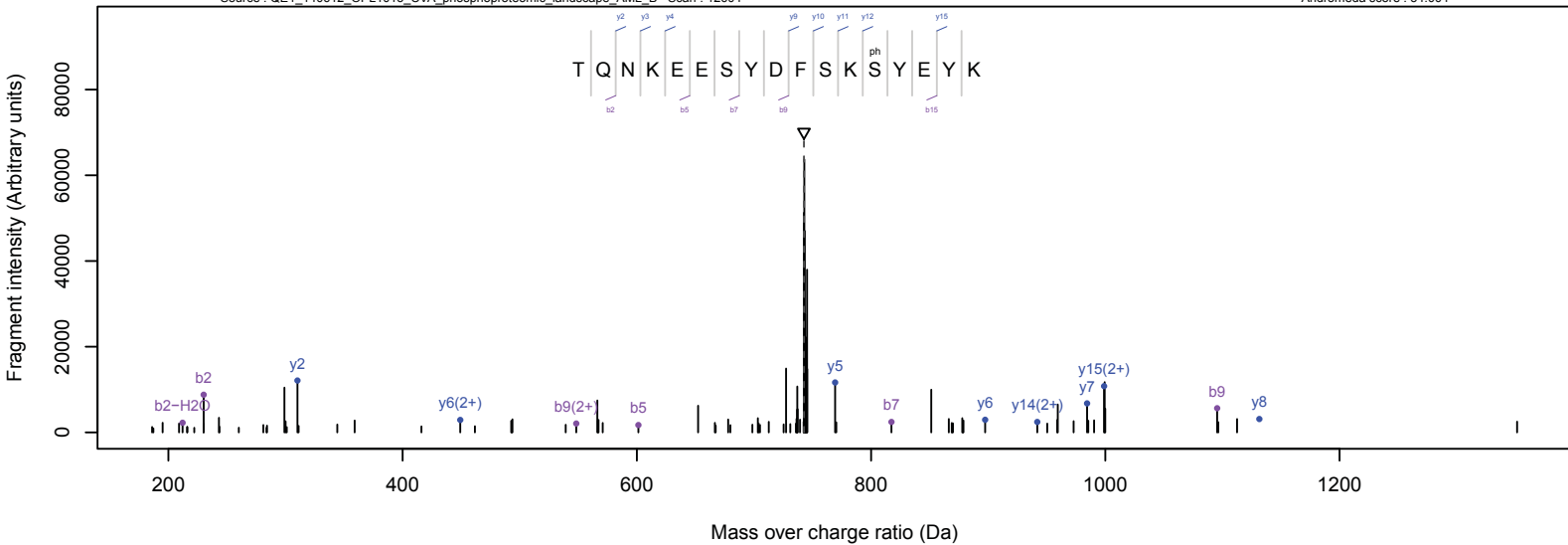
Andromeda score : 54.007



## TQNKEESYDFS(ph)YEEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12604

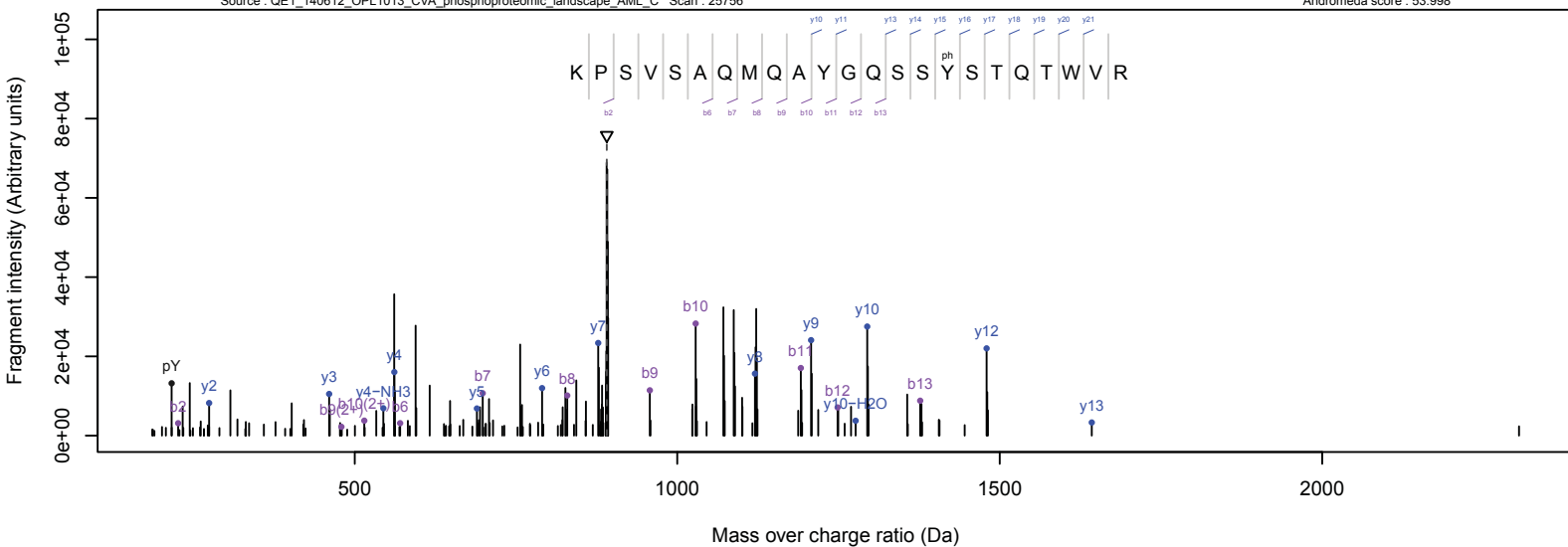
Andromeda score : 54.004



## KPSVSAQMAYGQSS(ph)STQWVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25756

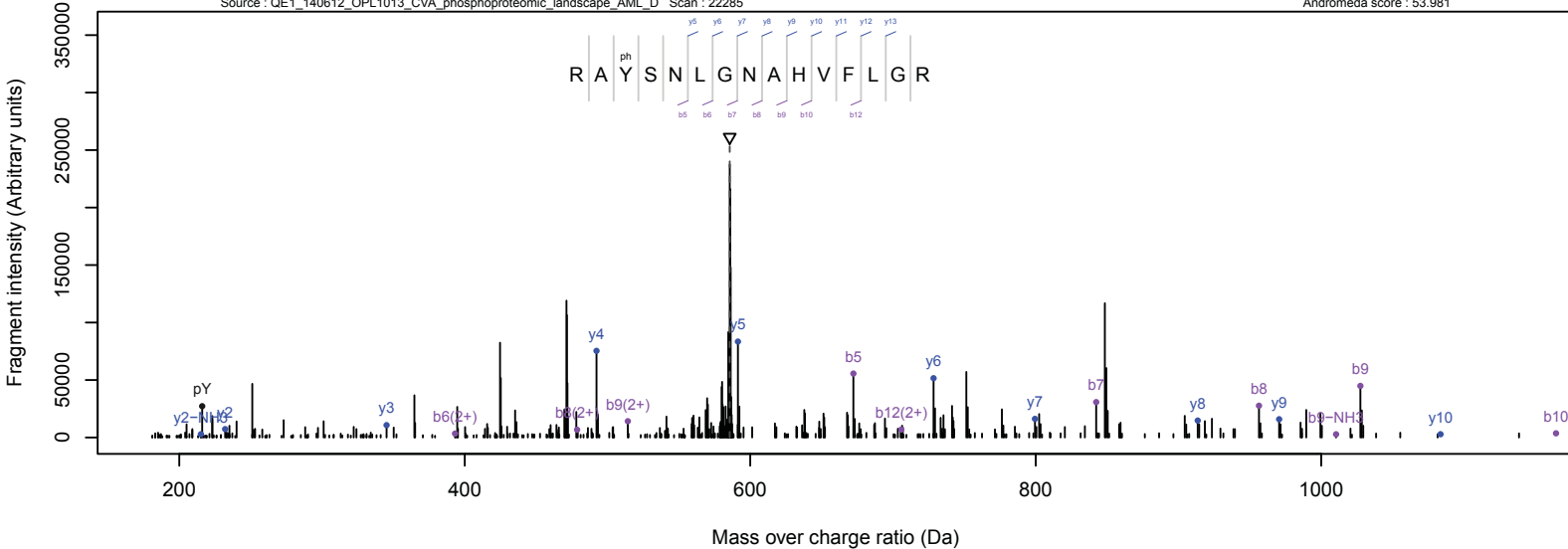
Andromeda score : 53.998



## RAY(ph)SNLGNHVFVFLGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22285

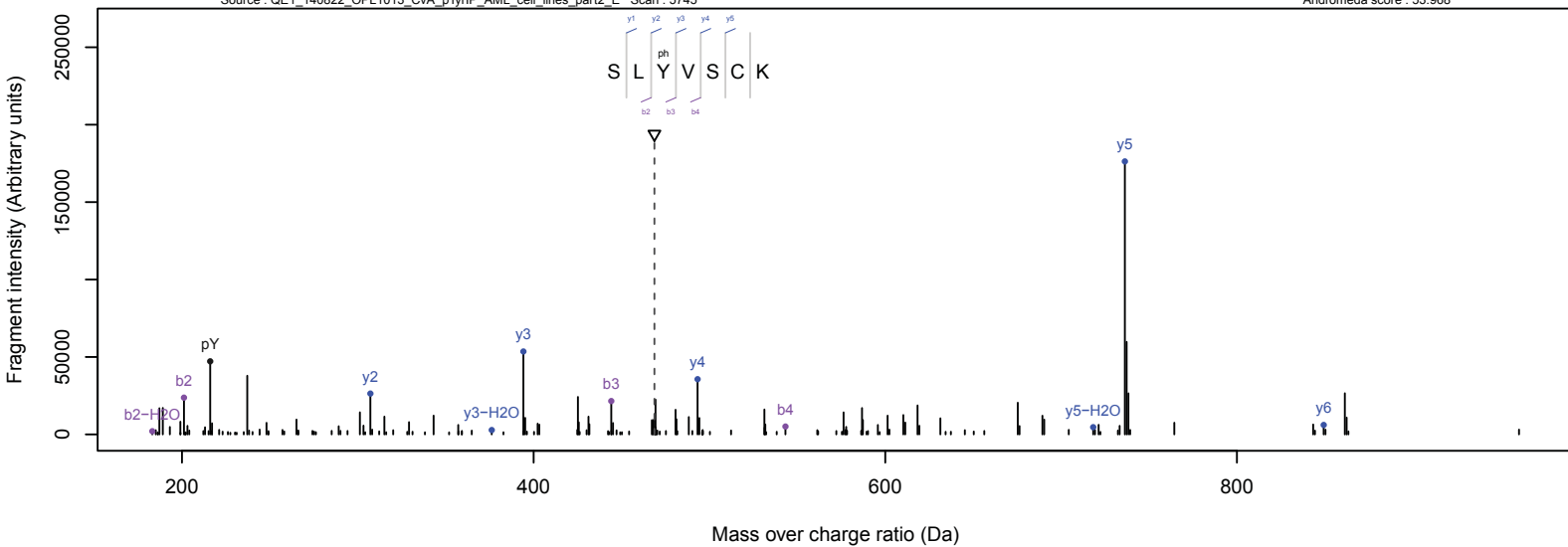
Andromeda score : 53.981



## SLY(ph)VSCK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 5745

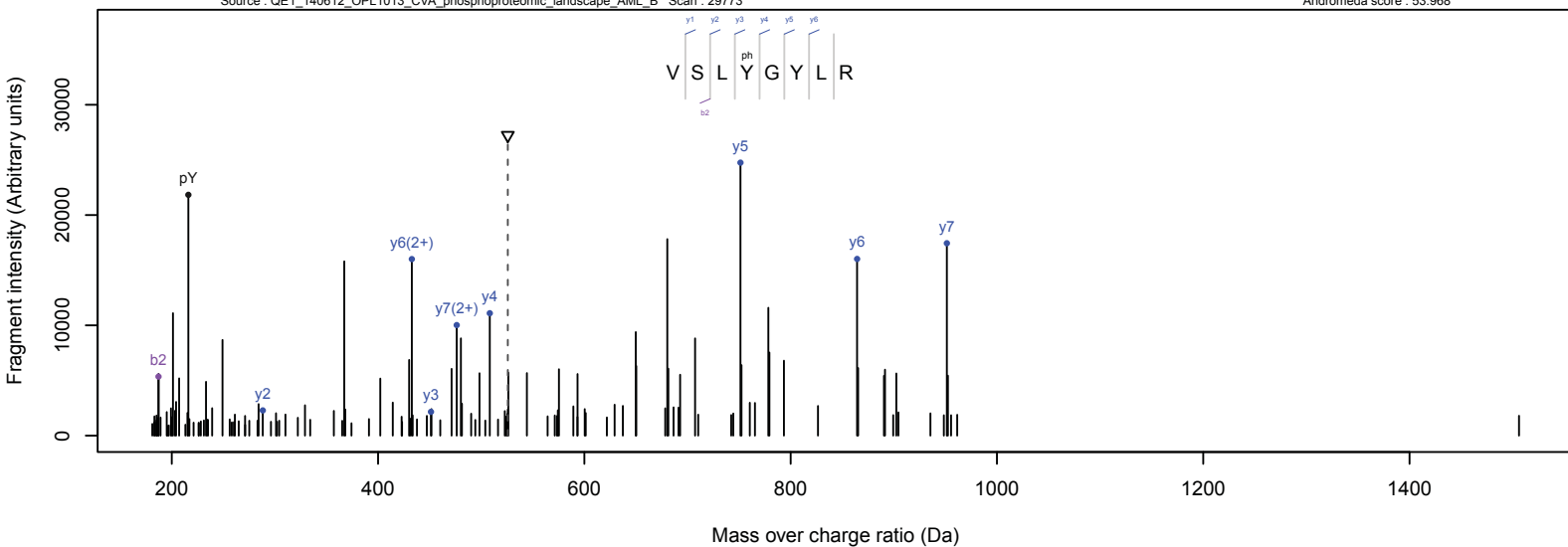
Andromeda score : 53.968



## VSLY(ph)GYLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29773

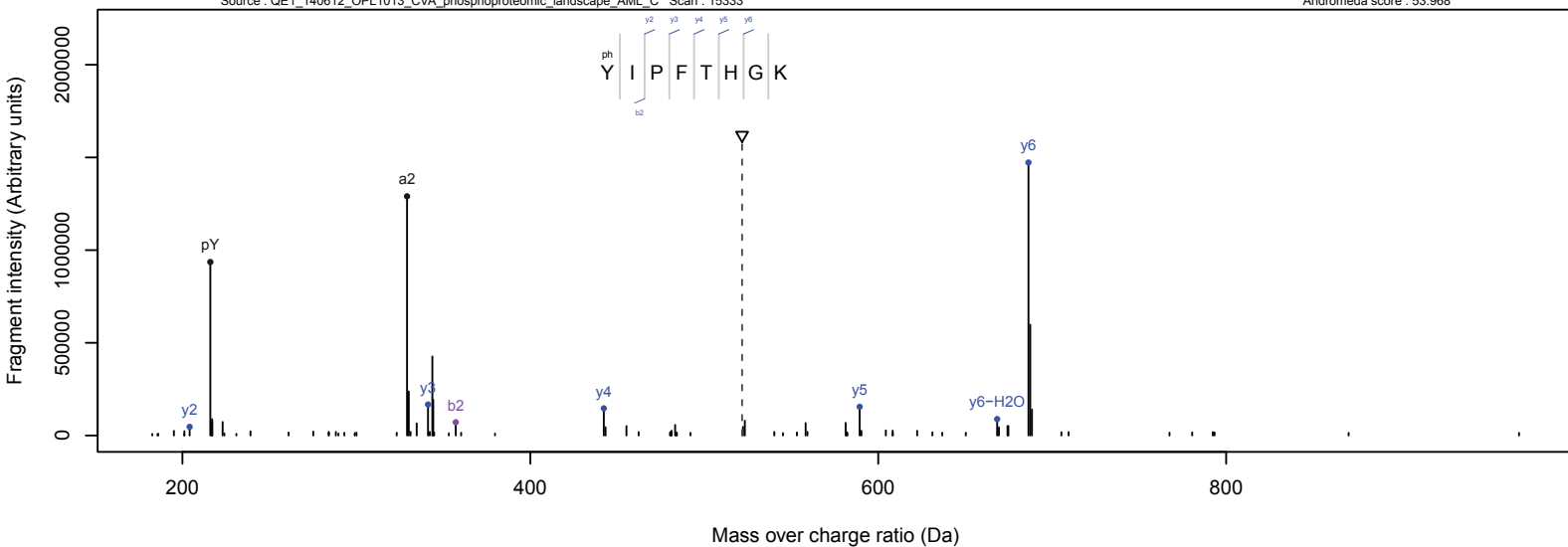
Andromeda score : 53.968



## Y(ph)IPFTHGK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15333

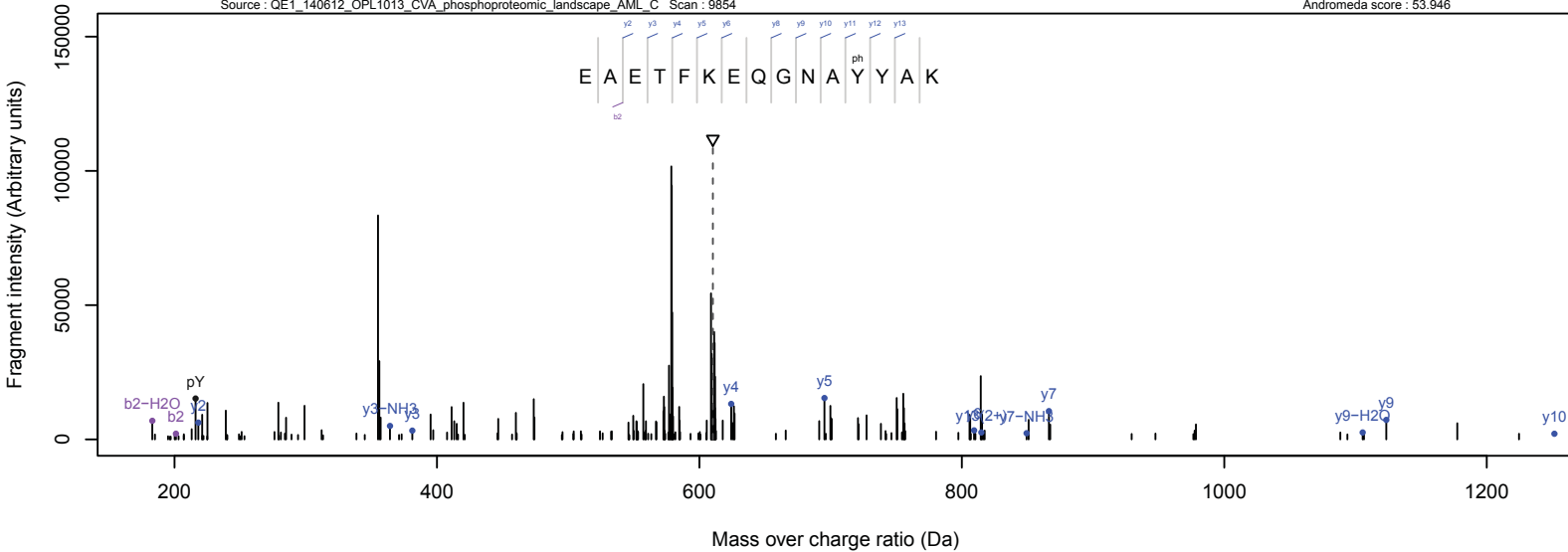
Andromeda score : 53.968



## EAETFKEQGNAY(ph)YAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9854

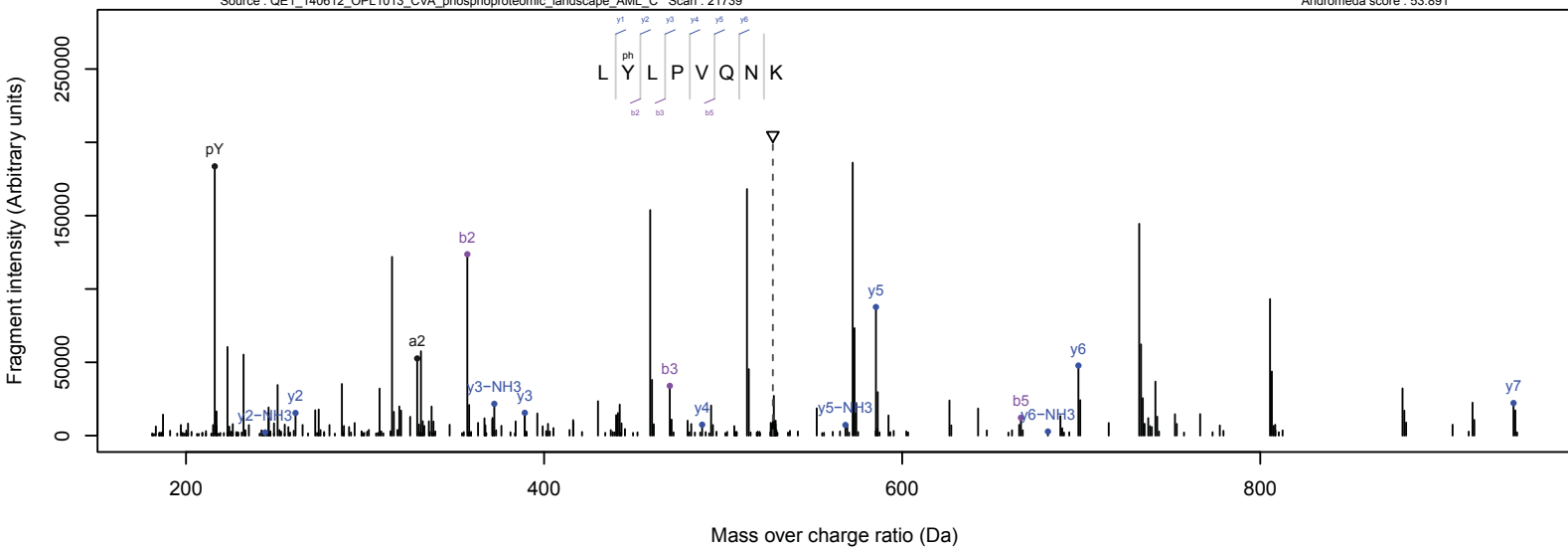
Andromeda score : 53.946



## LY(ph)LPVQNK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21739

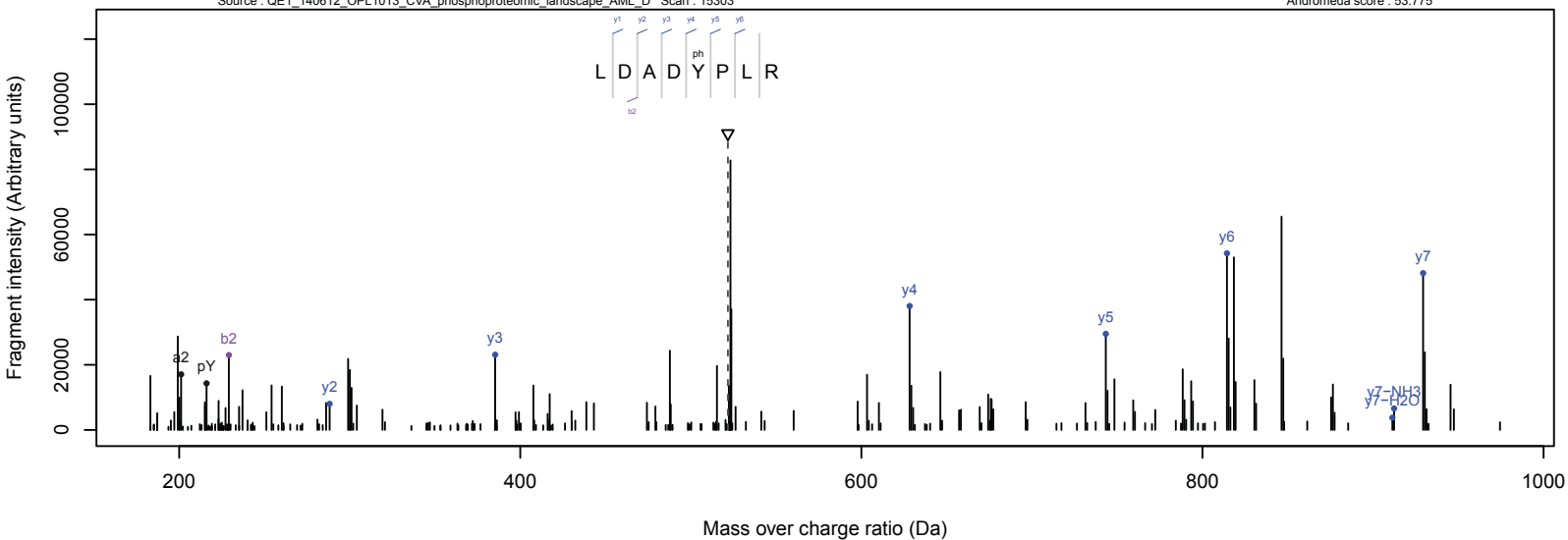
Andromeda score: 53.891



## LDADY(ph)PLR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 15303

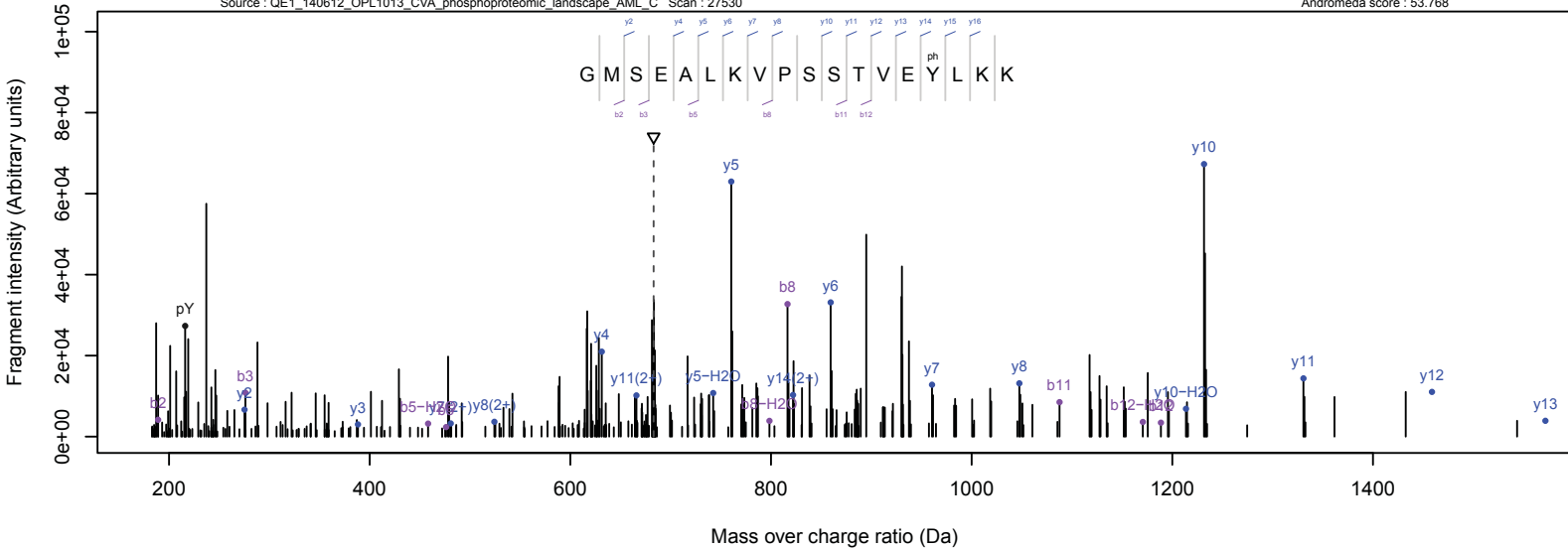
Andromeda score: 53.775



## GMSEALKVPSSTVEY(ph)LKK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 27530

Andromeda score: 53.768

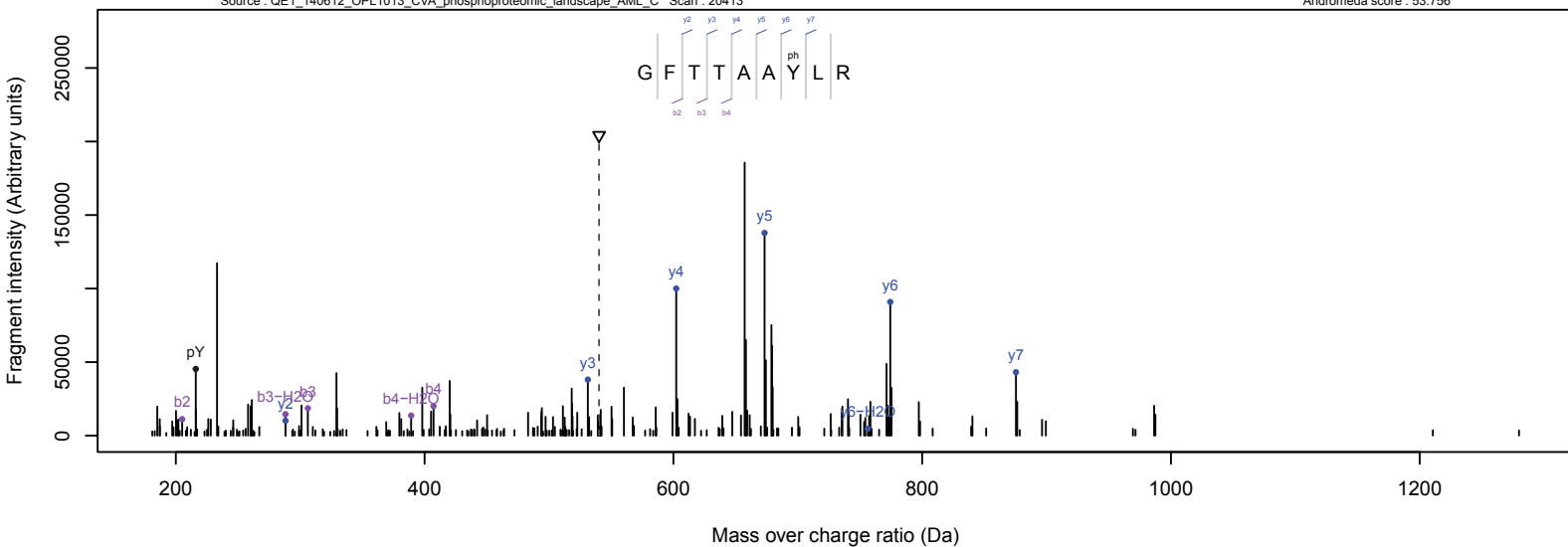




## GFTTAY(ph)LR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20413

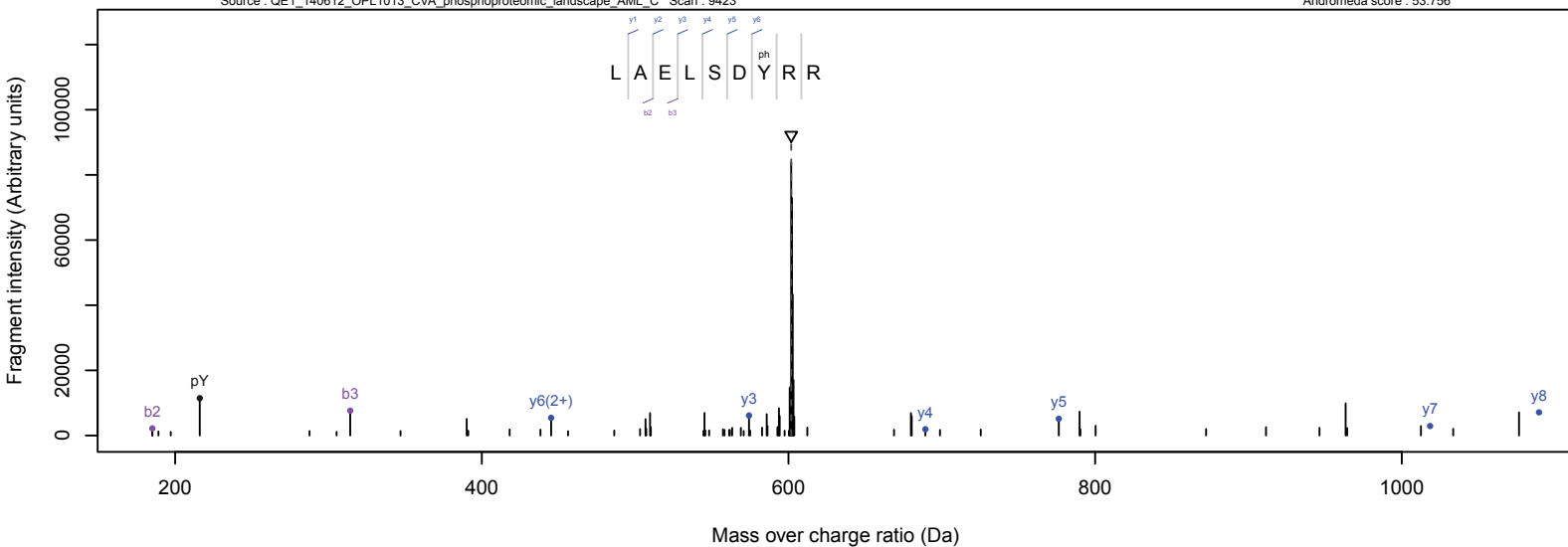
Andromeda score : 53.756



## LAELSDY(ph)RR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9423

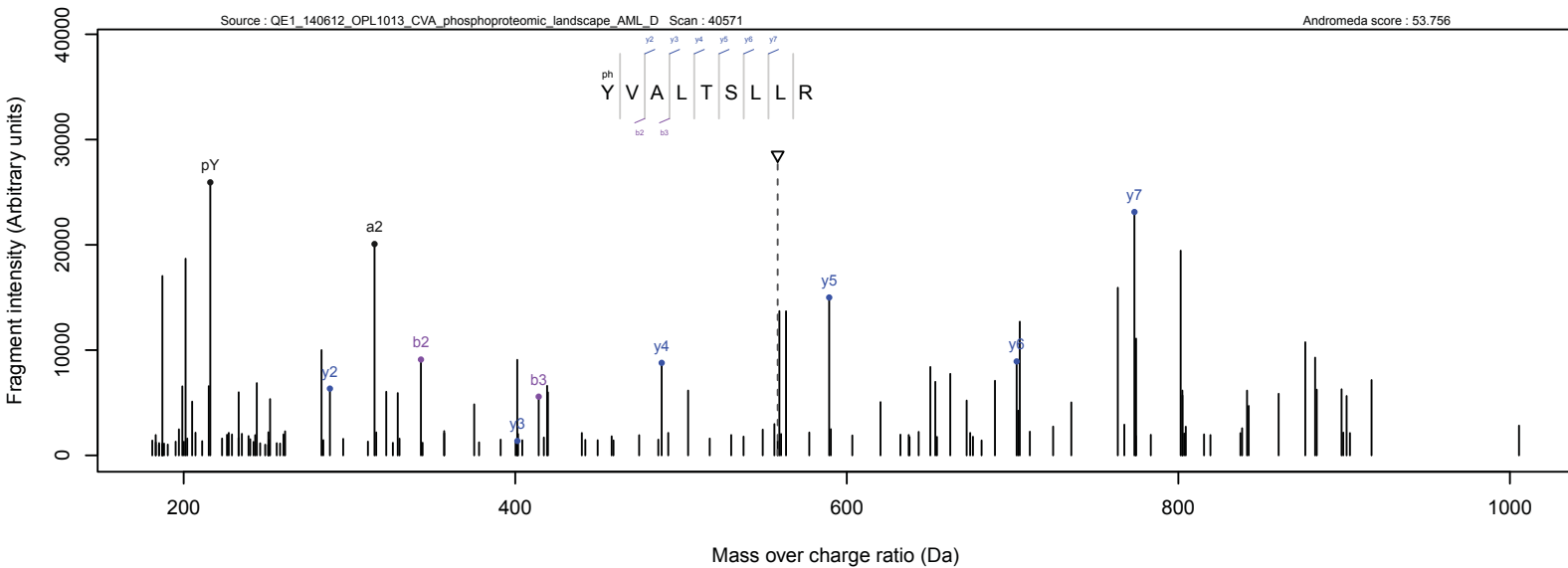
Andromeda score : 53.756



## Y(ph)VALTSLLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 40571

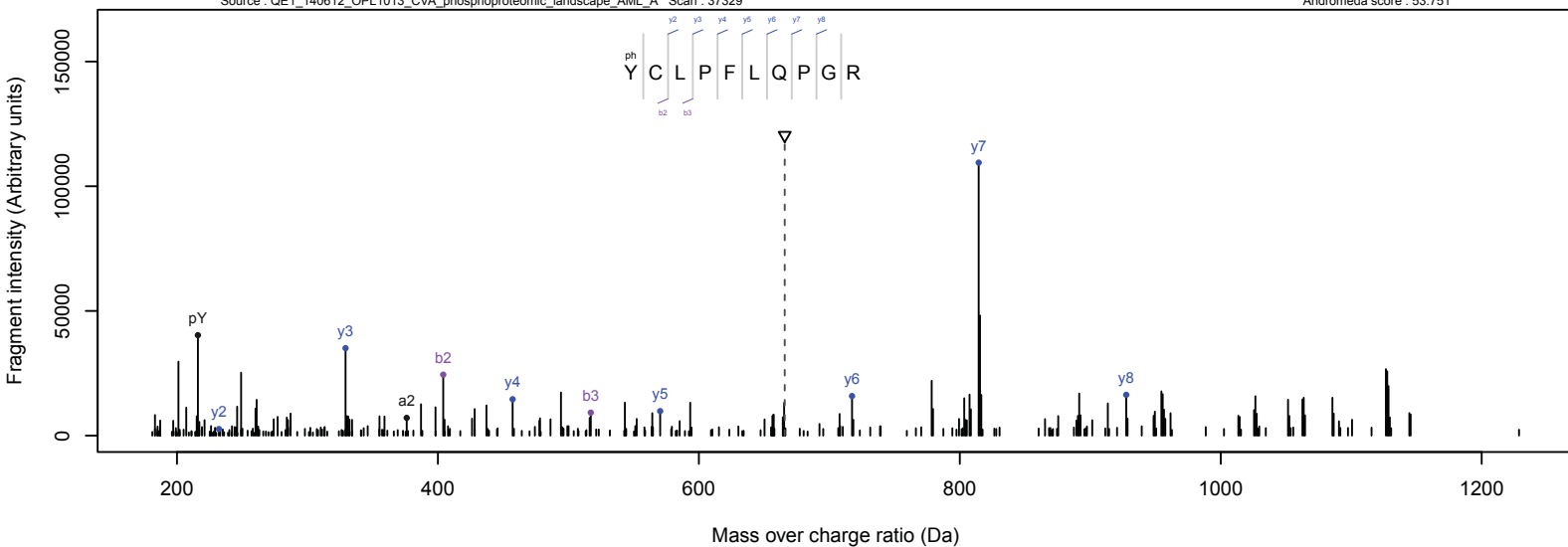
Andromeda score : 53.756



## Y(ph)CLPFLQPGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 37329

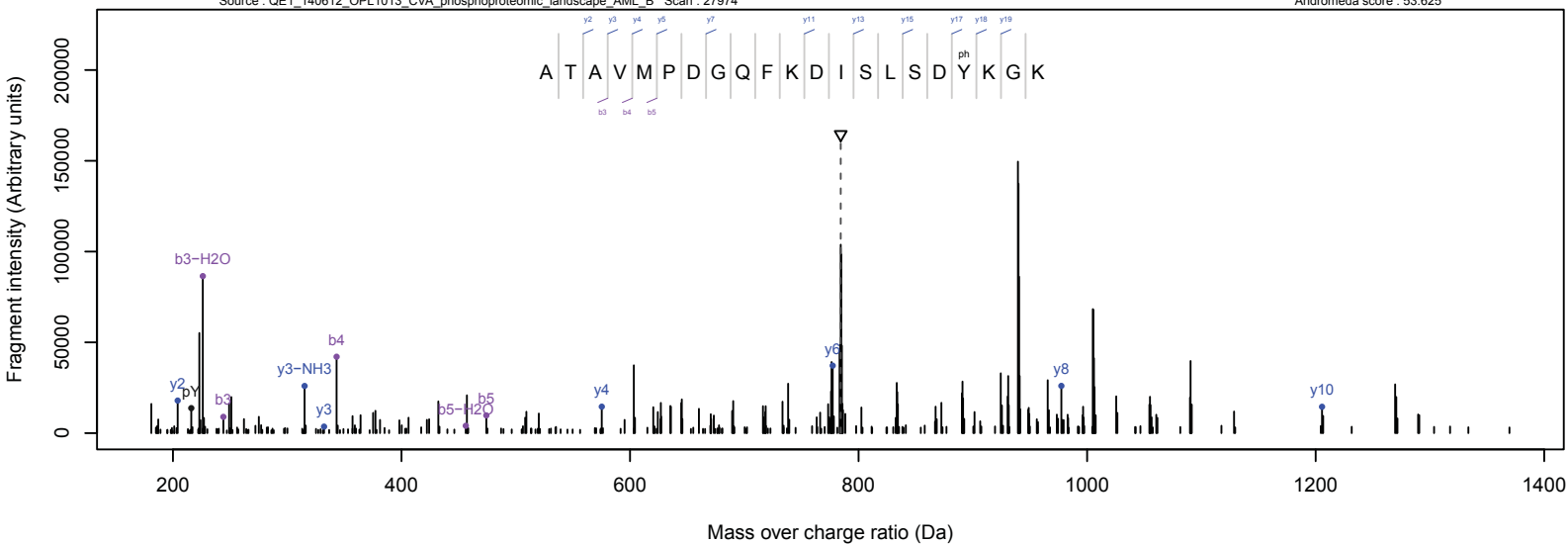
Andromeda score : 53.751



## ATAVMPDGGFKDISLSDY(ph)KGGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27974

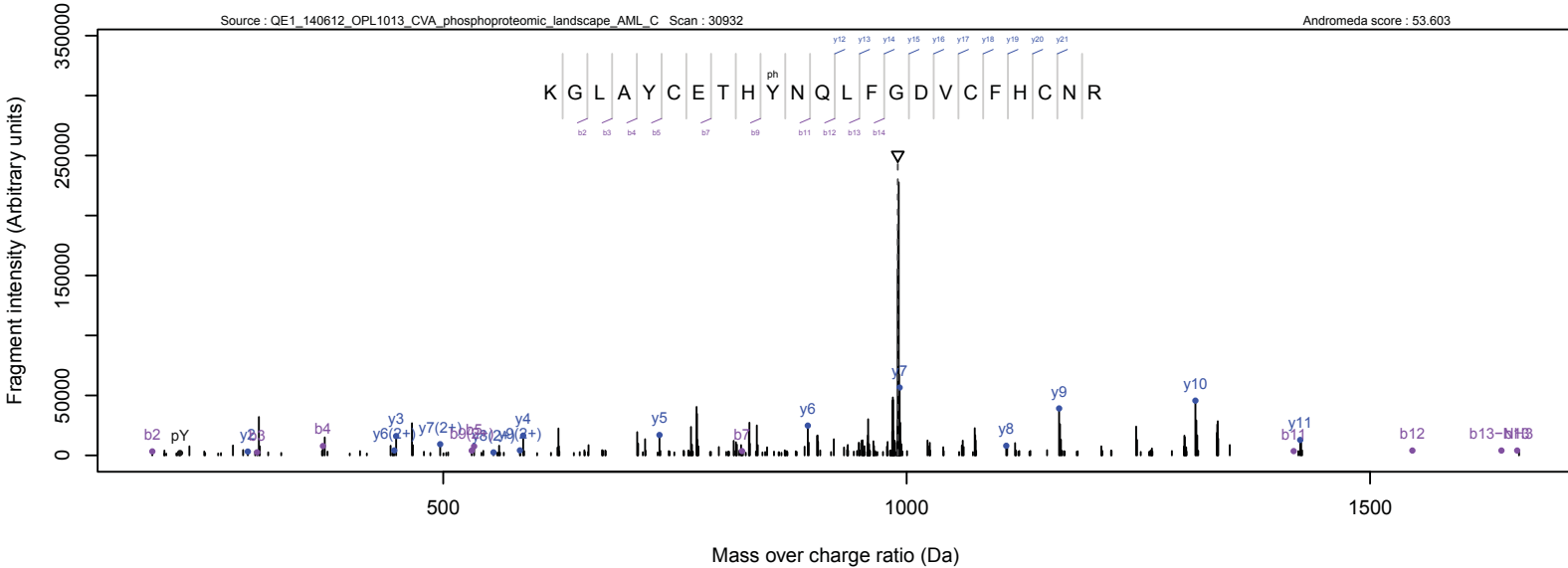
Andromeda score : 53.625



## KGLAYCETHY(ph)NQLFGDVCFHCNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30932

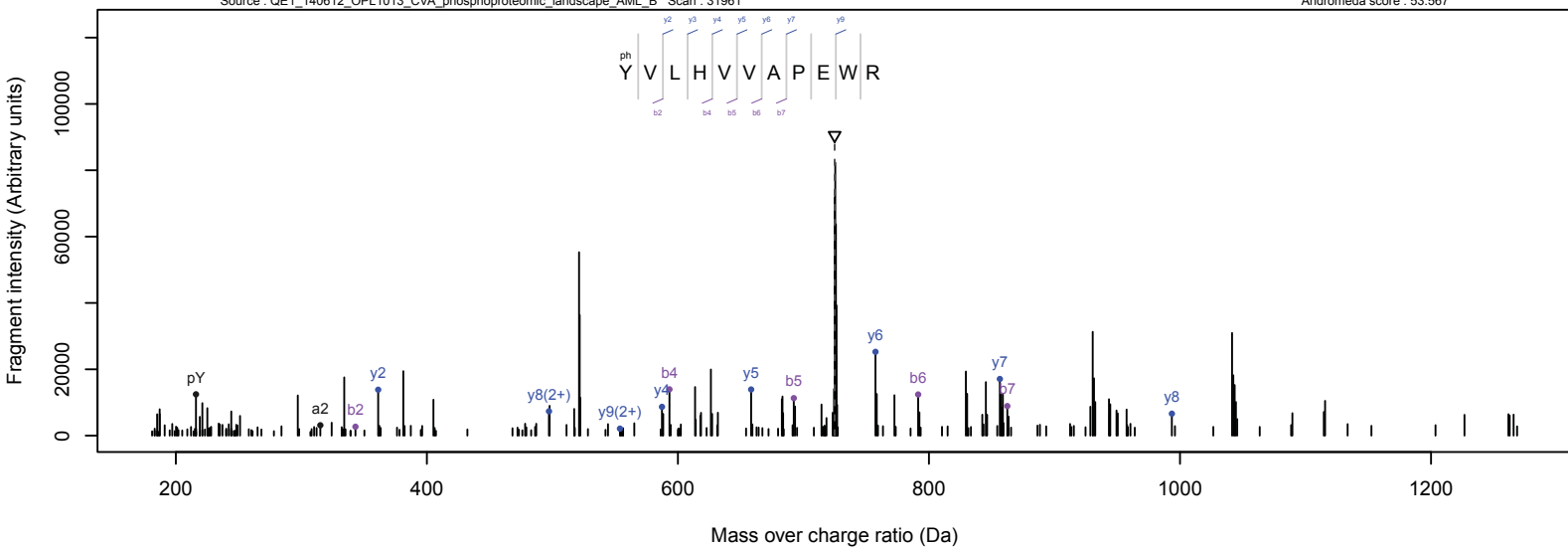
Andromeda score : 53.603



## Y(ph)VLHVVAPEWR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31961

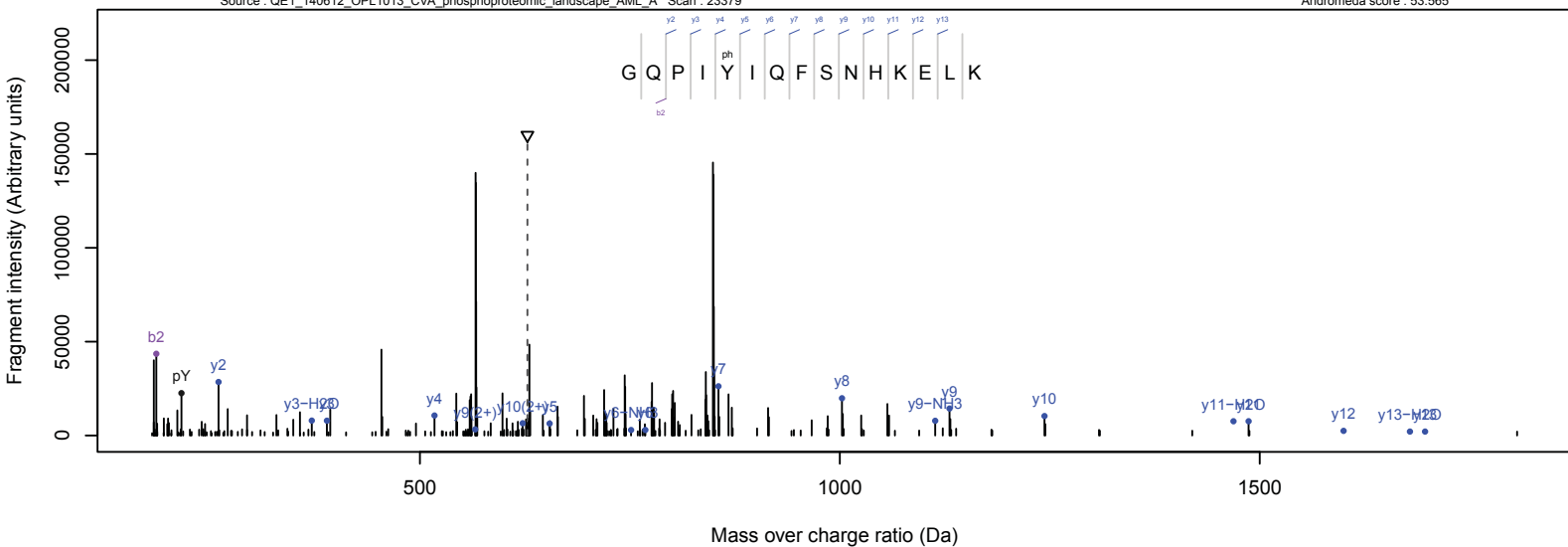
Andromeda score : 53.567



## GQPIY(ph)IQFSNHKELK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 23379

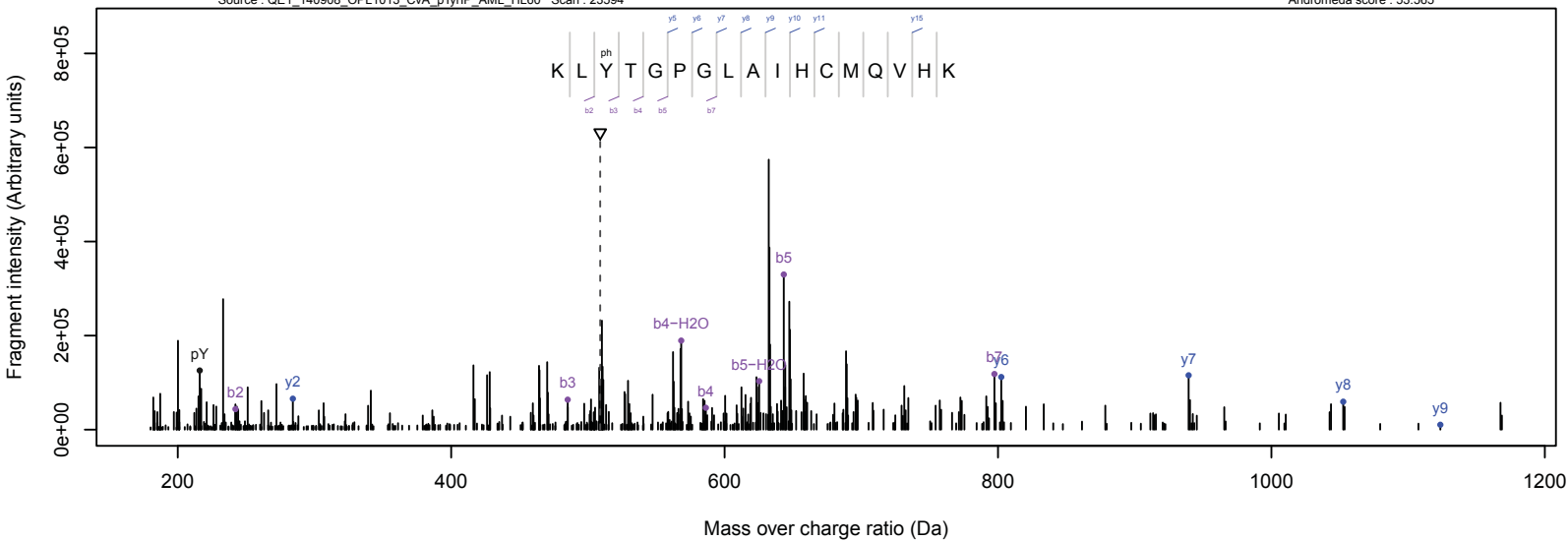
Andromeda score : 53.565



## KLY(ph)TGPLAIHCMQVHK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 23594

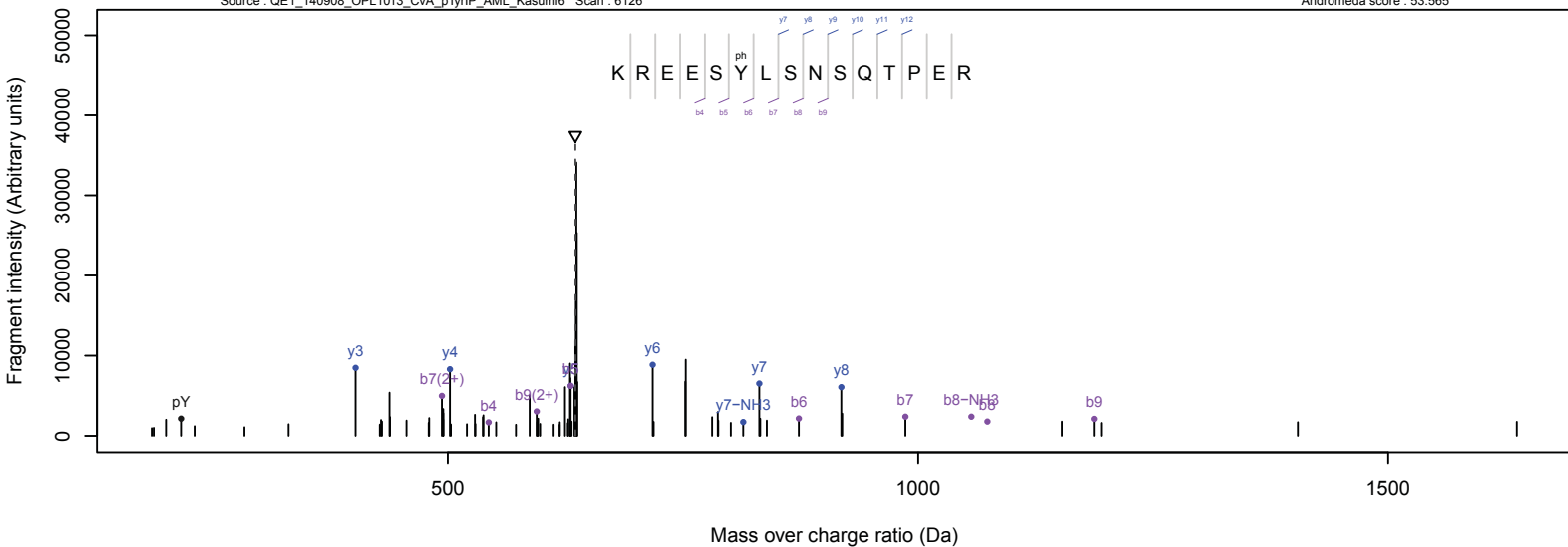
Andromeda score : 53.565



**KREESY(ph)LSNSQTPER-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 6126

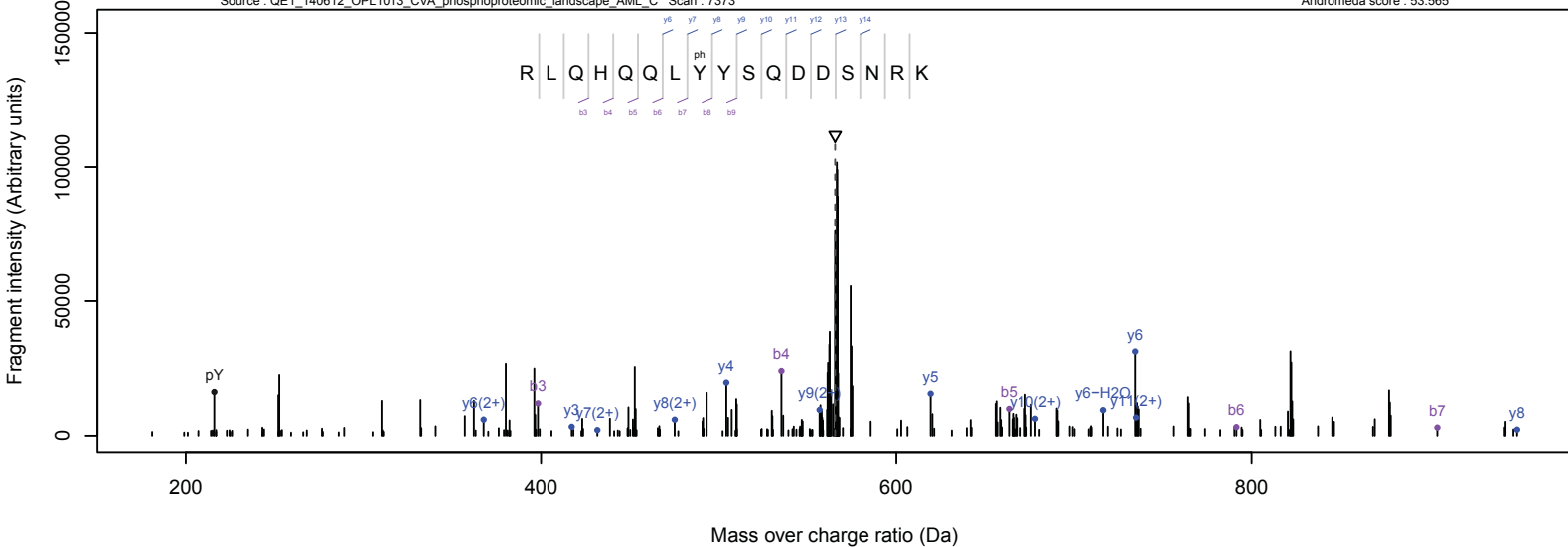
Andromeda score : 53.565



**RLQHQLY(ph)YSQDDSNRK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7373

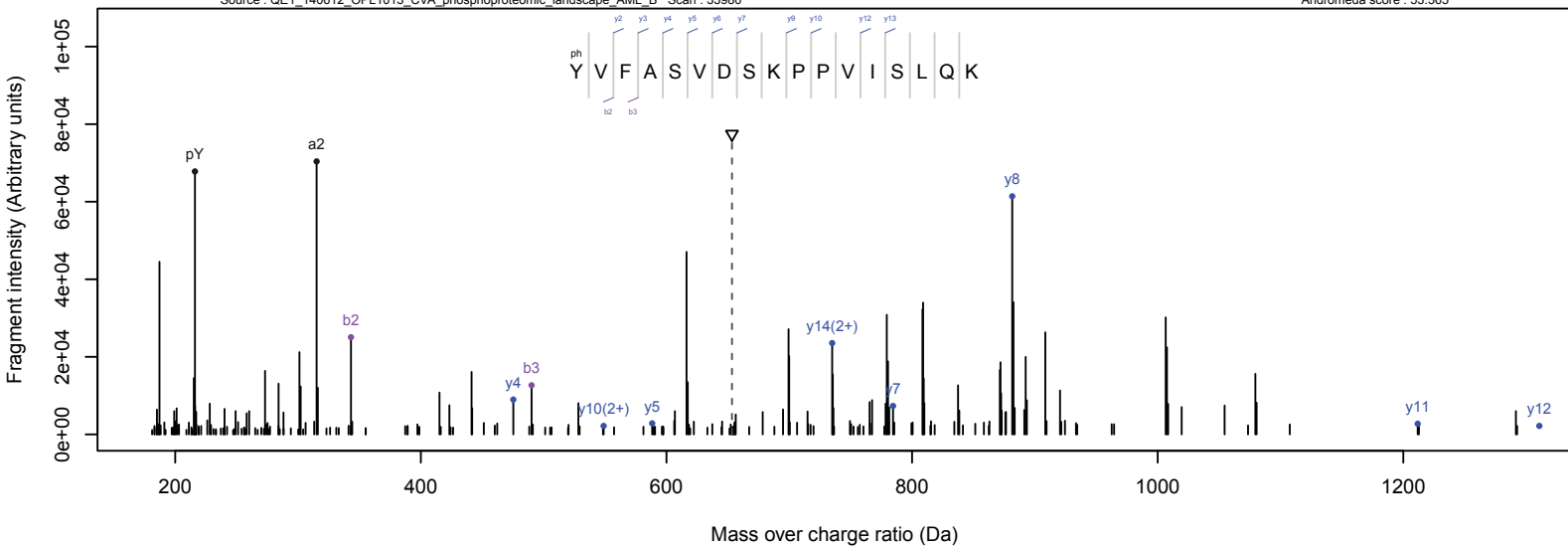
Andromeda score : 53.565

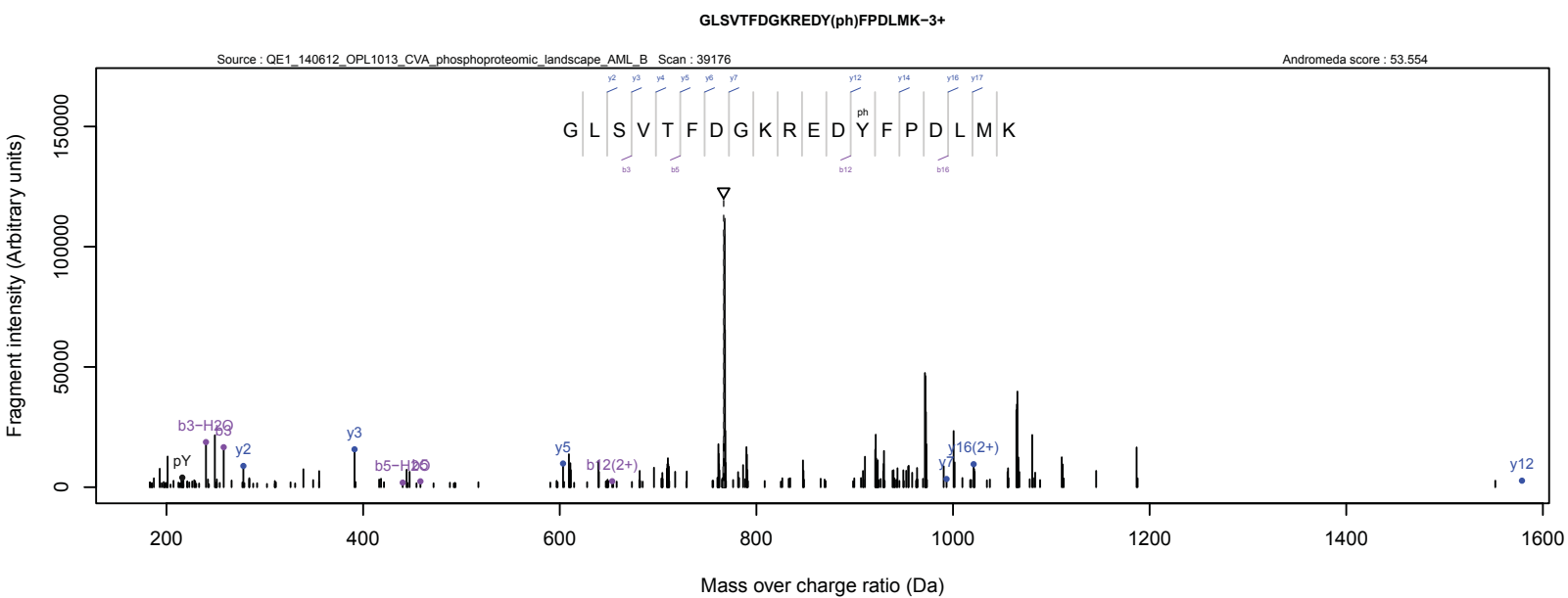
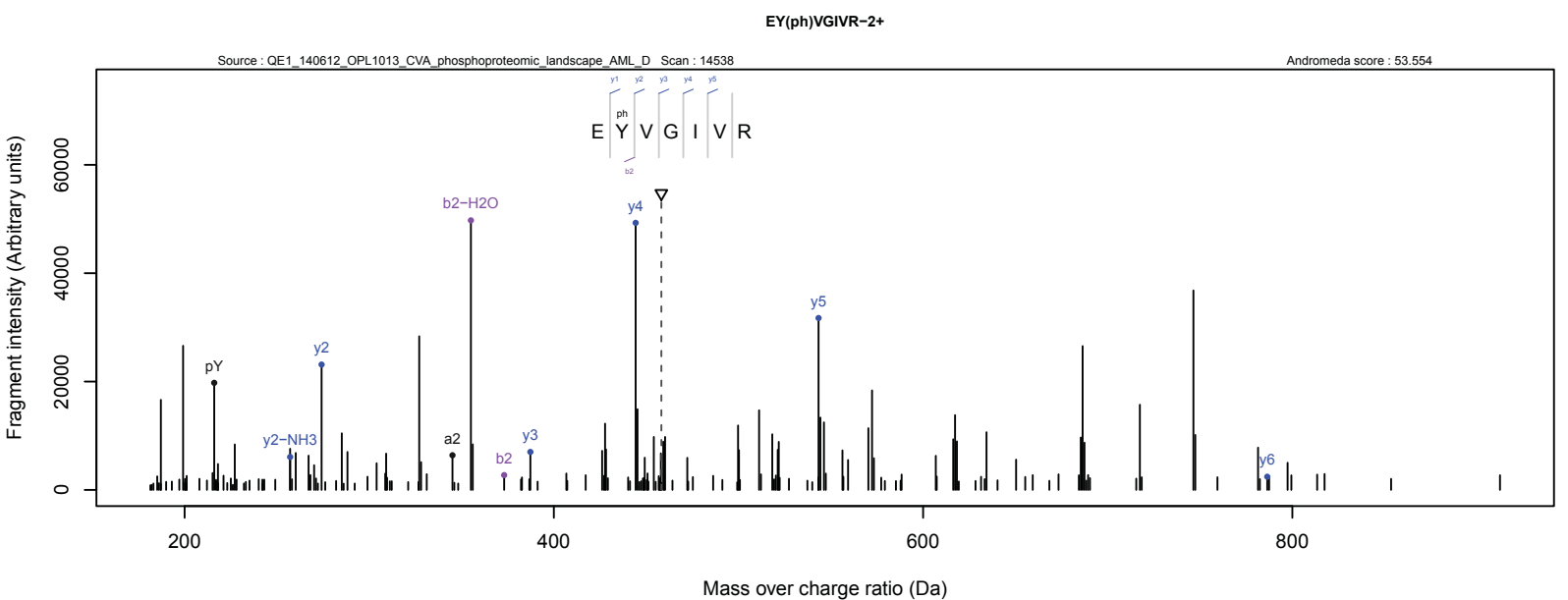
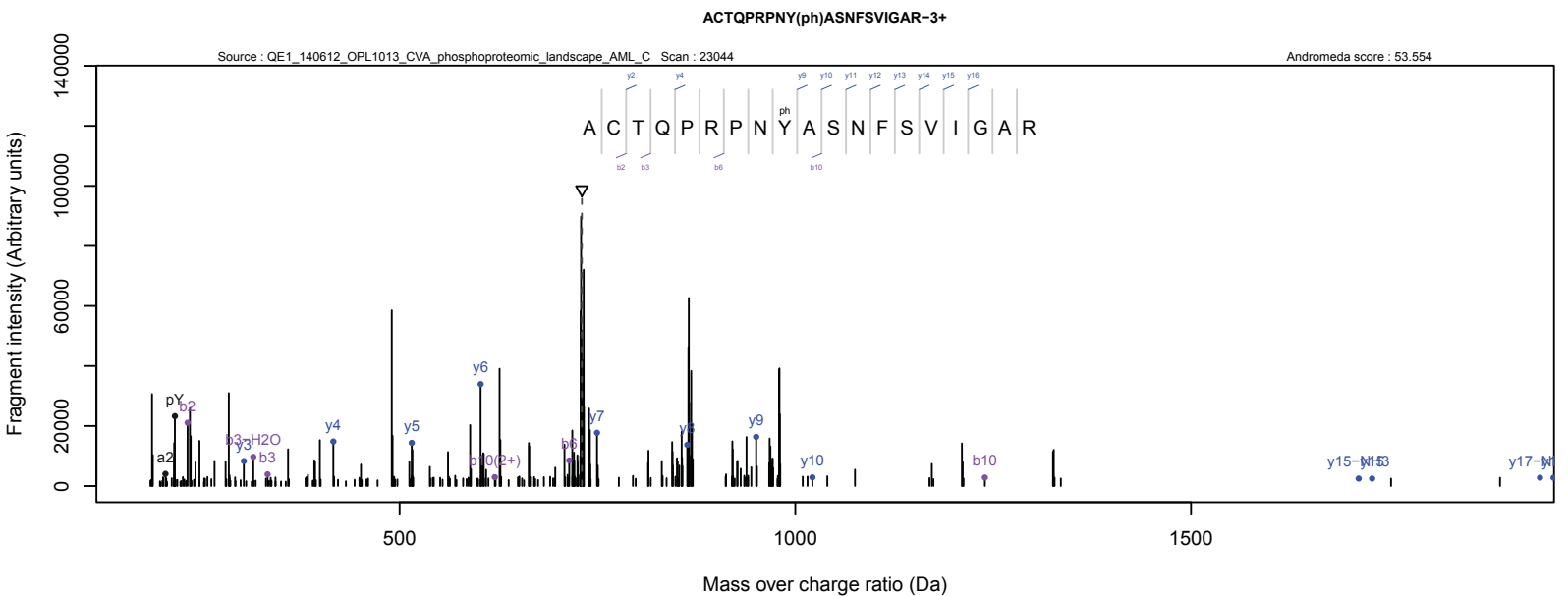


**Y(ph)VFASVDSKPPVISLQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33980

Andromeda score : 53.565

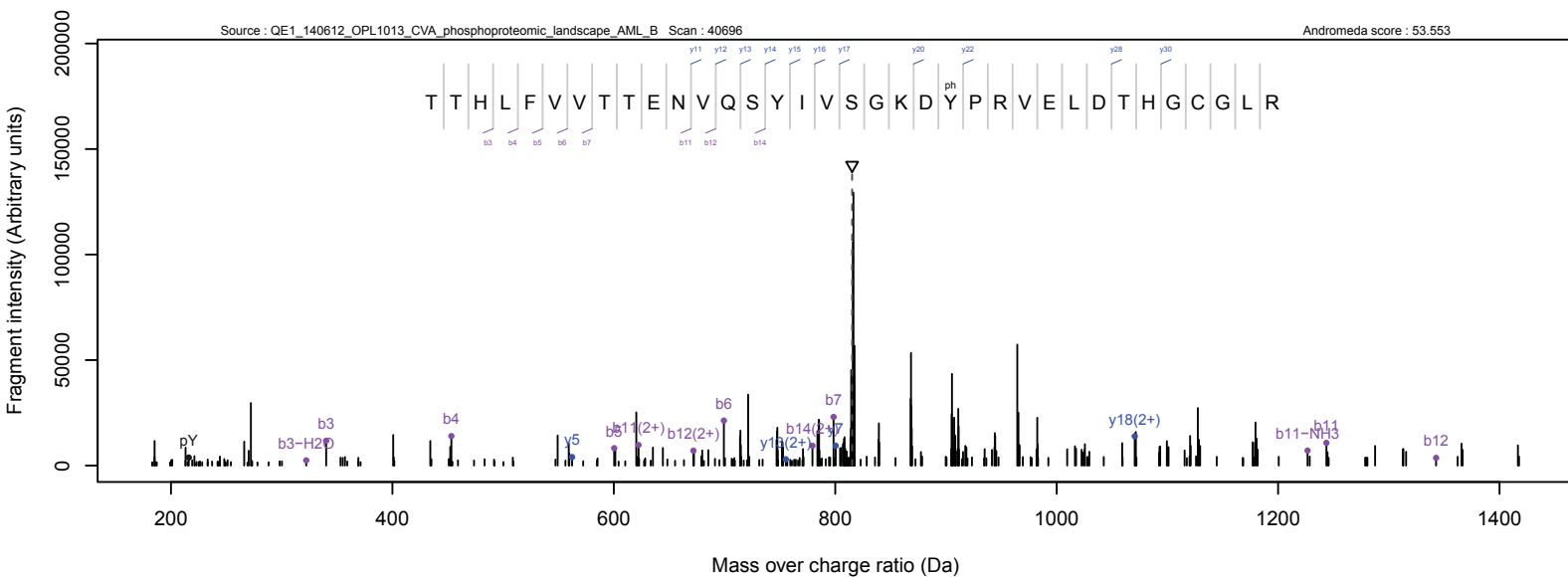




TTHLFVVTENQSYIVSGKDY(ph)PRVELDTHGCGLR-5+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 40696

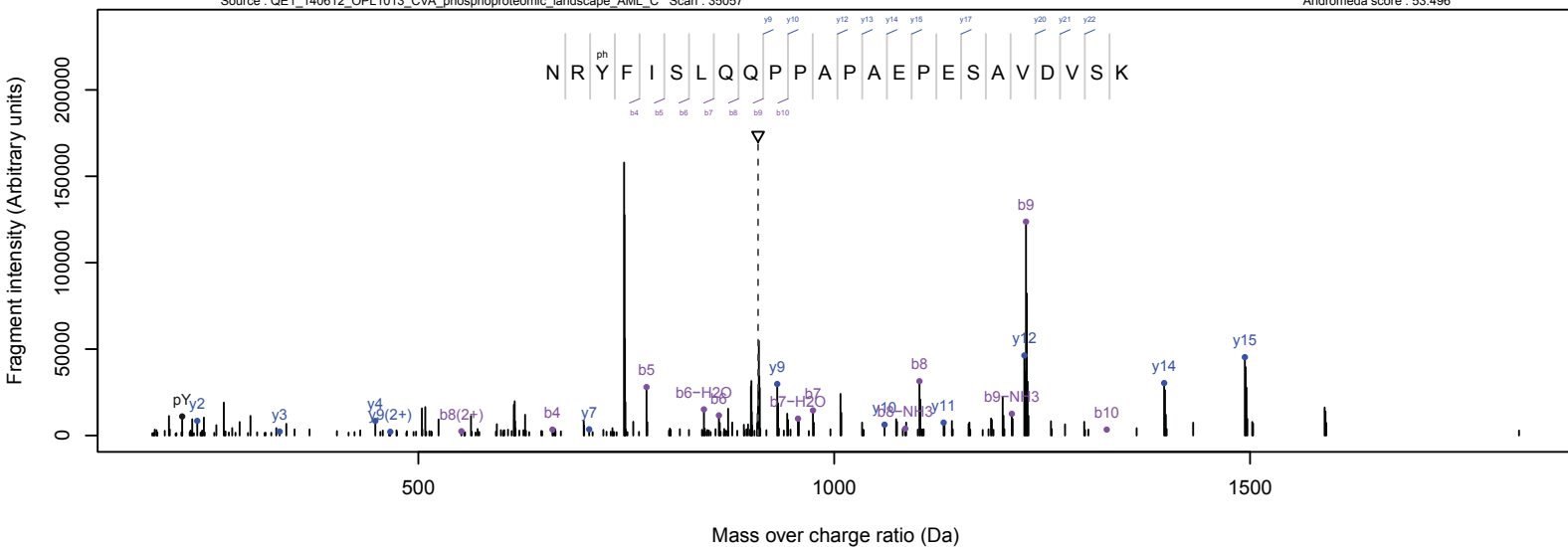
Andromeda score : 53.553



NRY(ph)FISLQQPPAPEESAVDVS K-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35057

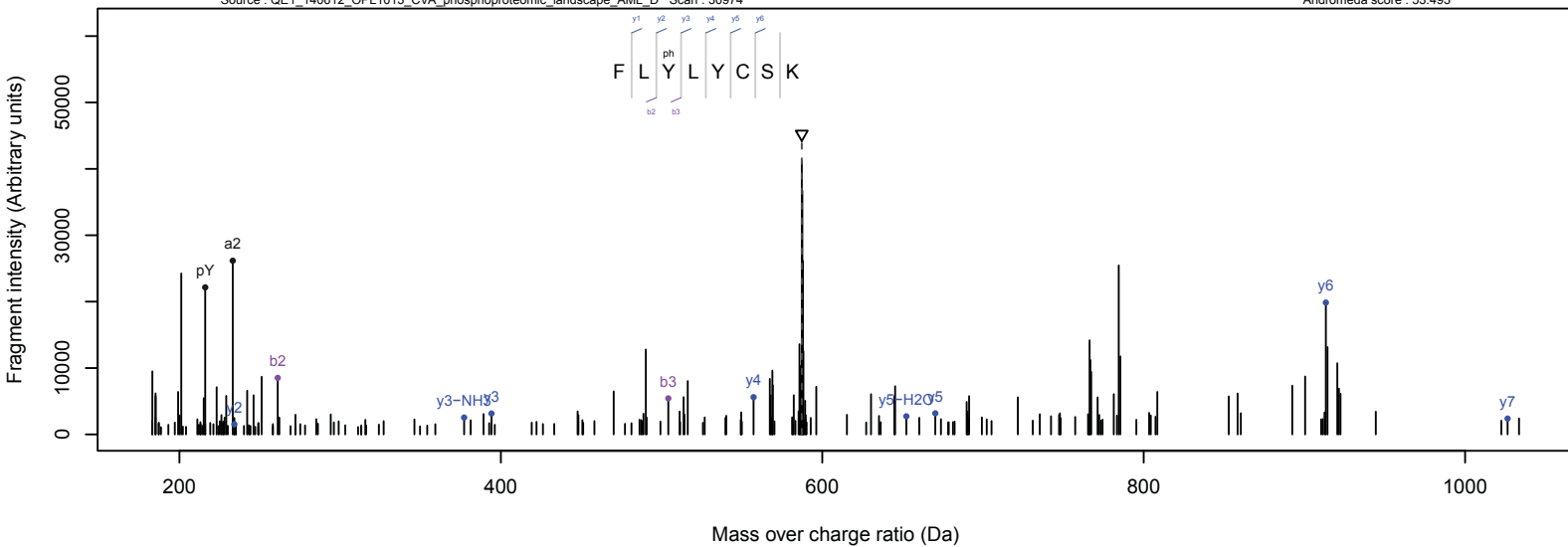
Andromeda score : 53.496



FLY(ph)LYCSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 30974

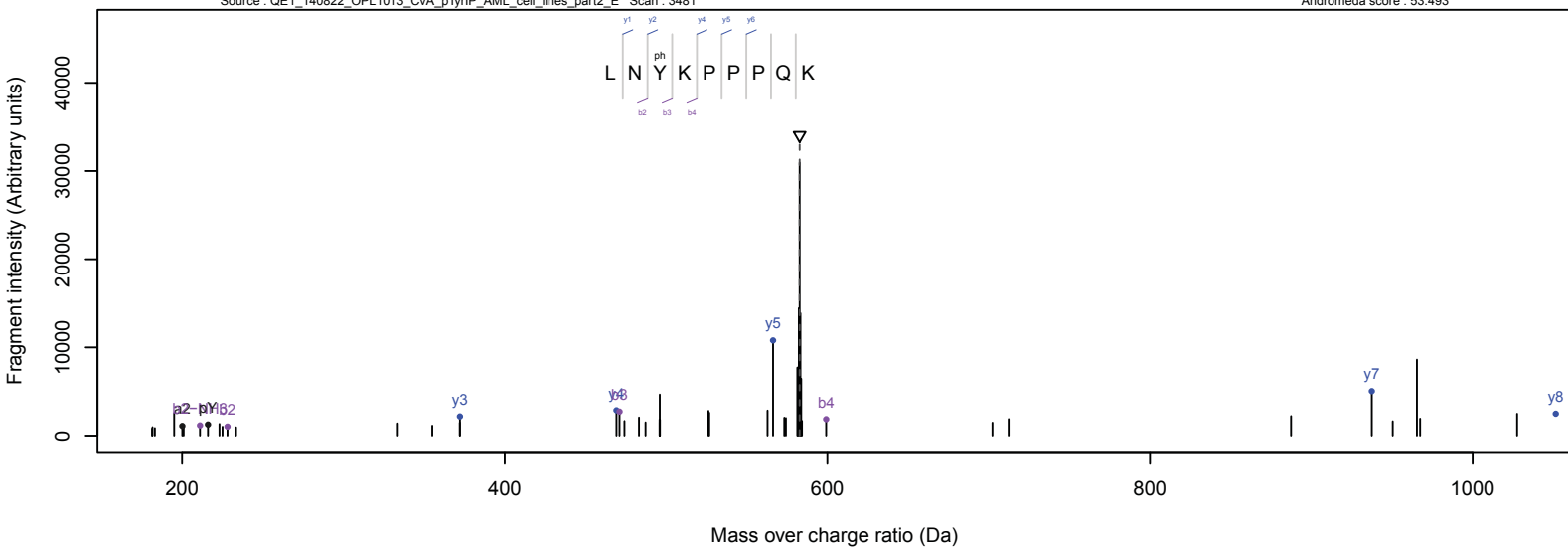
Andromeda score : 53.493



LNY(ph)KPPPQK-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 3481

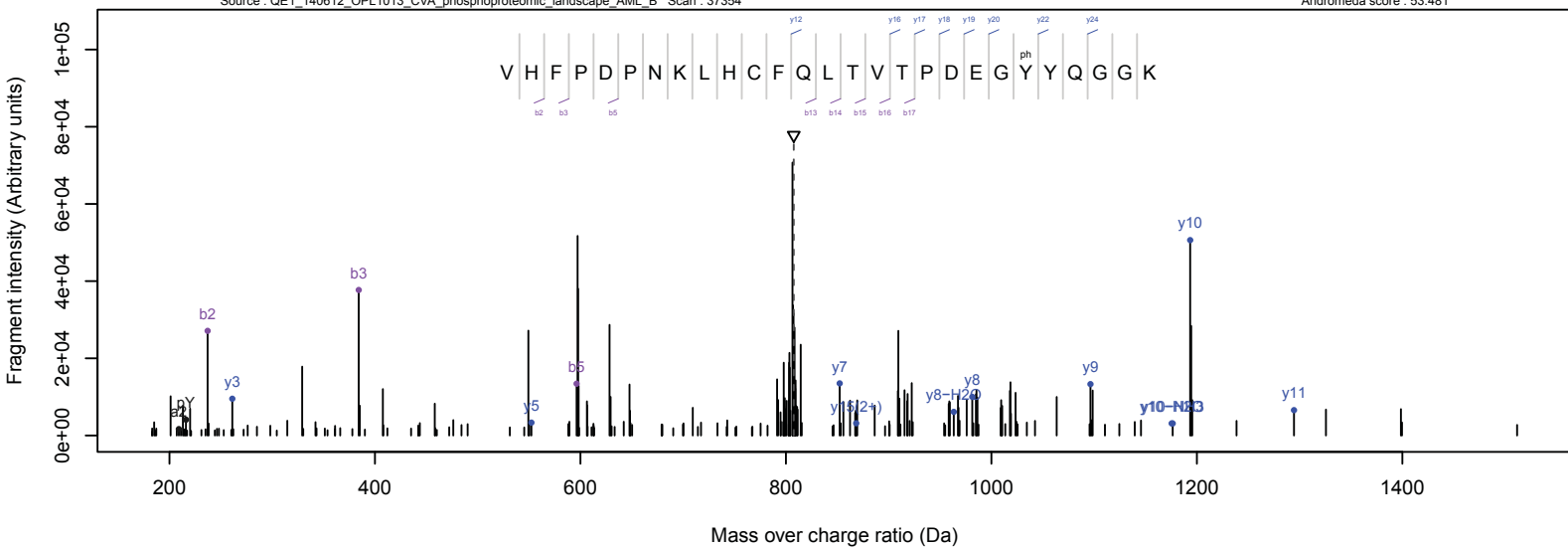
Andromeda score : 53.493



VHFPDPNKLHCFQLTVPDEGY(ph)YQGK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37354

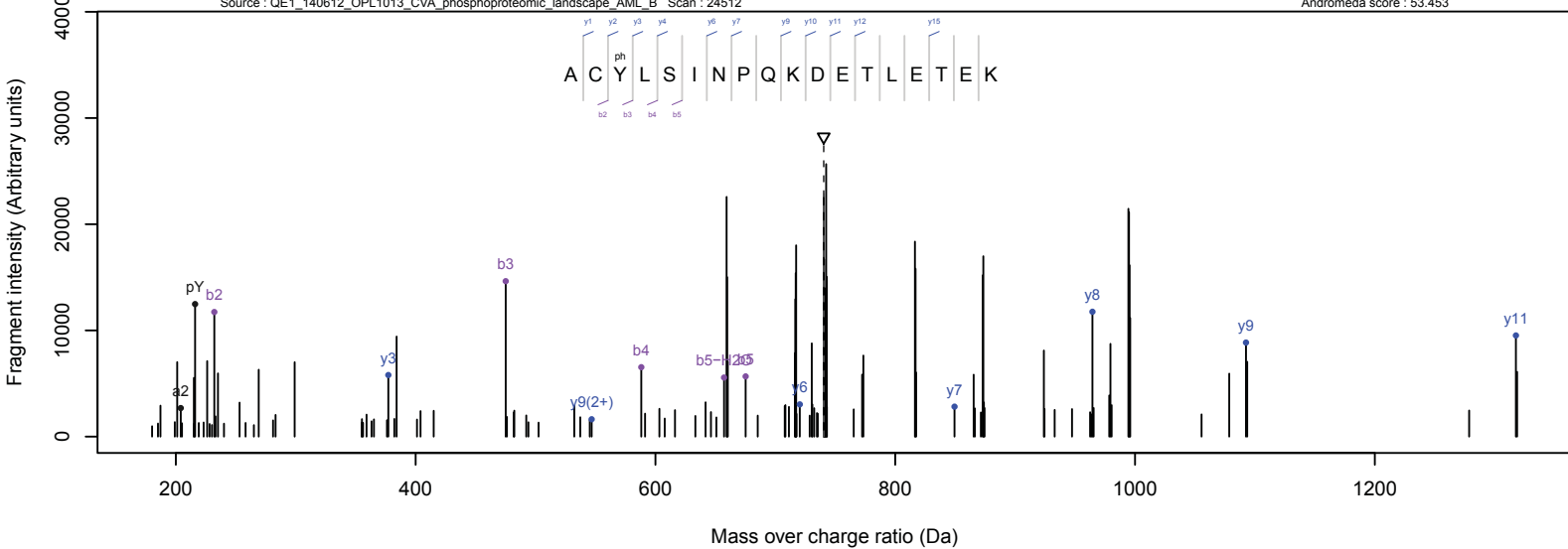
Andromeda score : 53.481



ACY(ph)LSINPQKDETLETEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24512

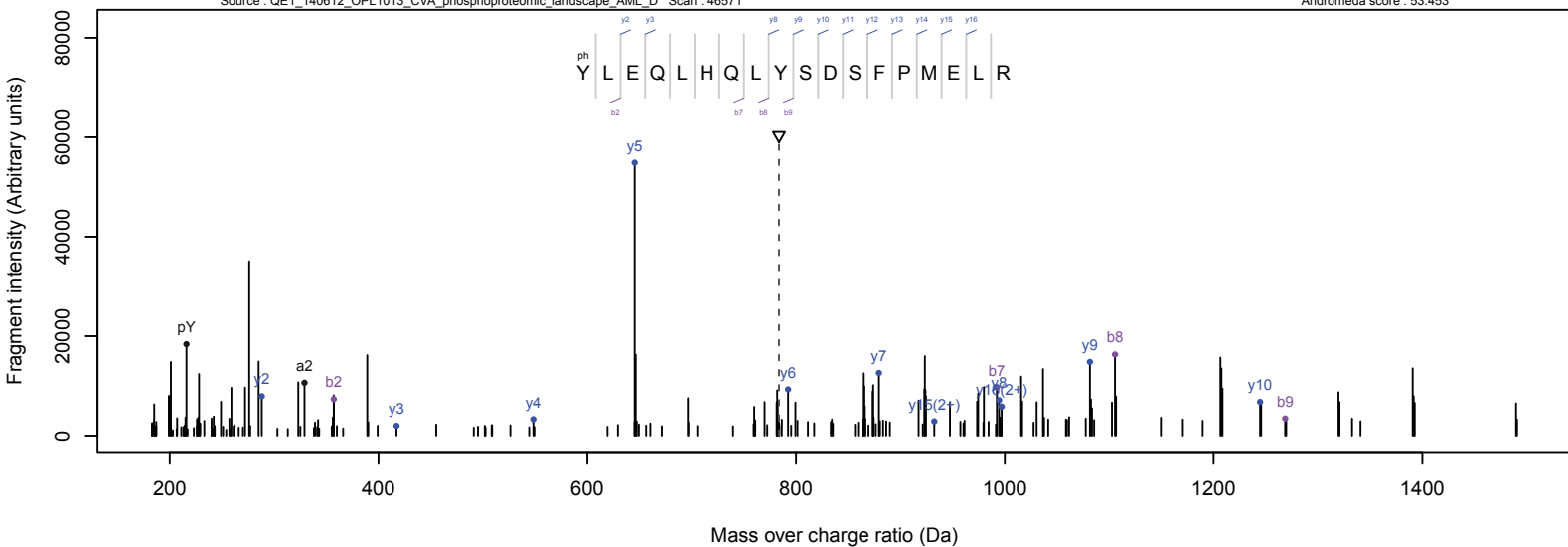
Andromeda score : 53.453



## Y(ph)LEQLHQLYSDSFPMELR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 46571

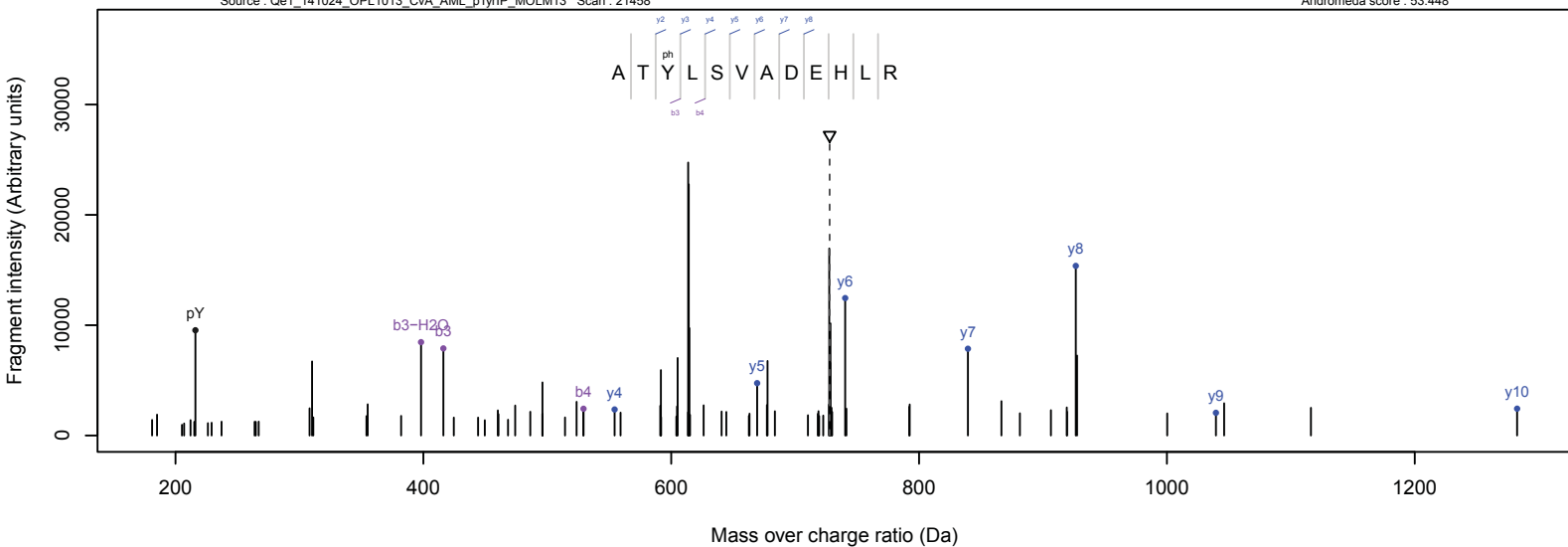
Andromeda score : 53.453



## ATY(ph)LSVADEHLR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 21458

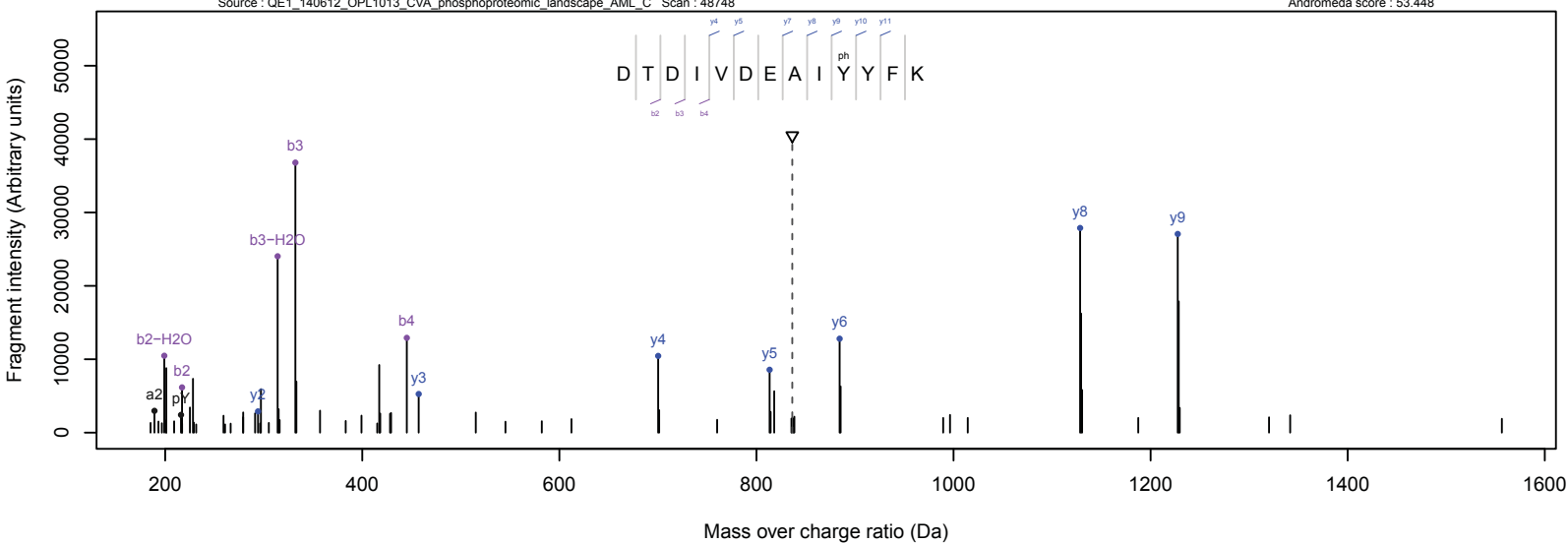
Andromeda score : 53.448



## DTDIVDEAIYFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 48748

Andromeda score : 53.448

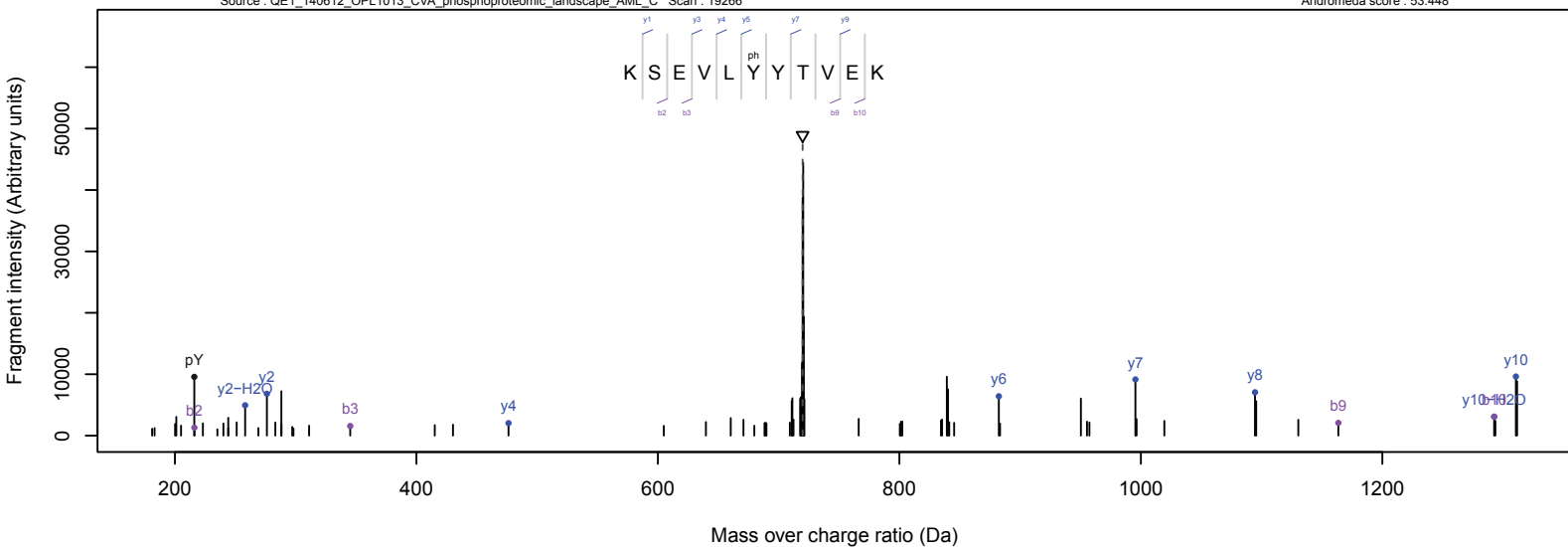




**KSEVLY(ph)YTVEK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19266

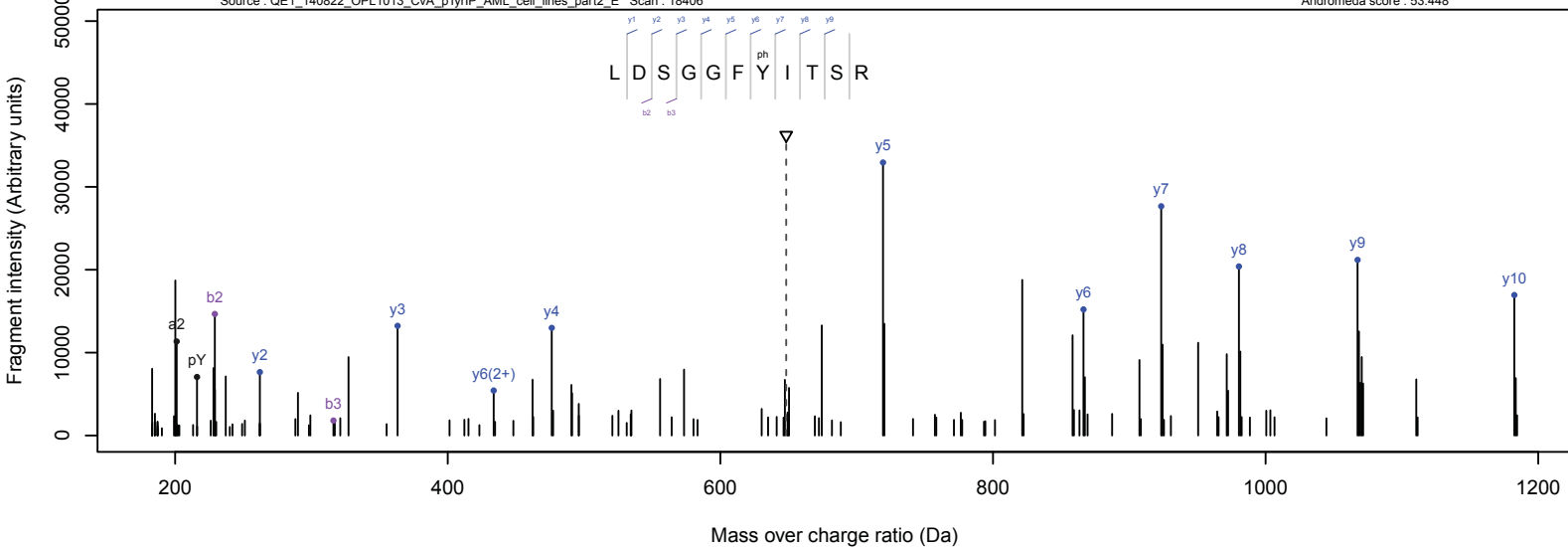
Andromeda score : 53.448



**LDSGGFY(ph)ITSR-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 18406

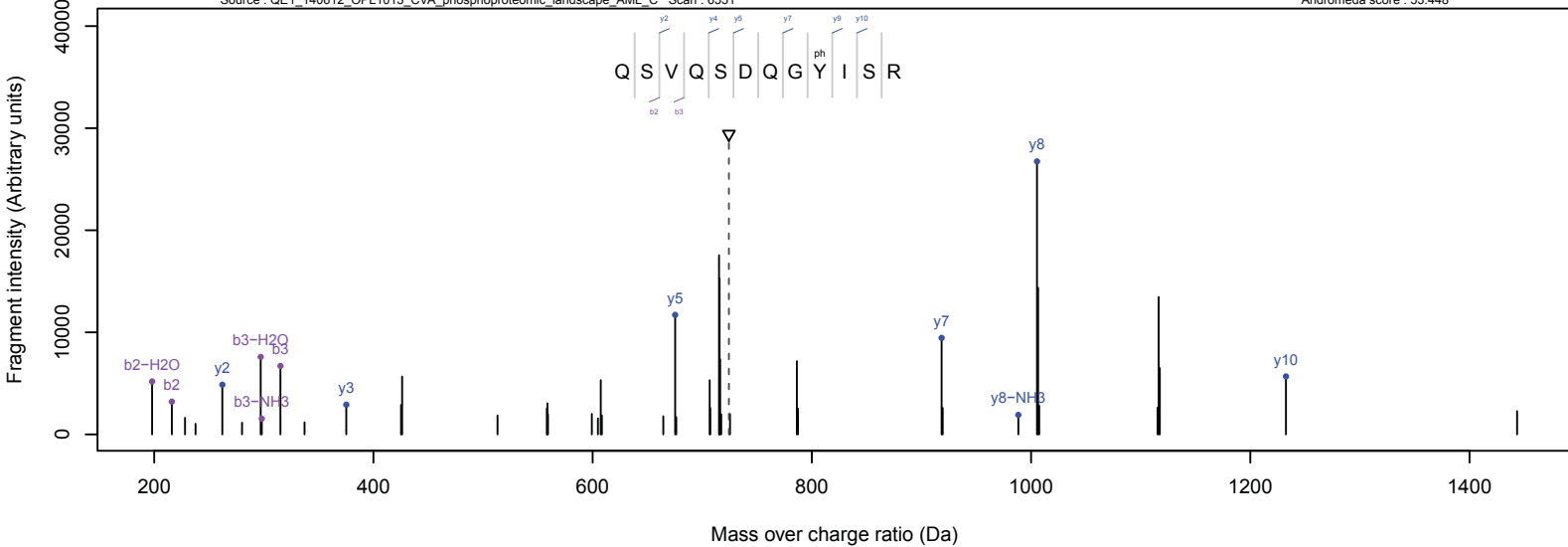
Andromeda score : 53.448



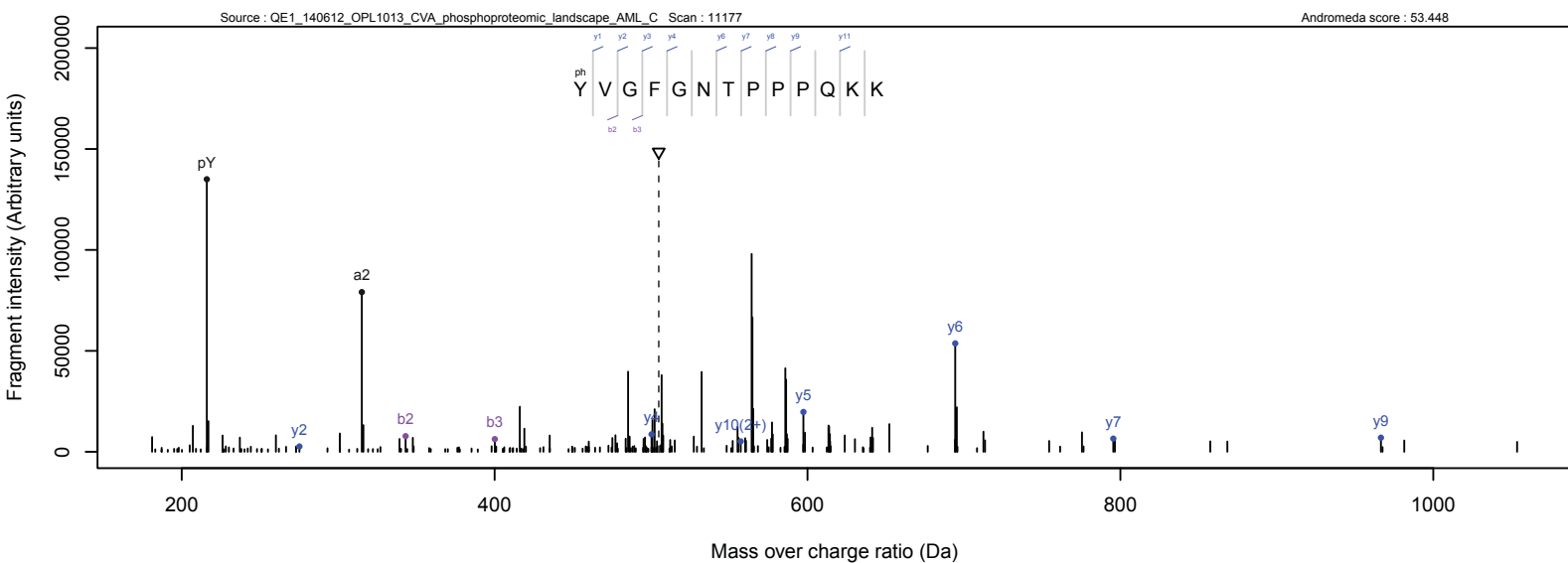
**QSVQSDQGY(ph)ISR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6551

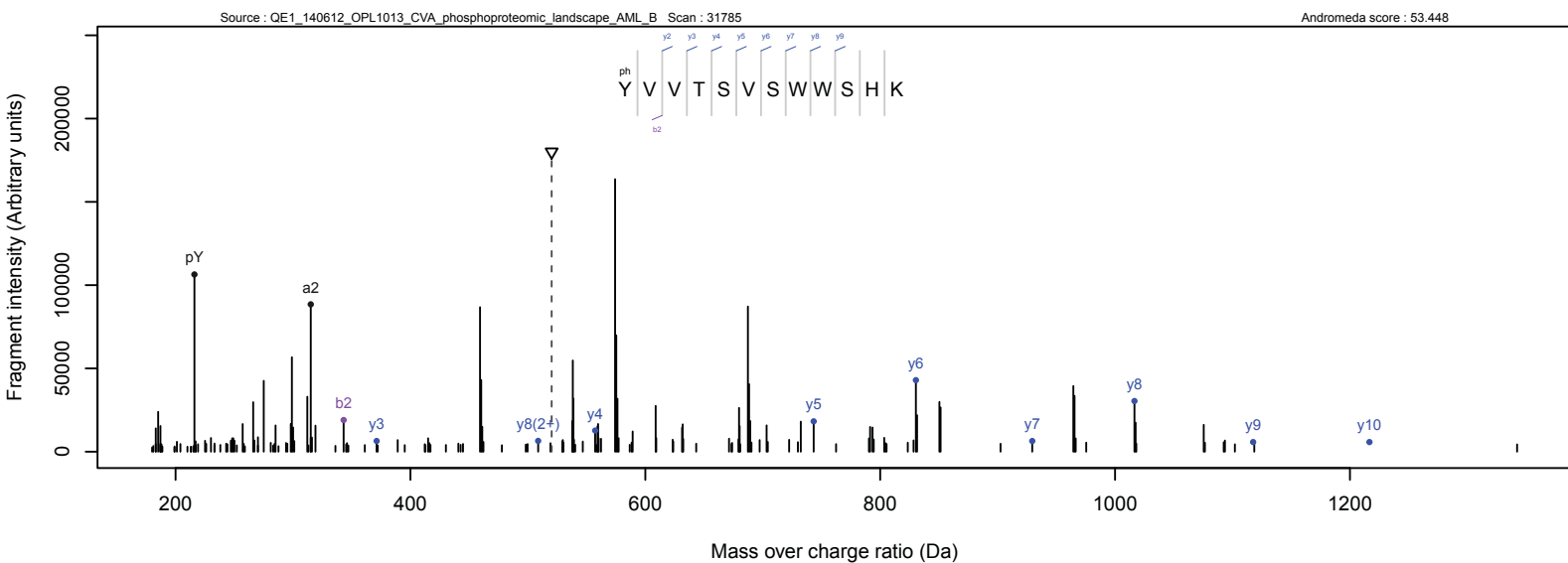
Andromeda score : 53.448



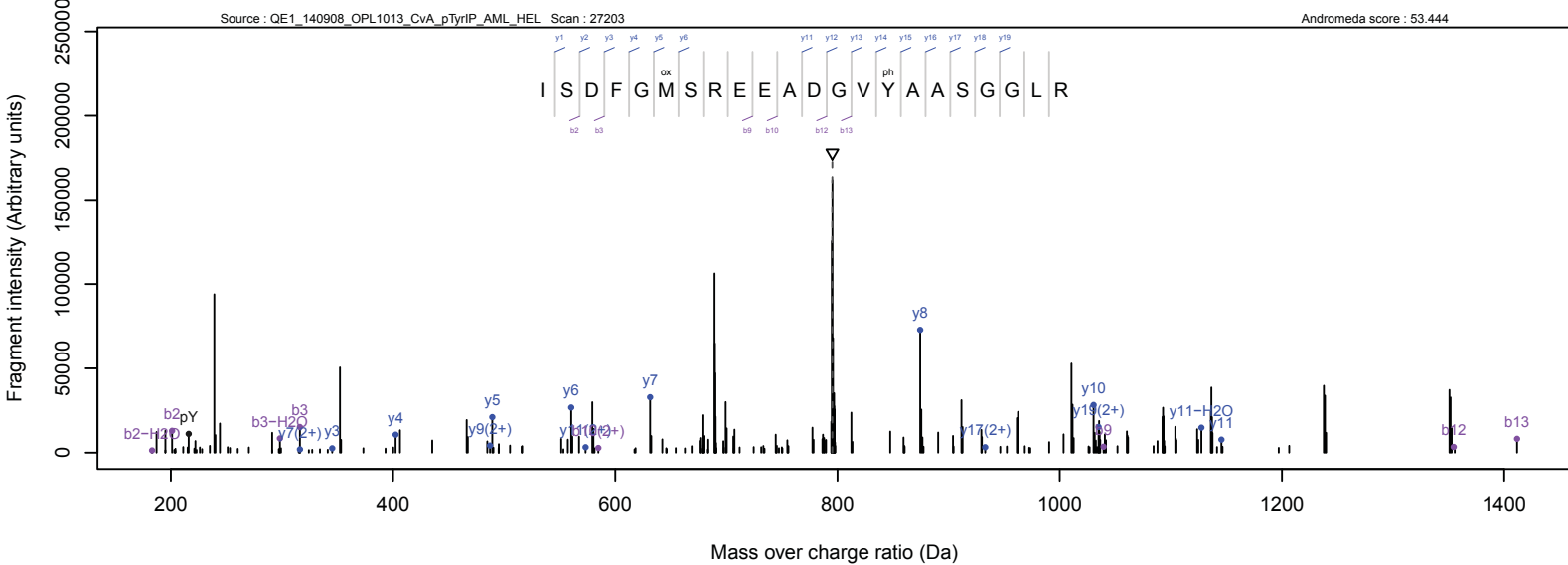
**Y(ph)VGFGNTPPPQKK-3+**



**Y(ph)VVTSVSWWSHK-3+**



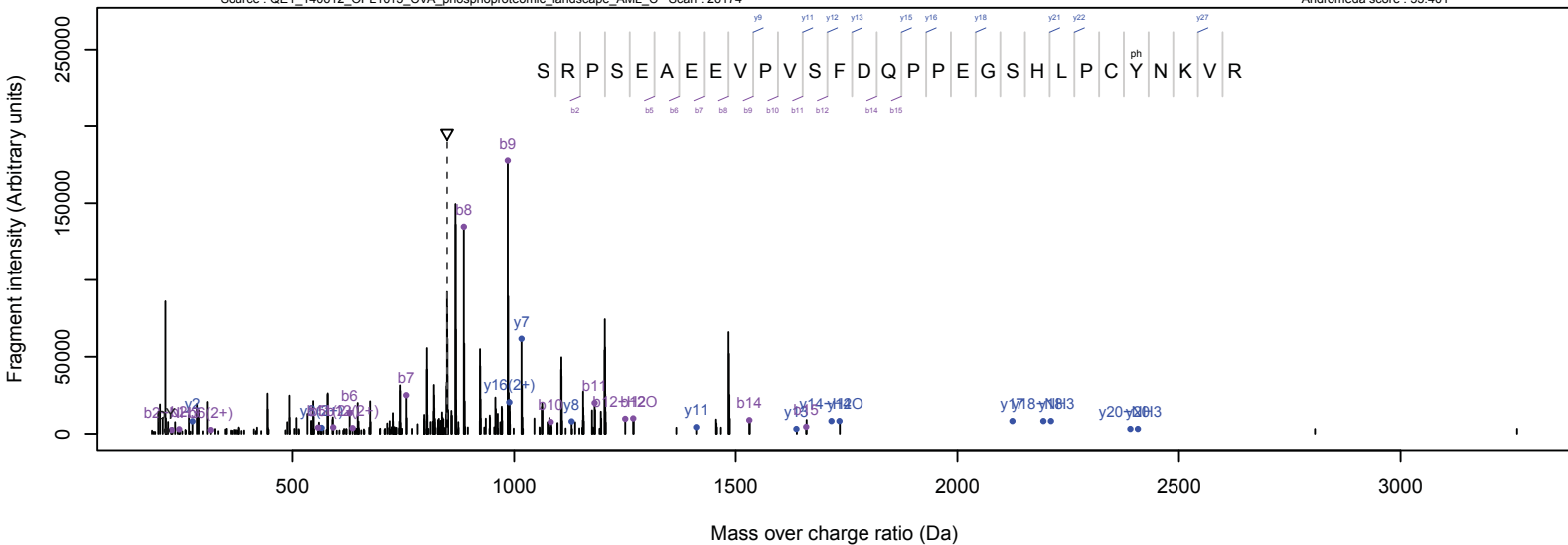
**ISDFGM(ox)SREEADGVY(ph)AASGGLR-3+**



SRPSEAEVPV<sup>S</sup>FDQPPEGSHLPC<sup>Y</sup>(ph)NKVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26174

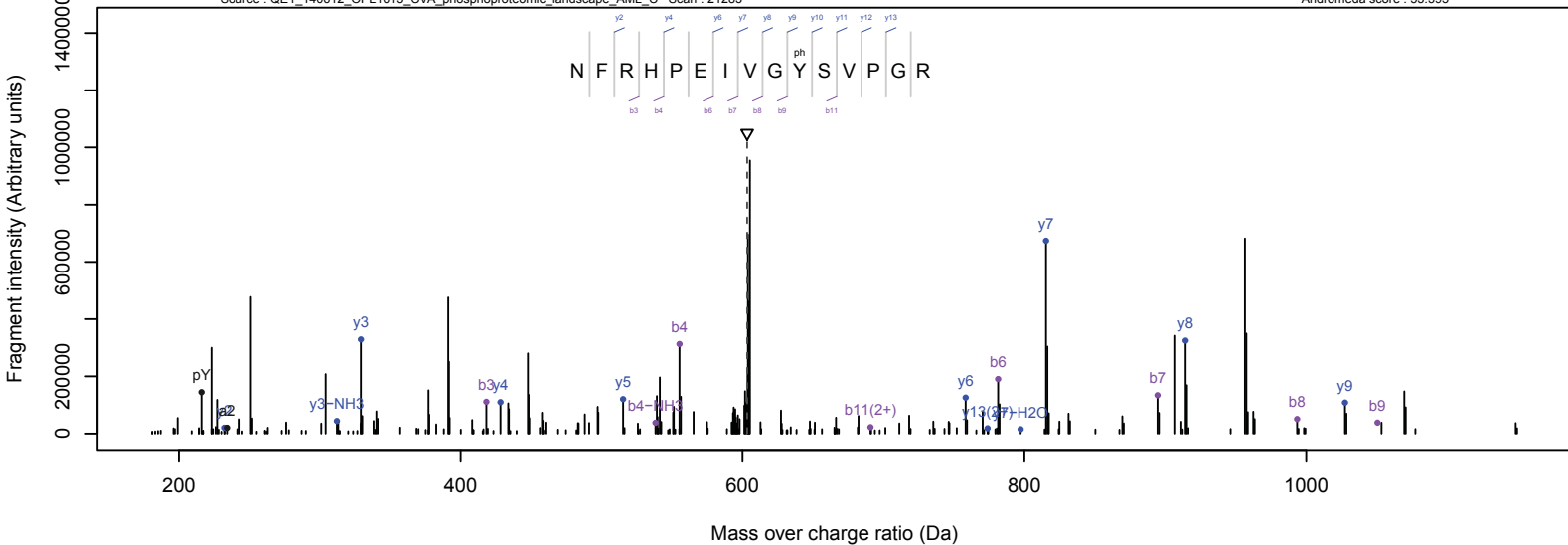
Andromeda score : 53.401



NFRHPEIVG<sup>Y</sup>SV<sup>P</sup>GR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21283

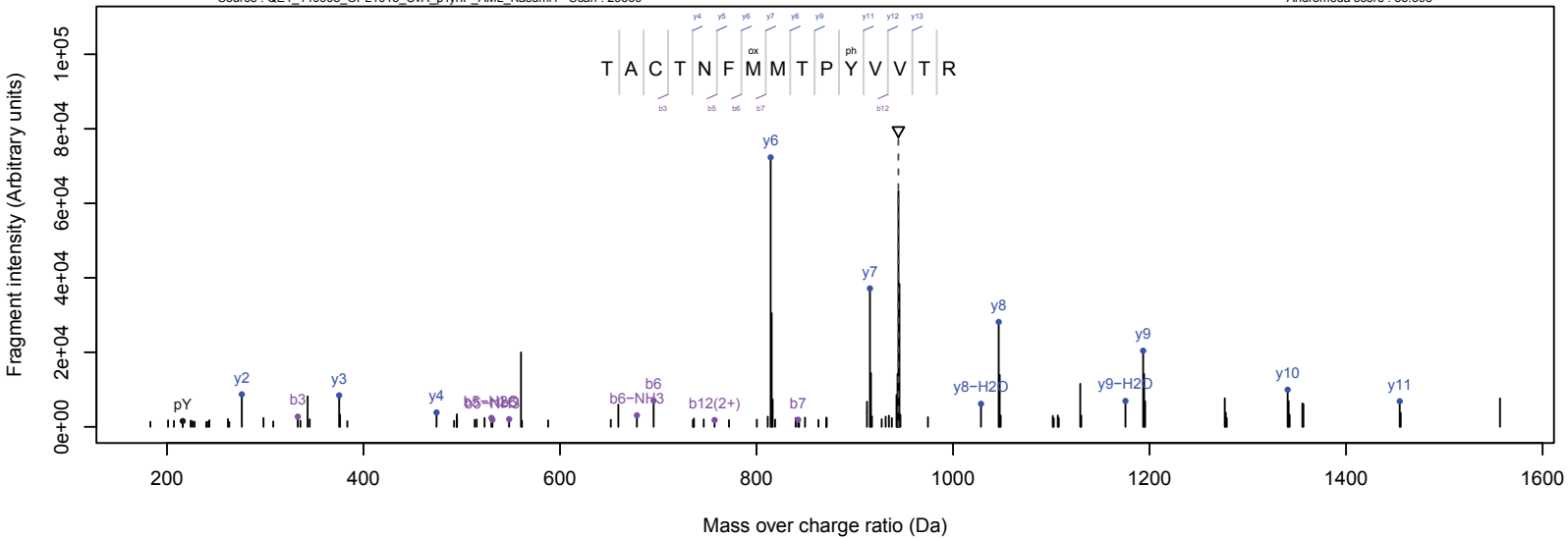
Andromeda score : 53.395



TACTNFM<sup>ox</sup>MTPY<sup>ph</sup>VVTR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 29569

Andromeda score : 53.395

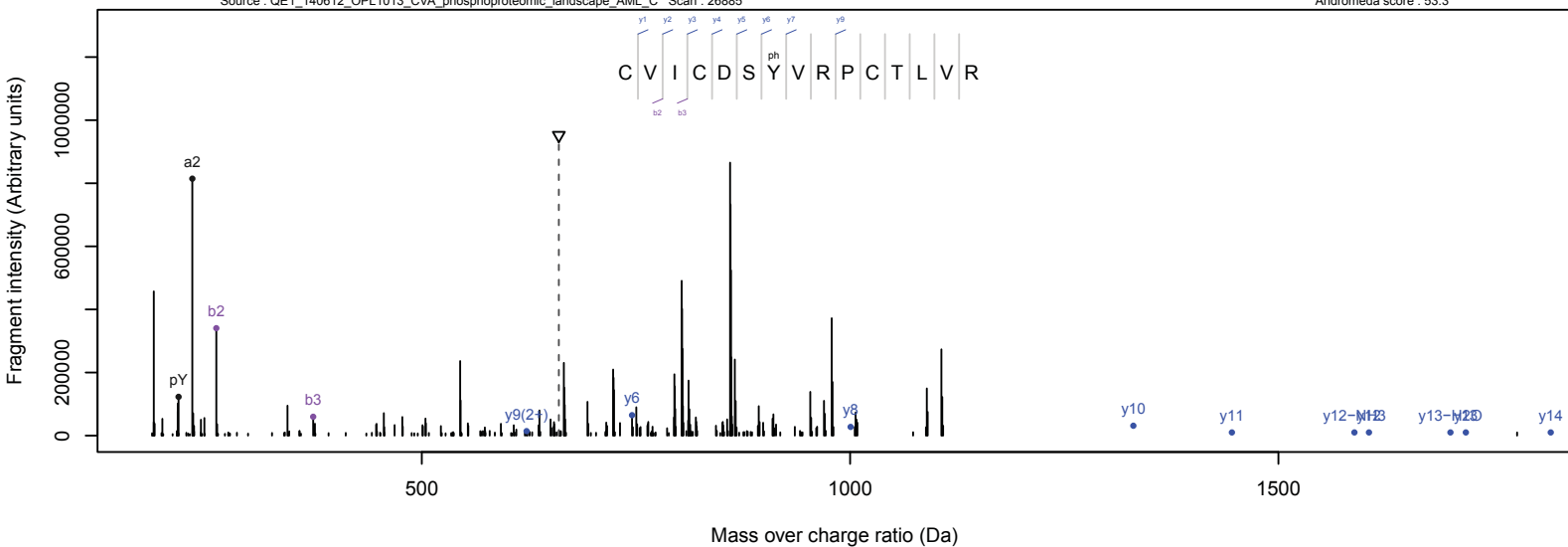




CVICDSY(ph)VRPCTLVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26885

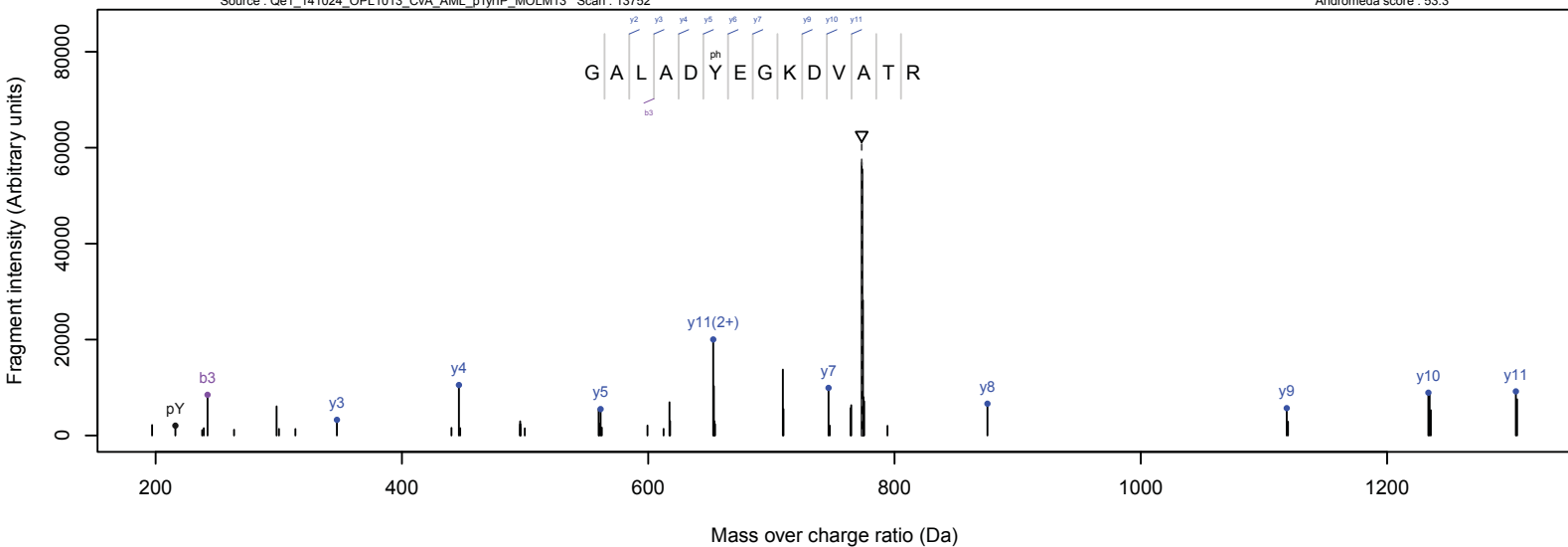
Andromeda score : 53.3



GALADY(ph)EGKDVATR-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 13752

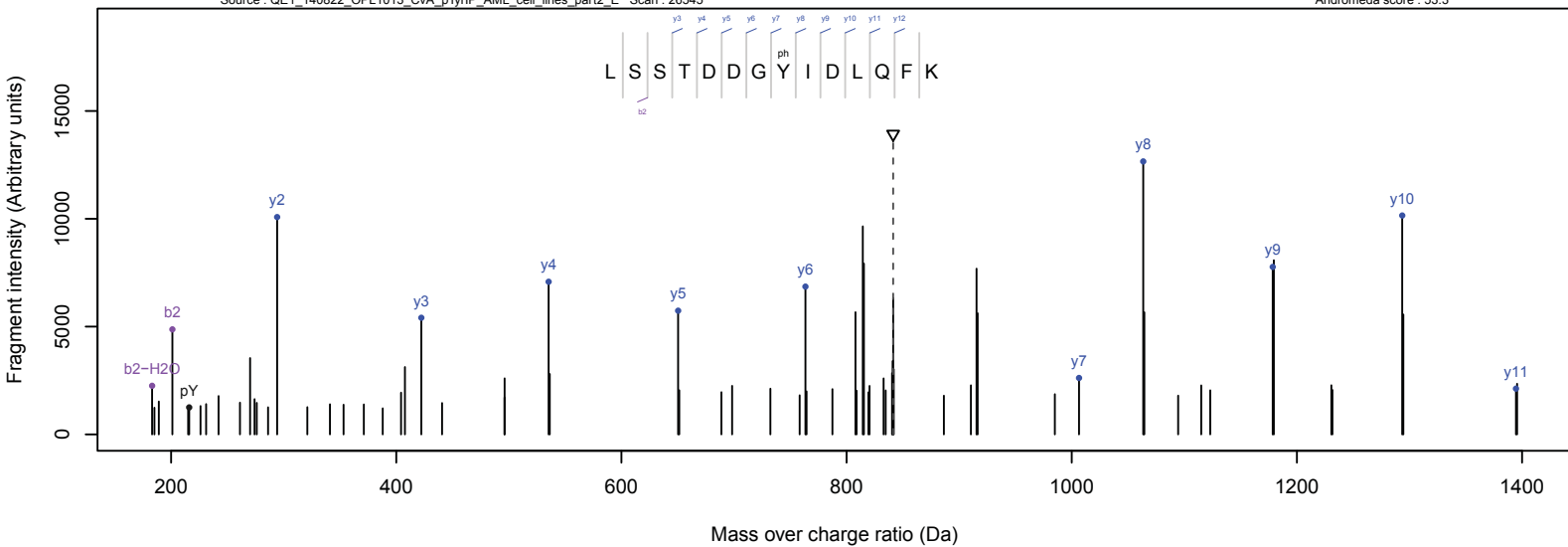
Andromeda score : 53.3



LSSTDDG(ph)IDLQFK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 26545

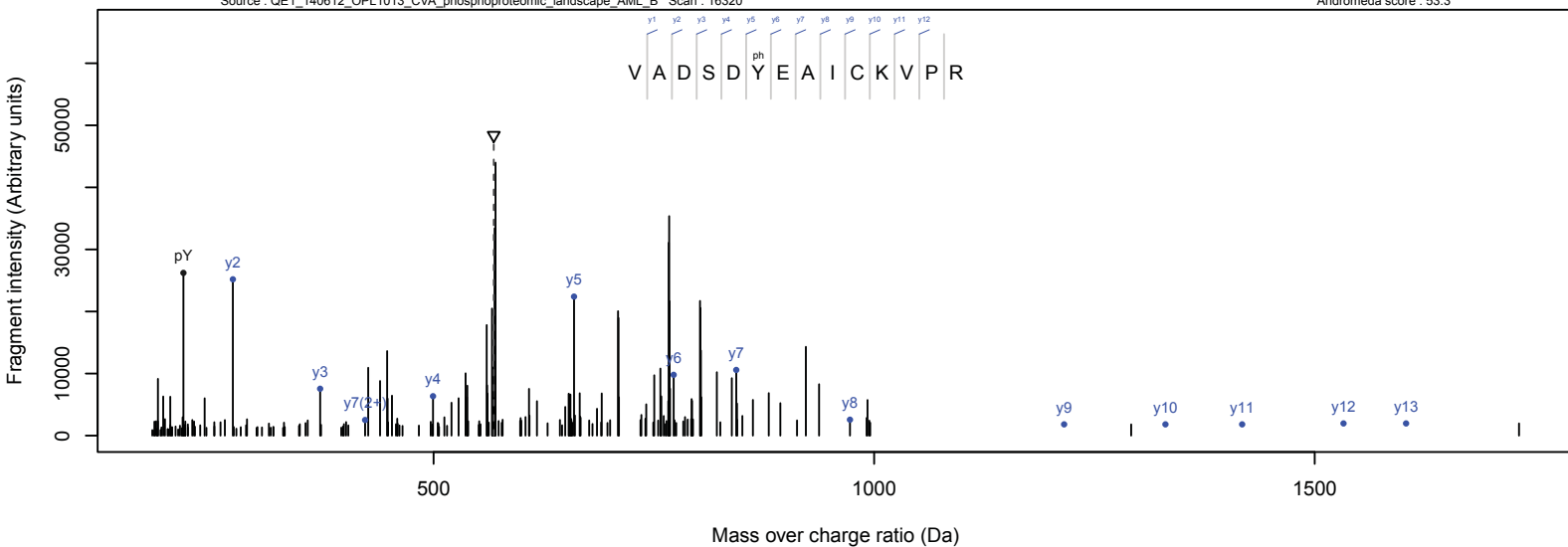
Andromeda score : 53.3



VADSDY(ph)EAICKVPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16320

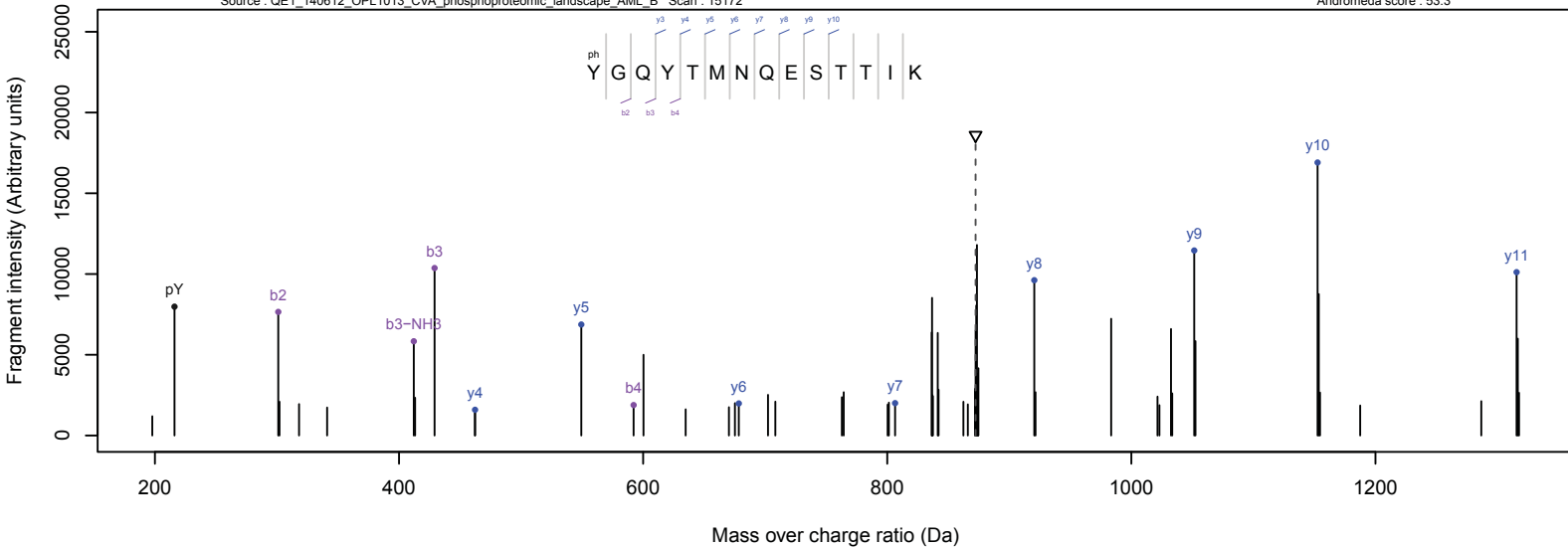
Andromeda score : 53.3



Y(ph)GQYTMNQESTTIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15172

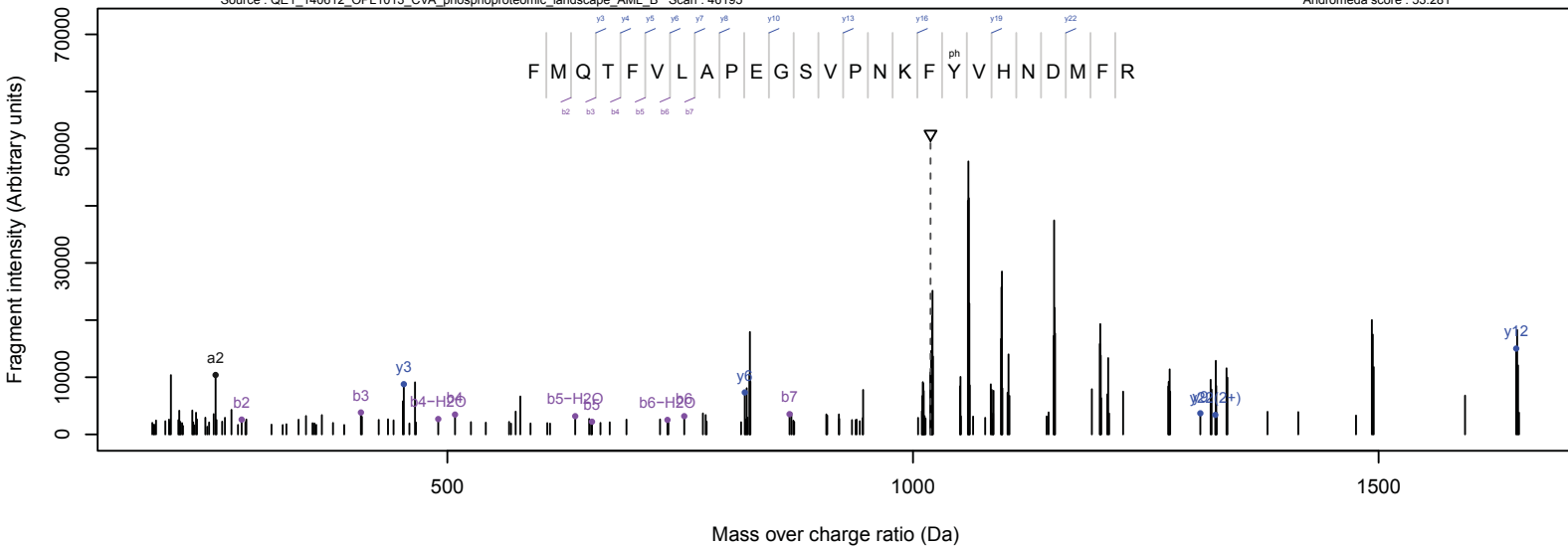
Andromeda score : 53.3



FMQTFVLAPEGSVPNKFYVHNDMFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46195

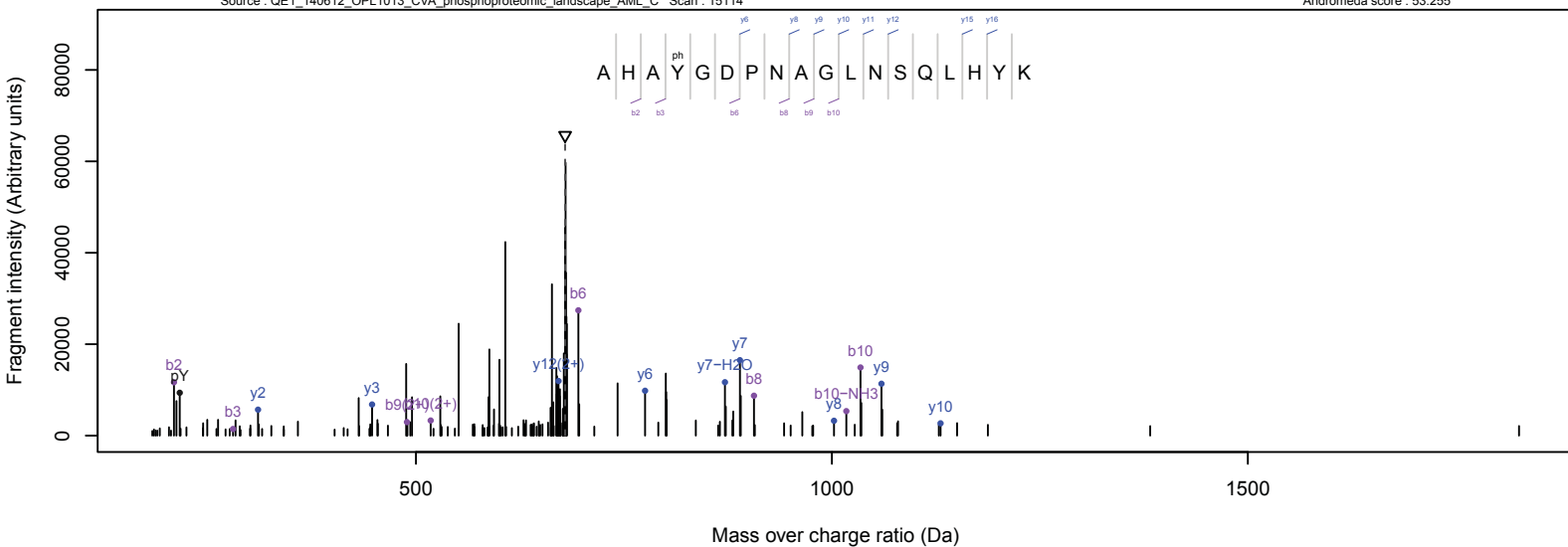
Andromeda score : 53.281



AHAY(ph)GDPNAGLNSQLHYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15114

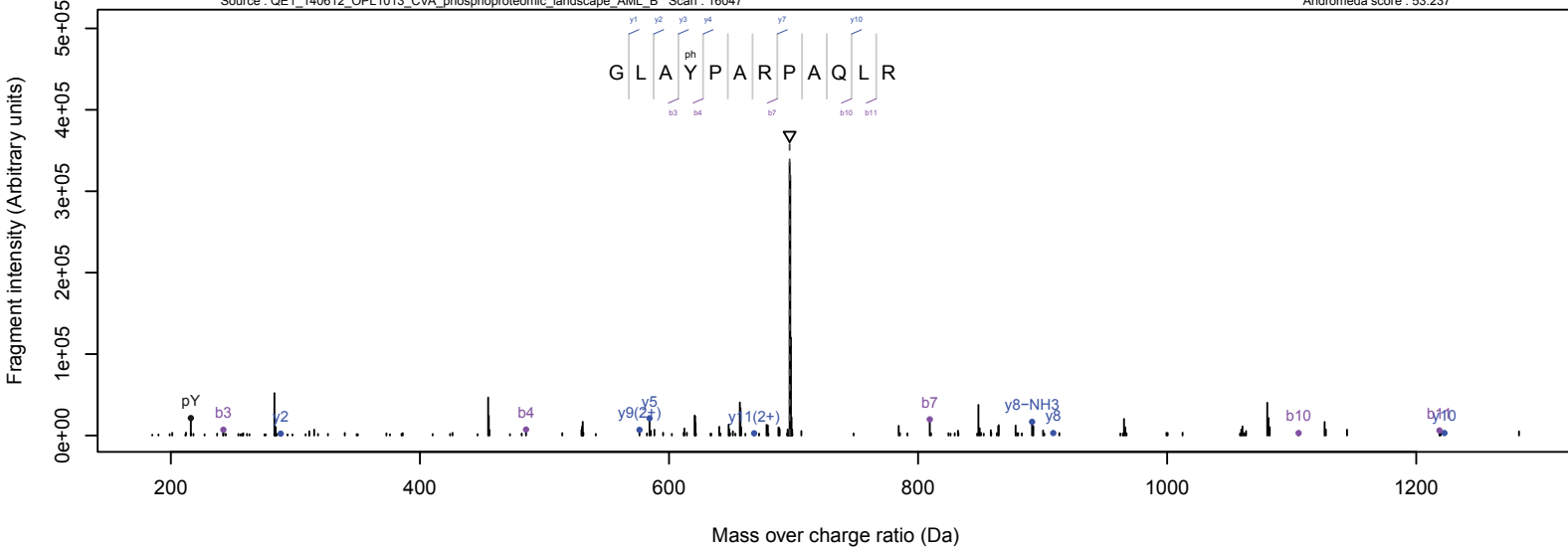
Andromeda score : 53.255



GLAY(ph)PARPAQLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 16047

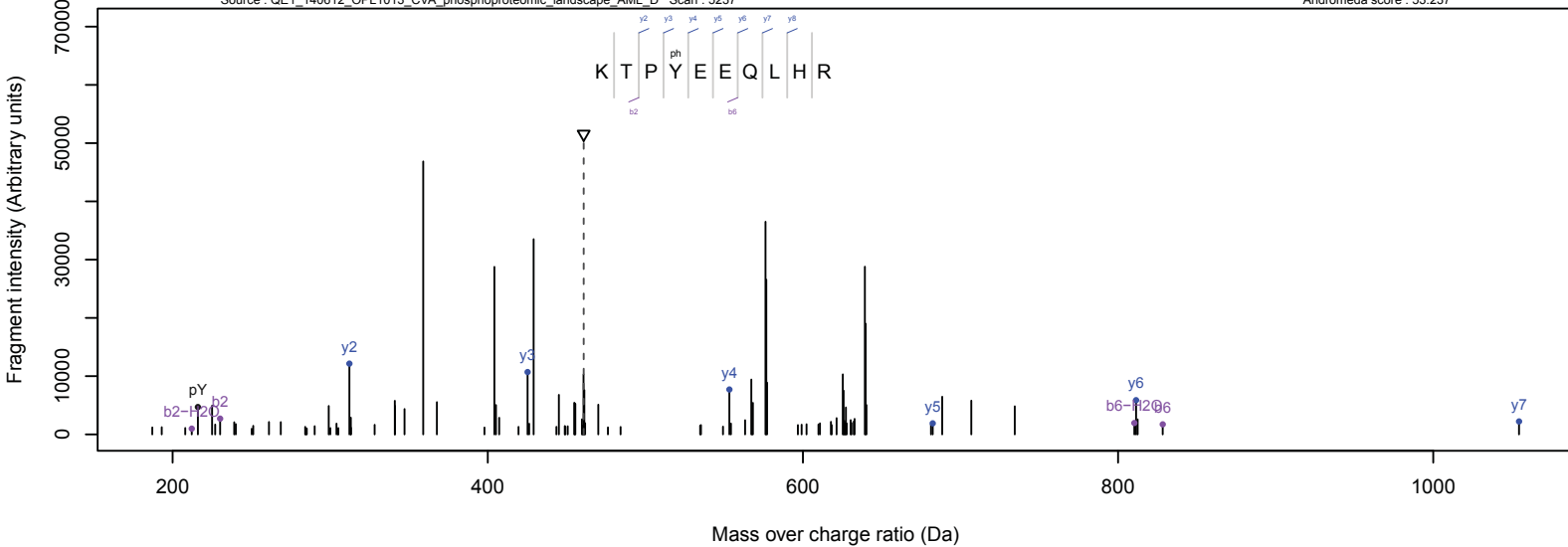
Andromeda score : 53.237



KTPY(ph)EEQLHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 5237

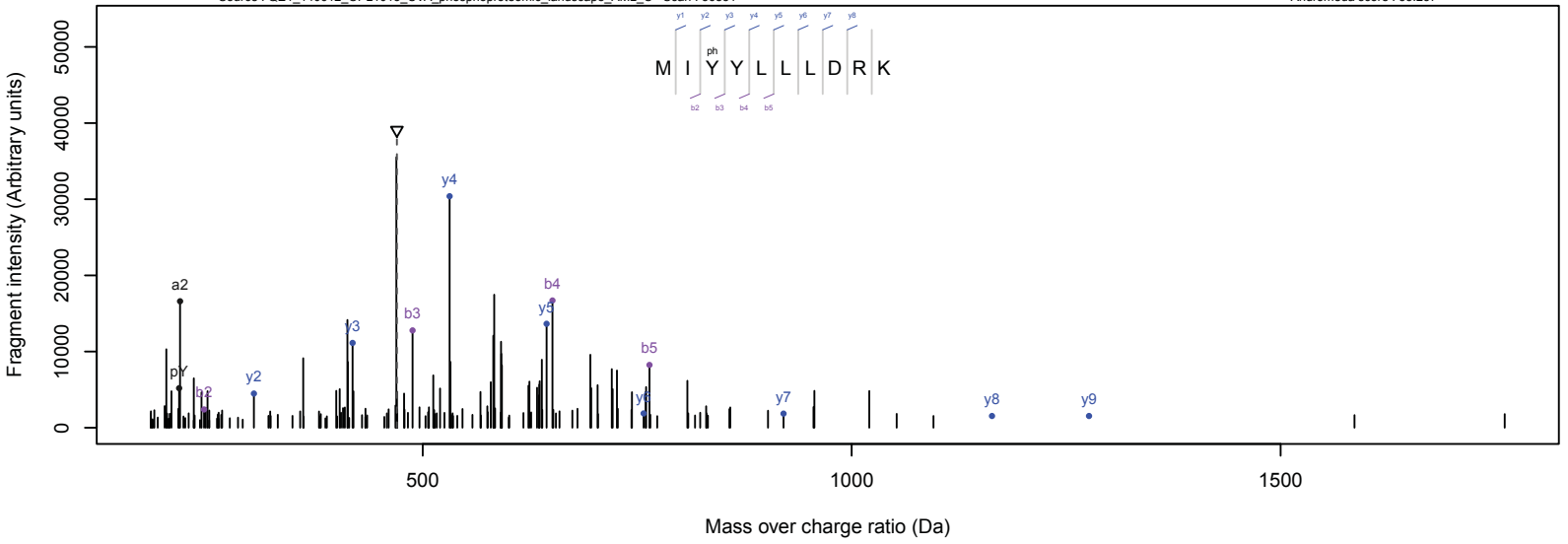
Andromeda score : 53.237



MIY(ph)YLLDRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36851

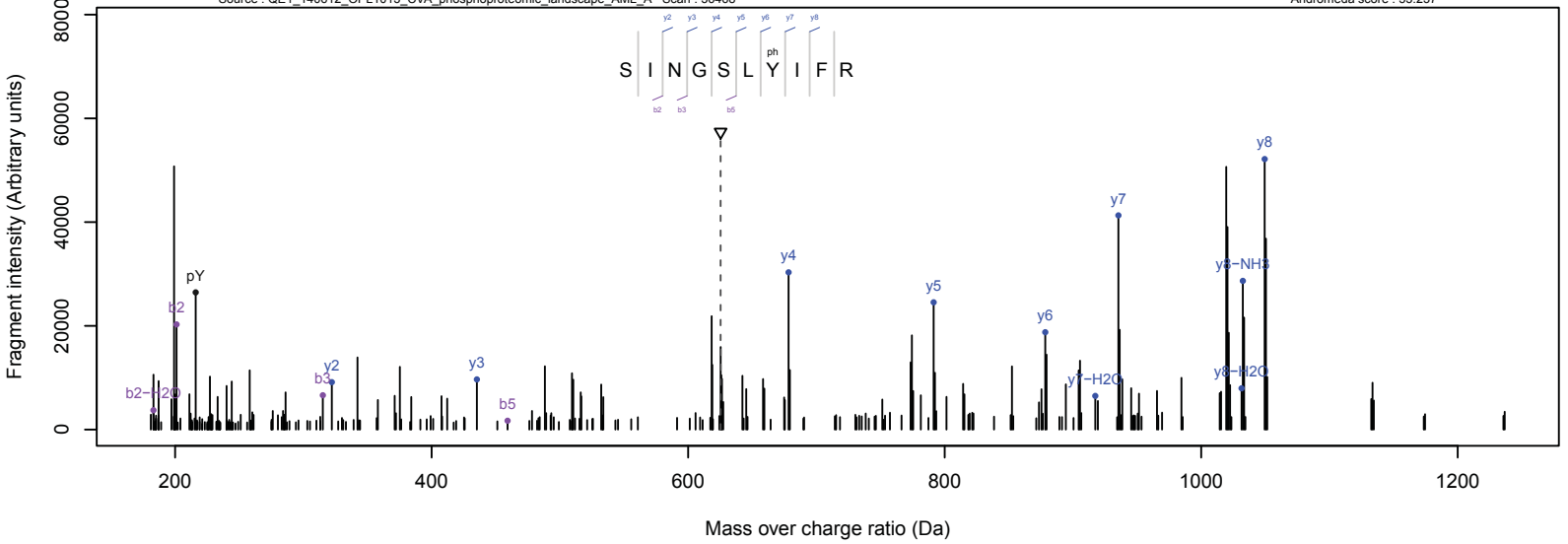
Andromeda score : 53.237



SINGSLY(ph)IFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 36468

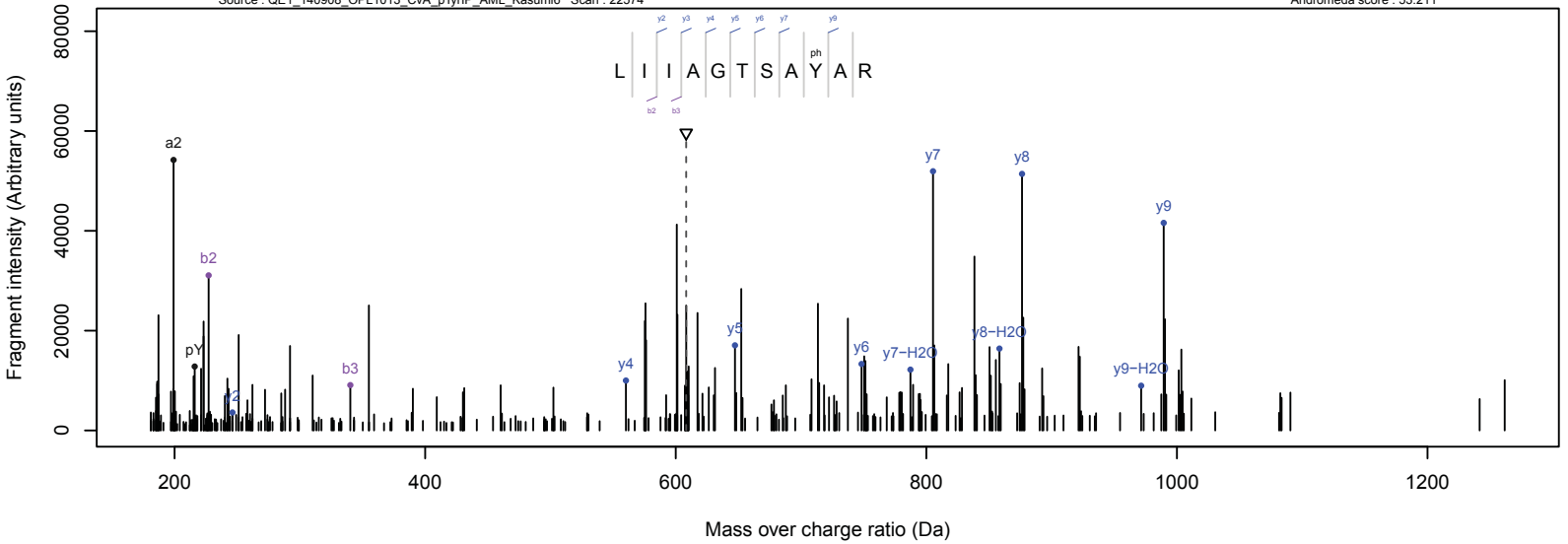
Andromeda score : 53.237



LIAGTSAY(ph)AR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 22574

Andromeda score : 53.211

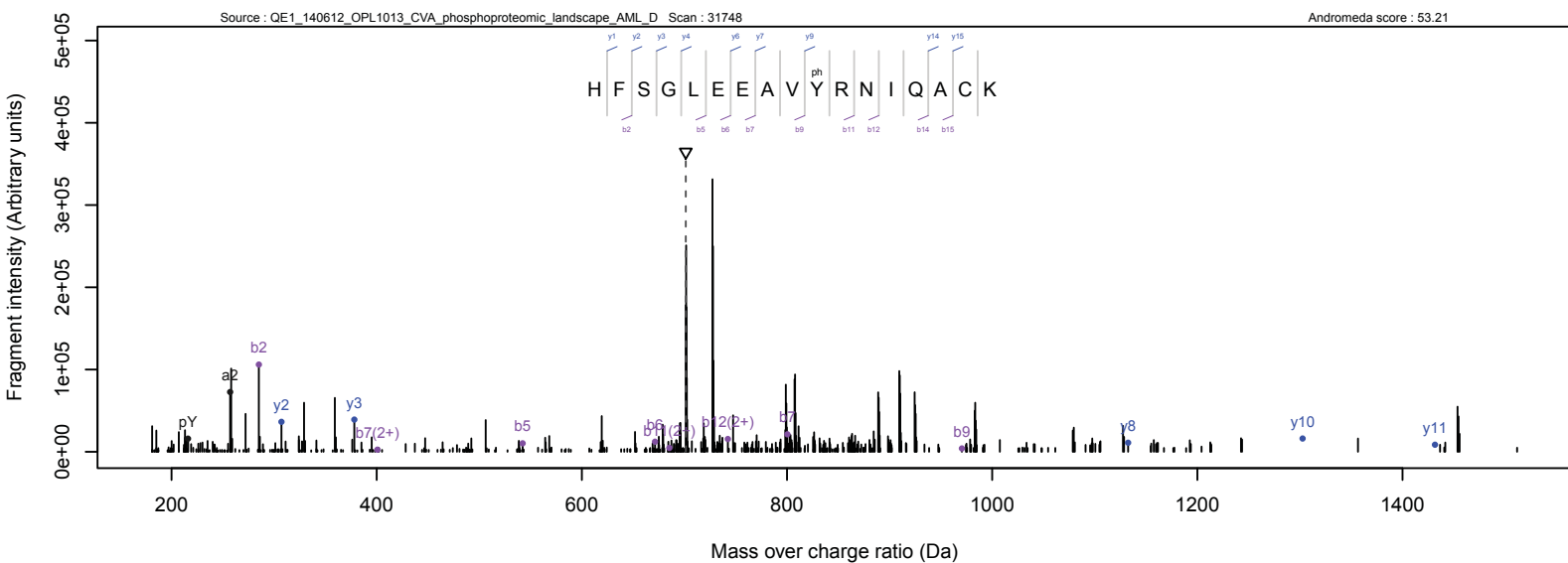




## HFSGLEEAVY(ph)RNIQACK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 31748

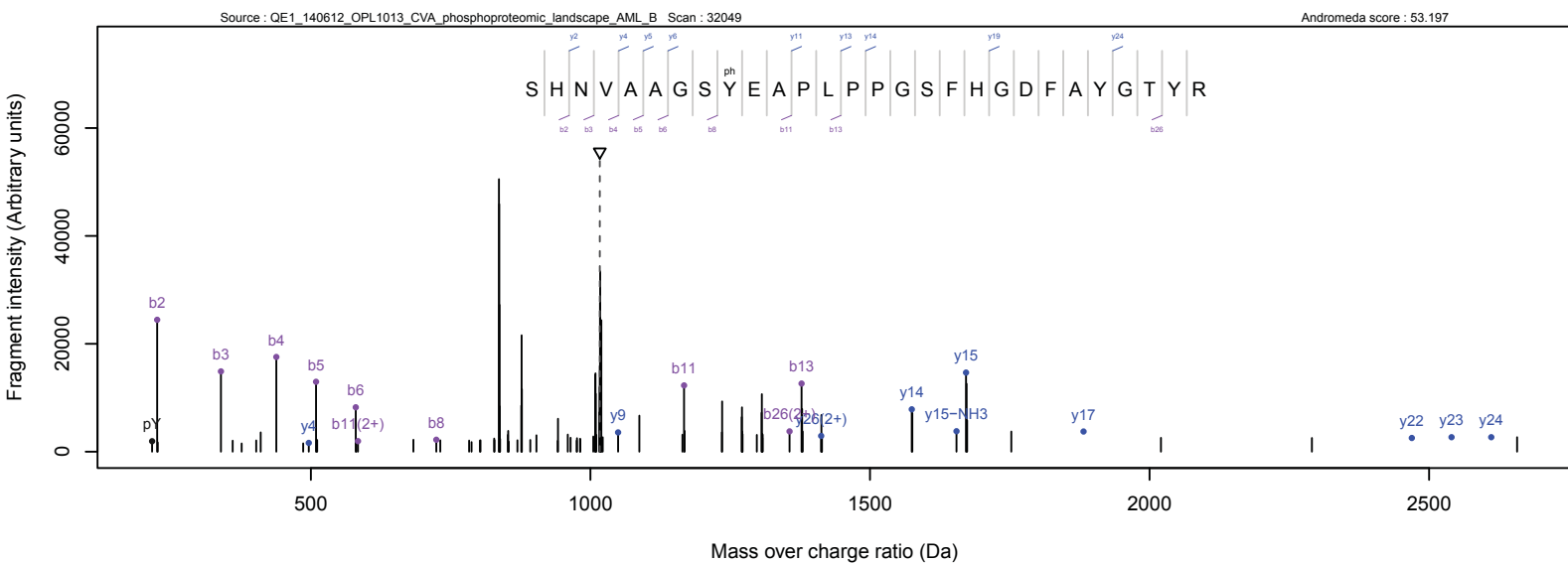
Andromeda score: 53.21



## SHNVAAGSY(ph)EAPLPPGSFHGDFAYGTYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 32049

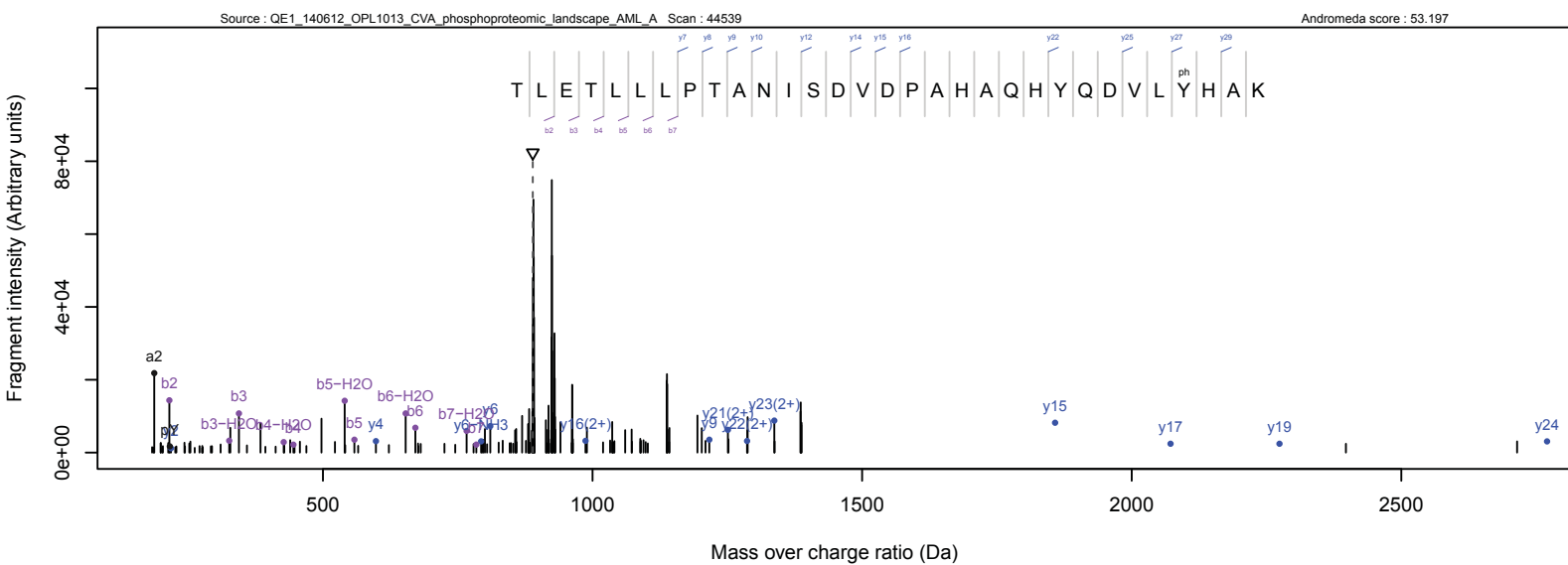
Andromeda score: 53.197



## TLETLLPTANISDVPAAHAQH YQDVLY(ph)HAK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 44539

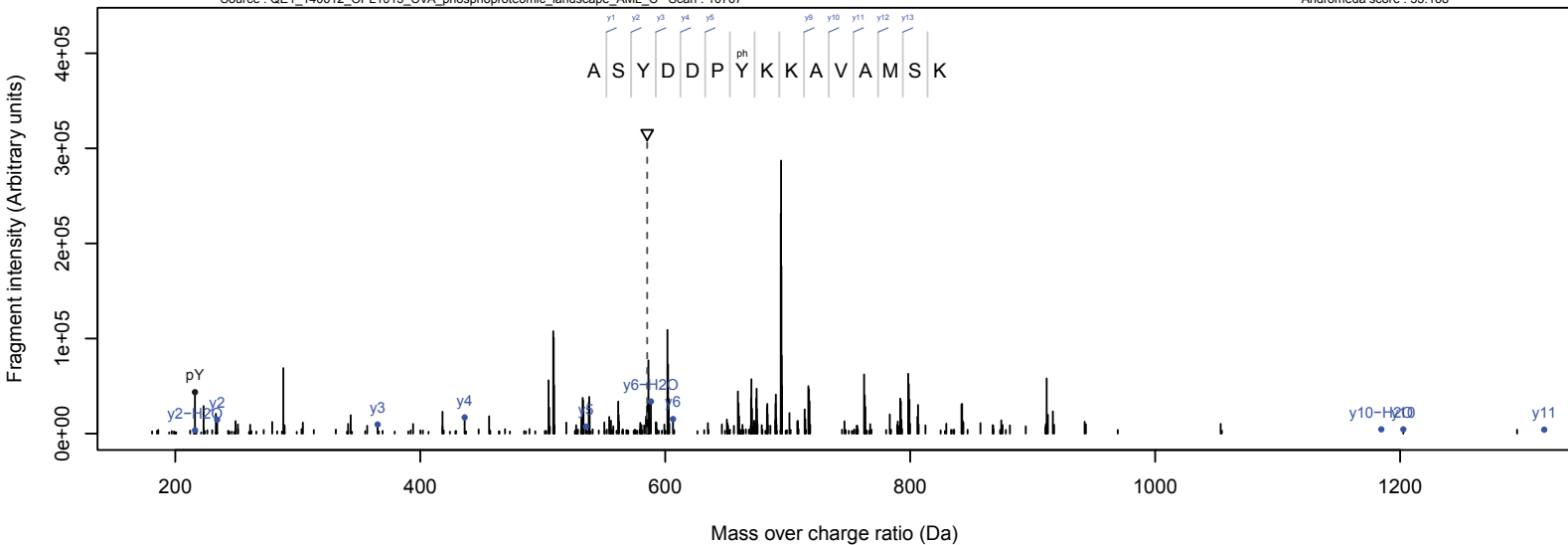
Andromeda score: 53.197



## ASYDDPY(ph)KKAVAMSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10767

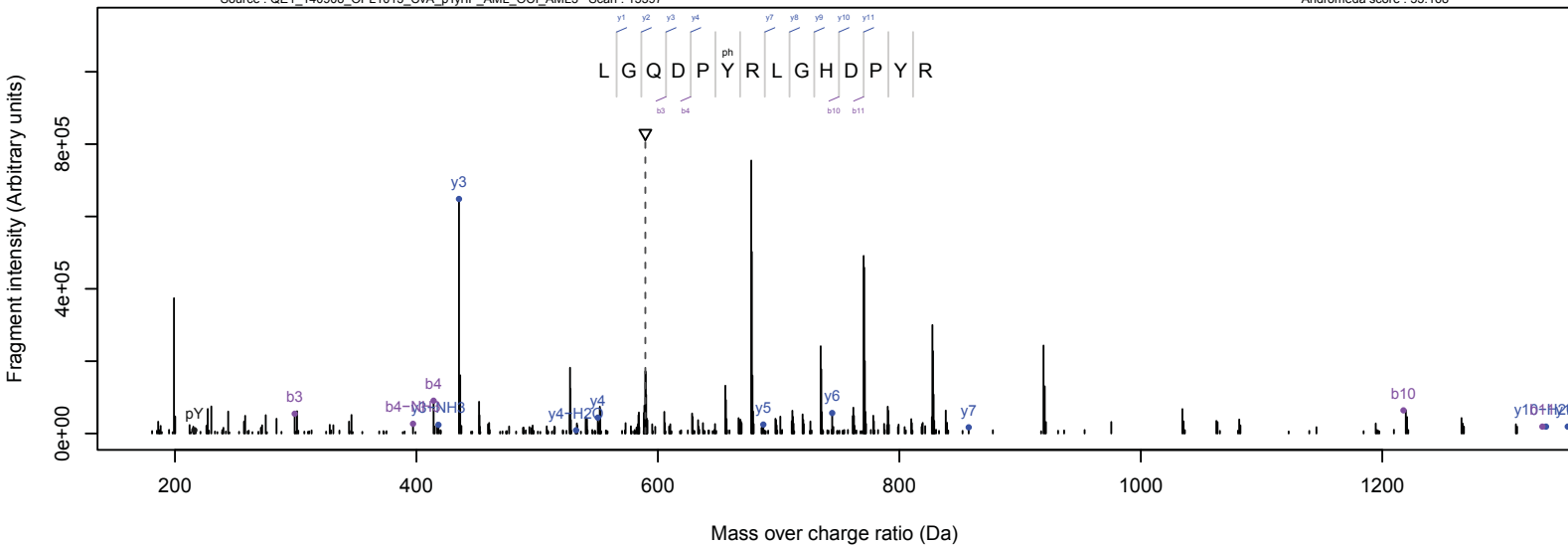
Andromeda score: 53.168



## LGQDPY(ph)RLGHDPYR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 15397

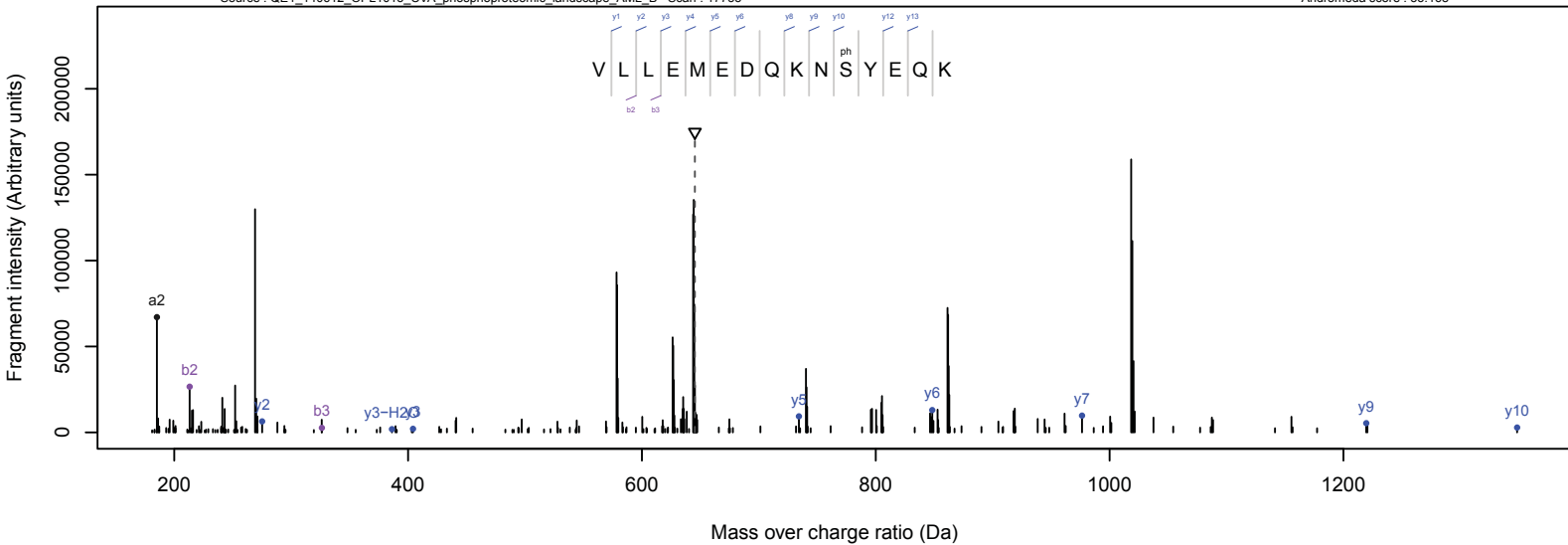
Andromeda score: 53.168



## VLEMEDQKNS(ph)YEQK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 17755

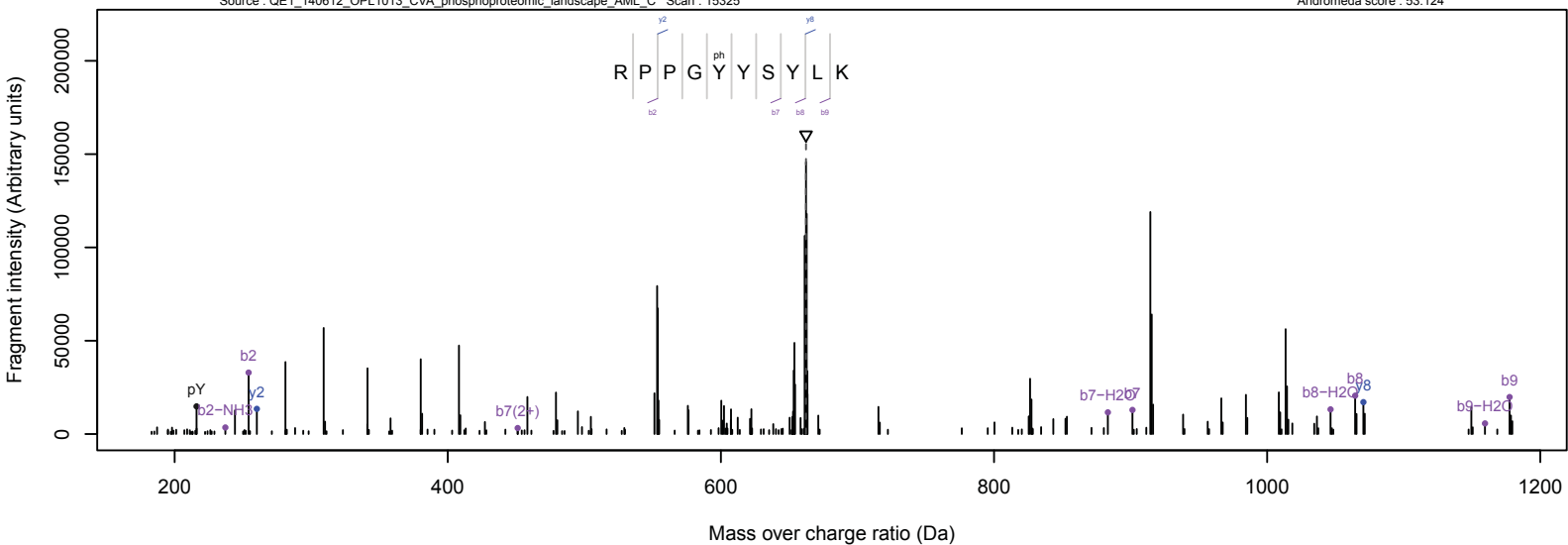
Andromeda score: 53.168



RPPGYYSYLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15325

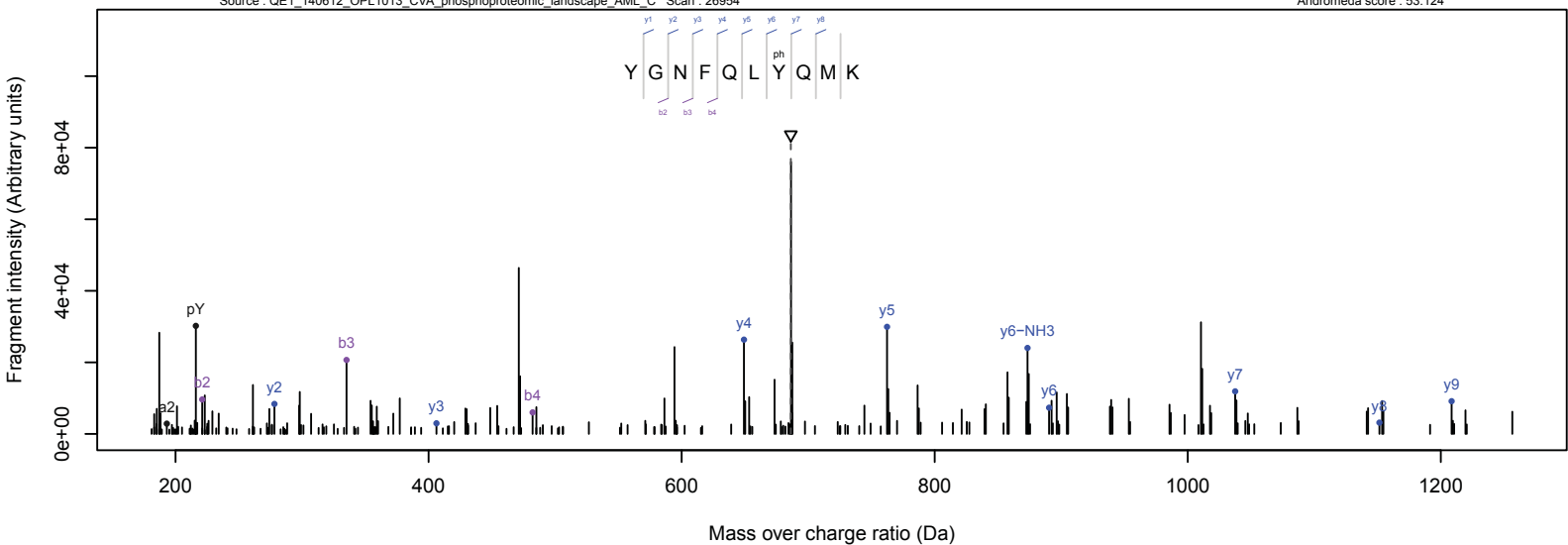
Andromeda score : 53.124



YGNFQLYQMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26954

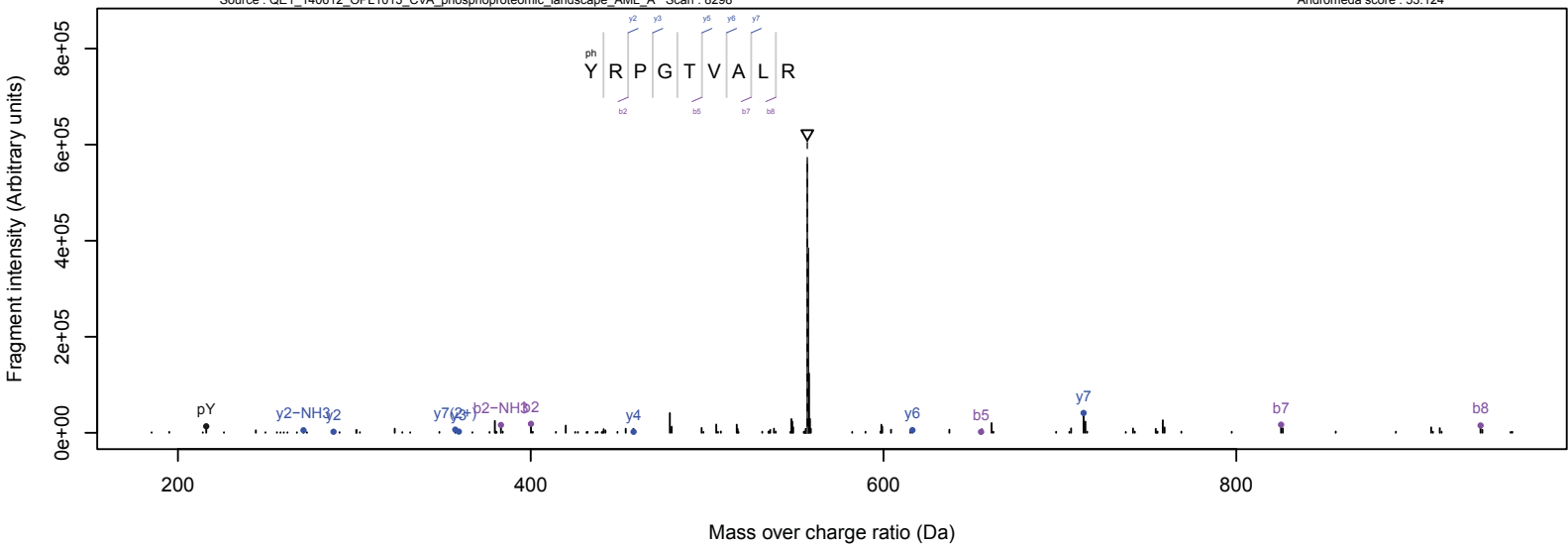
Andromeda score : 53.124



Y(ph)RPGTVLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 8298

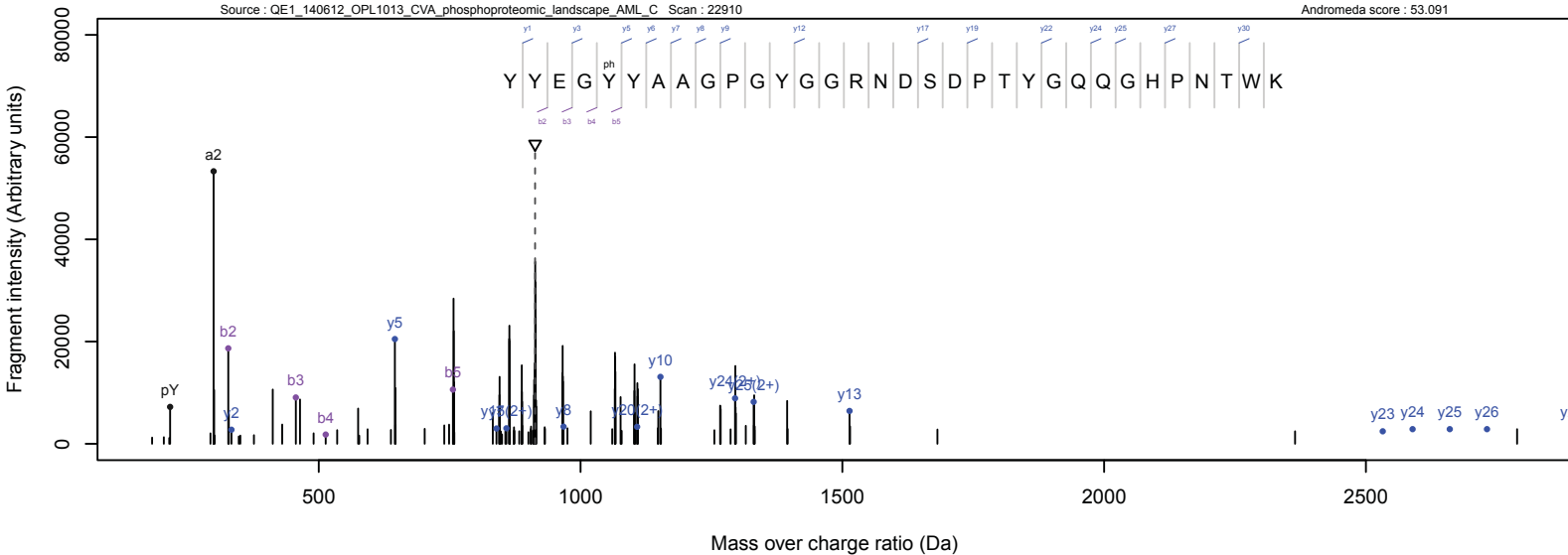
Andromeda score : 53.124



YYEGY(ph)YAAGPGYGGRRNDSPTYGQQGHPNTWK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22910

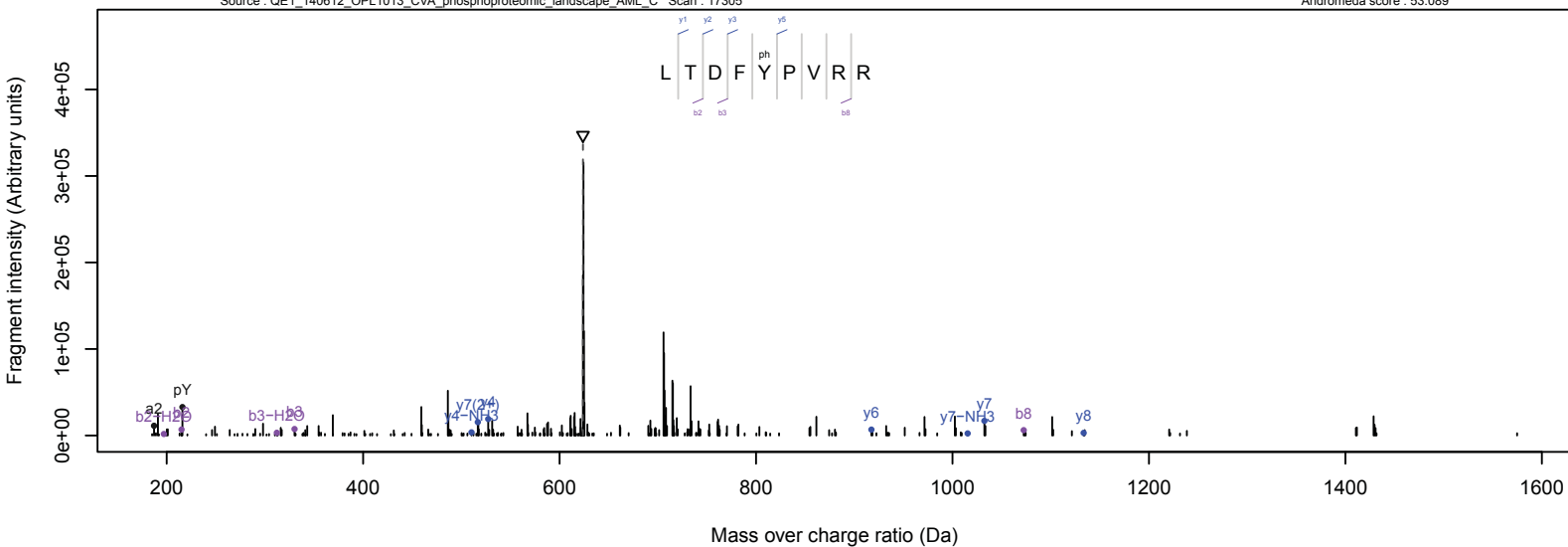
Andromeda score : 53.091



LTDIFY(ph)PVRR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17305

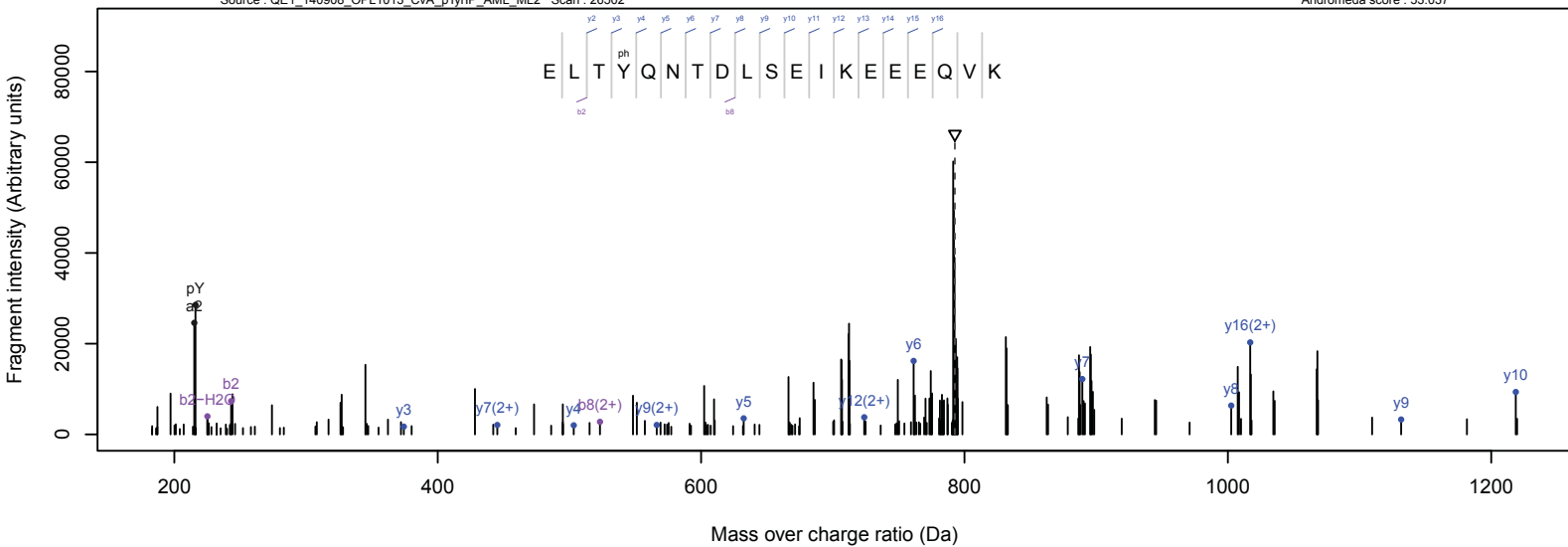
Andromeda score : 53.089



ELTY(ph)QNTDLSEIKEEEQVK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 26502

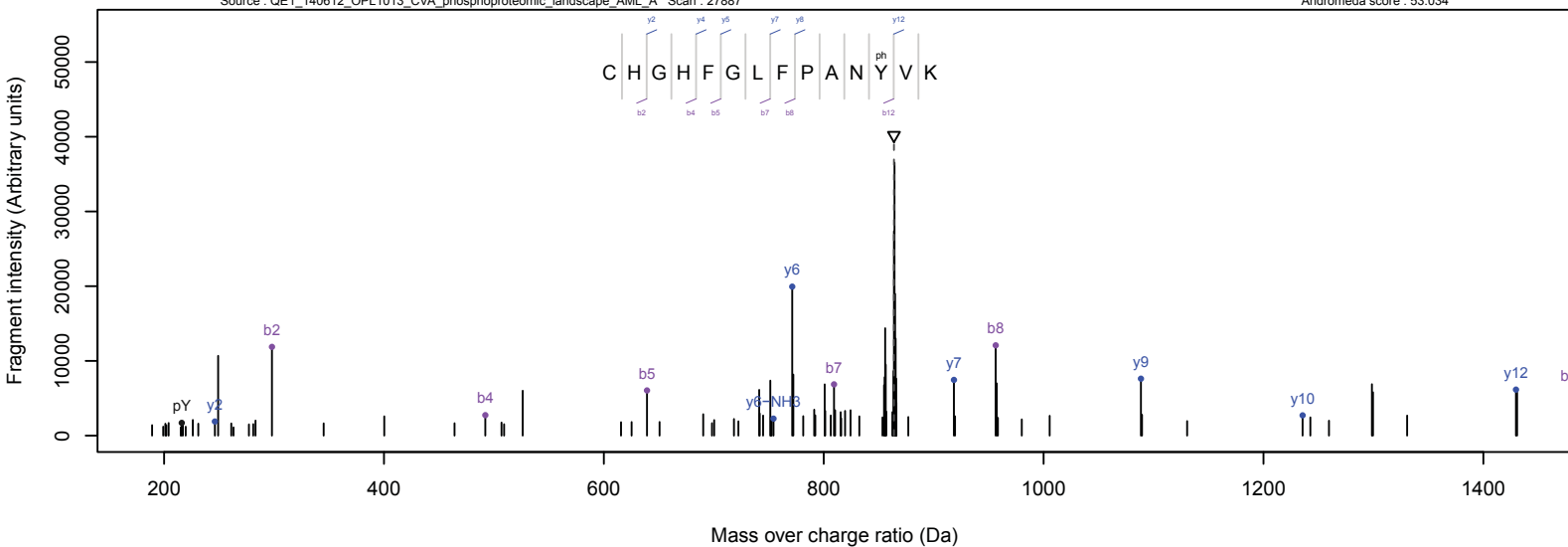
Andromeda score : 53.037



CHGHFGLFPANY(ph)VK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27887

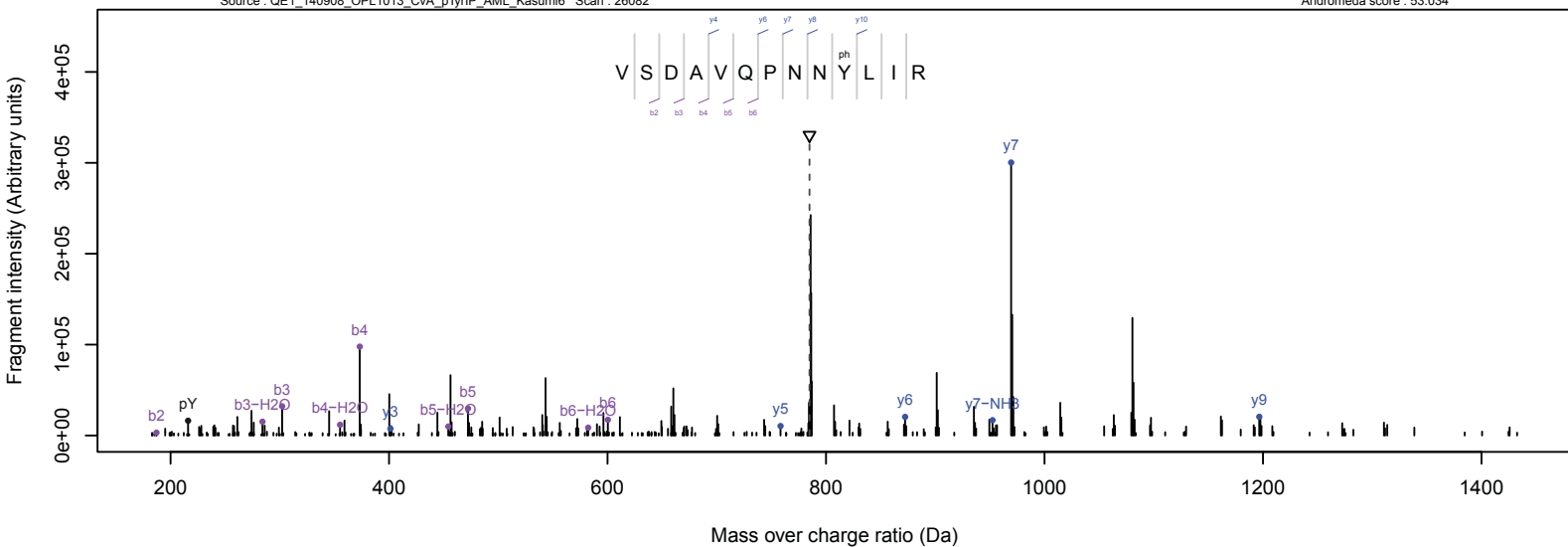
Andromeda score : 53.034



VSDAVQPNNY(ph)LIR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 26082

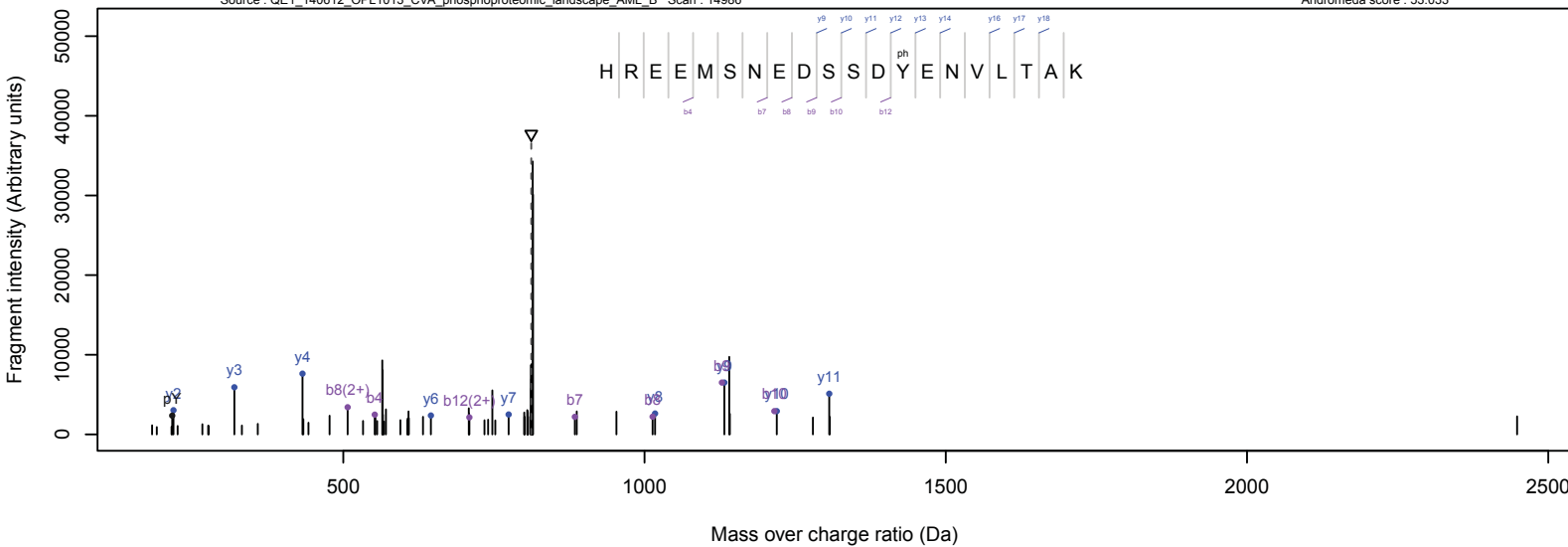
Andromeda score : 53.034



HREEMSNESSDY(ph)ENVLTAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14986

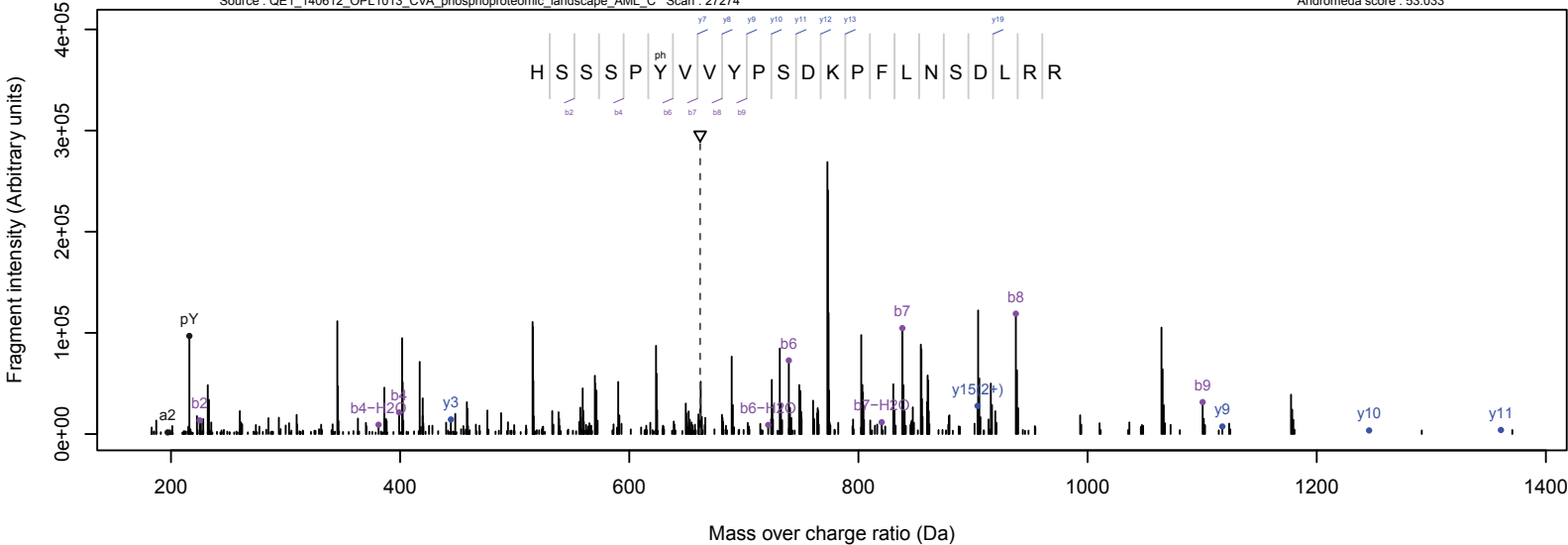
Andromeda score : 53.033



## HSSSPY(ph)VVYPSDKPFLNSDLRR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27274

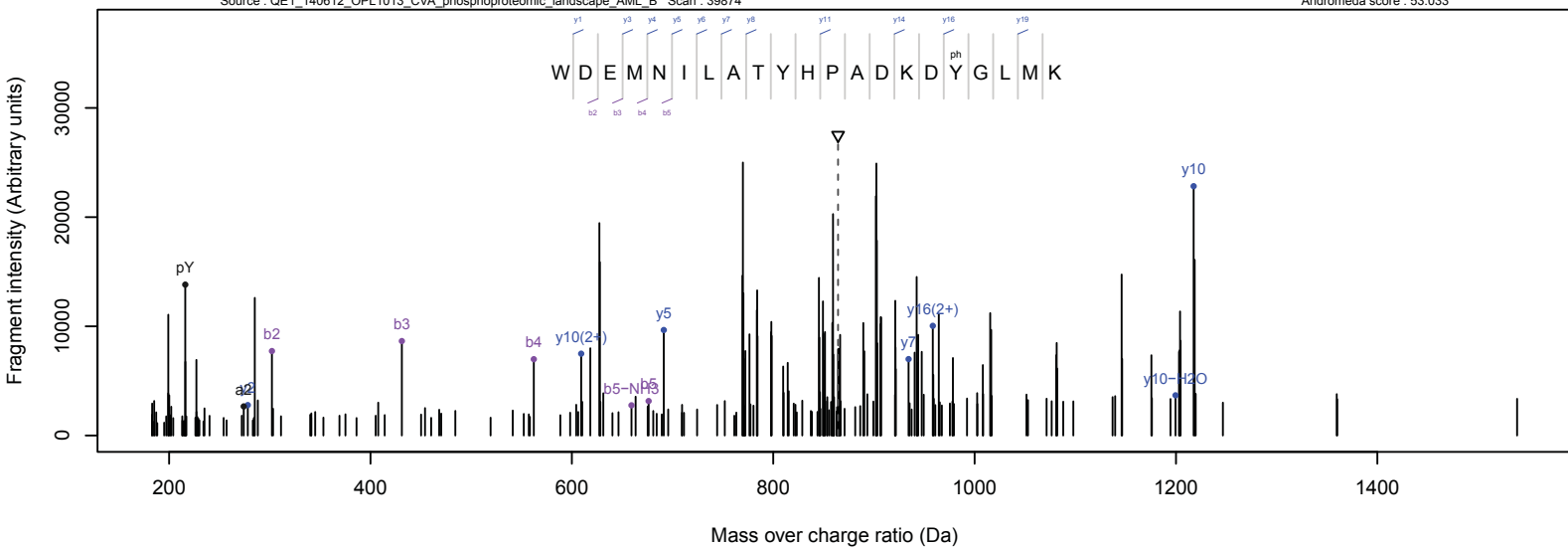
Andromeda score : 53.033



## WDEMNIATYHPADKDYGLMK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39874

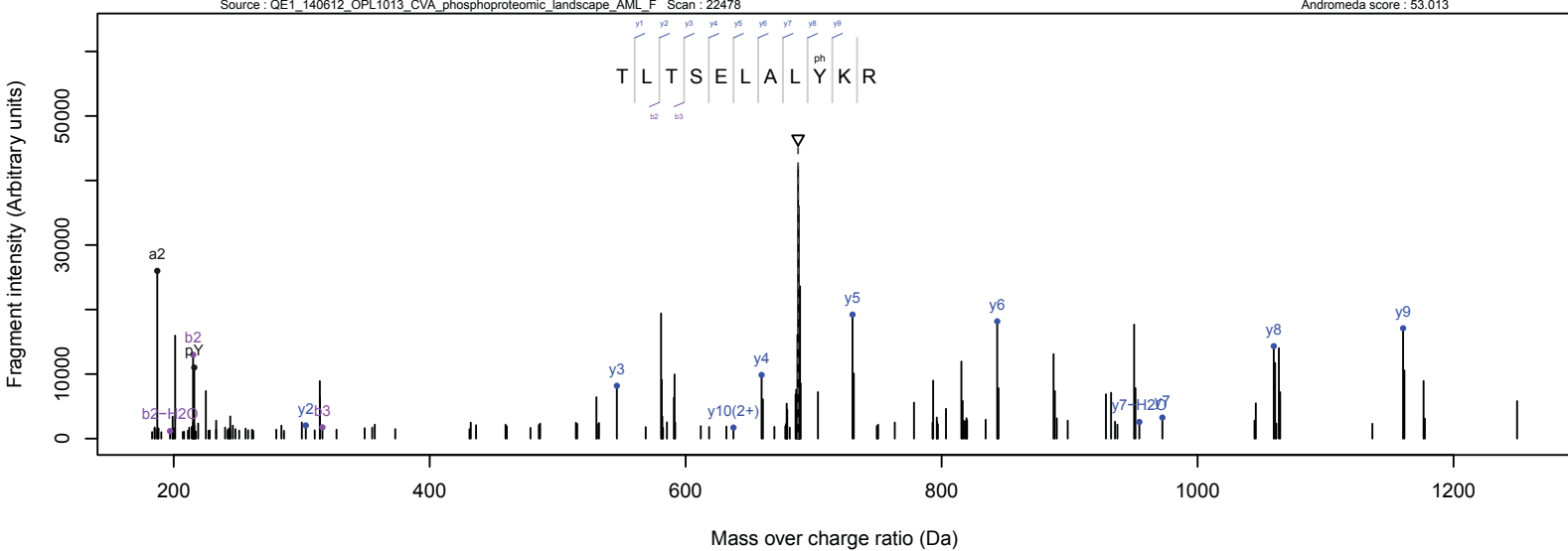
Andromeda score : 53.033



## TLTSELALYKR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 22478

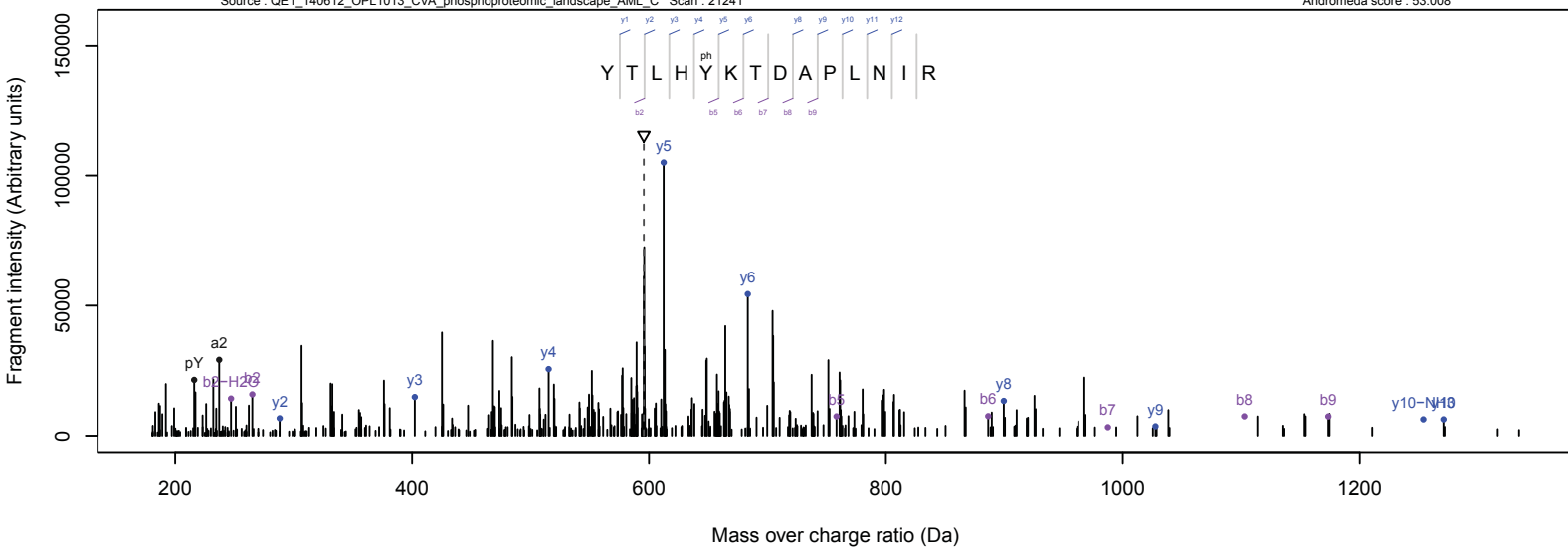
Andromeda score : 53.013



## YTLHY(ph)KTDAPLNIR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 21241

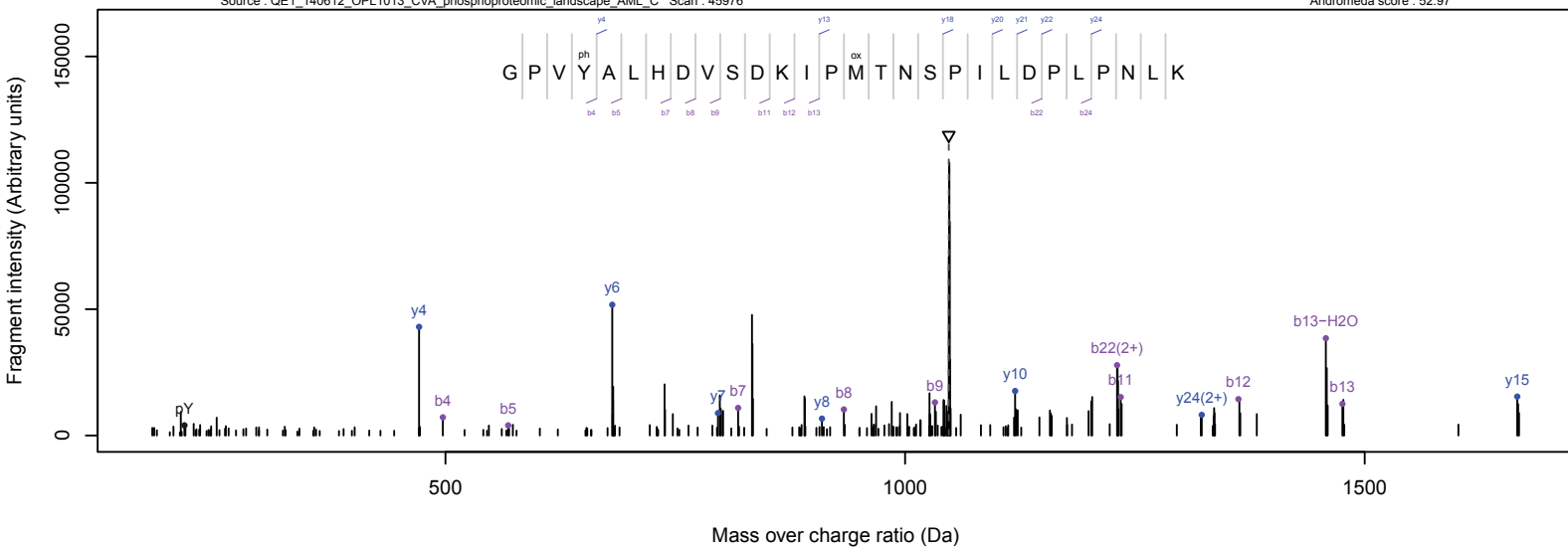
Andromeda score: 53.008



## GPVY(ph)ALHDVSDKIPMTNSPILDPLPNLK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 45976

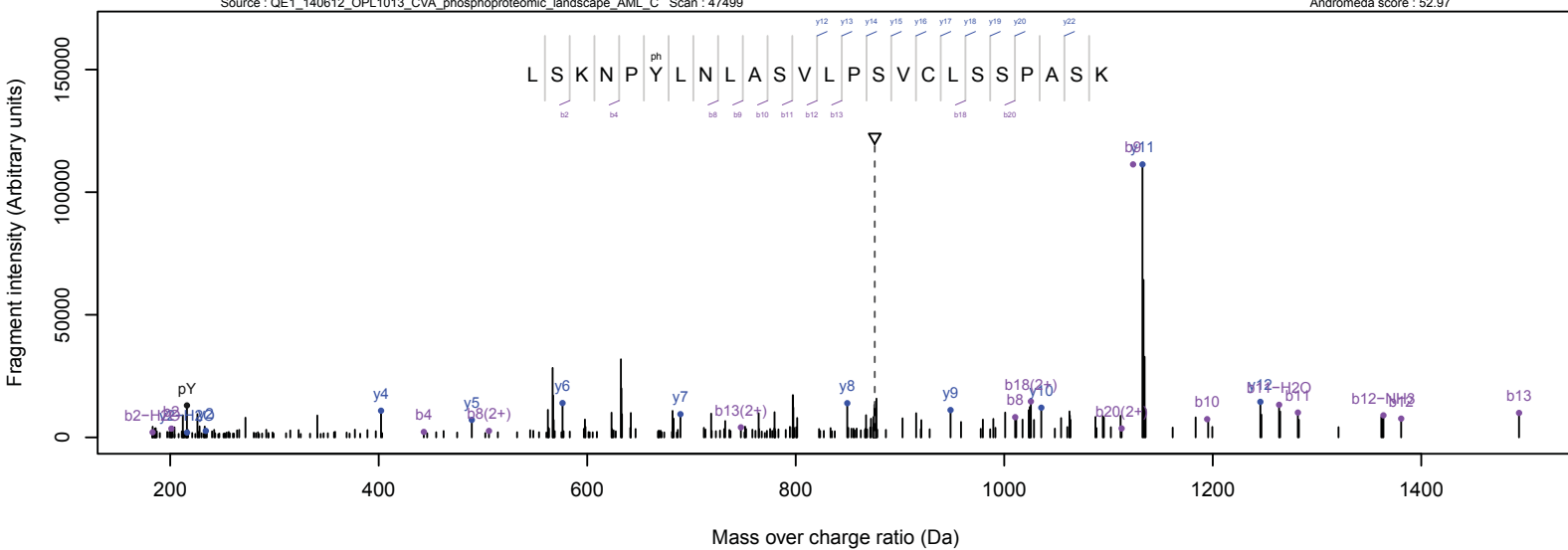
Andromeda score: 52.97



## LSKNPY(ph)LNLASVLPVCLSSPASK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 47499

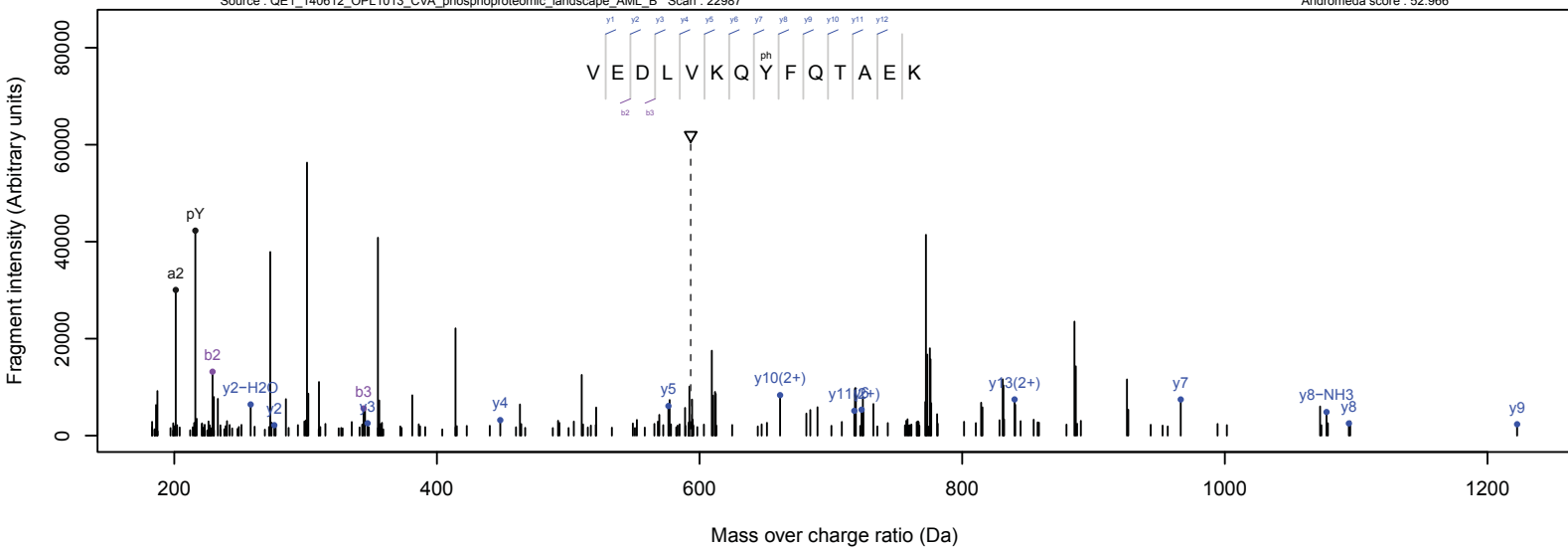
Andromeda score: 52.97



VEDLVKQY(ph)FQTAEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22987

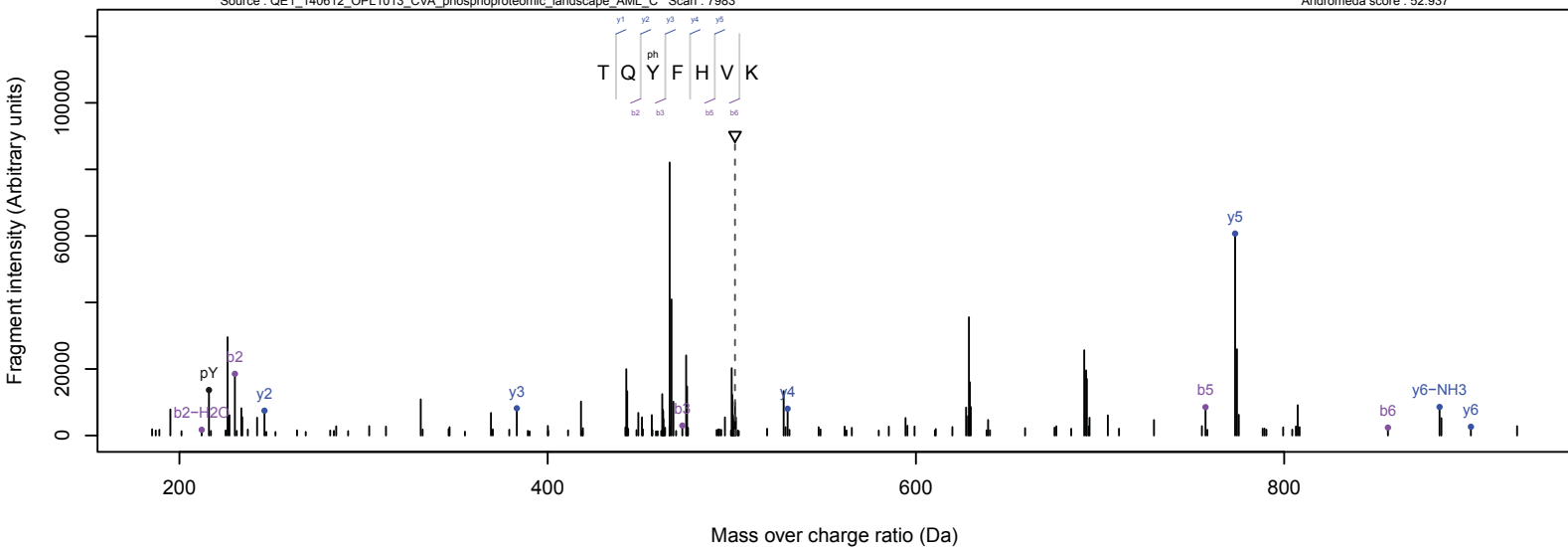
Andromeda score : 52.966



TQY(ph)FHVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7983

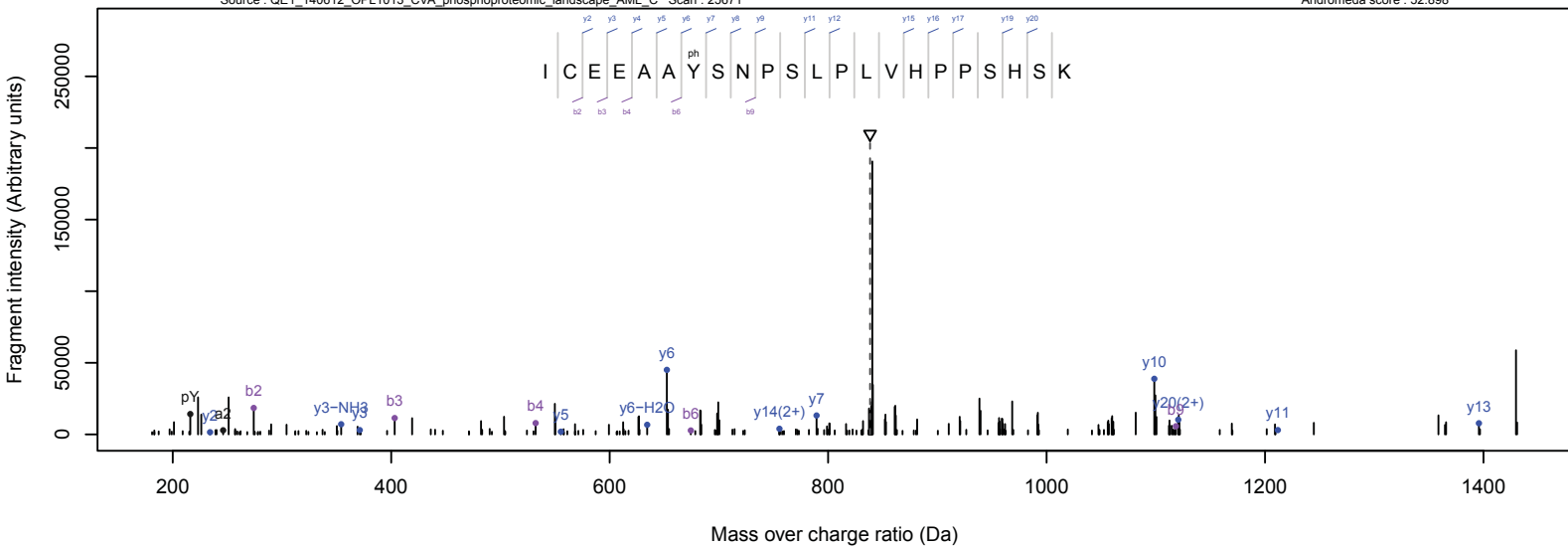
Andromeda score : 52.937



ICEEAAY(ph)SNPSLPLVHPPSHSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25671

Andromeda score : 52.898

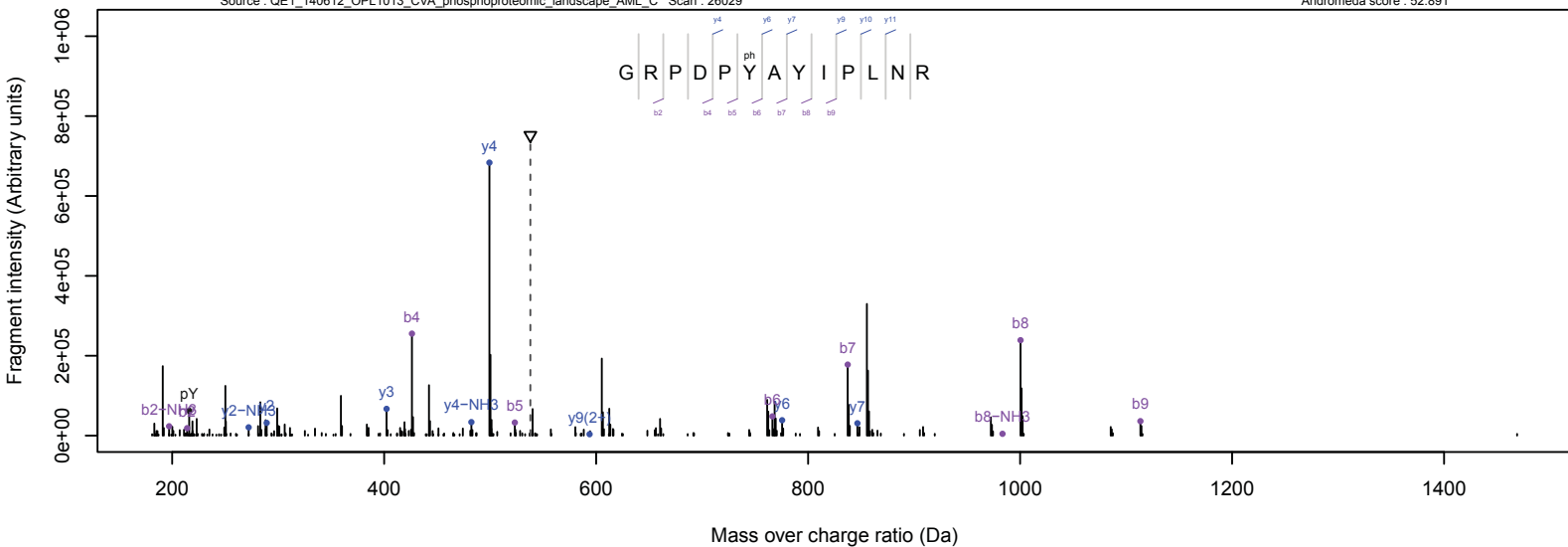




**GRDPY(ph)AYIPLNR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26029

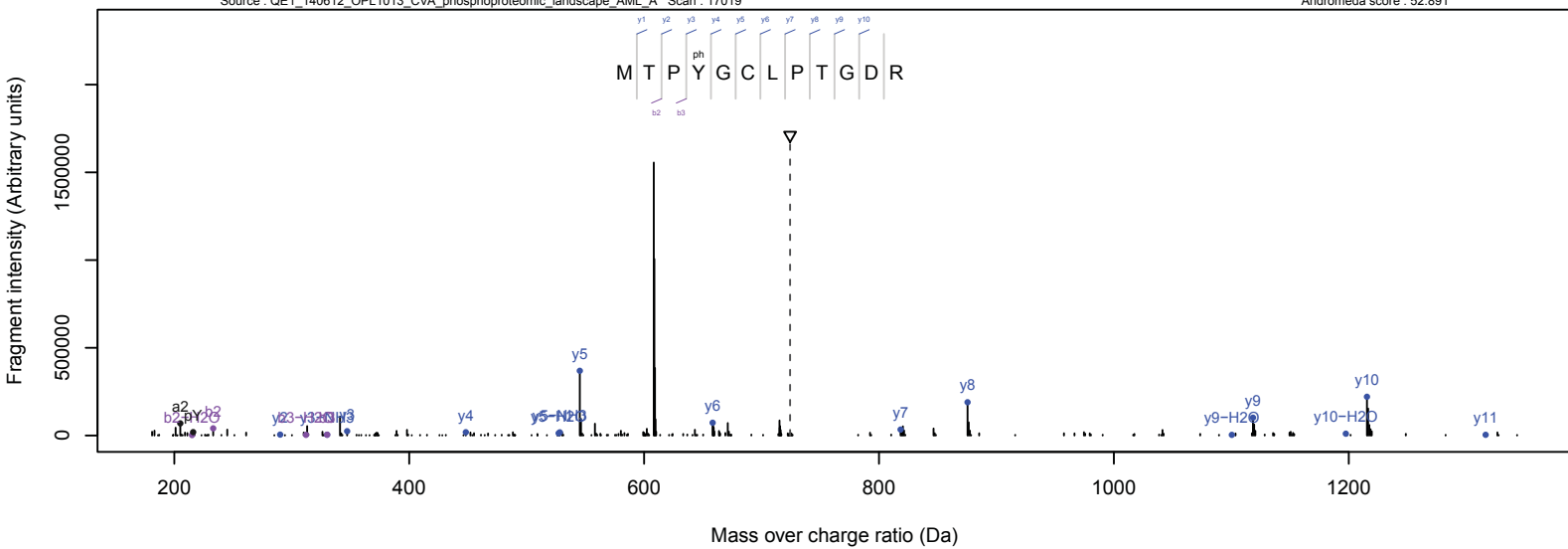
Andromeda score : 52.891



**MTPY(ph)GCLPTGDR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 17019

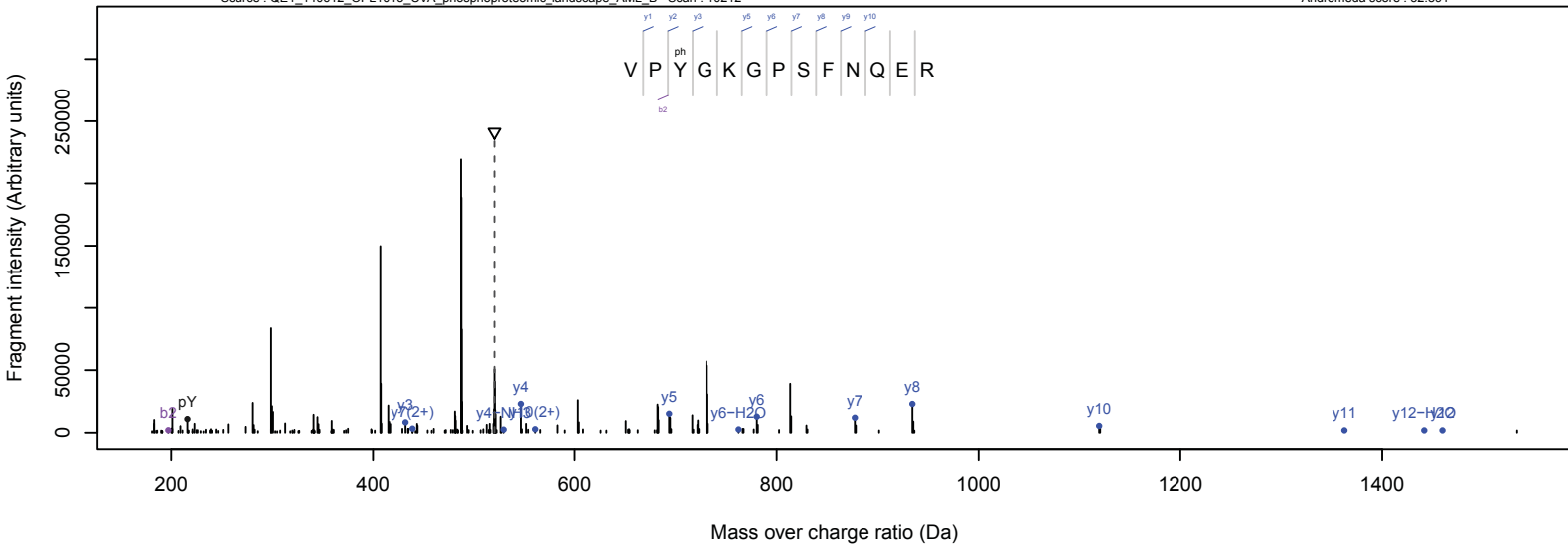
Andromeda score : 52.891



**VPY(ph)GKGPSFNQER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10212

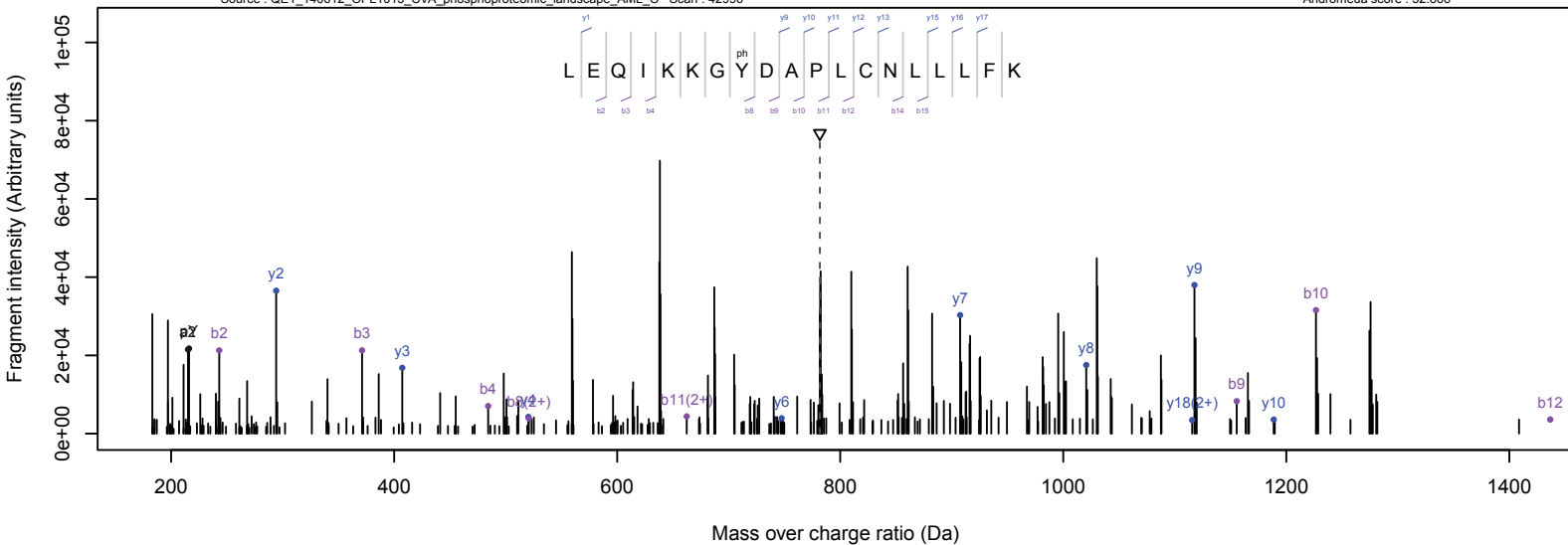
Andromeda score : 52.891



## LEQIKKGY(ph)DAPLCNLLLFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42996

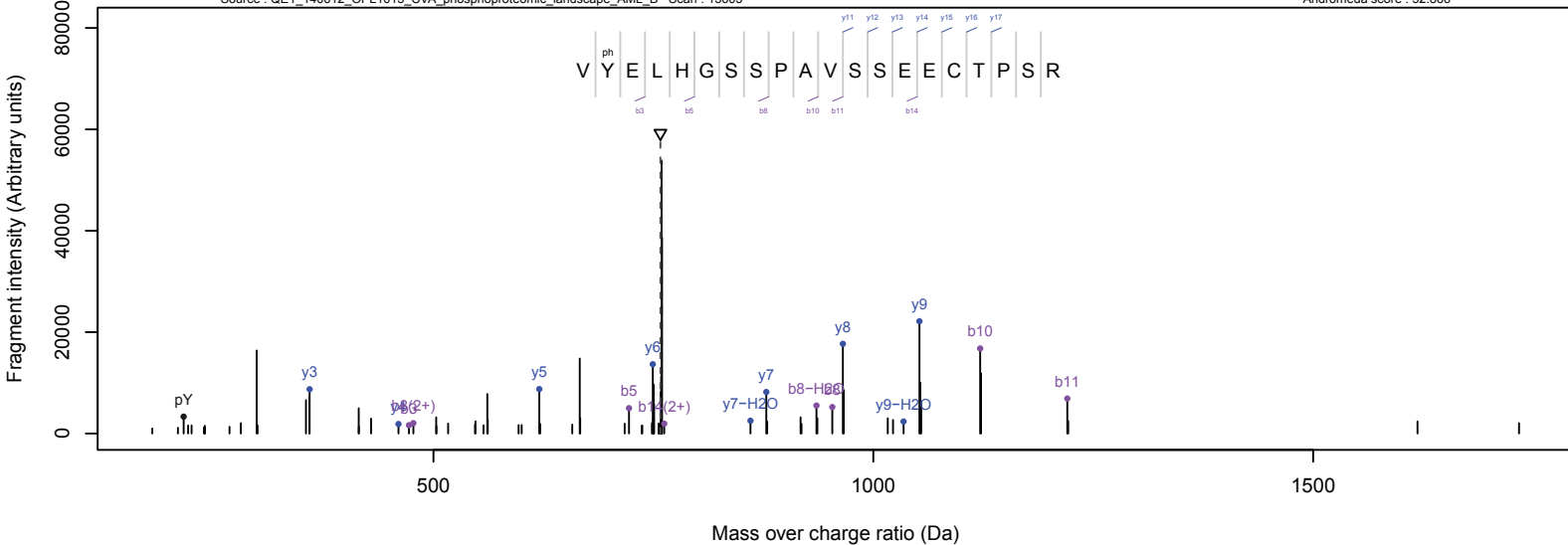
Andromeda score : 52.866



## VY(ph)ELHGSSPAVSSSEECTPSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13603

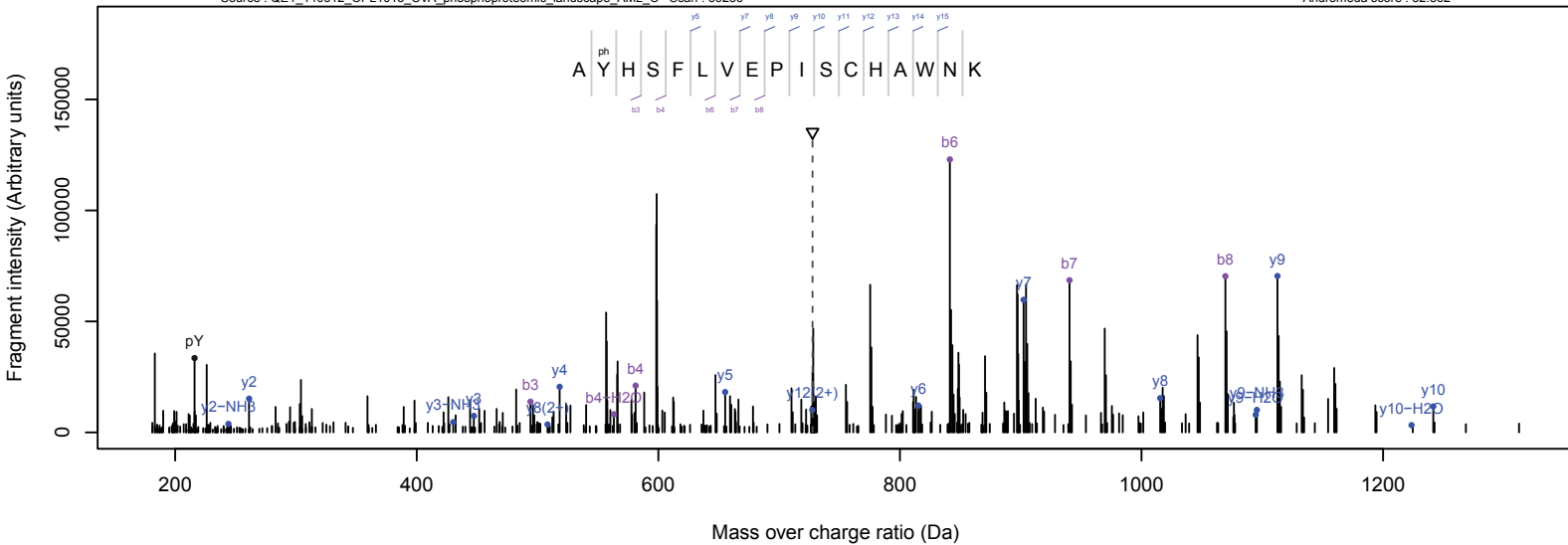
Andromeda score : 52.866



## (ac)AY(ph)HSFLVEPISCHAWNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39236

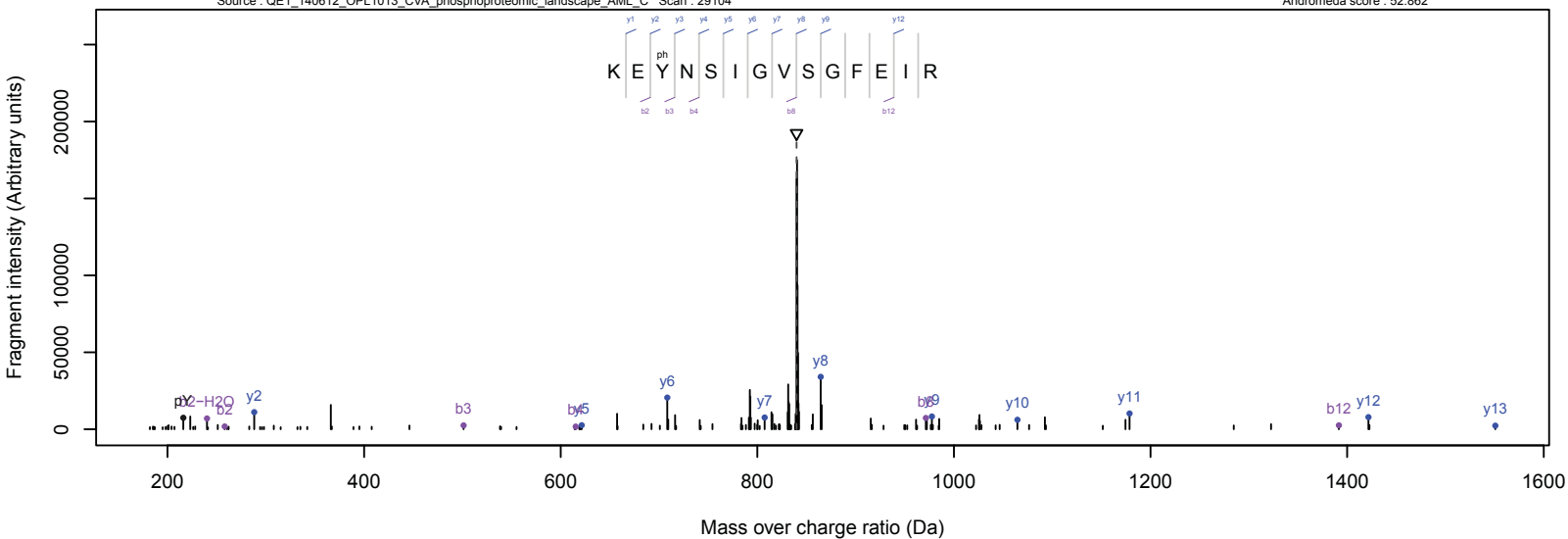
Andromeda score : 52.862



## KEY(ph)NSIGVSGFEIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29104

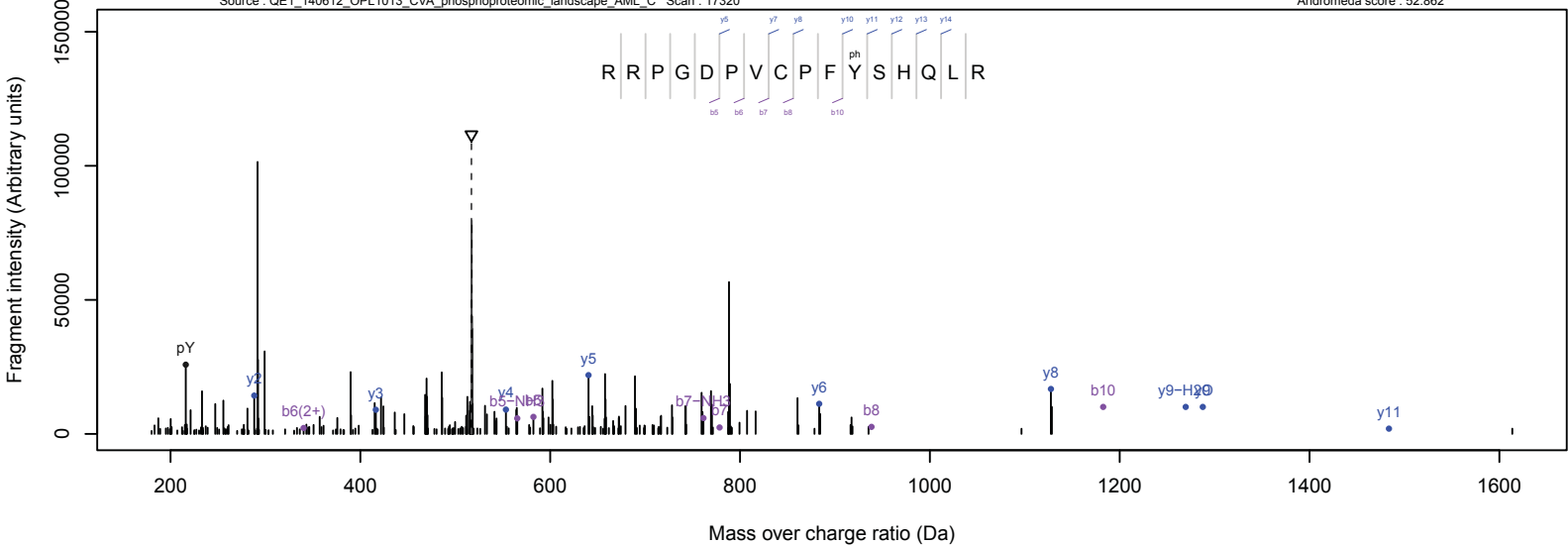
Andromeda score : 52.862



## RRPGDPVCPF(ph)SHQLR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17320

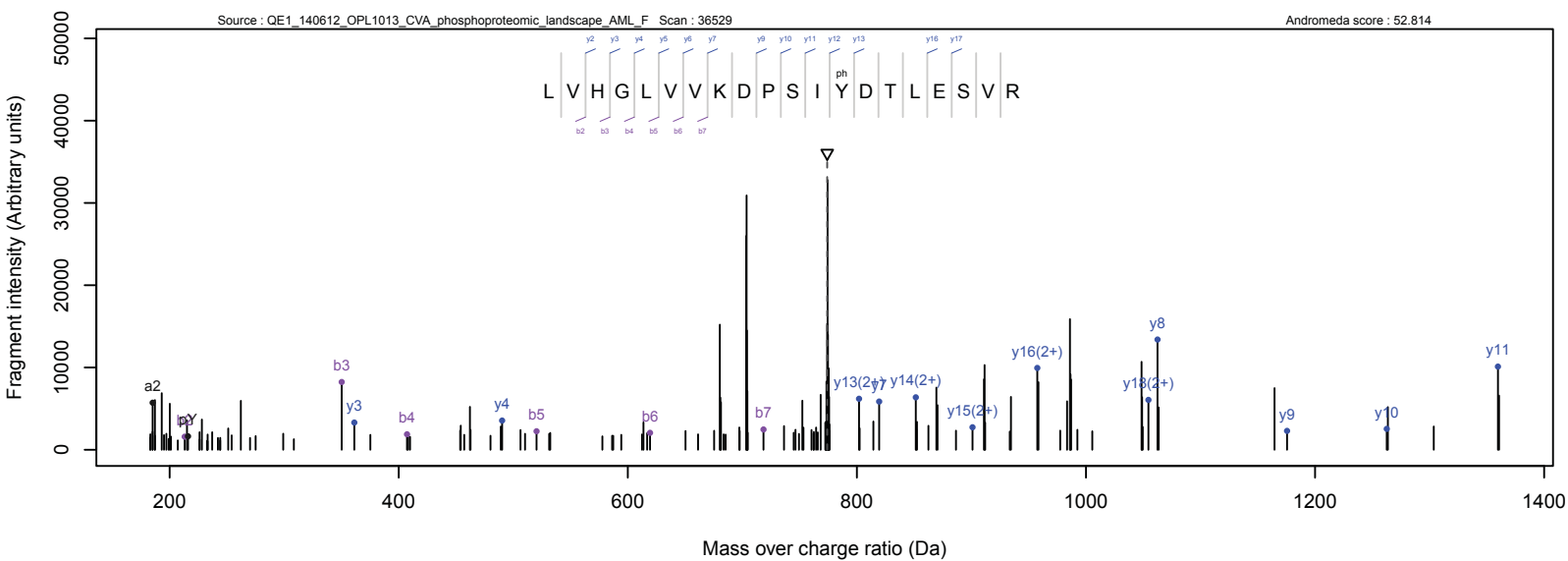
Andromeda score : 52.862



## LVHGLVVKDPSIYDTLESVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 36529

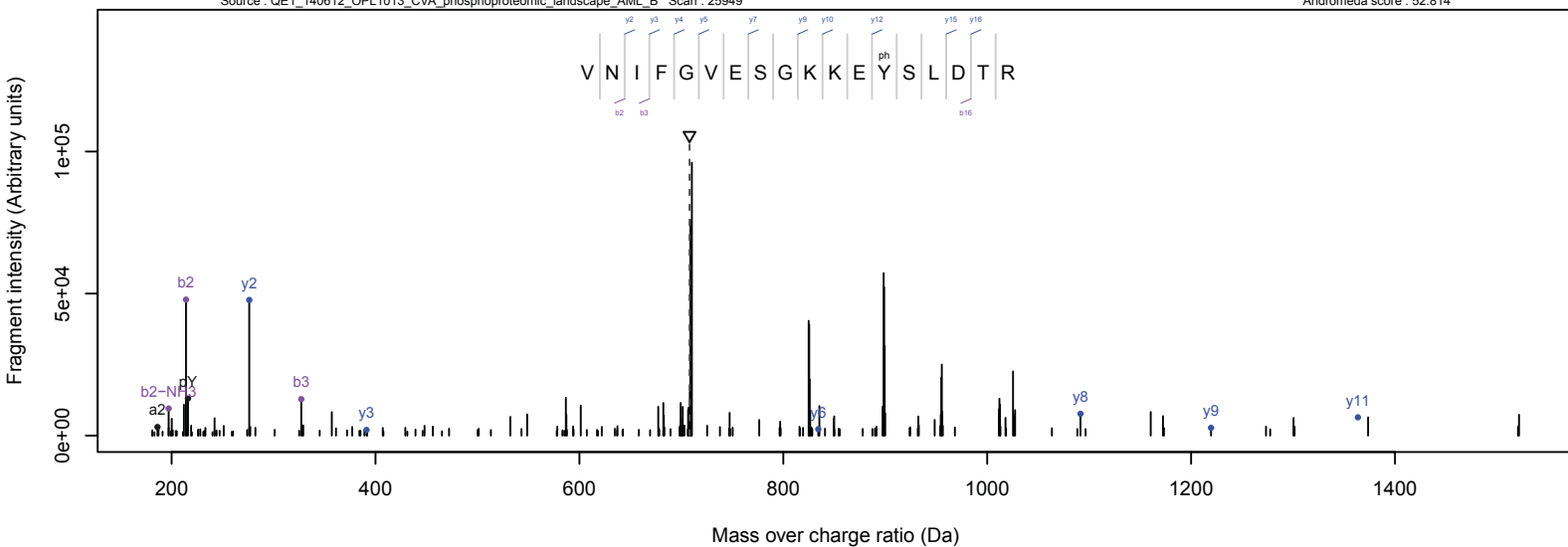
Andromeda score : 52.814



## VNIFGVESGKKEY(ph)SLDTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25949

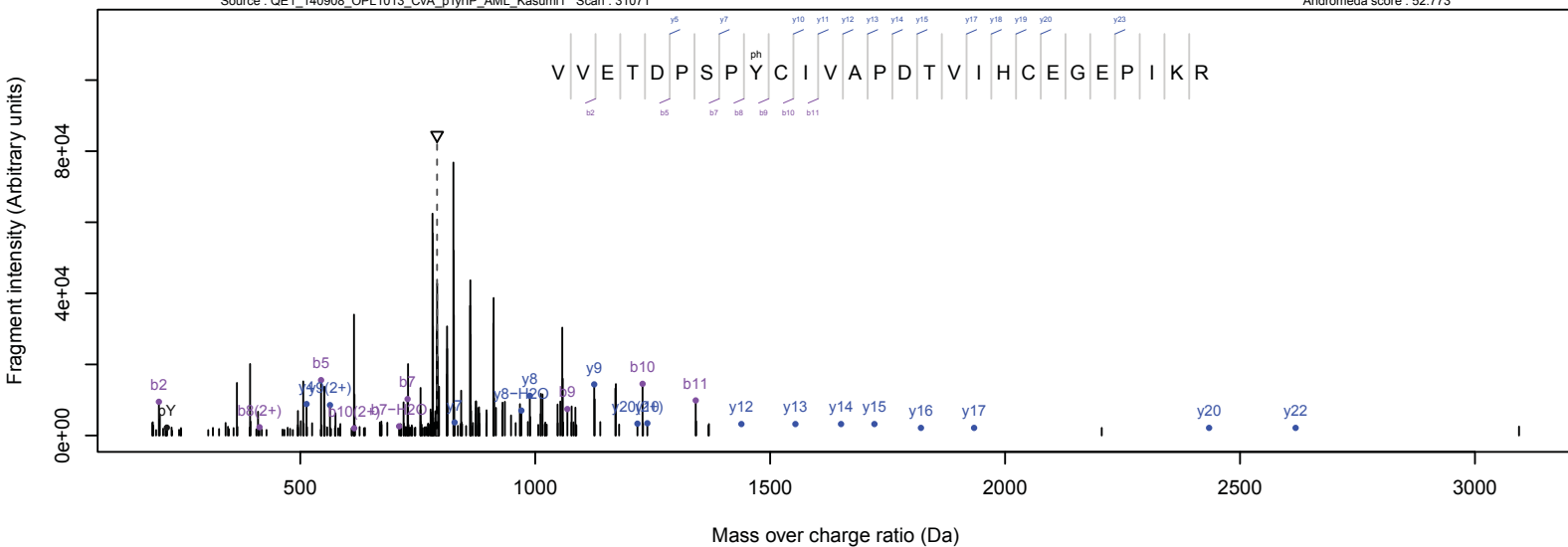
Andromeda score : 52.814



## VVETDPSY(ph)CIVAPDTVIHCEGEPIKR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 31071

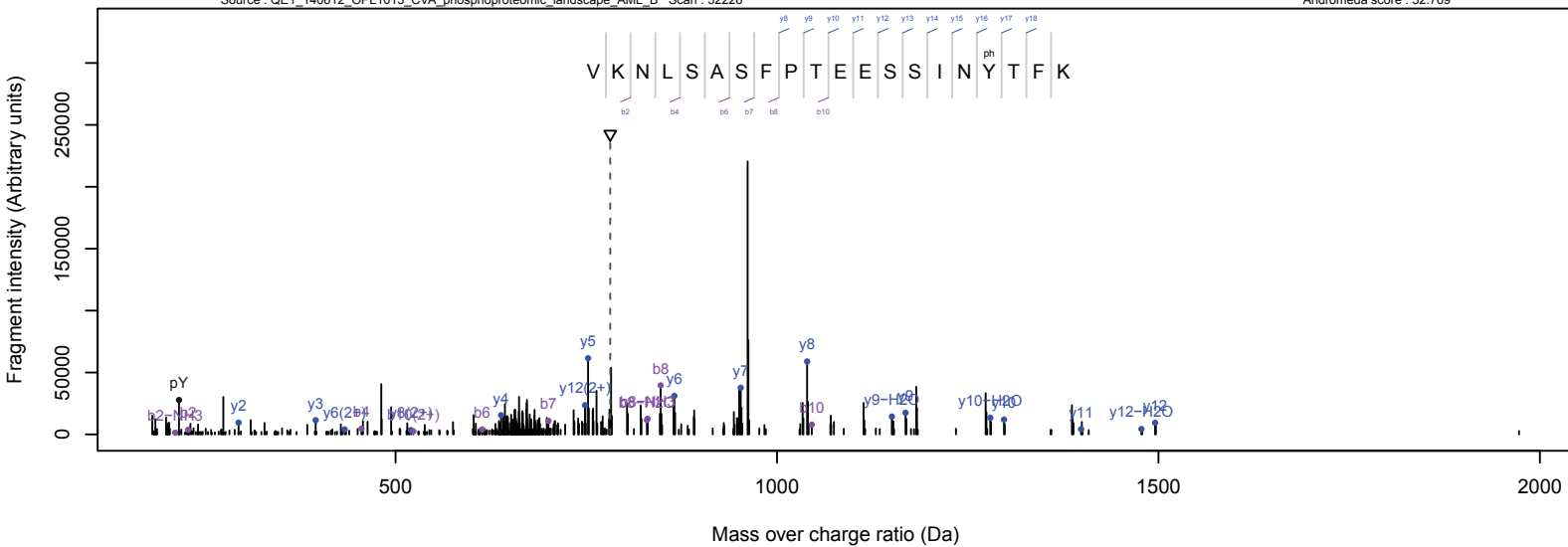
Andromeda score : 52.773



## VKNLSASFPTTEESSINY(ph)TFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32226

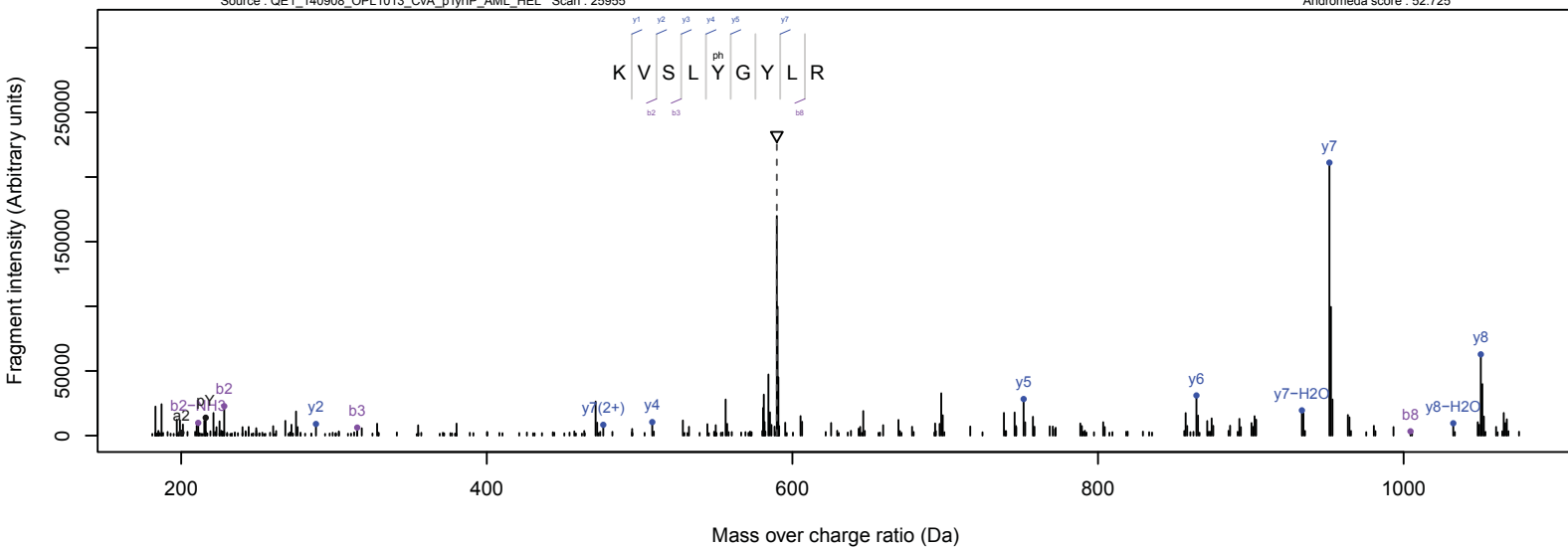
Andromeda score : 52.769



## KVSLY(ph)GYLR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 25955

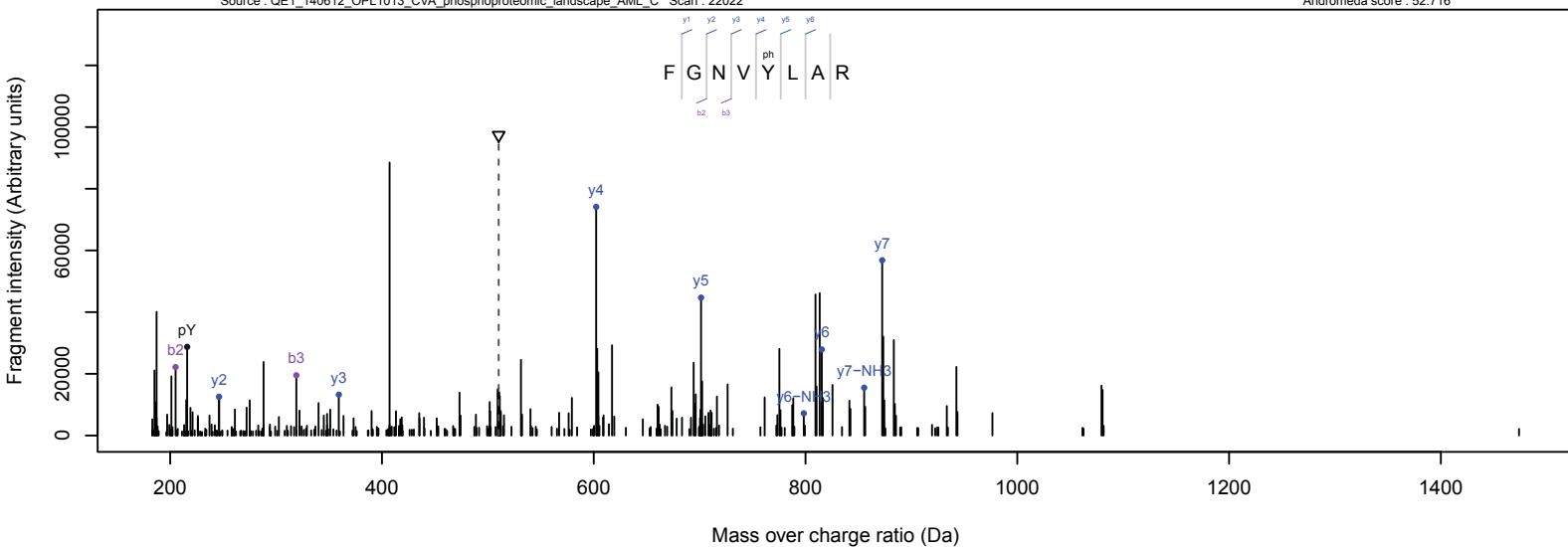
Andromeda score: 52.725



## FGNVY(ph)LAR-2+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 22022

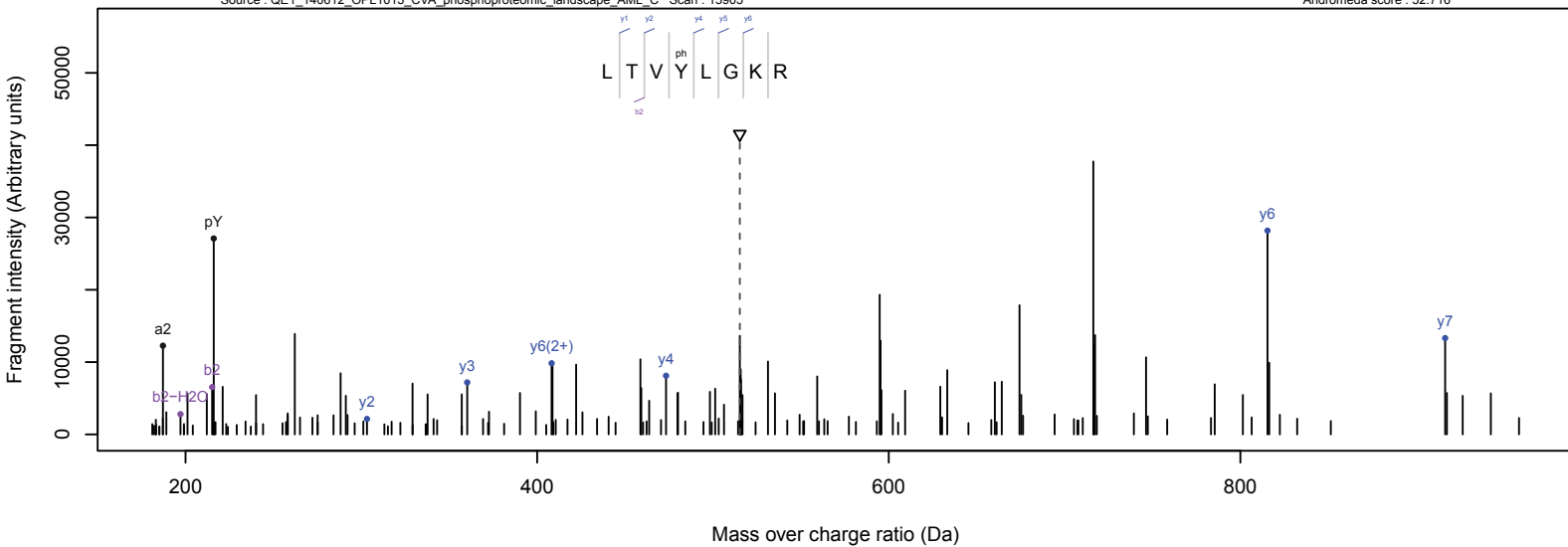
Andromeda score: 52.716



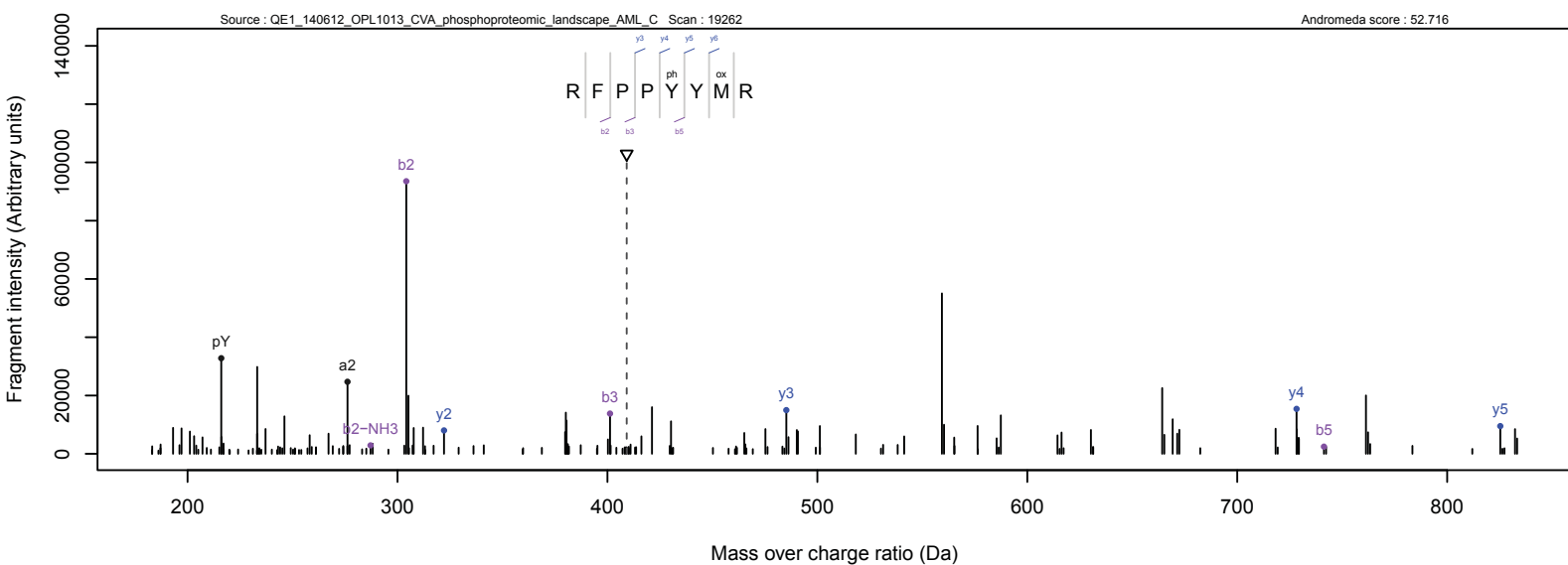
## LTVY(ph)LGKR-2+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 15905

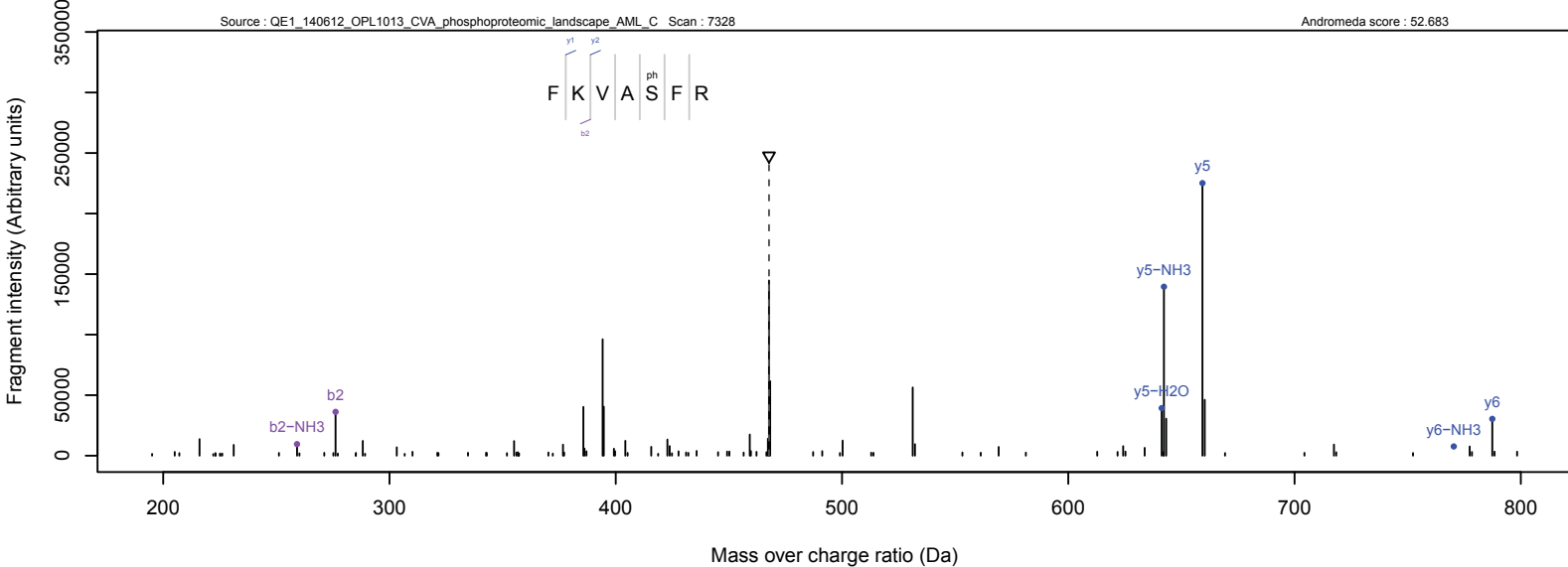
Andromeda score: 52.716



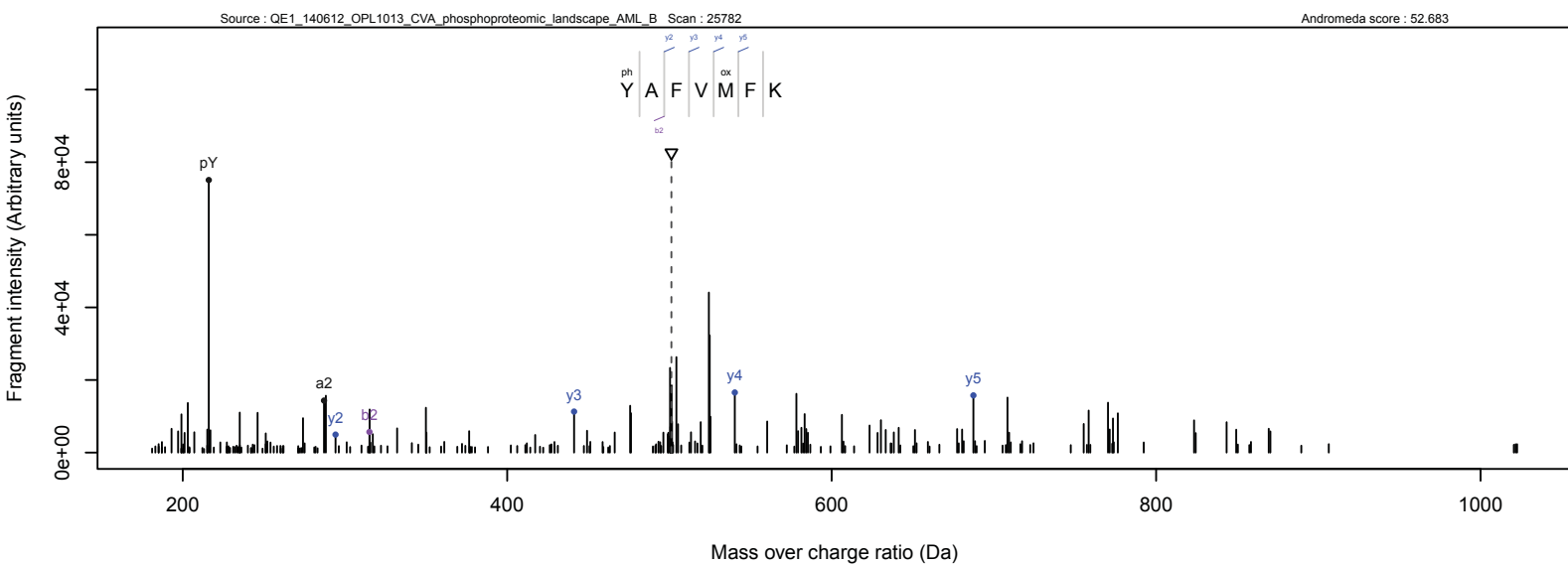
## RFPPY(ph)YM(ox)R-3+



## FKVAS(ph)FR-2+



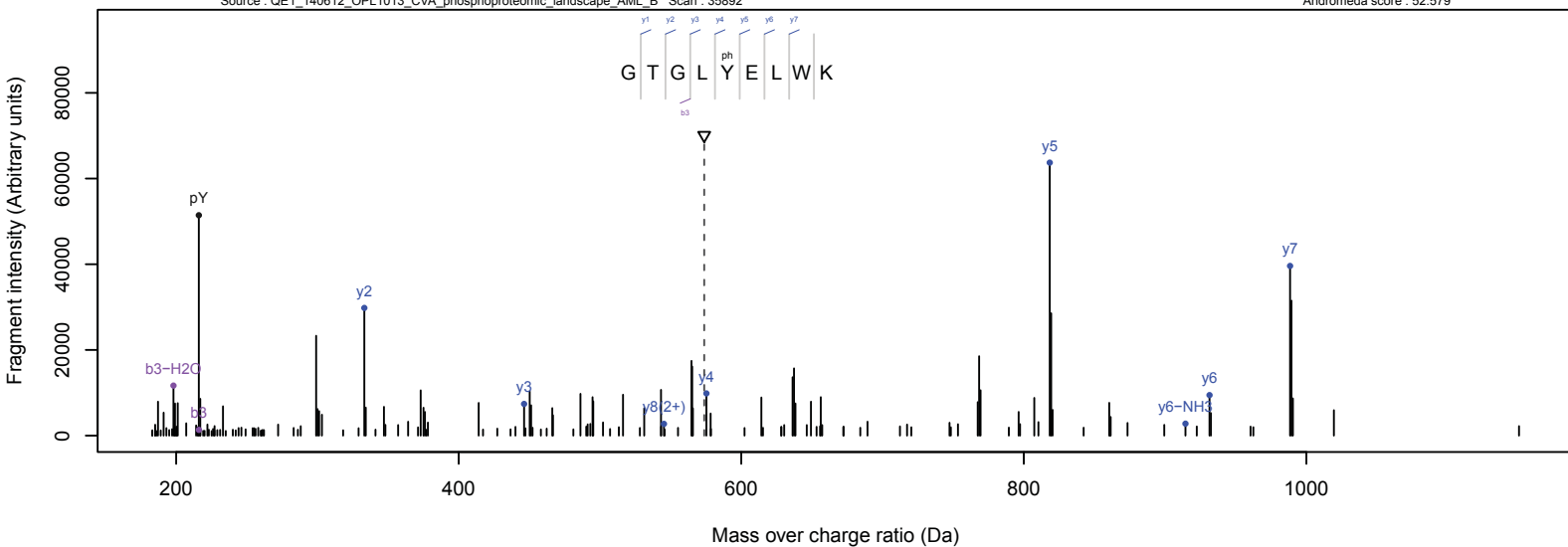
## Y(ph)AFVM(ox)FK-2+



## GTGLY(ph)ELWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 35892

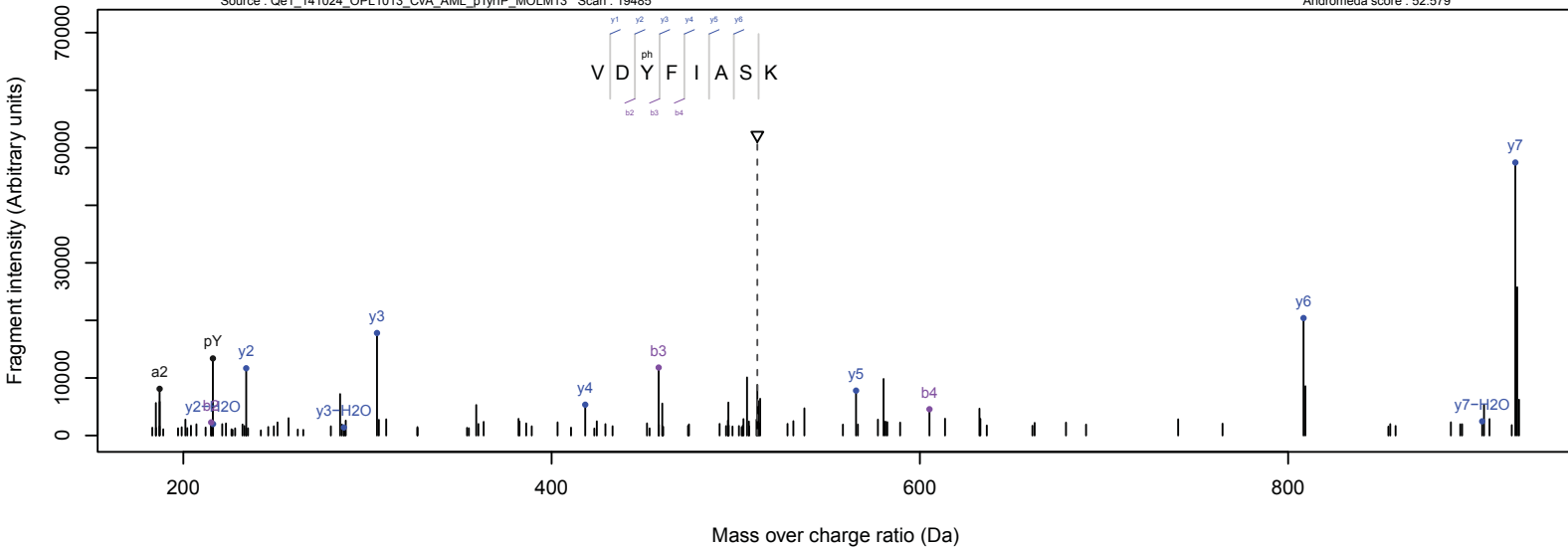
Andromeda score : 52.579



## VDY(ph)FIASK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 19485

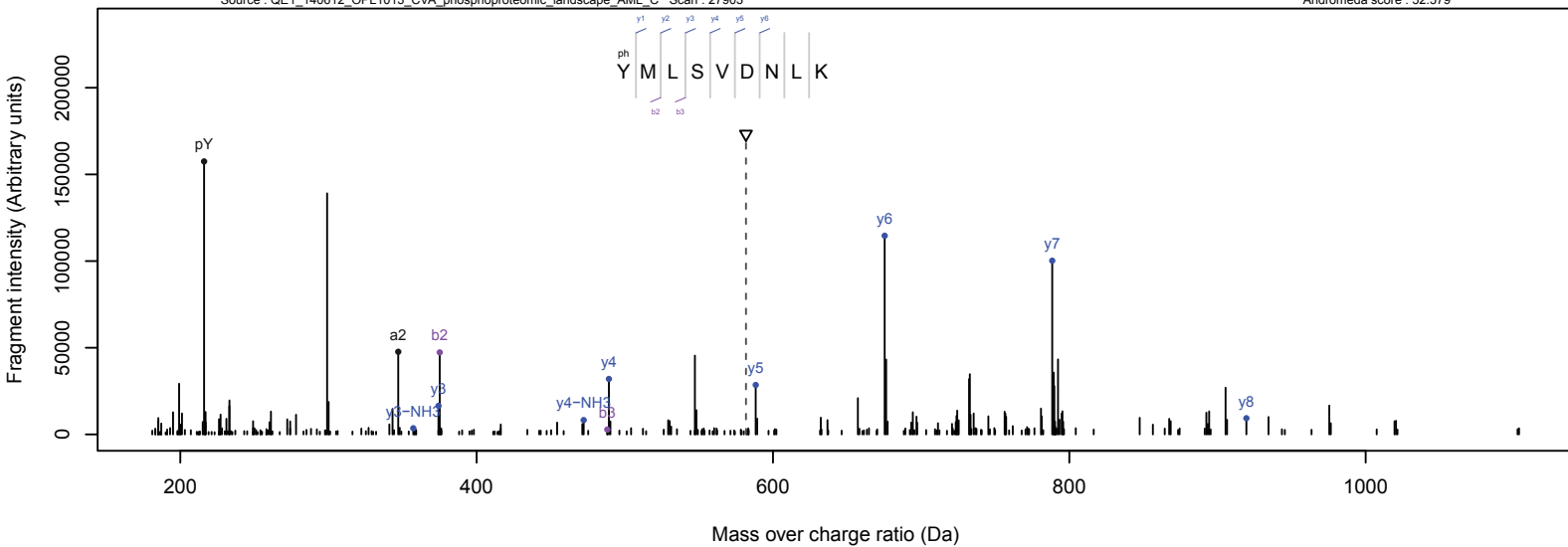
Andromeda score : 52.579



## Y(ph)MLSVDNLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27903

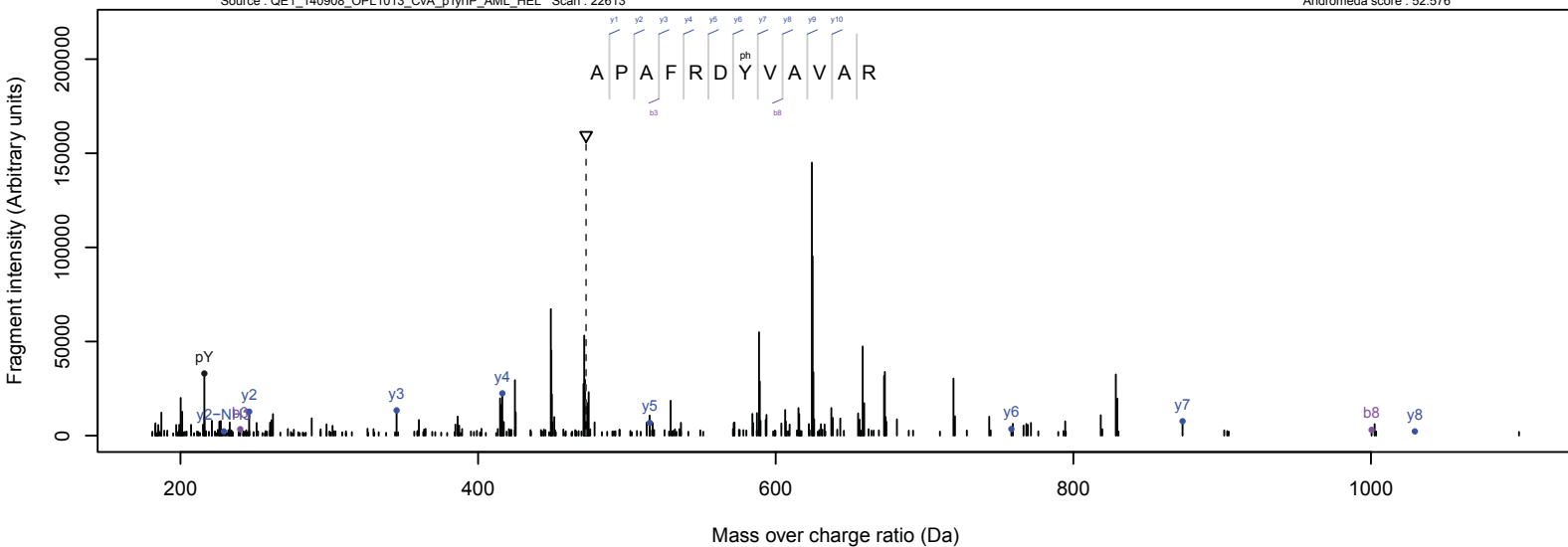
Andromeda score : 52.579



## APAFRDY(ph)VAVAR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 22613

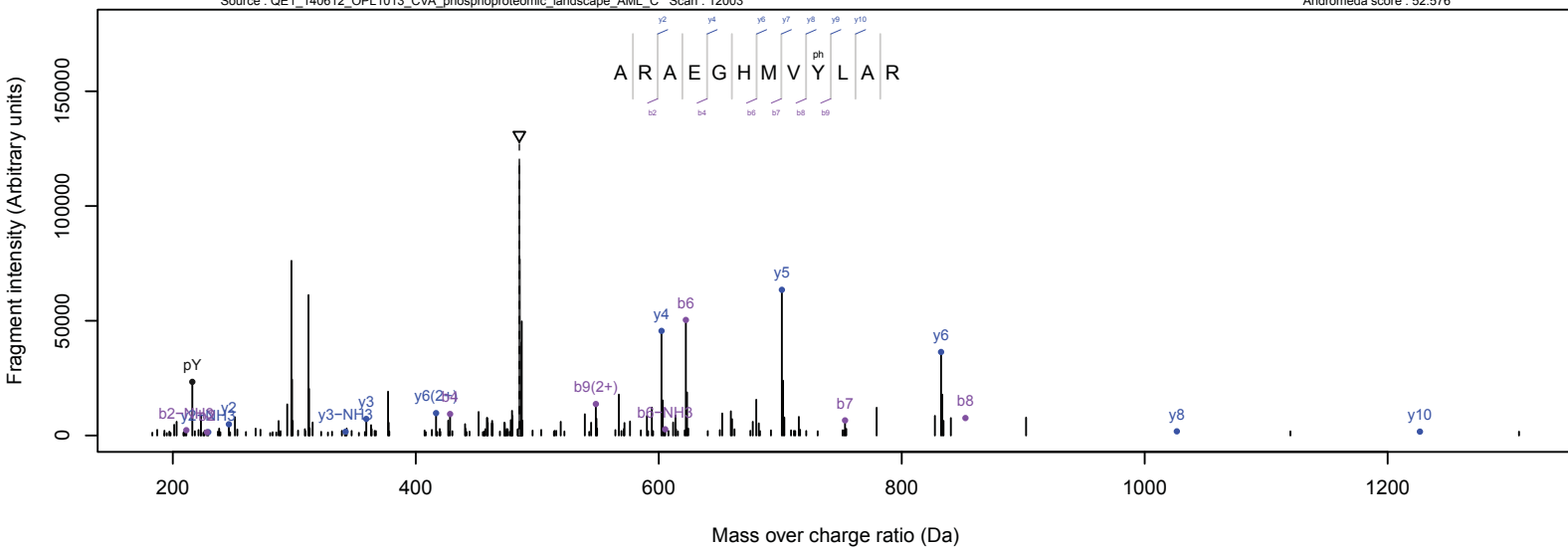
Andromeda score: 52.576



## ARAEGHMVY(ph)LAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12003

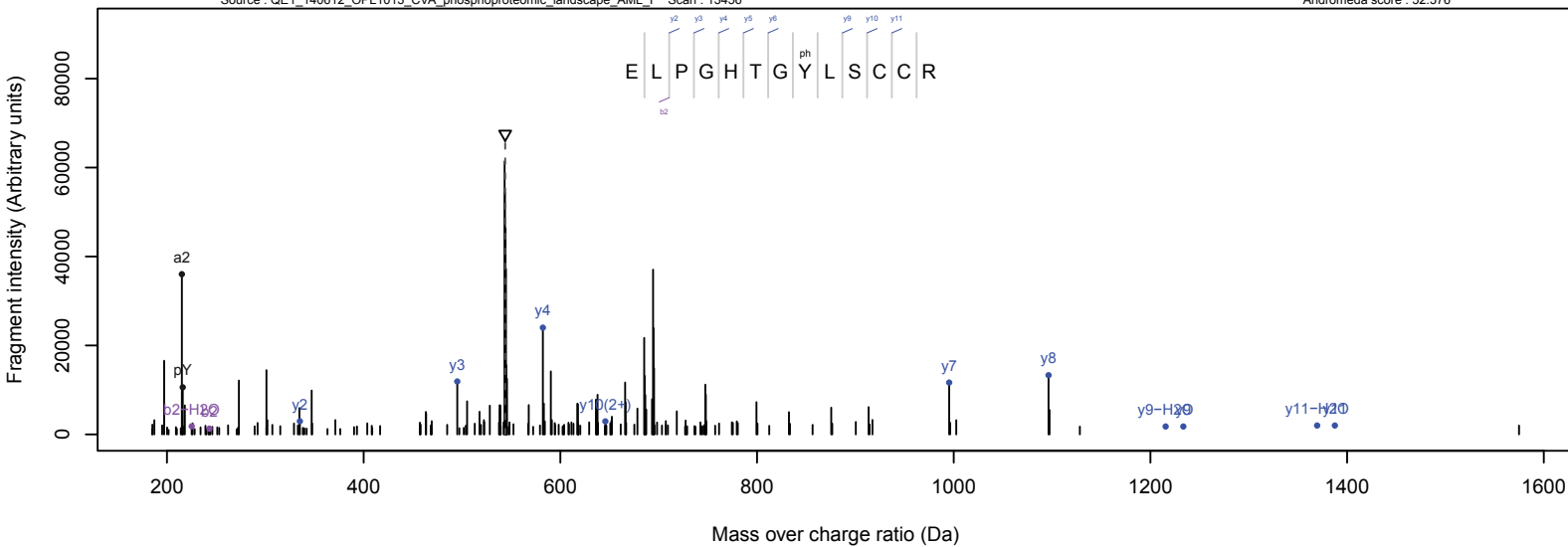
Andromeda score: 52.576



## ELPGHTGY(ph)LSCCR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan: 13456

Andromeda score: 52.576

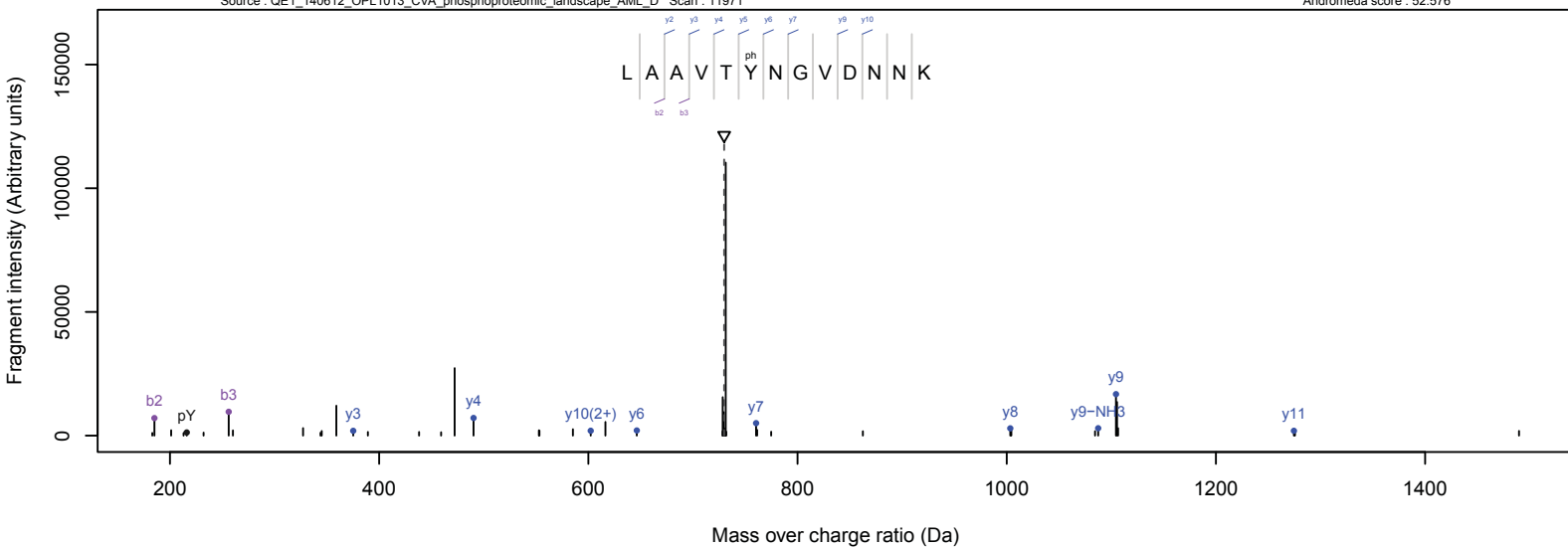




## LAAVY(ph)NGVDNKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 11971

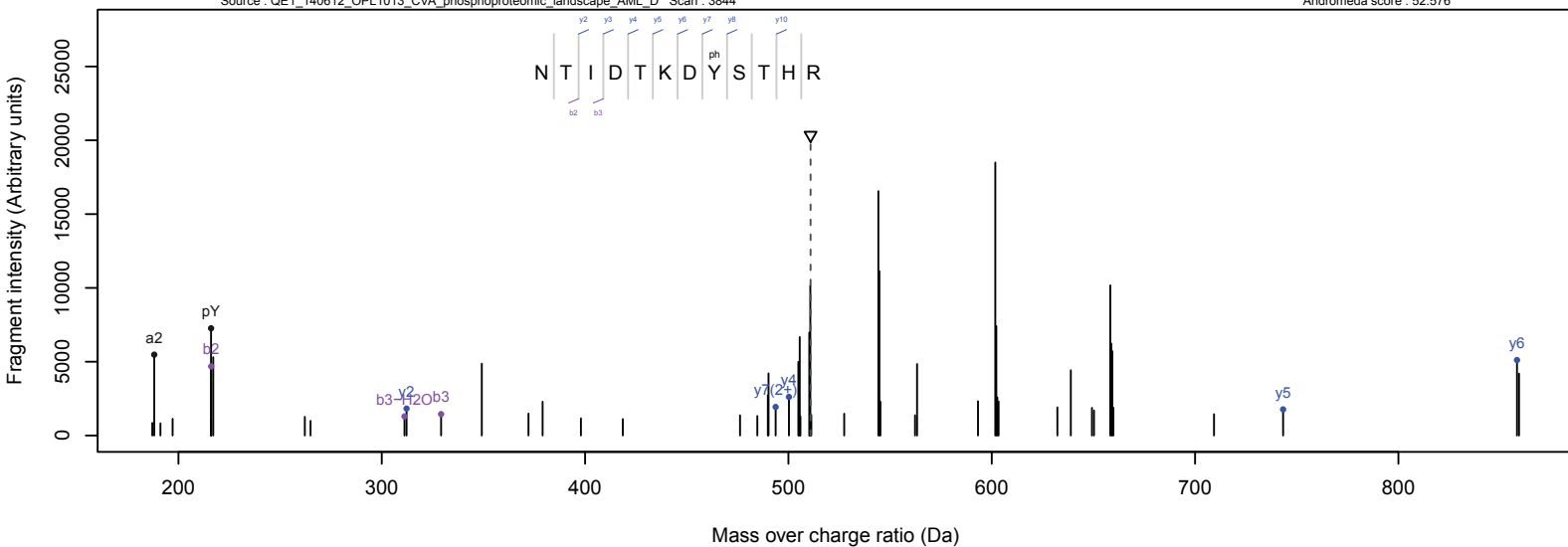
Andromeda score : 52.576



## NTIDTKDY(ph)STHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 3844

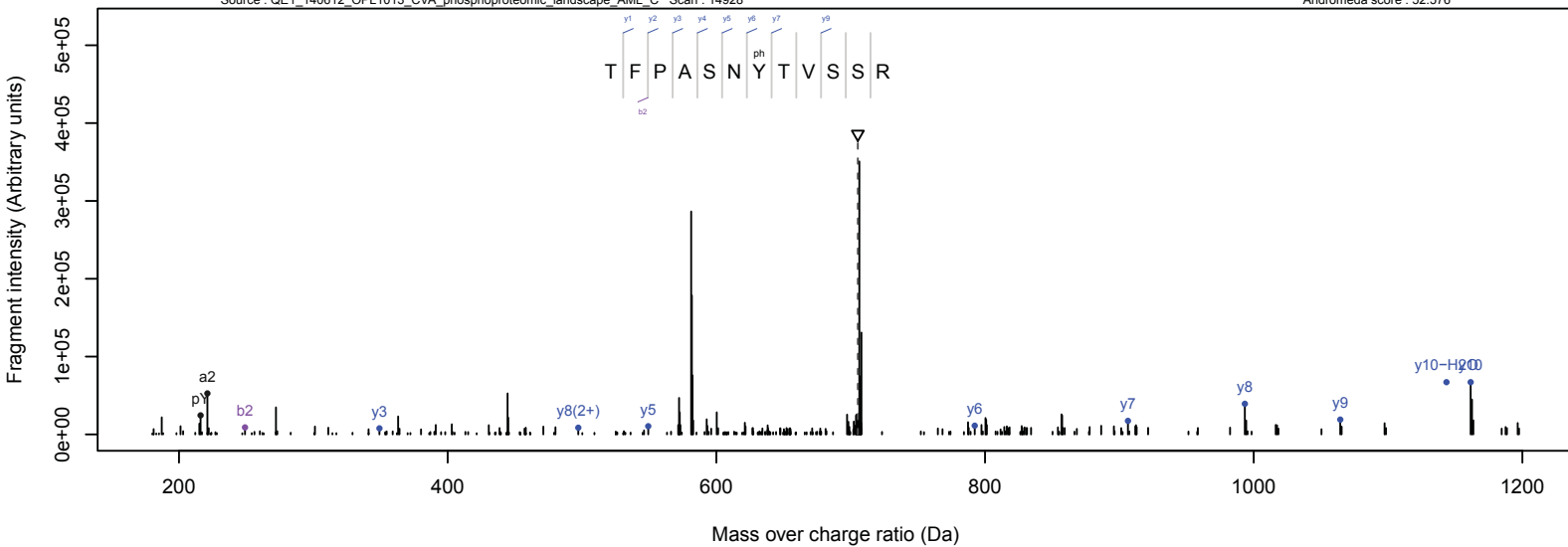
Andromeda score : 52.576



## TFPASNY(ph)TVSSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14928

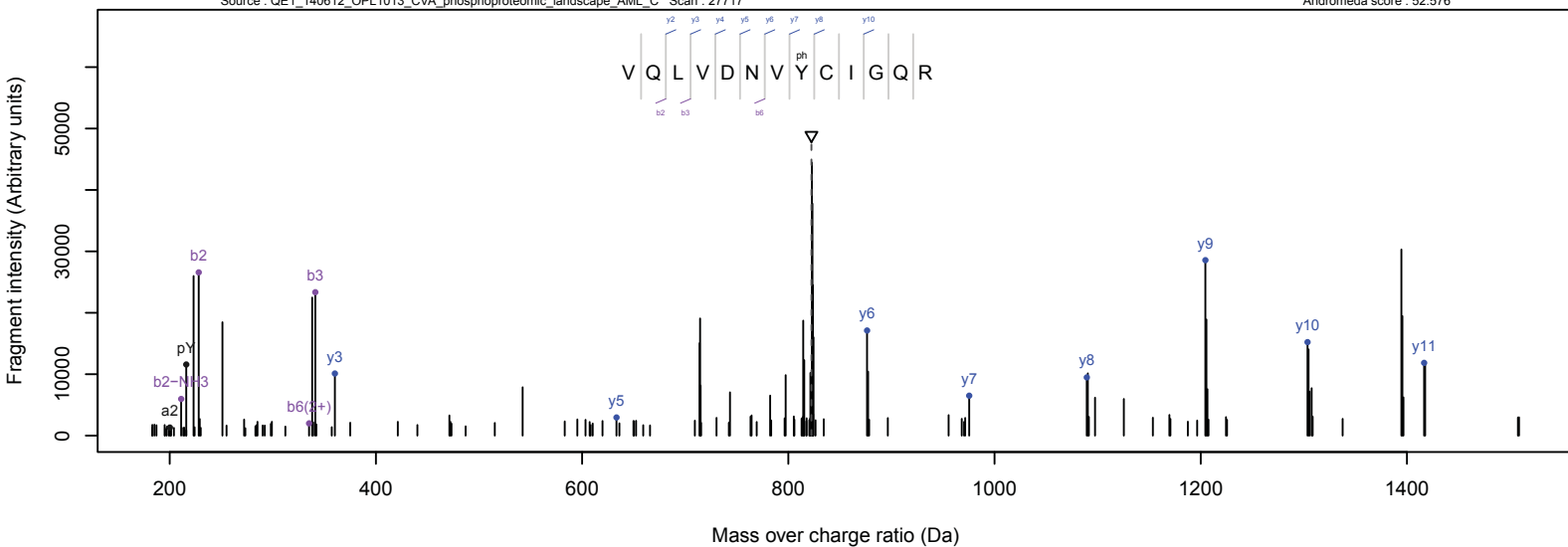
Andromeda score : 52.576



## VQLVDNVY(ph)CIGQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27717

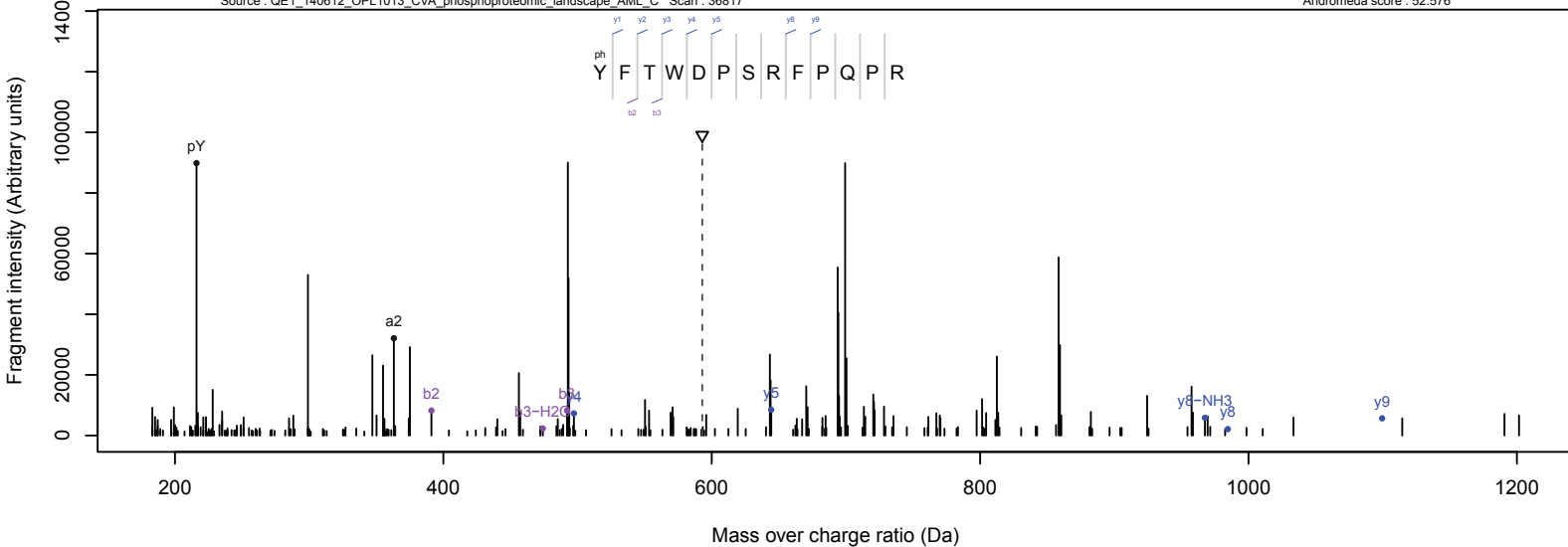
Andromeda score : 52.576



## Y(ph)FTWDPSPRFQPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36817

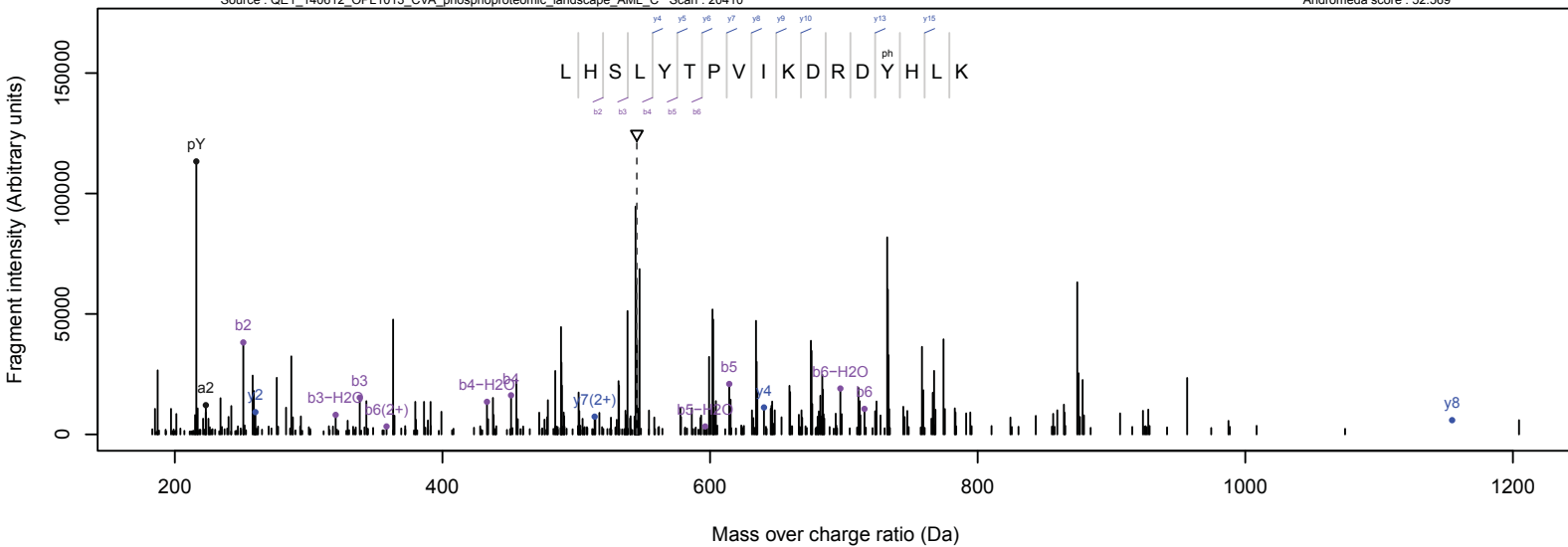
Andromeda score : 52.576



## LHSLYTPVIKDRDYHLK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20410

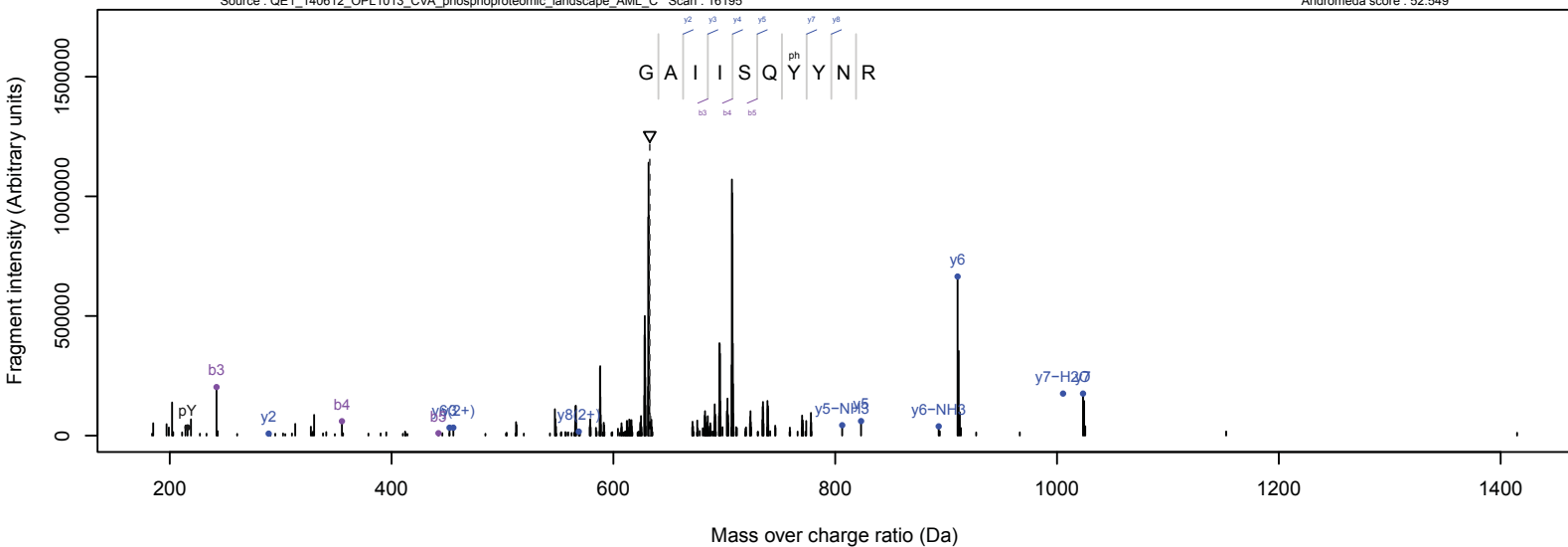
Andromeda score : 52.569



**GAIISQ(ph)YNR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 16195

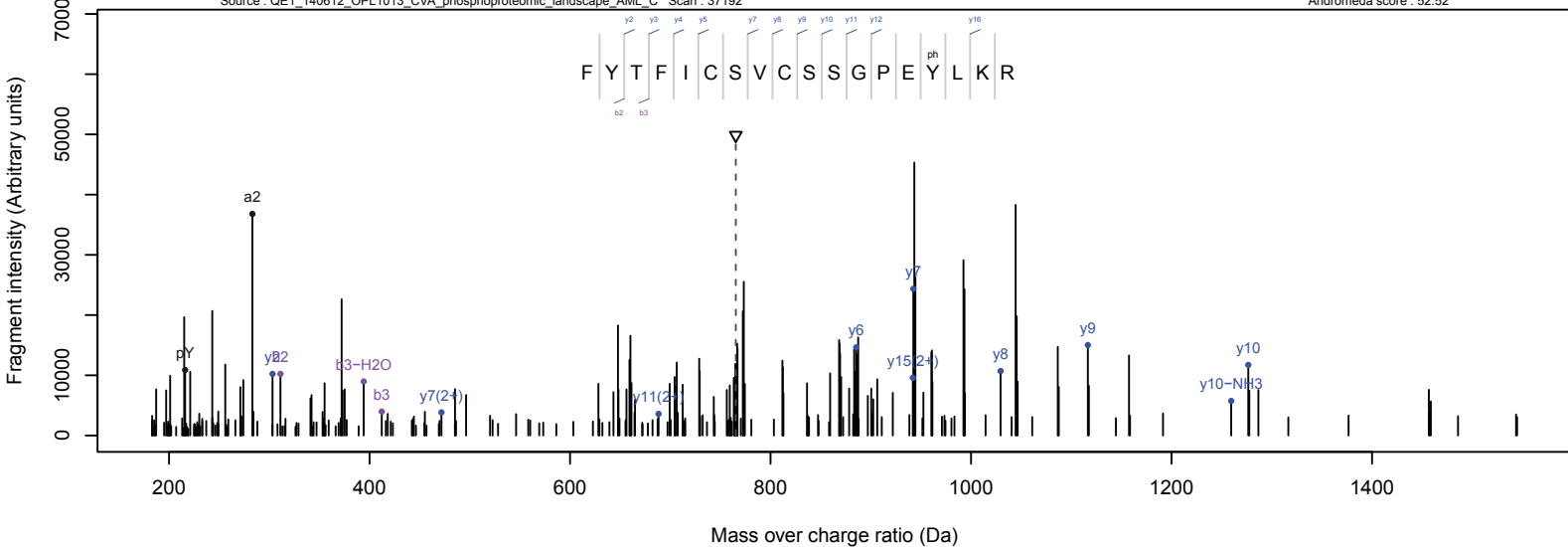
Andromeda score: 52.549



**FYTFICSVCSGPEY(L)KR-3+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37192

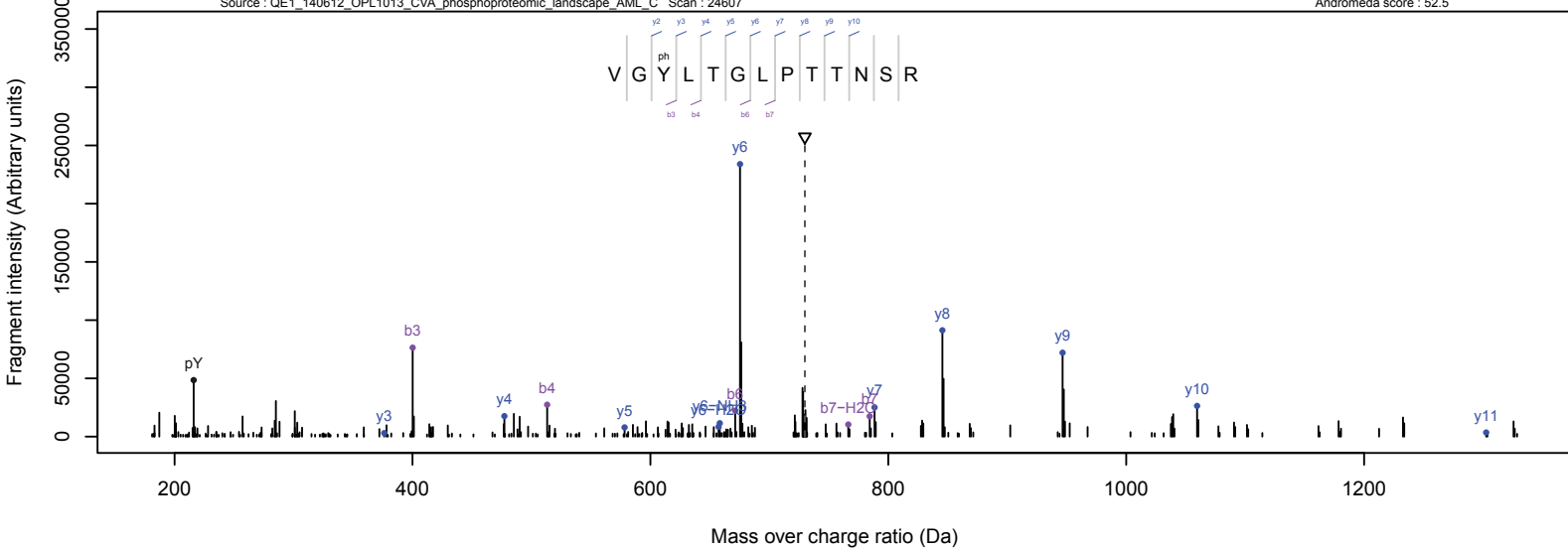
Andromeda score: 52.52



**VGY(ph)LTGLPTTNSR-2+**

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24607

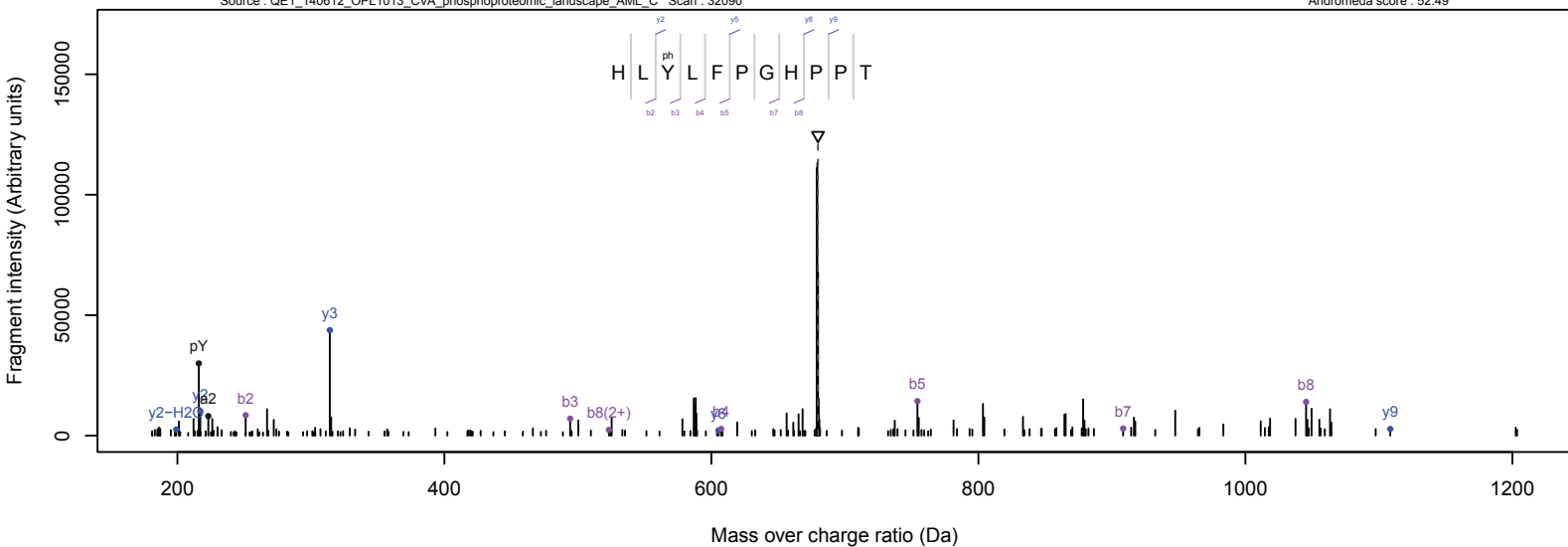
Andromeda score: 52.5



HL<sup>Y</sup>(ph)LFP<sup>G</sup>HPPT-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32090

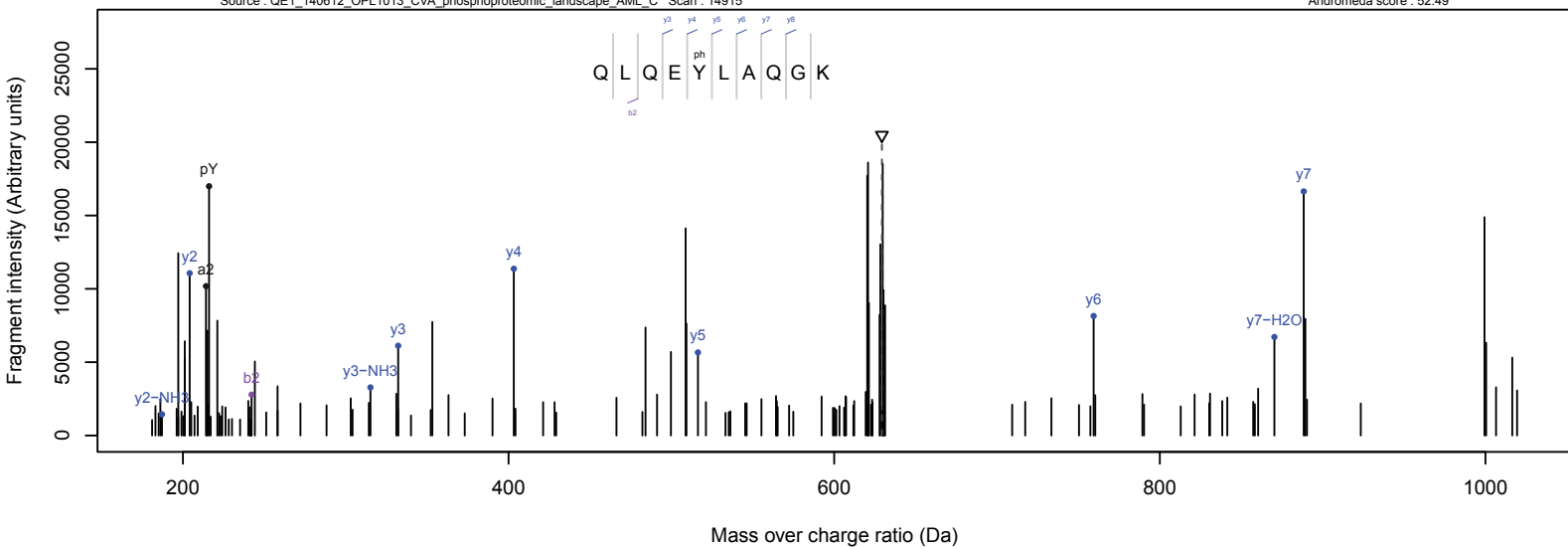
Andromeda score : 52.49



QLQE<sup>Y</sup>(ph)LAQ<sup>G</sup>KK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14915

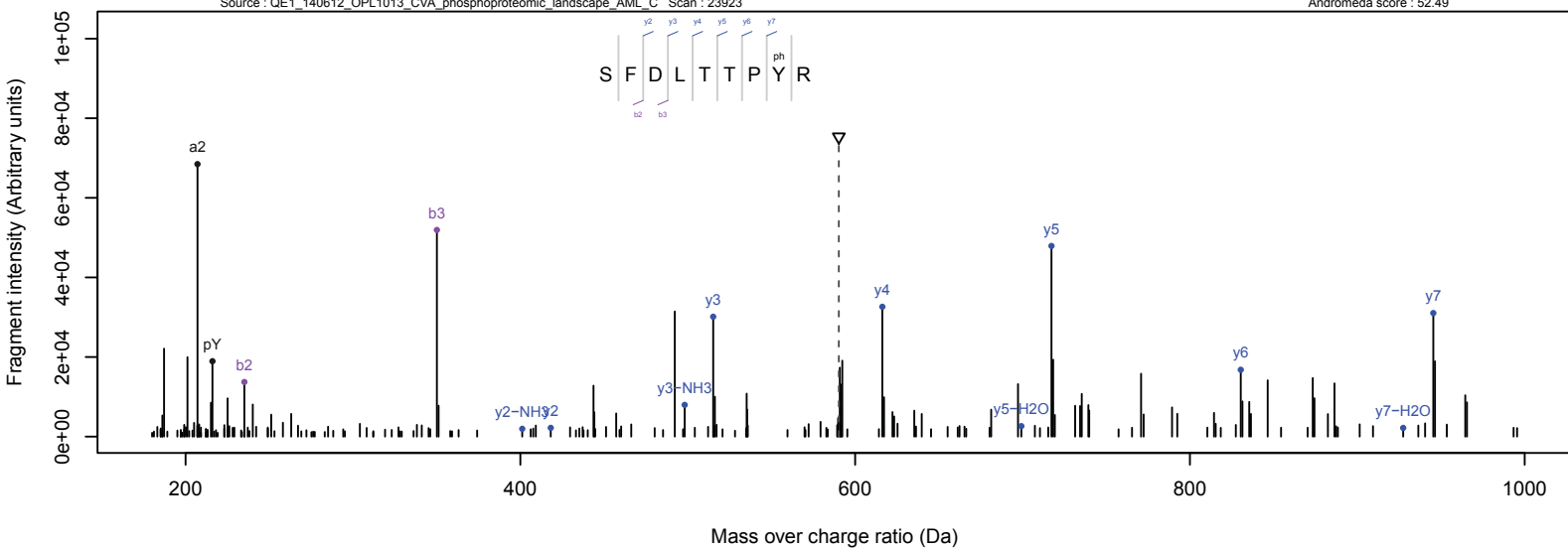
Andromeda score : 52.49



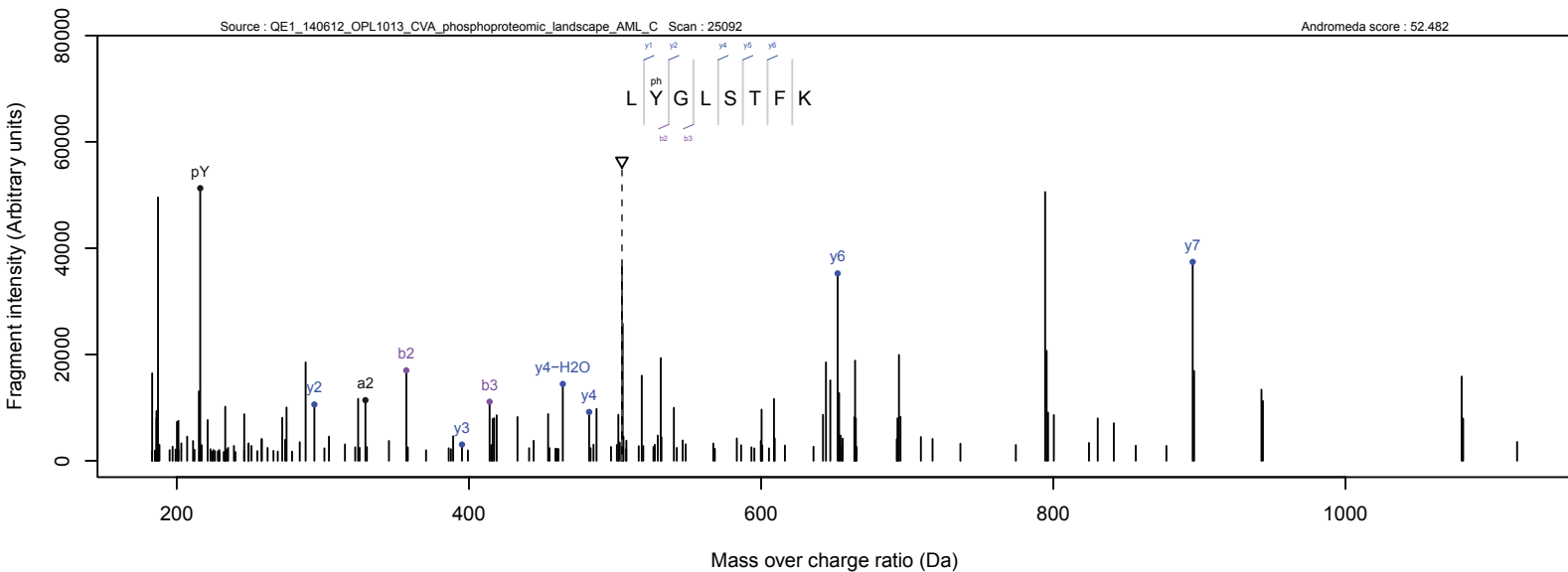
SFDLTTP<sup>Y</sup>(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23923

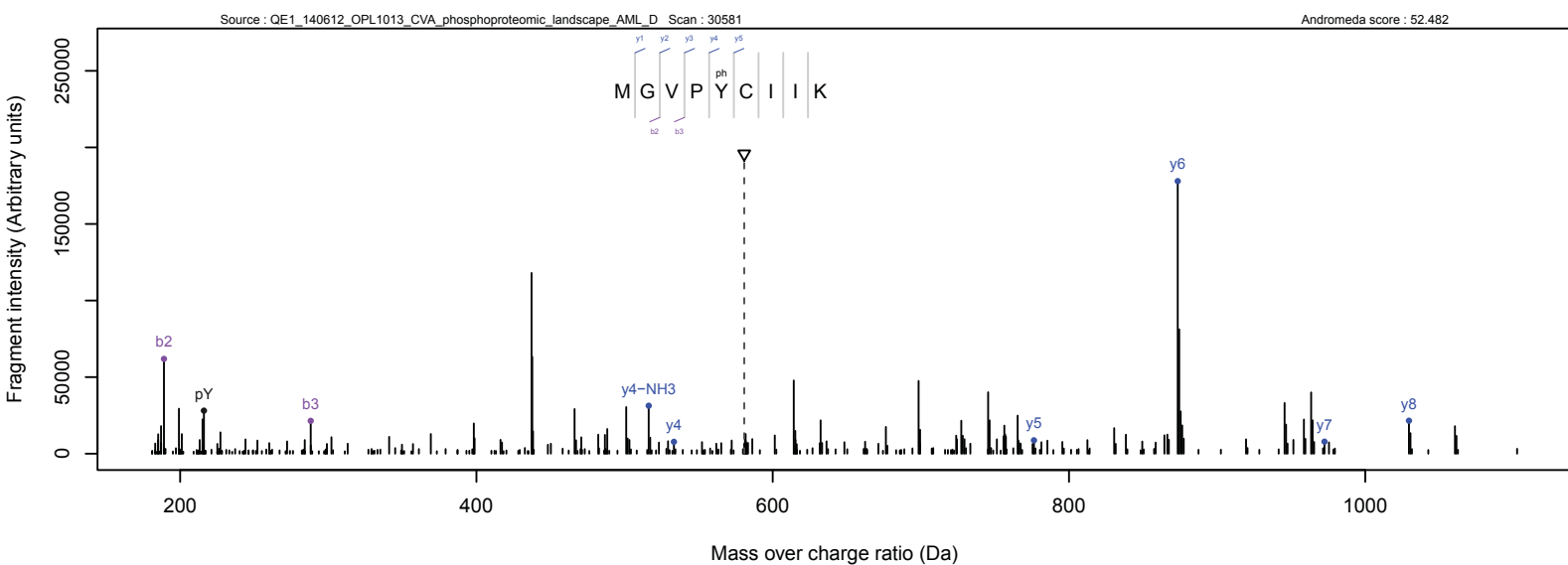
Andromeda score : 52.49



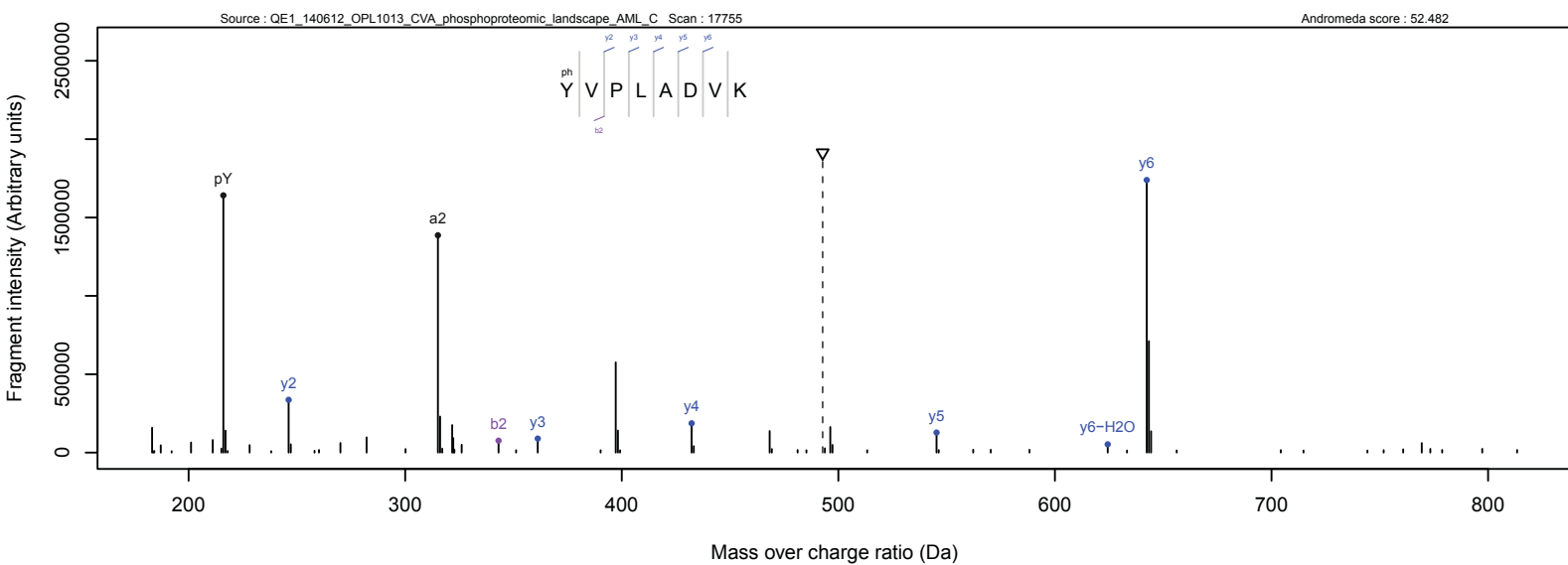
## LY(ph)GLSTFK-2+



## MGVPY(ph)CIK-2+



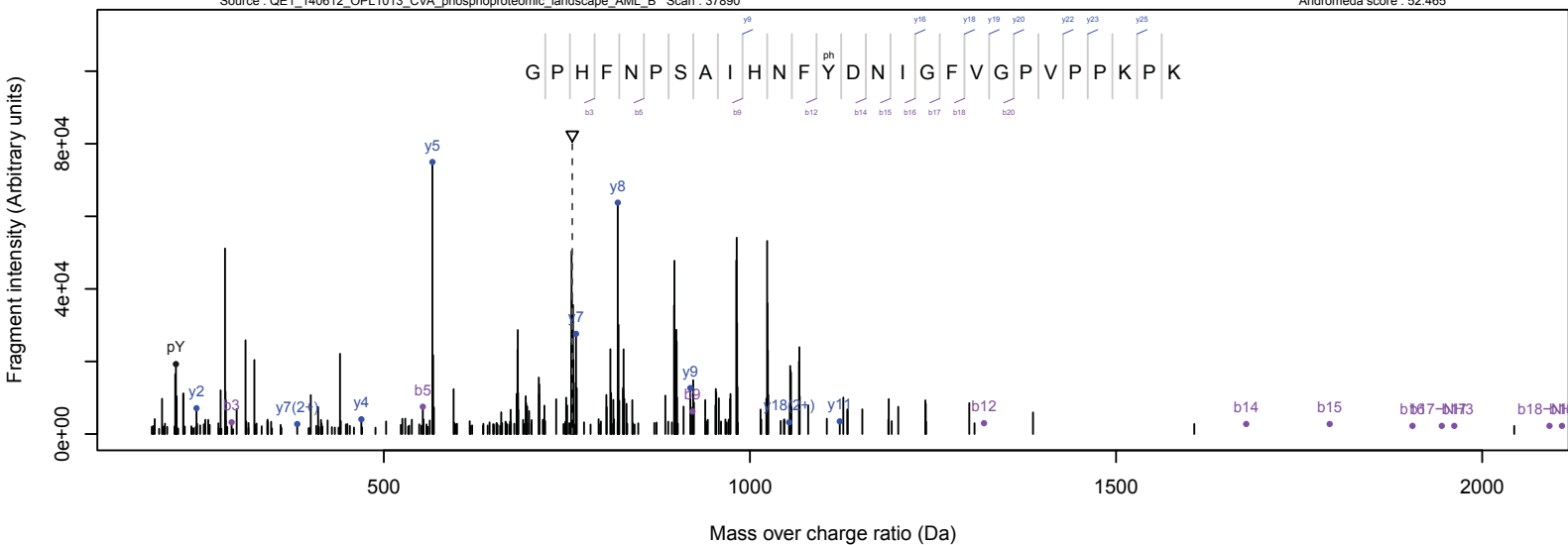
## Y(ph)VPLADVK-2+



## GPHFNPSAIHNFY(ph)DNIGFVGPVPPKPK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 37890

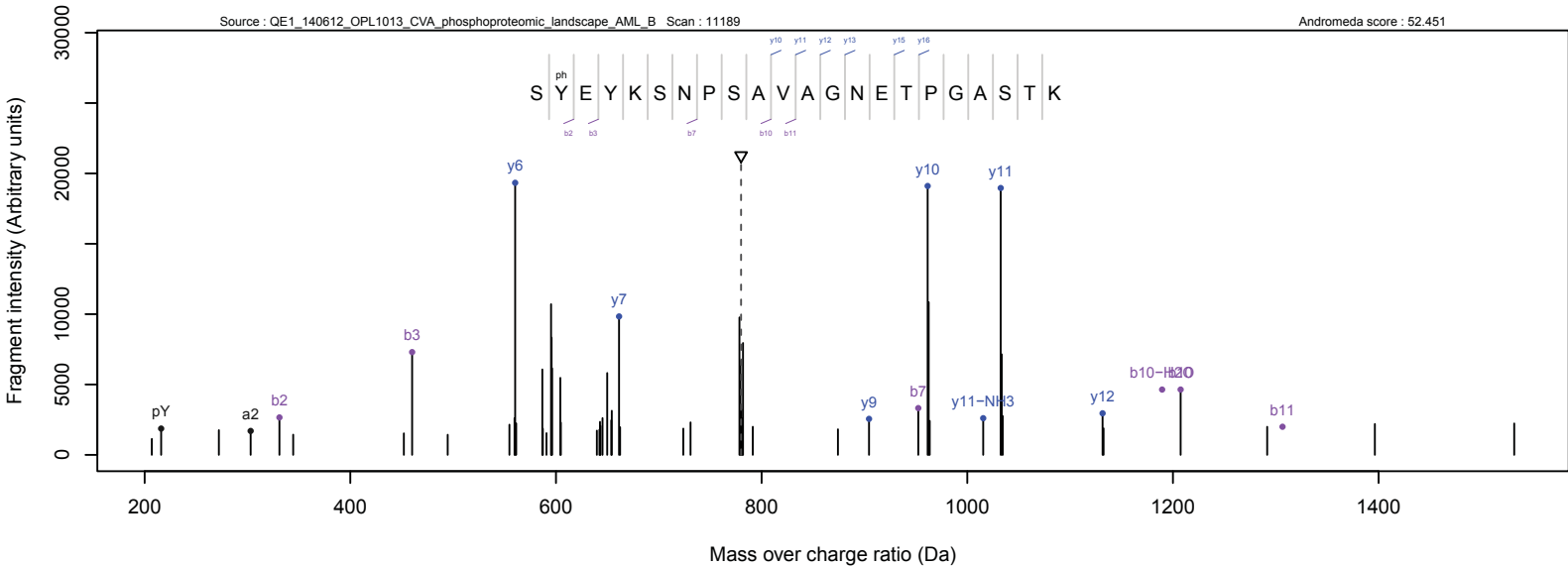
Andromeda score: 52.465



## SY(ph)EYKSNPSAVAGNETPGASTK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 11189

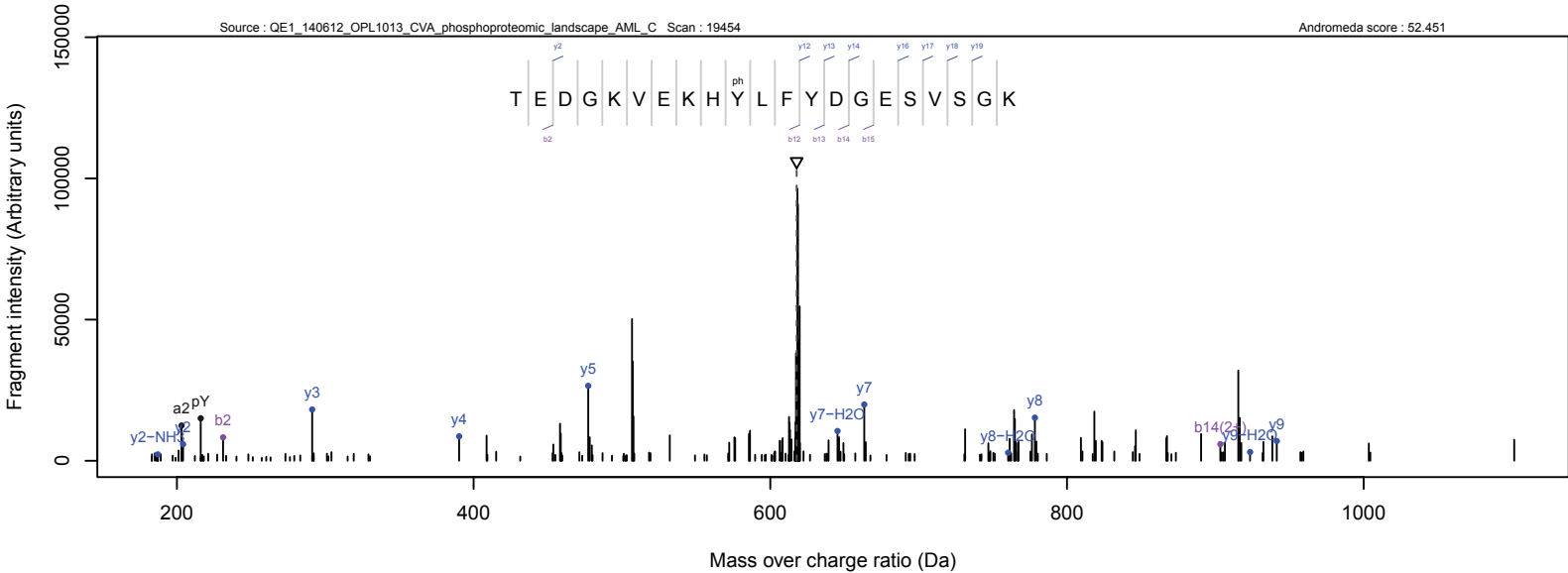
Andromeda score: 52.451



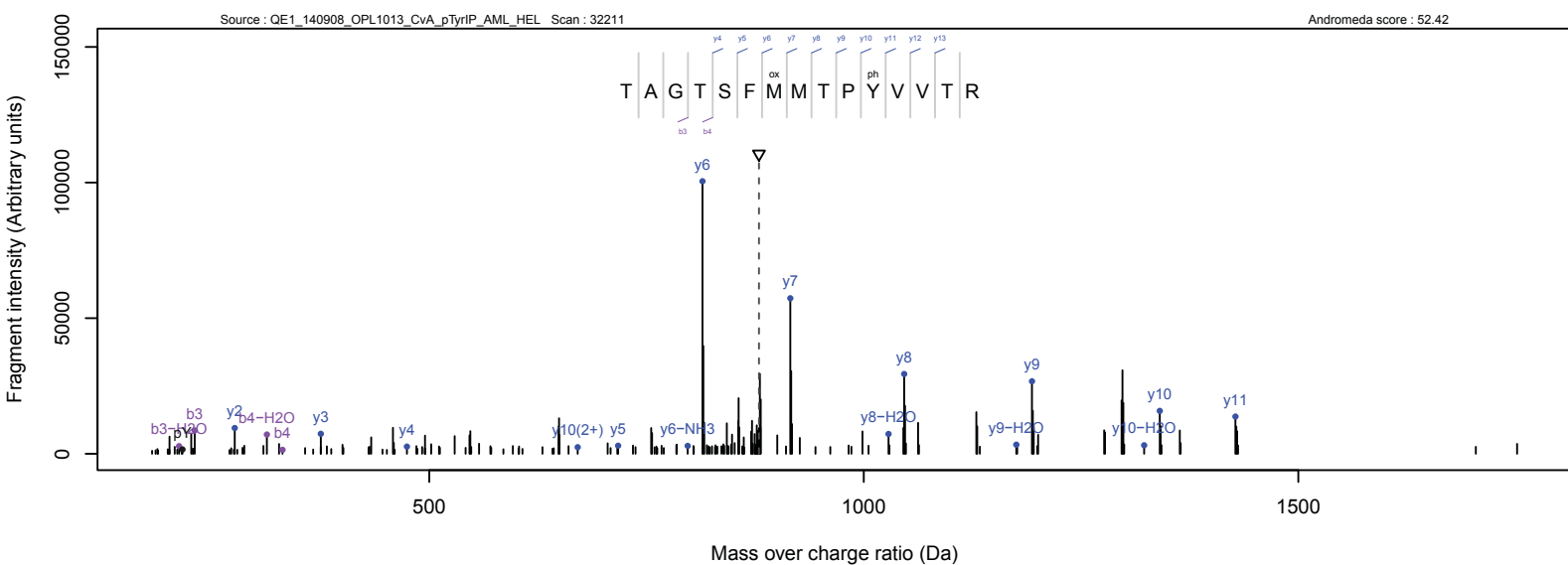
## TEDGKVEKHY(ph)LFYDGESVSGK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19454

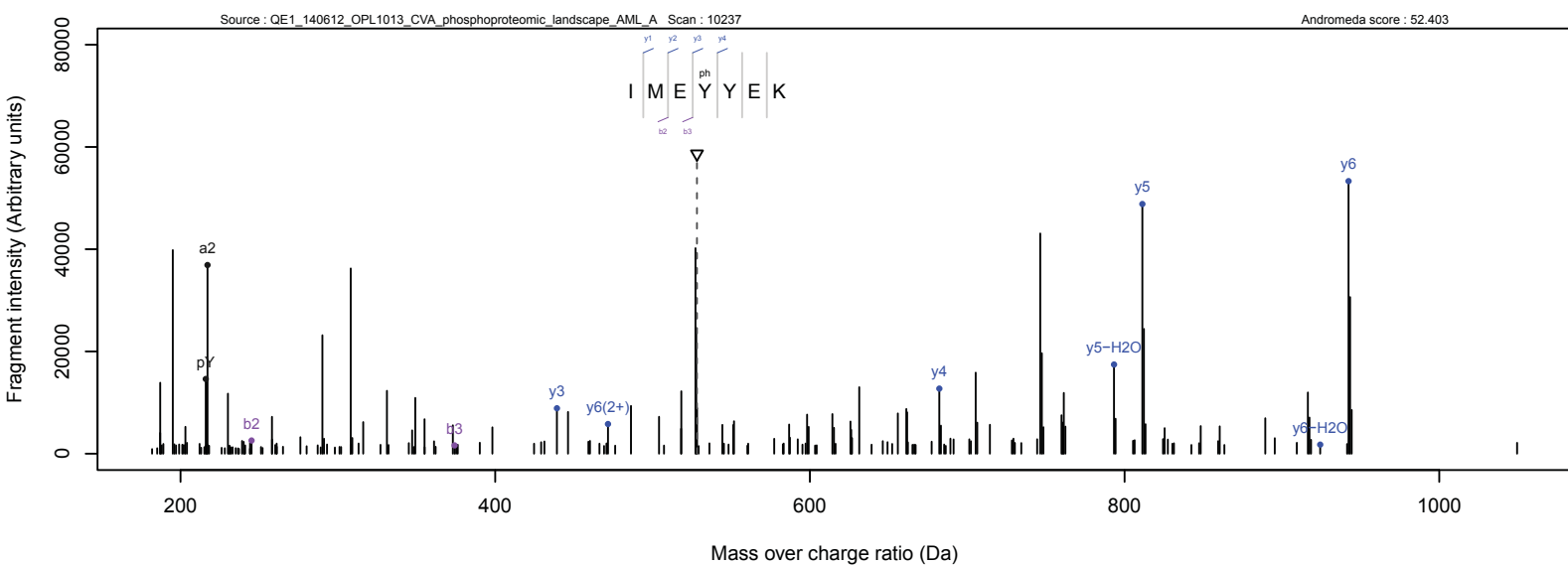
Andromeda score: 52.451



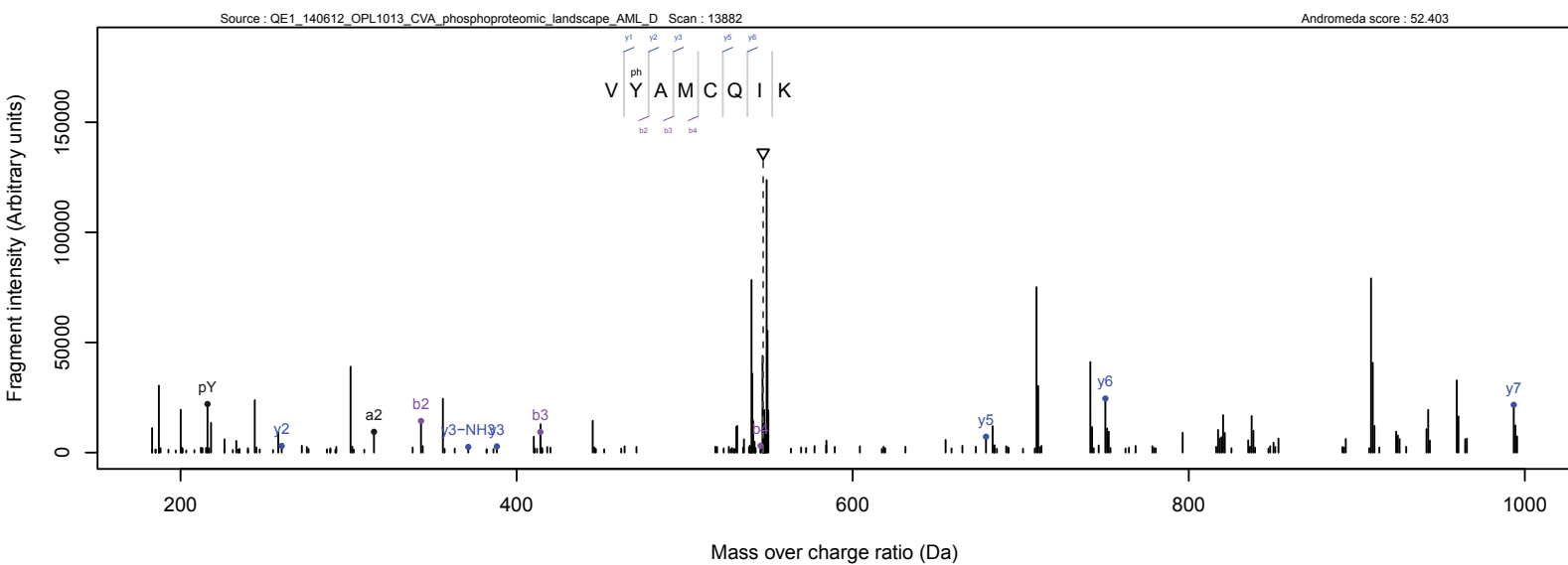
## TAGTSFM(ox)MTPY(ph)VVTR-2+

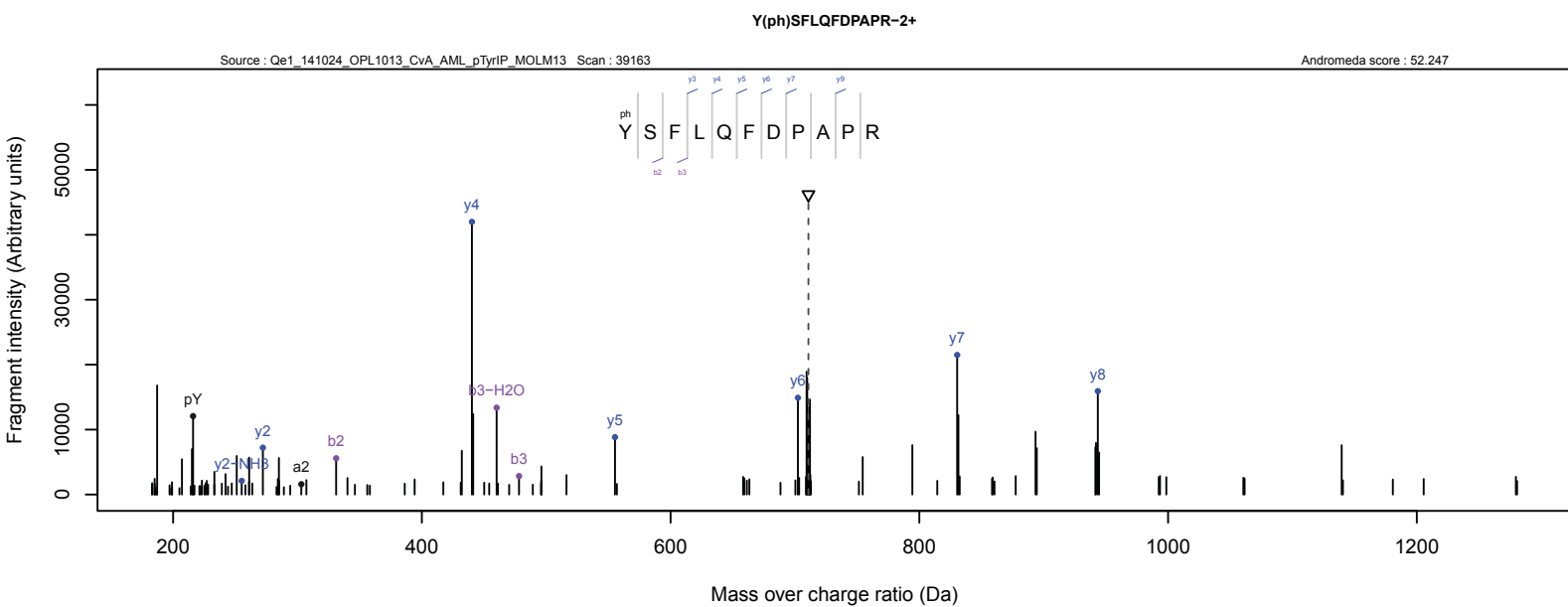
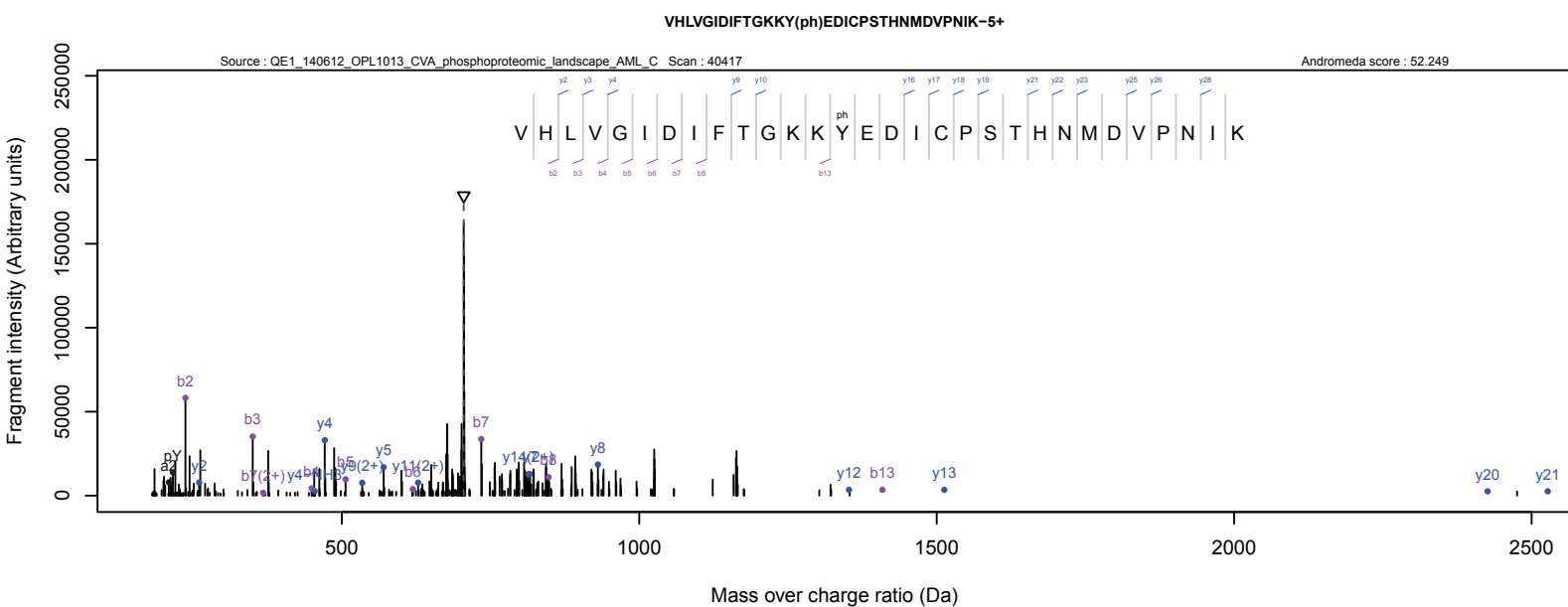
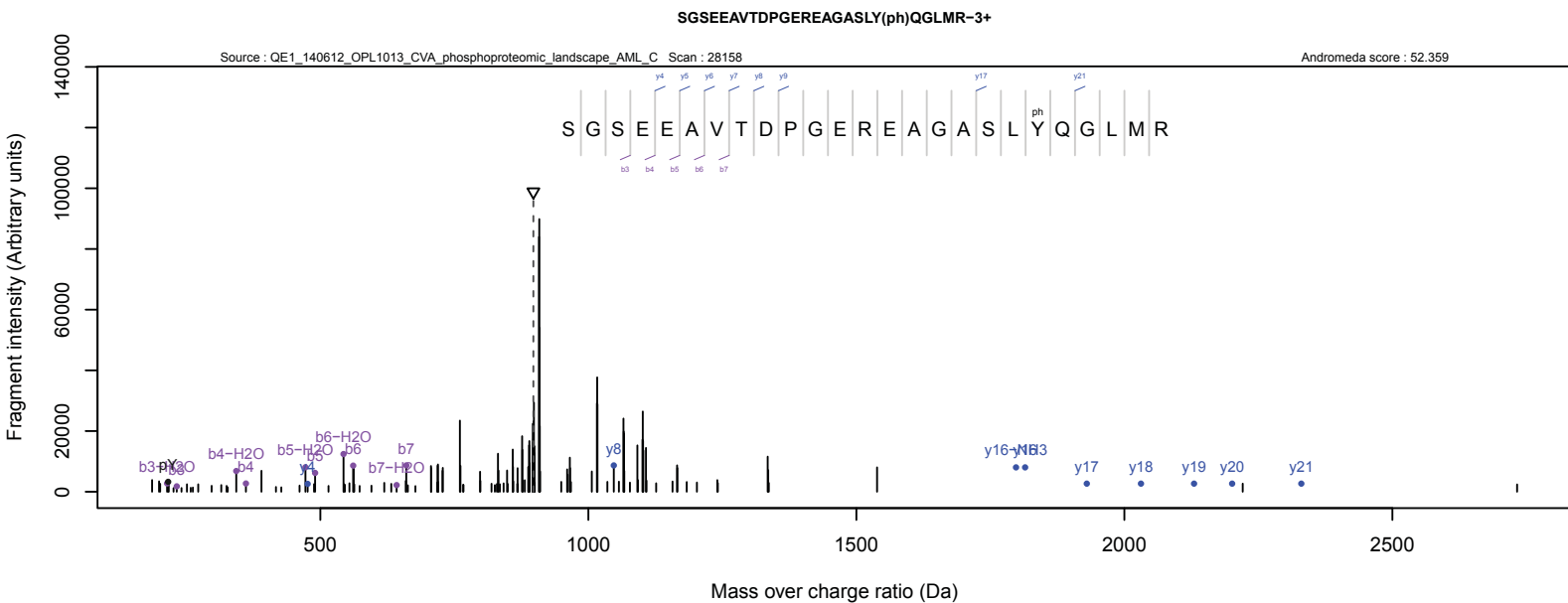


## IMEY(ph)YEK-2+



## VY(ph)AMCQIK-2+



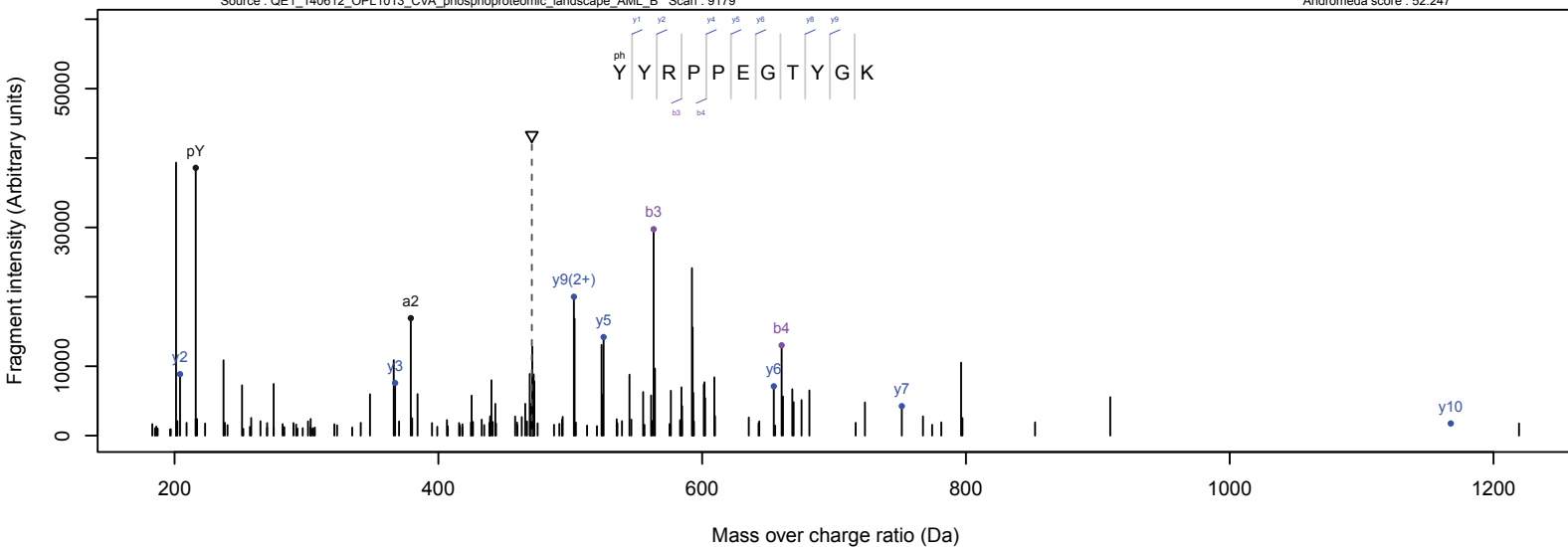




**Y(ph)YRPPEGTYGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 9179

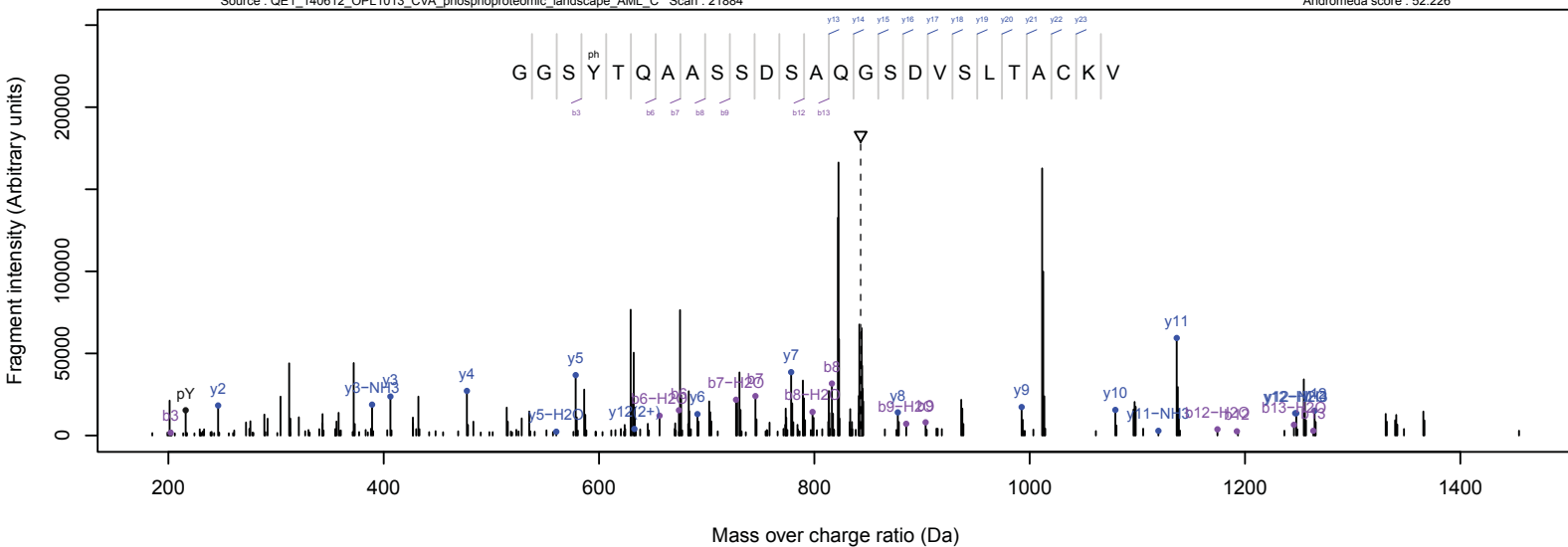
Andromeda score : 52.247



**GGSY(ph)TQAASSDAQGSDVSLTACKV-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21884

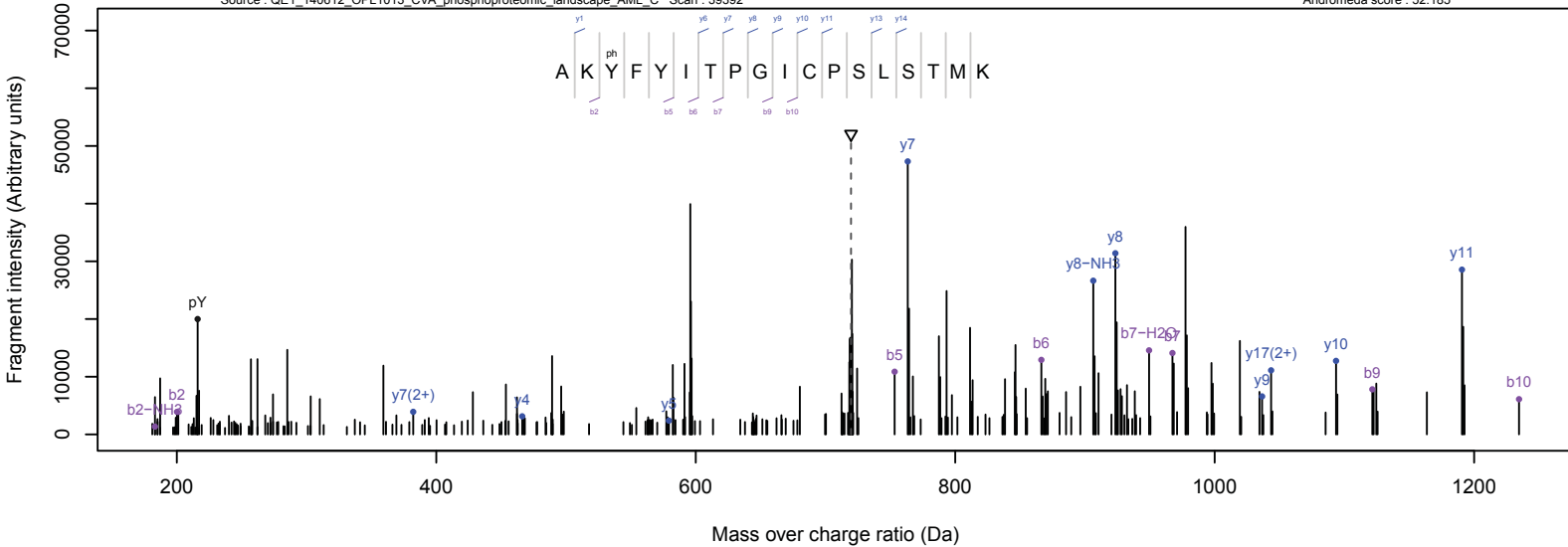
Andromeda score : 52.226



**AKY(ph)FYITPGICPSLSTMK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39392

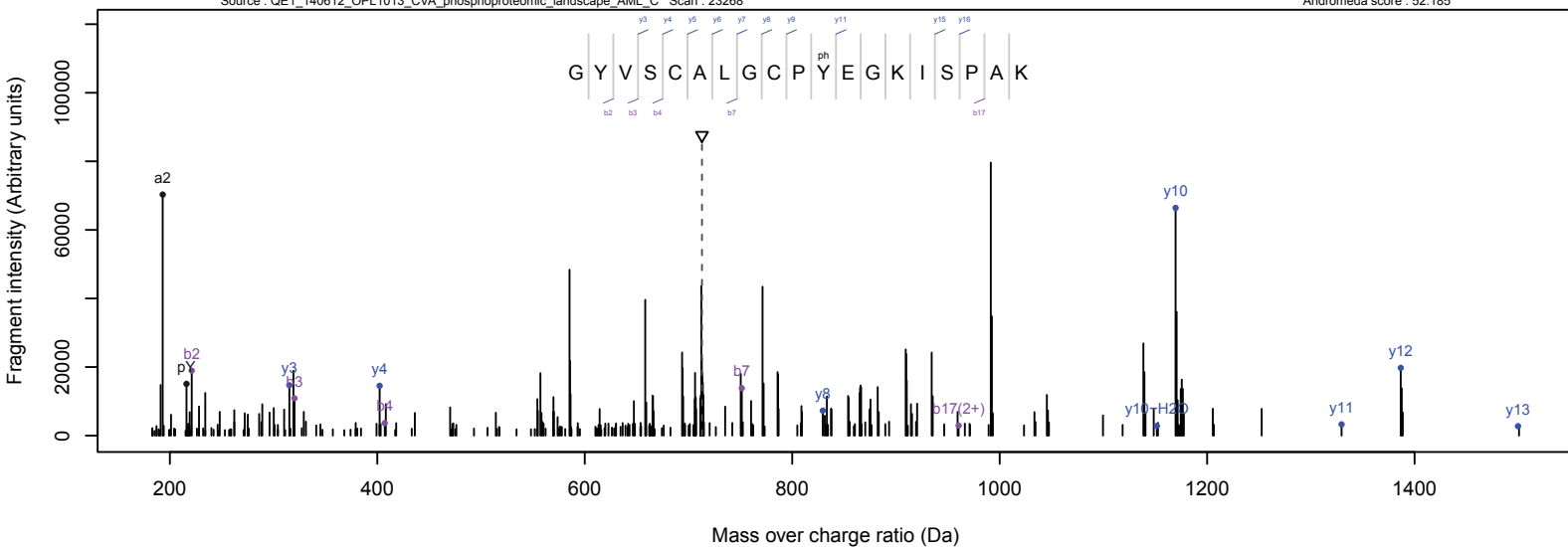
Andromeda score : 52.185



## GYVSCALGCPY(ph)EGKISPAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23268

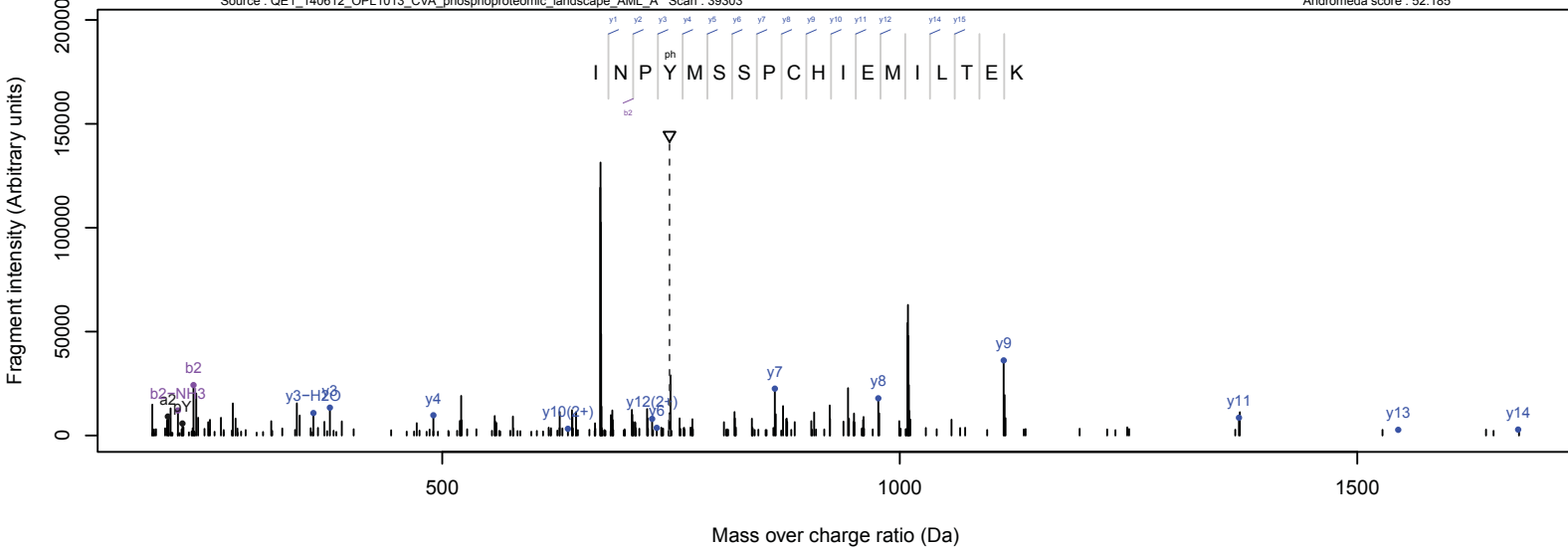
Andromeda score : 52.185



## INPY(ph)MSSPCHIEMLTEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 39303

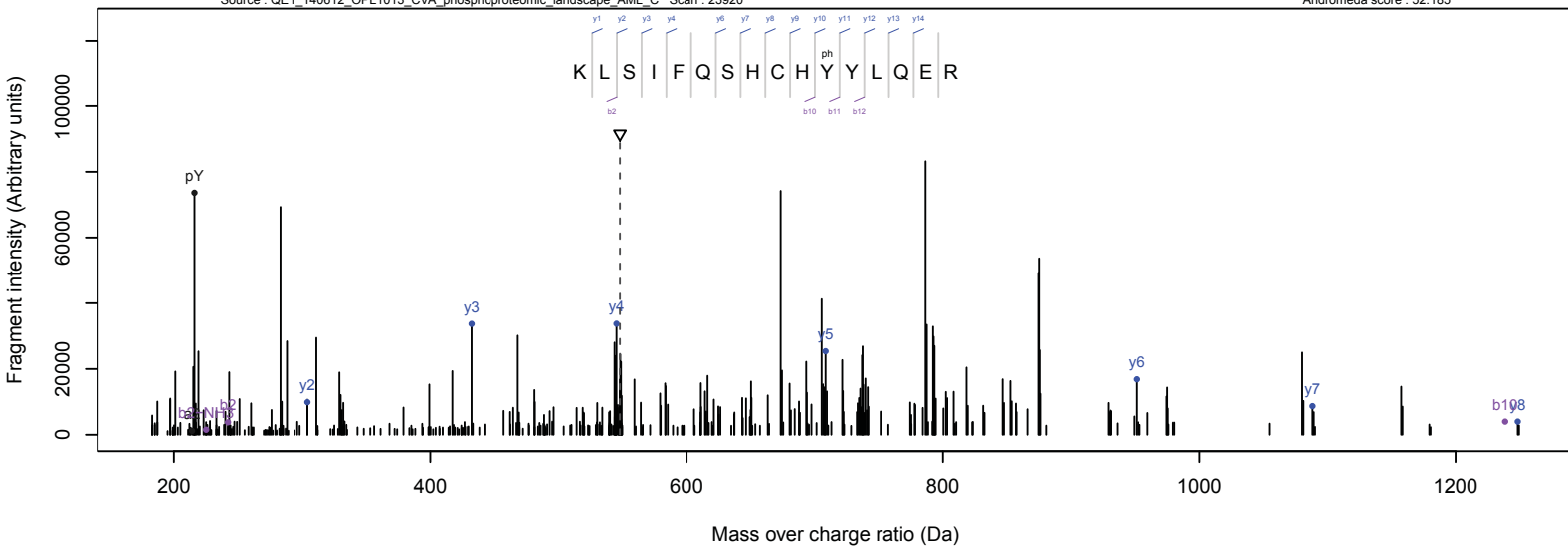
Andromeda score : 52.185



## KLSIFQSHCHY(ph)YLQER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23920

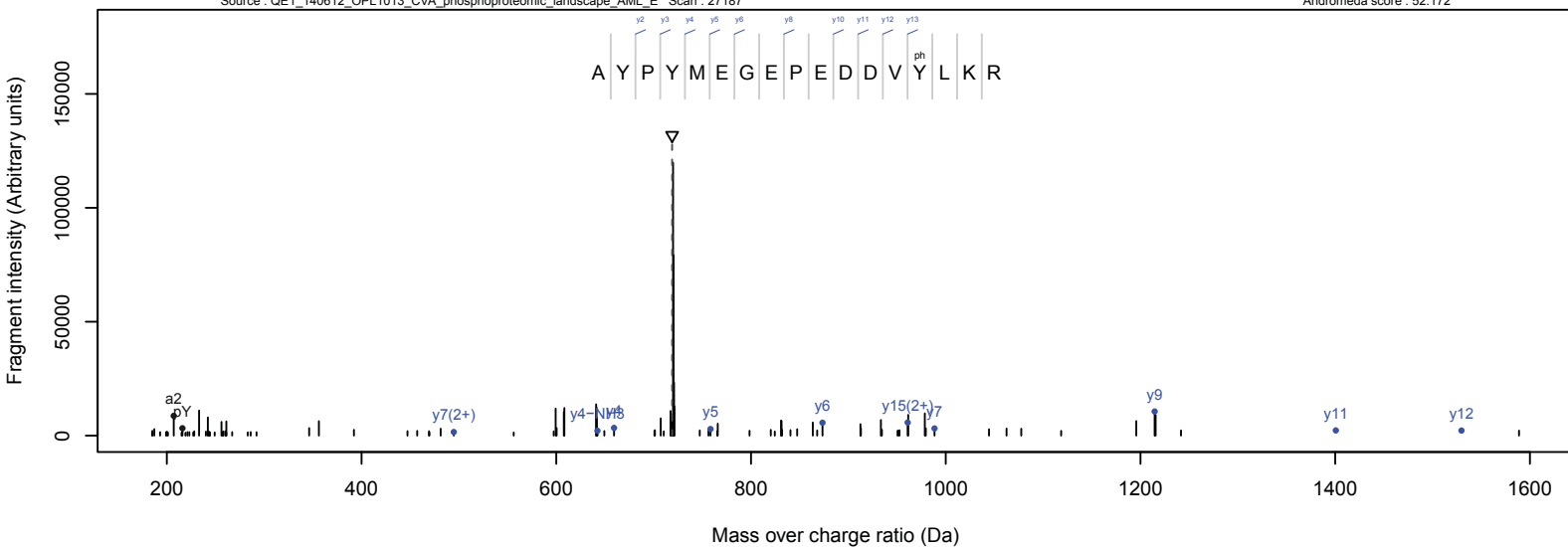
Andromeda score : 52.185



AYPYMEGEPEDDVY(ph)LKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 27187

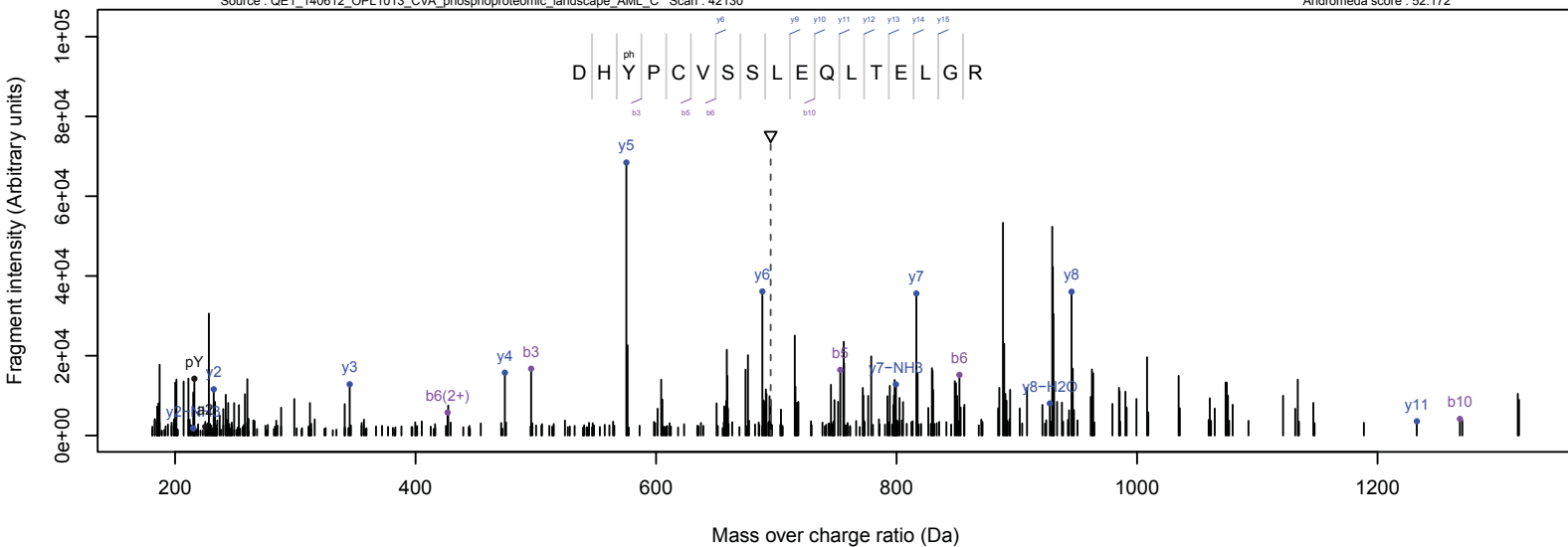
Andromeda score : 52.172



DHY(ph)PCVSSLEQLTELGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42130

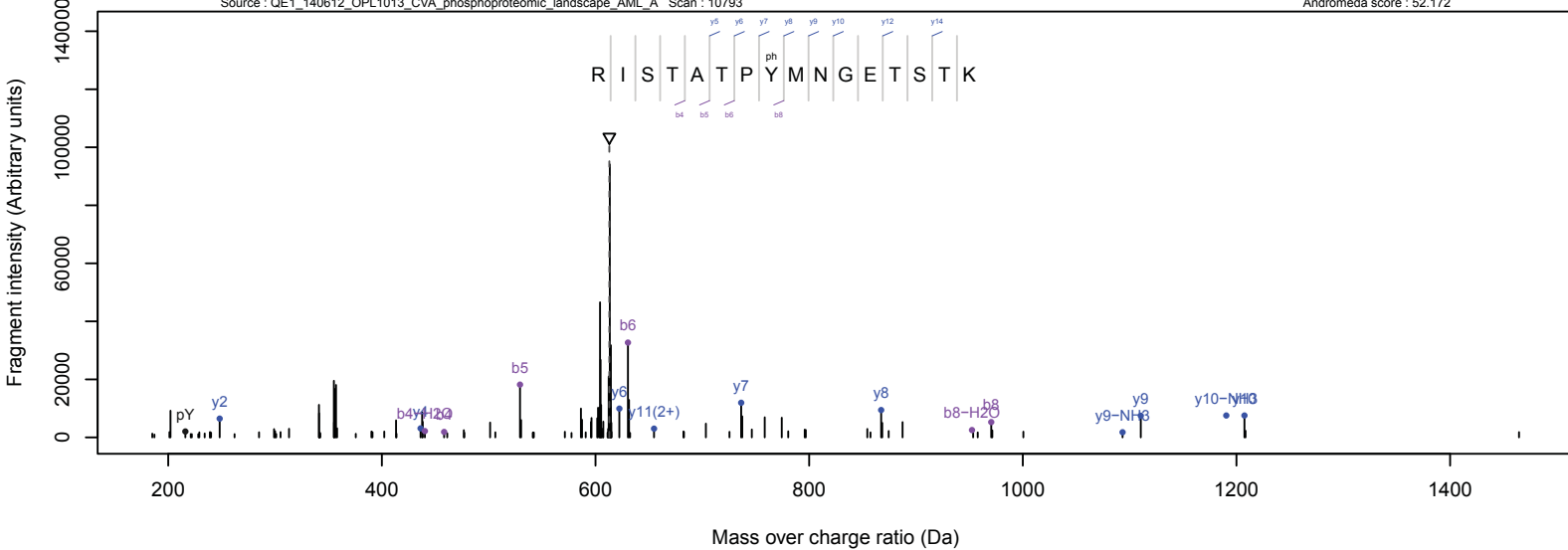
Andromeda score : 52.172



RISTATPY(ph)MNGETSTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 10793

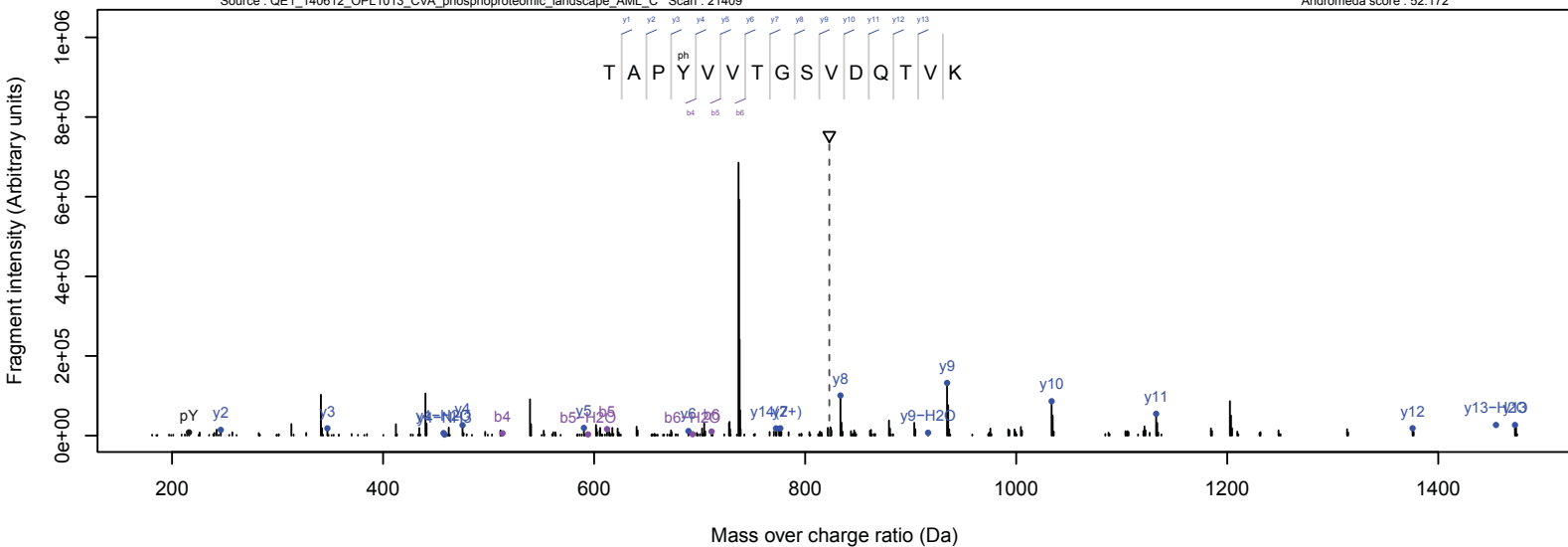
Andromeda score : 52.172



## TAPY(ph)VVTGSVDQTVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21409

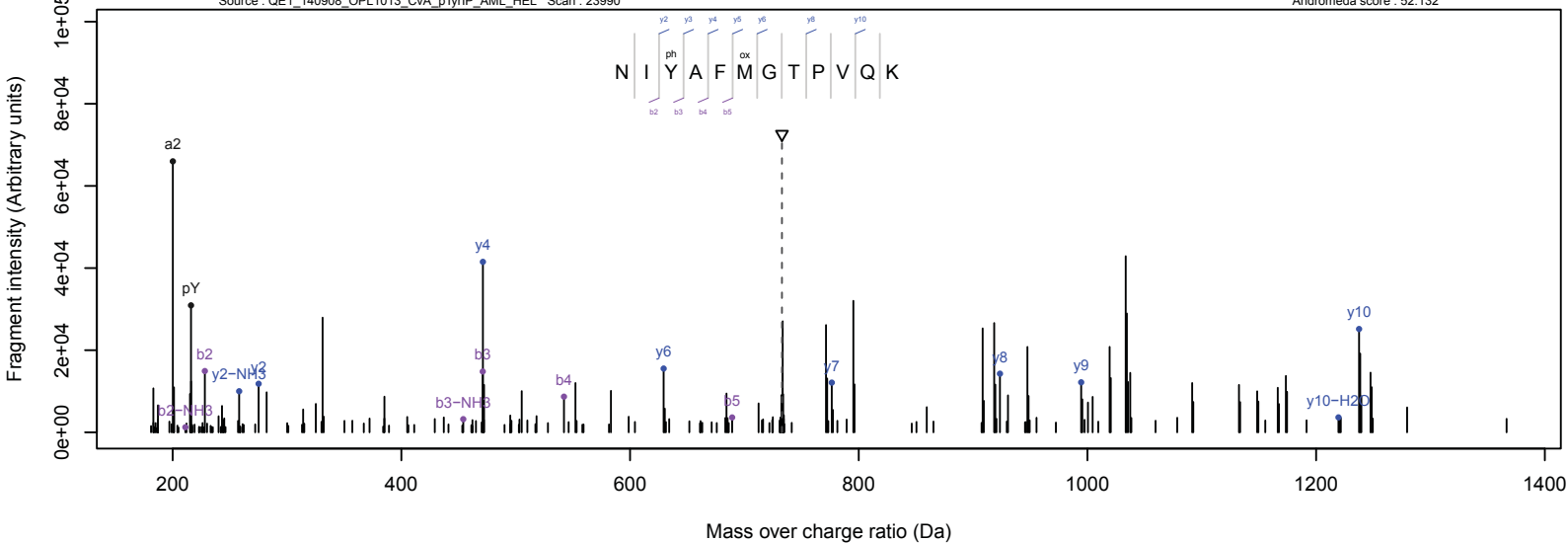
Andromeda score : 52.172



## NIY(ph)AFM(ox)GTPVQK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 23990

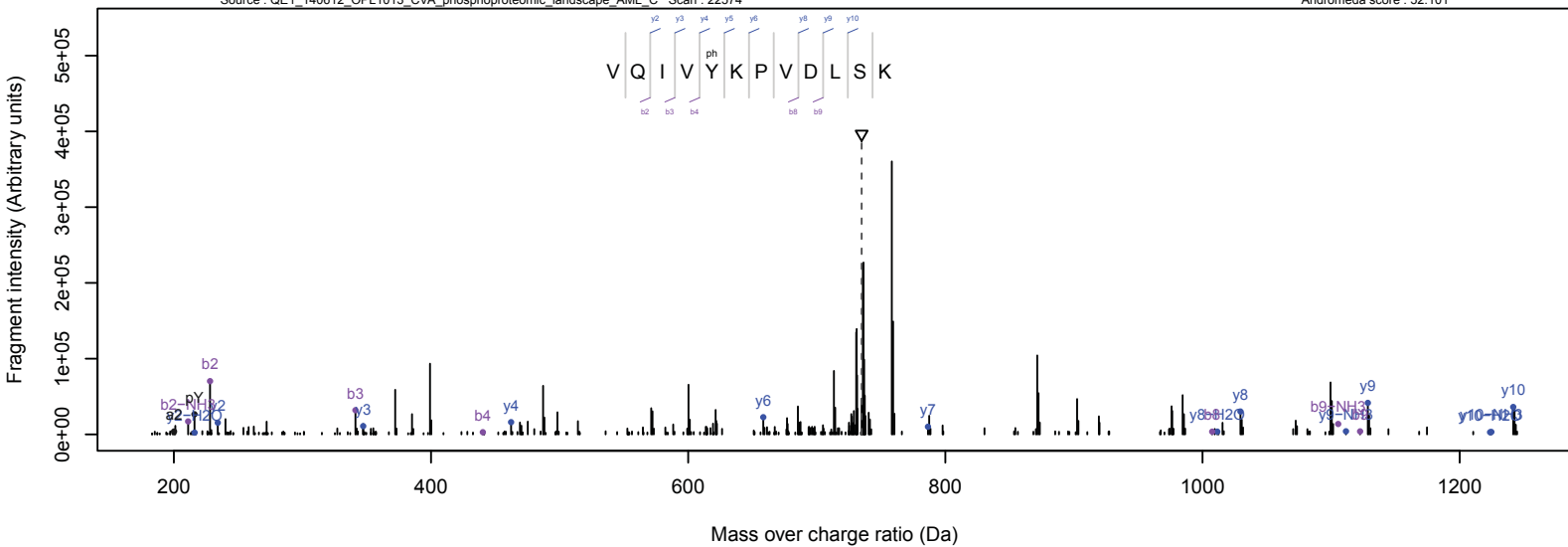
Andromeda score : 52.132

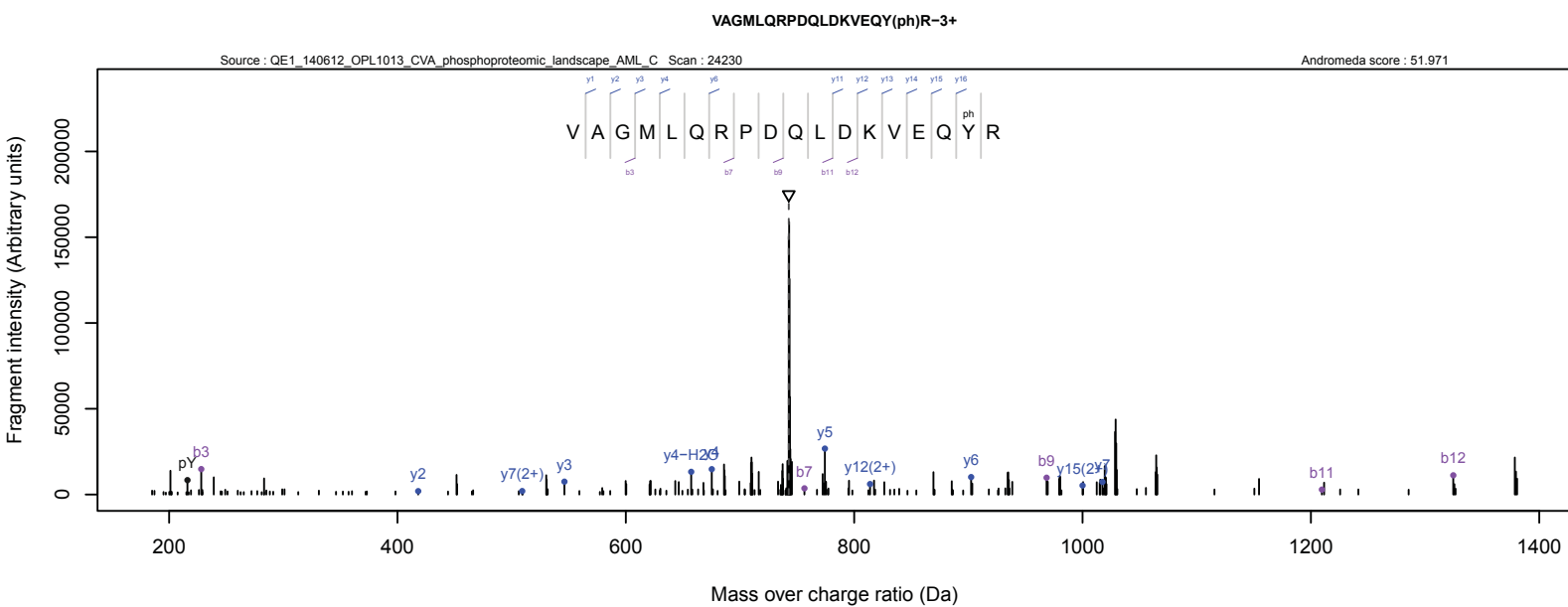
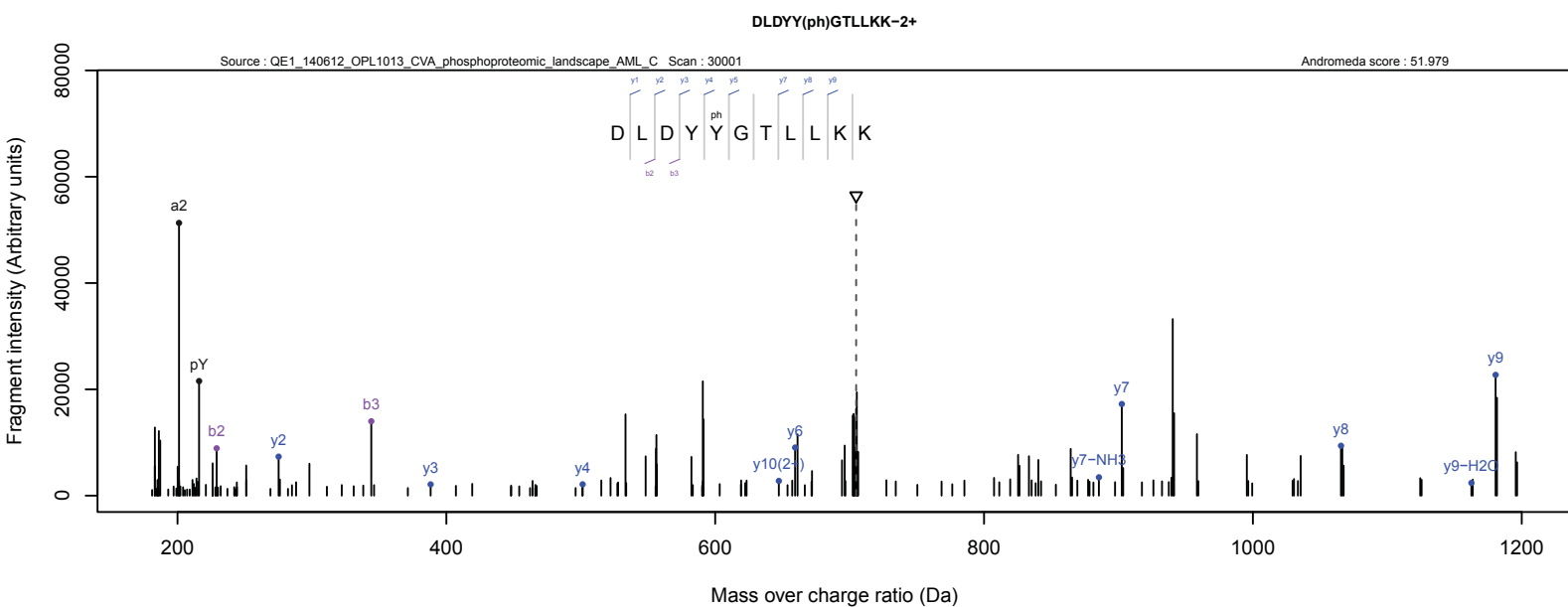
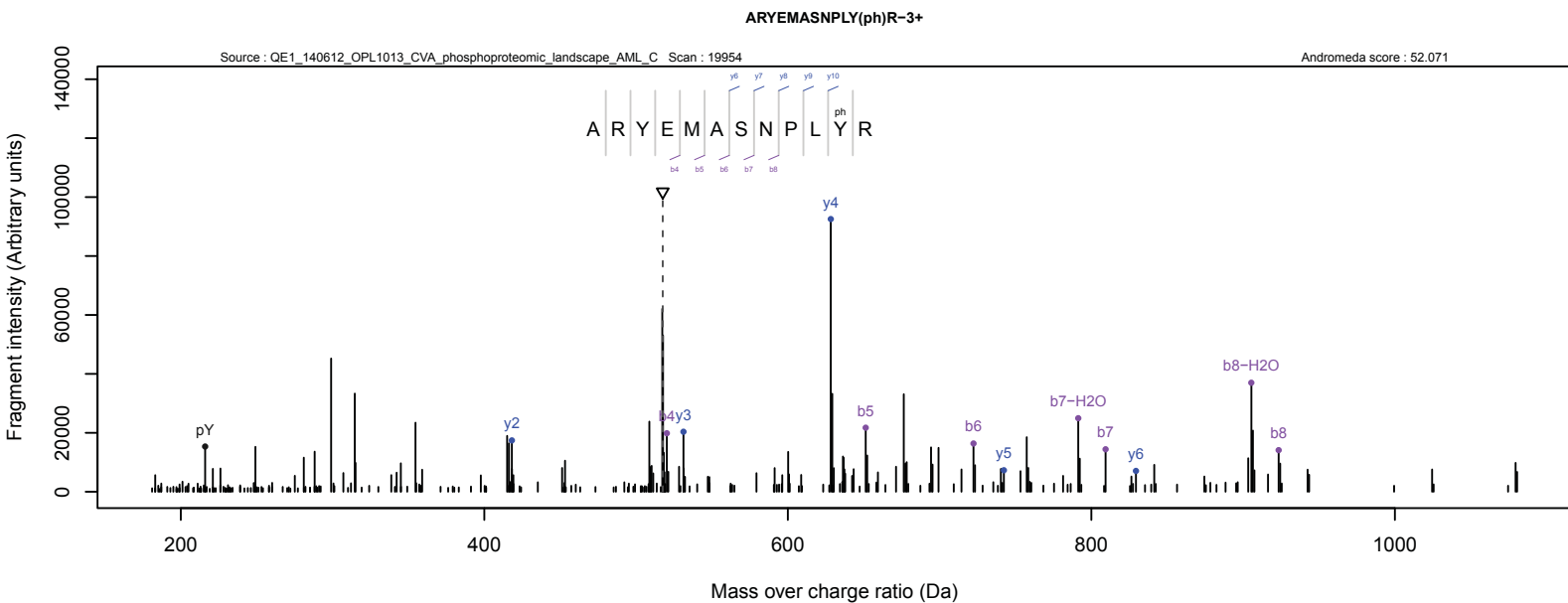


## VQIVY(ph)KPVDSLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22574

Andromeda score : 52.101

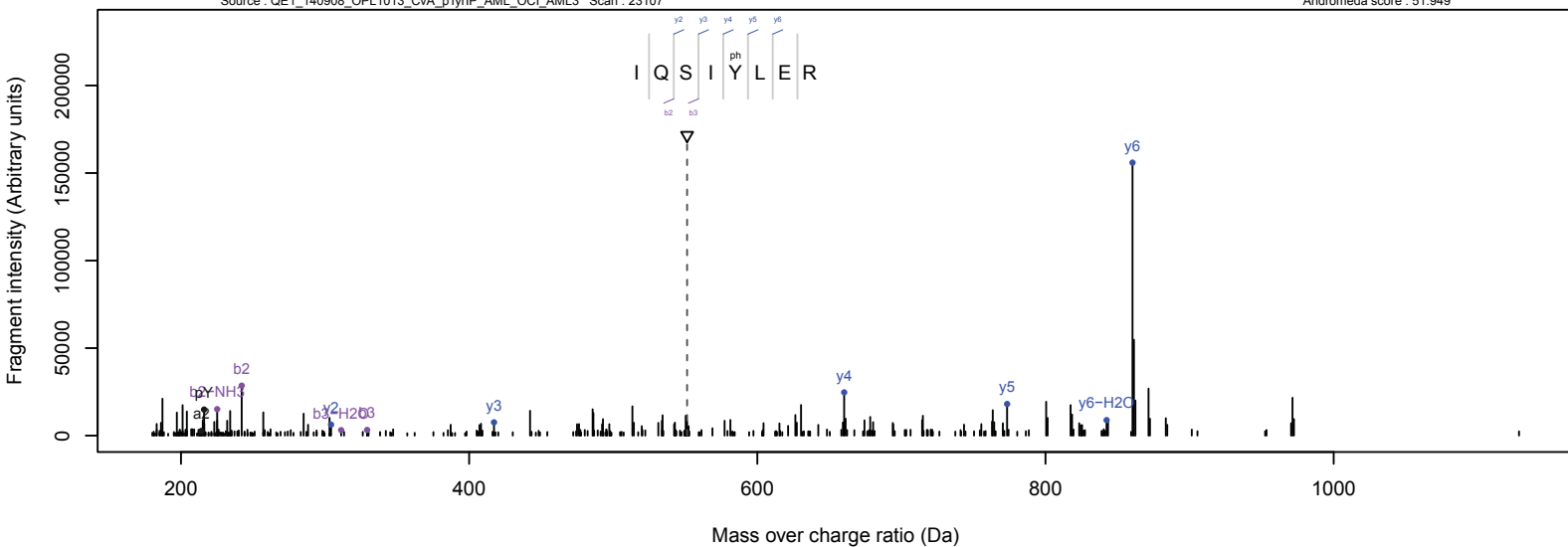




**IQSIY(ph)LER-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 23107

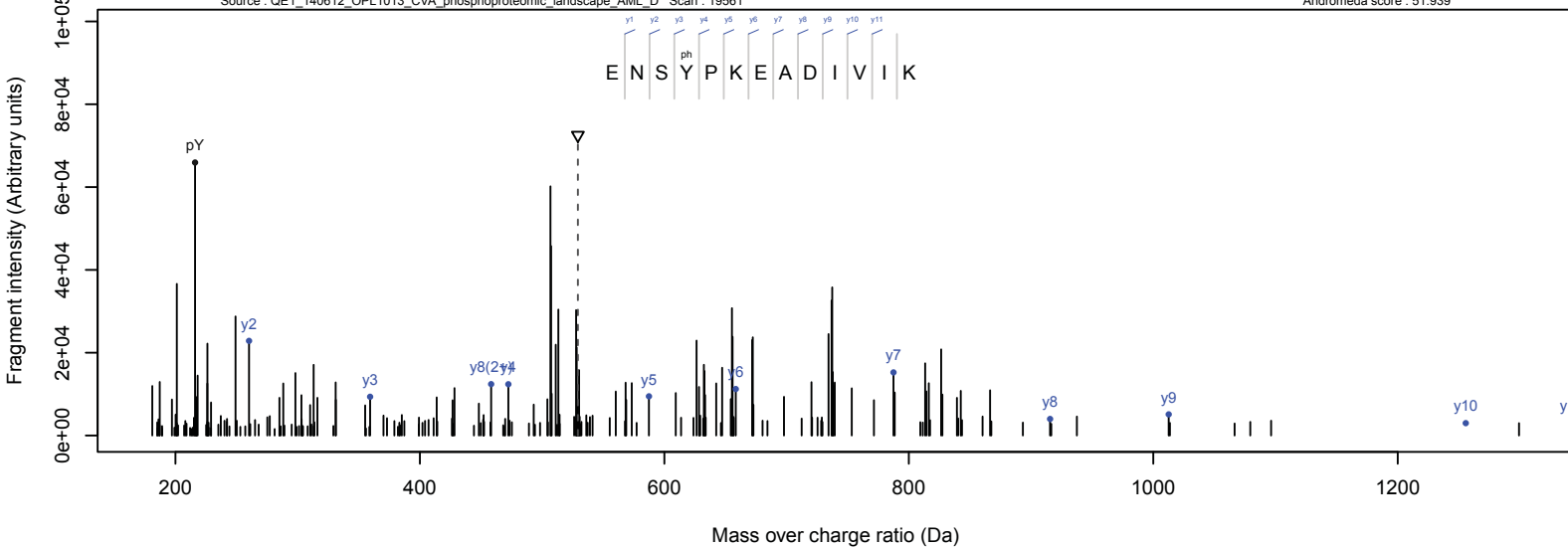
Andromeda score : 51.949



**ENSY(ph)PKEADIVIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 19561

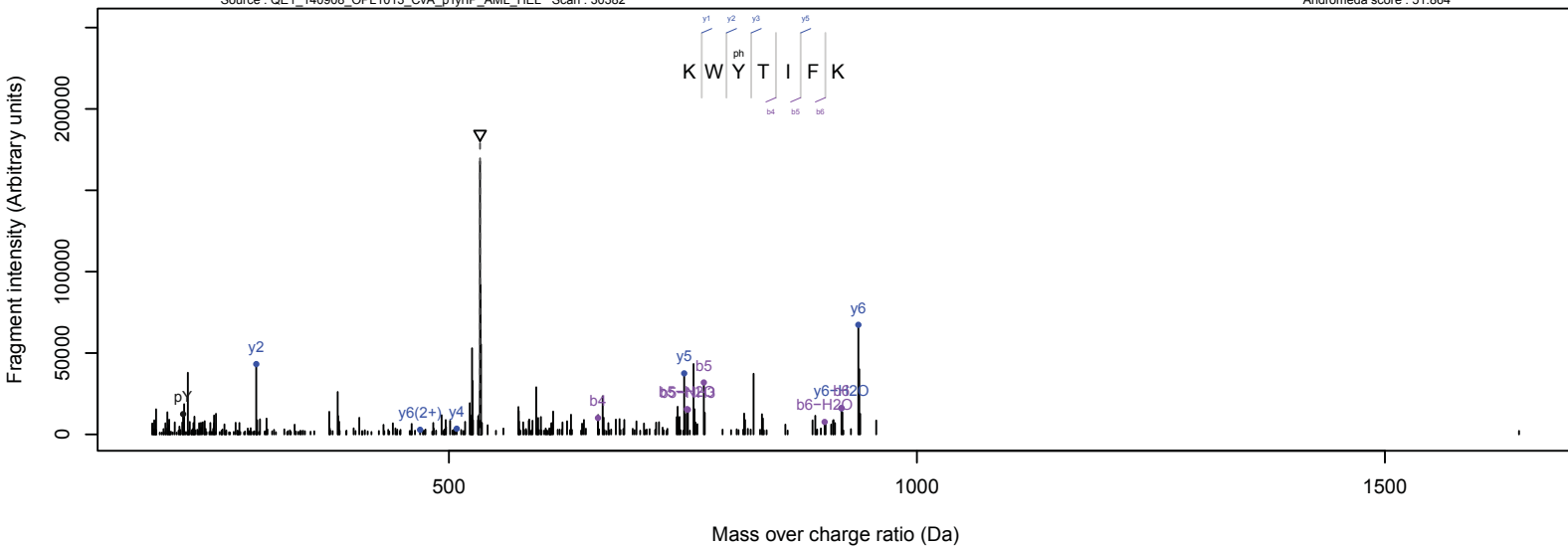
Andromeda score : 51.939



**KWY(ph)TIFK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 30382

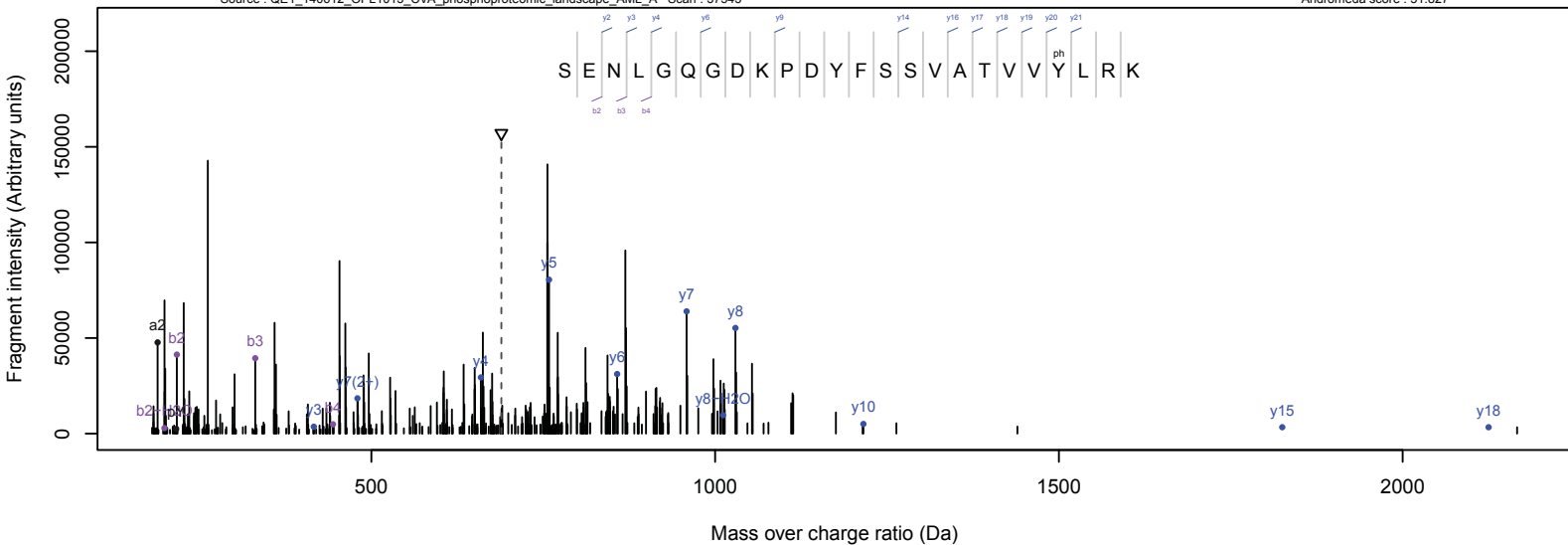
Andromeda score : 51.864



SENLGGDKPDYFSSVATVVYLRK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 37543

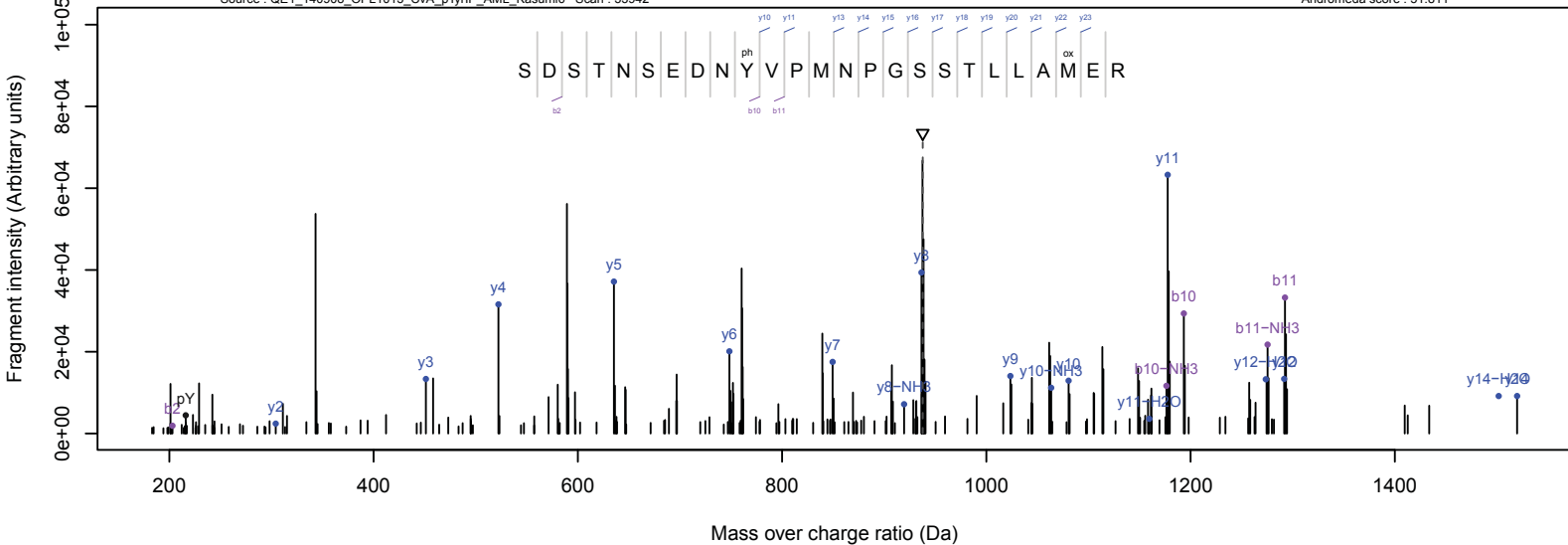
Andromeda score : 51.827



SDSTNSEDNYPMPNPGSSTLLAMER-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 33942

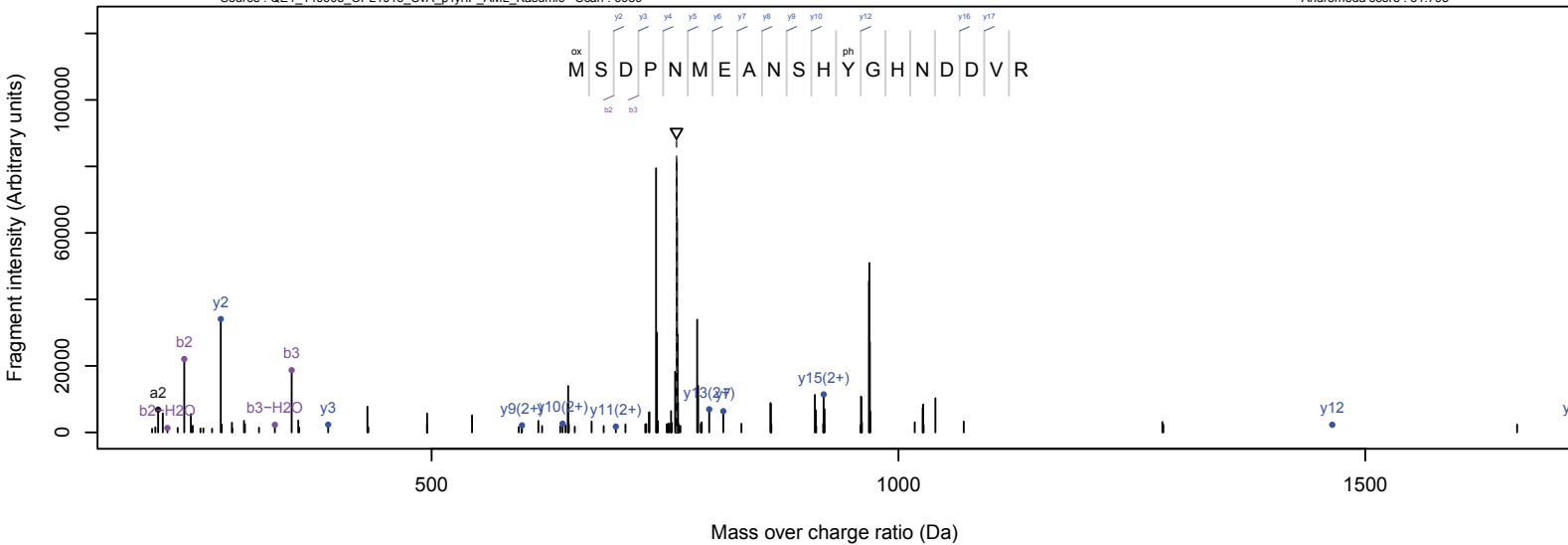
Andromeda score : 51.811



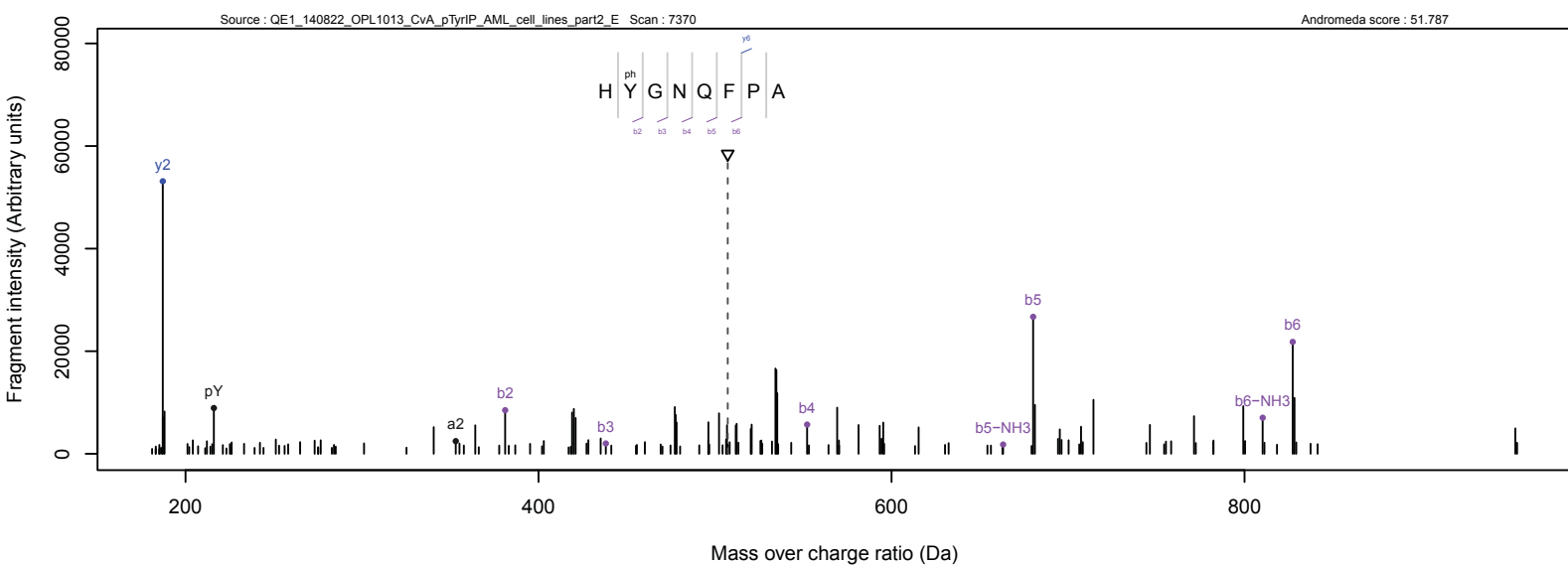
M(ox)SDPNMEANSHYGHNDVVR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 6989

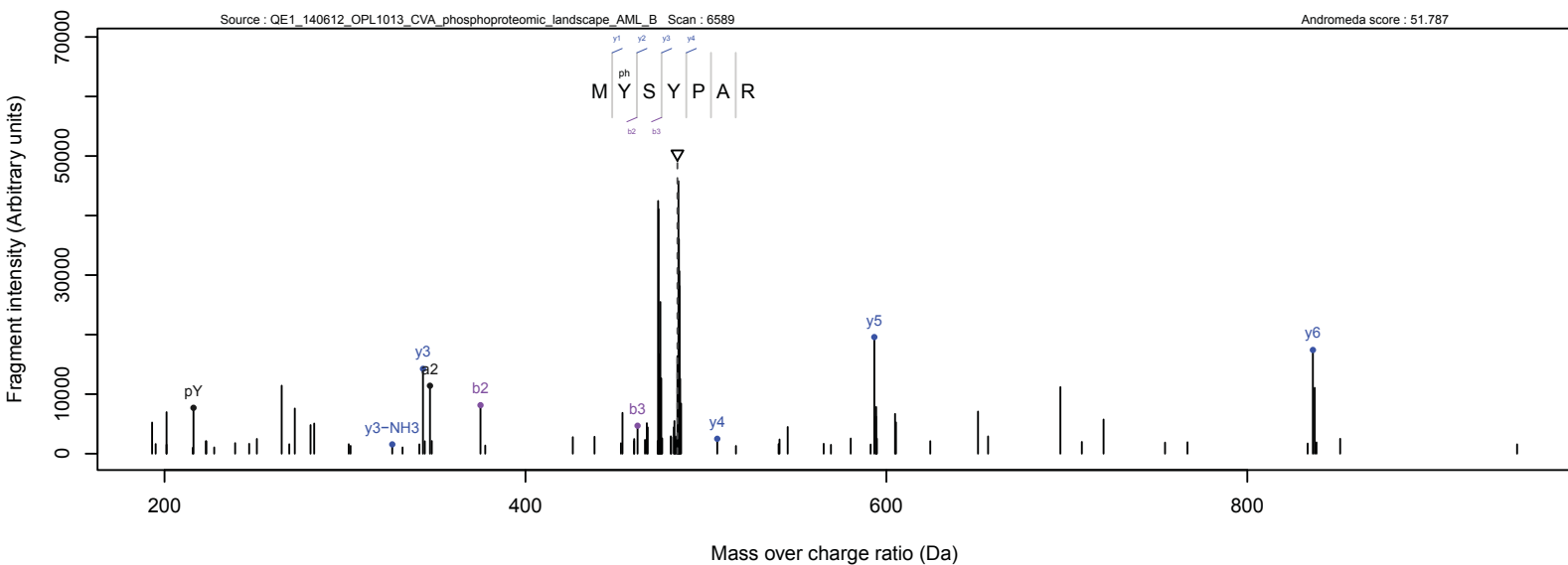
Andromeda score : 51.798



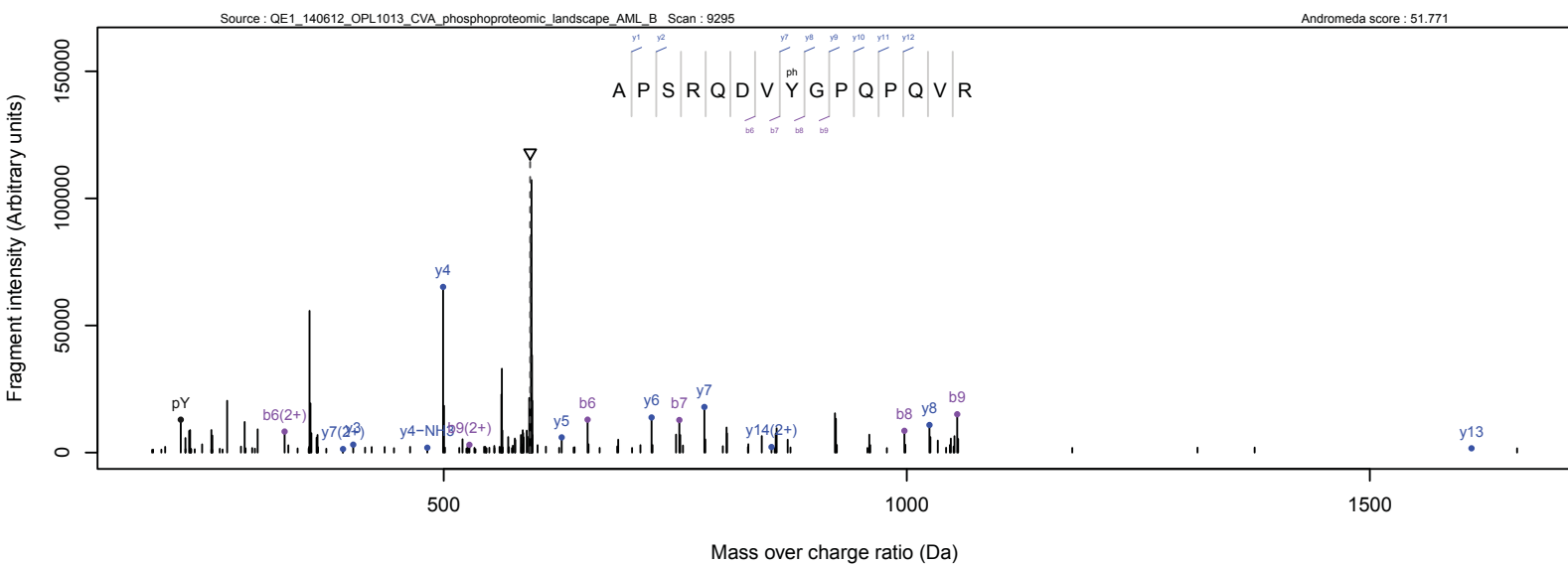
## HY(ph)GNQFPA-2+



## MY(ph)SYPAR-2+



## APSRQDVY(ph)GPQQR-3+

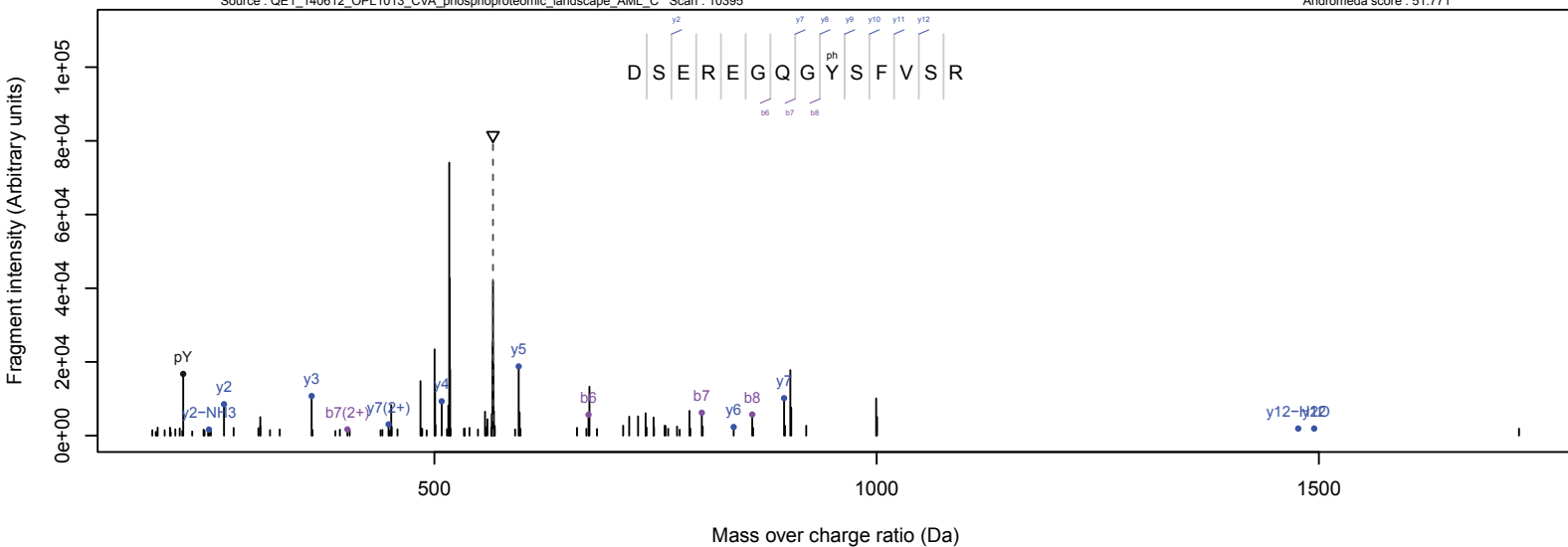




**DSEREGQGY(ph)SFVSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10395

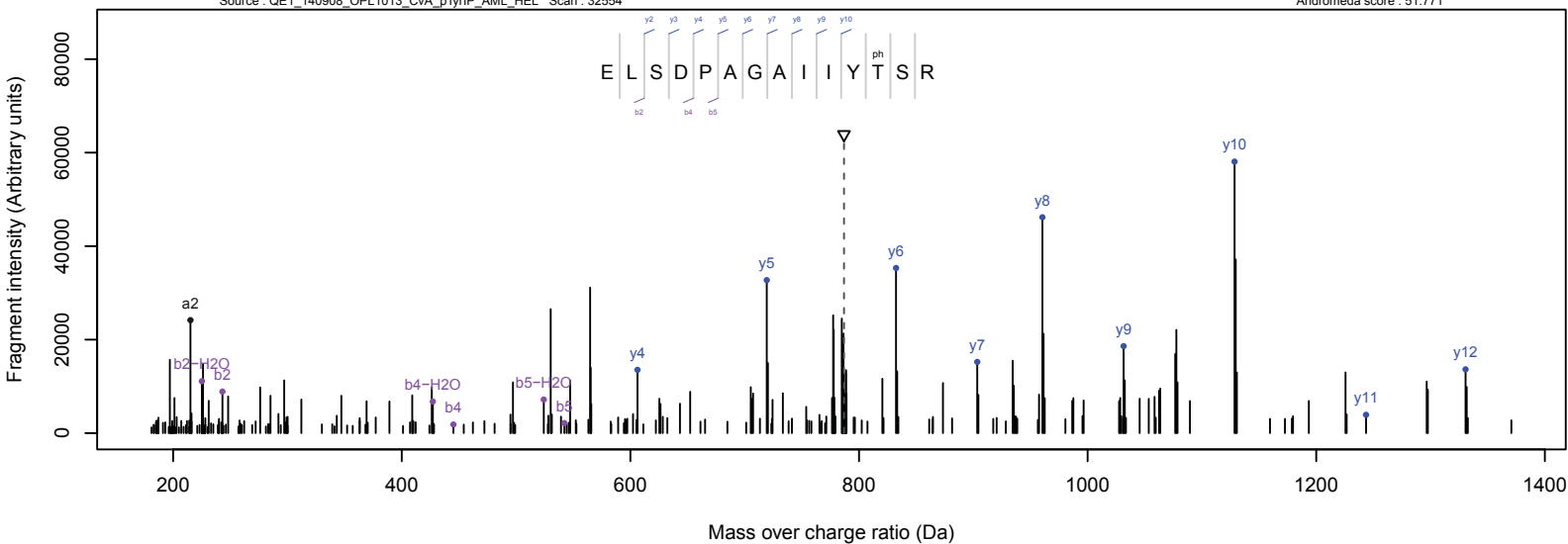
Andromeda score : 51.771



**ELSDPAGAIYT(ph)SR-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32554

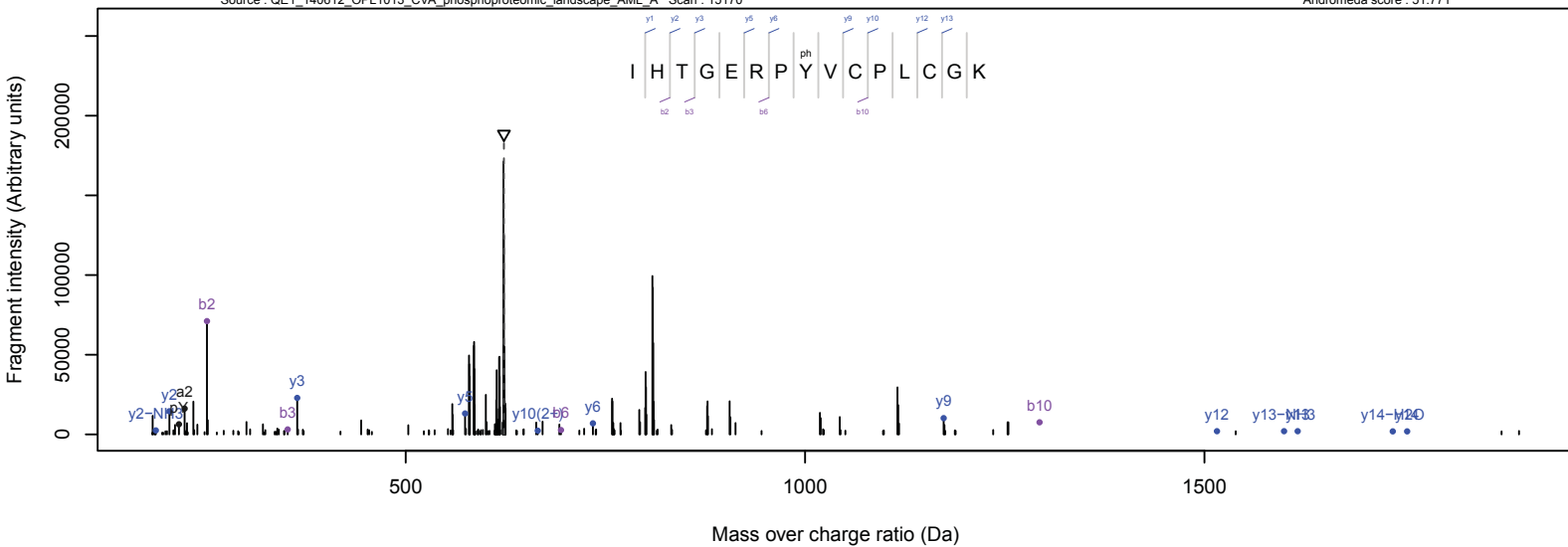
Andromeda score : 51.771



**IHTGERPY(ph)VCPLGK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 15170

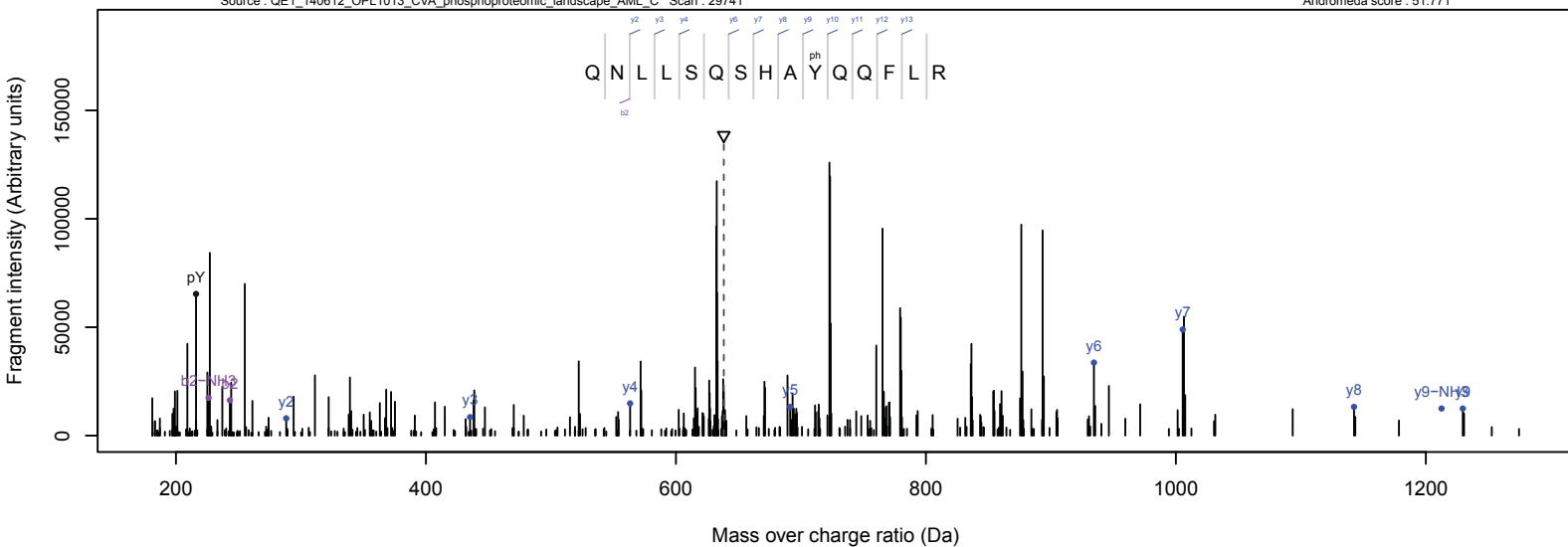
Andromeda score : 51.771



## QNLLSQSHAY(ph)QQFLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 29741

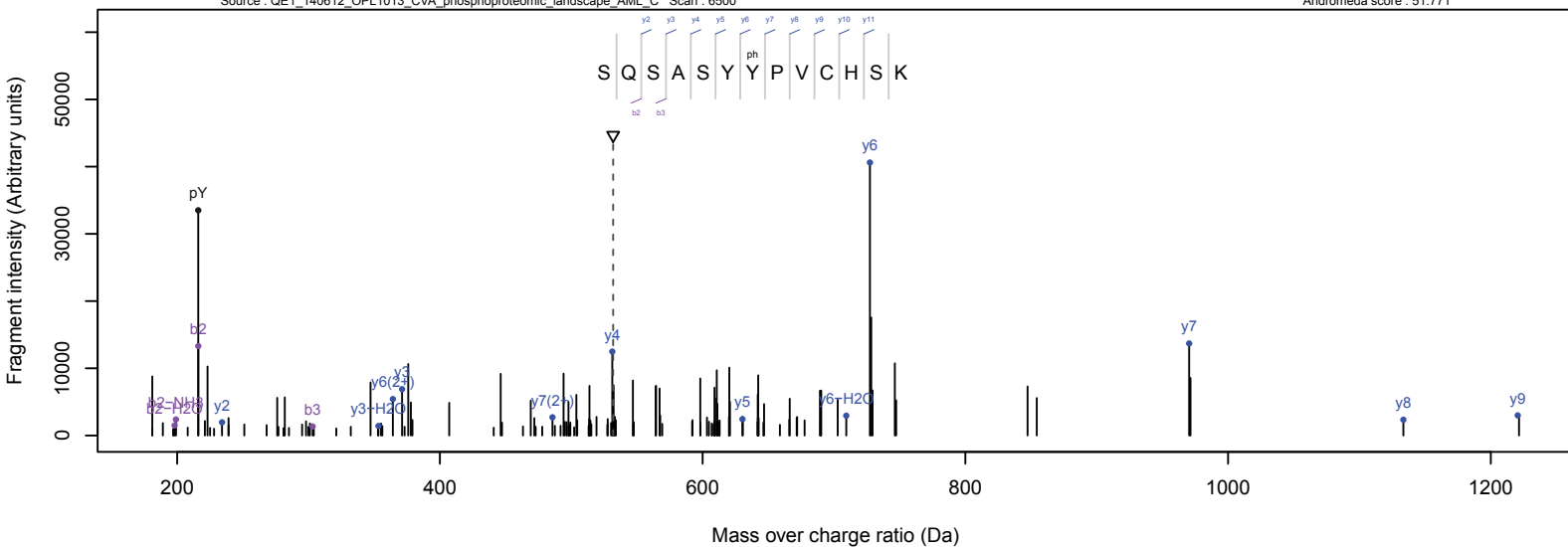
Andromeda score: 51.771



## SQSASY(ph)PVCHSK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 6500

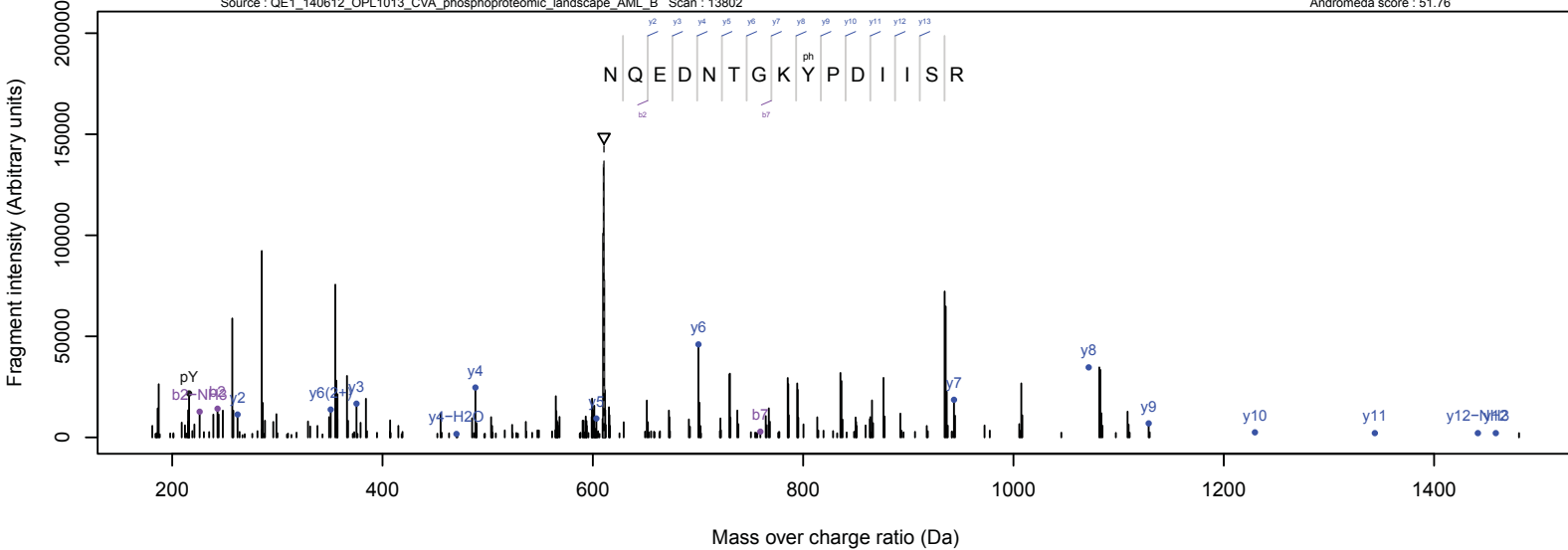
Andromeda score: 51.771



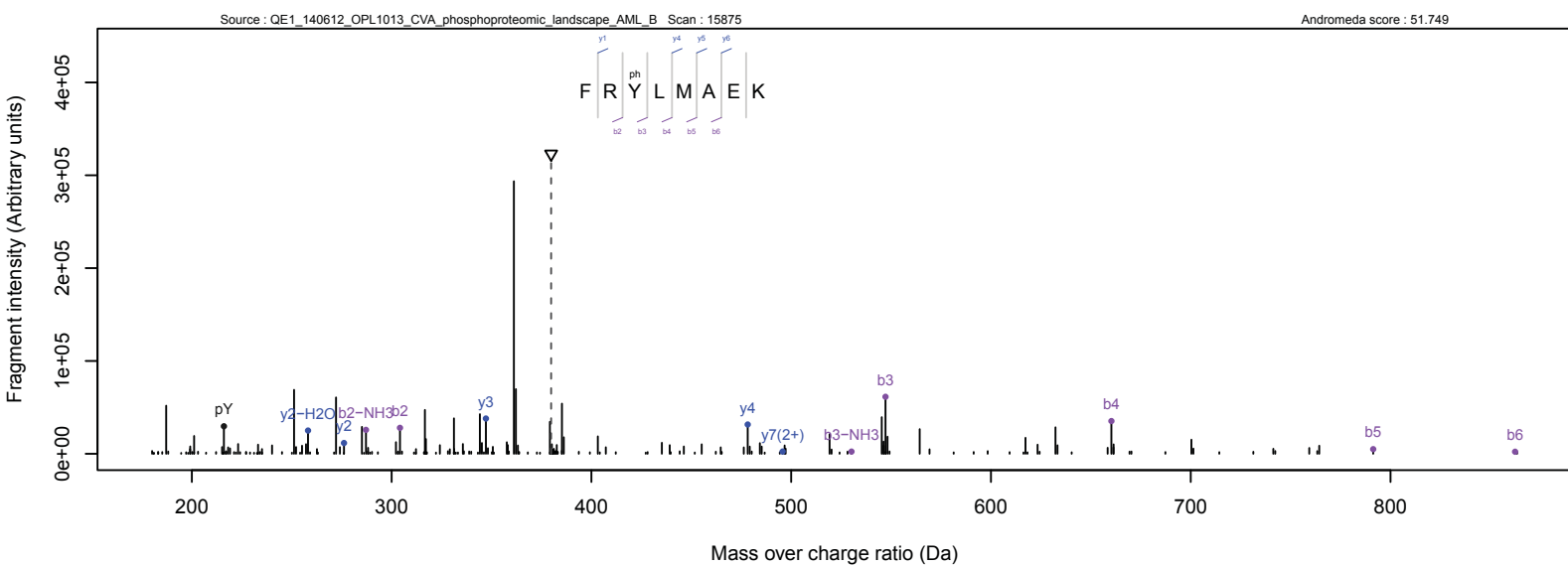
## NQEDNTGKY(ph)PDIISR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 13802

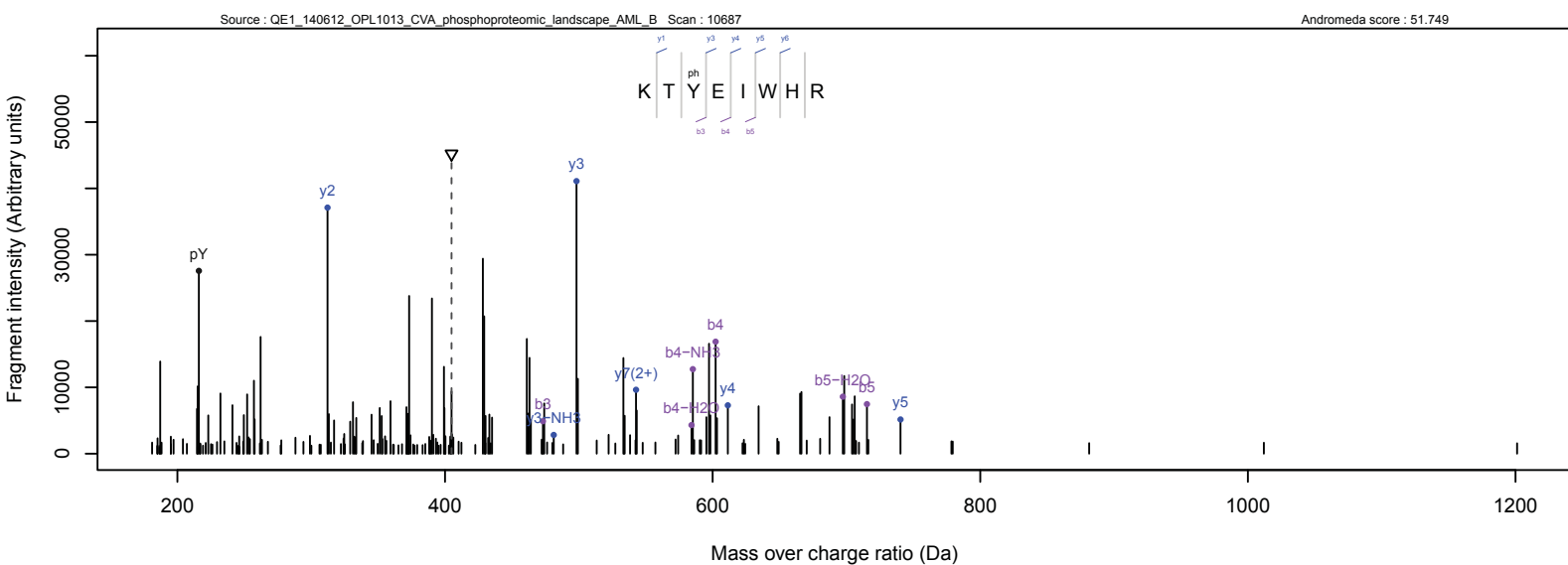
Andromeda score: 51.76



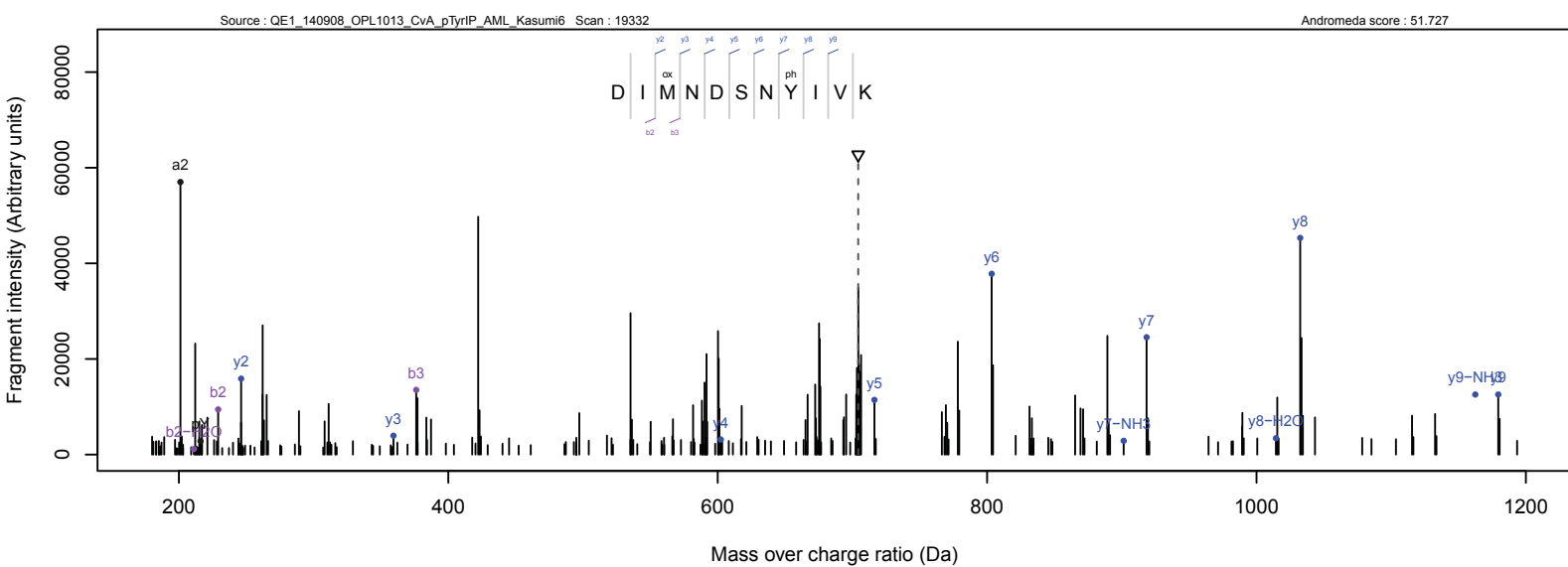
## FRY(ph)LMAEK-3+



## KTY(ph)EIWHR-3+



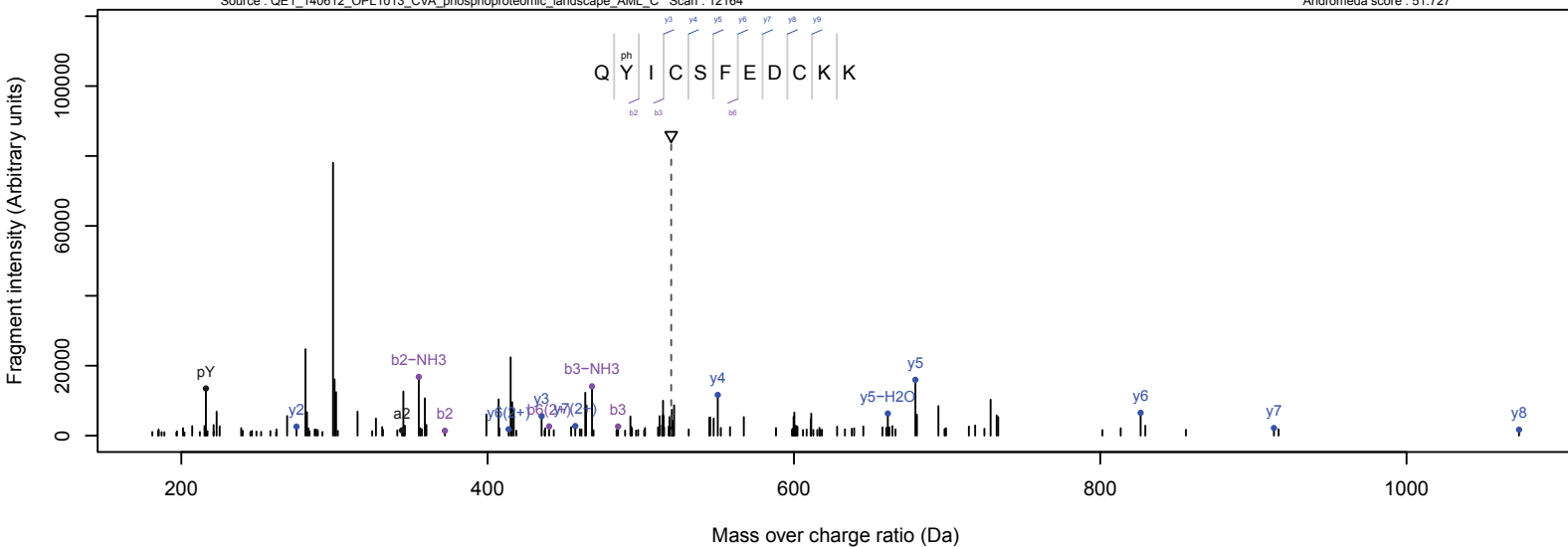
## DIM(ox)NDSNY(ph)IVK-2+



## QY(ph)CSFEDCKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12164

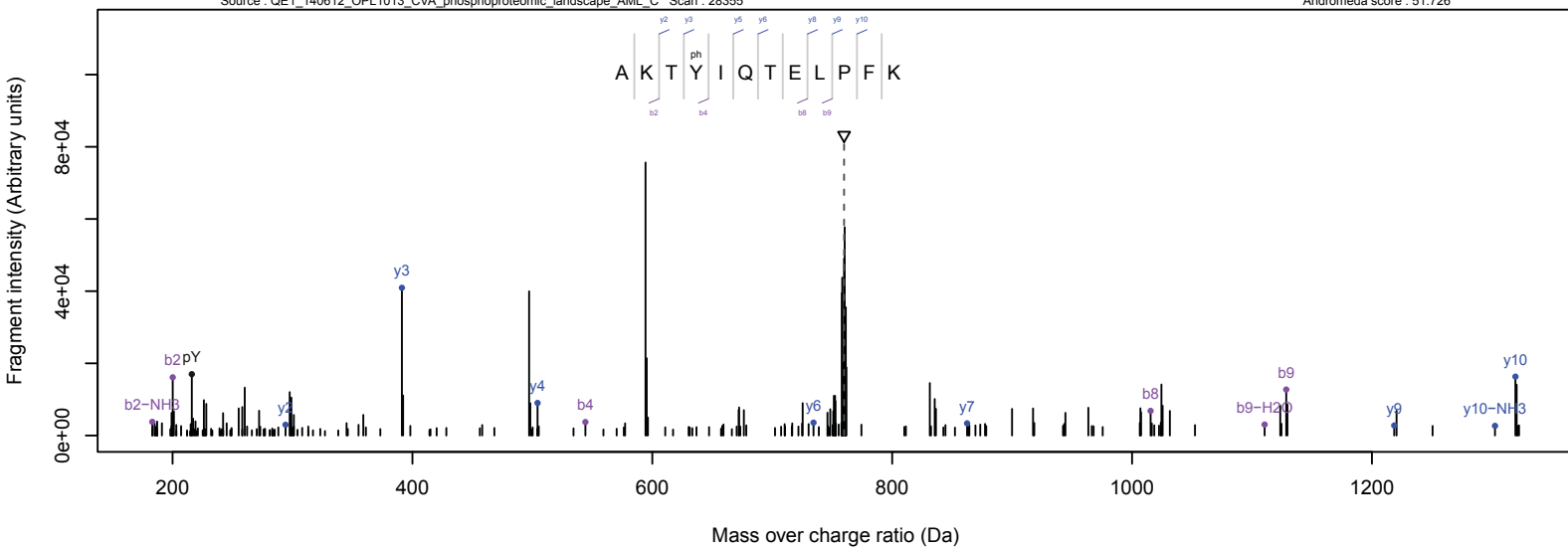
Andromeda score : 51.727



## AKTY(ph)QTELPFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28355

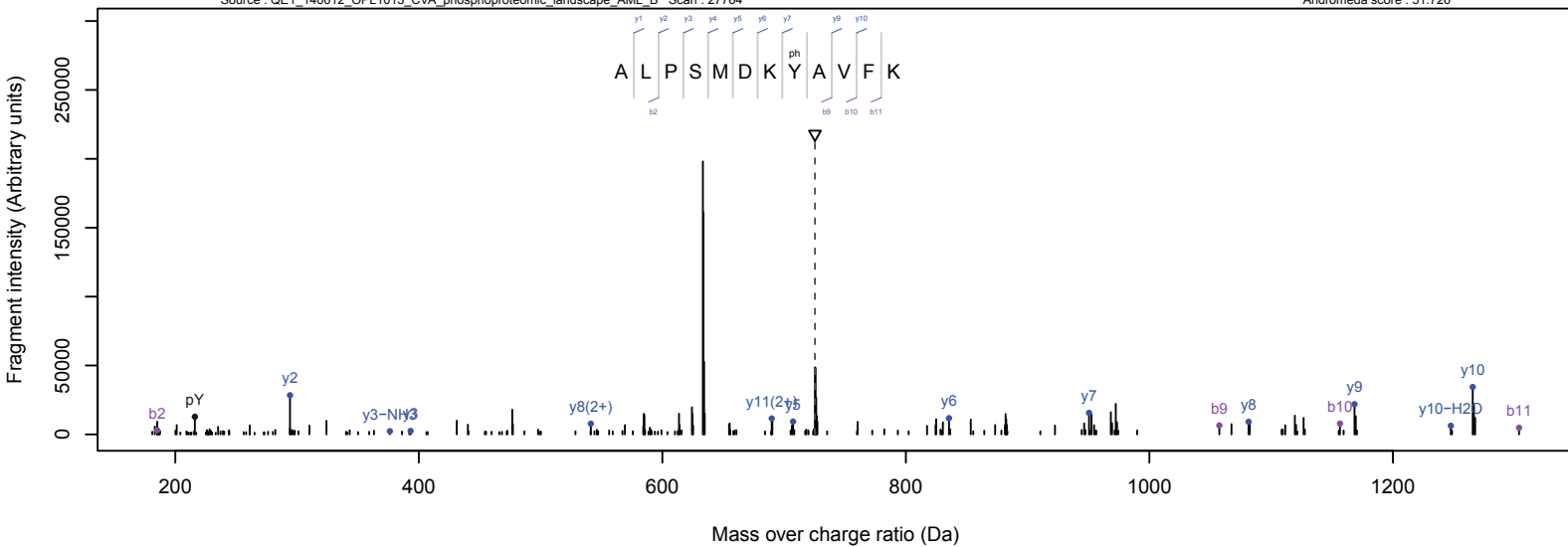
Andromeda score : 51.726



## ALPSMDKY(ph)AVFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27764

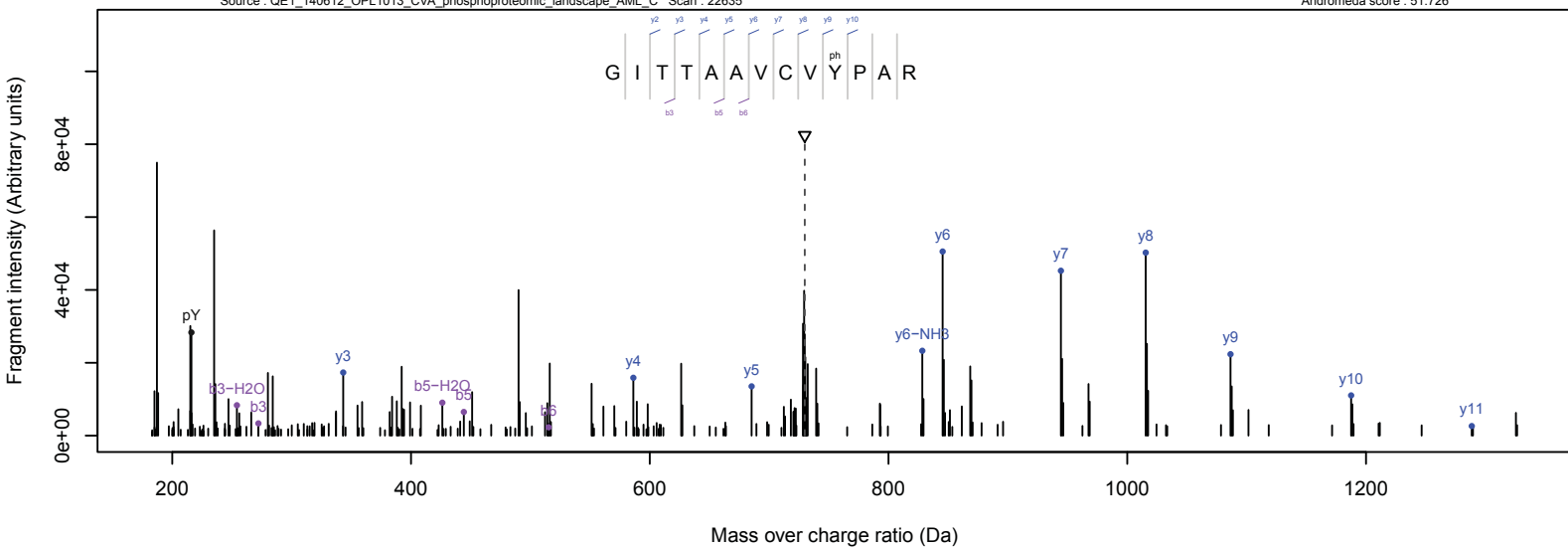
Andromeda score : 51.726



## GITTAACVY(ph)PAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22635

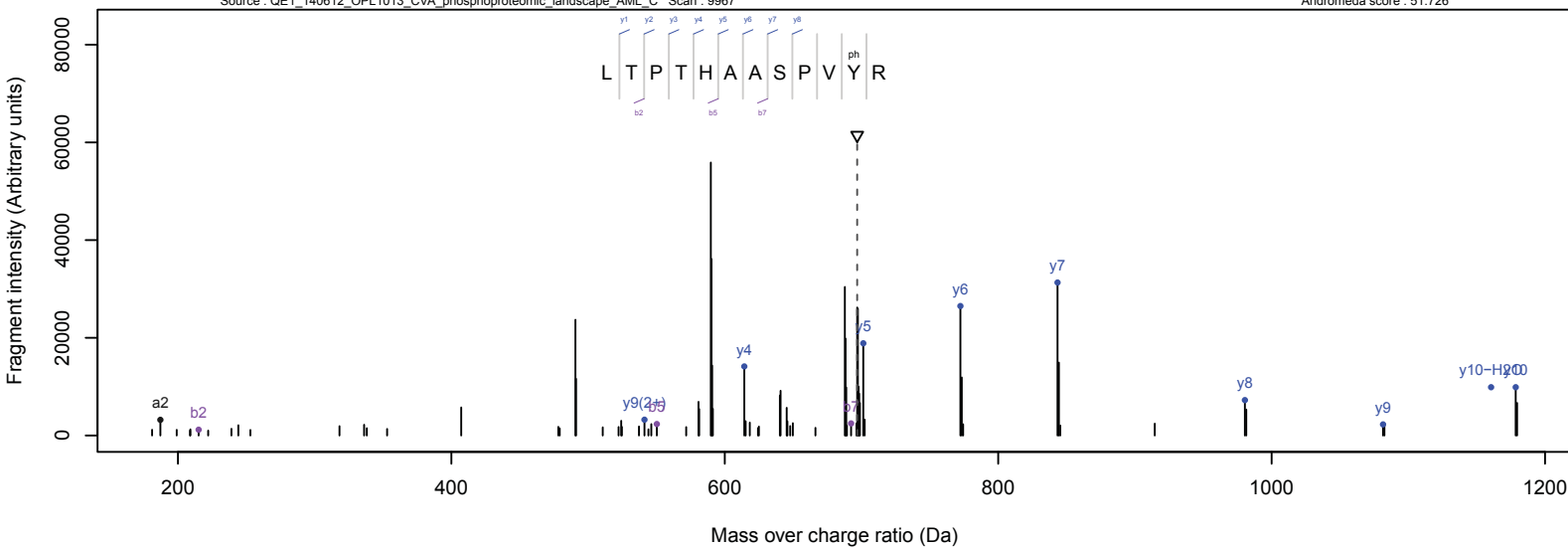
Andromeda score : 51.726



## LTPTHAASPVY(ph)R-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9967

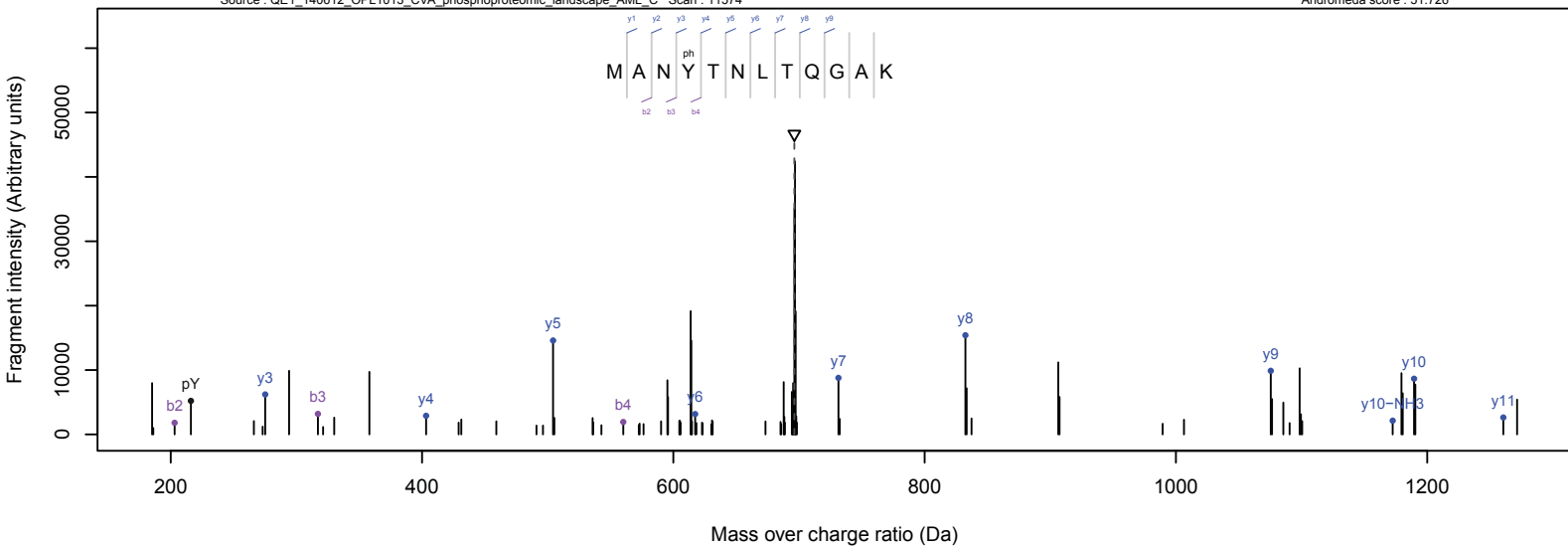
Andromeda score : 51.726



## MANY(ph)TNLTQGAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11574

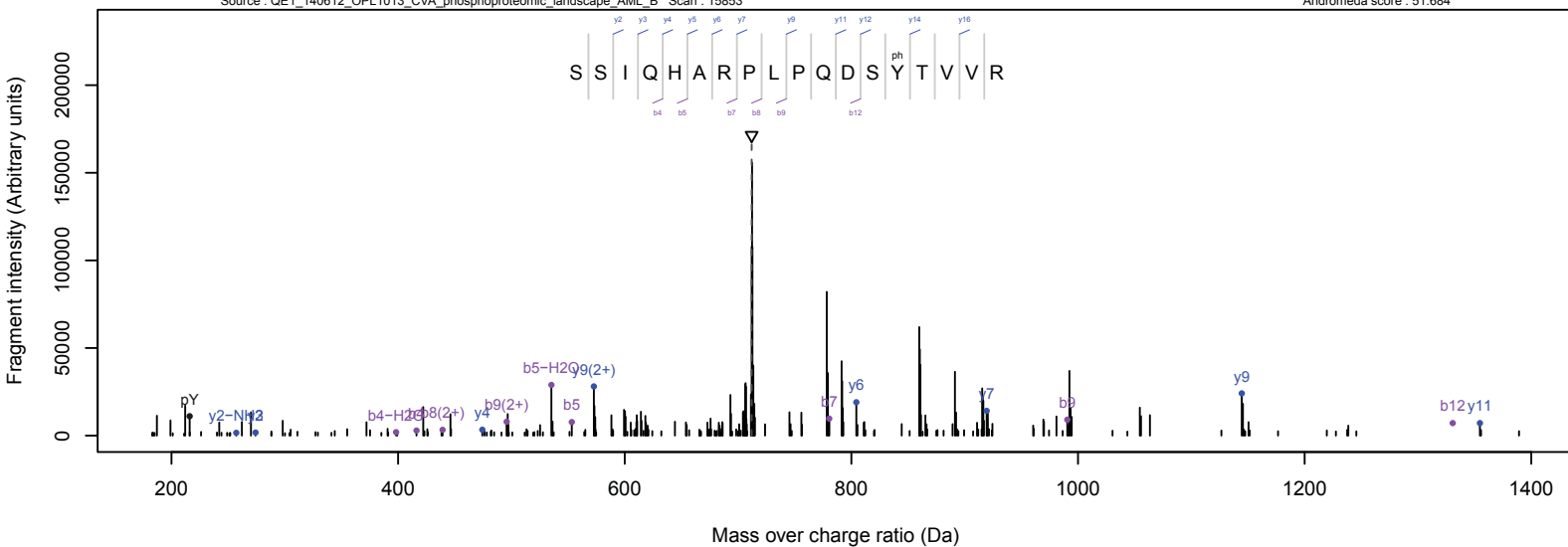
Andromeda score : 51.726



## SSIQHARLPQDSY(ph)TVVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15853

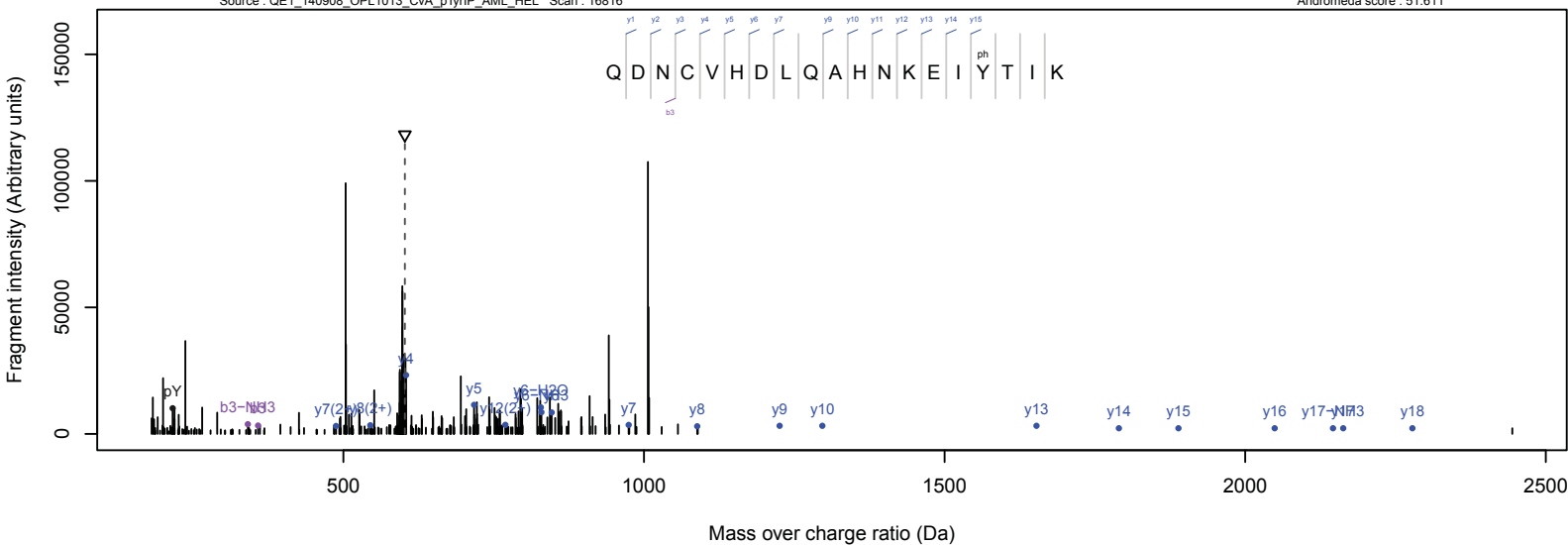
Andromeda score : 51.684



## QDNCVHDLQAHNKEIY(ph)TIK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 16816

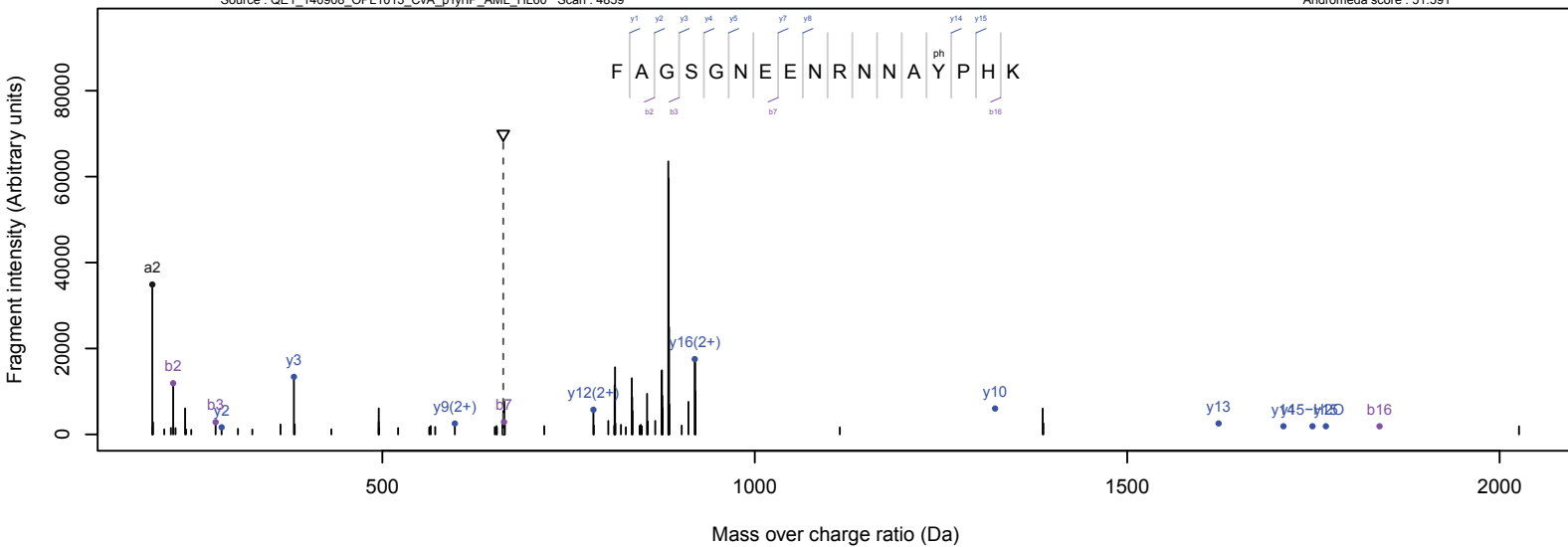
Andromeda score : 51.611



## FAGSGNEENNAY(ph)PHK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 4859

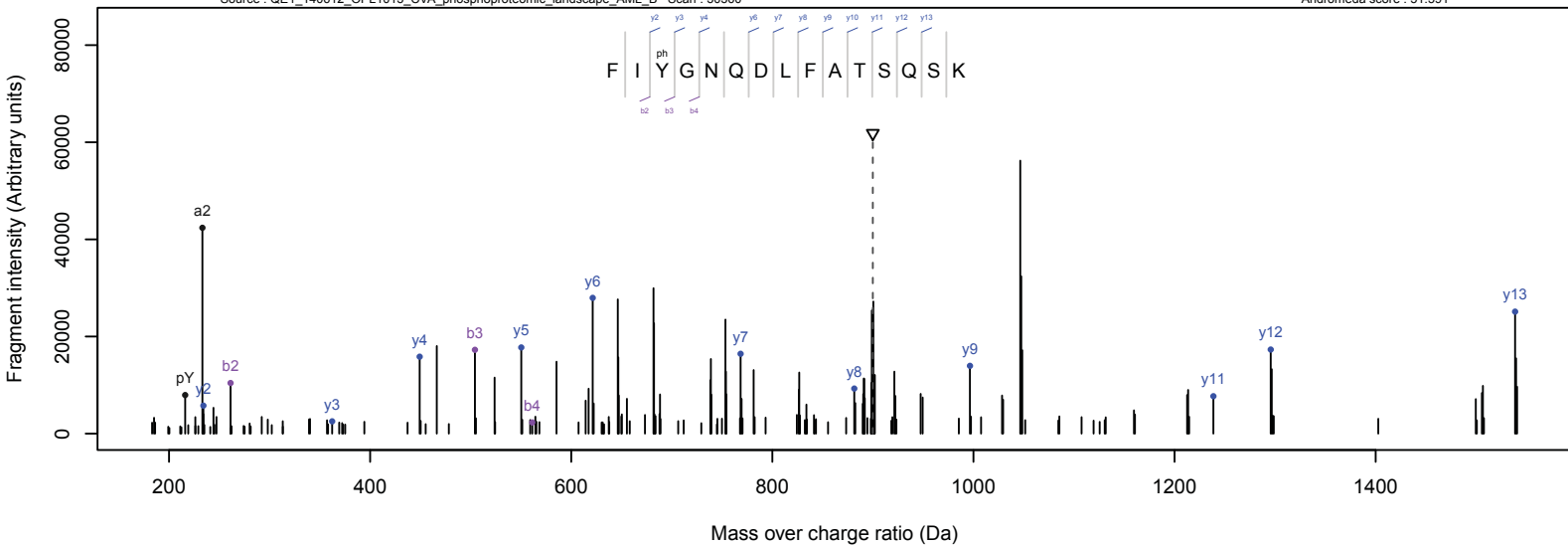
Andromeda score : 51.591



## FIY(ph)GNQLFATSQSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30360

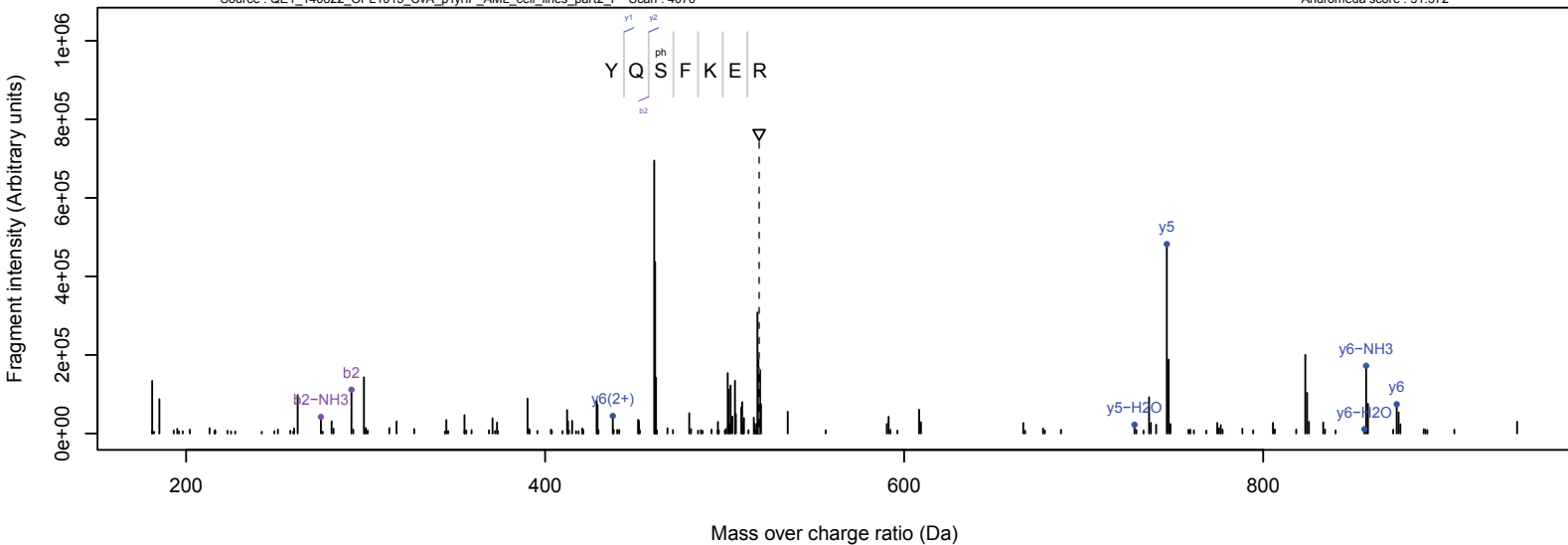
Andromeda score : 51.591



## YQS(ph)FKER-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 4070

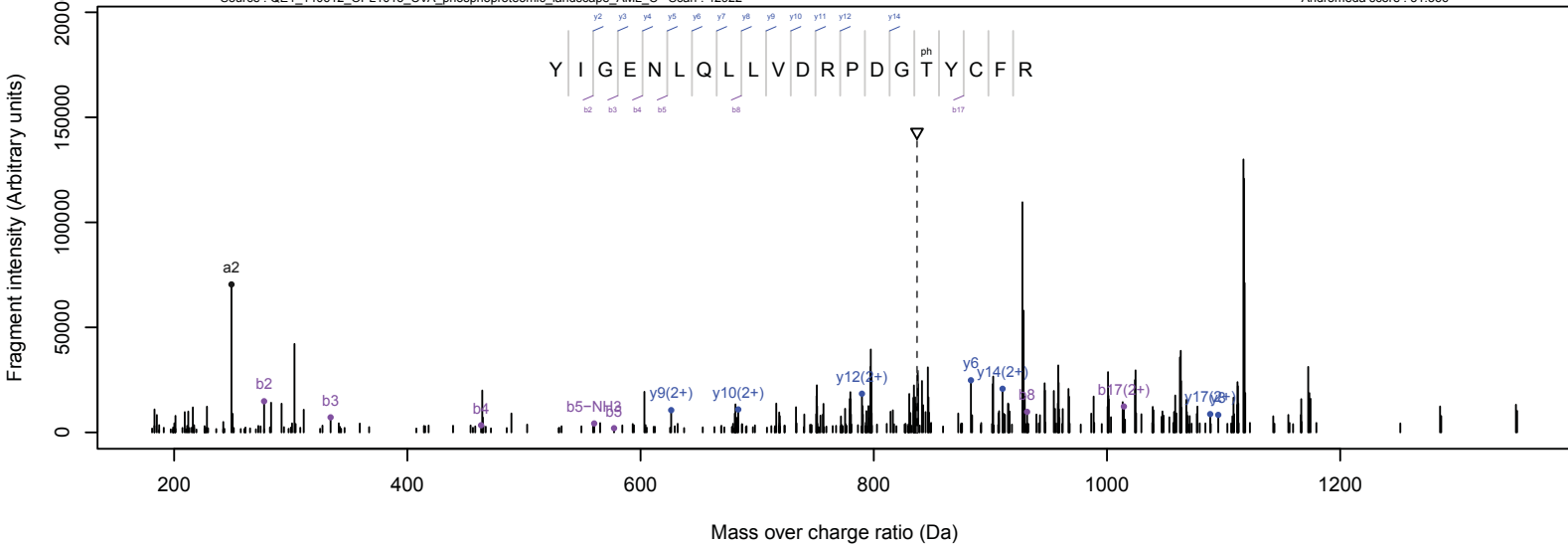
Andromeda score : 51.572



## YIGENLQLLVDRPDGT(ph)YCFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42922

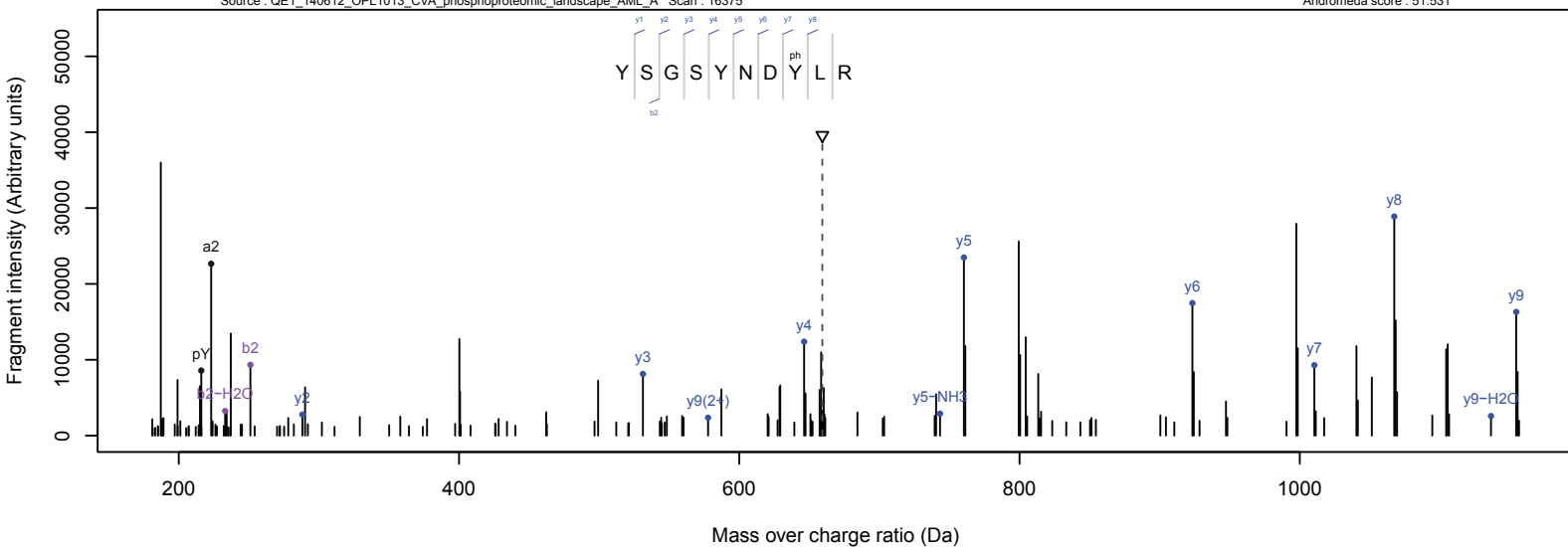
Andromeda score : 51.566



## YSGSYNDY(ph)LR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 16375

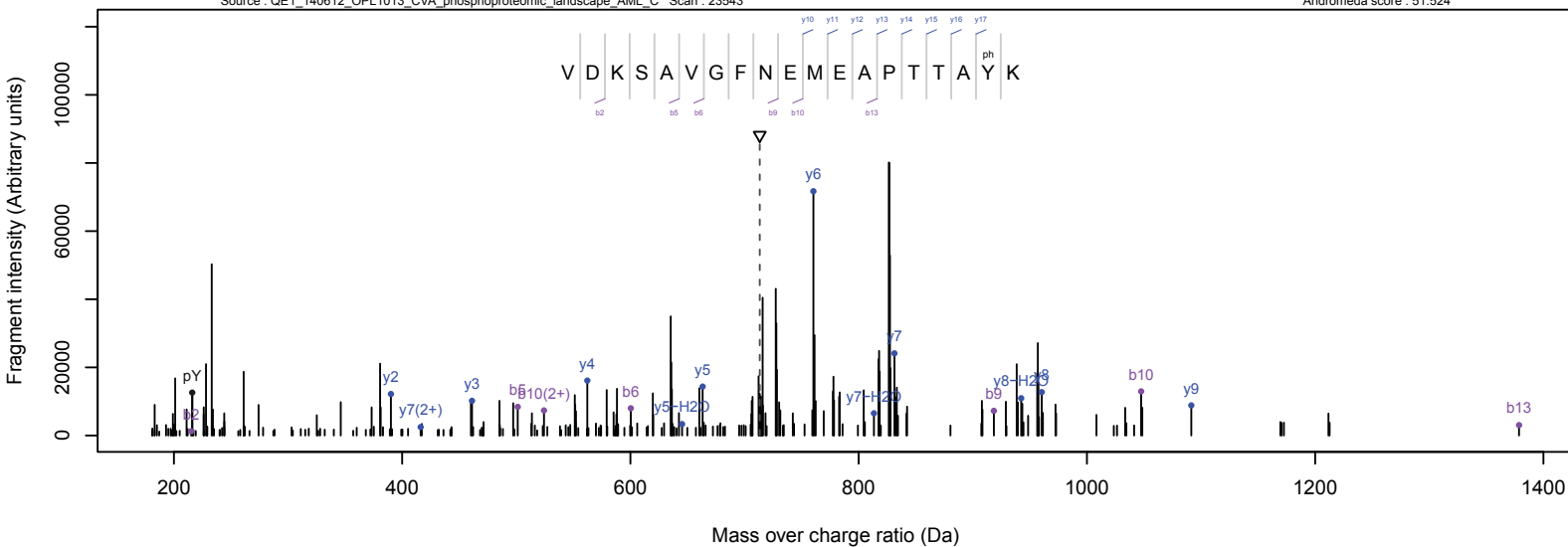
Andromeda score : 51.531



## VDKSAVGFNEMEAPTAYK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23543

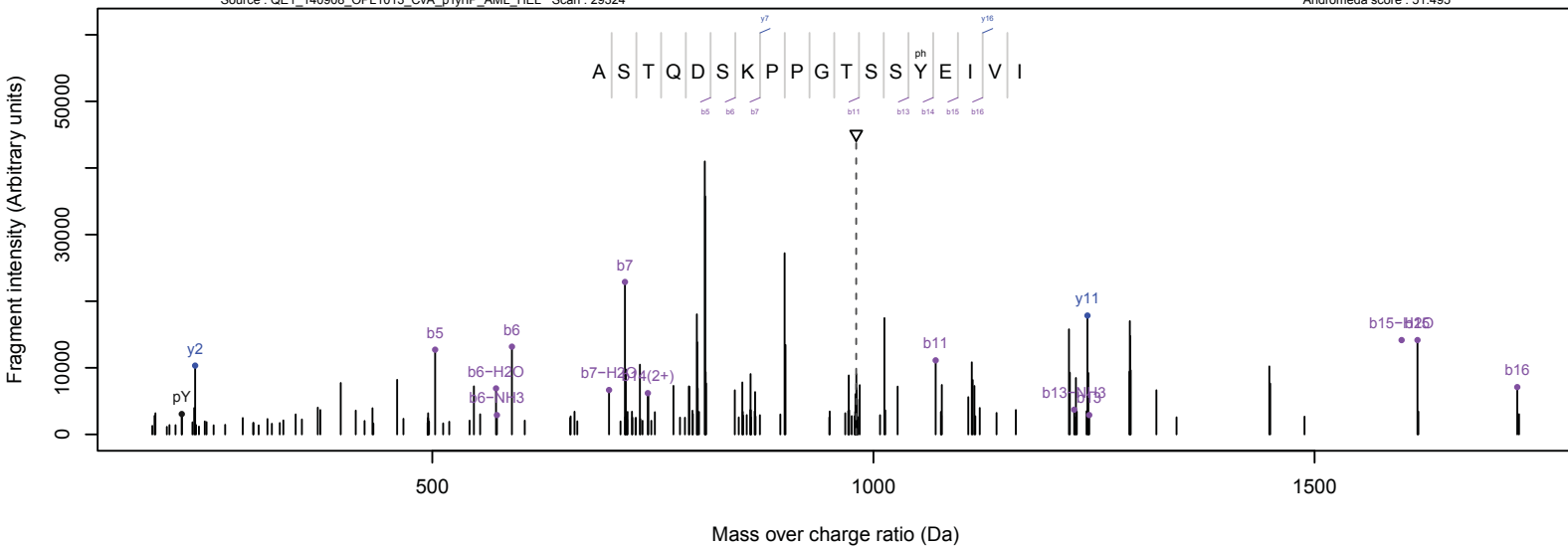
Andromeda score : 51.524



## ASTQSKPPGTSSY(ph)EIVI-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29324

Andromeda score : 51.495



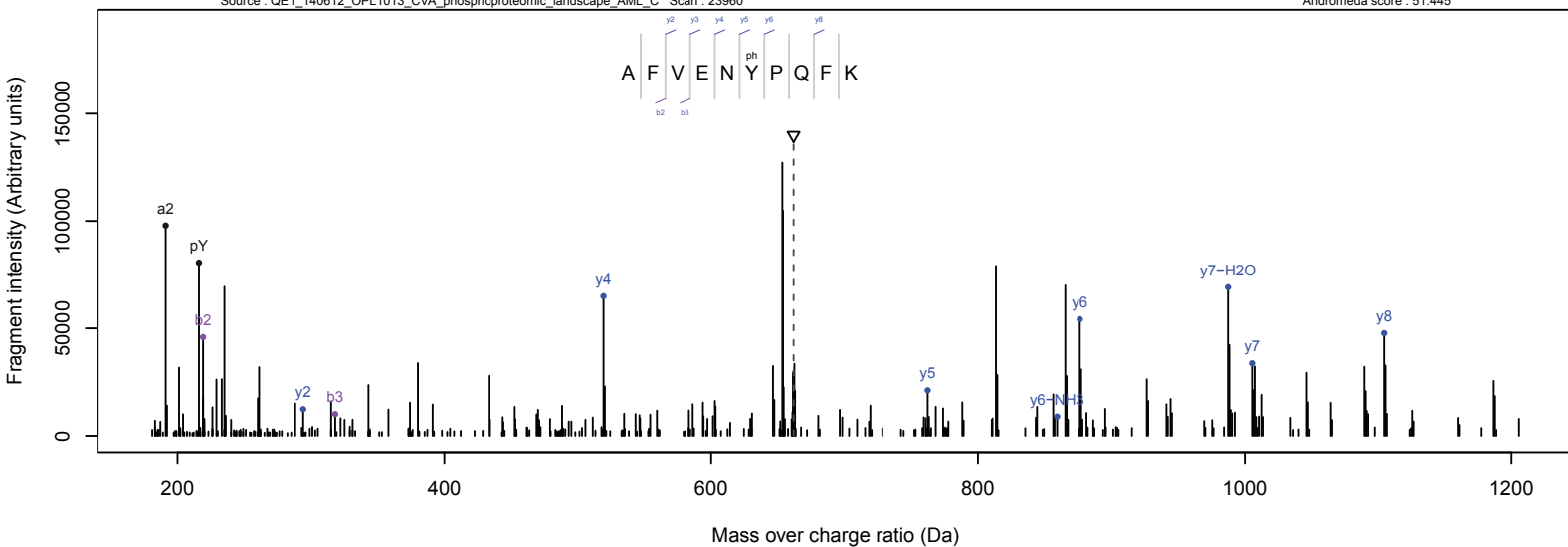




## AFVENVY(ph)PQFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23980

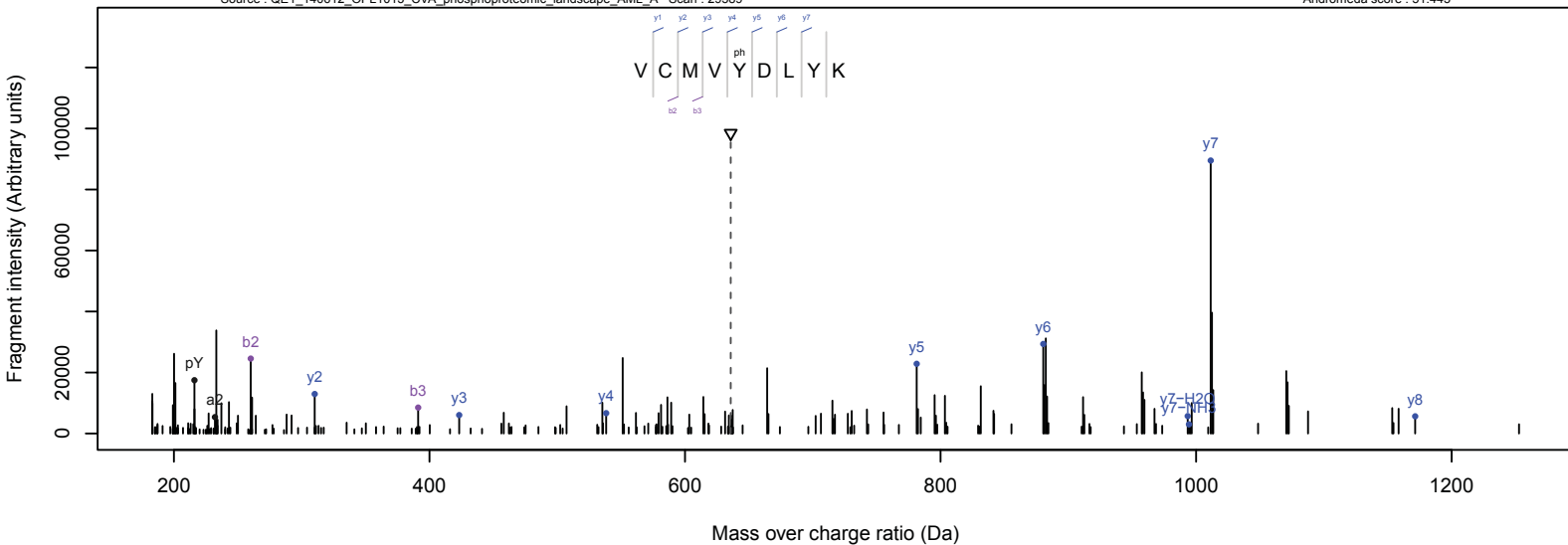
Andromeda score : 51.445



## VCMVY(ph)DLYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29589

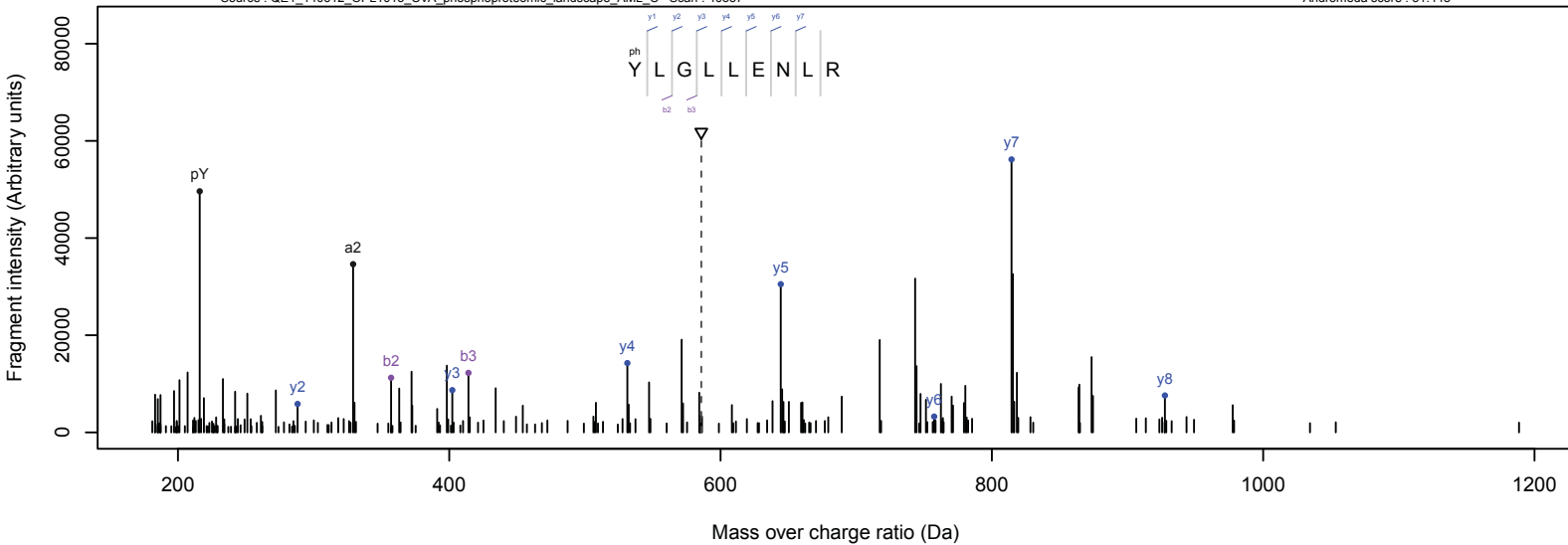
Andromeda score : 51.445



## Y(ph)LGLENLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40337

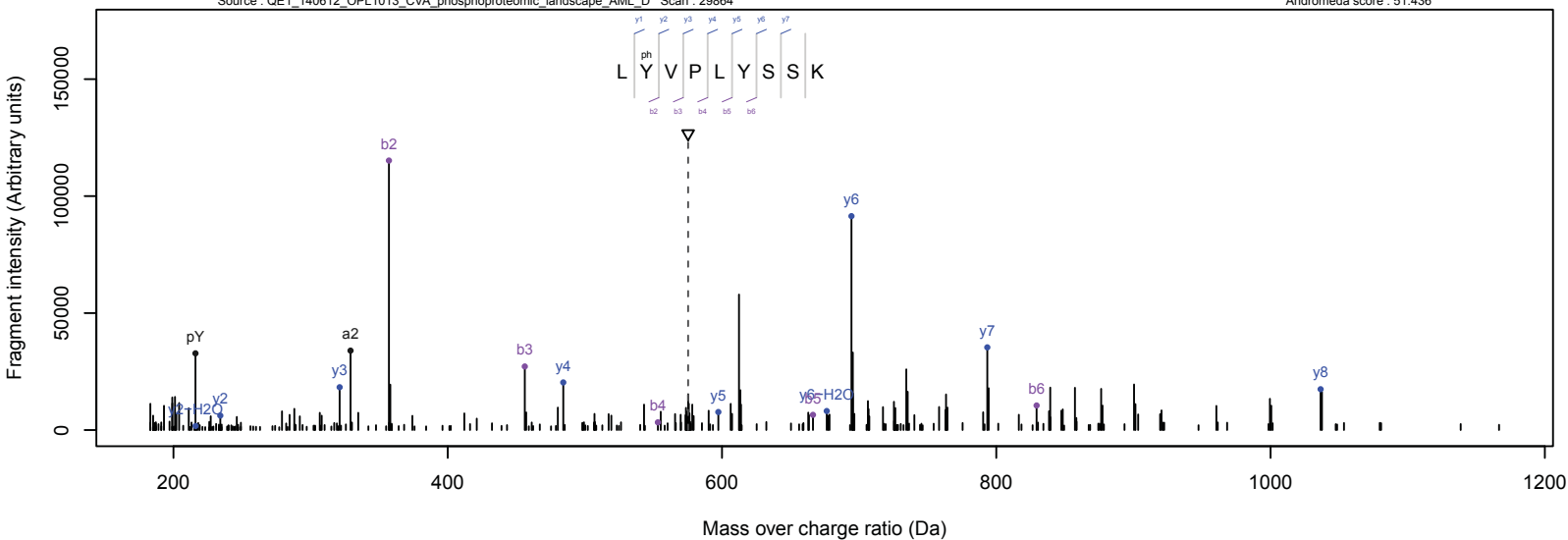
Andromeda score : 51.445



## LY(ph)VPLYSSK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 29884

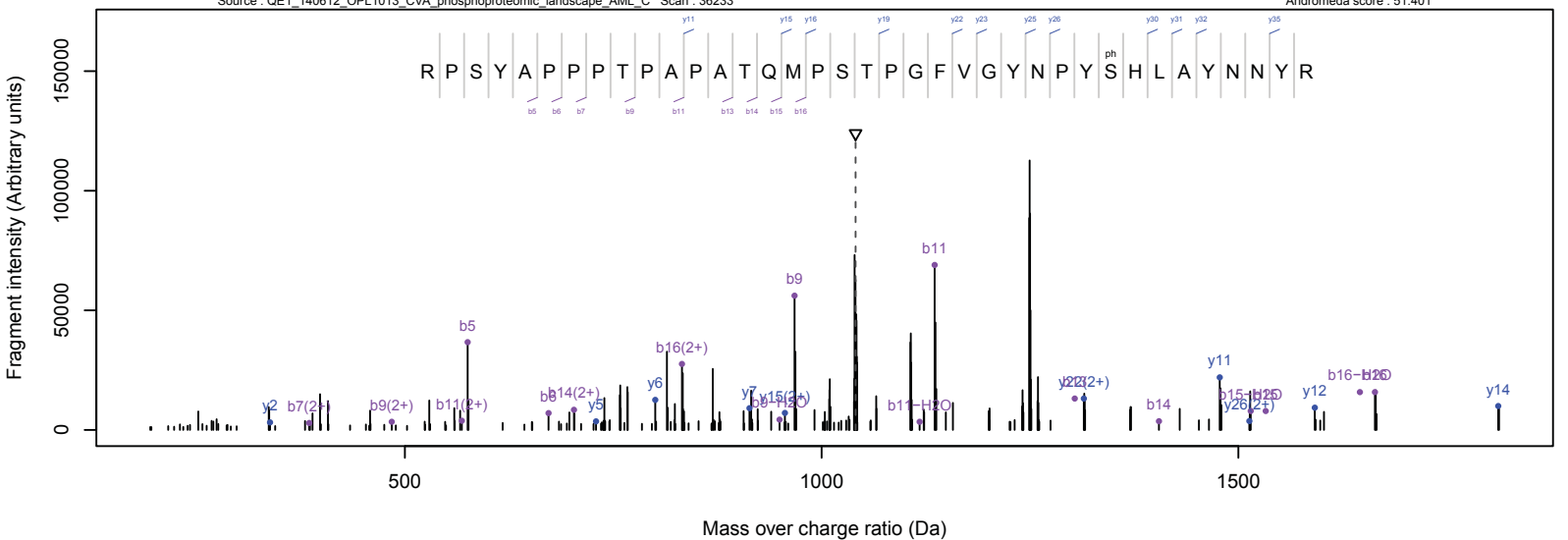
Andromeda score : 51.436



## RPSYAPPPTAPATQMPSTPGFVGYNPYS(ph)HLAYNNYR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36233

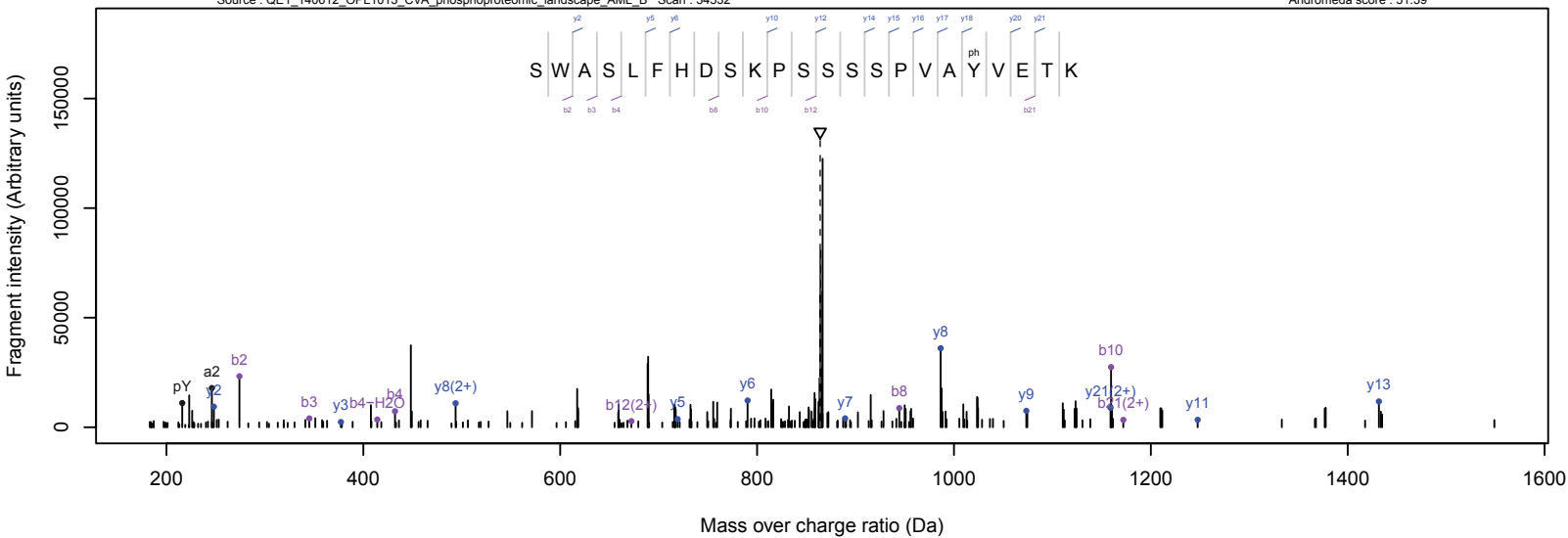
Andromeda score : 51.401



## SWASLFHDSKPSSSSPVAY(ph)VETK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 34532

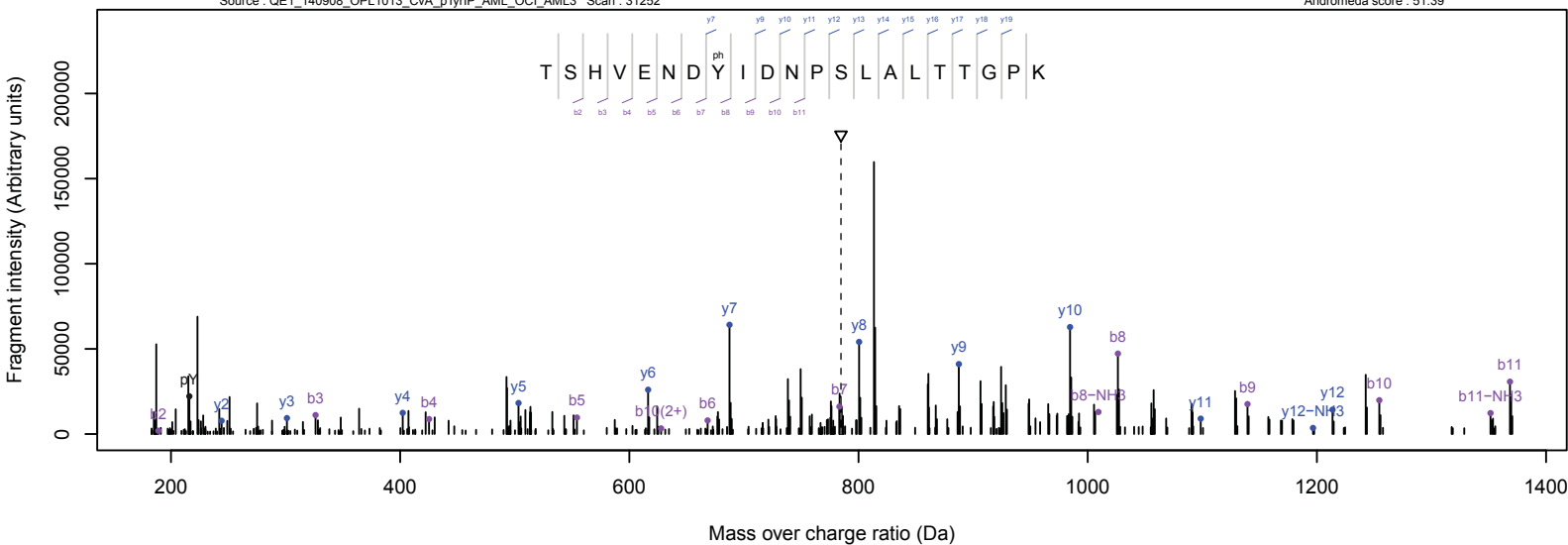
Andromeda score : 51.39



TSHVENDY(ph)IDNPSLALTTGPK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_OCI\_AML3 Scan : 31252

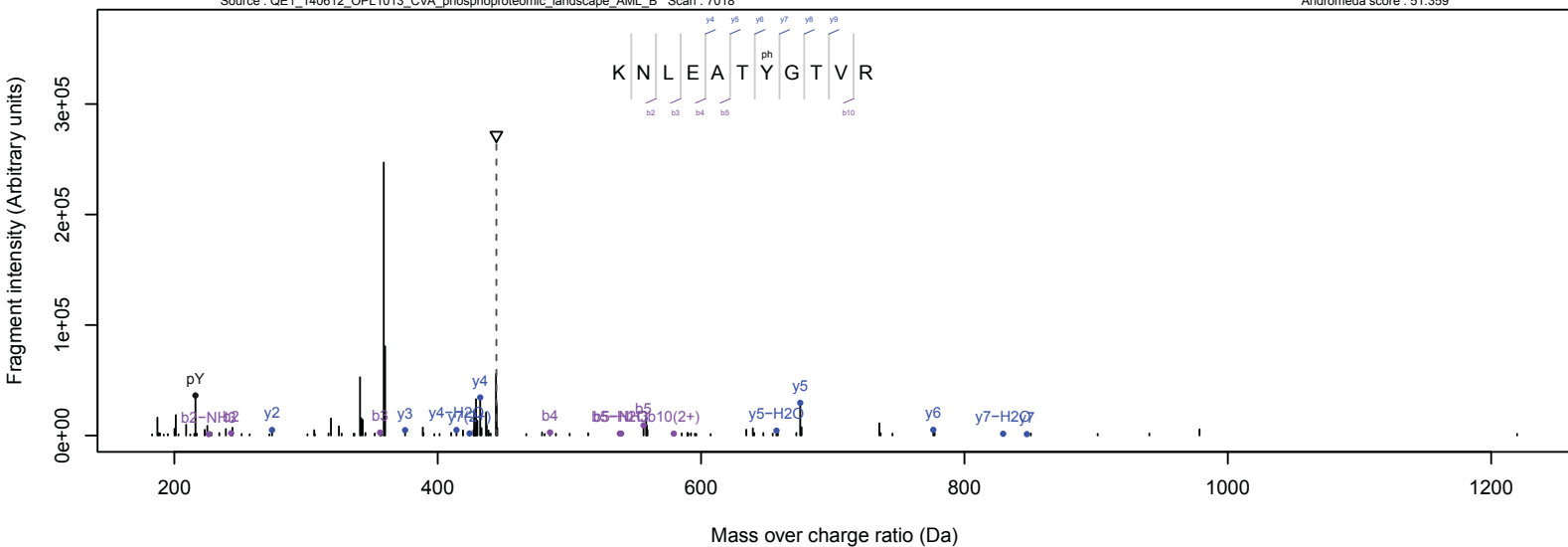
Andromeda score : 51.39



KNLEATY(ph)GTVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7018

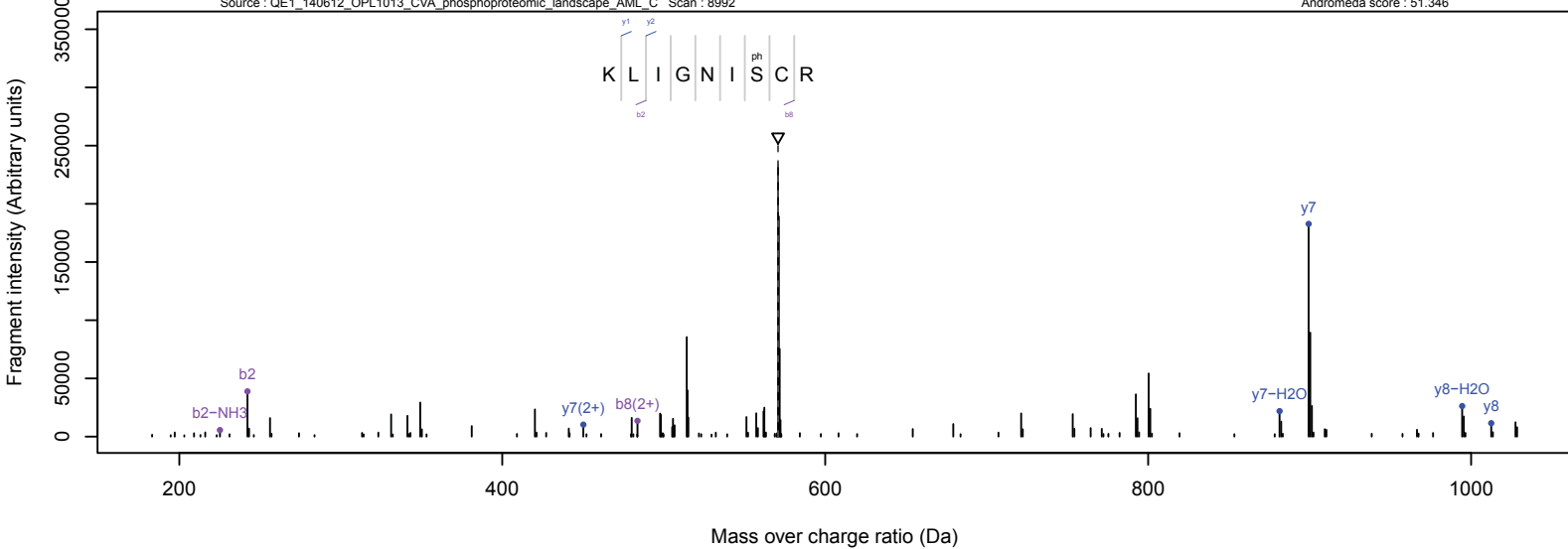
Andromeda score : 51.359



KLIGNIS(ph)CR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8992

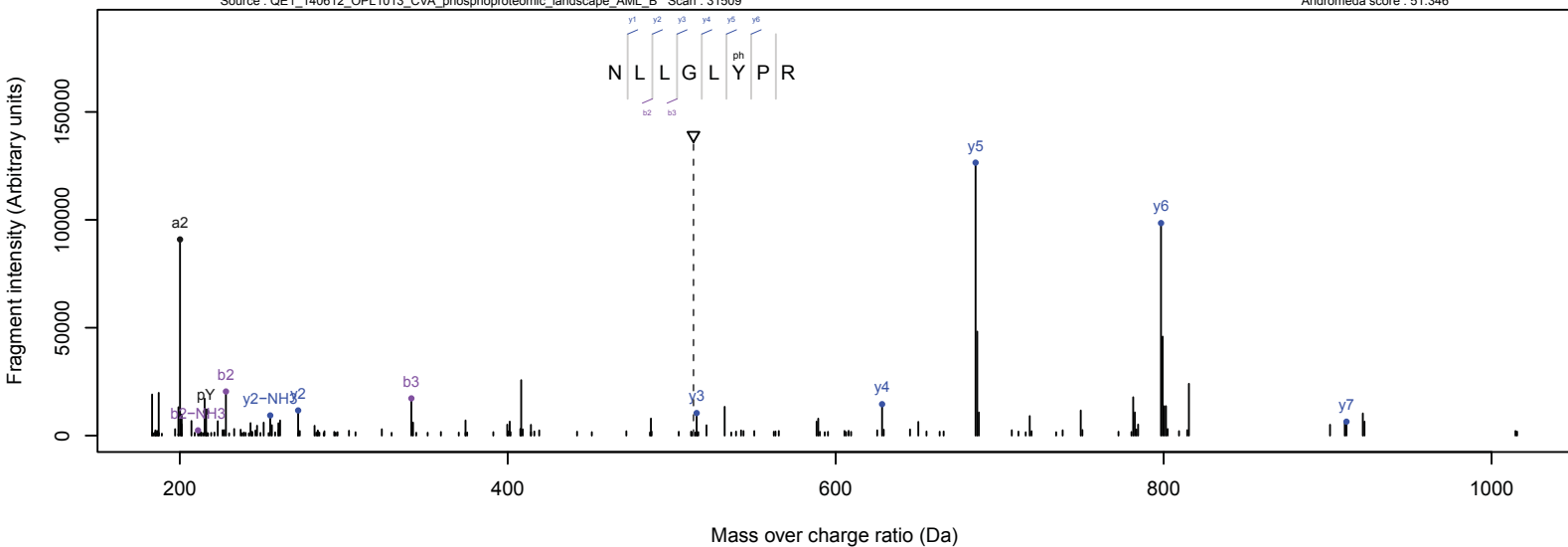
Andromeda score : 51.346



NLLGLY(ph)PR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 31509

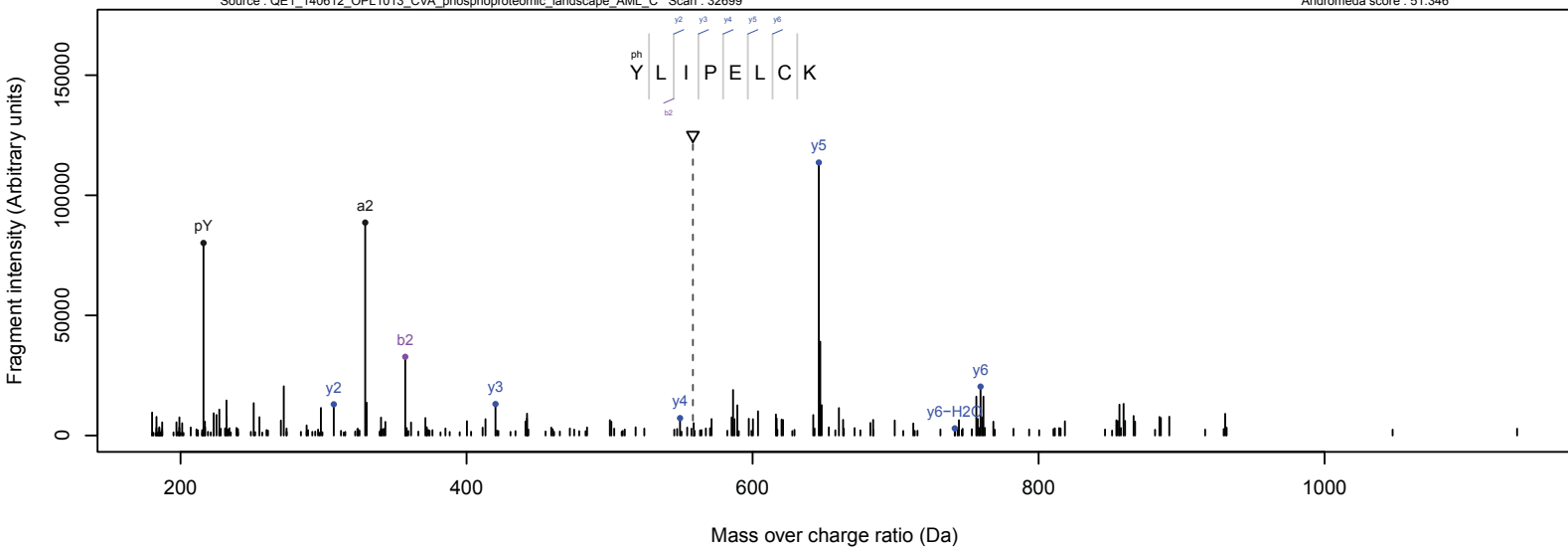
Andromeda score : 51.346



Y(ph)LIPELCK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32699

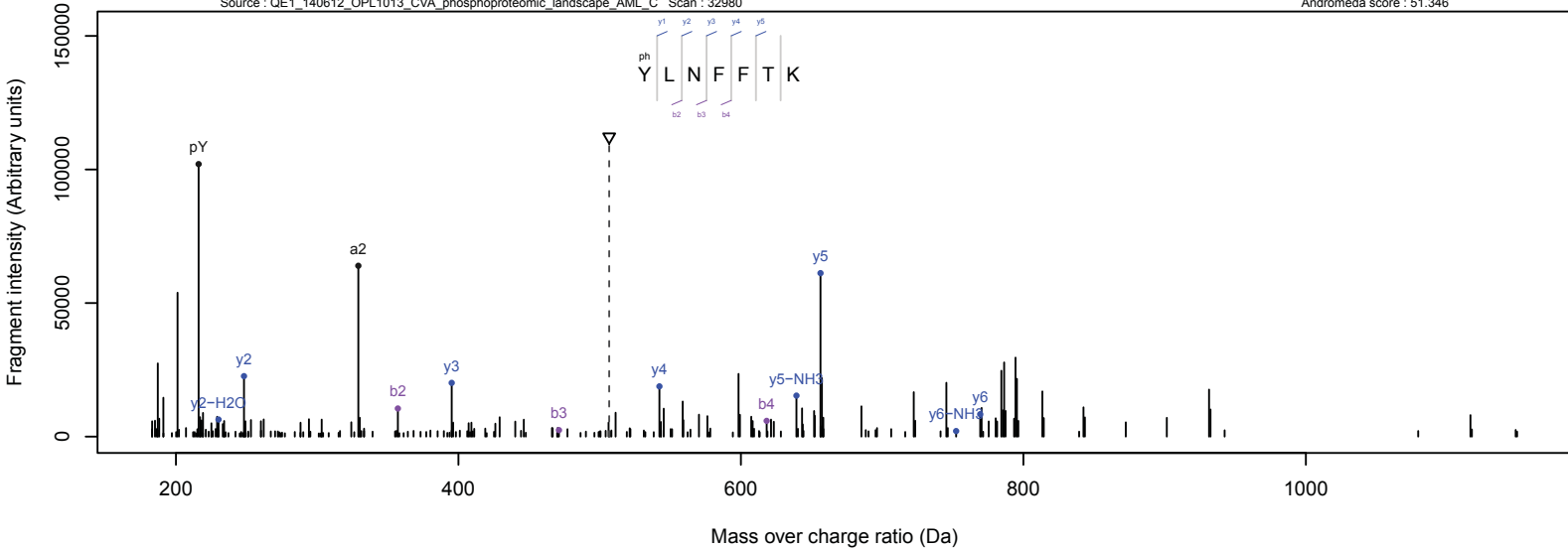
Andromeda score : 51.346



Y(ph)LNFFTK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32980

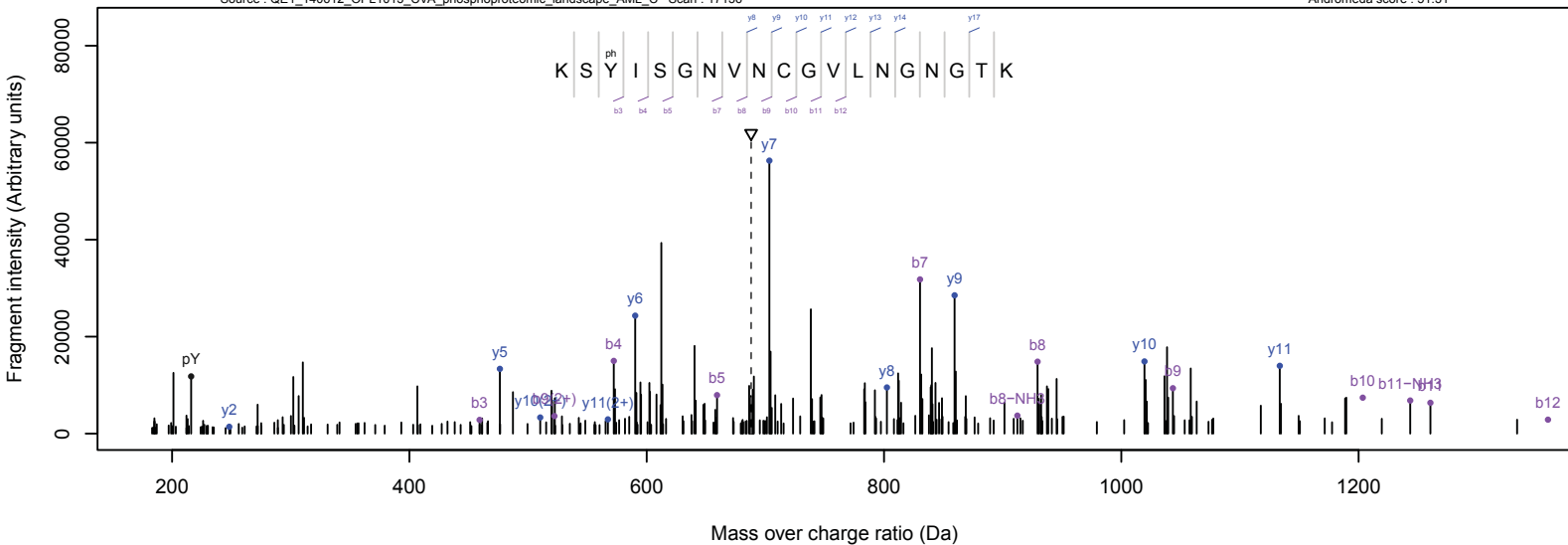
Andromeda score : 51.346



## KSY(ph)ISGNVNCGLVNLNGTK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17136

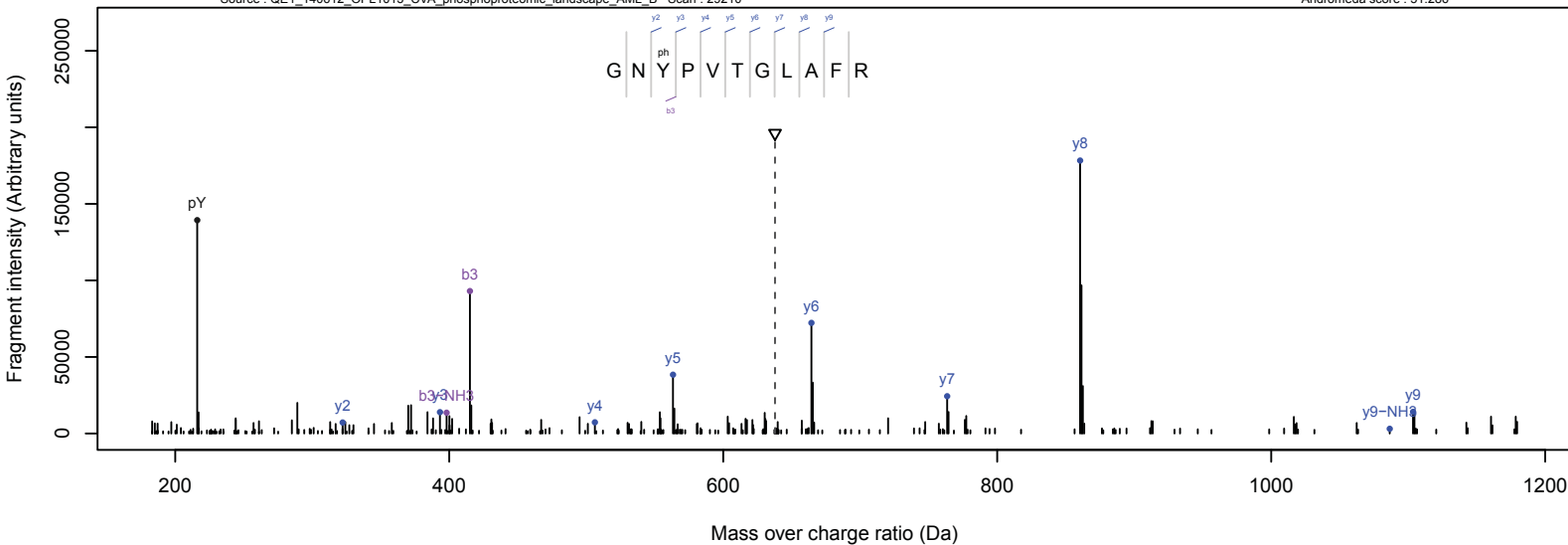
Andromeda score: 51.31



## GNYPVTGLAFR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 29210

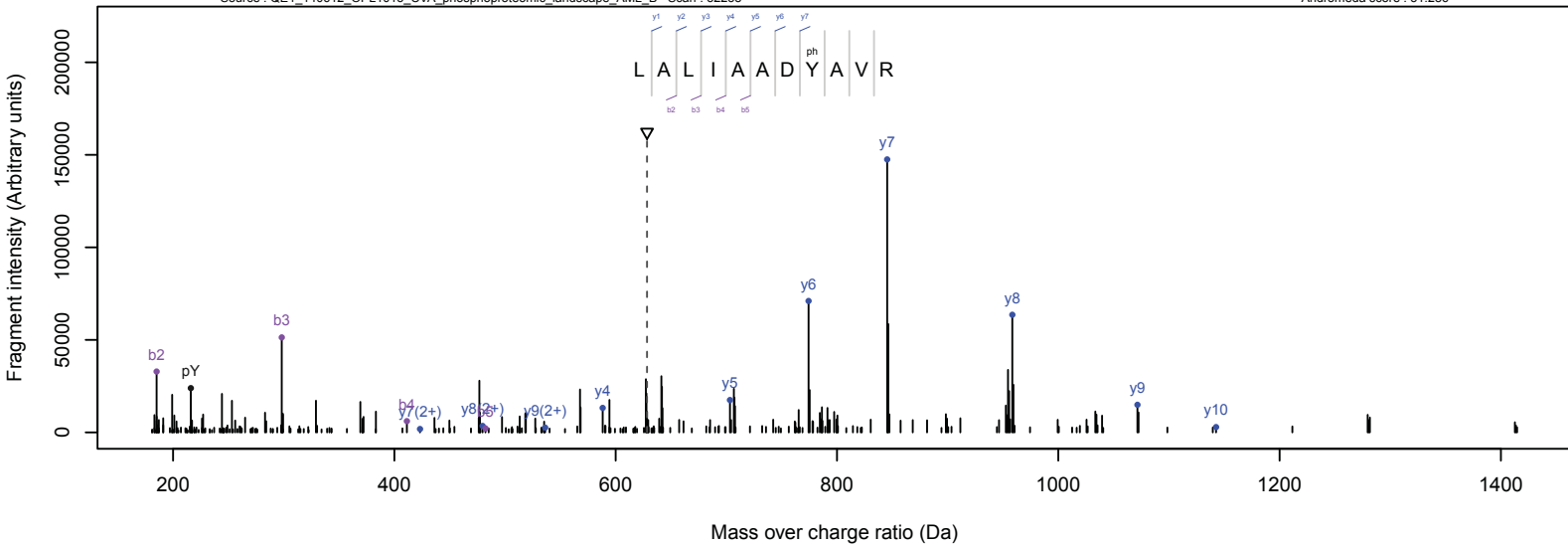
Andromeda score: 51.286



## LALIAADYAVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 32258

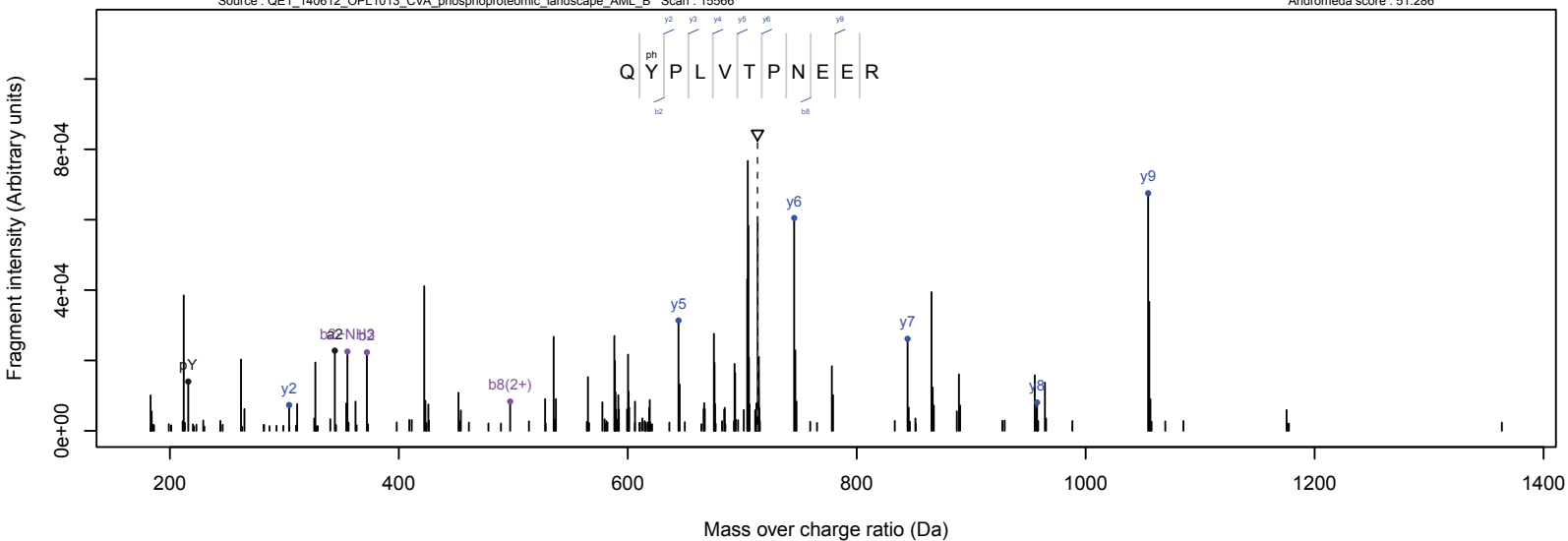
Andromeda score: 51.286



## QY(ph)PLVTPNEER-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15566

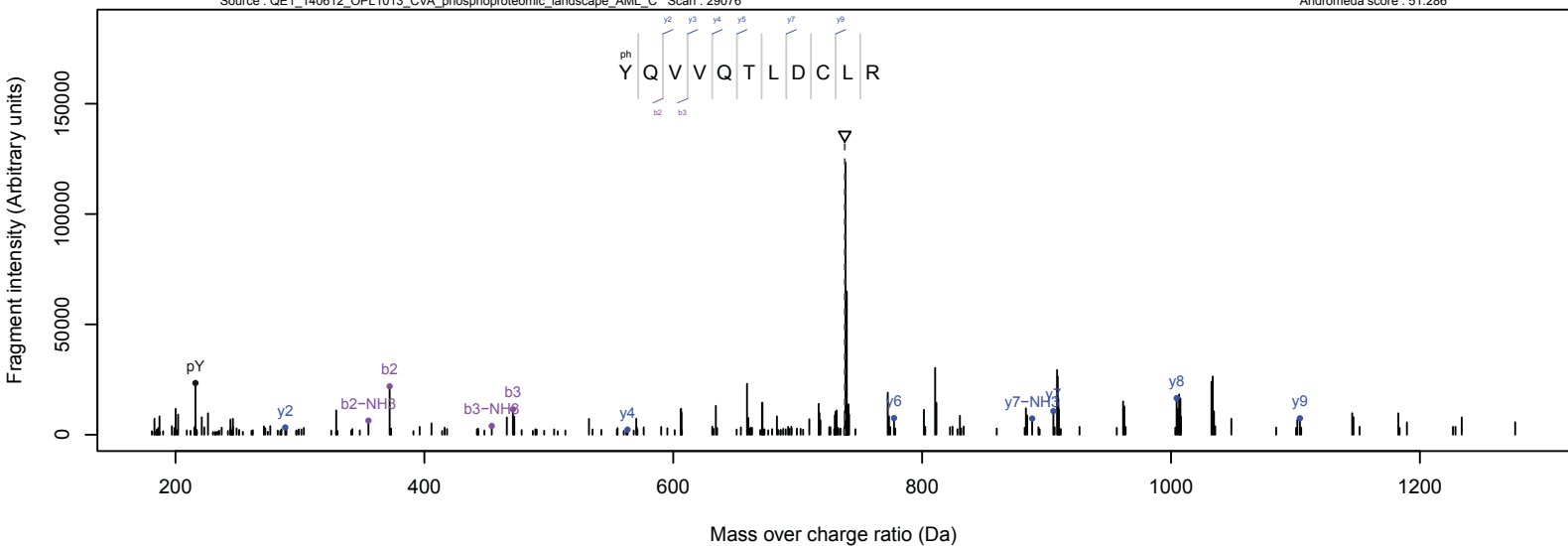
Andromeda score : 51.286



## Y(ph)QVVQTLDCLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29076

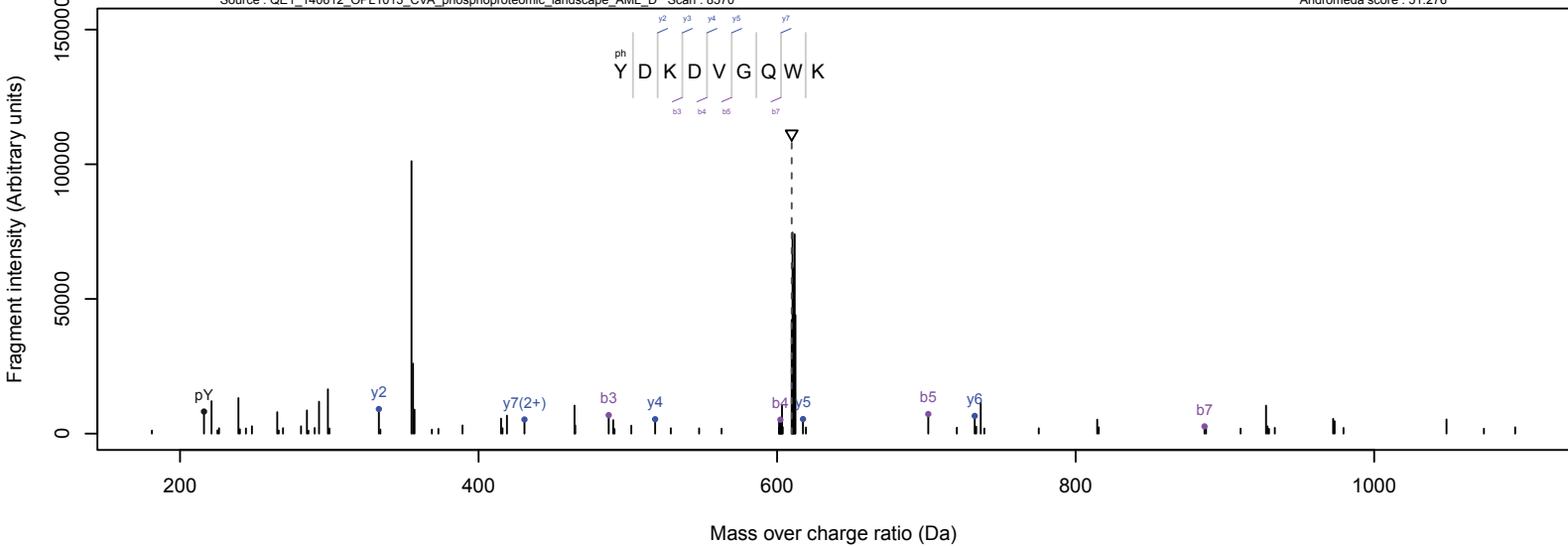
Andromeda score : 51.286



## Y(ph)DKDVGQWK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 8570

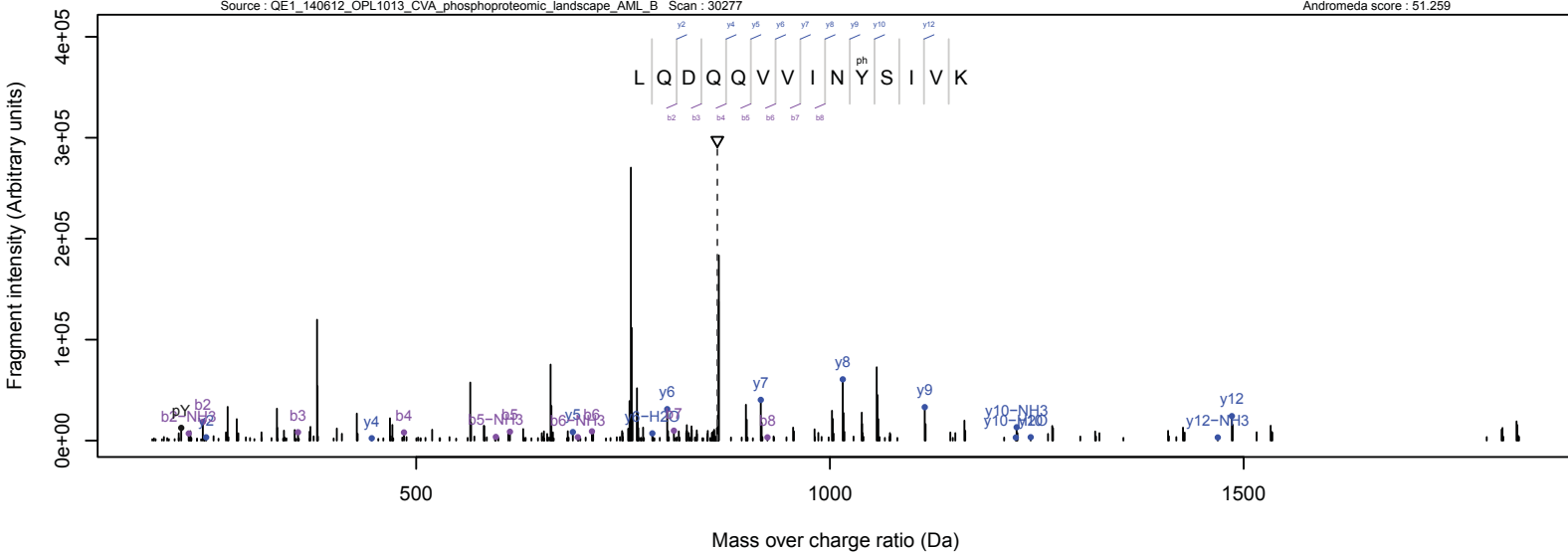
Andromeda score : 51.276



## LQDQQVINY(ph)SIVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30277

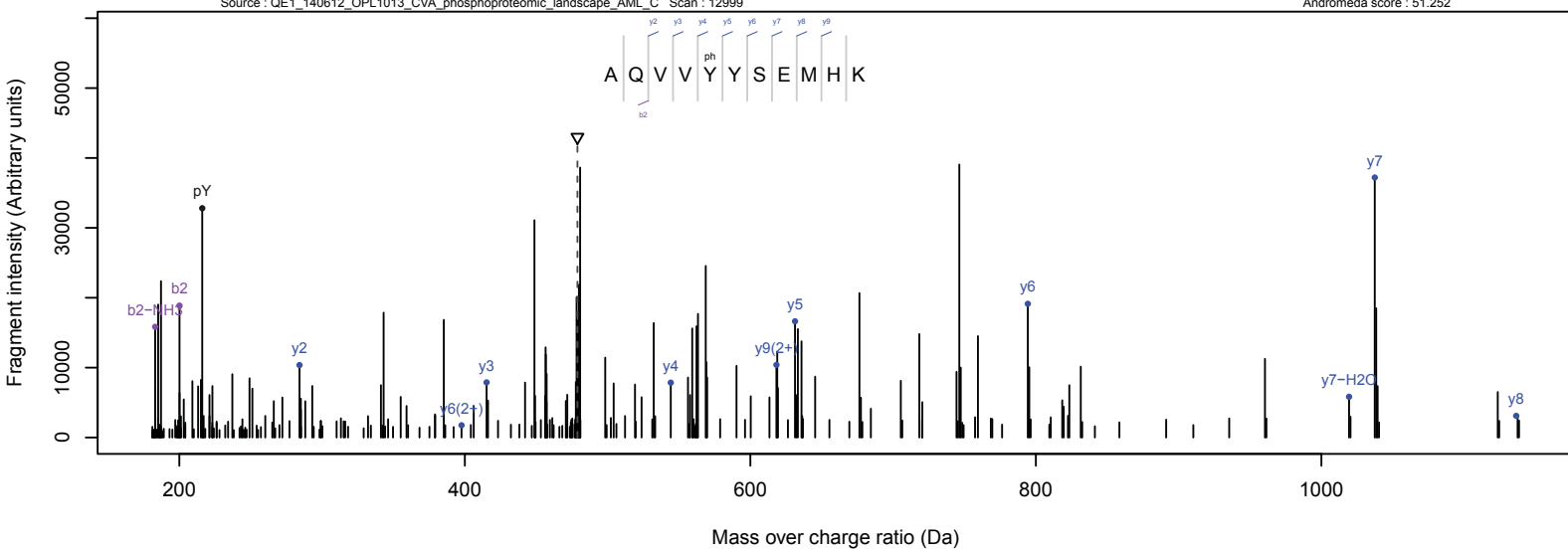
Andromeda score : 51.259



## AQVVY(ph)YSEMHK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12999

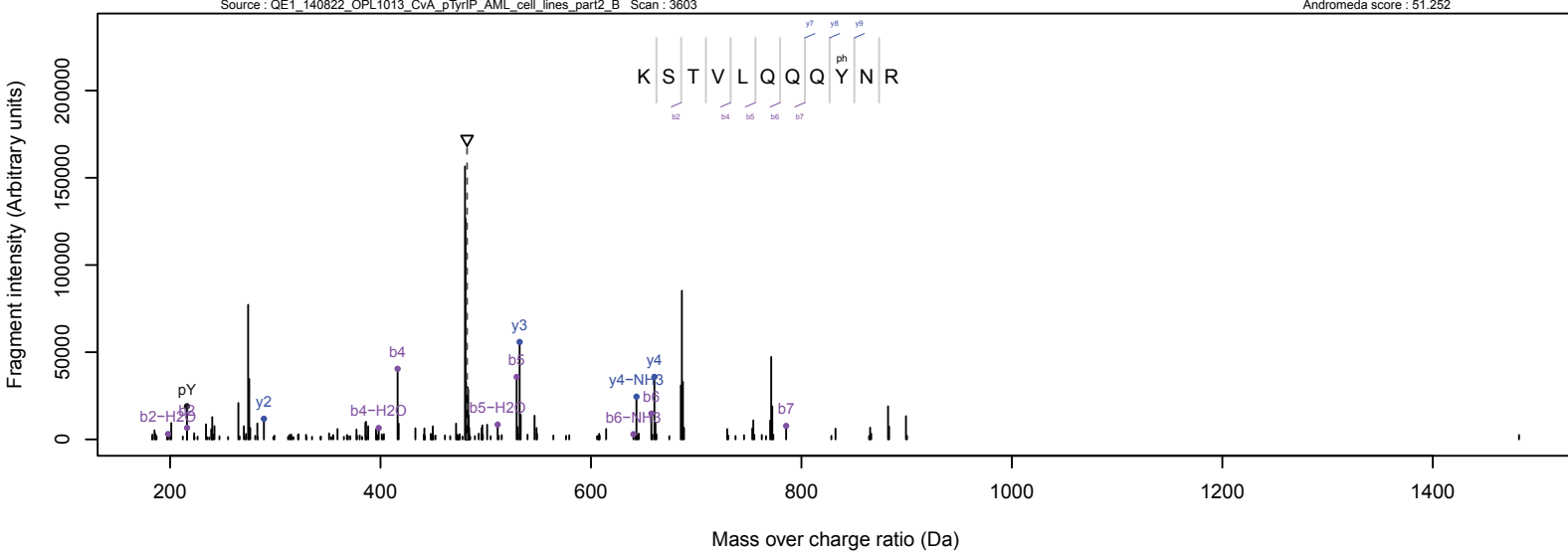
Andromeda score : 51.252



## KSTVLQQQY(ph)NR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3603

Andromeda score : 51.252

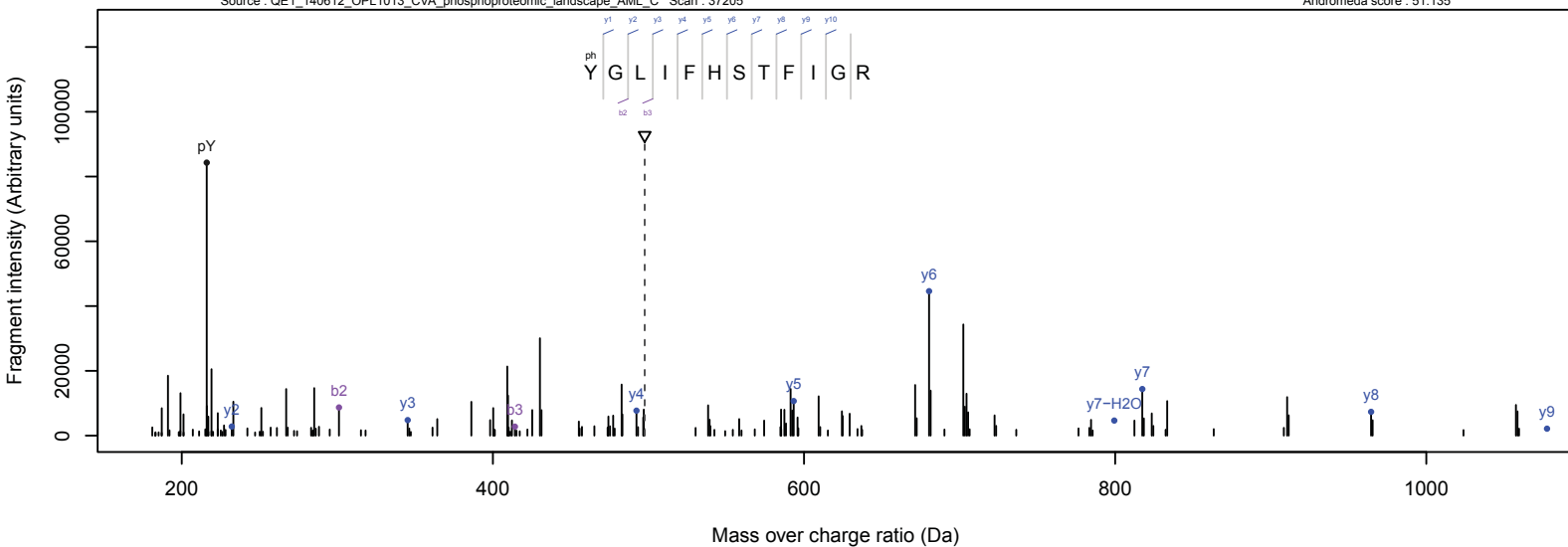




## Y(ph)GLIFHSTFIGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 37205

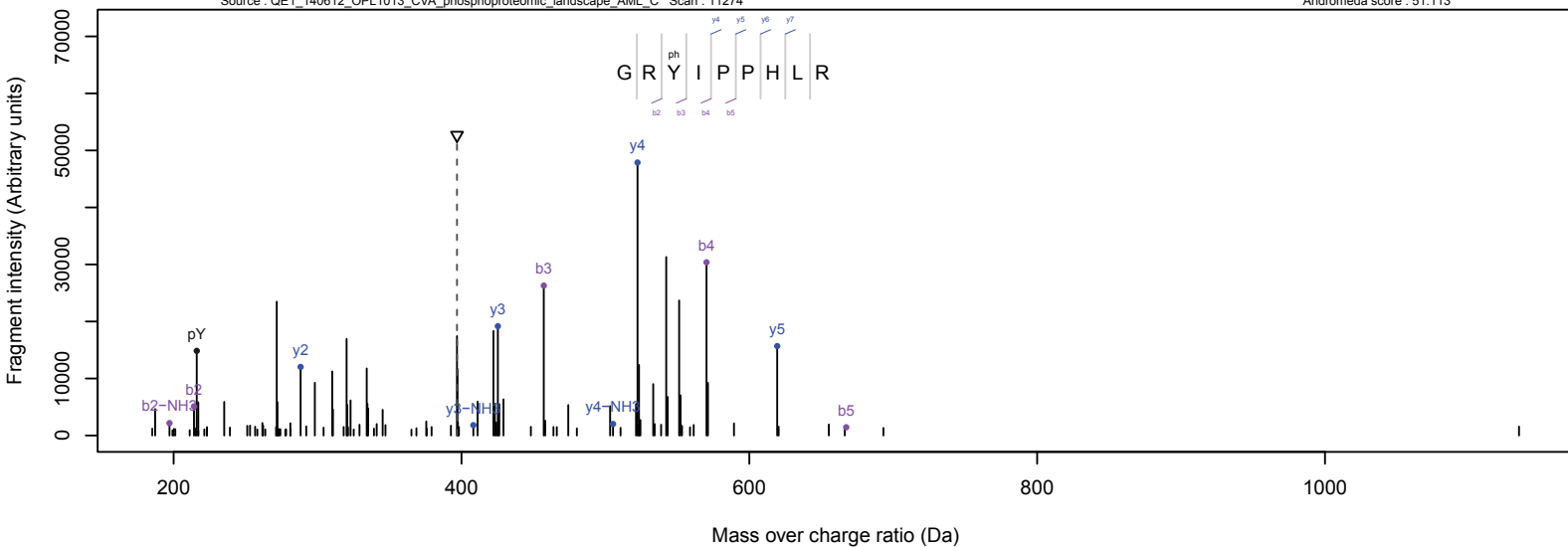
Andromeda score: 51.135



## GRY(ph)IPPHLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 11274

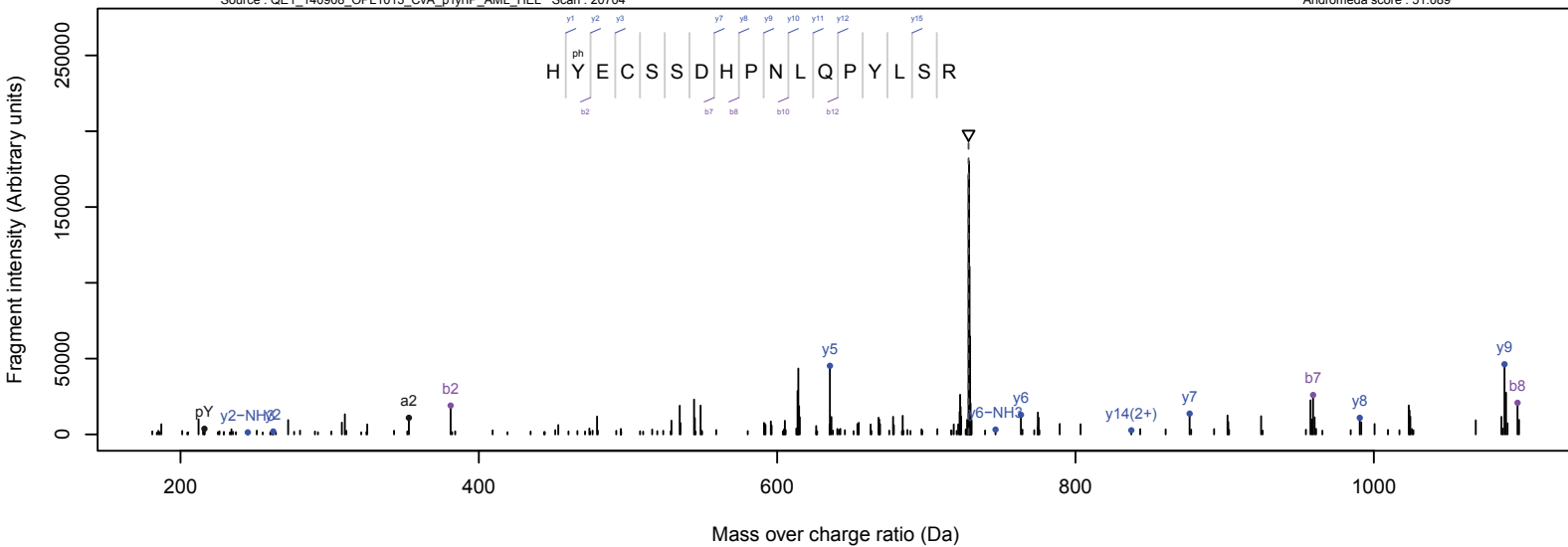
Andromeda score: 51.113

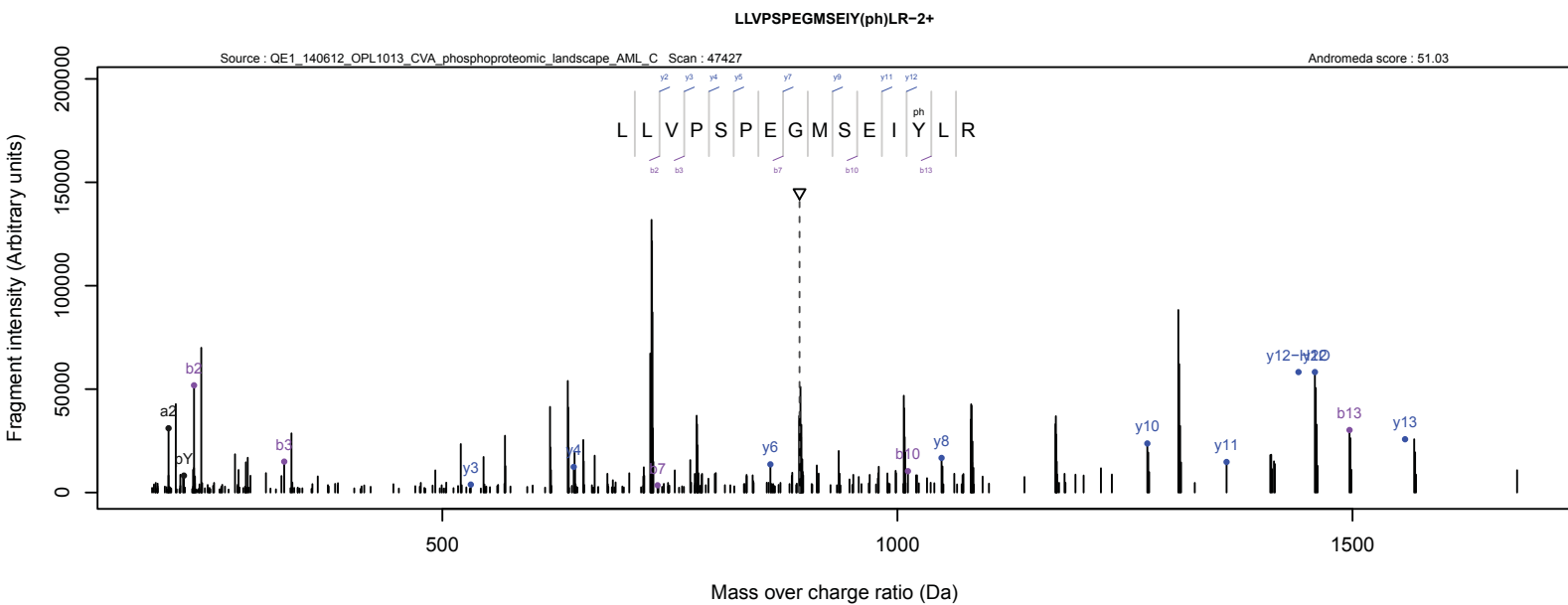
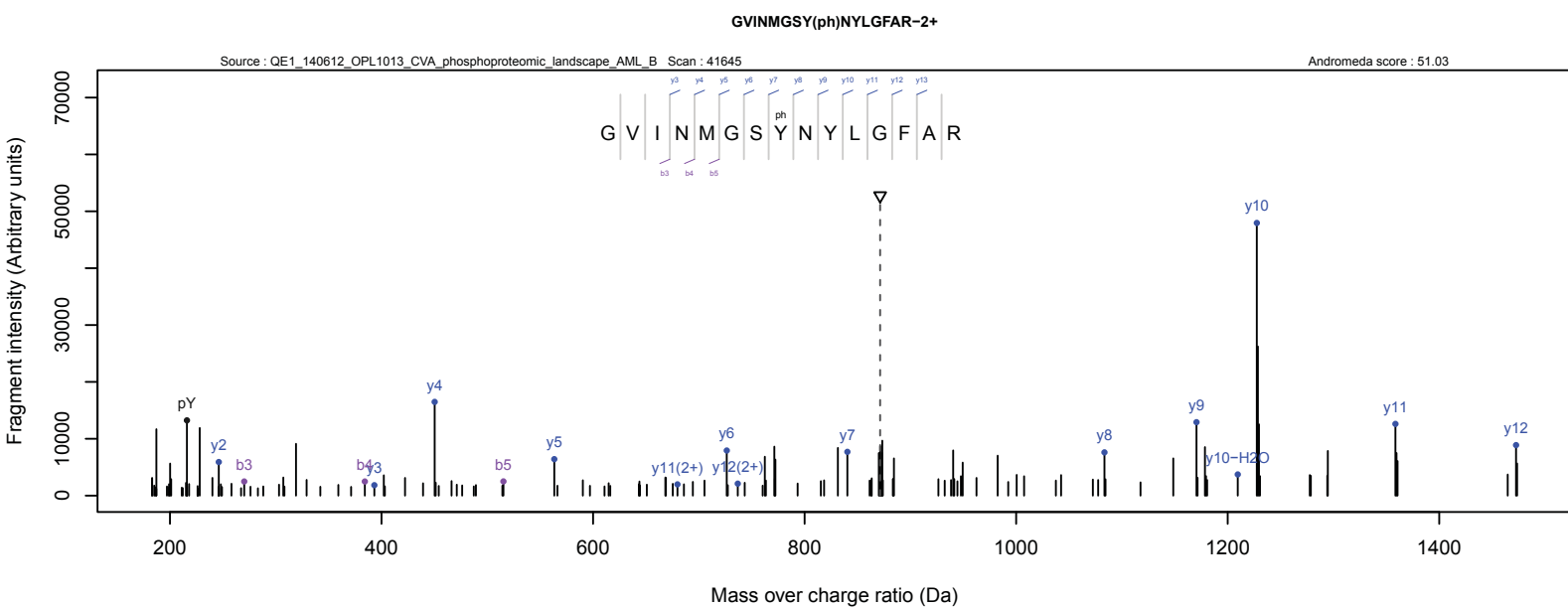
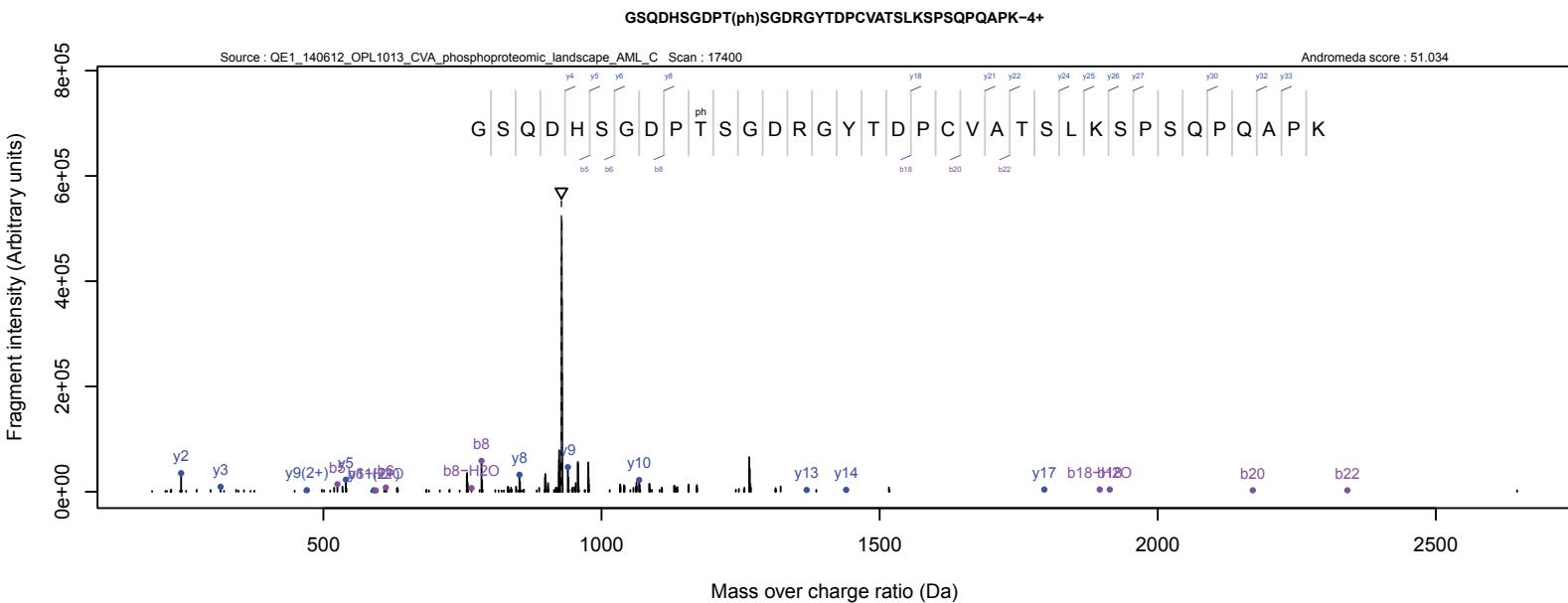


## HY(ph)ECSSDHPNLQPYLSR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 20704

Andromeda score: 51.089

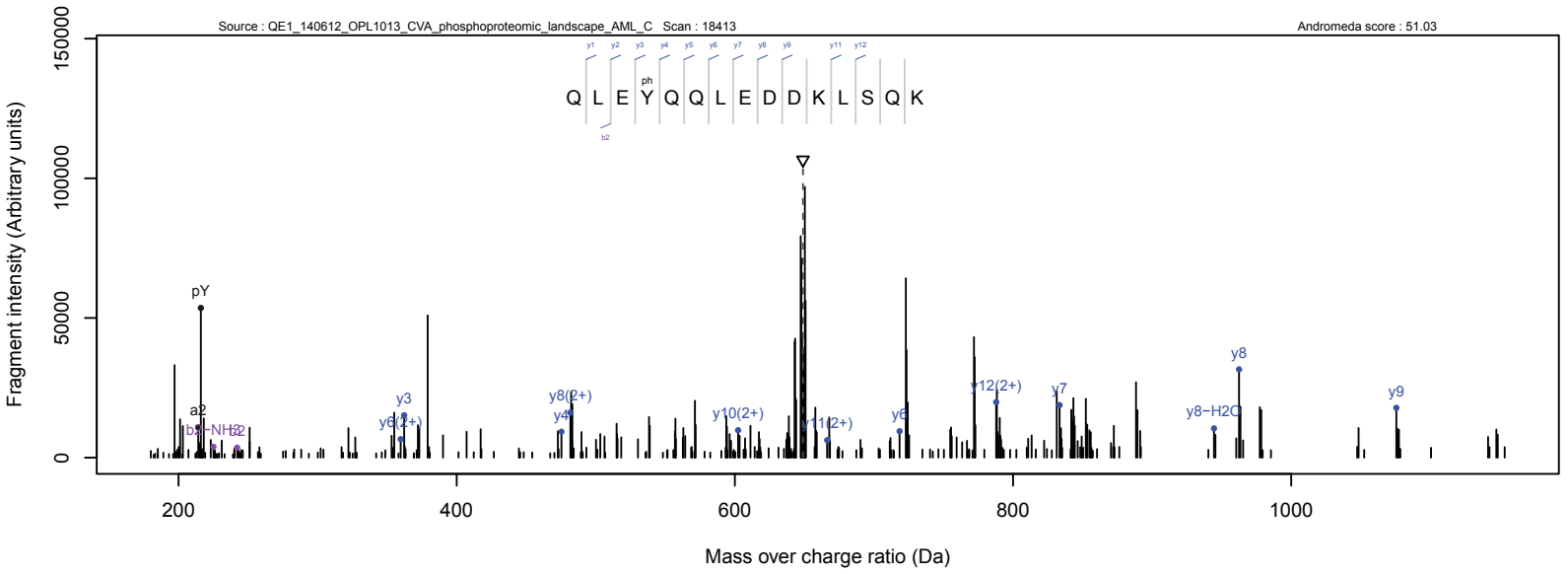




**QLEY(ph)QLEDDKLSQK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18413

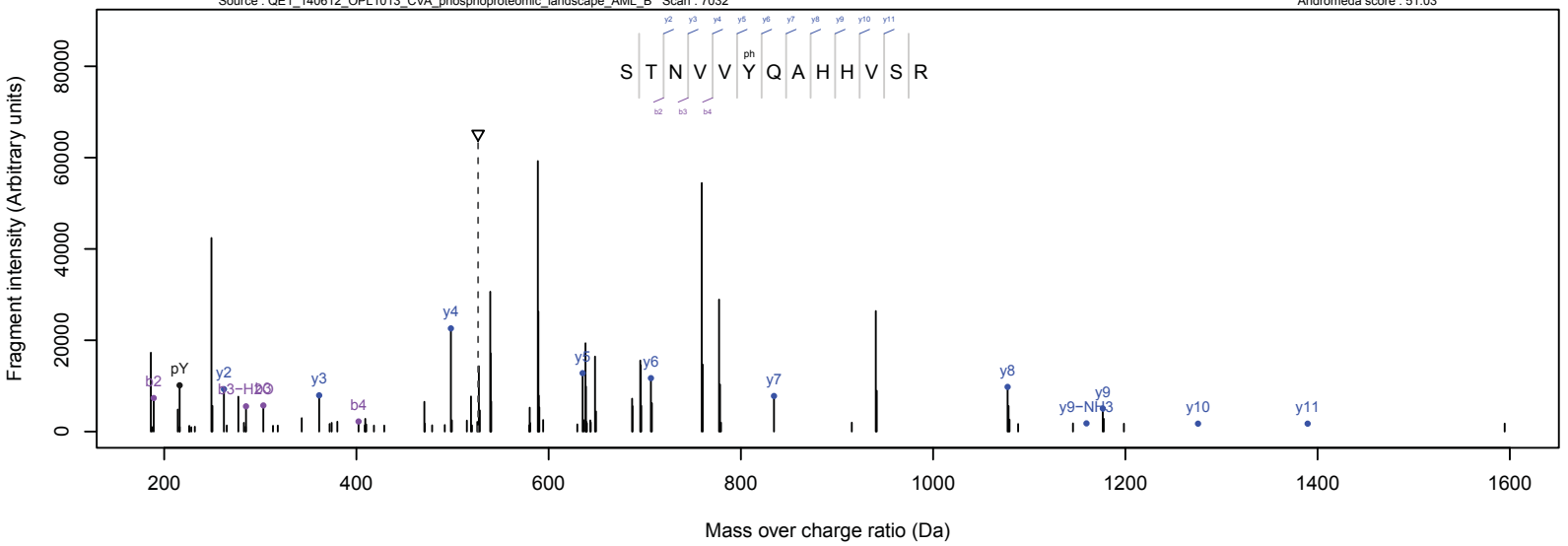
Andromeda score : 51.03



**STNVVY(ph)QAHHVSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7032

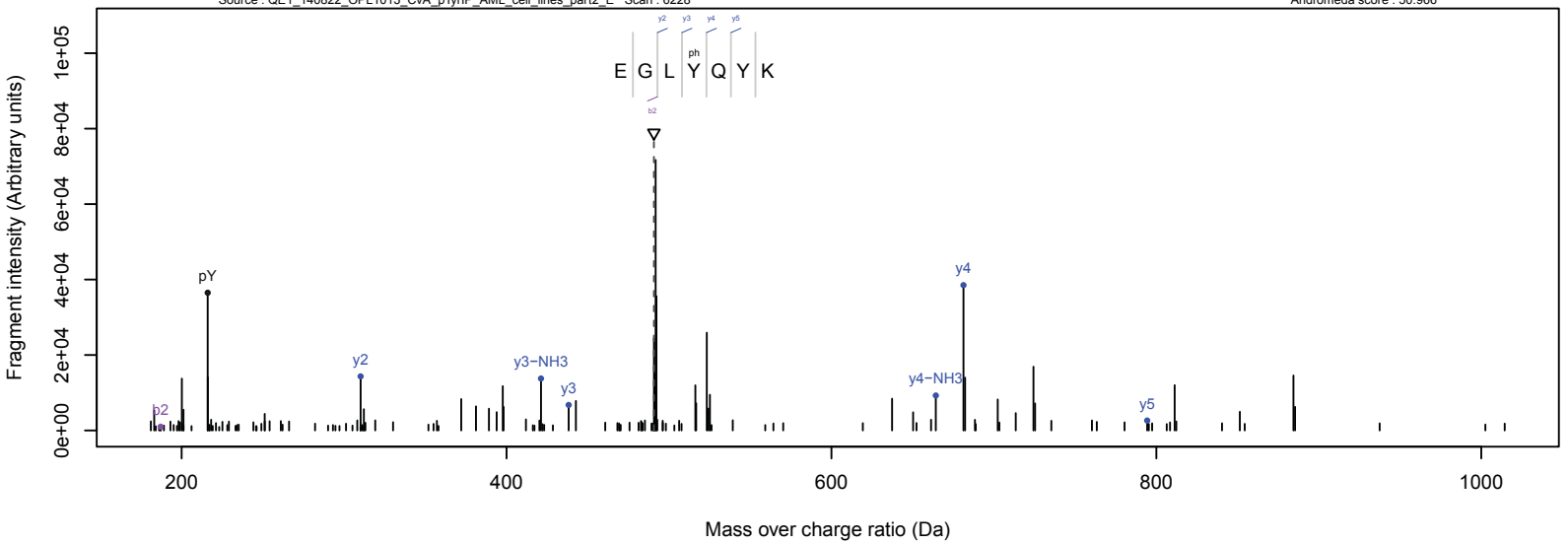
Andromeda score : 51.03



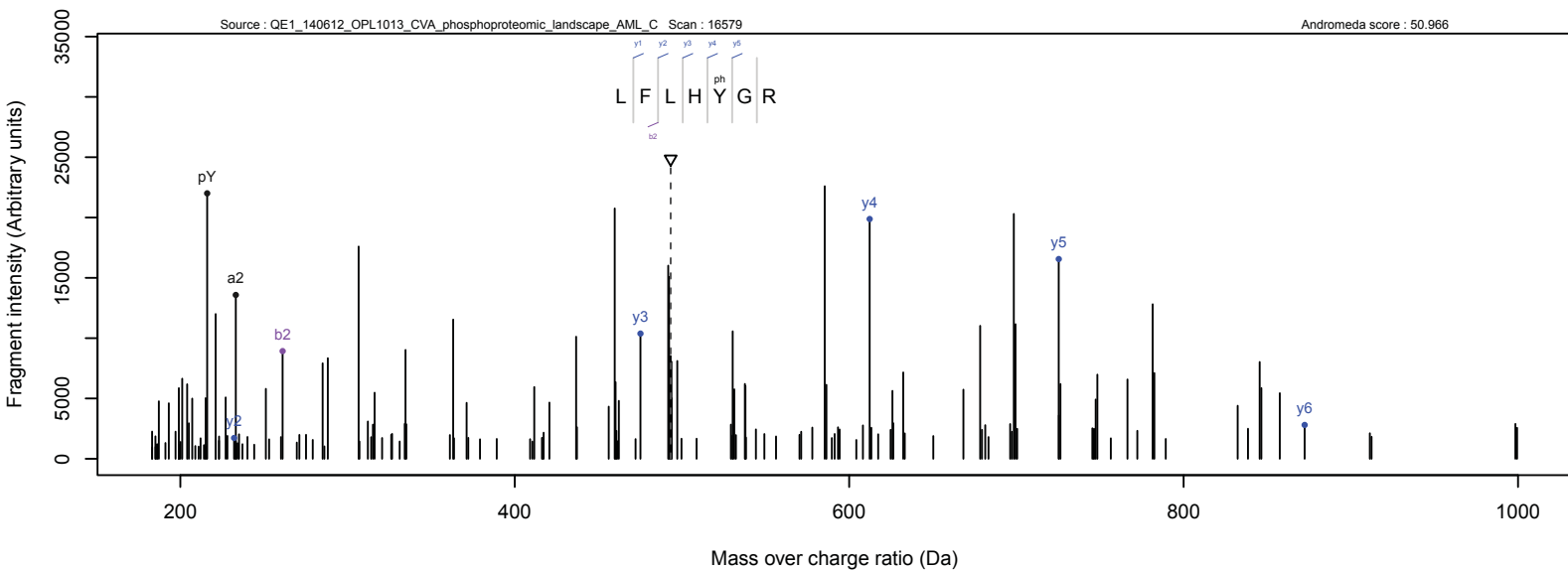
**EGLY(ph)QYK-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 6228

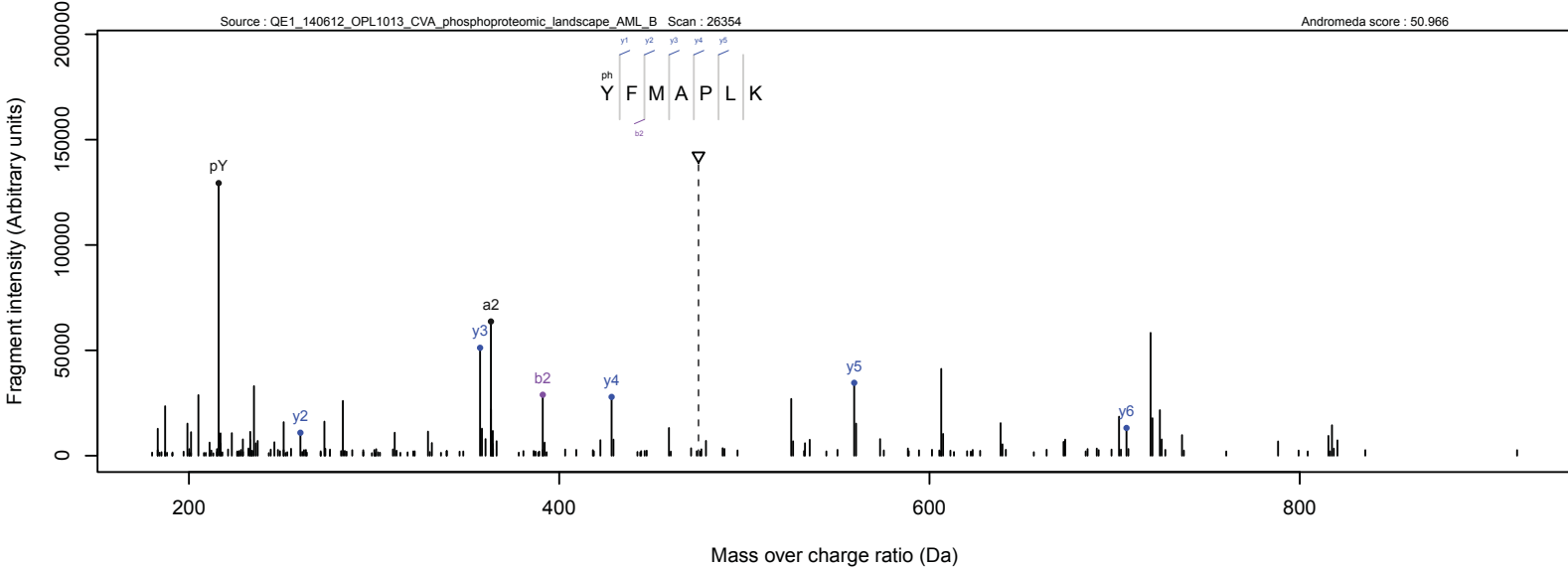
Andromeda score : 50.966



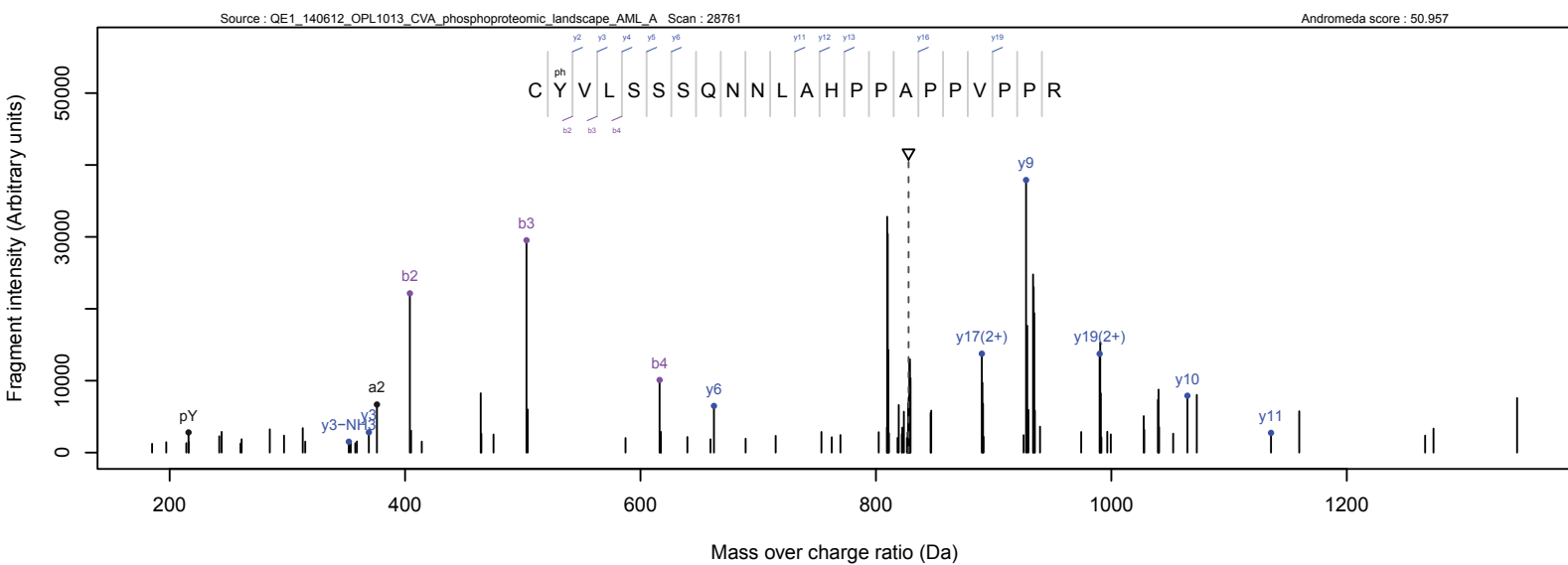
LFLHY(ph)GR-2+



Y(ph)FMAPLK-2+



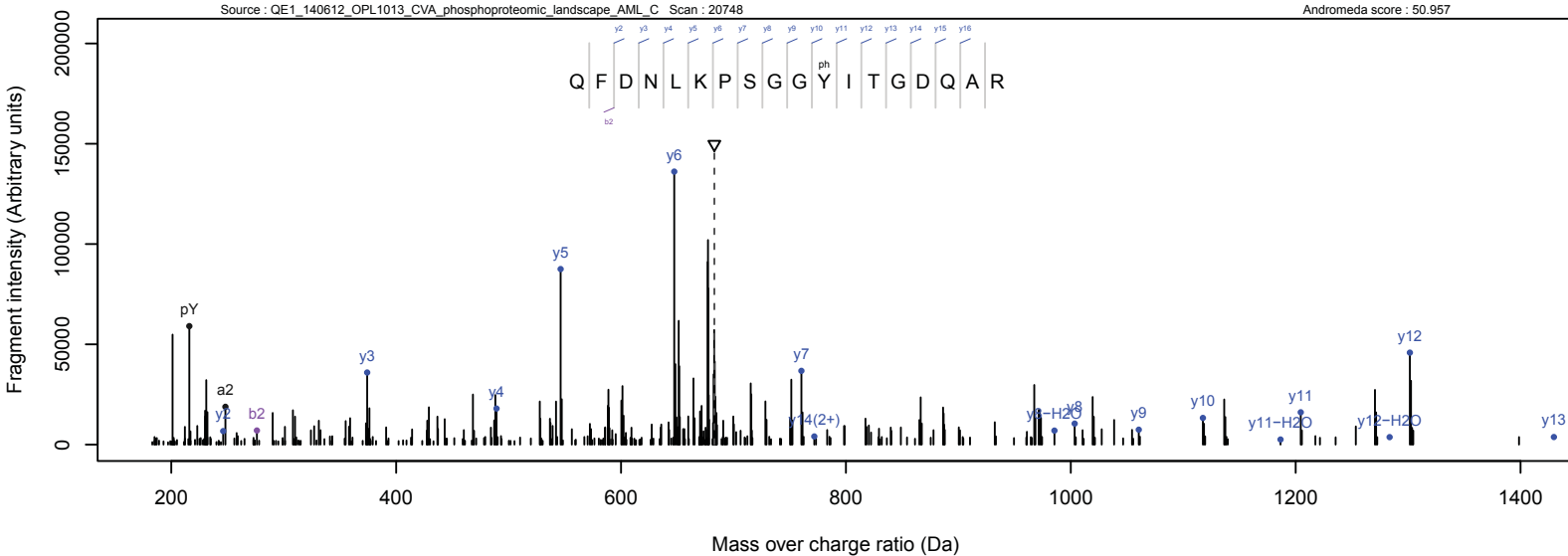
CY(ph)VLSSQNLAHPAPPVPPR-3+



## QFDNLKPSGGY(ph)ITGDQAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 20748

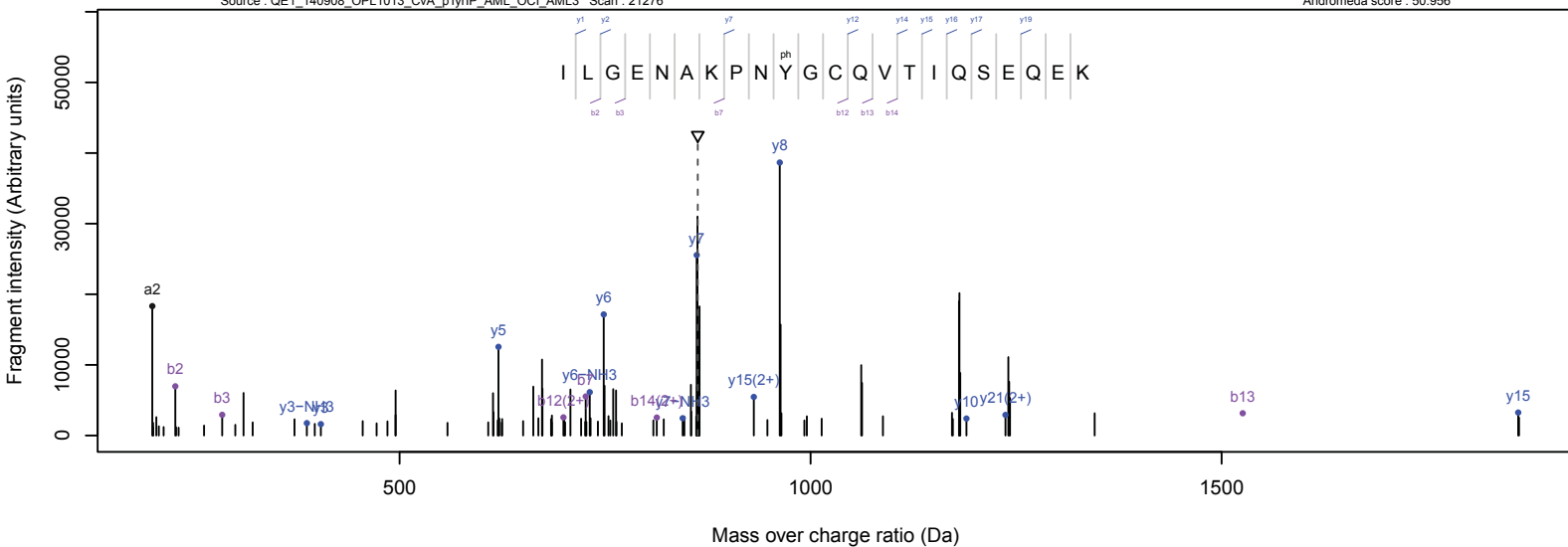
Andromeda score: 50.957



## ILGENAKPNY(ph)GCQVTIQSEQEK-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan: 21276

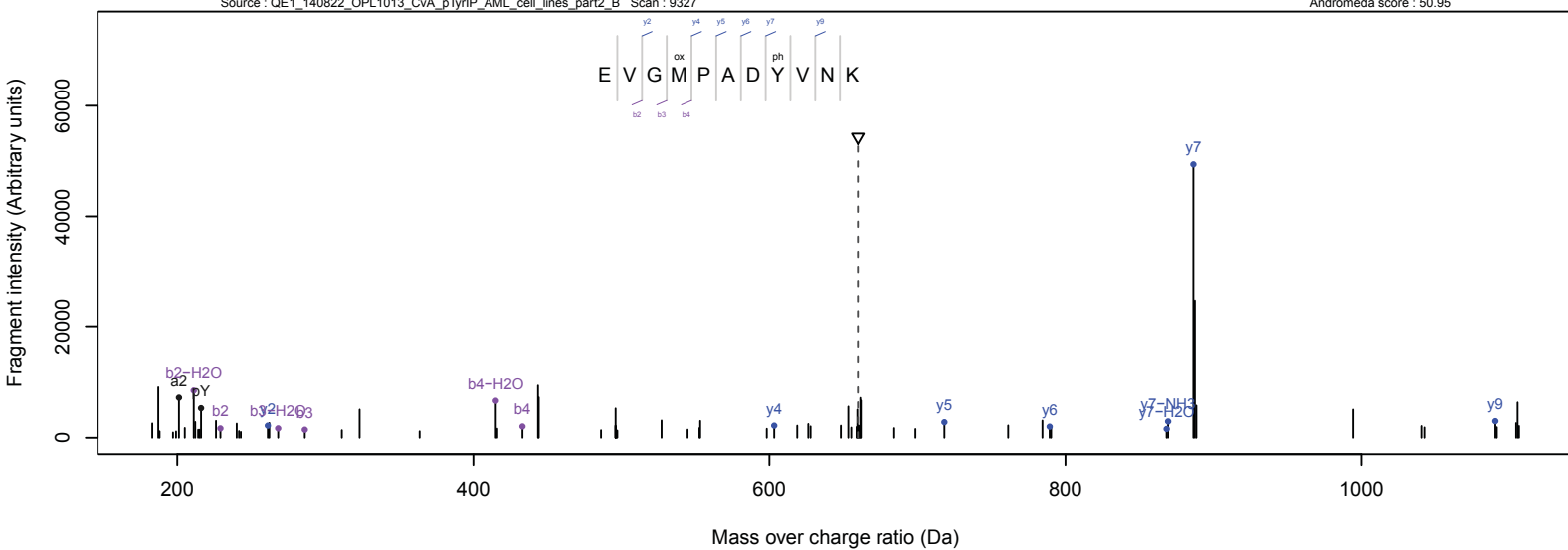
Andromeda score: 50.956



## EVGM(ox)PADY(ph)VNK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 9327

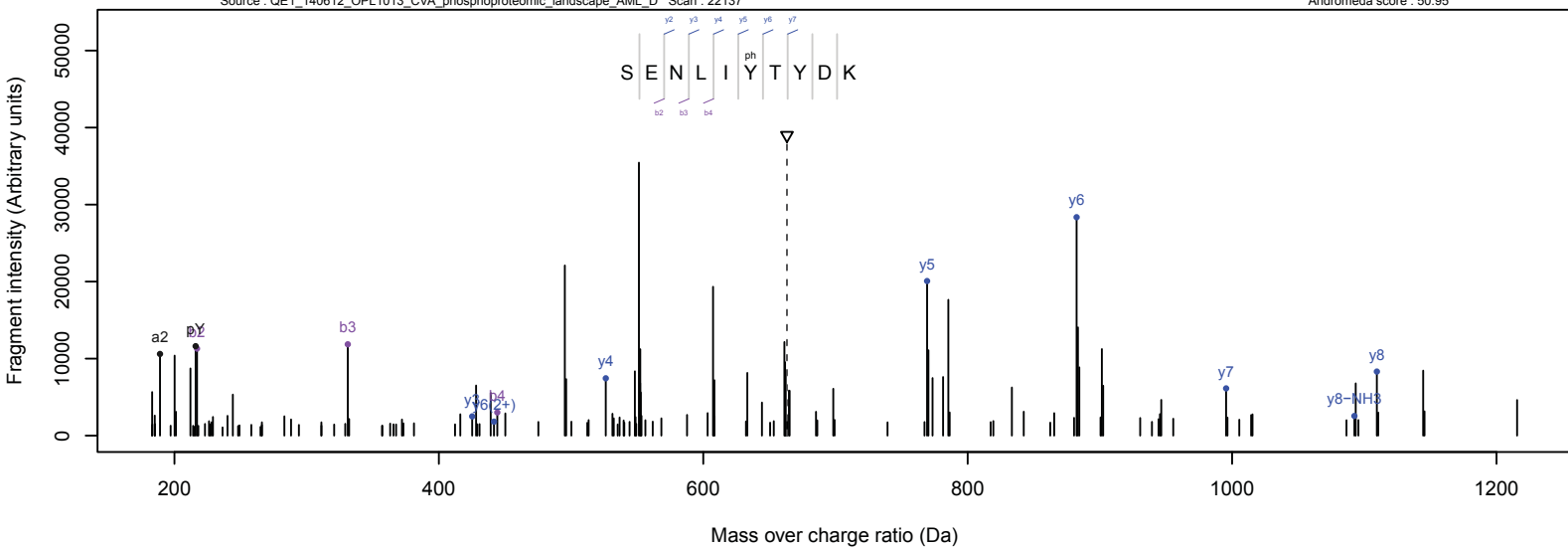
Andromeda score: 50.95



## SENLIY(ph)TYDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 22137

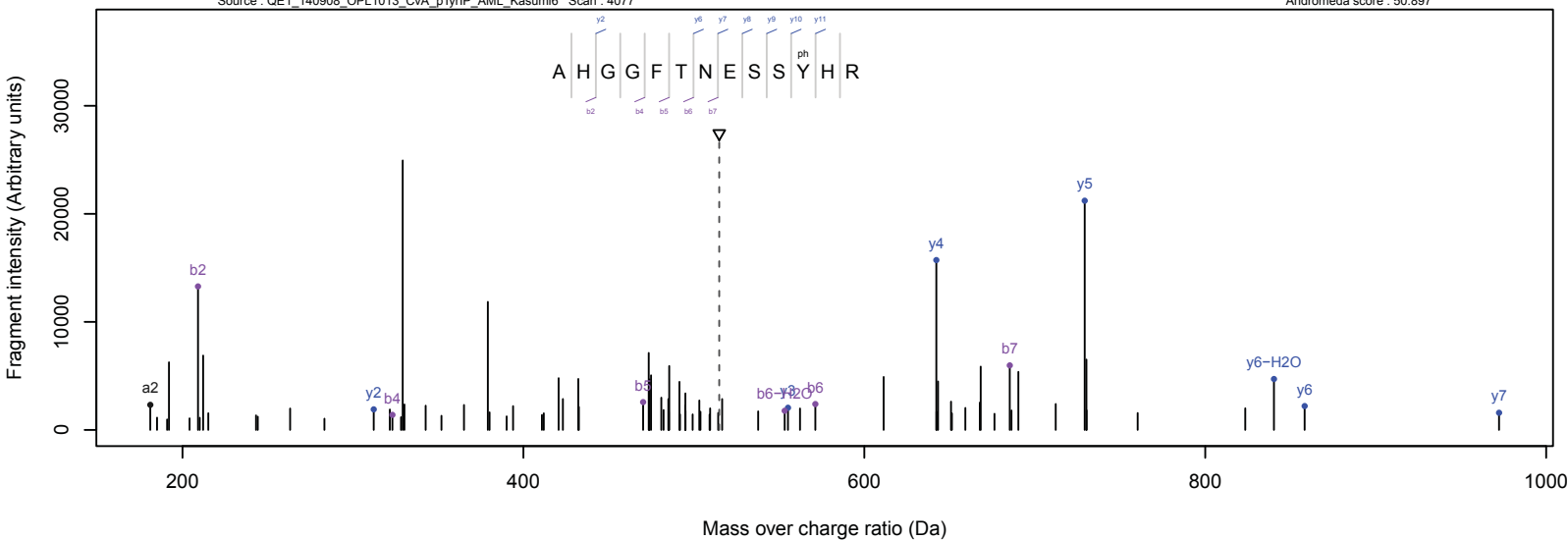
Andromeda score : 50.95



## AHGGFTNESSY(ph)HR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 4077

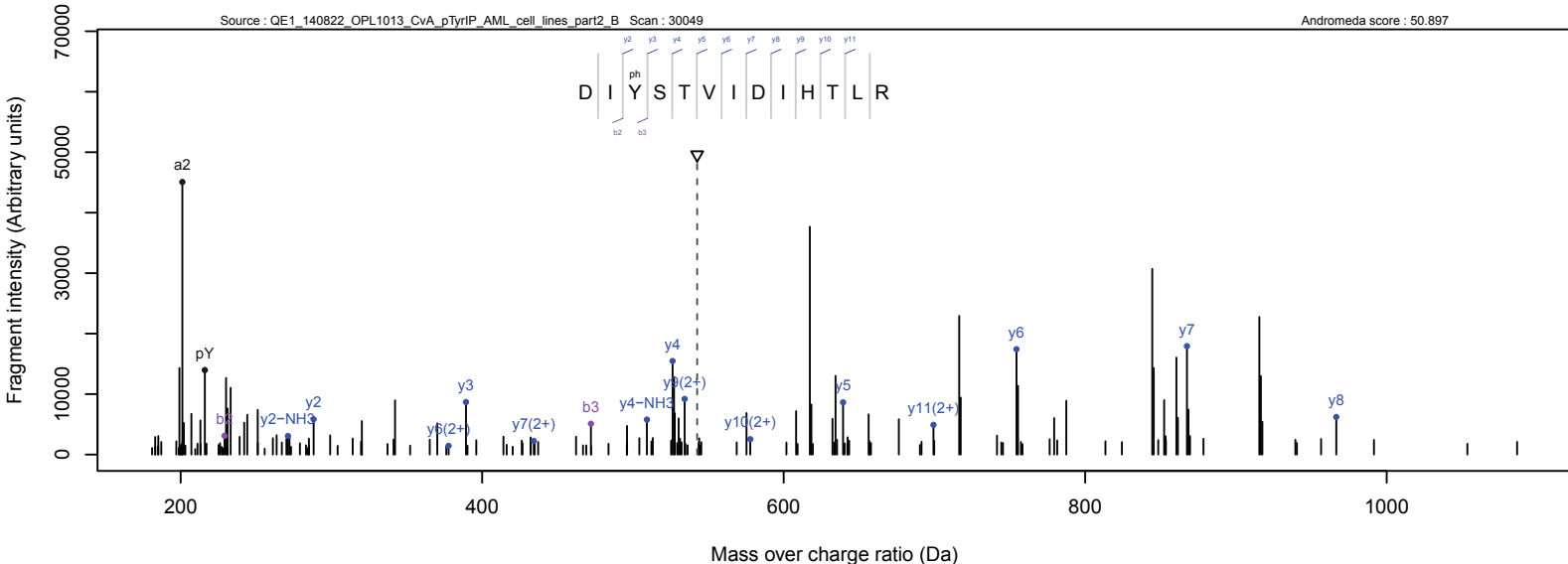
Andromeda score : 50.897



## DIY(ph)STVIDIHLR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 30049

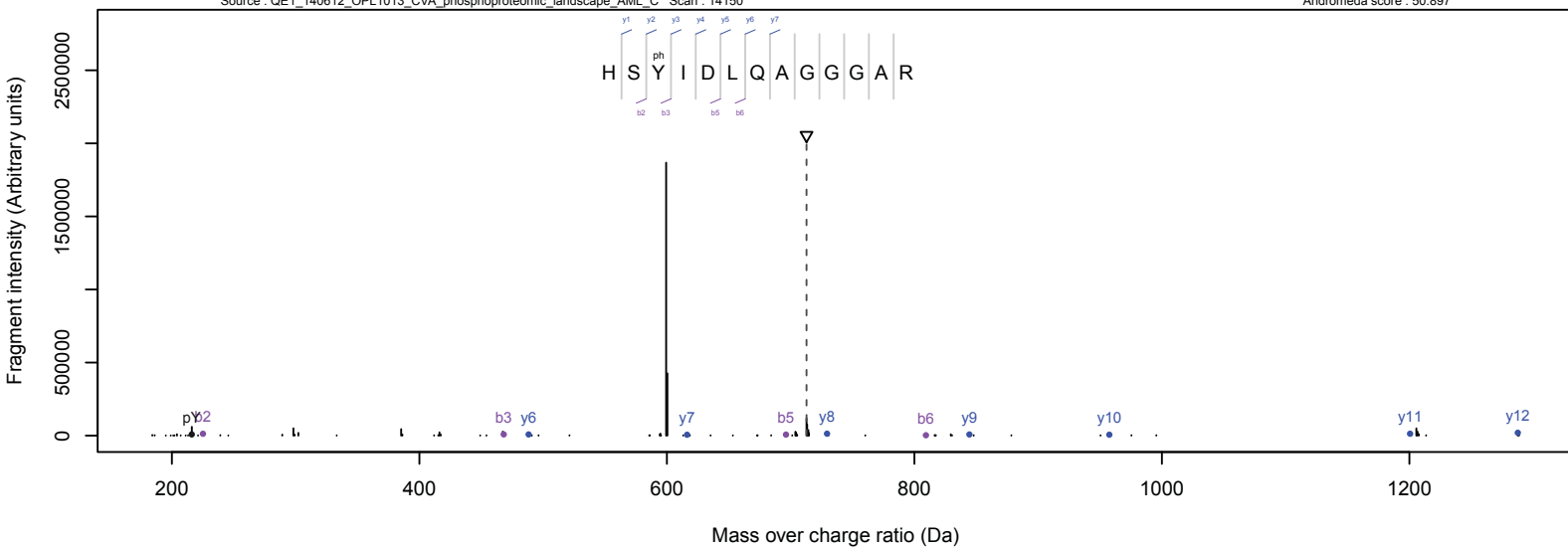
Andromeda score : 50.897



## HSY(ph)IDLQAGGAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14150

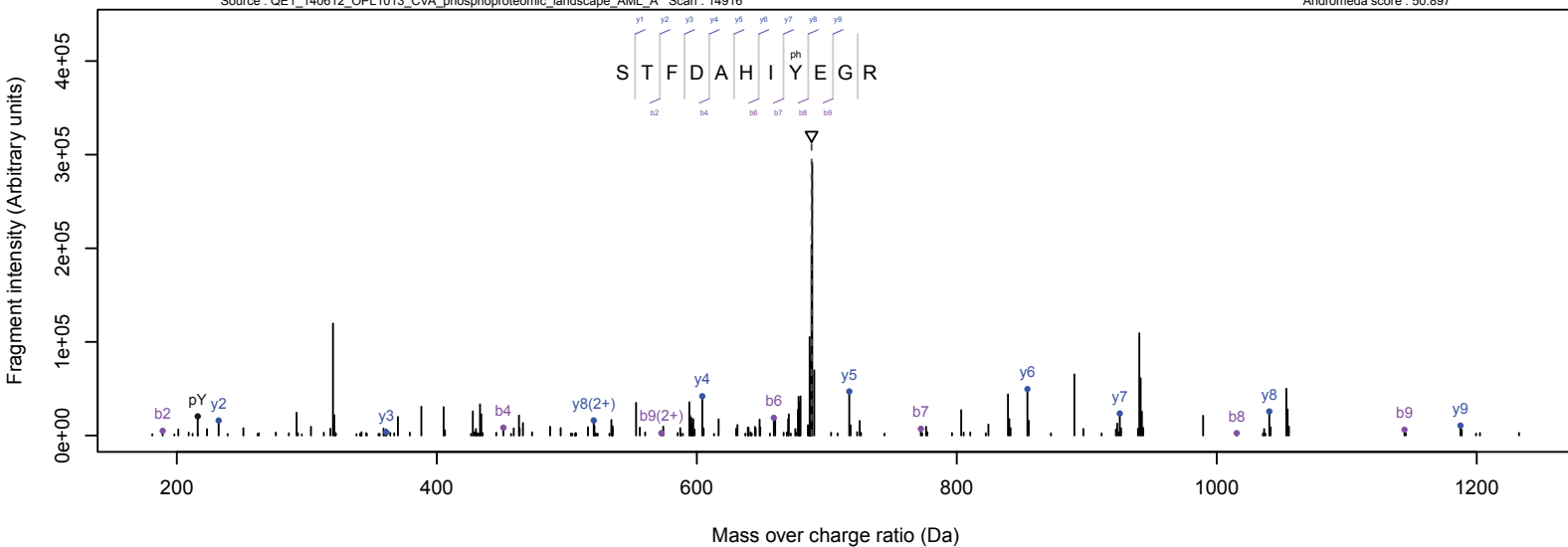
Andromeda score : 50.897



## STFDAHIY(ph)EGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 14916

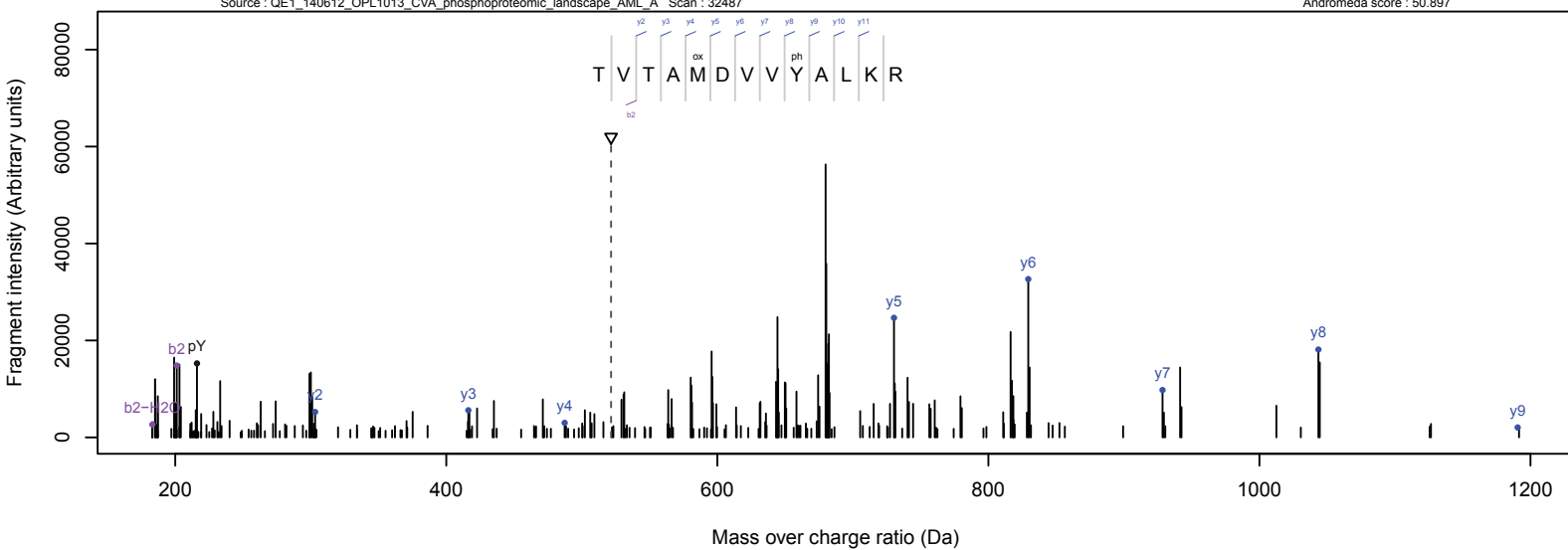
Andromeda score : 50.897



## TVTAM(ox)DVVY(ph)ALKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 32487

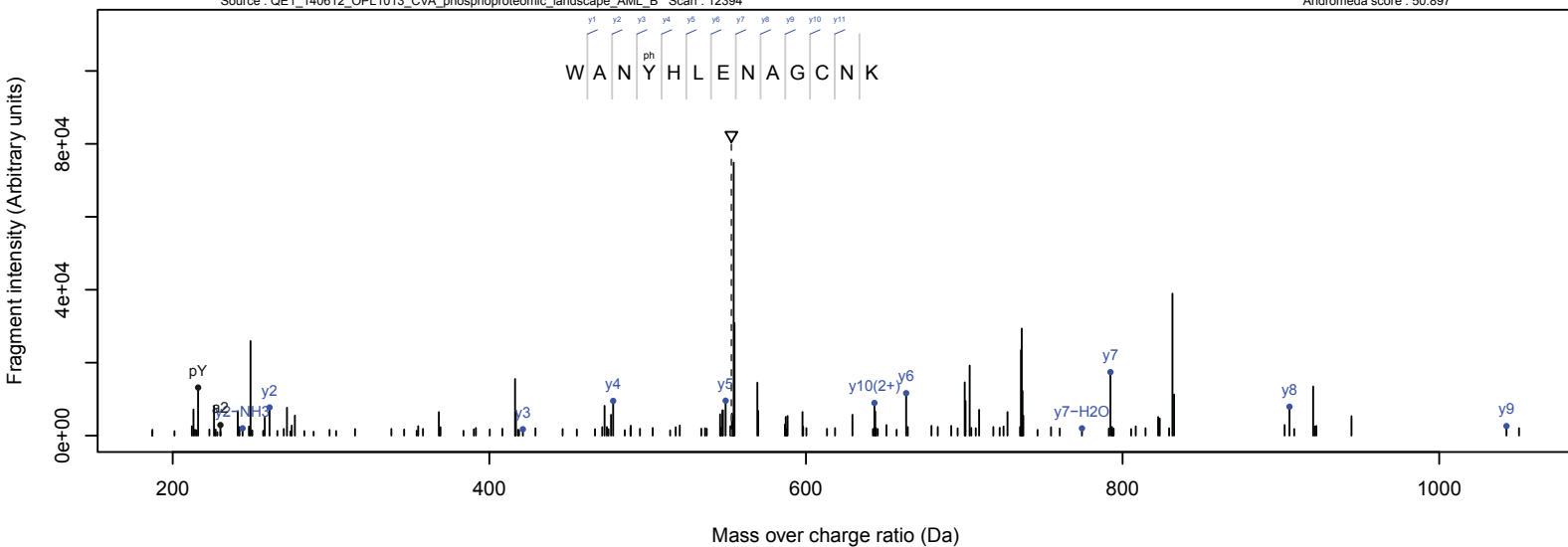
Andromeda score : 50.897



## WANY(ph)HLENAGCNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12394

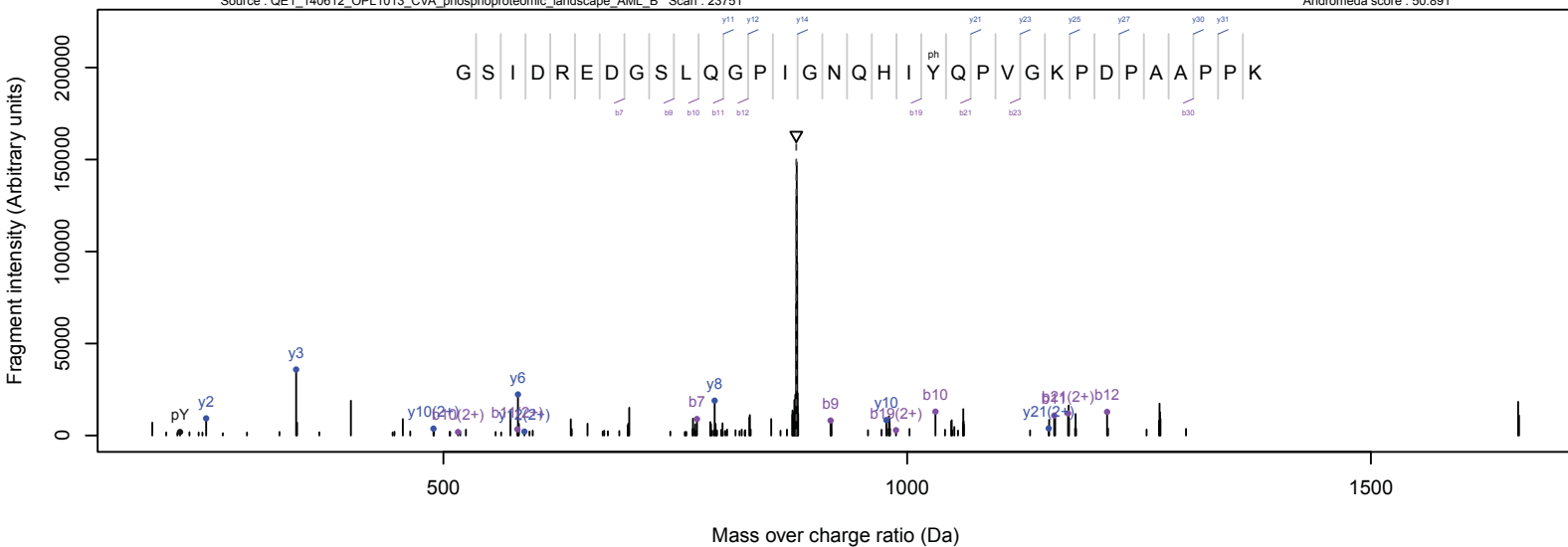
Andromeda score : 50.897



## GSIDREDGSLQGPIGNQH(ph)QPVGKPDPAAPPK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23751

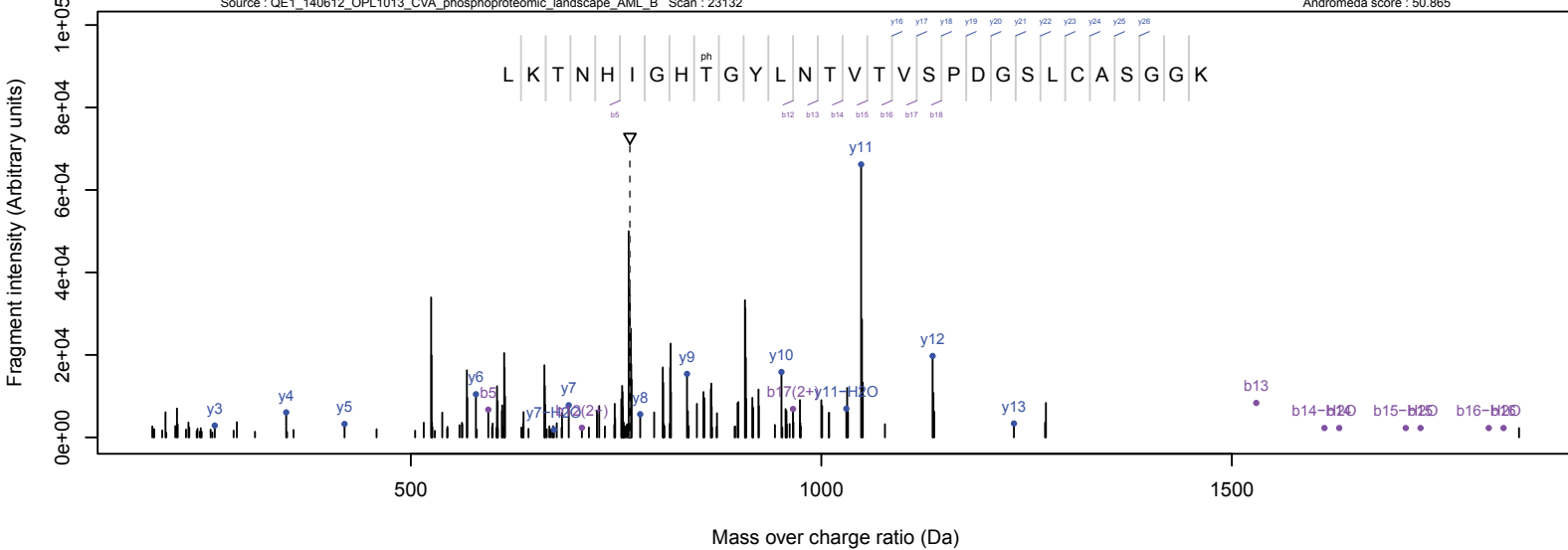
Andromeda score : 50.891



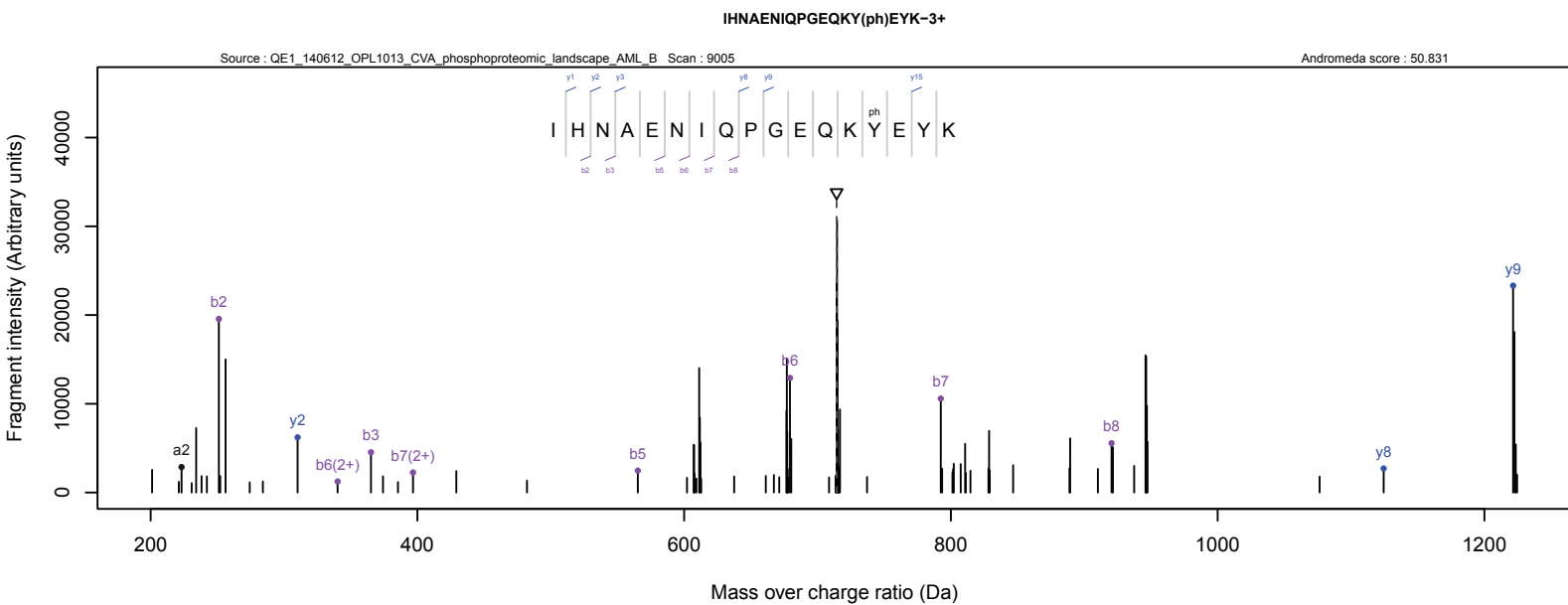
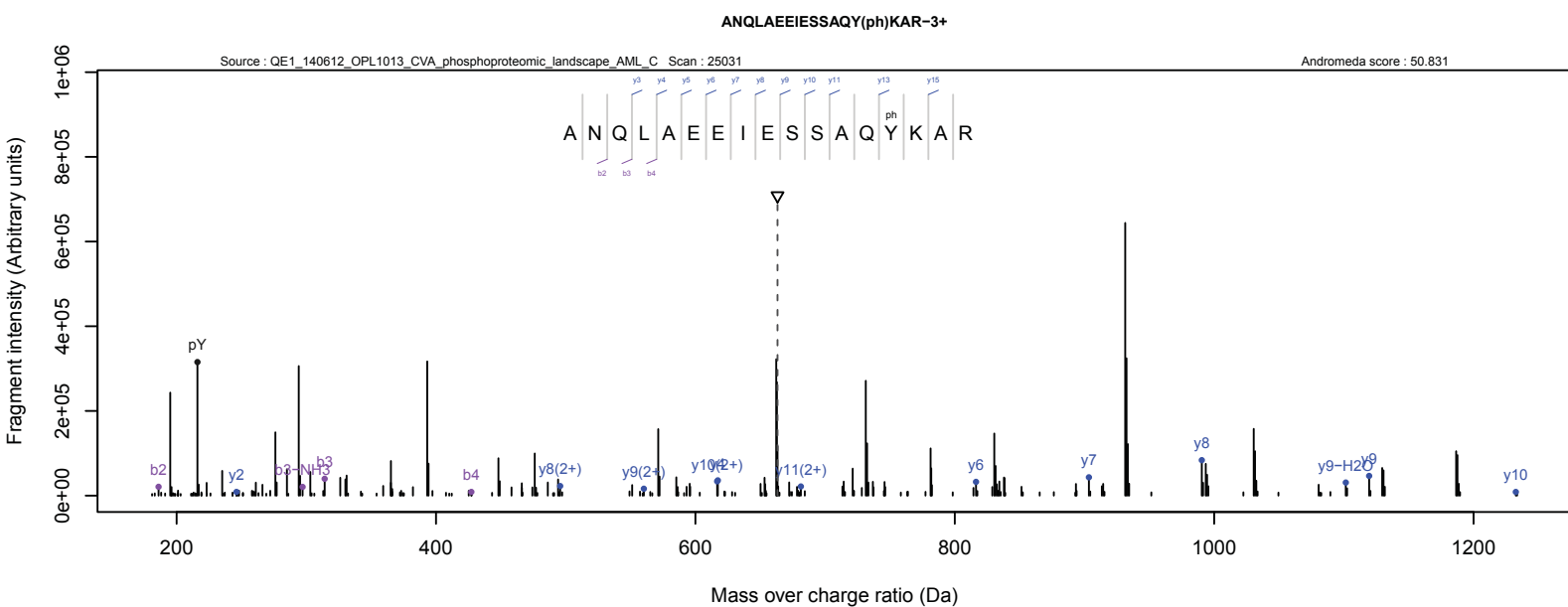
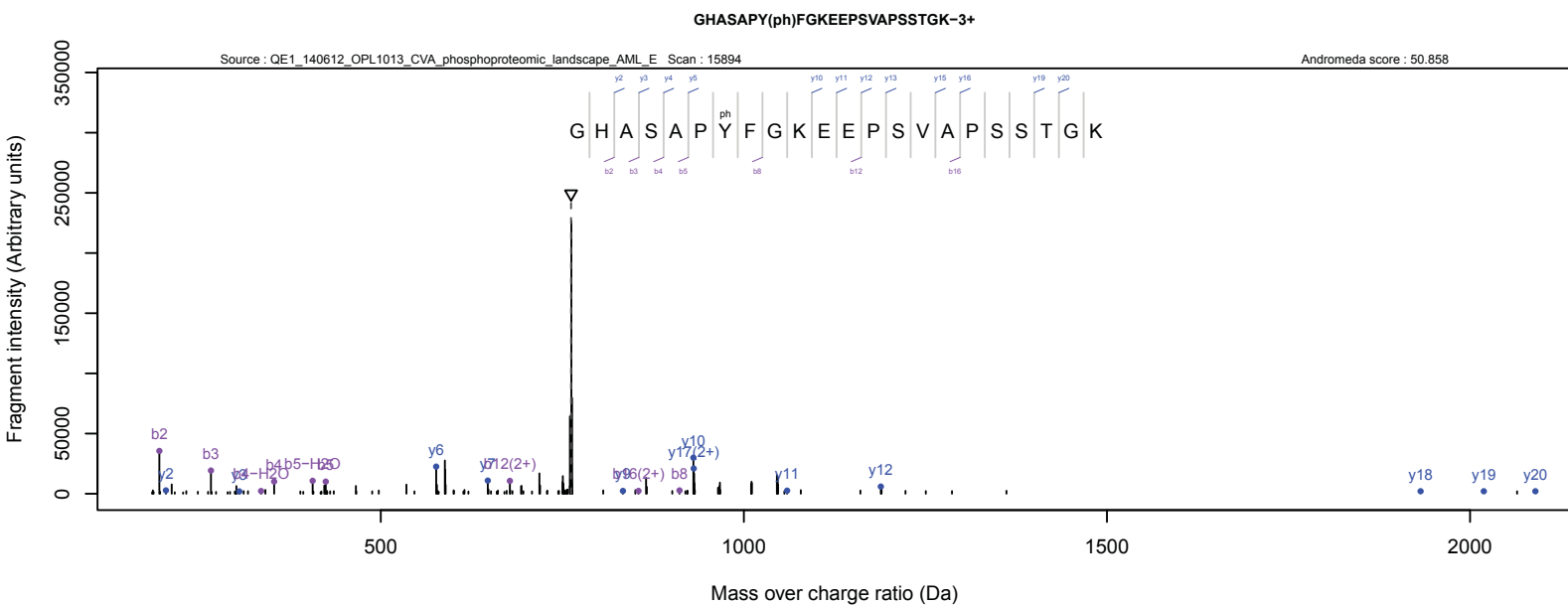
## LKTNHIGHT(ph)GYLNTVTVSPDGLCASGGK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23132

Andromeda score : 50.865



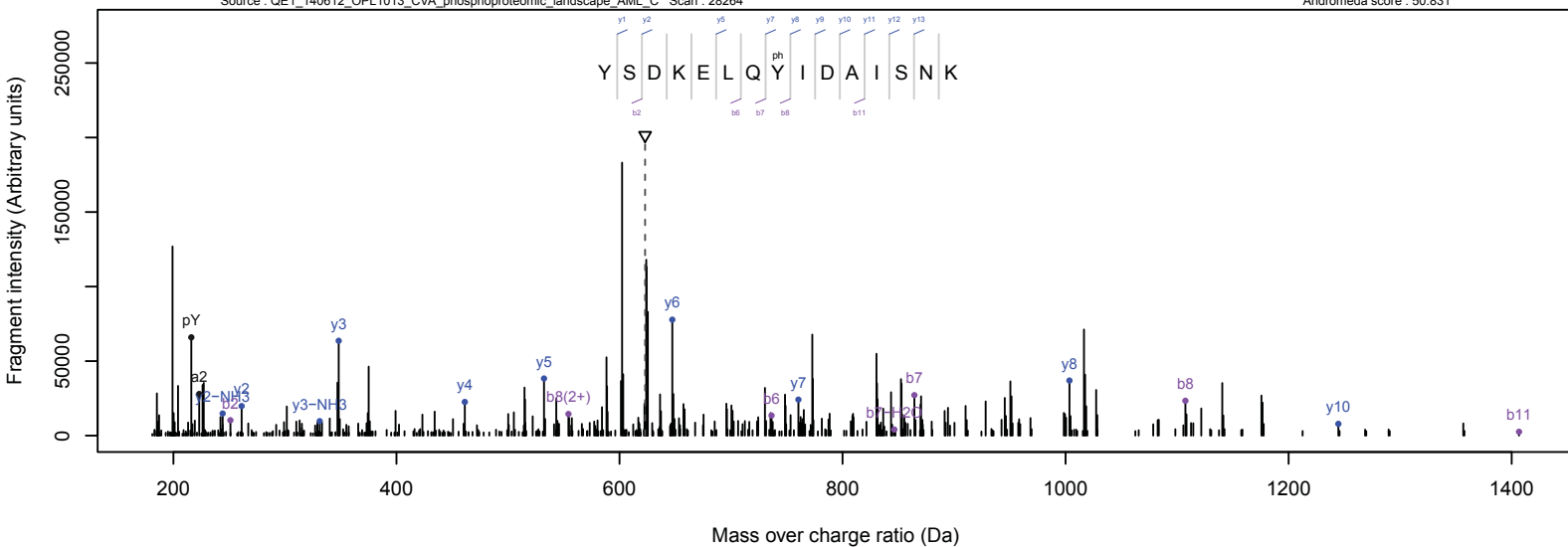




## YSDKELQY(ph)IDAI SNK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 28264

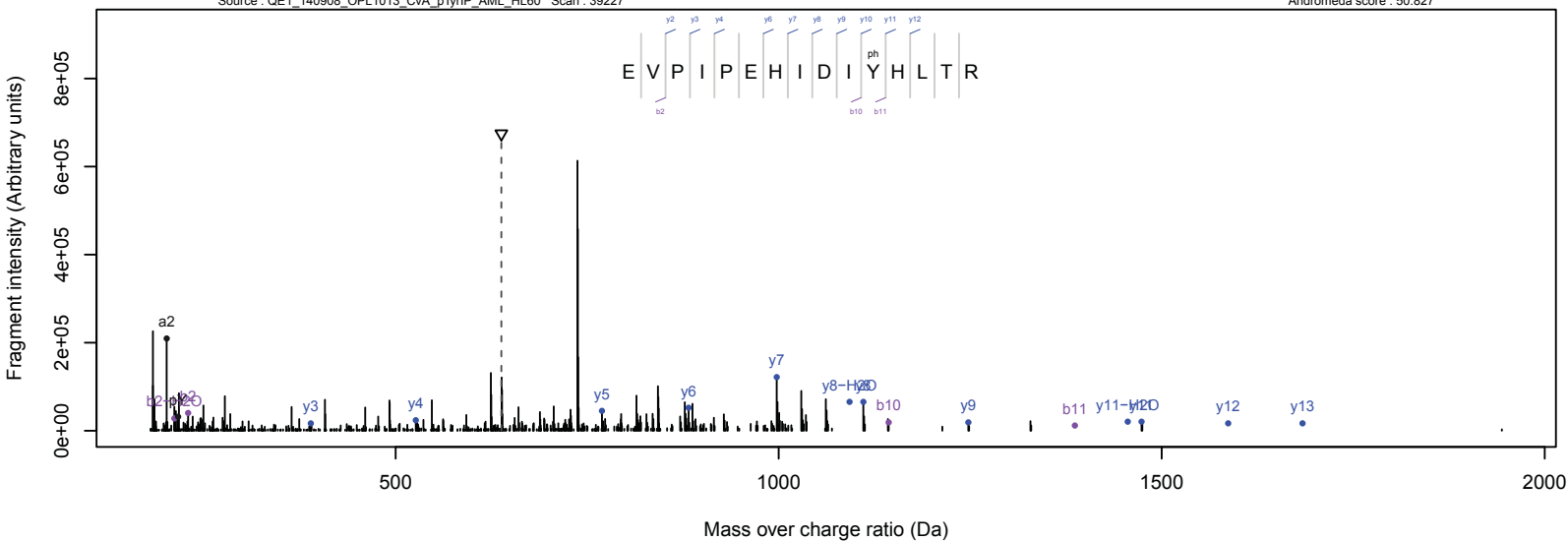
Andromeda score: 50.831



## EVPIPEHIDYHLTR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan: 39227

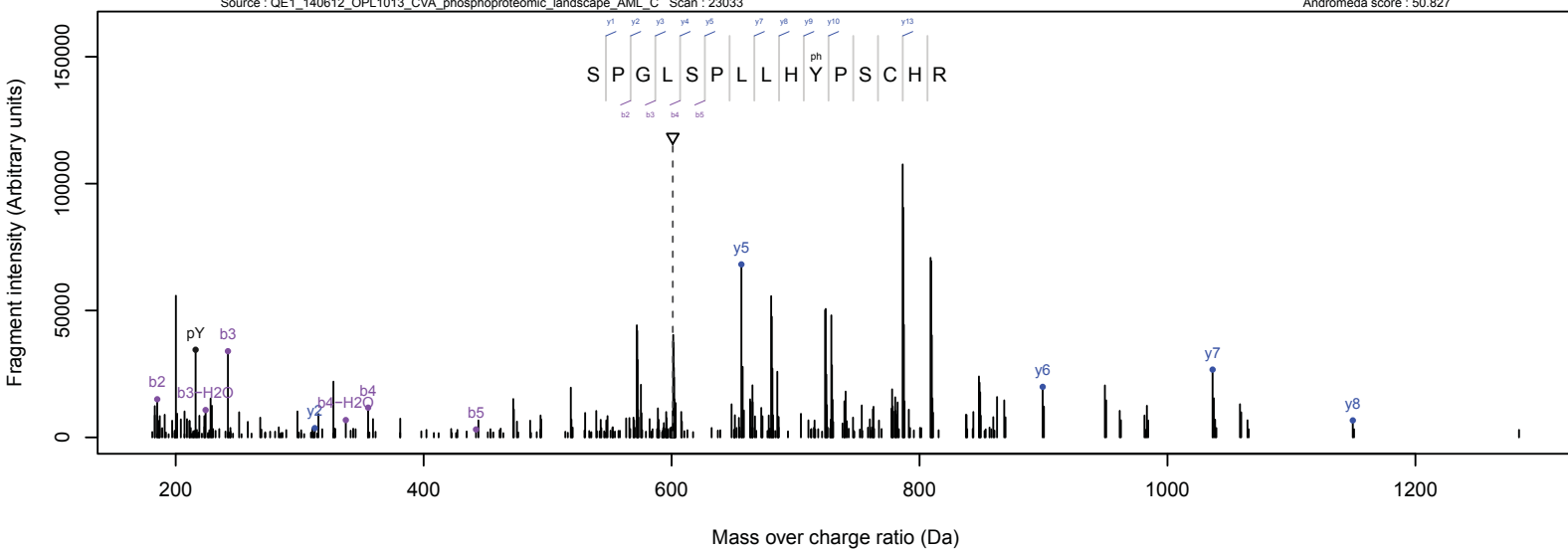
Andromeda score: 50.827



## SPGLSPLLHYPSCHR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23033

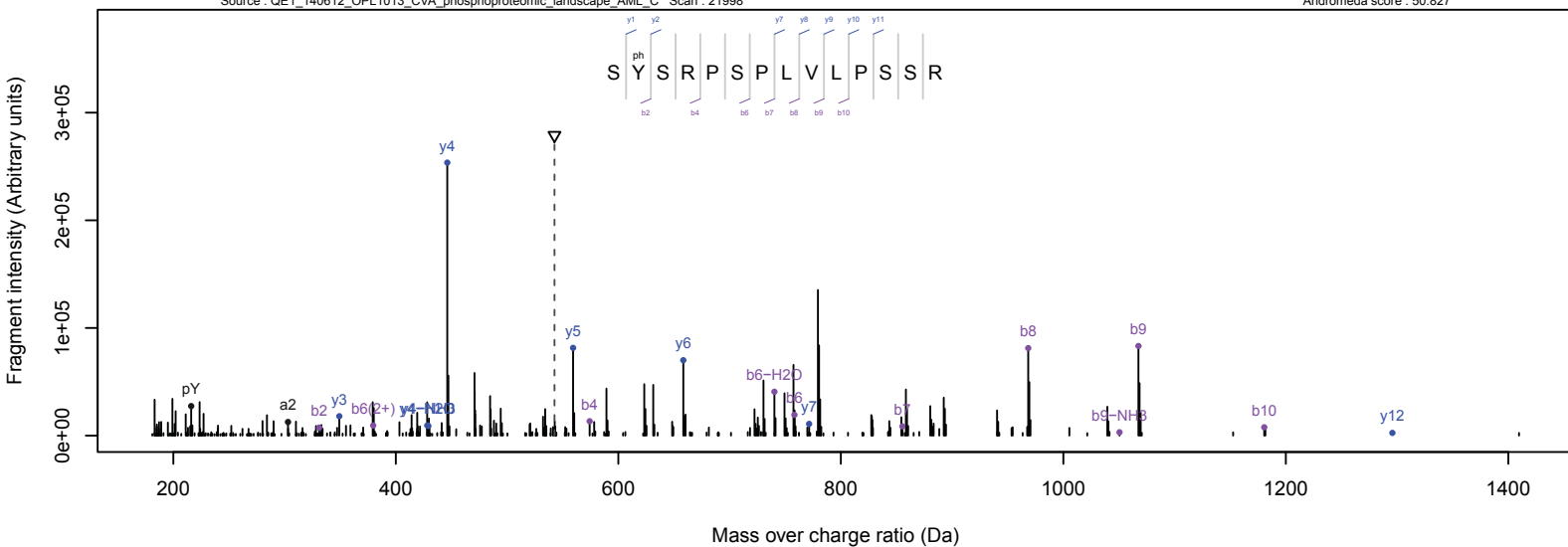
Andromeda score: 50.827



**SY(ph)SRPSPLVLPSSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21998

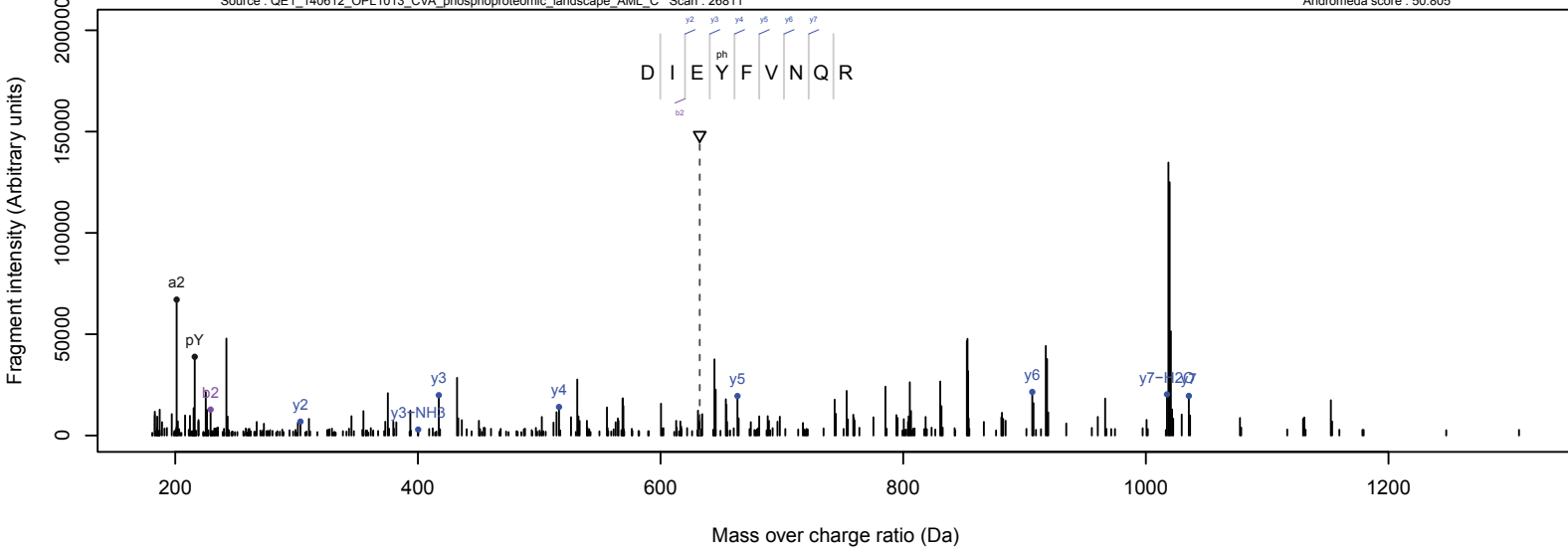
Andromeda score : 50.827



**DIEY(ph)FVNQR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26811

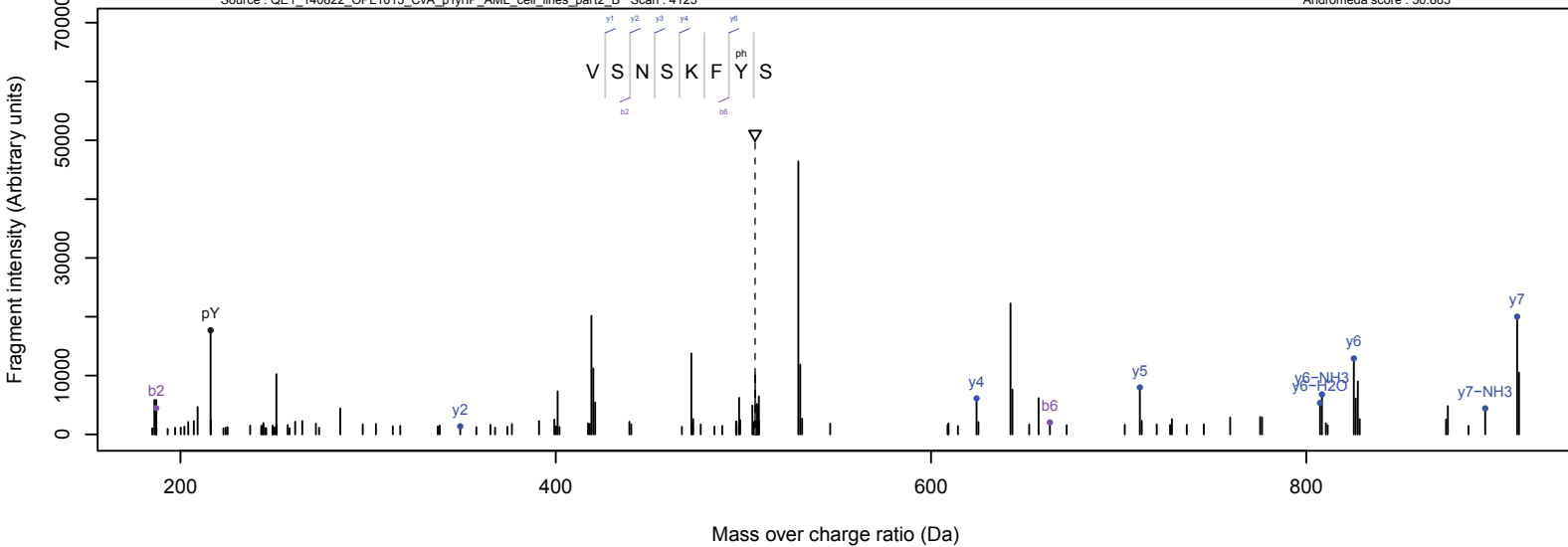
Andromeda score : 50.805



**VSNSKFY(ph)S-2+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 4125

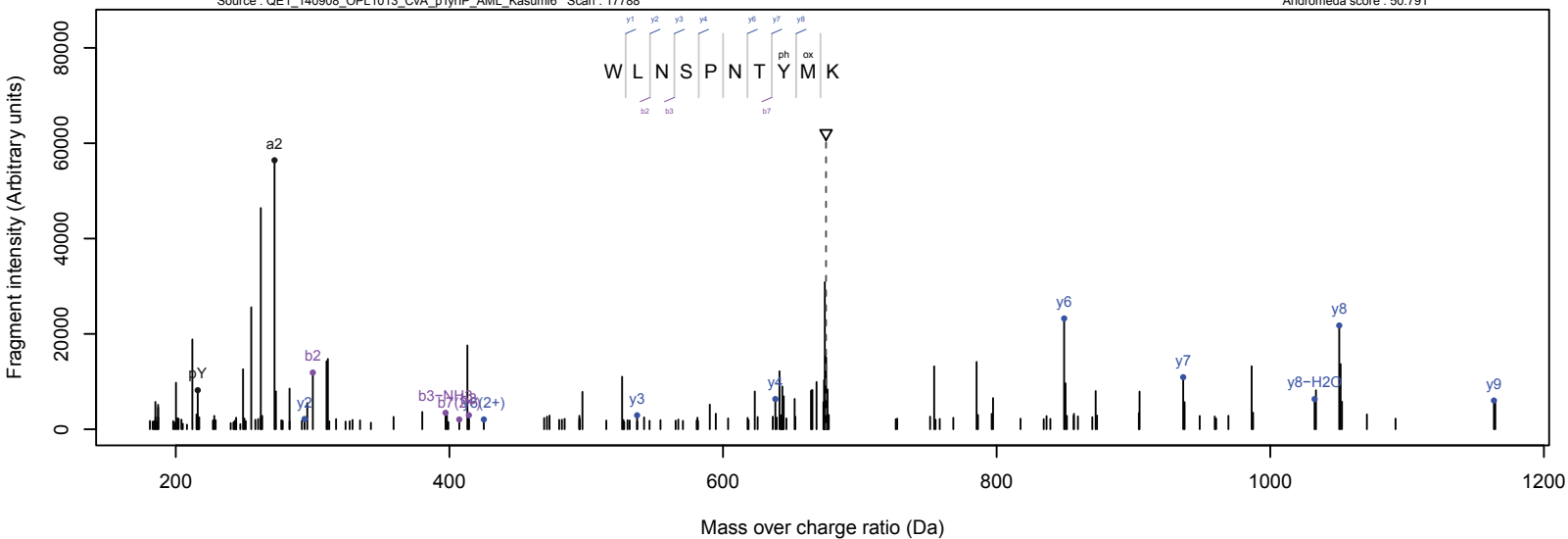
Andromeda score : 50.805



## WLNSPNTY(ph)M(ox)K-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 17788

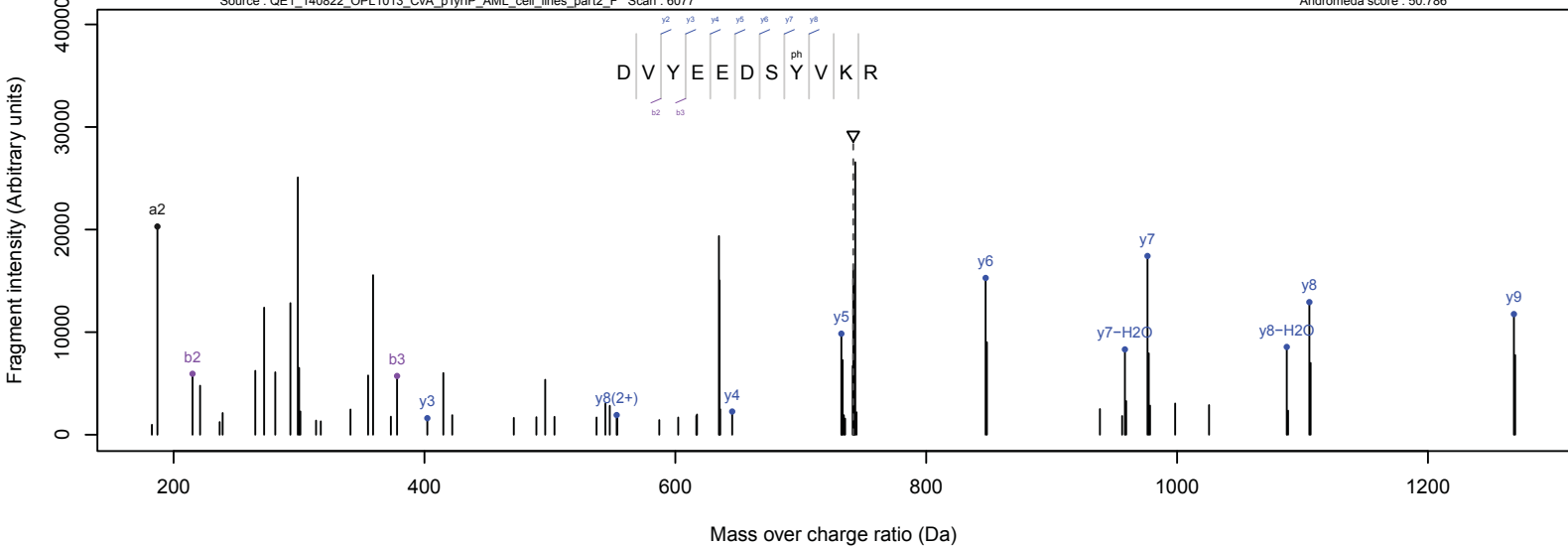
Andromeda score: 50.791



## DVYEEDSY(ph)VKR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan: 6077

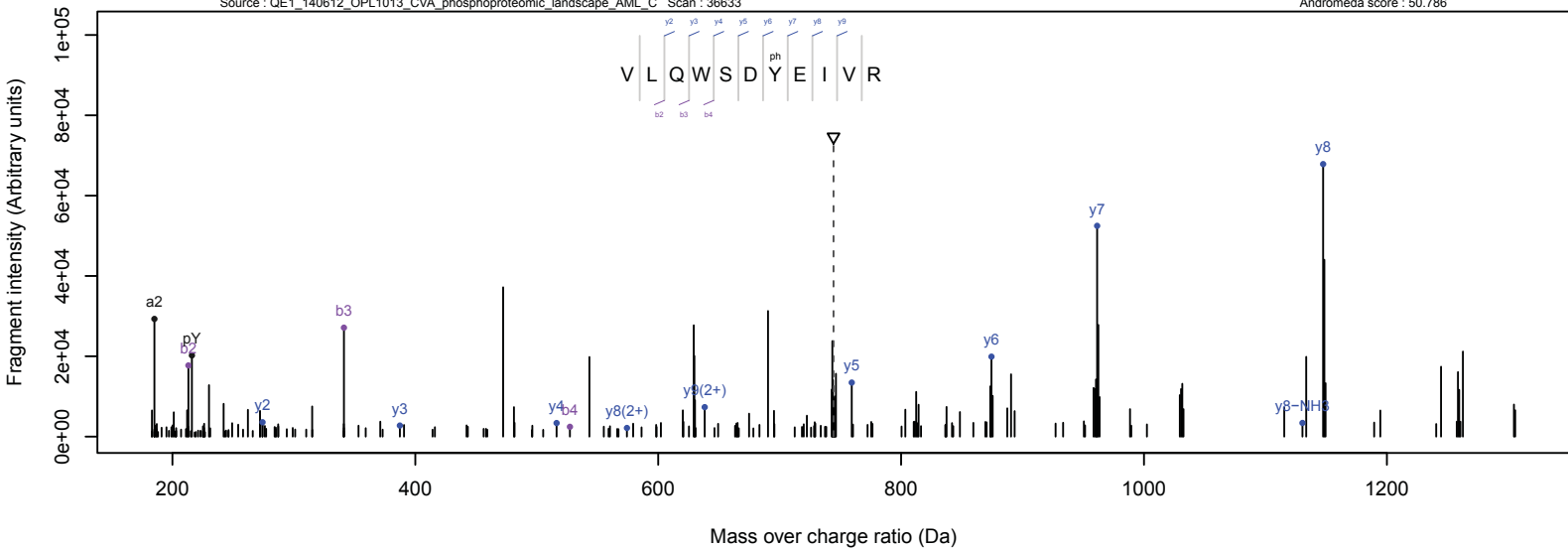
Andromeda score: 50.786



## VLQWSDY(ph)EIVR-2+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 36633

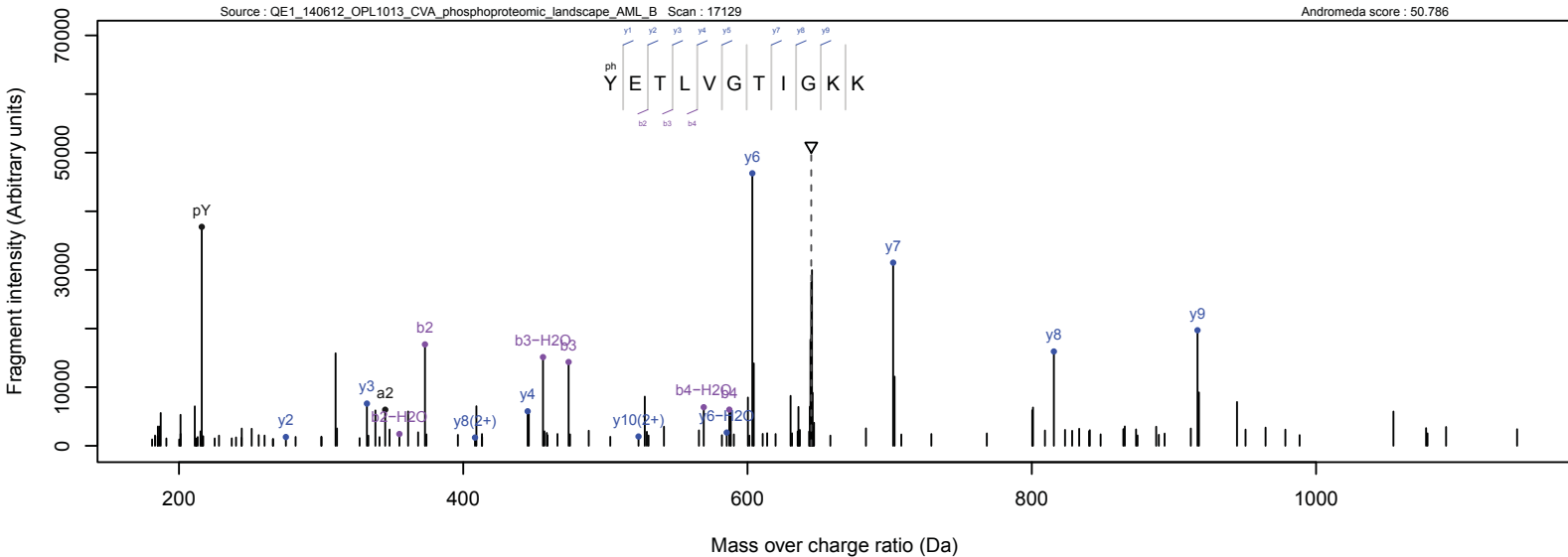
Andromeda score: 50.786



**Y(ph)ETLVGTIGKK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17129

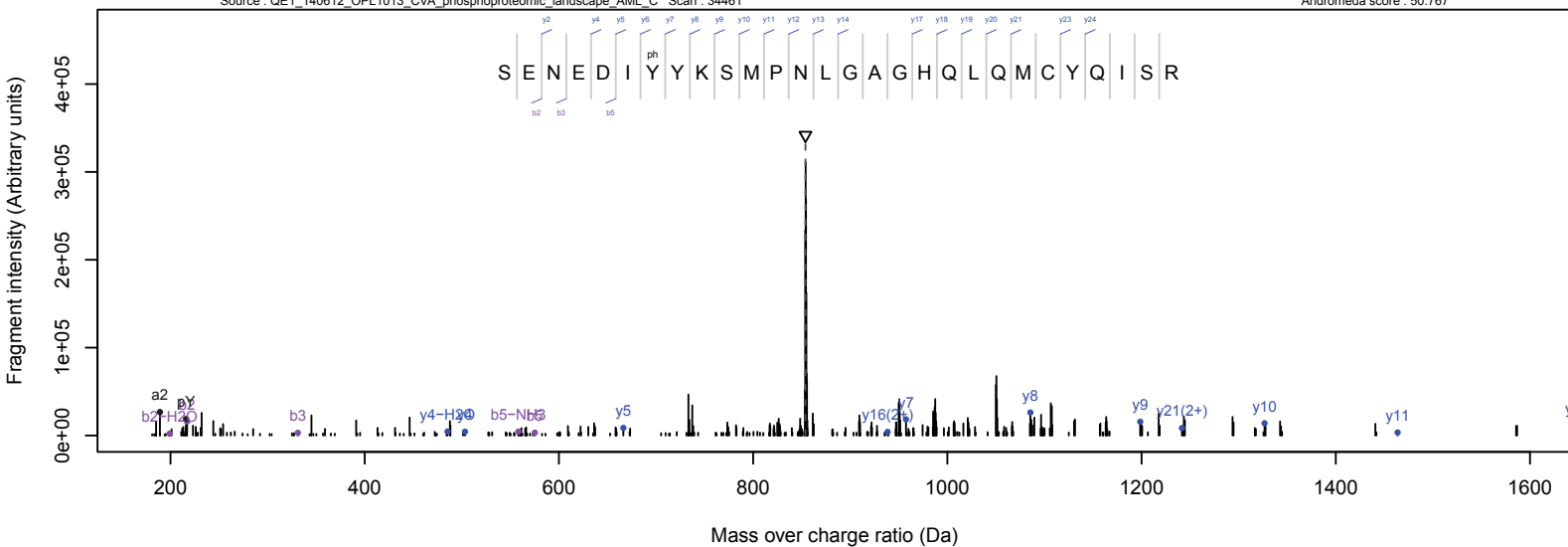
Andromeda score : 50.786



**SENEIDIYKSMPLNGAGHQLQMCYQISR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34461

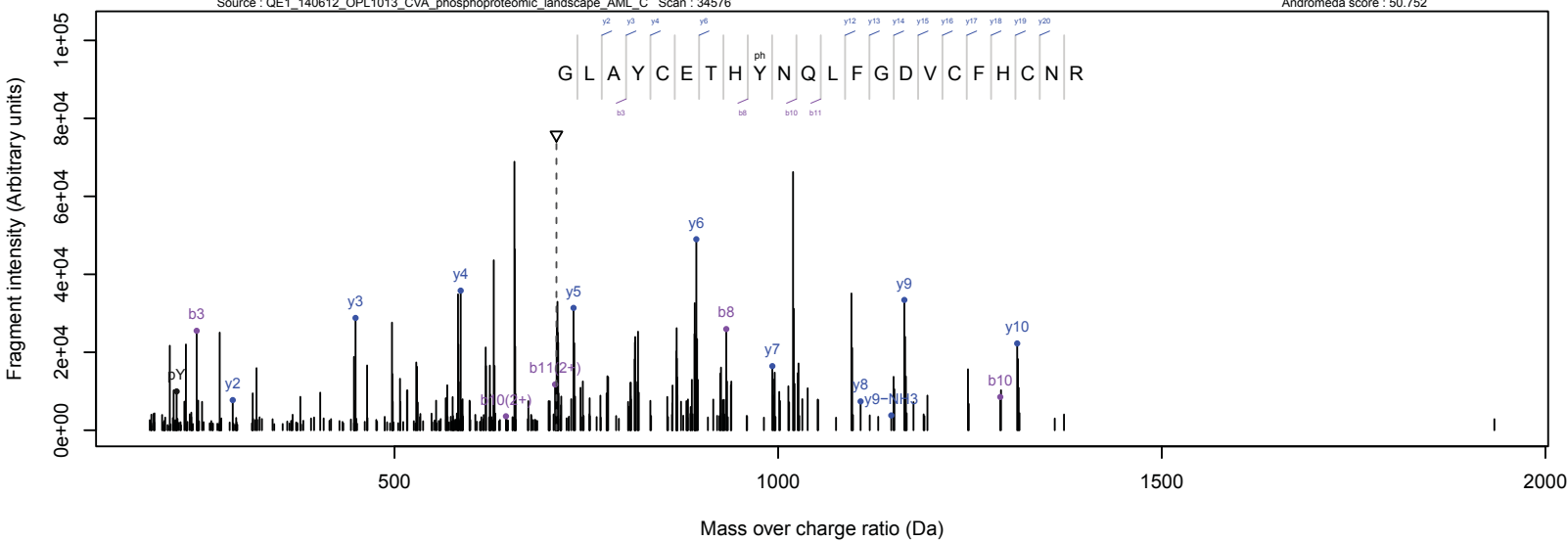
Andromeda score : 50.767

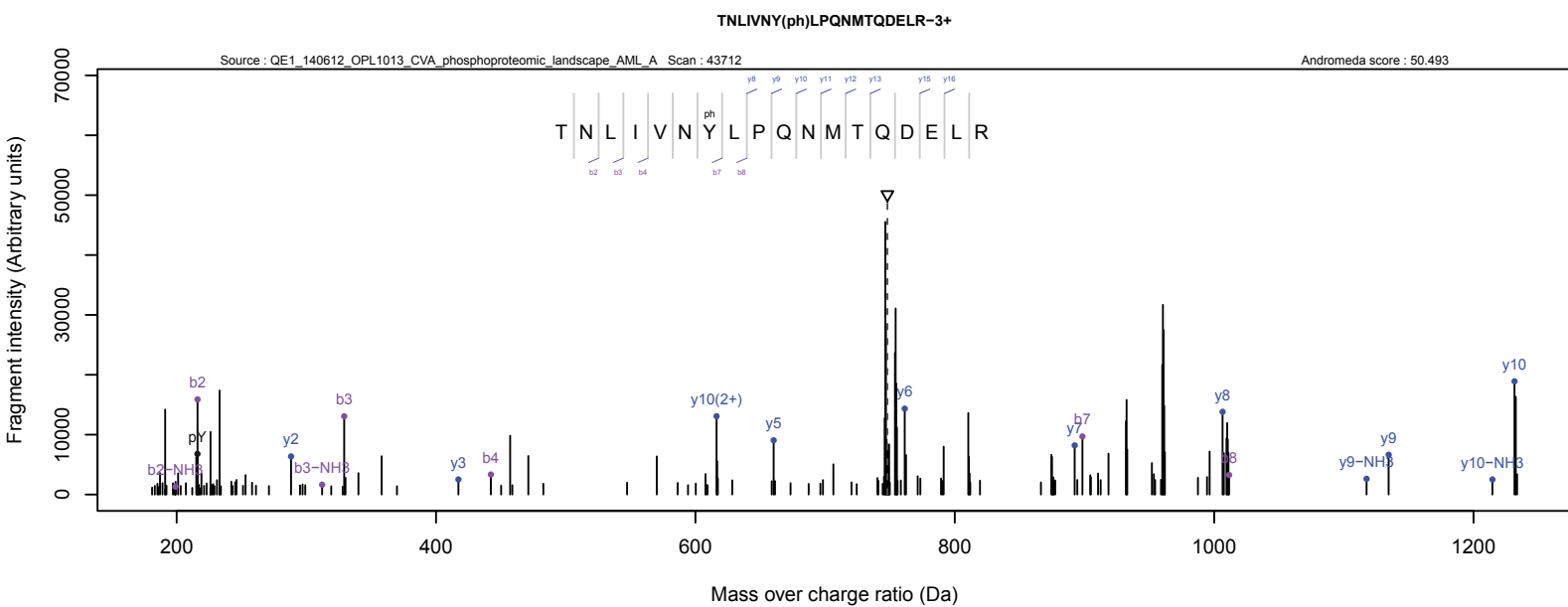
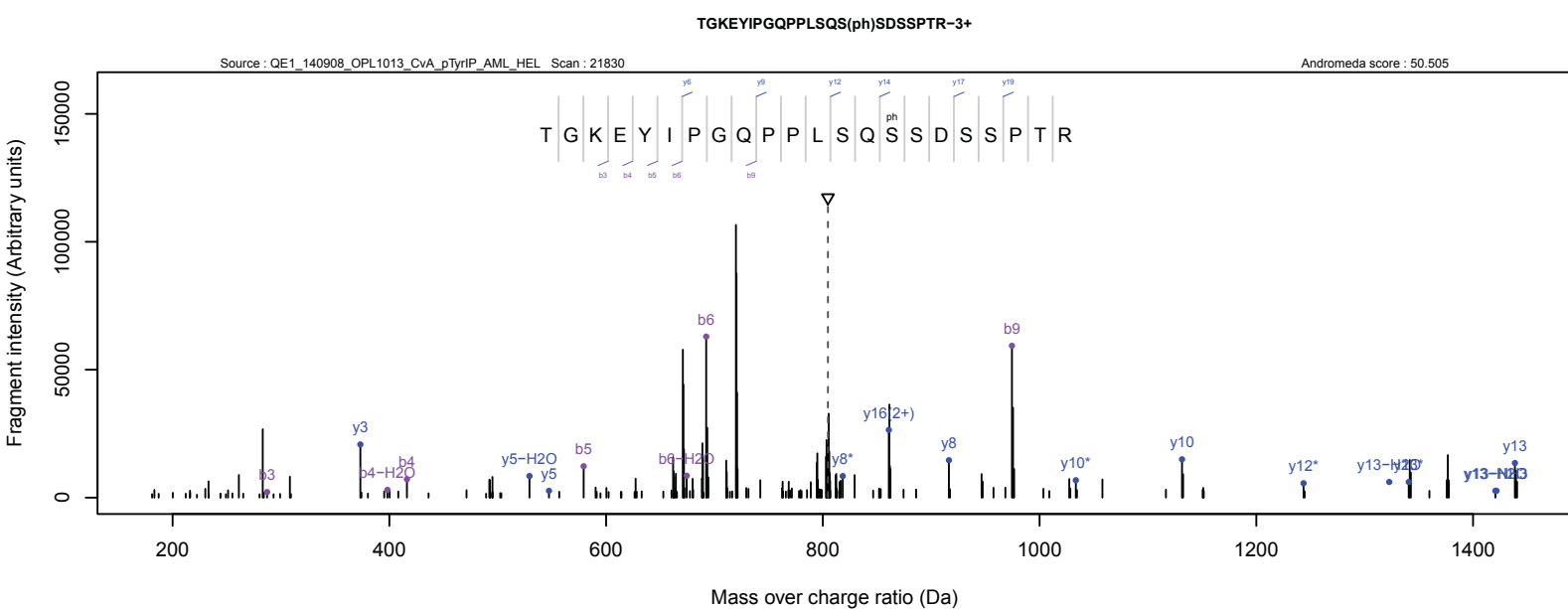
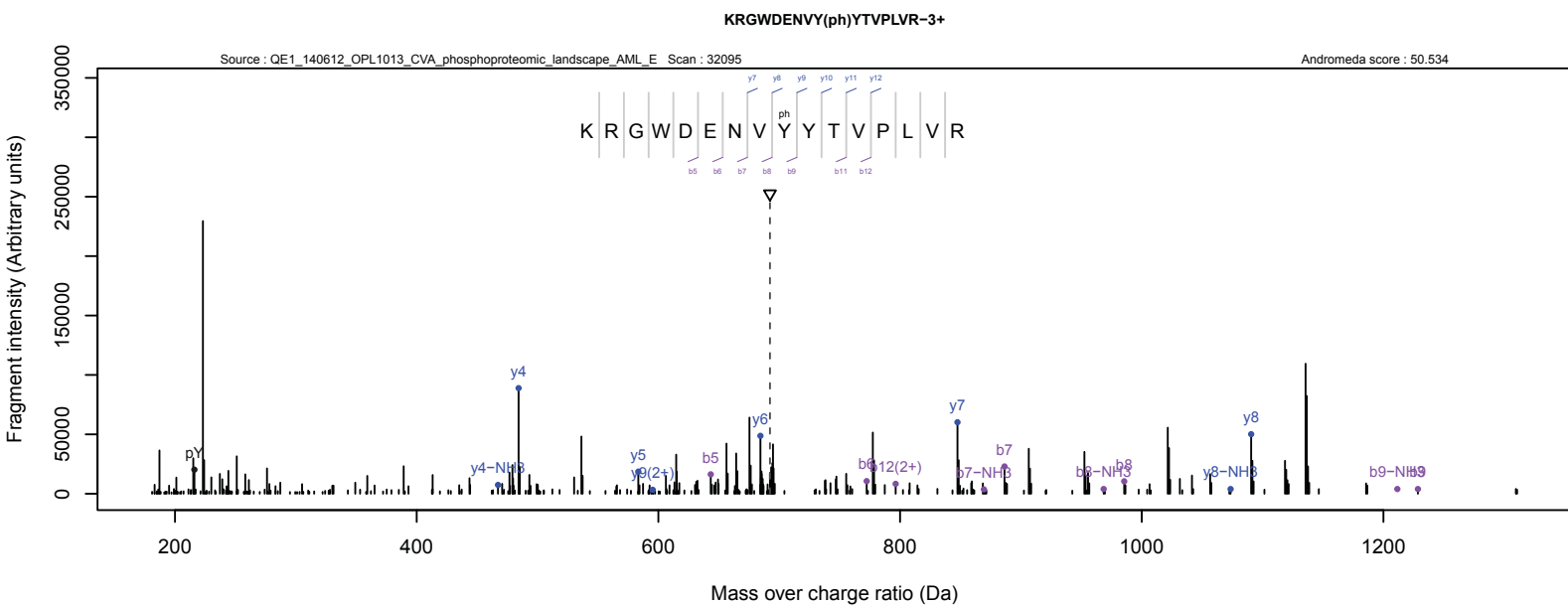


**GLAYCETHY(N)QLFGDVCFHCNR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34576

Andromeda score : 50.752

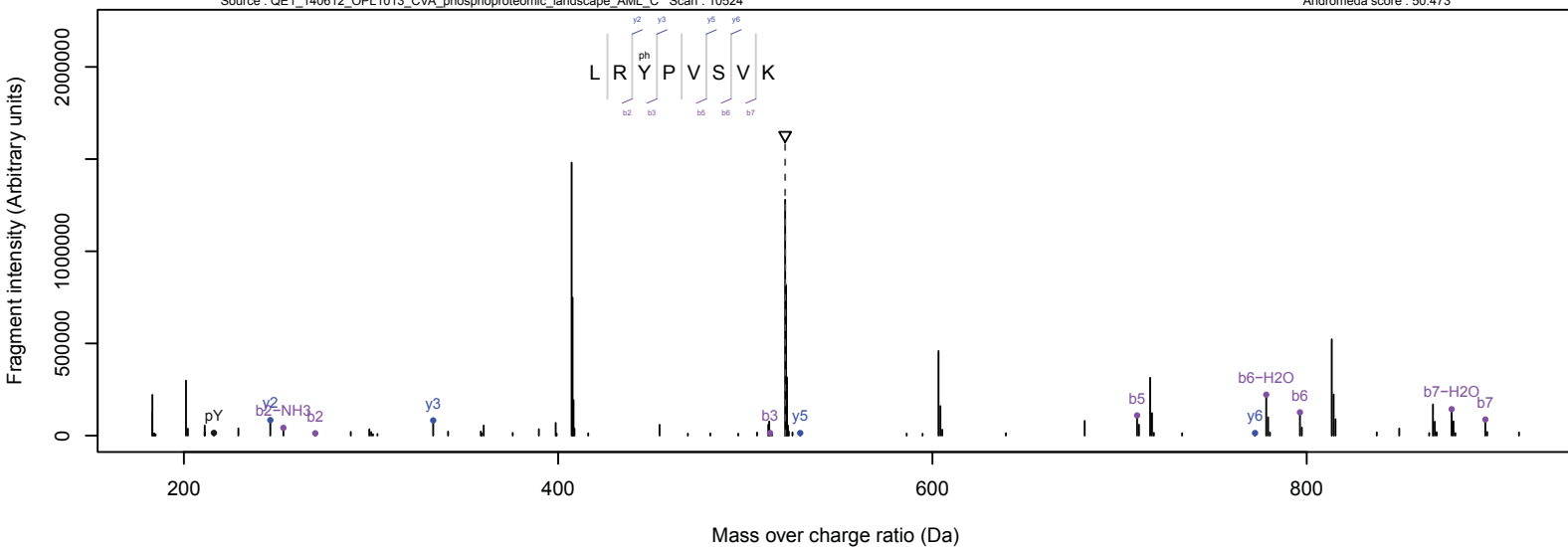




## LRY(ph)PVSVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10524

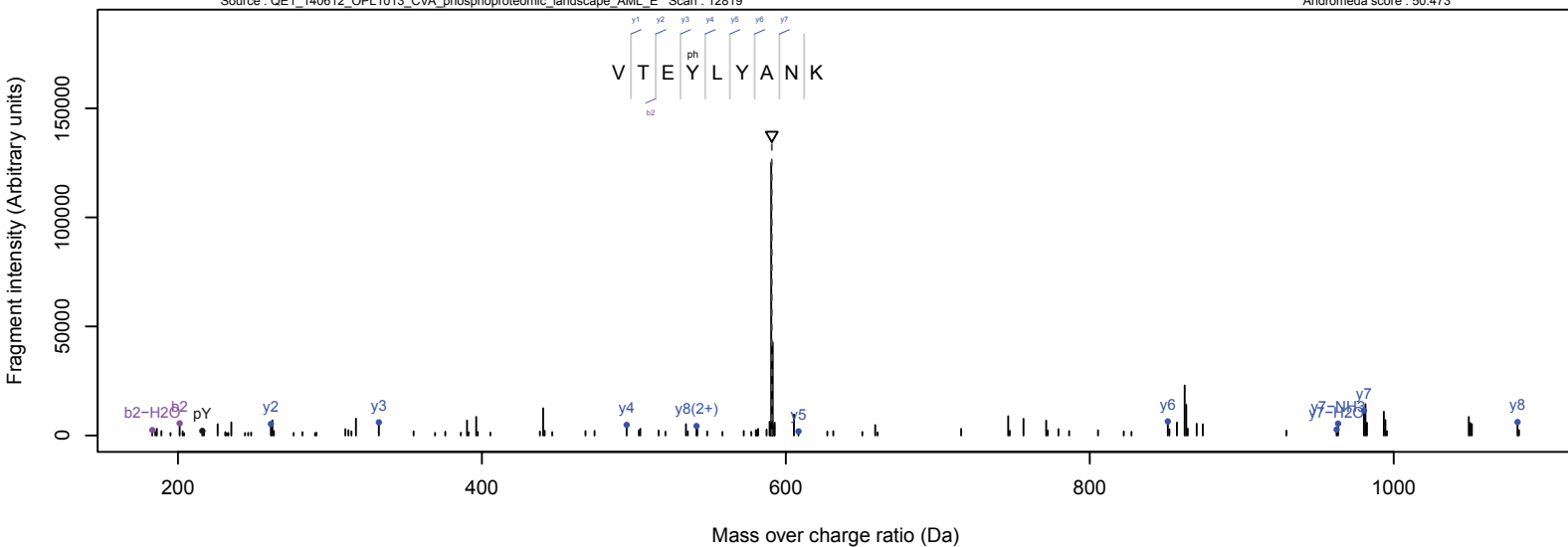
Andromeda score : 50.473



## VTEY(ph)LYANK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 12819

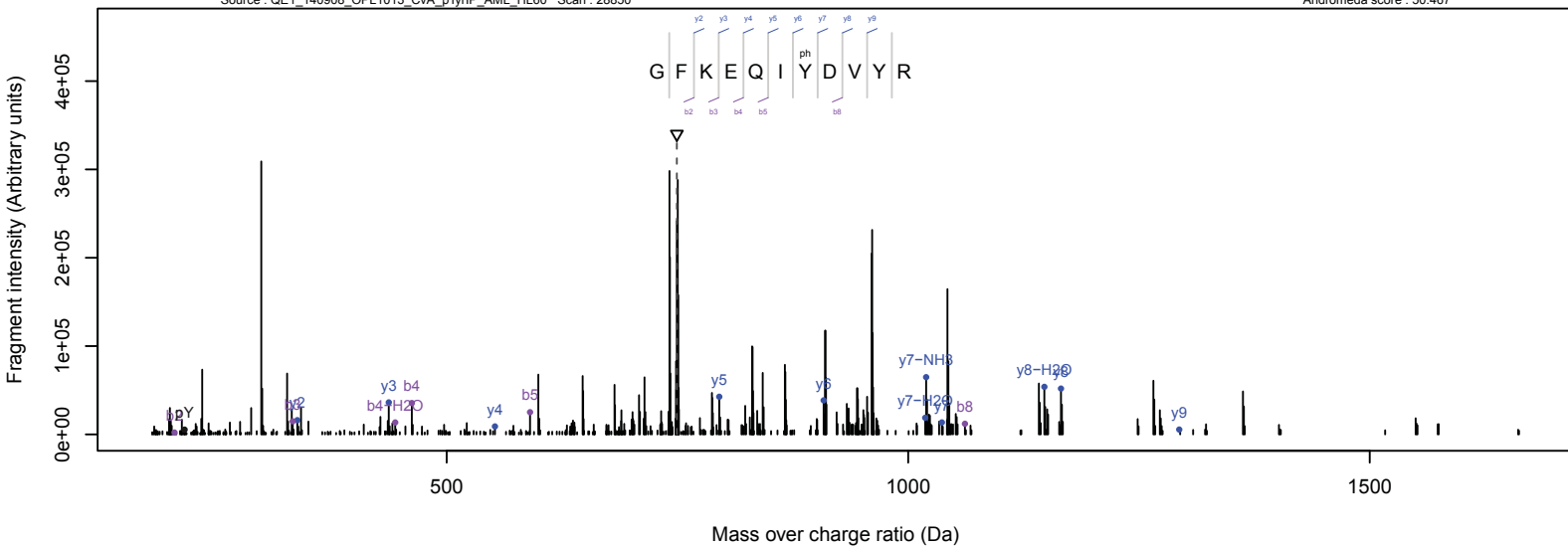
Andromeda score : 50.473



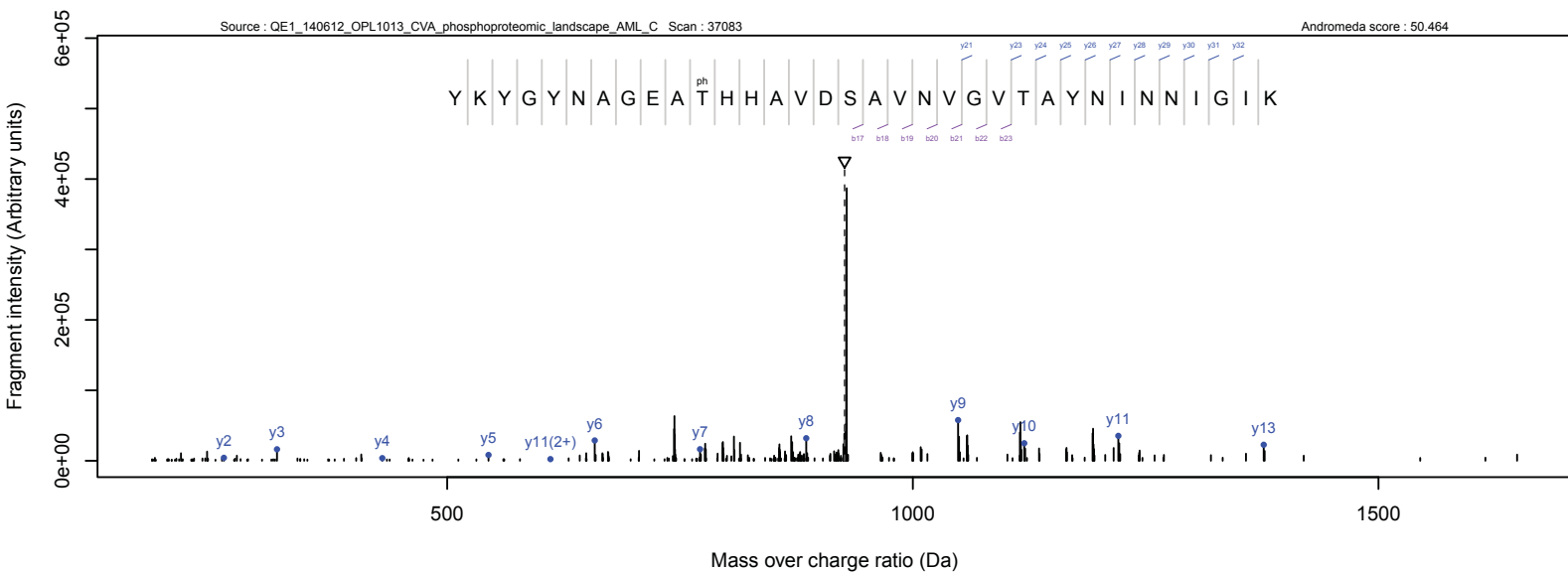
## GFKEQIY(ph)DVYR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 28850

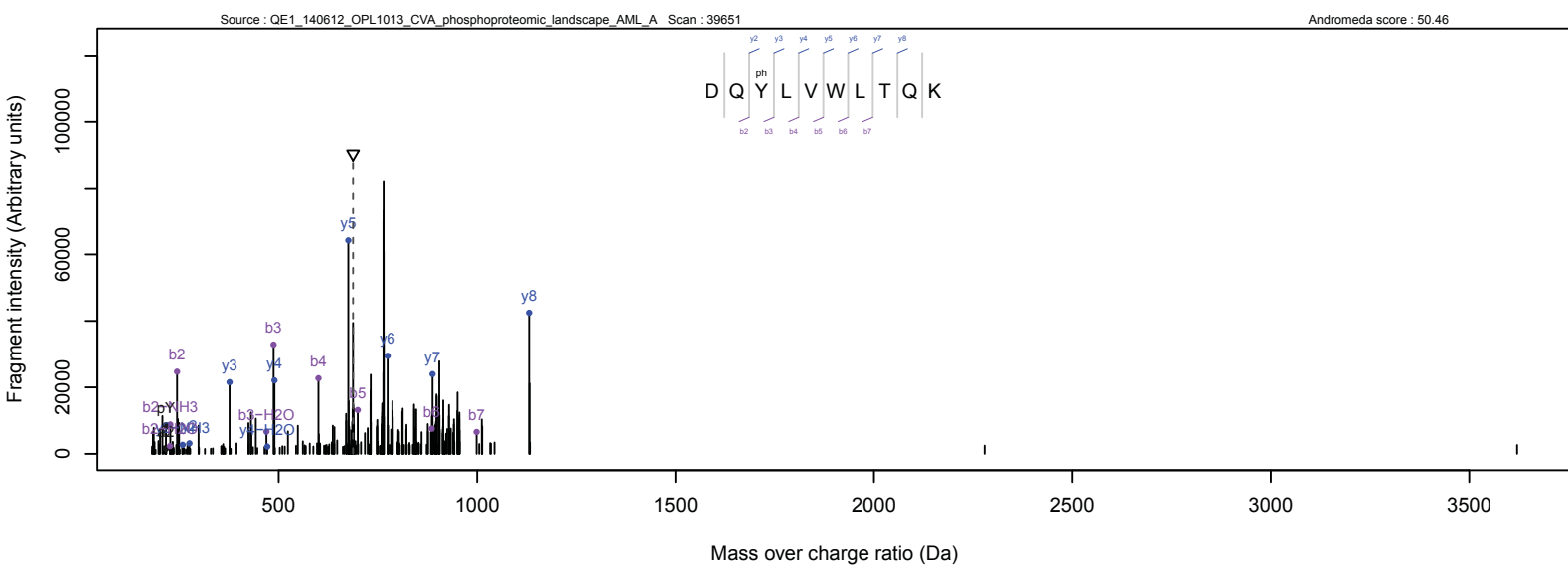
Andromeda score : 50.467



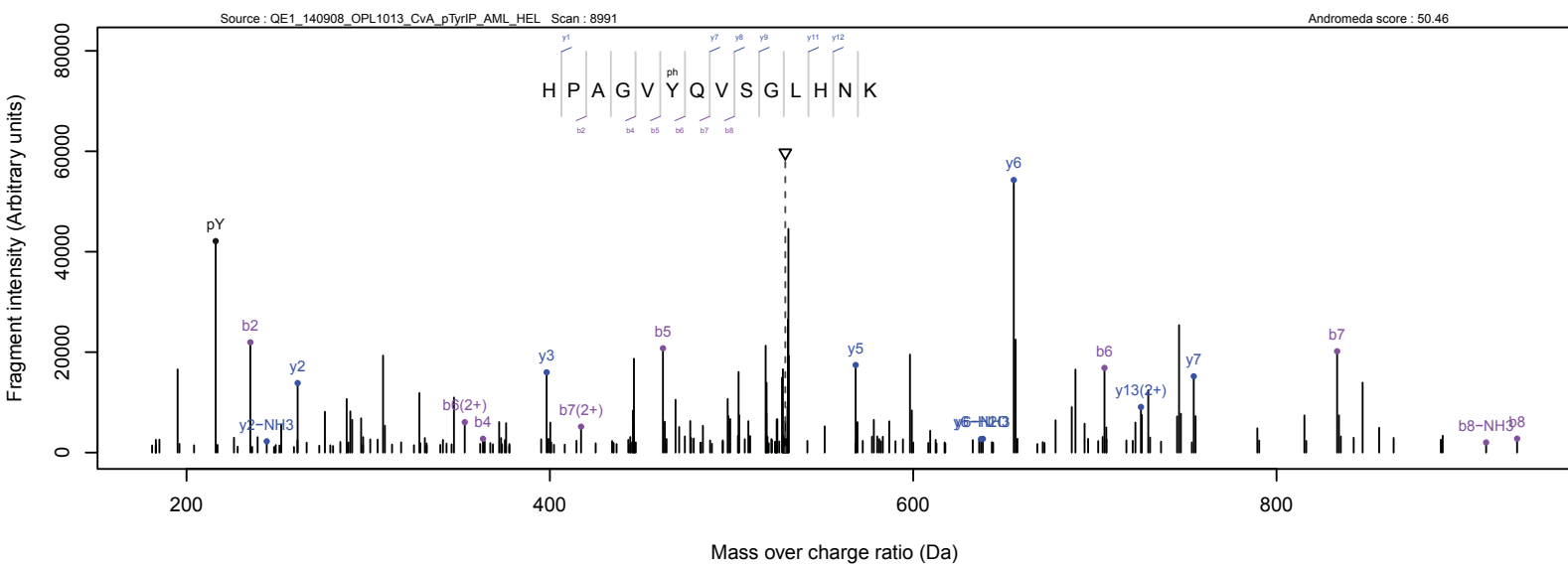
## YKYGNAGEAT(ph)HHAVDSDAVNVGVTA YNINNIGIK-4+



## DQY(ph)LVWLTQK-2+



## HPAGVY(ph)QVSGLHNK-3+

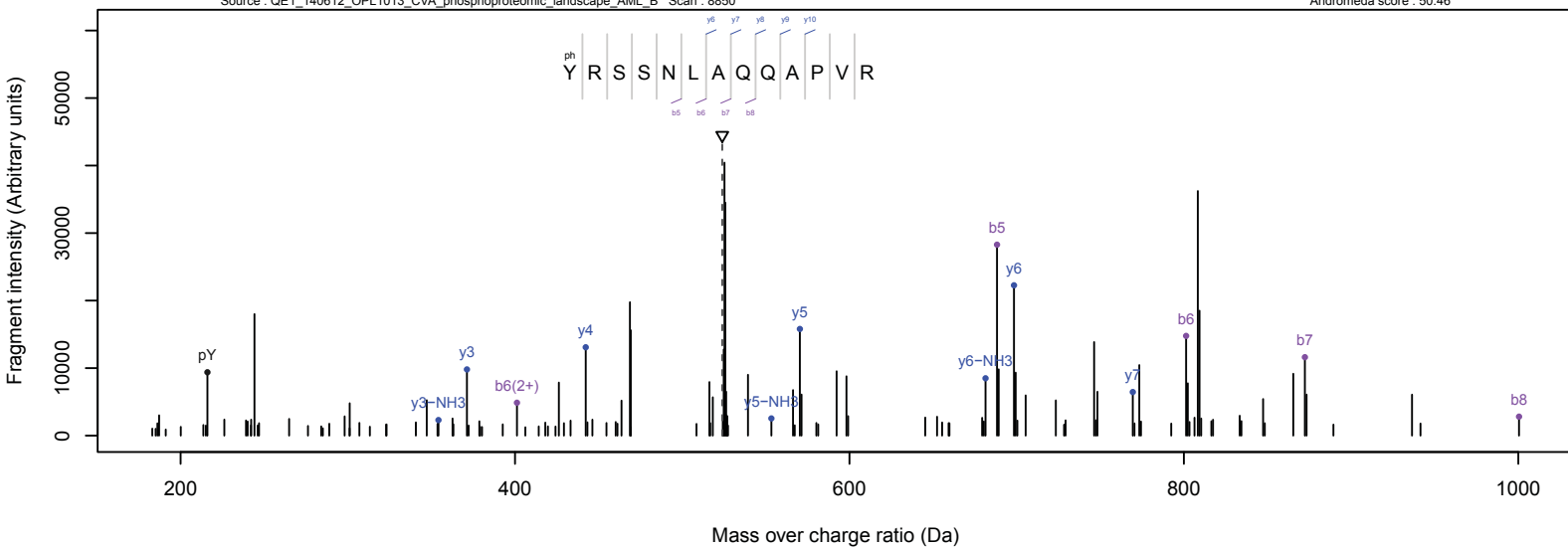




**Y(ph)RSSNLAQQAPVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 8850

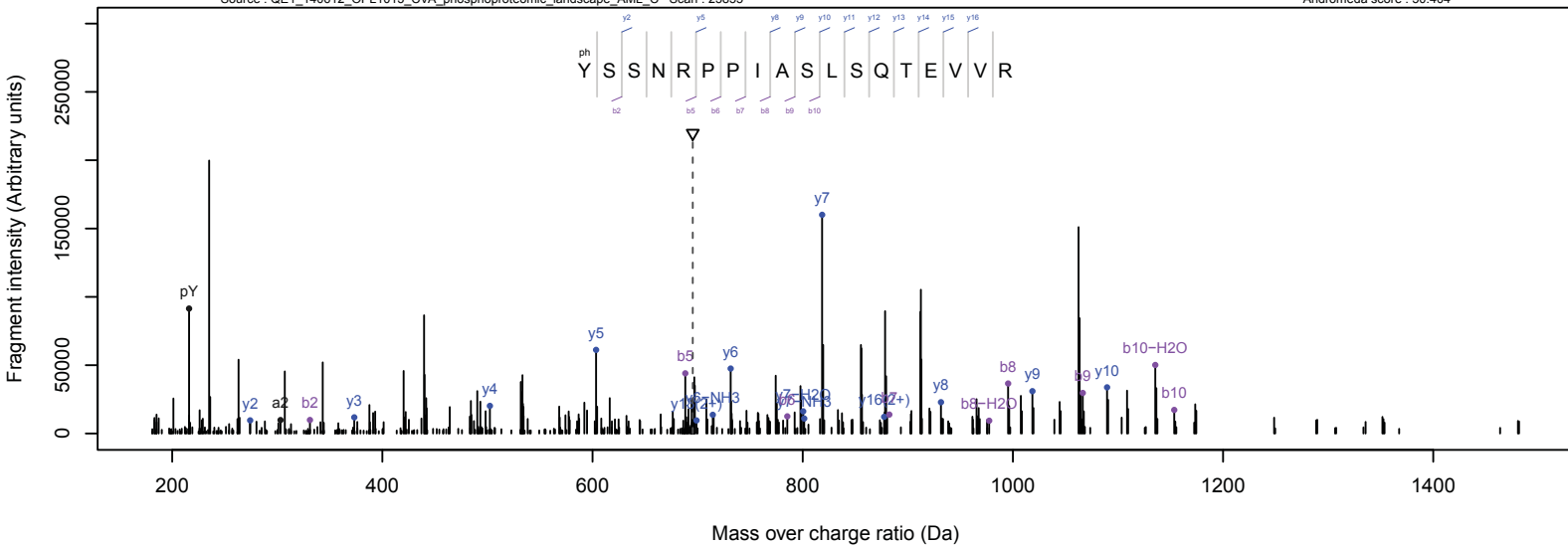
Andromeda score : 50.46



**Y(ph)SSNRPIIASLSQTEVVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25855

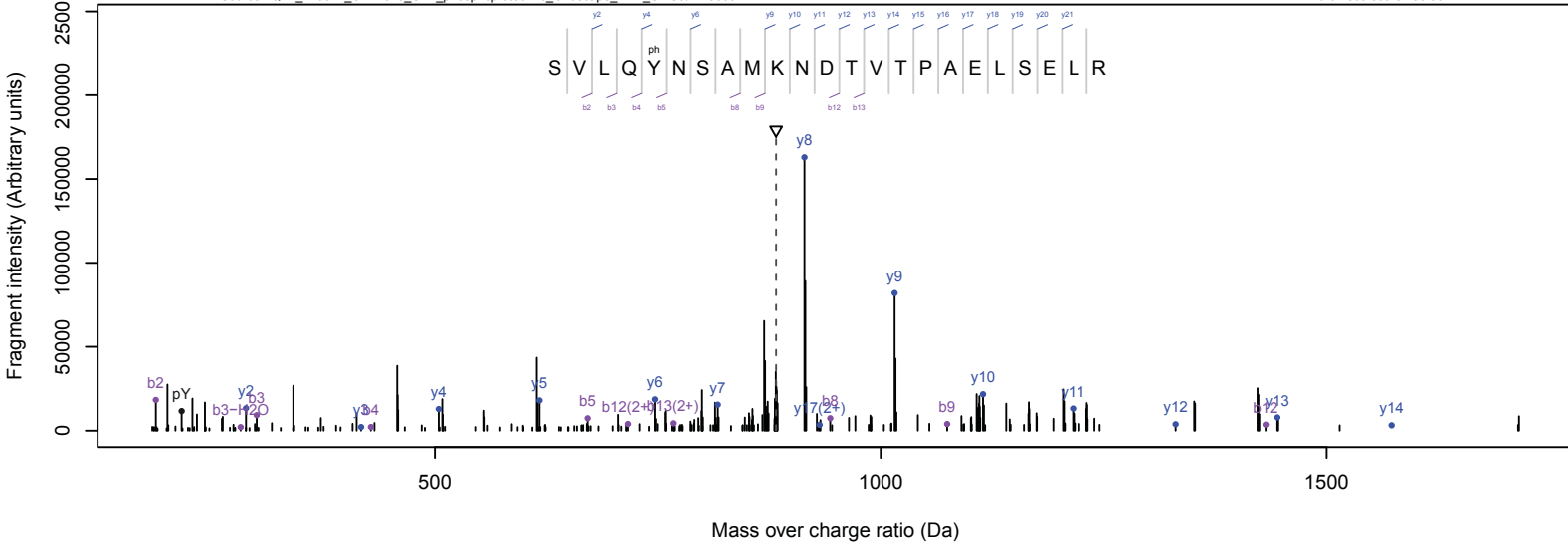
Andromeda score : 50.404



**SVLQY(ph)NSAMKNDTVTPAELSELR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36031

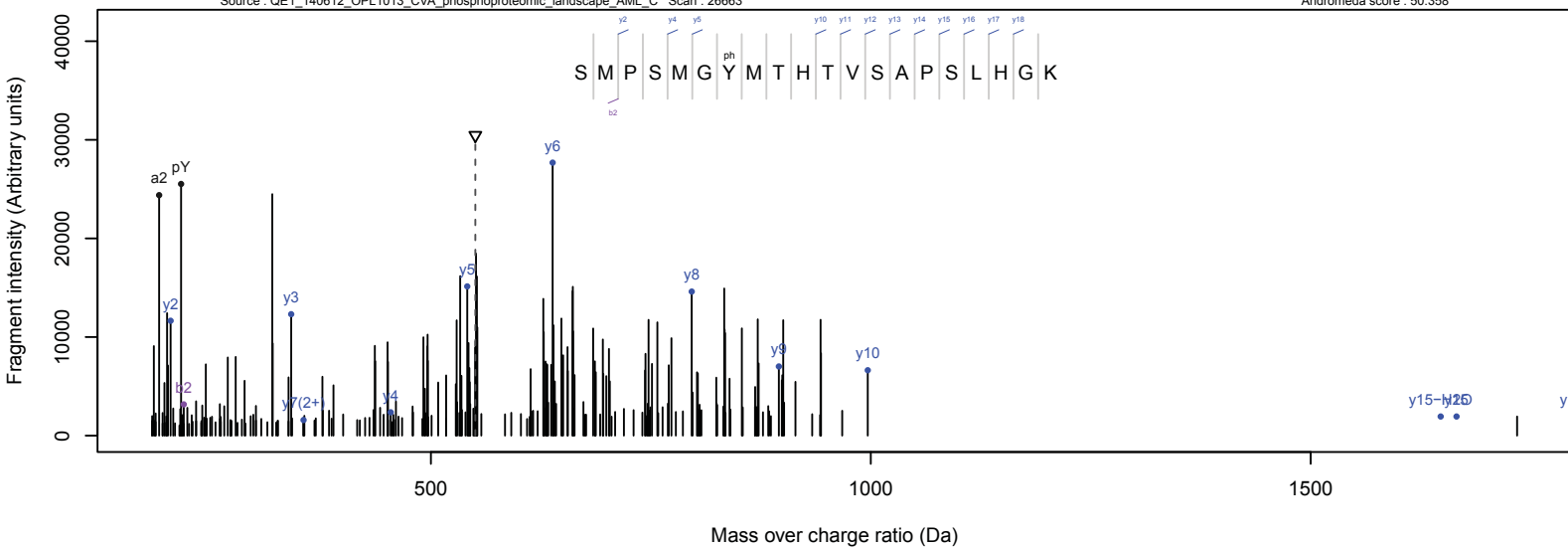
Andromeda score : 50.364



**SMPSMGY(ph)MHTVVSAPSLHGK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26663

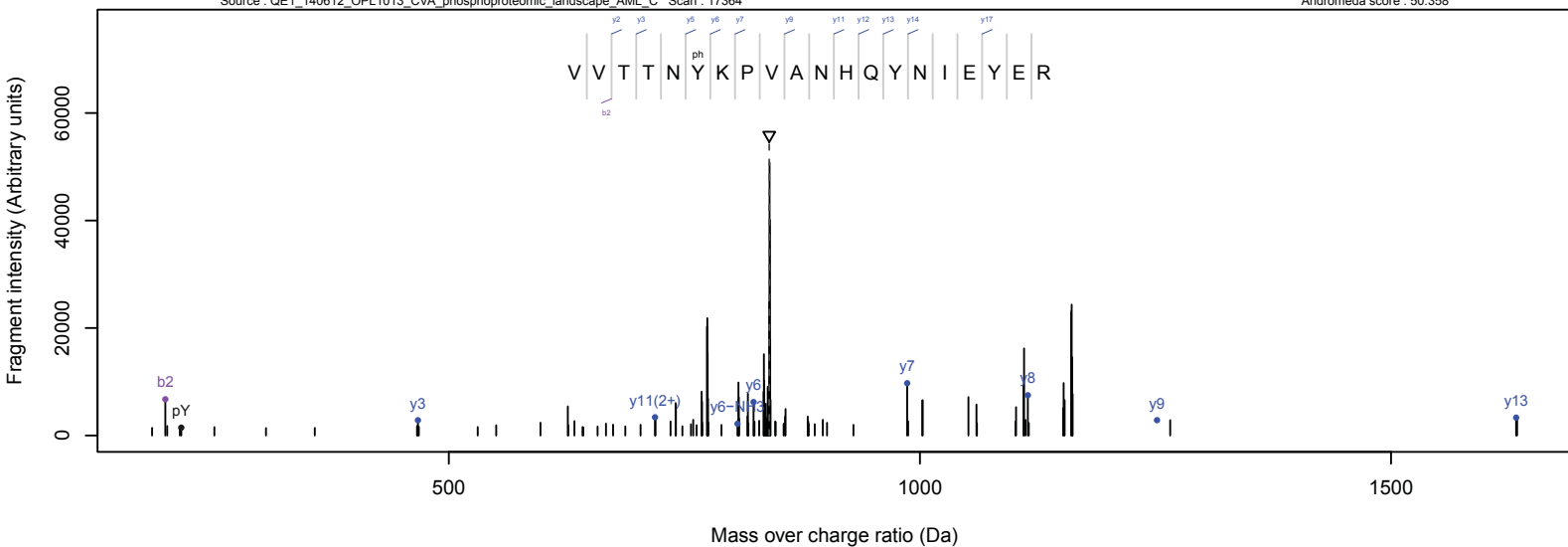
Andromeda score : 50.358



**VVTNNY(ph)KPVANHQYNI EYER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17364

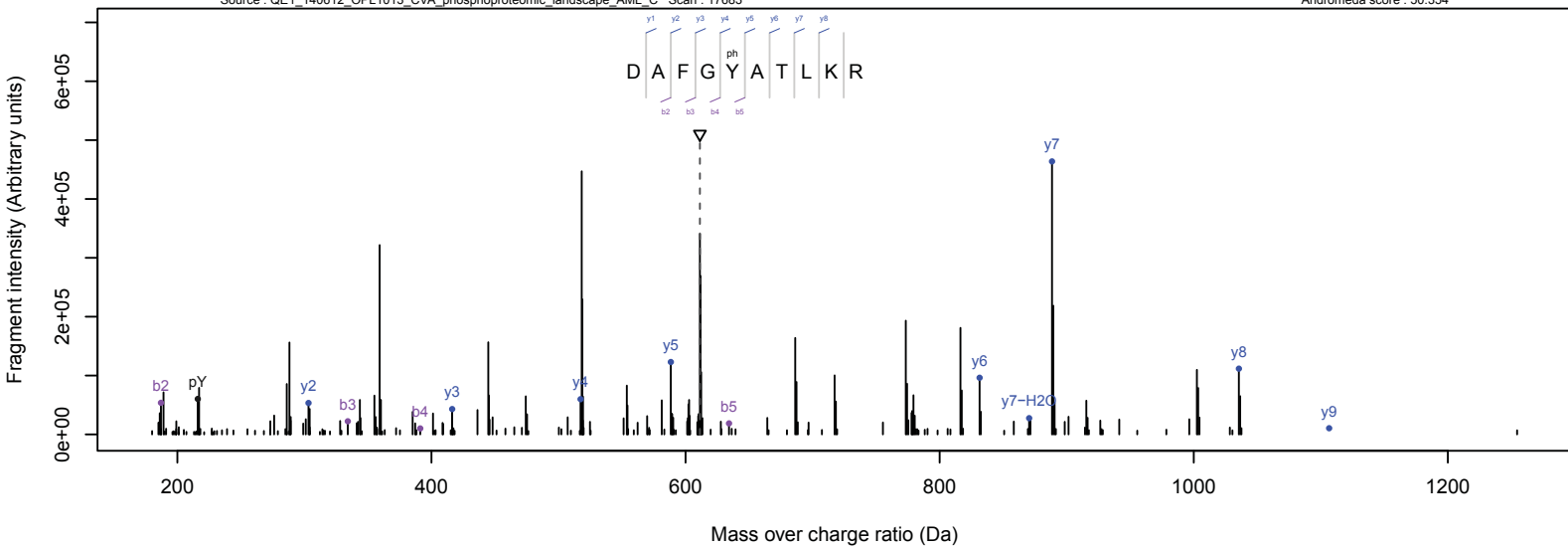
Andromeda score : 50.358

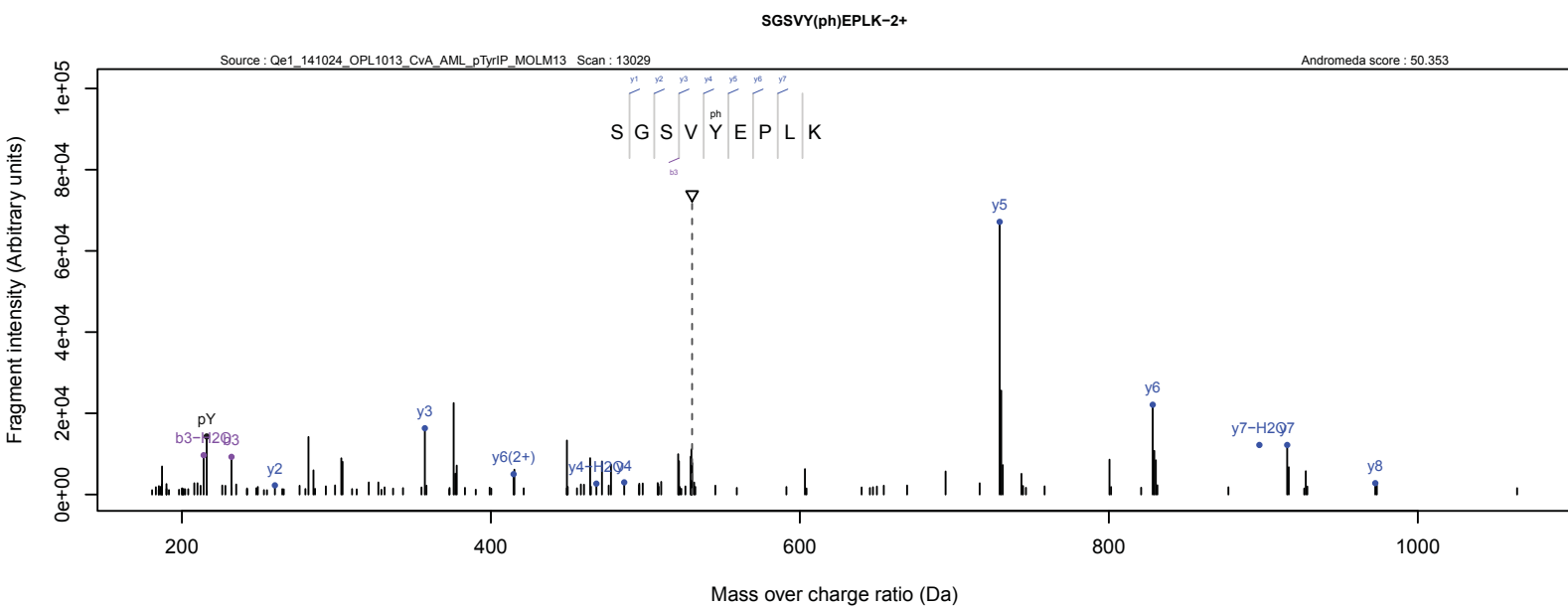
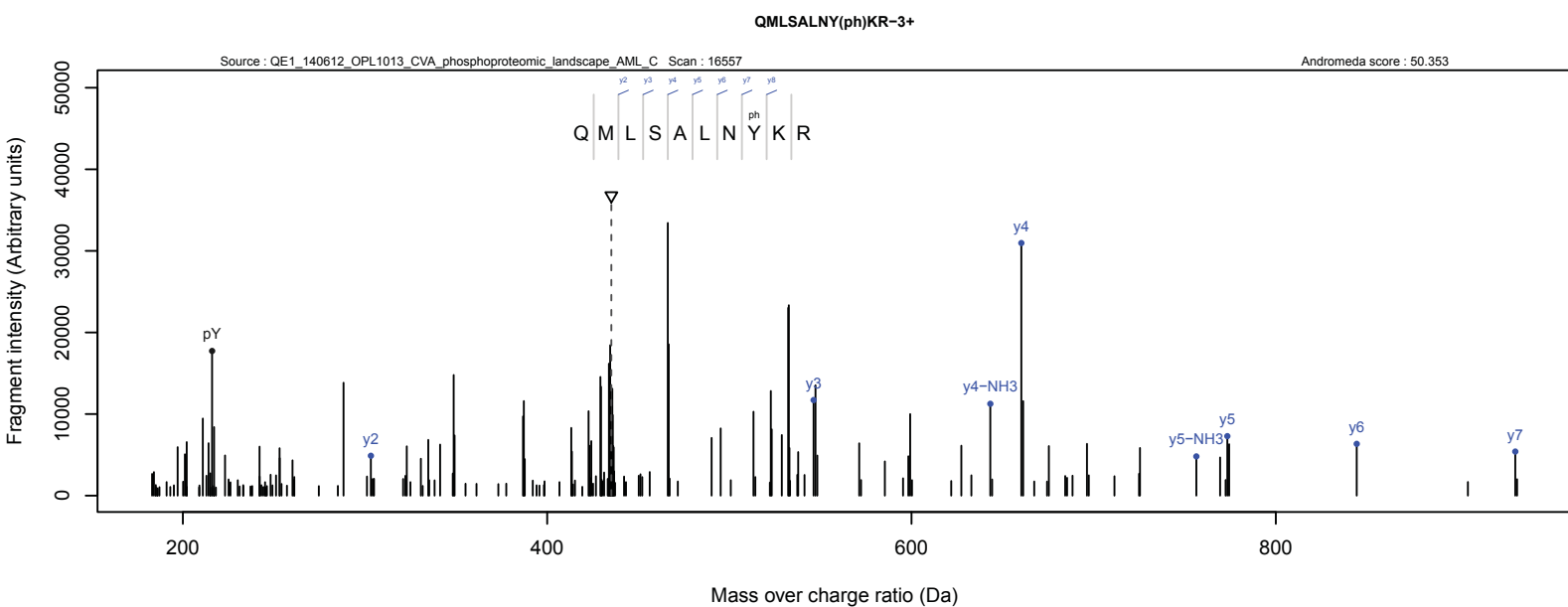
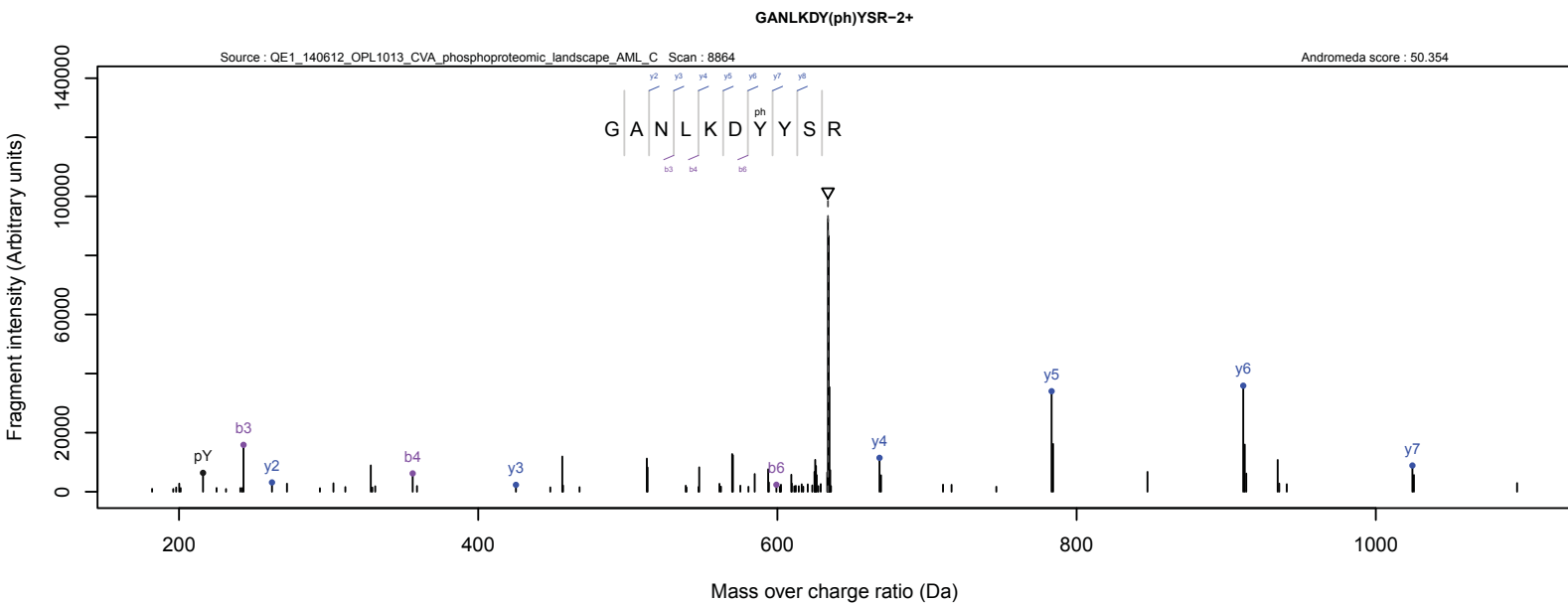


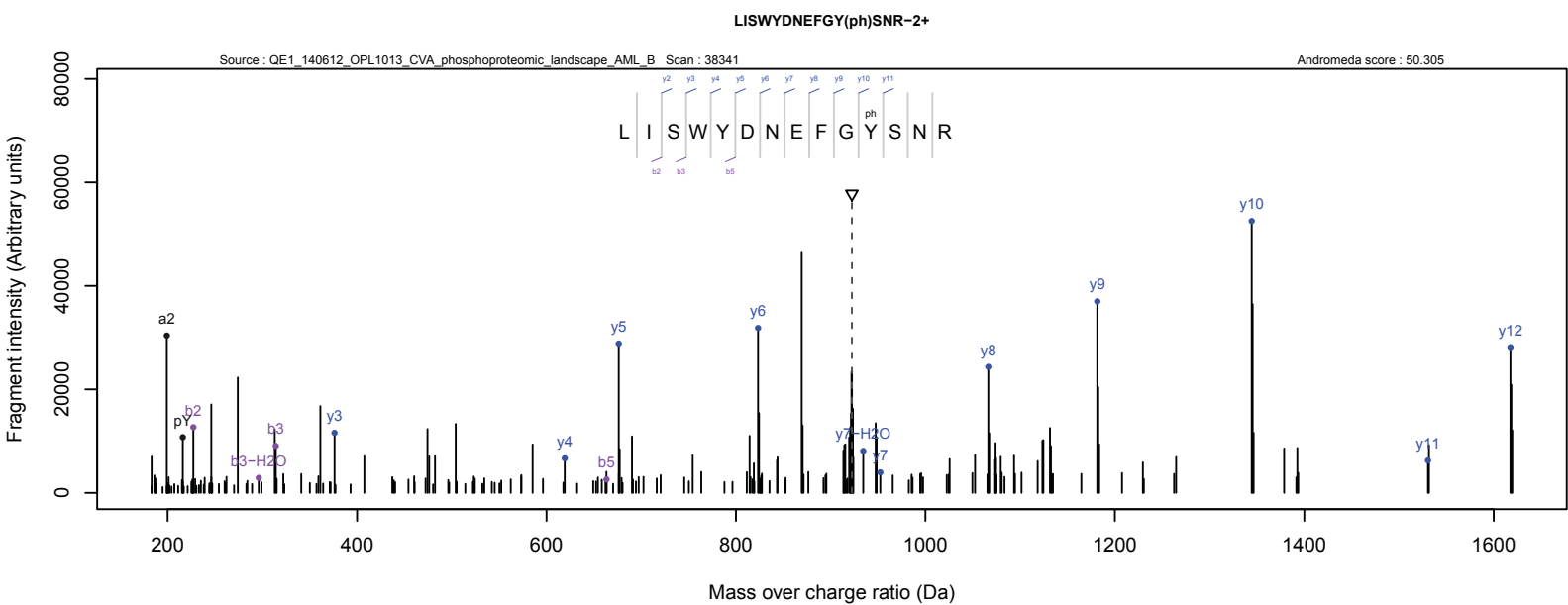
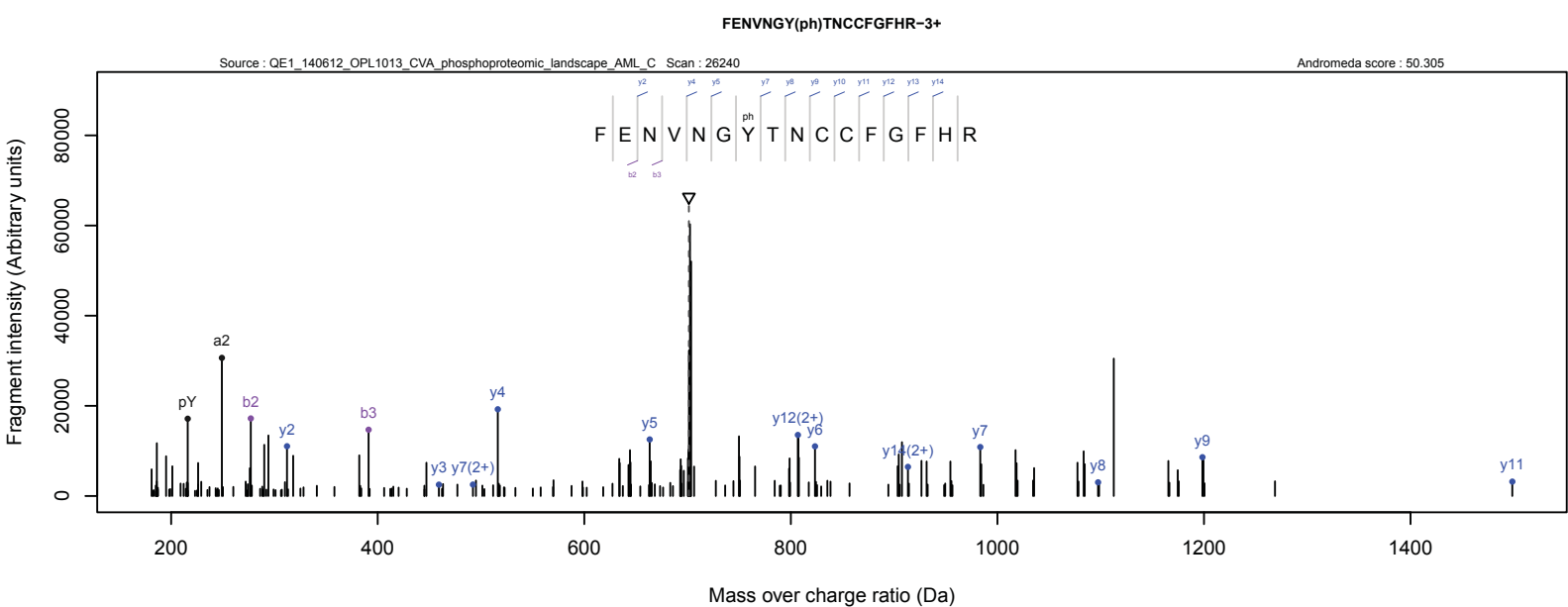
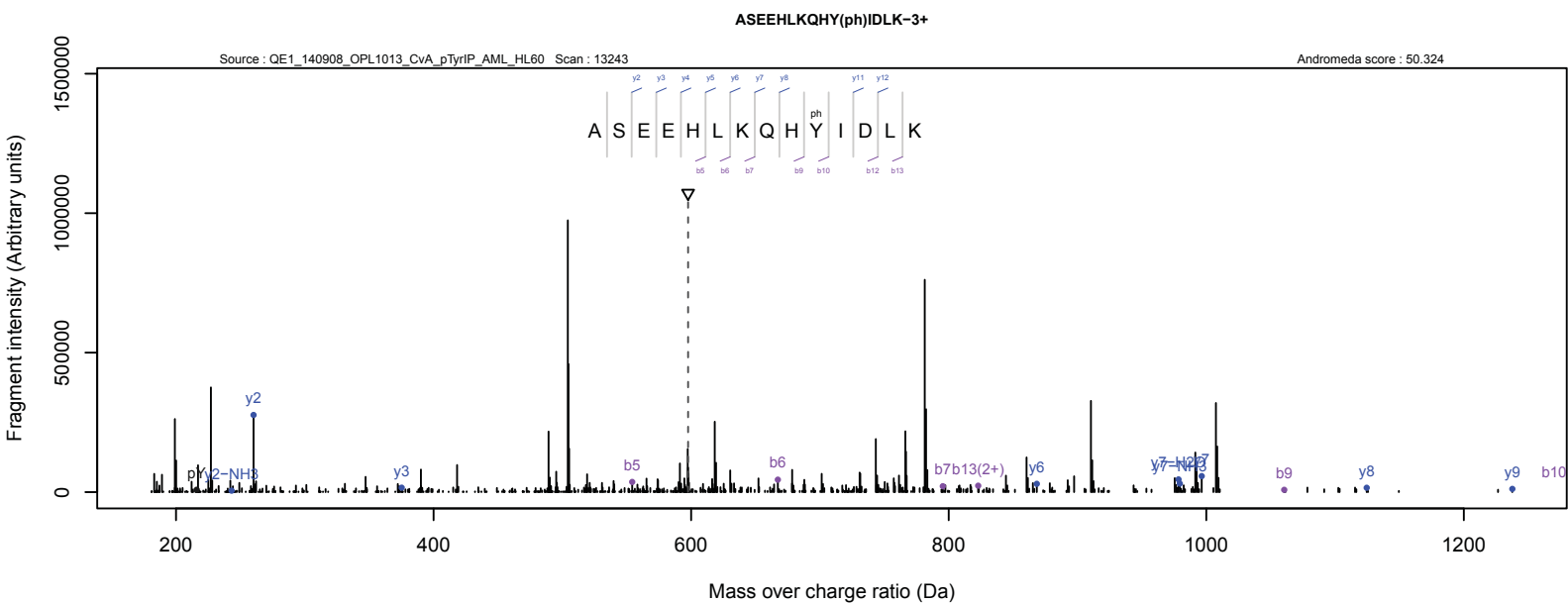
**DAFGY(ph)ATLKR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17683

Andromeda score : 50.354



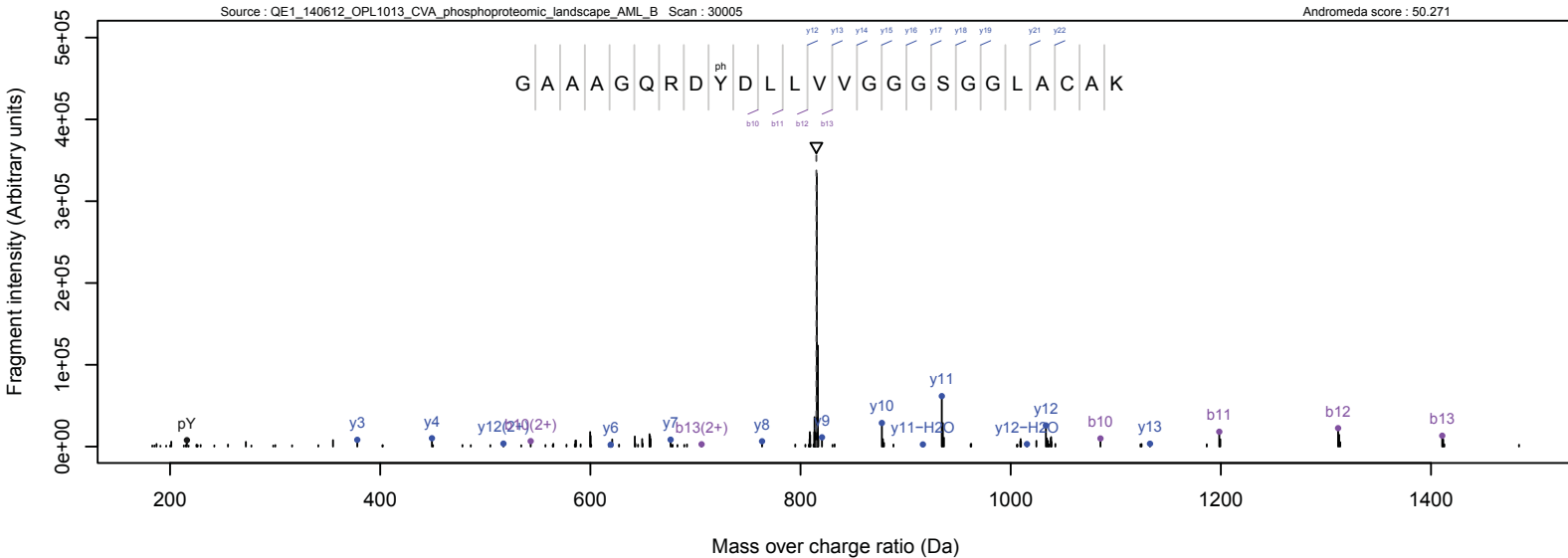




GAAAGQRDY(ph)DLLVVGGSGLCAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30005

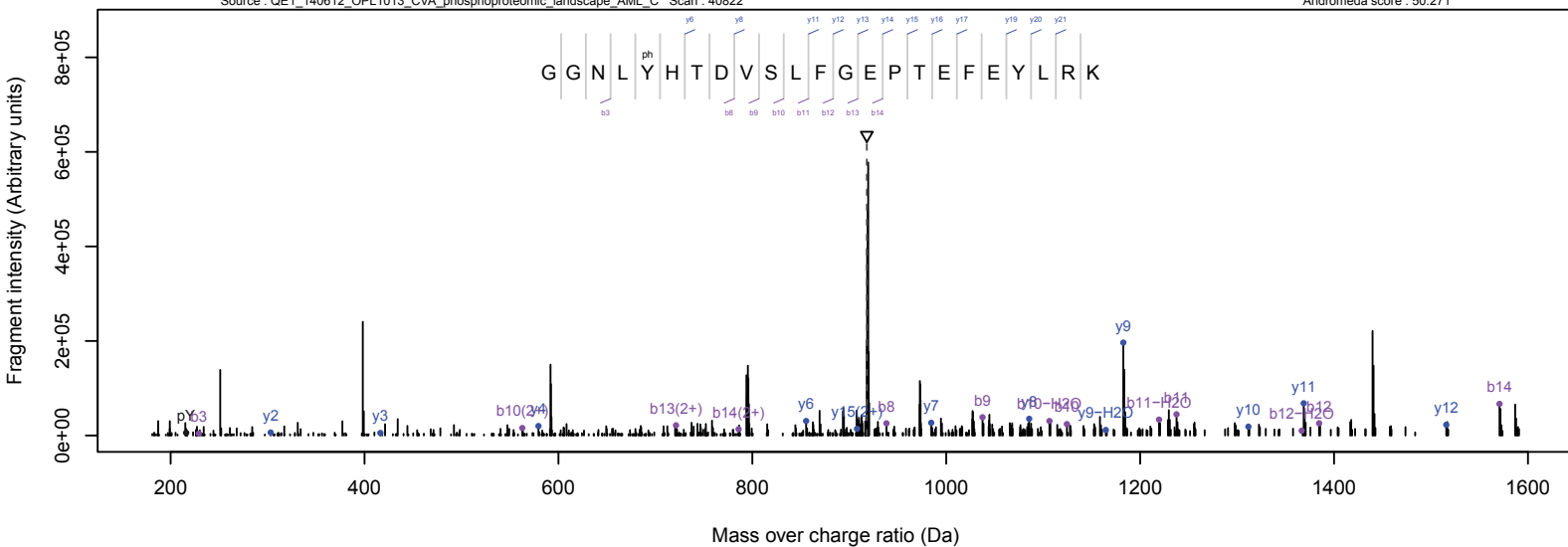
Andromeda score : 50.271



GGNLY(ph)HTDVSFLGEPTEFEYLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40822

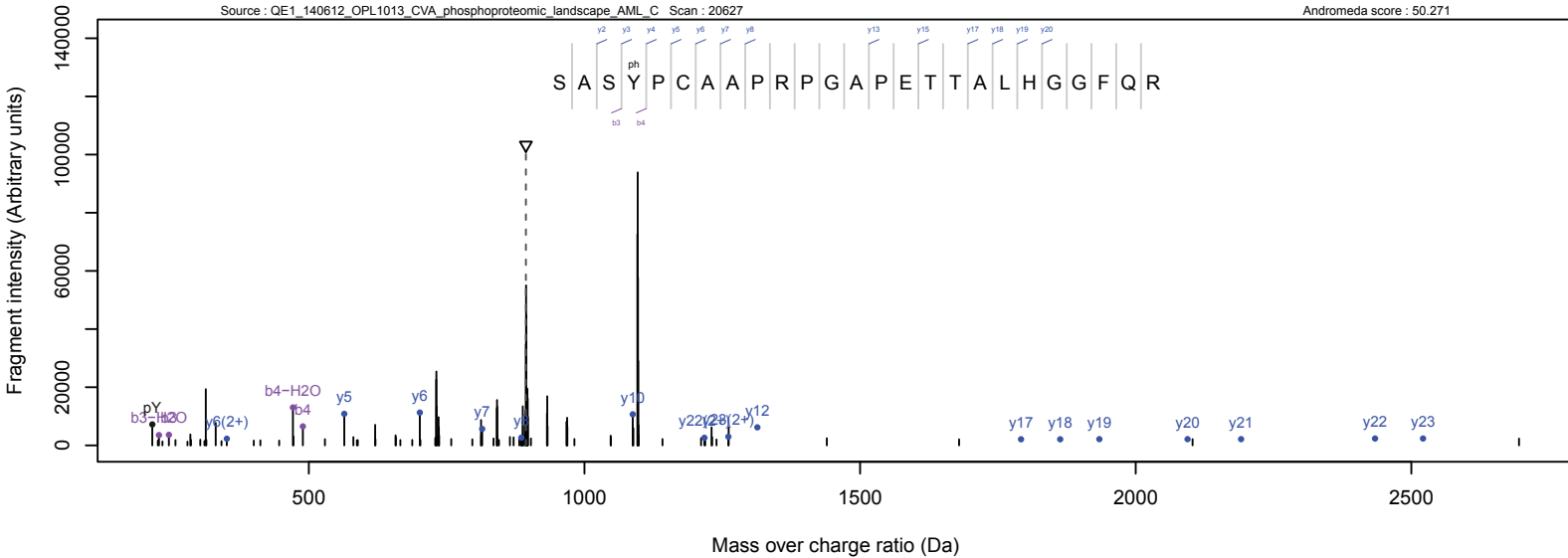
Andromeda score : 50.271



SASY(ph)PCAAPRPGAPETLHGGFQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20627

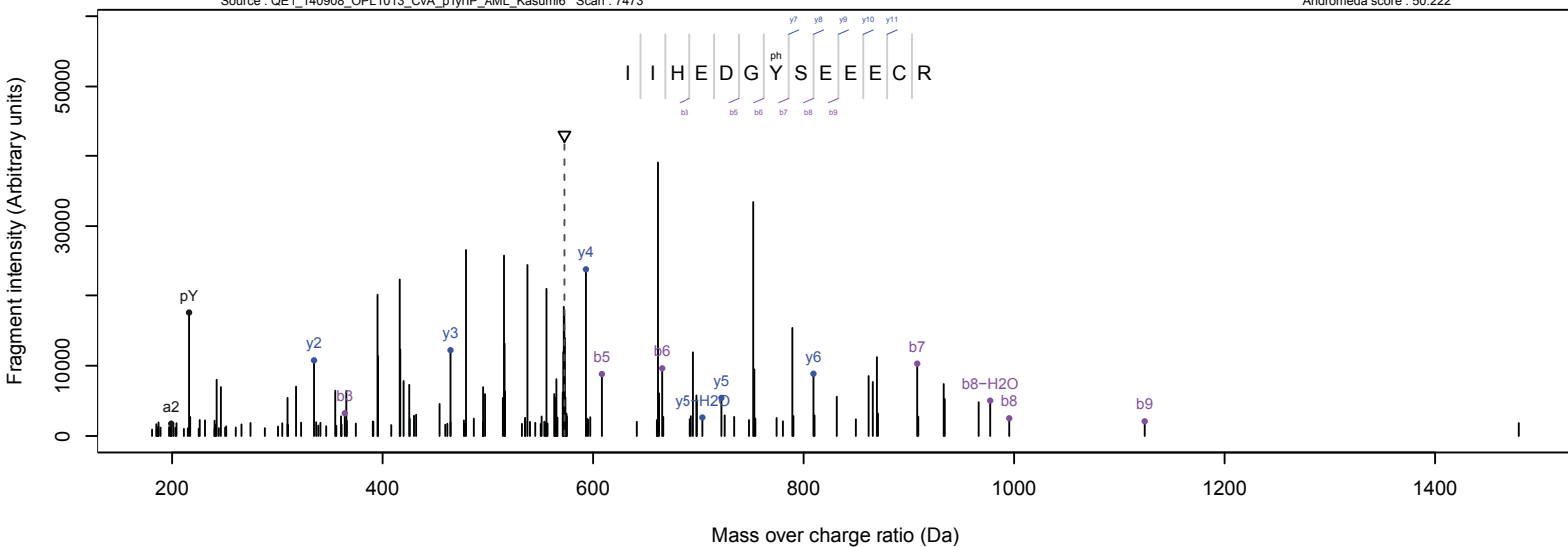
Andromeda score : 50.271



IIEHDGY(ph)SEEECR-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 7473

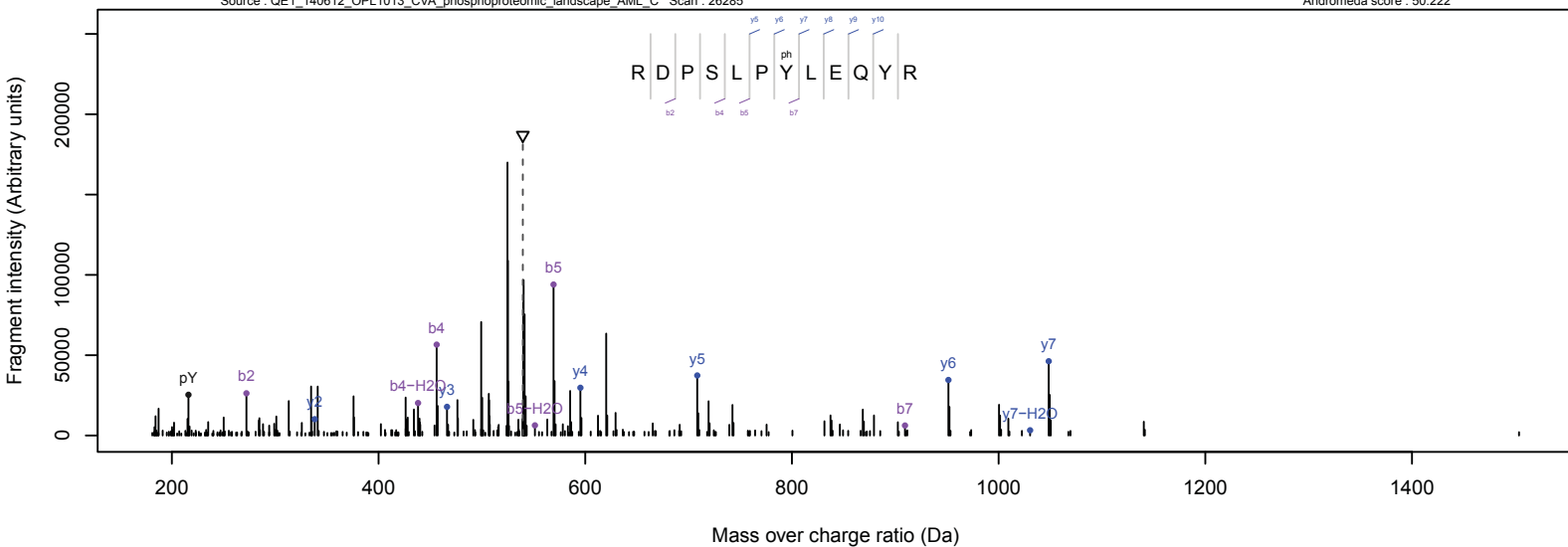
Andromeda score : 50.222



RDPSLPY(ph)LEQYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26285

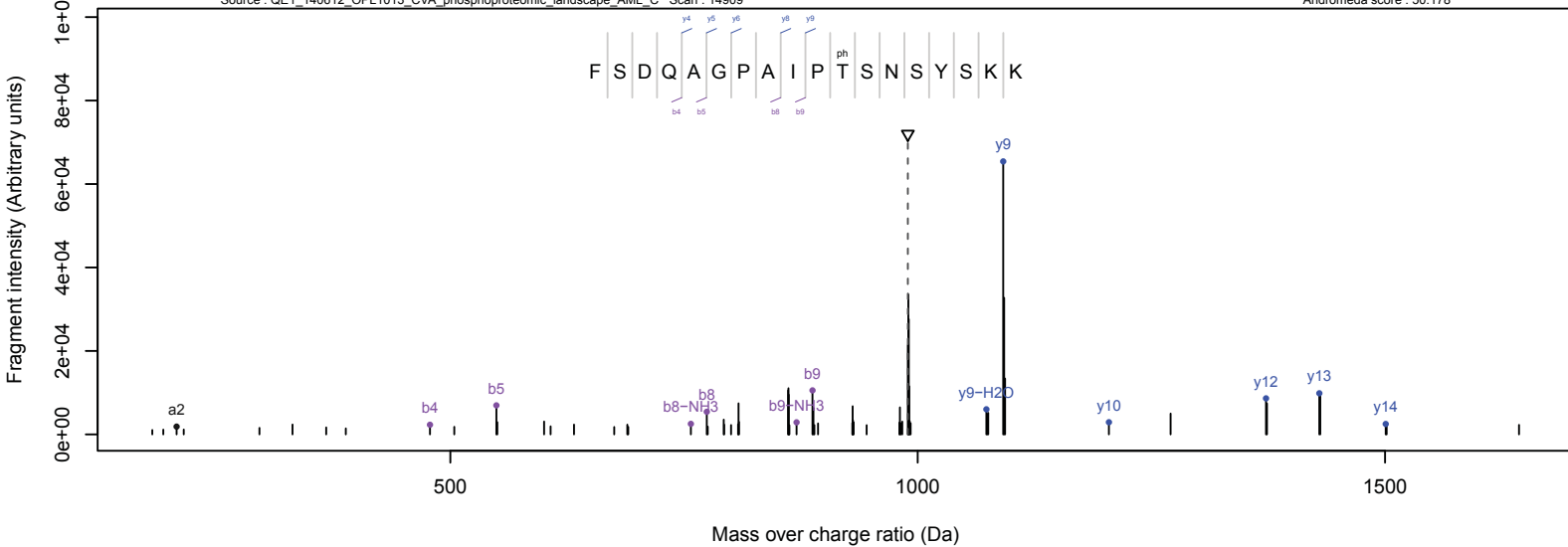
Andromeda score : 50.222



FSDQAGPAIP(ph)SNSYSKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14909

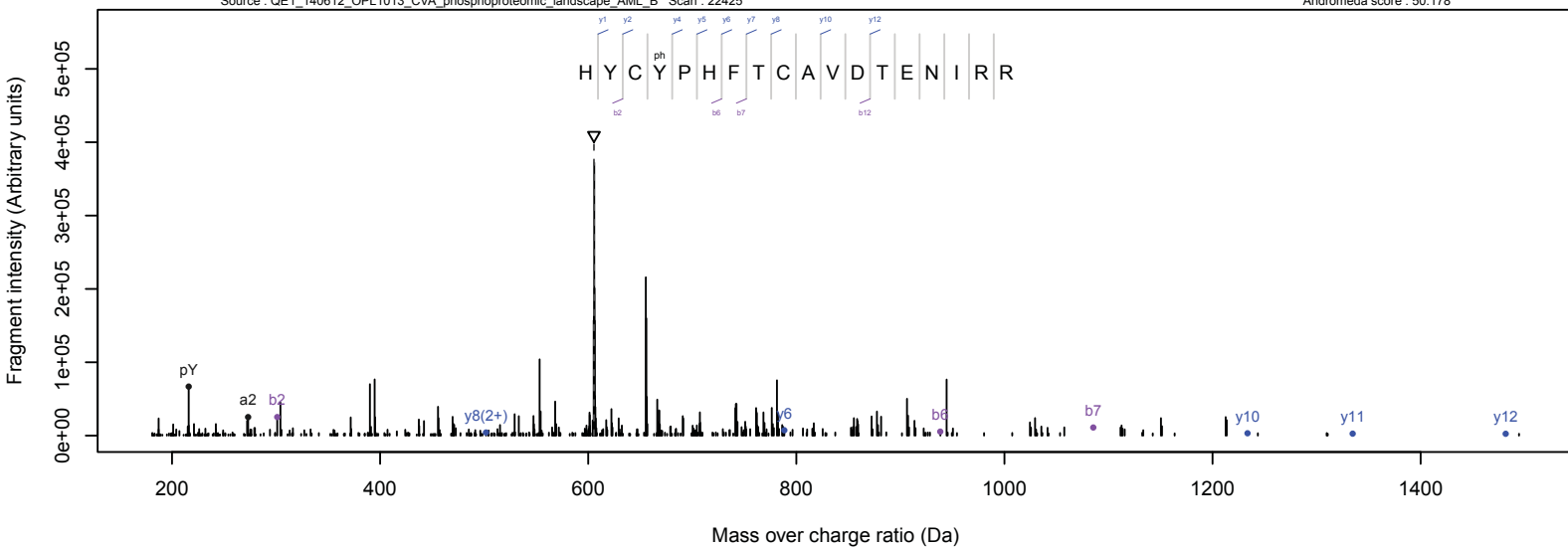
Andromeda score : 50.178



## HYCY(ph)PHFTCAVDTENIRR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22425

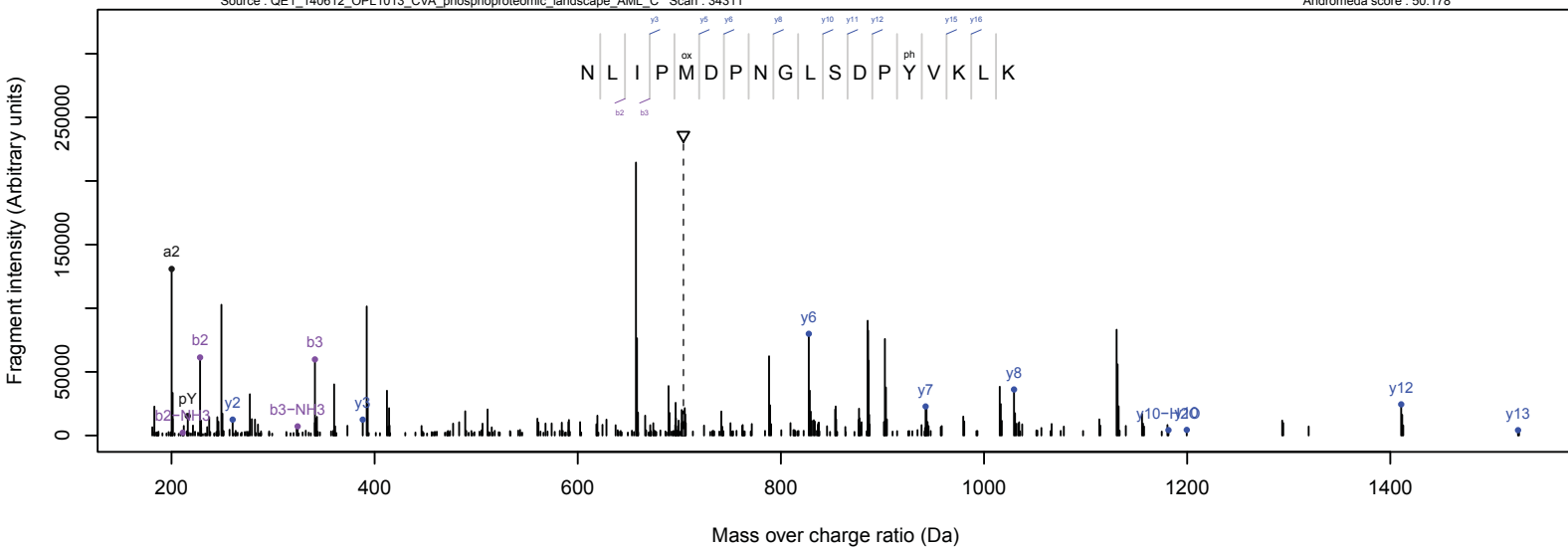
Andromeda score : 50.178



## NLIPM(ox)DPNGLSDPY(ph)VKLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34311

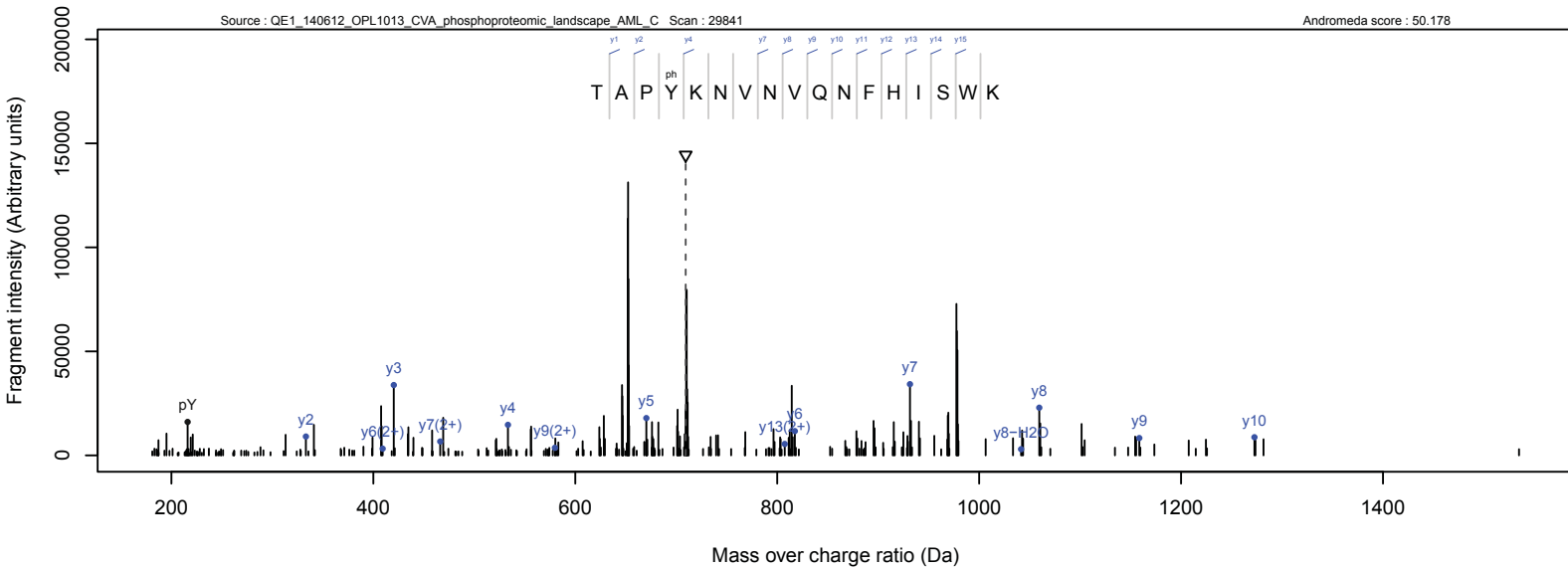
Andromeda score : 50.178

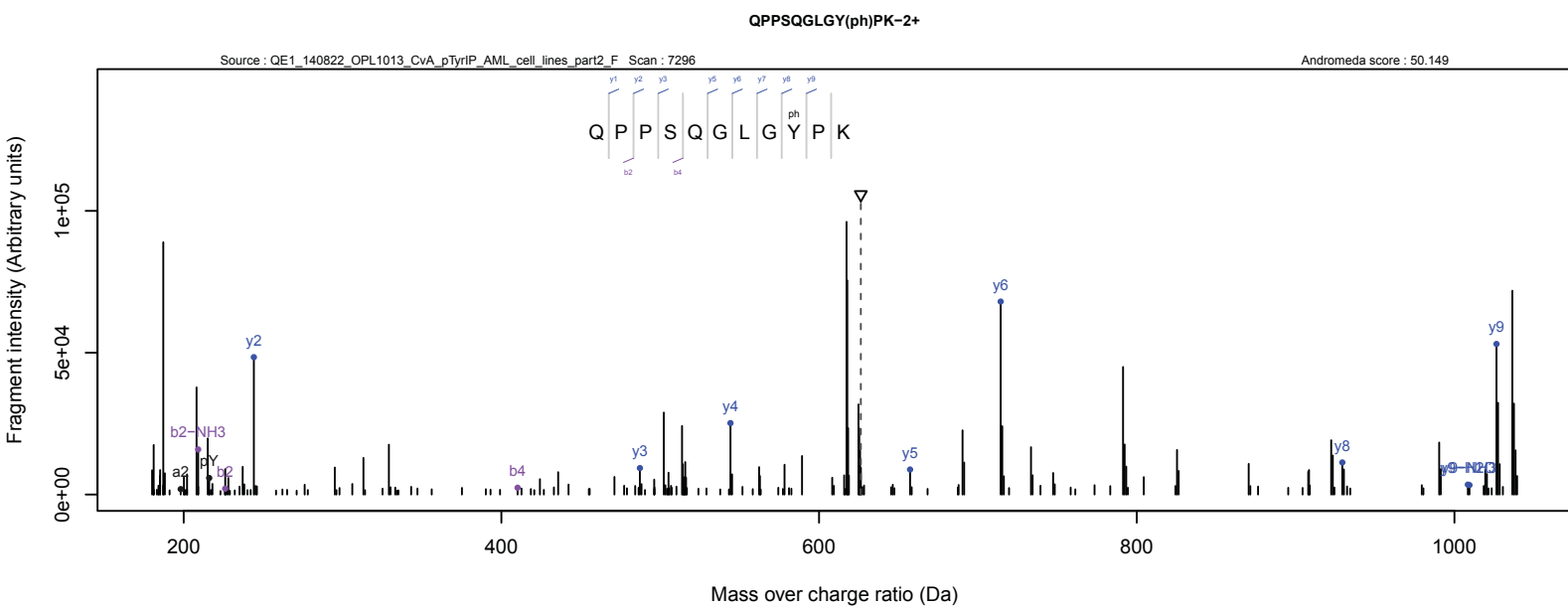
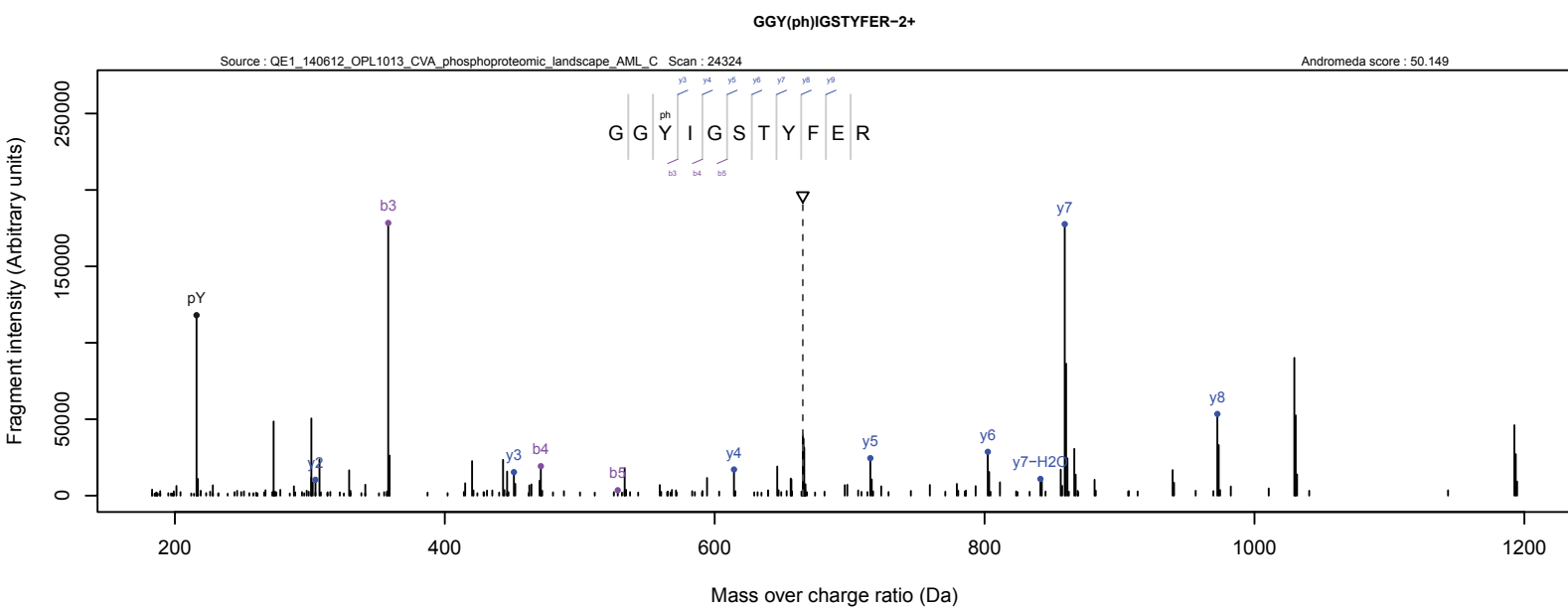
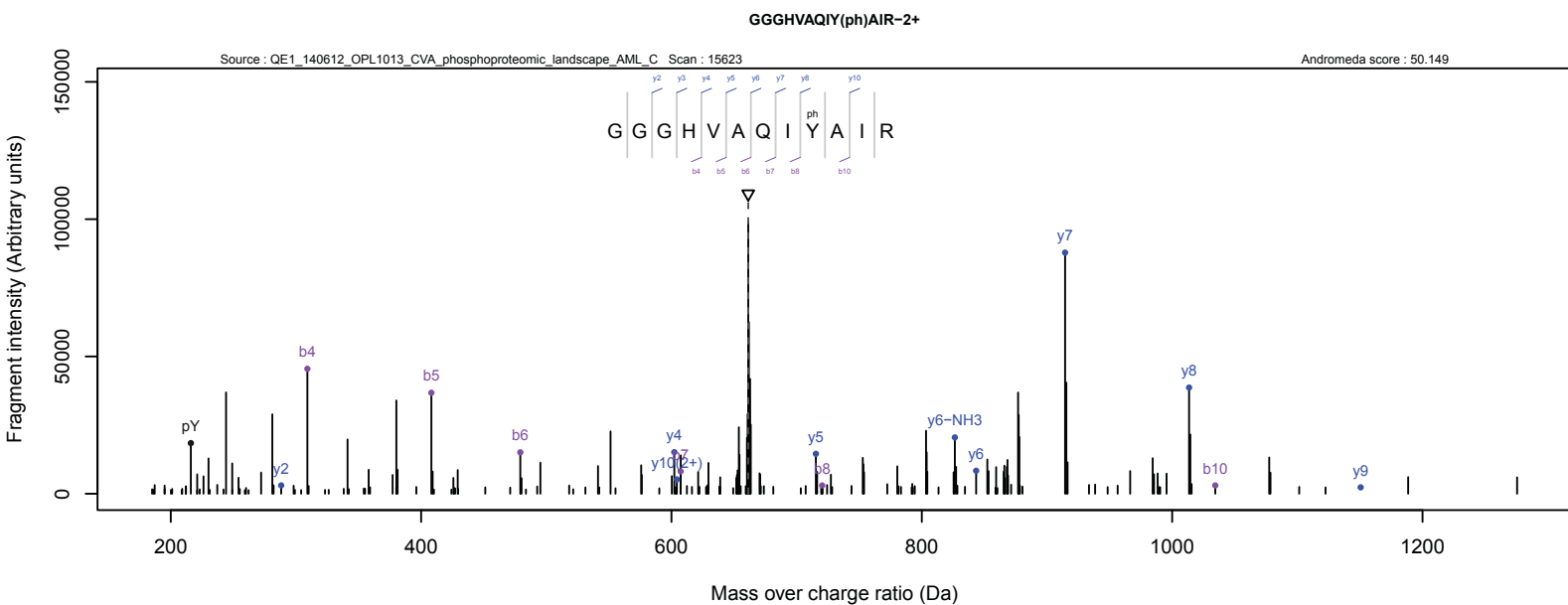


## TAPY(ph)KNVNVQNFHISWK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29841

Andromeda score : 50.178



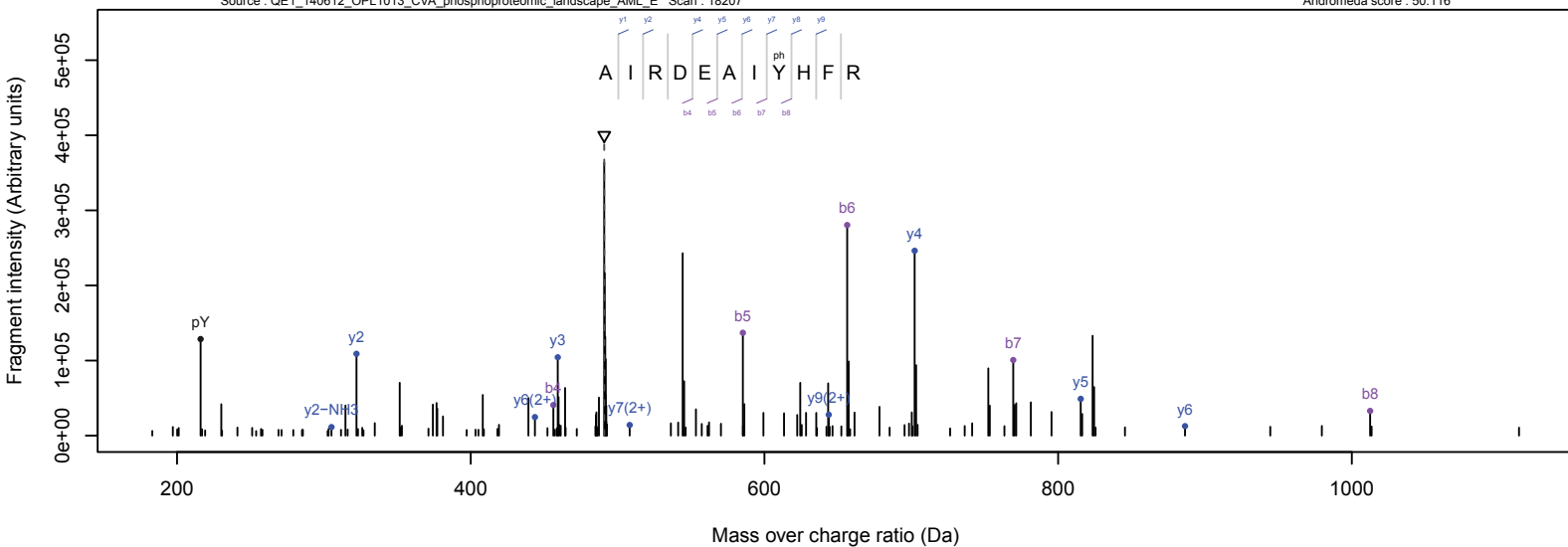




## AIRDEAIY(ph)HFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 18207

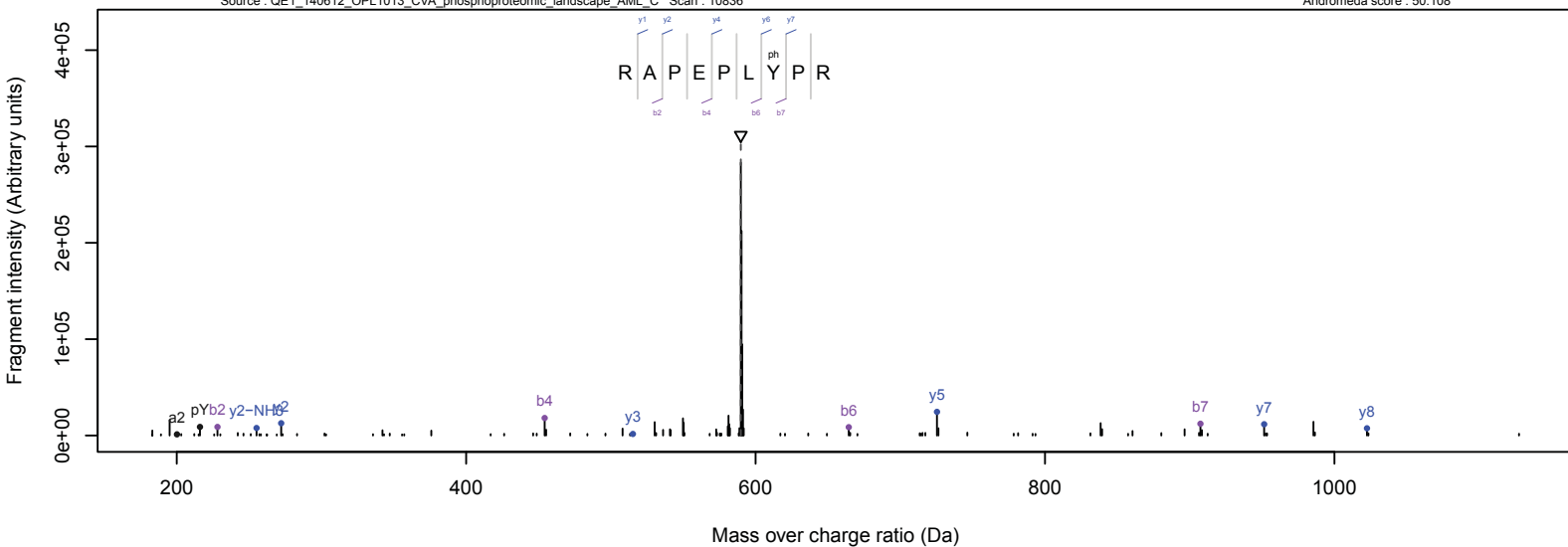
Andromeda score : 50.116



## RAPEPLY(ph)PR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10836

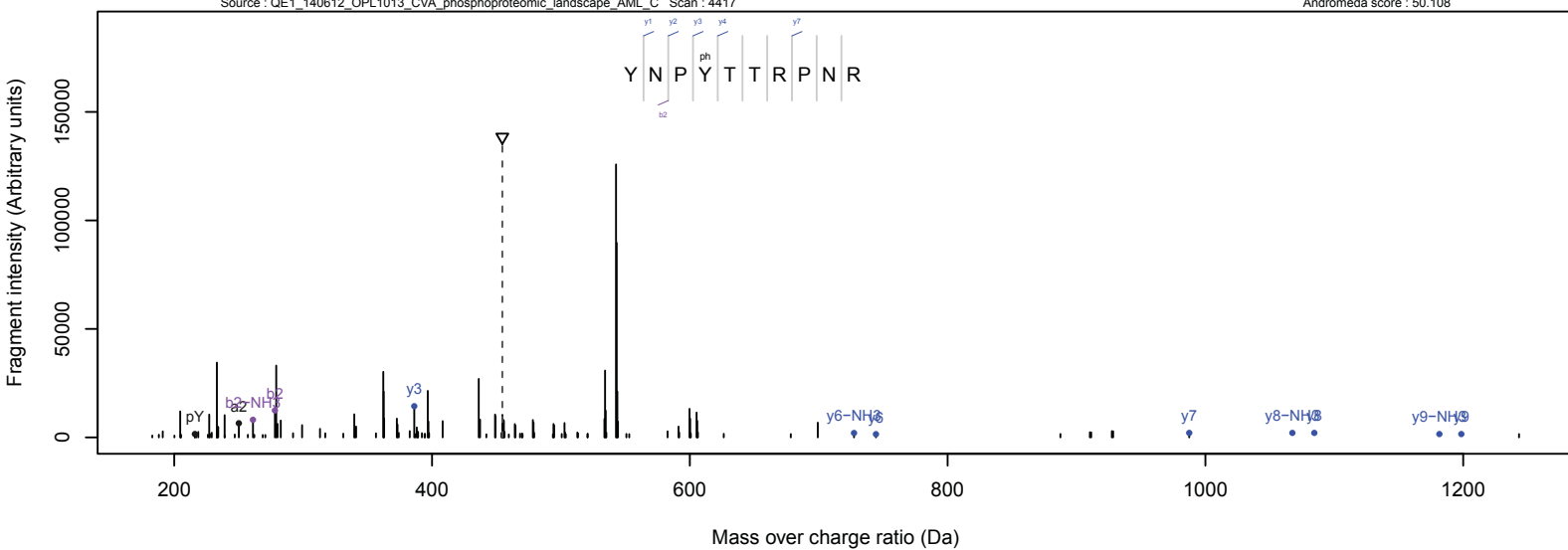
Andromeda score : 50.108



## YNPY(ph)TTRPNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 4417

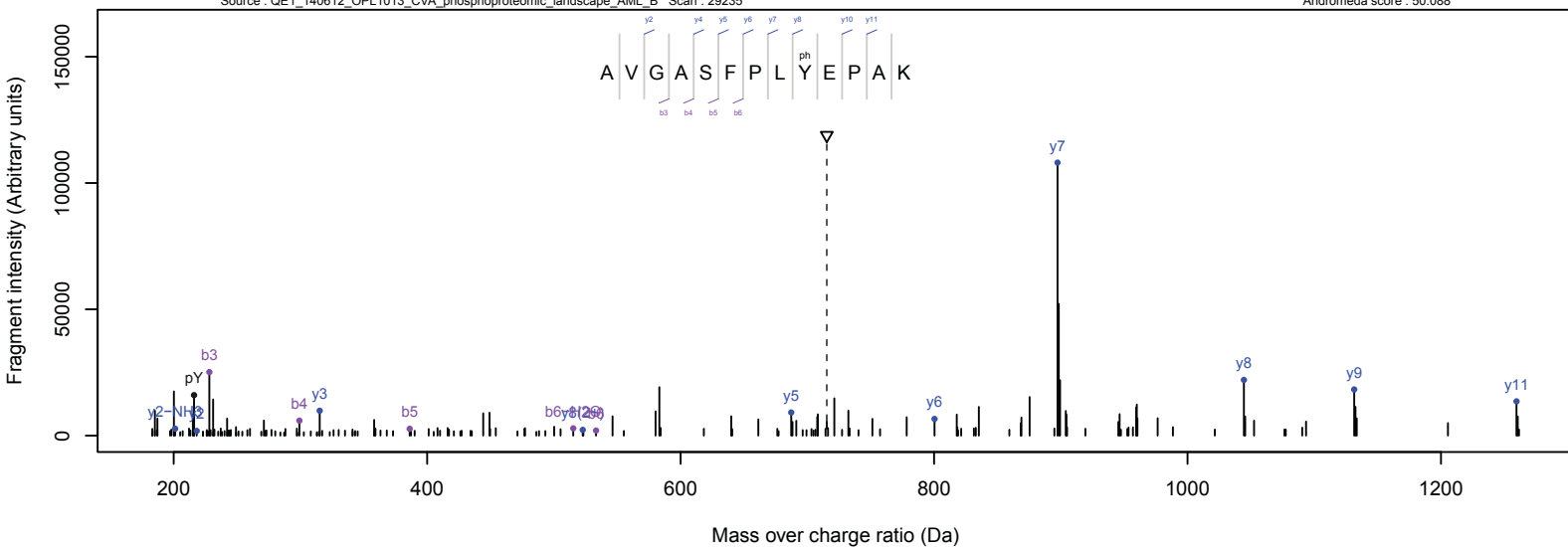
Andromeda score : 50.108



AVGASFLY(ph)EPAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29235

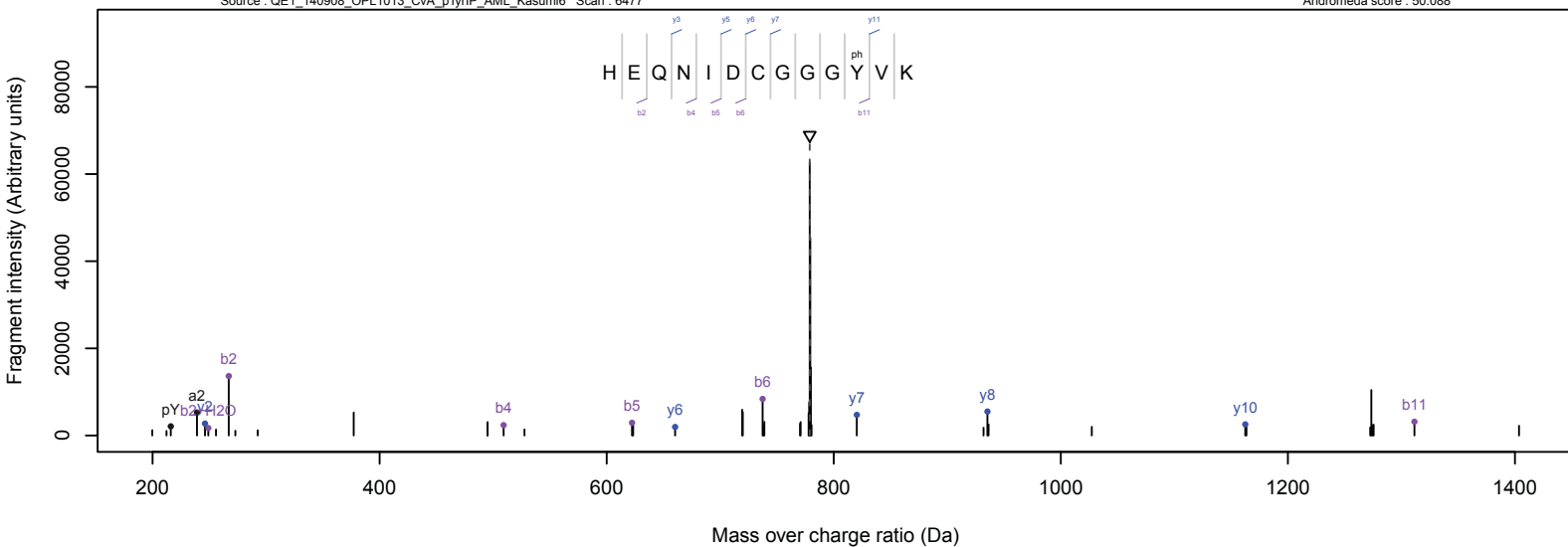
Andromeda score : 50.088



HEQNIDCGGY(ph)VK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 6477

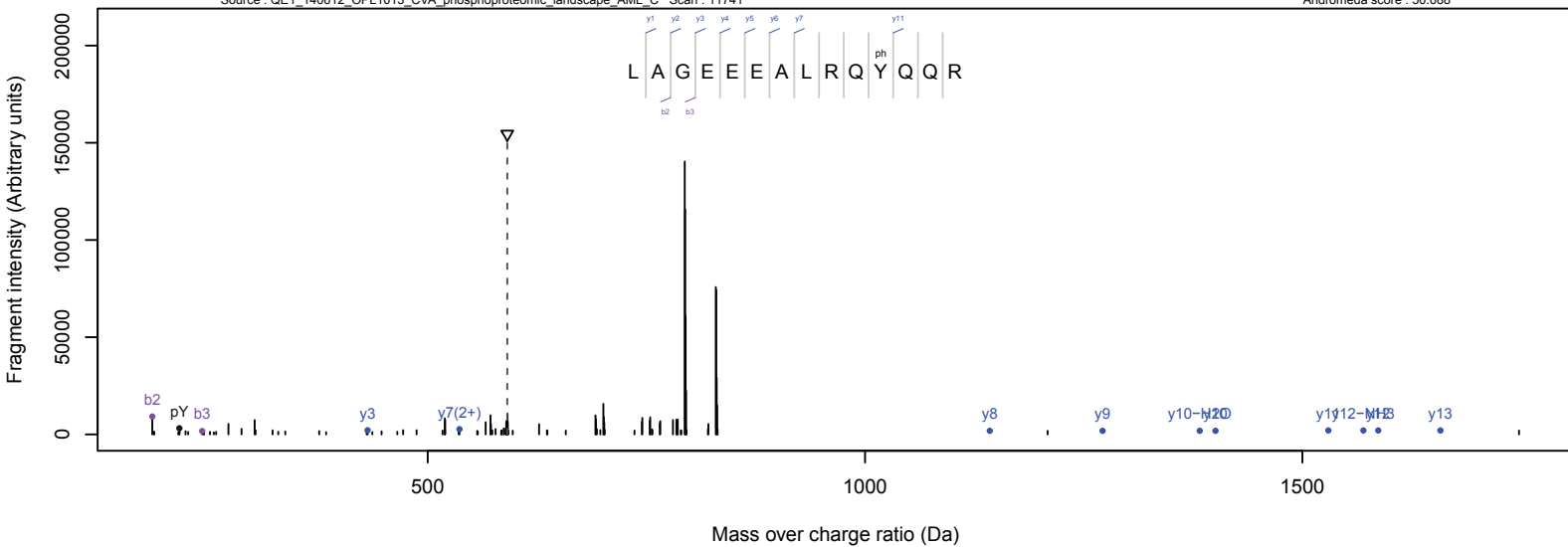
Andromeda score : 50.088



LAGEEEEALRQY(ph)QQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11741

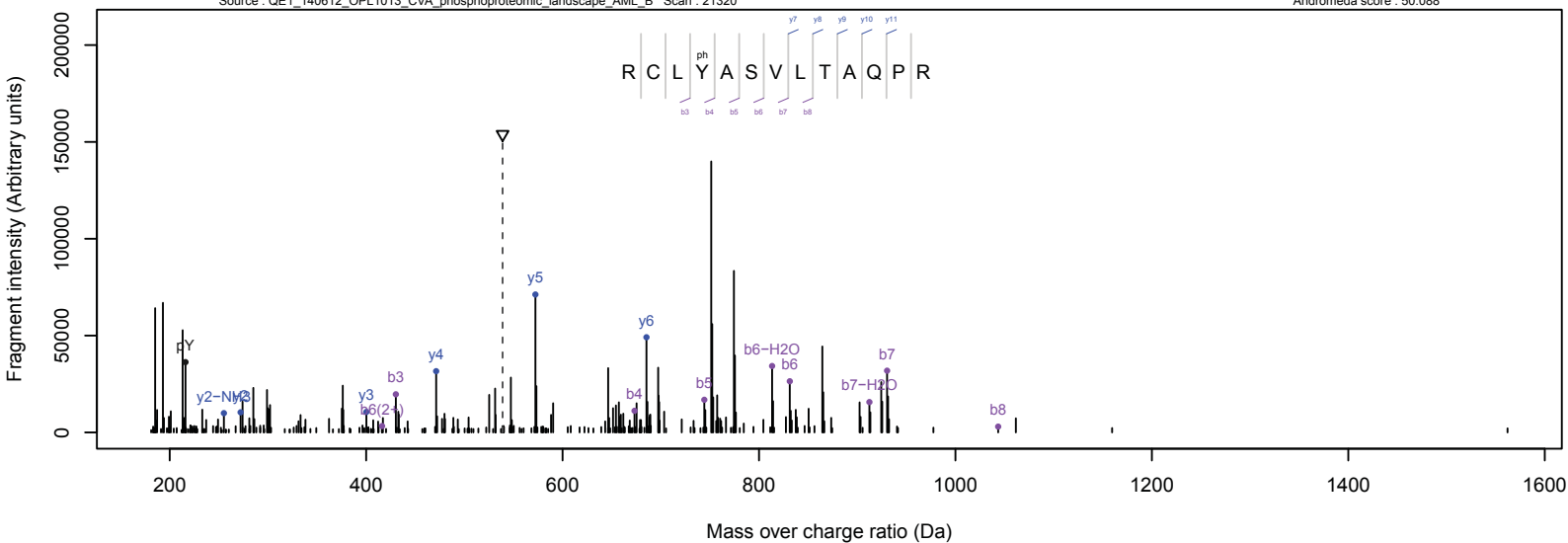
Andromeda score : 50.088



RCLY(ph)ASVLTQPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21320

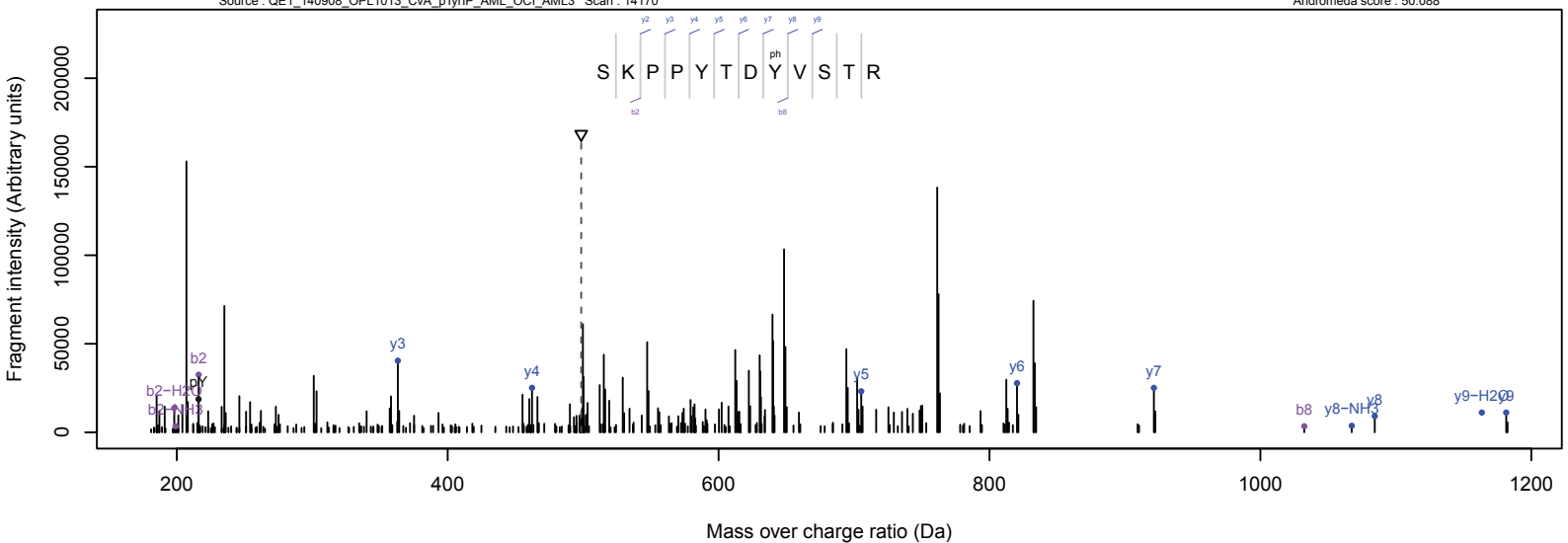
Andromeda score : 50.088



SKPPYTDY(ph)VSTR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 14170

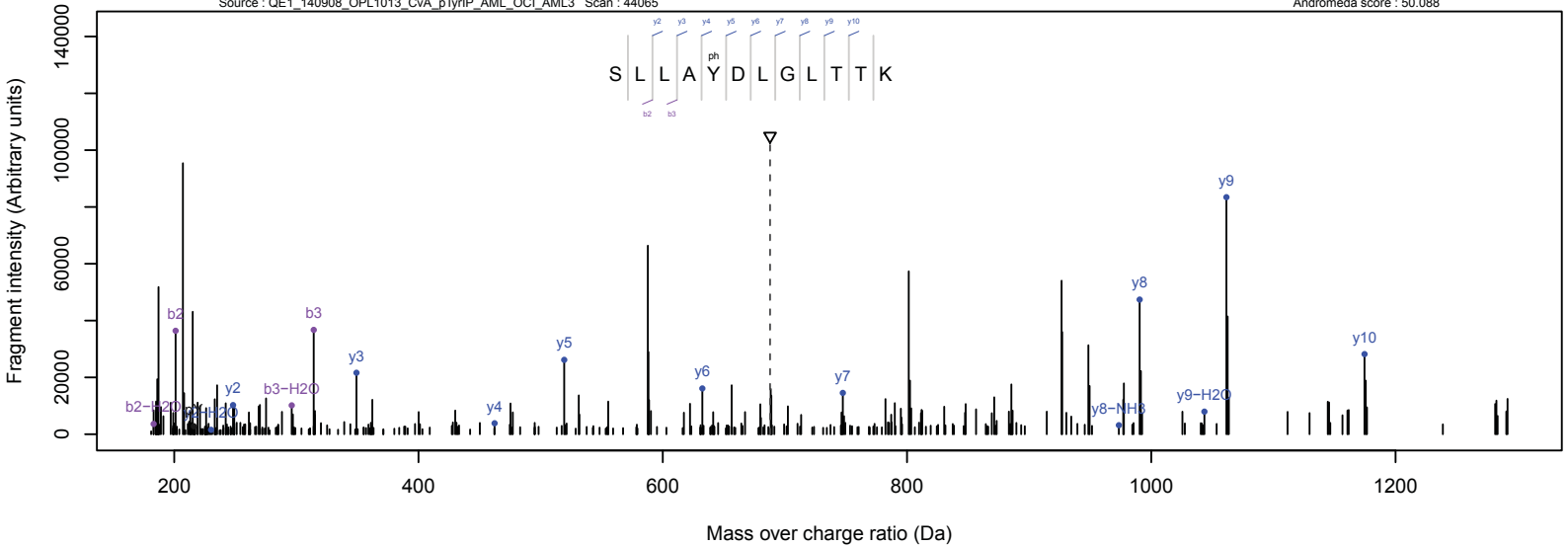
Andromeda score : 50.088



SLLAY(ph)DLGLTTK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 44065

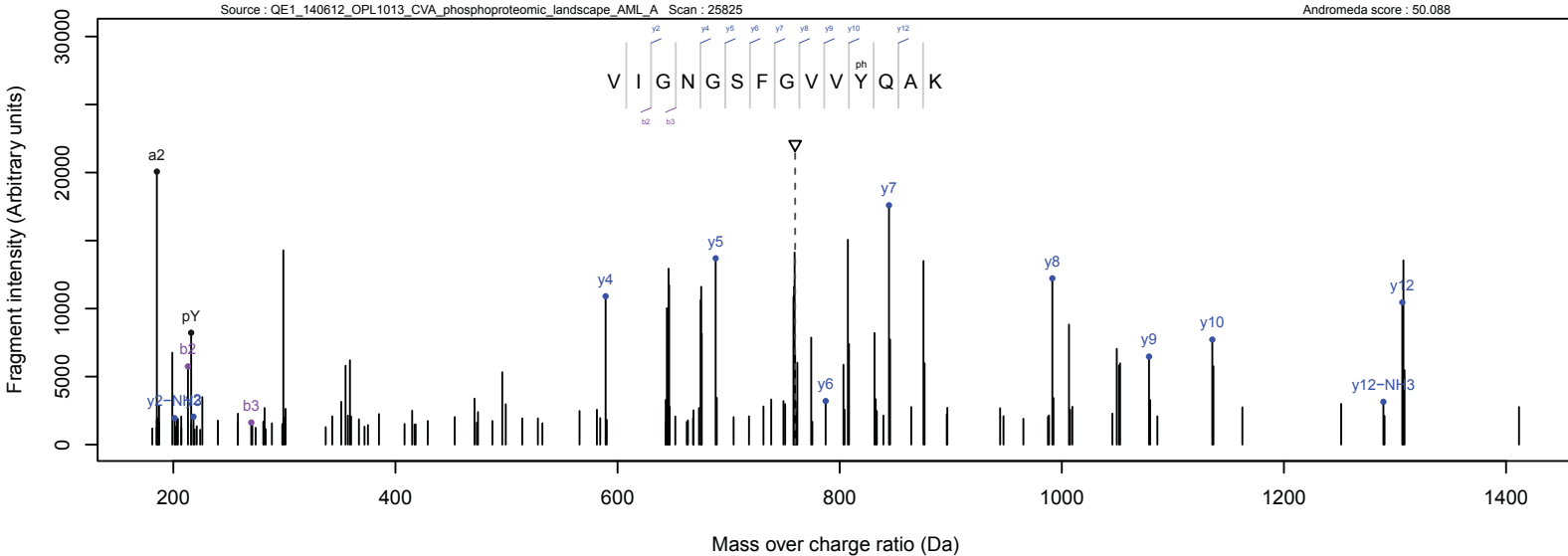
Andromeda score : 50.088



VIGNSGVVY(ph)QAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 25825

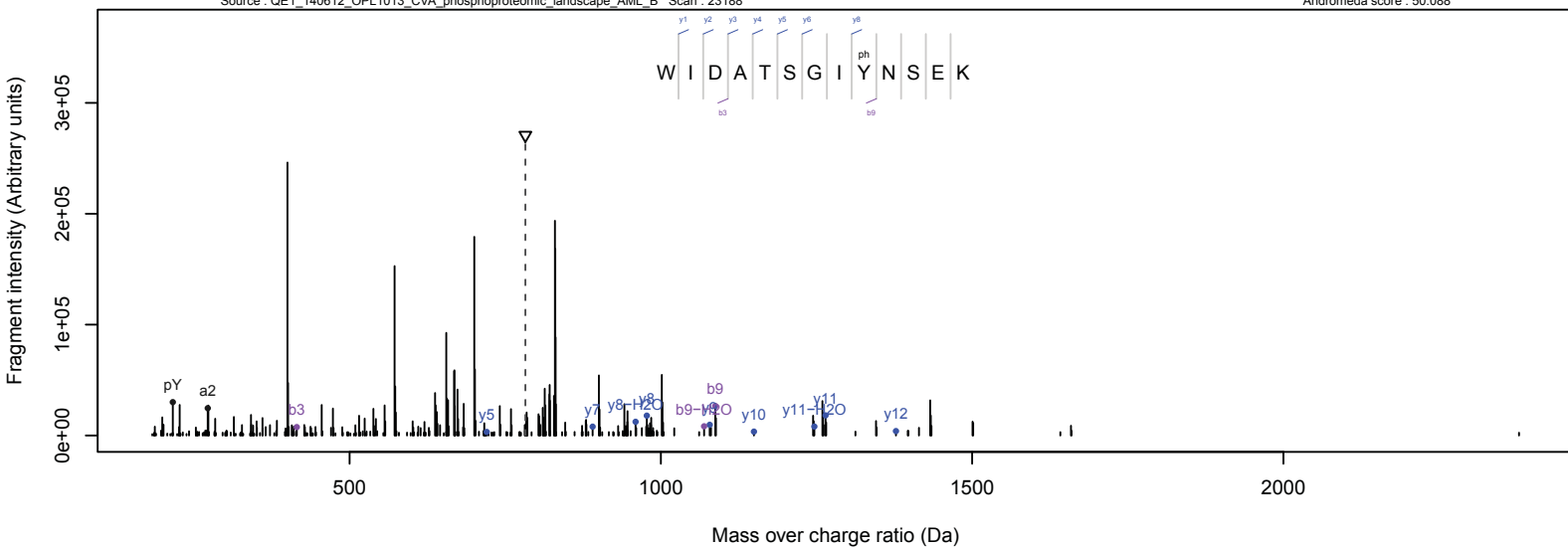
Andromeda score : 50.088



WIDATSGIY(ph)NSEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23188

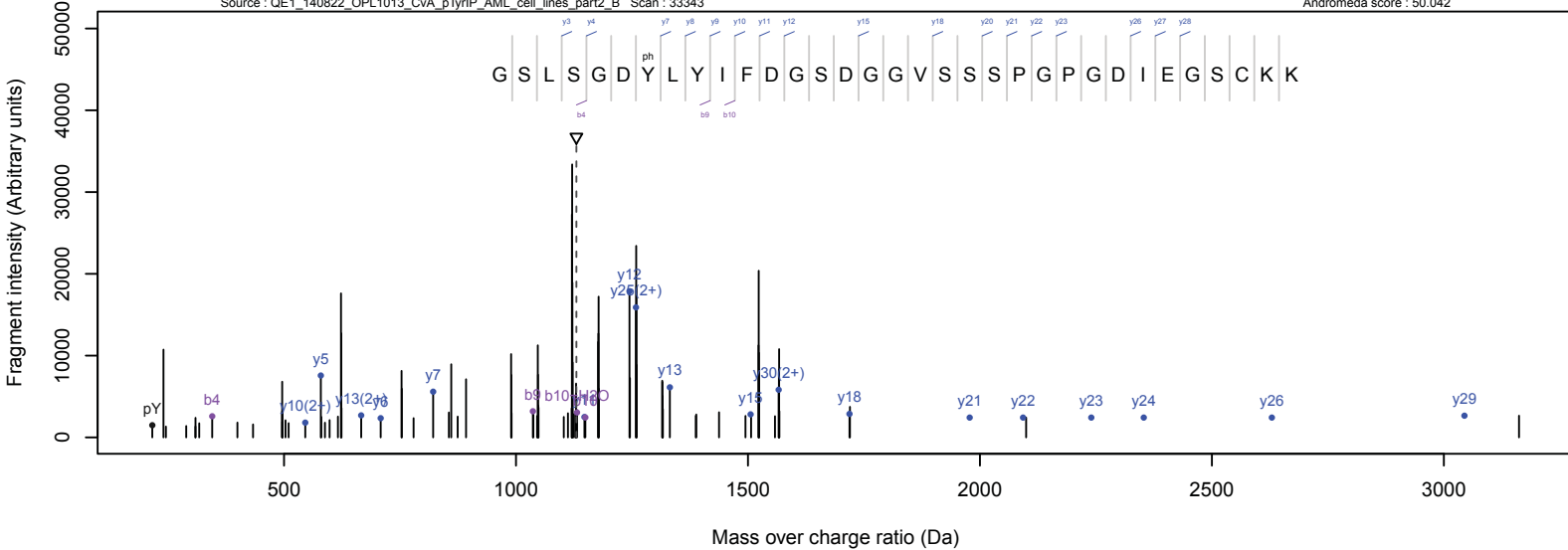
Andromeda score : 50.088



GSLSGDY(ph)LYIFDGS DGGVSSSPGPGDIEGSKK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 33343

Andromeda score : 50.042

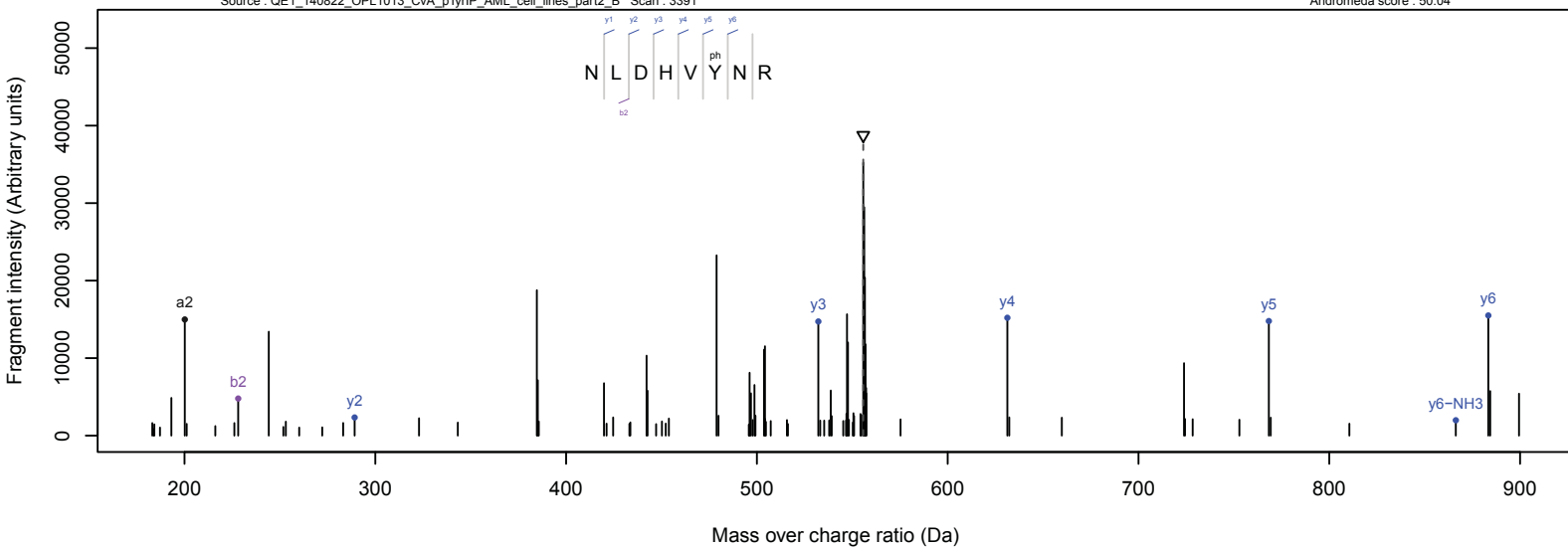




## NLDHVY(ph)NR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 3391

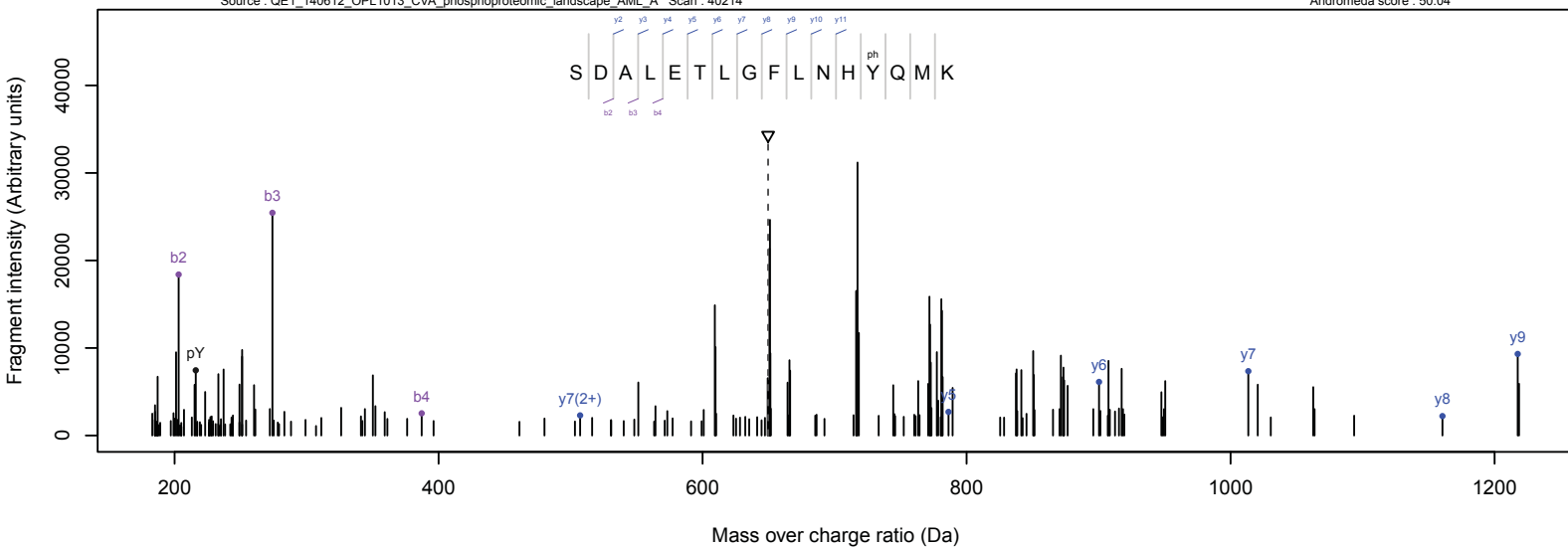
Andromeda score : 50.04



## SDALETGLFNHY(ph)QMK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 40214

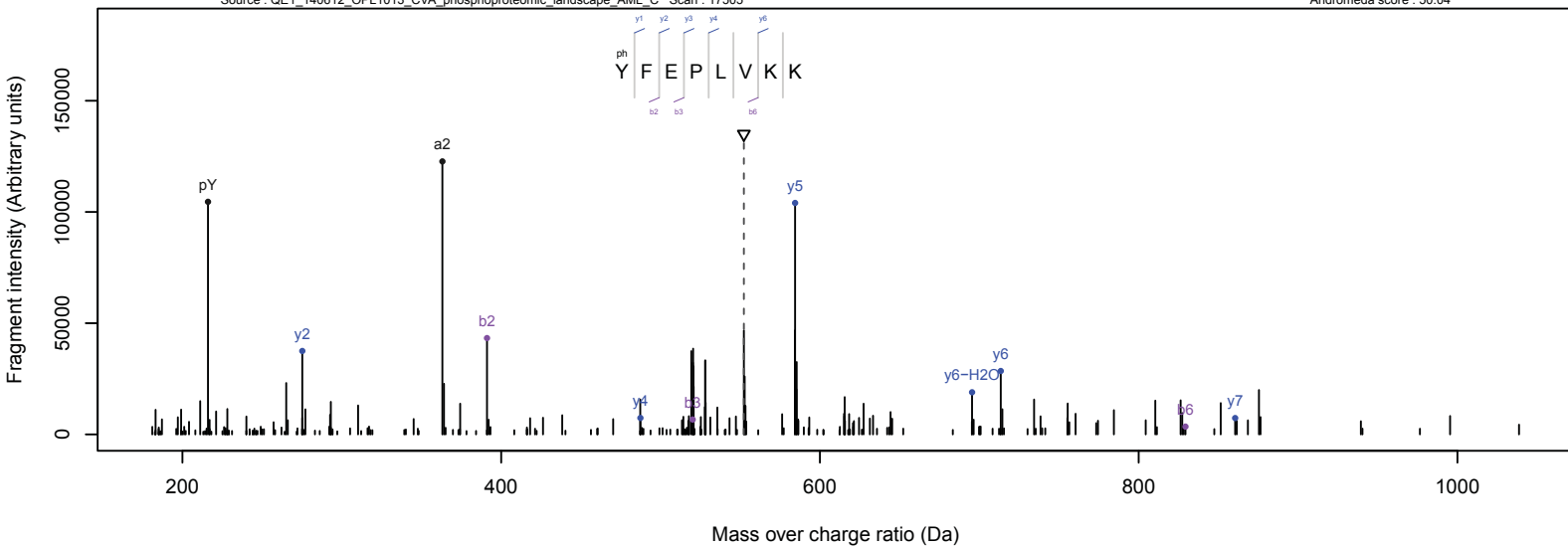
Andromeda score : 50.04



## Y(ph)FEPLVKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17505

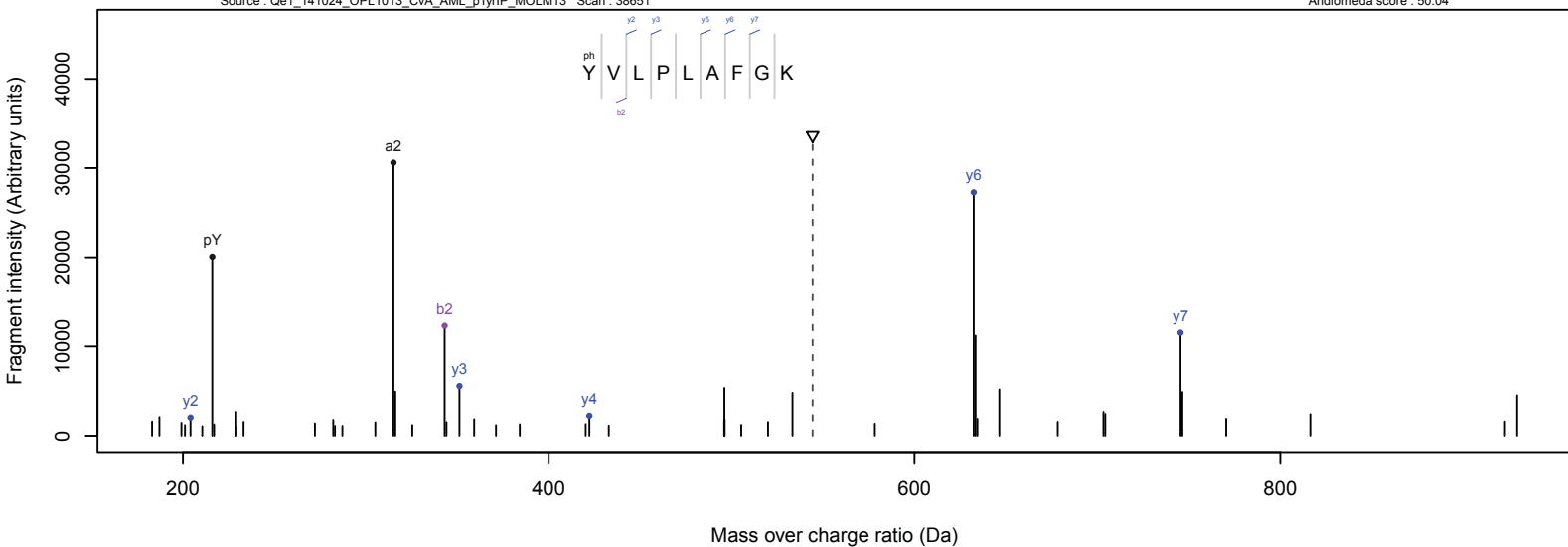
Andromeda score : 50.04



Y(ph)VLPLAFGK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 38651

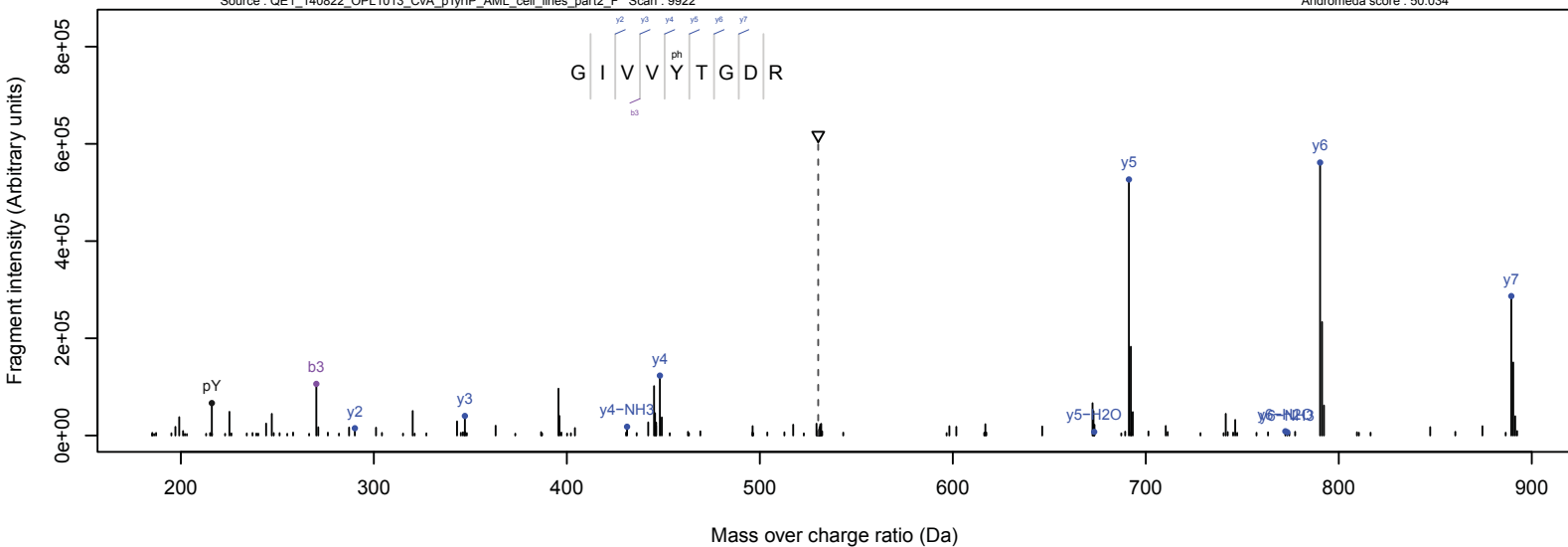
Andromeda score : 50.04



GIVVY(ph)TGDR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_F Scan : 9922

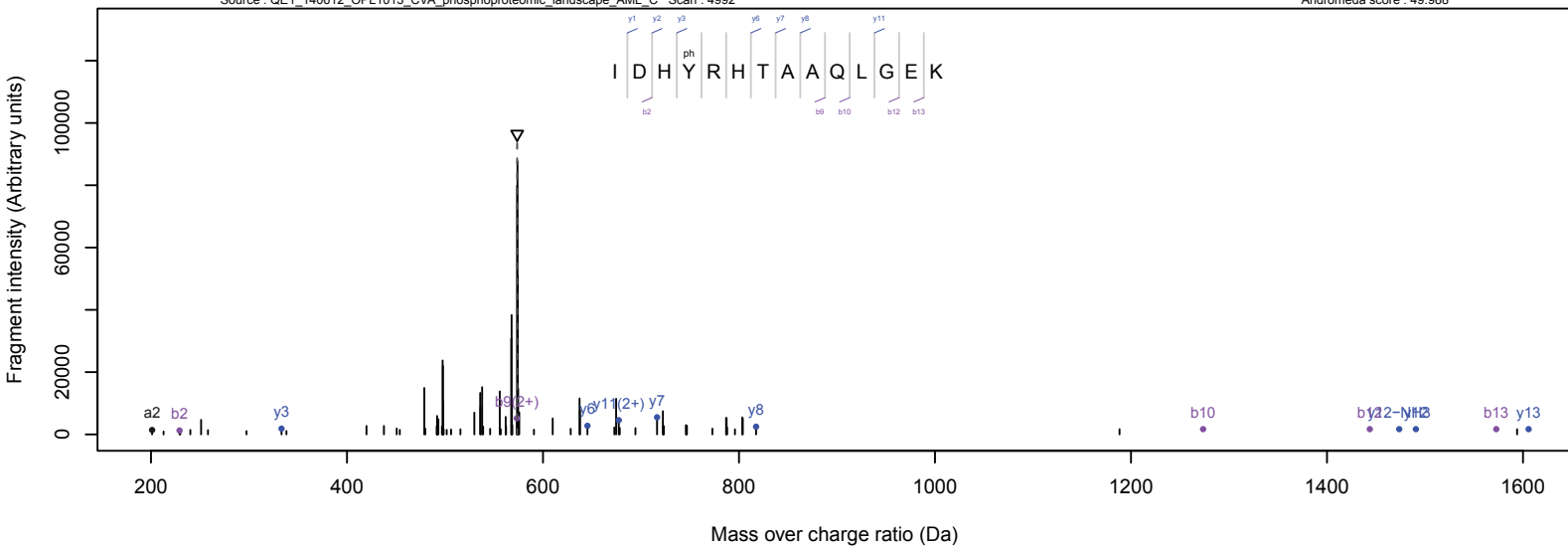
Andromeda score : 50.034



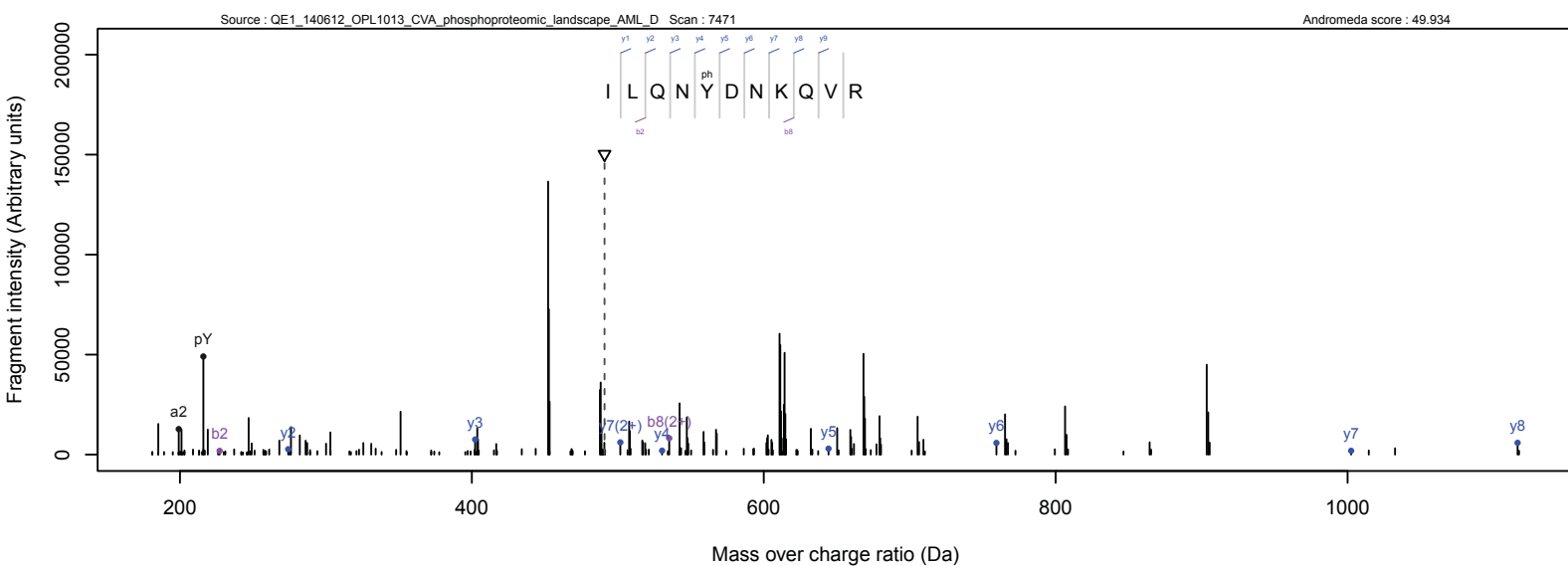
IDHY(ph)RHTAAQLGEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 4992

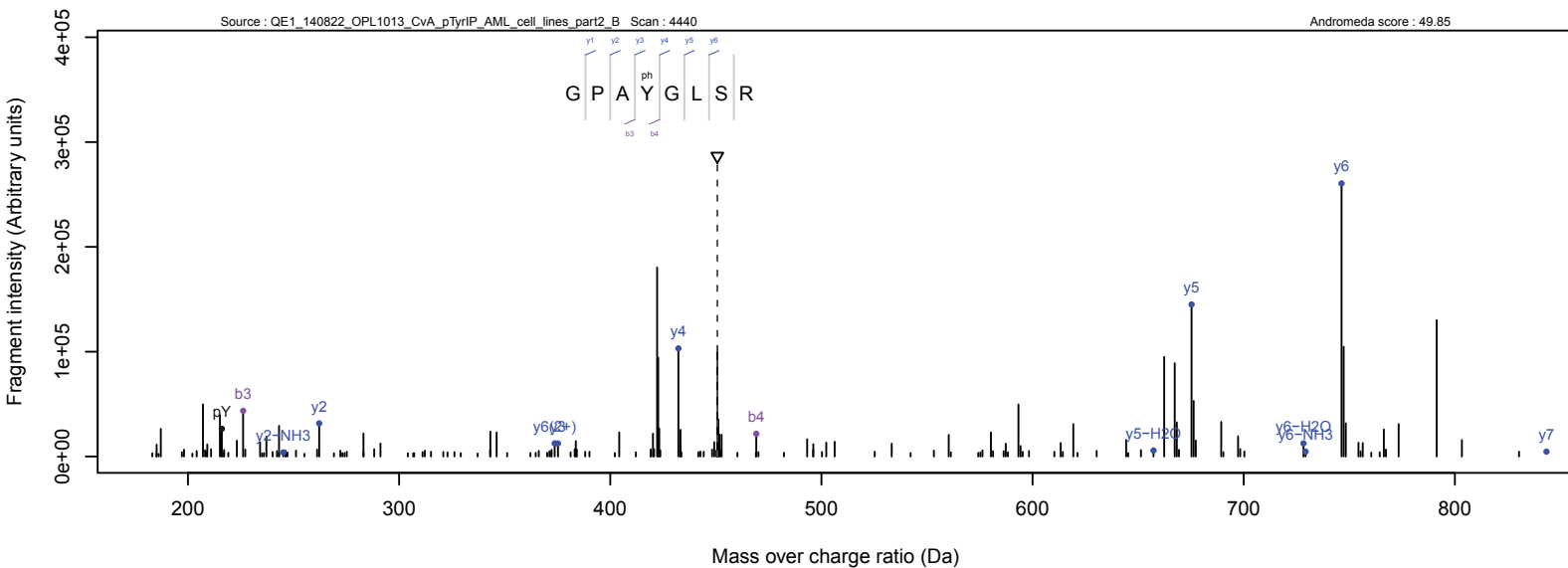
Andromeda score : 49.988



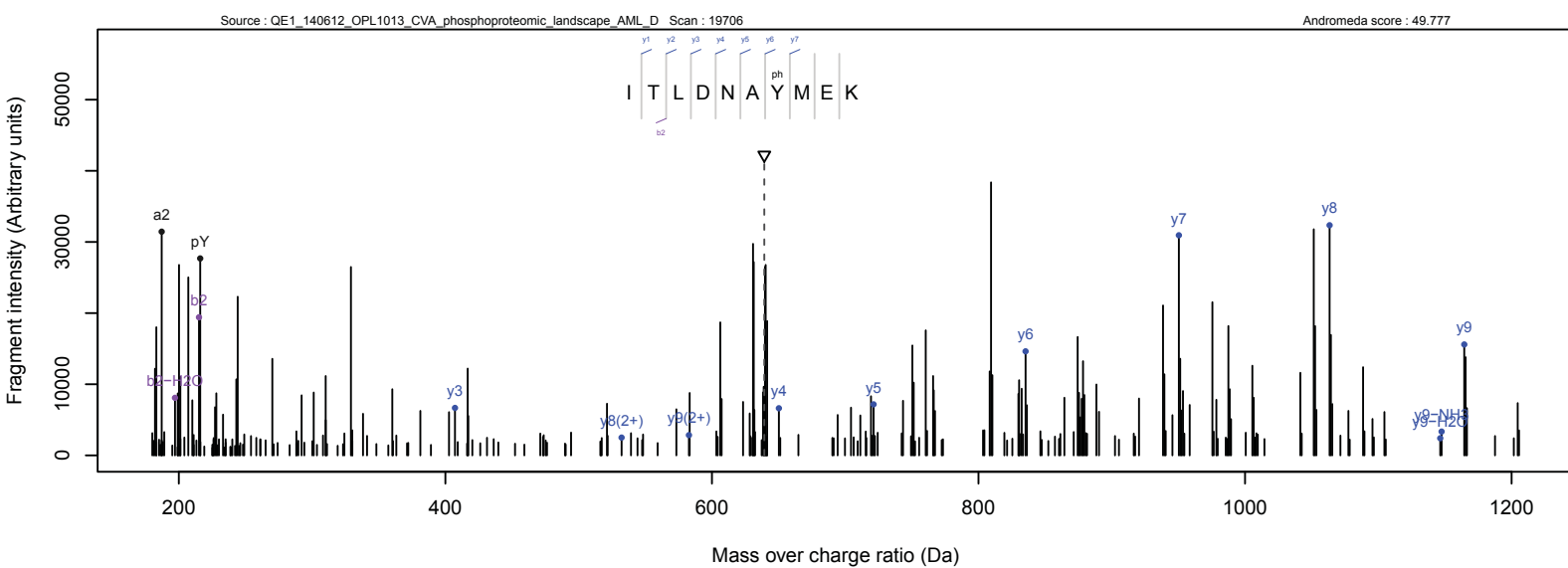
## ILQNY(ph)DNKQVR-3+



## GPAY(ph)GLSR-2+



## ITLDNAY(ph)MEK-2+

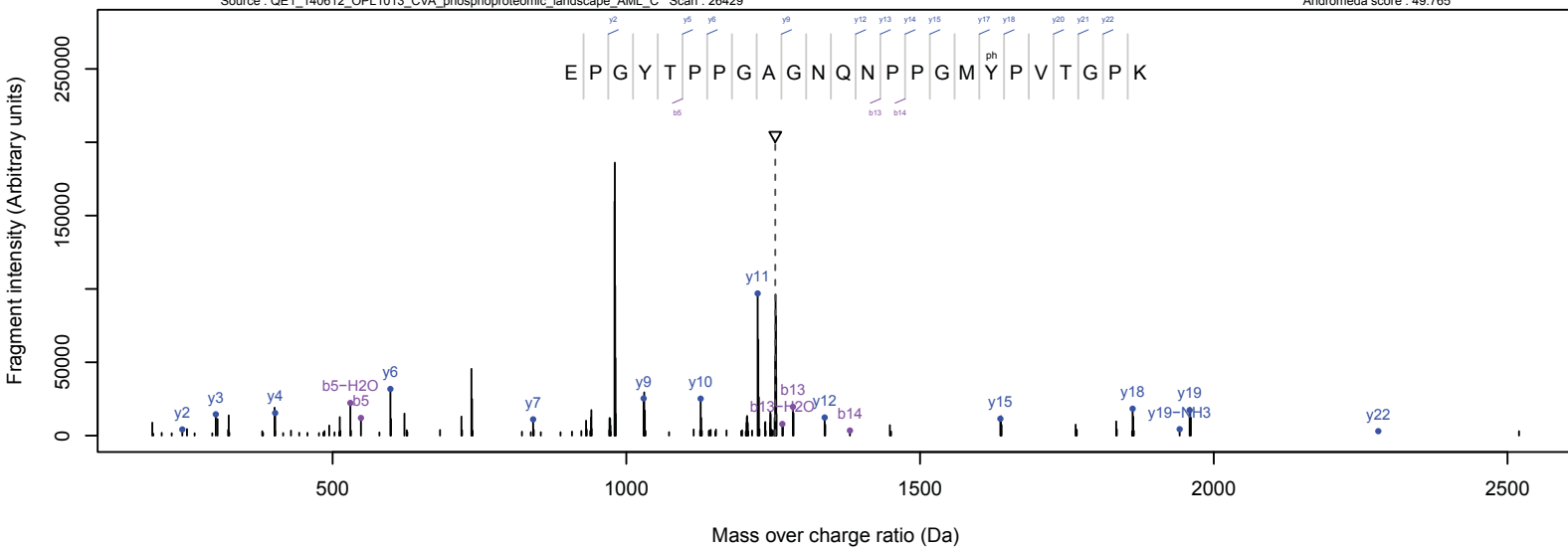




## EPGYTPPGAGNQNPpGMYpVVTGPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 26429

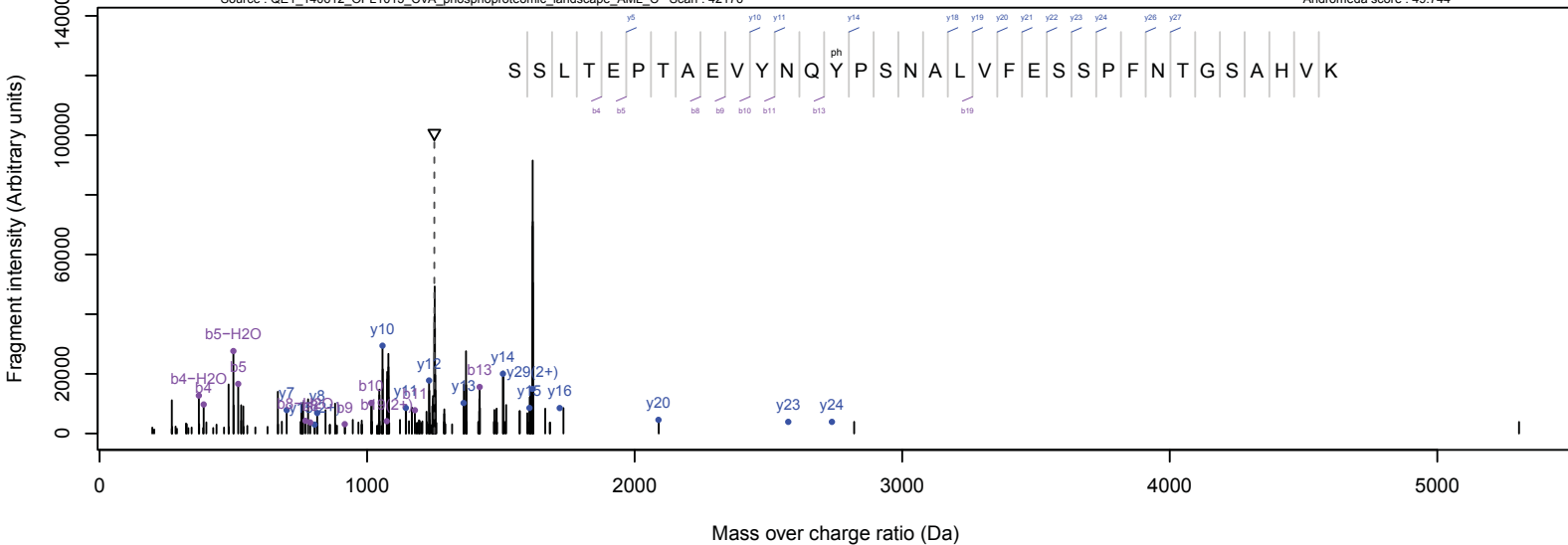
Andromeda score: 49.765



## SSLTEPTAEVYNQpYPSNALVFESSPFNTGSAHVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 42176

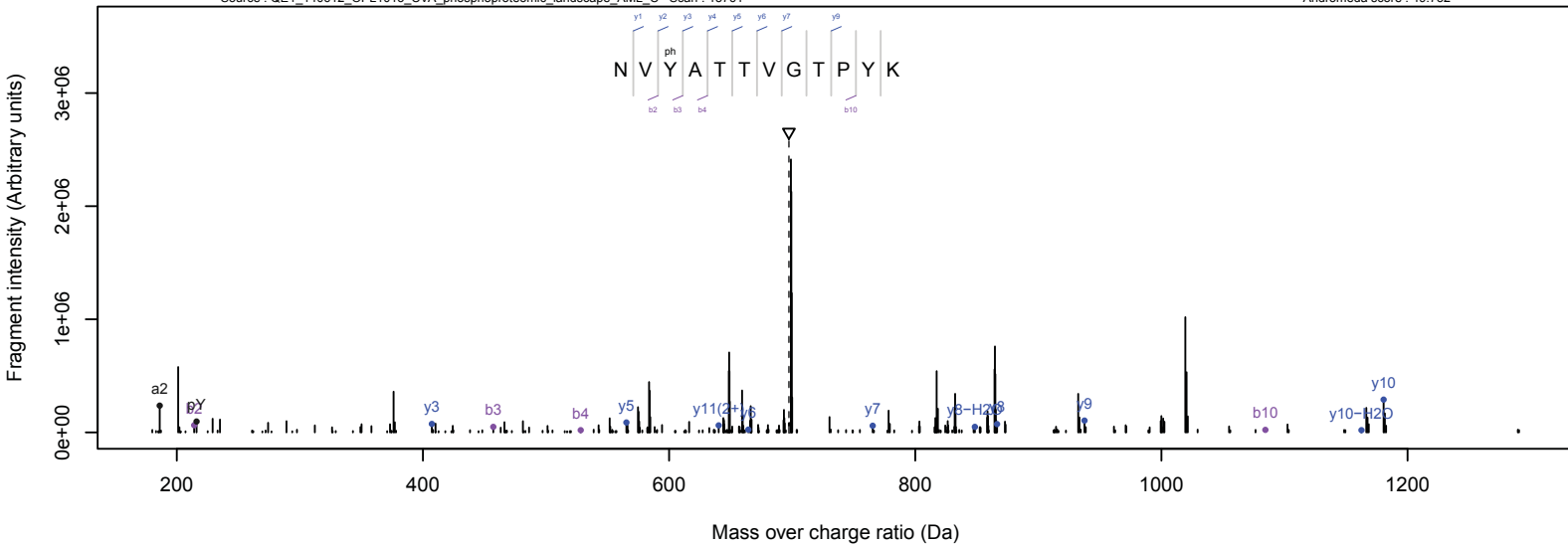
Andromeda score: 49.744



## NVYpATTVGTpYK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15761

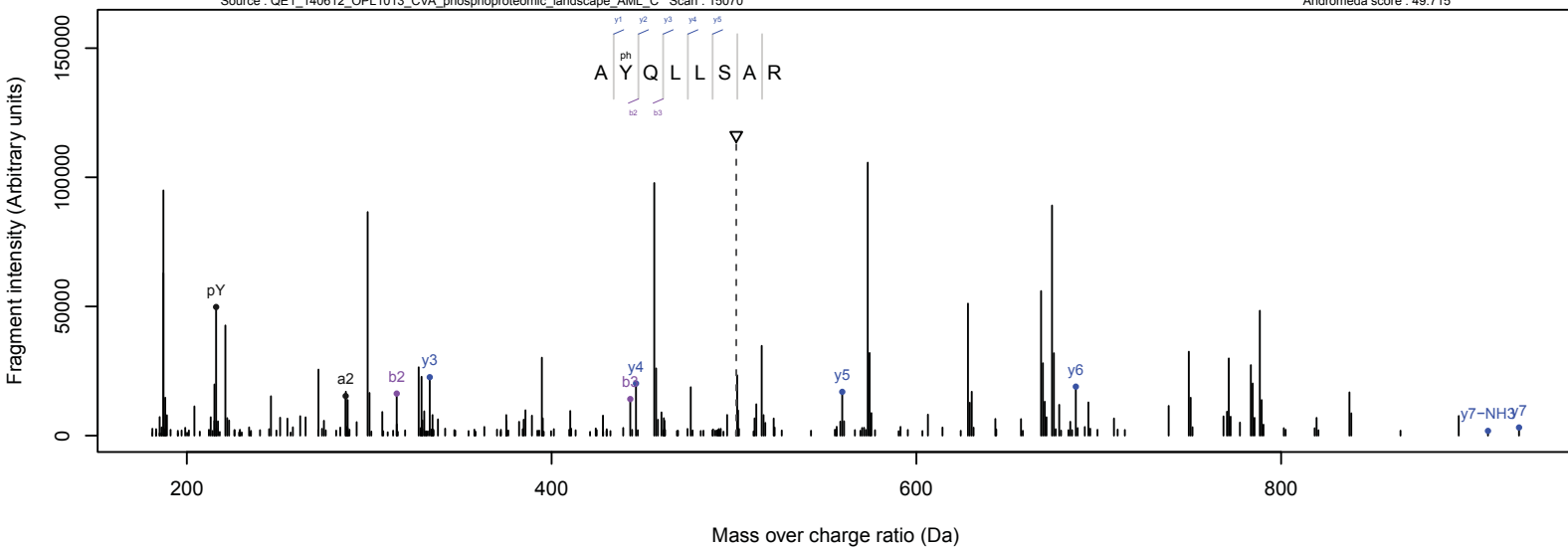
Andromeda score: 49.732



## AY(ph)QLLSAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15070

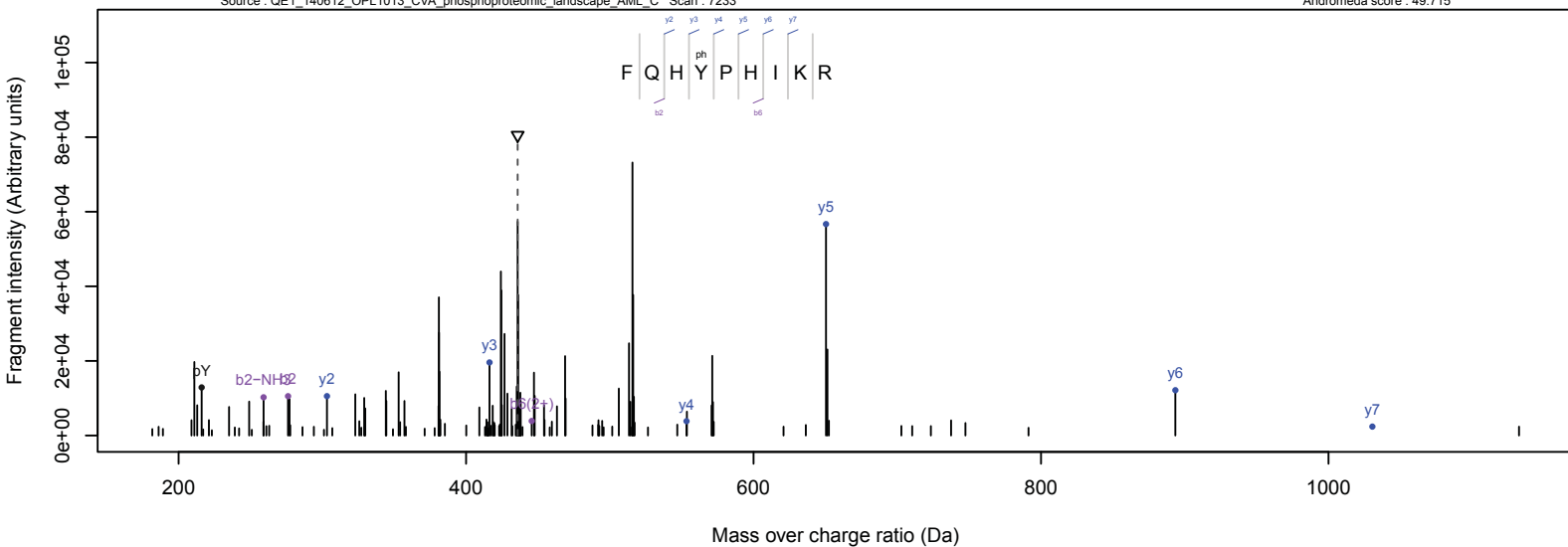
Andromeda score : 49.715



## FQHY(ph)PHIKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7233

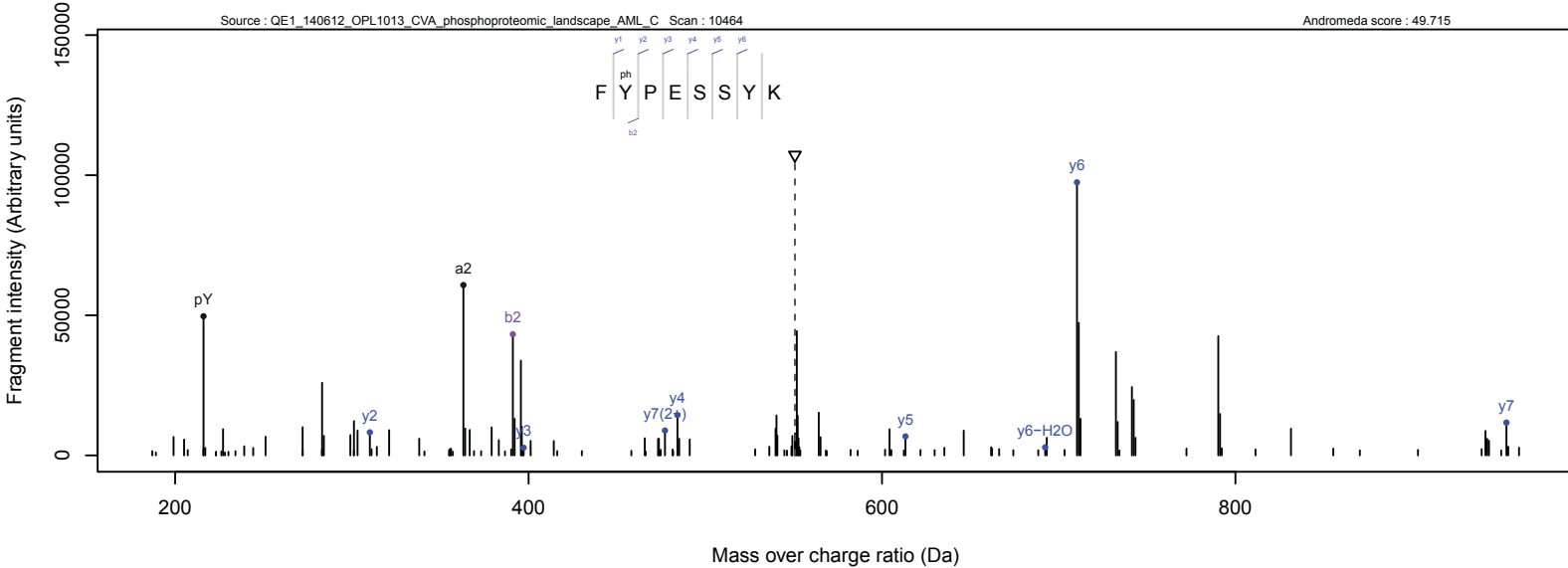
Andromeda score : 49.715



## FY(ph)PESSYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10464

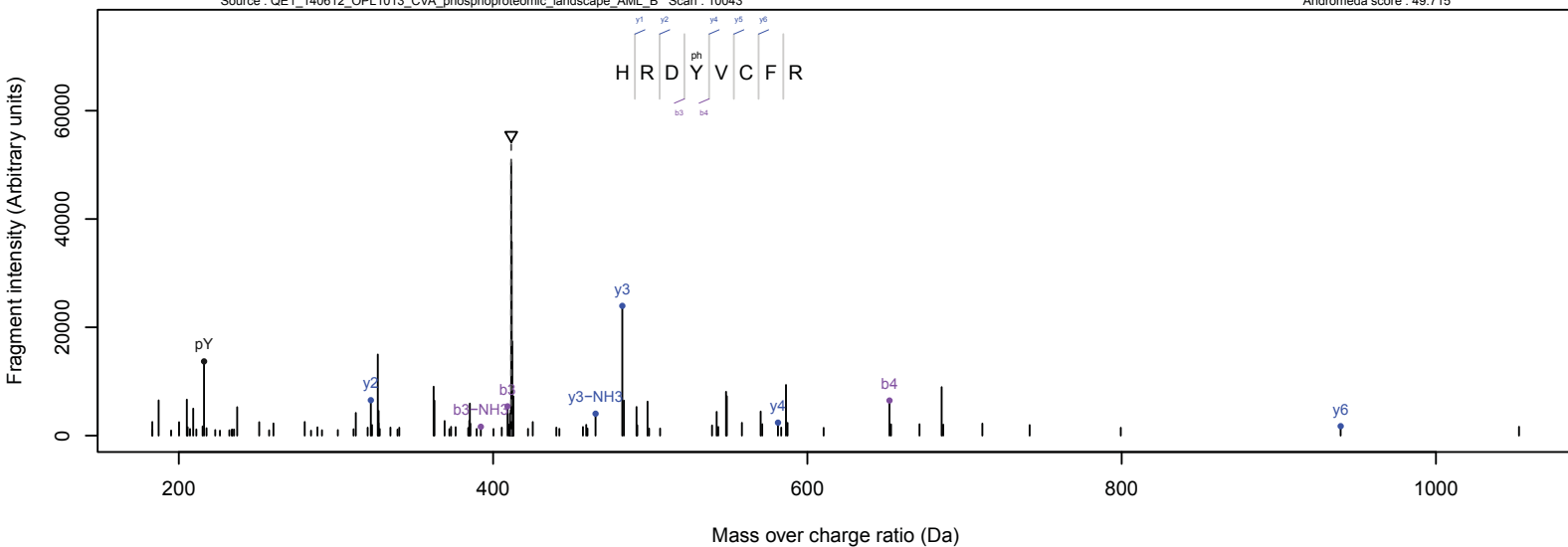
Andromeda score : 49.715



## HRDY(ph)VCFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10043

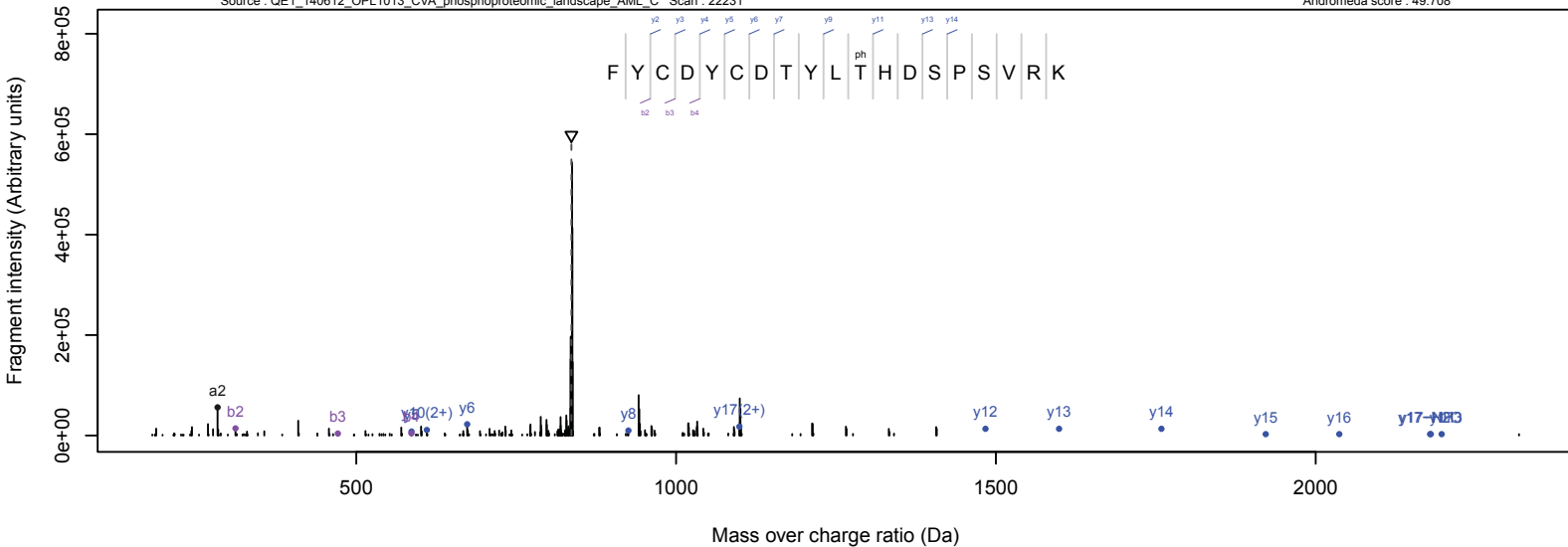
Andromeda score : 49.715



## FYCDYCDTYLT(ph)HDSPSVRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22231

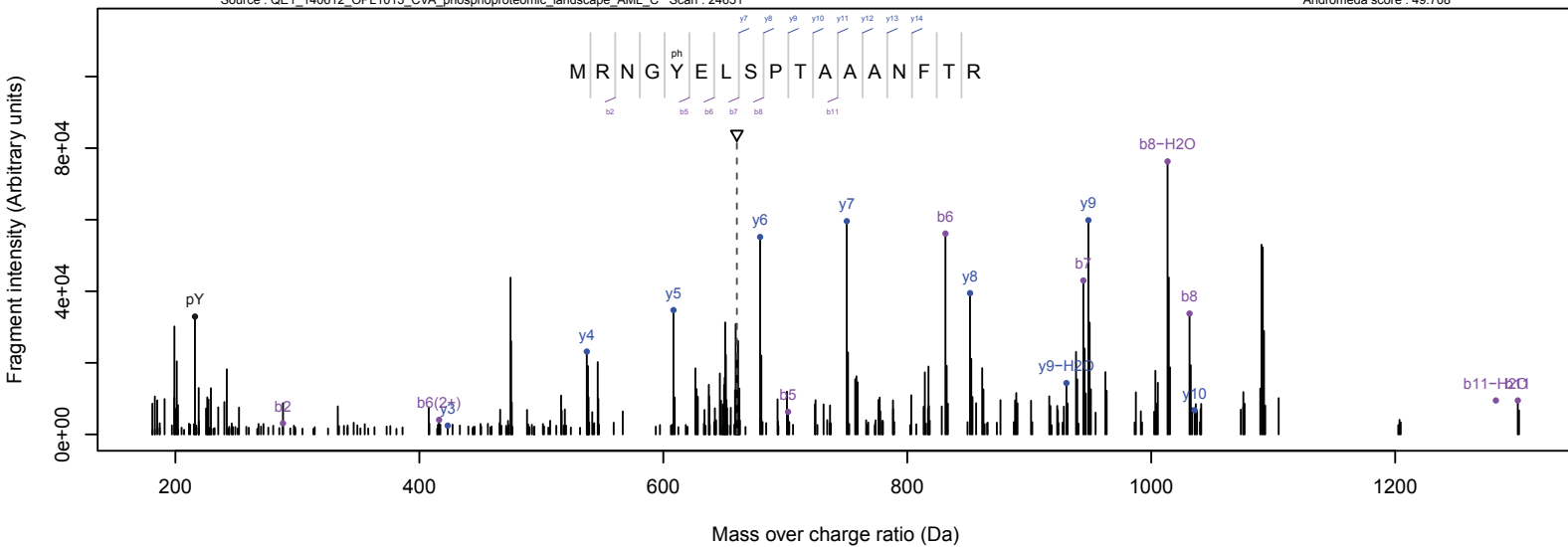
Andromeda score : 49.708



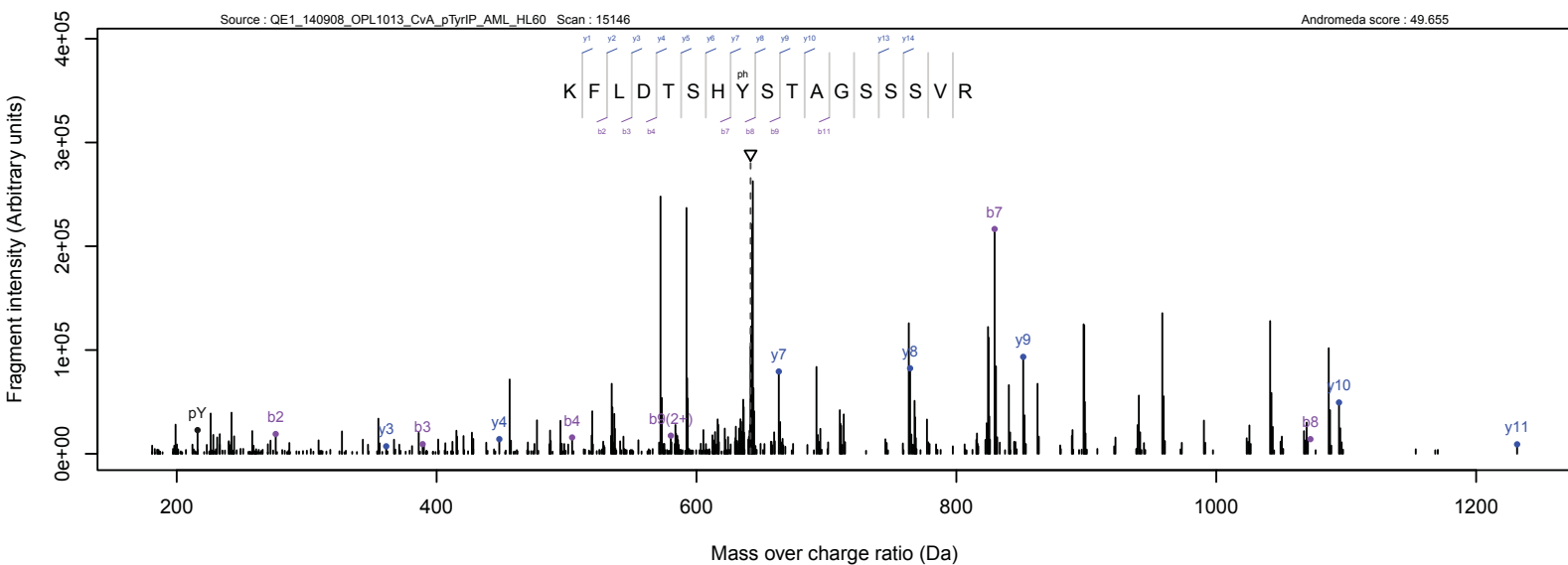
## MRNGY(ph)ELSPTAAANFTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24651

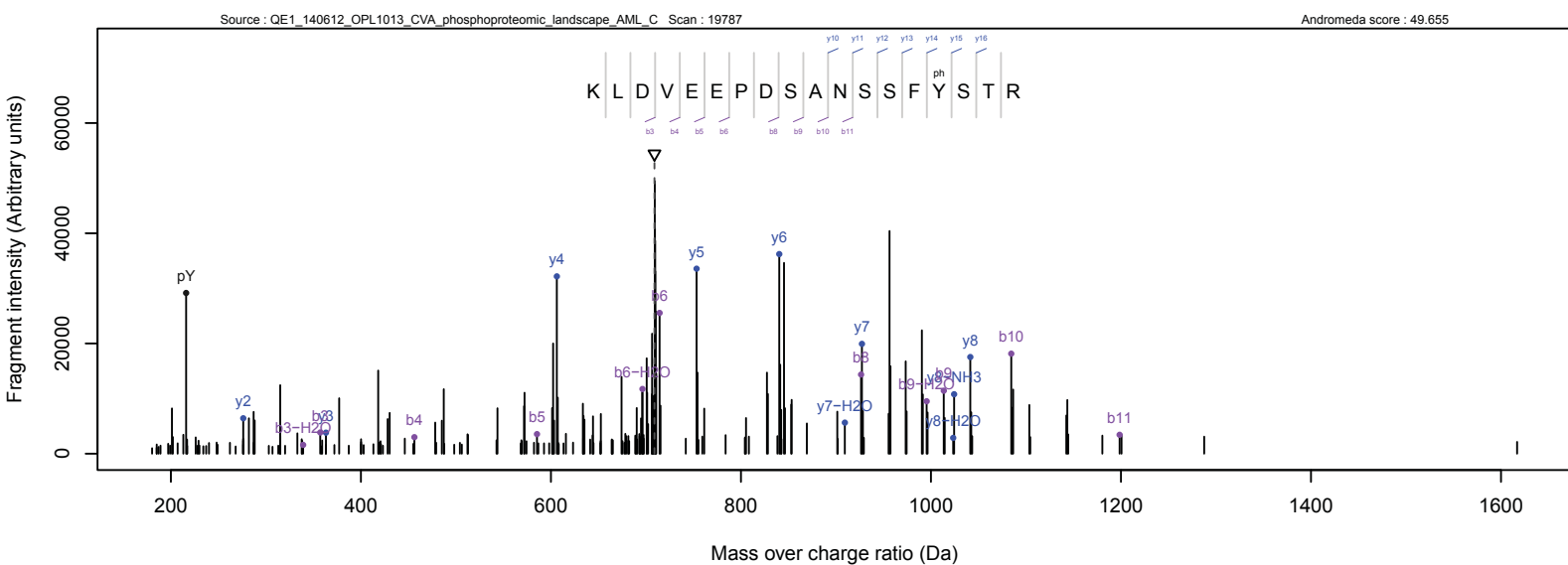
Andromeda score : 49.708



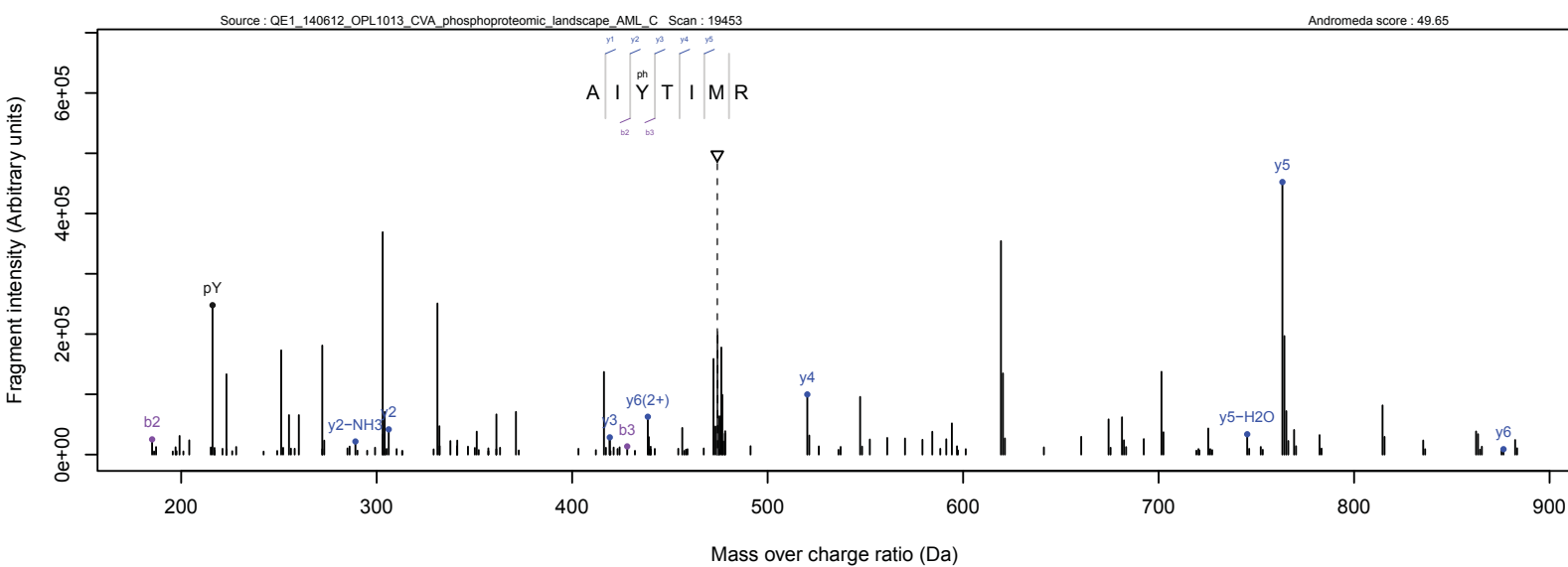
## KFLDTSHY(ph)STAGSSSVR-3+



## KLDVEEPDSANSS(ph)STR-3+



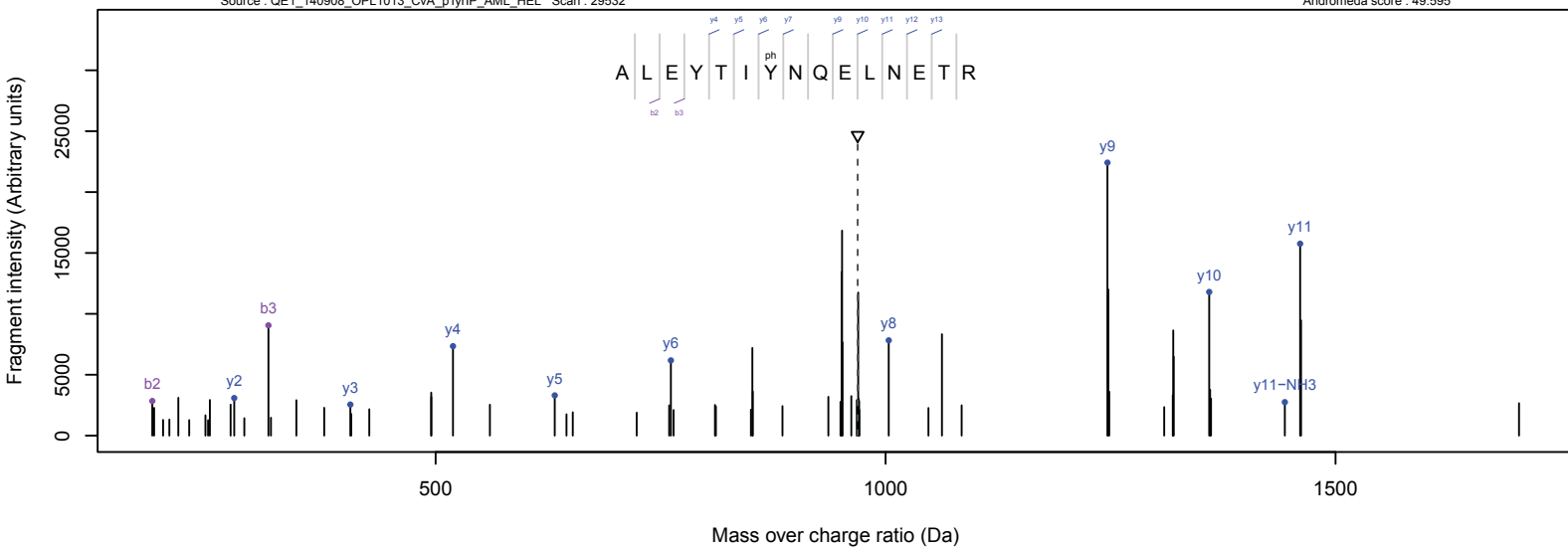
## AIY(ph)TMR-2+



## ALEYTIY(ph)NQLNETR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29532

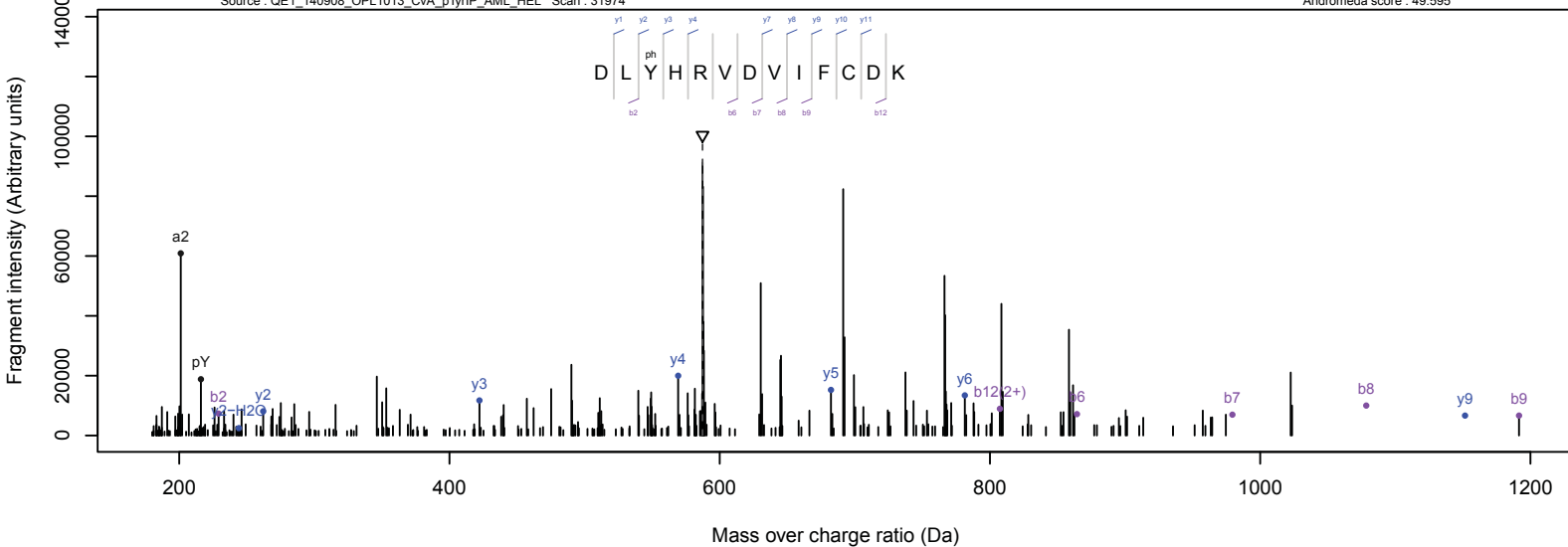
Andromeda score : 49.595



## DLY(ph)HRVDVIFCDK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 31974

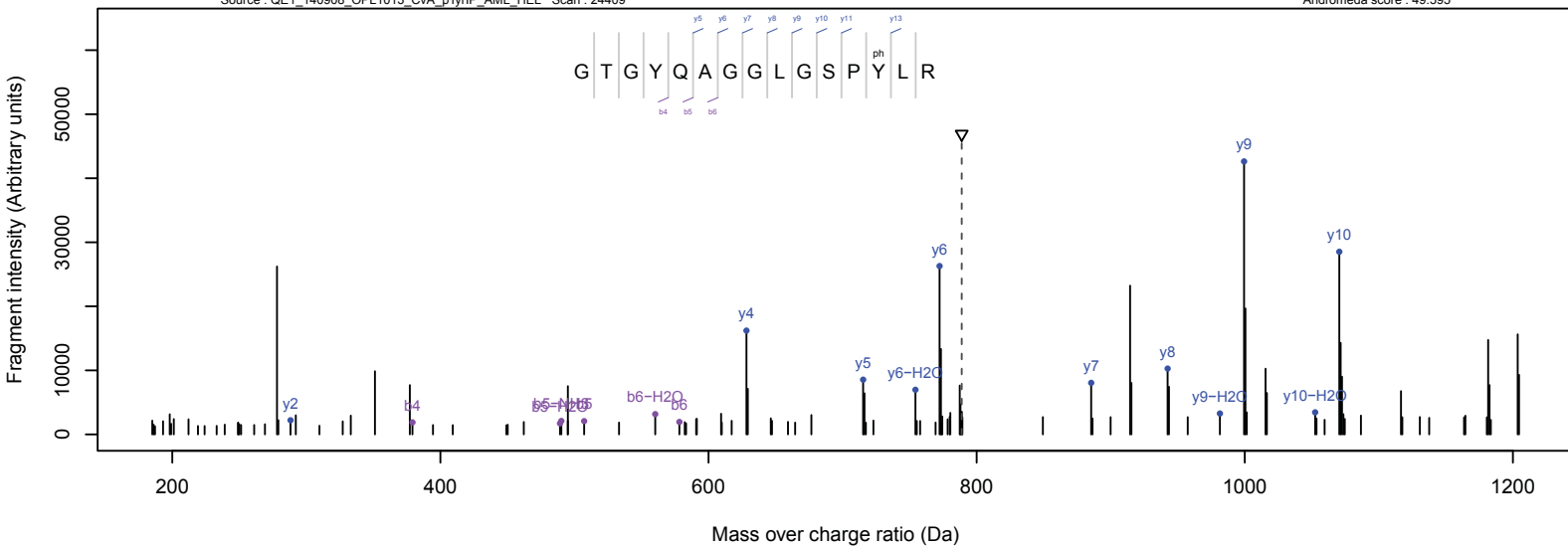
Andromeda score : 49.595



## GTGYQAGGLSPY(ph)LR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24409

Andromeda score : 49.595

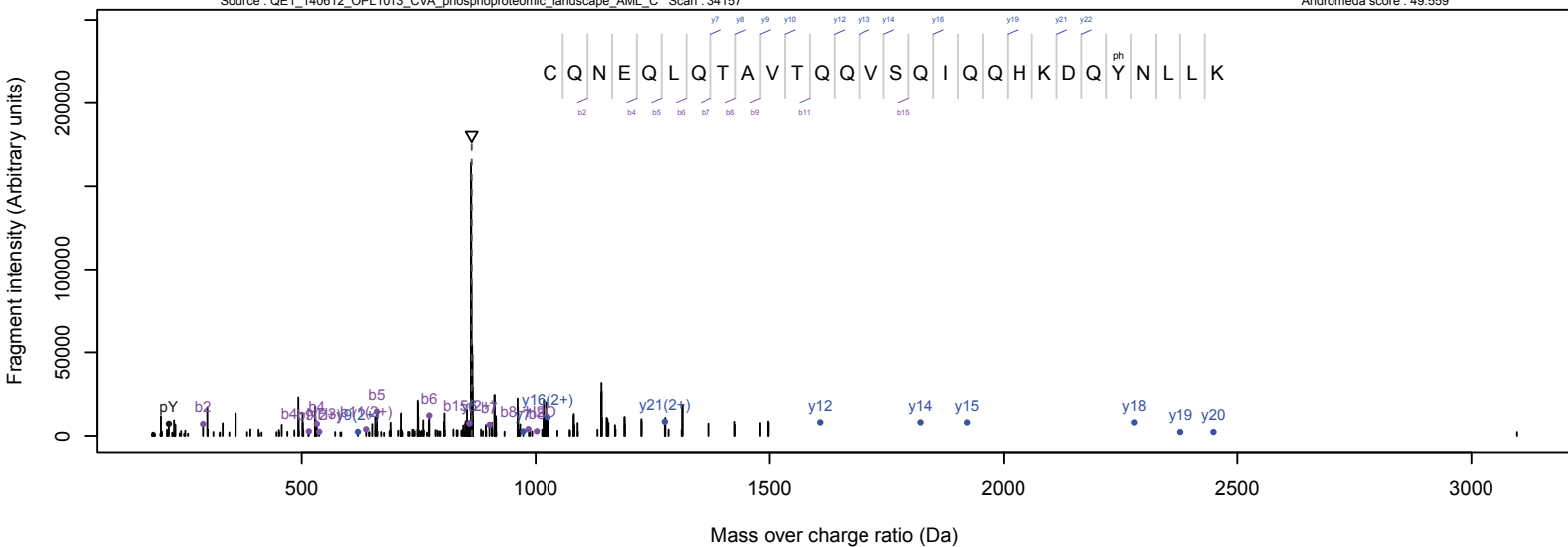




## CQNEQLTAVTQQVSIQQHKDQY(ph)NLLK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34157

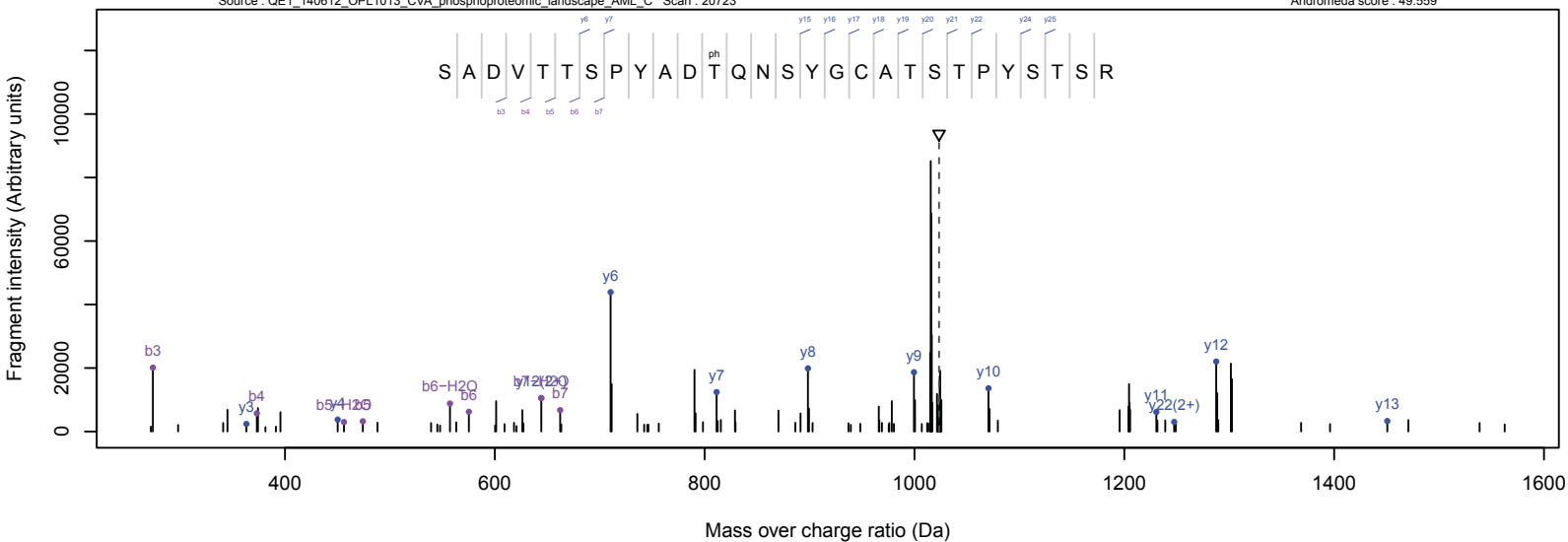
Andromeda score: 49.559



## SADVTTSPYADT(ph)QNSYGCATSTPYSTR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 20723

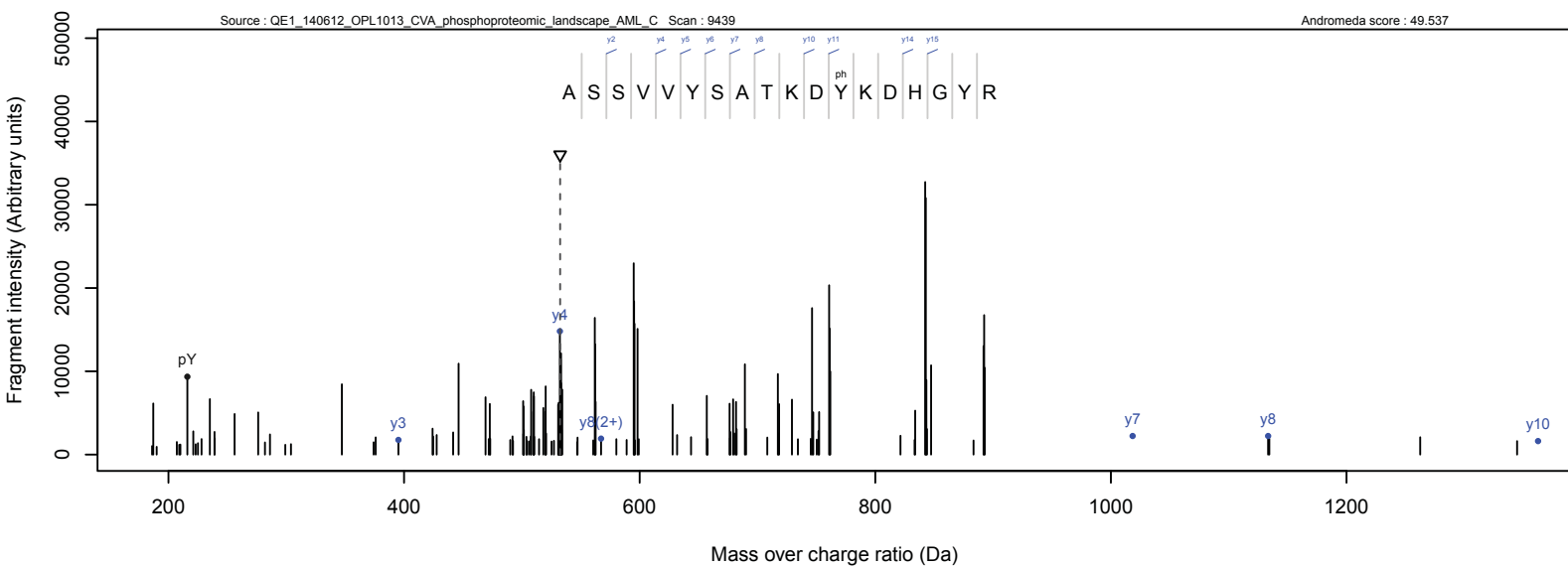
Andromeda score: 49.559



## ASSVVYSATKDY(ph)KDHYR-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 9439

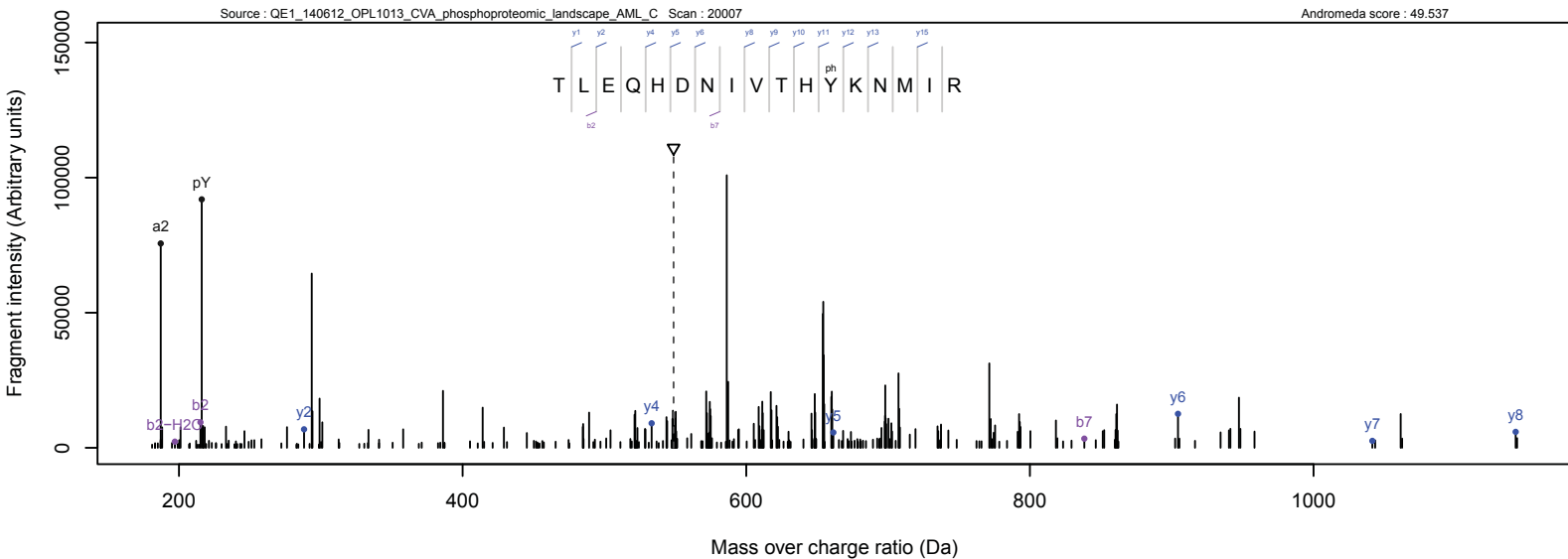
Andromeda score: 49.537



TLEQHDNIVTHYK<sup>ph</sup>NMIR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20007

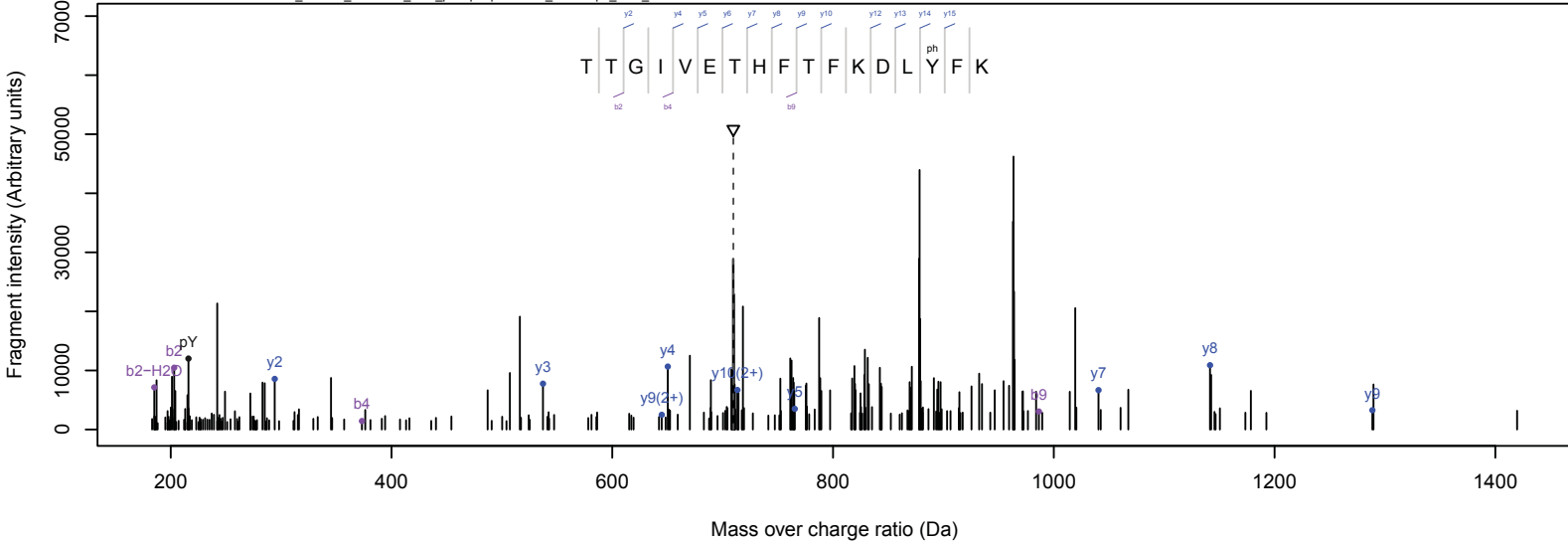
Andromeda score : 49.537



TTGIVETHFTKDL<sup>ph</sup>YFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39921

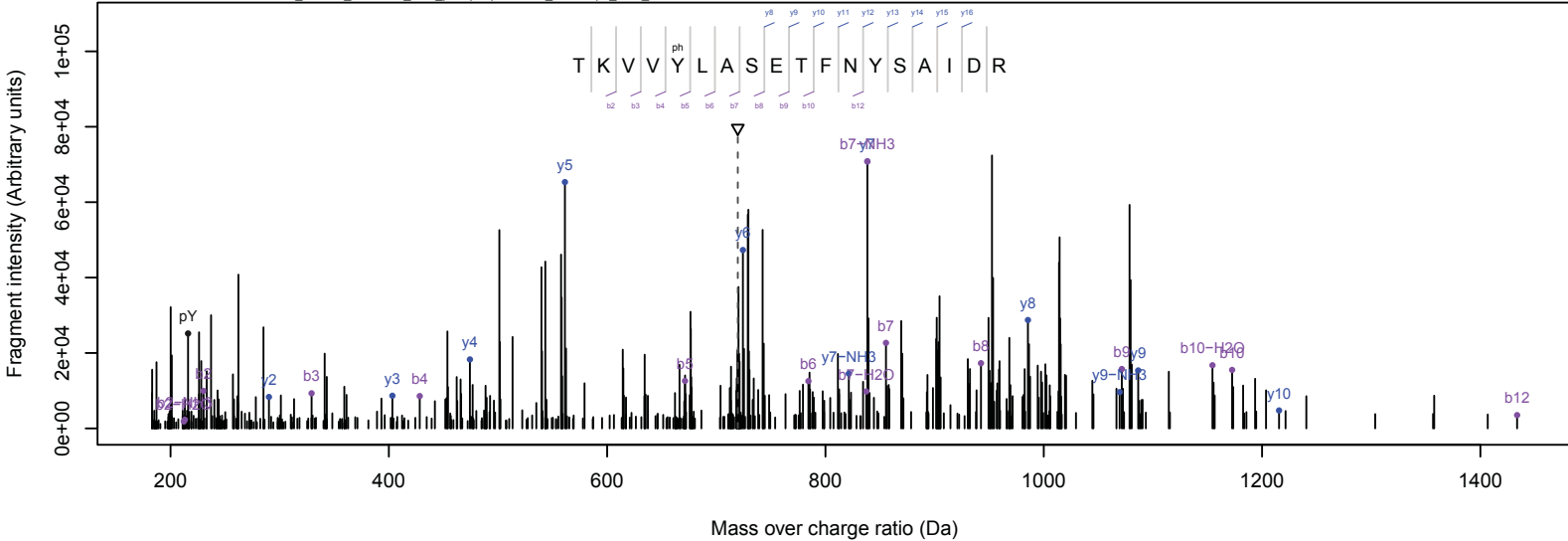
Andromeda score : 49.537



TKVVY<sup>ph</sup>LASETFNYSIDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37772

Andromeda score : 49.528

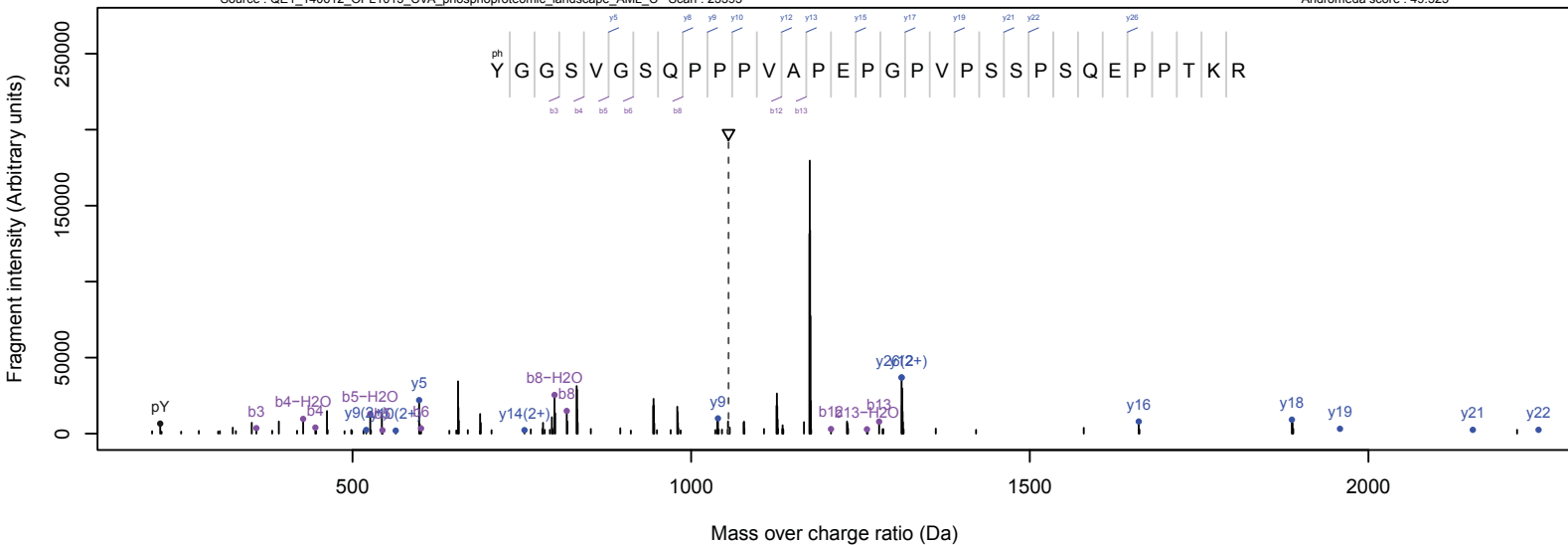




**Y(ph)GGVSGSQPPPVAPEPGVPSSPSQEPPTKR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23395

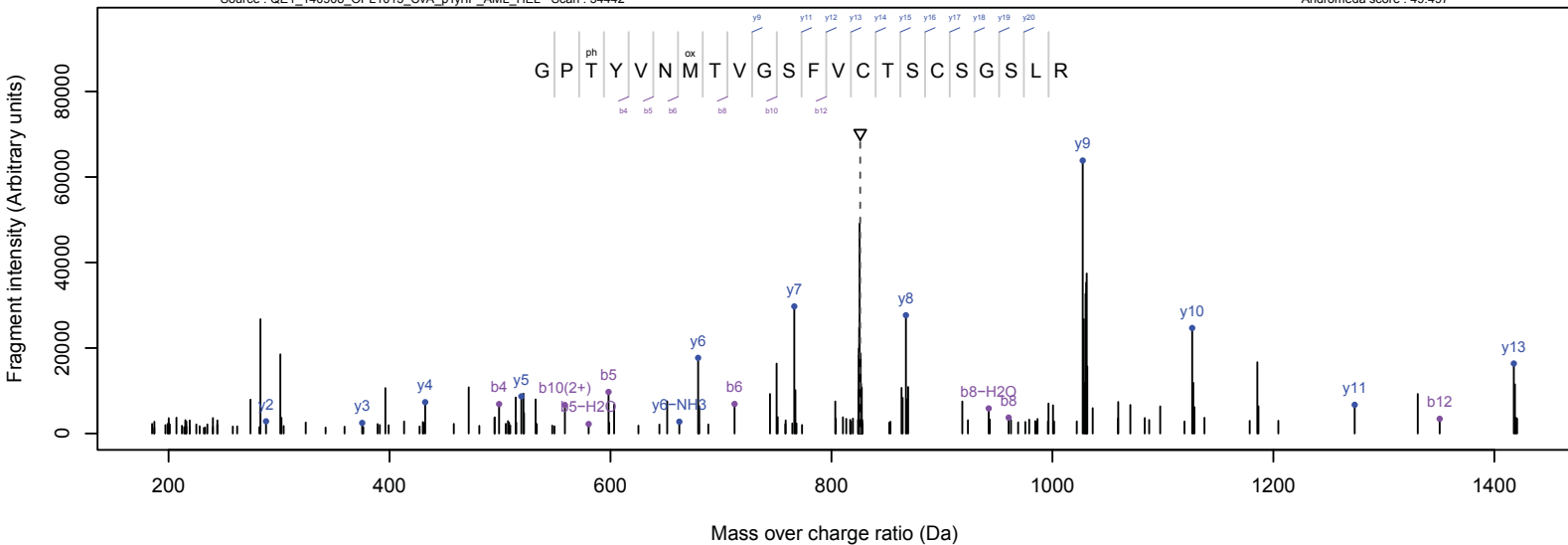
Andromeda score : 49.525



**GPT(ph)YVNM(ox)TVGSFVCTSCSGSLR-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 34442

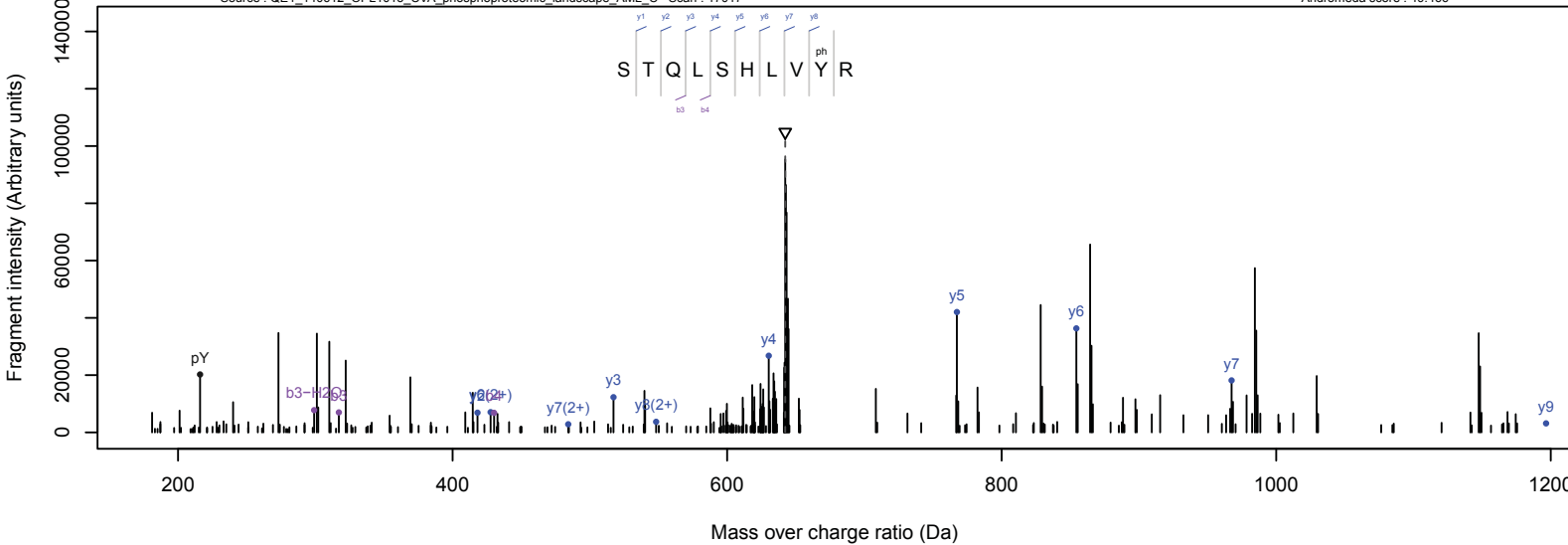
Andromeda score : 49.497



**STQLSHLVY(ph)R-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17017

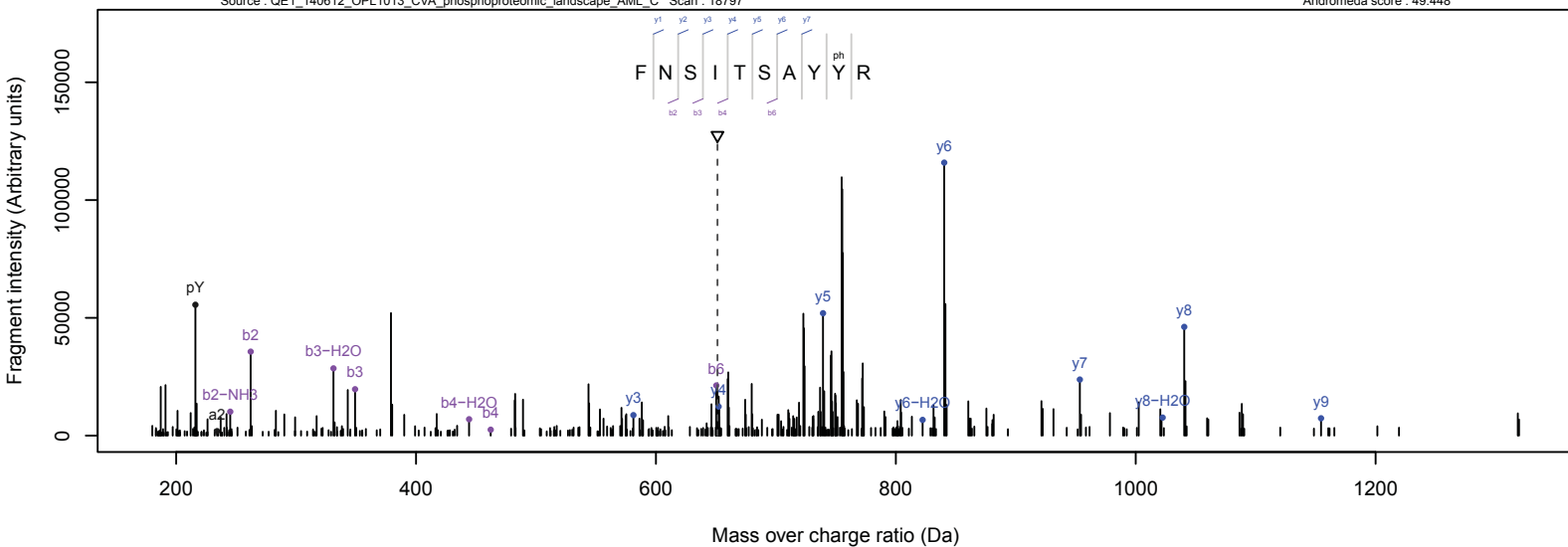
Andromeda score : 49.466



## FNSITSAY(ph)R-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 18797

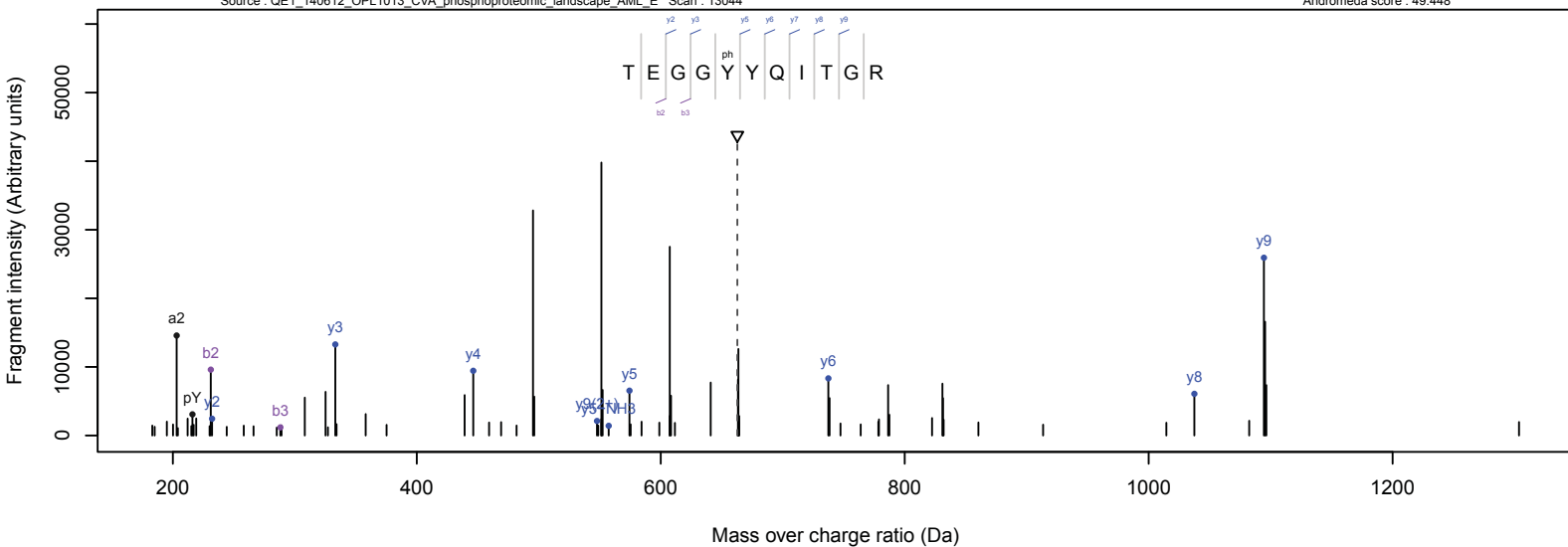
Andromeda score: 49.448



## TEGGY(ph)YQITGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 13044

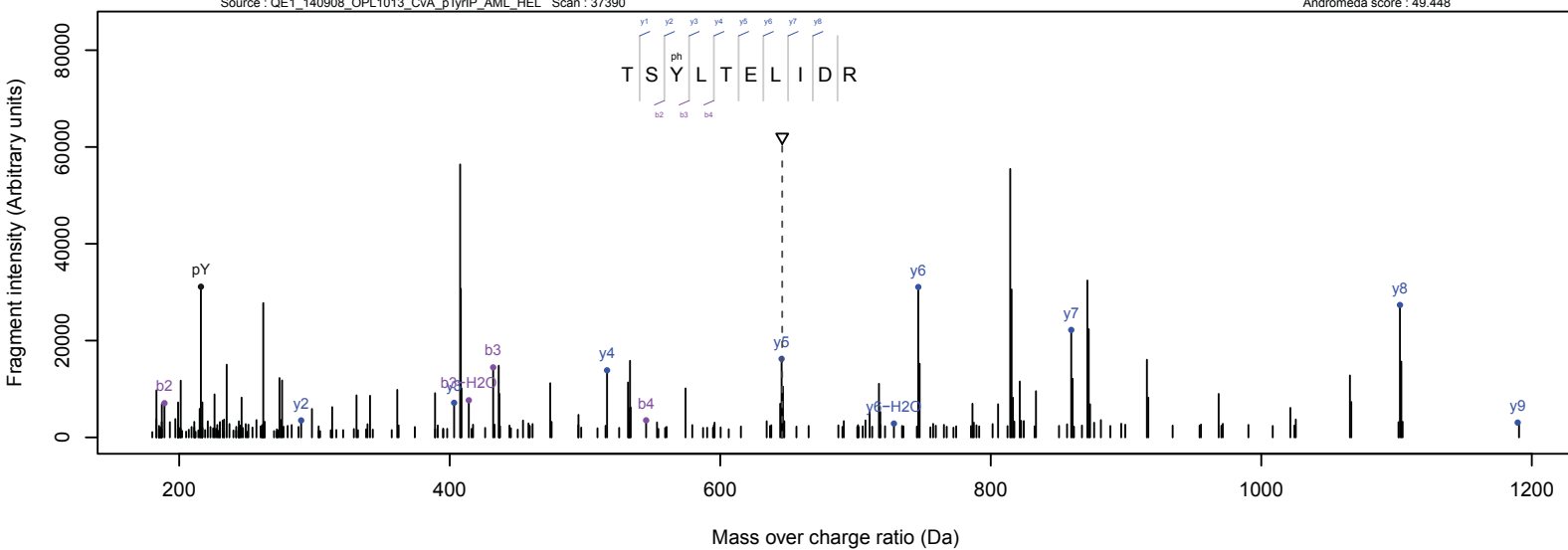
Andromeda score: 49.448



## TSY(ph)LTELIDR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 37390

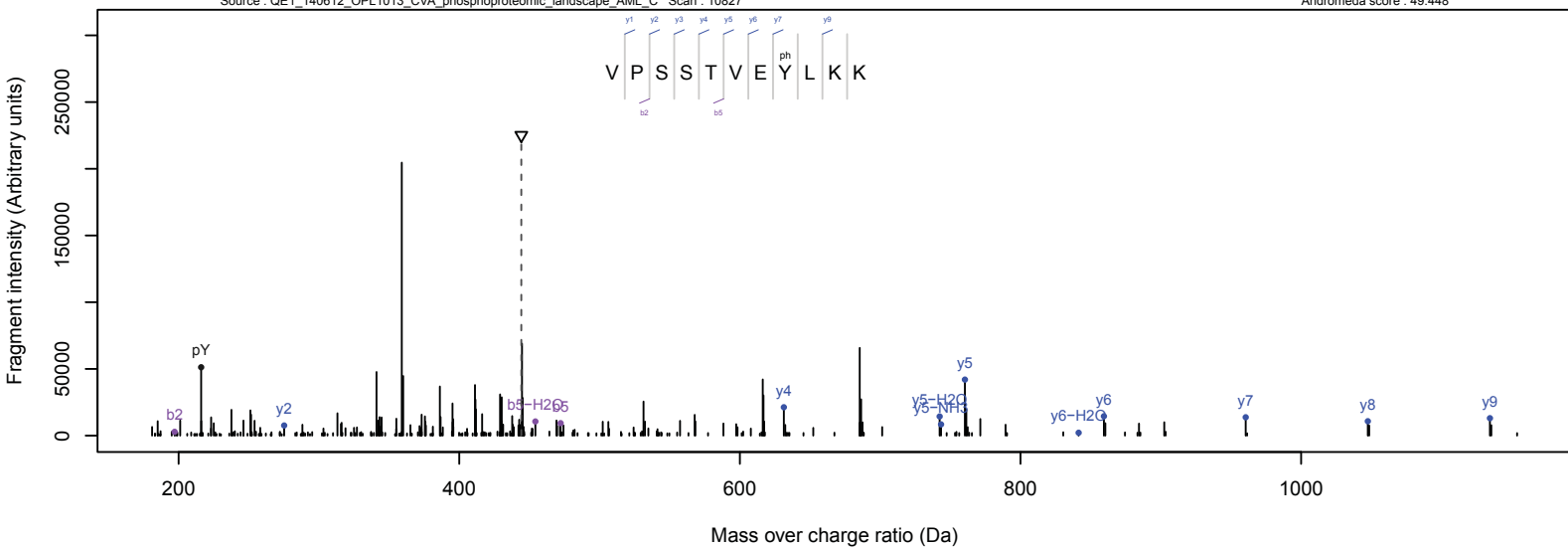
Andromeda score: 49.448



## VPSSTVEY(ph)LKK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10827

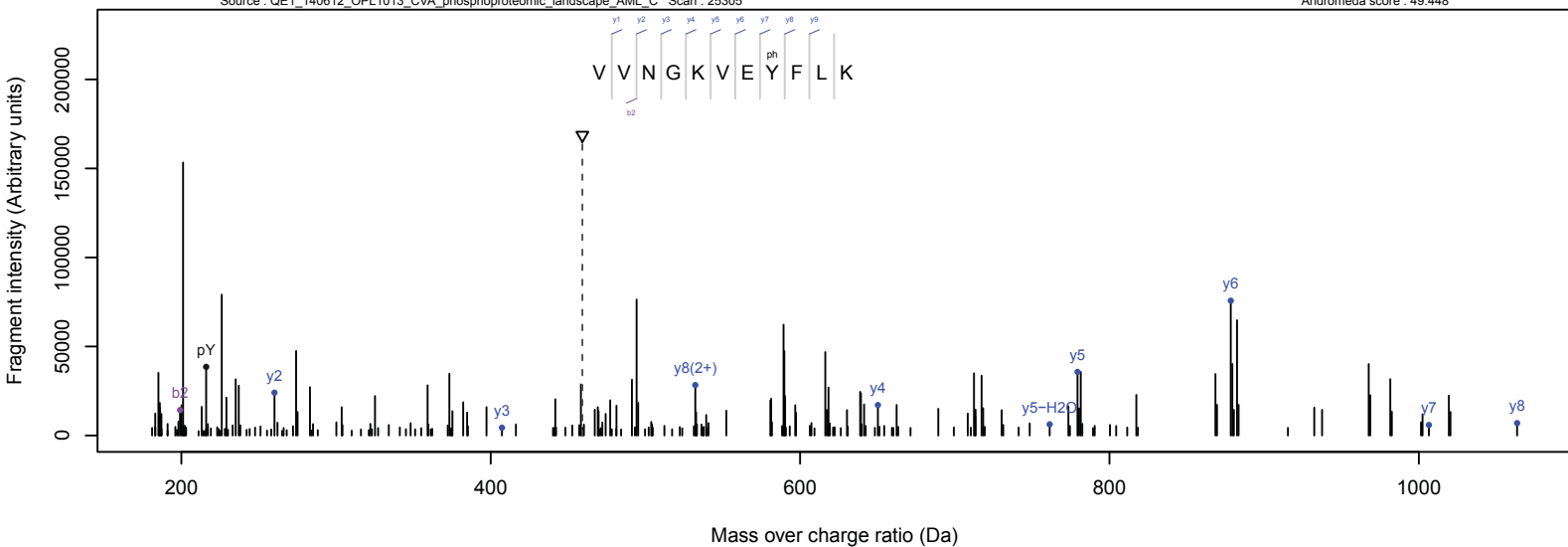
Andromeda score: 49.448



## VVNGKVEY(ph)FLK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 25305

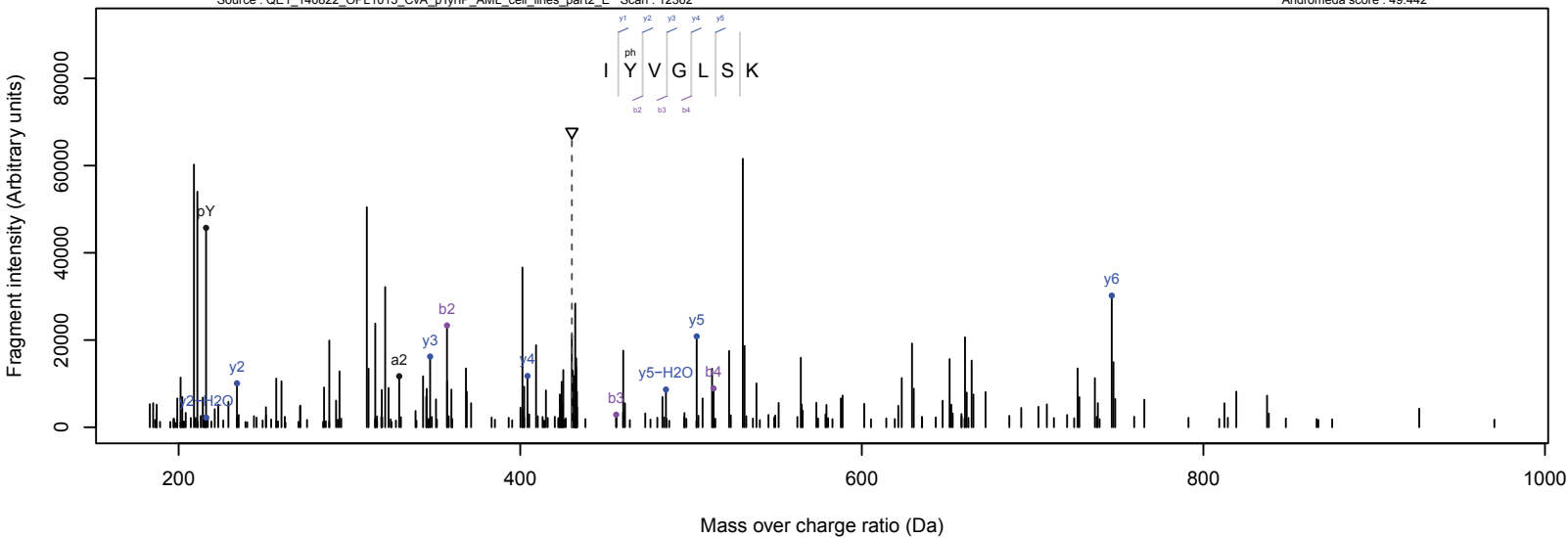
Andromeda score: 49.448



## IY(ph)VGLSK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 12362

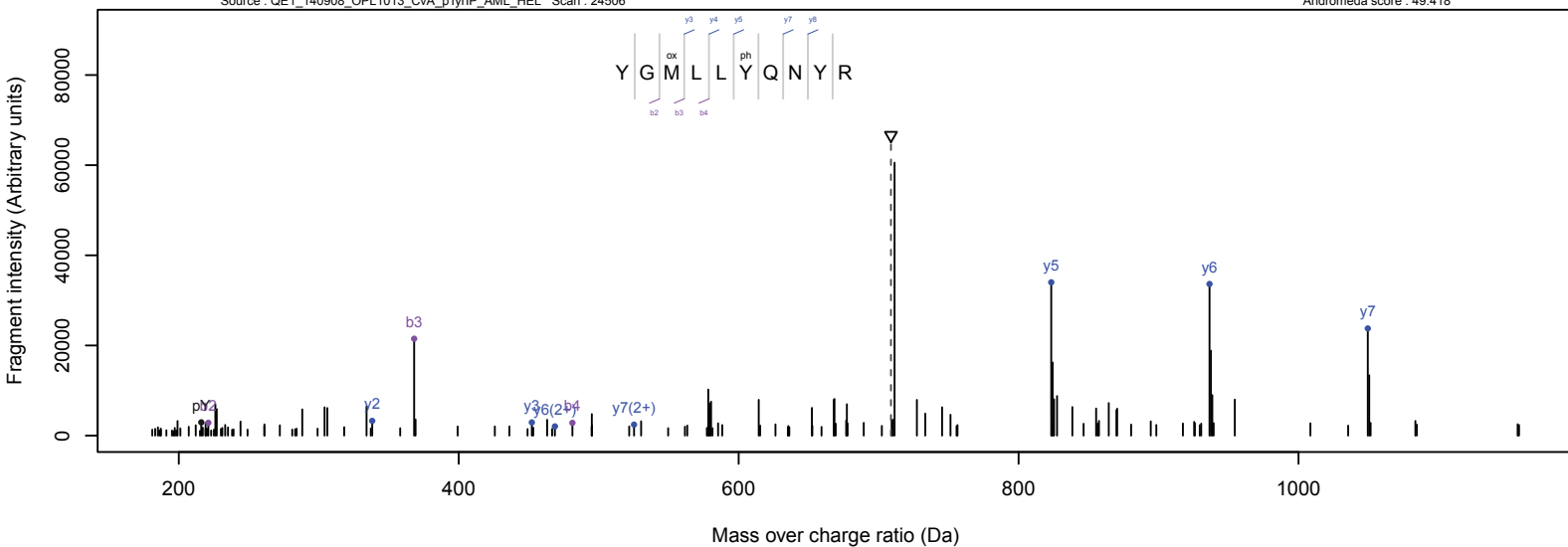
Andromeda score: 49.442



## YGM(ox)LLY(ph)QNYR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24506

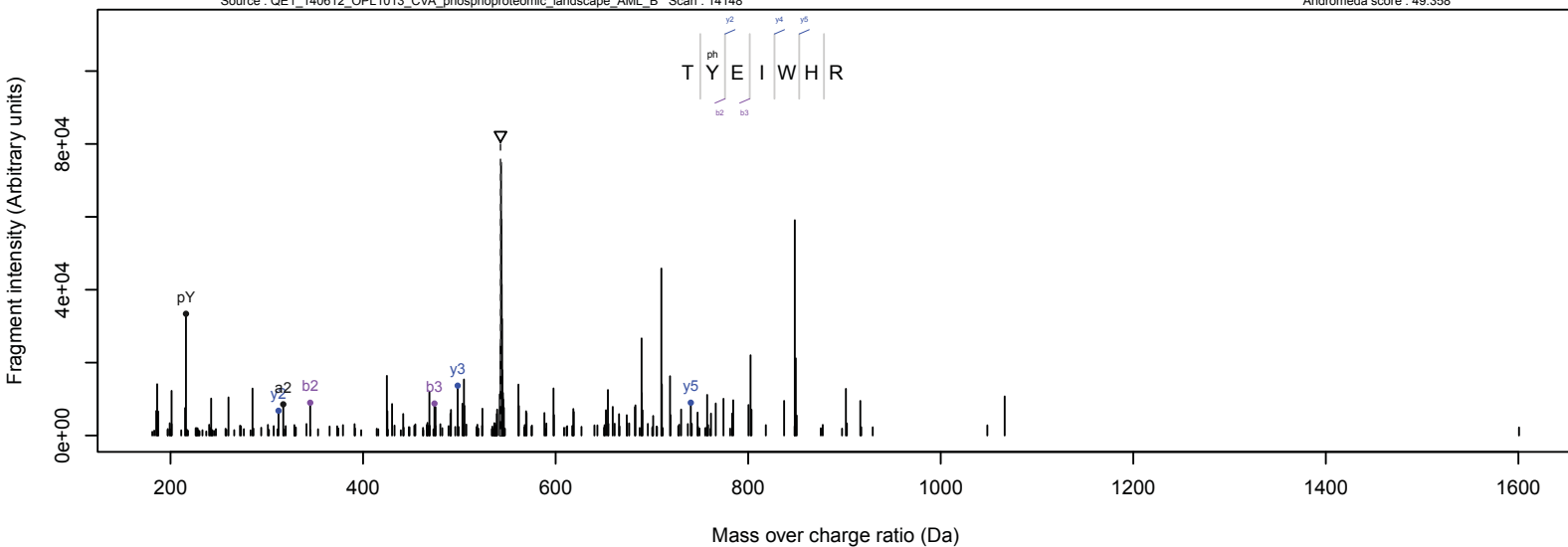
Andromeda score : 49.418



## TY(ph)EIWHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14148

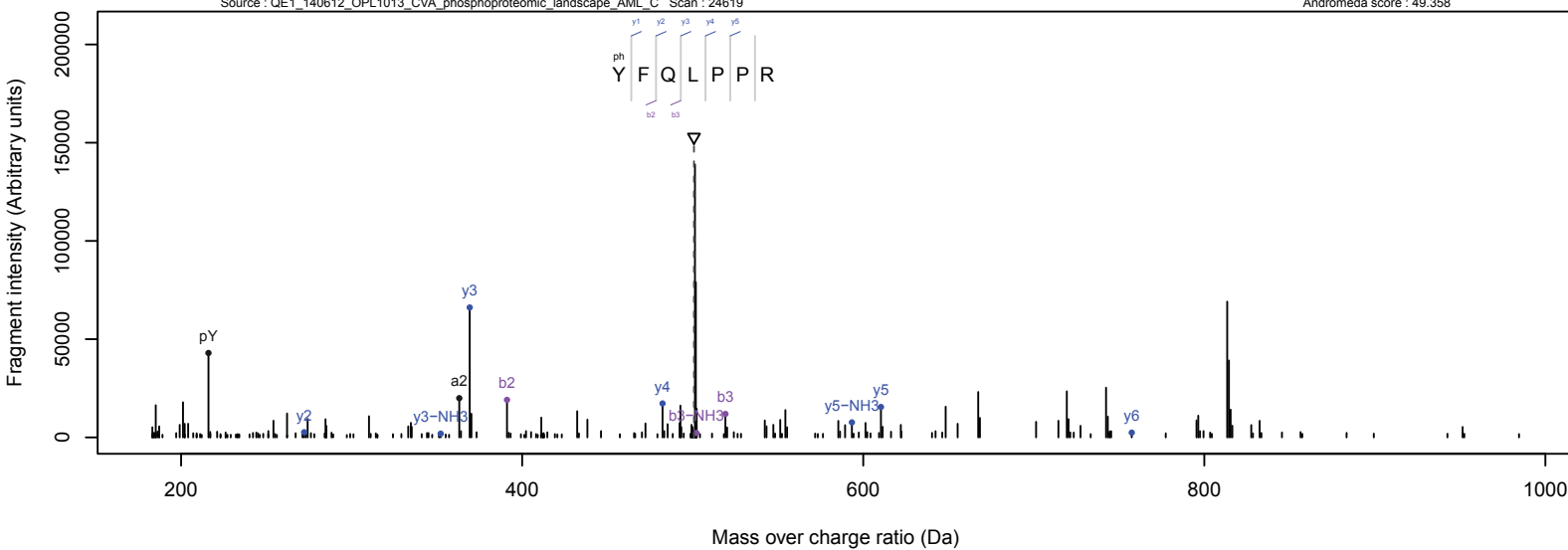
Andromeda score : 49.358



## Y(ph)FQLPPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24619

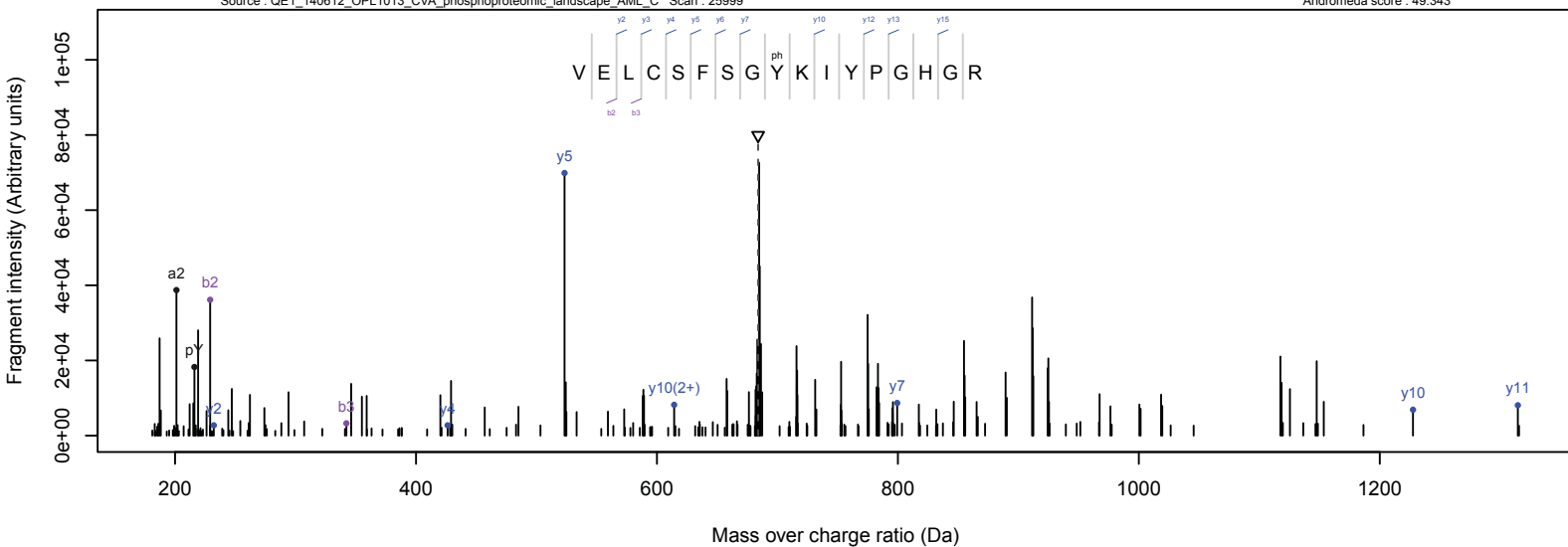
Andromeda score : 49.358



## VELCSFGY(ph)KIYPGHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25999

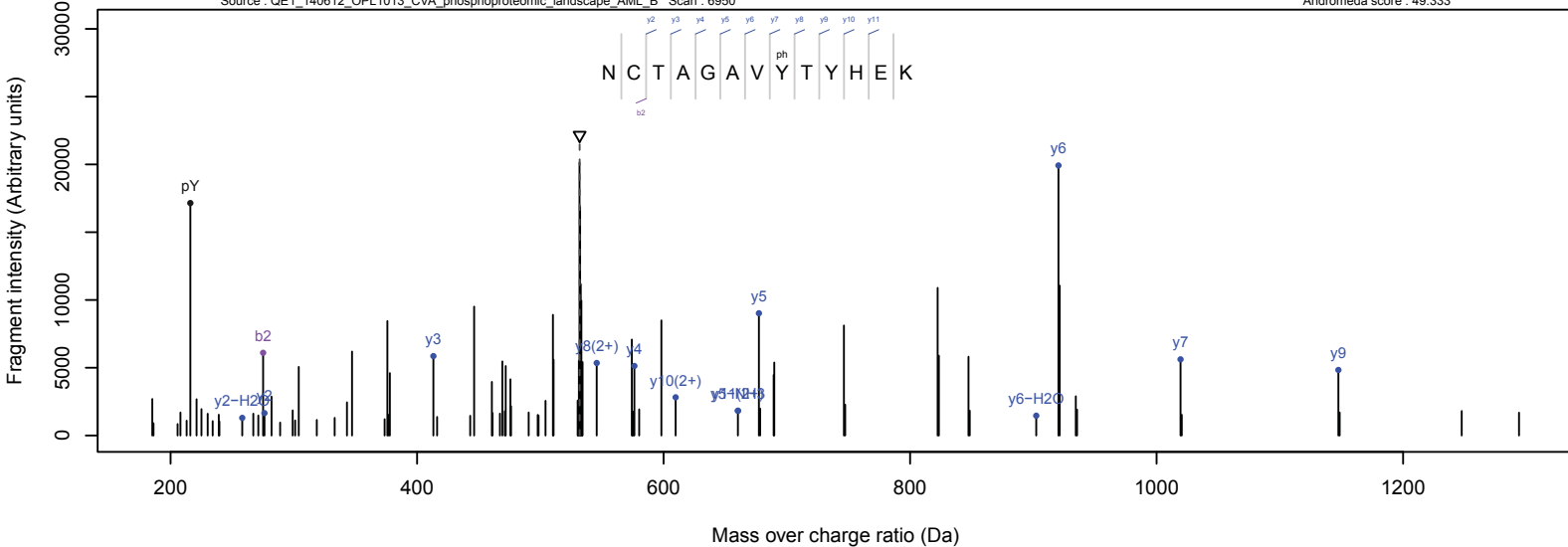
Andromeda score : 49.343



## NCTAGAVY(ph)TYHEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 6950

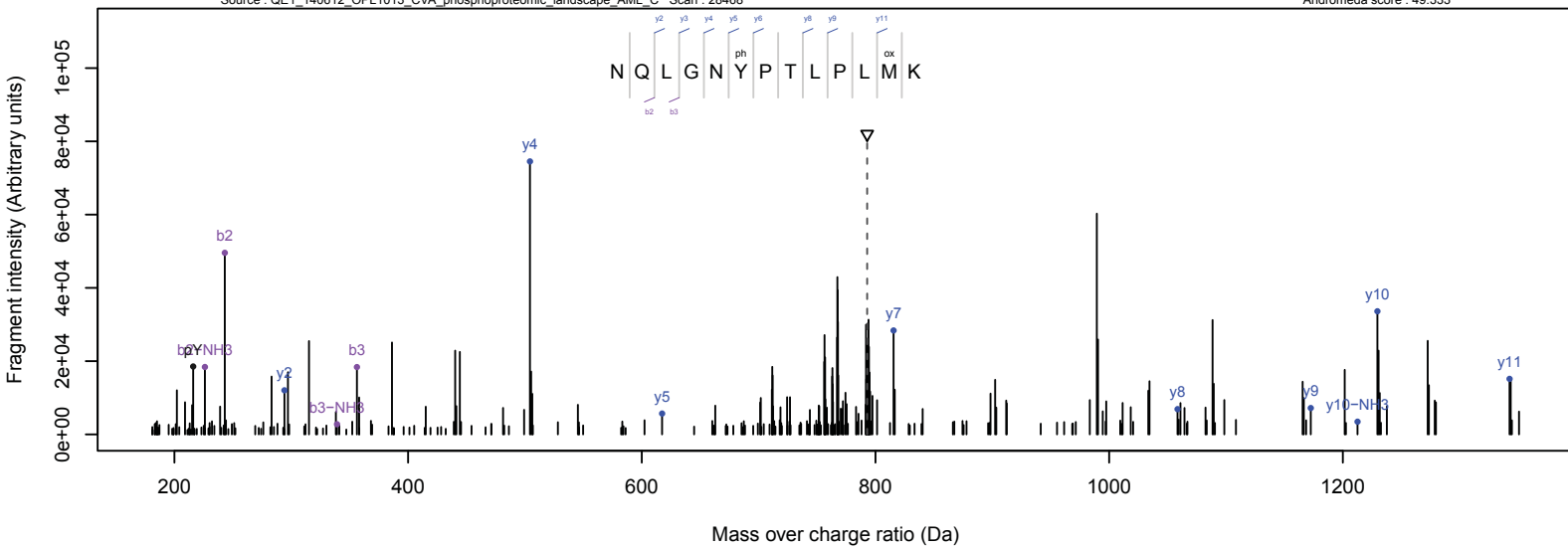
Andromeda score : 49.333



## NQLGNY(ph)PTLPLM(ox)K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28468

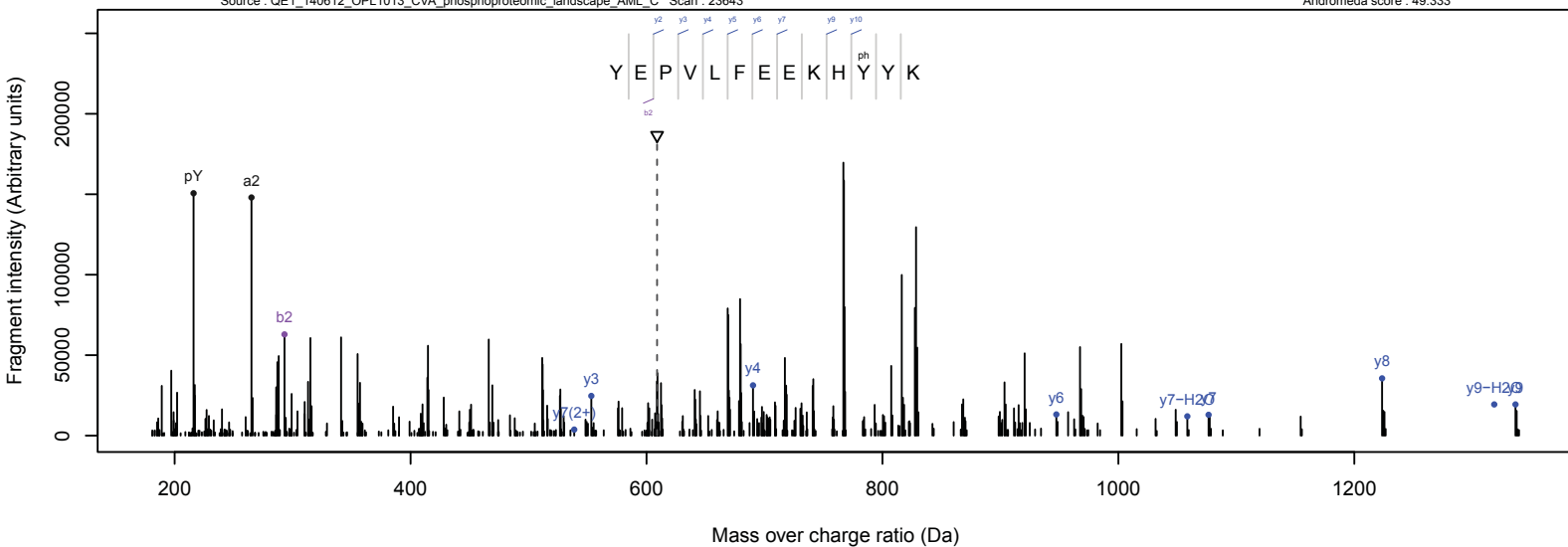
Andromeda score : 49.333



## YEPVLFEEKHYK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23643

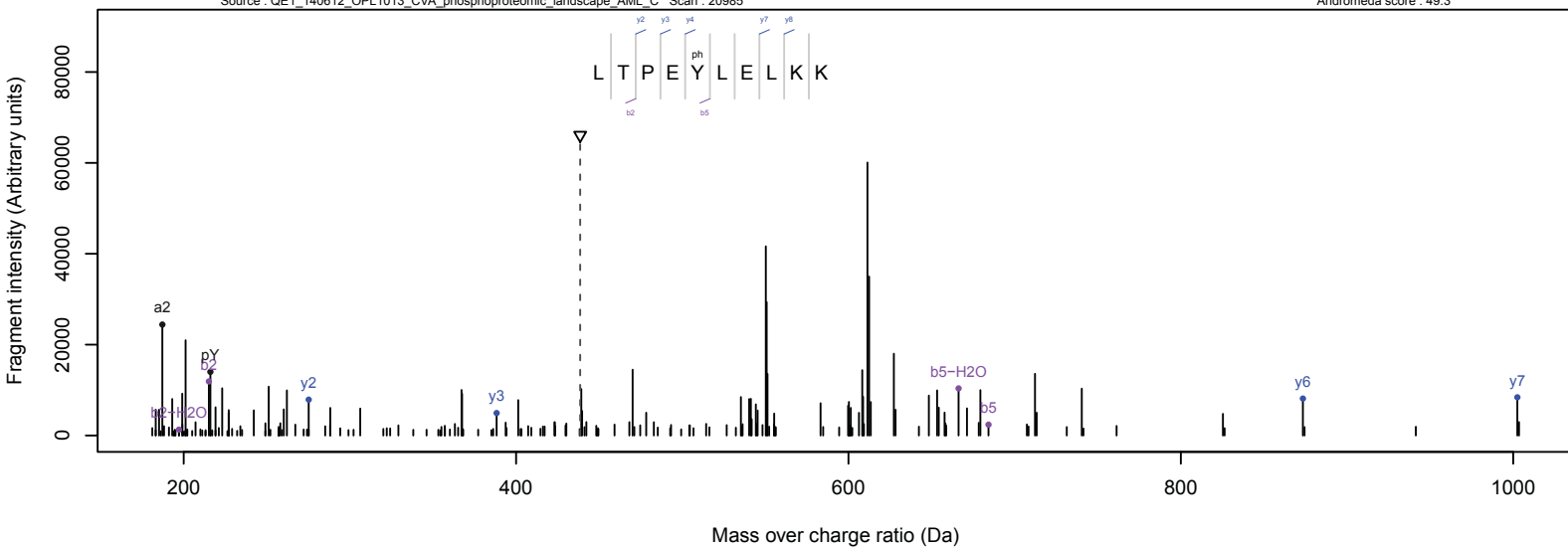
Andromeda score: 49.333



## LTPEYLELKK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 20985

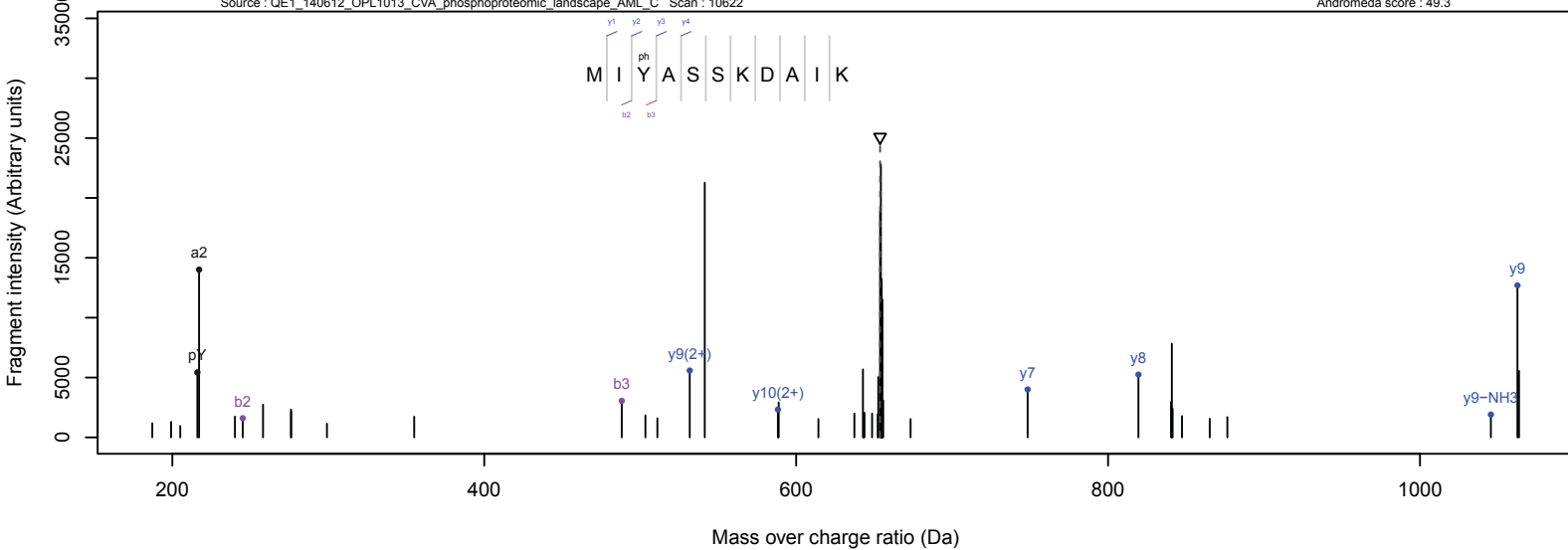
Andromeda score: 49.3



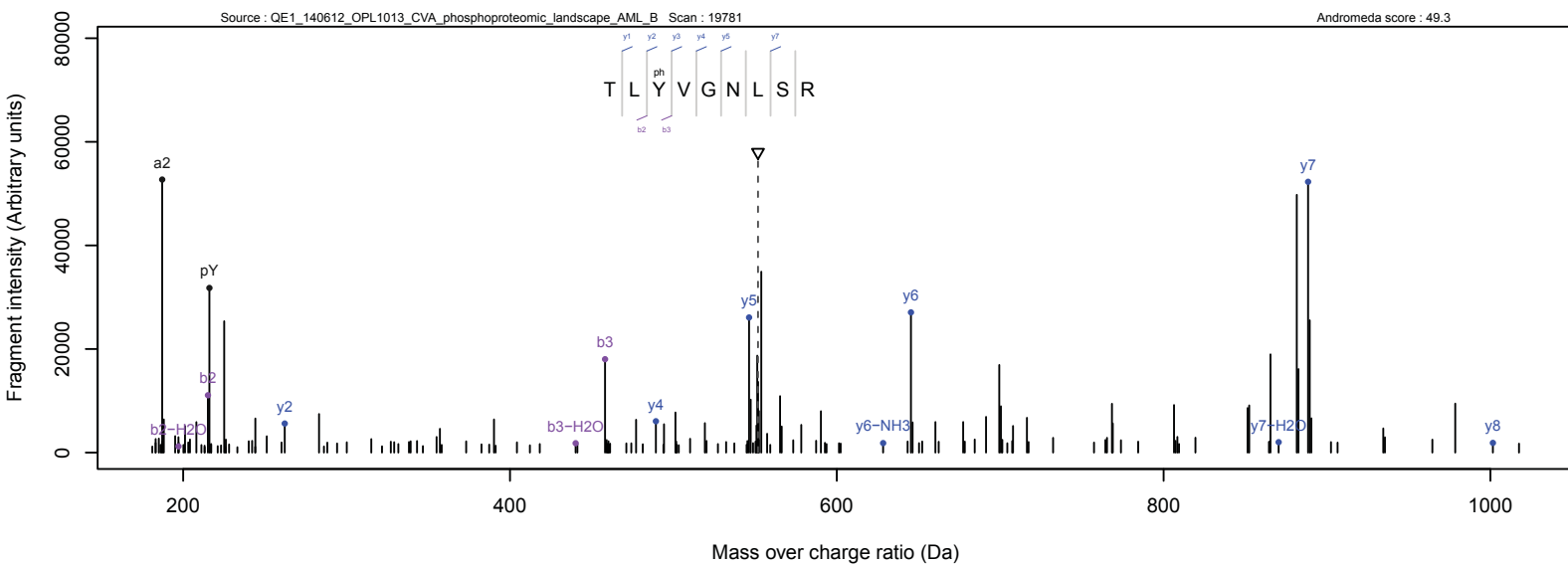
## MIY(ph)ASSKDAIK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10622

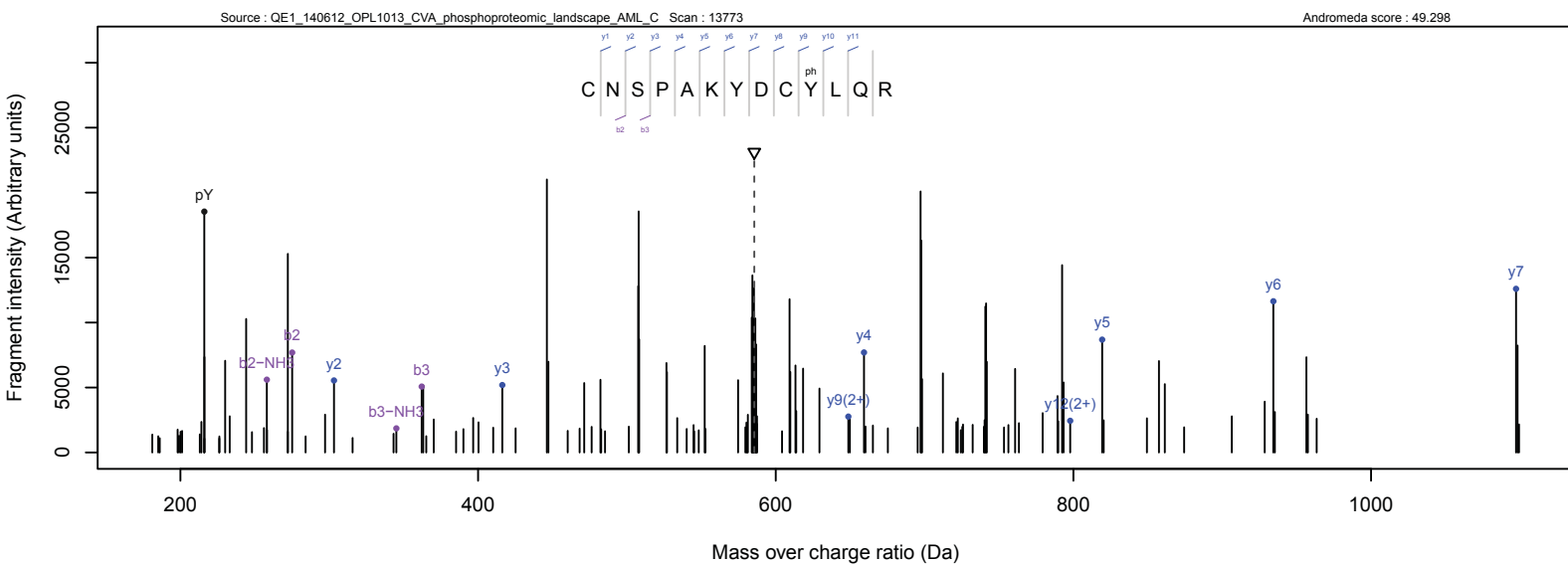
Andromeda score: 49.3



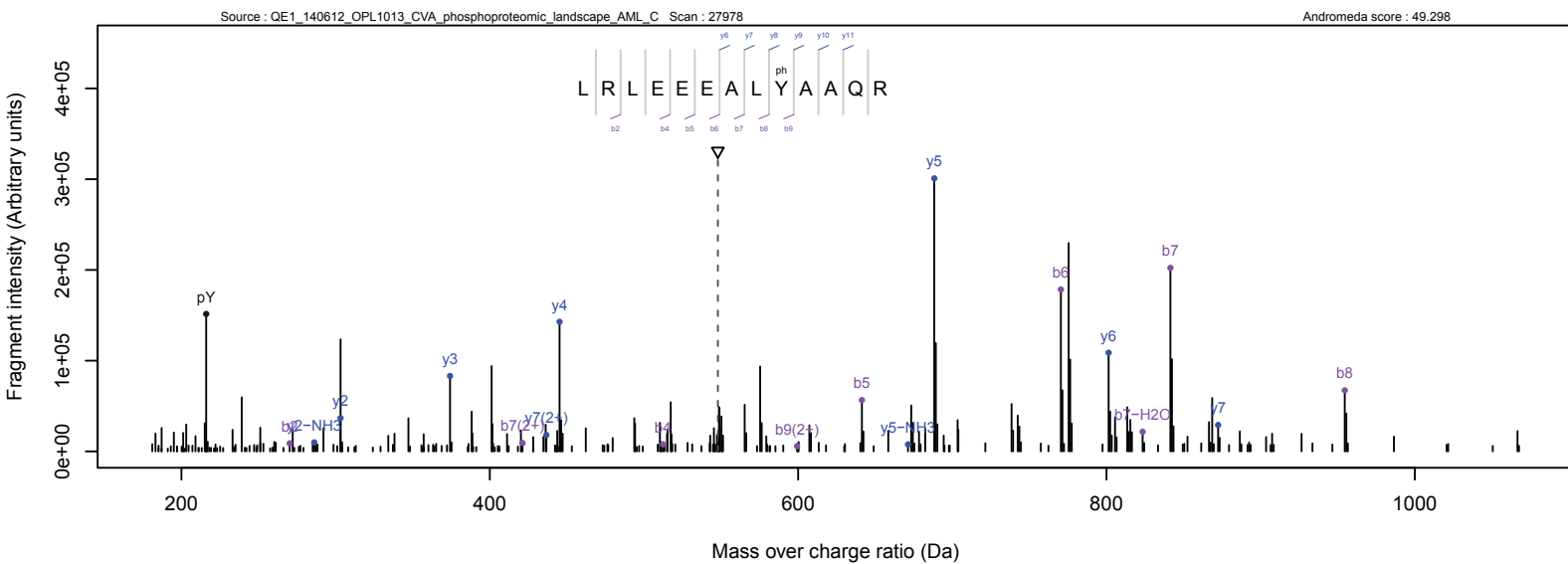
TLY(ph)VGNLSR-2+

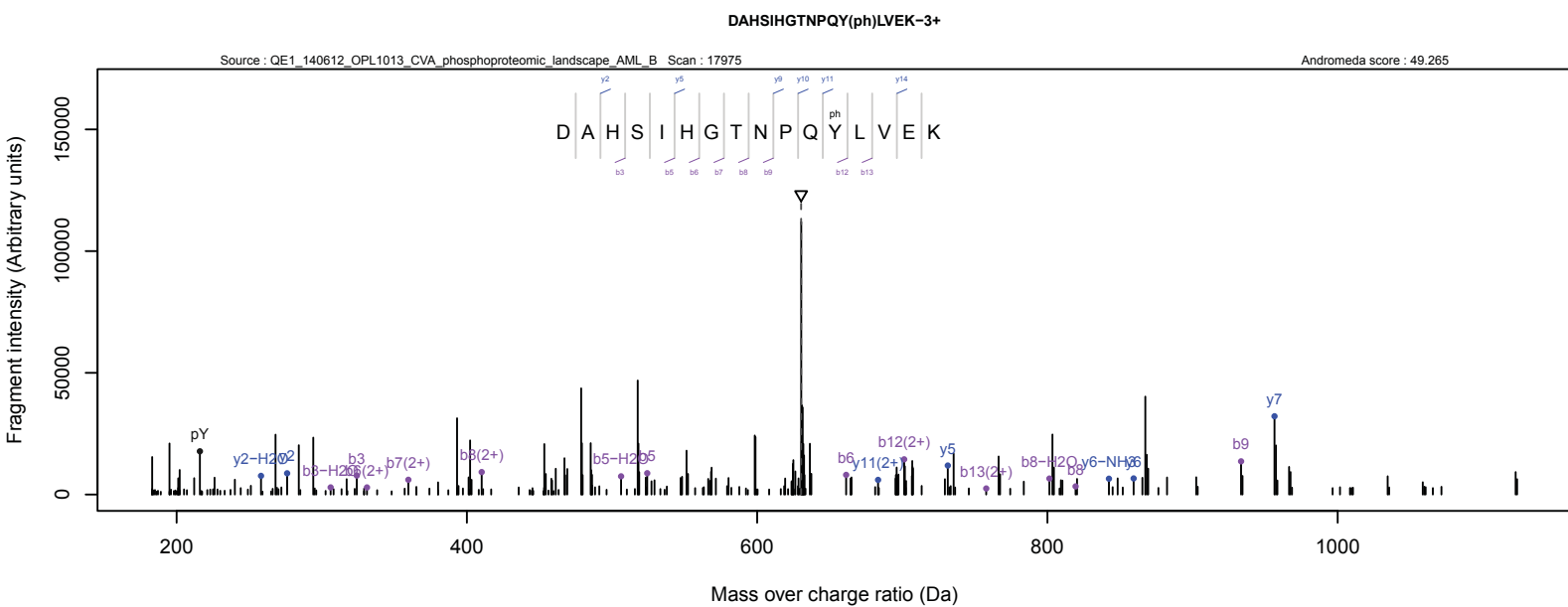
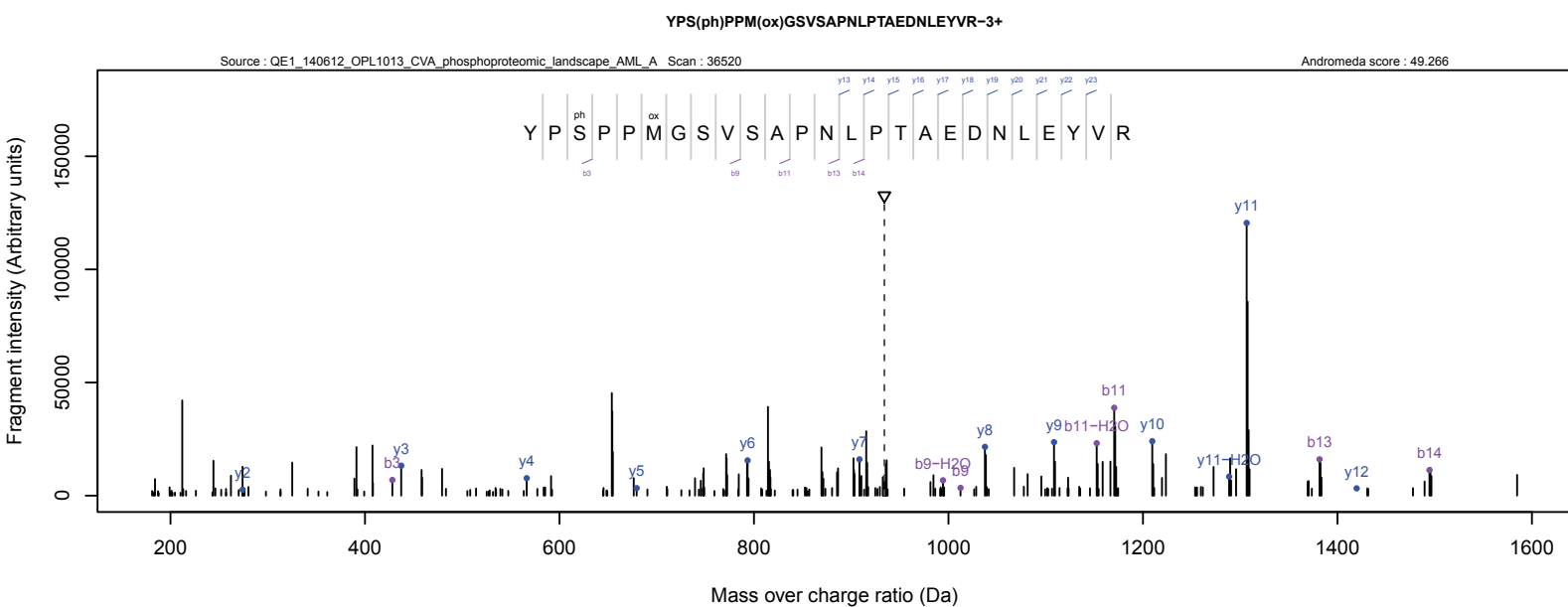
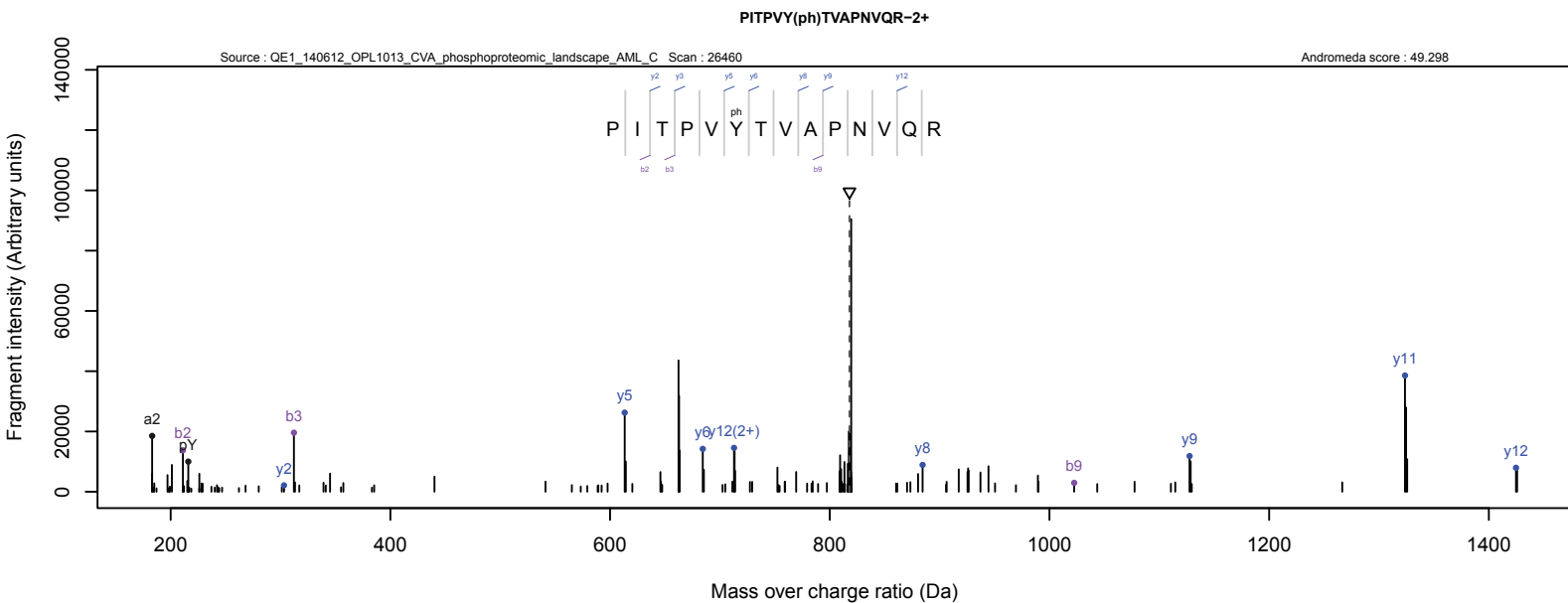


CNSPAKYDCY(ph)LQR-3+



LRLEEEALY(ph)AAQR-3+



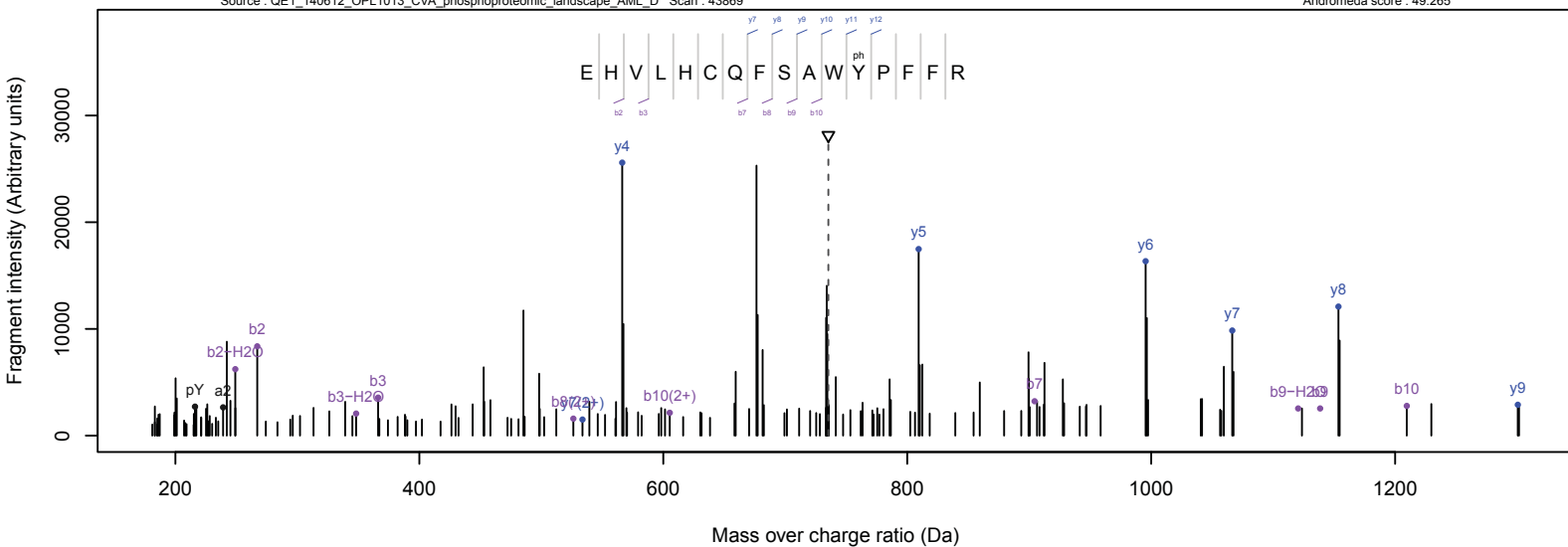




**EHVLCQFSAWY(ph)PFFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 43889

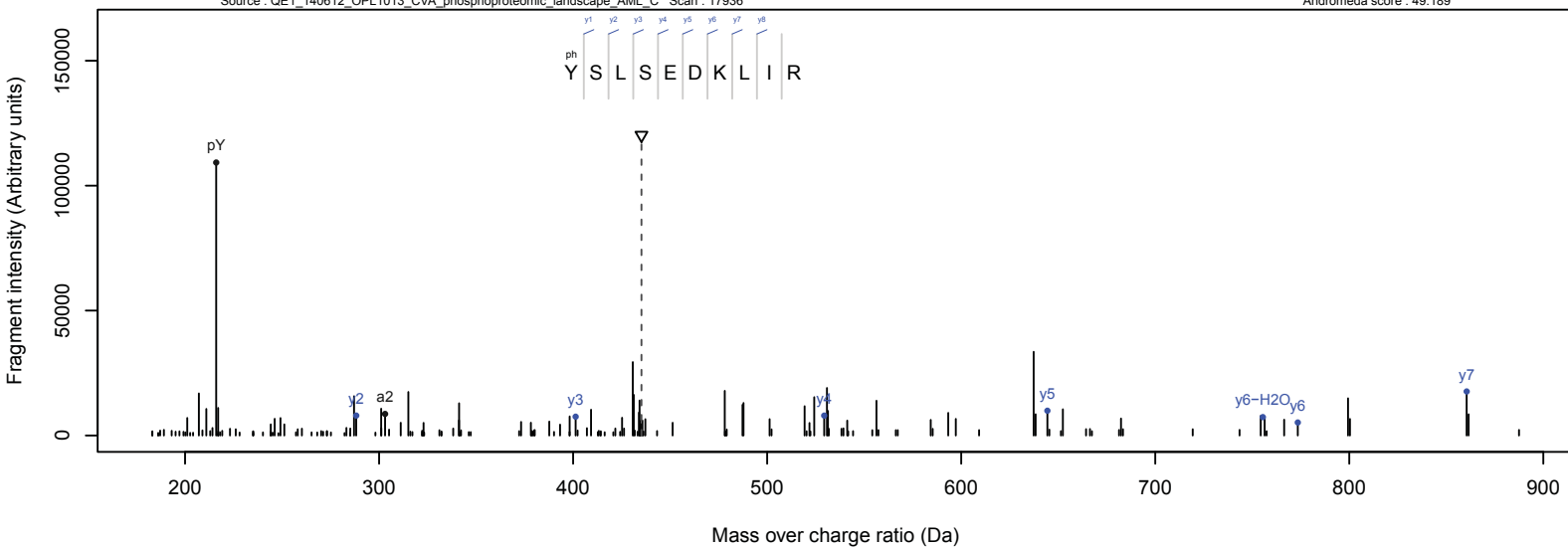
Andromeda score : 49.265



**Y(ph)SLSEDKLIR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17936

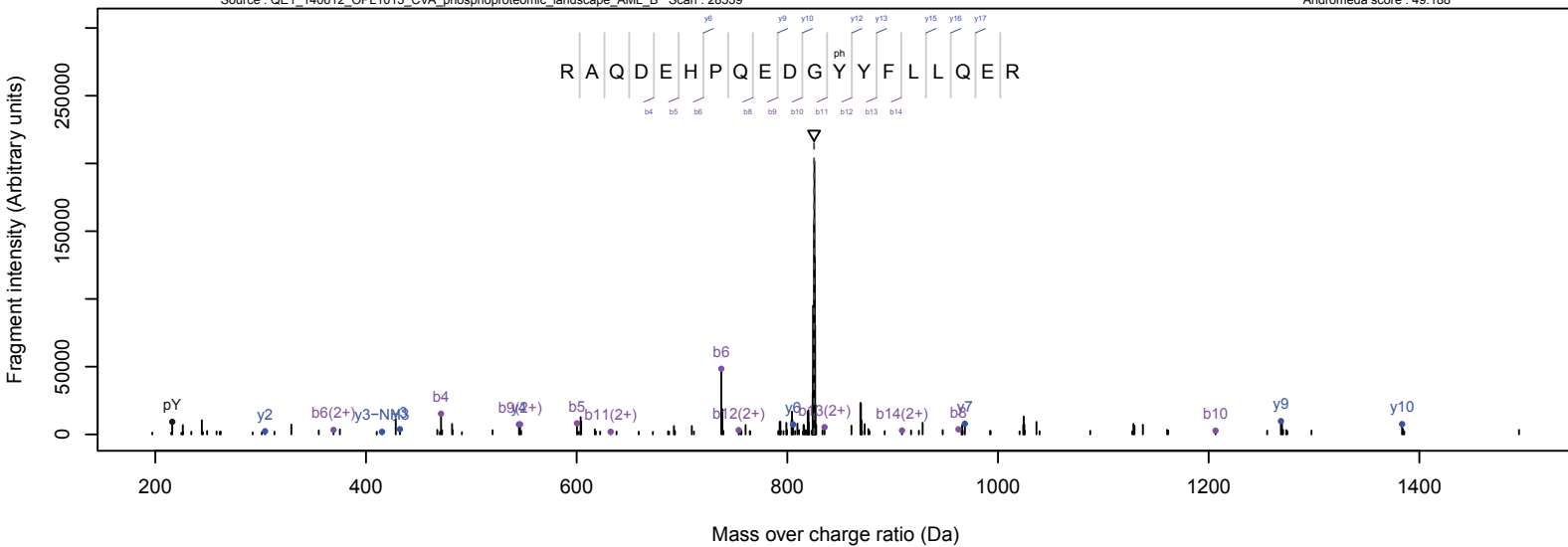
Andromeda score : 49.189



**RAQDEHPQEDGY(ph)YLLQER-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28539

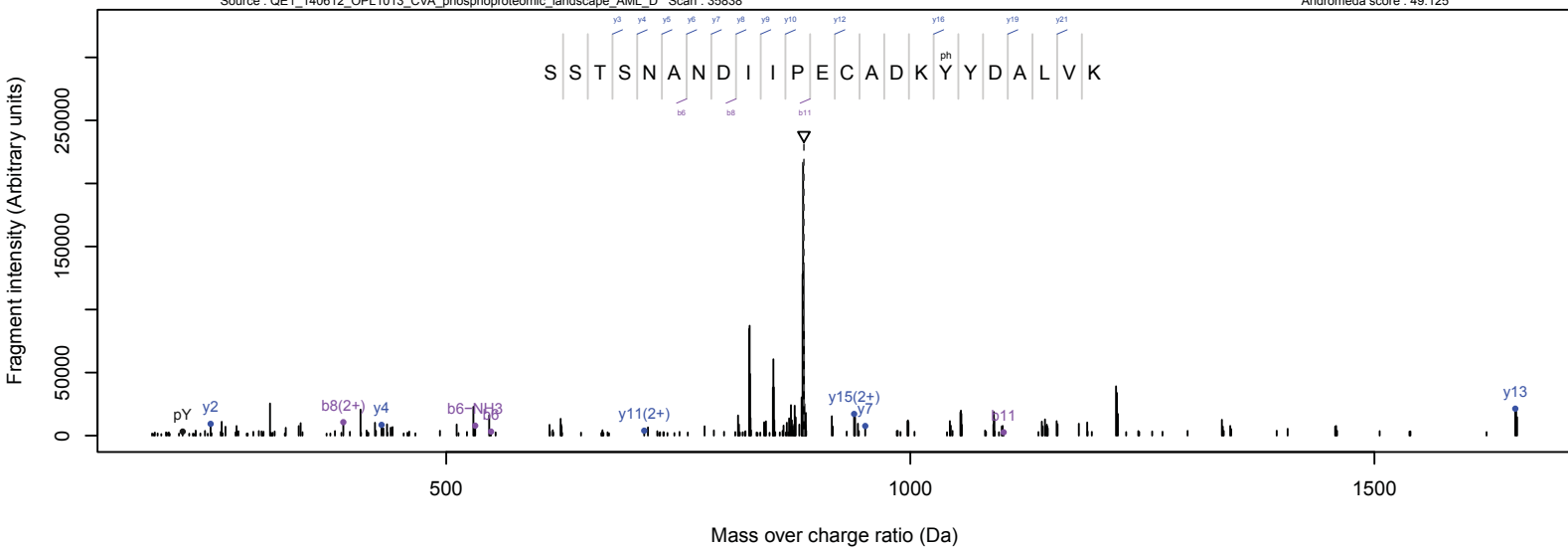
Andromeda score : 49.188



## SSTSNANDIPECADKY(ph)YDALVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 35838

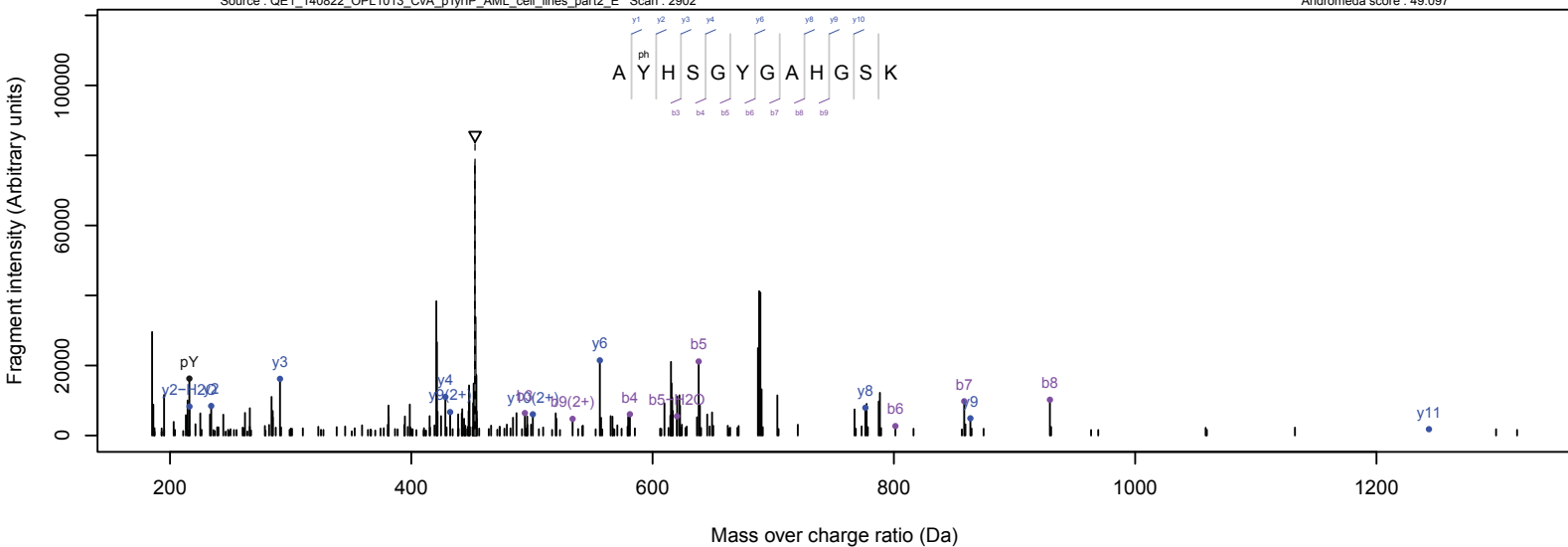
Andromeda score : 49.125



## (ac)AY(ph)HSGYGAHGSK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 2902

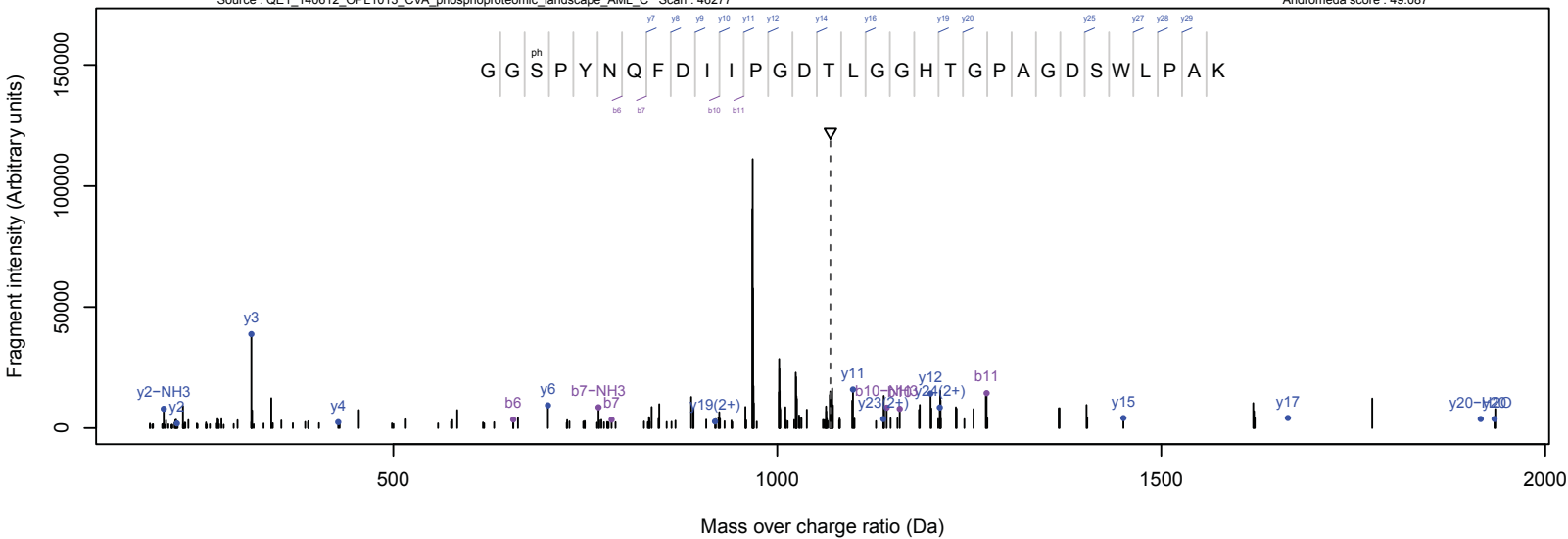
Andromeda score : 49.097



## GGS(ph)PYNQFDIIPGDTLGGHTGPAGDSWLPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46277

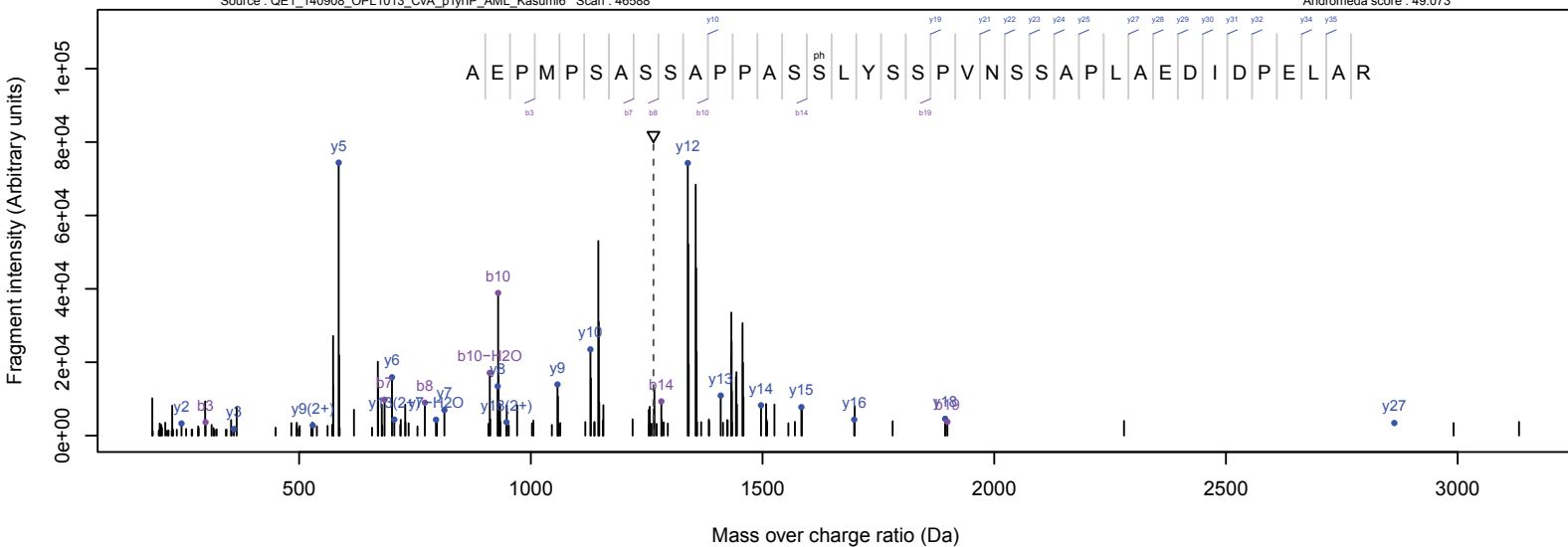
Andromeda score : 49.087



**AEPMPASSAPPASS(ph)LYSSPVNSSAPLAEDIDPELAR-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIPL\_AML\_Kasumi6 Scan : 46588

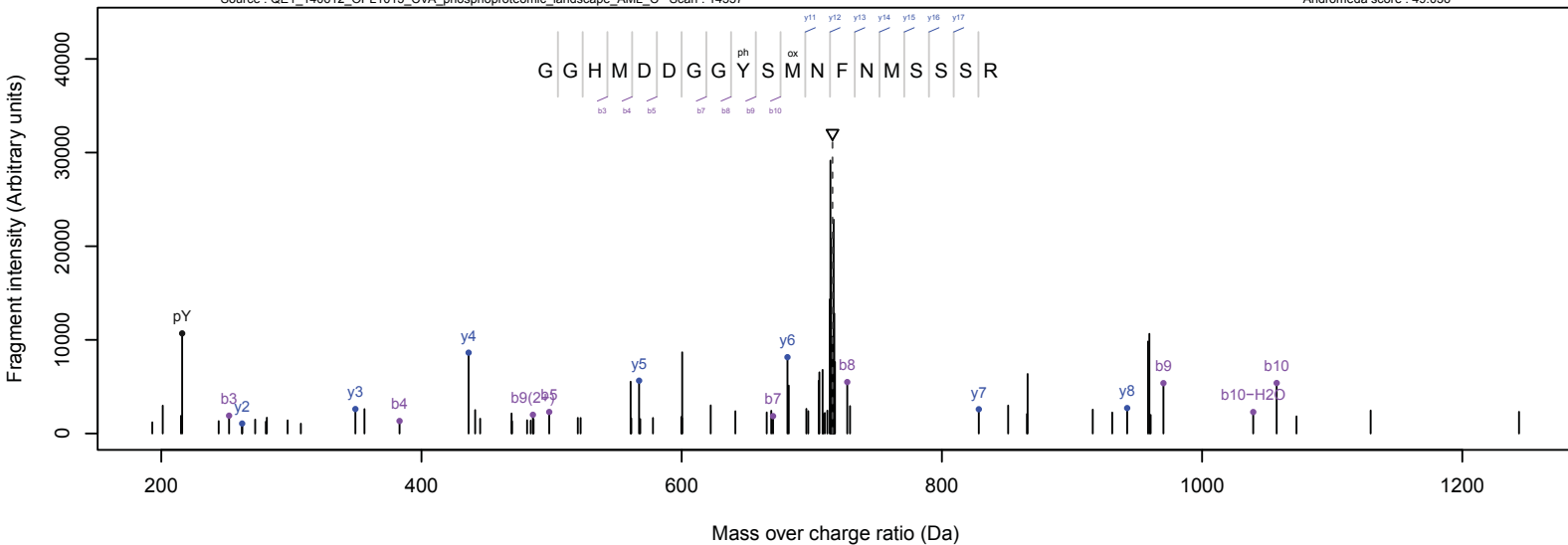
Andromeda score : 49.073



**GGHMDDGGY(ph)SM(ox)NFMSSSR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14557

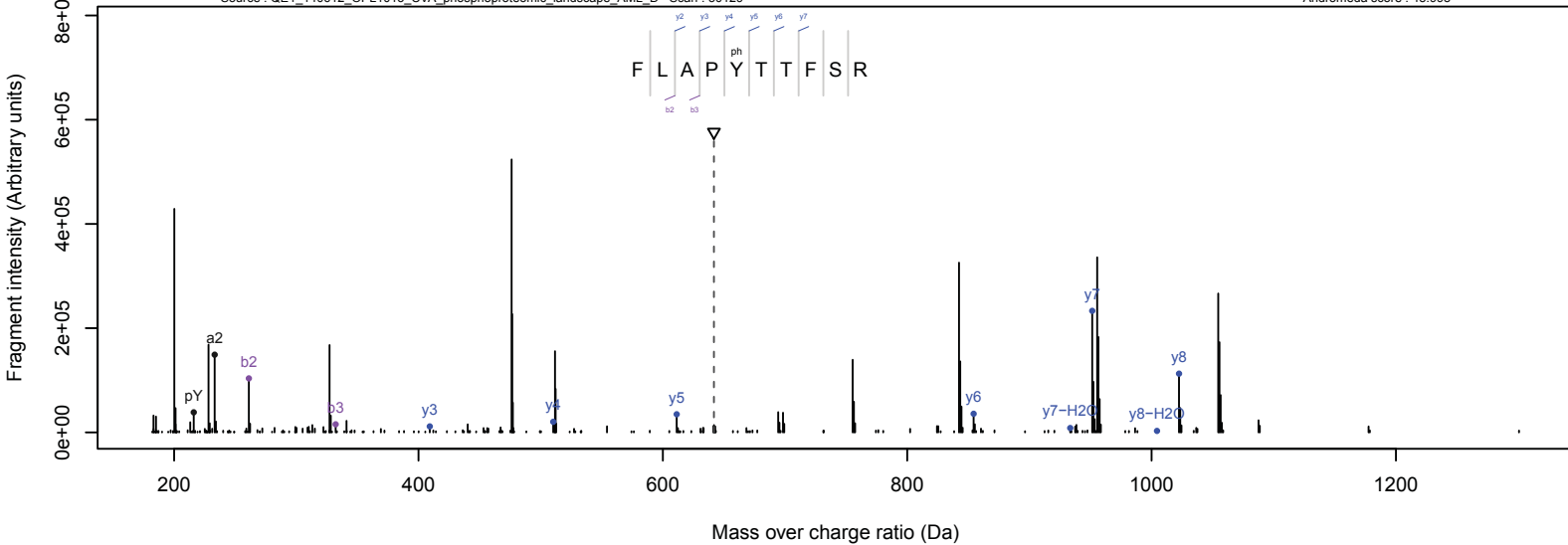
Andromeda score : 49.036



**FLAPY(ph)TTFSR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 30129

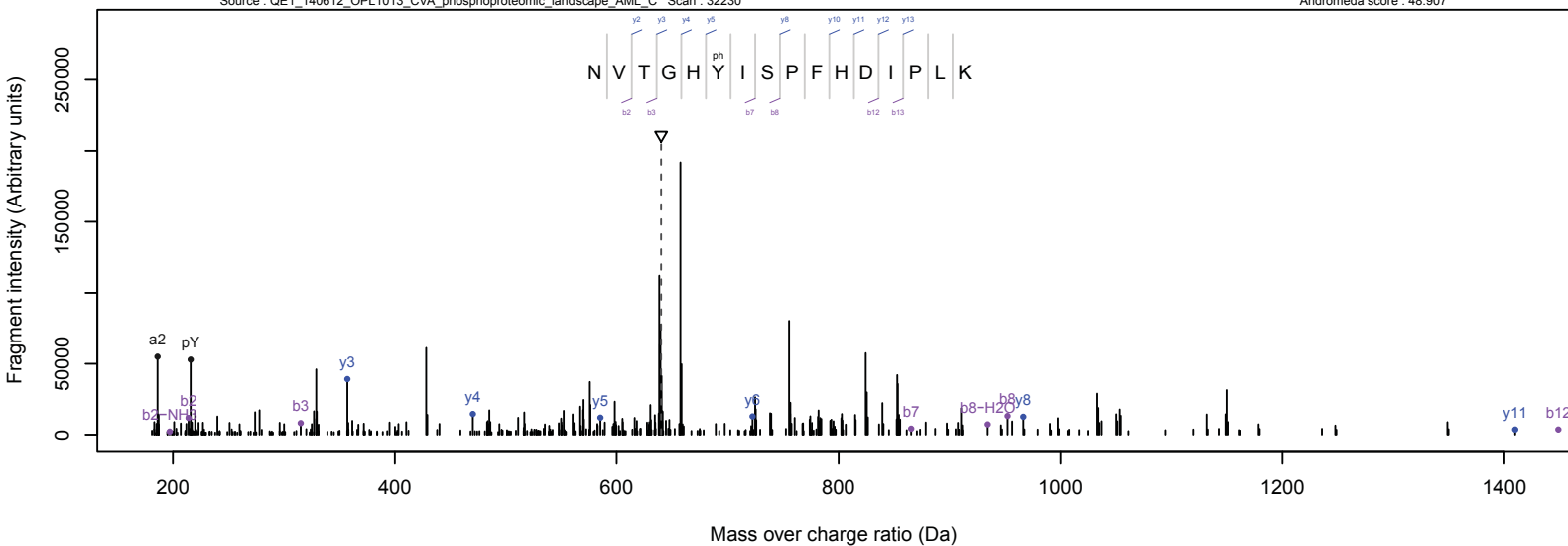
Andromeda score : 48.998



## NVTGHY(ph)ISPFHDIPLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32230

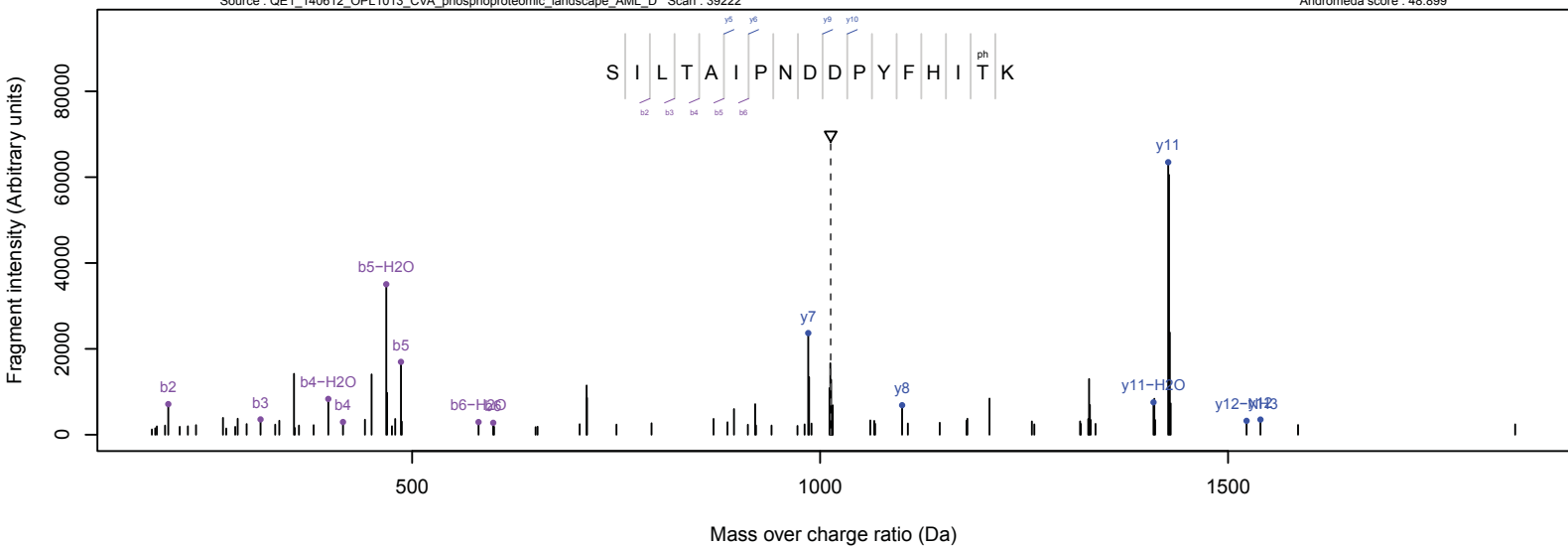
Andromeda score : 48.907



## SILTAIPNDPYPFHITK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 39222

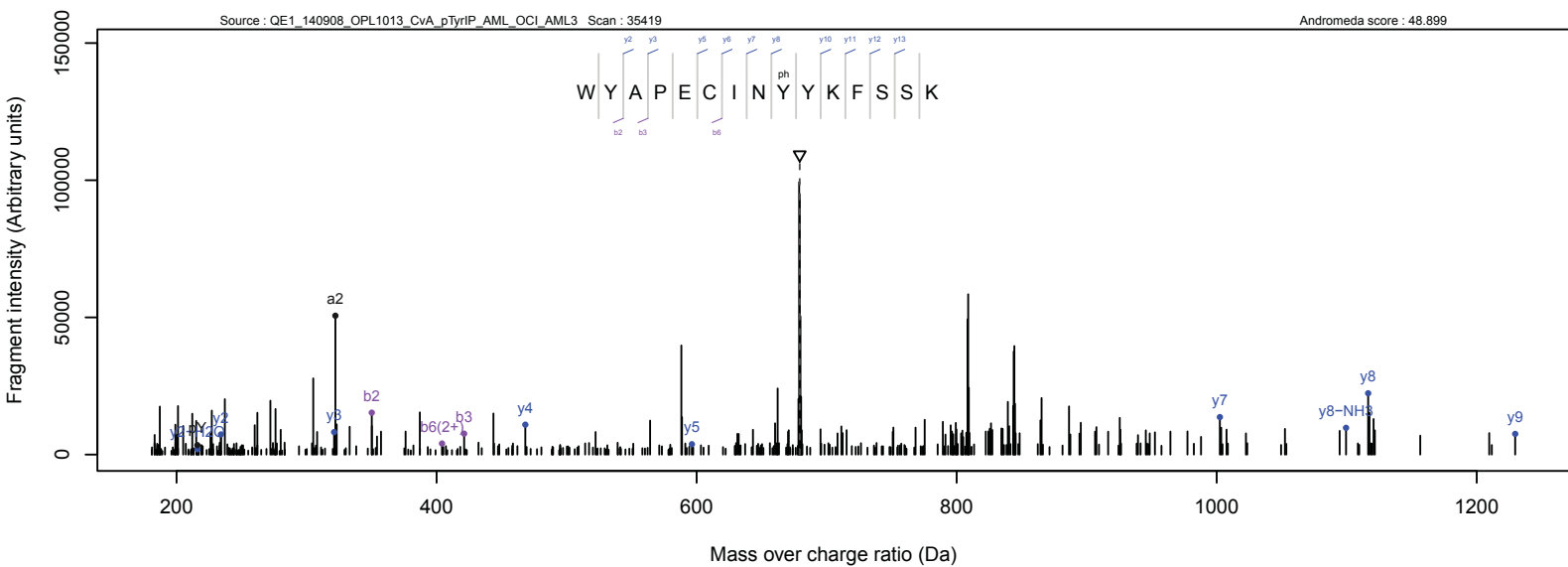
Andromeda score : 48.899



## WYAPECINYKFSK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 35419

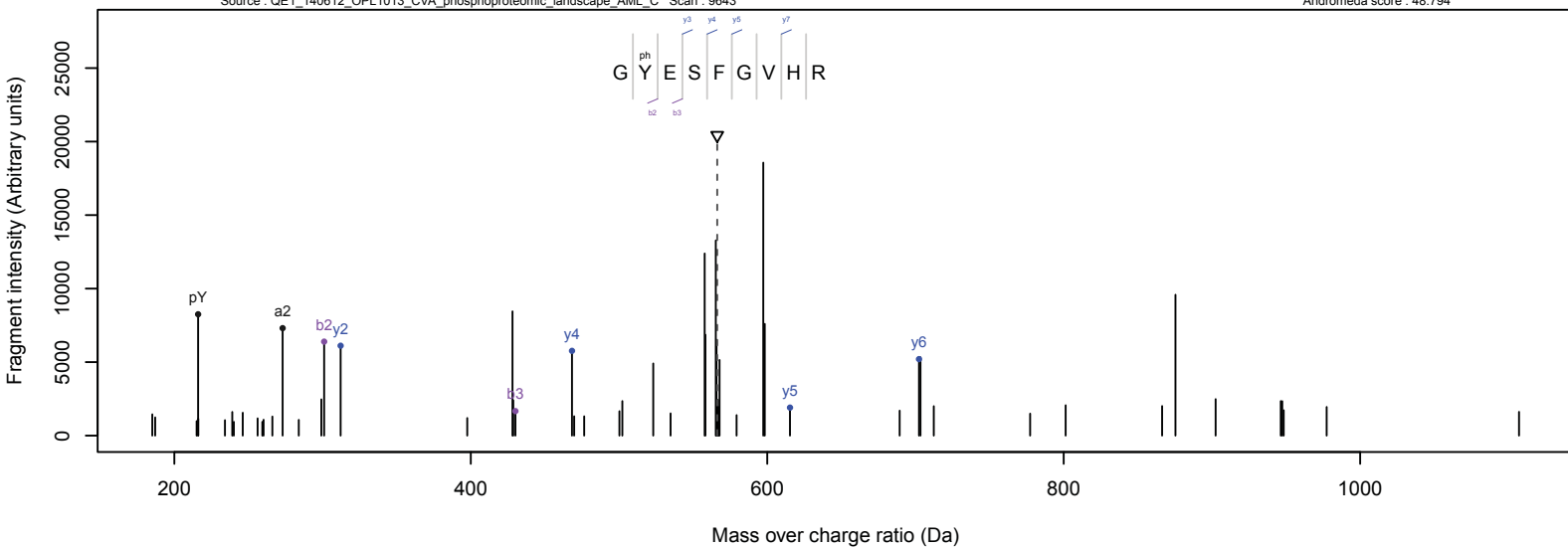
Andromeda score : 48.899



## GY(ph)ESFGVHR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9643

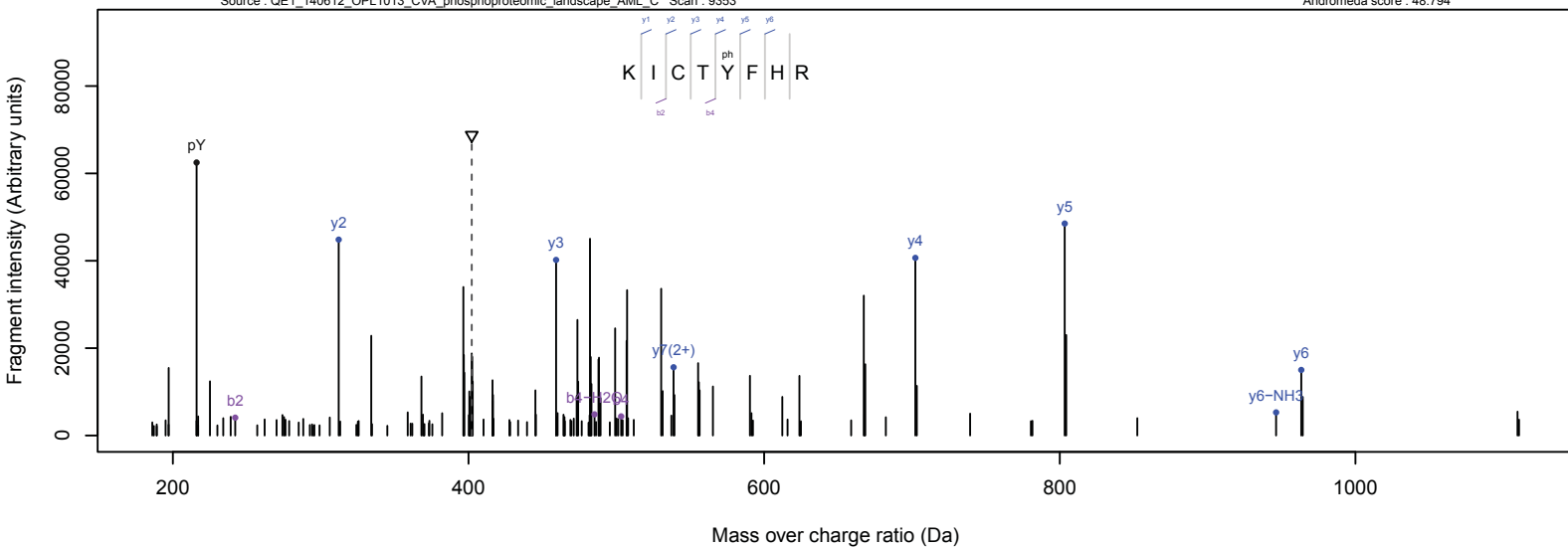
Andromeda score : 48.794



## KICTY(ph)FHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9353

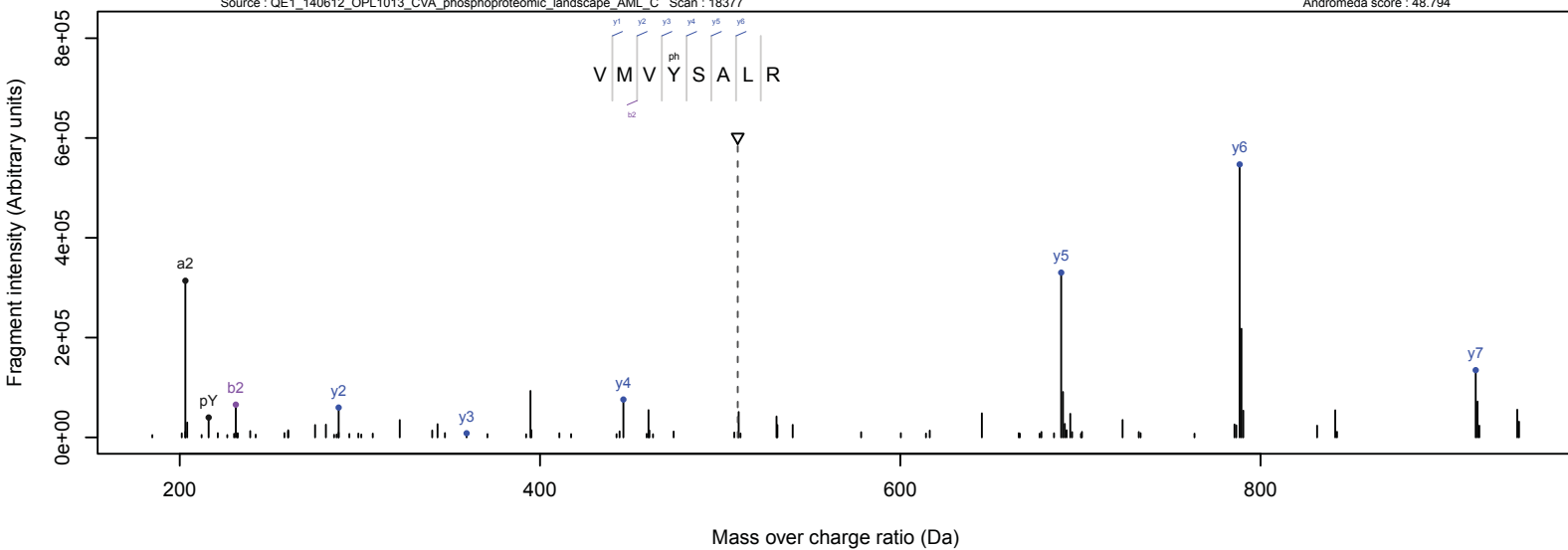
Andromeda score : 48.794



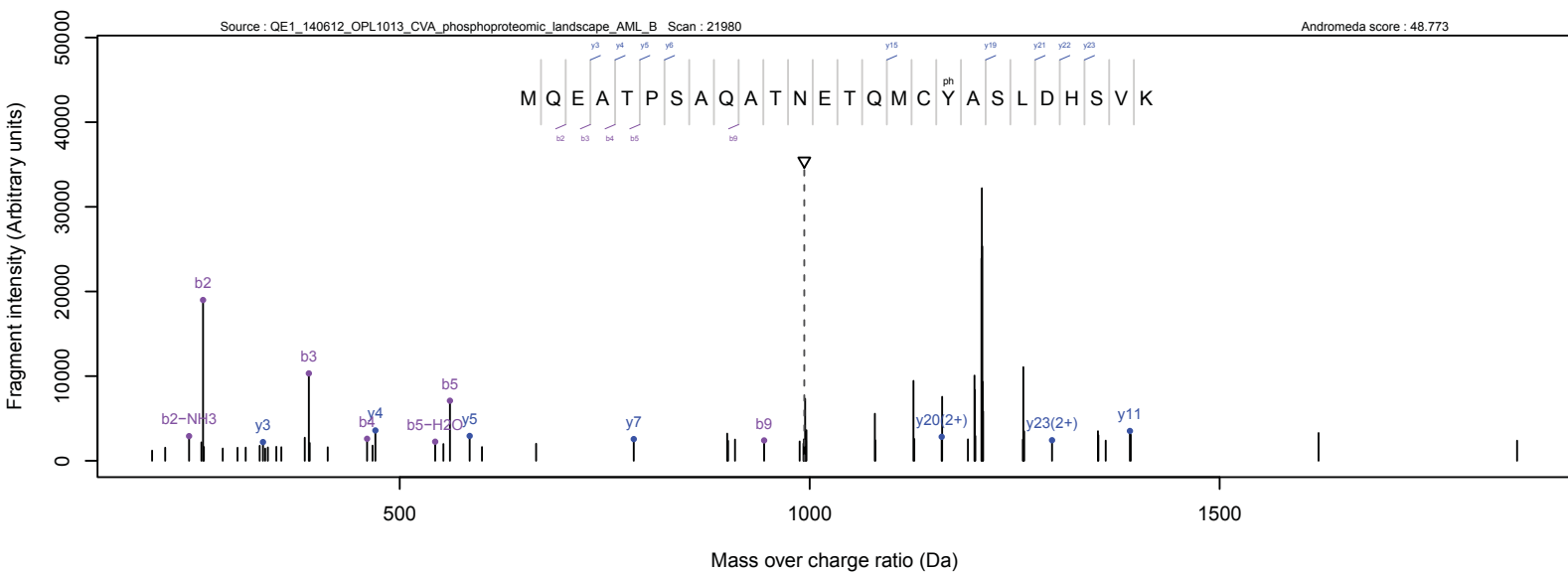
## VMVY(ph)SALR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18377

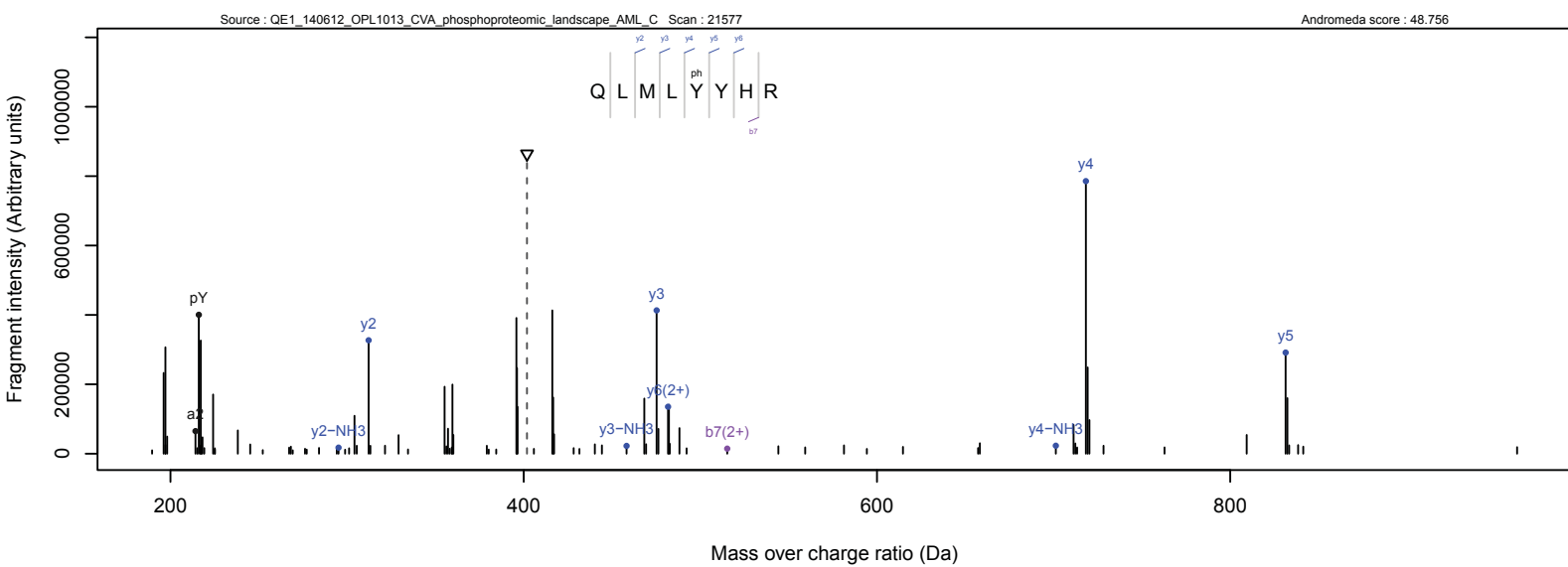
Andromeda score : 48.794



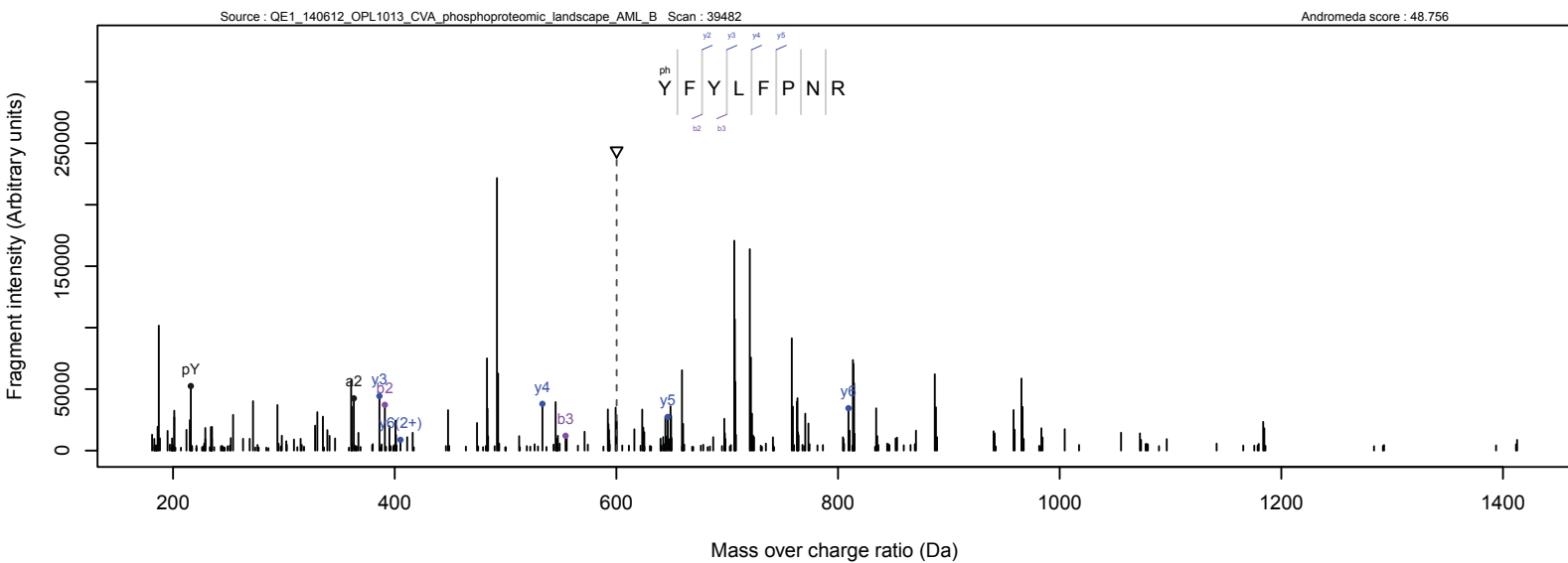
MQEATPSAQATNETQMCY<sup>ph</sup>ASLDH<sup>ph</sup>SVK-3+

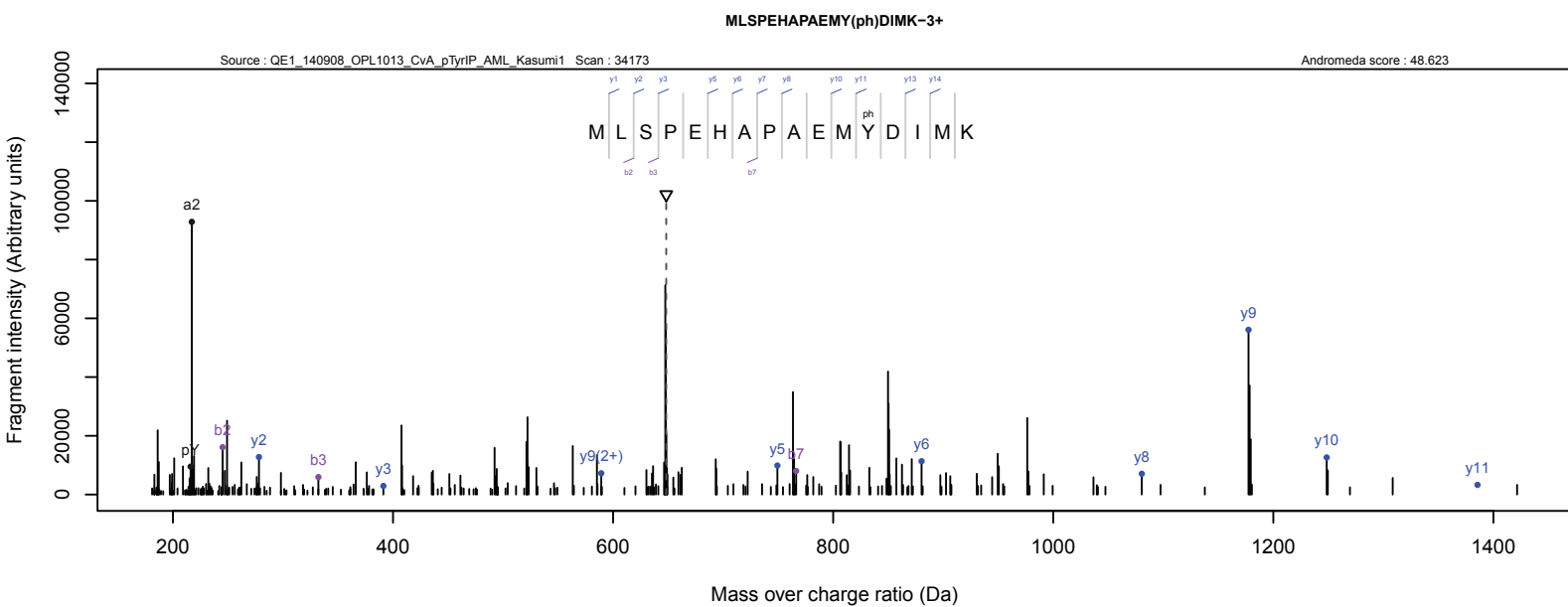
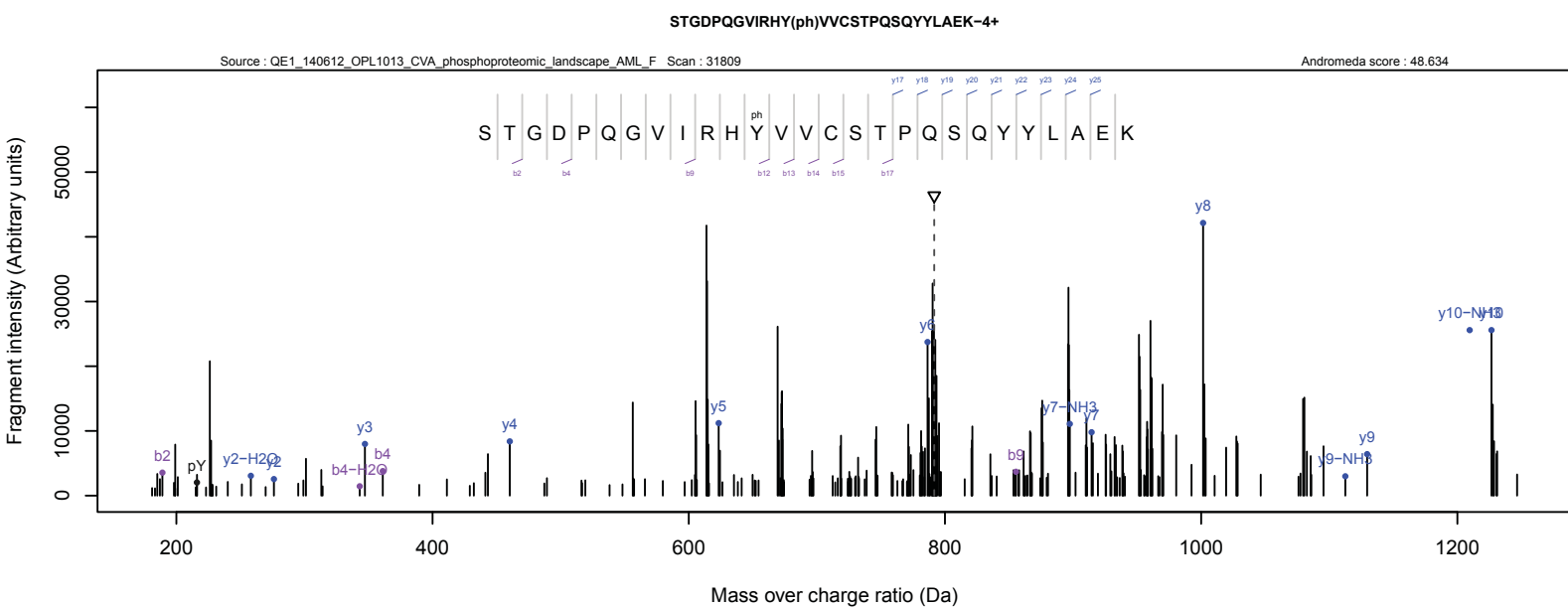
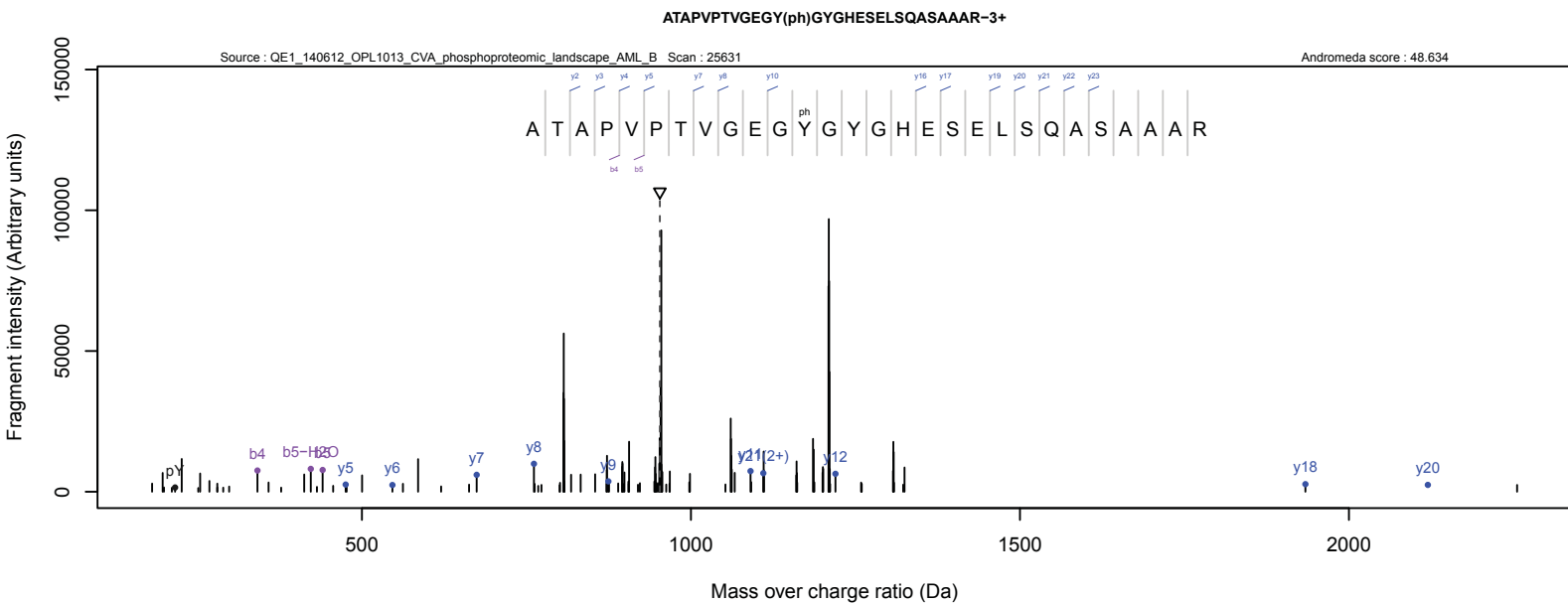


QLMLY<sup>ph</sup>YHR-3+

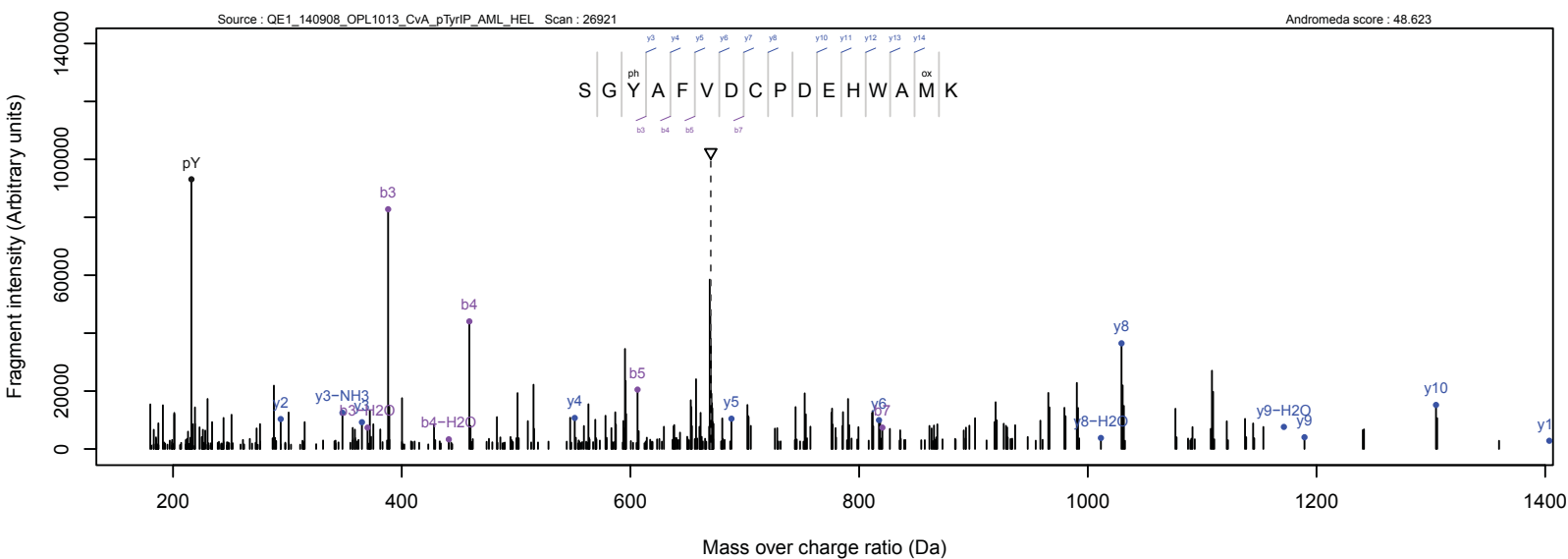


Y<sup>ph</sup>FYLF<sup>ph</sup>PNR-2+

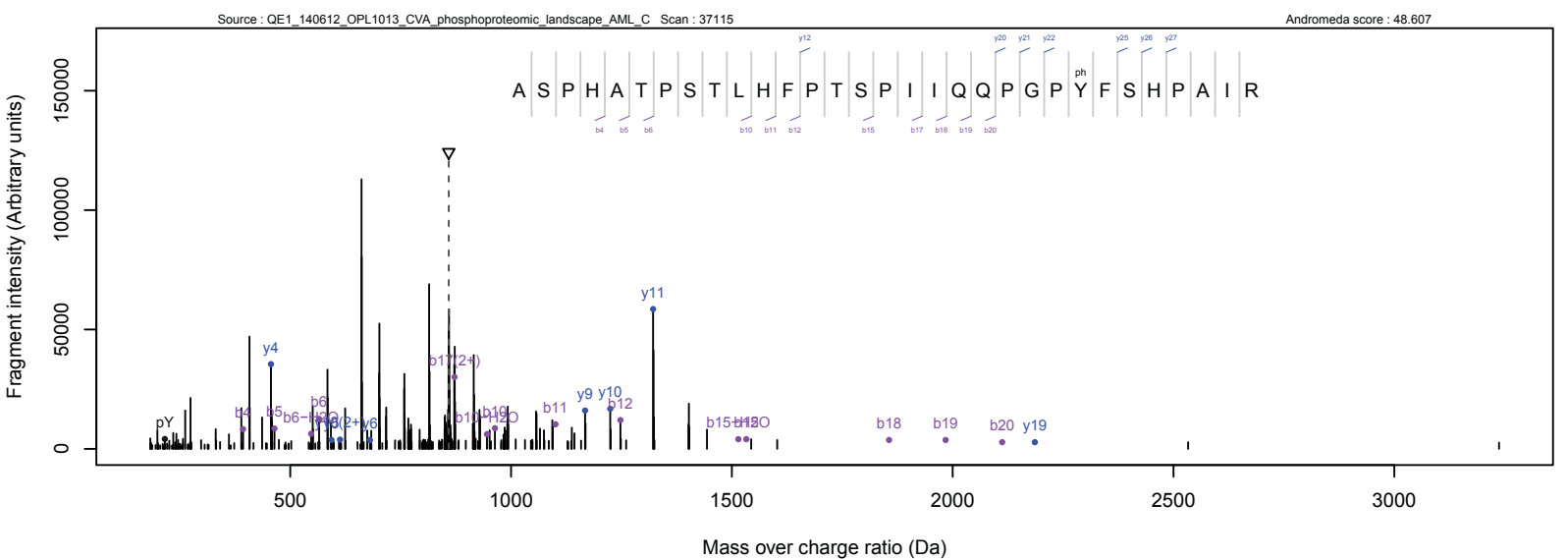




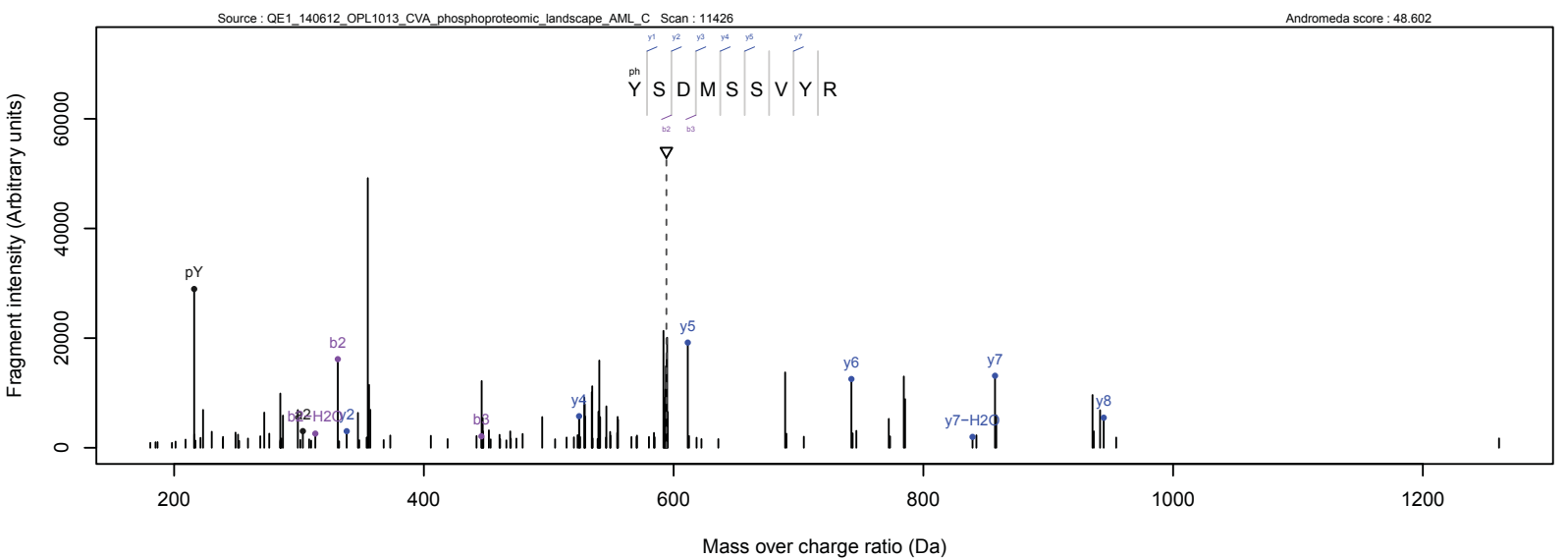
SGY(ph)AFVDCPDEHWAM(ox)K-3+



ASPHATPSTLHFTSPPIIQPGPY(ph)FSPHAIR-4+



Y(ph)SDMSSVYR-2+

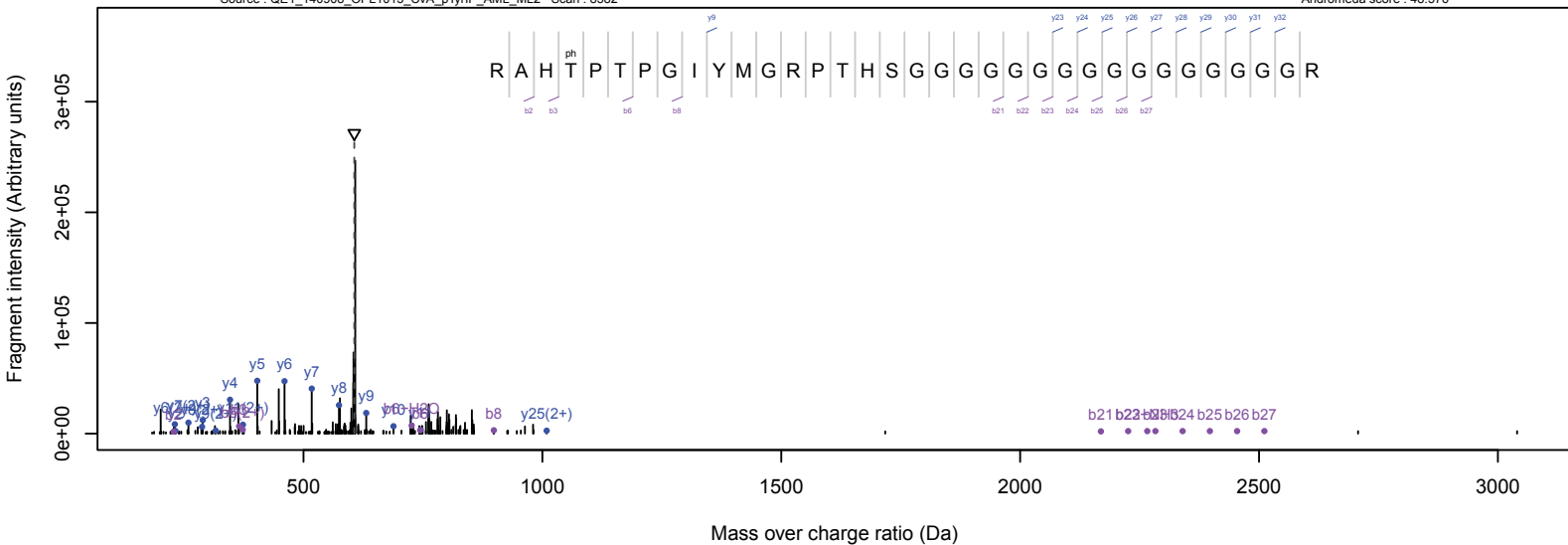




RAHT(ph)PTPGIYMGRRPTHTSGGGGGGGGGGGGR-5+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 8582

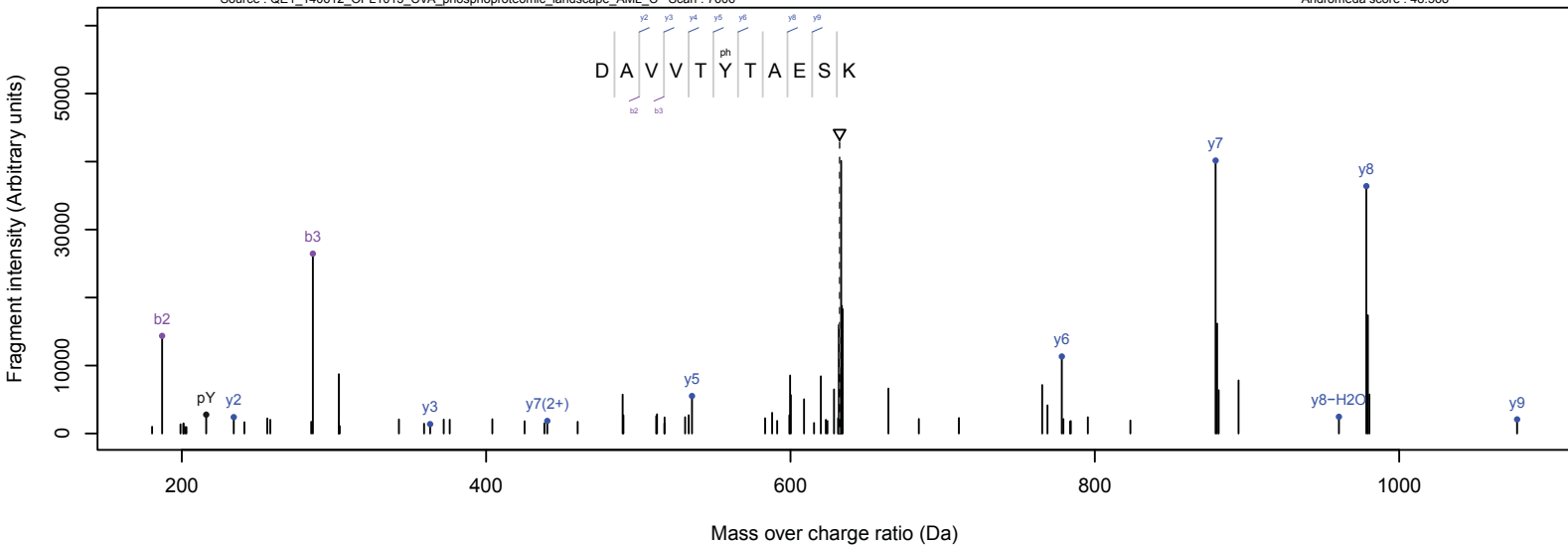
Andromeda score : 48.576



DAVVY(ph)TAESK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7606

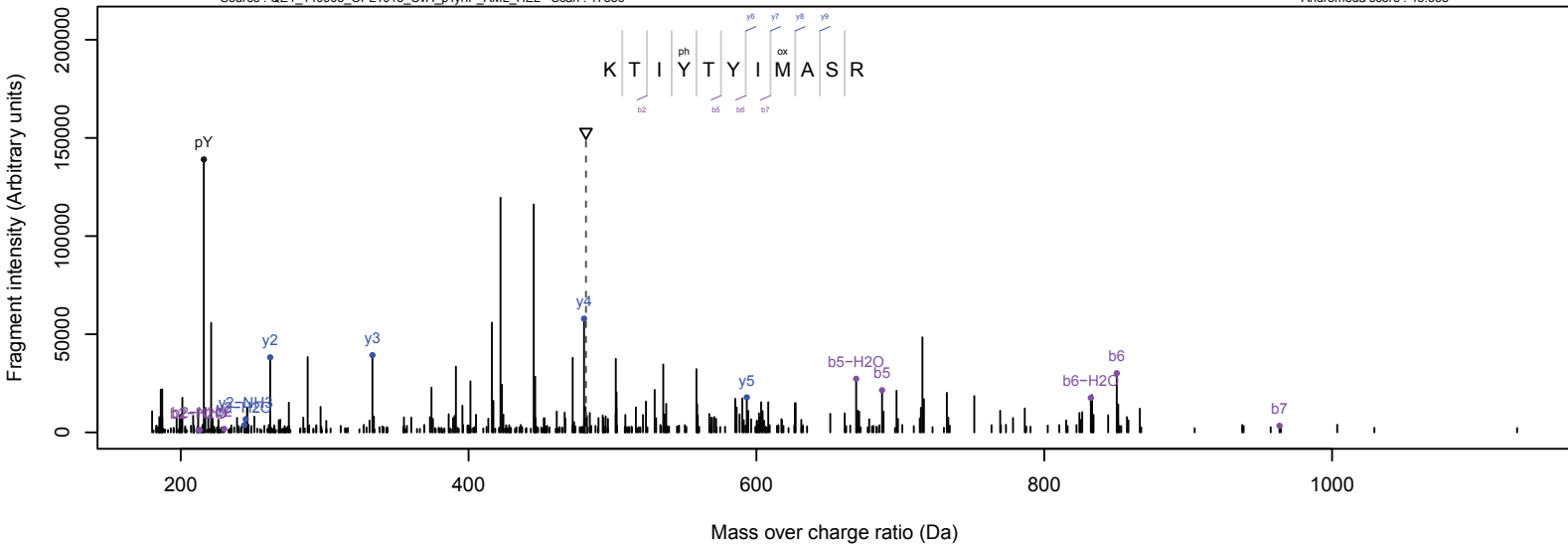
Andromeda score : 48.568



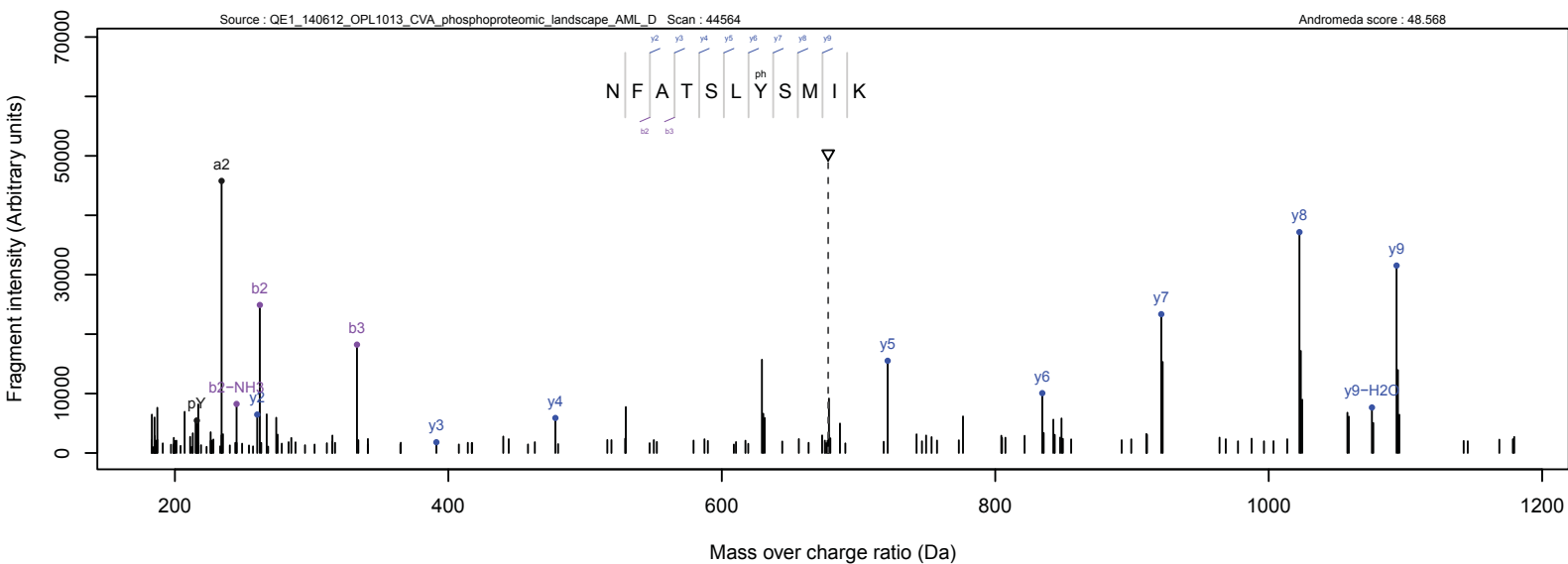
KTIY(ph)TYIM(ox)ASR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 17836

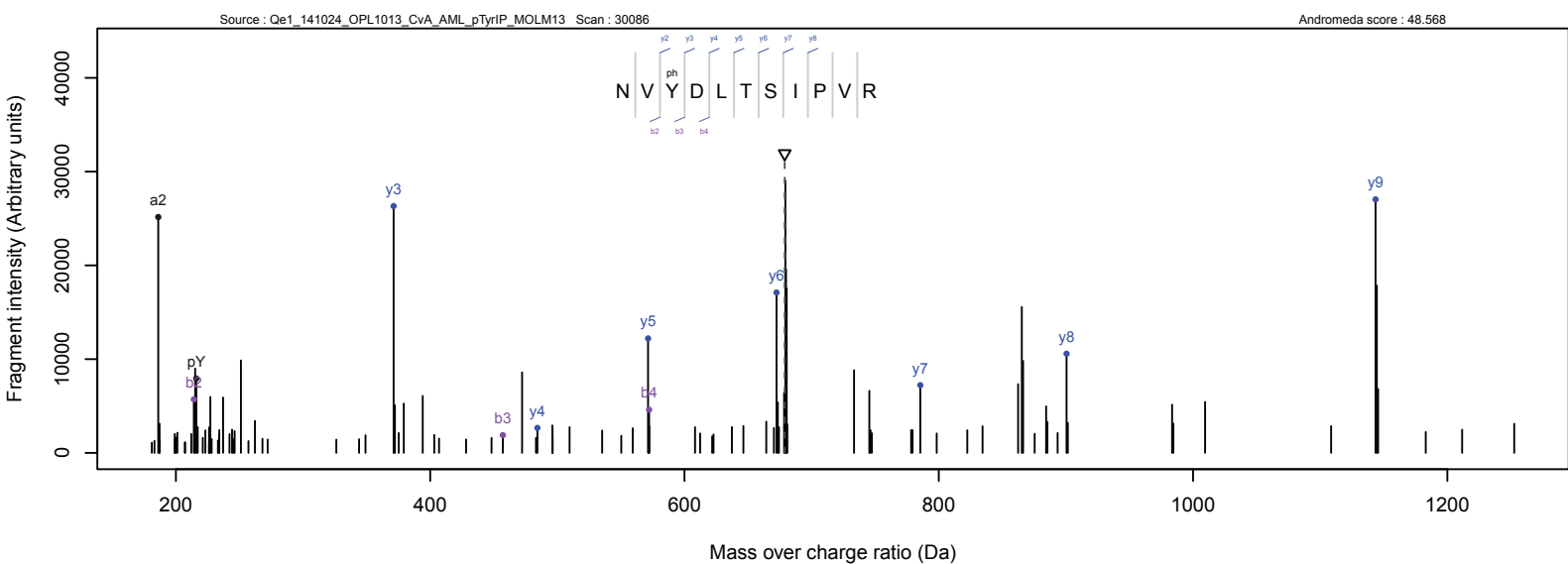
Andromeda score : 48.568



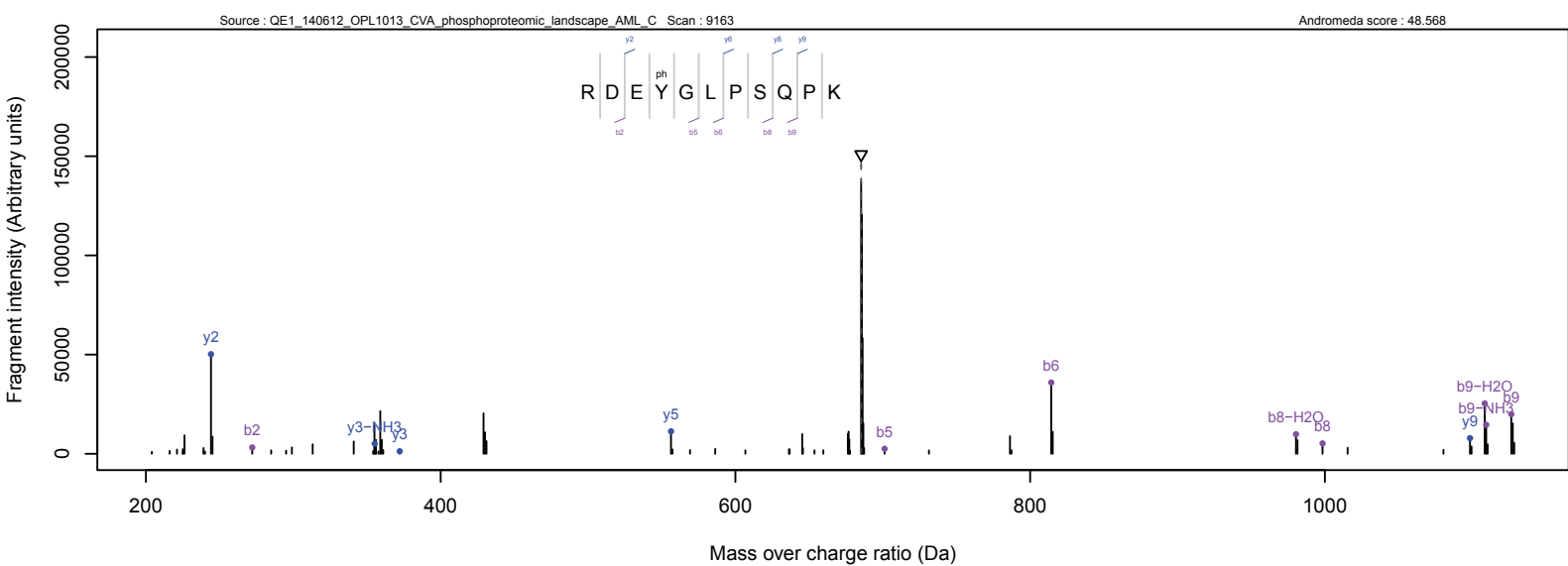
## NFATSLY(ph)SMIK-2+



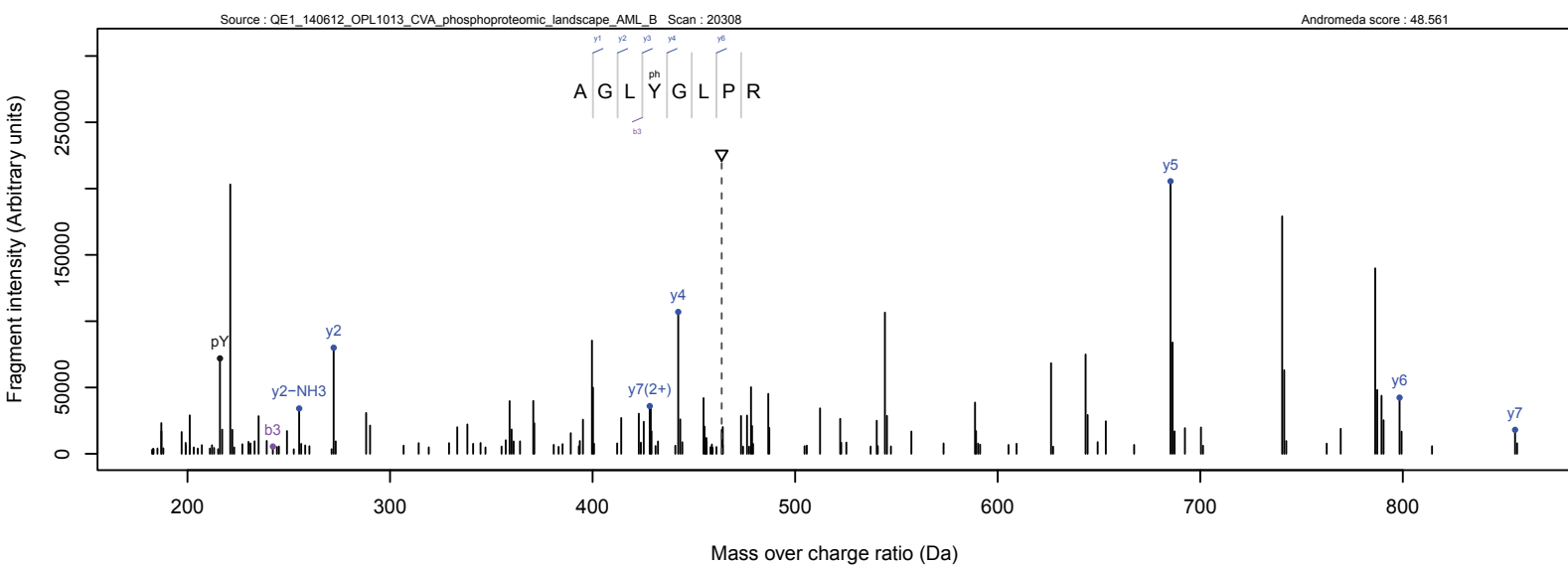
## NVY(ph)DLTSIPVR-2+



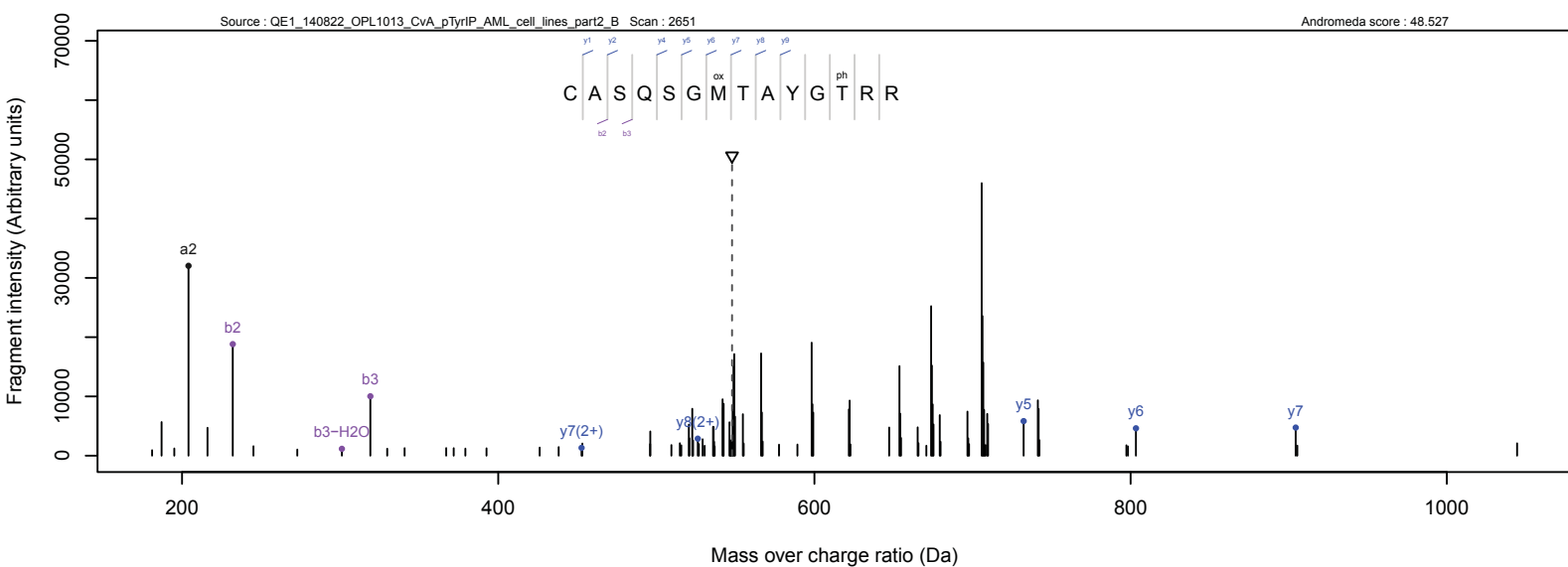
## RDEY(ph)GLPSQPK-2+



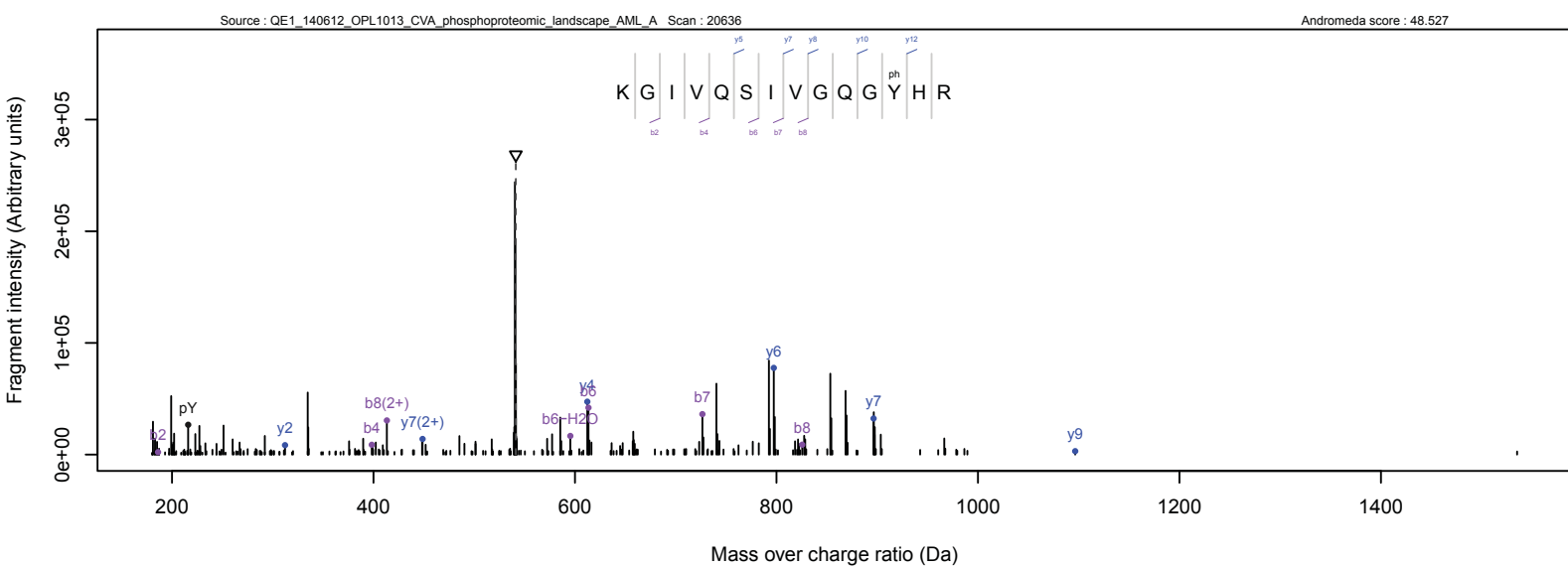
## AGLY(ph)GLPR-2+



## CASQSGM(ox)TAYG(ph)RR-3+



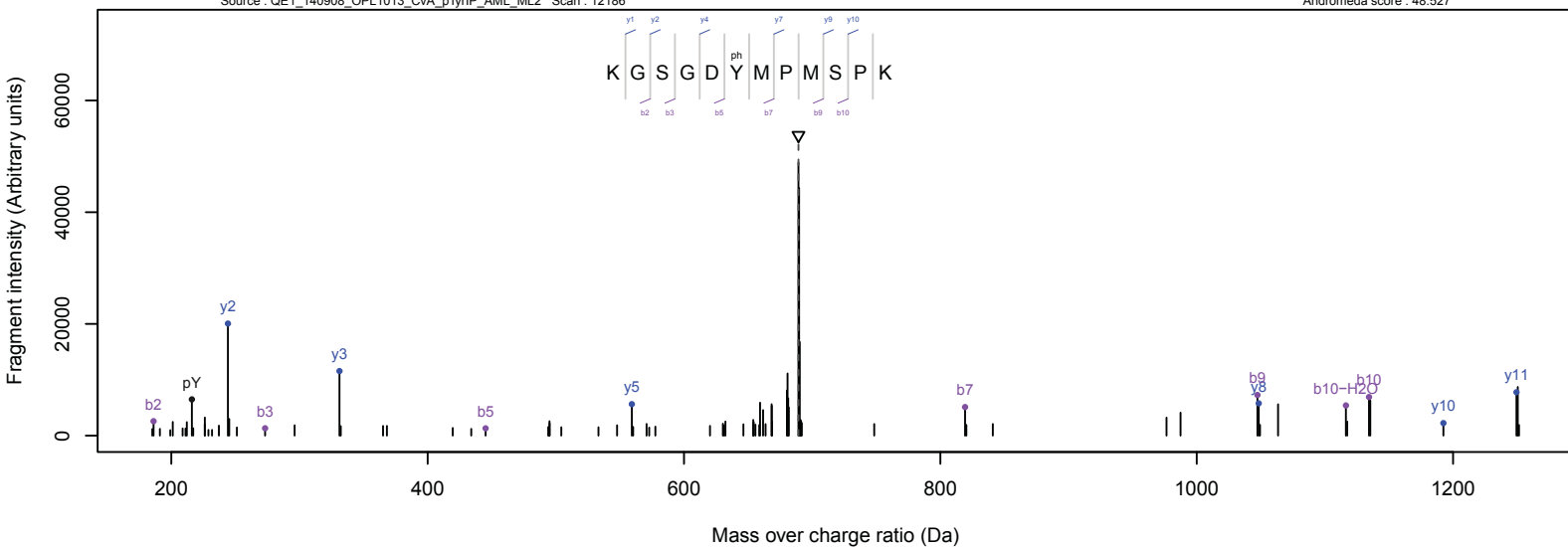
## KGIVSQGY(ph)HR-3+



**KGSGDY(ph)MPMSPK-2+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_ML2 Scan : 12186

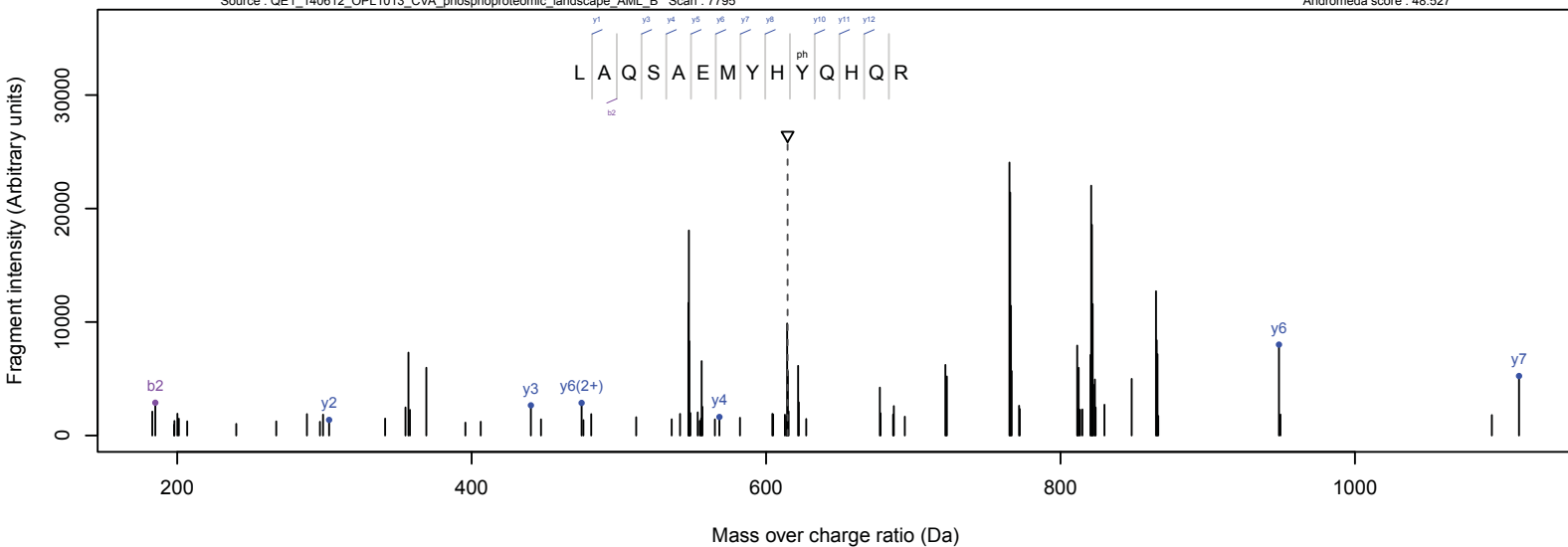
Andromeda score : 48.527



**LAQSAEMYH(ph)QHQR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7795

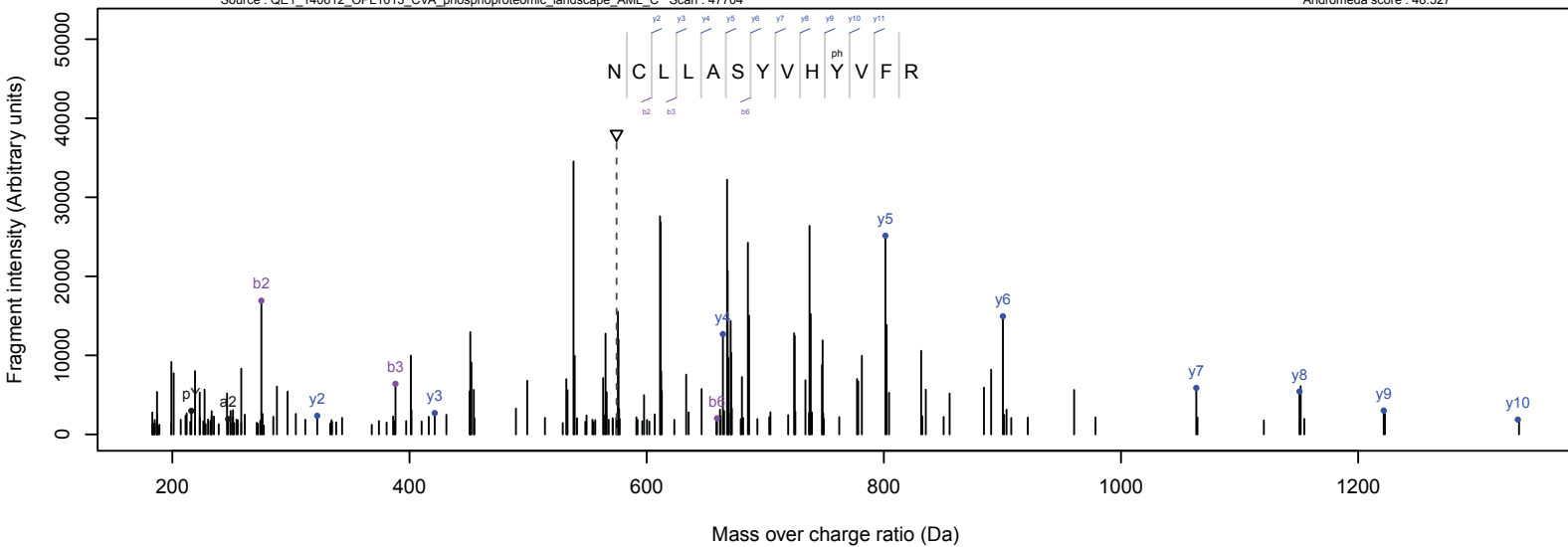
Andromeda score : 48.527



**NCLLASVHY(ph)VFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47704

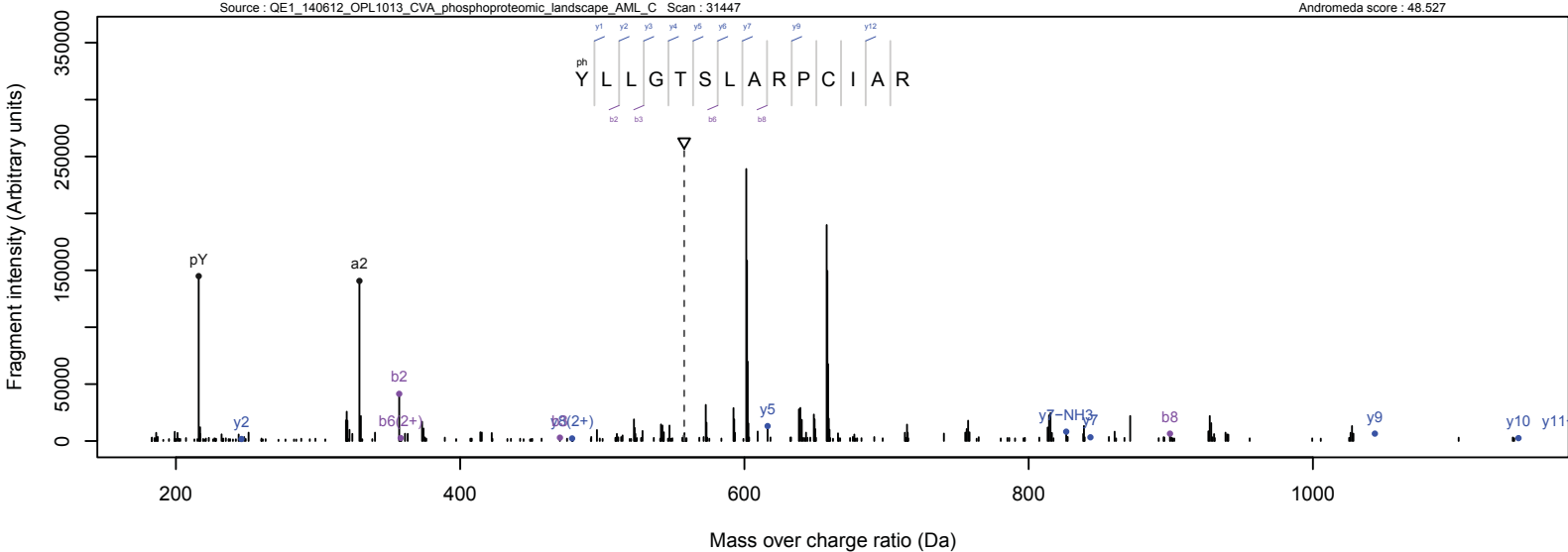
Andromeda score : 48.527



Y(ph)LLGTSLARPCIA R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 31447

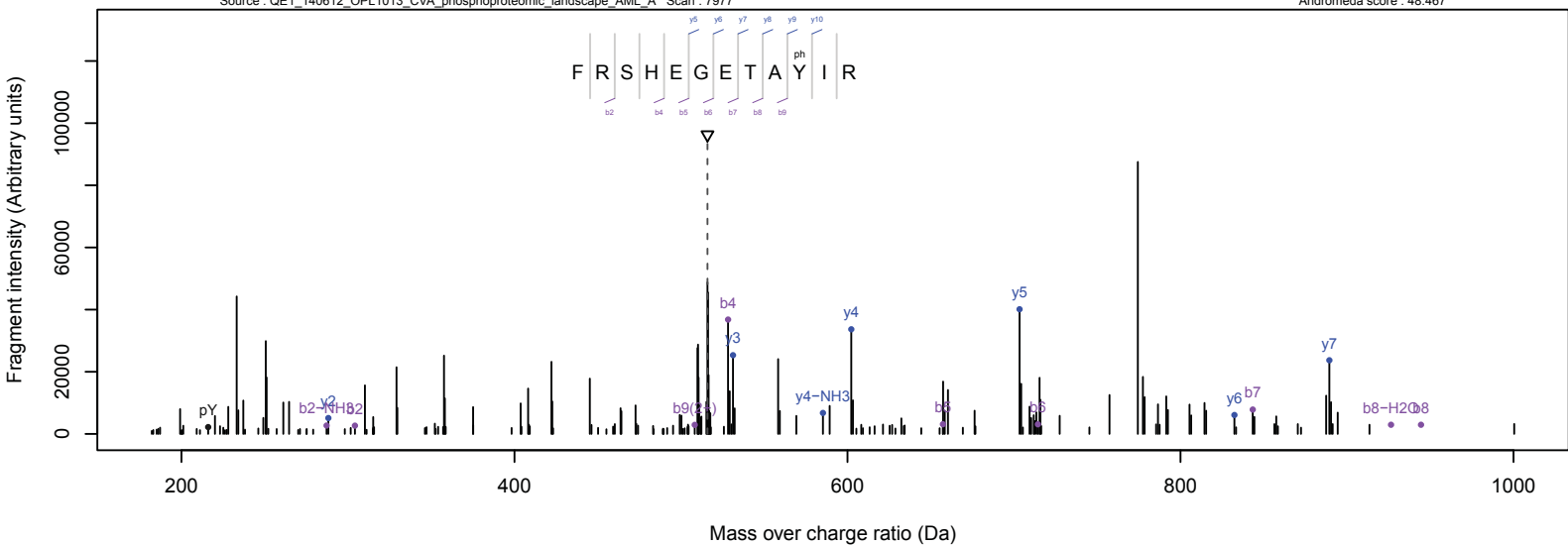
Andromeda score : 48.527



FRSHEGETAY(ph)IR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 7977

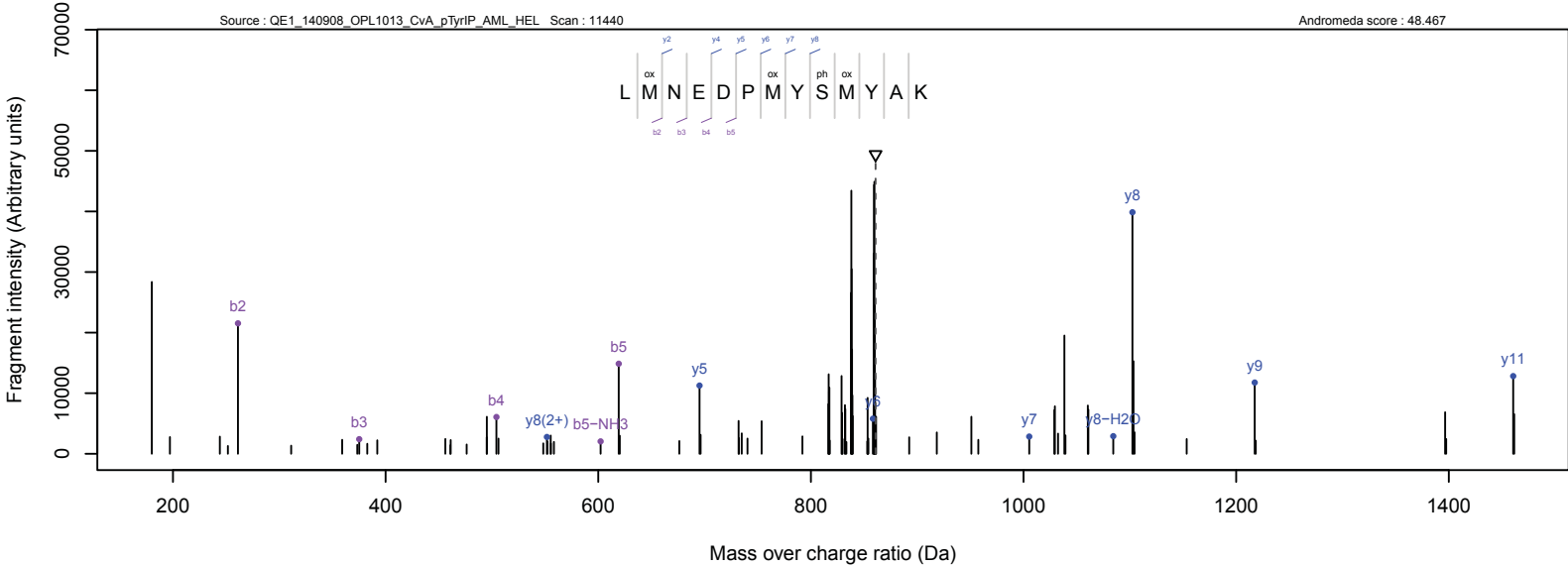
Andromeda score : 48.467



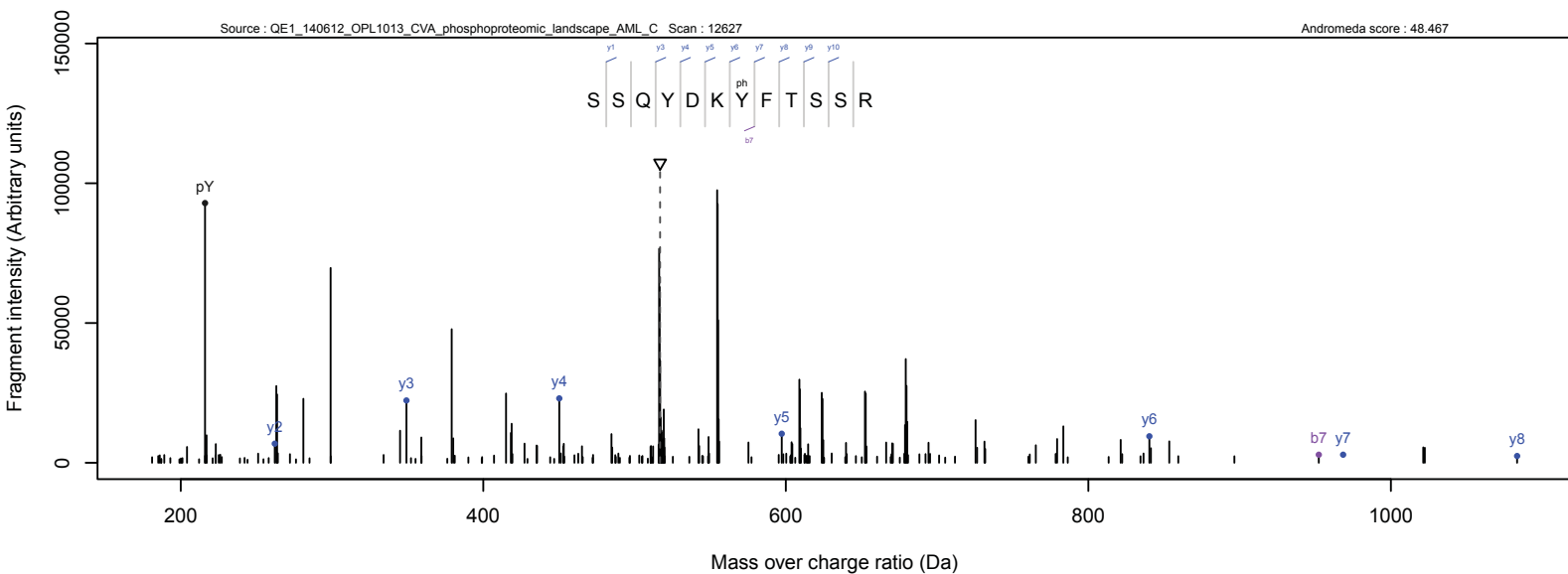
LM(ox)NEDPM(ox)YS(ph)M(ox)YAK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 11440

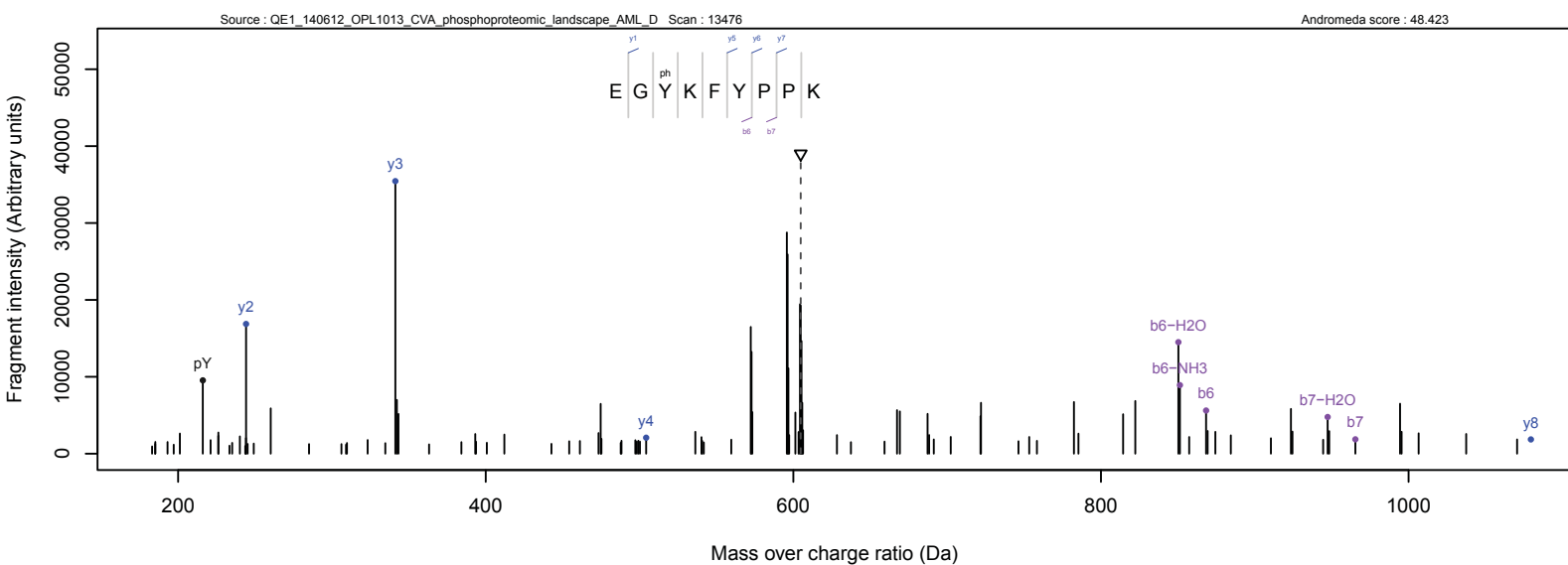
Andromeda score : 48.467



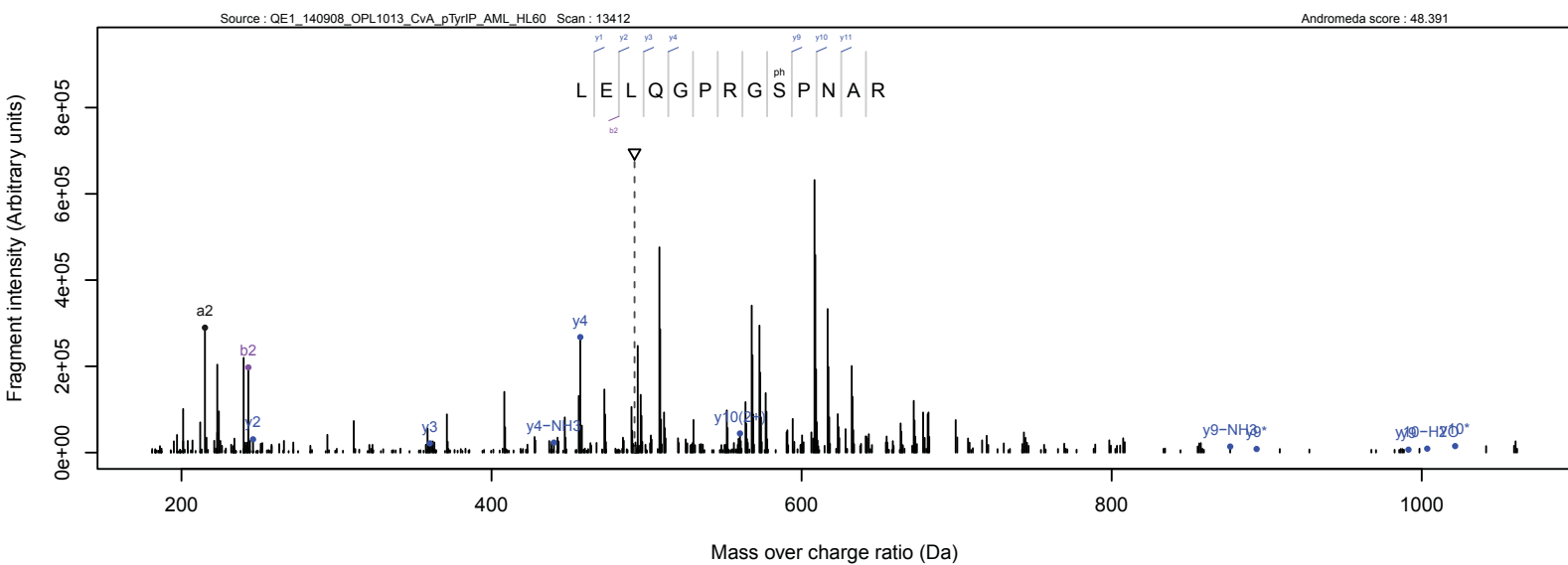
## SSQYDKY(ph)FTSSR-3+



## EGY(ph)KFYPPK-2+



## LELQGRGS(ph)PNAR-3+

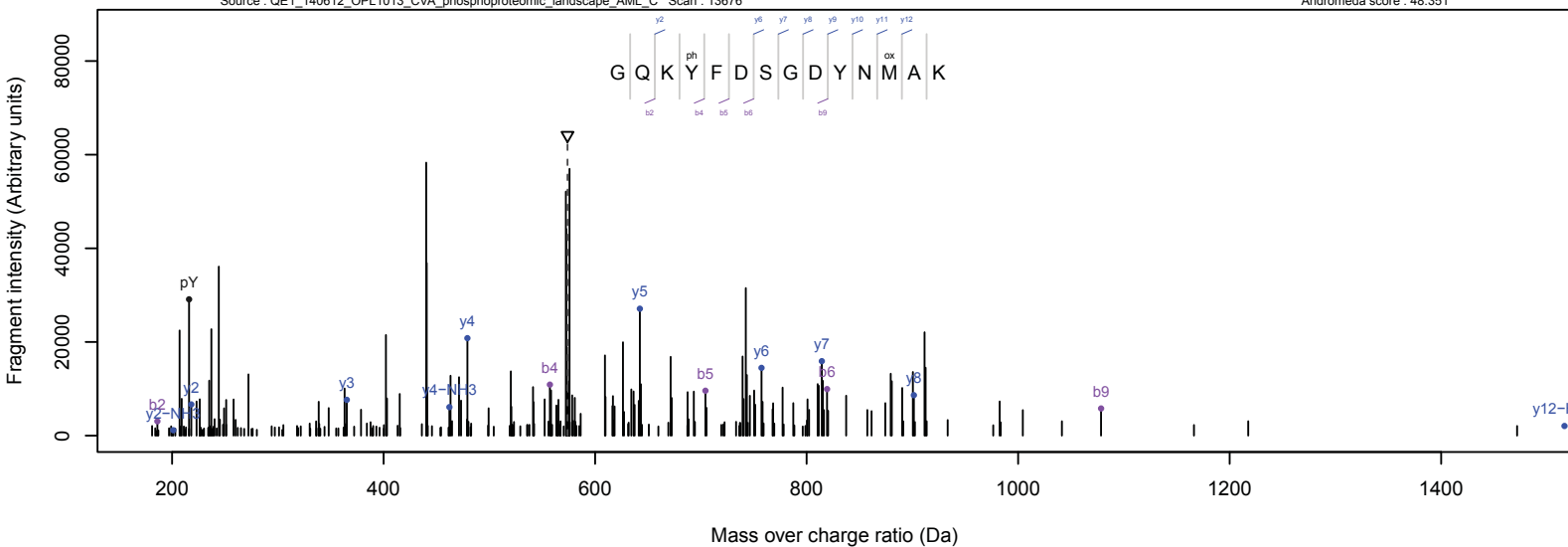




## GQKY(ph)FDSDYNM(ox)AK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13676

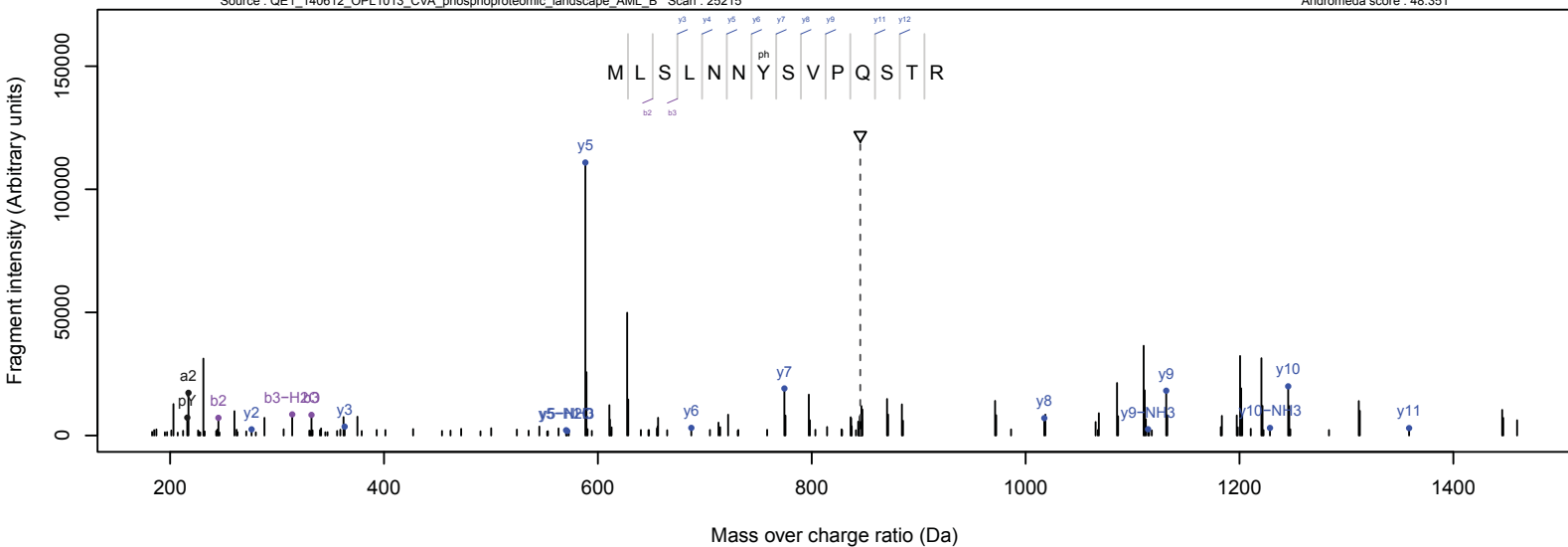
Andromeda score : 48.351



## MLSLNNY(ph)SVPQSTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25215

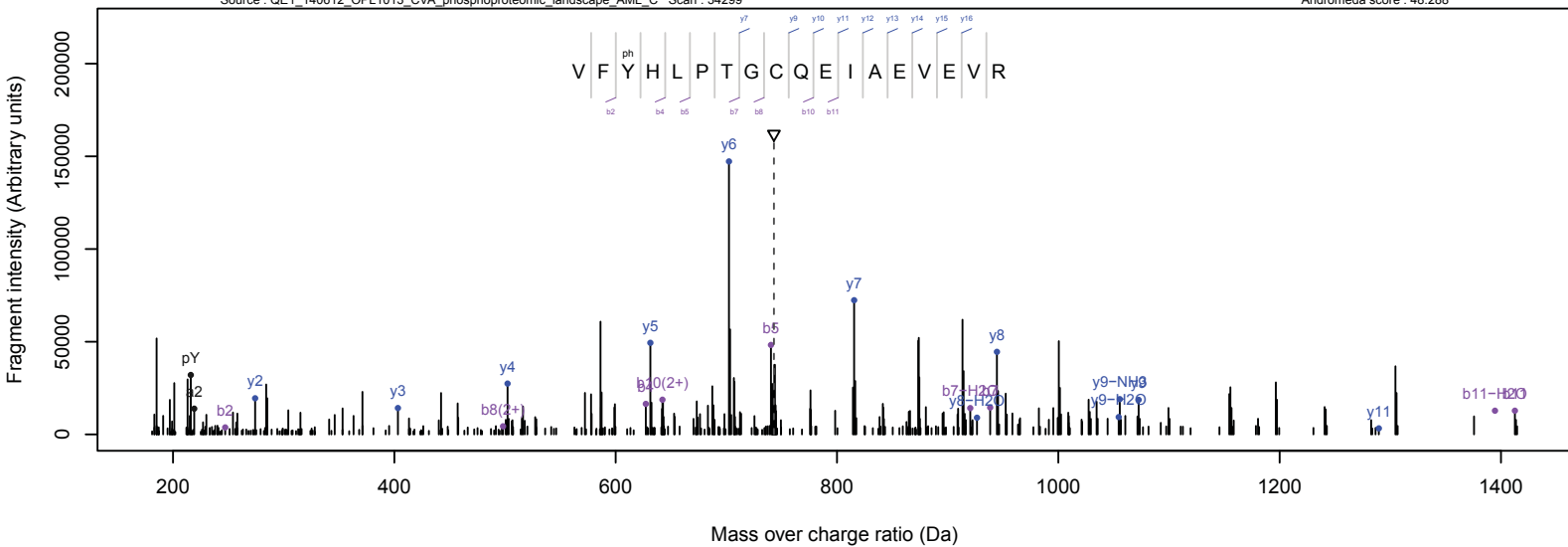
Andromeda score : 48.351



## VFY(ph)HLPTGCQEIAEEVVR-3+

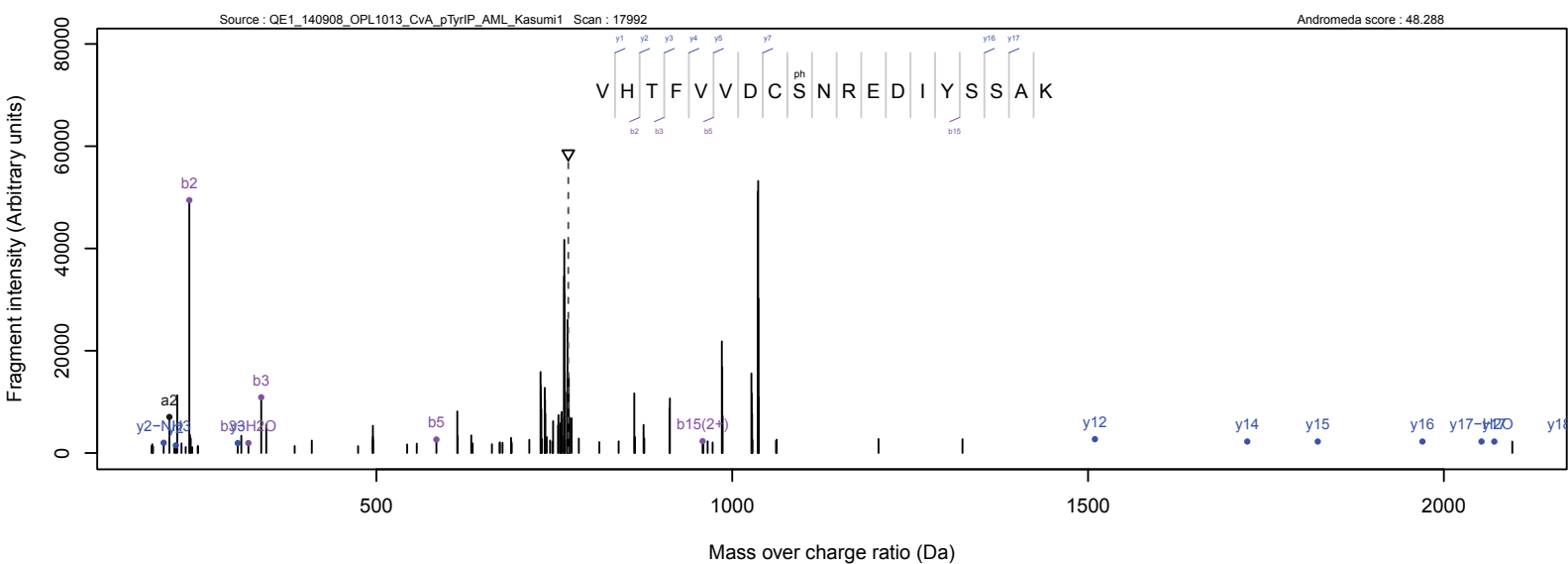
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34299

Andromeda score : 48.288

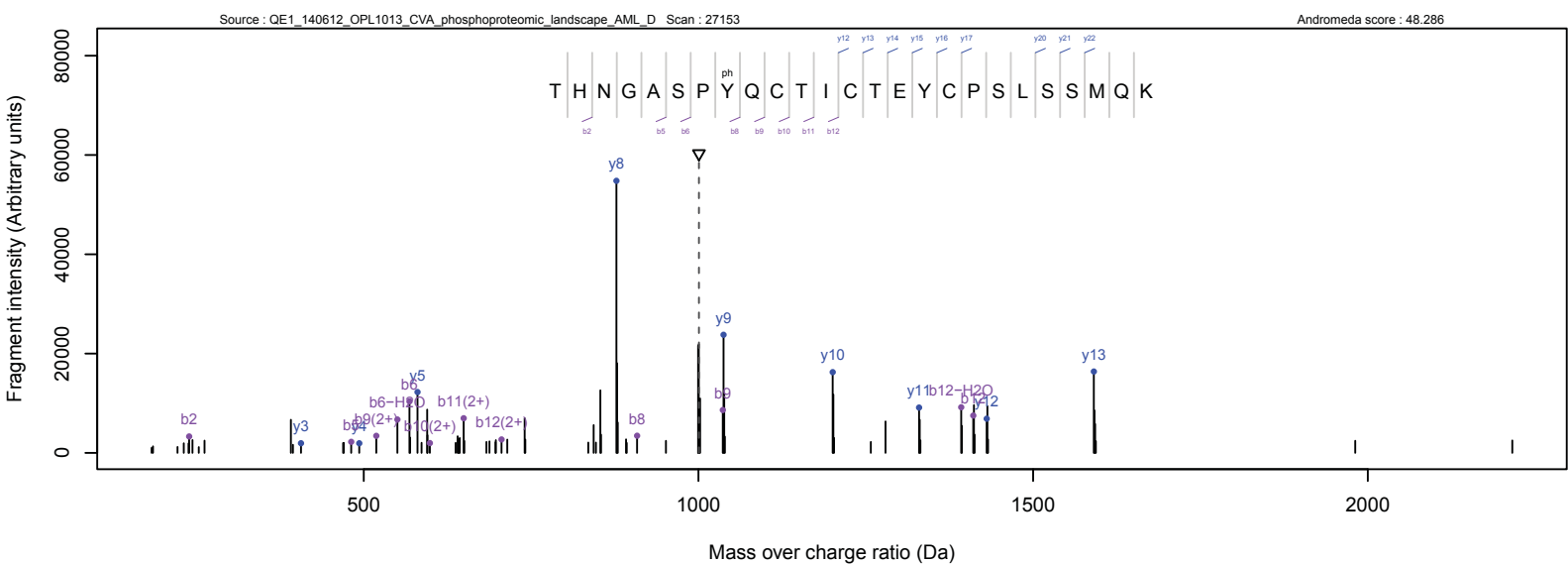




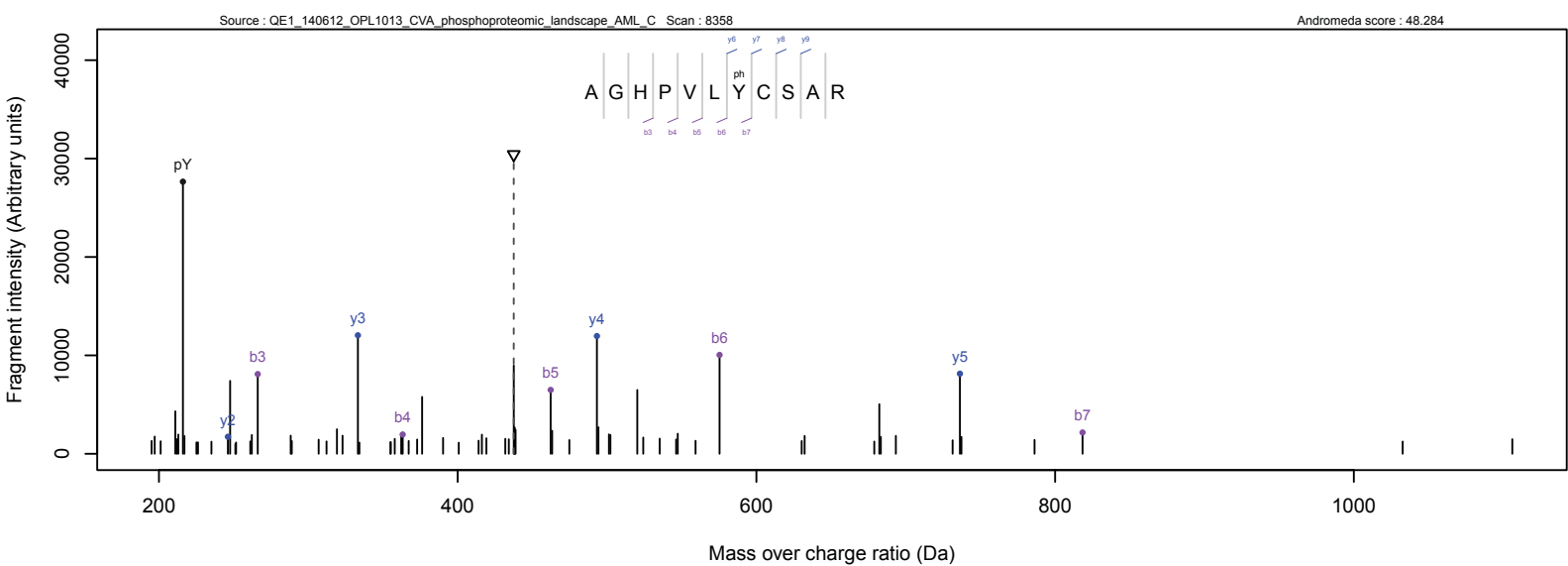
## VHTFVDCS(ph)NREDIYSSAK-3+



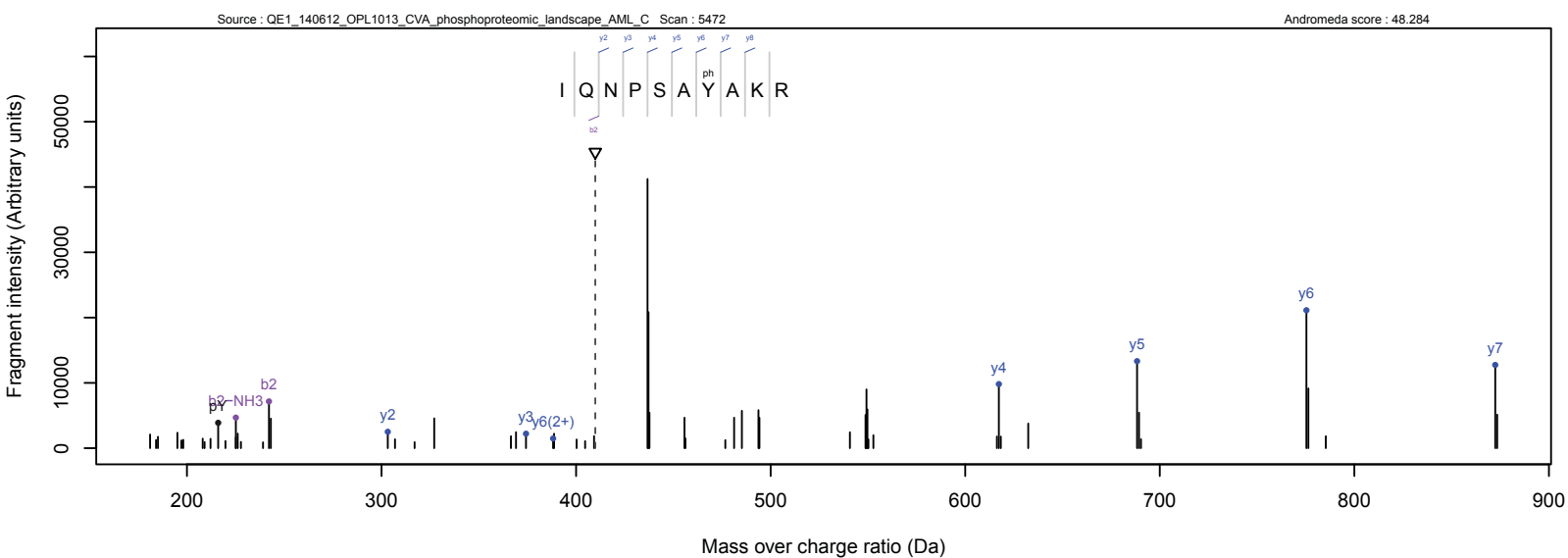
## THNGASPY(ph)QCTICTEYCPSSLSSMQK-3+



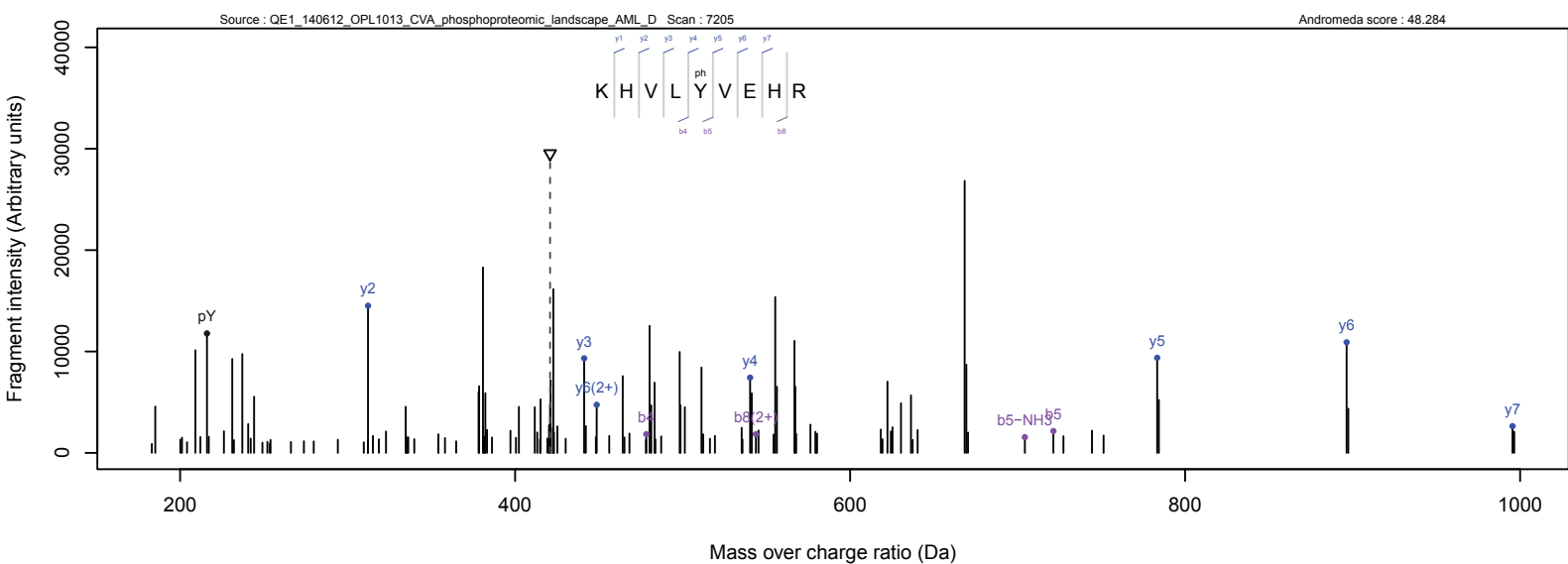
## AGHPVLY(ph)CSAR-3+



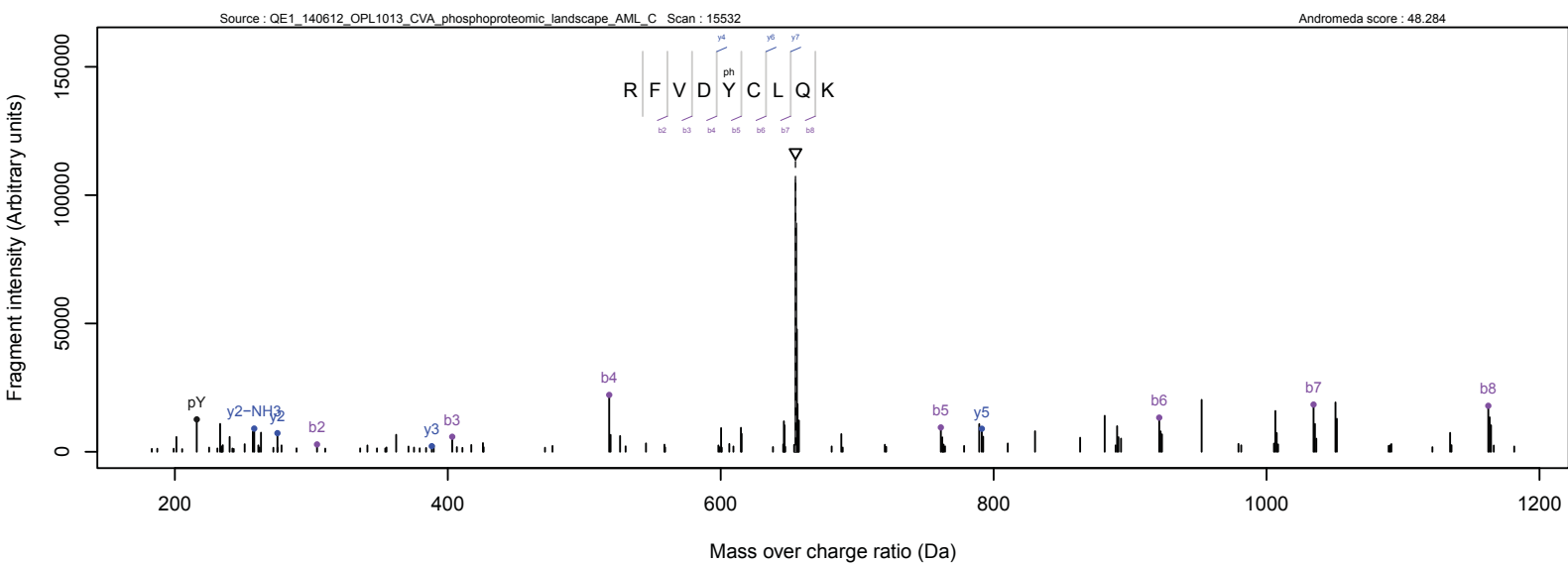
## IQNPSAY(ph)AKR-3+



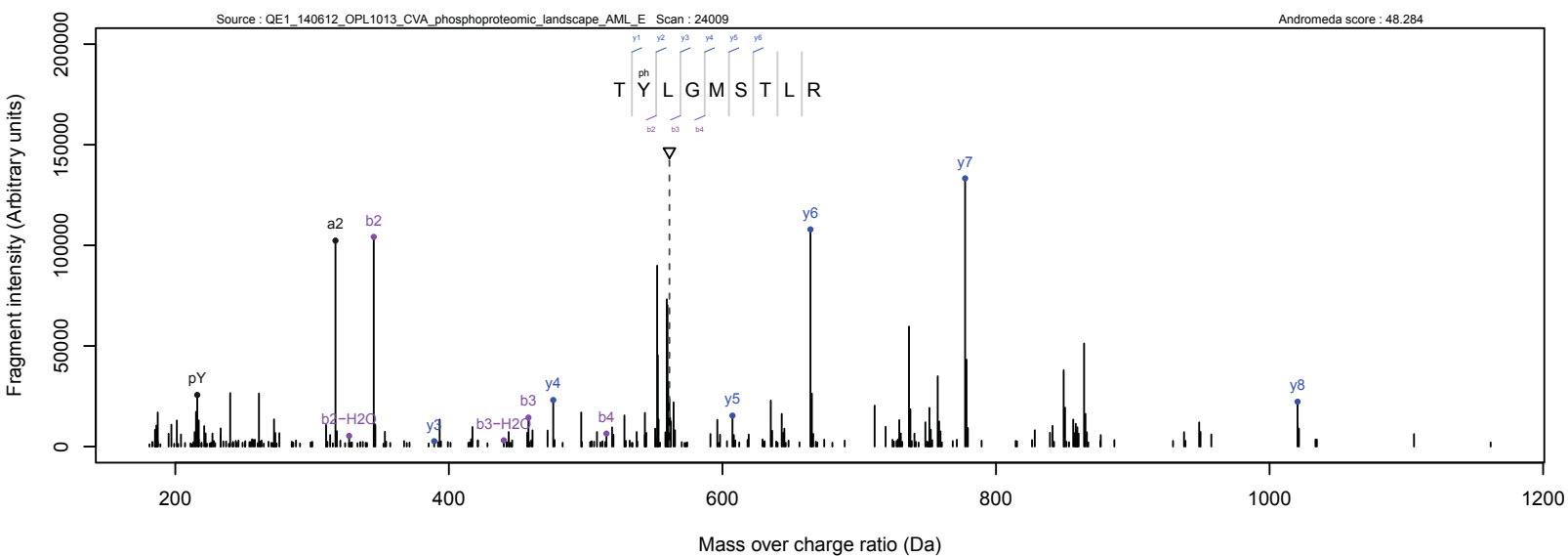
## KHVLY(ph)VEHR-3+



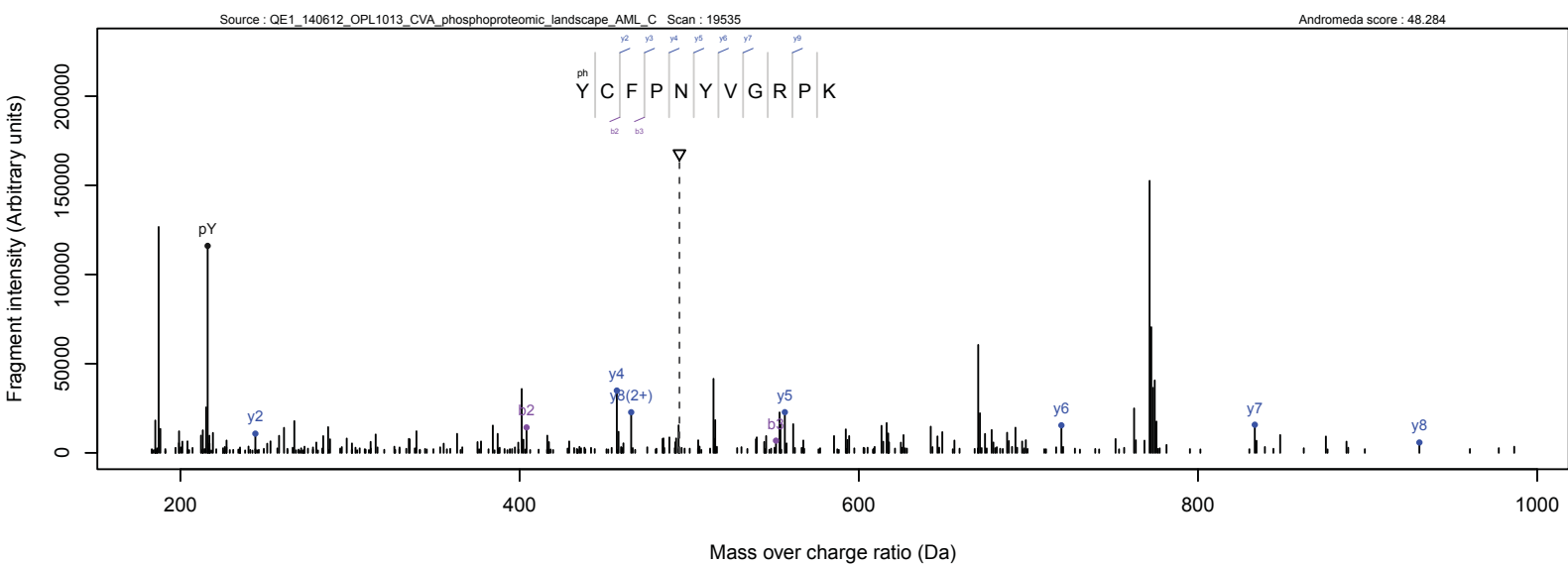
## RFVDY(ph)CLQK-2+



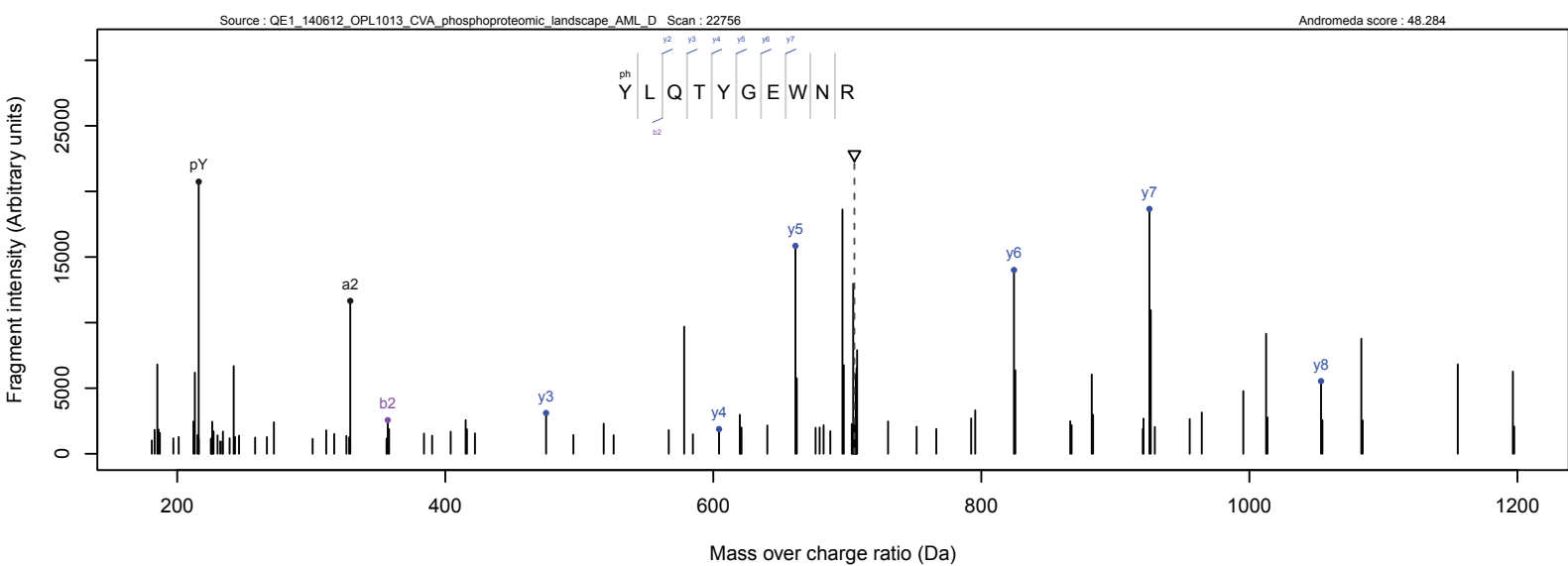
## TY(ph)LGMSTLR-2+



## Y(ph)CFPNYVGRPK-3+



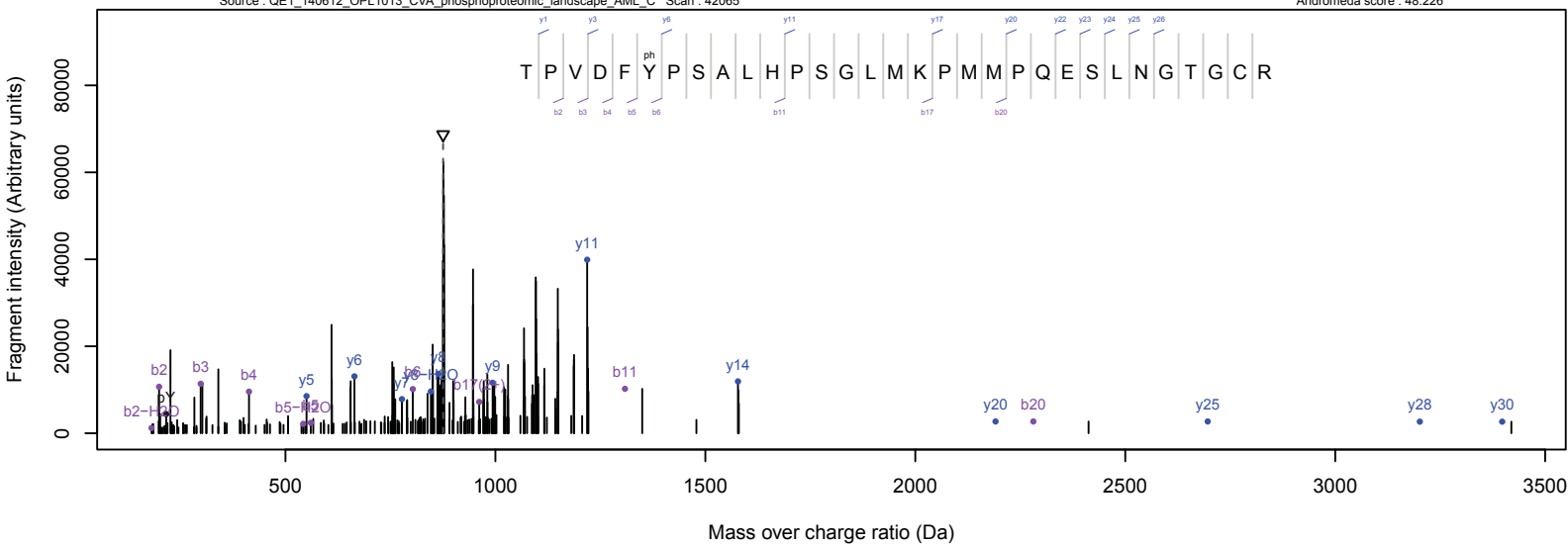
## Y(ph)LQTYGEWNR-2+



## TPVDFY(ph)PSALHPSGLMKPMPQESLNGTGCR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 42065

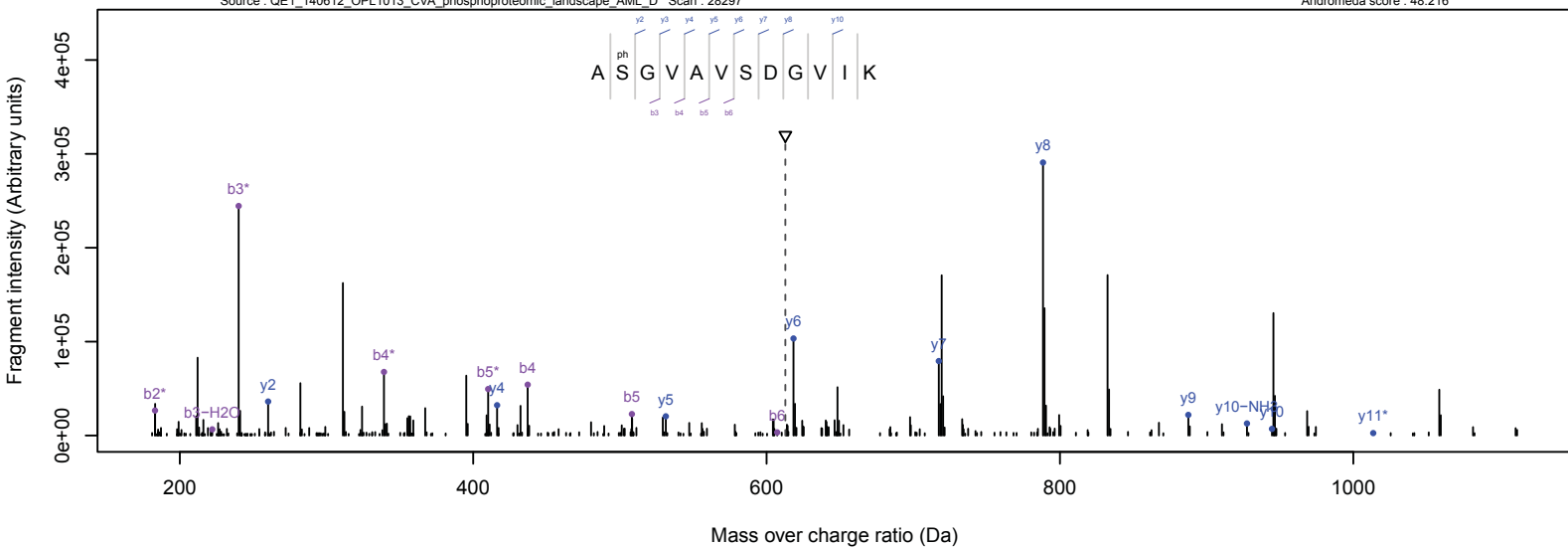
Andromeda score : 48.226



## (ac)AS(ph)GVAVSDGVIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28297

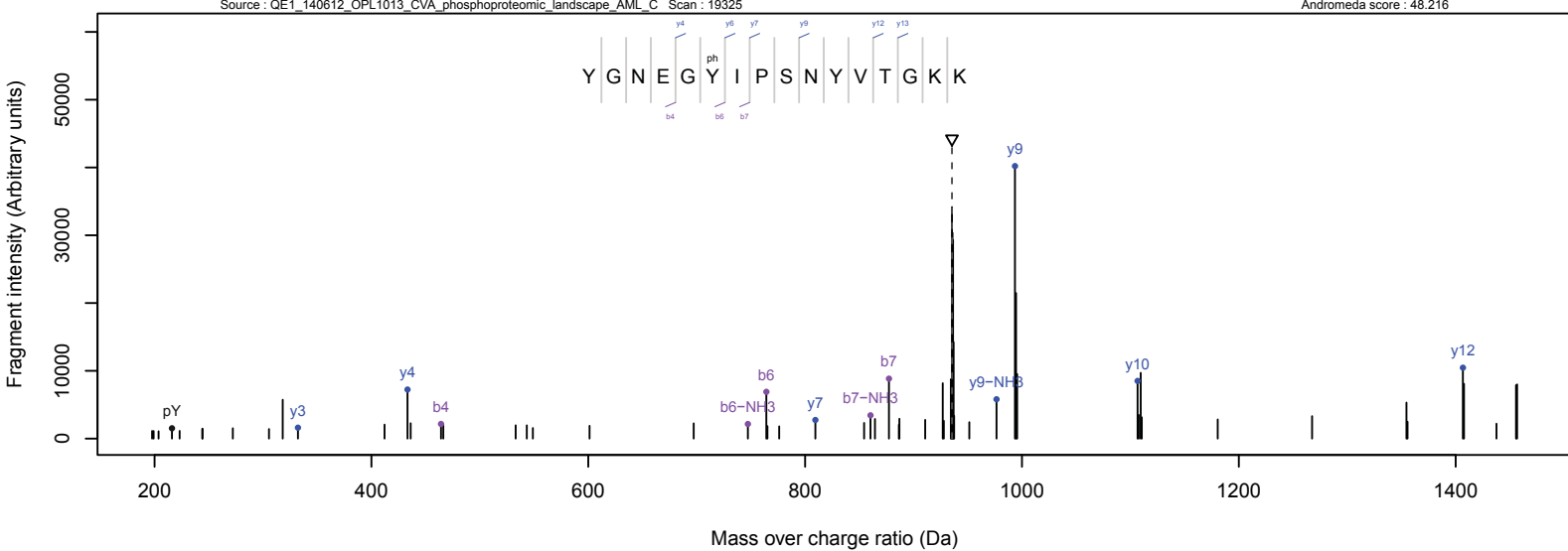
Andromeda score : 48.216



## YGNEY(ph)IPSNYVTGKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19325

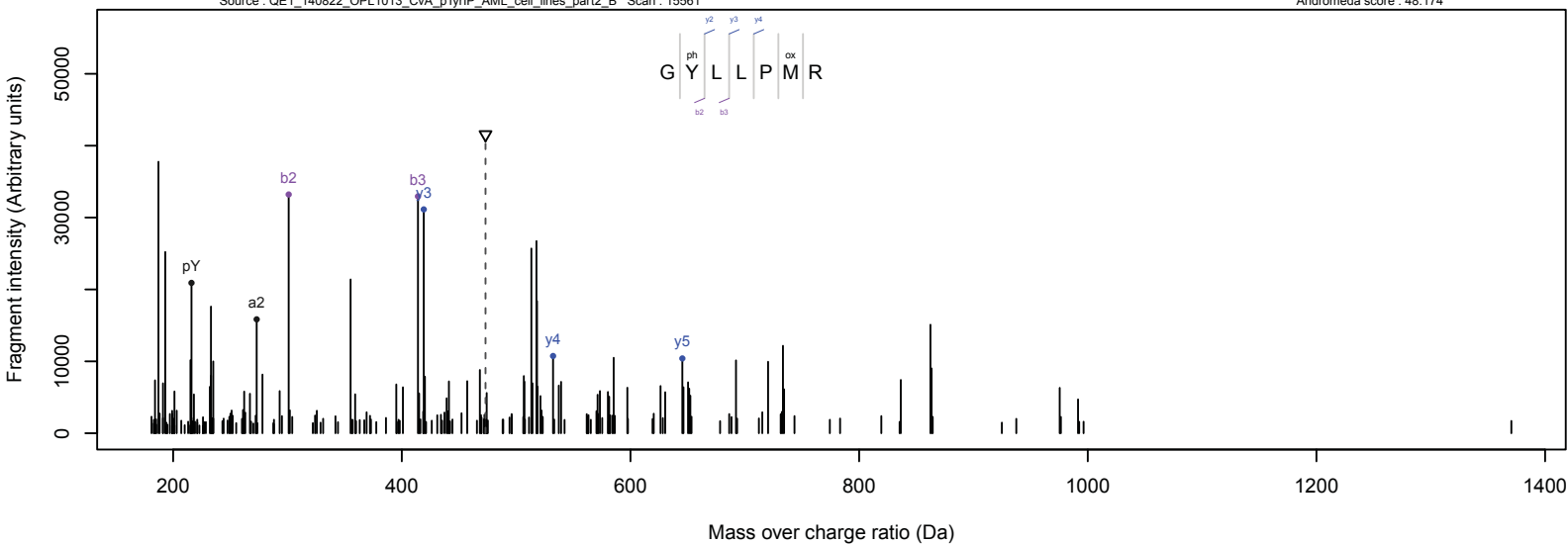
Andromeda score : 48.216



GY(ph)LLPM(ox)R-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 15561

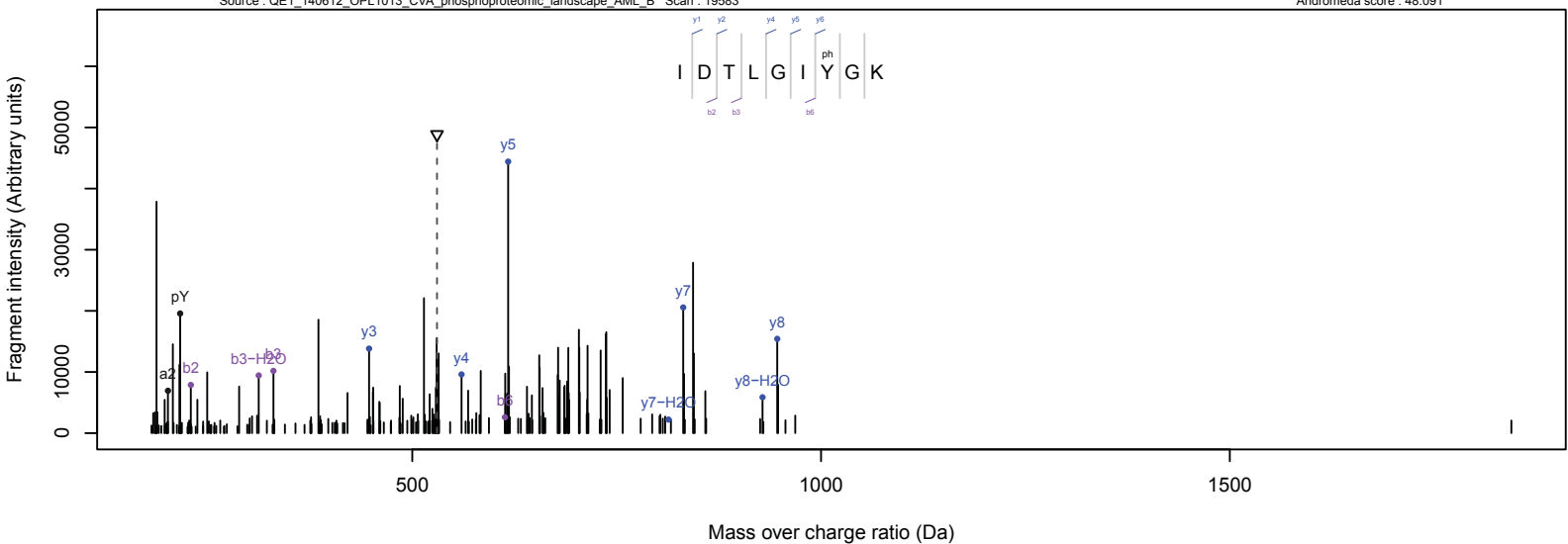
Andromeda score: 48.174



IDTLGIY(ph)GK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 19583

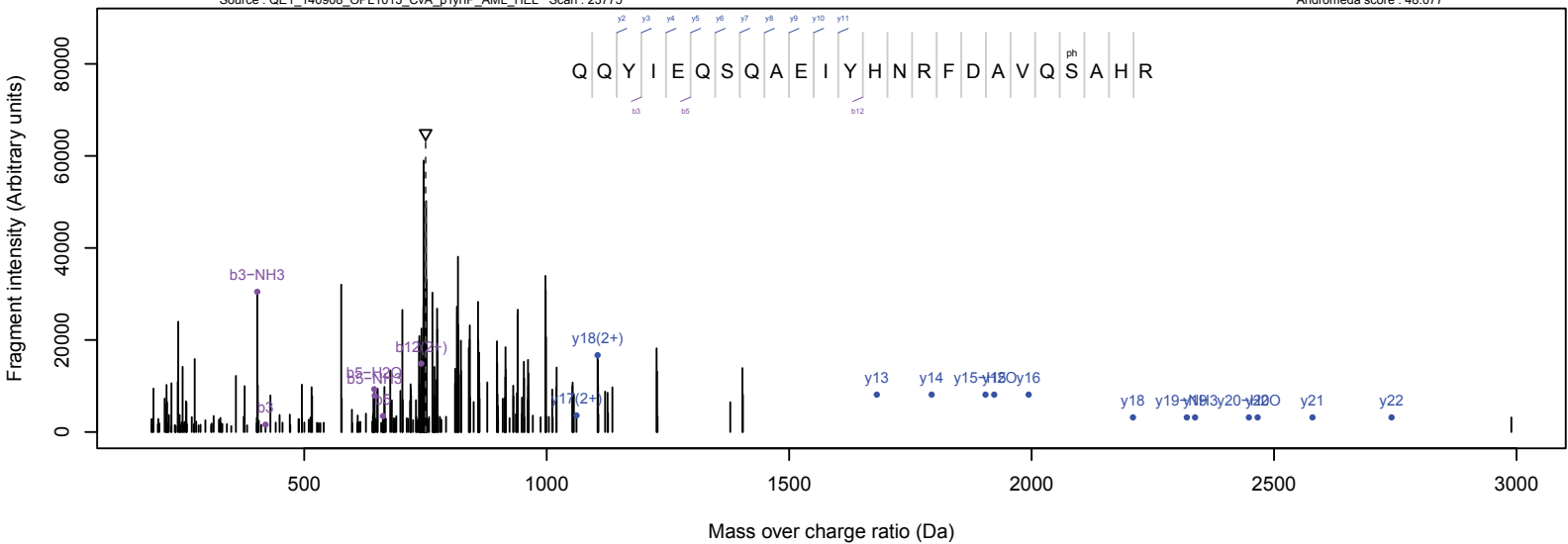
Andromeda score: 48.091



QQYIEQSQA E I Y H N R F D A V Q S A H R-4+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan: 23775

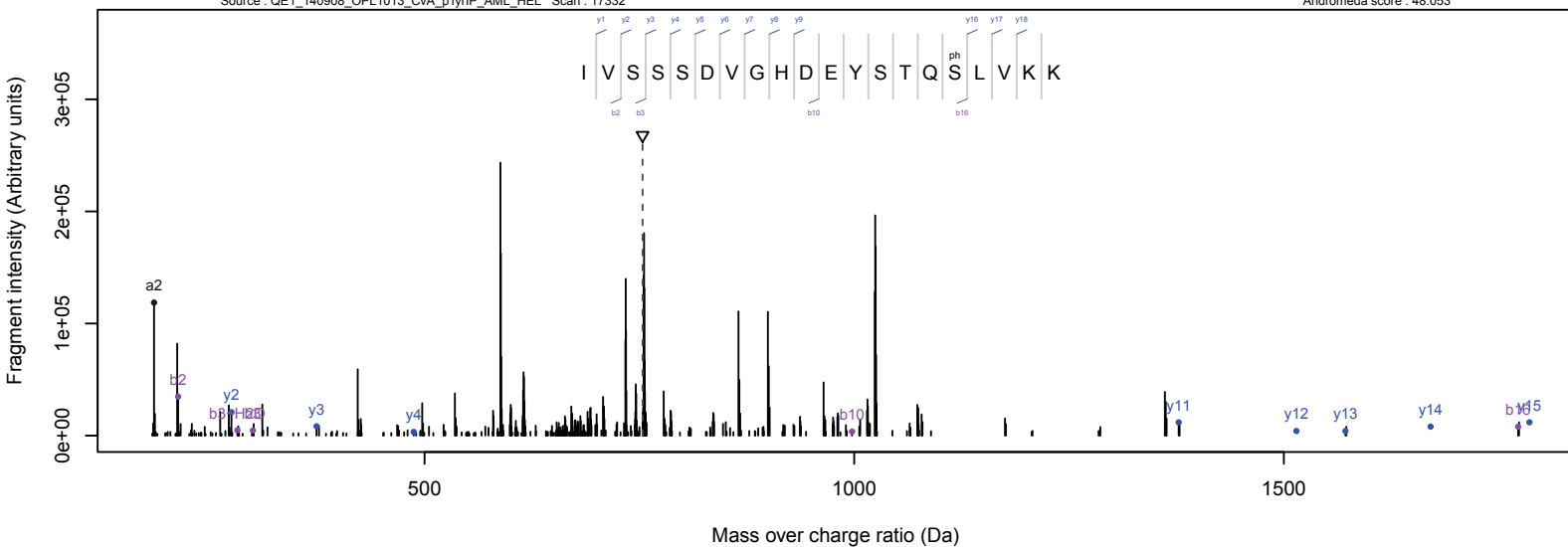
Andromeda score: 48.077



## IVSSSDVGHDEYSTQ(ph)LVKK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 17332

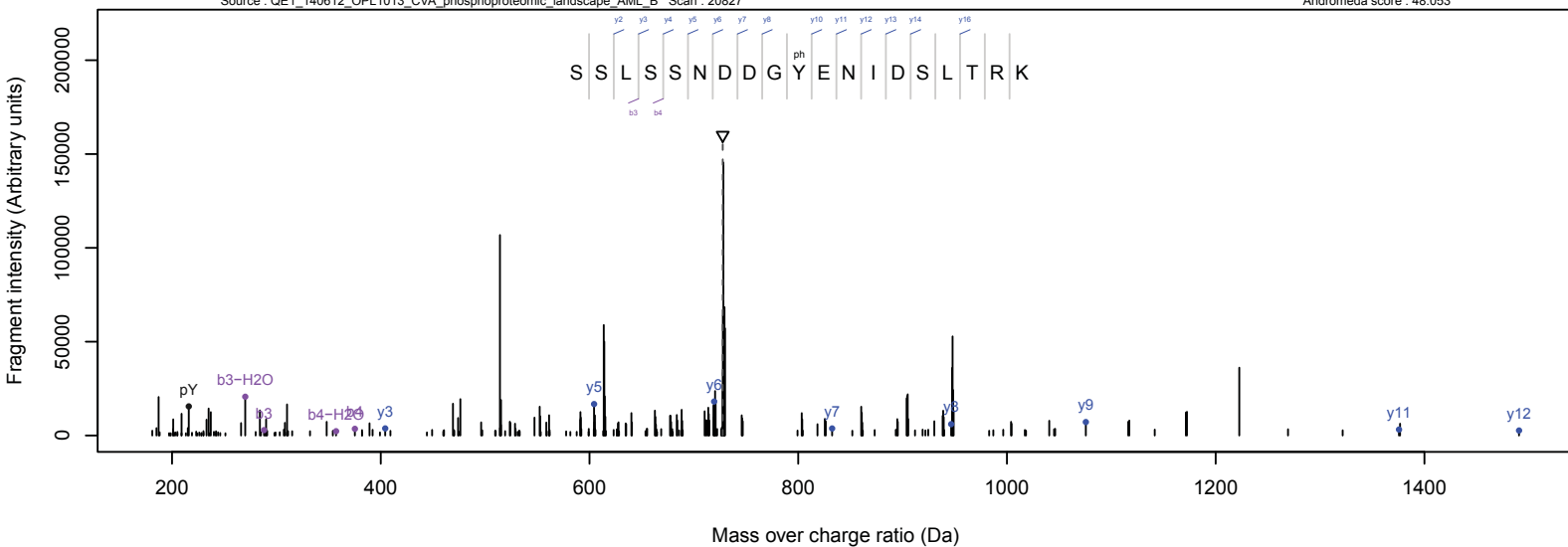
Andromeda score : 48.053



## SSLSSNDG(ph)ENIDSLTRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20827

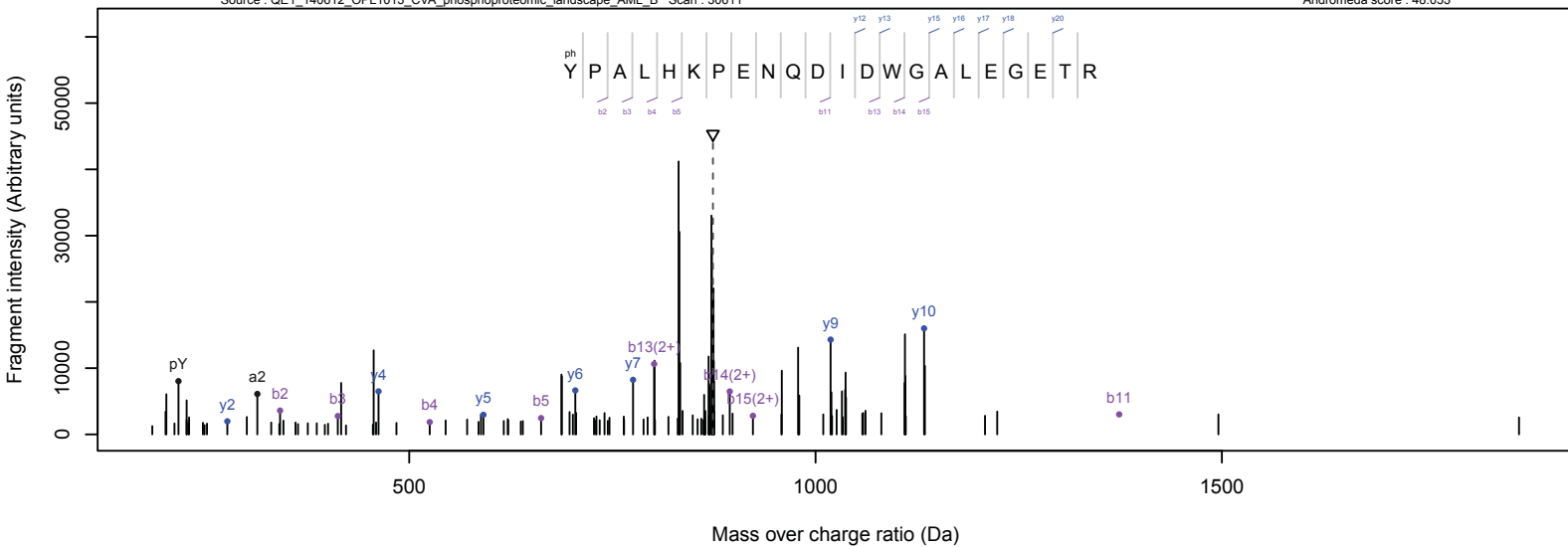
Andromeda score : 48.053



## Y(ph)PALHKPENQDIDWGALEGETR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30611

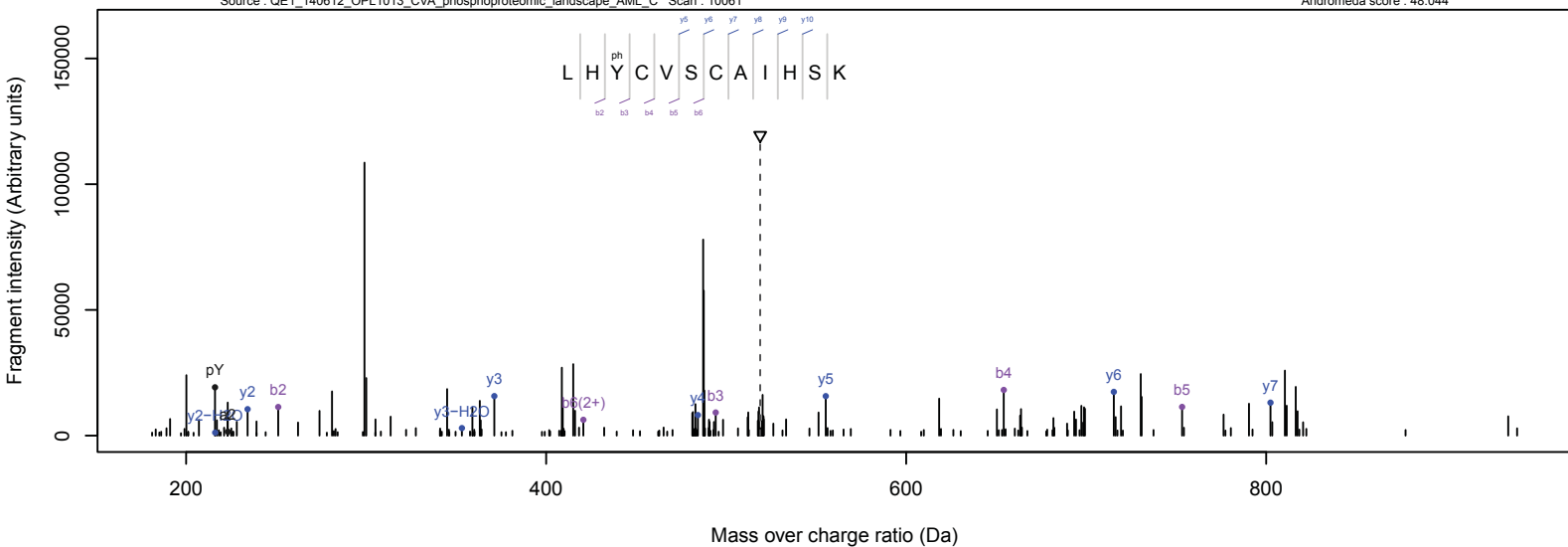
Andromeda score : 48.053



LHY(ph)CVSCAIHSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10061

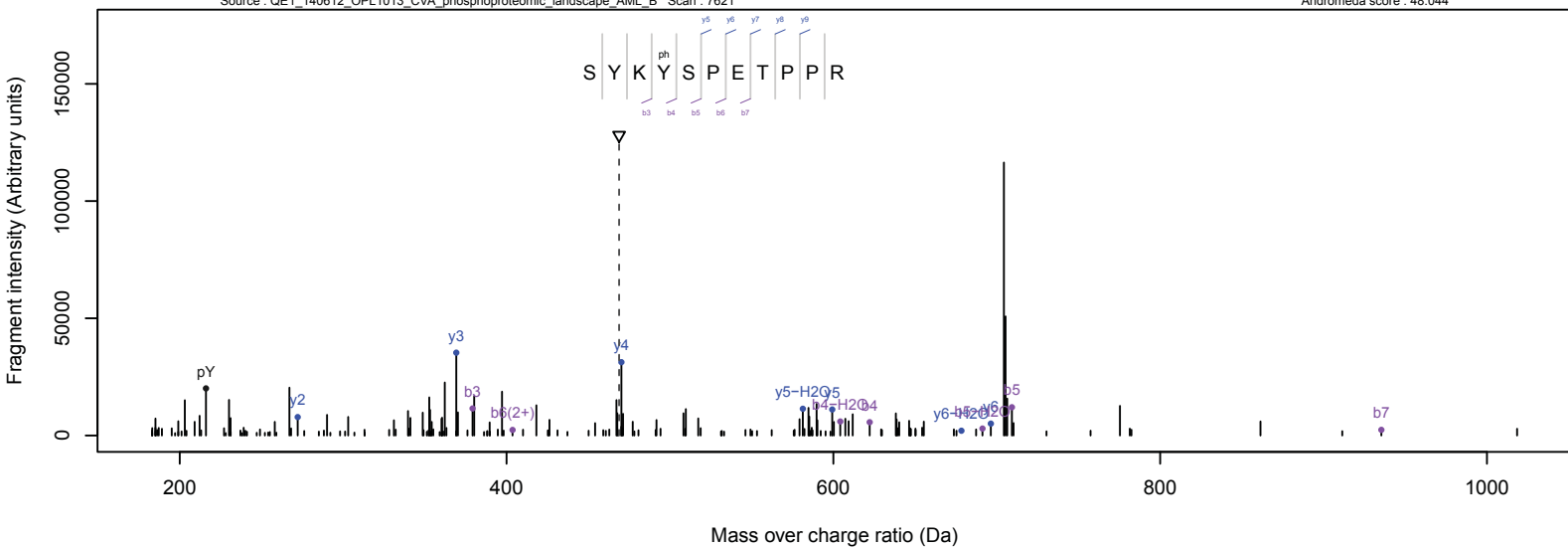
Andromeda score : 48.044



SYKY(ph)SPETPPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 7621

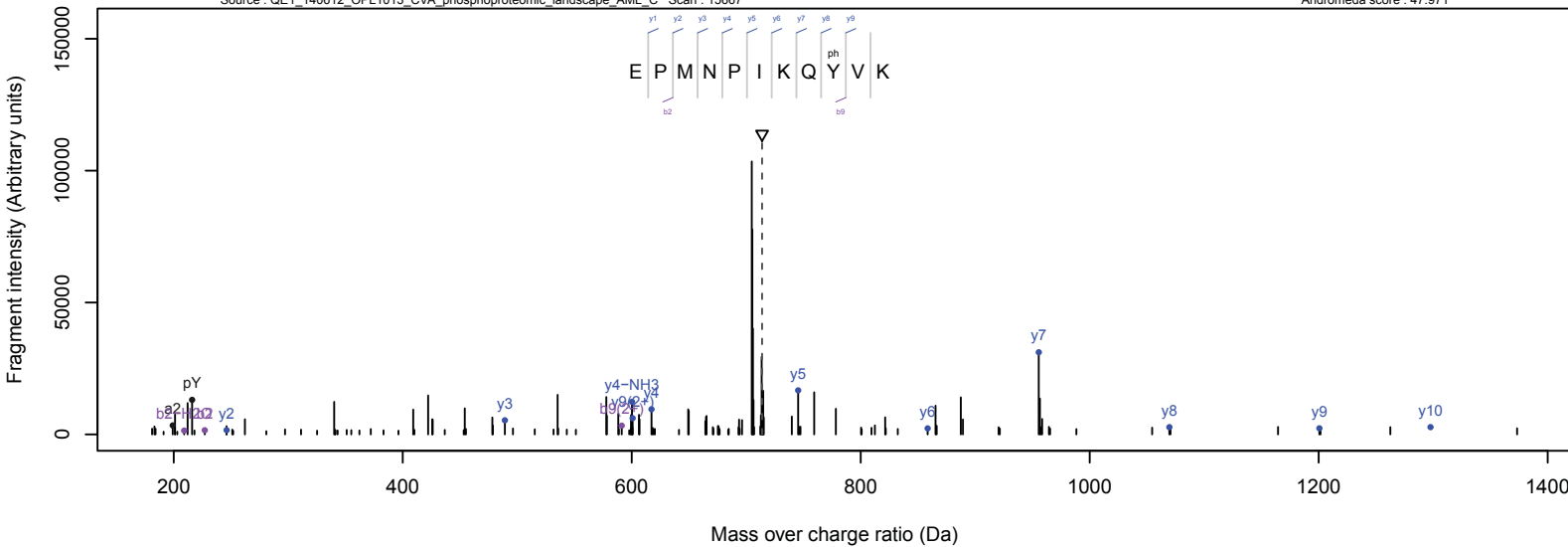
Andromeda score : 48.044



EPMNPIKQYVVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15667

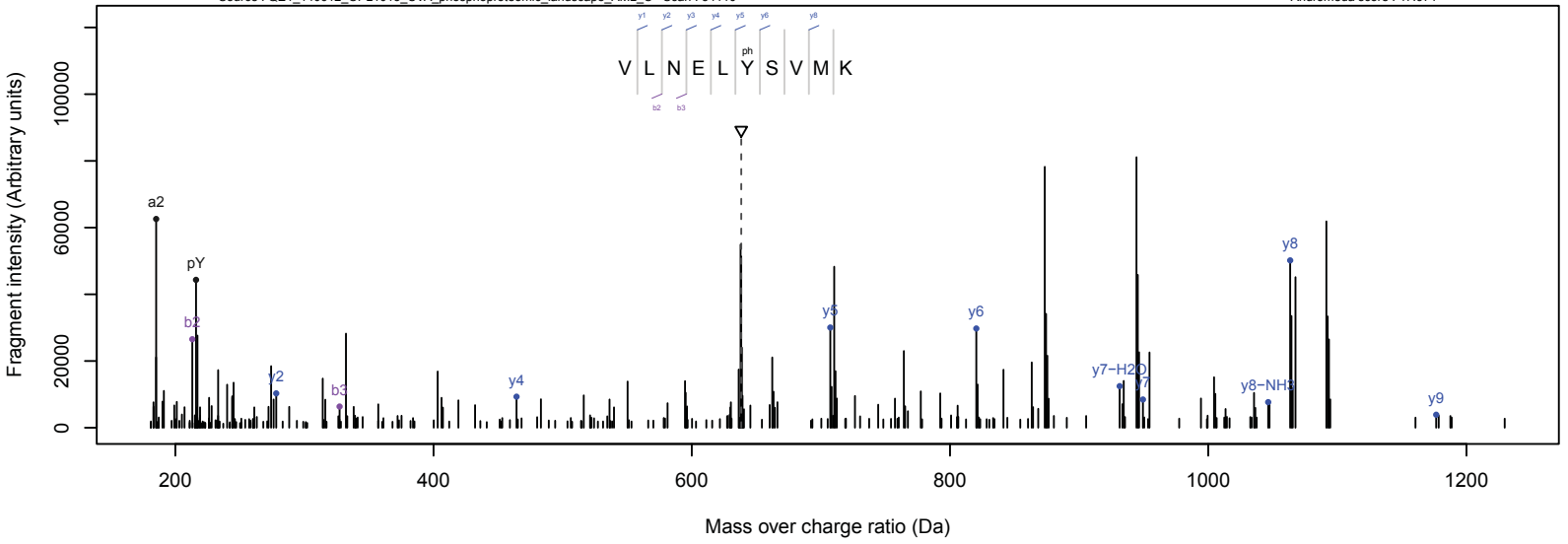
Andromeda score : 47.971



**VLNELY(ph)SVMK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34110

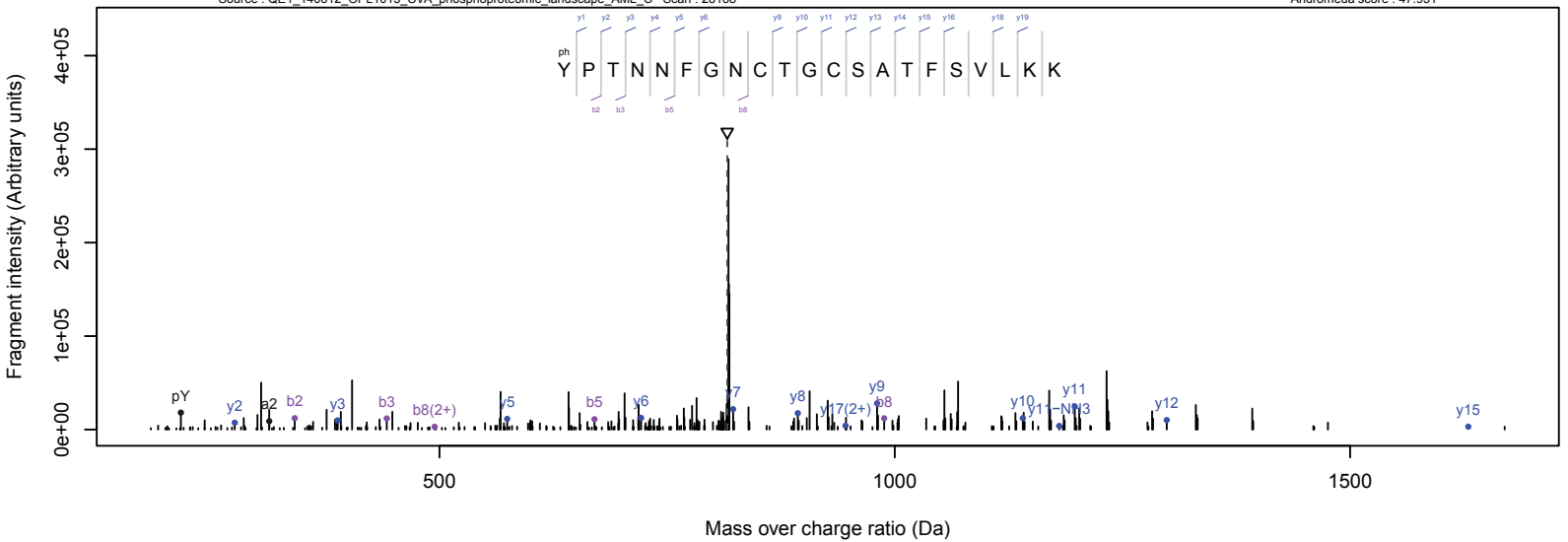
Andromeda score : 47.971



**Y(ph)PTNNGNCTGSATFSVLKK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28188

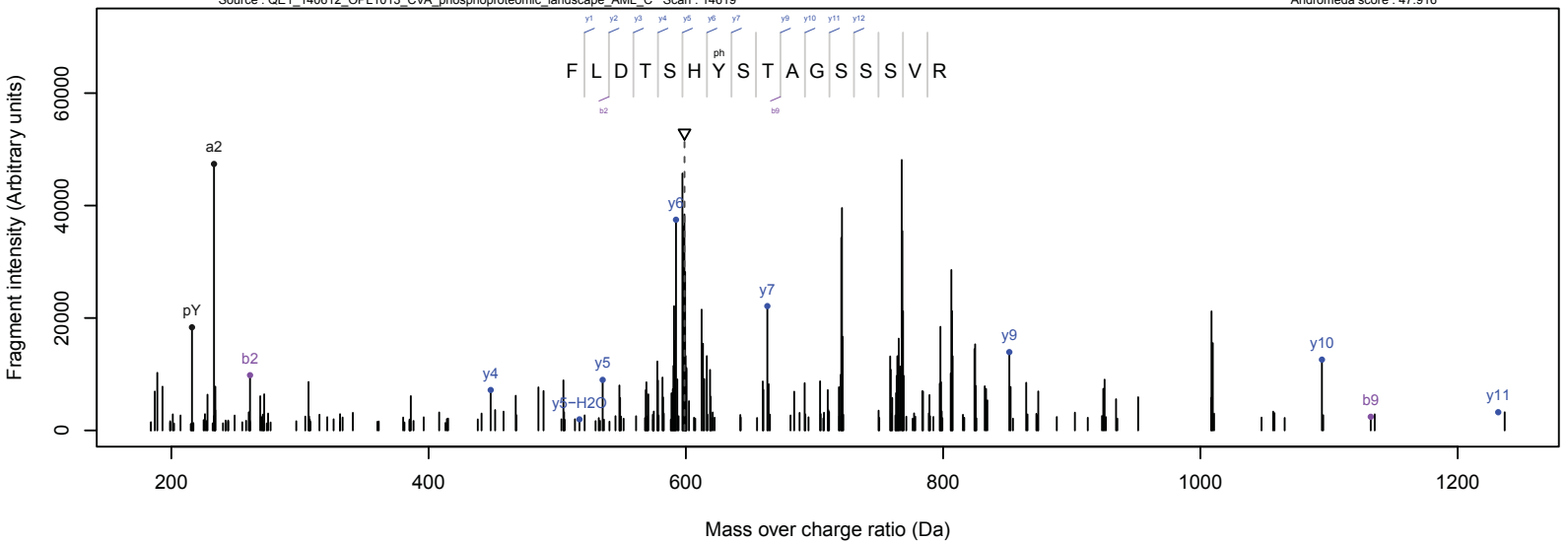
Andromeda score : 47.931



**FLDTSHY(ph)STAGSSSVR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14619

Andromeda score : 47.916

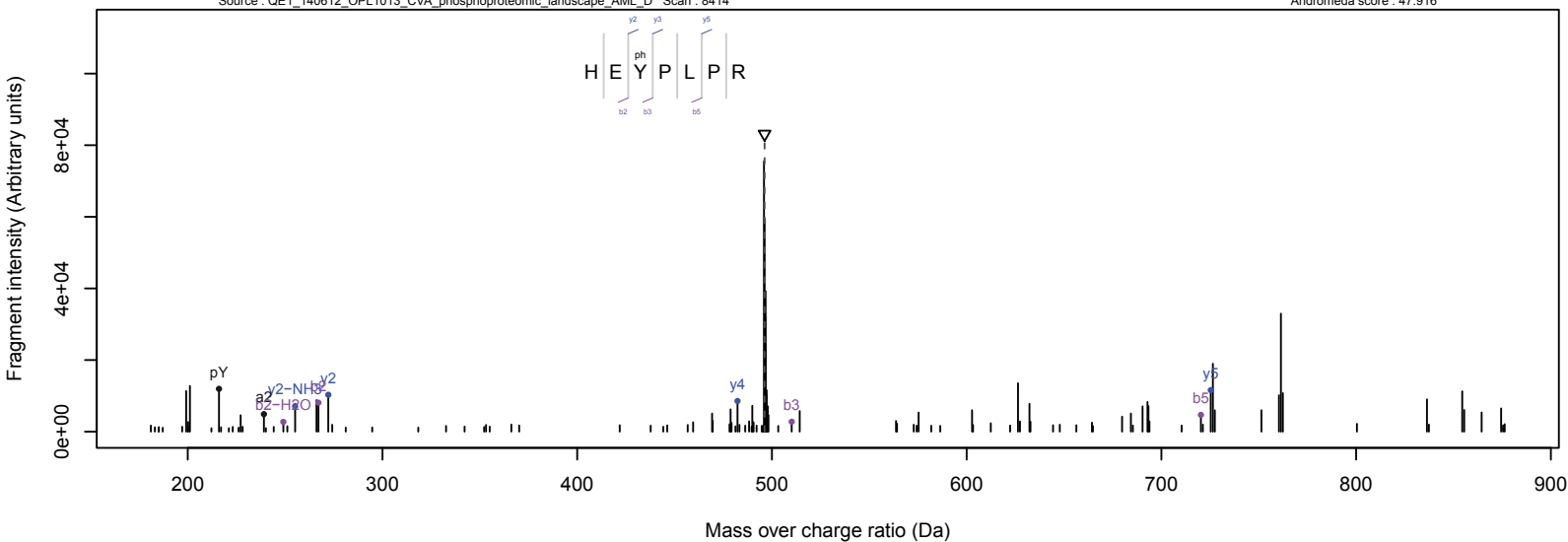




## HEY(ph)PLPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 8414

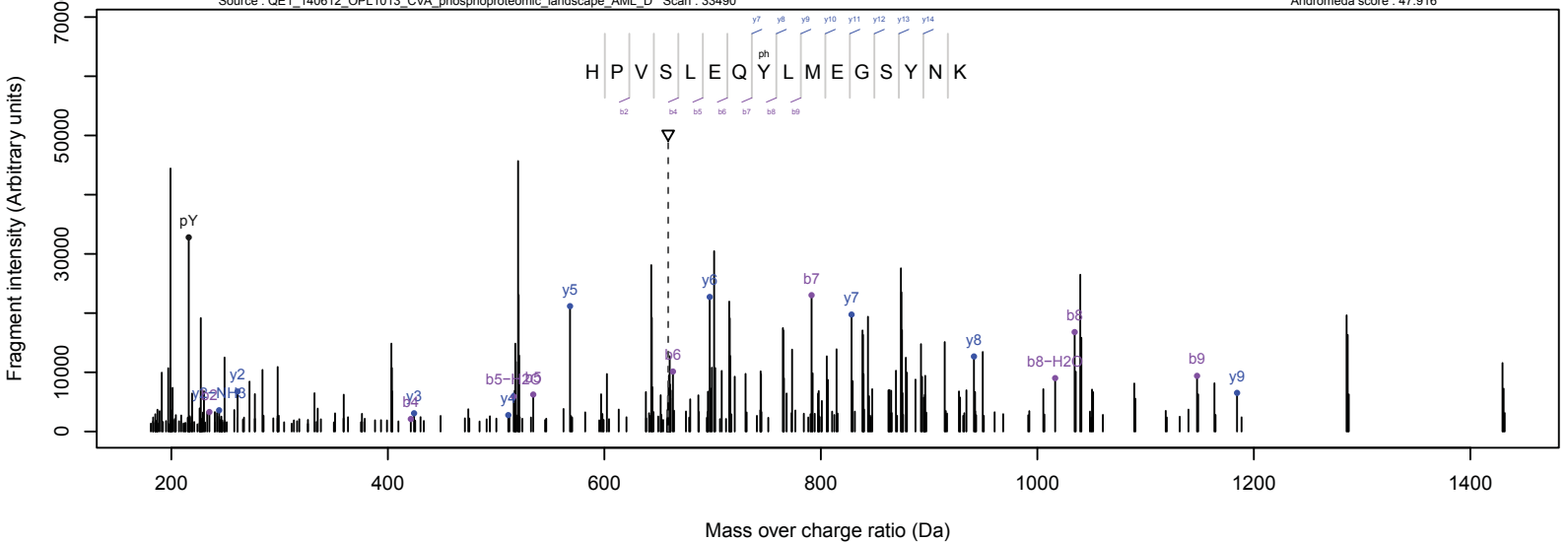
Andromeda score: 47.916



## HPVSLQY(ph)LMEGSYNK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 33490

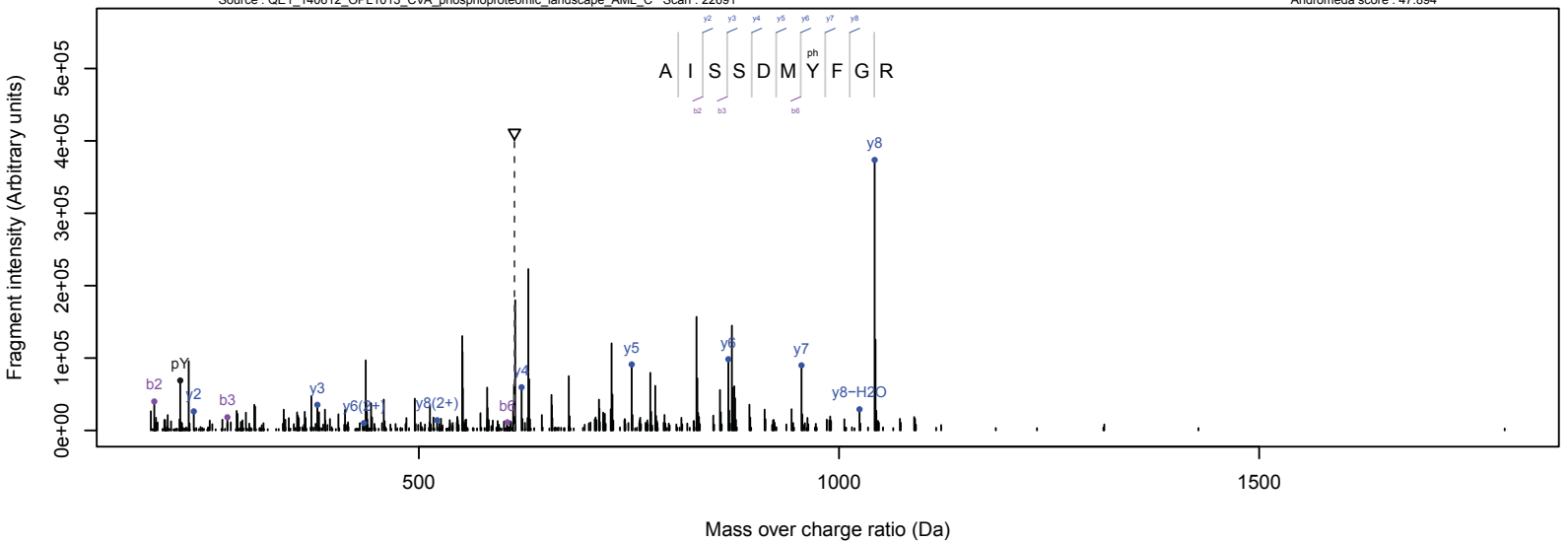
Andromeda score: 47.916

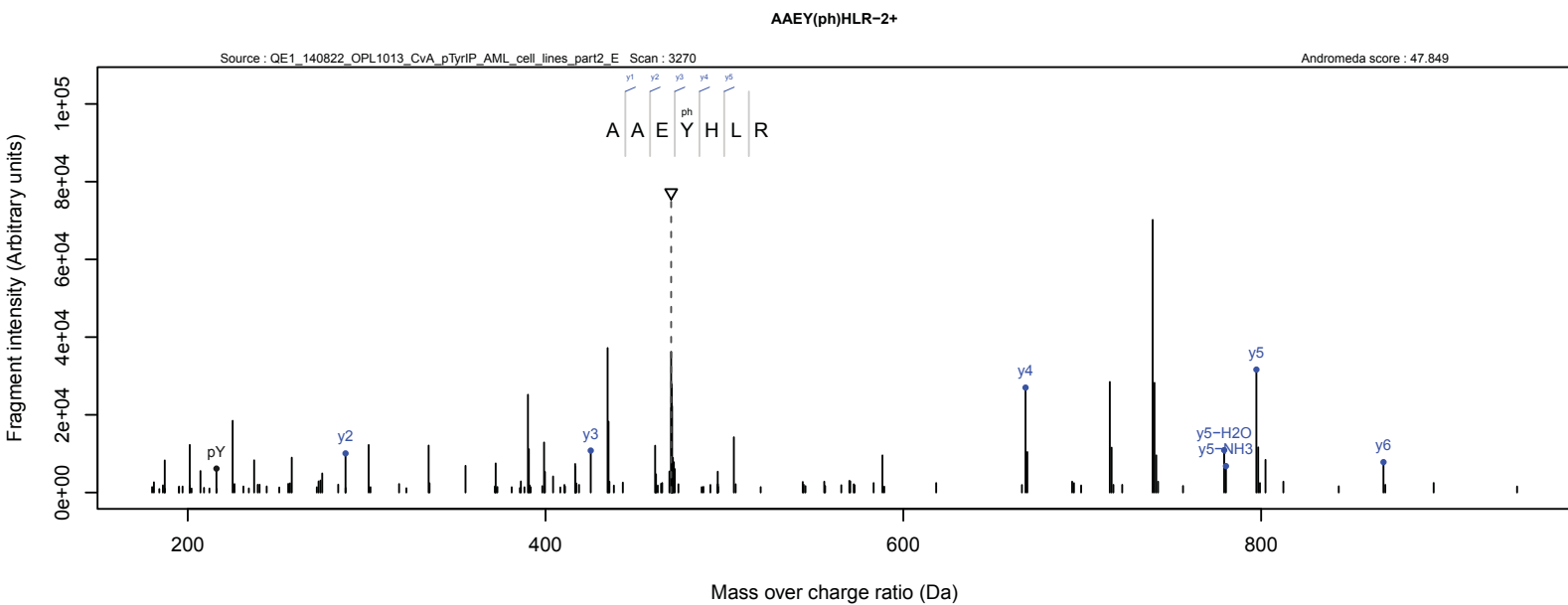
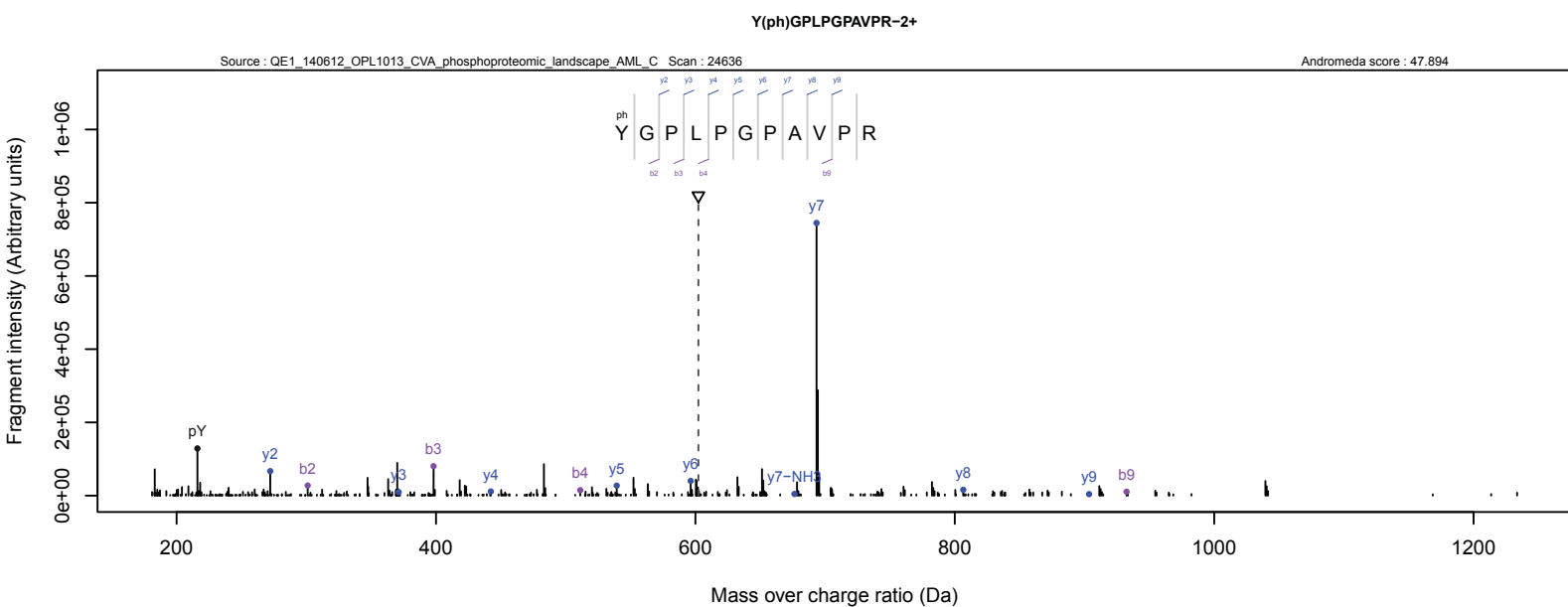
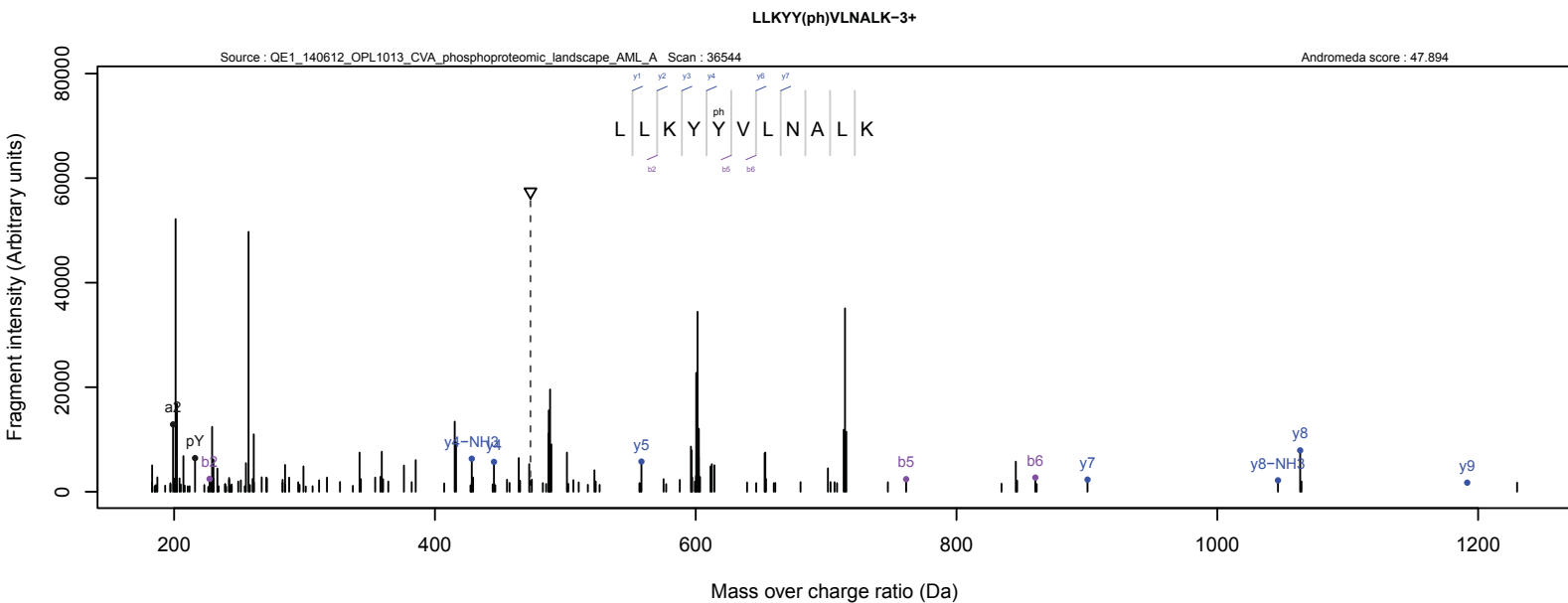


## AISSDMY(ph)FGR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 22691

Andromeda score: 47.894

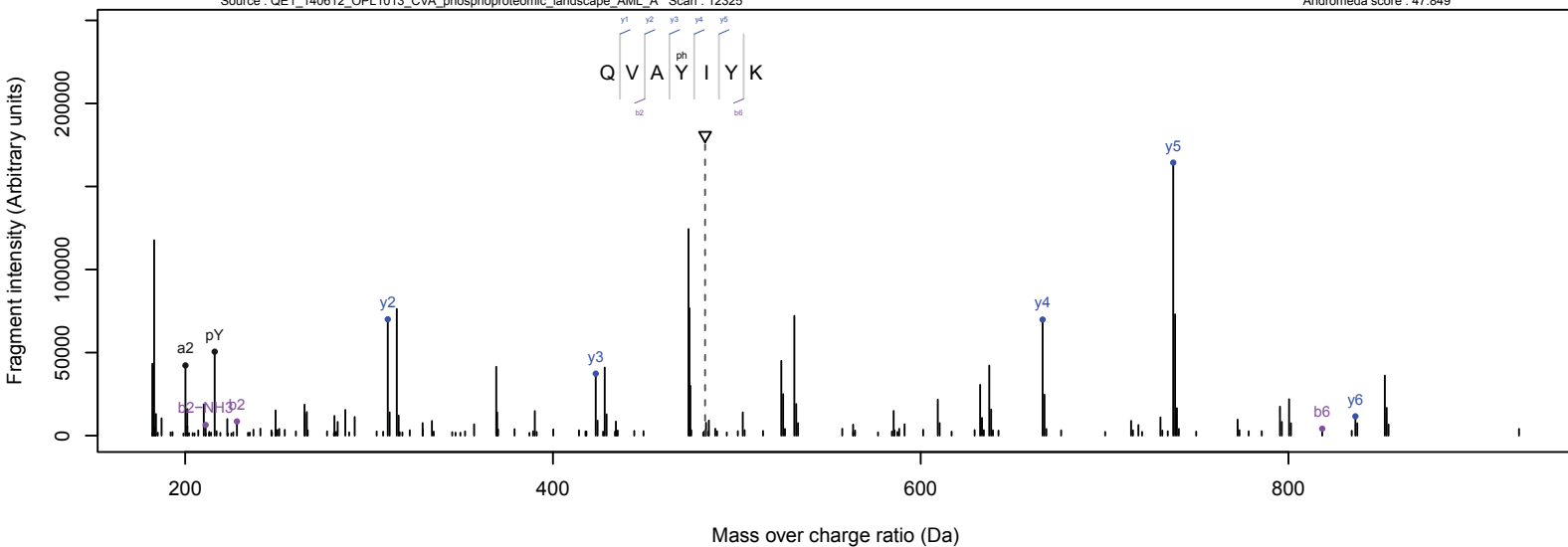




## QVAY(ph)IYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12325

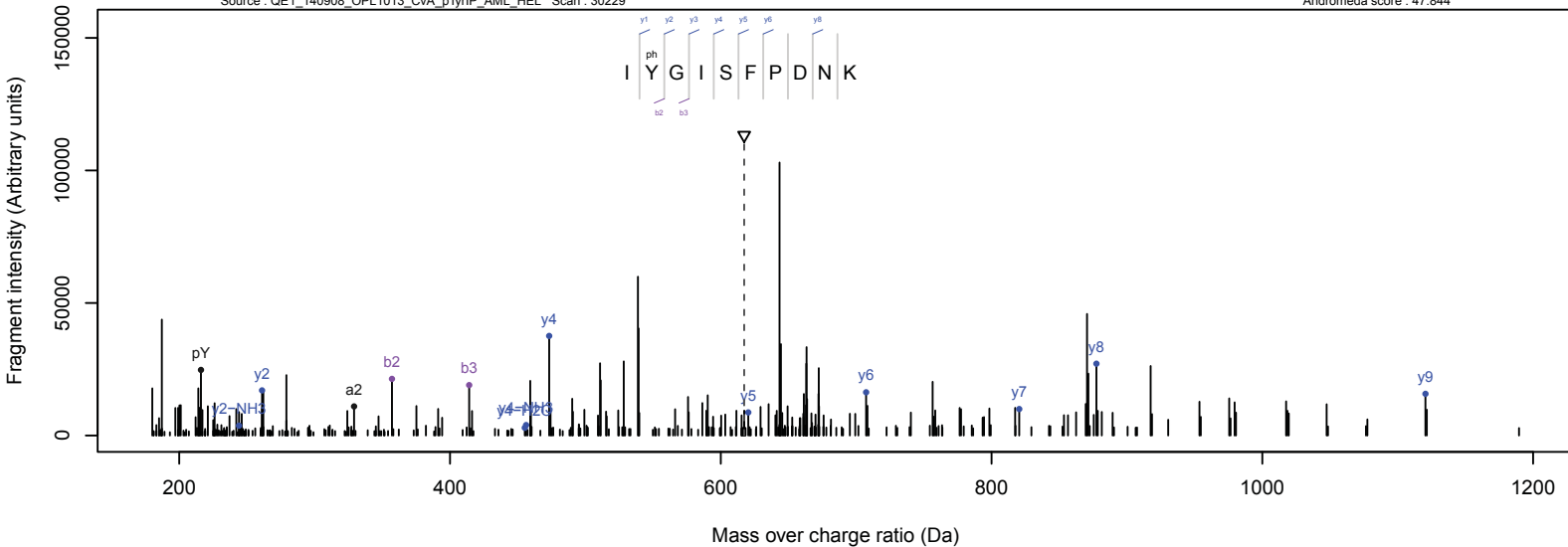
Andromeda score : 47.849



## IY(ph)GISFPDNK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 30229

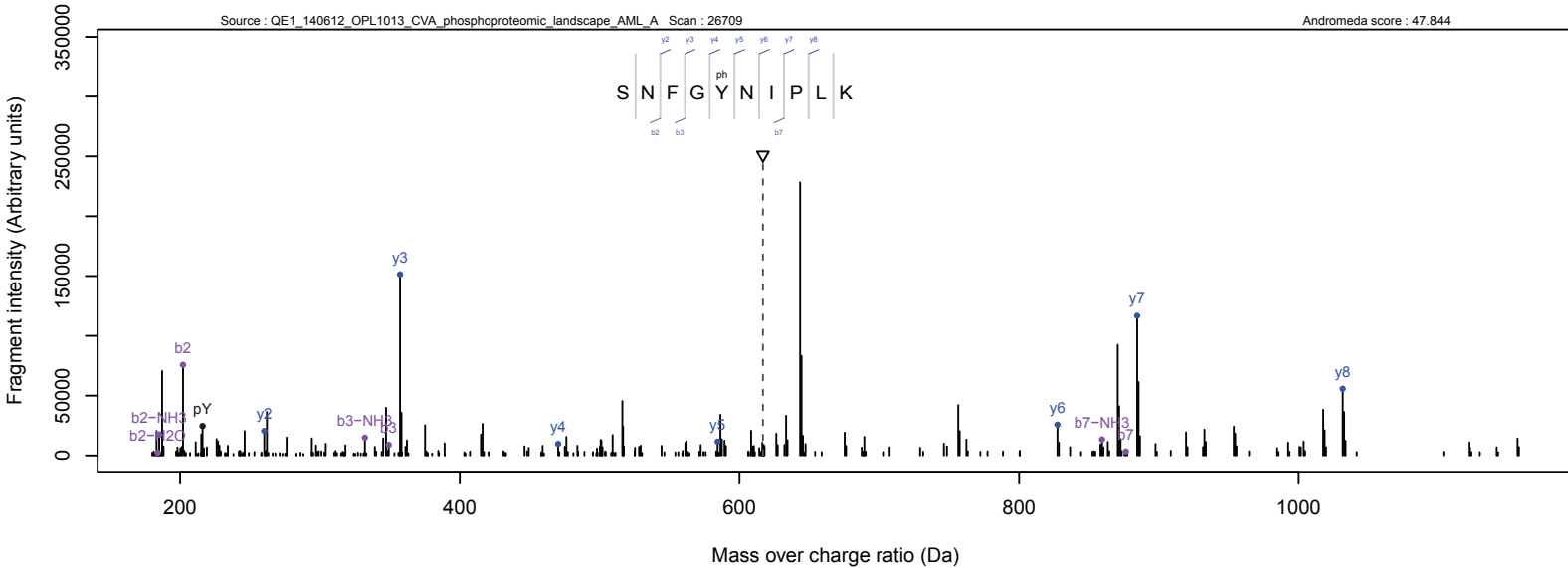
Andromeda score : 47.844

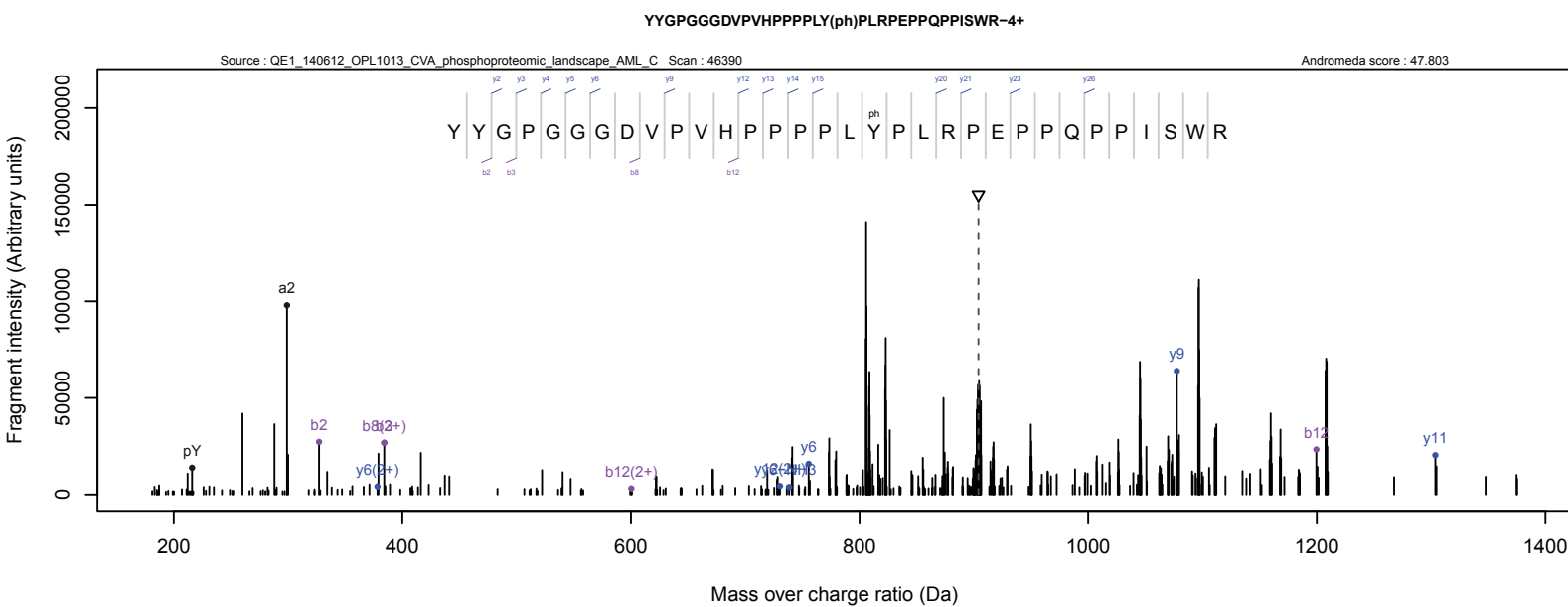
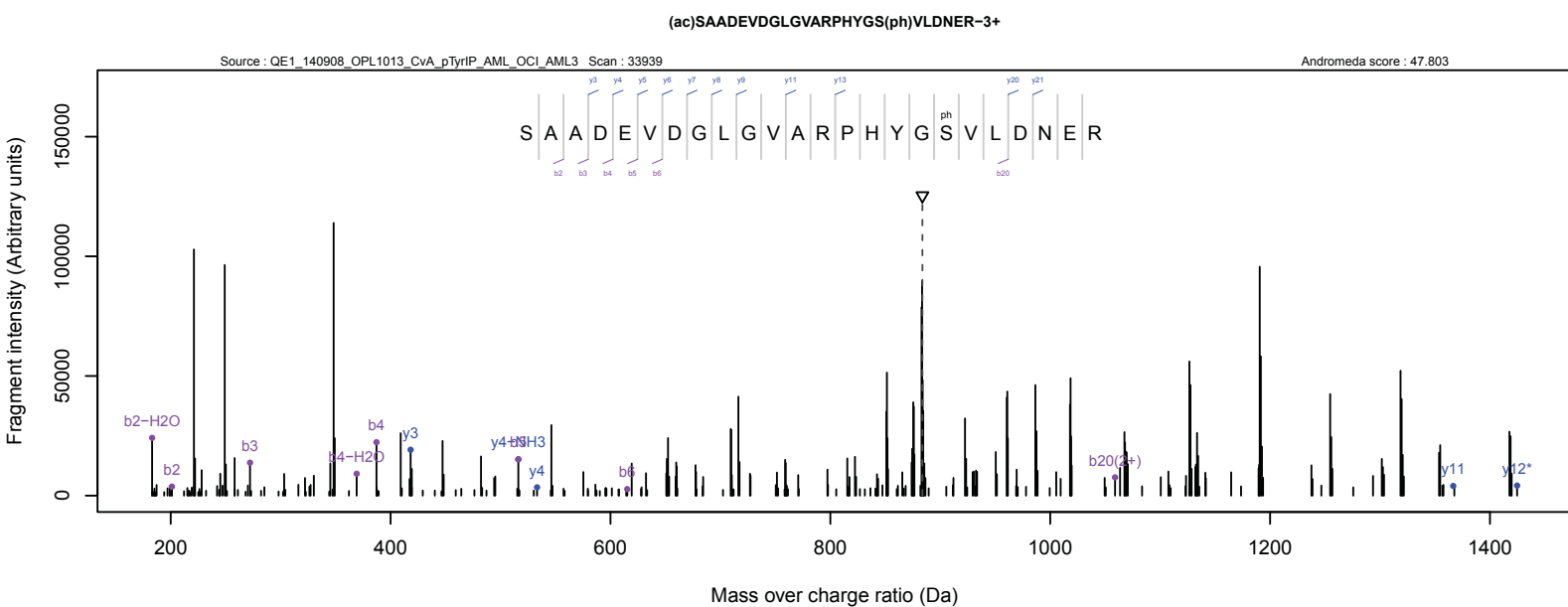
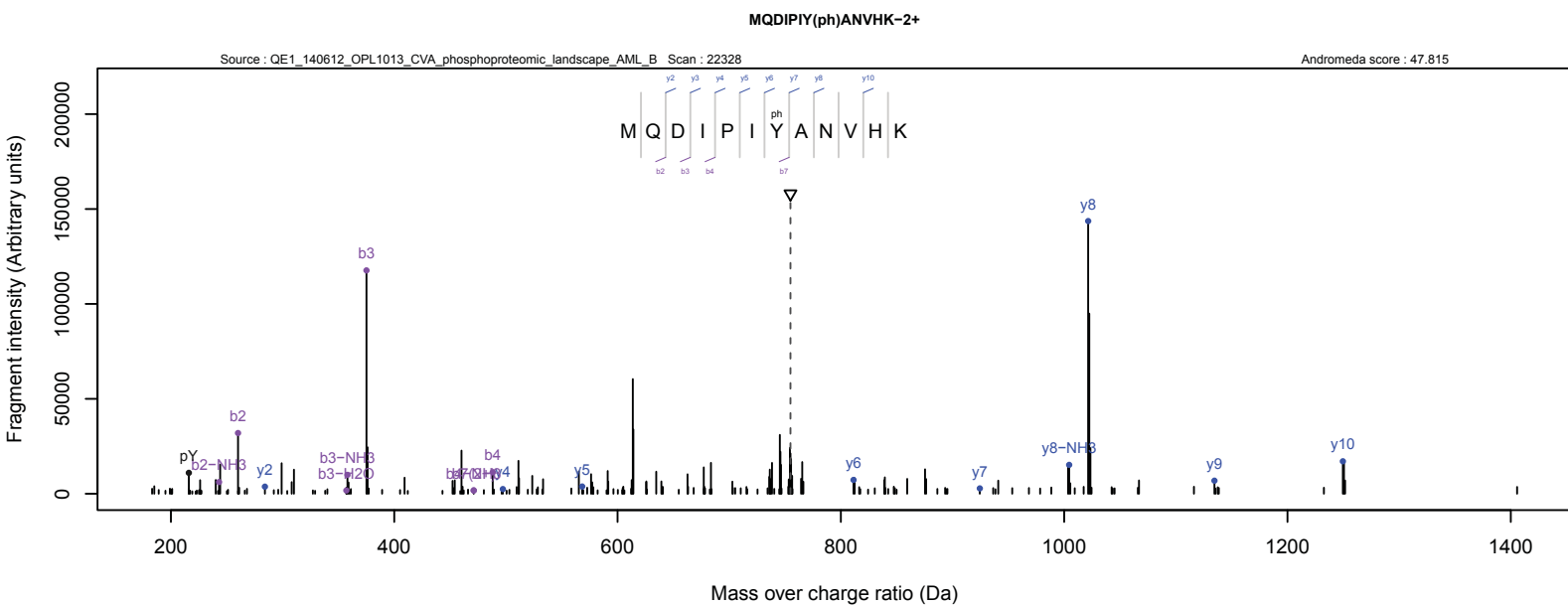


## SNFGY(ph)NIPLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 26709

Andromeda score : 47.844

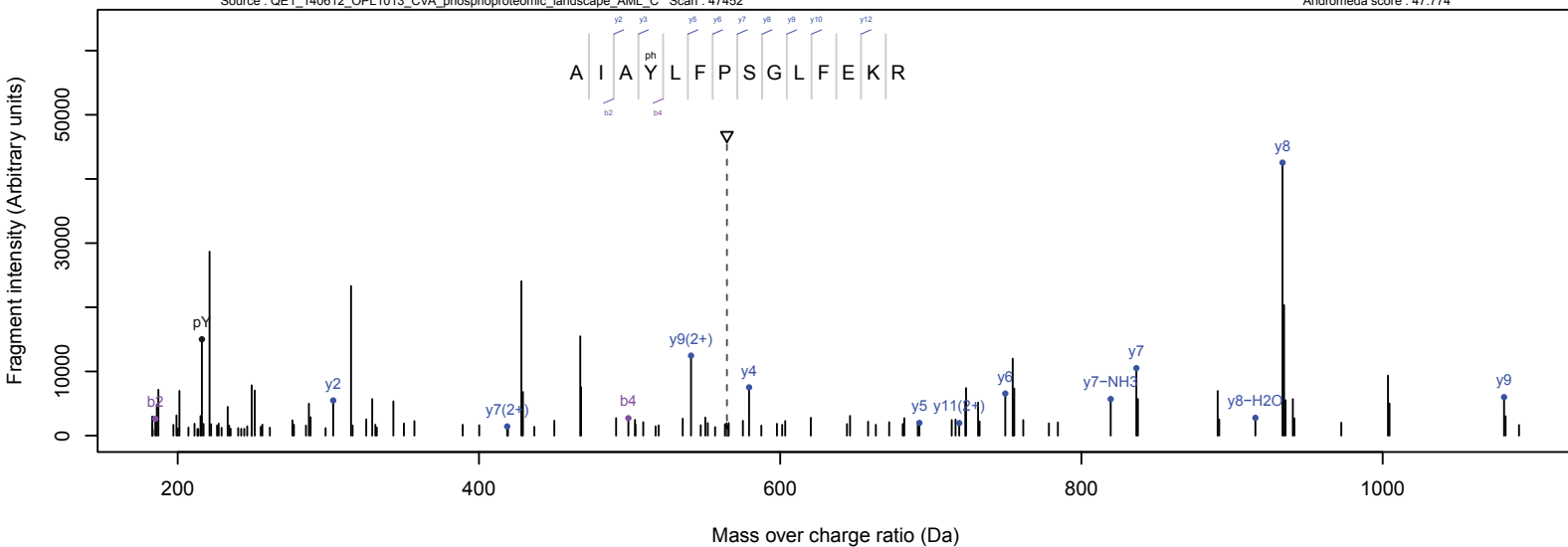




## AIAY(ph)LFPSGLFEKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47452

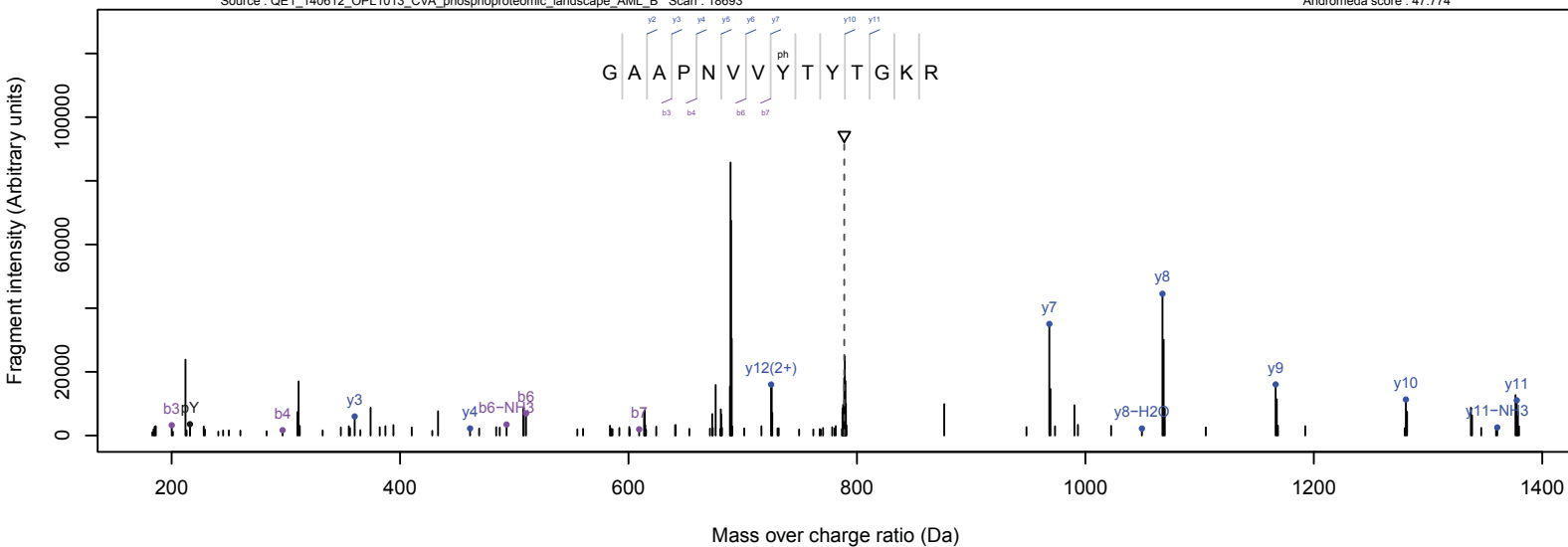
Andromeda score : 47.774



## GAAPNVVY(ph)TYTGKR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18693

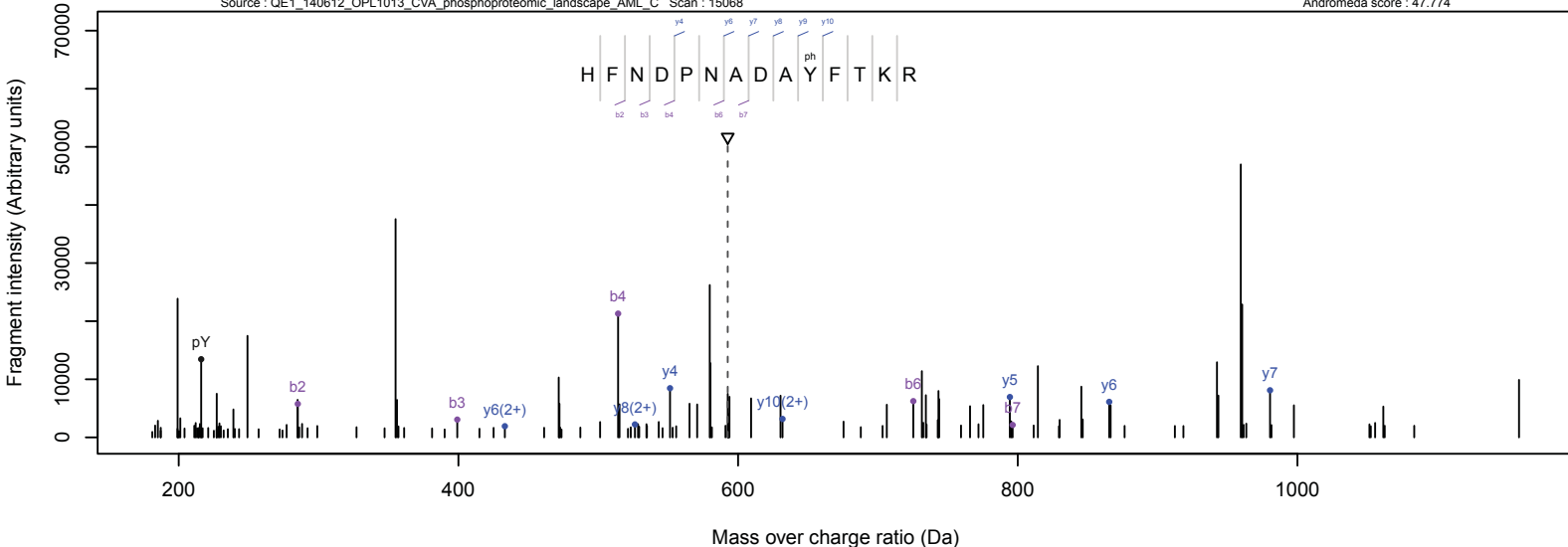
Andromeda score : 47.774



## HFNDPNA D A Y F T K R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15068

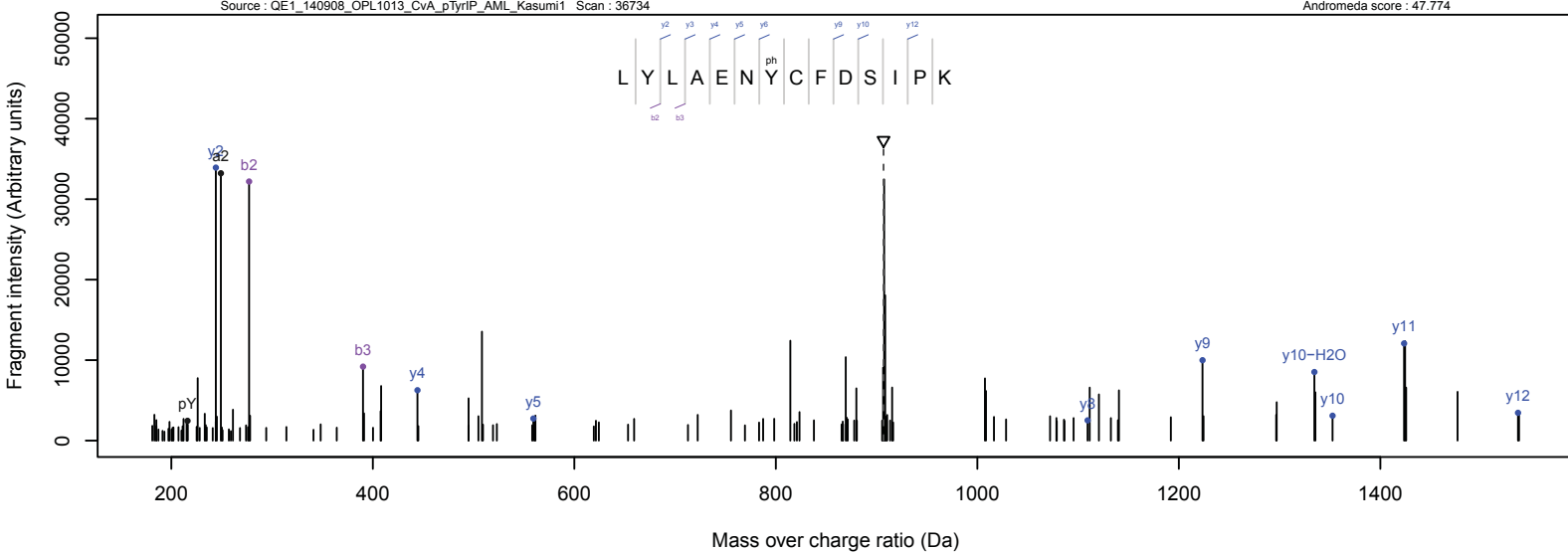
Andromeda score : 47.774



## LYLAENY(ph)CFDSIPK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 36734

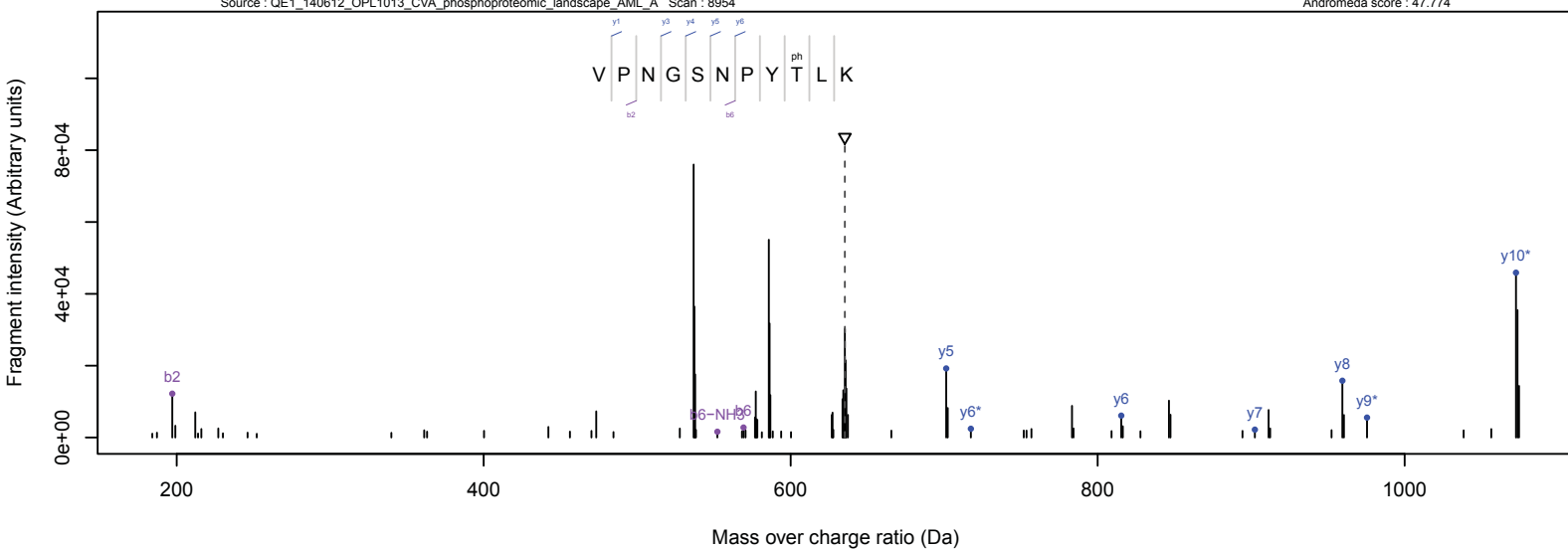
Andromeda score : 47.774



## VPNGSNPYT(ph)LK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 8954

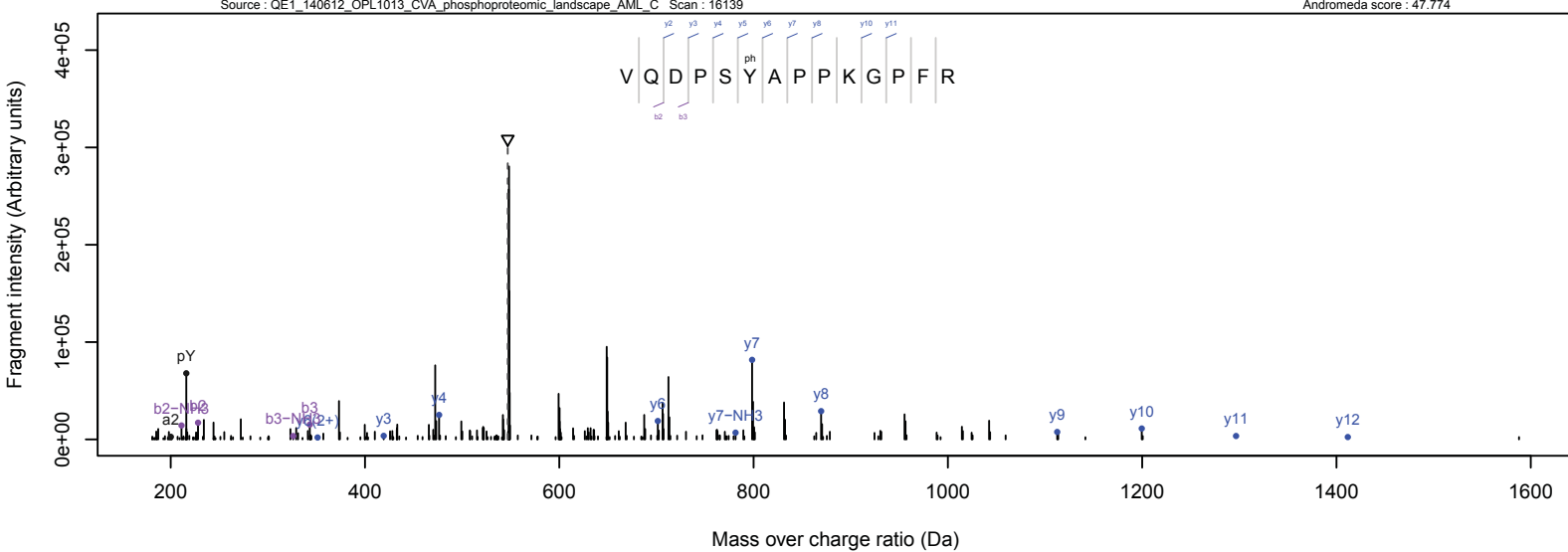
Andromeda score : 47.774



## VQDPSY(ph)APKGPFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16139

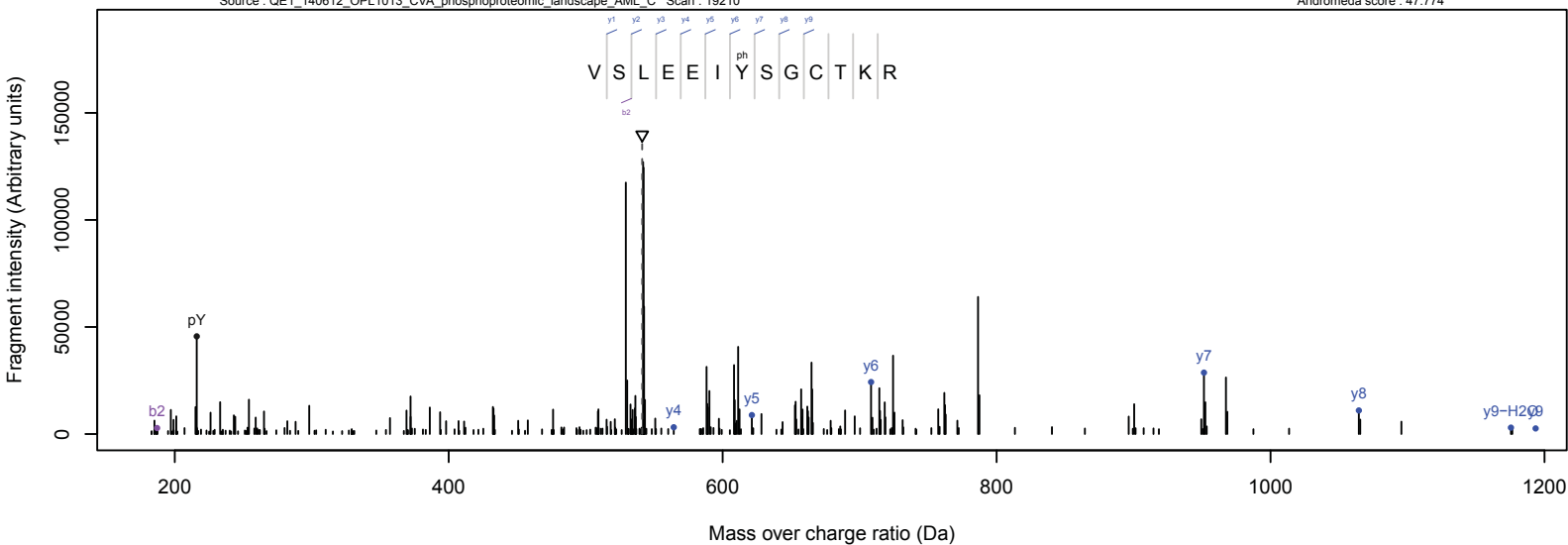
Andromeda score : 47.774



## VSLEEIY(ph)SGCTKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19210

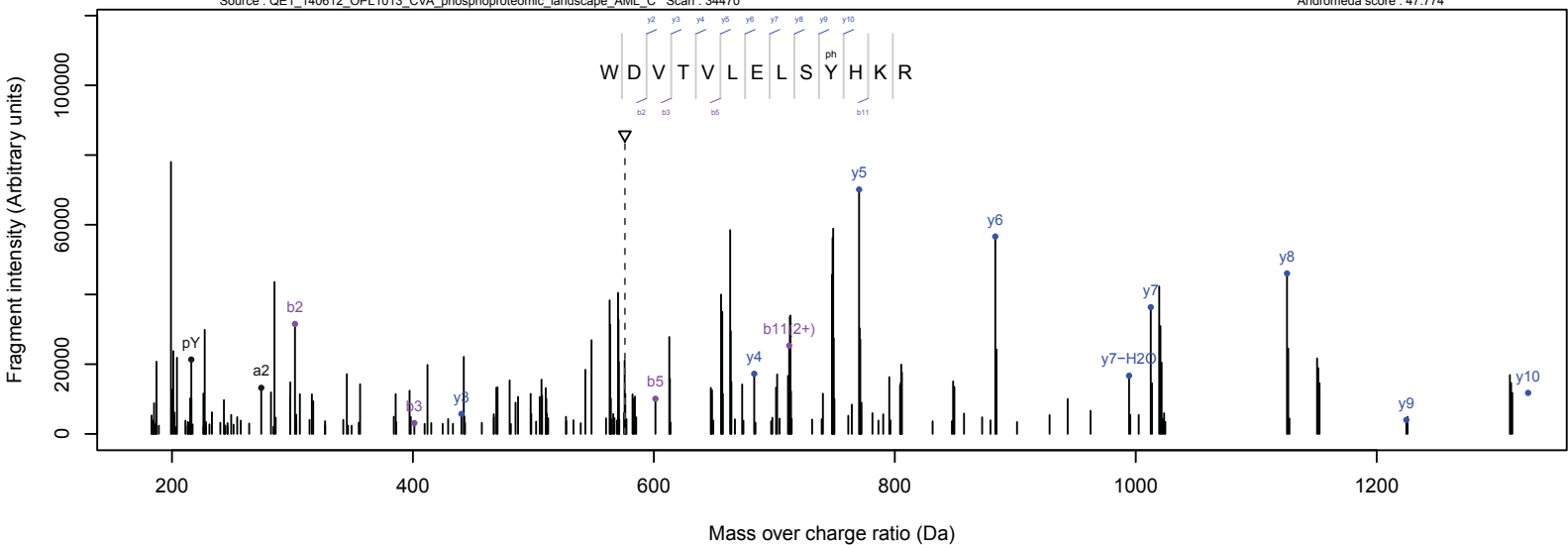
Andromeda score : 47.774



## WDVTVLELSY(ph)HKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34470

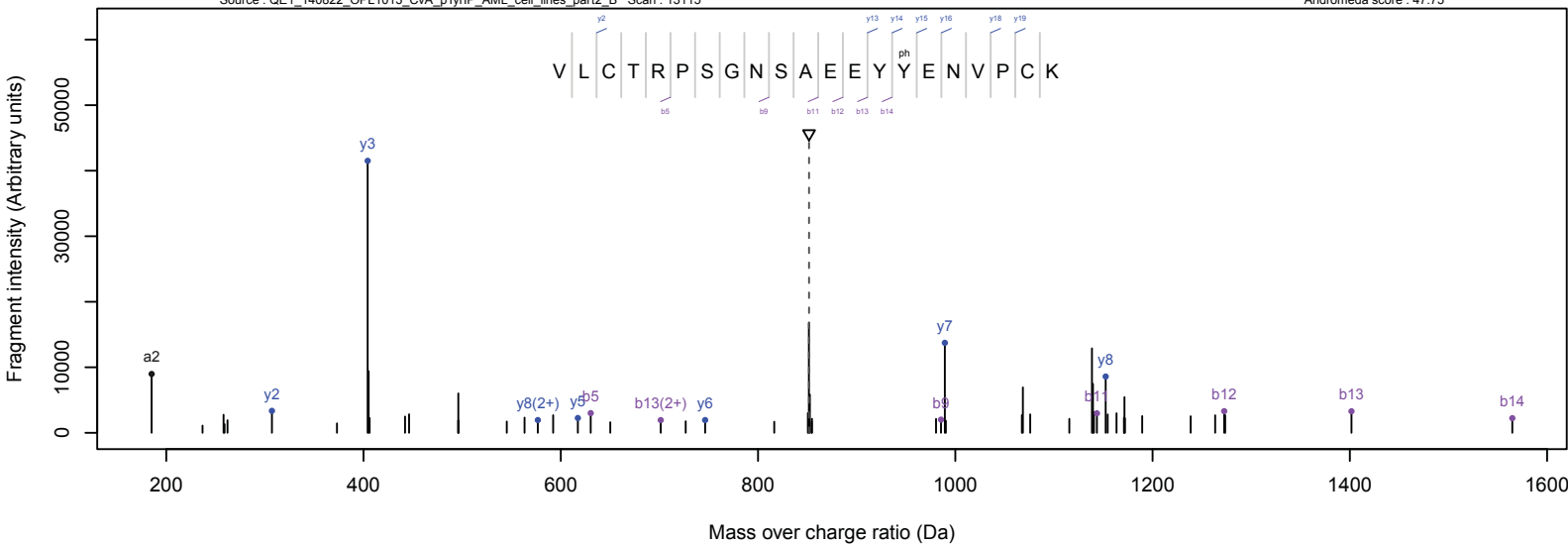
Andromeda score : 47.774



## VLCTRPSGNSAEEY(ph)ENVPCK-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 13115

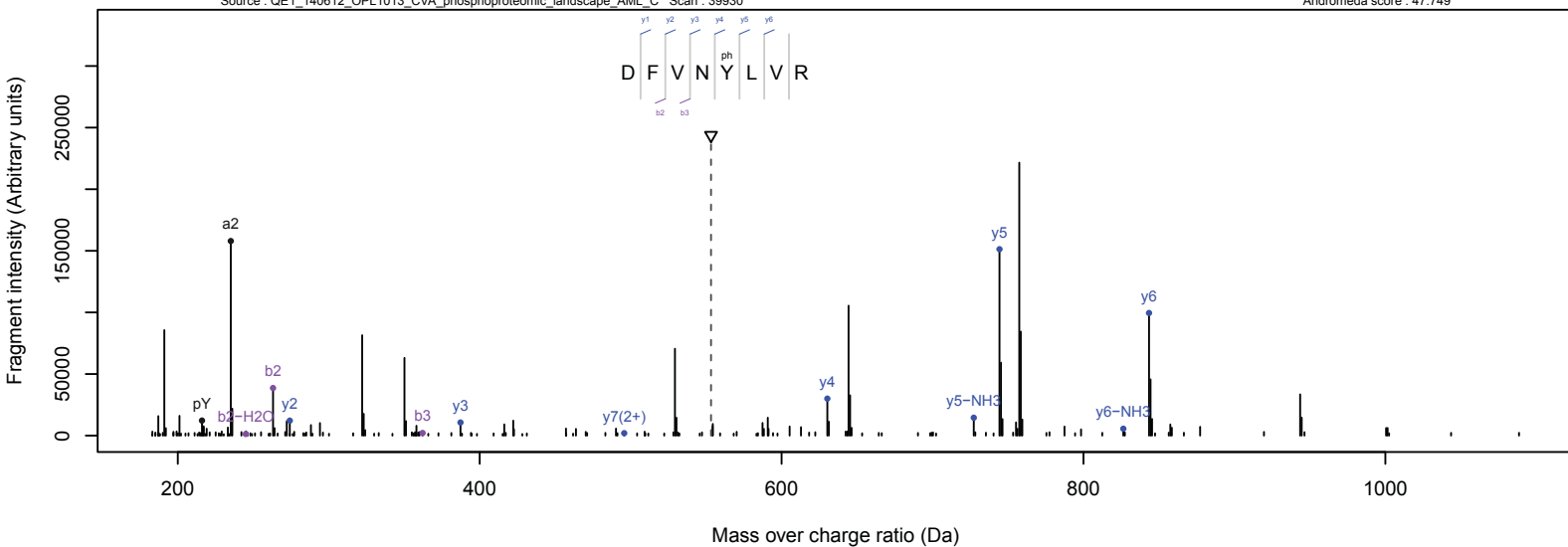
Andromeda score : 47.75



## DFVNY(ph)LVR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 39930

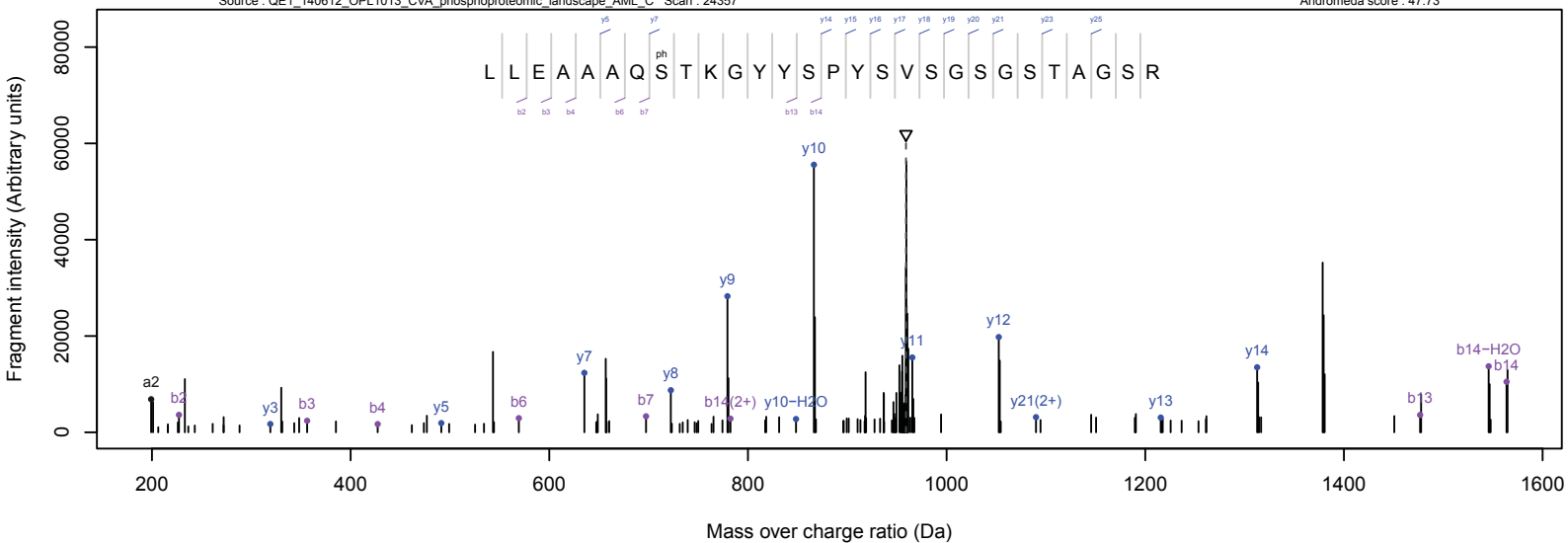
Andromeda score: 47.749



## LLEAAQ(S)TKGYYSYPVSGSGSTAGSR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24357

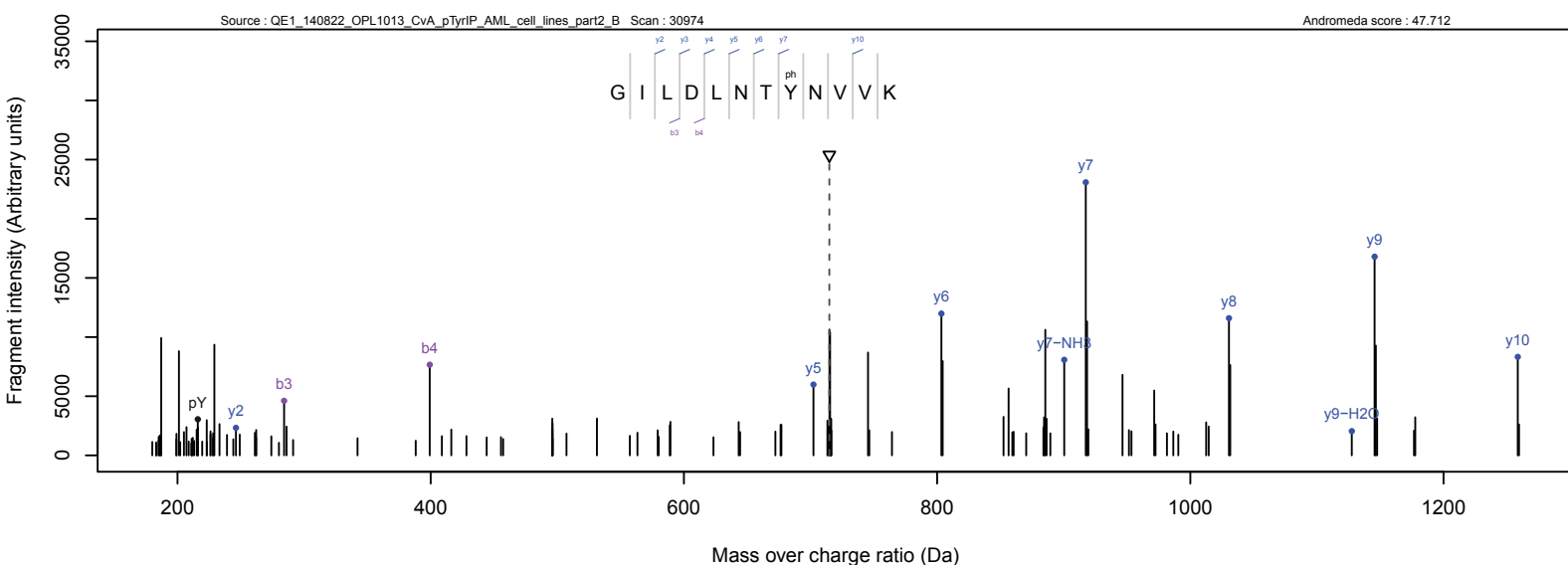
Andromeda score: 47.73



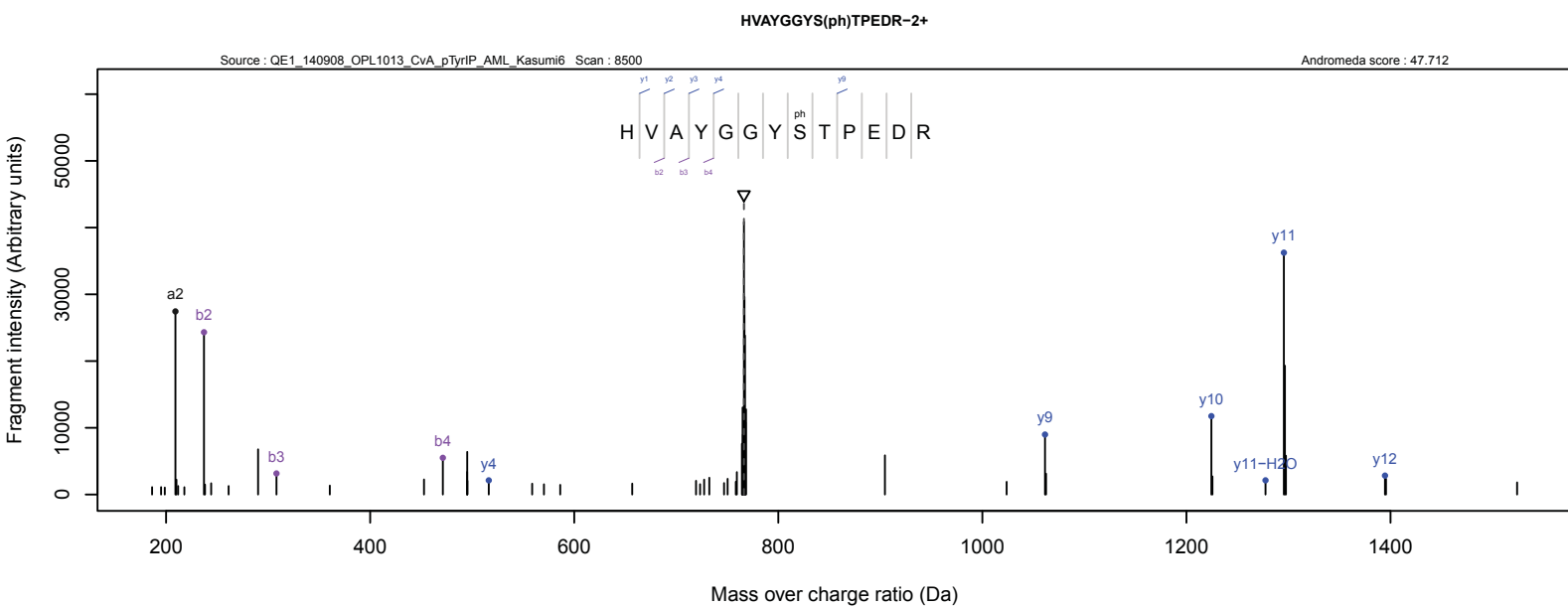
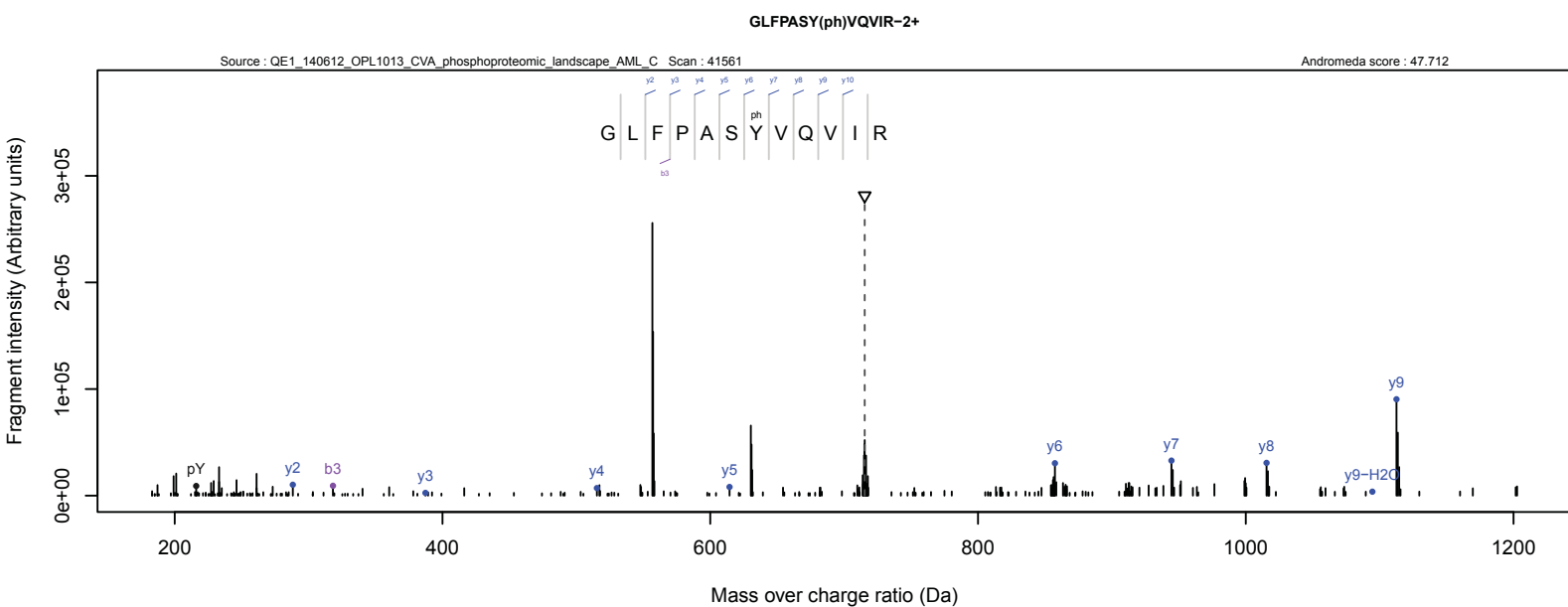
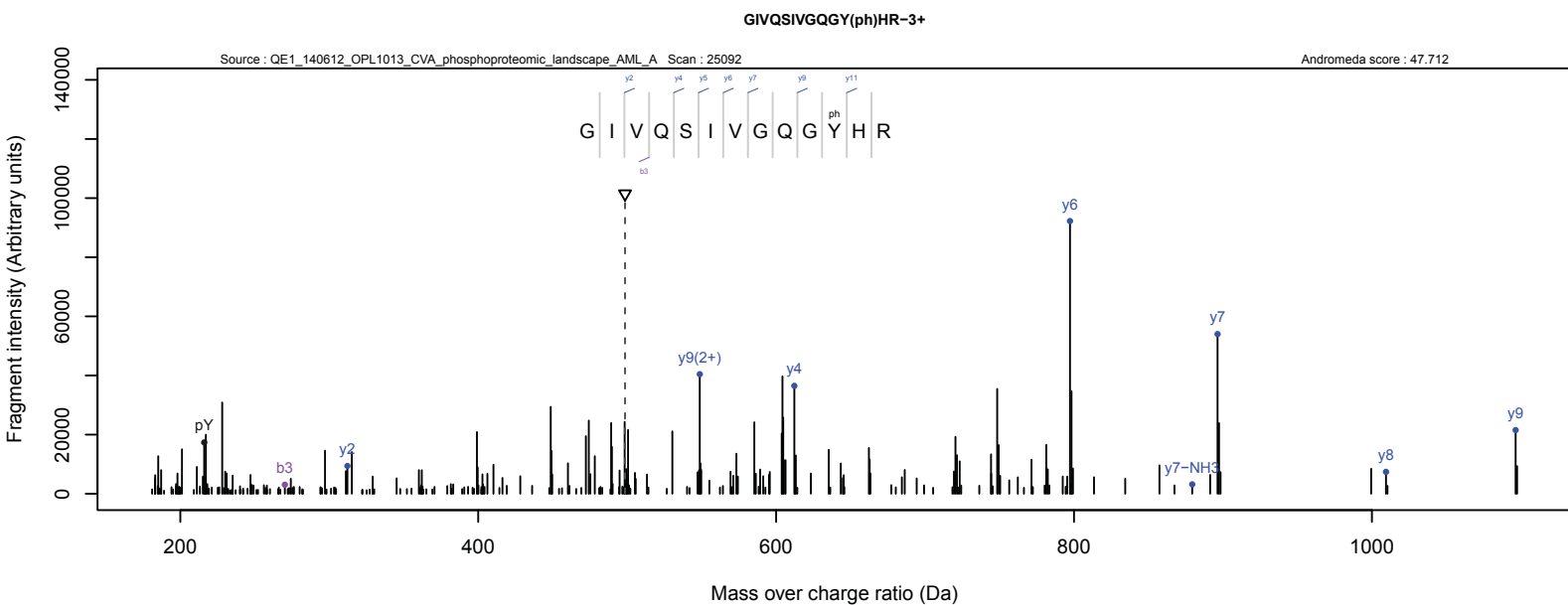
## GILDNTY(ph)NVVK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 30974

Andromeda score: 47.712



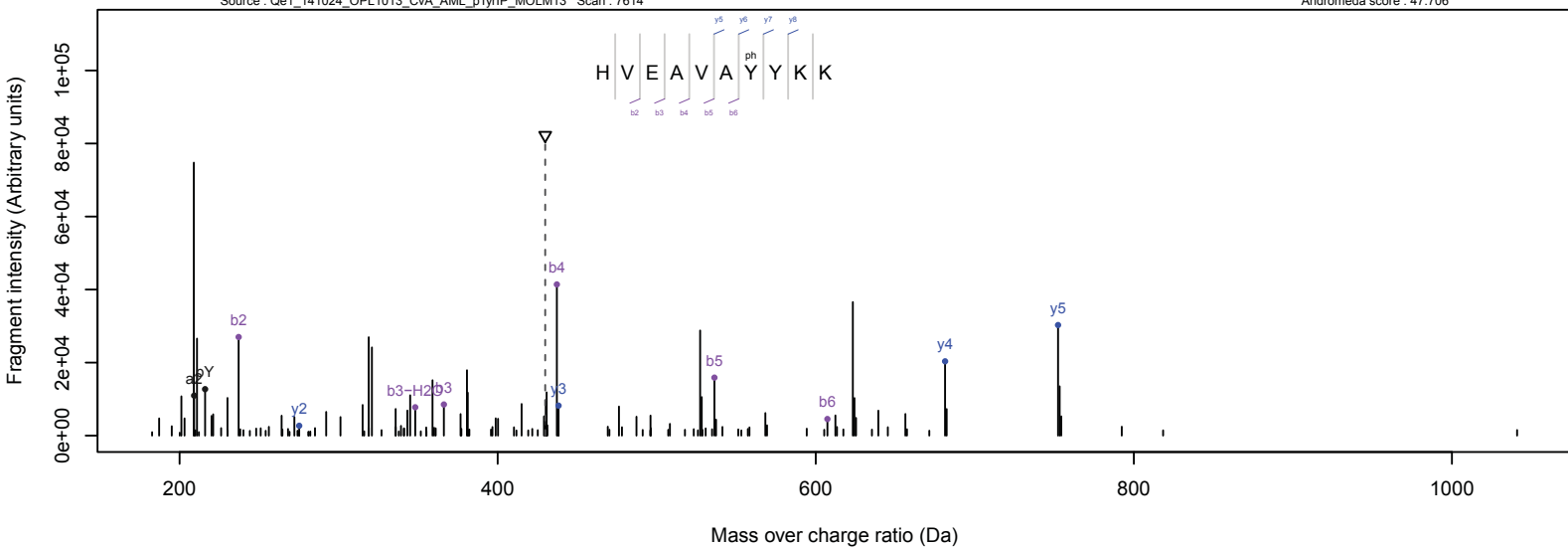




HVEAVAY(ph)YKK-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 7614

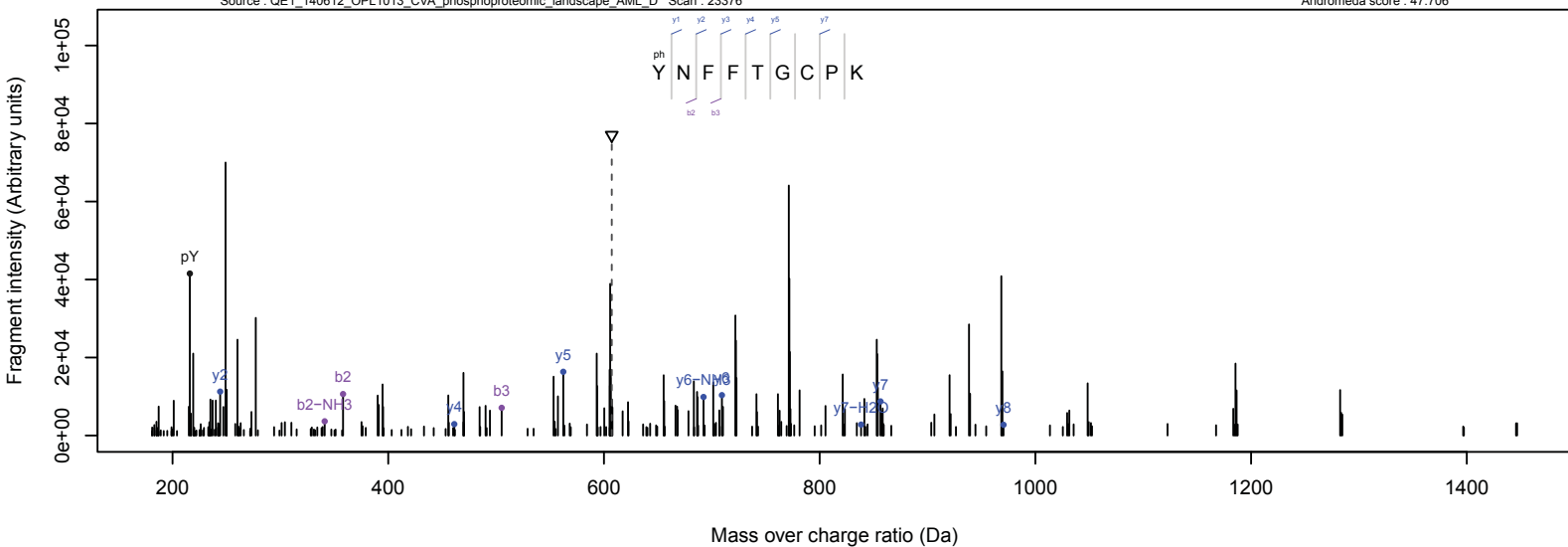
Andromeda score : 47.706



Y(ph)NFFTGPCPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23376

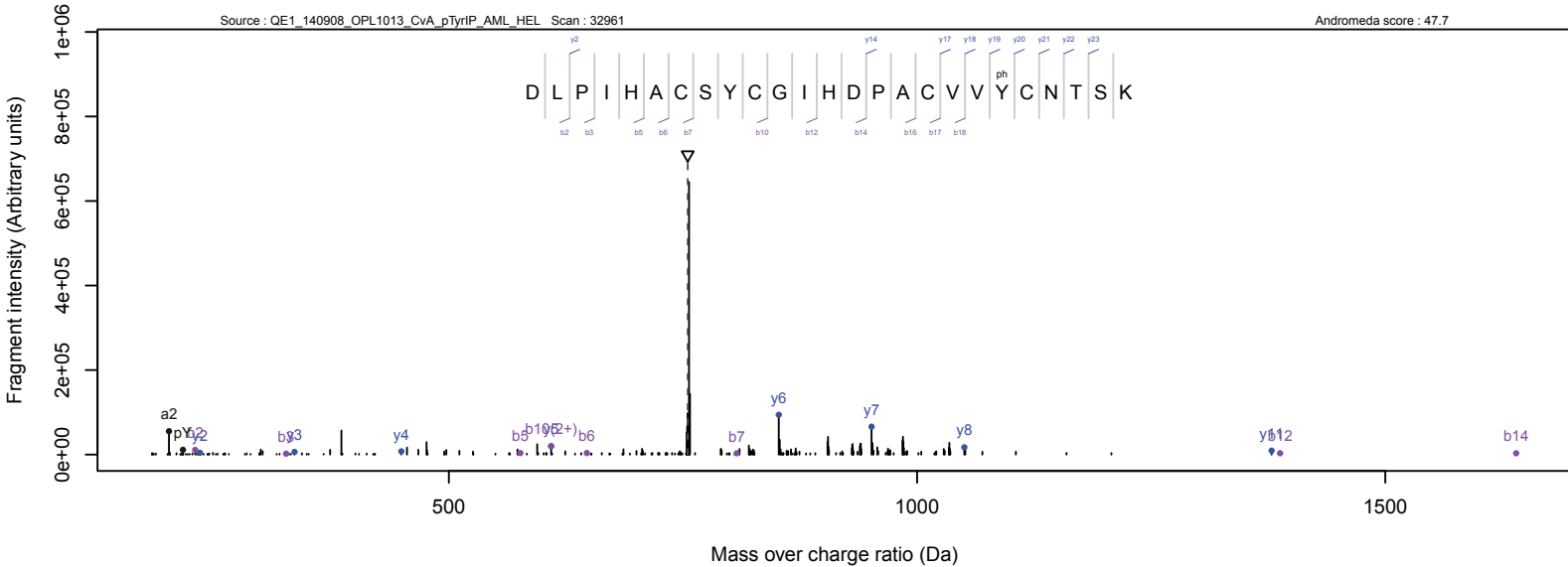
Andromeda score : 47.706



DLPIHACSYGIIHDPACVVY(ph)CNTSK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32961

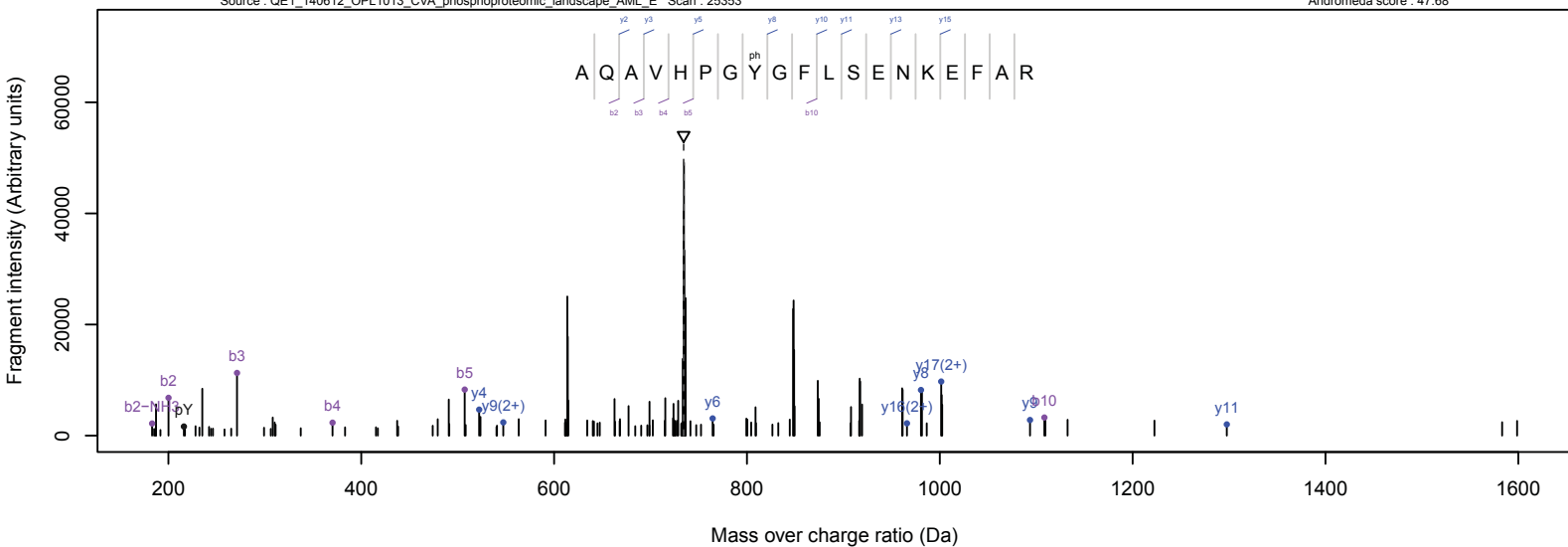
Andromeda score : 47.7



## AQAVHPG(ph)GFLSENKEFAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 25353

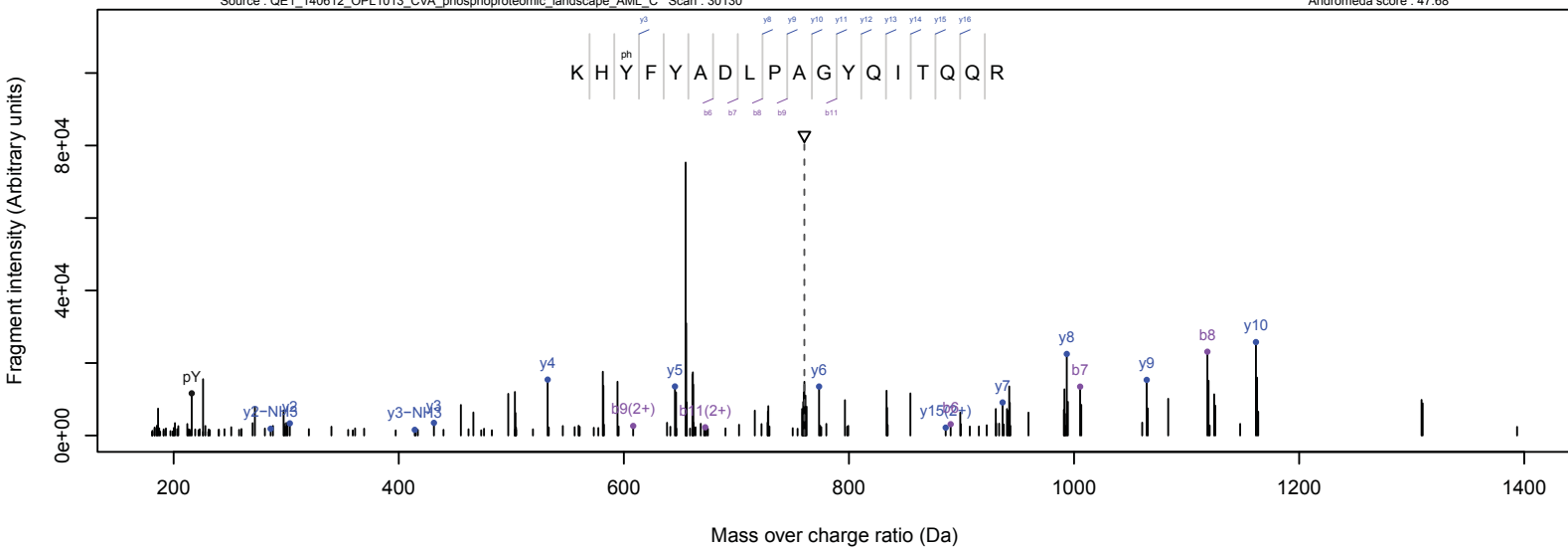
Andromeda score: 47.68



## KHYPYADLPAGYQITQQR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 30130

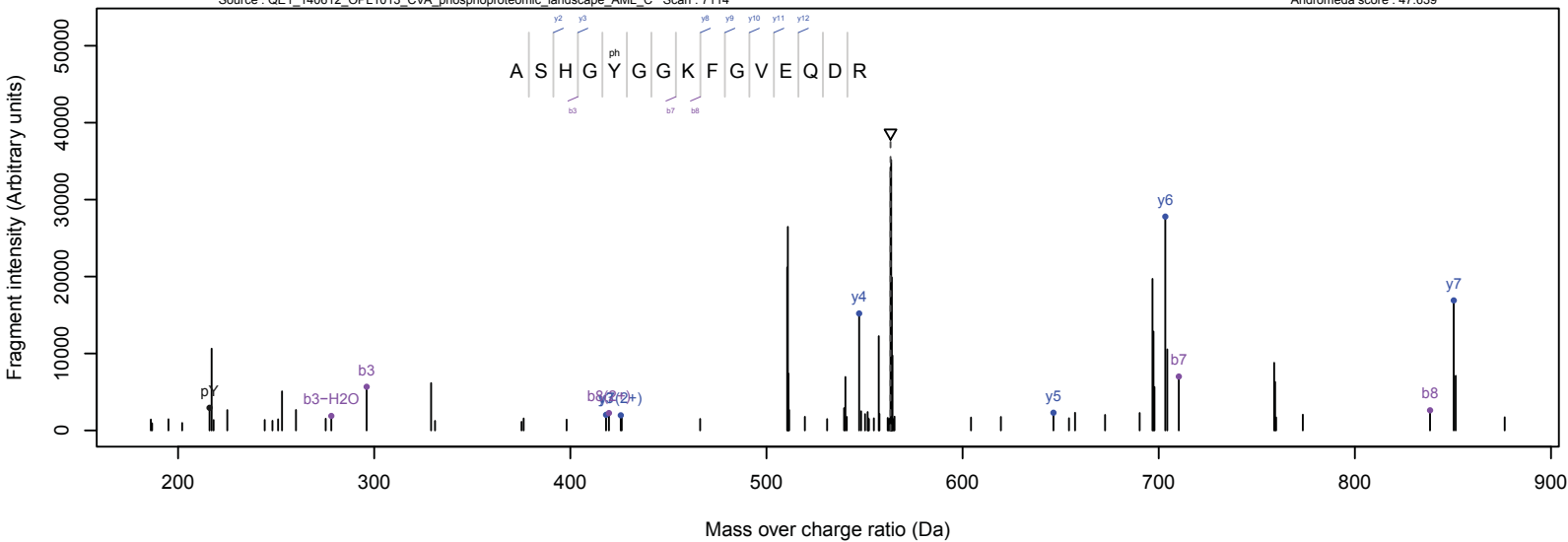
Andromeda score: 47.68



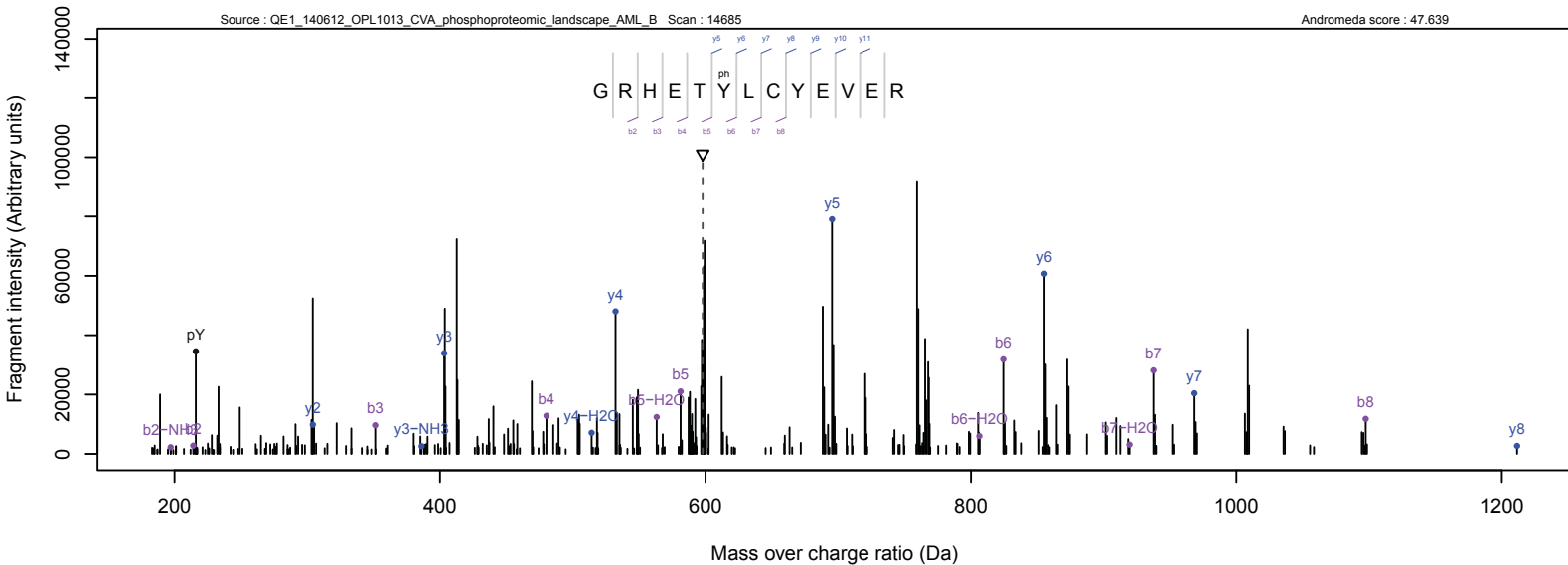
## ASHGY(ph)GGKFGVEQDR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 7114

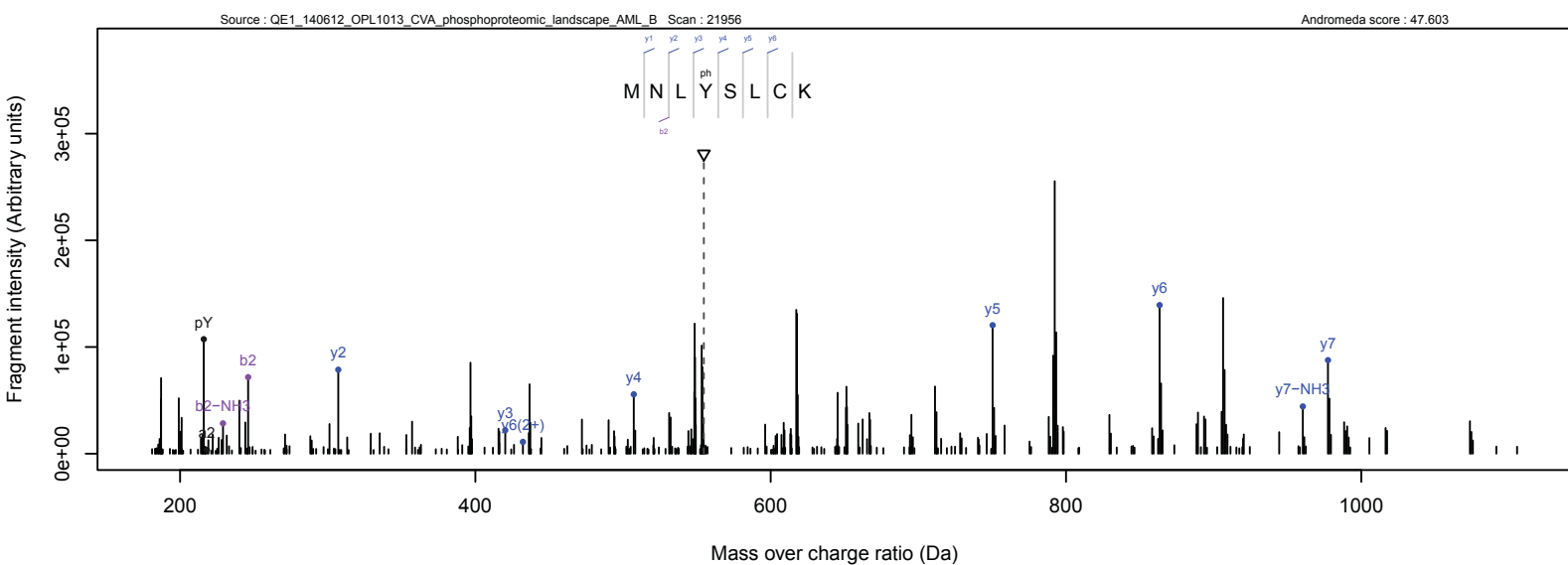
Andromeda score: 47.639



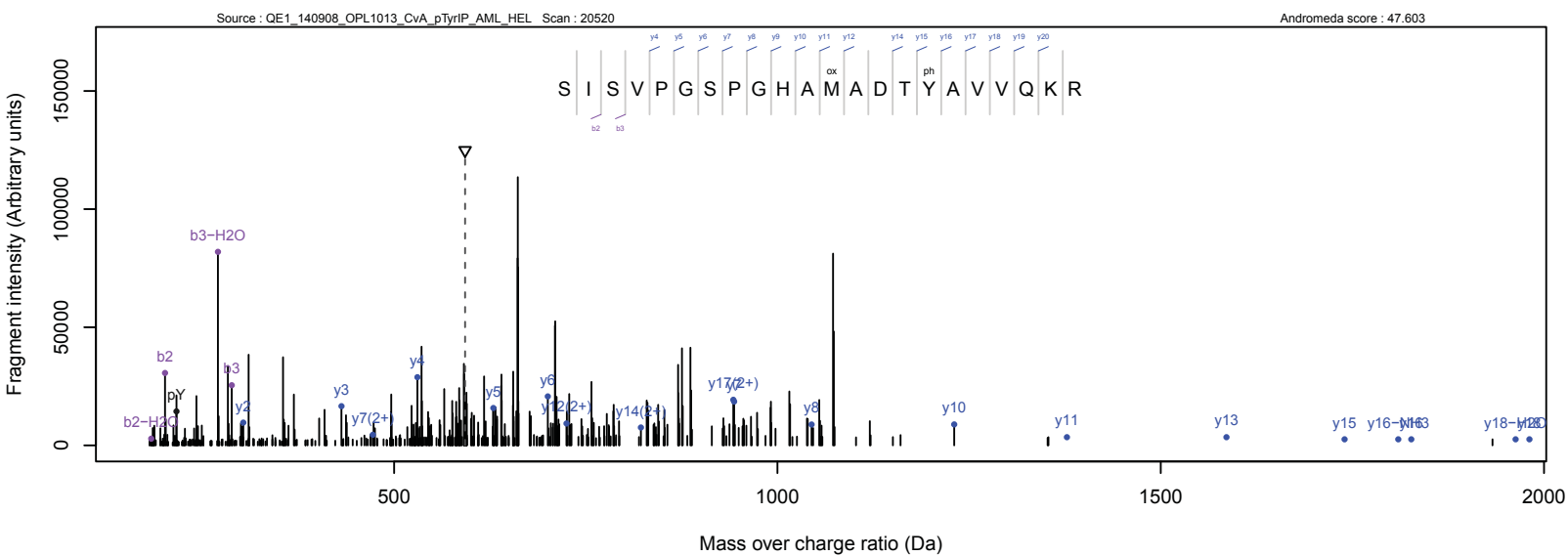
GRHETY(ph)LCYEVEVER-3+



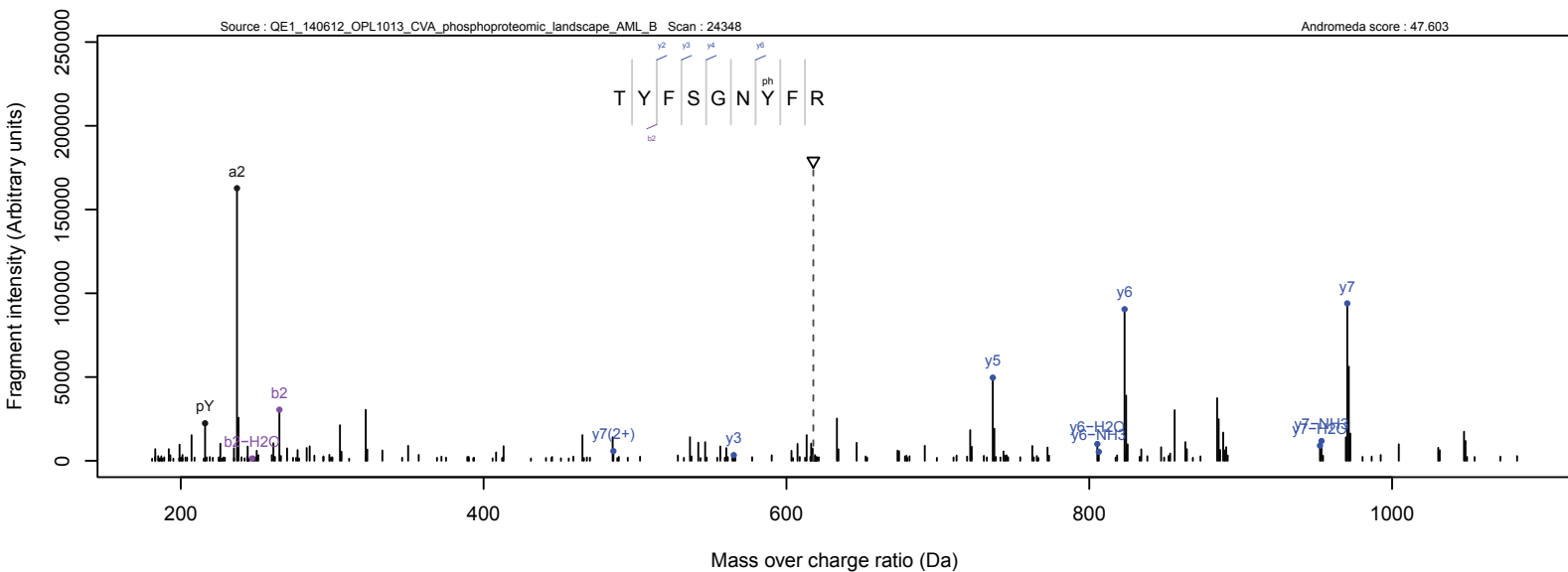
MNLY(ph)SLCK-2+



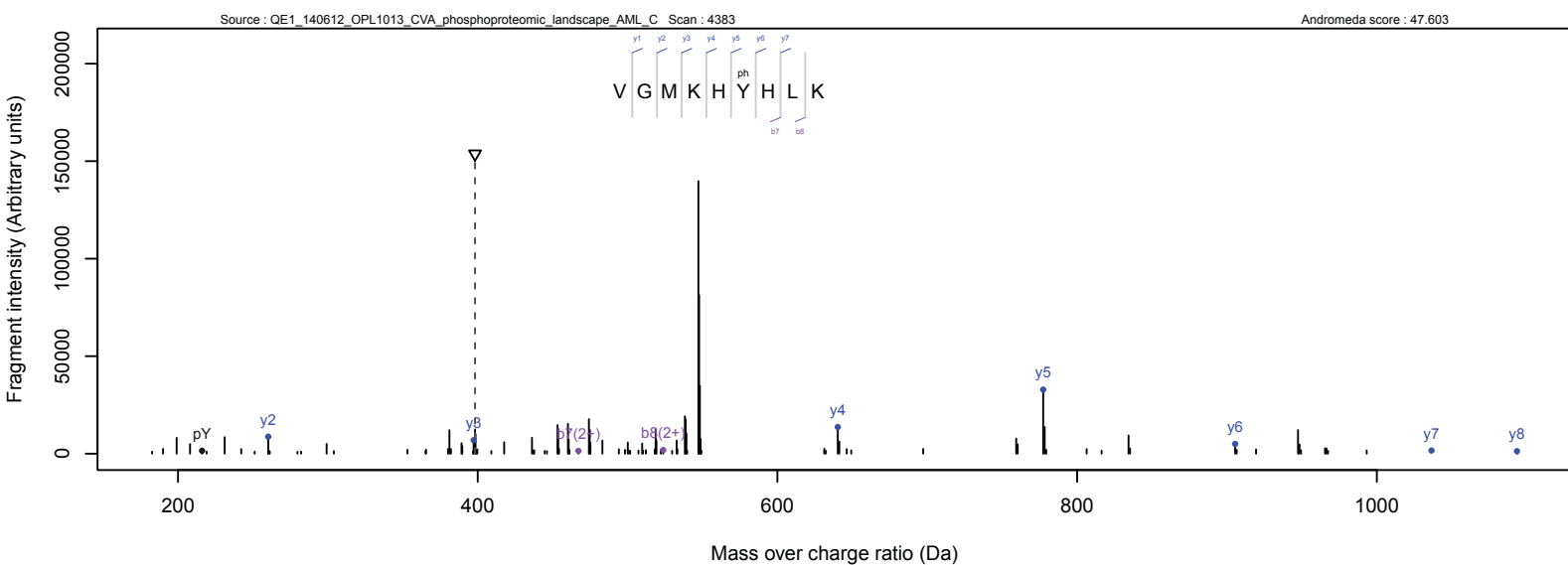
SISVPGSPGHAM(ox)ADTY(ph)AVVQKR-4+



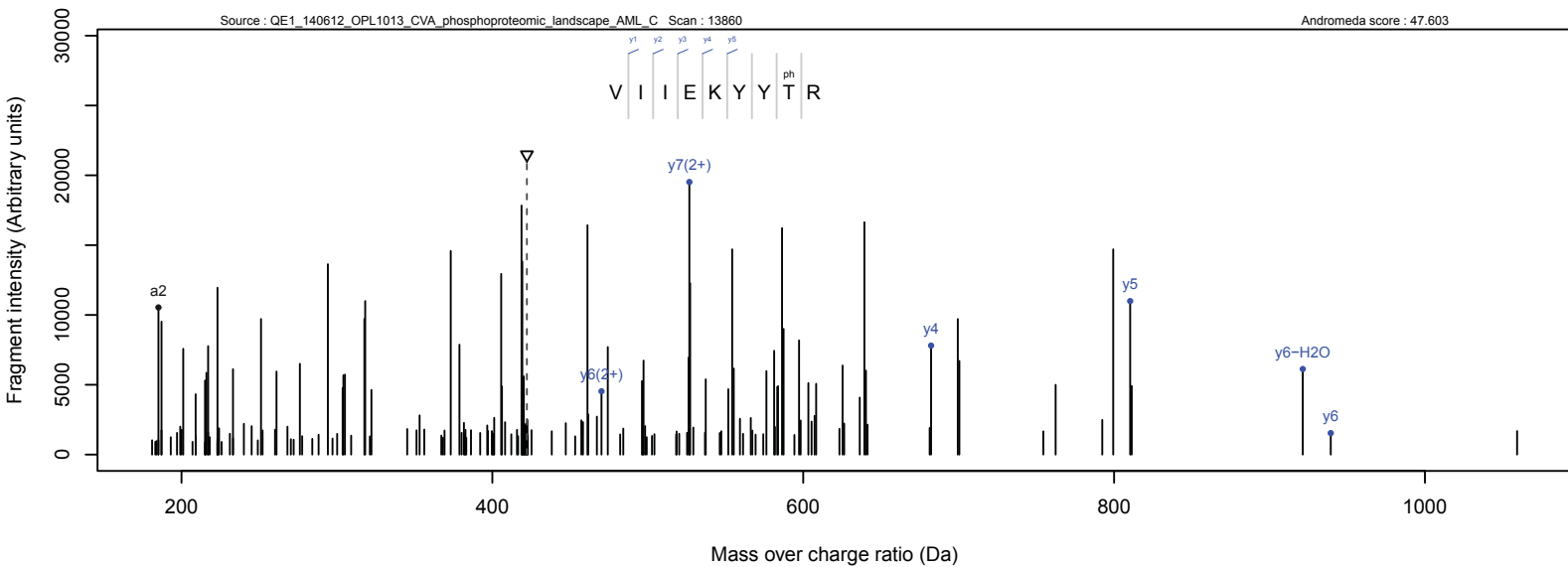
## TYFSGNY(ph)FR-2+



## VGMKHY(ph)HLK-3+



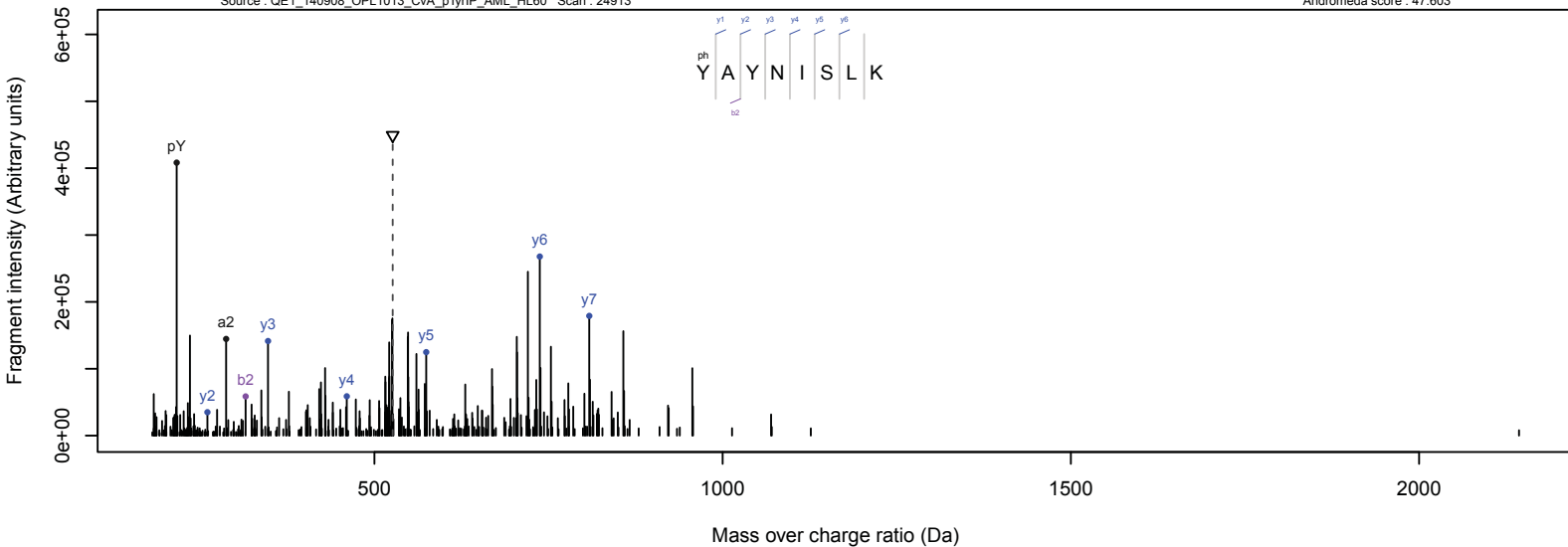
## VIIEKYT(ph)R-3+



Y(ph)AYNISLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 24913

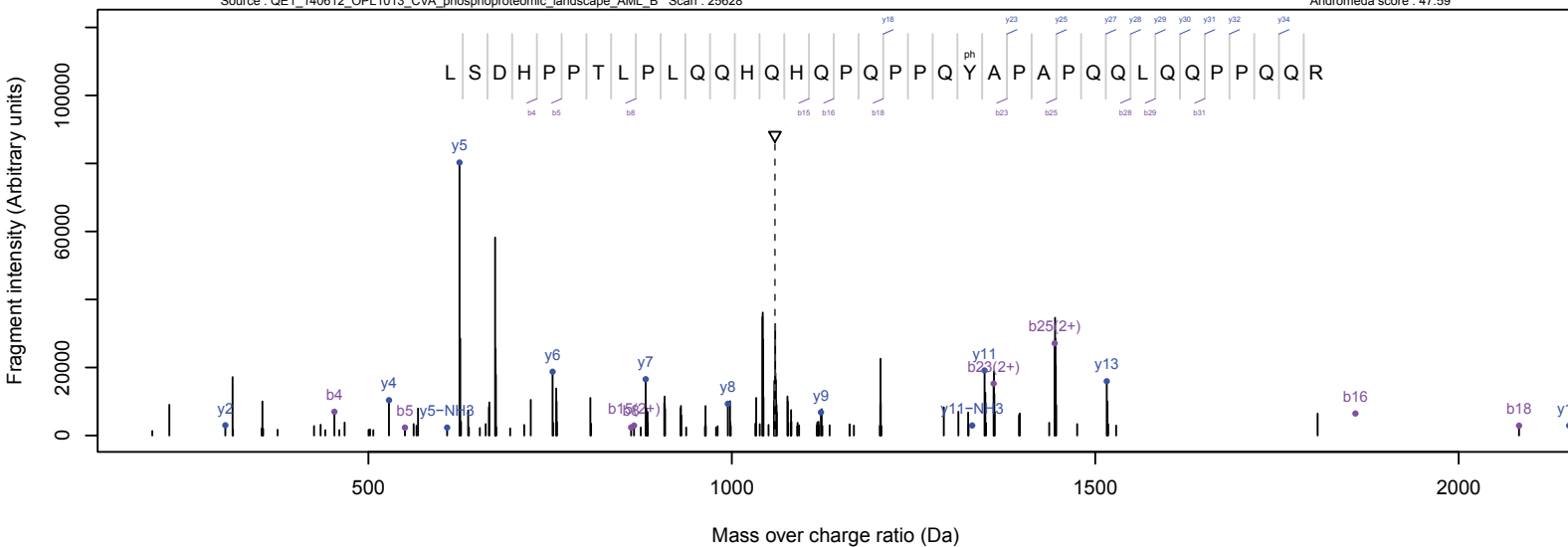
Andromeda score : 47.603



LSDHPPTLPLQQHQHPQPQY(ph)APAPQLQPPQQR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25628

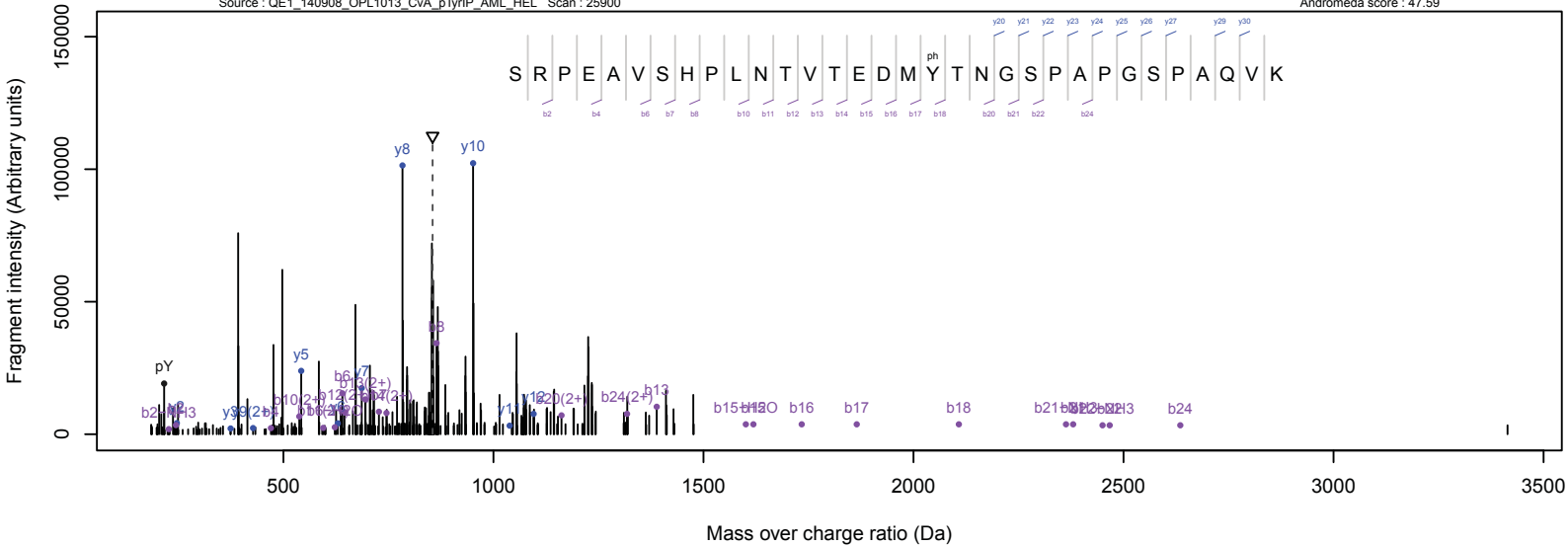
Andromeda score : 47.59



SRPEAVSHPLNTVTEDMY(ph)TNGSPAPGSPAQVK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 25900

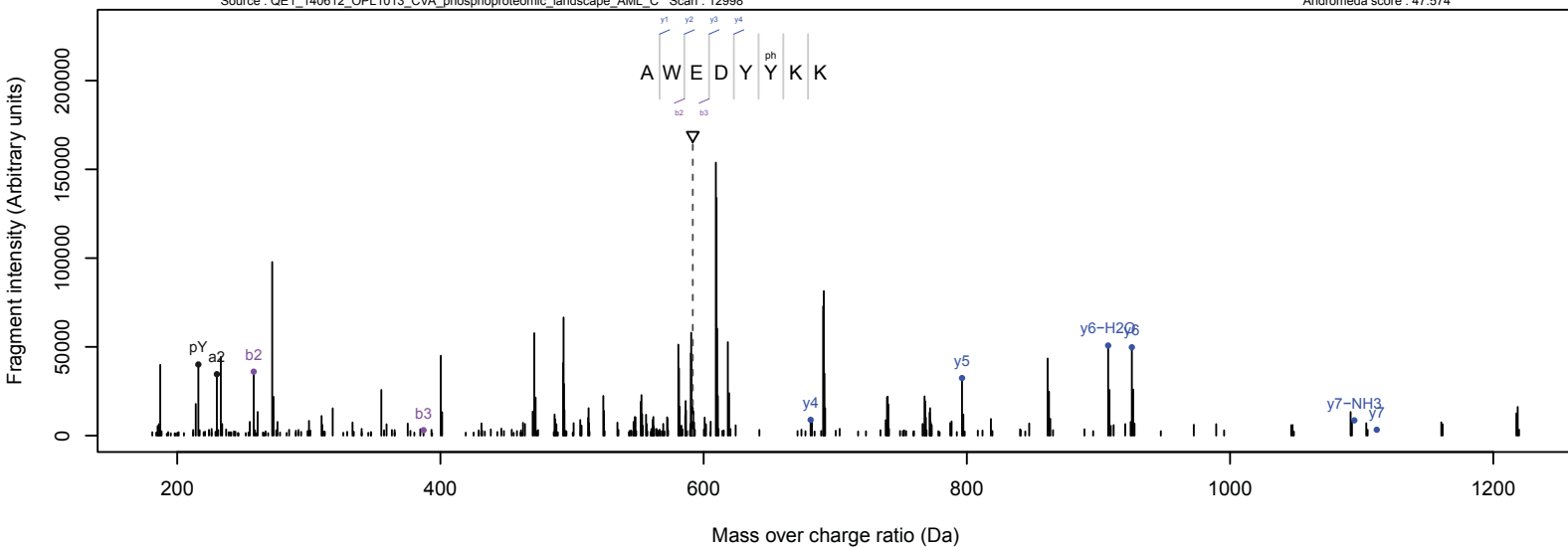
Andromeda score : 47.59



AWEDYY(ph)KK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12998

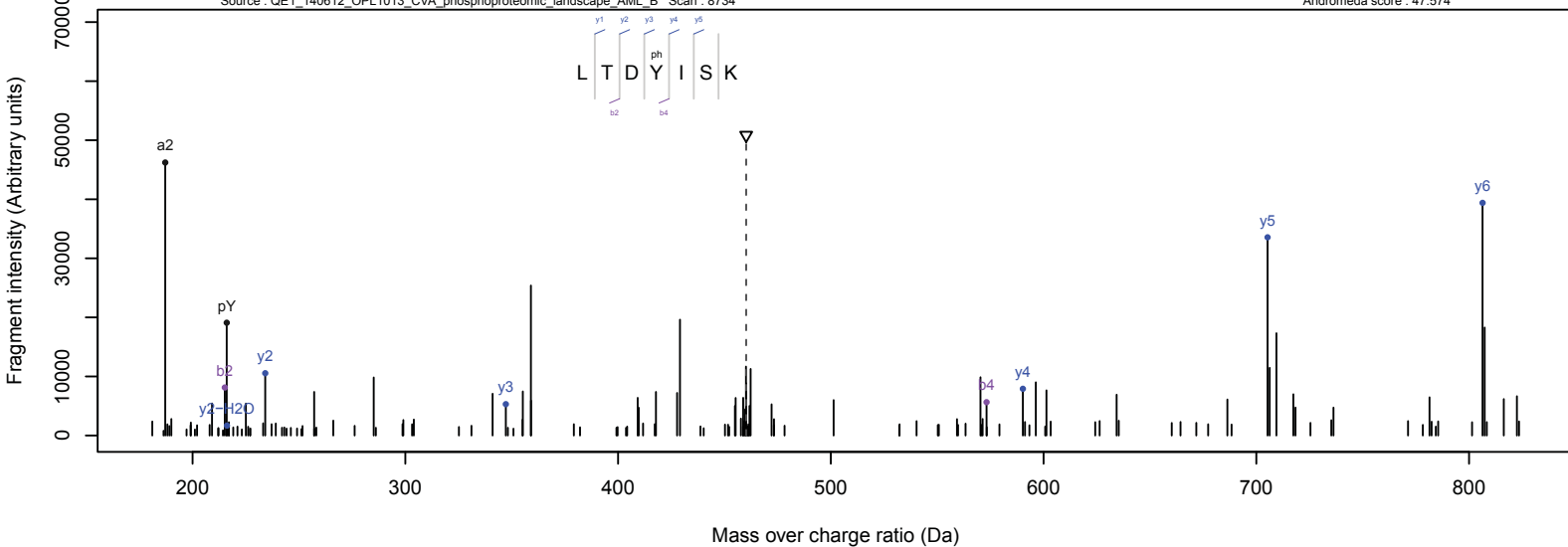
Andromeda score : 47.574



LTDY(ph)ISK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 8734

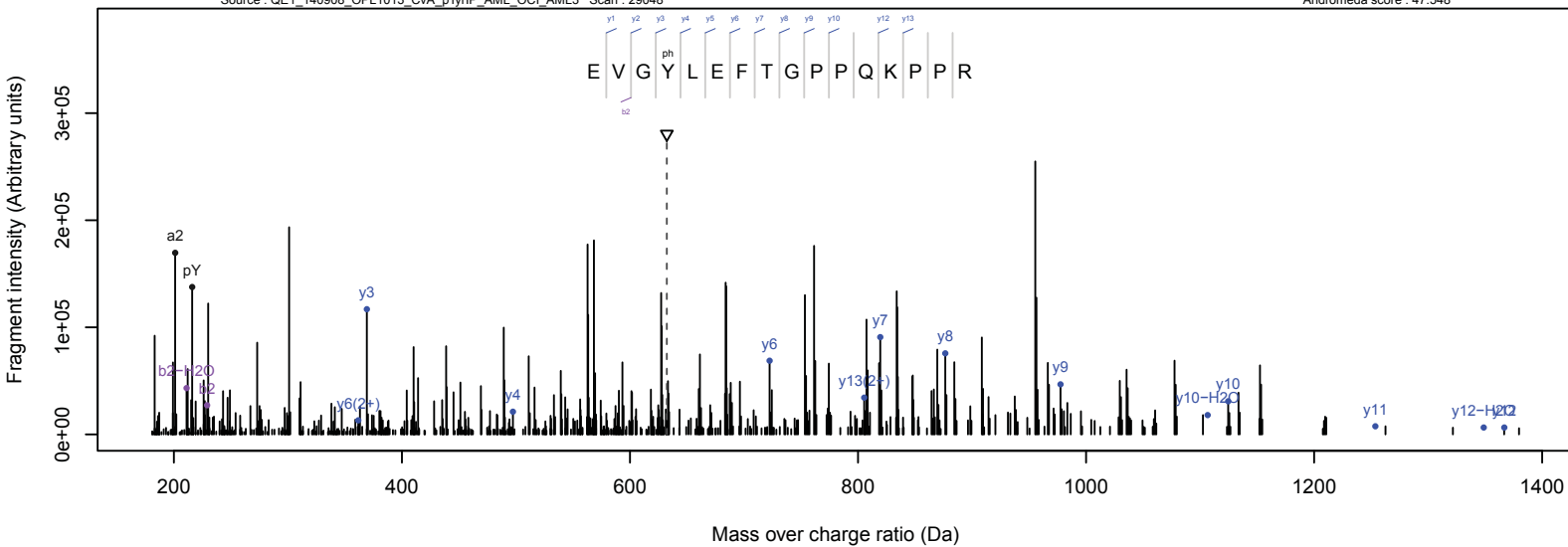
Andromeda score : 47.574



EVGY(ph)LEFTGPPQKPPR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 29048

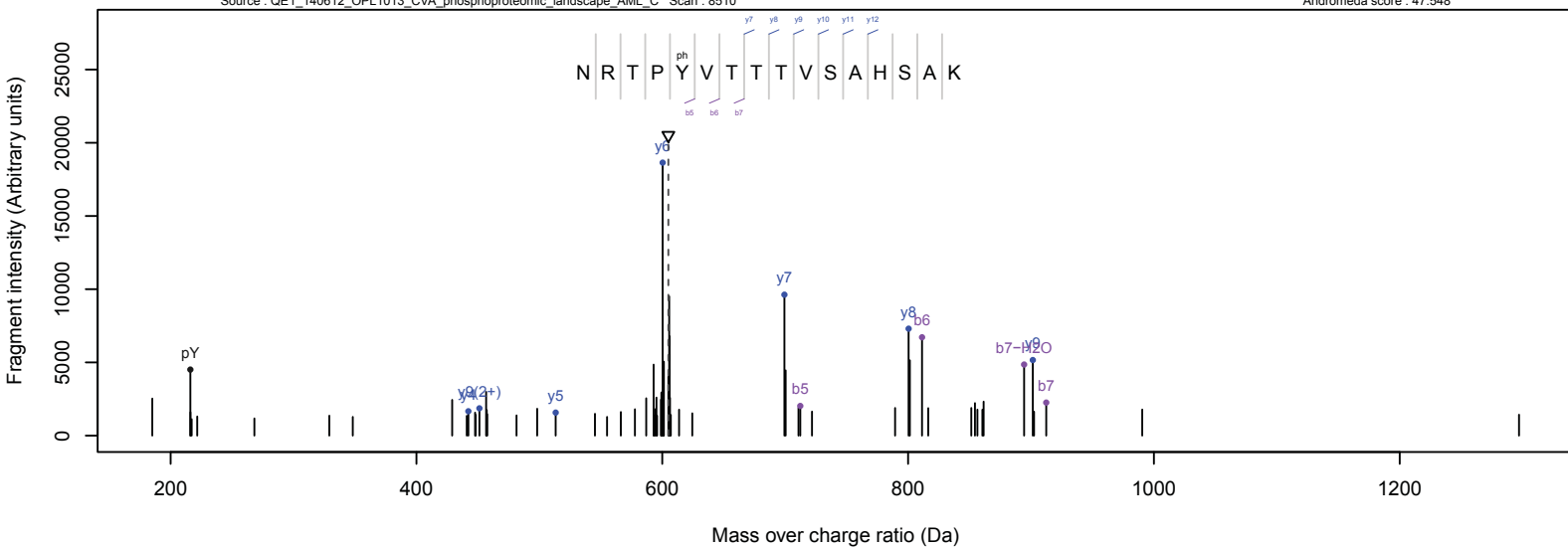
Andromeda score : 47.548



**NRTPY(ph)VTTTSAHSAK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 8510

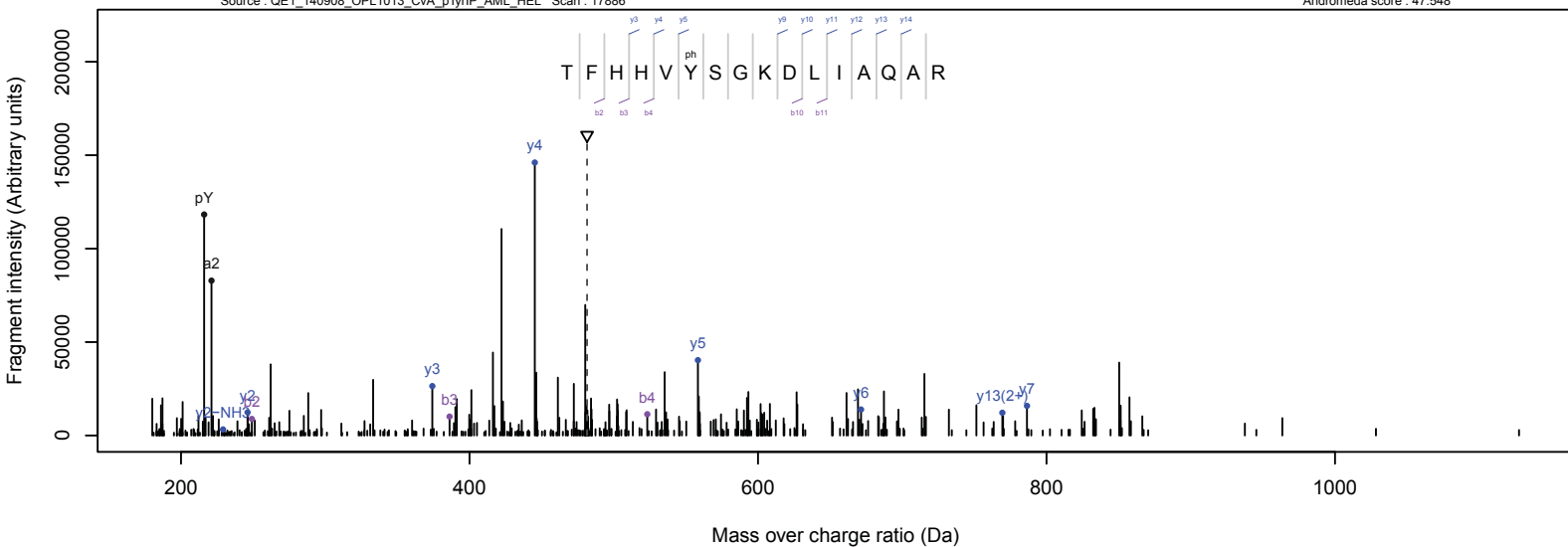
Andromeda score : 47.548



**TFHHVY(ph)SGKDLIAQAR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 17886

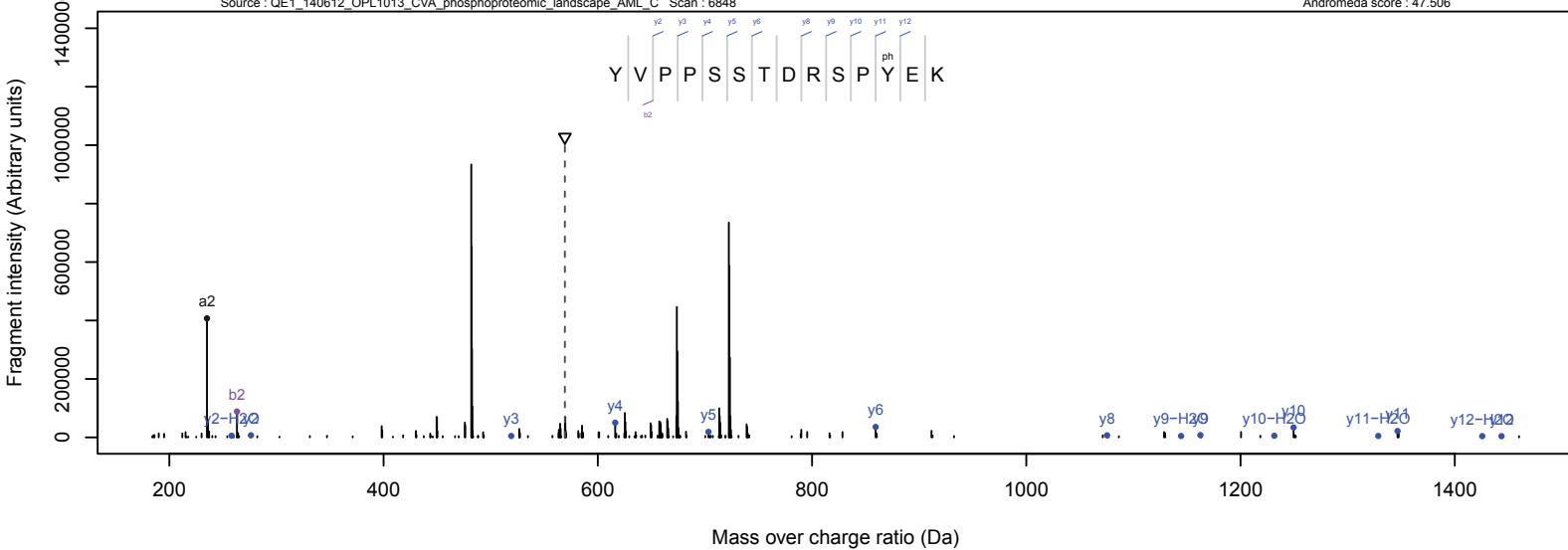
Andromeda score : 47.548



**YVPPSSTRSPY(ph)EK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 6848

Andromeda score : 47.506

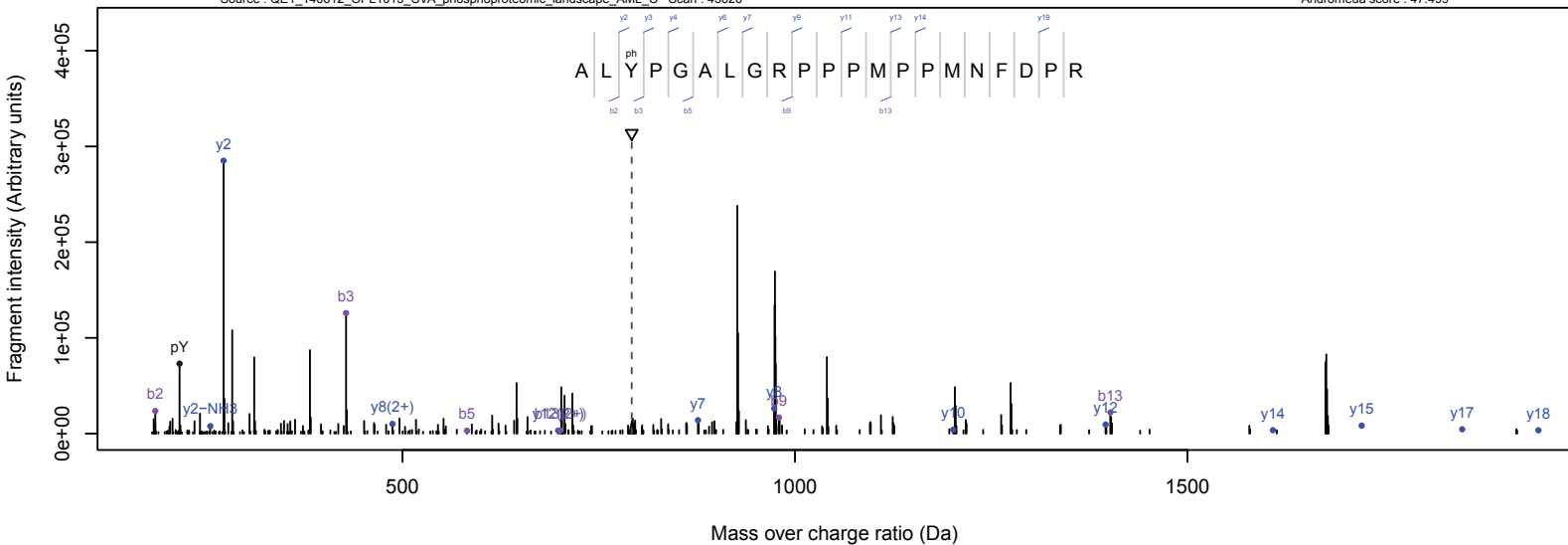




ALY(ph)PGALGRPPMPMNFDP-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43626

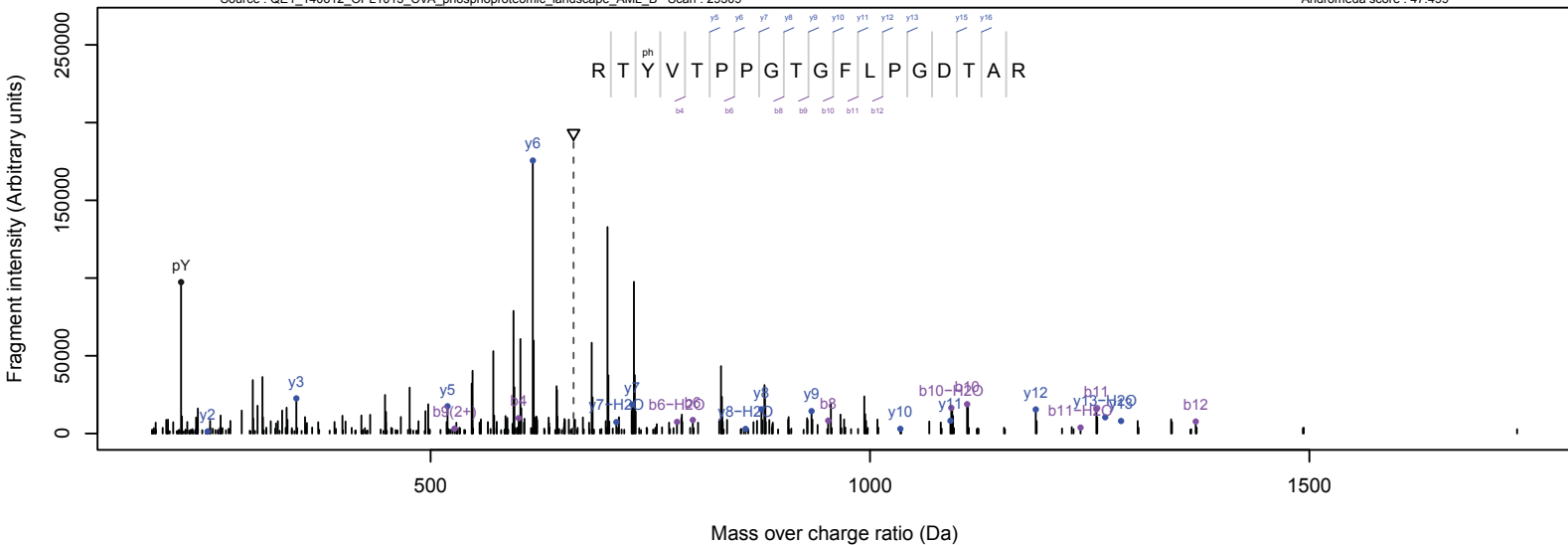
Andromeda score : 47.499



RTY(ph)VTPPGTGFLPGDTAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 29309

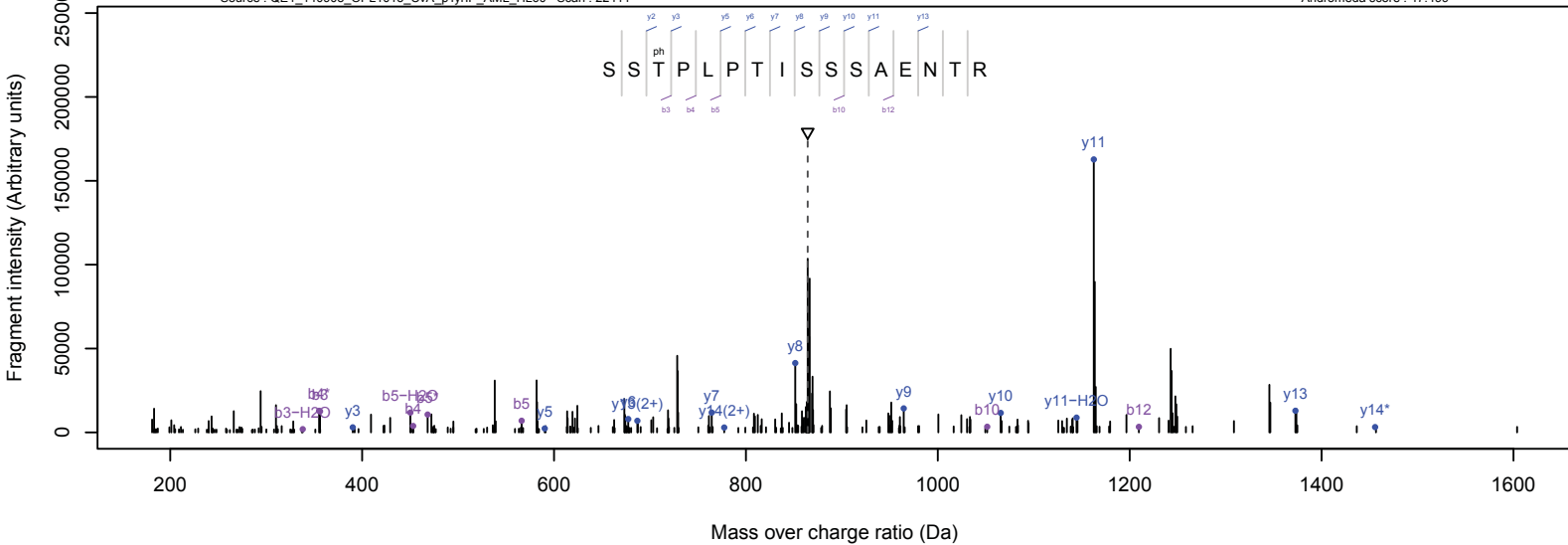
Andromeda score : 47.499



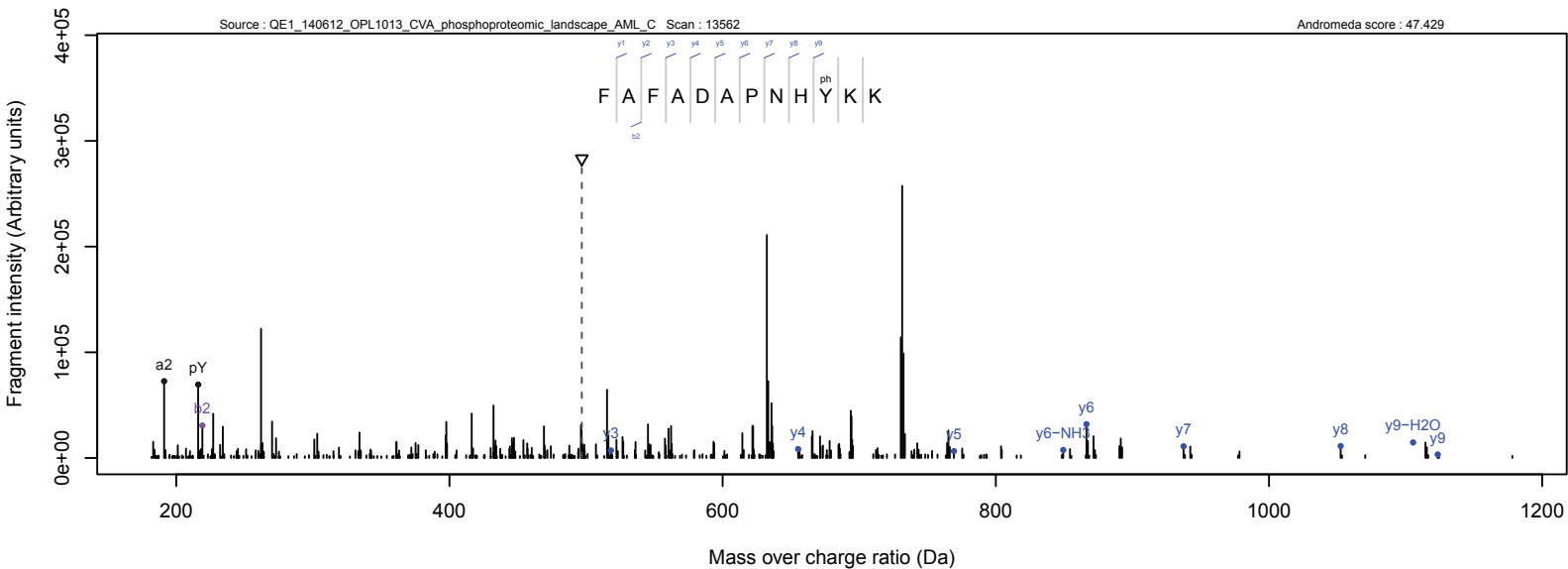
SST(ph)PLPTISSAENTR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 22411

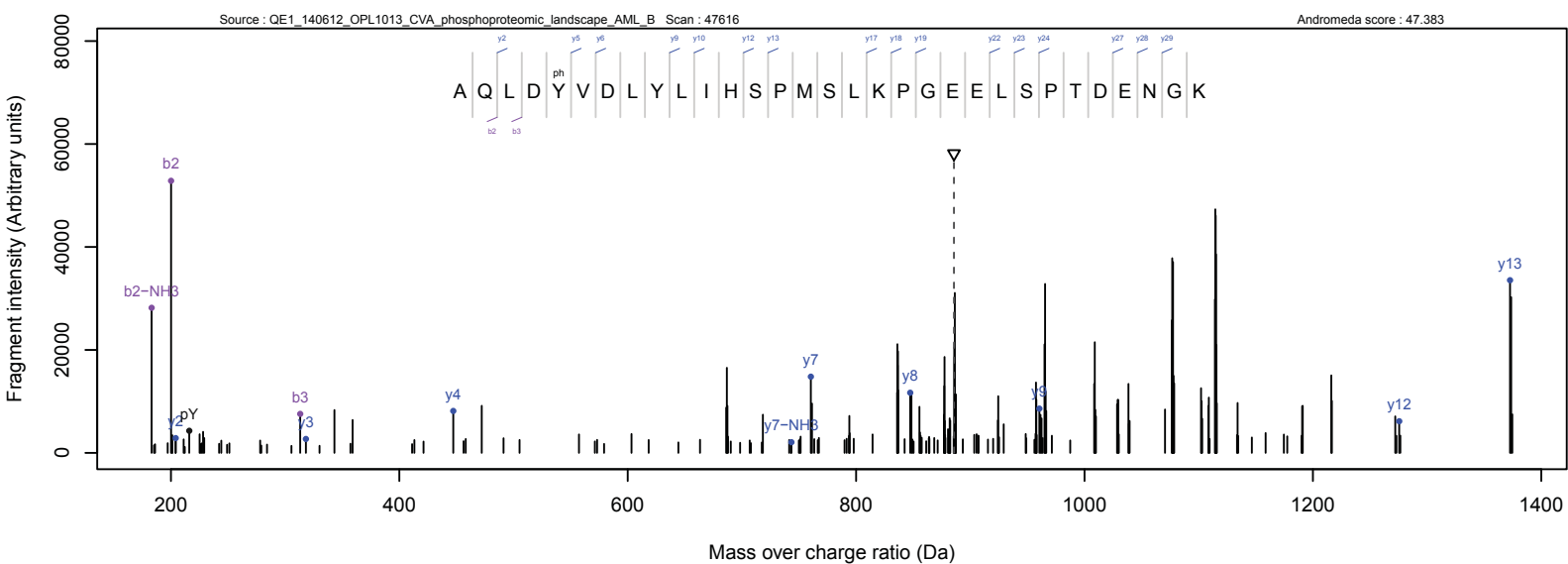
Andromeda score : 47.499



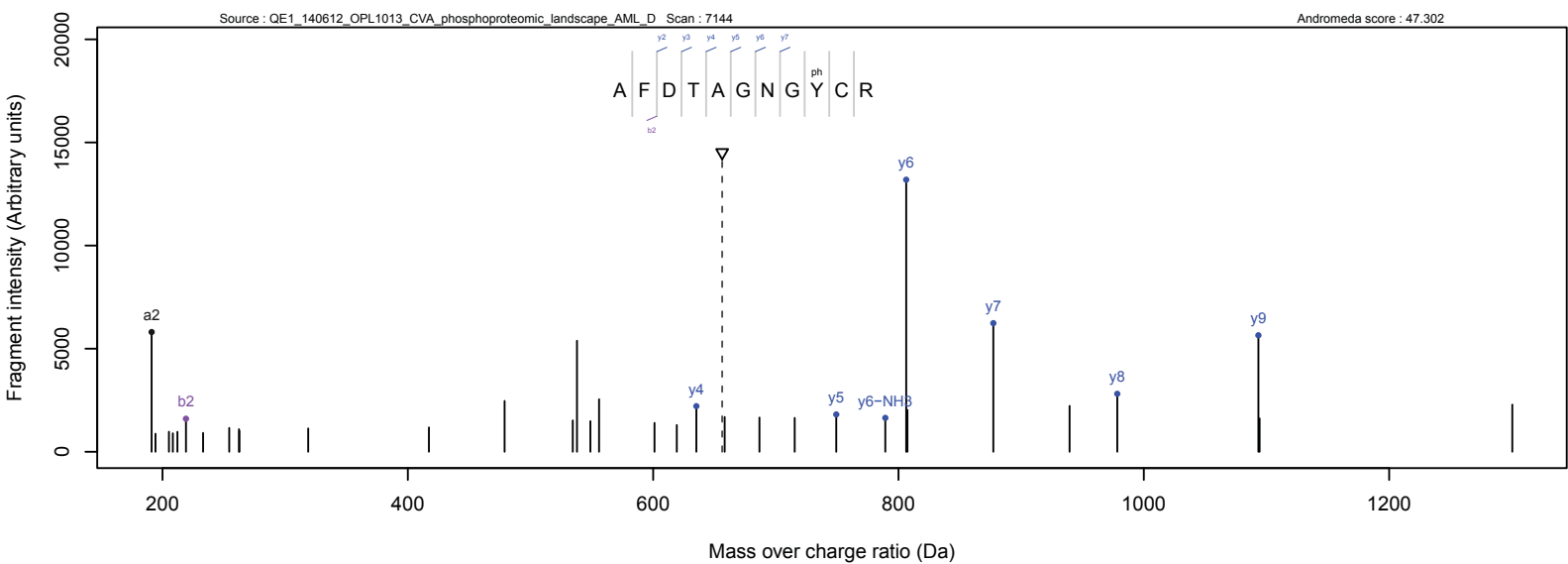
**FAFADAPNHY(ph)KK-3+**



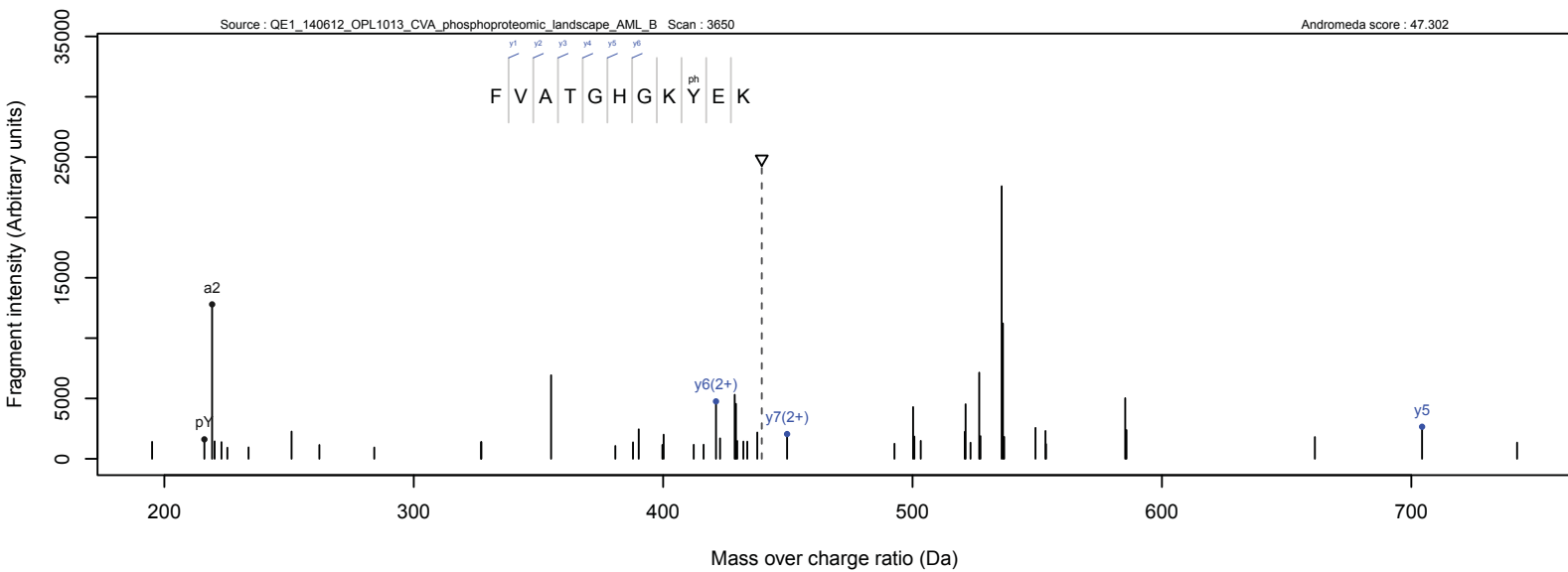
**AQLDY(ph)VDLYLHSPMSLKPGEEELSPTDENGK-4+**



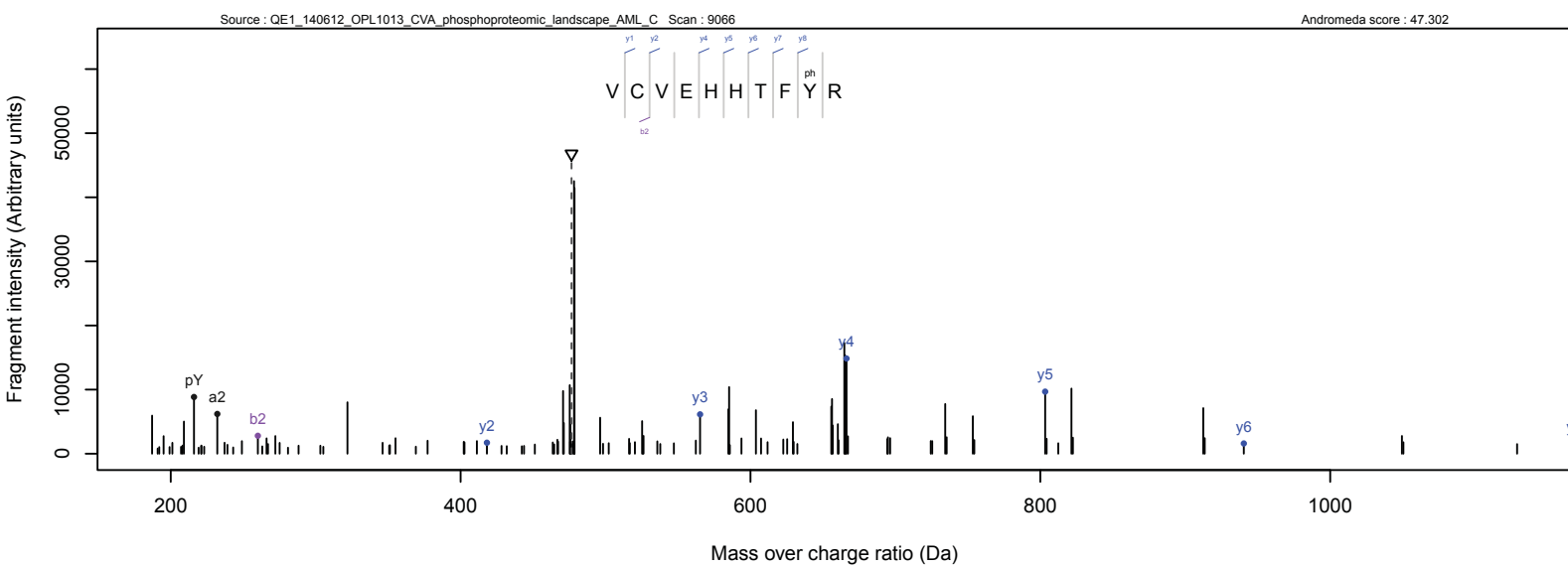
**AFDTAGNGY(ph)CR-2+**



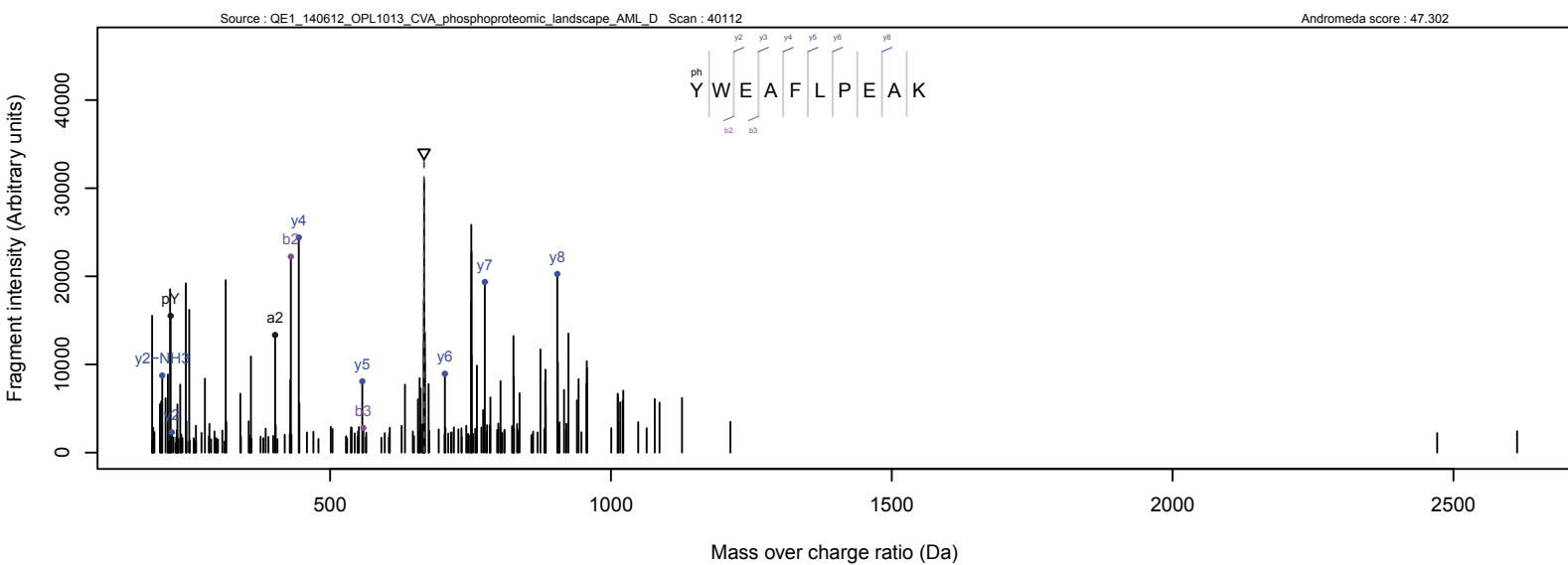
## FVATGHGKY(ph)EK-3+



## VCVEHHTFY(ph)R-3+



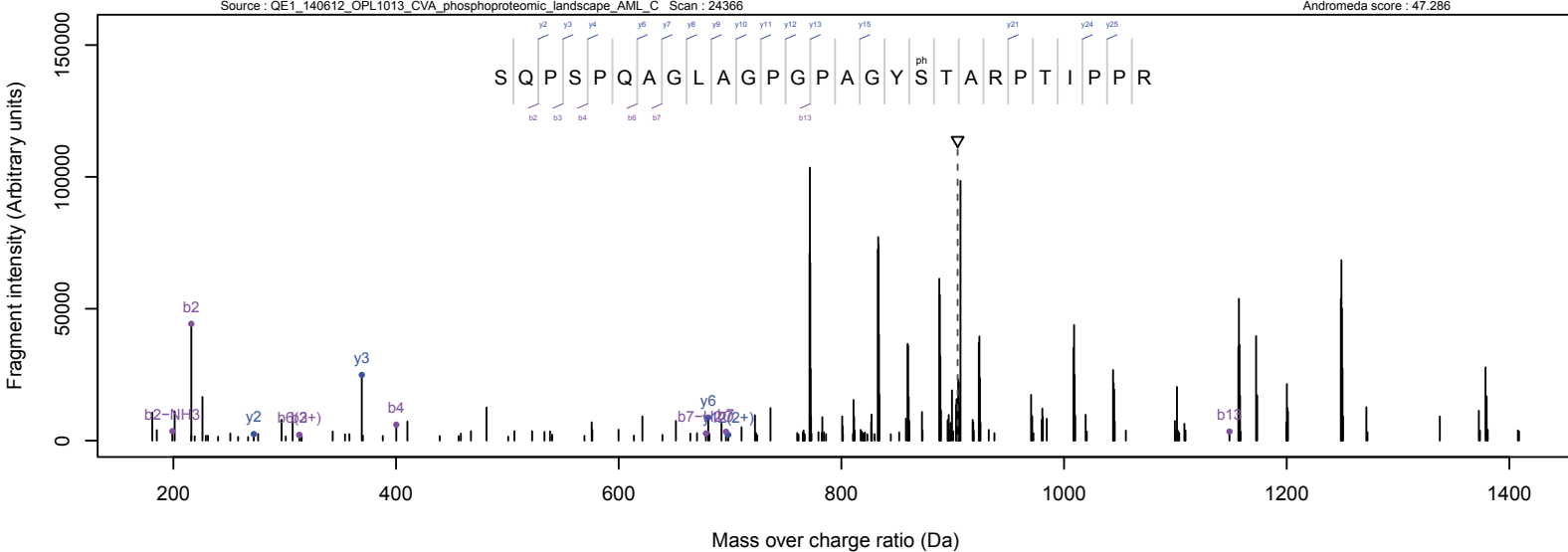
## Y(ph)WEAFLPEAK-2+



## SQPSPQAGLAGPGAGYS(ph)TARPTIPPR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 24366

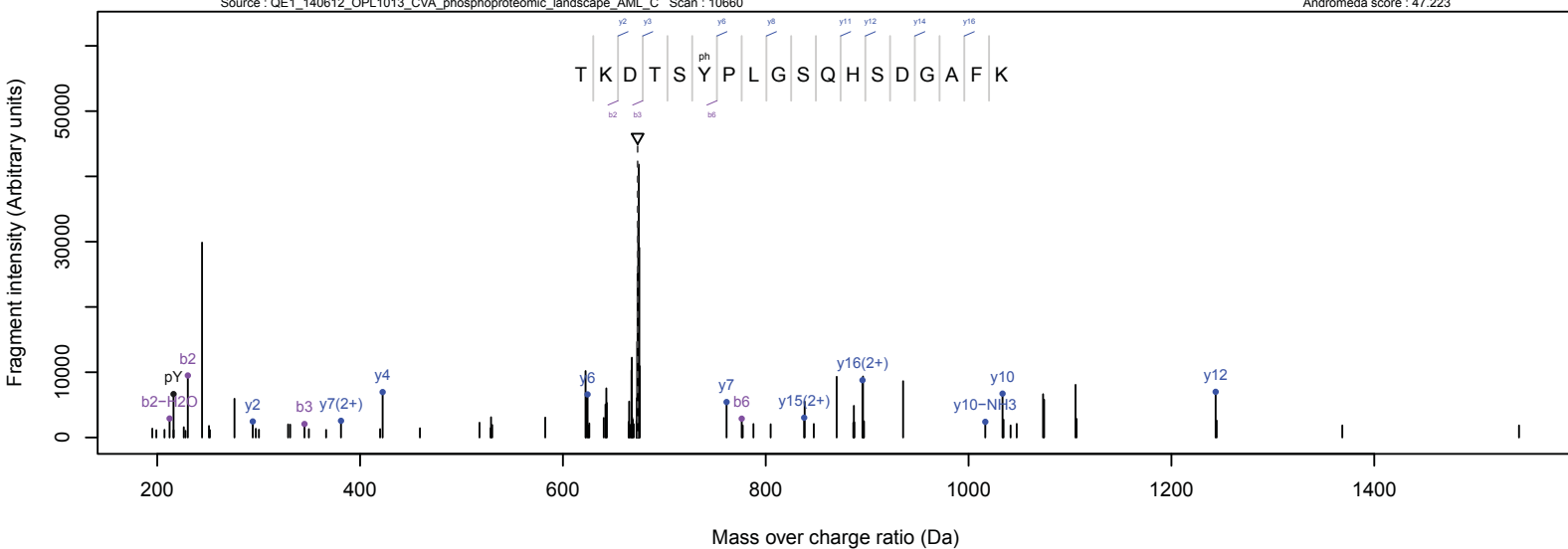
Andromeda score: 47.286



## TKDTSY(ph)PLGSQHS DGA FK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 10660

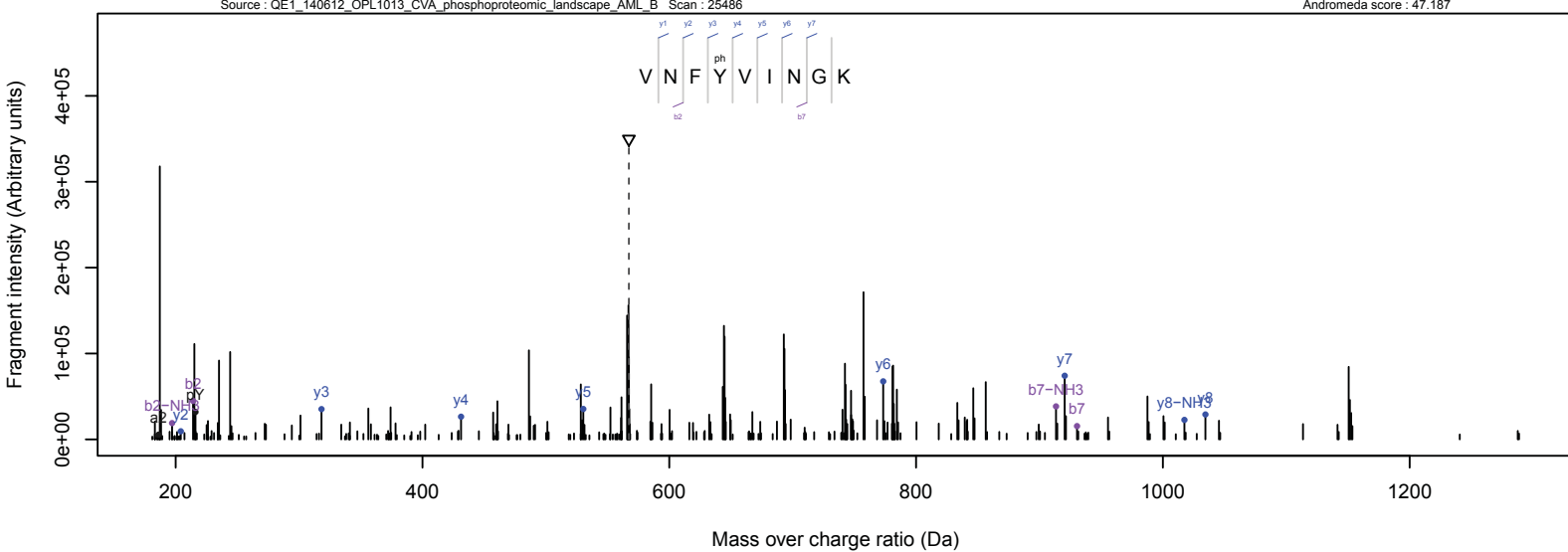
Andromeda score: 47.223



## VNFY(ph)VINGK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 25486

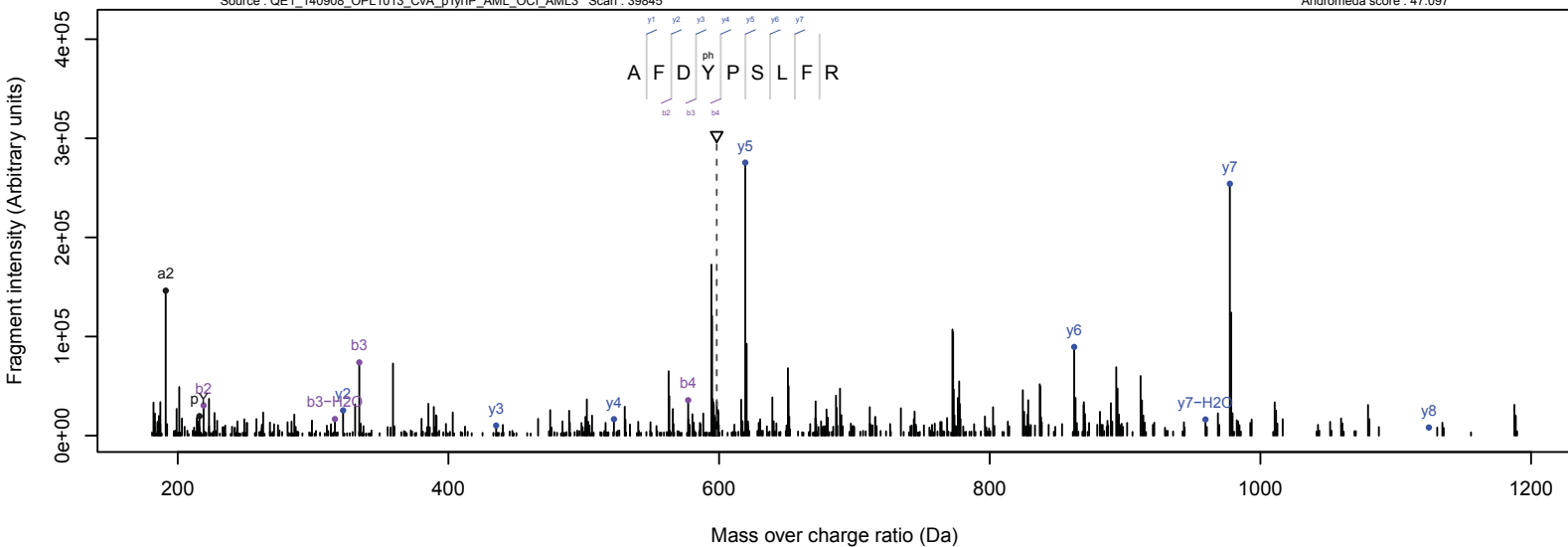
Andromeda score: 47.187



## AFDY(ph)PSLFR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 39845

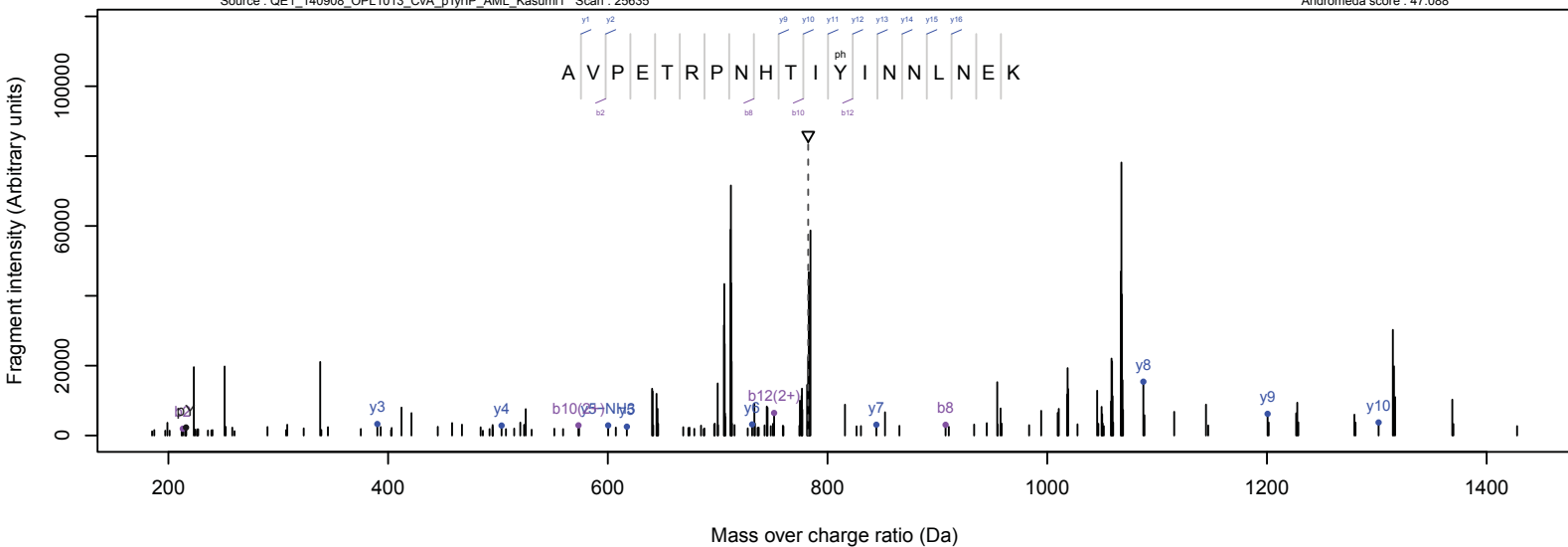
Andromeda score : 47.097



## (ac)AVPETRPNHTIY(ph)INNLNEK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 25635

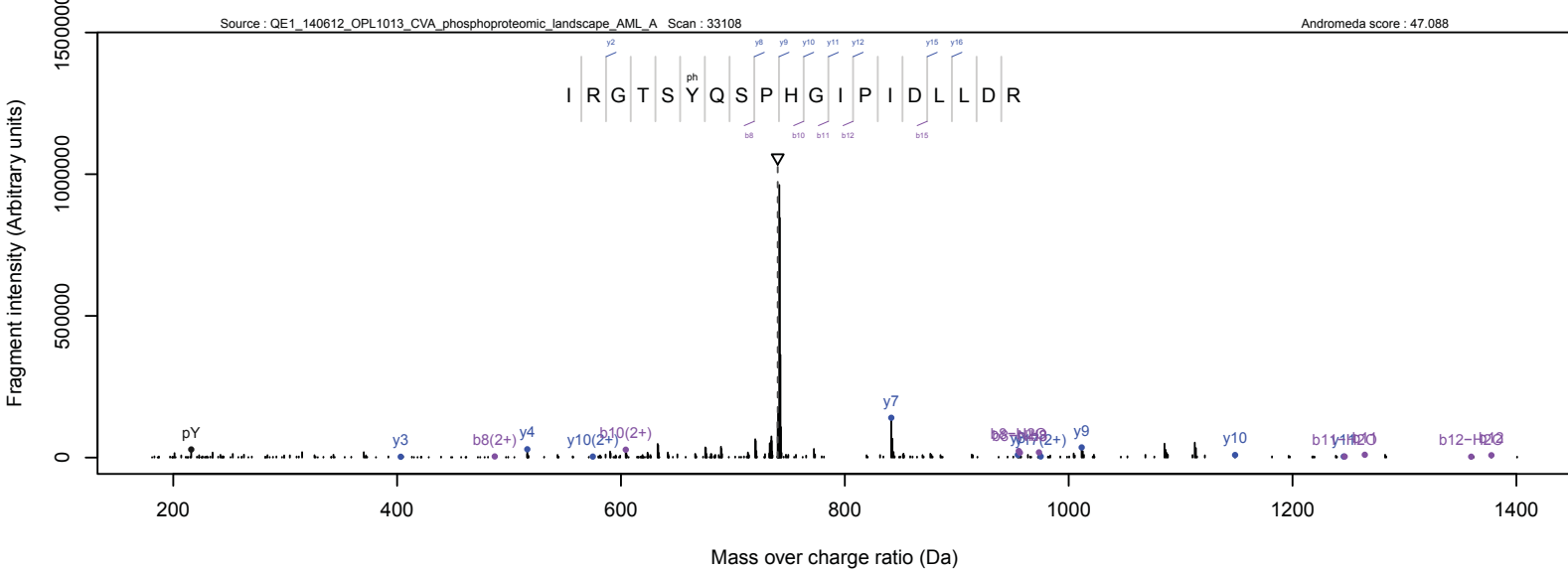
Andromeda score : 47.088



## IRGTSY(ph)QSPHGIPIDLLDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33108

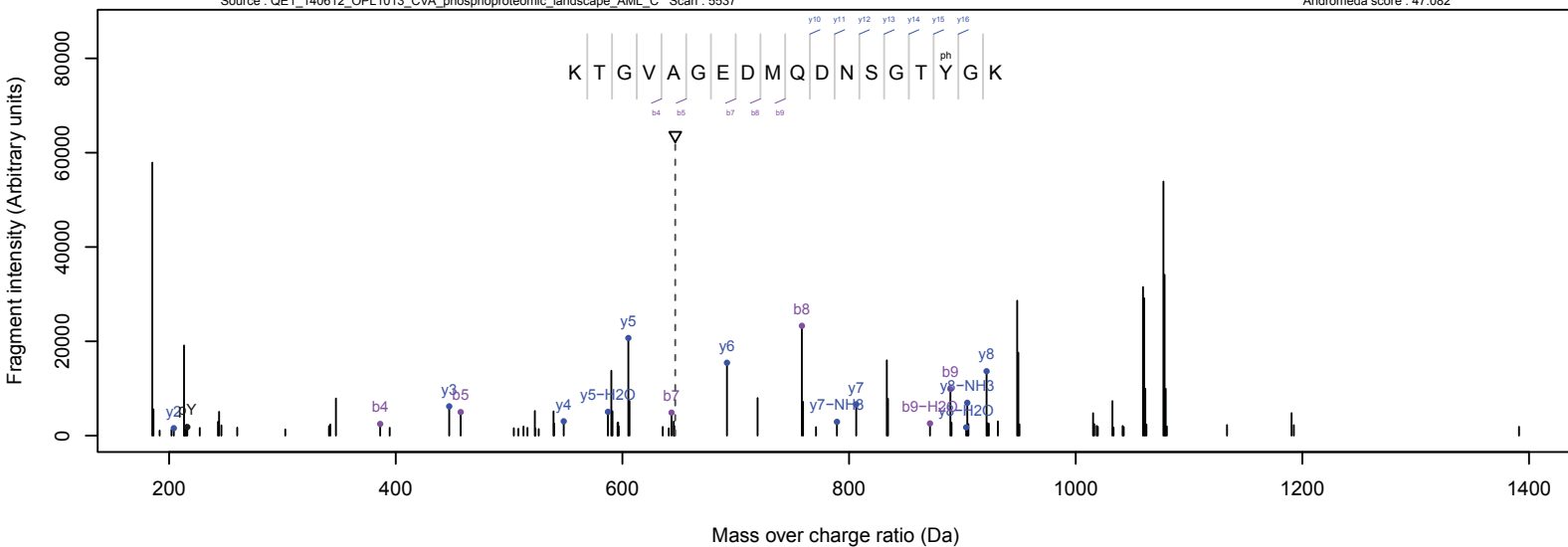
Andromeda score : 47.088



## KTGVAGEDMQDMSGTY(ph)GK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 5537

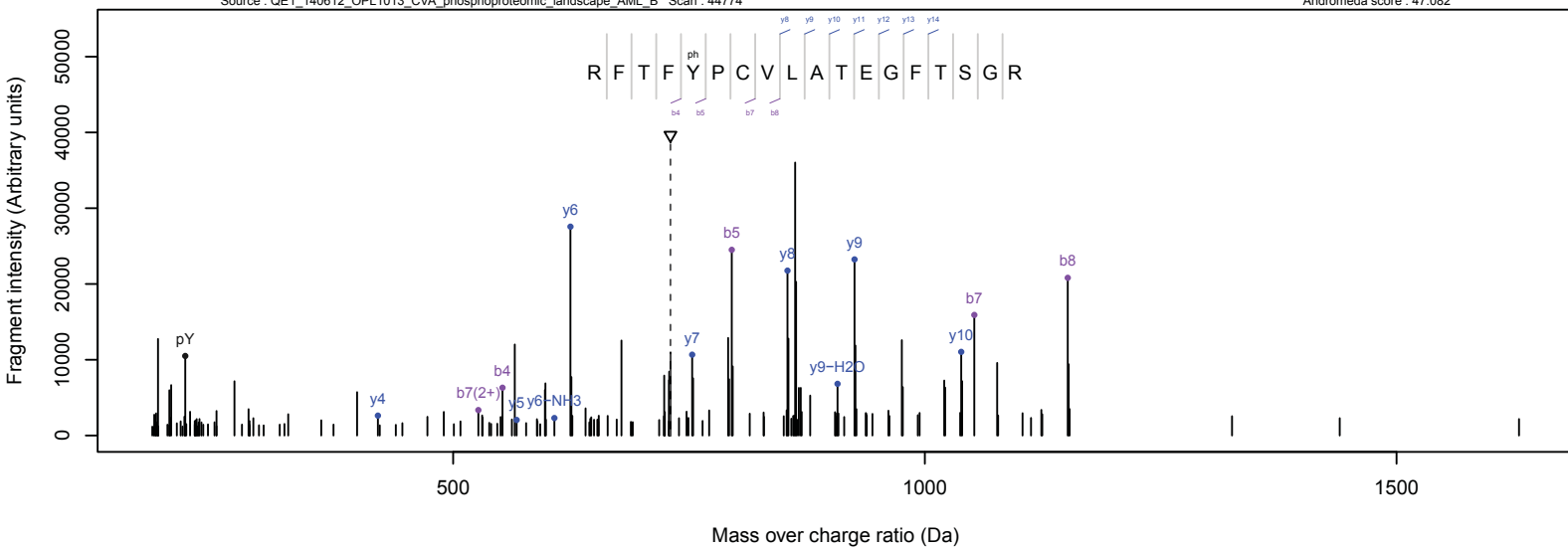
Andromeda score : 47.082



## RFTFY(ph)PCVLATEGFTSGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44774

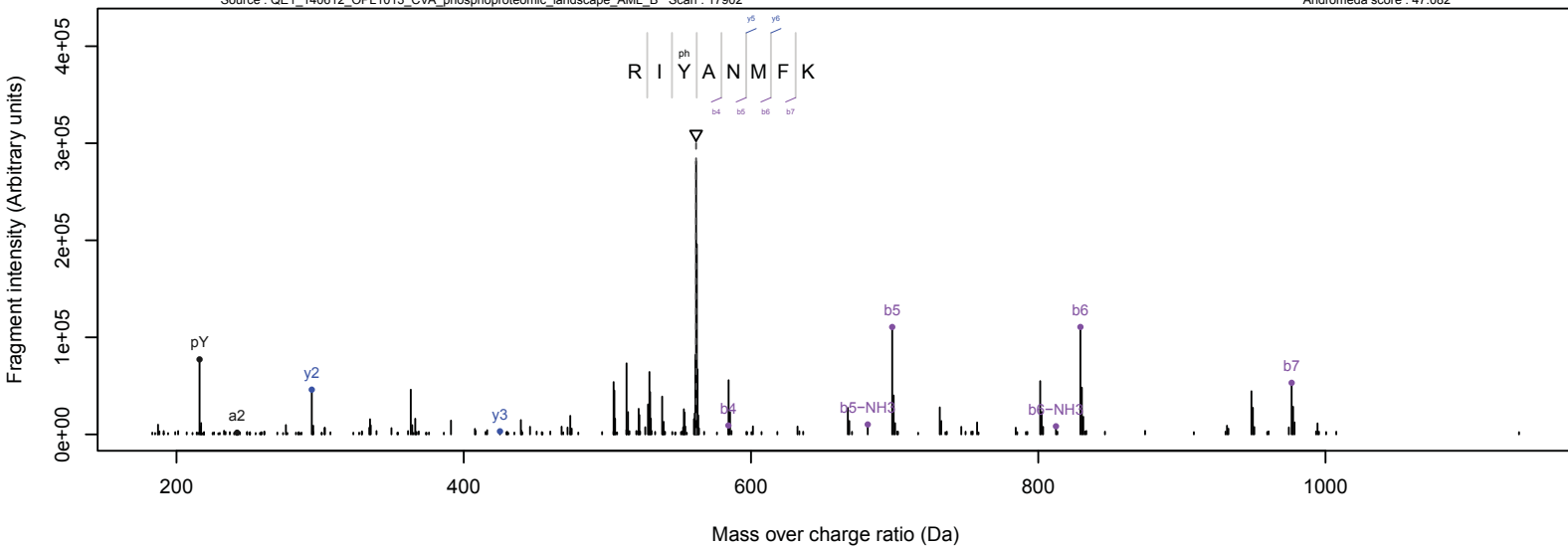
Andromeda score : 47.082



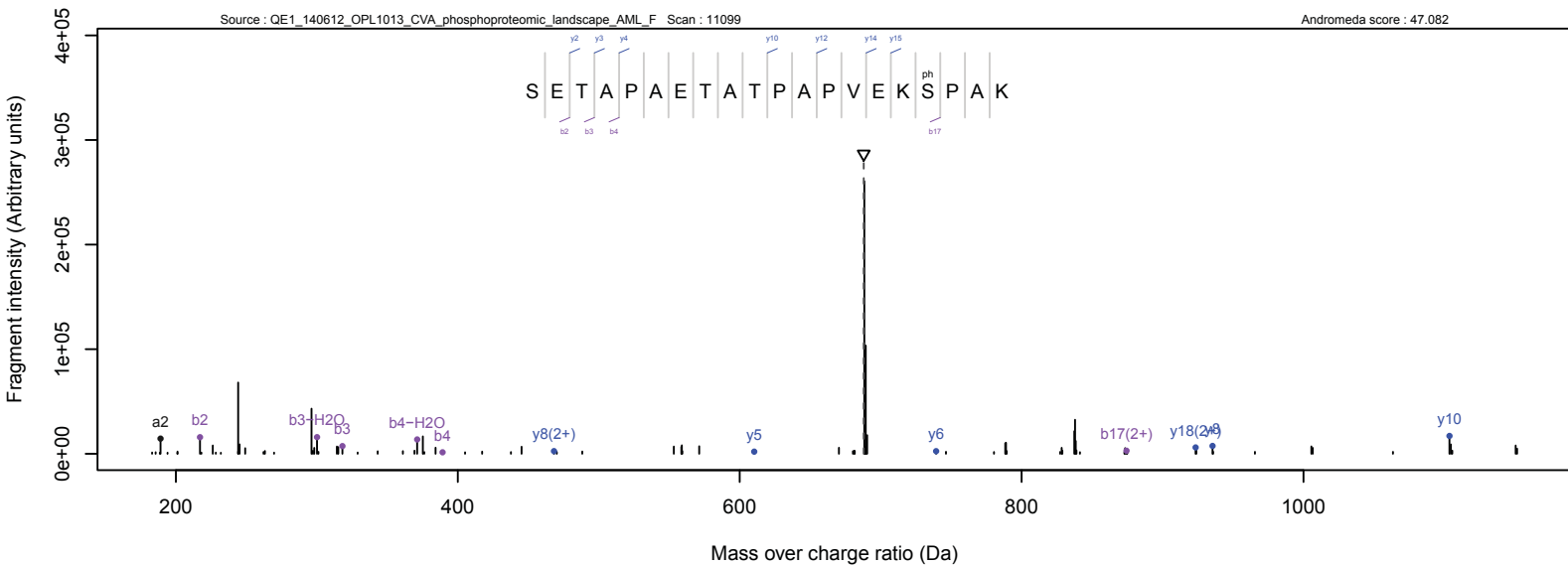
## RIY(ph)ANMFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17902

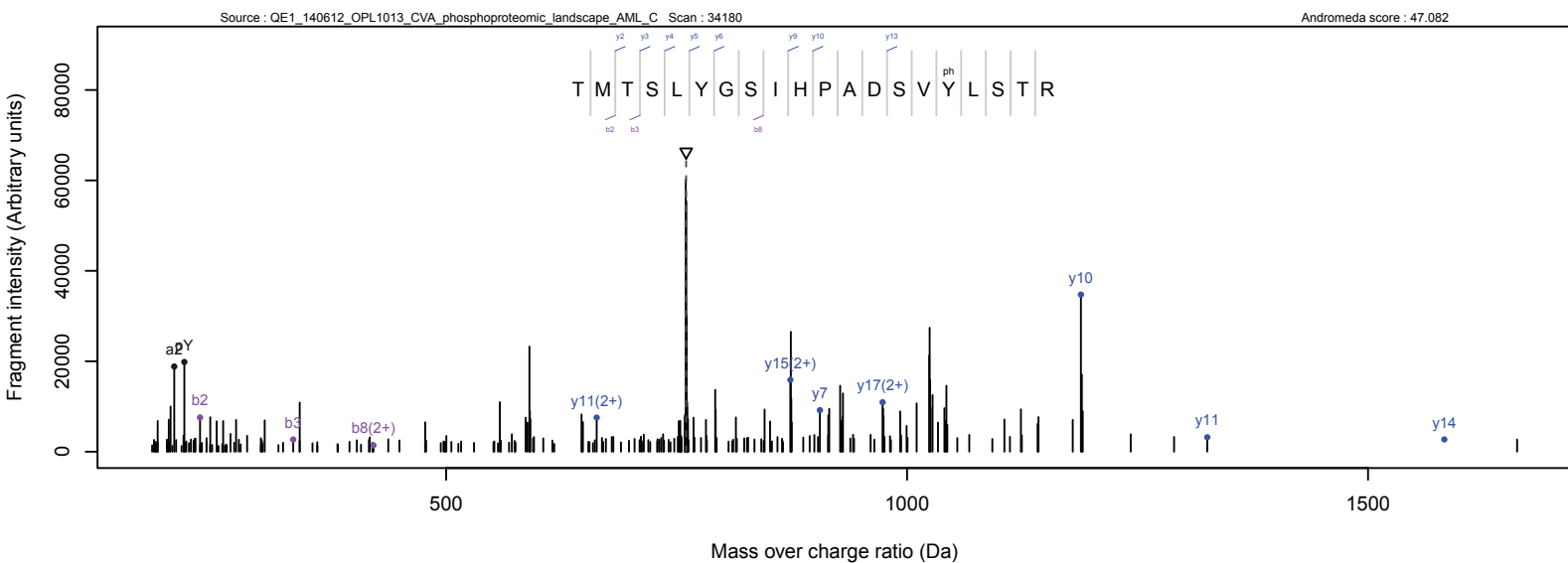
Andromeda score : 47.082



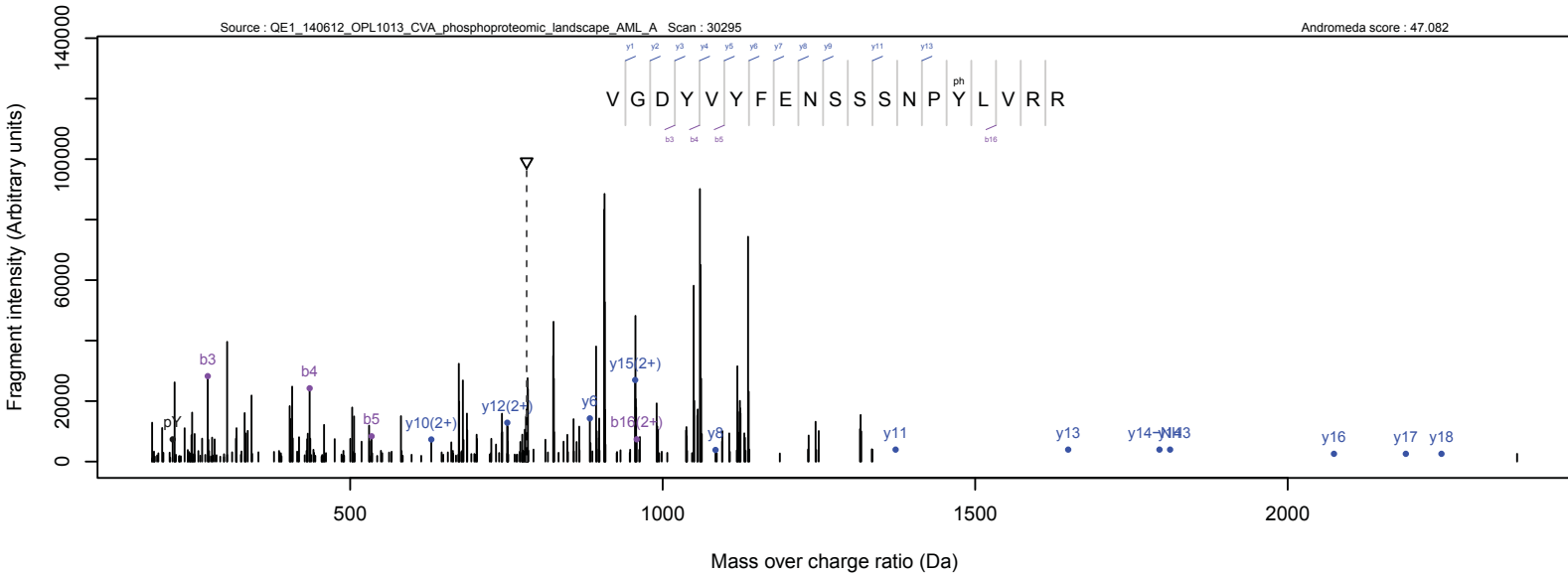
SETAPAETATPAPVEKS(ph)PAK-3+



TMTSLYGSIH PADS VY(ph) LSTR-3+



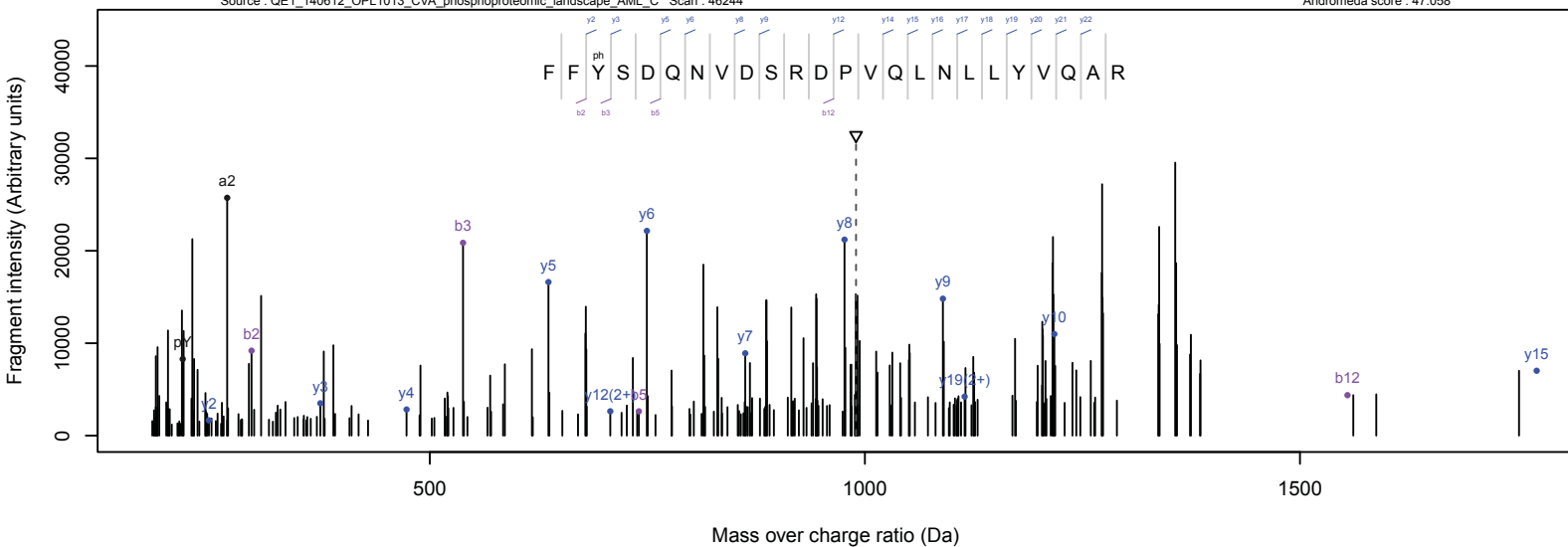
VDYVYFENSSNPY(ph)LVRR-3+



## FFY(ph)SDQNVDSRDPVQLNLLYQAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46244

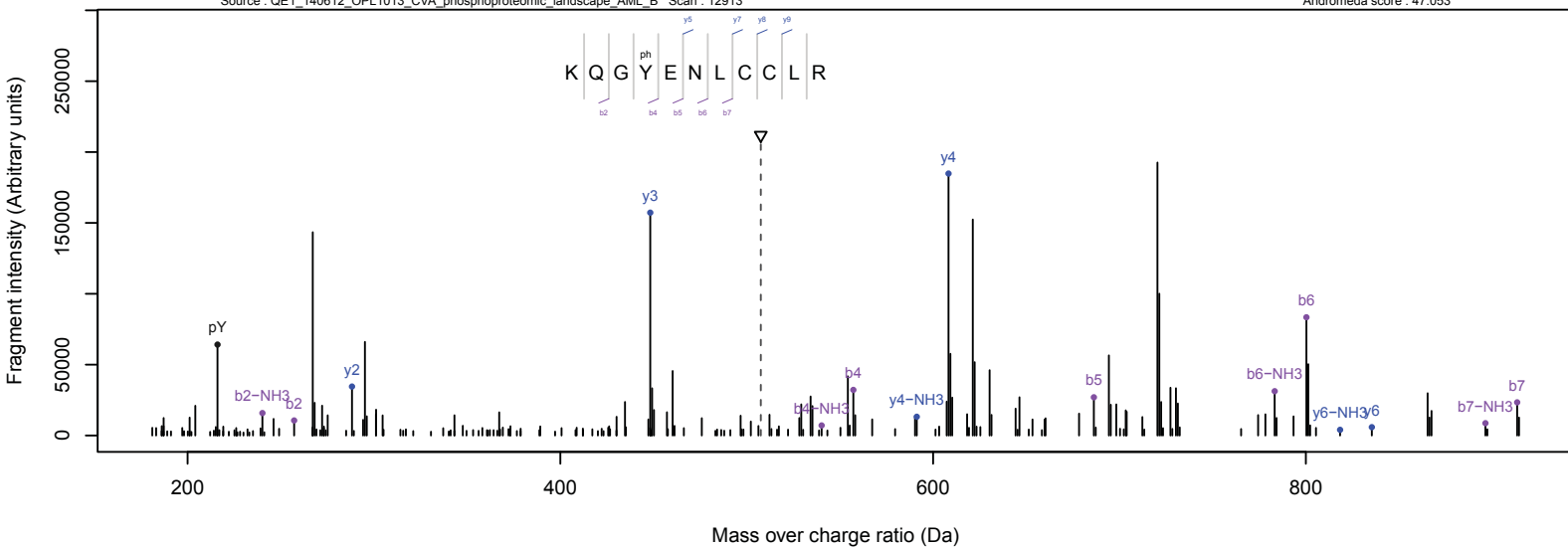
Andromeda score : 47.058



## KQGY(ph)ENLCLLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12913

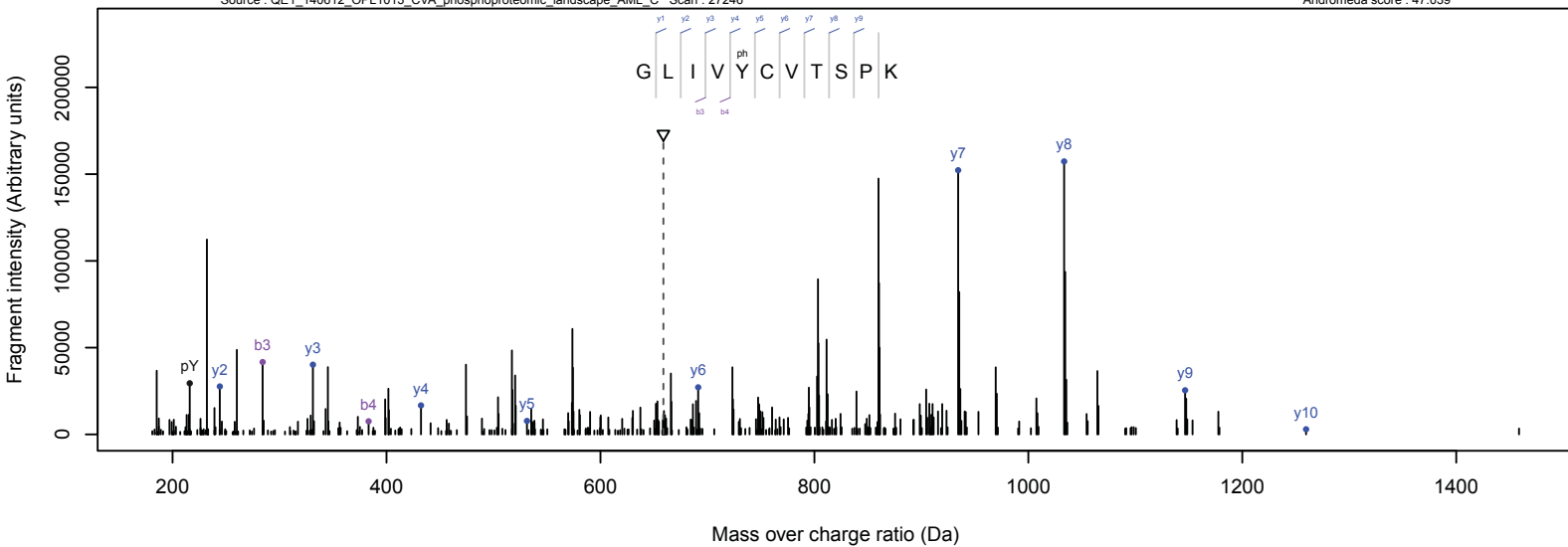
Andromeda score : 47.053



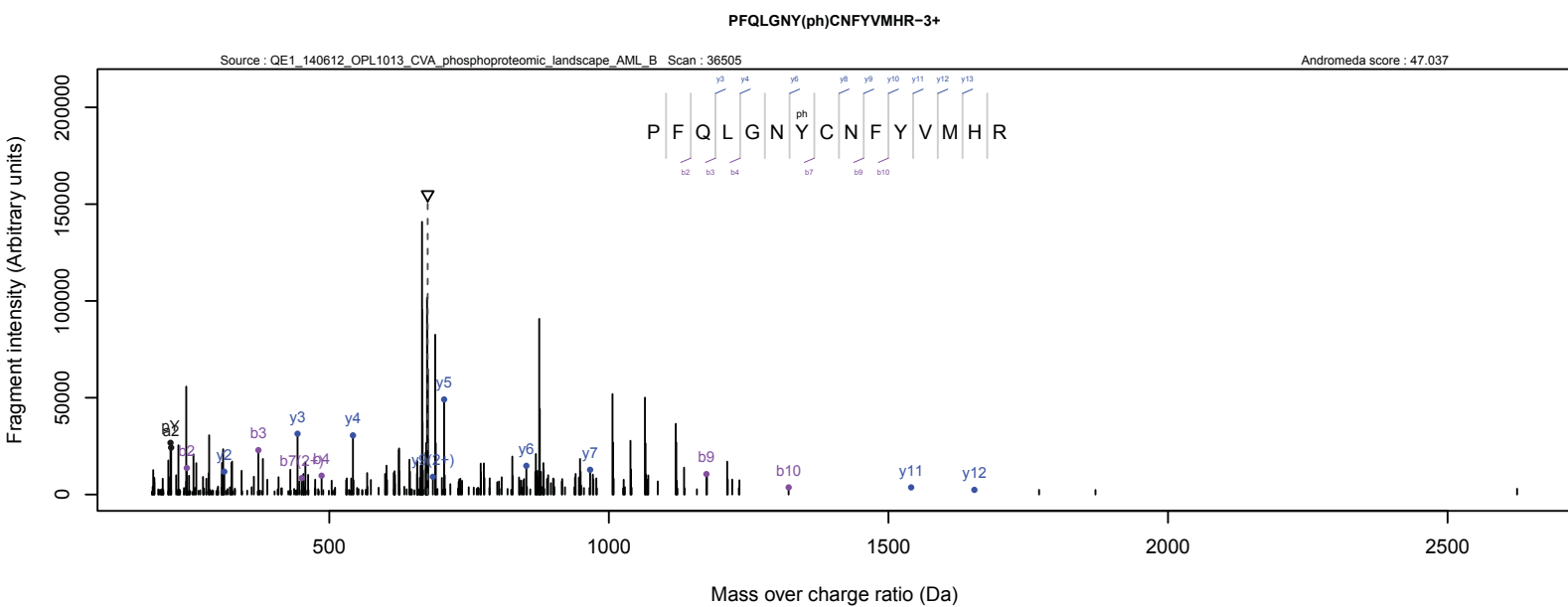
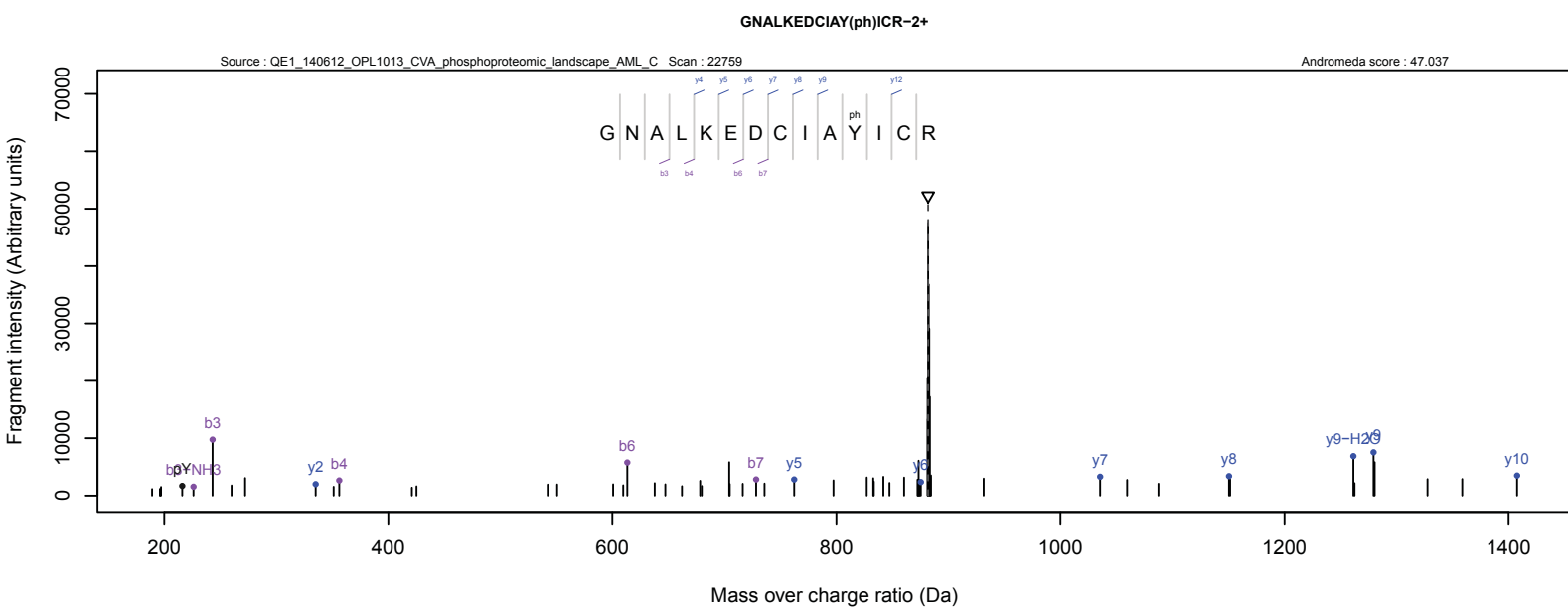
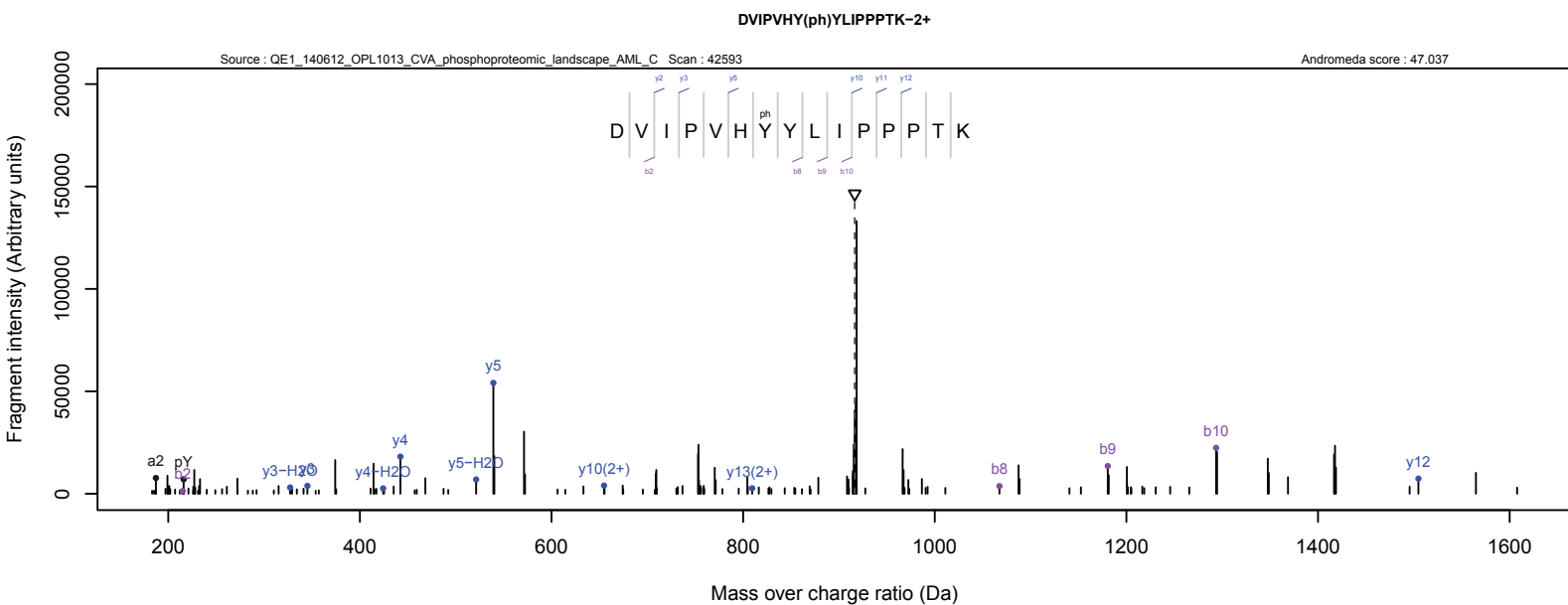
## GLIVY(ph)CVTSPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27246

Andromeda score : 47.039



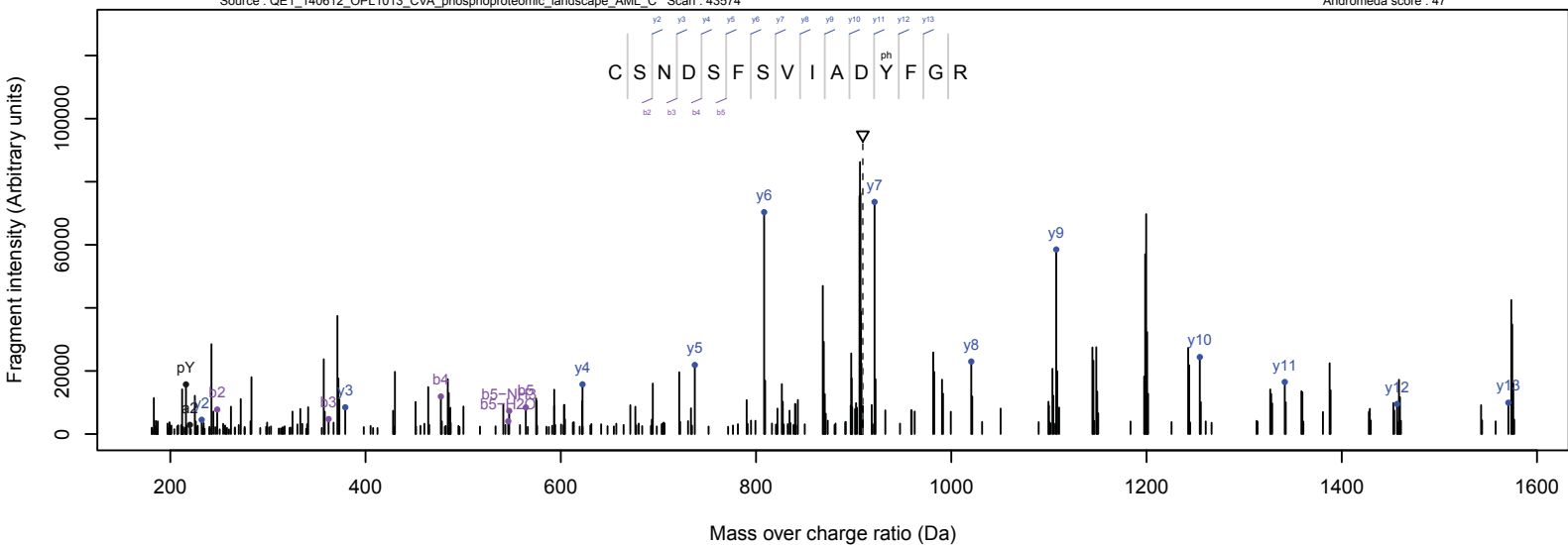




## CSNDSFSVIADY(ph)FGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43574

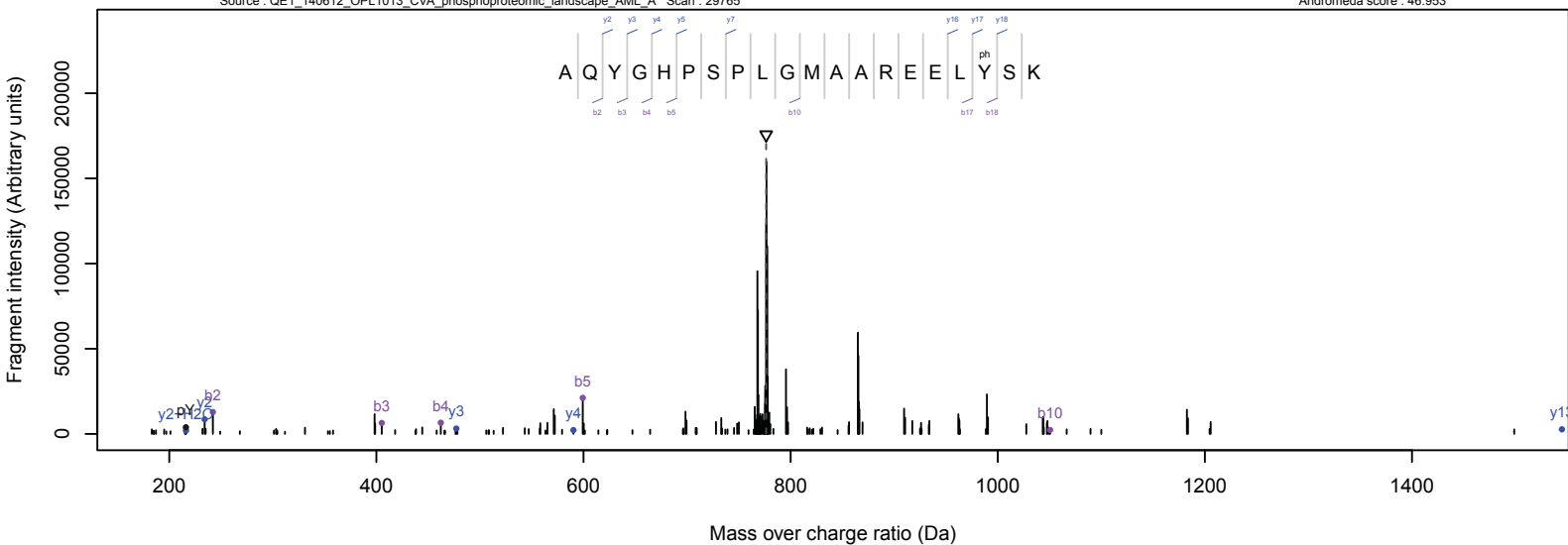
Andromeda score : 47



## (ac)AQYGHPSPLGMAAREELY(ph)SK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29765

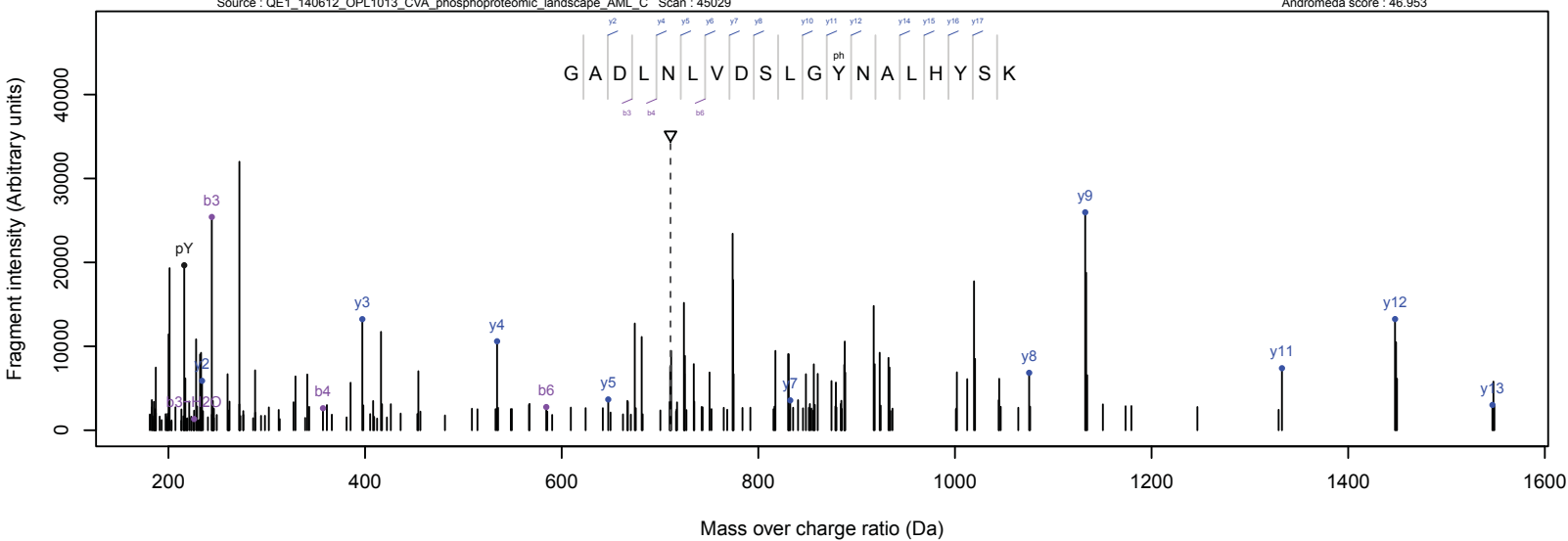
Andromeda score : 46.953

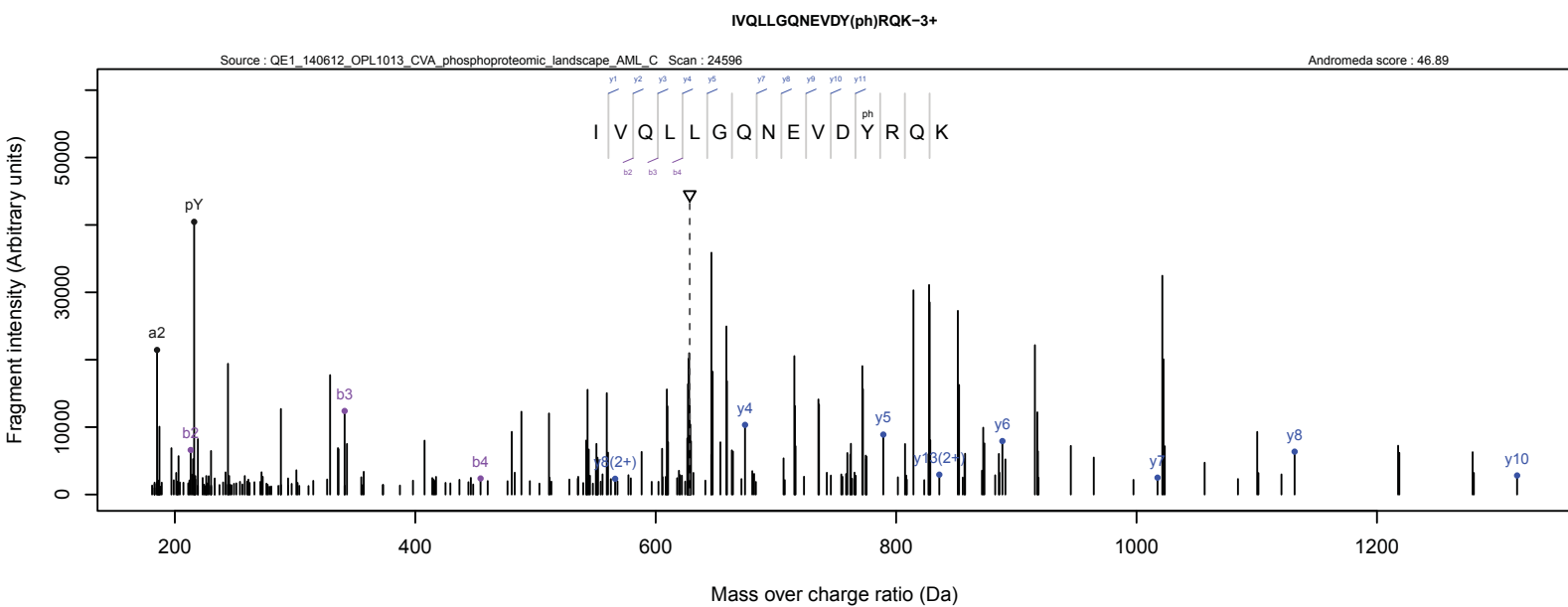
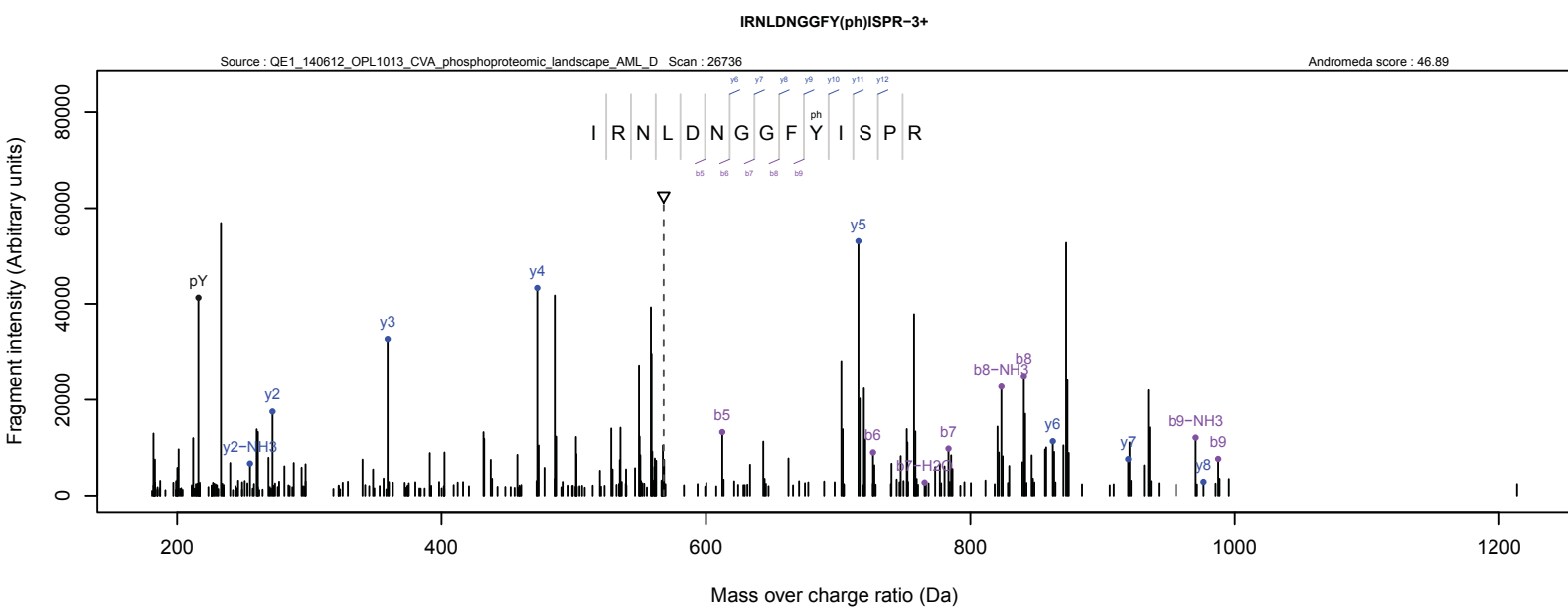
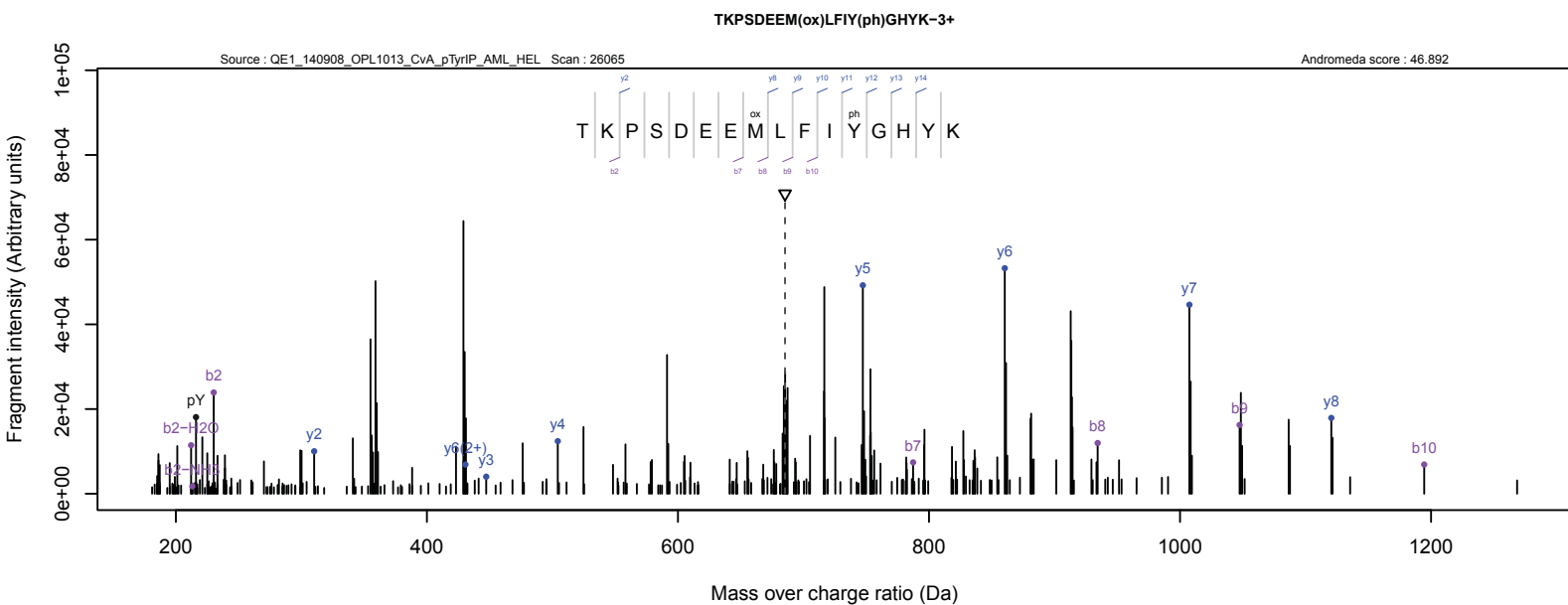


## GADLNLVDSLGY(ph)NALHYSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45029

Andromeda score : 46.953

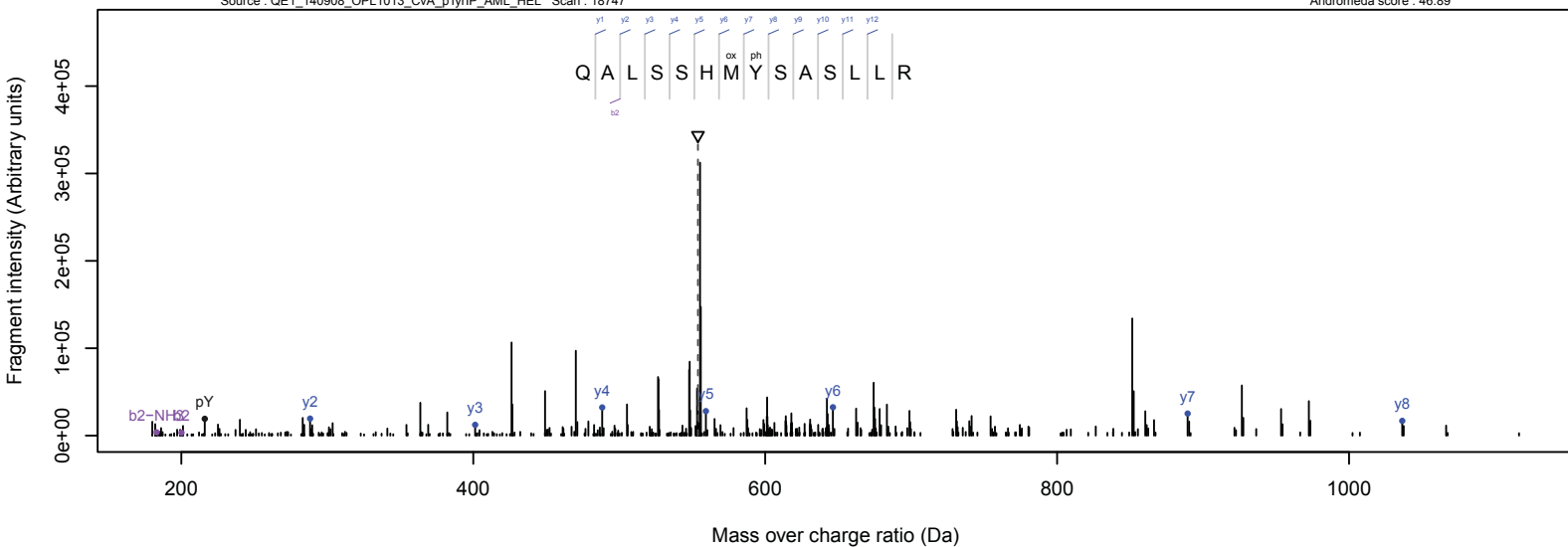




## QALSSHM(ox)Y(ph)SASLLR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 18747

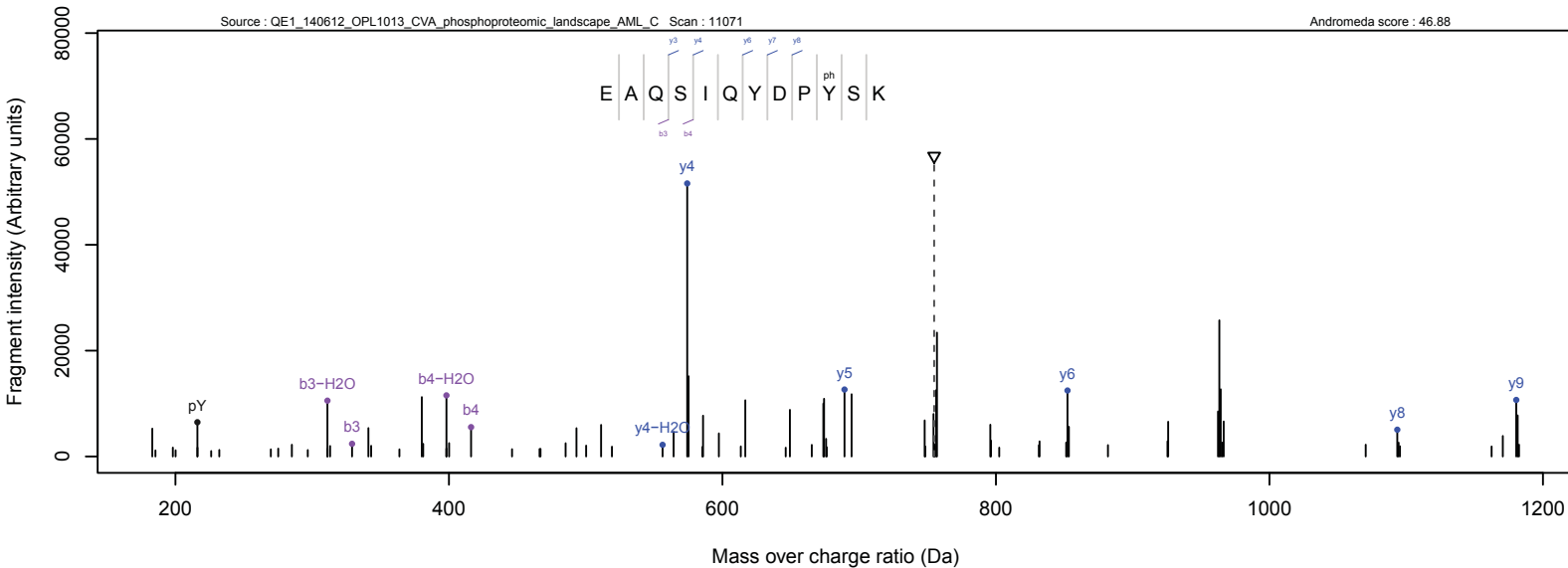
Andromeda score : 46.89



## EAQSIQYDPY(ph)SK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11071

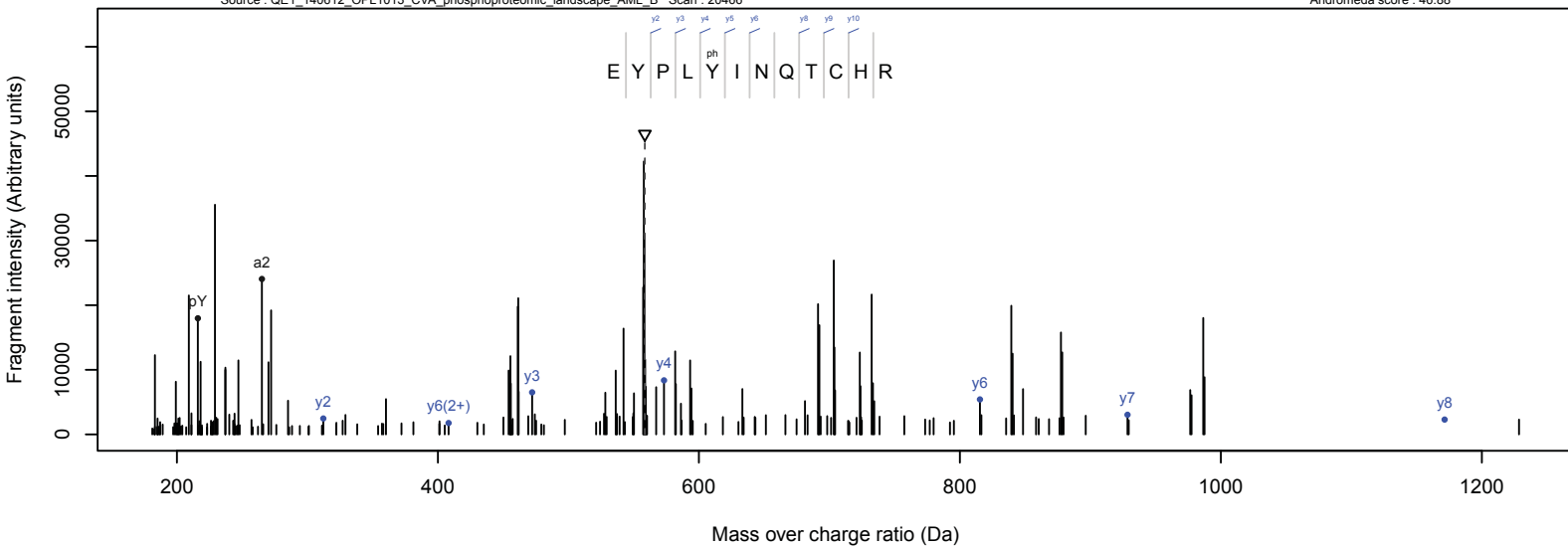
Andromeda score : 46.88

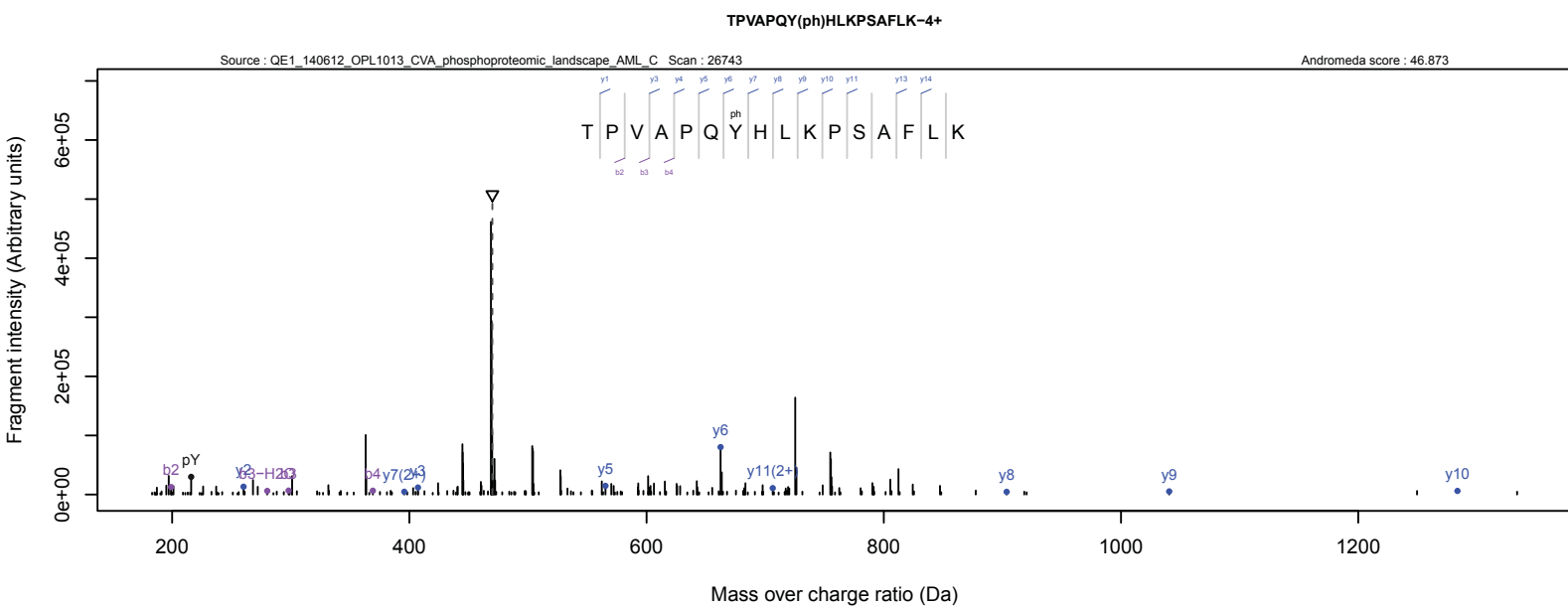
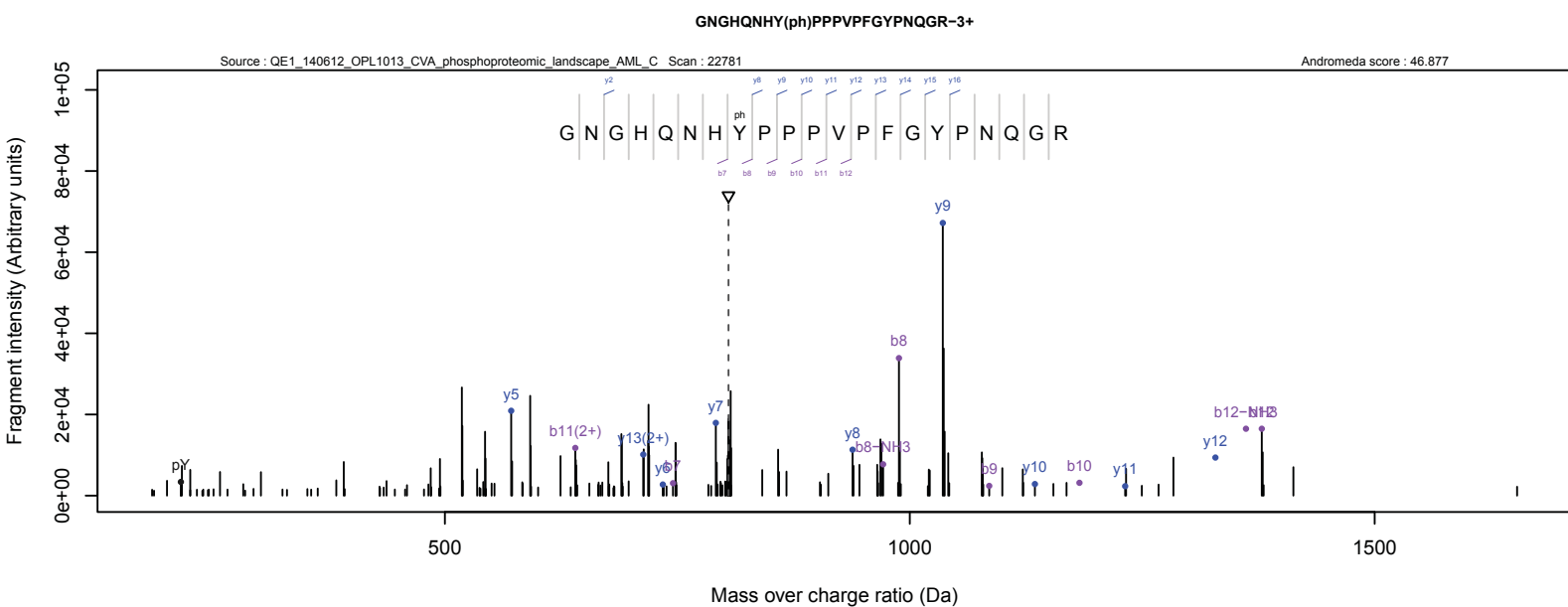
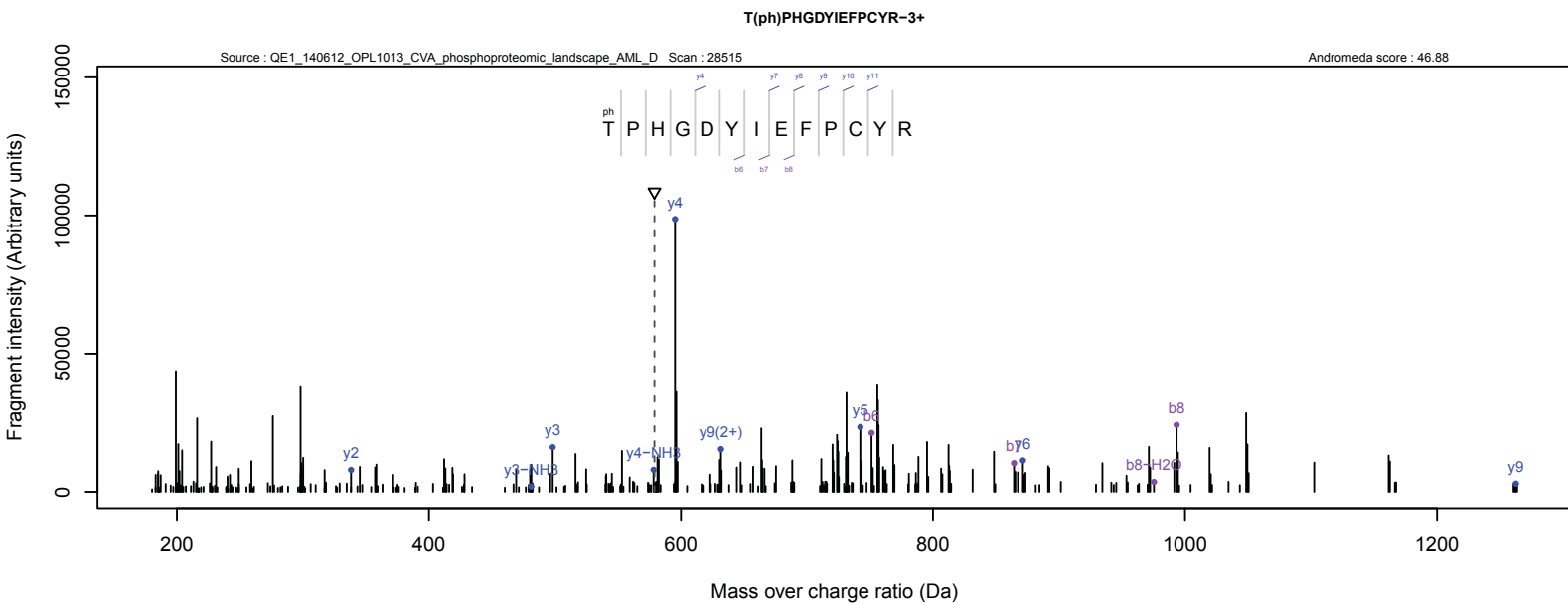


## EYPLY(ph)INQTCR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20466

Andromeda score : 46.88

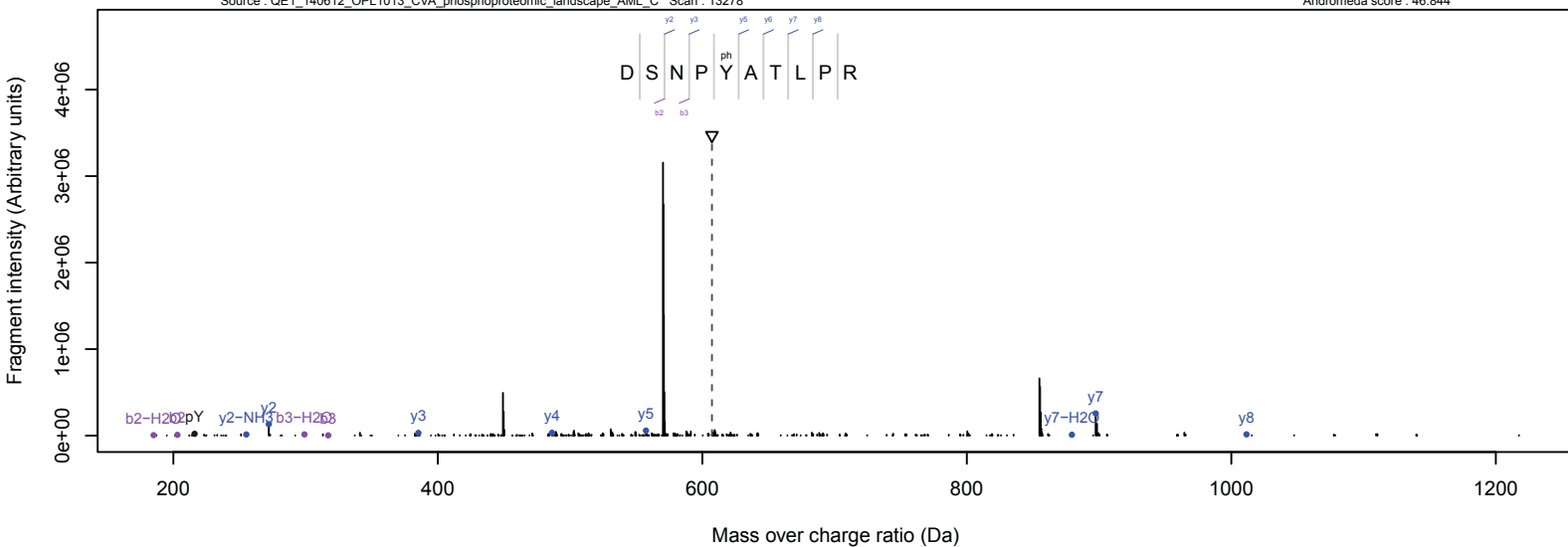




## DSNPY(ph)ATLPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 13278

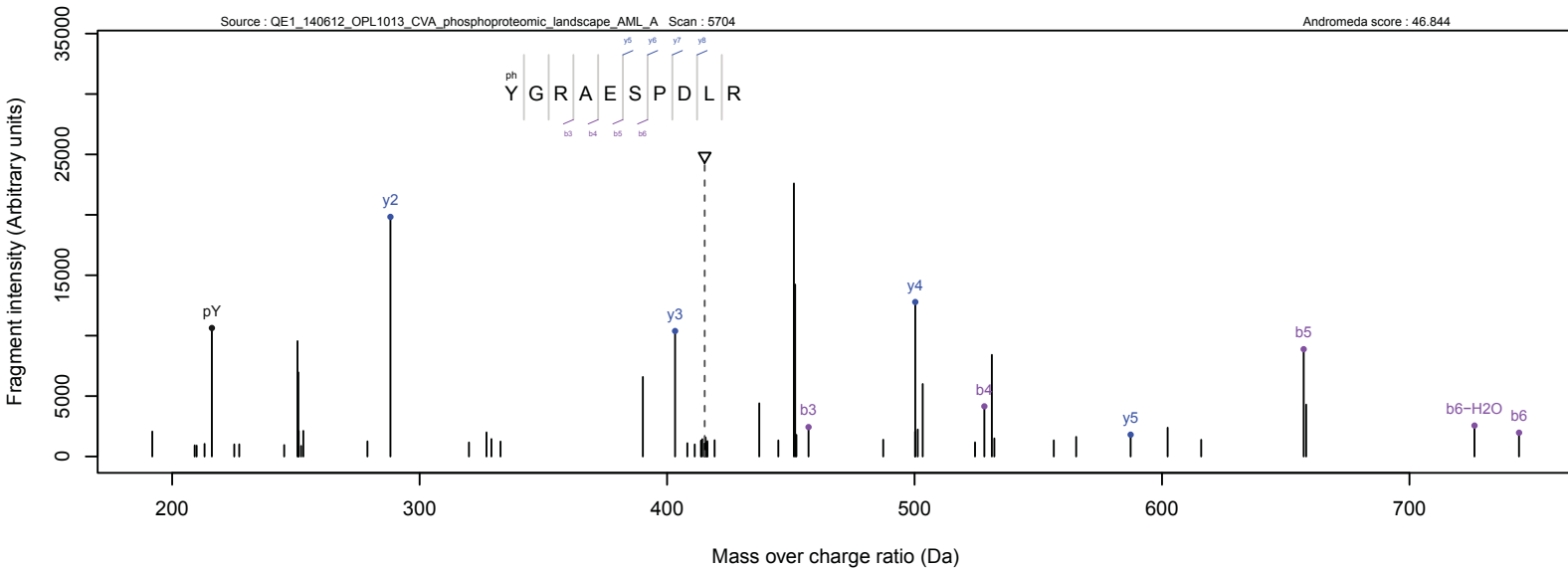
Andromeda score: 46.844



## Y(ph)GRAESPDLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 5704

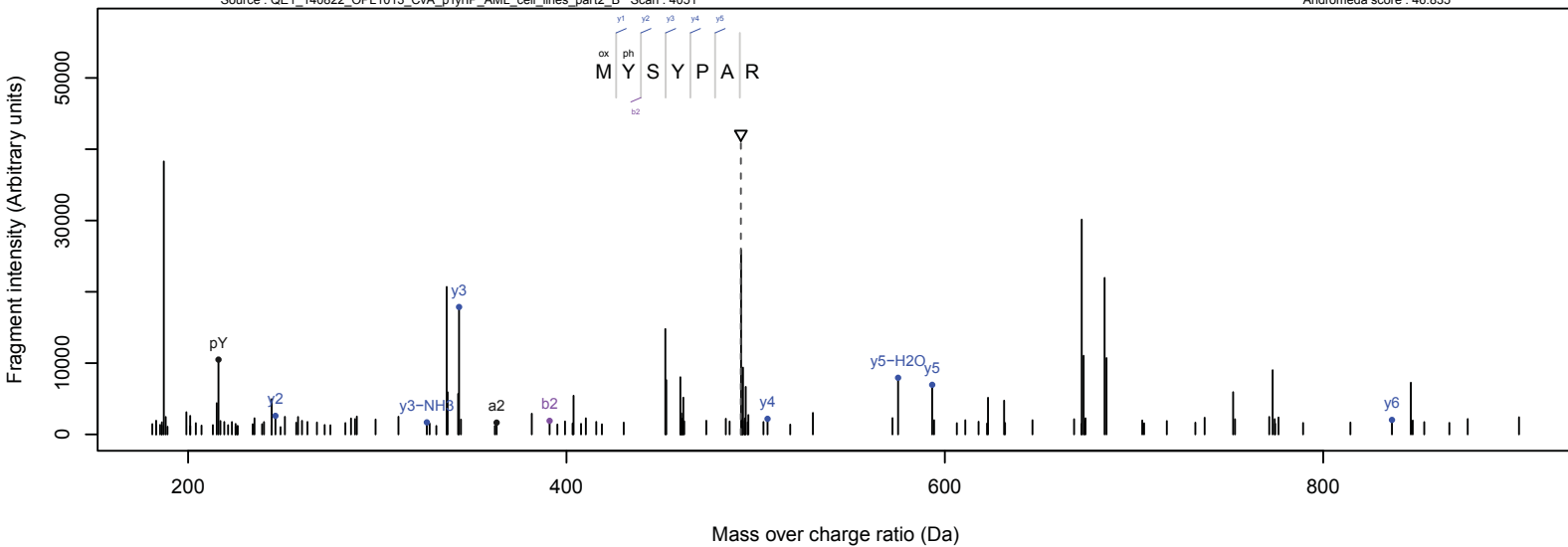
Andromeda score: 46.844



## M(ox)Y(ph)SYPAR-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 4051

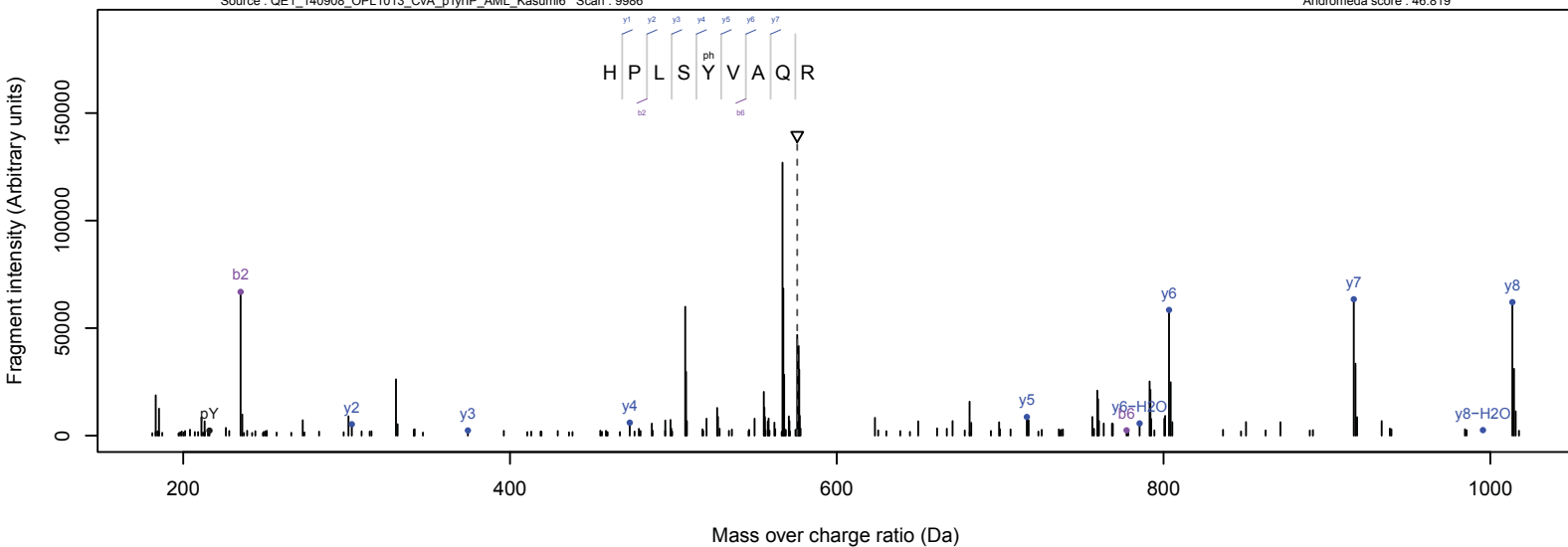
Andromeda score: 46.835



## HPLSY(ph)VAQR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 9986

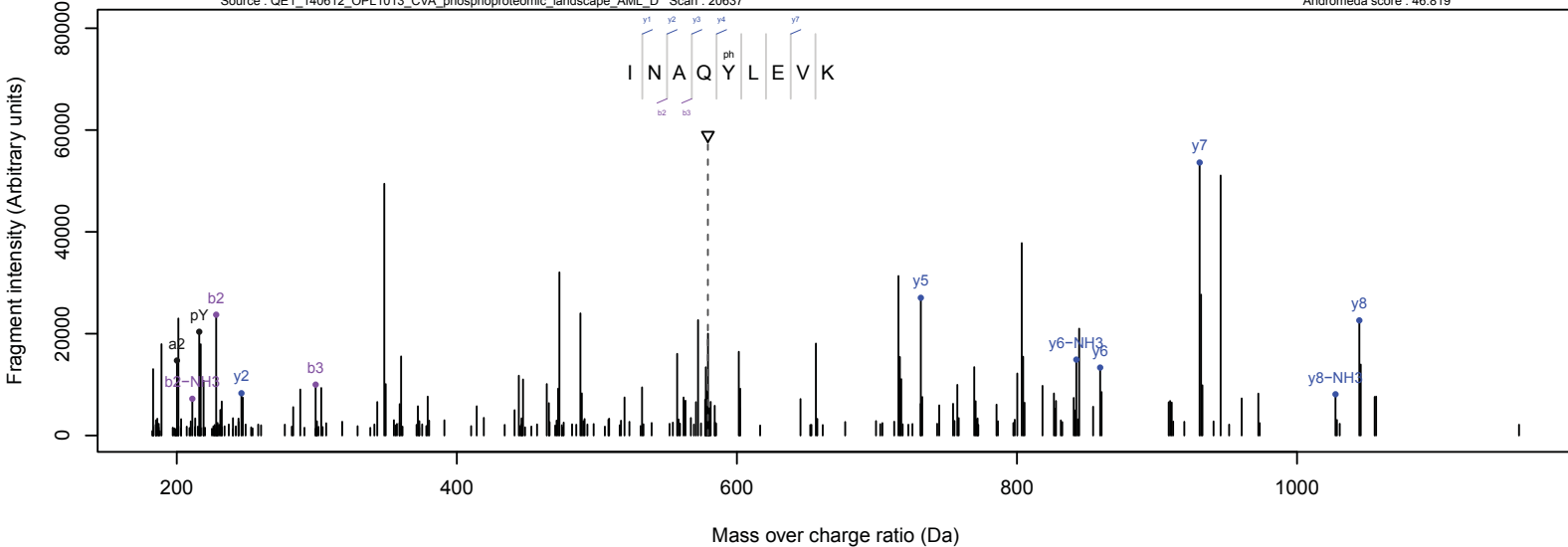
Andromeda score : 46.819



## INAQY(ph)LEVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 20637

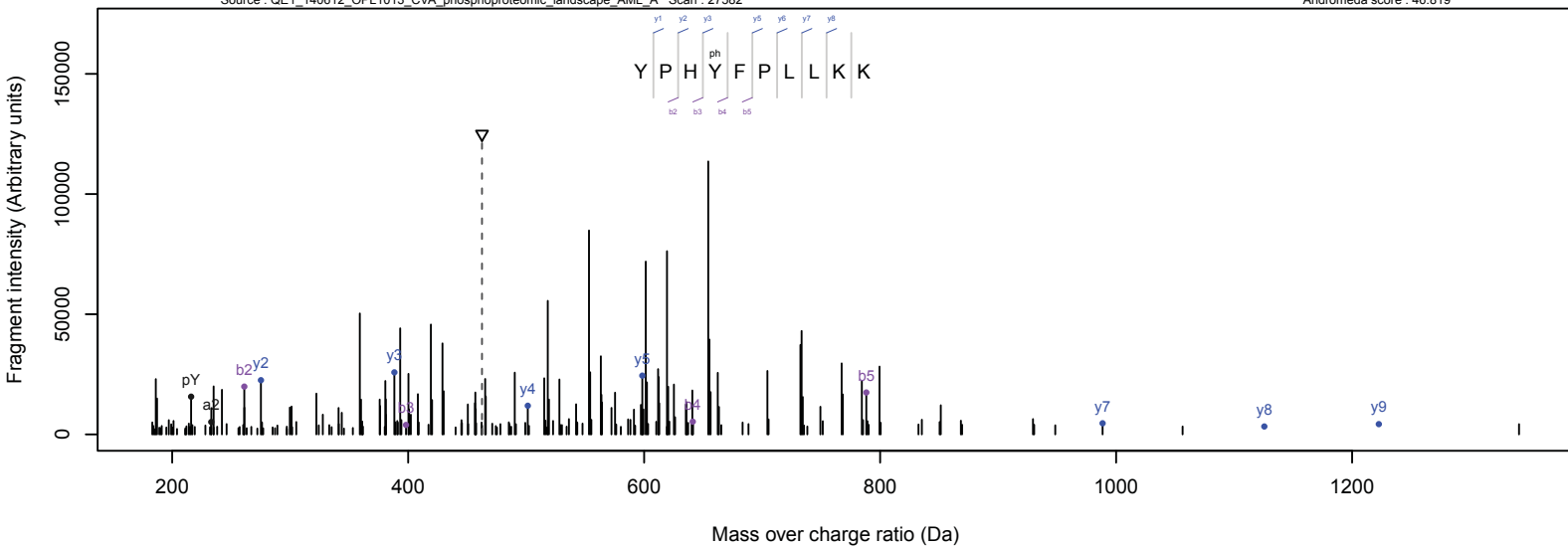
Andromeda score : 46.819



## YPHY(ph)FPLLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27582

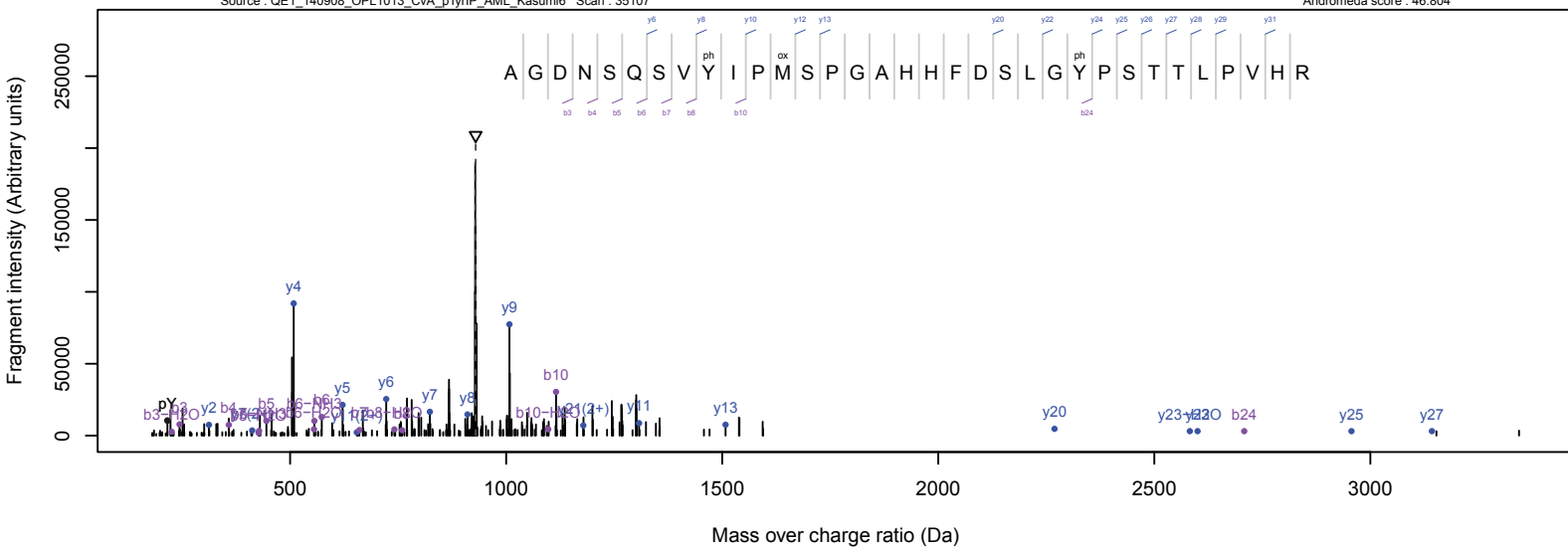
Andromeda score : 46.819



AGDNSQSVY(ph)IPM(ox)SPGAHHFDSLGY(ph)PSTTLPVHR-4+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan: 35107

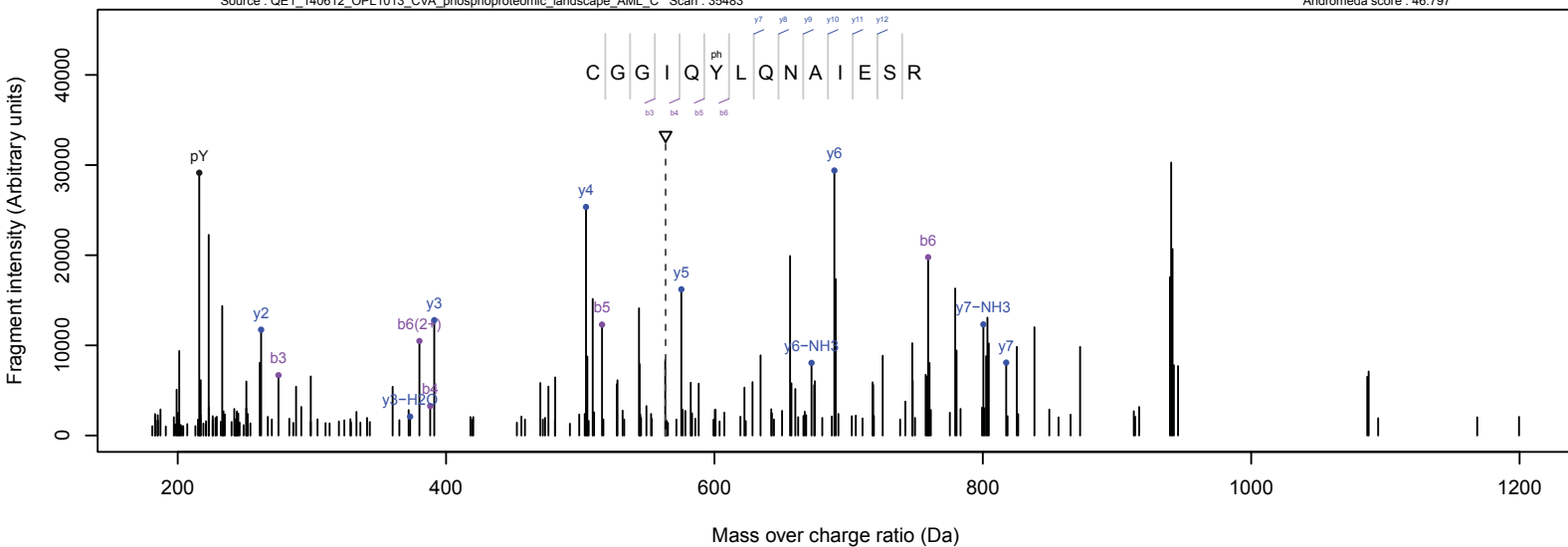
Andromeda score: 46.804



CGGIQY(ph)LQNAIESR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 35483

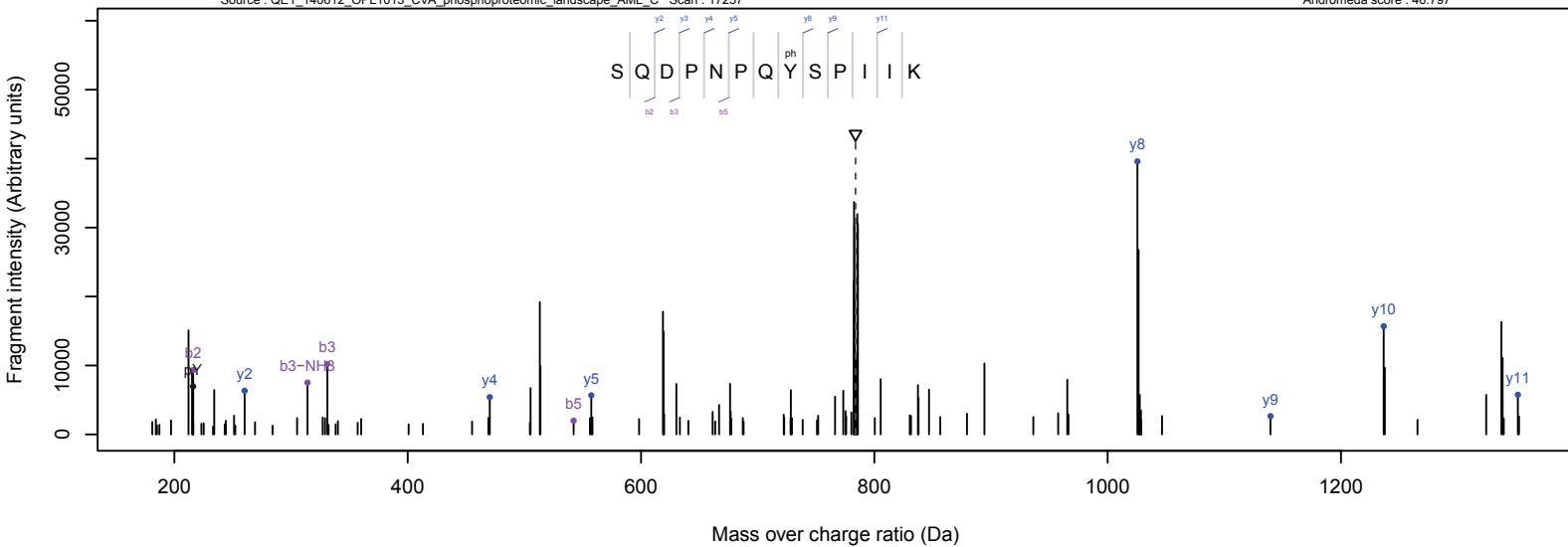
Andromeda score: 46.797



SQDPNPQY(ph)SPIIK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17257

Andromeda score: 46.797

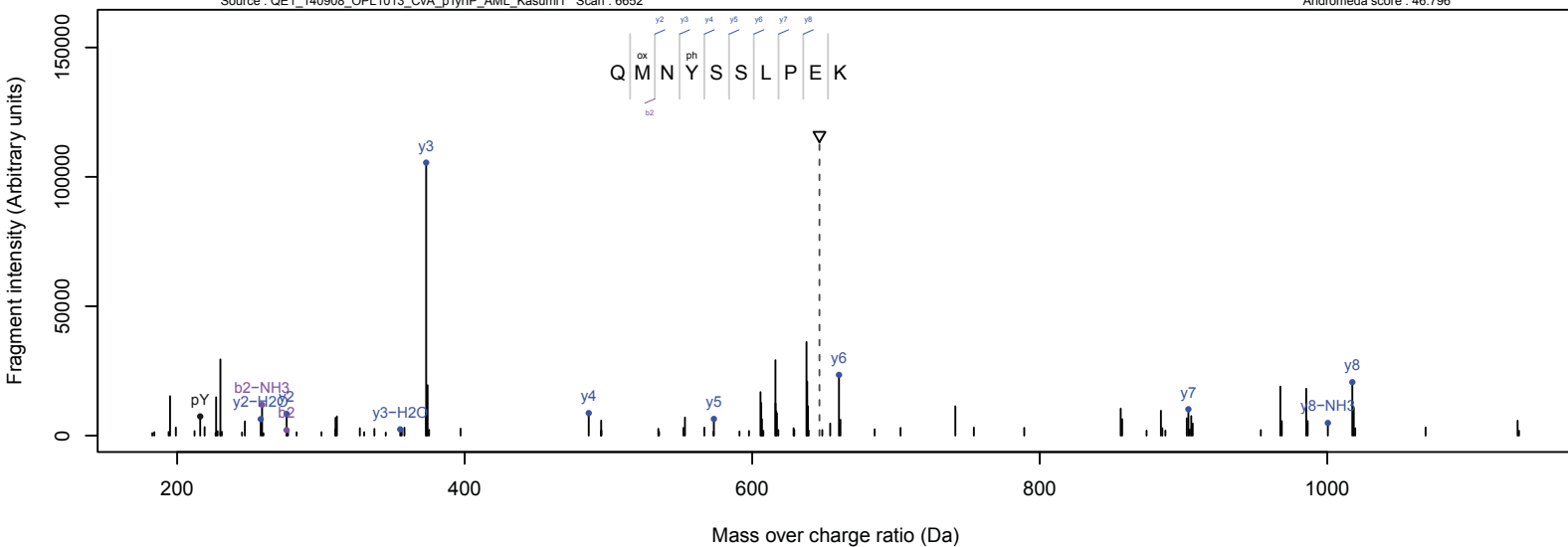




QM(ox)NY(ph)SSLPEK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi1 Scan : 6652

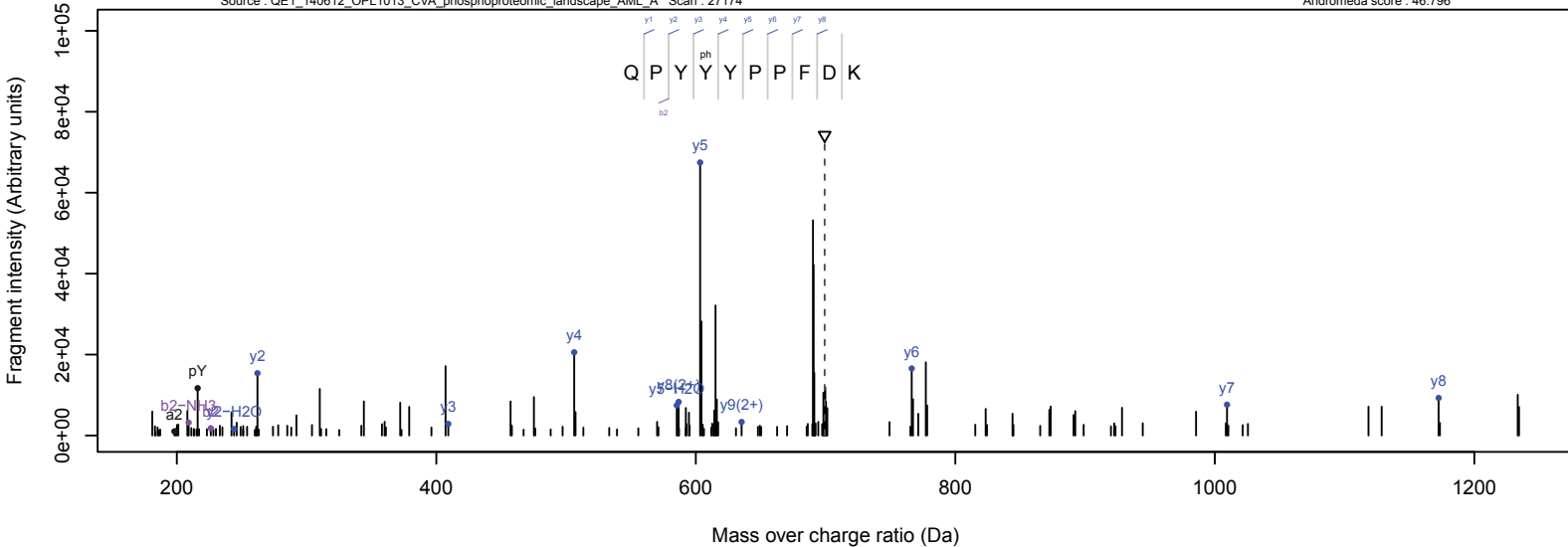
Andromeda score : 46.796



QPYY(ph)YPPFDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 27174

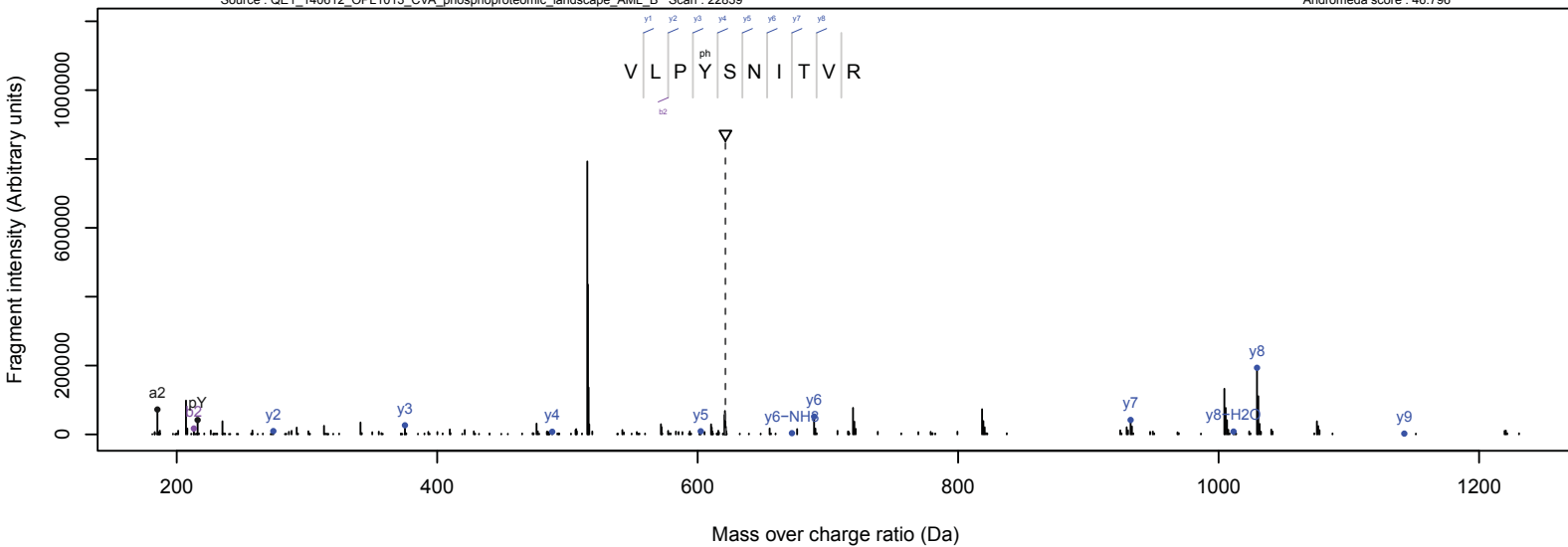
Andromeda score : 46.796



VLPY(ph)SNITVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22839

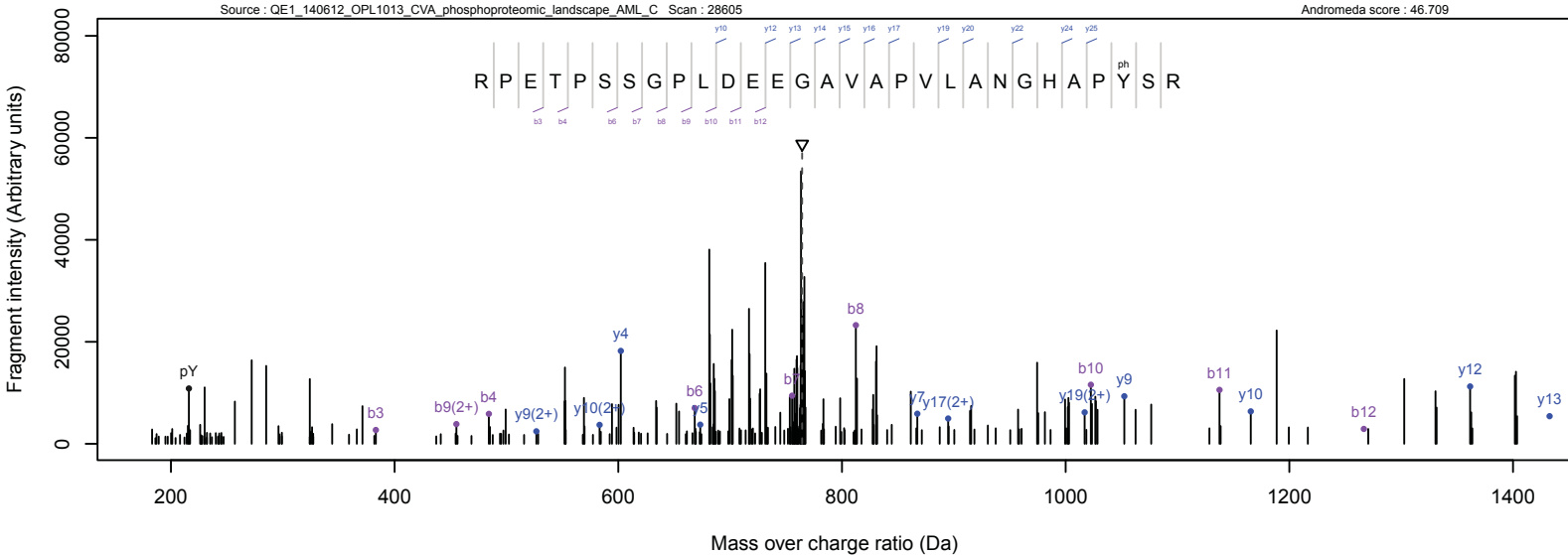
Andromeda score : 46.796



RPETPSSGLDEEGAVPVLANGHAPY<sup>ph</sup>SR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 28605

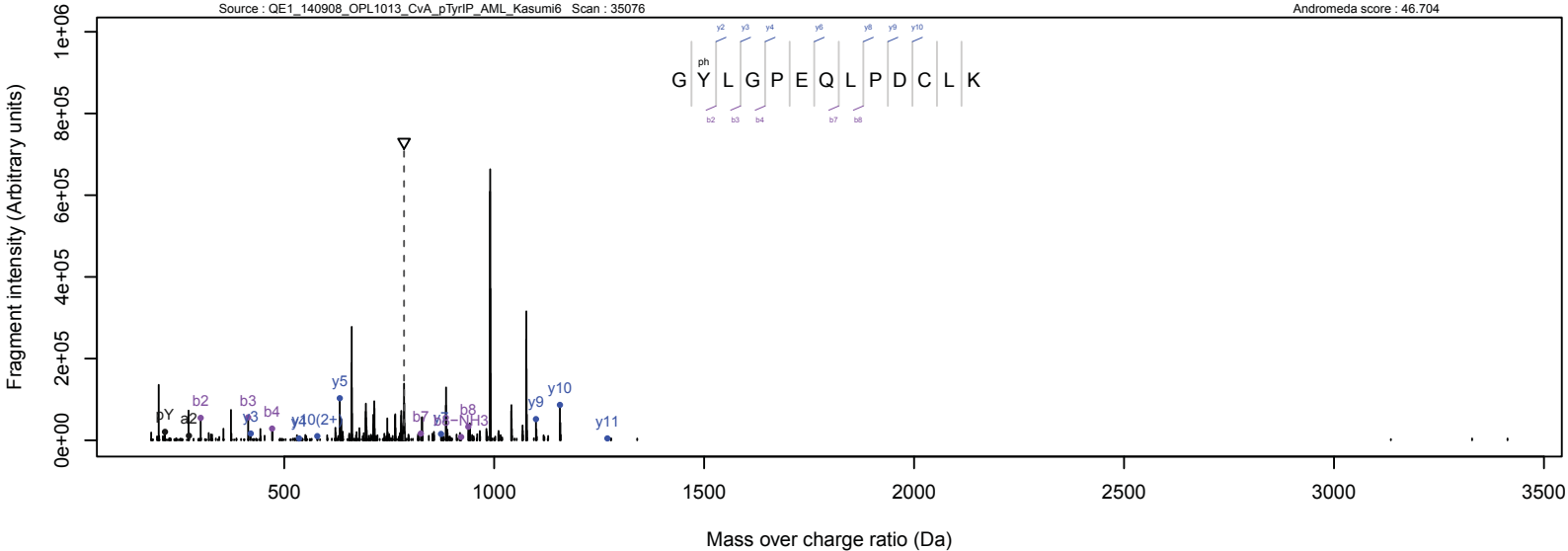
Andromeda score : 46.709



GY<sup>ph</sup>LGPEQLPDCLK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 35076

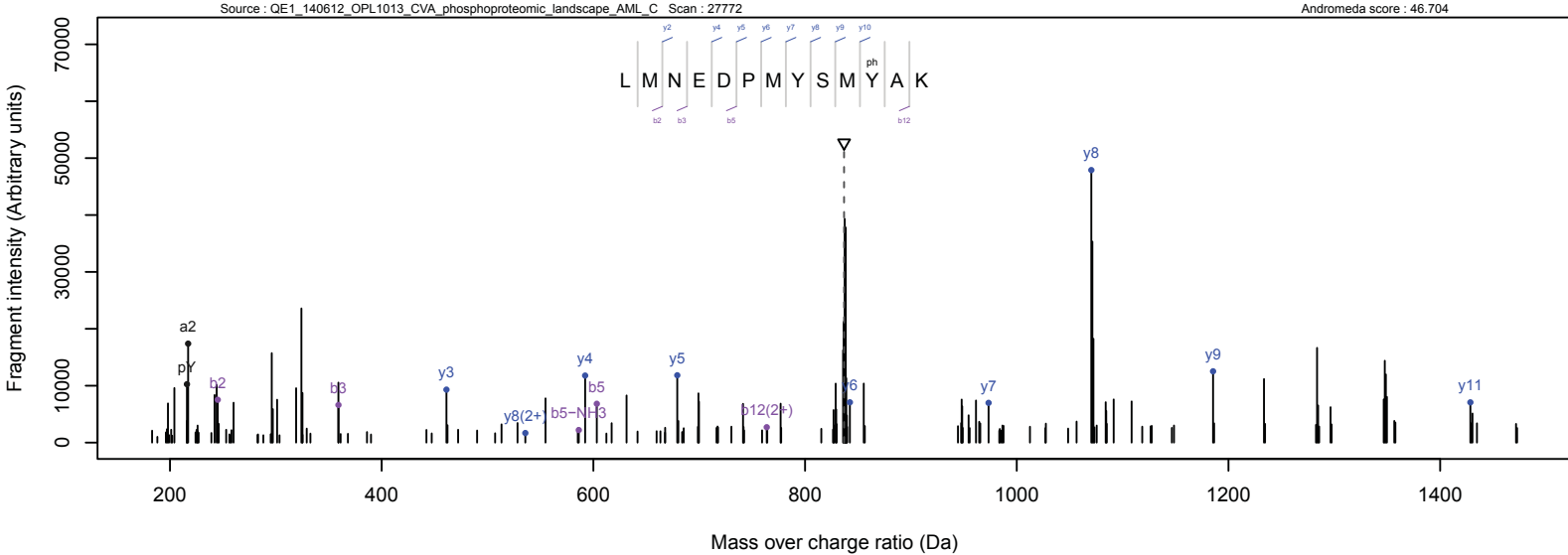
Andromeda score : 46.704



LMNEDPMSMY<sup>ph</sup>AK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27772

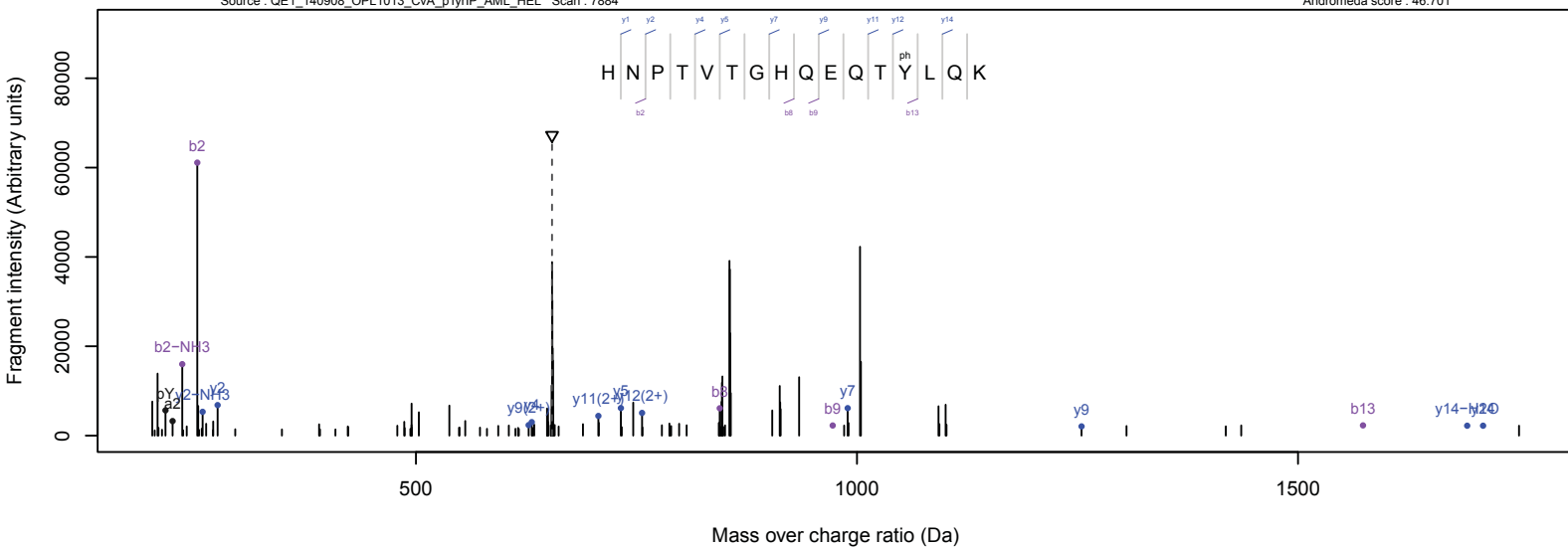
Andromeda score : 46.704



HNPTVTGHQEQT<sup>(ph)</sup>LQK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 7884

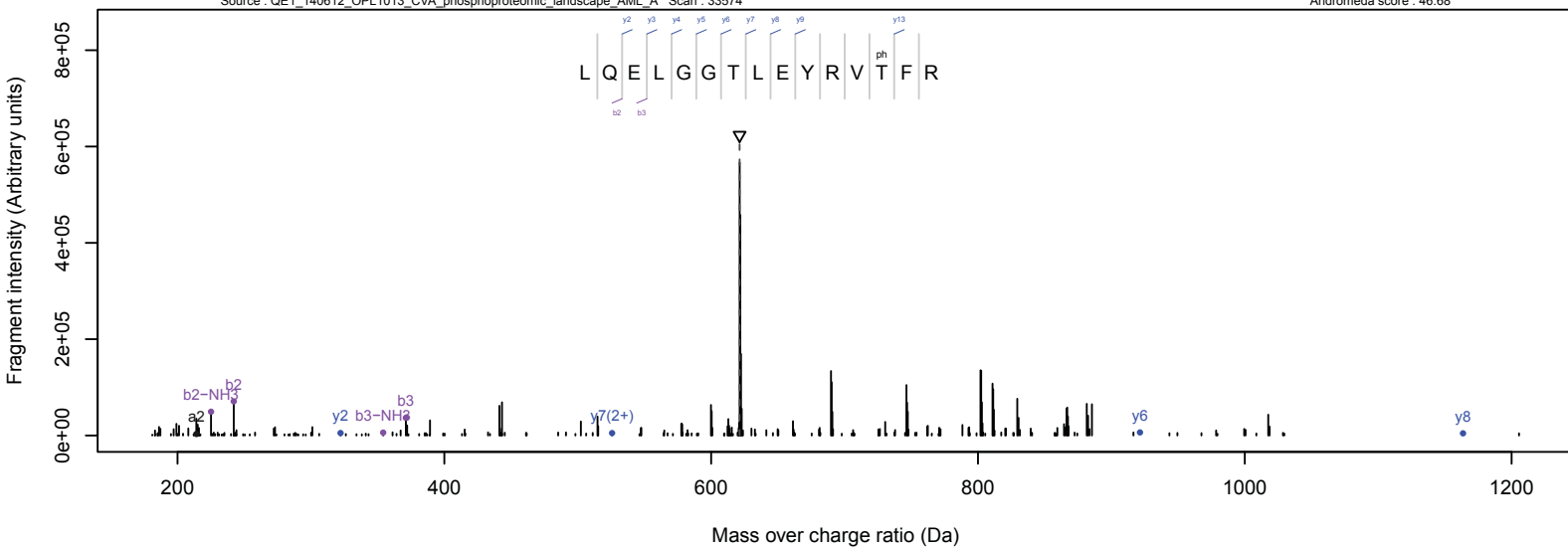
Andromeda score : 46.701



LQELGGTLEYR<sup>(ph)</sup>VTFR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 33574

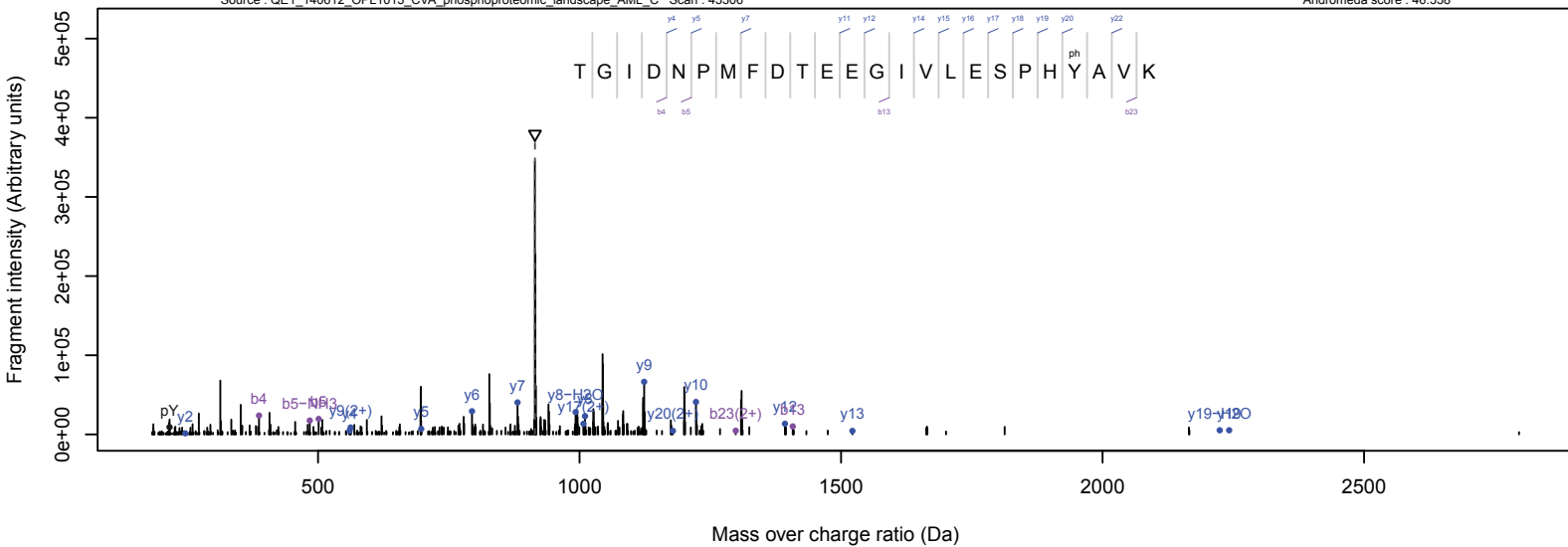
Andromeda score : 46.68



TGIDNPMFDTEEGIVLES<sup>(ph)</sup>PHYAVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43306

Andromeda score : 46.558

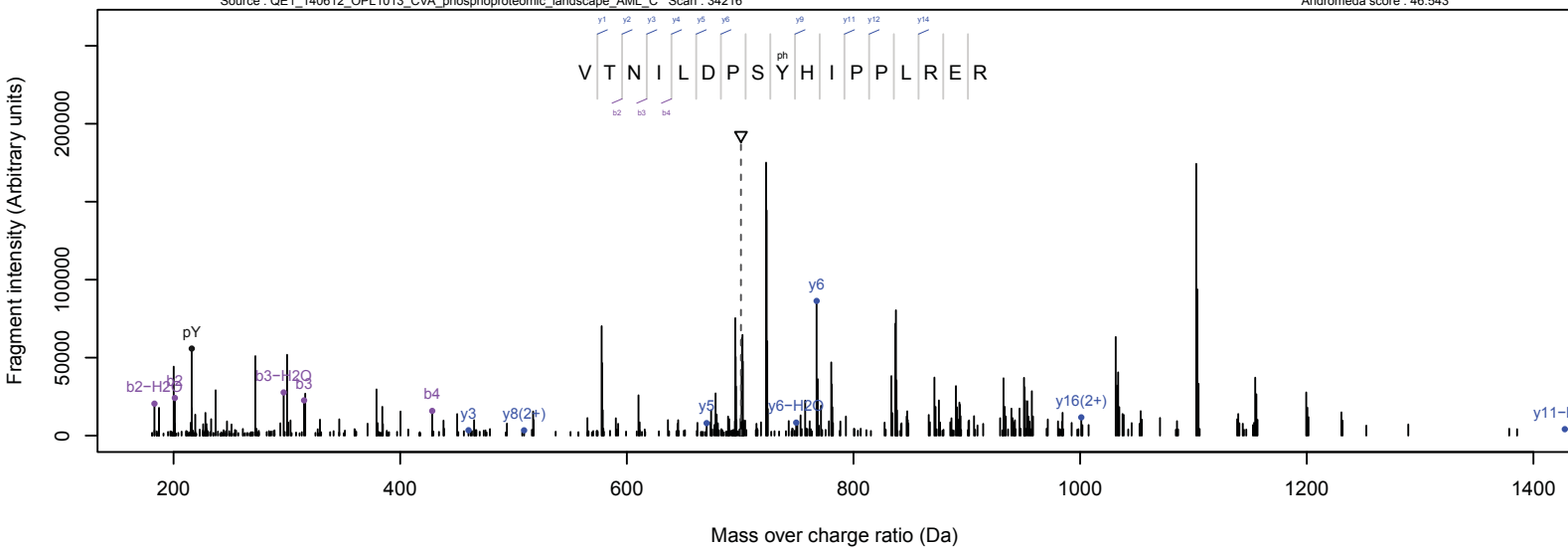




## VTNILDPSY(ph)HIPPLRER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34216

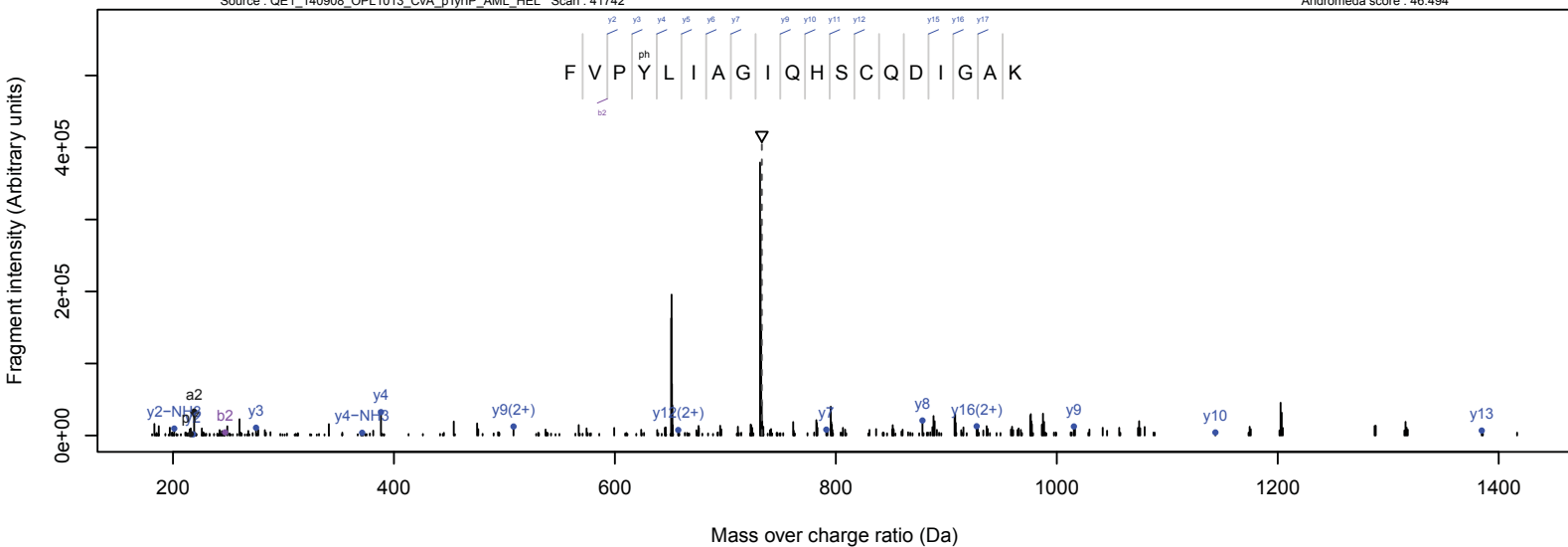
Andromeda score : 46.543



## FVPY(ph)LIAGIQHSCQDIGAK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 41742

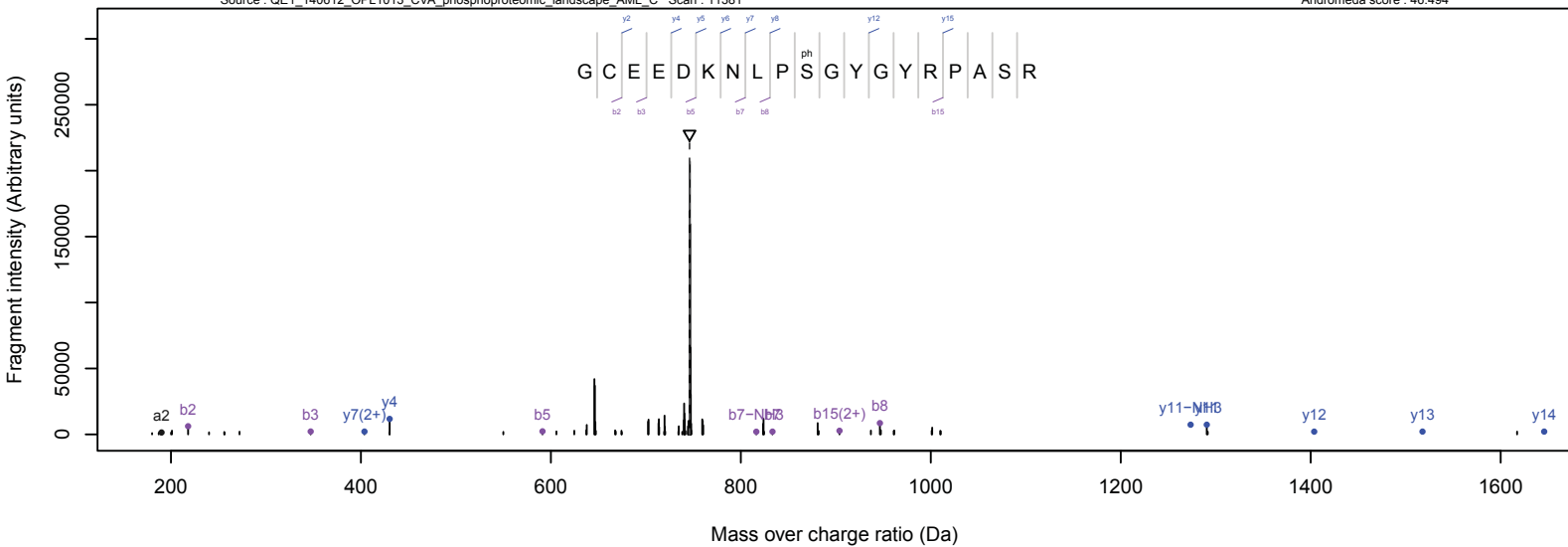
Andromeda score : 46.494



## GCEEDKNLPS(ph)GYRPARSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11381

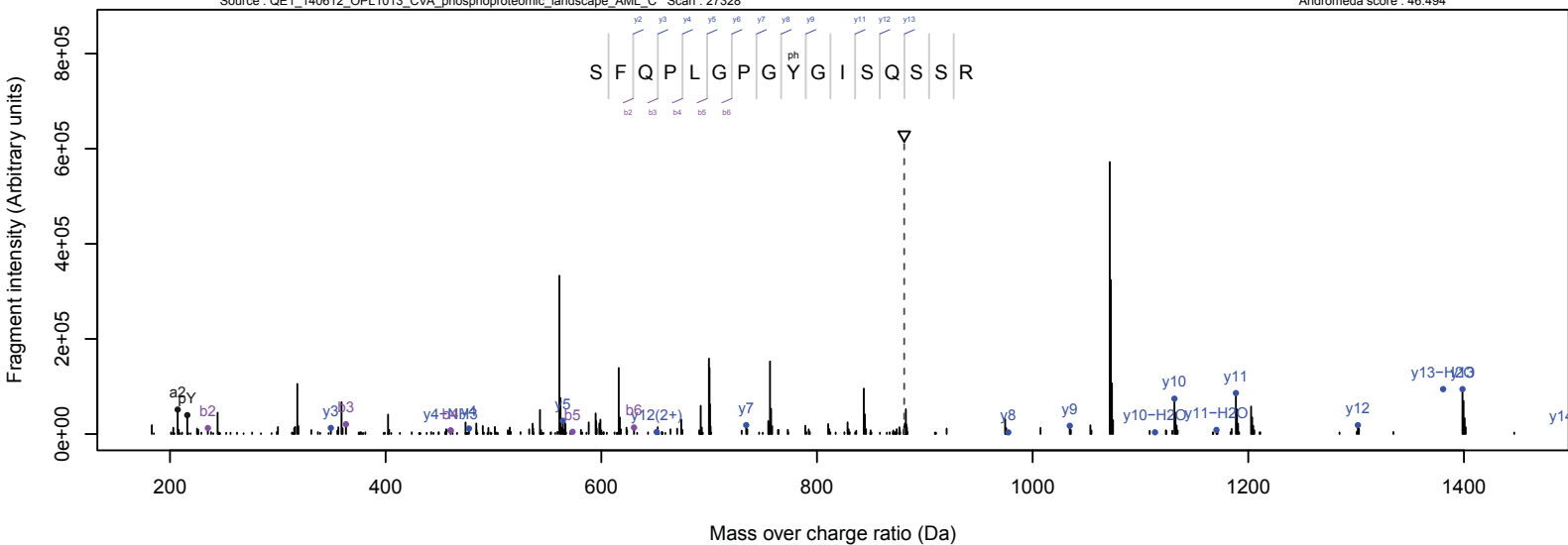
Andromeda score : 46.494



SFQPLGPGY(ph)GISQSSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27328

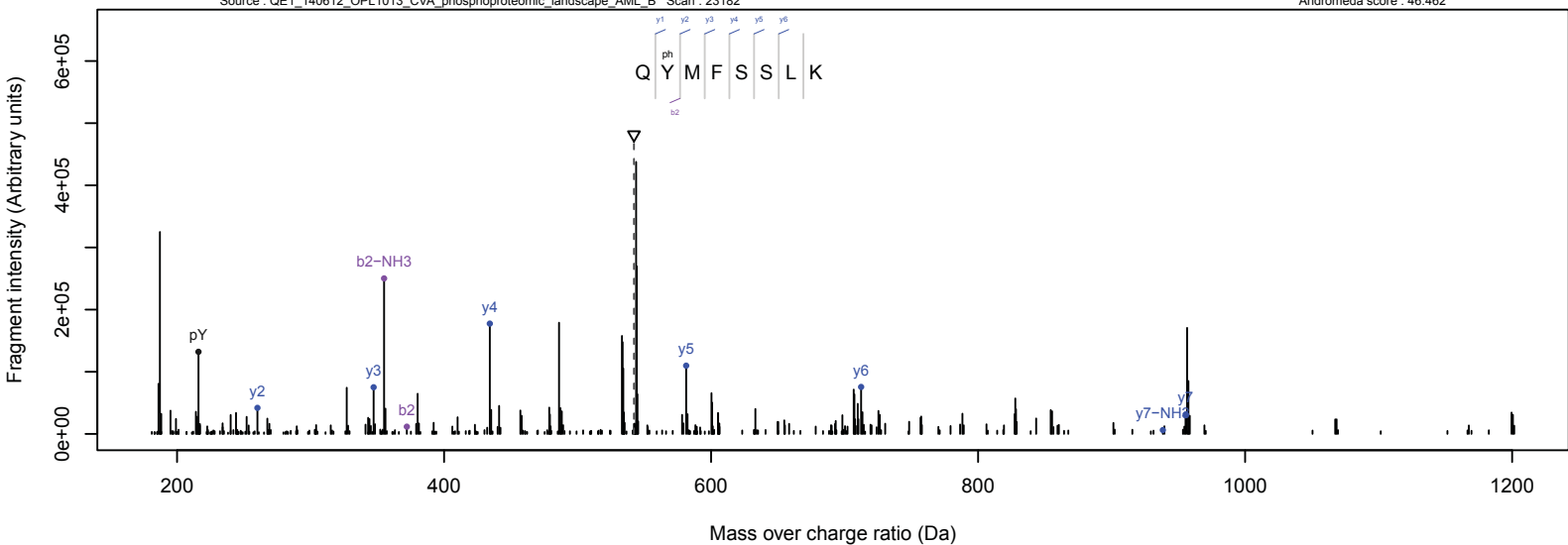
Andromeda score : 46.494



QY(ph)MFSSLK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23182

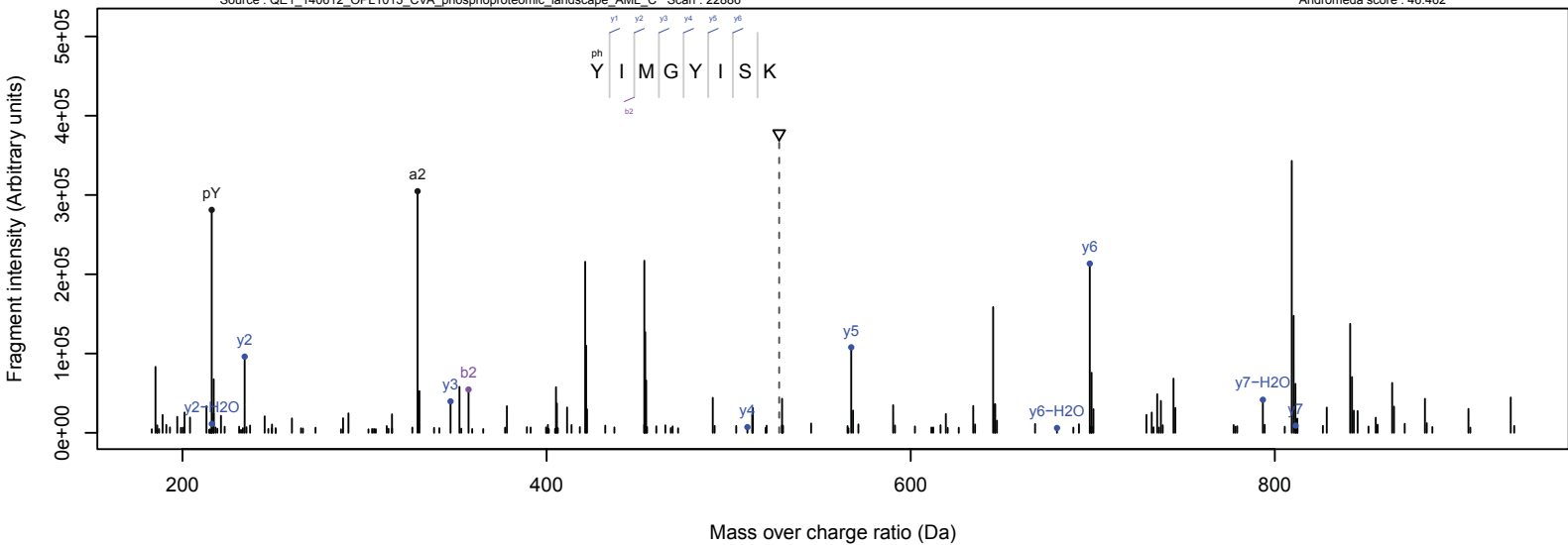
Andromeda score : 46.462



Y(ph)IMGYISK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22886

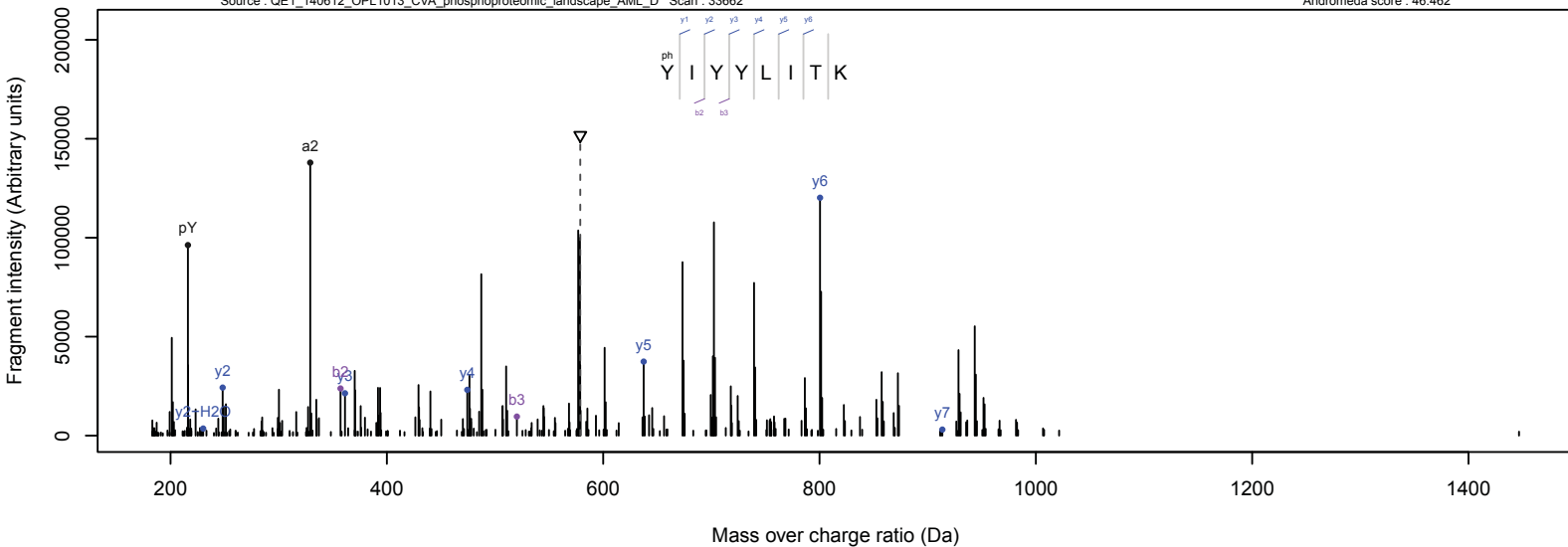
Andromeda score : 46.462



Y(ph)IYYLITK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 33662

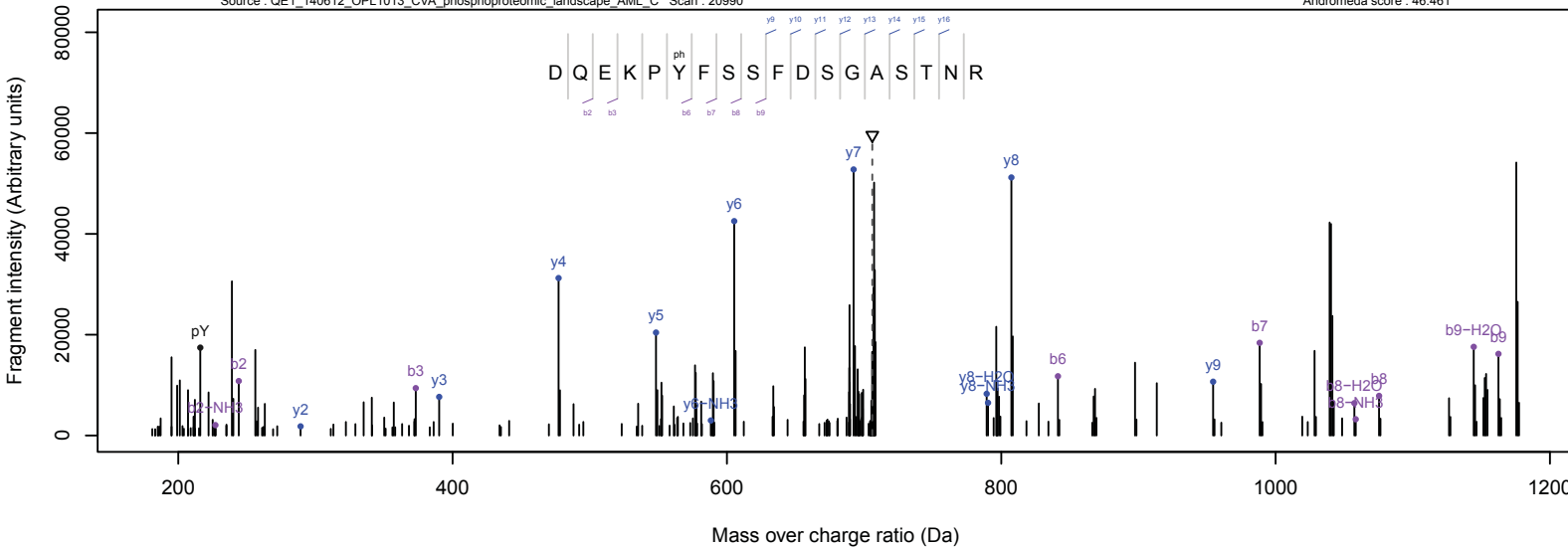
Andromeda score : 46.462



DQEKP(ph)FSSFDGASTNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20990

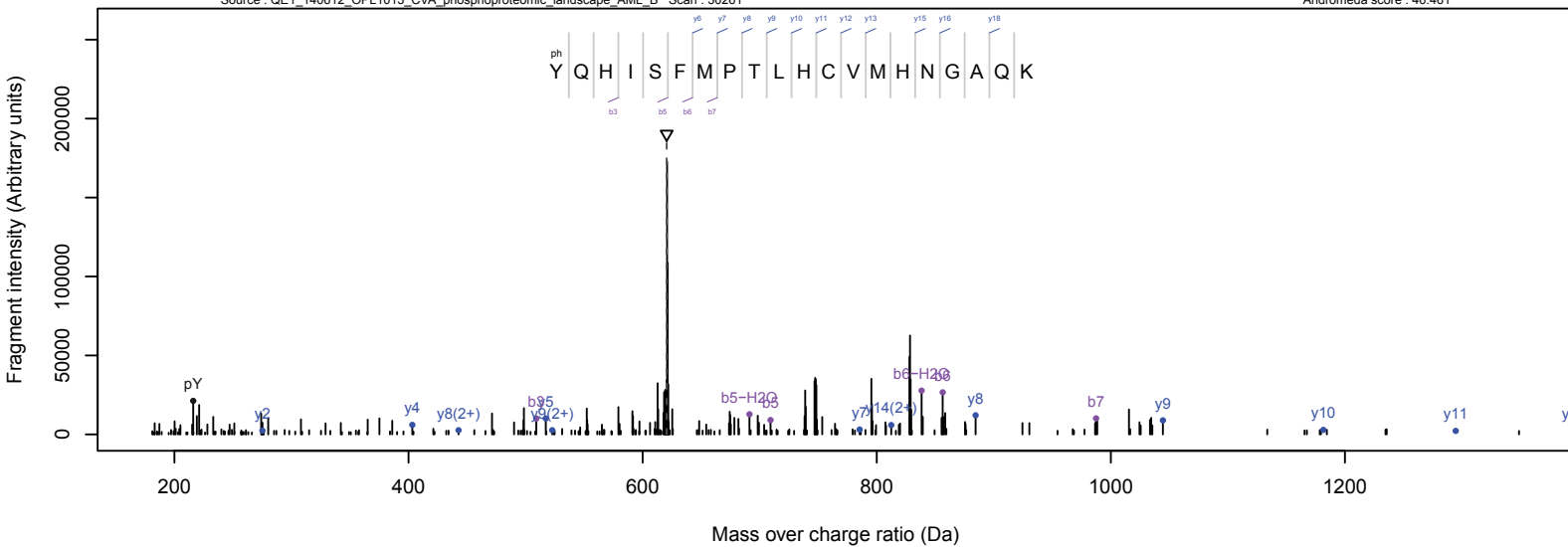
Andromeda score : 46.461



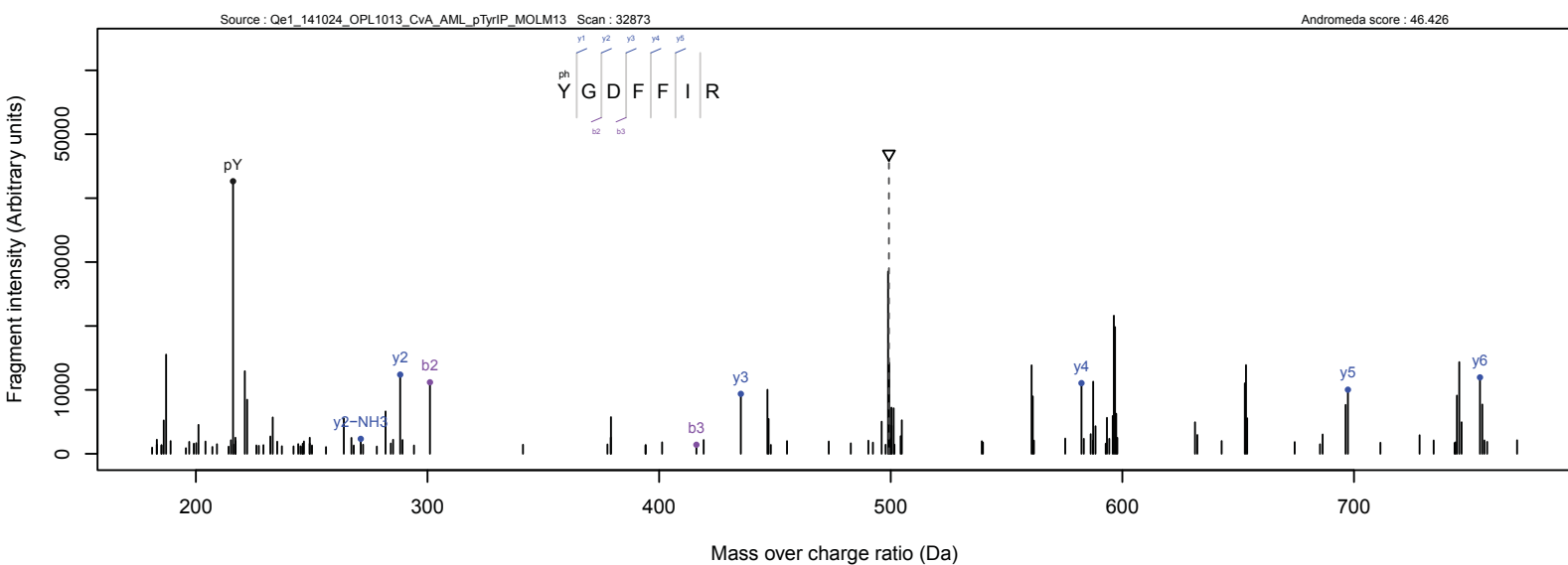
Y(ph)QHISFMPTLHCVMHNGAQK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30261

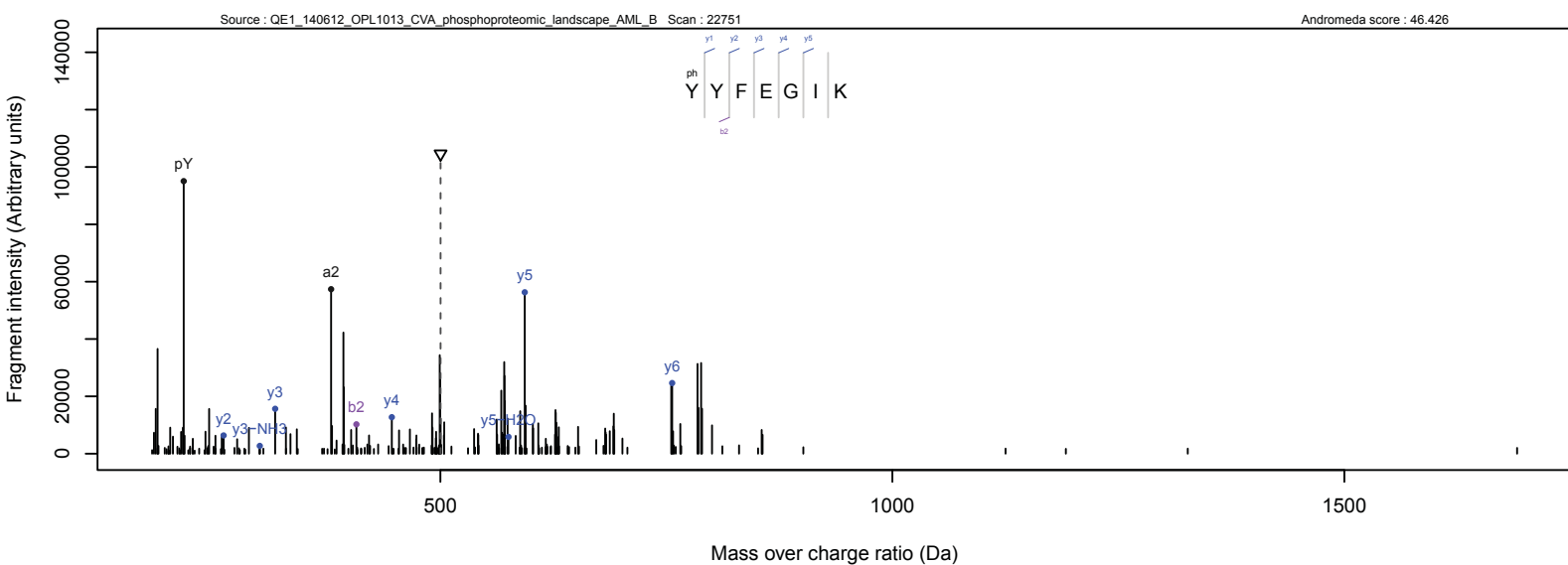
Andromeda score : 46.461



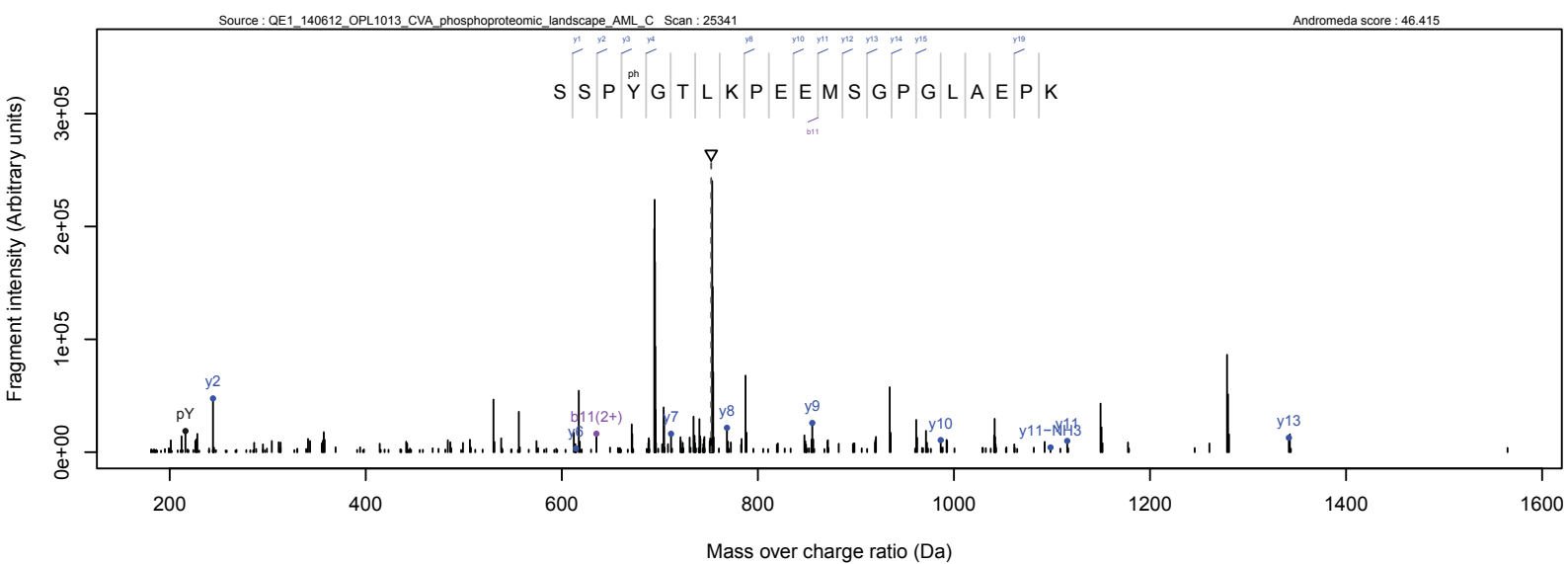
## Y(ph)GDF FIR-2+



## Y(ph)YFEGIK-2+



## SSPY(ph)GTLKPEEMSGPGLAEPK-3+



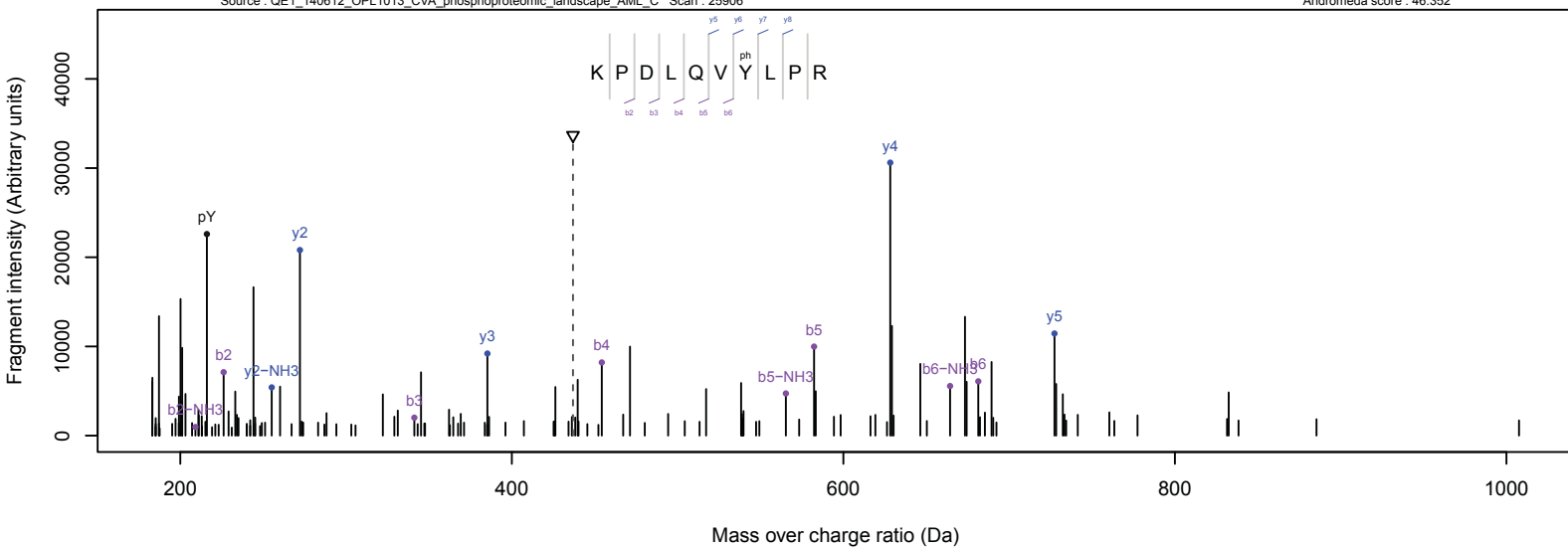




**KPDLQVY(ph)LPR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25906

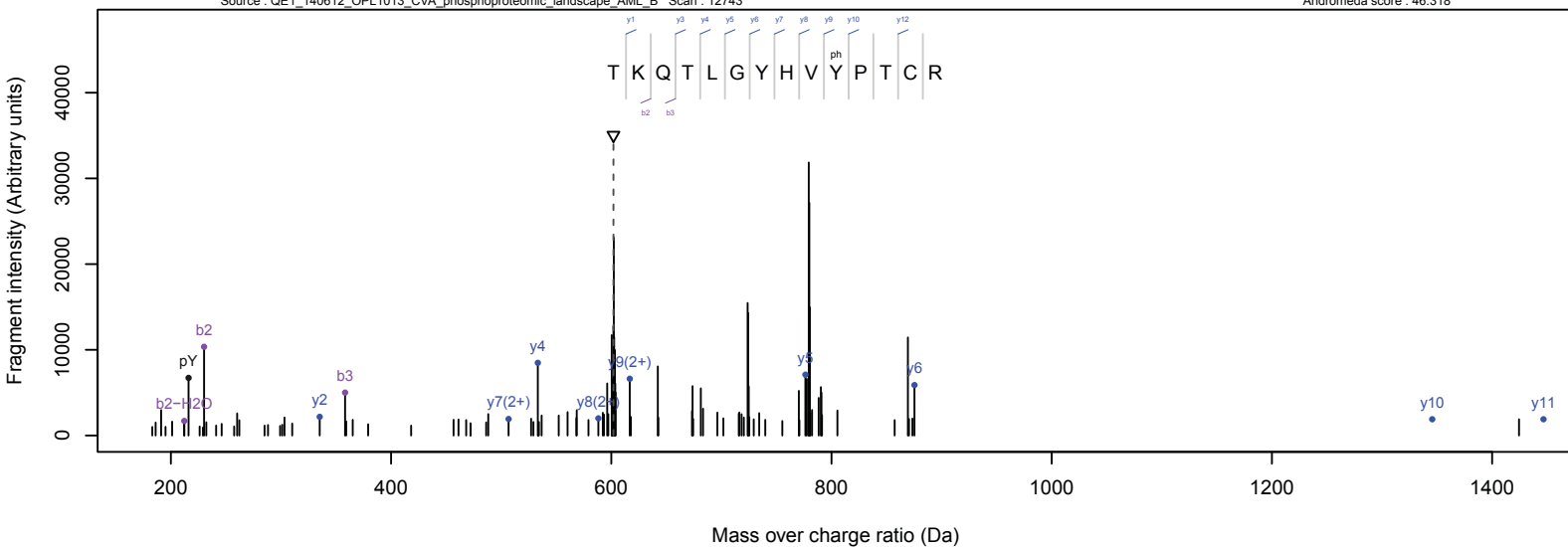
Andromeda score : 46.352



**TKQTLGYHVY(ph)PTCR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12743

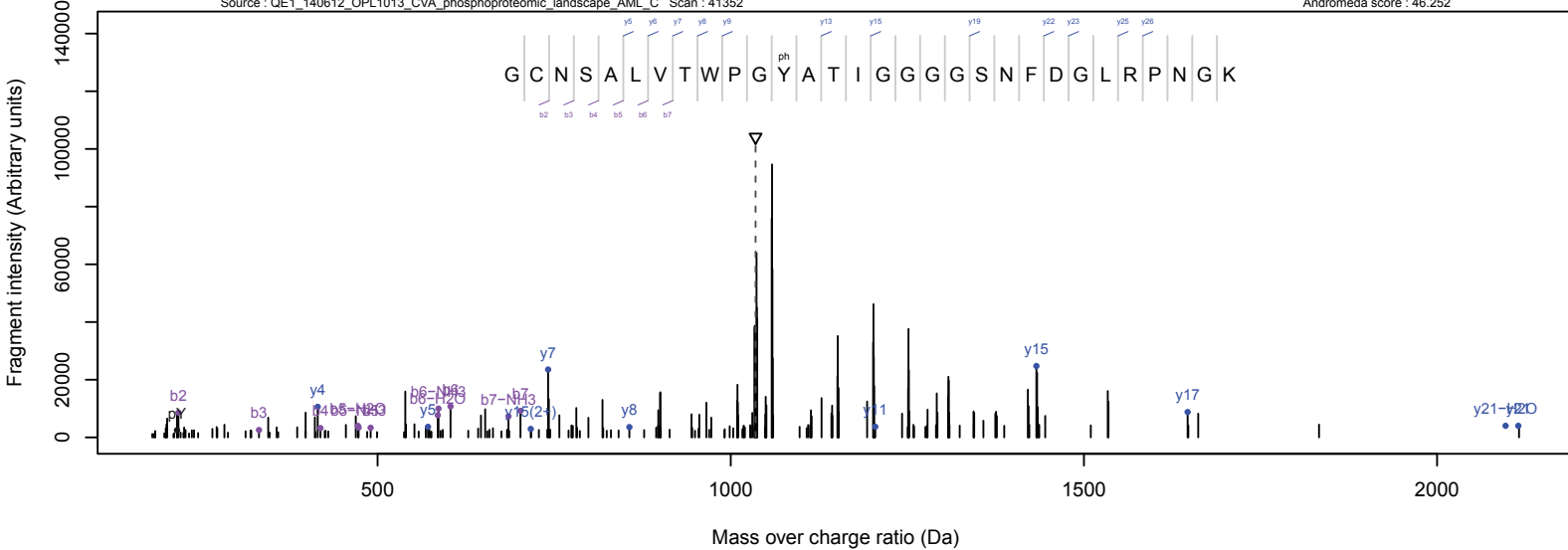
Andromeda score : 46.318

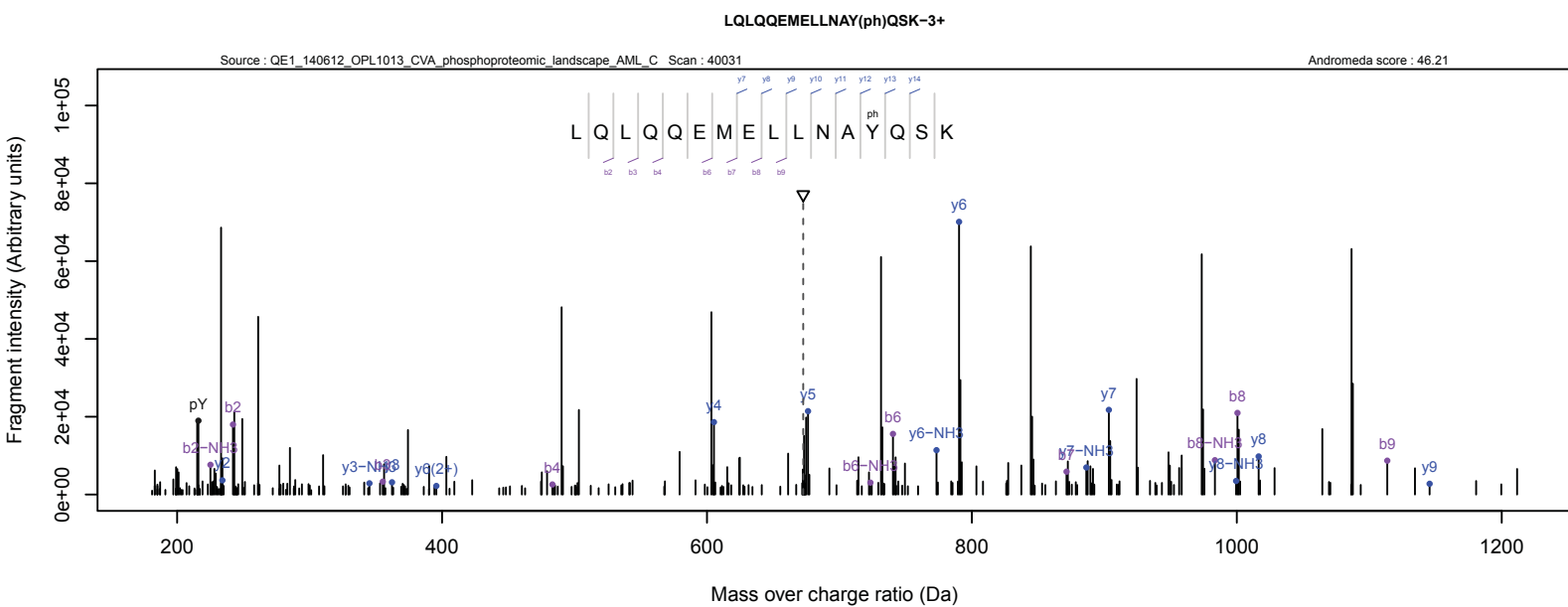
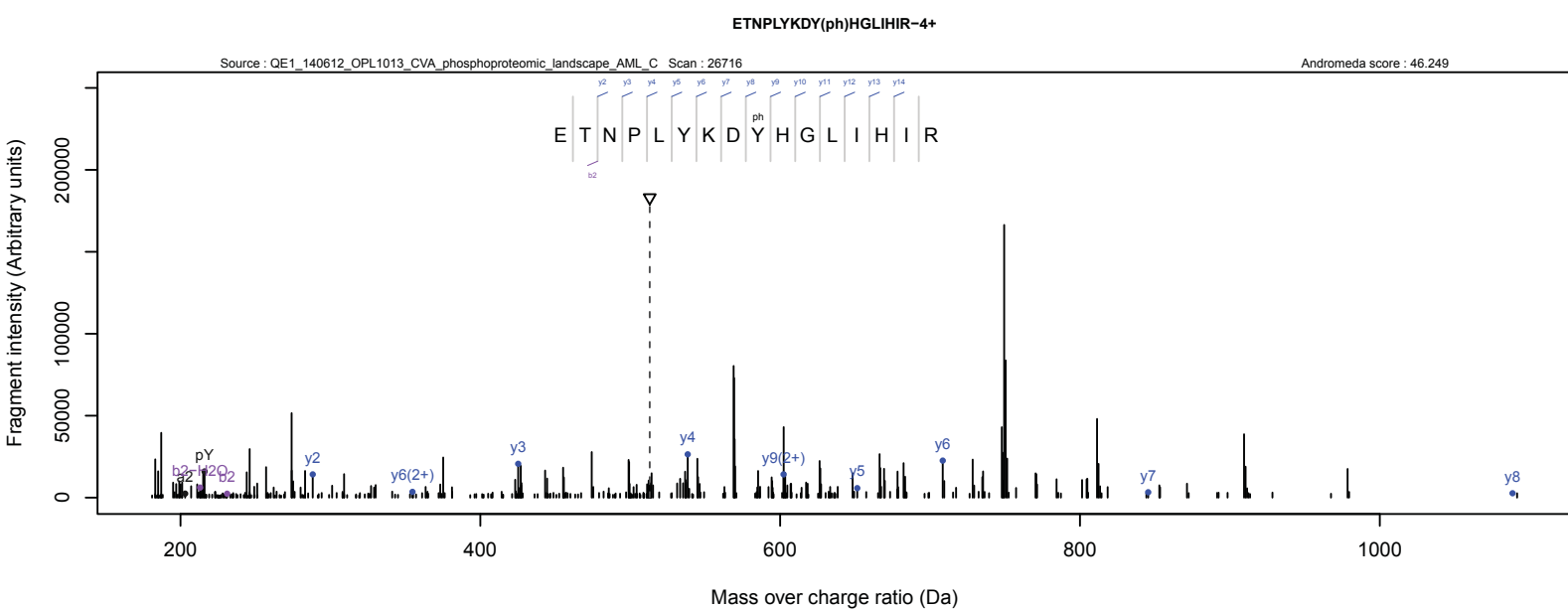
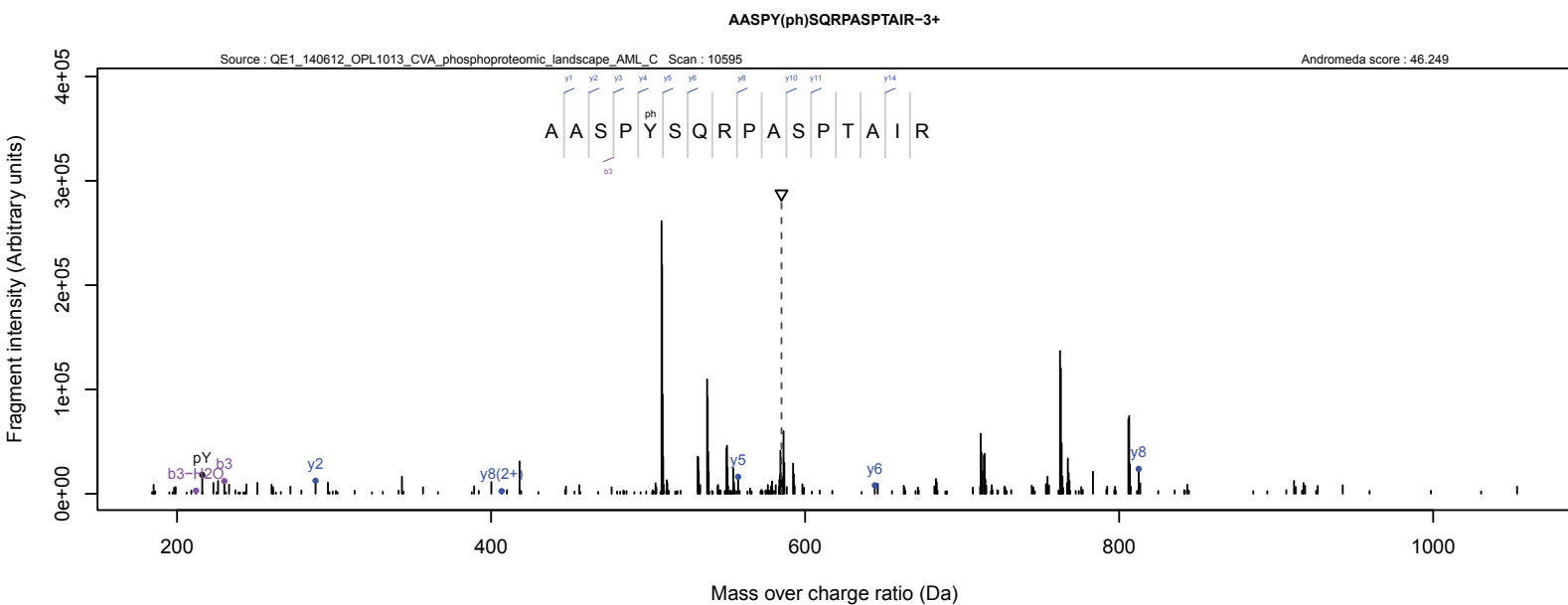


**GCNSALVTPGY(ph)ATIGGGGSNFDGLRPN GK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 41352

Andromeda score : 46.252

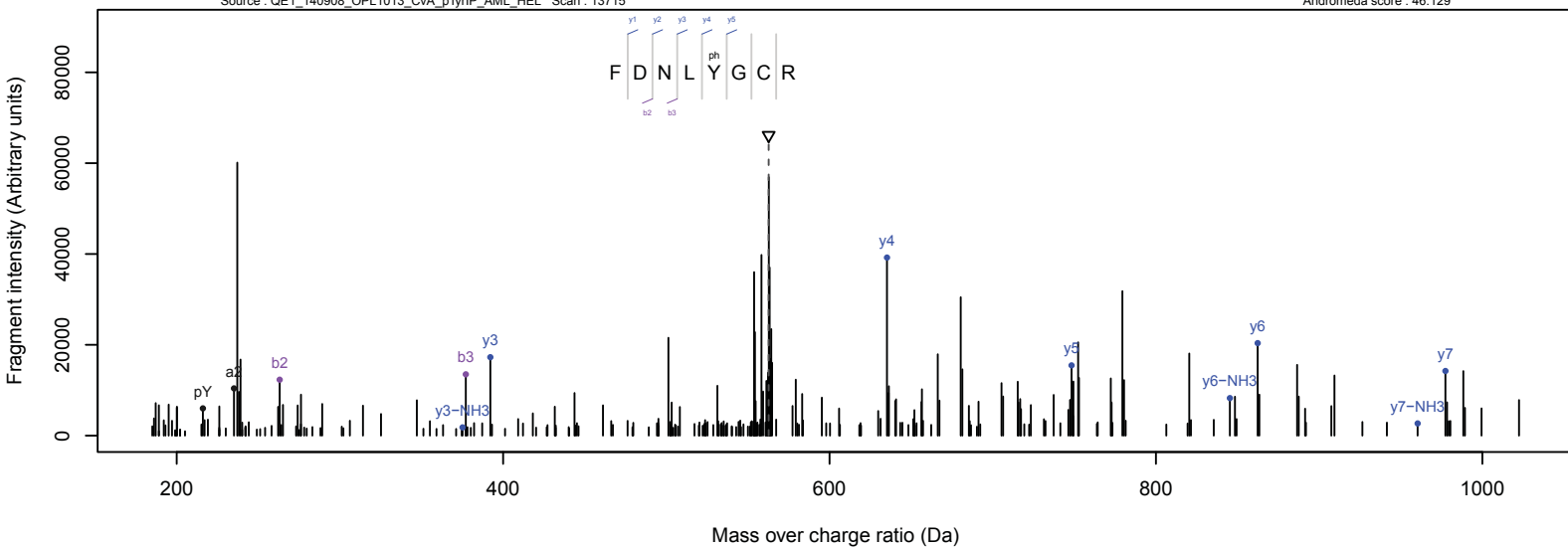




## FDNLY(ph)GCR-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 13715

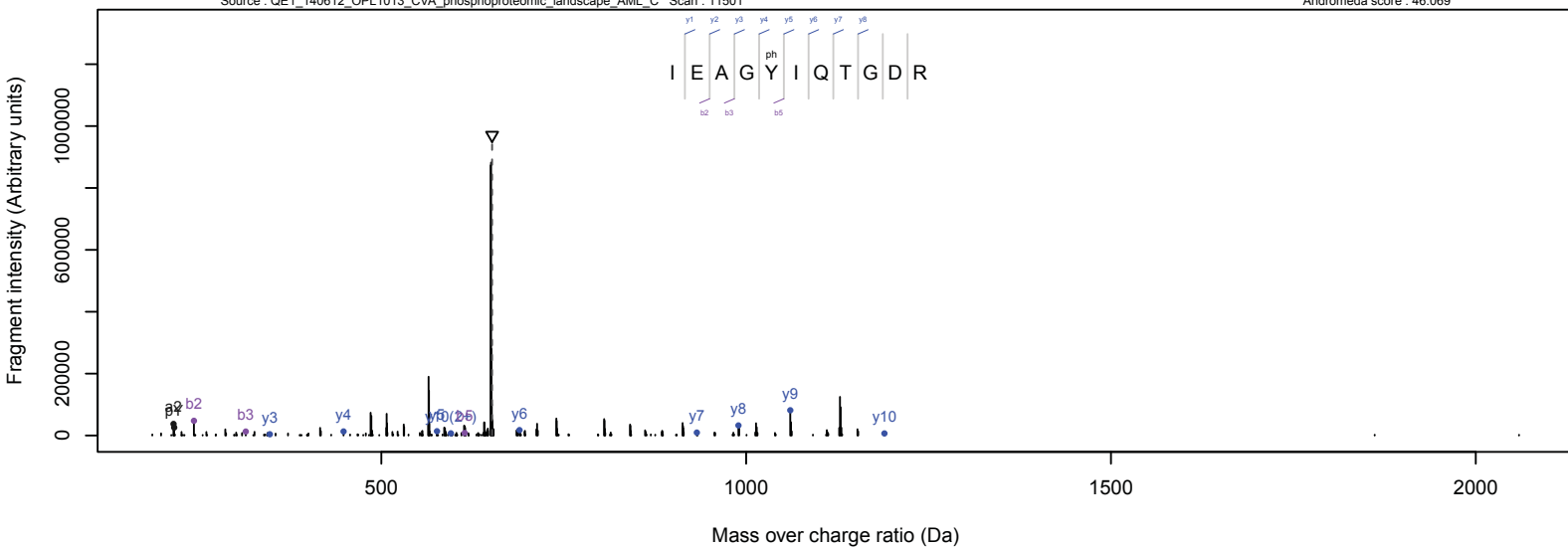
Andromeda score : 46.129



## IEAGY(ph)IQTGDR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11501

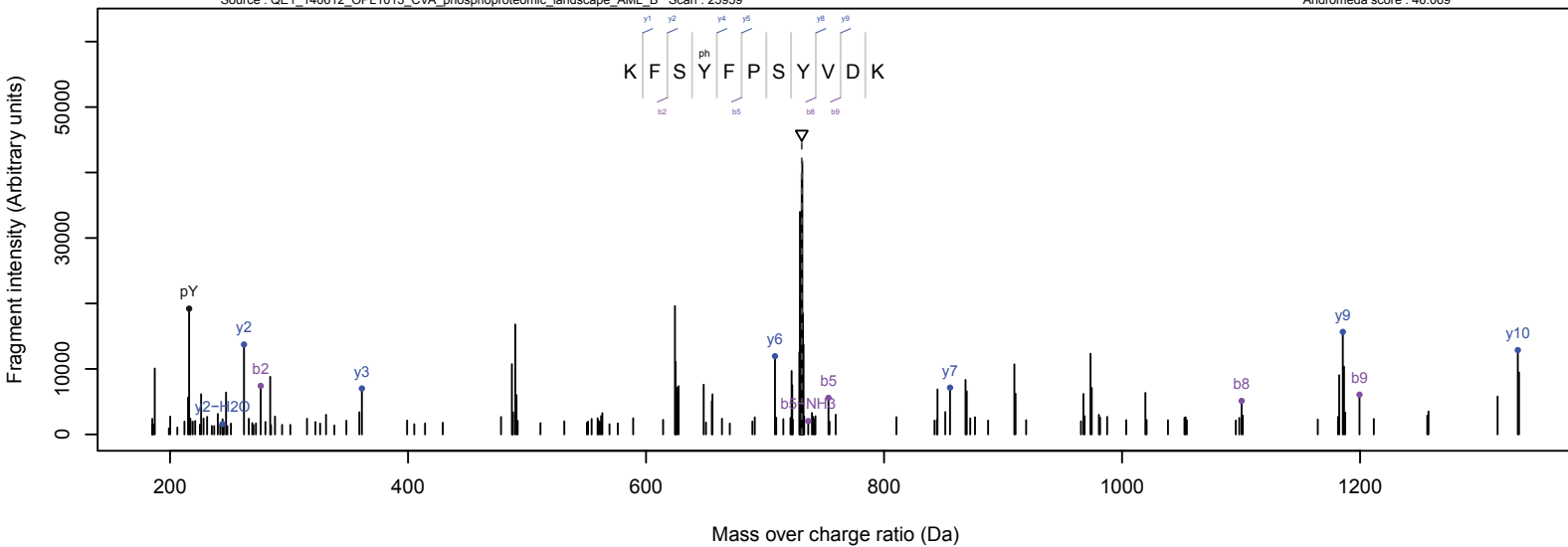
Andromeda score : 46.069



## KFSY(ph)FPSYVDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25959

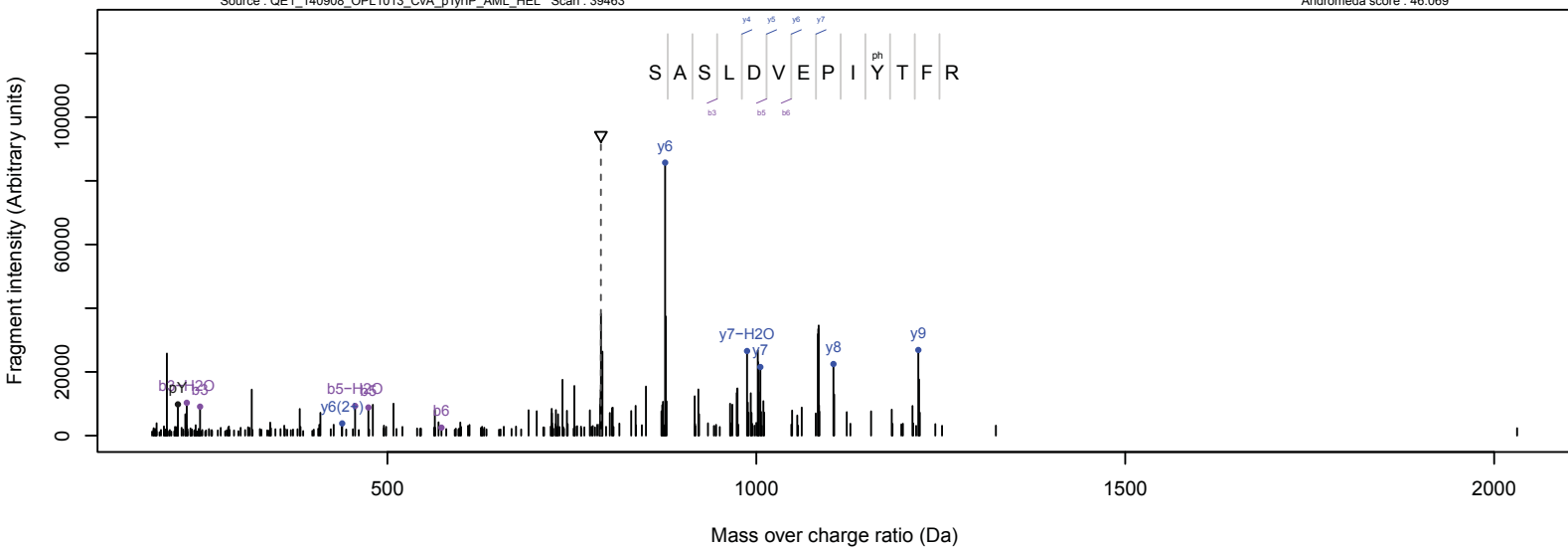
Andromeda score : 46.069



SASLDVEPIY(ph)TFR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 39463

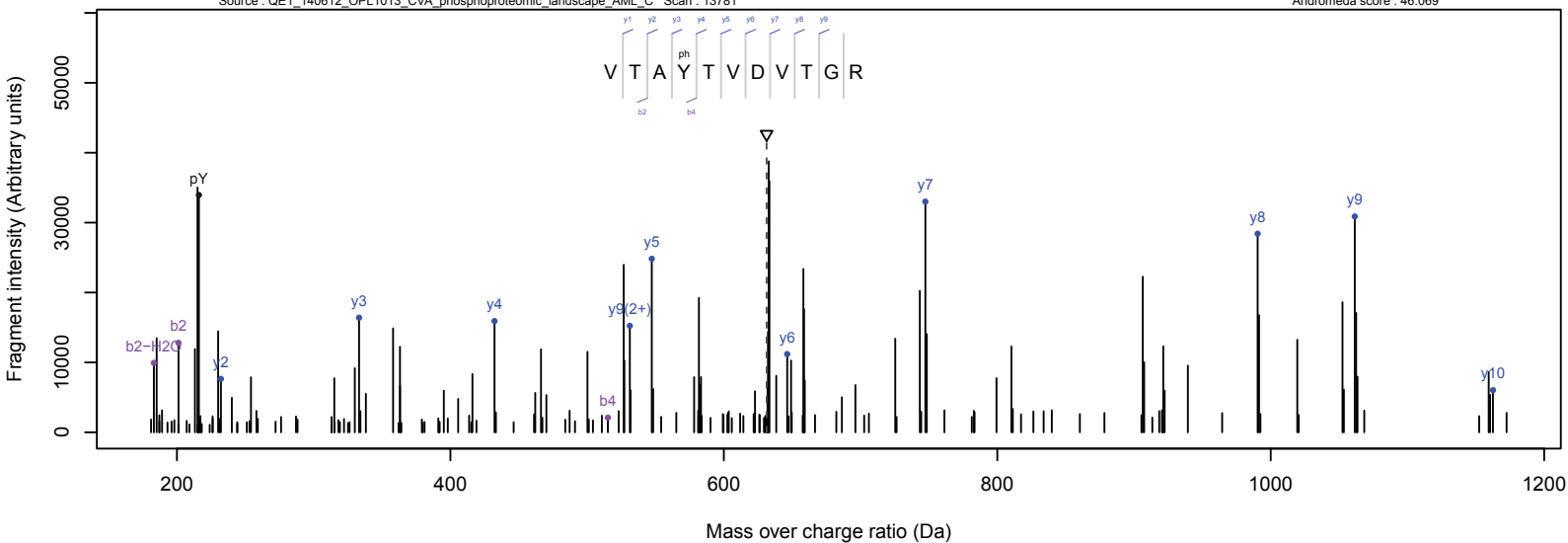
Andromeda score : 46.069



VTAY(ph)TVDTVTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13781

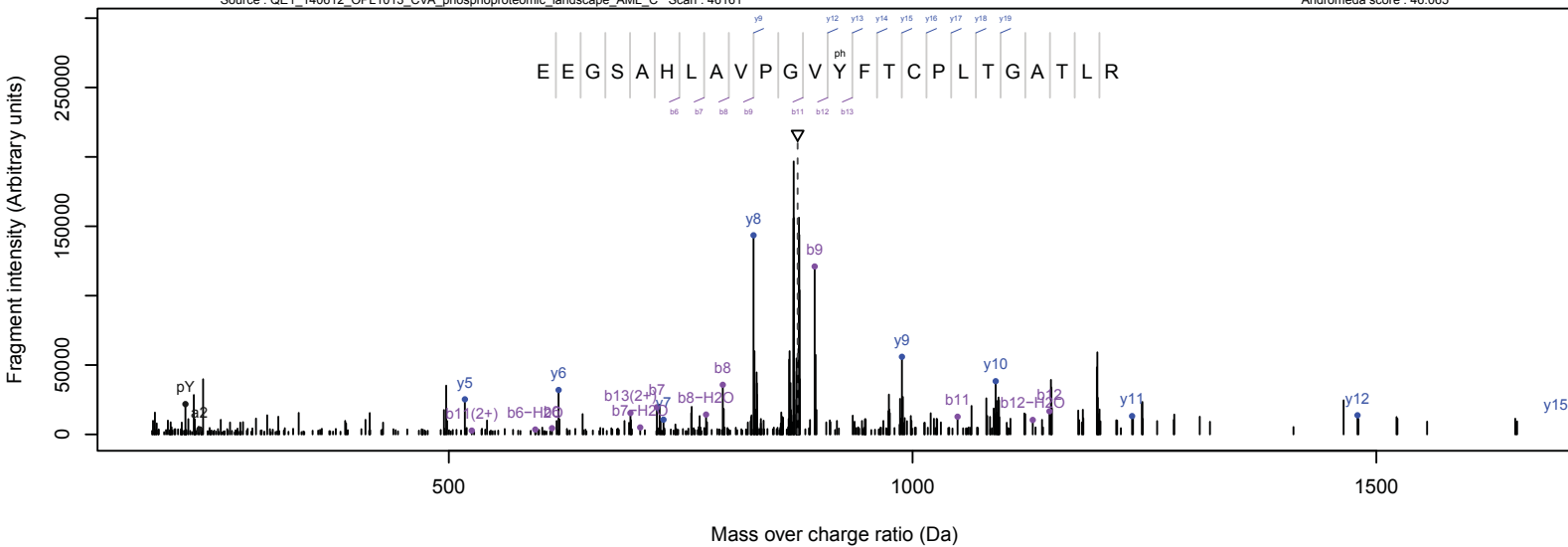
Andromeda score : 46.069



EEGSAHLAVPGVY(ph)FTCLTGATLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46161

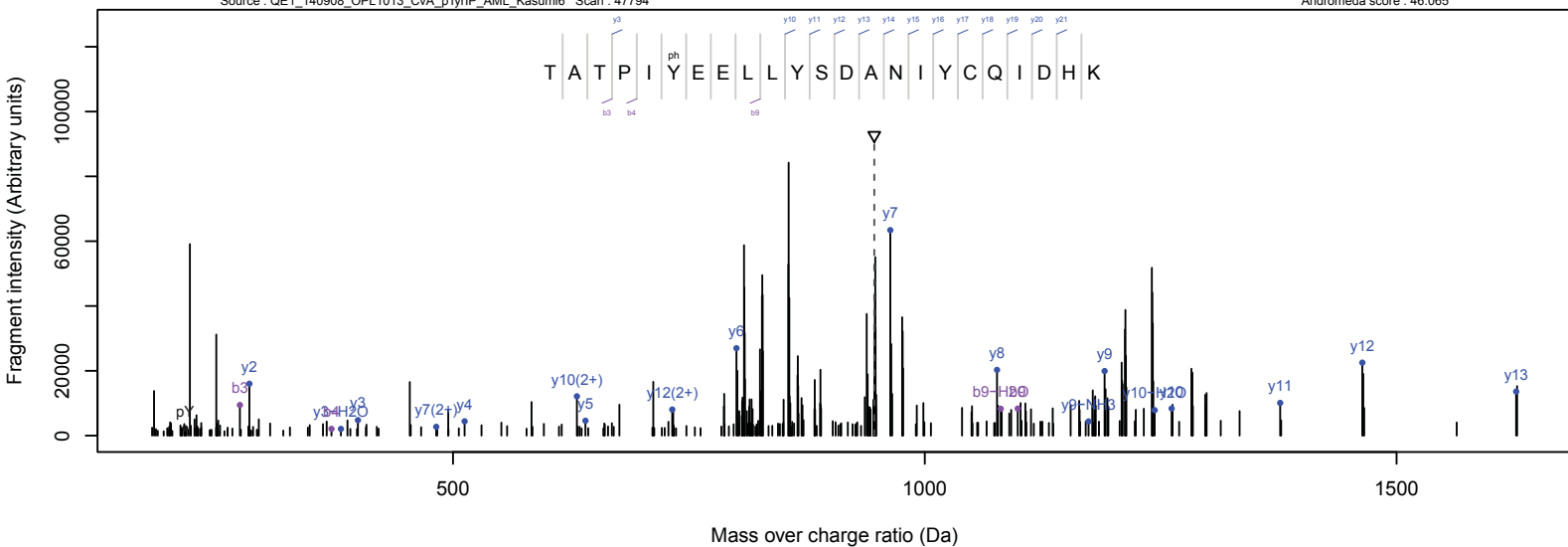
Andromeda score : 46.065



## TATPIY(ph)EELLYSDANIYQCIDHK-3+

Source: QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan: 47794

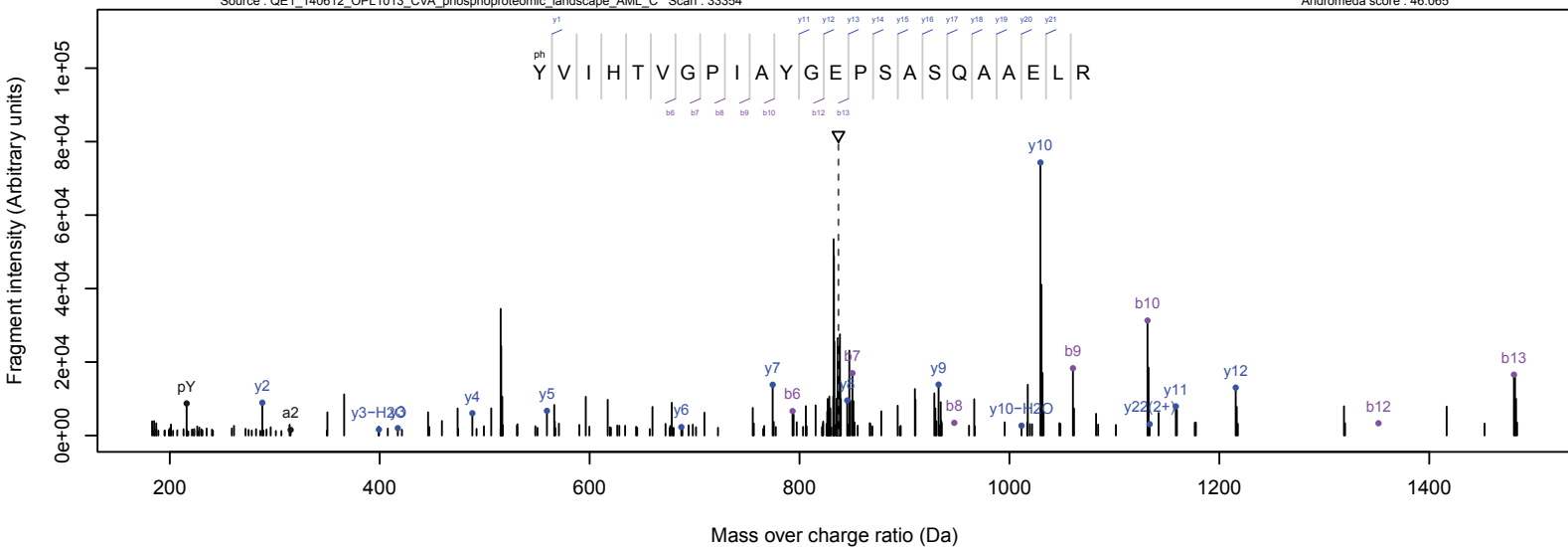
Andromeda score: 46.065



## Y(ph)VIHTVGPVIAAYGEPSSAQAAELR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 33354

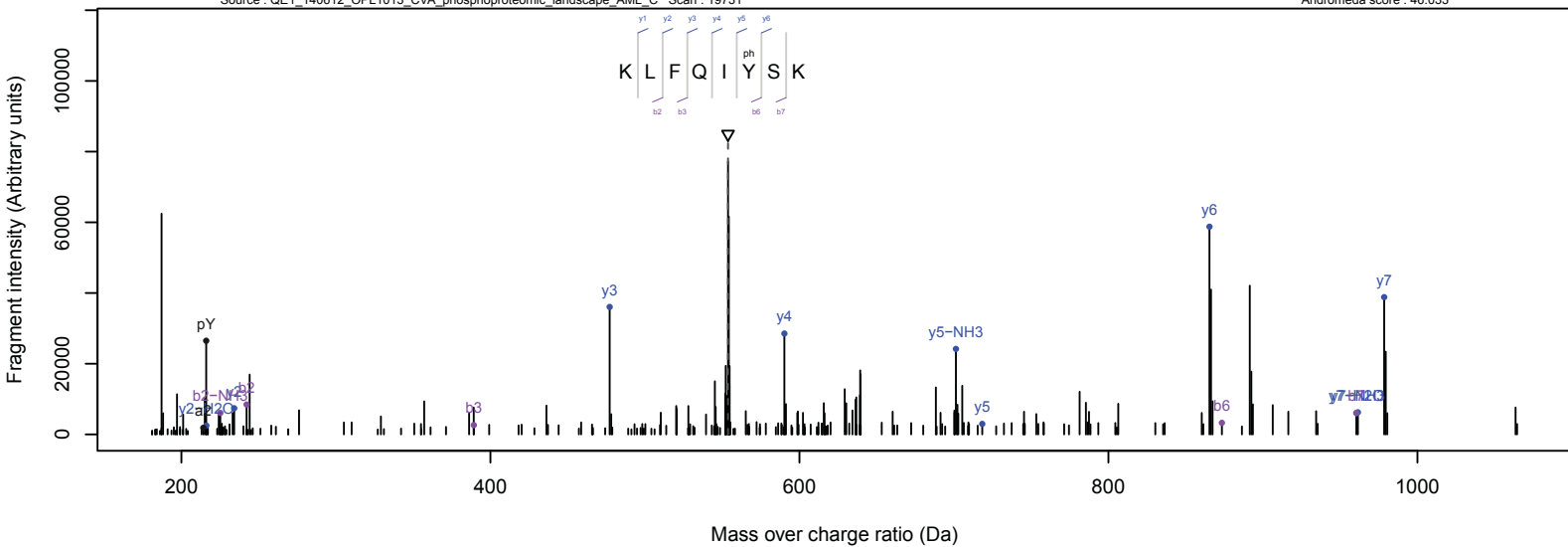
Andromeda score: 46.065



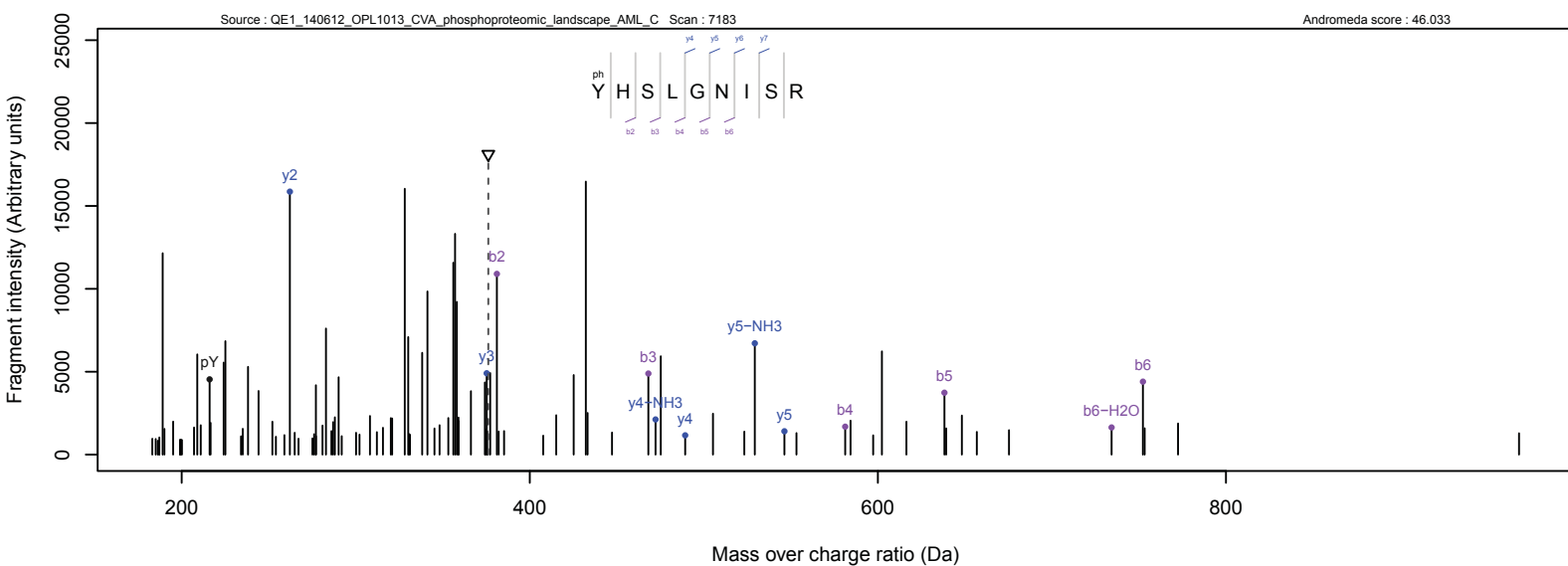
## KLFQIY(ph)SK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 19731

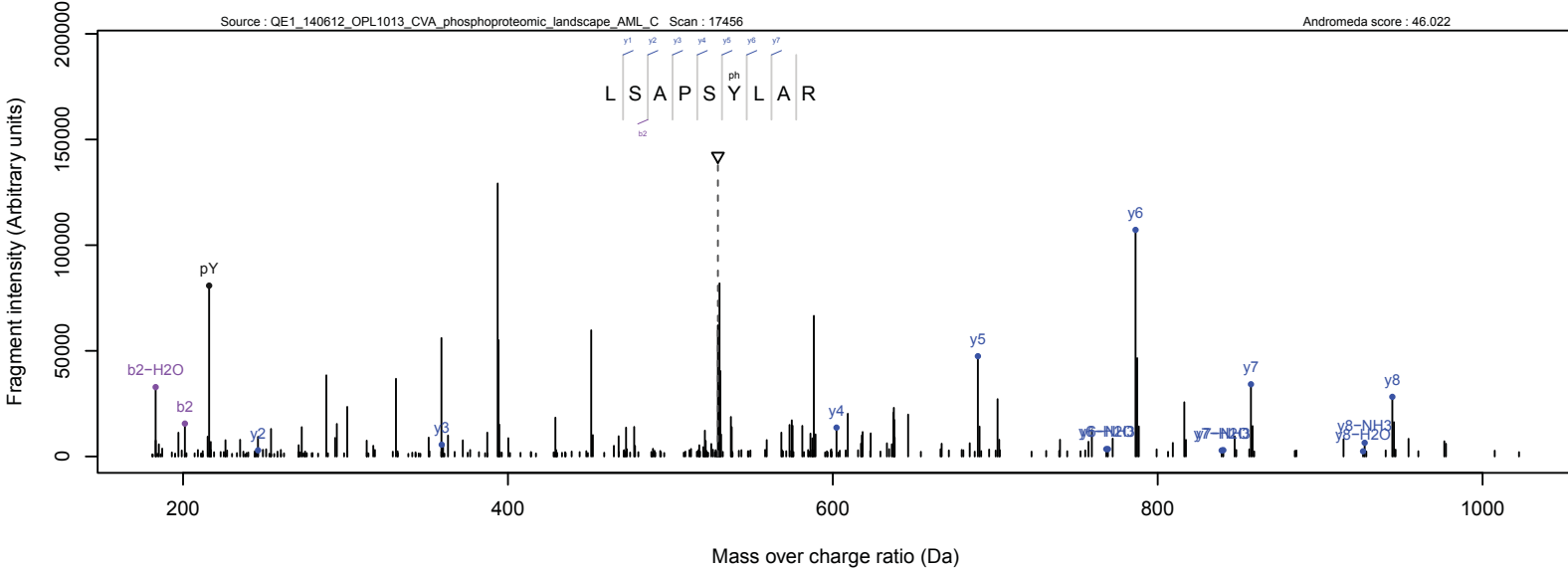
Andromeda score: 46.033



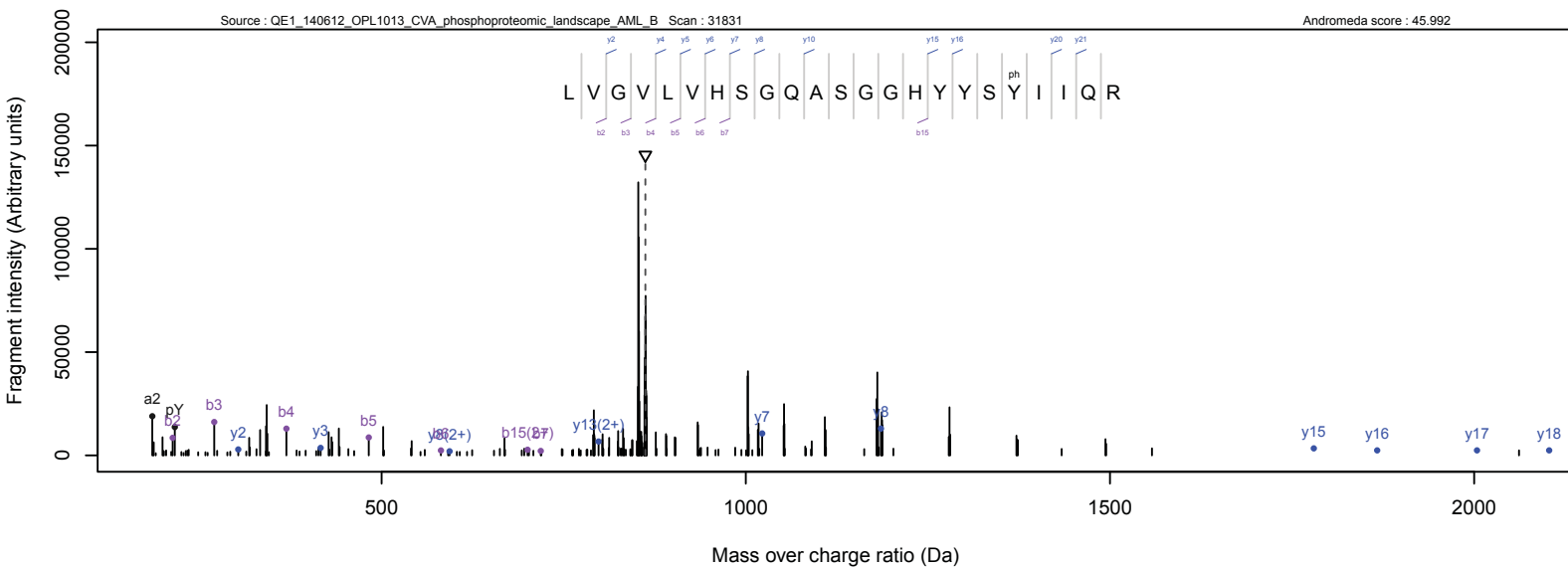
Y(ph)HSLGNISR-3+



LSAPSY(ph)LAR-2+



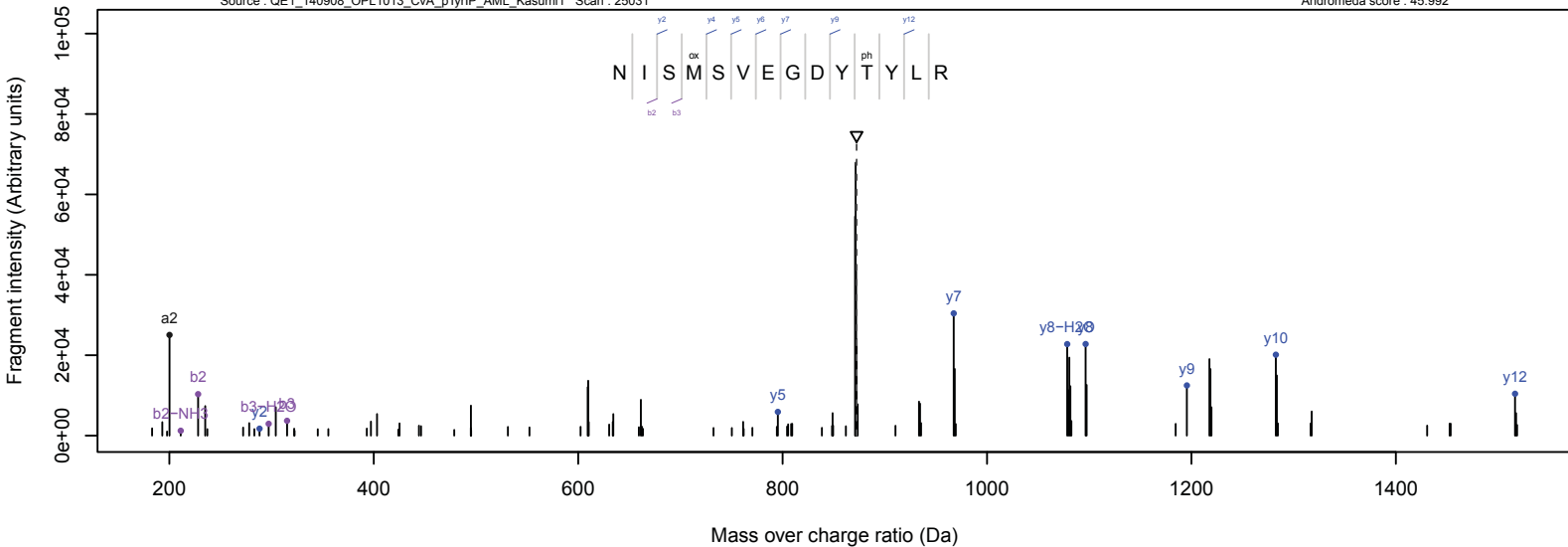
LVGLVHSGQASGGHYYS(ph)IIQR-3+



NISM(ox)SVEGDYT(ph)YLR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 25031

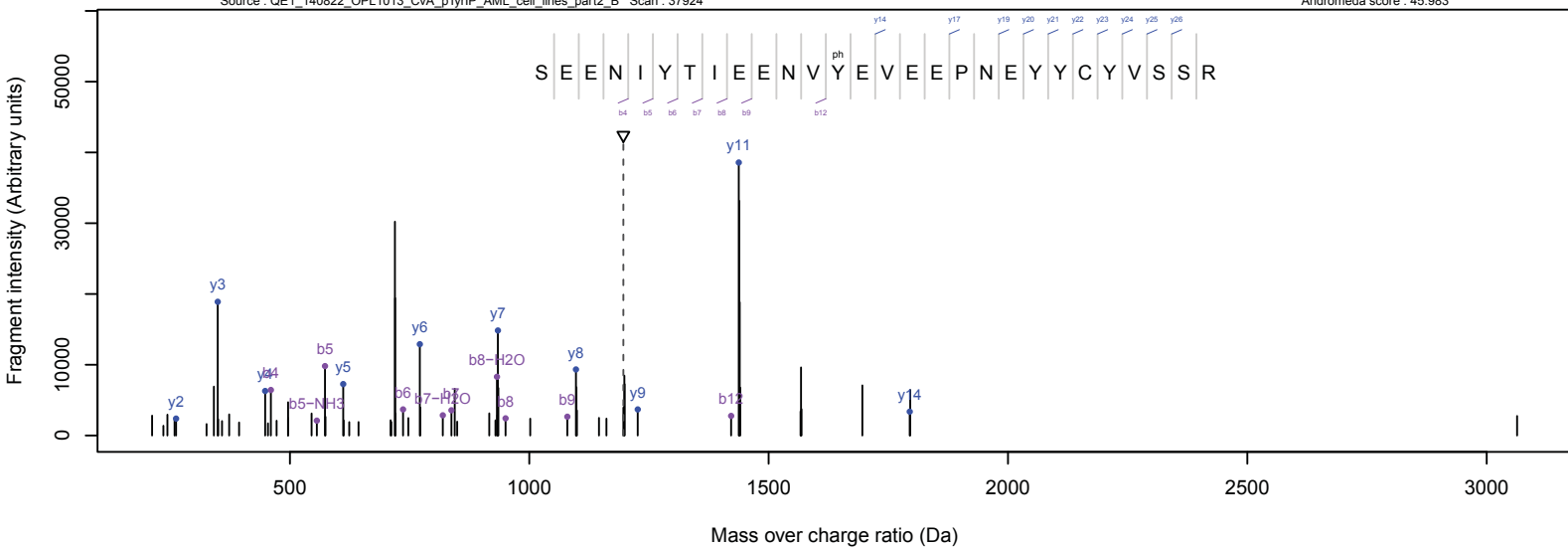
Andromeda score : 45.992



SEENIYTIENVY(ph)EVEEPNEYCYVSSR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 37924

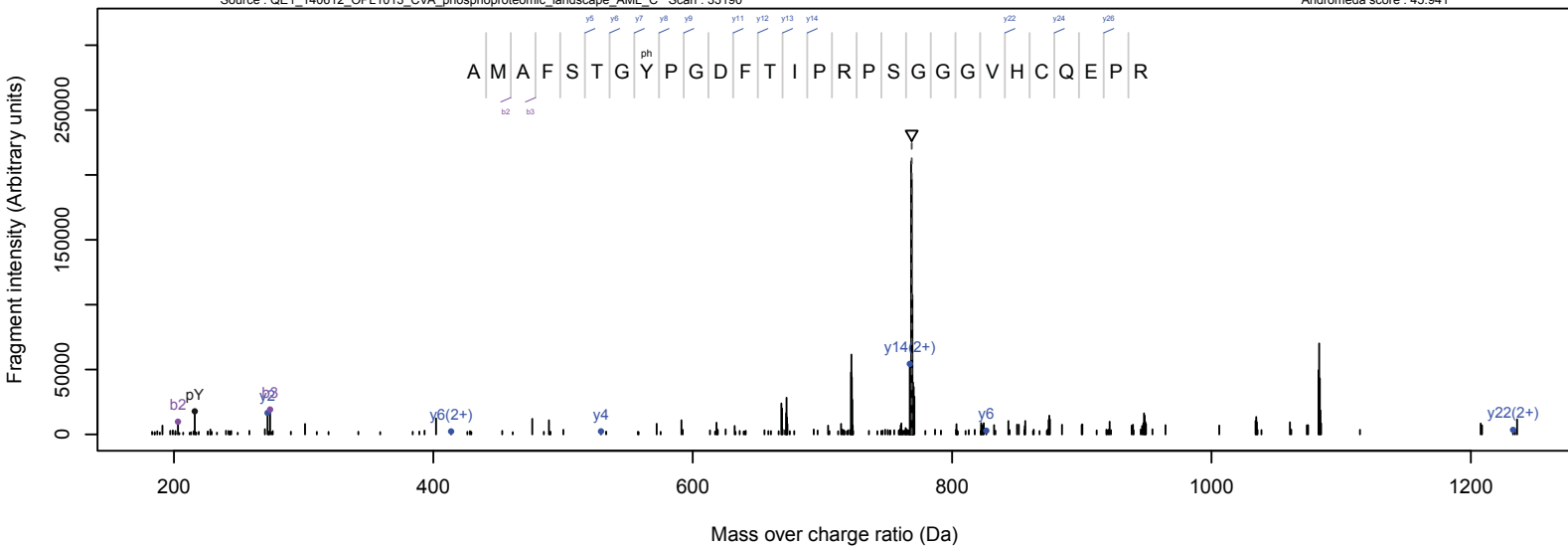
Andromeda score : 45.983



AMAFSTGY(ph)PGDFTIPRPSGGGVHQCQEP-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33190

Andromeda score : 45.941

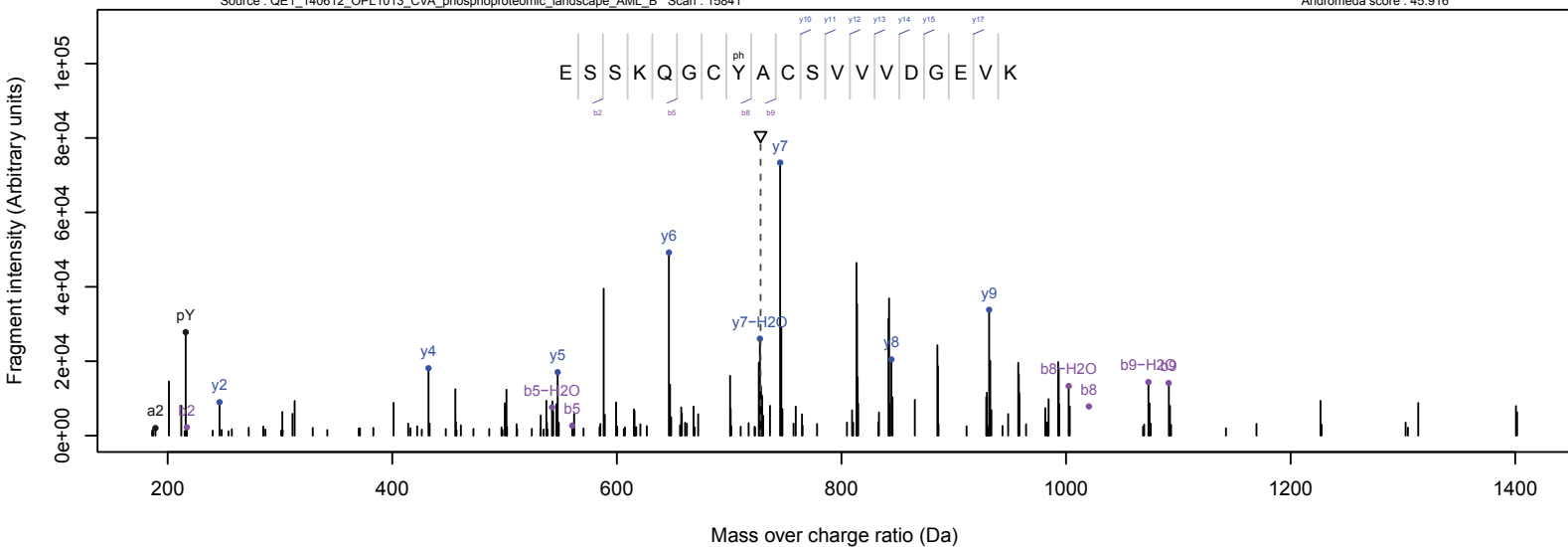




ESSKQGCY(ph)ACSVVDGEVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15841

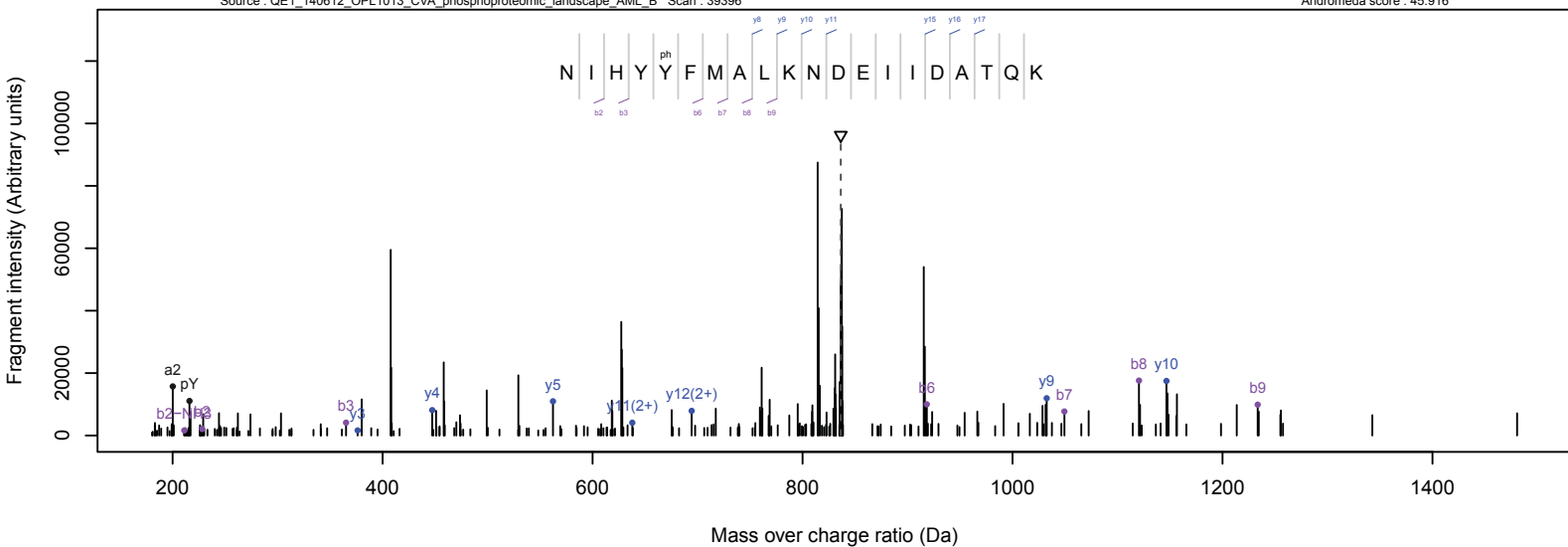
Andromeda score : 45.916



NIHYYP(ph)FMALKNDEIIDATQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 39396

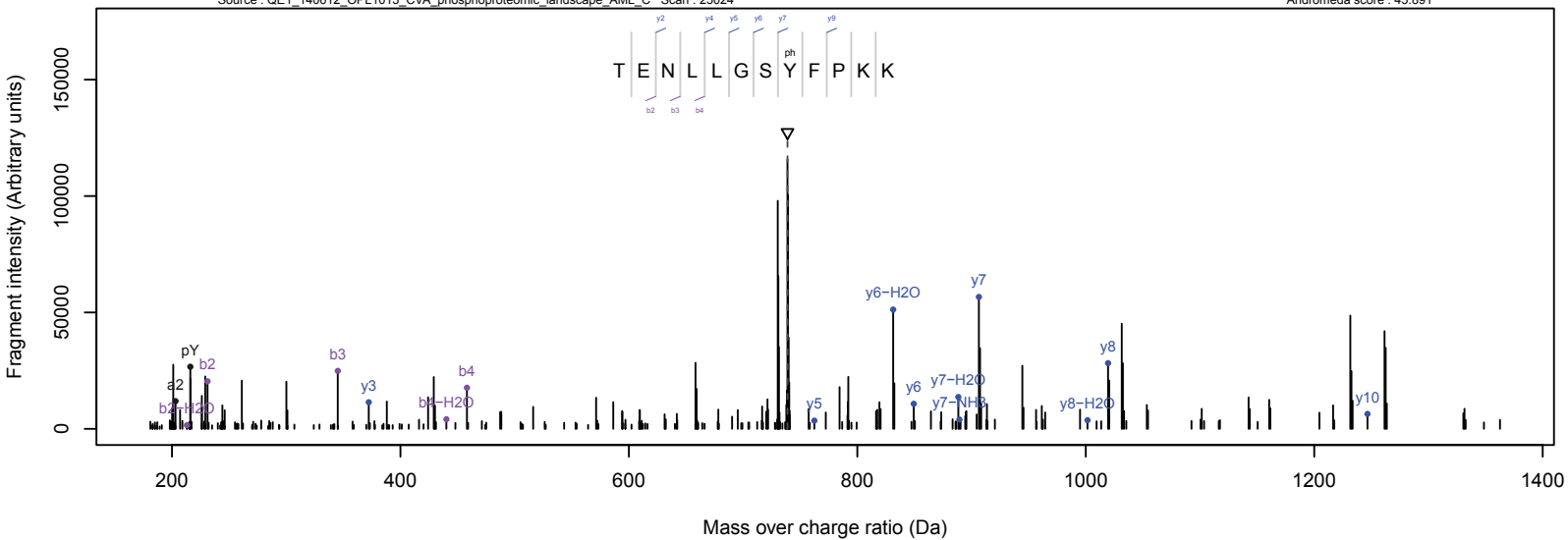
Andromeda score : 45.916



TENLLGSYP(ph)FPKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 25024

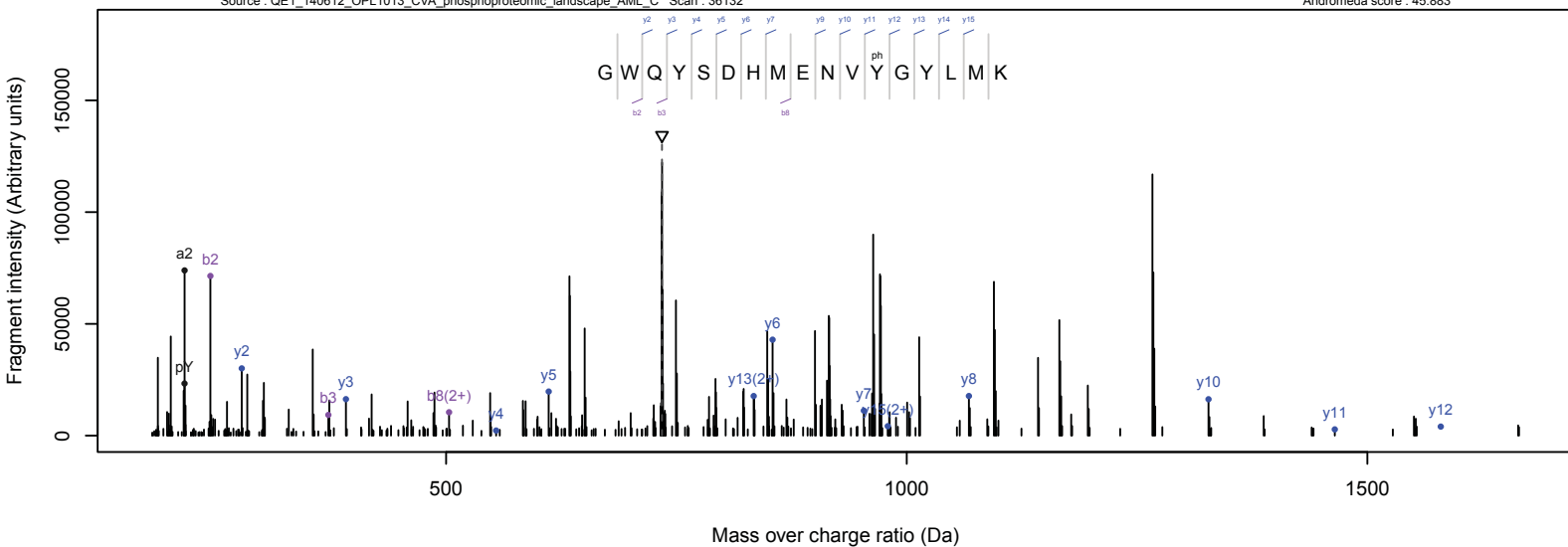
Andromeda score : 45.891



## GWQYSDHMENVY(ph)GYLMK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36132

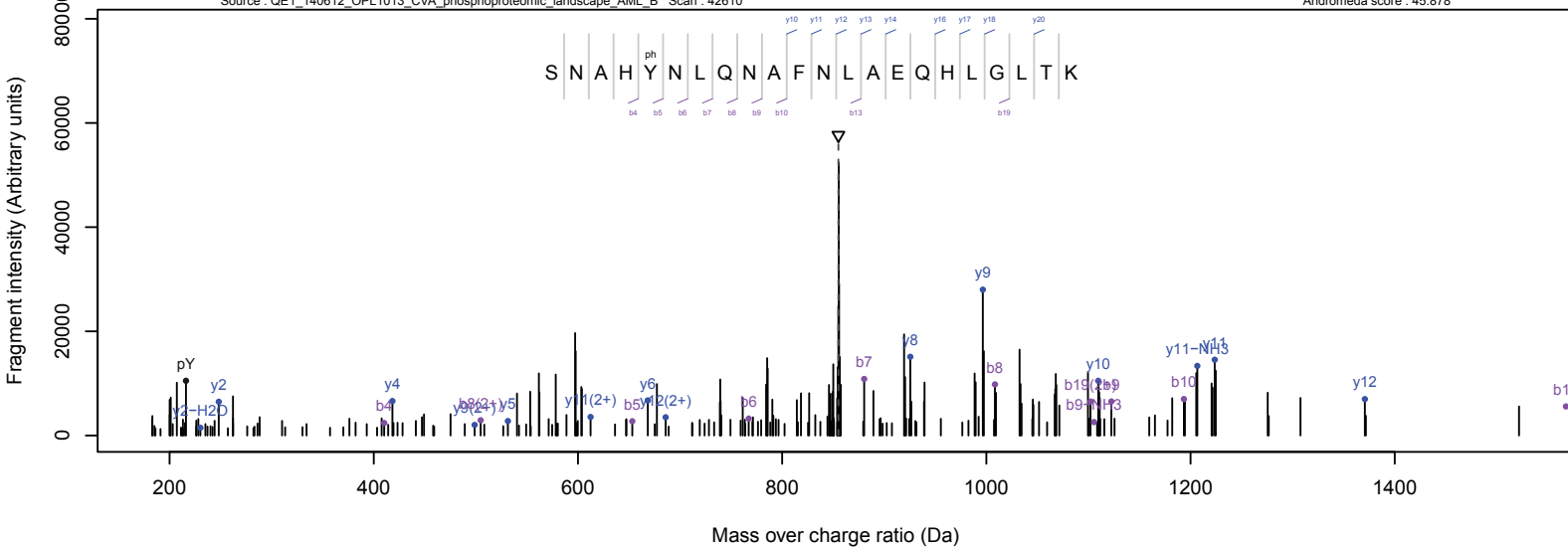
Andromeda score : 45.883



## SNAHY(ph)NLQNAFLAEQHLGLTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 42610

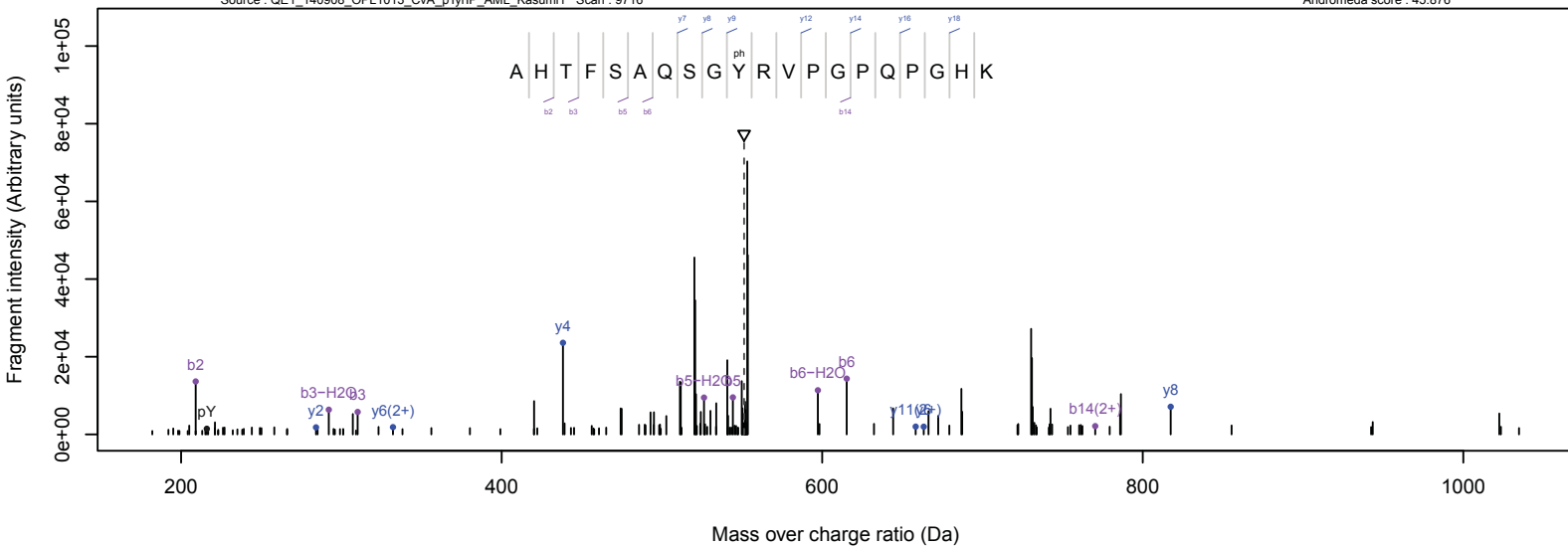
Andromeda score : 45.878



## AHTFSAQSGY(ph)RVPGPQGHK-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 9716

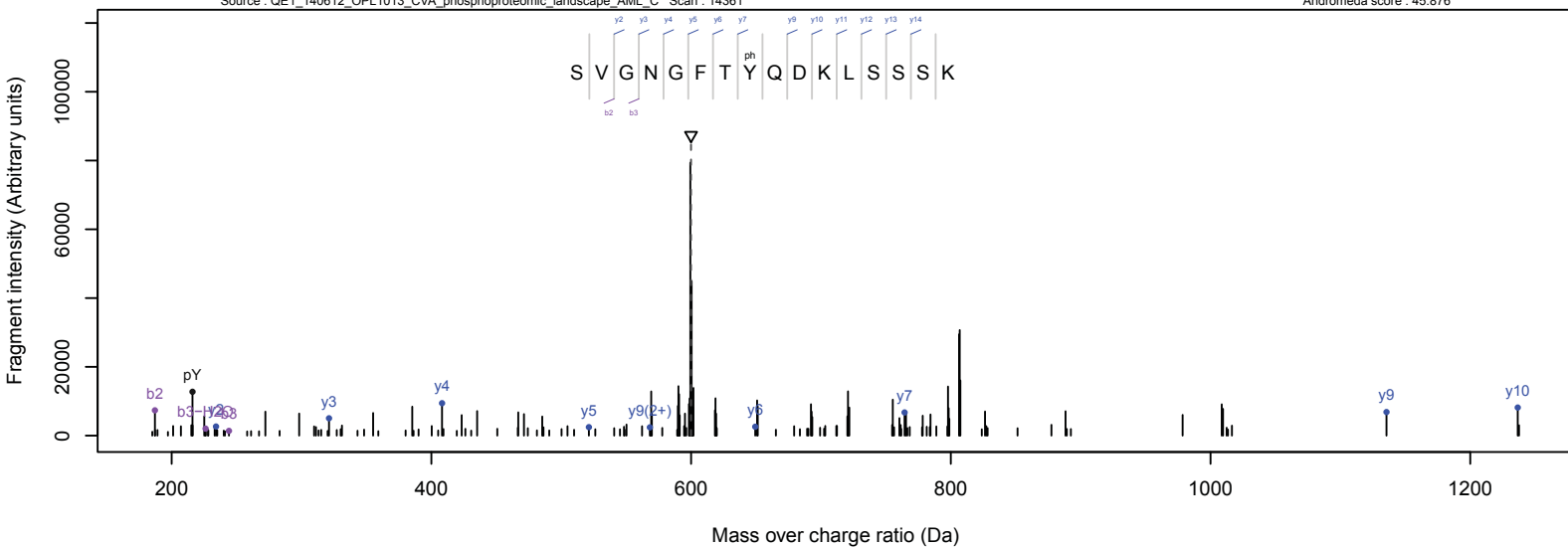
Andromeda score : 45.876



## SVNGFTY(ph)QDKLSSSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14361

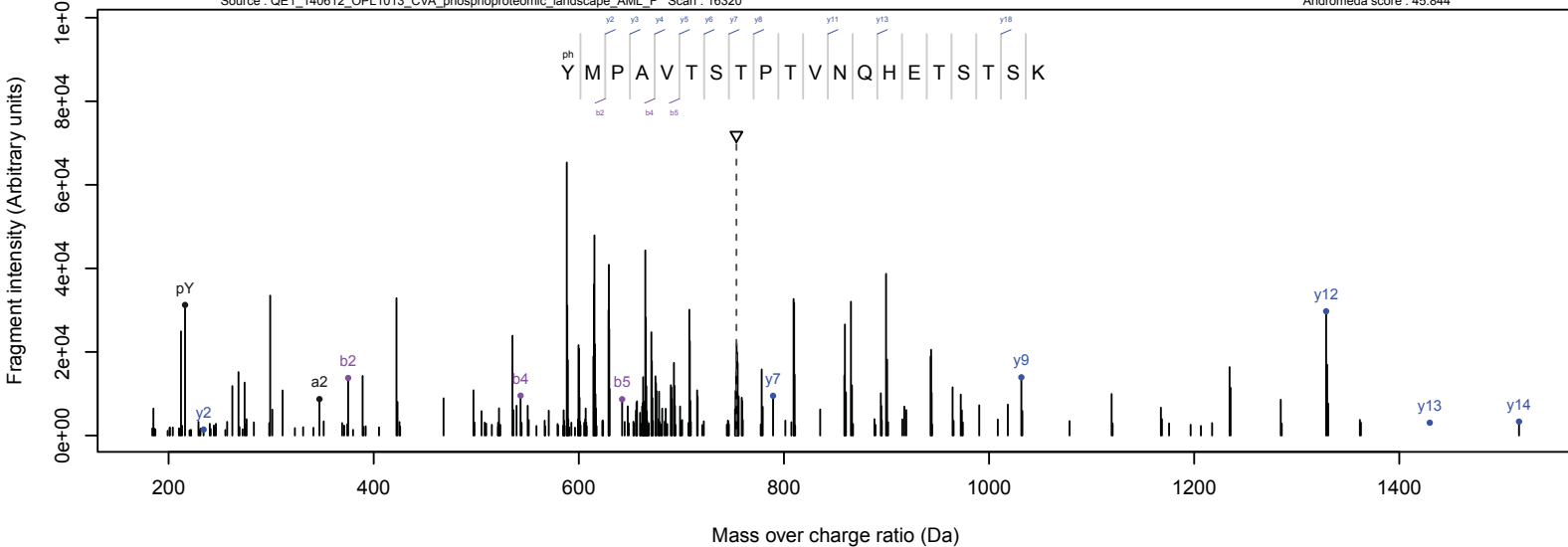
Andromeda score : 45.876



## Y(ph)MPAVTSTPTVNQHETSTSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 16320

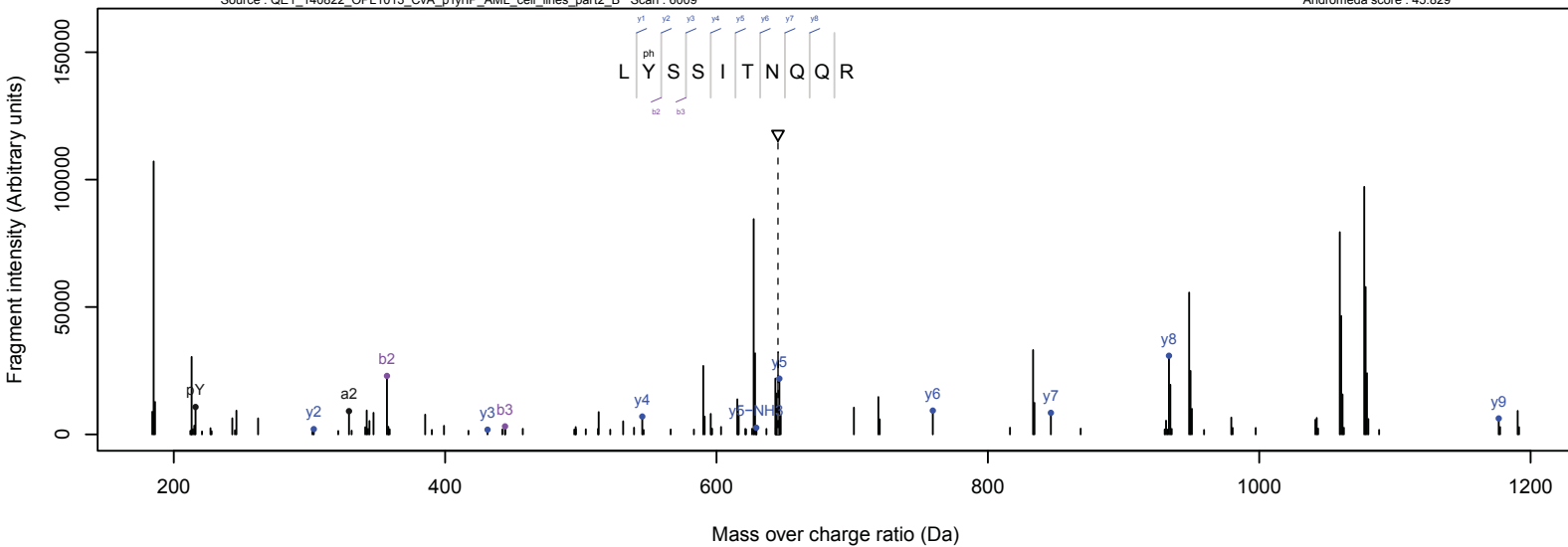
Andromeda score : 45.844

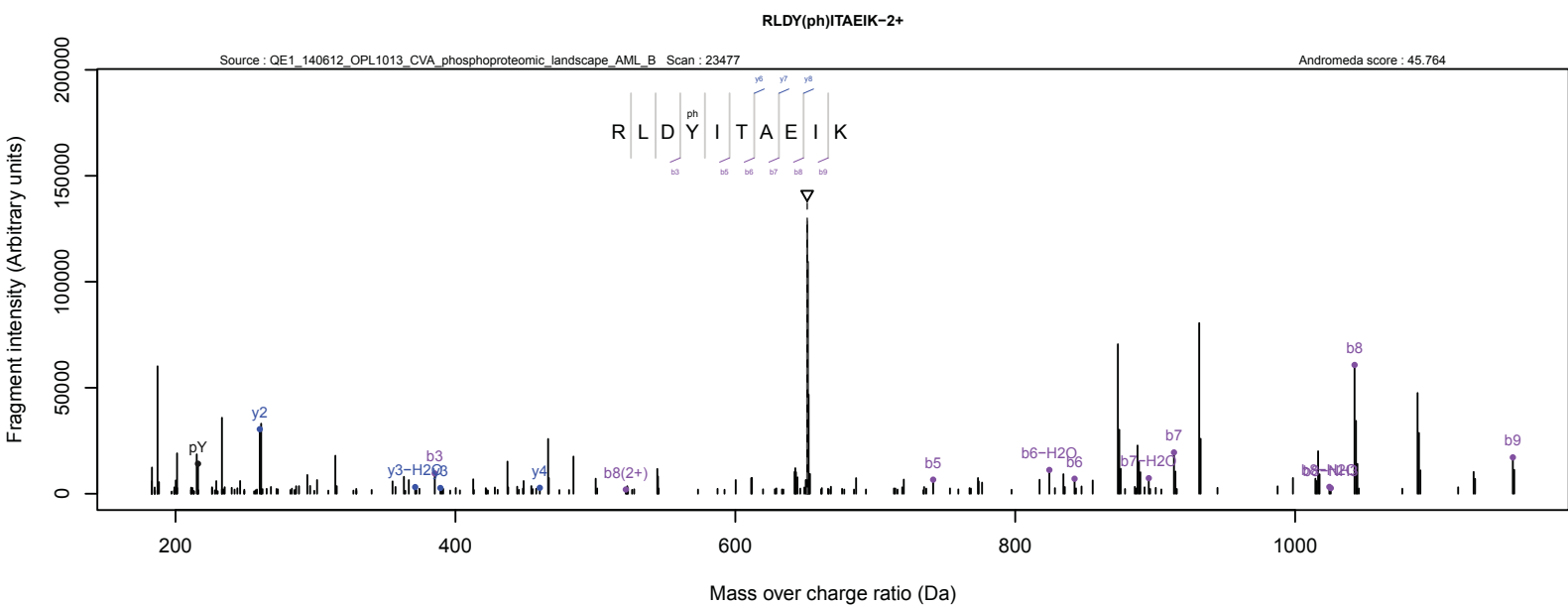
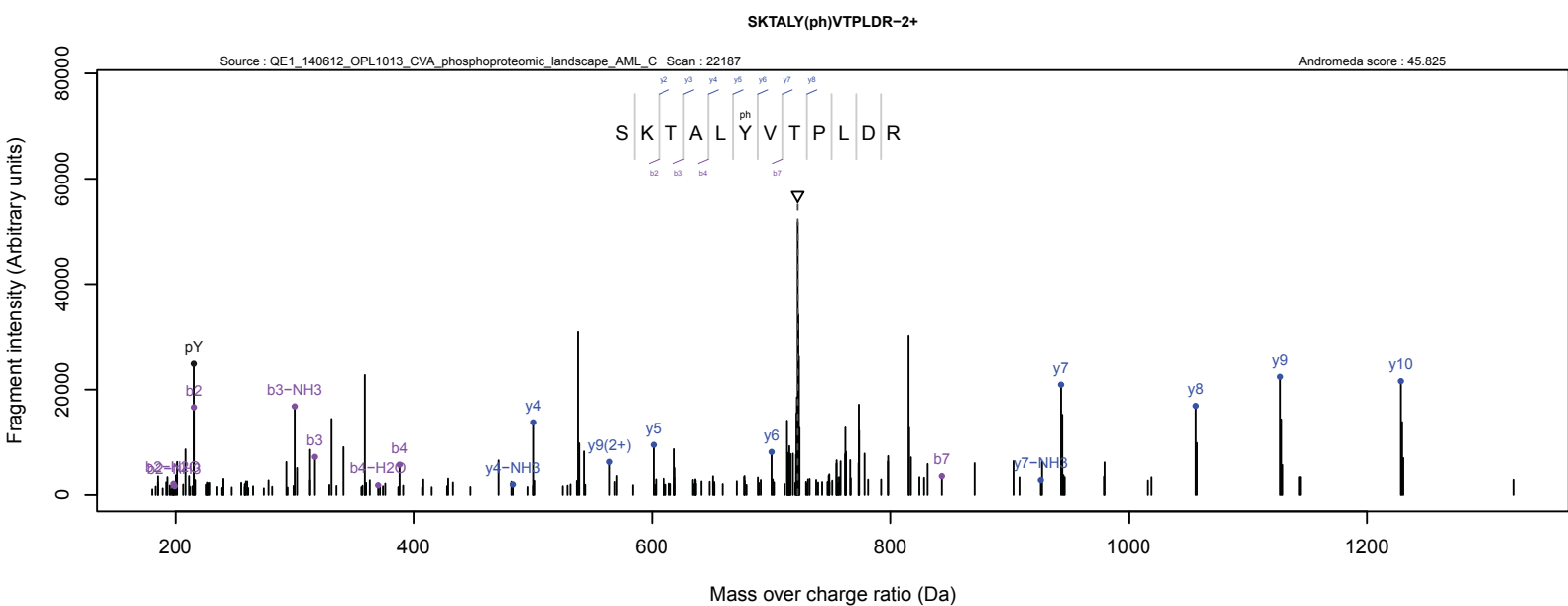
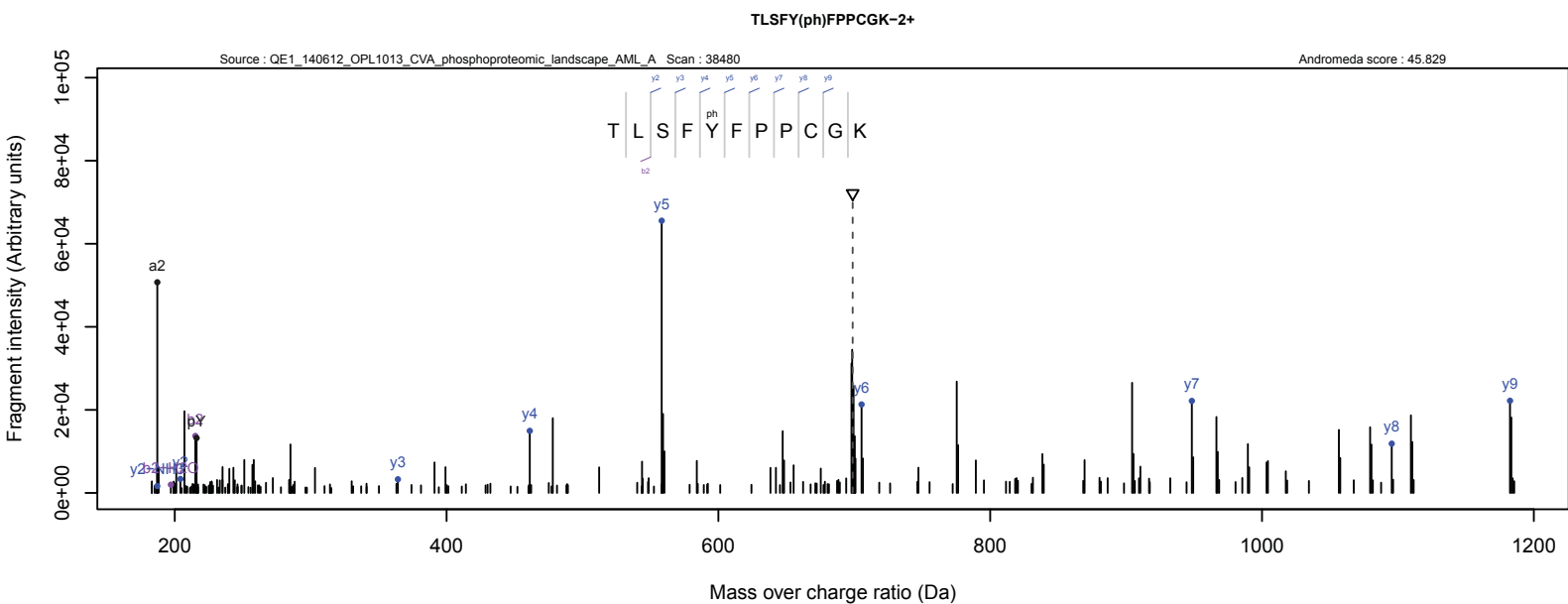


## LY(ph)SSITNQR-2+

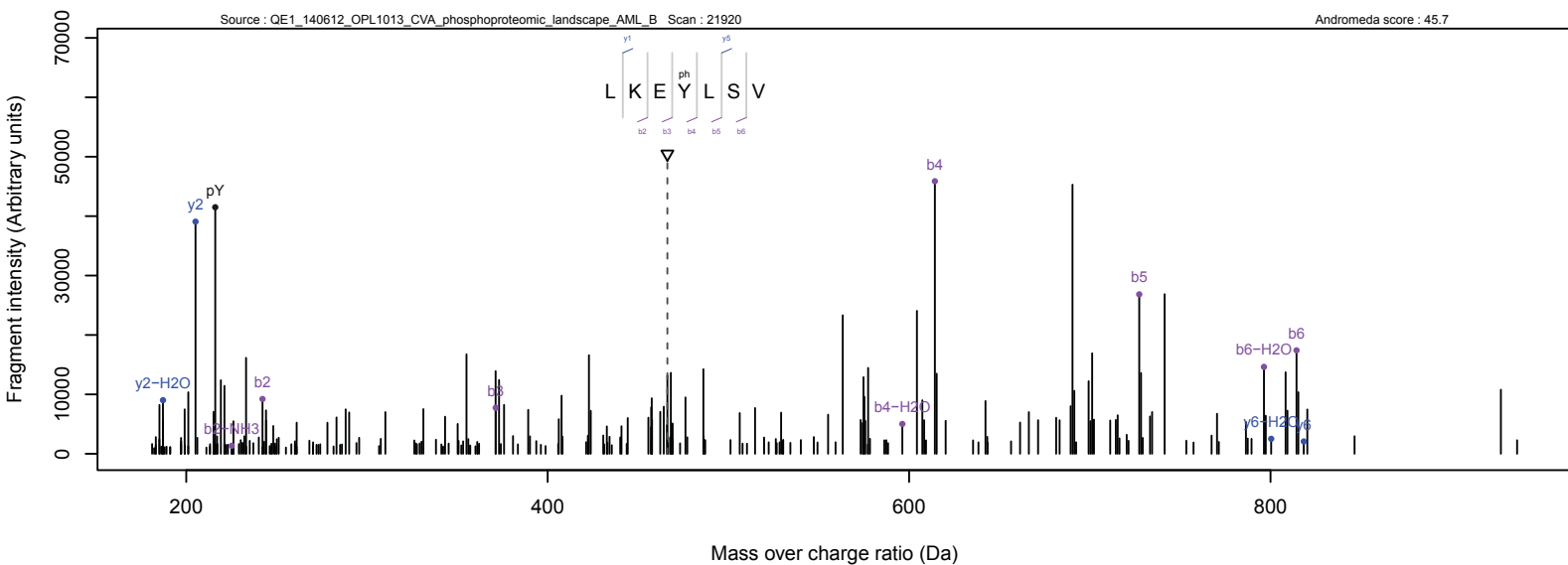
Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 6009

Andromeda score : 45.829

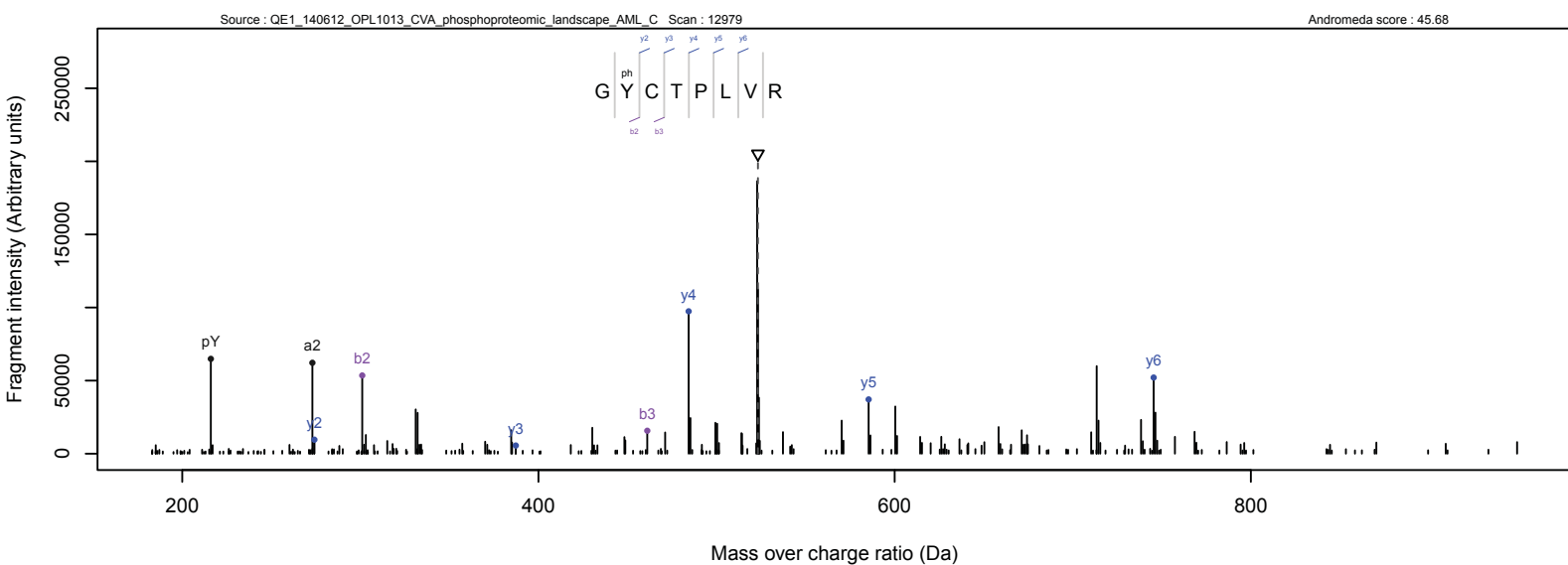




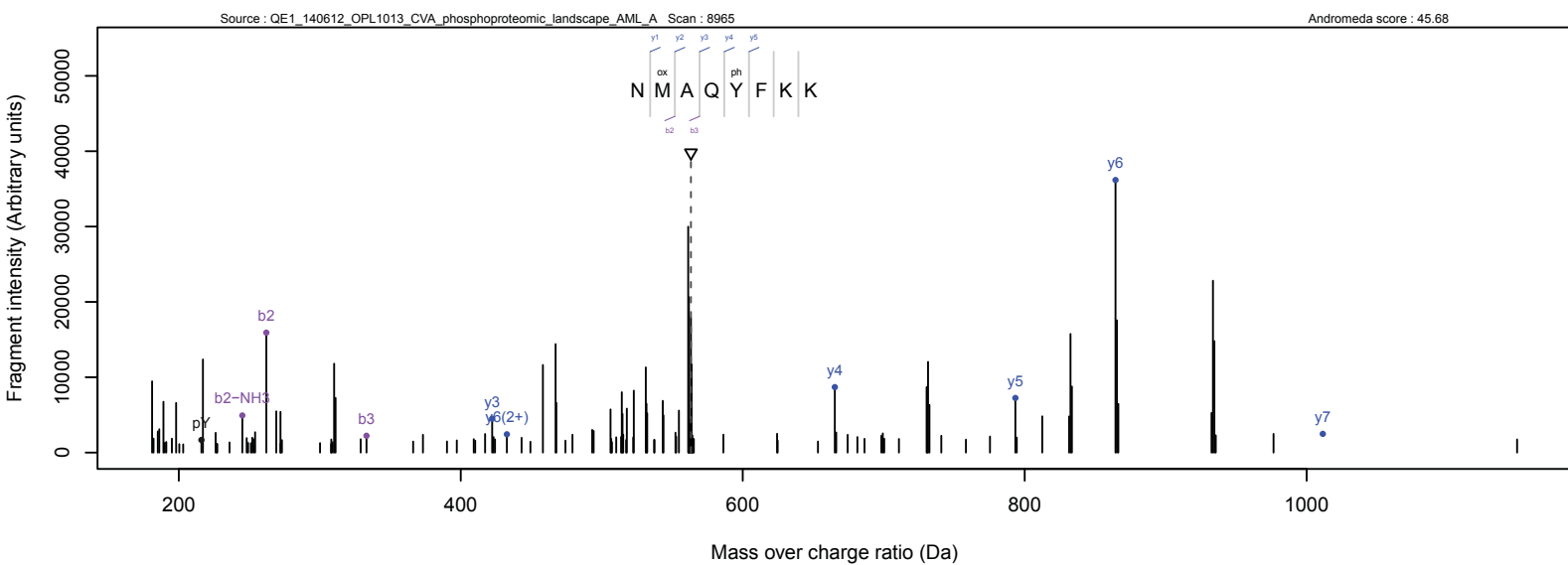
## LKEY(ph)LSV-2+



## GY(ph)CTPLVR-2+



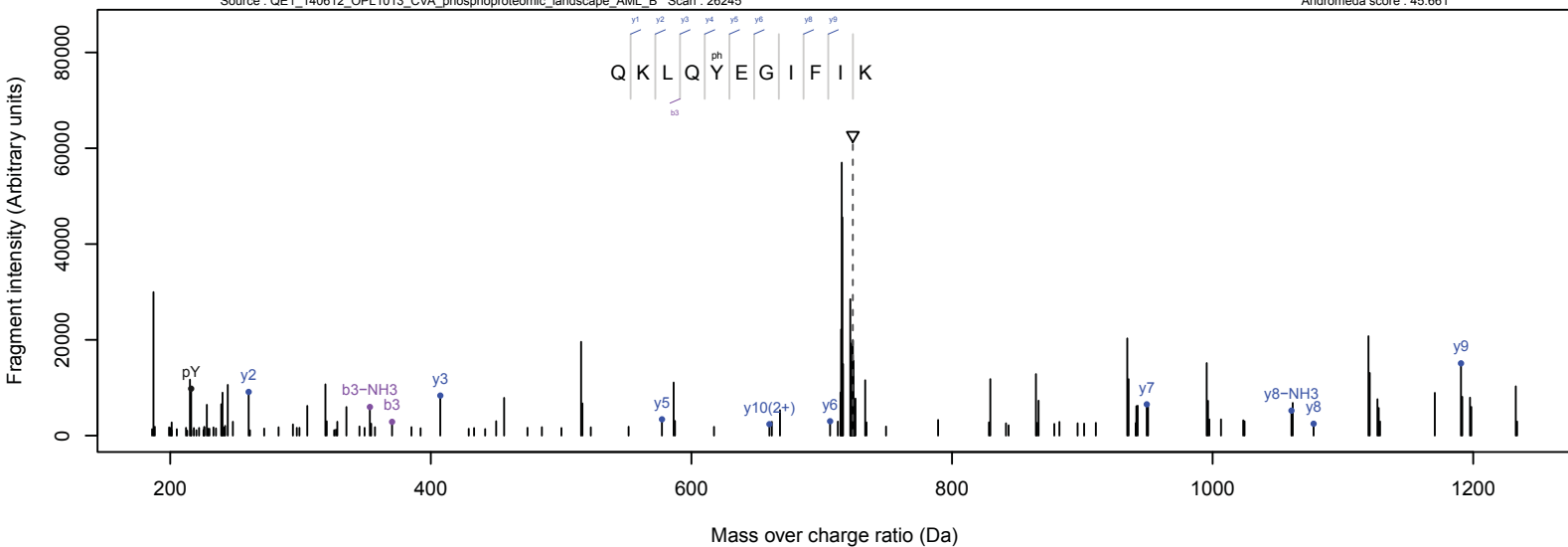
## NM(ox)AQY(ph)FKK-2+



## QKLQY(ph)EGIFIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26245

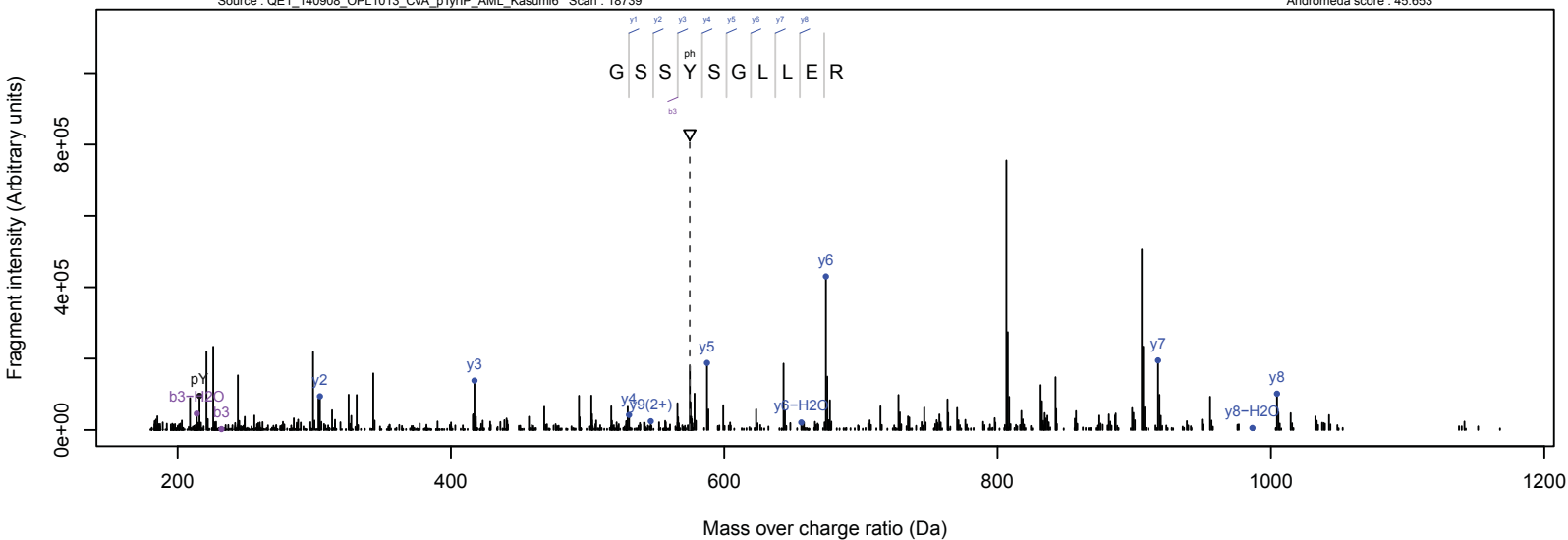
Andromeda score : 45.661



## GSSY(ph)SGLLER-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 18739

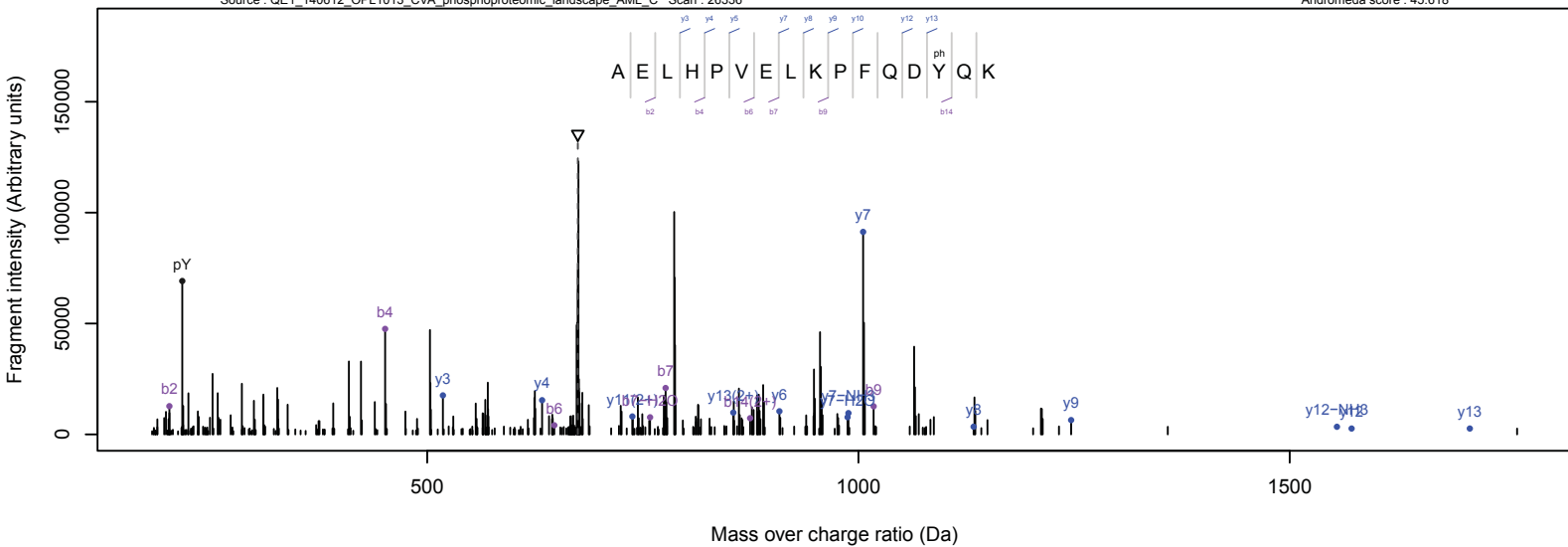
Andromeda score : 45.653



## AELHPVELKPFQDYQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26336

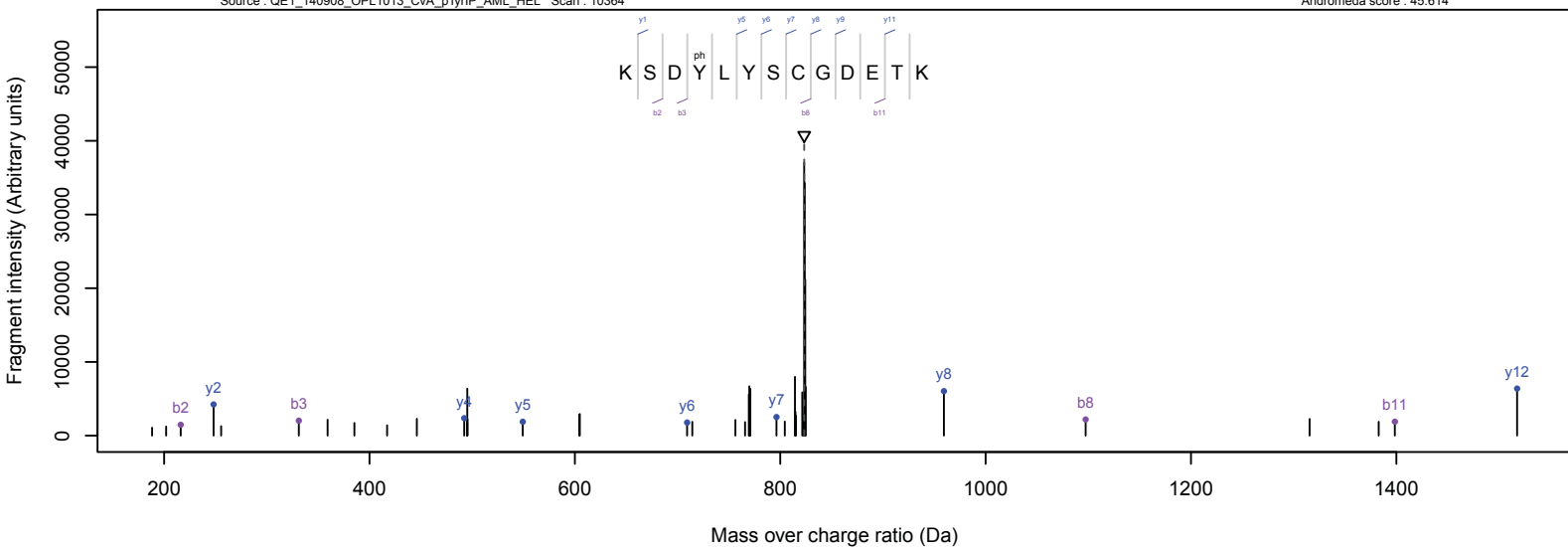
Andromeda score : 45.618



**KSDY(ph)LYSCGDETK-2+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 10364

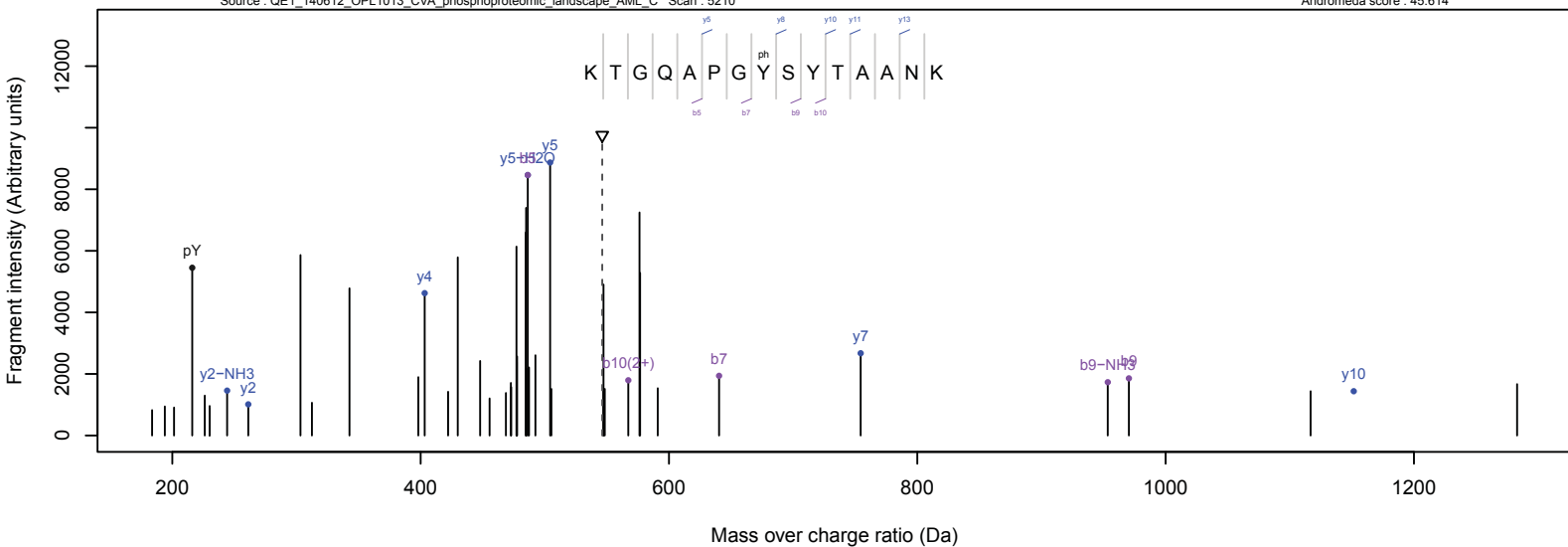
Andromeda score : 45.614



**KTGQAPG(ph)SYTAAANK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 5210

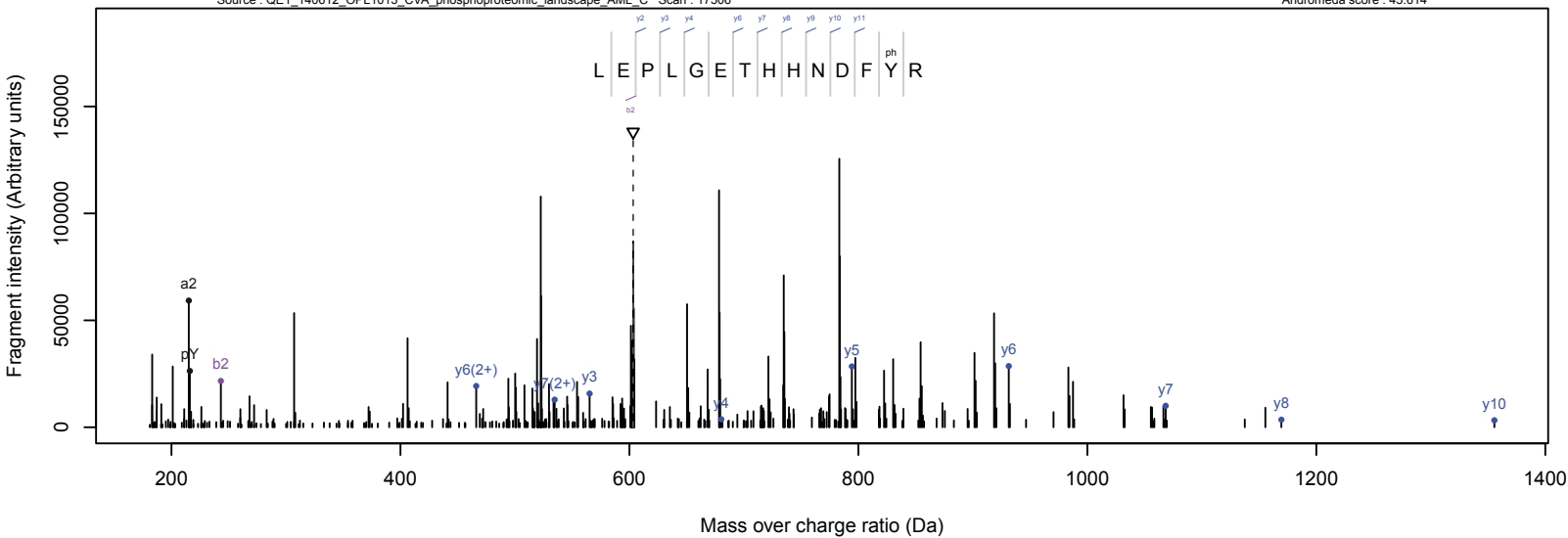
Andromeda score : 45.614



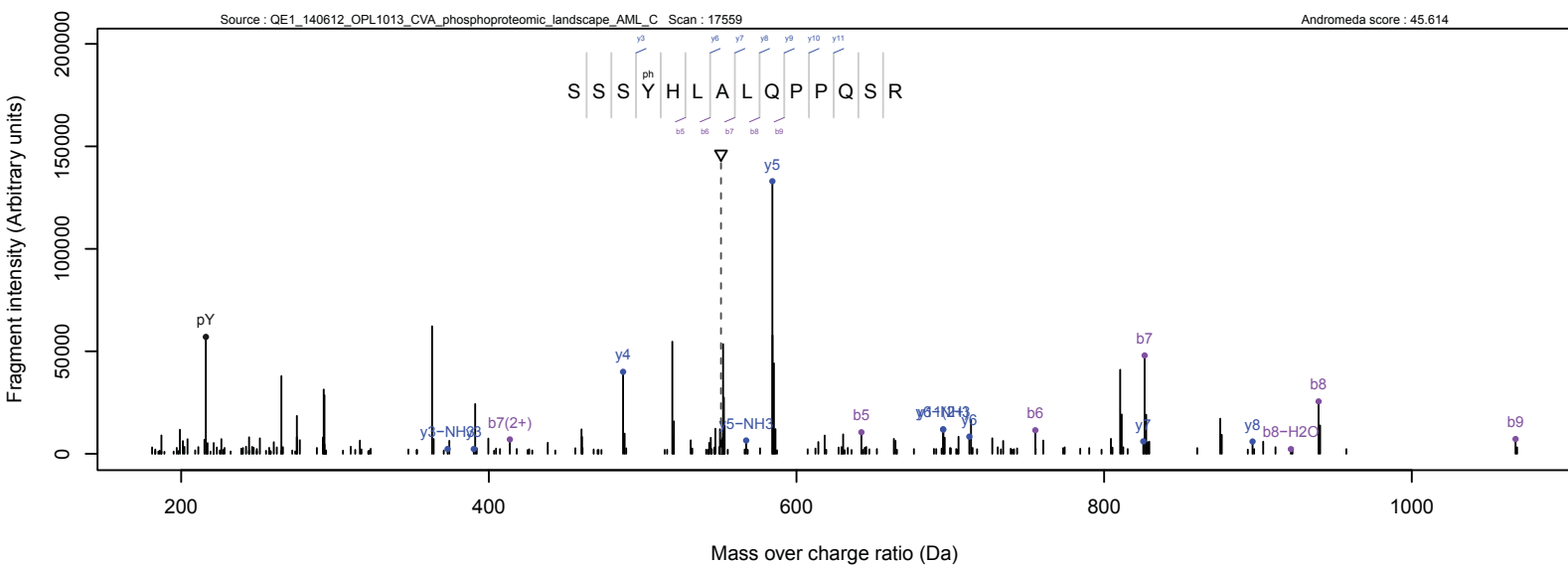
**LEPLGETHHNDFY(ph)R-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17506

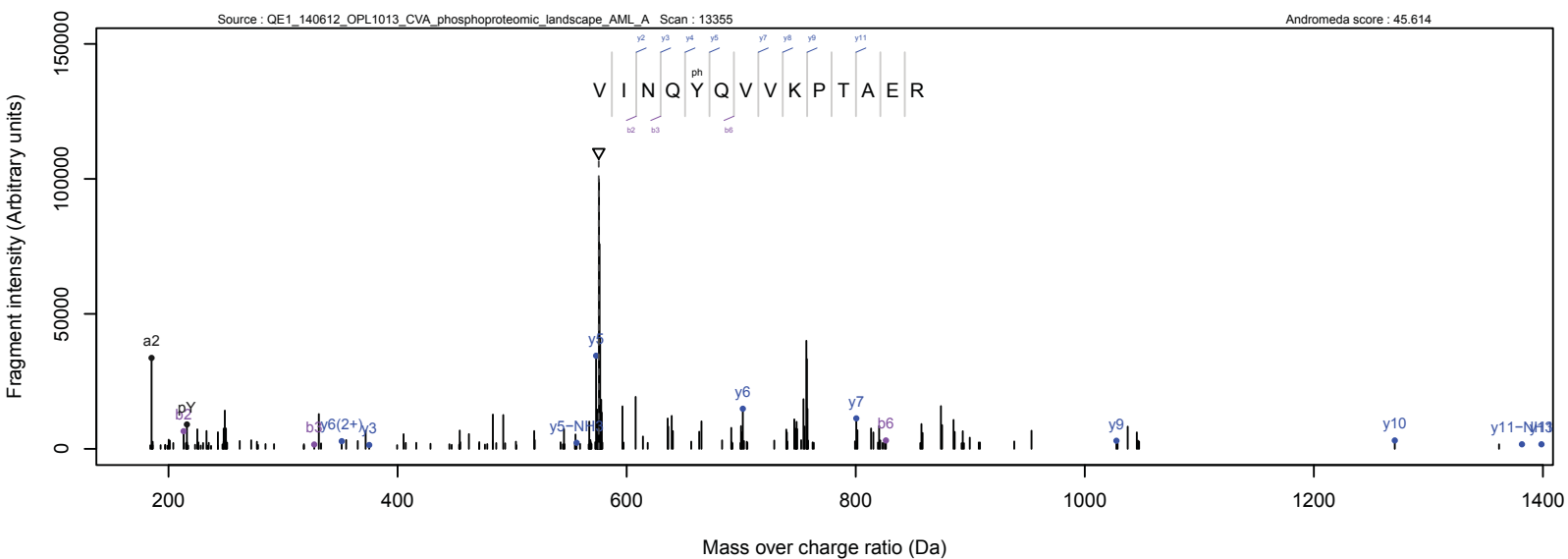
Andromeda score : 45.614



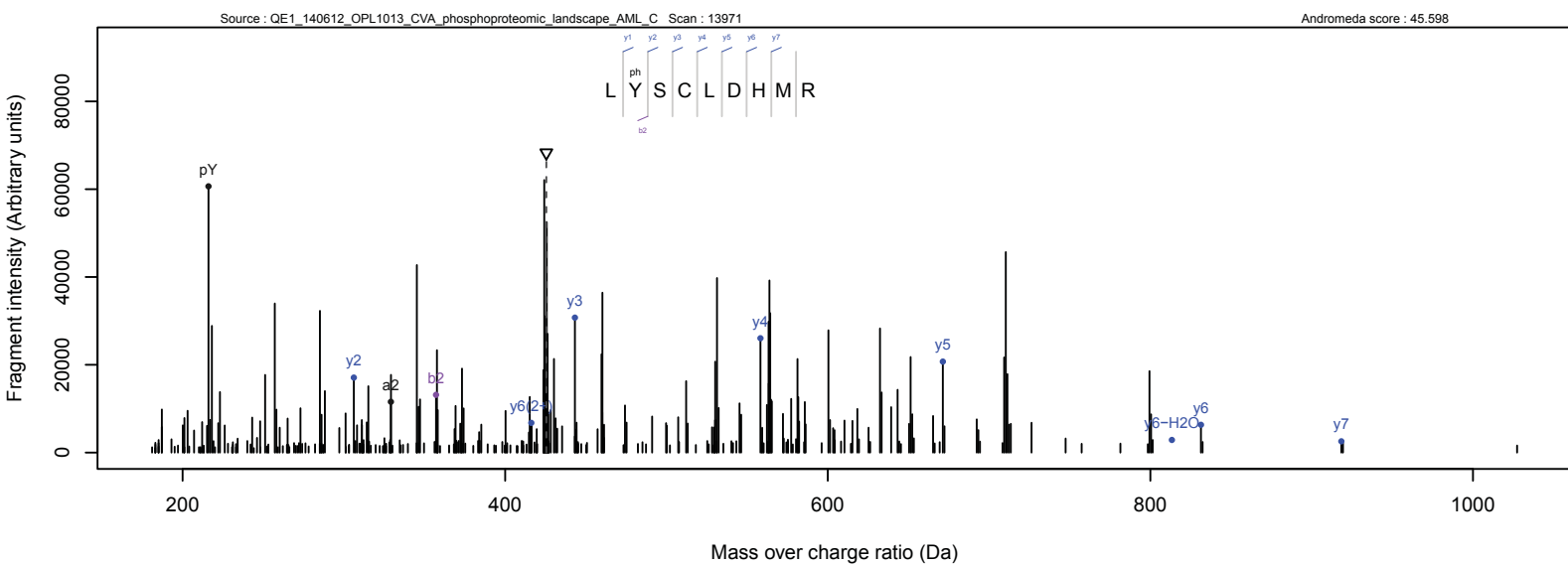
SSSY(ph)HLALQPPQR-3+



VINQY(ph)QVVKPTAER-3+

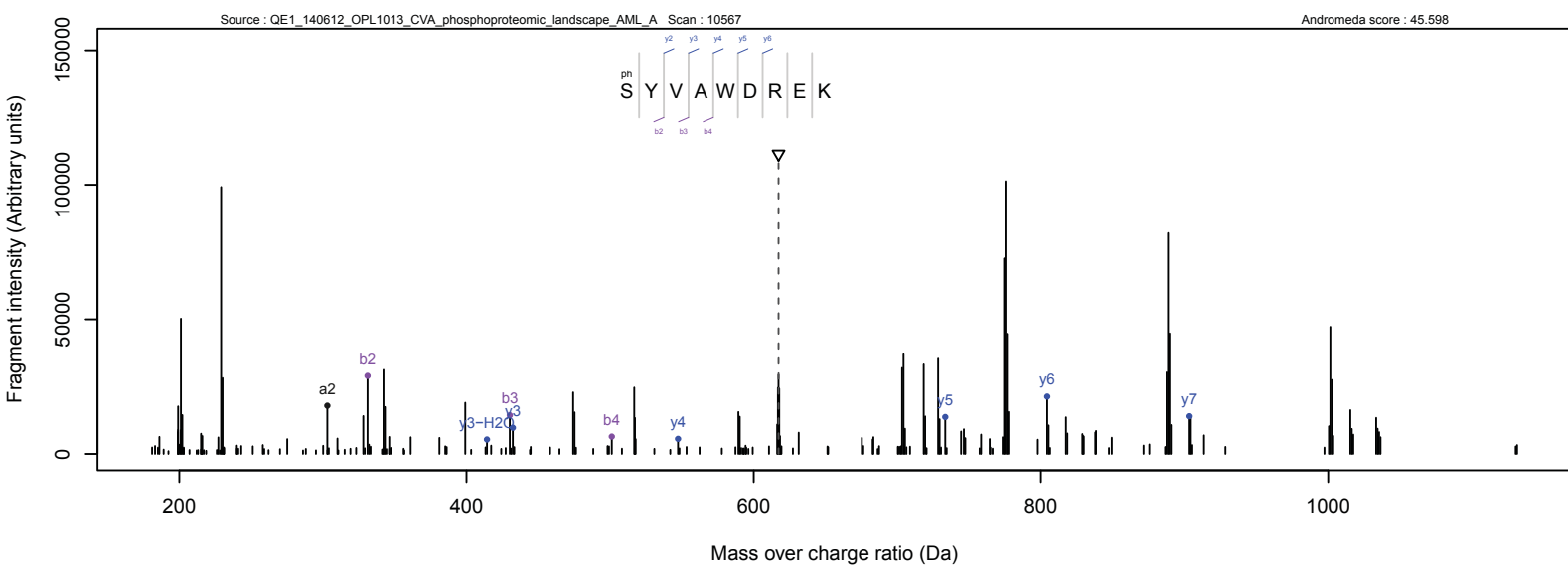


LY(ph)SCLDHR-3+

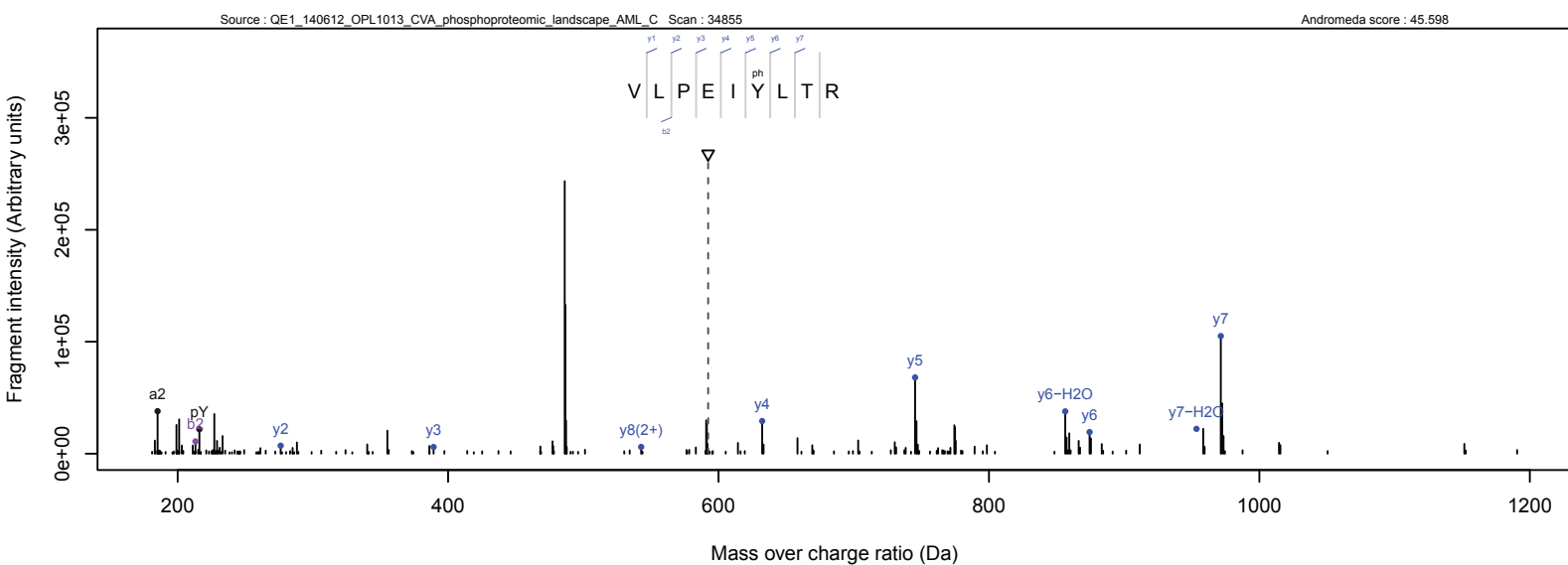




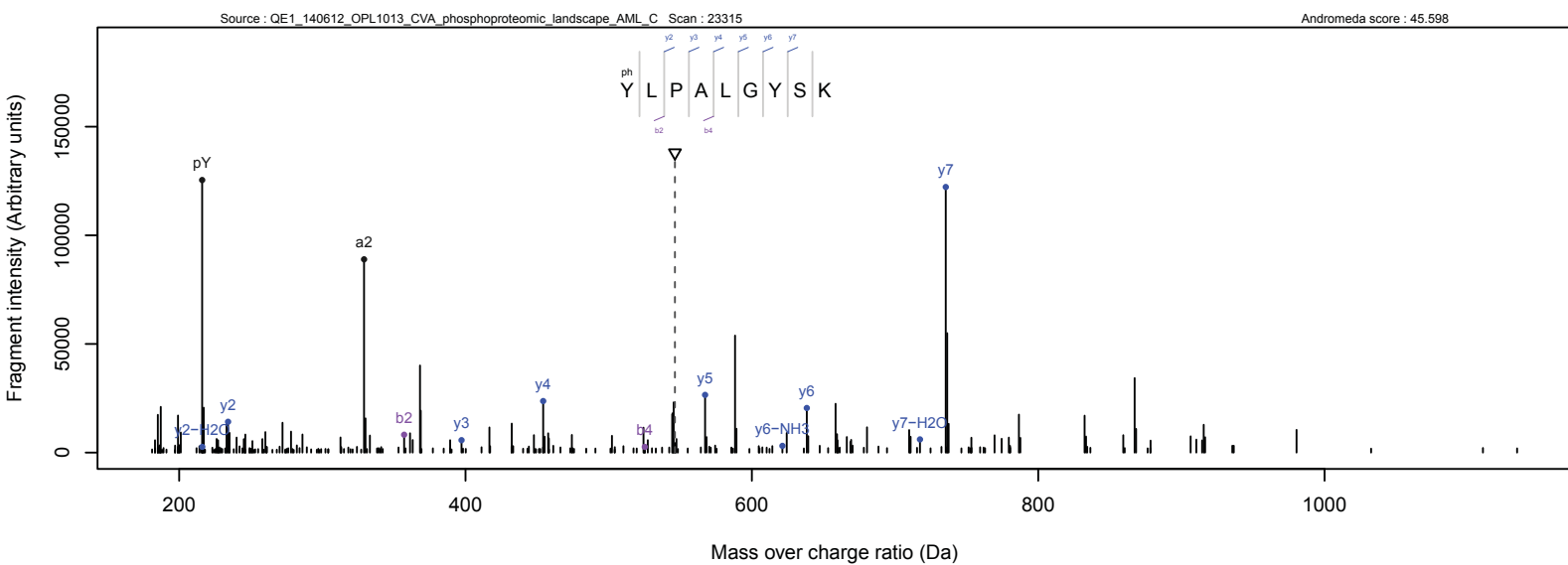
## S(ph)YVAWDREK-2+

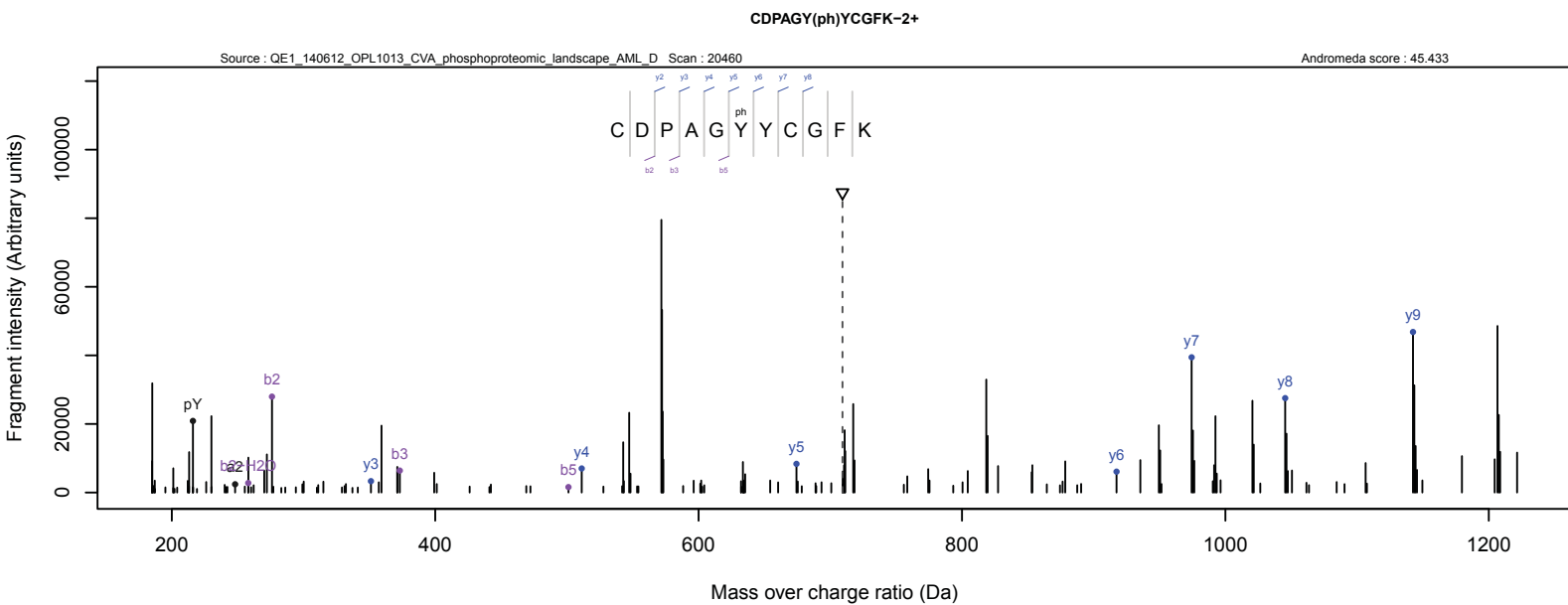
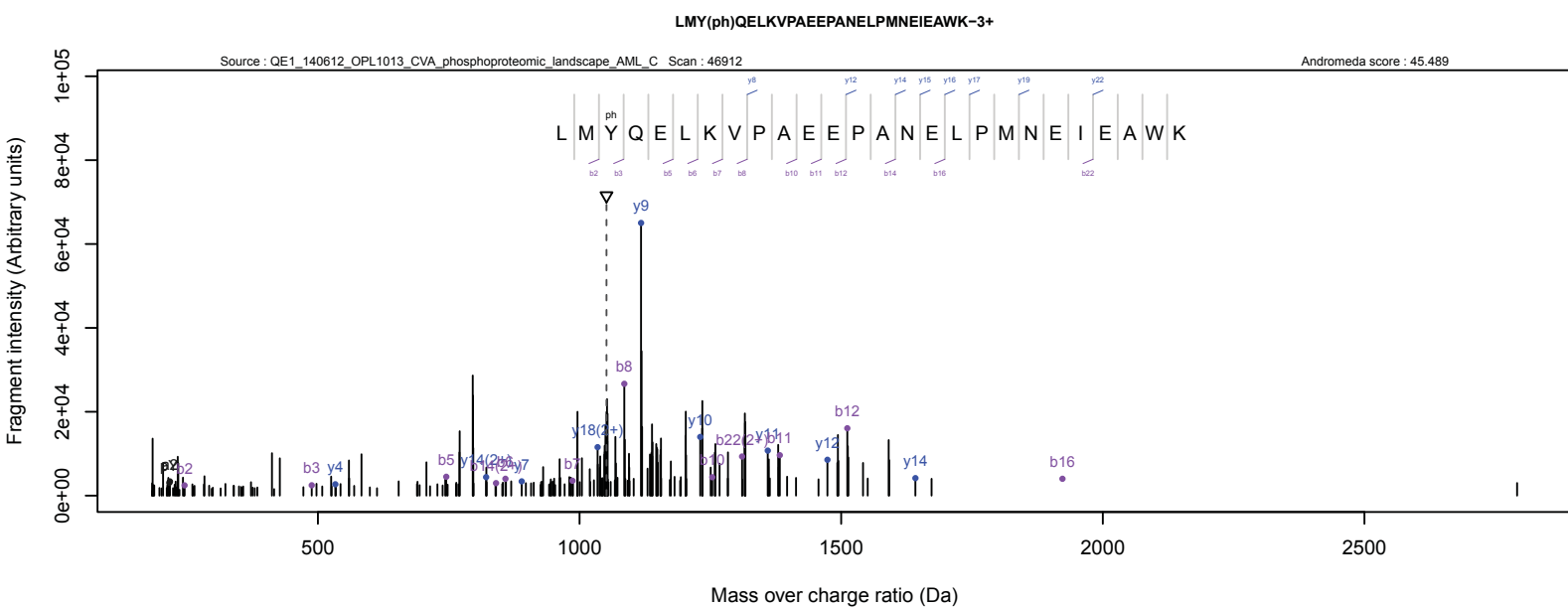
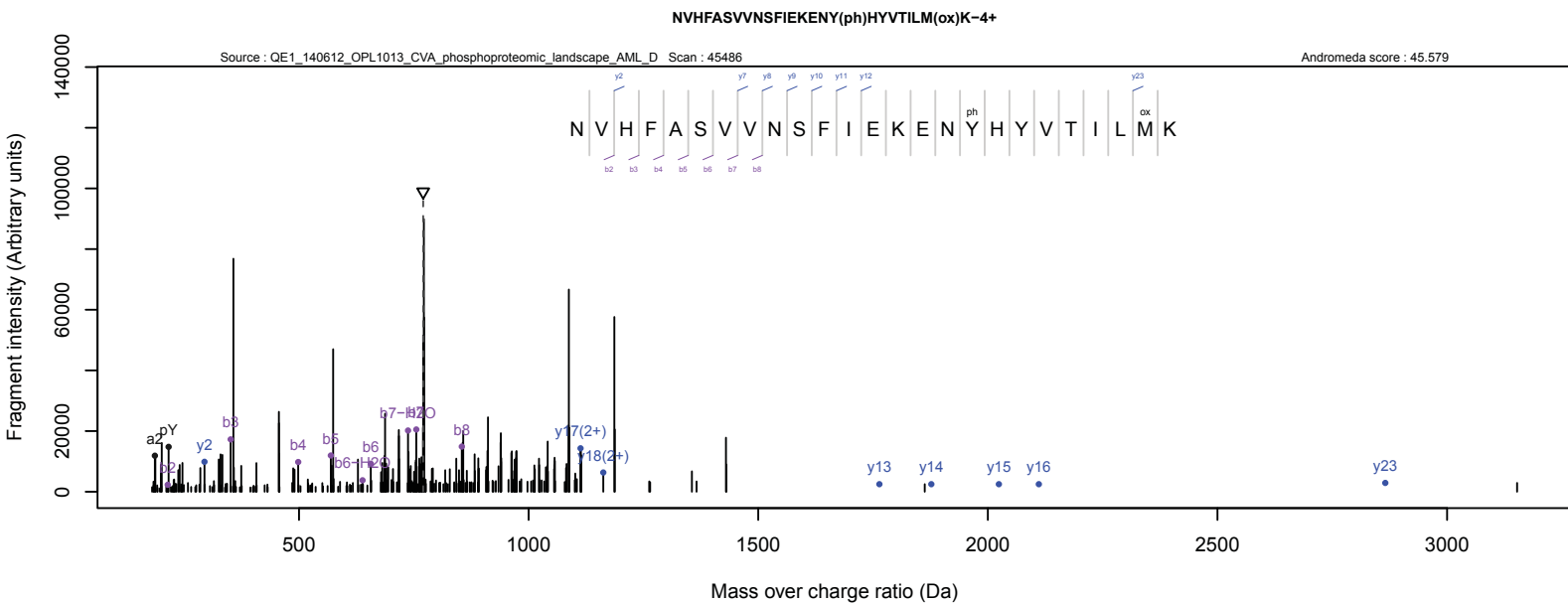


## VLPEIY(ph)LTR-2+



## Y(ph)LPALGYSK-2+

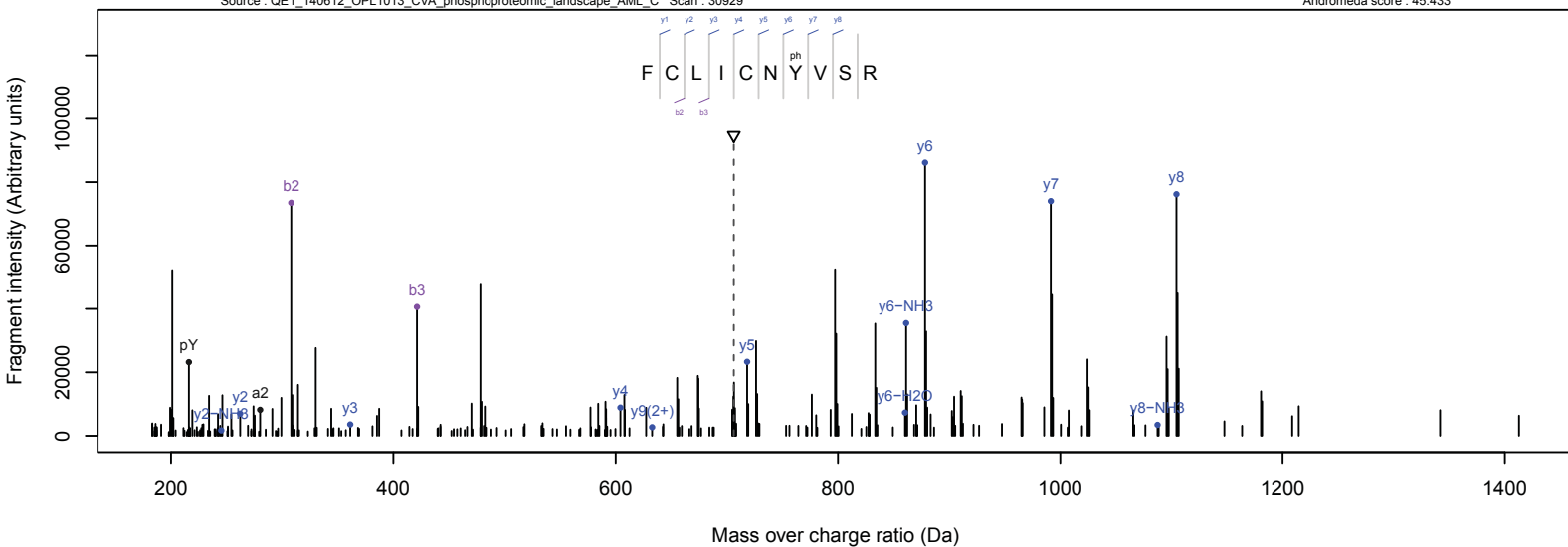




## FCLICNY(ph)VSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30929

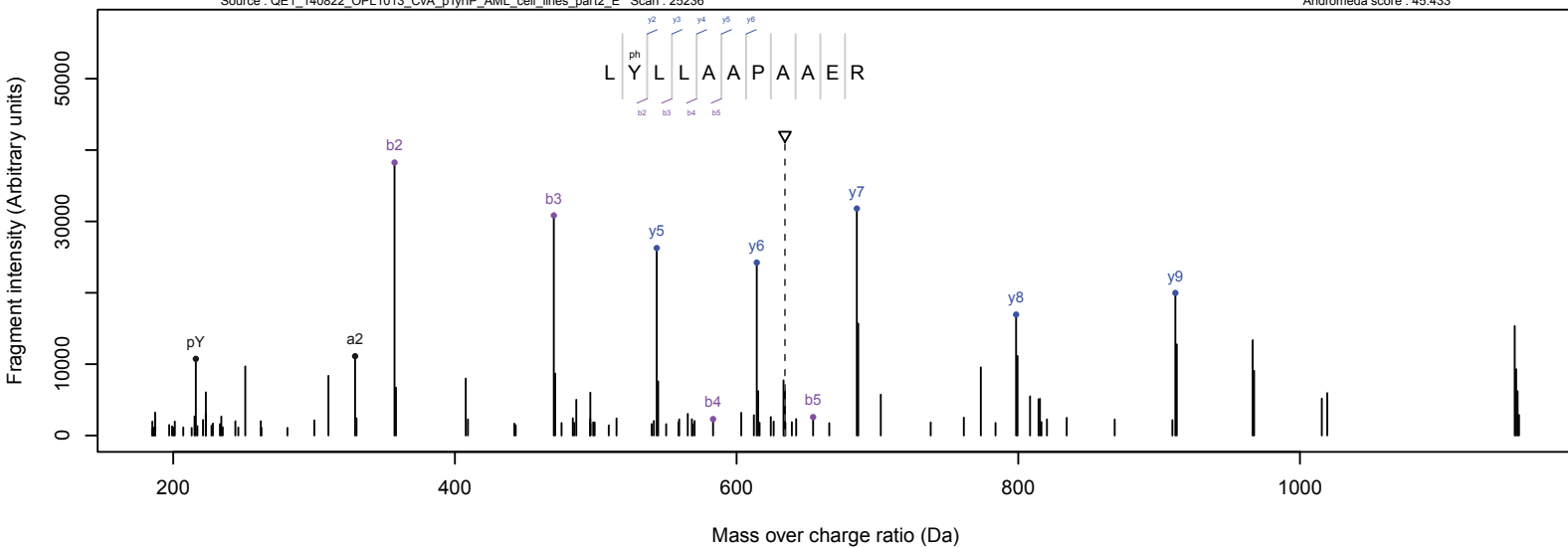
Andromeda score : 45.433



## LY(ph)LLAAPAAER-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 25236

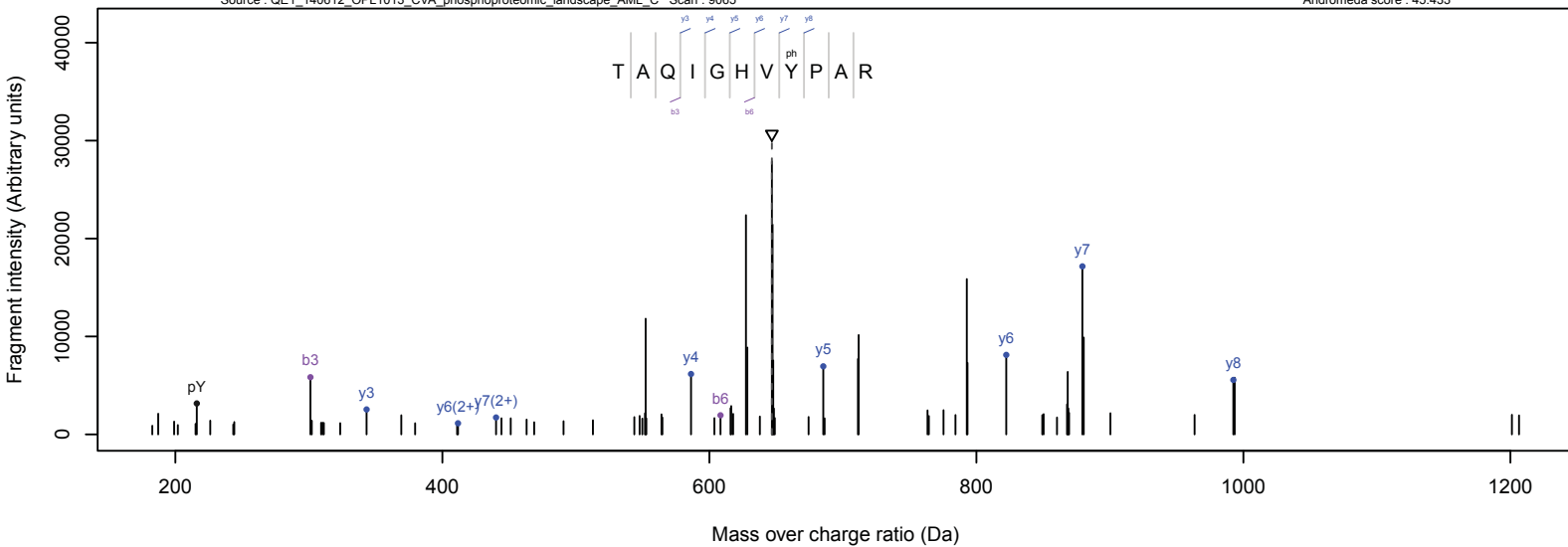
Andromeda score : 45.433



## TAQIGHVY(ph)PAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 9065

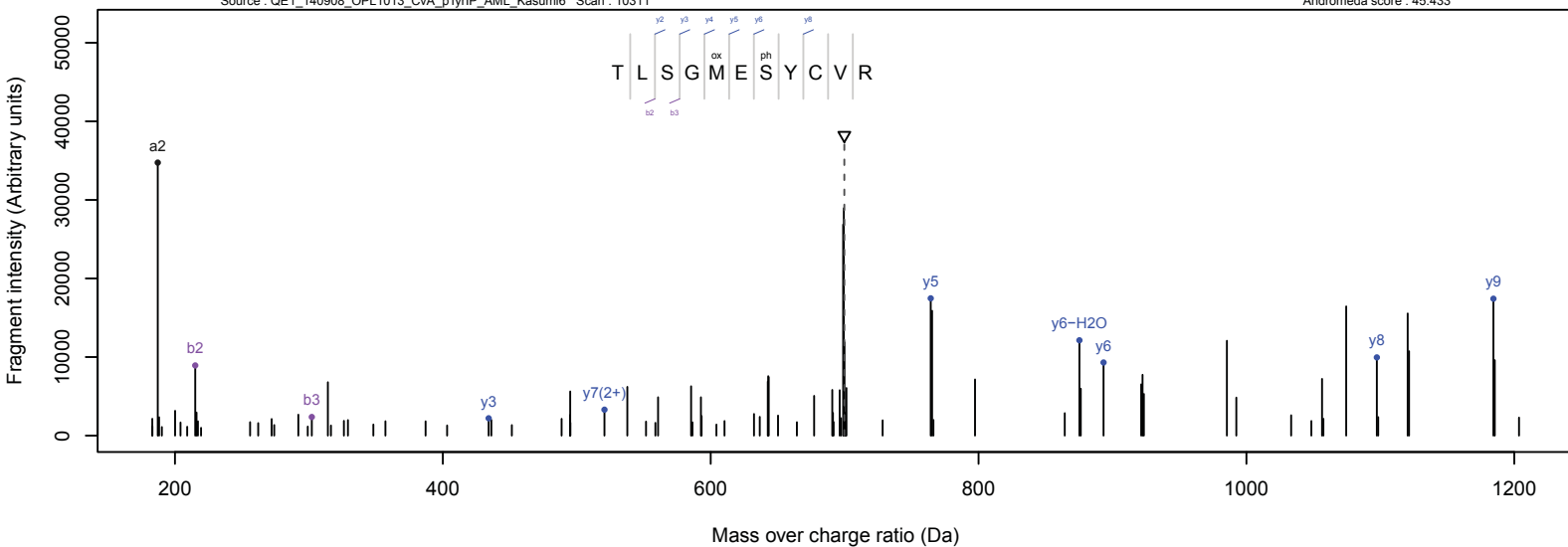
Andromeda score : 45.433



## TLSGM(ox)ES(ph)YCVR-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_Kasumi6 Scan : 10311

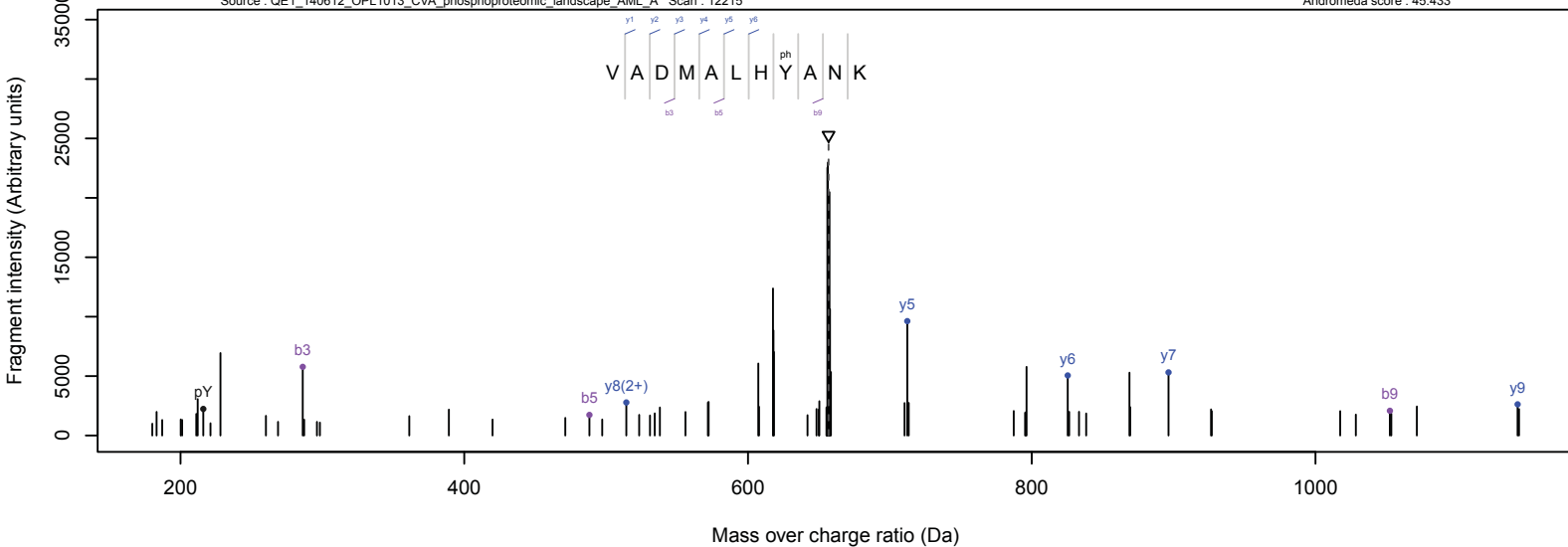
Andromeda score : 45.433



## VADMALHY(ph)ANK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12215

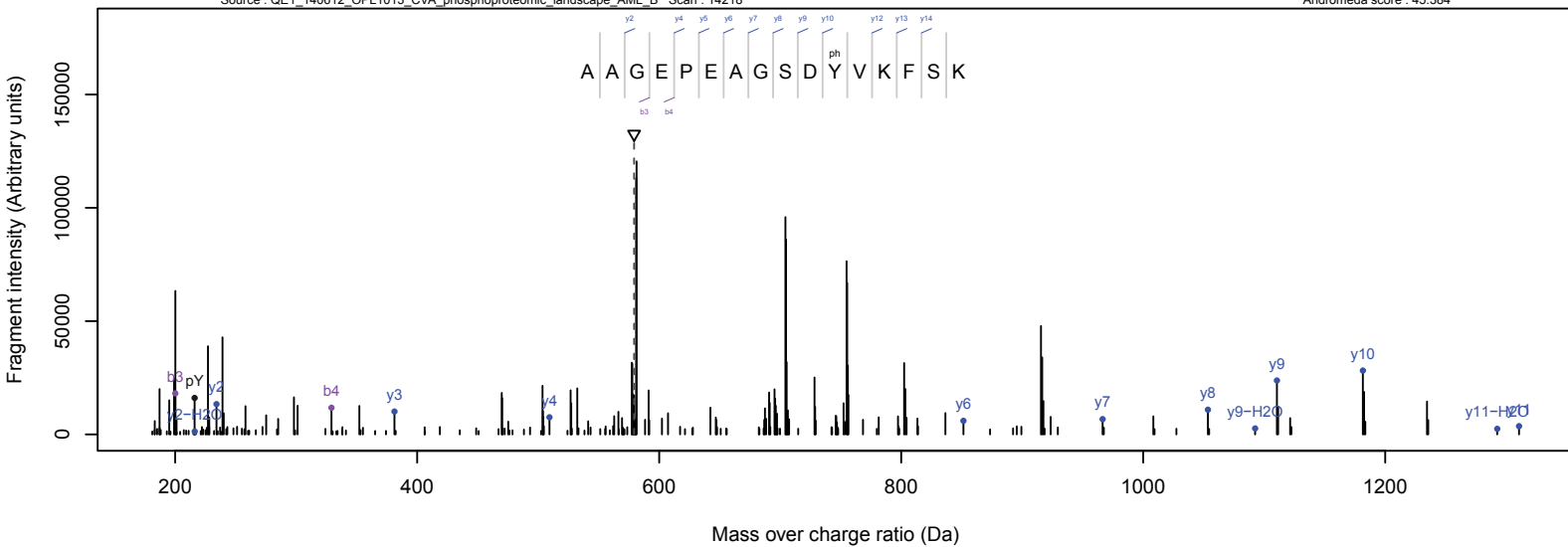
Andromeda score : 45.433



## AAGEPEAGSDY(ph)VKFSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14218

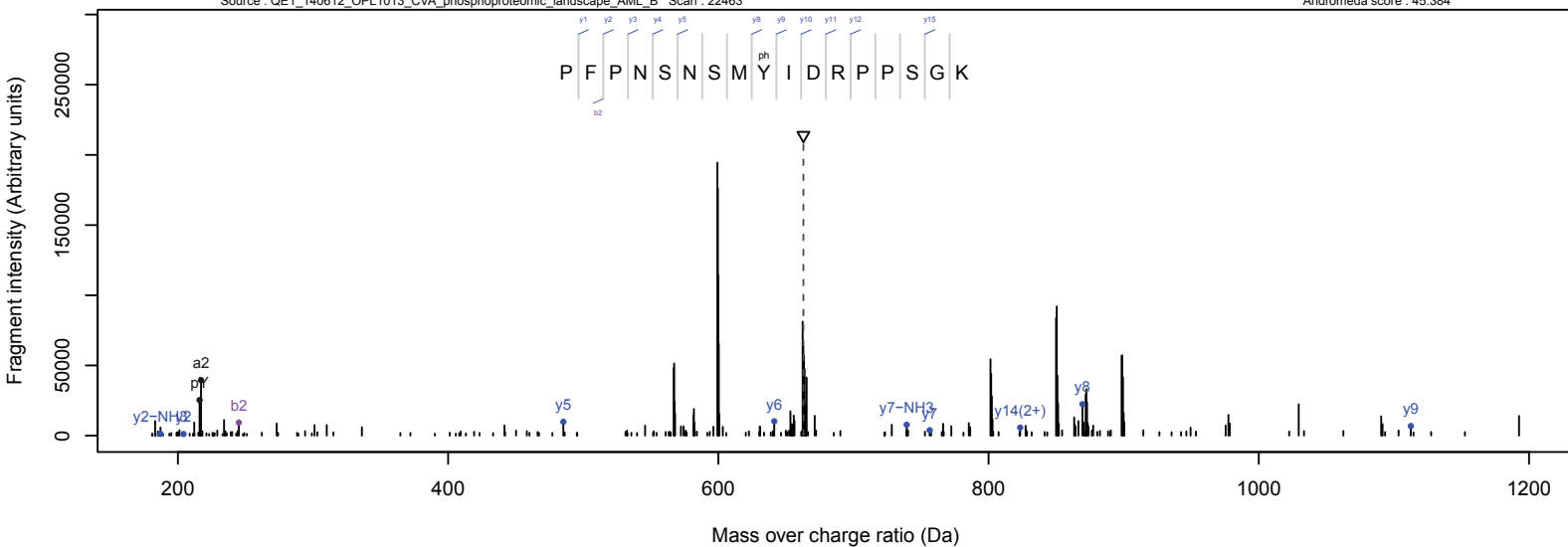
Andromeda score : 45.384



## PFPNSMY(ph)DRPPSGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 22463

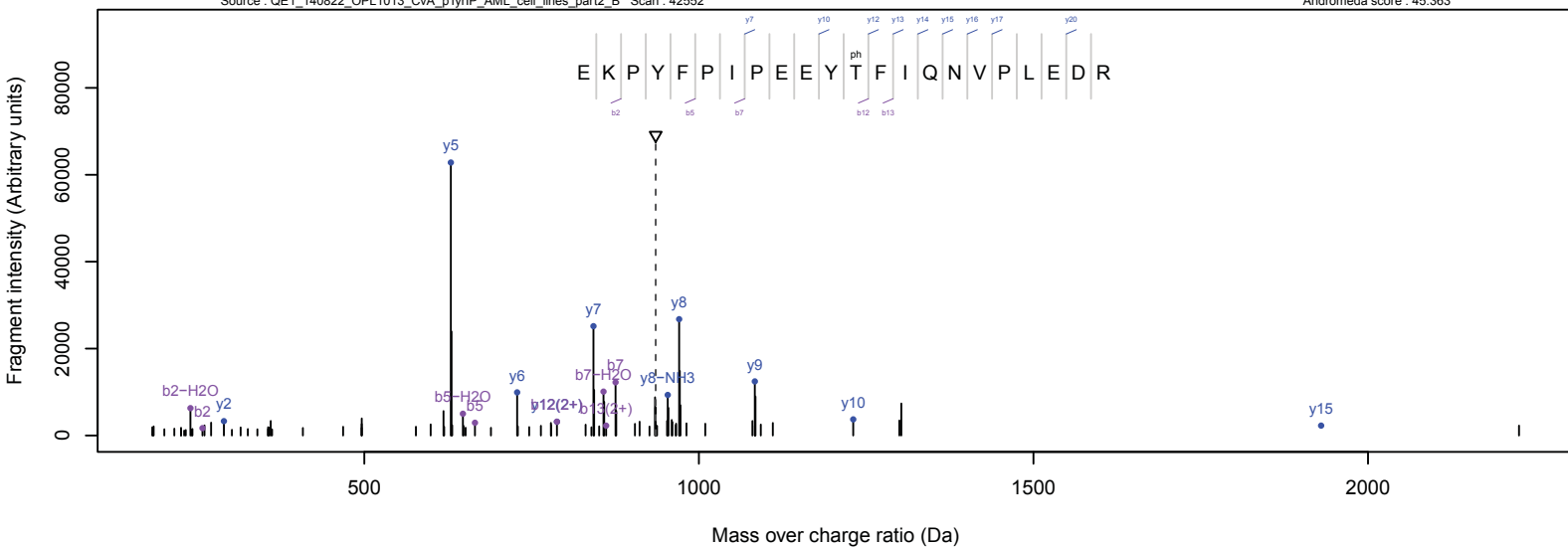
Andromeda score : 45.384



## EKPYFPIPEEY(ph)FIQNVPLEDR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 42552

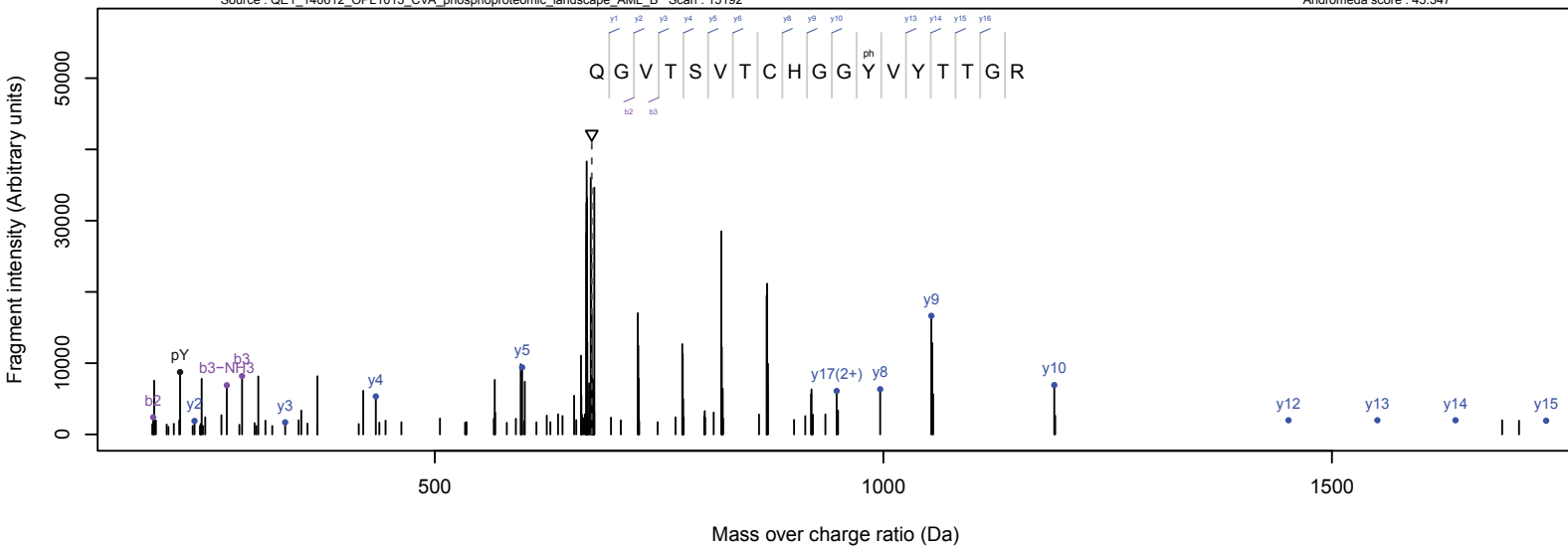
Andromeda score : 45.363



## QGVTSVTCGGY(ph)VYTTGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 15192

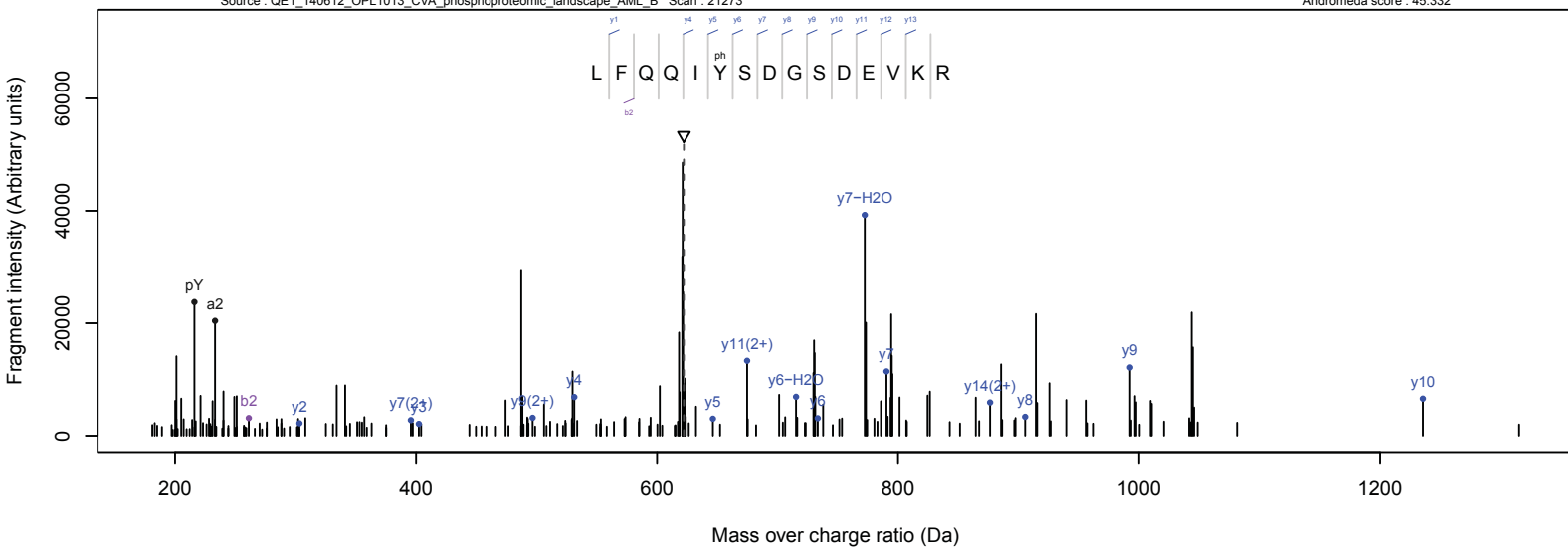
Andromeda score : 45.347



## LFQIQ(ph)SDGSDEVKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21273

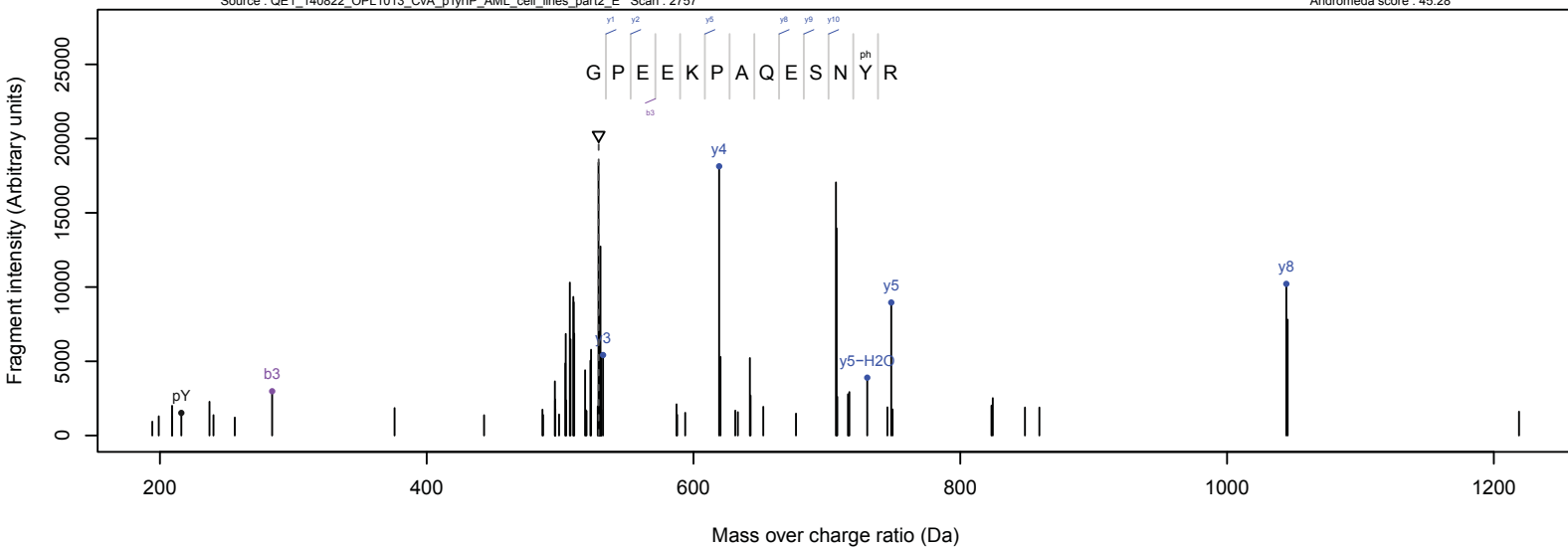
Andromeda score : 45.332



## GPEEKPAQESNY(ph)R-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 2757

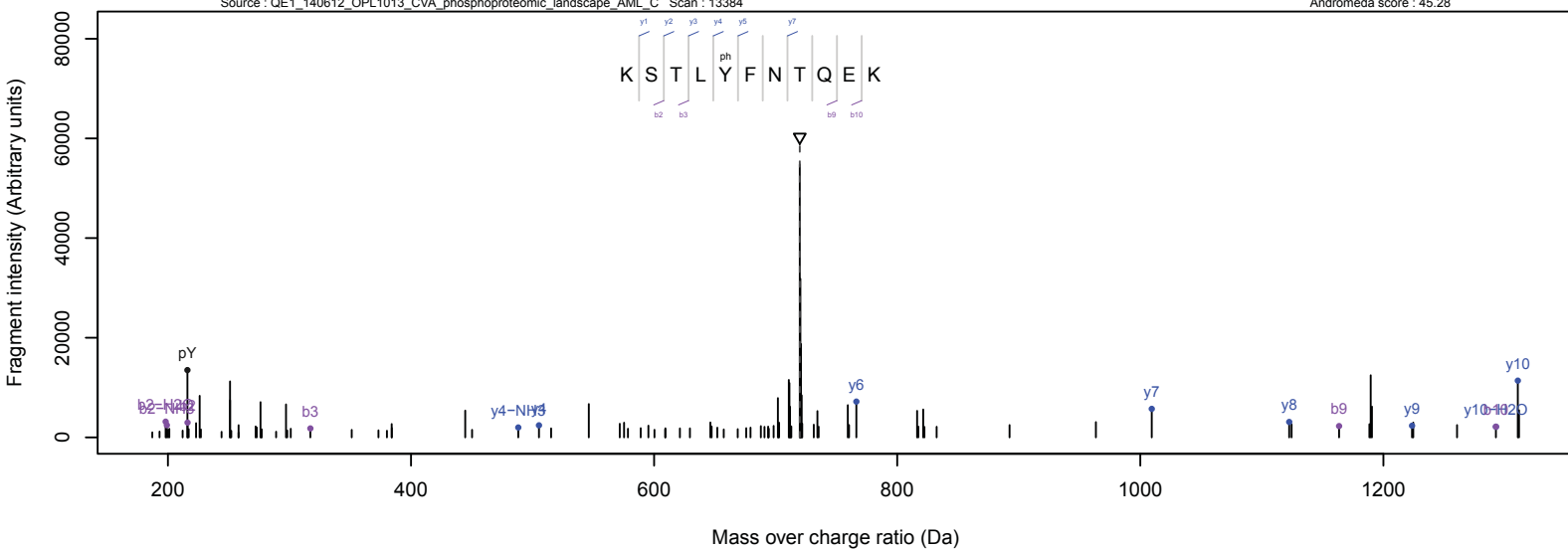
Andromeda score : 45.28



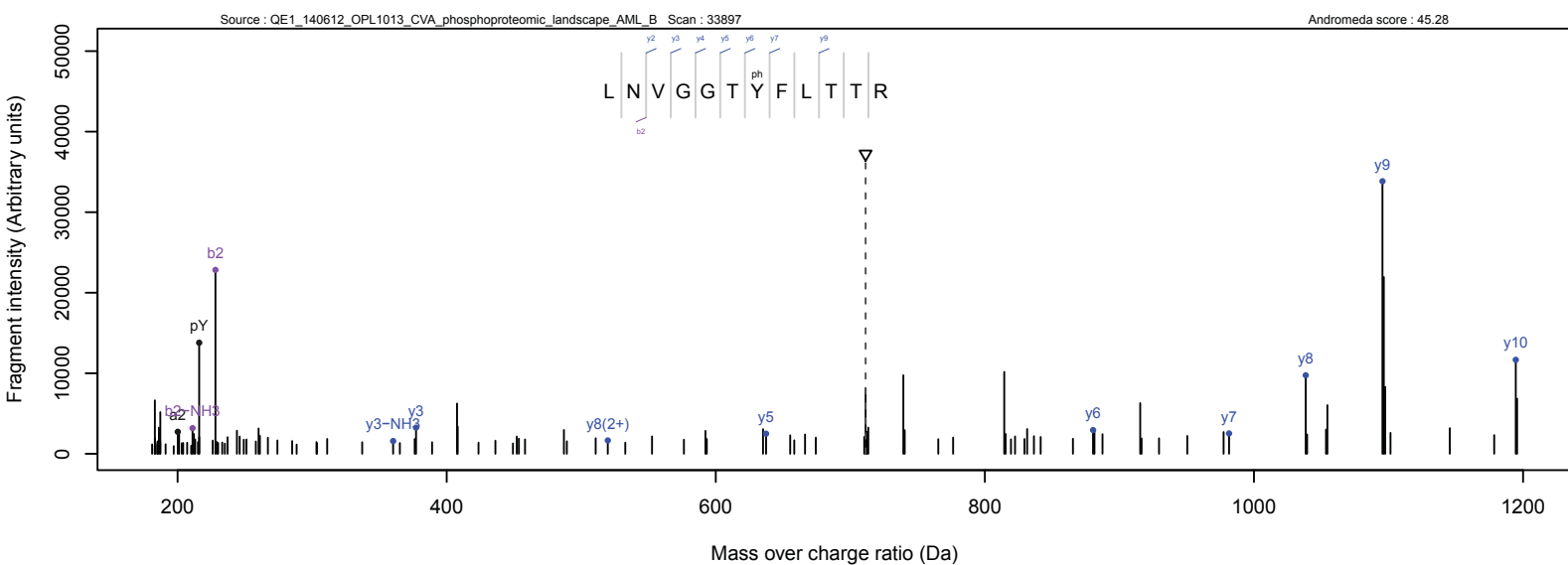
## KSTLY(ph)FNTQEK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13384

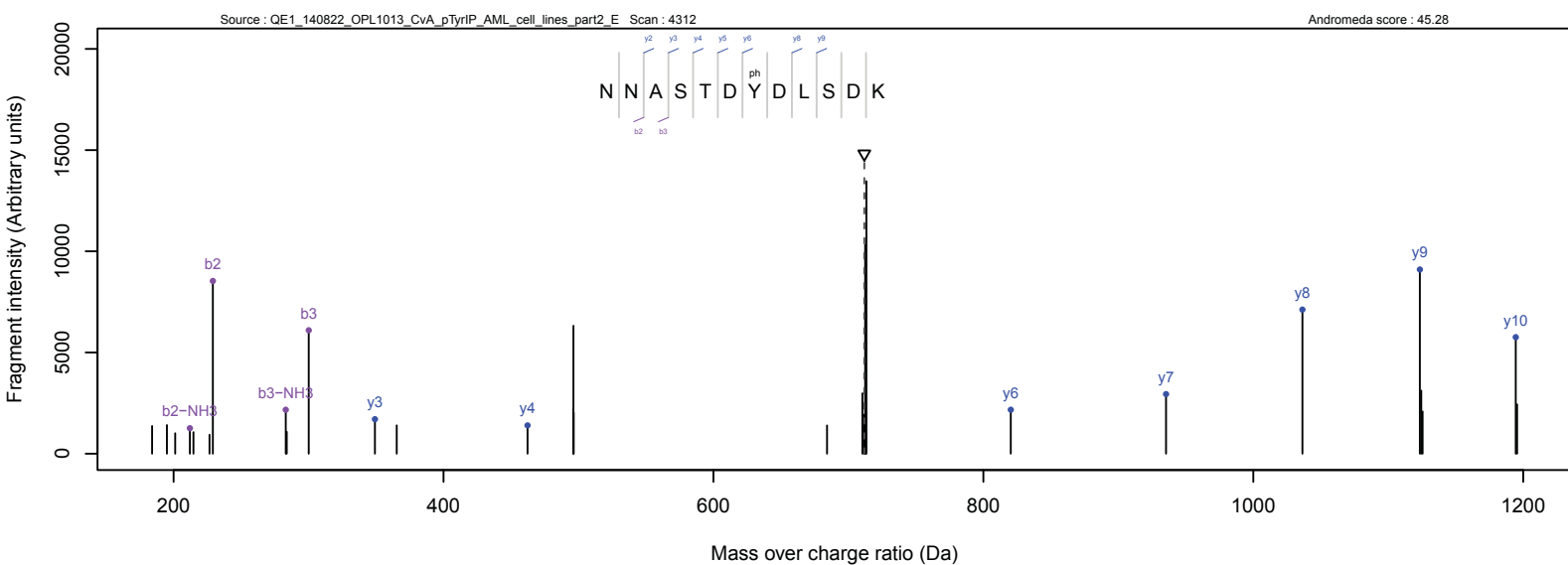
Andromeda score : 45.28



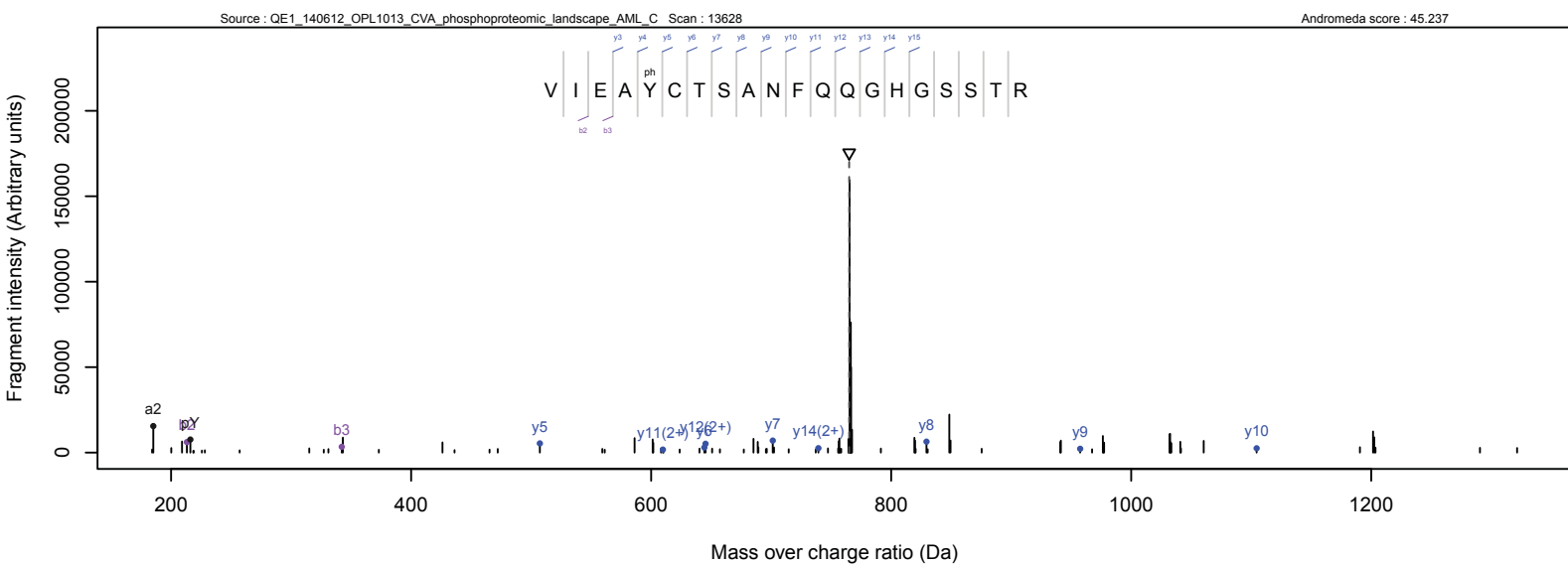
## LNVGGTY(ph)FLTR-2+



## NNASTDY(ph)DLSDK-2+



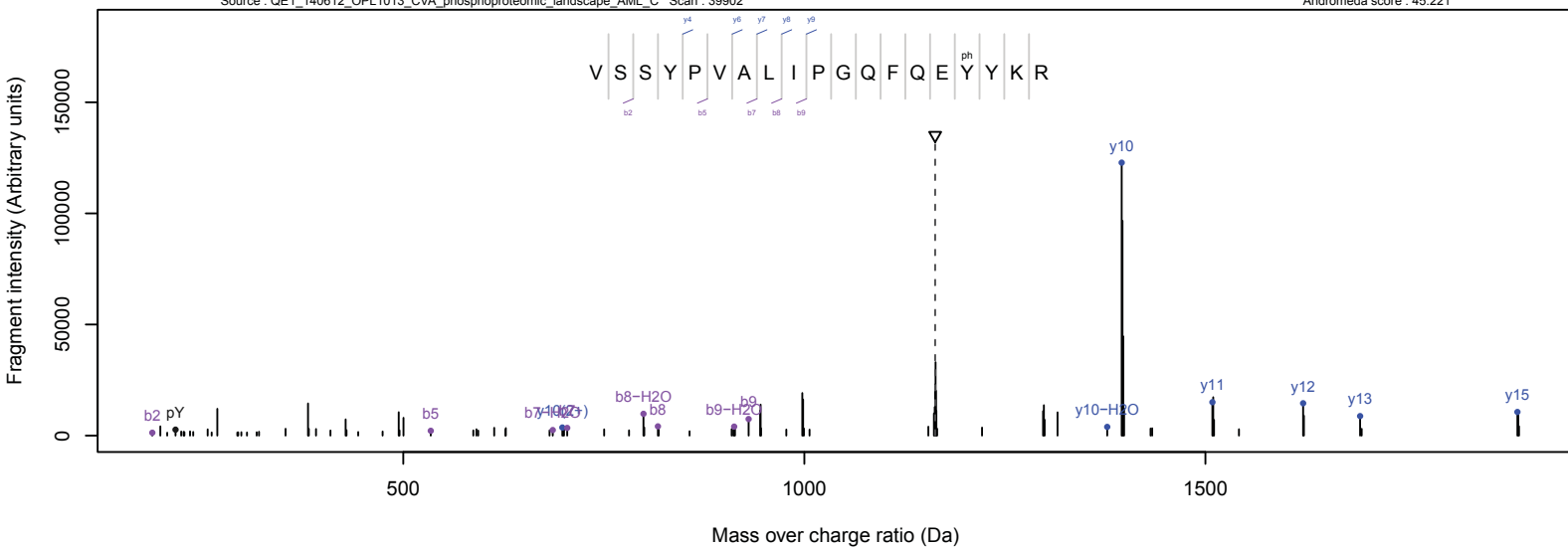
## VIEAY(ph)CTSANFQQGHGSSTR-3+



VSSYPVALIPGQFQEY<sup>ph</sup>YKR-2+

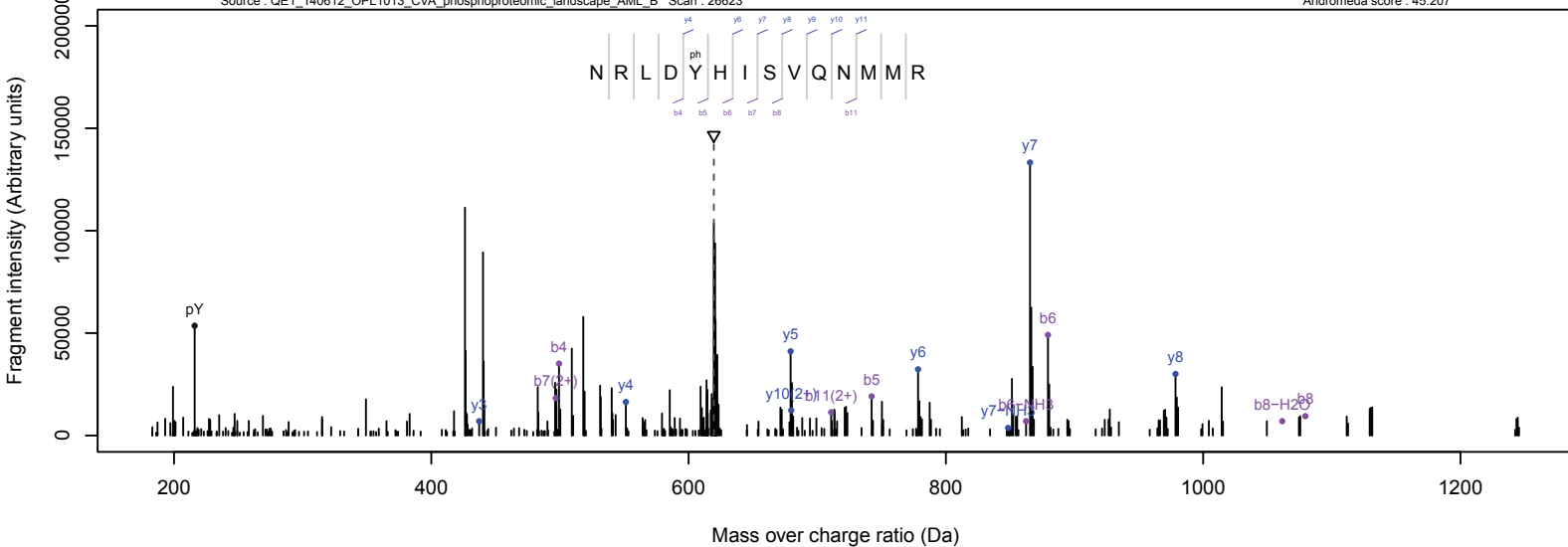
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 39902

Andromeda score : 45.221

NRLDY<sup>ph</sup>HISVQNMMR-3+

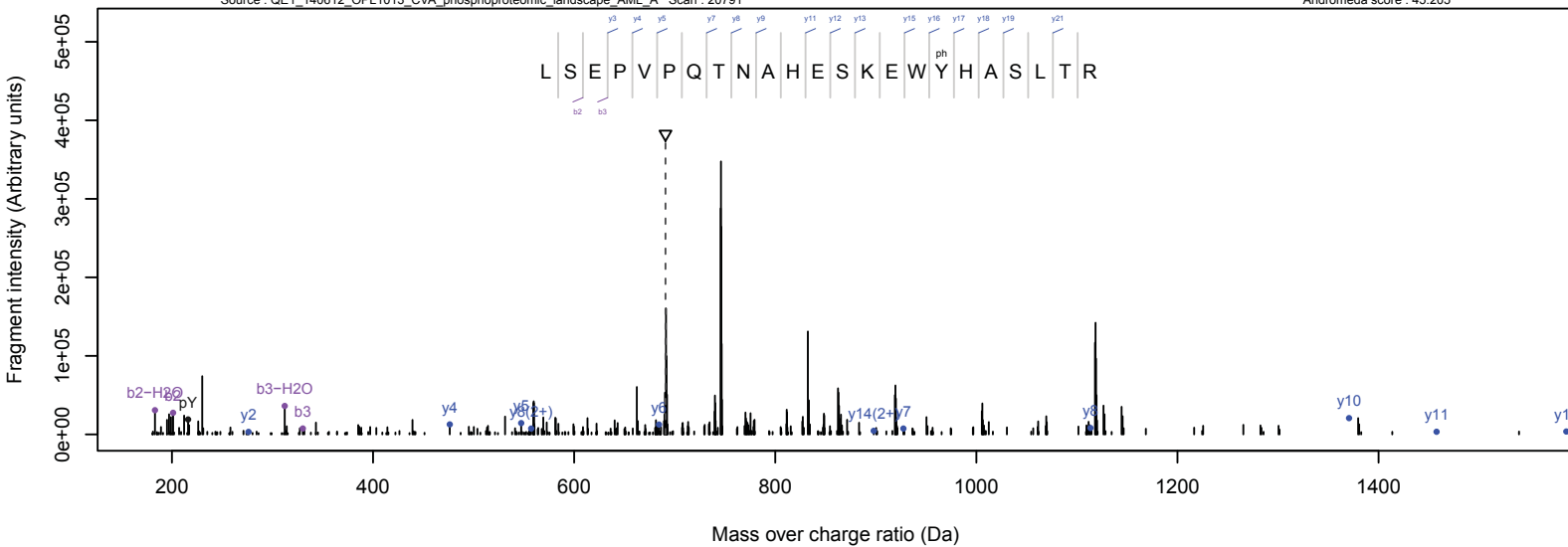
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26623

Andromeda score : 45.207

LSEPVPQTNAHESKEWY<sup>ph</sup>HASLTR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 20791

Andromeda score : 45.205

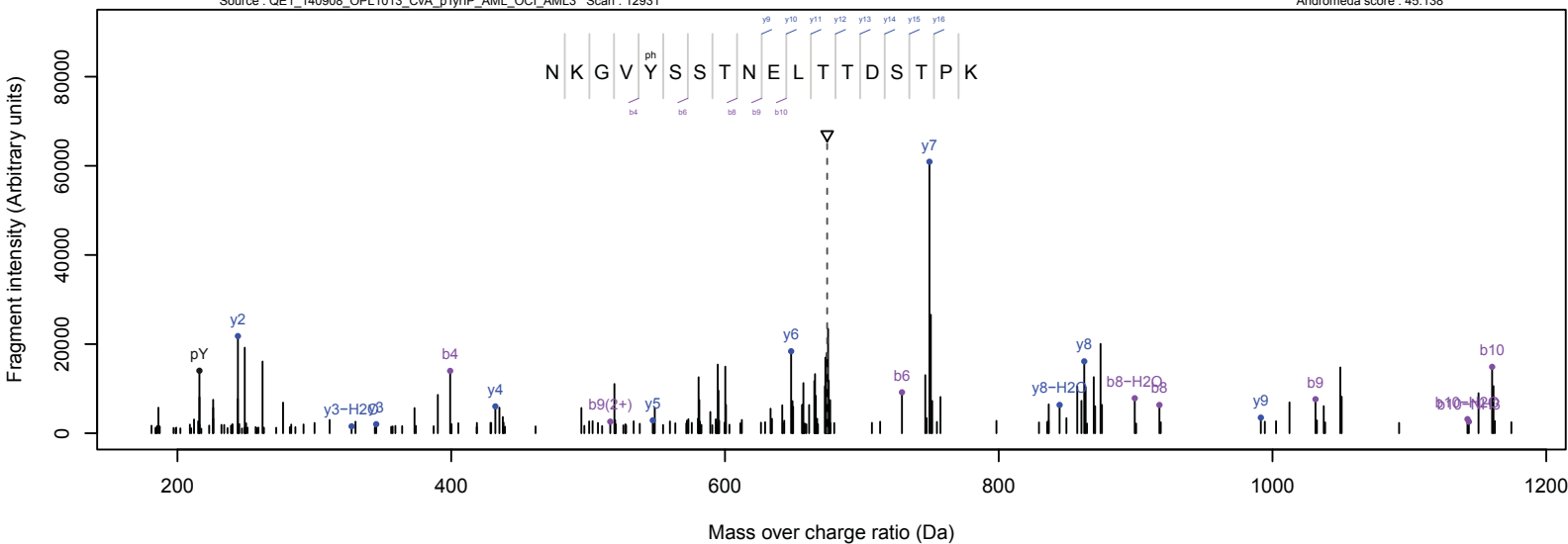




**NKGVY(ph)SSTNELTSTDSTPK-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 12931

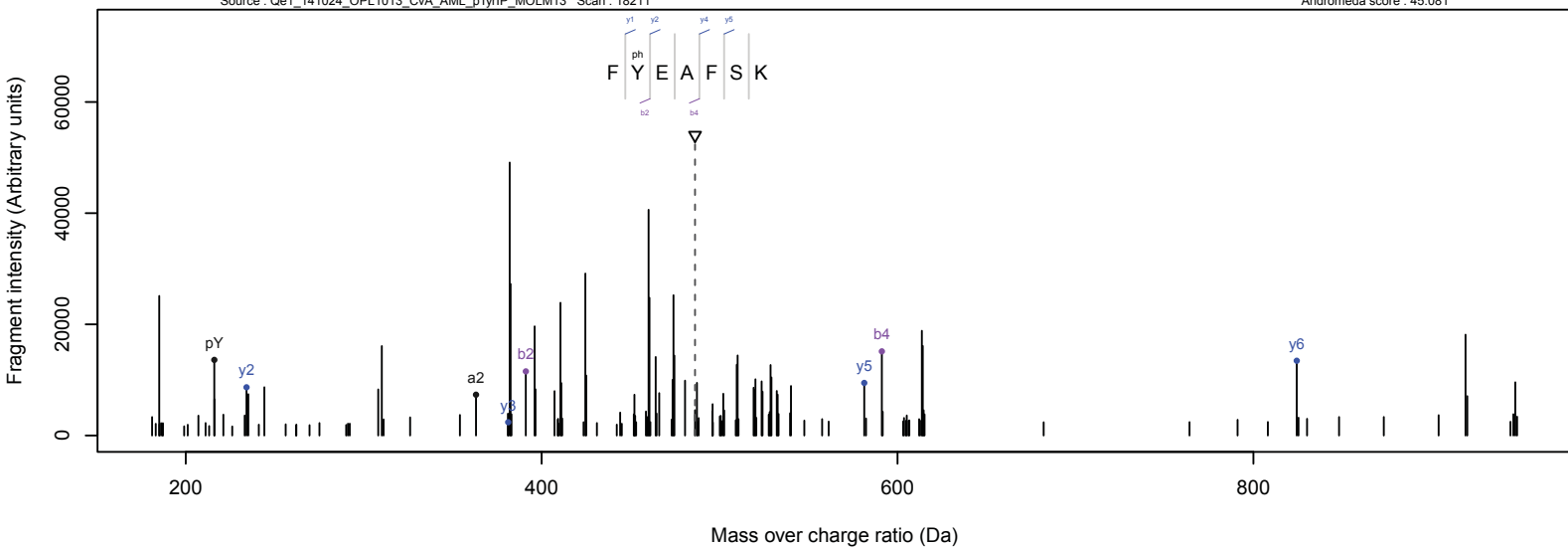
Andromeda score : 45.138



**FY(ph)EAFSK-2+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 18211

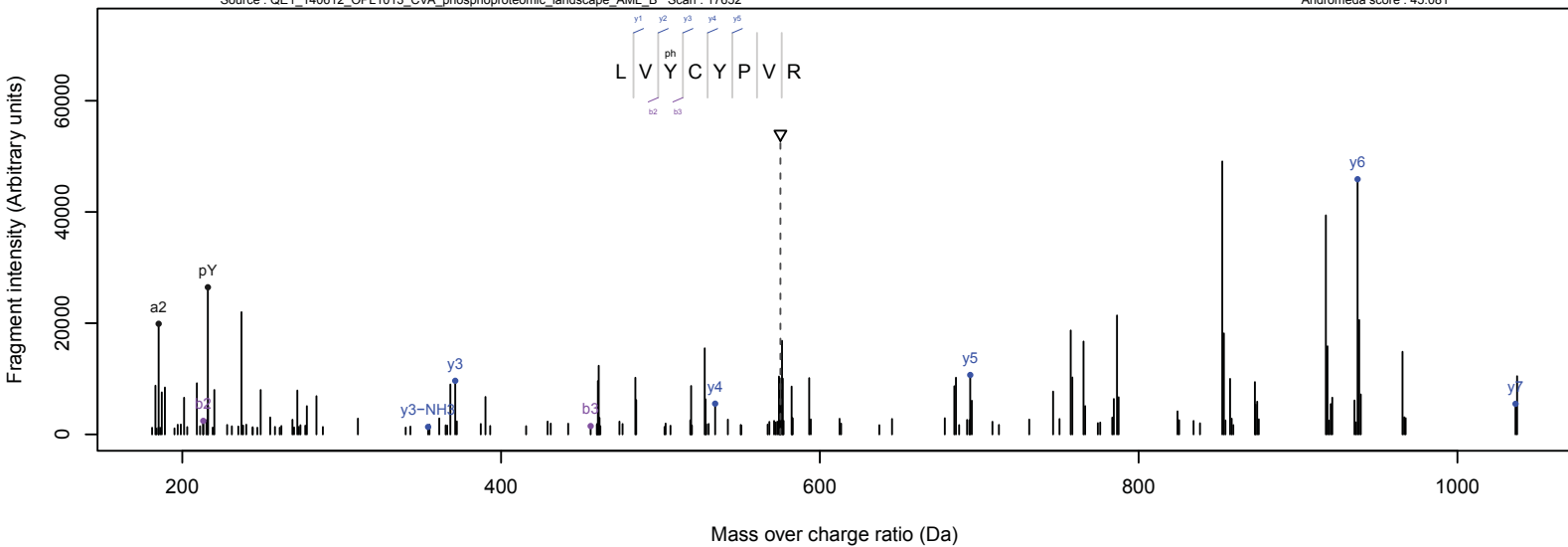
Andromeda score : 45.081



**LVY(ph)CYPVR-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17652

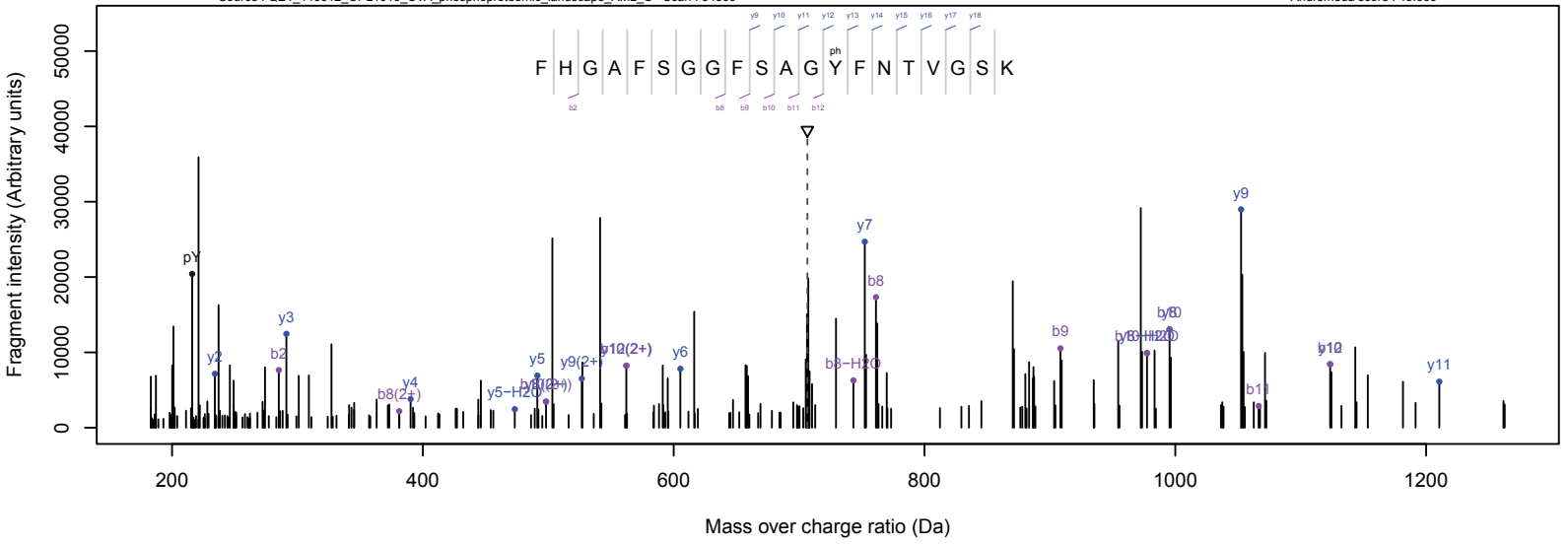
Andromeda score : 45.081



**FHGAFSGGFSAGYFNTVGSK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34309

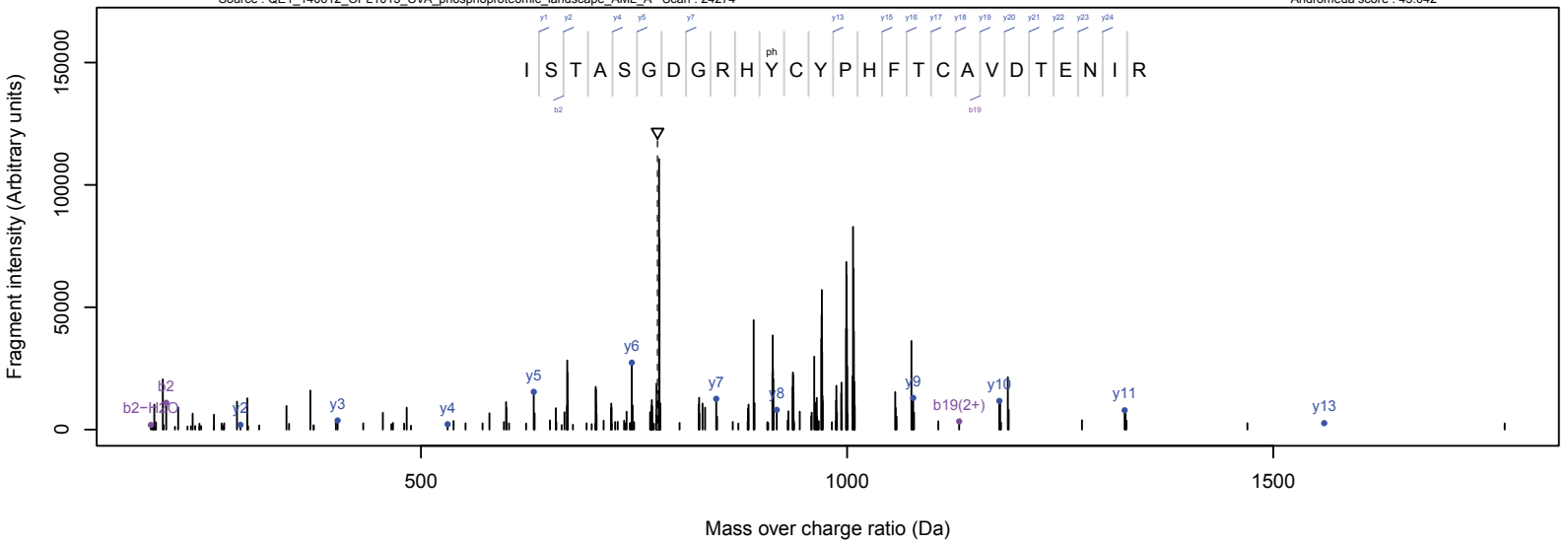
Andromeda score : 45.069



**ISTASGDGRHYCYPHFTCAVDTENIR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 24274

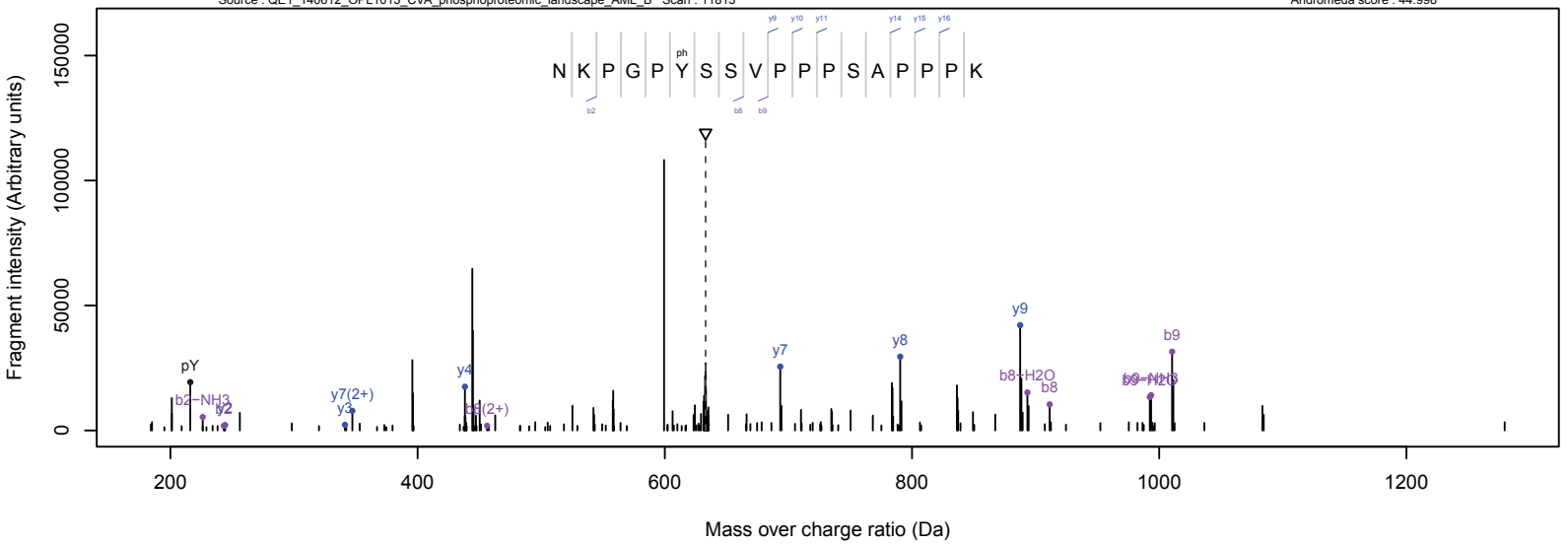
Andromeda score : 45.042



**NKPGPYSSVPPPSAPPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11813

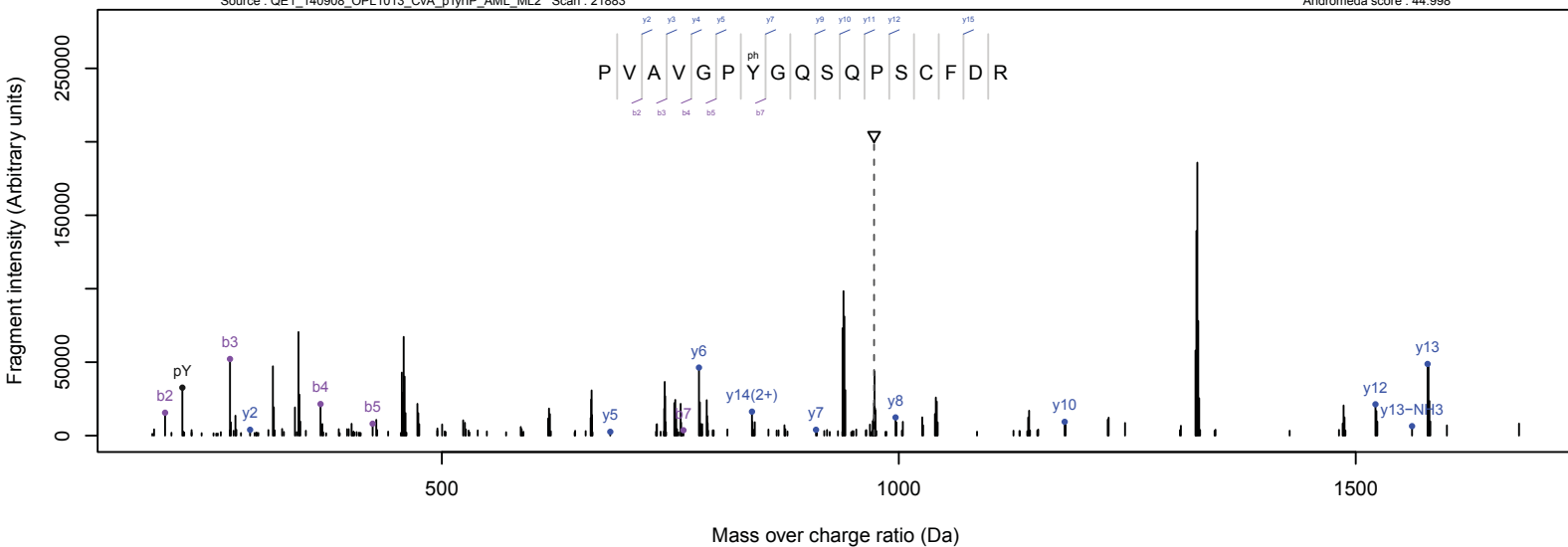
Andromeda score : 44.998



PVAVGPY(ph)GQSQPSCFDR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 21883

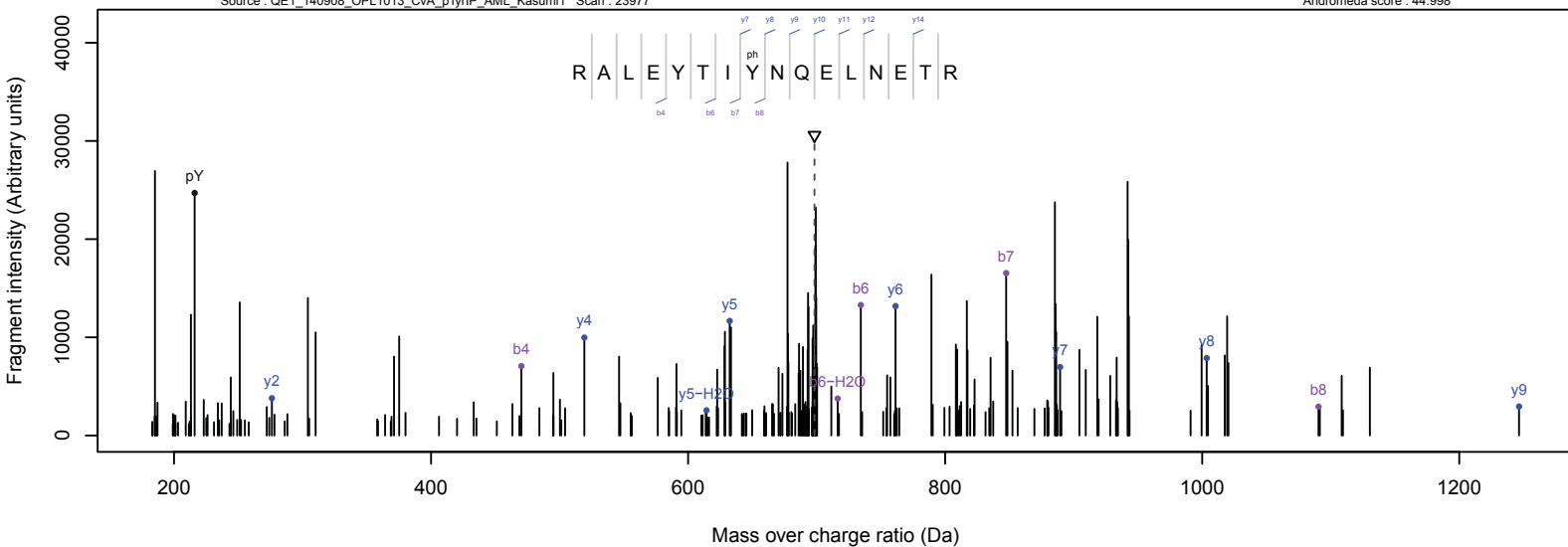
Andromeda score : 44.998



RALEYTIY(ph)NQELETNR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 23977

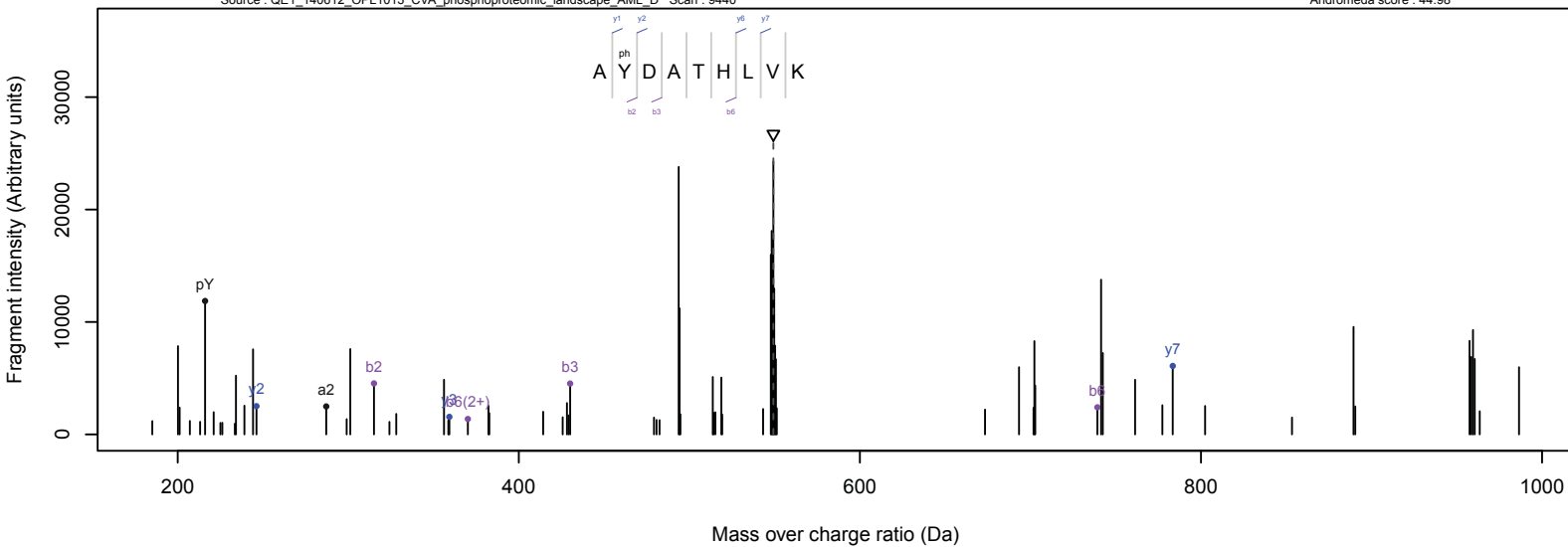
Andromeda score : 44.998



AY(ph)DATHLVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9440

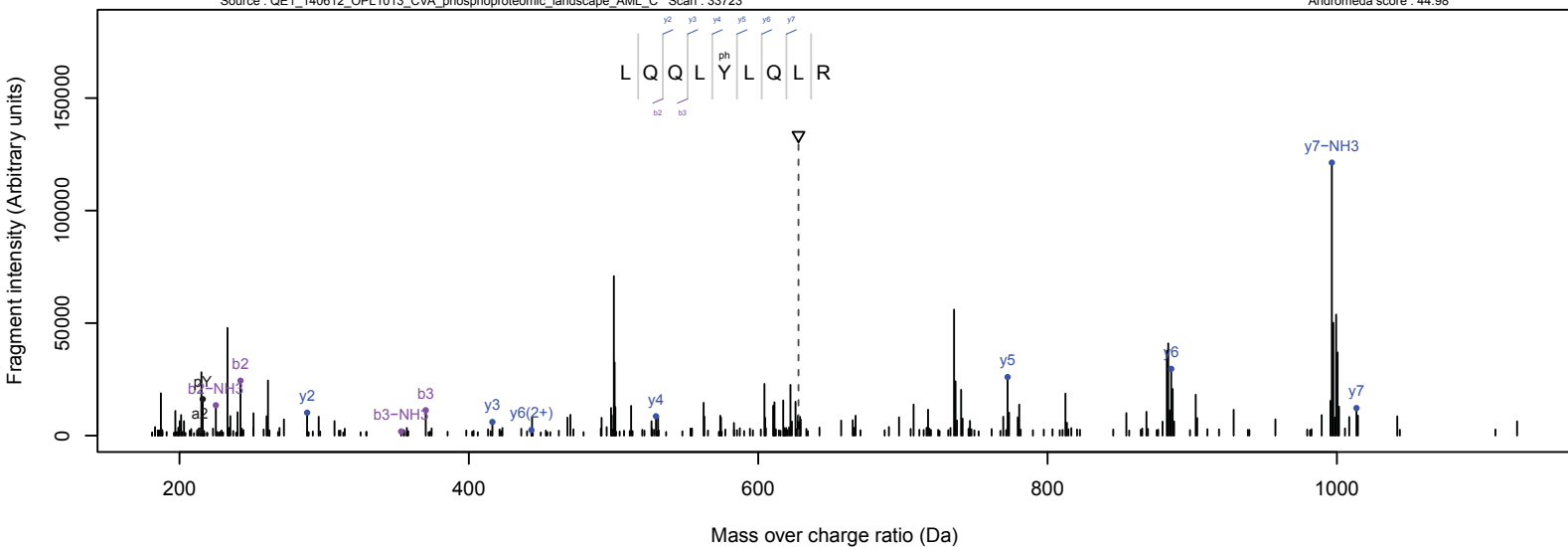
Andromeda score : 44.98



## LQQLY(ph)LQLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33723

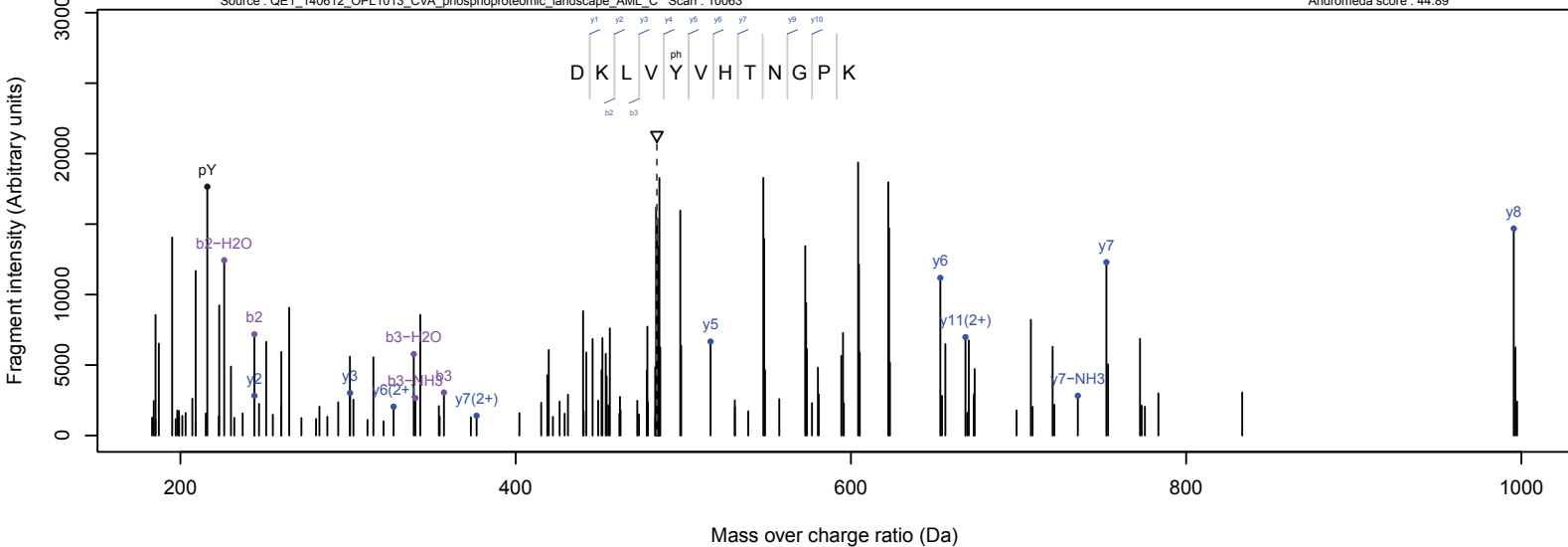
Andromeda score : 44.98



## DKLVY(ph)VHTNGPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10063

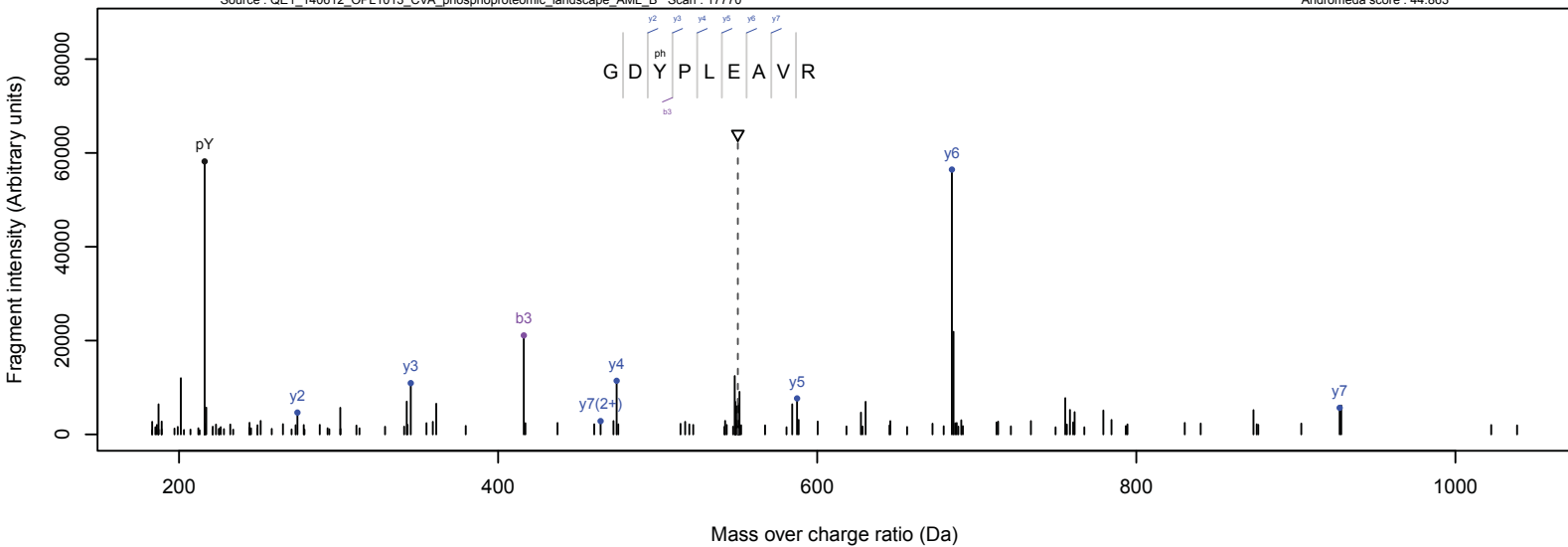
Andromeda score : 44.89



## GDY(ph)PLEAVR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17770

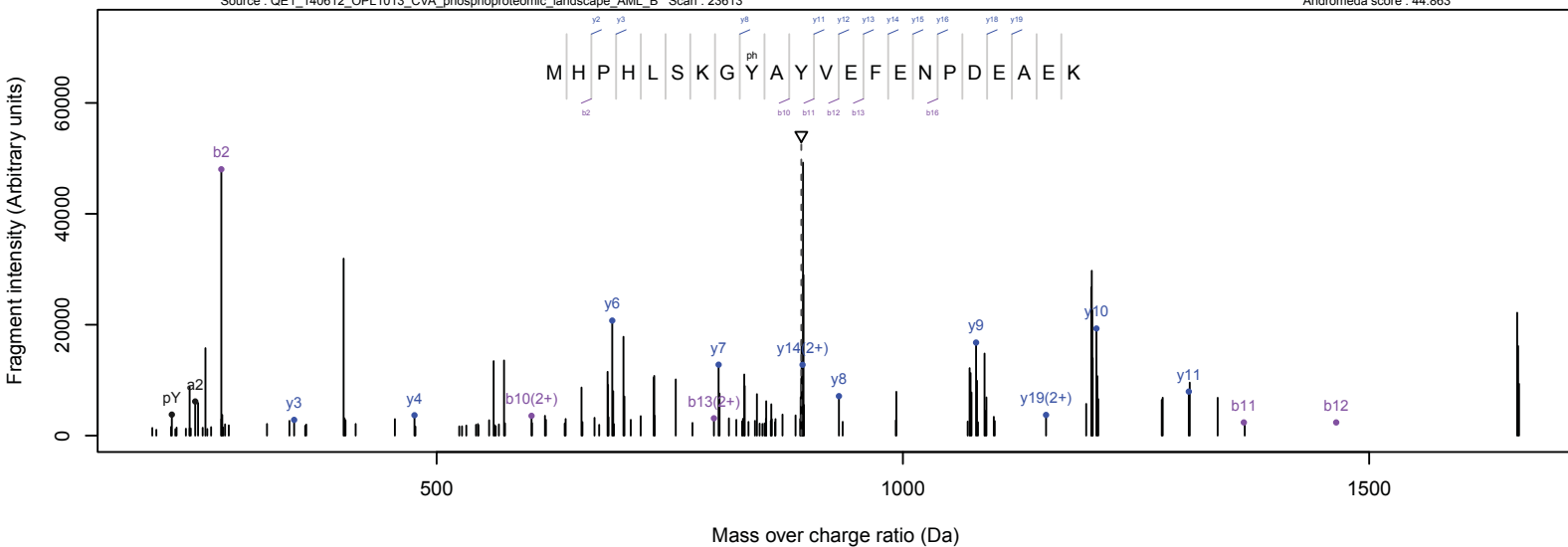
Andromeda score : 44.863



MPHLSKGY(ph)AYVEFENPDEAEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23613

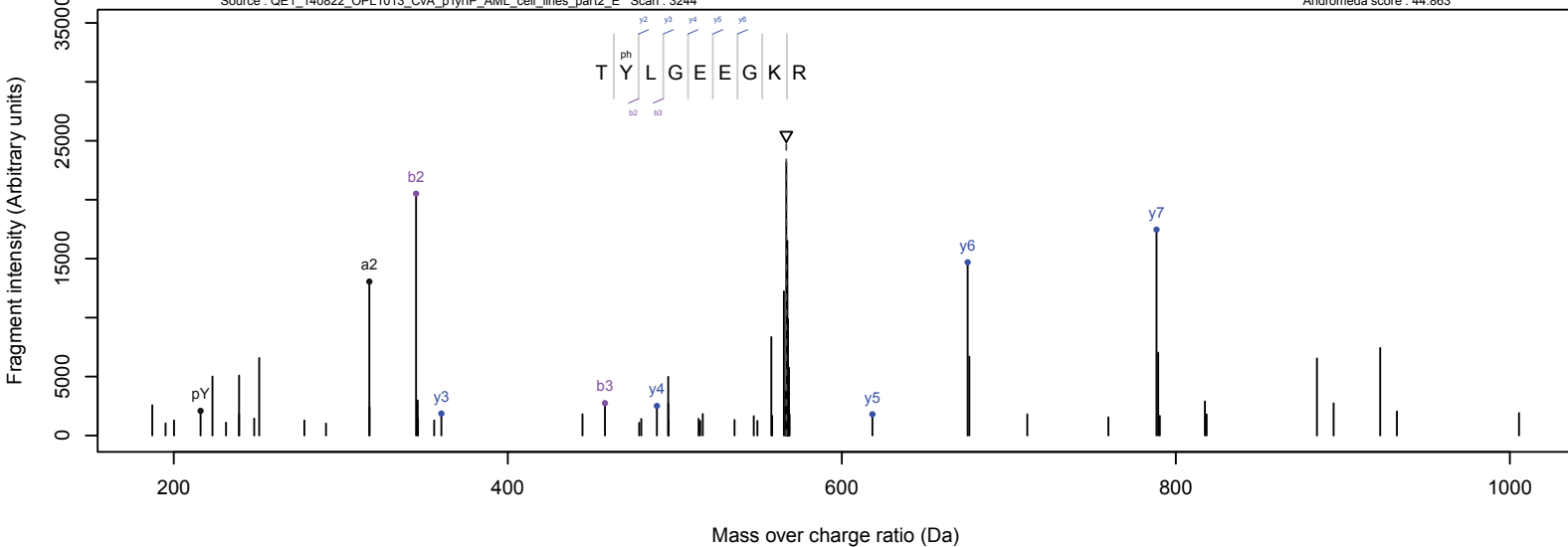
Andromeda score : 44.863



TY(ph)LGEEGKR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 3244

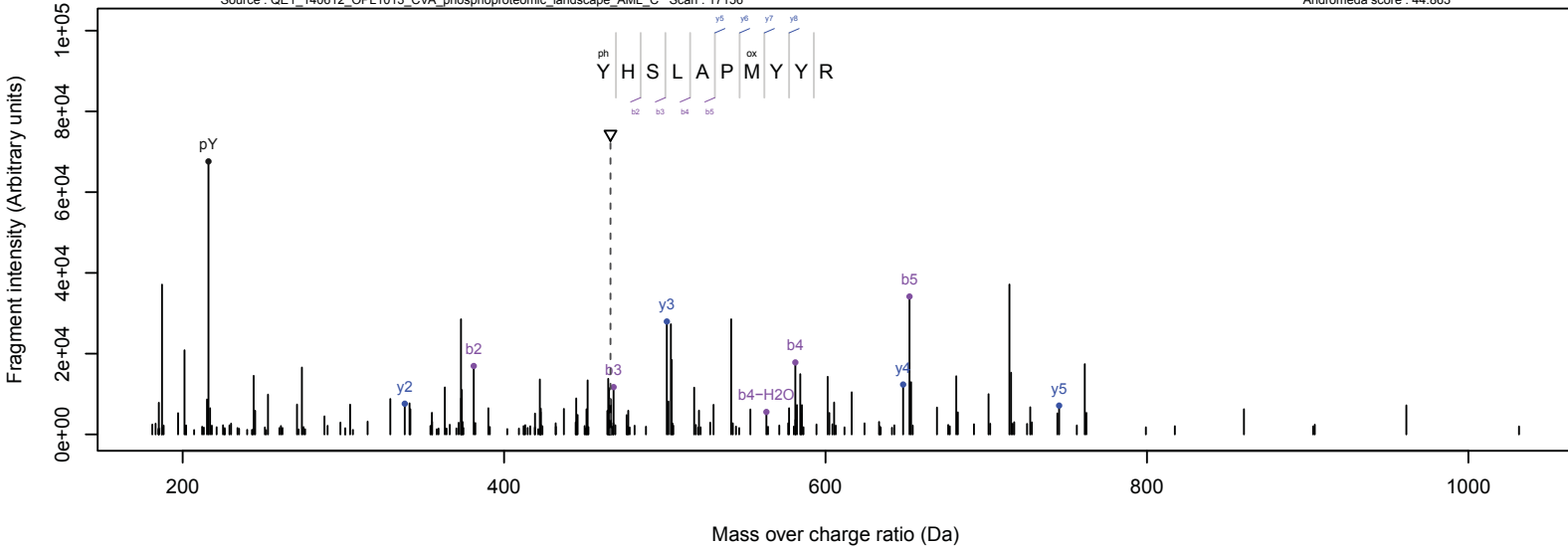
Andromeda score : 44.863



Y(ph)HSLAPM(ox)YYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17156

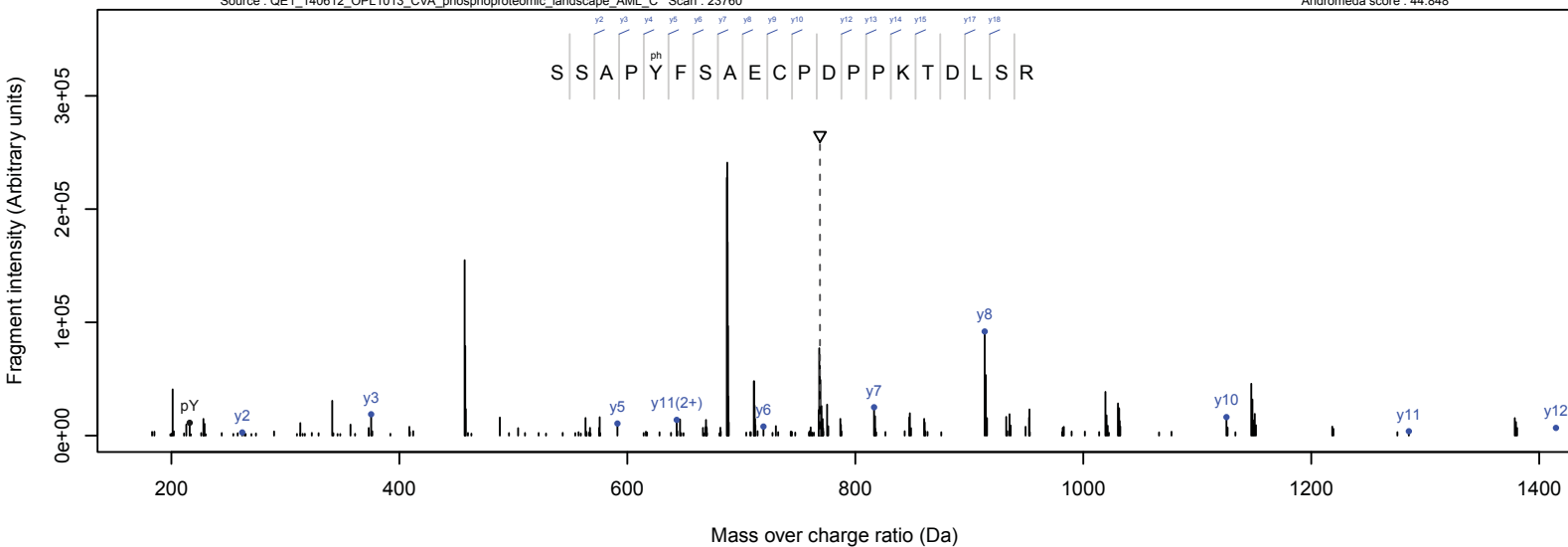
Andromeda score : 44.863



SSAPY(ph)FSAECPDPPKTLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23760

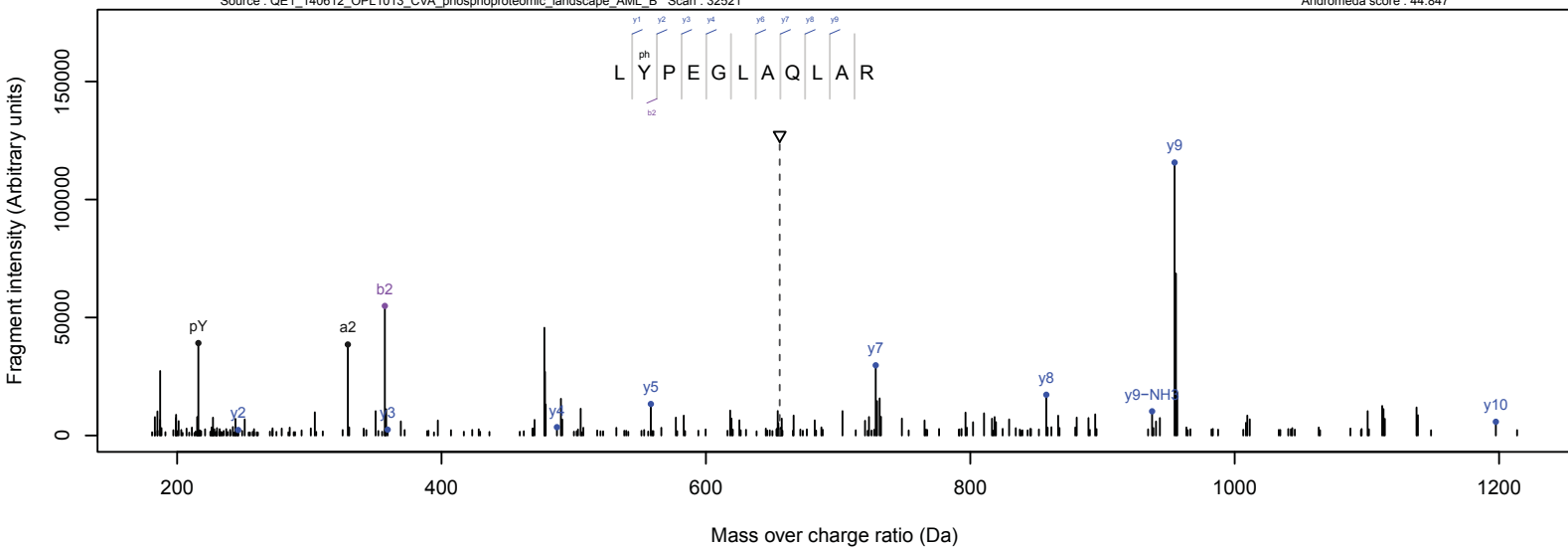
Andromeda score : 44.848



LY(ph)PEGLAQLAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32521

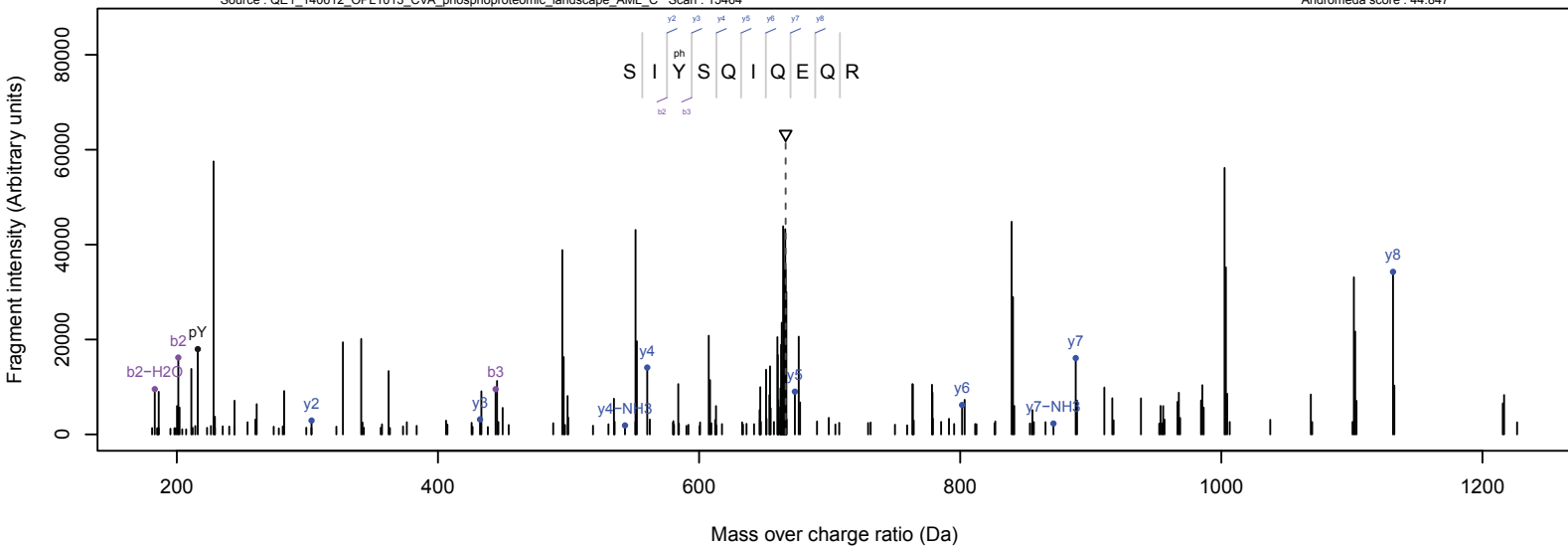
Andromeda score : 44.847



SIY(ph)SQIQEQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15464

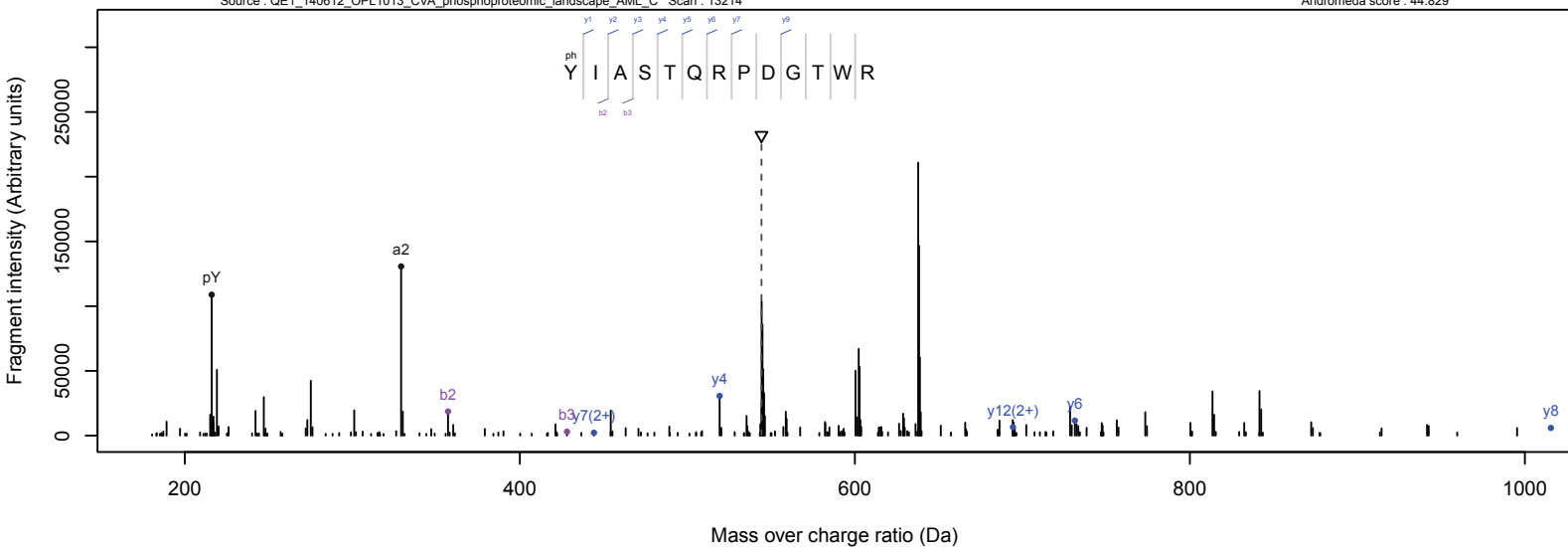
Andromeda score : 44.847



## Y(ph)IASTQRPDGTWR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13214

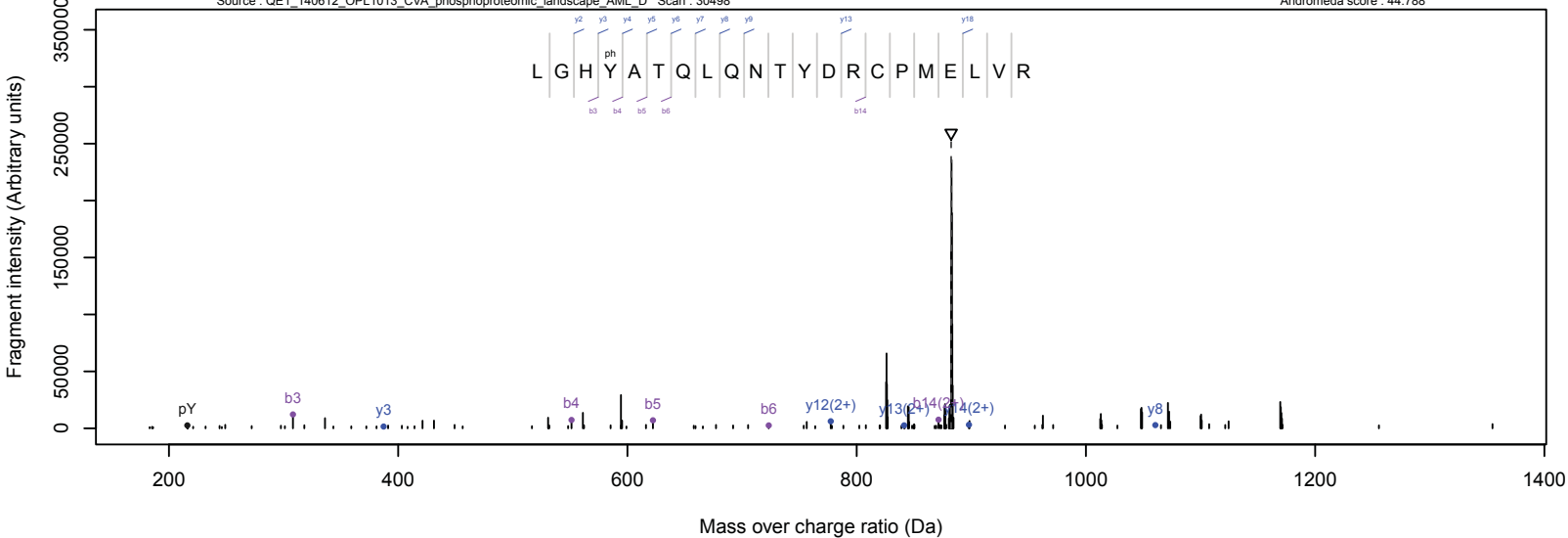
Andromeda score : 44.829



## LGHY(ph)ATQLQNTYDRCPMELVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 30498

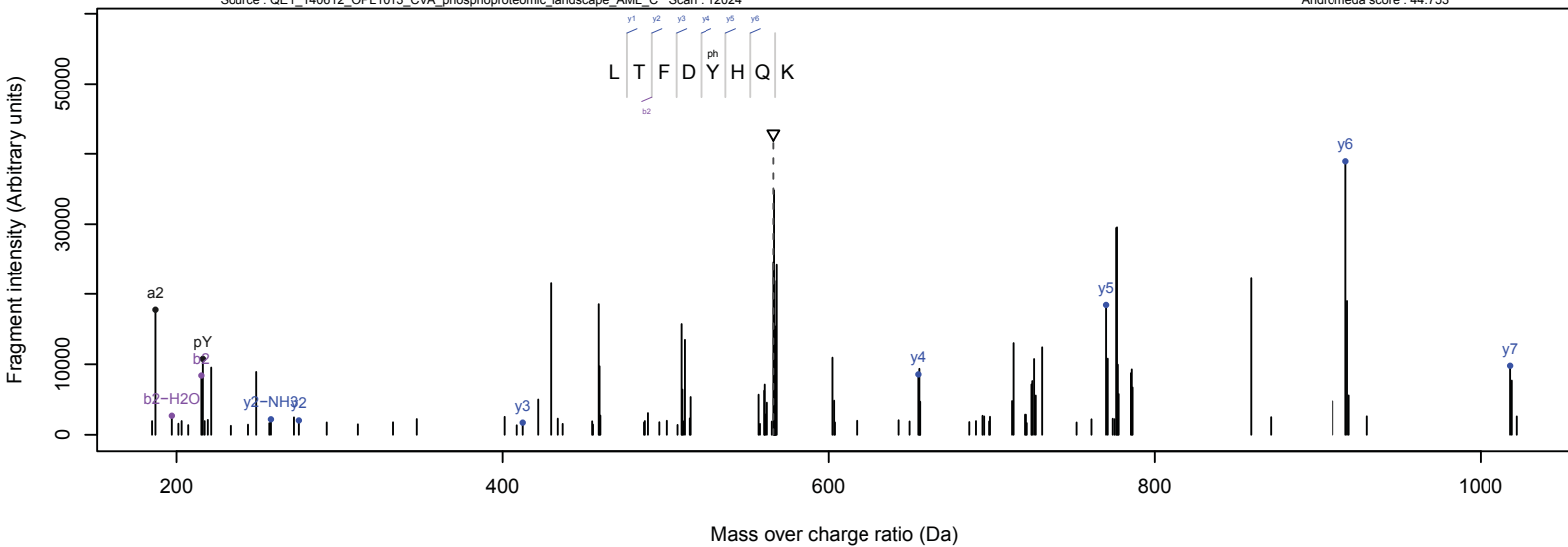
Andromeda score : 44.788



## LTFDY(ph)HQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12024

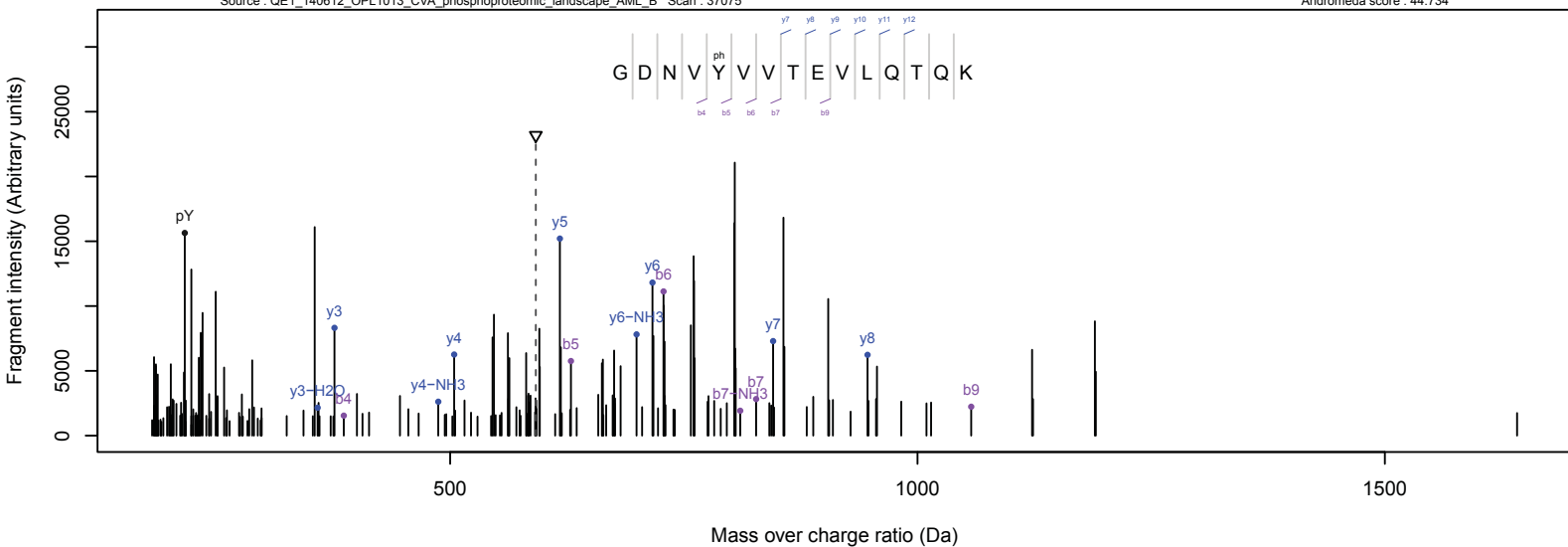
Andromeda score : 44.753



## GDNVY(ph)VVTEVLQTQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37075

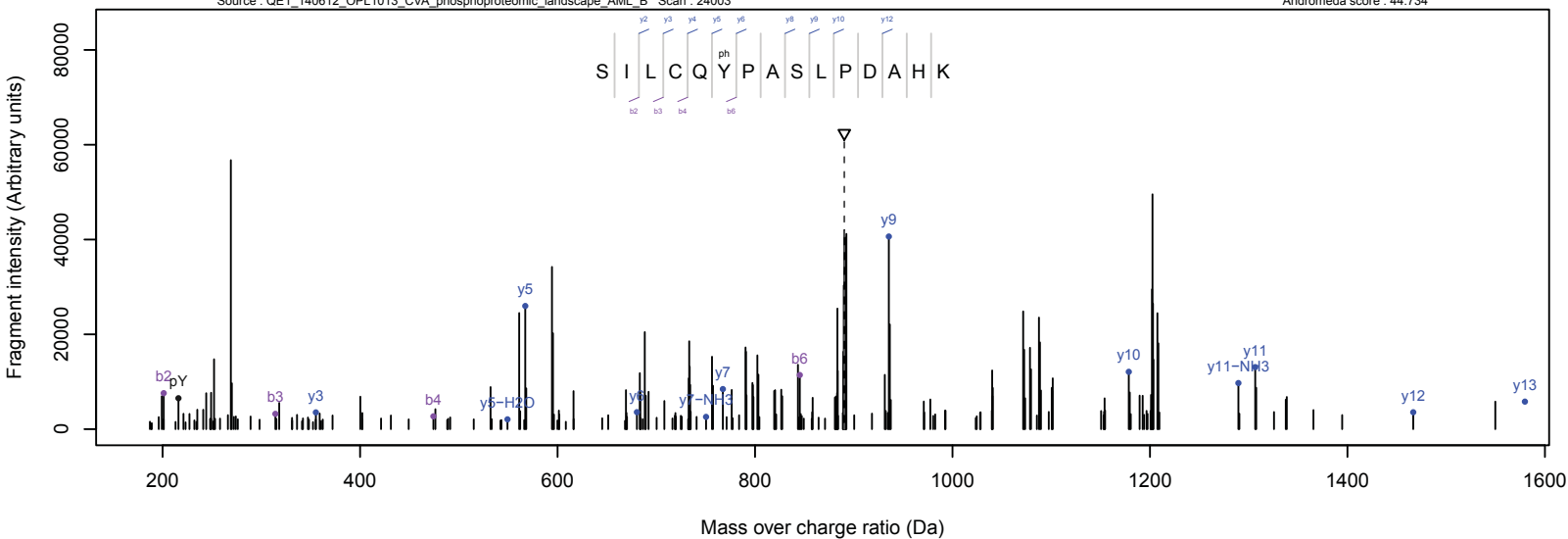
Andromeda score : 44.734



## SILCQY(ph)PASPDAHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24003

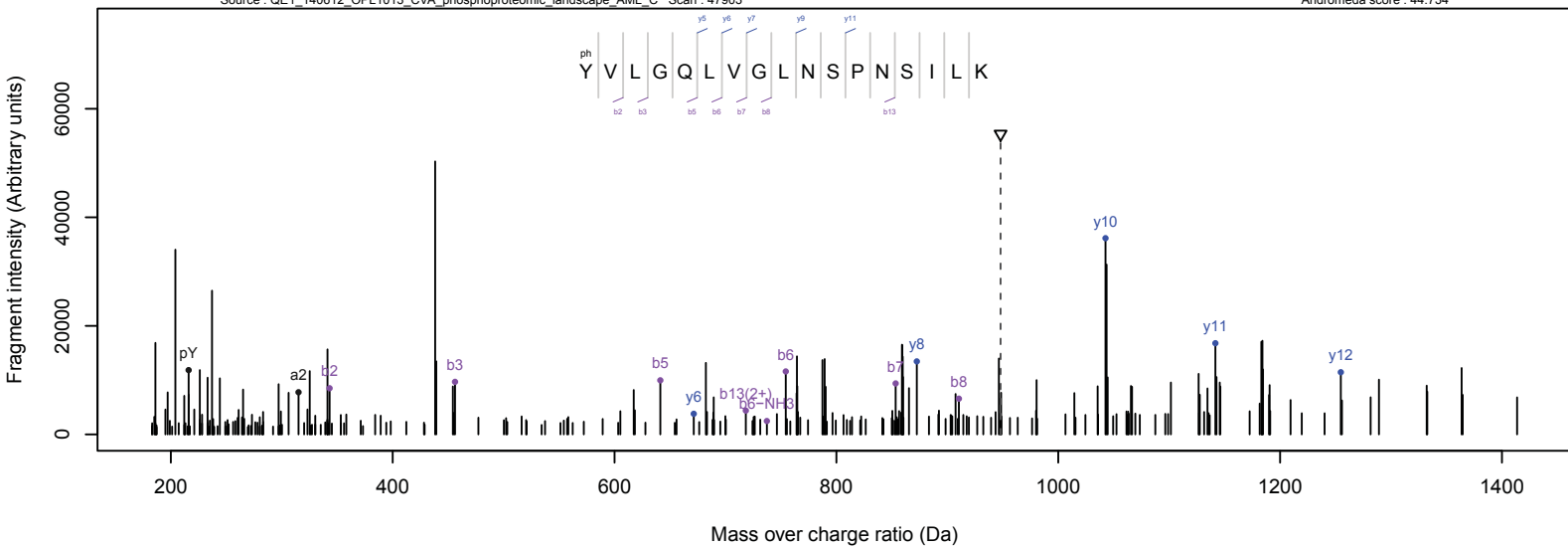
Andromeda score : 44.734



## Y(ph)VLGQLVGLNSPNSILK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47903

Andromeda score : 44.734

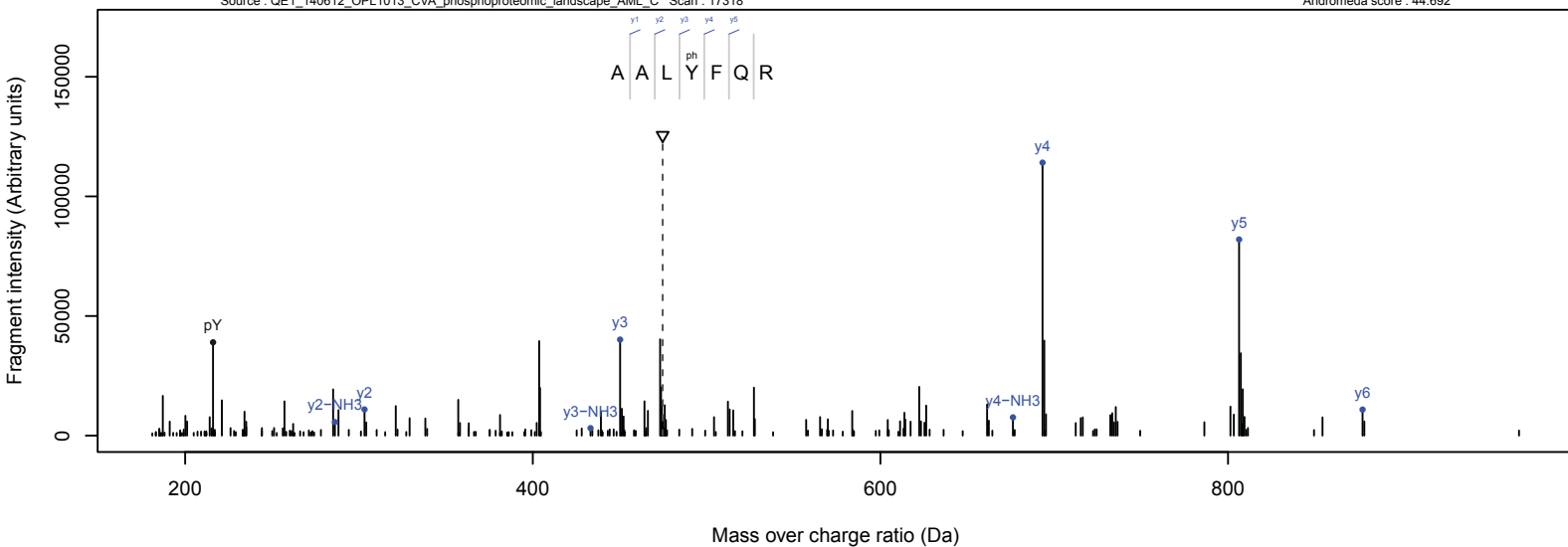




## AALY(ph)FQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17318

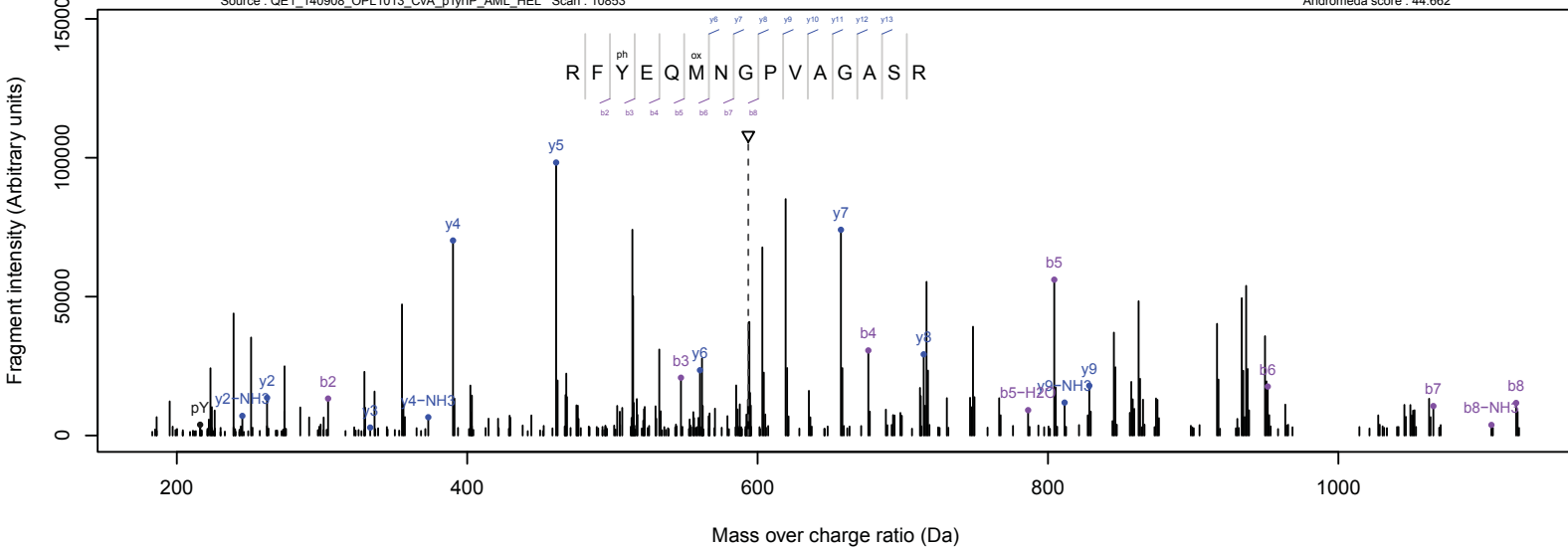
Andromeda score : 44.692



## RFY(ph)EQM(ox)NGPVAGASR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 10853

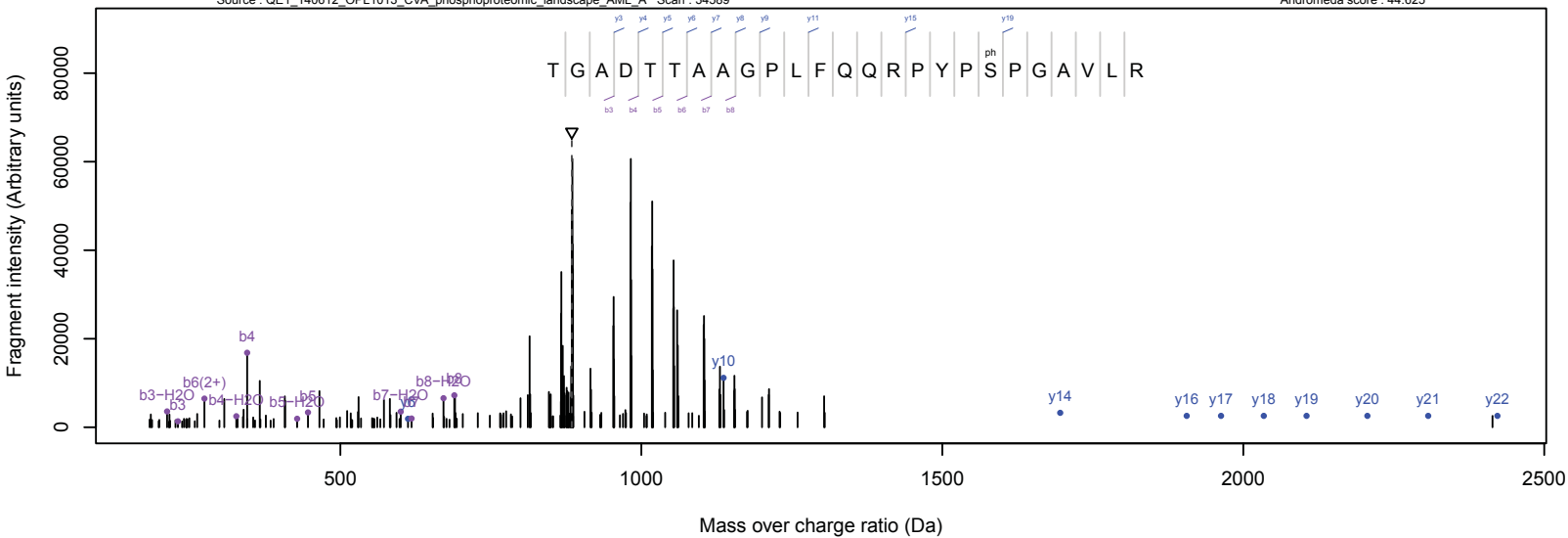
Andromeda score : 44.662



## TGADTTAAGPLFQQRYPSPGAVLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 34589

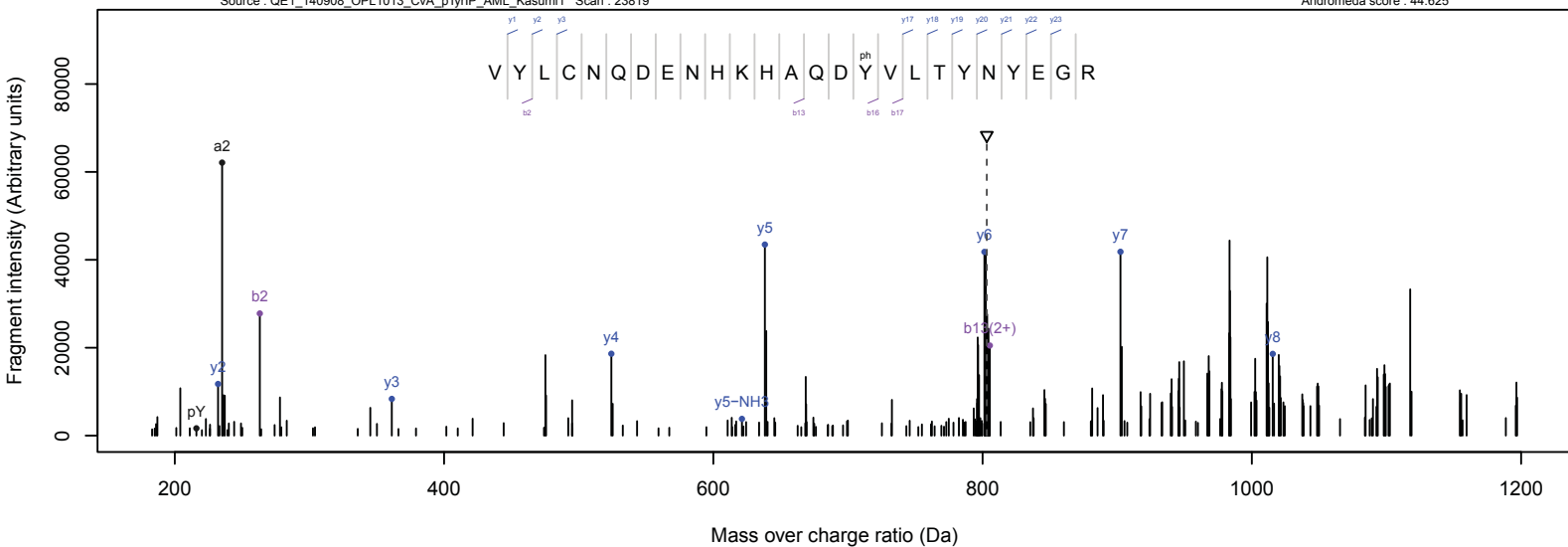
Andromeda score : 44.625



VYLCNQDENHKHAQDY(ph)VLTNYEGR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 23819

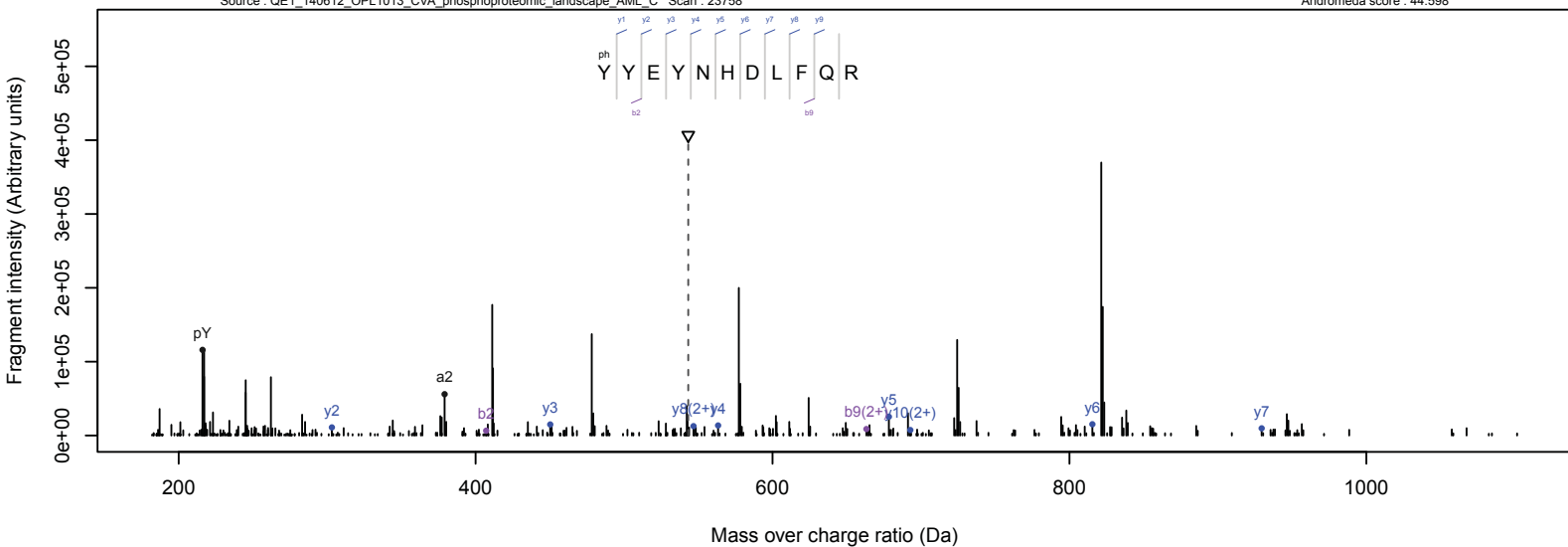
Andromeda score : 44.625



Y(ph)YEYNHDLFQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23758

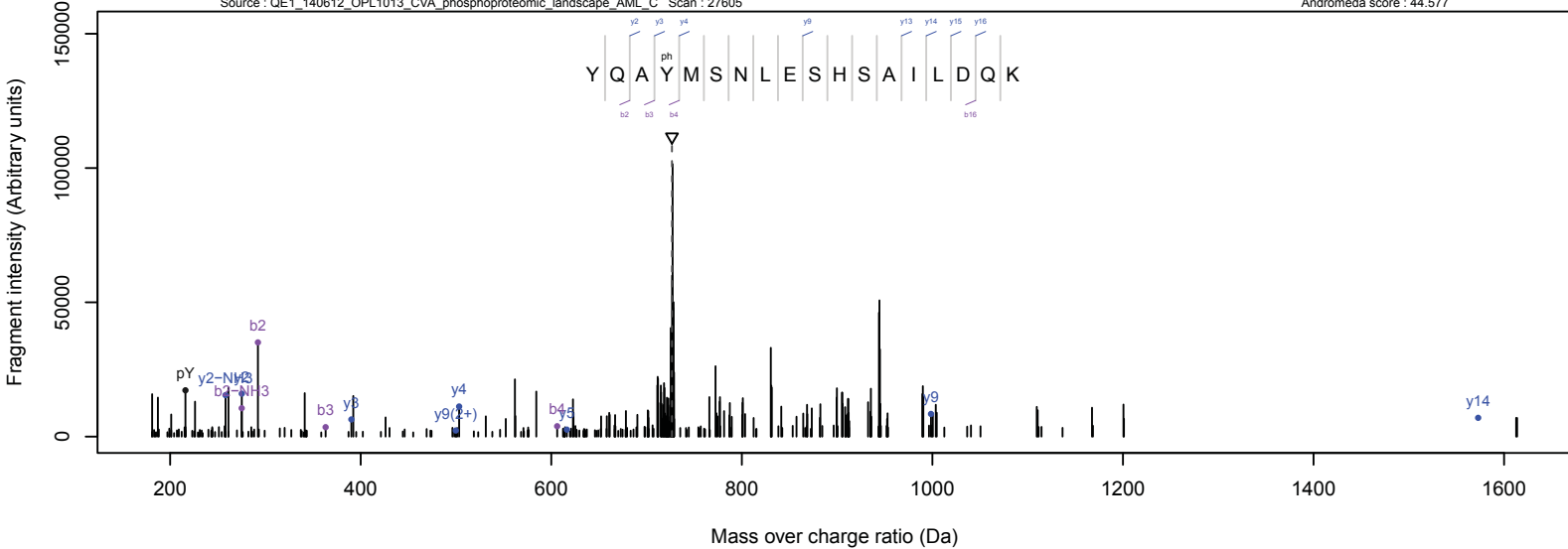
Andromeda score : 44.598



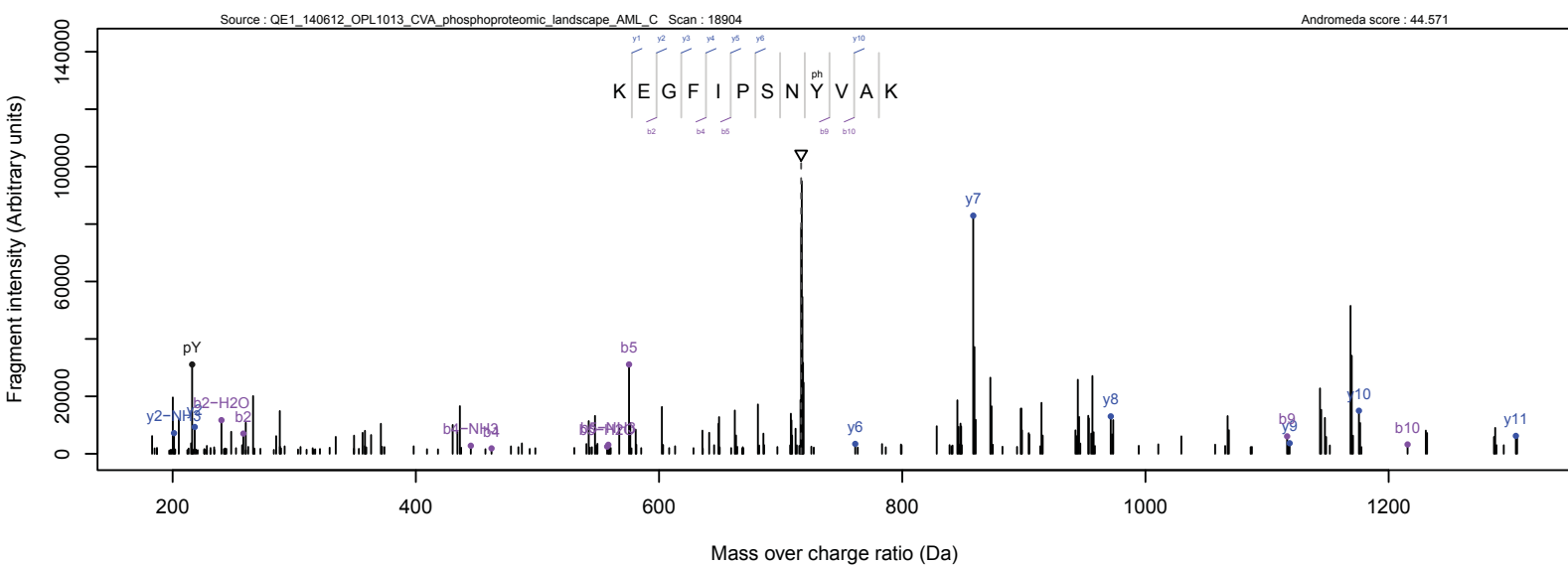
YQAY(ph)MSNLESHSAILDQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27605

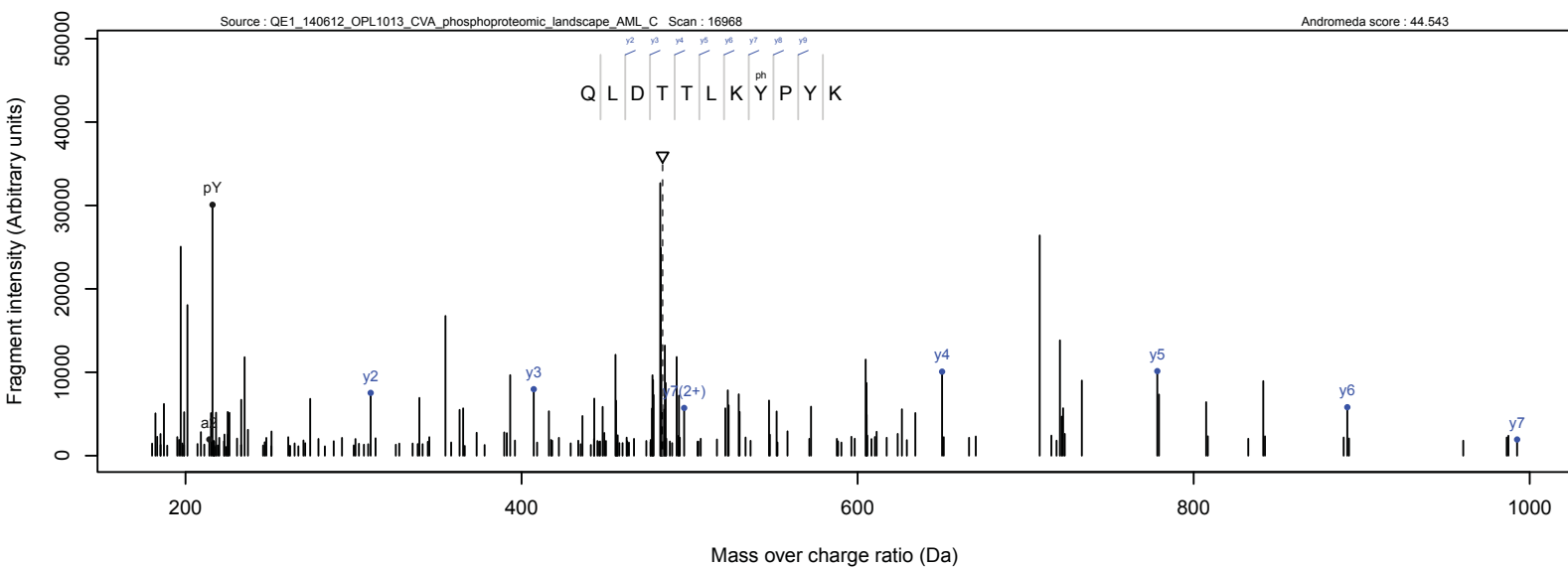
Andromeda score : 44.577



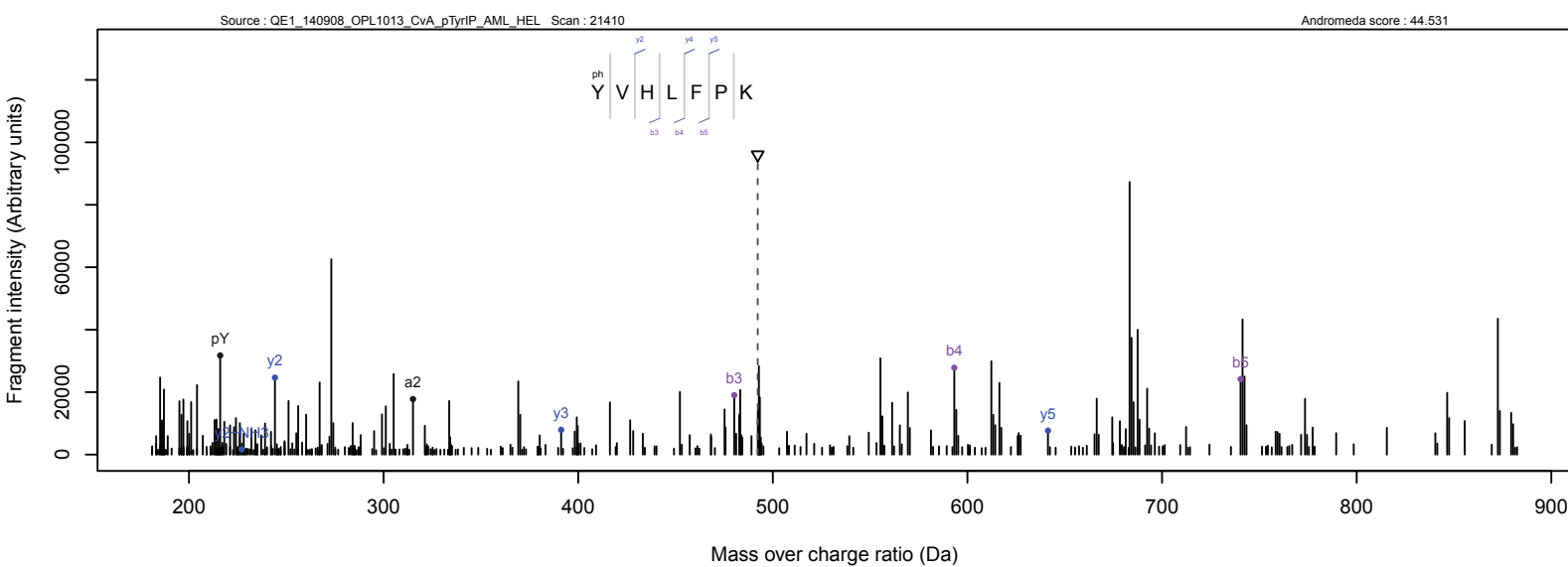
## KEGFIPSNY(ph)VAK-2+



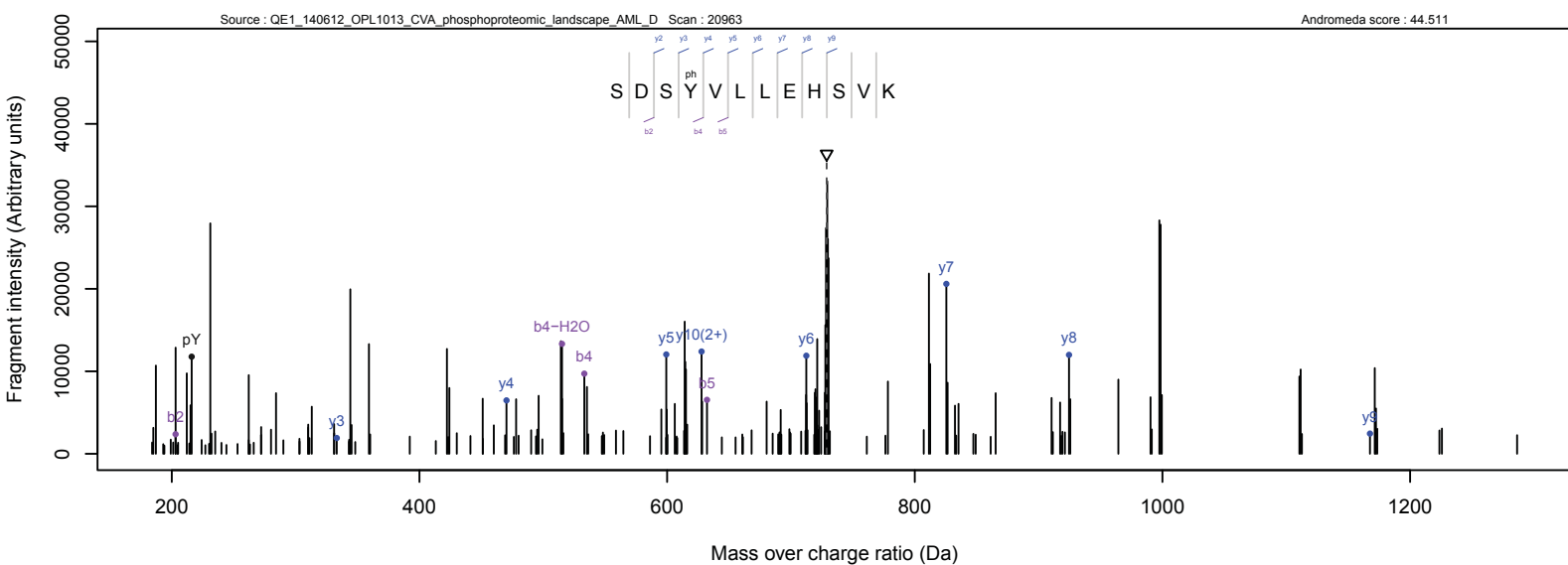
## QLDRTLKY(ph)PYK-3+



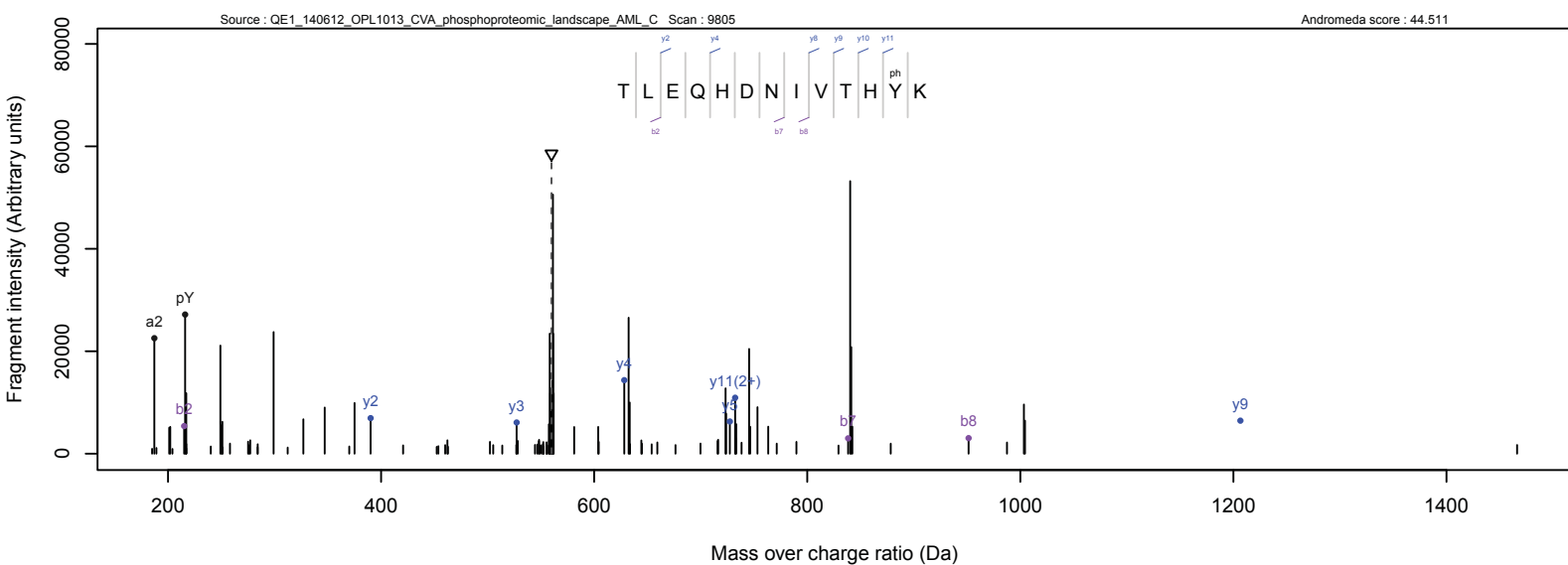
## Y(ph)VHLFPK-2+



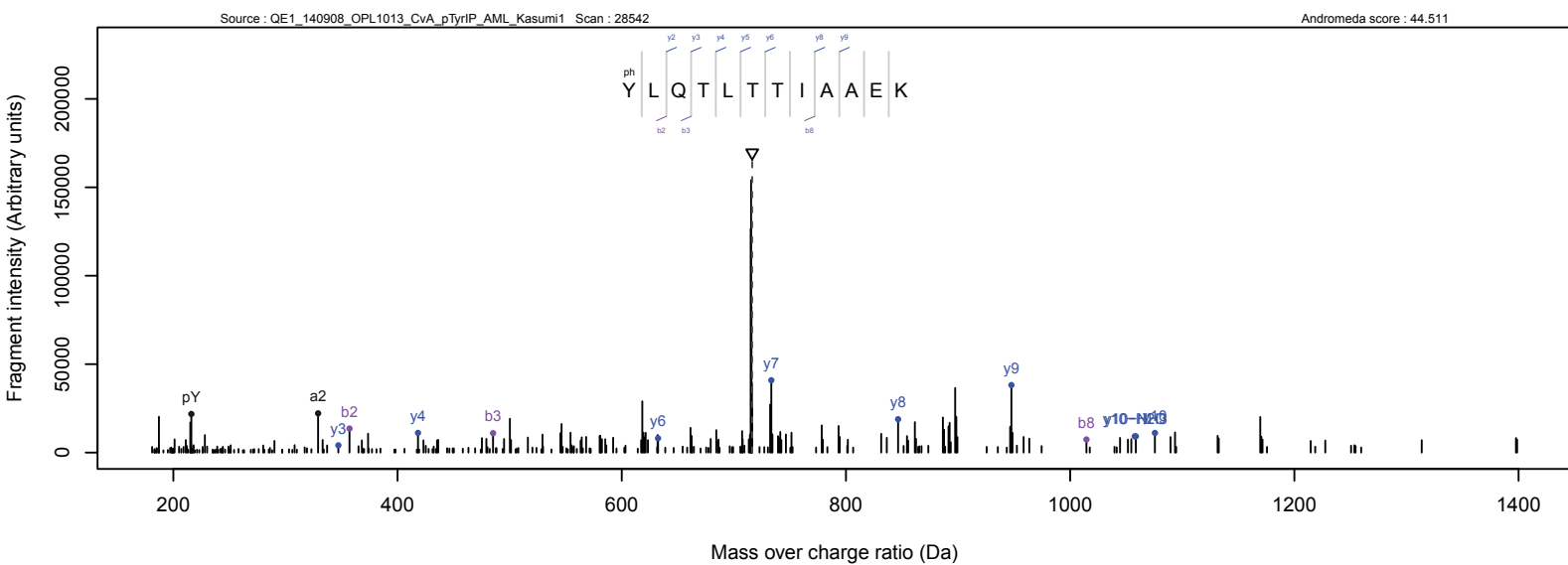
## SDSY(ph)VLLHSVK-2+



## TLEQHDNIVTHYK-3+



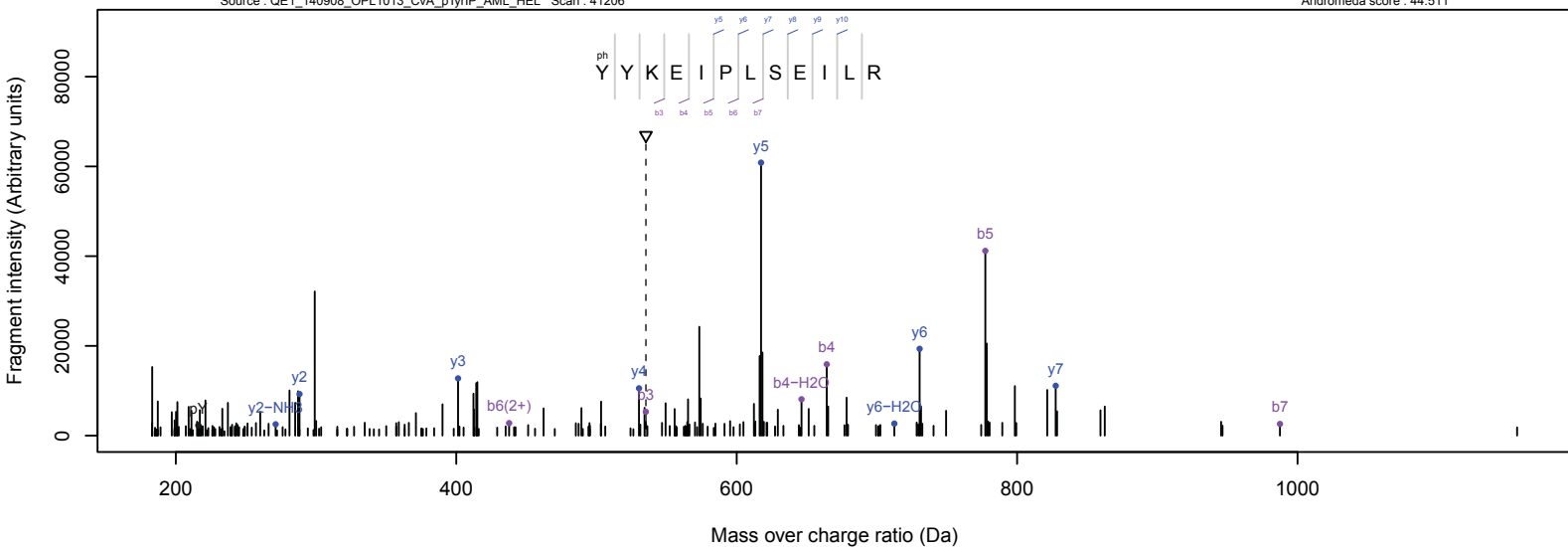
## Y(ph)LQTLTIAAEK-2+



## Y(ph)YKEIPLSEILR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 41206

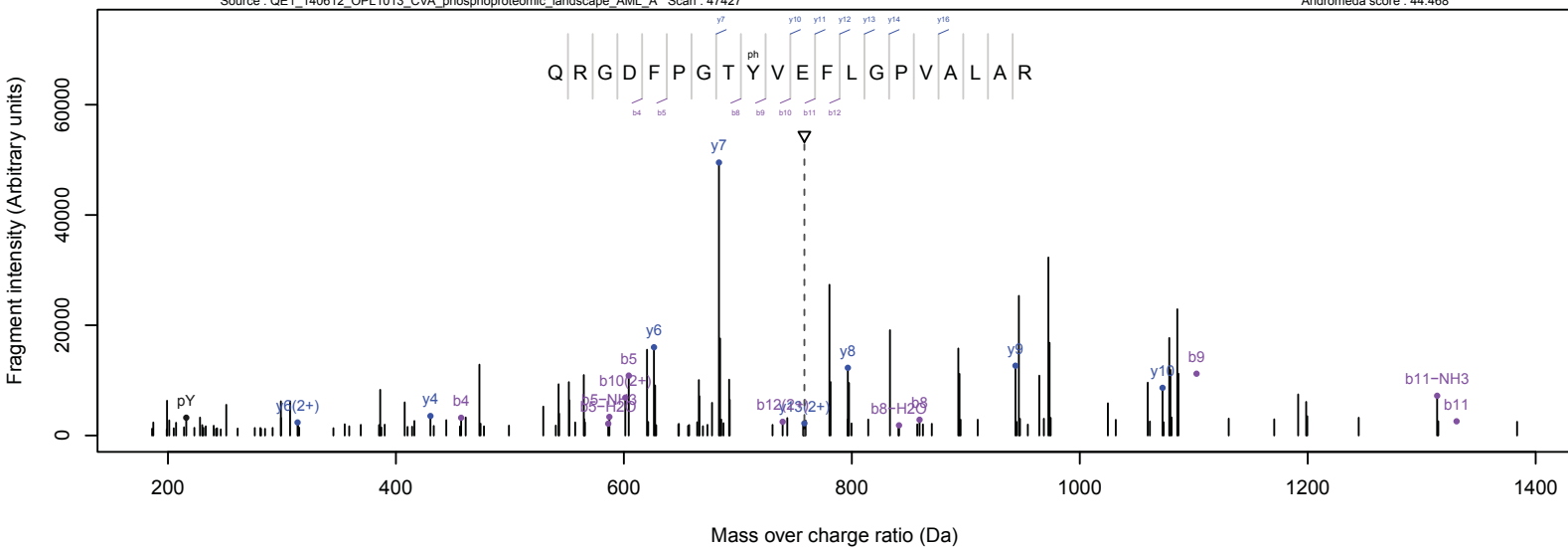
Andromeda score : 44.511



## QRGDFPGTY(ph)VEFLGPVALAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 47427

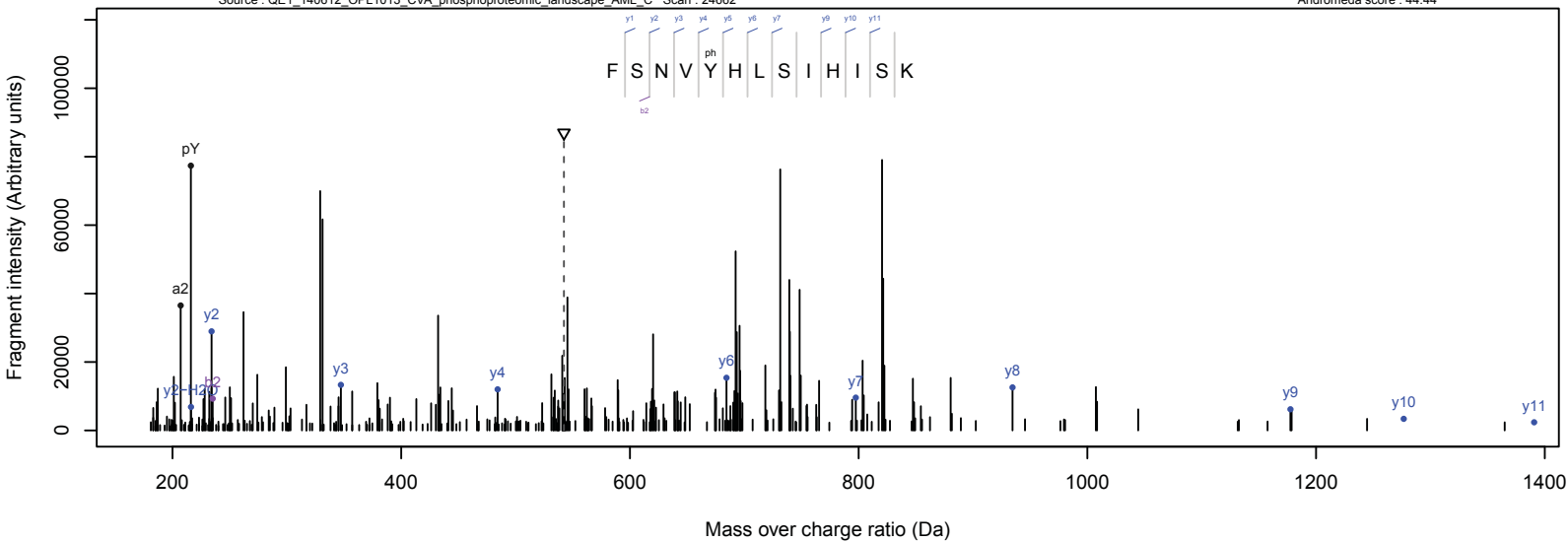
Andromeda score : 44.468



## FSNVY(ph)HLSIHISK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24662

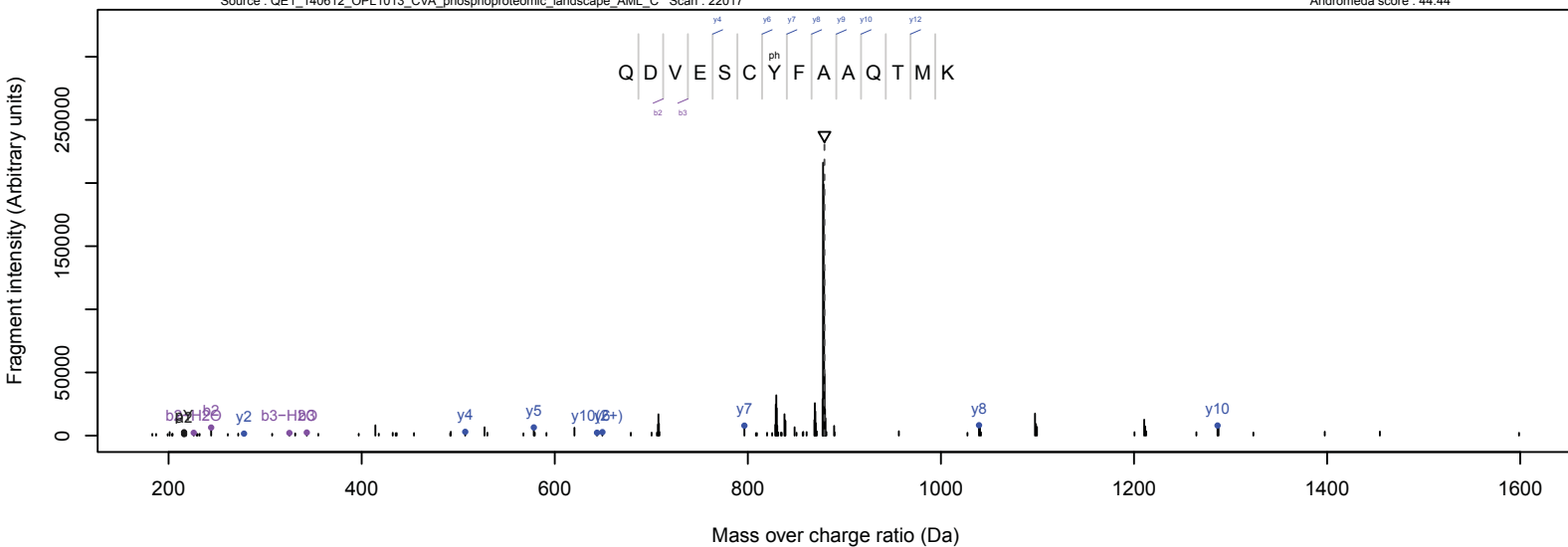
Andromeda score : 44.44



## QDVESCY(ph)FAAQTMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22017

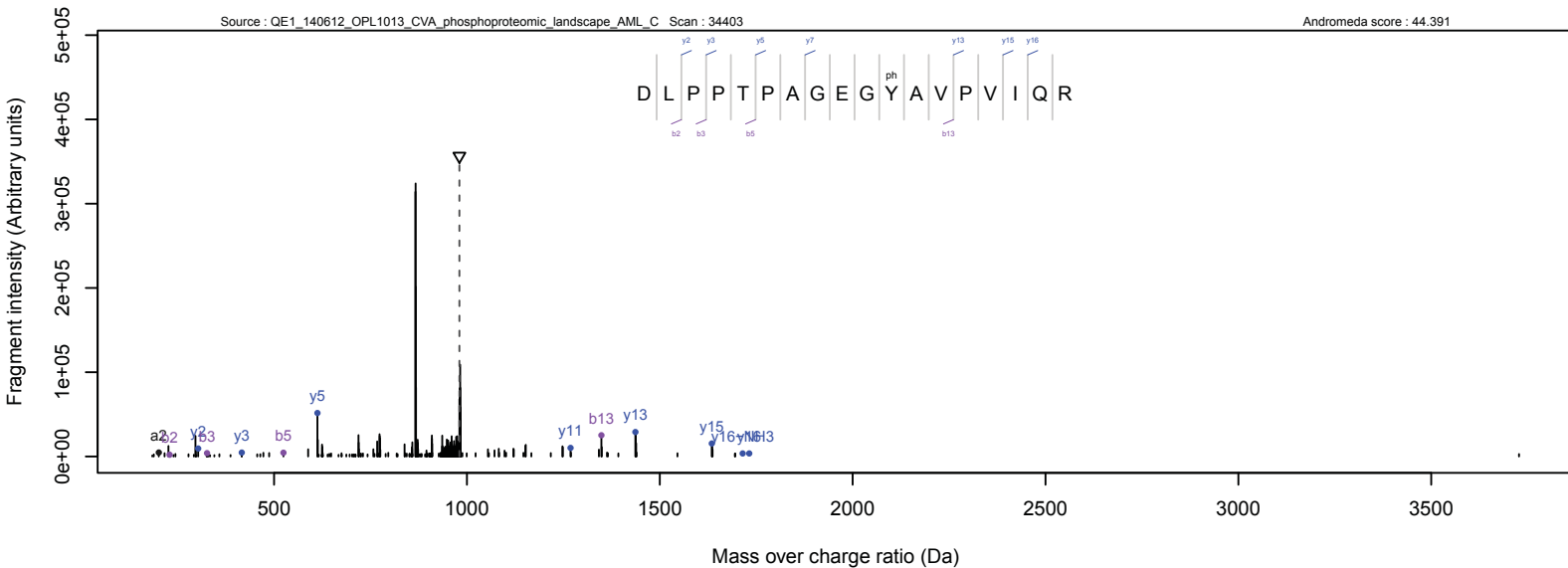
Andromeda score : 44.44



## DLPPTPAGEGY(ph)AVPVIQR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 34403

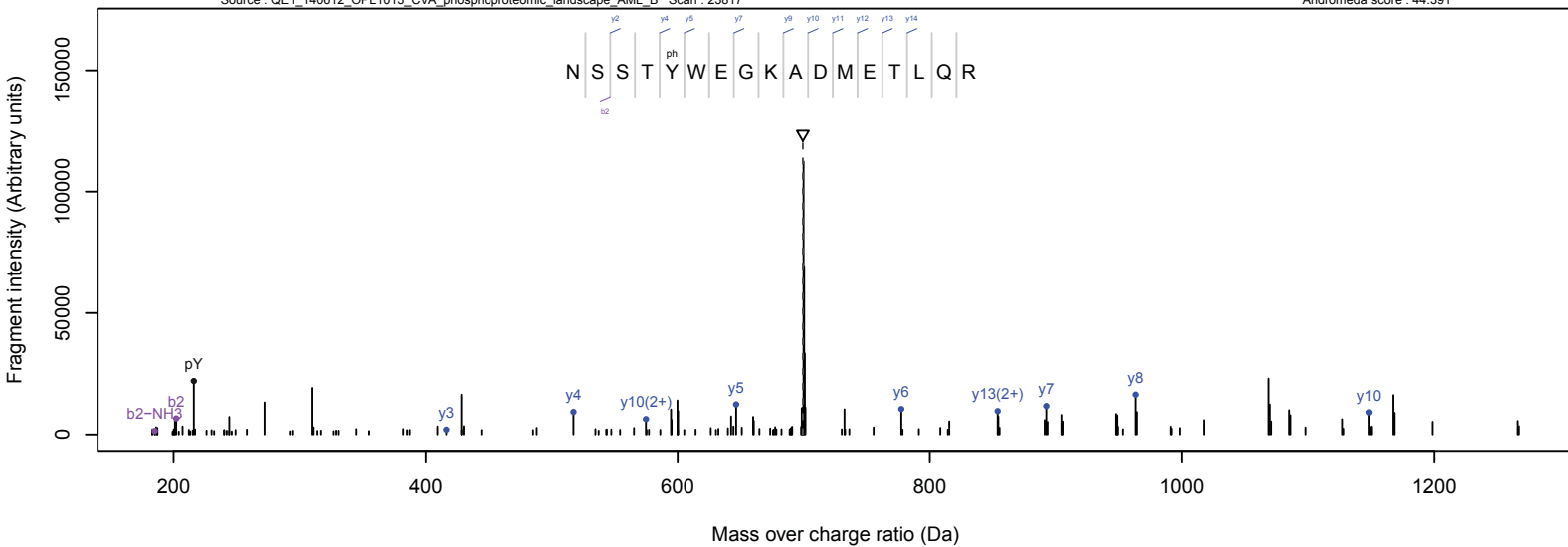
Andromeda score : 44.391



## NSSTY(ph)WEGKADMETLQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 23817

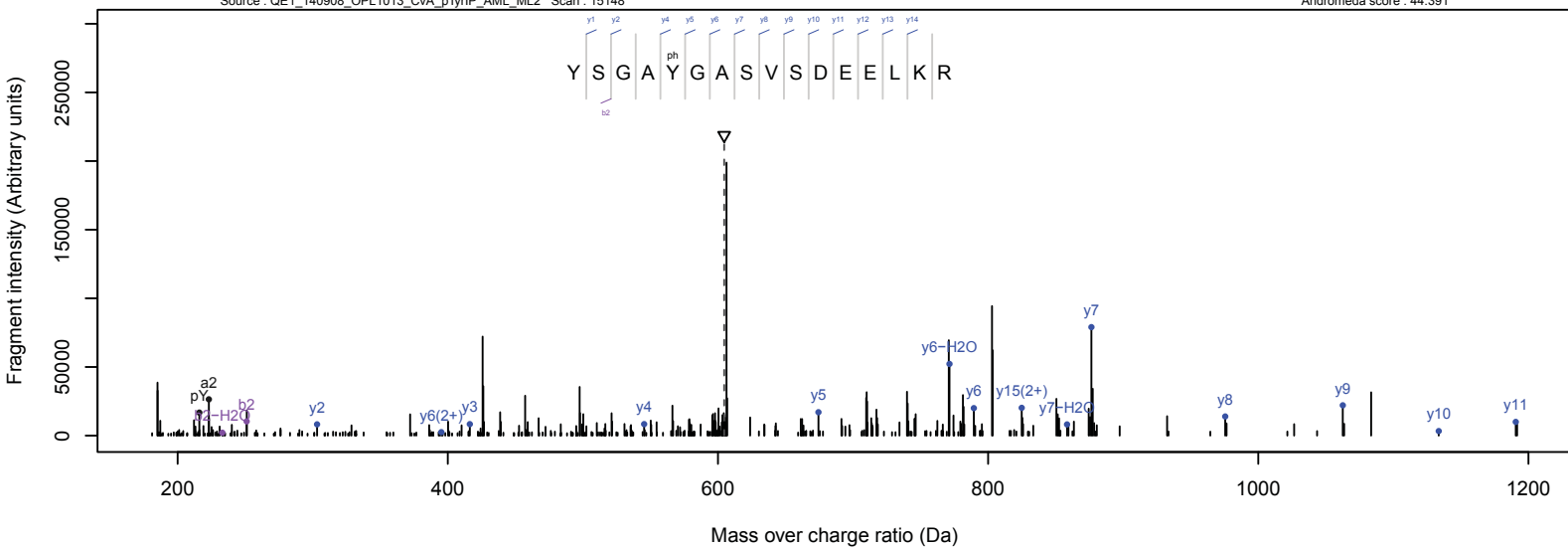
Andromeda score : 44.391



YSGAY(ph)GASVSDEELKR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 15148

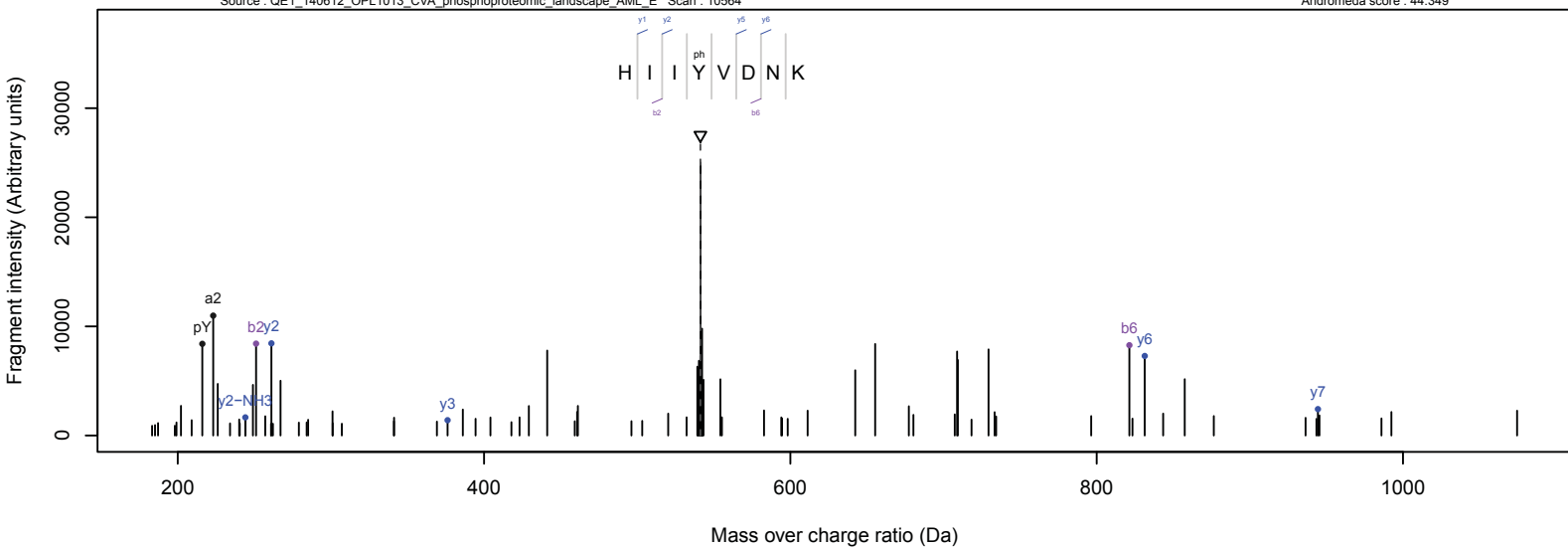
Andromeda score : 44.391



HIYY(ph)VDNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 10564

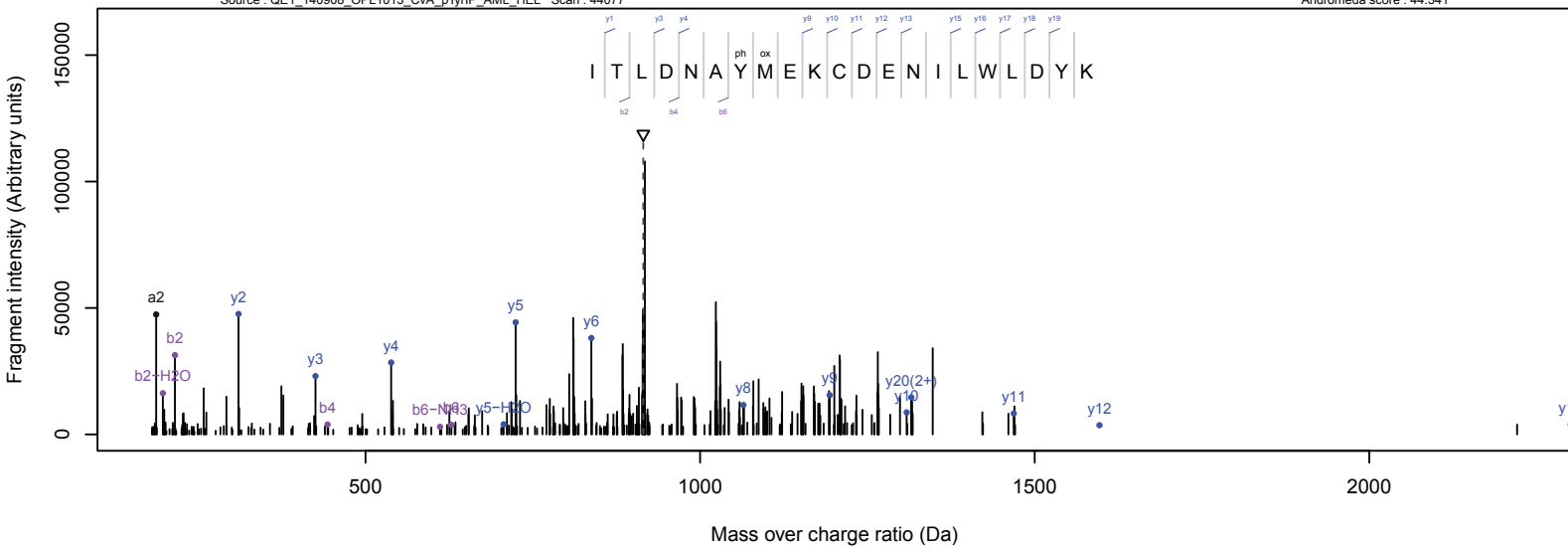
Andromeda score : 44.349



ITLDNAY(ph)M(ox)EKCDENILWLDYK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 44077

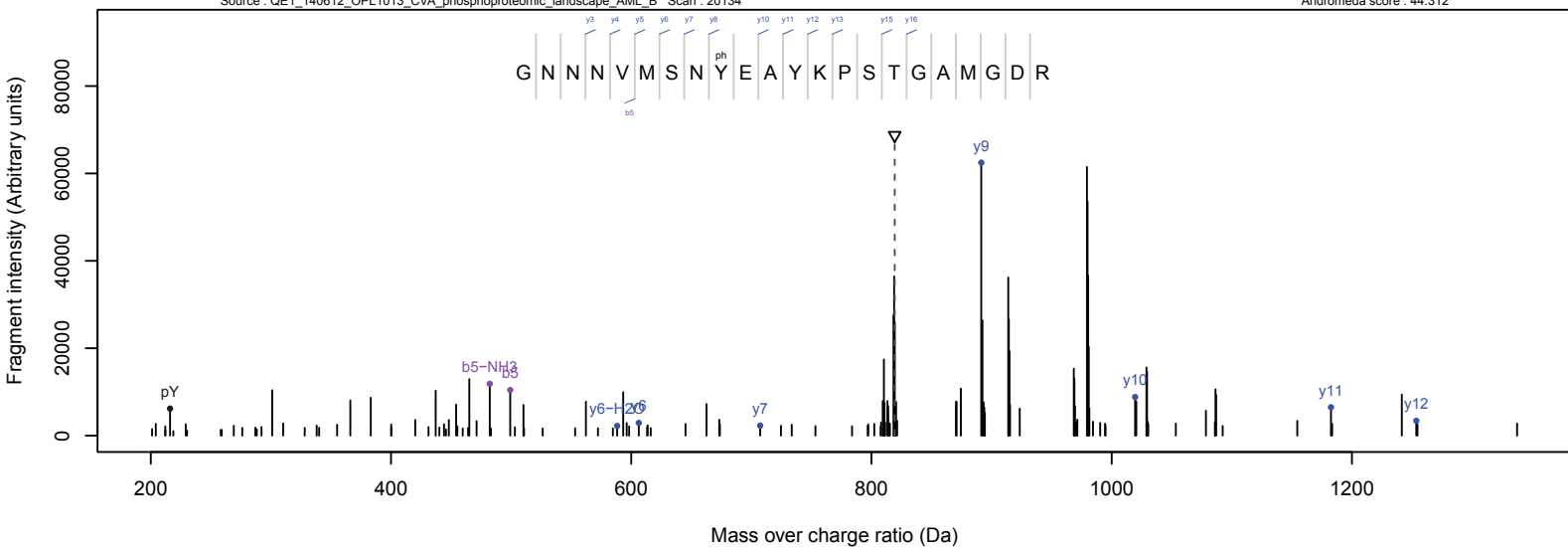
Andromeda score : 44.341



**GNNVMSNY(ph)EAYKPSTGAMGDR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 20134

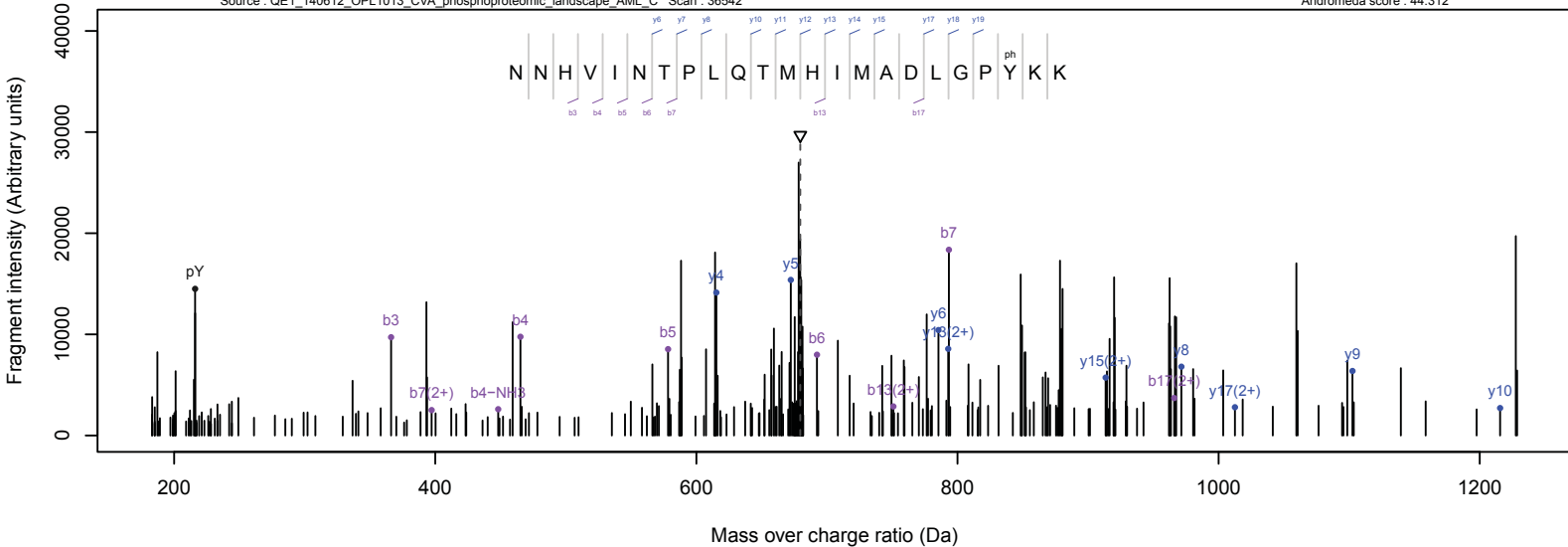
Andromeda score : 44.312



**NNHVINTPLQTMHIMADLGPYK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36542

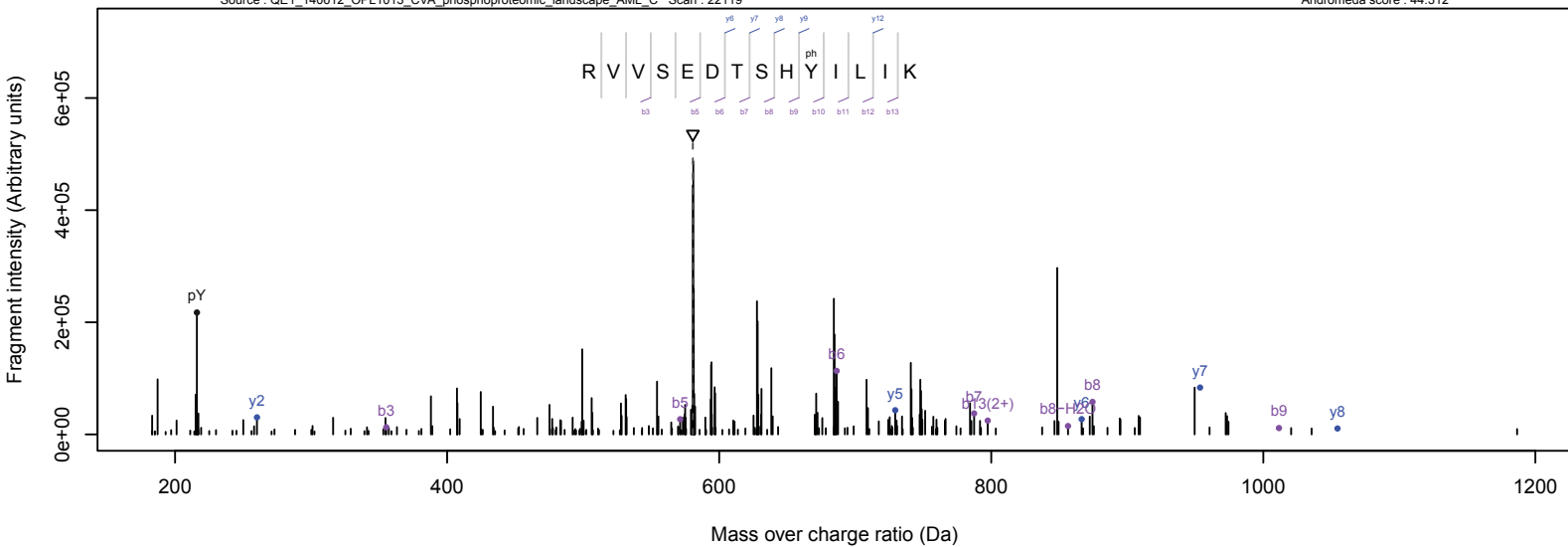
Andromeda score : 44.312



**RVVSEDTSHYILIK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22119

Andromeda score : 44.312

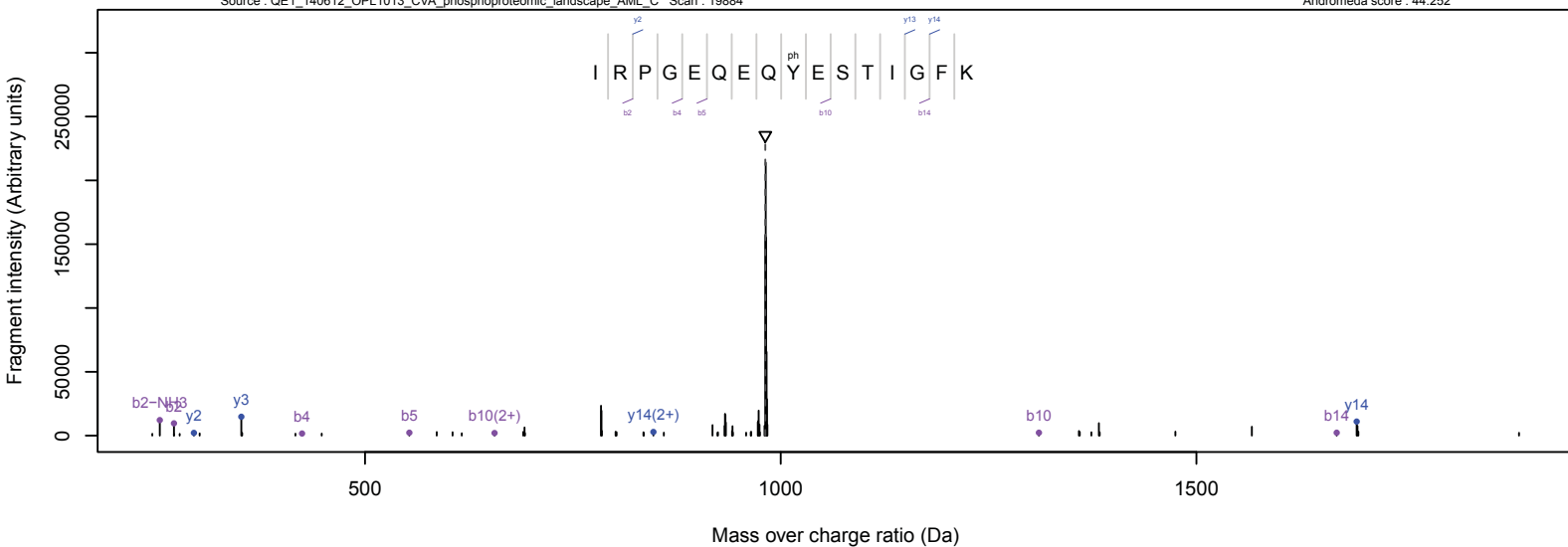




## IRPGEQEQY(ph)ESTIGFK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19884

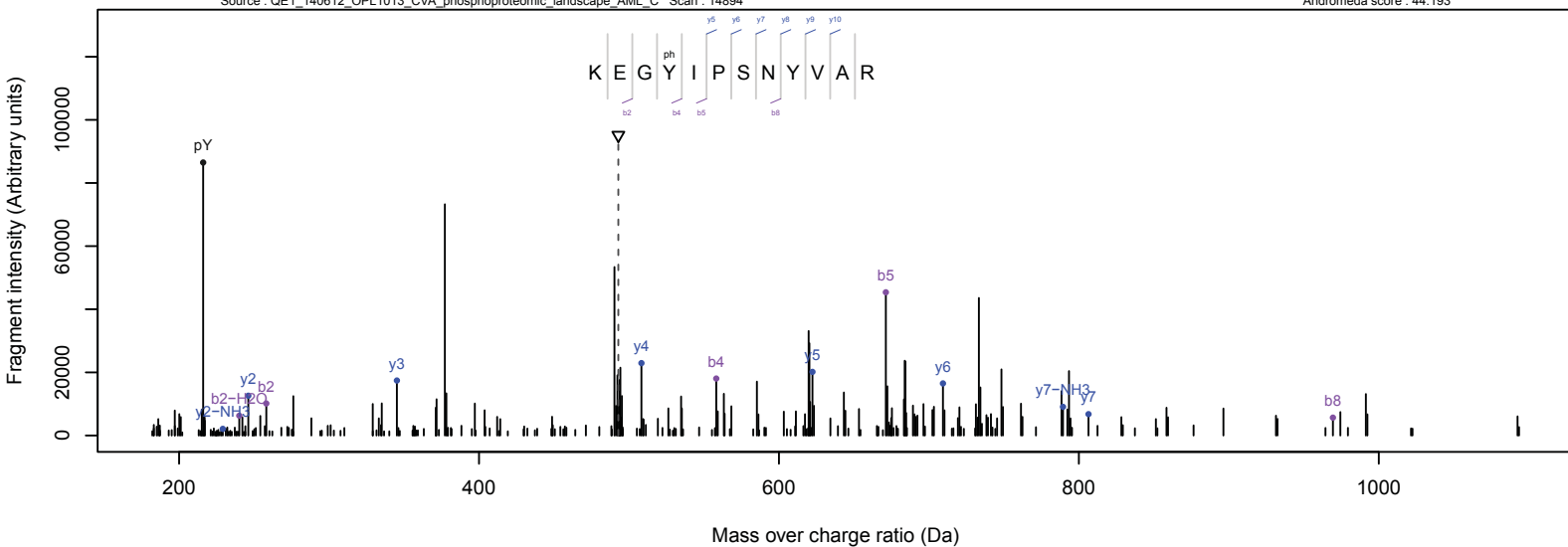
Andromeda score : 44.252



## KEGY(ph)IPSNYVAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14894

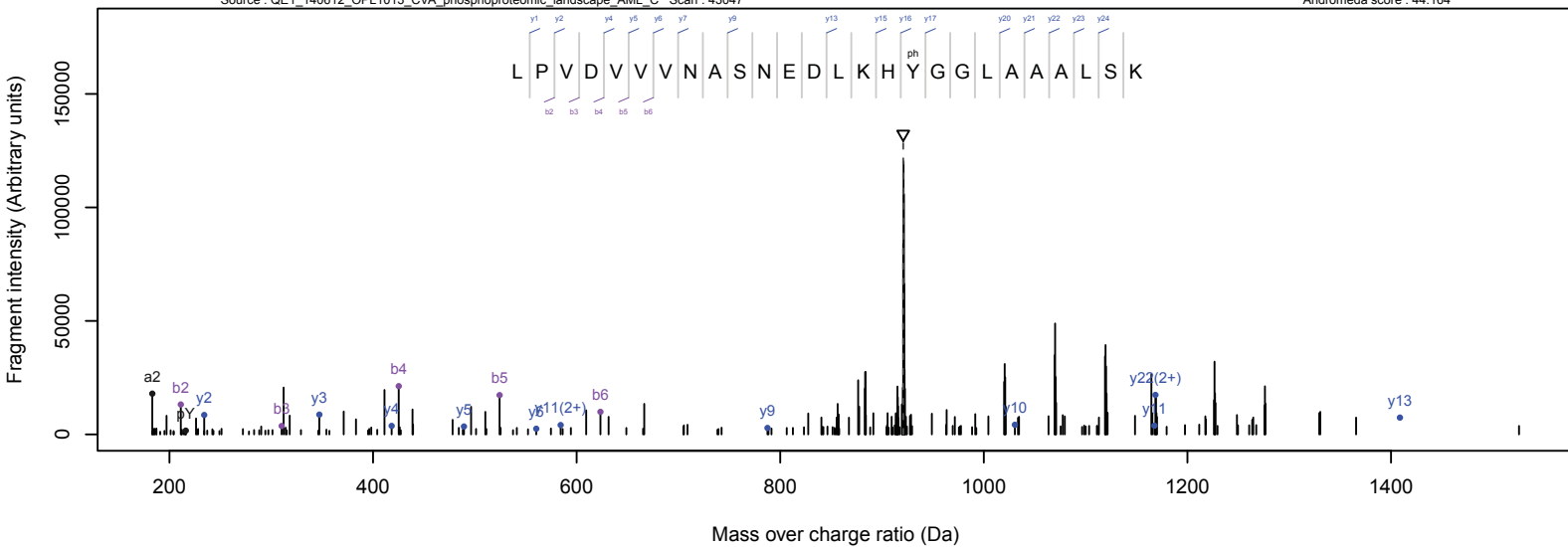
Andromeda score : 44.193



## LPVDVVNASNEDLKHY(ph)GGLAALSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 43047

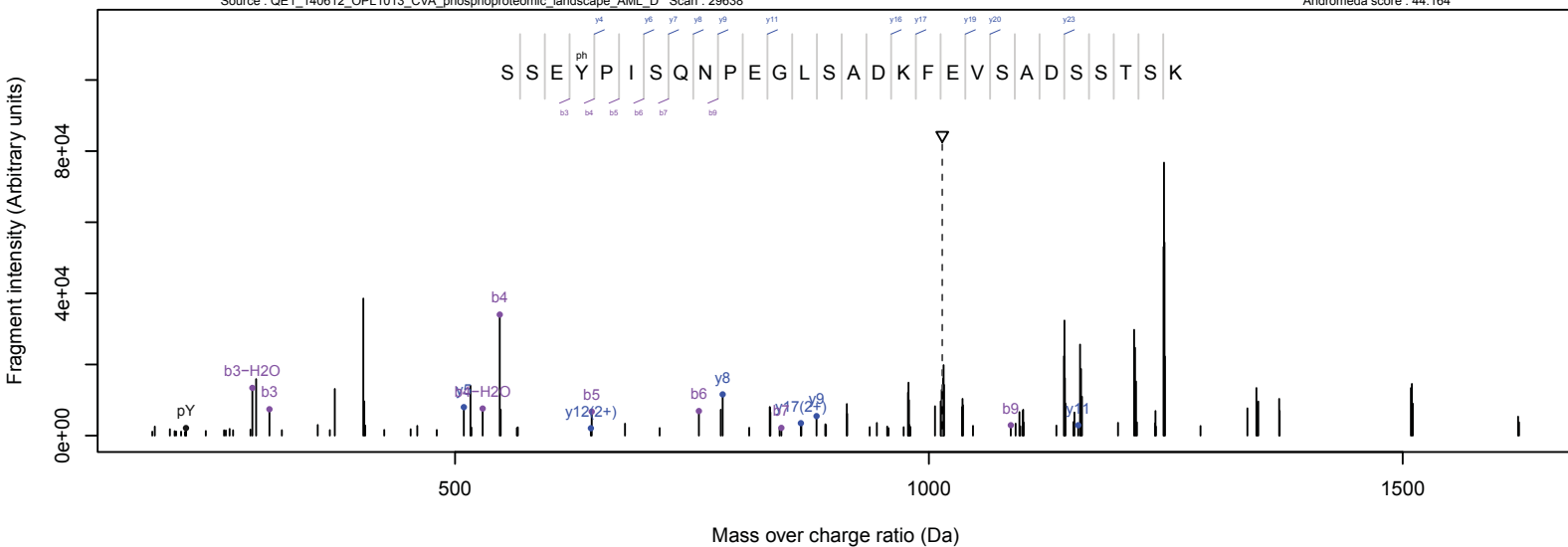
Andromeda score : 44.164



## SSEY(ph)PISQNP EGLSADKFEV SADSSTSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 29638

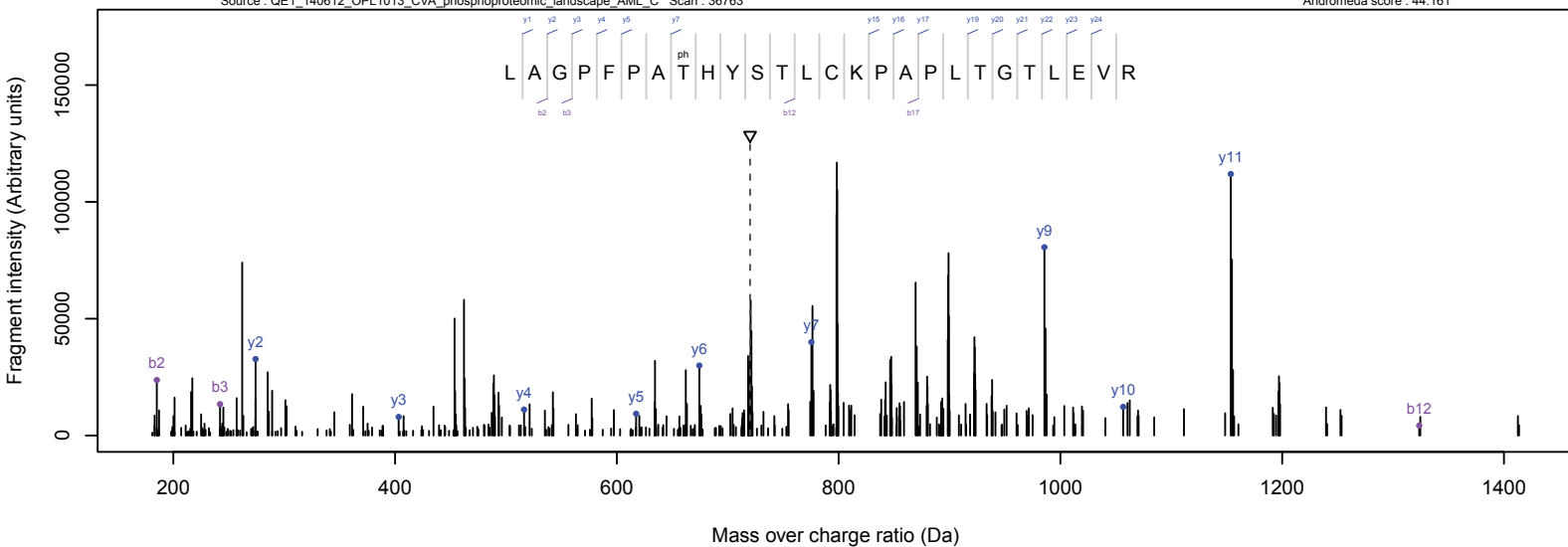
Andromeda score : 44.164



## LAGFPAT(ph)HYSTLCKPAPLTGTLEVR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36763

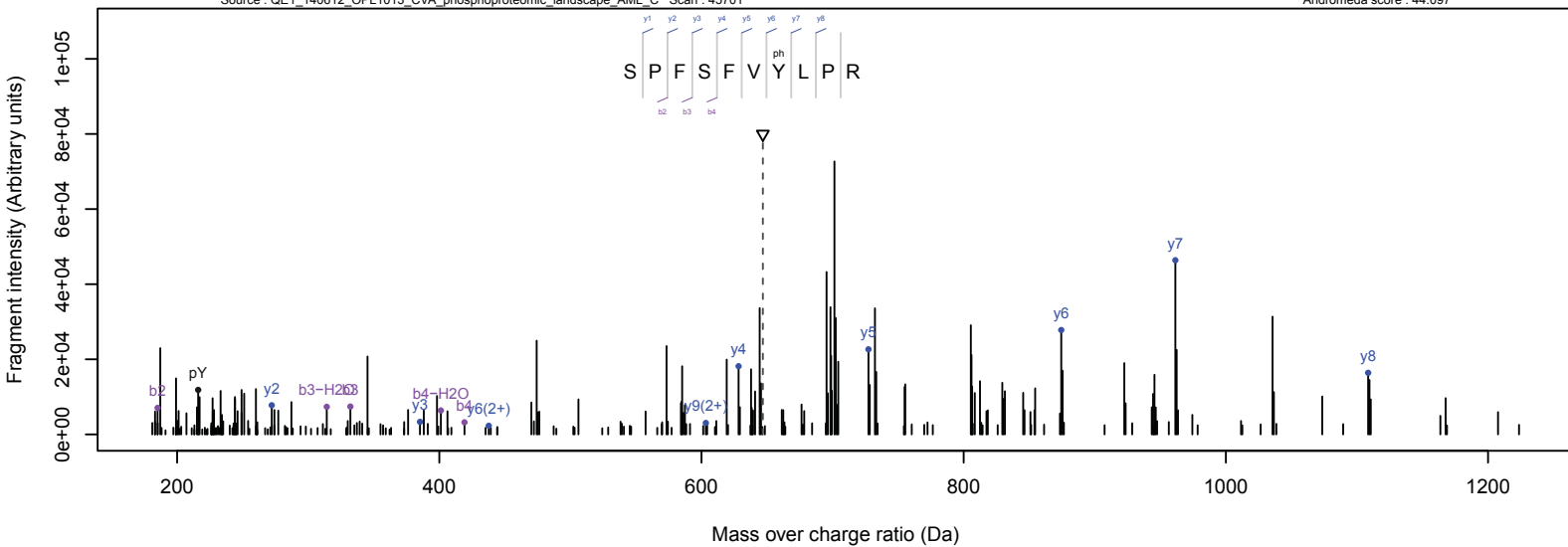
Andromeda score : 44.161



## SPFSFVY(ph)LPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45701

Andromeda score : 44.097

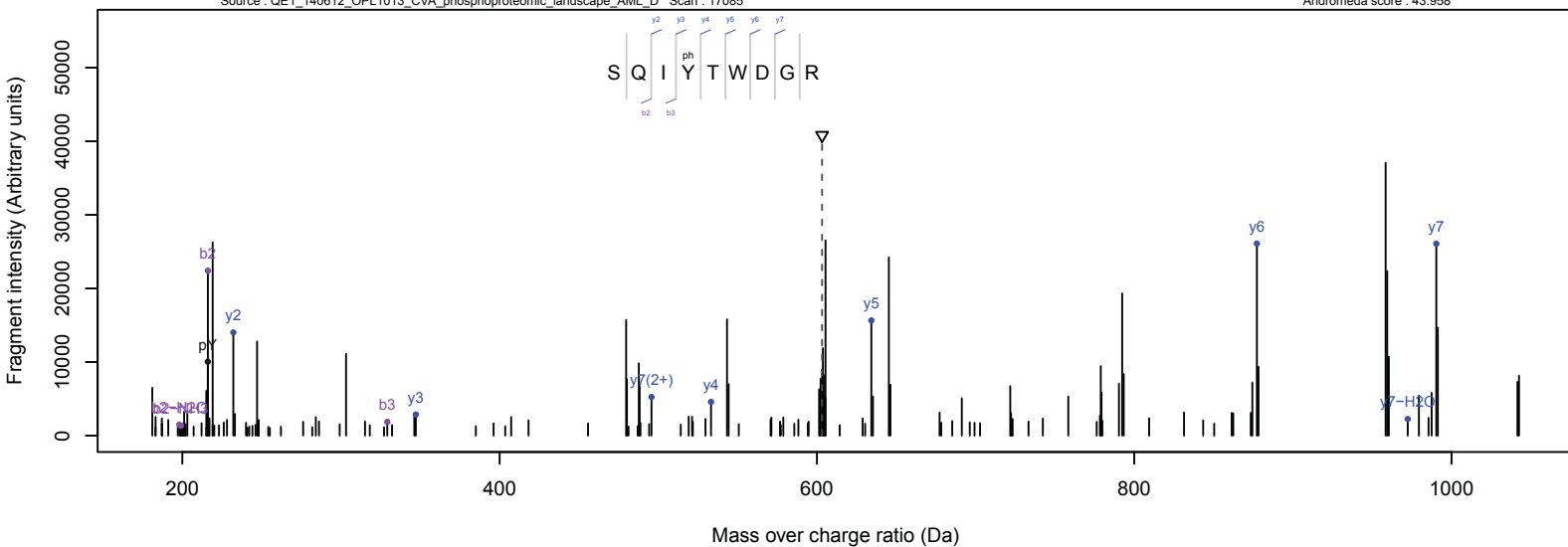




## SQIY(ph)TWDGR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 17085

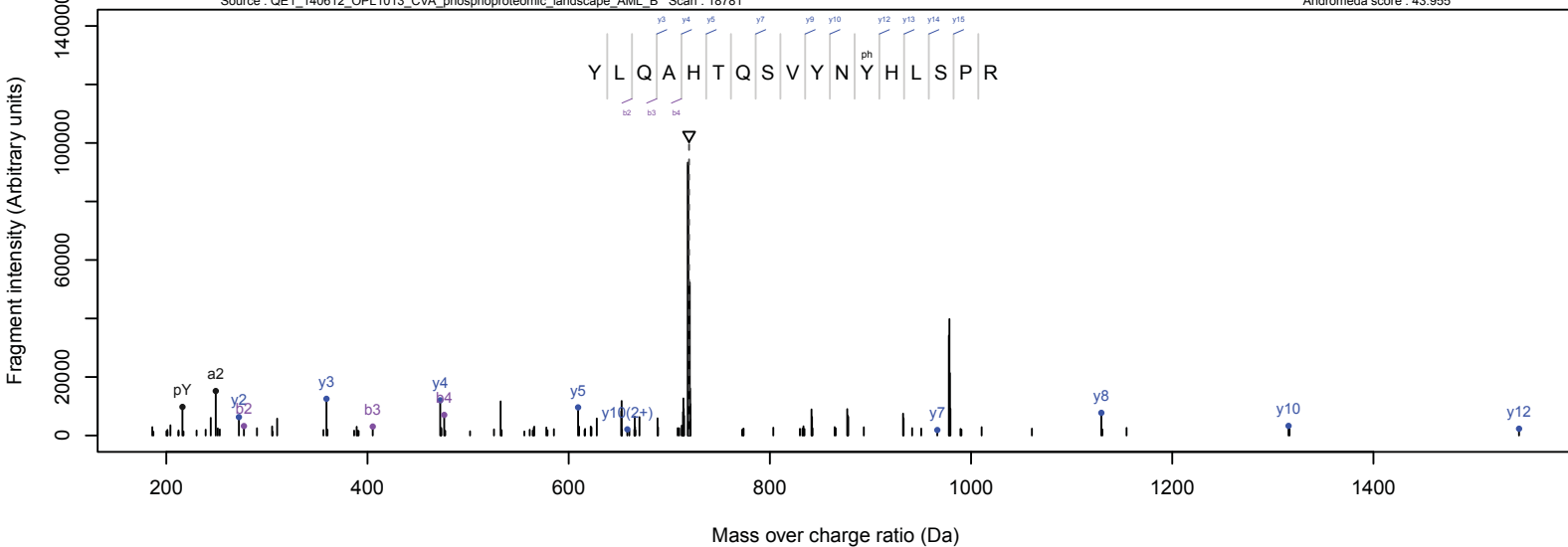
Andromeda score : 43.958



## YLQAHTQSVYNY(ph)HLSPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18781

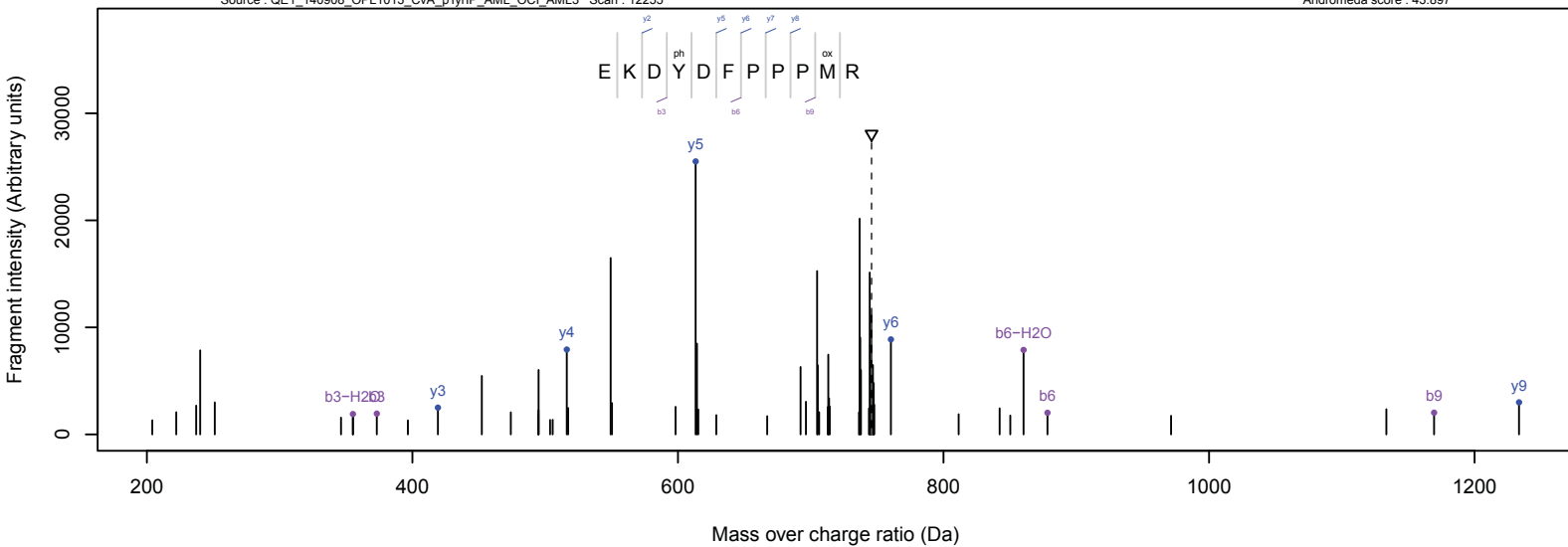
Andromeda score : 43.955



## EKDY(ph)DFPPM(ox)R-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 12255

Andromeda score : 43.897

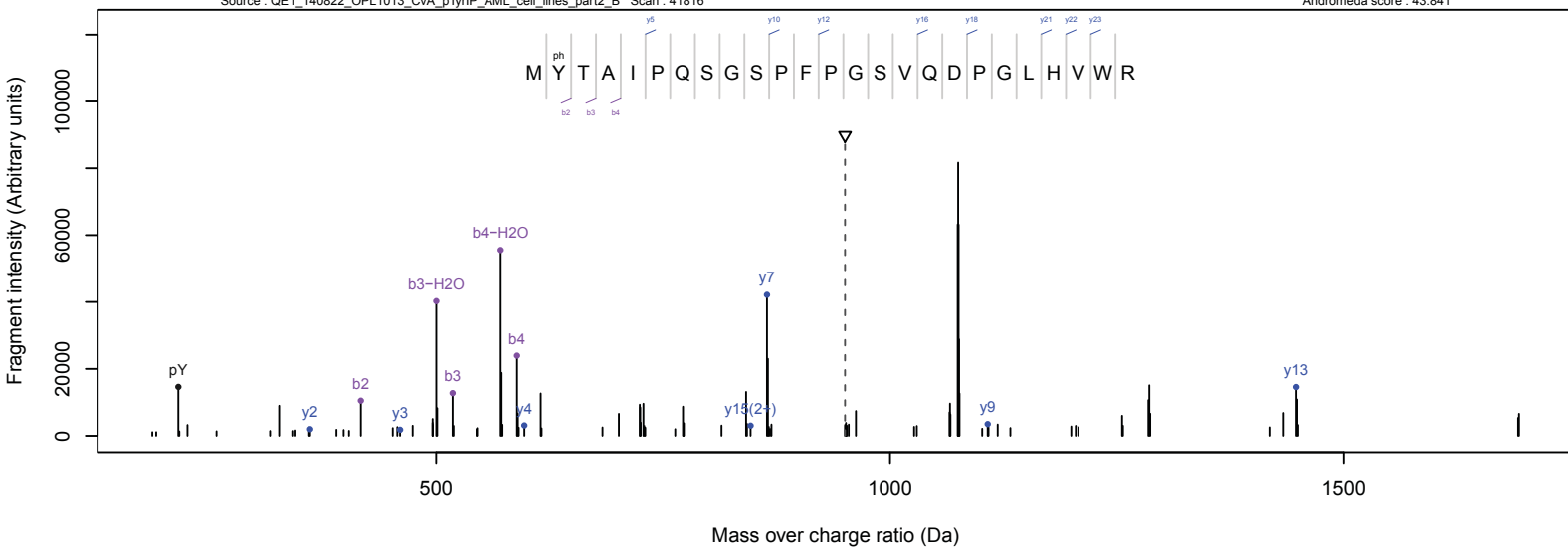




## (ac)MY(ph)TAIPQSGSPFPGSVQDPLHVWR-3+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 41816

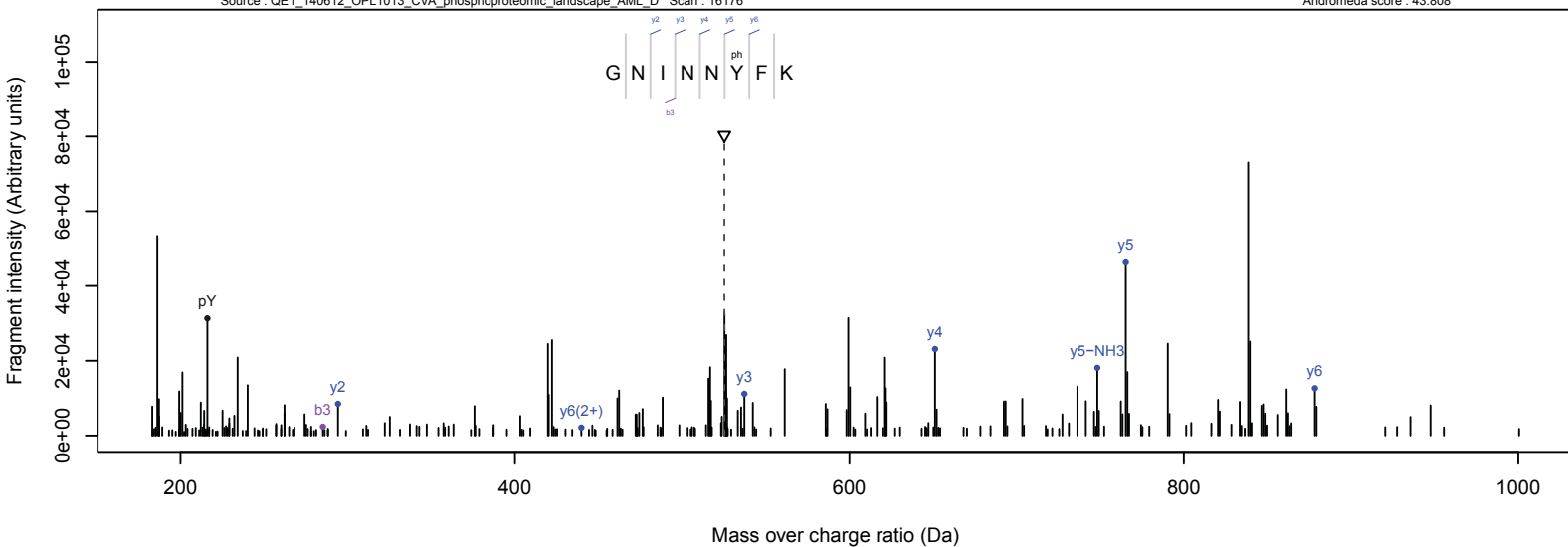
Andromeda score : 43.841



## GNINNY(ph)FK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 16176

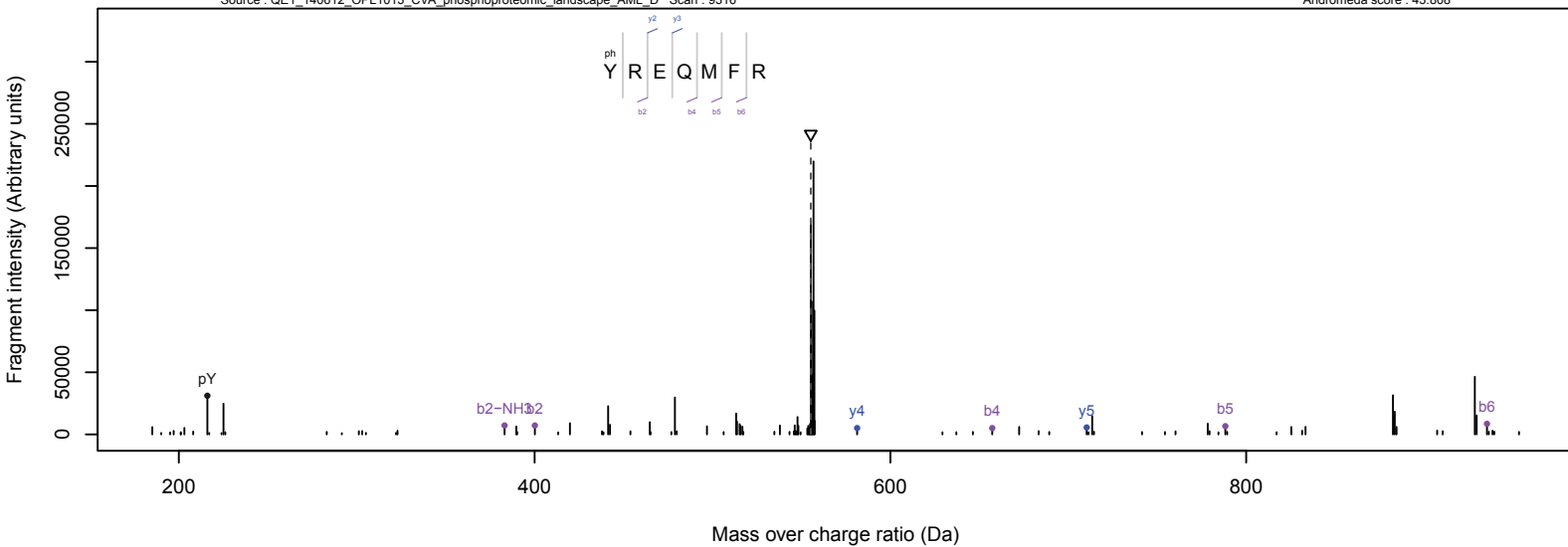
Andromeda score : 43.808



## Y(ph)REQMFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9316

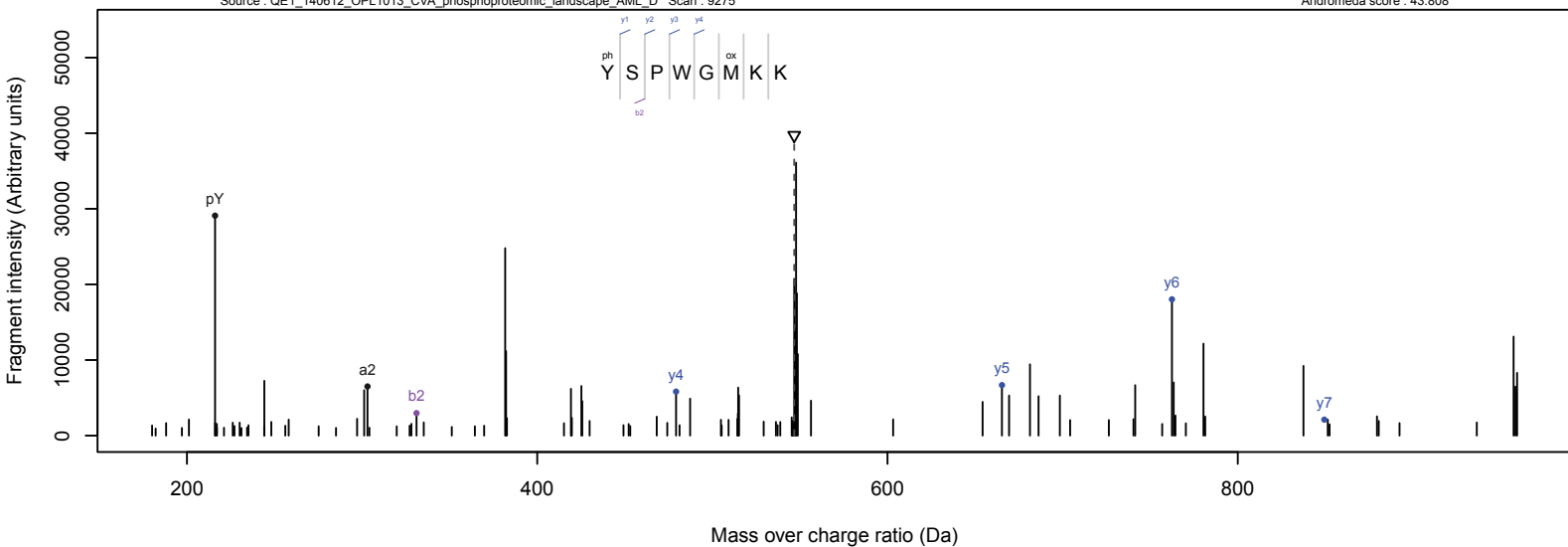
Andromeda score : 43.808



Y(ph)SPWGM(ox)KK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9275

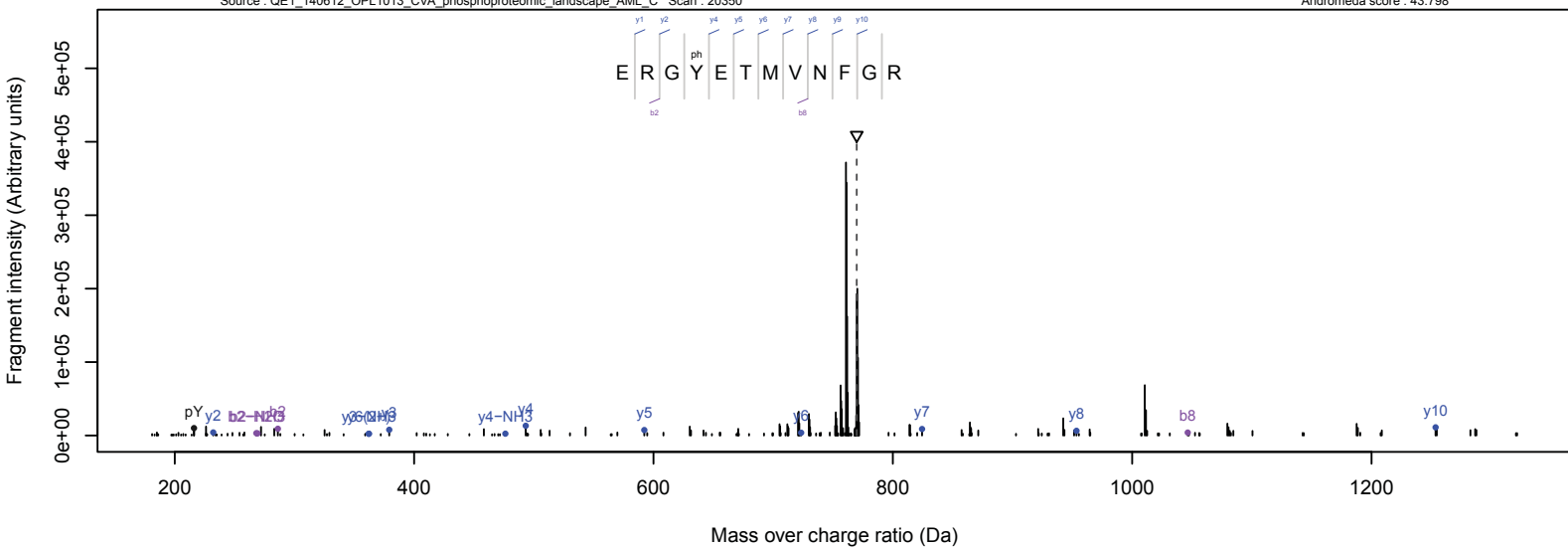
Andromeda score : 43.808



ERGY(ph)ETMVNFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 20350

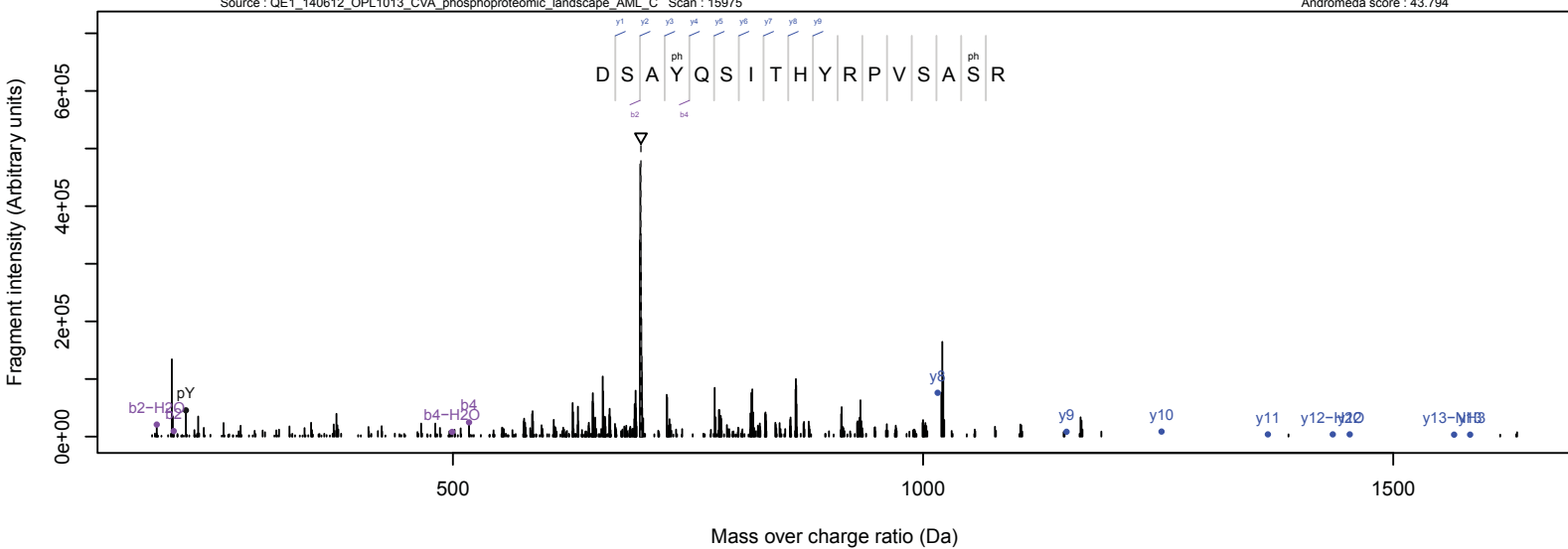
Andromeda score : 43.798

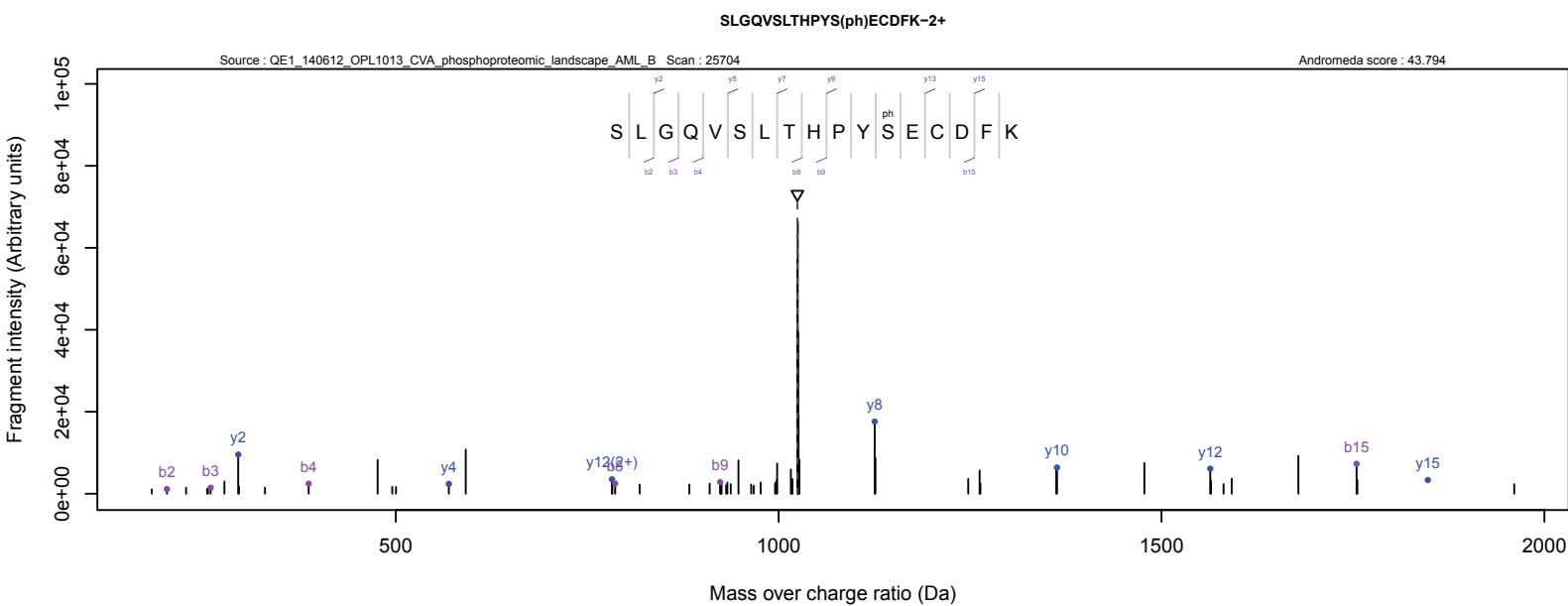
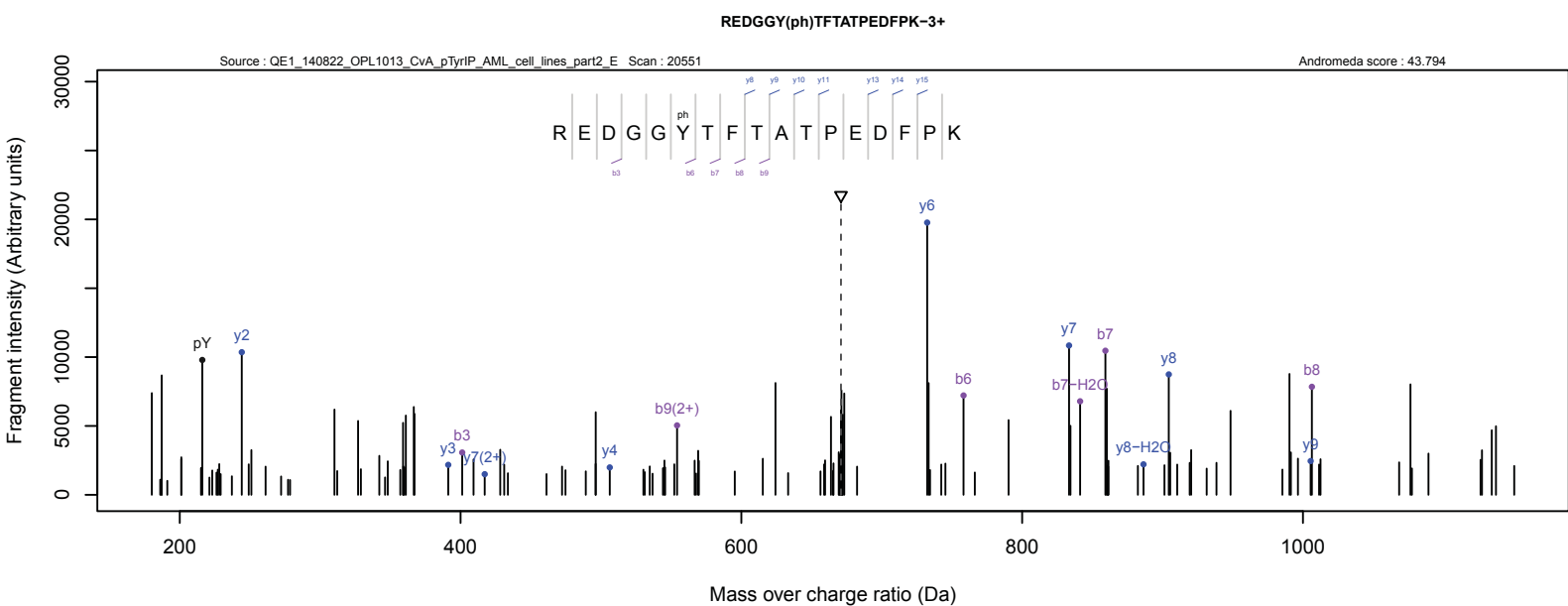
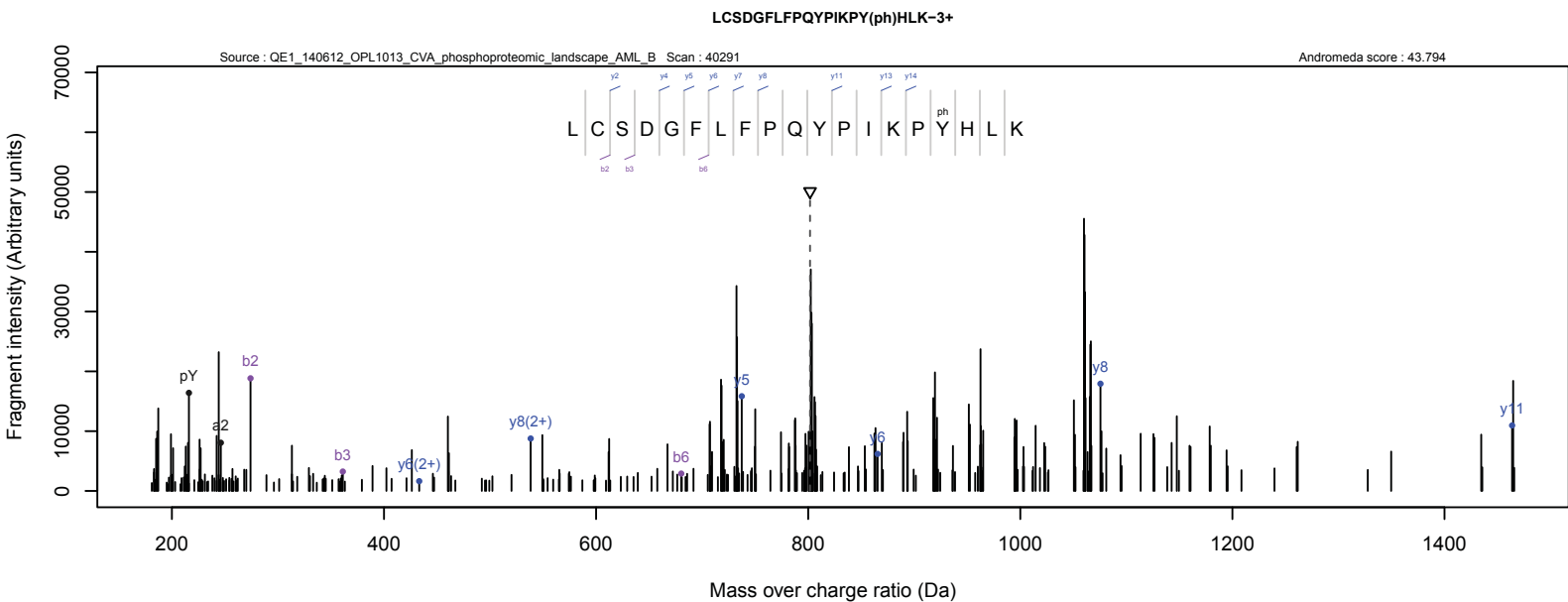


DSAY(ph)QSITHYRPVSAS(ph)R-3+

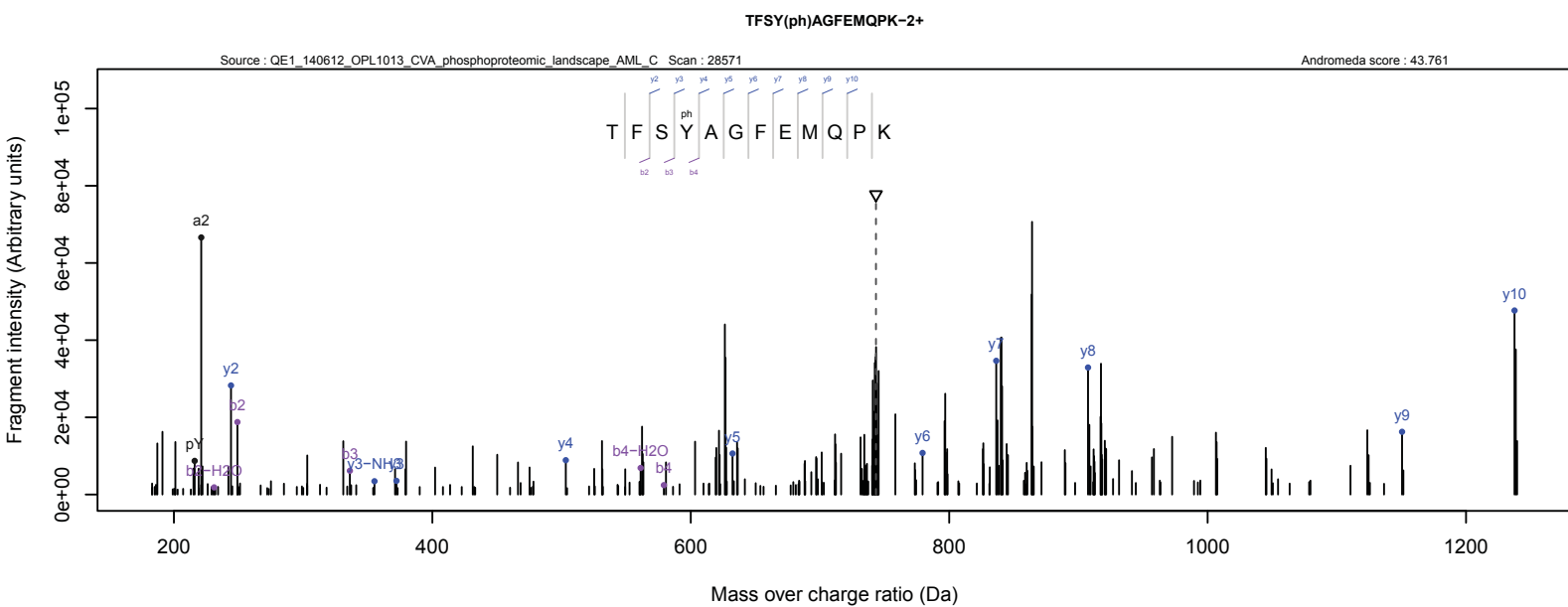
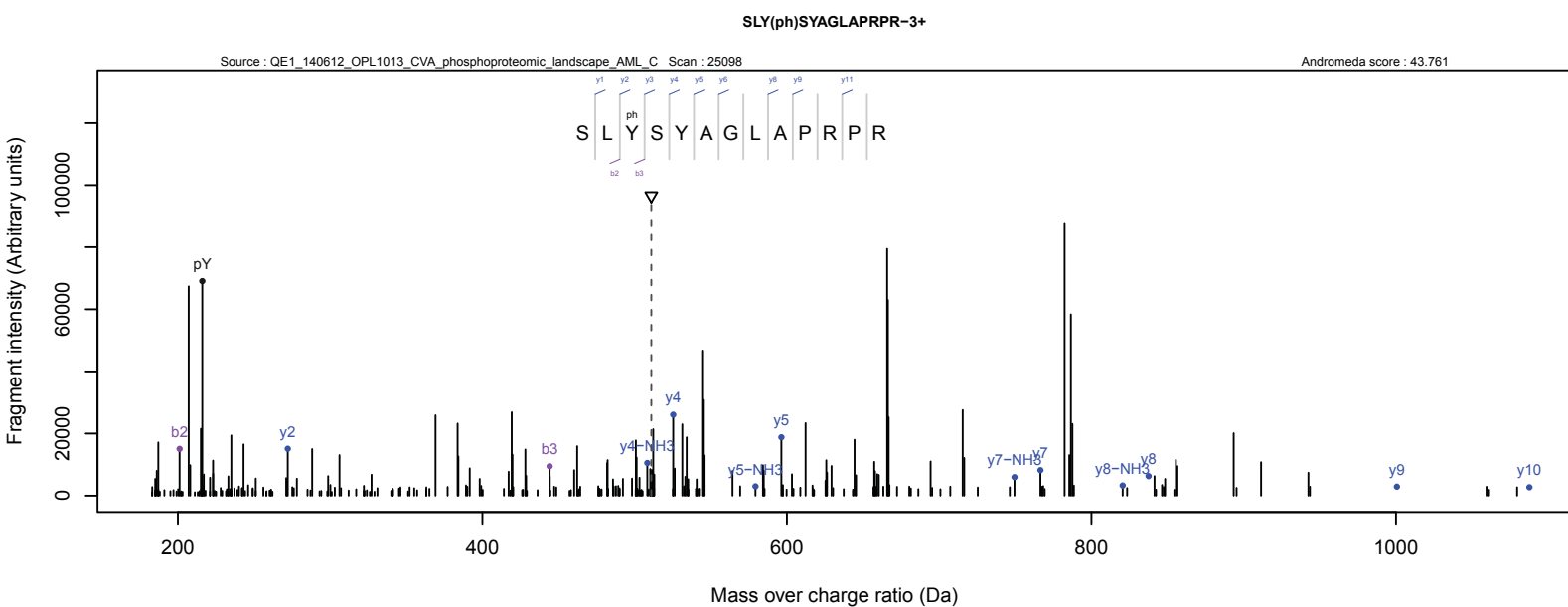
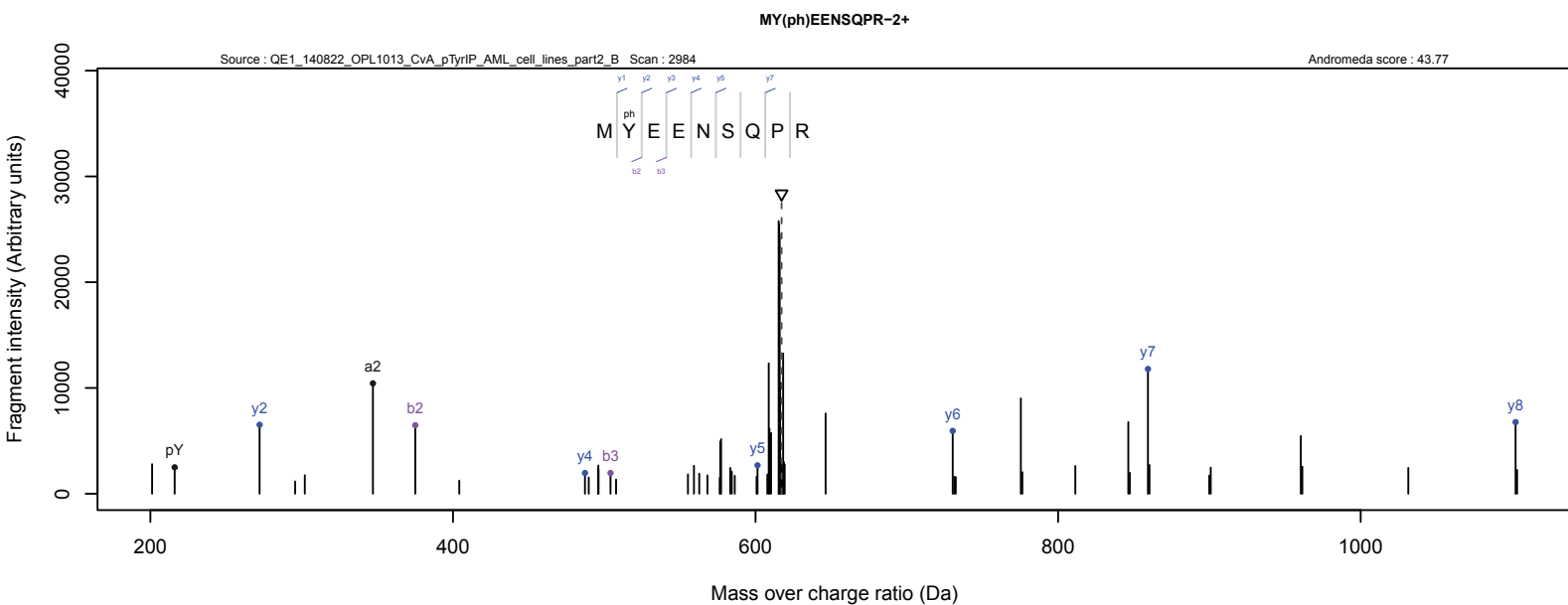
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15975

Andromeda score : 43.794





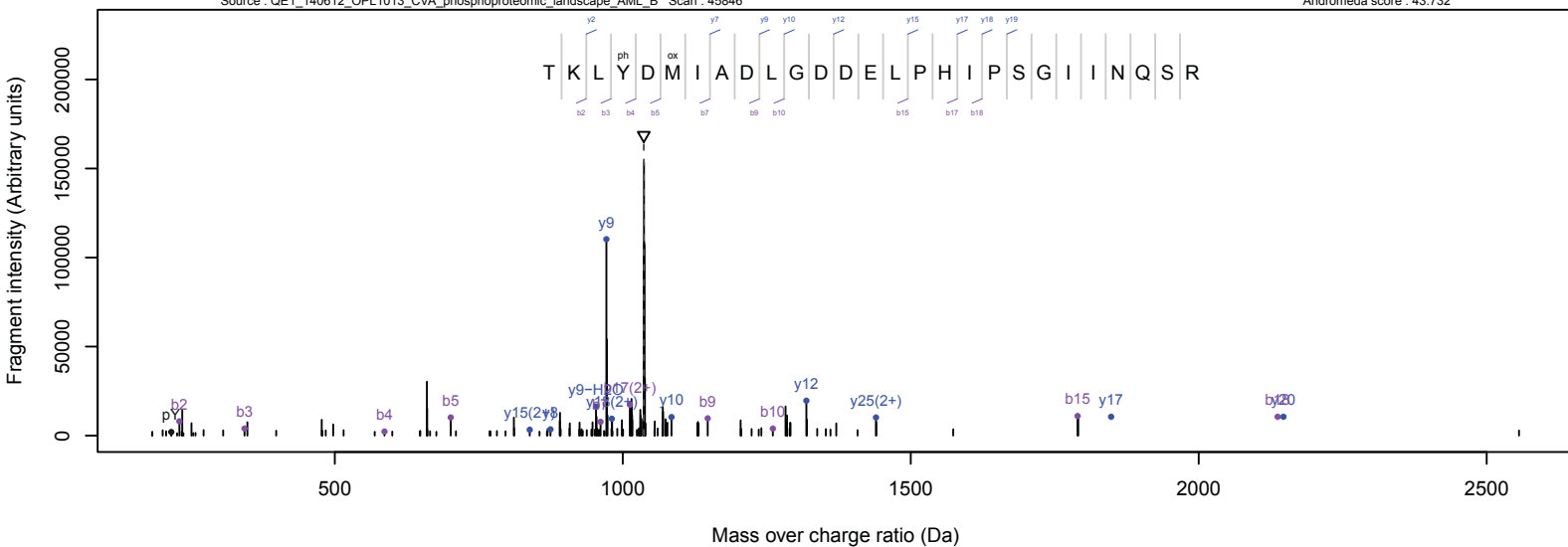




## TKLY(ph)DM(ox)IADLGDDDELPHIPSGIINQSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45846

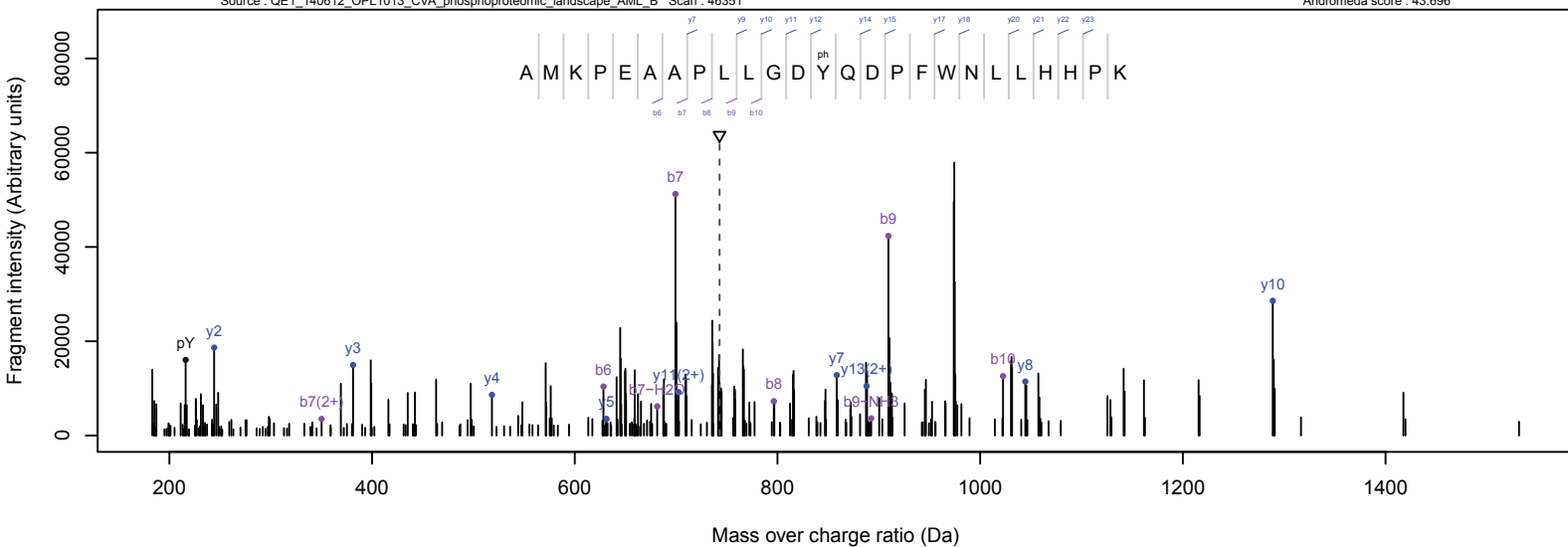
Andromeda score : 43.732



## AMKPEAAPLLGDY(ph)QDPFWNLLHHPK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 46351

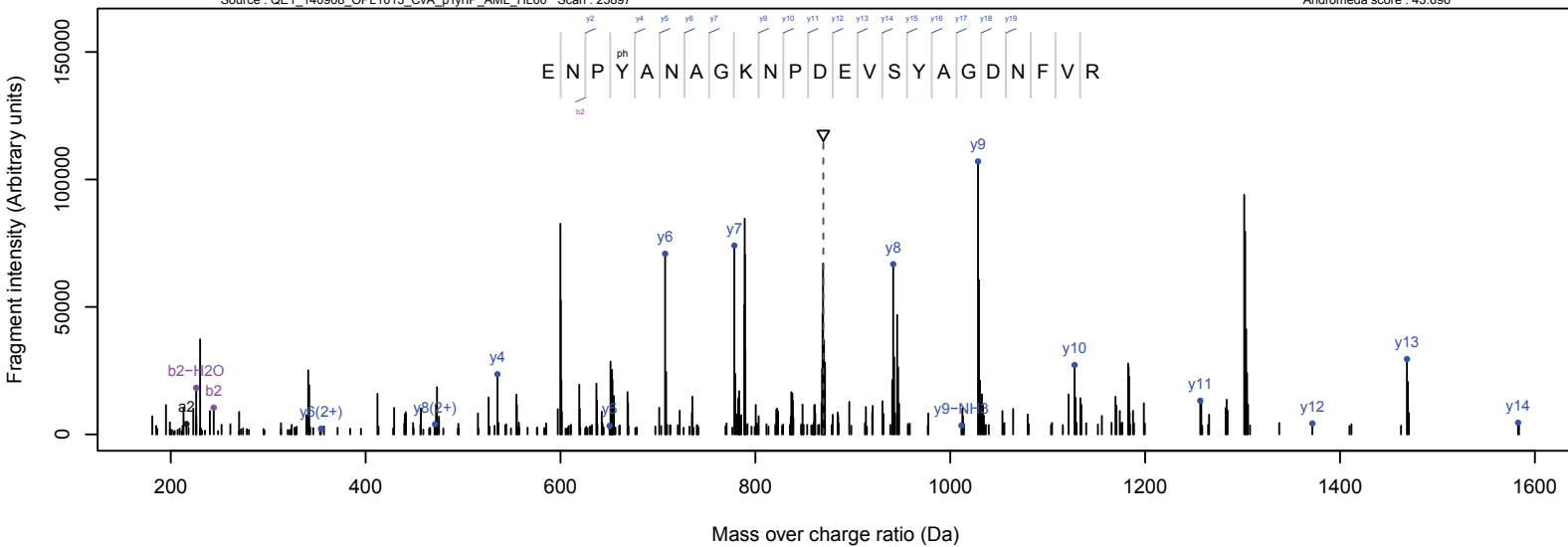
Andromeda score : 43.696

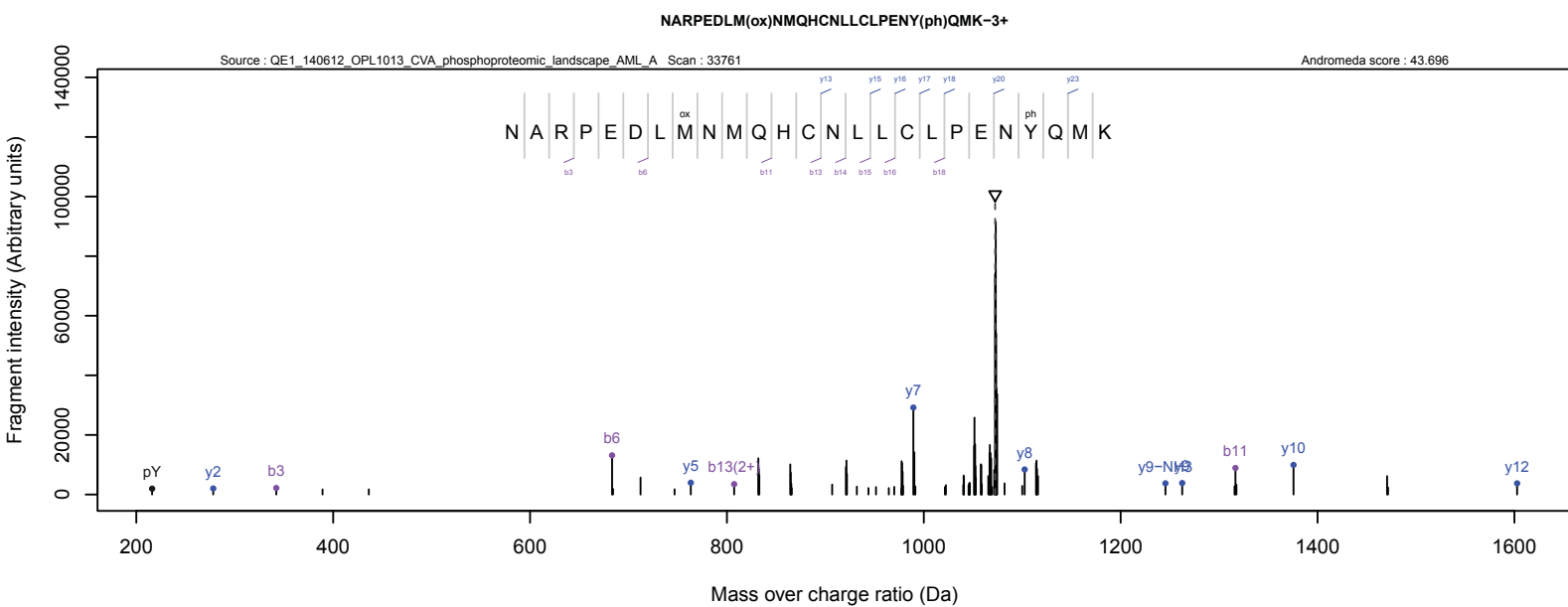
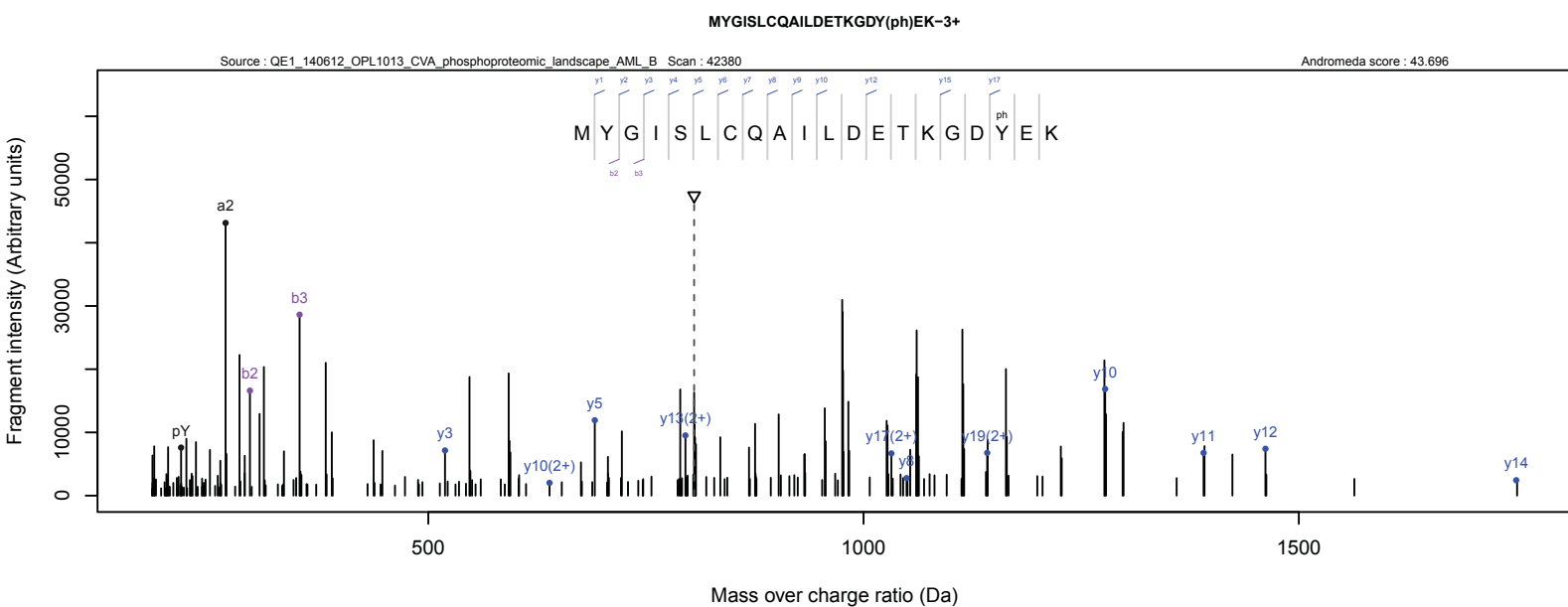
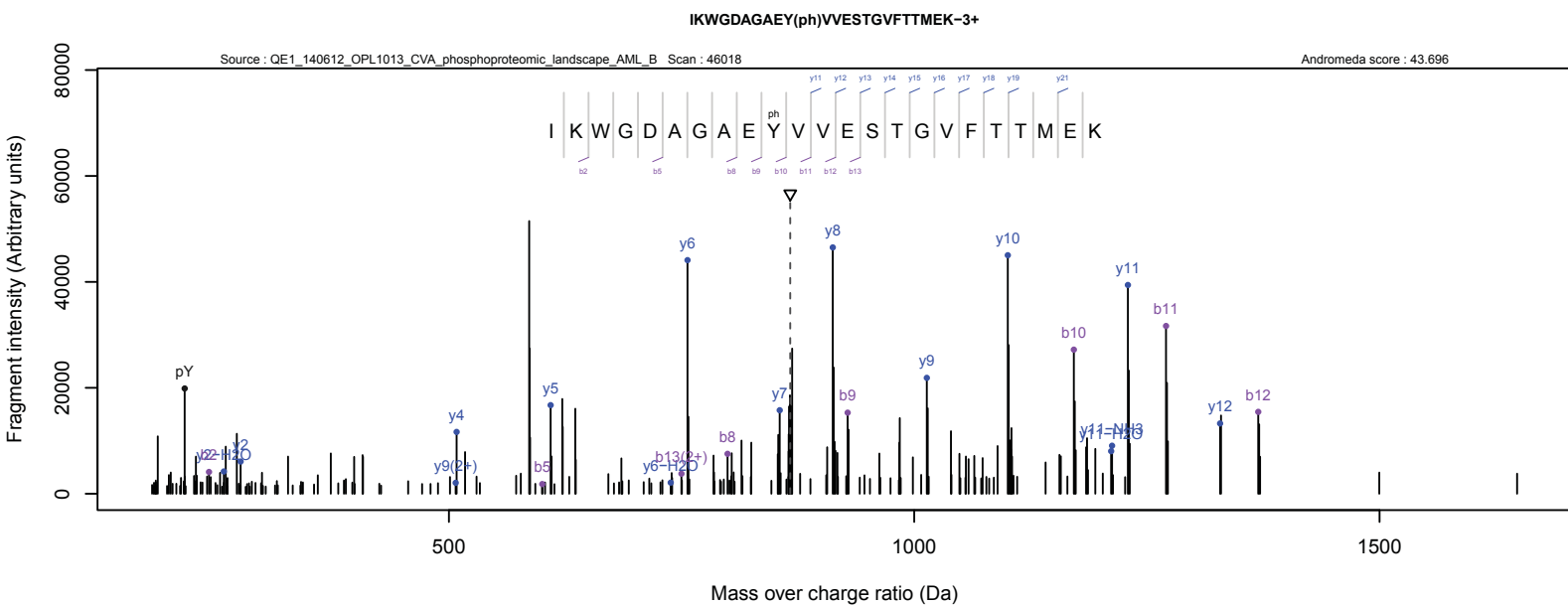


## ENPY(ph)ANAGKNPDEVSYAGDNFVR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 25897

Andromeda score : 43.696

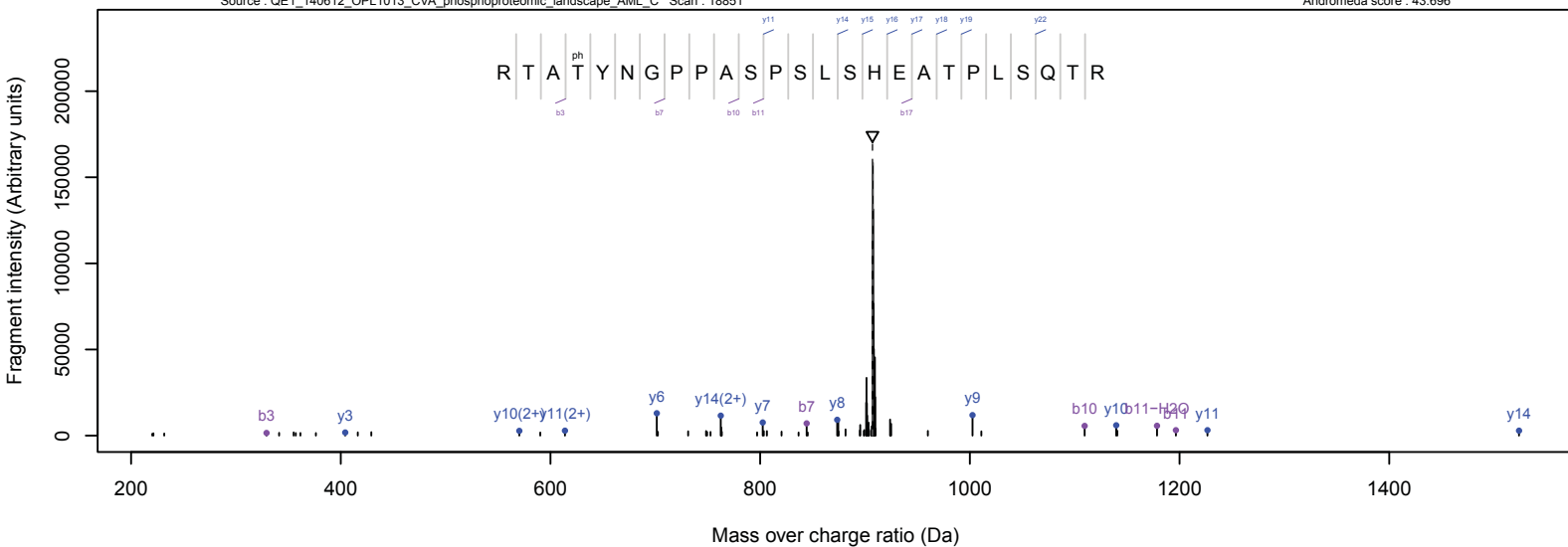




## RTAT(ph)YNGPPASPSLSHEATPLSQTR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18851

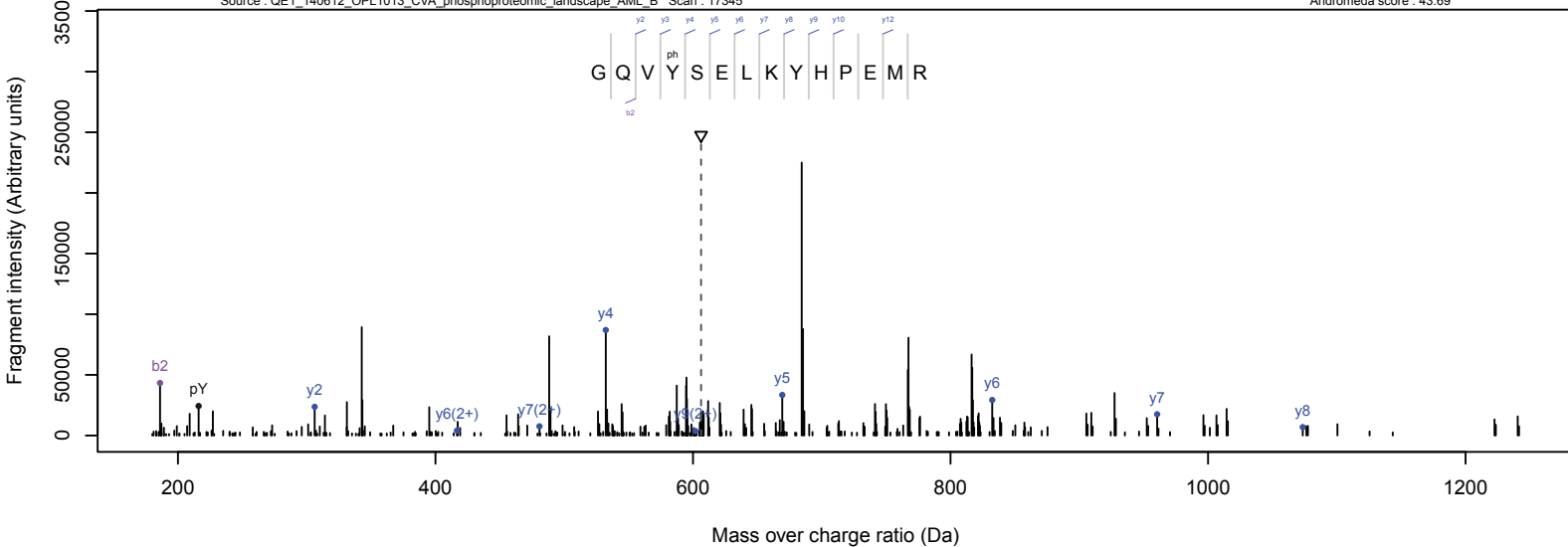
Andromeda score : 43.696



## GQVY(ph)SELKYHPEMR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17345

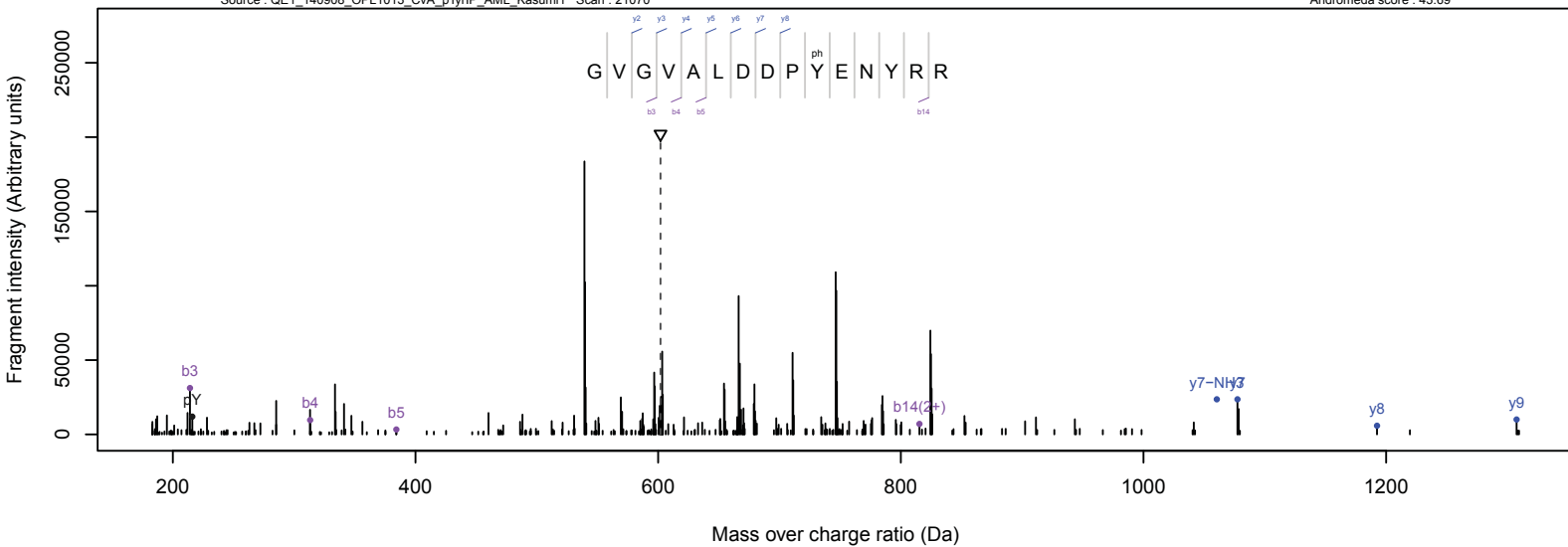
Andromeda score : 43.69

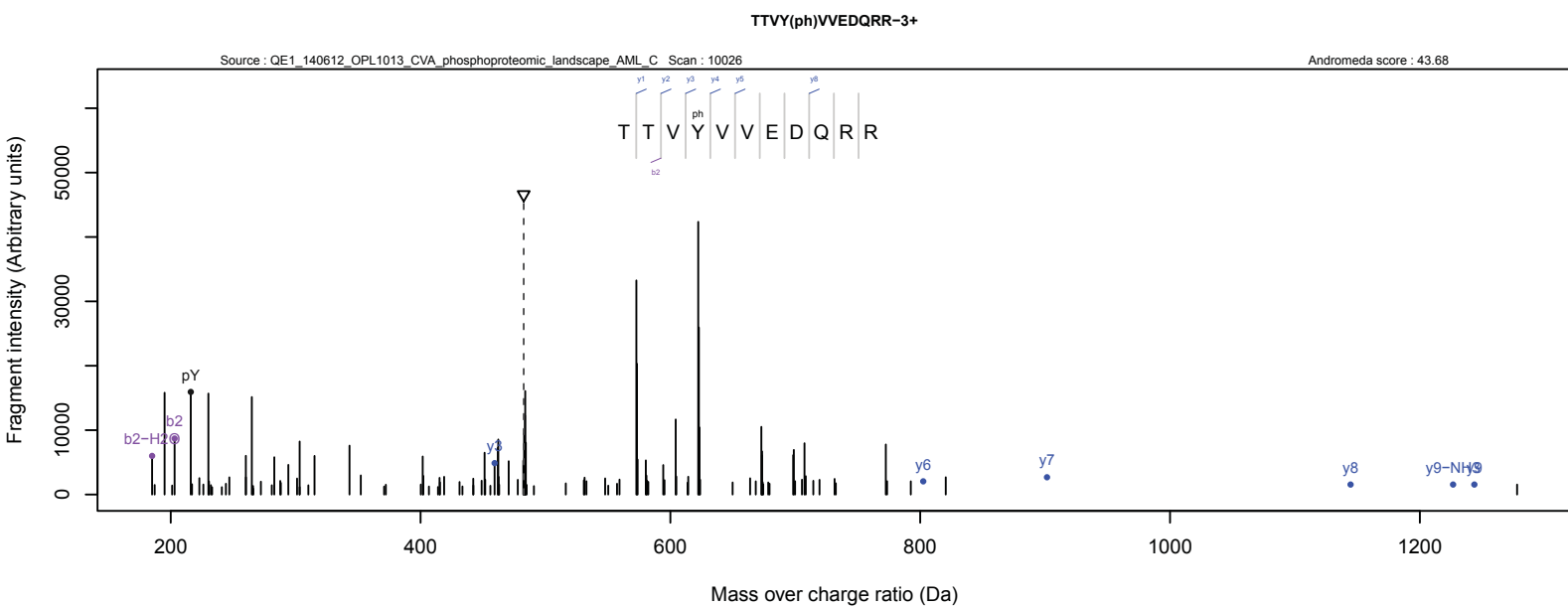
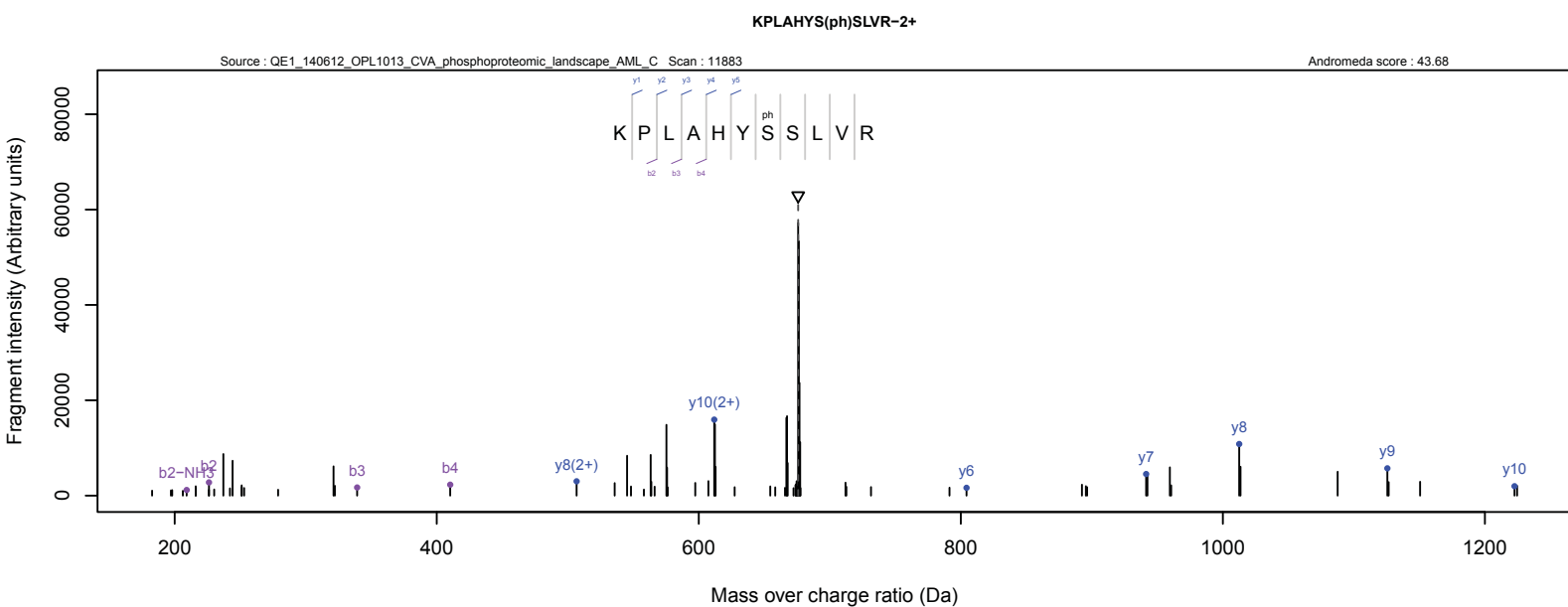
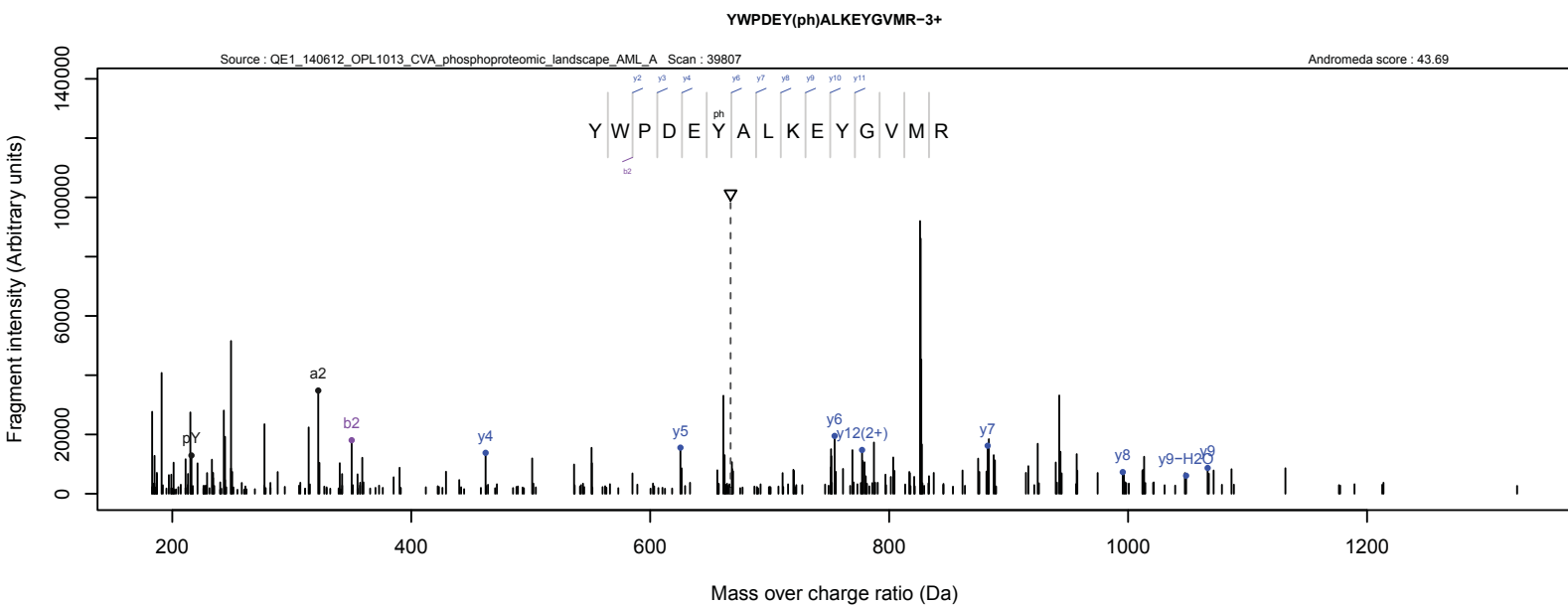


## GVGVALDDPY(ph)ENYRR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 21070

Andromeda score : 43.69

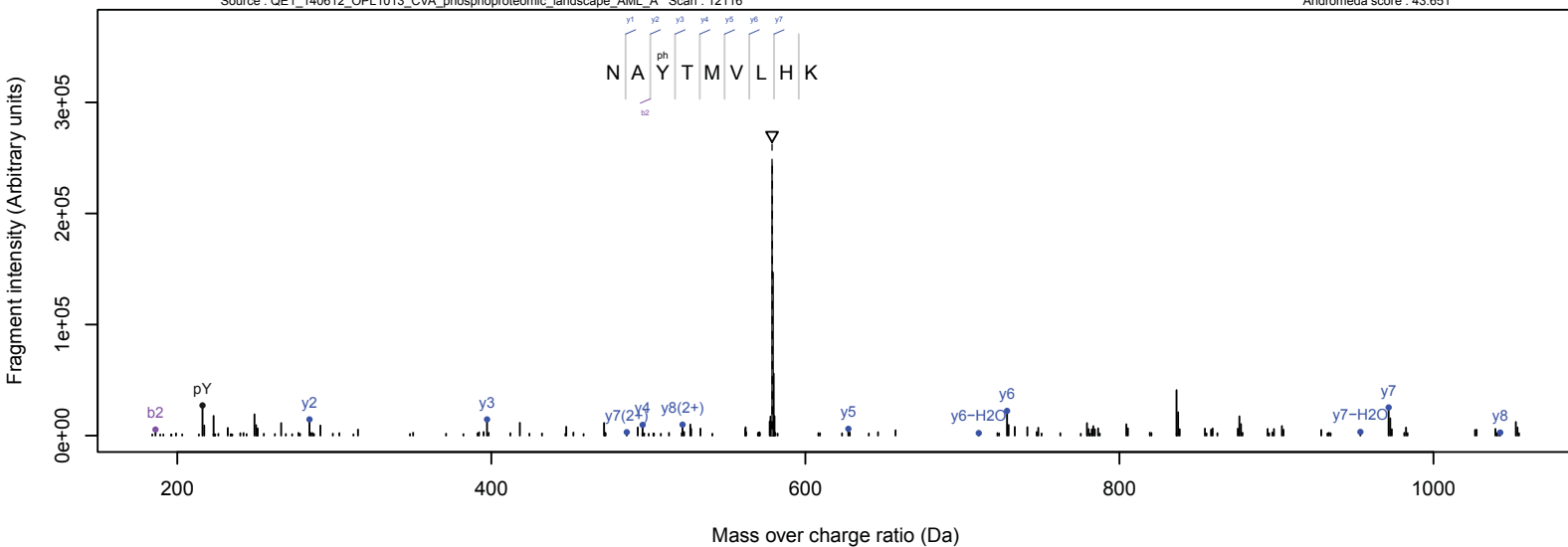




## NAY(ph)TMVLHK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 12116

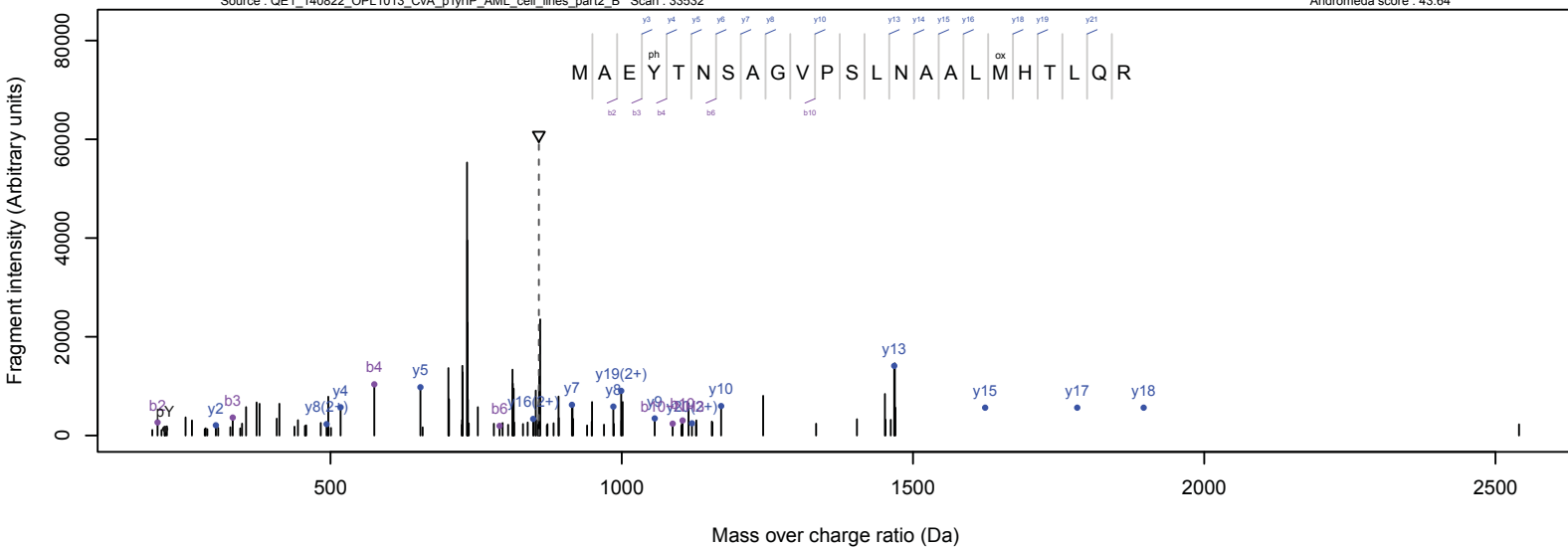
Andromeda score : 43.651



## MAEY(ph)TNSAGVPSLNAALM(ox)HTLQR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 33532

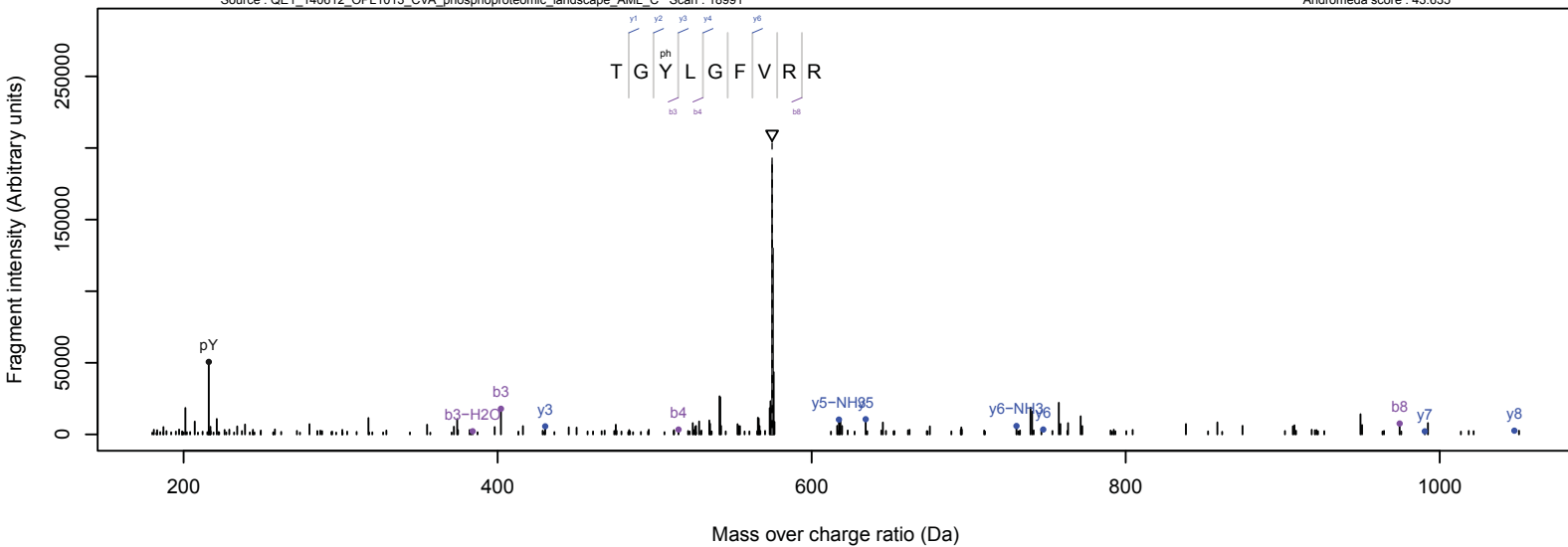
Andromeda score : 43.64



## TGY(ph)LGFVRR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18991

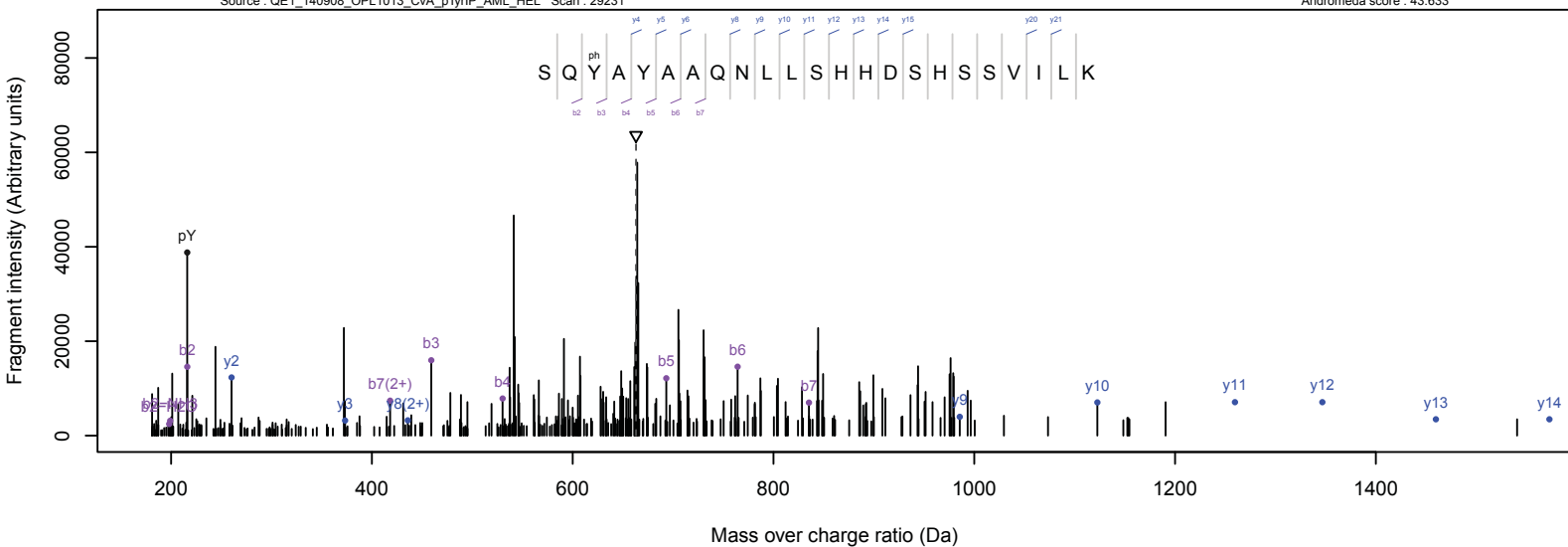
Andromeda score : 43.635



**SQY(ph)AYAAQNLLSHHDSHSSVILK-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29231

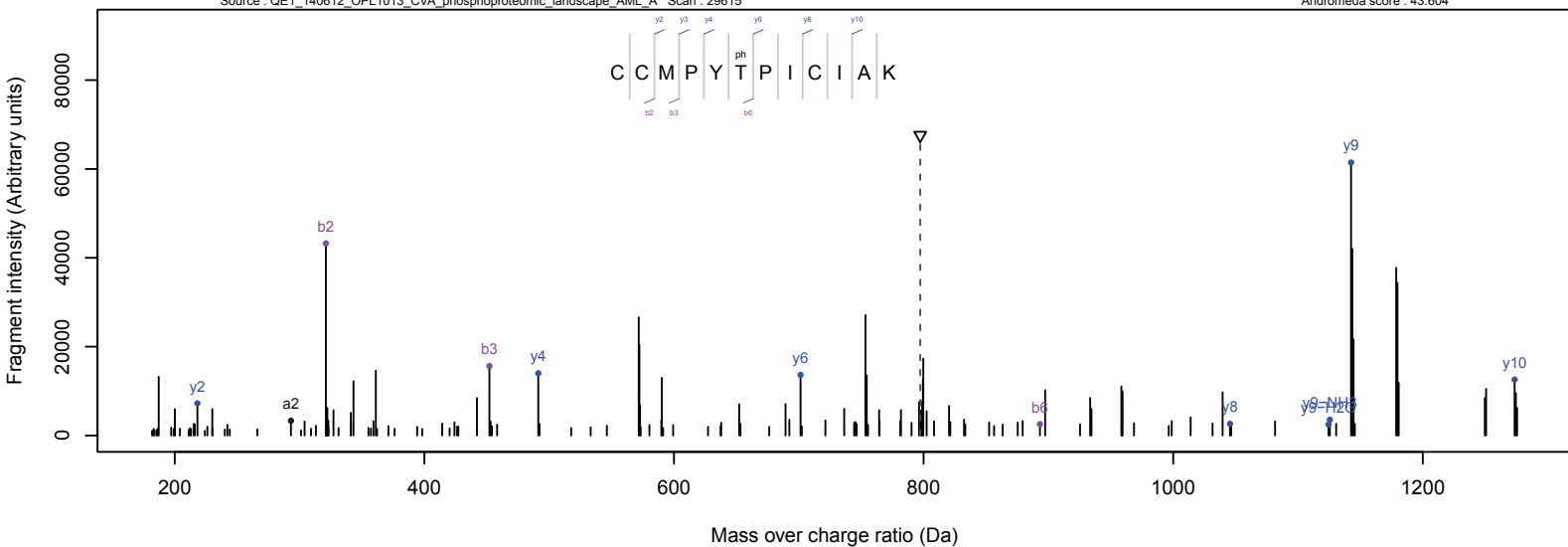
Andromeda score : 43.633



**CCMPYT(ph)PICI AK-2+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29615

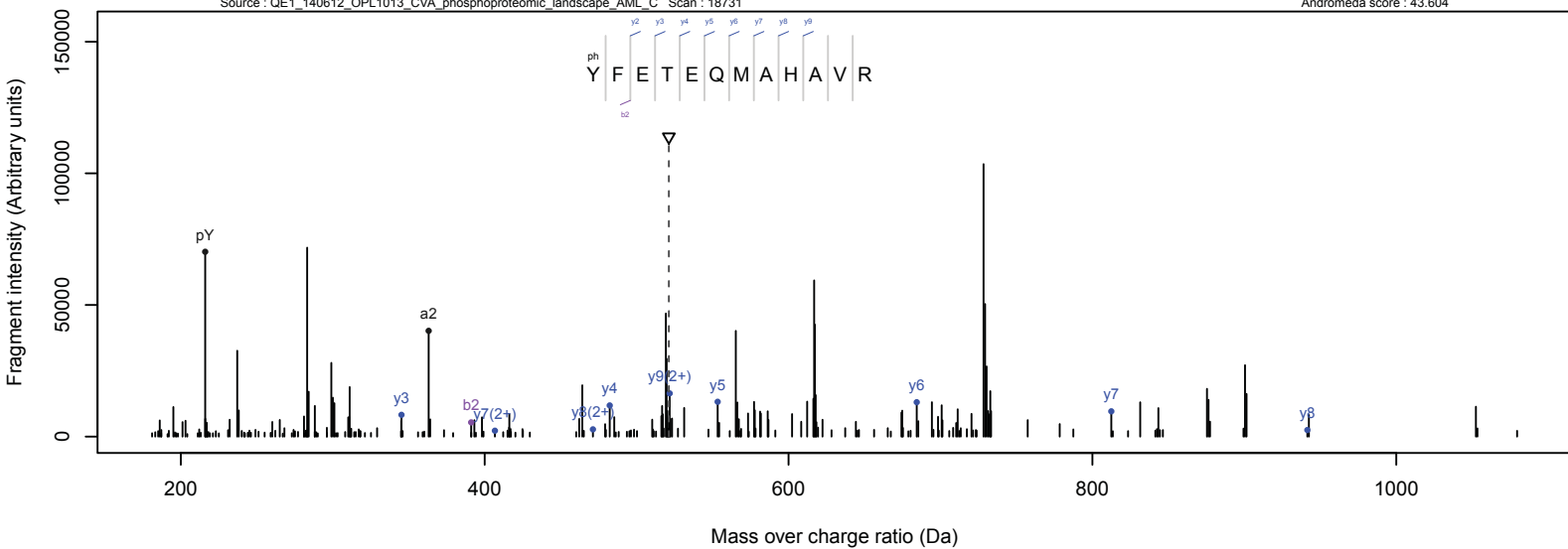
Andromeda score : 43.604



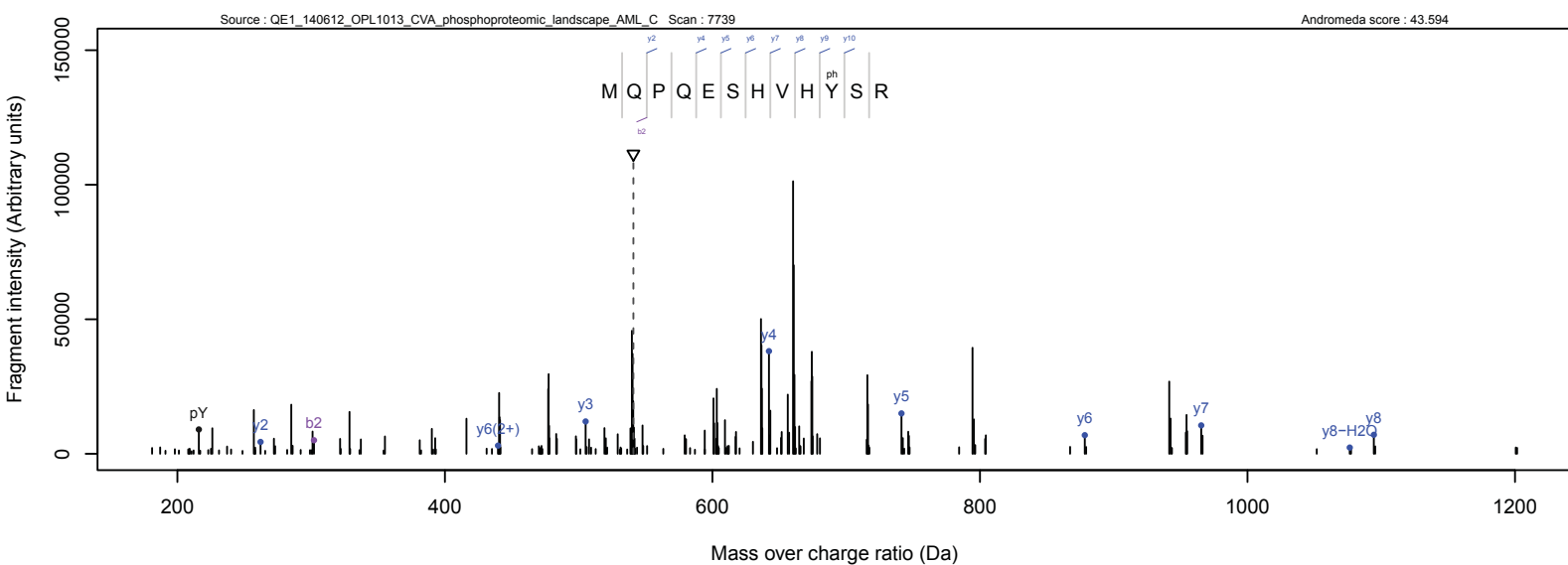
**Y(ph)FETEQAHA VR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 18731

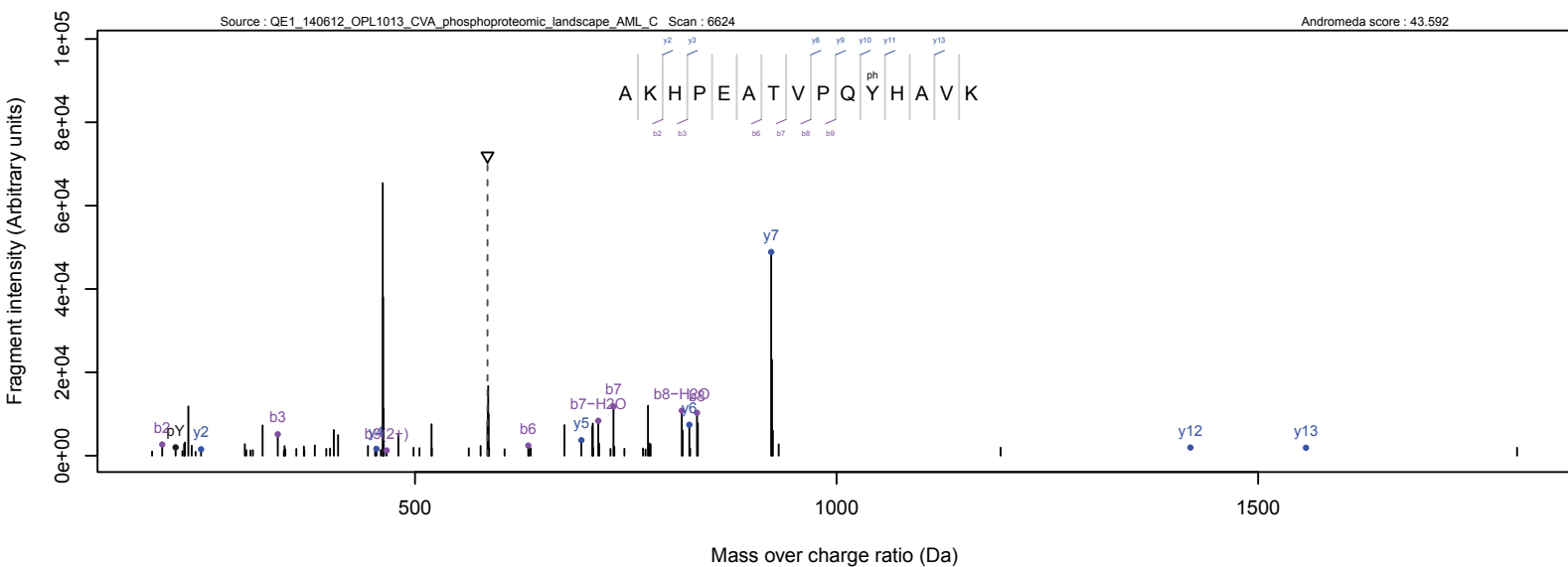
Andromeda score : 43.604



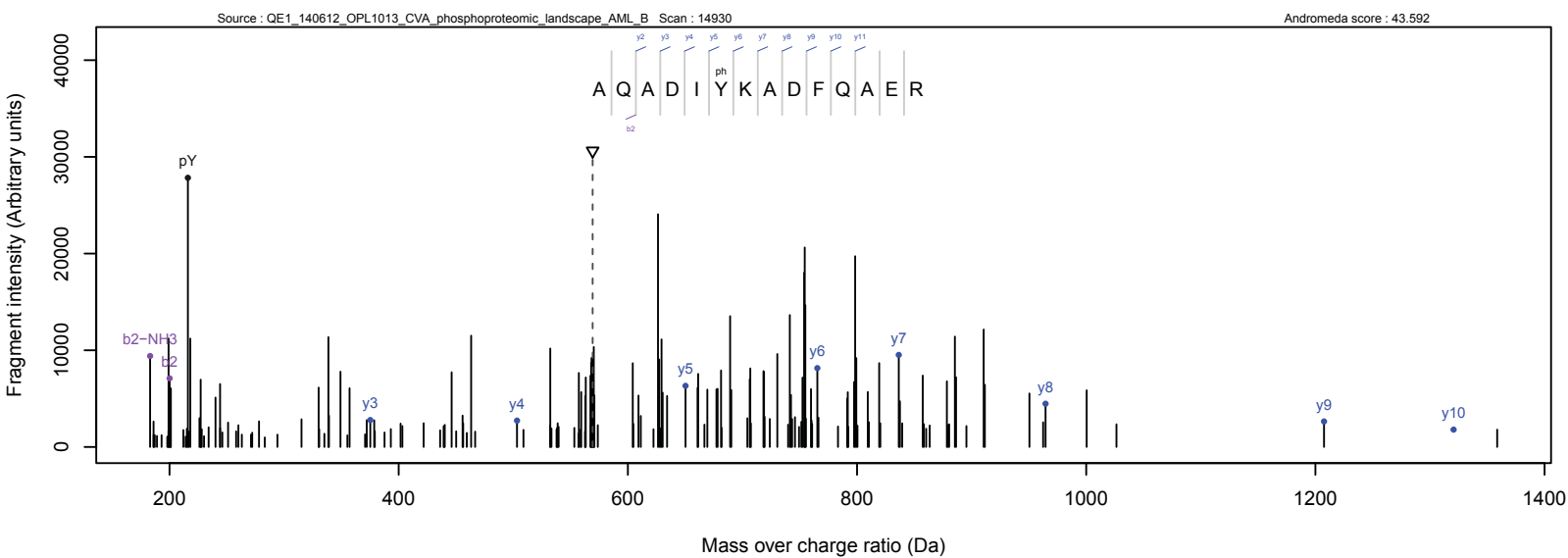
## (ac)MQPQESHVHY(ph)SR-3+



## AKHPEATVPQY(ph)HAVK-3+



## AQADIY(ph)KADFQAER-3+

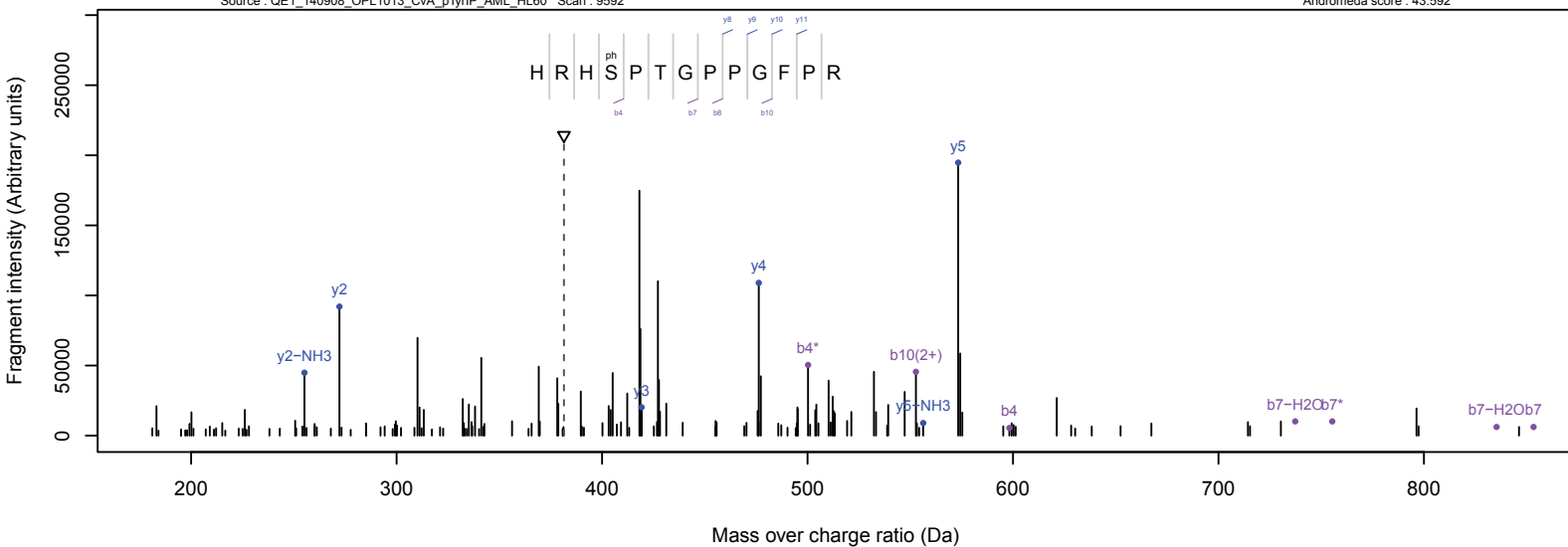




HRHS(ph)PTGPPGFPR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 9592

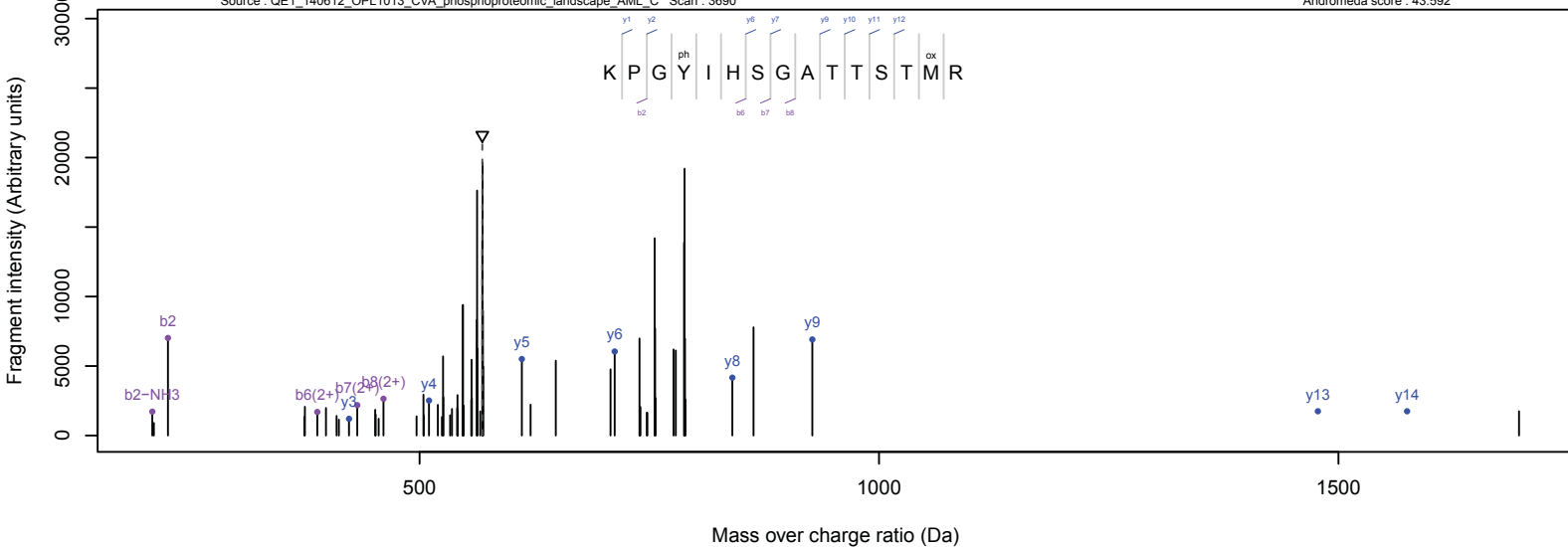
Andromeda score : 43.592



KPGY(ph)HSGATTSTM(ox)R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3690

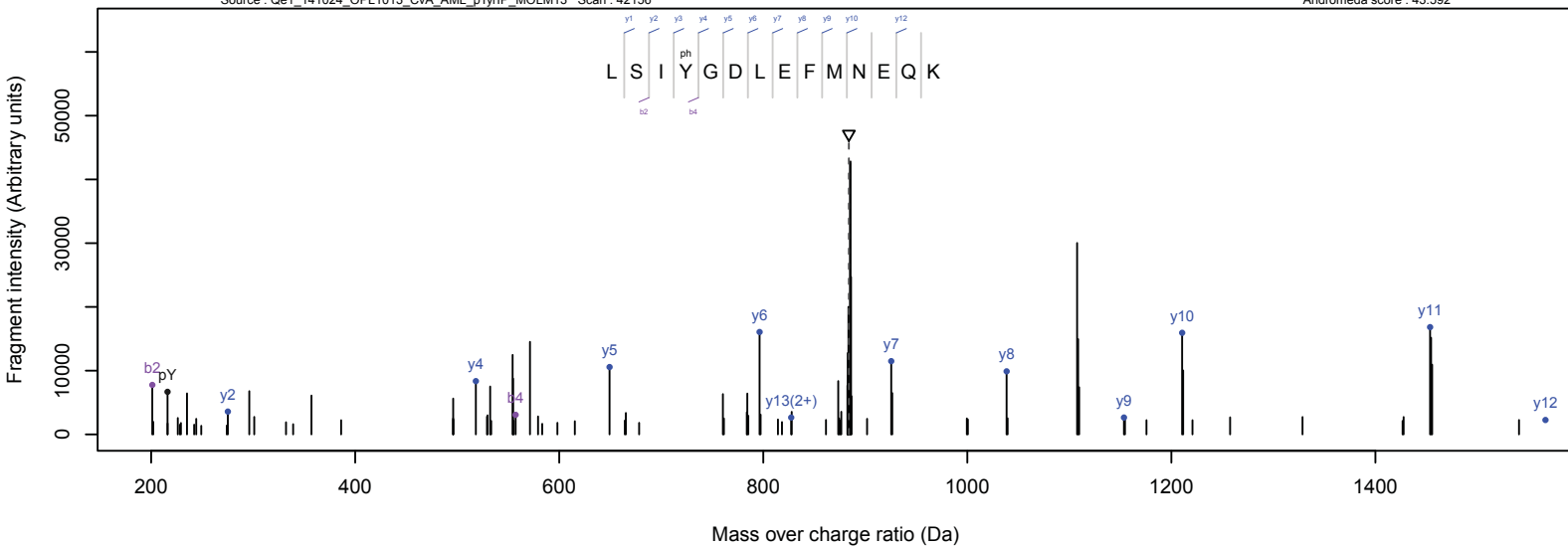
Andromeda score : 43.592



LSIY(ph)GDLEFMNEQK-2+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 42156

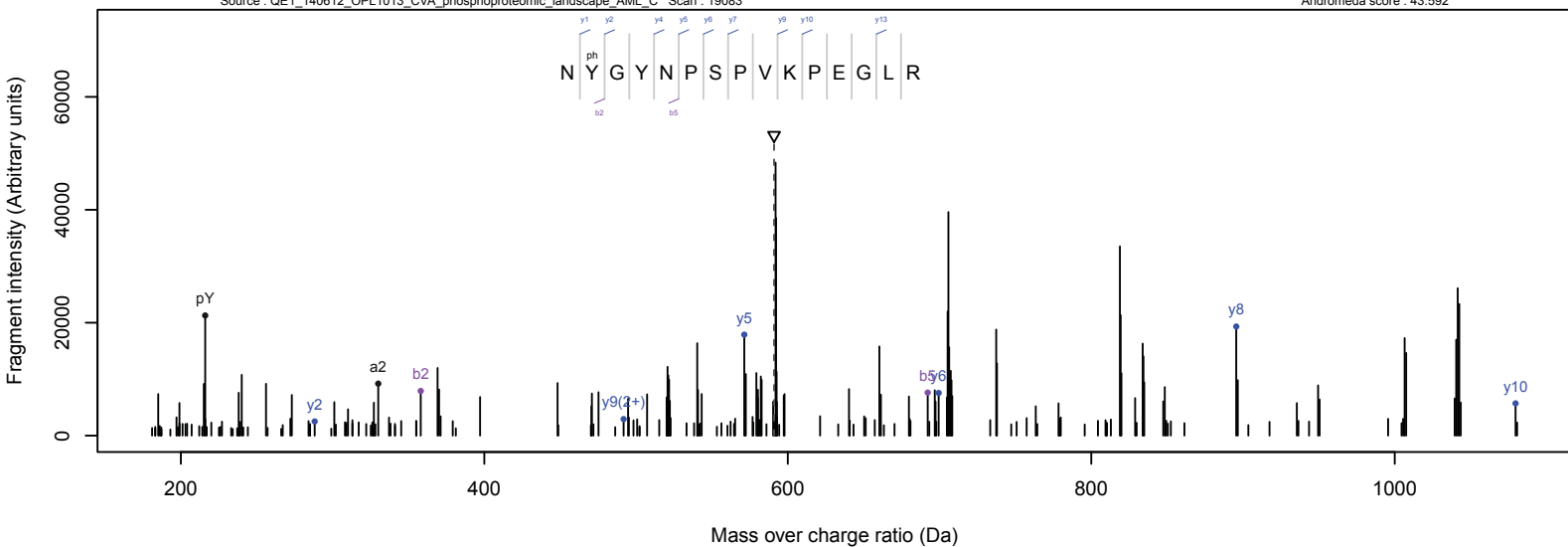
Andromeda score : 43.592



## NY(ph)GYNPSPVKPEGLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19083

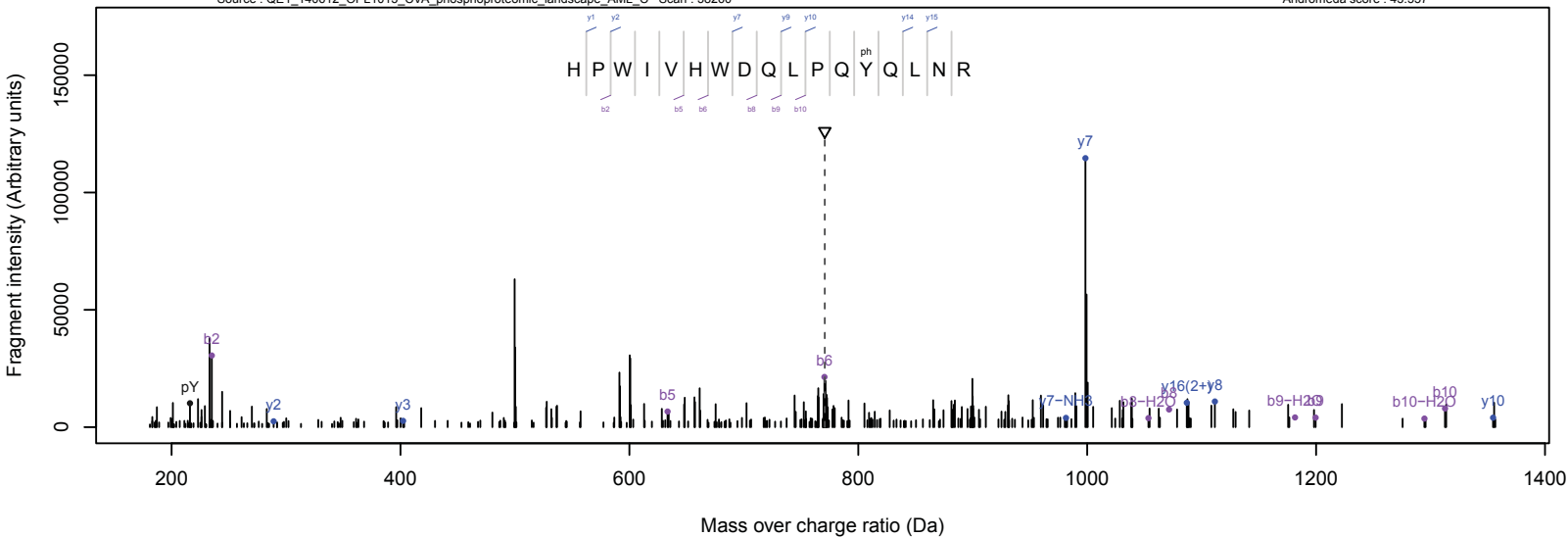
Andromeda score : 43.592



## HPWIVHWDQLPQYQLNR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38260

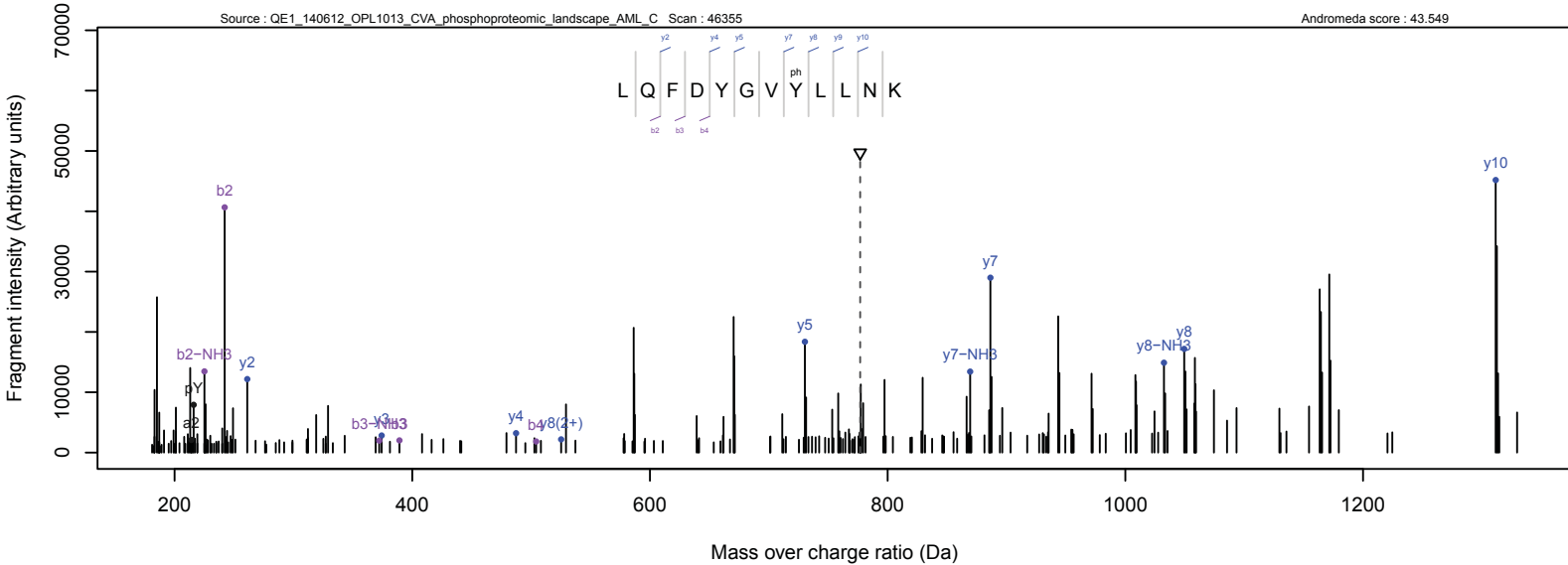
Andromeda score : 43.557



## LQFDYGVYLLNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 46355

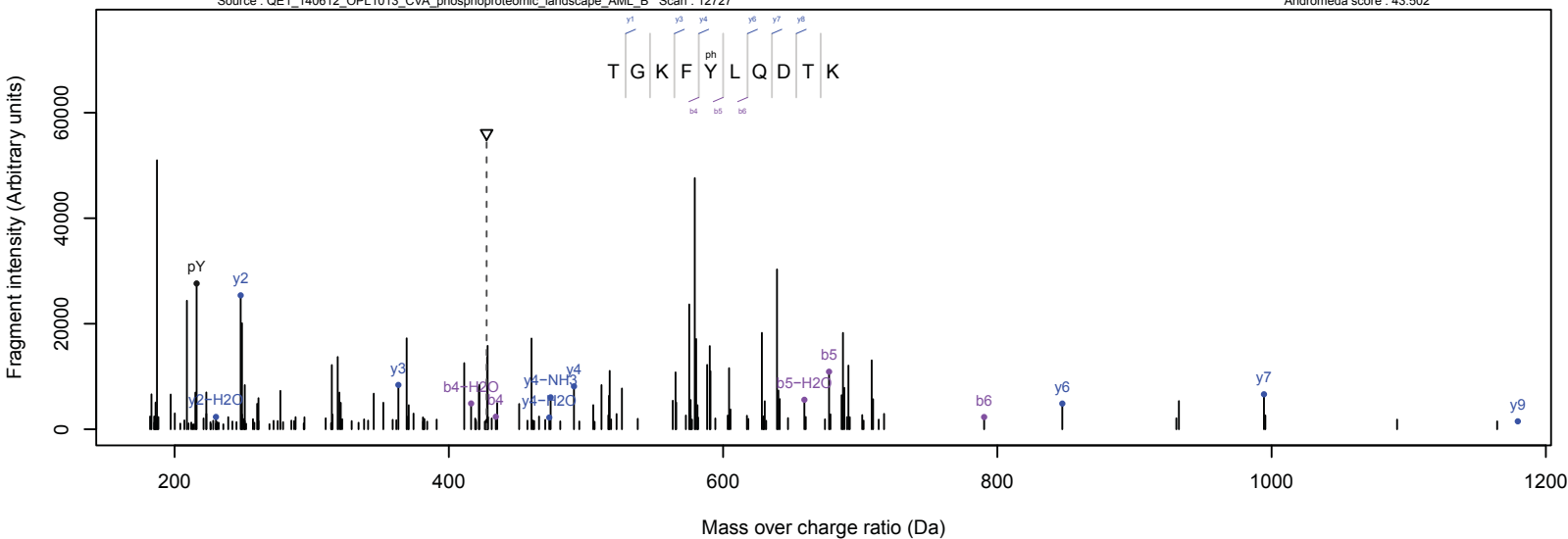
Andromeda score : 43.549



TGKFY(ph)LQDTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 12727

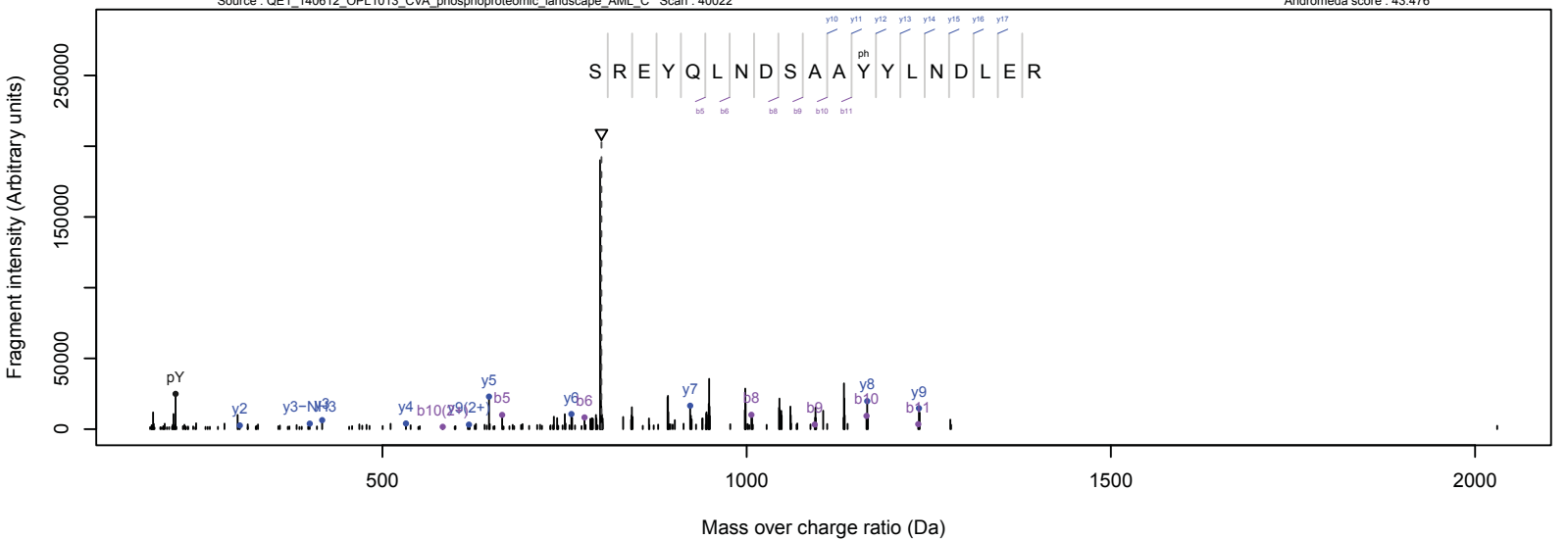
Andromeda score : 43.502



SREYQLNDSAAY(ph)YLNDLER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40022

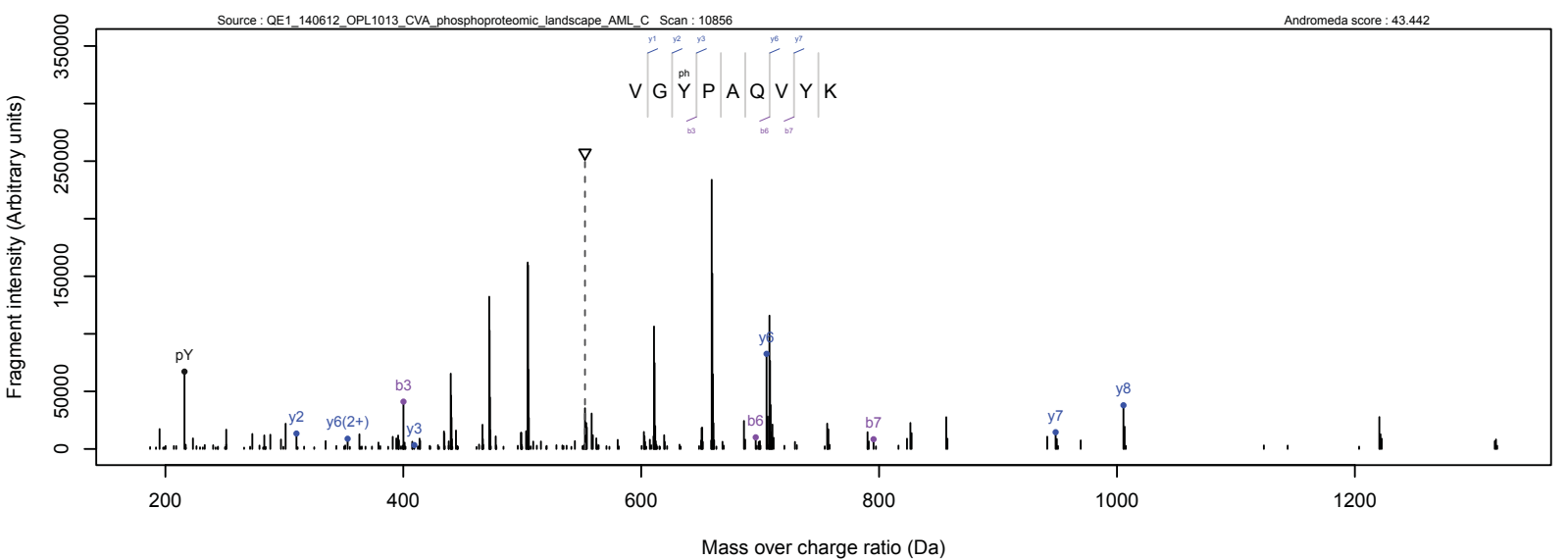
Andromeda score : 43.476



VGYP A Q V Y K-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 10856

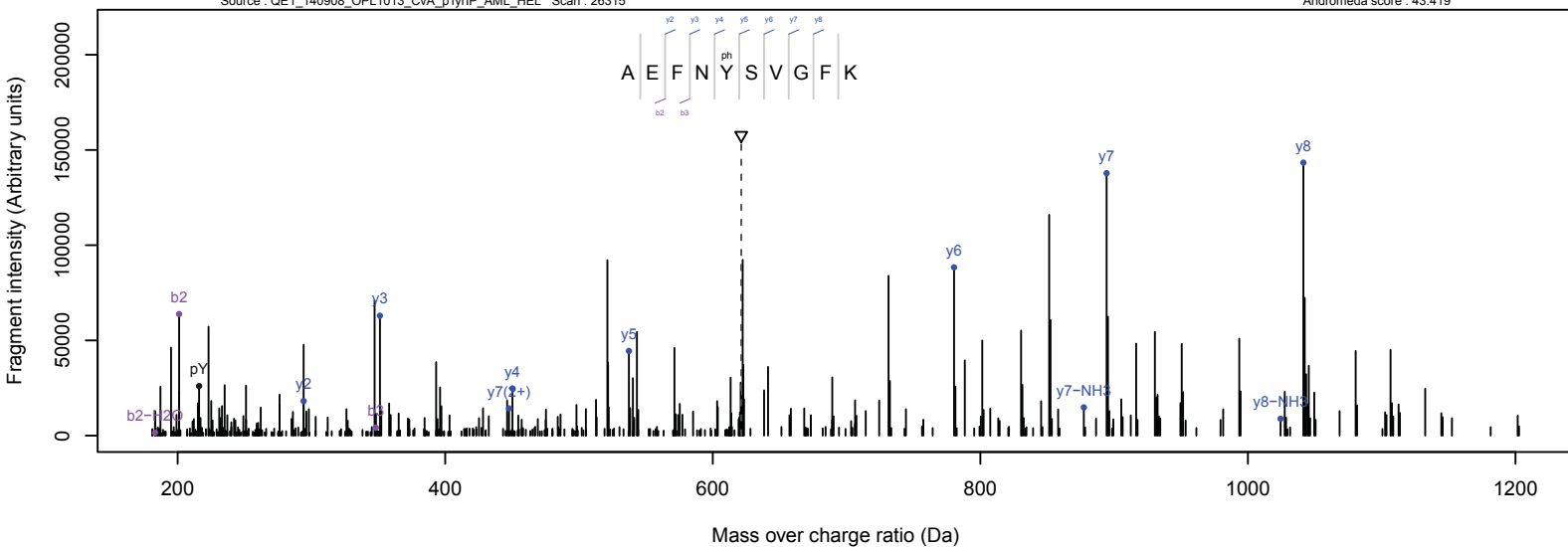
Andromeda score : 43.442



## AEFNY(ph)SVGFK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 26315

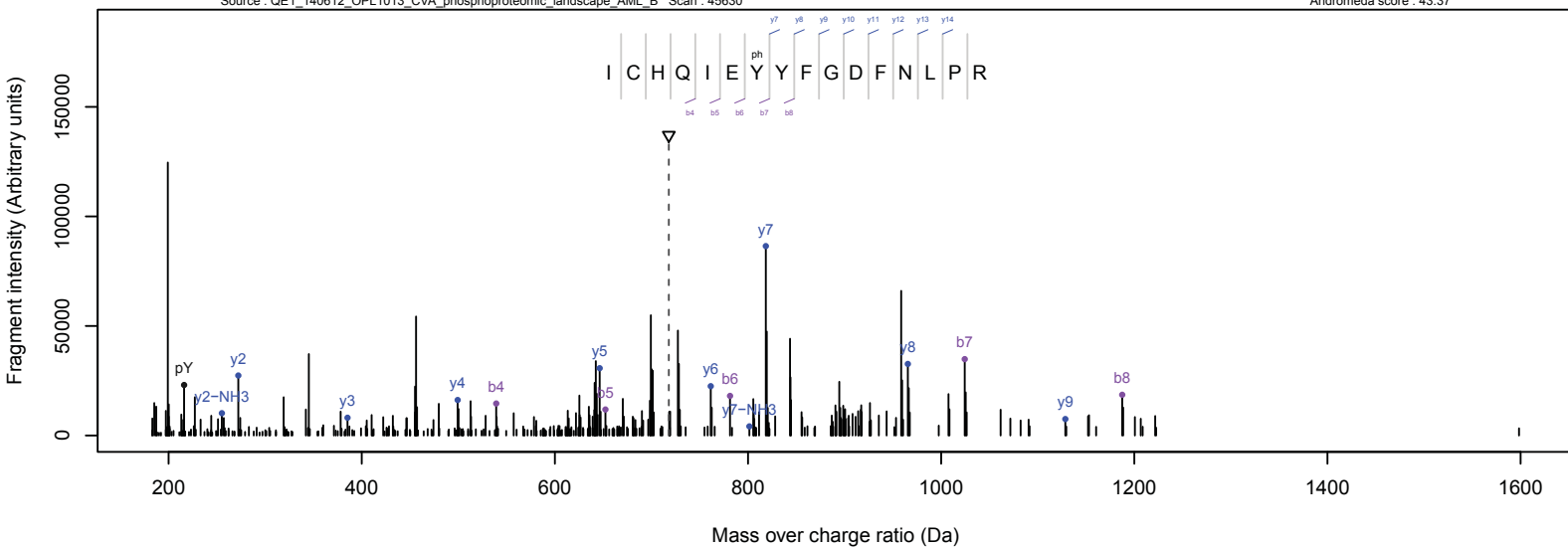
Andromeda score : 43.419



## ICHQIEY(ph)YFGDFNLPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45630

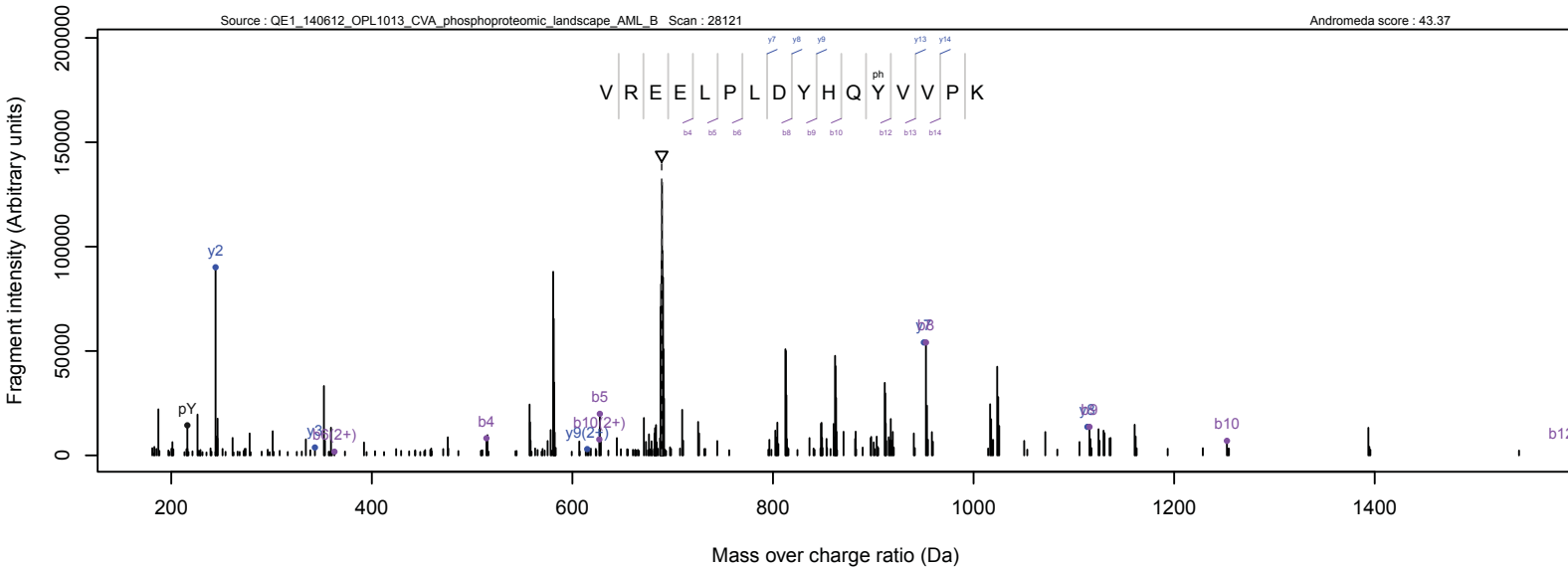
Andromeda score : 43.37



## VREELPDYHQY(ph)VVPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 28121

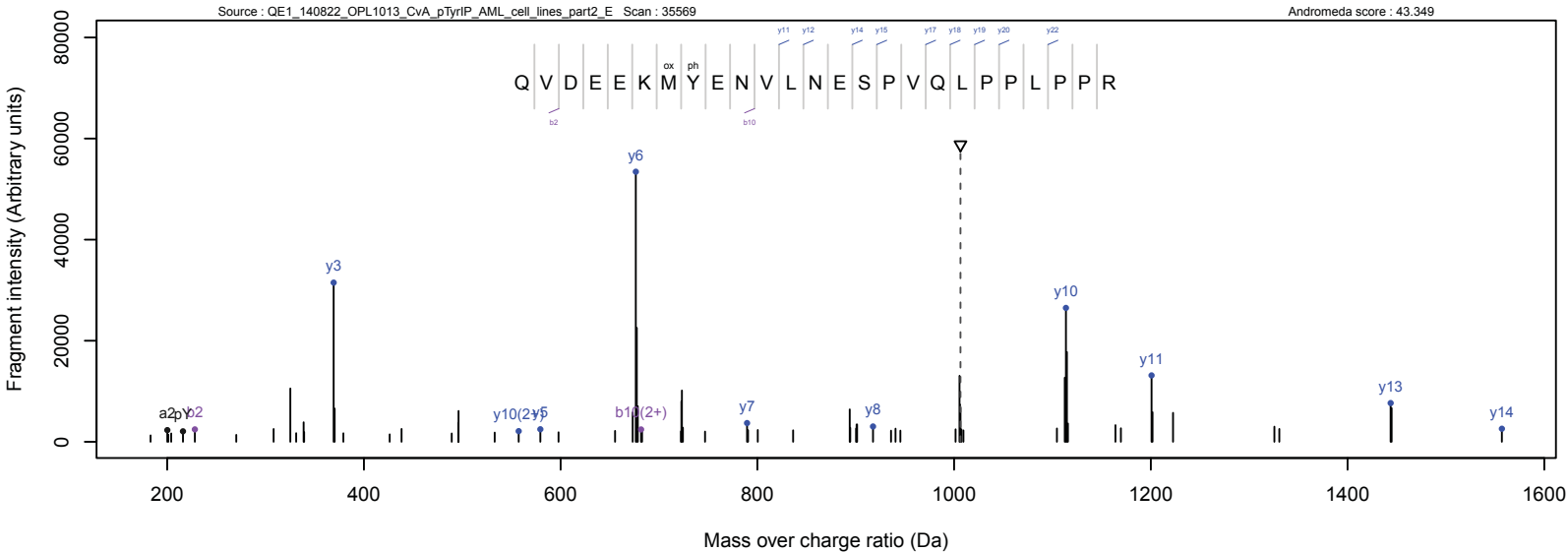
Andromeda score : 43.37



**QVDEEK(ox)Y(ph)ENLVNESPVQLPPLPPR-3+**

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 35569

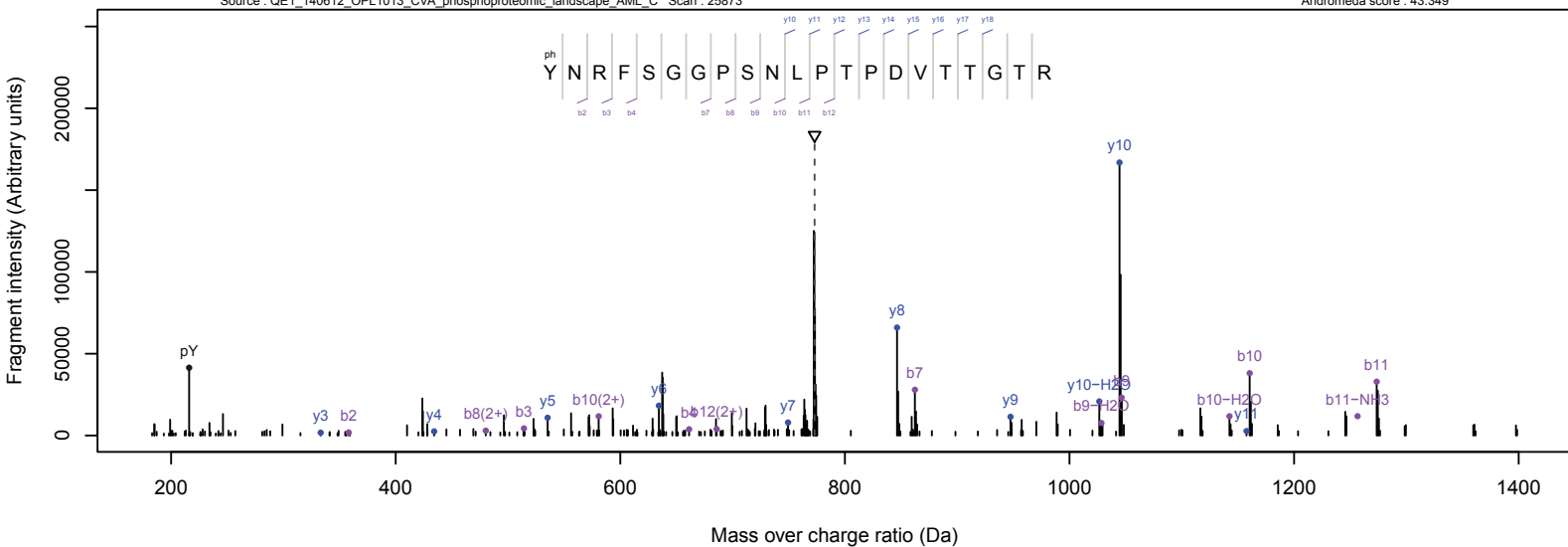
Andromeda score : 43.349



**Y(ph)NRFSGGPNLPTDVTTR-3+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 25873

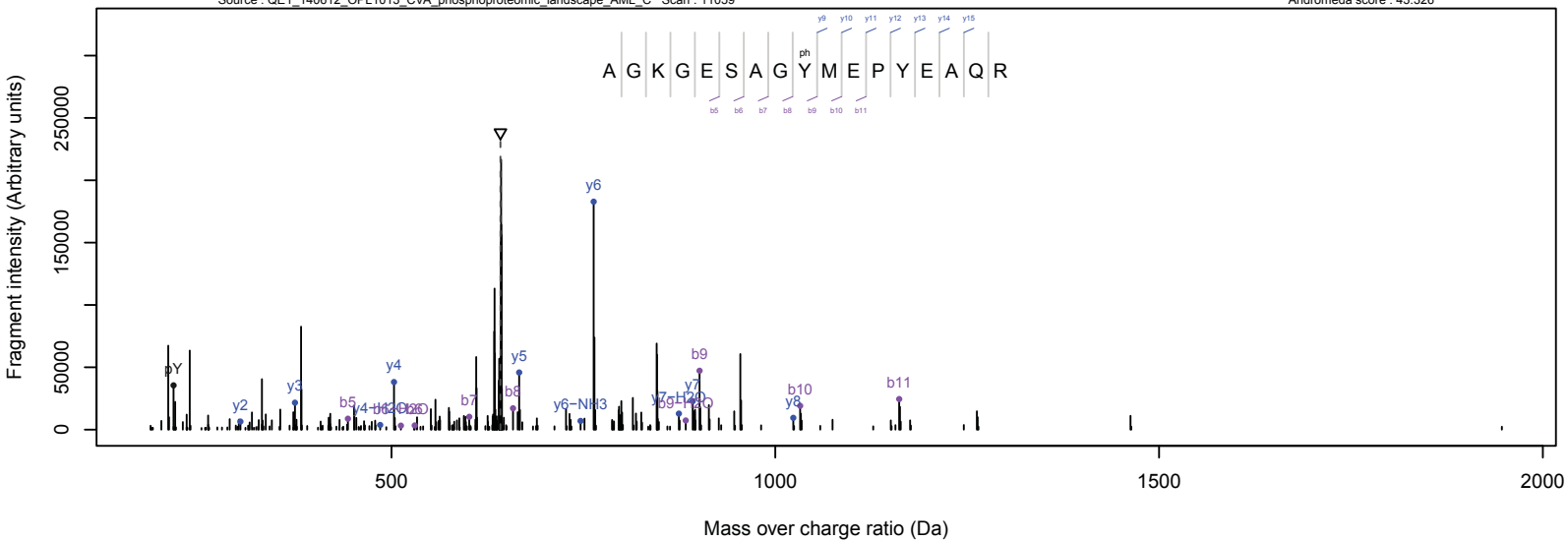
Andromeda score : 43.349



**AGKGESAGY(ph)MEPYEAQR-3+**

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 11059

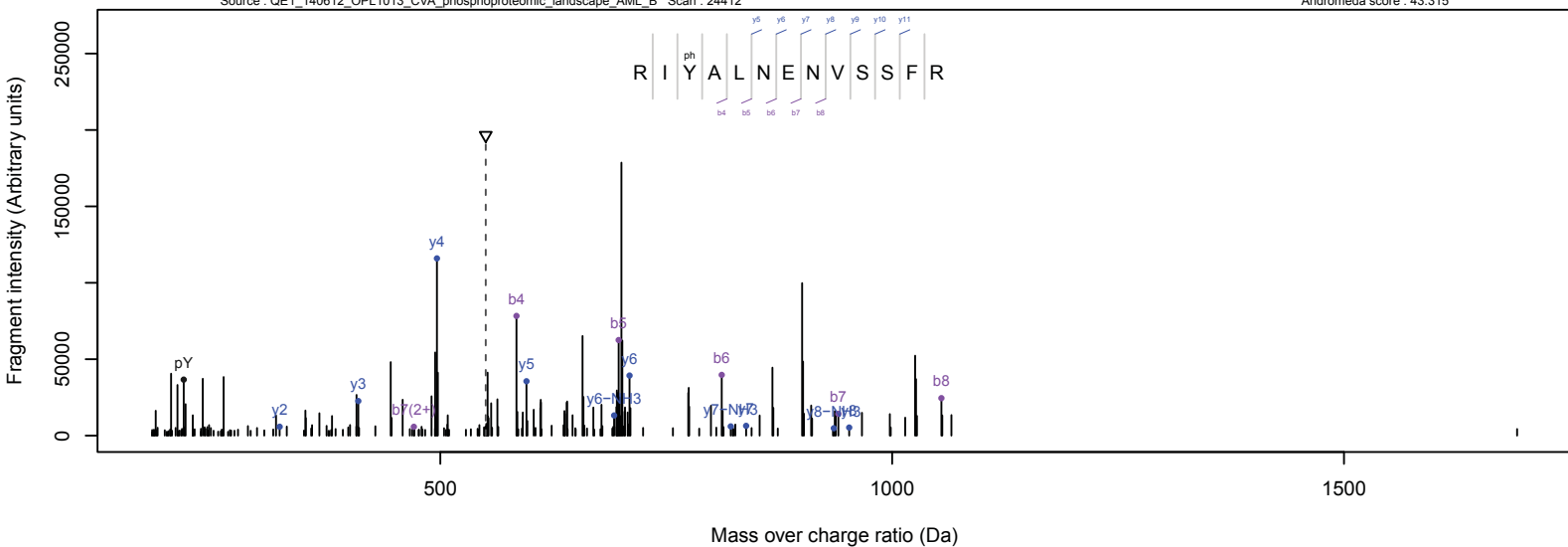
Andromeda score : 43.326



**RIY(ph)ALNENVSSFR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24412

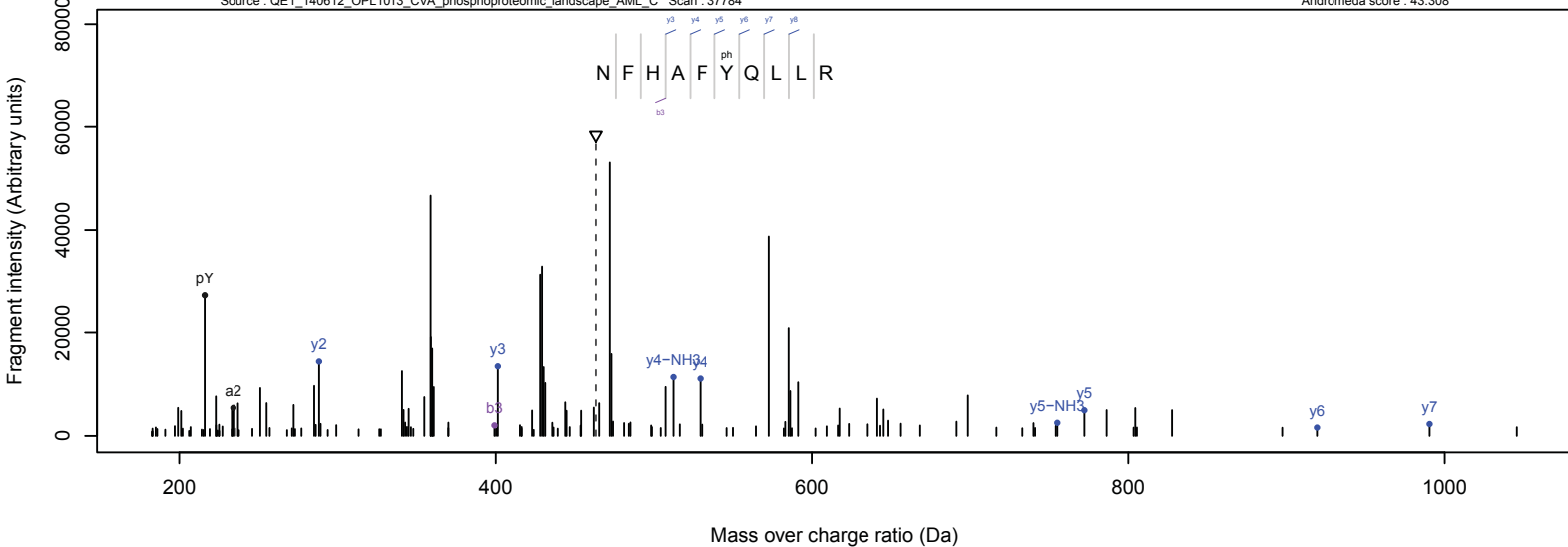
Andromeda score : 43.315



**NFHAFY(ph)QLLR-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37784

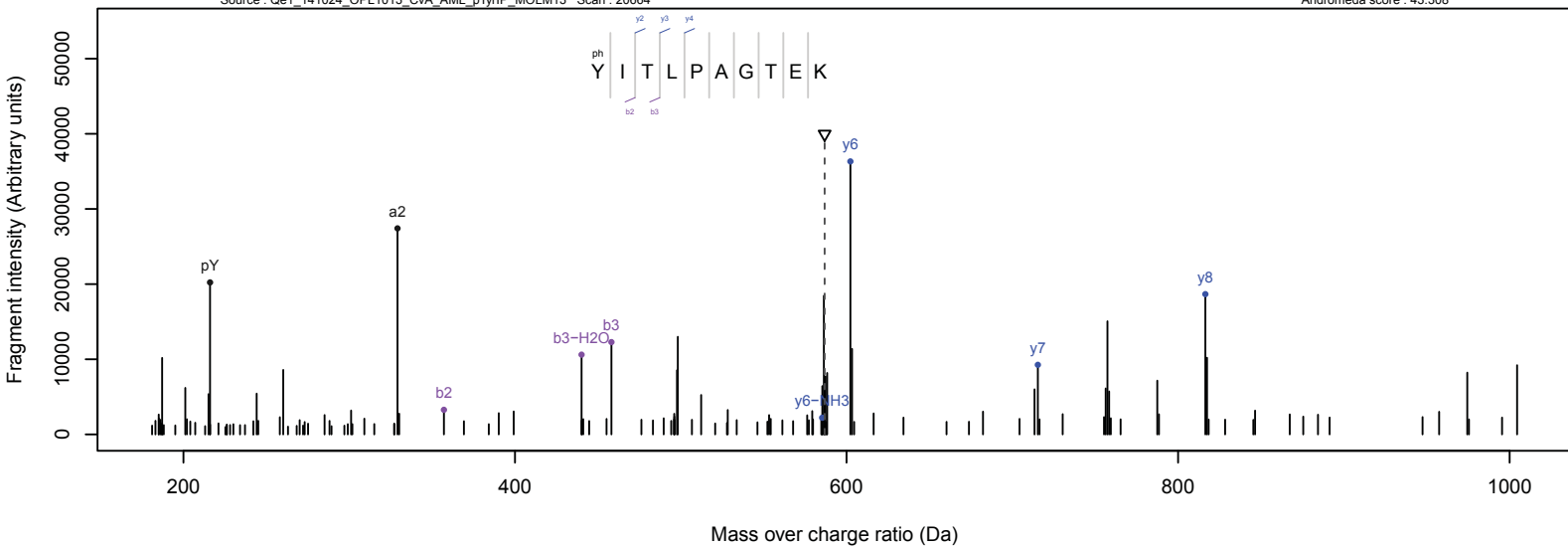
Andromeda score : 43.308



**Y(ph)ITLPAGTEK-2+**

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 20664

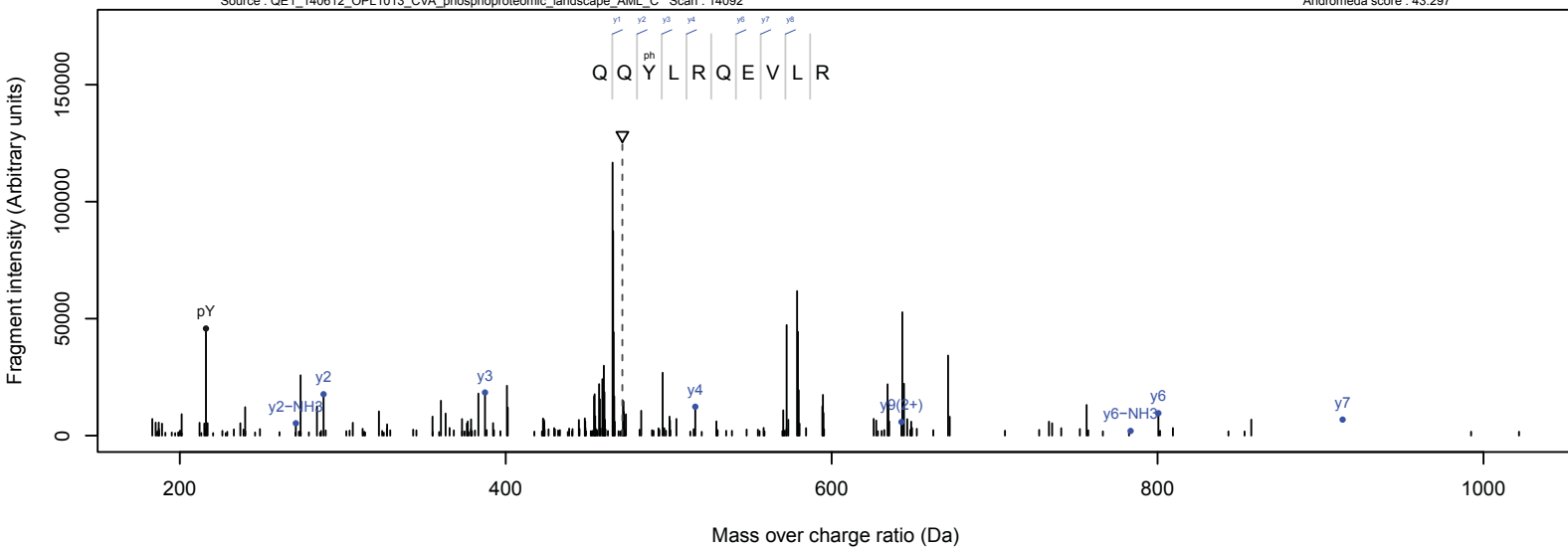
Andromeda score : 43.308



## QQY(ph)LRQEVLR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14092

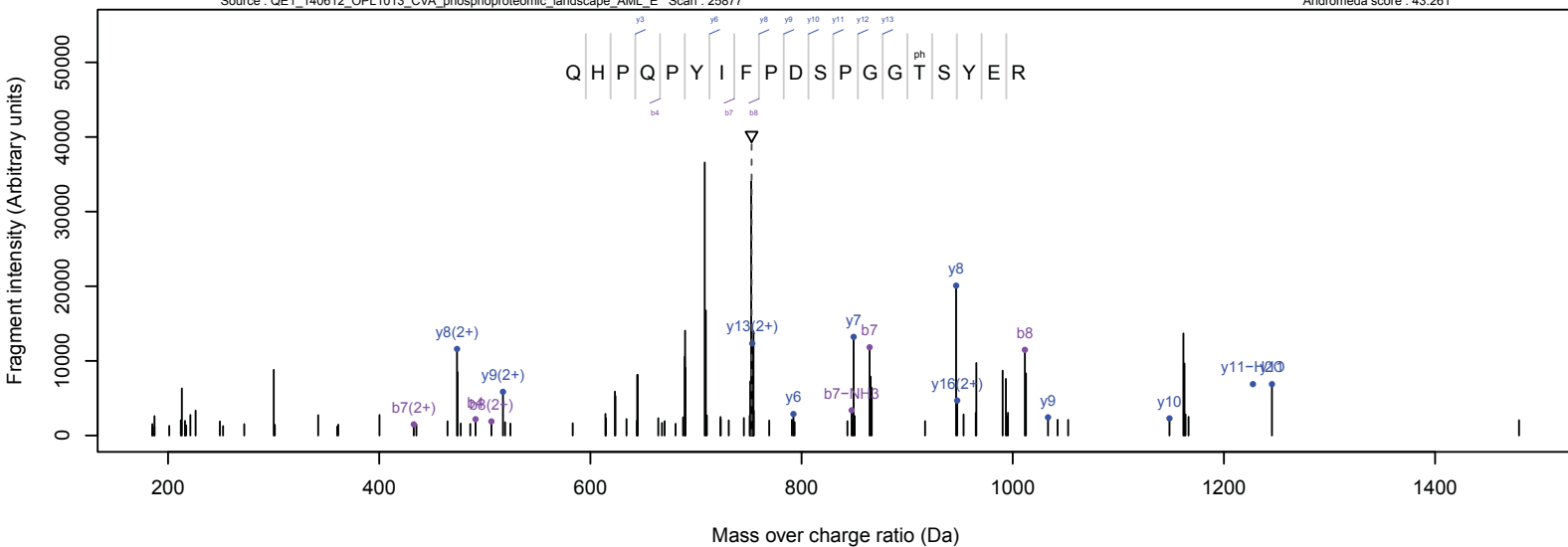
Andromeda score : 43.297



## QHPQPYIFPDSPPG(ph)TSYER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 25877

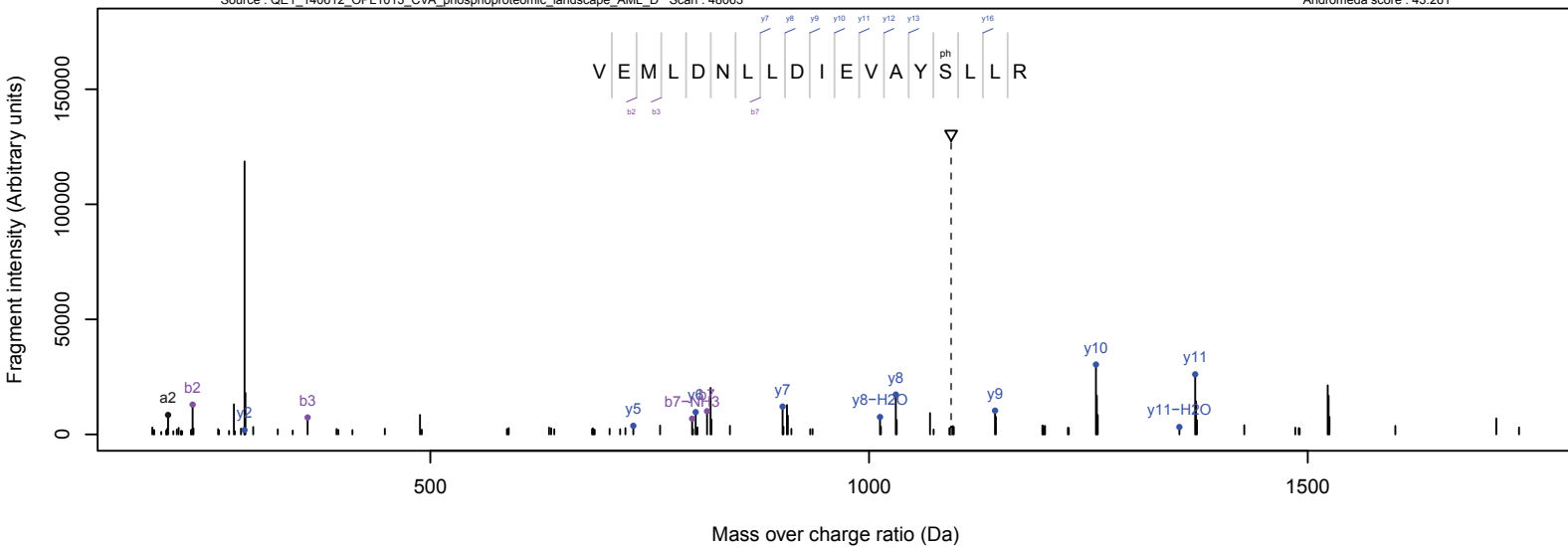
Andromeda score : 43.261



## VEMLDNLLDIEVAYS(ph)LLR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 48063

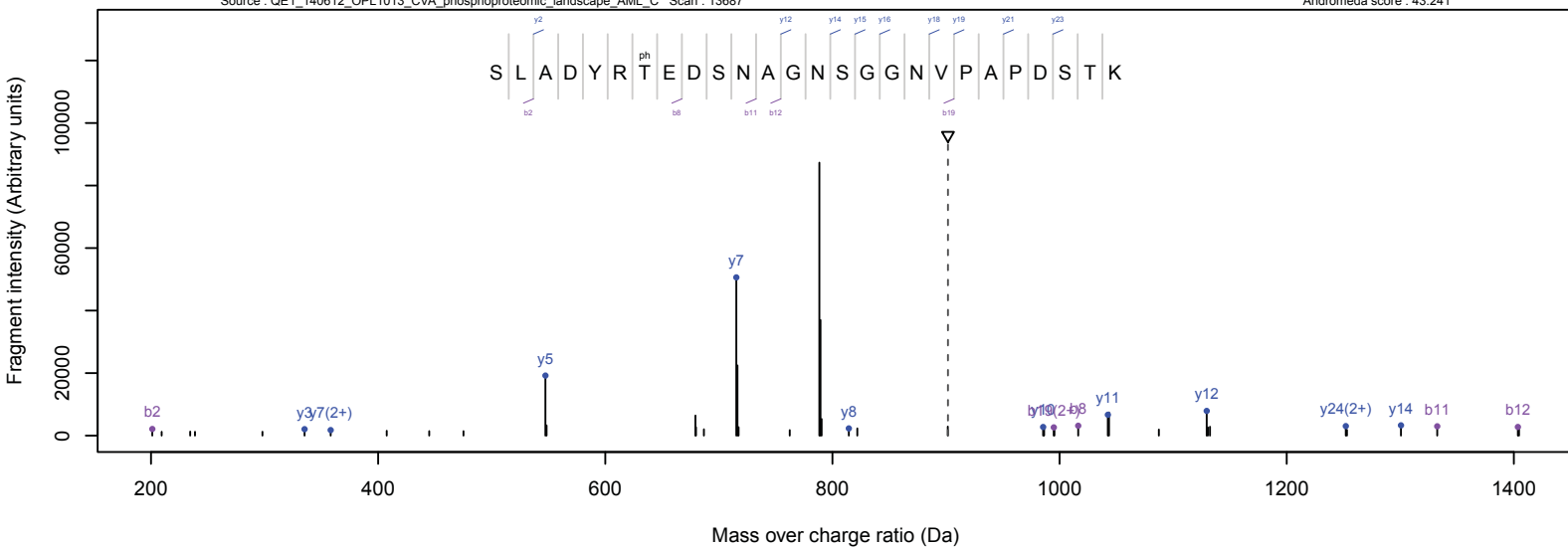
Andromeda score : 43.261



## SLADYRT(ph)EDSNAGNSGGNVPA PDSTK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13687

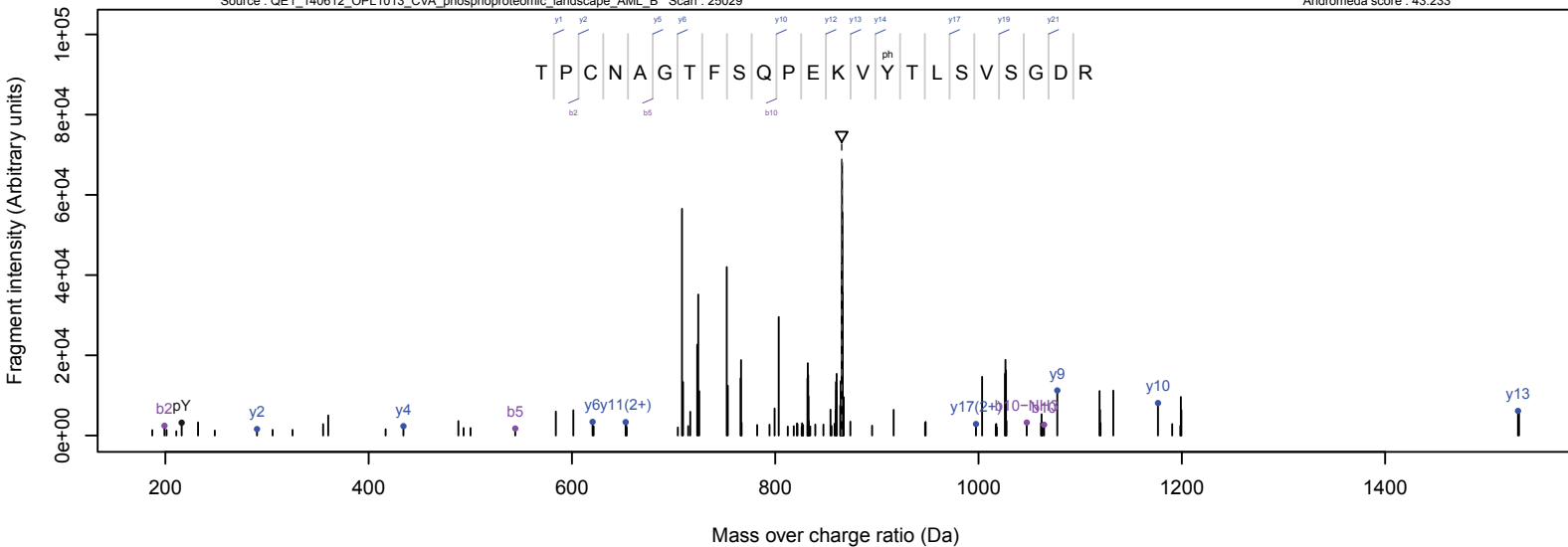
Andromeda score : 43.241



## TPCNAGTFSQPEKV(ph)YTLVSGDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25029

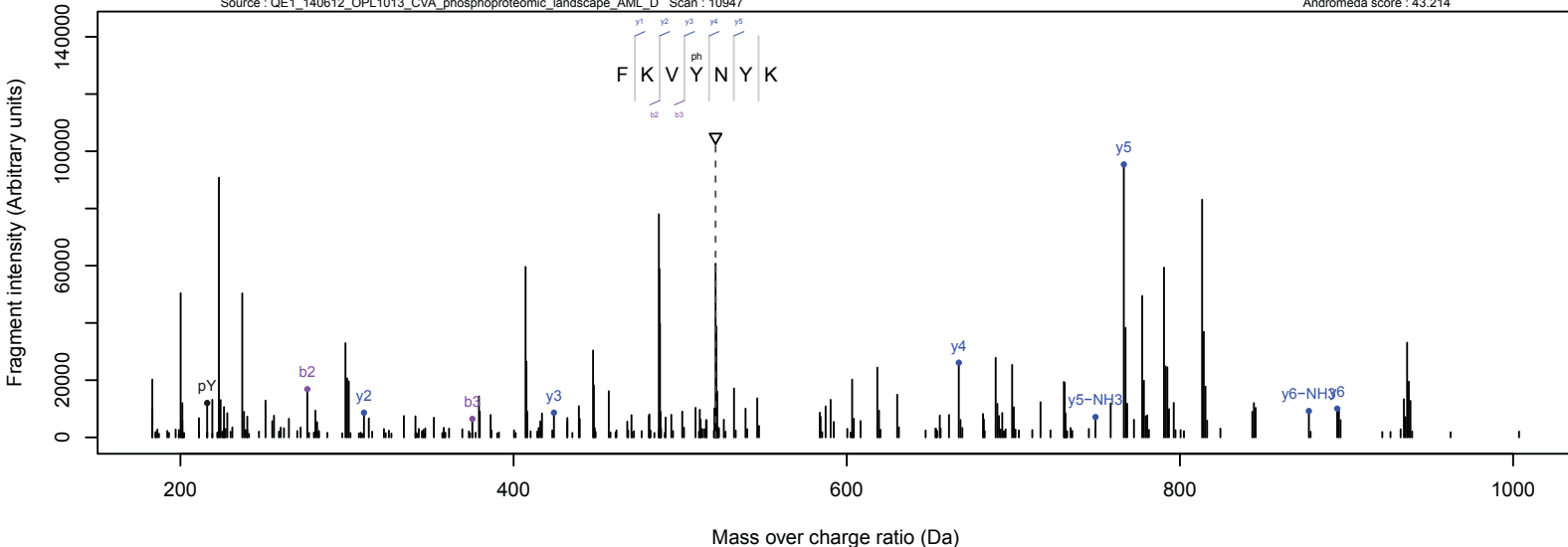
Andromeda score : 43.233



## FKVY(ph)NYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 10947

Andromeda score : 43.214

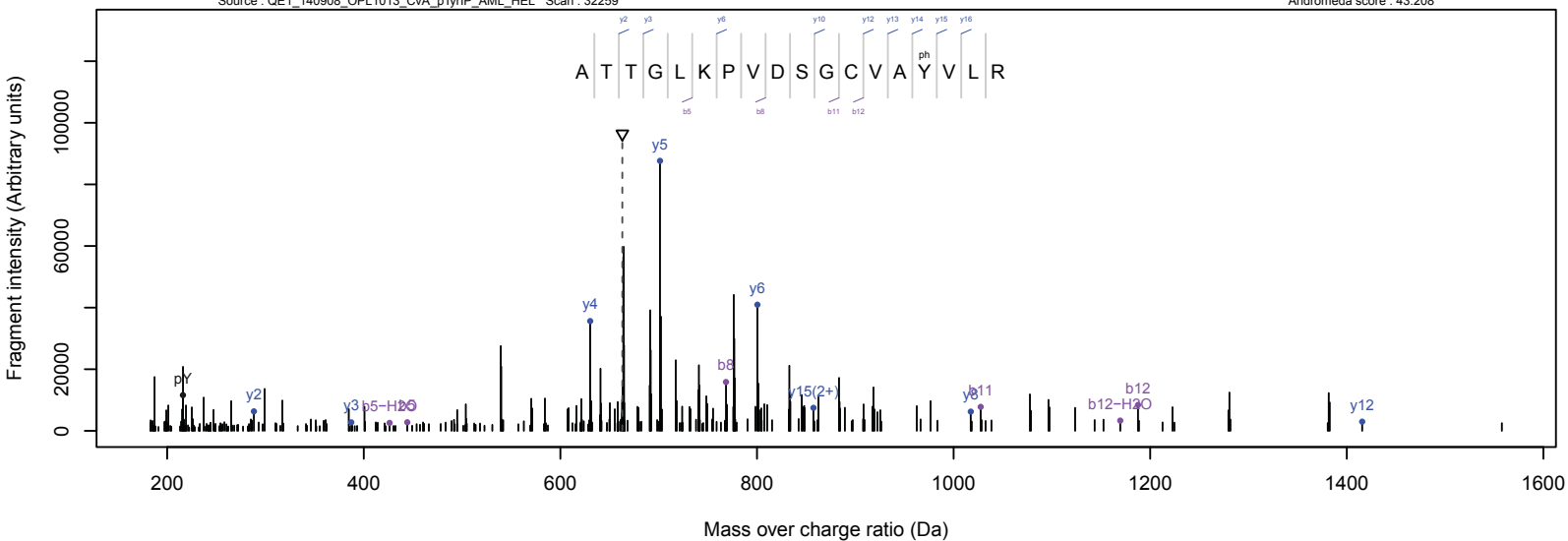




## ATTGLKPVDSGCVAY(ph)VLR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 32259

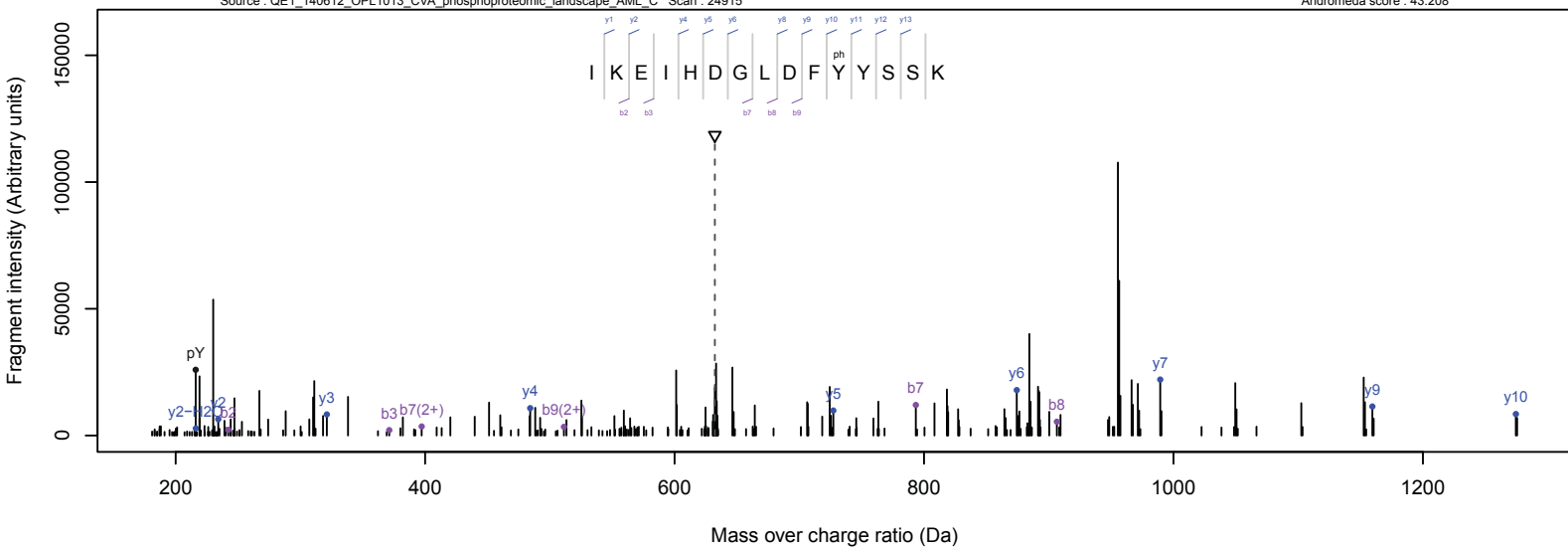
Andromeda score : 43.208



## IKEIHDGLDFYSSK-3+

Source : QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan : 24915

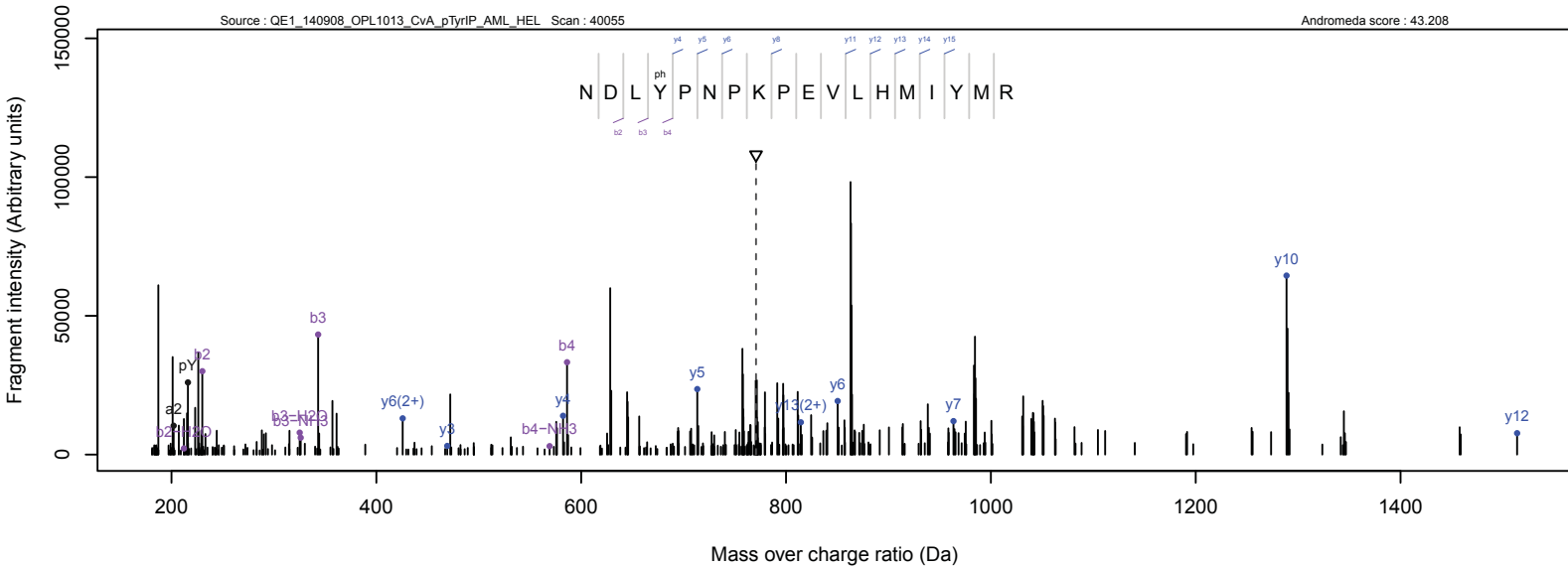
Andromeda score : 43.208

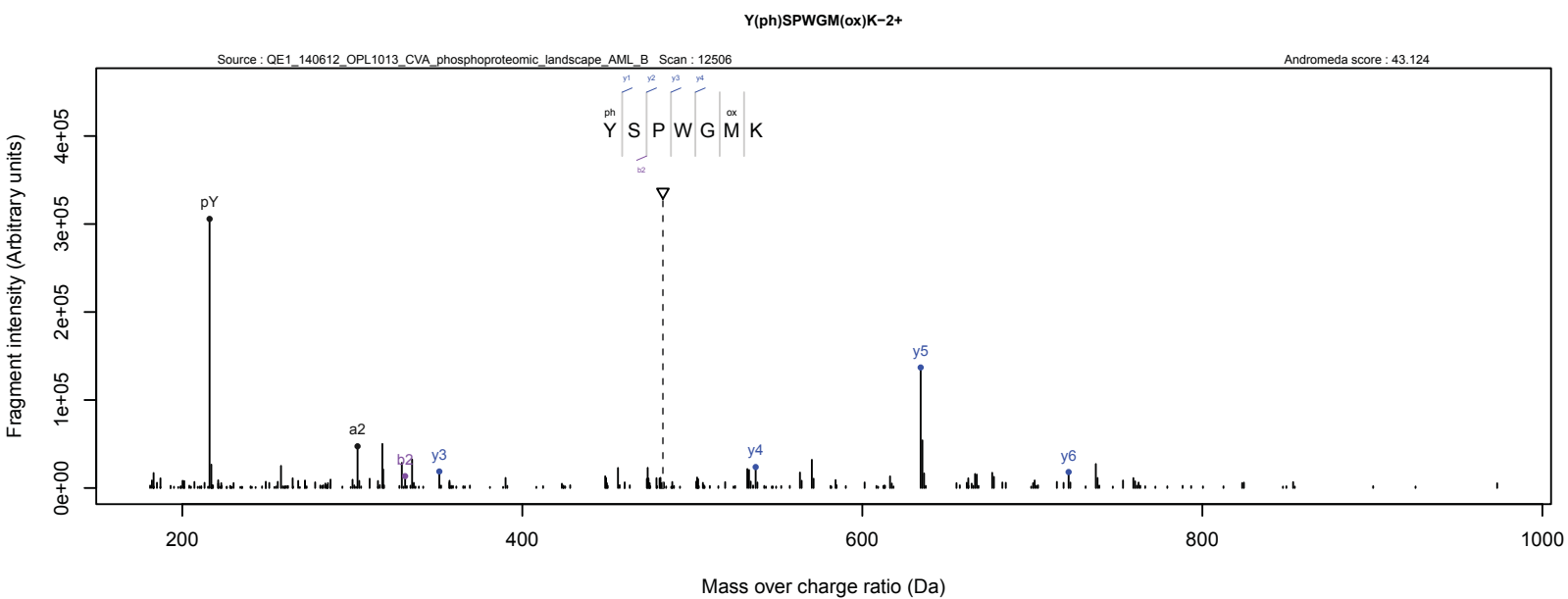
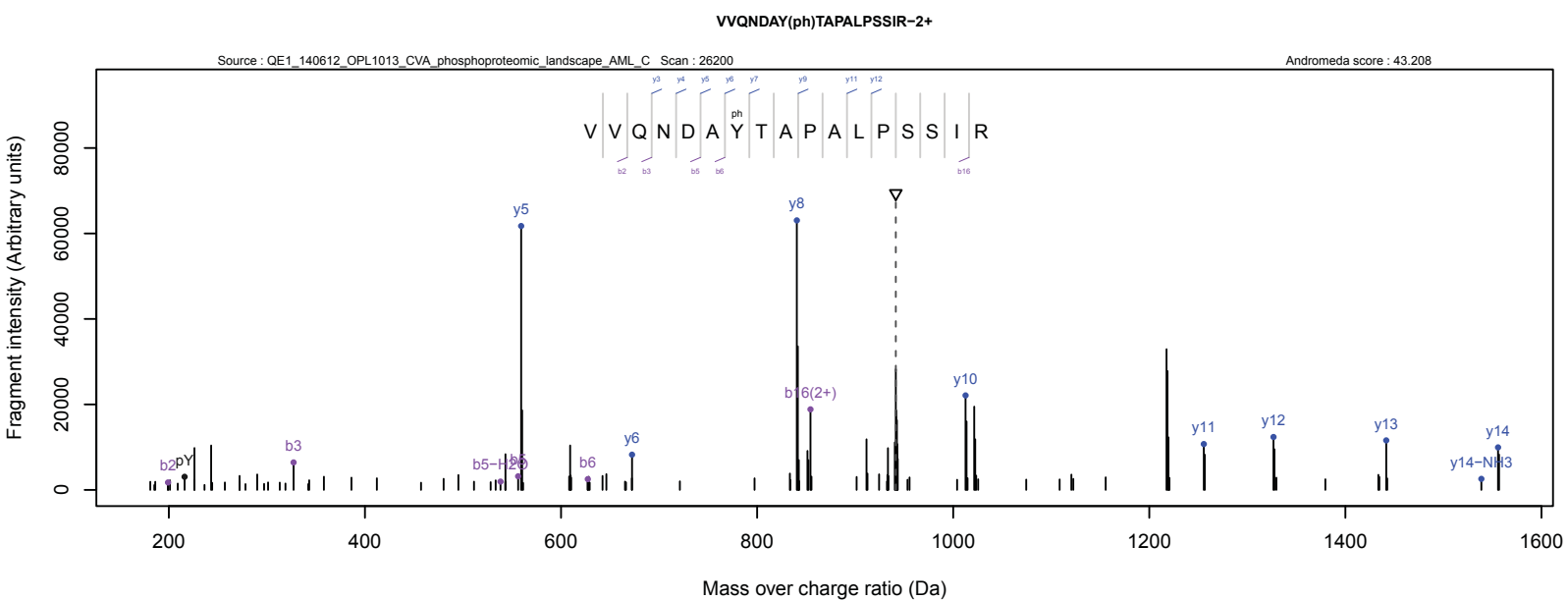
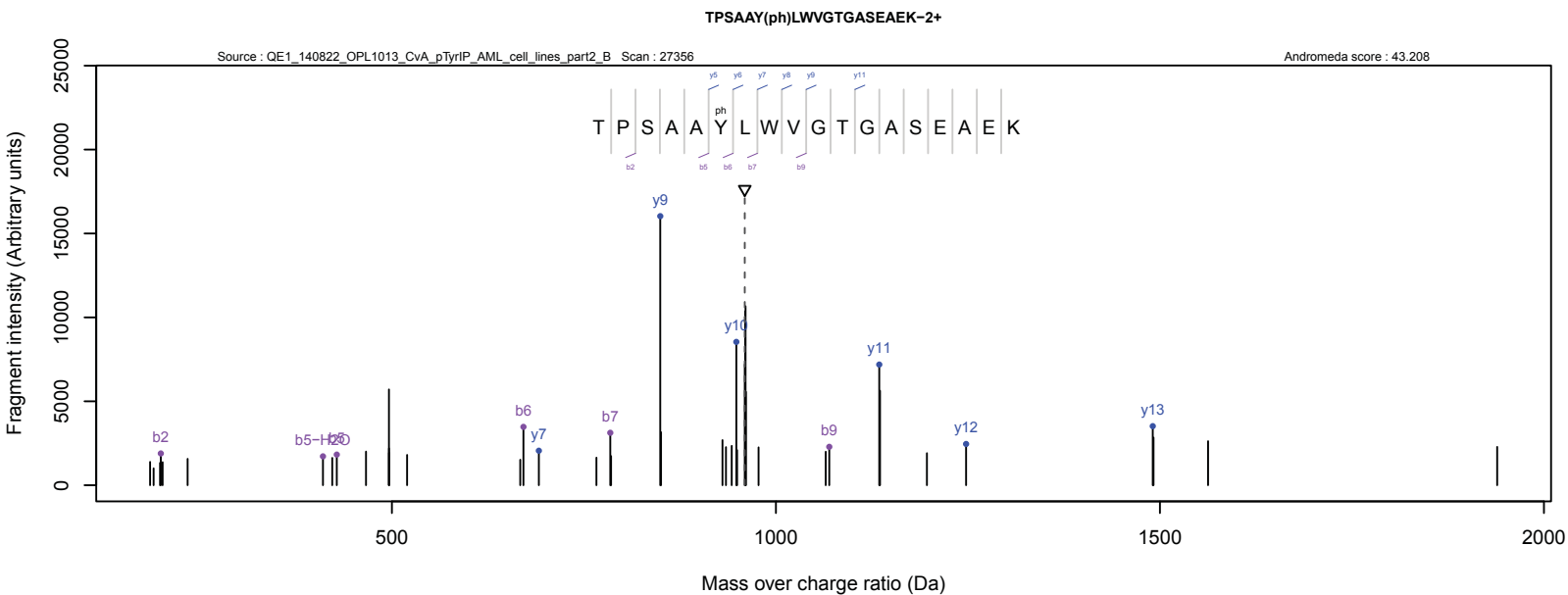


## NDLY(ph)PNPKPEVLHMIYMR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 40055

Andromeda score : 43.208

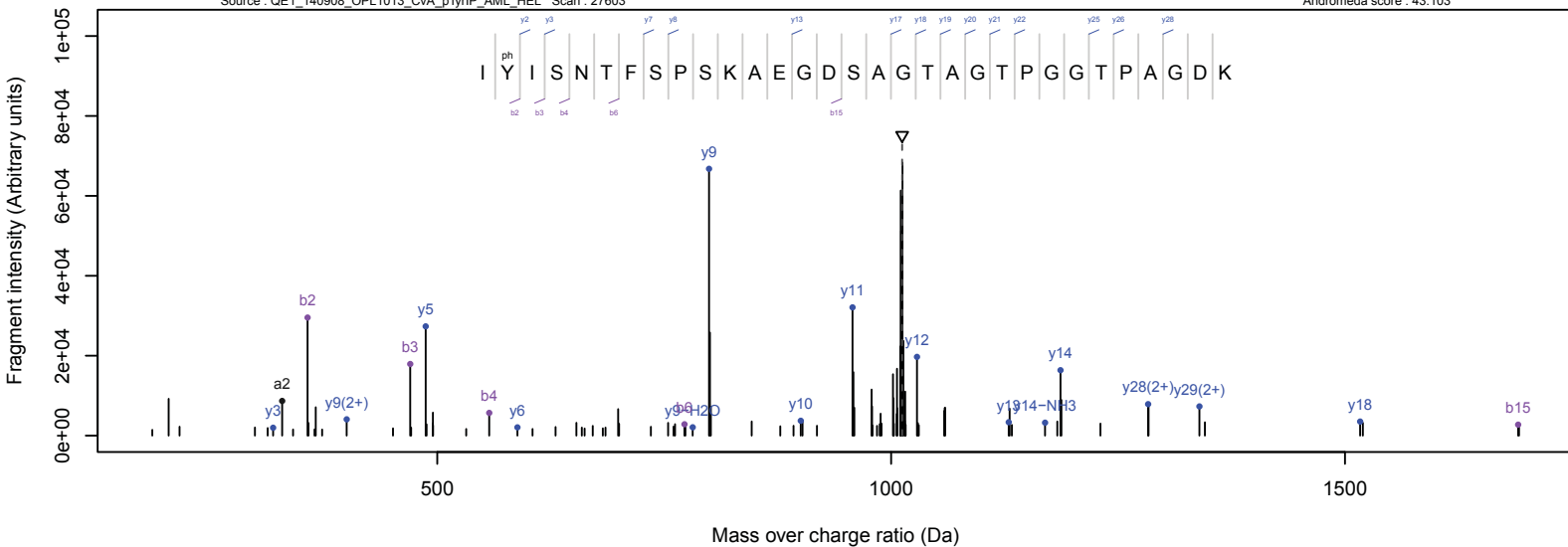




IY(ph)ISNTFSPSKAEGDSAGTAPGGTPAGDK-3+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 27603

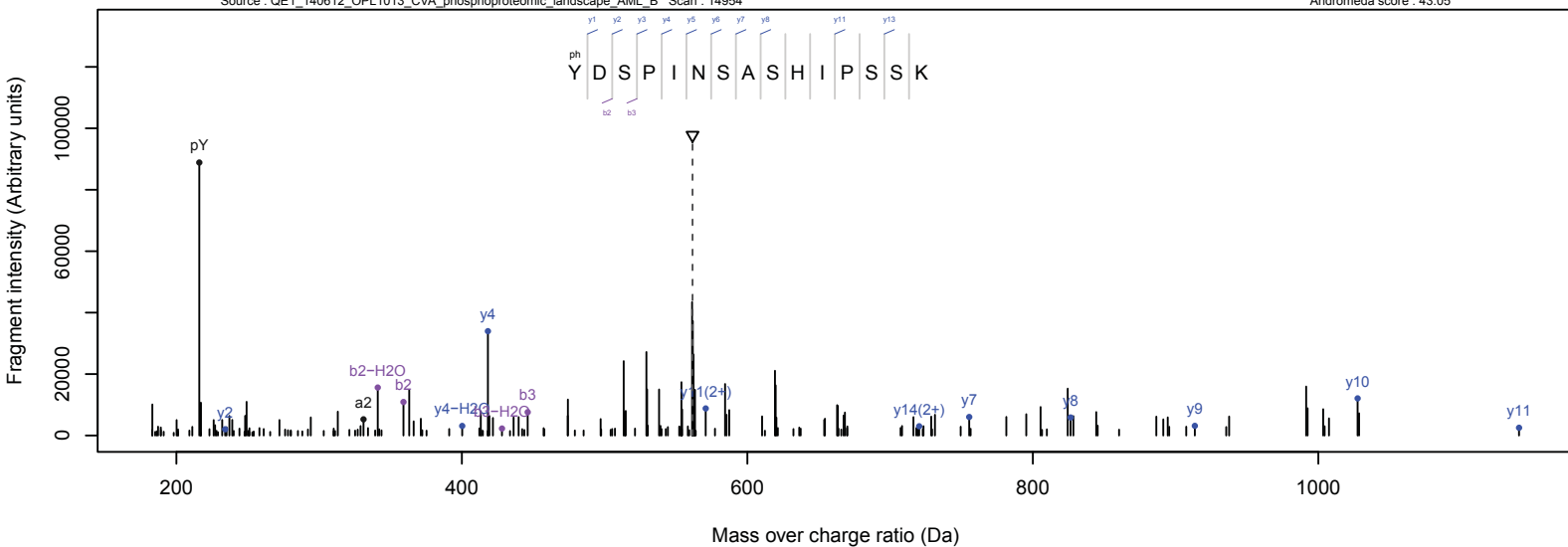
Andromeda score : 43.103



Y(ph)DSPINSASHIPSSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 14954

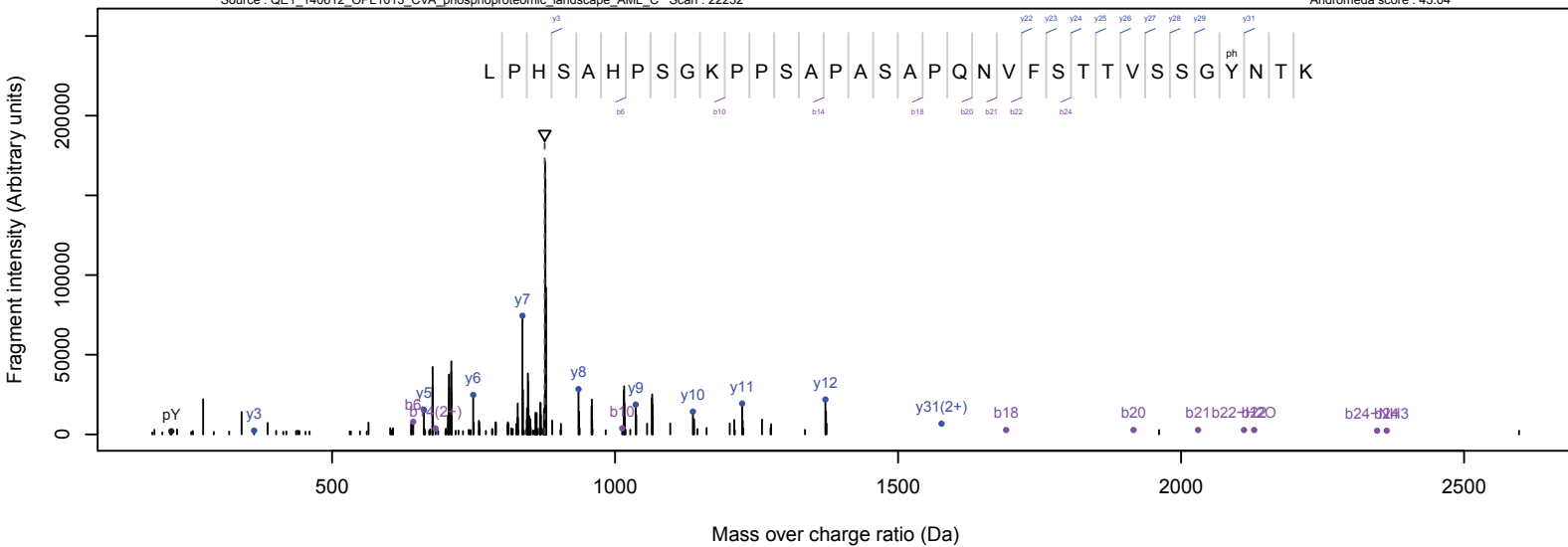
Andromeda score : 43.05



LPHSAHPGKPPSAPASAPQNVFSTTVSSGY(ph)NTK-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22232

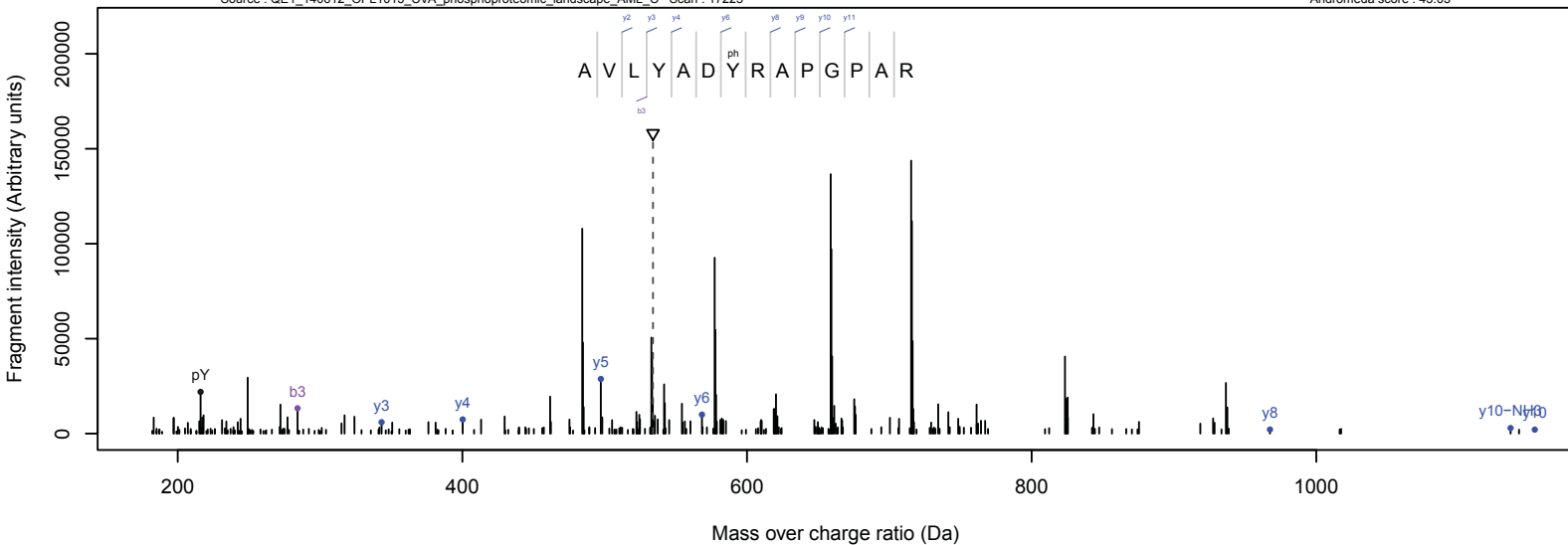
Andromeda score : 43.04



## AVLYADY(ph)RAPGAR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 17225

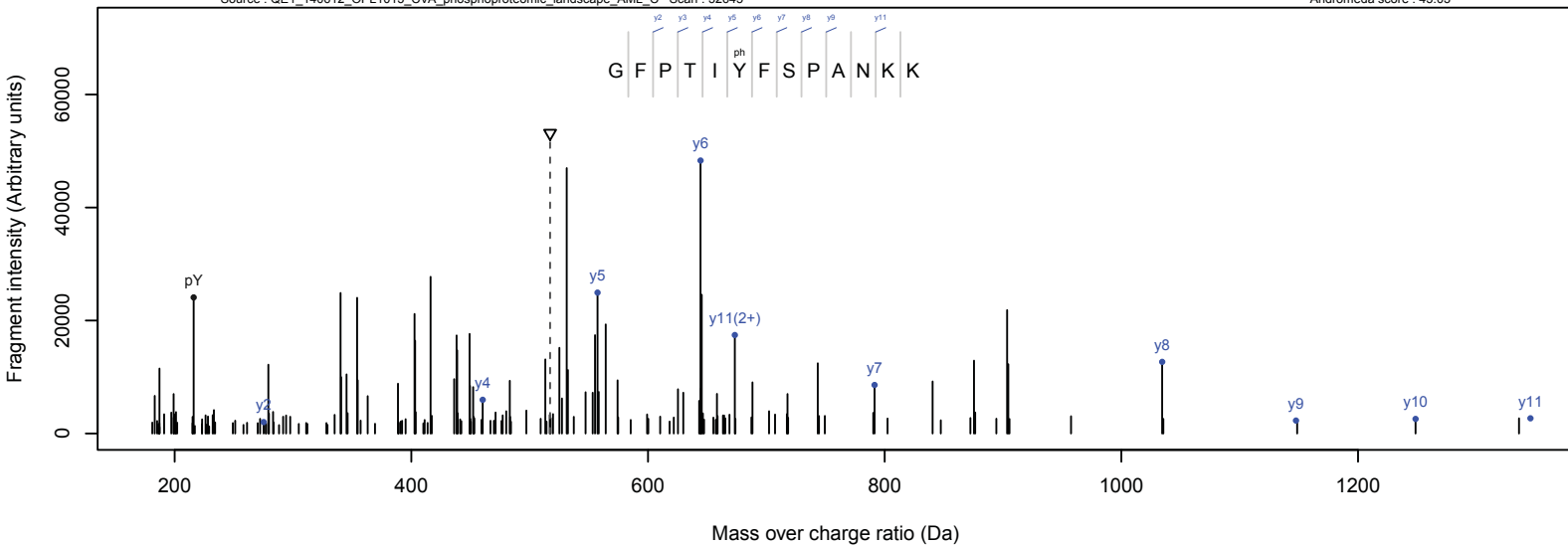
Andromeda score: 43.03



## GFPTIY(ph)FSPANKK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 32645

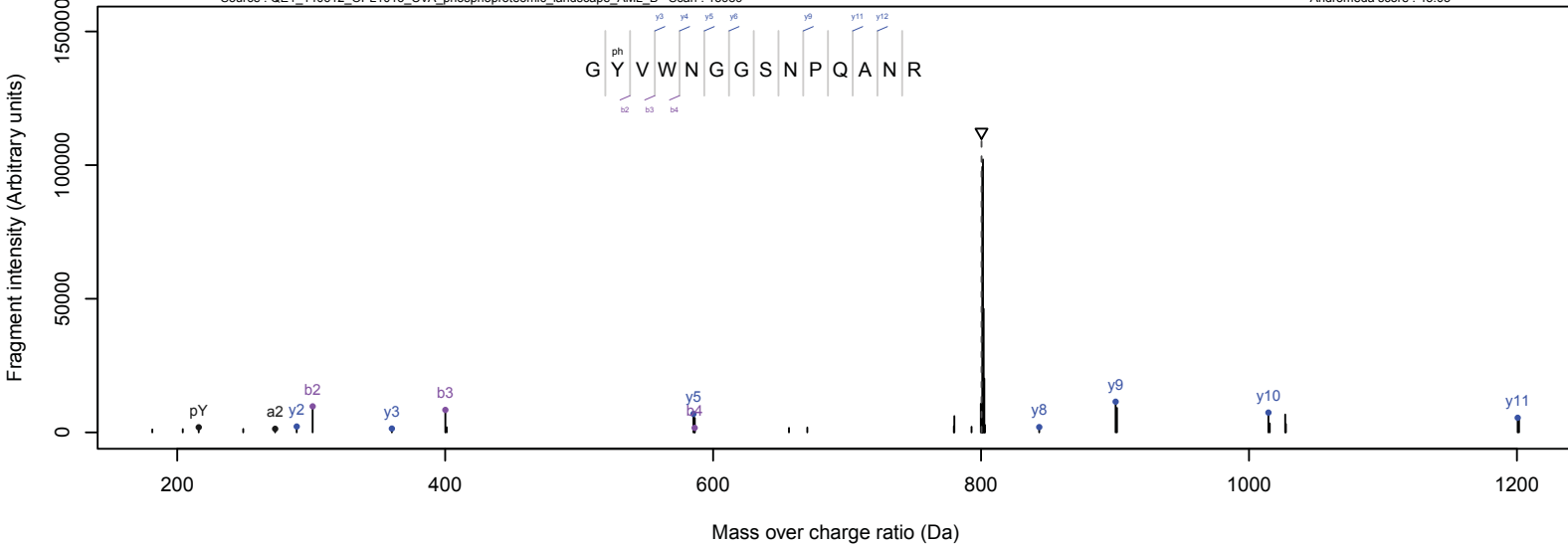
Andromeda score: 43.03



## GY(ph)VWNGSNPQANR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 13989

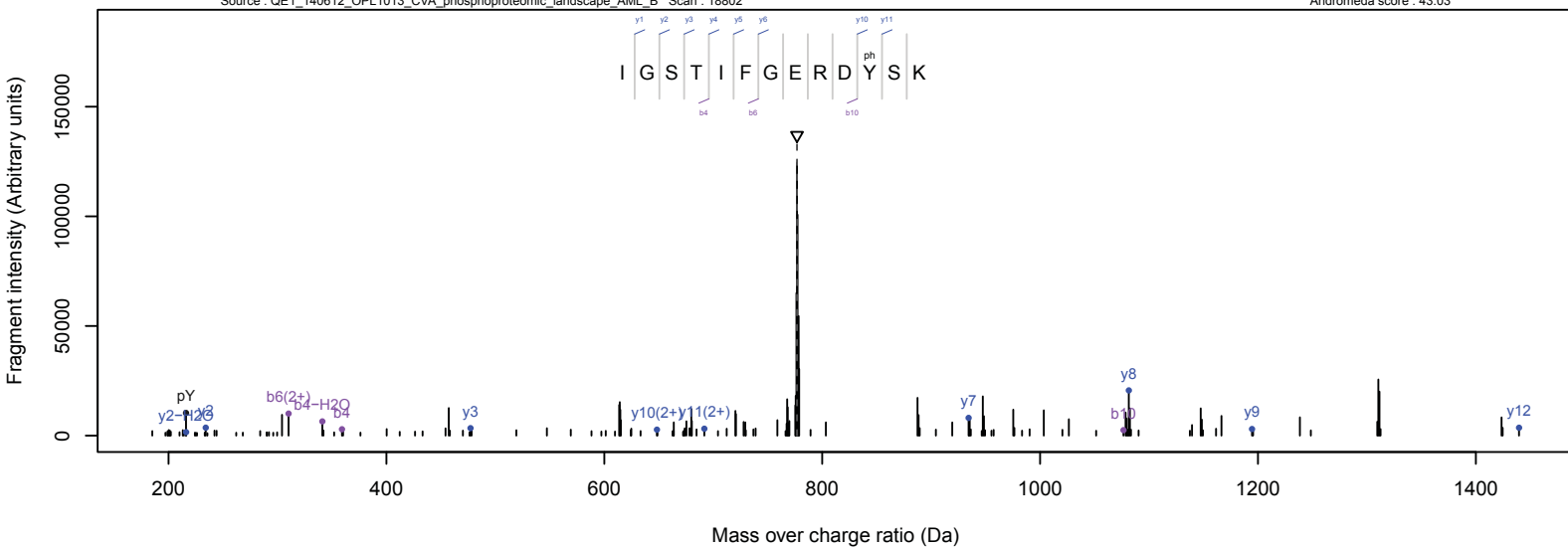
Andromeda score: 43.03



## IGSTIFGERDY(ph)SK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18802

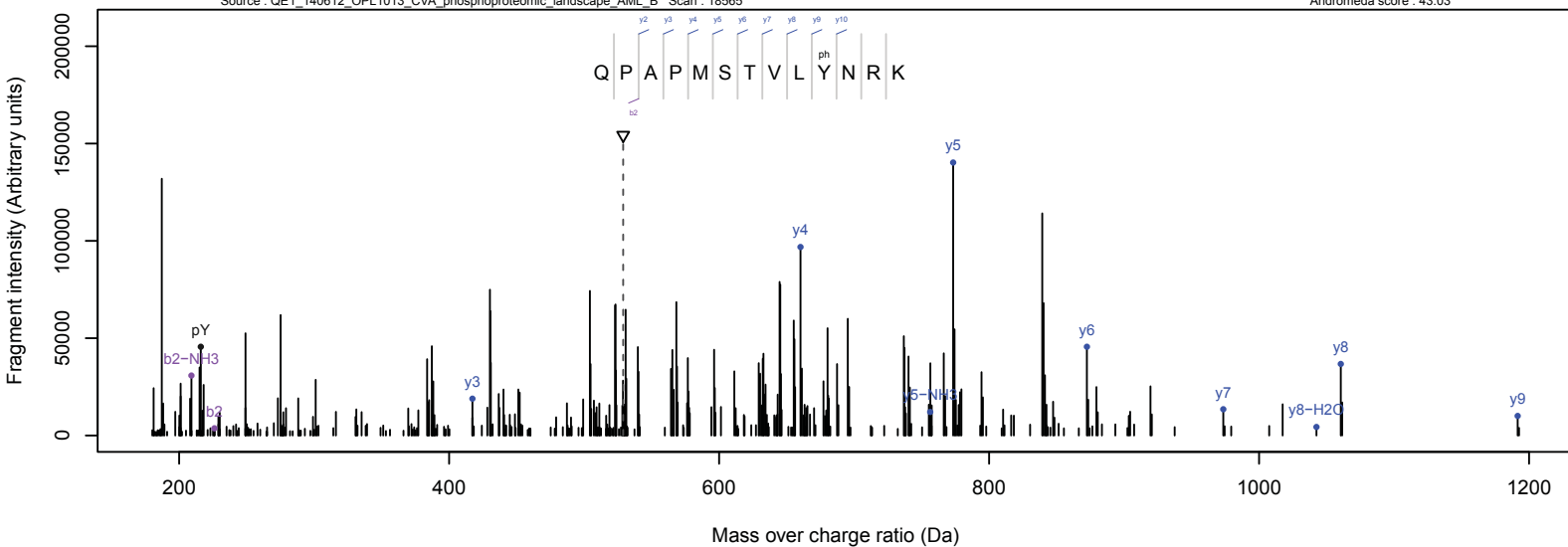
Andromeda score: 43.03



## QPAPMSTVLY(ph)NRK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18565

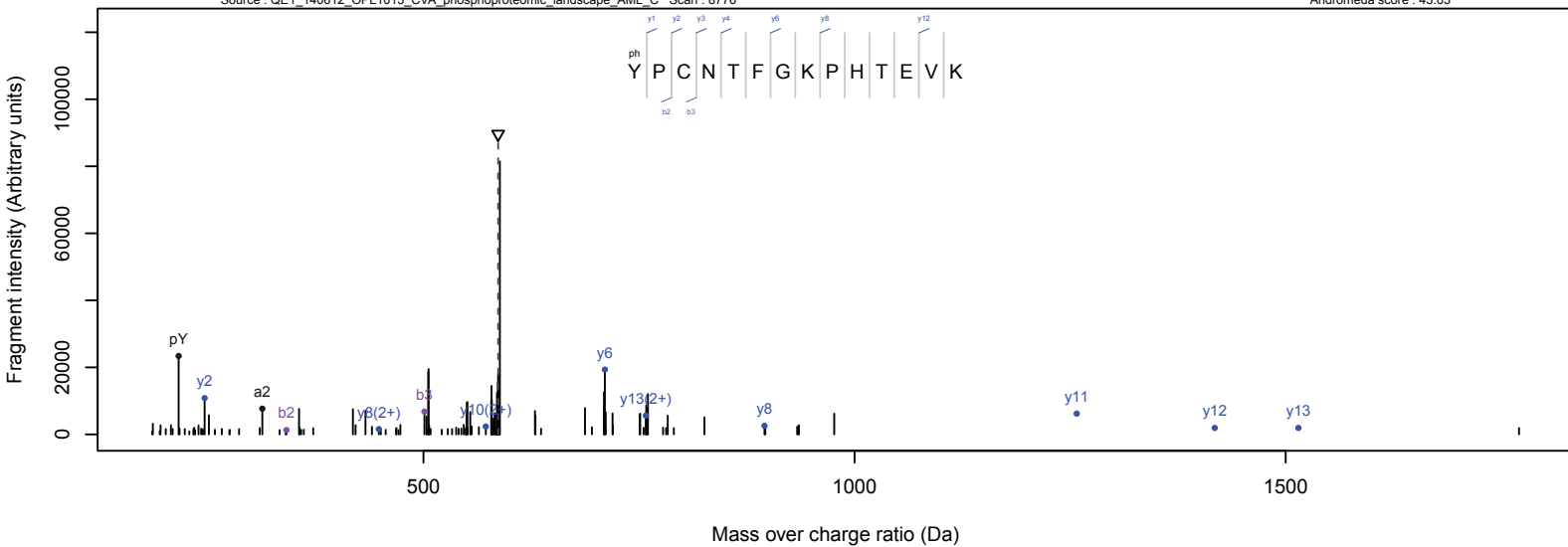
Andromeda score: 43.03



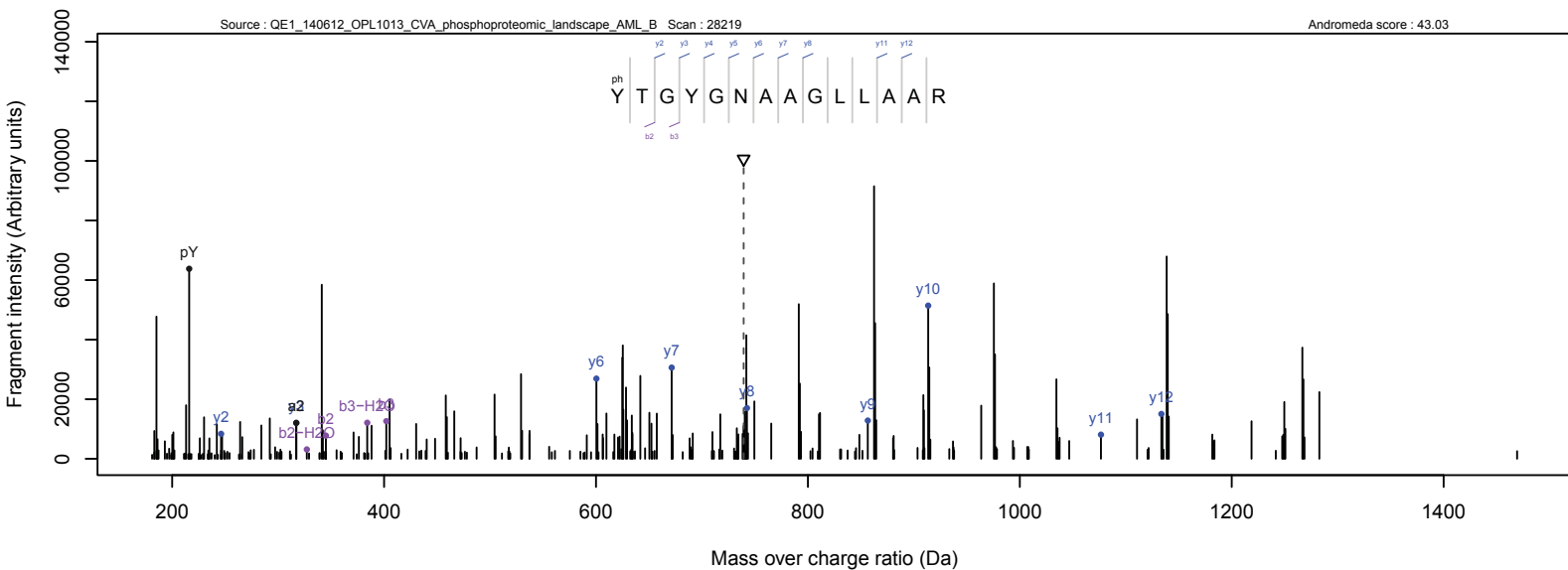
## Y(ph)PCNTFGKPHTEVK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 8776

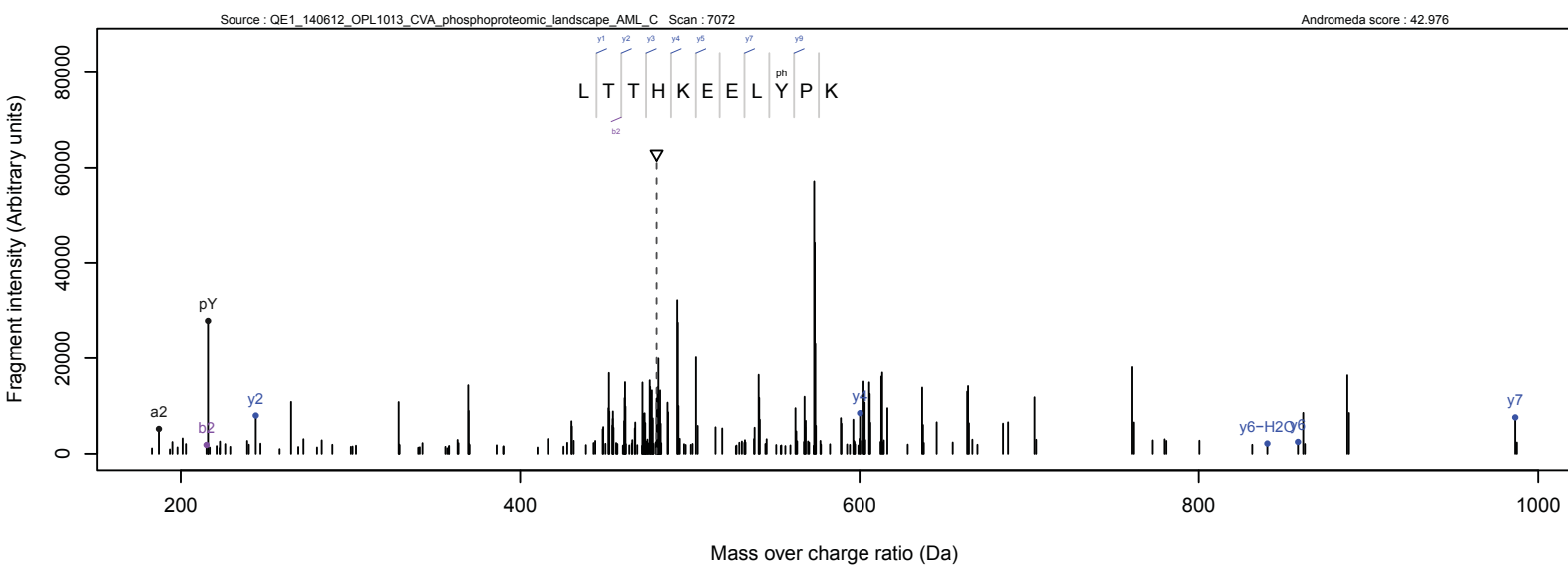
Andromeda score: 43.03



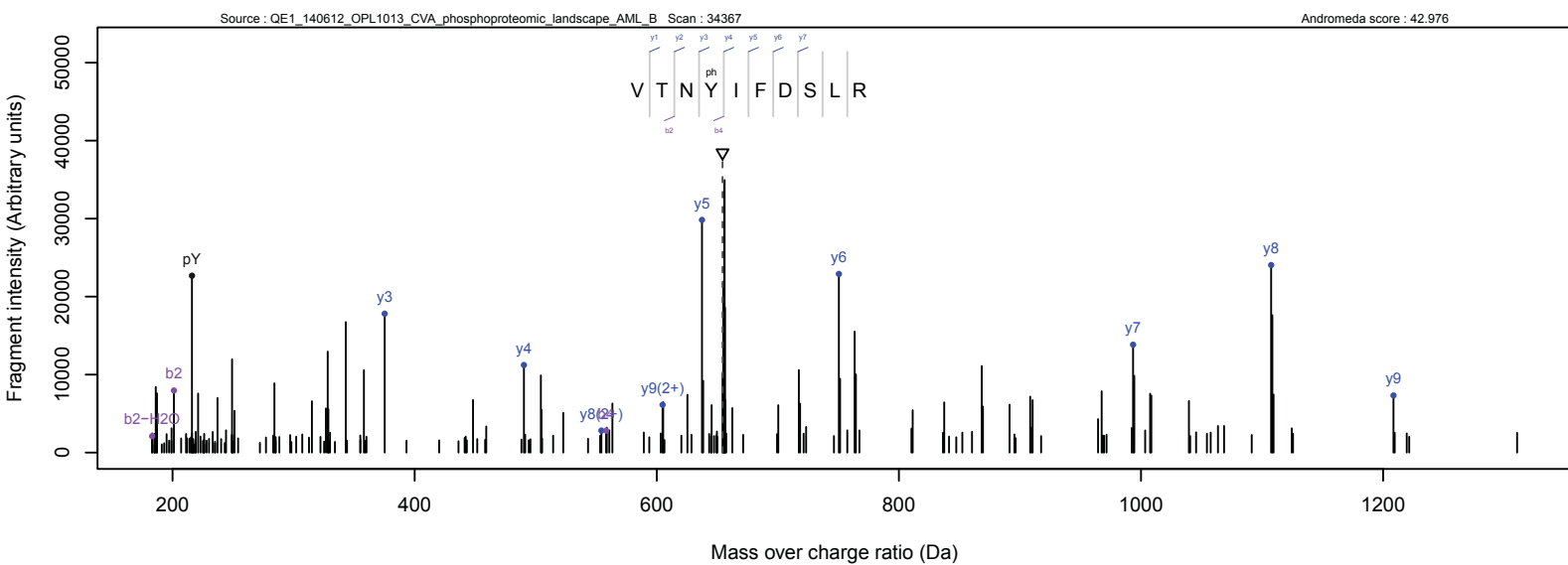
## Y(ph)TGYNAAGLLAAR-2+



## LTTHKEELY(ph)PK-3+



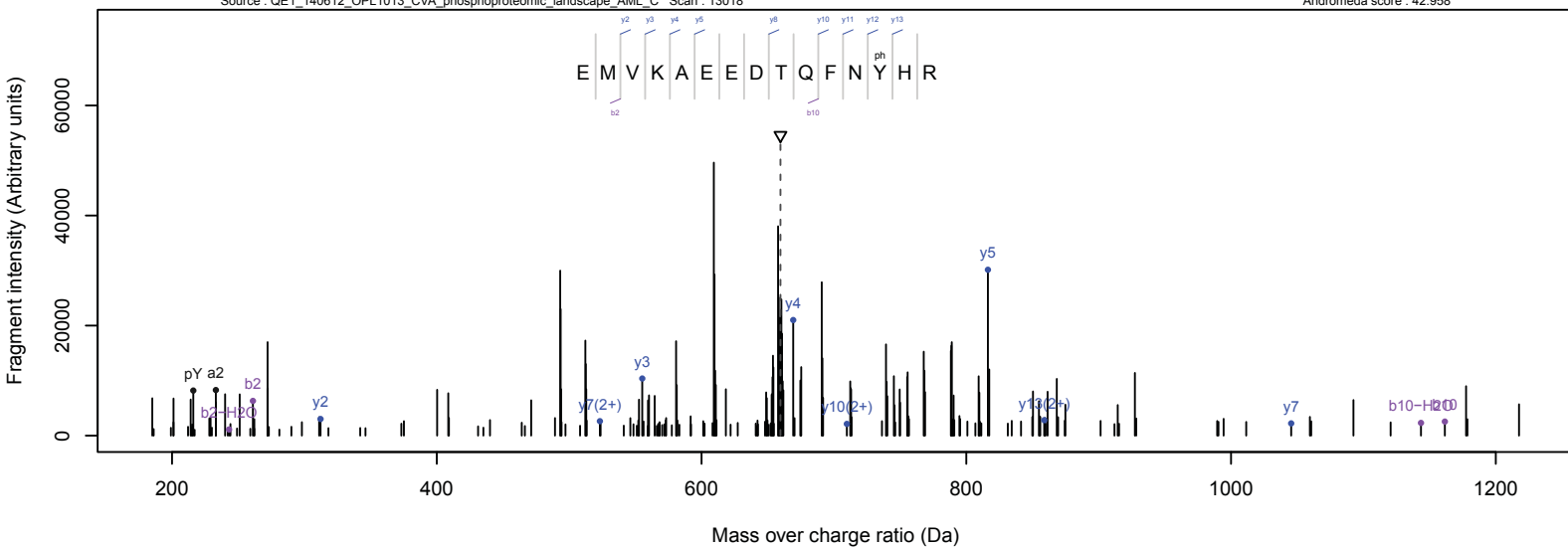
## VTNY(ph)IFDSLRL-2+



## EMVKAEDTQFNYPHHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 13018

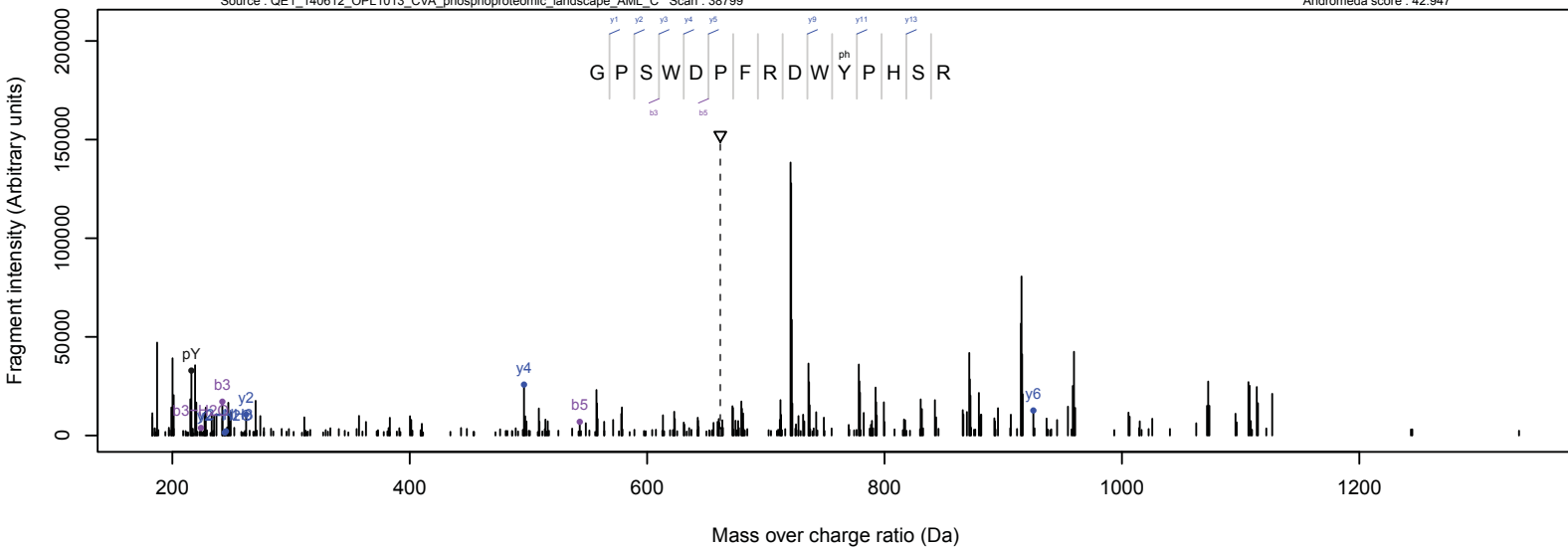
Andromeda score : 42.958



## GPSWDPFRDWYPHSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 38799

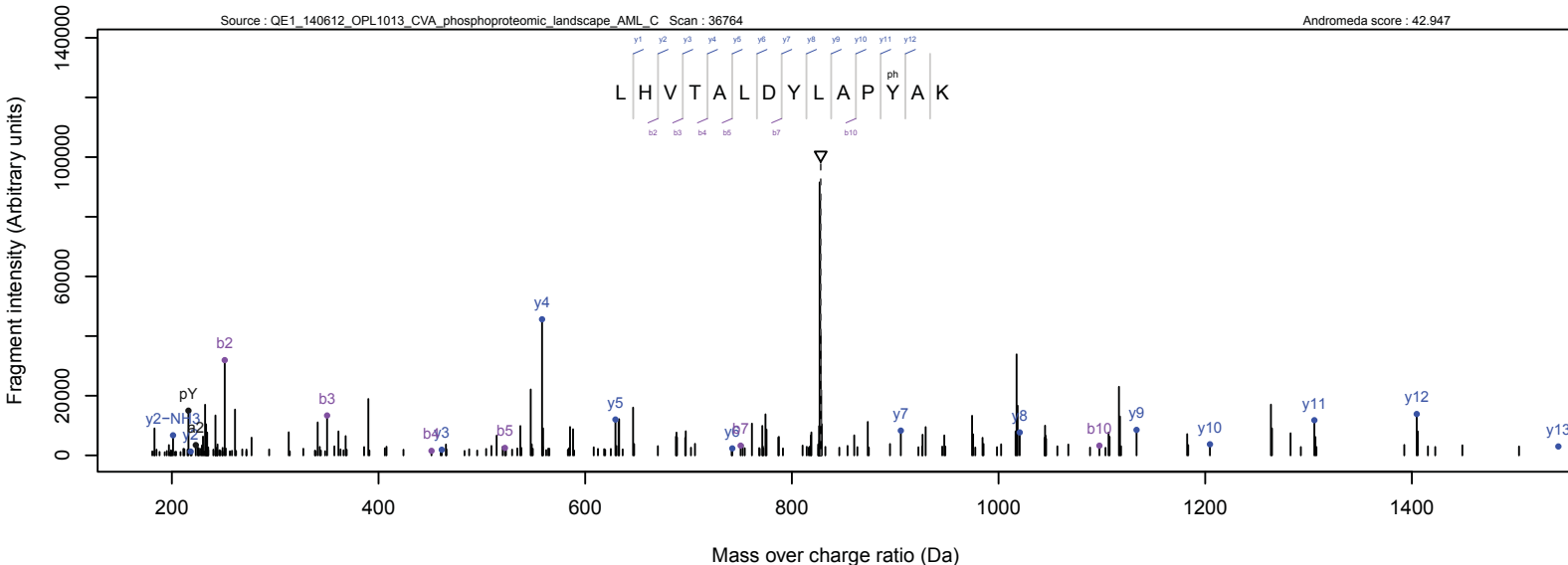
Andromeda score : 42.947



## LHVTALDYLAOPYAK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 36764

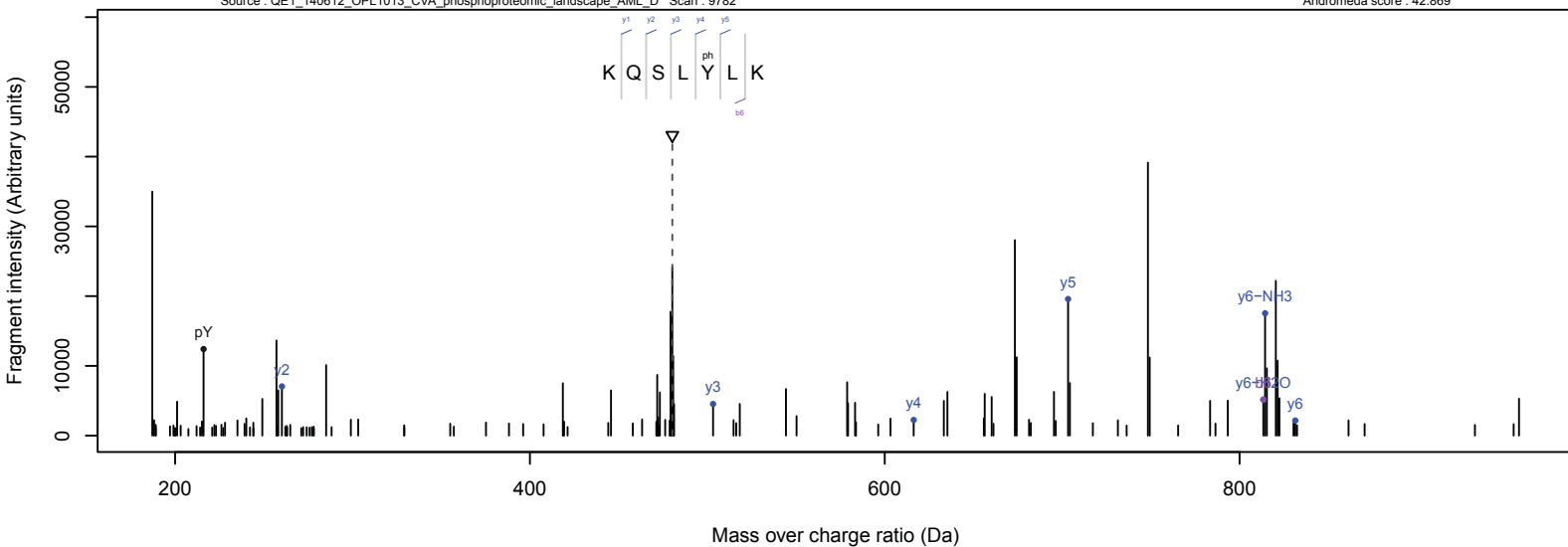
Andromeda score : 42.947



## KQSLY(ph)LK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 9782

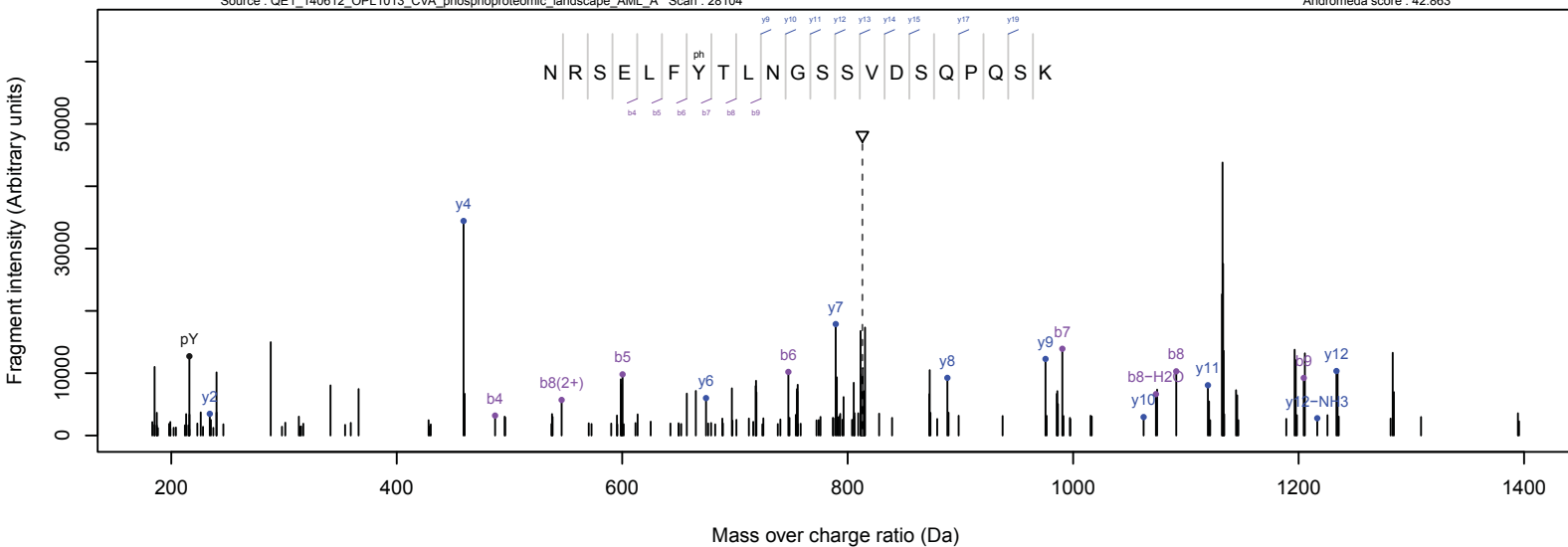
Andromeda score : 42.869



## NRSELY(ph)TLNGSSVDSQPQSK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 28104

Andromeda score : 42.863

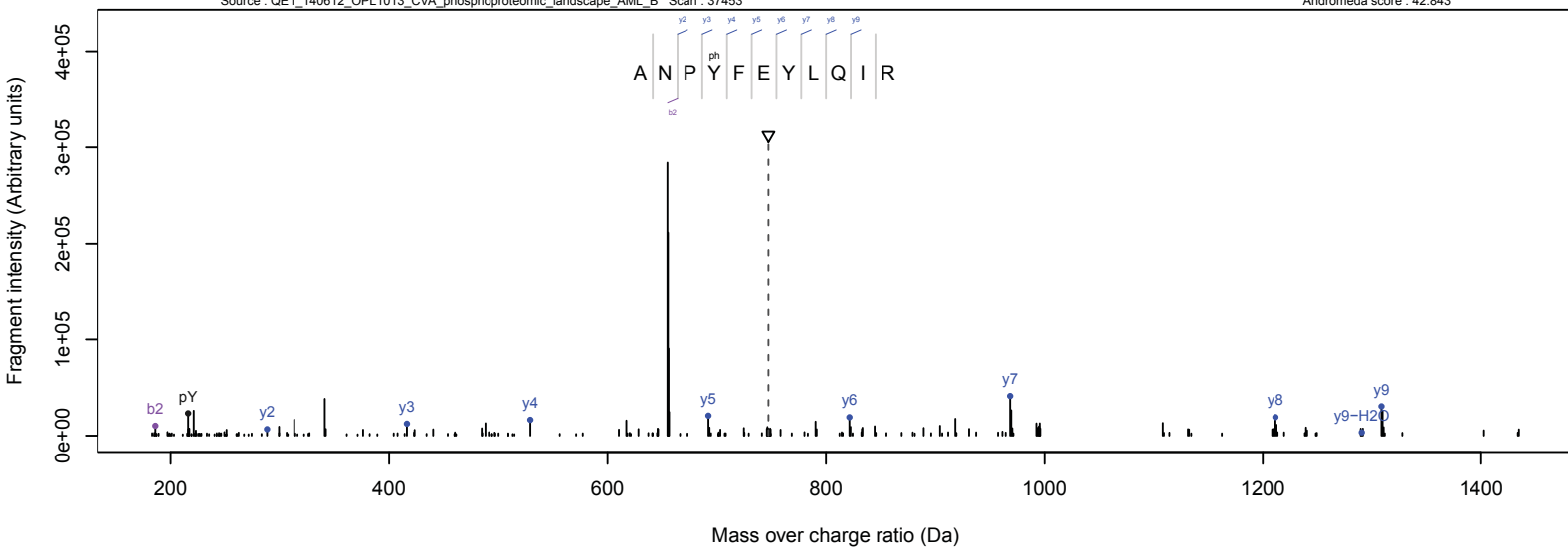




## ANPY(ph)FEYLQIR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 37453

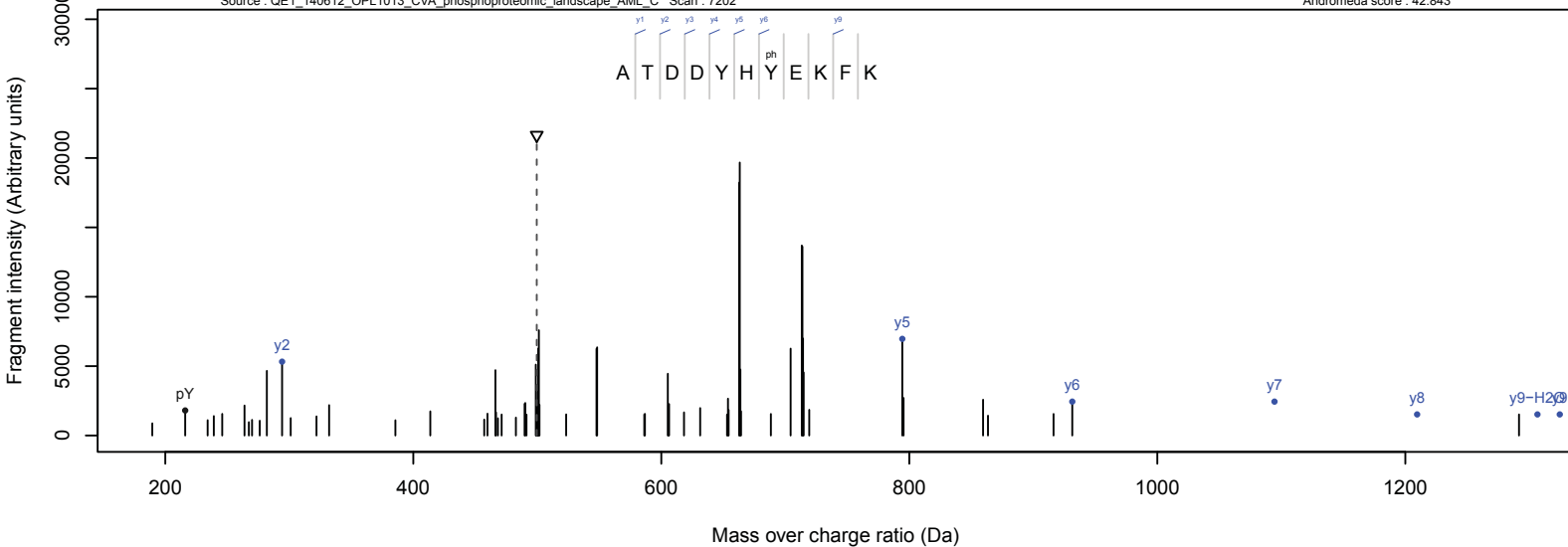
Andromeda score : 42.843



## ATDDYH(ph)EKFK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7202

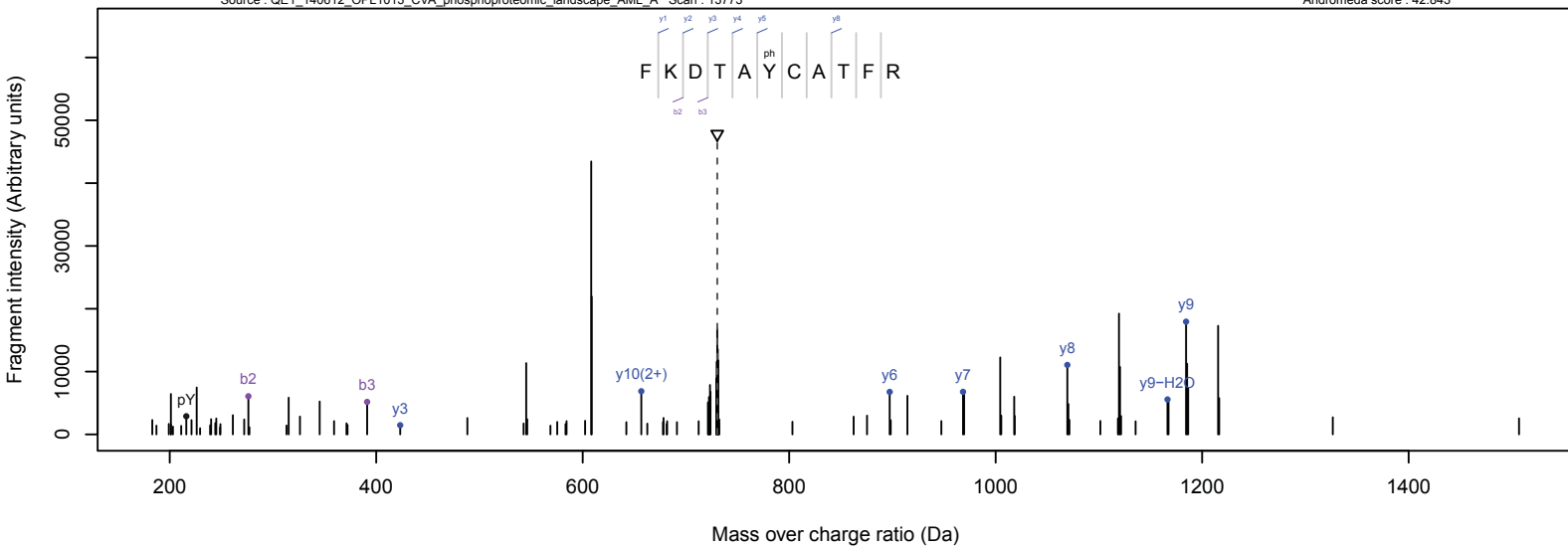
Andromeda score : 42.843



## FKDTAY(ph)CATFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 13773

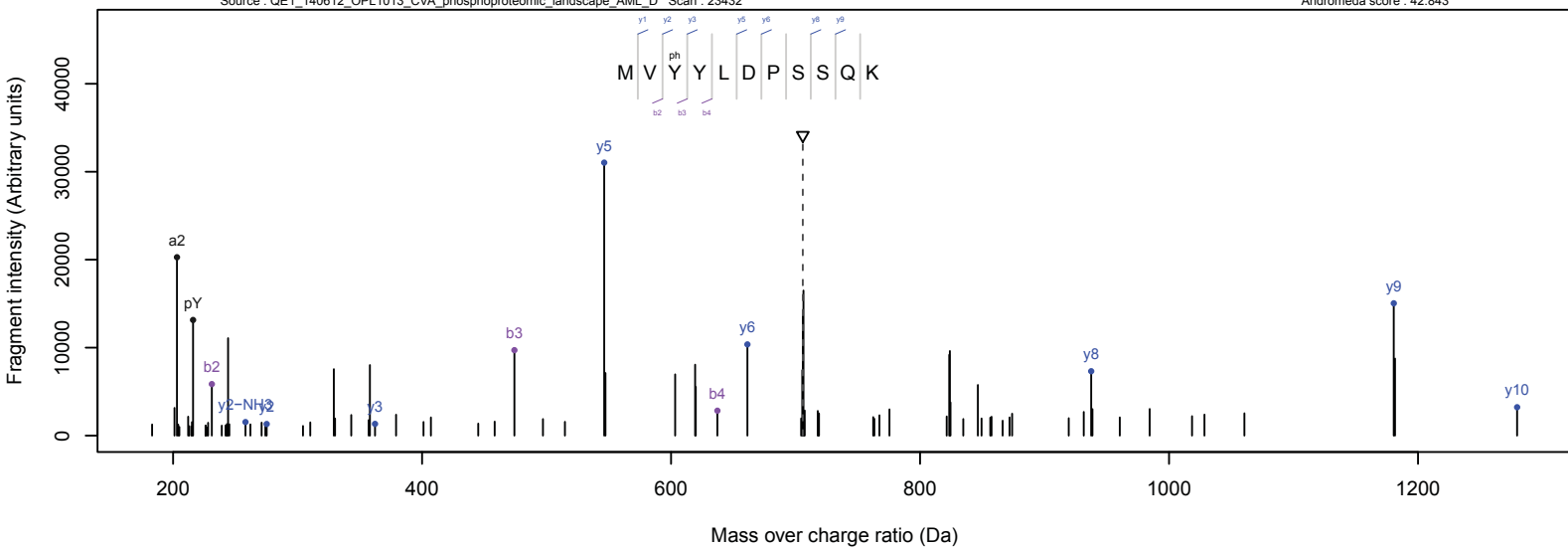
Andromeda score : 42.843



MVY(ph)YLDPSSQK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 23432

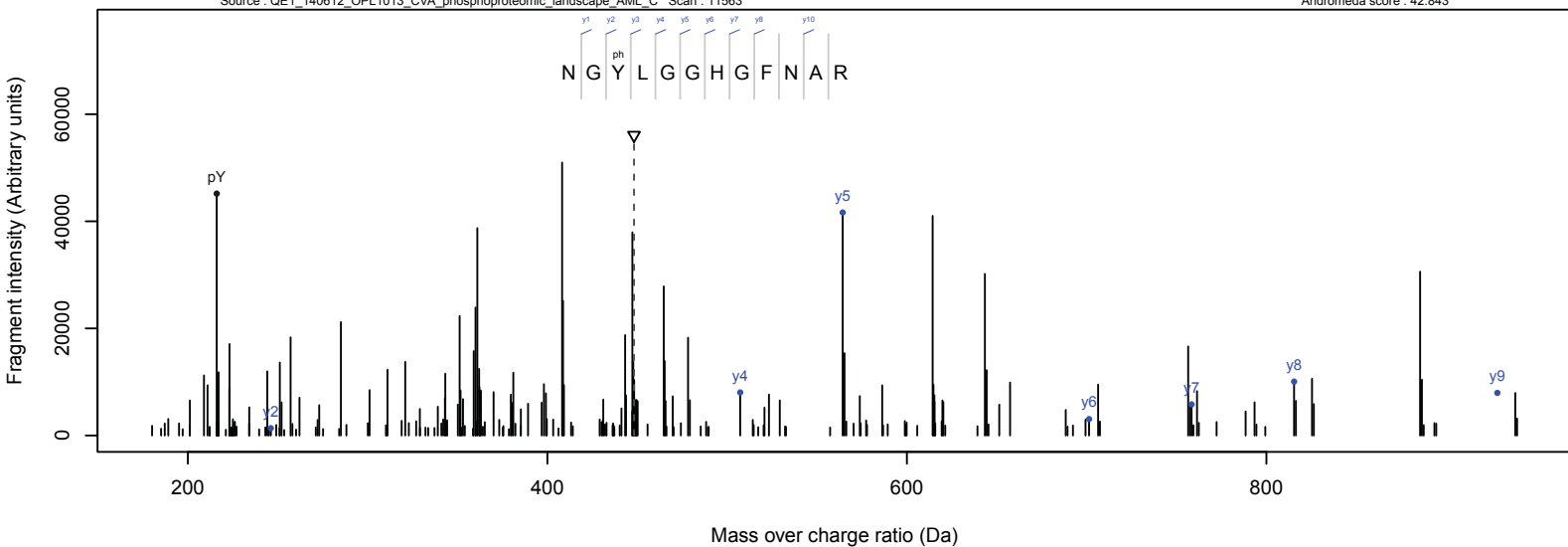
Andromeda score : 42.843



NGY(ph)LGGHGFNAR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11563

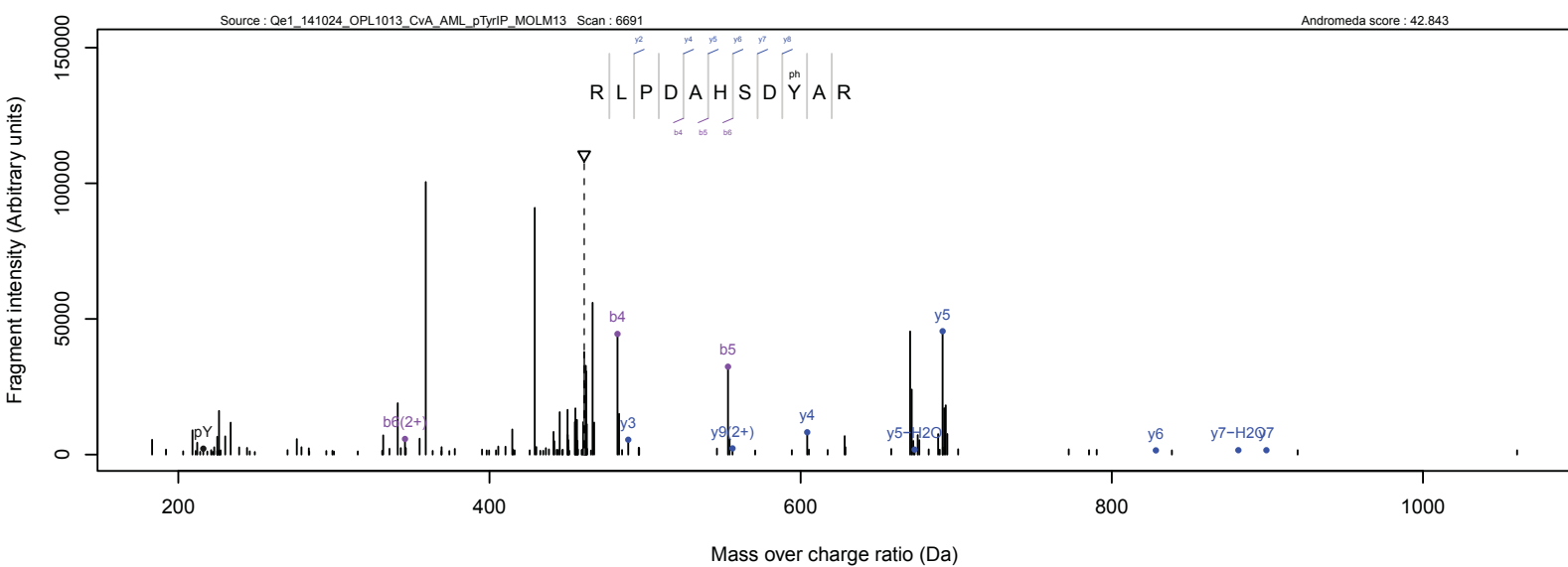
Andromeda score : 42.843



RLPDAHSDYAR-3+

Source : Qe1\_141024\_OPL1013\_CvA\_AML\_pTyrIP\_MOLM13 Scan : 6691

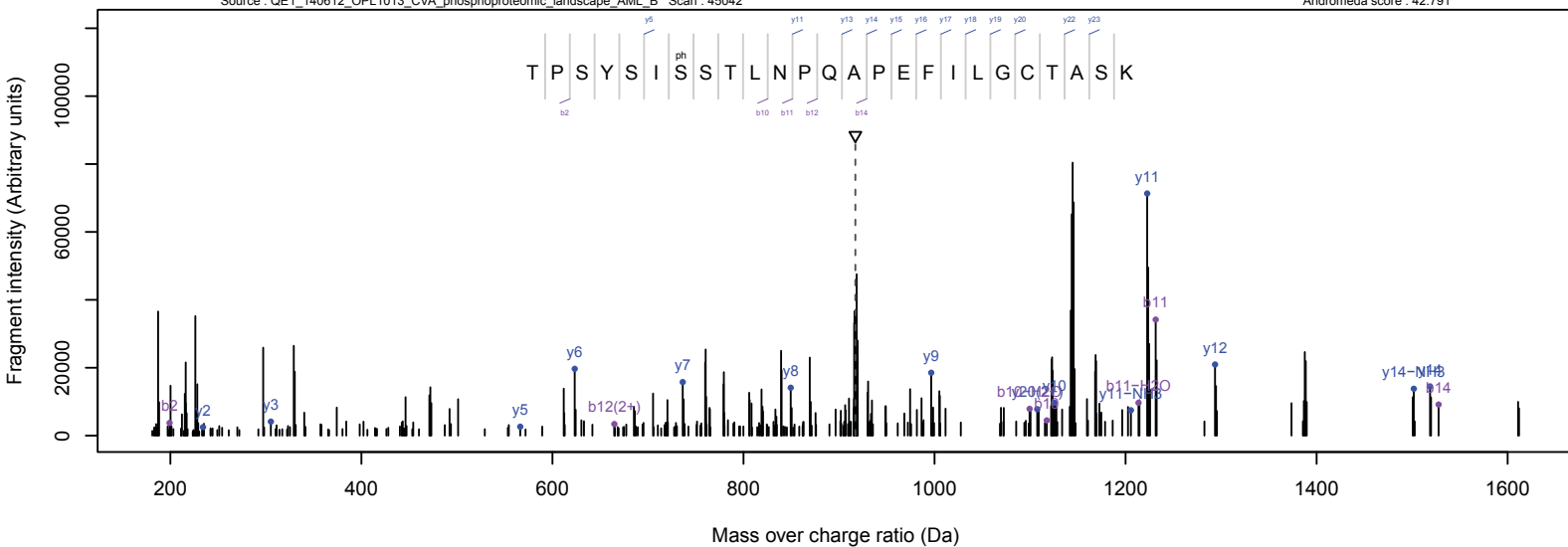
Andromeda score : 42.843



TPSYSIS(ph)STLNPQAPEFILGCTASK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 45042

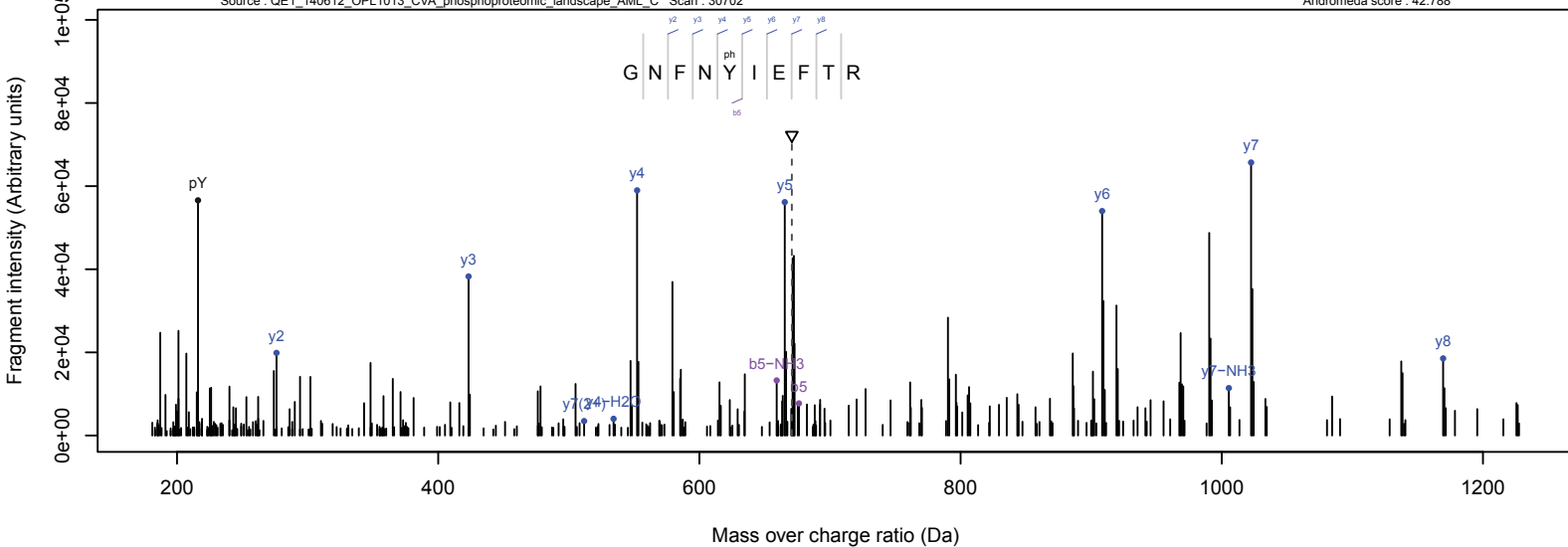
Andromeda score : 42.791



GNFNYPHIEFTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30702

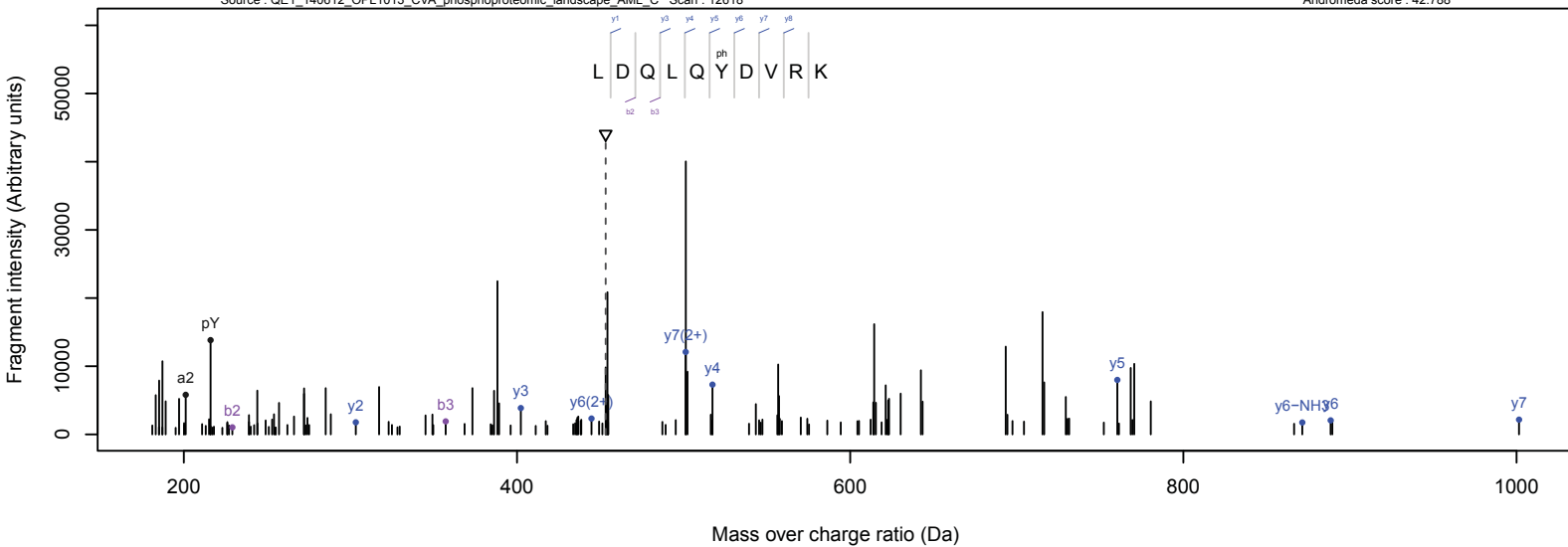
Andromeda score : 42.788



LDQLQY(ph)DVRK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12618

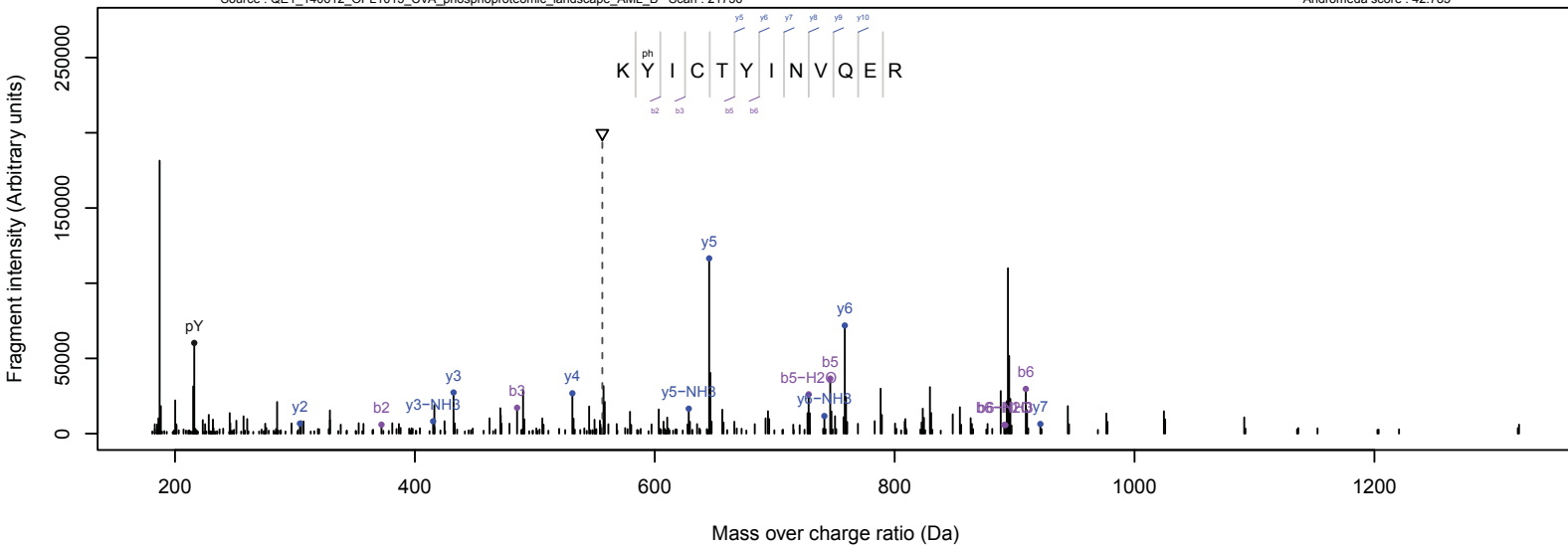
Andromeda score : 42.788



## KY(ph)ICTYINVQER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 21736

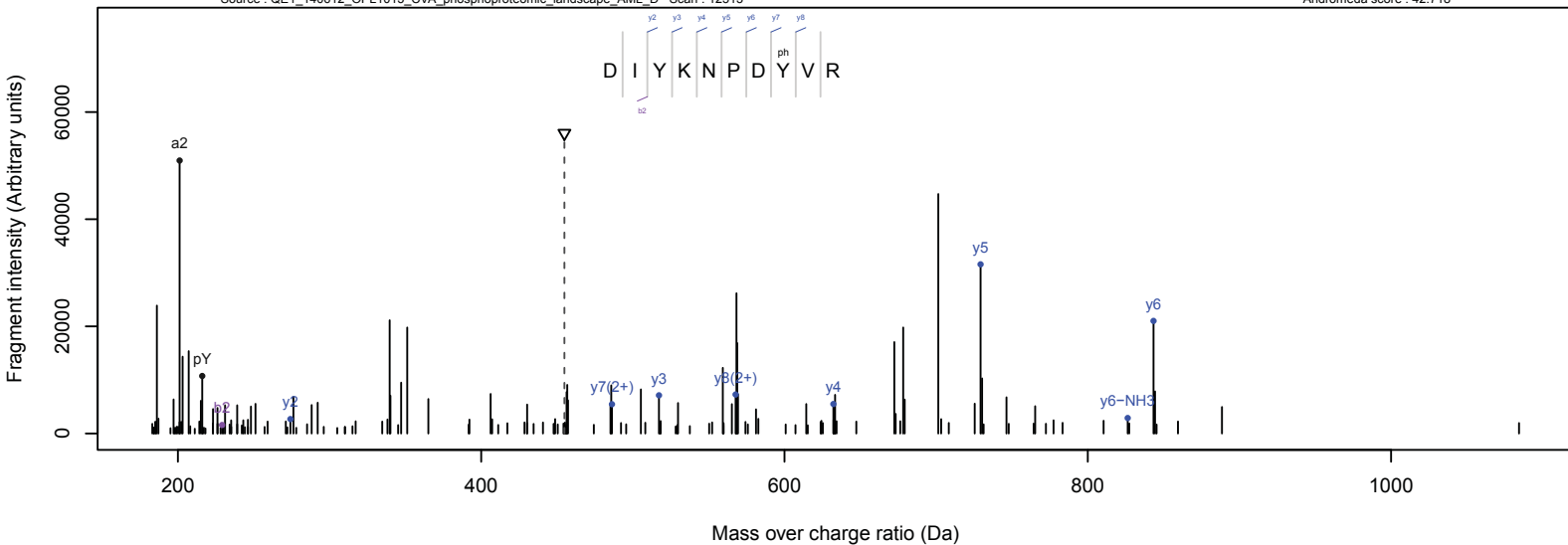
Andromeda score : 42.785



## DIYKNPDYVR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12513

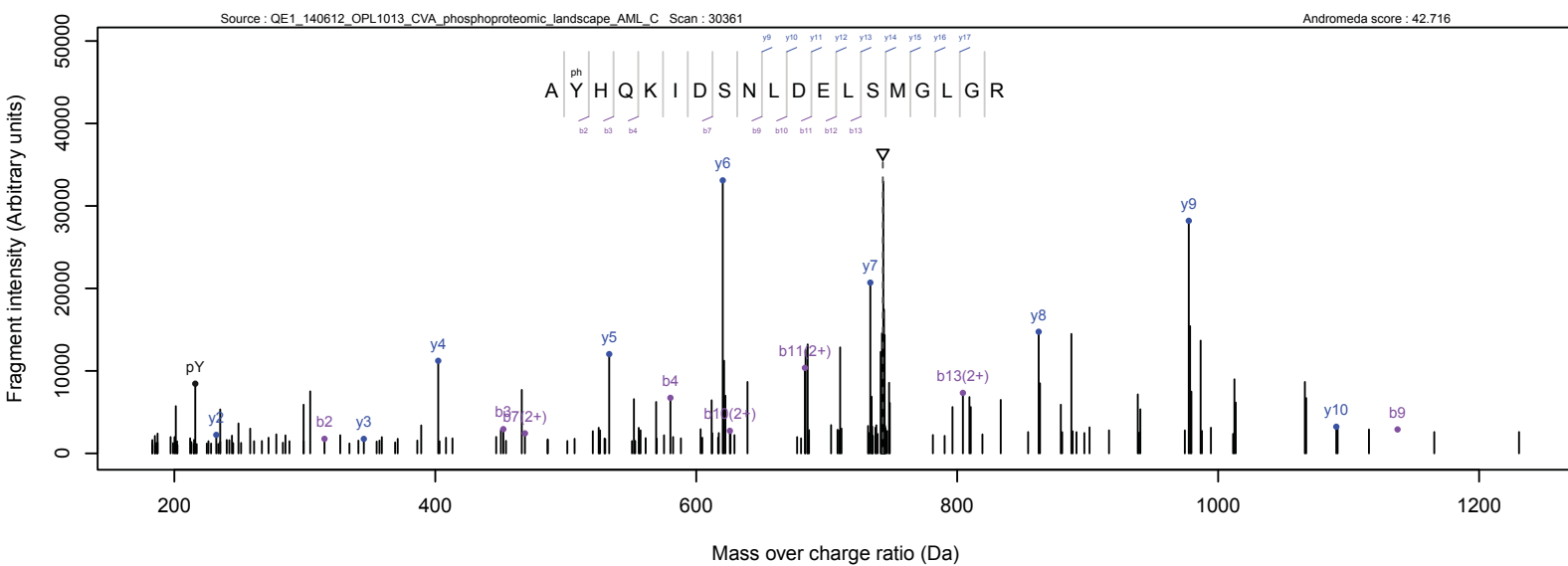
Andromeda score : 42.718



## AY(ph)HQKIDSLDEL S MGLGR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30361

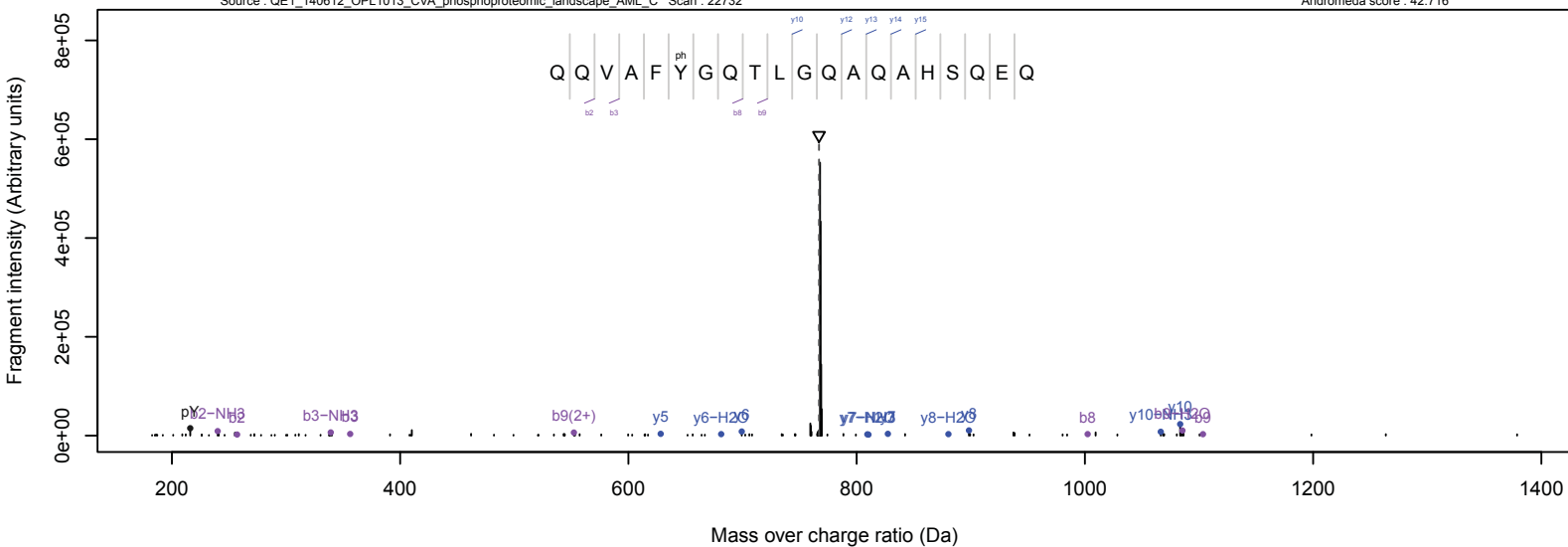
Andromeda score : 42.716



QQVAFY(ph)GQTLGQAQAHSEQ-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 22732

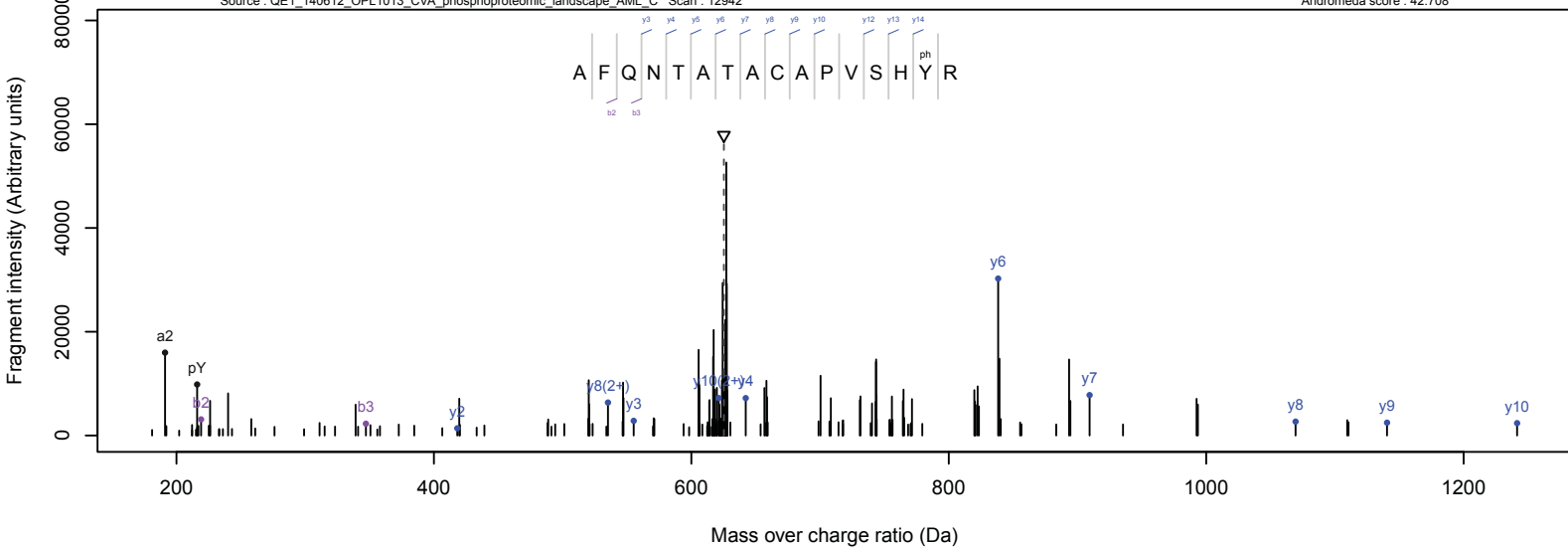
Andromeda score : 42.716



AFQNTATACAPVSHY(ph)R-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 12942

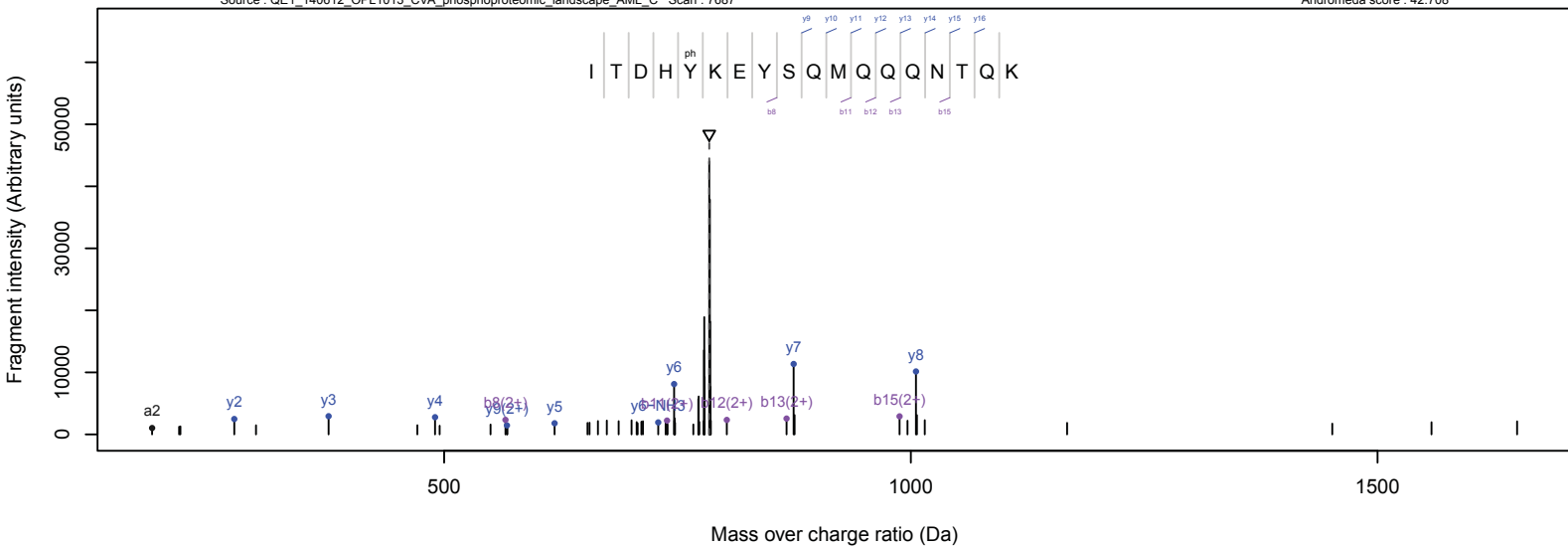
Andromeda score : 42.708



ITDHY(ph)KEYSQMQQNNTQK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 7687

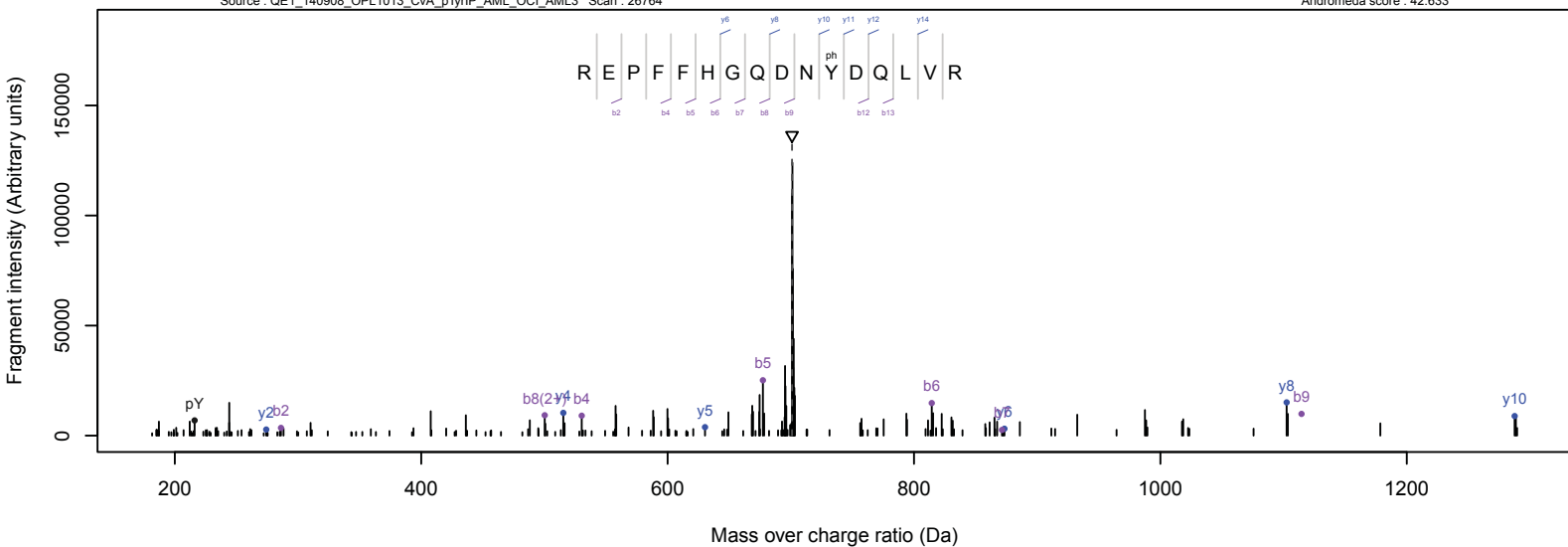
Andromeda score : 42.708



REPFHGDNY(ph)DQLVR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 26764

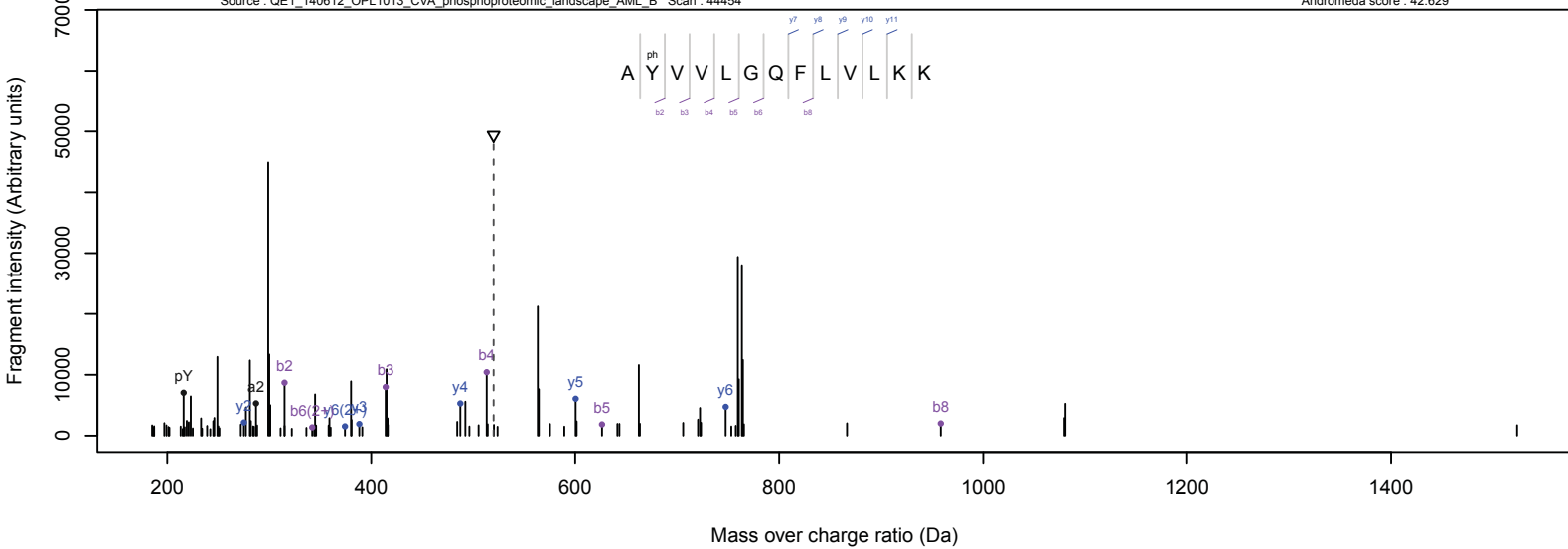
Andromeda score : 42.633



AY(ph)VVLGQFLVLLKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44454

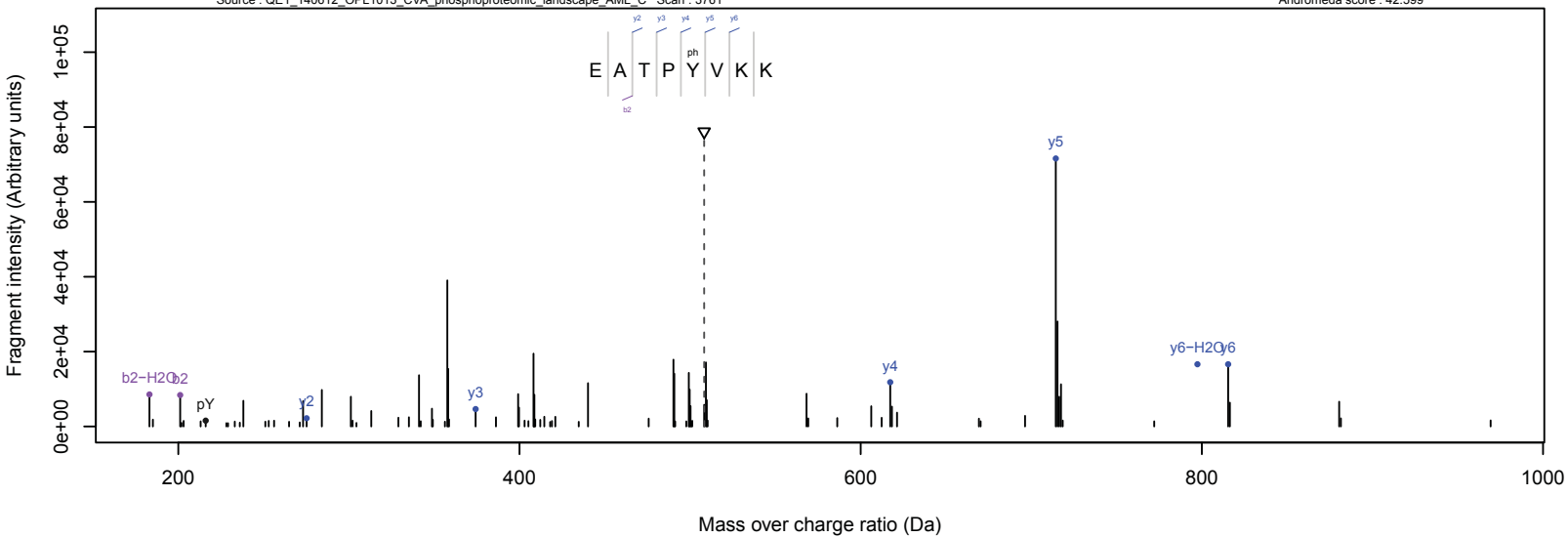
Andromeda score : 42.629



EATPY(ph)VKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 3761

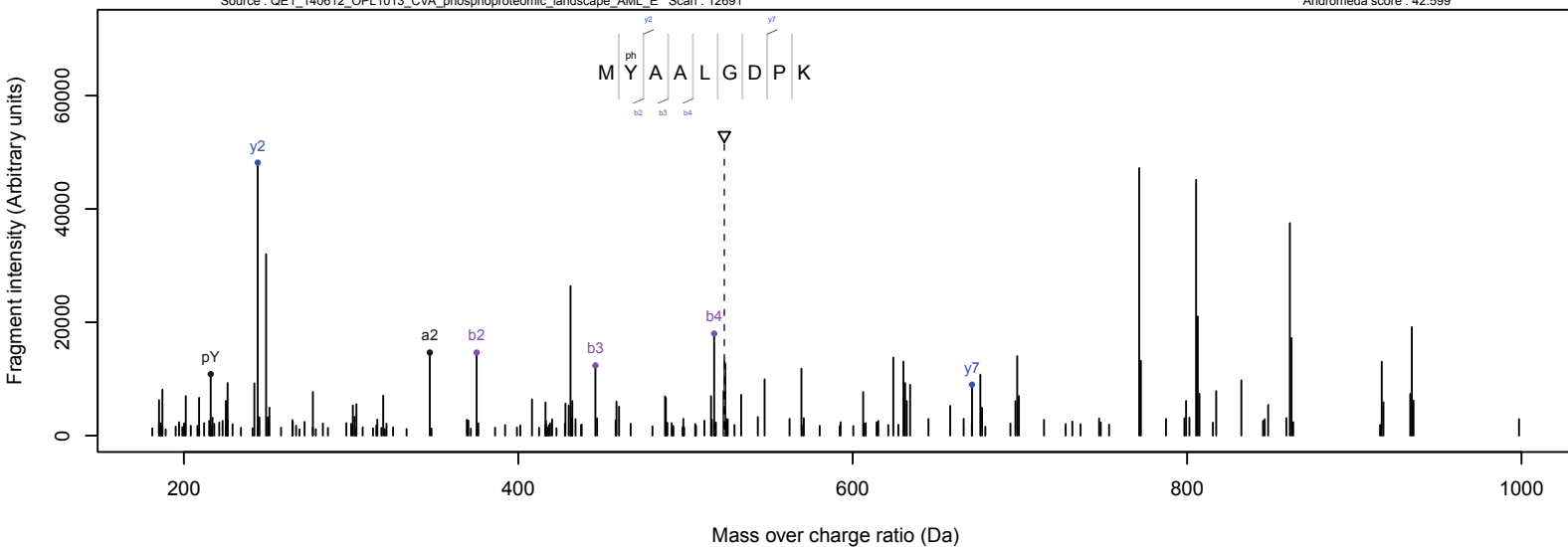
Andromeda score : 42.599



## MY(ph)AALGDPK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 12691

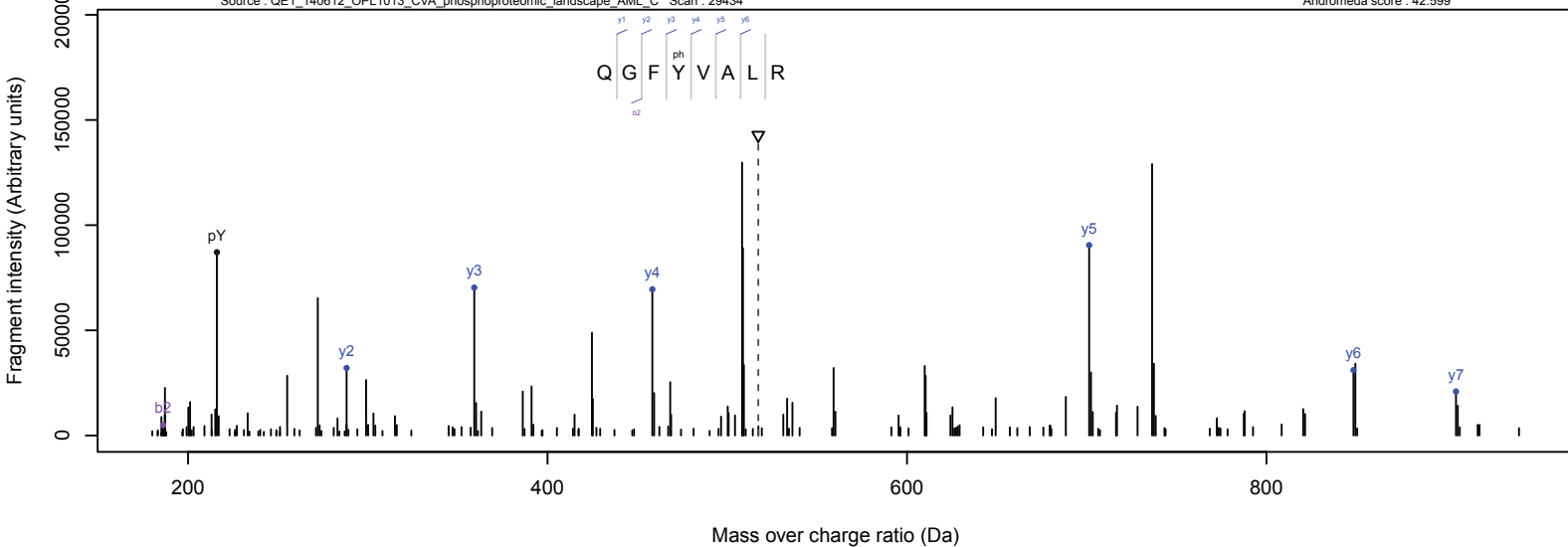
Andromeda score : 42.599



## QGFY(ph)VALR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29434

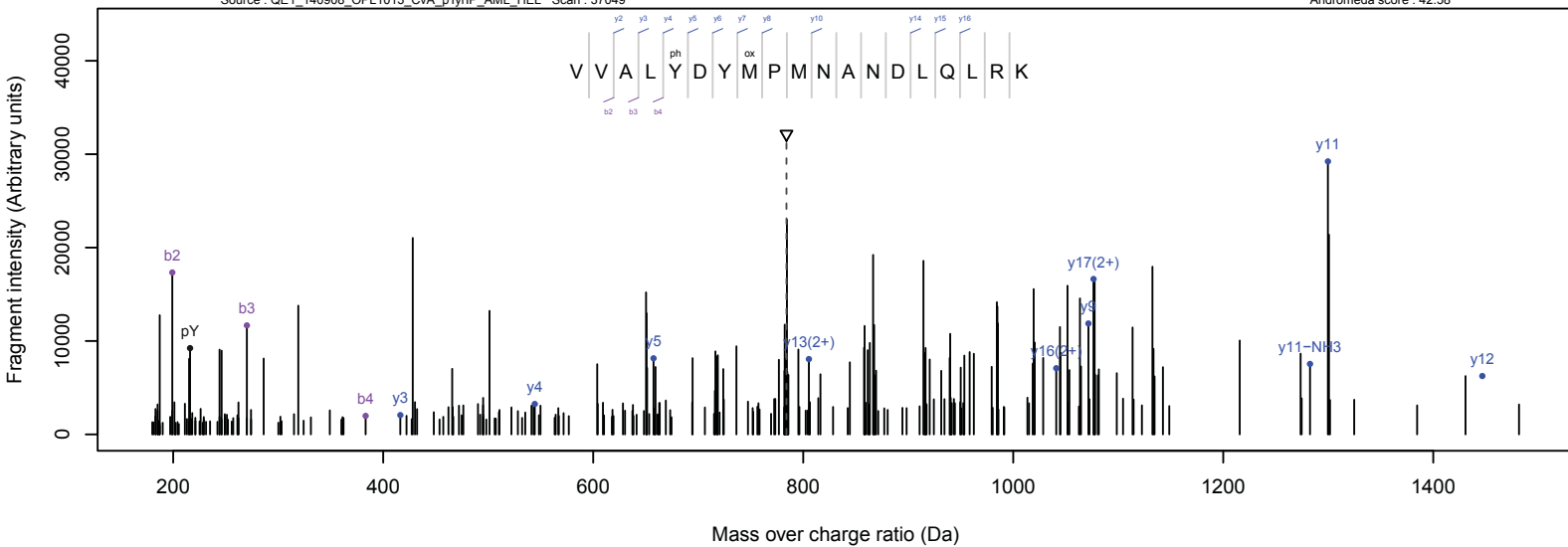
Andromeda score : 42.599



## VVALY(ph)DYM(ox)PMNANDLQLRK-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 37049

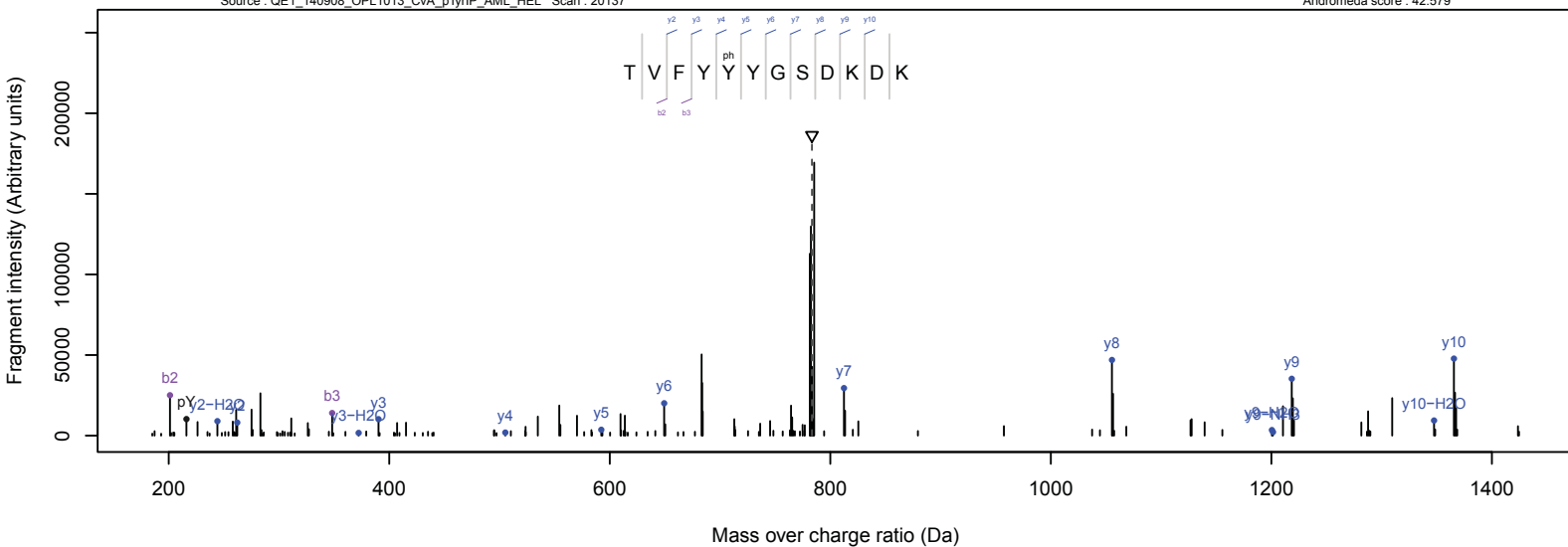
Andromeda score : 42.58



TVFY(ph)YGSDDK-2+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 20137

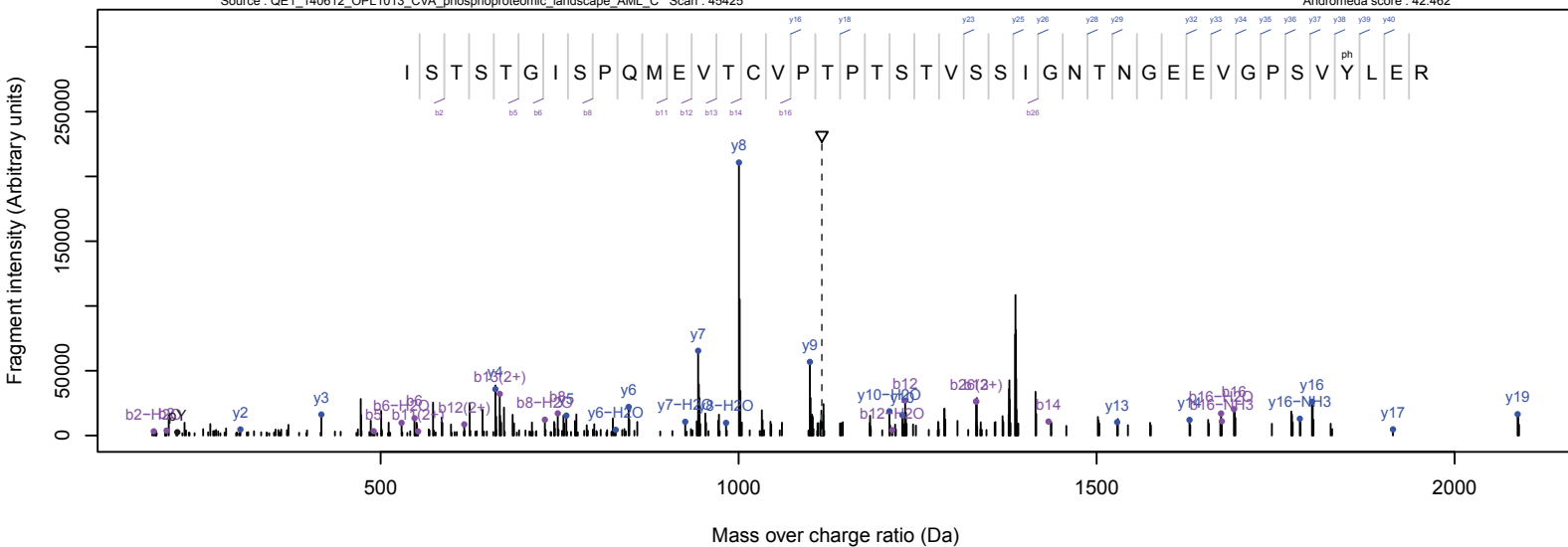
Andromeda score : 42.579



ISTSTGISQMEVTCVPTPTSTVSSIGNTNGEEVGPSVY(ph)LER-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 45425

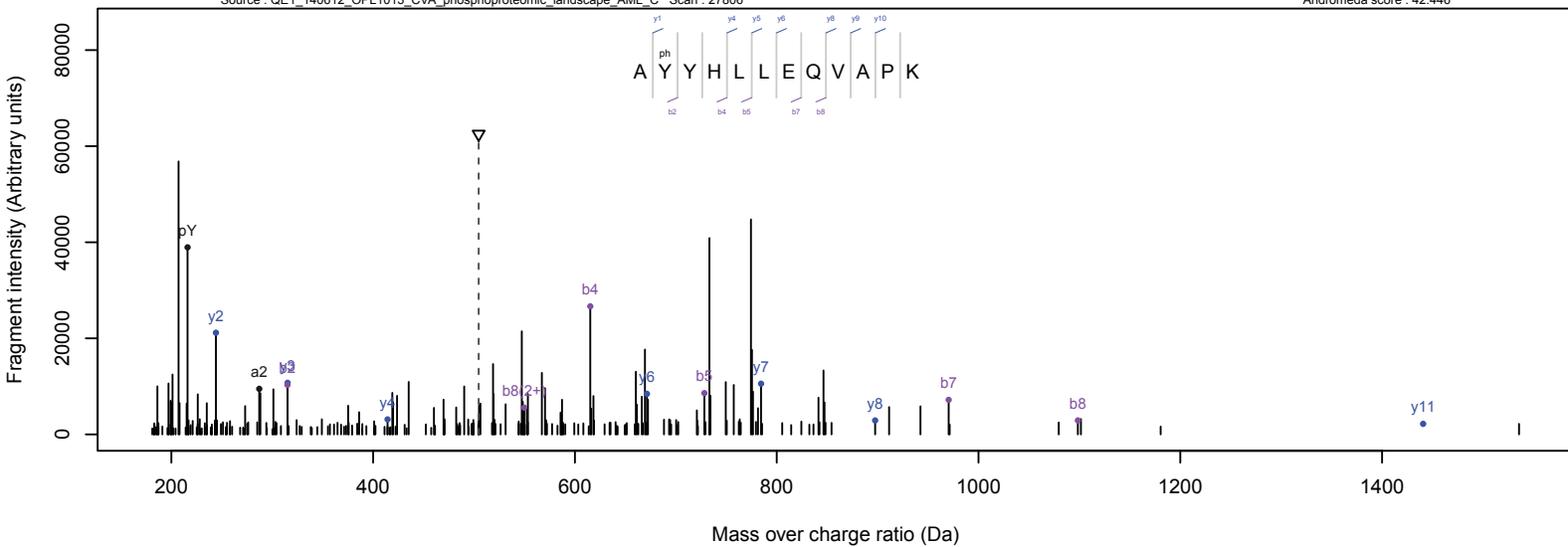
Andromeda score : 42.462



AY(ph)YHLLQVAPK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 27806

Andromeda score : 42.446

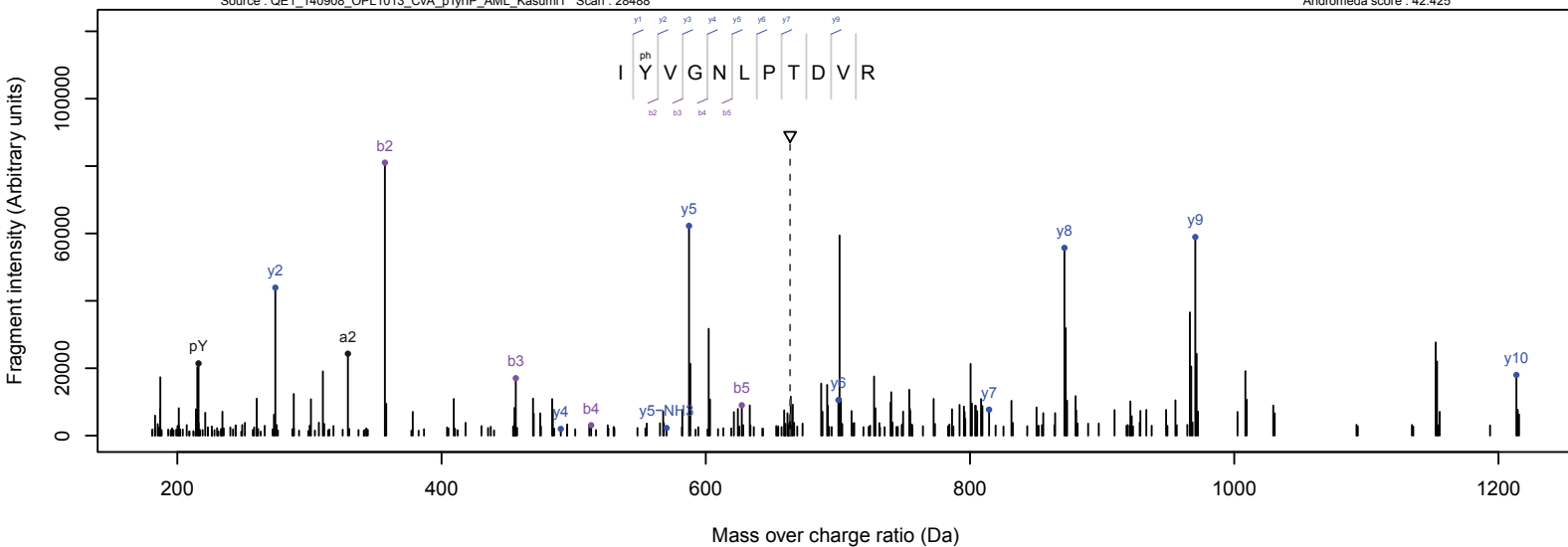




## IY(ph)VGNLPTDVR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 28488

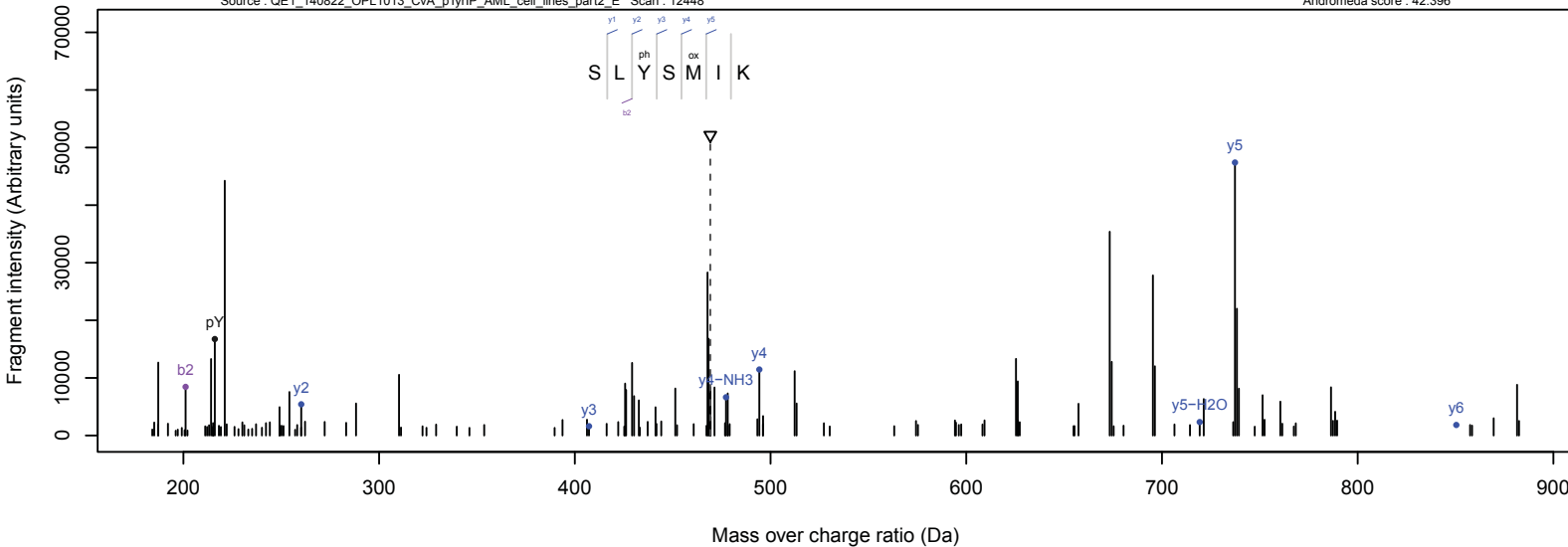
Andromeda score: 42.425



## SLY(ph)SM(ox)IK-2+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 12448

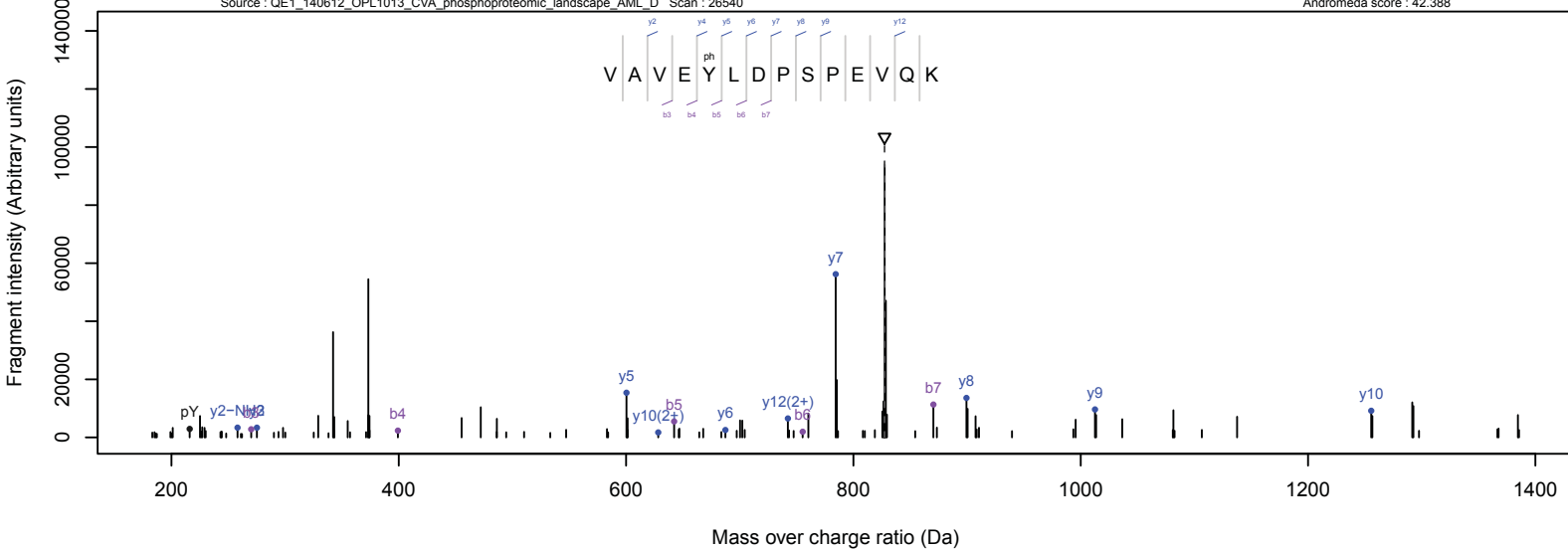
Andromeda score: 42.396



## VAVEY(ph)LDPSPEVQK-2+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_D Scan: 26540

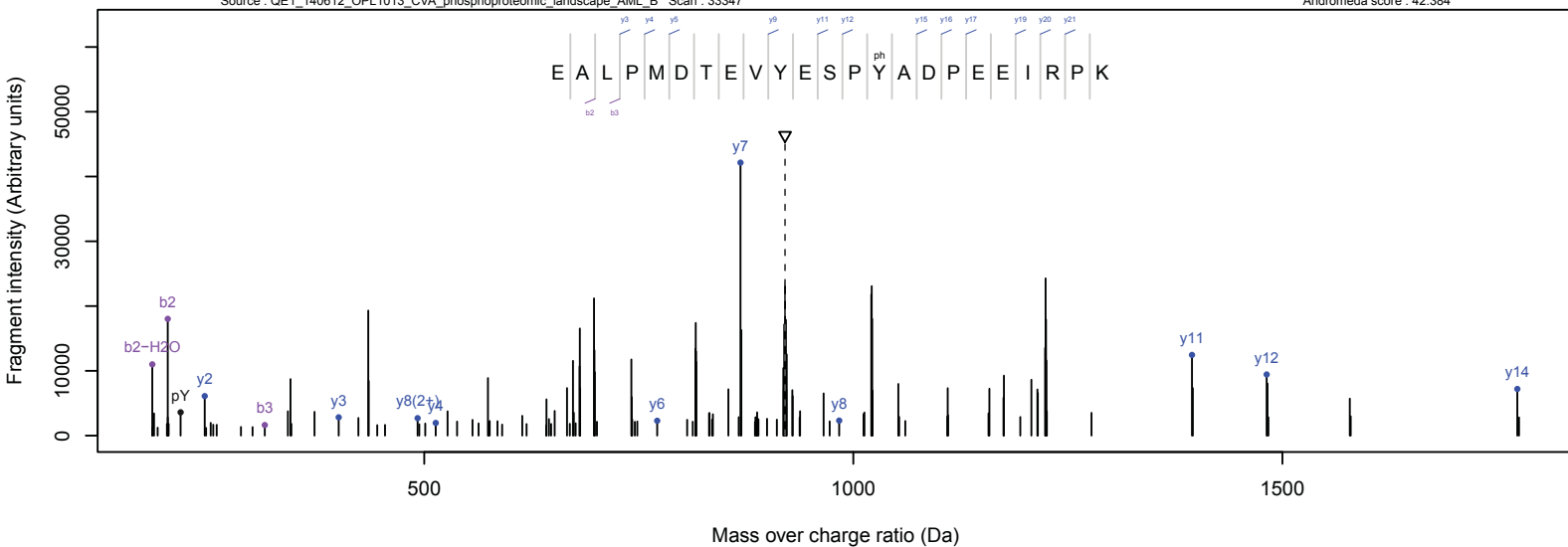
Andromeda score: 42.388



**EALPMDTEVYESPY(ph)ADPEEIRPK-3+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33347

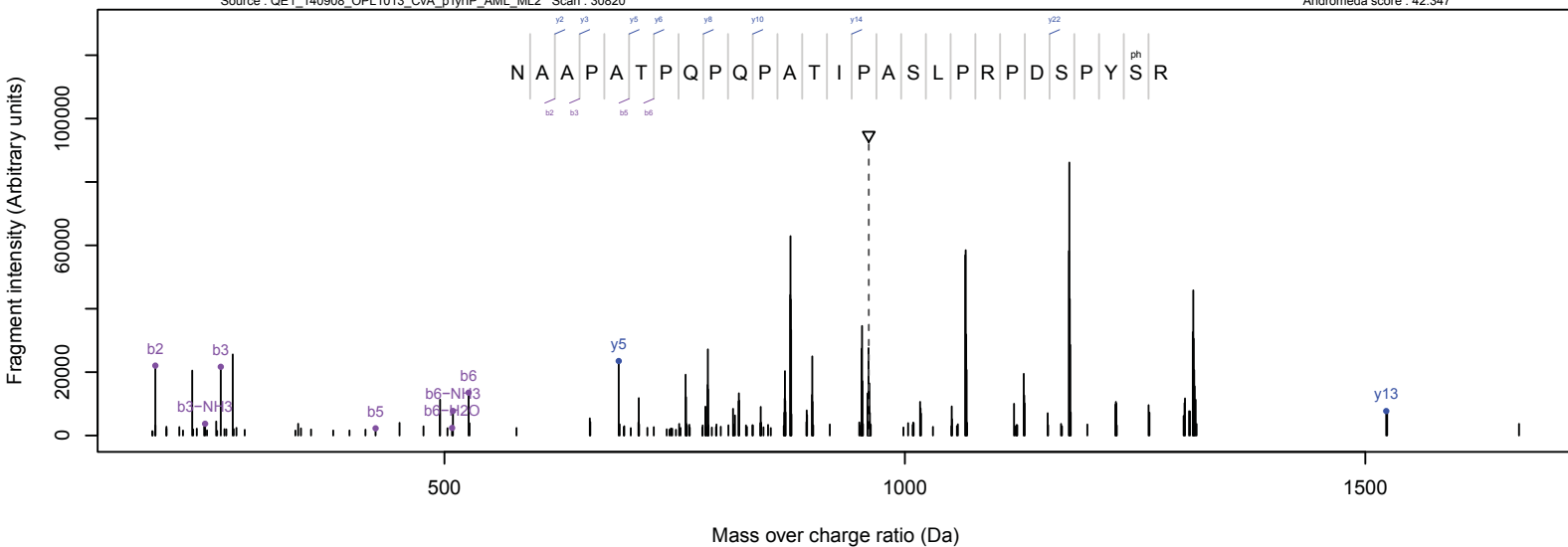
Andromeda score : 42.384



**NAAPATPQPQPATIPASLRPDPSPYS(ph)R-3+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 30820

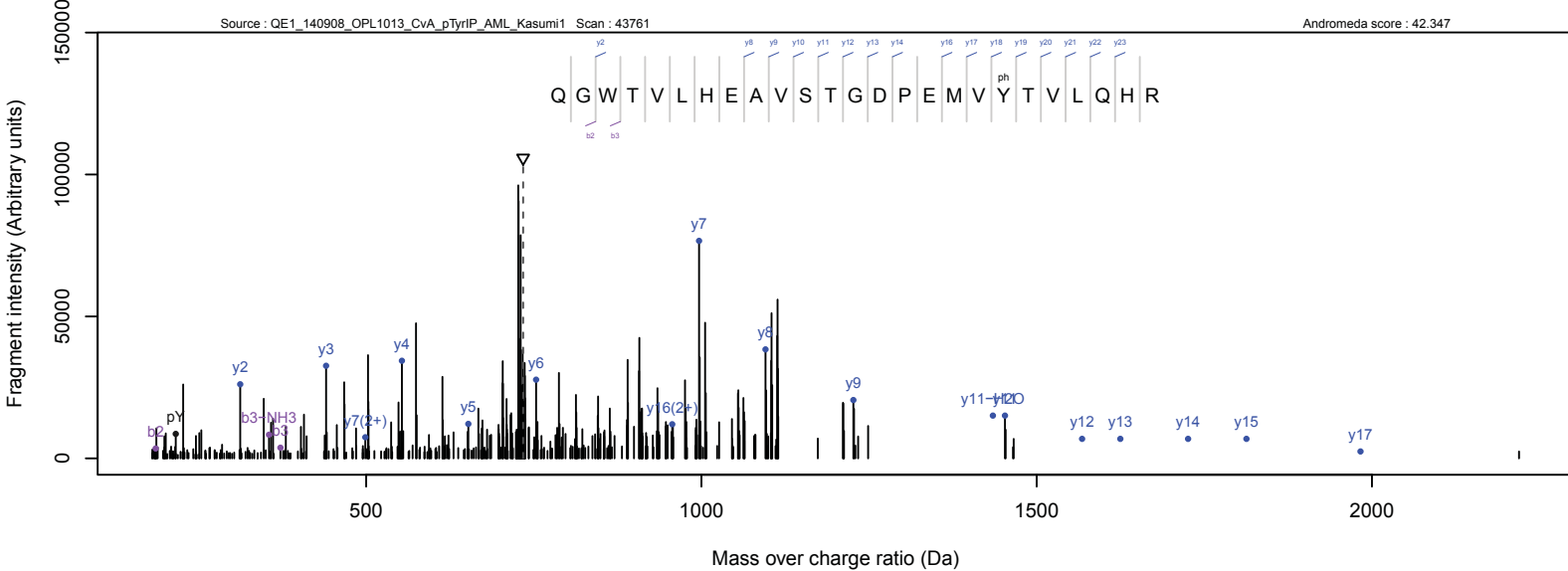
Andromeda score : 42.347



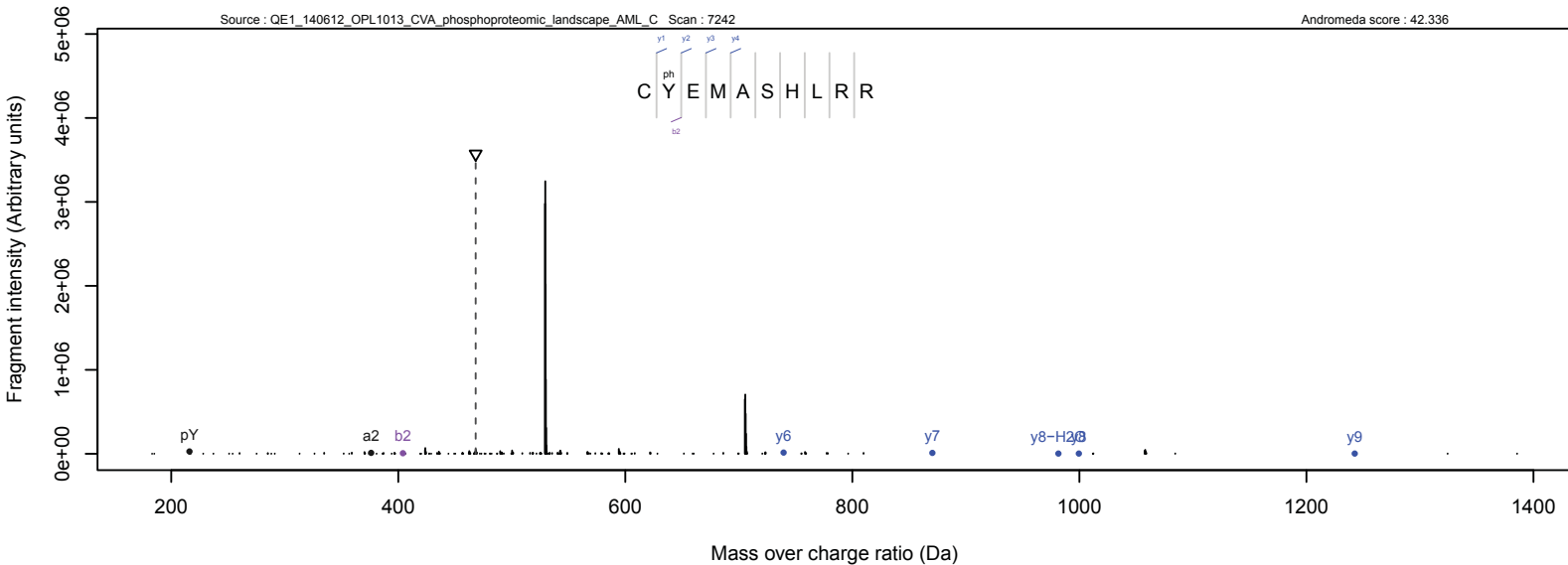
**QGWTVLHEAVSTGDPVMVY(ph)TVLQHR-4+**

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 43761

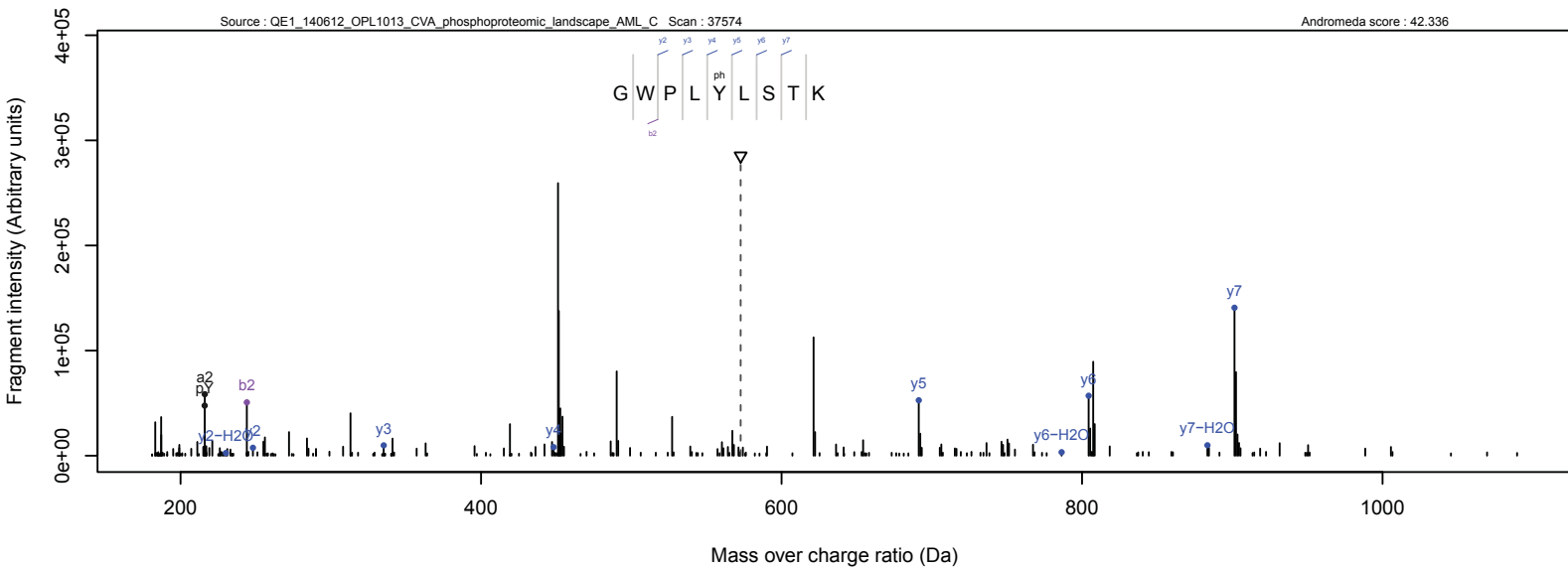
Andromeda score : 42.347



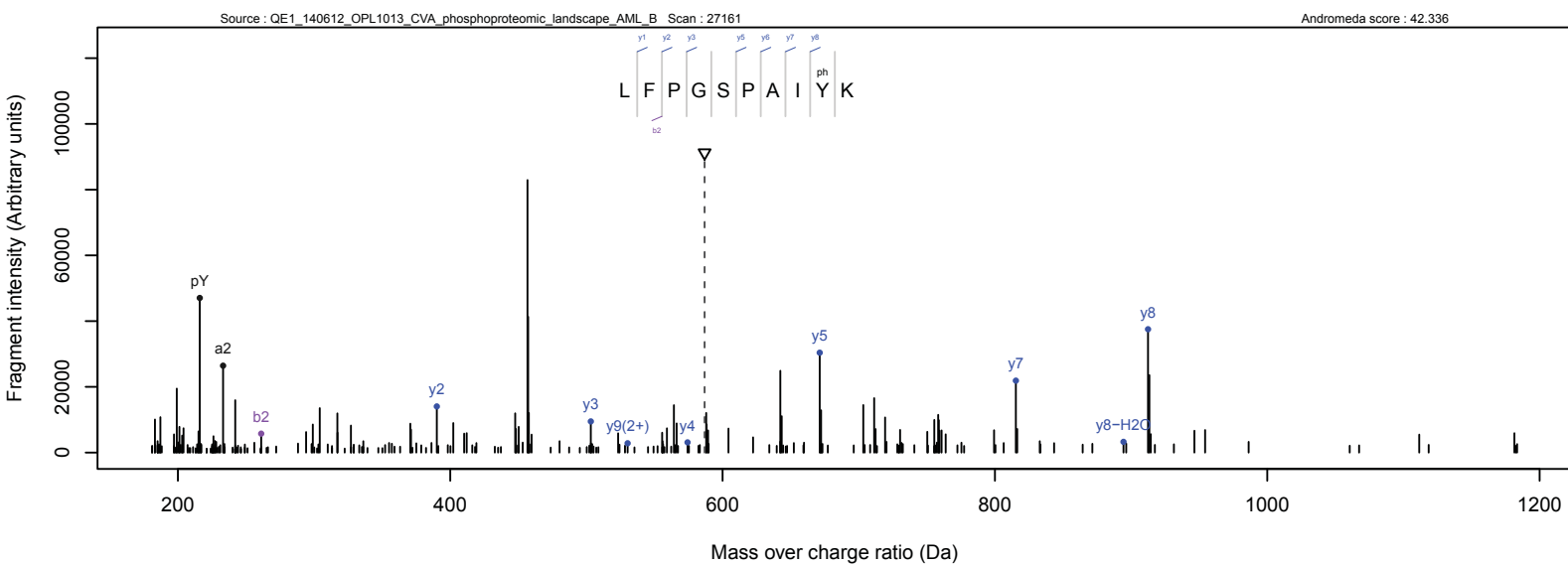
CY(ph)EMASHLRR-3+



GWPLY(ph)LSTK-2+



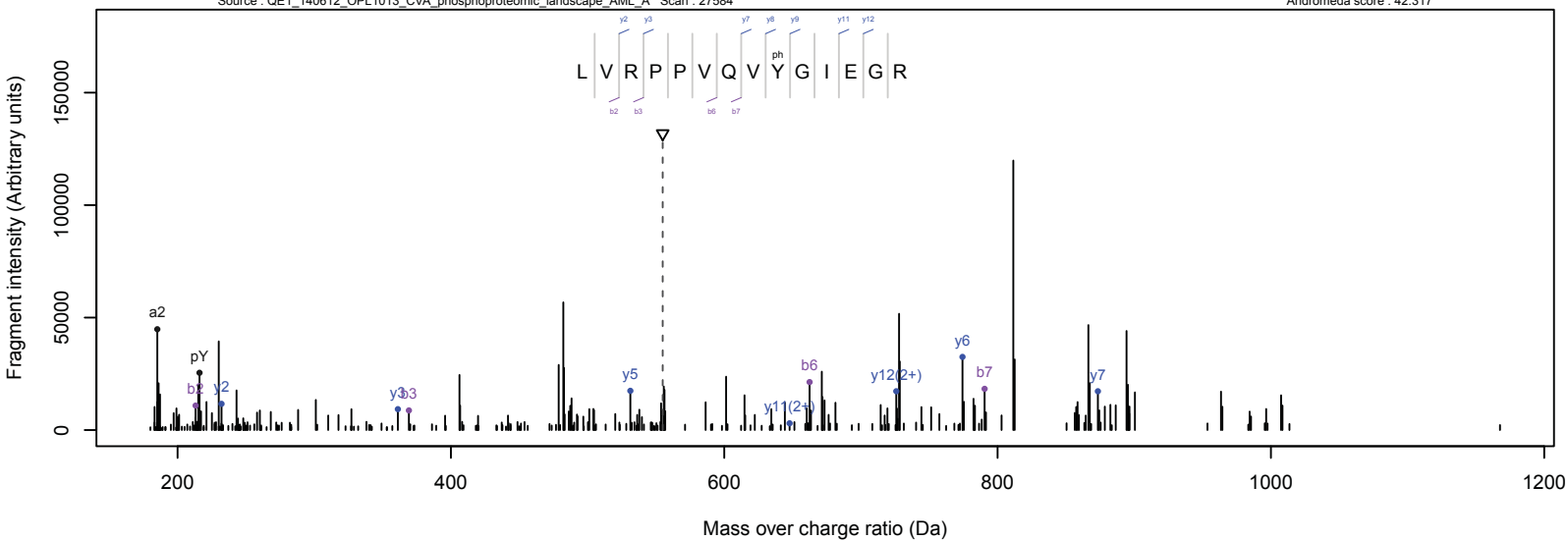
LFGSPAIIYK-2+



## LVRPPQVY(ph)GIEGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 27584

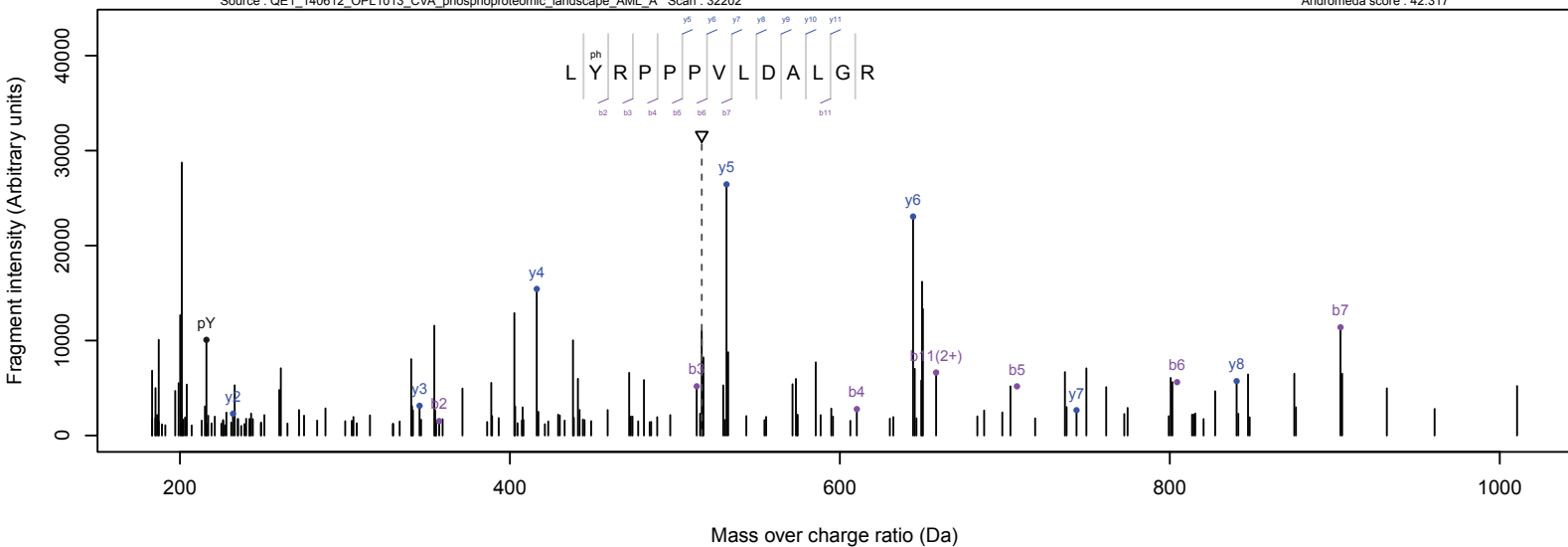
Andromeda score: 42.317



## LY(ph)RPPVLDALGR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 32202

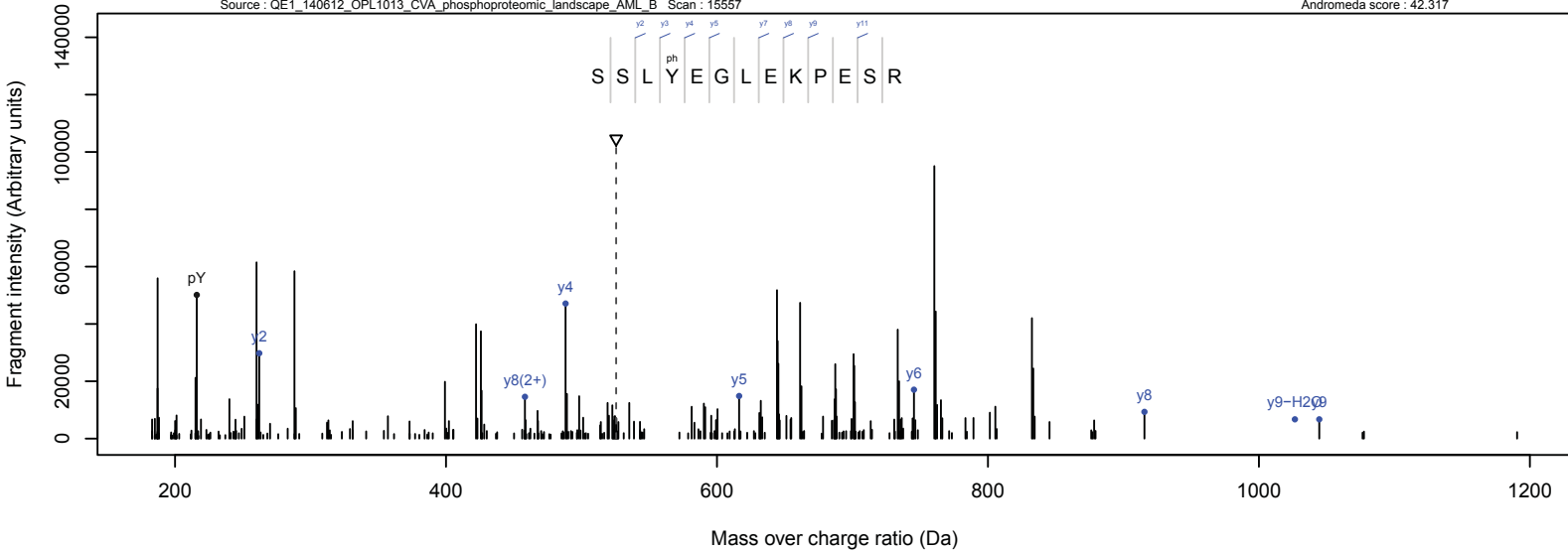
Andromeda score: 42.317



## SSLY(ph)EGLEKPEESR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 15557

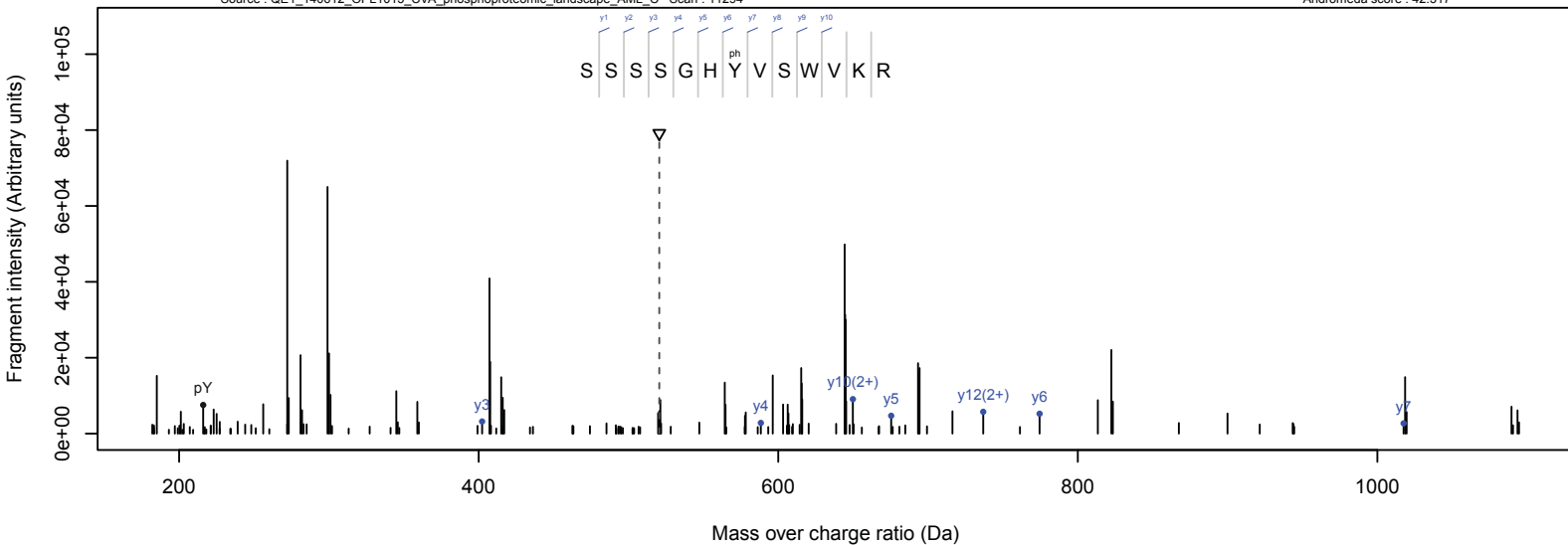
Andromeda score: 42.317



## SSSSGHY(ph)VSWVK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 11294

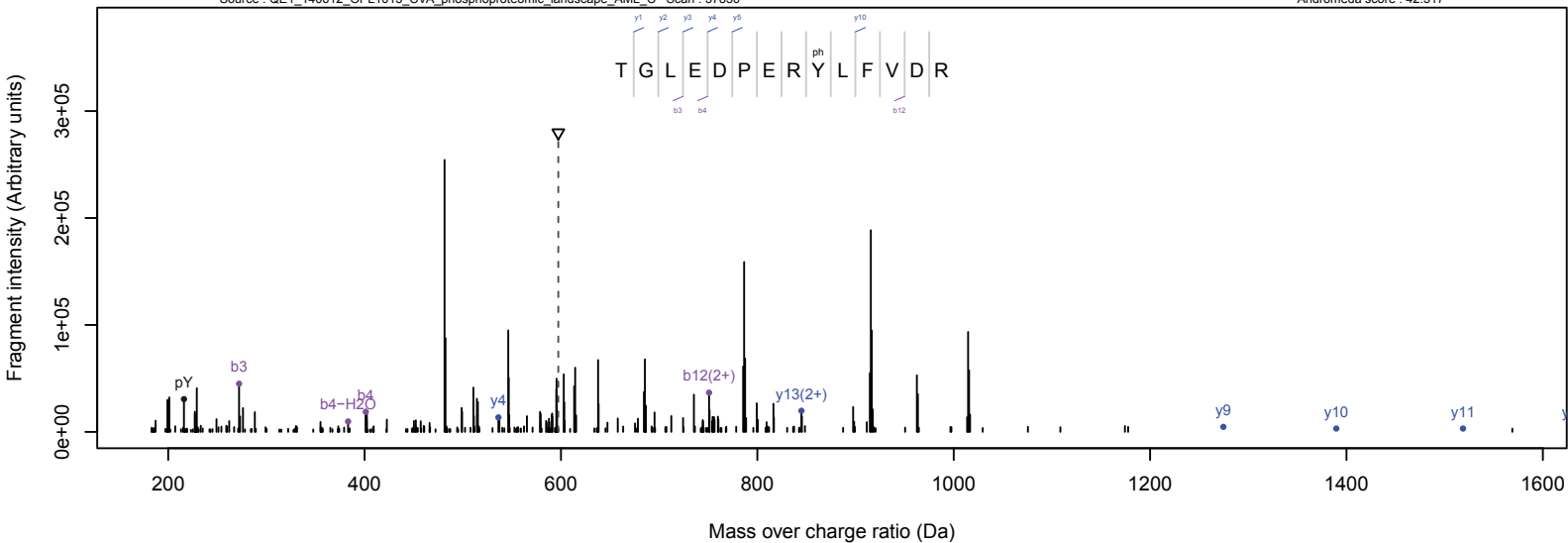
Andromeda score : 42.317



## TGLEDPERY(ph)LFVDR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 37830

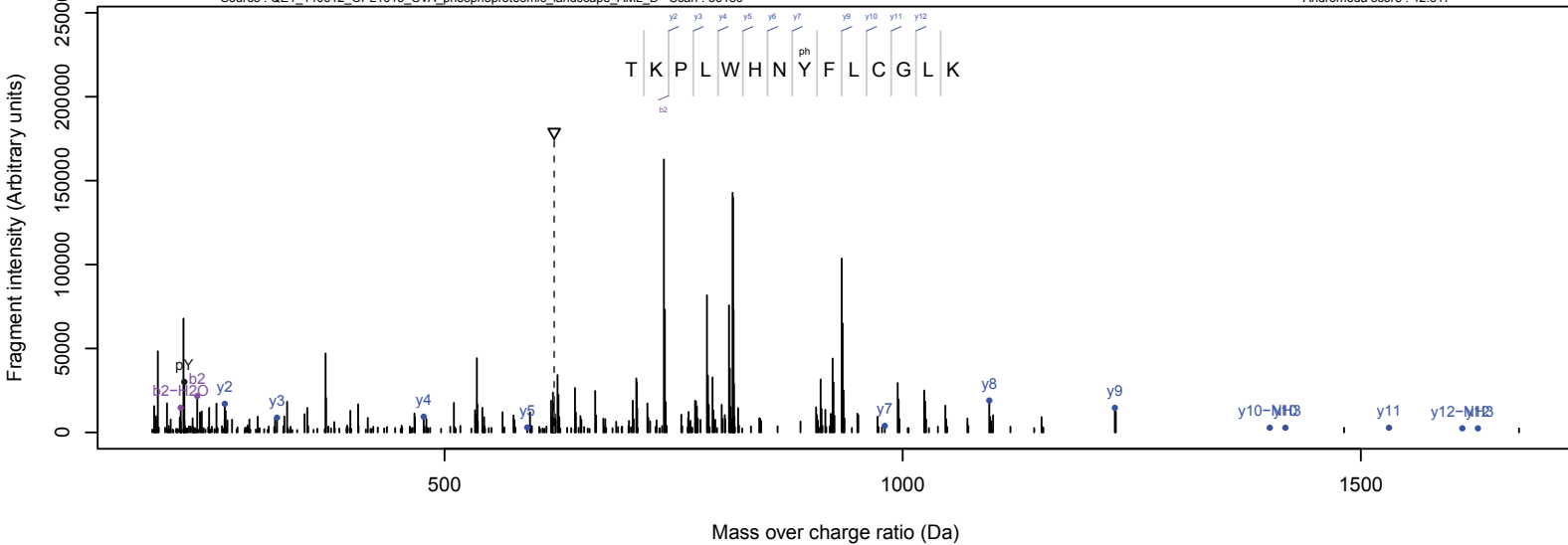
Andromeda score : 42.317



## TKPLWHNY(ph)FLCGLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 36159

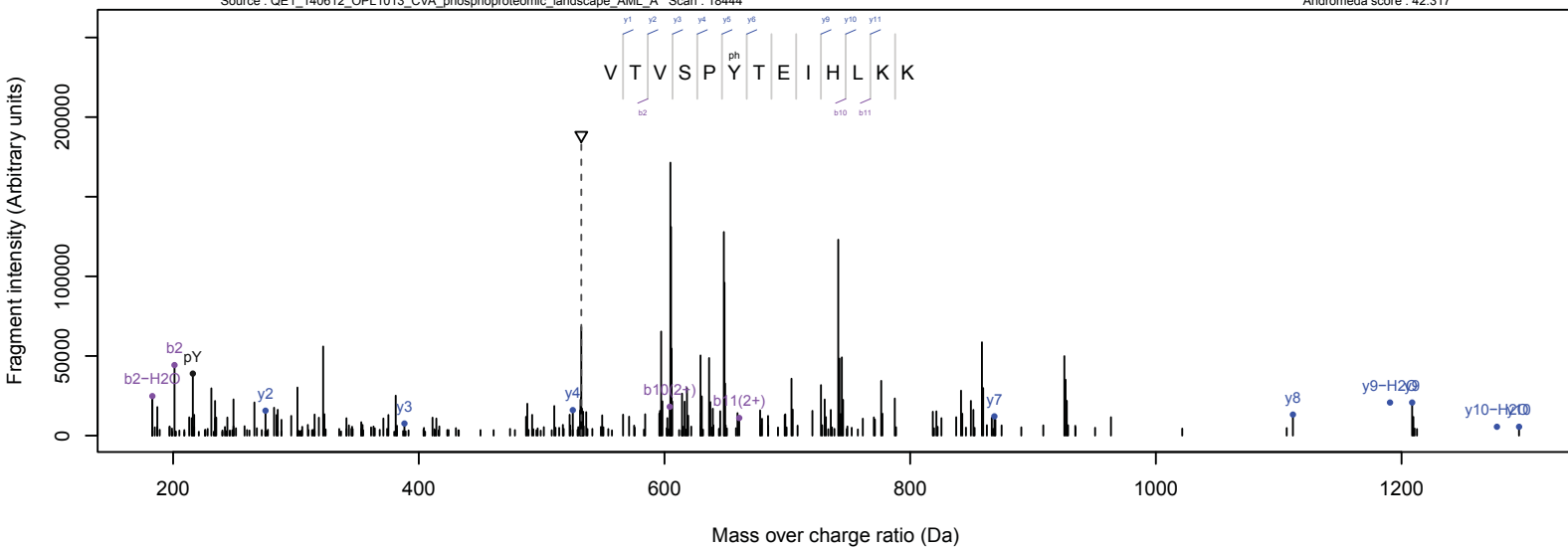
Andromeda score : 42.317



## VTVSPY(ph)TEIHLKK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 18444

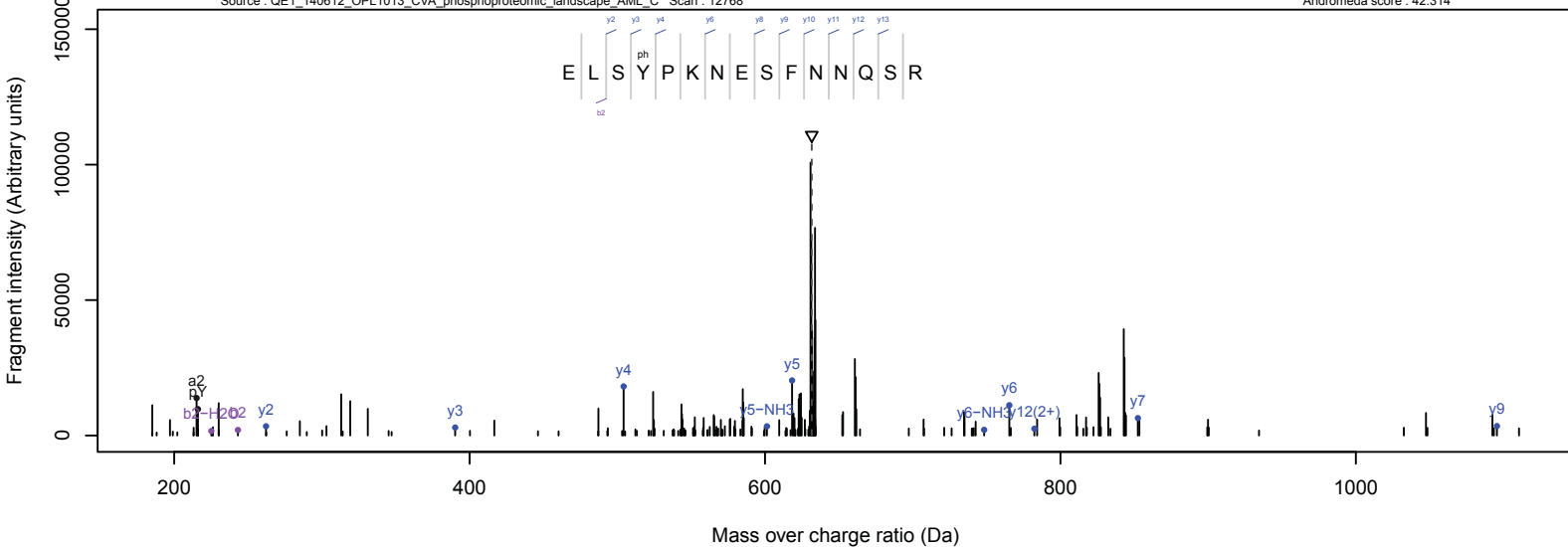
Andromeda score: 42.317



## ELSY(ph)PKNESFNQSR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 12768

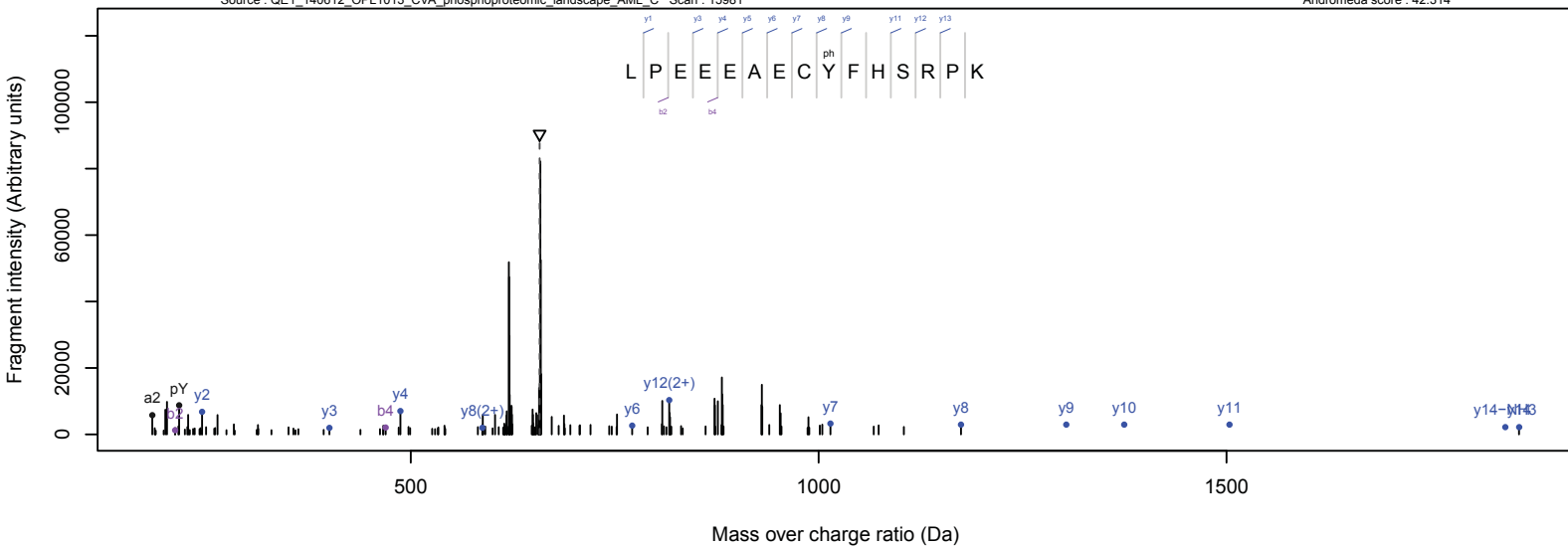
Andromeda score: 42.314



## LPEEEAECY(ph)FHSRPK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 15981

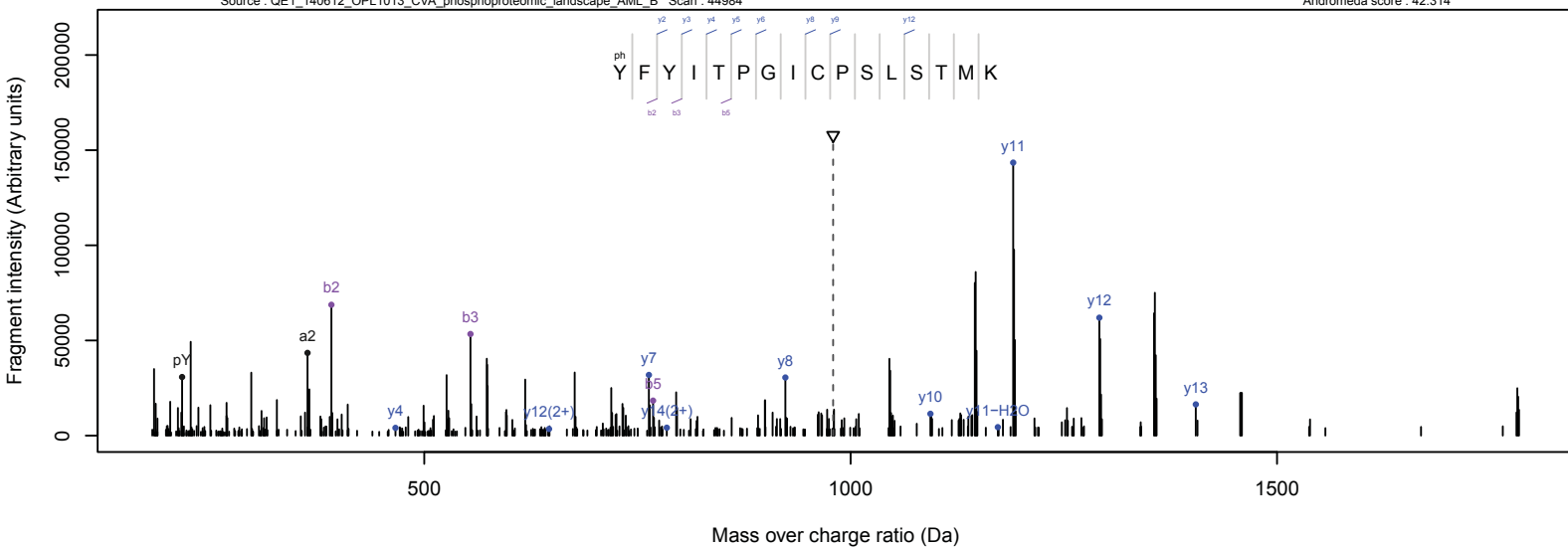
Andromeda score: 42.314



## Y(ph)FYITPGICPSLSTMK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 44984

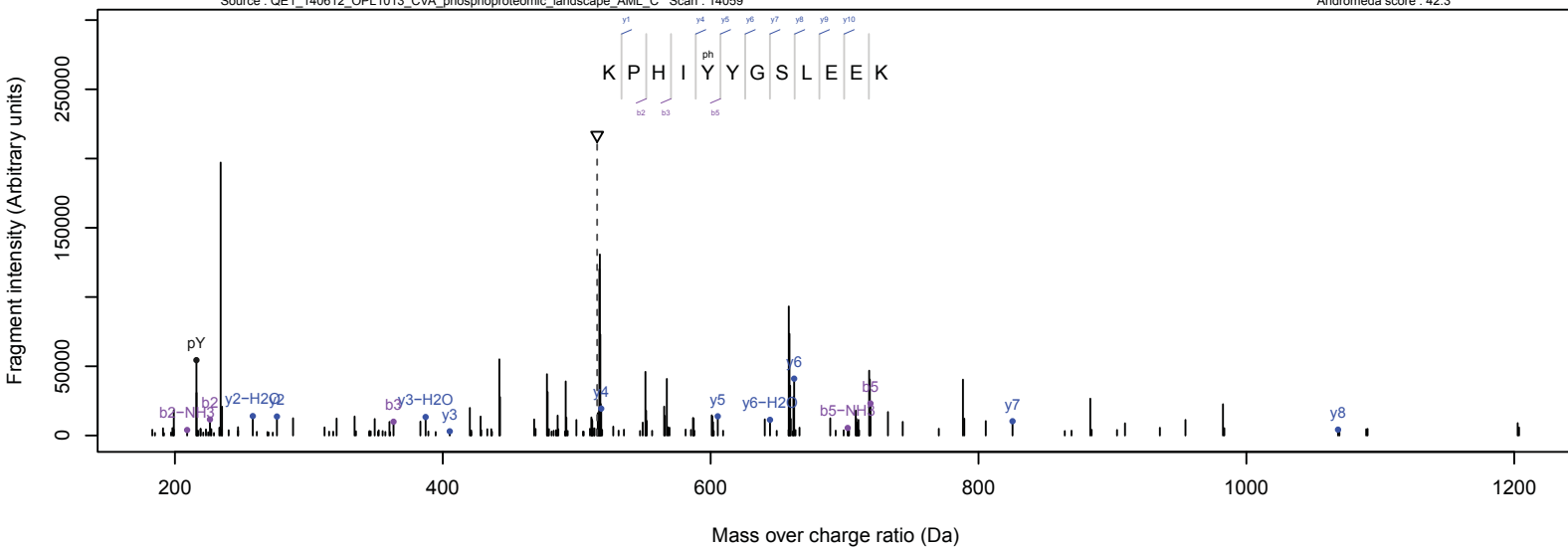
Andromeda score : 42.314



## KPHIY(ph)YGSLEEK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14059

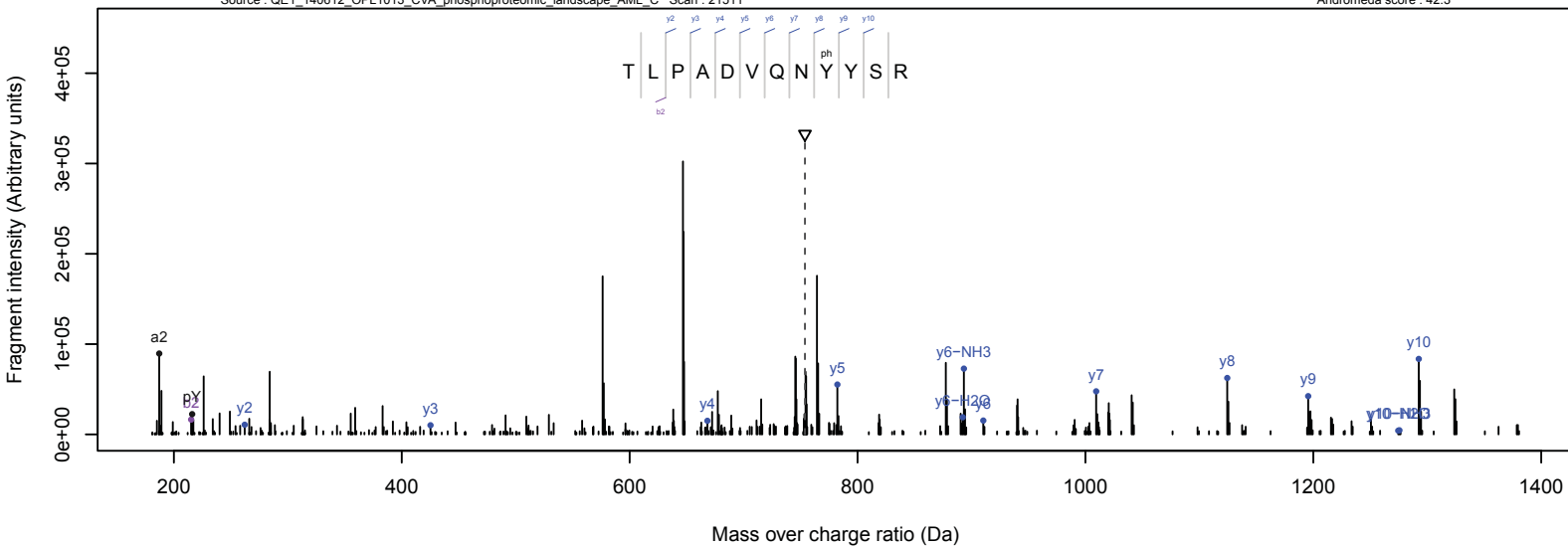
Andromeda score : 42.3



## TLPADVQNYYSR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21511

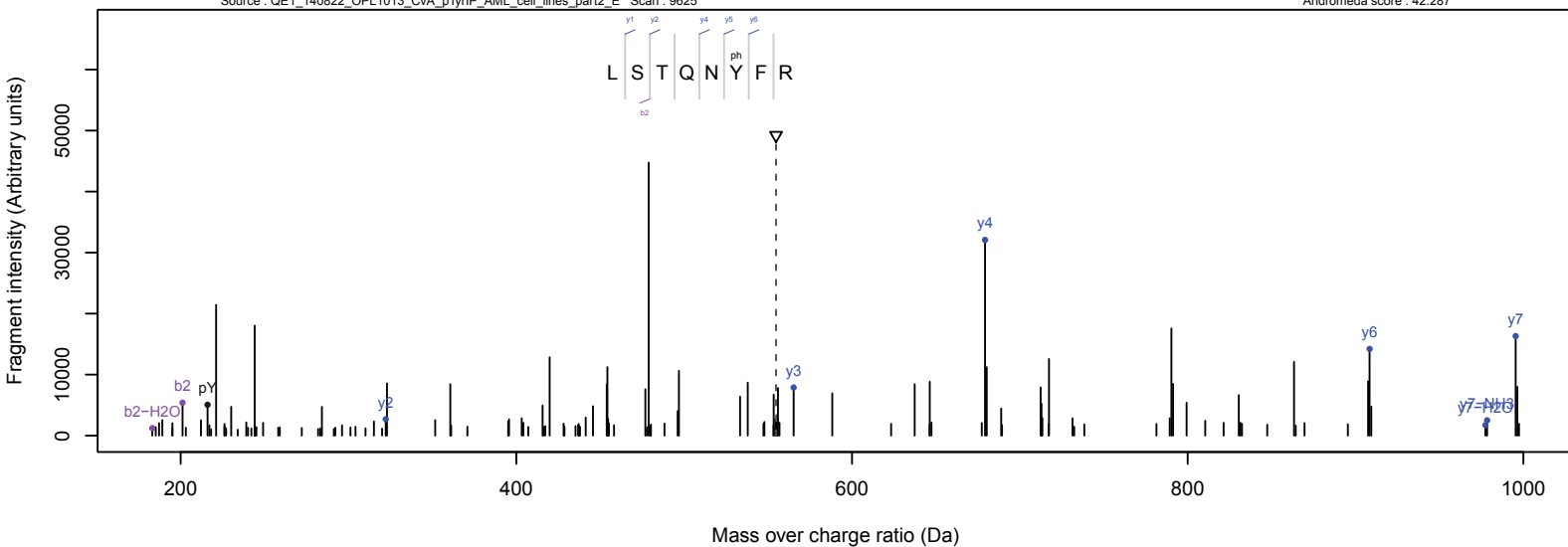
Andromeda score : 42.3



LSTQNY(ph)FR-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan : 9625

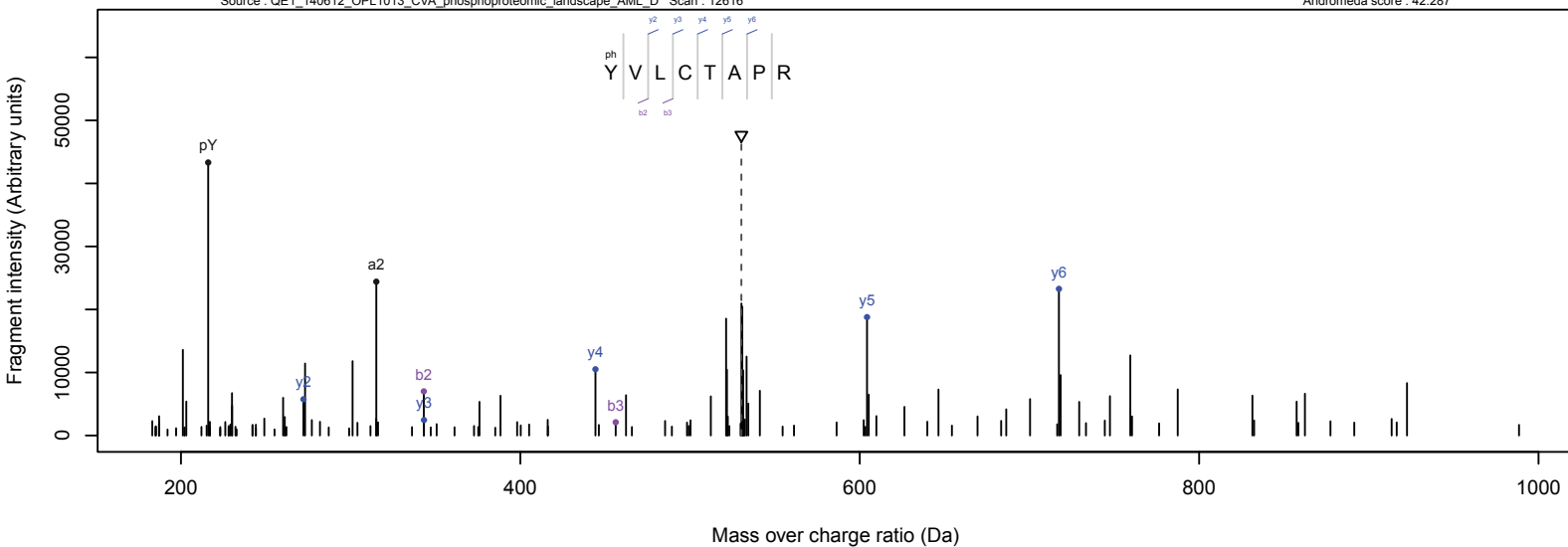
Andromeda score : 42.287



Y(ph)VLCTAPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12616

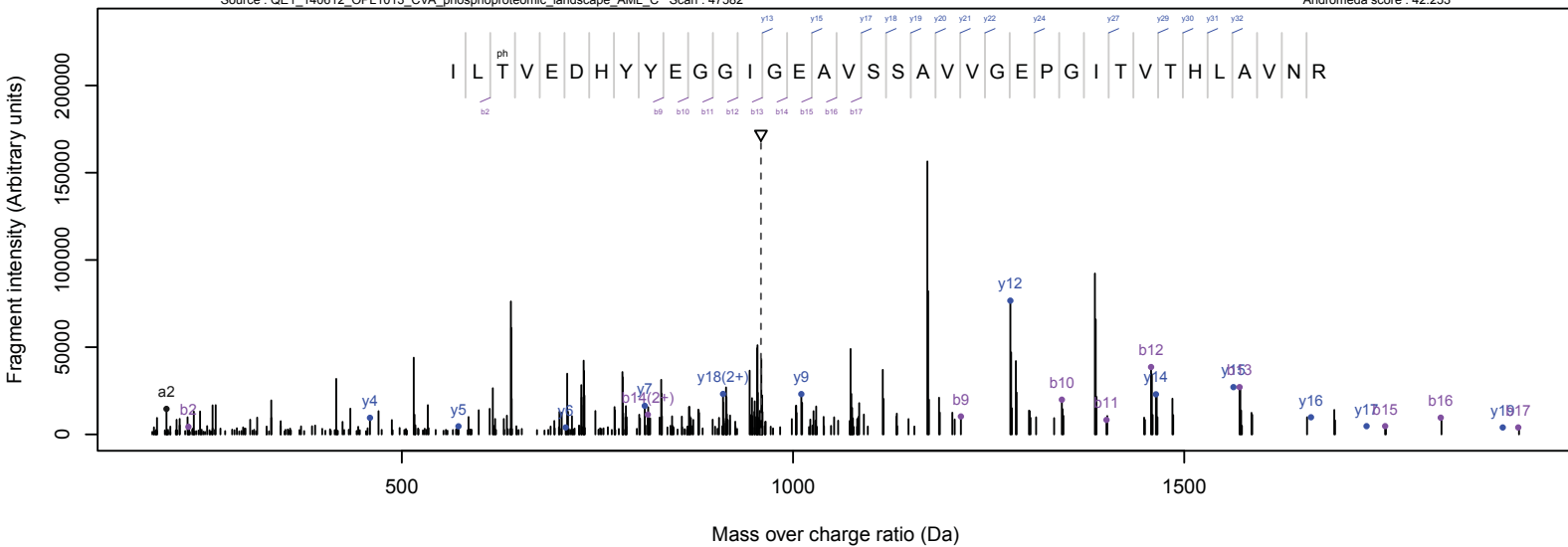
Andromeda score : 42.287



ILT(ph)VEDHYEGGIGEAVSSAVVGEPIVTHLAVNR-4+

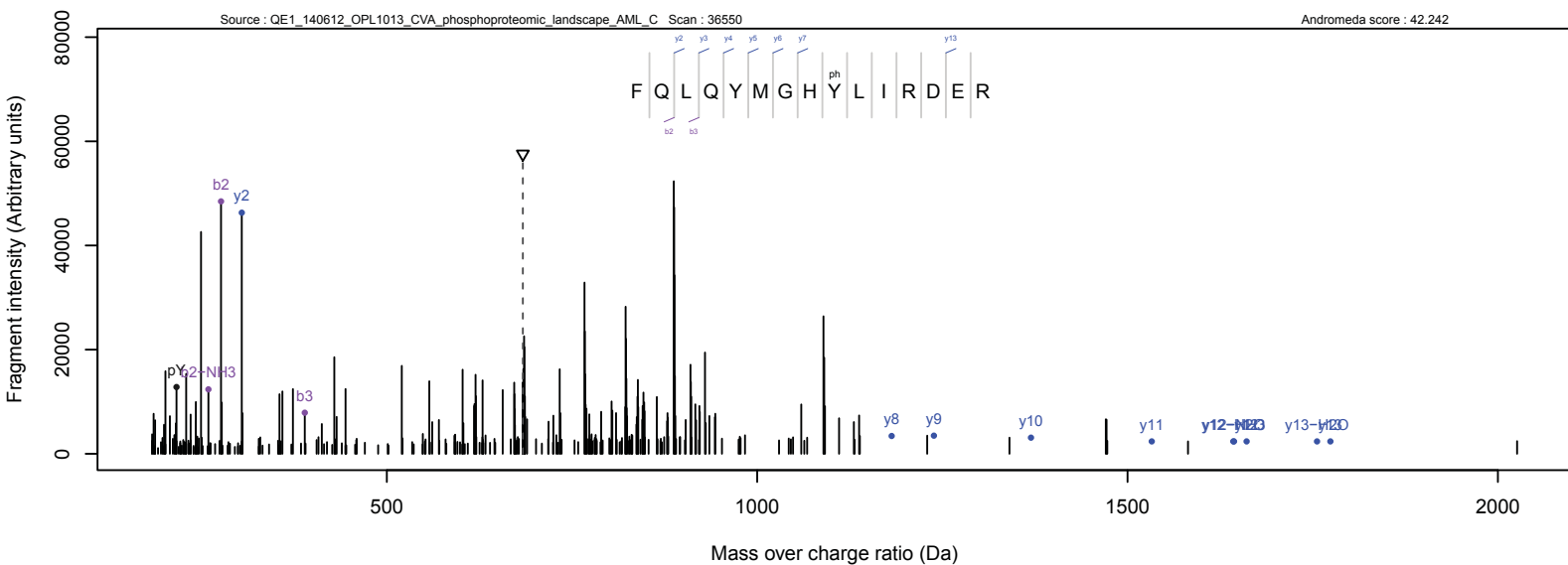
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 47582

Andromeda score : 42.253

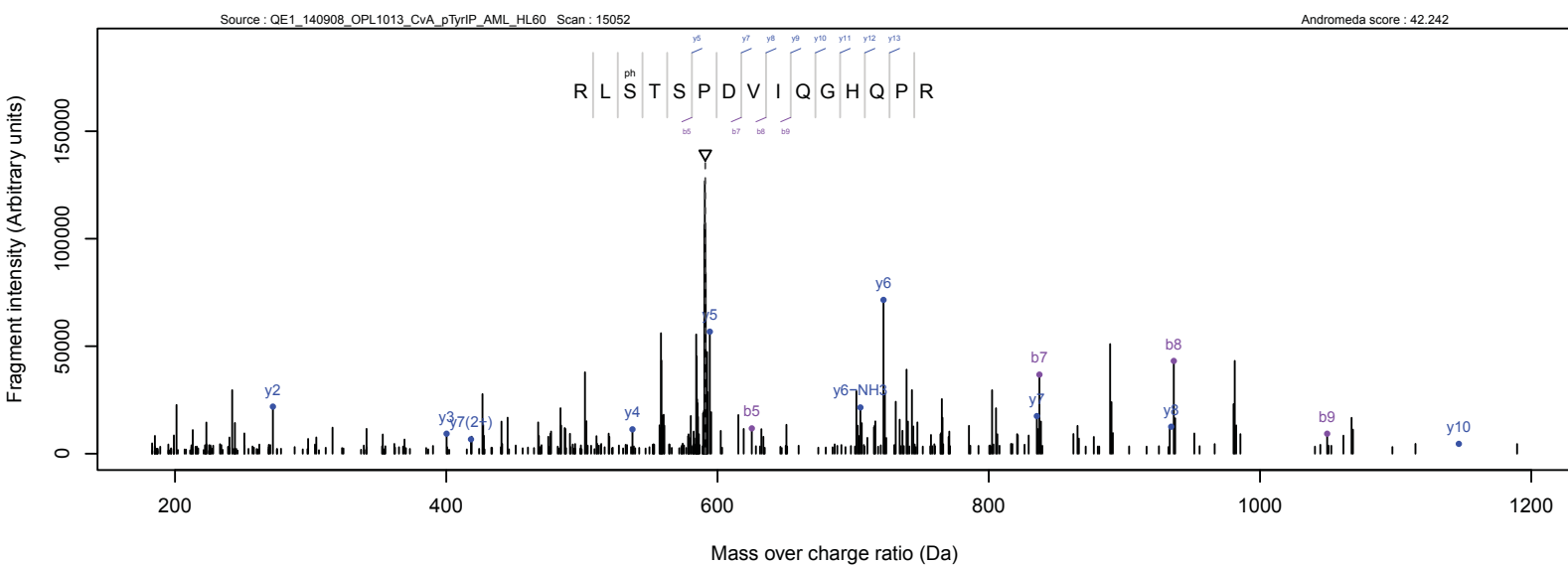




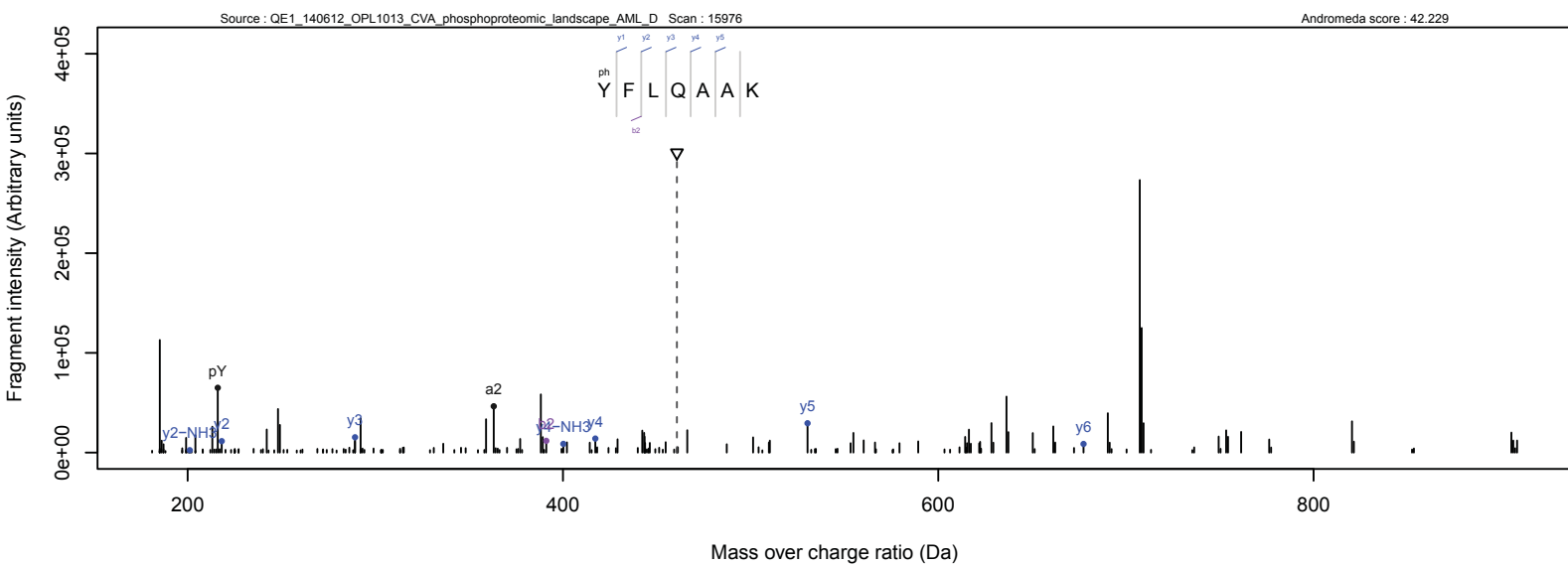
FQLQYMGHY(ph)LIRDER-3+



RLS(ph)TSPDVIQGHQPR-3+



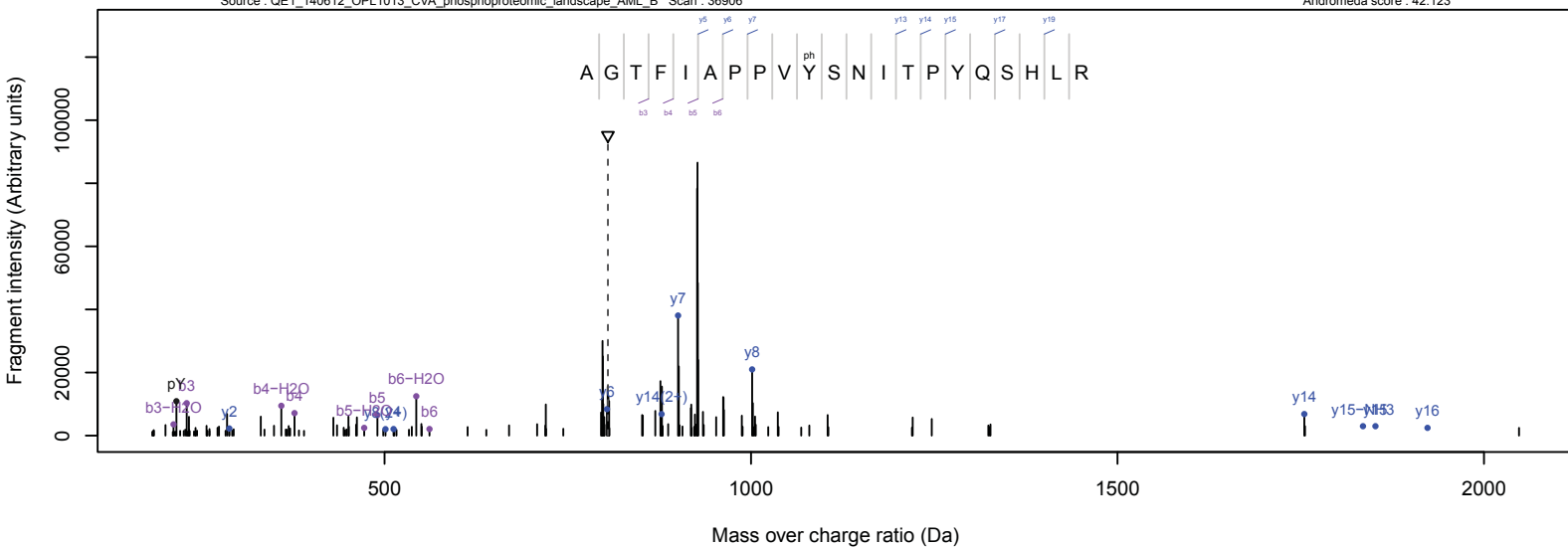
Y(ph)FLQAAK-2+



## AGTFIAPPVY(ph)SNITPYQSHLR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 36906

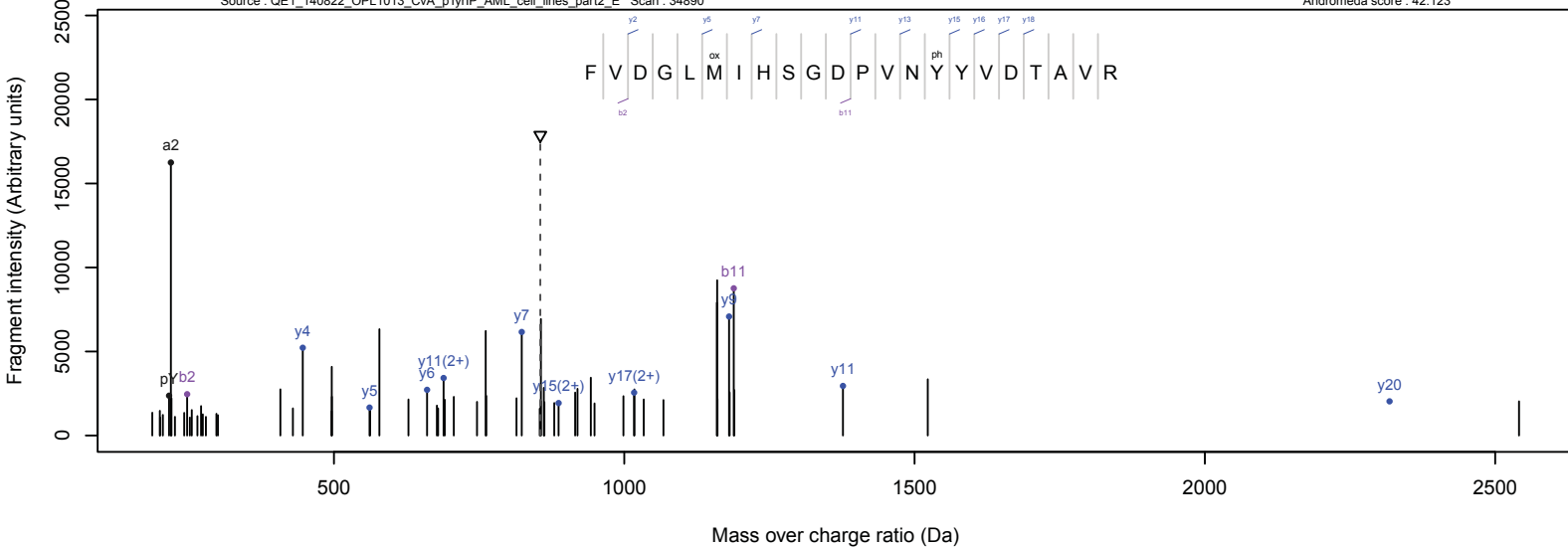
Andromeda score: 42.123



## FVDGLM(ox)IHSGDPVNY(ph)YVDTAVR-3+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 34890

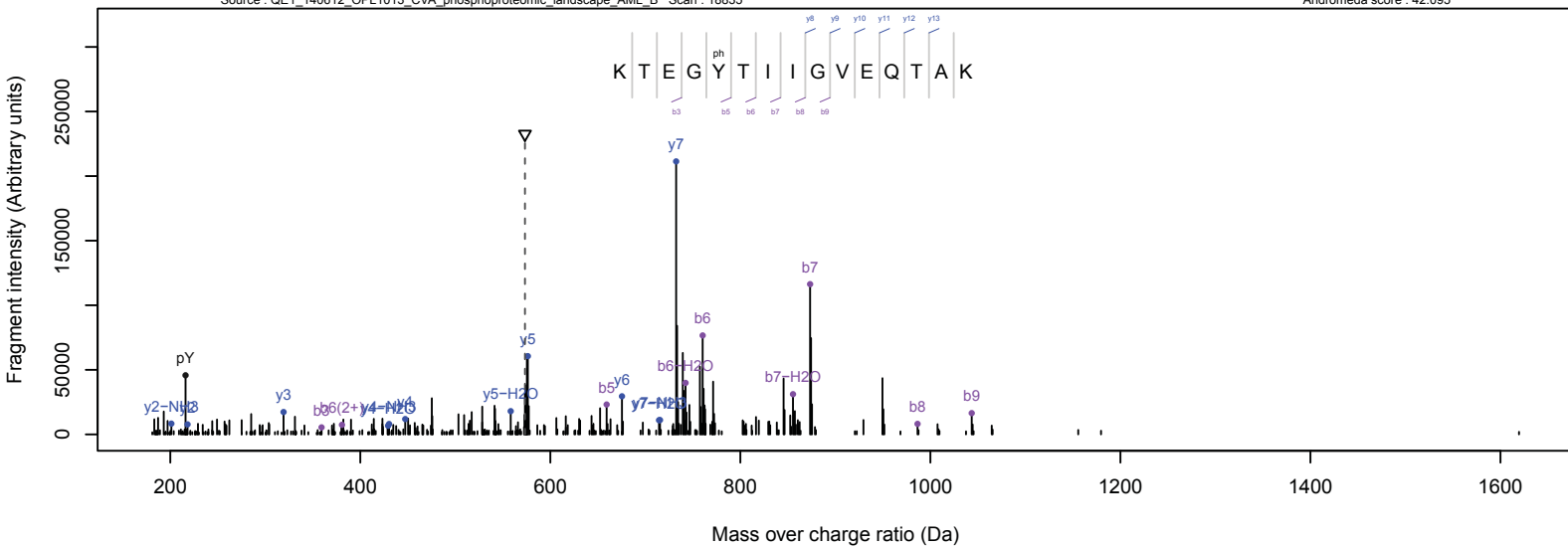
Andromeda score: 42.123



## KTEGY(ph)TIIGVEQTAK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18835

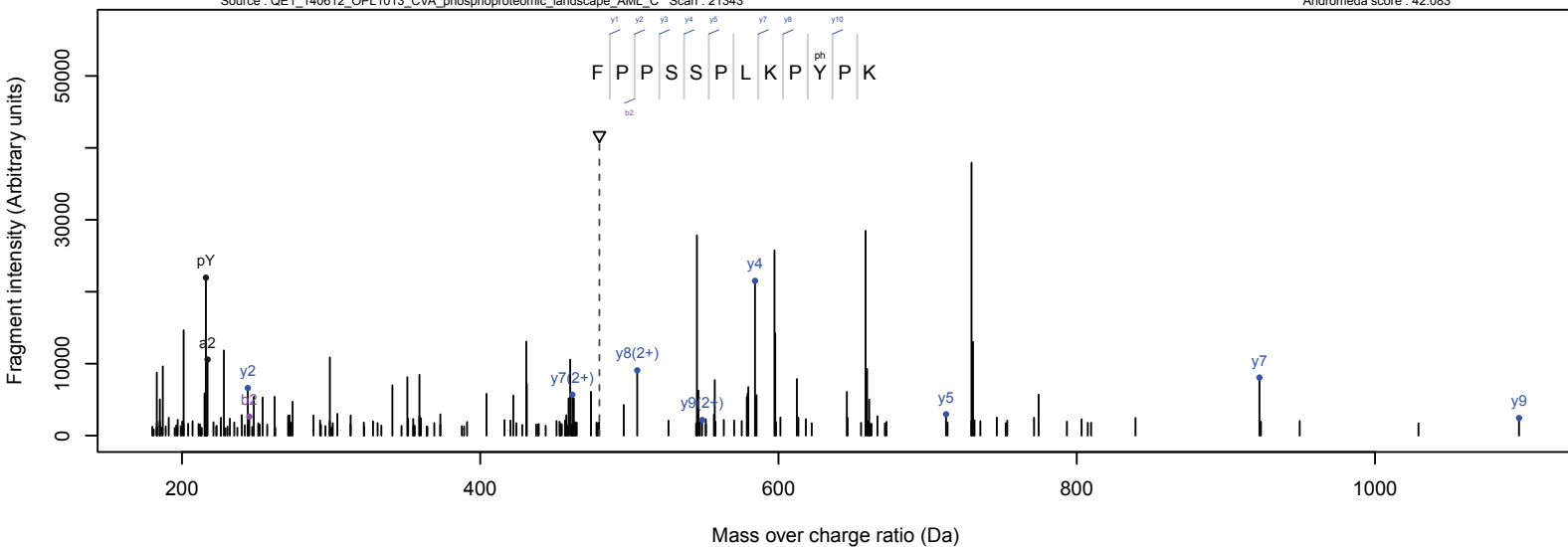
Andromeda score: 42.095



## FPSSPLKPY(ph)PK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21343

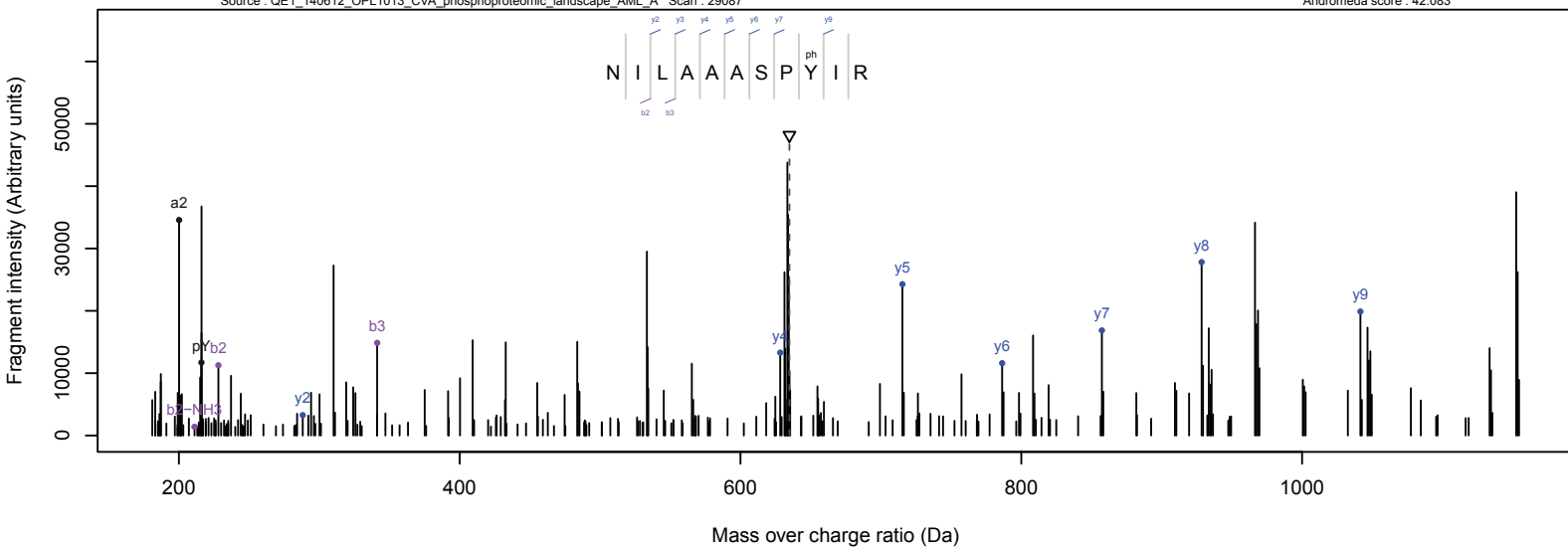
Andromeda score : 42.083



## NILAASPY(ph)IR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 29087

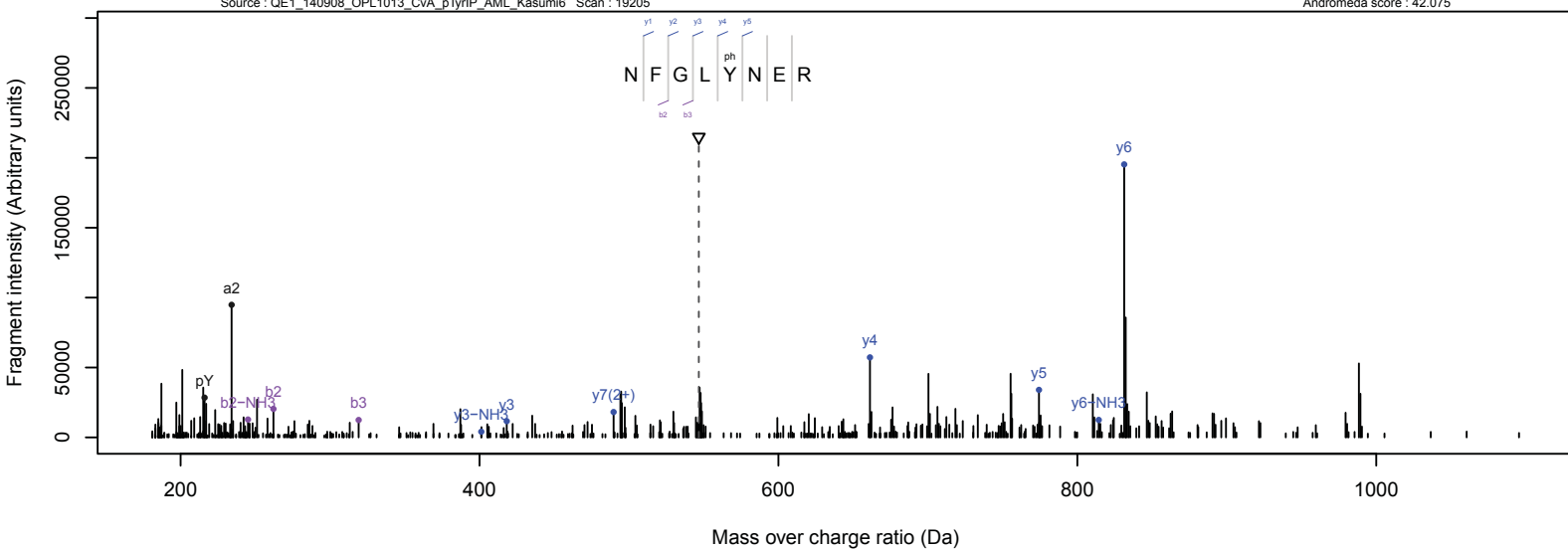
Andromeda score : 42.083



## NFGLY(ph)NER-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 19205

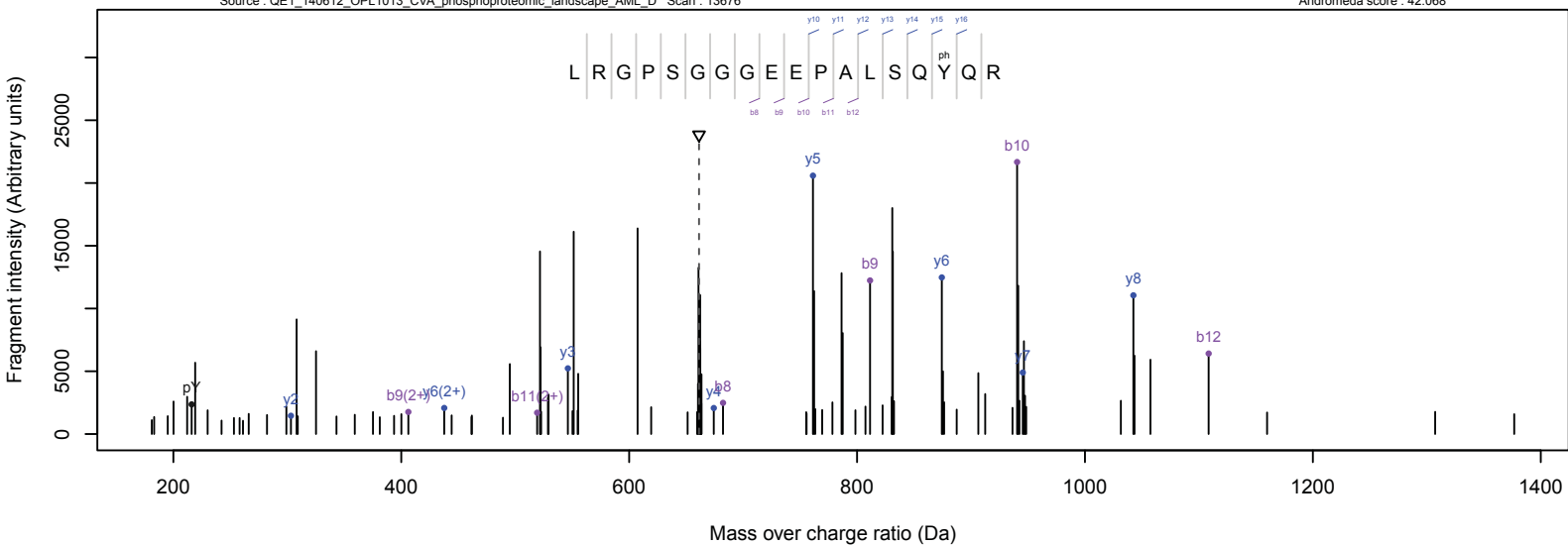
Andromeda score : 42.075



LRGPGSGGEEPALSYQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 13676

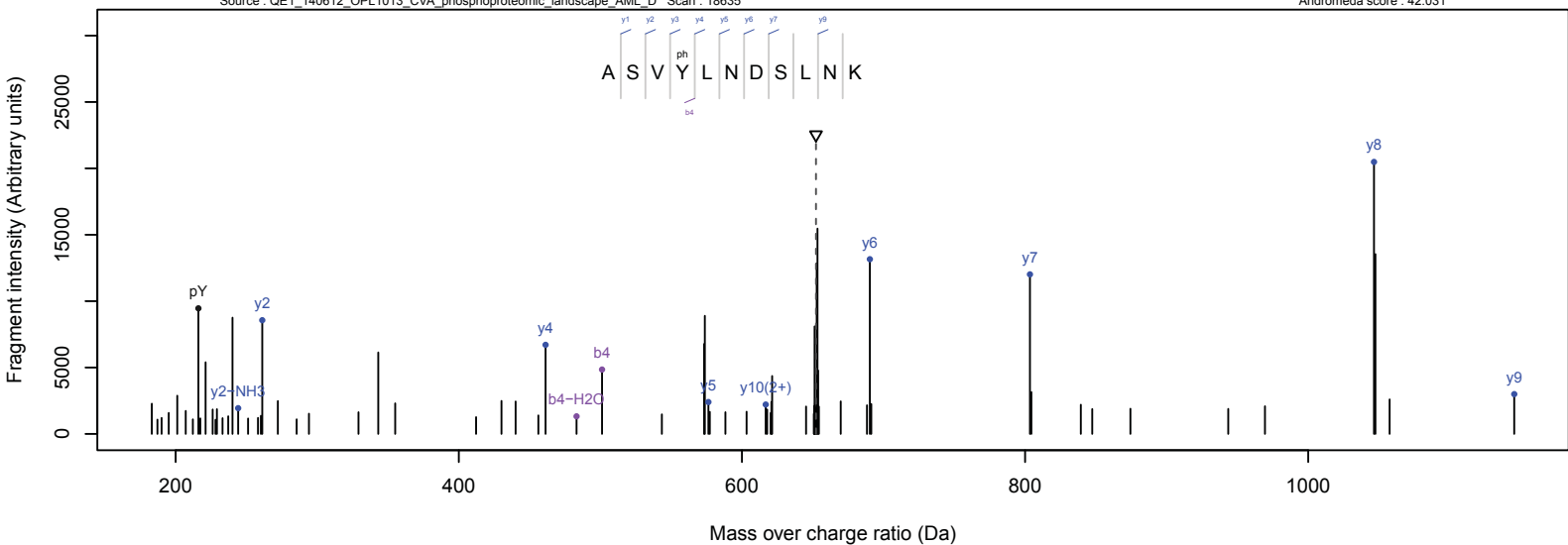
Andromeda score : 42.068



ASVY(ph)LNDSLNK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 18635

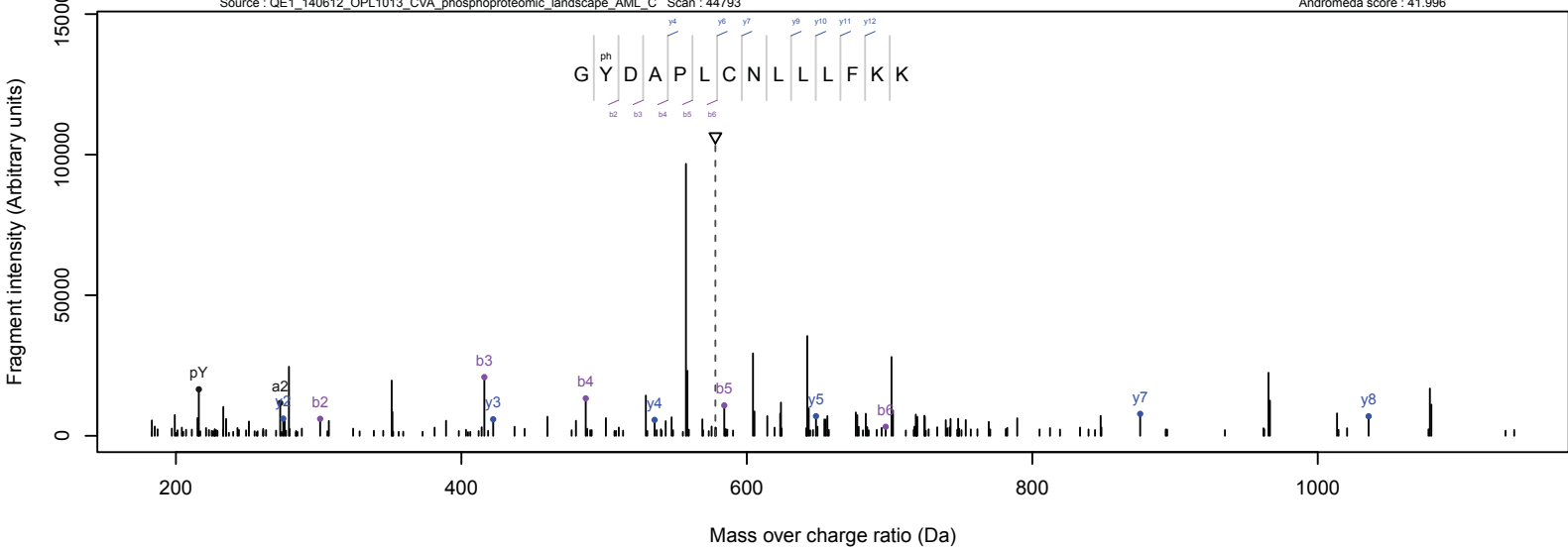
Andromeda score : 42.031

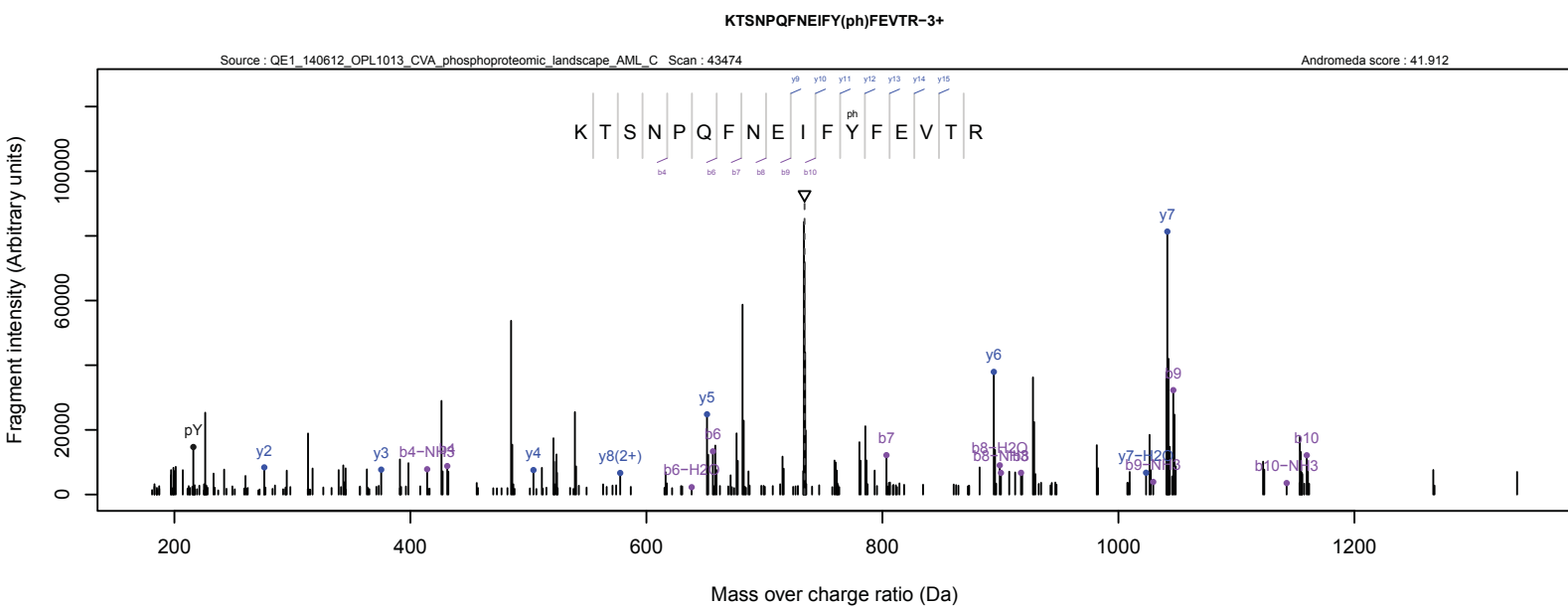
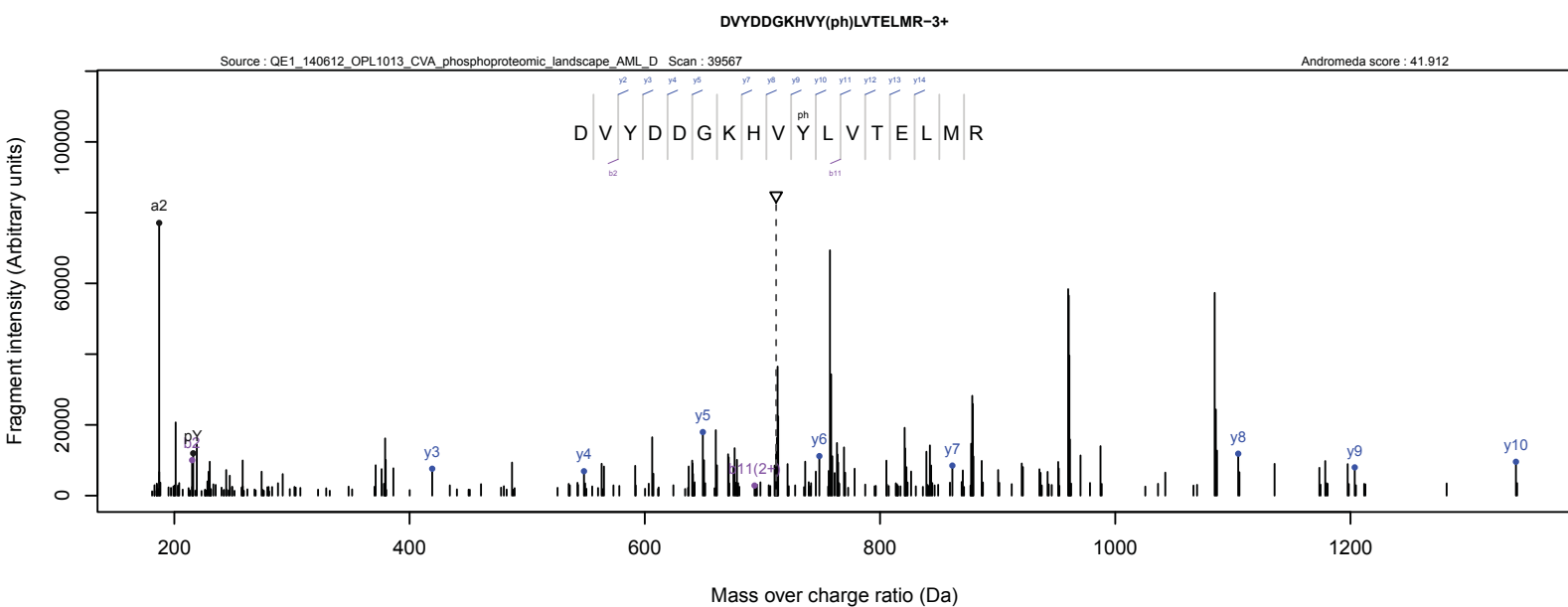
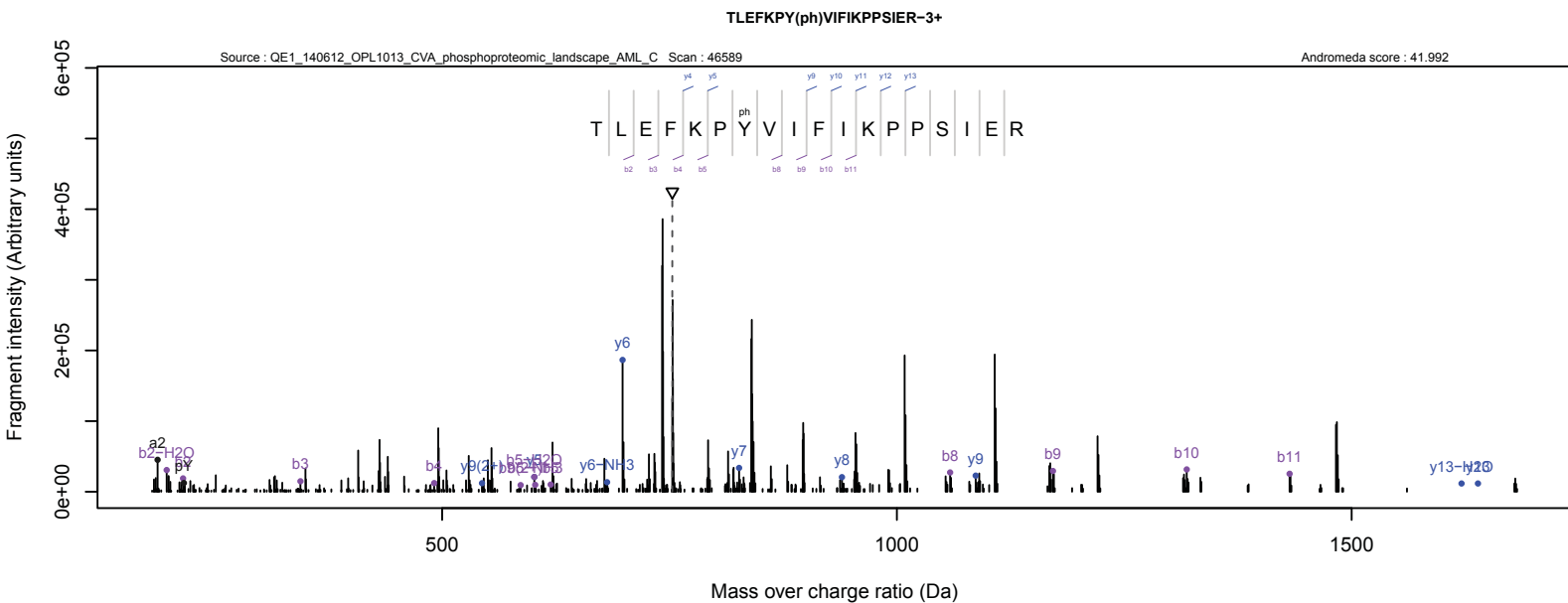


GY(ph)DAPLCLNLLFKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 44793

Andromeda score : 41.996

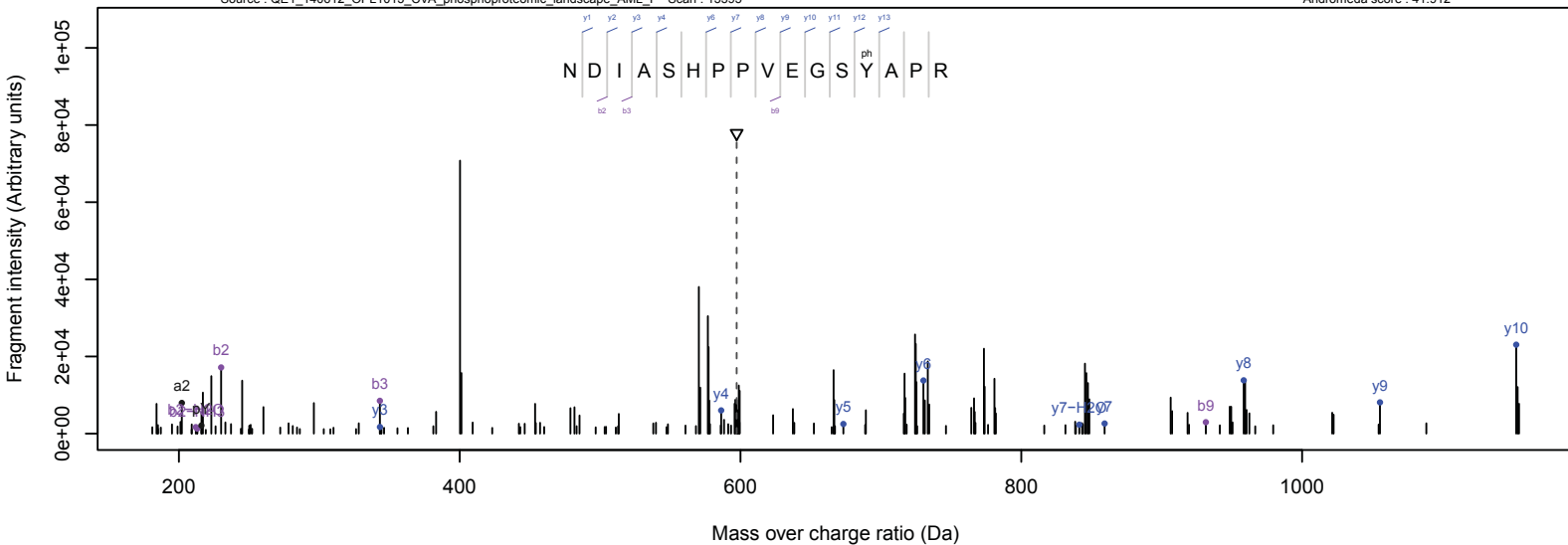




## NDIASHPPEGSY(ph)APR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_F Scan : 13395

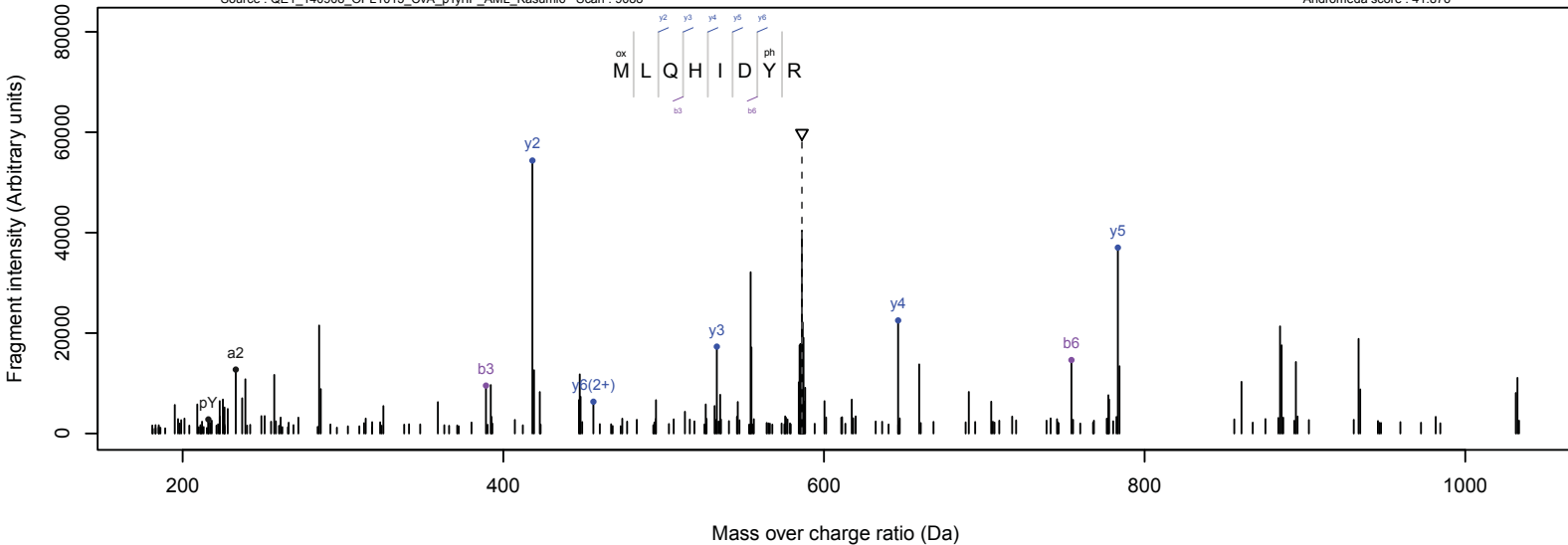
Andromeda score : 41.912



## M(ox)LQHIDY(ph)R-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 9088

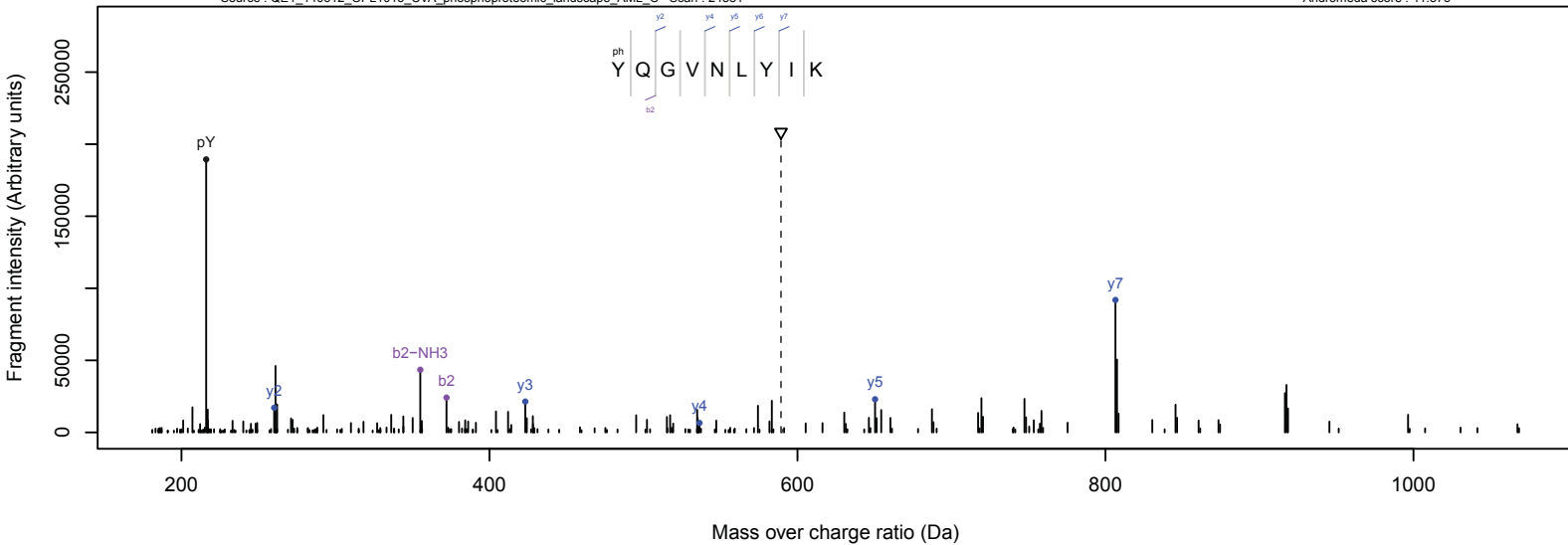
Andromeda score : 41.876



## Y(ph)QGVNLYIK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 24881

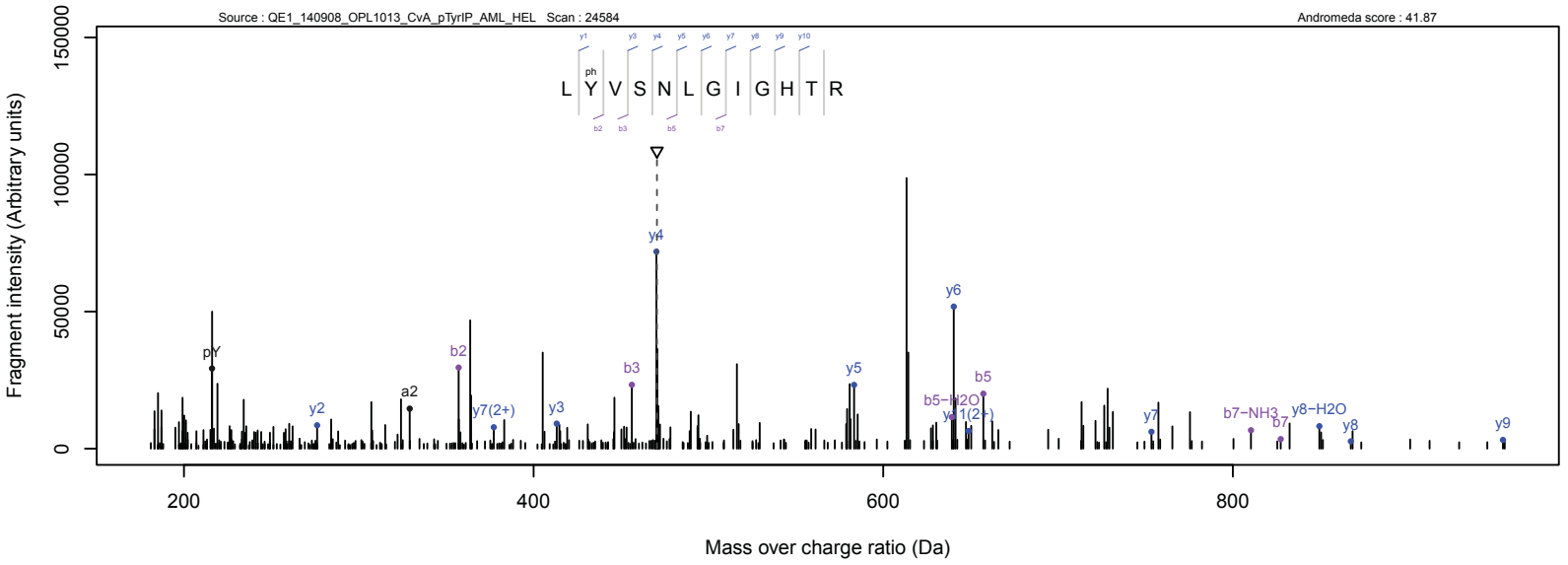
Andromeda score : 41.876



**LY(ph)VSNLGIGHTR-3+**

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 24584

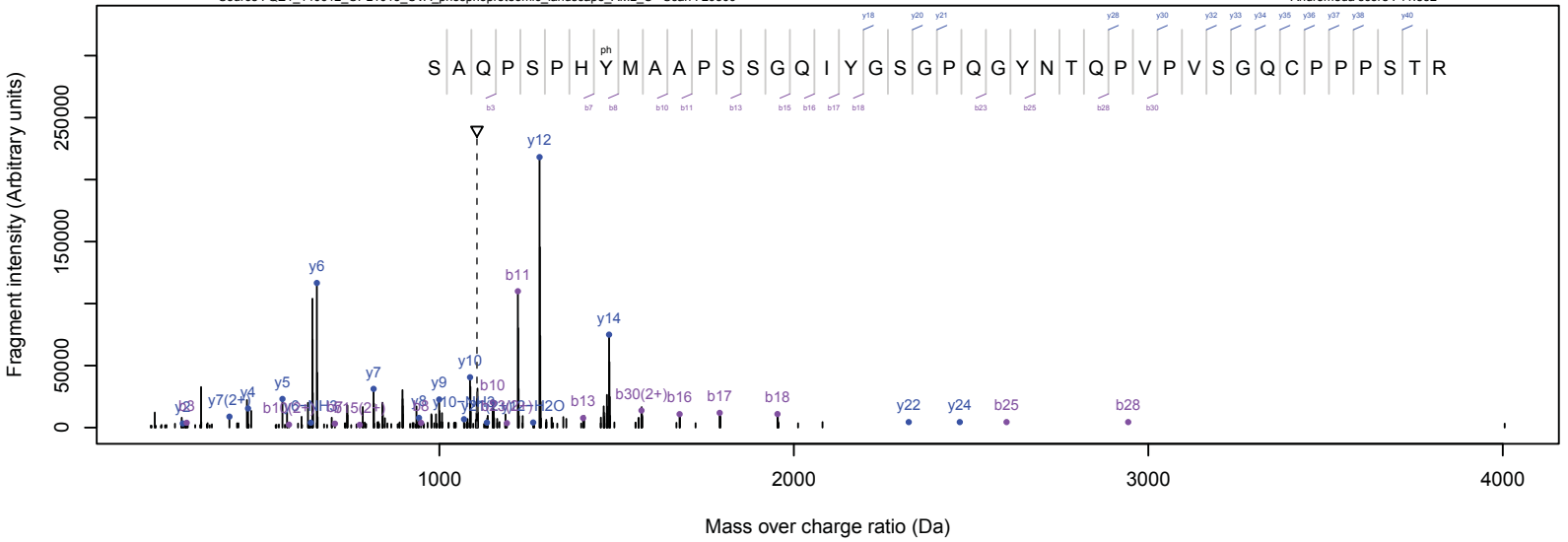
Andromeda score : 41.87



**SAQPSPHY(ph)MAAPSSGQIYGSGPQGYNTQPVPVSGQCPPPSTR-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29309

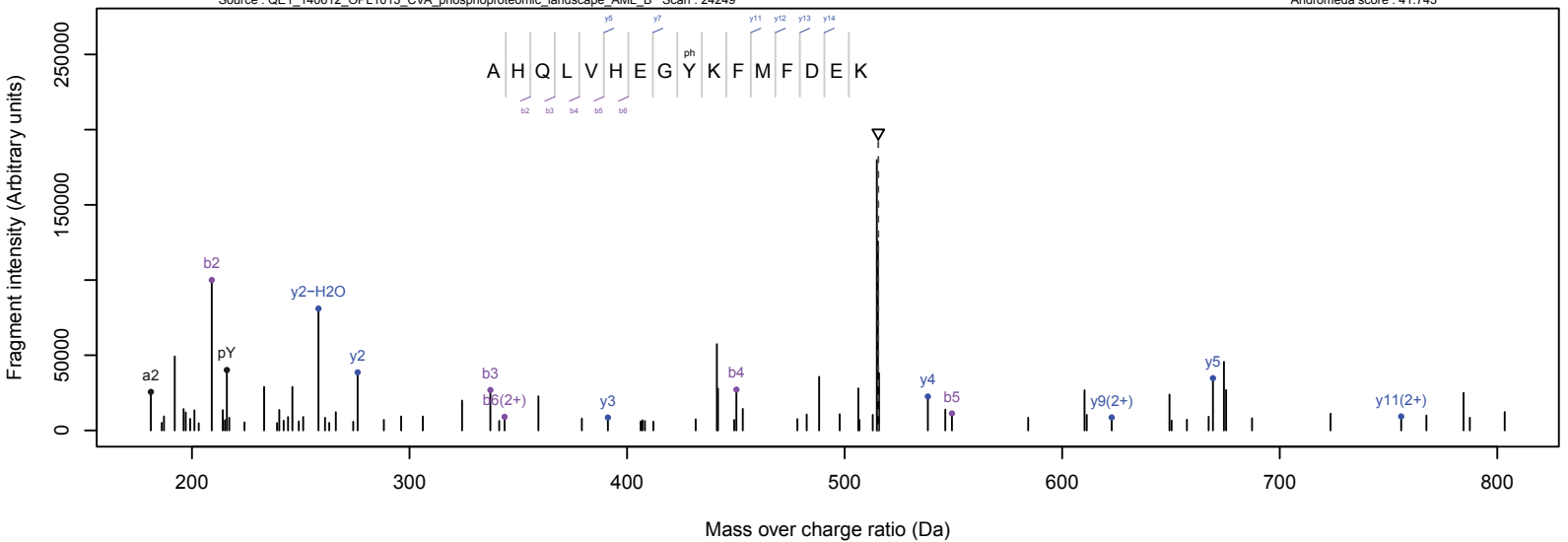
Andromeda score : 41.832



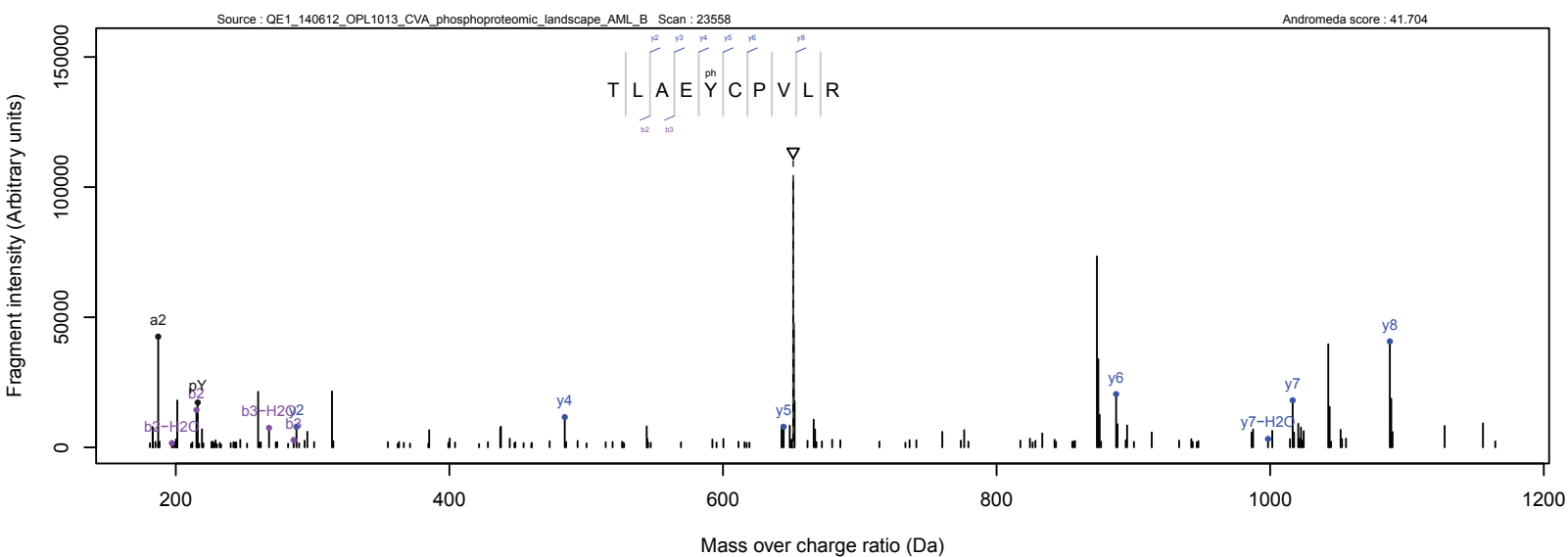
**AHQLVHEGY(ph)KFMFDEK-4+**

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 24249

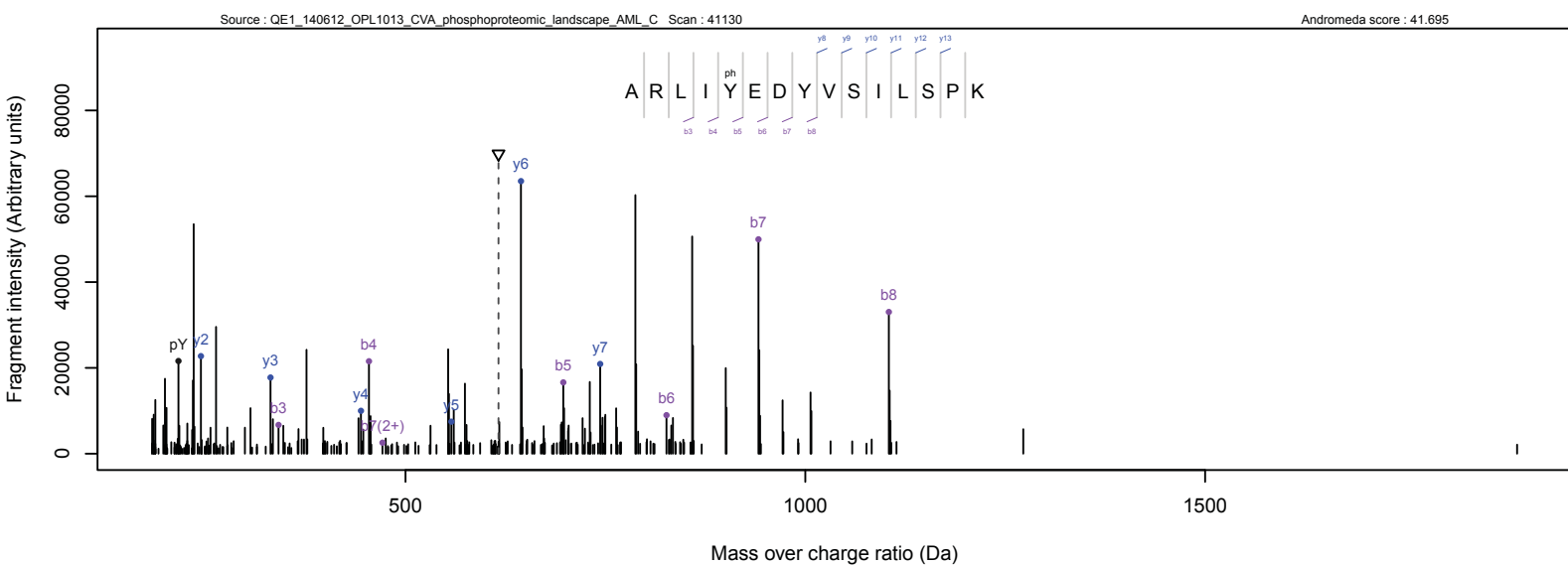
Andromeda score : 41.743



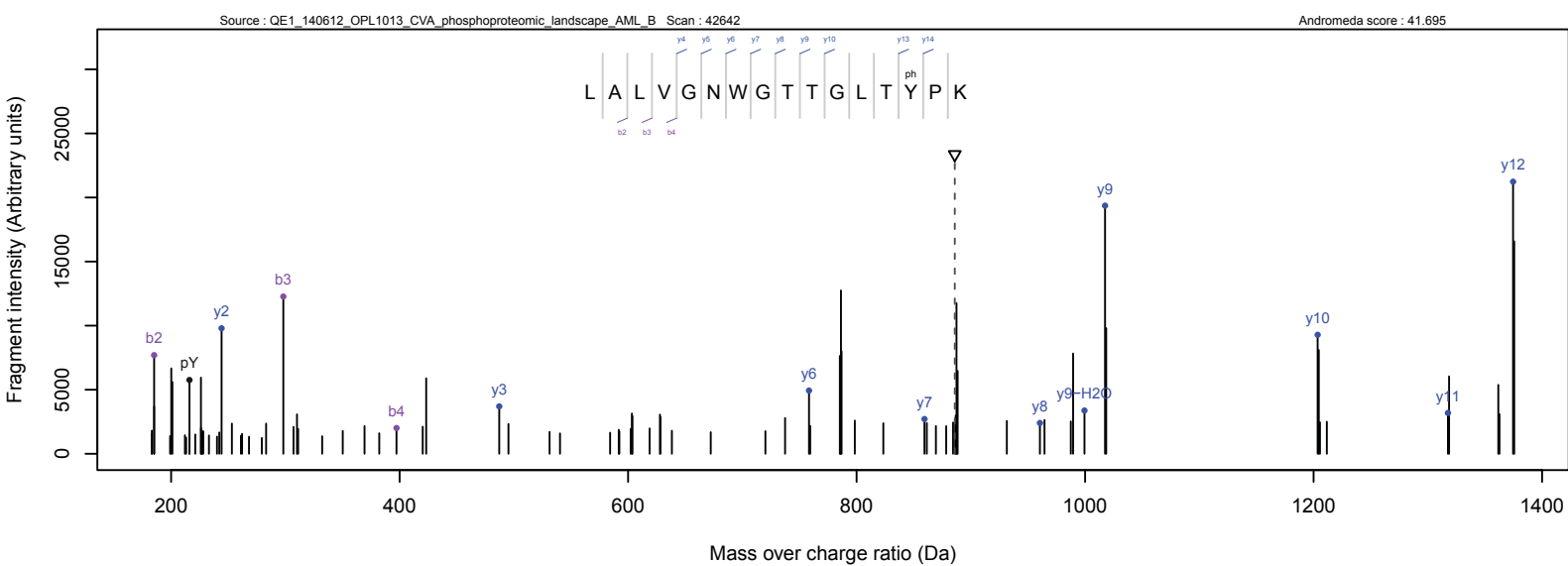
## TLAEY(ph)CPVLR-2+



## ARLIY(ph)EDYVSILSPK-3+



## LALVGNWGTGLTY(ph)PK-2+

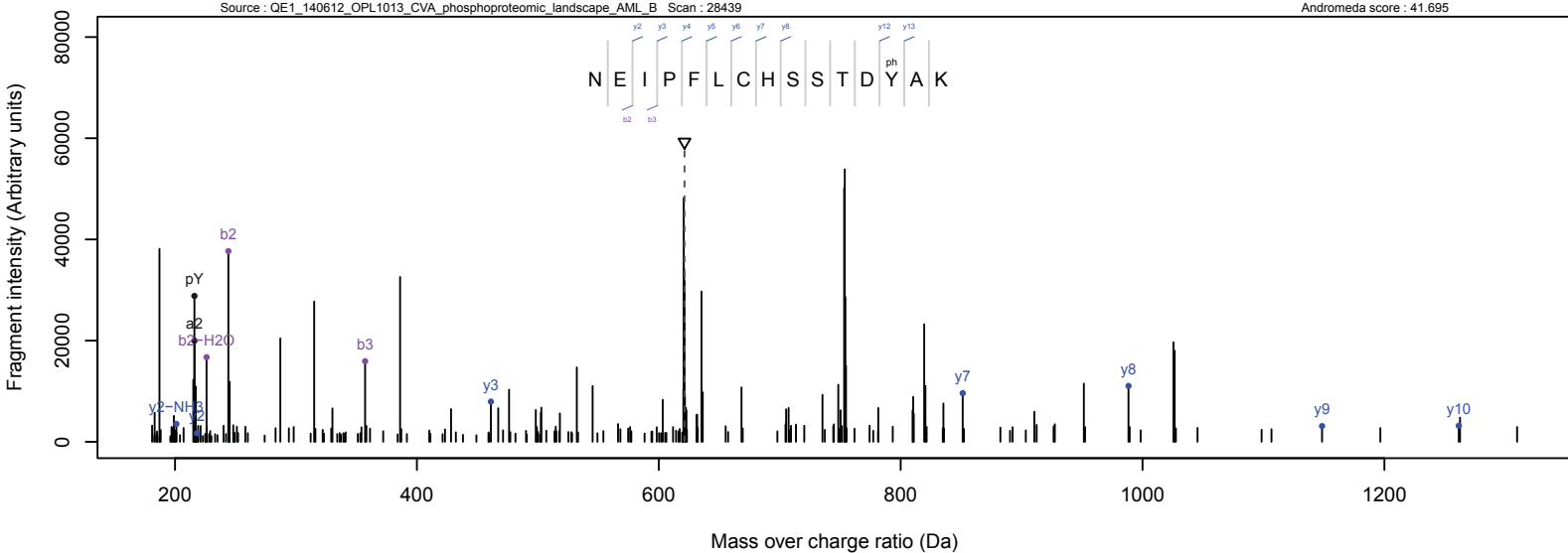




## NEIPFLCHSSTDY(ph)AK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 28439

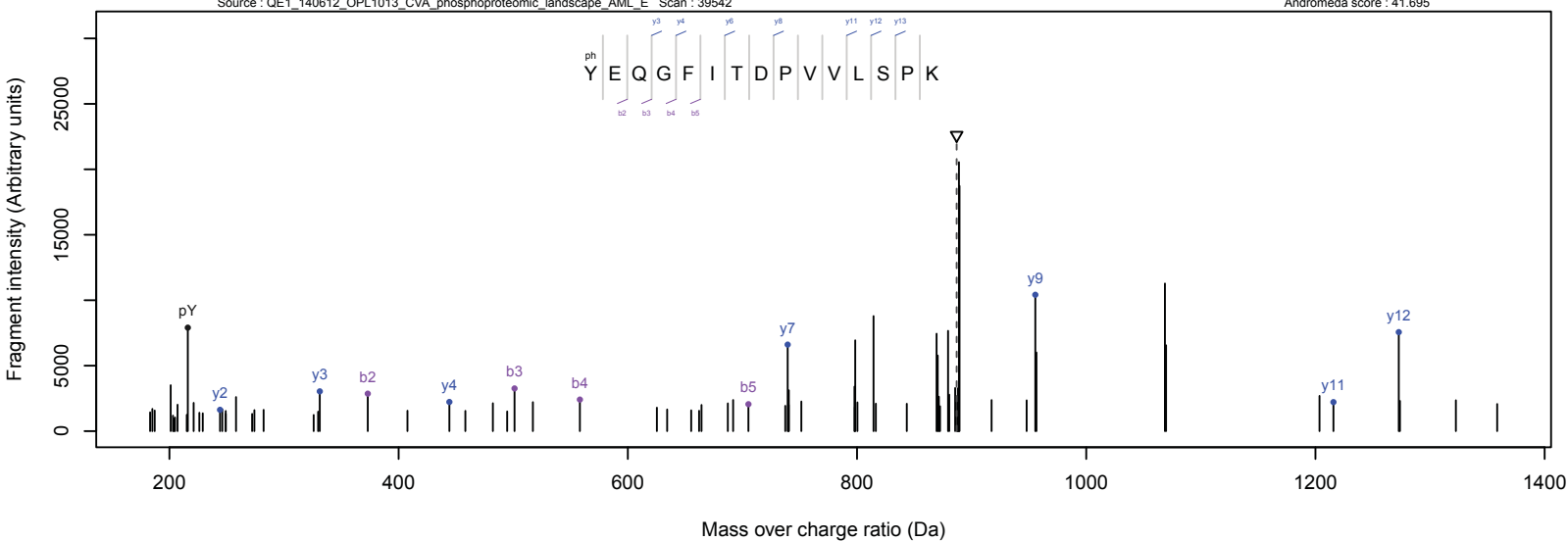
Andromeda score: 41.695



## Y(ph)EQGFITDPVVLSPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan: 39542

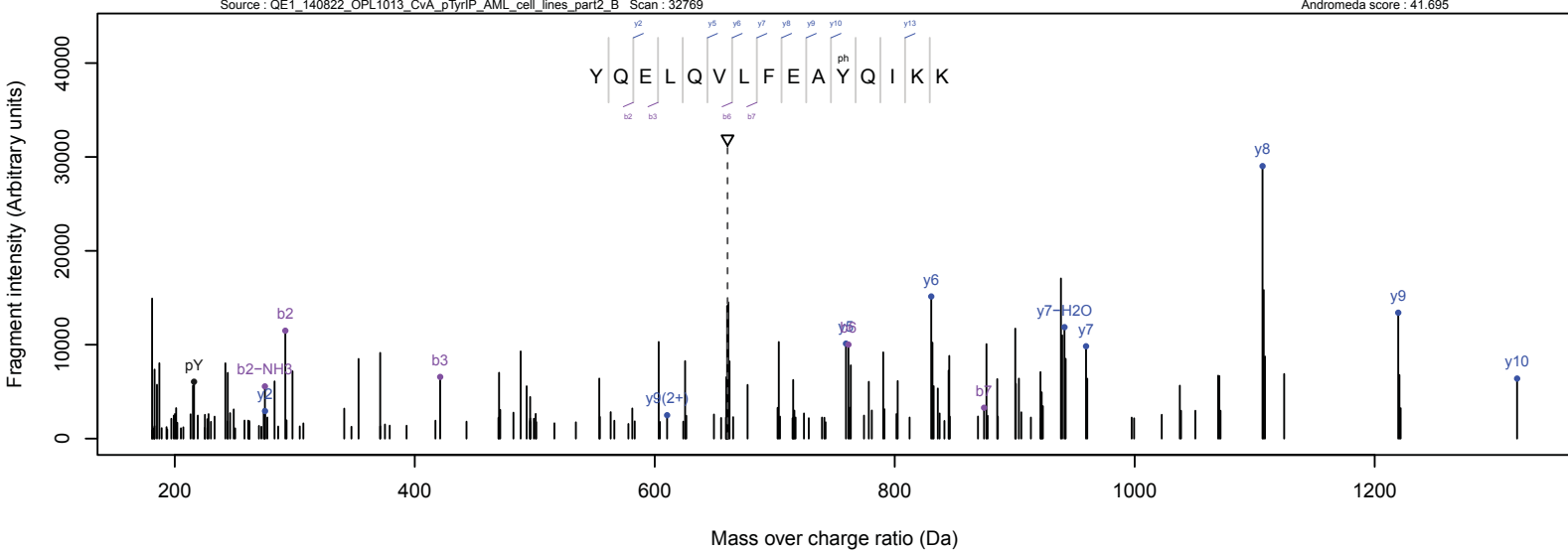
Andromeda score: 41.695



## YQELQVLF EAYQIKK-3+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan: 32769

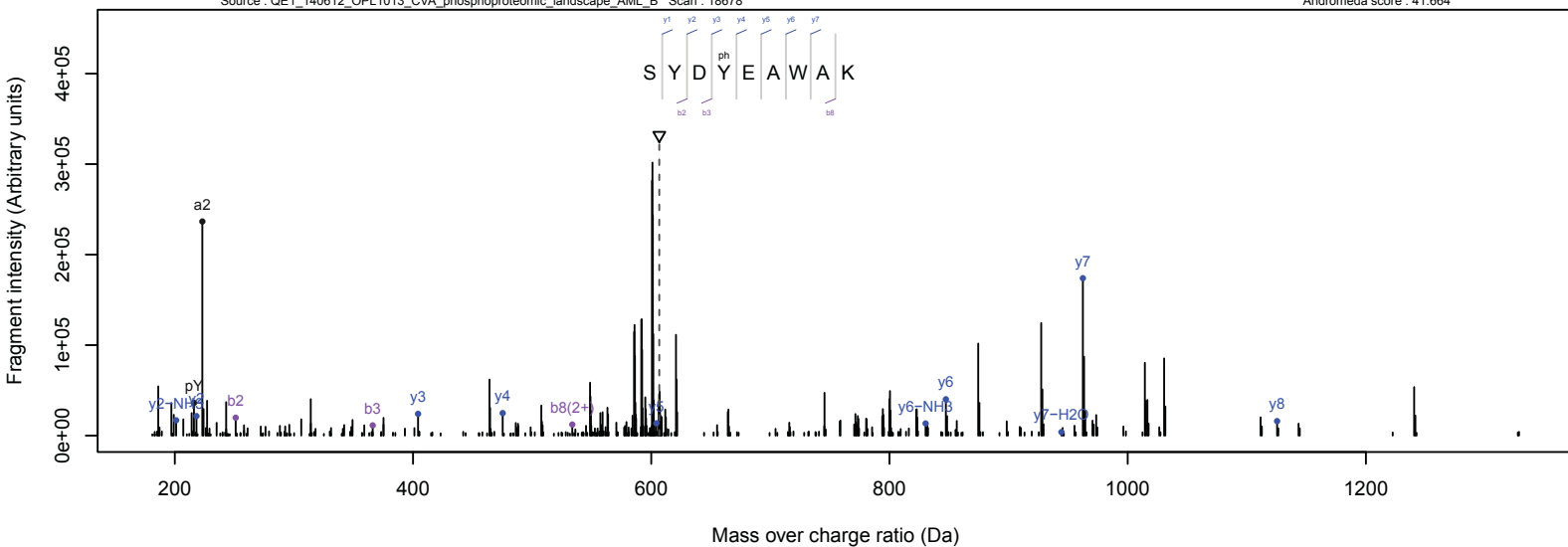
Andromeda score: 41.695



## SYDY(ph)EAWAK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 18678

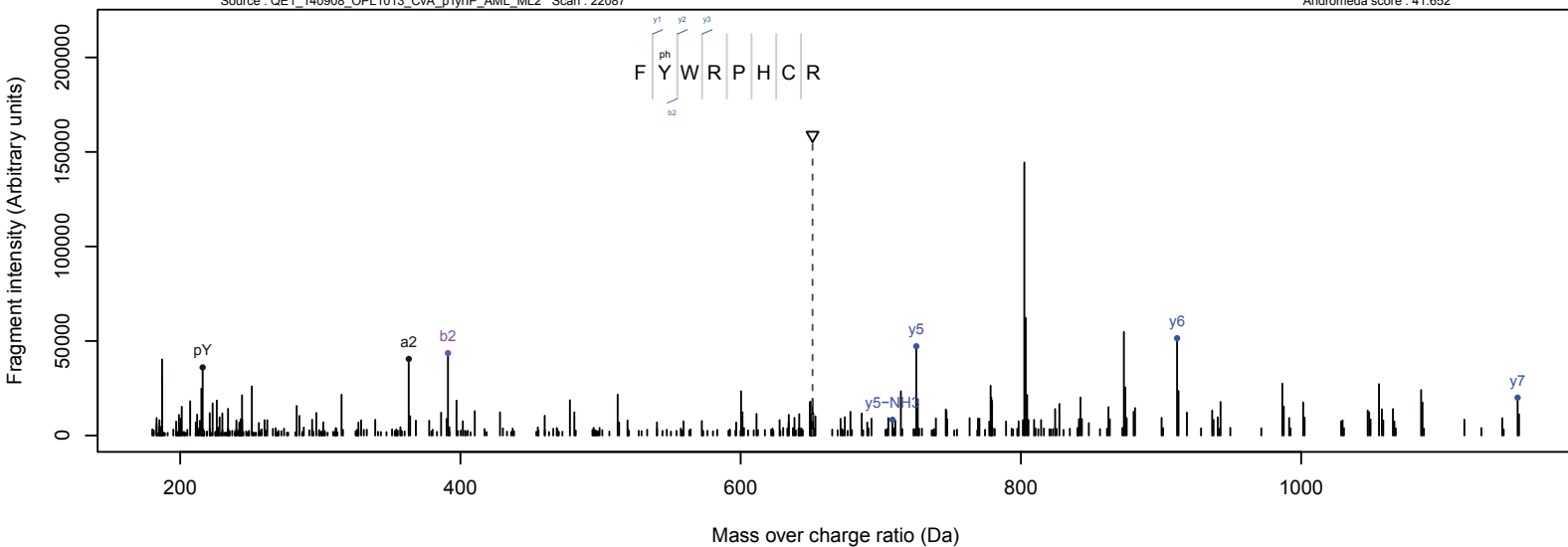
Andromeda score: 41.664



## FY(ph)WRPHCR-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan: 22087

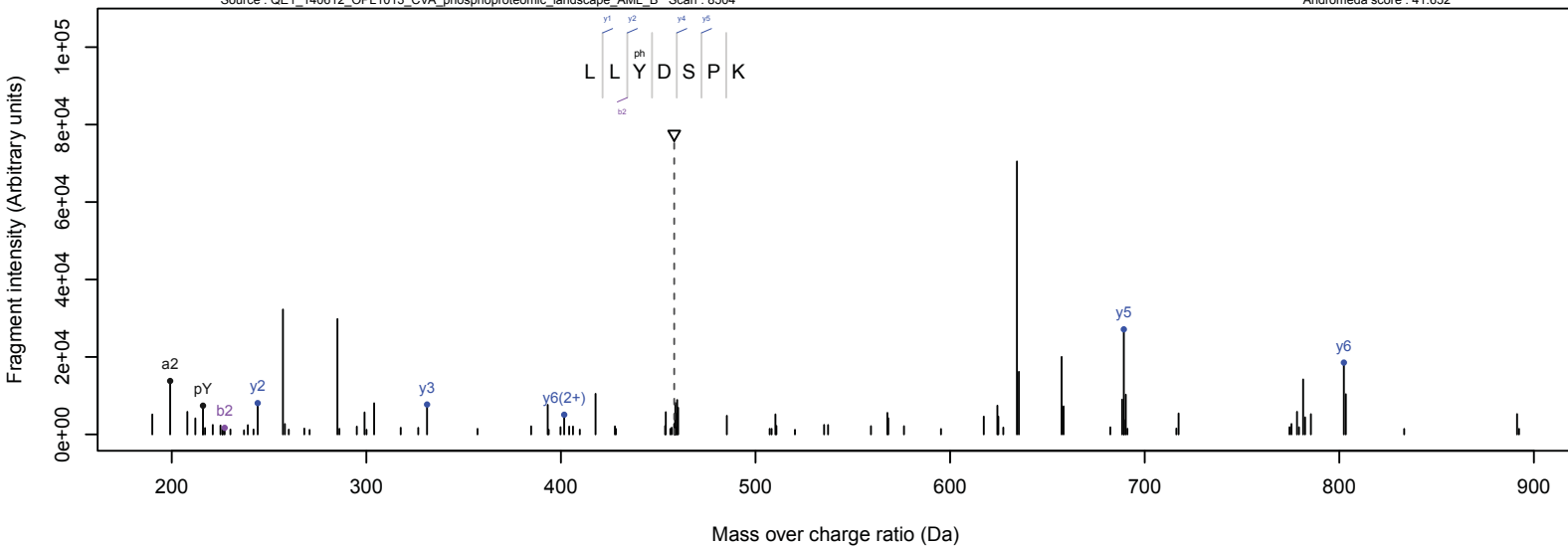
Andromeda score: 41.652



## LLY(ph)DSPK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 8504

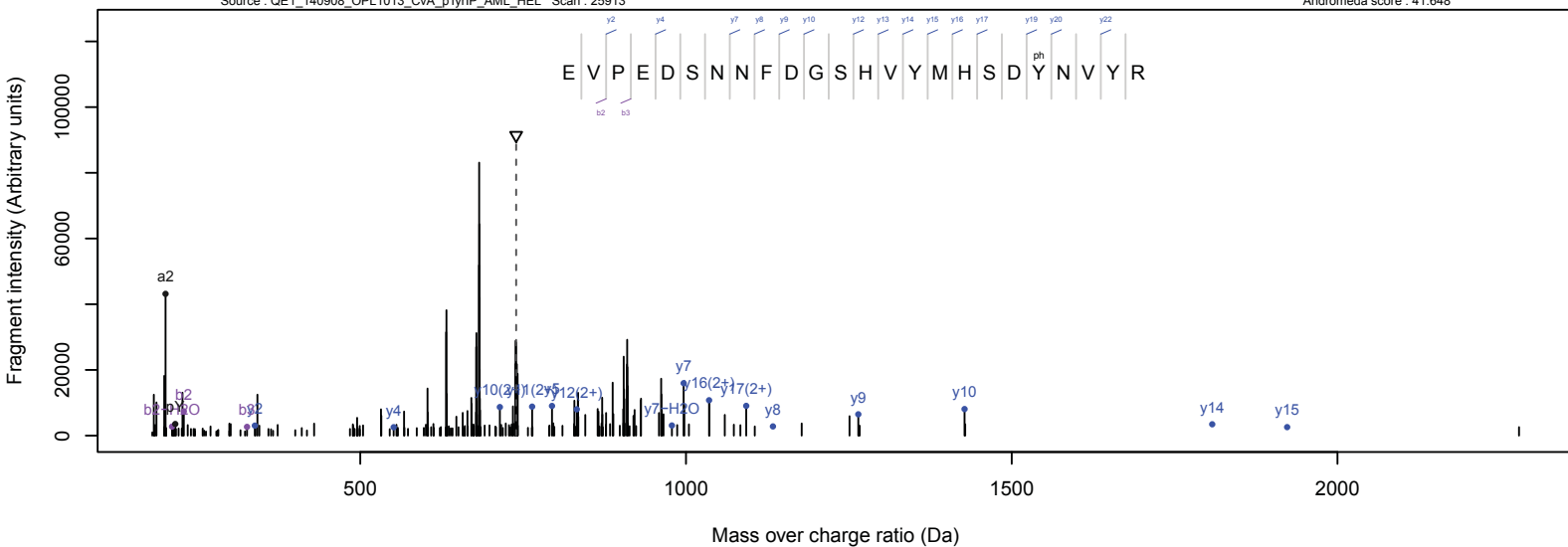
Andromeda score: 41.652



EVPEDSNNFDGSHVYMHSY(ph)NVYR-4+

Source : QE1\_140908\_OPL1013\_CVA\_pTyrIP\_AML\_HEL Scan : 25913

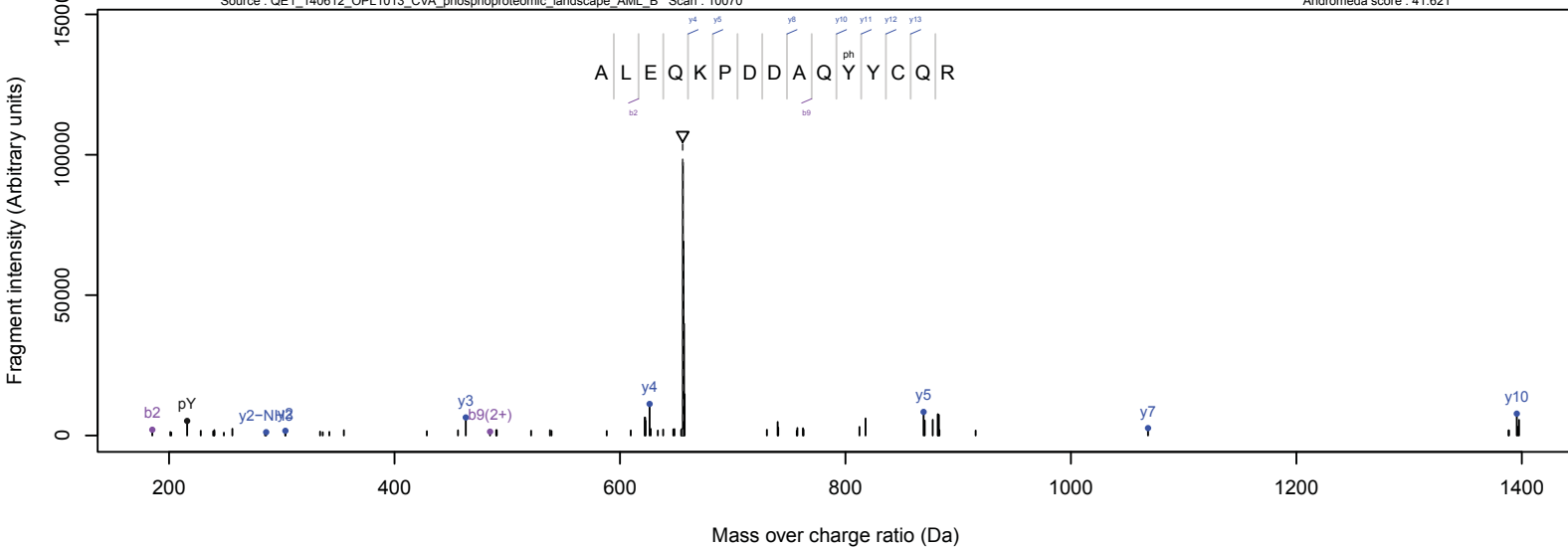
Andromeda score : 41.648



ALEQKPDDAQY(ph)YCQR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 10070

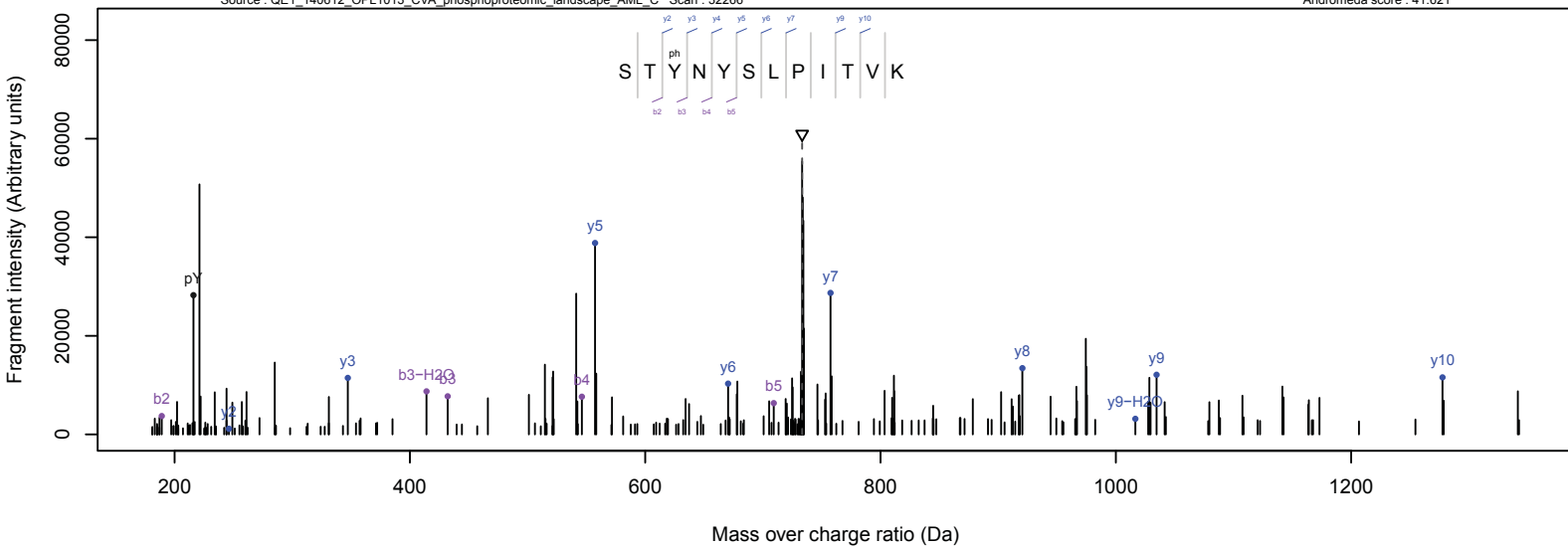
Andromeda score : 41.621



STY(ph)NYSLPITVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 32266

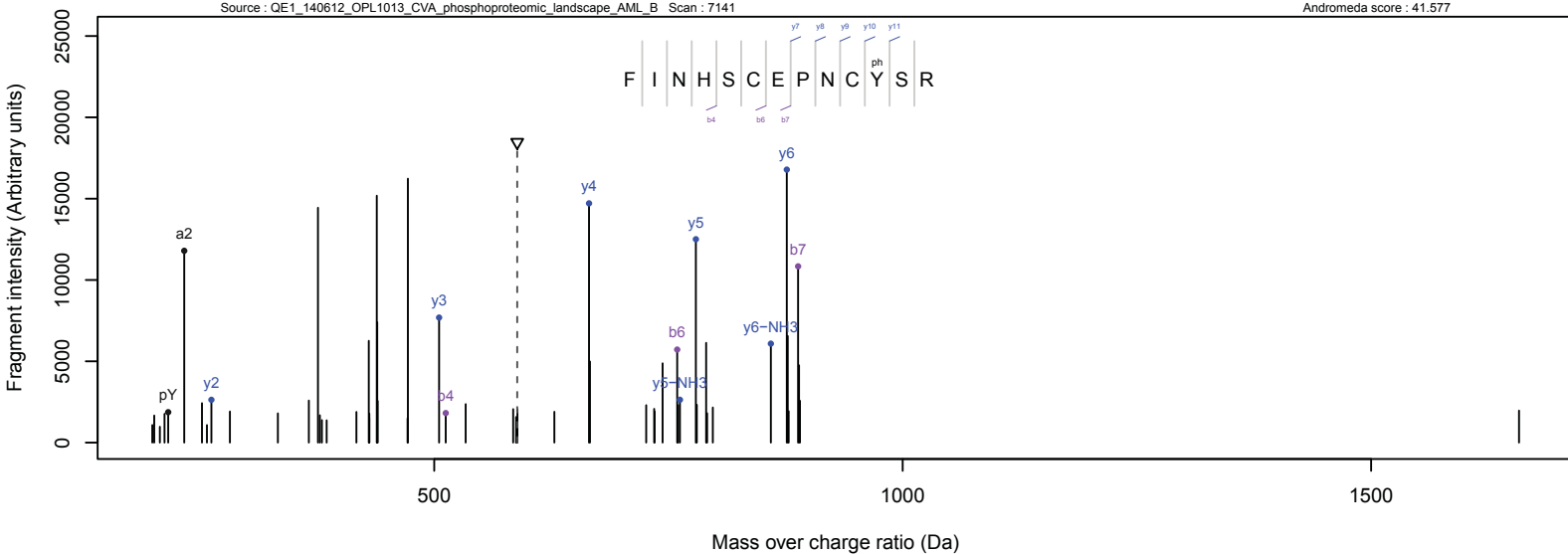
Andromeda score : 41.621



FINHSCEPN<sup>ph</sup>YSR-3+

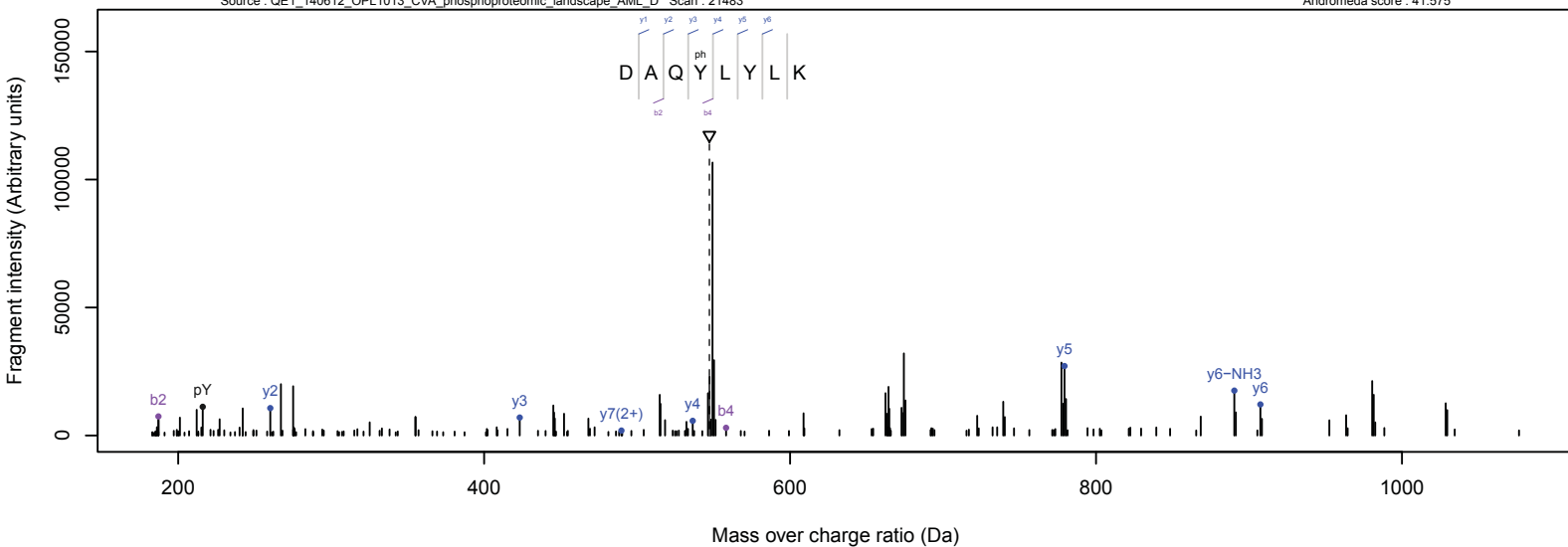
Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 7141

Andromeda score: 41.577

DAQY<sup>ph</sup>LYLK-2+

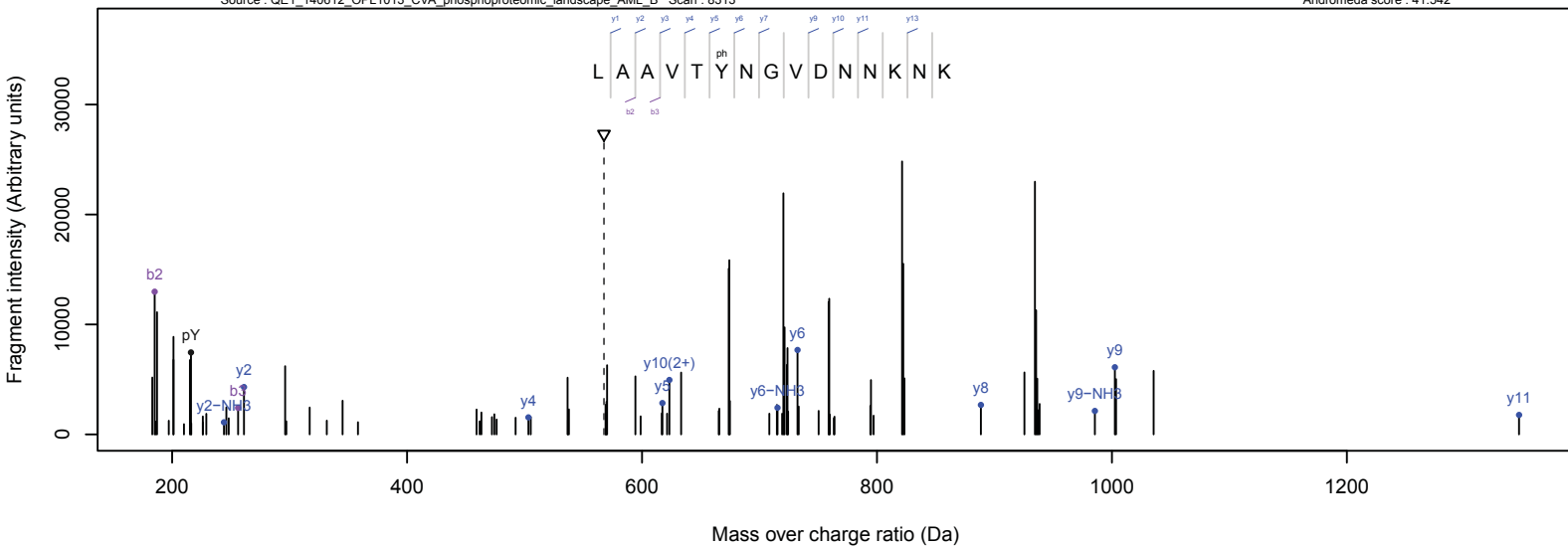
Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 21483

Andromeda score: 41.575

LAAVTY<sup>ph</sup>NGVDNKNK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 8313

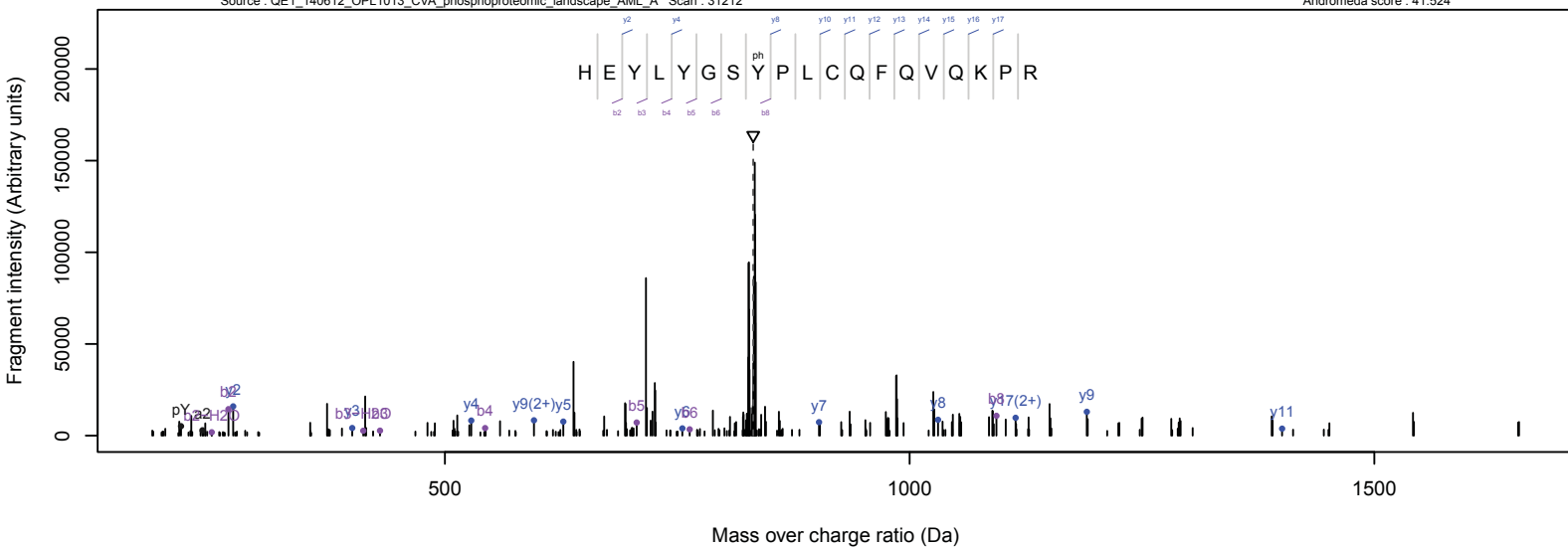
Andromeda score: 41.542



## HEYLYGSY(ph)PLCQFVQKPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 31212

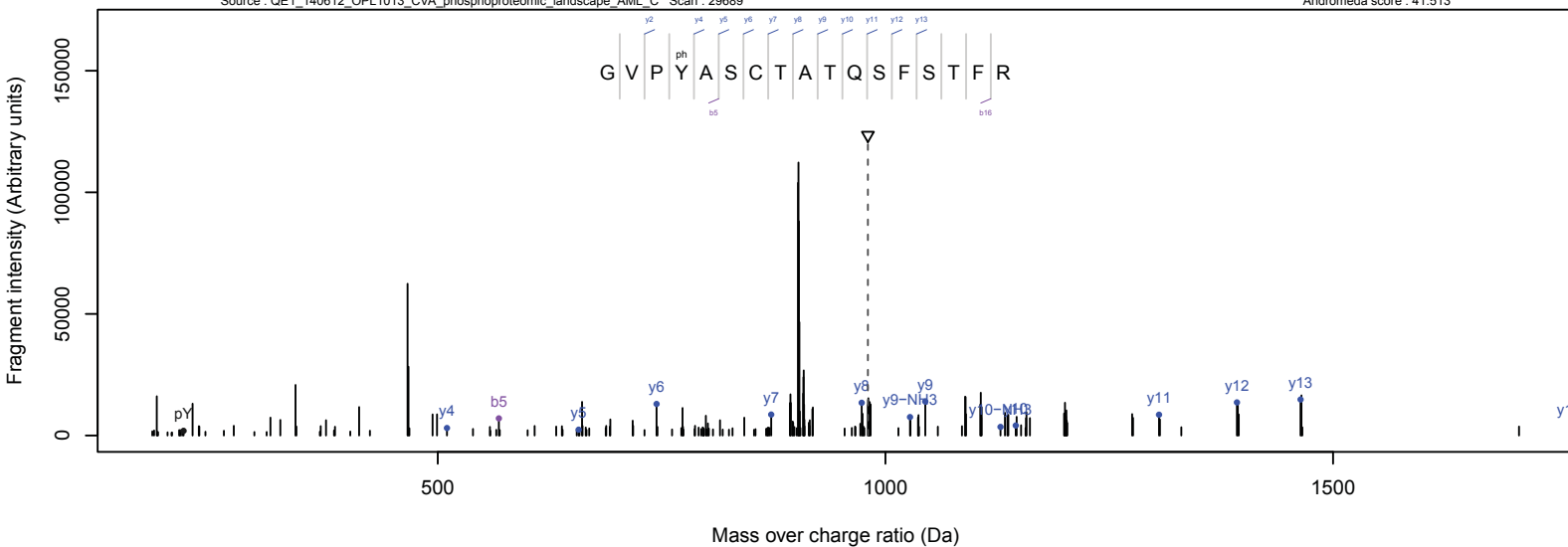
Andromeda score : 41.524



## GVPLYASCTATQSFSTR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29689

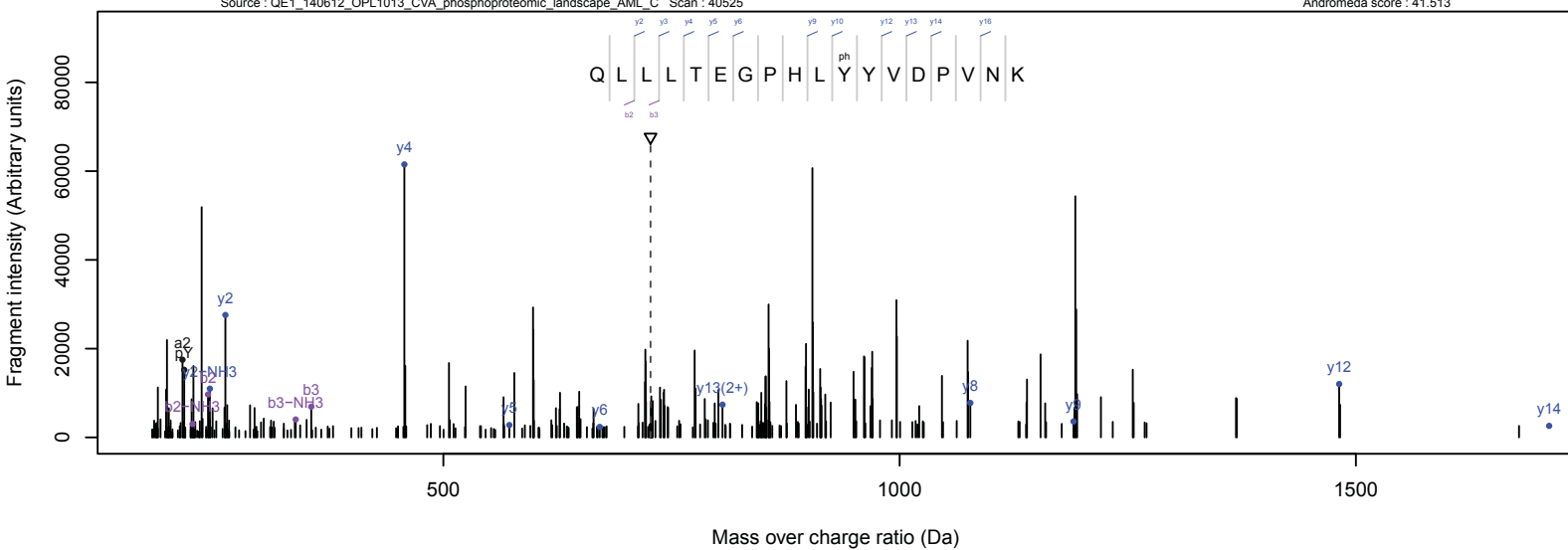
Andromeda score : 41.513



## QLLLETPHLYYVDPVVK-3+

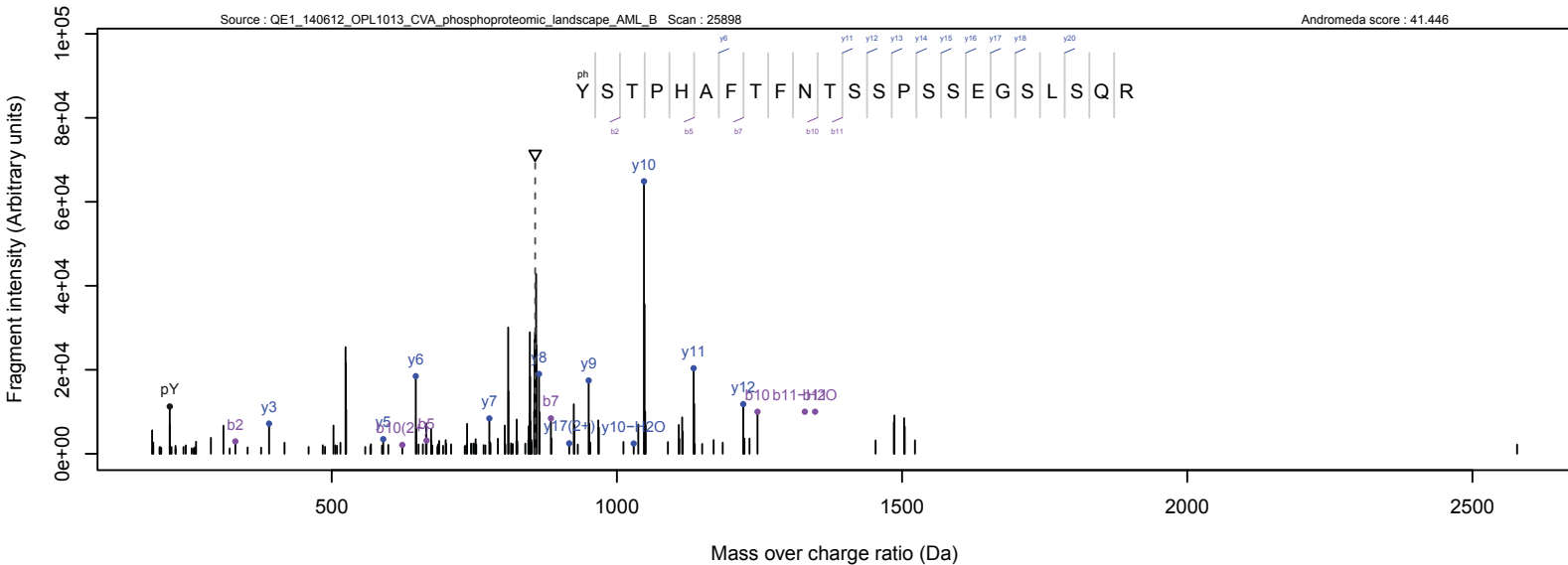
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 40525

Andromeda score : 41.513

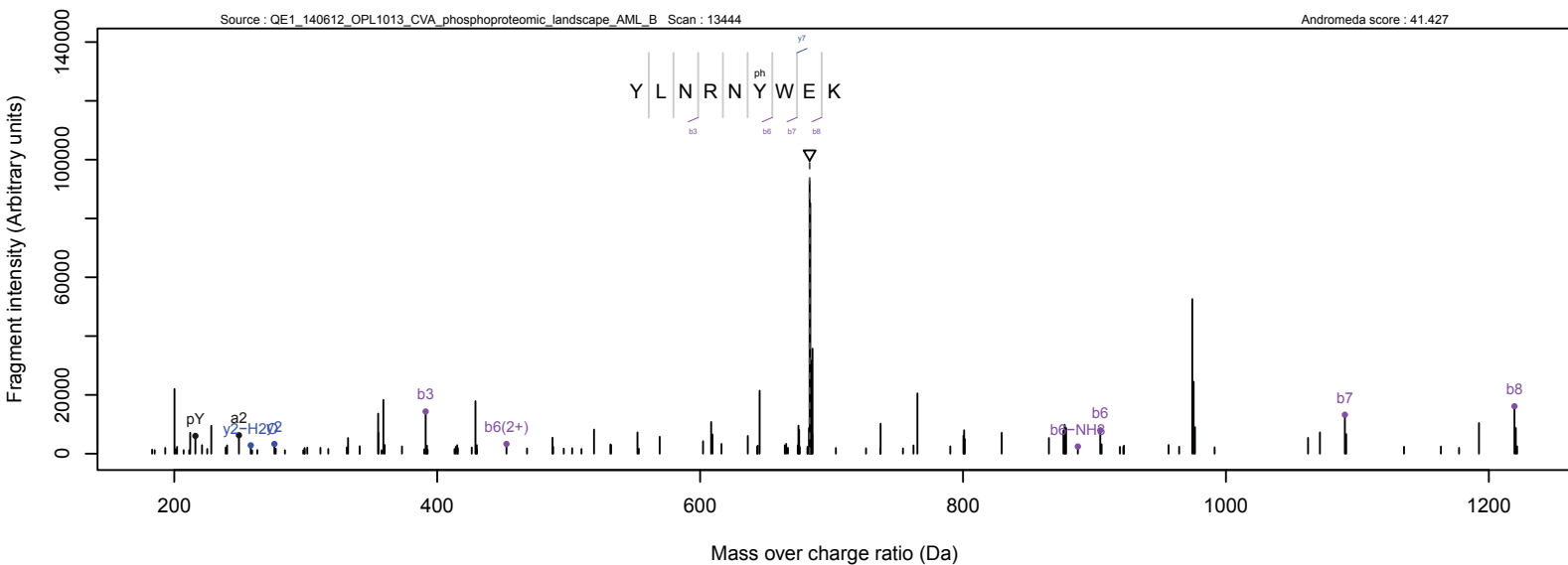




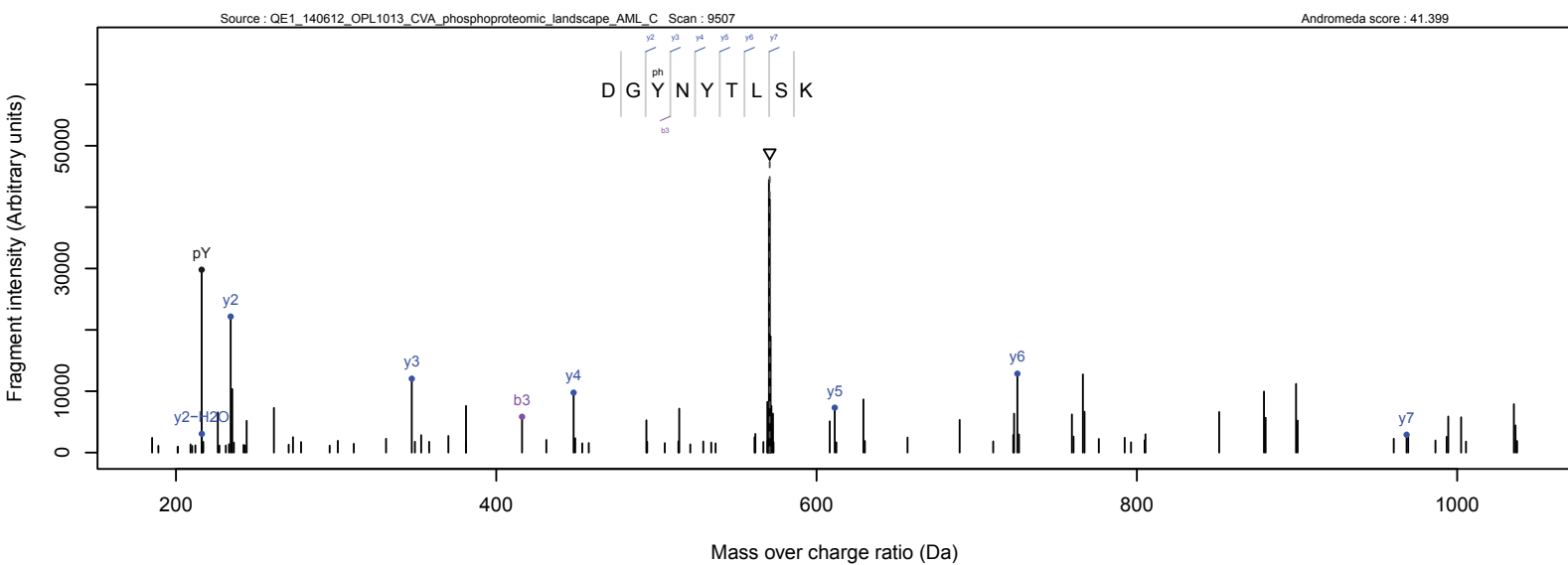
## Y(ph)STPHAFNTSSPSSEGLSQR-3+



## YLNRNY(ph)WEK-2+



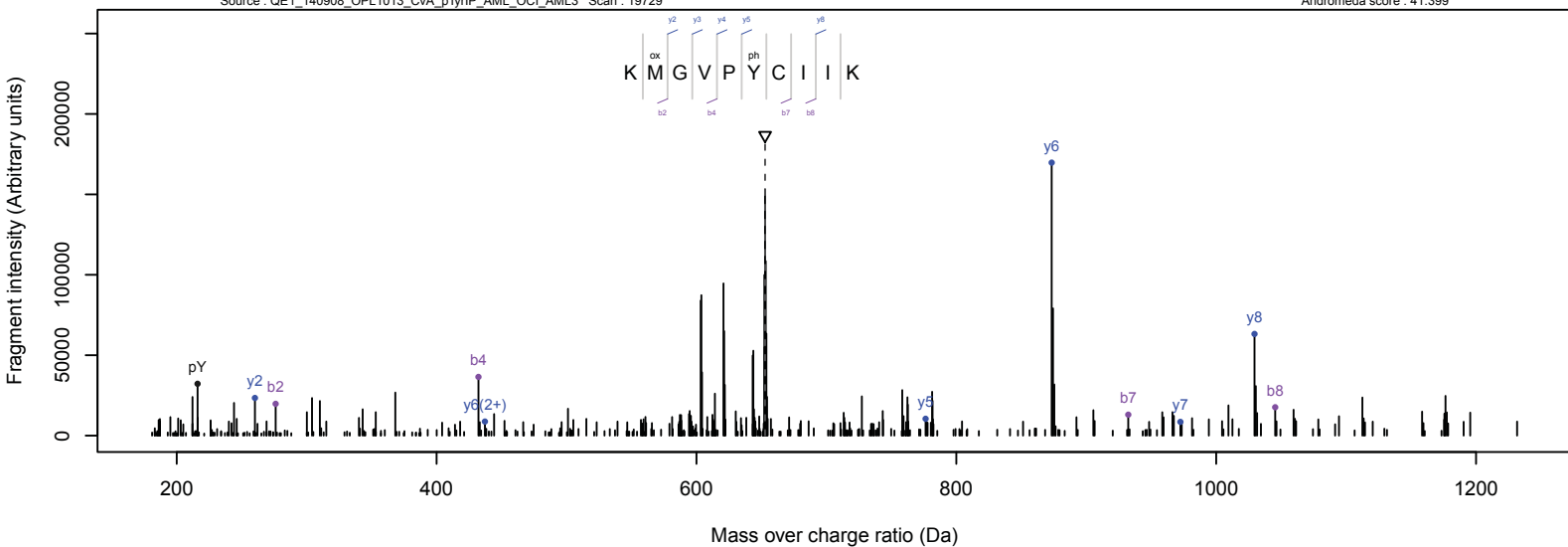
## DGY(ph)NYTLISK-2+



## KM(ox)GVPY(ph)CIK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 19729

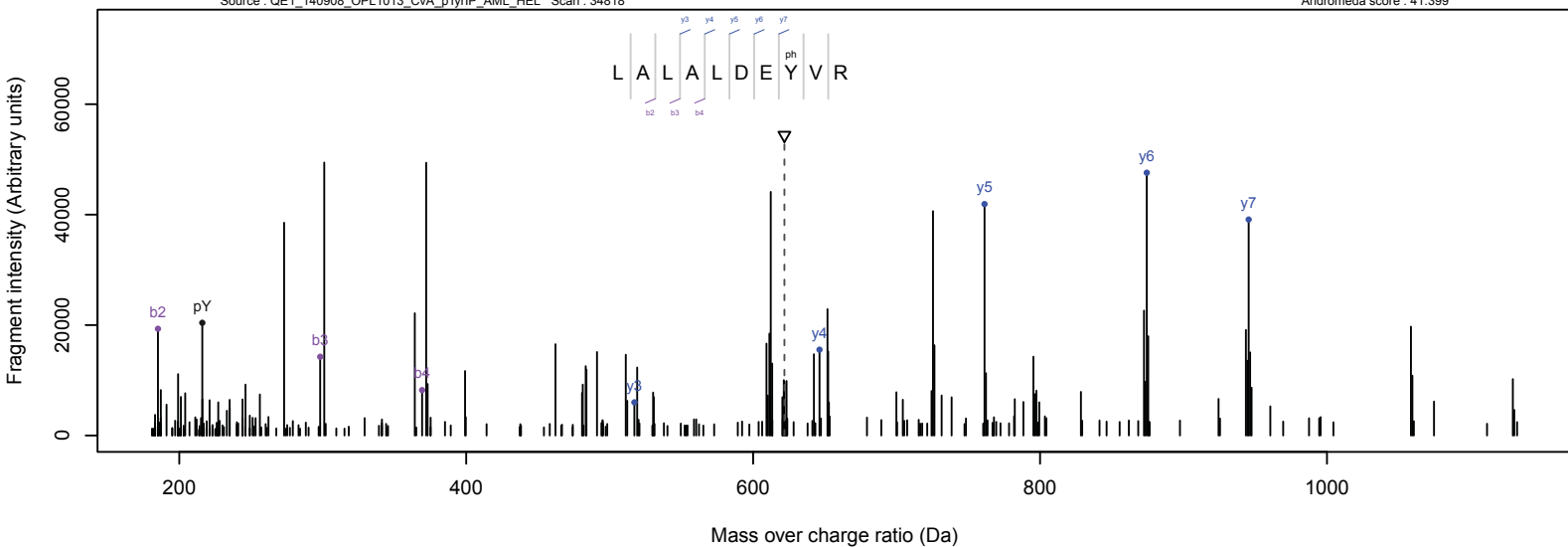
Andromeda score : 41.399



## LALALDEY(ph)VR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 34818

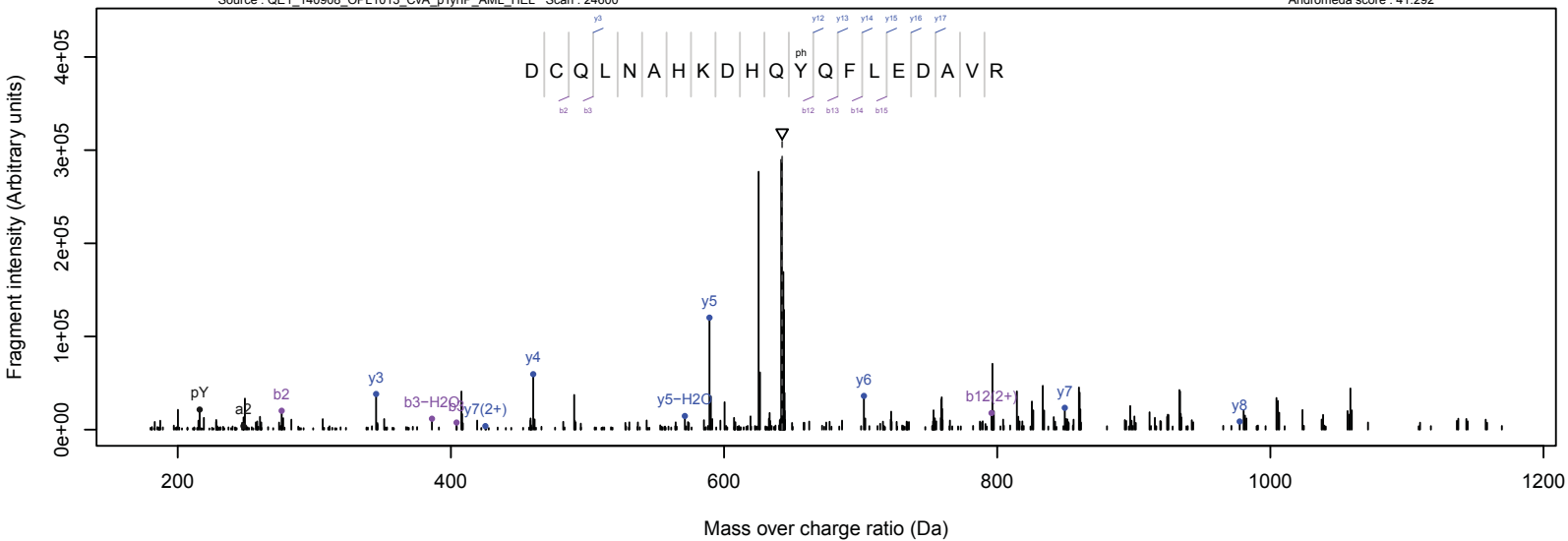
Andromeda score : 41.399



## DCQLNAHKDHQ(ph)QFLEDAVR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 24600

Andromeda score : 41.292

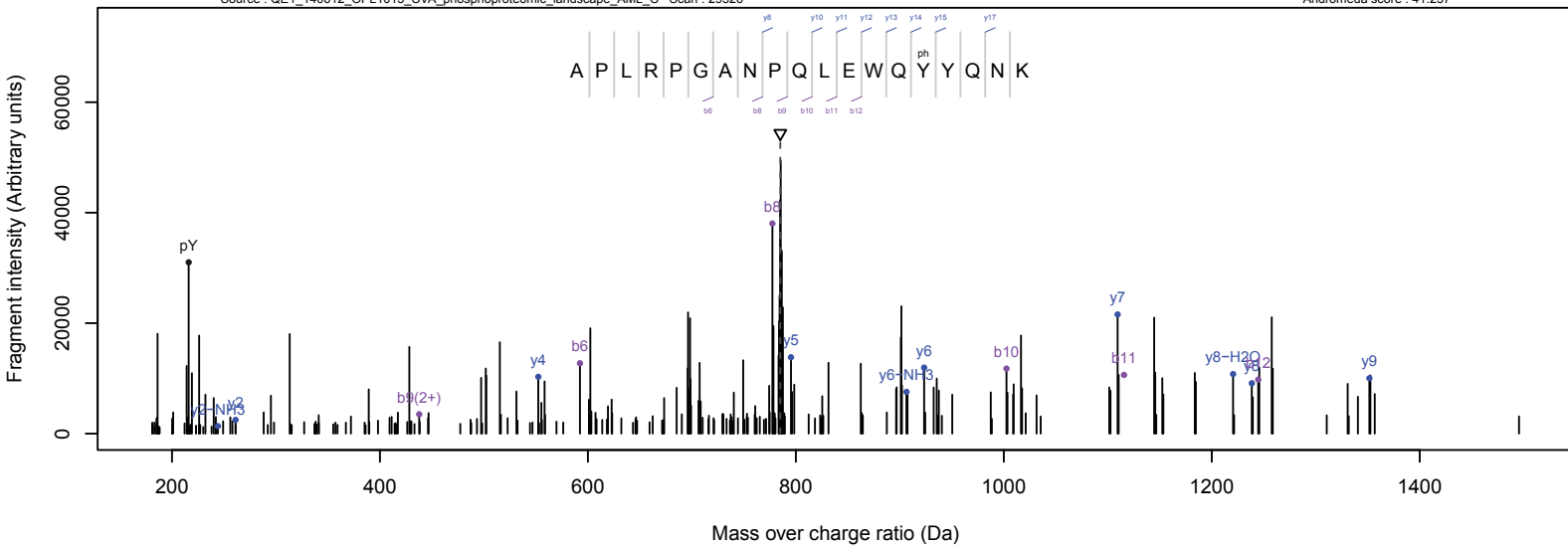




APLRPGANPQLEWQY(ph)YQNK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 29526

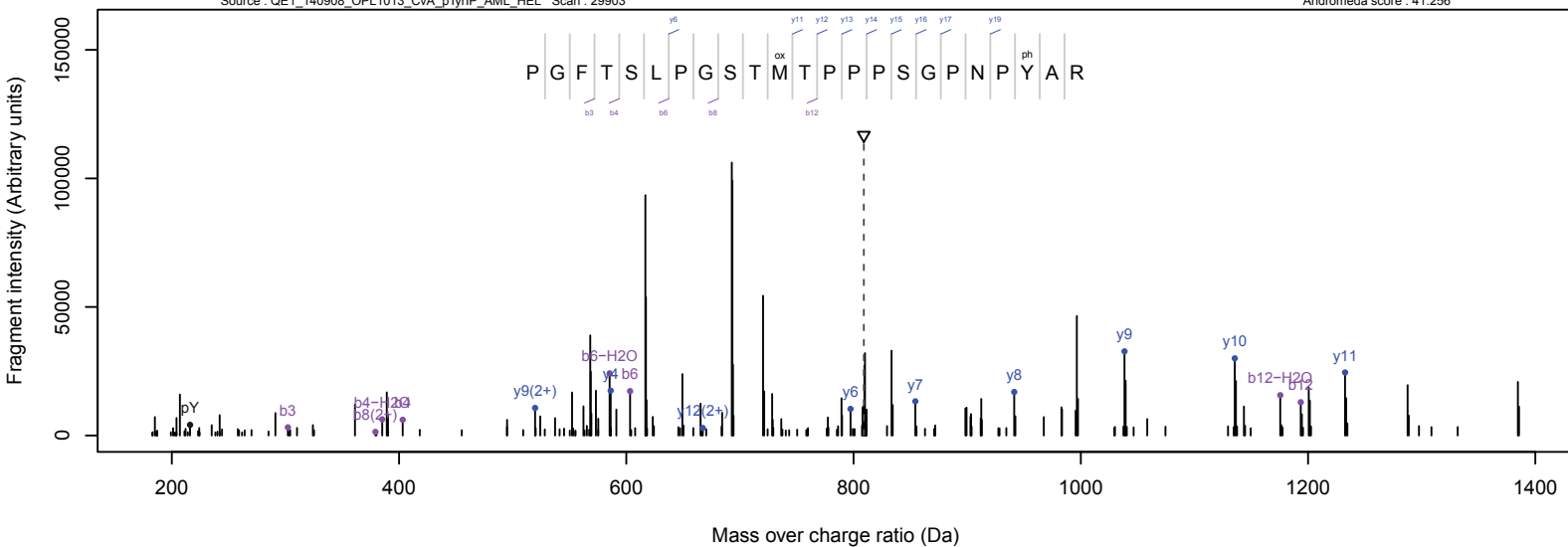
Andromeda score : 41.257



PGFTSLPGSTM(ox)TPPPSGPNP(ph)AR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 29903

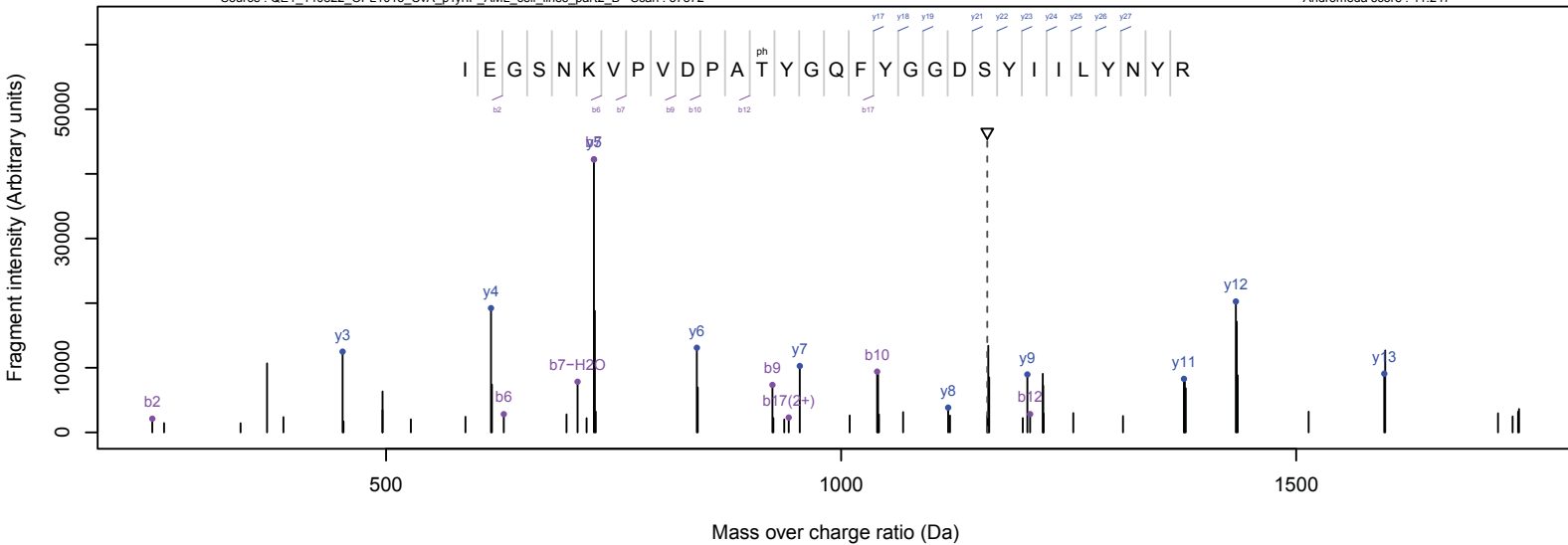
Andromeda score : 41.256



IEGSNKVPVDPAT(ph)YGQFYGGDSYIILYNYR-3+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 37872

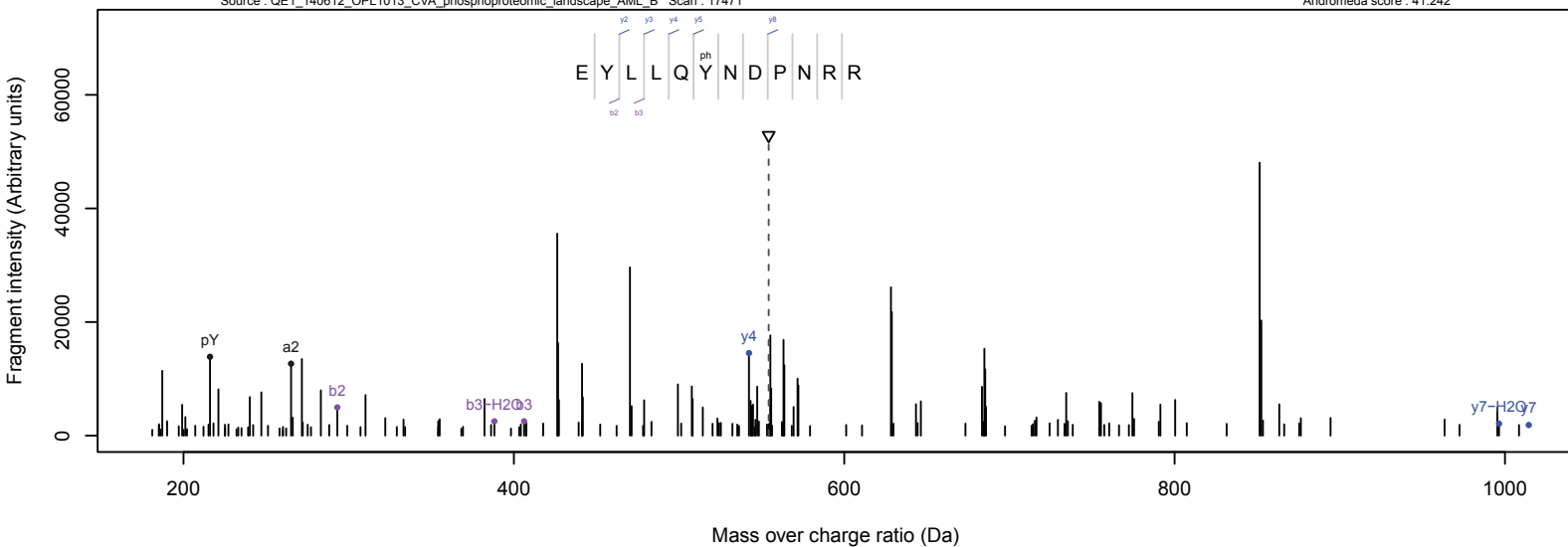
Andromeda score : 41.247



## EYLLQY(ph)NDPNRR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 17471

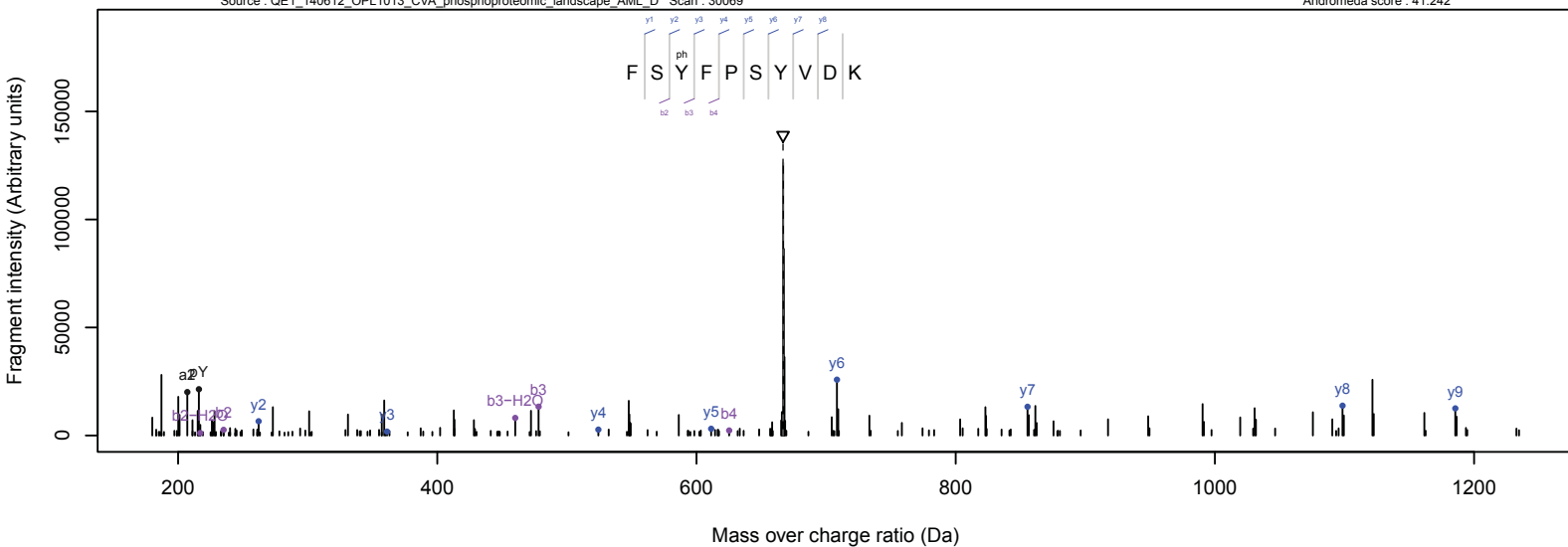
Andromeda score : 41.242



## FSY(ph)FPSYVDK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 30069

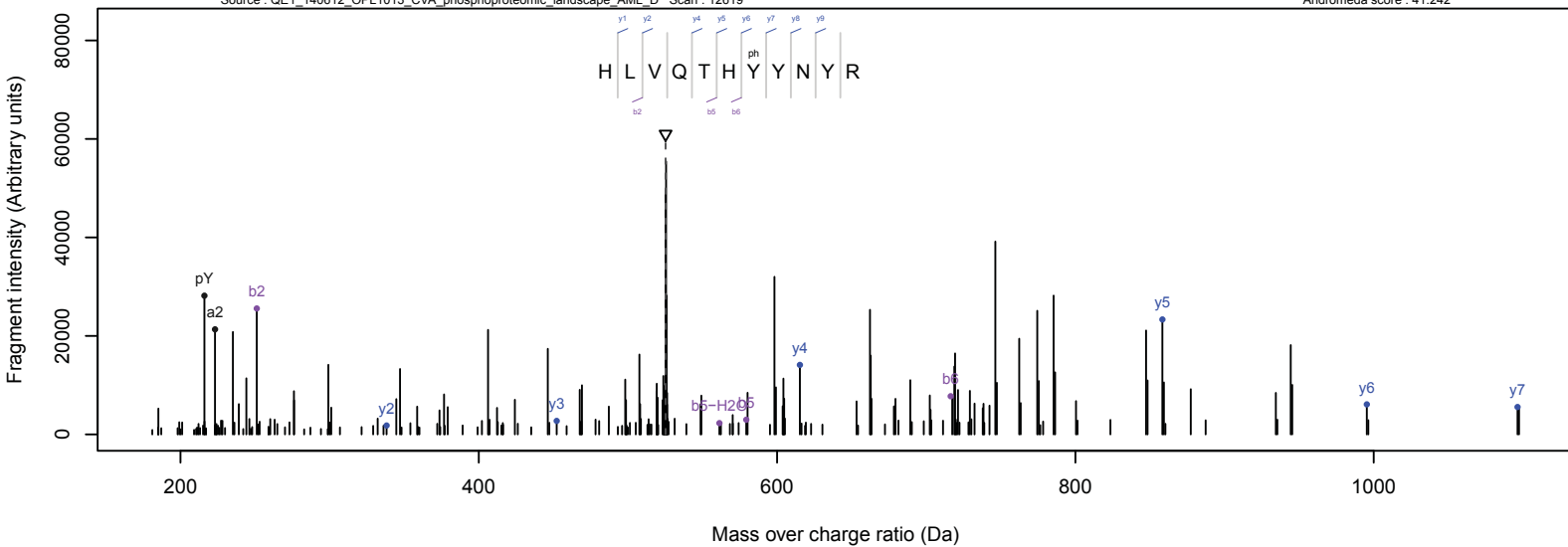
Andromeda score : 41.242



## HLVQTHY(ph)YNYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 12619

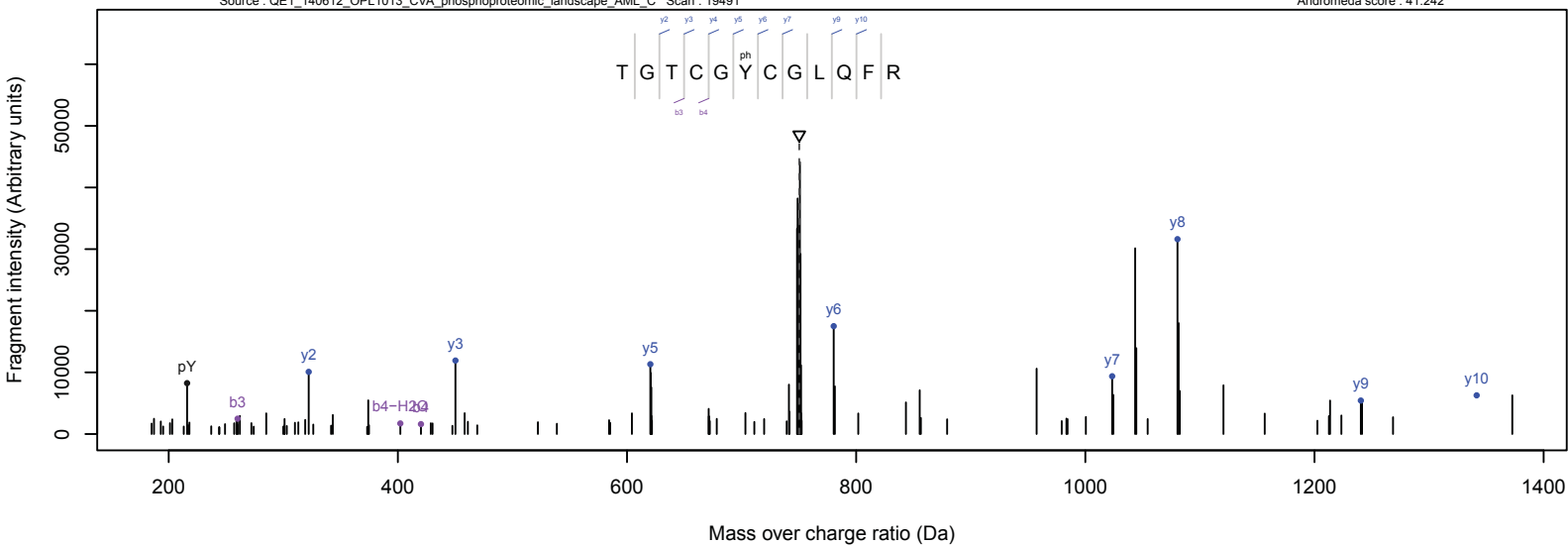
Andromeda score : 41.242



## TGTCGY(ph)CGLQFR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 19491

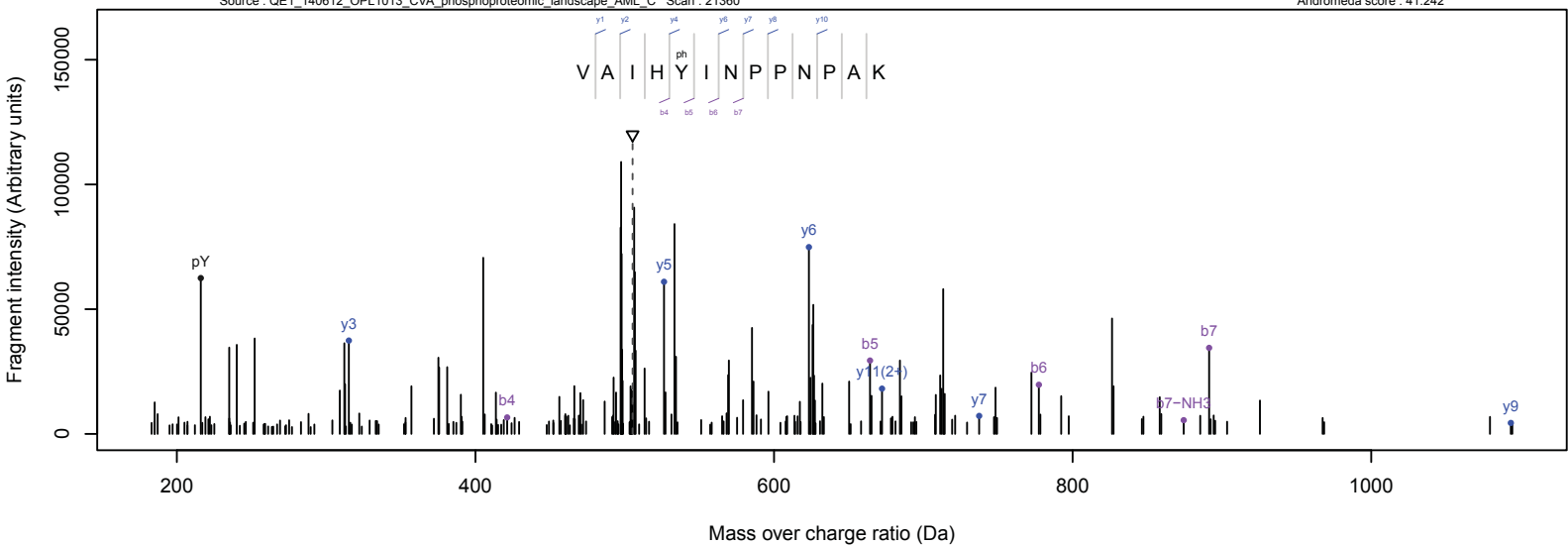
Andromeda score : 41.242



## VAIHYP(h)INPPPAK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 21360

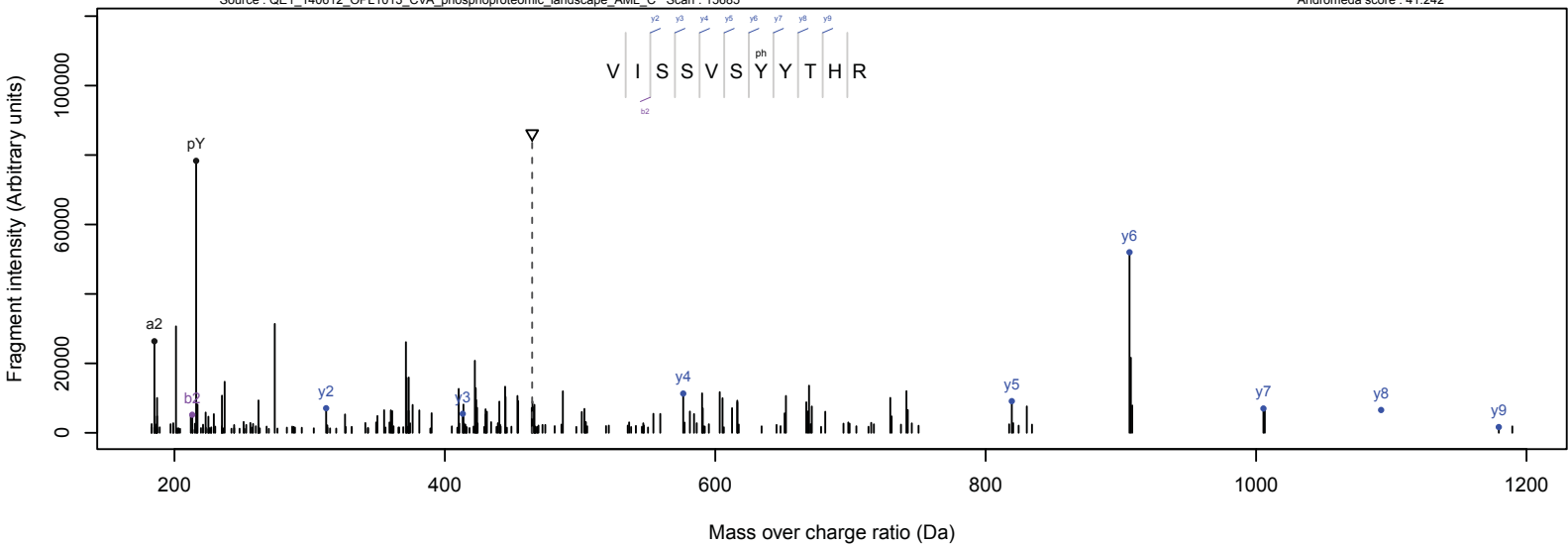
Andromeda score : 41.242



## VISSVY(ph)YTHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15685

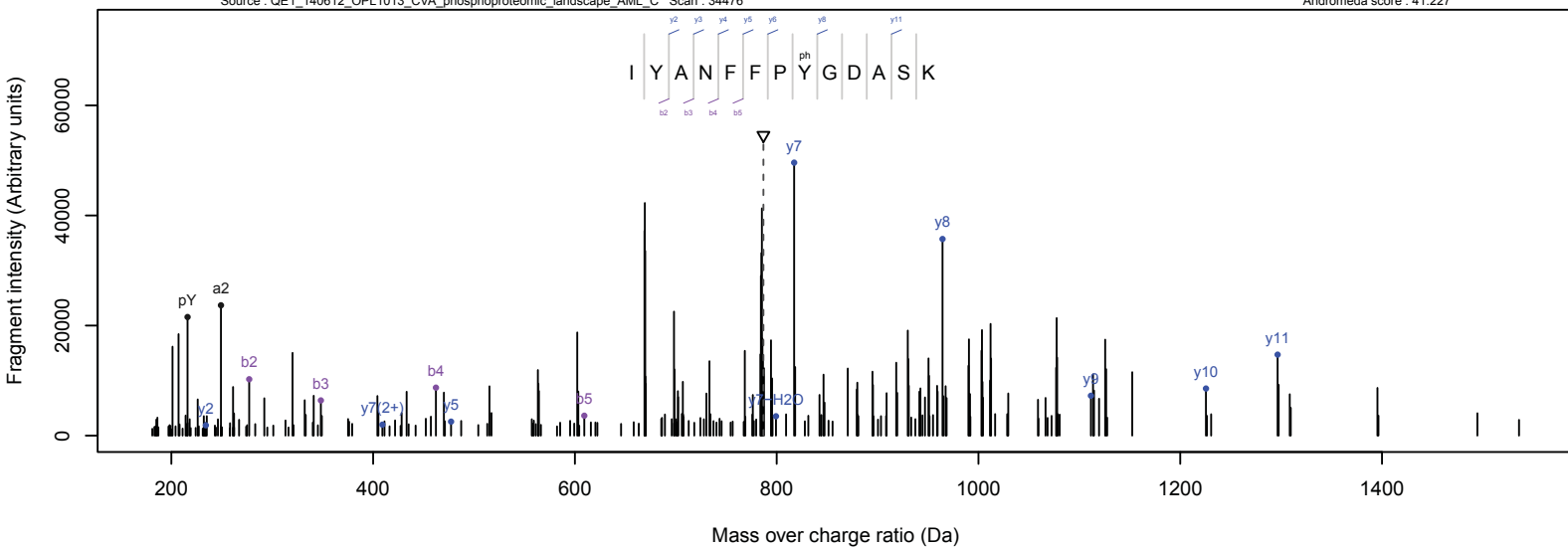
Andromeda score : 41.242



## IYANFFYPYGDASK-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 34476

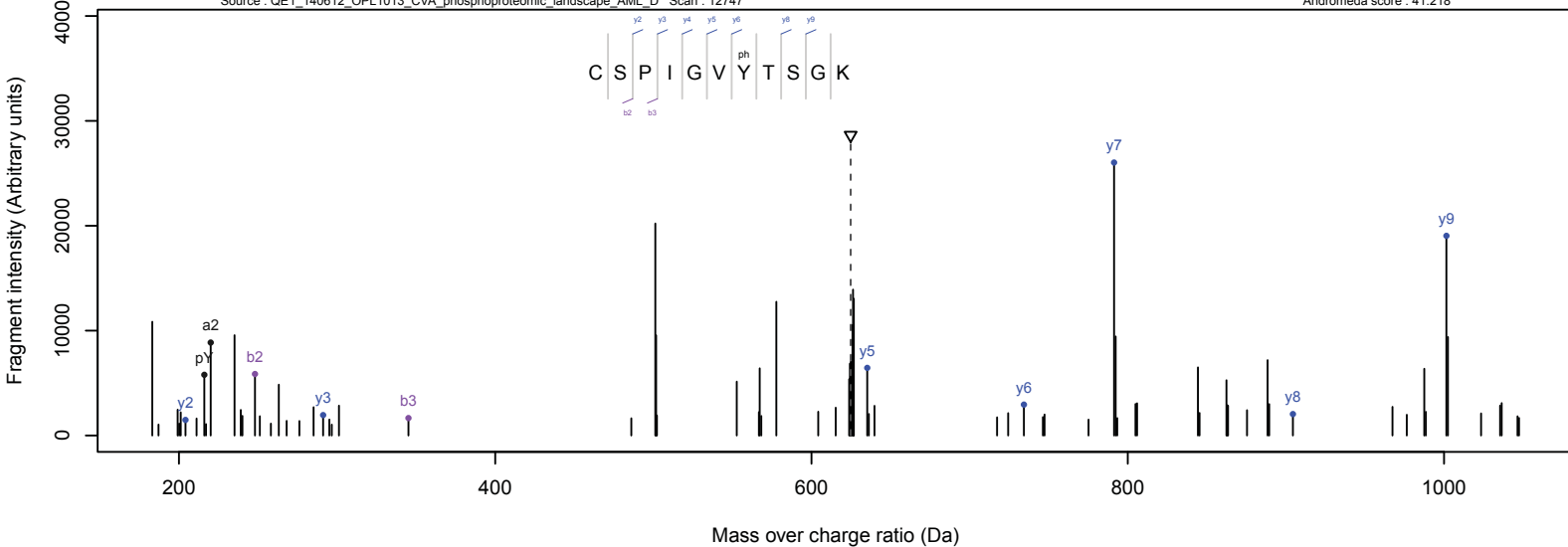
Andromeda score: 41.227



## CSPIGVY(TSGK)-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan: 12747

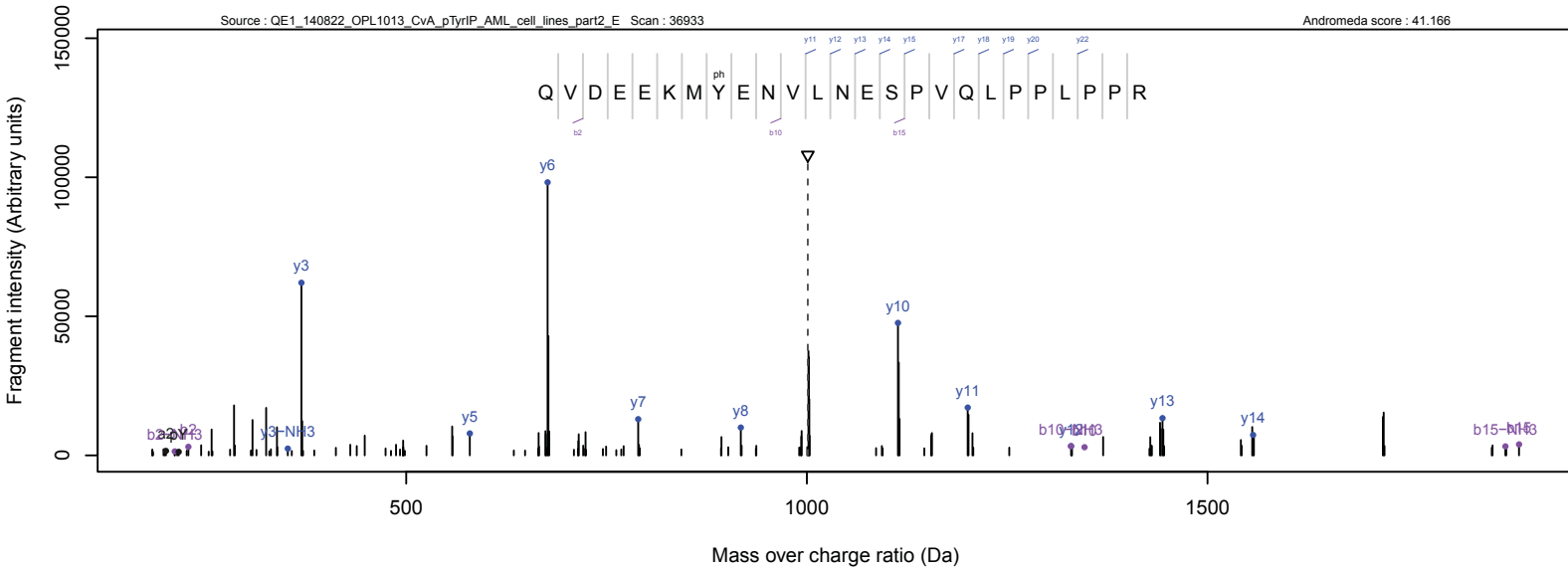
Andromeda score: 41.218



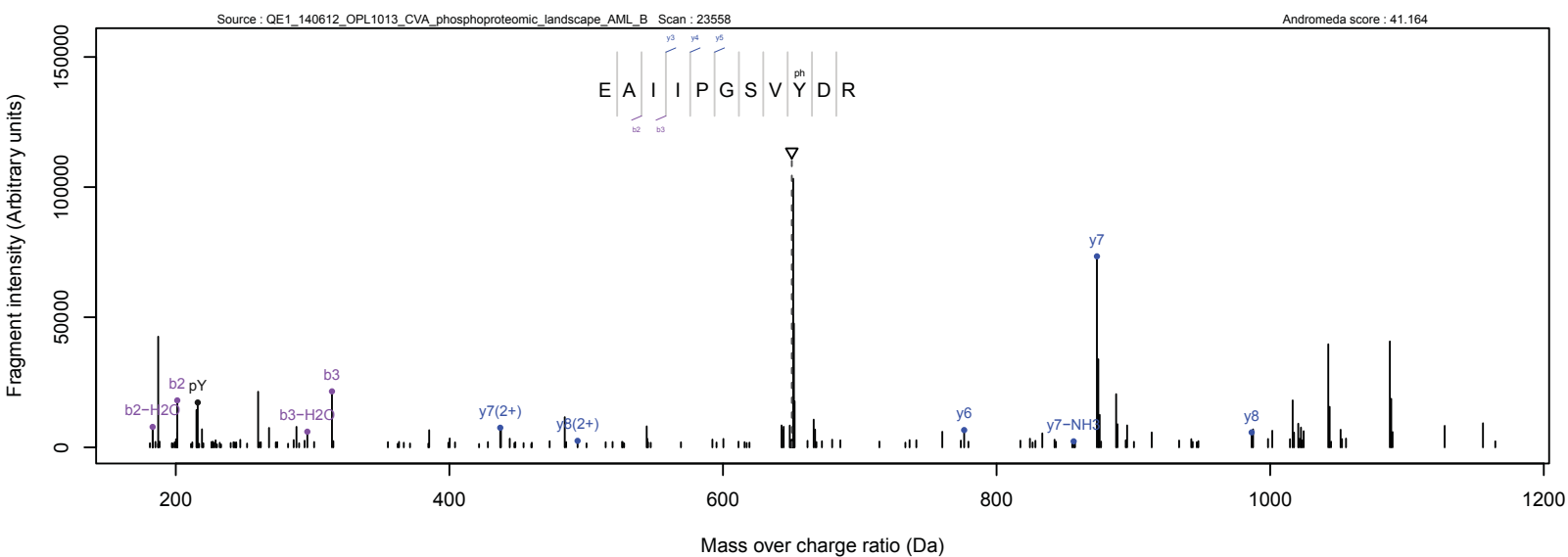
## QVDEEKMY(ph)ENVLNEPVLPLPPR-3+

Source: QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_E Scan: 36933

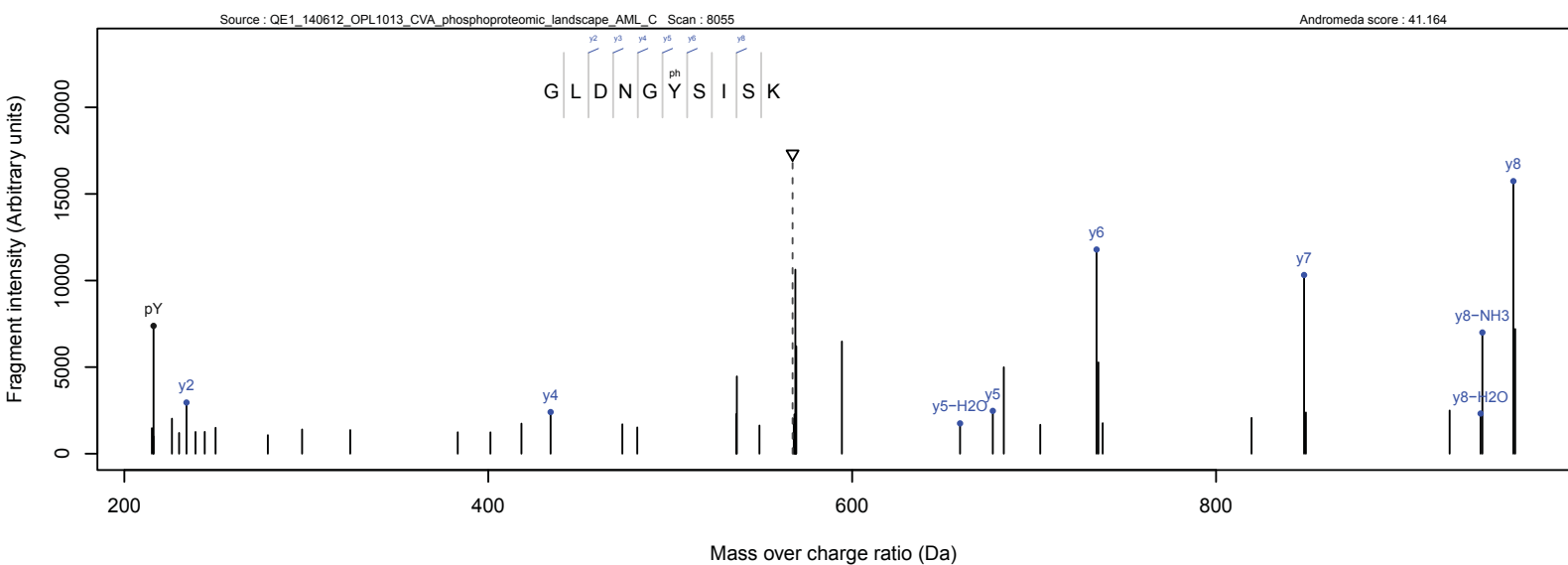
Andromeda score: 41.166



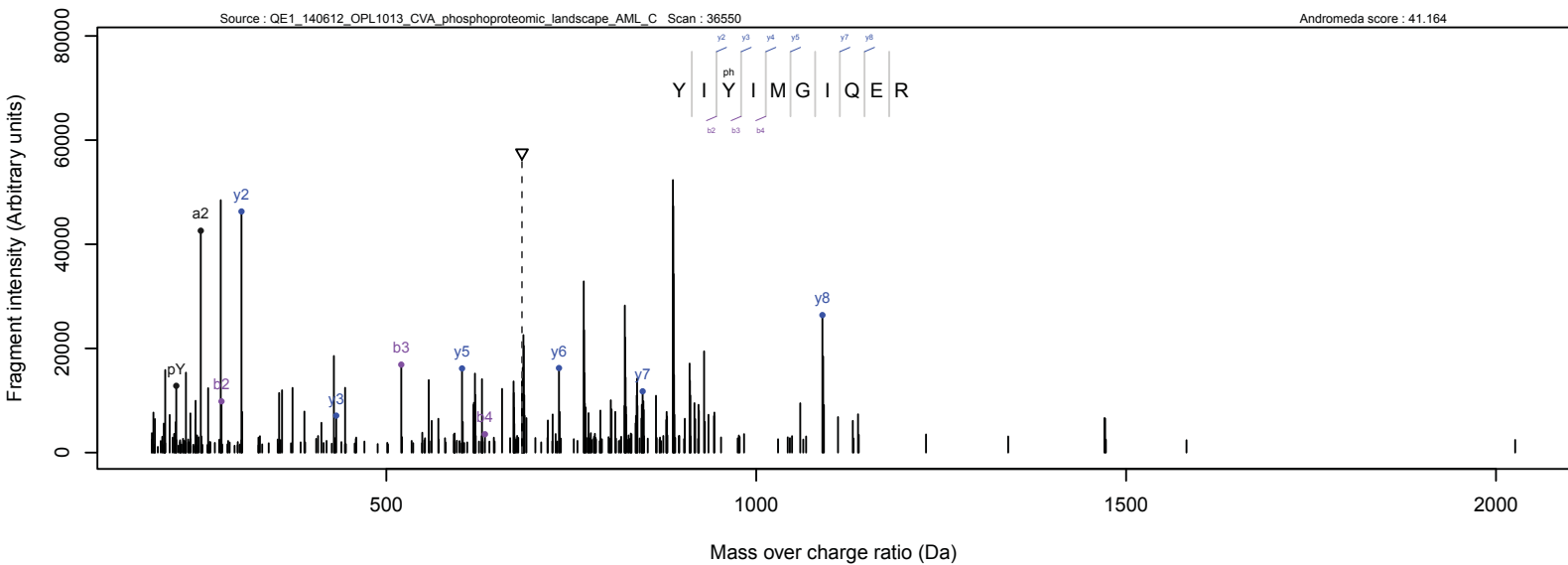
## EAIIPGSVY(ph)DR-2+



## GLDNGY(ph)SISK-2+



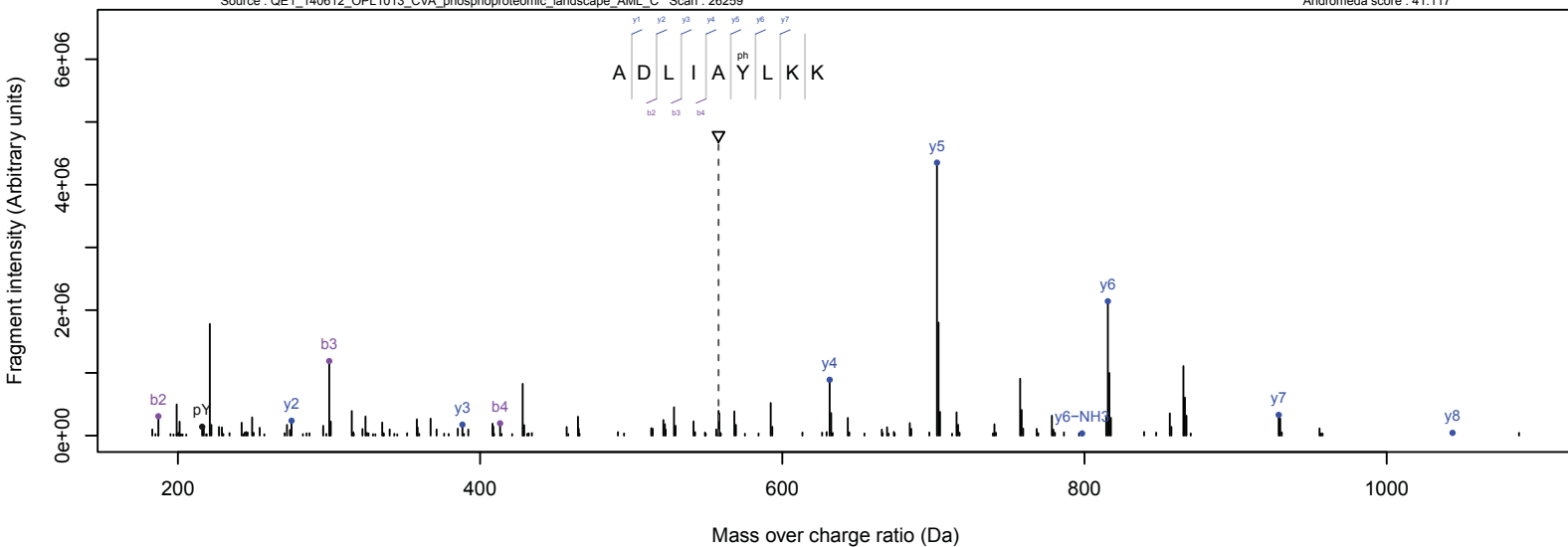
## YIYIMGIQER-2+



## ADLIAY(ph)LKK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 26259

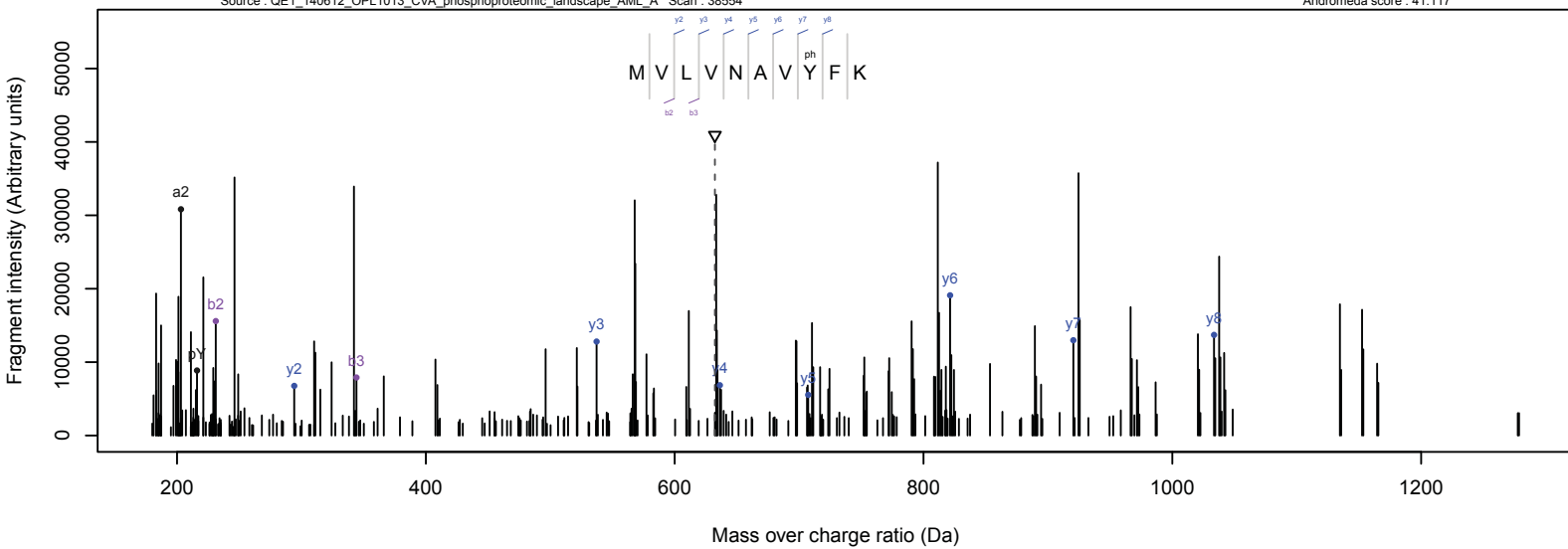
Andromeda score : 41.117



## MVLNAVY(ph)FK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 38554

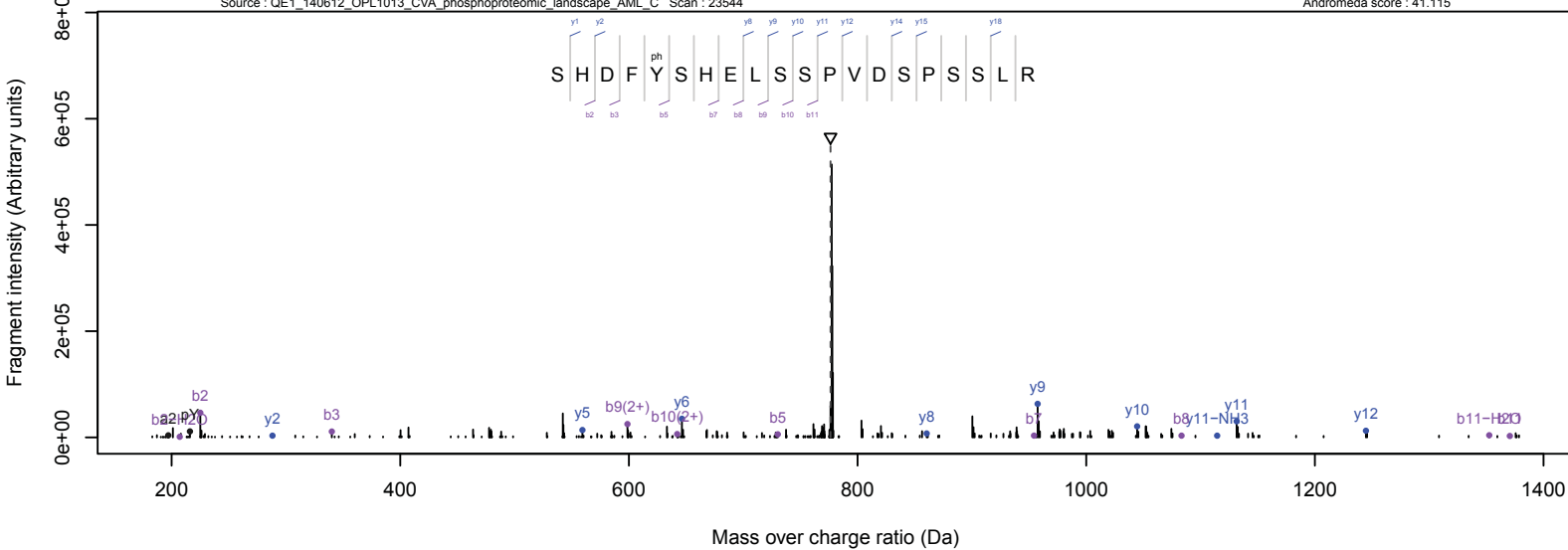
Andromeda score : 41.117



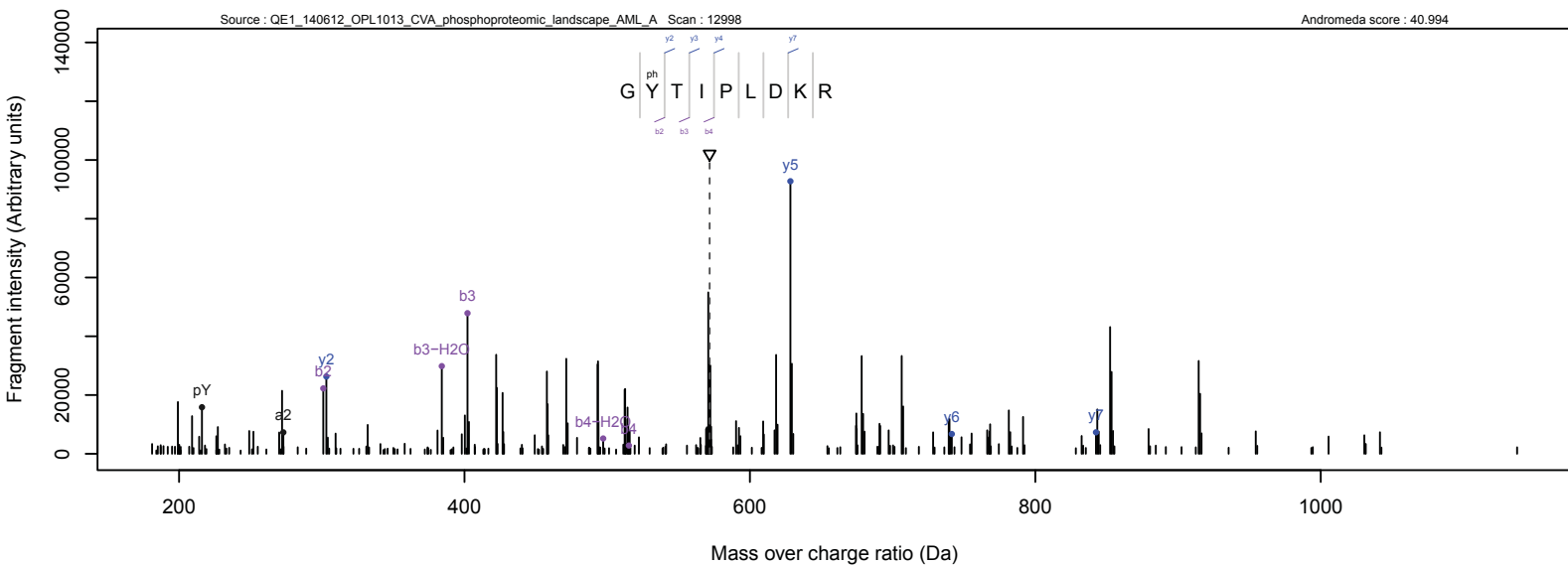
## SHDFY(ph)SHELSSPVDSPLSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23544

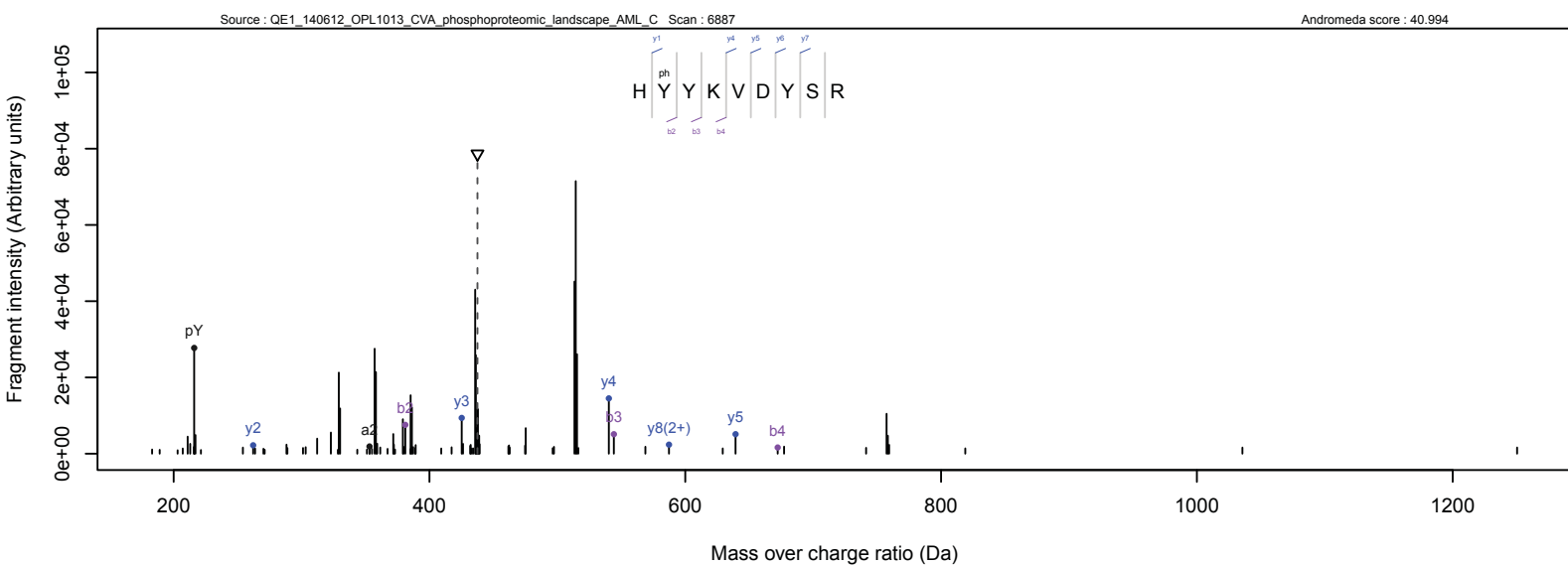
Andromeda score : 41.115



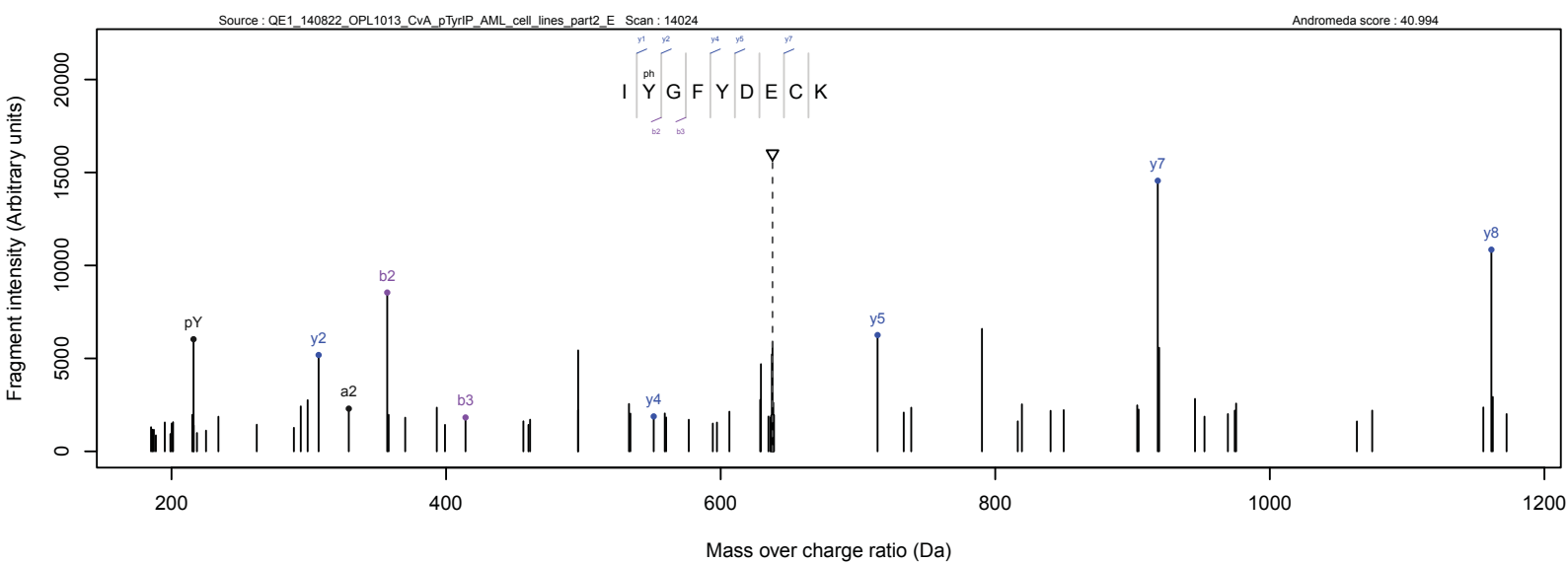
## GY(ph)TIPLDKR-2+



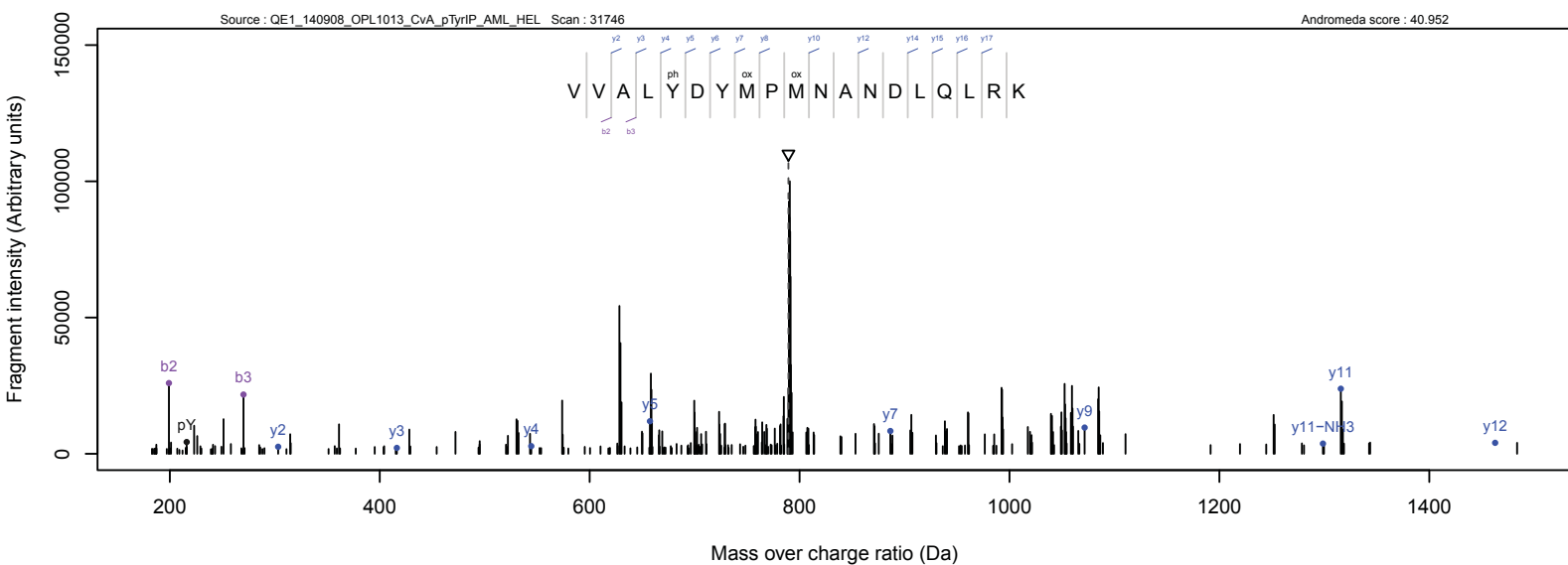
## HY(ph)YKVDYSR-3+



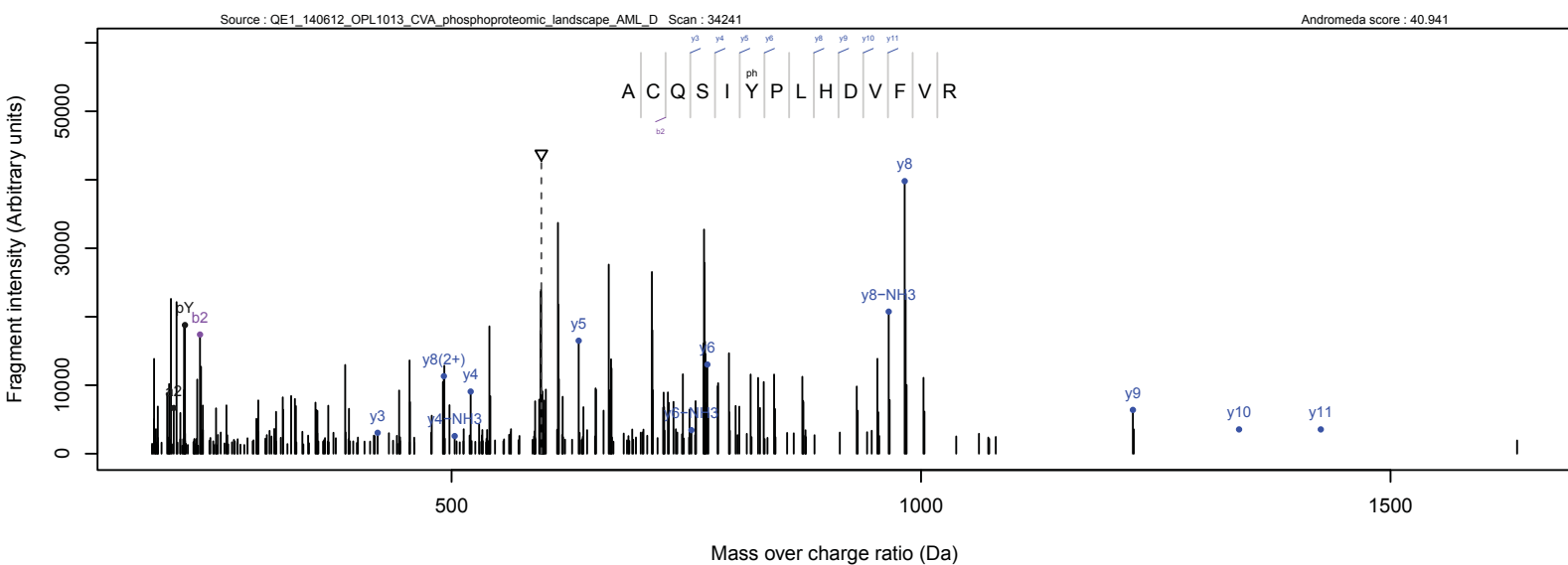
## IY(ph)GFYDECK-2+



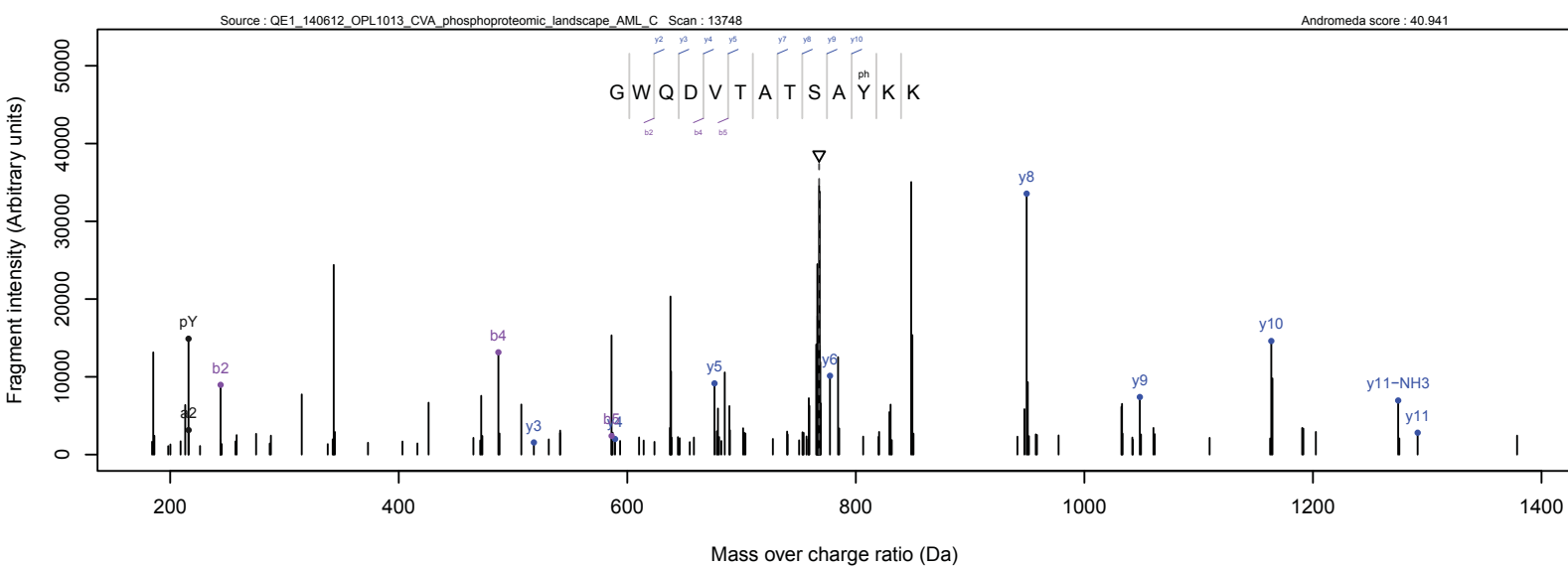
## VVALY(ph)DYM(ox)PM(ox)NANDLQLRK-3+



## ACQSIY(ph)PLHDVFVR-3+

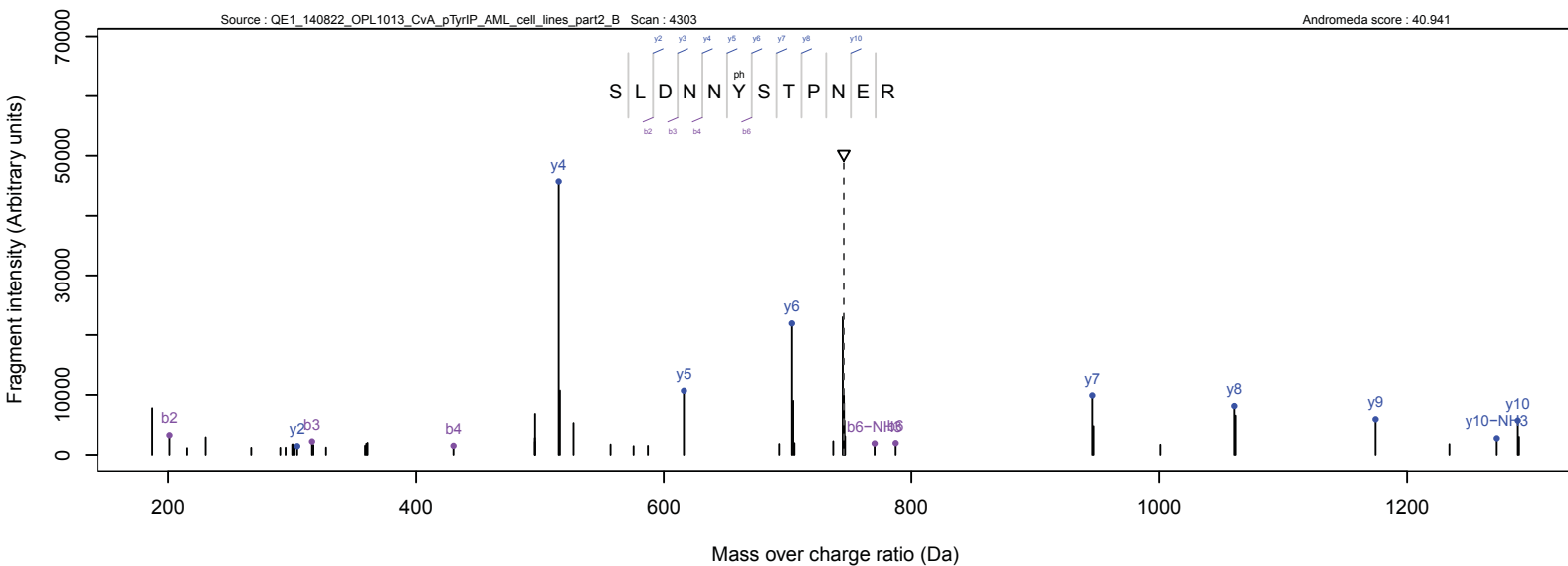


## GWQDVTATSAY(ph)KK-2+

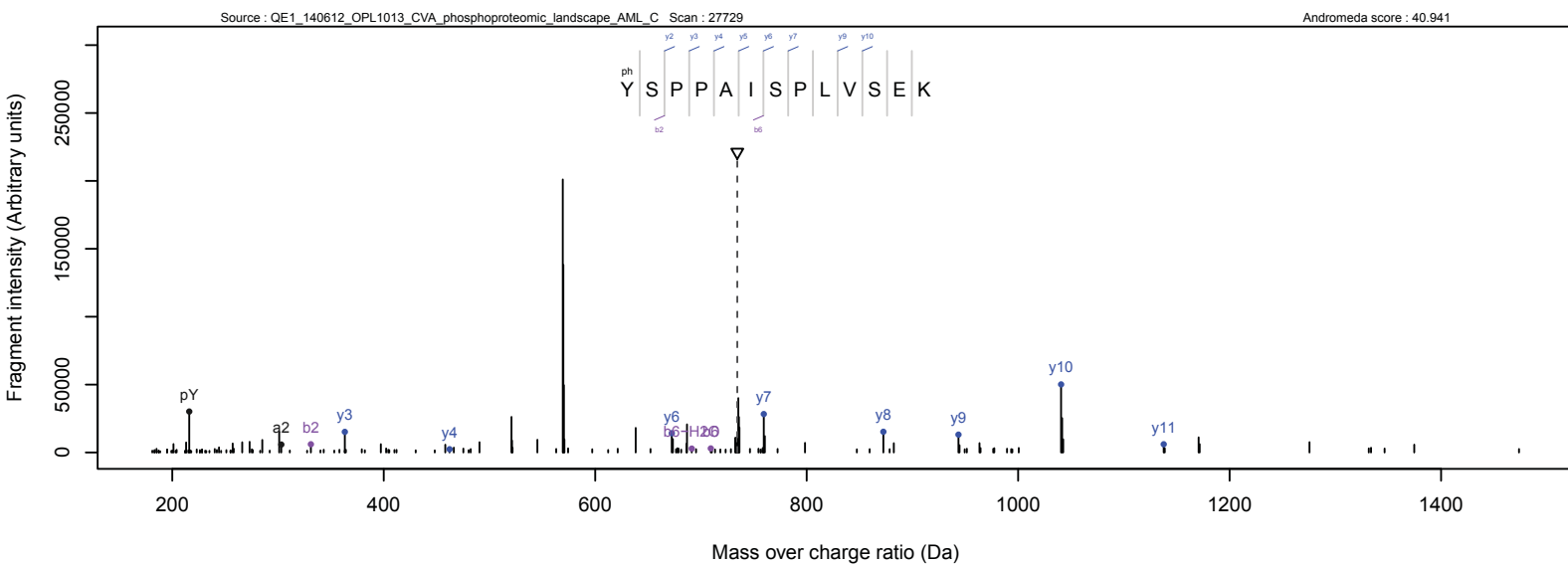




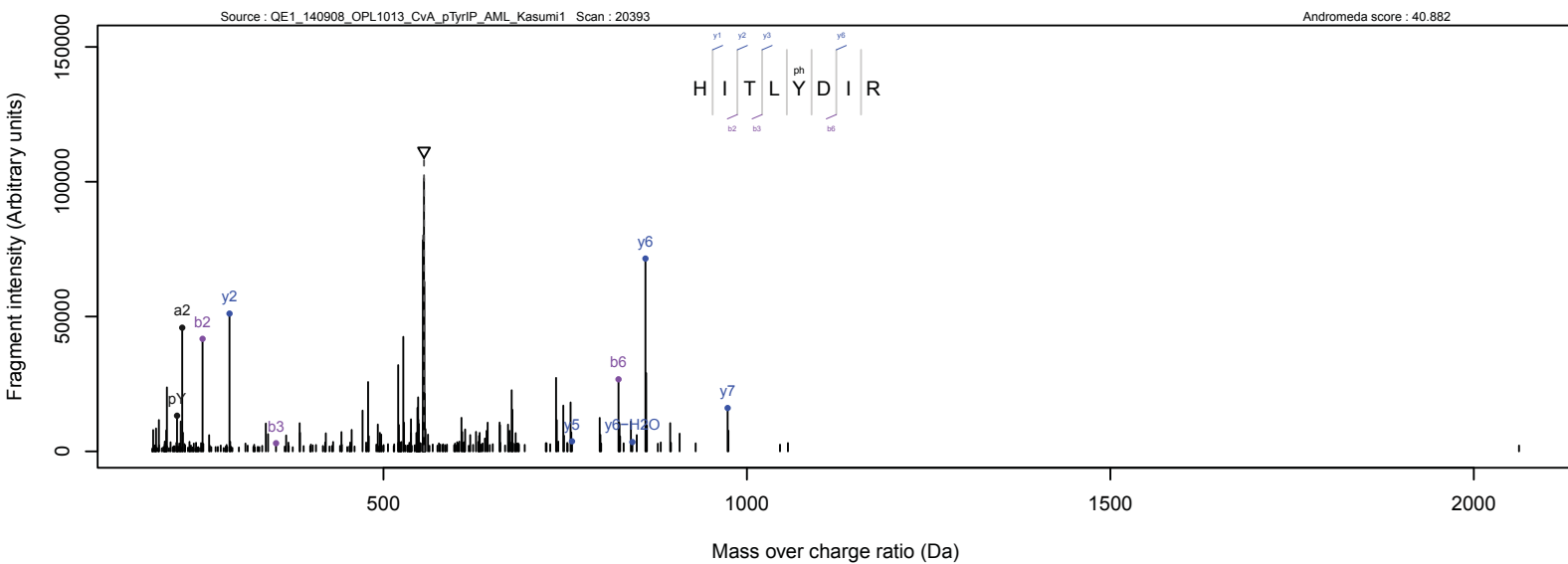
SLDNNY(ph)STPNER-2+



Y(ph)SPPAISPLVSEK-2+



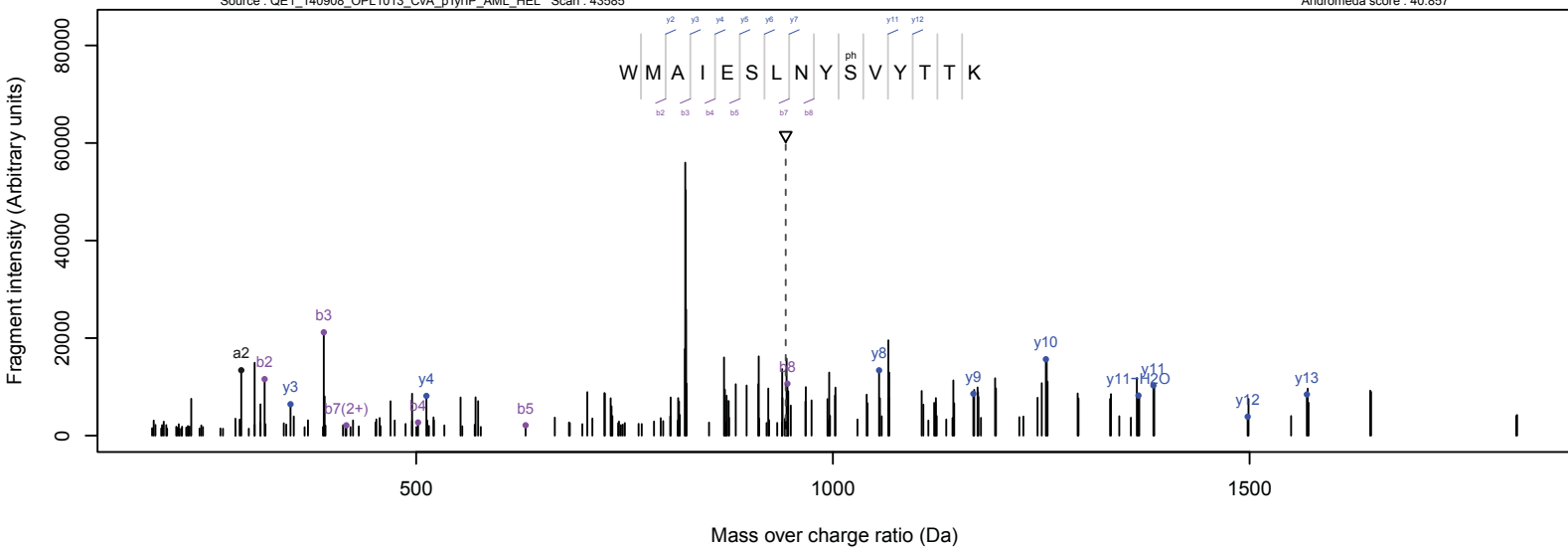
HITLY(ph)DIR-2+



## WMAIESLNYS(ph)VYTTK-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HEL Scan : 43585

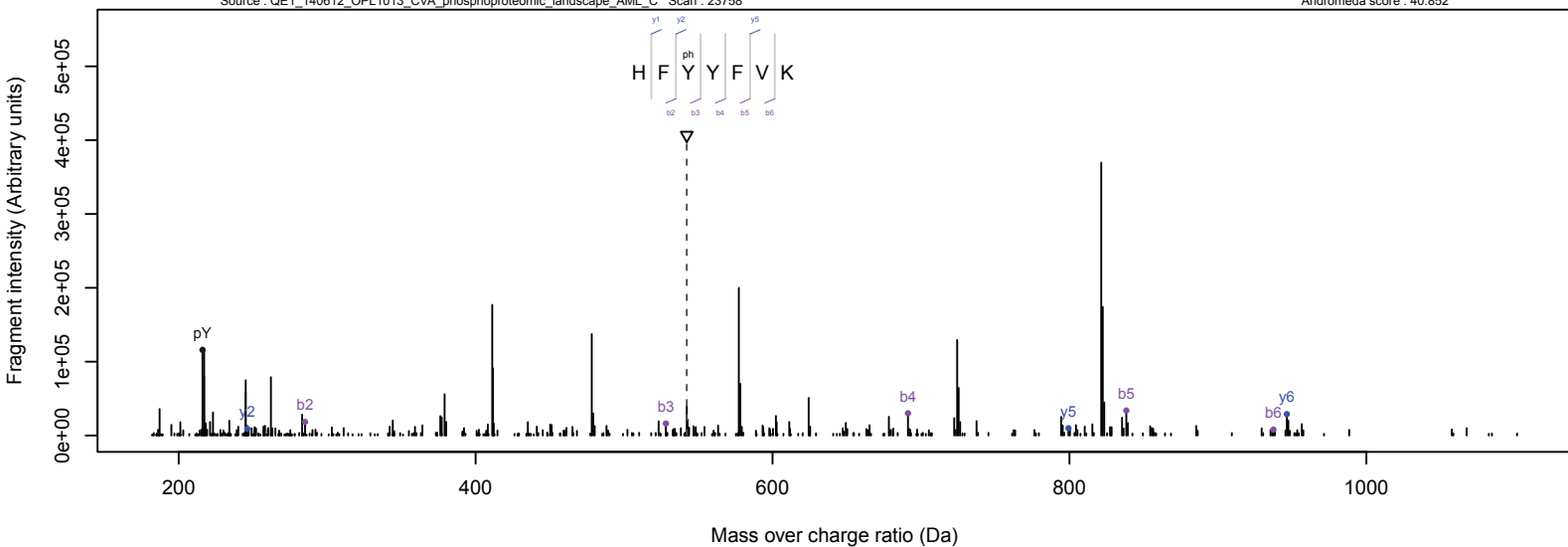
Andromeda score : 40.857



## HFY(ph)YFVK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23758

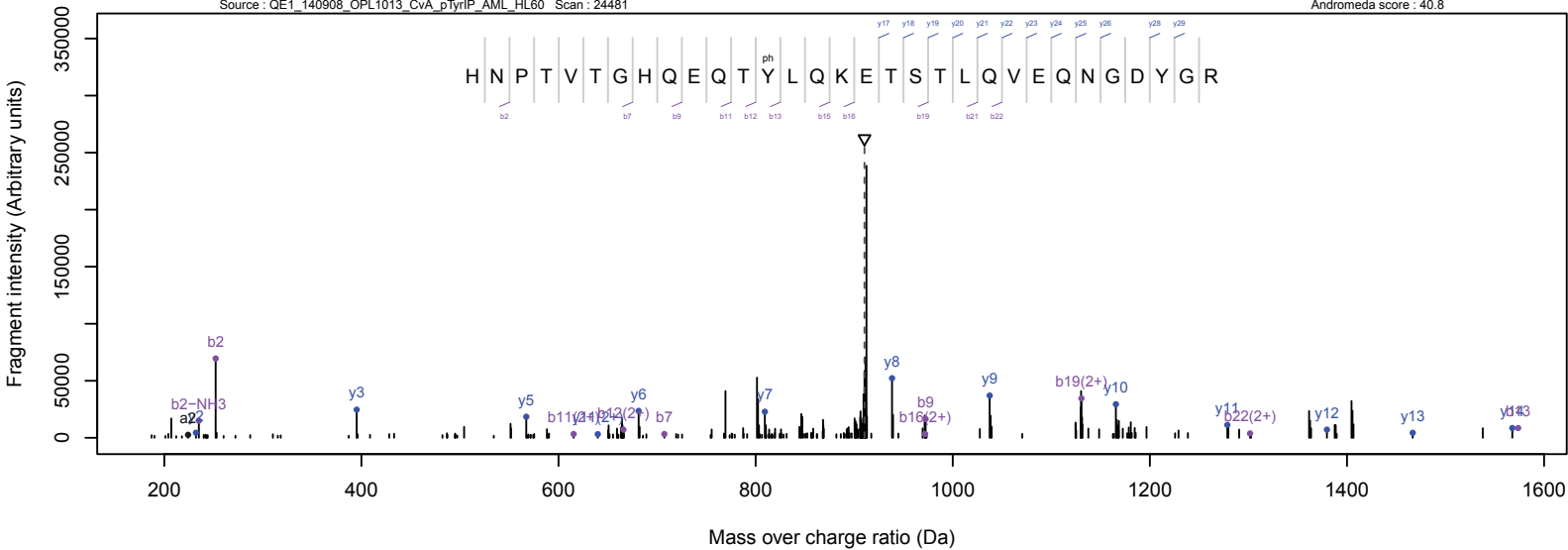
Andromeda score : 40.852



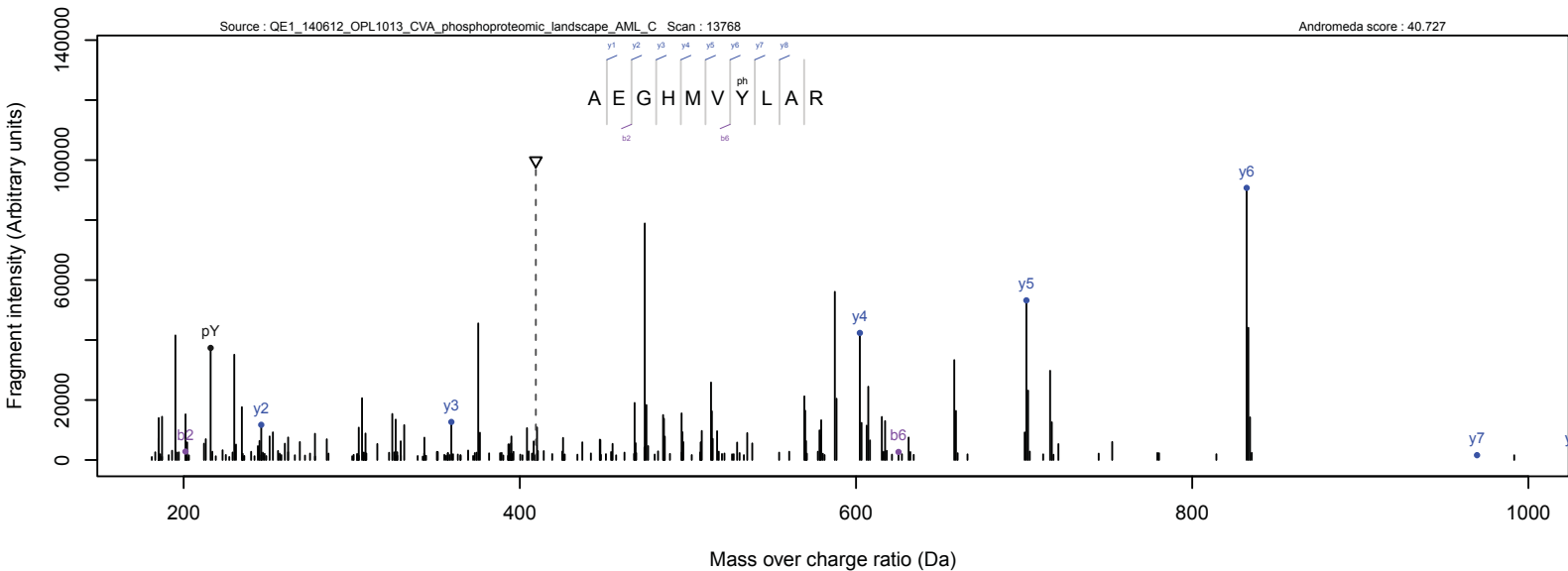
## HNPTVTGHQEQT(ph)LQKETSTLQVEQNGDYGR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_HL60 Scan : 24481

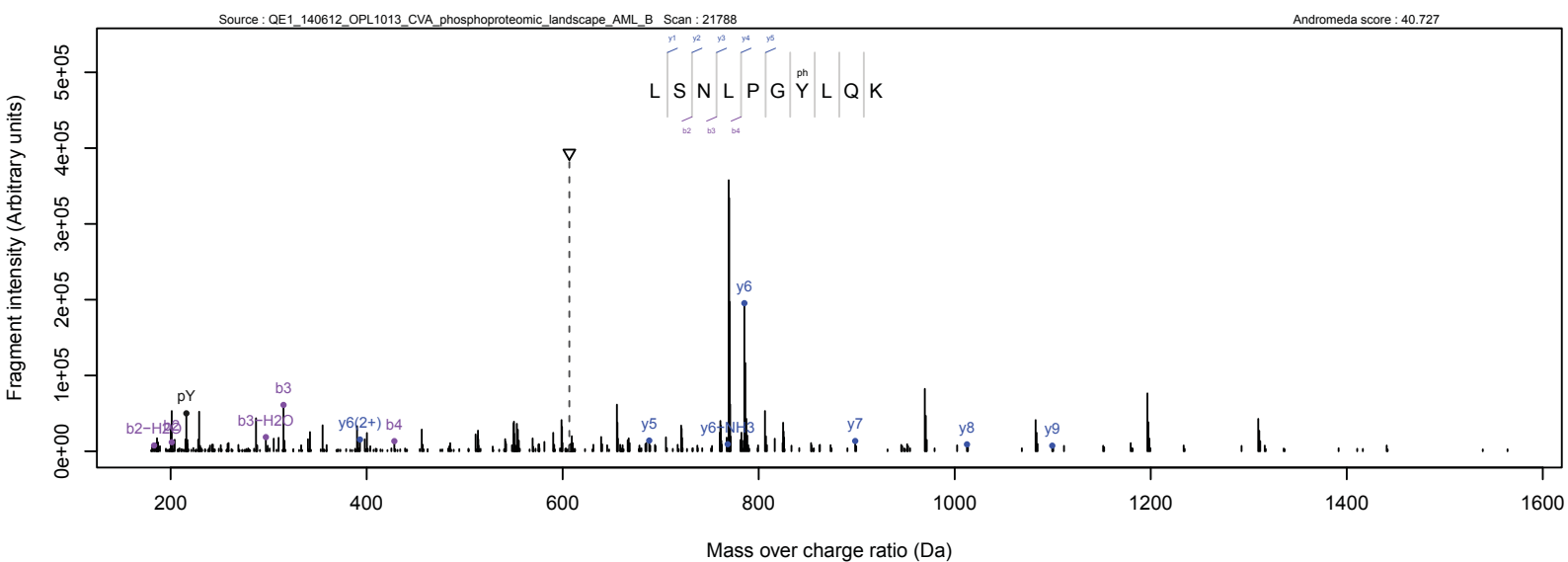
Andromeda score : 40.8



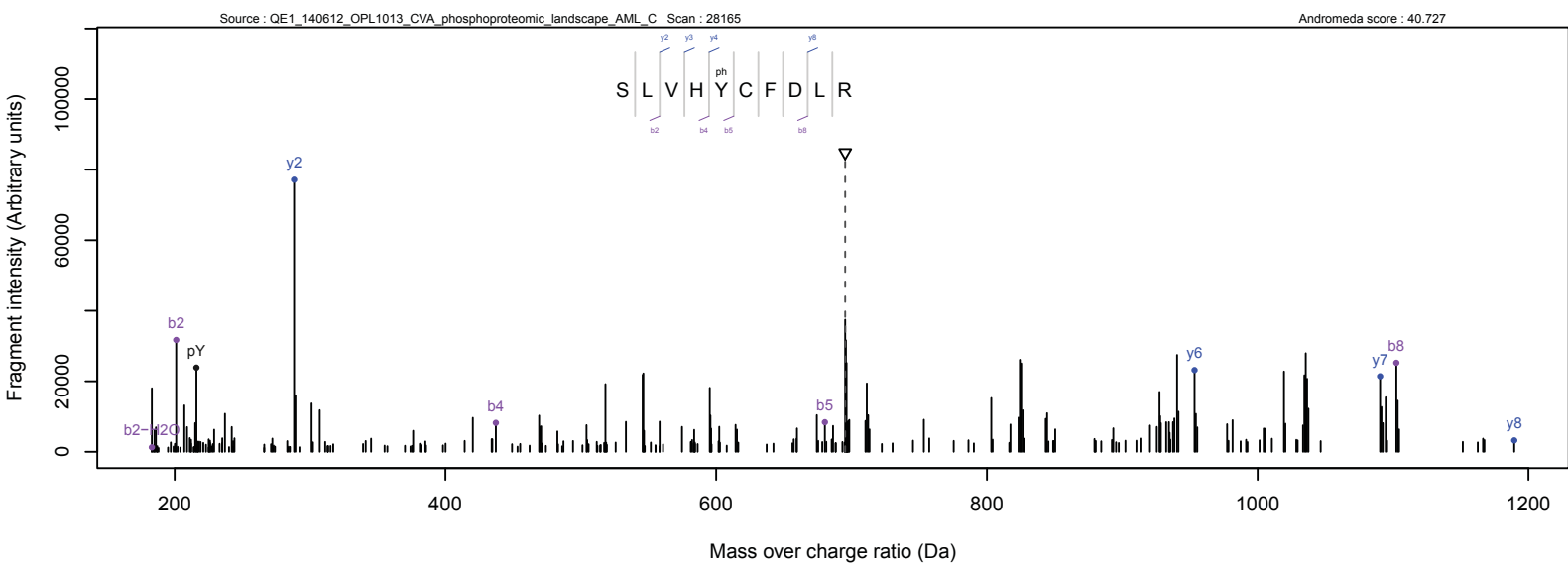
**AEGHVMY(ph)LAR-3+**



**LSNLPGY(ph)LQK-2+**



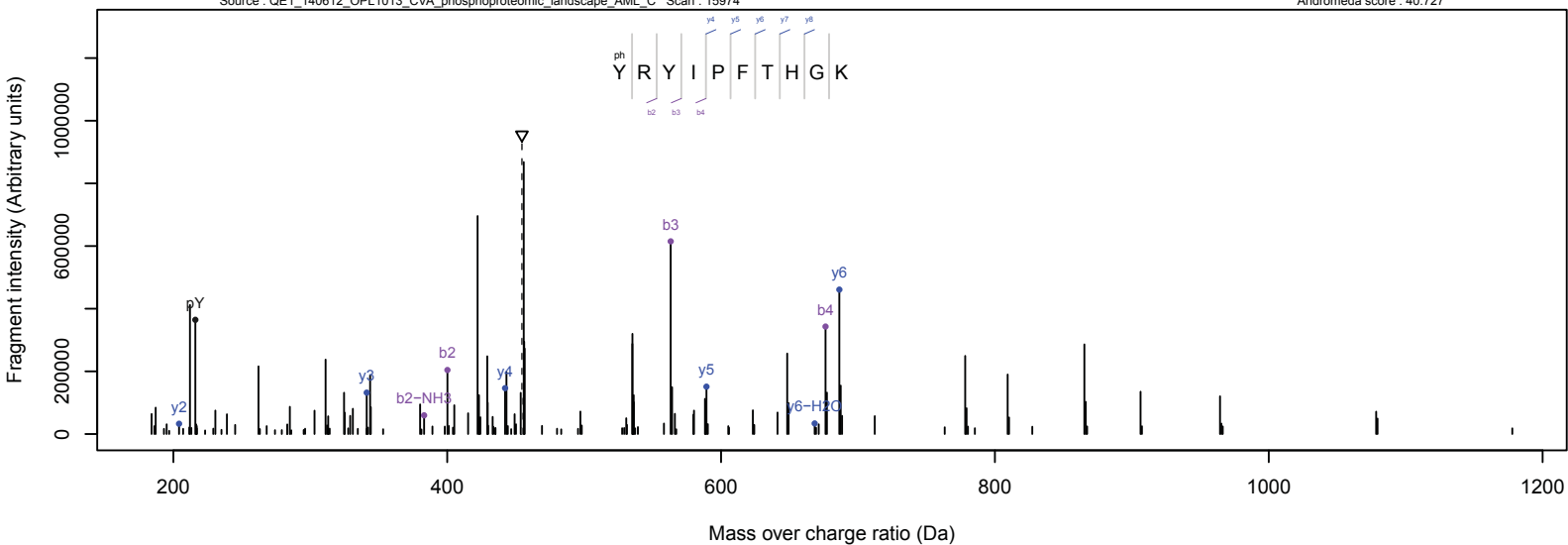
**SLVHY(ph)CFDLR-2+**



Y(ph)RYPFTHGK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15974

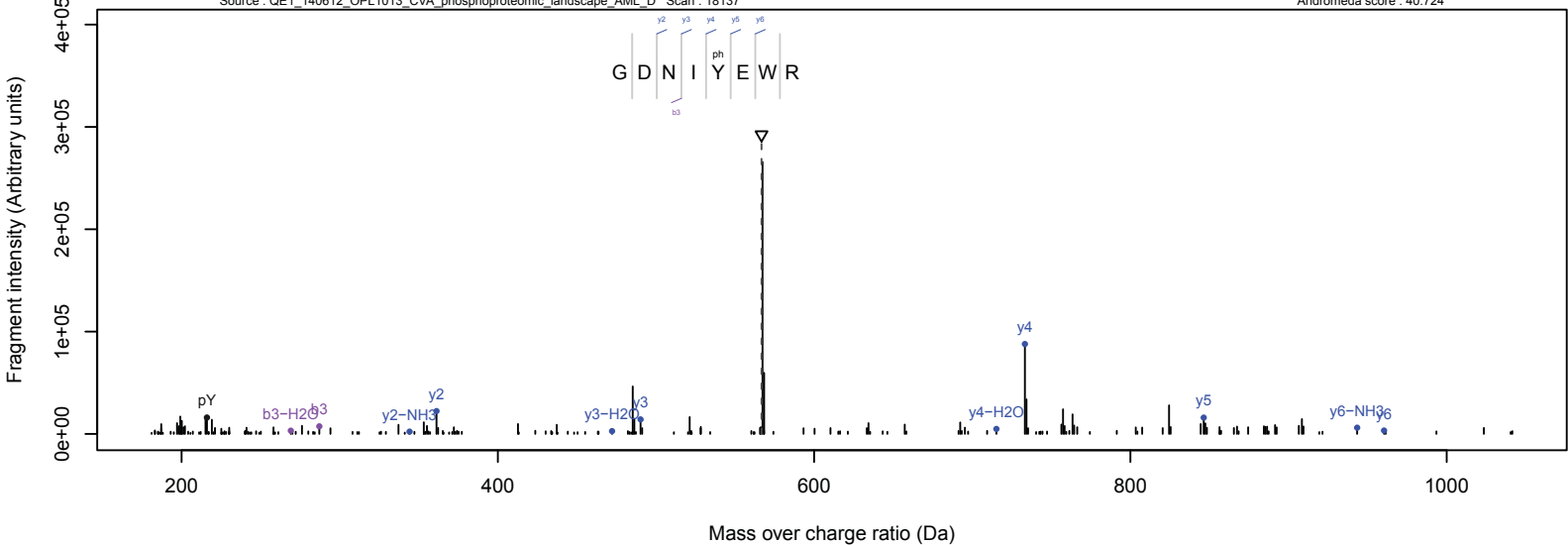
Andromeda score : 40.727



GDNIY(ph)EWR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 18137

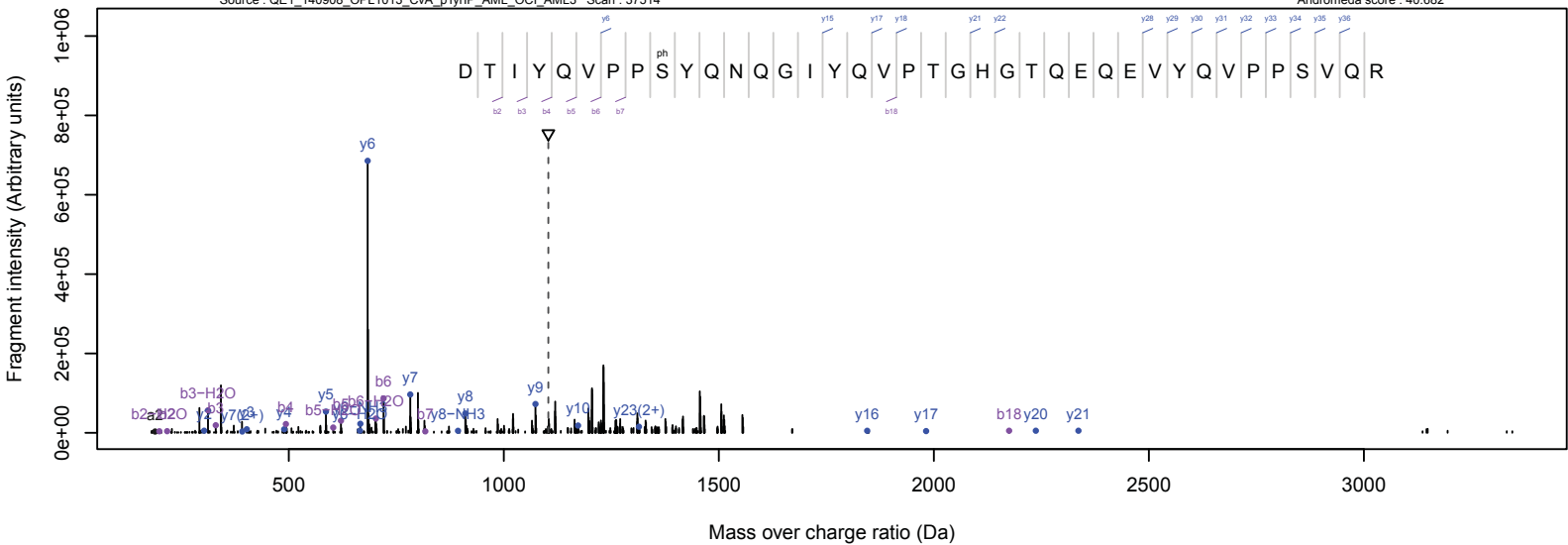
Andromeda score : 40.724



DTIQVPPS(ph)YQNQGIYQVPTGHGTQEVEYQVPPSVQR-4+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_OCI\_AML3 Scan : 37514

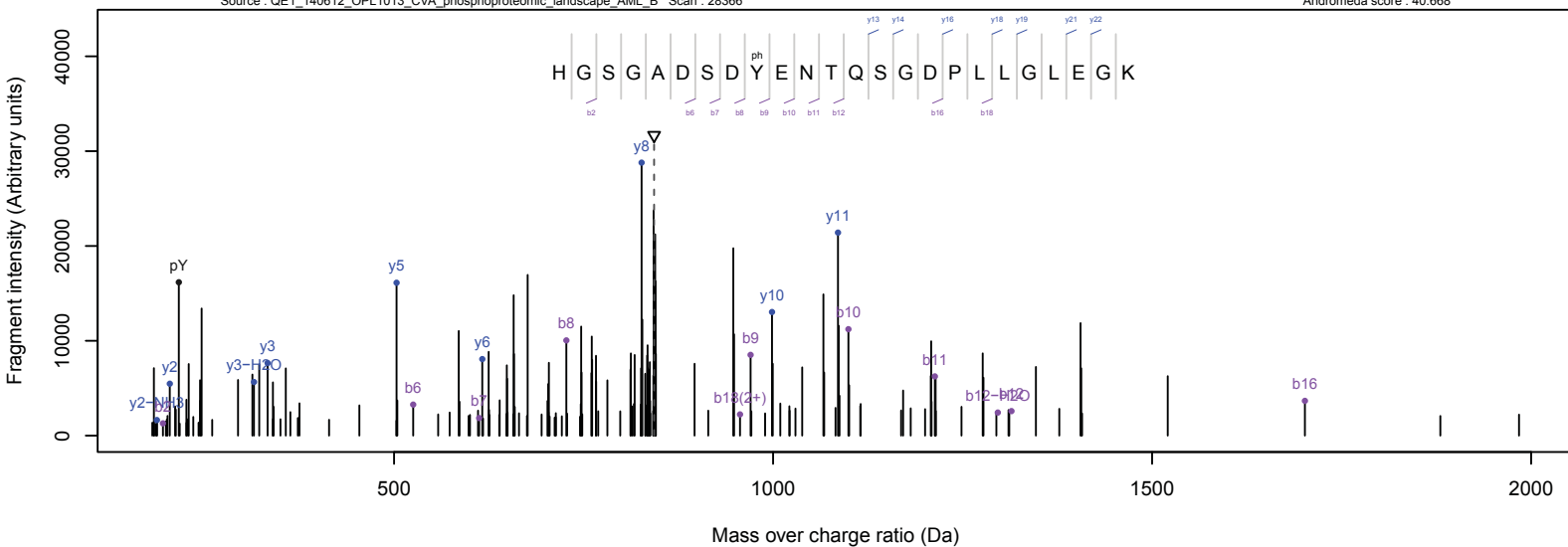
Andromeda score : 40.682



## HGSGADSDY(ph)ENTQSGDPLLEGLK-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 28366

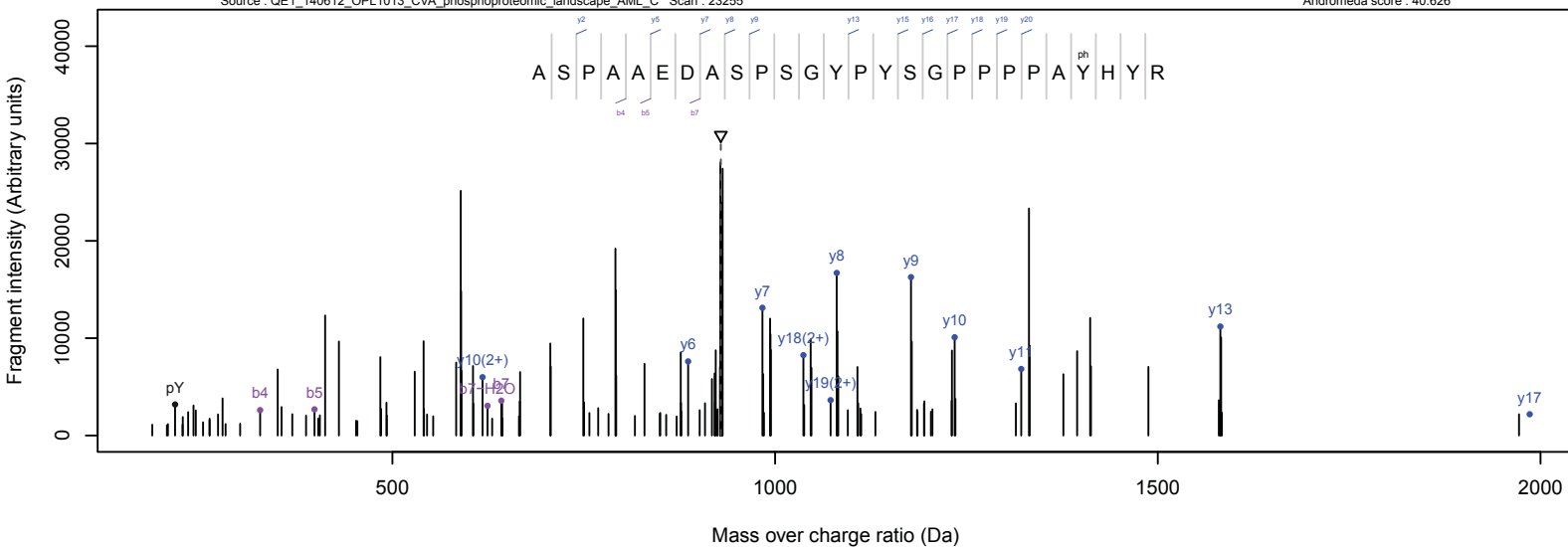
Andromeda score: 40.668



## ASPAEADSPSGYPYSGPPPPAY(ph)HYR-3+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 23255

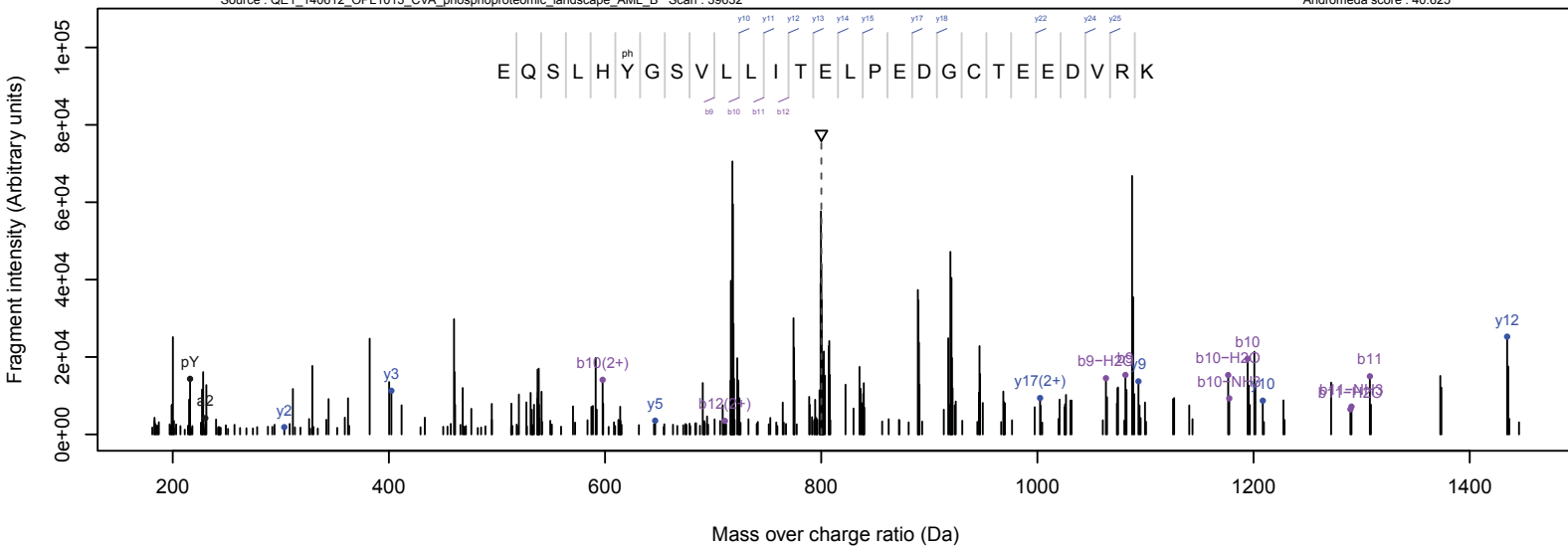
Andromeda score: 40.626



## EQSLHY(ph)GSVLLITELPEDGCTEEDVRK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan: 39632

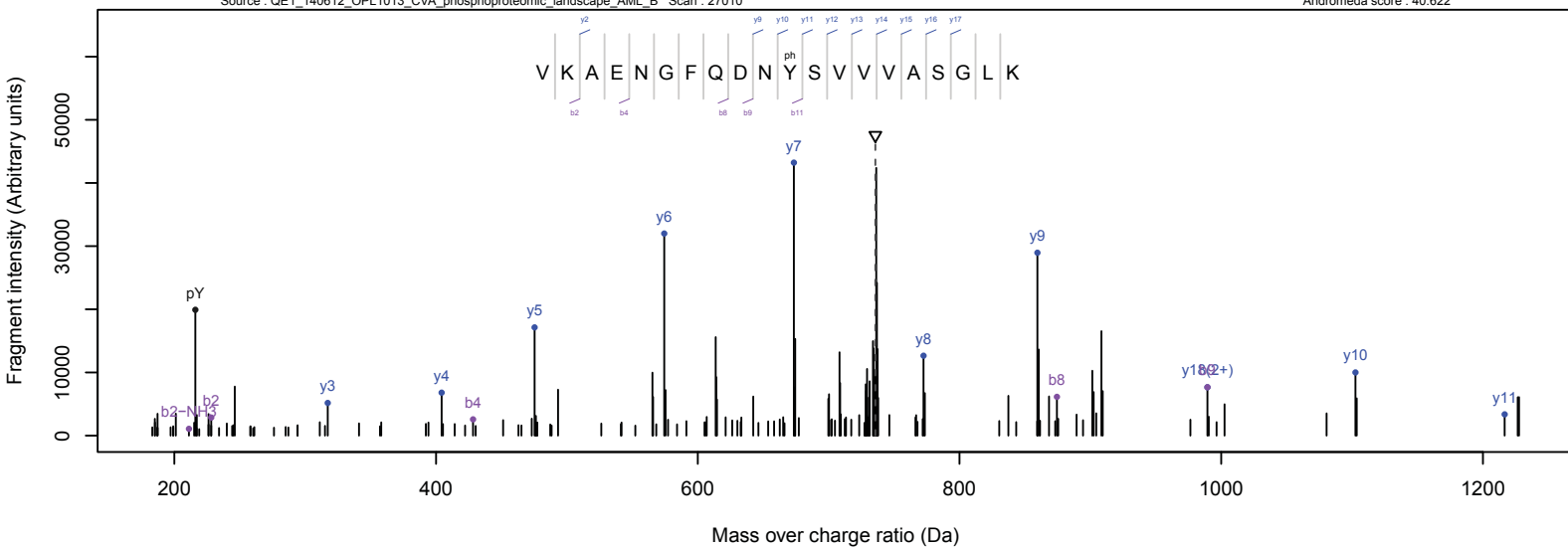
Andromeda score: 40.625



## VKAENGFQDNY(ph)SVVASGLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 27010

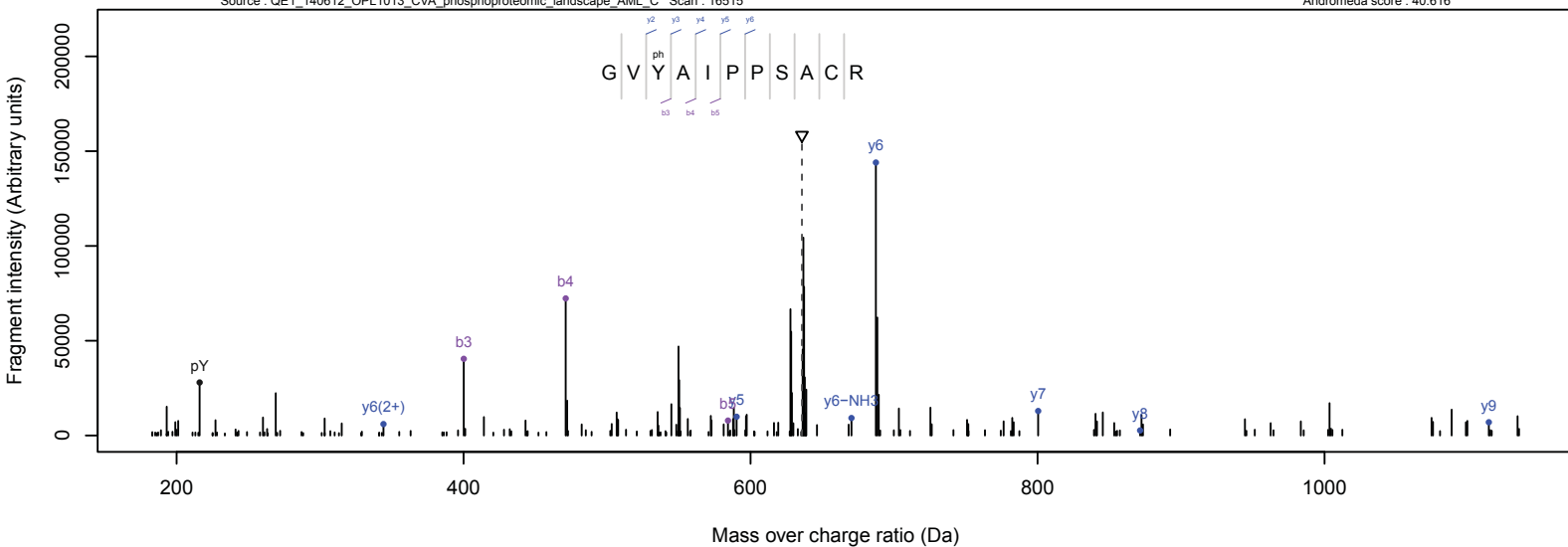
Andromeda score : 40.622



## GVY(ph)AIPPSACR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 16515

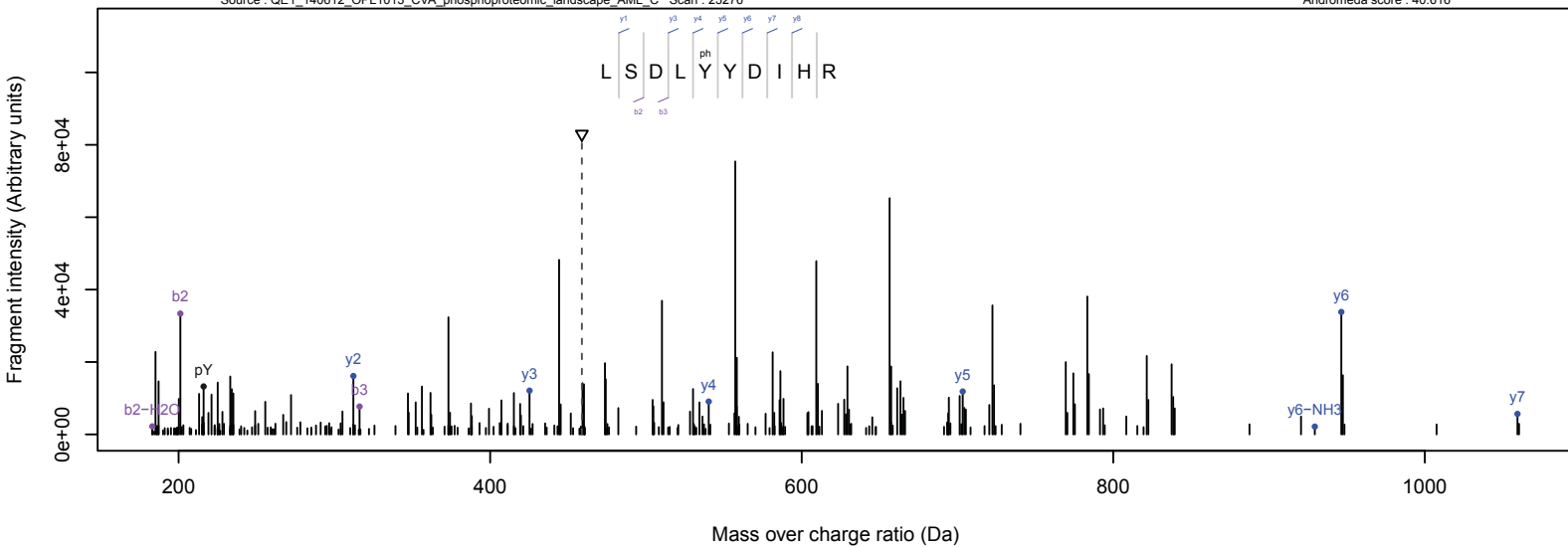
Andromeda score : 40.616



## LSDLY(ph)YDIHR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 23276

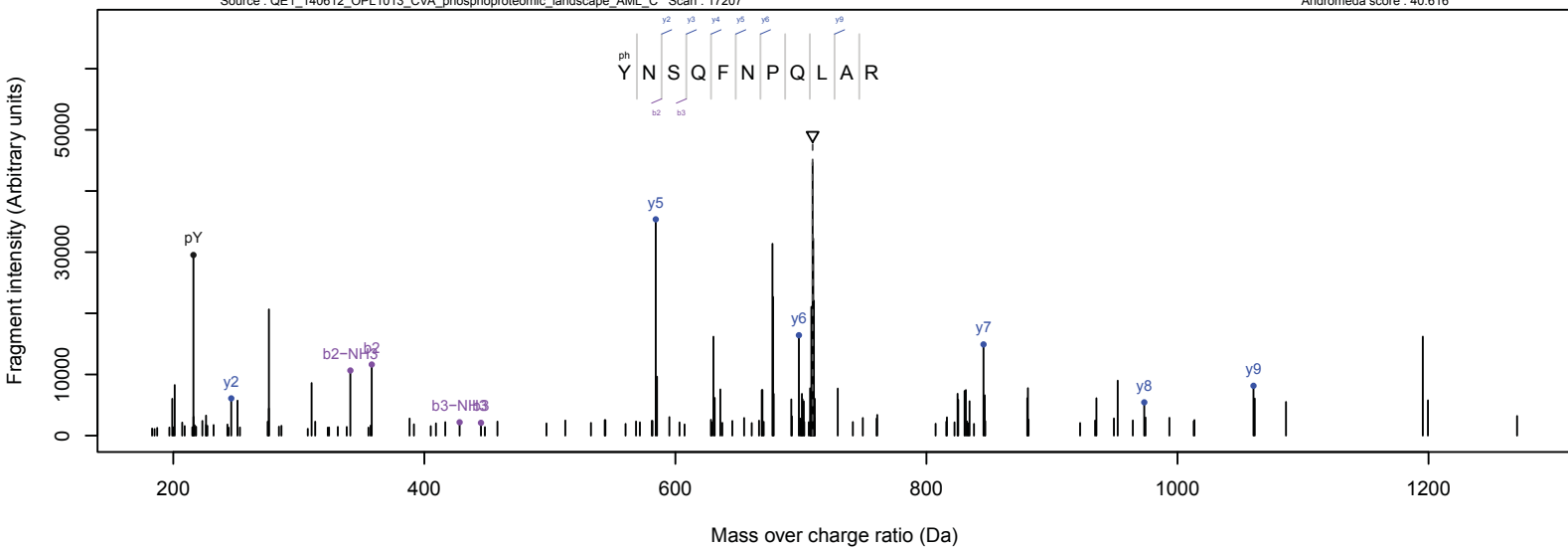
Andromeda score : 40.616



Y(ph)NSQFNPQLAR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 17207

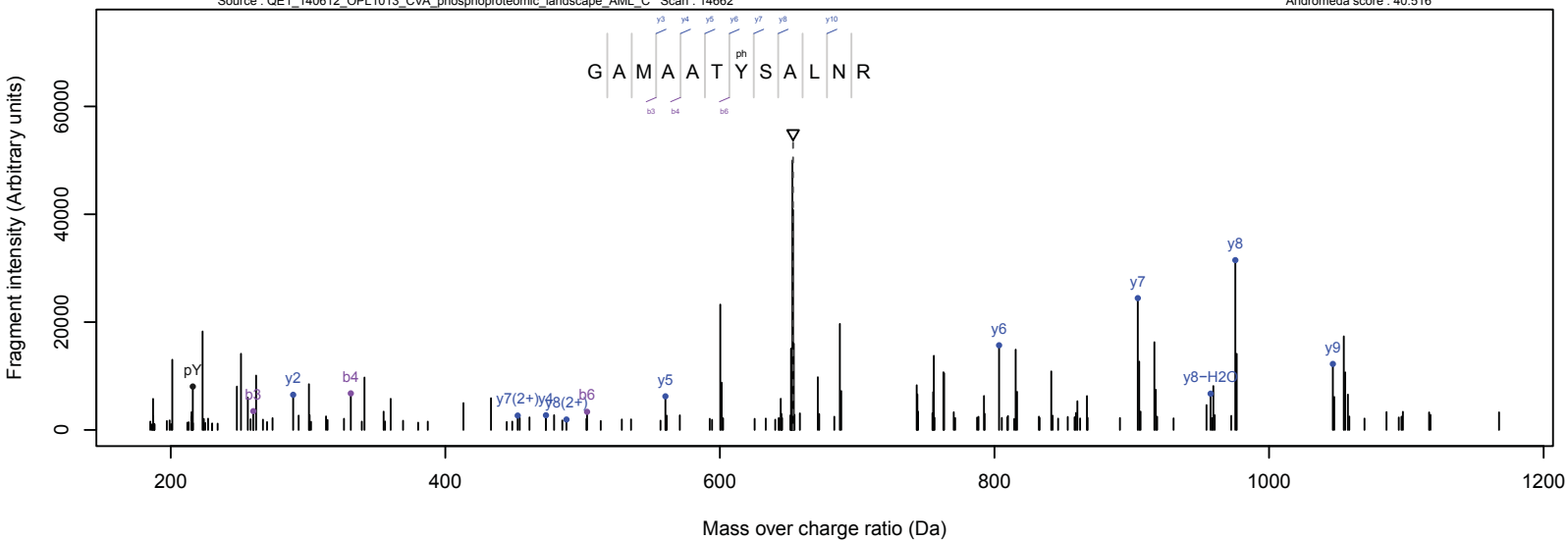
Andromeda score : 40.616



GAMAATY(ph)SALNR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 14662

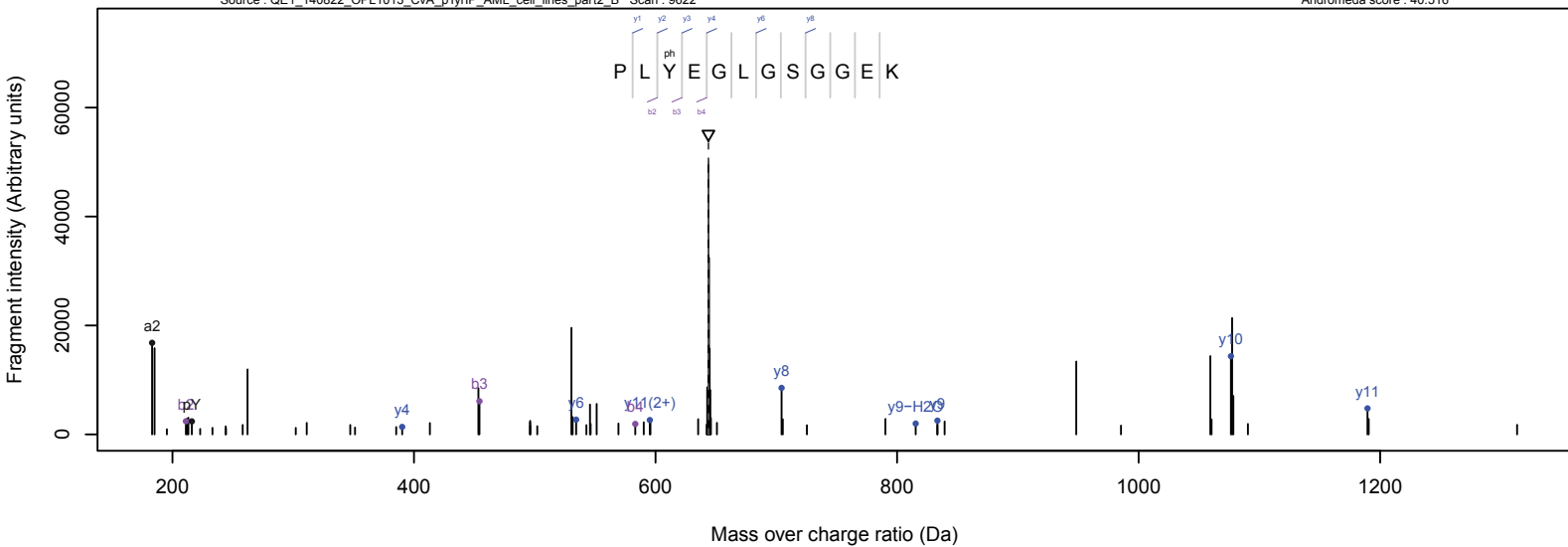
Andromeda score : 40.516



PLY(ph)EGLSGGGEK-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 9622

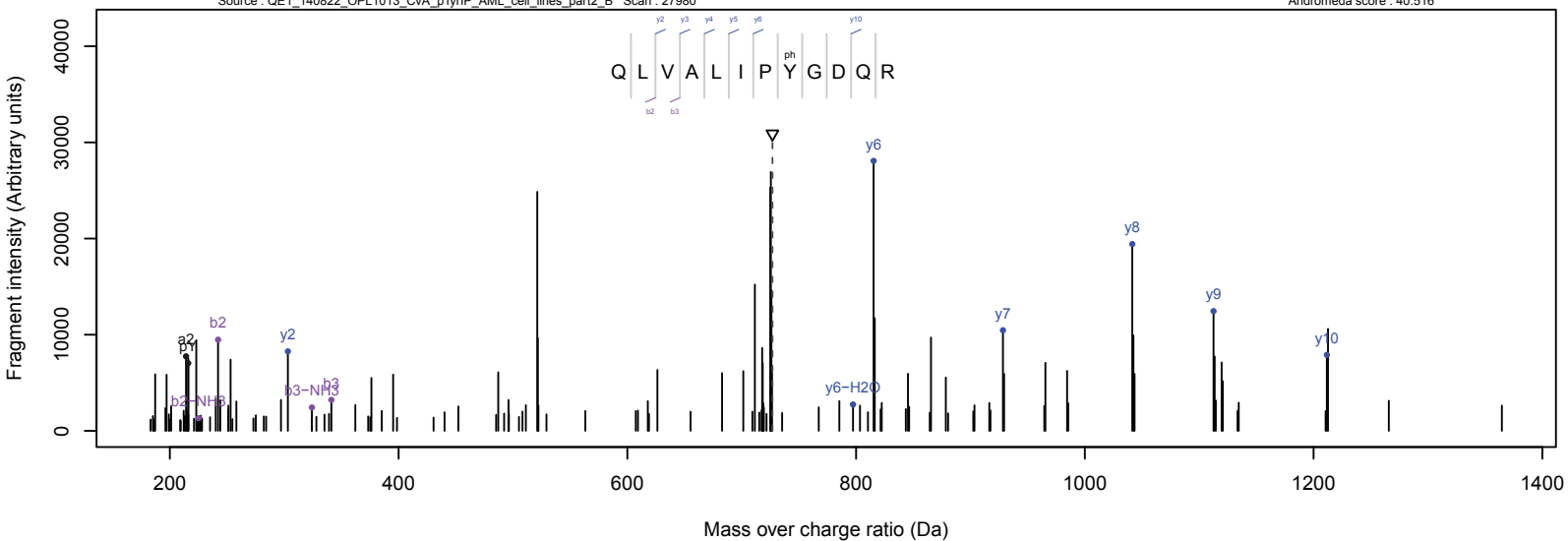
Andromeda score : 40.516



QLVALIPY(ph)GDQR-2+

Source : QE1\_140822\_OPL1013\_CVA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 27980

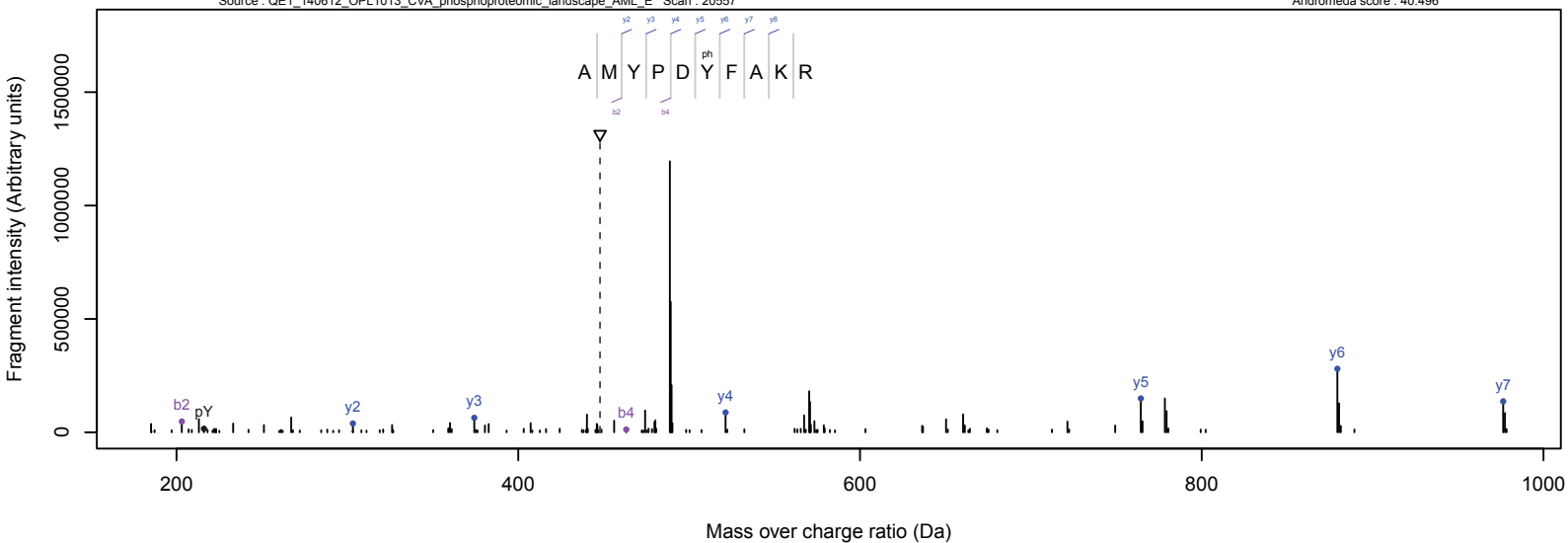
Andromeda score : 40.516



AMYPDY(ph)FAKR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_E Scan : 20557

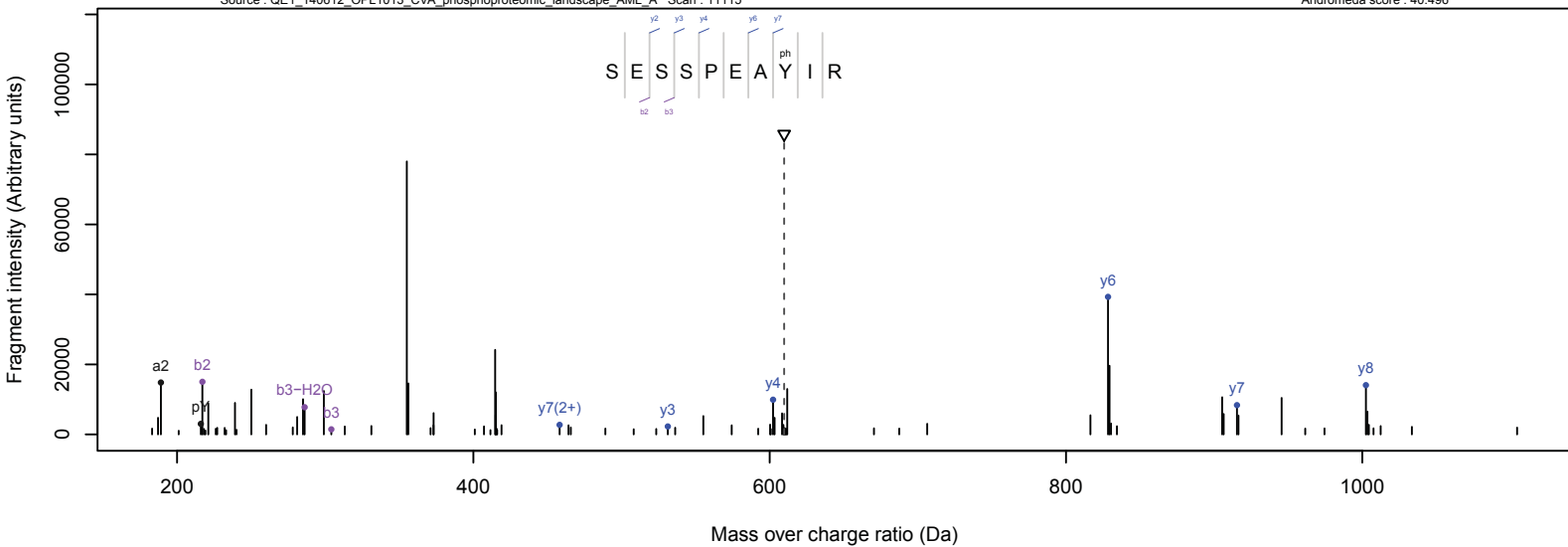
Andromeda score : 40.496



SESSPEAY(ph)IR-2+

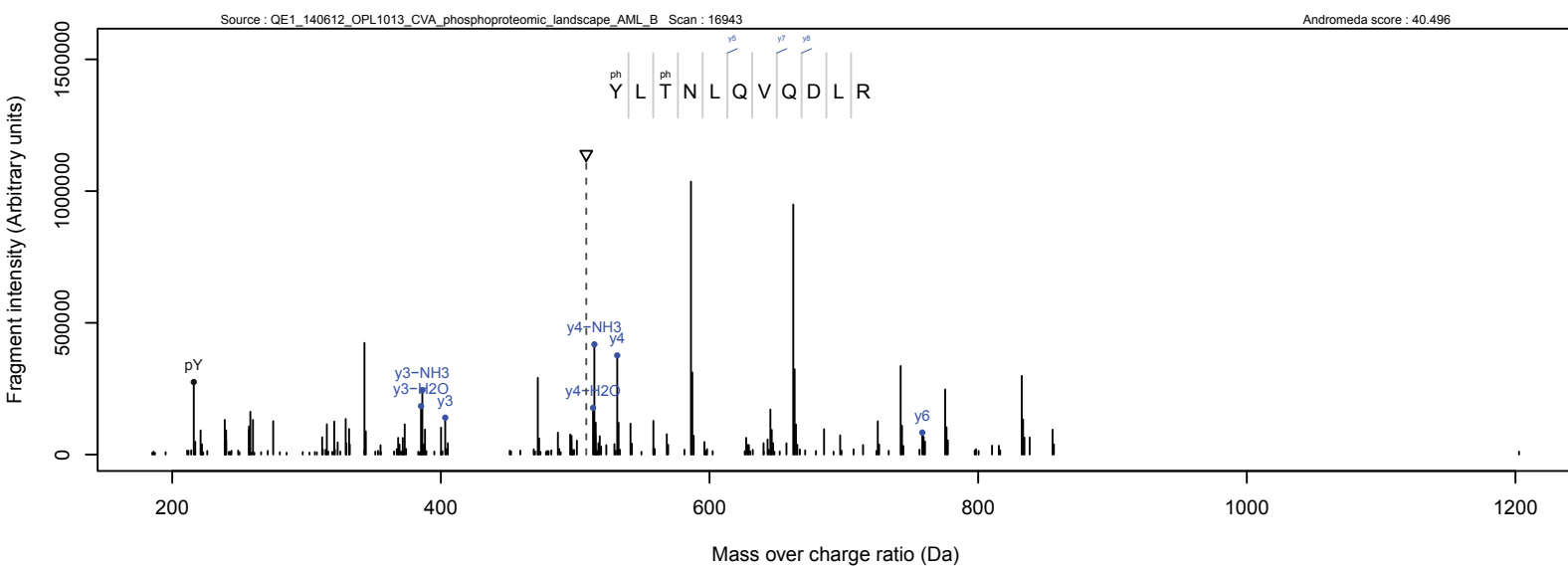
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan : 11115

Andromeda score : 40.496

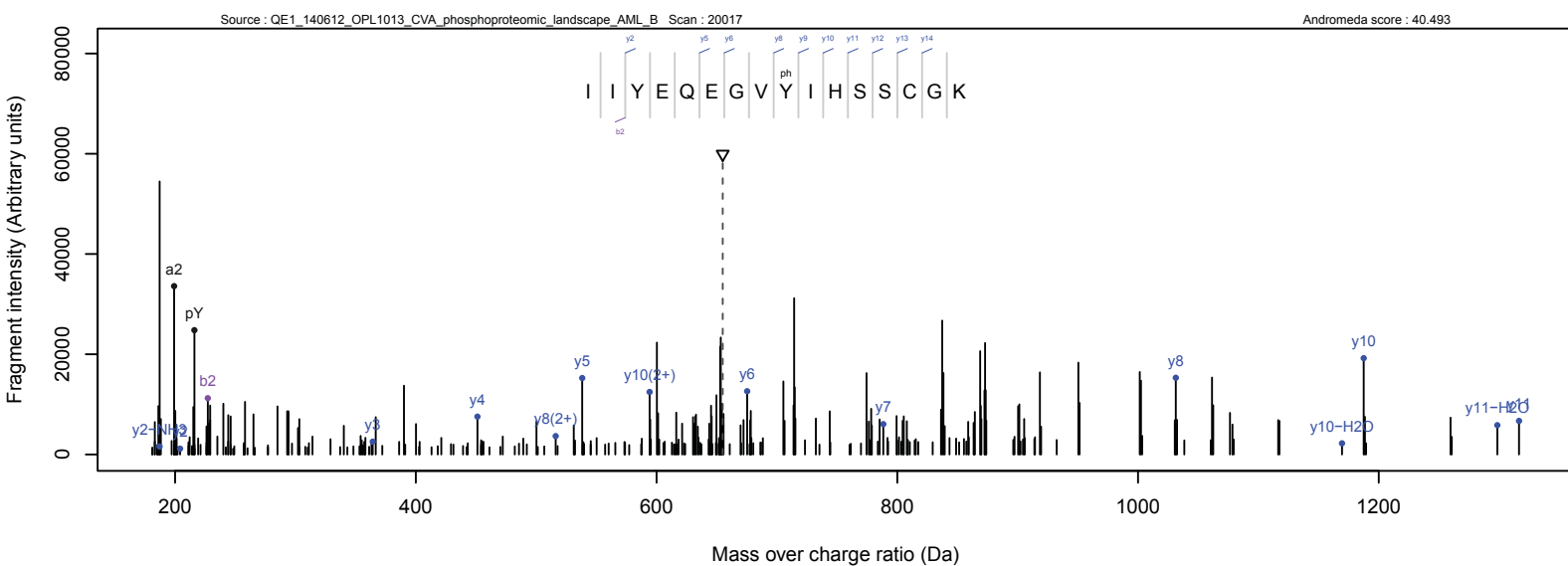




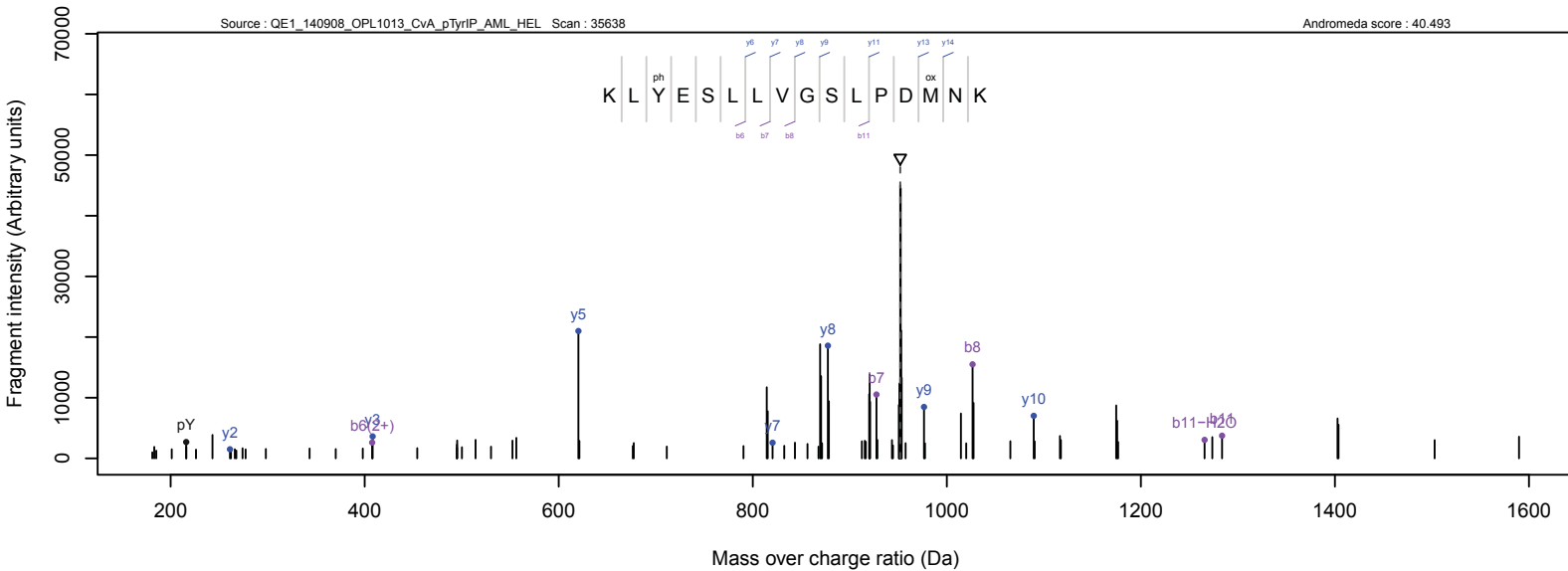
## Y(ph)LT(ph)NLQVQDLR-3+



## IIYEQGVY(ph)HSSCGK-3+



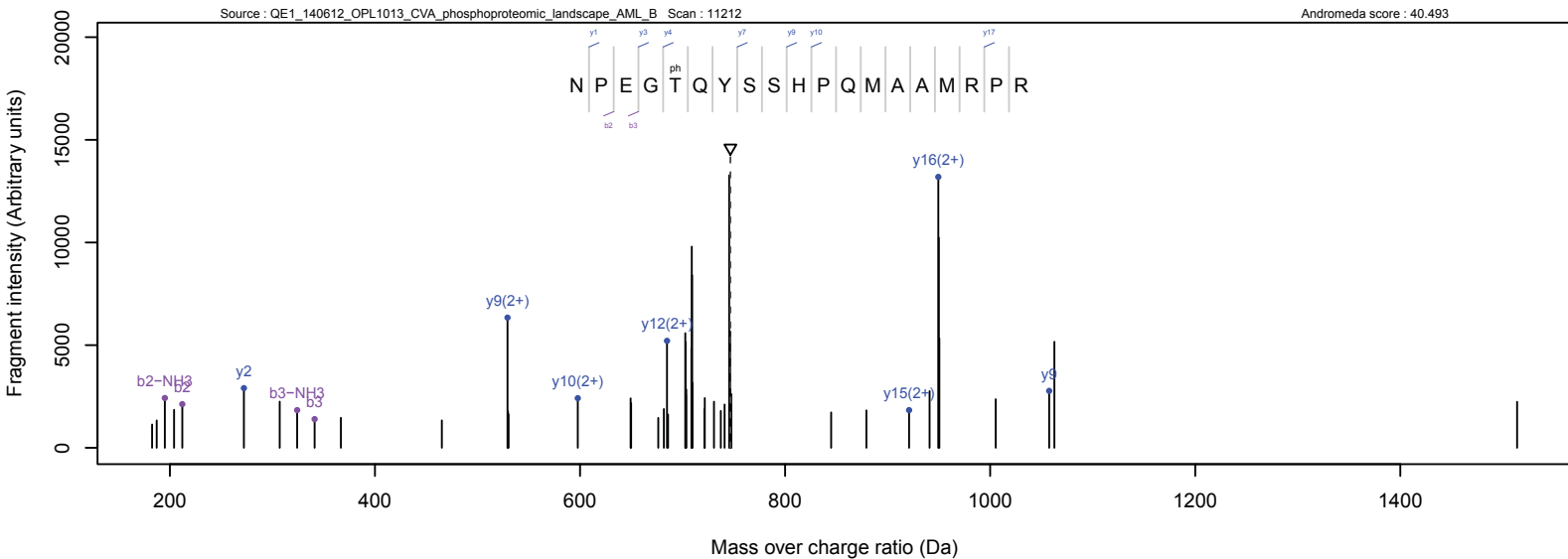
## KLY(ph)ESLLVGLPDM(ox)NK-2+



## NPEGT(ph)QYSSHPQMAAMRPR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 11212

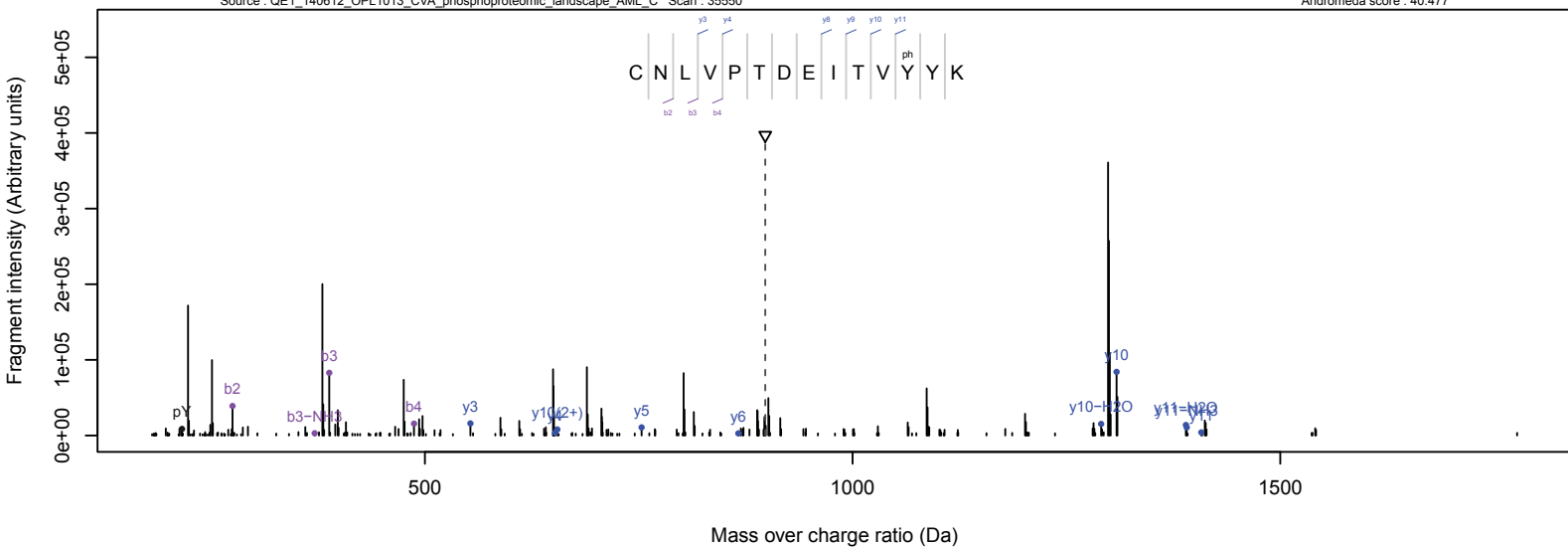
Andromeda score : 40.493



## CNLVPTDEITVYK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 35550

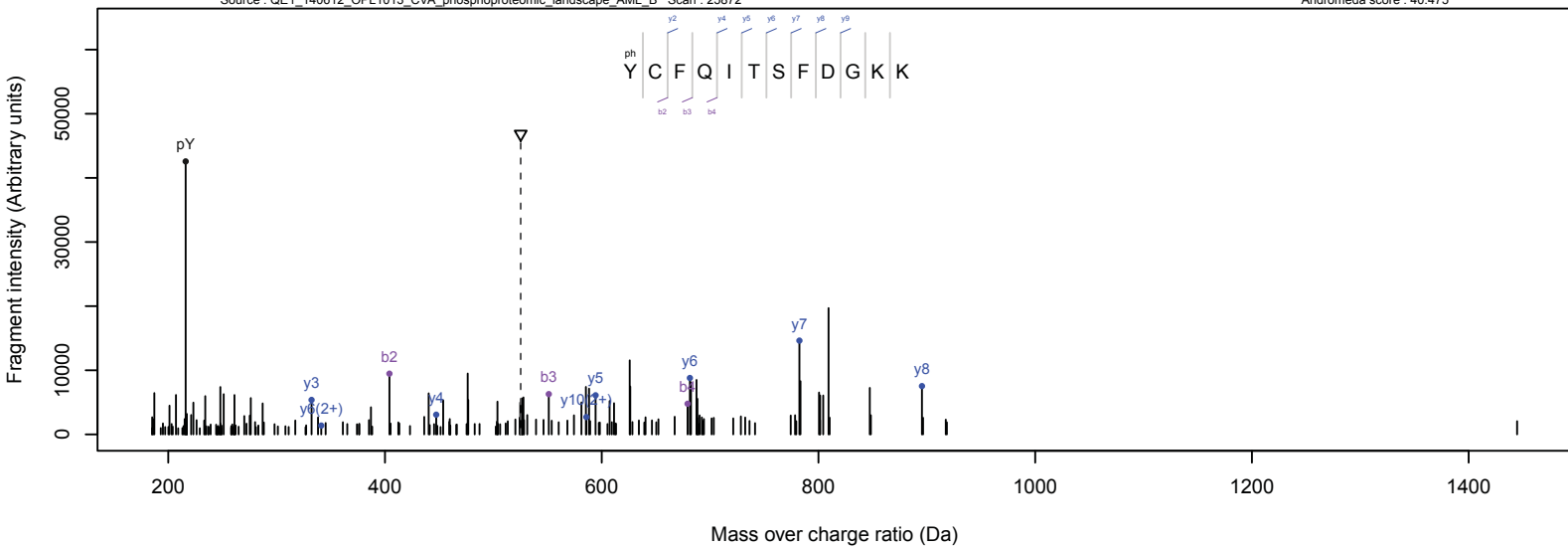
Andromeda score : 40.477



## Y(ph)CFQITSFDGKK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 25872

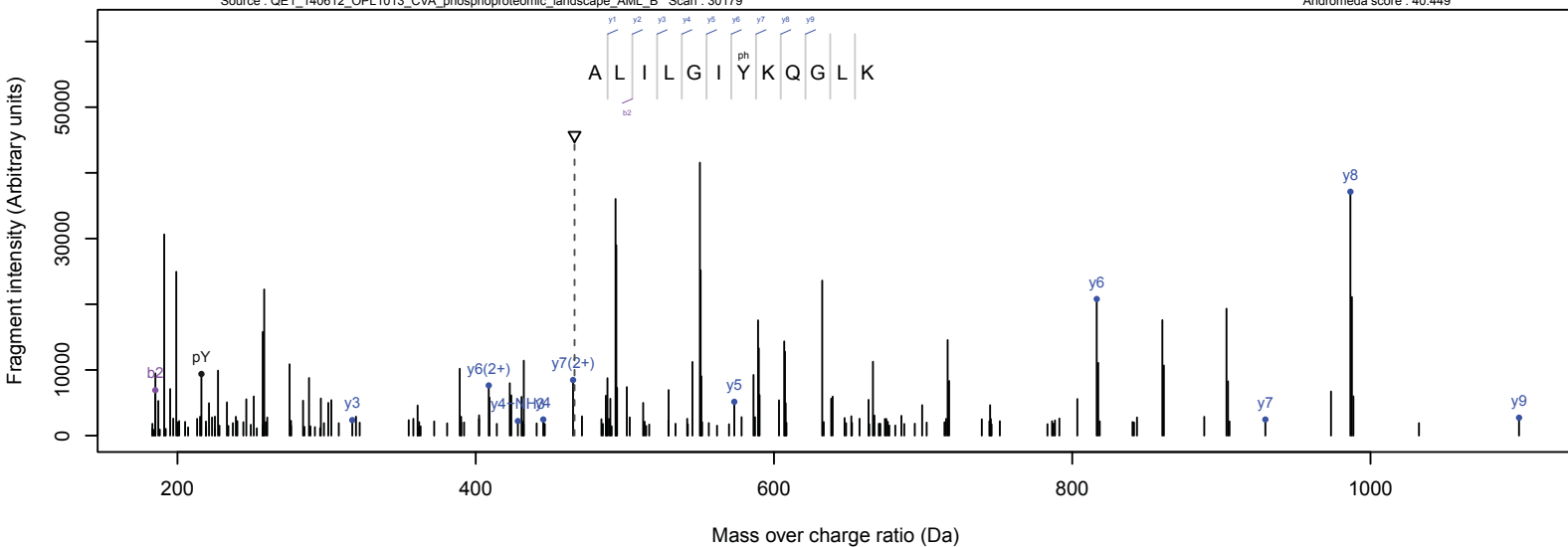
Andromeda score : 40.475



ALILGIY(ph)KQGLK-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 30179

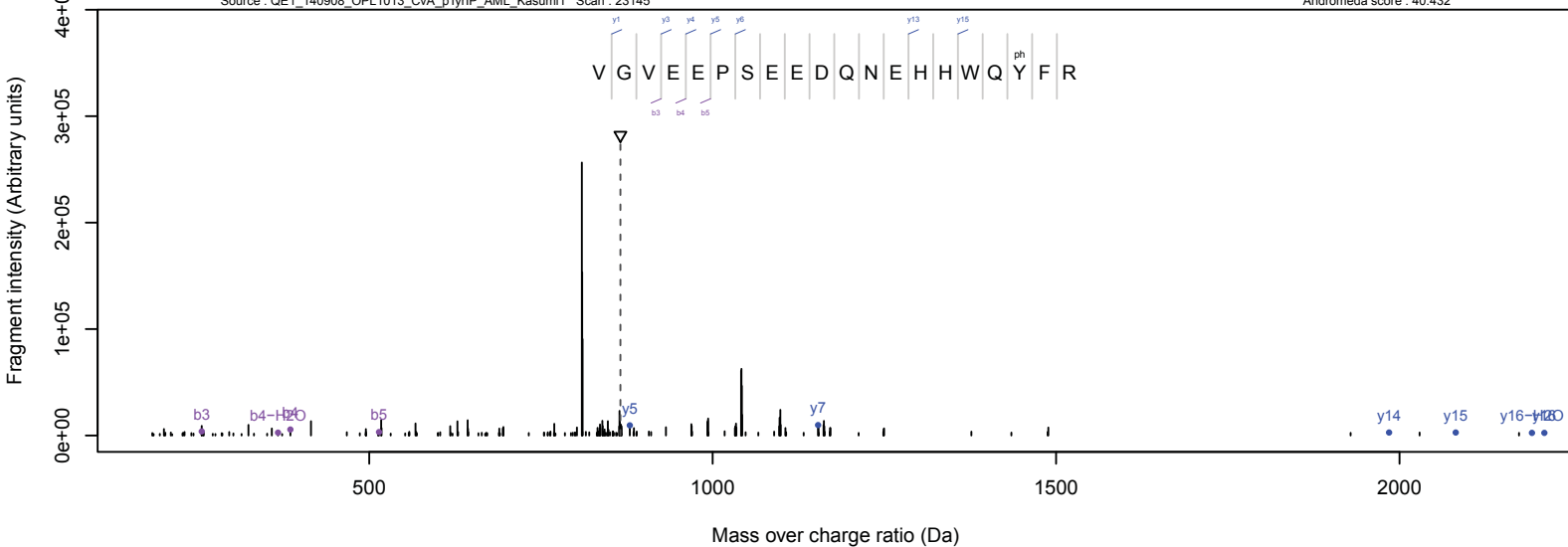
Andromeda score : 40.449



VGVEEPSEEDQNEHHWQYFR-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan : 23145

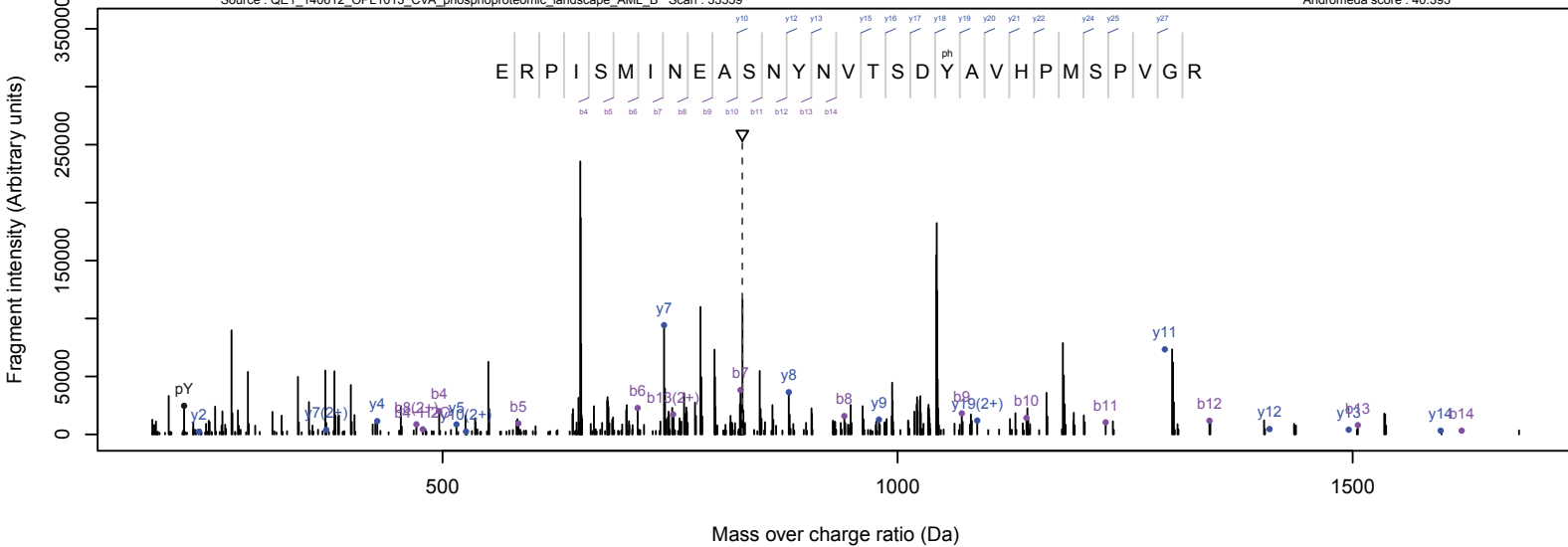
Andromeda score : 40.432



ERPISMINEASNYNTSDYAVHPMSPVGR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 33559

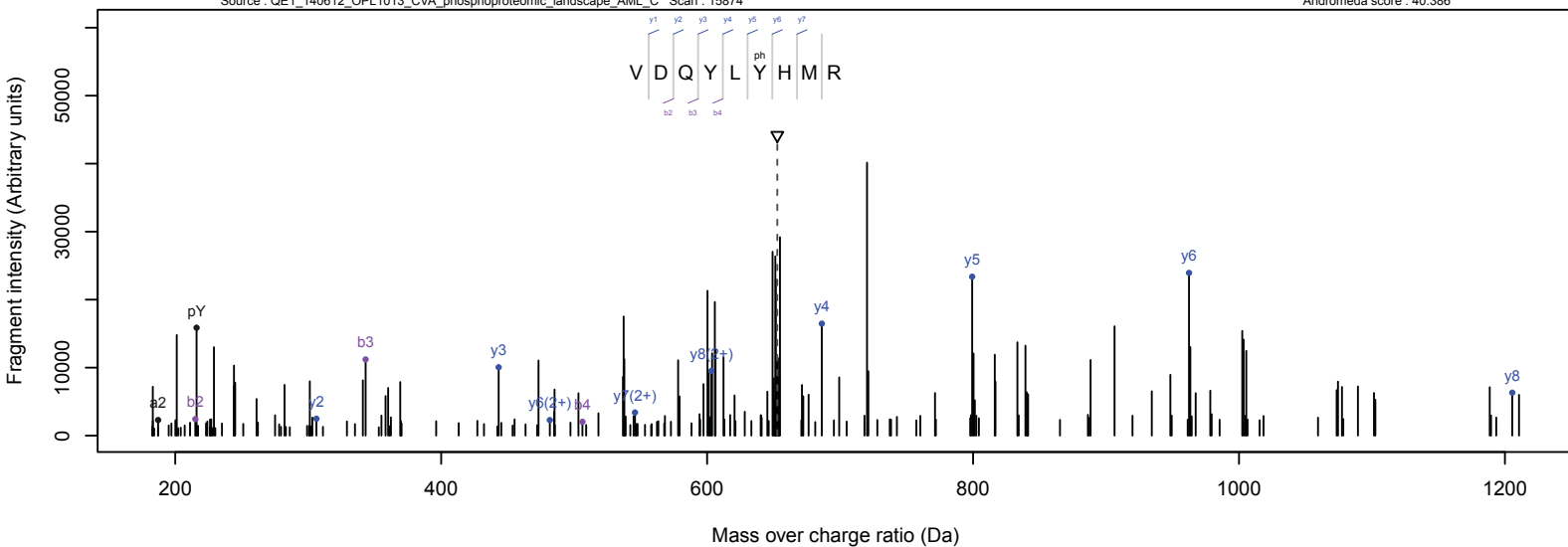
Andromeda score : 40.393



## VDQYLY(ph)HMR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 15874

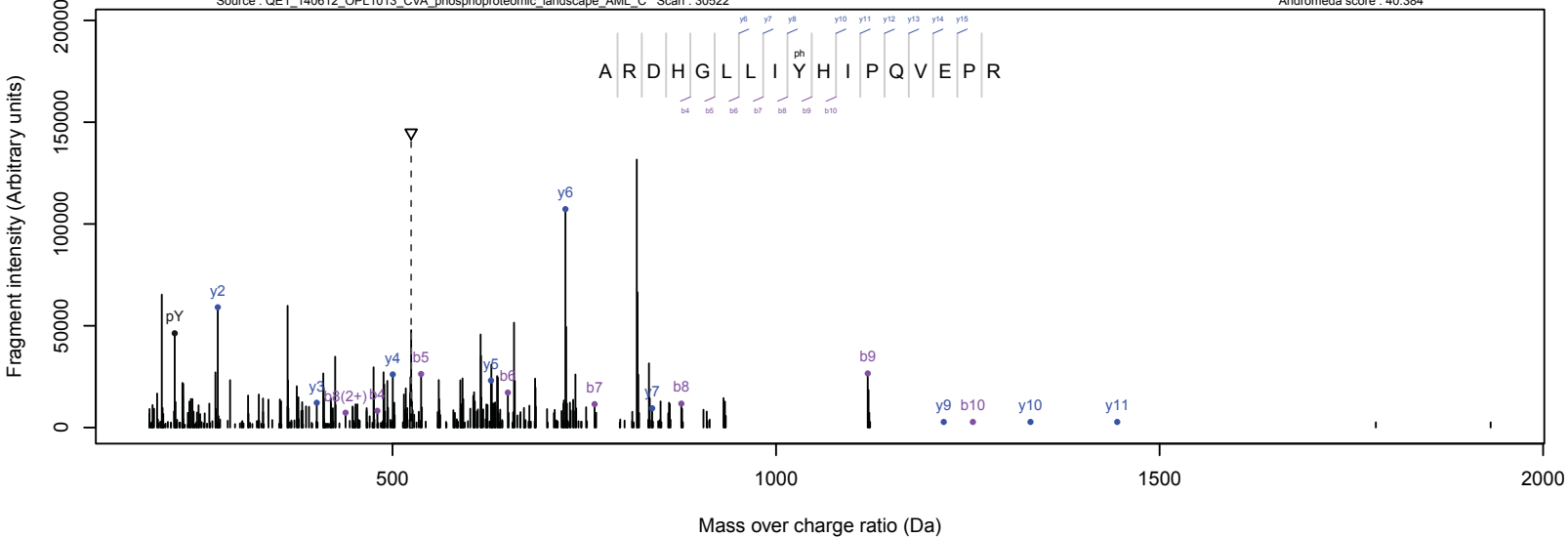
Andromeda score : 40.386



## ARDHGLLIY(ph)HIPQVEPR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30522

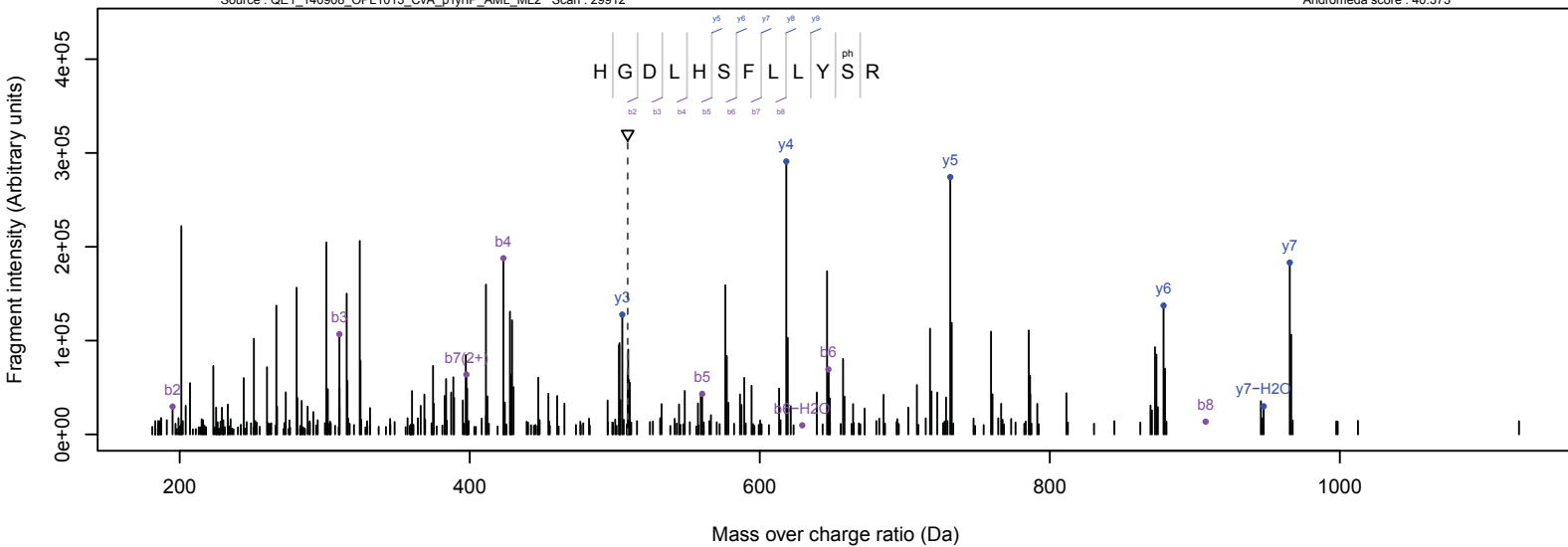
Andromeda score : 40.384



## HGDLSFLLYS(ph)R-3+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_ML2 Scan : 29912

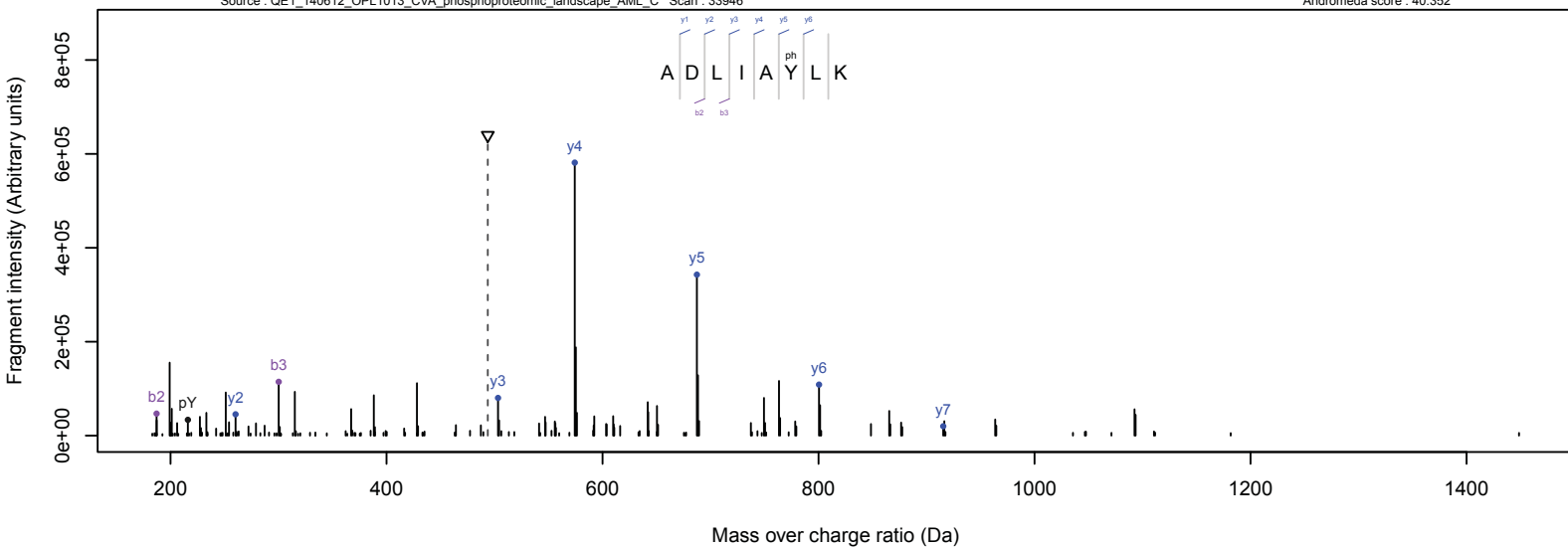
Andromeda score : 40.373



ADLIAY(ph)LK-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 33946

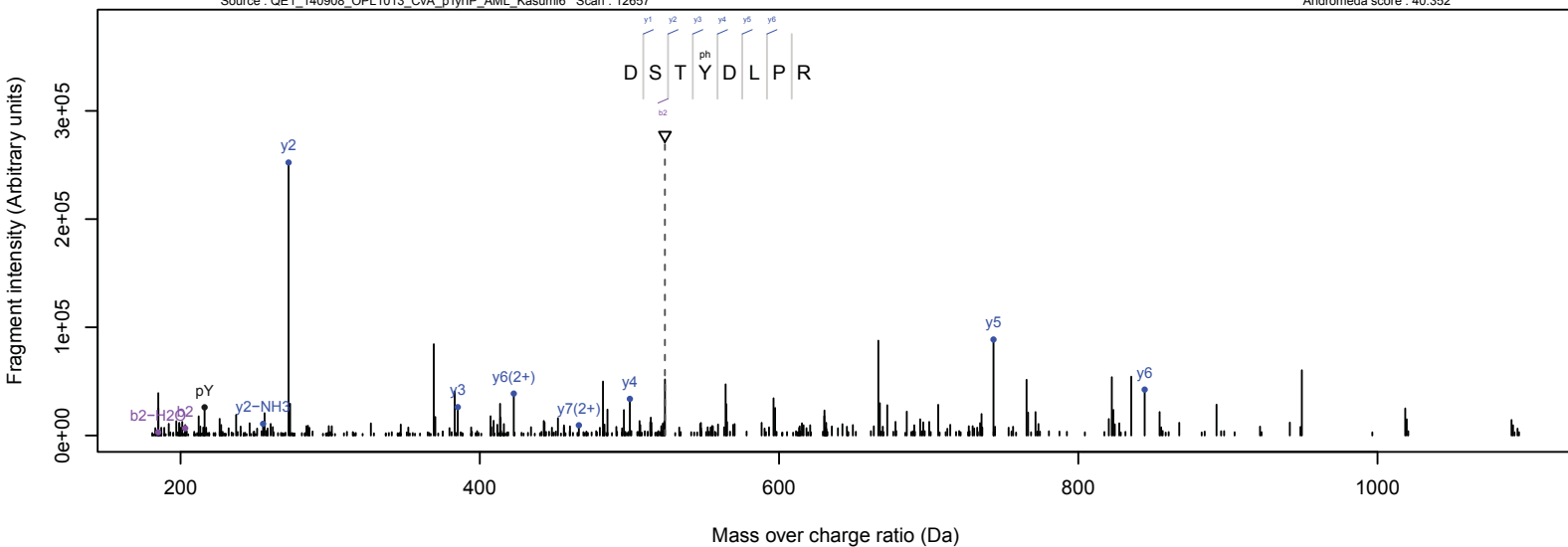
Andromeda score : 40.352



DSTY(ph)DLPR-2+

Source : QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan : 12657

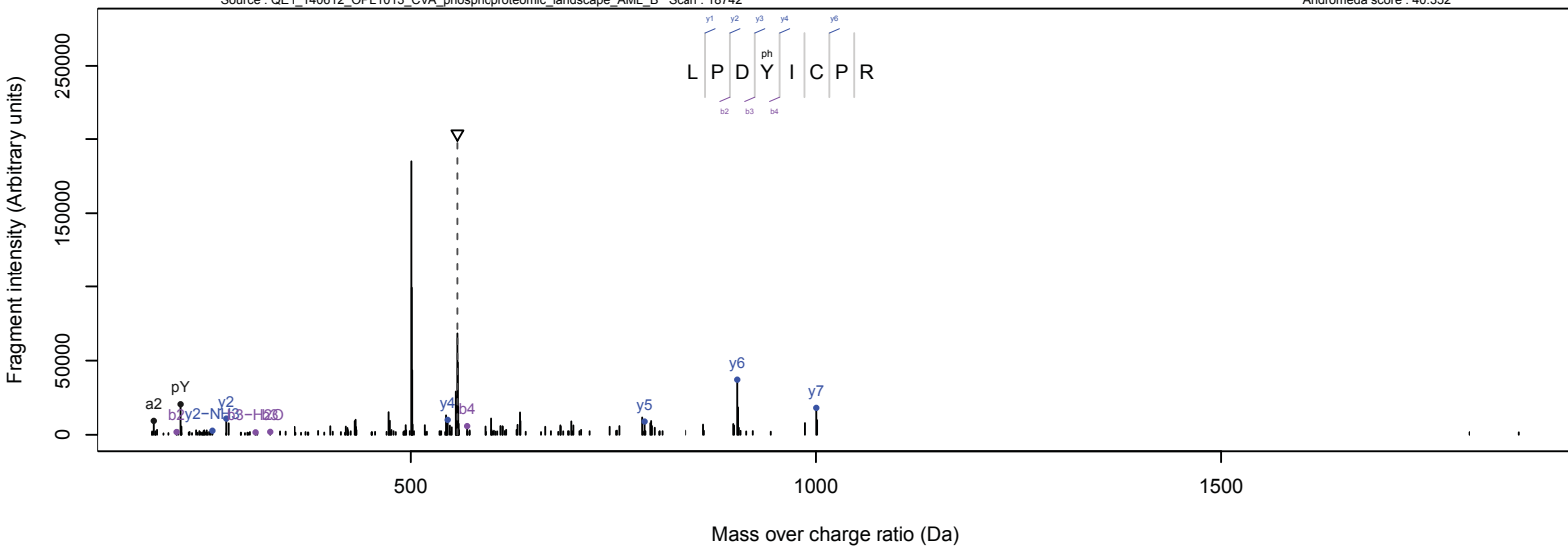
Andromeda score : 40.352



LPDY(ph)ICPR-2+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 18742

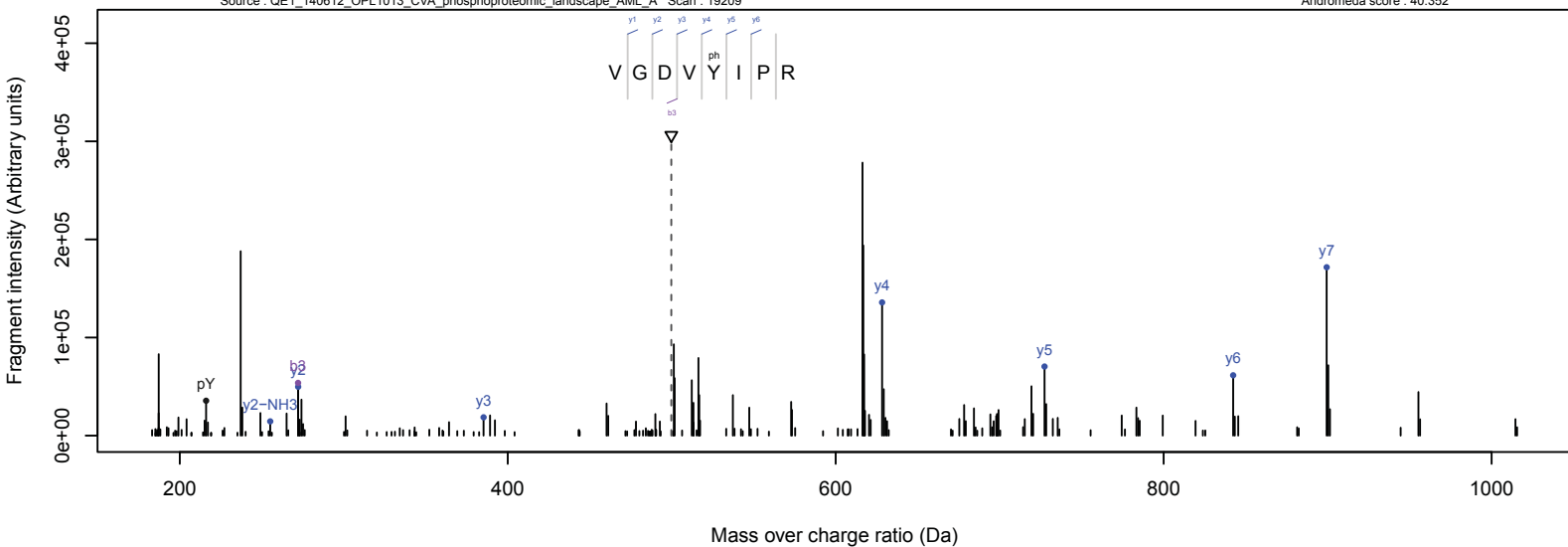
Andromeda score : 40.352



VGDVY(ph)IPR-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_A Scan: 19209

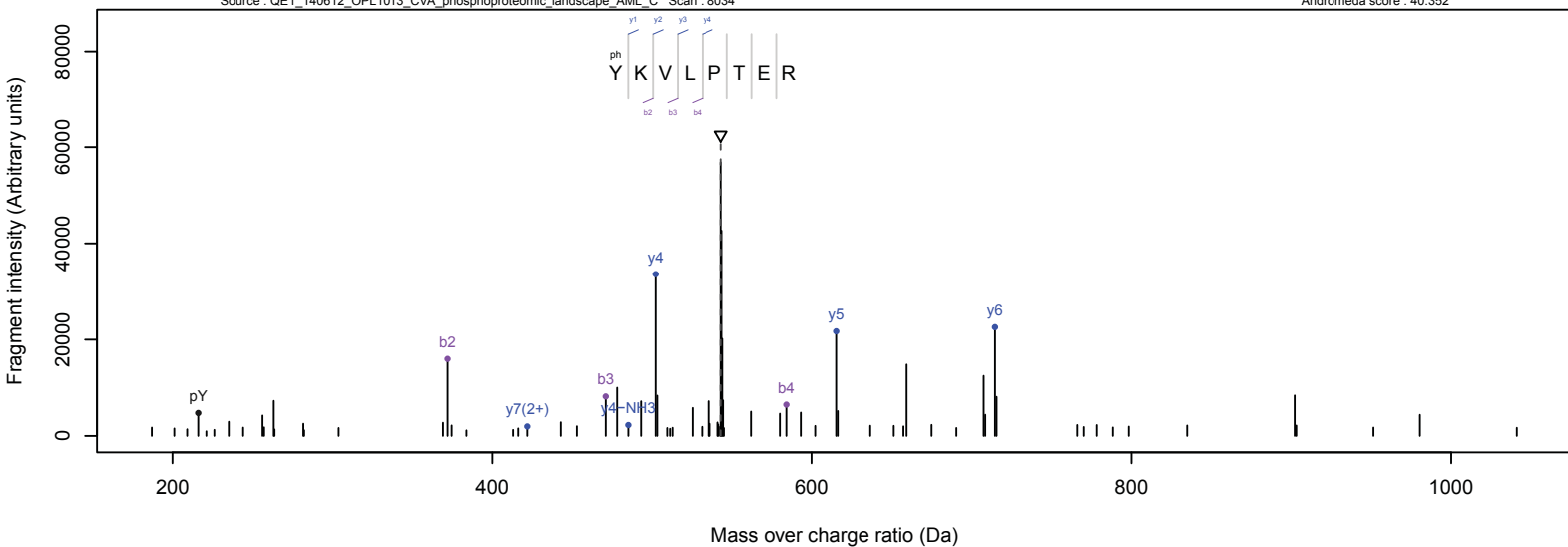
Andromeda score: 40.352



Y(ph)KVLPTER-2+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 8034

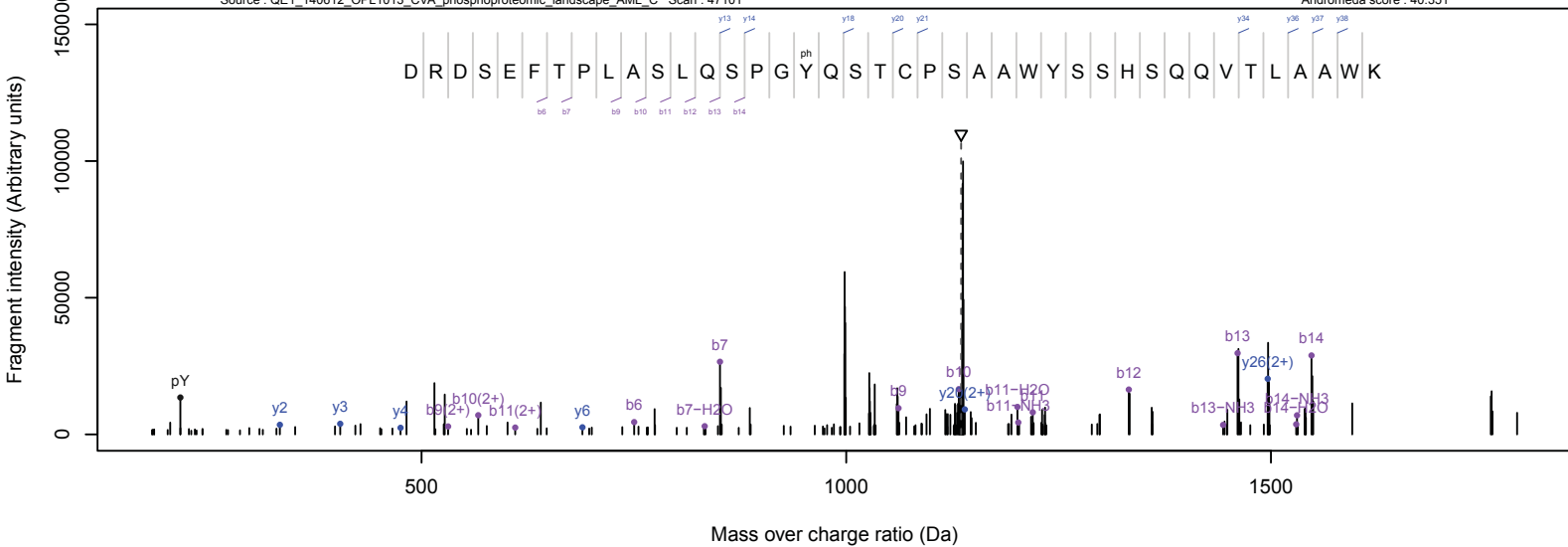
Andromeda score: 40.352



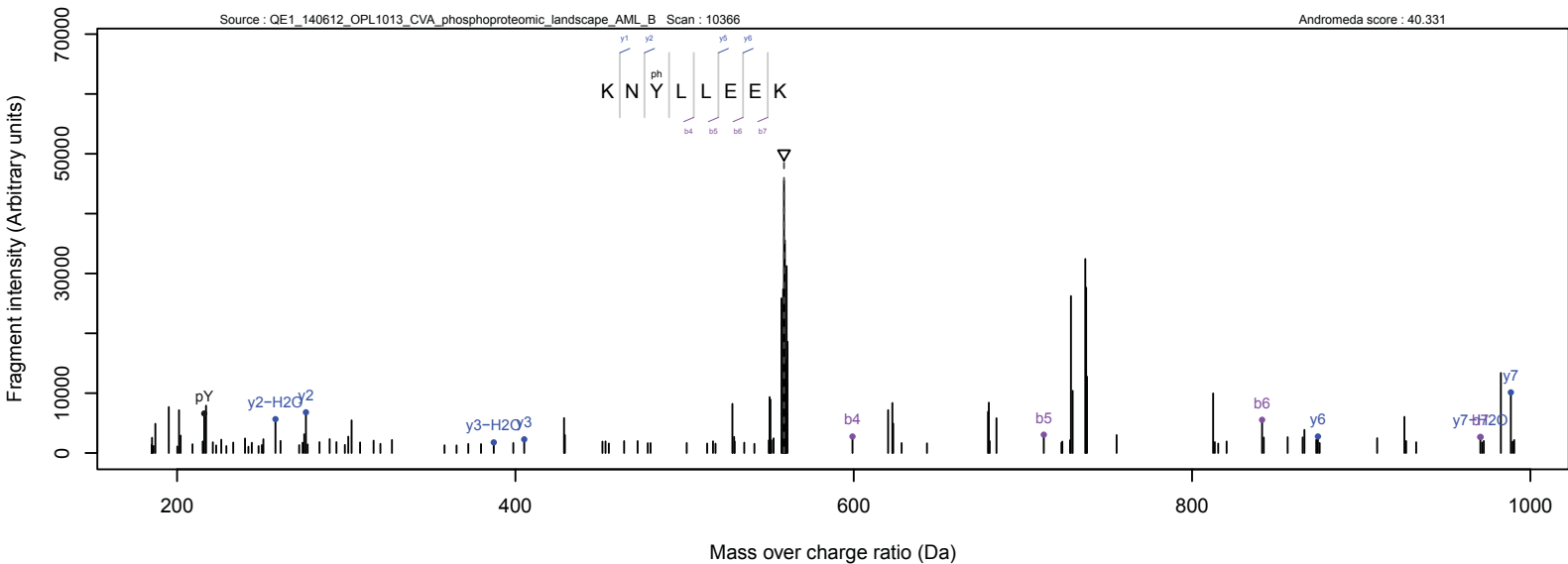
DRDSEFTPLASLQSPGY(ph)QSTCPSAAWYSSHSQQVTLAAWK-4+

Source: QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan: 47101

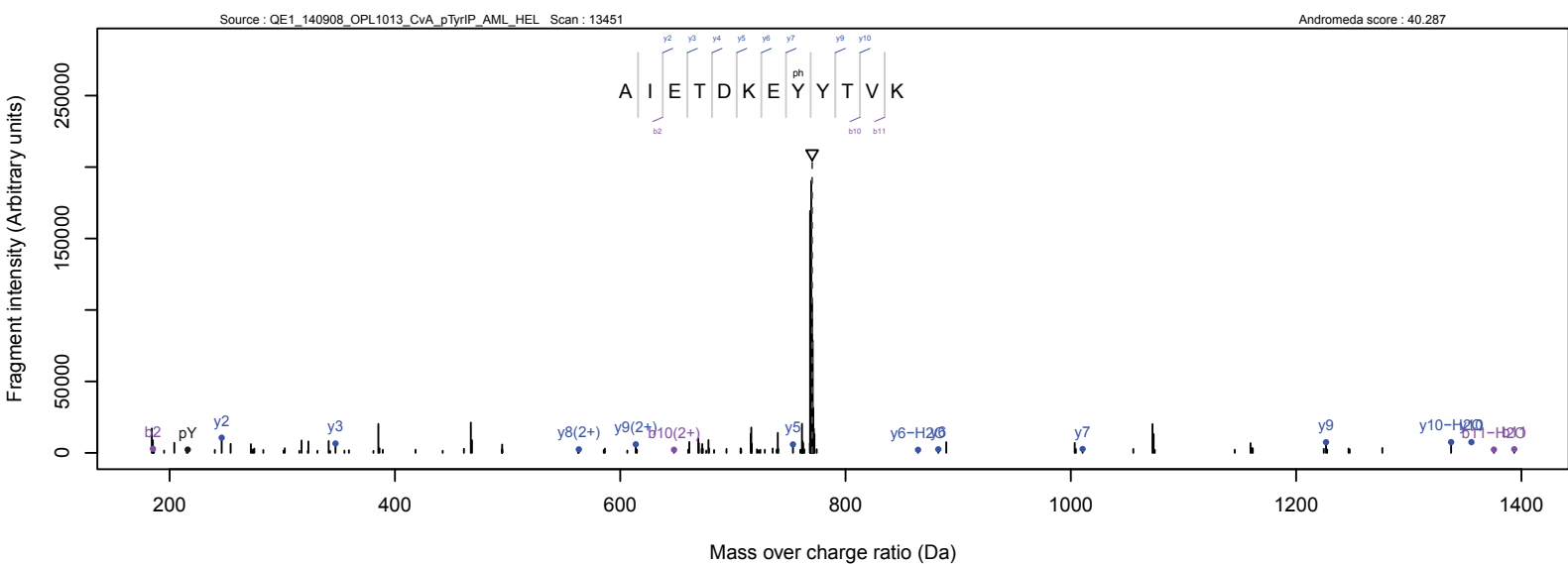
Andromeda score: 40.351



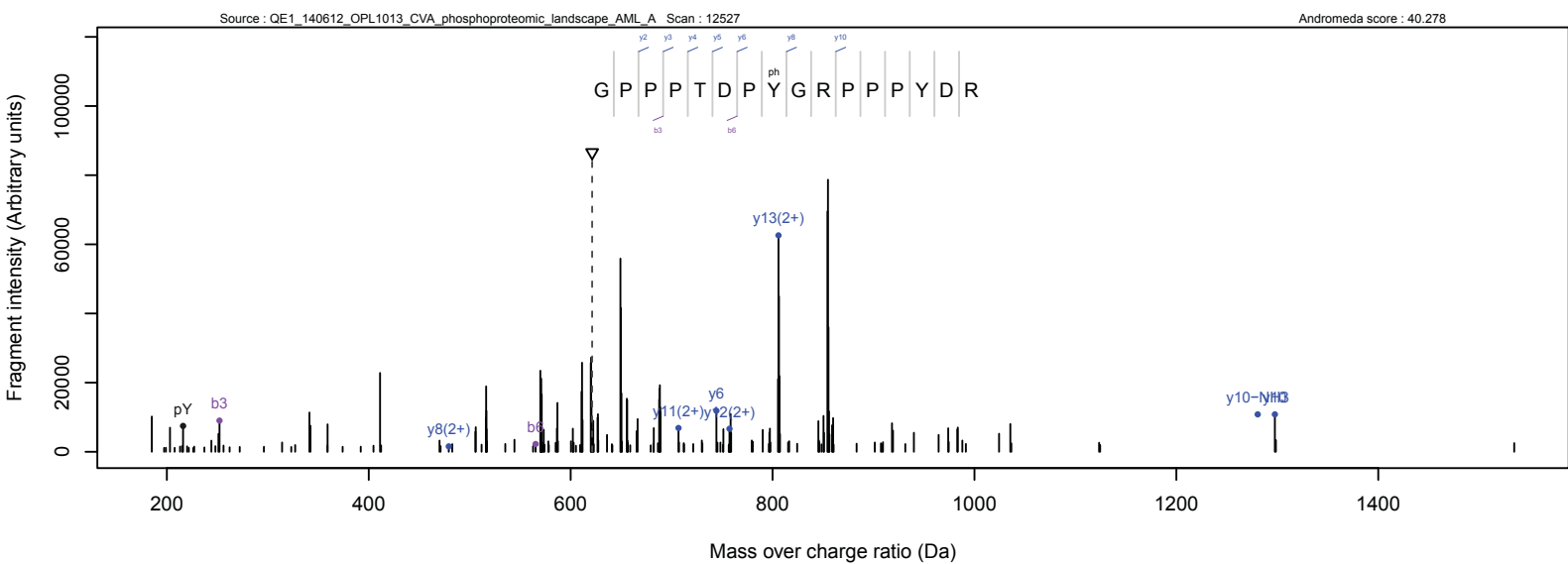
**KNY(ph)LLEEK-2+**



**AIETDKEY(ph)YTVK-2+**



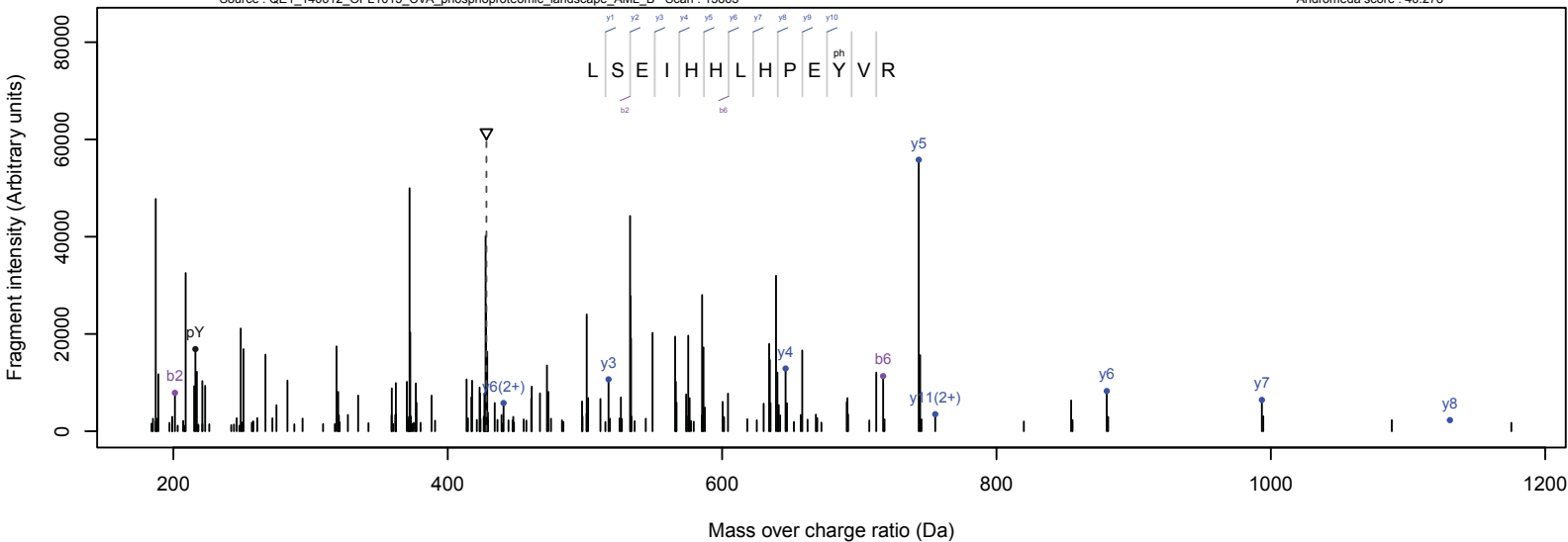
**GPPPTDPY(ph)GRPPPYDR-3+**



## LSEIHLHPEY(ph)VR-4+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 13863

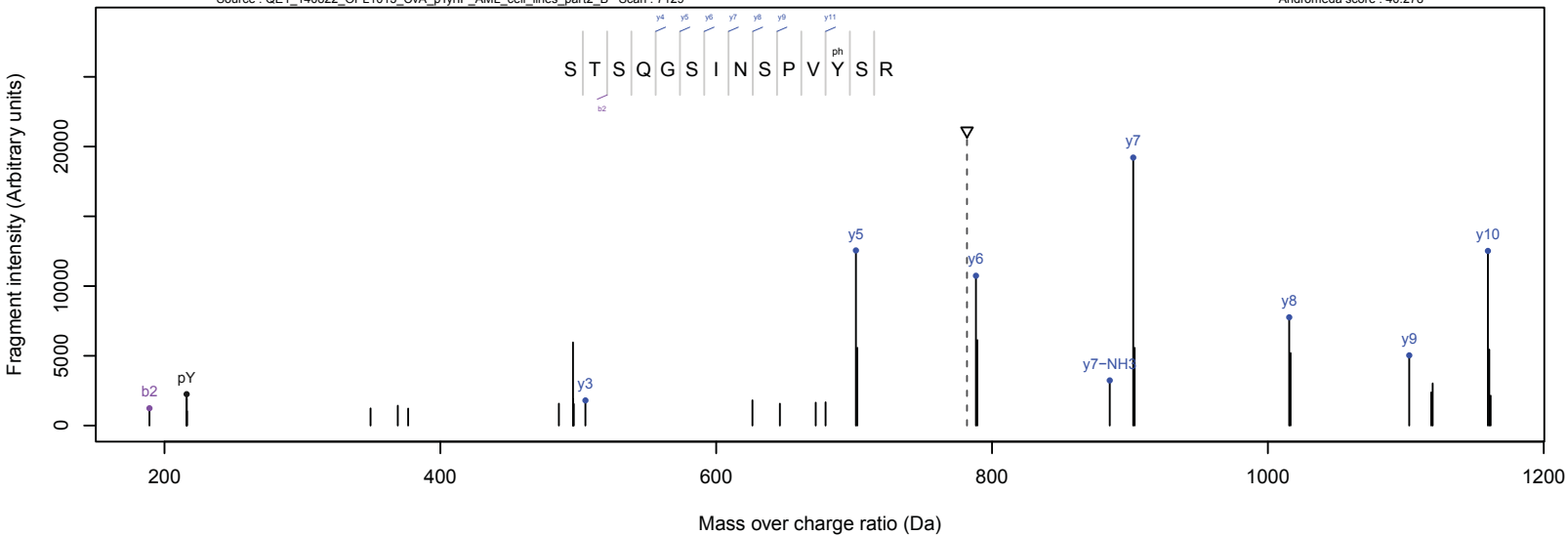
Andromeda score : 40.278



## STSQGSINSPV(ph)SR-2+

Source : QE1\_140822\_OPL1013\_CvA\_pTyrIP\_AML\_cell\_lines\_part2\_B Scan : 7129

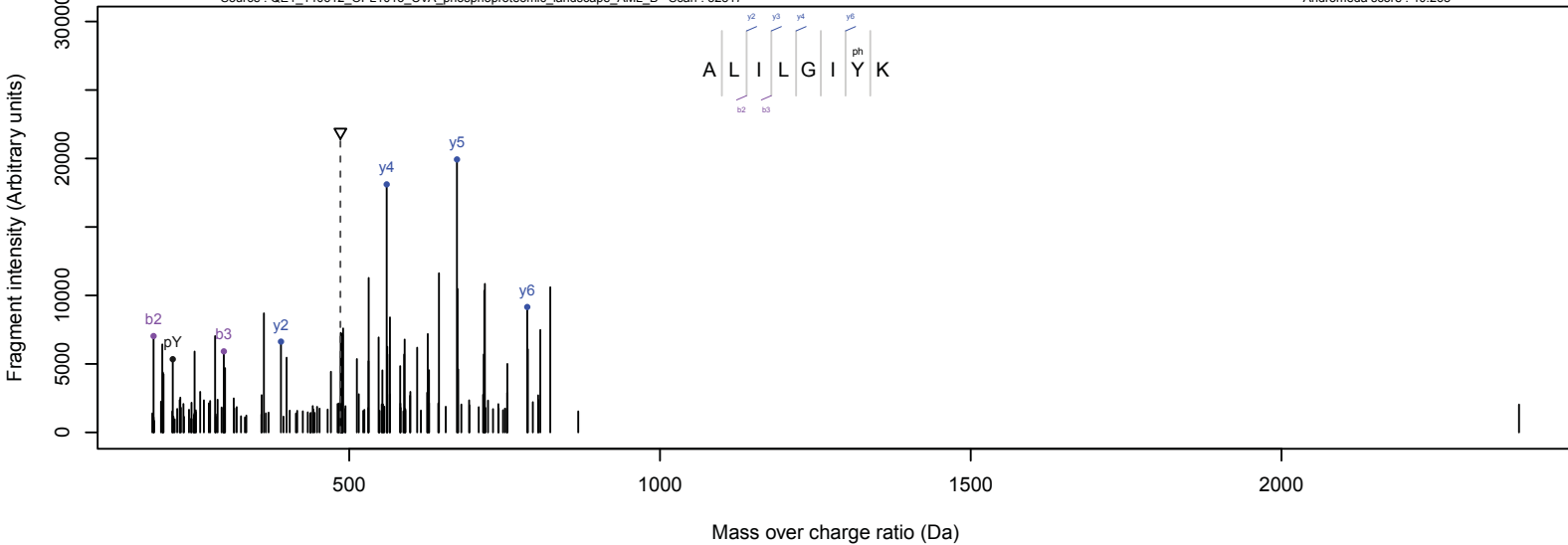
Andromeda score : 40.278



## ALILGIY(ph)K-2+

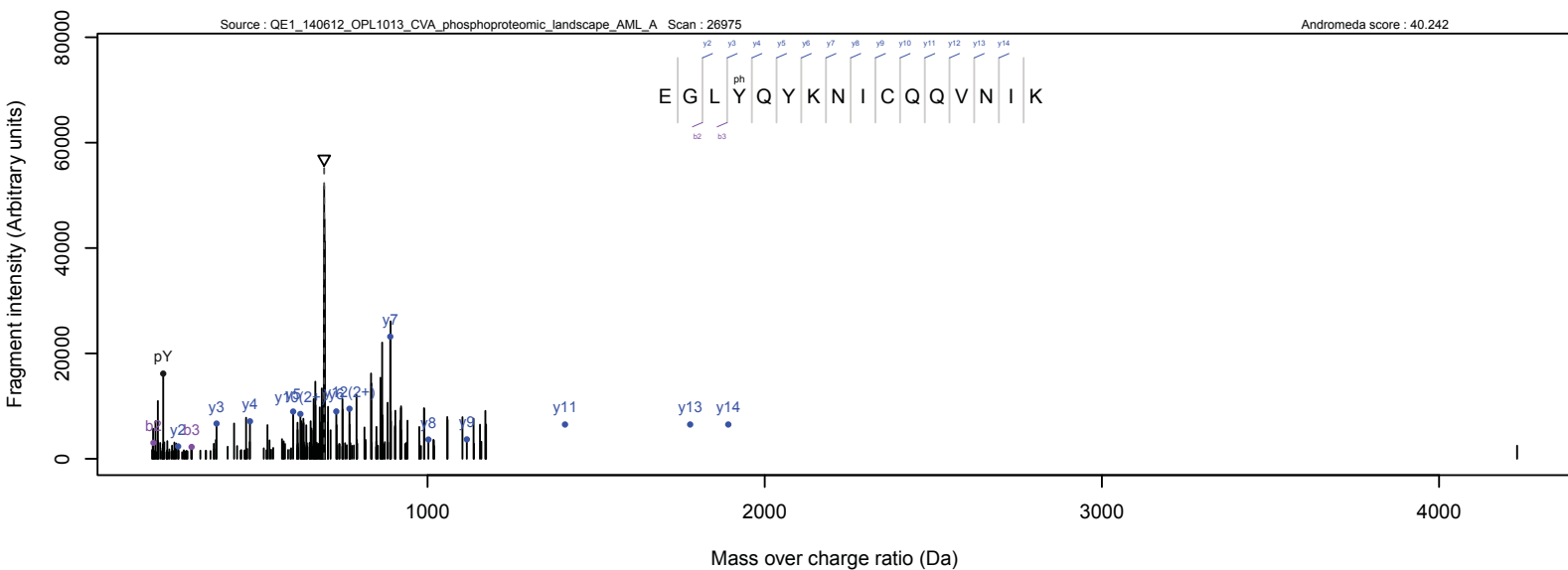
Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 32517

Andromeda score : 40.268

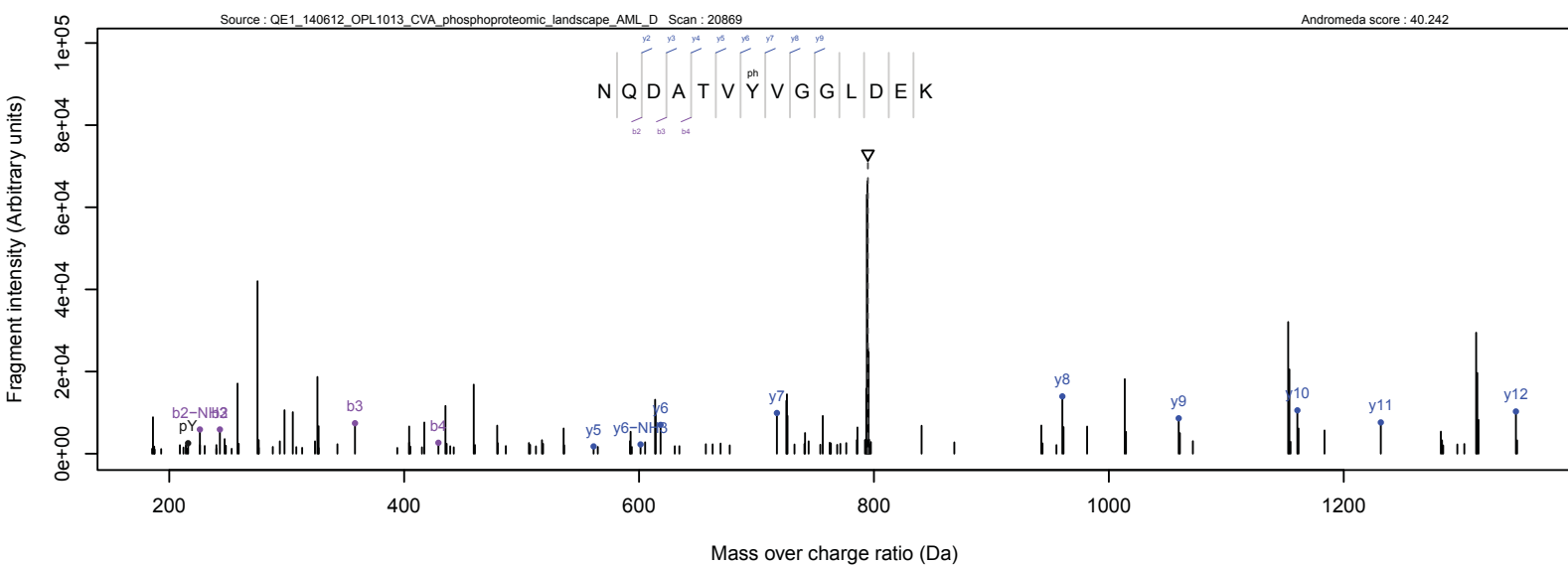




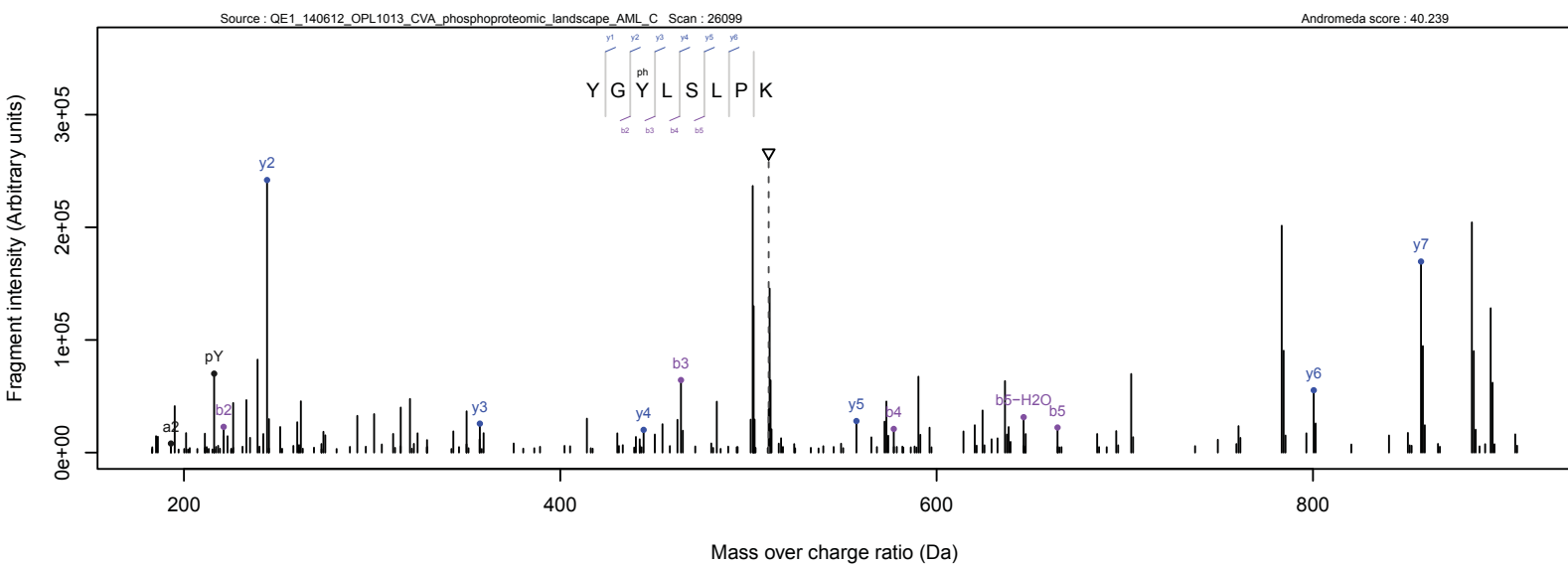
**EGLY(ph)QYKNICQQVNIK-3+**



**NQDATVY(ph)VGGLEDEK-2+**



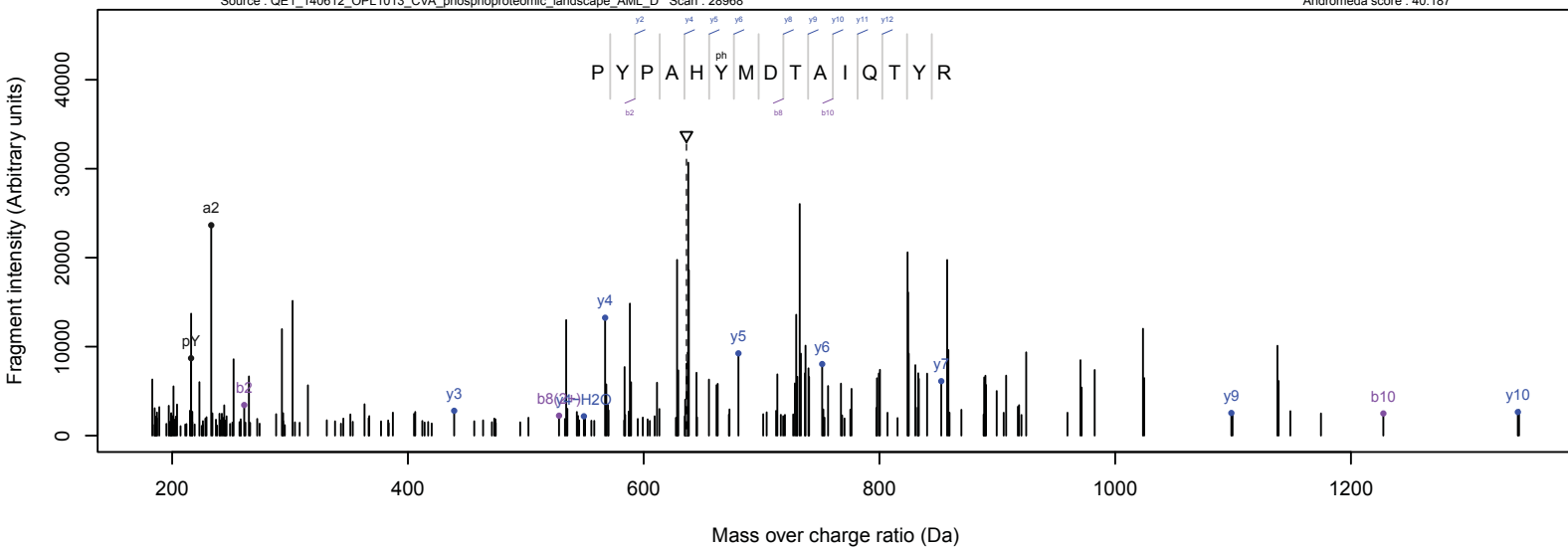
**YGY(ph)LSLPK-2+**



## PYPAHY(ph)MDTAIQTYR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_D Scan : 28968

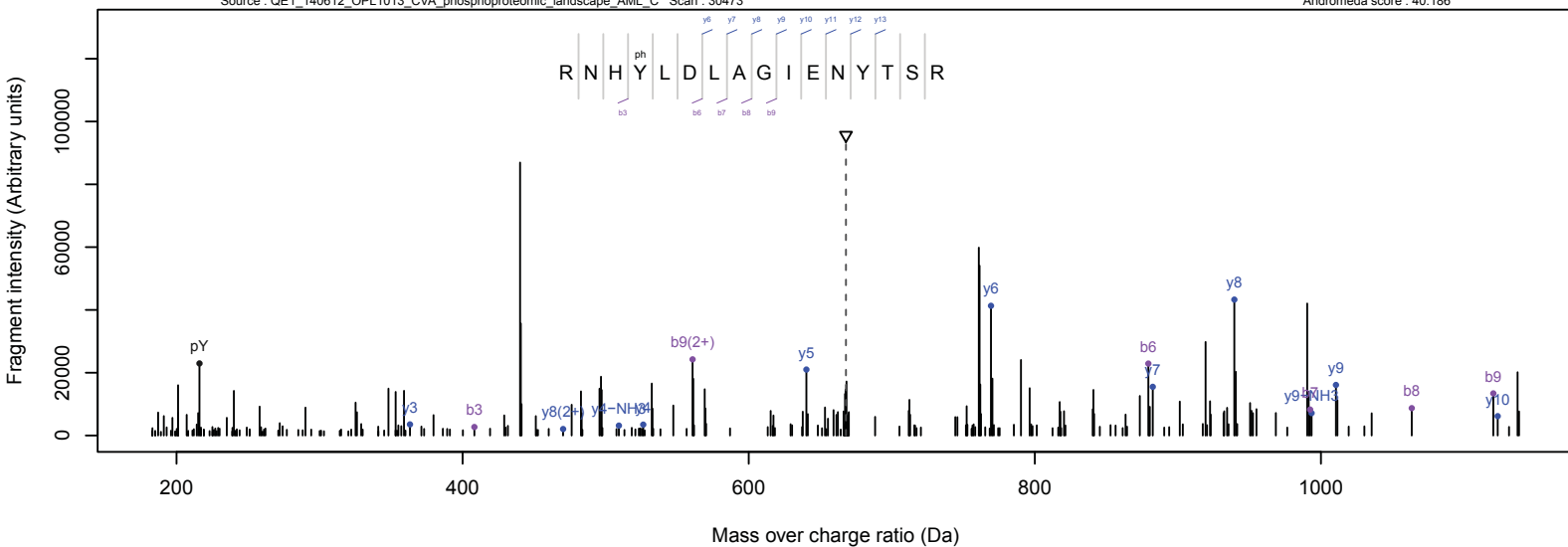
Andromeda score : 40.187



## RNHY(ph)LDLAGIENYTSR-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_C Scan : 30473

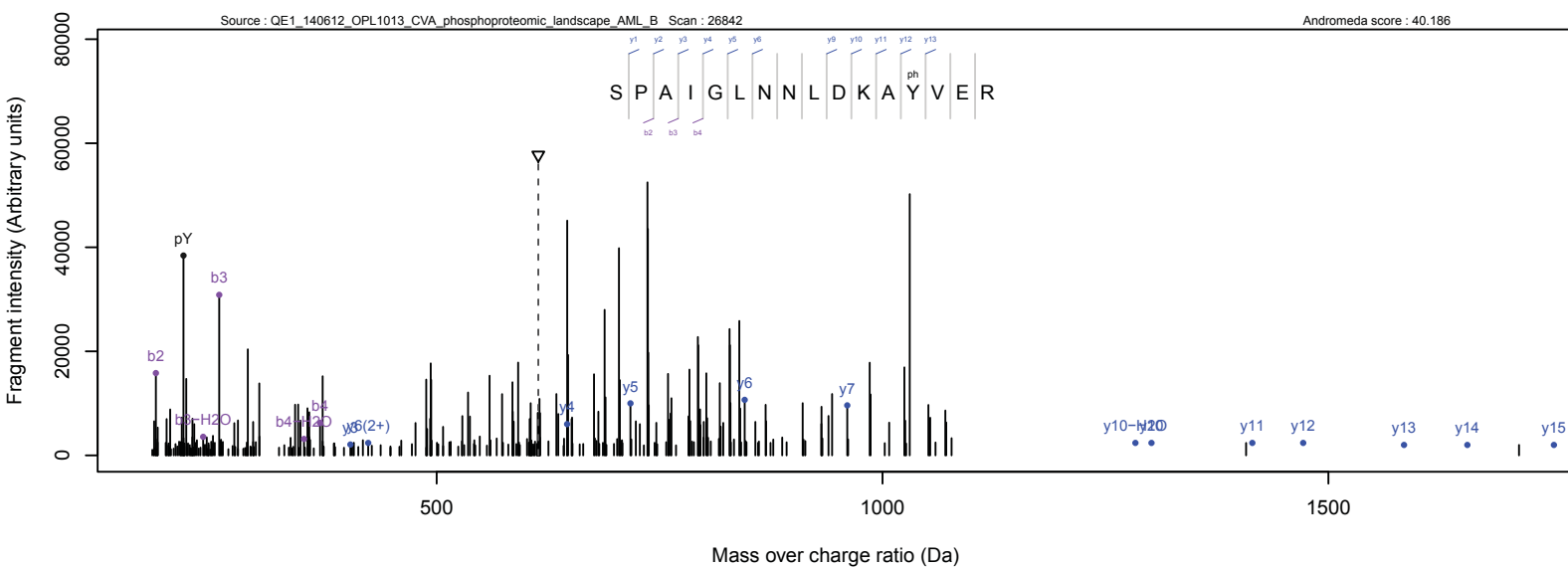
Andromeda score : 40.186



## SPAIGLNNLDKAY(ph)VER-3+

Source : QE1\_140612\_OPL1013\_CVA\_phosphoproteomic\_landscape\_AML\_B Scan : 26842

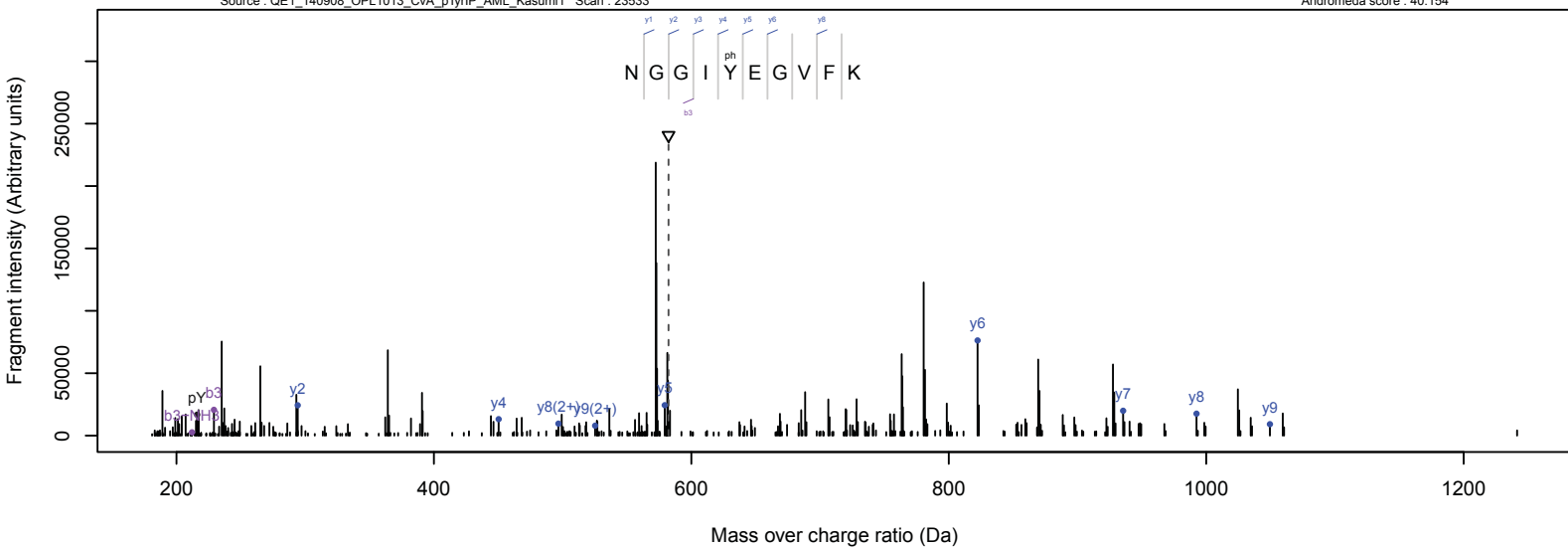
Andromeda score : 40.186



## NGGIY(ph)EGVFK-2+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi1 Scan: 23533

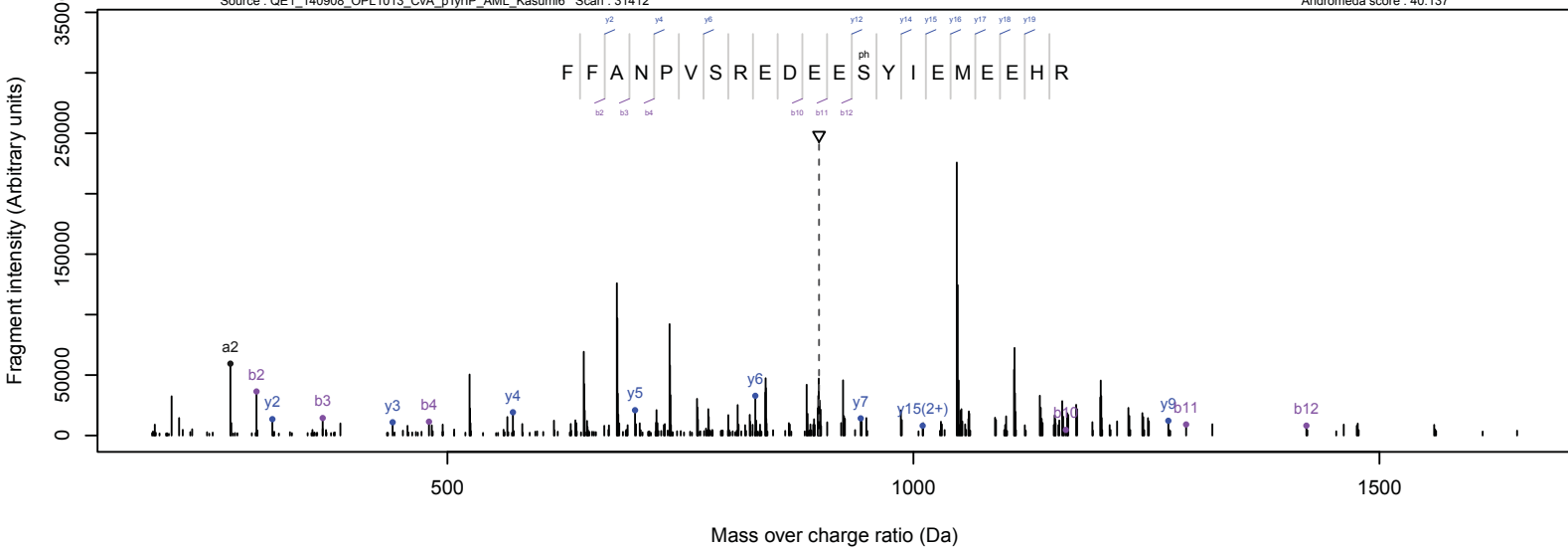
Andromeda score: 40.154



## FFANPVSREDEES(ph)YIEMEEHR-3+

Source: QE1\_140908\_OPL1013\_CvA\_pTyrIP\_AML\_Kasumi6 Scan: 31412

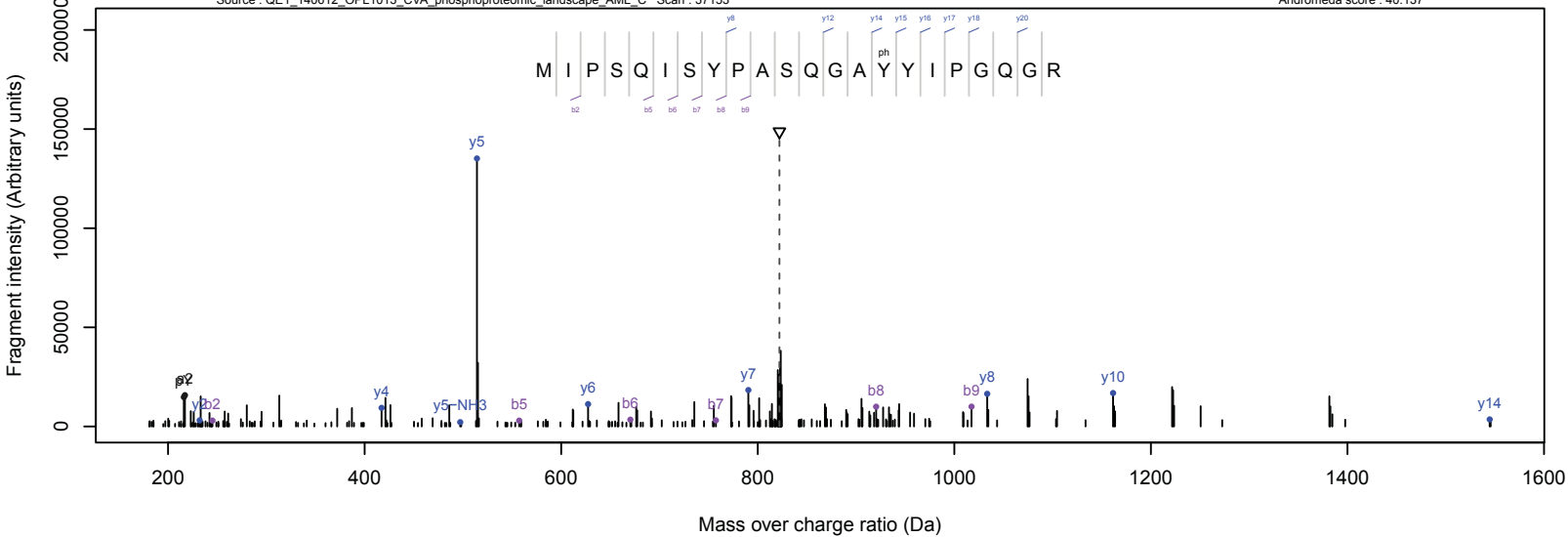
Andromeda score: 40.137



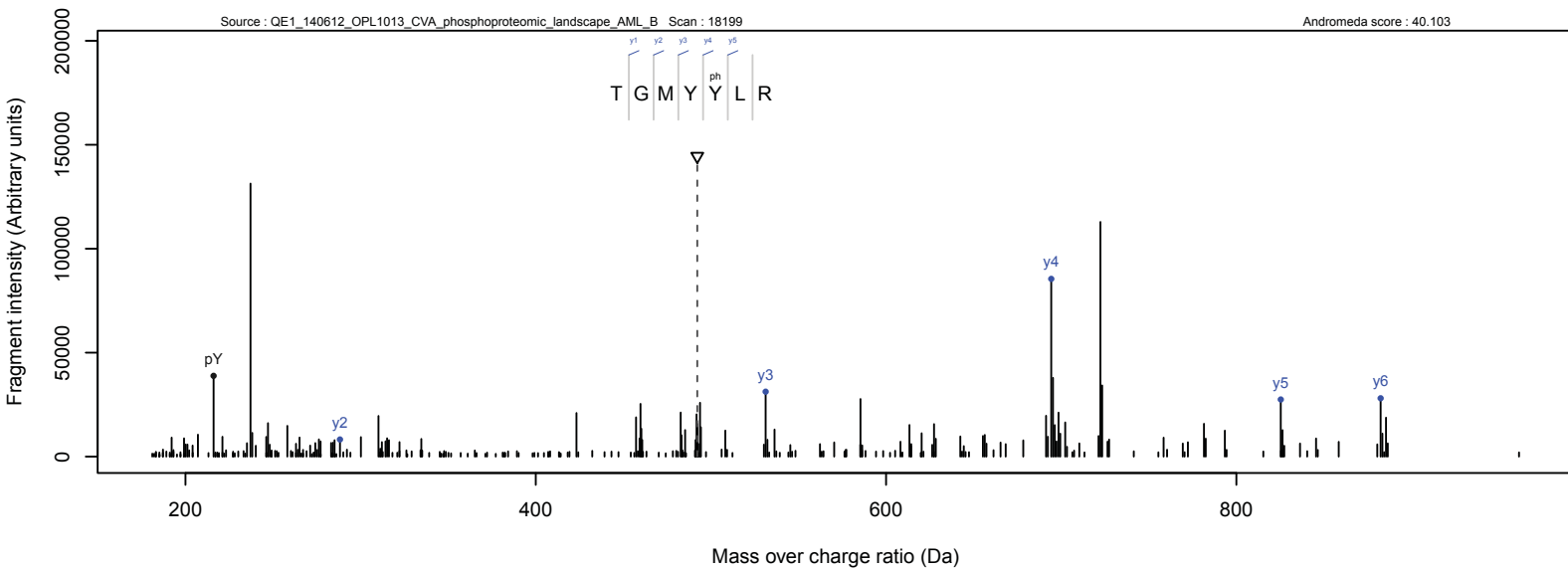
## MIPSQISYPASQGA(ph)YIPGQR-3+

Source: QE1\_140612\_OPL1013\_CvA\_phosphoproteomic\_landscape\_AML\_C Scan: 37153

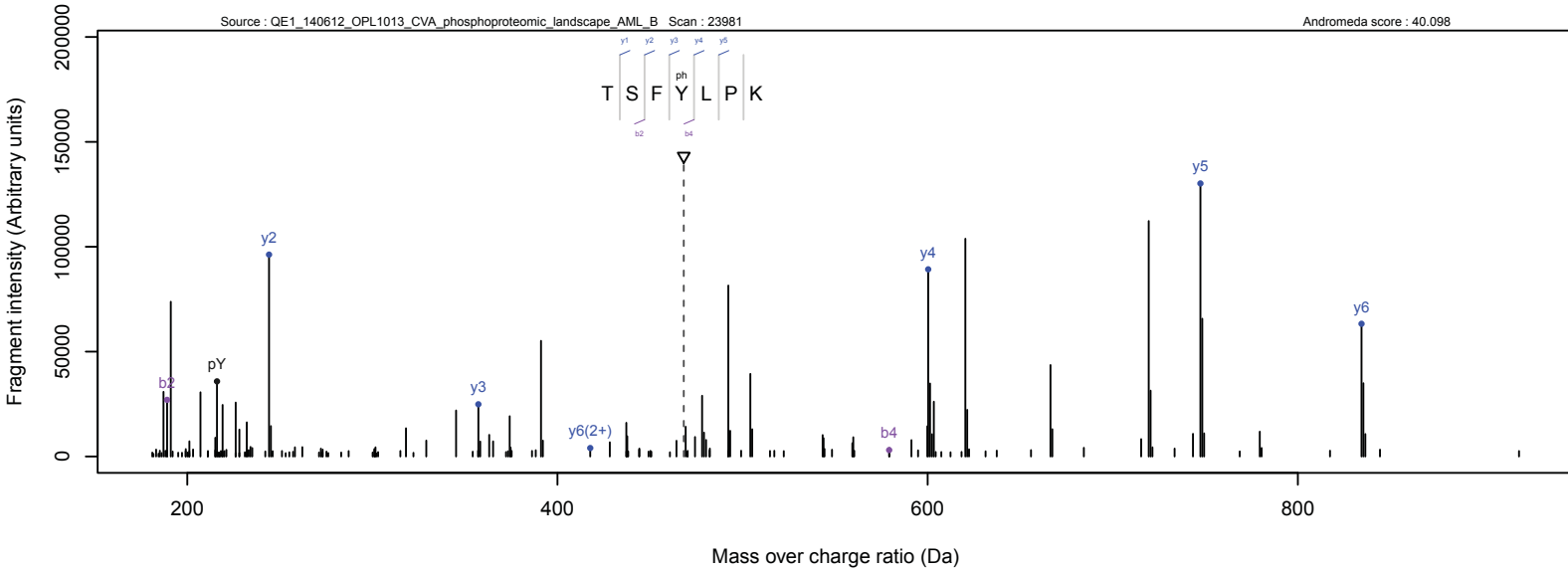
Andromeda score: 40.137



## TGMYY(ph)LR-2+



## TSFY(ph)LPK-2+



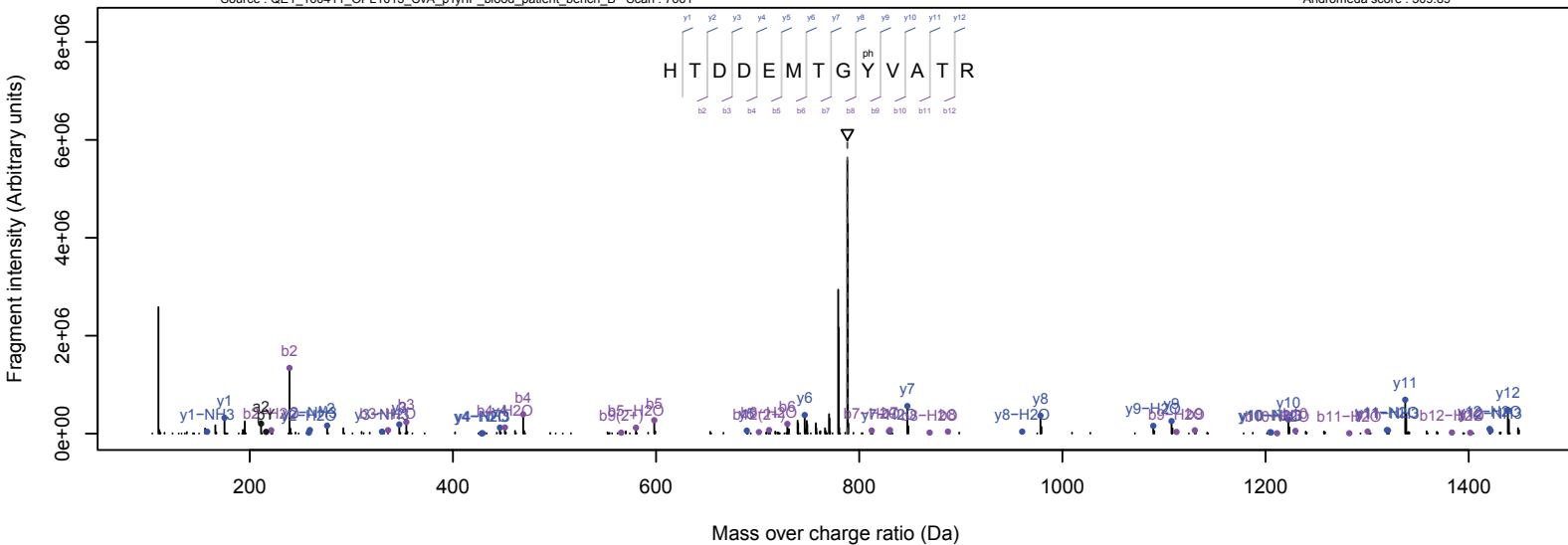
# Supplemental Figure 9

Annotated MS/MS spectra of phosphopeptides  
identified in clinical AML patient samples

## HTDDEMTGY(ph)VATR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 7661

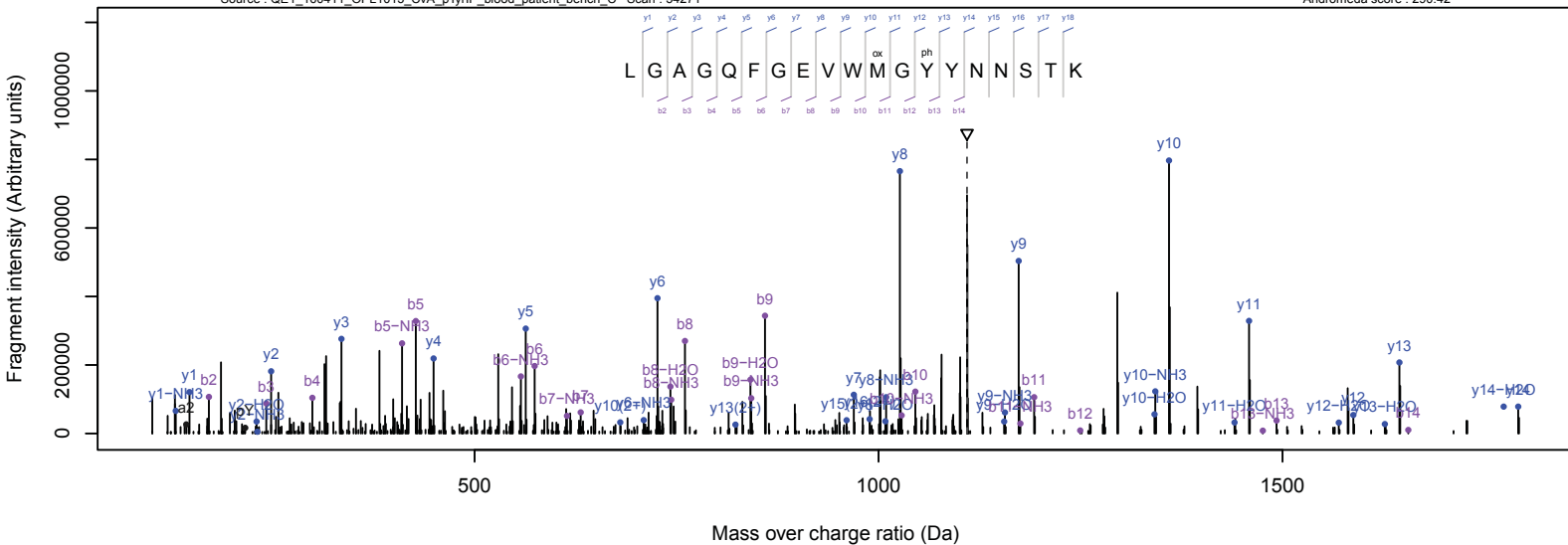
Andromeda score: 309.89



## LGAGQFGEVWM(ox)GY(ph)YNNSTK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 34271

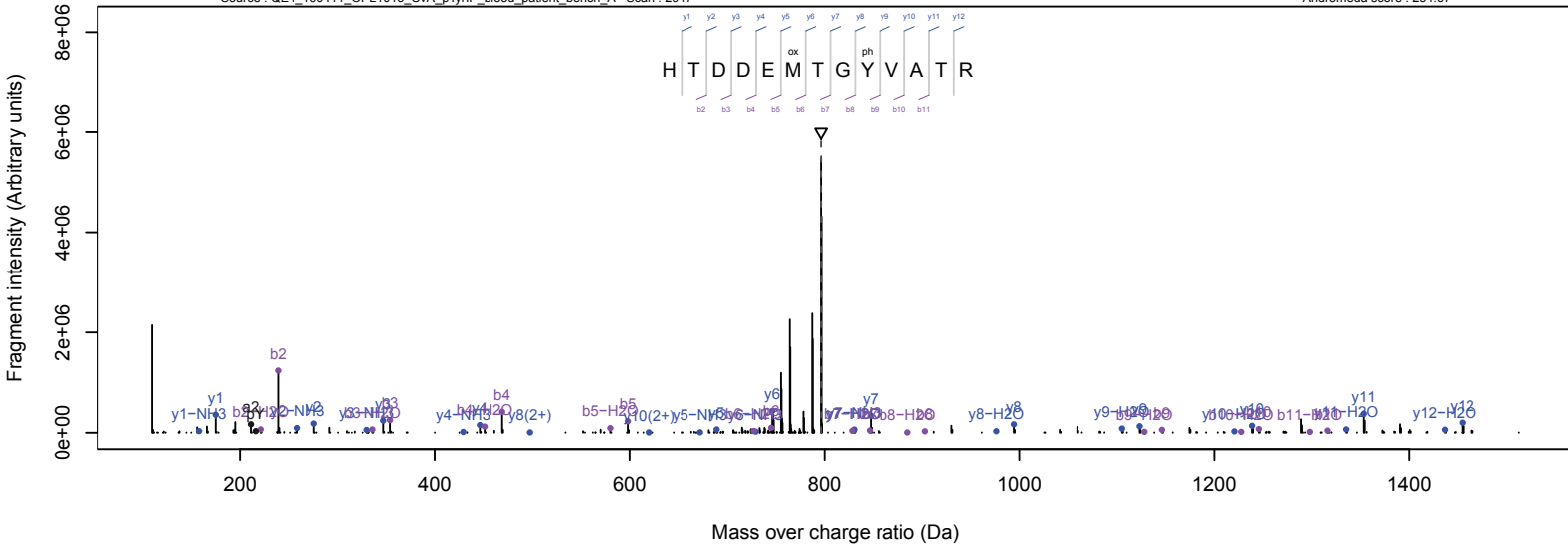
Andromeda score: 290.42



## HTDDEM(ox)TGY(ph)VATR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 2917

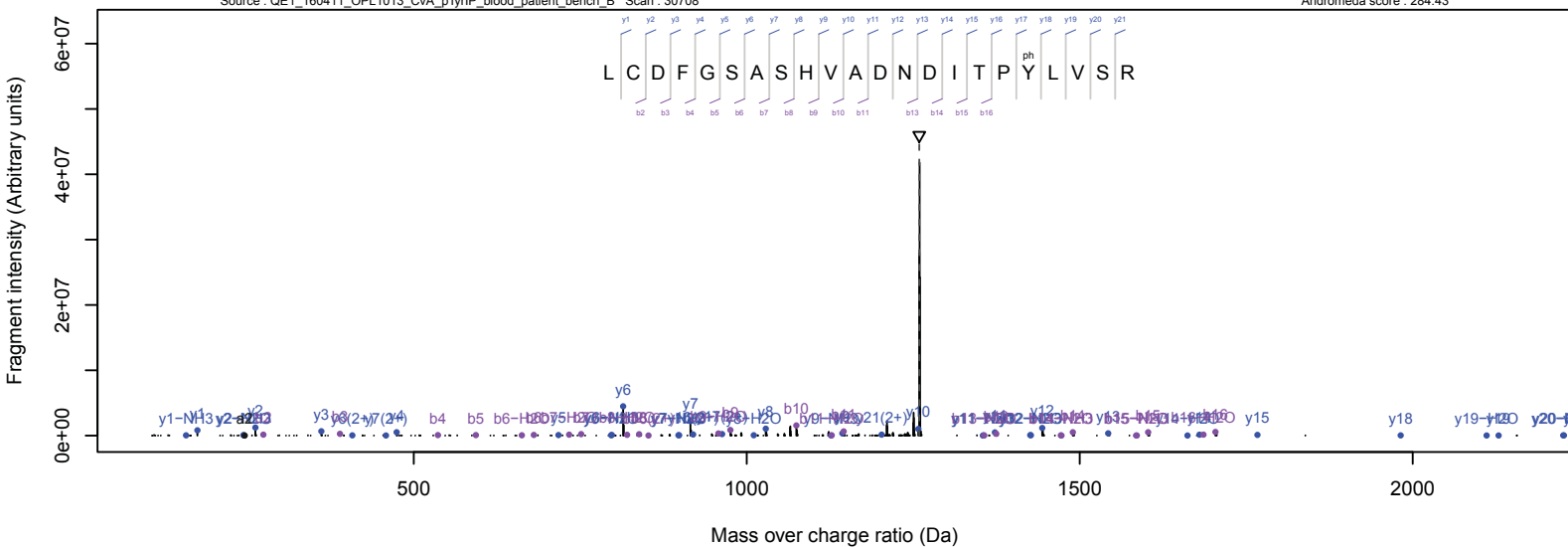
Andromeda score: 284.67



**LCDFGSASHVADNDITPYLVSR-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 30708

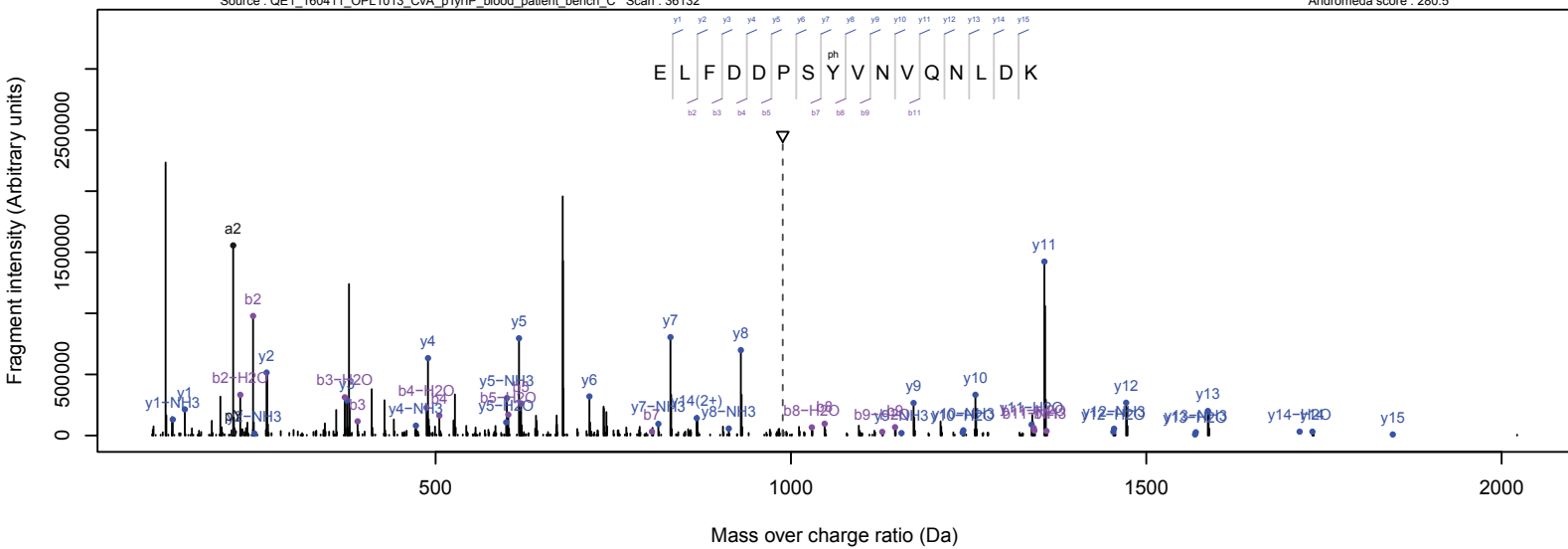
Andromeda score: 284.43



**ELFDDPSYVNVQLDK-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 36132

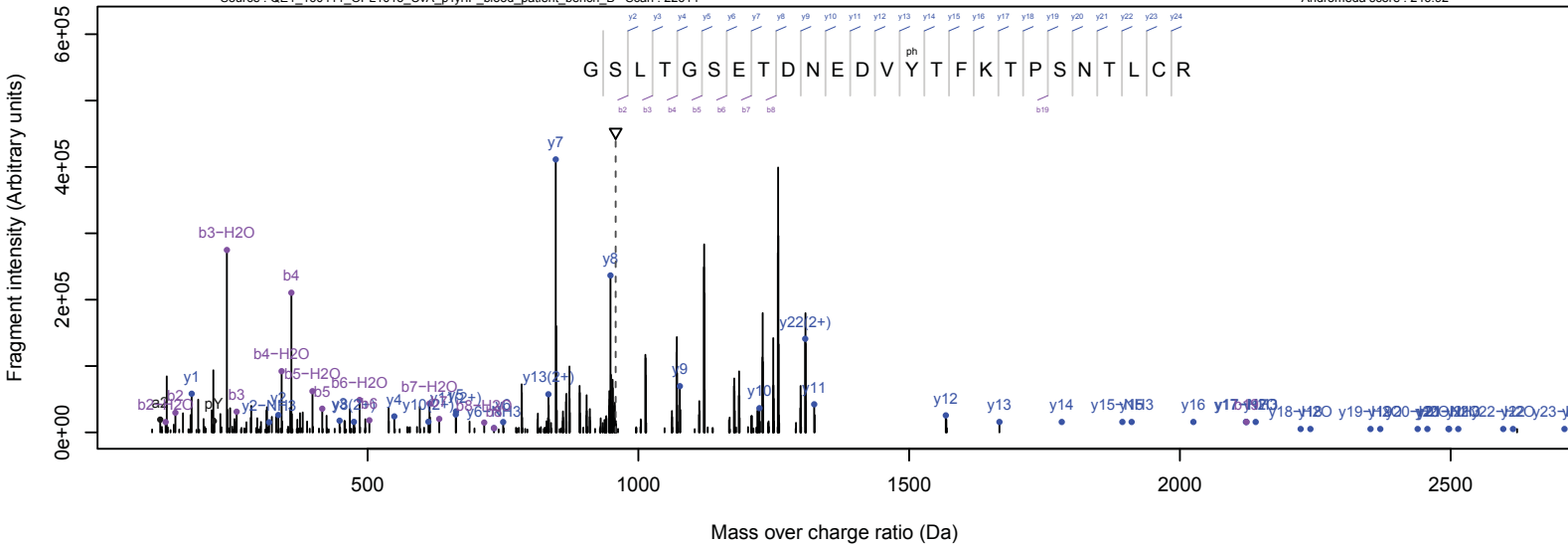
Andromeda score: 280.5



**GSLTGSETDNEDVYTFKTPSNTLCR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 22614

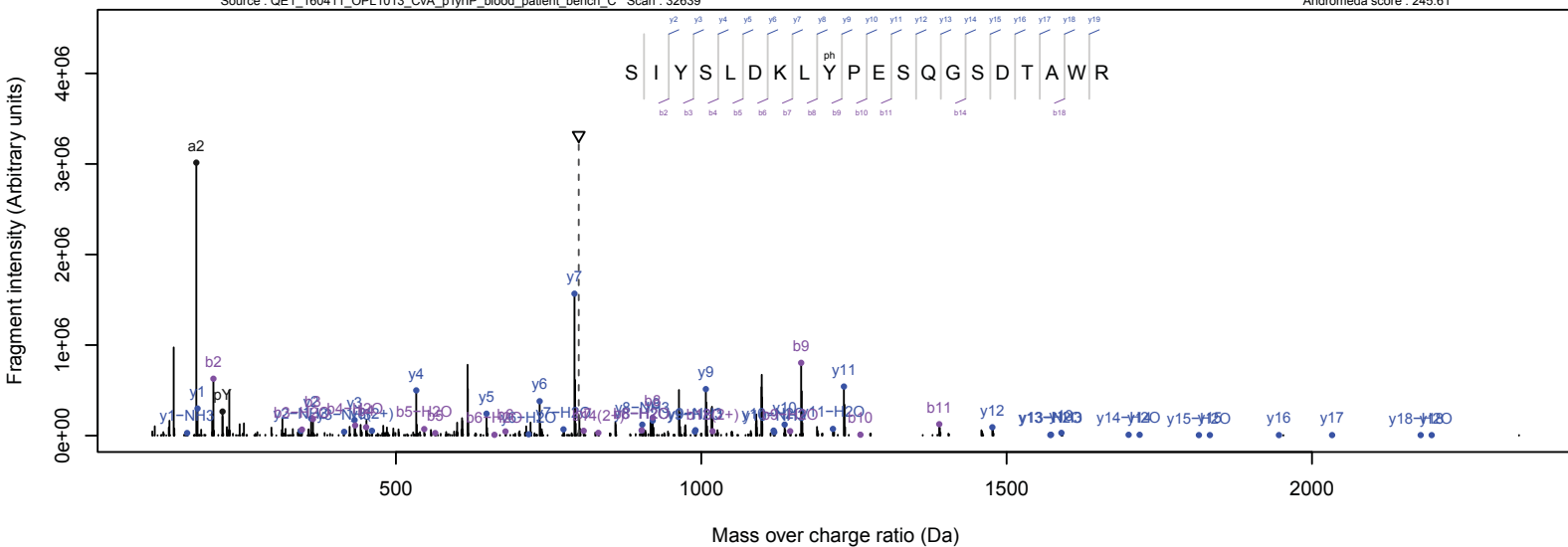
Andromeda score: 246.92



SIYSLDKLY(ph)PESQGSDTAWR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 32639

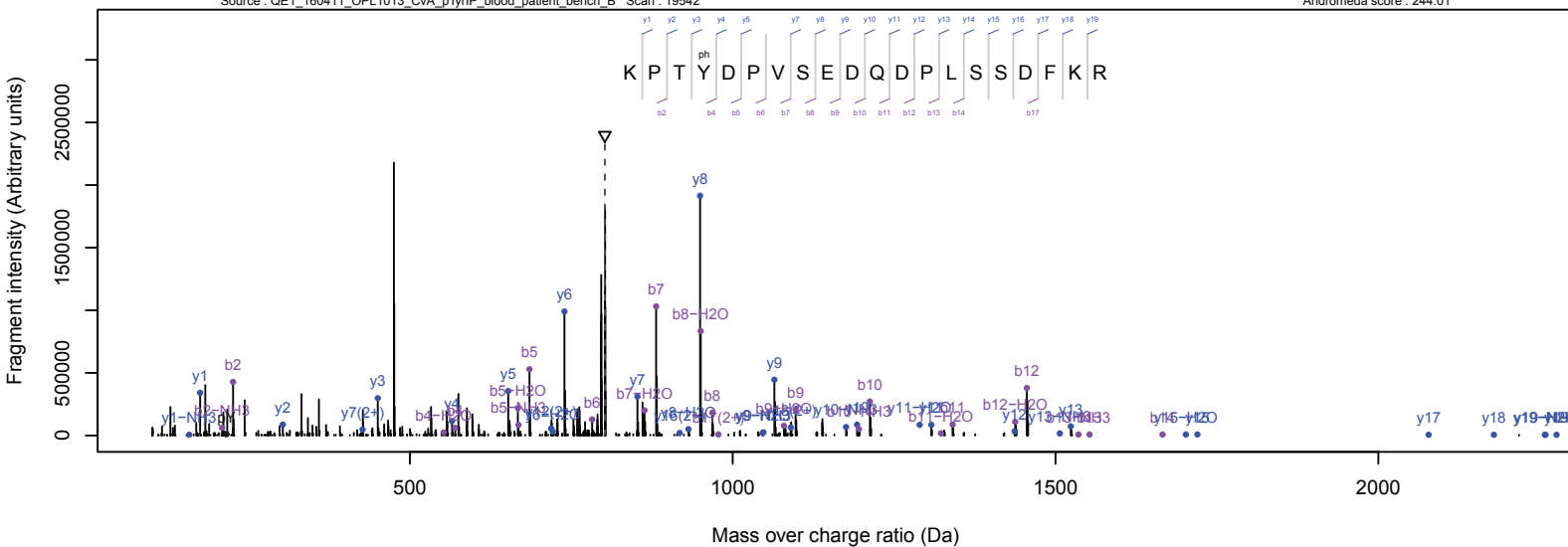
Andromeda score : 245.61



KPTY(ph)DPVSEDQDPLSSDFKR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 19542

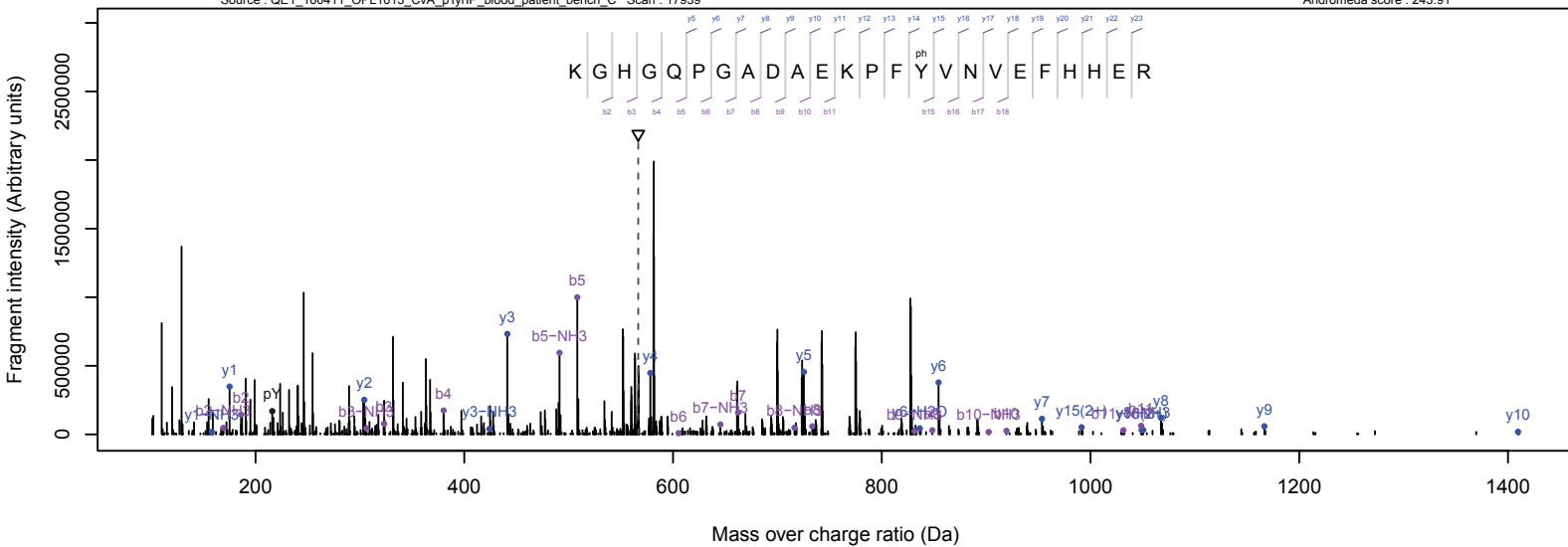
Andromeda score : 244.01



KGHGQPGADAEKPHYVNVFHHHER-5+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 17939

Andromeda score : 243.91

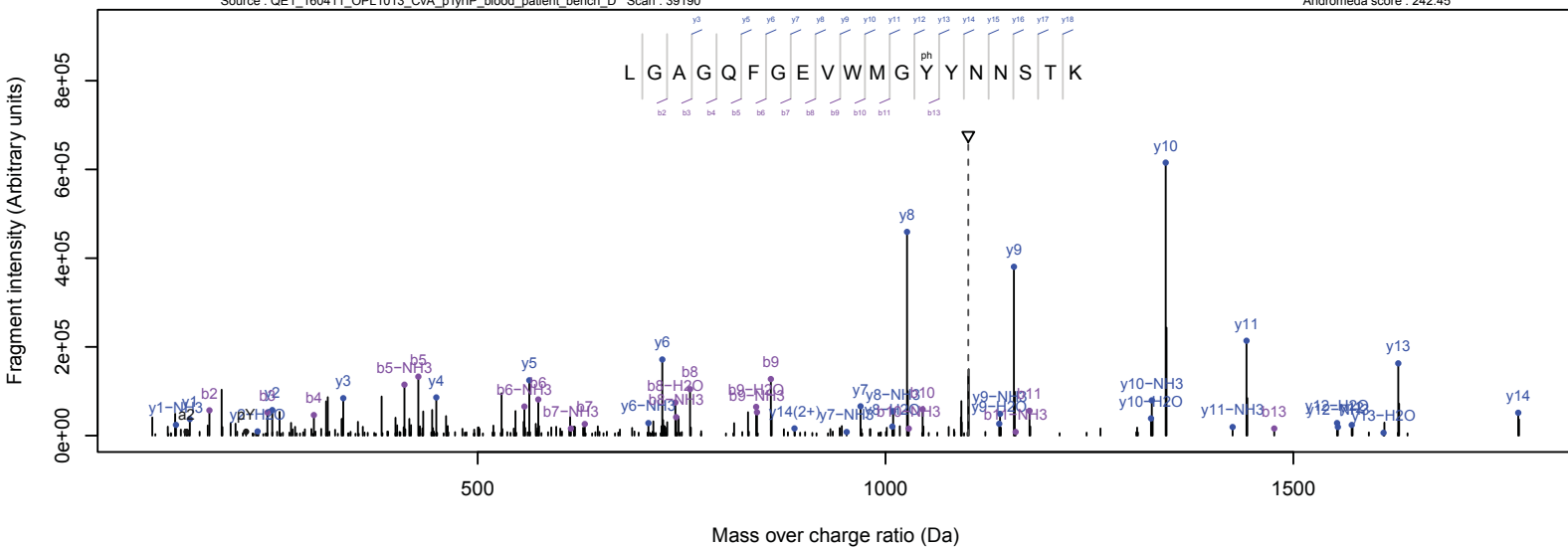




LGAGQFGEVWMGY(ph)YNNSTK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 39190

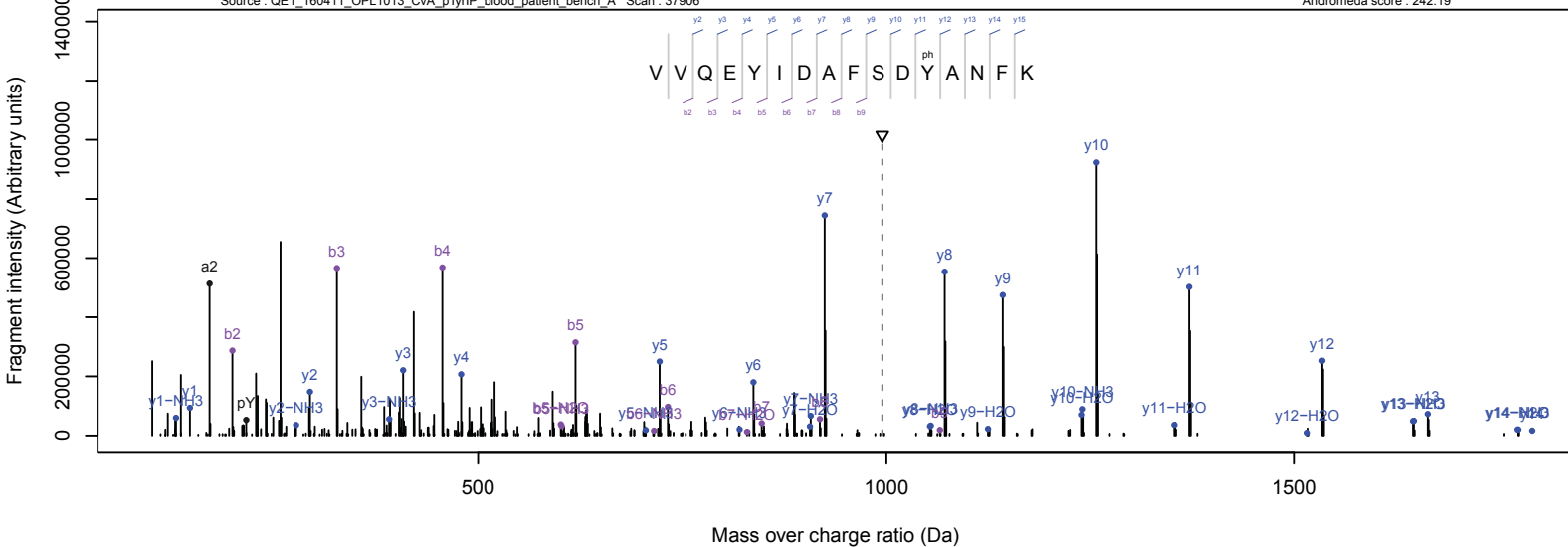
Andromeda score: 242.45



VVQEYIDAFSDY(ph)ANFK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 37906

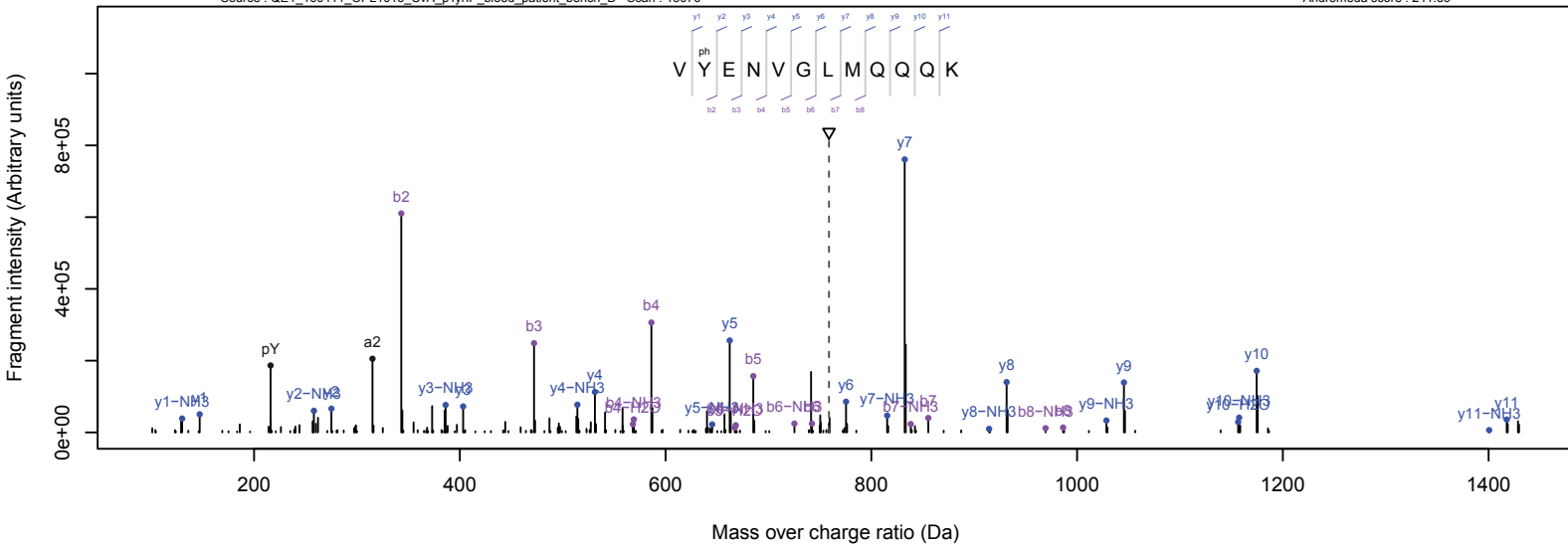
Andromeda score: 242.19



VY(ph)ENVGLMQQK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 15076

Andromeda score: 241.39

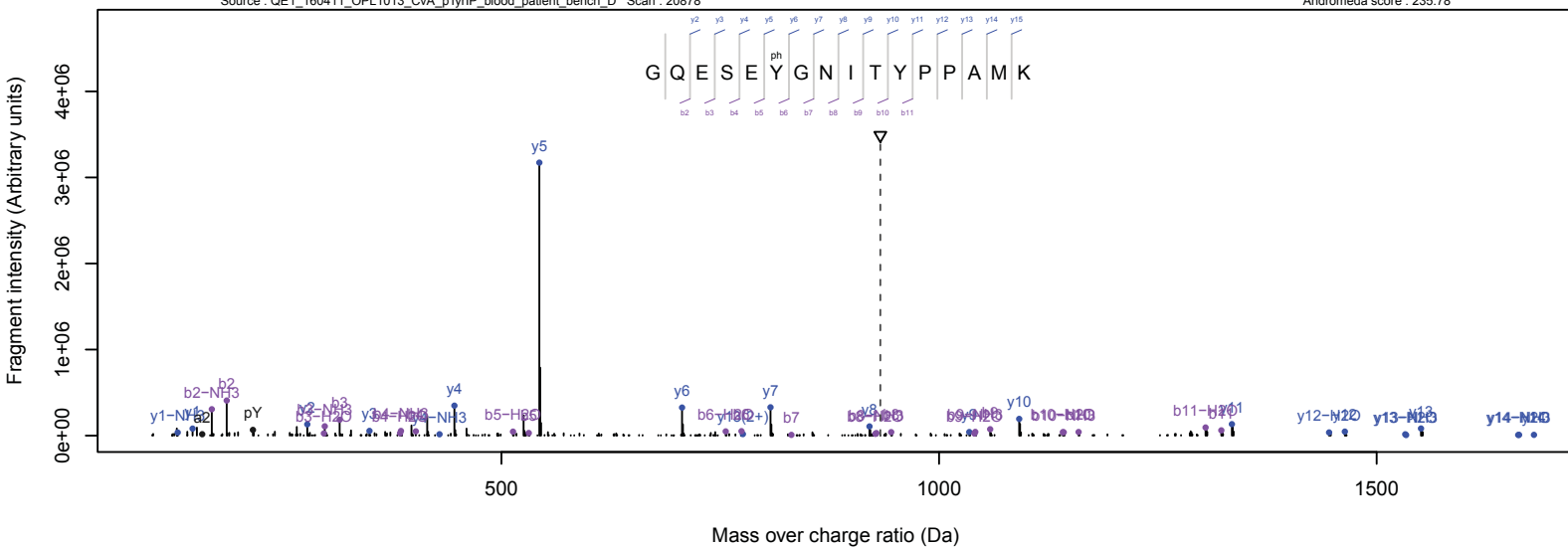




**GQESEY(ph)GNITYPPAMK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 20878

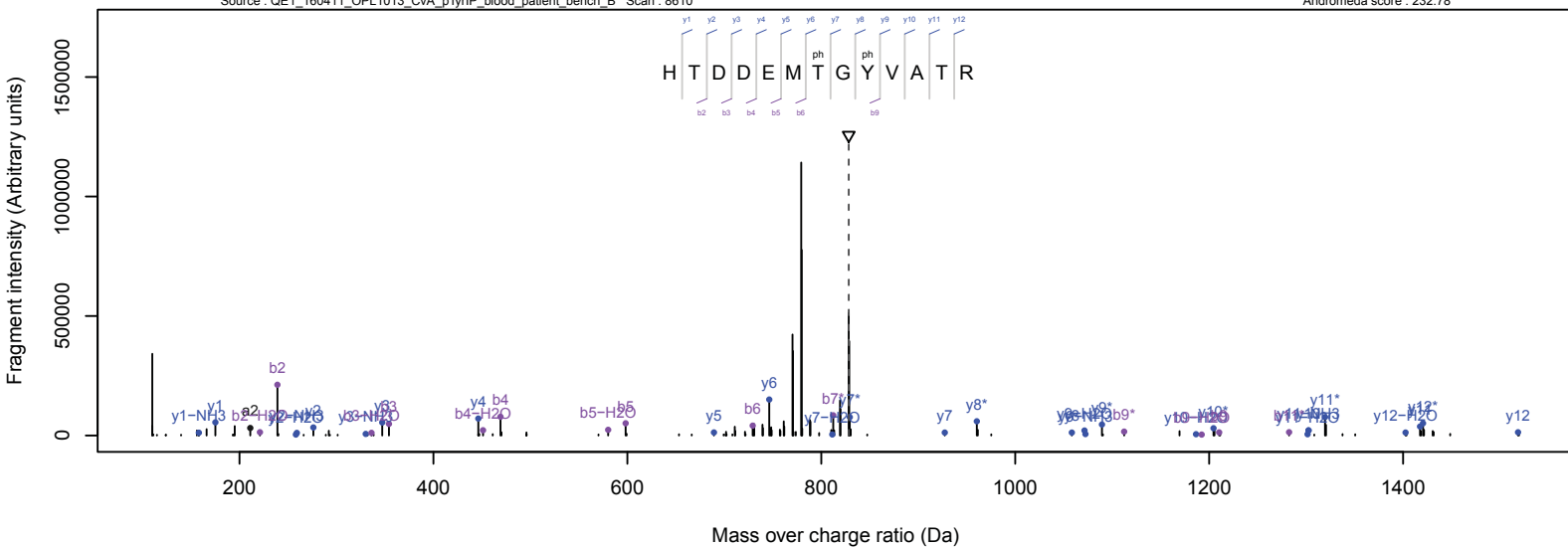
Andromeda score : 235.78



**HTDDEMT(ph)GYVATR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 8610

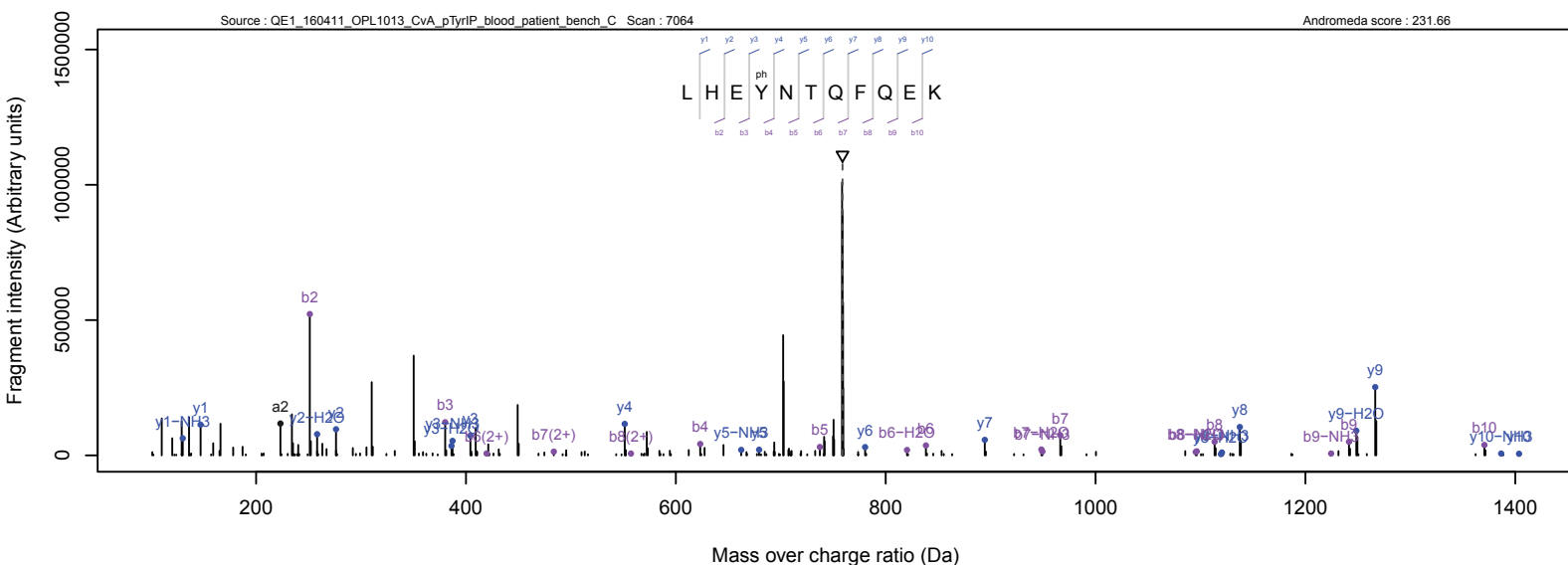
Andromeda score : 232.78



**LHEY(ph)NTQFQEK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 7064

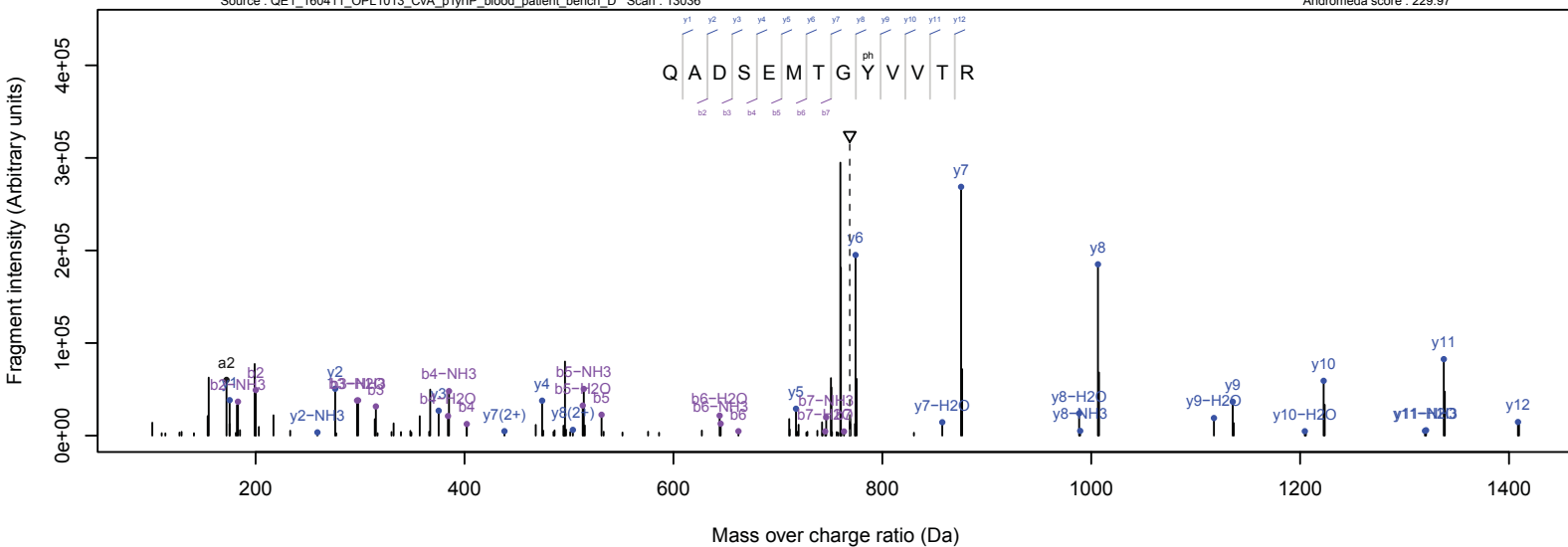
Andromeda score : 231.66



**QADSEMTGY(ph)VVTR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 13036

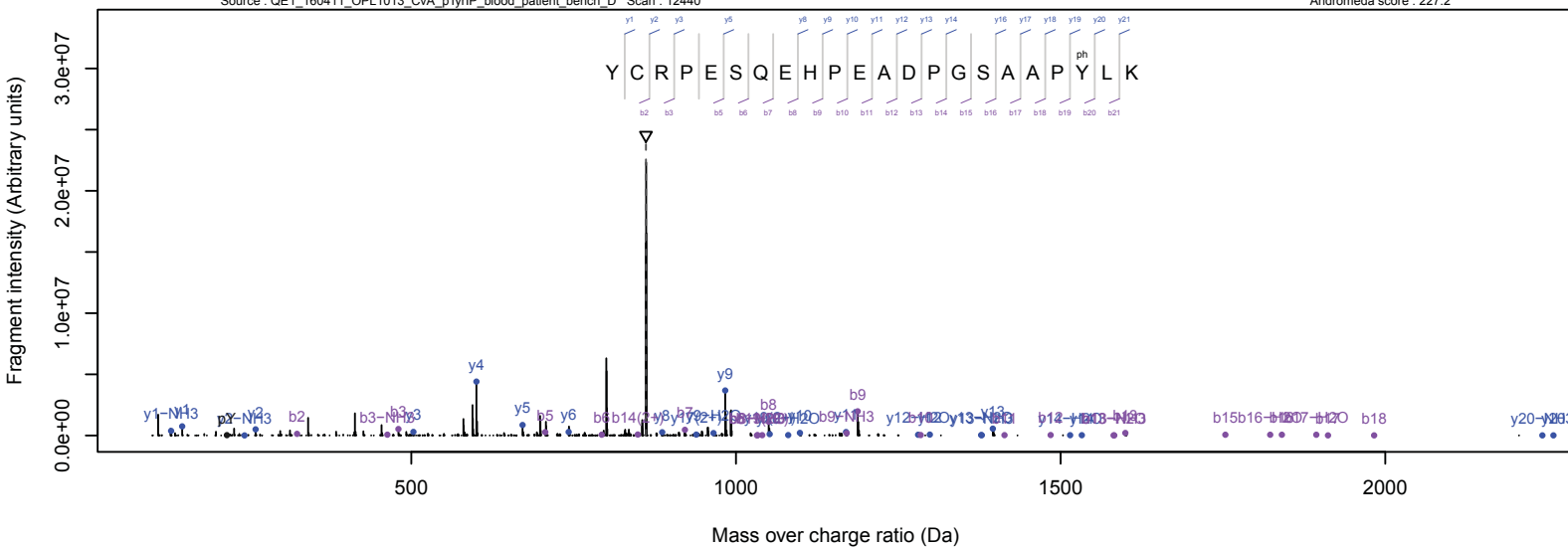
Andromeda score : 229.97



**YCRPESQEHPEADPGSAAPY(ph)LK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12440

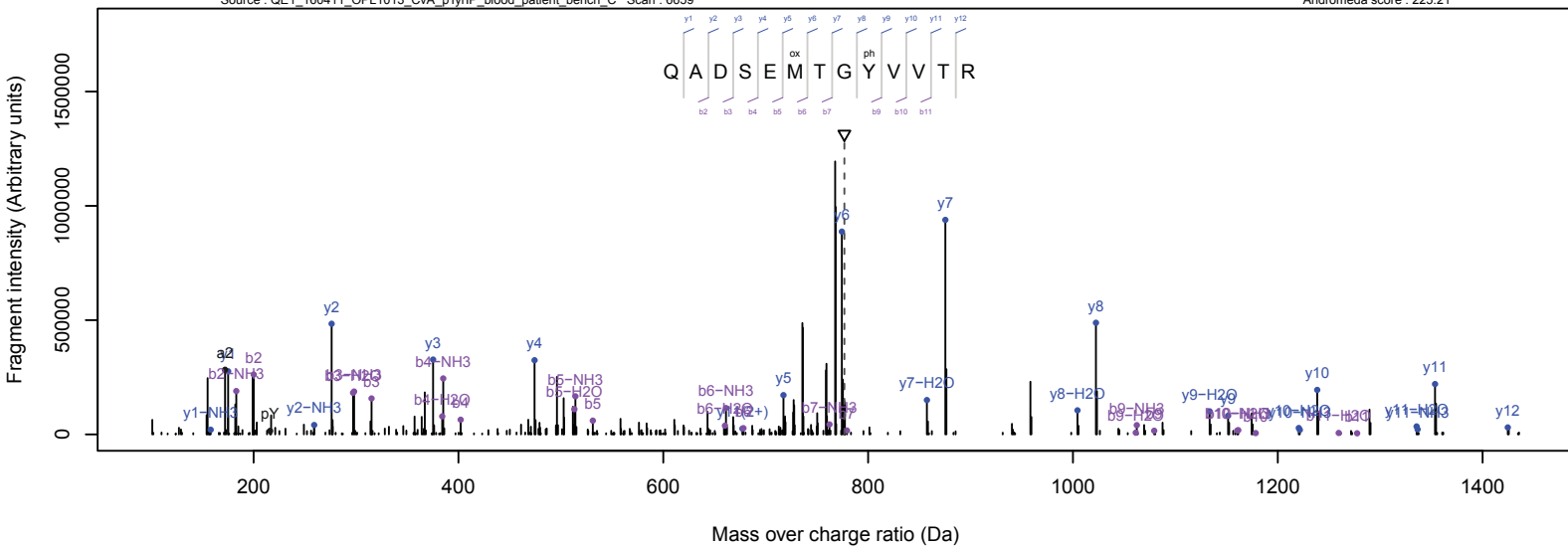
Andromeda score : 227.2

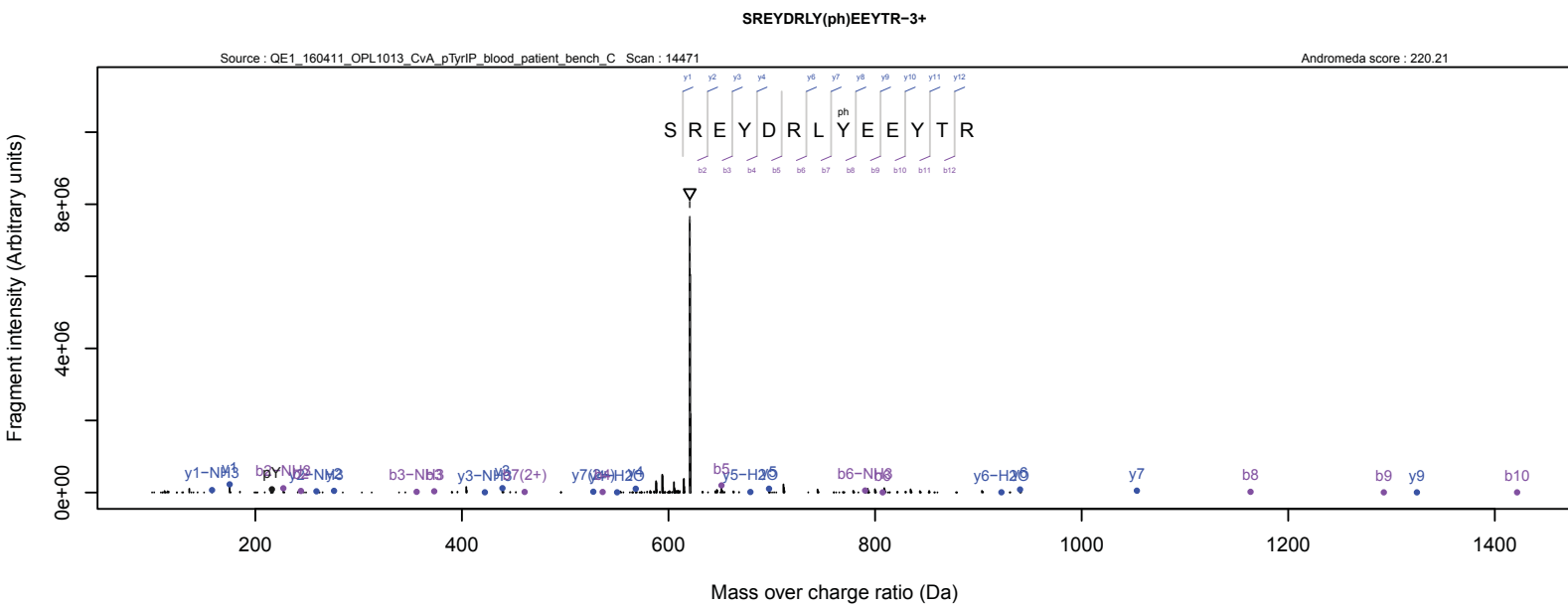
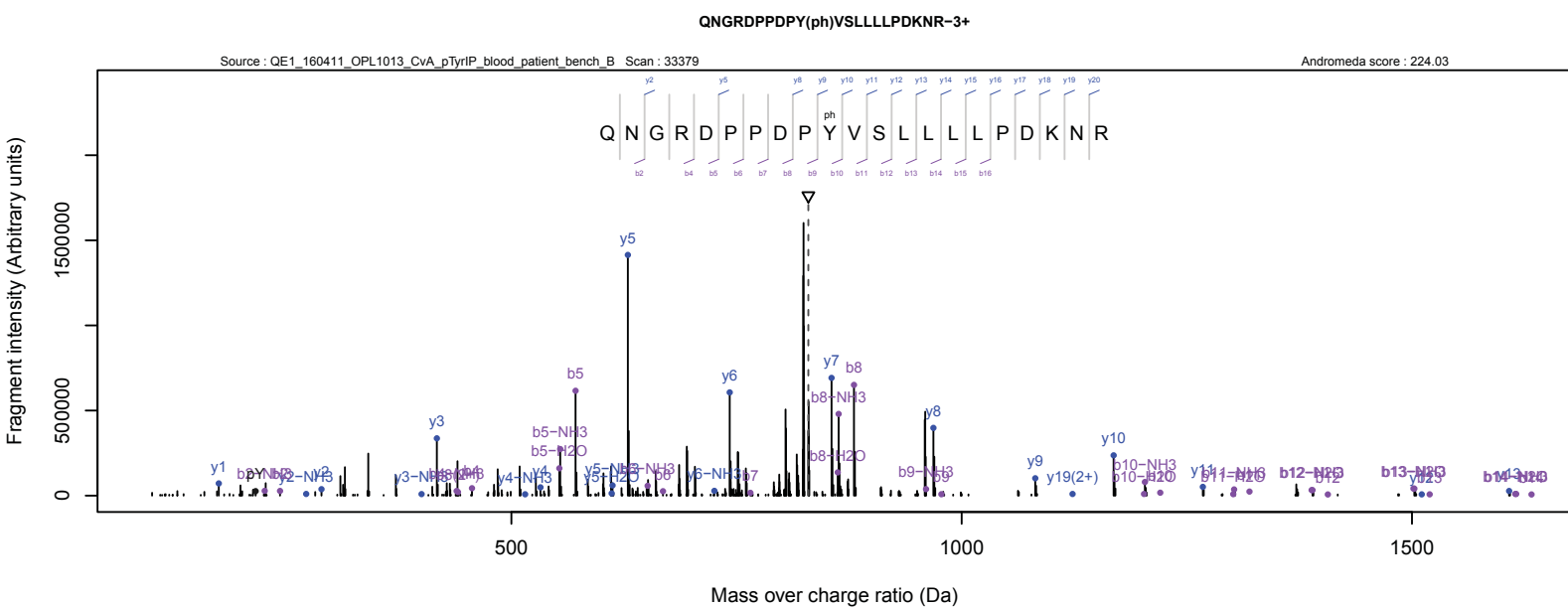
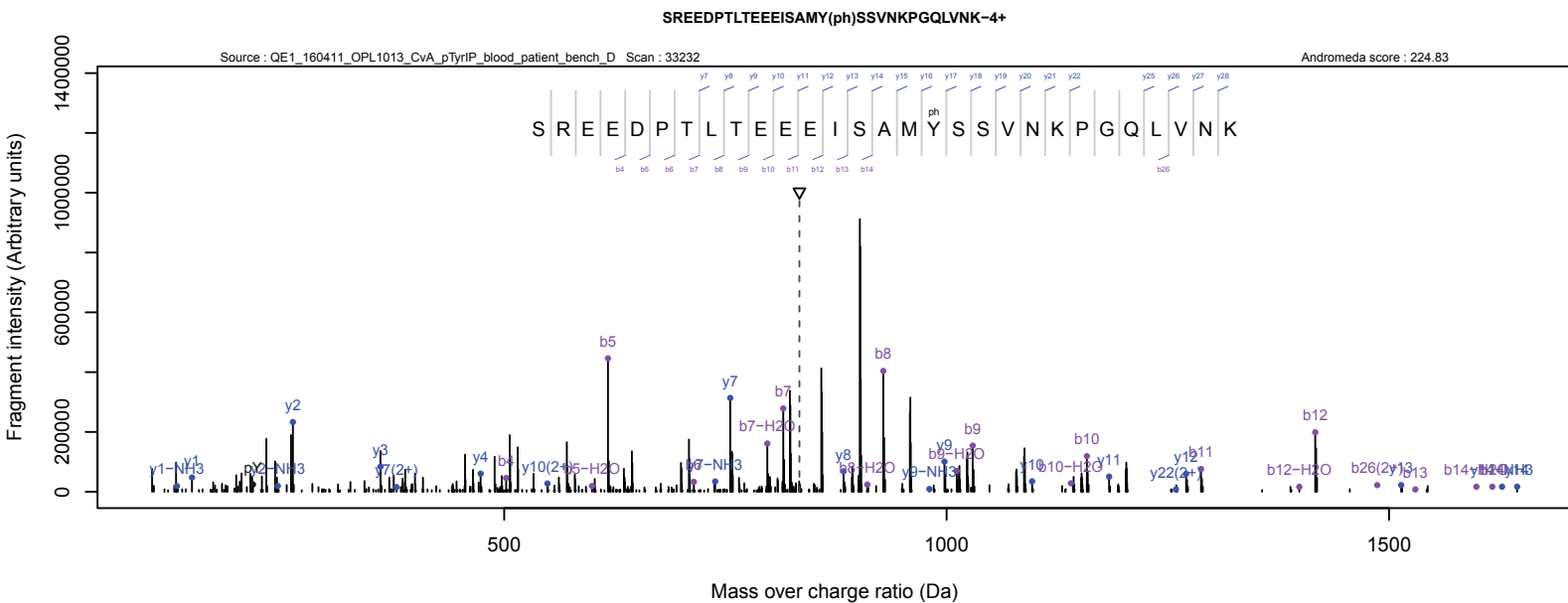


**QADSEM(ox)TGY(ph)VVTR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 6659

Andromeda score : 225.21



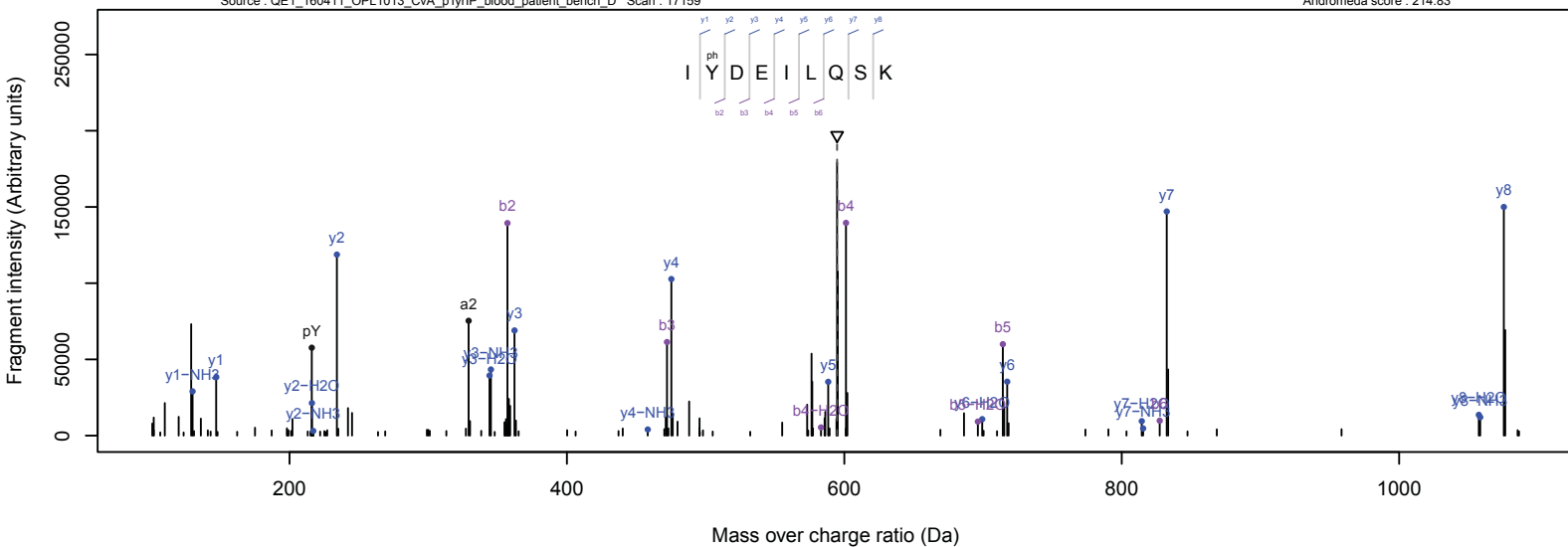




## IY(ph)DEILQSK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 17159

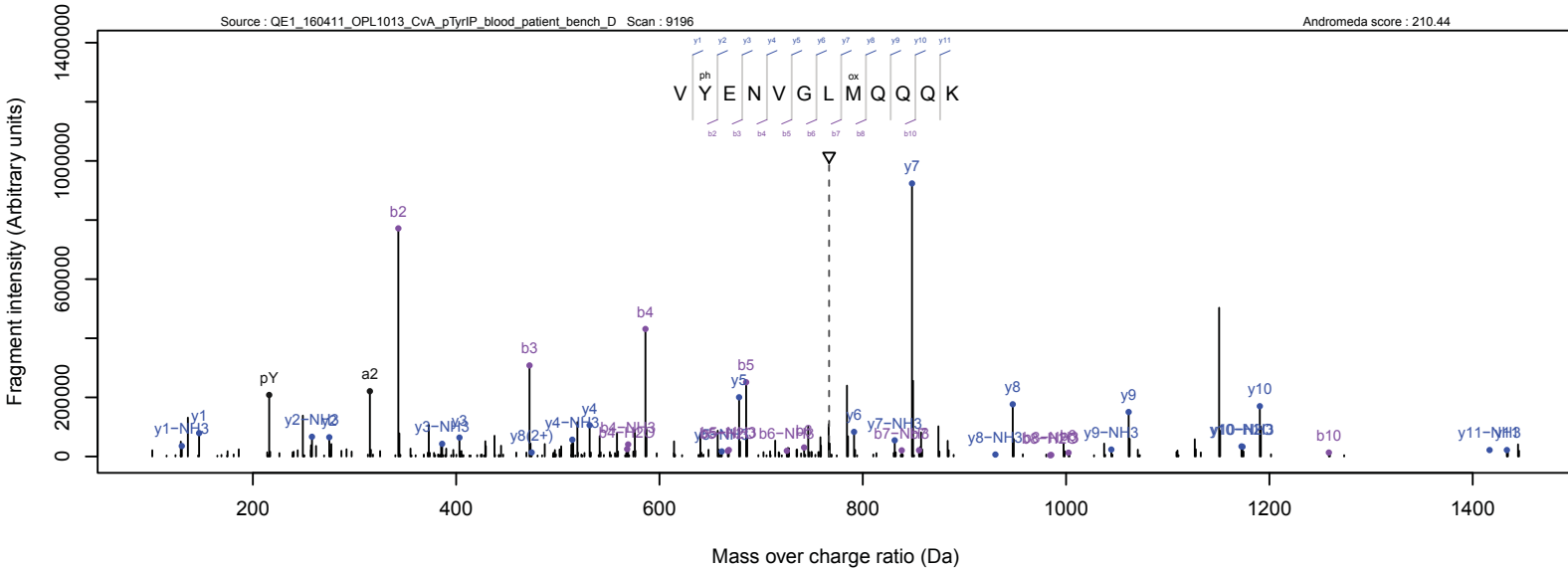
Andromeda score: 214.83



## VY(ph)ENVGLM(ox)QQQK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 9196

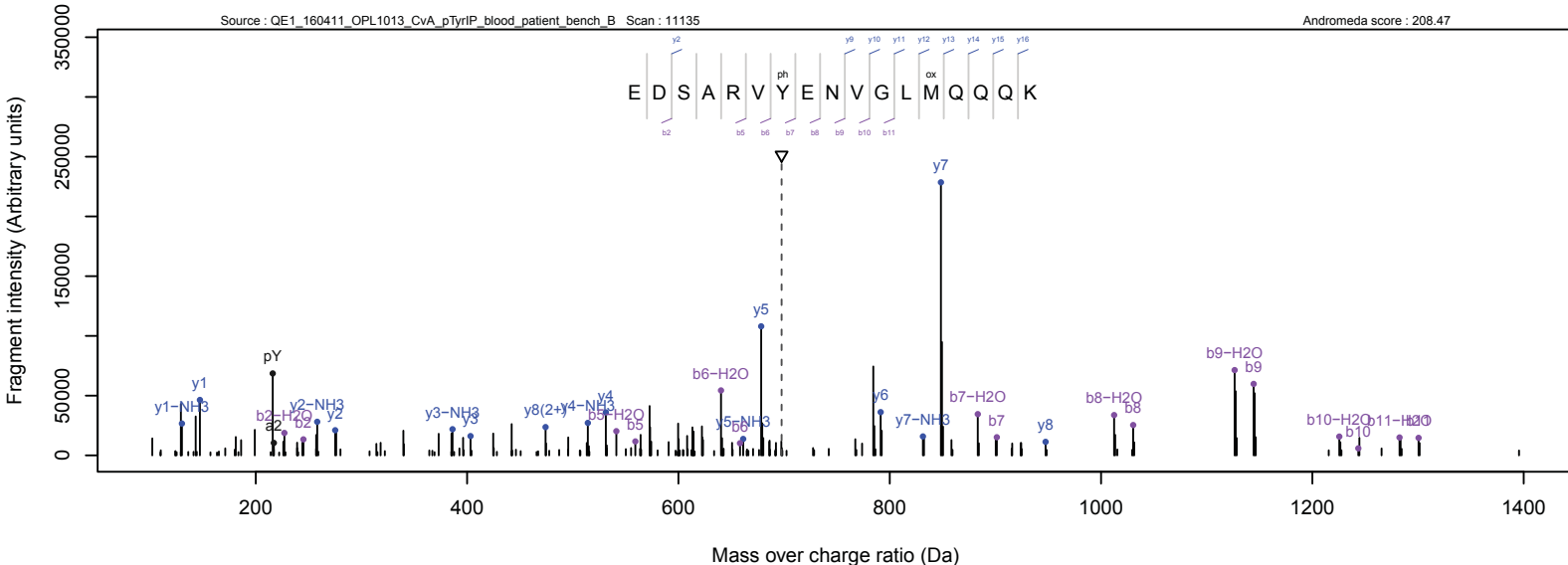
Andromeda score: 210.44



## EDSARVY(ph)ENVGLM(ox)QQQK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 11135

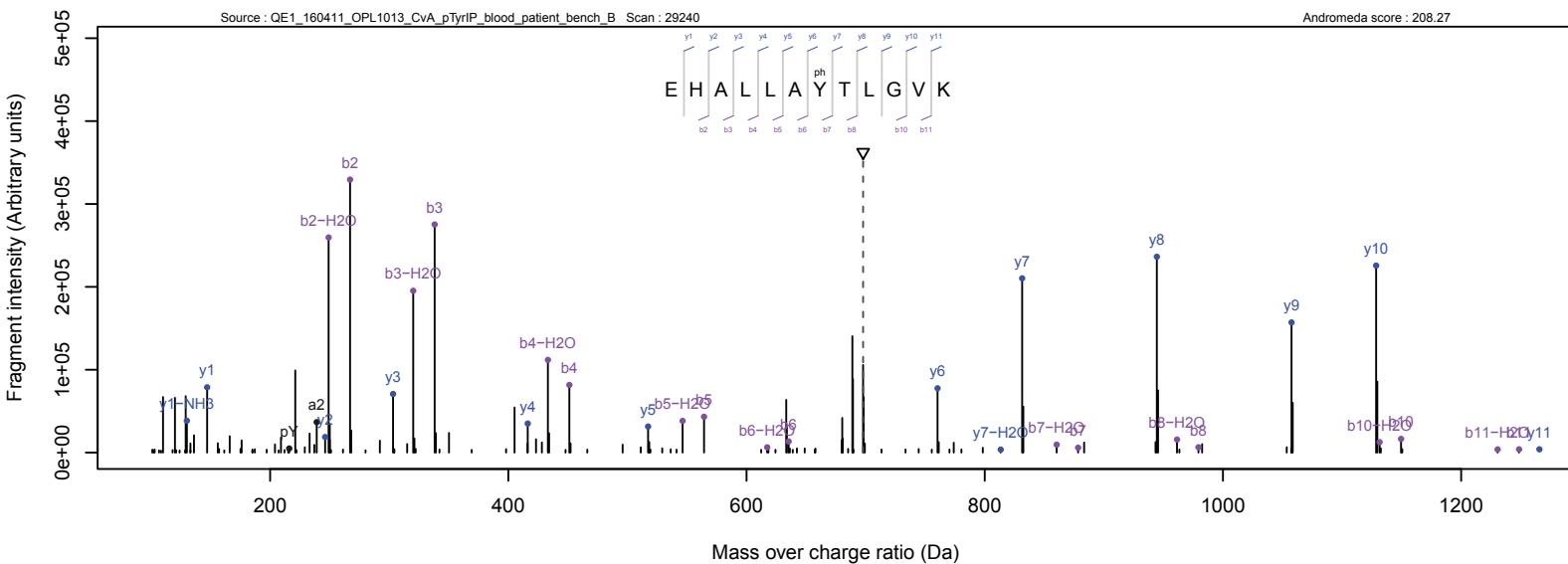
Andromeda score: 208.47



**EHALLAY(ph)TLGVK-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 29240

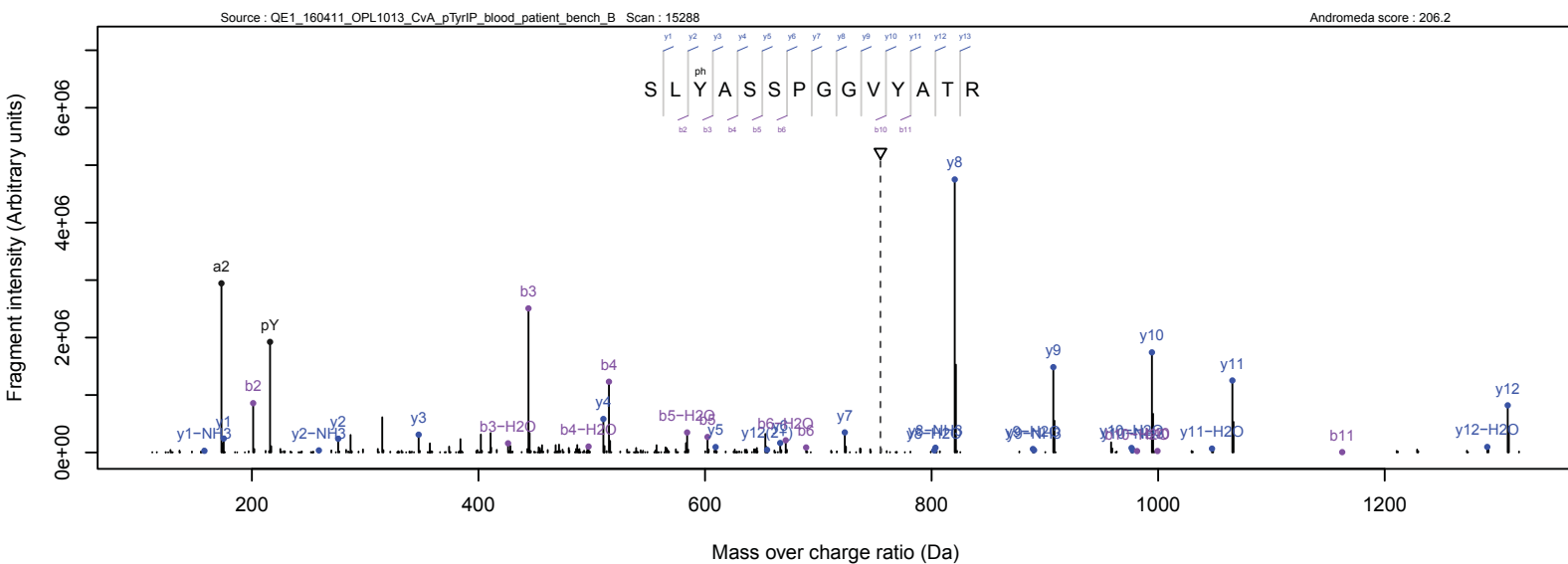
Andromeda score: 208.27



**SLY(ph)ASSPGVYATR-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 15288

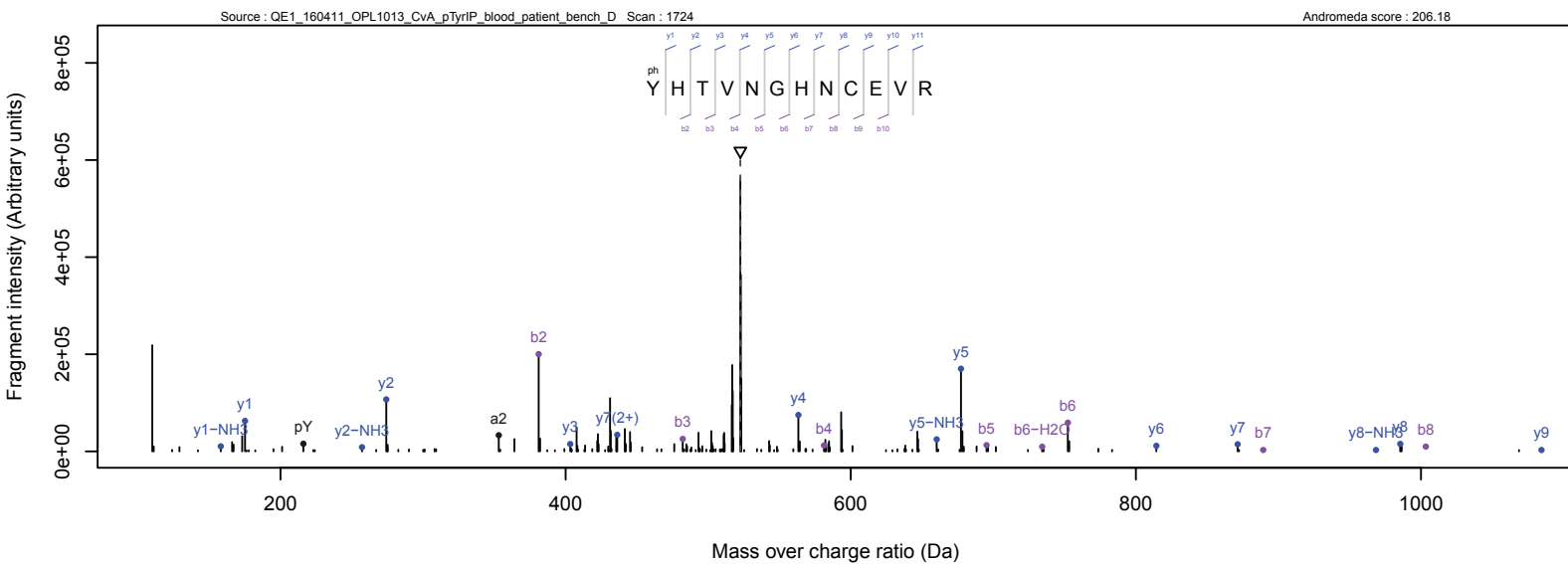
Andromeda score: 206.2



**Y(ph)HTVNGHNCVVR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 1724

Andromeda score: 206.18



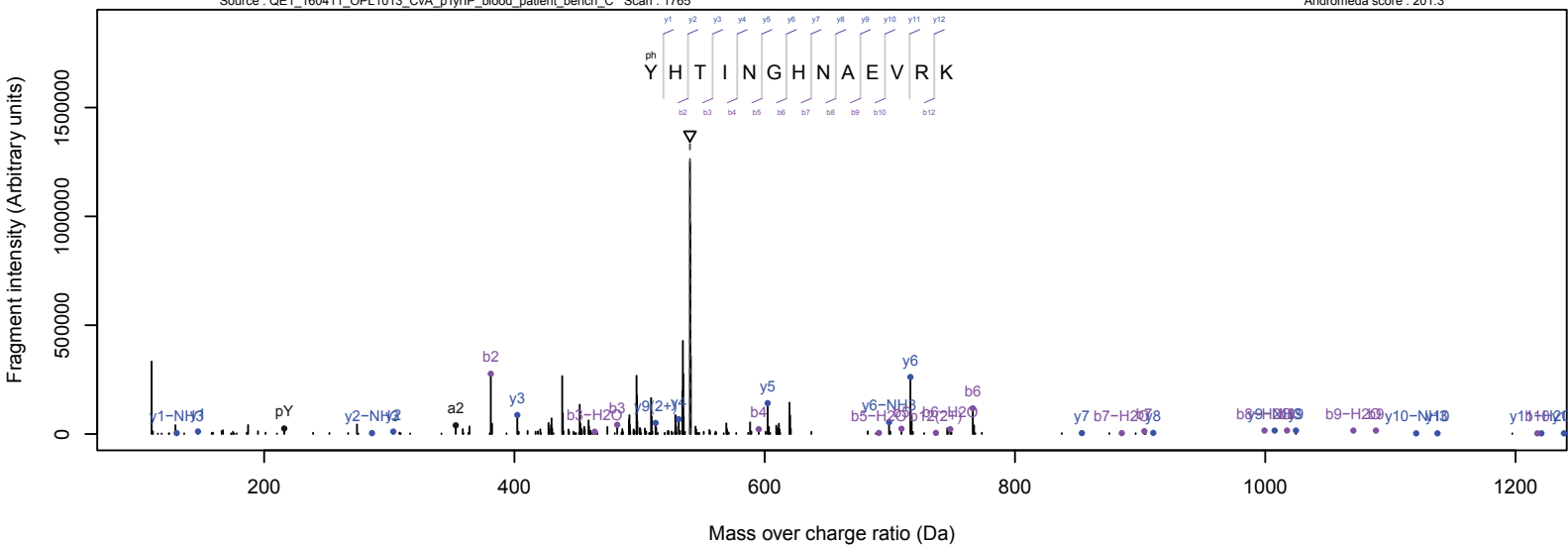




## Y(ph)HTINGHNAEVRK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 1765

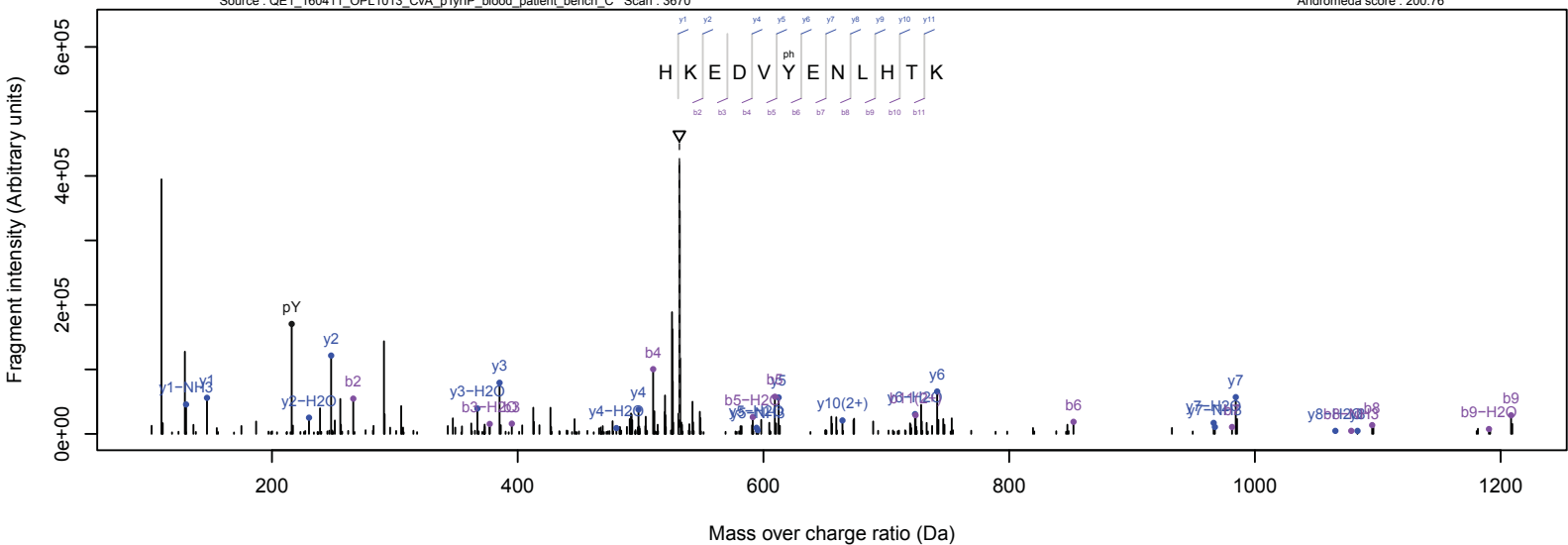
Andromeda score: 201.3



## HKEDVY(ph)ENLHTK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 3670

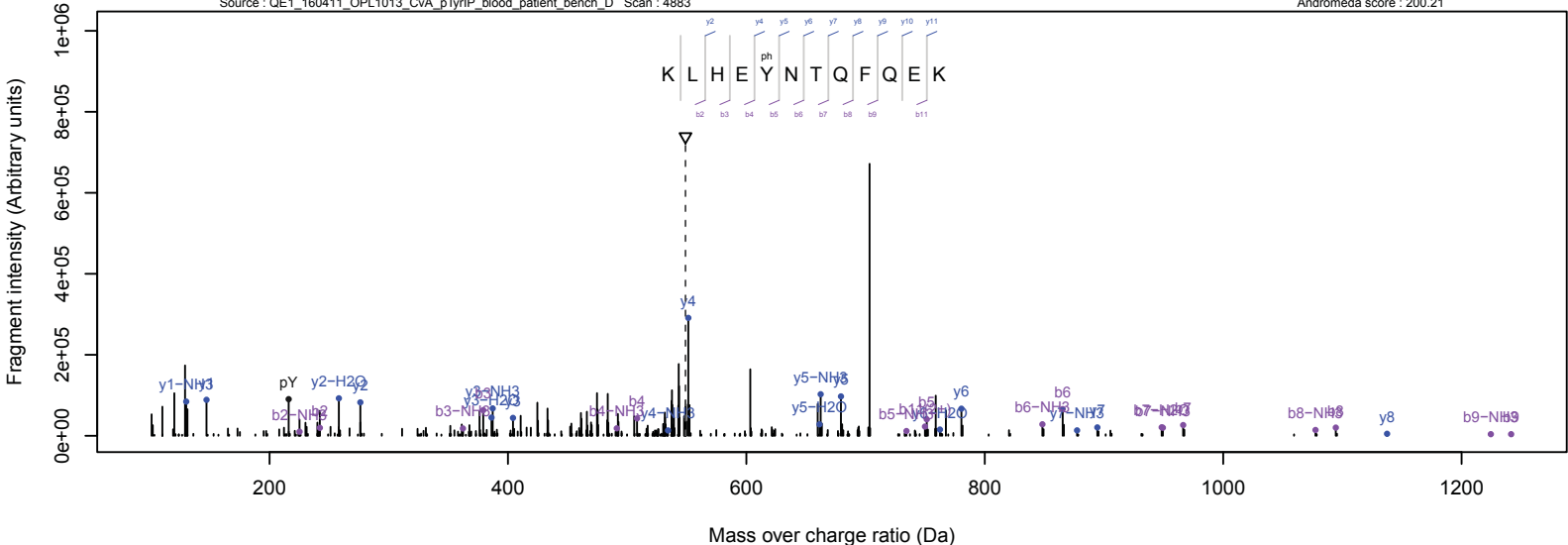
Andromeda score: 200.76



## KLHEY(ph)NTQFQEK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 4883

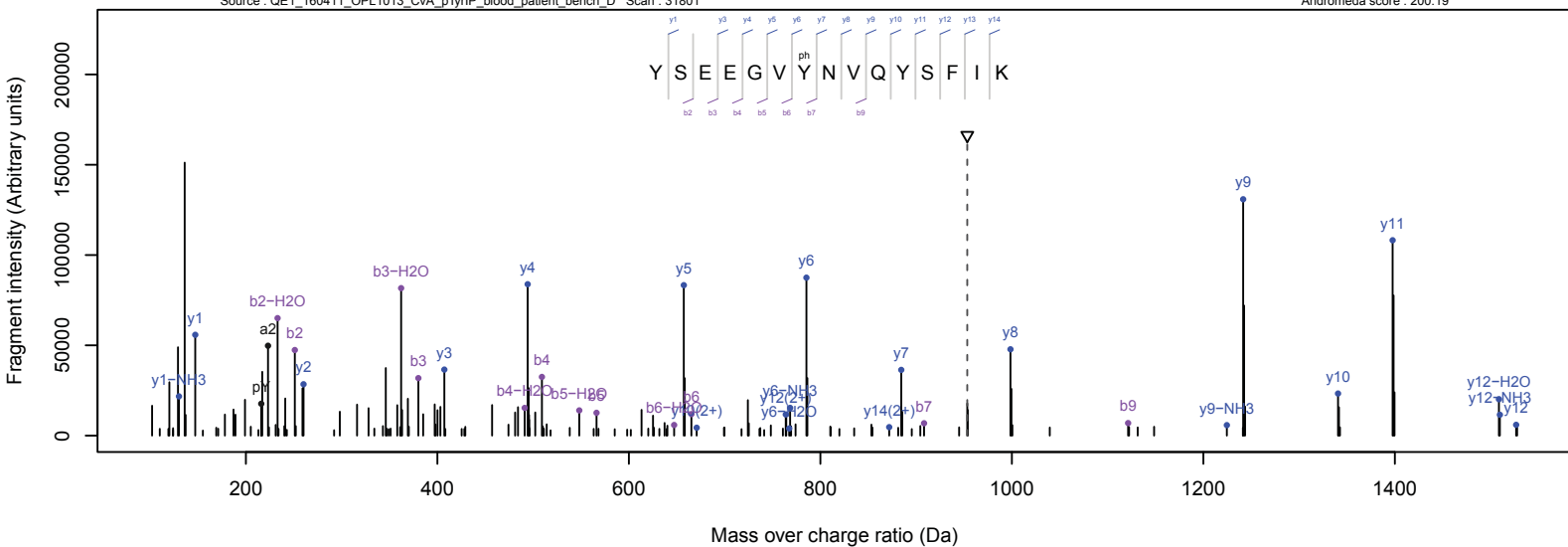
Andromeda score: 200.21



**YSEEGVY(ph)NVQYSFIK-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 31801

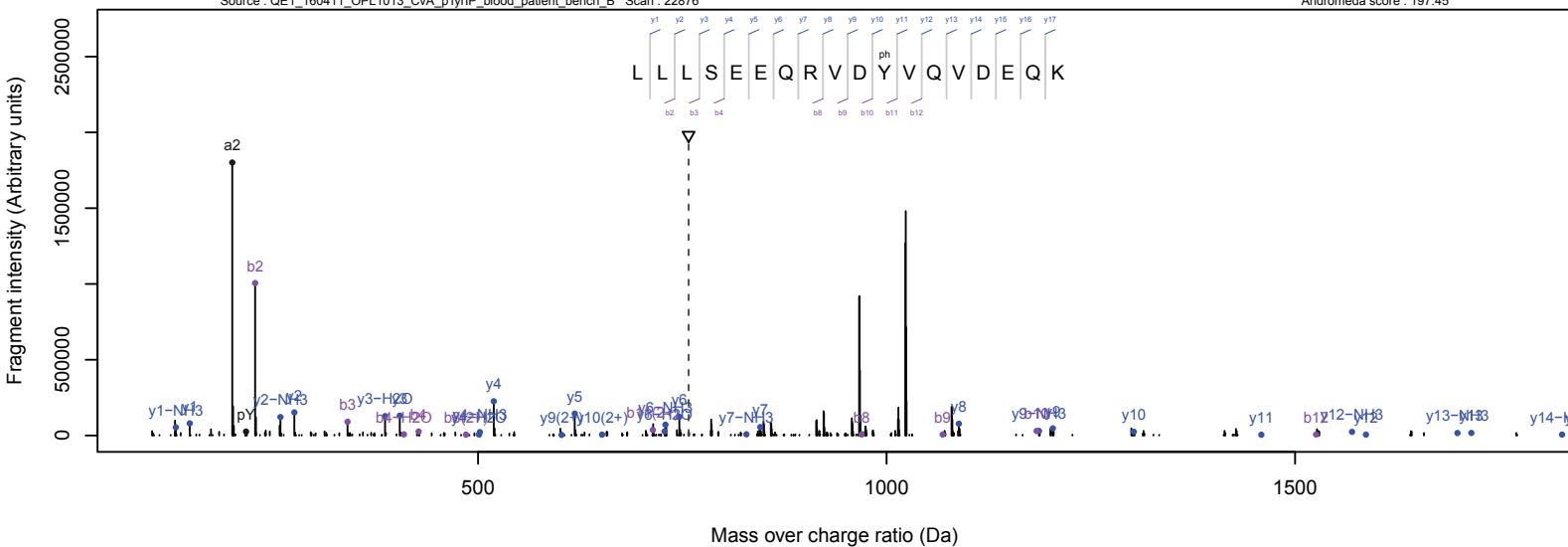
Andromeda score: 200.19



**LLLSEEQRVYD(ph)VQVDEQK-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 22876

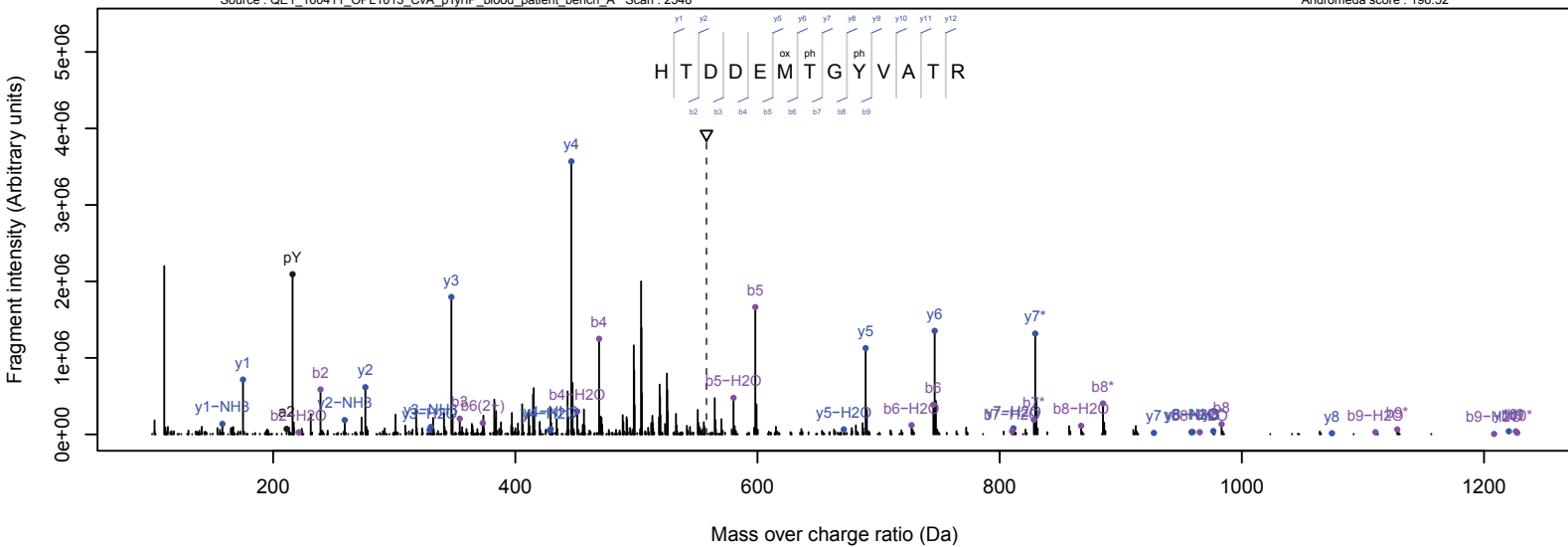
Andromeda score: 197.45



**HTDDEM(ox)T(ph)GYVATR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 2548

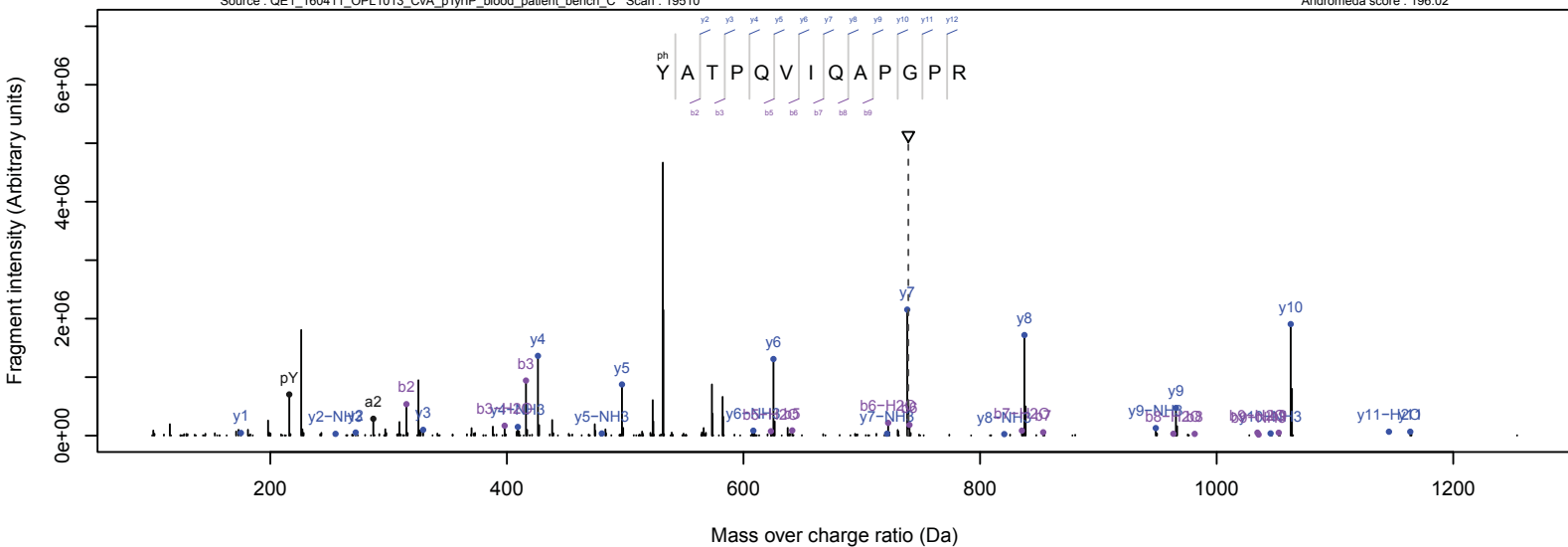
Andromeda score: 196.52



## Y(ph)ATPQVIQAPGR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 19510

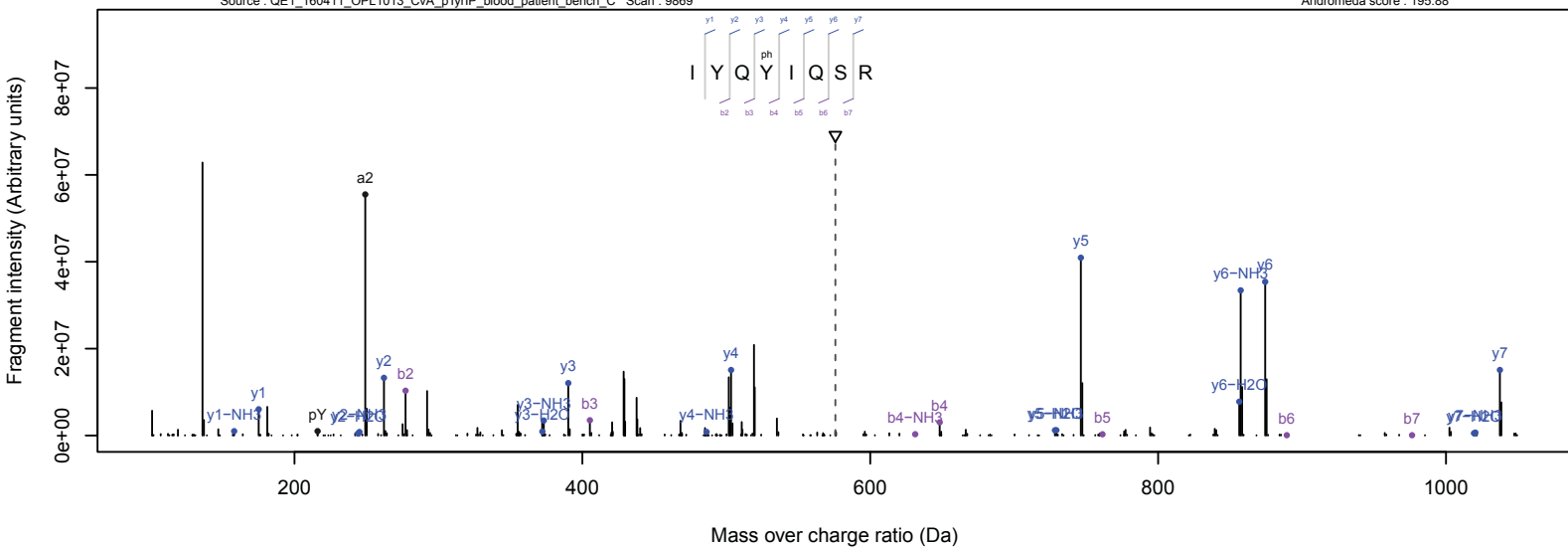
Andromeda score : 196.02



## IYQY(ph)IQSR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 9869

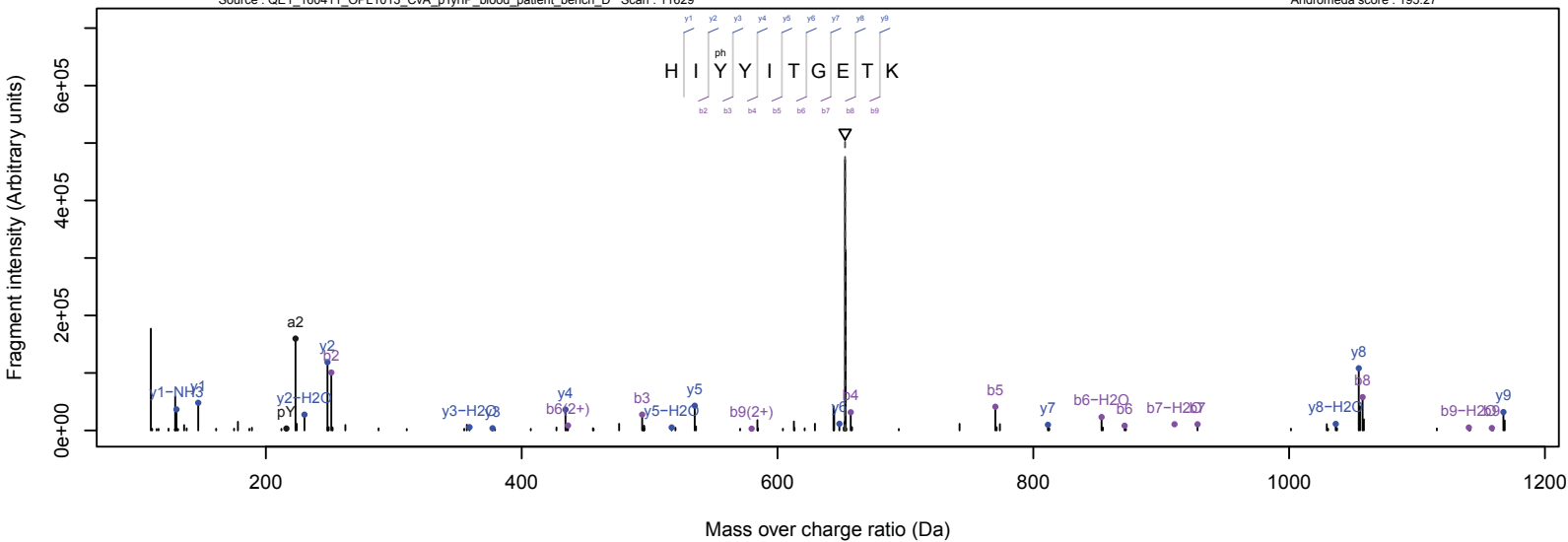
Andromeda score : 195.88



## HIY(ph)YITGETK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11629

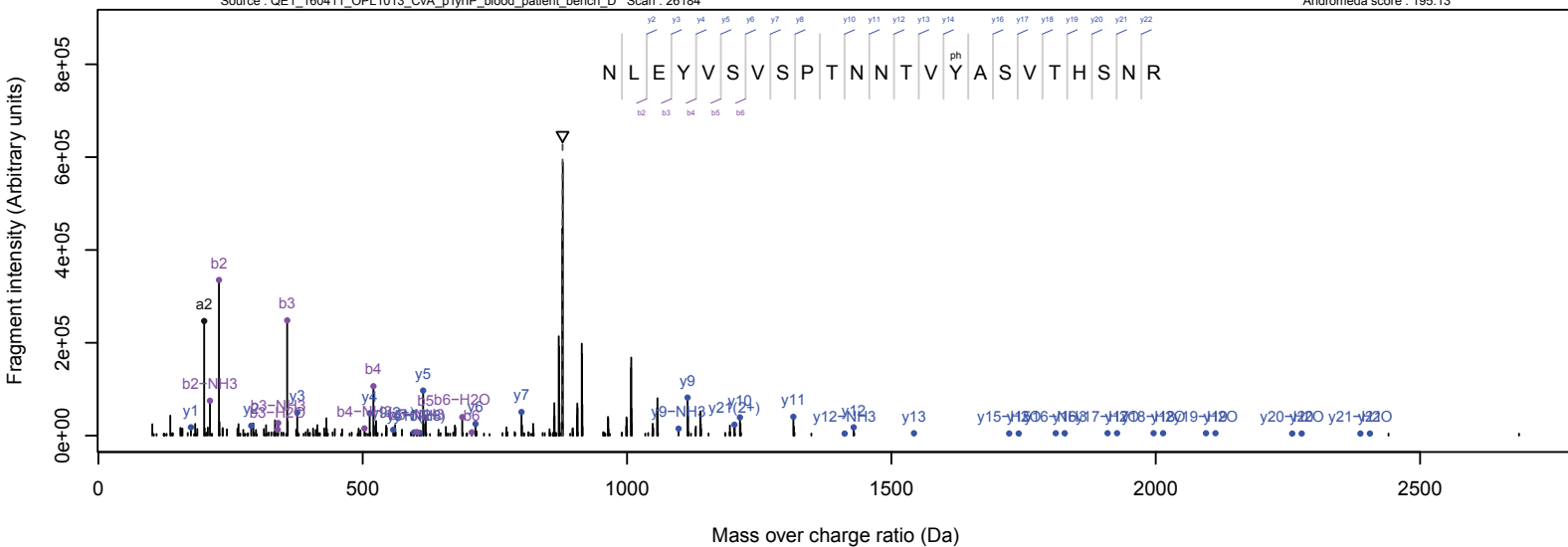
Andromeda score : 195.27



NLEYVSVSPTNNTVY(ph)ASVTHSNR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 26184

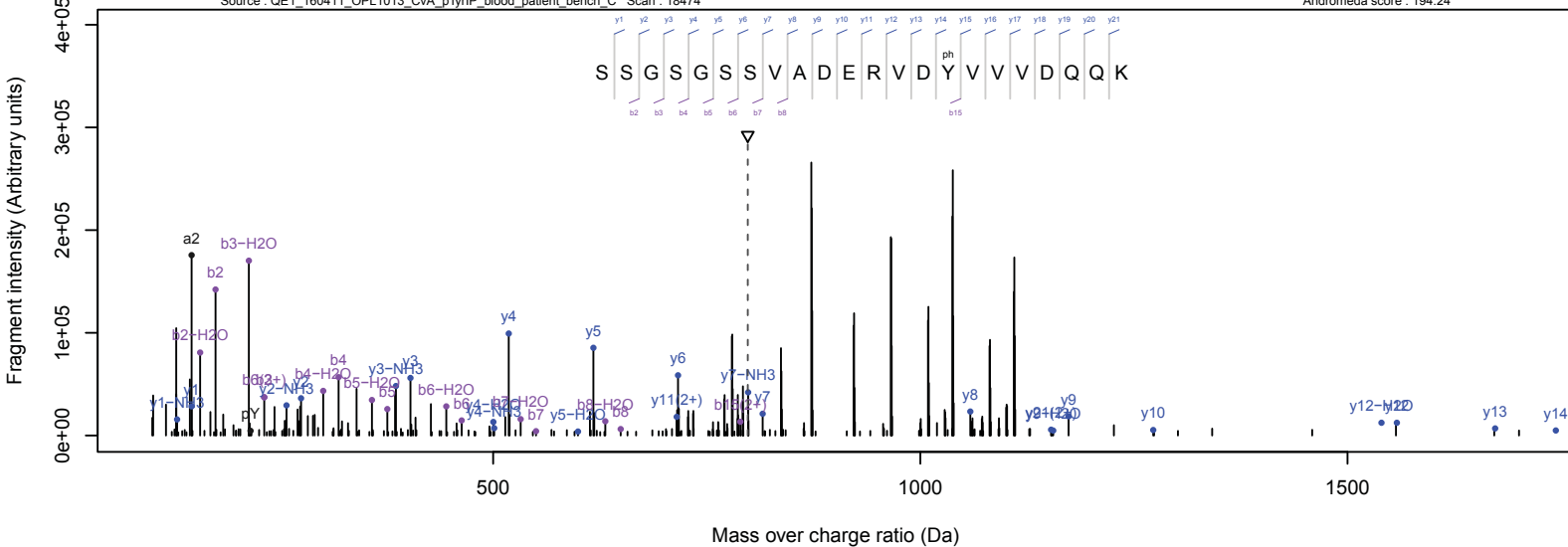
Andromeda score: 195.13



SSGSGSSVADERVDY(ph)VVDQK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 18474

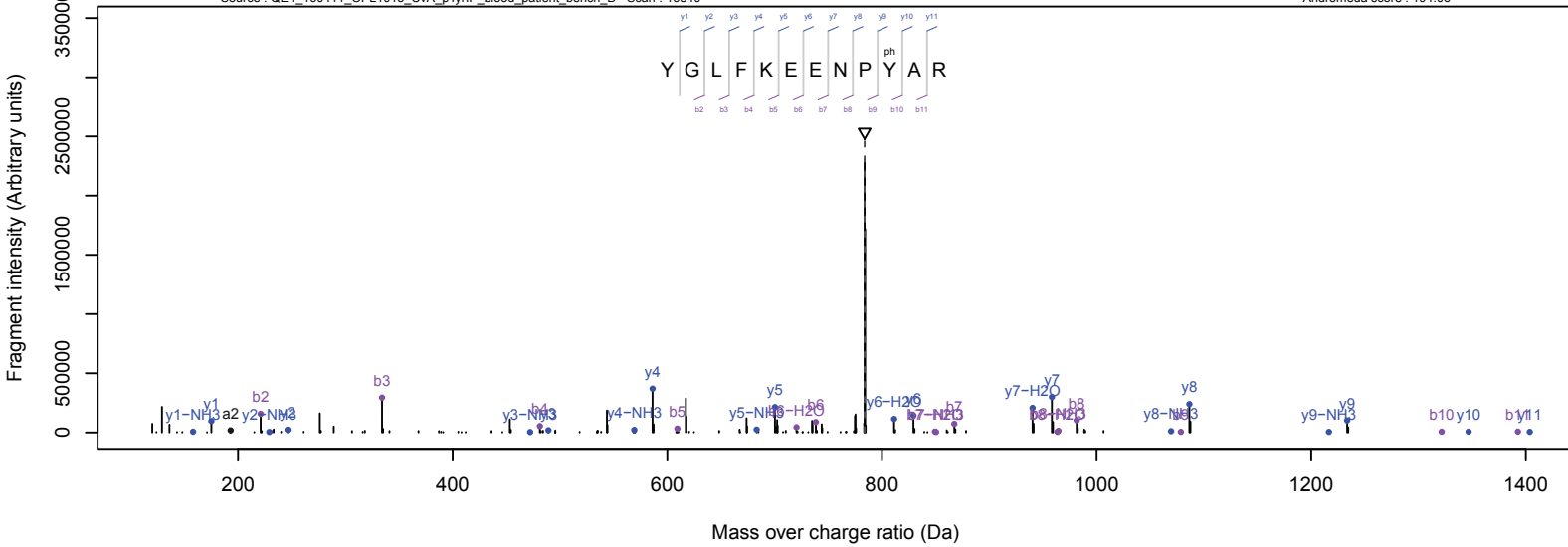
Andromeda score: 194.24



YGLFKEENPYAR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 16840

Andromeda score: 194.06

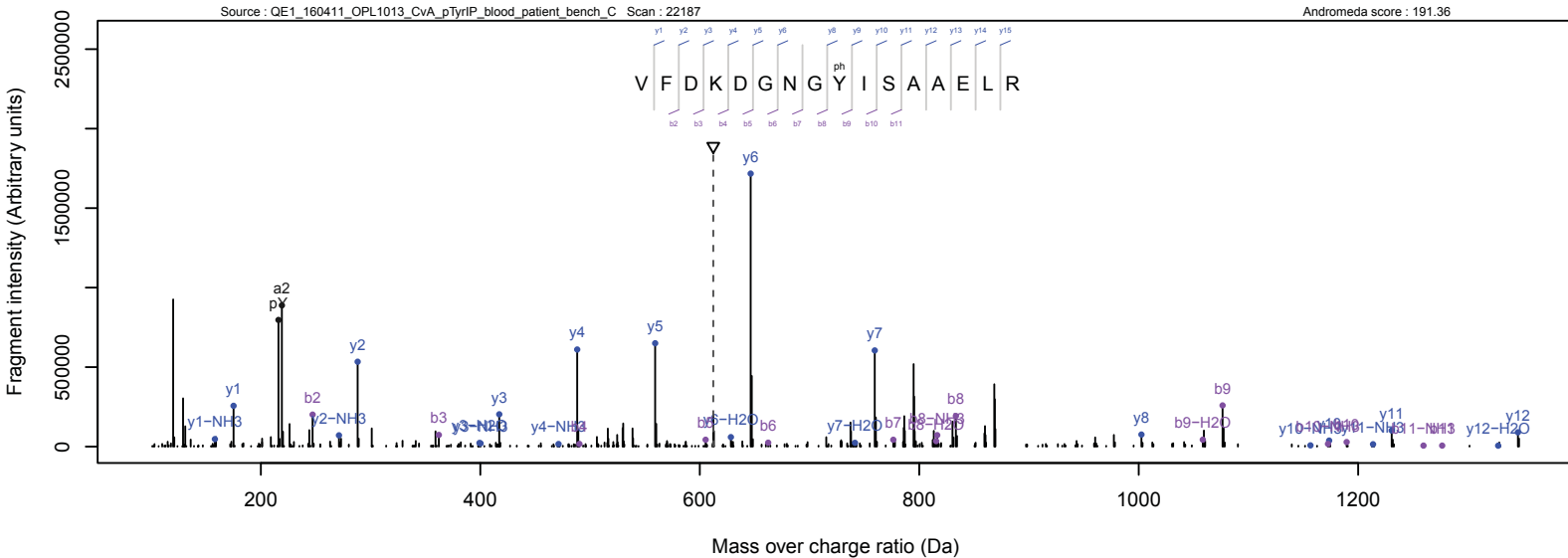




VFDKDGNGY(ph)ISAAELR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 22187

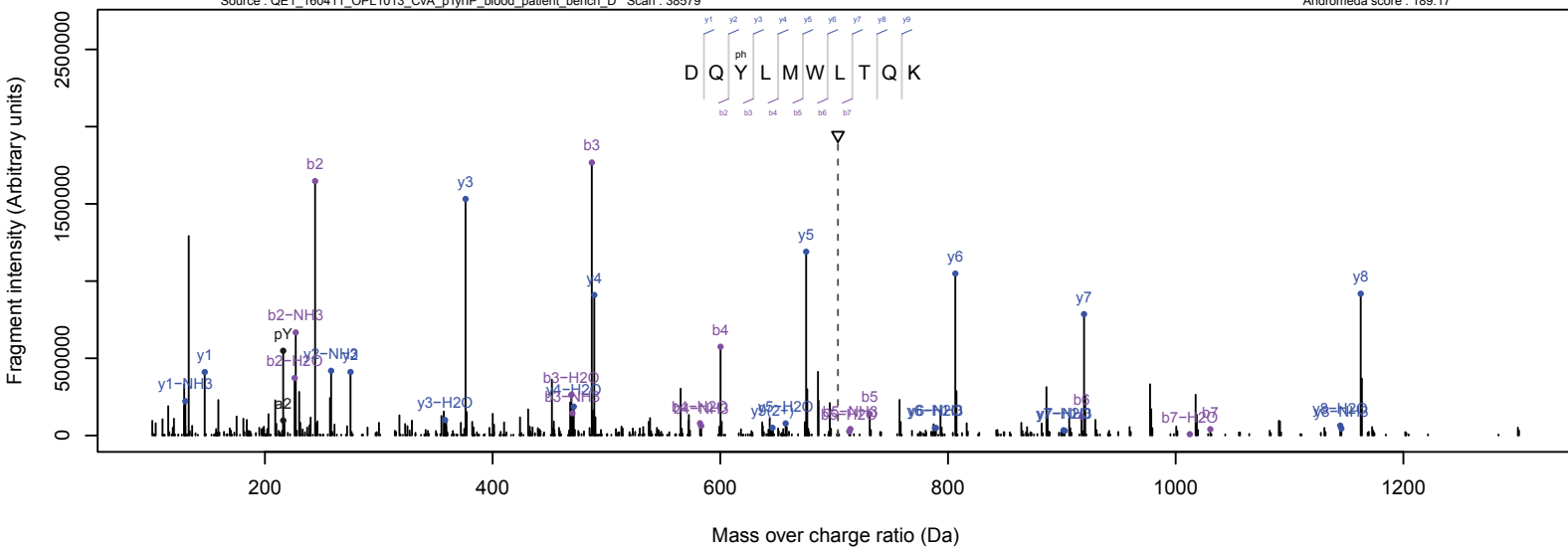
Andromeda score : 191.36



DQY(ph)LMWLTQK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 38579

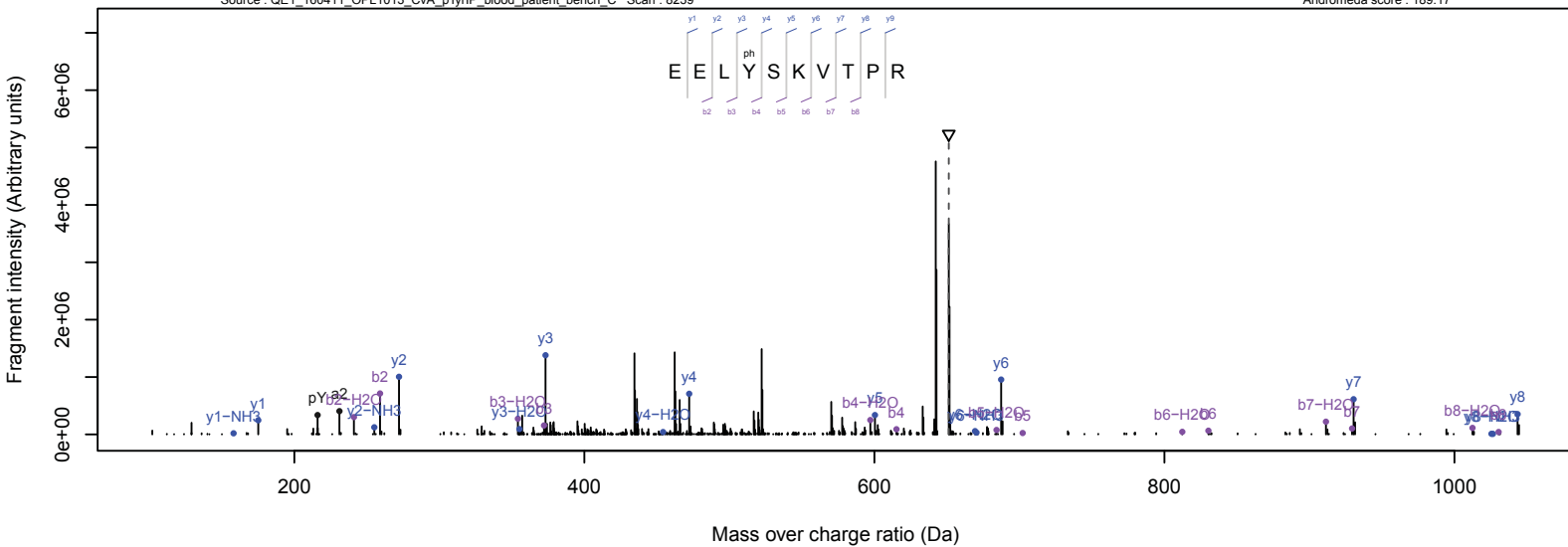
Andromeda score : 189.17



EELY(ph)SKVTPR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 8239

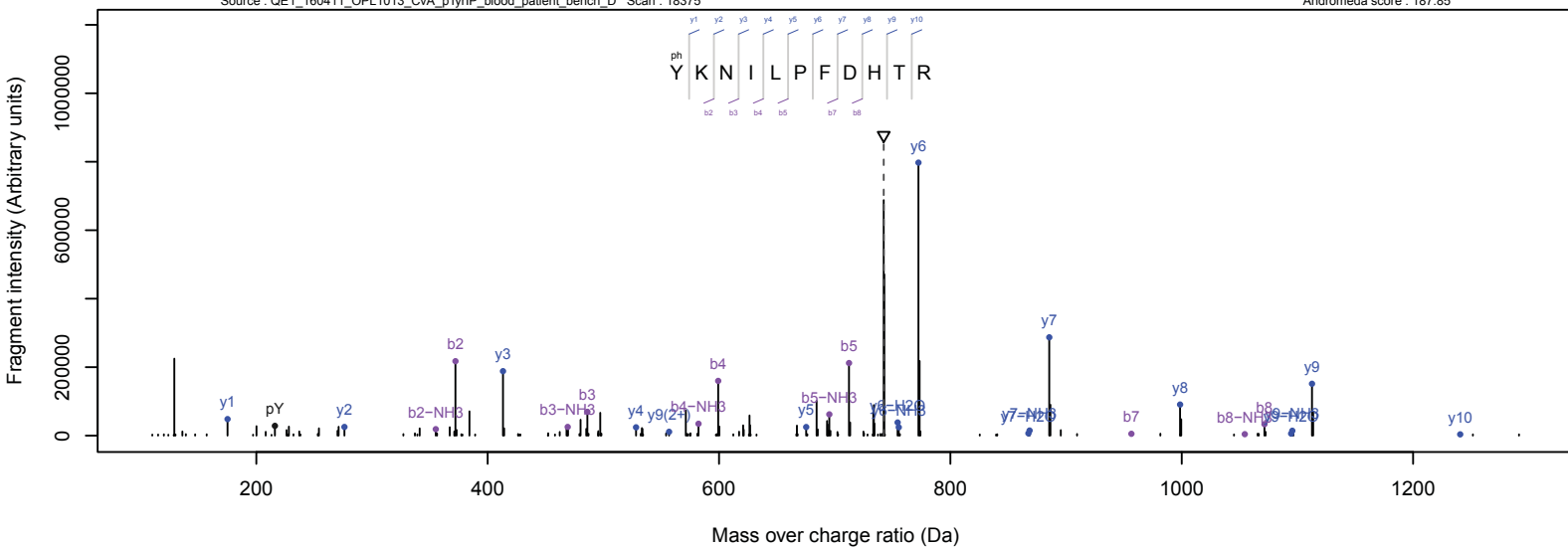
Andromeda score : 189.17



## Y(ph)KNILPFDHTR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 18375

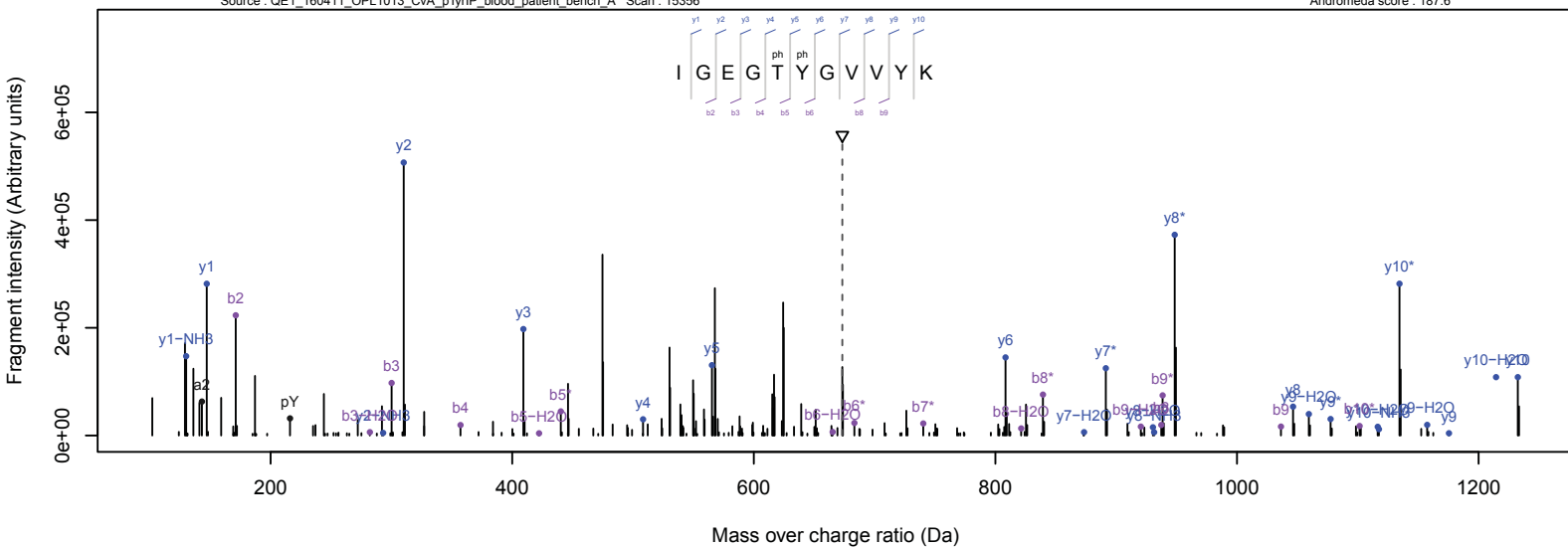
Andromeda score : 187.85



## IGEGT(ph)Y(ph)GVVYK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 15356

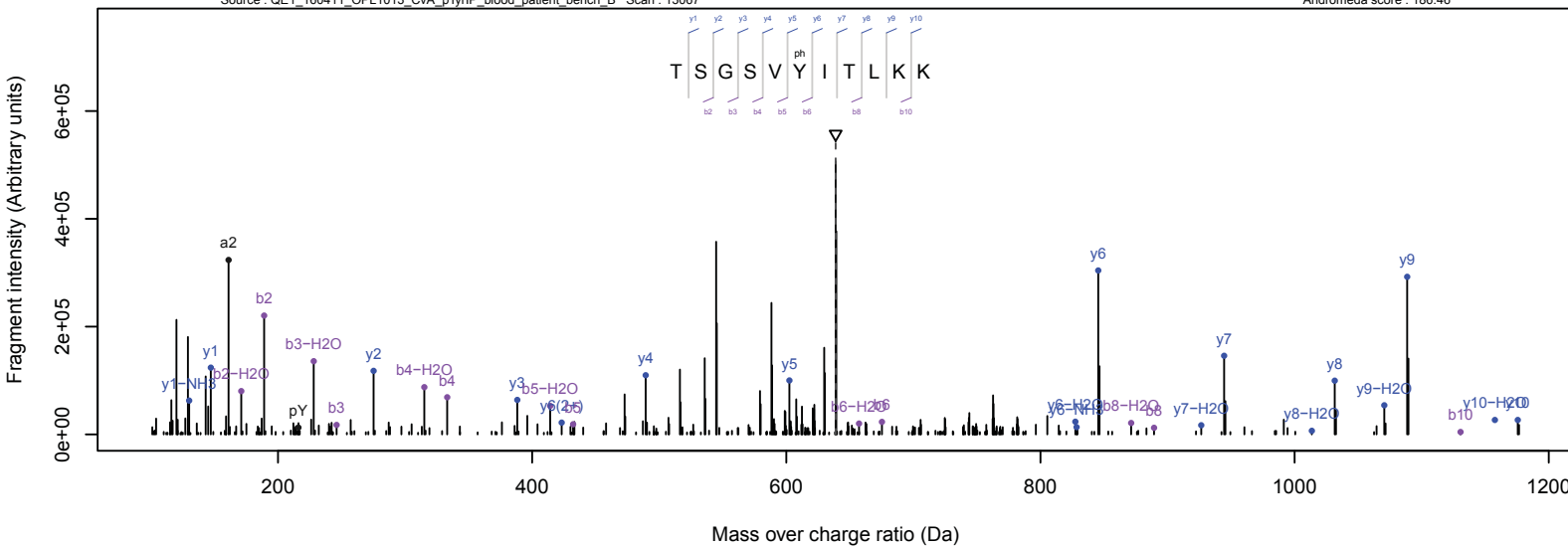
Andromeda score : 187.6



## TSGSVY(ph)ITLKK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 13067

Andromeda score : 186.46

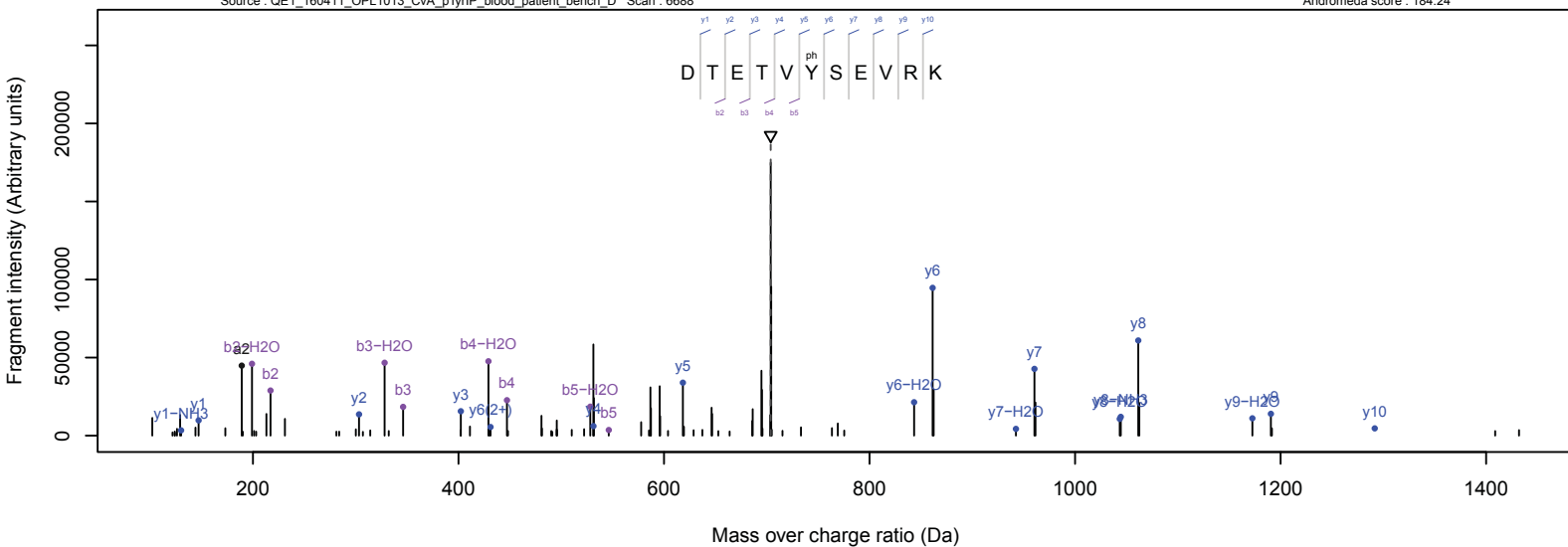




## DTETVY(ph)SEVRK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 6688

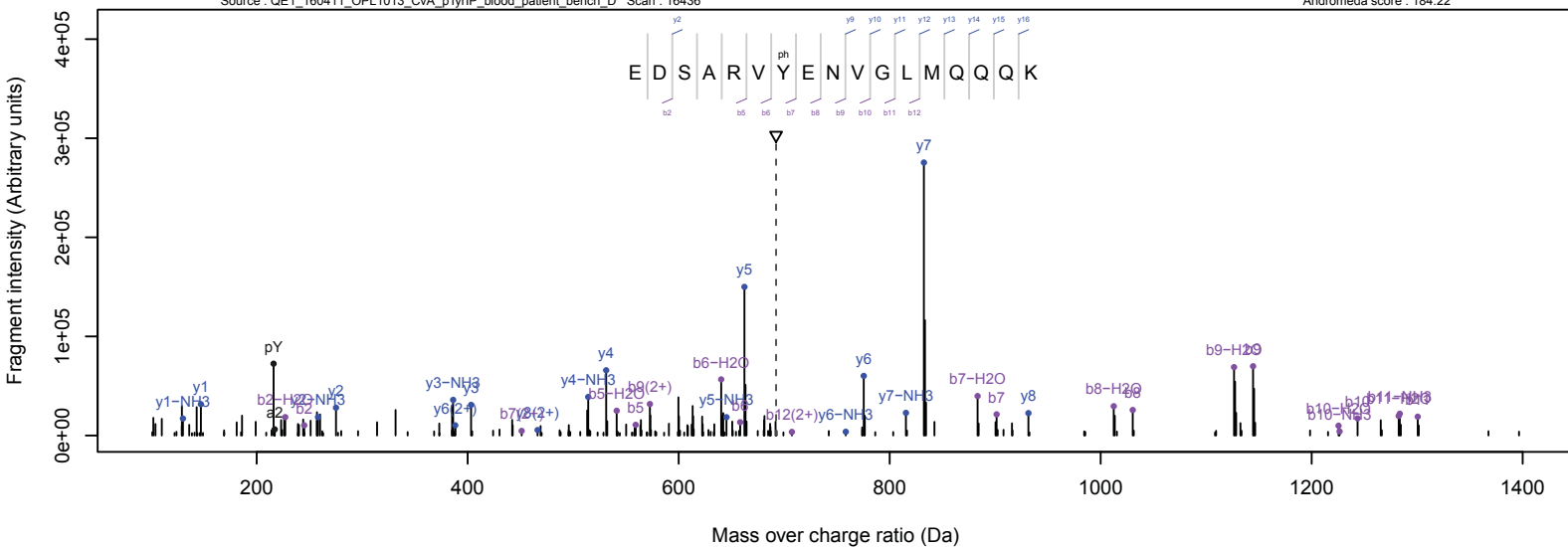
Andromeda score : 184.24



## EDSARVY(ph)ENVGLMQQK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 16436

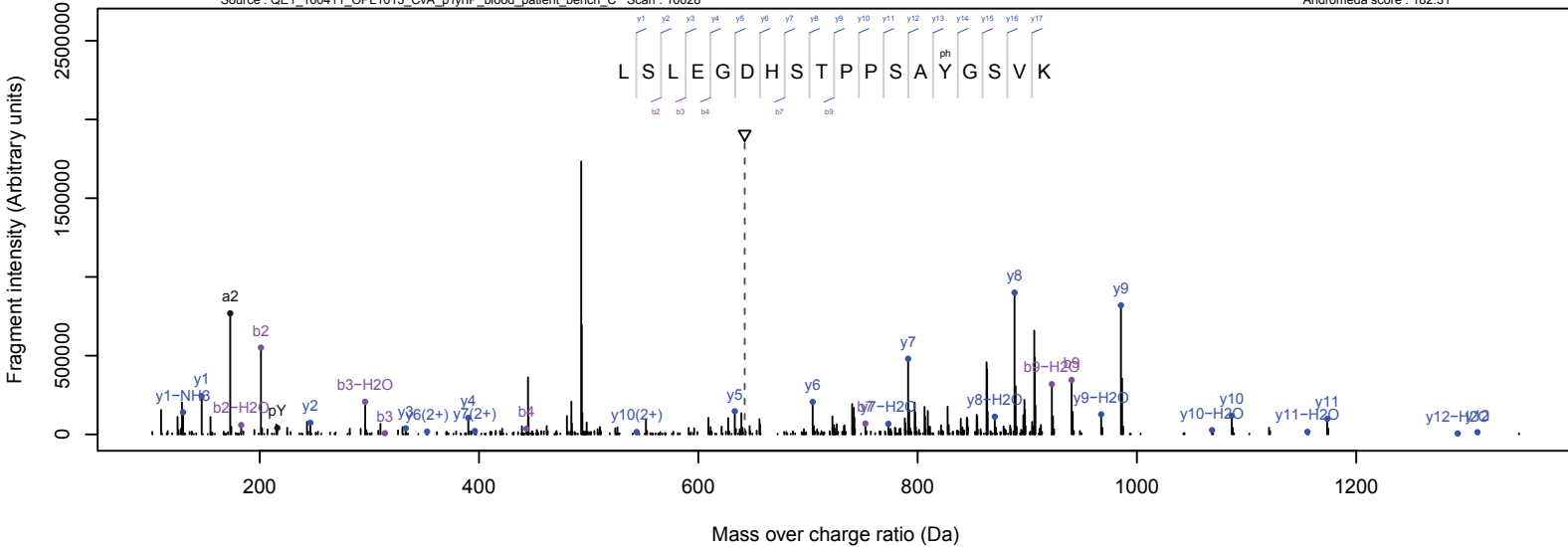
Andromeda score : 184.22



## LSLEGDHSTPPSAY(ph)GSVK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16028

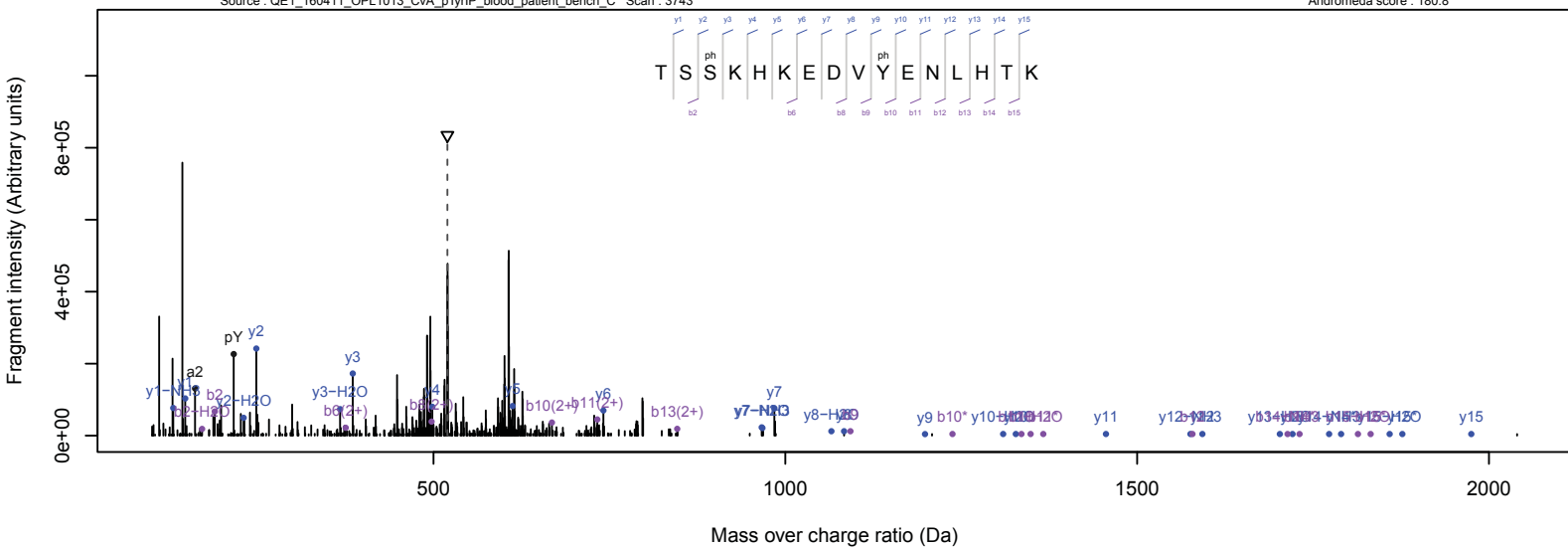
Andromeda score : 182.31



TSS(ph)KHKEDVY(ph)ENLHTK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3743

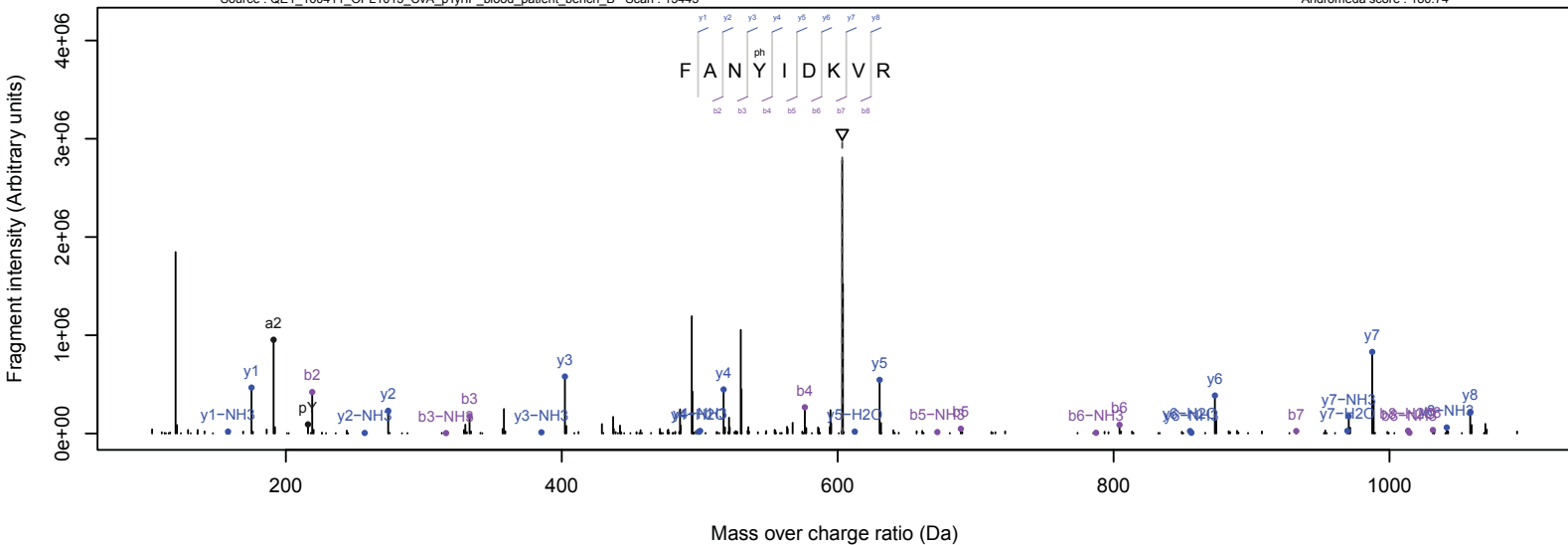
Andromeda score : 180.8



FANY(ph)IDKVR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 13445

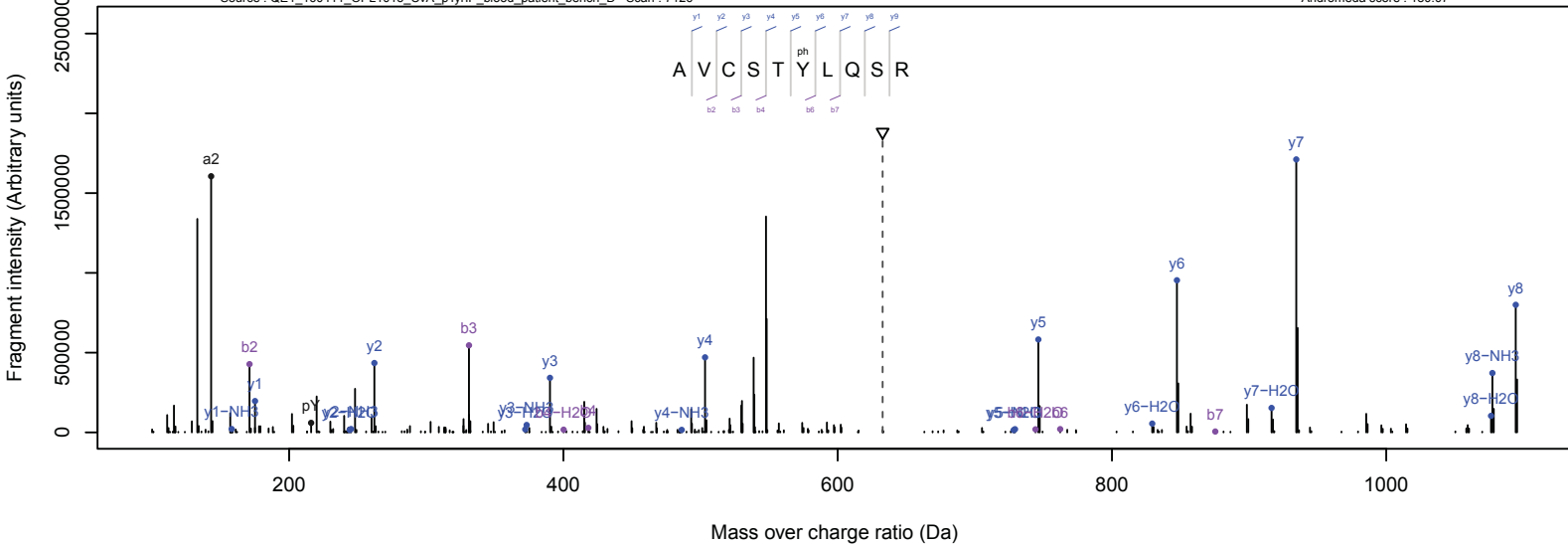
Andromeda score : 180.74



AVCSTY(ph)LQSR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7126

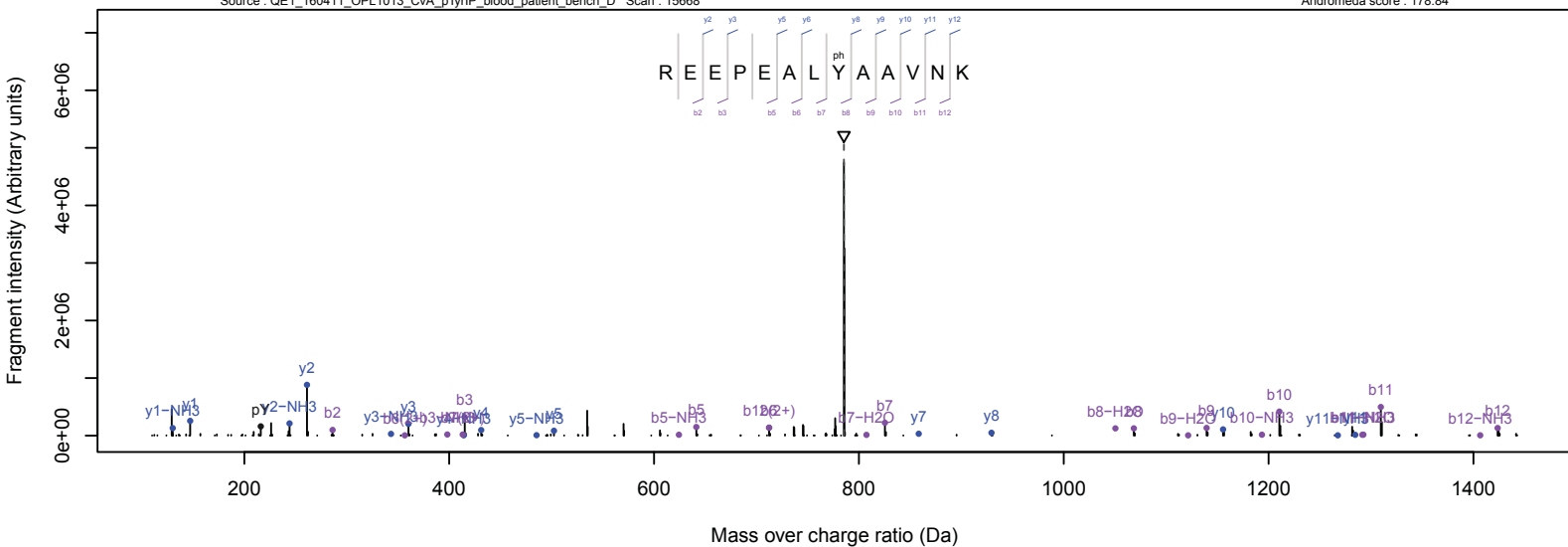
Andromeda score : 180.07



REEPEALY(ph)AAVNK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 15668

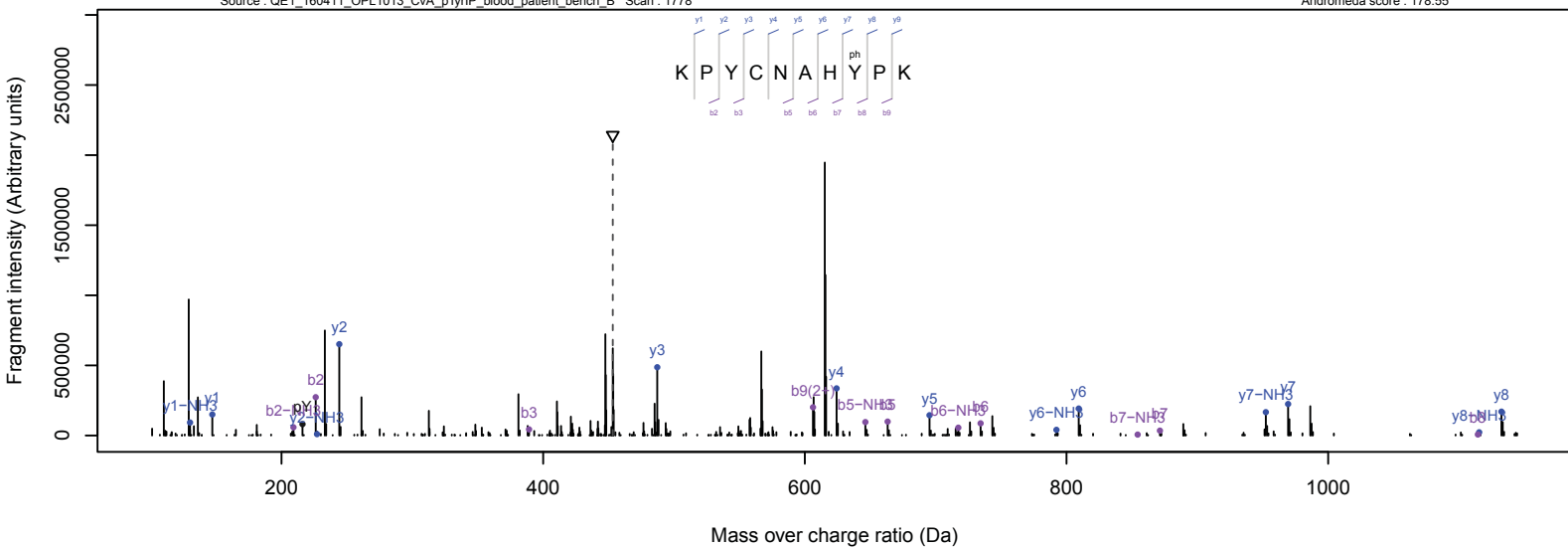
Andromeda score : 178.84



KPYCNAHY(ph)PK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 1778

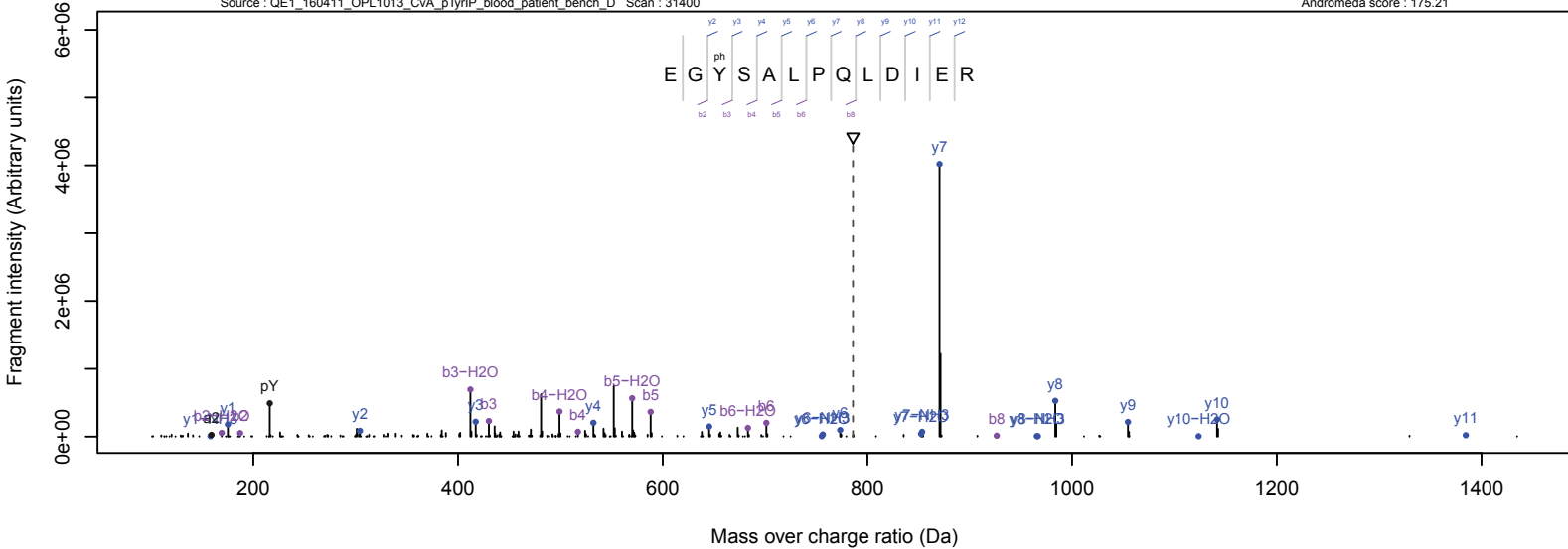
Andromeda score : 178.55



EGY(ph)SALPLDIER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 31400

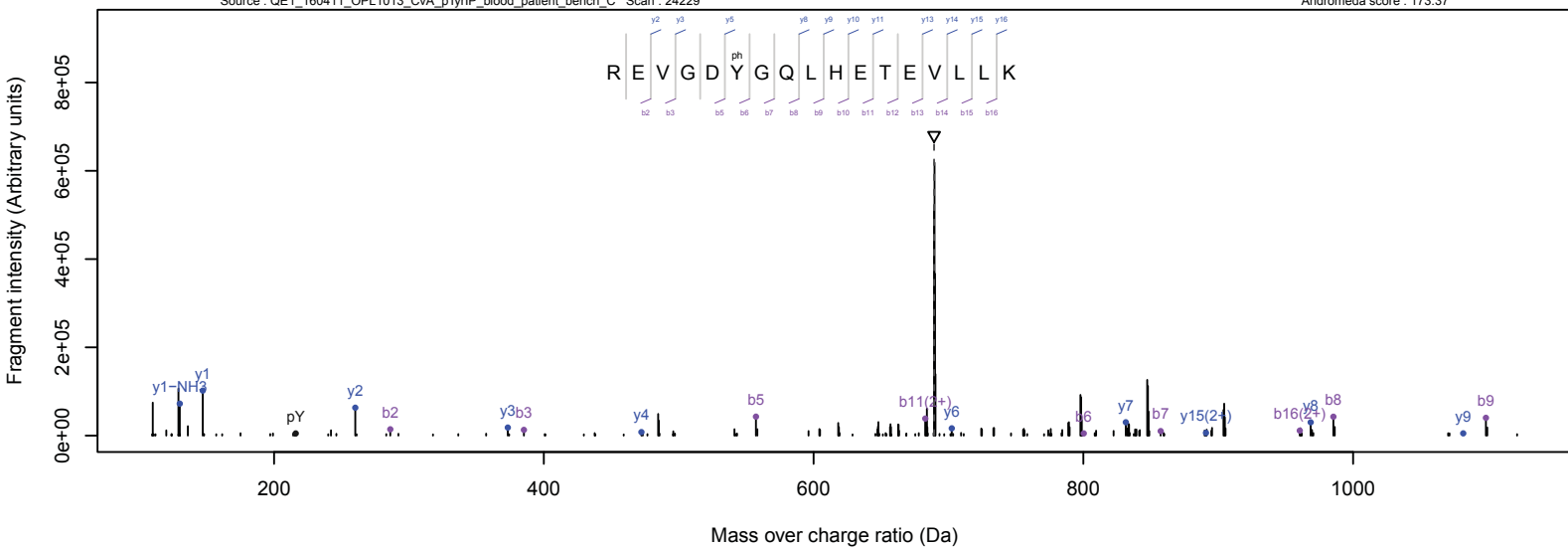
Andromeda score : 175.21



## REVGDY(ph)GQLHETEVLLK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 24229

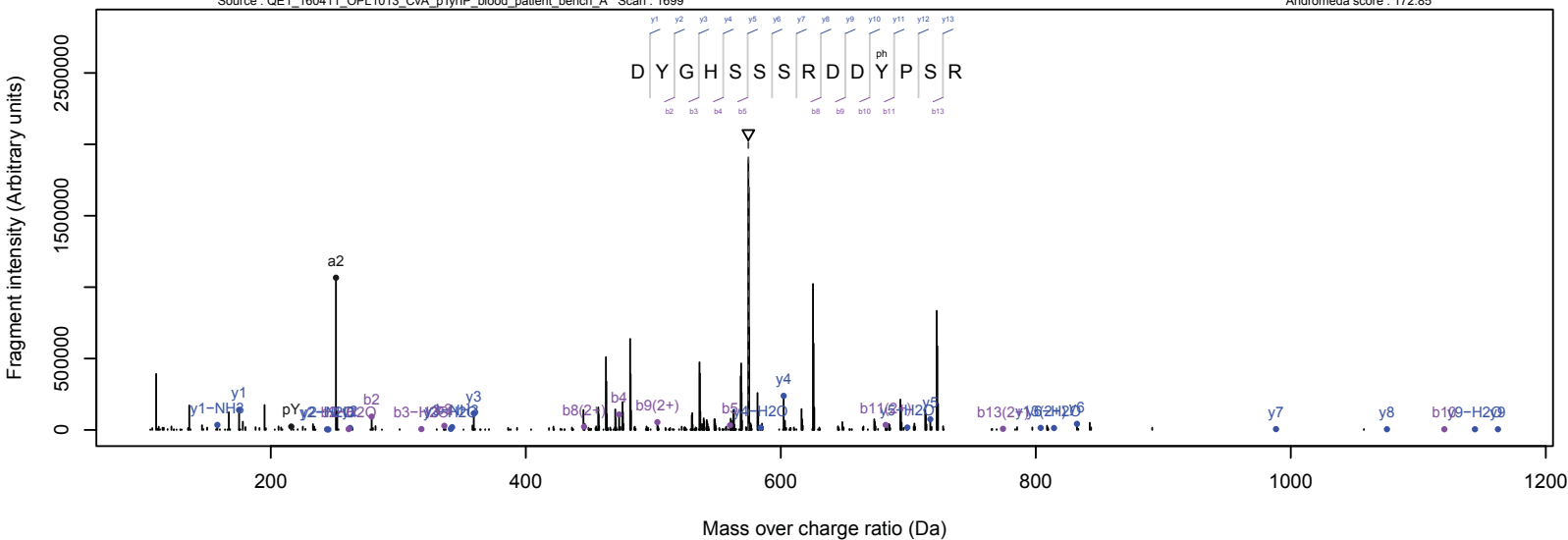
Andromeda score: 173.37



## DYGHSSRDYPSR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 1699

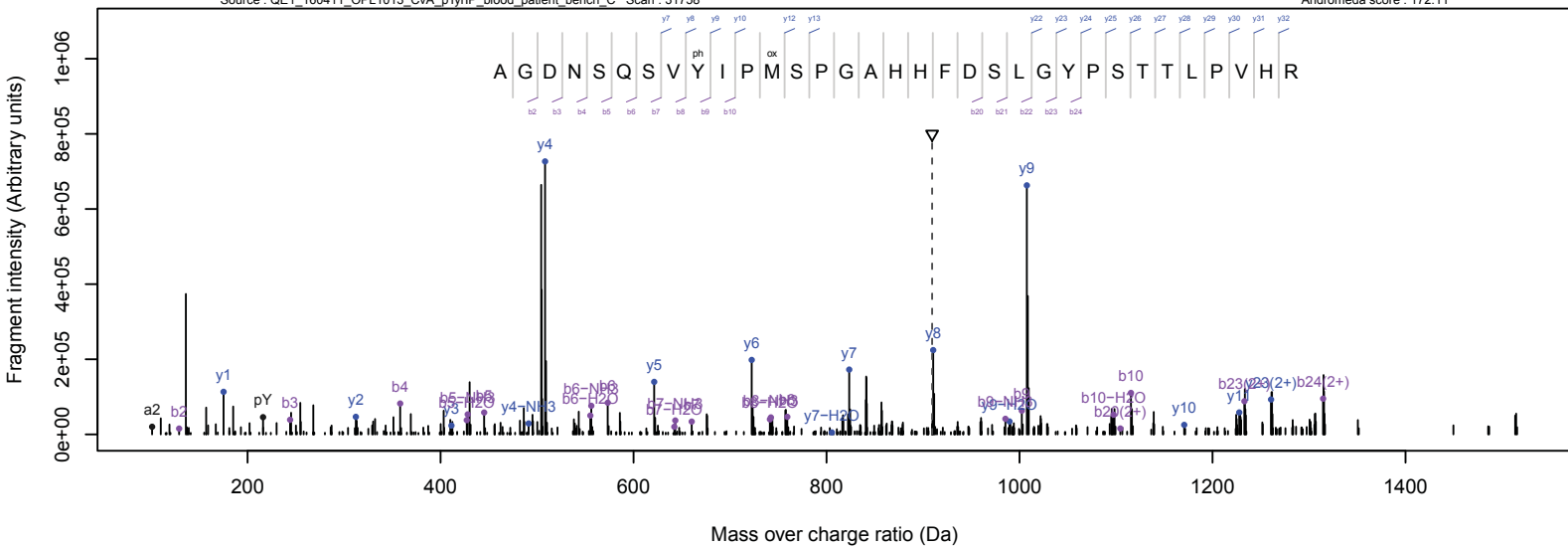
Andromeda score: 172.85

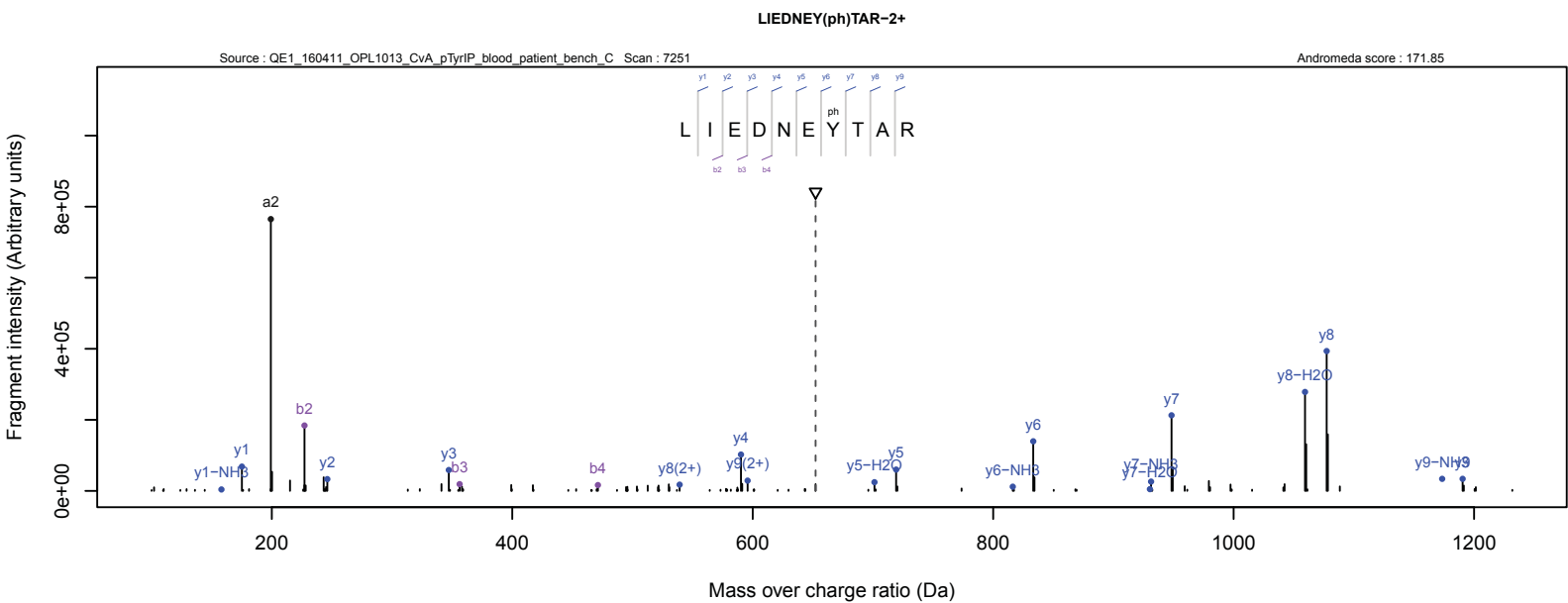
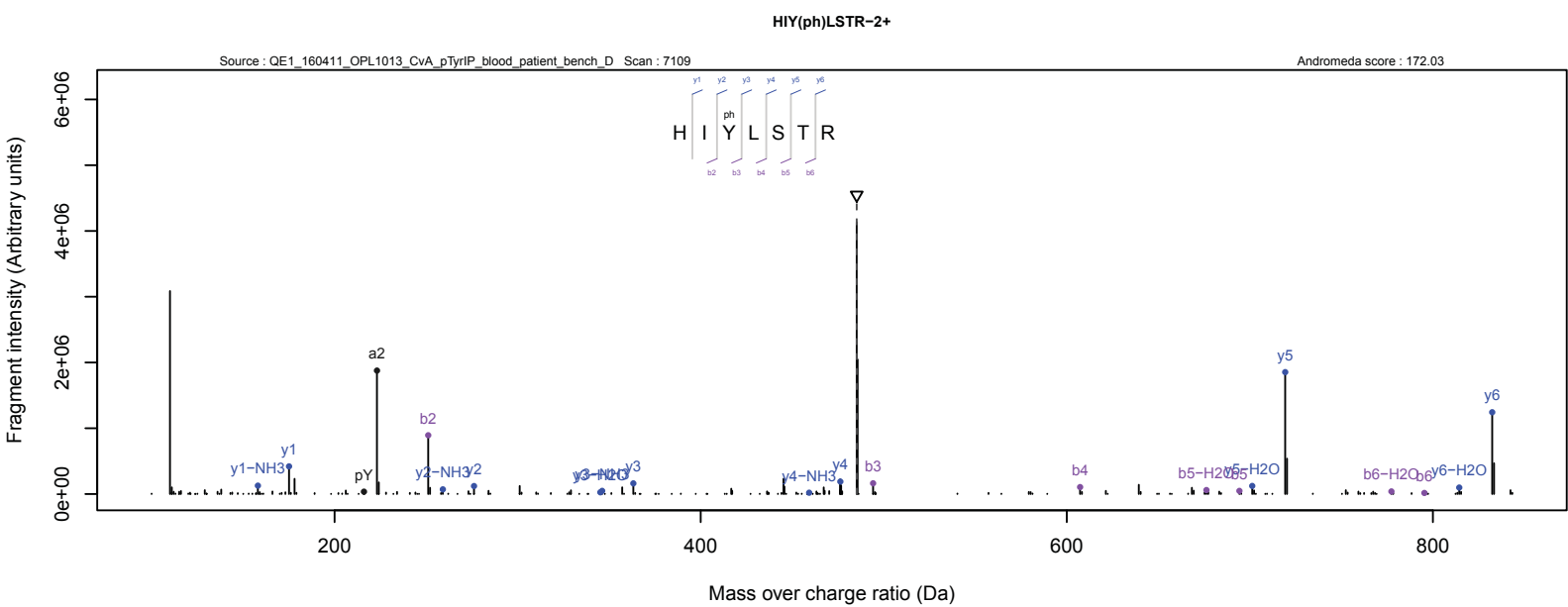
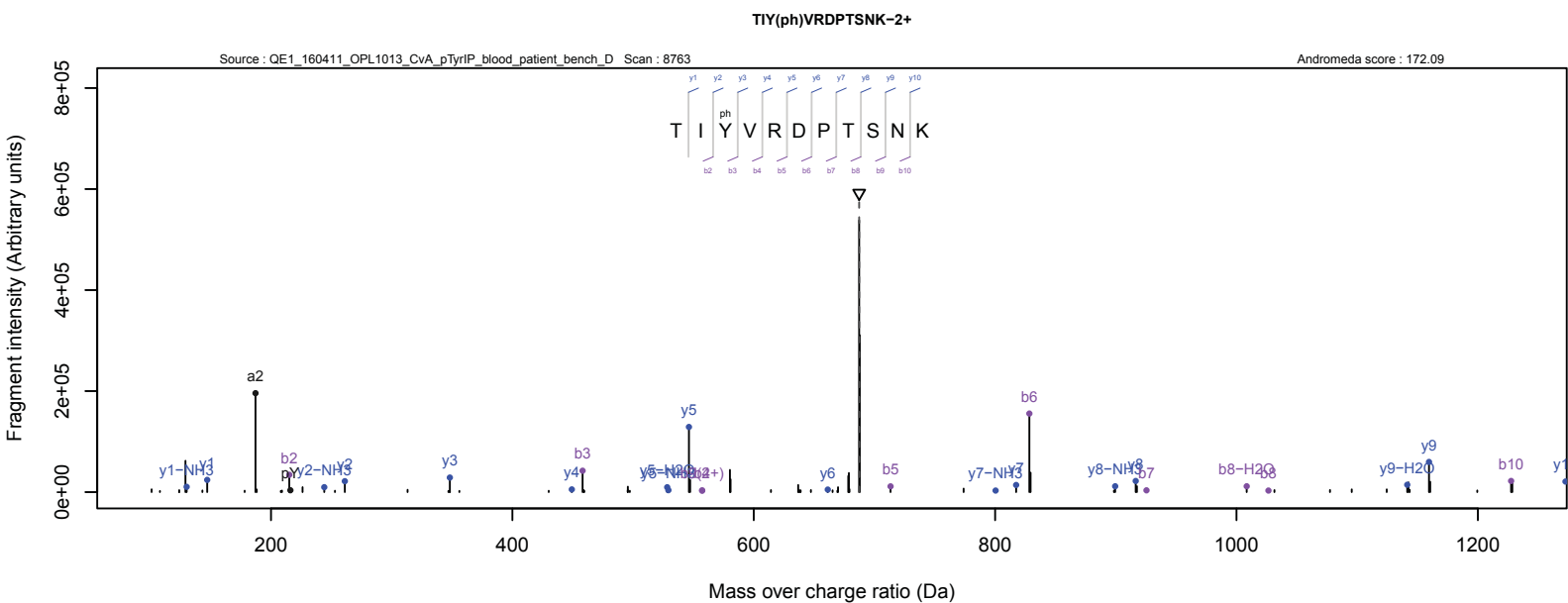


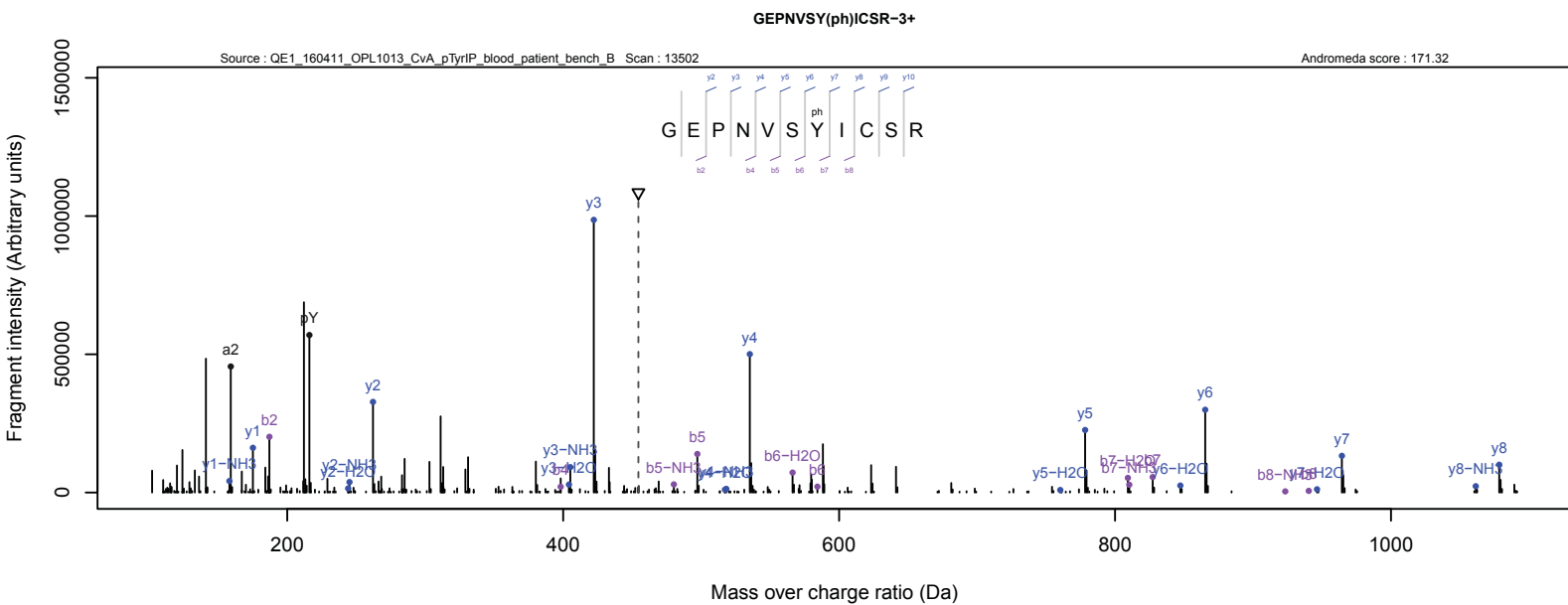
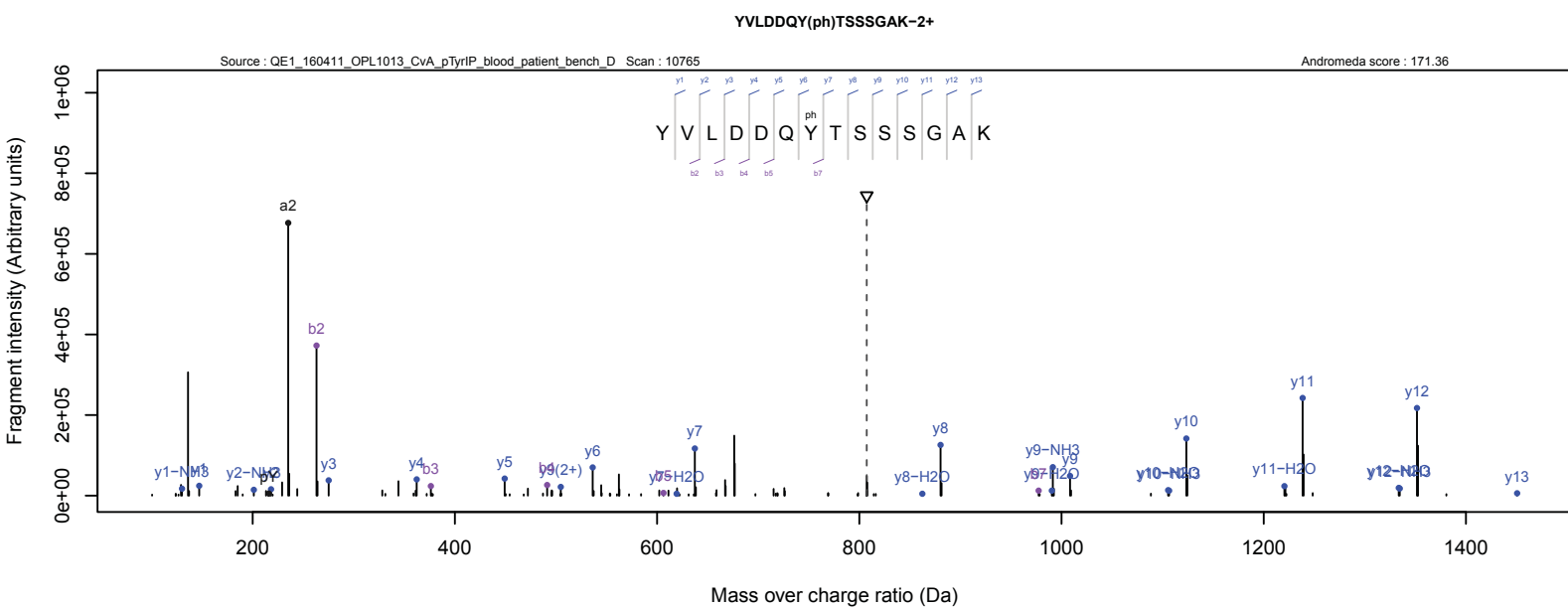
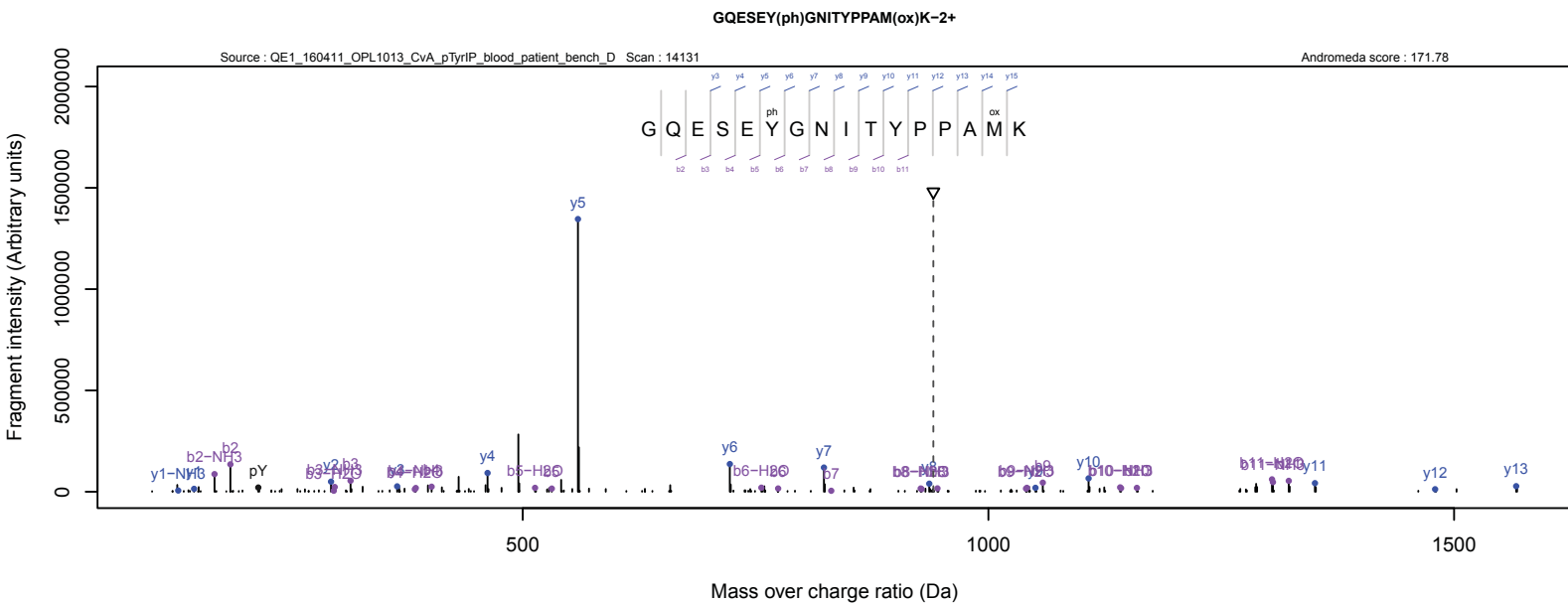
## AGDNSQSVY(ph)IPMSPGAHHFDLSLGY PSTTLPVHR-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 31758

Andromeda score: 172.11



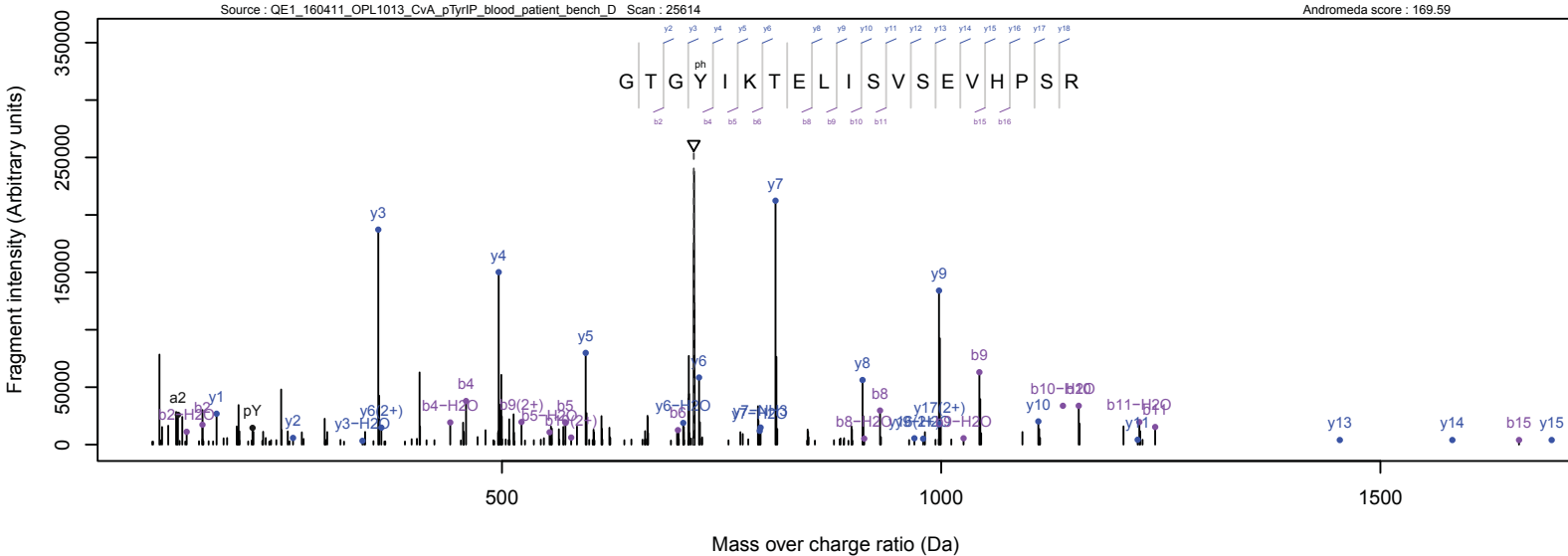




GTGY(ph)IKTELISVSEVHPSR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 25614

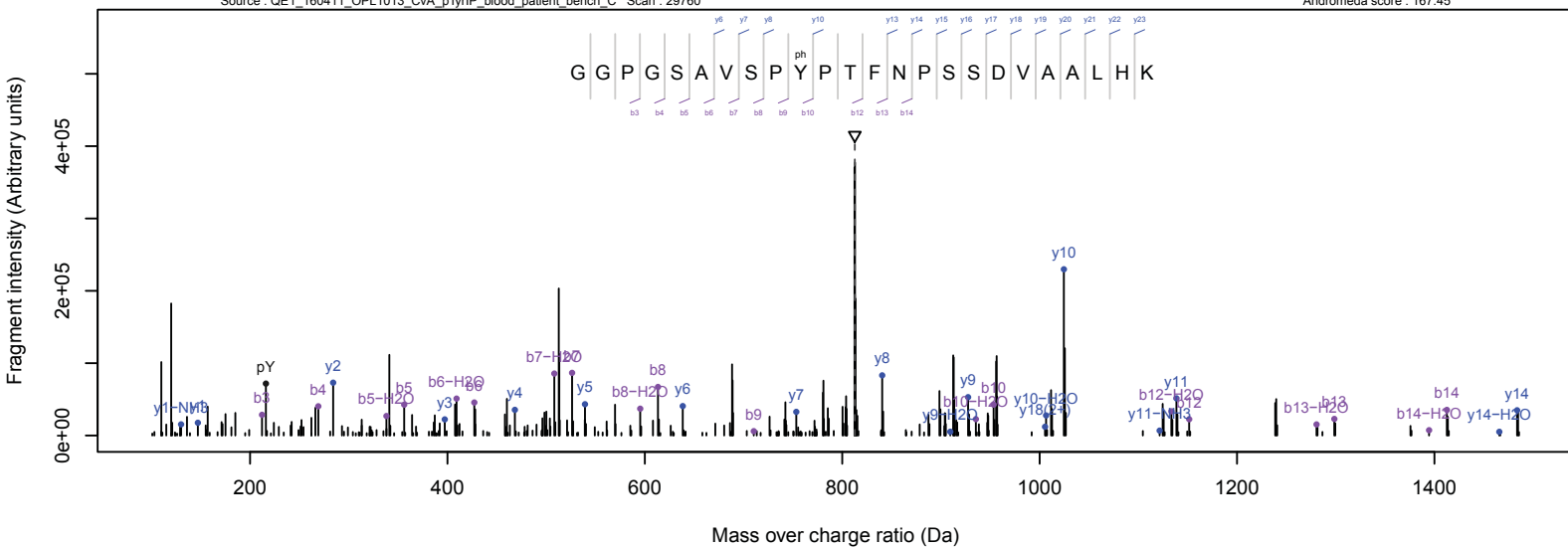
Andromeda score : 169.59



GGPGSAVSPY(ph)PTFNPSSDVAALHK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 29760

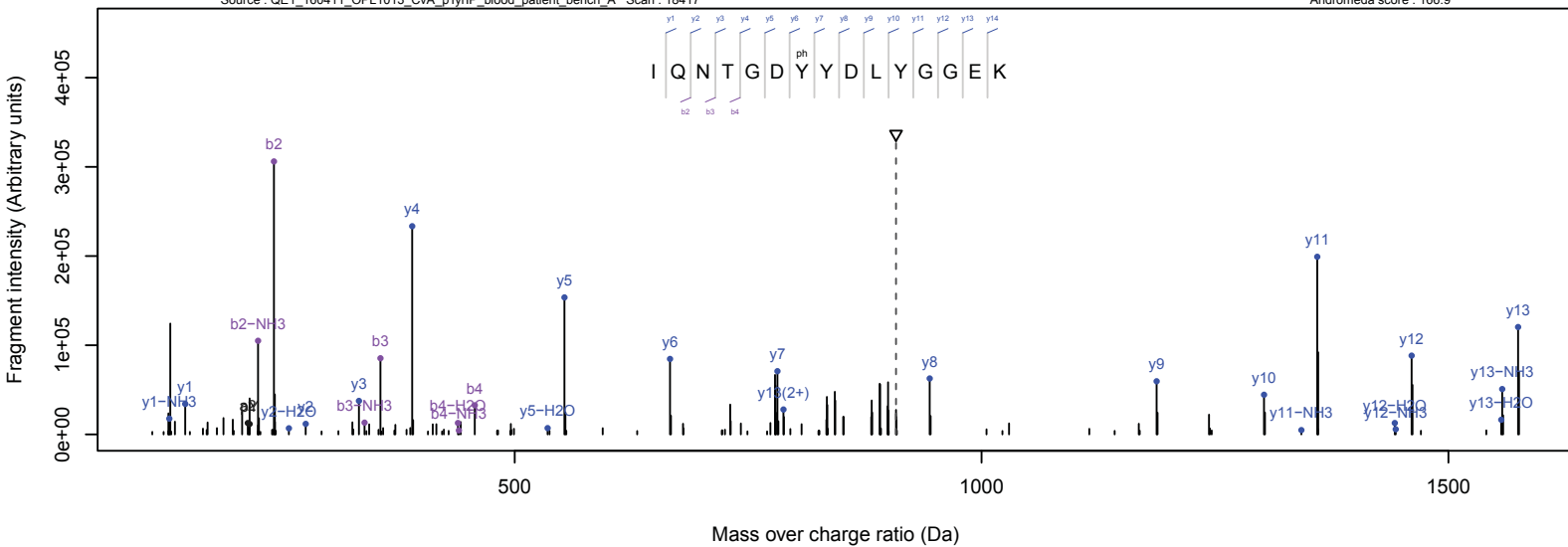
Andromeda score : 167.45



IQNTGDY(ph)YDLYGGEK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 18417

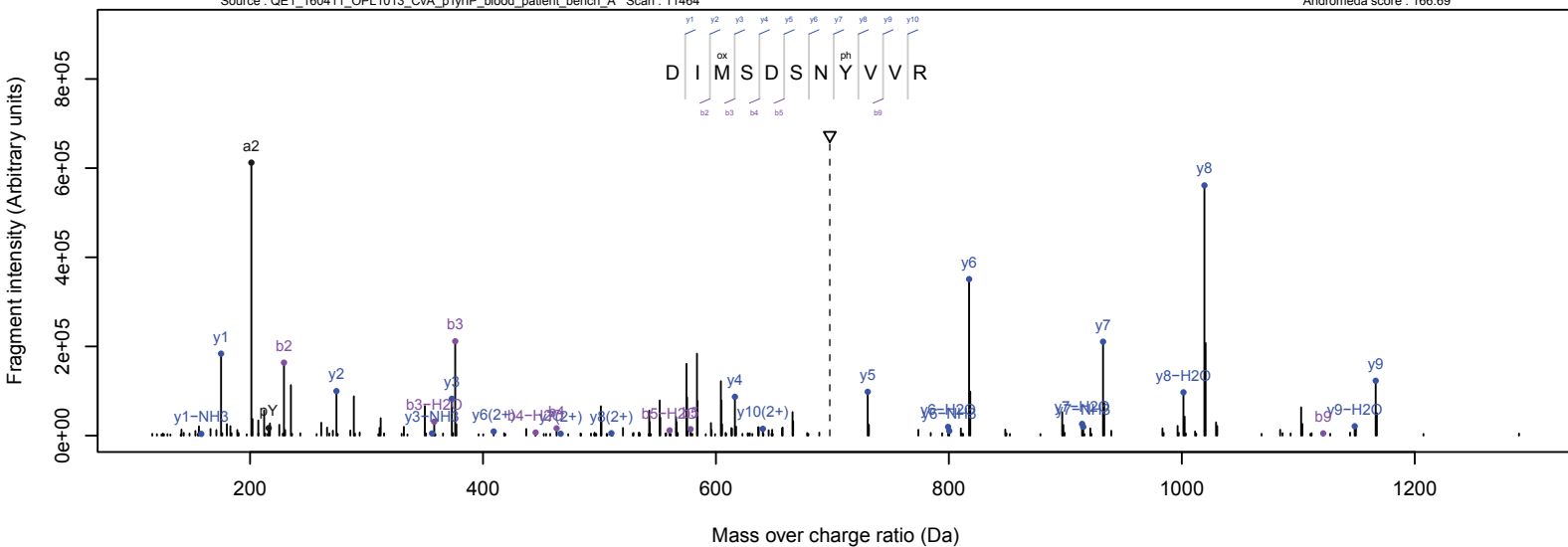
Andromeda score : 166.9



DIM( $\alpha$ )SDSNY(ph)VVR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 11464

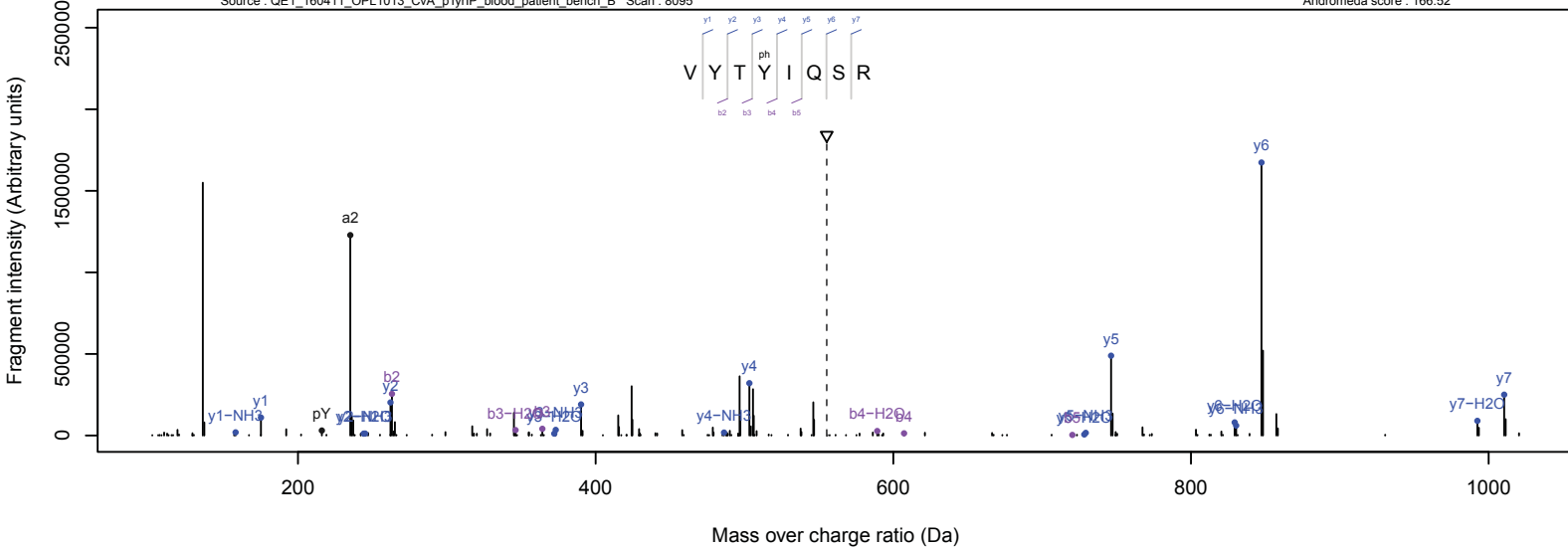
Andromeda score : 166.69



## VYTY(ph)QSR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 8095

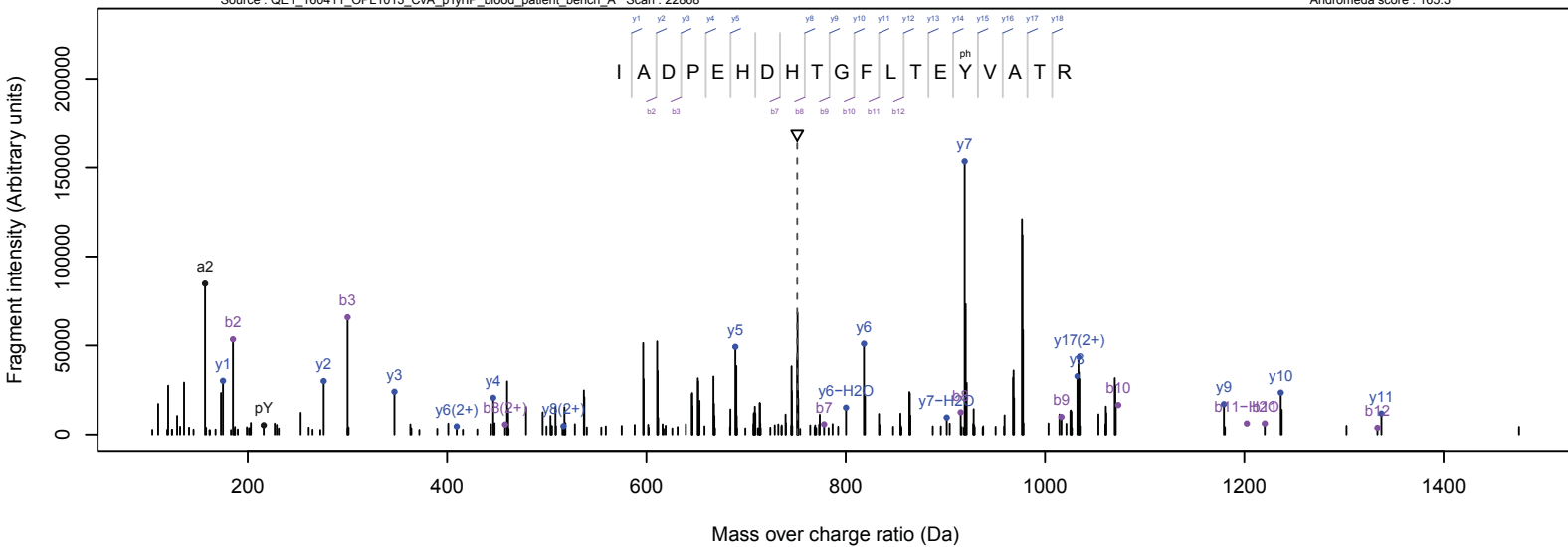
Andromeda score : 166.52



## IADPEHDHTGLTEY(ph)VATR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 22868

Andromeda score : 165.3

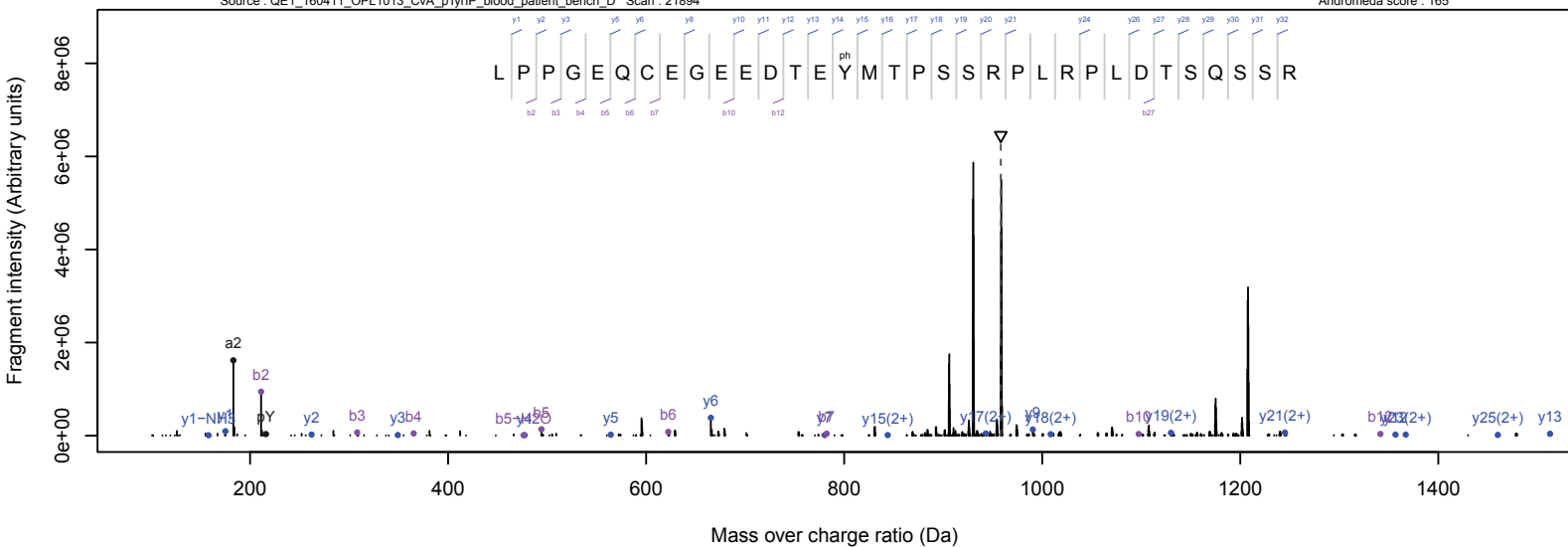




LPPGEQCEGEEDTEY(ph)MTPSSRPLRLDTSQSSR-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 21894

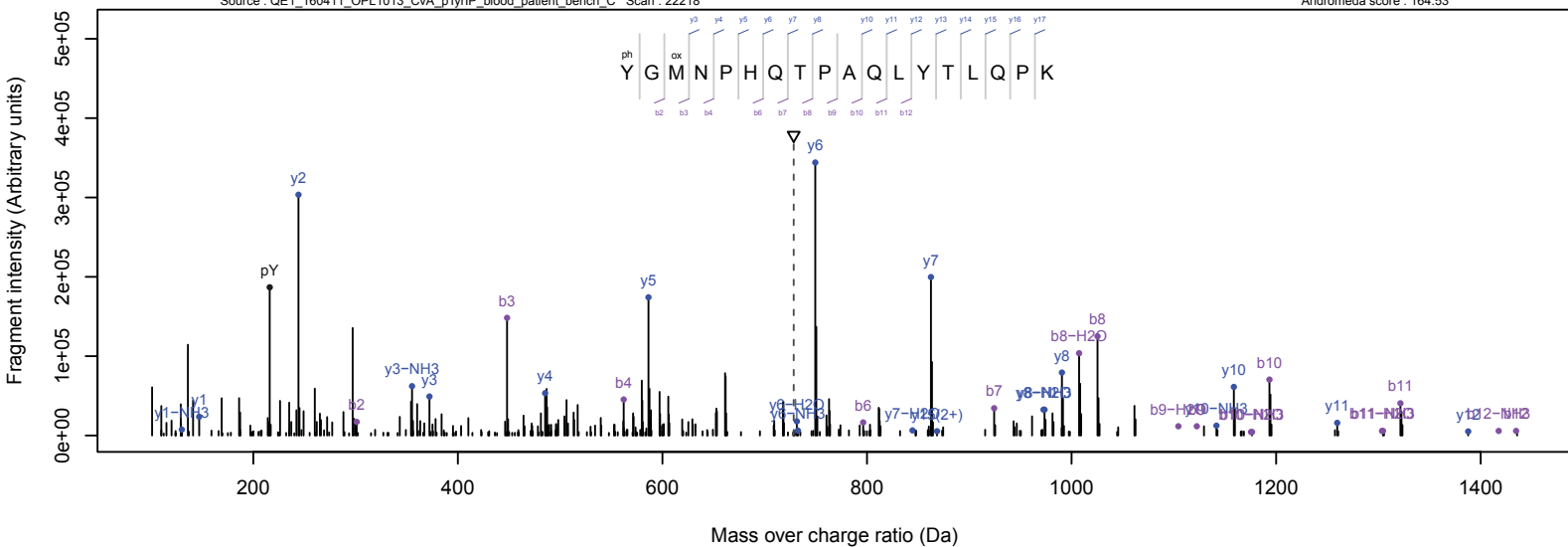
Andromeda score: 165



Y(ph)GM(ox)NPHQTPAQLYTLQPK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 22218

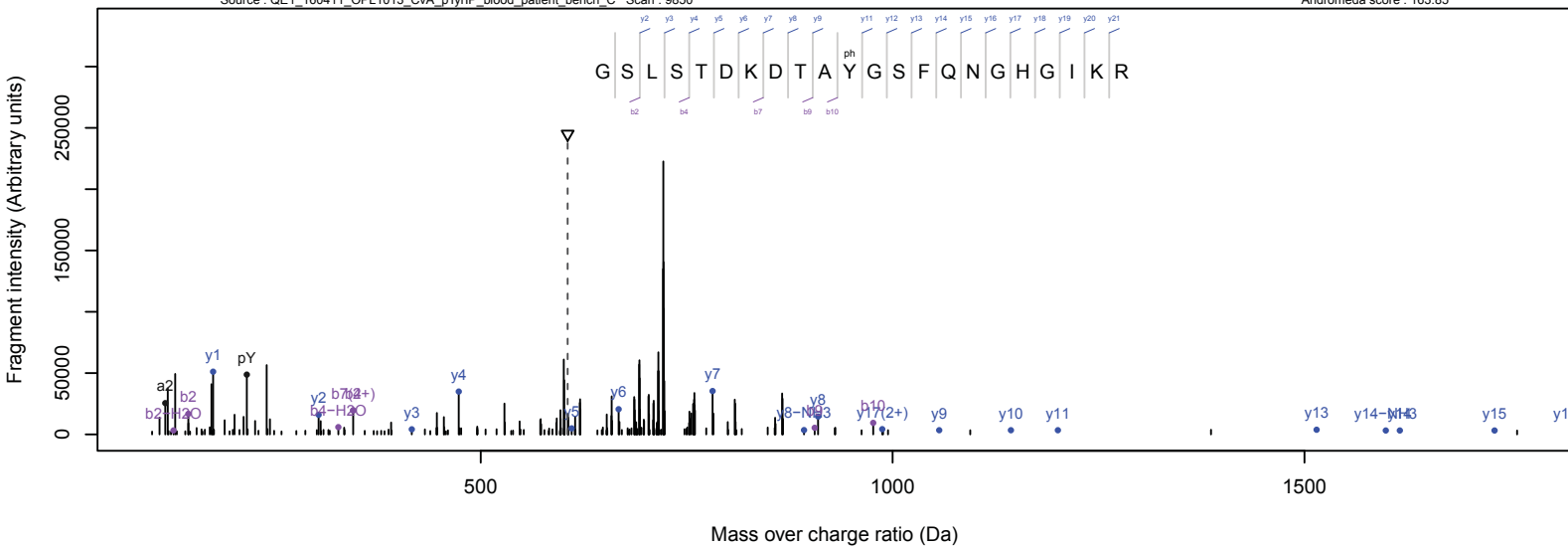
Andromeda score: 164.53



GSLSTDKDTAYG(SF)QNGHGIRK-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 9850

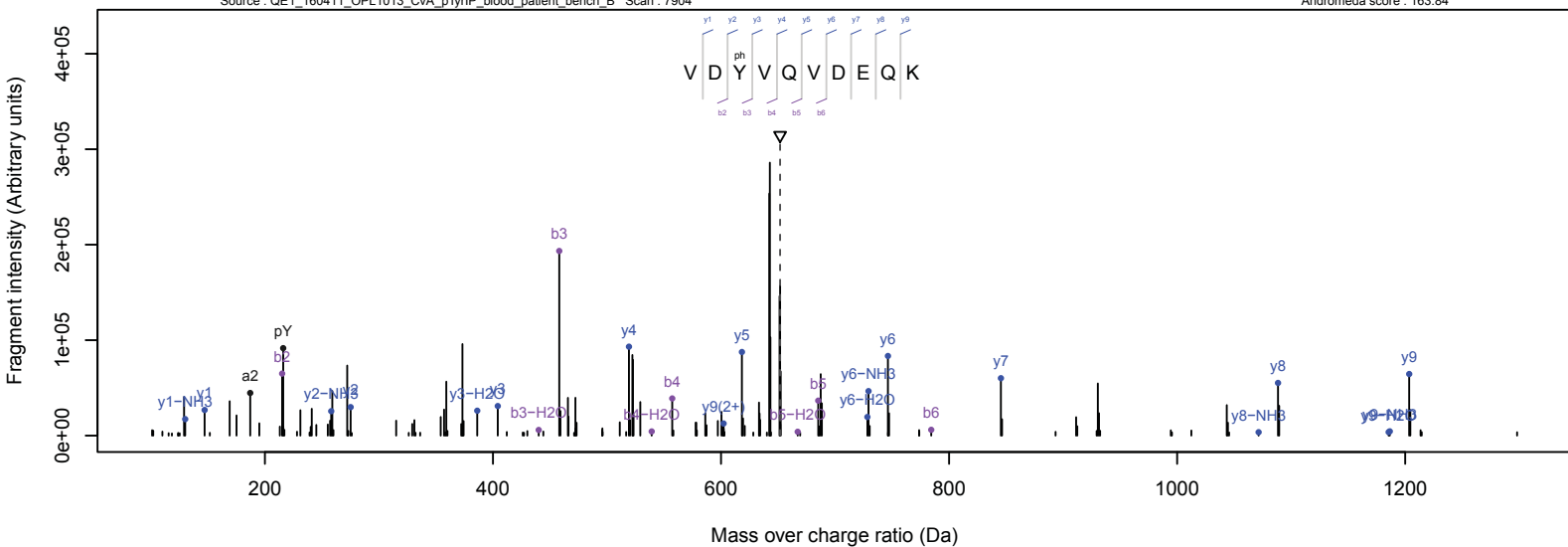
Andromeda score: 163.85



## VDY(ph)VQVDEQK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 7904

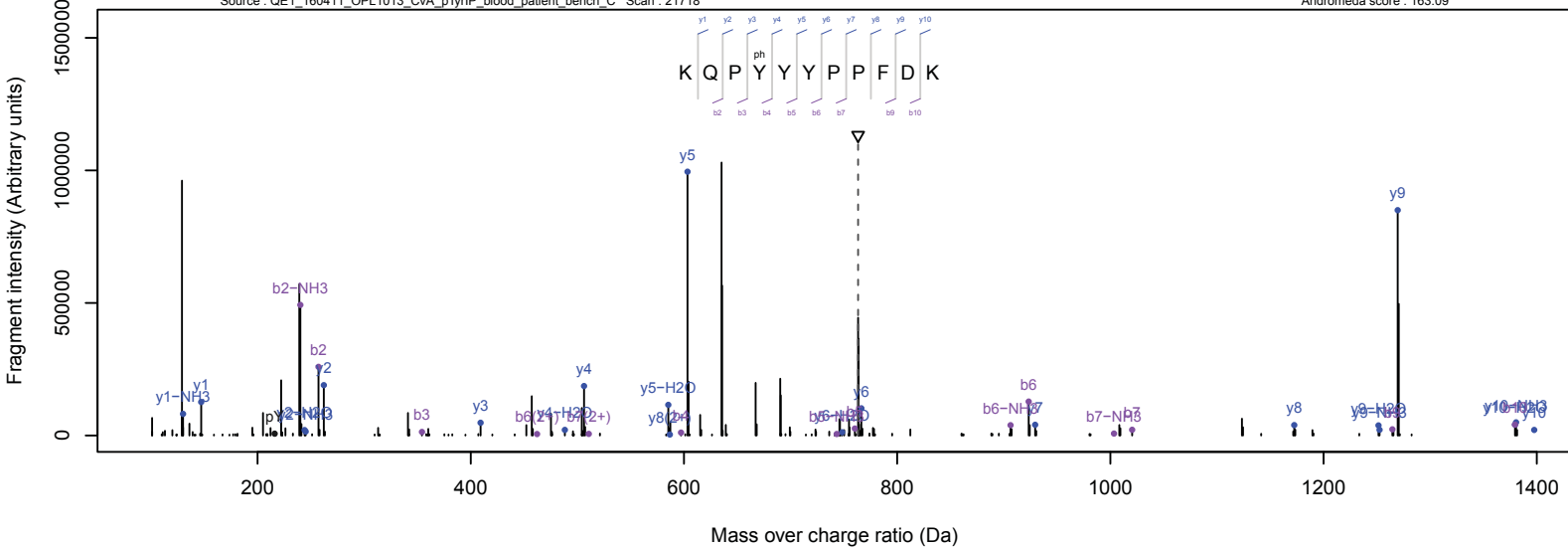
Andromeda score: 163.84



## KQPY(ph)YPPFDK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 21718

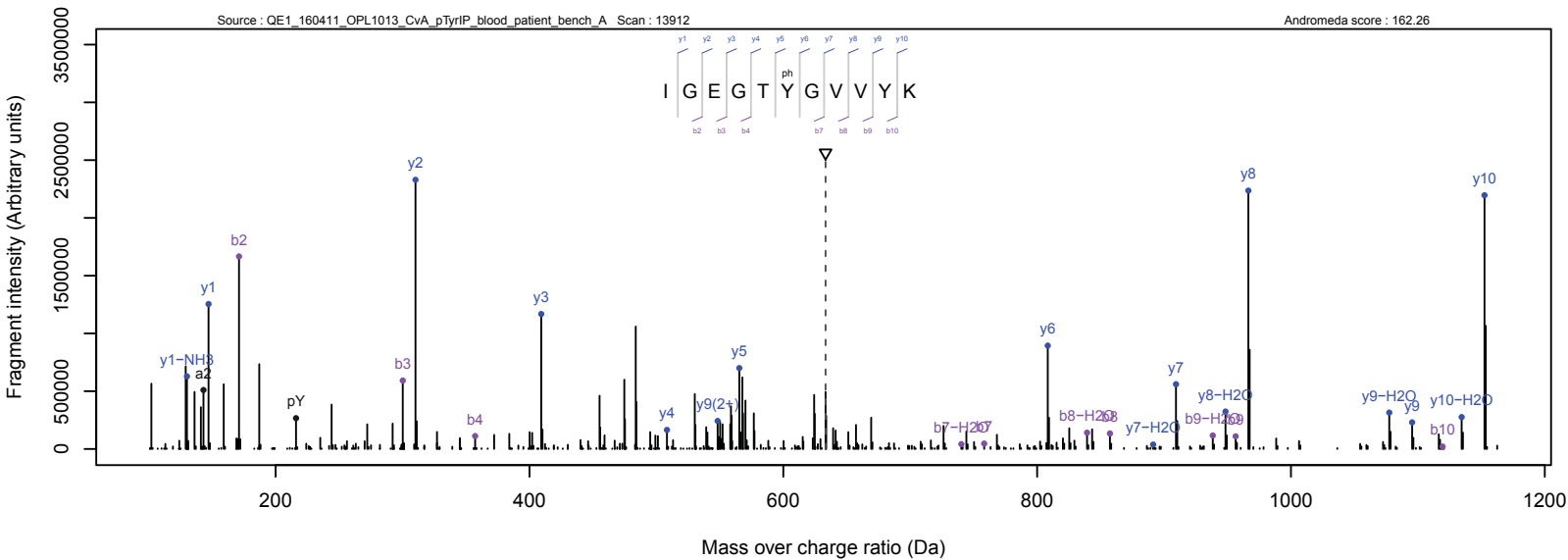
Andromeda score: 163.09

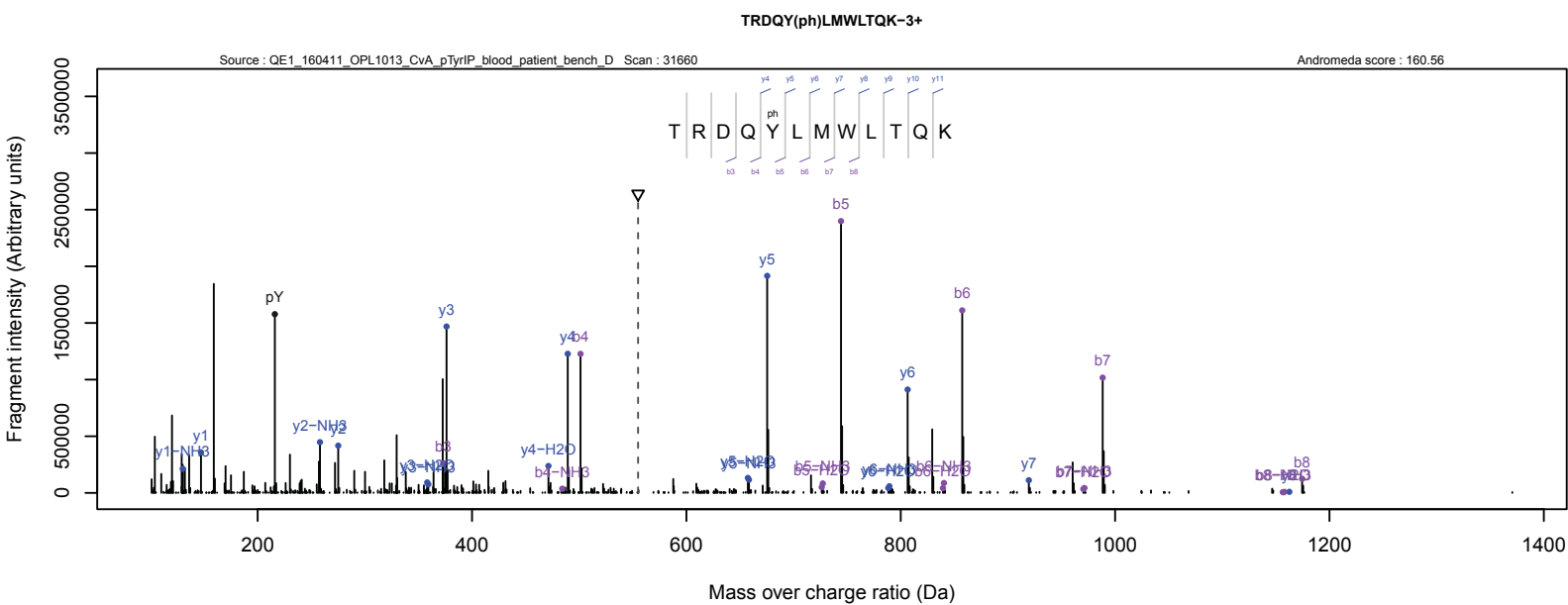
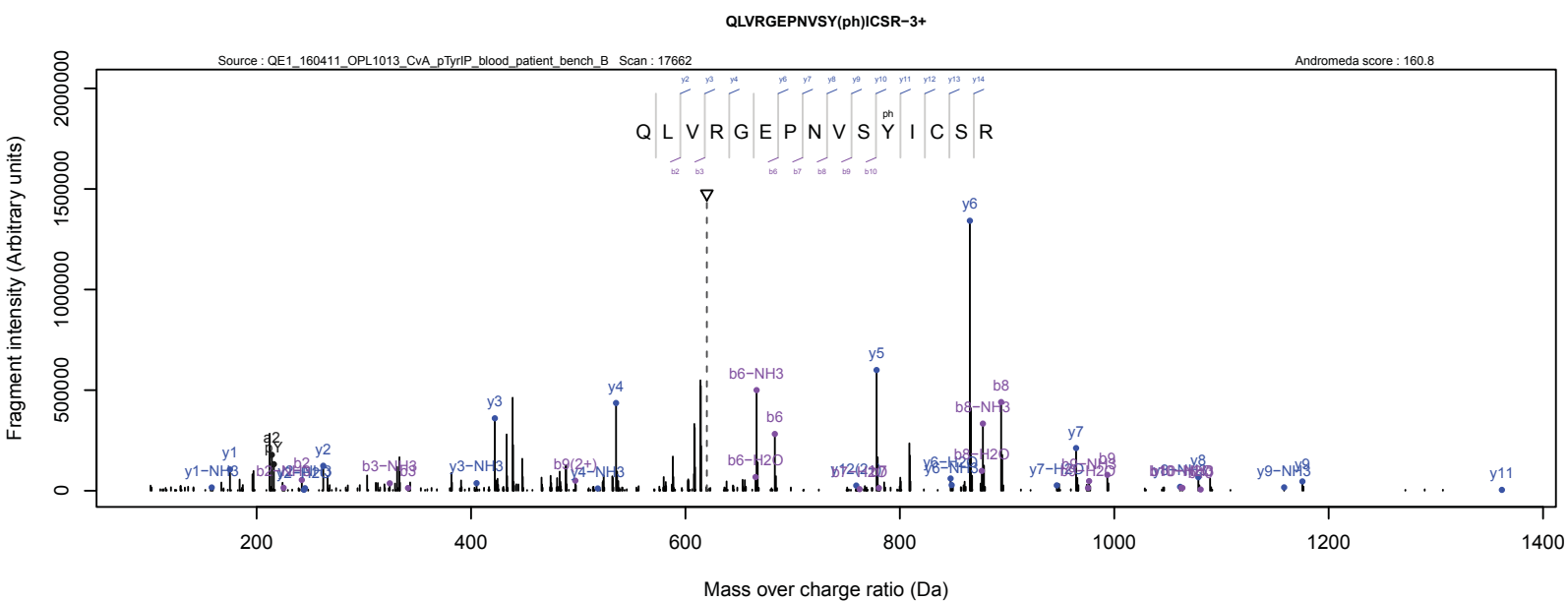
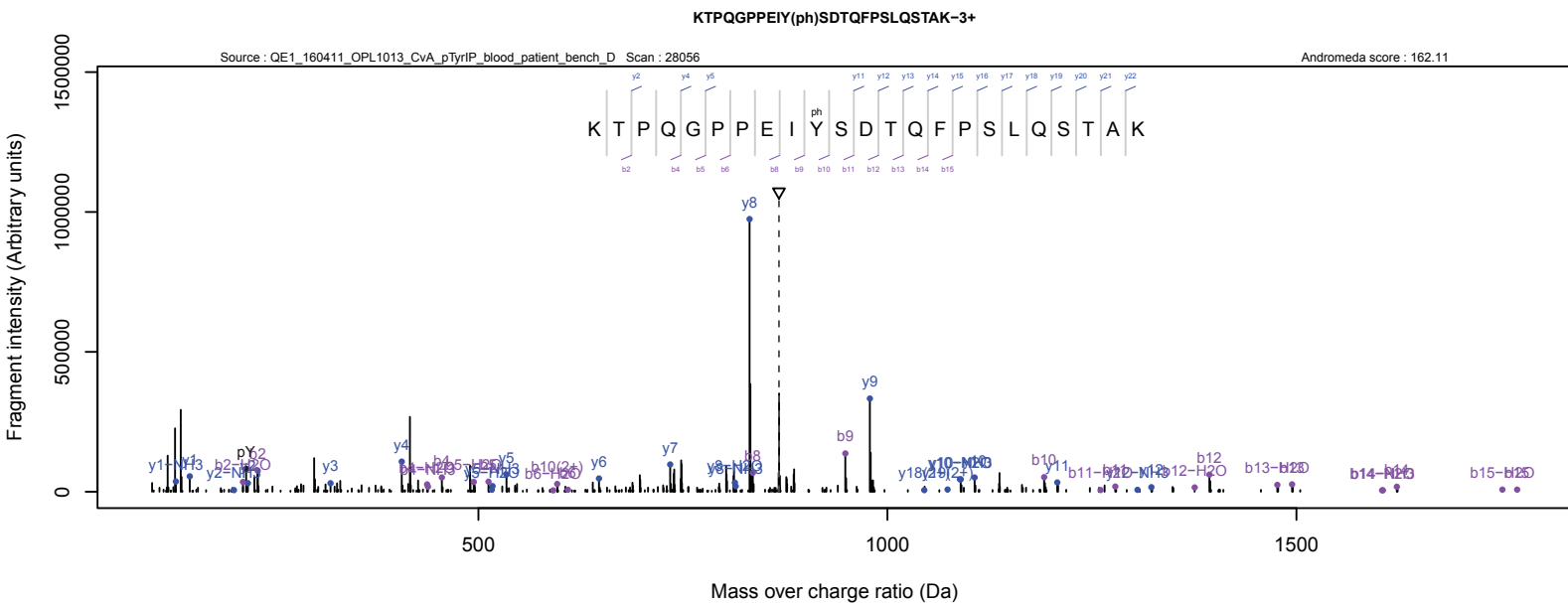


## IGEGTY(ph)GVVYK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 13912

Andromeda score: 162.26

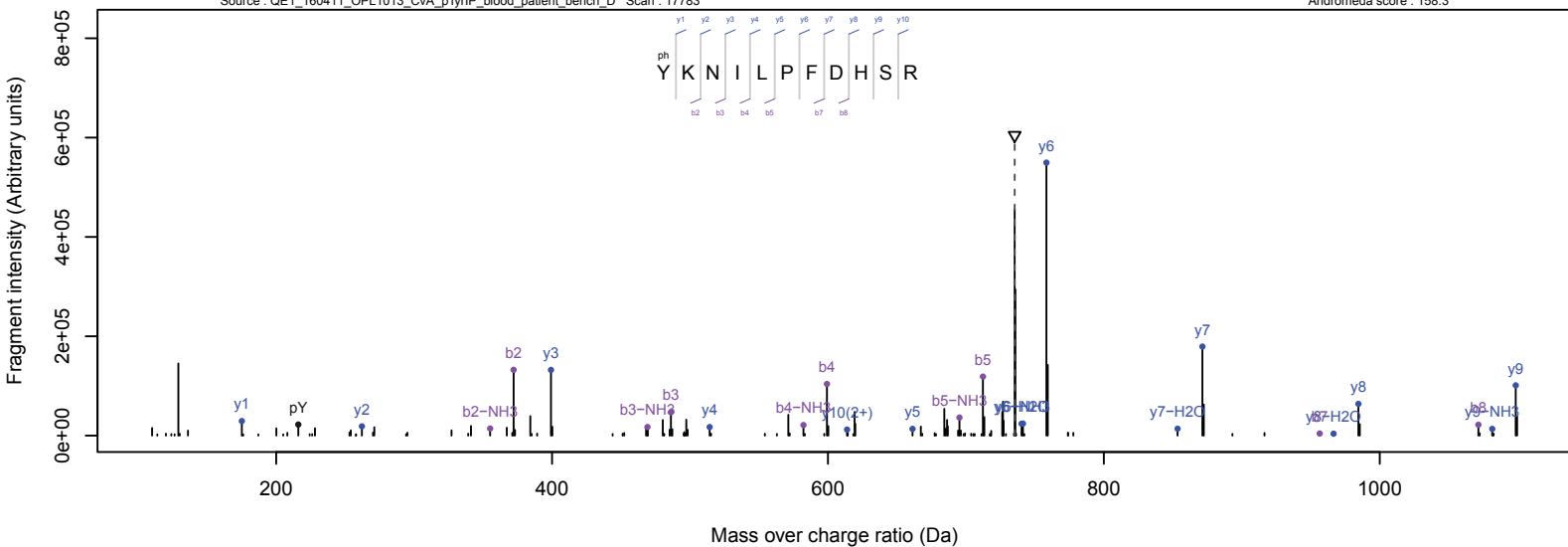




**Y(ph)KNILPFDHSR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17783

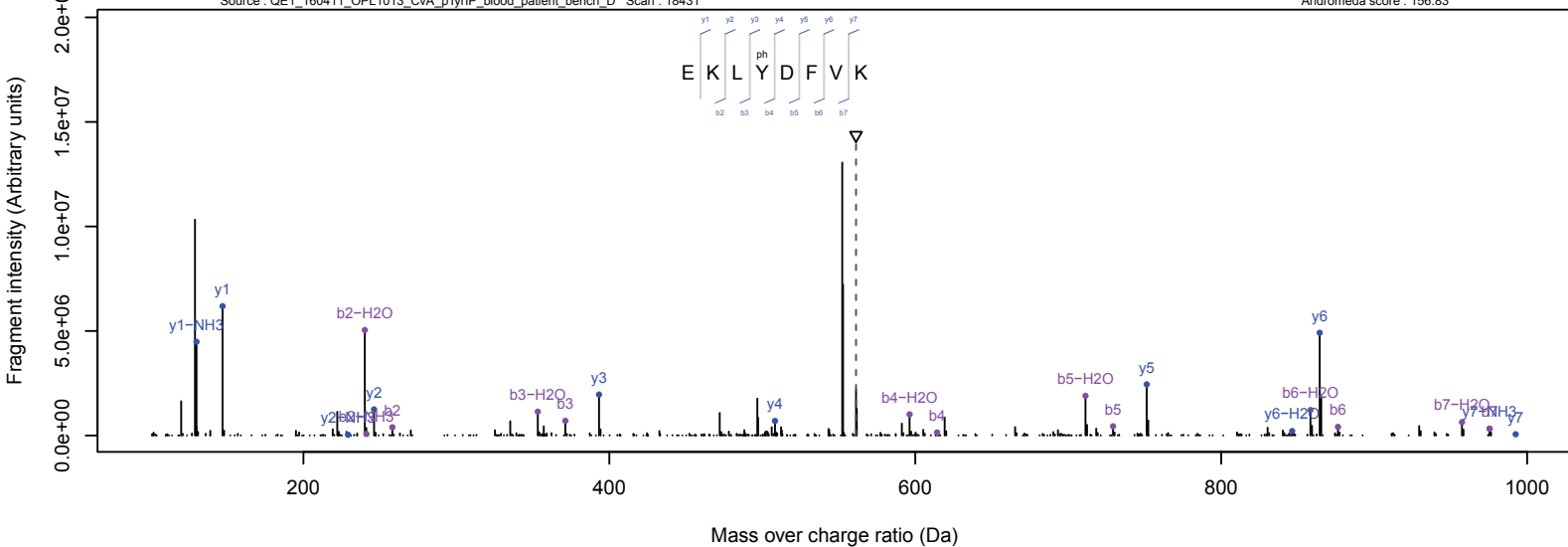
Andromeda score : 158.3



**EKLY(ph)DFVK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 18431

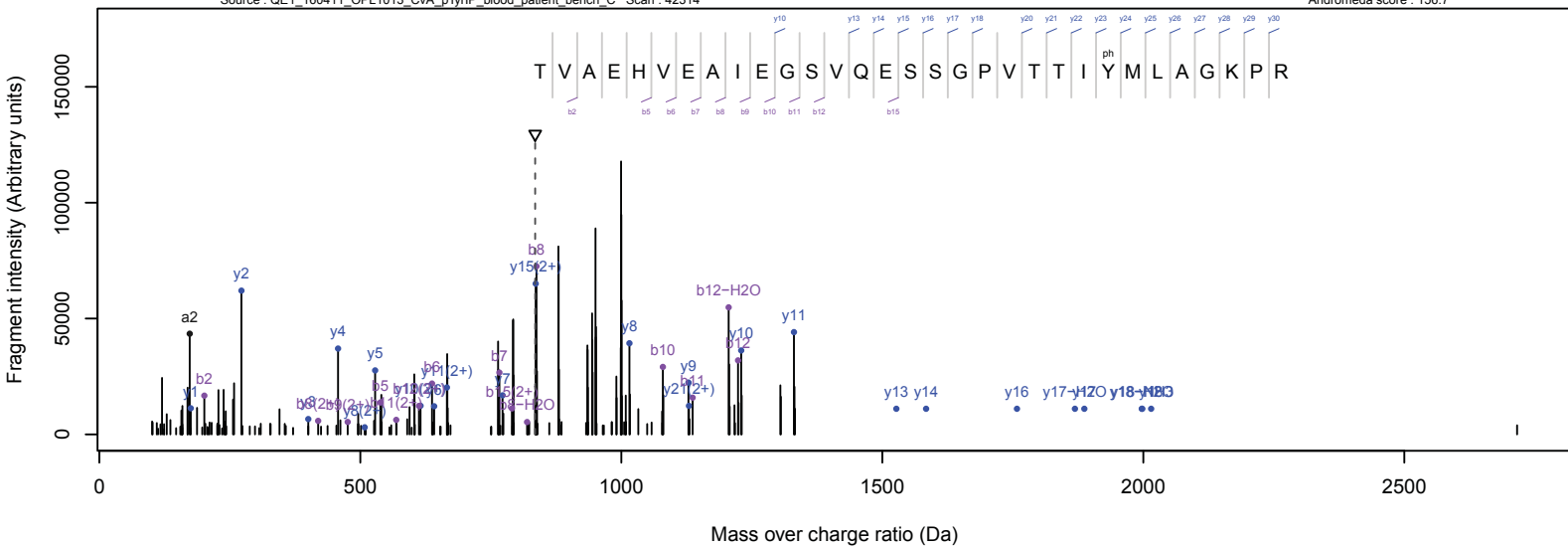
Andromeda score : 156.83



**TVAEHVEAIEGSGVPTTIY(ph)MLAGKPR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 42314

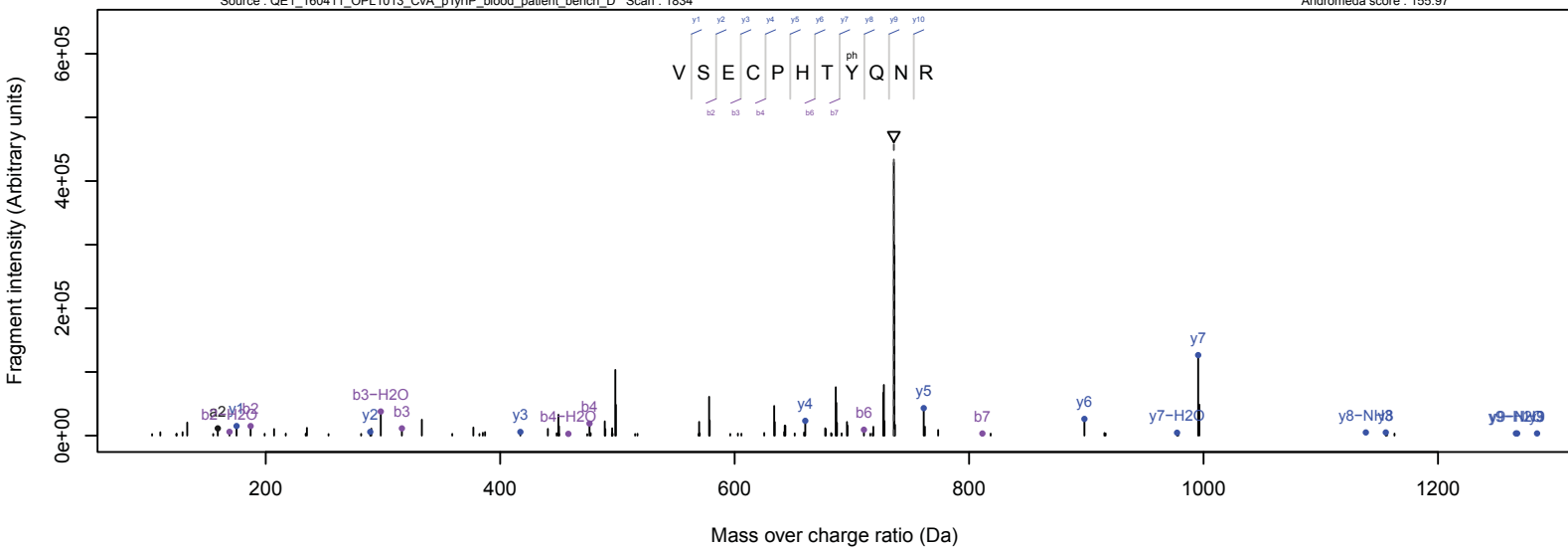
Andromeda score : 156.7



VSECPHTY(ph)QNR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 1834

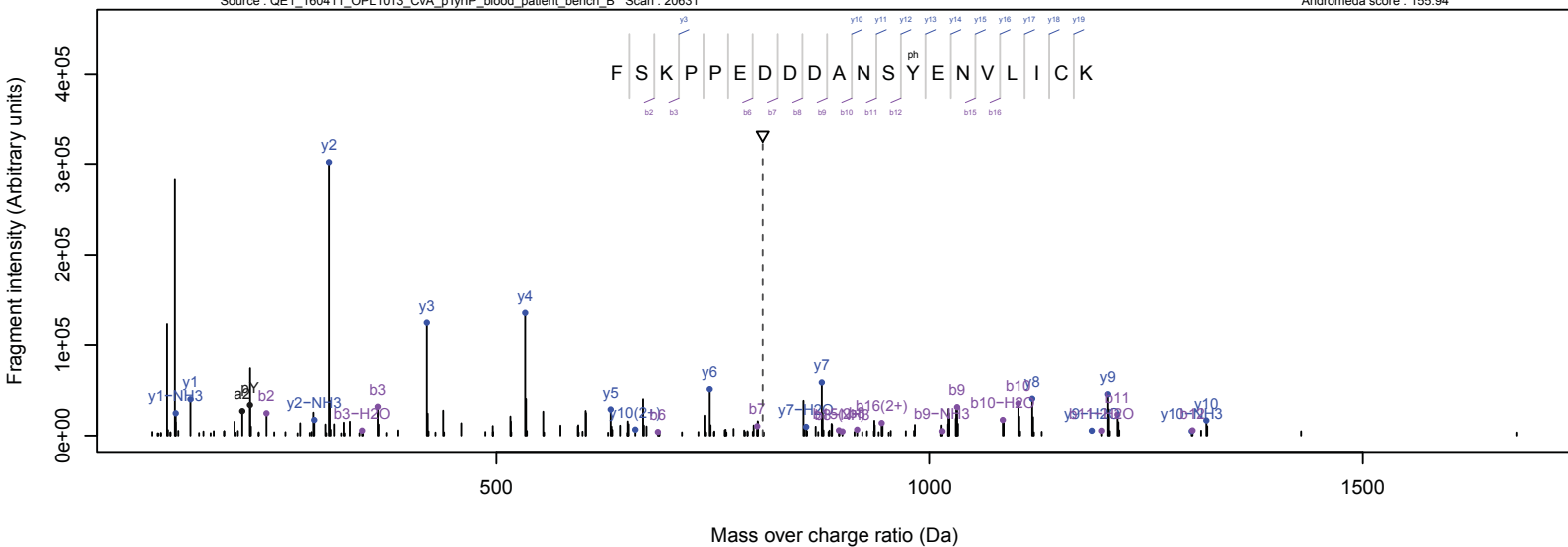
Andromeda score : 155.97



FSKPPEDDDANSY(ph)ENVLICK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 20631

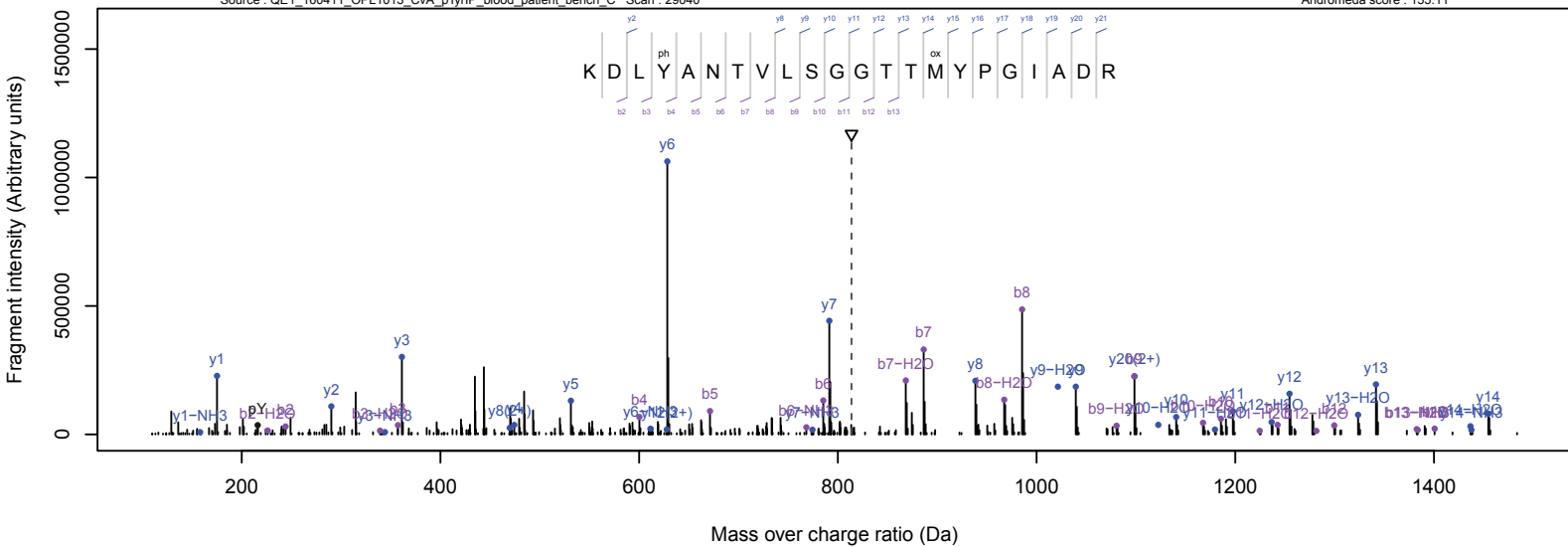
Andromeda score : 155.94



KDLY(ph)ANTVLSGGTTMYPGIADR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 29040

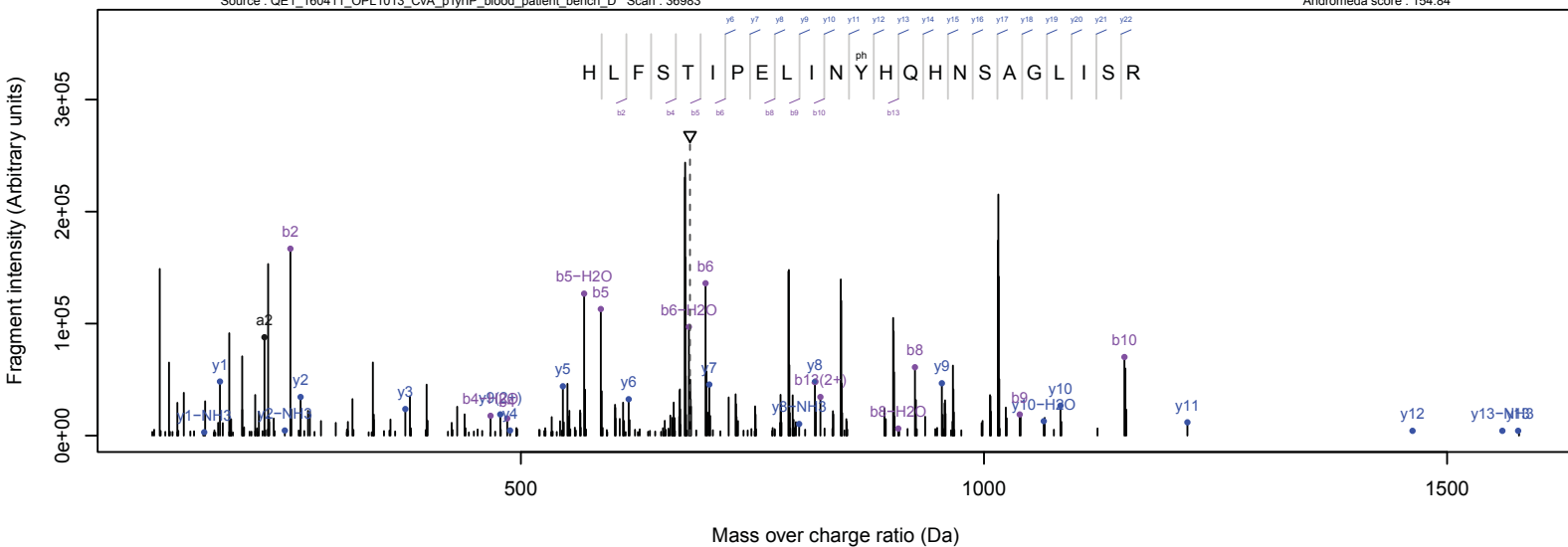
Andromeda score : 155.11



HLFSTIPELINY<sup>ph</sup>HQHNSAGLISR-4+

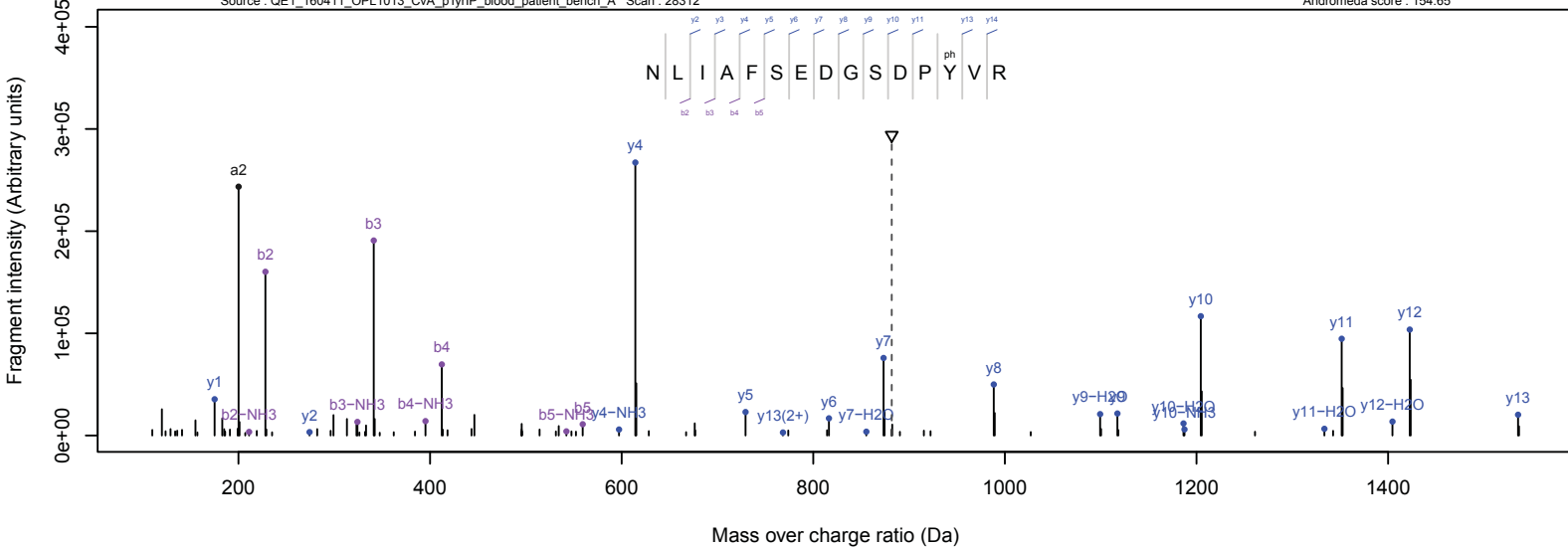
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 36983

Andromeda score: 154.84

NLIAFSEGDSPY<sup>ph</sup>VR-2+

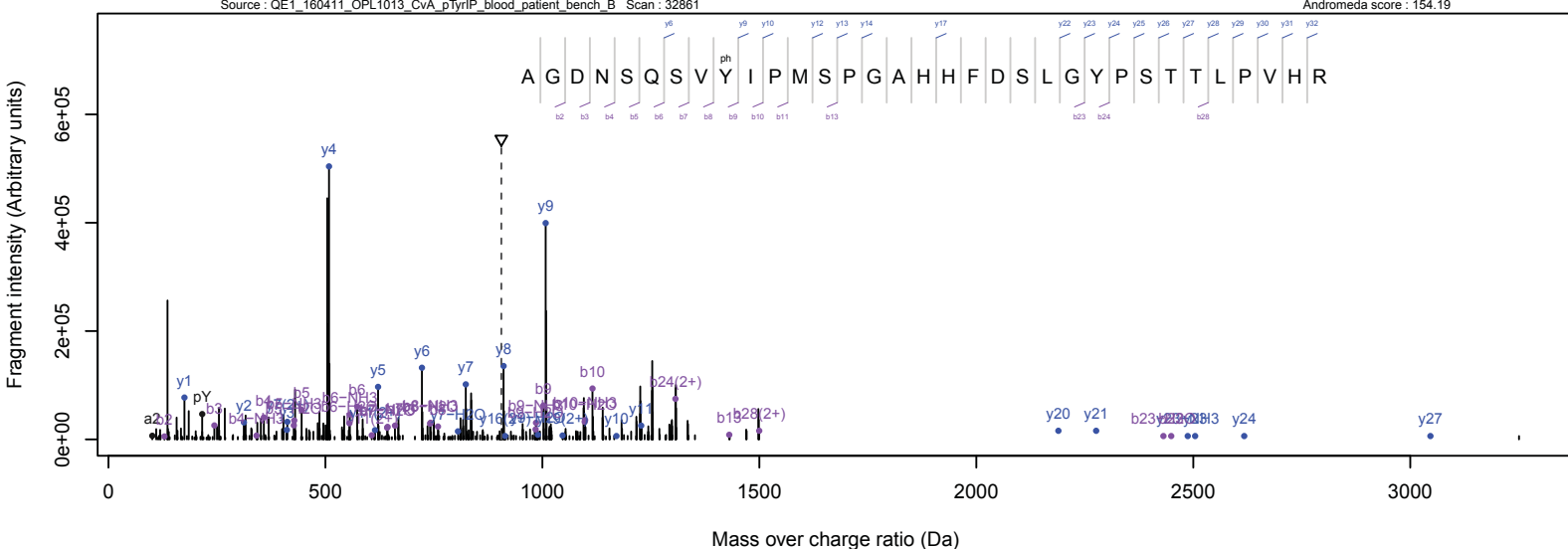
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 28312

Andromeda score: 154.65

AGDNSQSVY<sup>ph</sup>IPMSPGAHHFDSLGYPTTLPVHR-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 32861

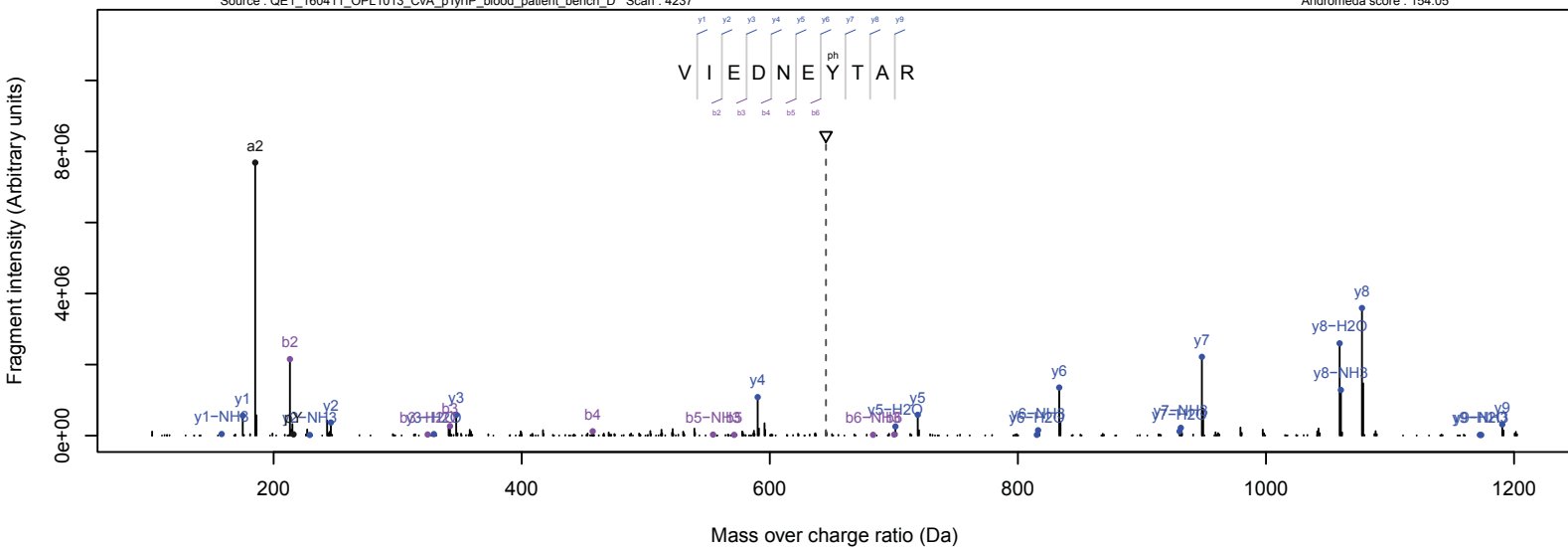
Andromeda score: 154.19



VIEDNEY(ph)TAR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 4237

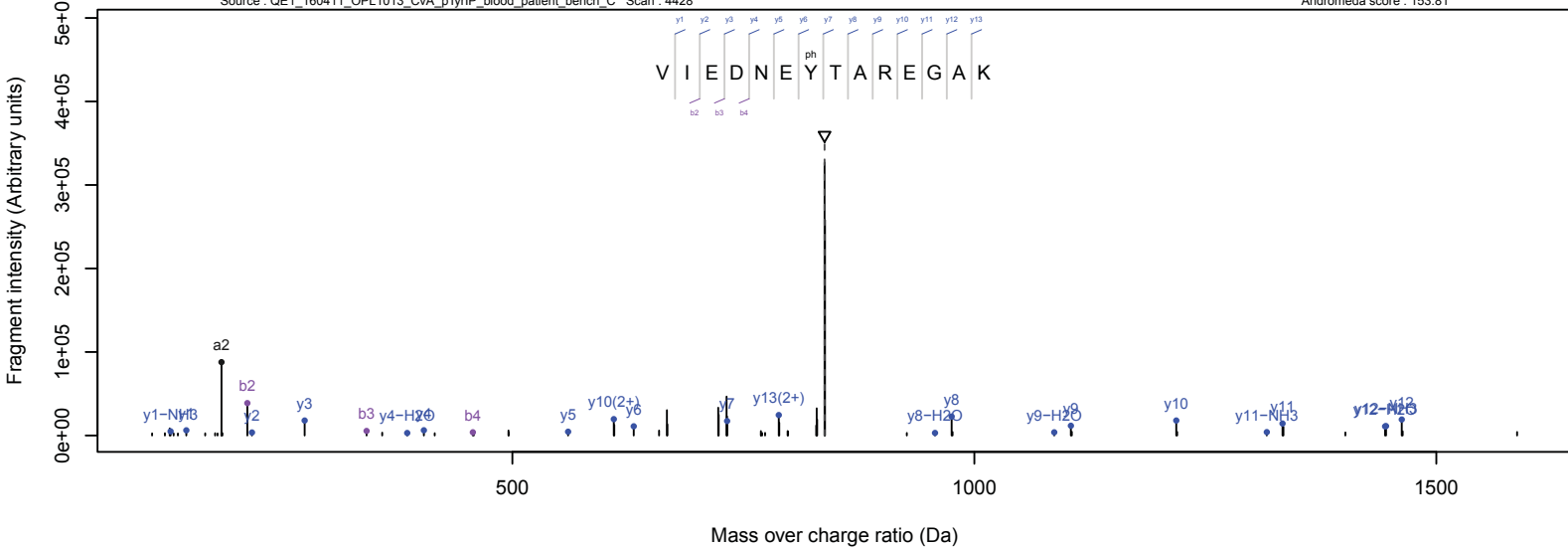
Andromeda score: 154.05



VIEDNEY(ph)TAREGAK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 4428

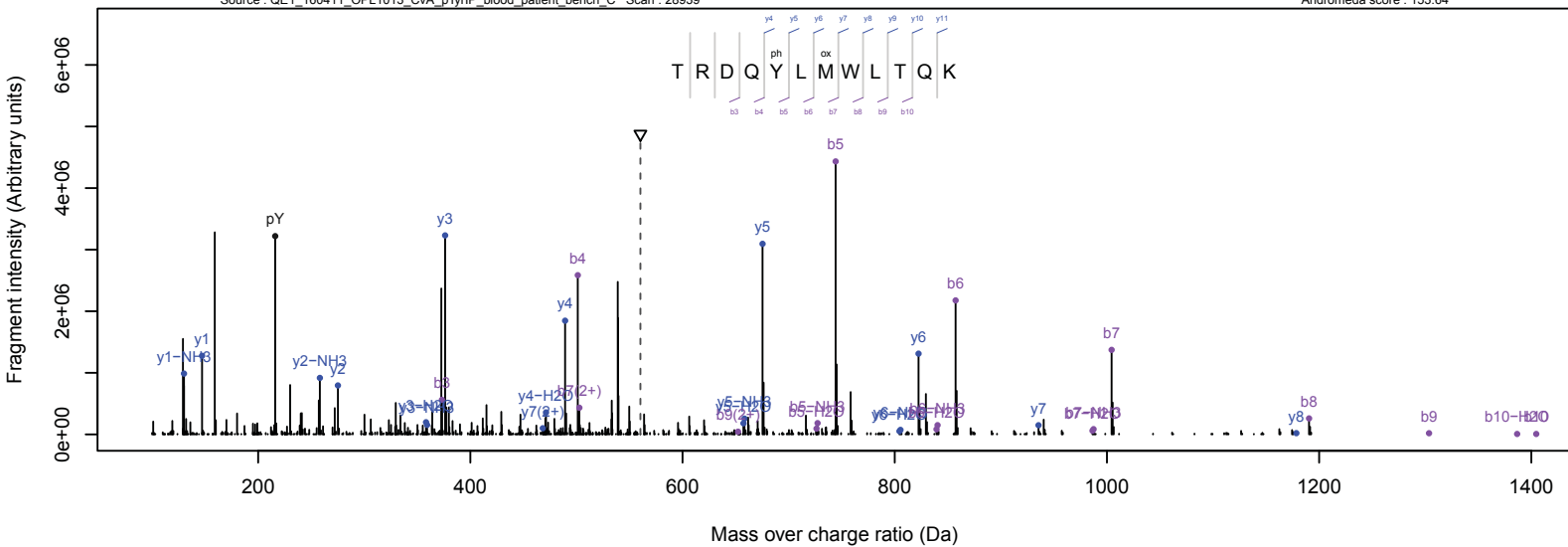
Andromeda score: 153.81

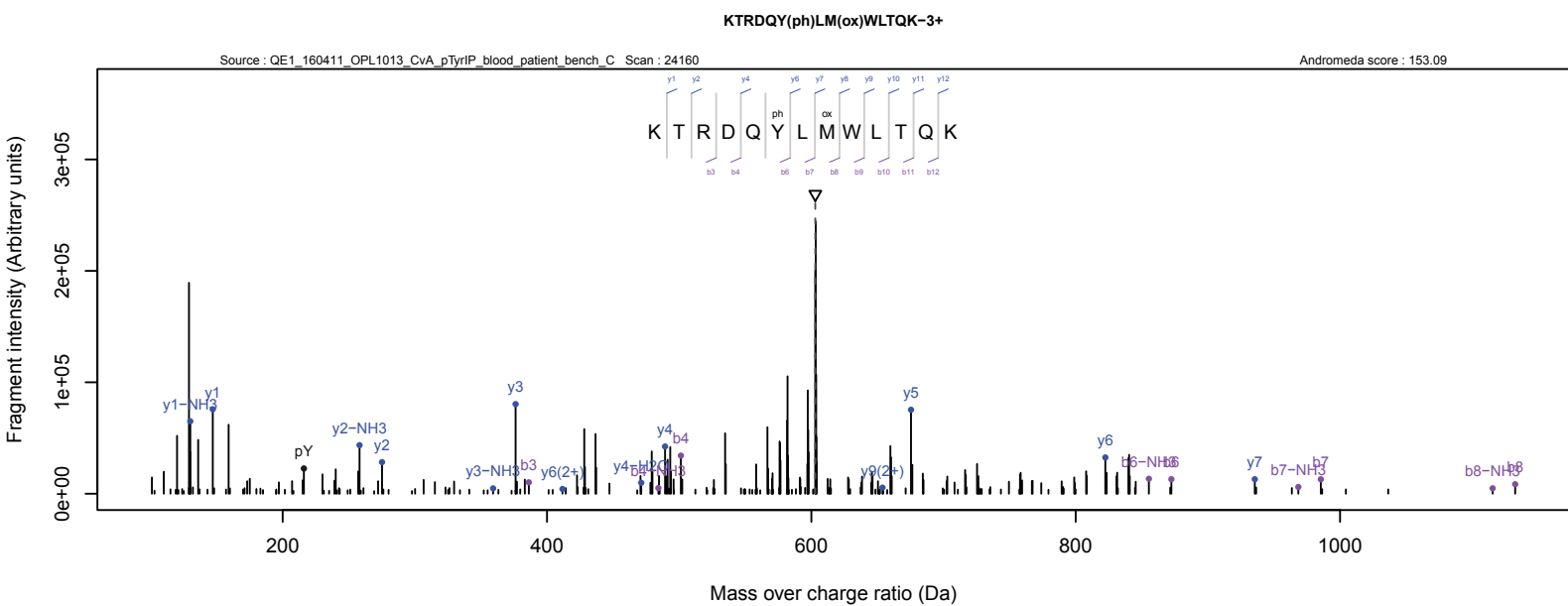
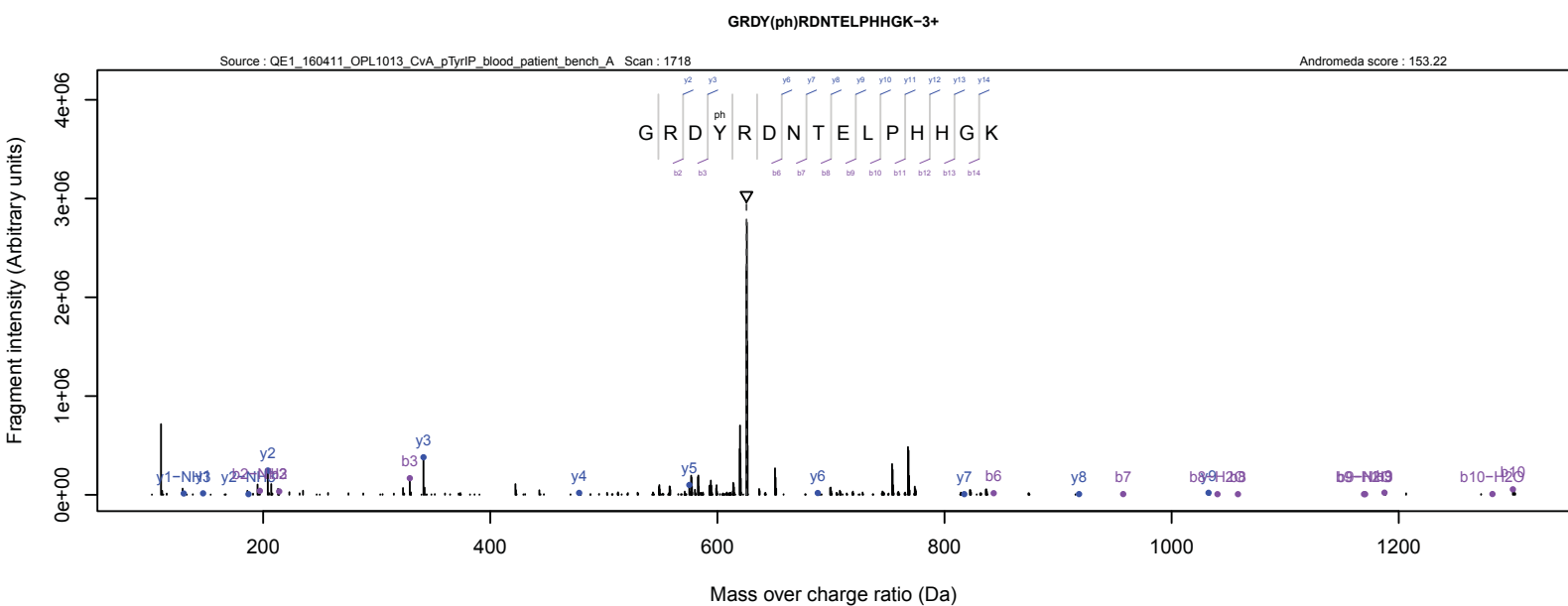
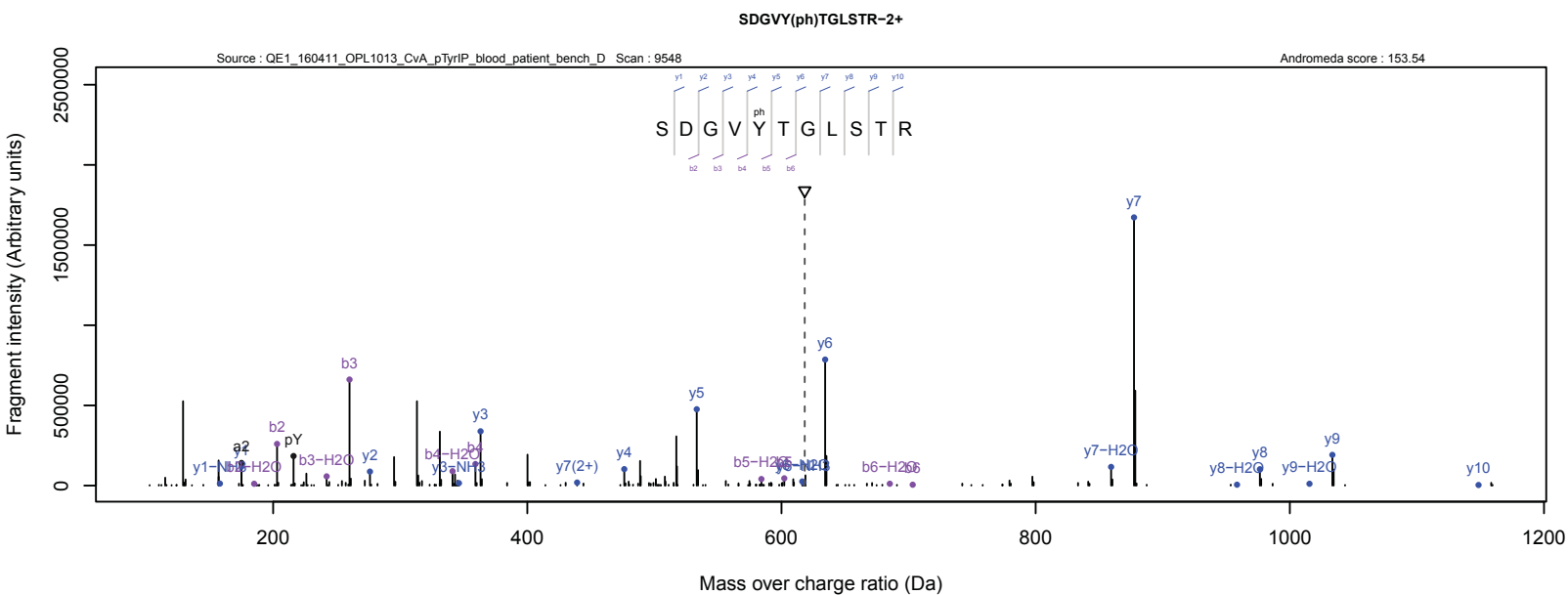


TRDQY(ph)LMWLTQK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 28939

Andromeda score: 153.64



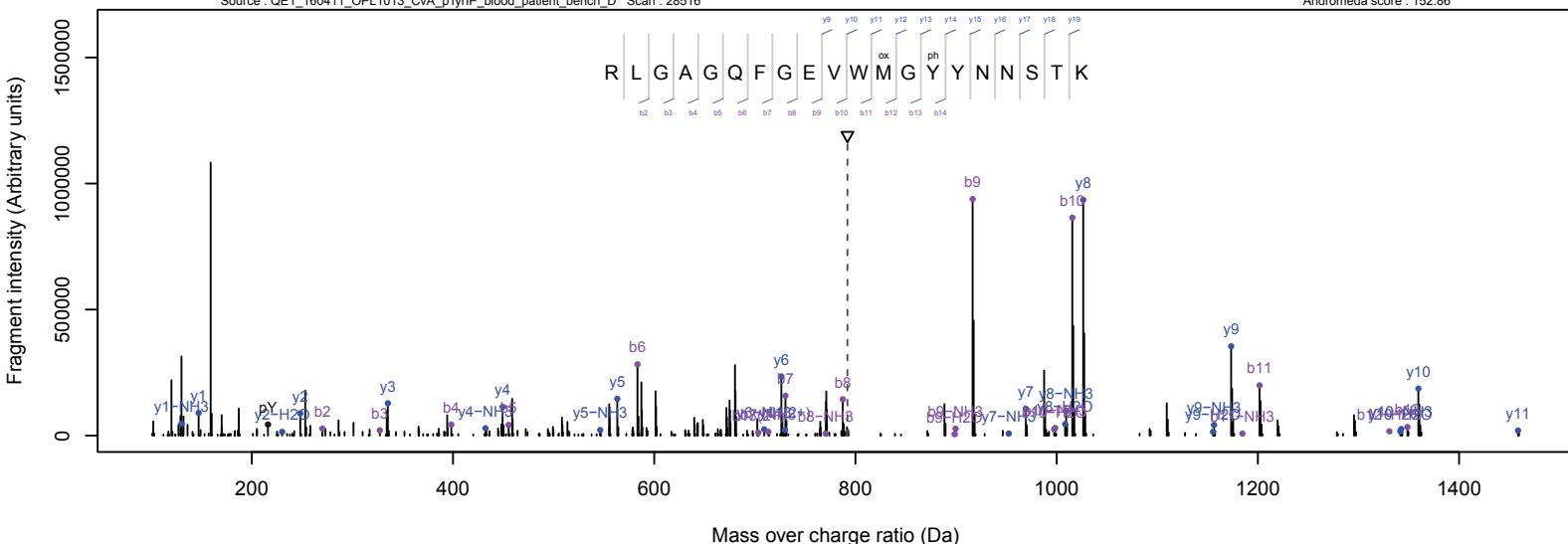




RLGAGQFGEVWMGYPHNNSTK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28516

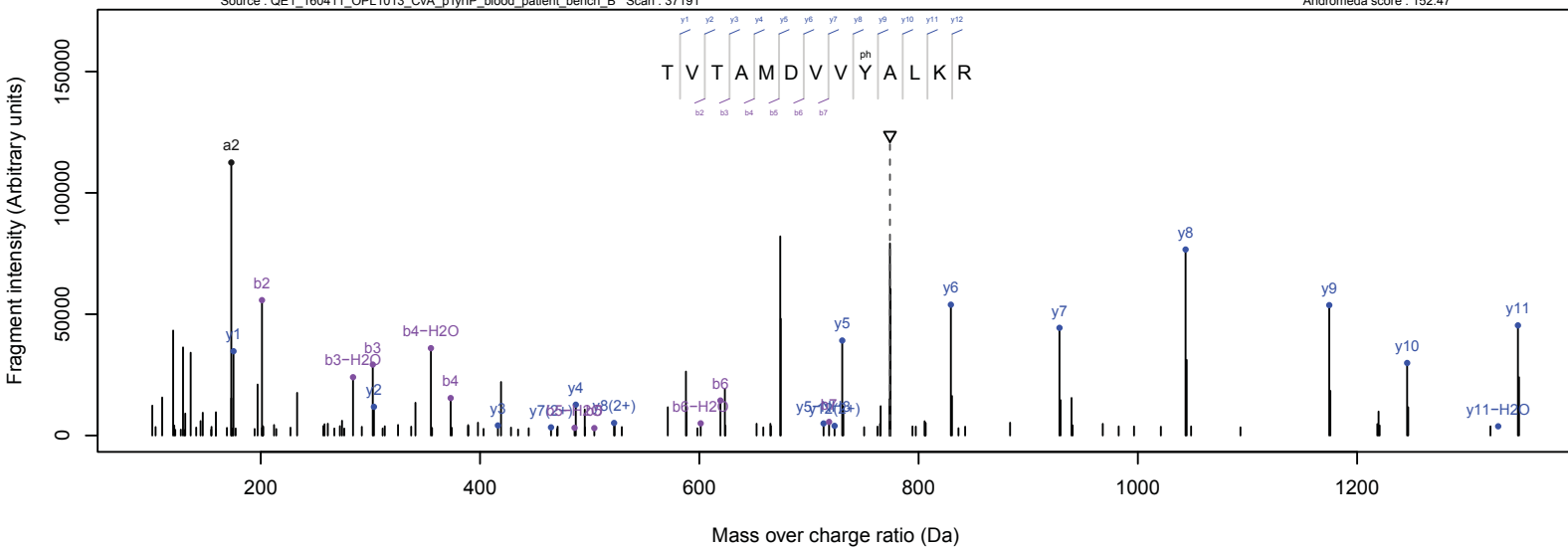
Andromeda score : 152.86



TVTAMDVVYPHALKR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 37191

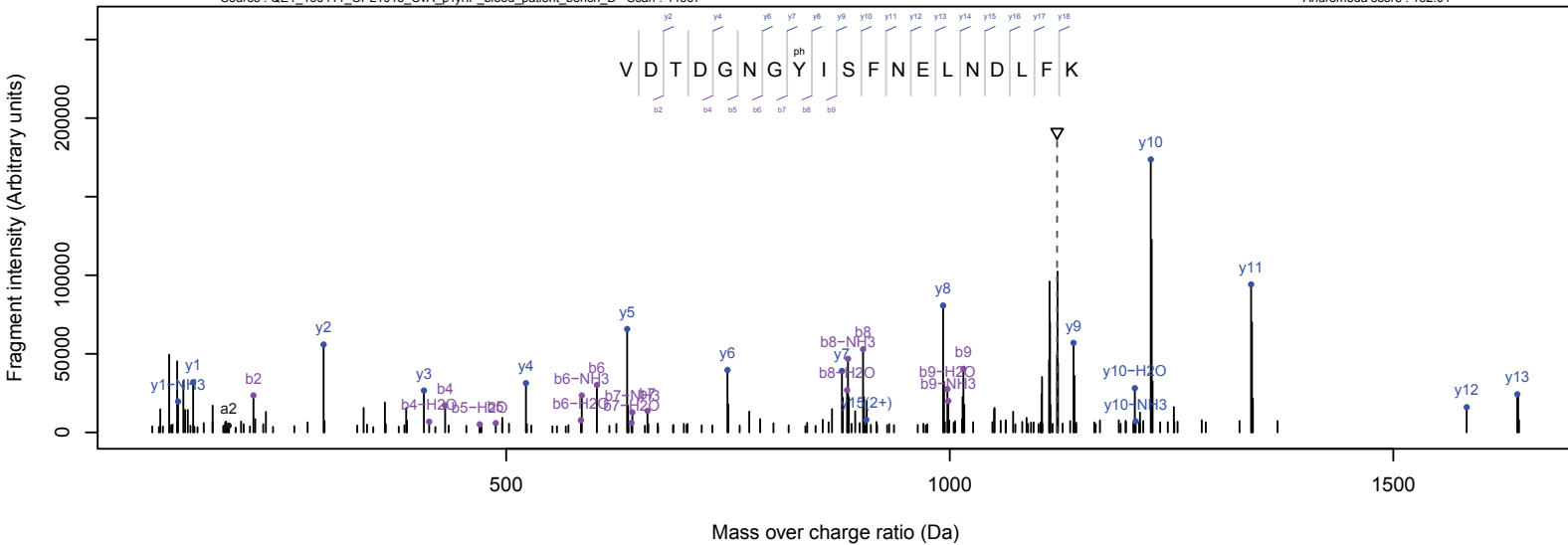
Andromeda score : 152.47



VDTDNGYIPHSFNELNDFK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 44367

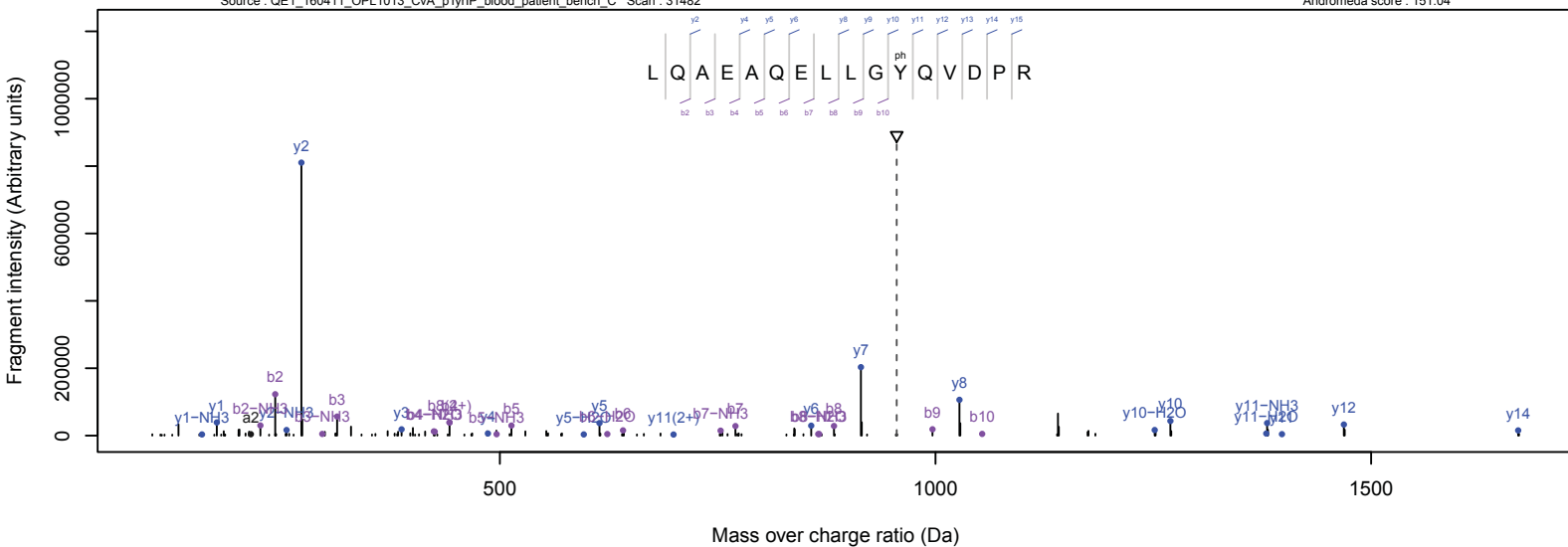
Andromeda score : 152.01



LQAEAQELLY(ph)QVDPR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 31482

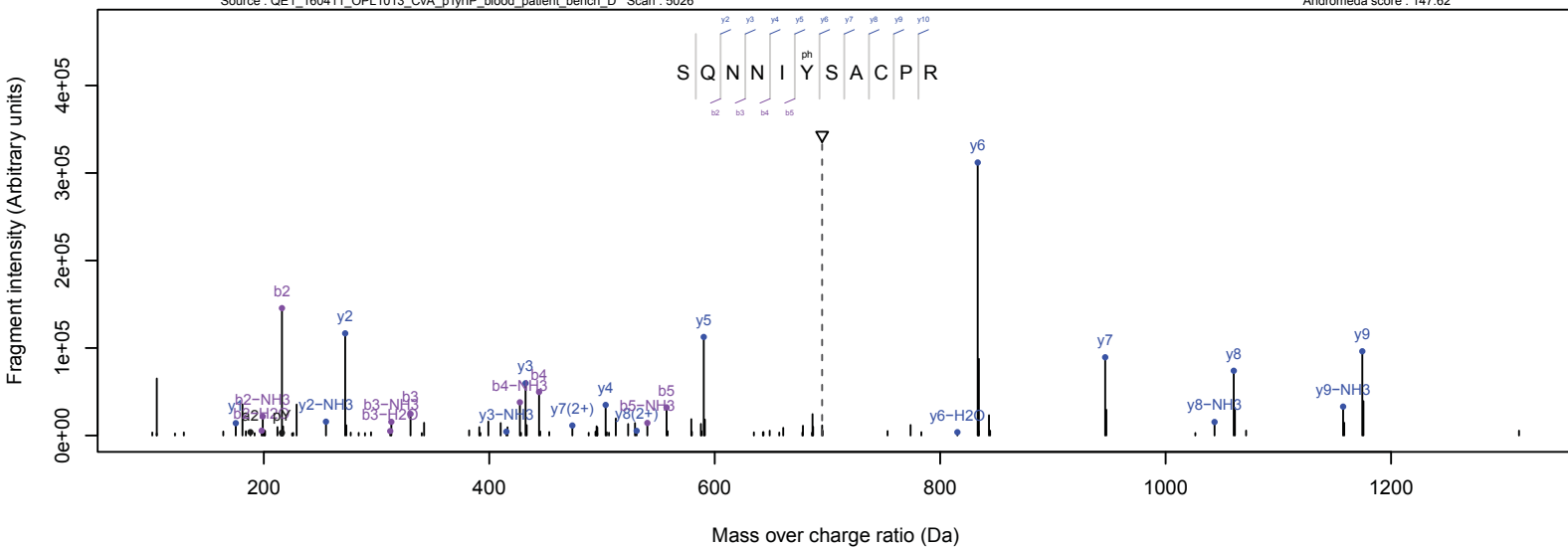
Andromeda score: 151.04



SQNNIY(ph)SACPR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 5026

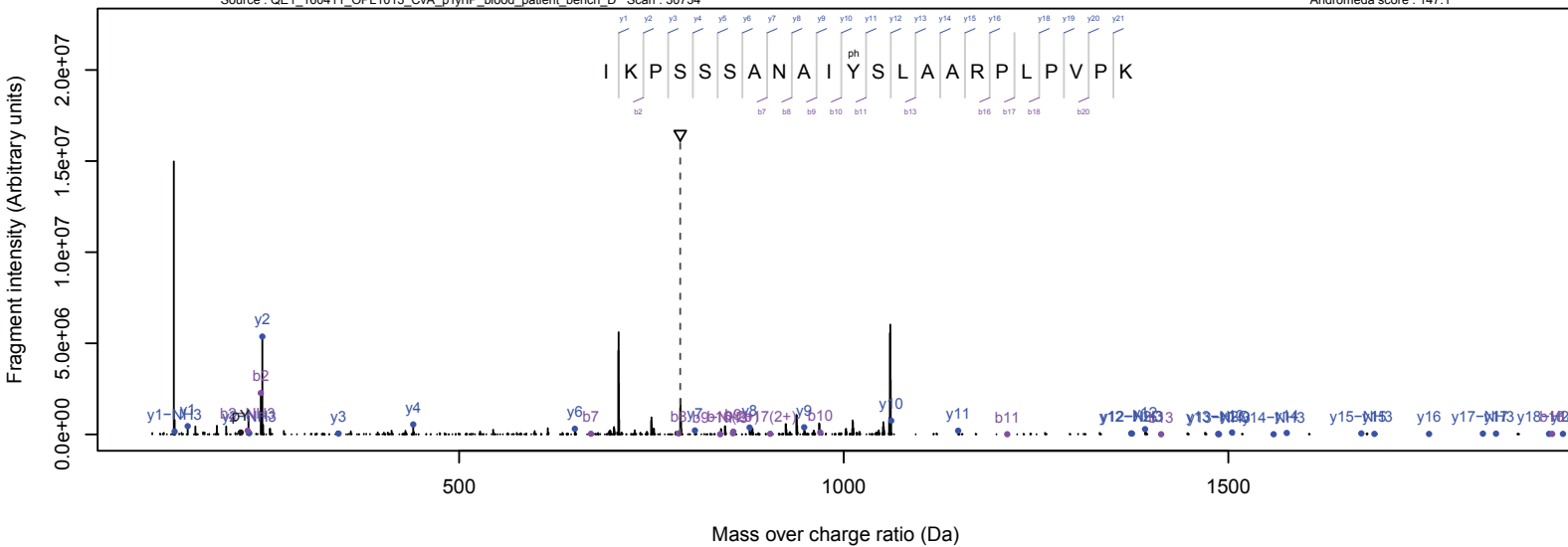
Andromeda score: 147.62



IKPSSANAIY(ph)SLAARPLVPK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 30734

Andromeda score: 147.1

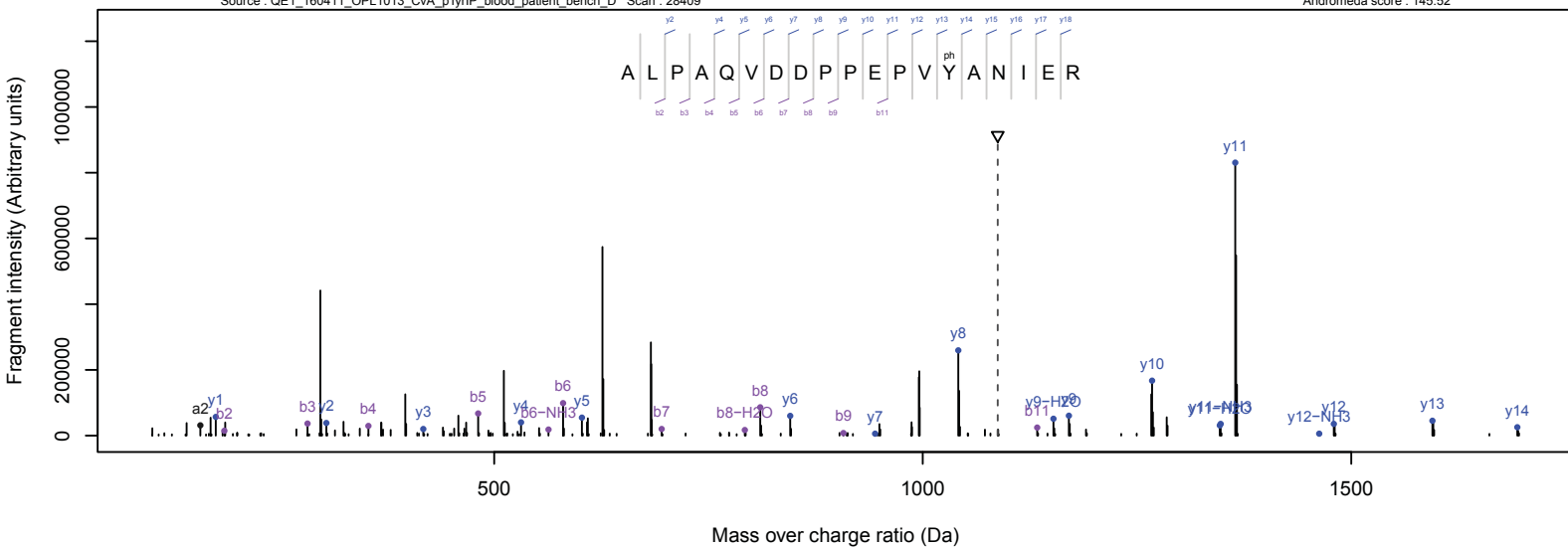




## ALPAQVDDPPEPVY(ph)ANIER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28409

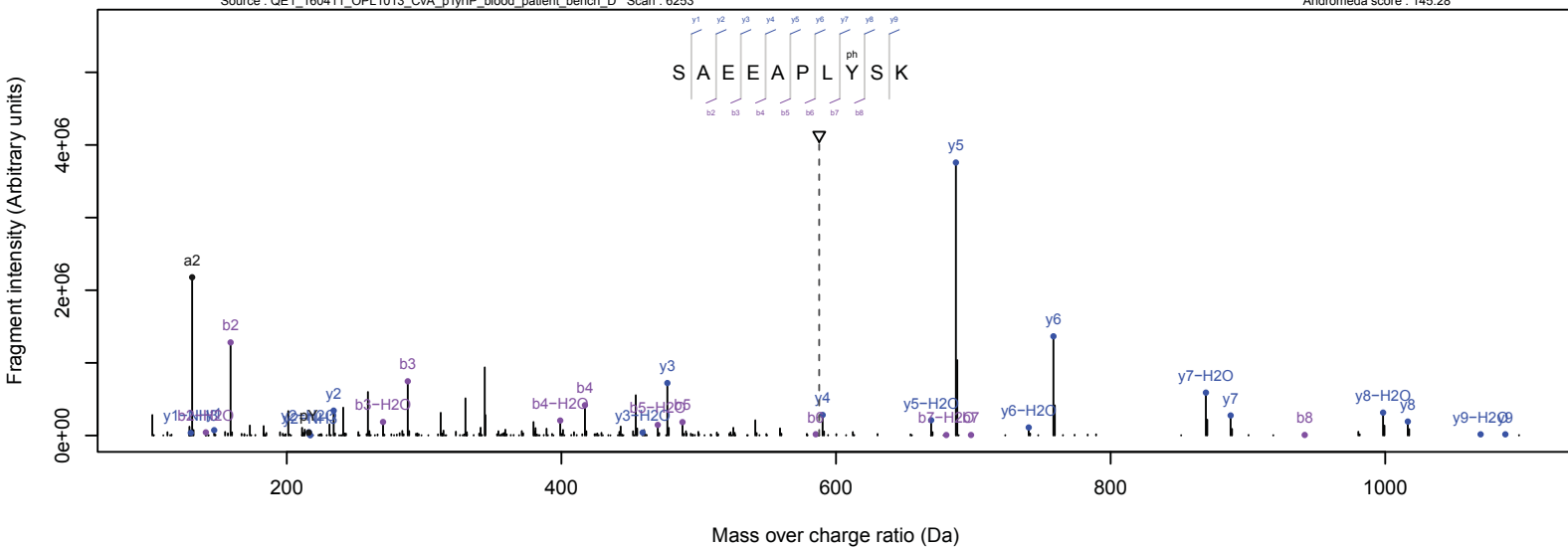
Andromeda score : 145.52



## SAEEAPLYSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 6253

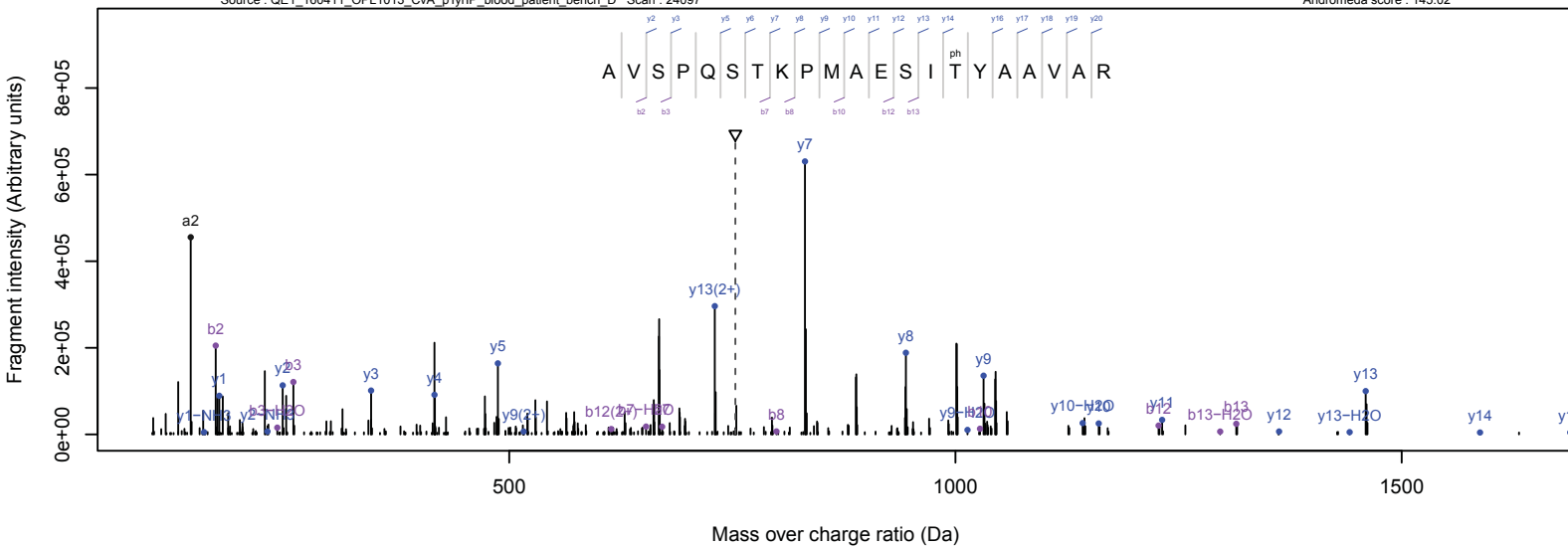
Andromeda score : 145.28

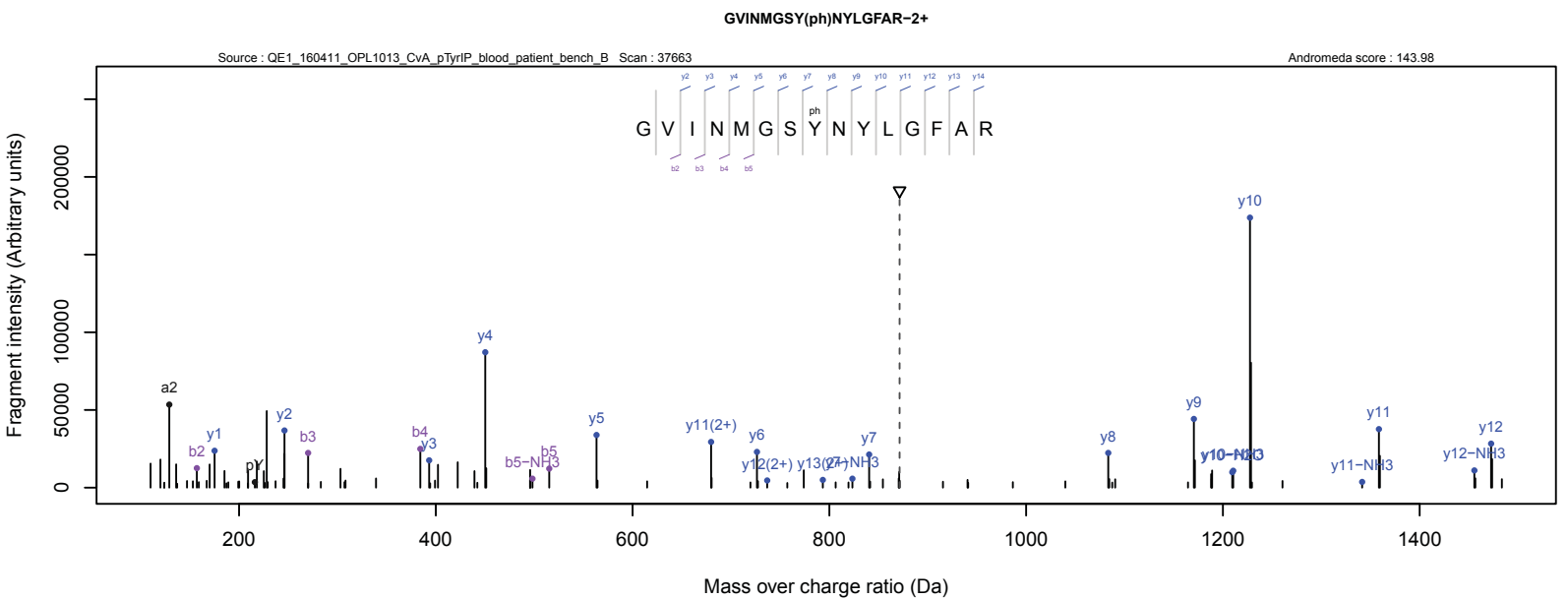
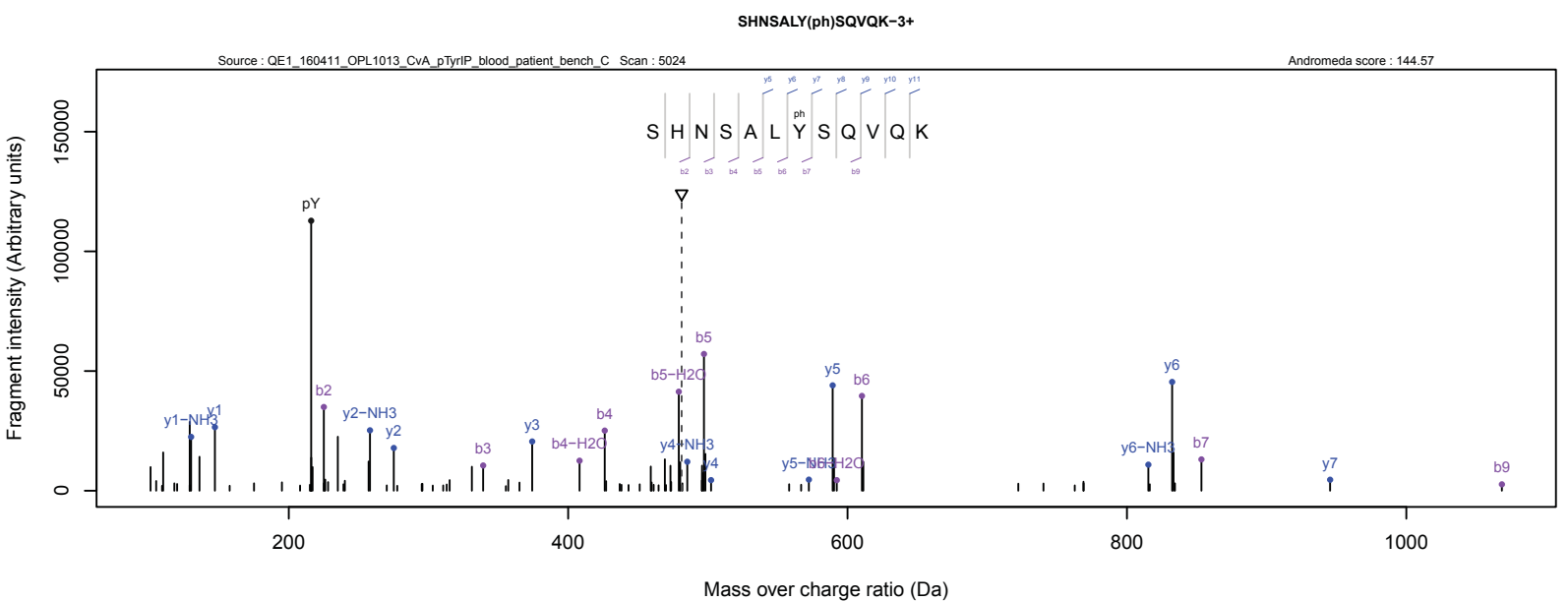
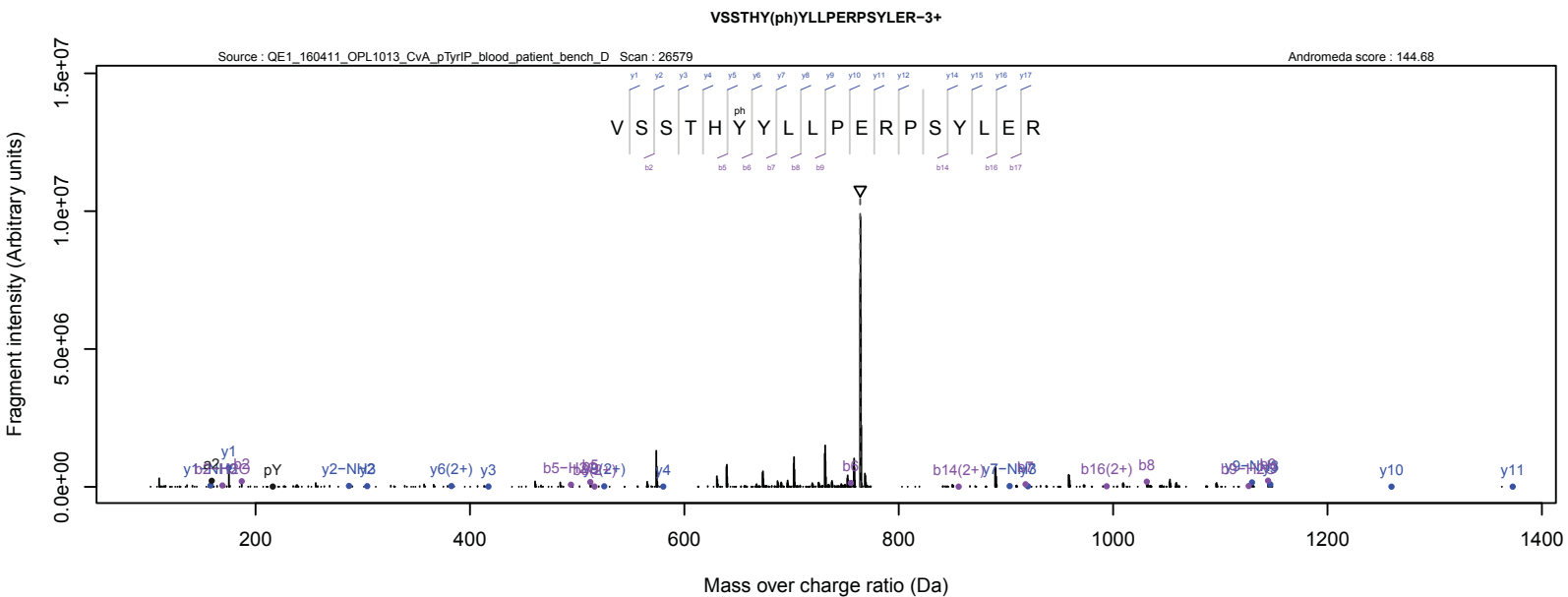


## AVSPQSTKPMAESIT(ph)YAAVAR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24097

Andromeda score : 145.02

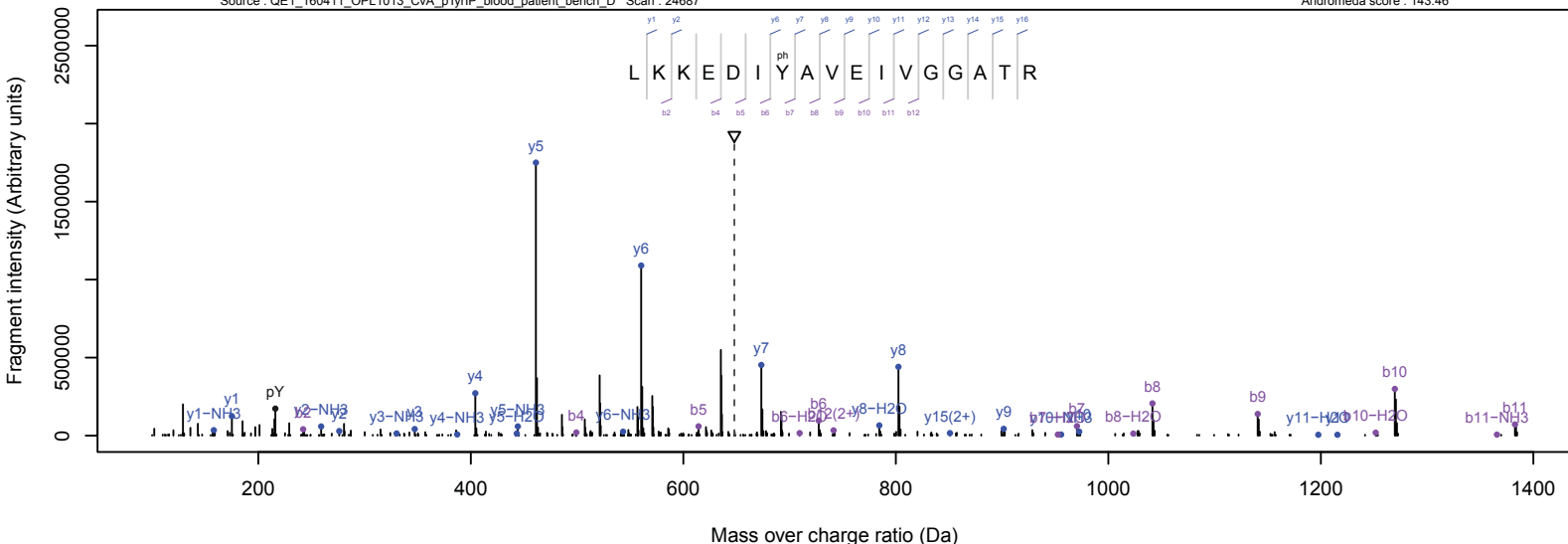




LKKEDIY(ph)AVEIVGGATR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24687

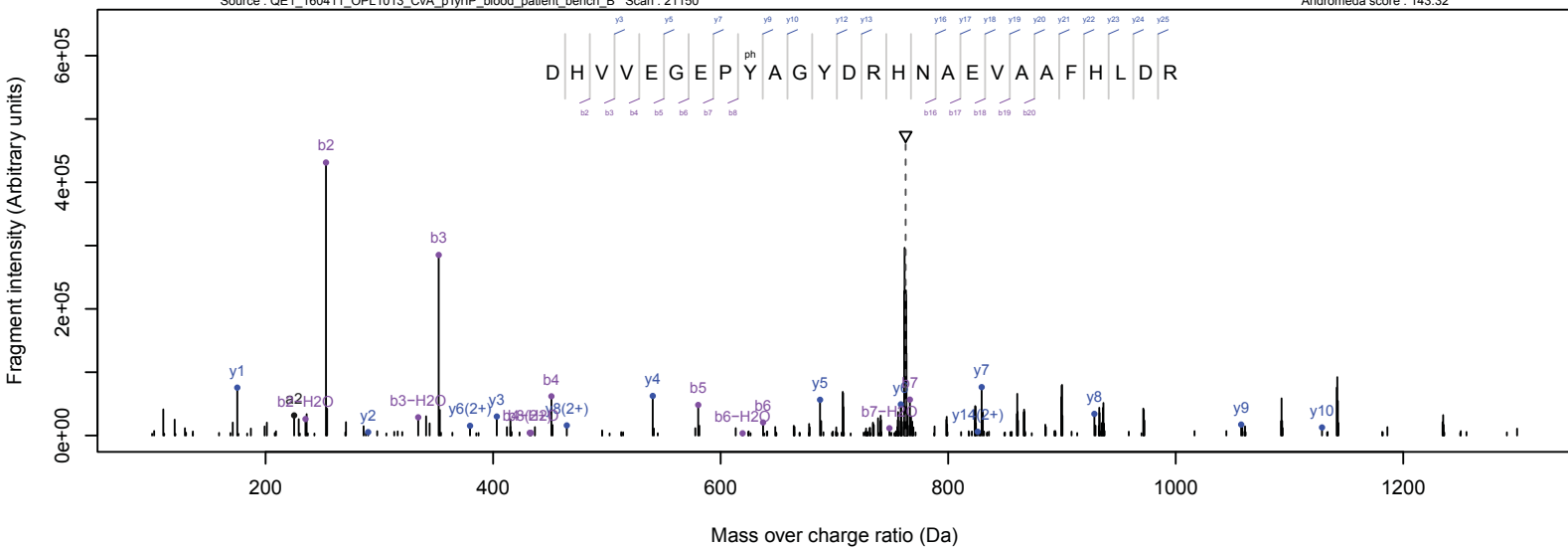
Andromeda score : 143.46



DHVVEGEPY(ph)AGYDRHNAEVAAFHLDR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 21150

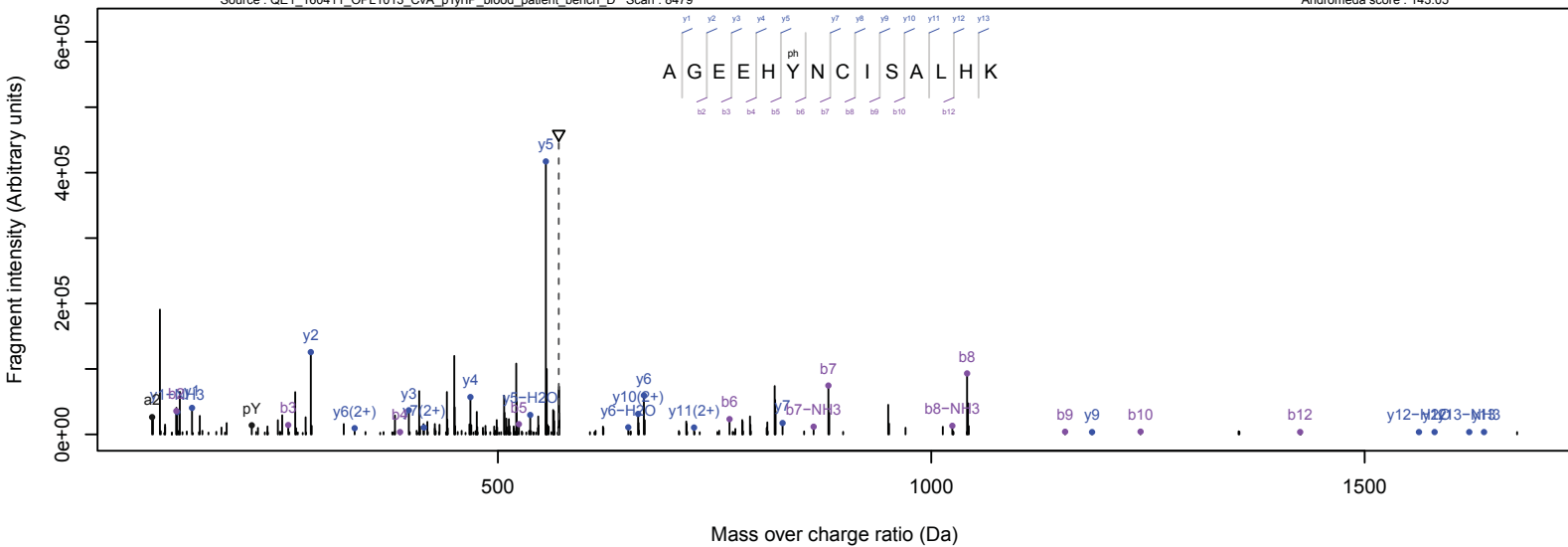
Andromeda score : 143.32



AGEEHY(ph)NCISALHK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 8479

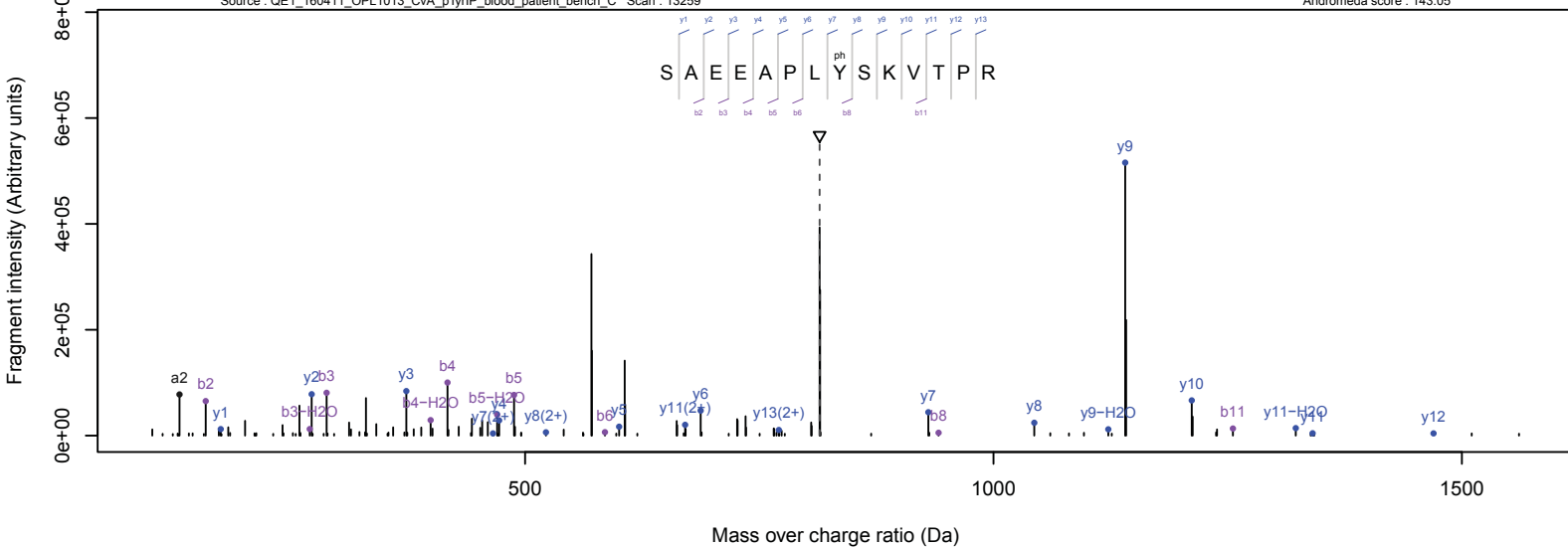
Andromeda score : 143.05



SAEEAPLY(ph)SKVTPR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 13259

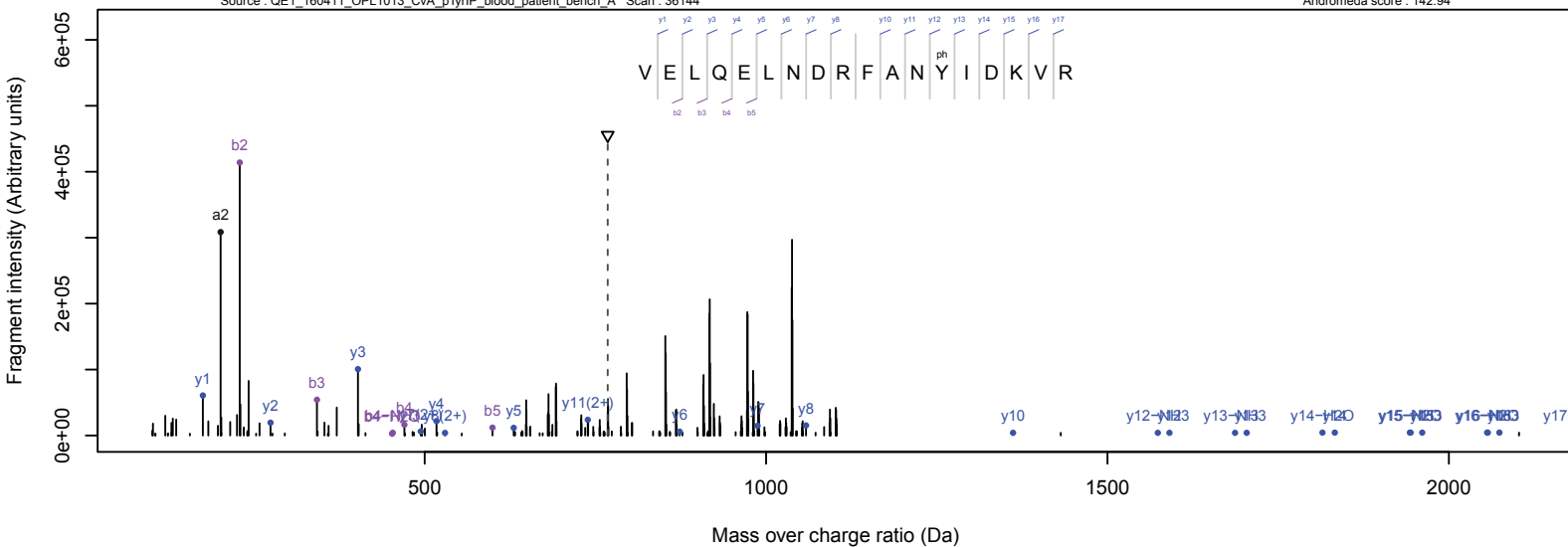
Andromeda score : 143.05



VELQELNDRFANY(ph)IDKVR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 36144

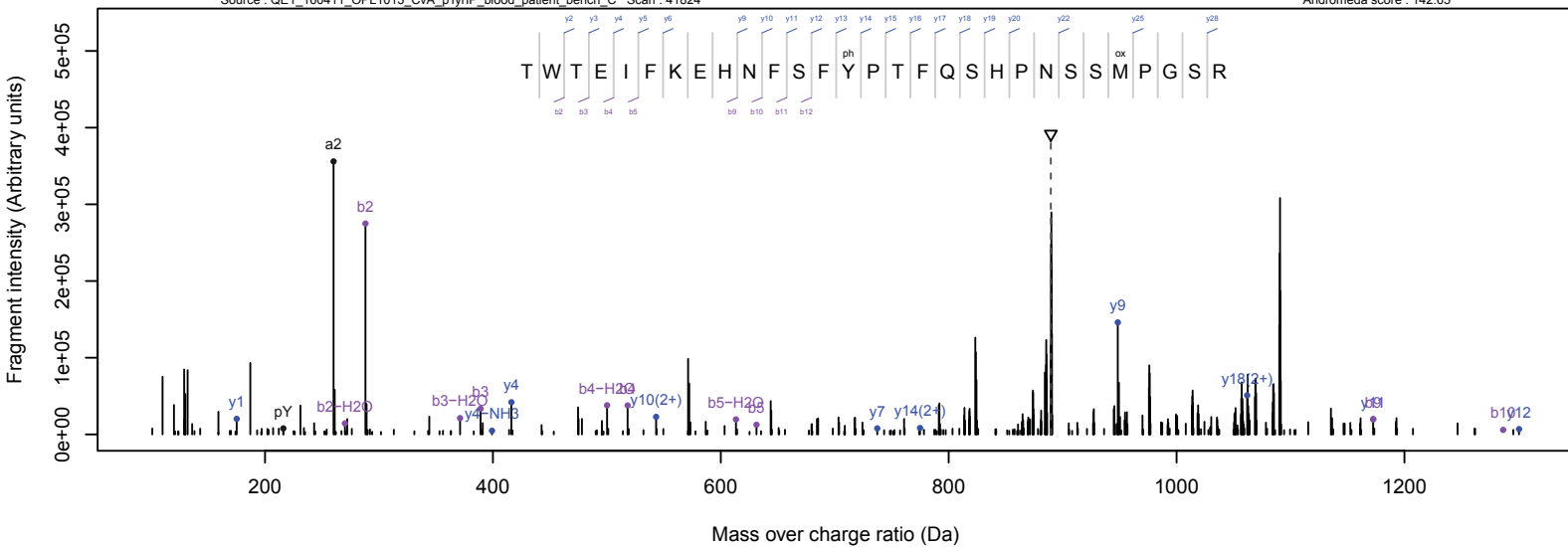
Andromeda score : 142.94



TWTEIFKEHNFSFY(ph)PTFQSHPNSSM(ox)PGSR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 41824

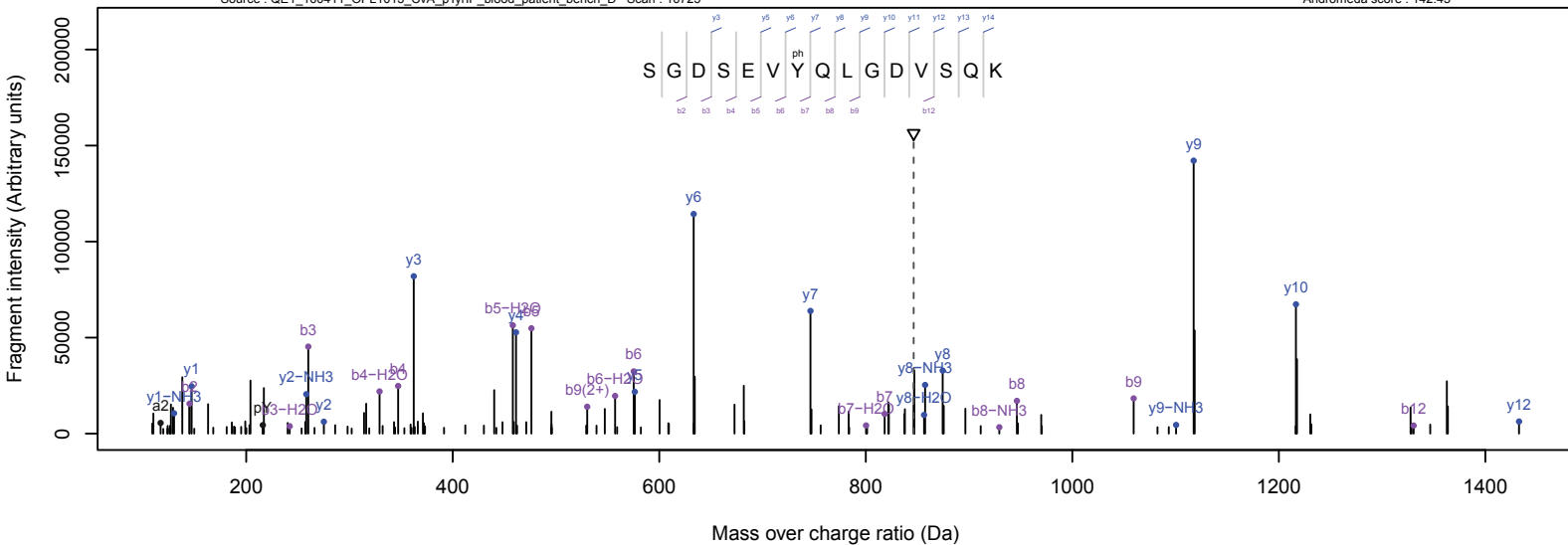
Andromeda score : 142.65



## SGDSEVY(ph)QLGDVSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 16725

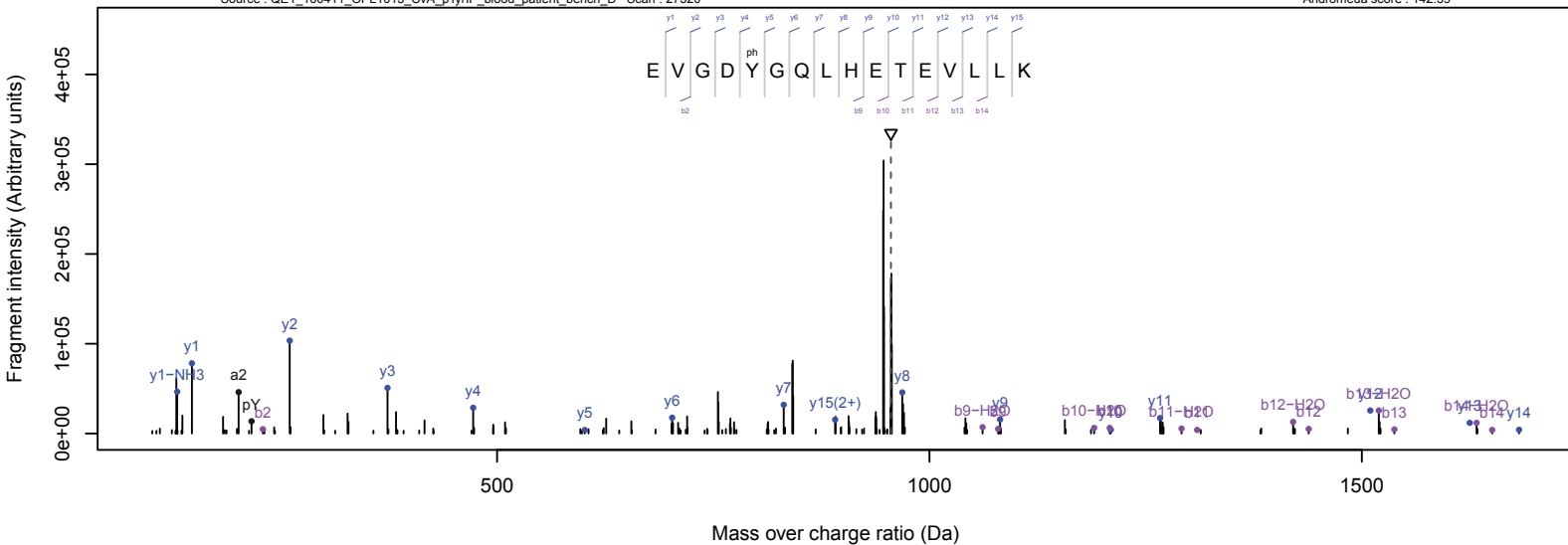
Andromeda score : 142.45



## EVGDY(ph)QLHETEVLK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 27320

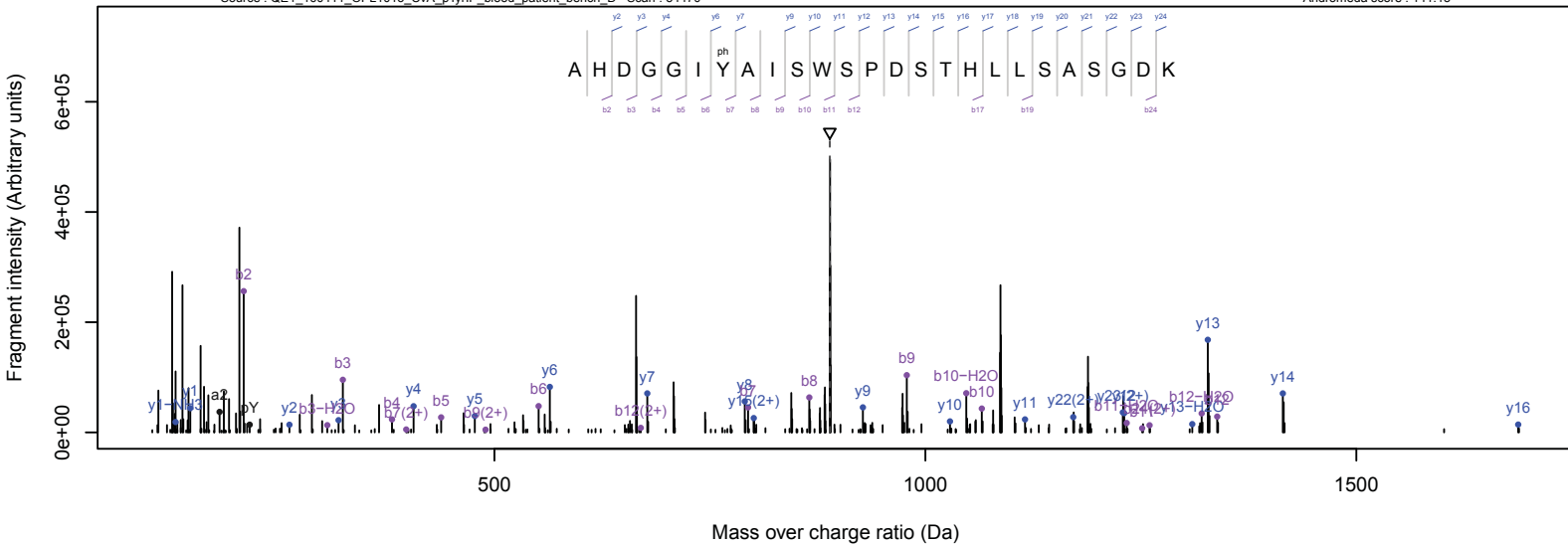
Andromeda score : 142.39



## AHDGGIY(ph)AISWSPDSTHLLSASGDK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 34470

Andromeda score : 141.15

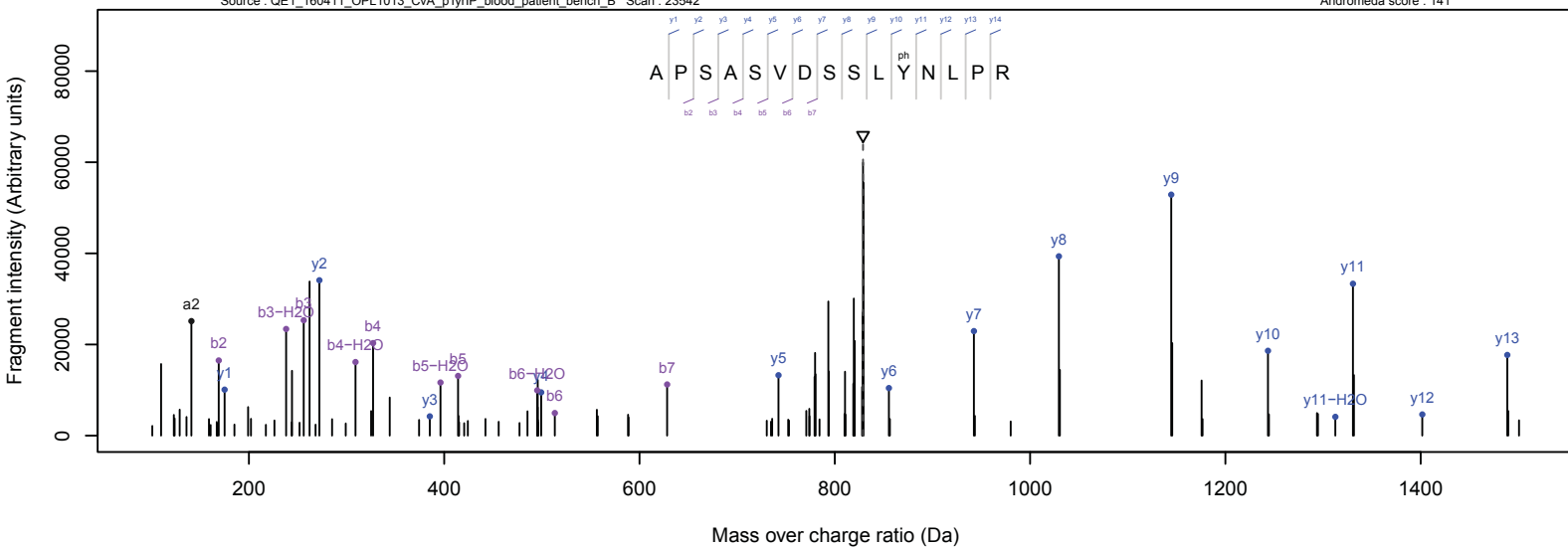




APSASVDSL(ph)NLPR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 23542

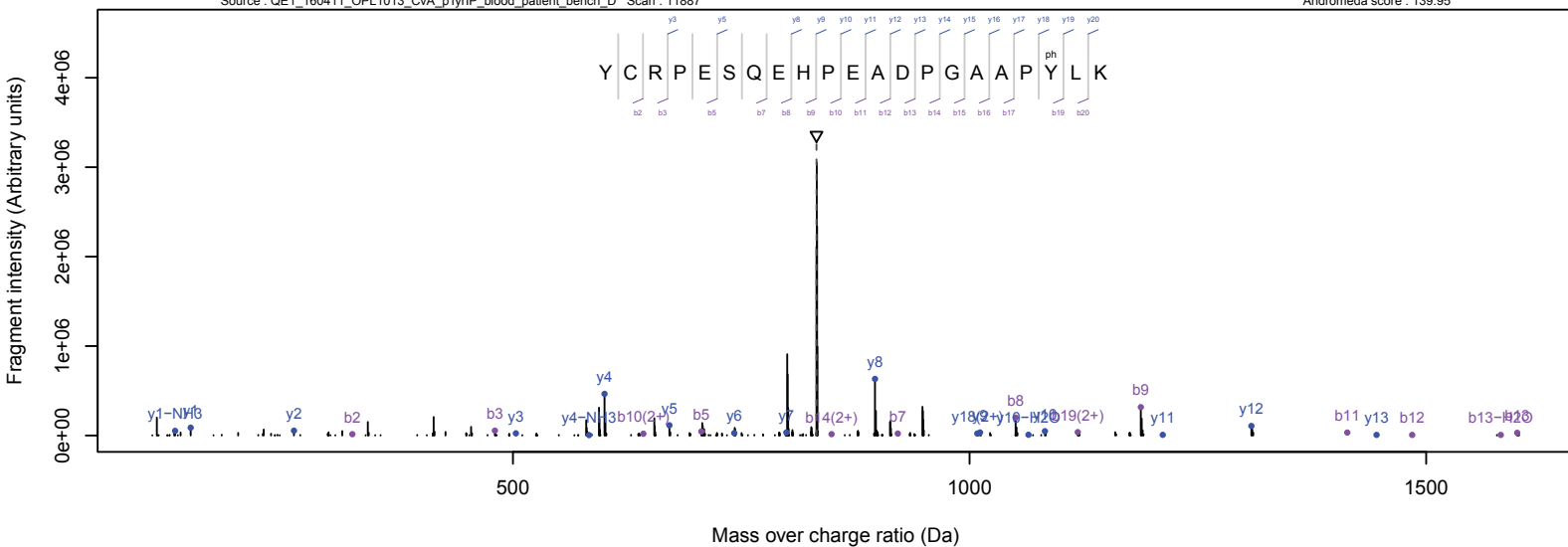
Andromeda score : 141



YCRPESQEHPEADPGAAPY(ph)LK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11887

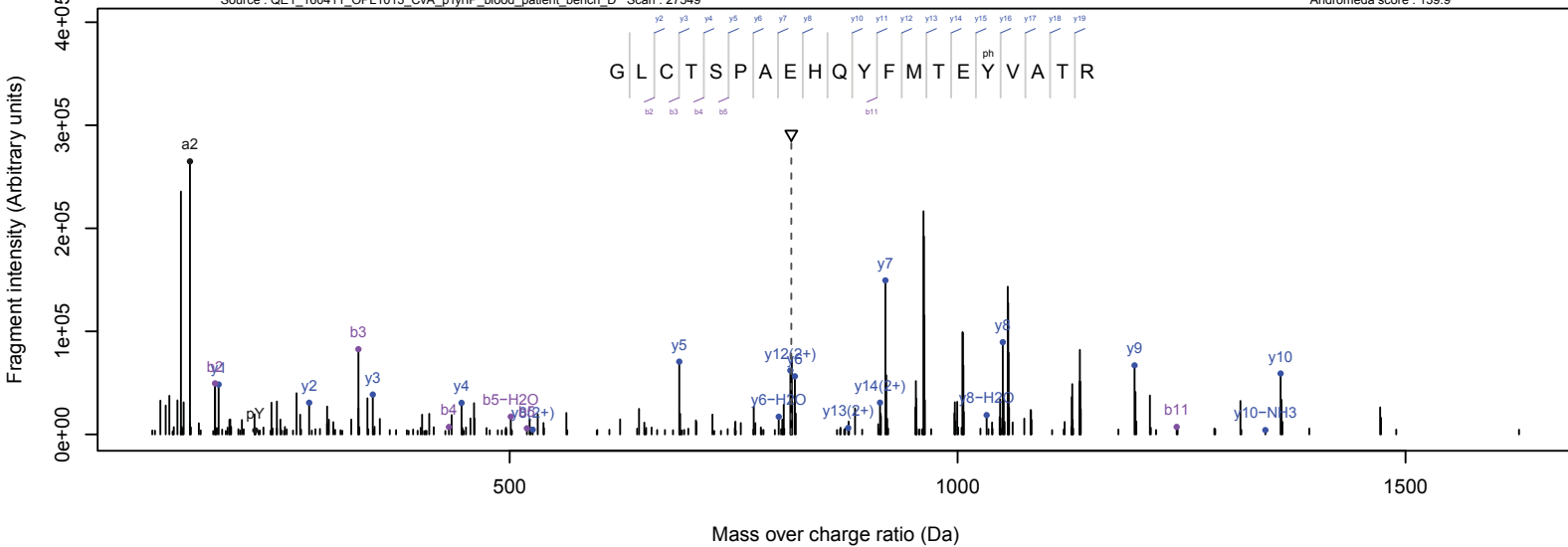
Andromeda score : 139.95



GLCTSPAETHQYFMTEY(ph)VATR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 27549

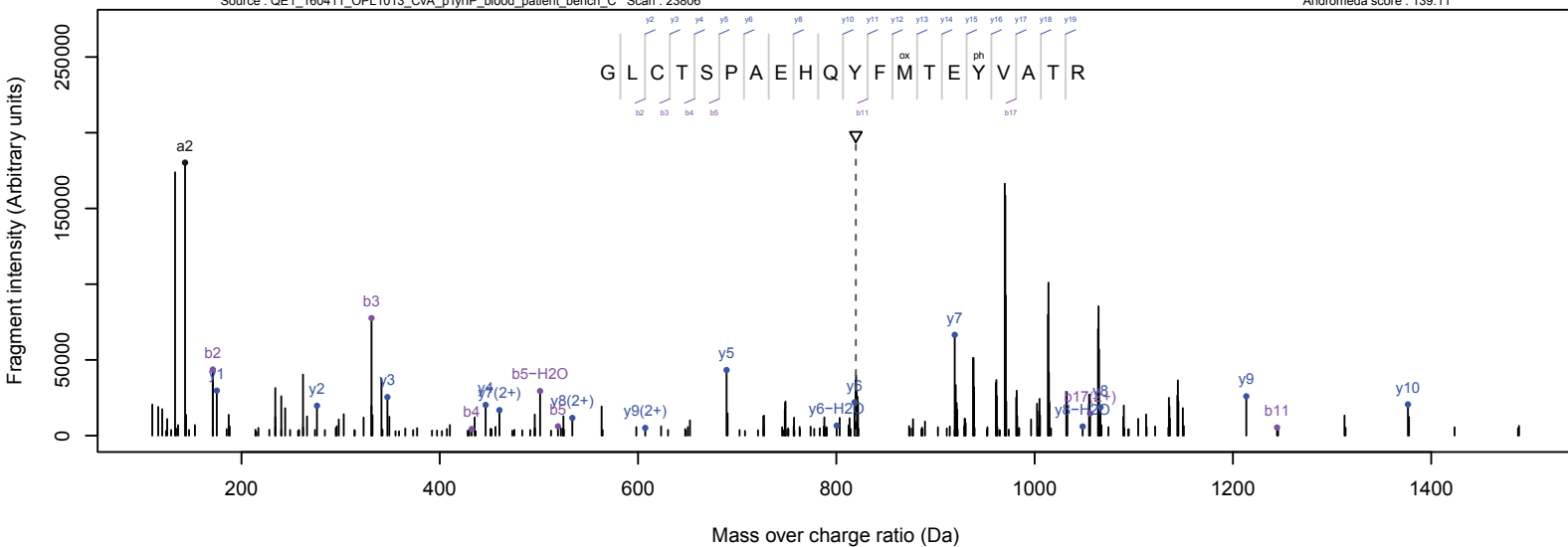
Andromeda score : 139.9



GLCTSPA<sup>ox</sup>EHQYFMTEYVATR-3+

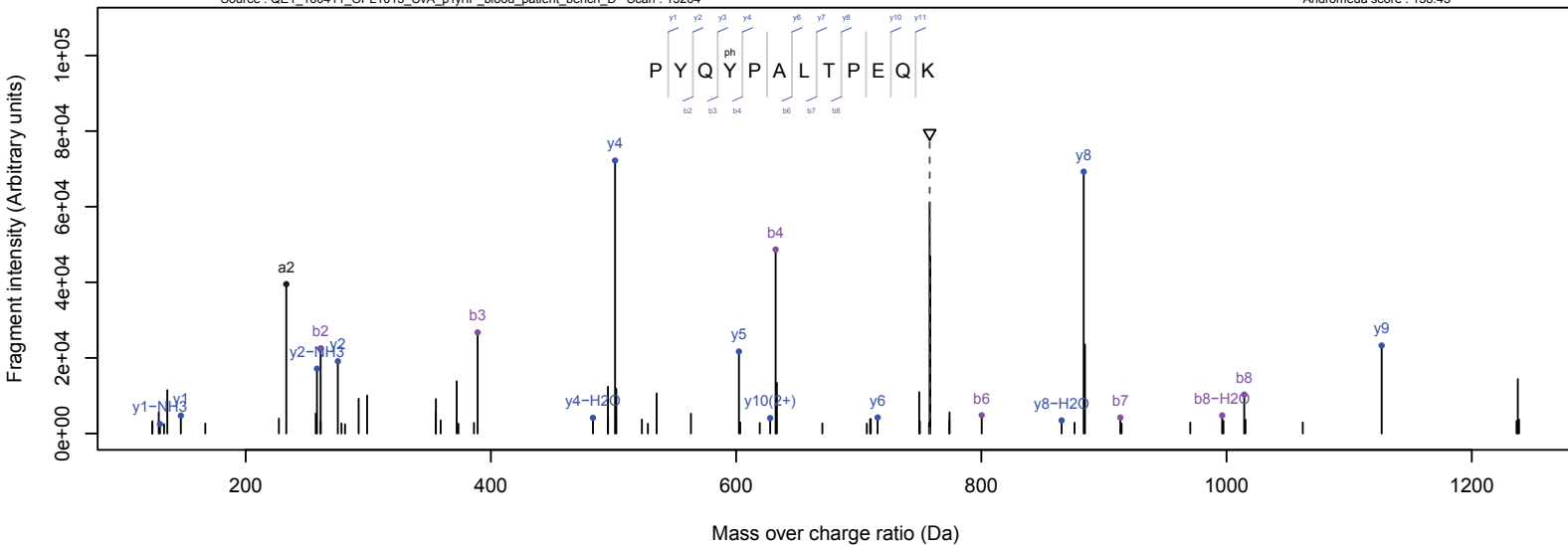
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 23806

Andromeda score: 139.11

PYQY<sup>ph</sup>PALTP<sup>ox</sup>EQK-2+

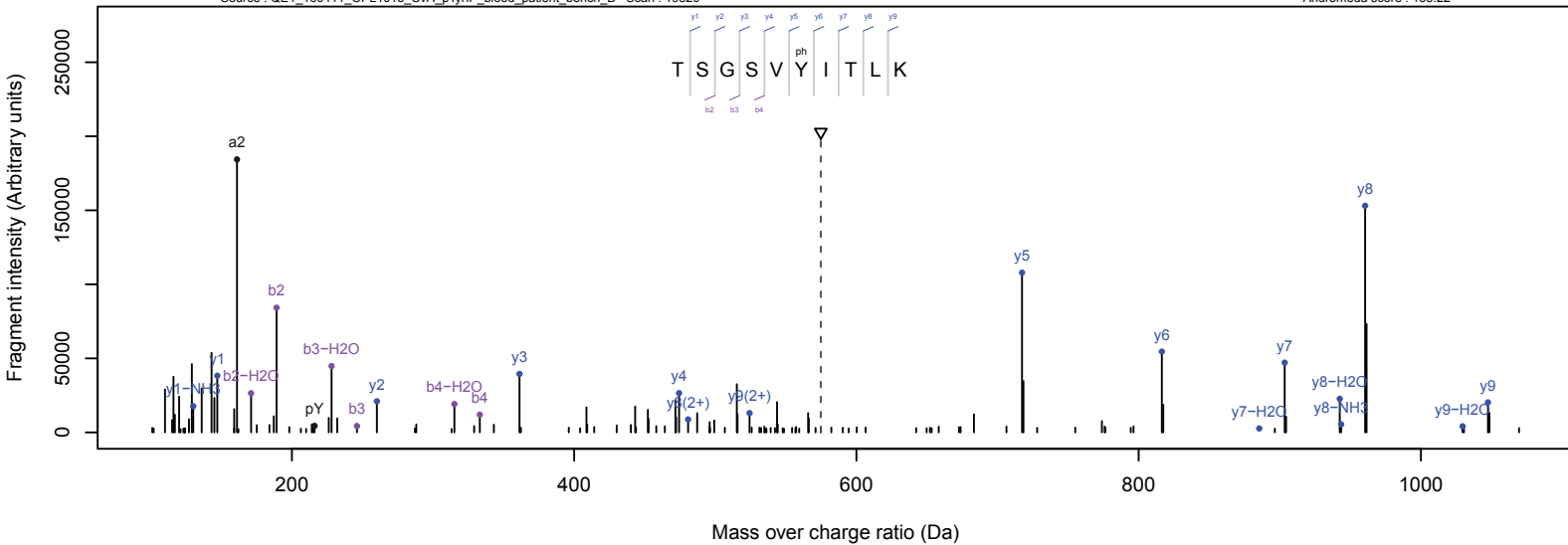
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 15264

Andromeda score: 138.45

TSGSVY<sup>ph</sup>ITLK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 19820

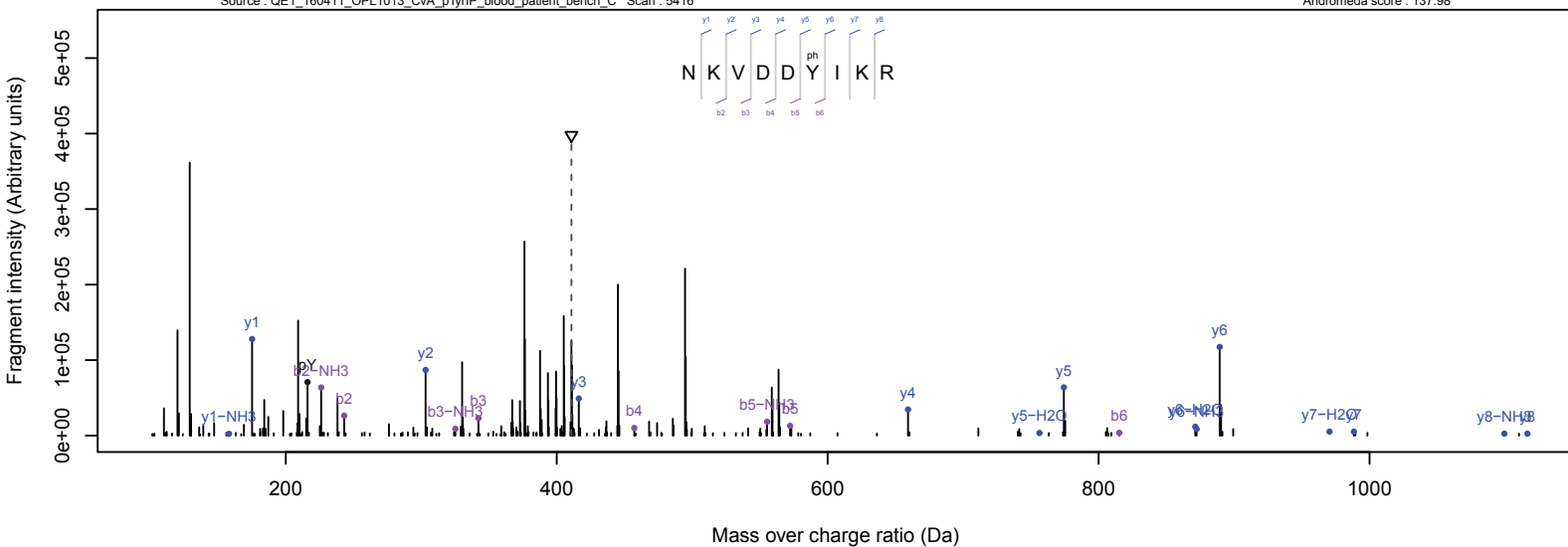
Andromeda score: 138.22



NKVVDDY(ph)IKR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 5416

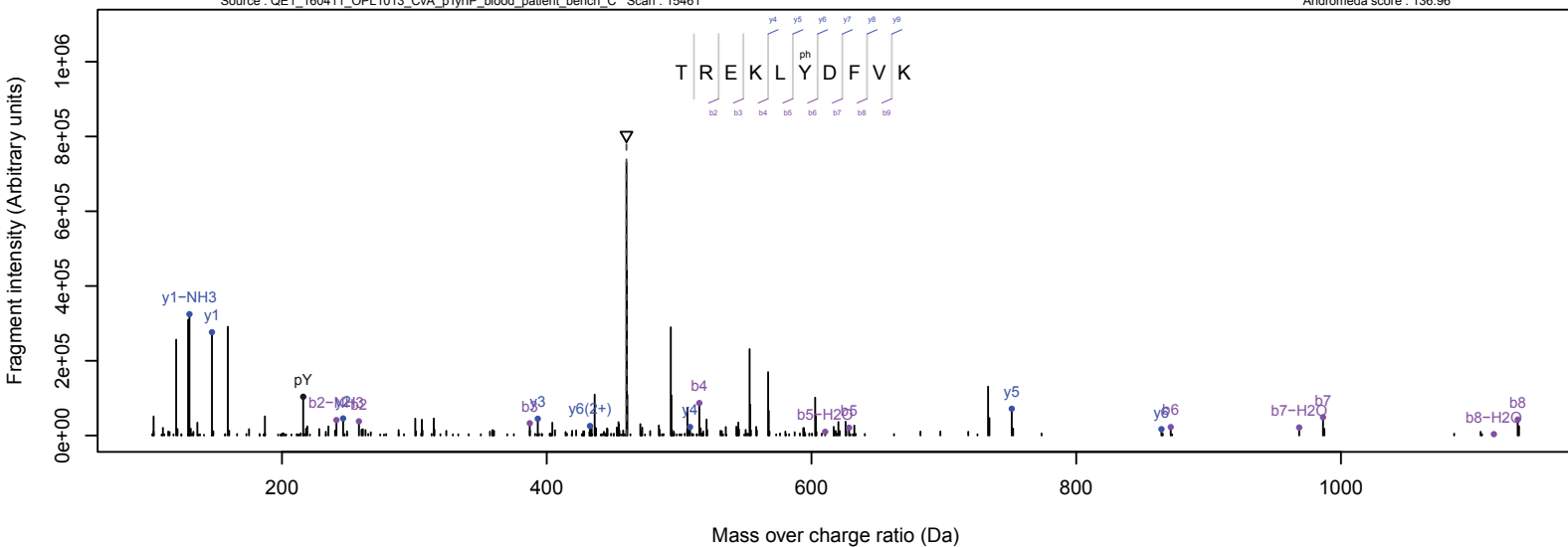
Andromeda score : 137.98



TREKLY(ph)DFVK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 15461

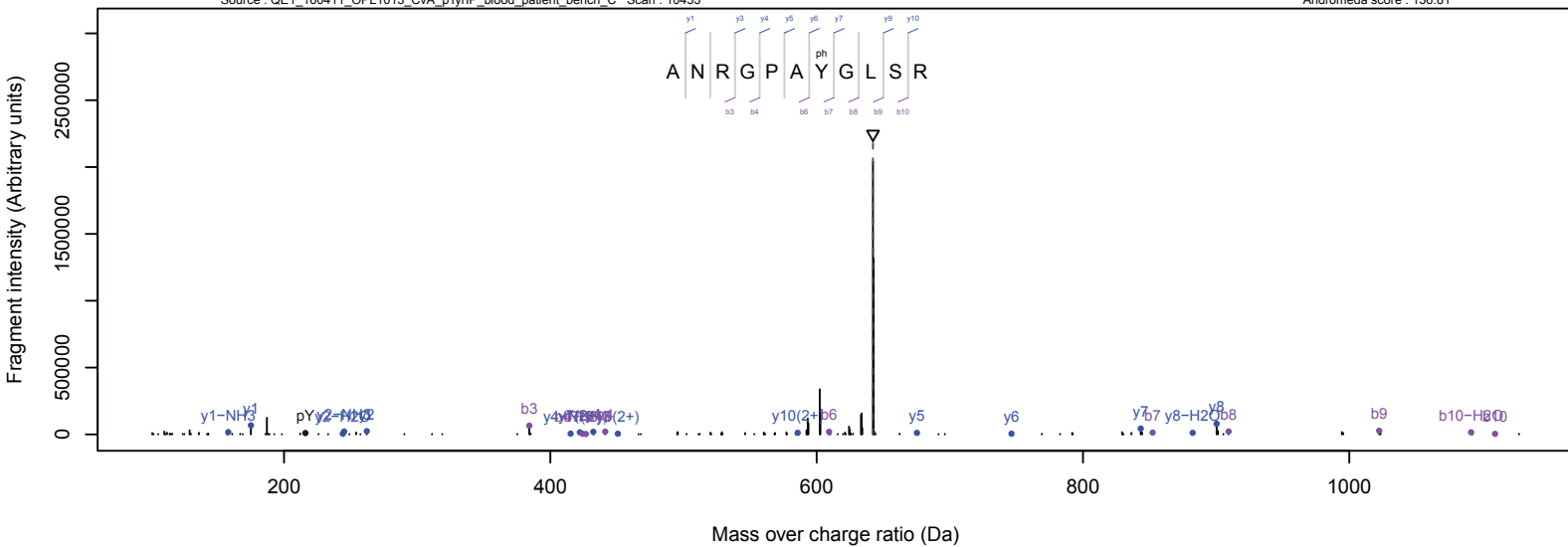
Andromeda score : 136.96



(ac)ANRGPAY(ph)GLSR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 10453

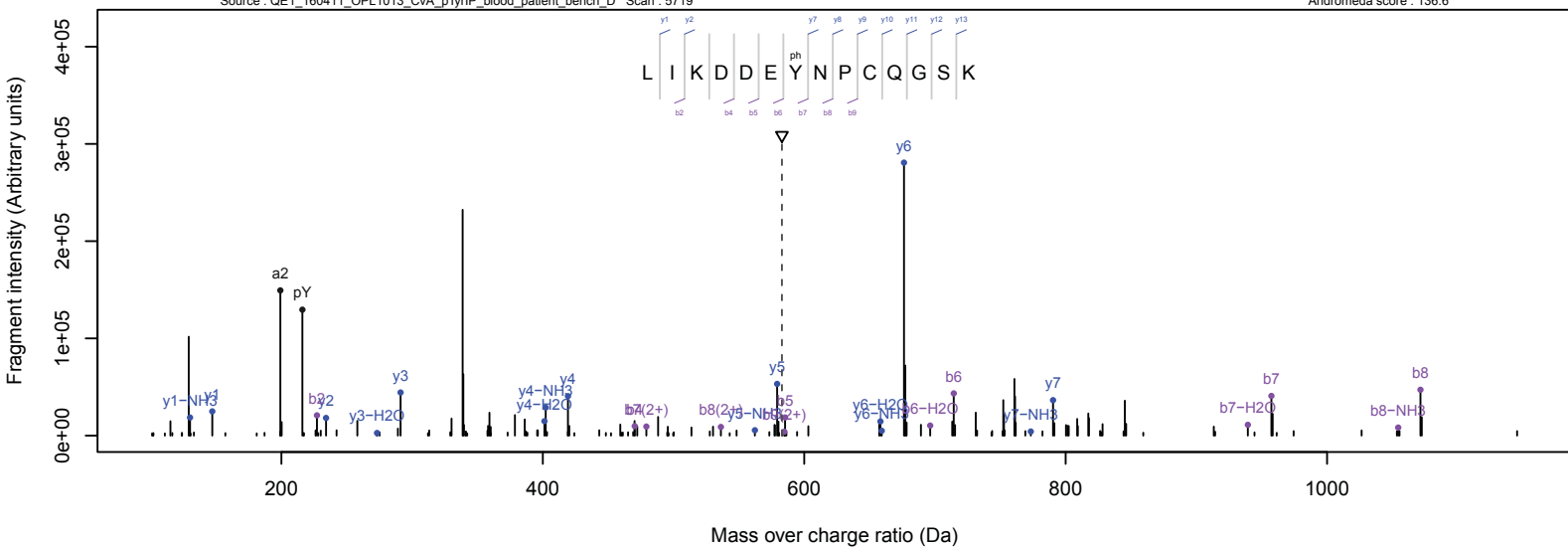
Andromeda score : 136.81



## LIKDDEY(ph)NPCQGSK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5719

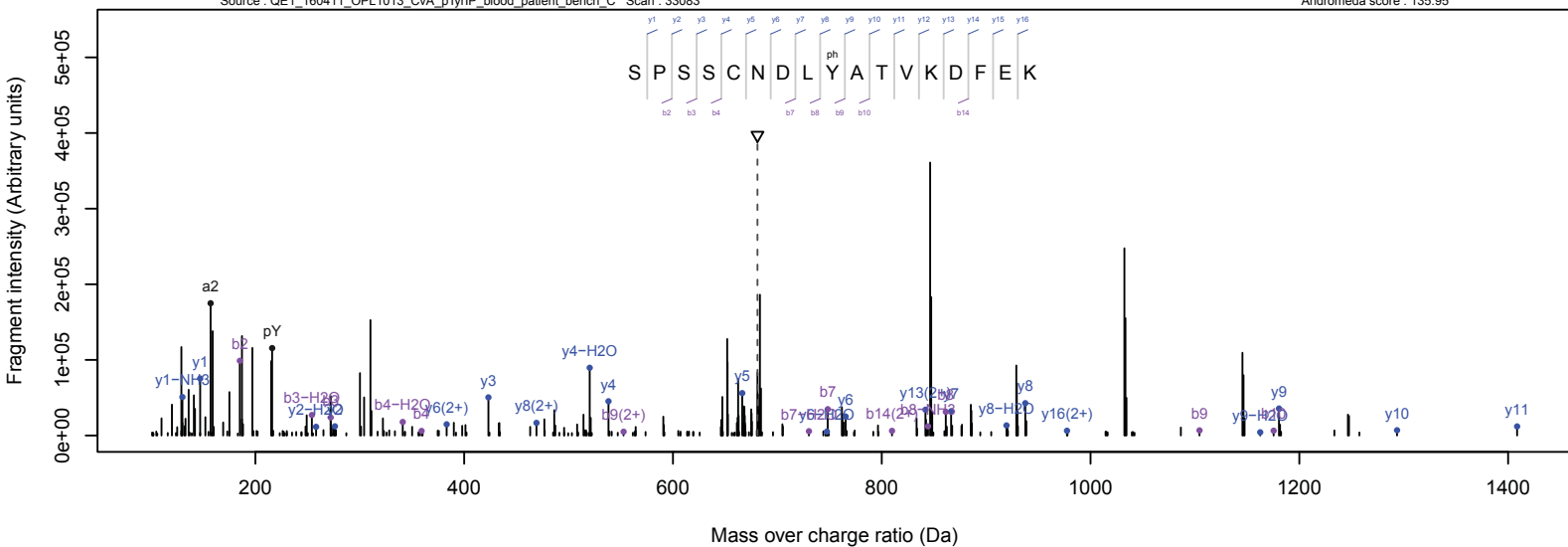
Andromeda score : 136.6



## SPSSCNLDY(ph)ATVKDFEK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 33083

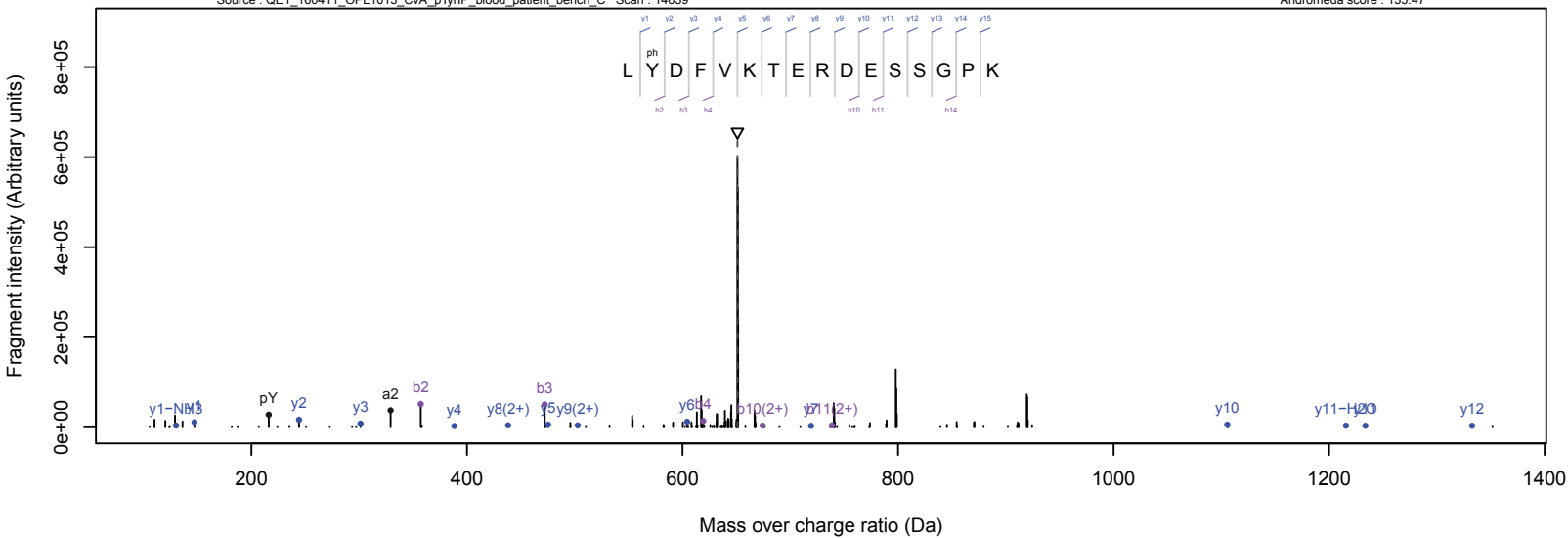
Andromeda score : 135.95



## LY(ph)DFVKTERDESSGPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 14639

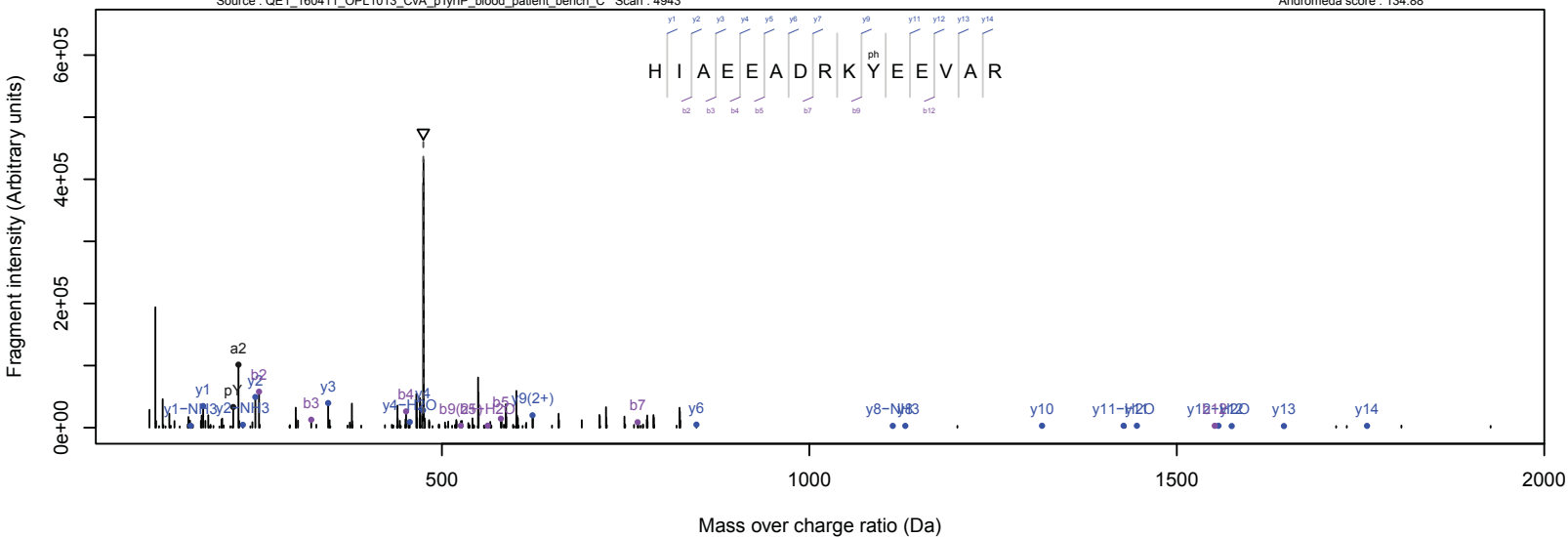
Andromeda score : 135.47



## HIAEEADRKY(ph)EEVAR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 4943

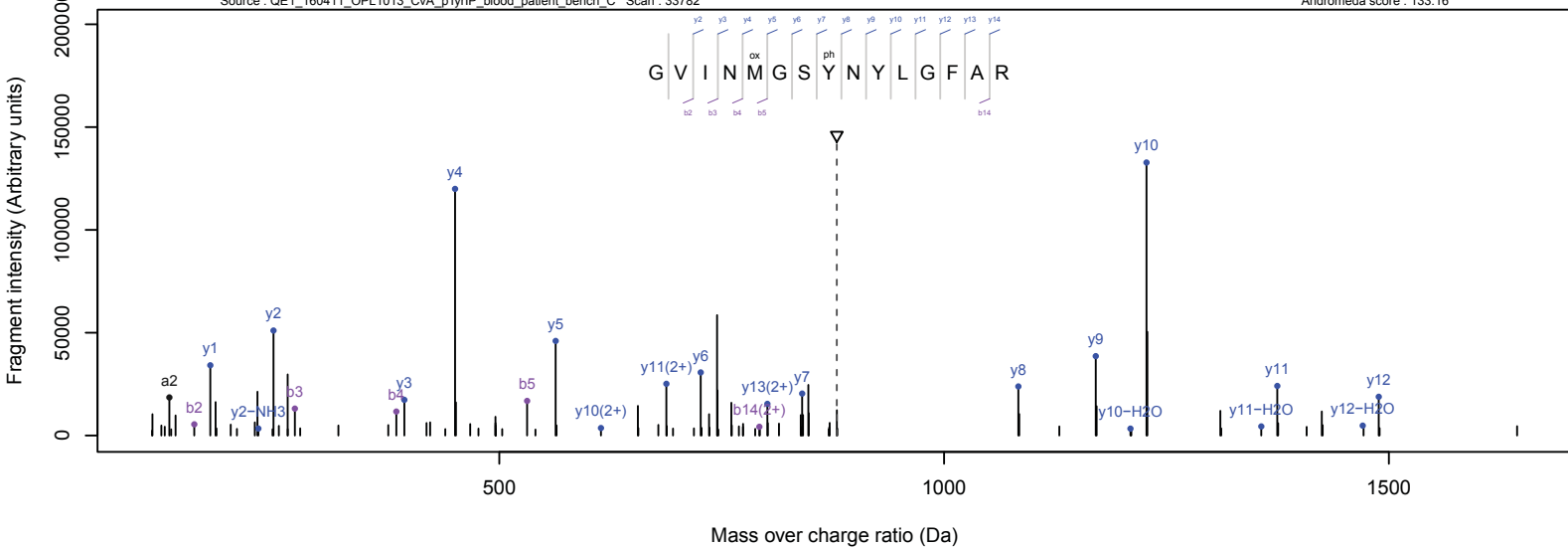
Andromeda score : 134.88



## GVINM(ox)GSY(ph)NYLGFAR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 33782

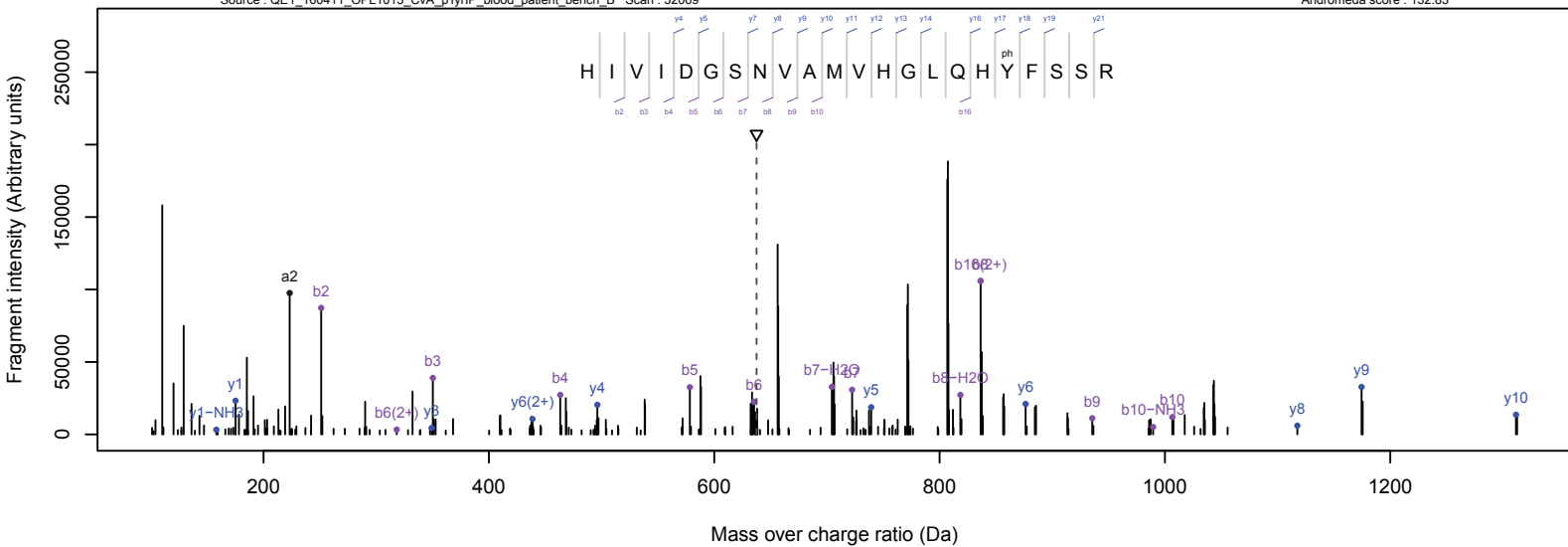
Andromeda score : 133.16



## HIVIDGSNVAMVHGLQHYP(ph)FSSR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 32069

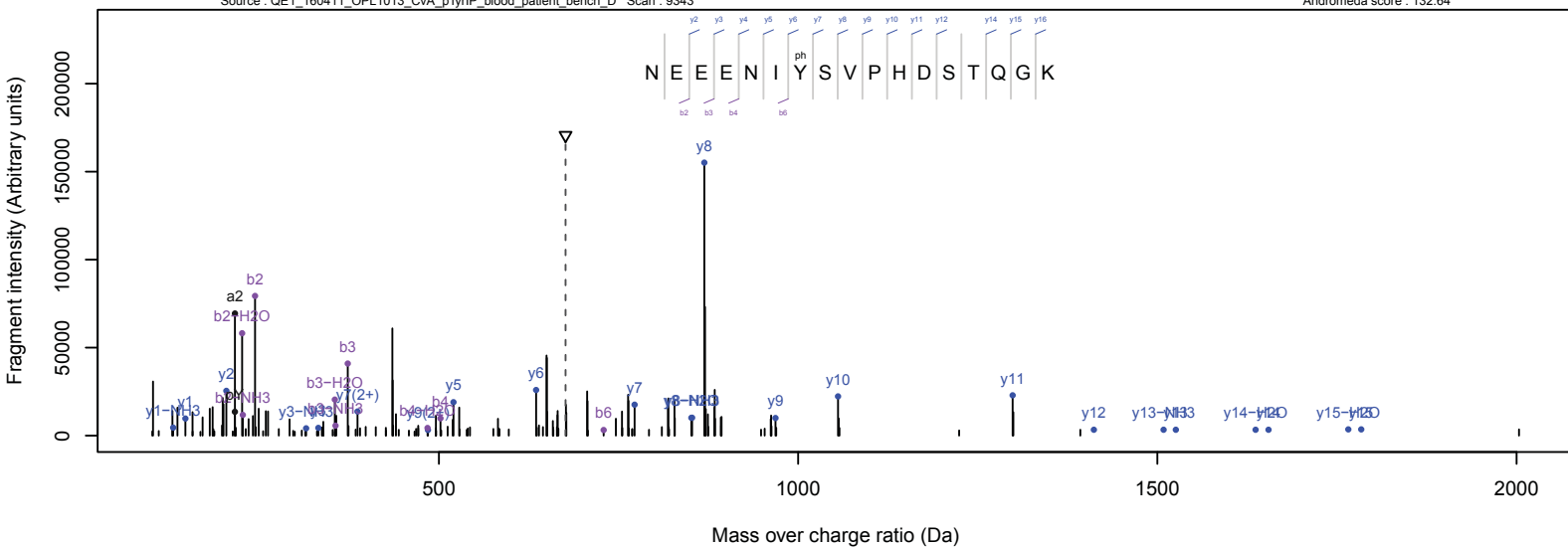
Andromeda score : 132.83



NEEENIY(ph)SVPHDSTQGK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 9343

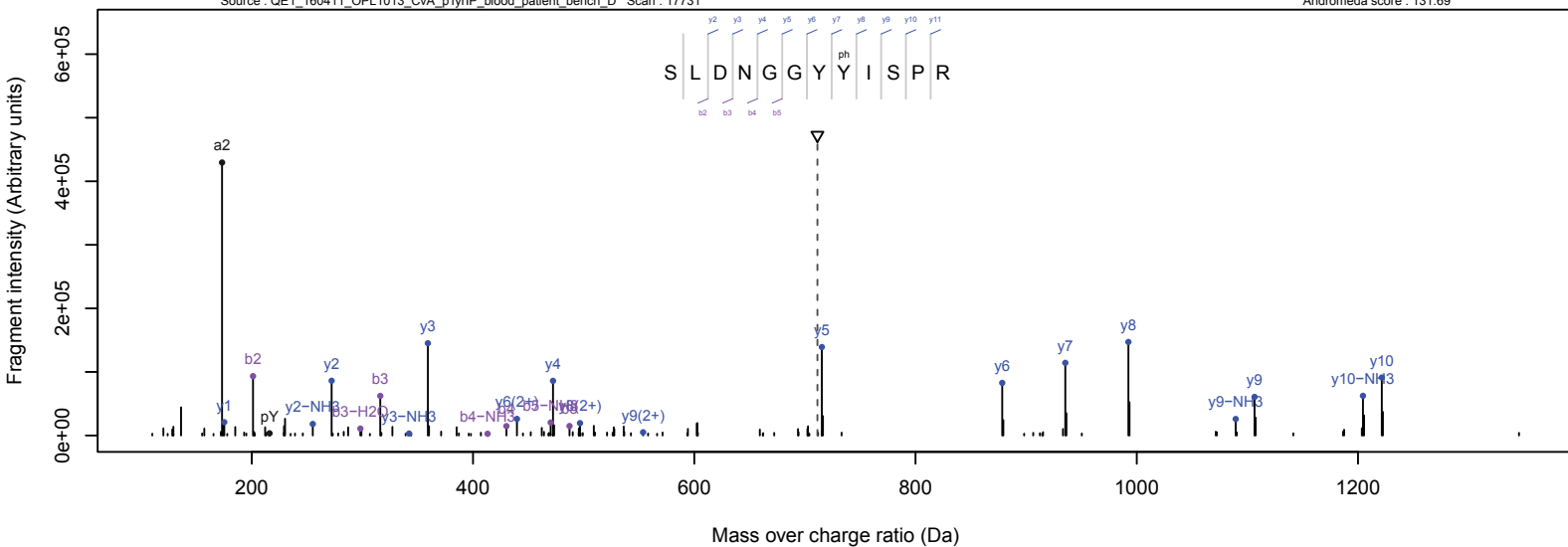
Andromeda score : 132.64



SLDNGGY(ph)ISPR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17731

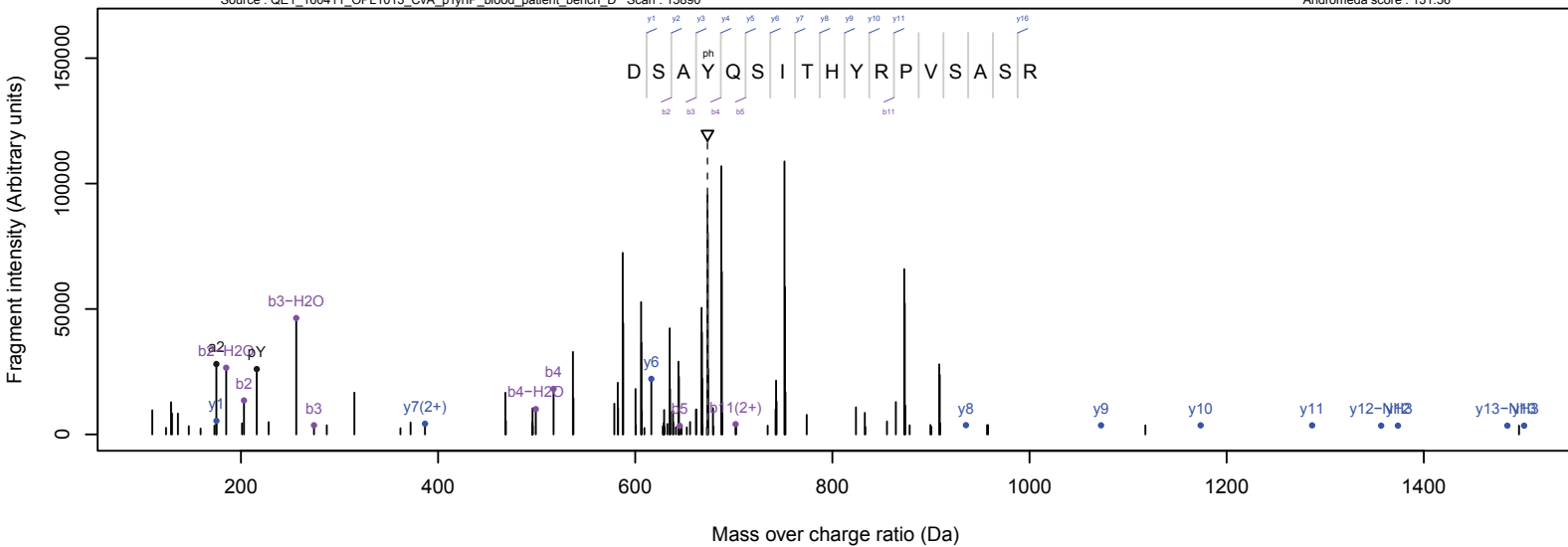
Andromeda score : 131.69



DSAY(ph)QSITHYRPVSA SR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 15890

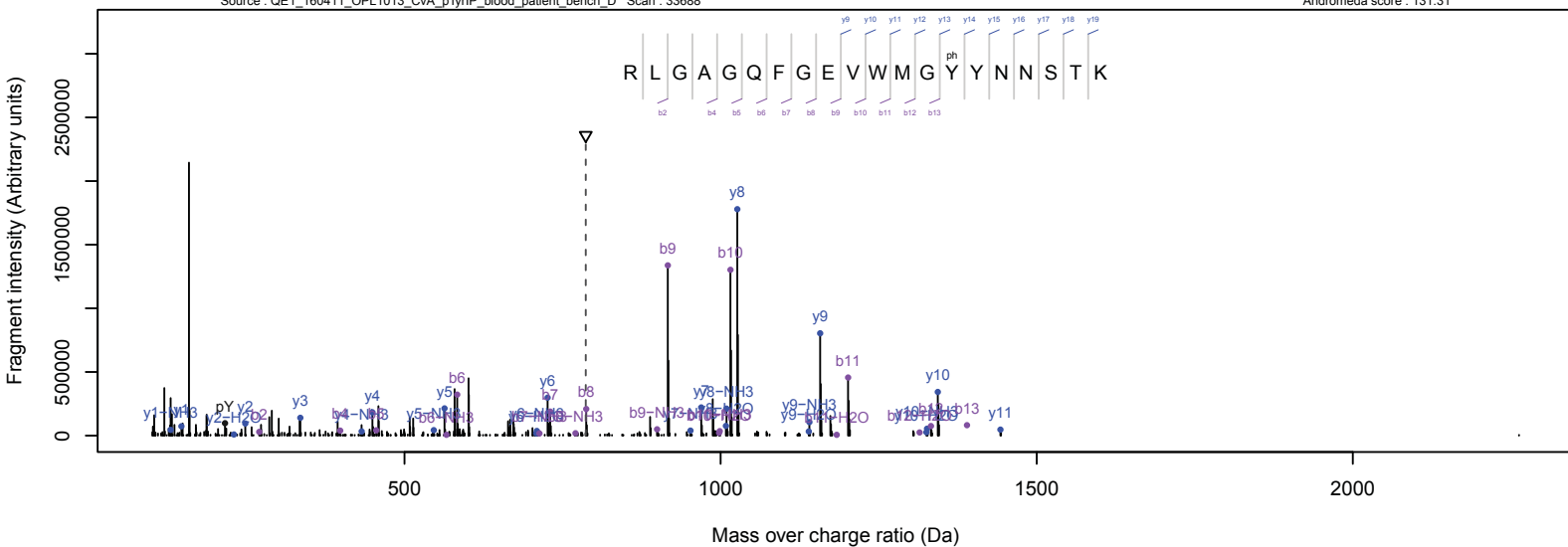
Andromeda score : 131.56



RLGAGQFGEVWVG<sup>ph</sup>YNNSTK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 33688

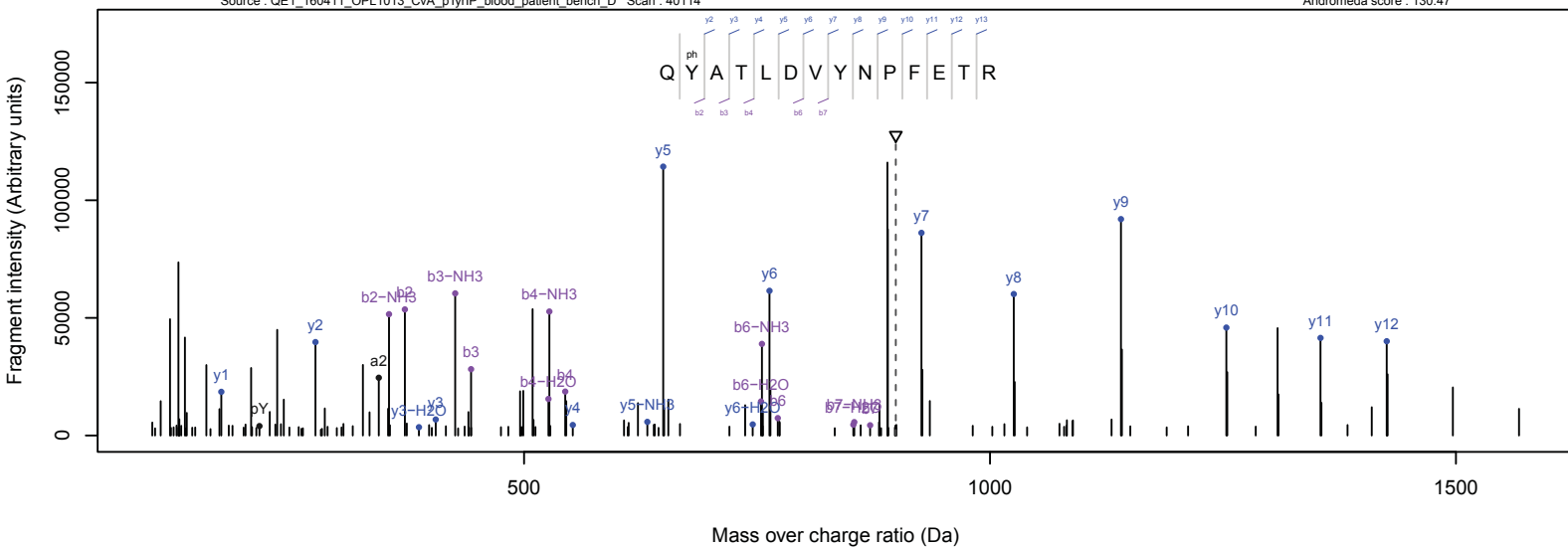
Andromeda score : 131.31



QY<sup>ph</sup>ATLDVYNPFETR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 40114

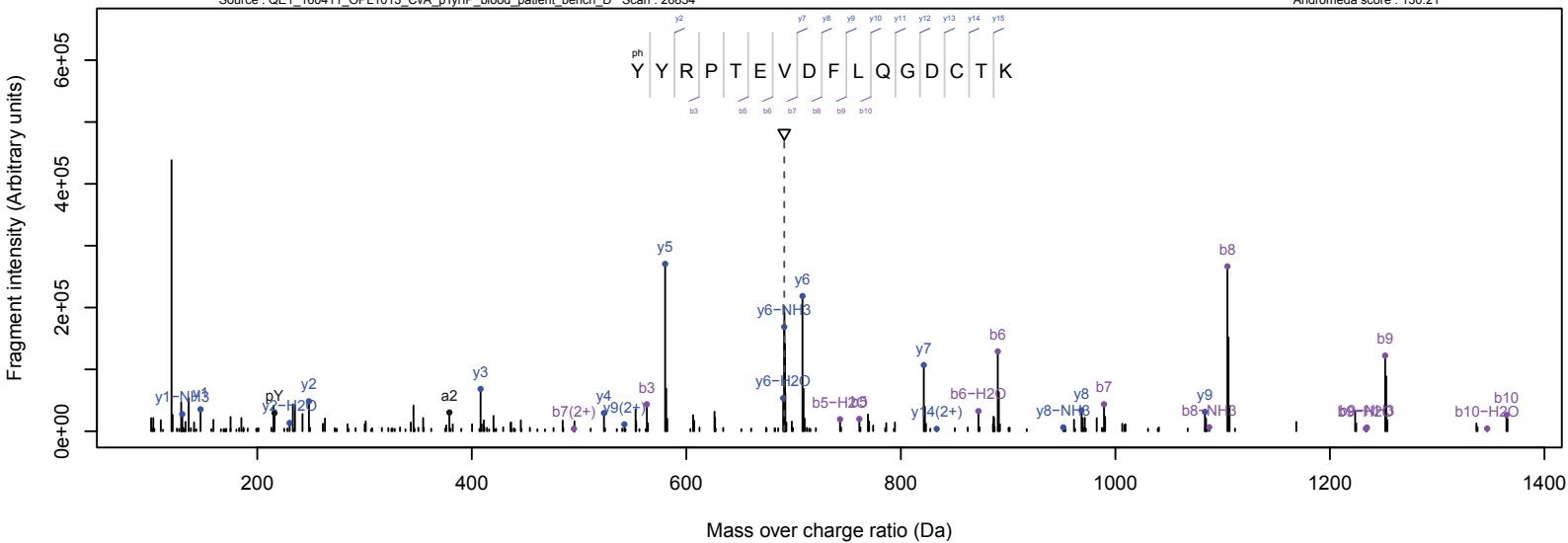
Andromeda score : 130.47



Y<sup>ph</sup>YRPTEVDFLQGDCTK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 26834

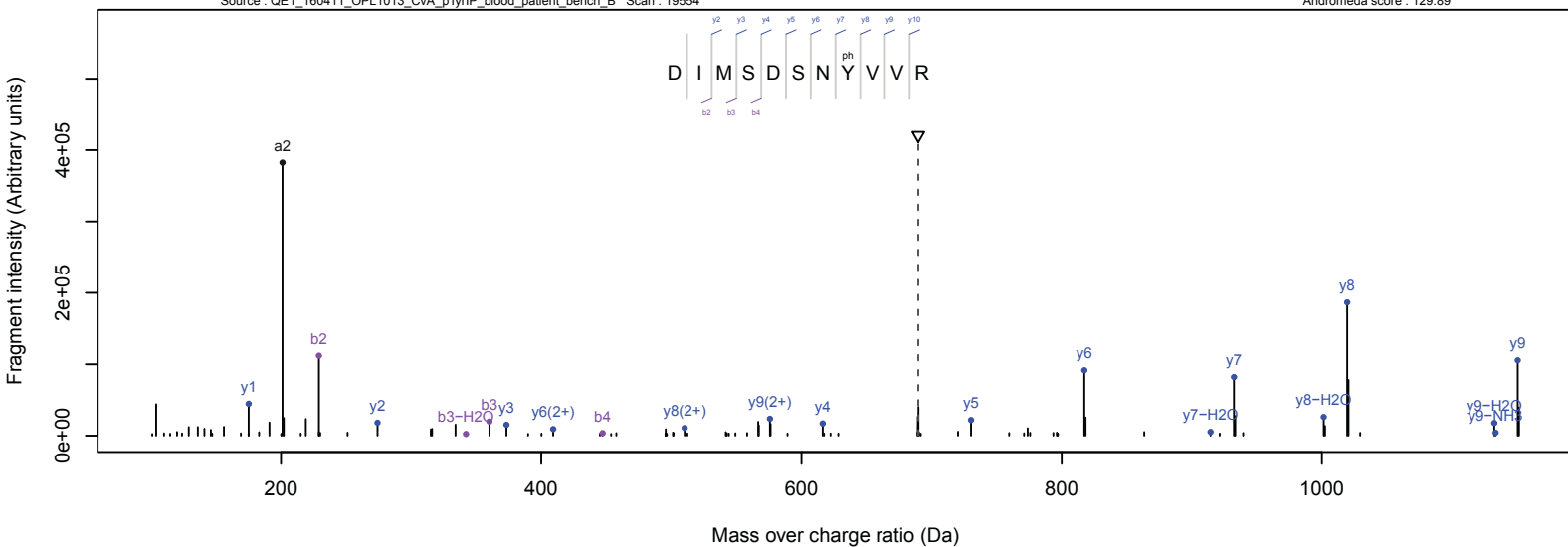
Andromeda score : 130.21



## DIMSDSNY(ph)VVR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 19554

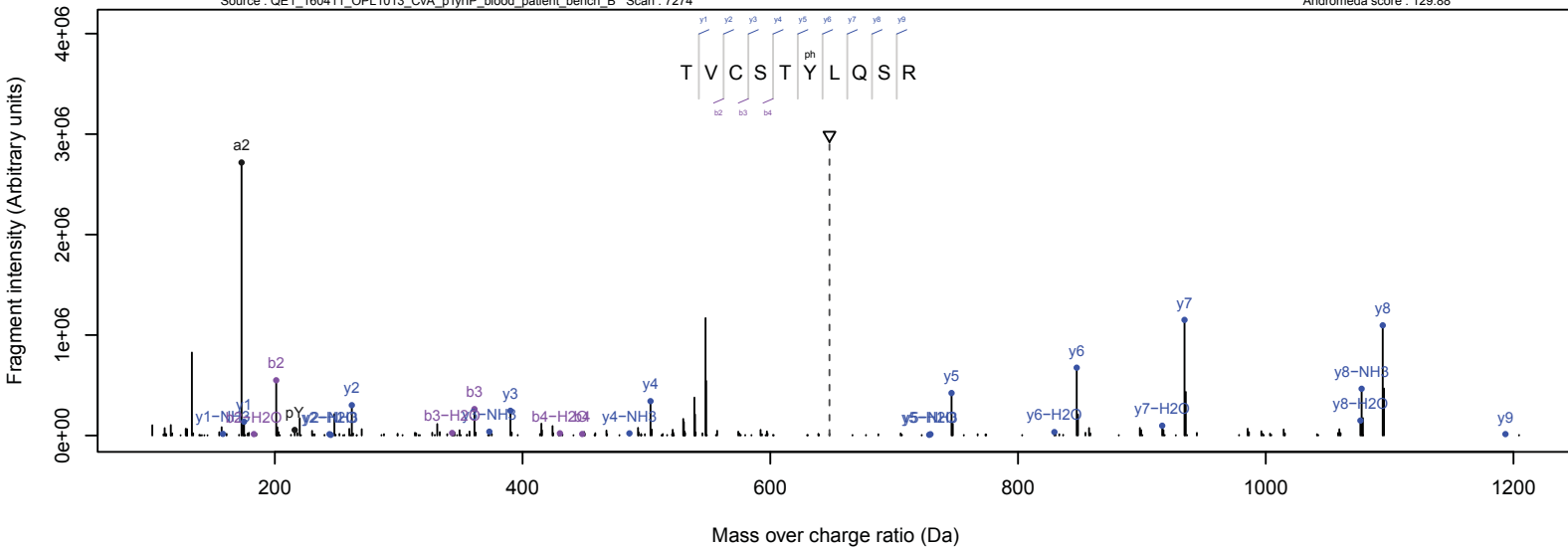
Andromeda score: 129.89



## TVCSSTY(ph)LQSR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 7274

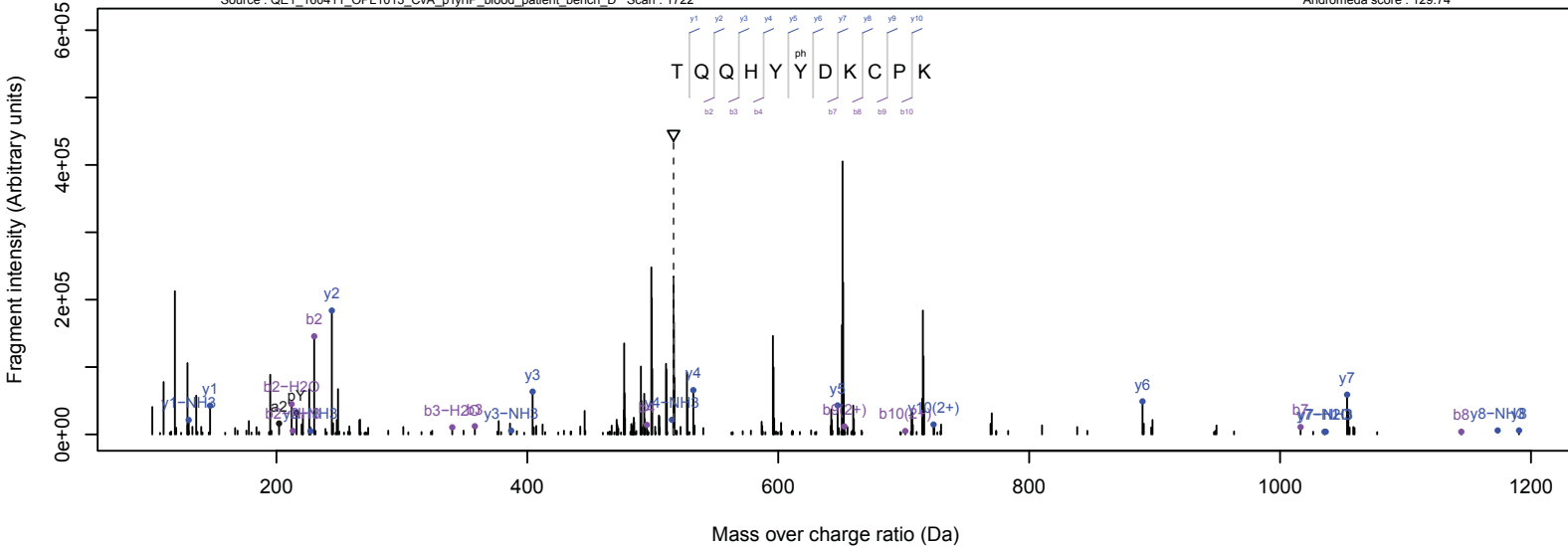
Andromeda score: 129.88



## TQQHYD(ph)KCPK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 1722

Andromeda score: 129.74

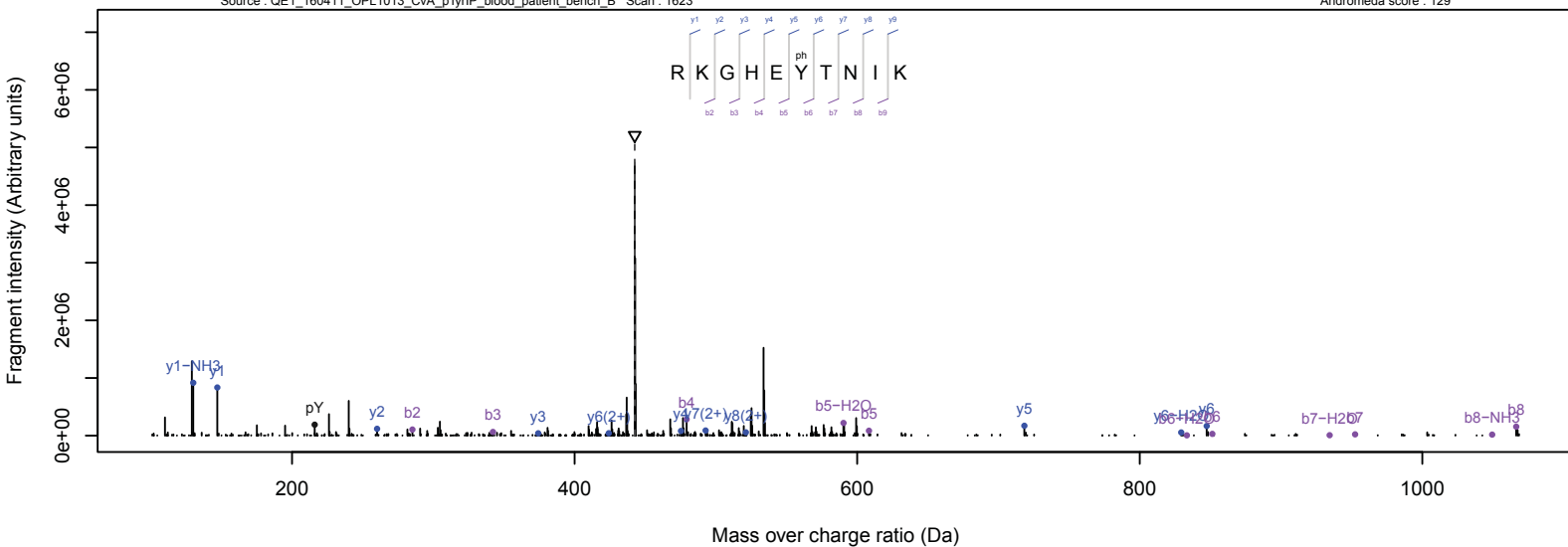




RKGHEY(ph)TNIK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 1623

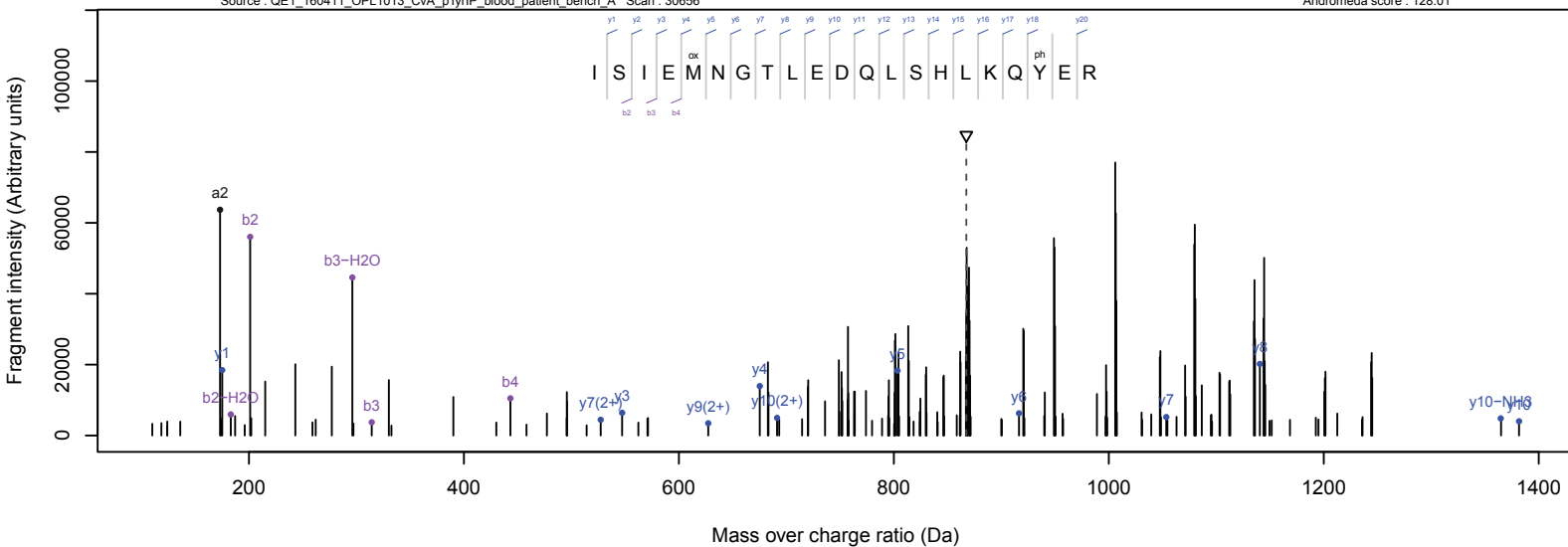
Andromeda score : 129



ISIEM(ox)NGTLEDQLSHLKQY(ph)ER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 30656

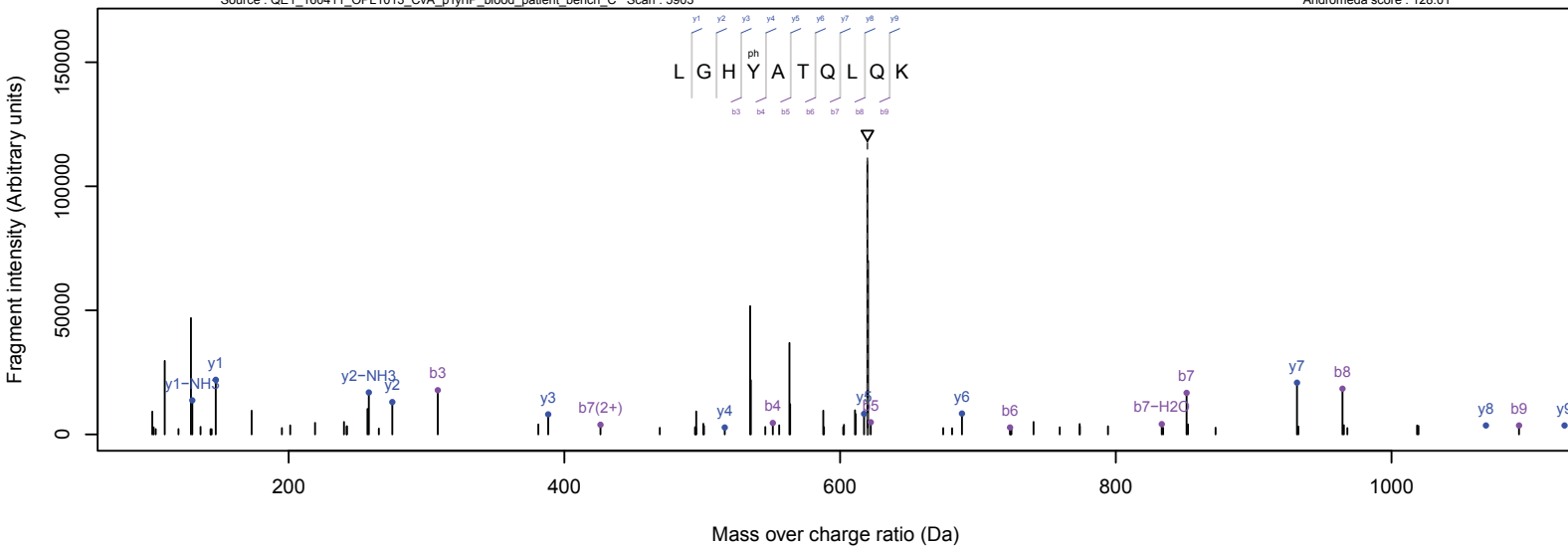
Andromeda score : 128.01



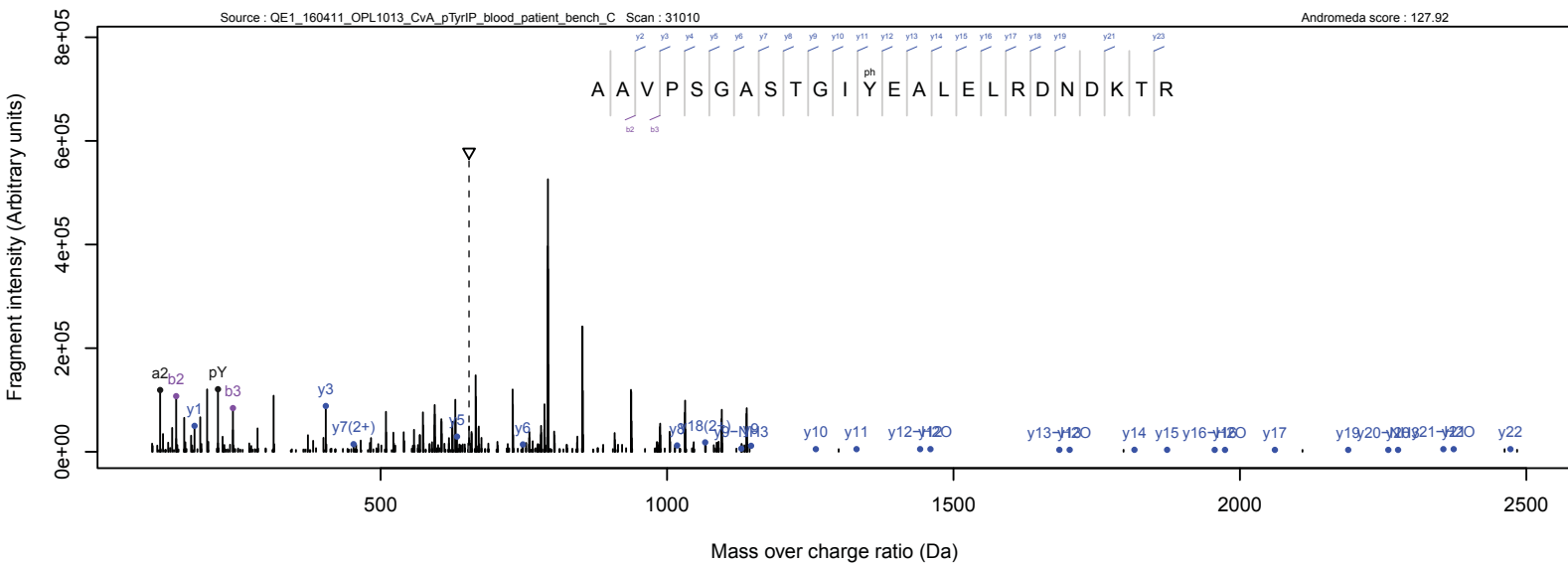
LGHY(ph)ATLQK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 5903

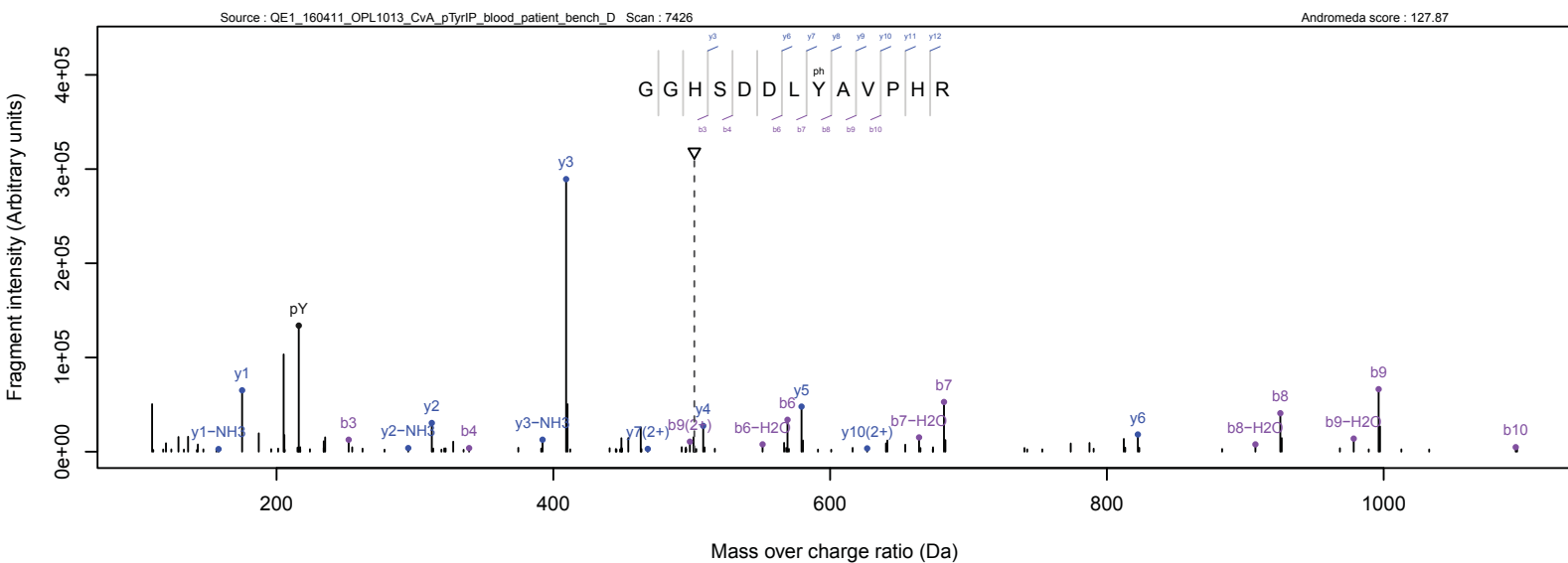
Andromeda score : 128.01



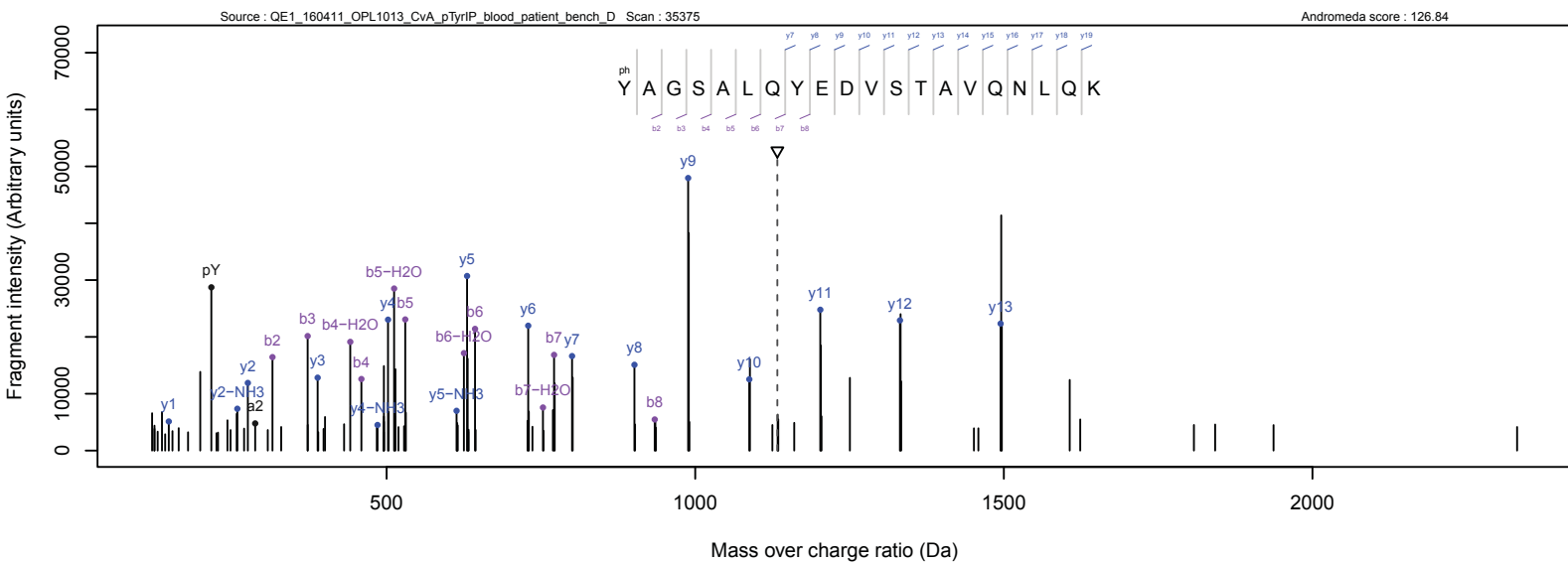
AAVPSGASTGIY(ph)EALELRDNDKTR-4+



GGHSDDL(ph)AVPHR-3+



Y(ph)AGSALQYEDVSTAVQNLQK-2+

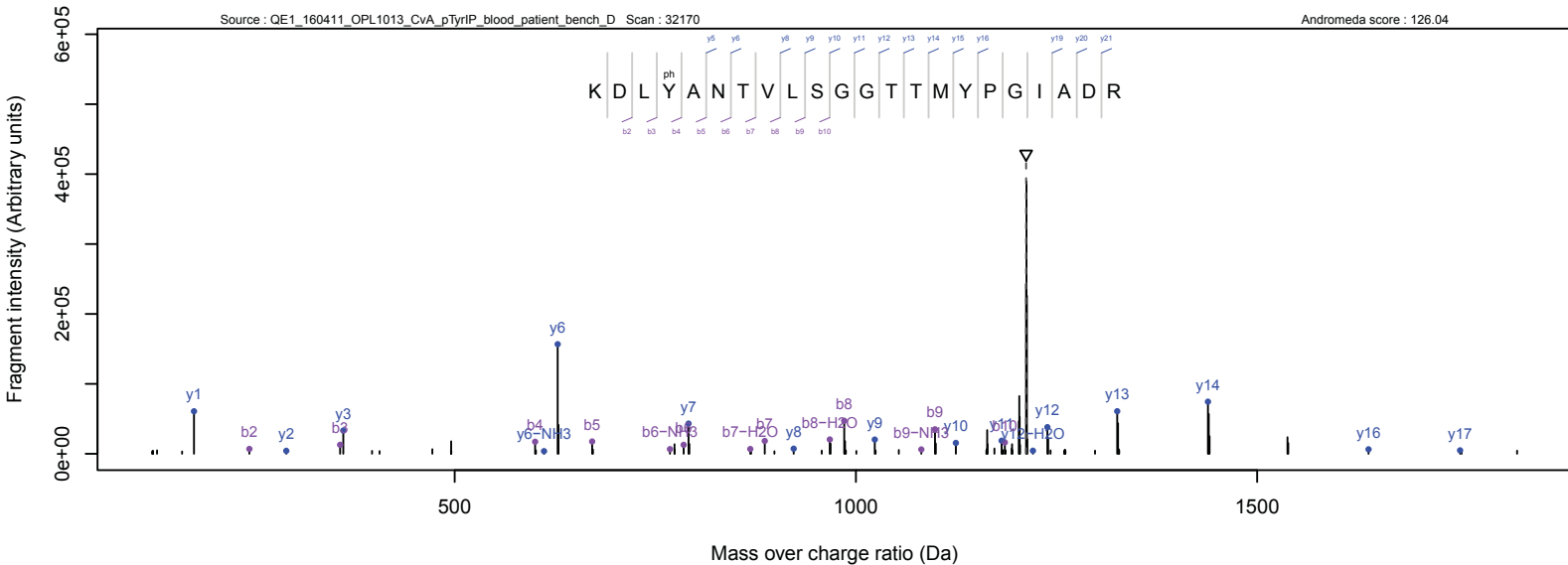




**KDLY(ph)ANTVLSGGTTMYPGIADR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 32170

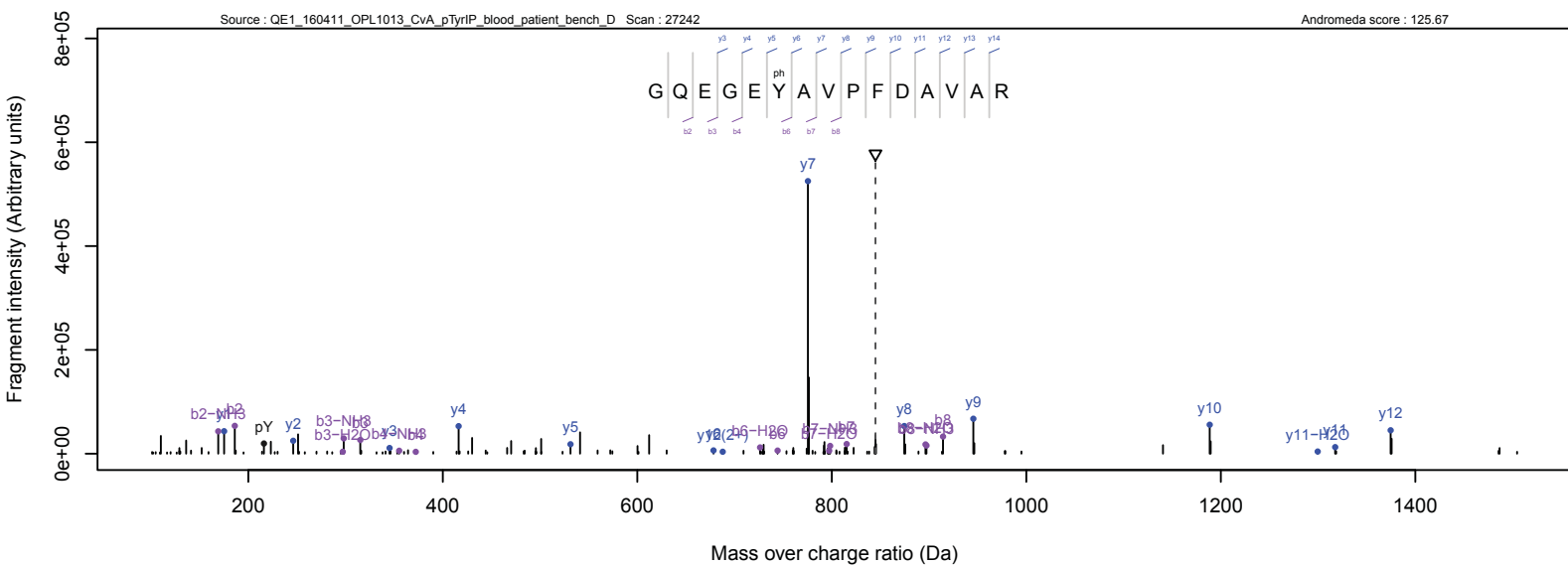
Andromeda score : 126.04



**GQEGEY(ph)AVPFDAVAR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 27242

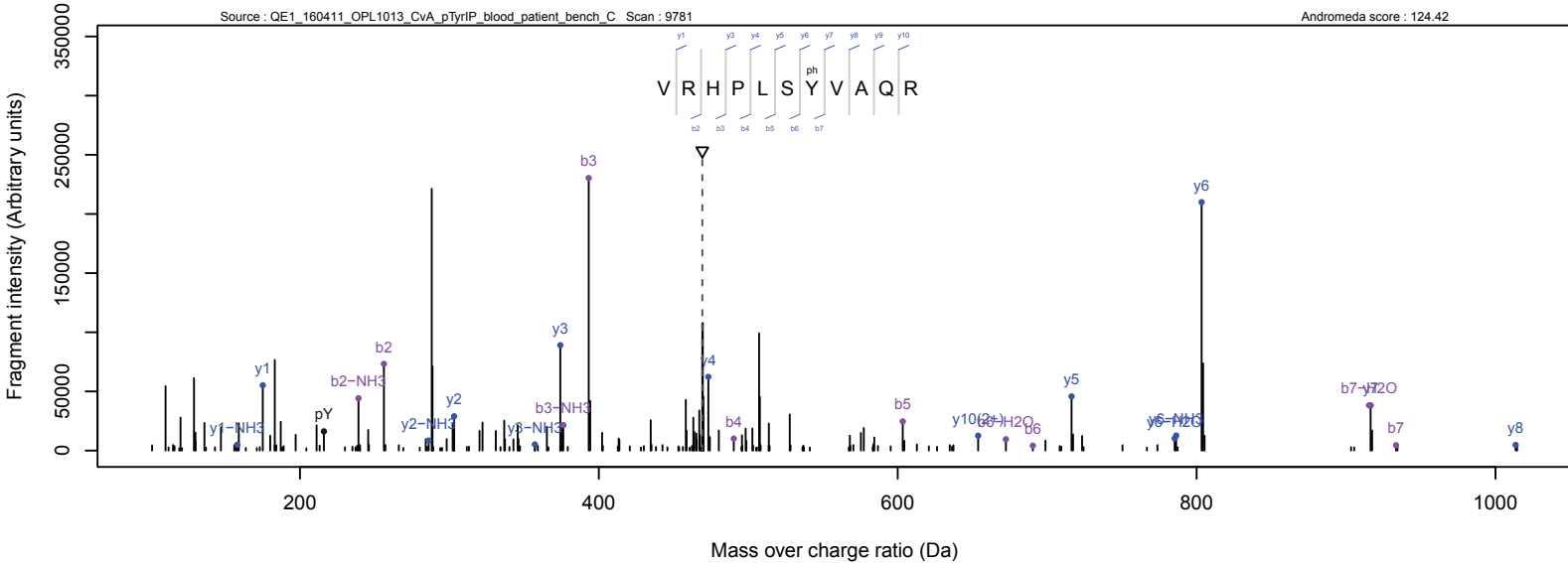
Andromeda score : 125.67



**VRHPLSY(ph)VAQR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 9781

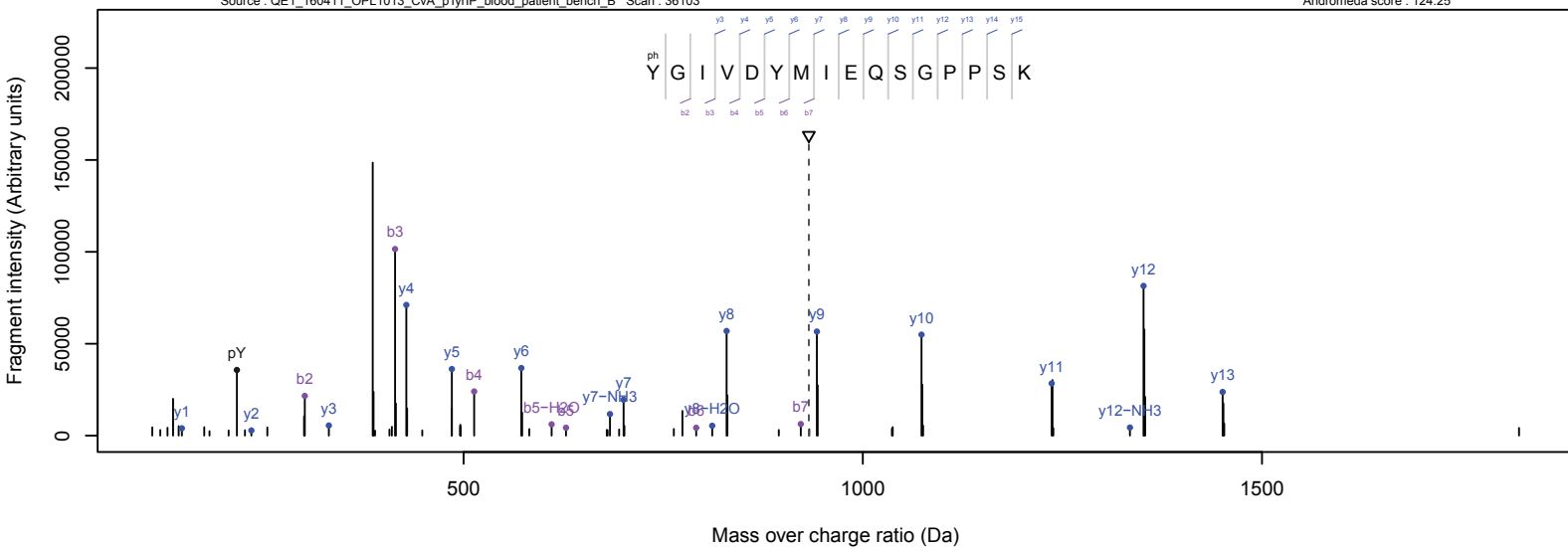
Andromeda score : 124.42



Y(ph)GIVDYMI EQSGPPSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 36103

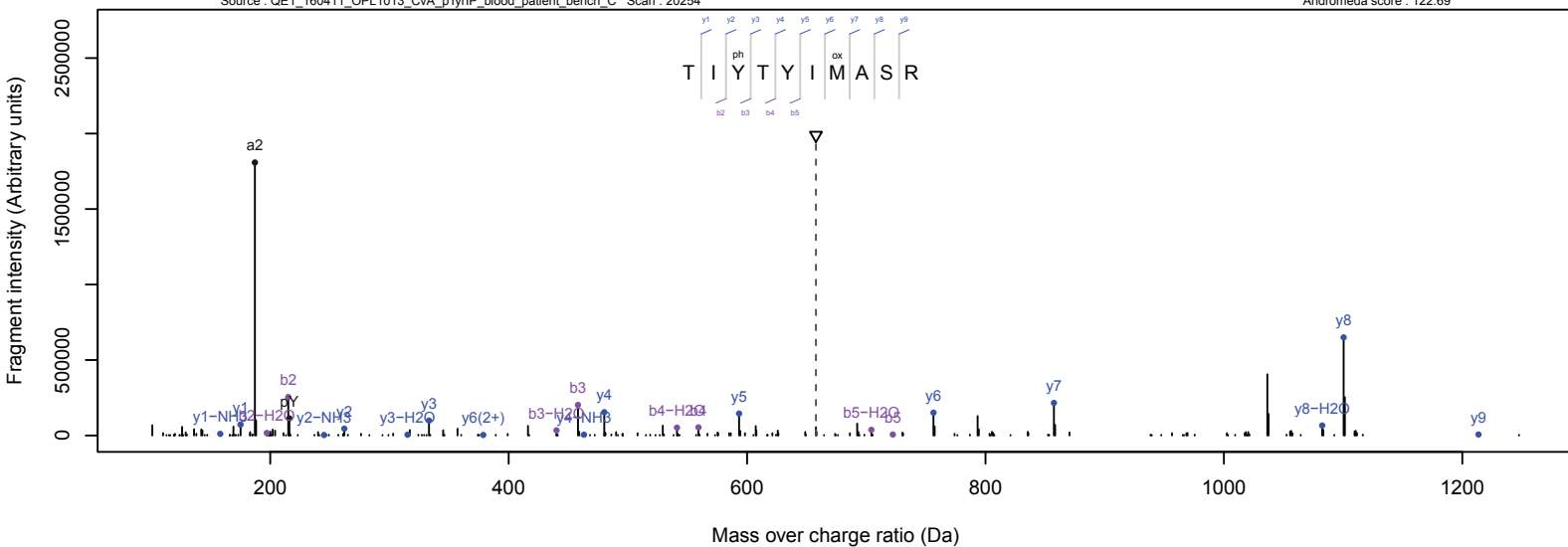
Andromeda score : 124.25



TIY(ph)TYIM(ox)ASR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 20254

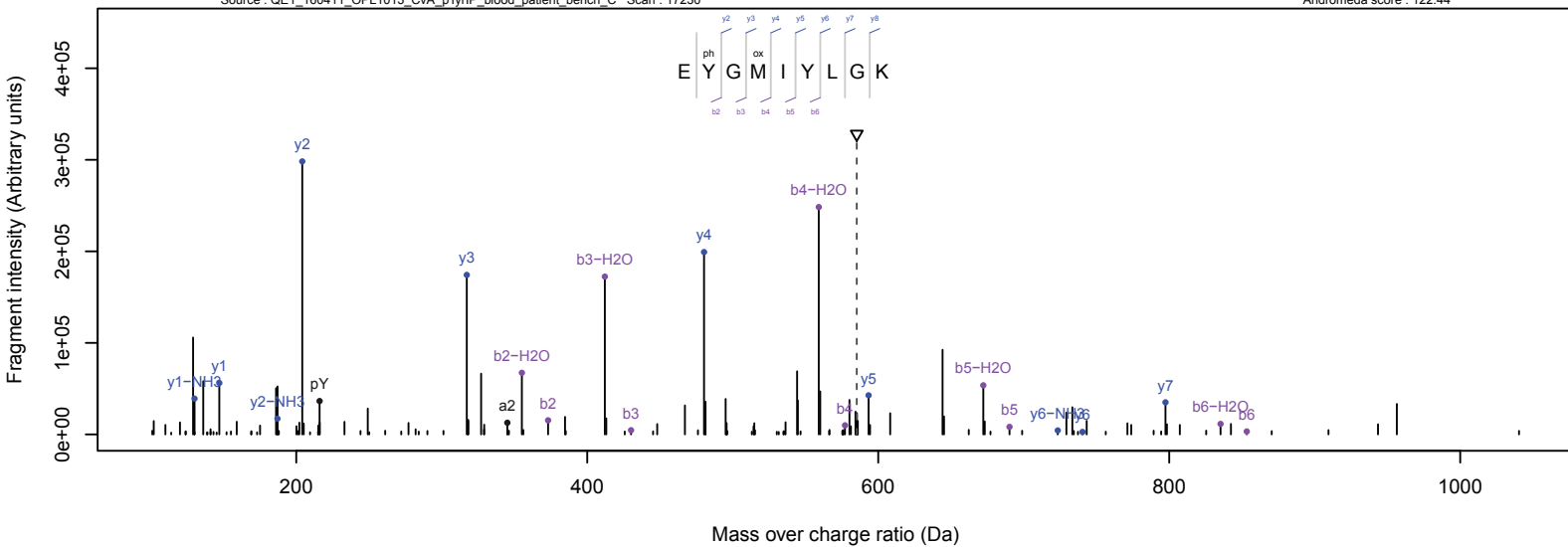
Andromeda score : 122.69



EY(ph)GM(ox)IYL GK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 17230

Andromeda score : 122.44

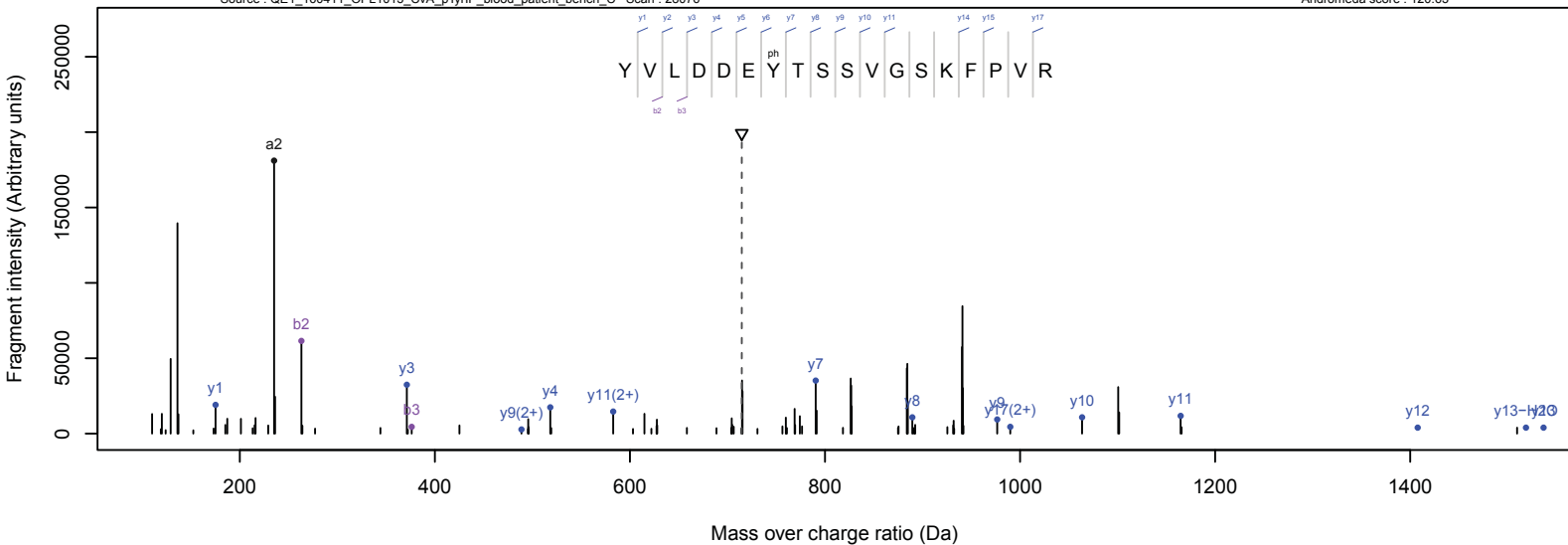




**YVLDDDEY(ph)TSSVGSKFPVR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 28076

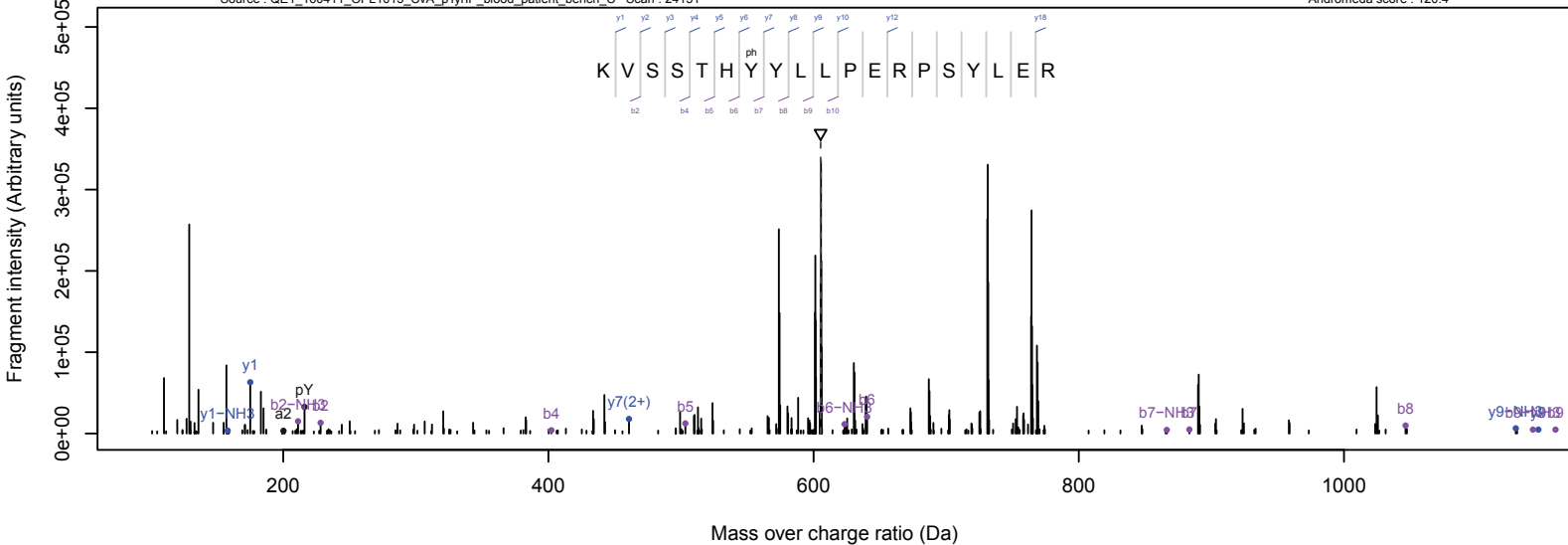
Andromeda score: 120.63



**KVSSTHY(ph)YLLPERPSYLER-4+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 24151

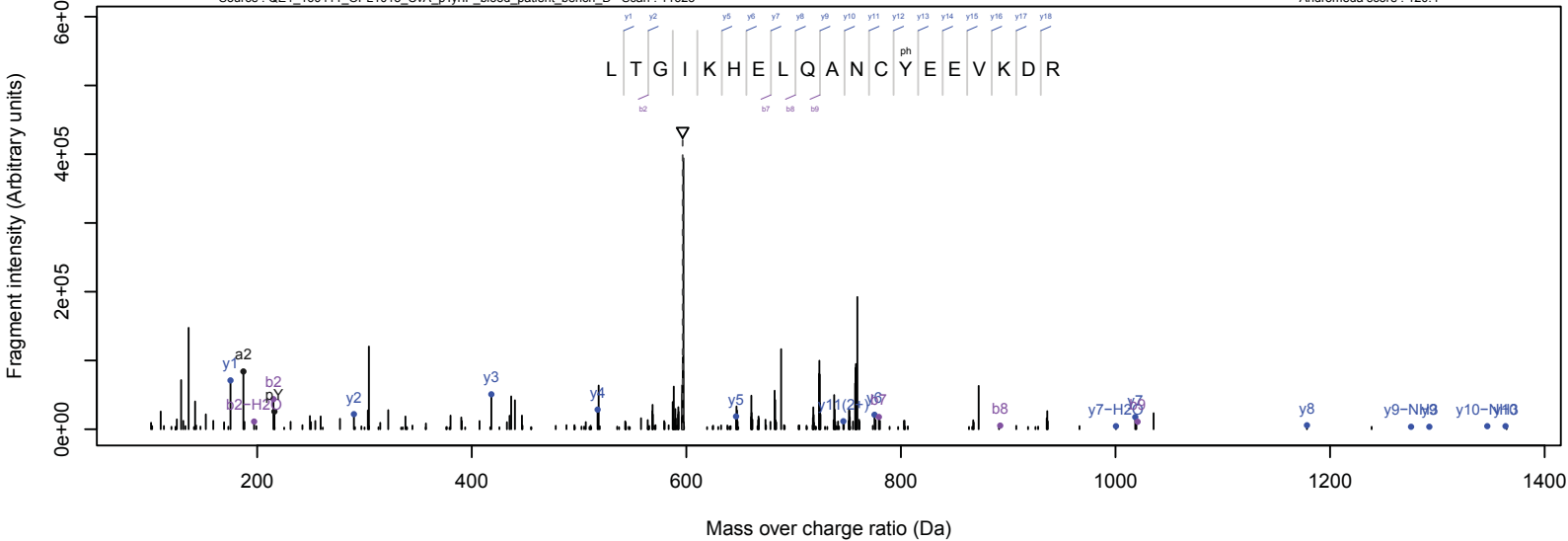
Andromeda score: 120.4



**LTGIKHELQANCY(ph)EEVKDR-4+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 11625

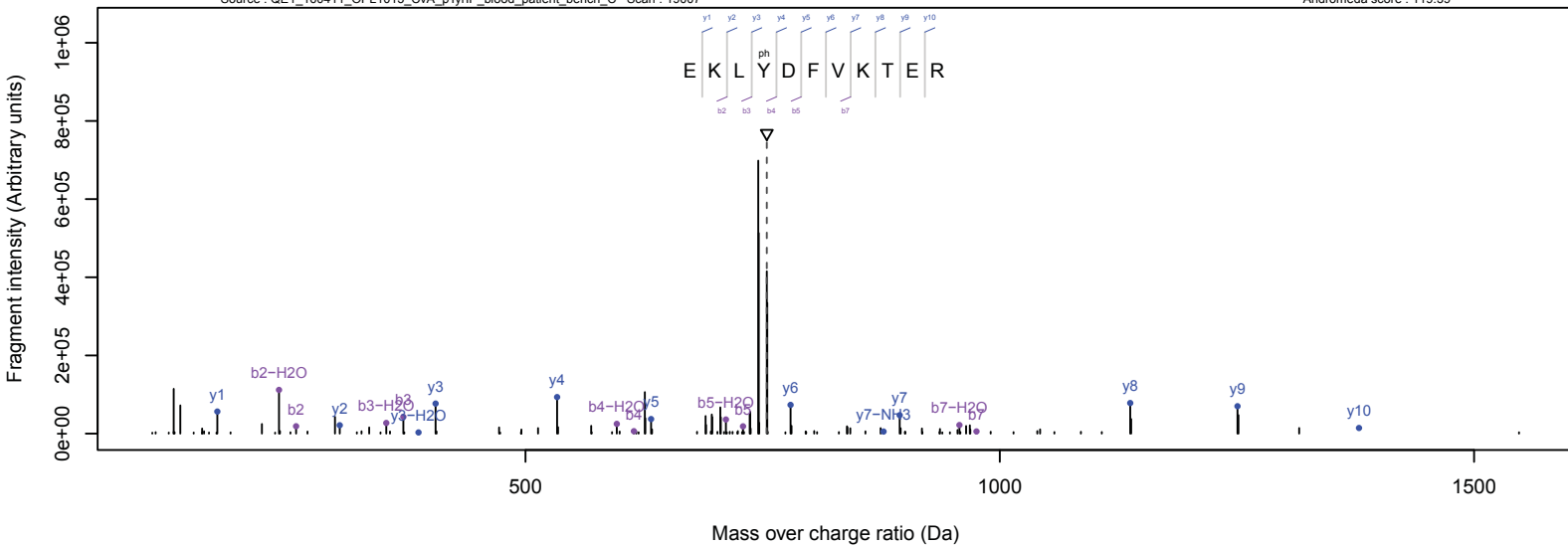
Andromeda score: 120.4



**EKLY(ph)DFVKTER-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 19007

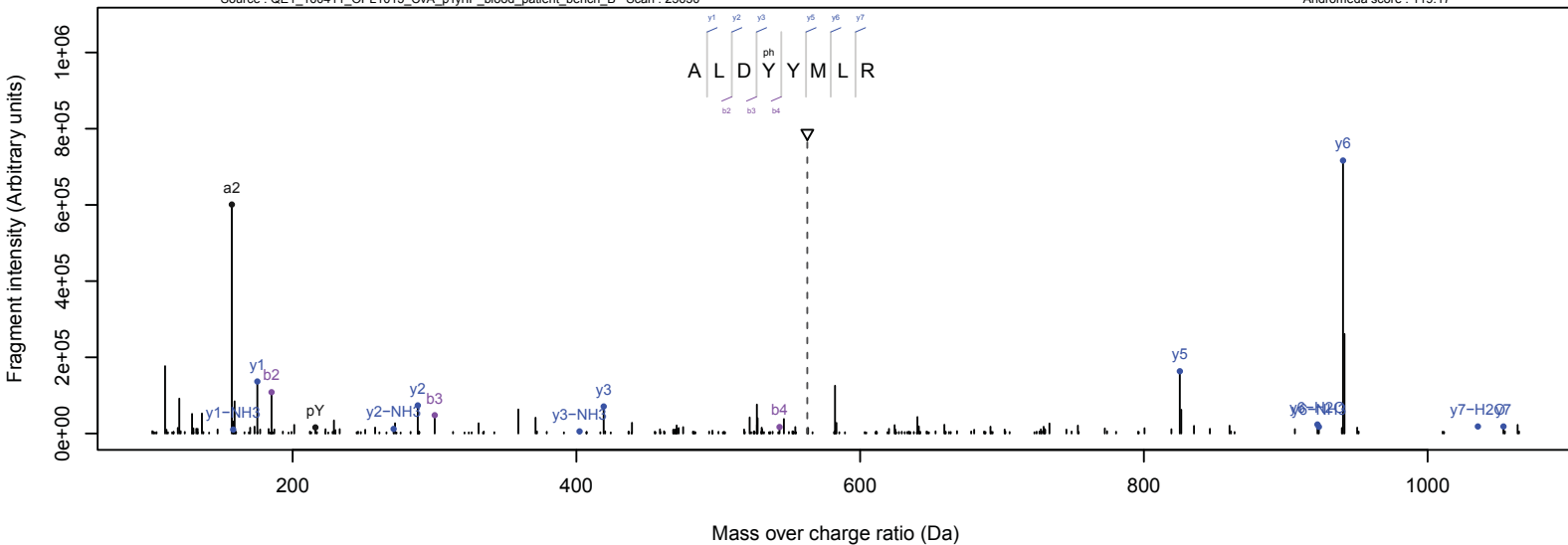
Andromeda score : 119.39



**ALDY(ph)YMLR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 25630

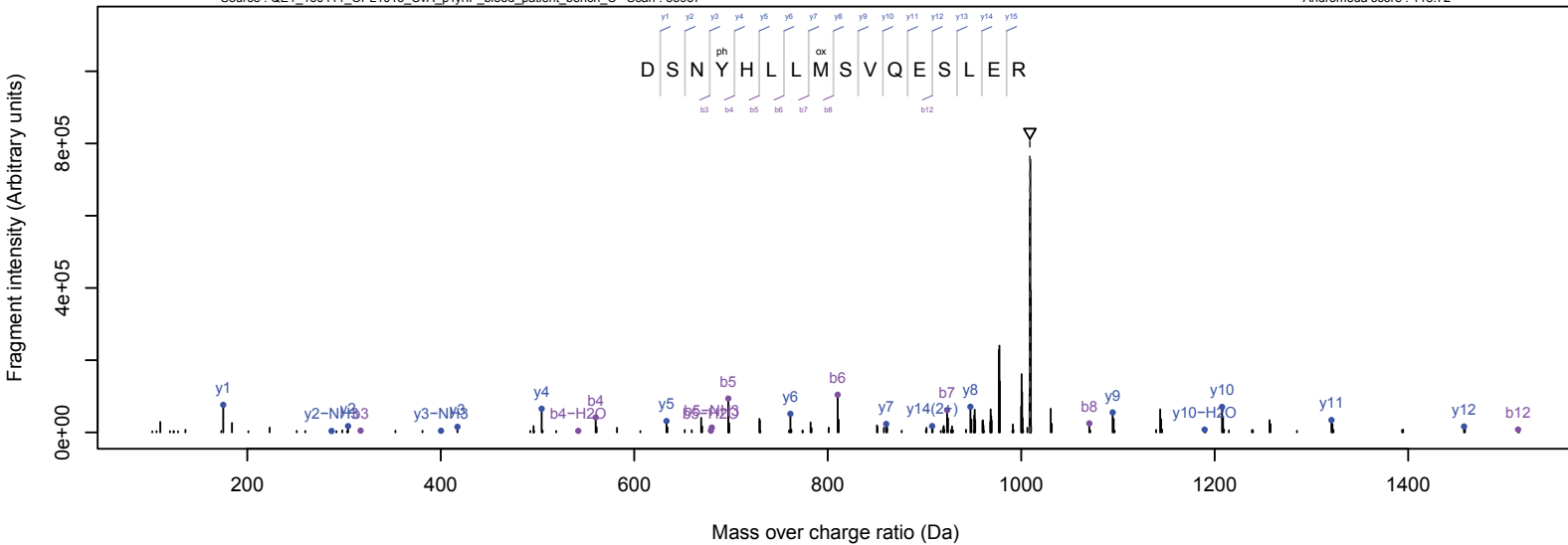
Andromeda score : 119.17



**DSNY(ph)HLLM(ox)SVQESLER-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 35937

Andromeda score : 118.72

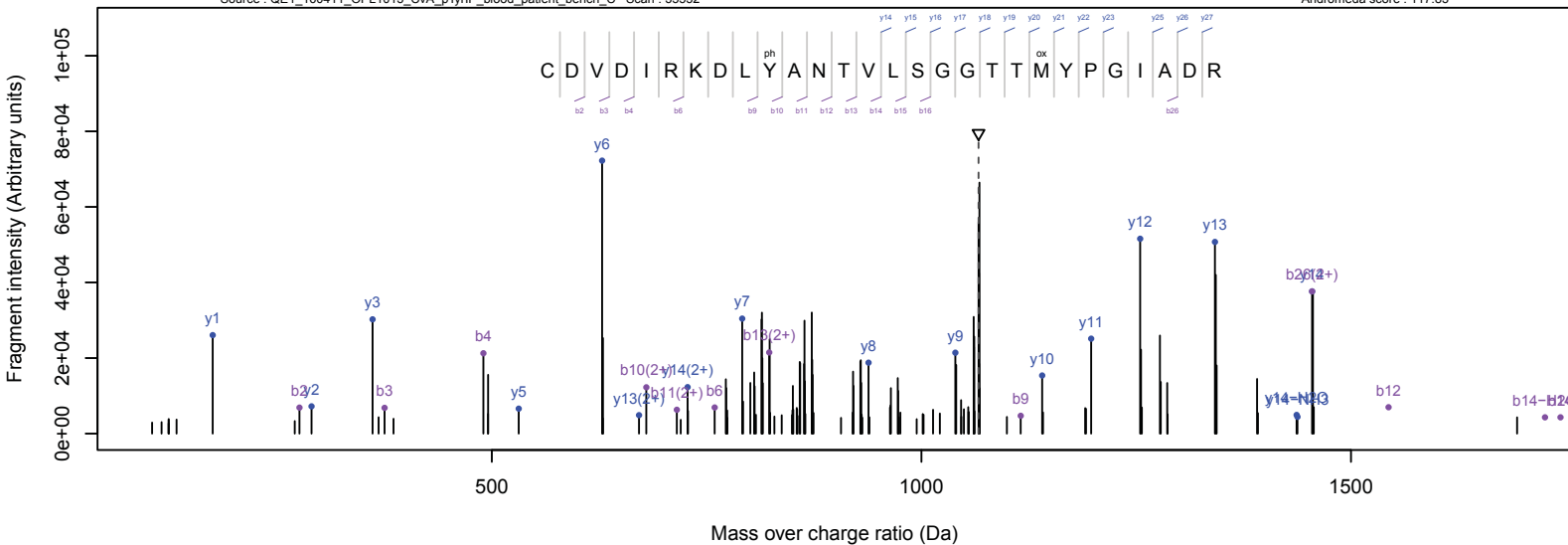




CDVDIRKDL(ph)ANTVLSGGTTM(ox)YPGIADR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 35532

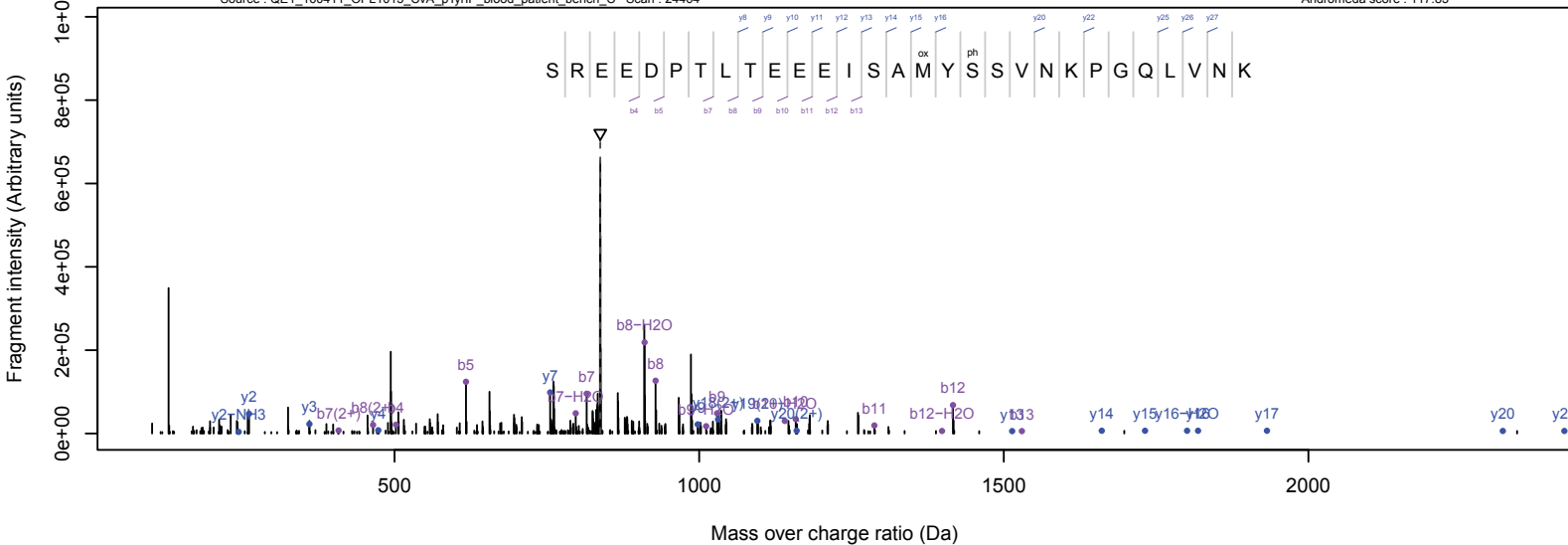
Andromeda score: 117.83



SREEDPTLTEEEISAM(ox)YS(ph)SVNKPGLVNK-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 24464

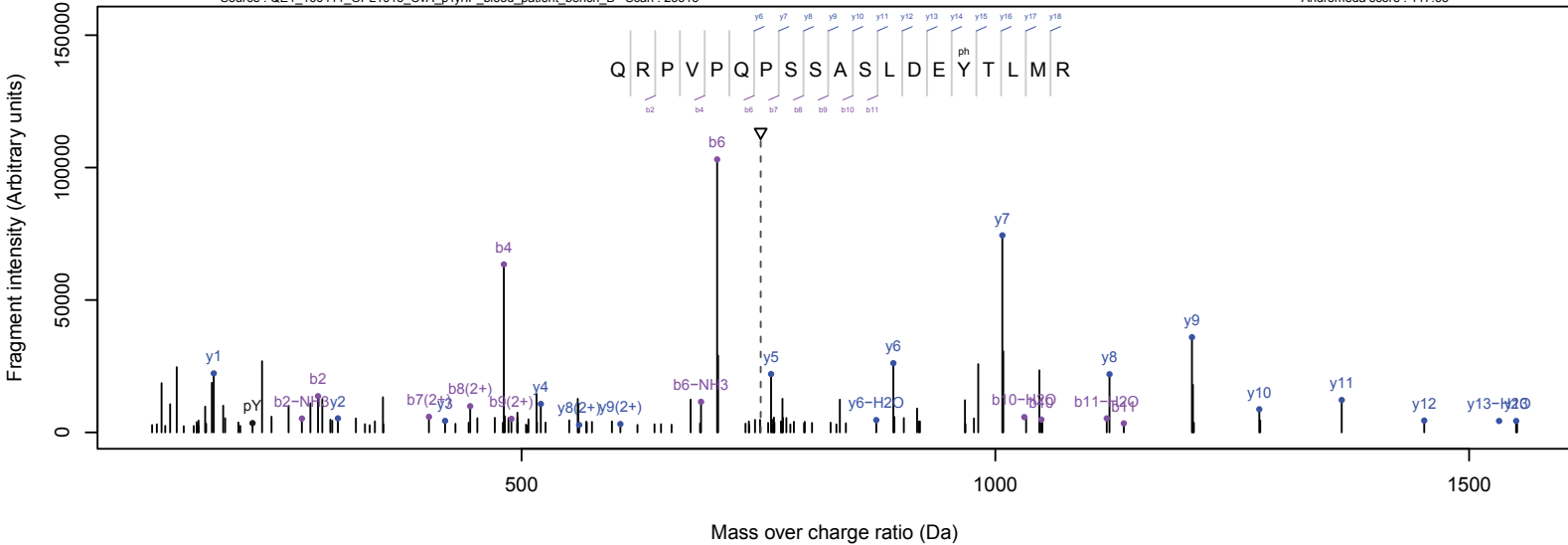
Andromeda score: 117.83



QRPVPQPSSASLDEYTLMR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 23313

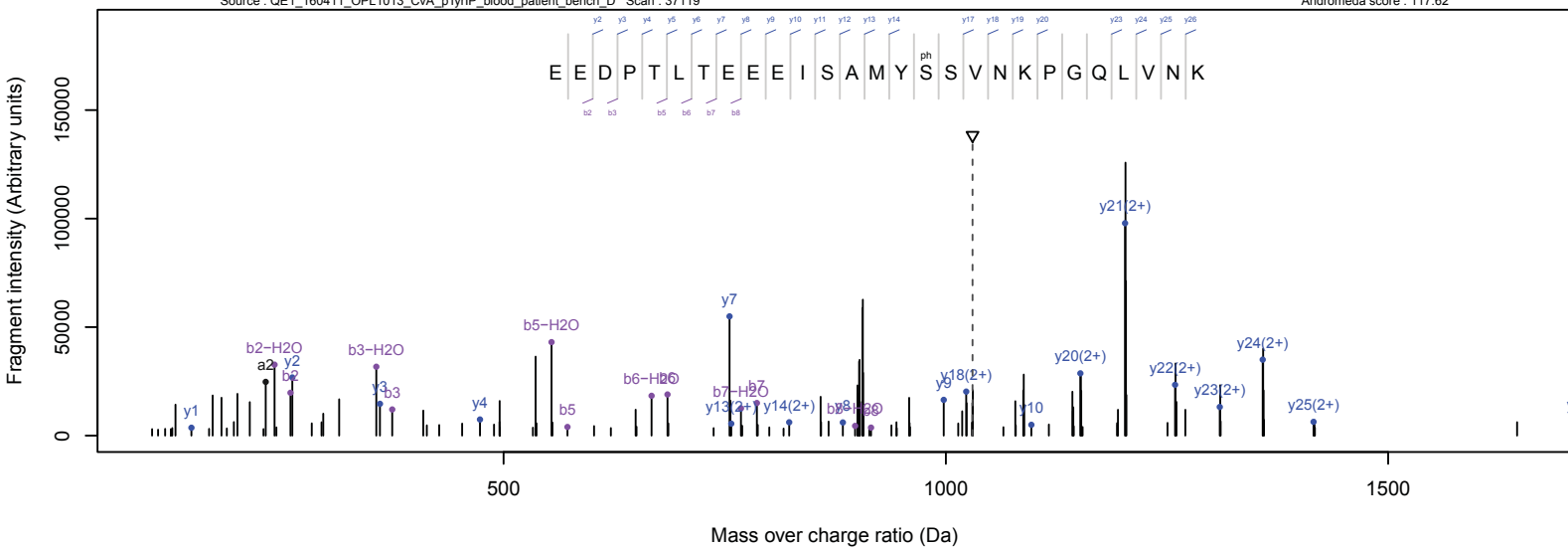
Andromeda score: 117.65



EEDPTLTEEISAMYS(ph)SVNKPGQLVNK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 37119

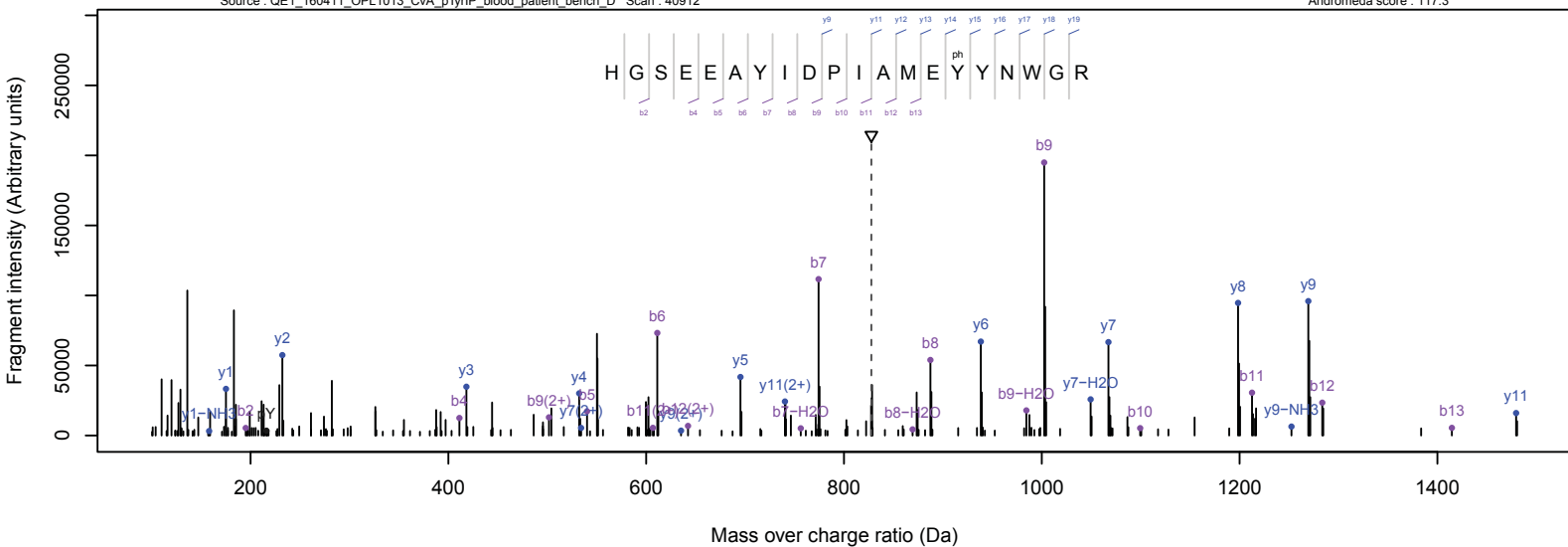
Andromeda score : 117.62



HGSEEA YIDPIAMEY(ph)YNWGR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 40912

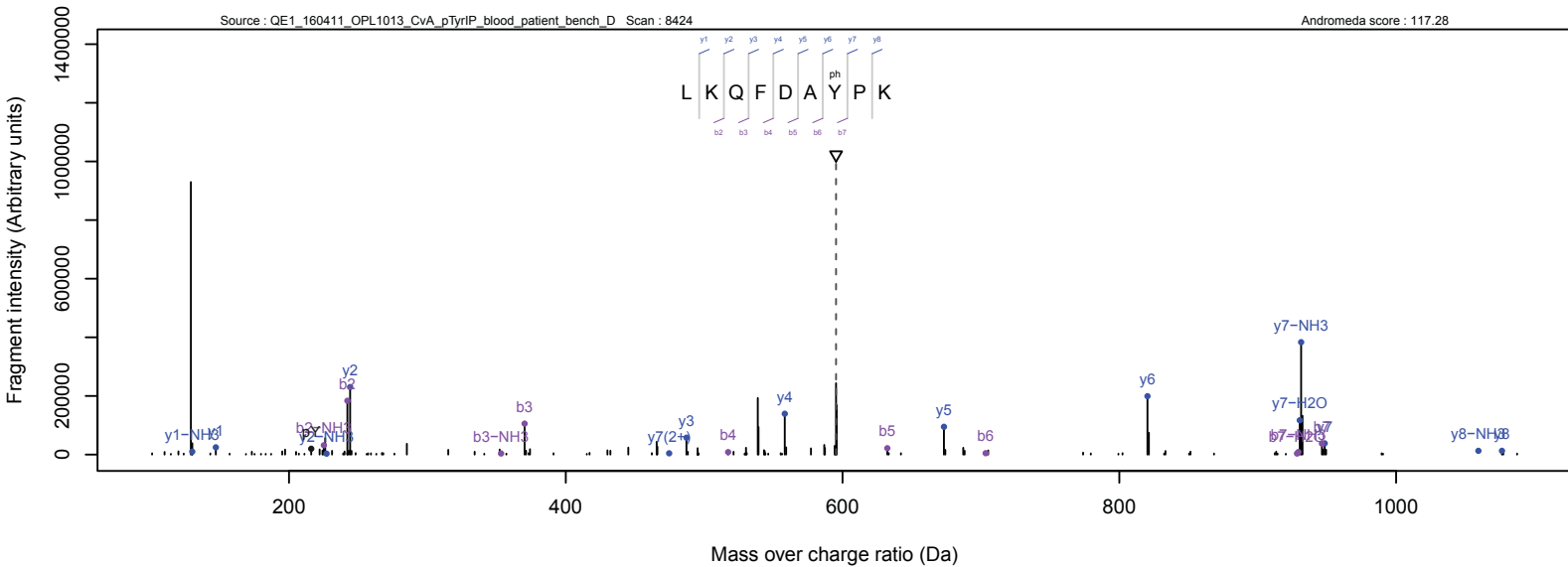
Andromeda score : 117.3



LKQFDAY(ph)PK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 8424

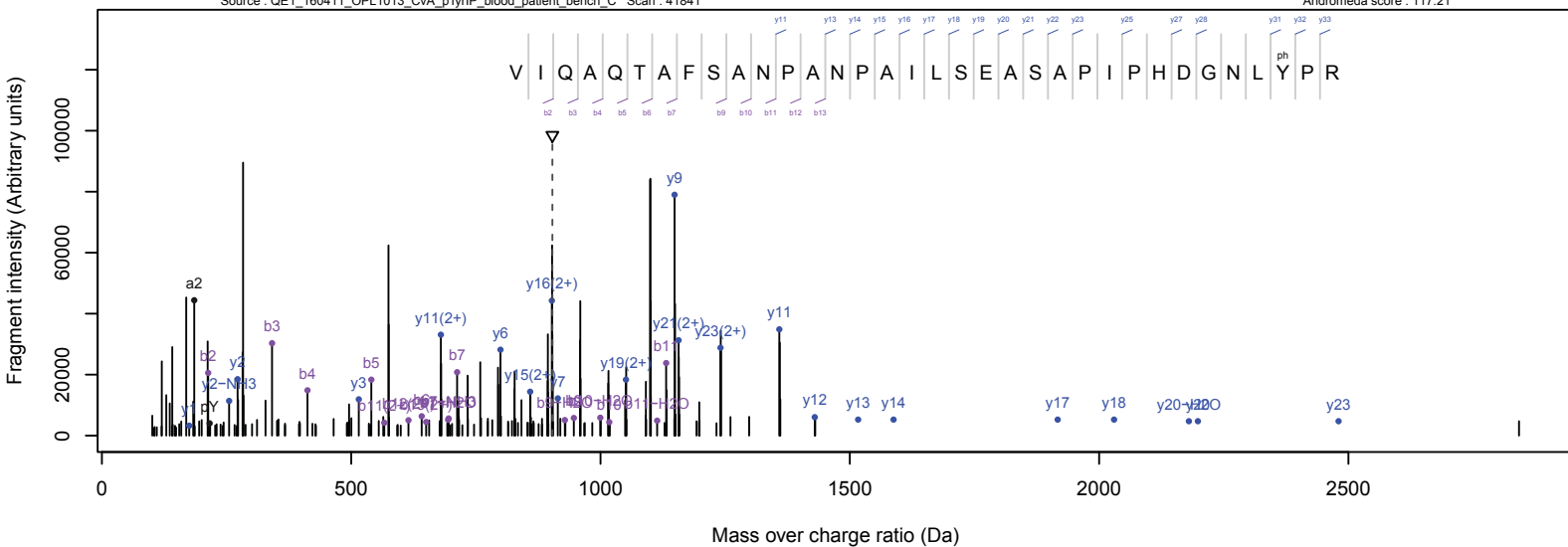
Andromeda score : 117.28



VIQAQTAFSANPANPAILSEASAPIPHDGNLYPR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 41841

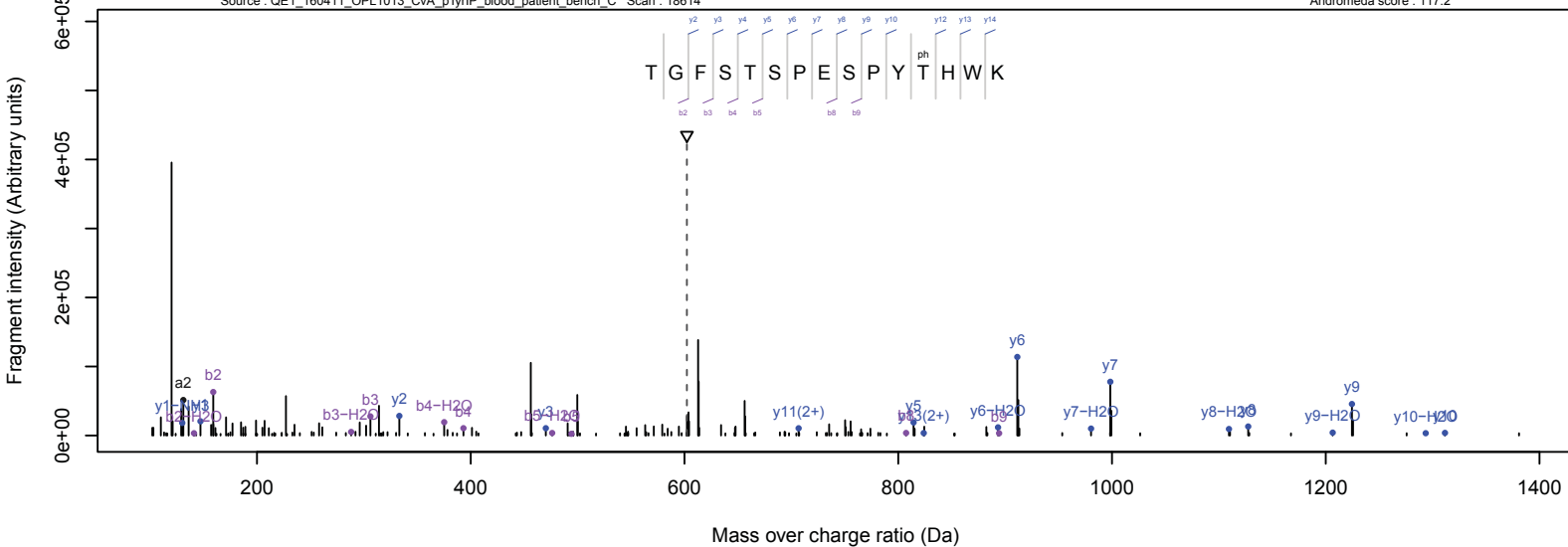
Andromeda score : 117.21



TGFSTSPESPYTHWK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18614

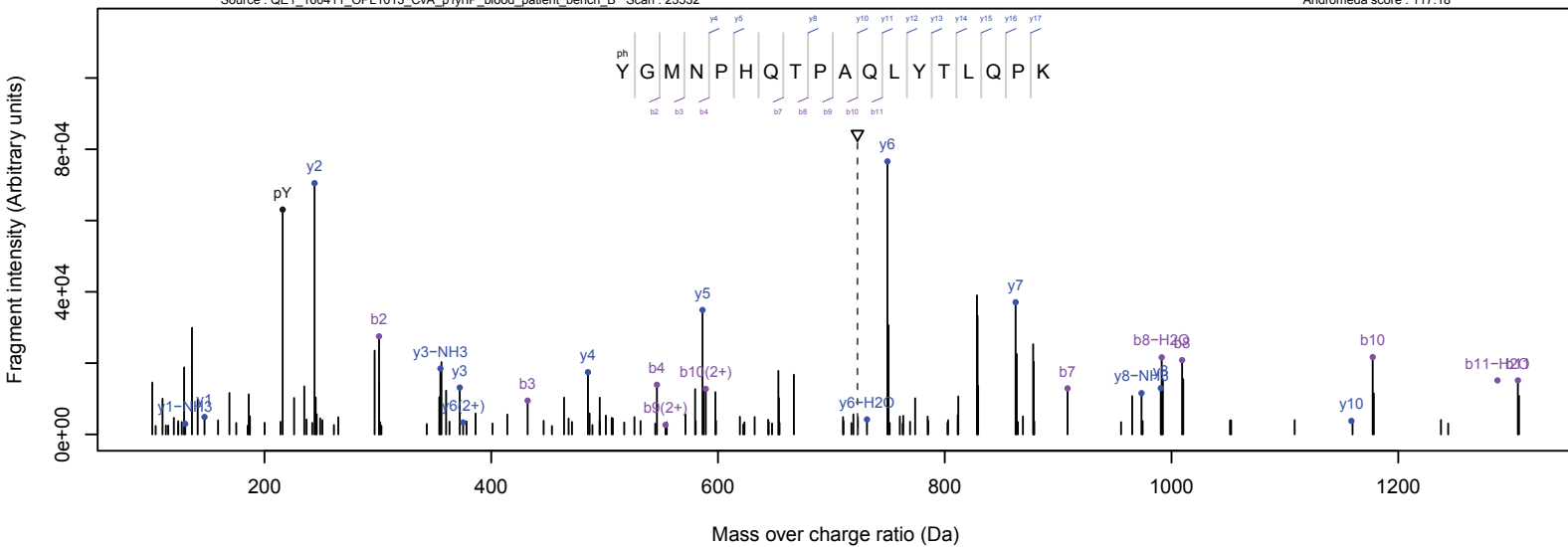
Andromeda score : 117.2



Y(ph)GMNPHQTPAQLYTLQPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 23532

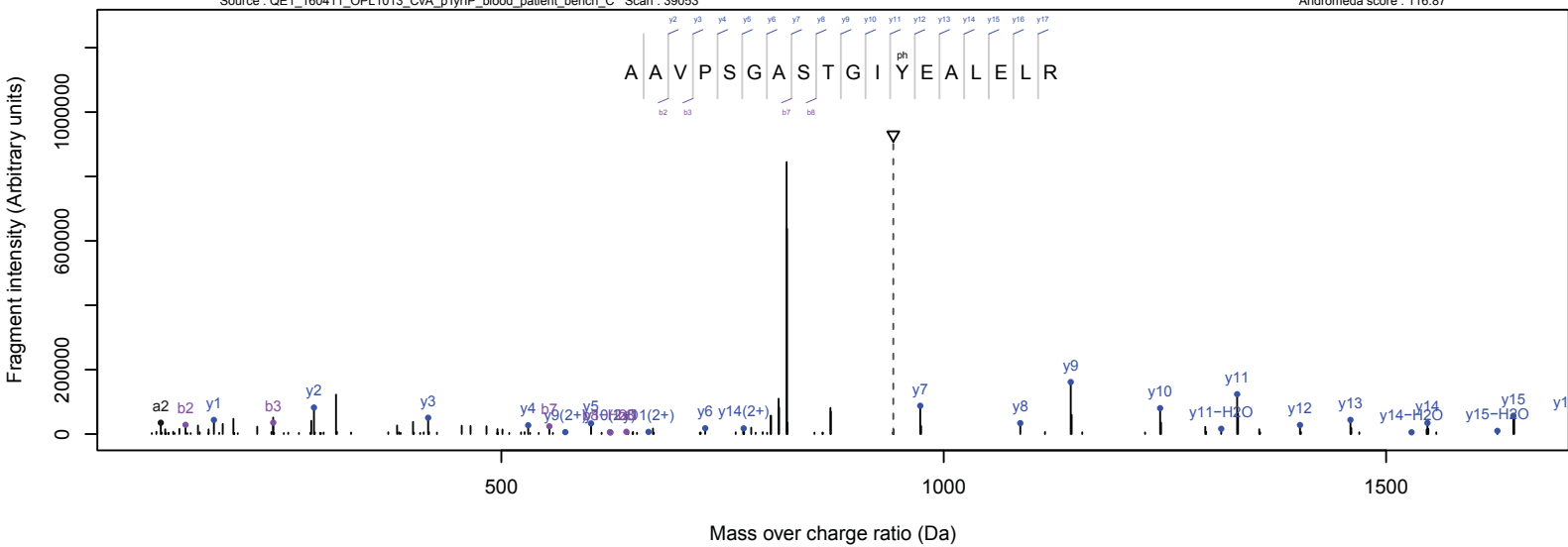
Andromeda score : 117.18



AAVPSGASTGIY(ph)EALELR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 39053

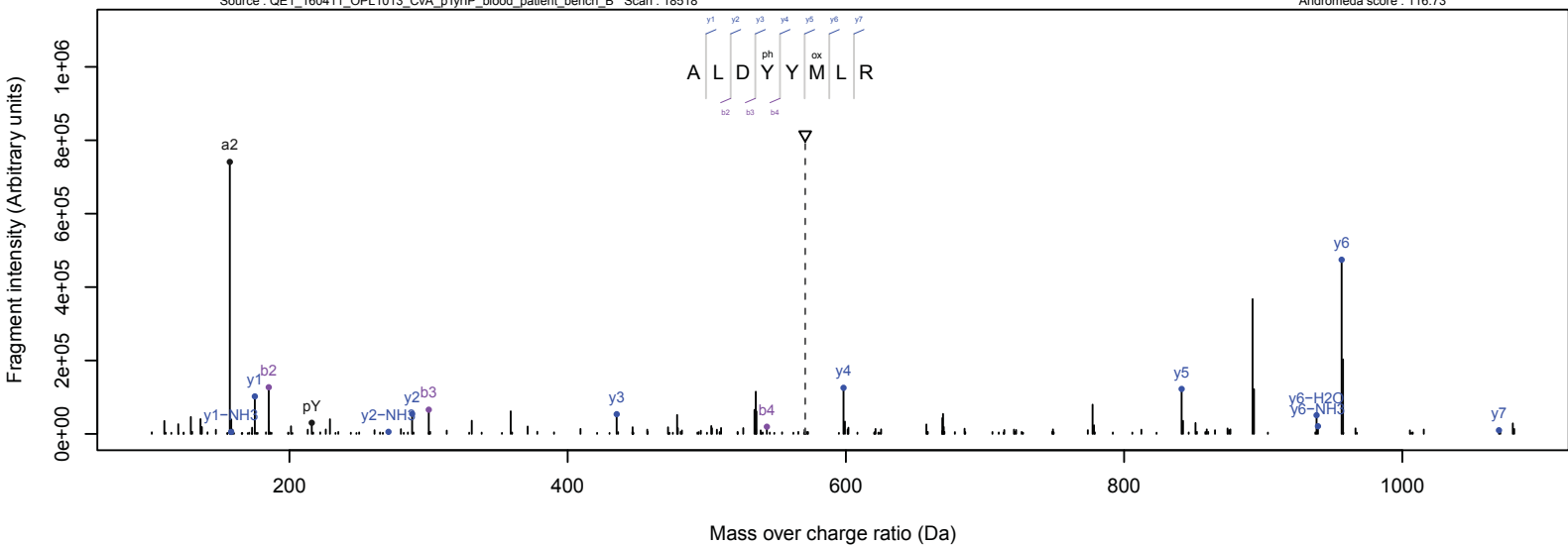
Andromeda score : 116.87



ALDY(ph)YM(ox)LR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 18518

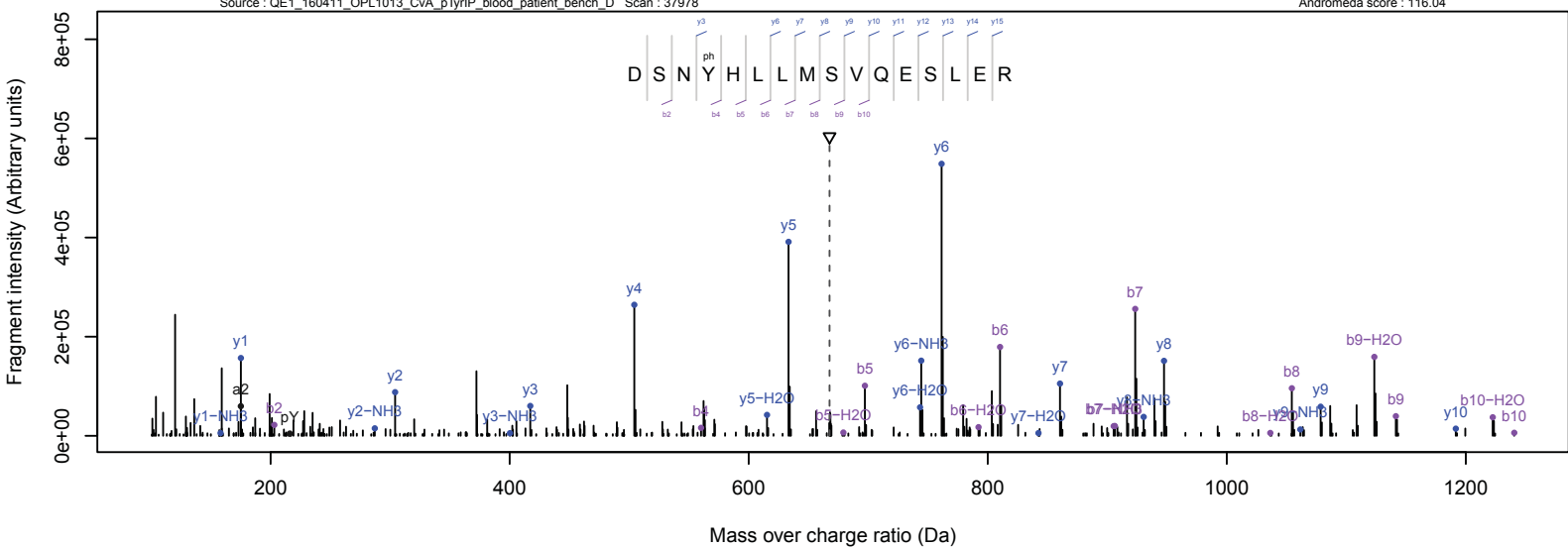
Andromeda score : 116.73



DSNY(ph)HLLMSVQESLER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 37978

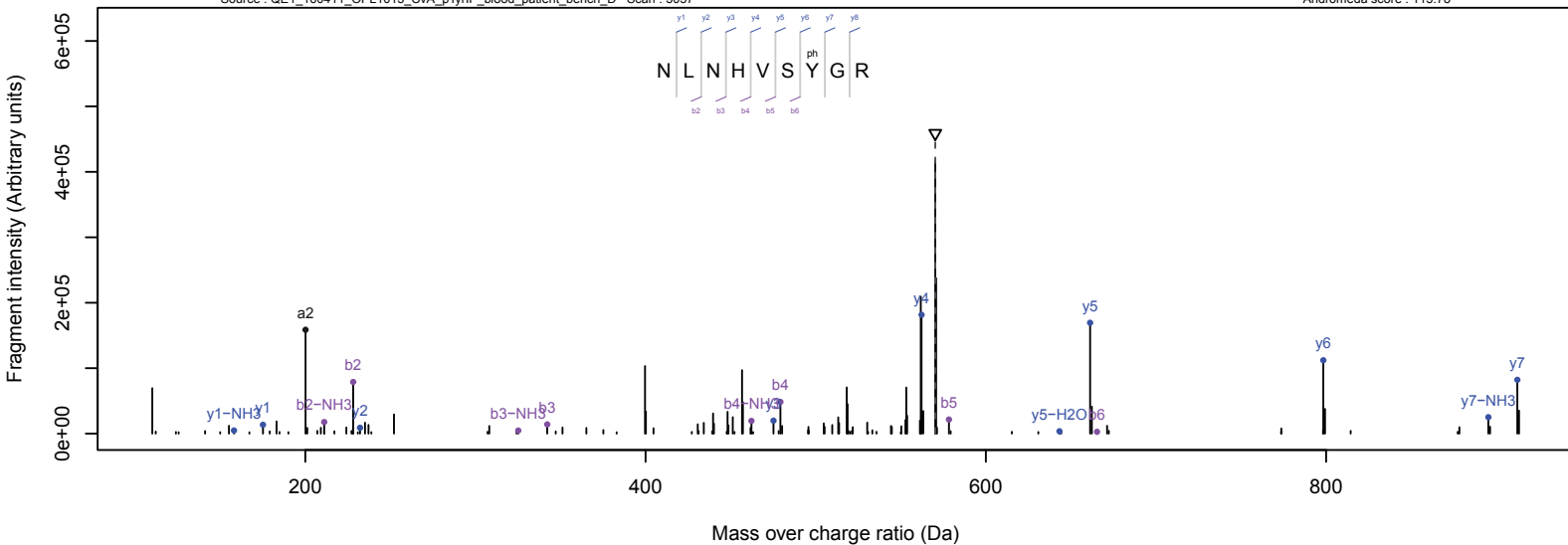
Andromeda score : 116.04



## NLNHVSY(ph)GR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 3057

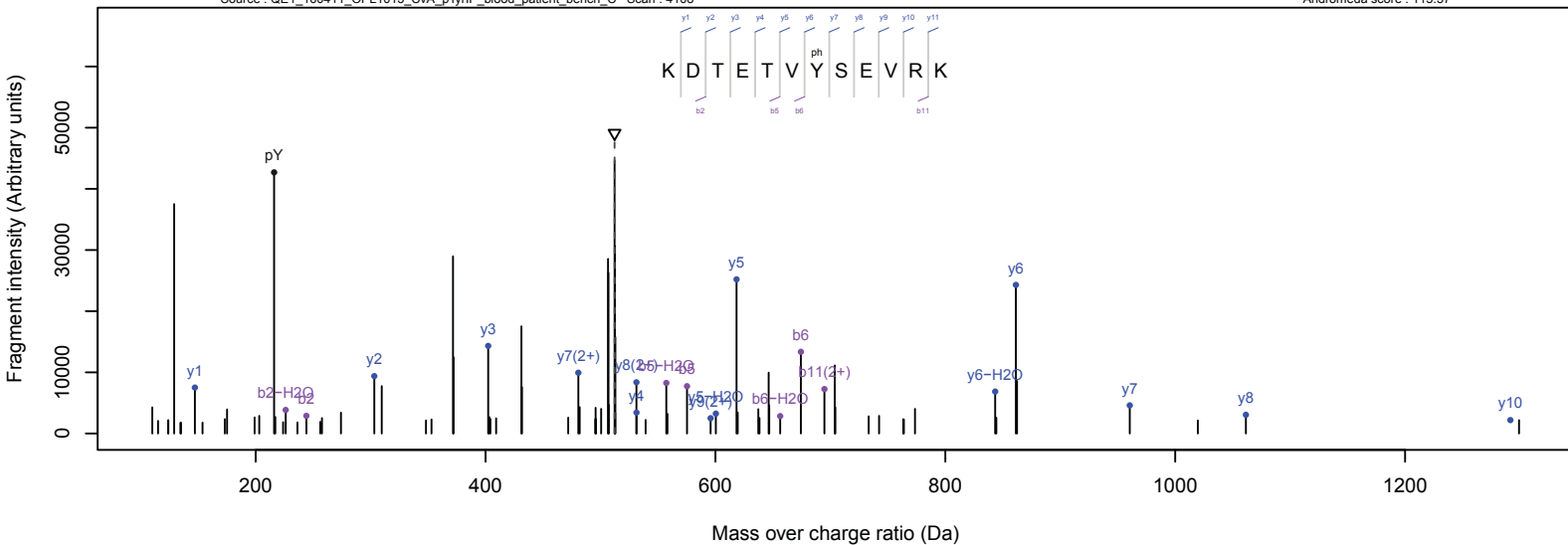
Andromeda score: 115.78



## KDTEVY(ph)SEVRK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 4108

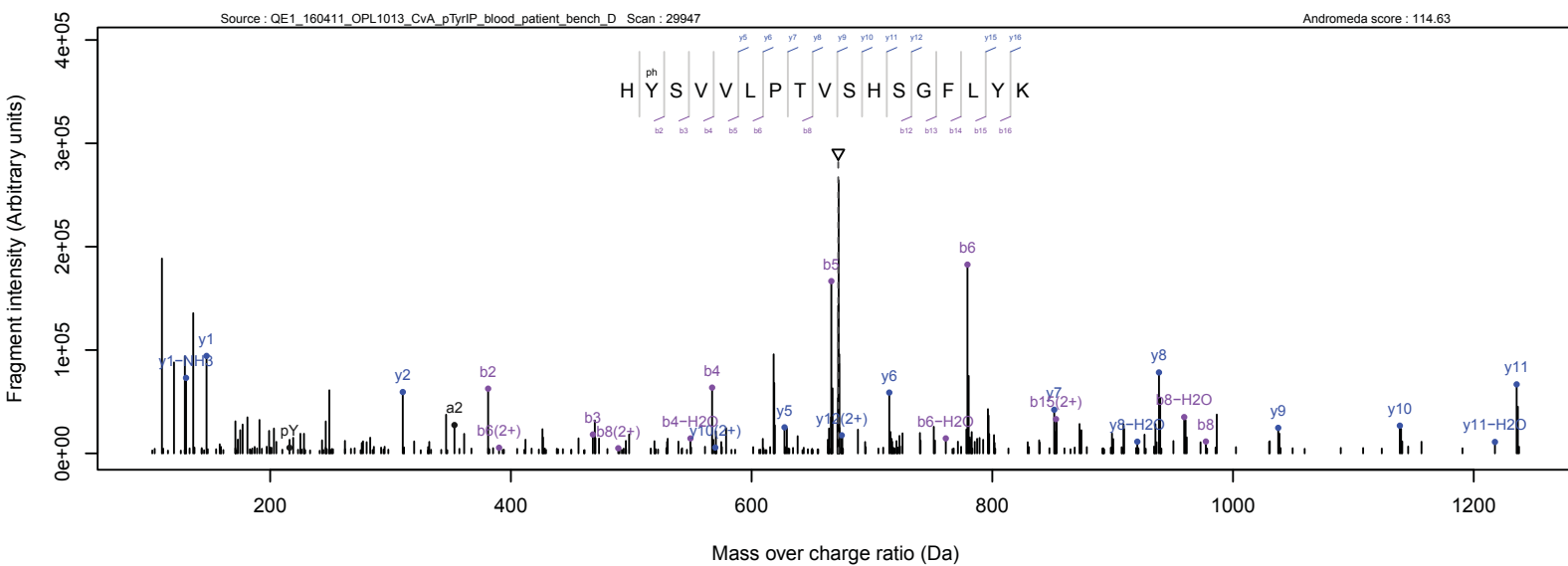
Andromeda score: 115.57



## HY(ph)SVLVPTVSHSGFLYK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 29947

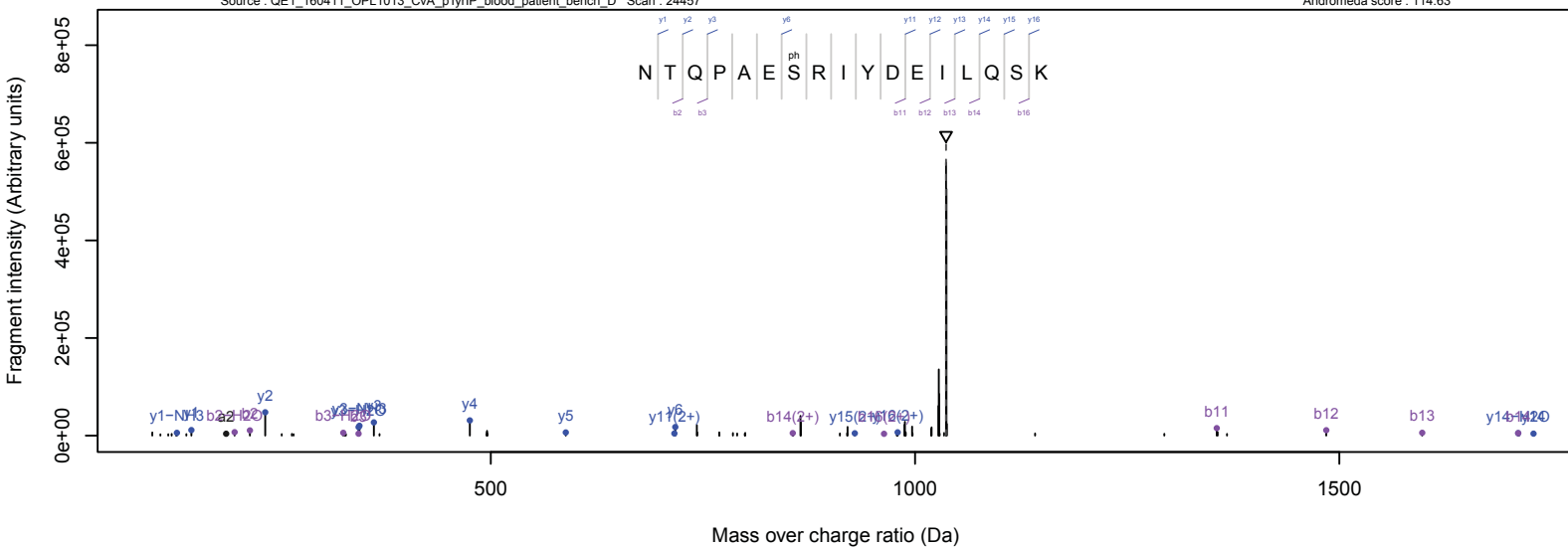
Andromeda score: 114.63



## NTQPAES(ph)RIYDEILQSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24457

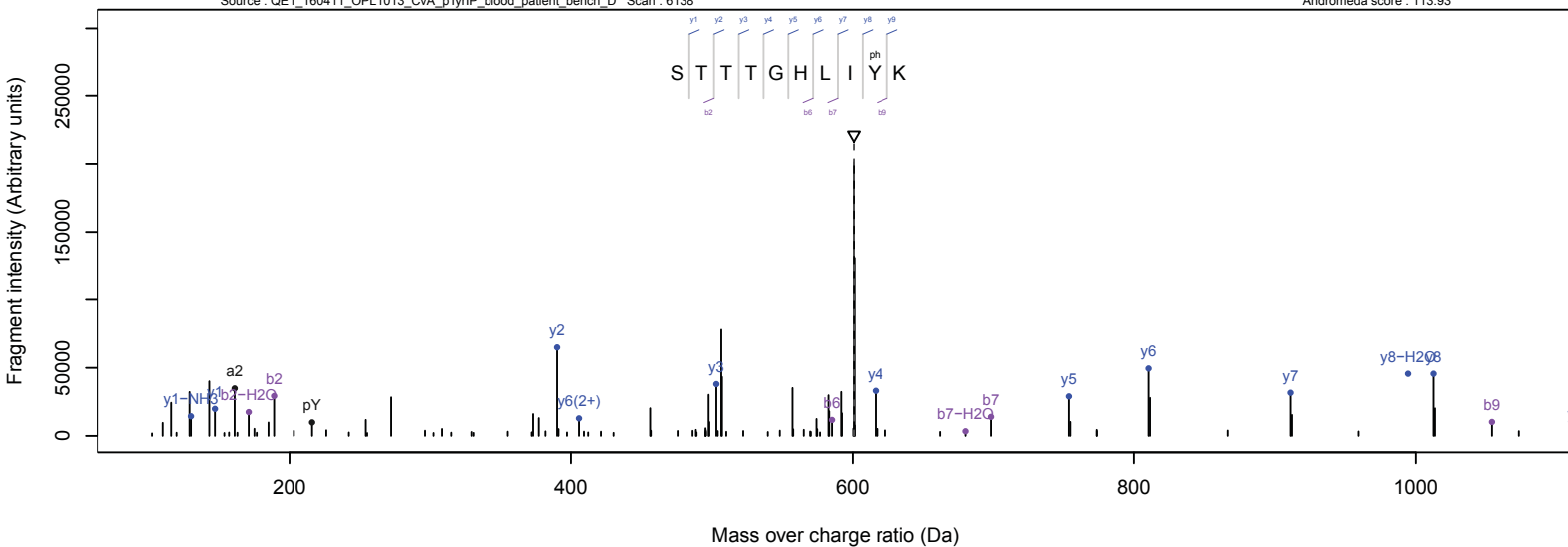
Andromeda score : 114.63



## STTTGHLIY(ph)K-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 6138

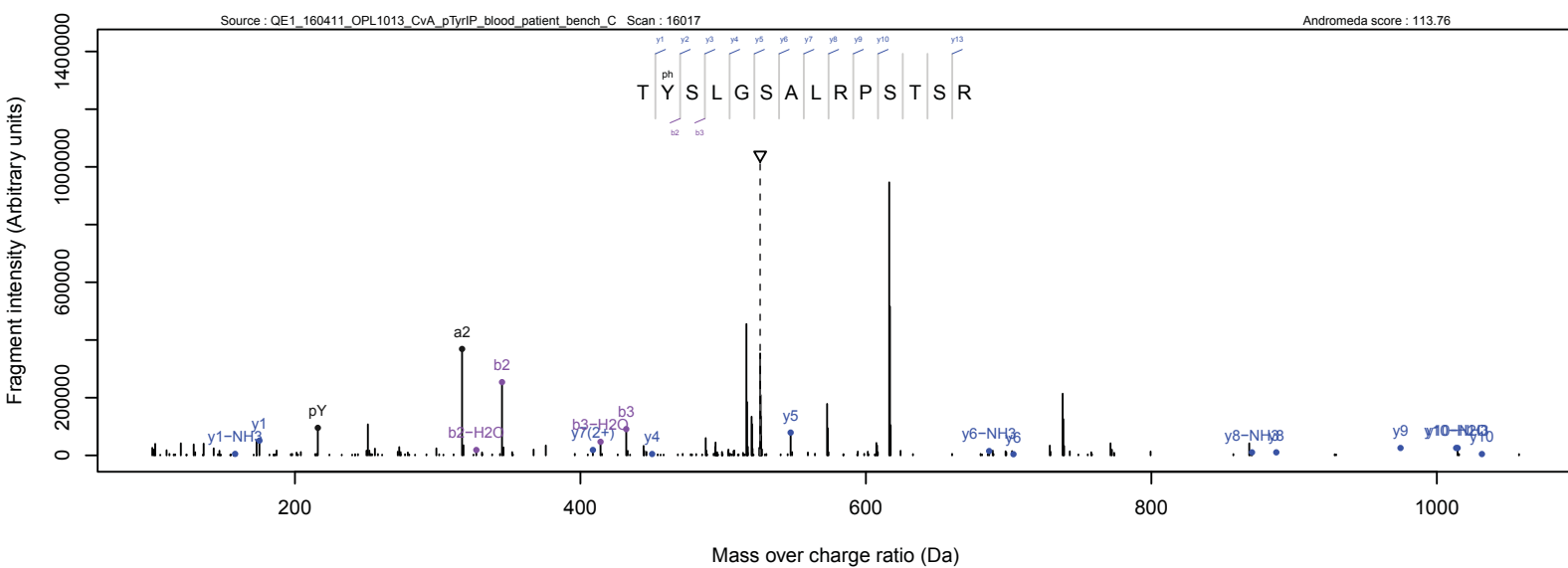
Andromeda score : 113.93



## TY(ph)SLGSALRPSTSR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16017

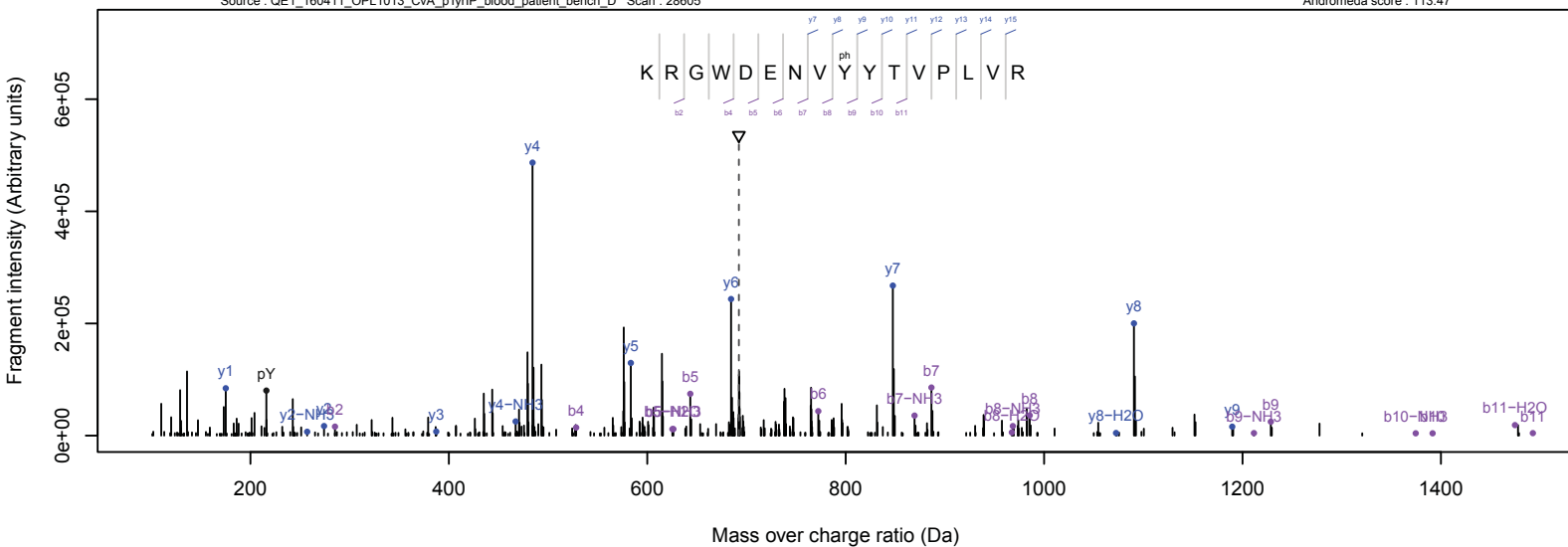
Andromeda score : 113.76



**KRGWDENVY<sup>ph</sup>YTVPLVR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28605

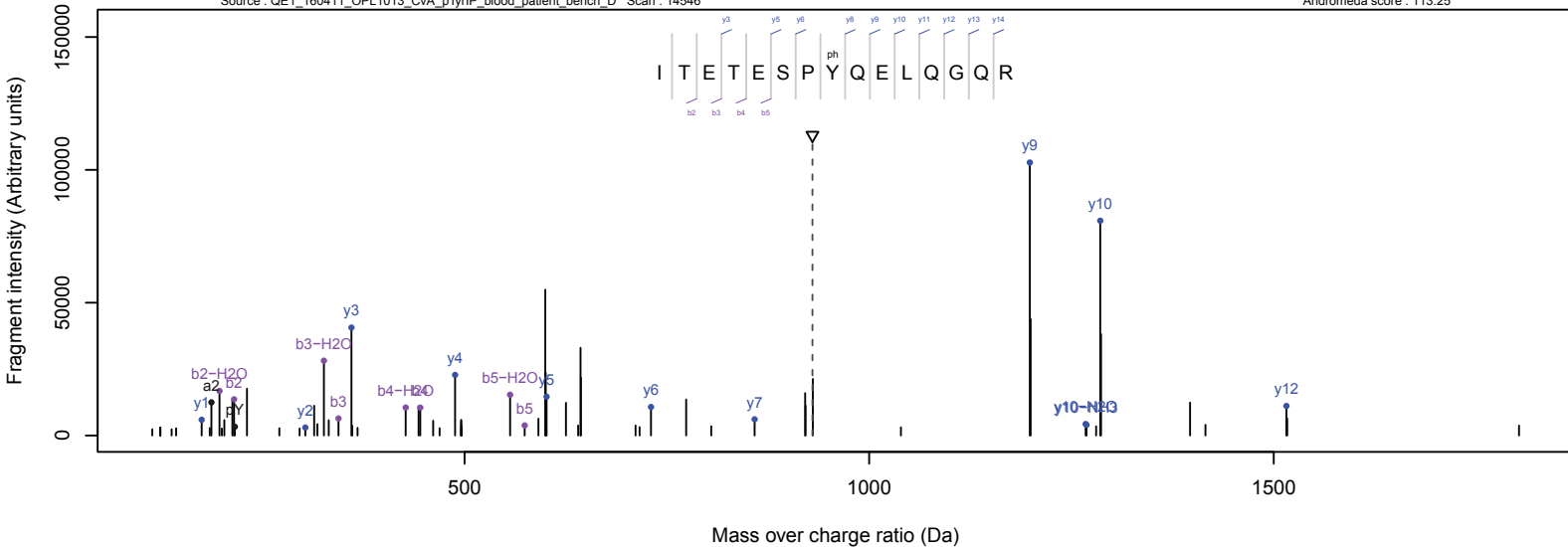
Andromeda score : 113.47



**ITETESPY<sup>ph</sup>QELQGQR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 14546

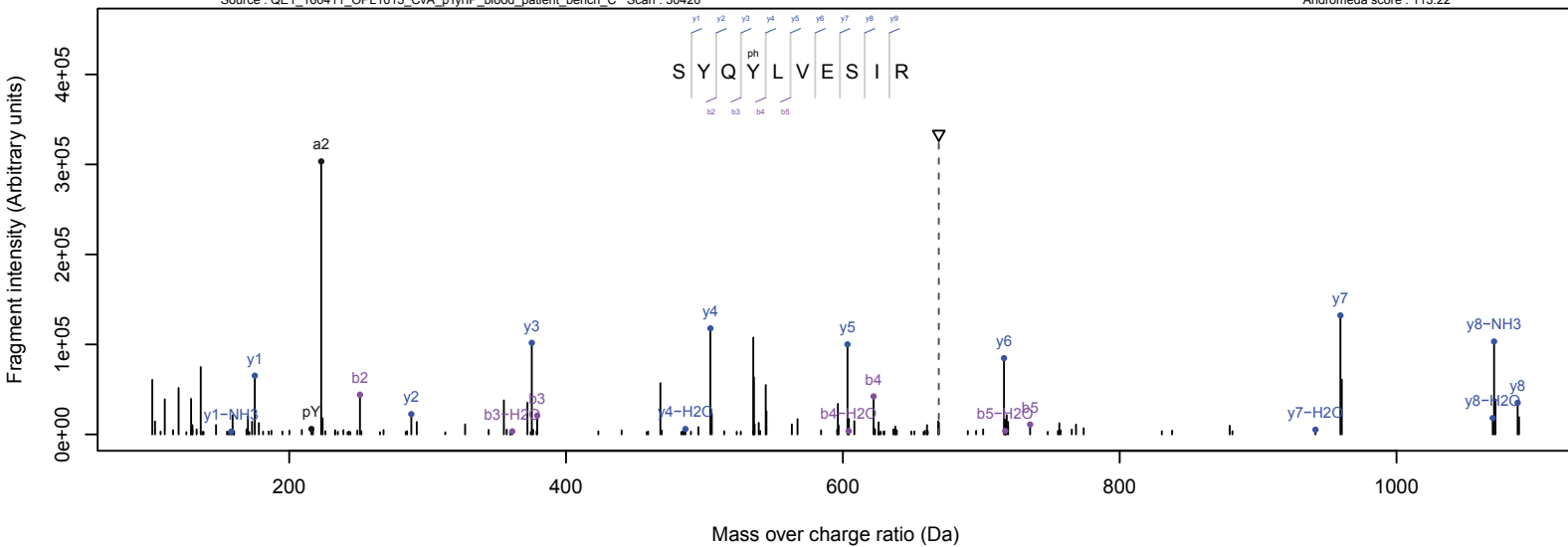
Andromeda score : 113.25



**SYQY<sup>ph</sup>LVESIR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 30420

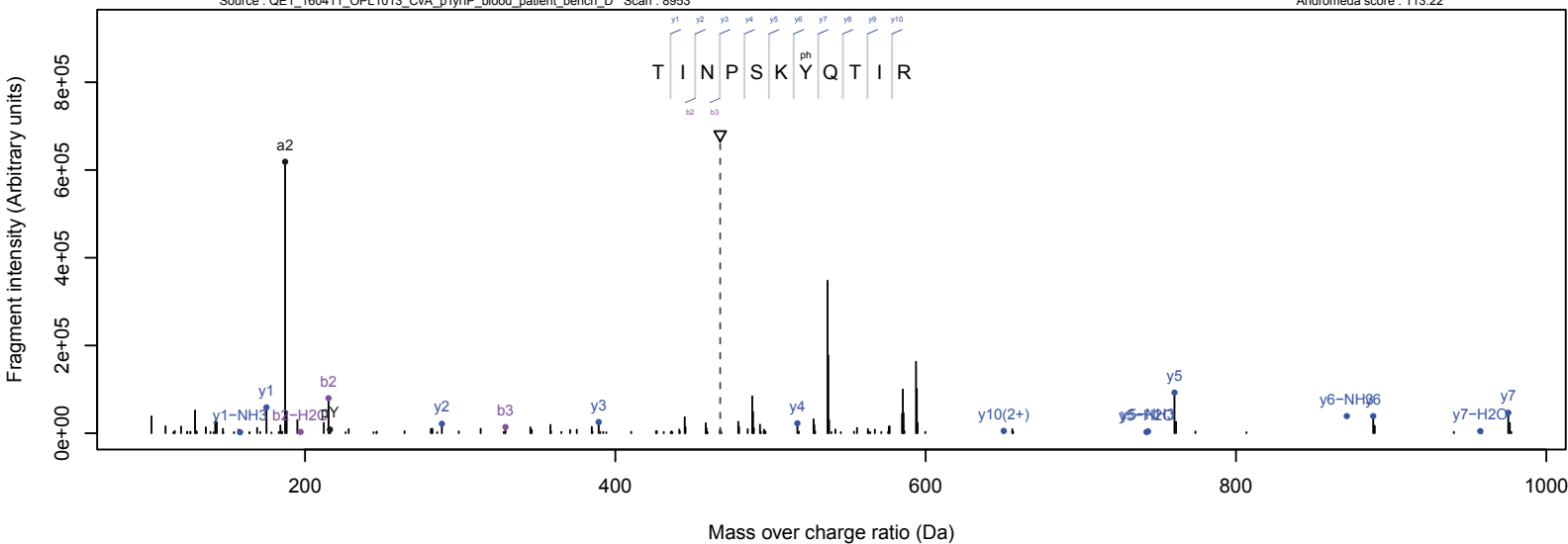
Andromeda score : 113.22



TINPSKY(ph)QTIR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 8953

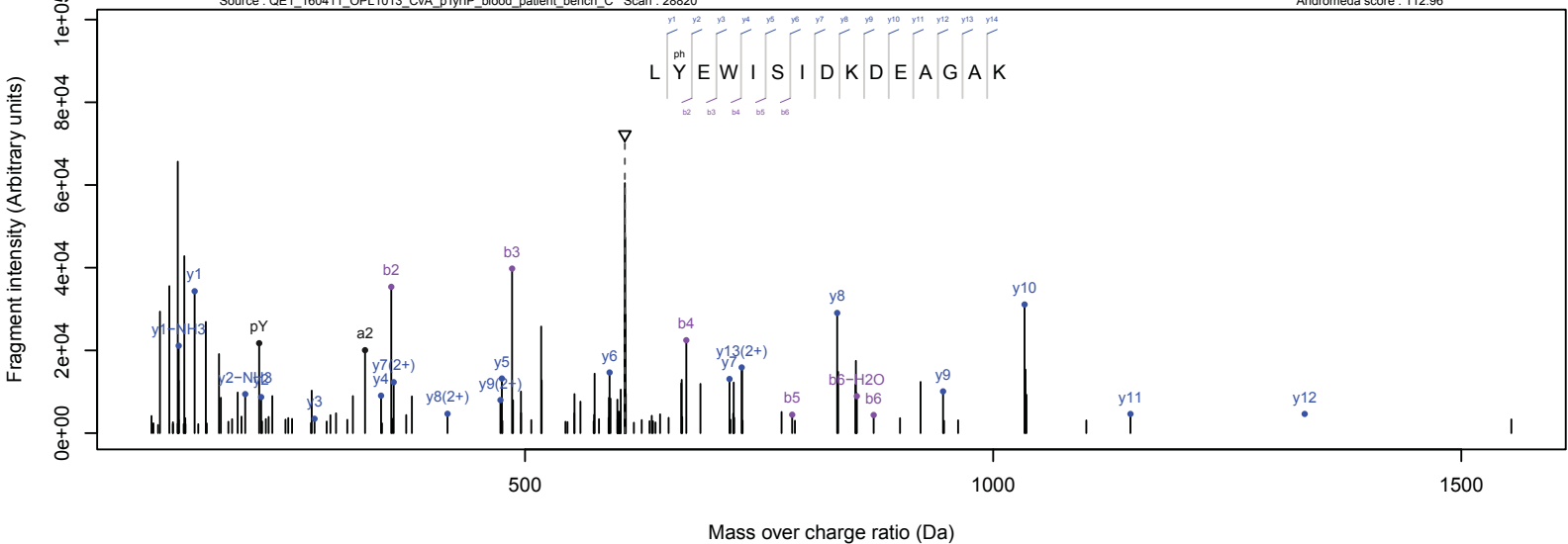
Andromeda score: 113.22



LY(ph)EWISIDKDEAGAK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 28820

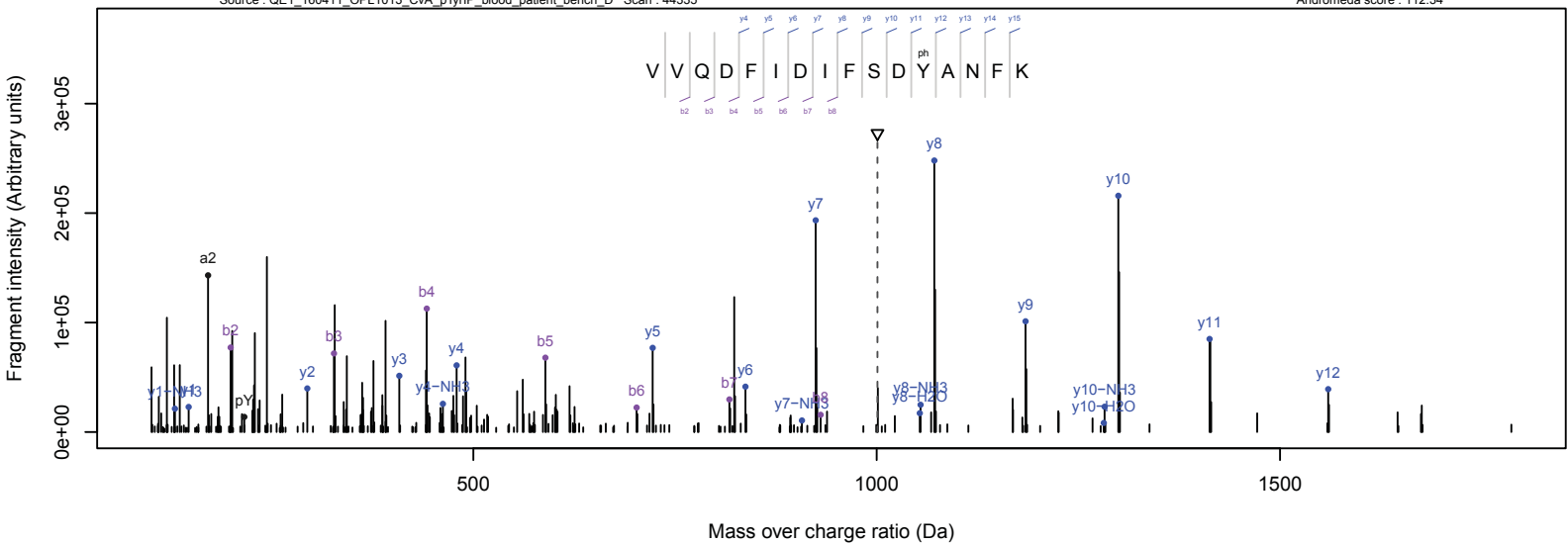
Andromeda score: 112.96



VVQDFIDIFSDYANF K-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 44335

Andromeda score: 112.34

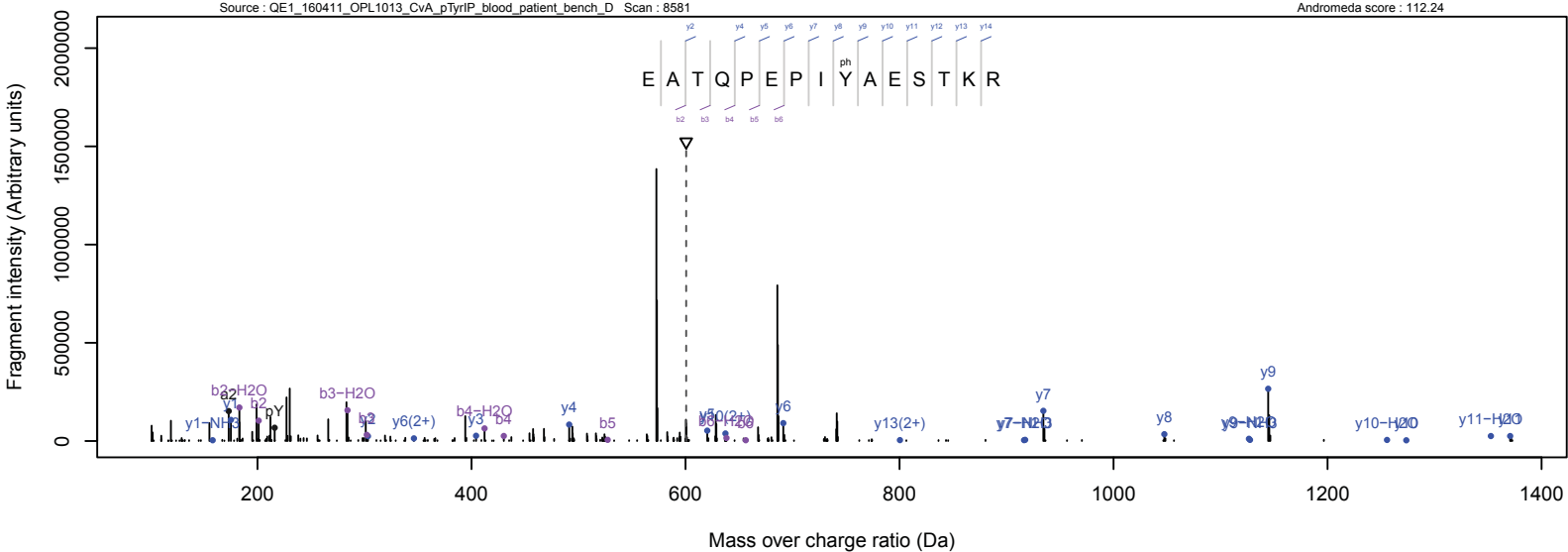




## EATQPEPIY(ph)AESTKR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 8581

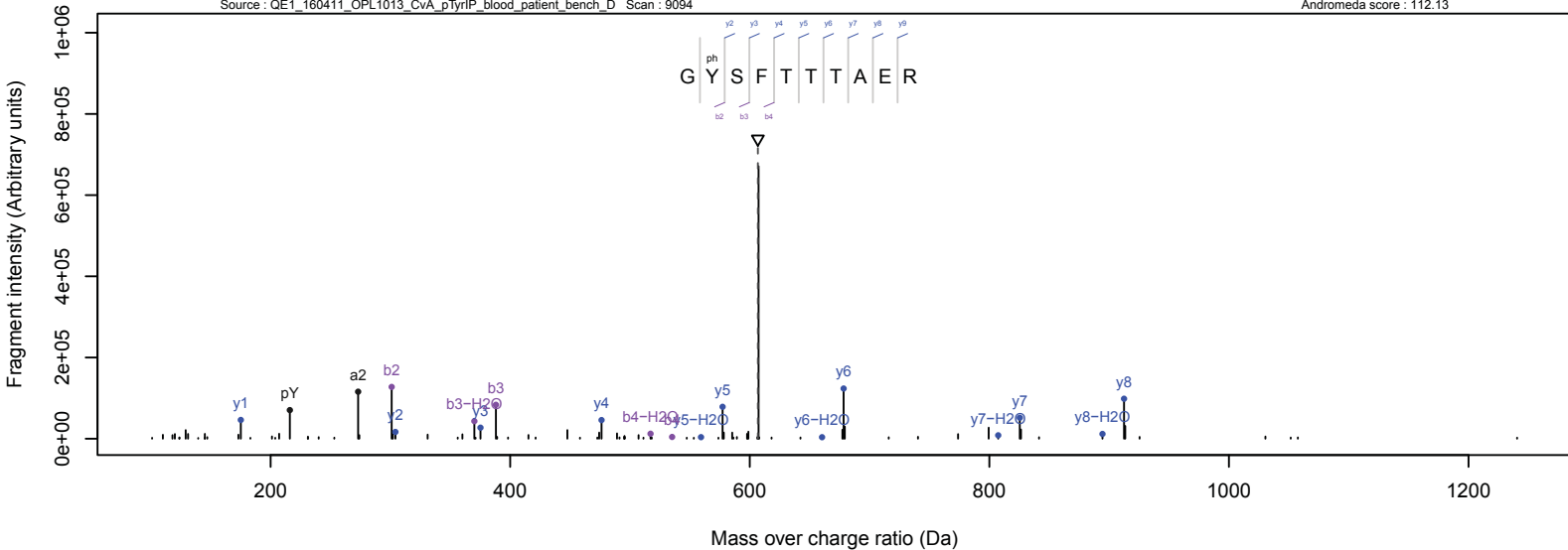
Andromeda score: 112.24



## GY(ph)SFTTTAER-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 9094

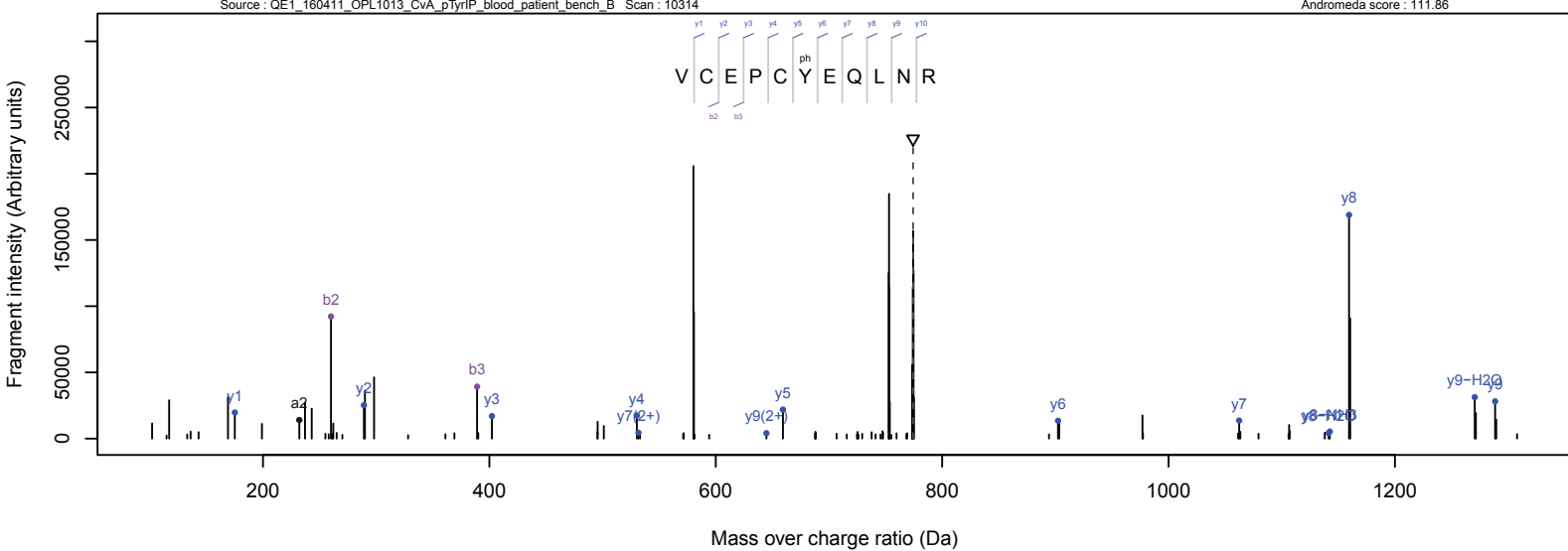
Andromeda score: 112.13



## VCEPCY(ph)EQLNR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 10314

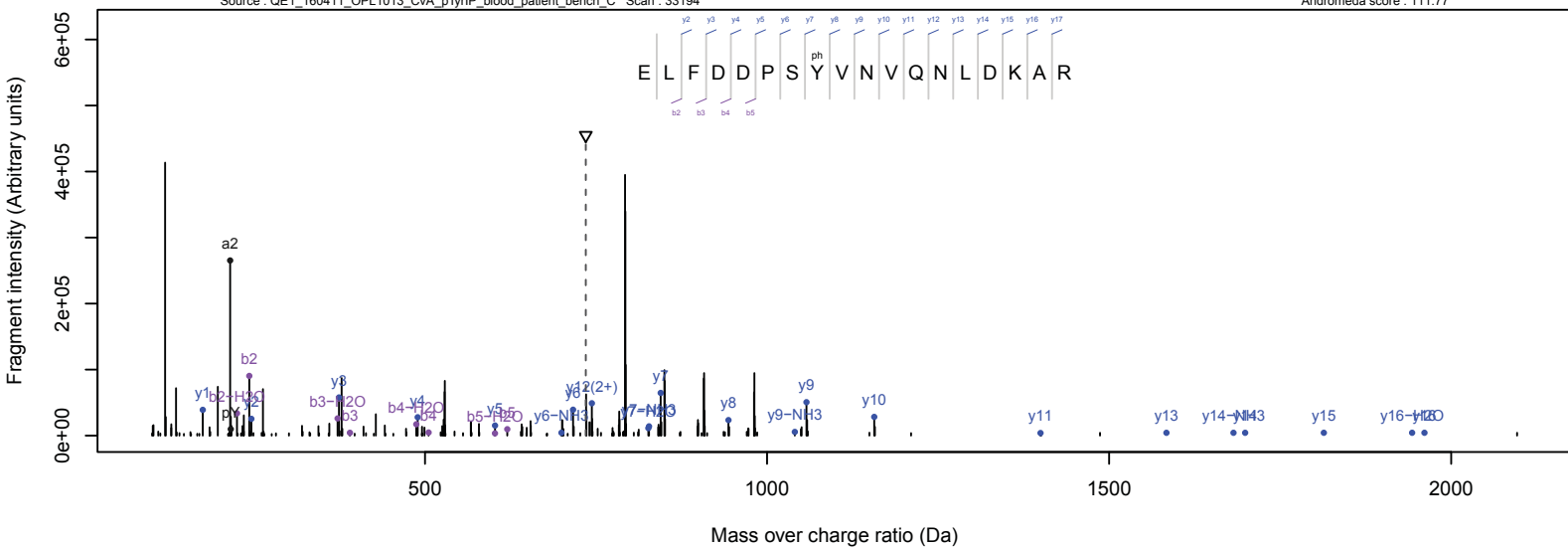
Andromeda score: 111.86



ELFDDPSY(ph)VNVQLDKAR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 33194

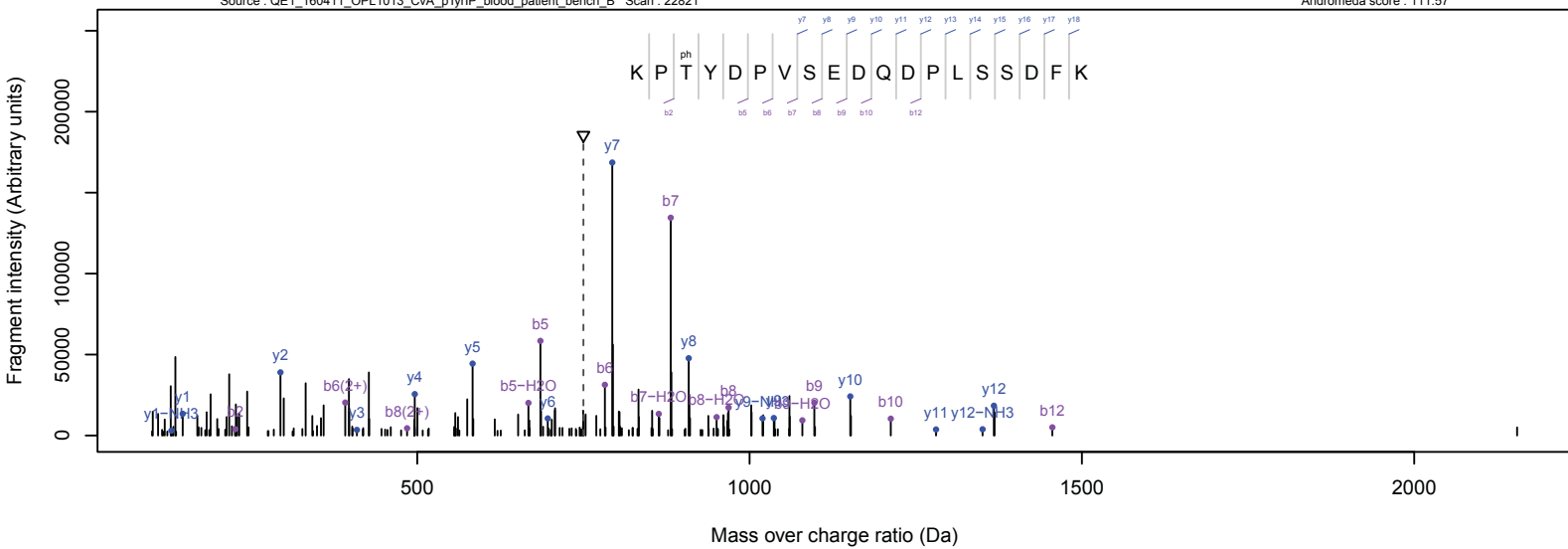
Andromeda score : 111.77



KPT(ph)YDPVSEDQDPLSSDFK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 22821

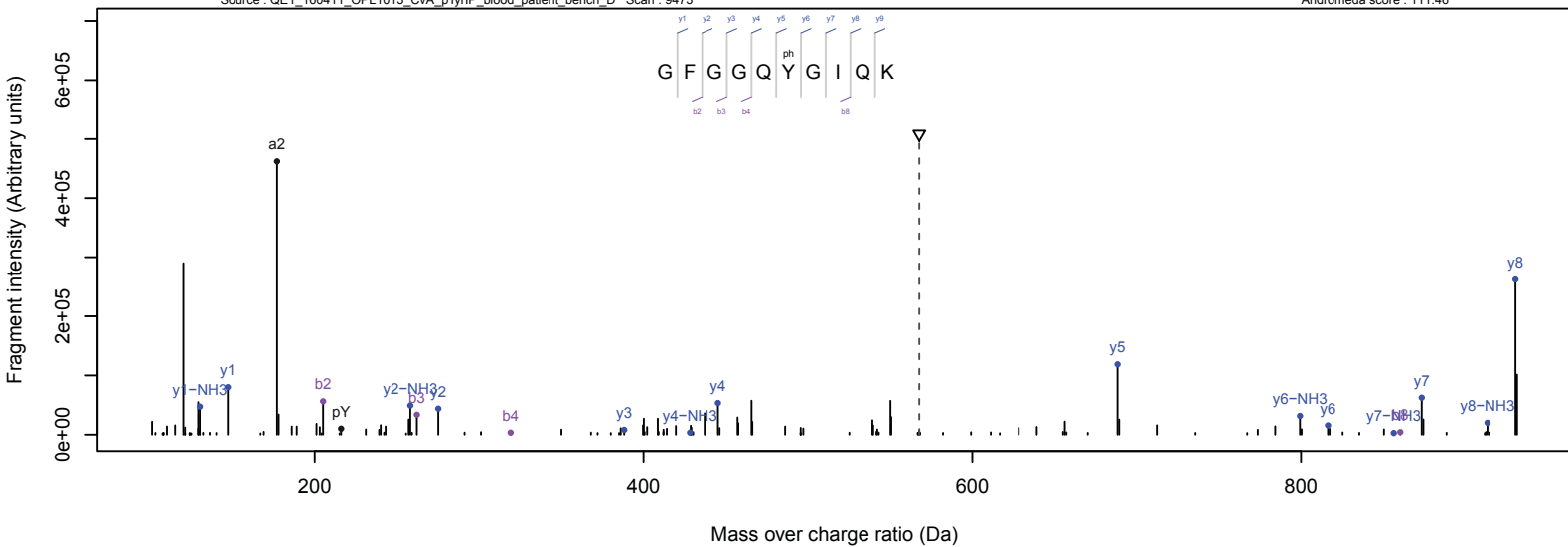
Andromeda score : 111.57



GFGGQY(ph)GIQK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 9473

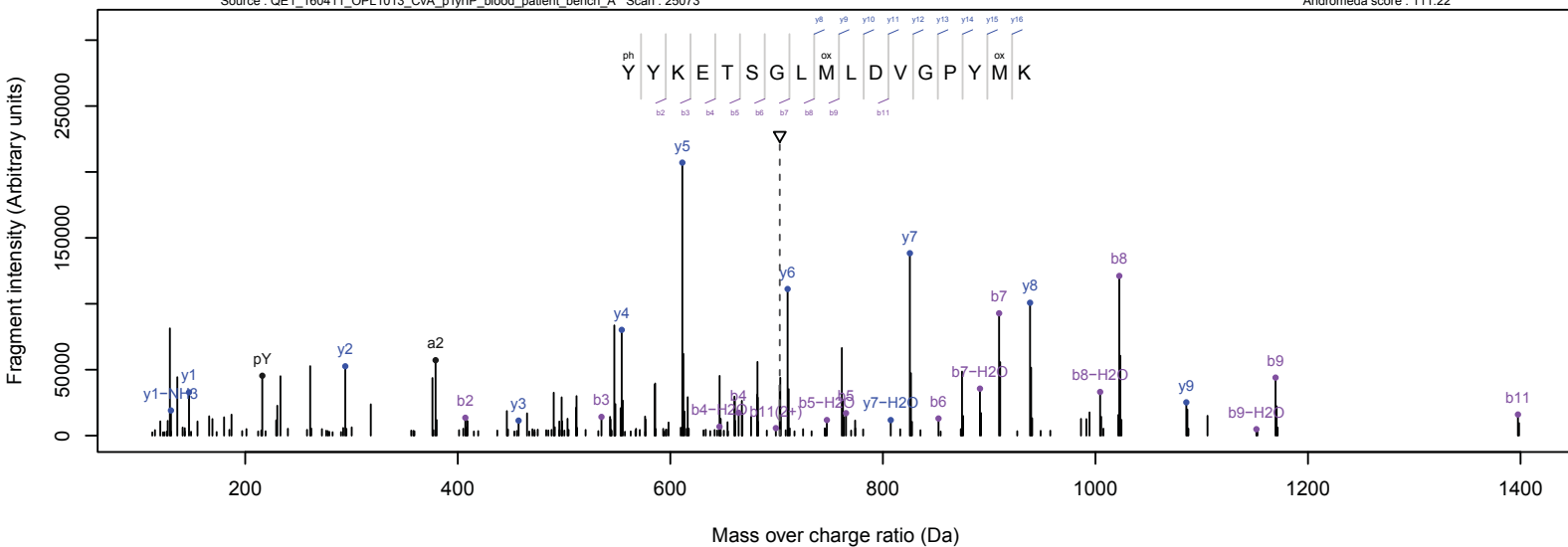
Andromeda score : 111.46



## Y(ph)YKETSGLM(ox)LDVGPYM(ox)K-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 25073

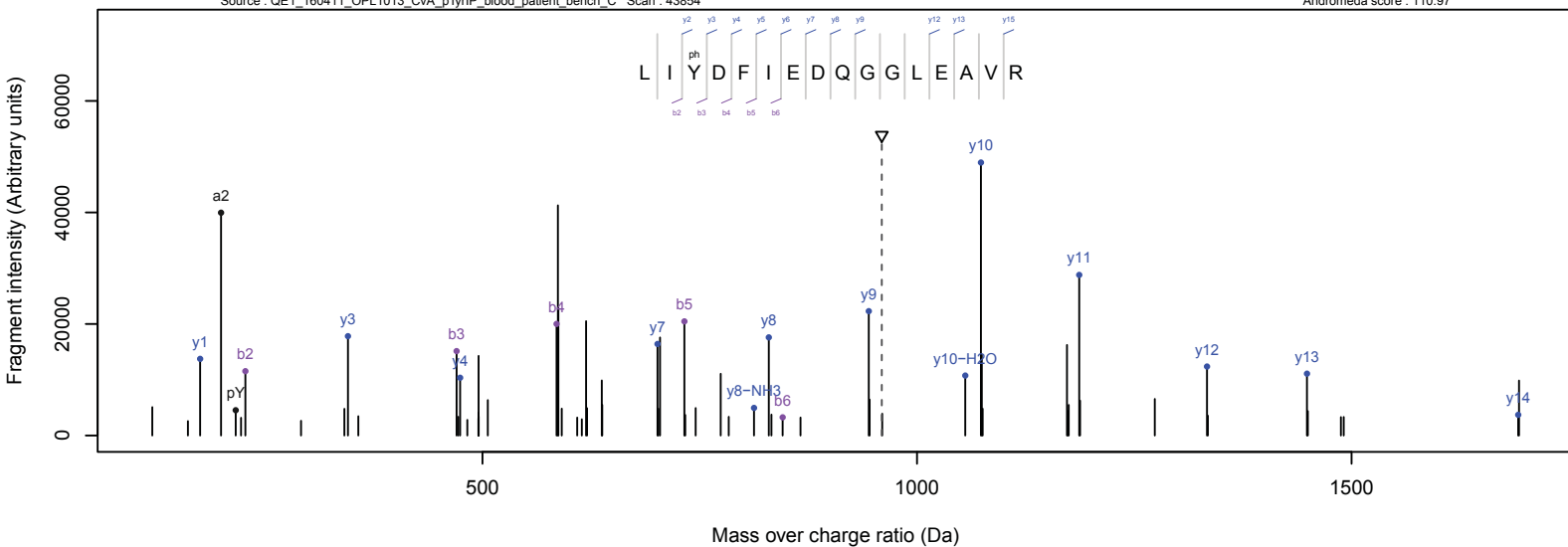
Andromeda score : 111.22



## LIY(ph)DFIEDQGGLEAVR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 43854

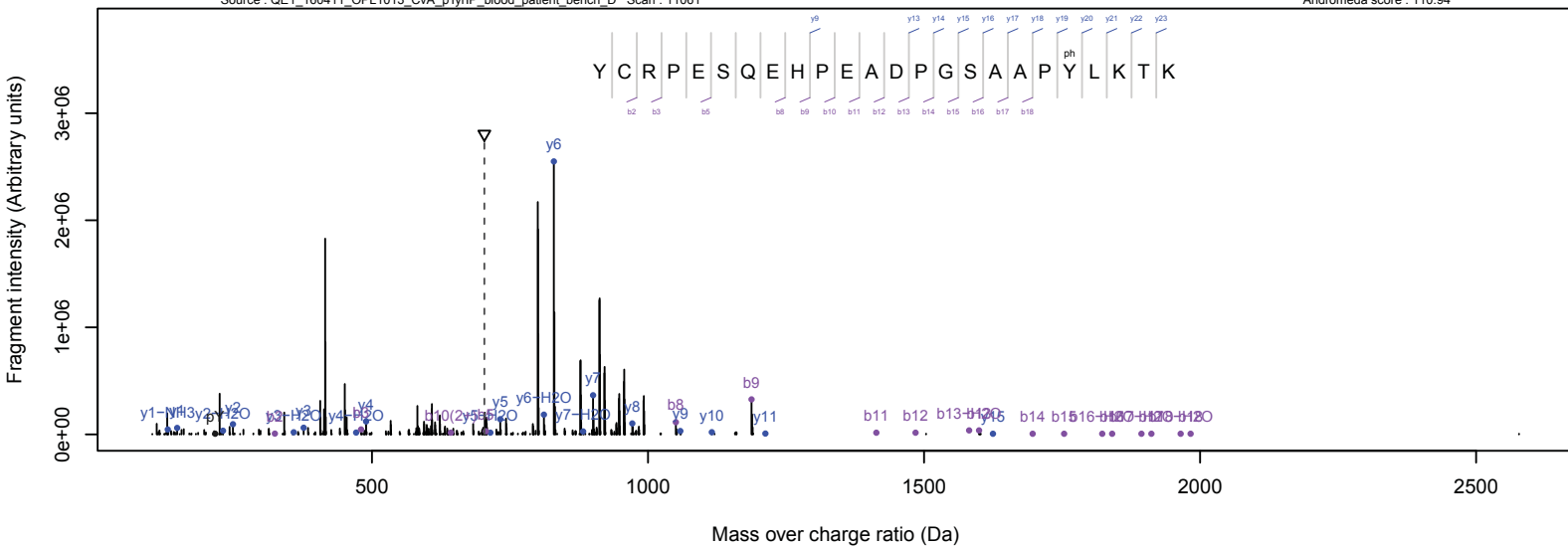
Andromeda score : 110.97



## YCRPESQEHPEADPGSAAPY(ph)LKTK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11061

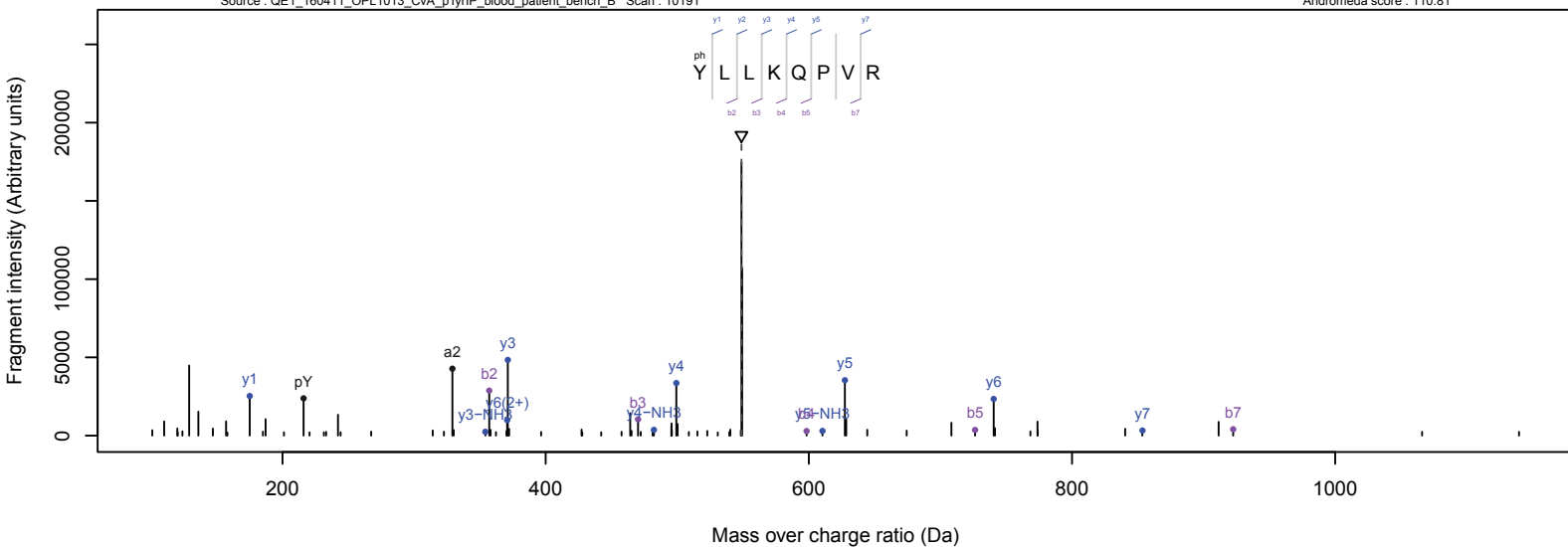
Andromeda score : 110.94



Y(ph)LLKQPVR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 10191

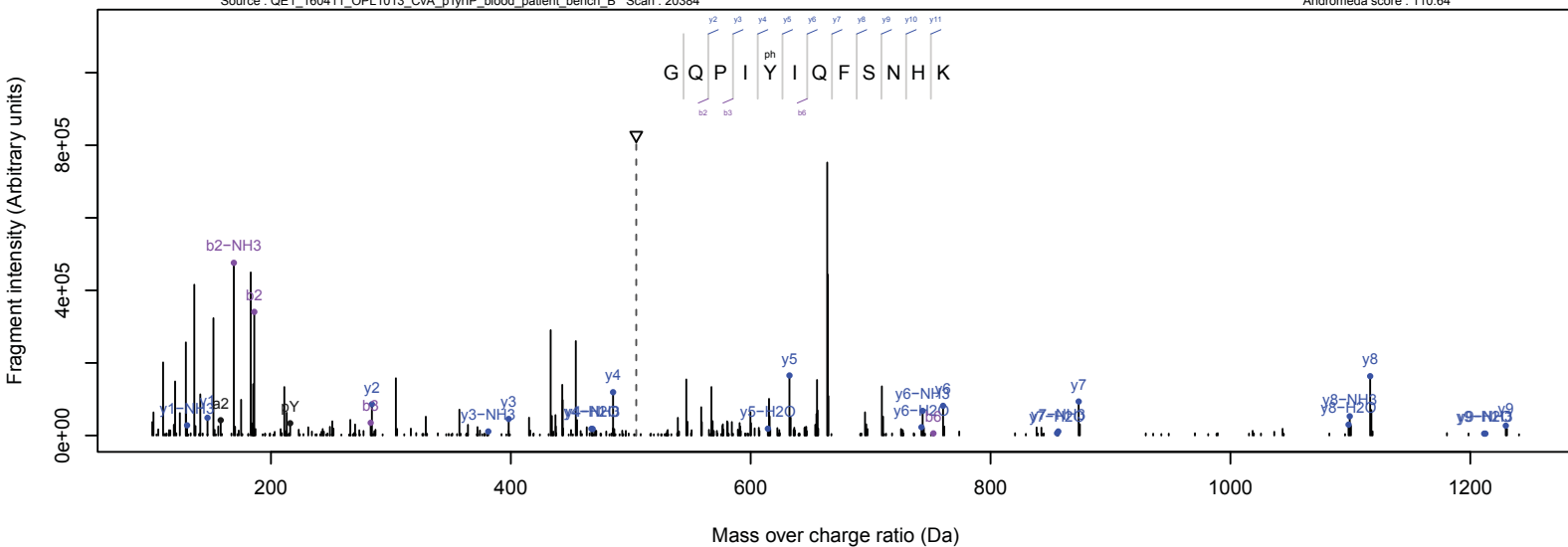
Andromeda score : 110.81



GQPIYIQFSNHK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 20384

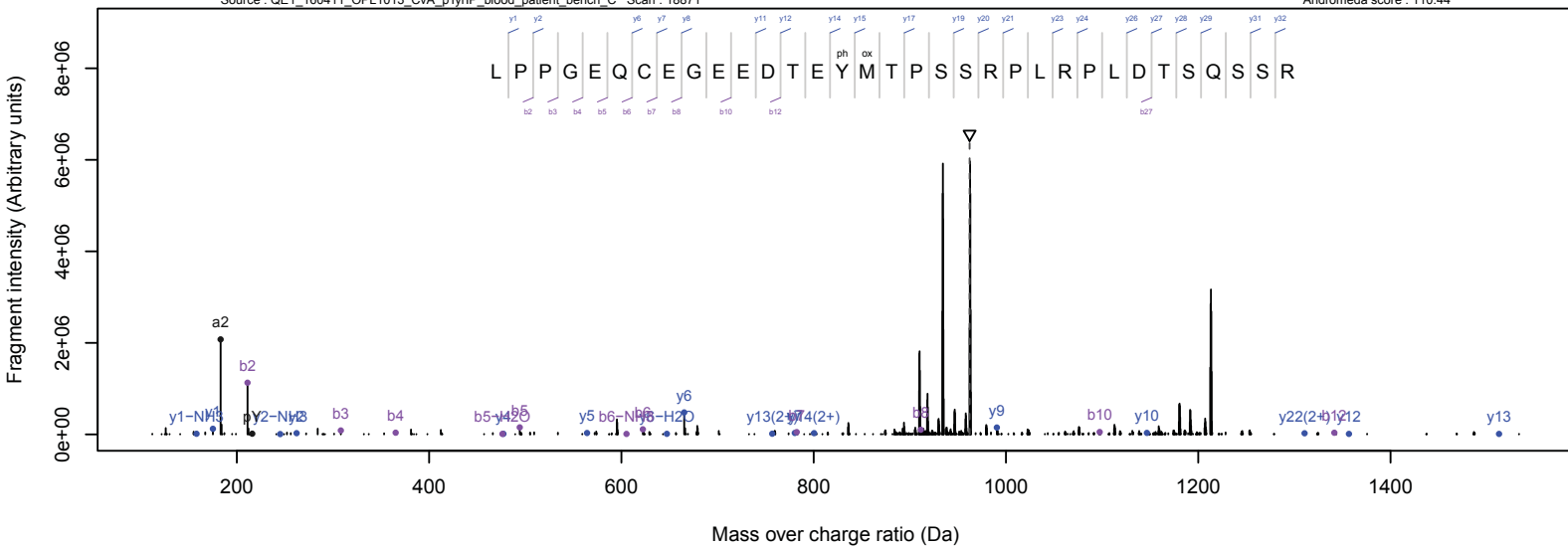
Andromeda score : 110.64



LPPGEQCEGEDTEY(ph)M(ox)TPSSRPLRPLDTSQSSR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18871

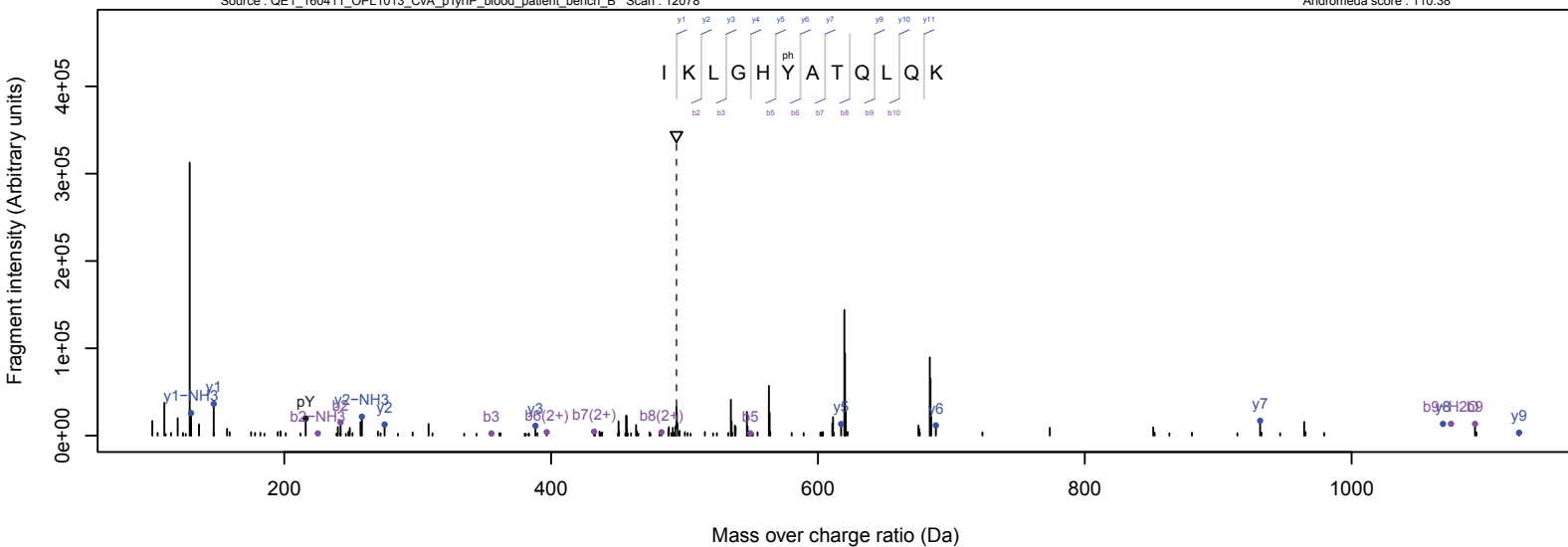
Andromeda score : 110.44



**IKLGHY(ph)ATQLQK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 12078

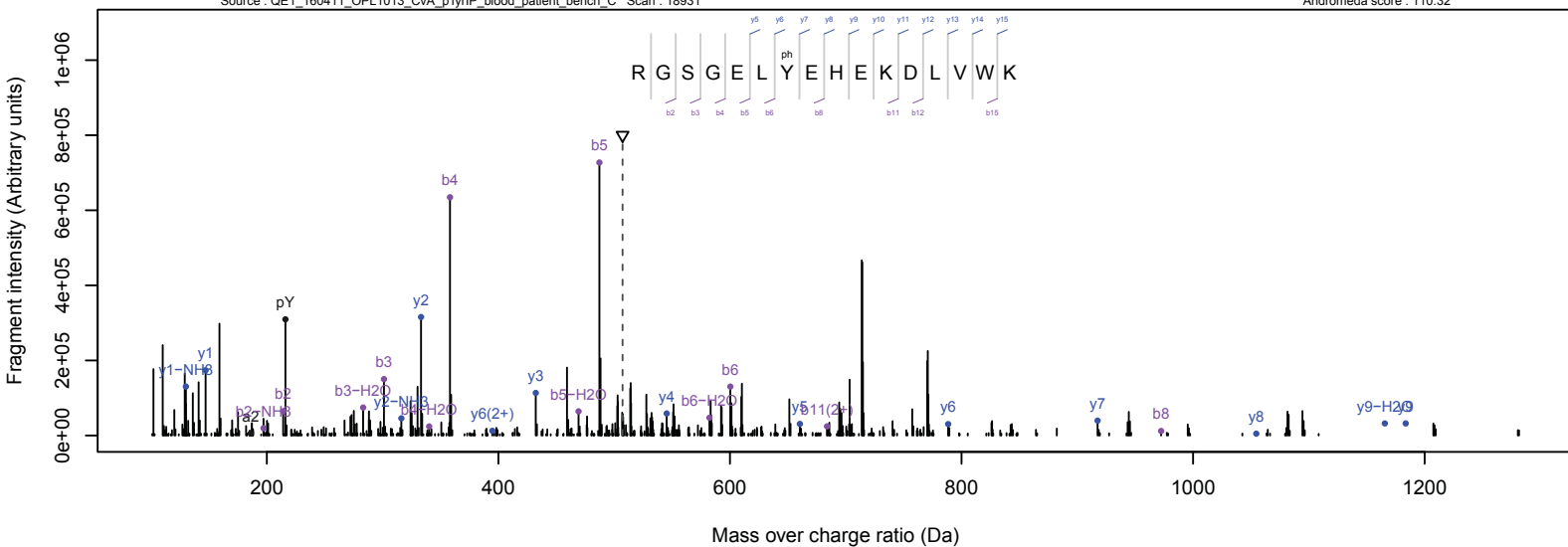
Andromeda score : 110.38



**RGSGELY(ph)EHEKDLVWK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18931

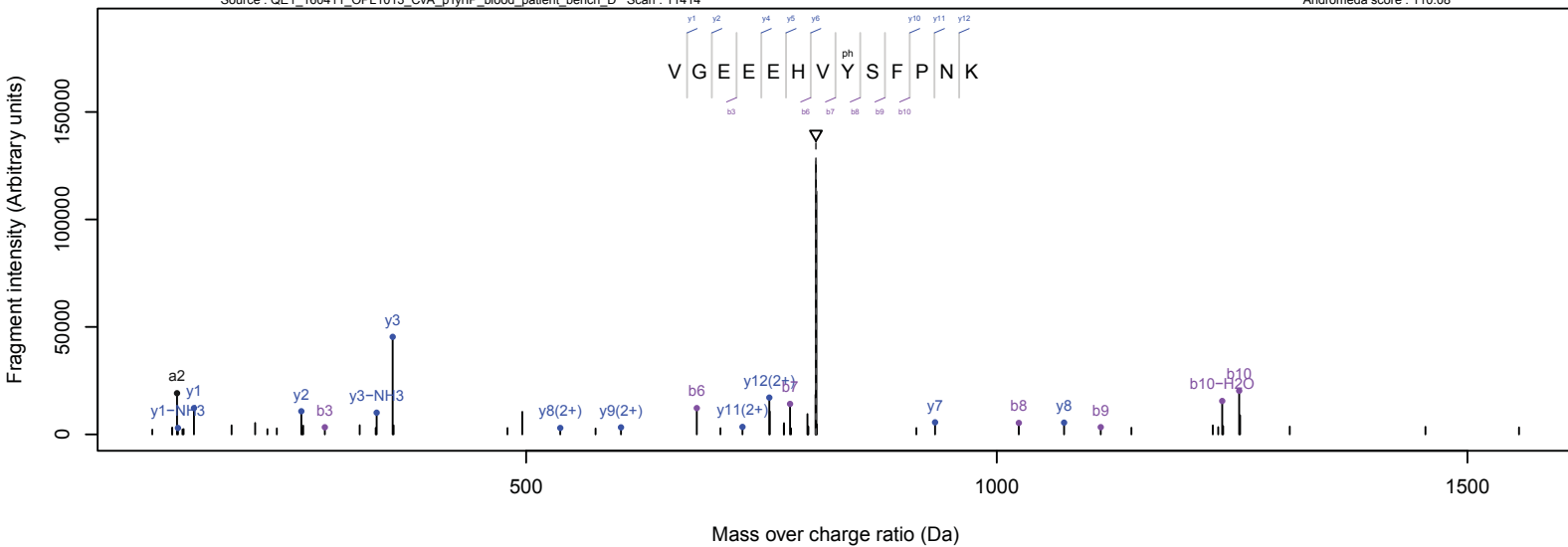
Andromeda score : 110.32



**VGEEEHVY(ph)SFPNK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11414

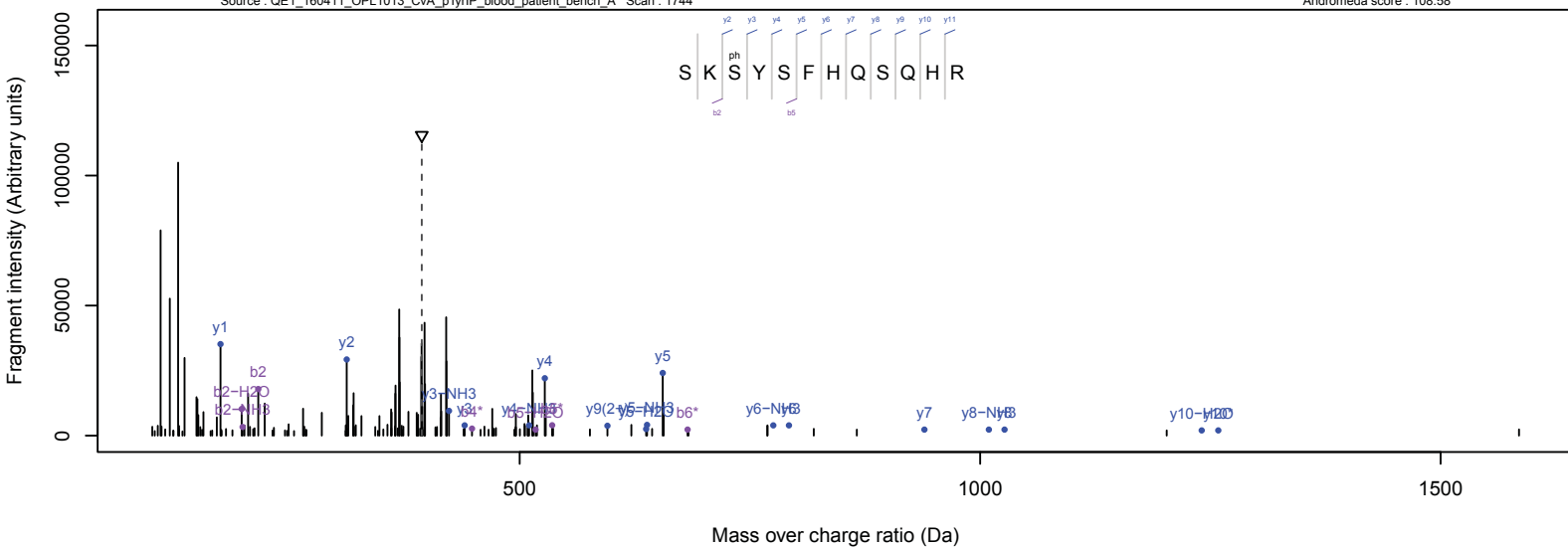
Andromeda score : 110.08



## SKS(ph)YSFHQSQR-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 1744

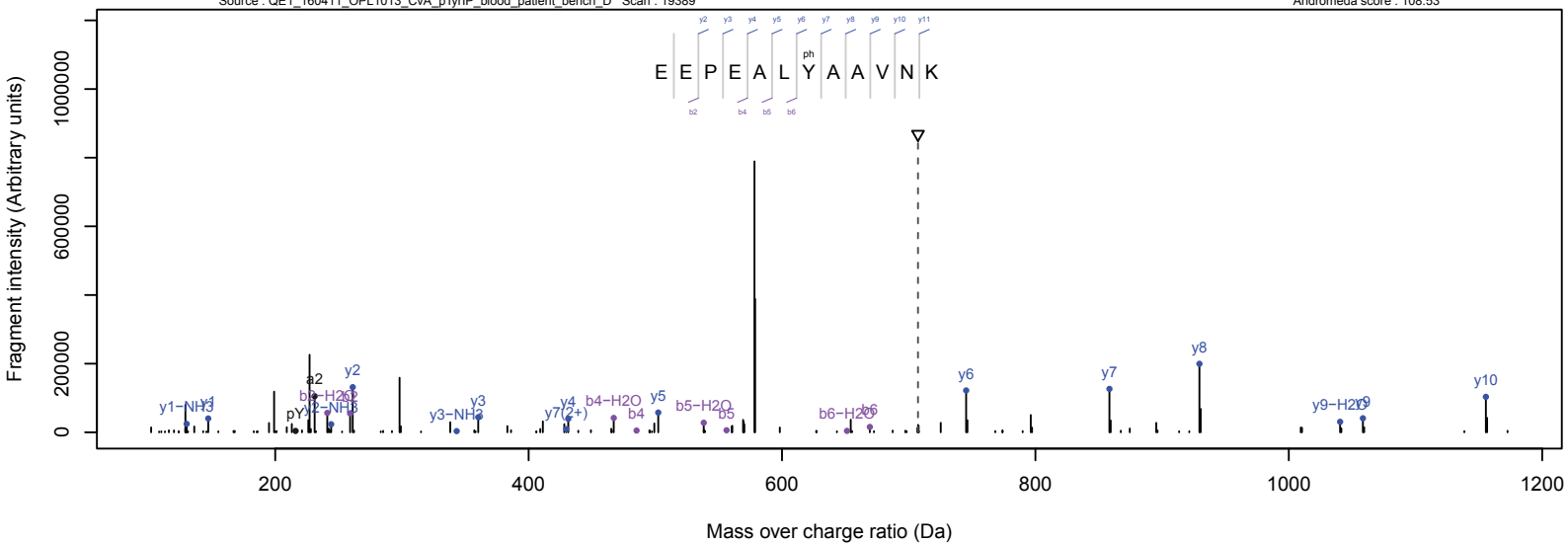
Andromeda score: 108.58



## EEPEALY(ph)AAVNK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 19389

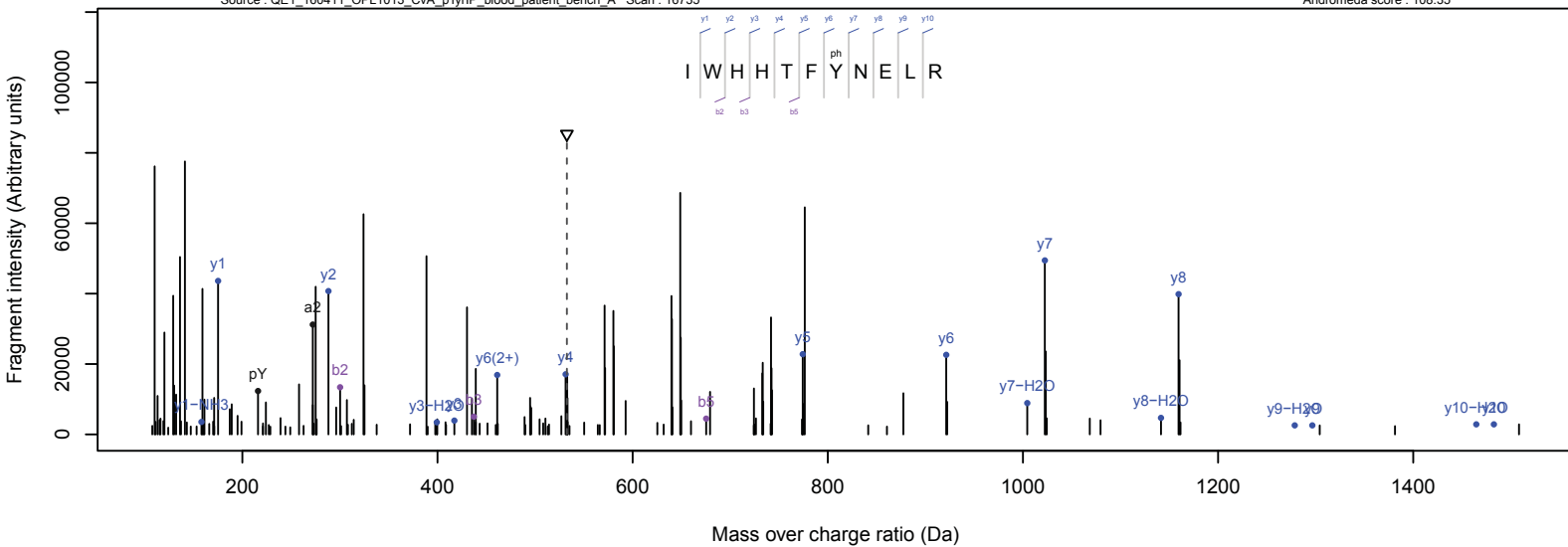
Andromeda score: 108.53



## IWHHTFY(ph)NELR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 16735

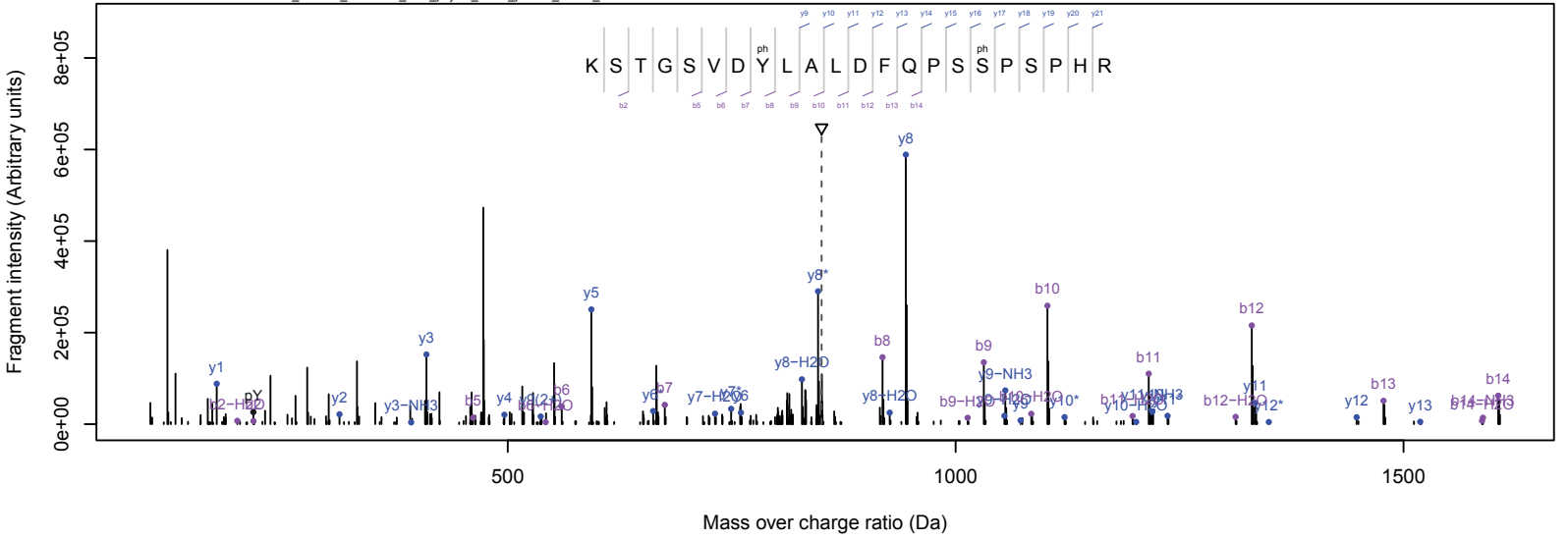
Andromeda score: 108.35



**KSTGSDY(ph)LALDFQPSS(ph)PSPHR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 29526

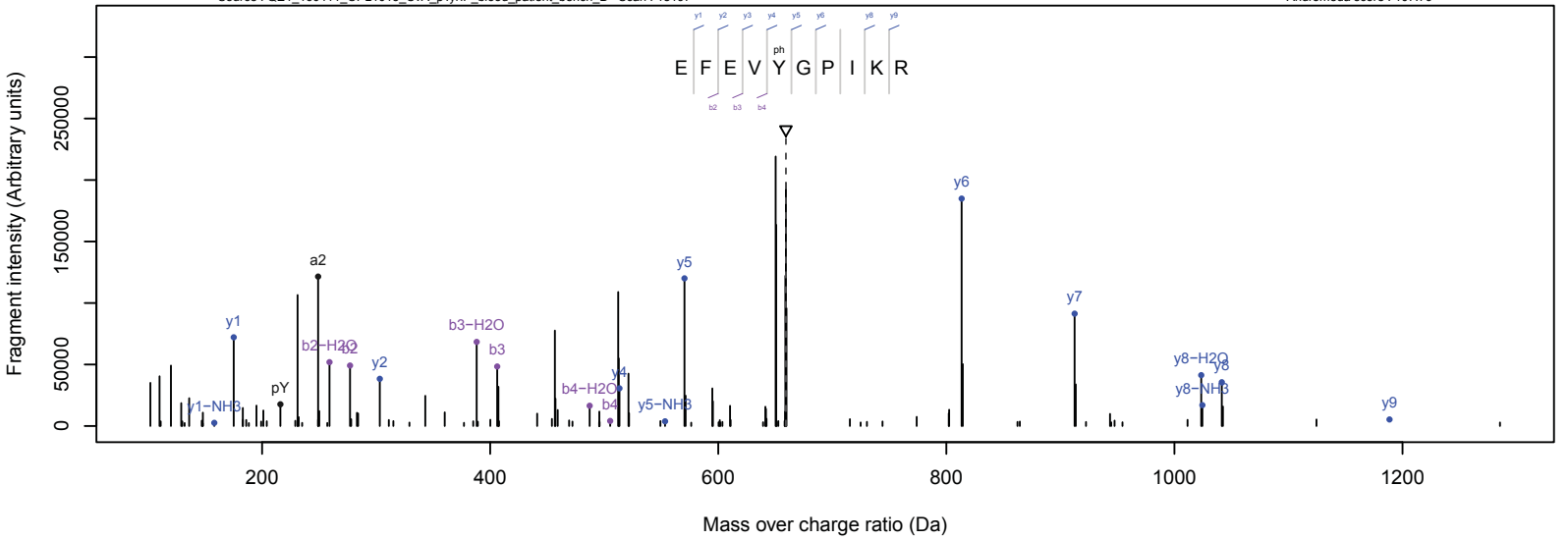
Andromeda score : 108.23



**EFEVY(ph)GPIKR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 18197

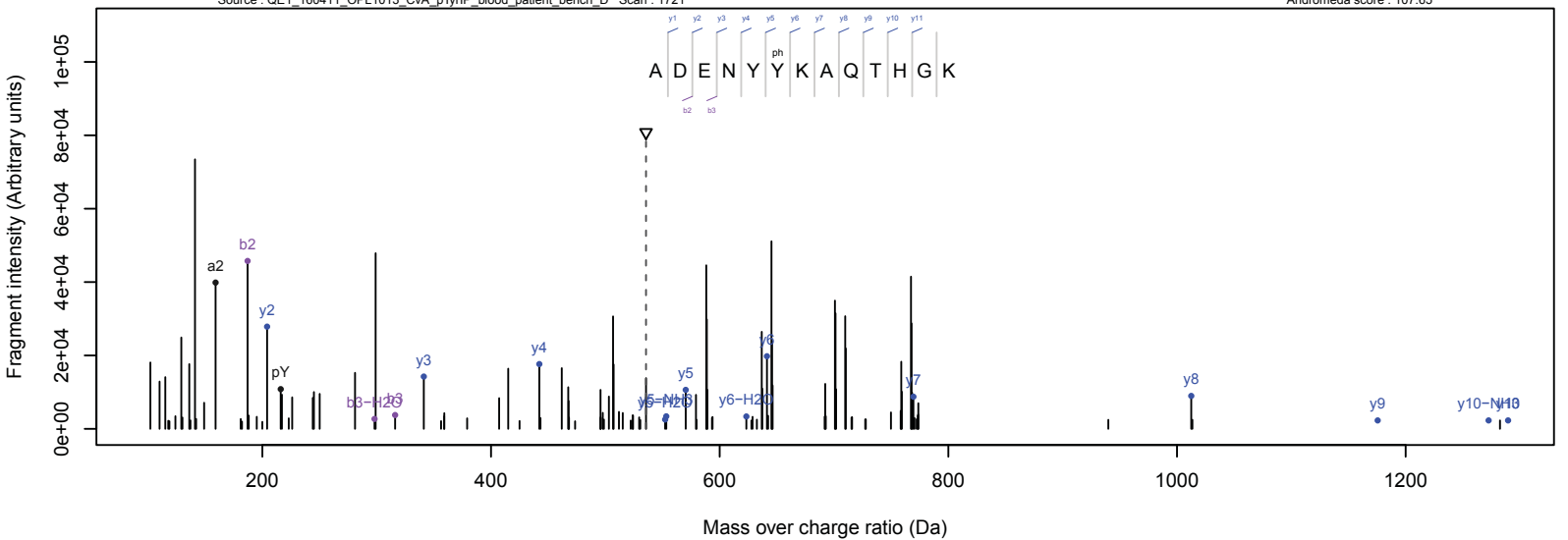
Andromeda score : 107.79



**ADENY(ph)KAQTHGK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 1721

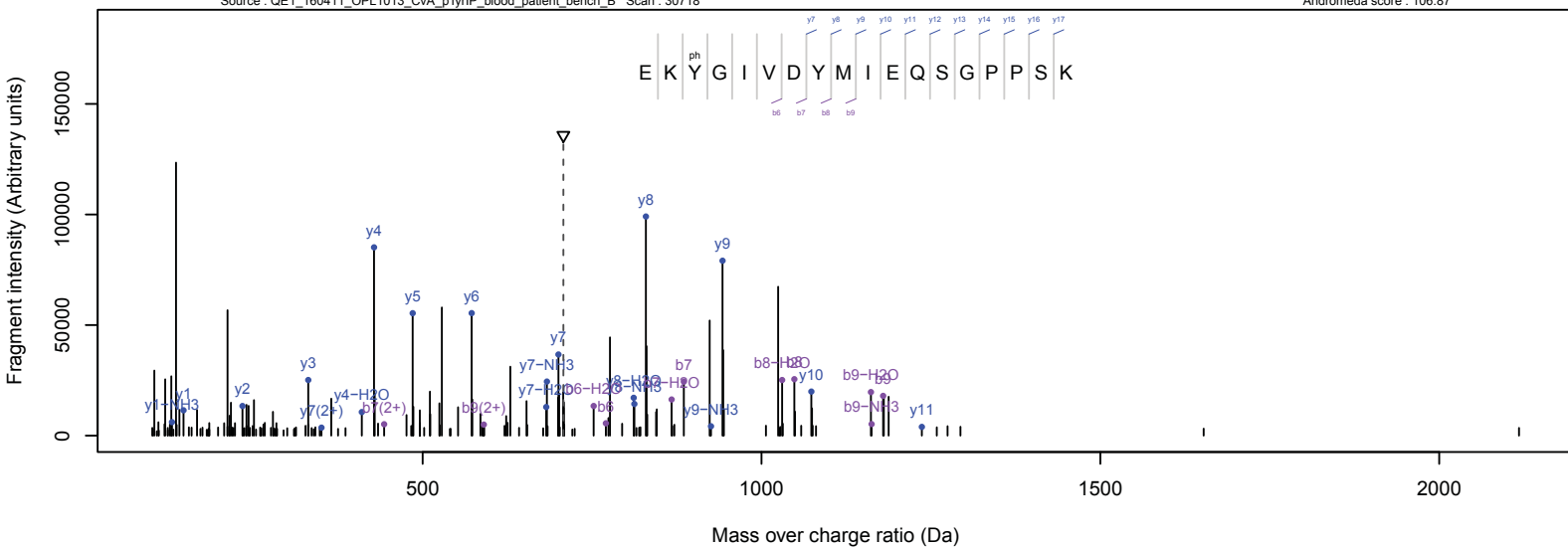
Andromeda score : 107.65



**EKY(ph)GIVDYMIEQSGPPSK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 30718

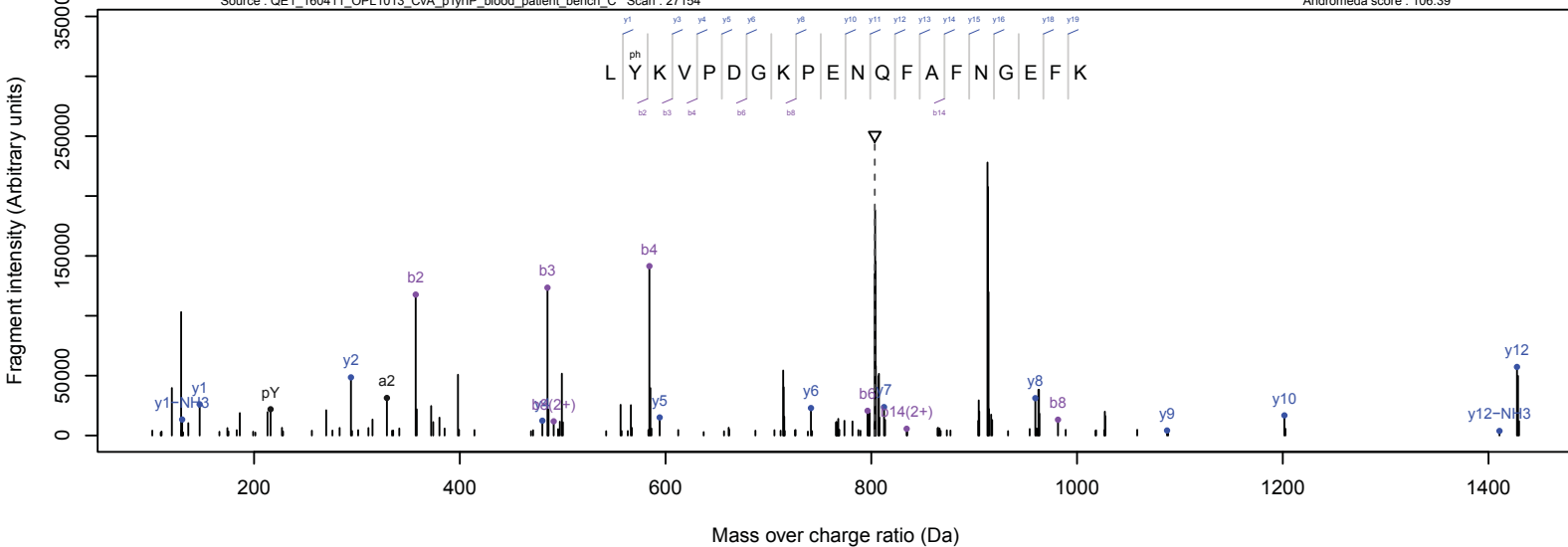
Andromeda score : 106.87



**LY(ph)KVPDGPENQFAFNGEFK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 27154

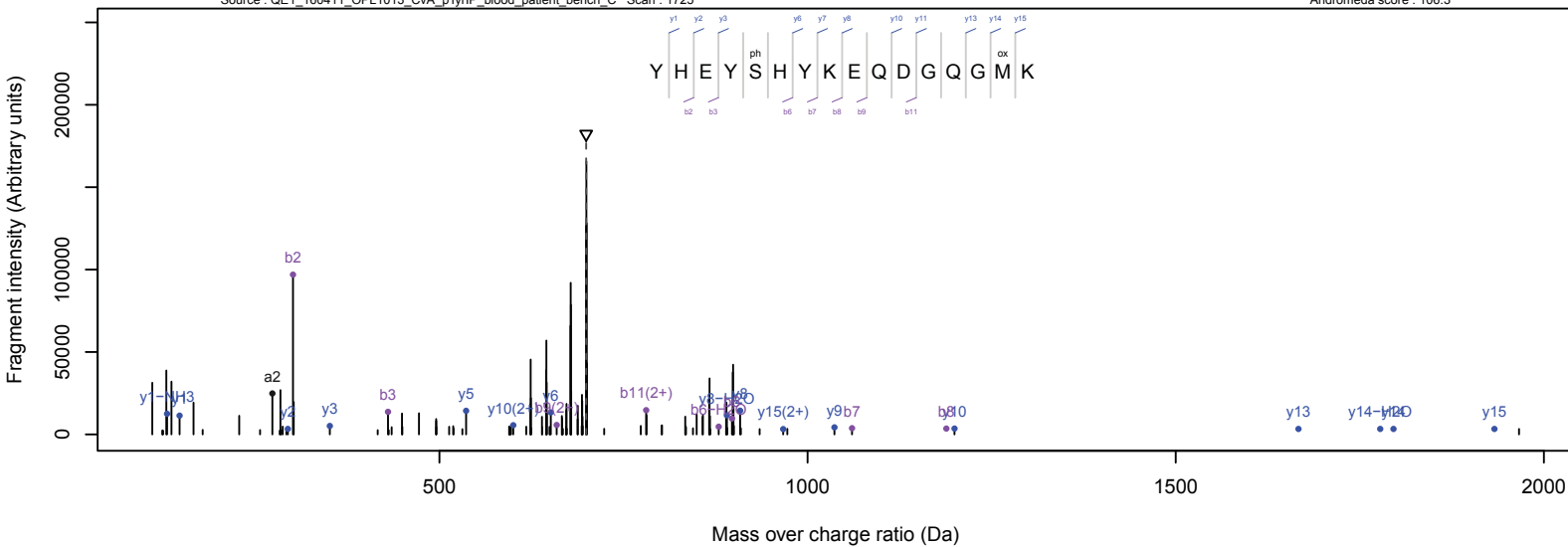
Andromeda score : 106.39



**YHEYS(ph)HYKEQDGGM(ox)K-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 1725

Andromeda score : 106.3

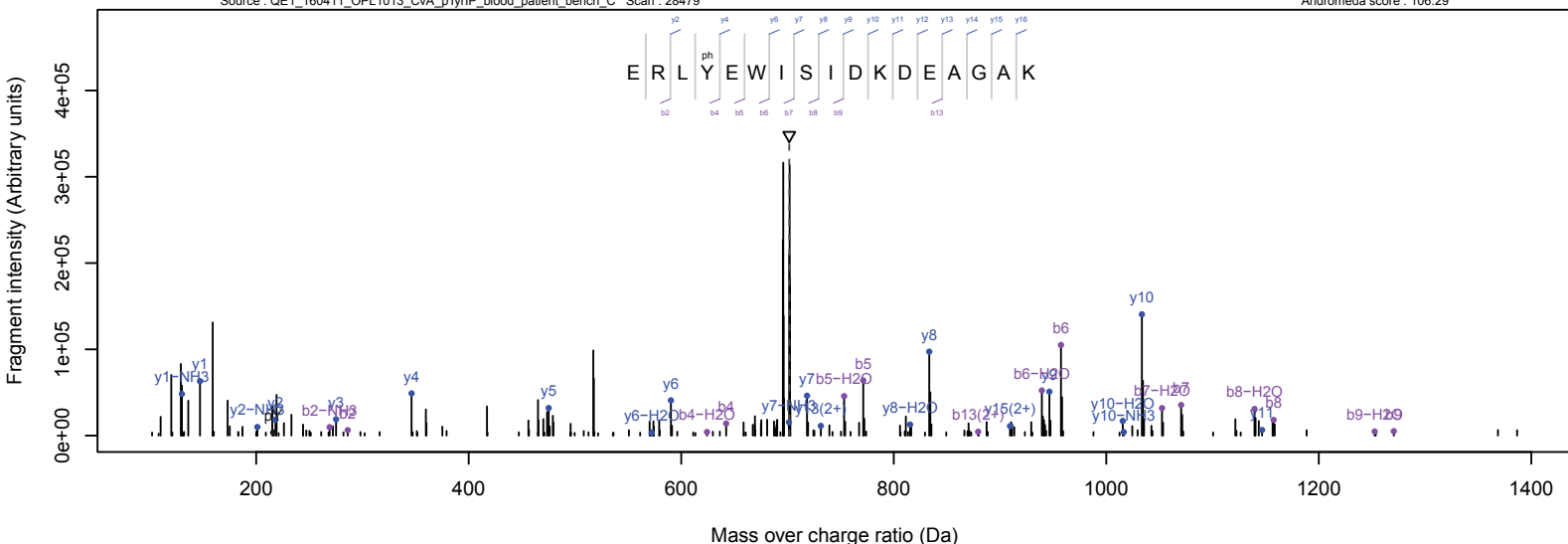




ERLY(ph)EWISIDKDEAGK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28479

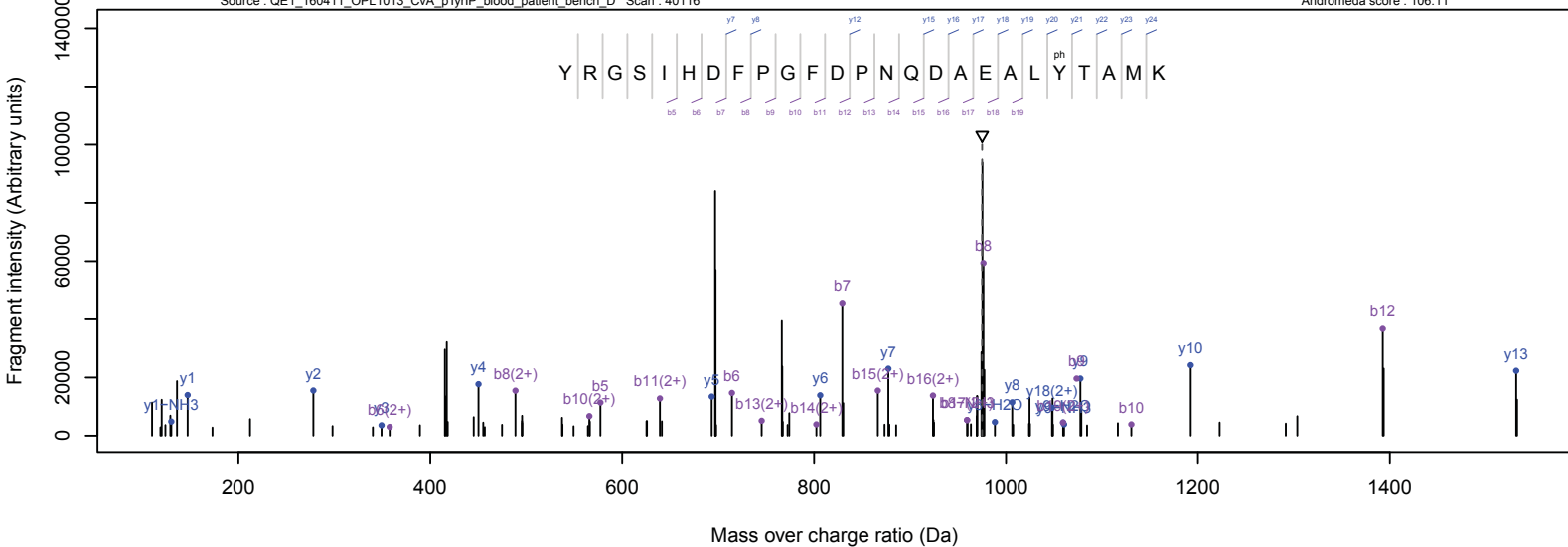
Andromeda score : 106.29



YRGSIHDFPGFDPNQDAEAL(ph)YTAMK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 40116

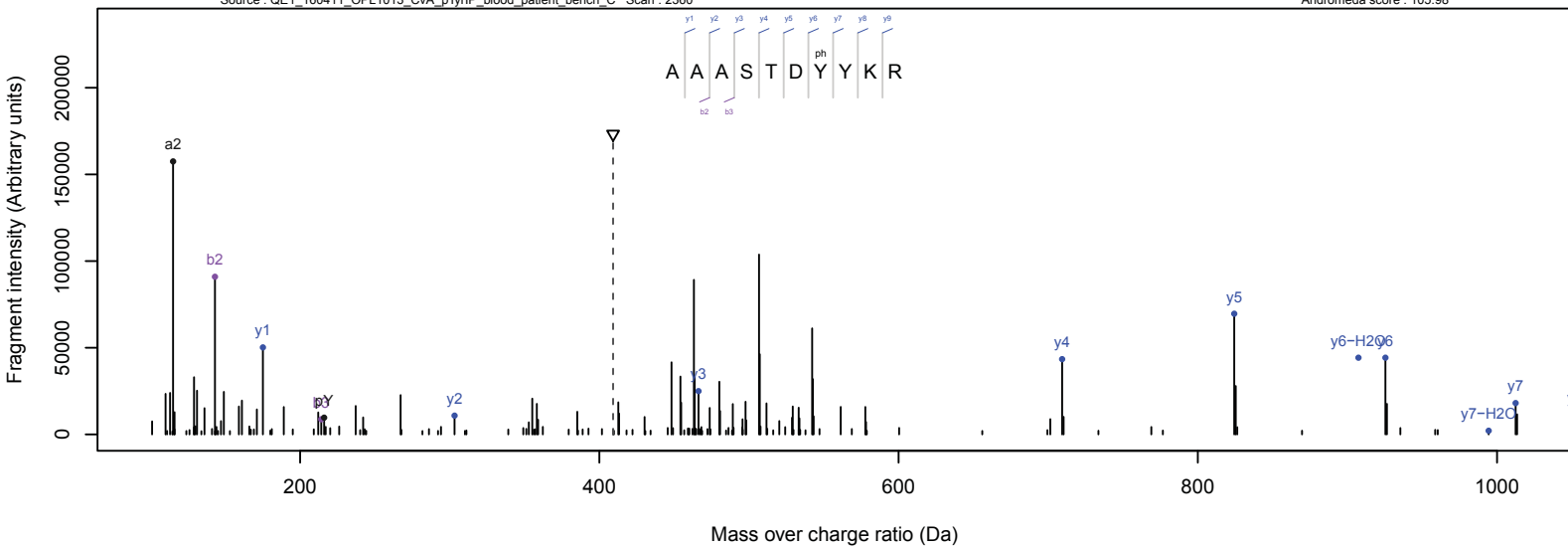
Andromeda score : 106.11

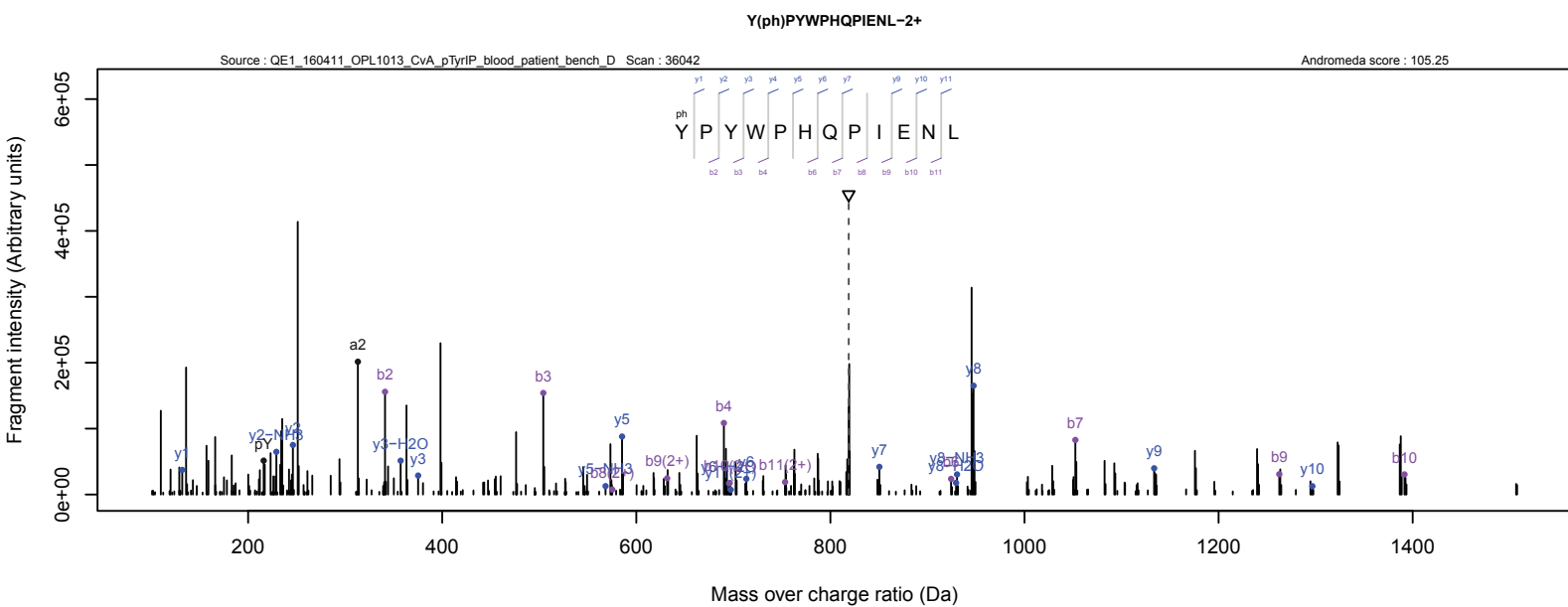
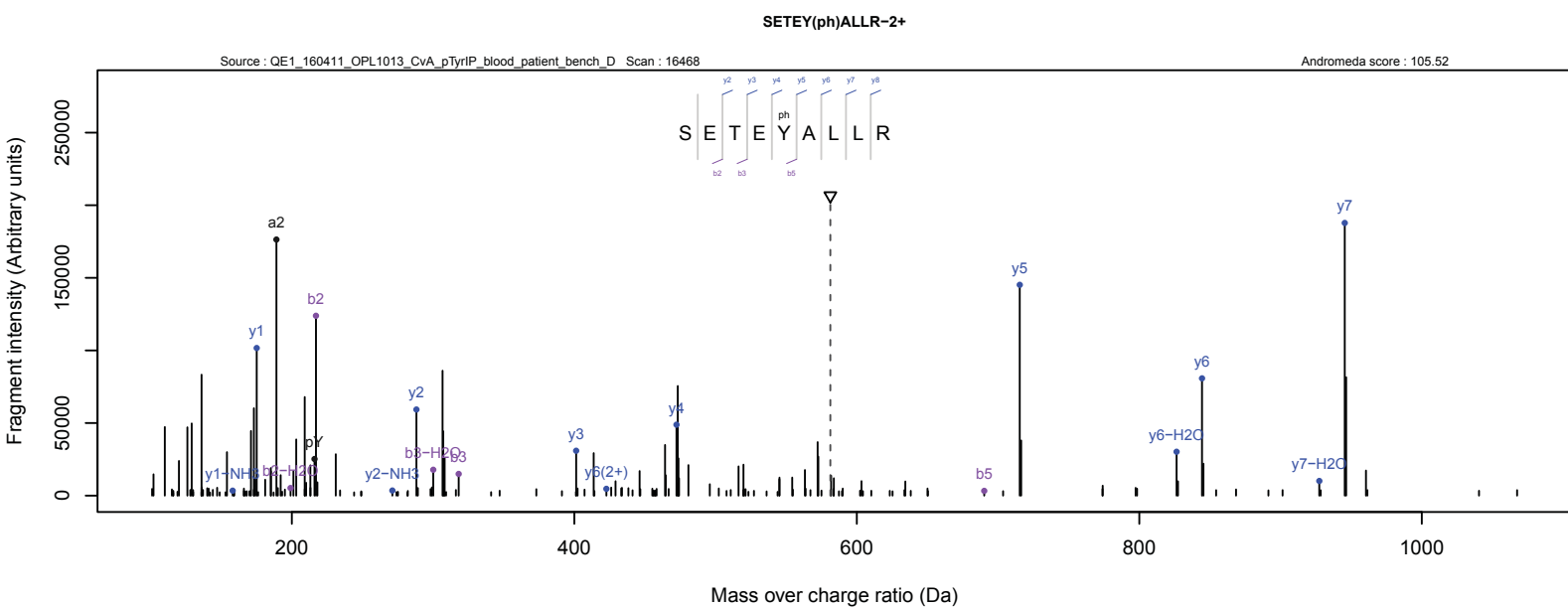
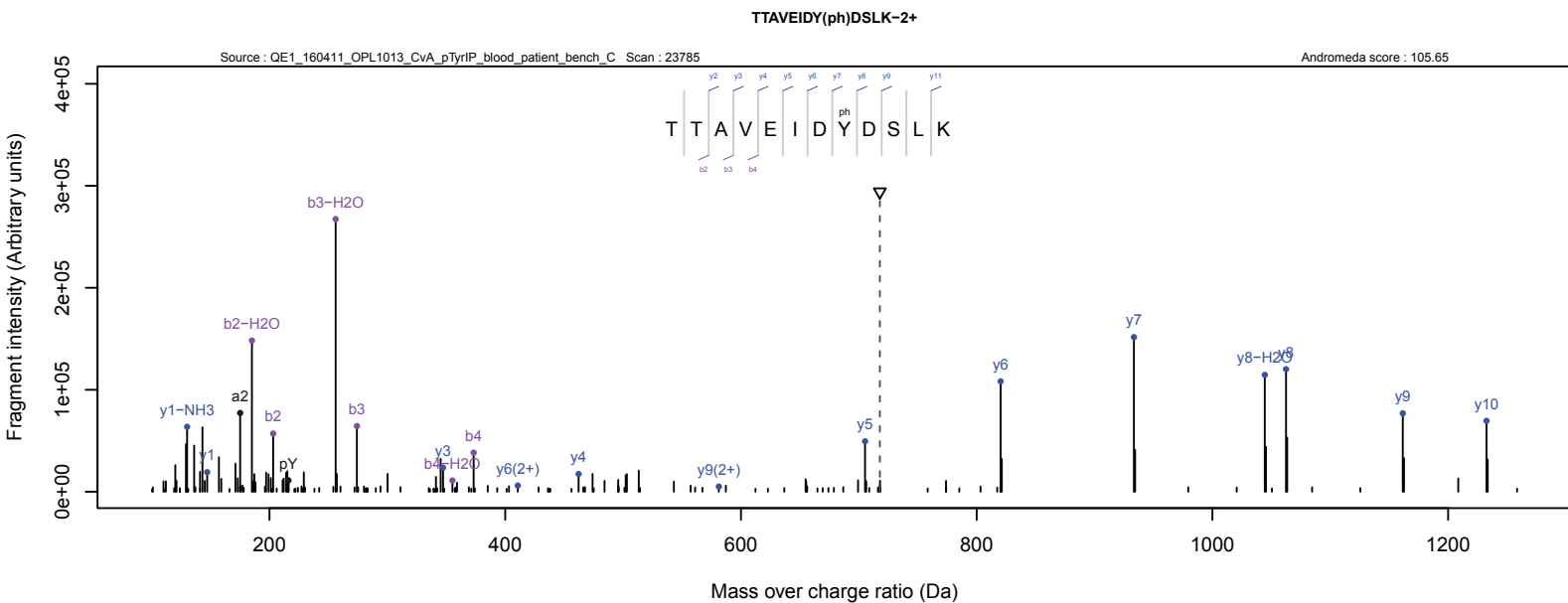


AAASTDY(ph)YKR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 2360

Andromeda score : 105.98



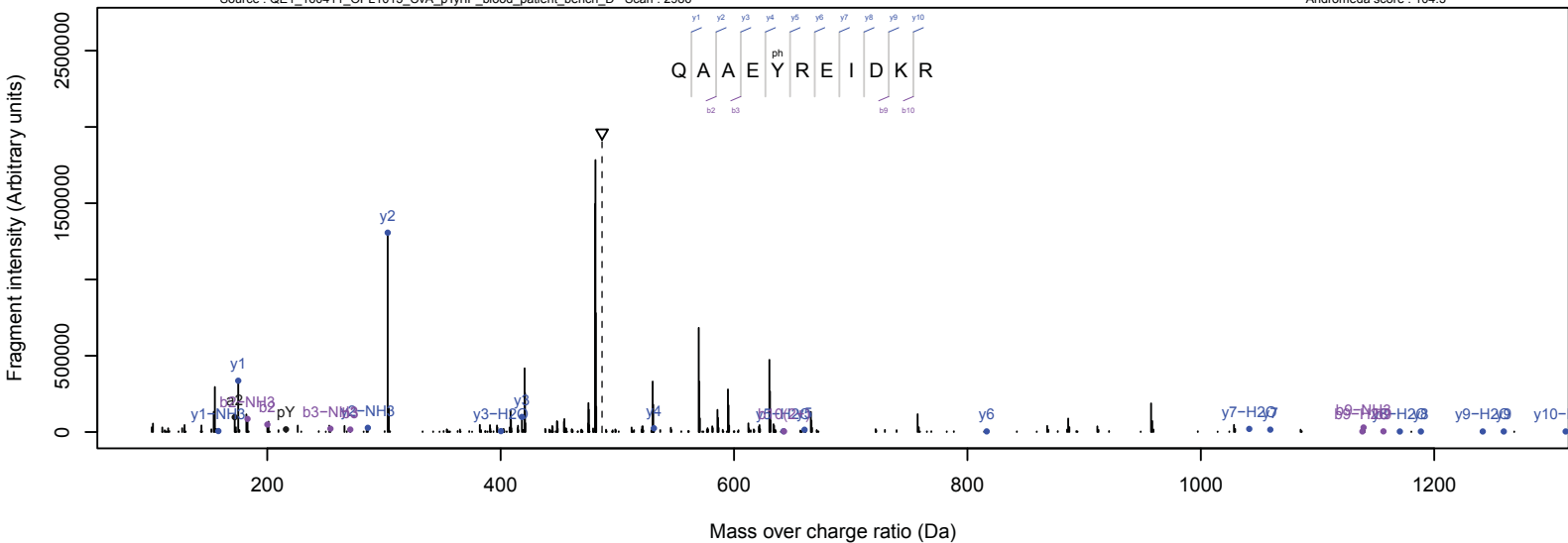




QAAEY(ph)REIDKR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 2986

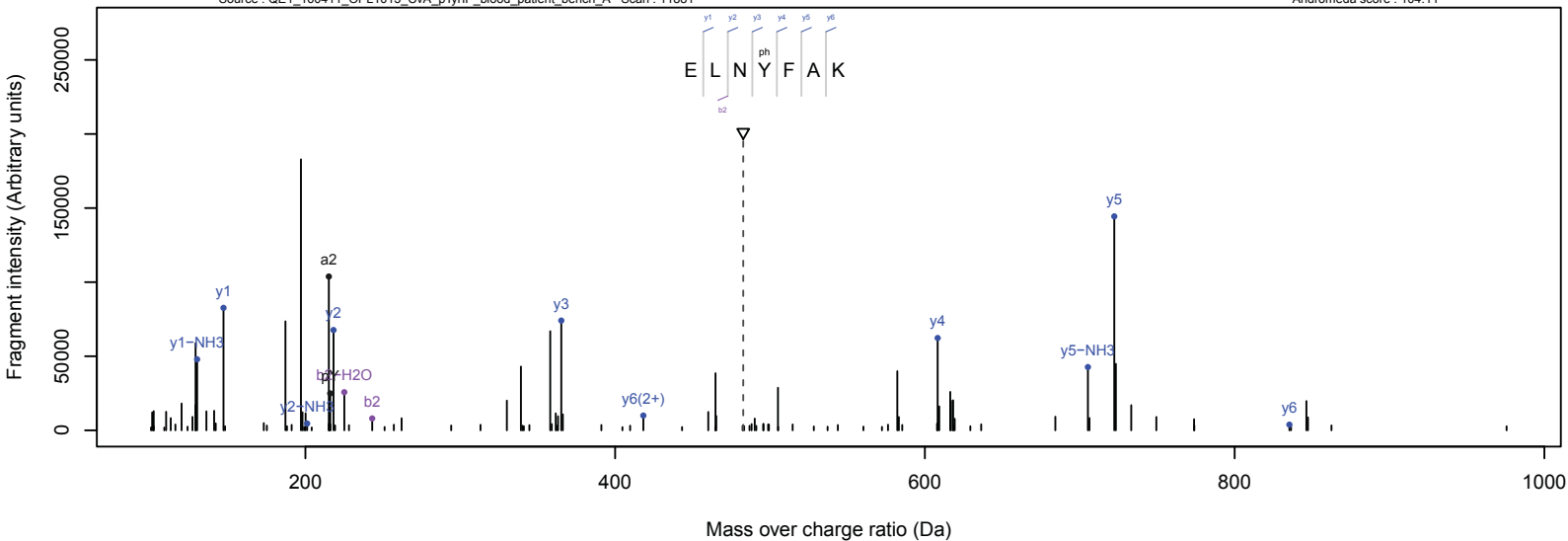
Andromeda score: 104.3



ELNY(ph)FAK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 11881

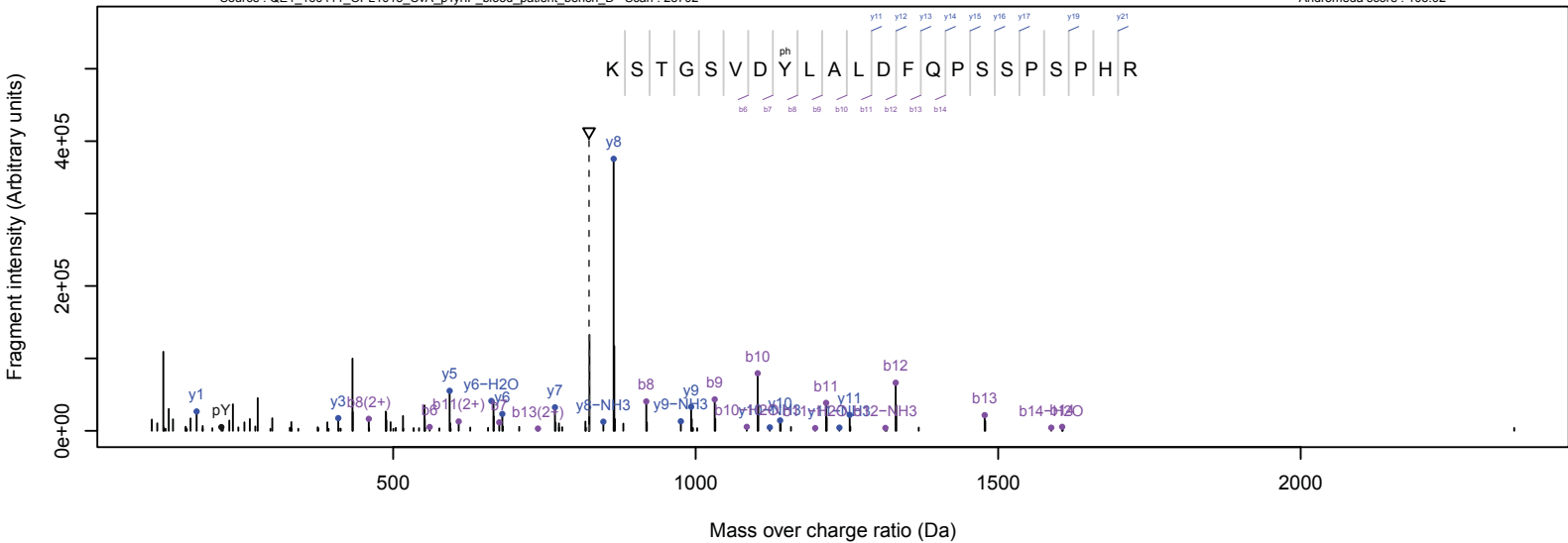
Andromeda score: 104.11



KSTGSVDY(ph)LALDFQPSSPSHR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 28702

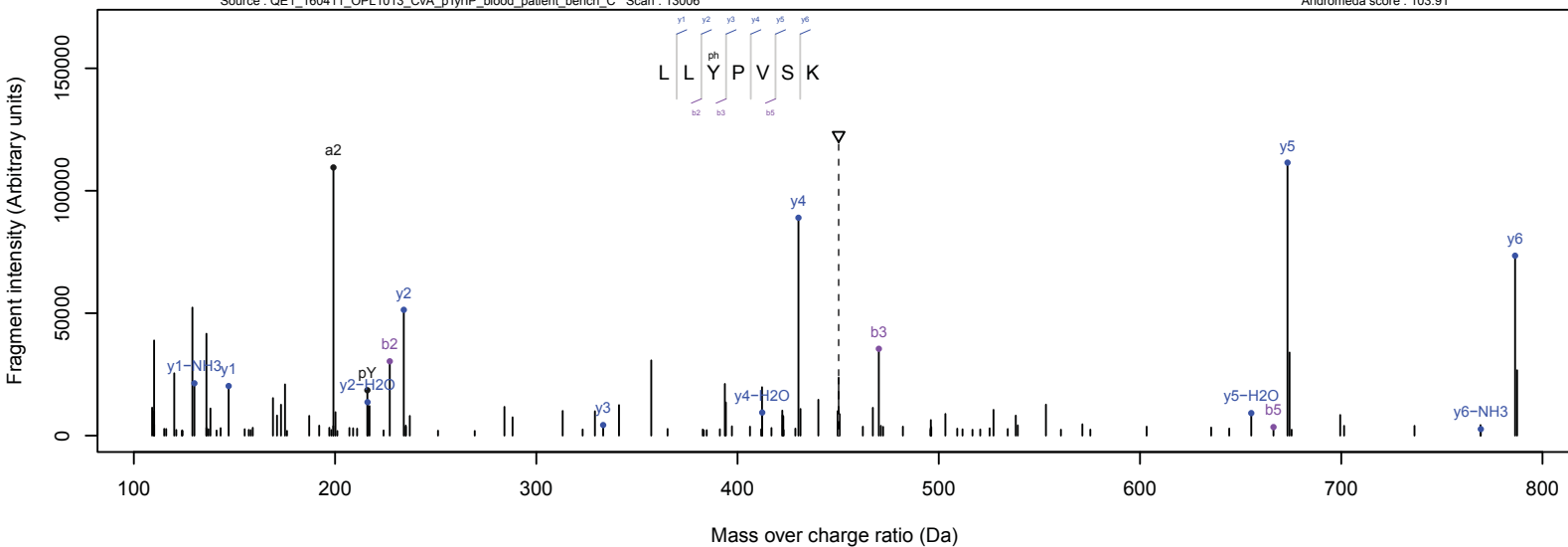
Andromeda score: 103.92



LLY(ph)PVSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 13006

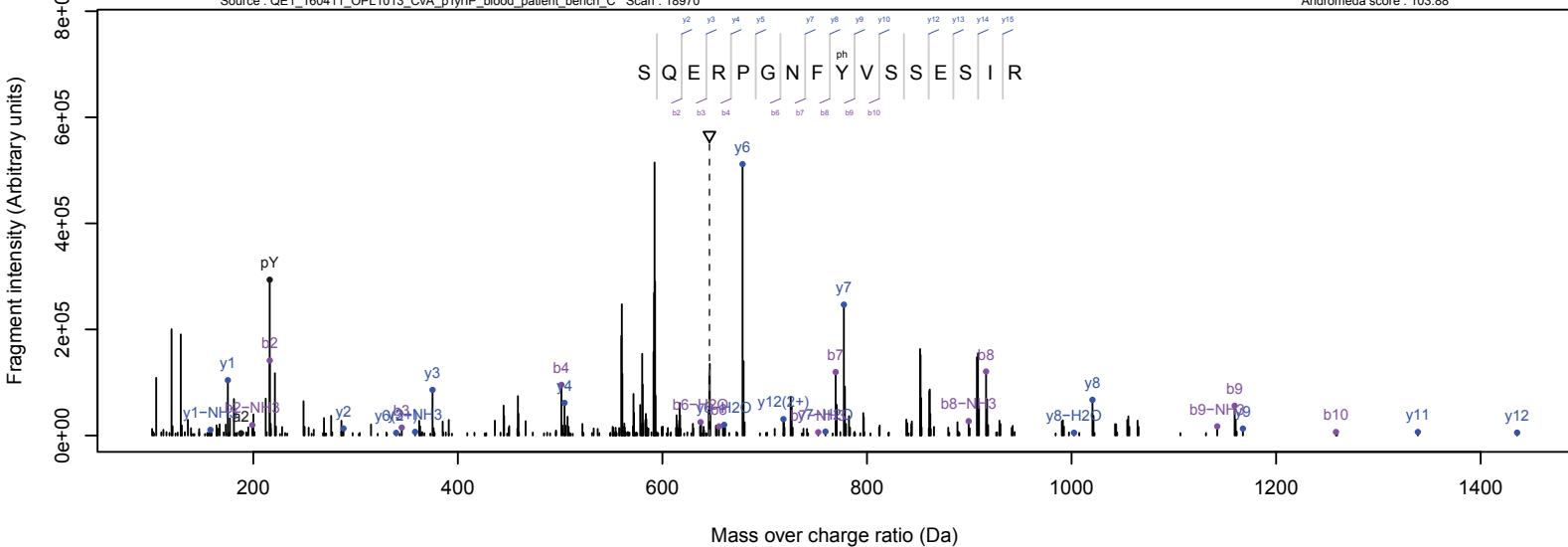
Andromeda score : 103.91



SQERPGNFY(ph)VSSEIR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18970

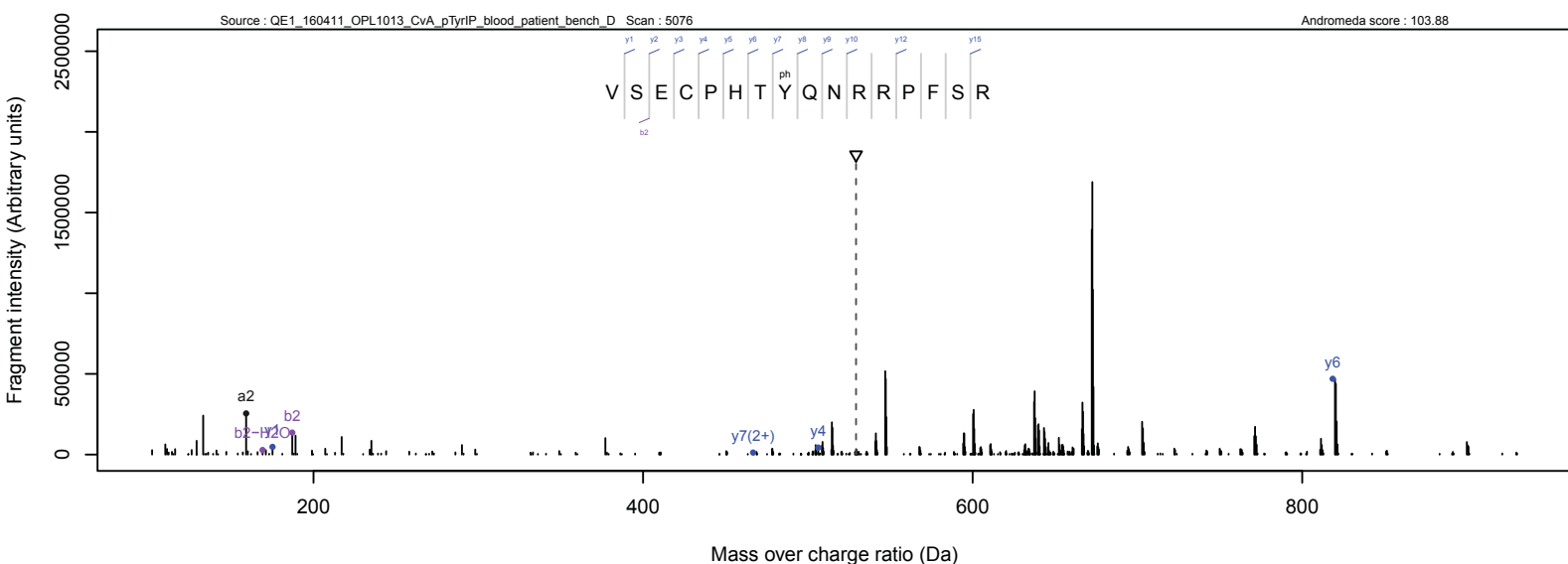
Andromeda score : 103.88



VSECPHTY(ph)QNRPF SR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5076

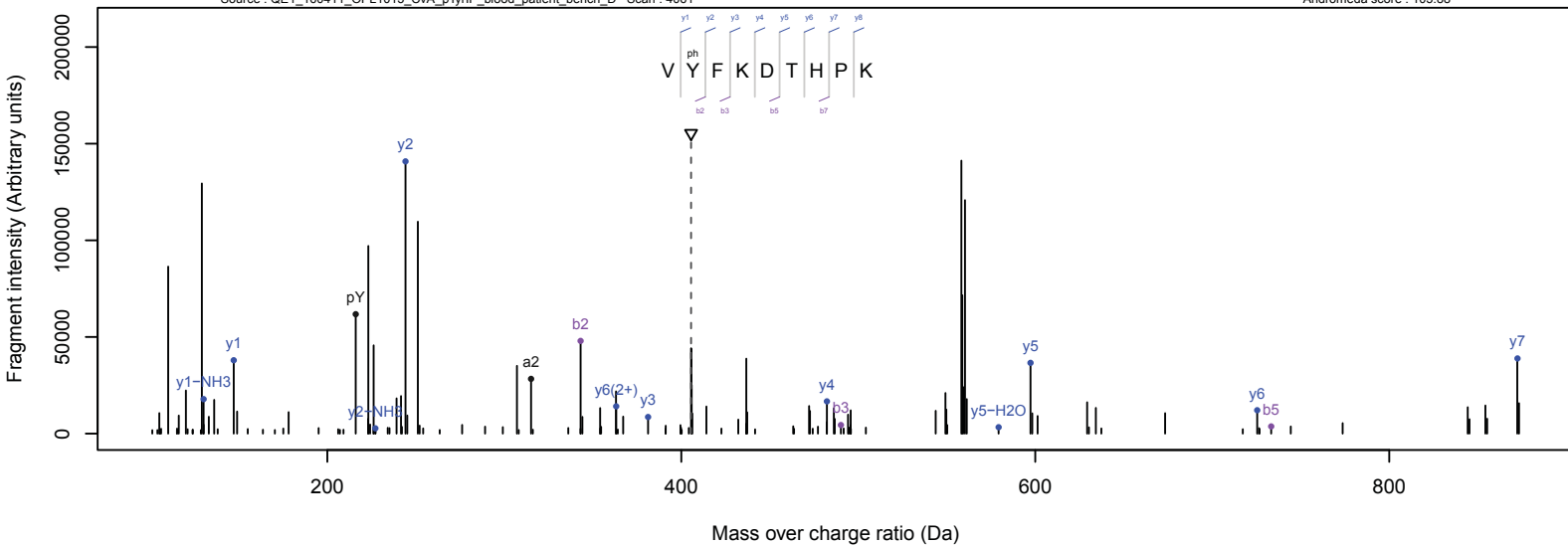
Andromeda score : 103.88



VY(ph)FKDTHPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 4061

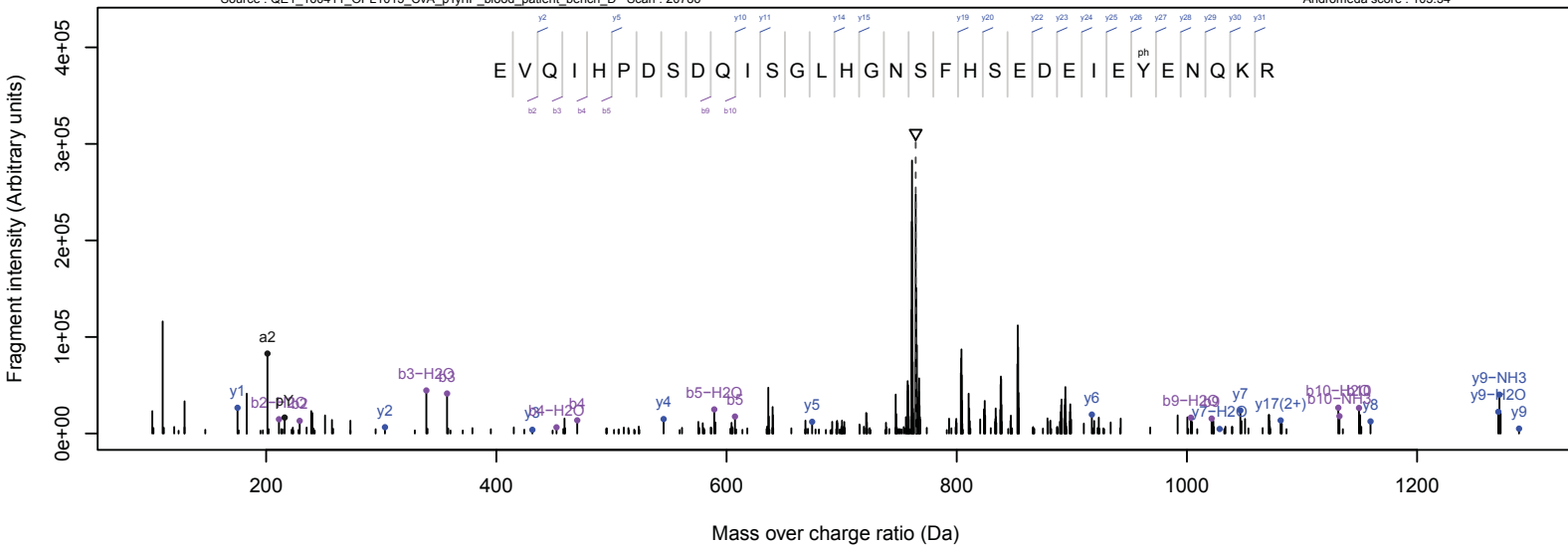
Andromeda score : 103.88



EVQIHPDSDQISGLHGNSFHSEDEIEY(ph)ENQKR-5+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 20786

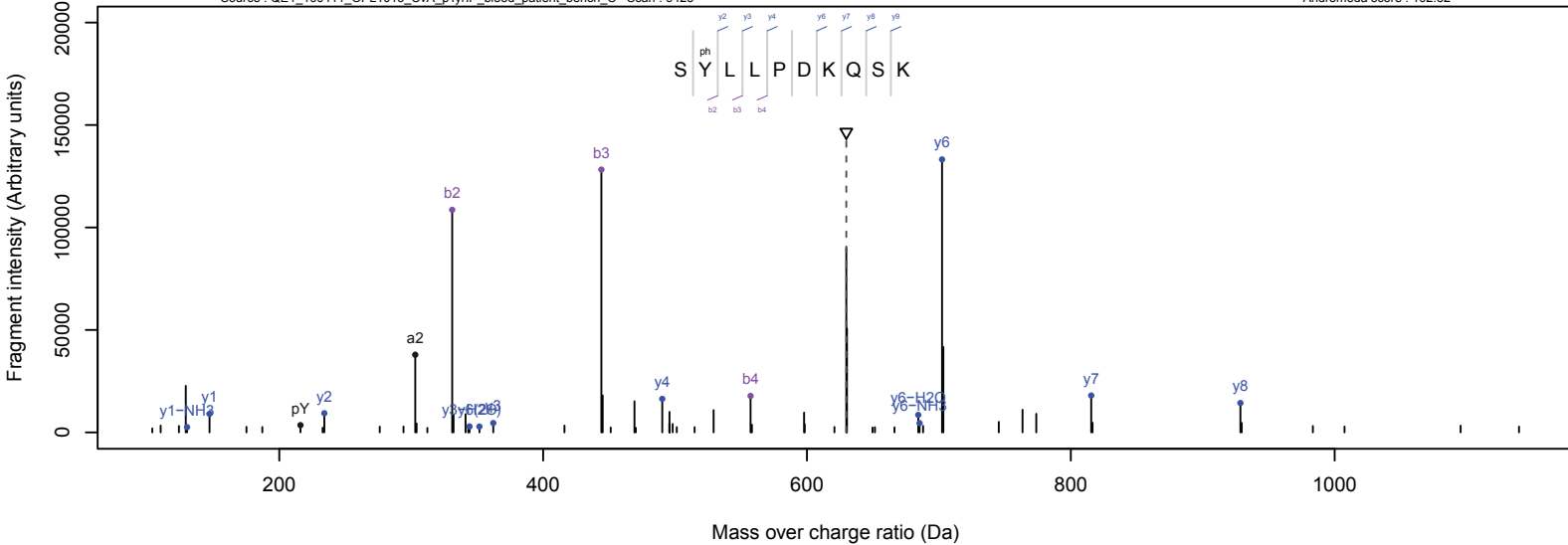
Andromeda score : 103.34



SY(ph)LLPDKQSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 9425

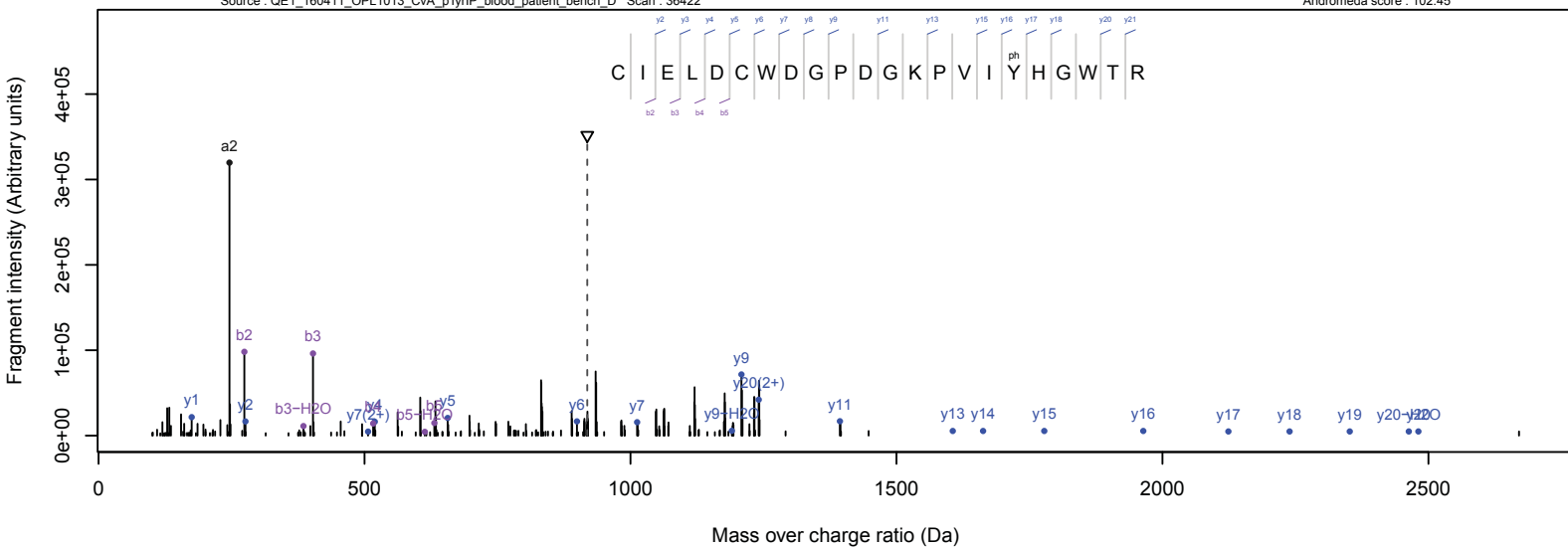
Andromeda score : 102.52



**CIELDCWDGPKVIY<sup>ph</sup>HGWTR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 36422

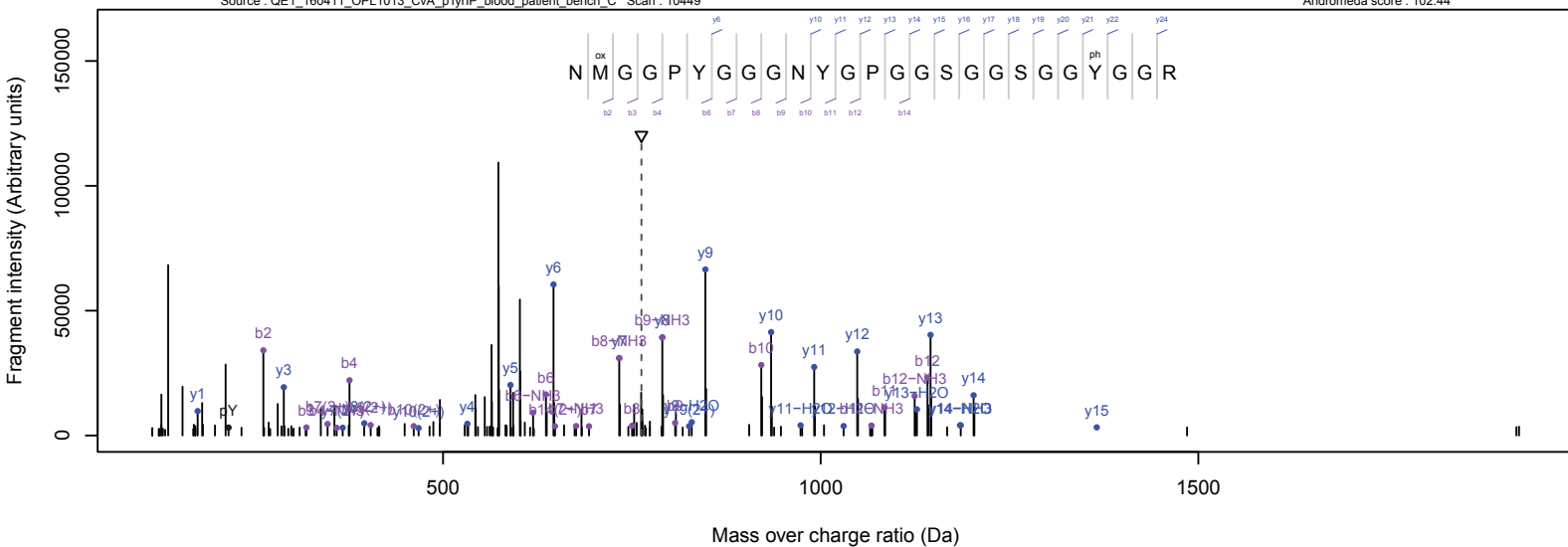
Andromeda score : 102.45



**NM(ox)GGPYGGNYGPGSGGGY<sup>ph</sup>GGR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 10449

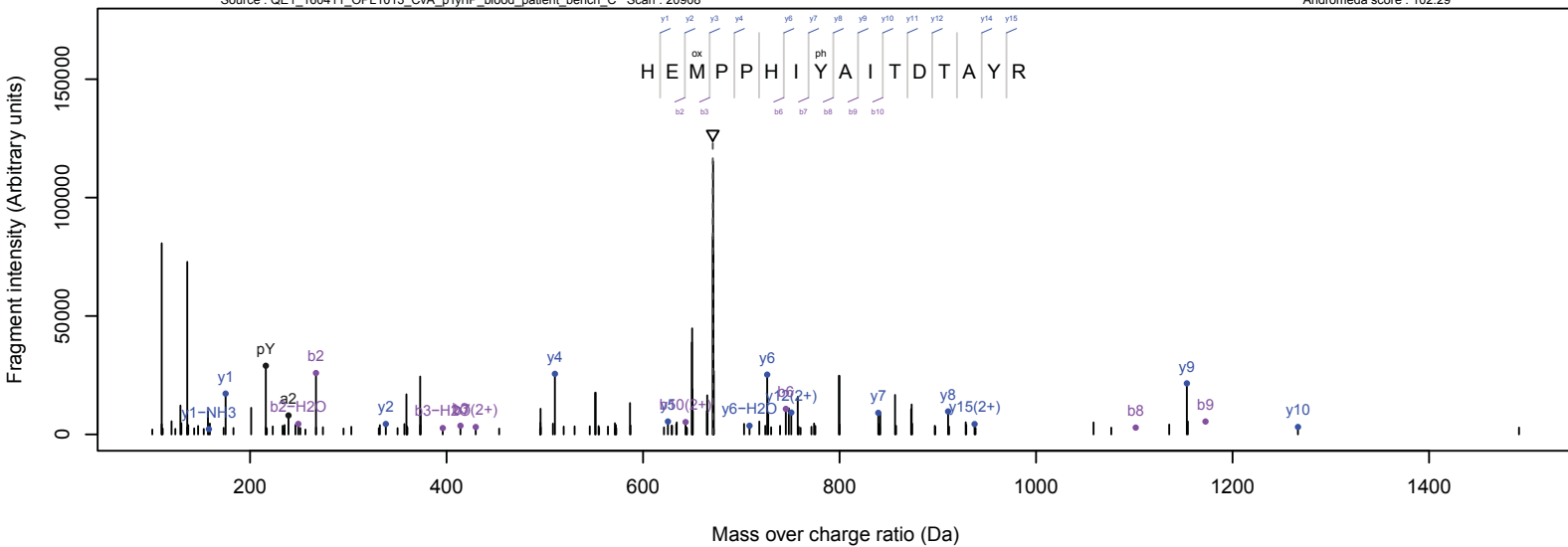
Andromeda score : 102.44



**HEM(ox)PPHIY<sup>ph</sup>AITDTAYR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 20908

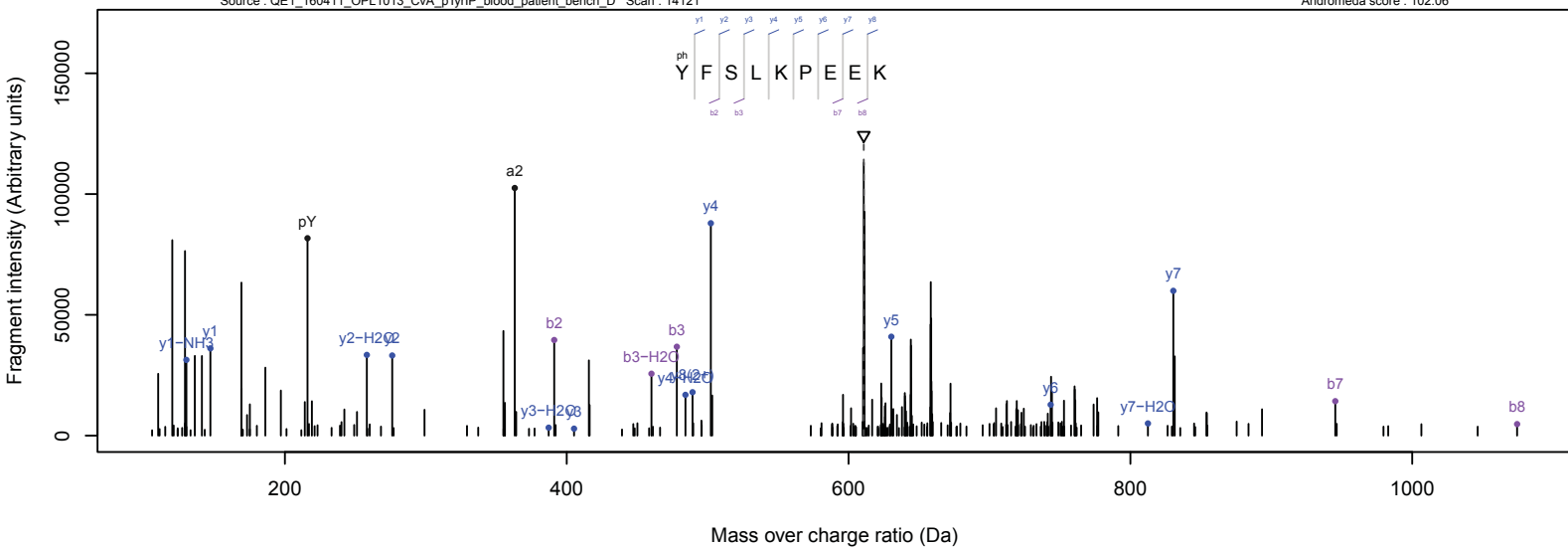
Andromeda score : 102.29



**Y(ph)FSLKPEEK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 14121

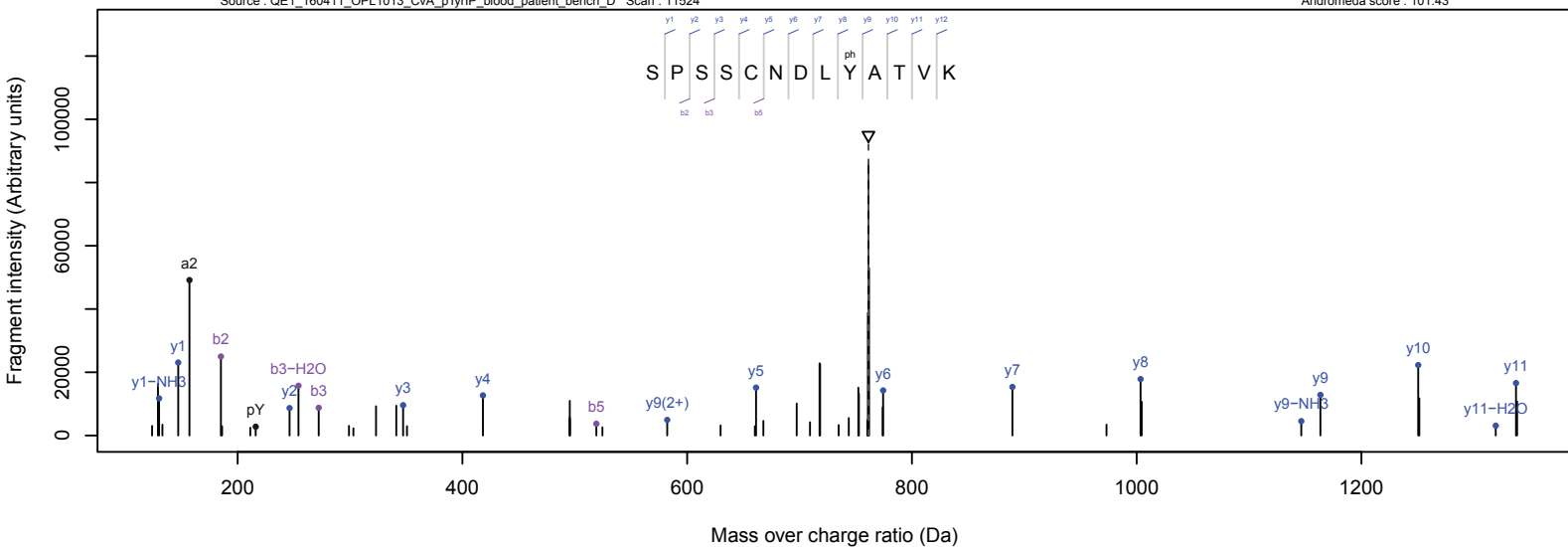
Andromeda score : 102.06



**SPSSCNDLY(ph)ATVK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11524

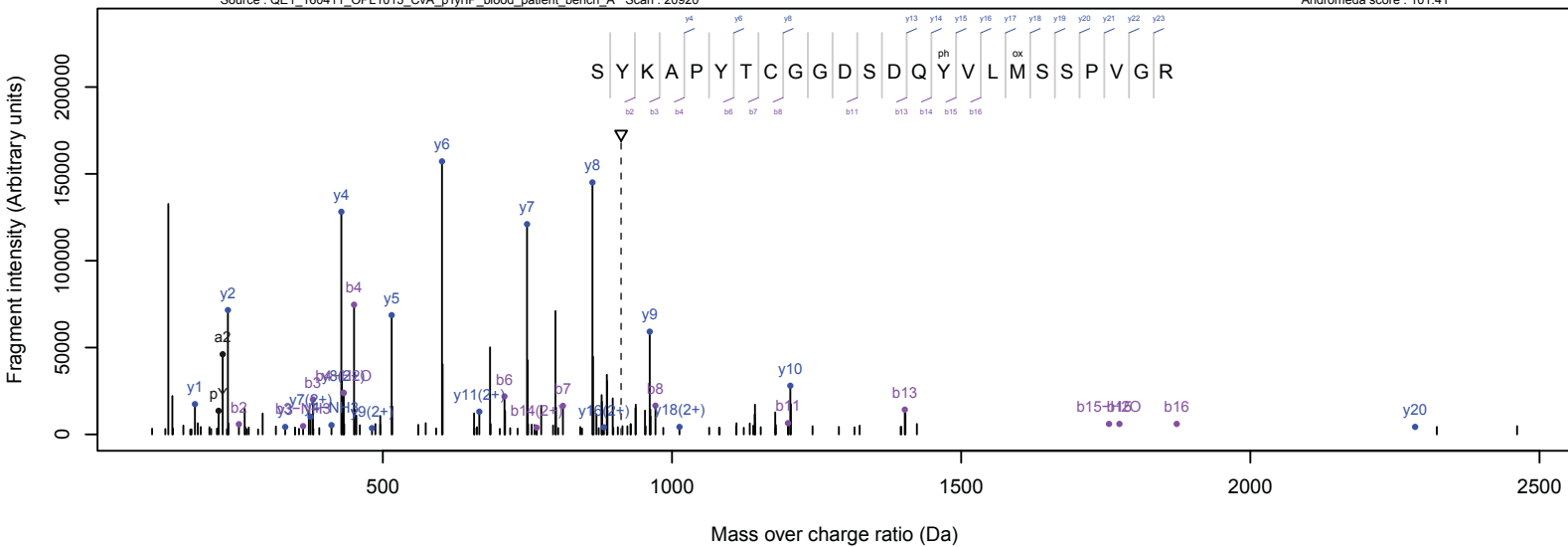
Andromeda score : 101.43



**SYKAPYTCGGDSQY(ph)VLMSSPVGR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 20920

Andromeda score : 101.41

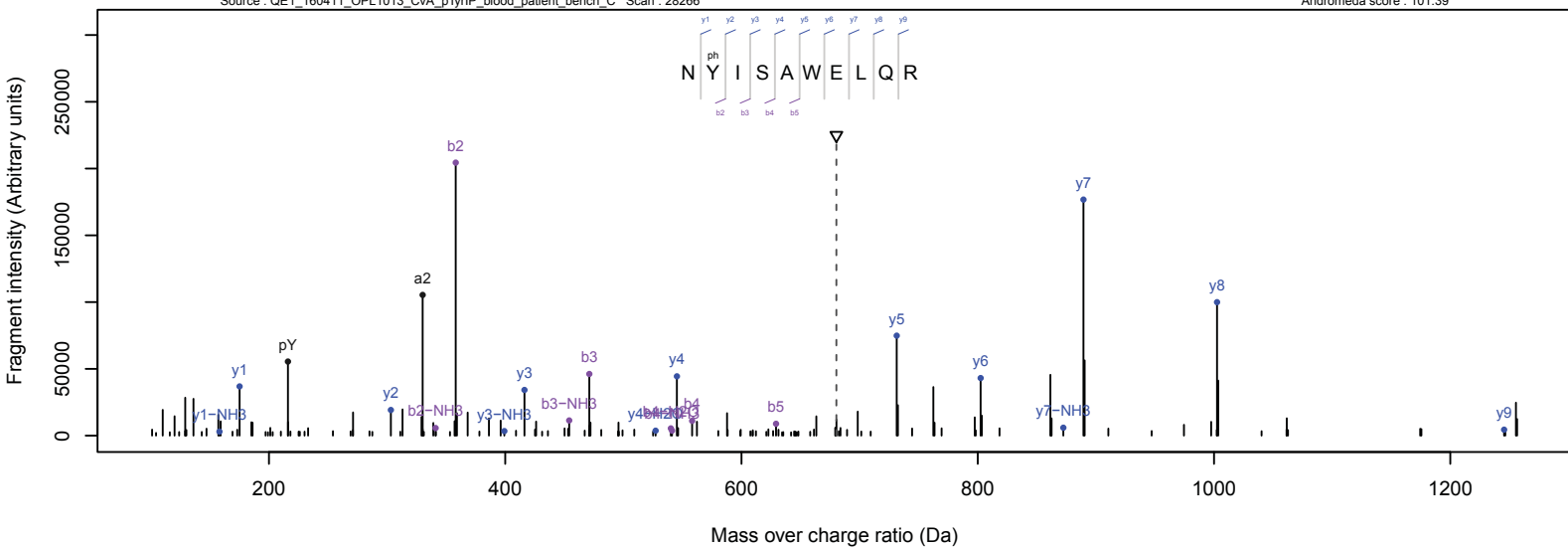




**NY(ph)ISAWELQR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28266

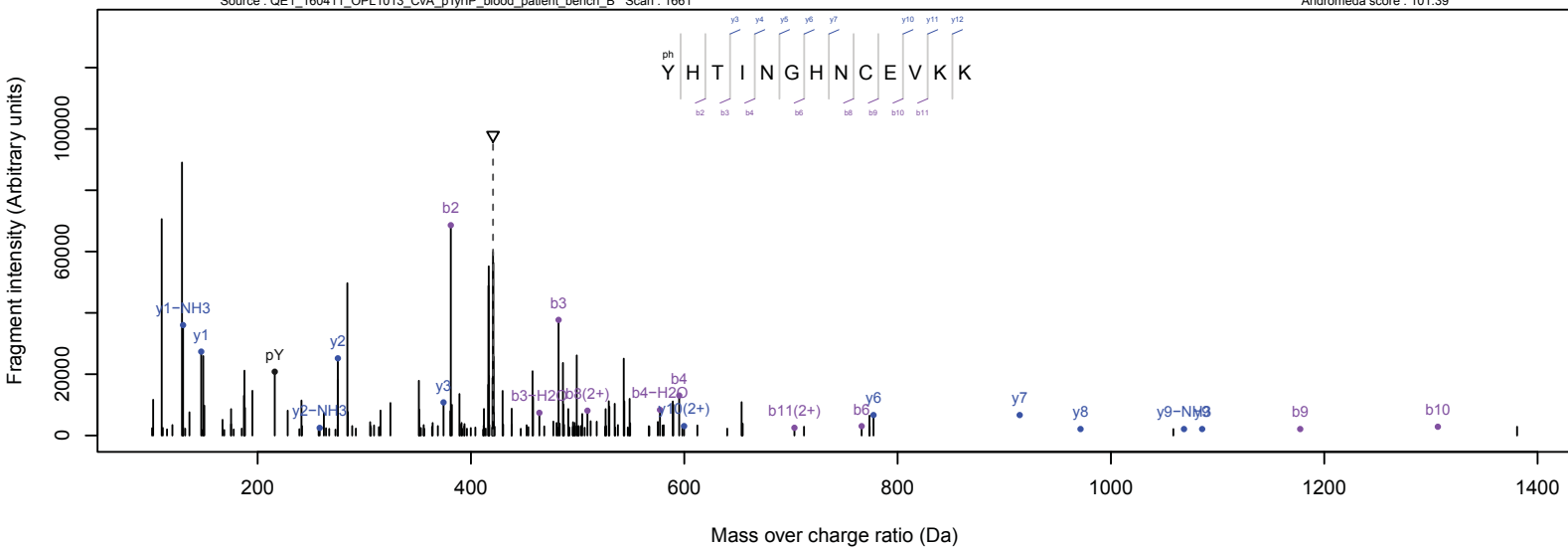
Andromeda score : 101.39



**Y(ph)HTINGHNCVKK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 1661

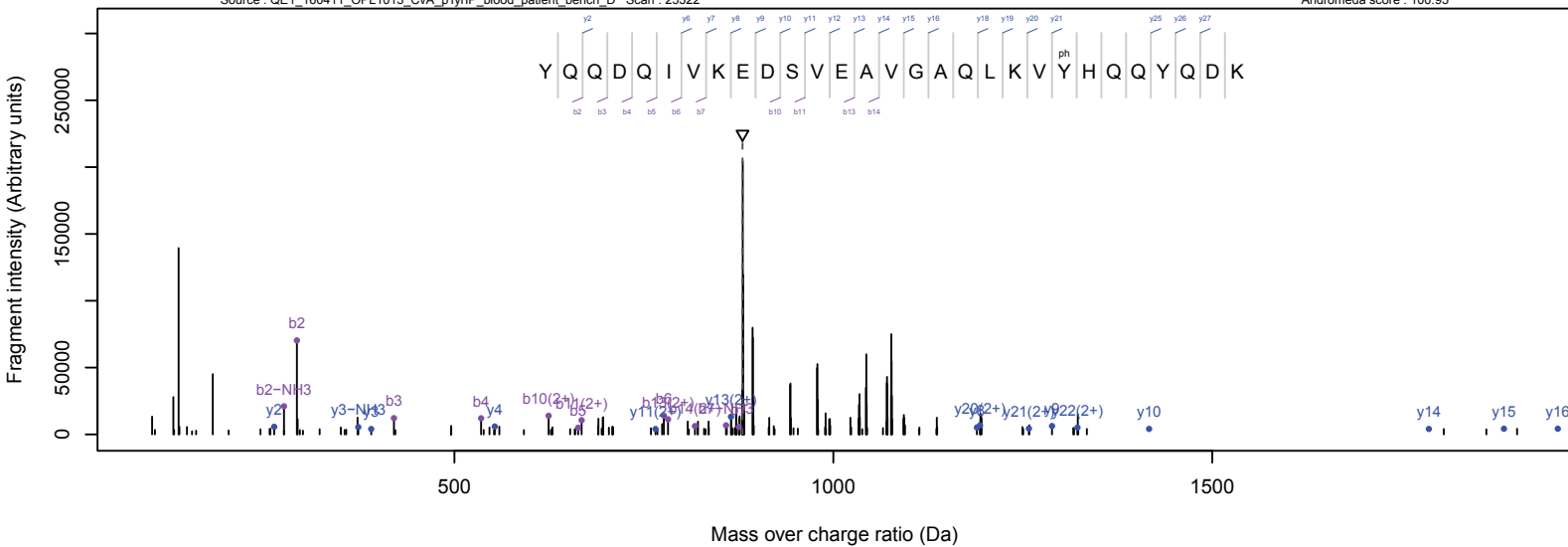
Andromeda score : 101.39



**YQQDQIVKEDSVEAVGAQLKV(ph)HQYQDK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 25322

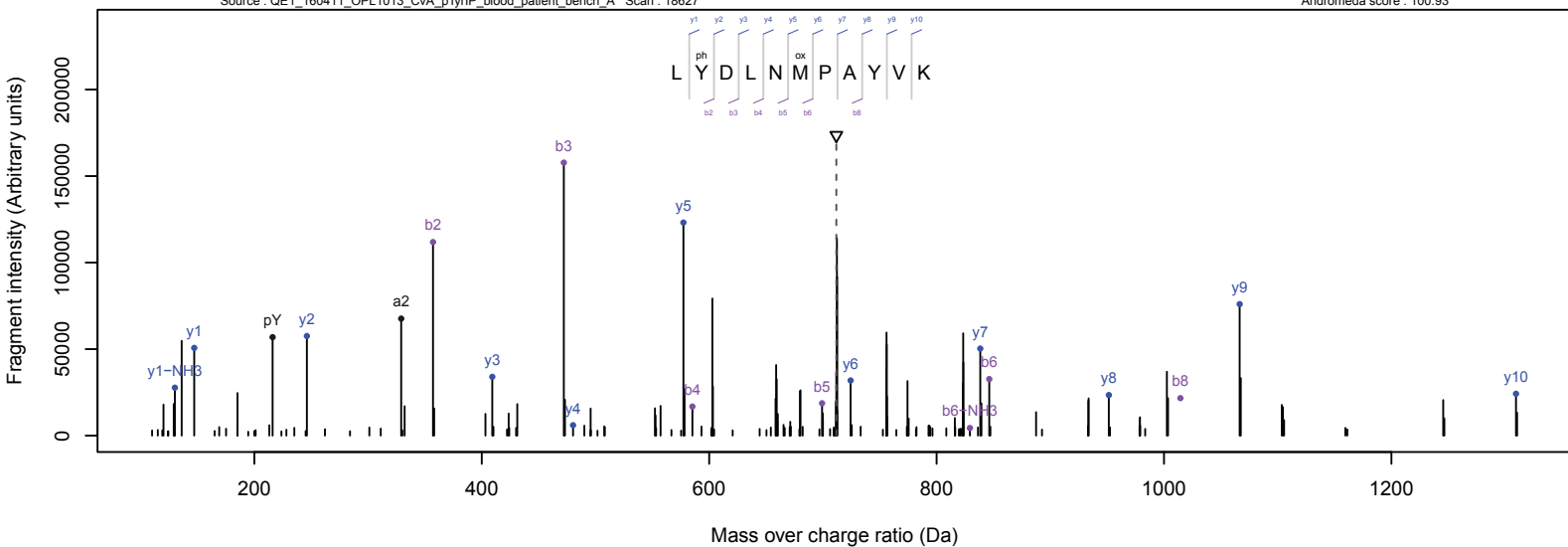
Andromeda score : 100.95



LY(ph)DLNM(ox)PAYVK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 18627

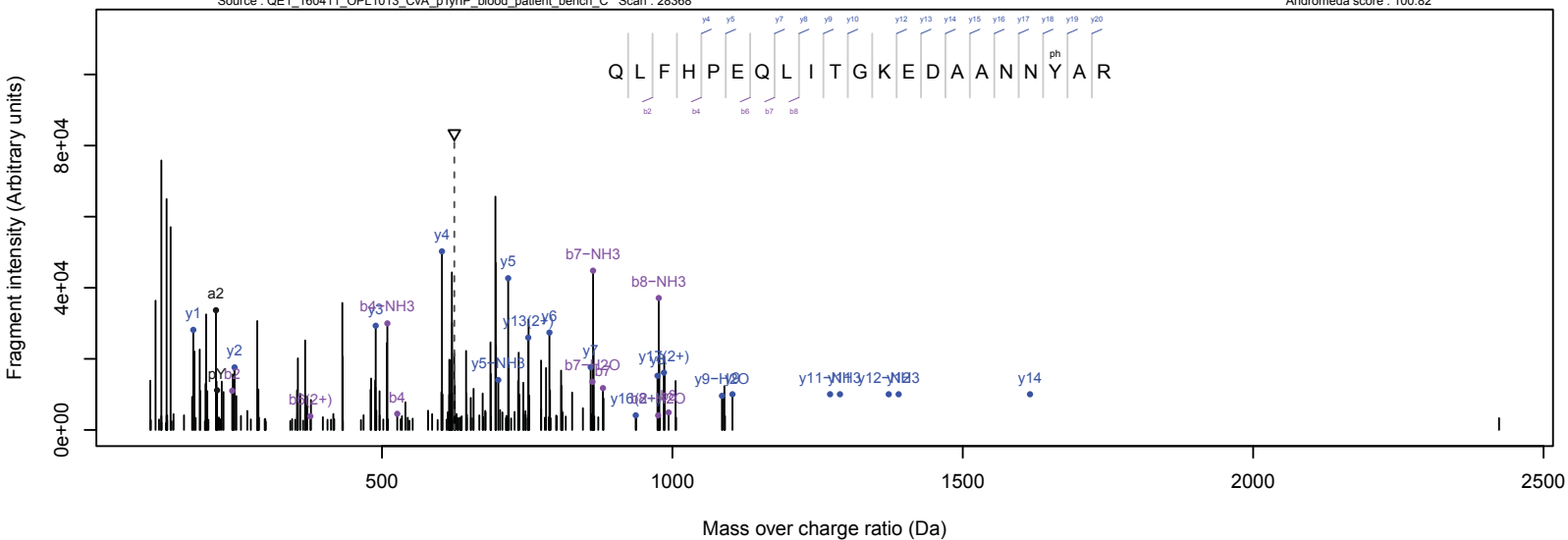
Andromeda score : 100.93



QLFHPEQLITGKEDAANNY(ph)AR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28368

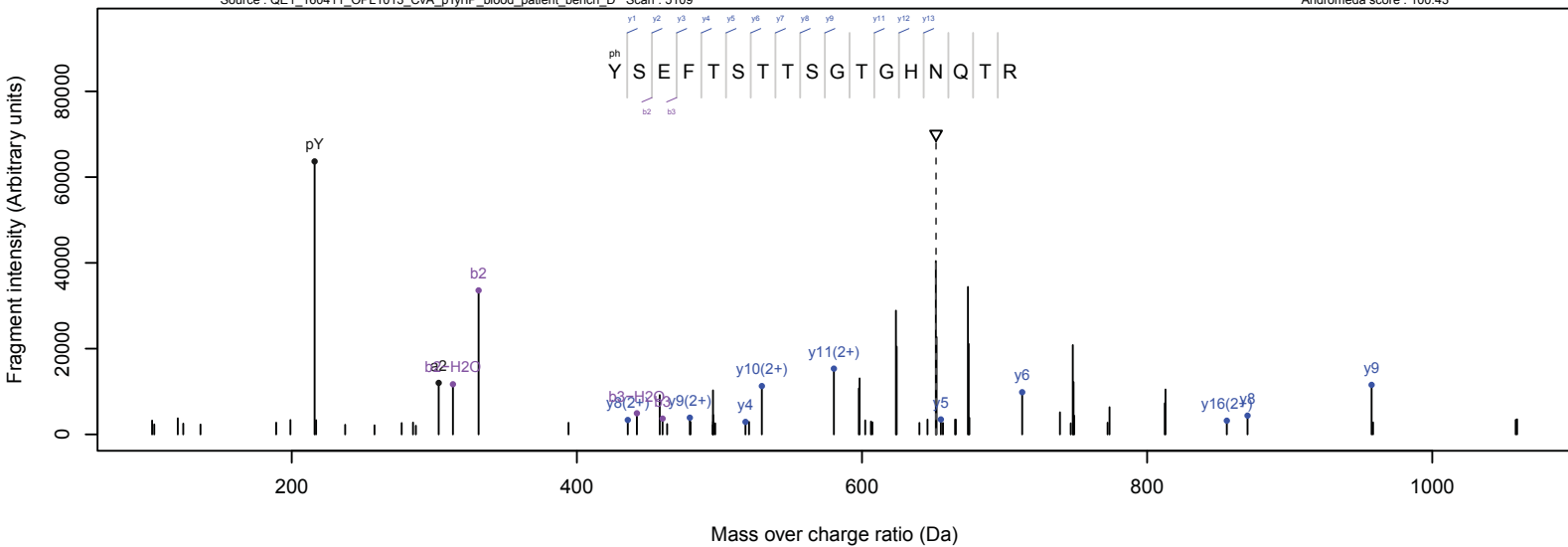
Andromeda score : 100.82



Y(ph)SEFTSTTSGTGHNQTR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5109

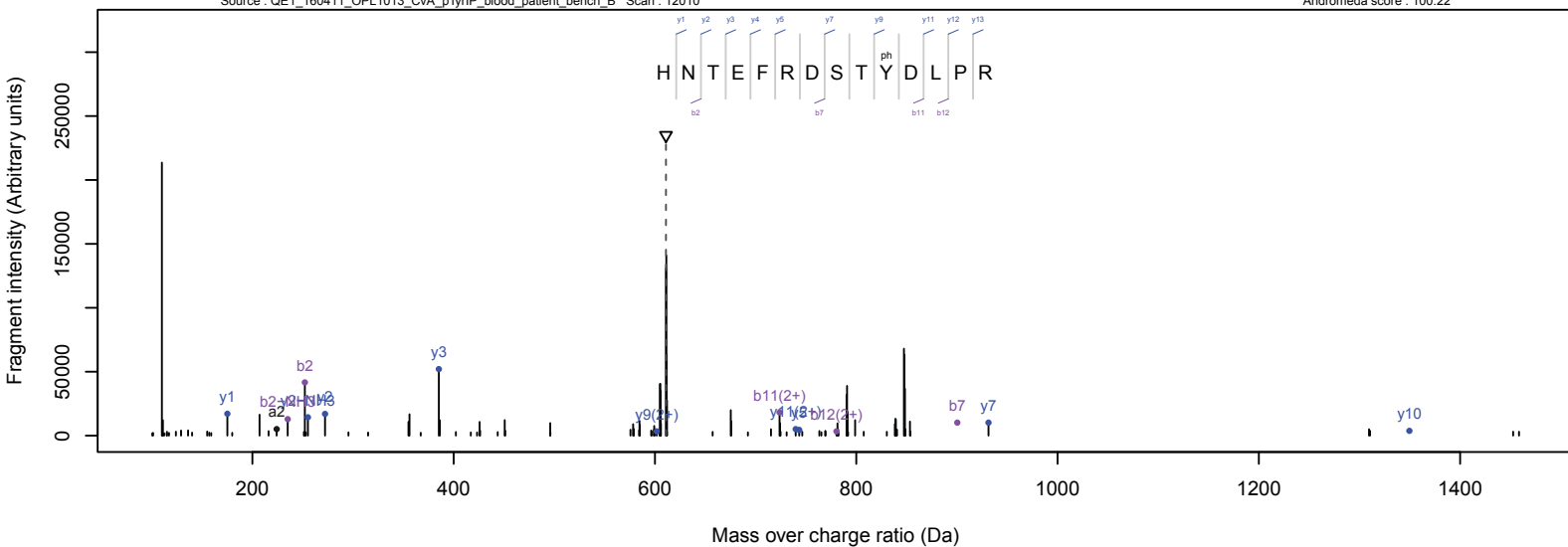
Andromeda score : 100.43



**HNTEFRDSTY(ph)DLPR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 12010

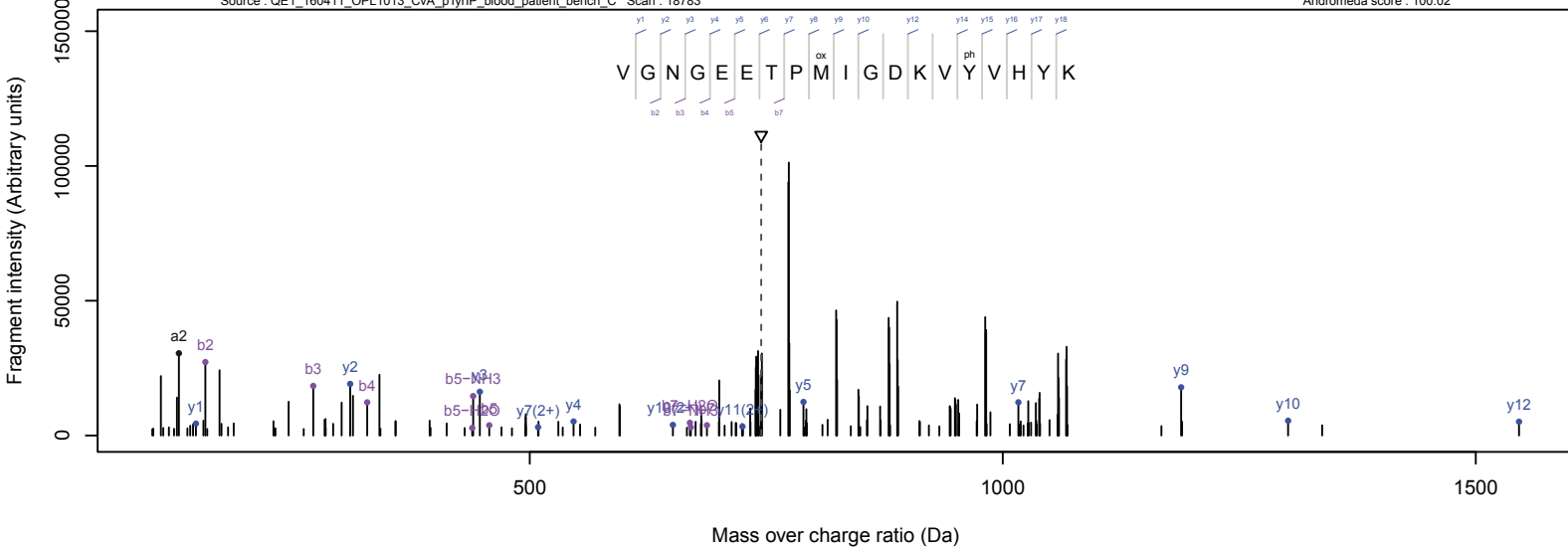
Andromeda score : 100.22



**VGNGEETPM(ox)IGDKVY(ph)VHYK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18783

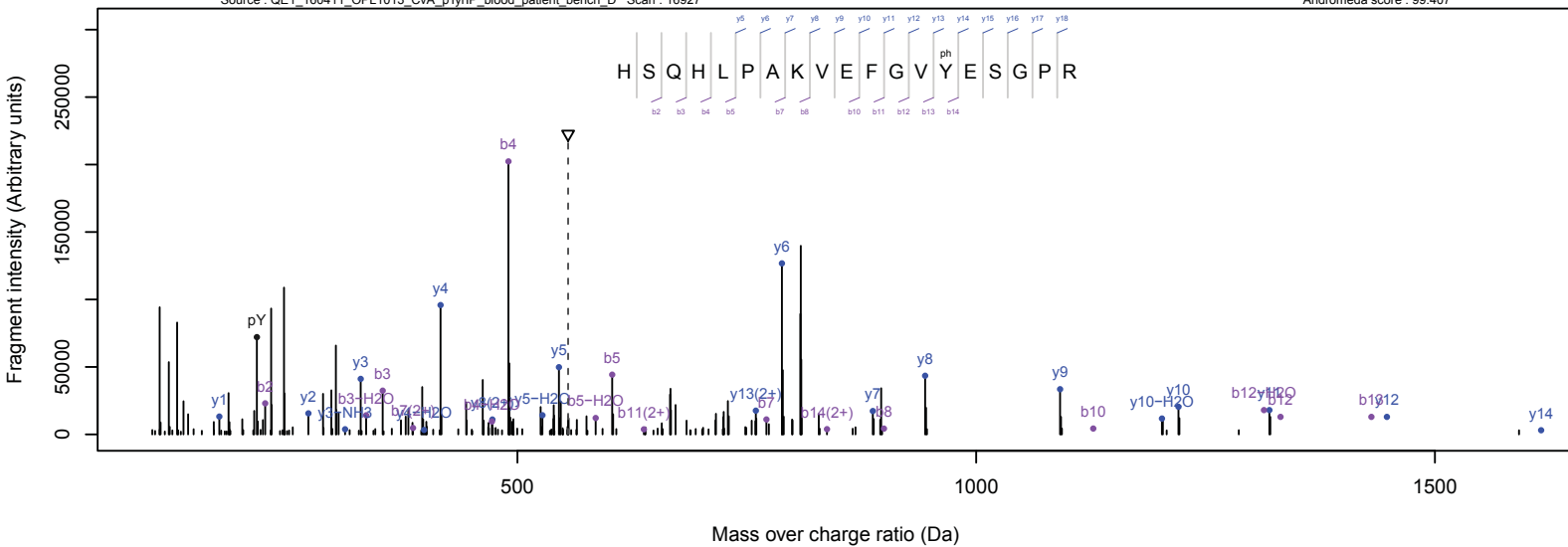
Andromeda score : 100.02



**HSQHLPKVEFGVY(ph)ESGPR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 16927

Andromeda score : 99.407

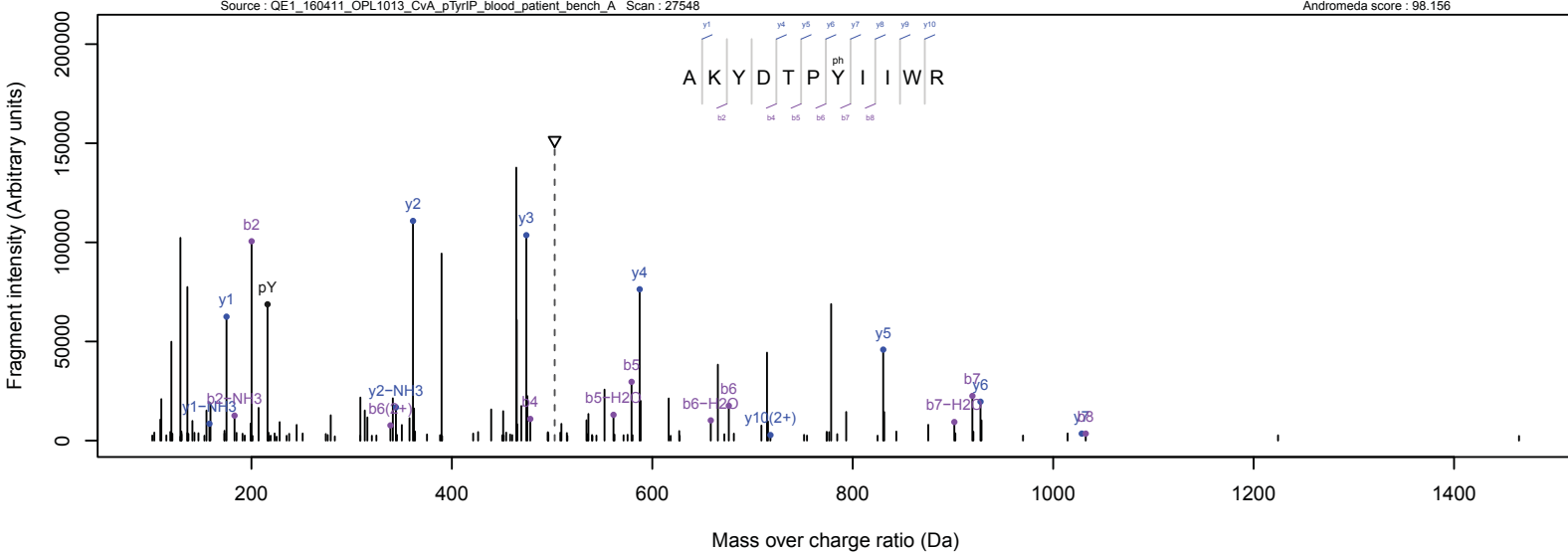




## AKYDTPY(ph)IWR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 27548

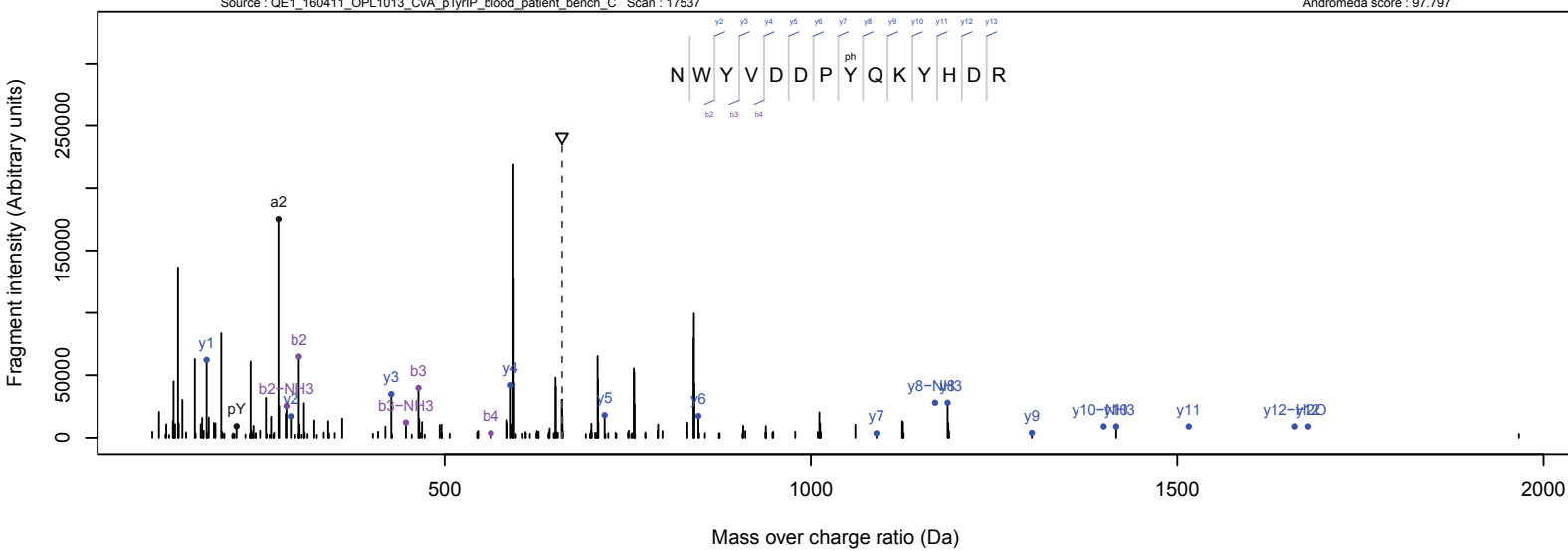
Andromeda score: 98.156



## NWYVDDPY(ph)QKYHDR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 17537

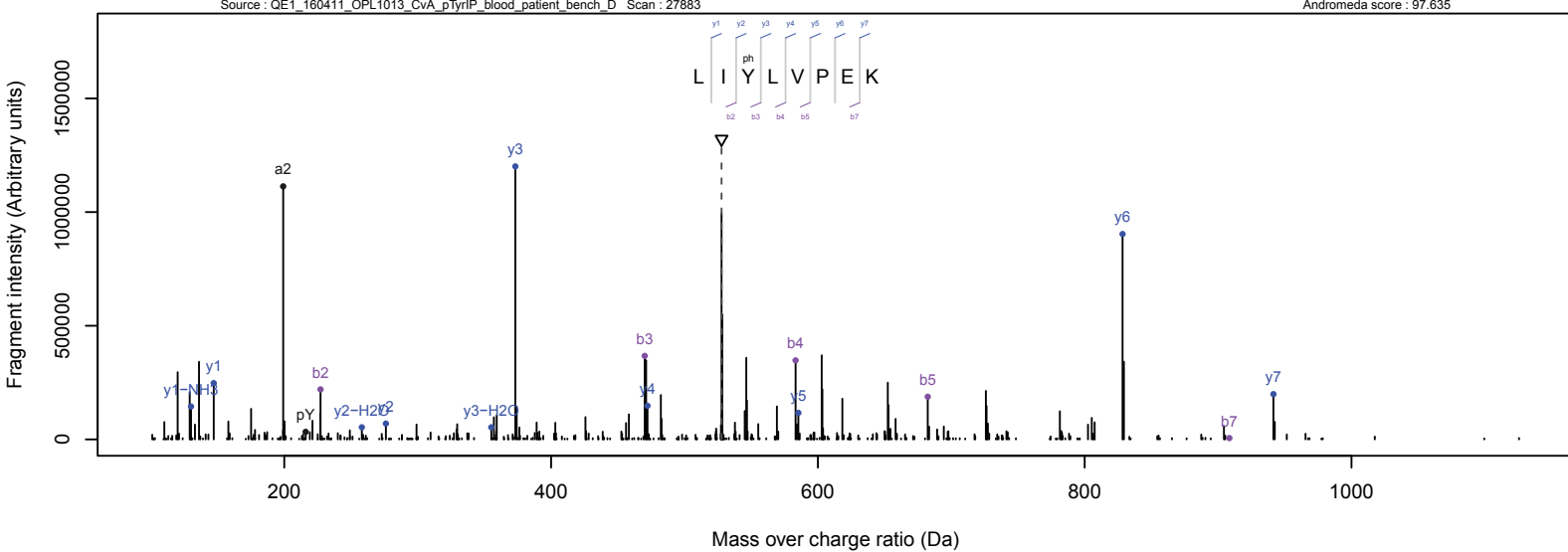
Andromeda score: 97.797



## LIY(ph)LVPEK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 27883

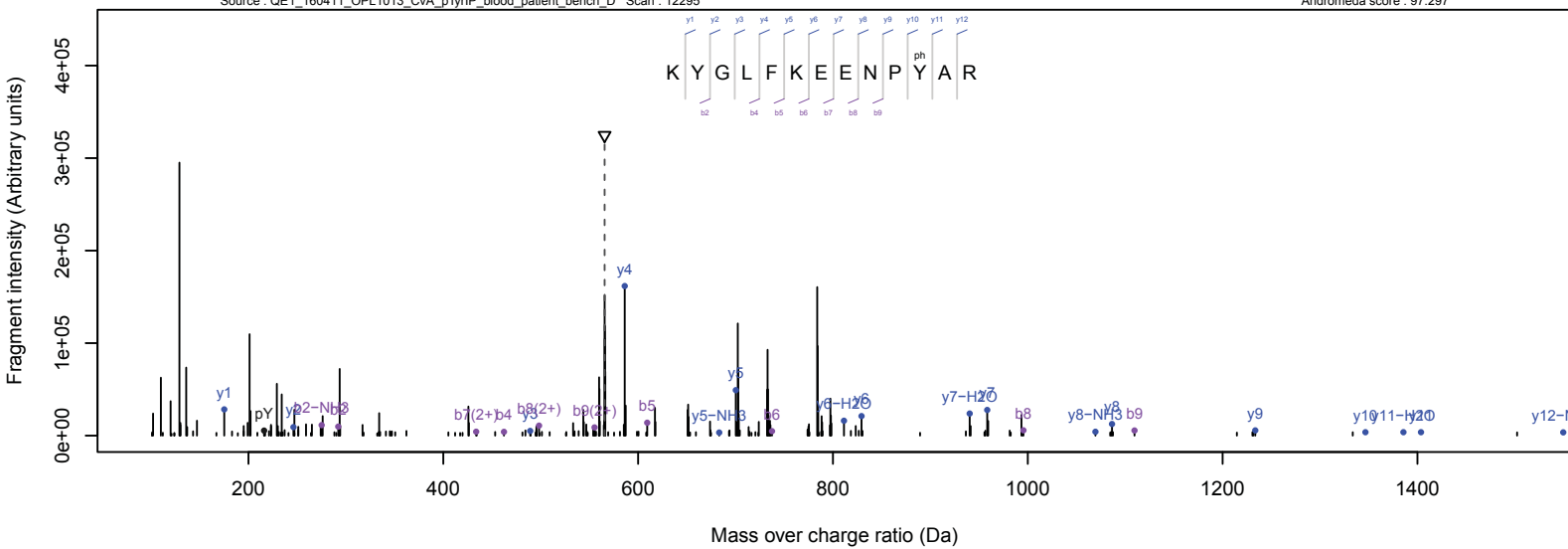
Andromeda score: 97.635



## KYGLFKEENPYAR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 12295

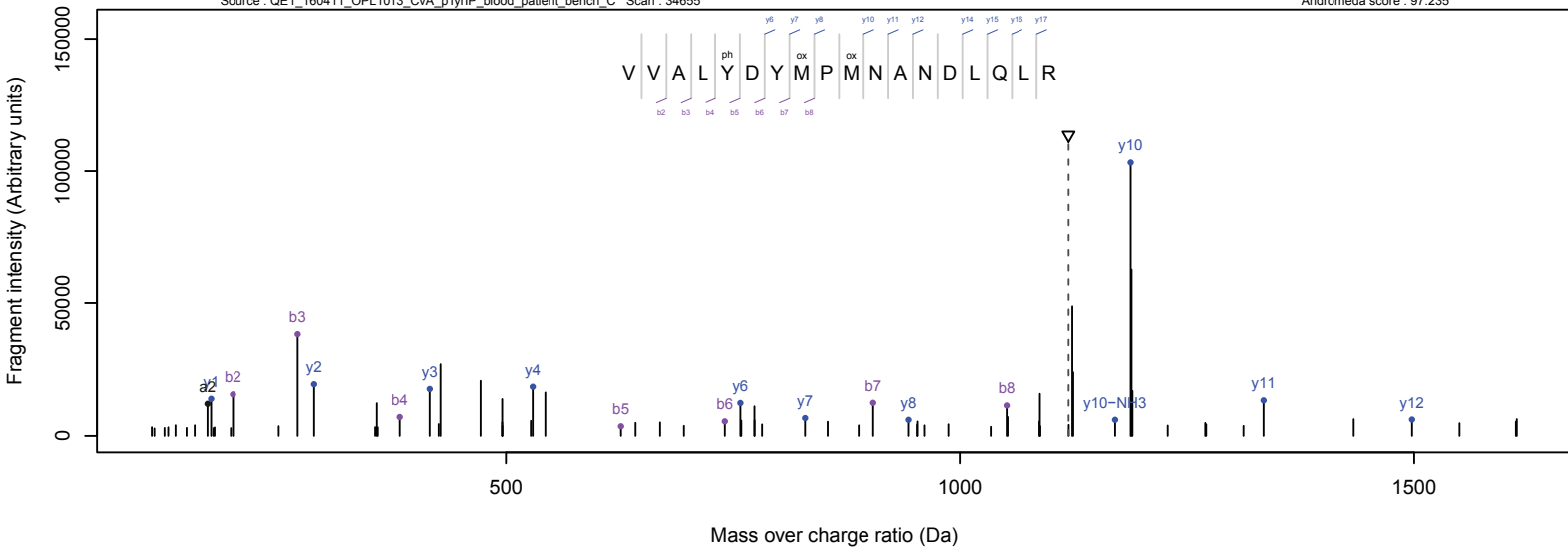
Andromeda score: 97.297



## VVALY(ph)DYM(ox)PM(ox)NANDLQLR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 34655

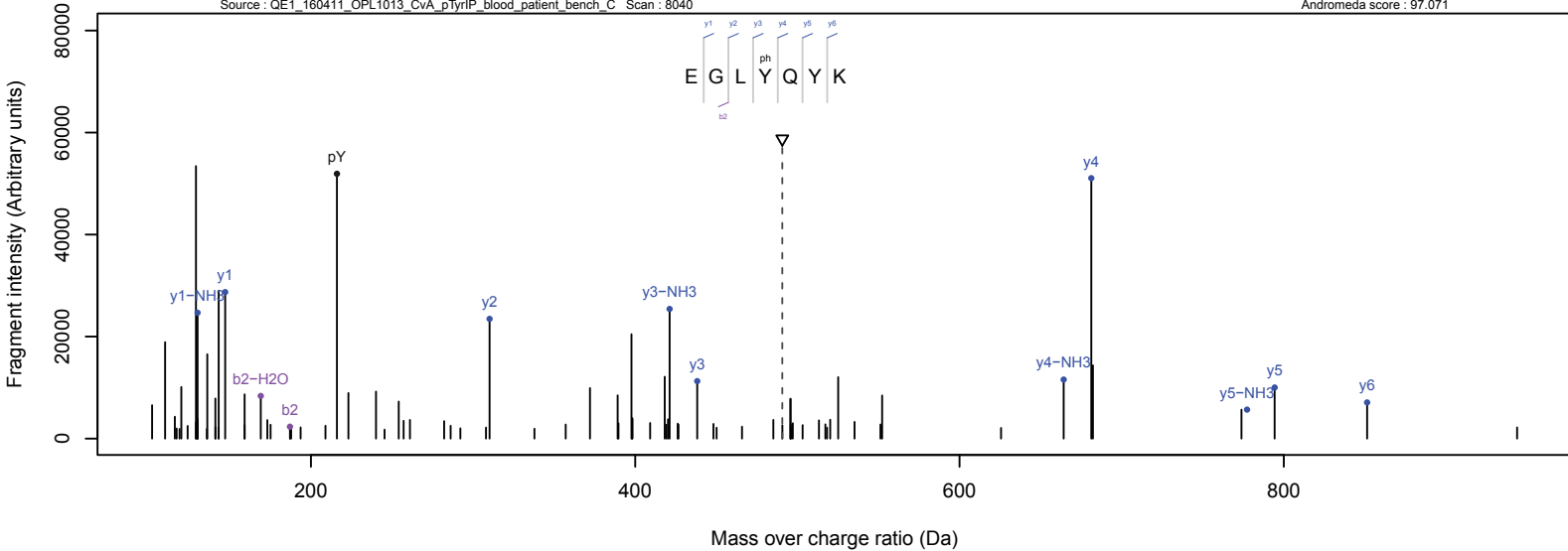
Andromeda score: 97.235



## EGLY(ph)QYK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 8040

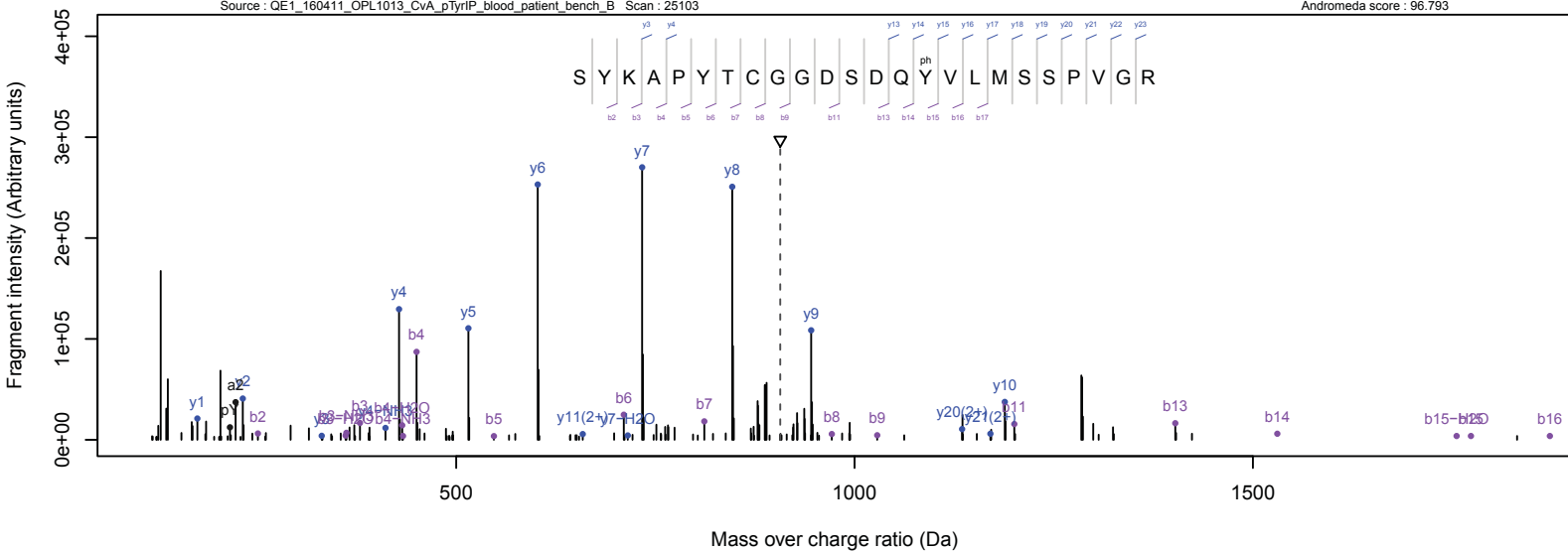
Andromeda score: 97.071



**SYKAPYTCGGDSDQY<sup>ph</sup>VLMSSPVGR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 25103

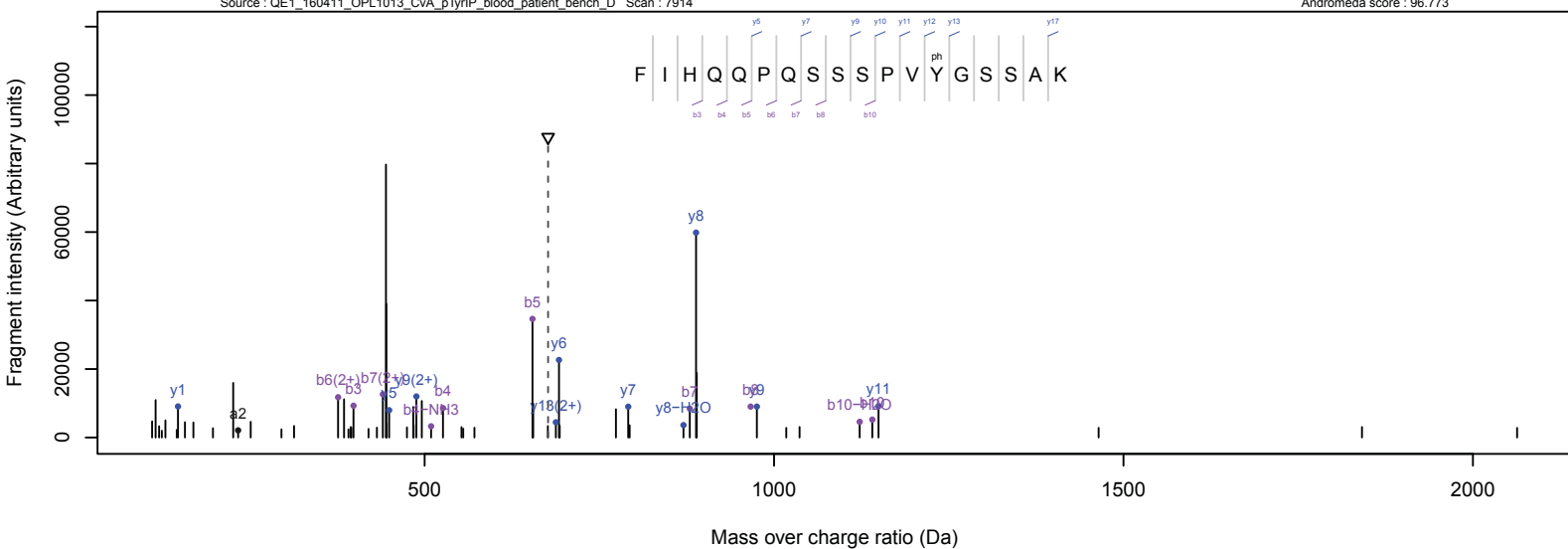
Andromeda score : 96.793



**FIHQQPQSSPVY<sup>ph</sup>GSSAK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7914

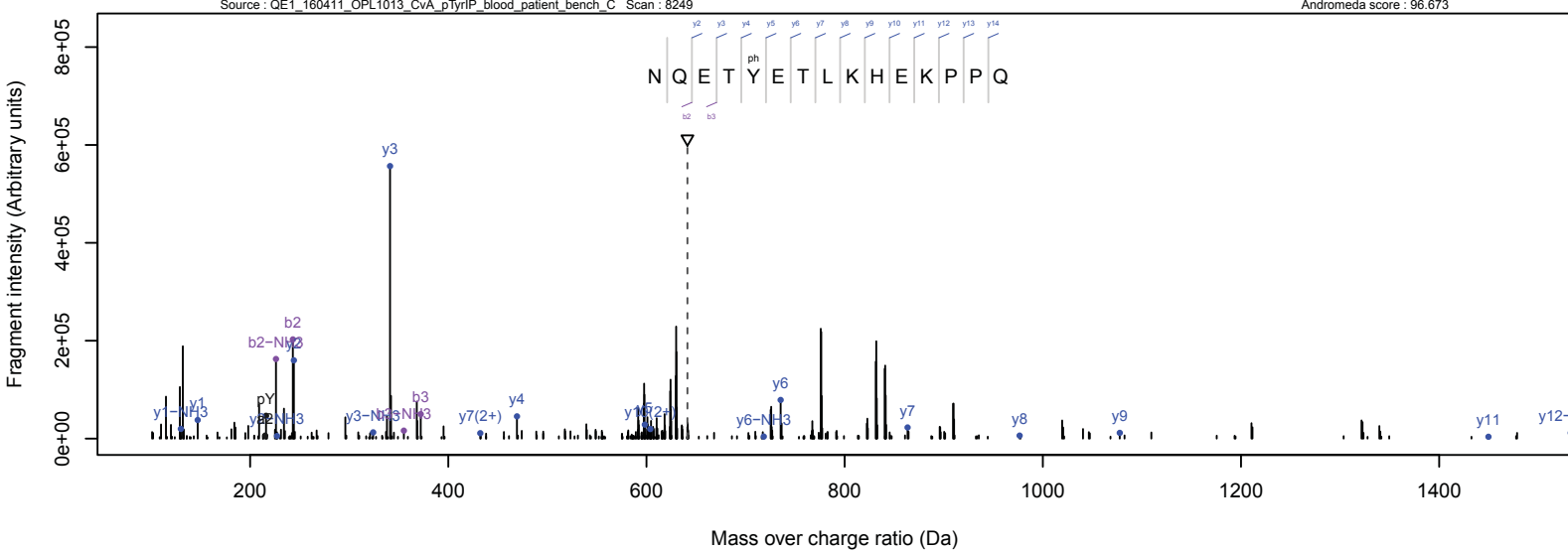
Andromeda score : 96.773

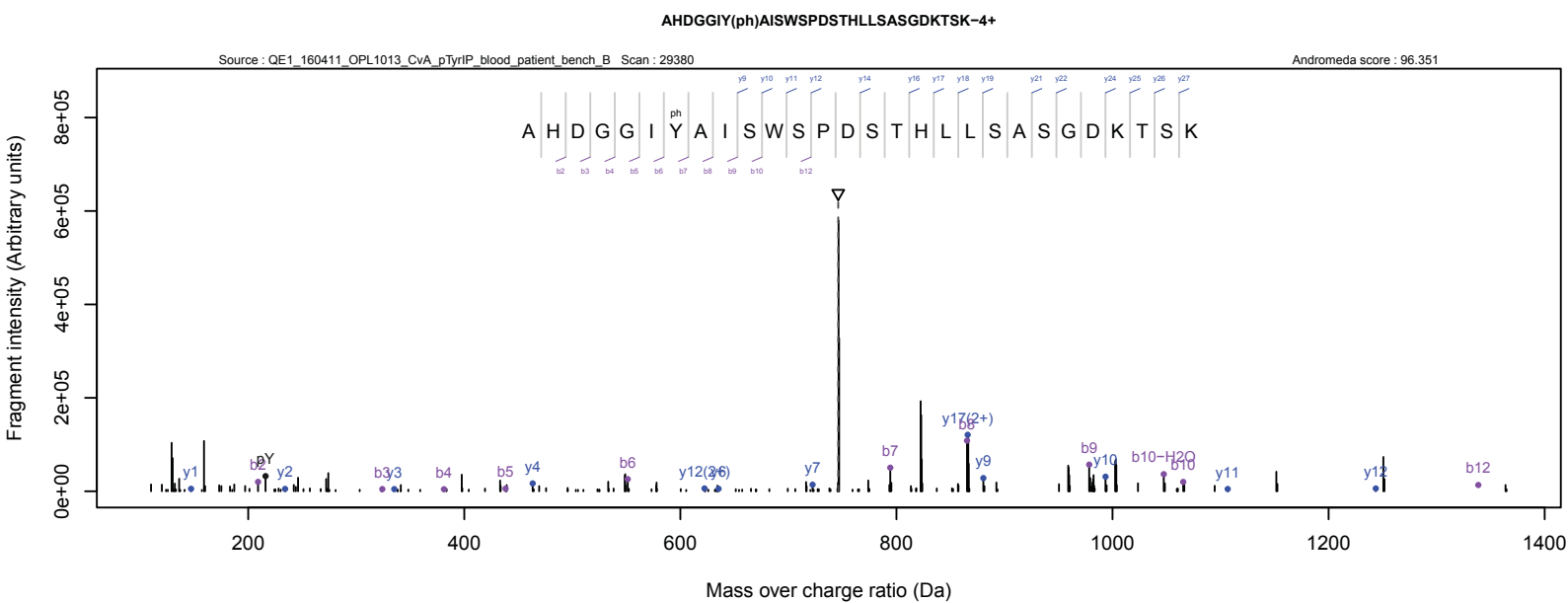
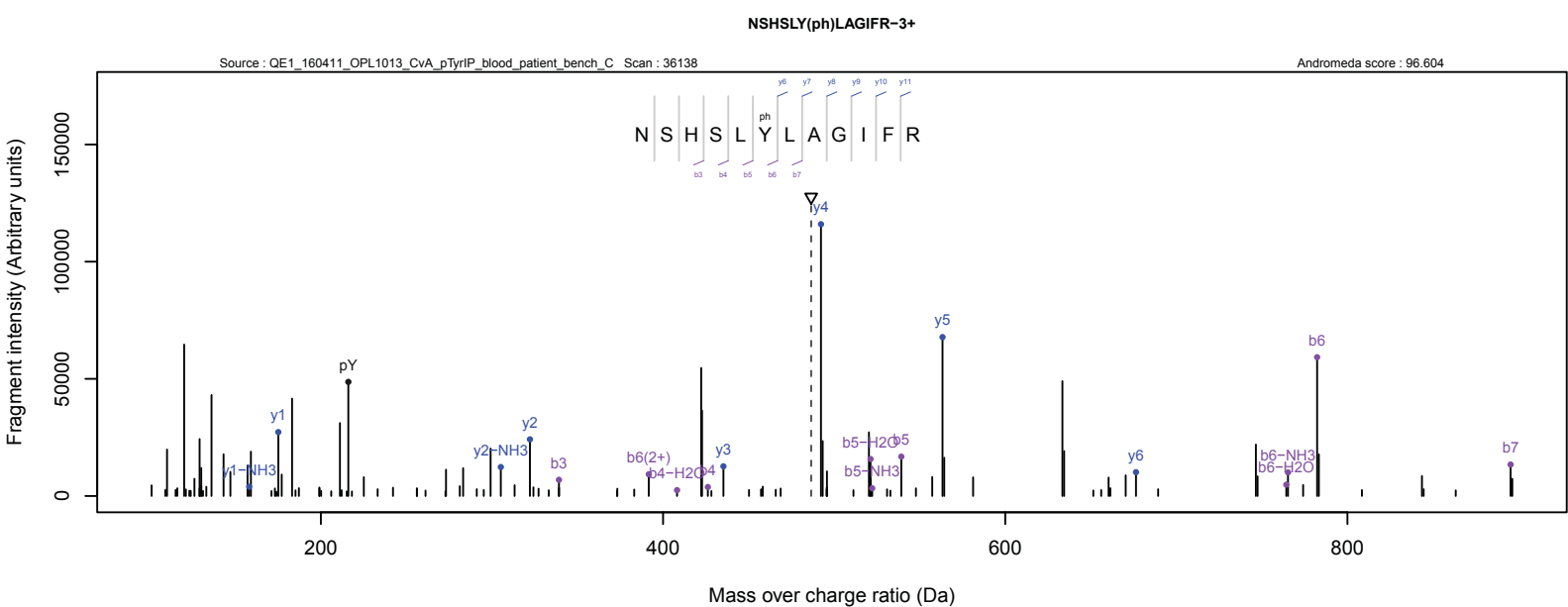
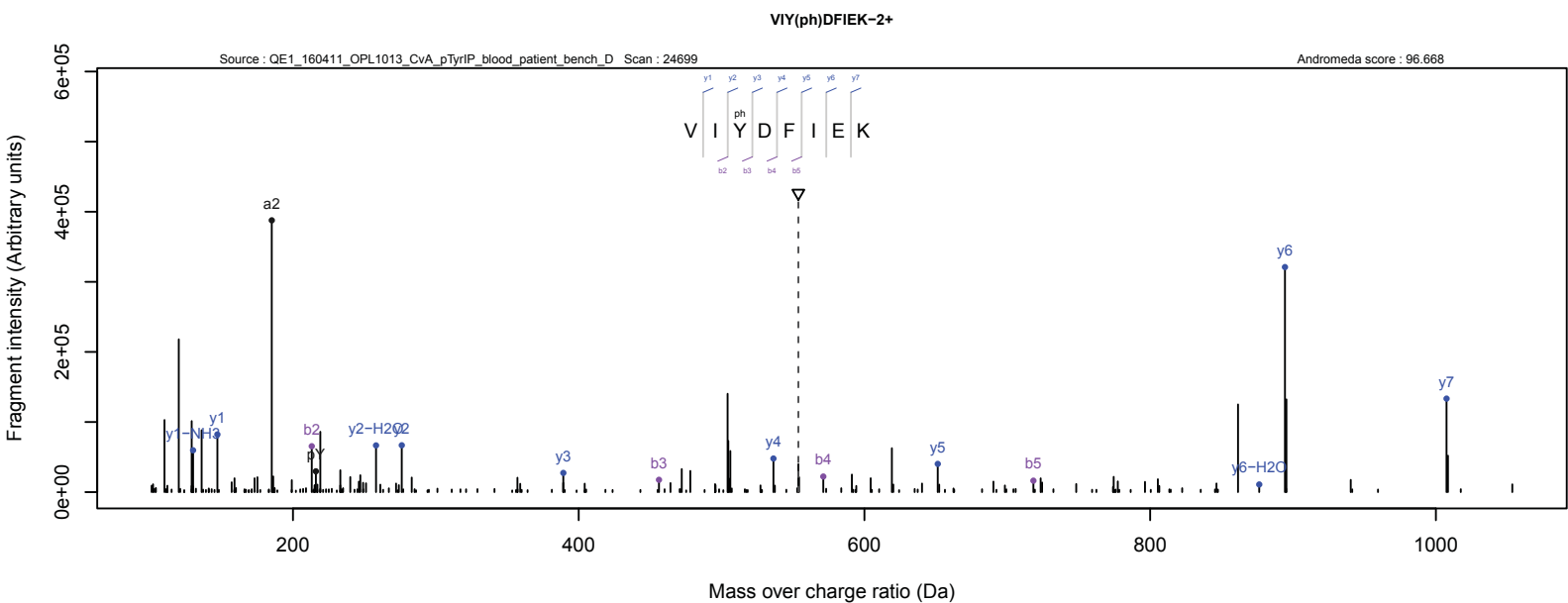


**NQETY<sup>ph</sup>ETLKHEKPPQ-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 8249

Andromeda score : 96.673



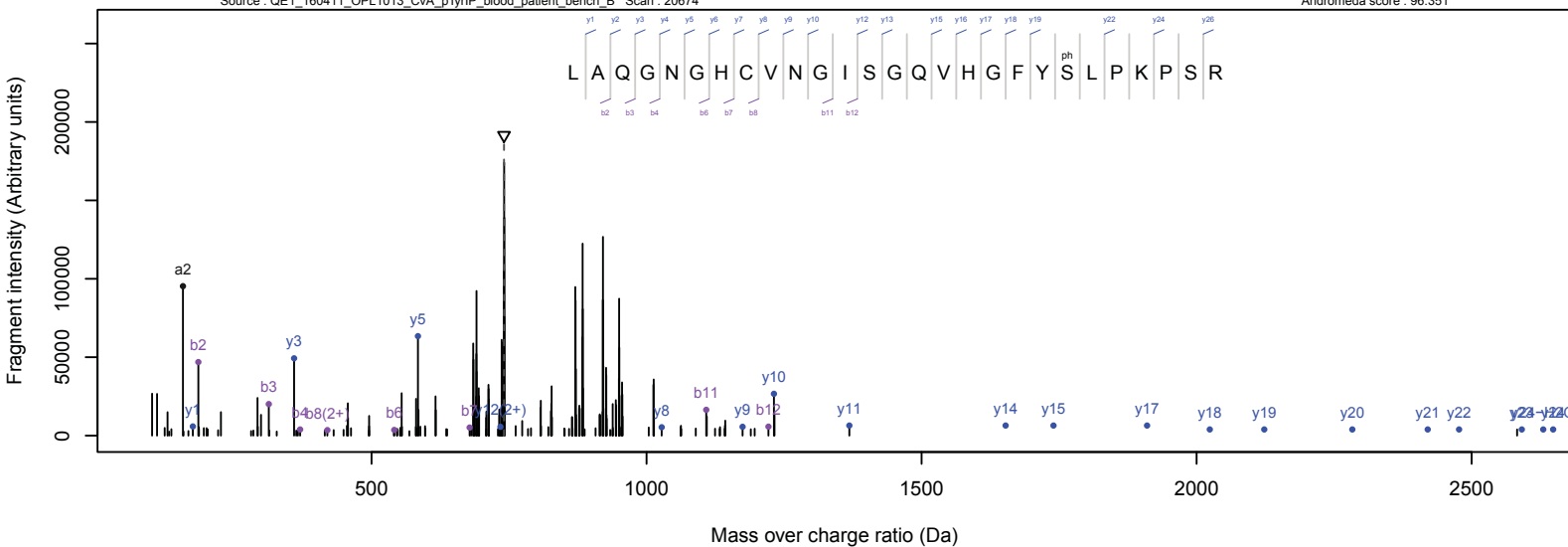




LAQNGHCVNGISGQVHGFYS(ph)LPKPSR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 20674

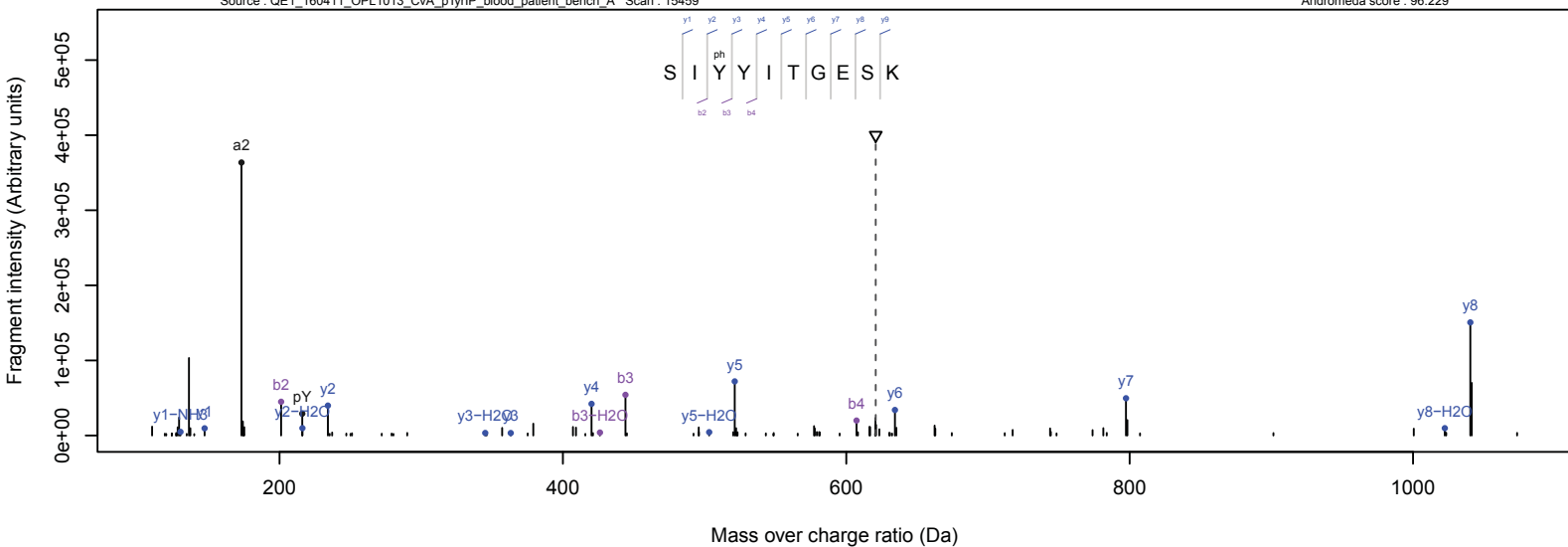
Andromeda score : 96.351



SIY(ph)YITGESK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 15459

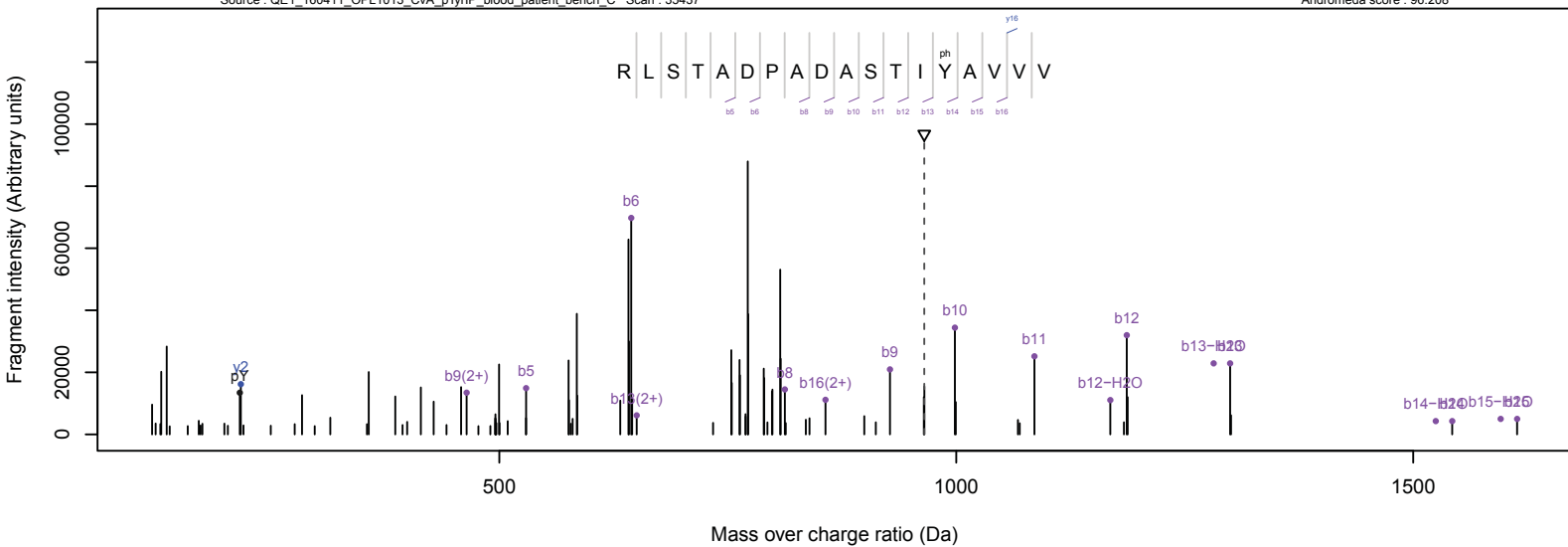
Andromeda score : 96.229



RLSTADPADASTIY(ph)AVVV-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 35437

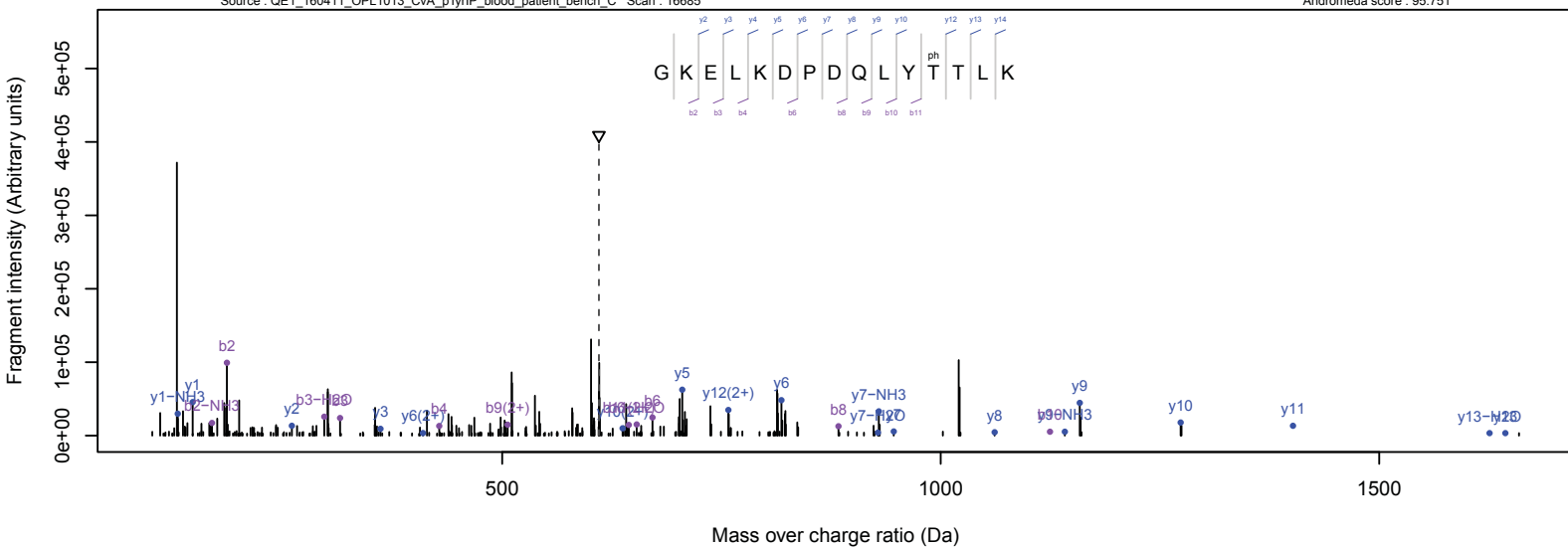
Andromeda score : 96.208



**GKELKDPDQLYT(ph)TLK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16685

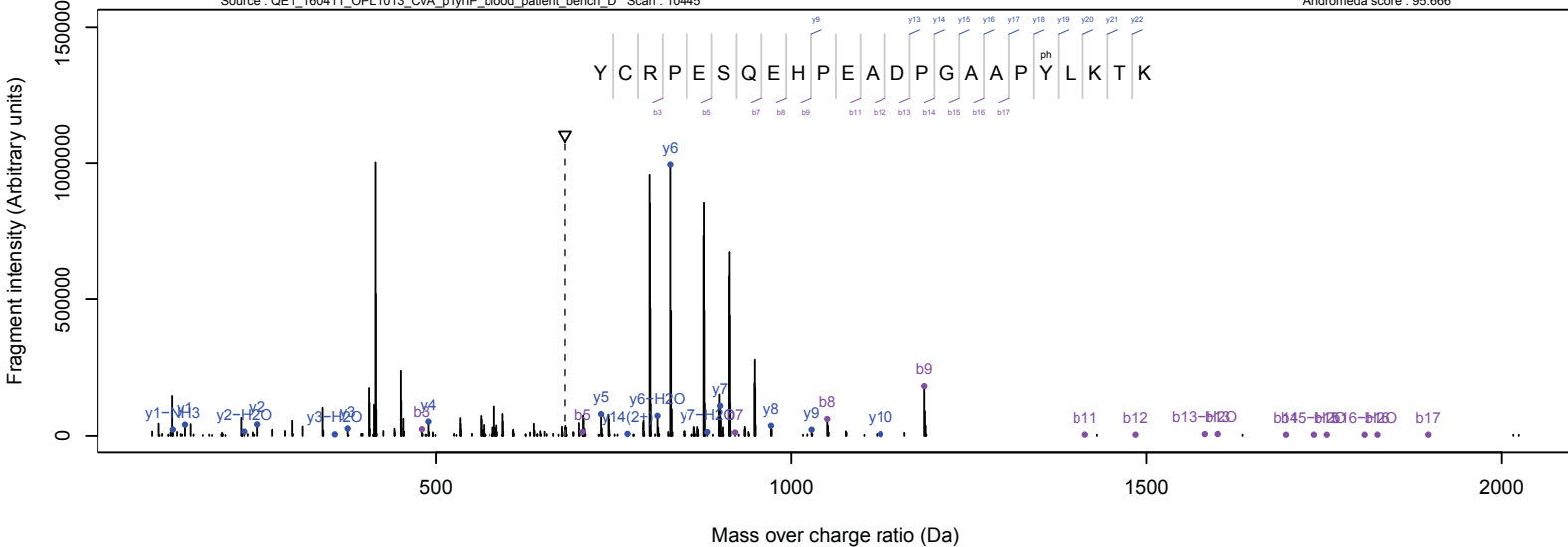
Andromeda score : 95.751



**YCRPESQEHPDPAAPY(ph)LKTK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 10445

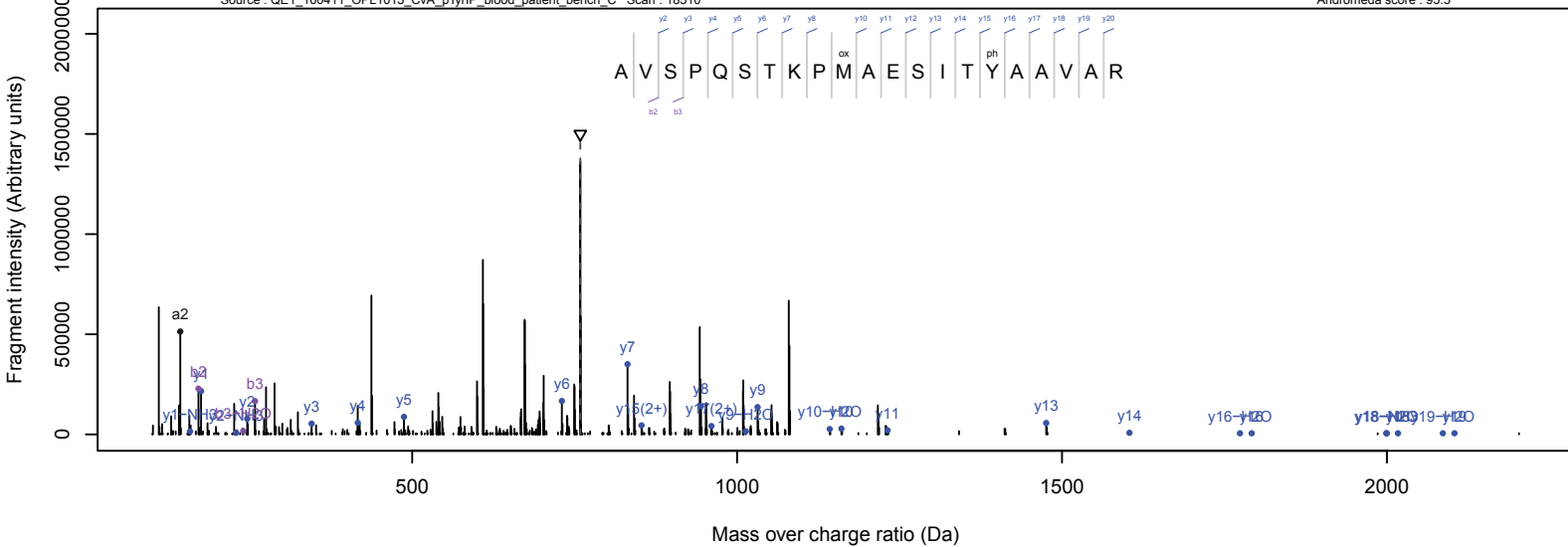
Andromeda score : 95.666



**AVSPQSTKPM(ox)AESITYA(AVAR)-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18510

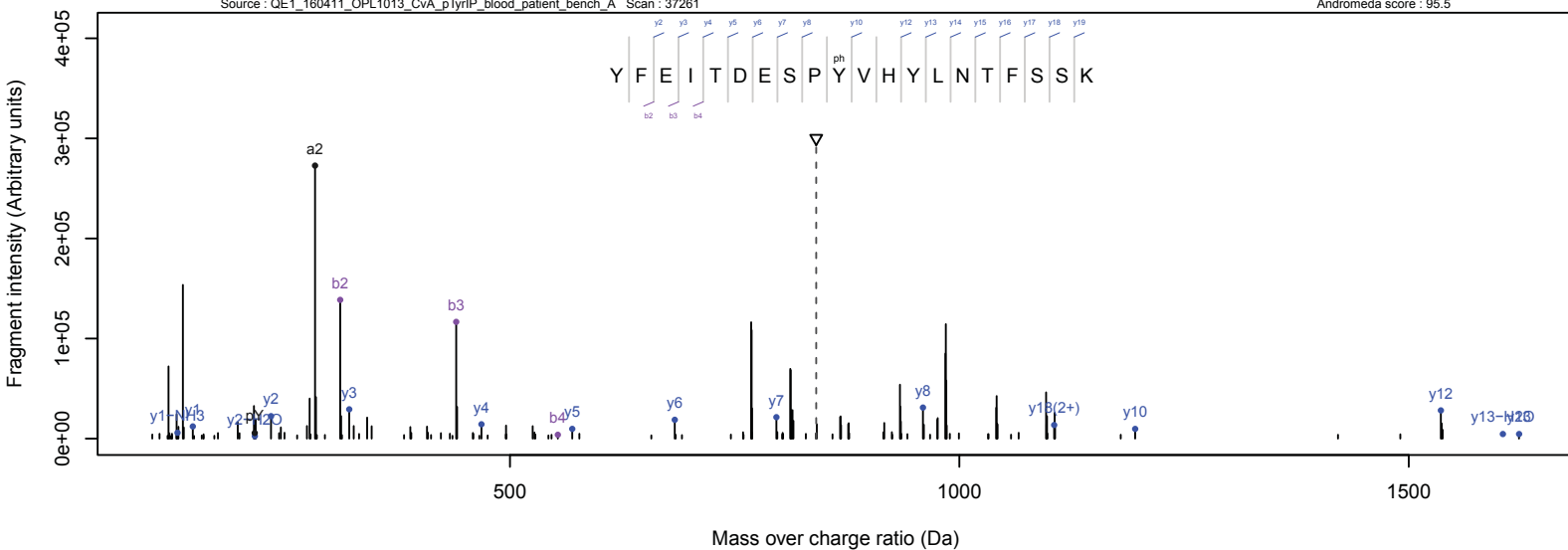
Andromeda score : 95.5



YFEITDESPY<sup>ph</sup>VHYLNTFSSK-3+

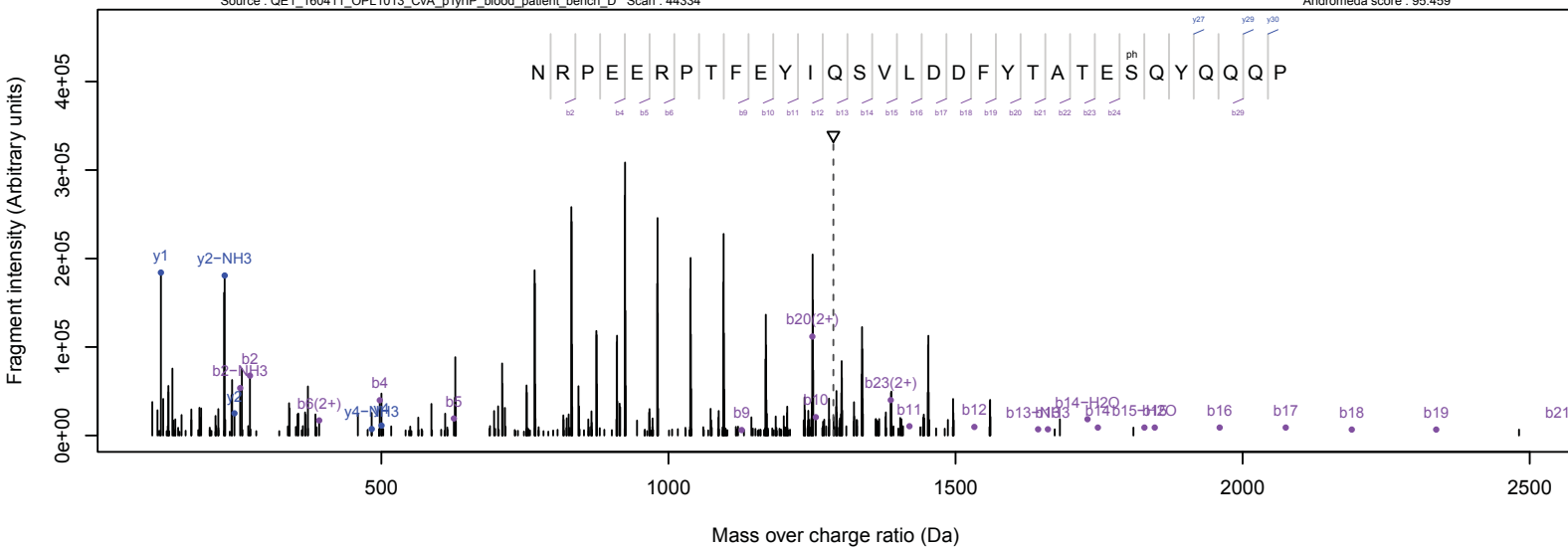
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 37261

Andromeda score: 95.5

NRPEERPTFEYIQSVLDDFYTATES<sup>ph</sup>QYQQP-3+

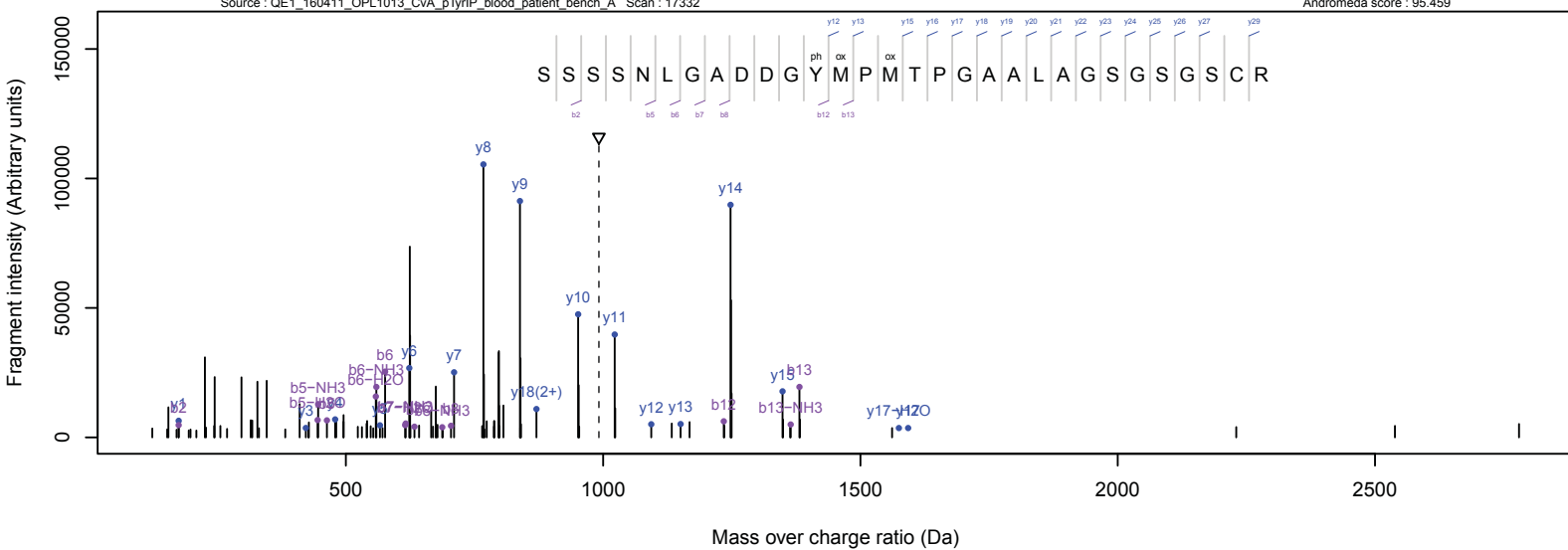
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 44334

Andromeda score: 95.459

SSSSNLGADDGY<sup>ph</sup>M<sup>(ox)</sup>PM<sup>(ox)</sup>TPGAALAGSGSGSCR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 17332

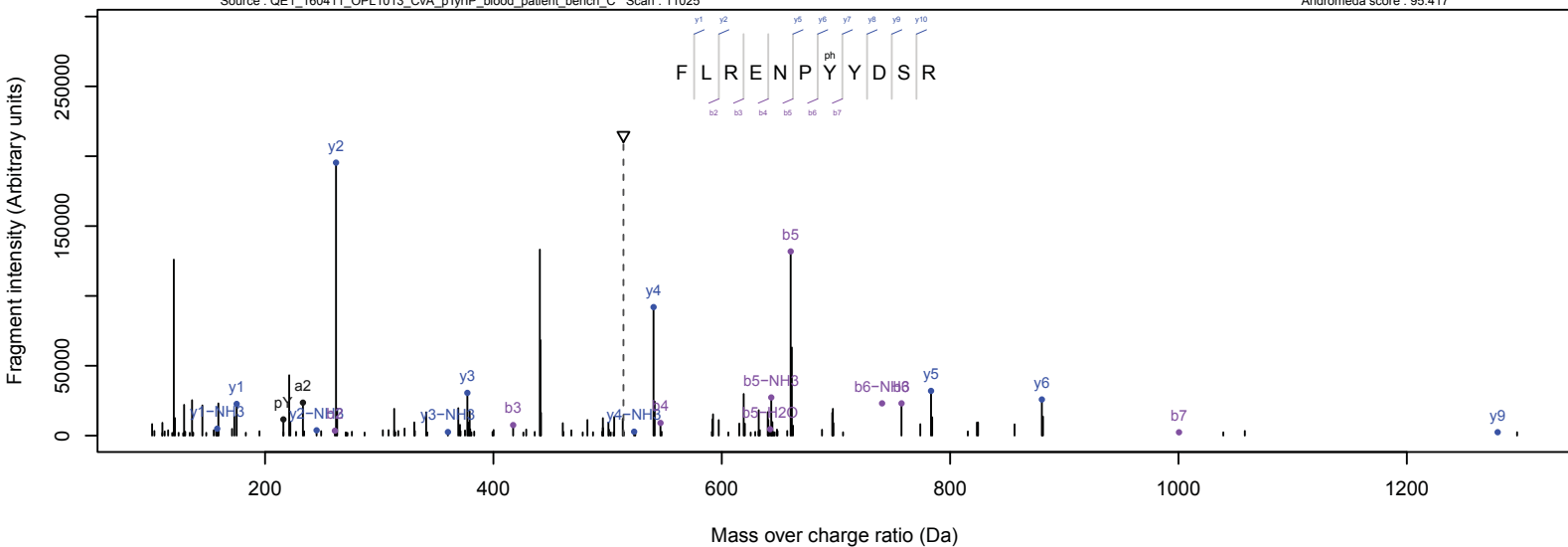
Andromeda score: 95.459



**FLRENPY(ph)YDSR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 11025

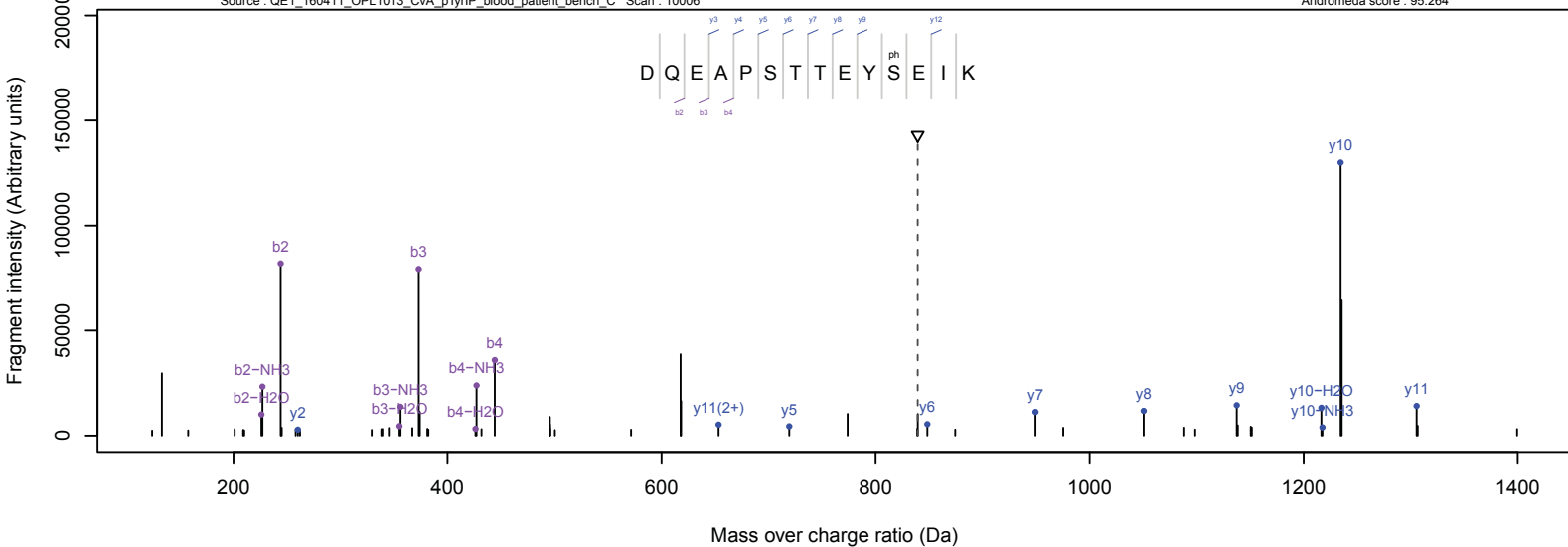
Andromeda score: 95.417



**DQEAPSTTEYS(ph)EIK-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 10006

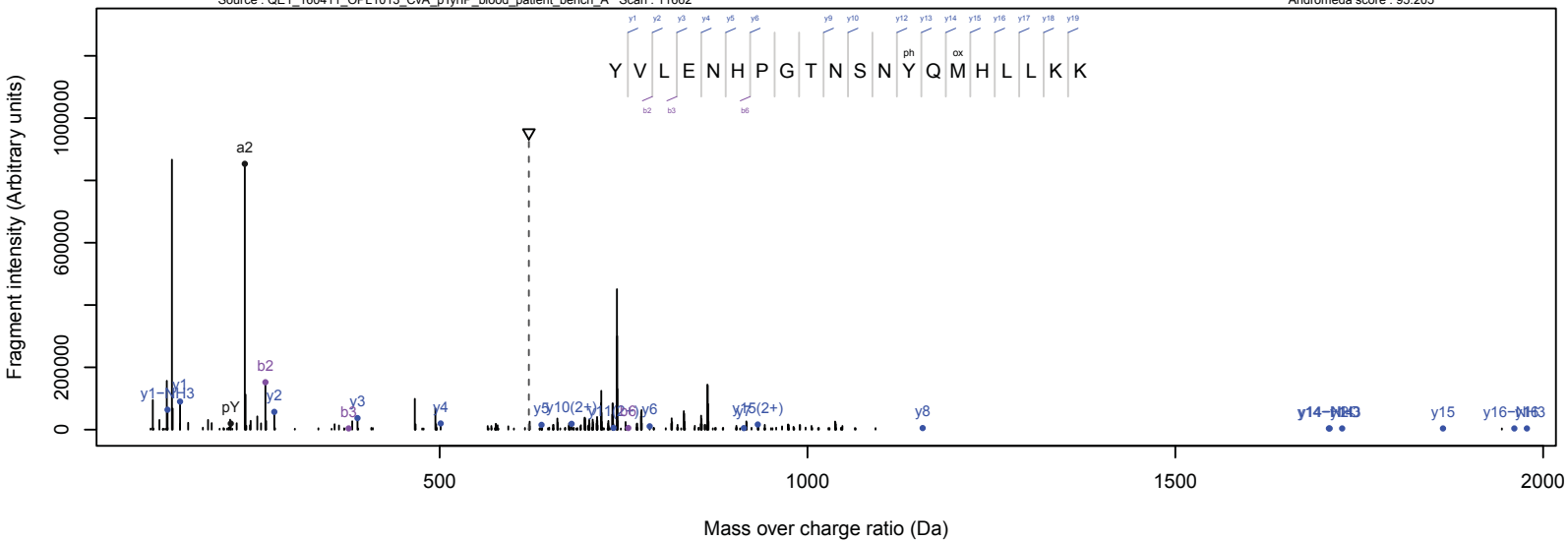
Andromeda score: 95.264



**YVLENHPGTSNY(ph)QM(ox)HLLKK-4+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 11662

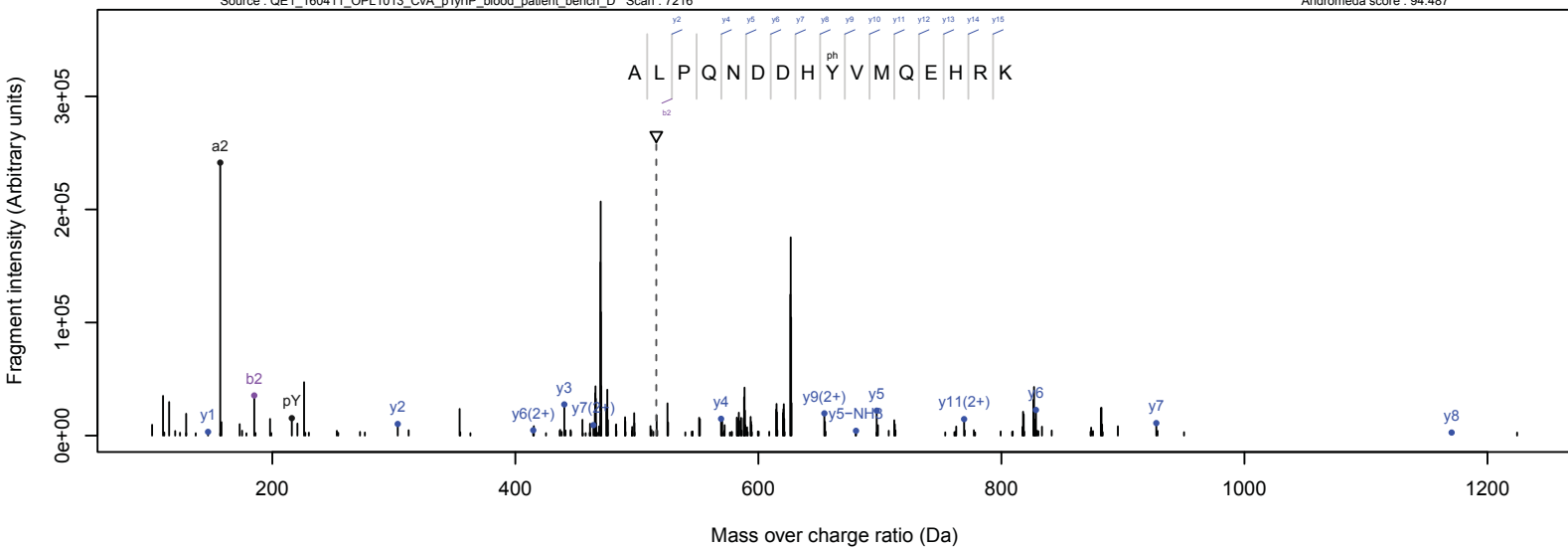
Andromeda score: 95.205



ALPQNDDHY(ph)VMQEHK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7216

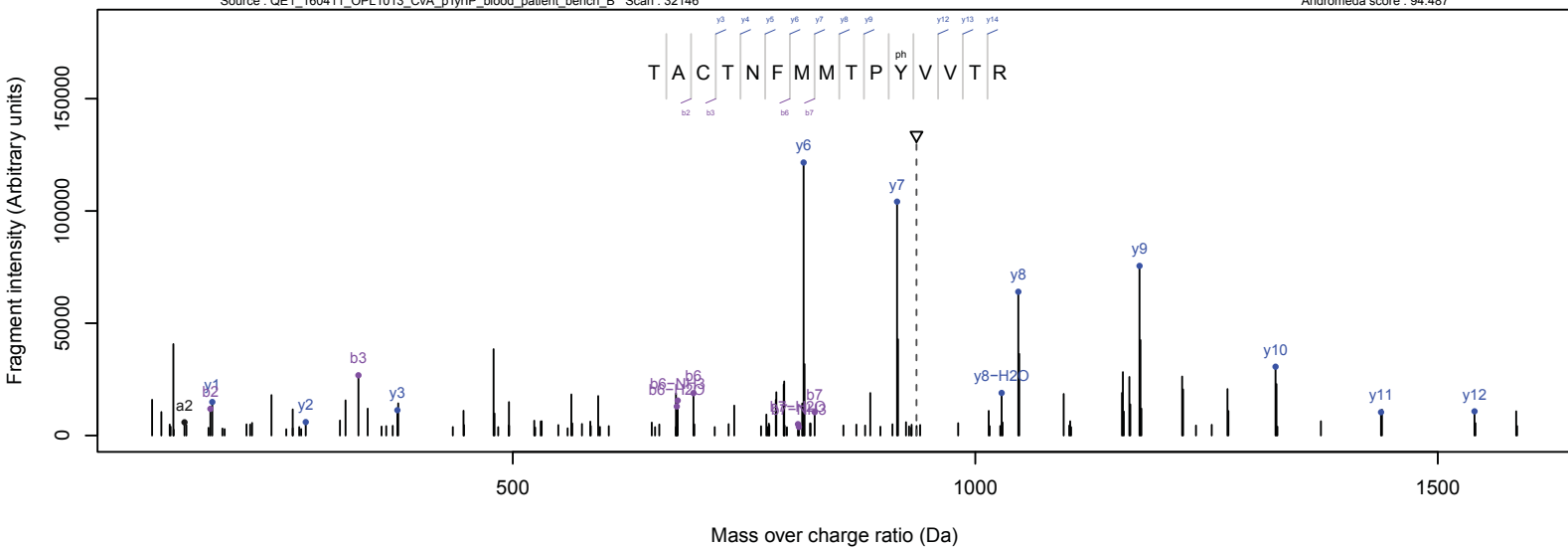
Andromeda score : 94.487



TACTNFMMPY(ph)VVTR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 32146

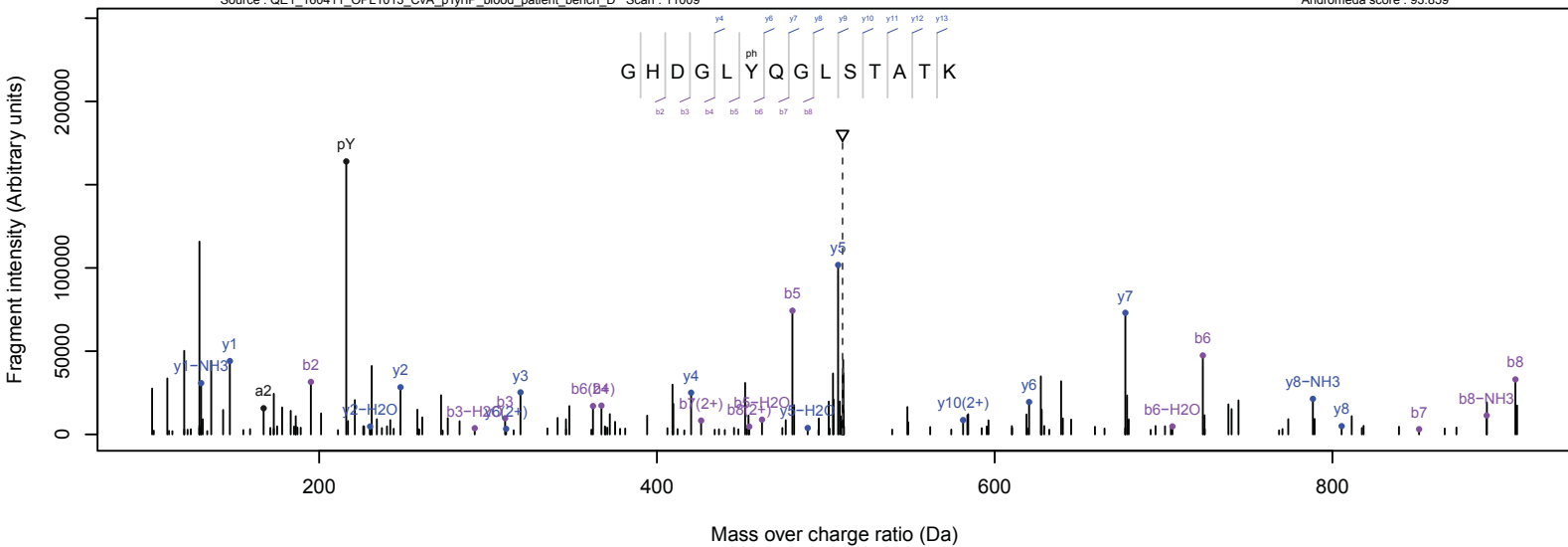
Andromeda score : 94.487



GHGGLY(ph)QLSTATK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11009

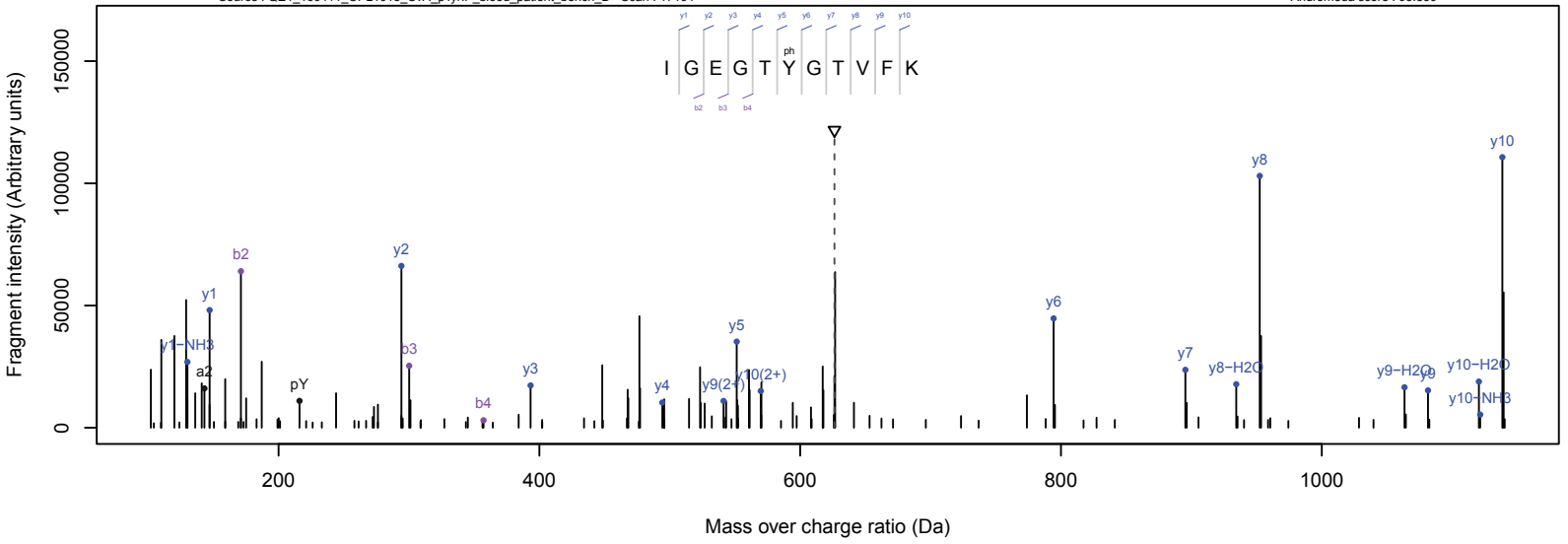
Andromeda score : 93.859



**IGEGTY(ph)GTVFK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17164

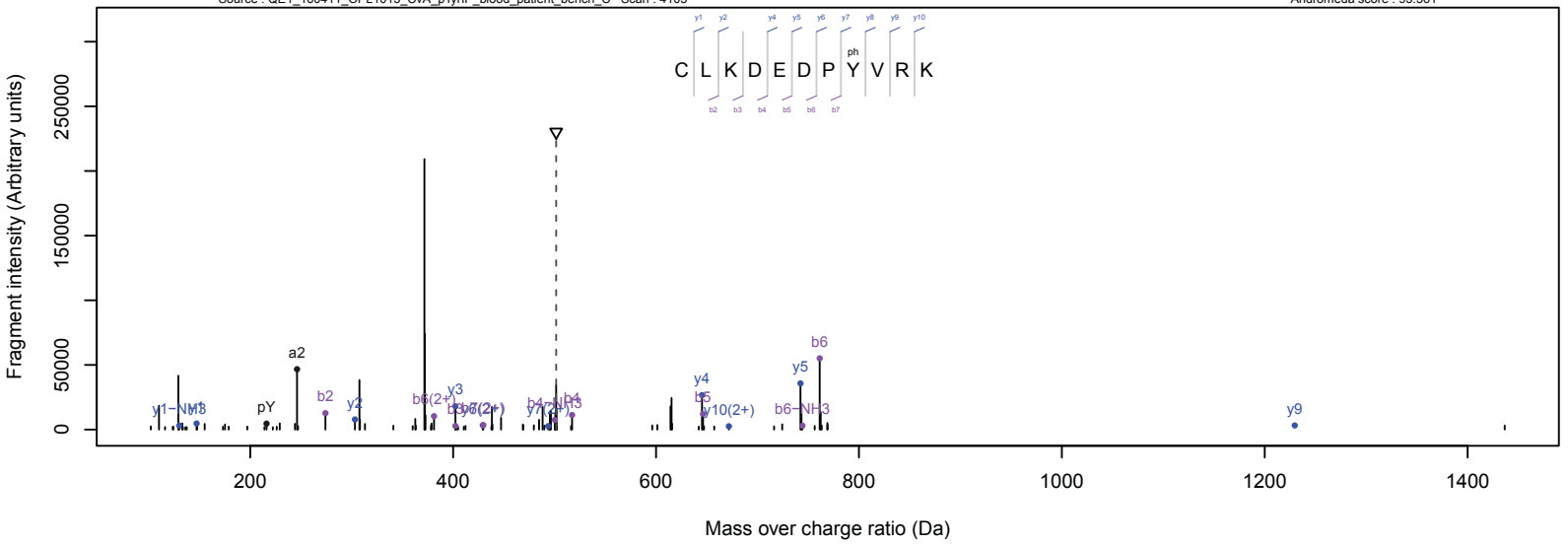
Andromeda score : 93.839



**CLKDEDPY(ph)VRK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 4105

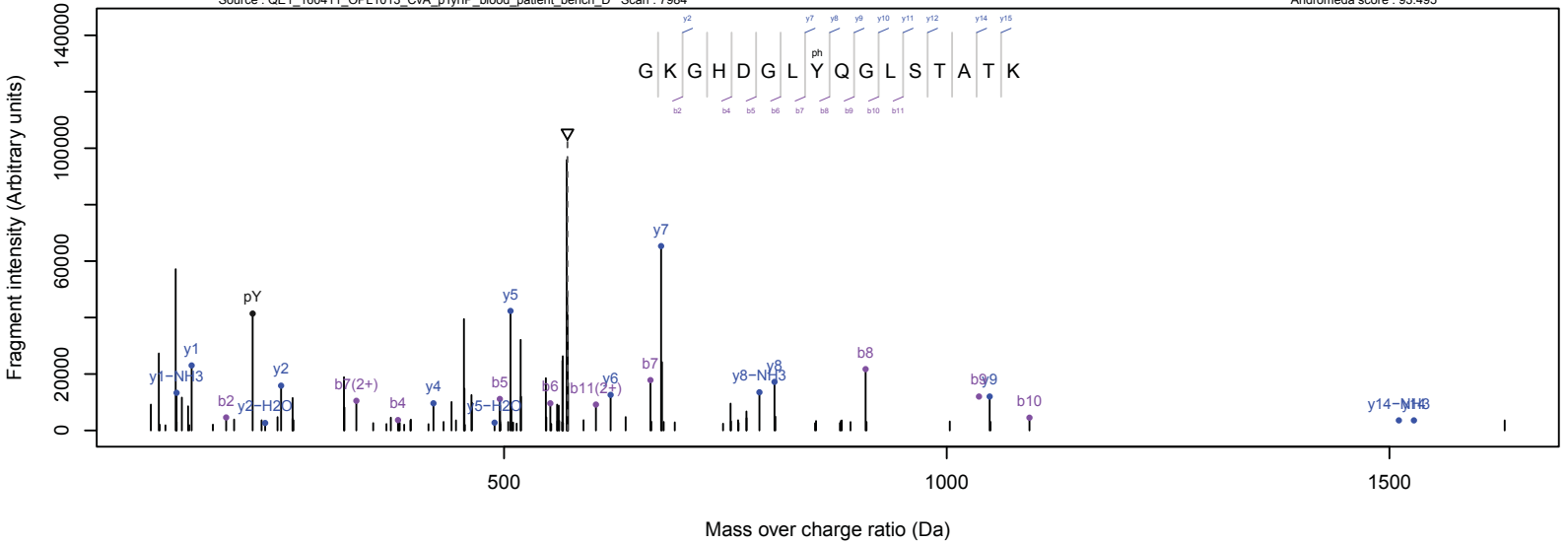
Andromeda score : 93.561



**GKGHDGLY(ph)QLSTATK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7984

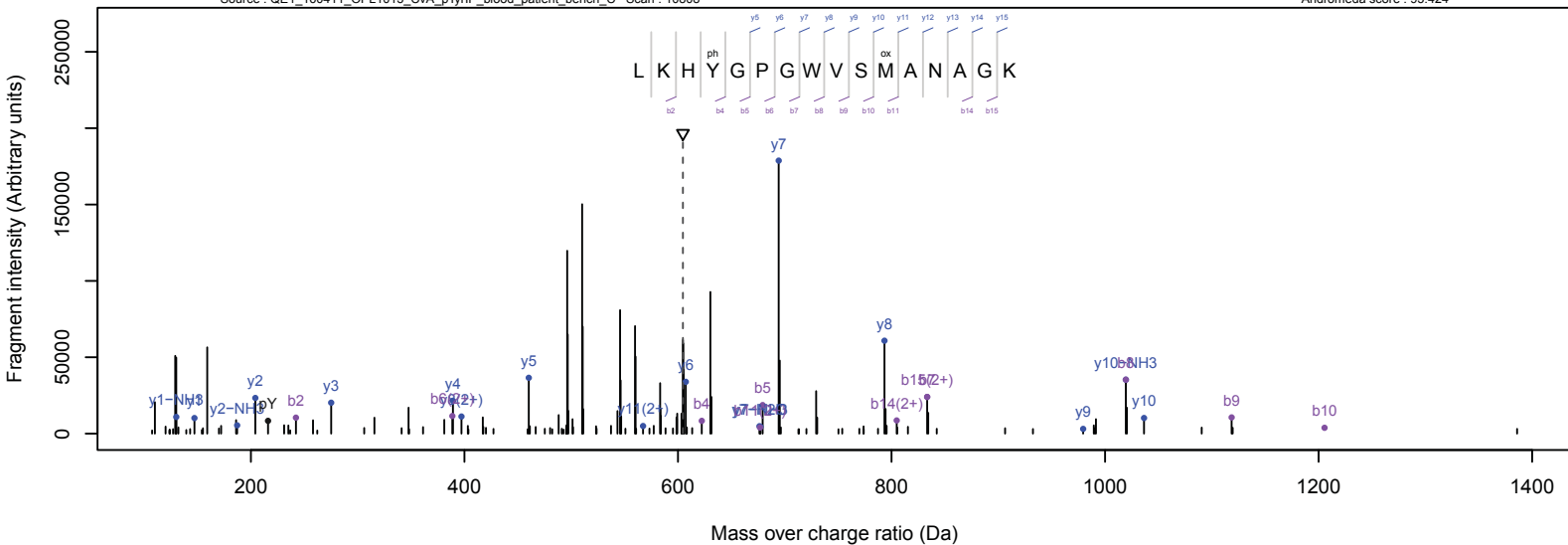
Andromeda score : 93.495



LKHY(ph)GPGWVSM(ox)ANAGK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 10808

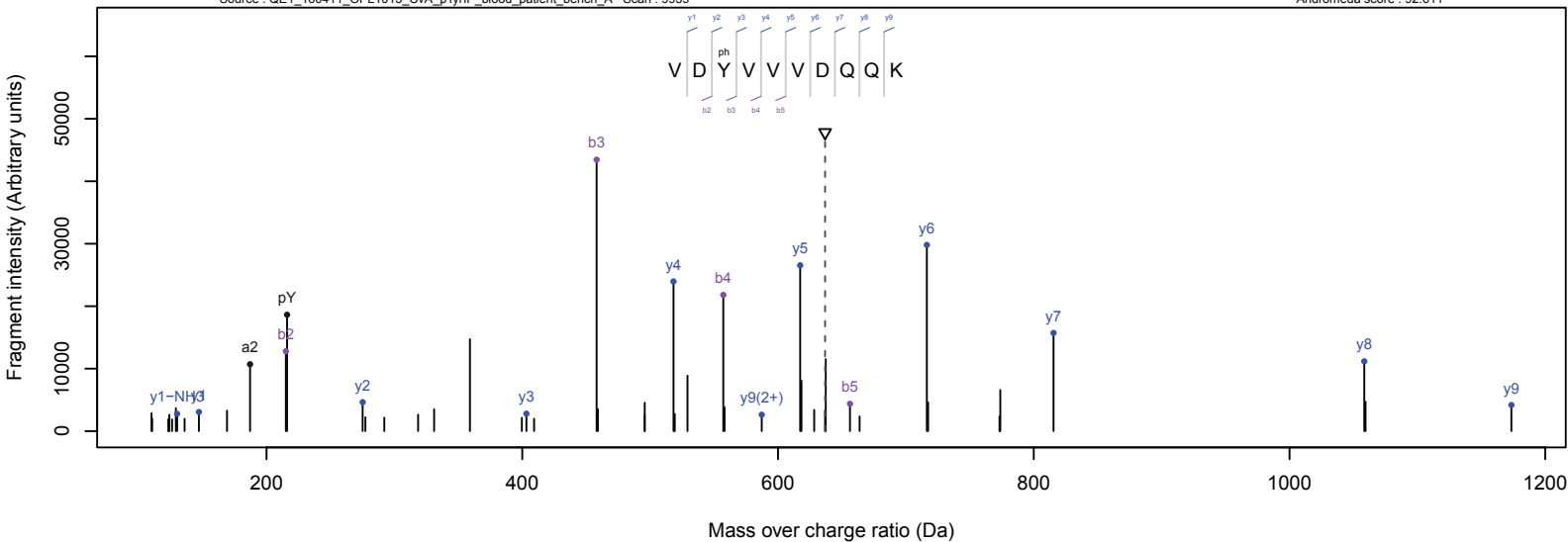
Andromeda score : 93.424



VDY(ph)VVDQK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 9939

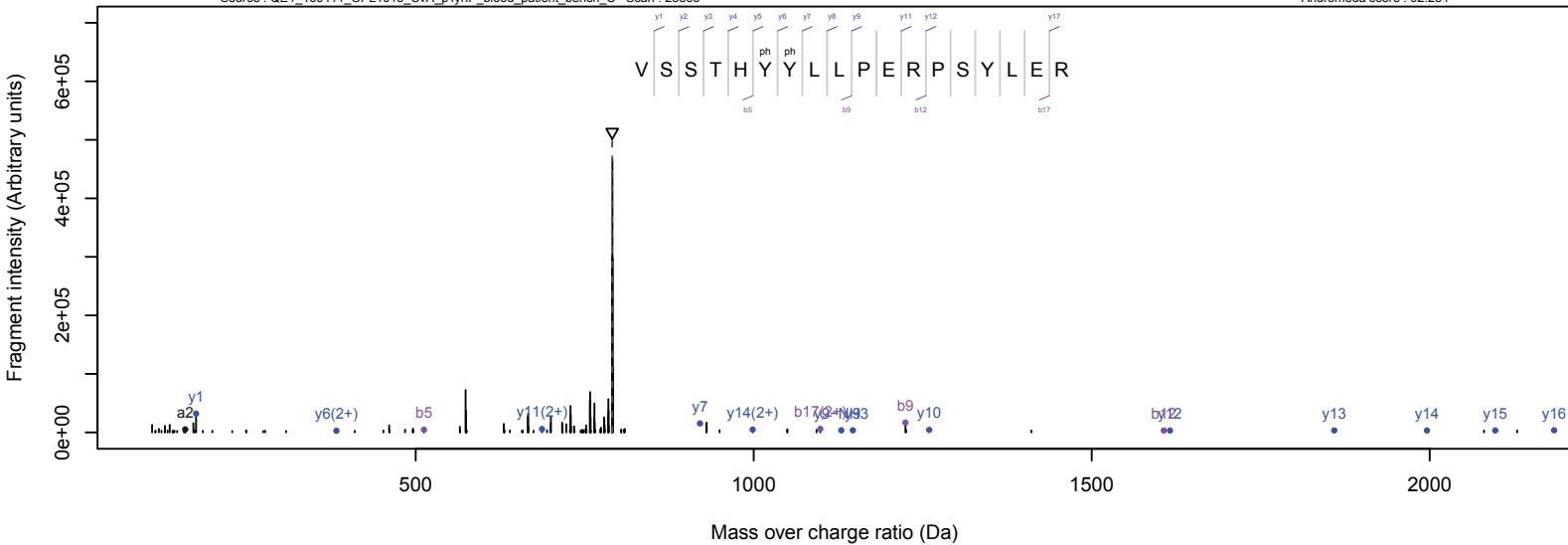
Andromeda score : 92.611



VSSTHY(ph)Y(ph)LLPERPSYLER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28863

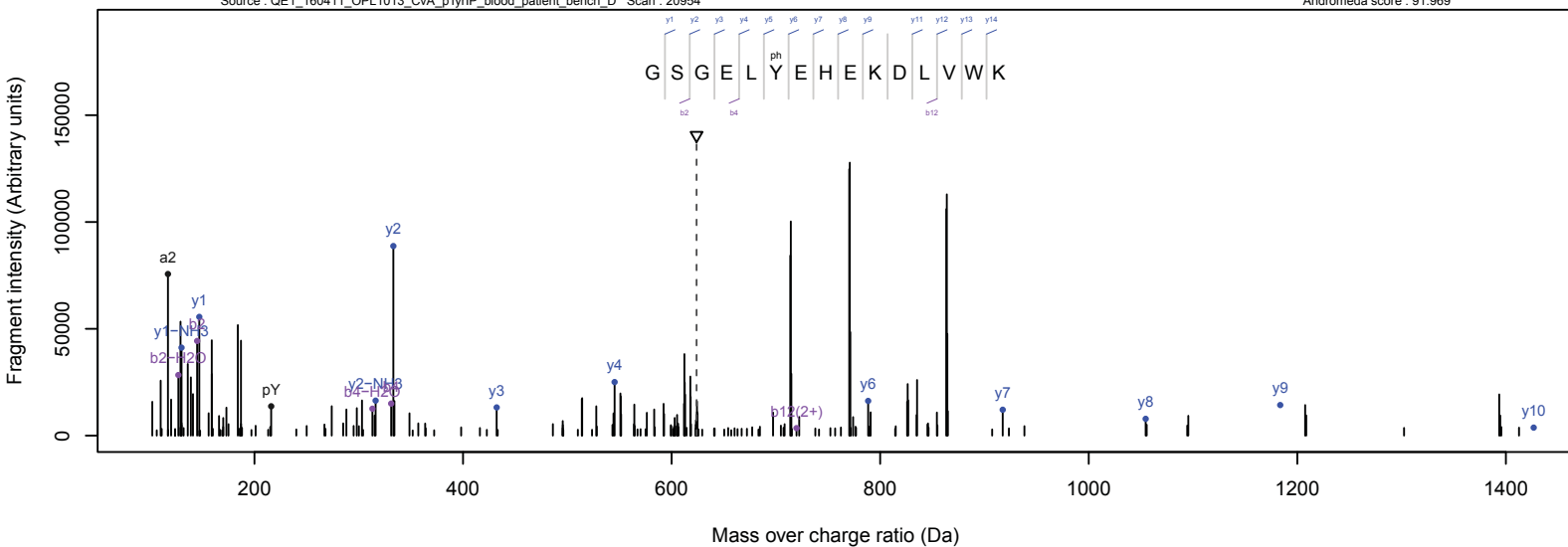
Andromeda score : 92.264



**GSGELY(ph)EHEKDLVWK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 20954

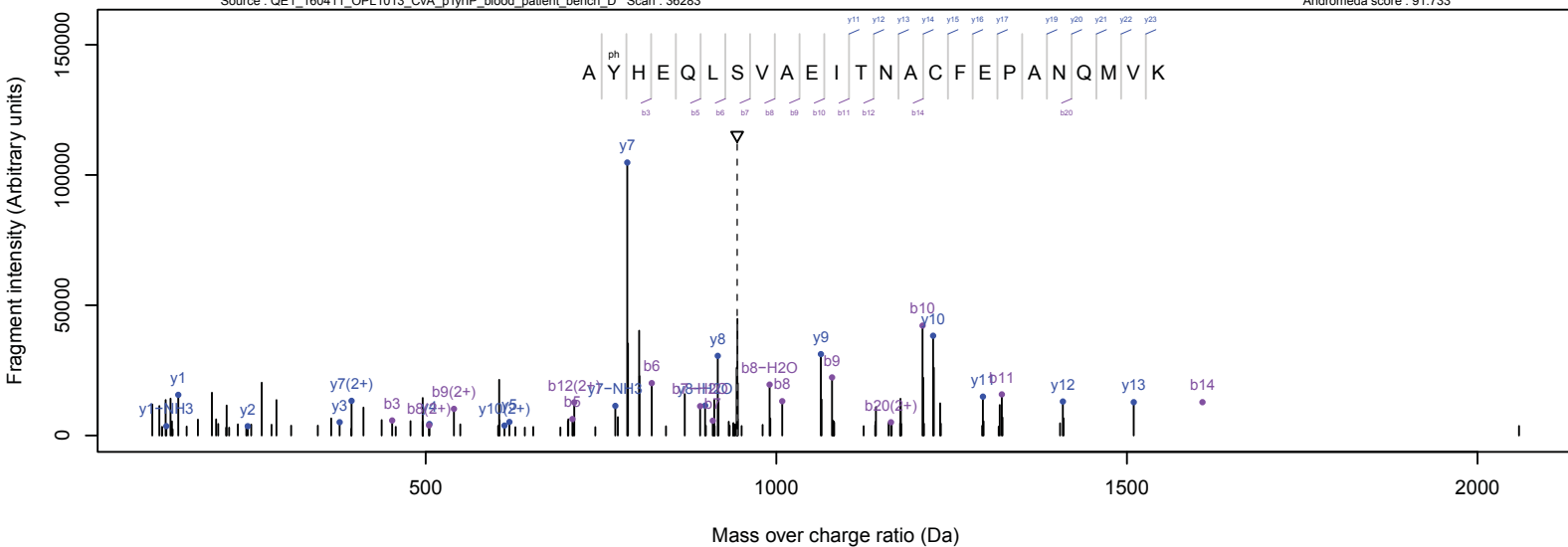
Andromeda score : 91.969



**AY(ph)HEQLSVAEITNACFEPANQMVK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 36283

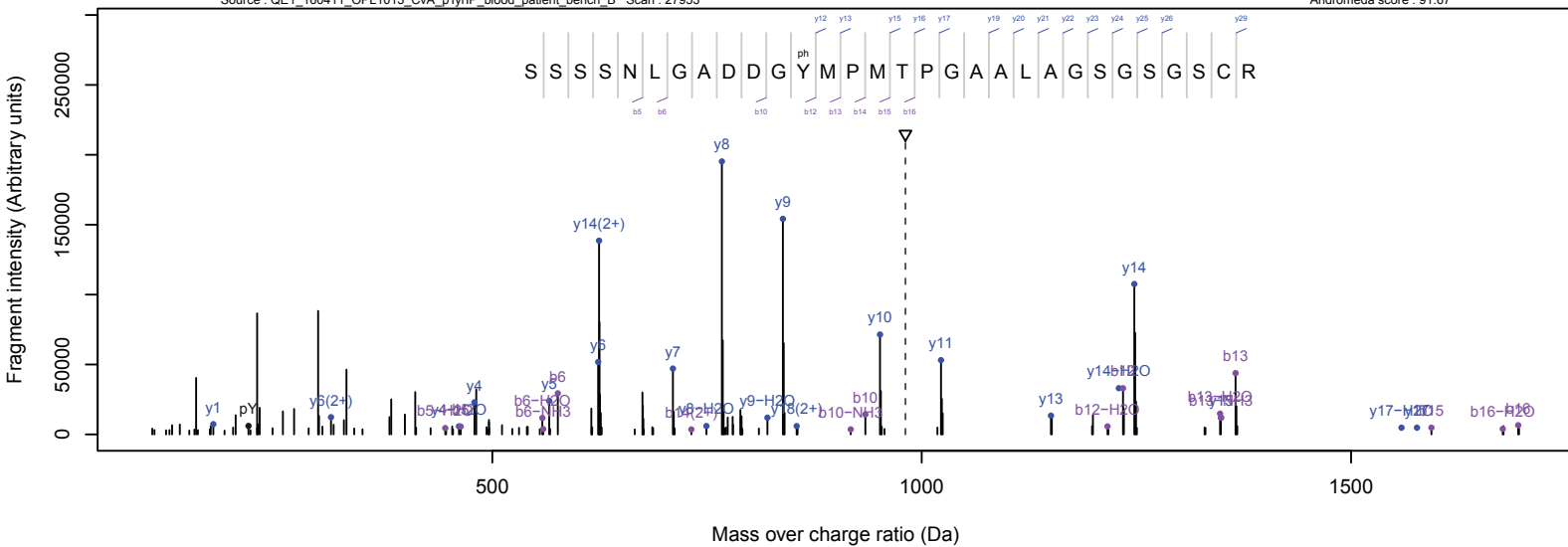
Andromeda score : 91.733



**SSSNLGADDG(ph)MPMPGAALAGSGSGSCR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 27953

Andromeda score : 91.67

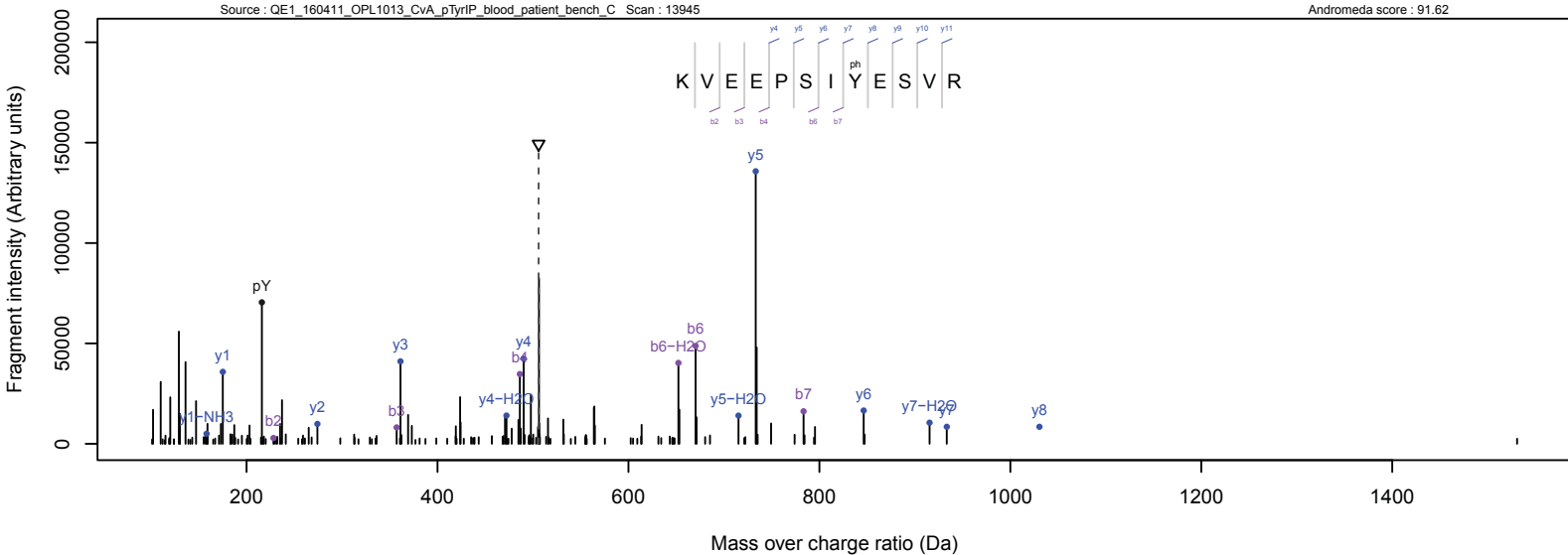




**KVEEPSIY(ph)ESVR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 13945

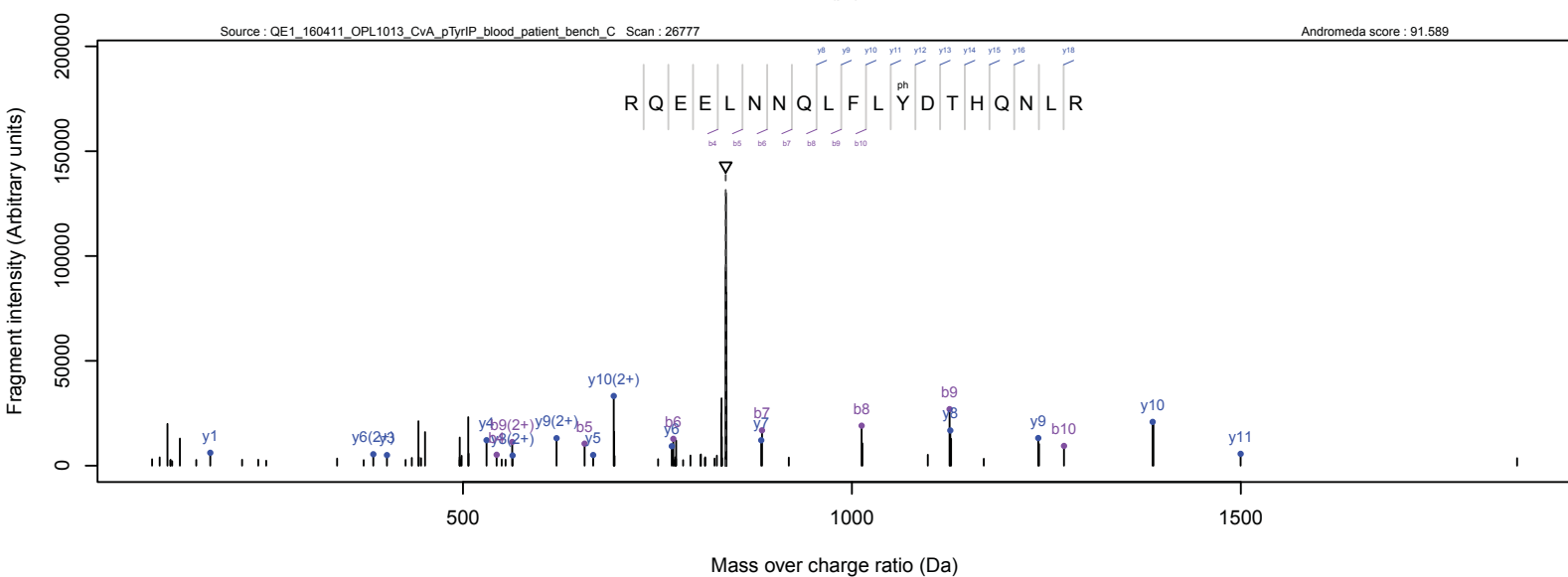
Andromeda score: 91.62



**RQEELNNQLFLY(ph)DTHQNL R-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 26777

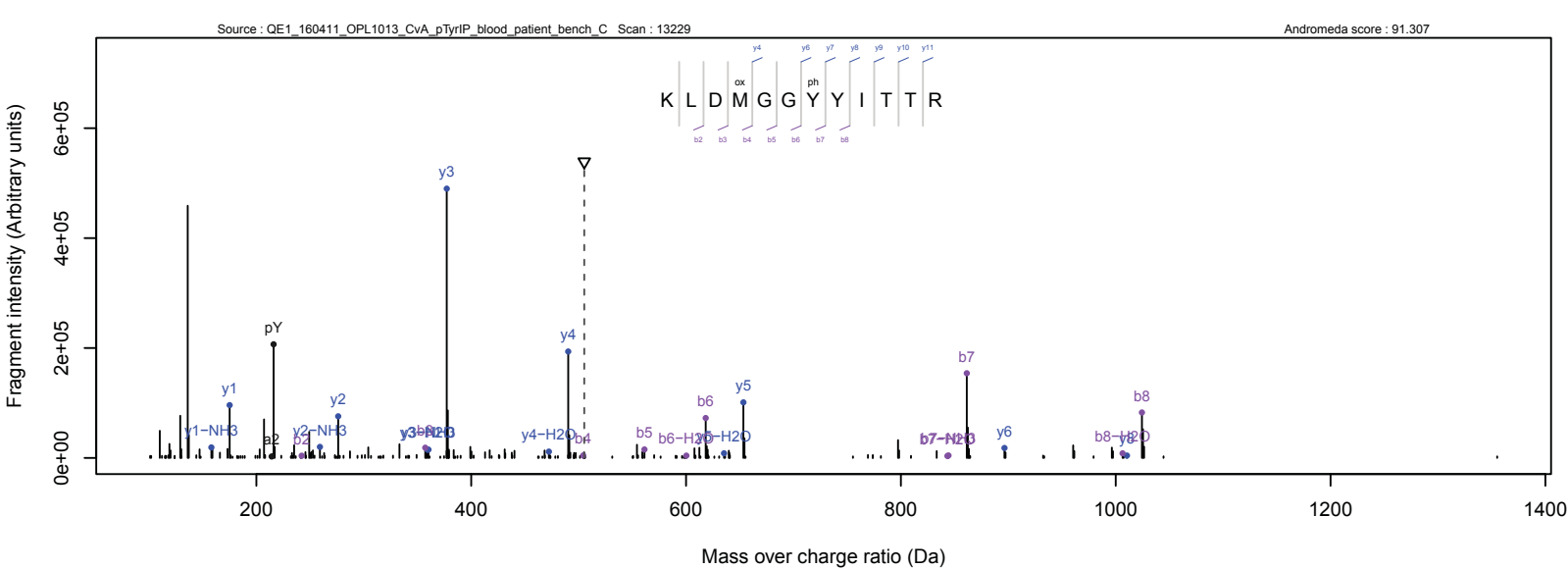
Andromeda score: 91.589



**KLDMGGY(ph)YITTR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 13229

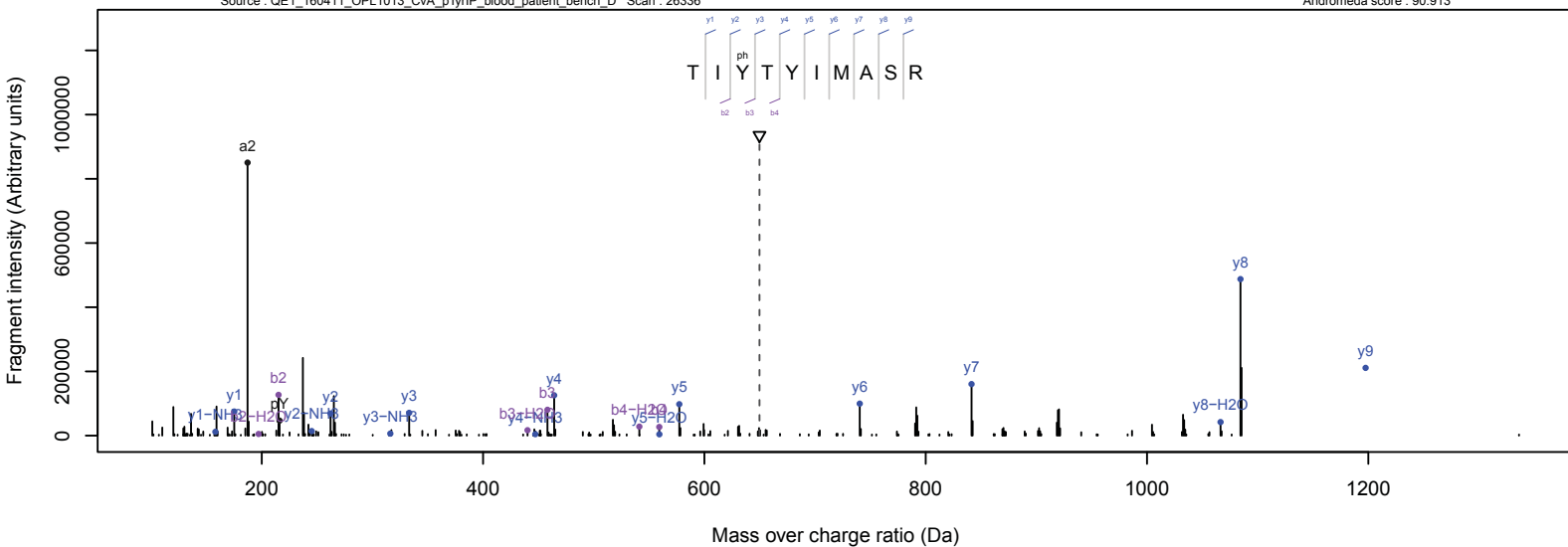
Andromeda score: 91.307



## TIY(ph)TYIMASR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 26336

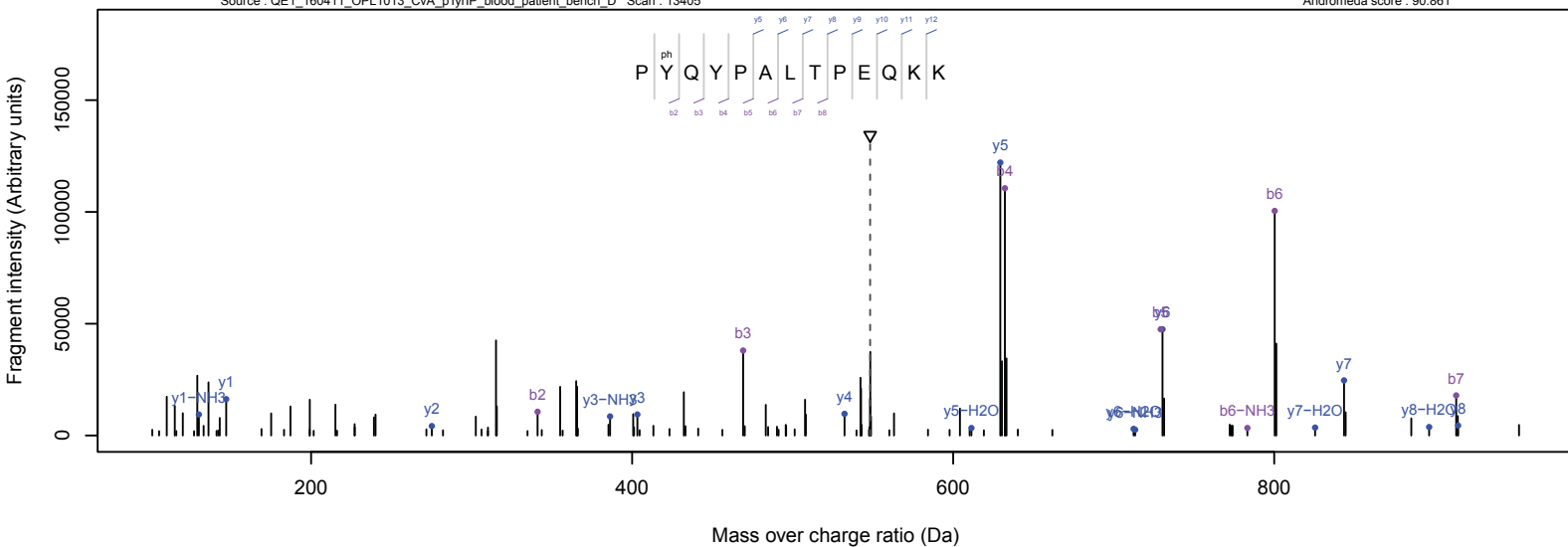
Andromeda score: 90.913



## PY(ph)QYPALTPQKK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 13405

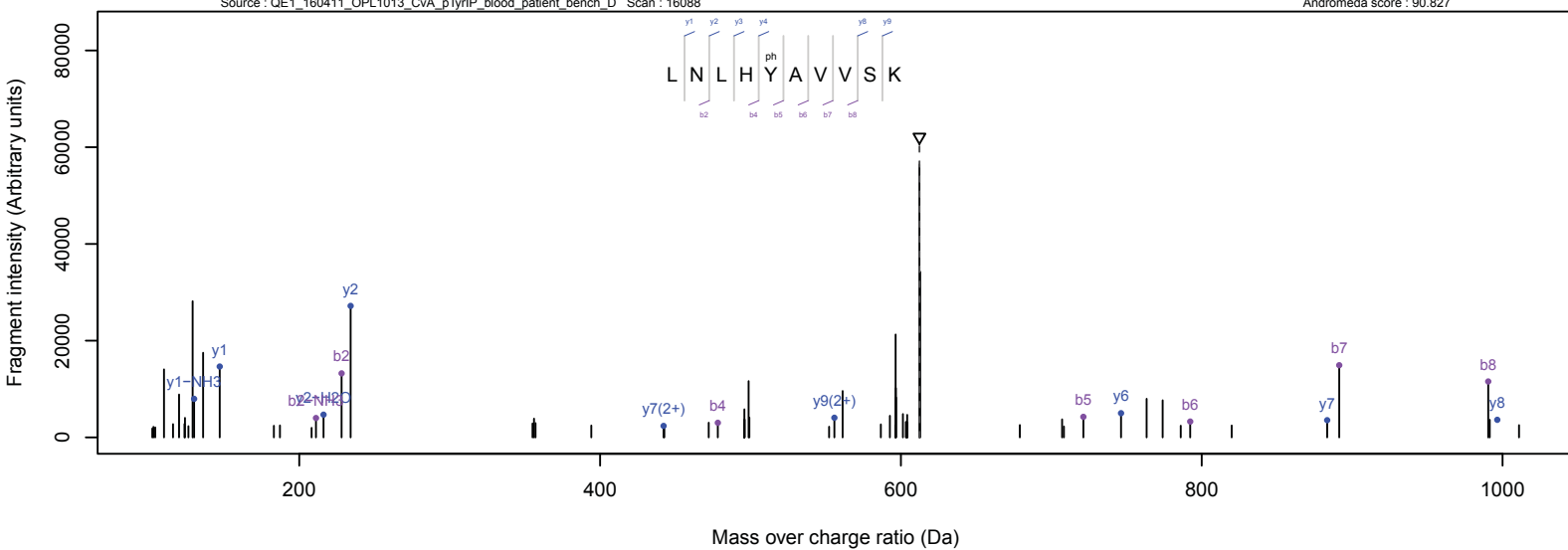
Andromeda score: 90.861



## LNLHY(ph)AVVSK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 16088

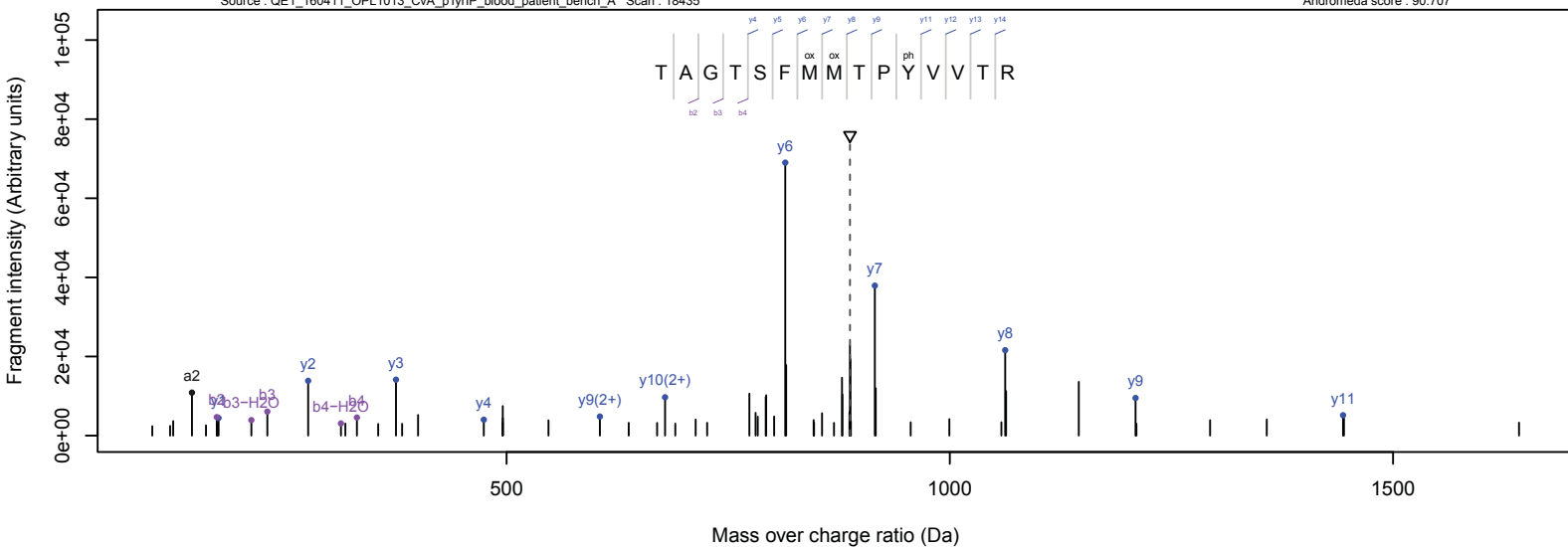
Andromeda score: 90.827



**TAGTSM(ox)M(ox)TPY(ph)VVTR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 18435

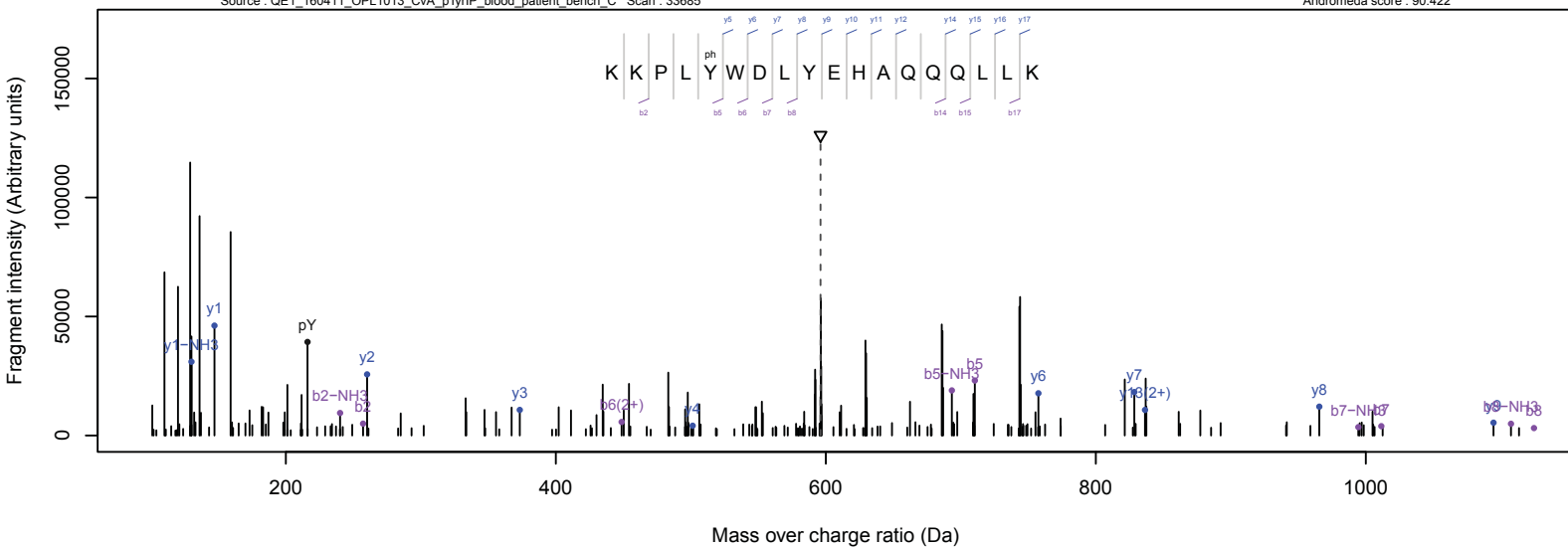
Andromeda score : 90.707



**KKPLY(ph)WDLYEHAQQQLK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 33685

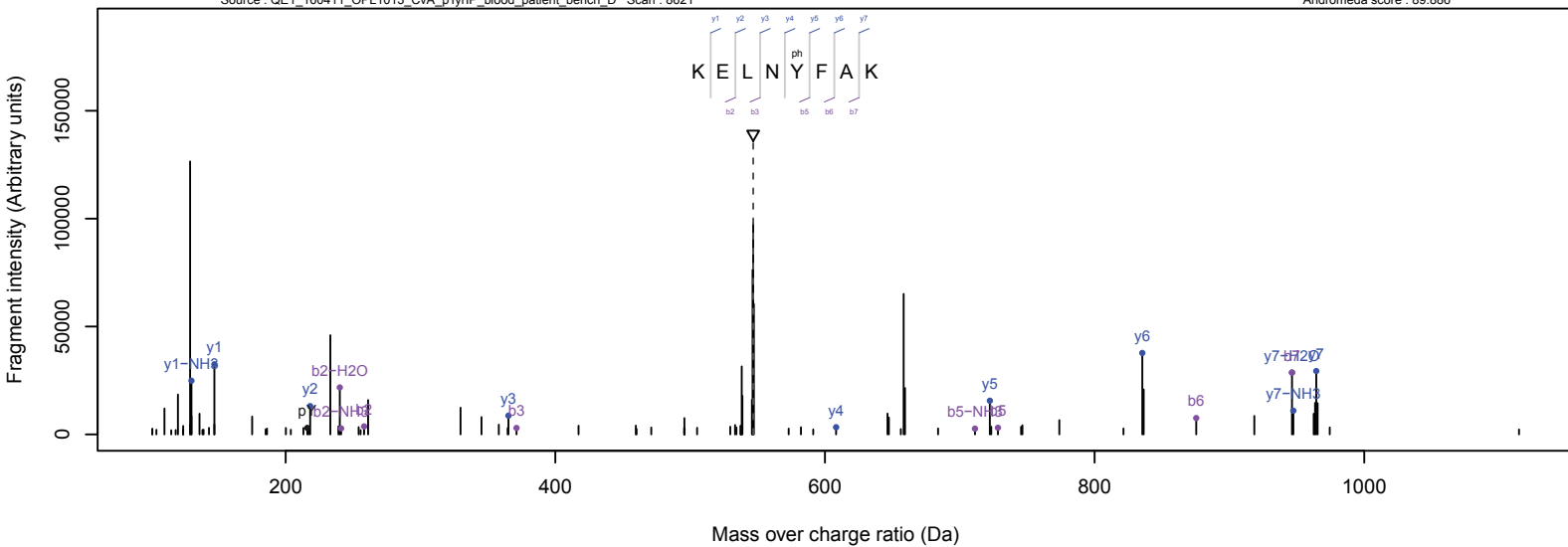
Andromeda score : 90.422



**KELNY(ph)FAK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 8621

Andromeda score : 89.886

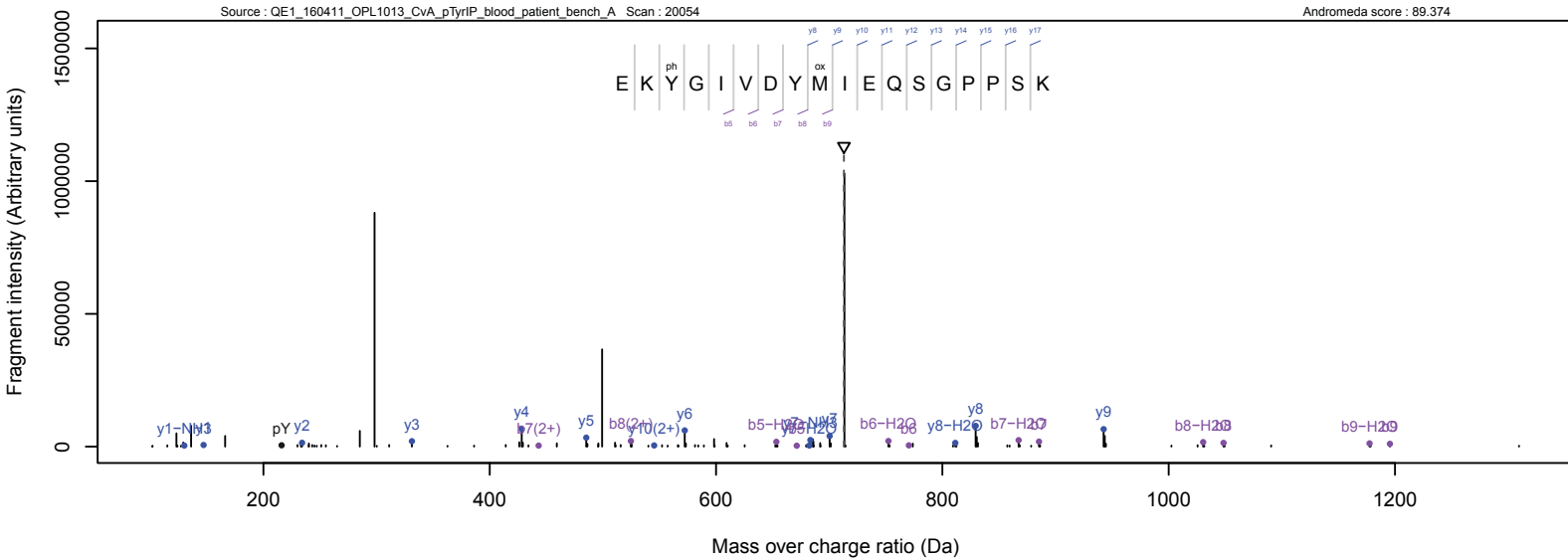




## EKY(ph)GIVDYM(ox)EQSGPPSK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 20054

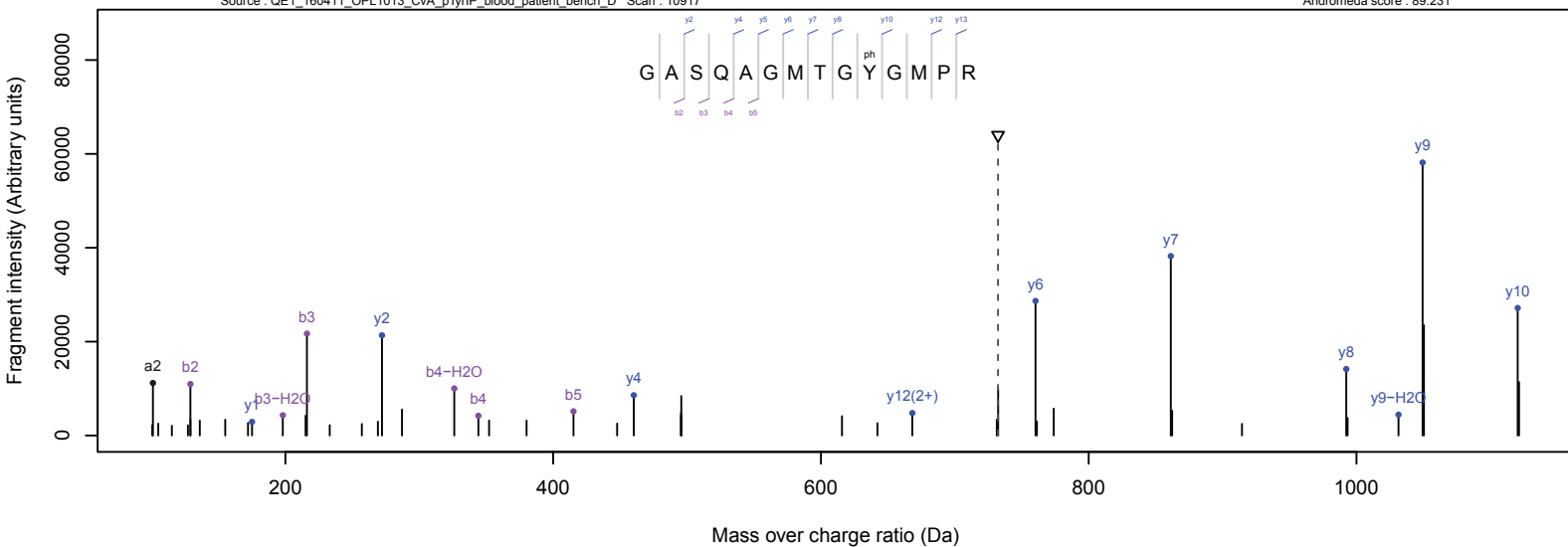
Andromeda score: 89.374



## GASQAGMTG(ph)GMPR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 10917

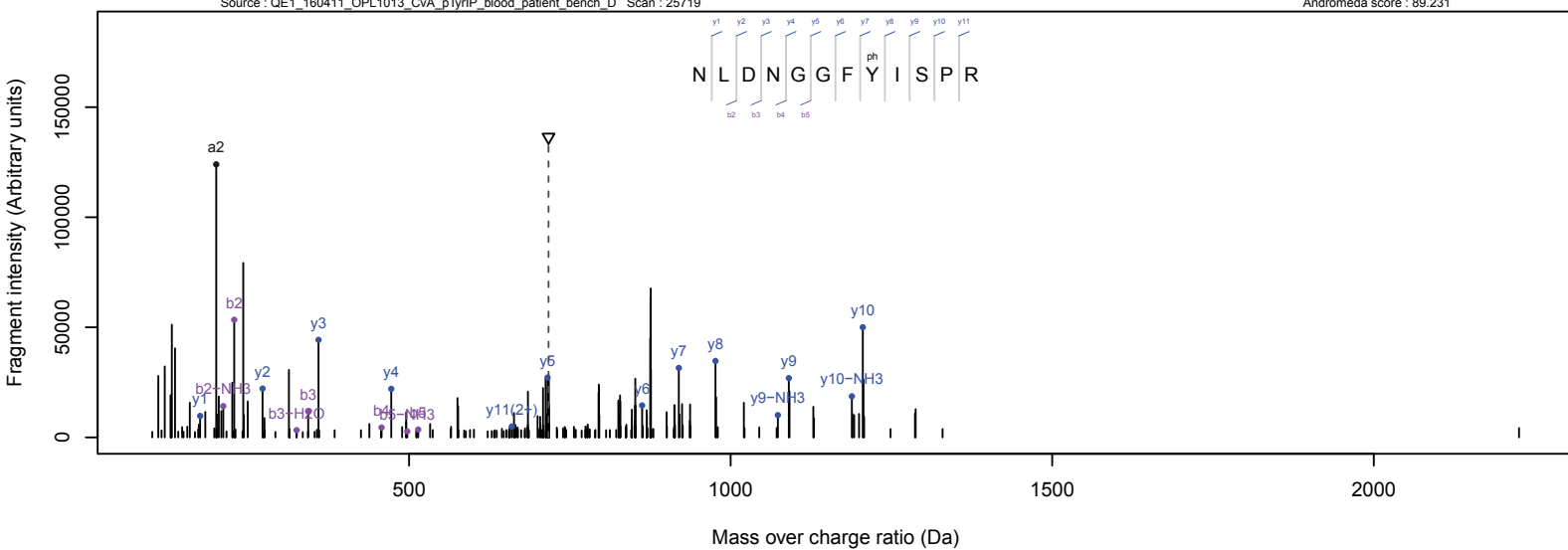
Andromeda score: 89.231



## NLDNGGFY(ph)ISPR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 25719

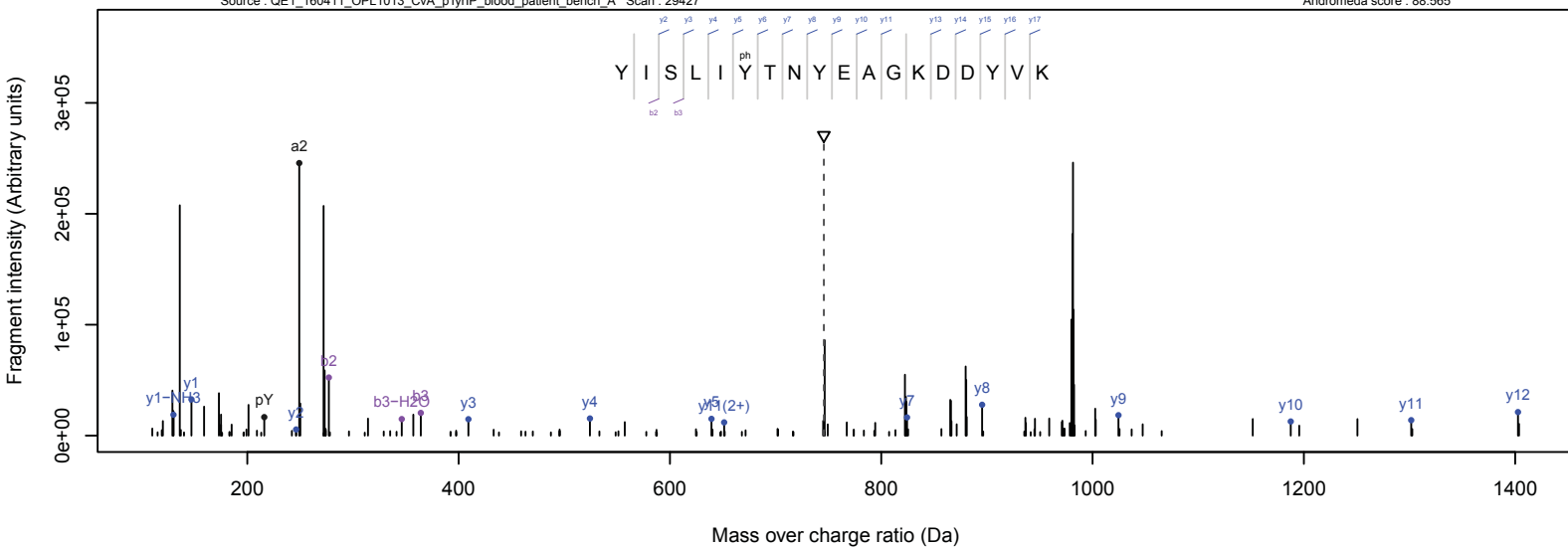
Andromeda score: 89.231



## YISLIY(ph)TNYEAGKDDYVK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 29427

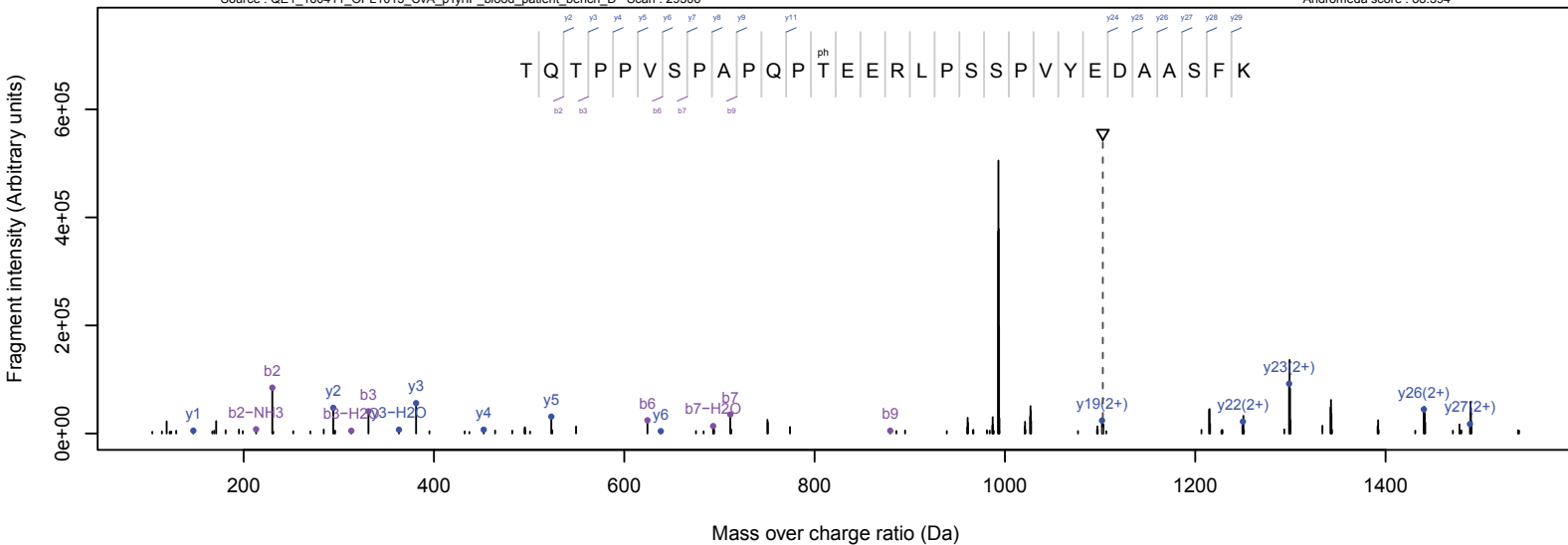
Andromeda score : 88.565



## TQTPPVPAPQPT(ph)EERLPSSPVYEDAASFK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 29306

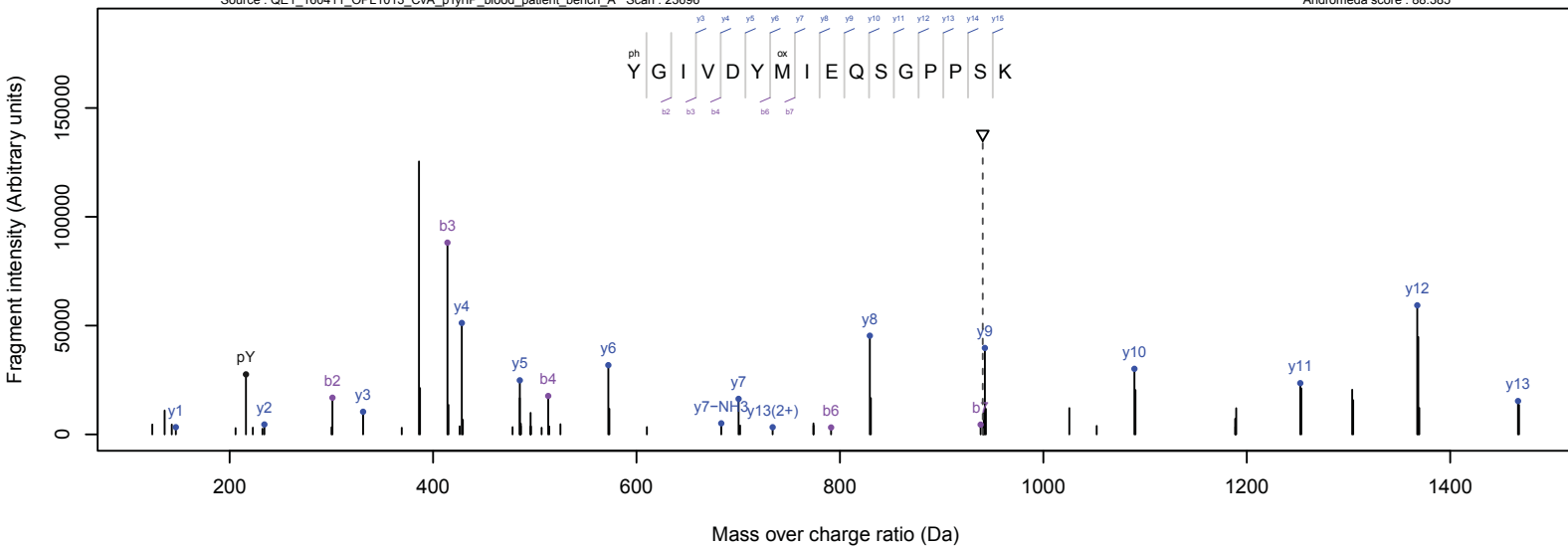
Andromeda score : 88.394



## Y(ph)GIVDYM(ox)IEQSGPPSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 23696

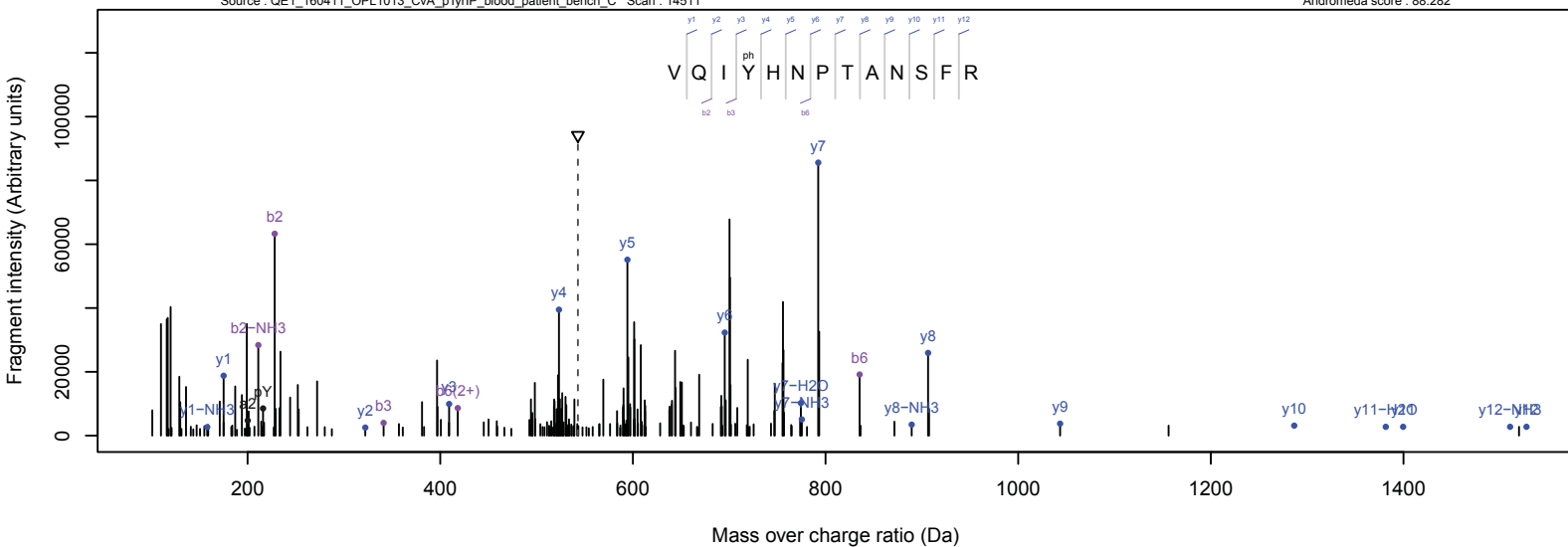
Andromeda score : 88.385



VQIY(ph)HNPTANSFR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 14511

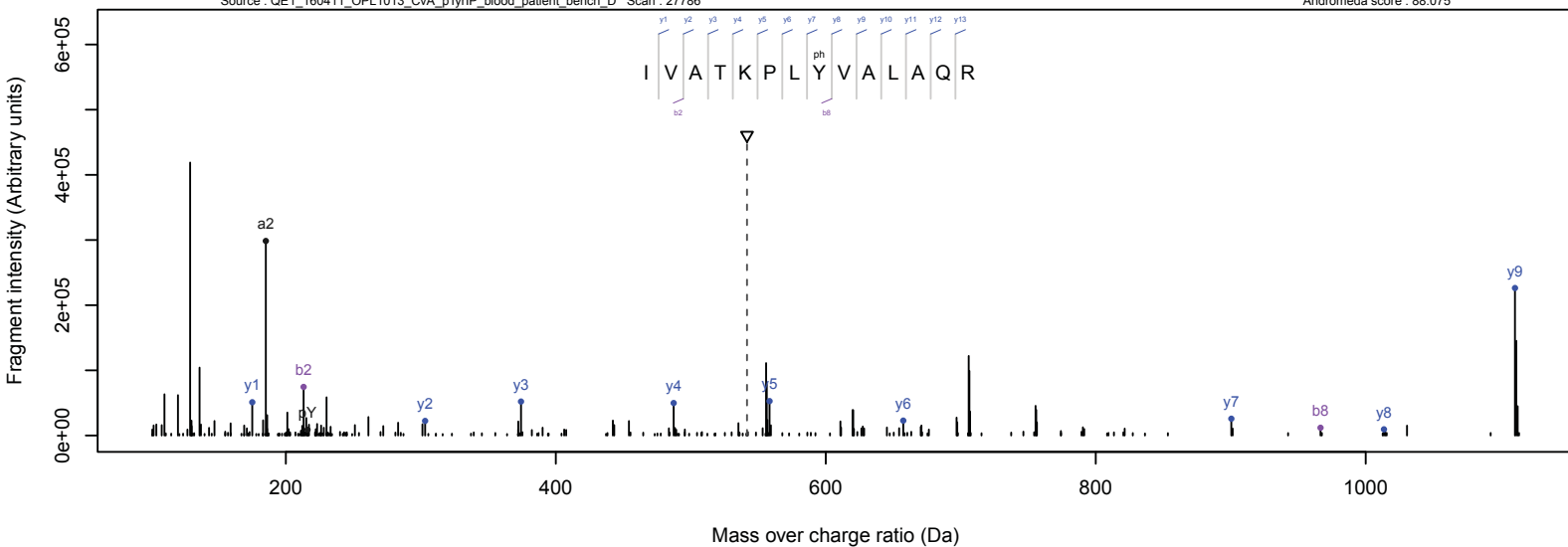
Andromeda score: 88.282



IVATKPLY(ph)VALAQR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 27786

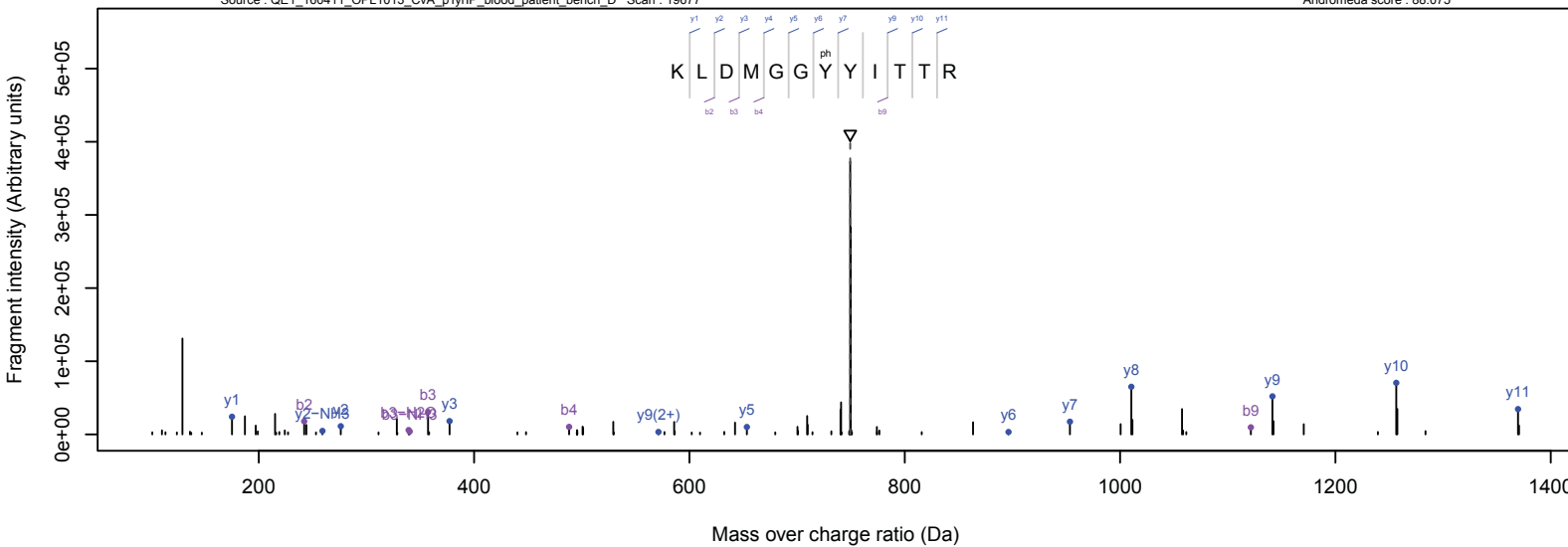
Andromeda score: 88.075



KLDMGY(ph)YITTR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 19677

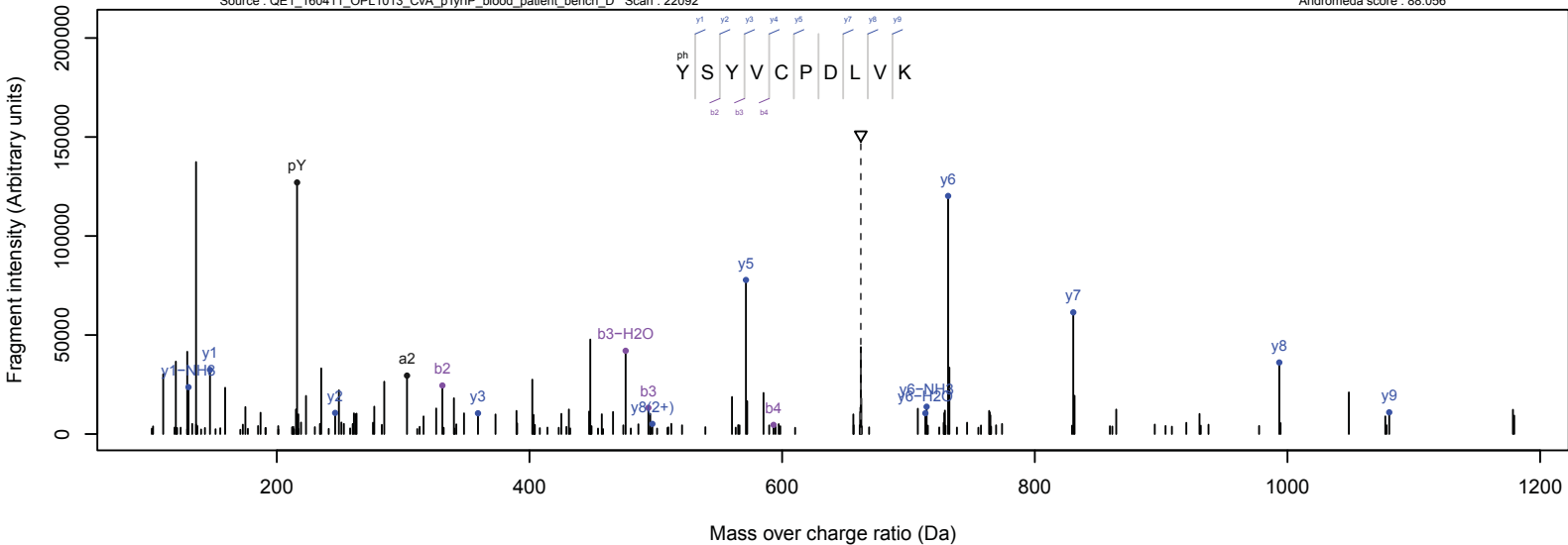
Andromeda score: 88.075



## Y(ph)SYVCPDLVK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 22092

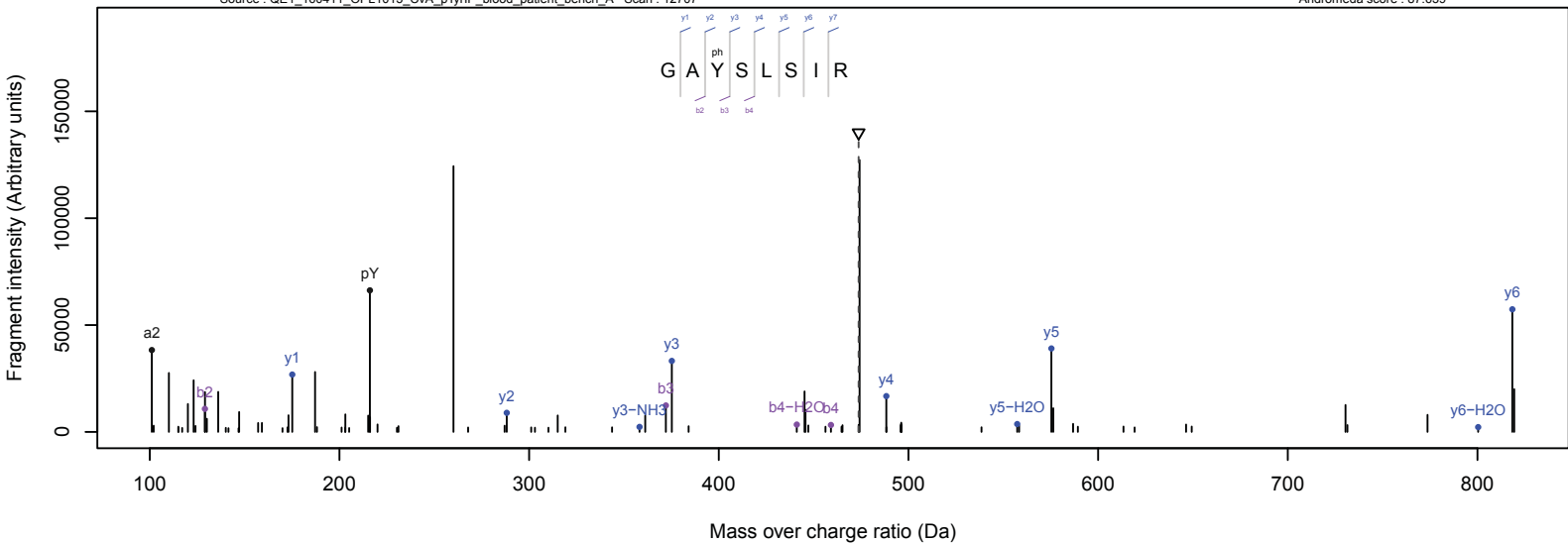
Andromeda score : 88.056



## GAY(ph)SLSIR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 12707

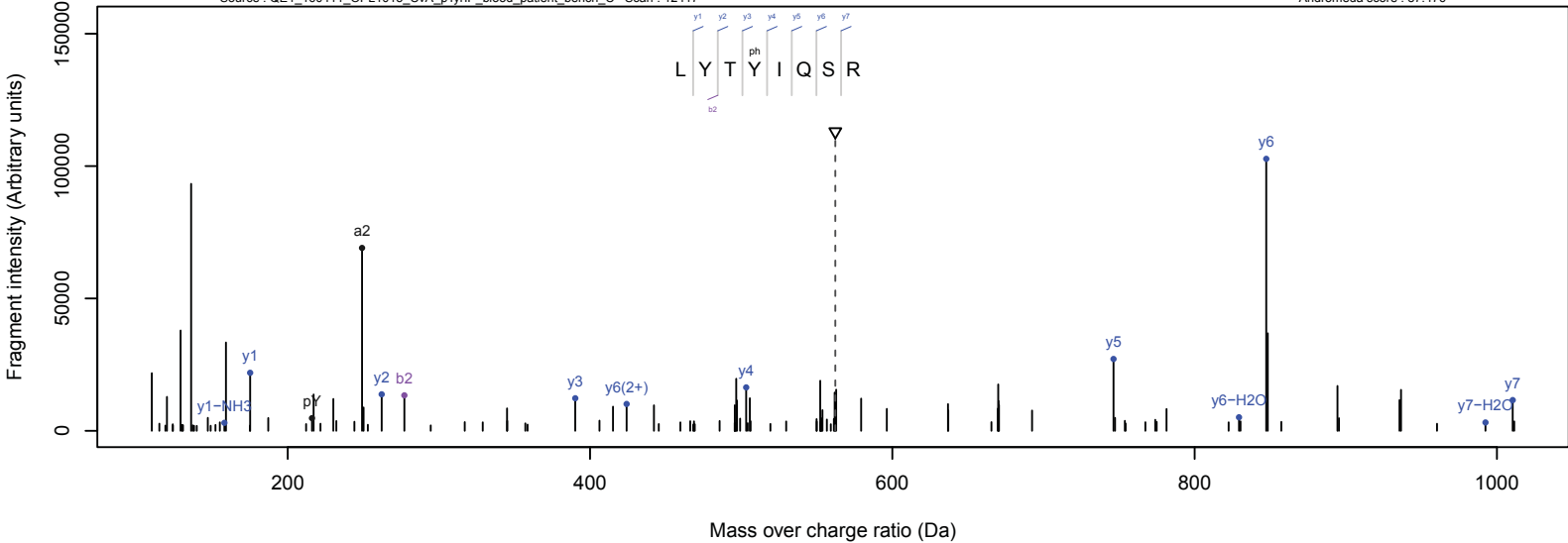
Andromeda score : 87.639



## LYTY(ph)QSR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 12417

Andromeda score : 87.476

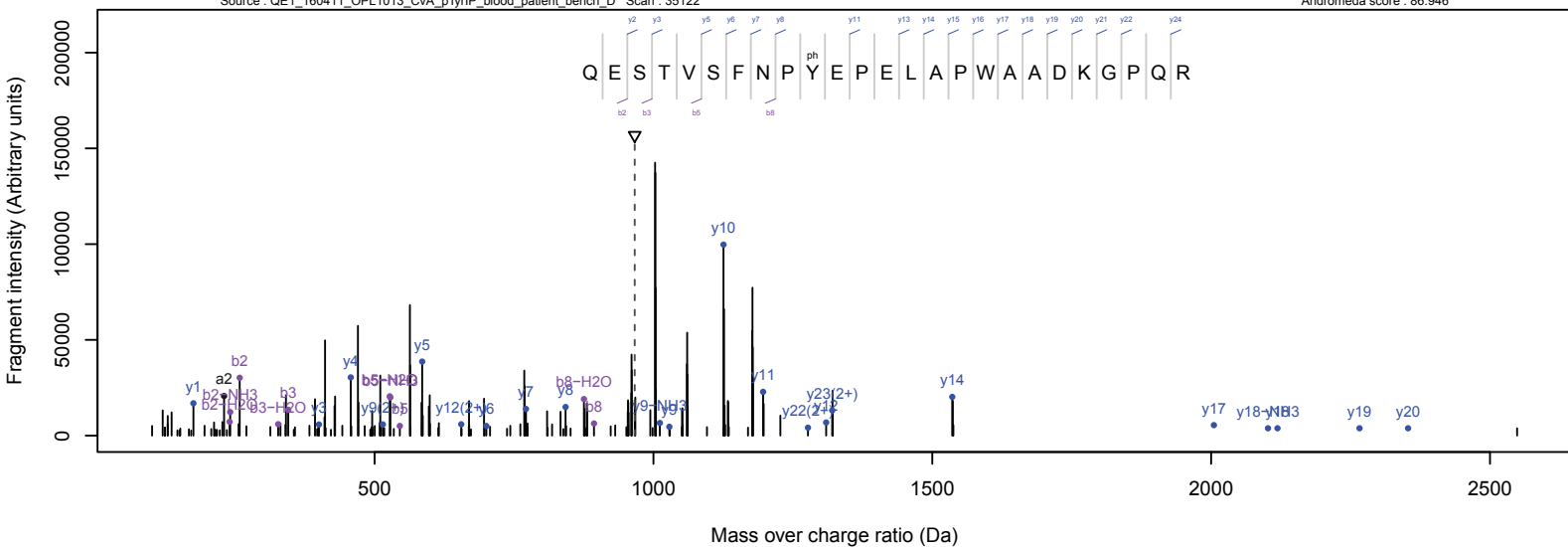




**QESTVSFNPY(ph)EPELAPWAADKGPQR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 35122

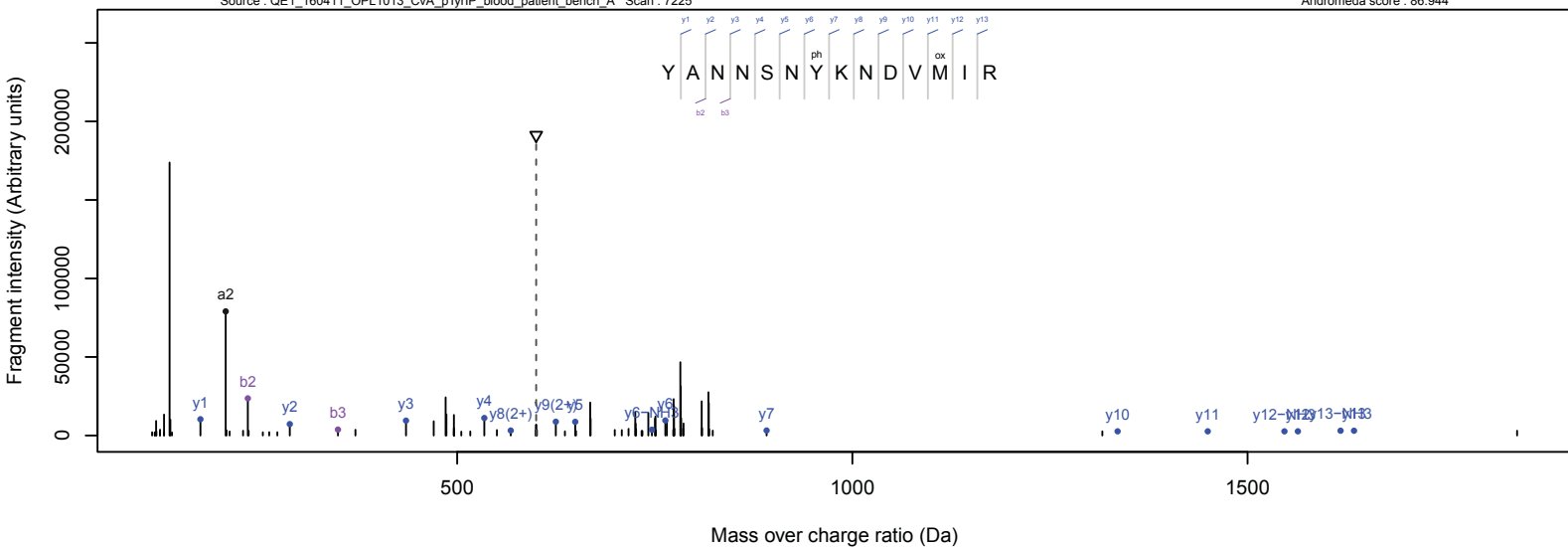
Andromeda score : 86.946



**YANNSNY(ph)KNDVMOIR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 7225

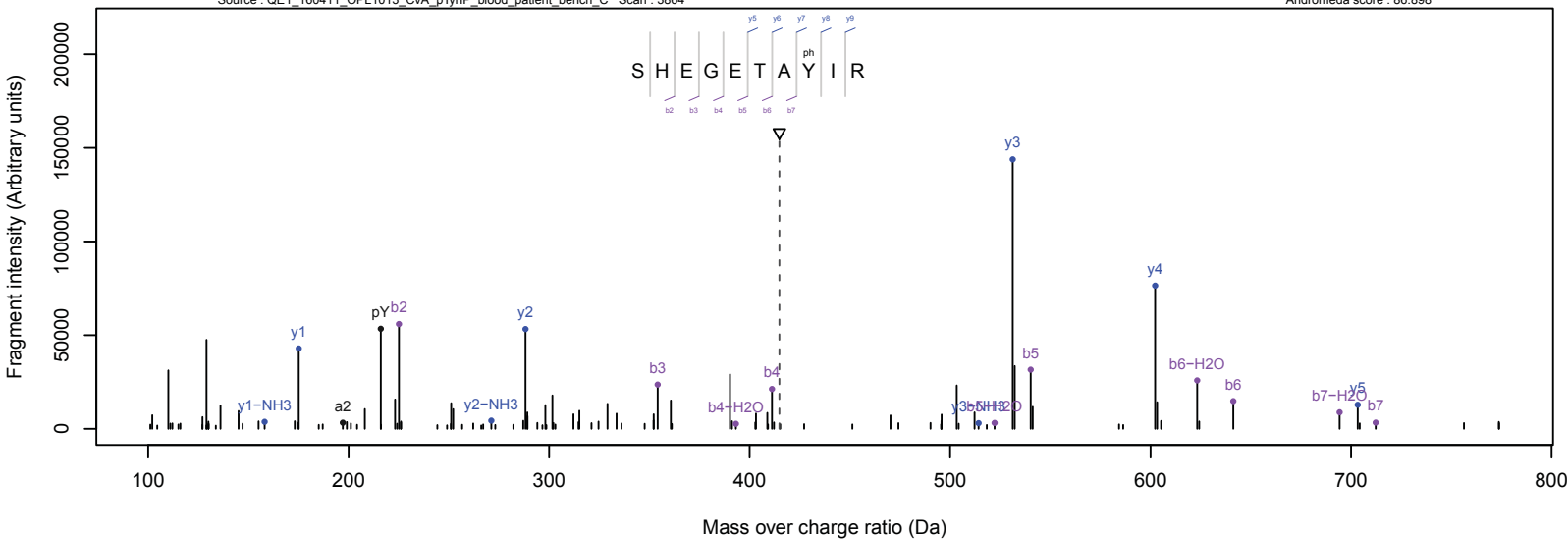
Andromeda score : 86.944



**SHEGETAY(ph)IR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3864

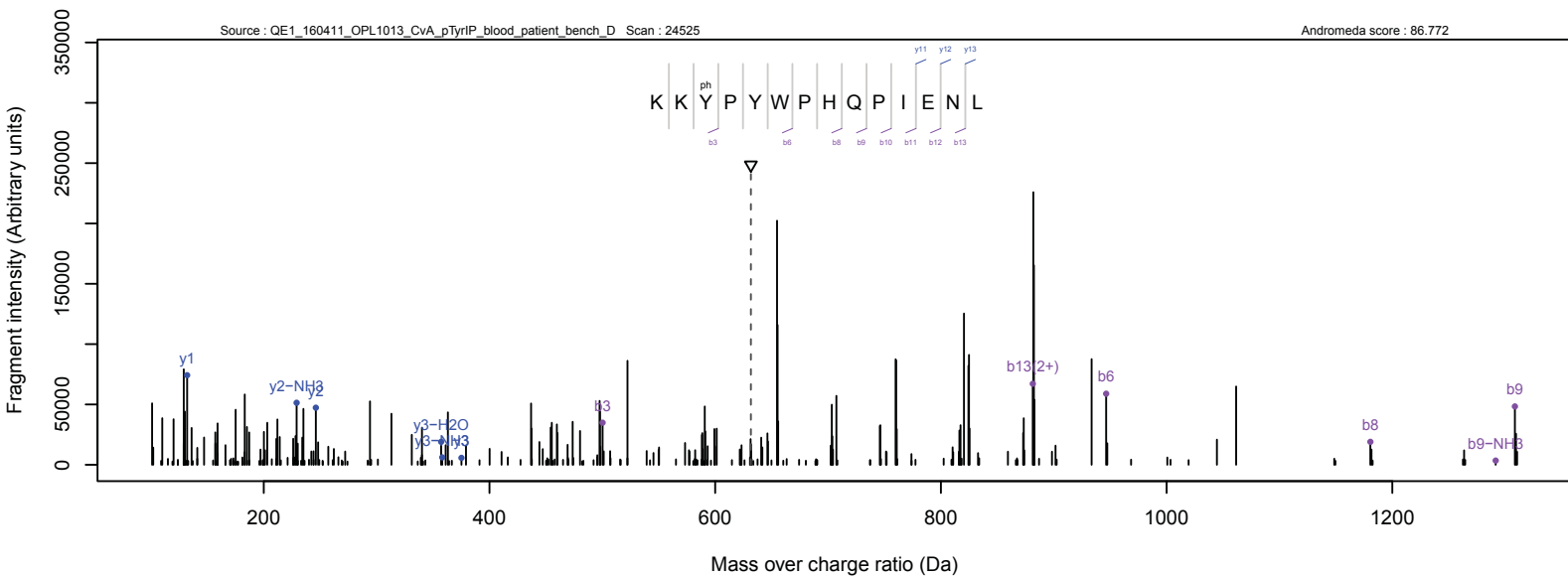
Andromeda score : 86.898



## KKY(ph)PYWPHQPIENL-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24525

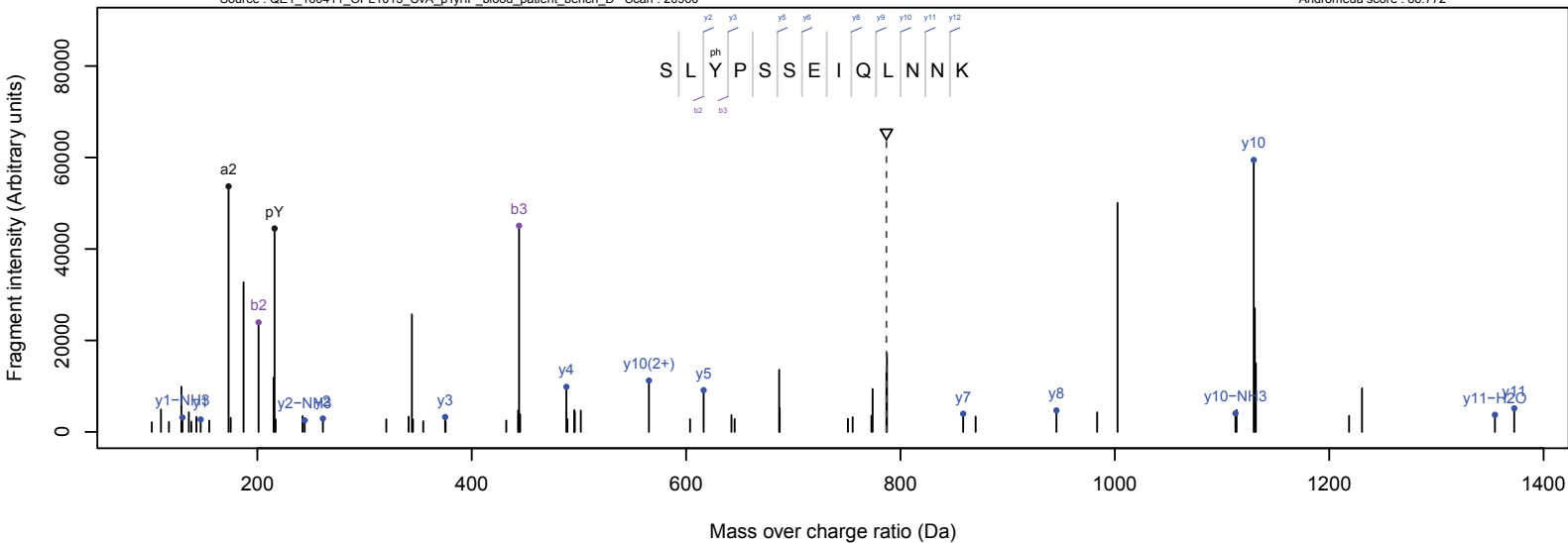
Andromeda score : 86.772



## SLY(ph)PSSEIQLNKK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 20960

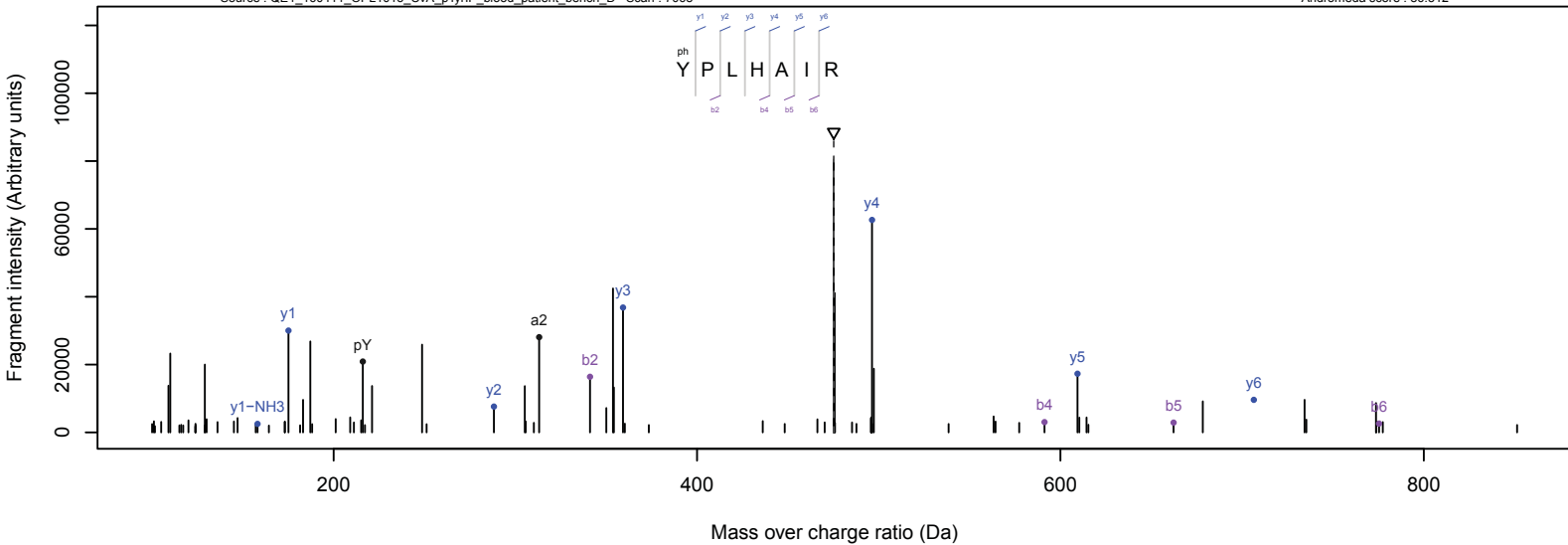
Andromeda score : 86.772

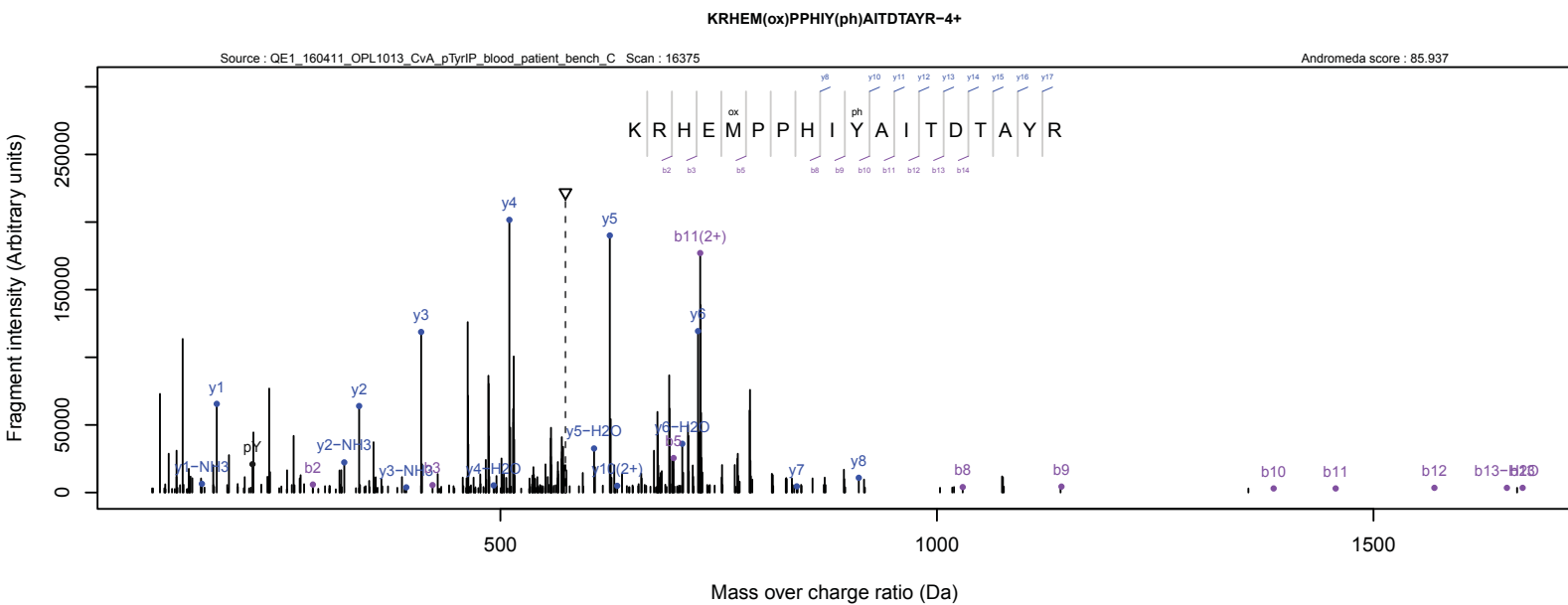
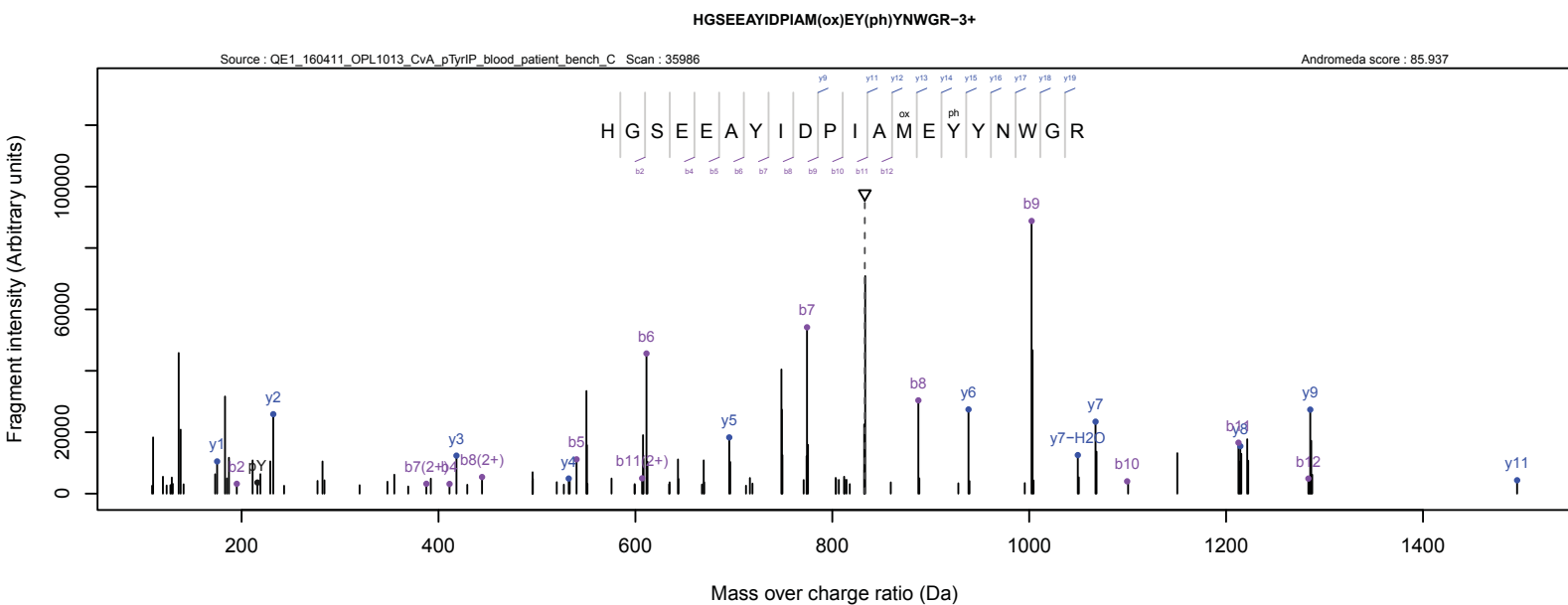
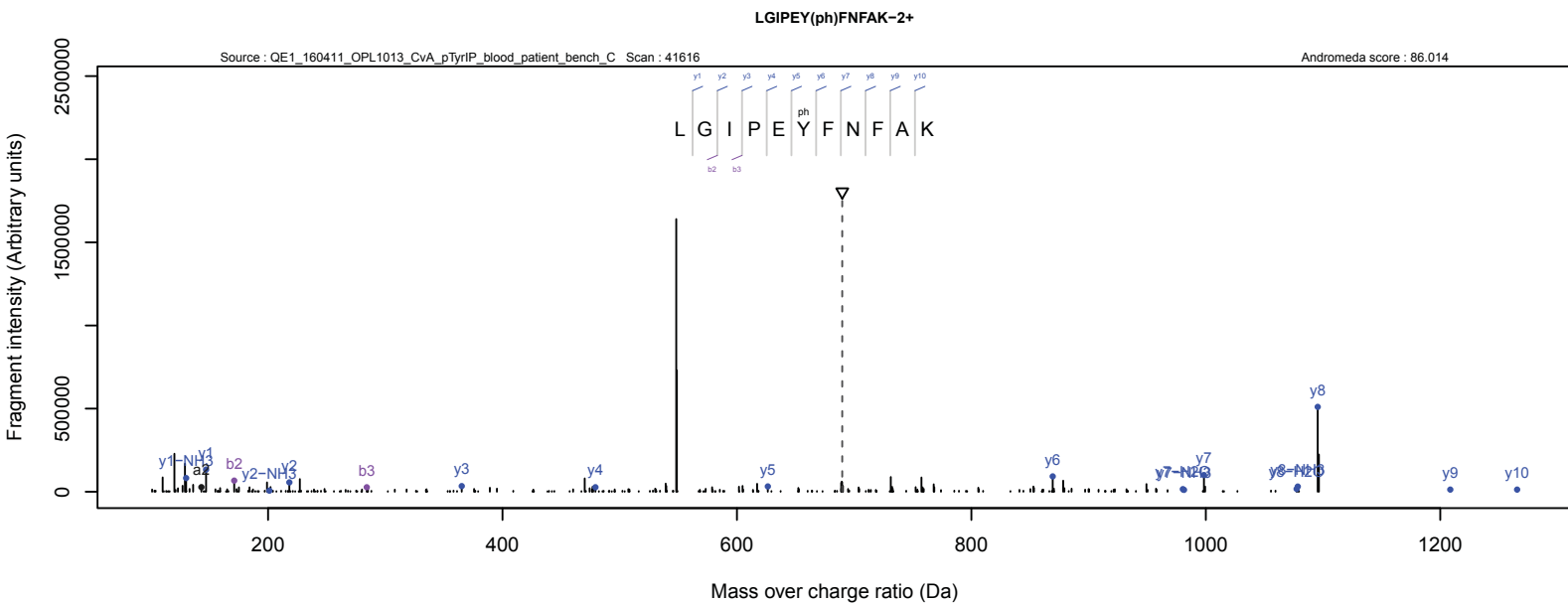


## Y(ph)PLHAIR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7068

Andromeda score : 86.512



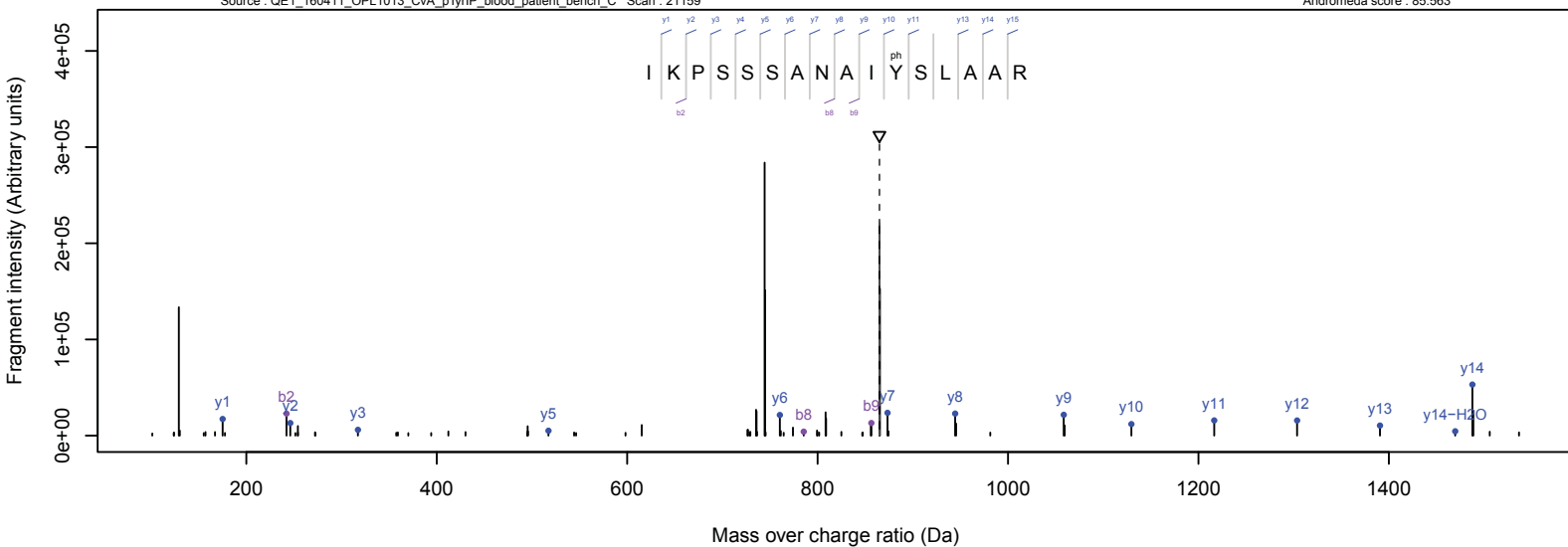




**IKPSSANAIY(ph)SLAAR-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 21159

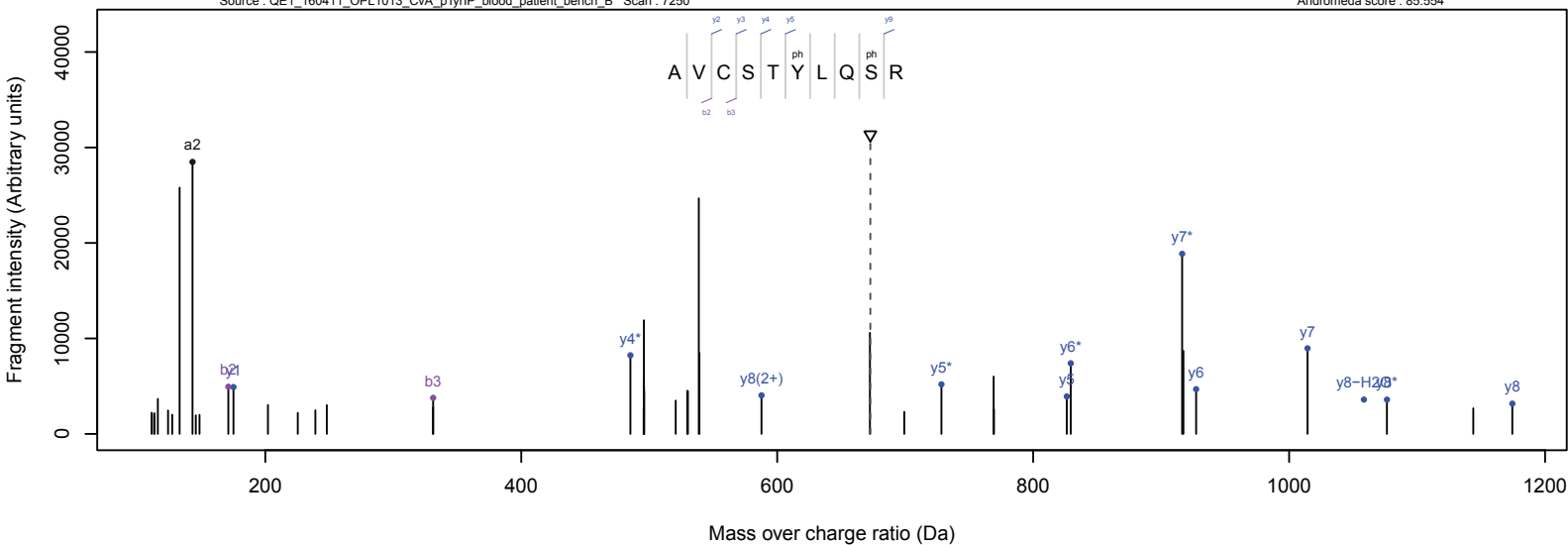
Andromeda score: 85.563



**AVCSTY(ph)LQS(ph)R-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 7250

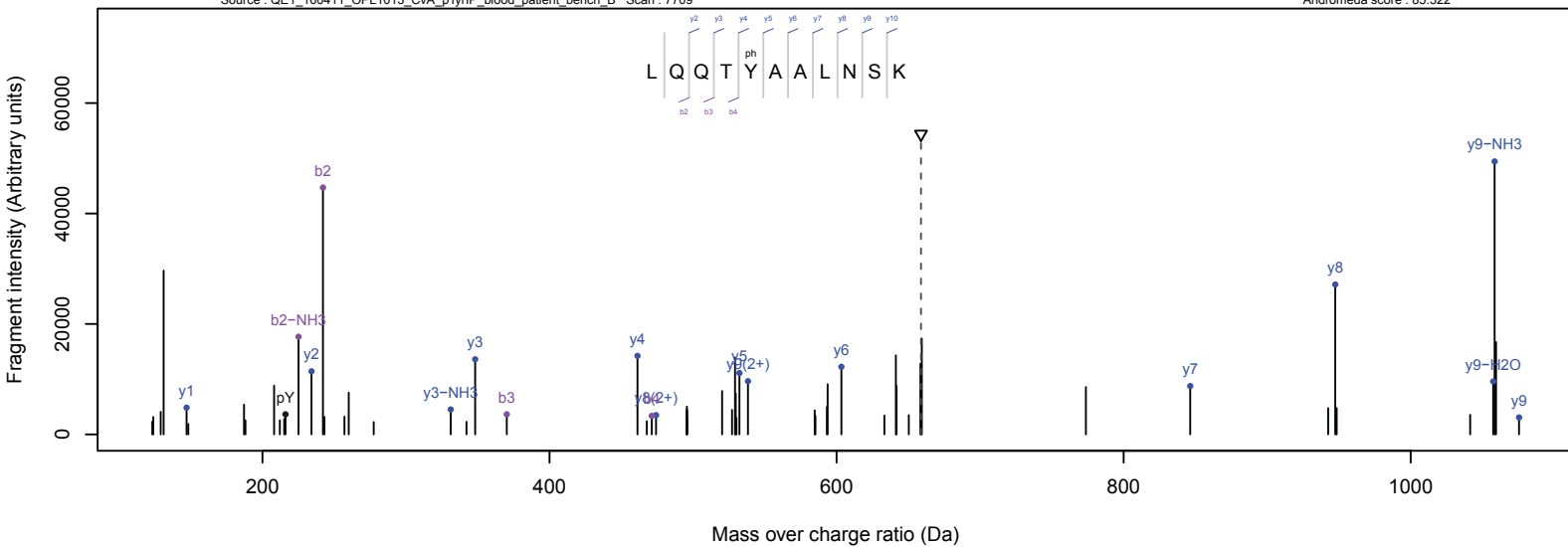
Andromeda score: 85.554



**LQQTY(ph)AALNSK-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 7709

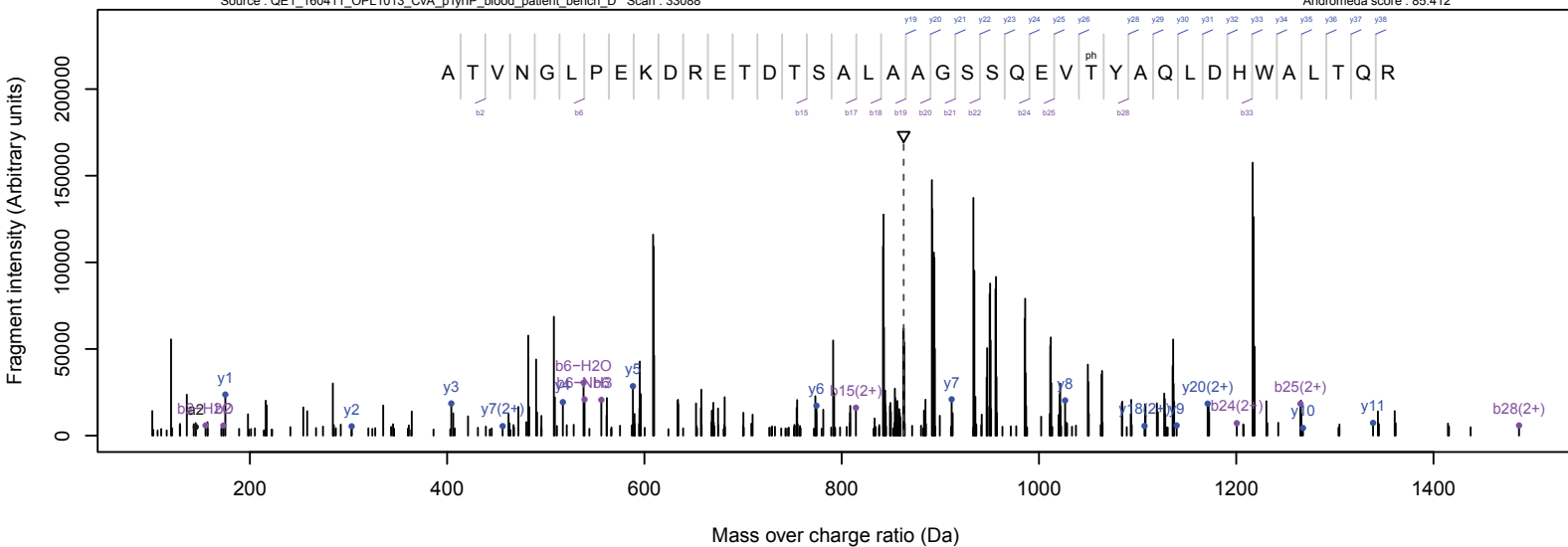
Andromeda score: 85.522



ATVNLPEKRETDTSALAAAGSSQEV<sup>ph</sup>YAQLDHWALTQR-5+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 33088

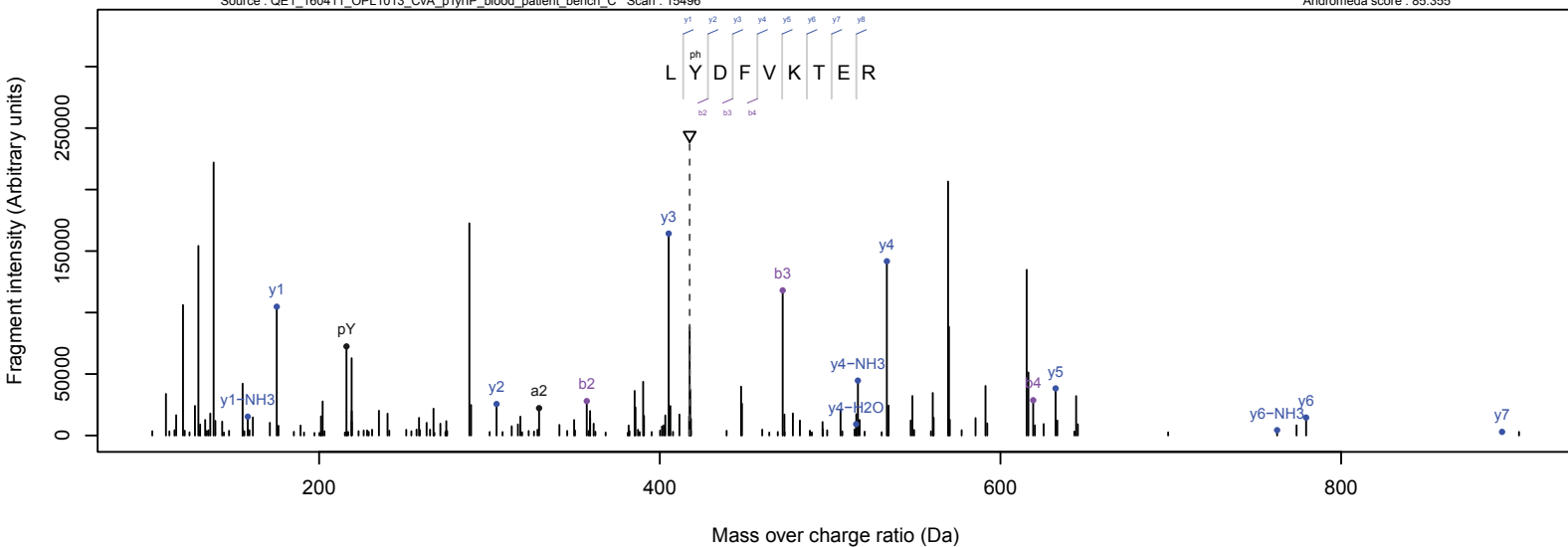
Andromeda score : 85.412



LY<sup>ph</sup>DFVKTER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 15496

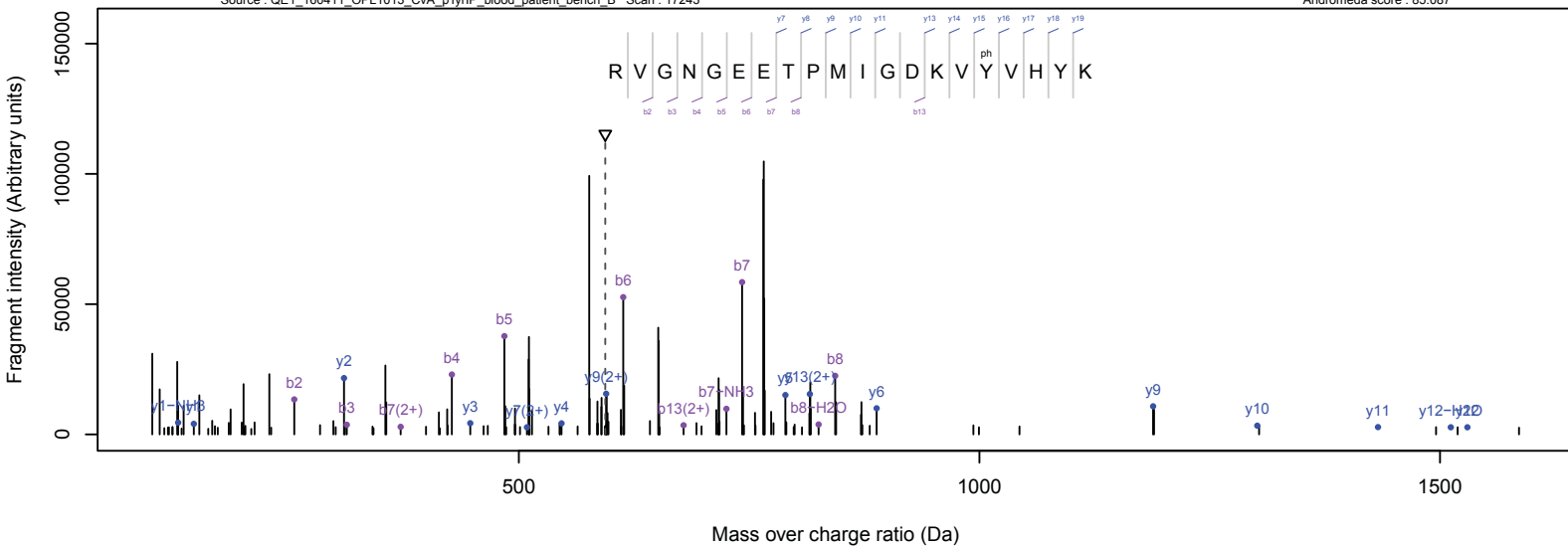
Andromeda score : 85.355



RVNGEETPMIGDKV<sup>ph</sup>VHYK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 17243

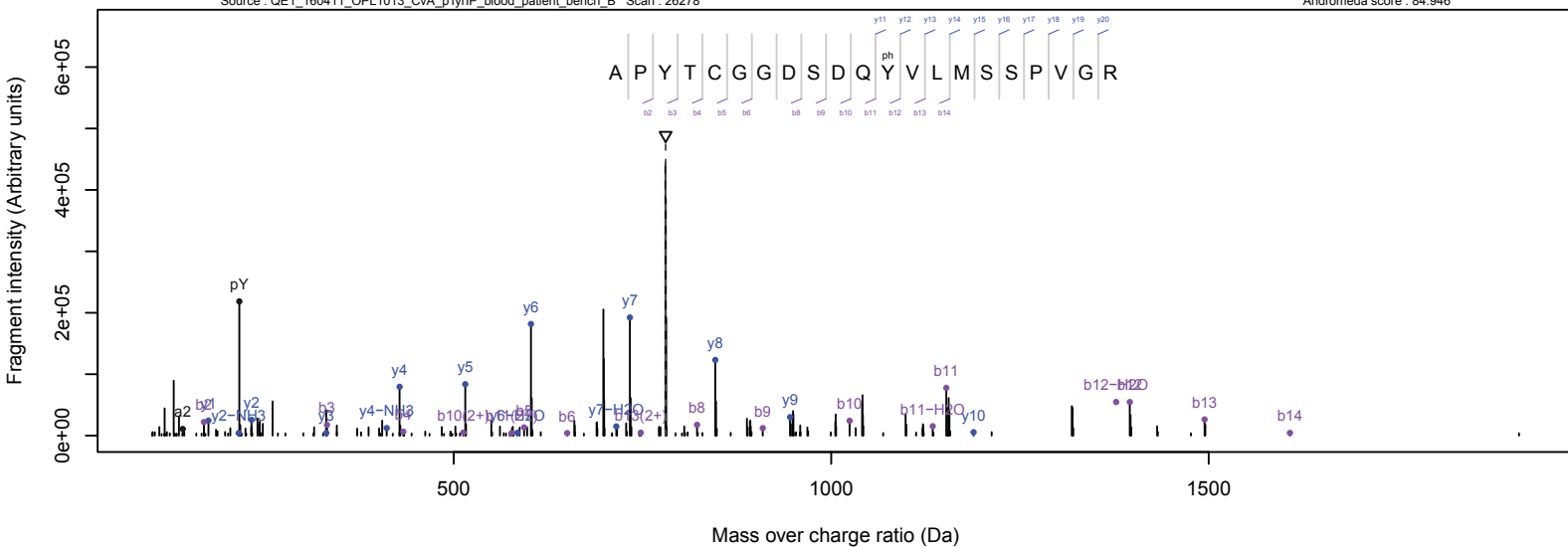
Andromeda score : 85.087



APYTCGGSDQY(ph)VLMSSPVGR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 26278

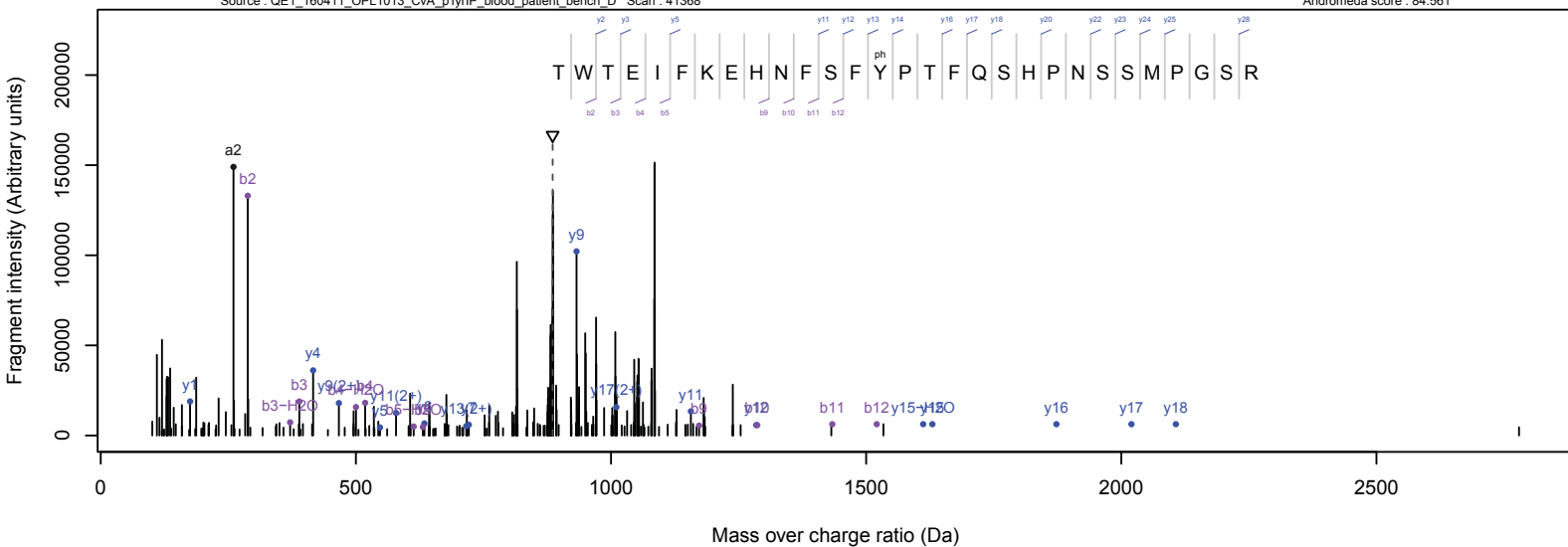
Andromeda score : 84.946



TWTEIFKEHNFSFY(ph)PTFQSHPNSSMPGSR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 41368

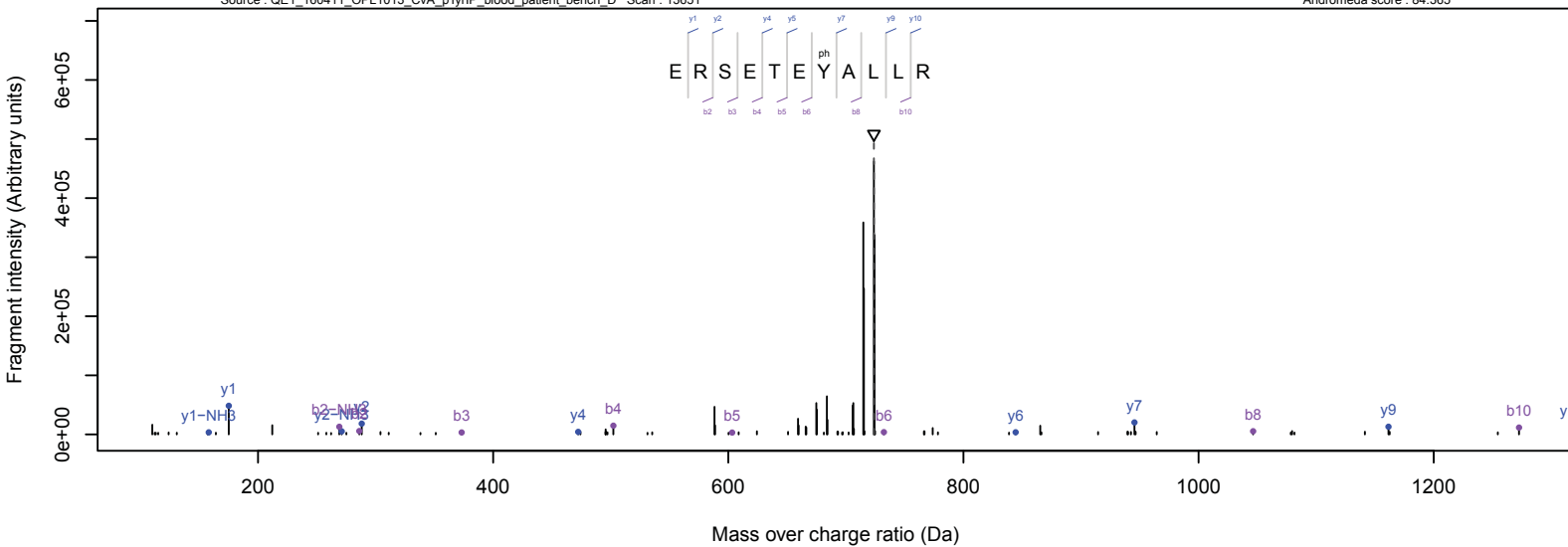
Andromeda score : 84.561



ERSETEY(ph)ALLR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 13651

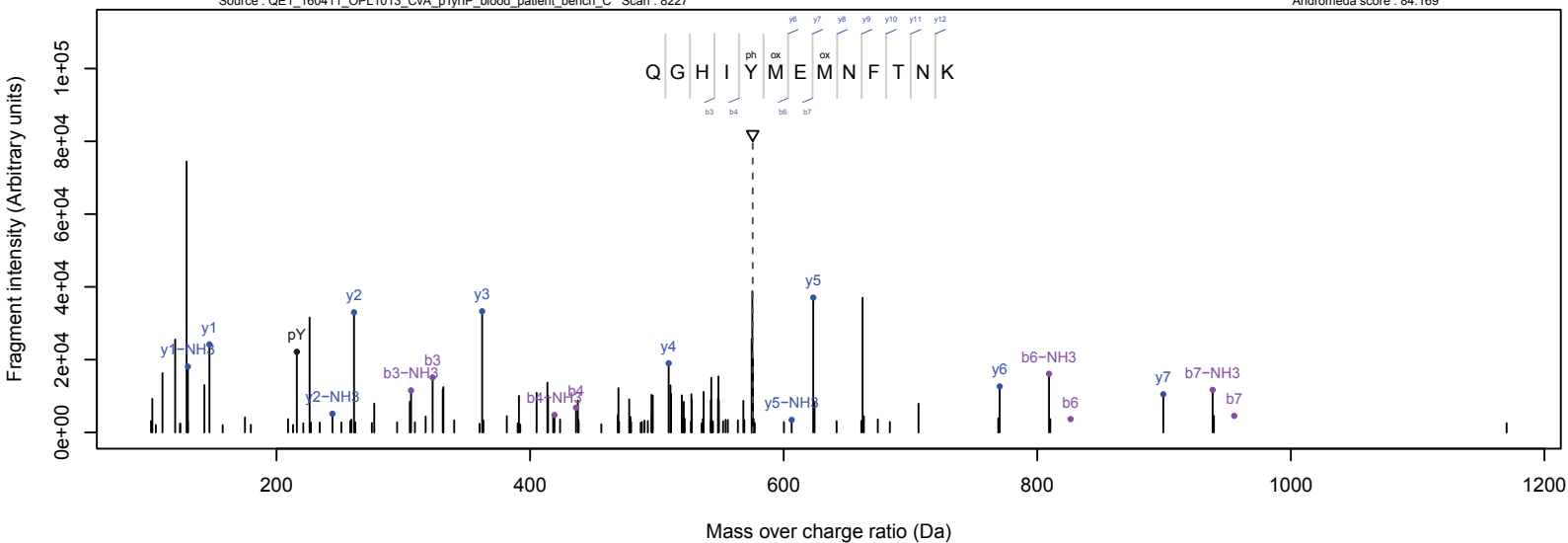
Andromeda score : 84.365



**QGHYIY<sup>ph</sup>M<sup>ox</sup>EM<sup>ox</sup>NFTNK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 8227

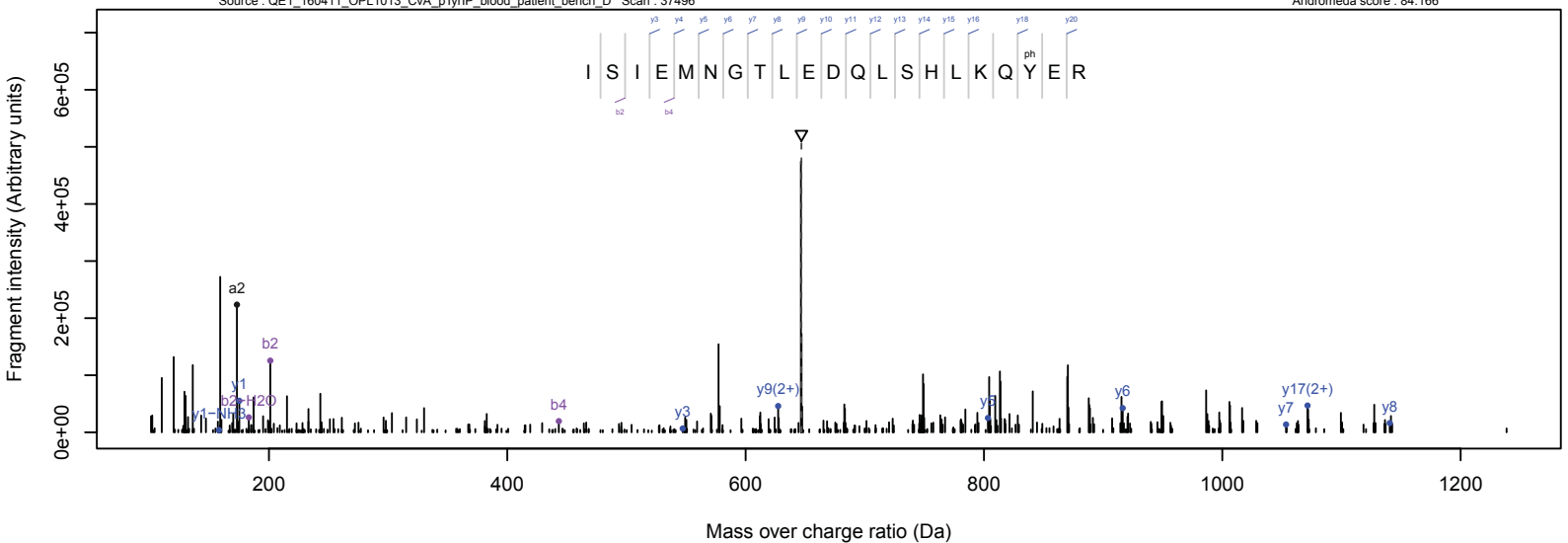
Andromeda score : 84.169



**ISIEMNGTLEDQLSHLKQY<sup>ph</sup>ER-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 37496

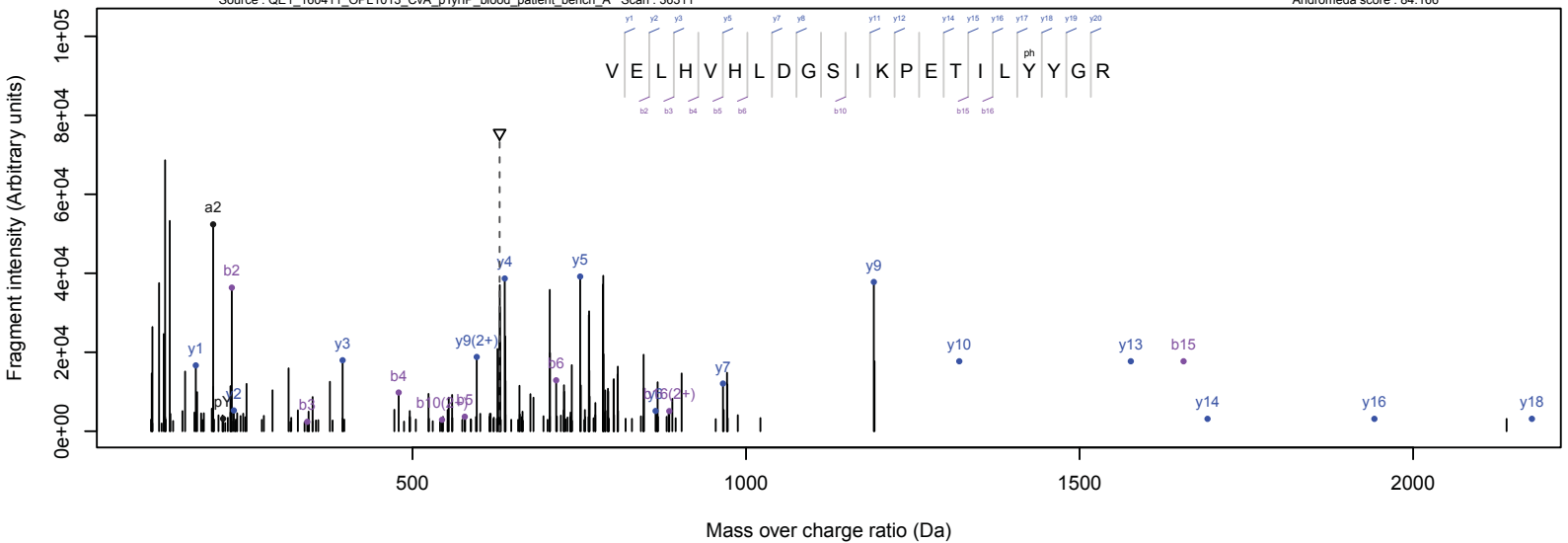
Andromeda score : 84.166



**VELHVHLDGSIKPETILY<sup>ph</sup>YGR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 36311

Andromeda score : 84.166

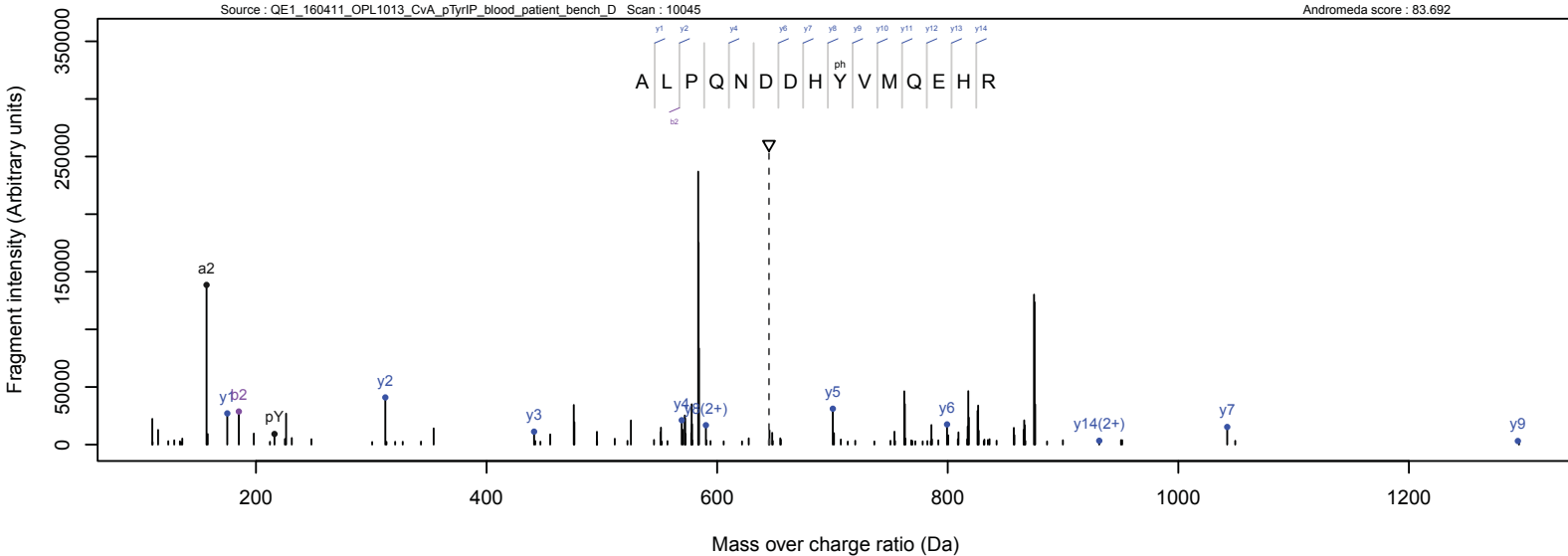




ALPQDDHY(ph)VMQEHR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 10045

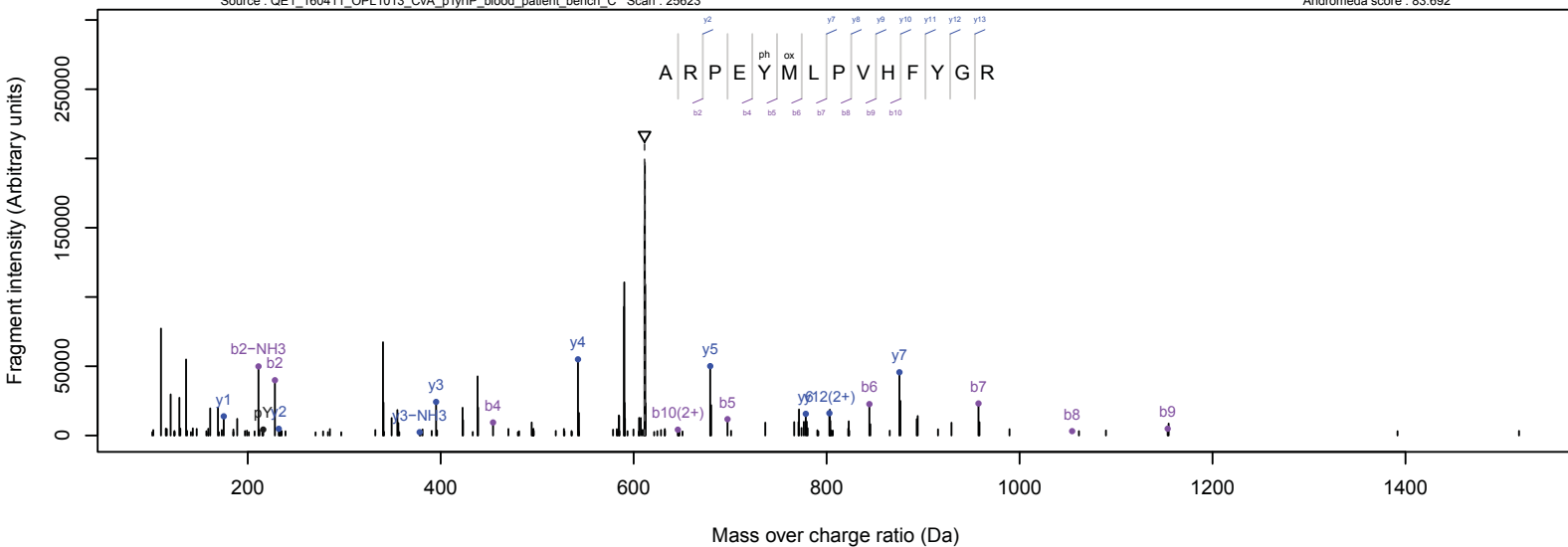
Andromeda score : 83.692



ARPEY(ph)M(ox)LPVHFYGR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 25623

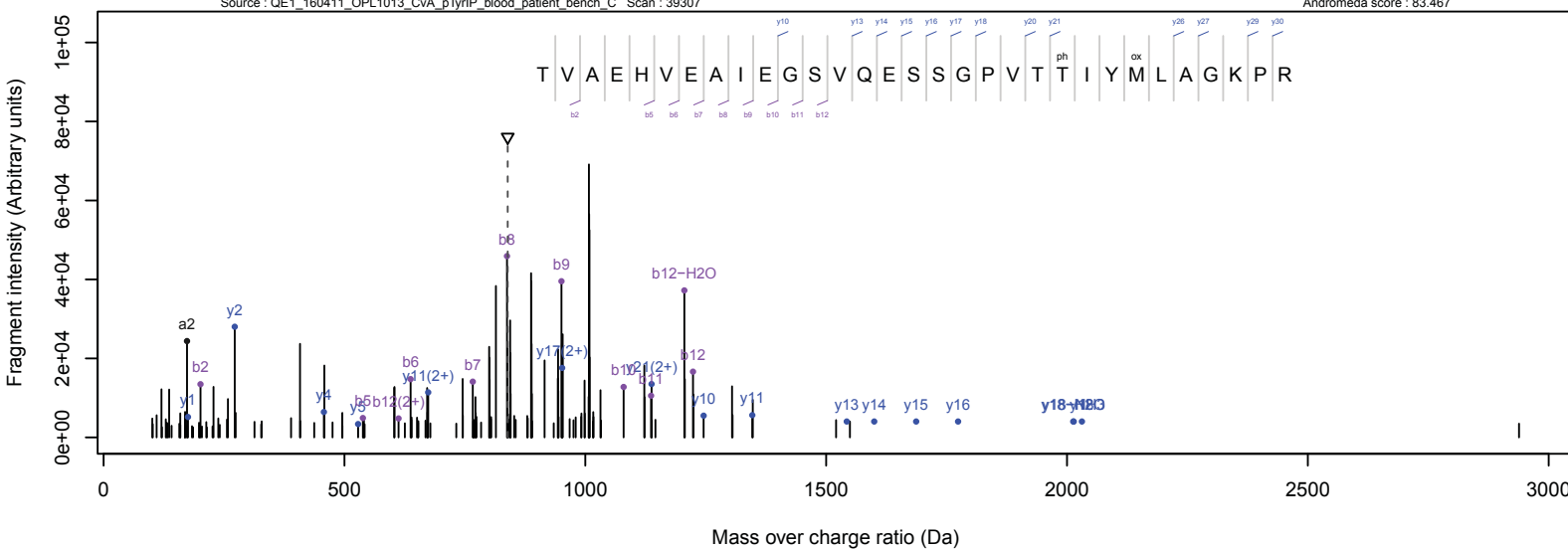
Andromeda score : 83.692



TVAEHVEAIEGSSGPVTT(ph)IYM(ox)LAGKPR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 39307

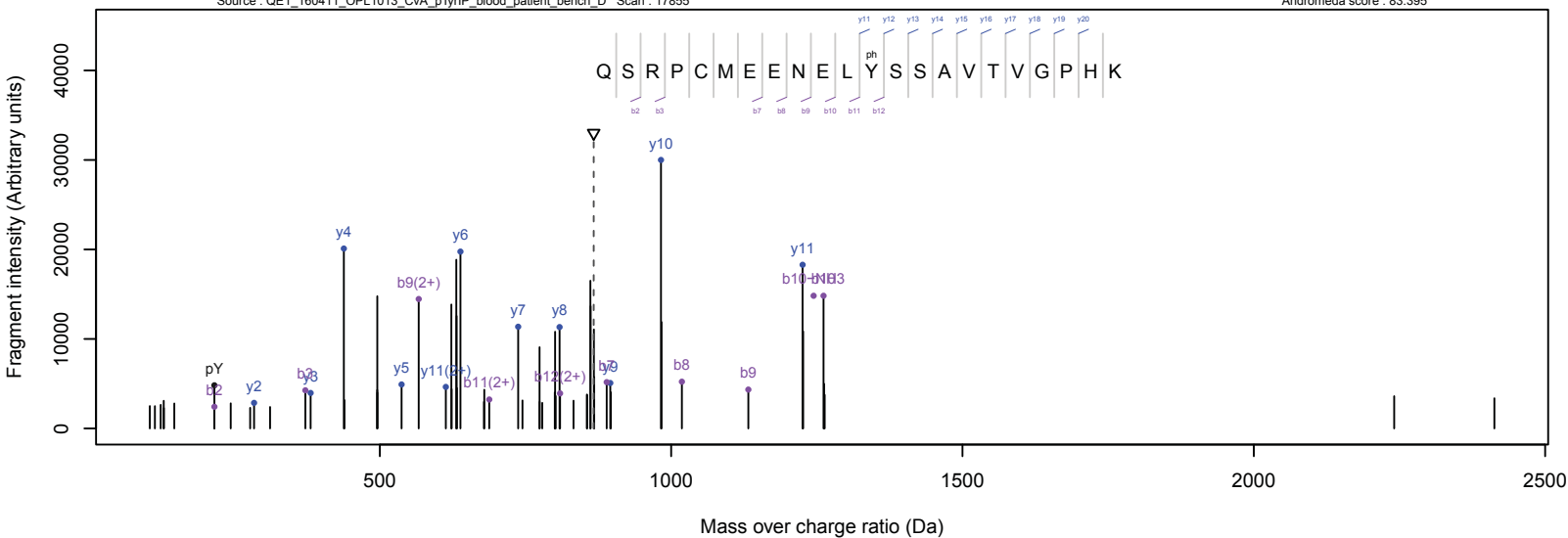
Andromeda score : 83.467



QSRPCMEENEELYS(A)SSAVTVGPHK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 17855

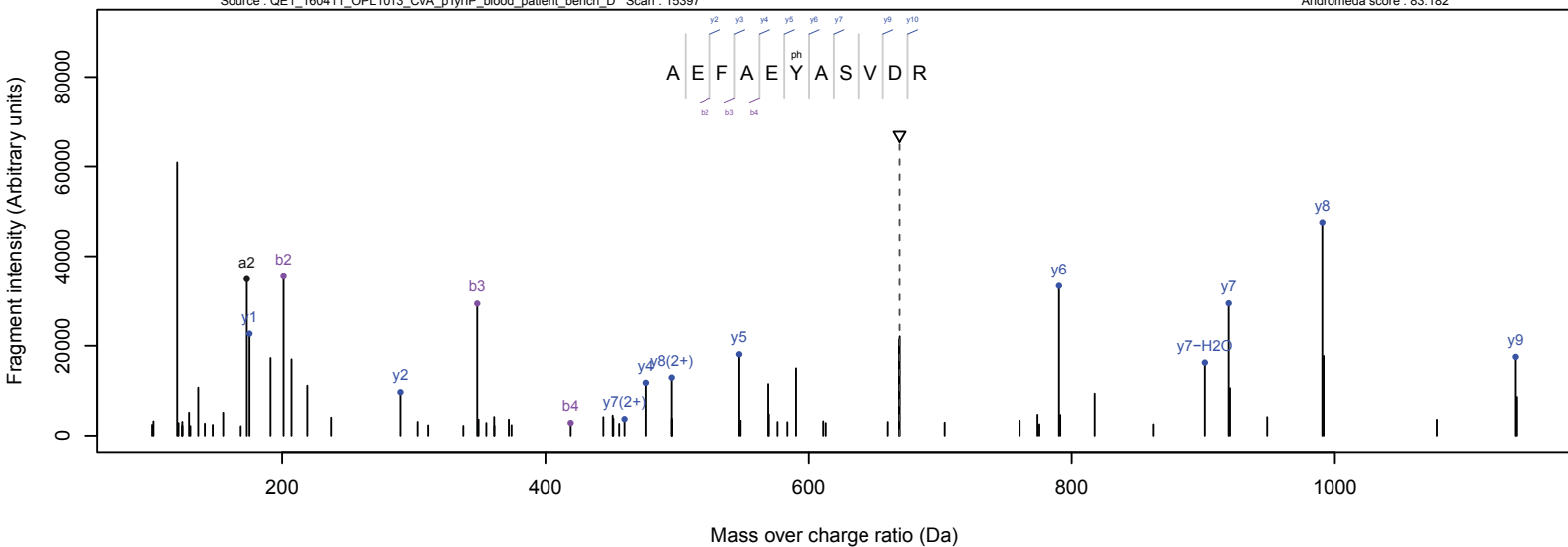
Andromeda score: 83.395



AEFAEY(A)ASVDR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 15397

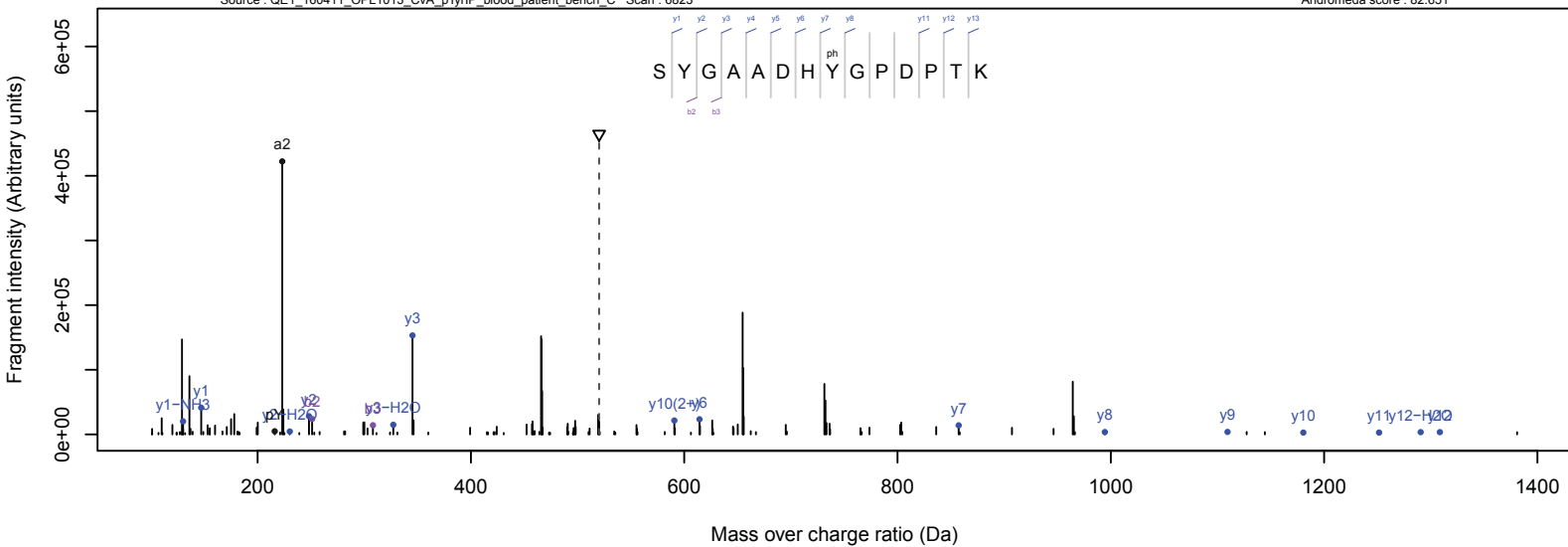
Andromeda score: 83.182



SYGAADHY(A)GPDPTK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 6823

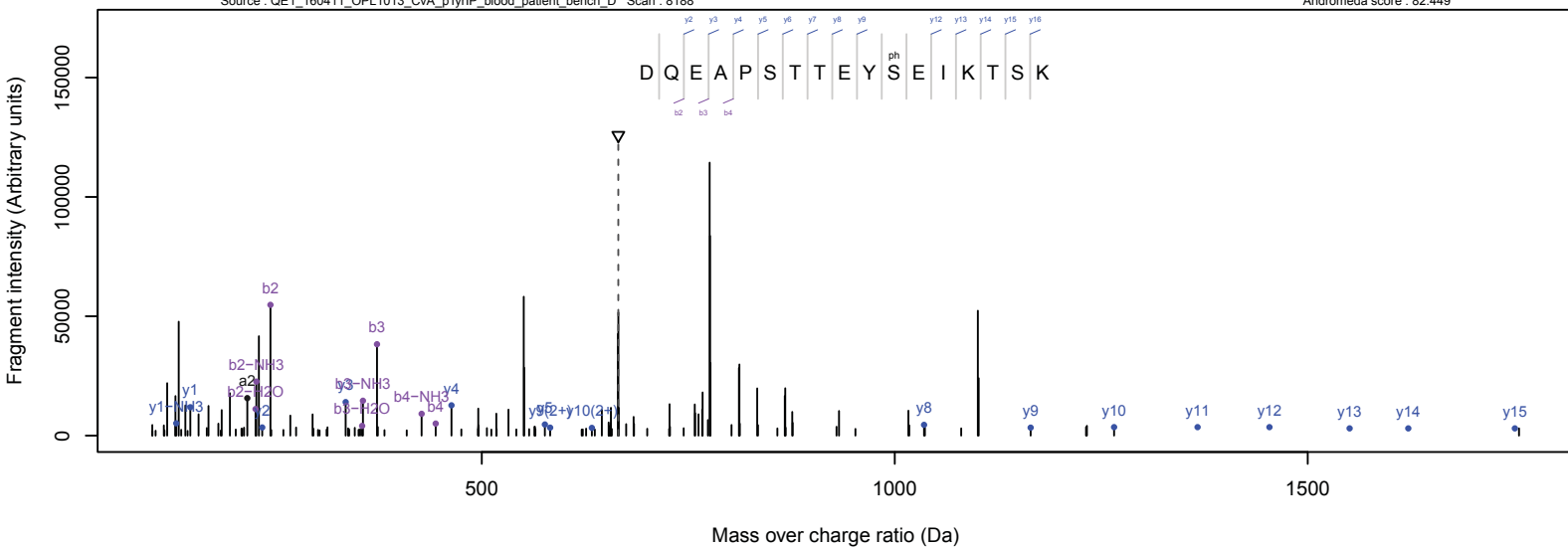
Andromeda score: 82.651



**DQEAPSTTEYS(ph)EIKTSK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 8188

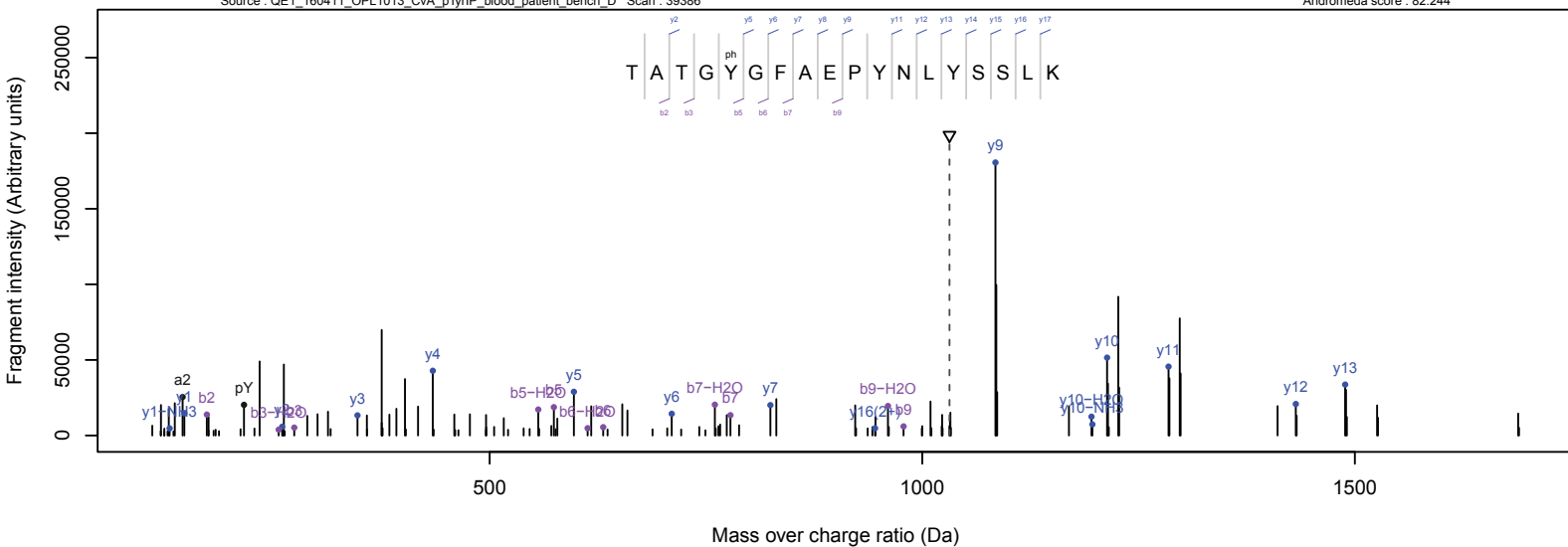
Andromeda score : 82.449



**TATGY(ph)GFAEPYNLYSSLK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 39386

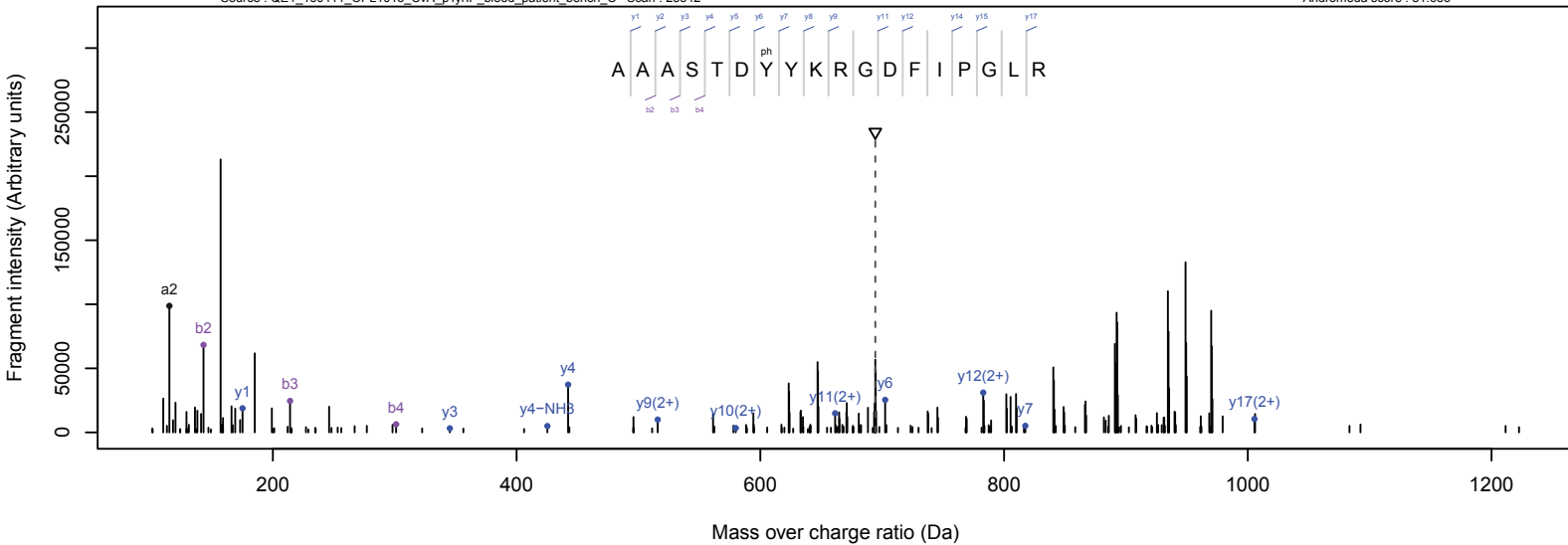
Andromeda score : 82.244



**AAASTDY(ph)YKRGDFIPGLR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 25542

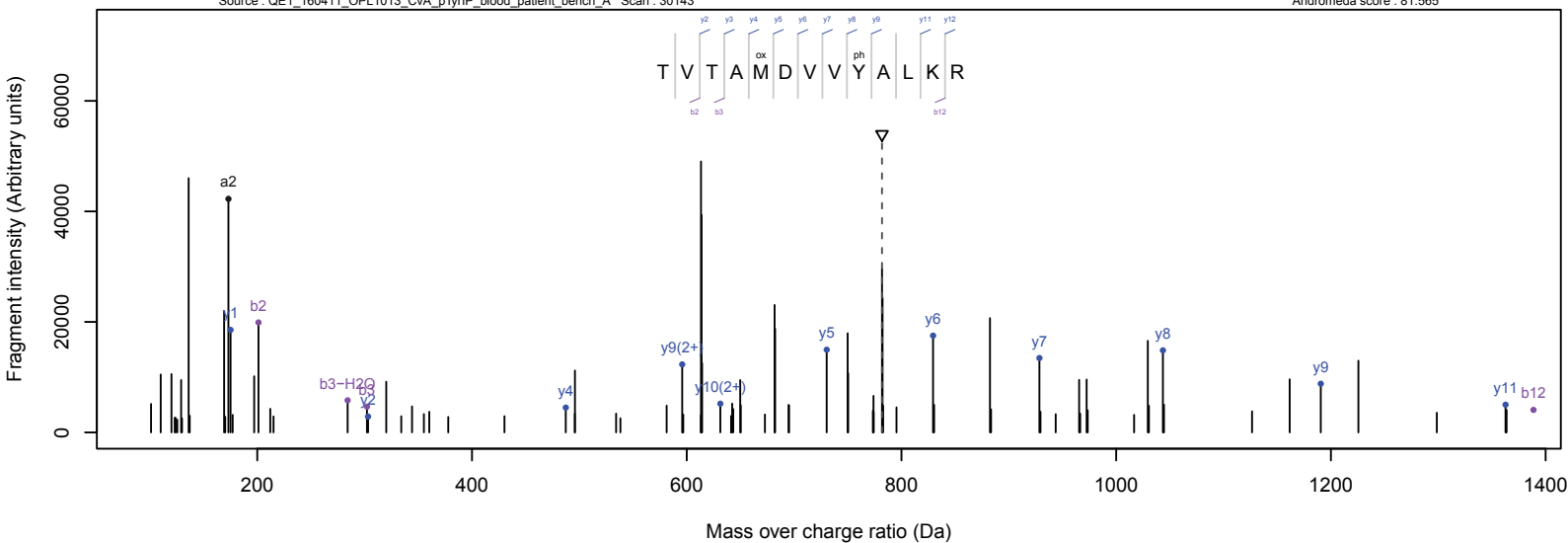
Andromeda score : 81.656



TVTAM(ox)DVVY(ph)ALKR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 30143

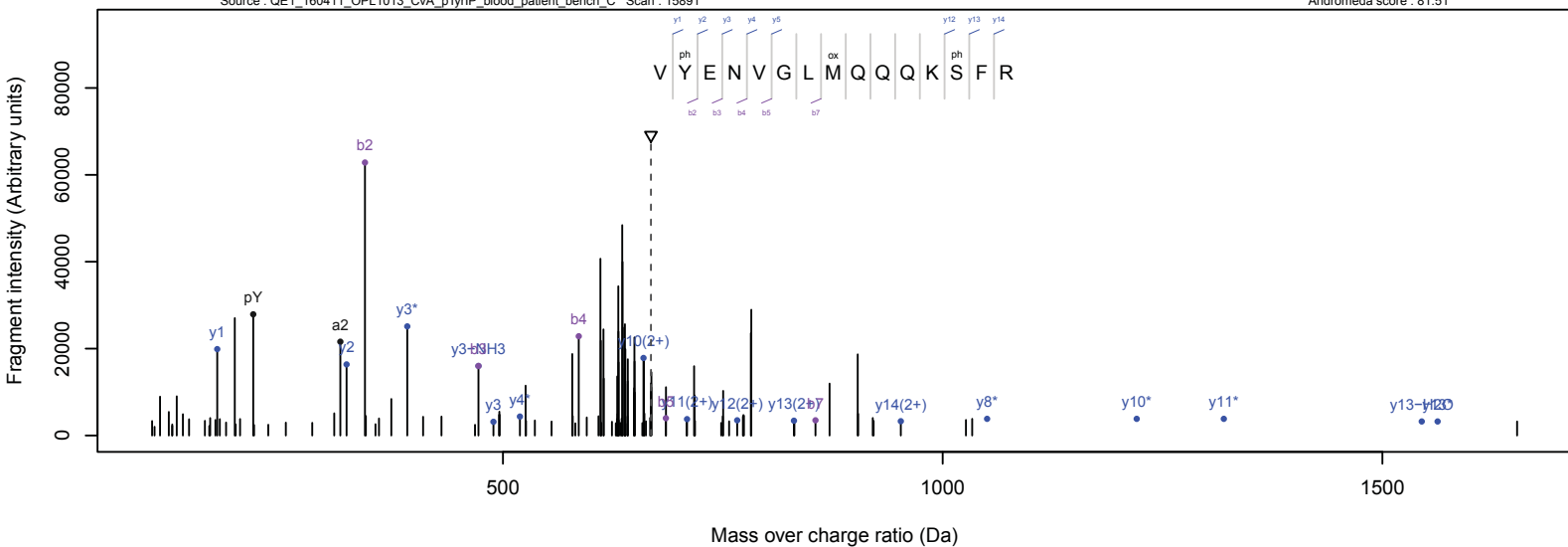
Andromeda score : 81.565



VY(ph)ENVGLM(ox)QQQS(ph)FR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 15891

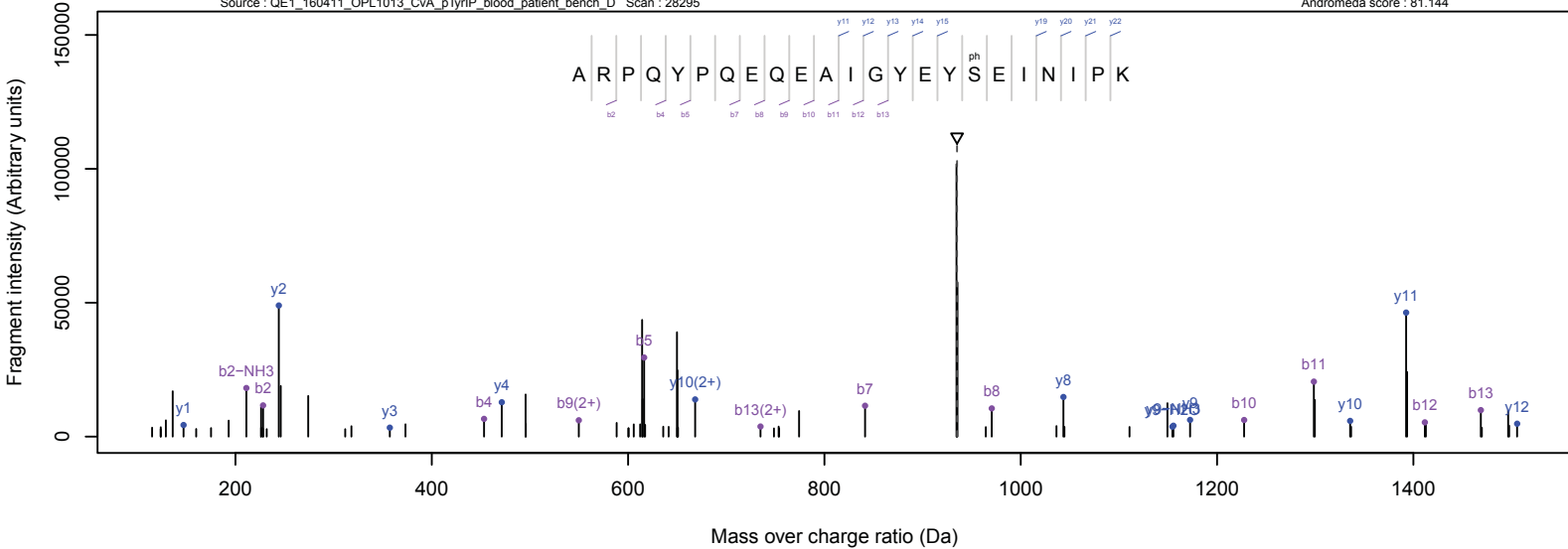
Andromeda score : 81.51



ARPQYPQEQAIGYEYS(ph)EINIPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28295

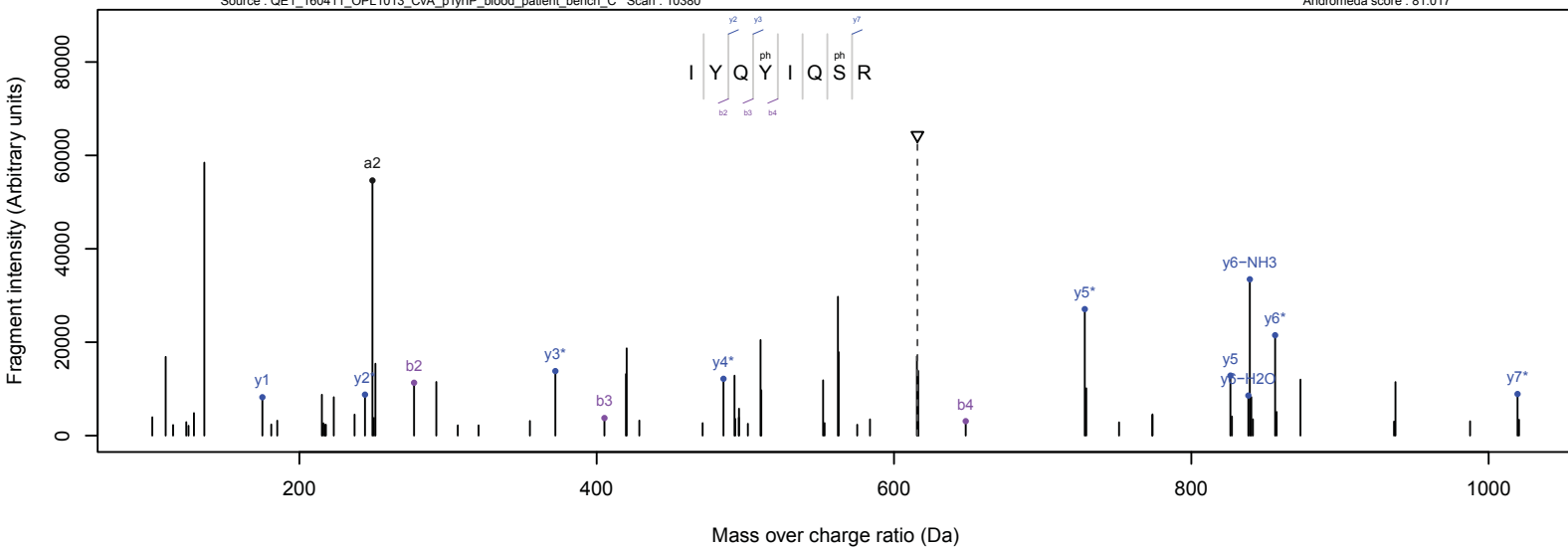
Andromeda score : 81.144



## IYQY(ph)IQS(ph)R-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 10380

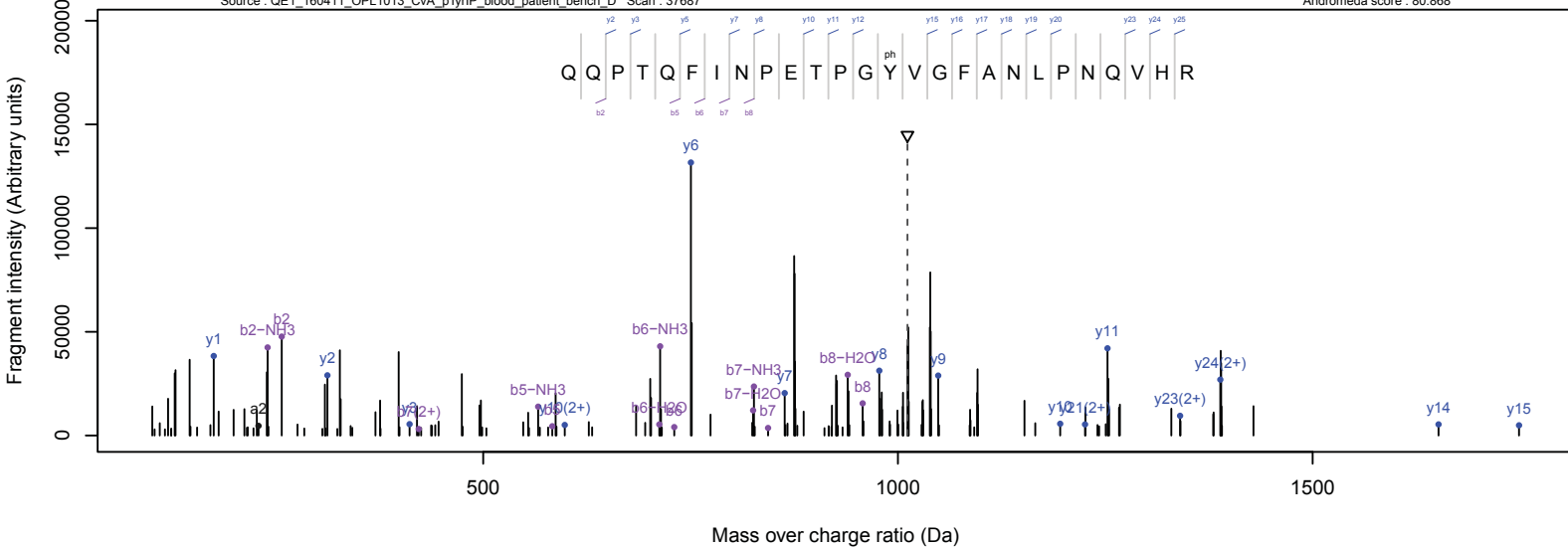
Andromeda score : 81.017



## QQPTQFINPETPGYVGFANLPNQVHR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 37687

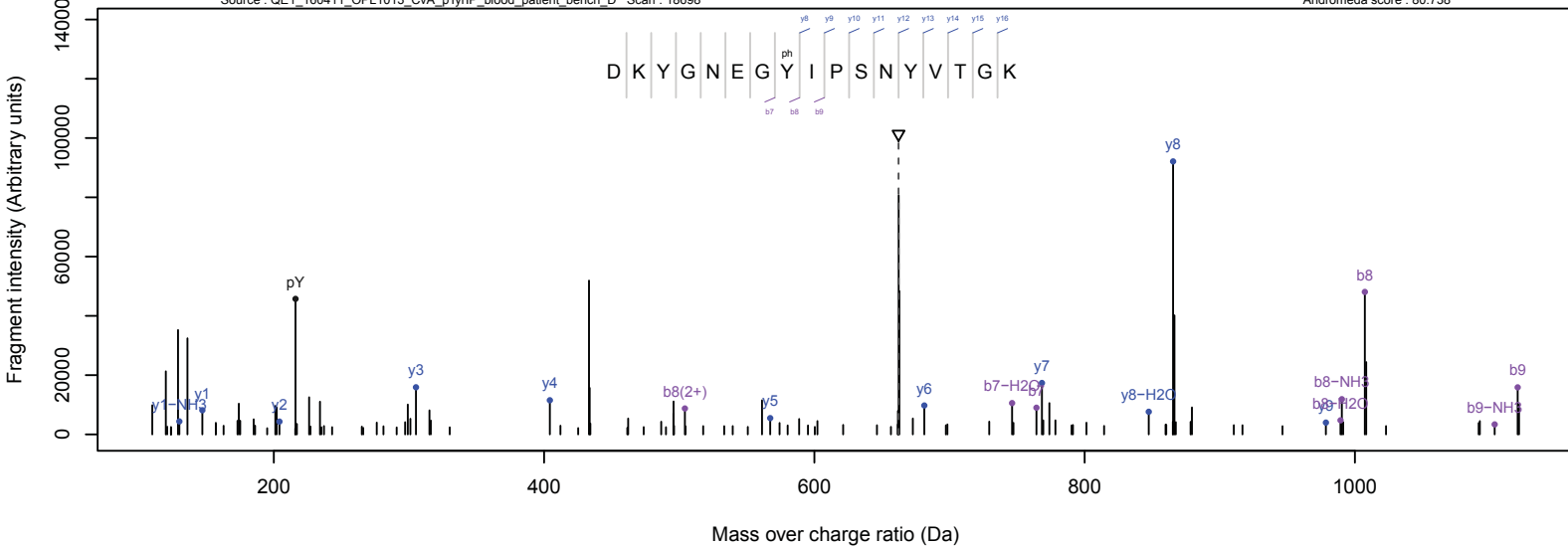
Andromeda score : 80.868



## DKYGENEYI PSNYVTGK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 18698

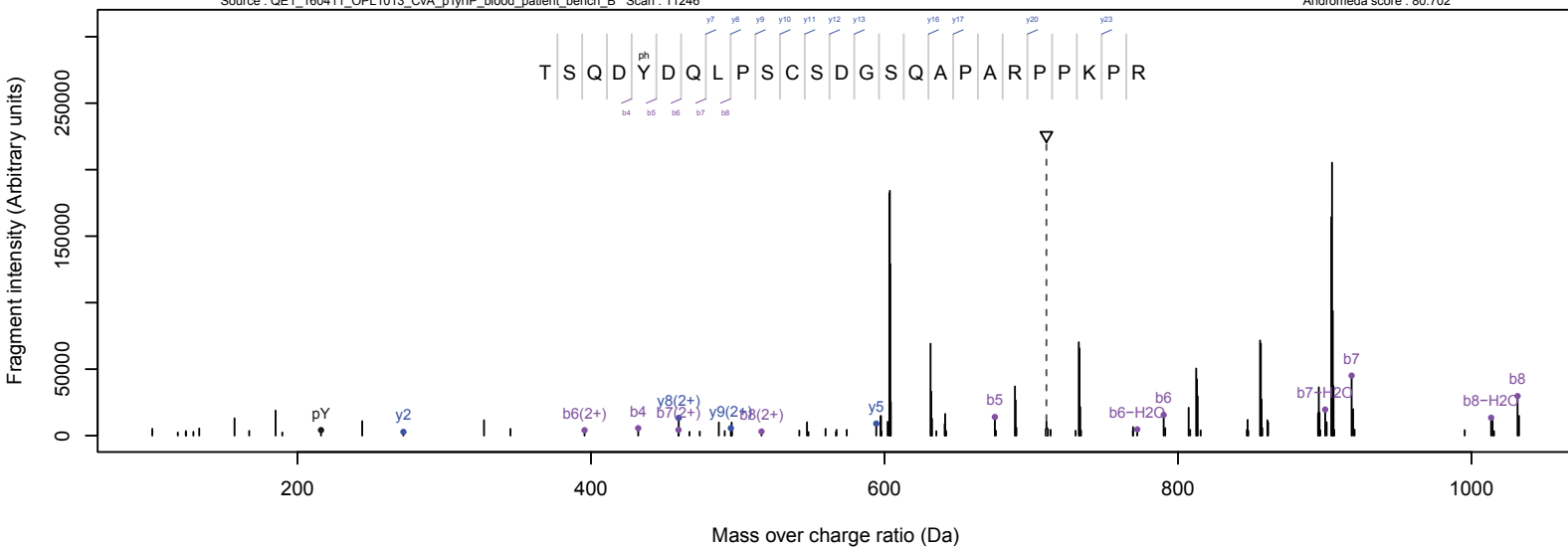
Andromeda score : 80.738



## TSQDY(ph)DQLPSCDGSQAPARPPKPR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 11246

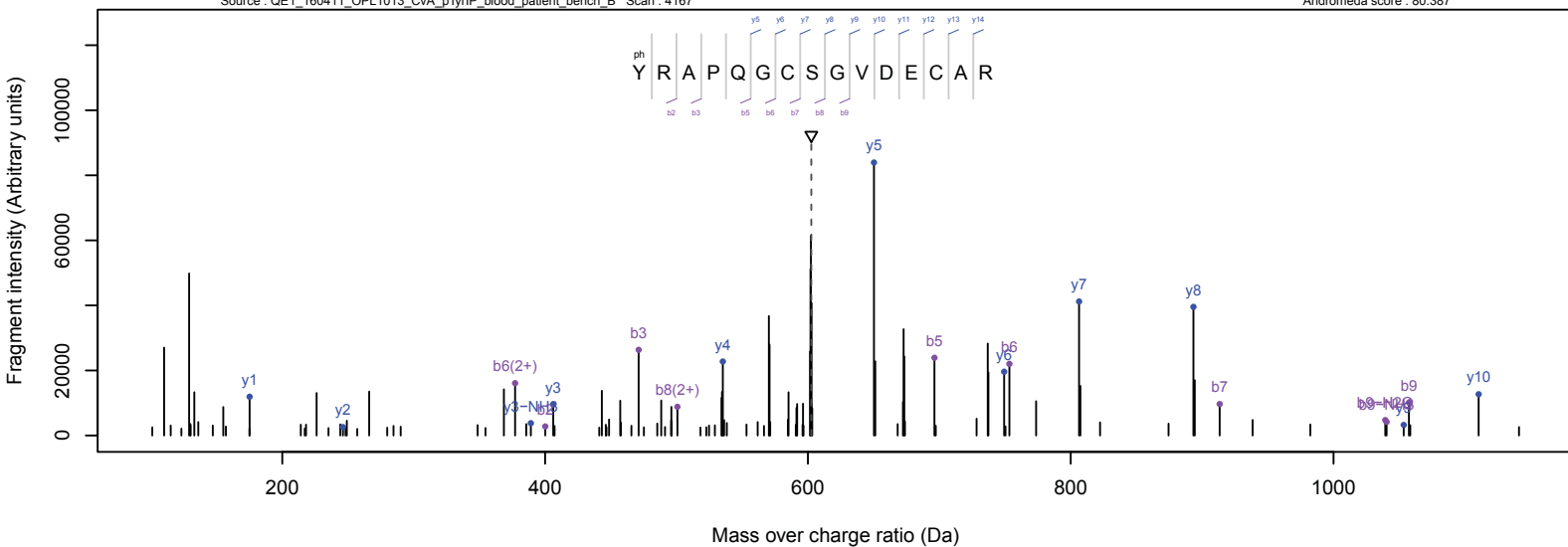
Andromeda score : 80.702



## Y(ph)RAPQGCSGVDECAR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 4167

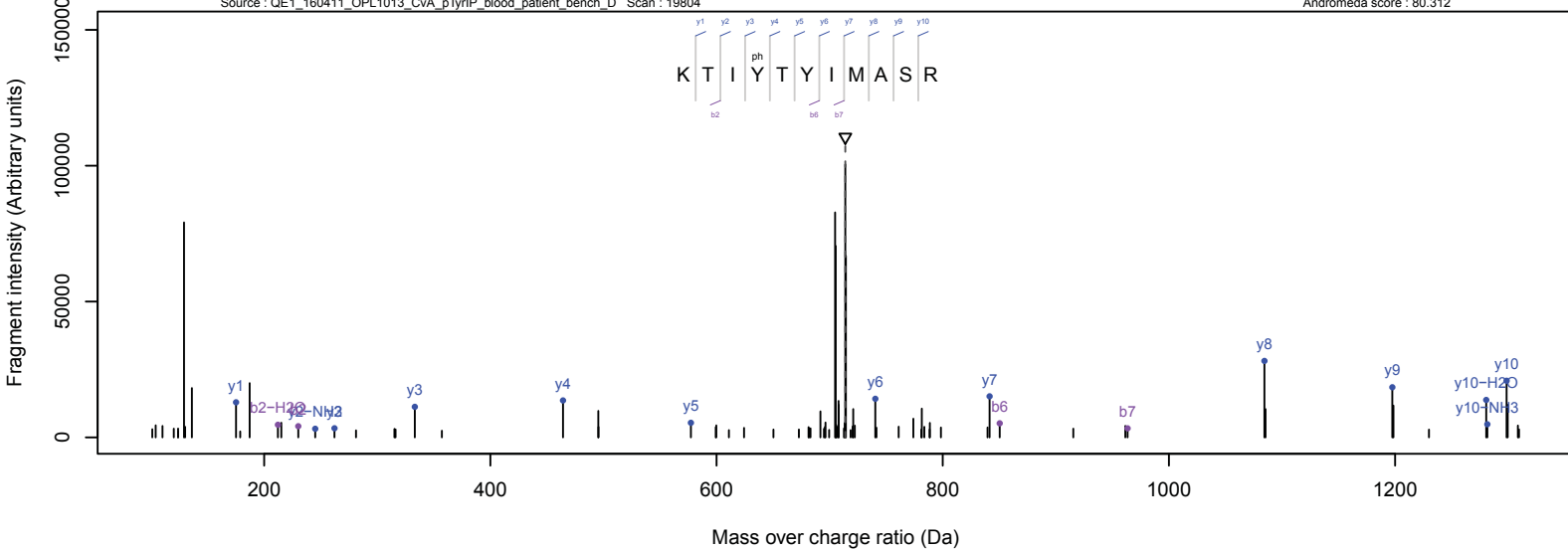
Andromeda score : 80.387



## KTIY(ph)TYIMASR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 19804

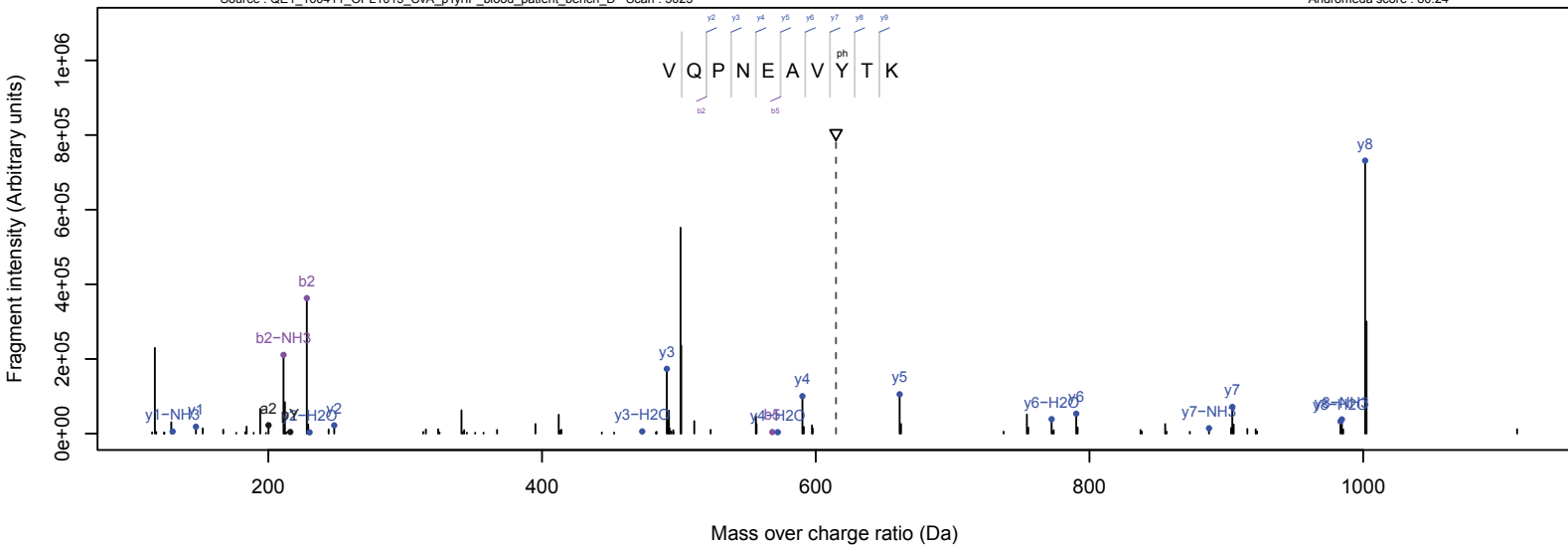
Andromeda score : 80.312



VQPNEAVYTK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 5025

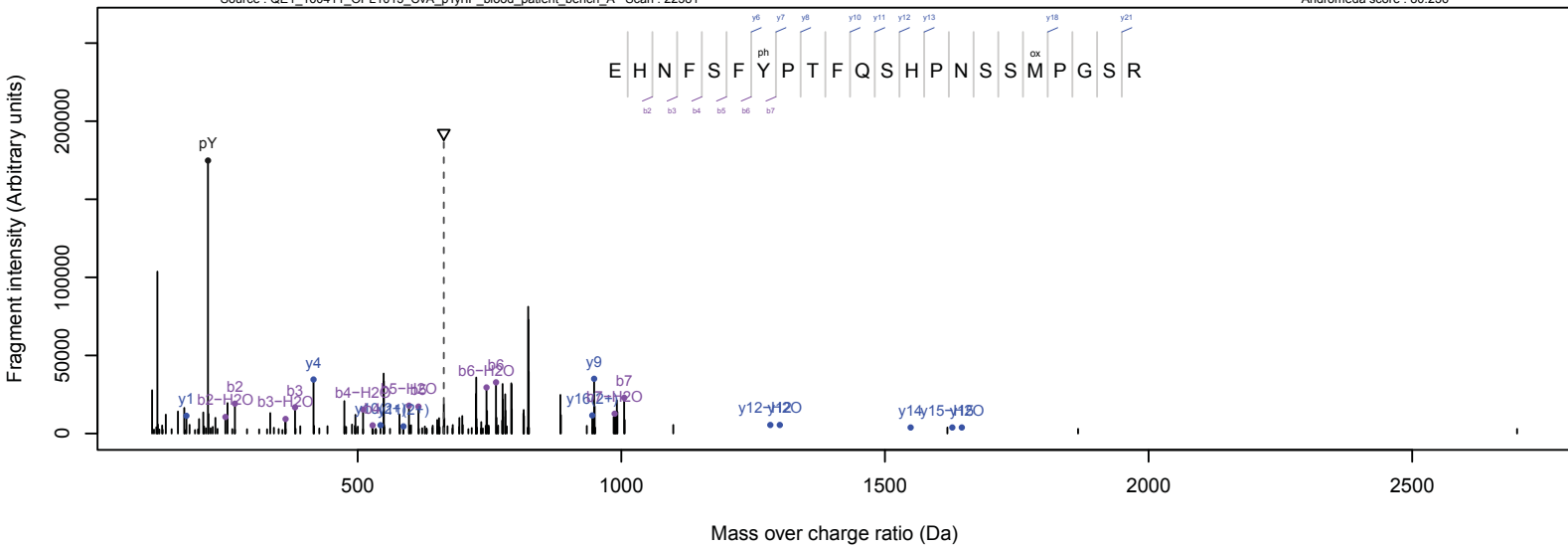
Andromeda score: 80.24



EHNFSFYPTFQSHPNSSMPGSR-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 22381

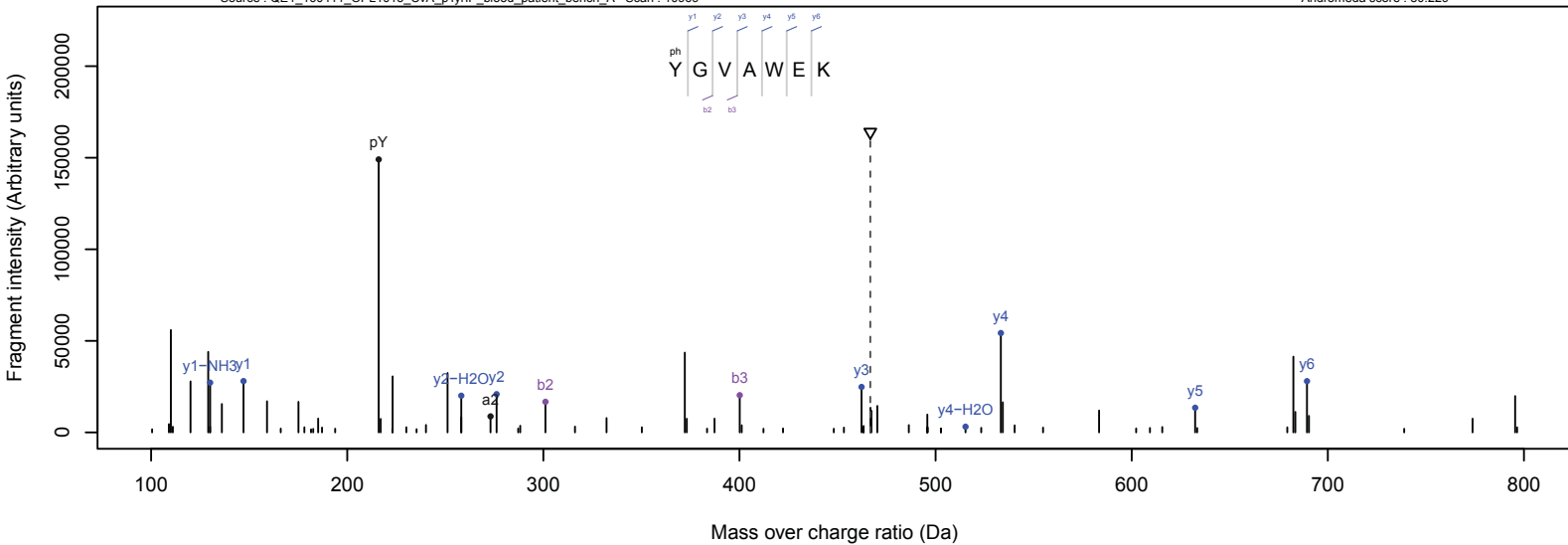
Andromeda score: 80.236



Y(ph)GVAWEK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 10969

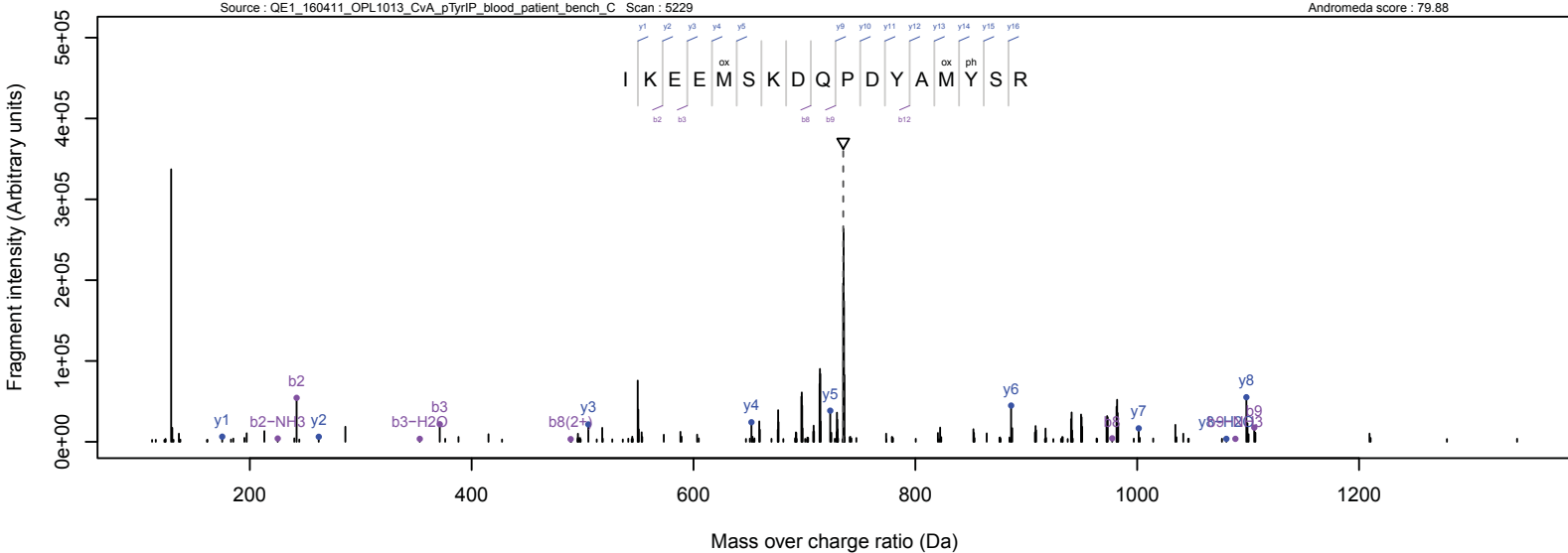
Andromeda score: 80.229



## IKEEM(ox)SKDQPDYAM(ox)Y(ph)SR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 5229

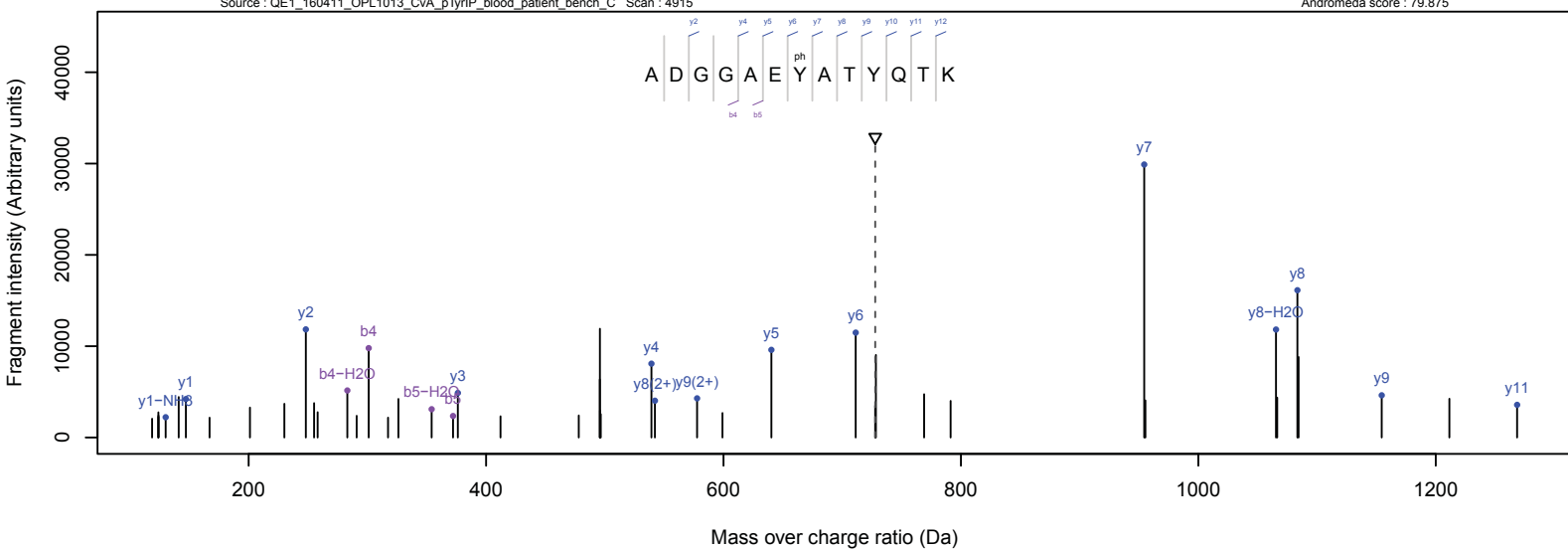
Andromeda score : 79.88



## ADGGAEY(ph)ATYQTK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 4915

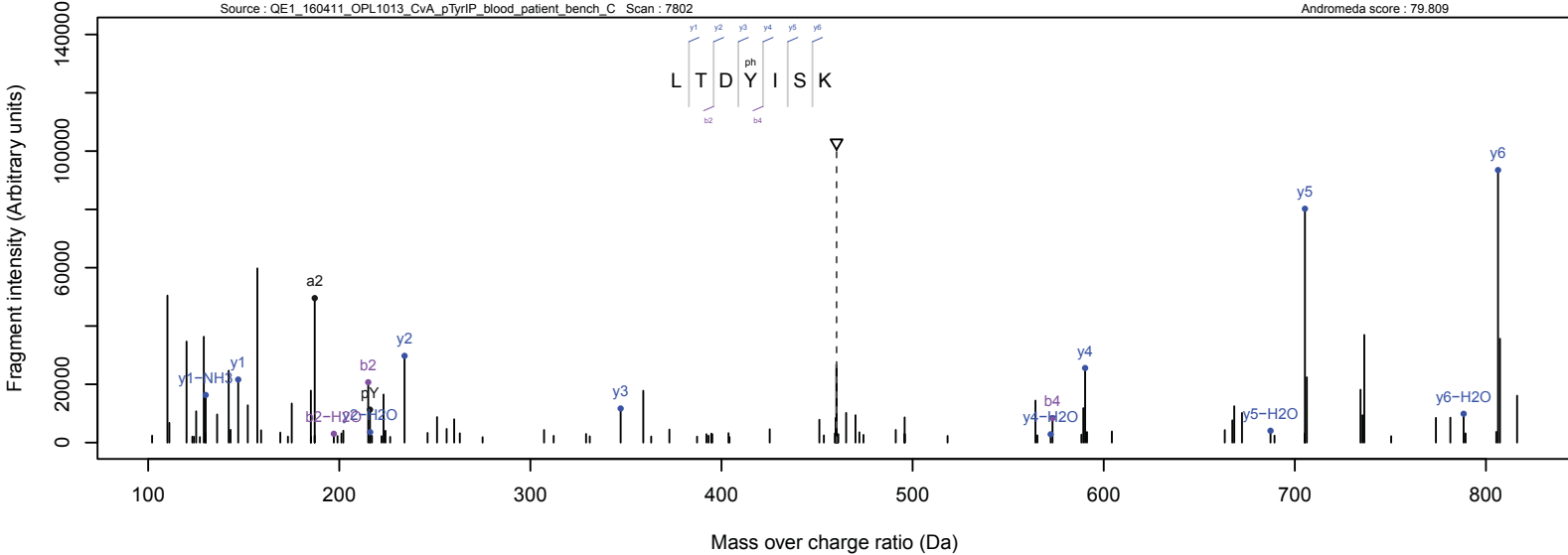
Andromeda score : 79.875



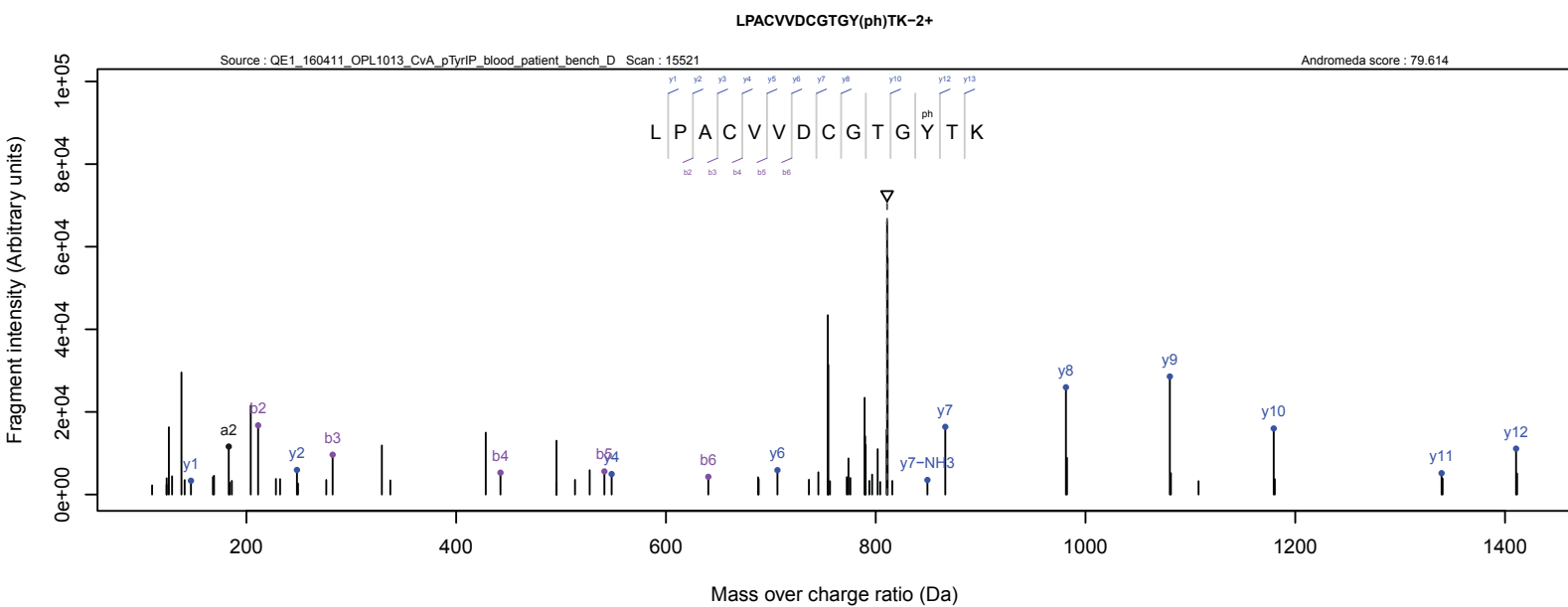
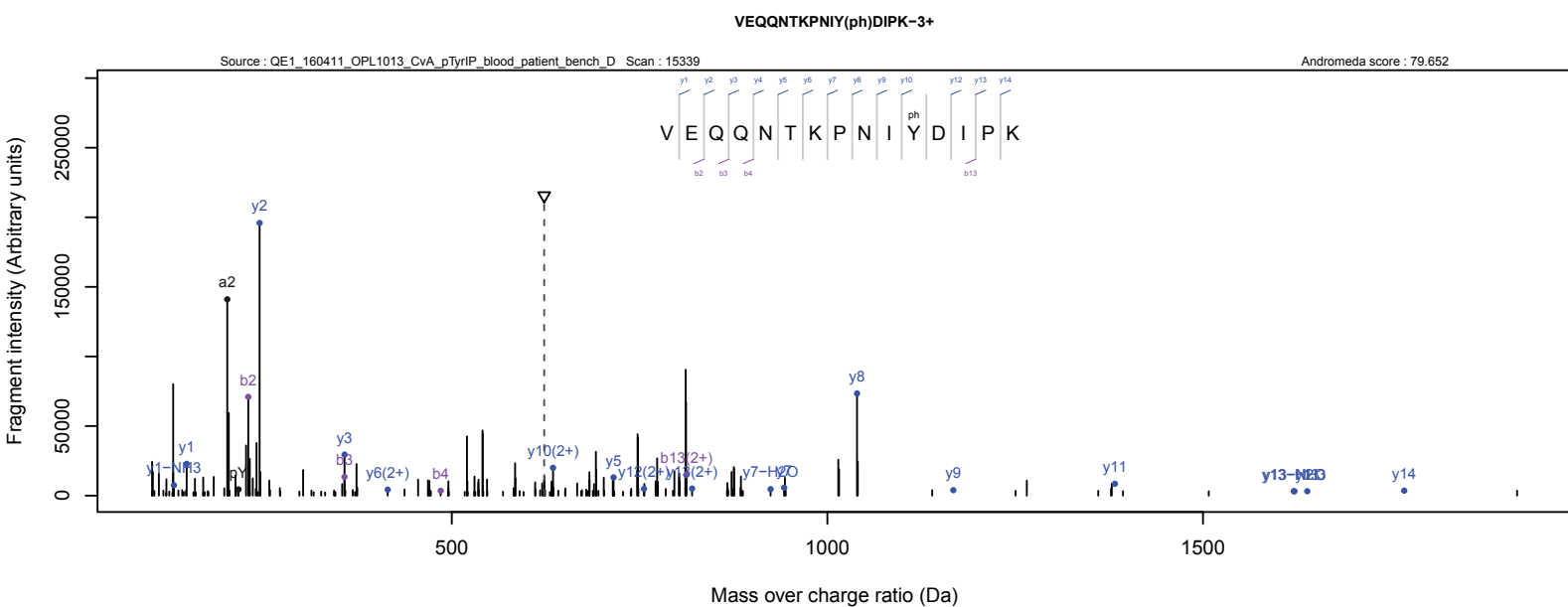
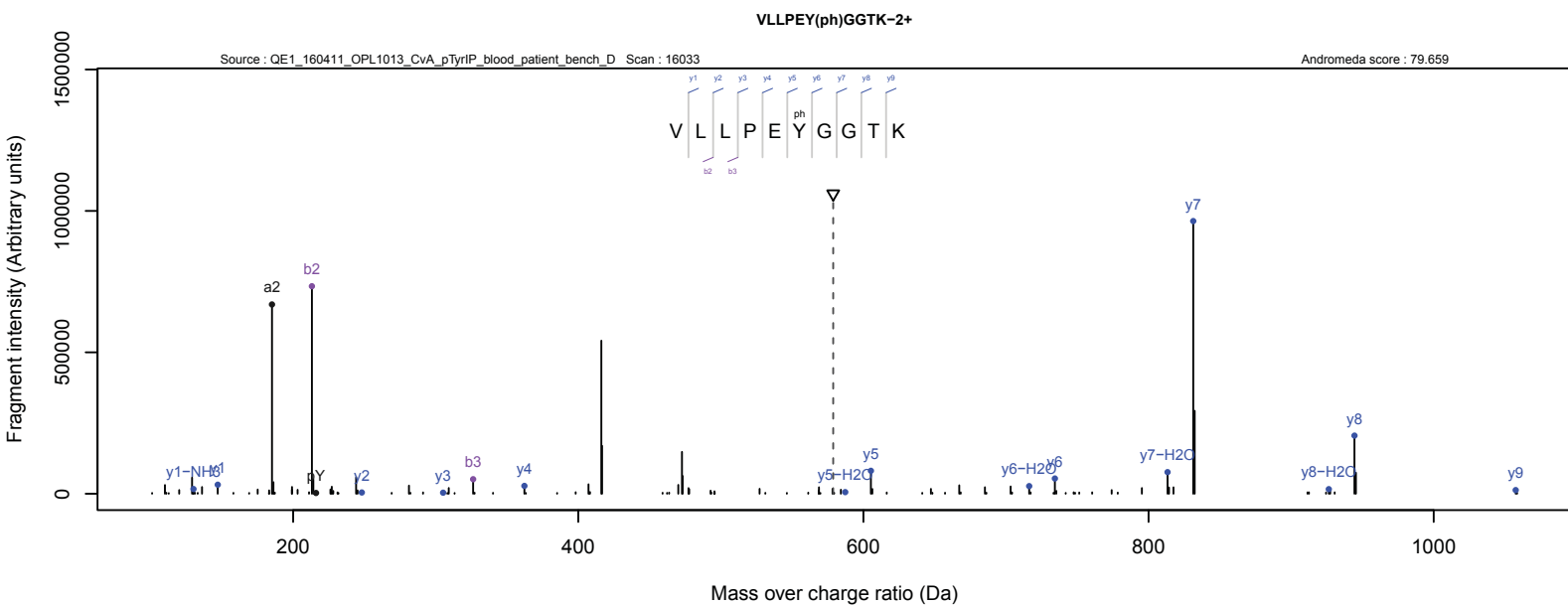
## LTDY(ph)ISK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 7802

Andromeda score : 79.809



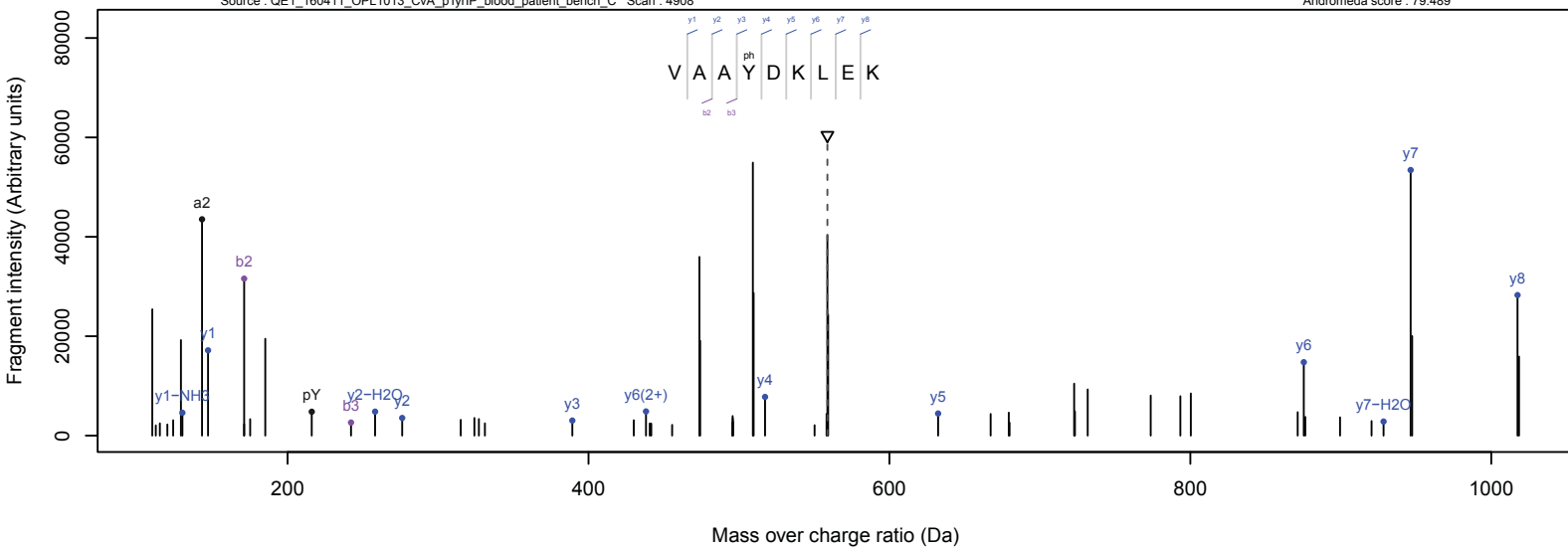




VAAY(ph)DKLEK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 4908

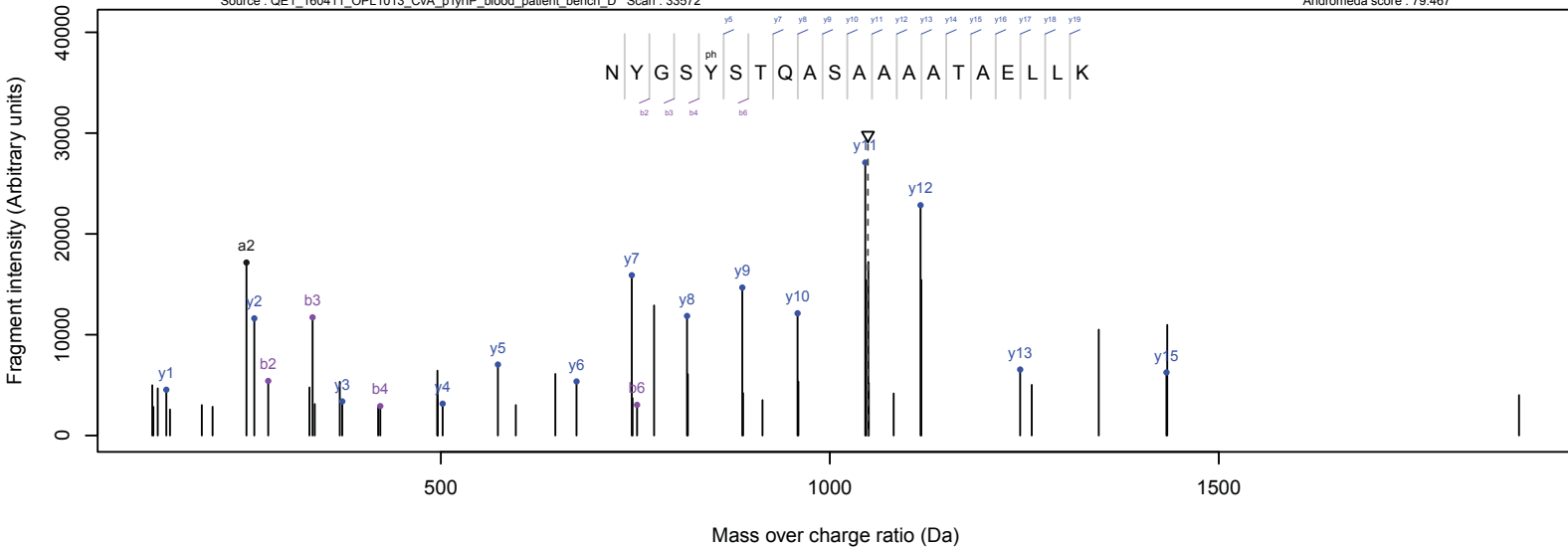
Andromeda score : 79.489



NYGSY(ph)STQASAAAATAELLK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 33572

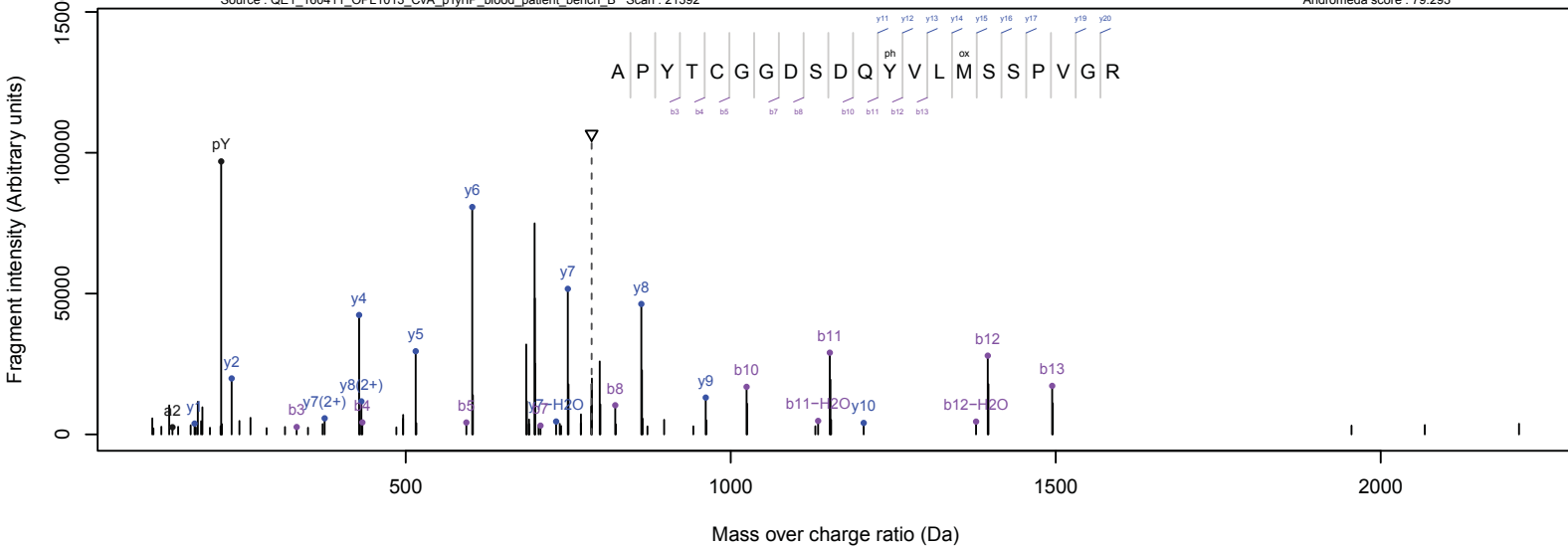
Andromeda score : 79.467

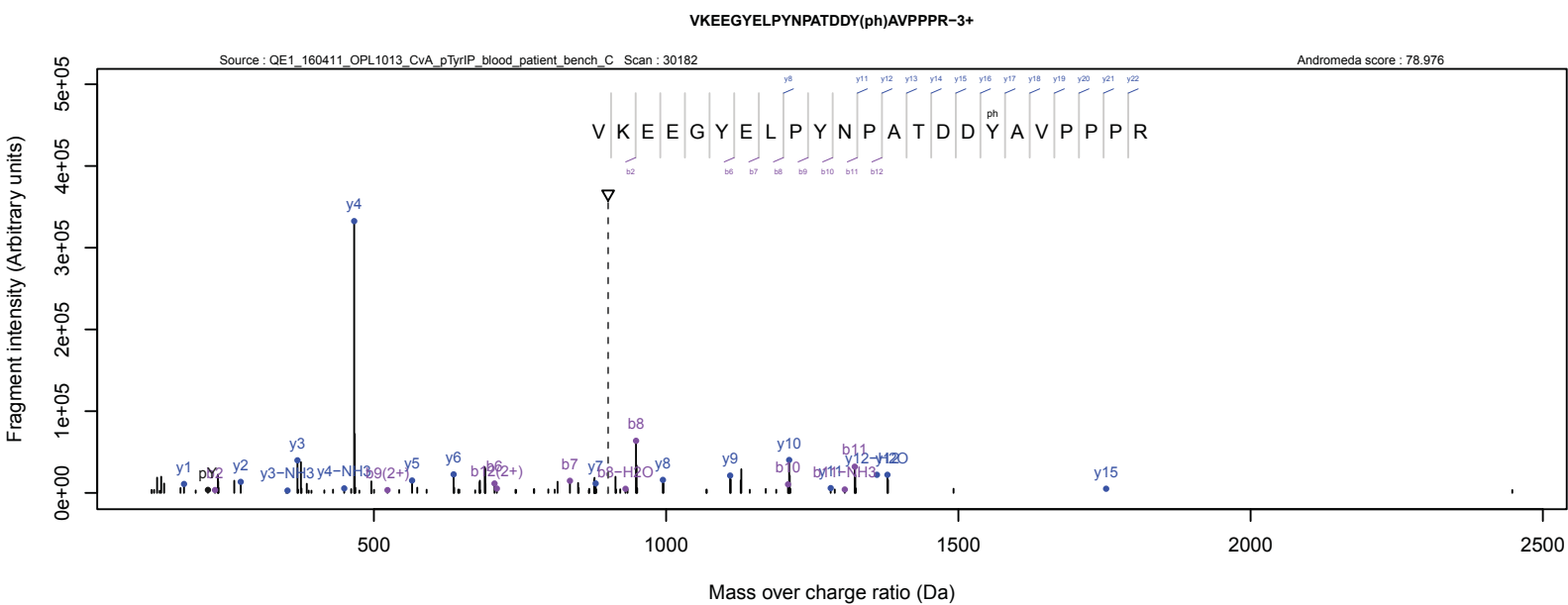
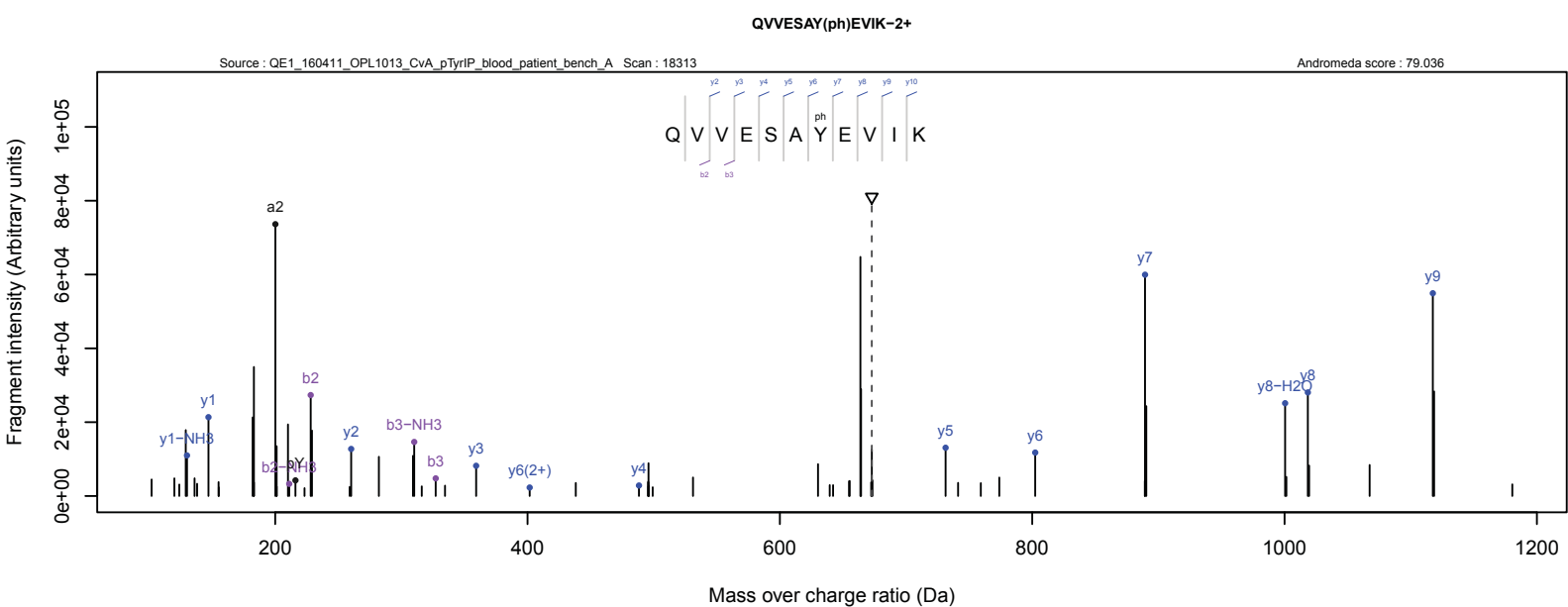
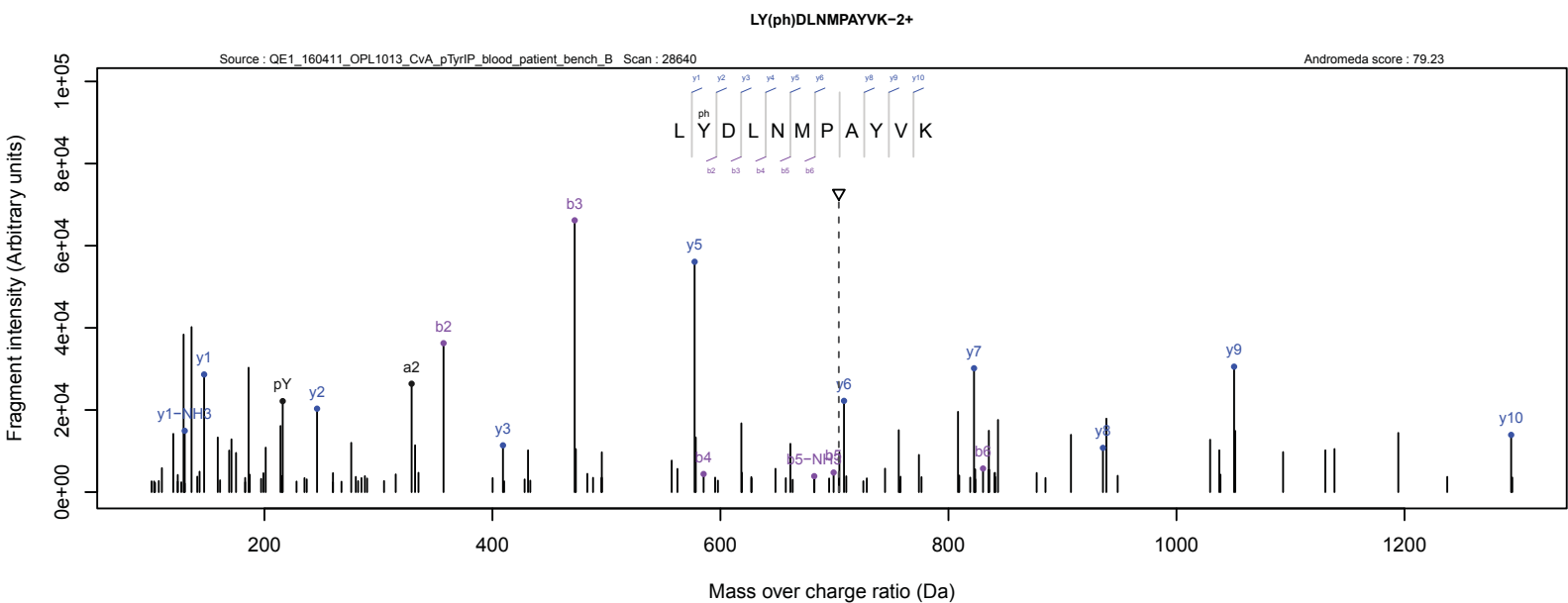


APYTCGGDSDQY(ph)VLMSSPVGGR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 21392

Andromeda score : 79.293

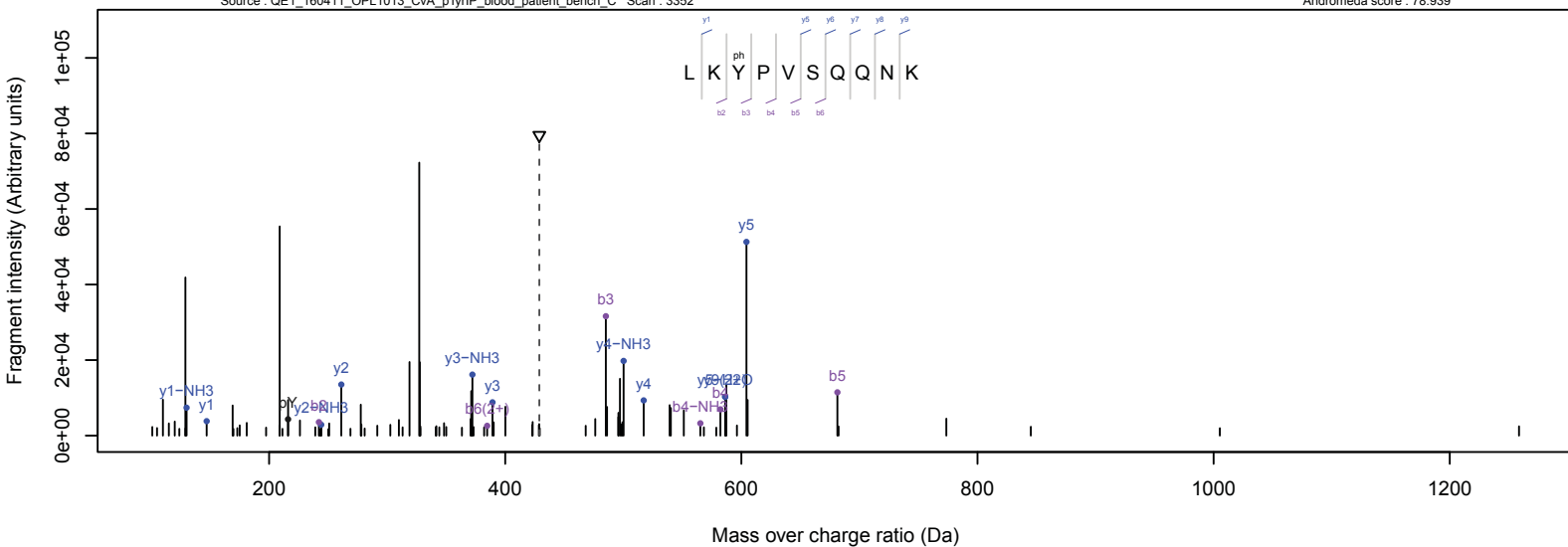




LKY(ph)PVSQQNK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3352

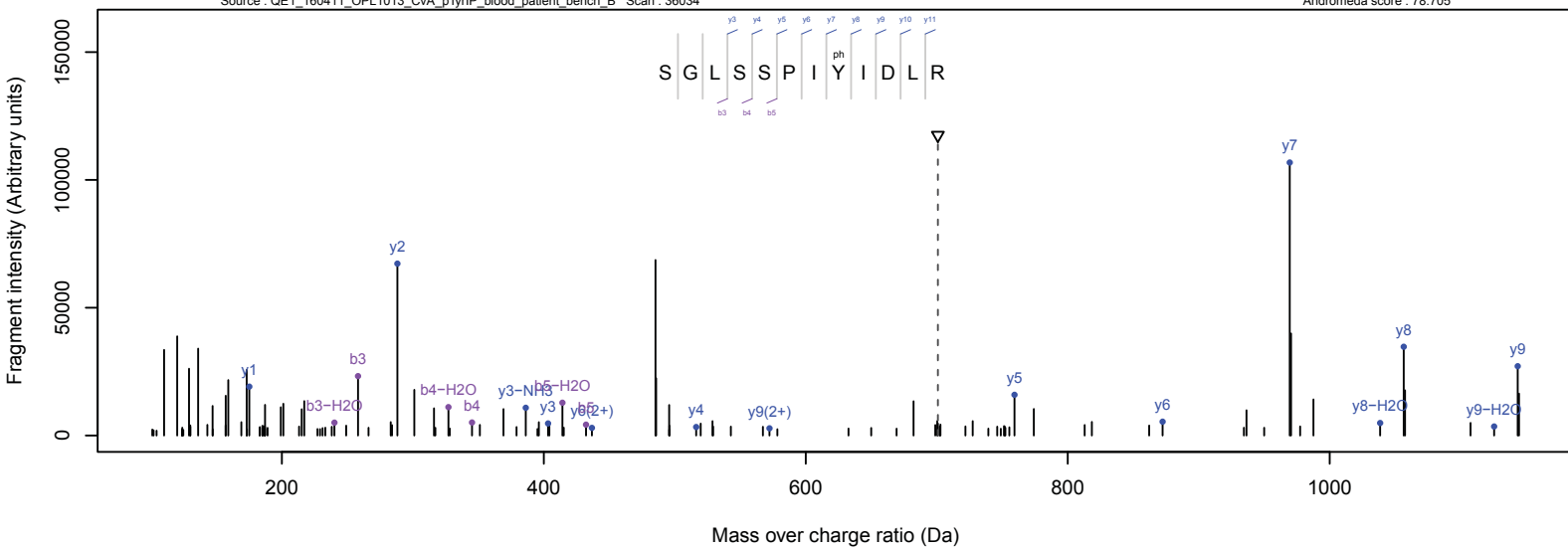
Andromeda score : 78.939



SGLSSPIY>IDLR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 36034

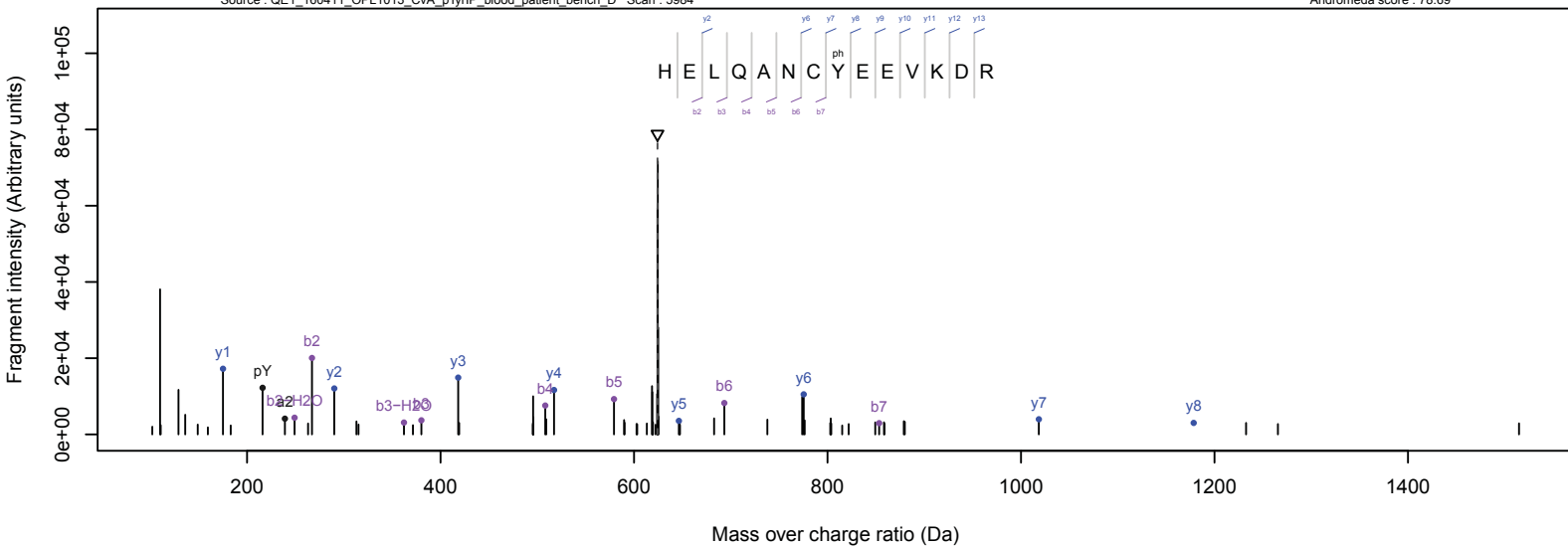
Andromeda score : 78.705



HELQANCY(ph)EEVKDR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5984

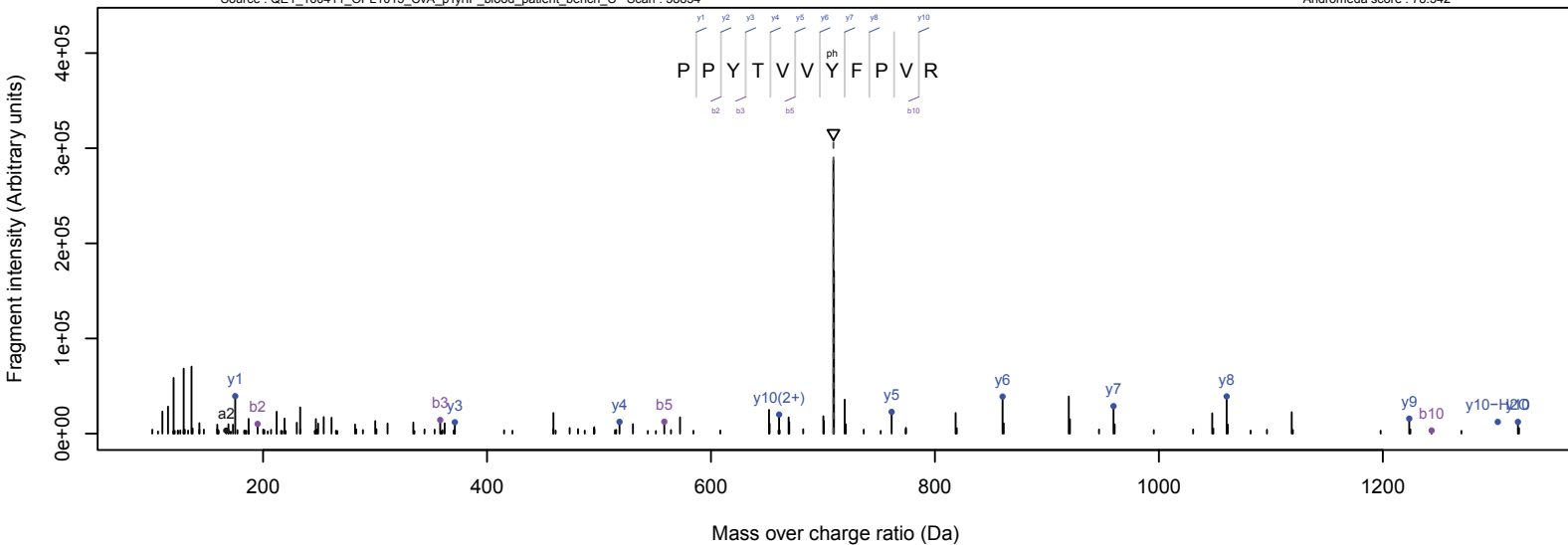
Andromeda score : 78.69



PPYTVVY(ph)FPVR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 38834

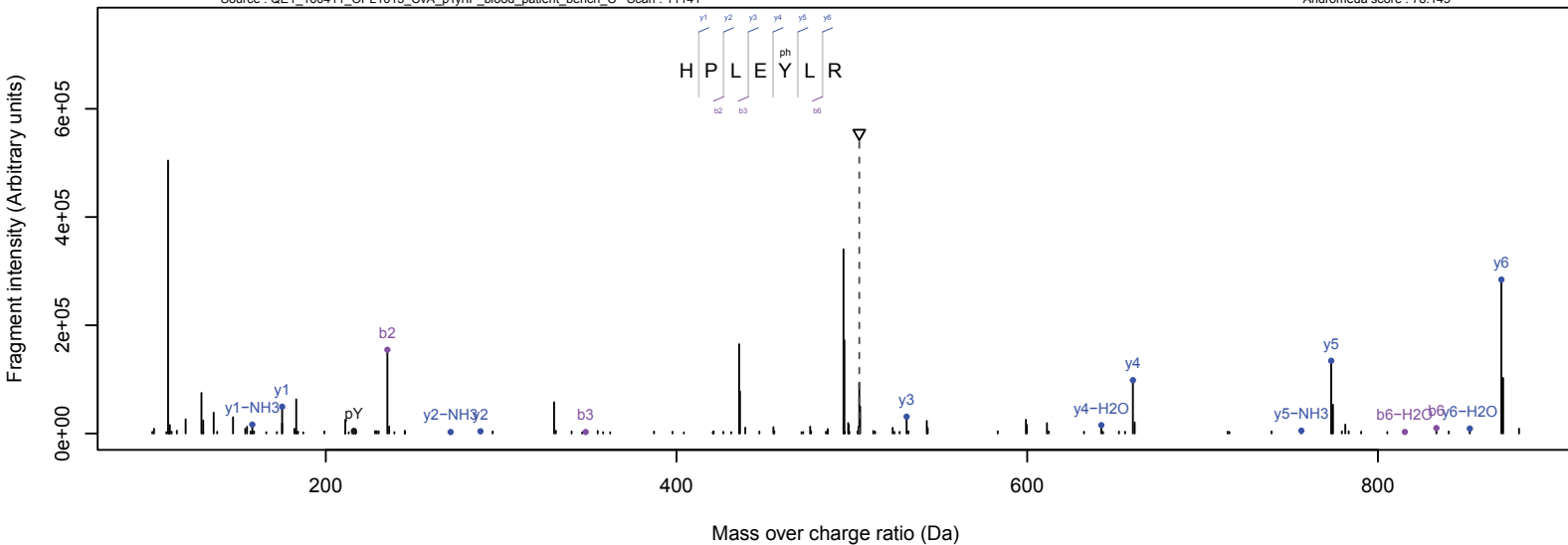
Andromeda score : 78.342



HPLEY(ph)LR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 11141

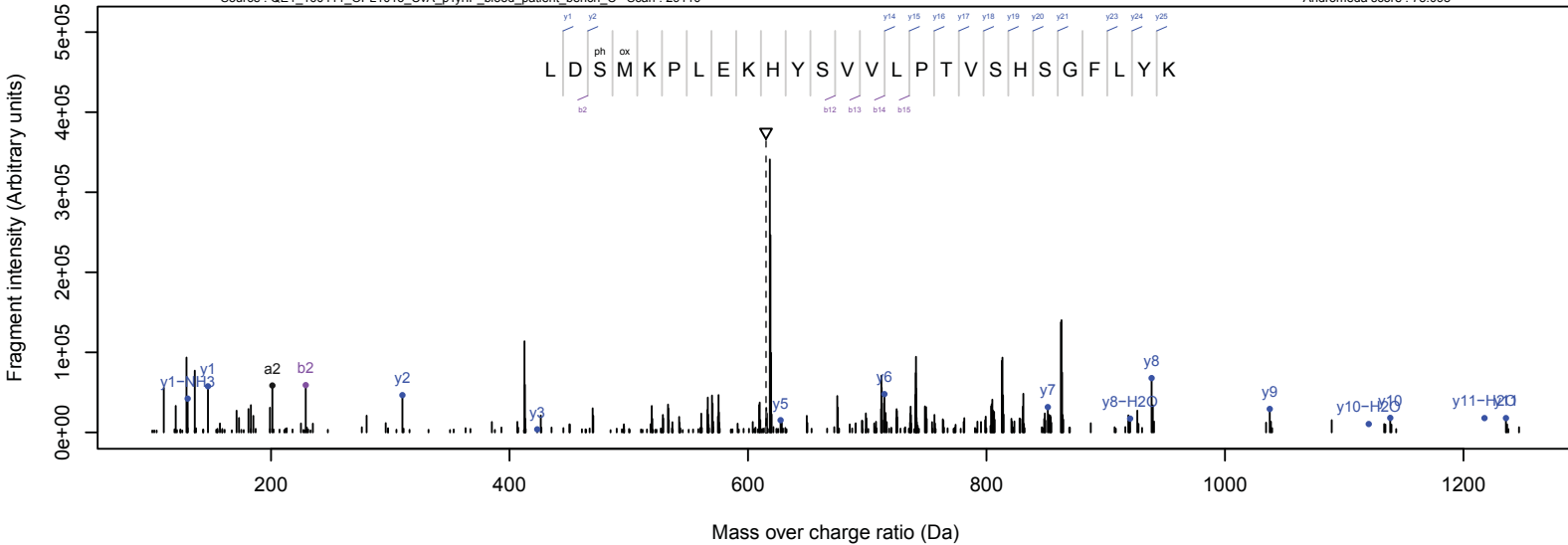
Andromeda score : 78.149



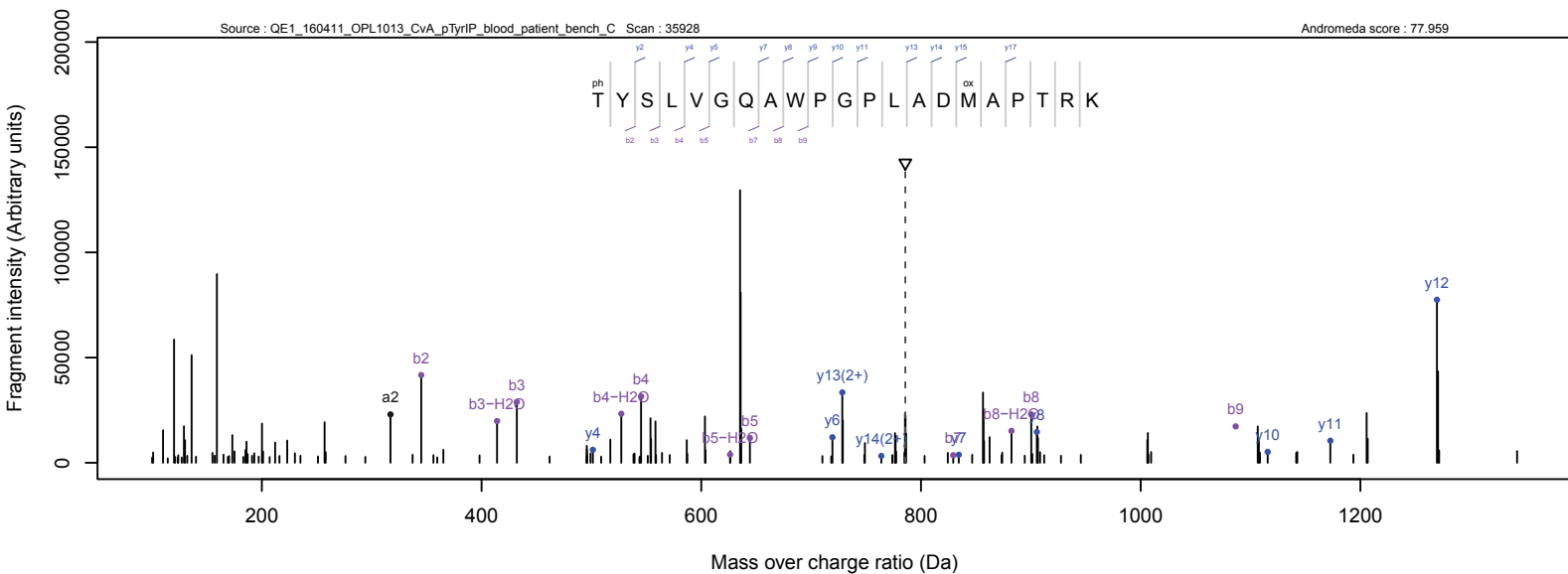
LDS(ph)M(ox)KPLEKHYSVVLPTVSHSGFLYK-5+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 29110

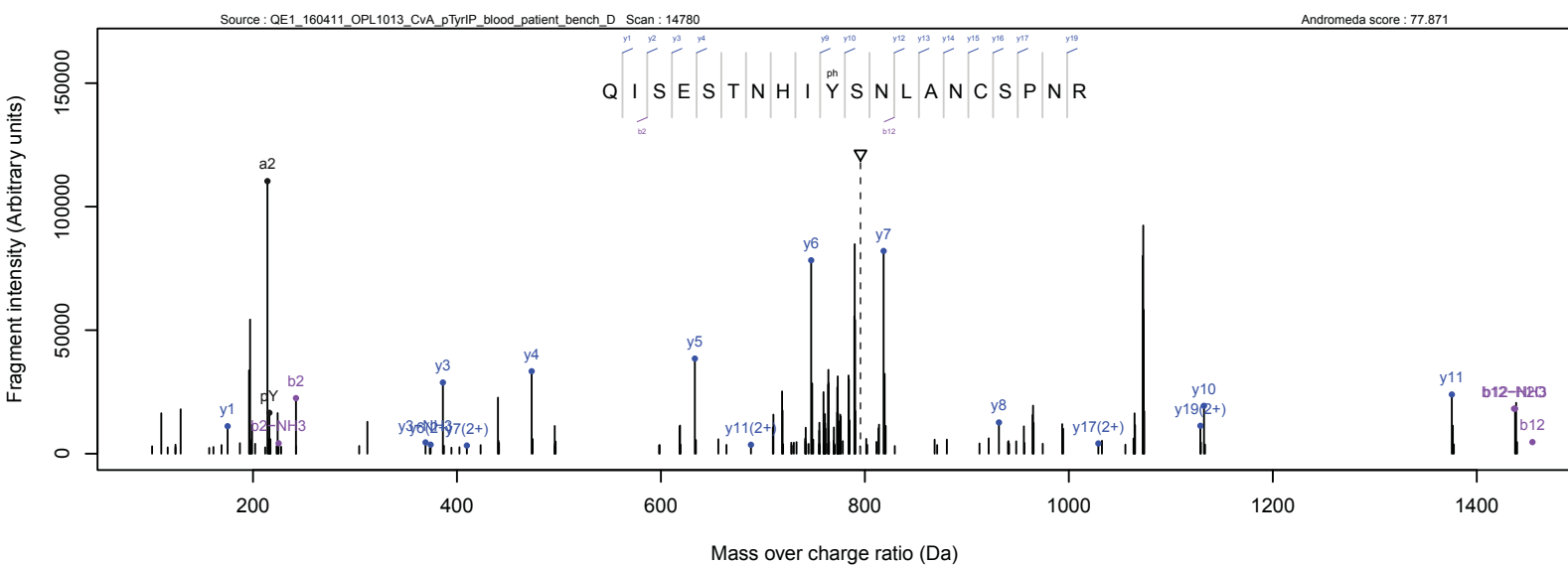
Andromeda score : 78.098



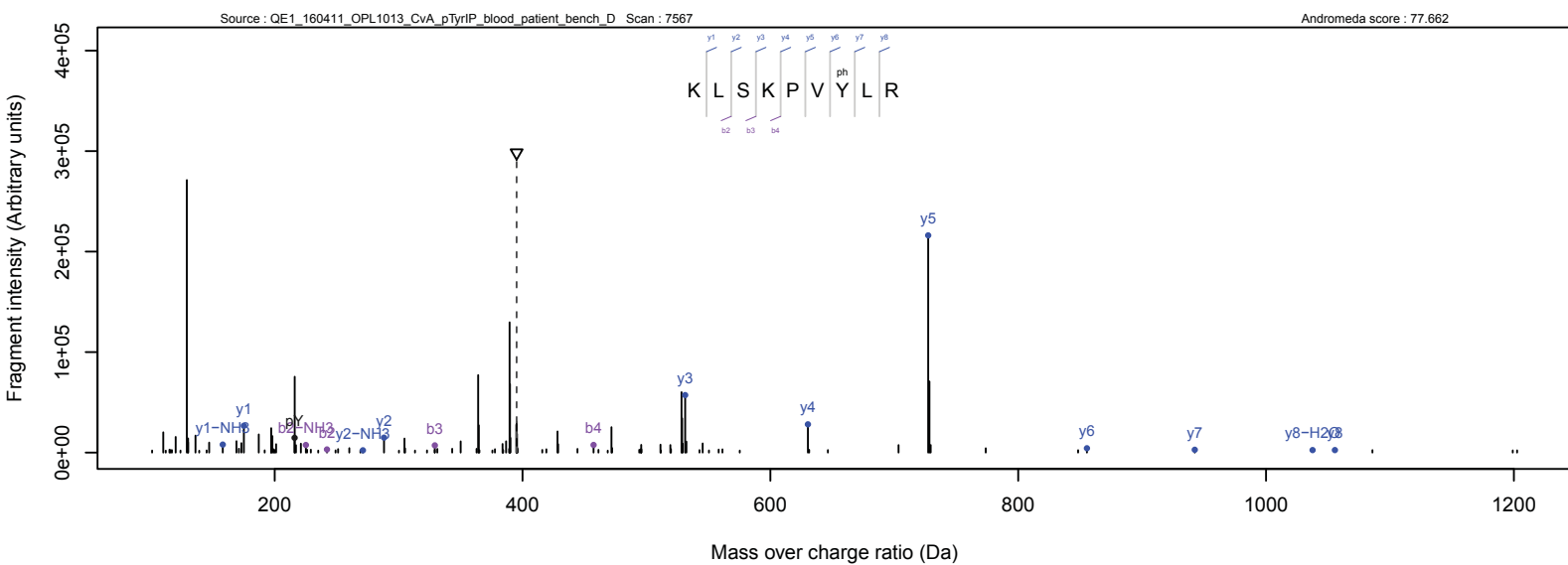
T(ph)YSLVGQAWPGPLADM(ox)APTRK-3+



QISESTNHIY(ph)SNLANCSPNR-3+



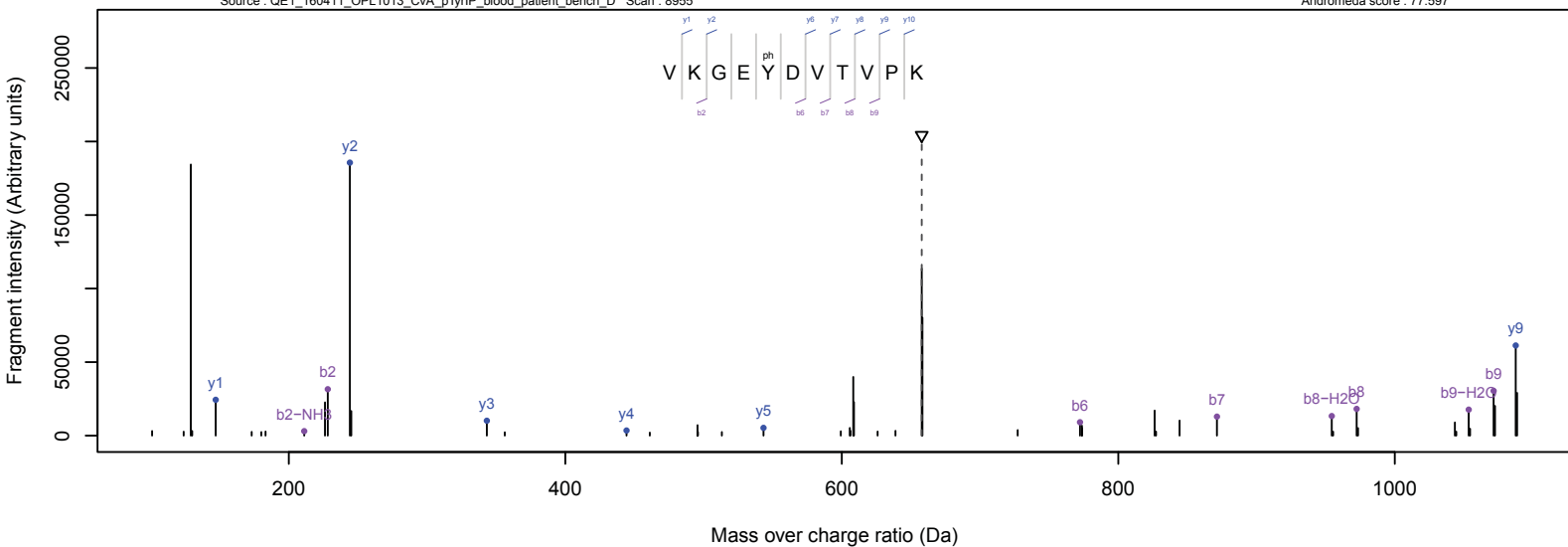
KLSKPVY(ph)LR-3+



VKGEY(ph)DVTVPK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 8955

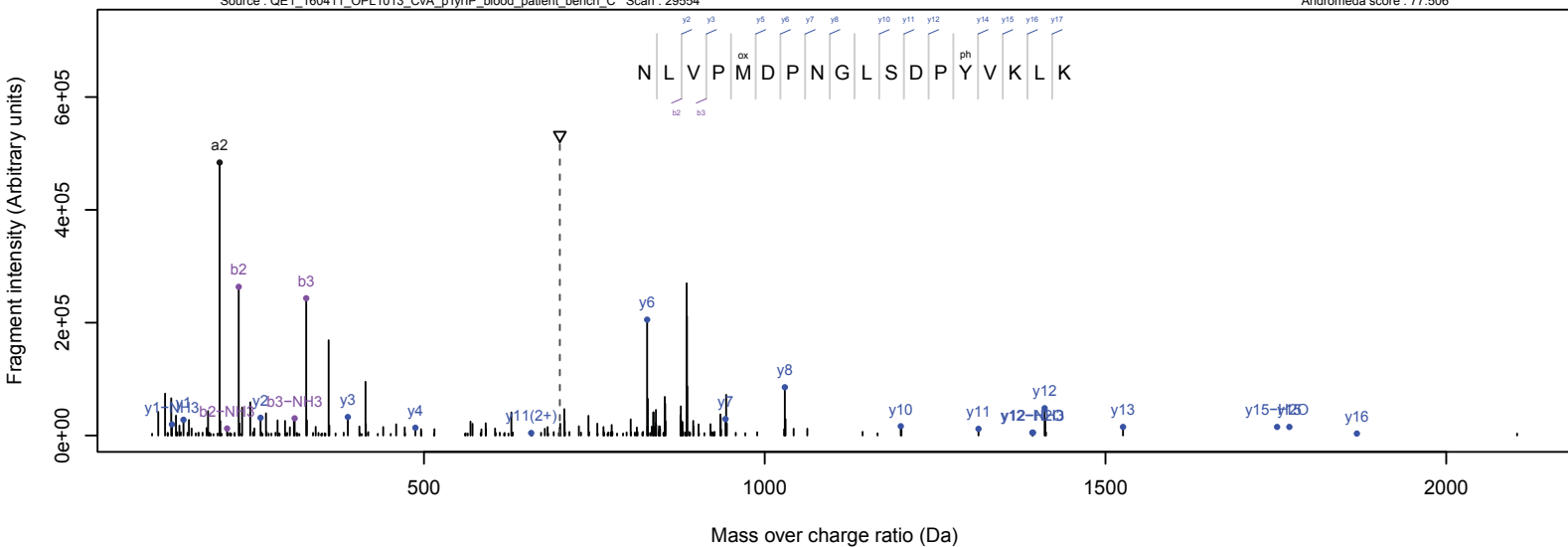
Andromeda score : 77.597



NLVPM(ox)DPNGLSDPY(ph)VKLK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 29554

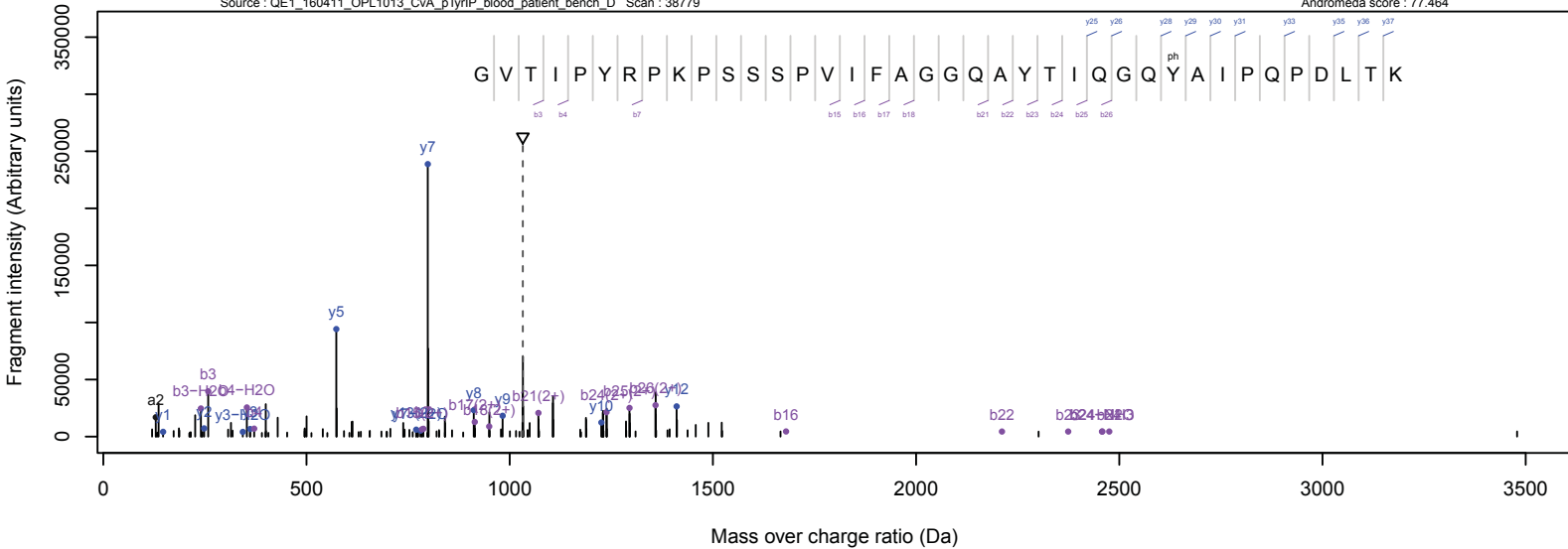
Andromeda score : 77.506



GVTIYRKPSSSPVIFAGGQAYTIQQGYAIPQDLTK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 38779

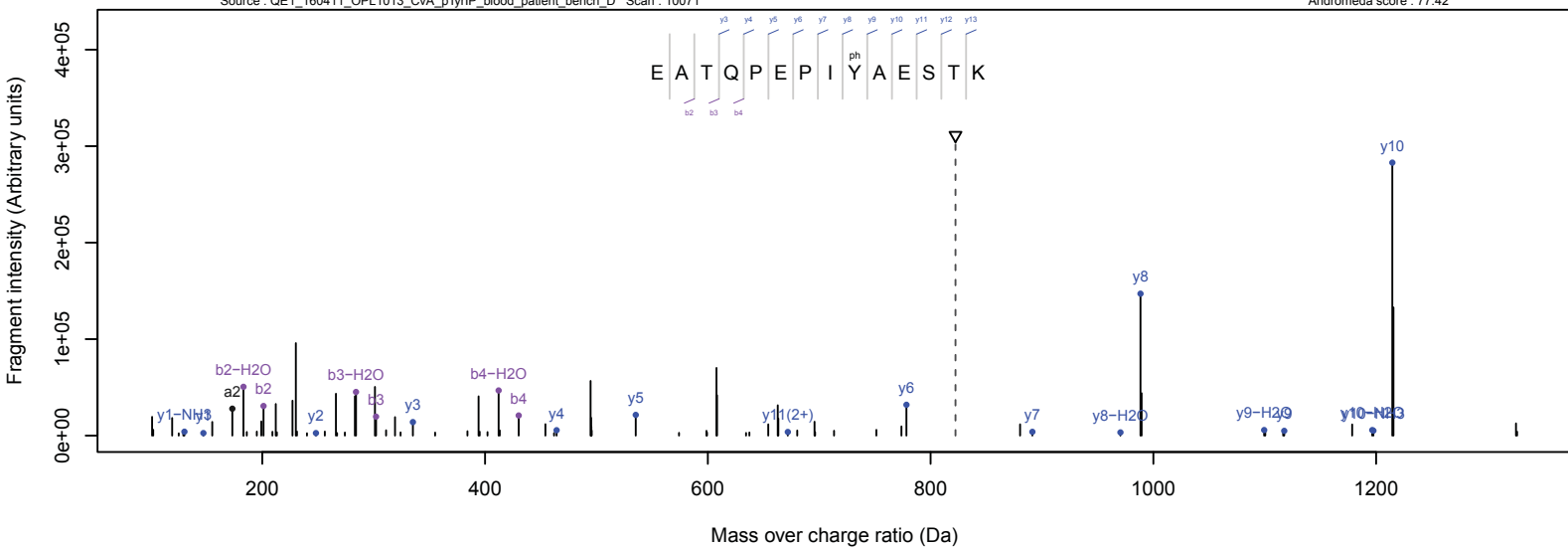
Andromeda score : 77.464



EATQPEIY(ph)AESTK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 10071

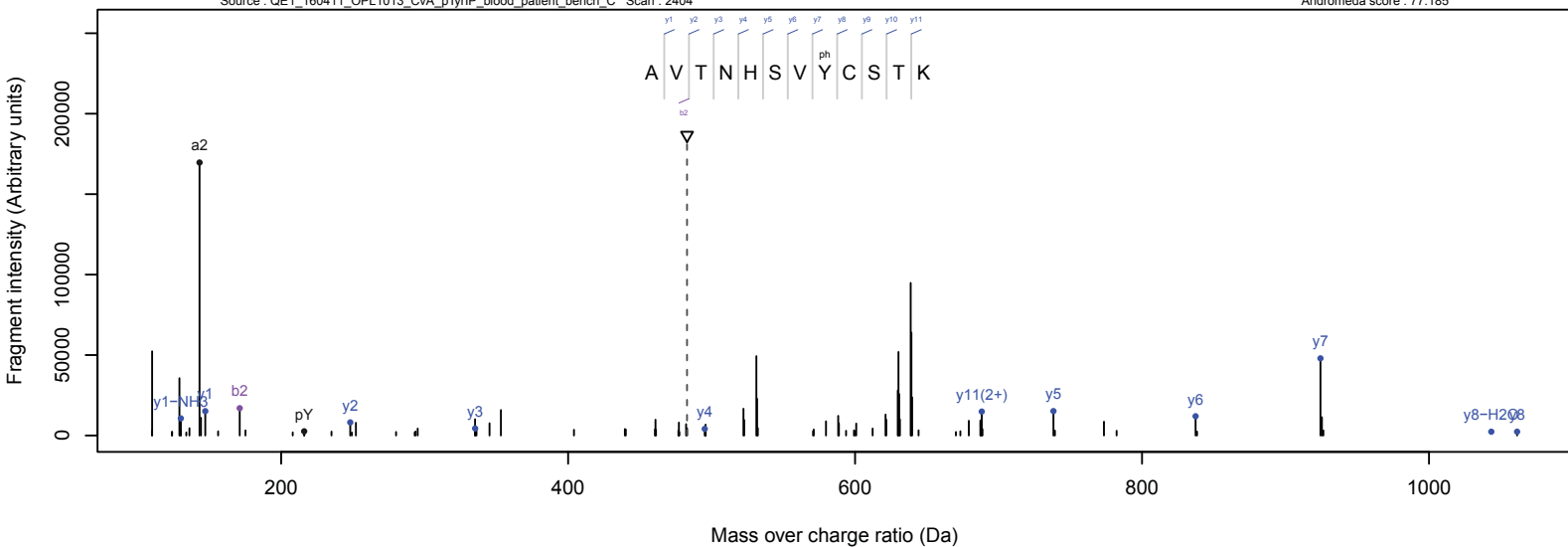
Andromeda score : 77.42



AVTNHSVY(ph)CSTK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 2404

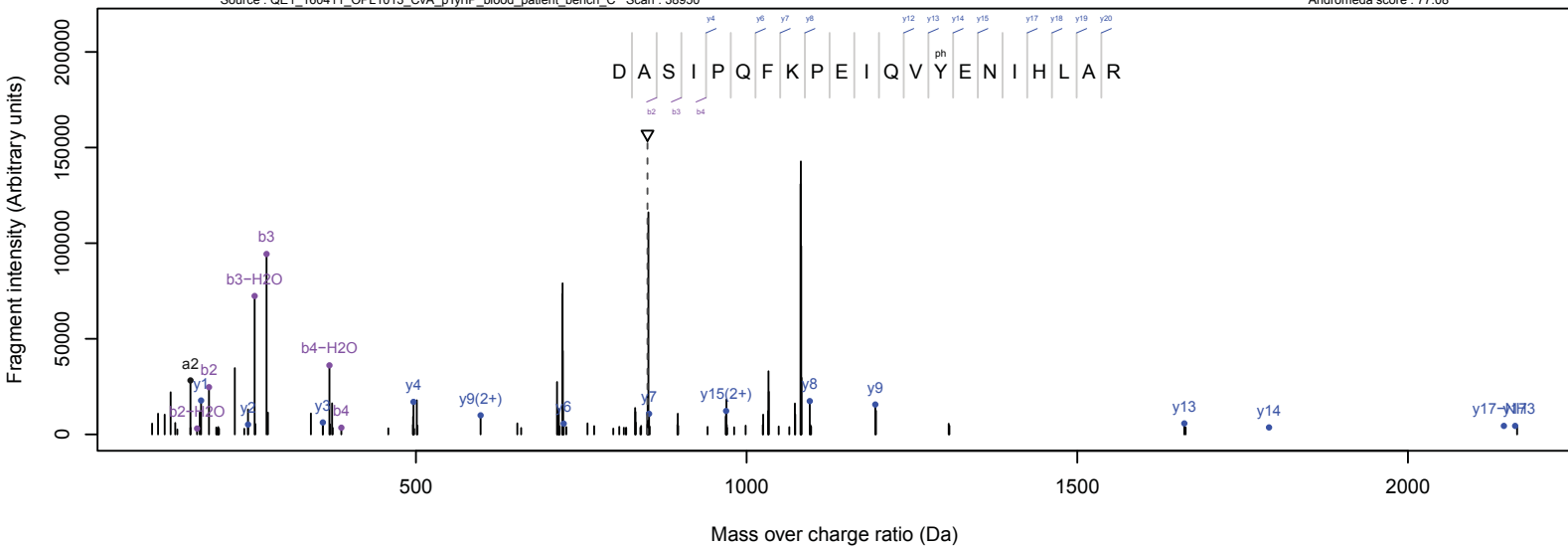
Andromeda score : 77.185



DASIPQFKPEIQVYENIHLAR-3+

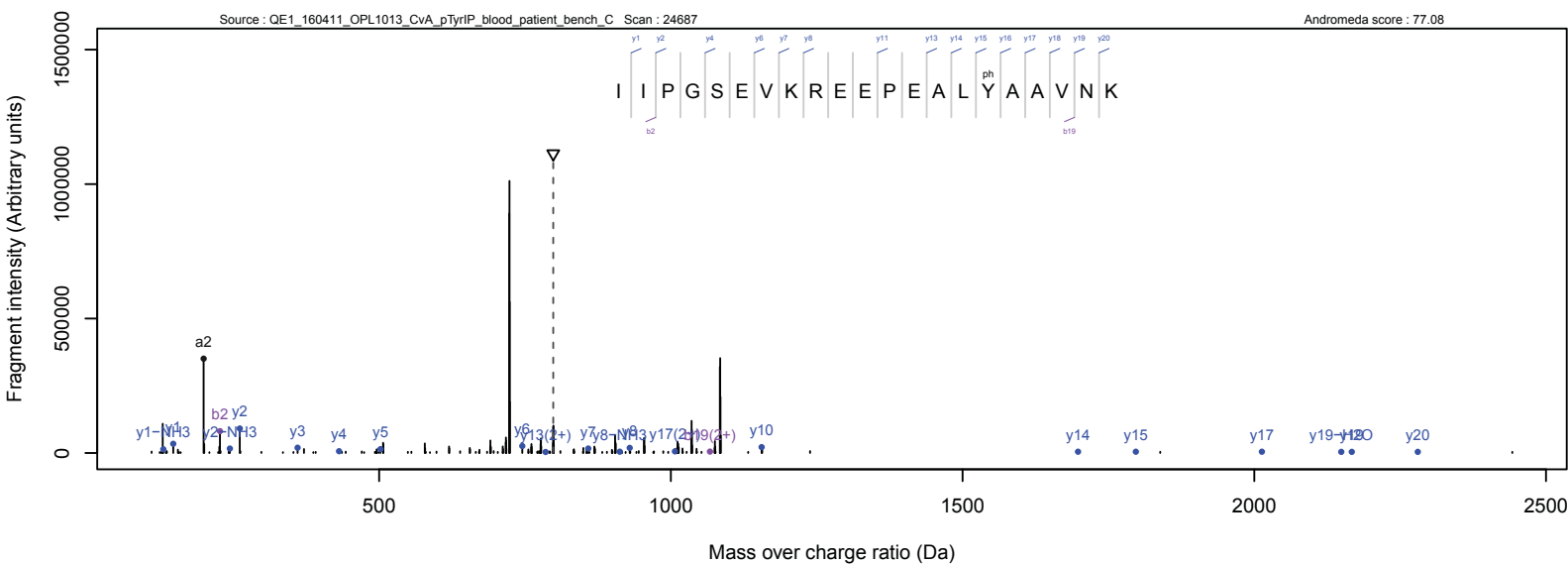
Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 38950

Andromeda score : 77.08

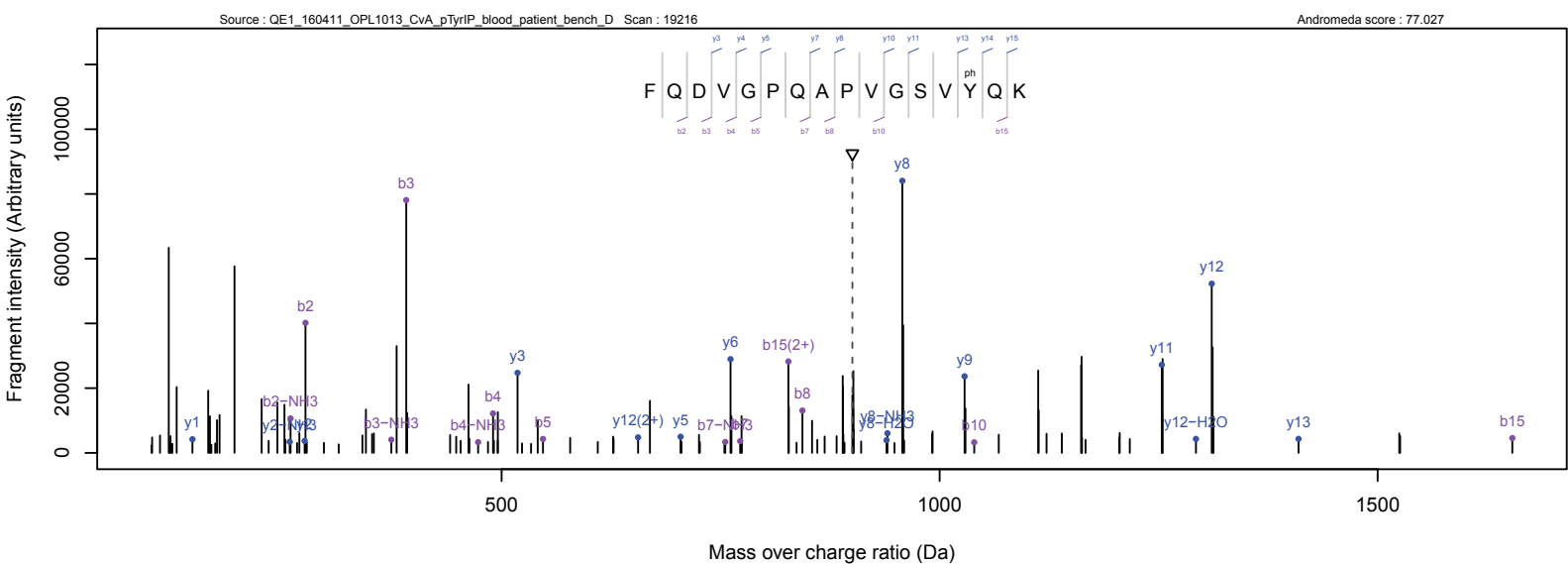




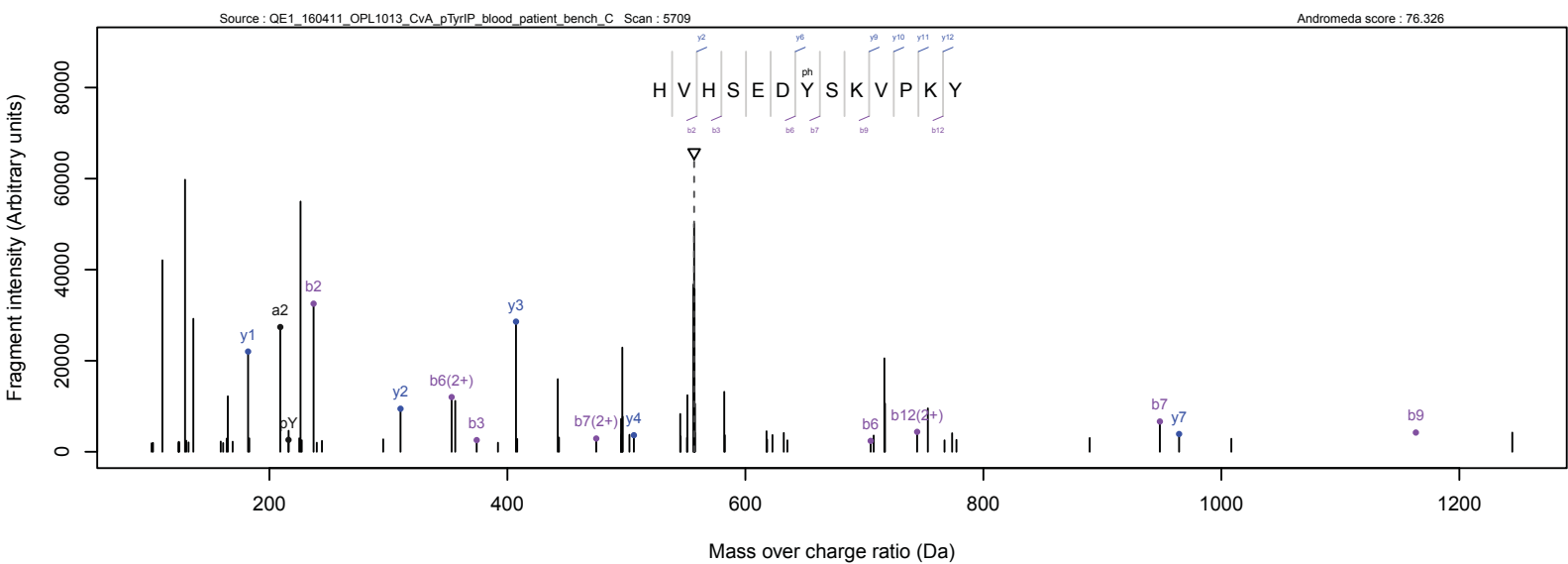
IIPGSEVKREEPEALY(ph)AAVNK-3+



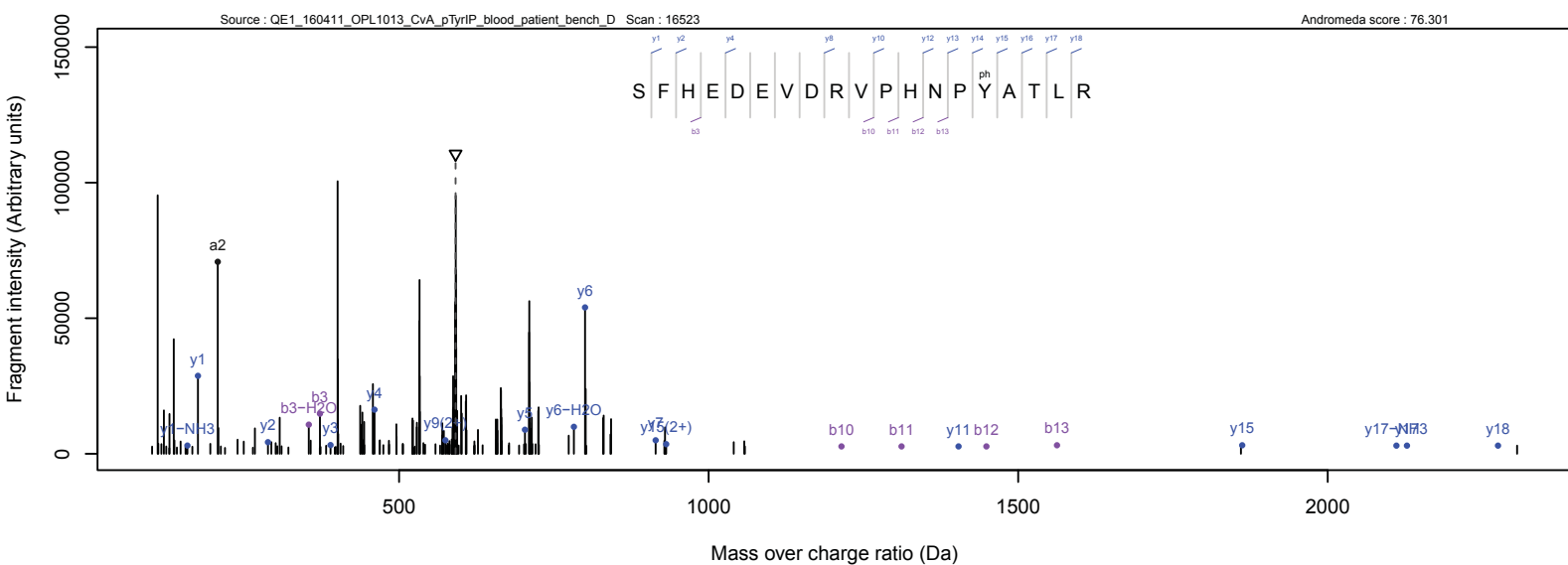
FQDVGPQAPVGSVY(ph)QK-2+



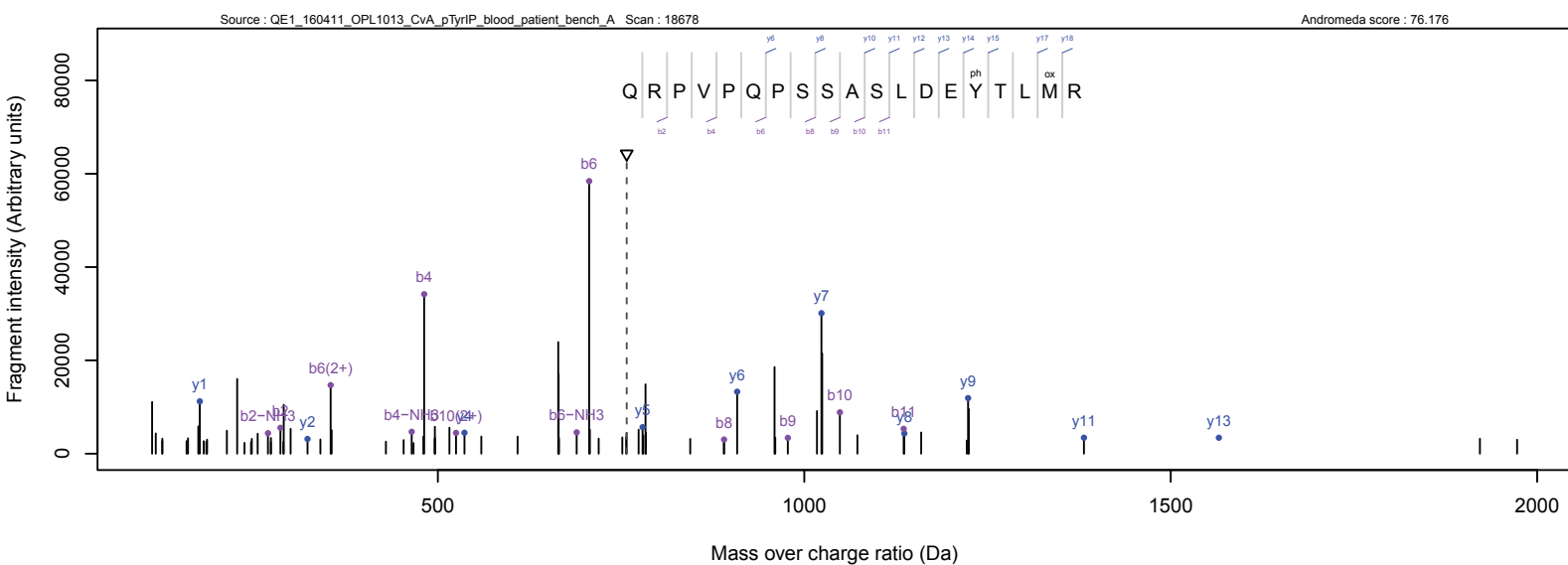
HVHSEY(ph)SKVPKY-3+



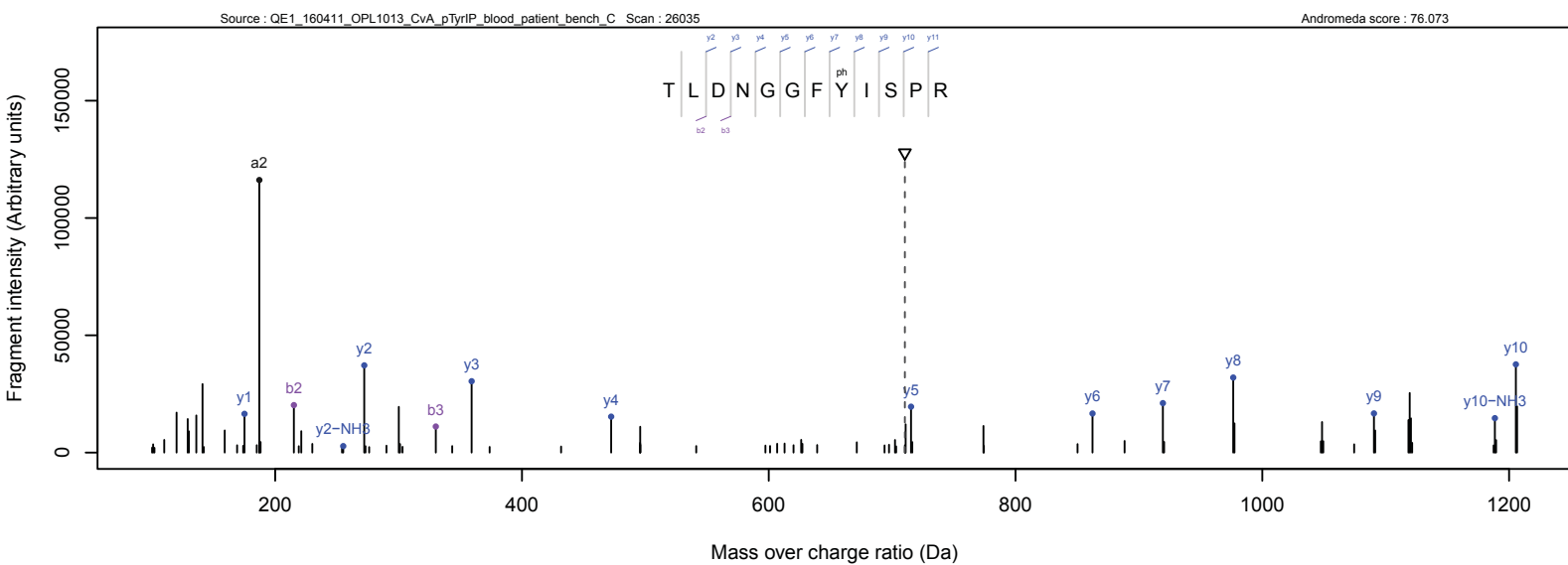
SFHEDEVDRVPHNPYATLR-4+



QRVPQPSSASLDEYTLMR-3+



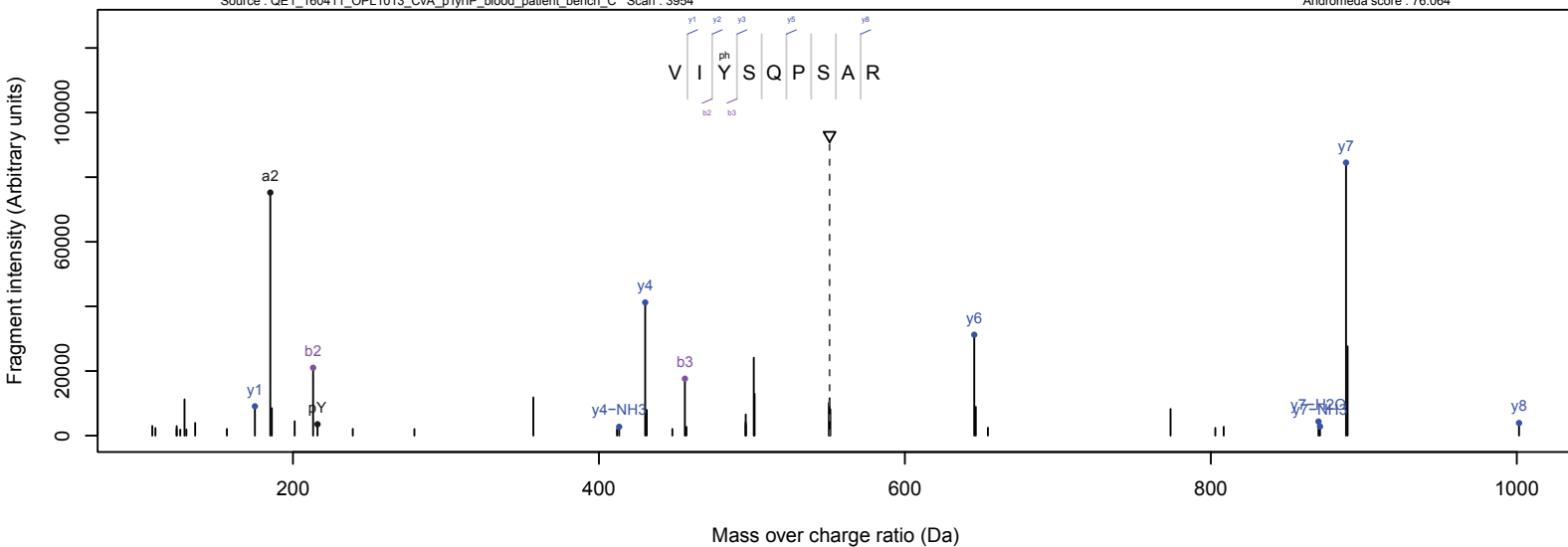
TLDNGGFYISPR-2+



## VIY(ph)SQPSAR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3954

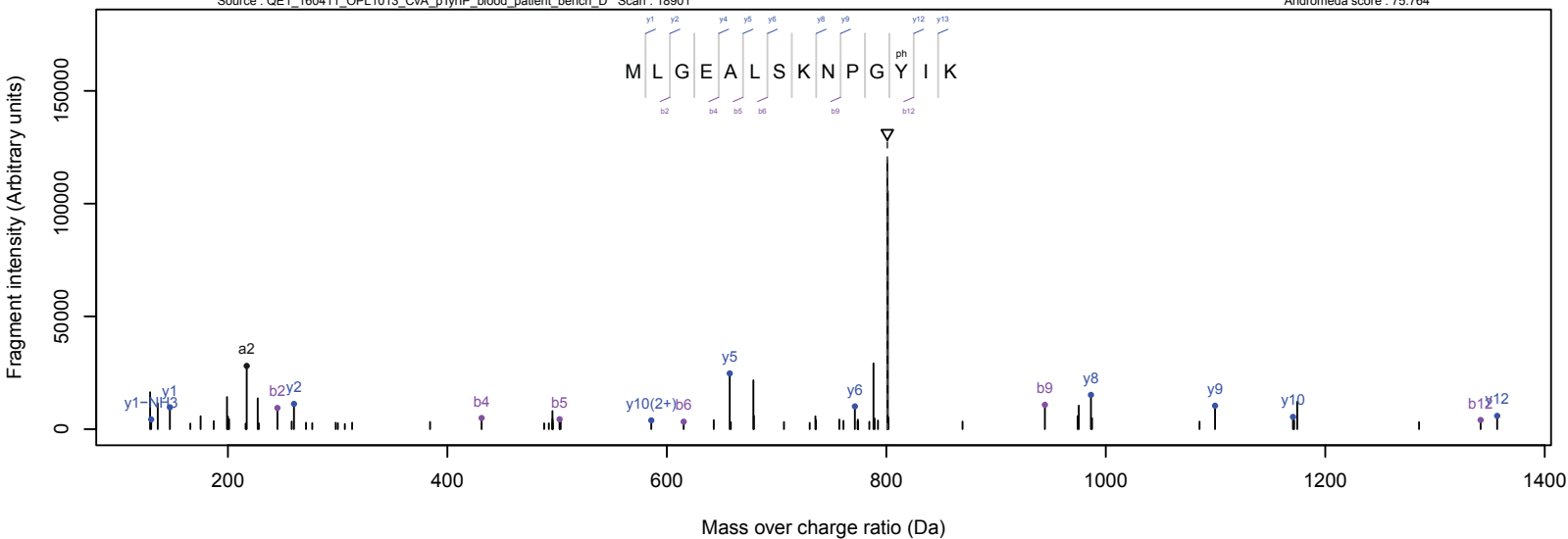
Andromeda score : 76.064



## MLGEALSKNPGY(ph)IK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 18901

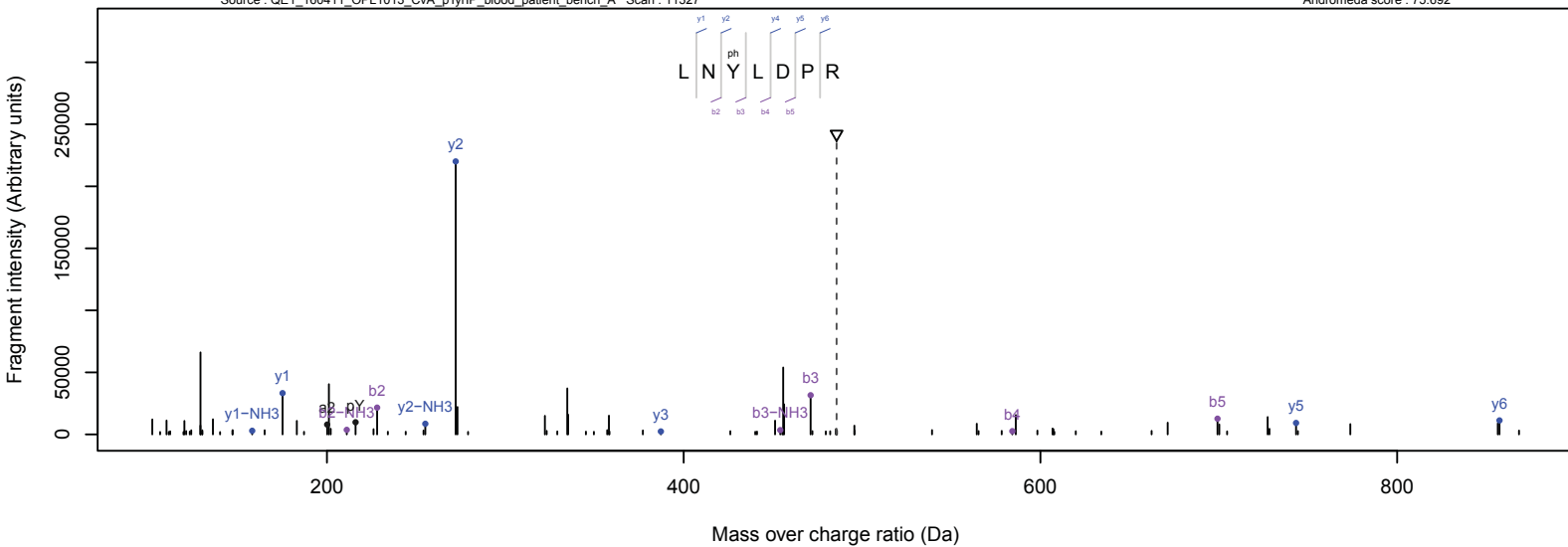
Andromeda score : 75.764



## LNY(ph)LDPR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 11327

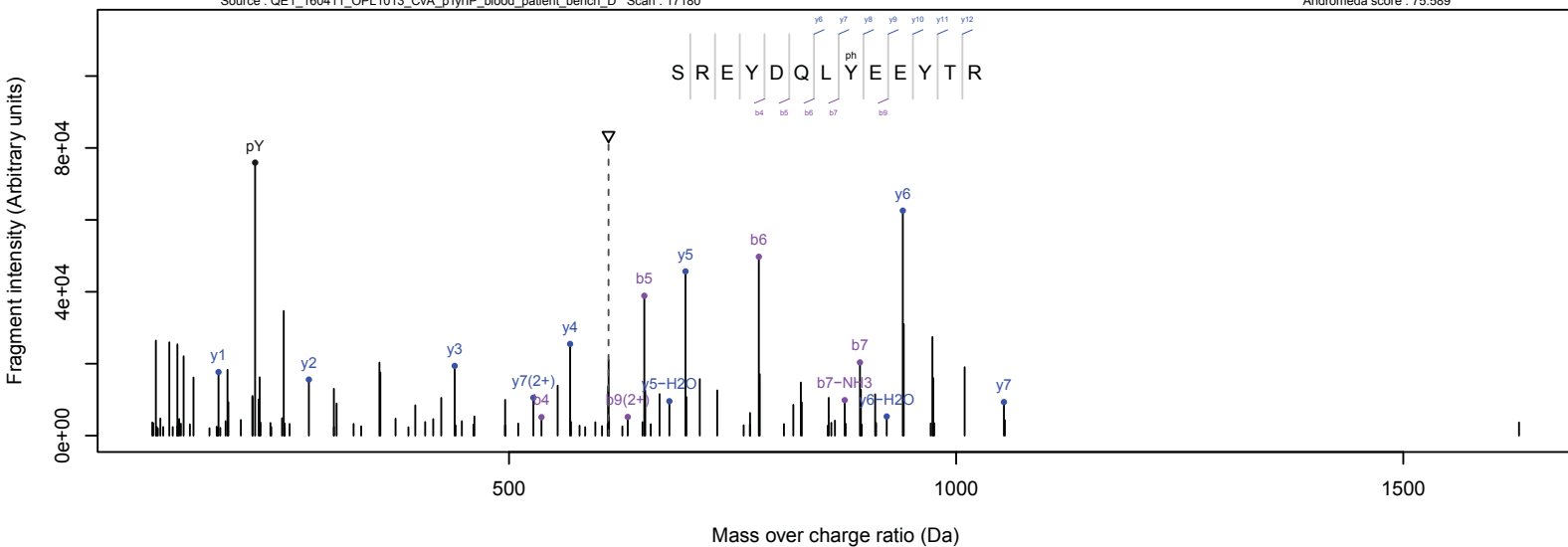
Andromeda score : 75.692



## SREYDQLY(ph)EEYTR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17180

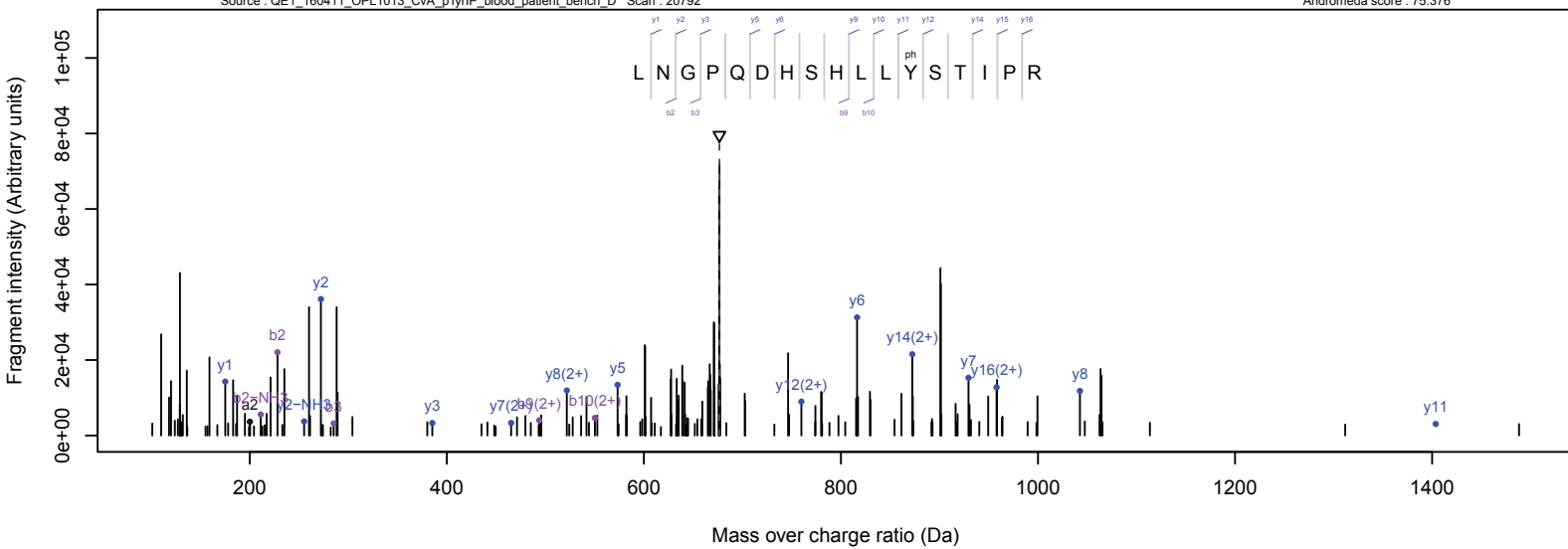
Andromeda score : 75.589



## LNGPQDHSLLY(ph)STIPR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 20792

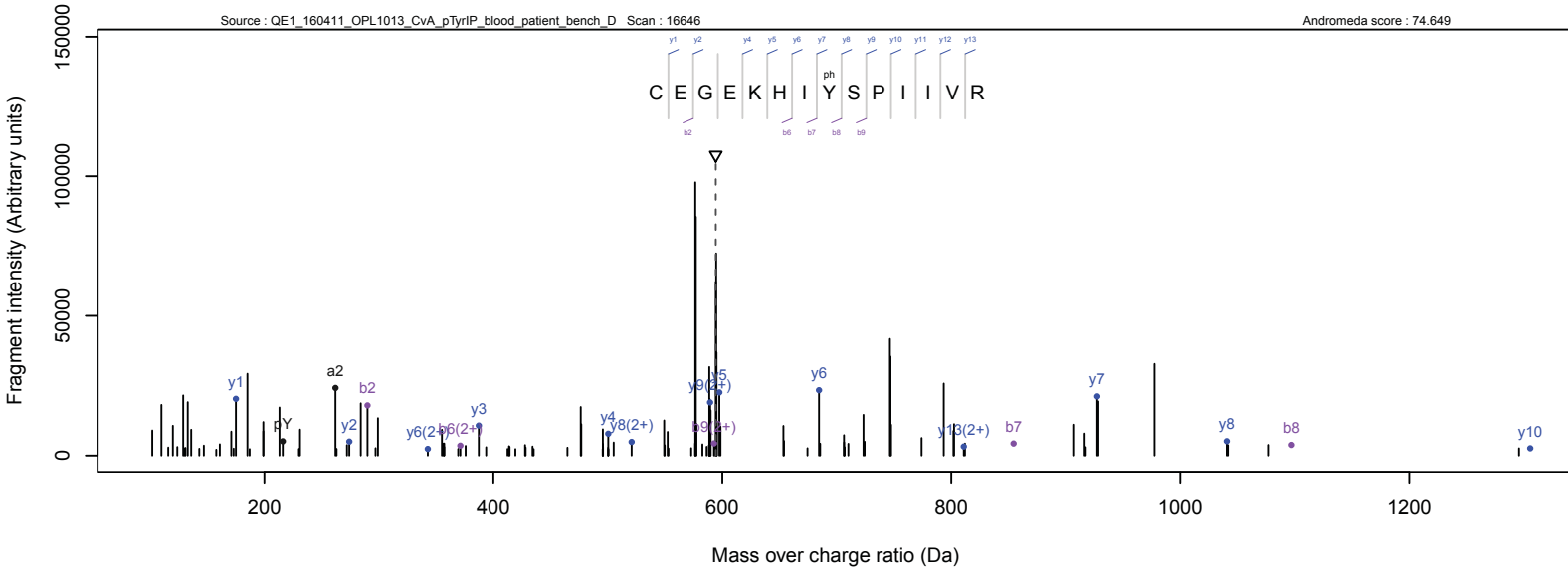
Andromeda score : 75.376



## CEGEKHIY(ph)SPIIVR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 16646

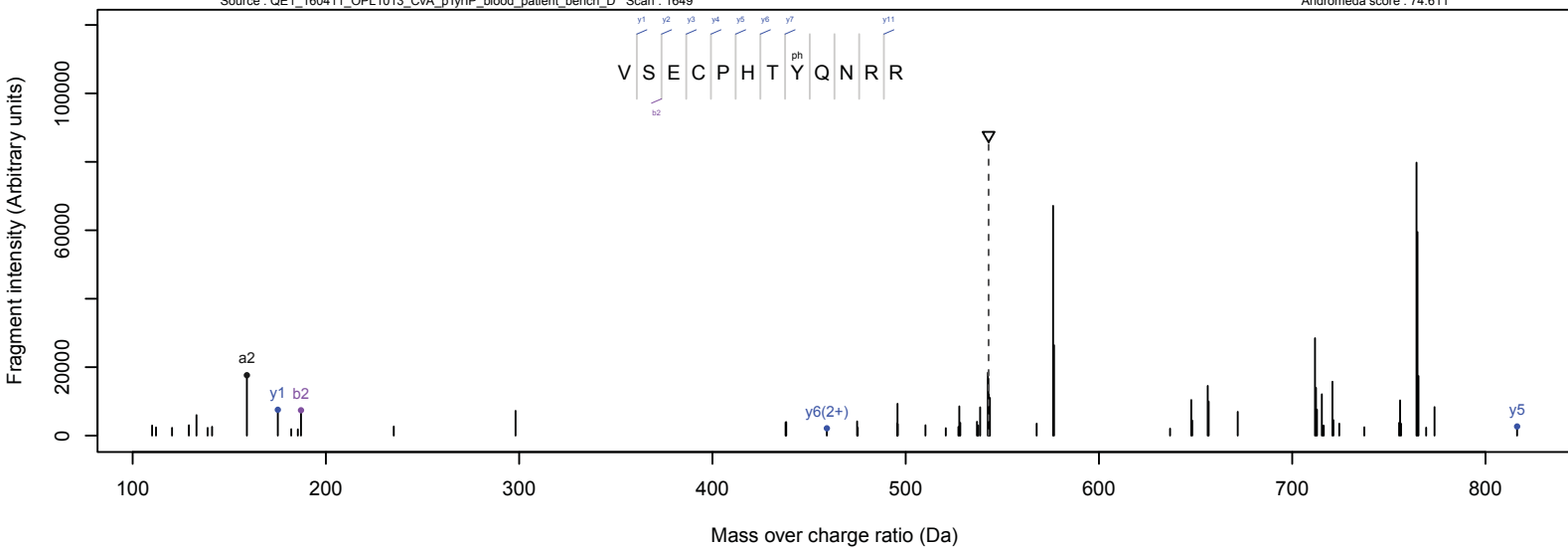
Andromeda score : 74.649



## VSECPHTY(ph)QNR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 1649

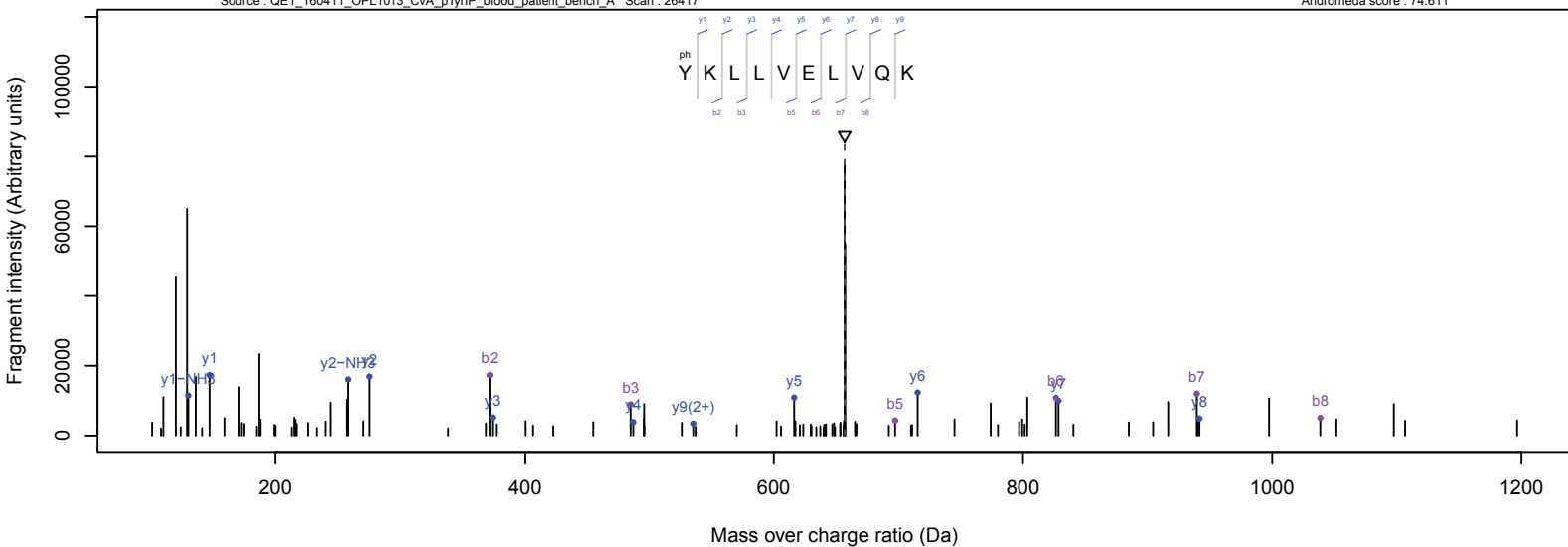
Andromeda score: 74.611



## Y(ph)KLLVELVQK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 26417

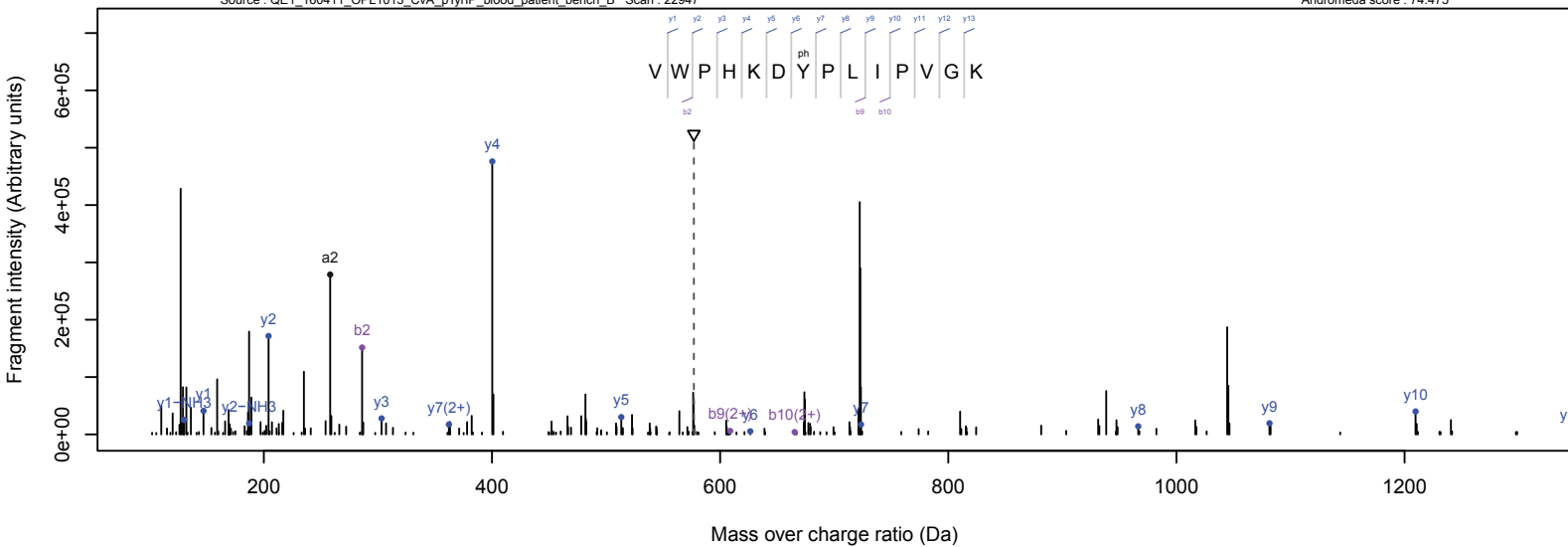
Andromeda score: 74.611



## VWPHKDY(ph)PLIPVGK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 22947

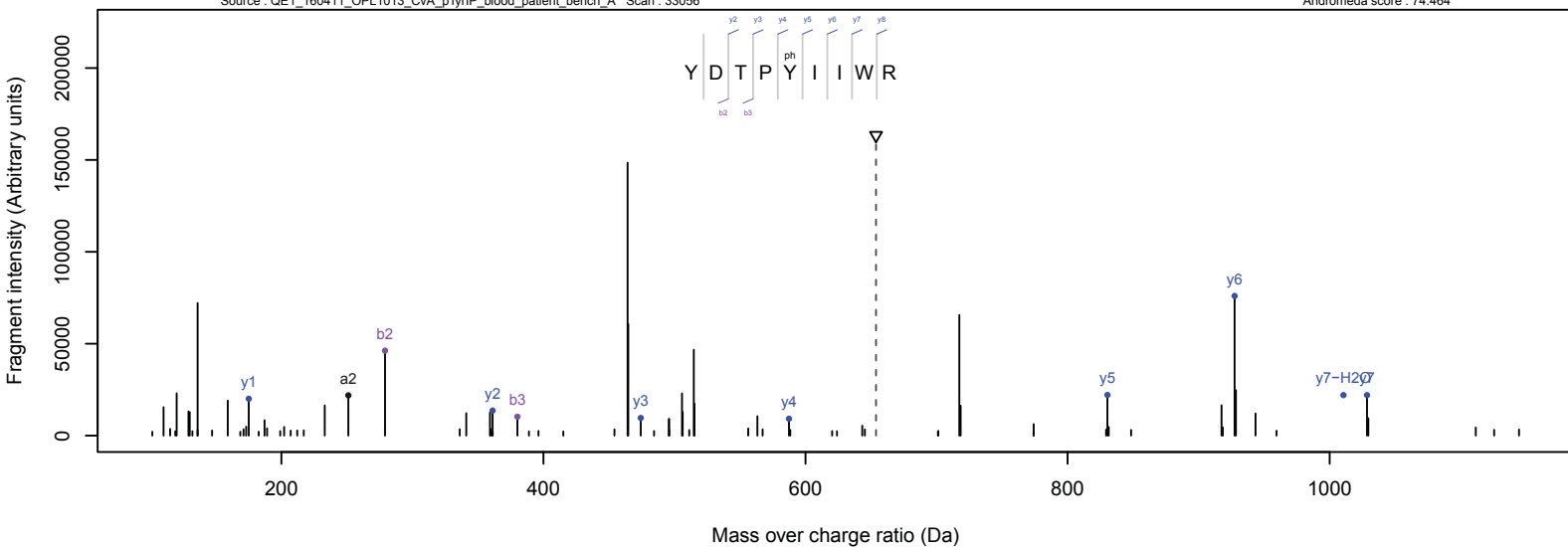
Andromeda score: 74.475



## YDTPY(ph)IWR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 33056

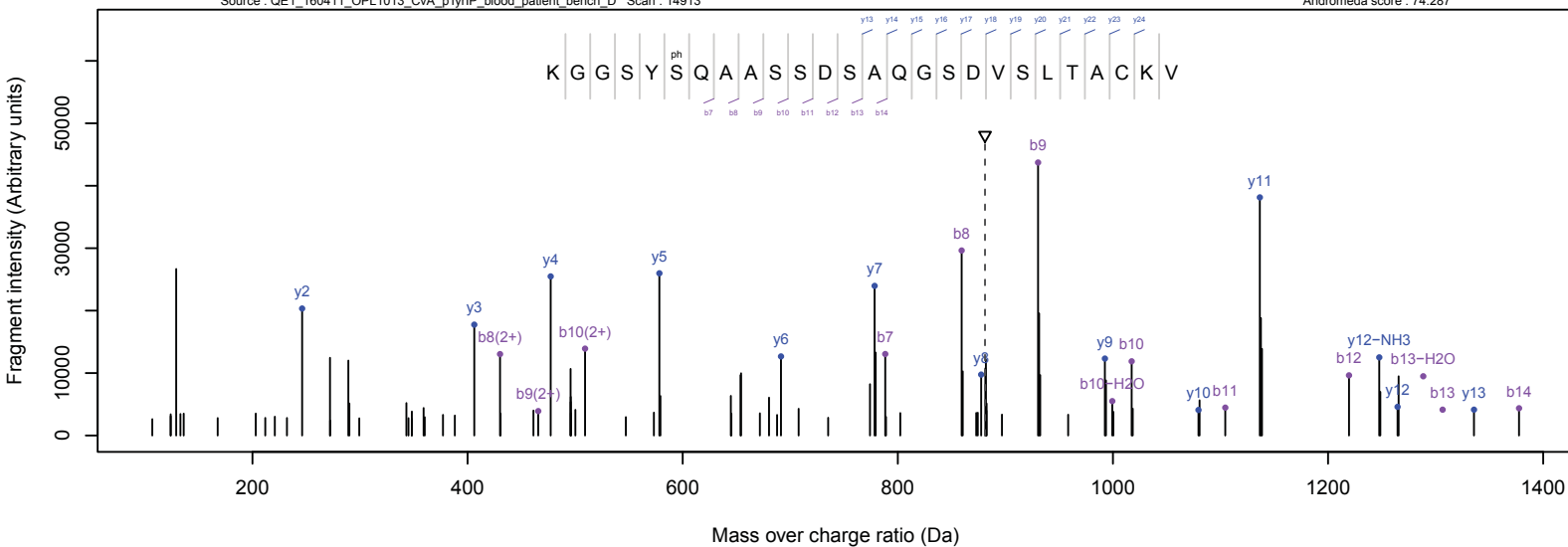
Andromeda score : 74.464



## KGGYS(ph)QAASSDAQSDVSLTACKV-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 14913

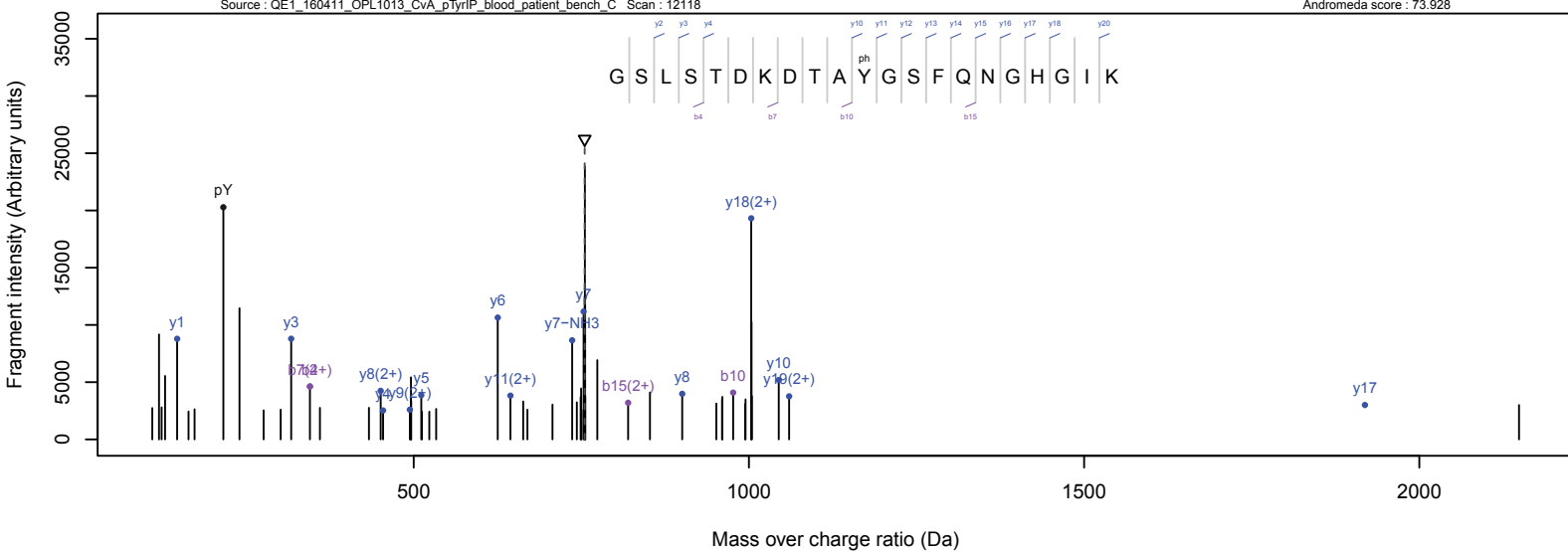
Andromeda score : 74.287



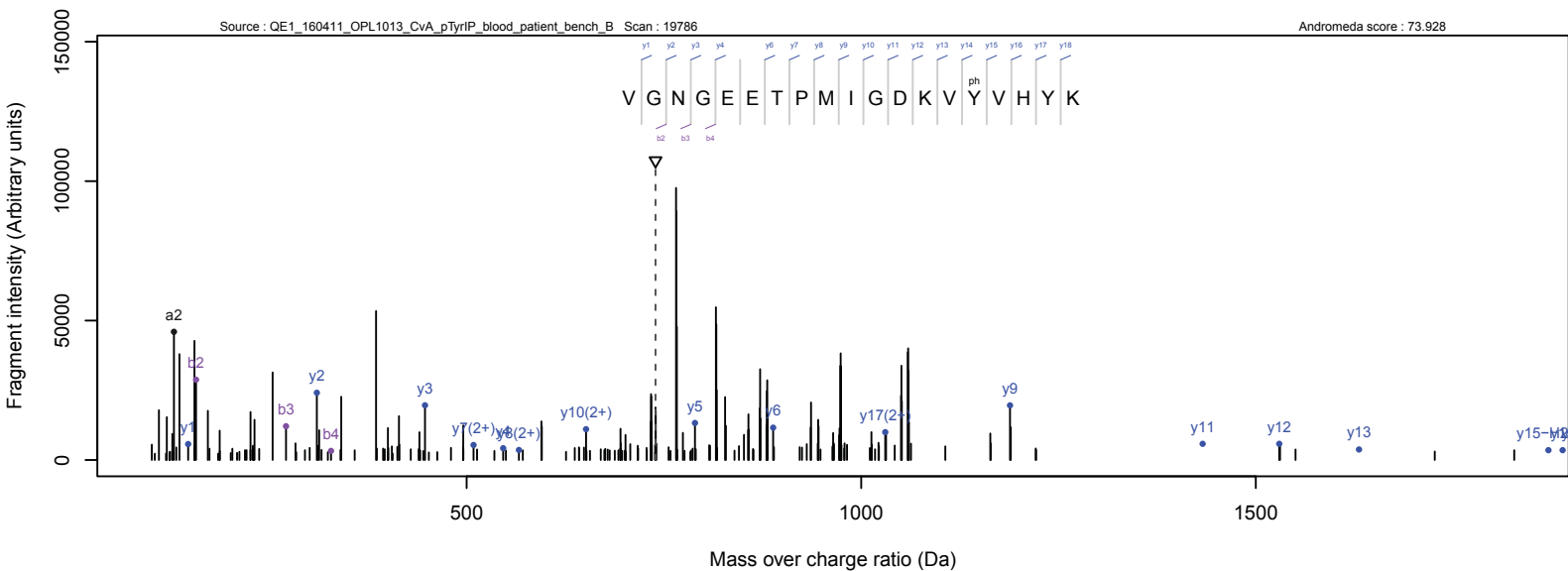
## GSLSTDKDTAY(ph)GSFQNGHGK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 12118

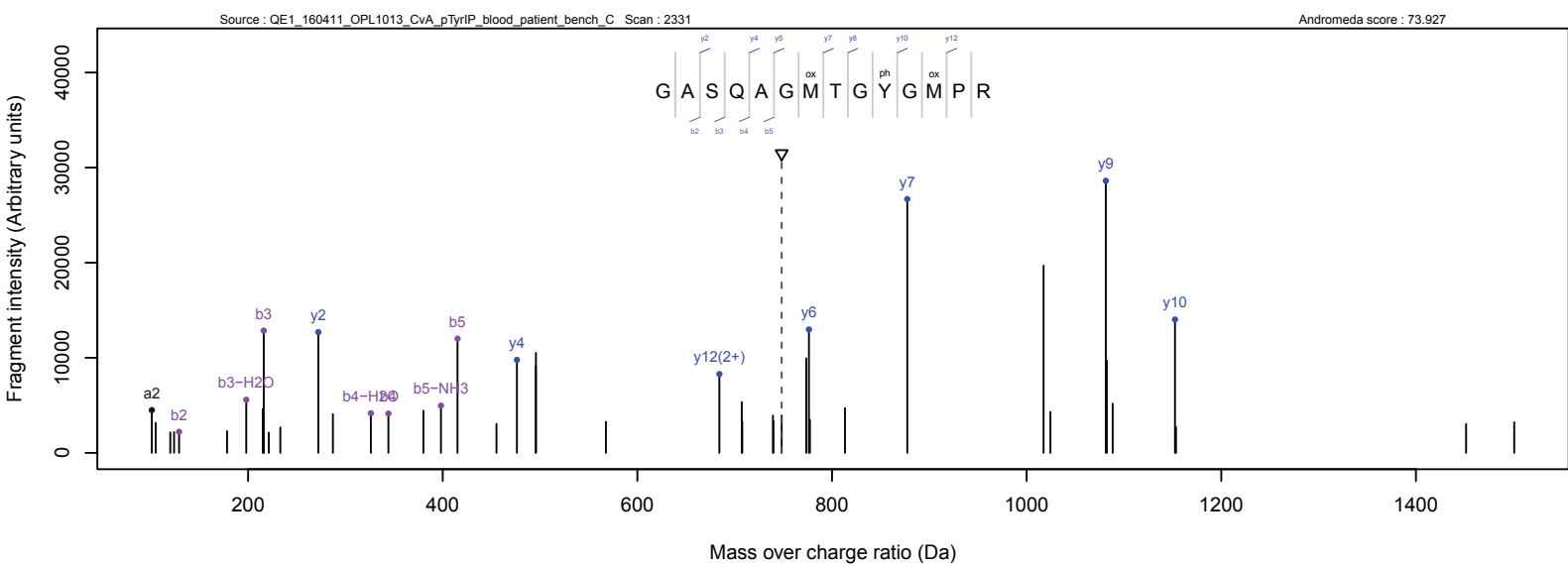
Andromeda score : 73.928



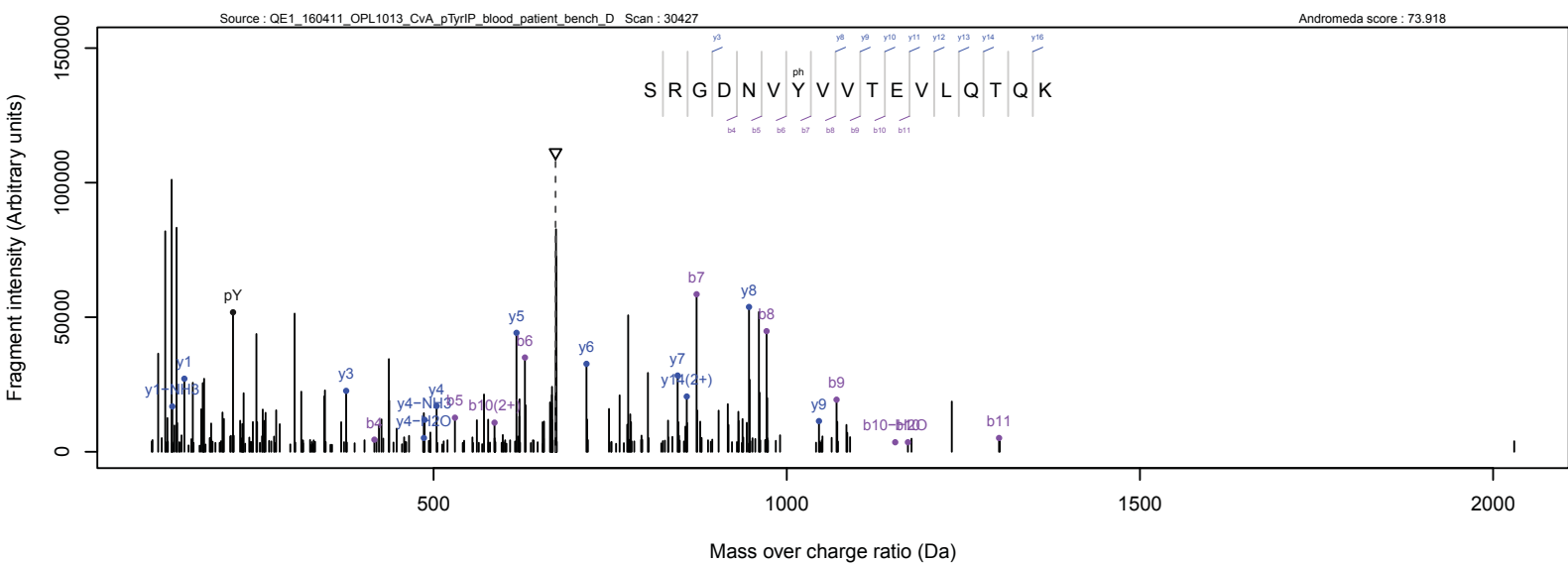
**VNGEETPMIGDKVY(ph)VHYK-3+**



**GASQAGM(oX)TGY(ph)GM(oX)PR-2+**



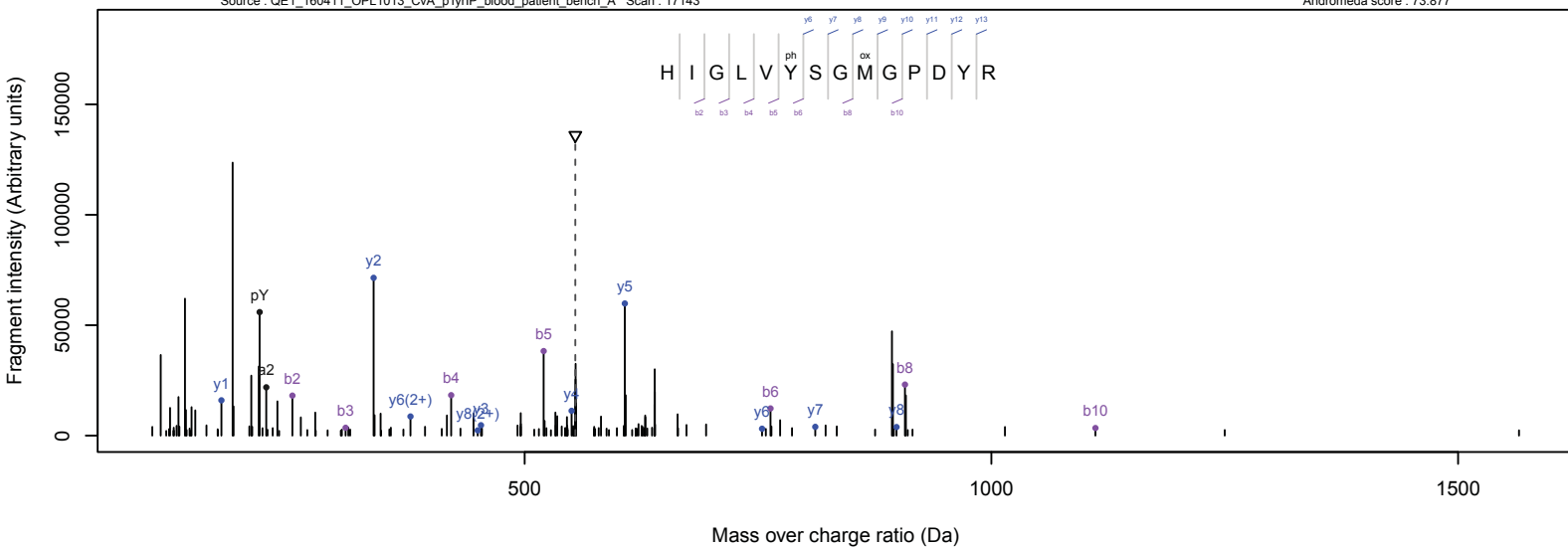
**SRGDNVY(ph)VVTEVLQTK-3+**



HIGLVY(ph)SGM(ox)GPDYR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 17143

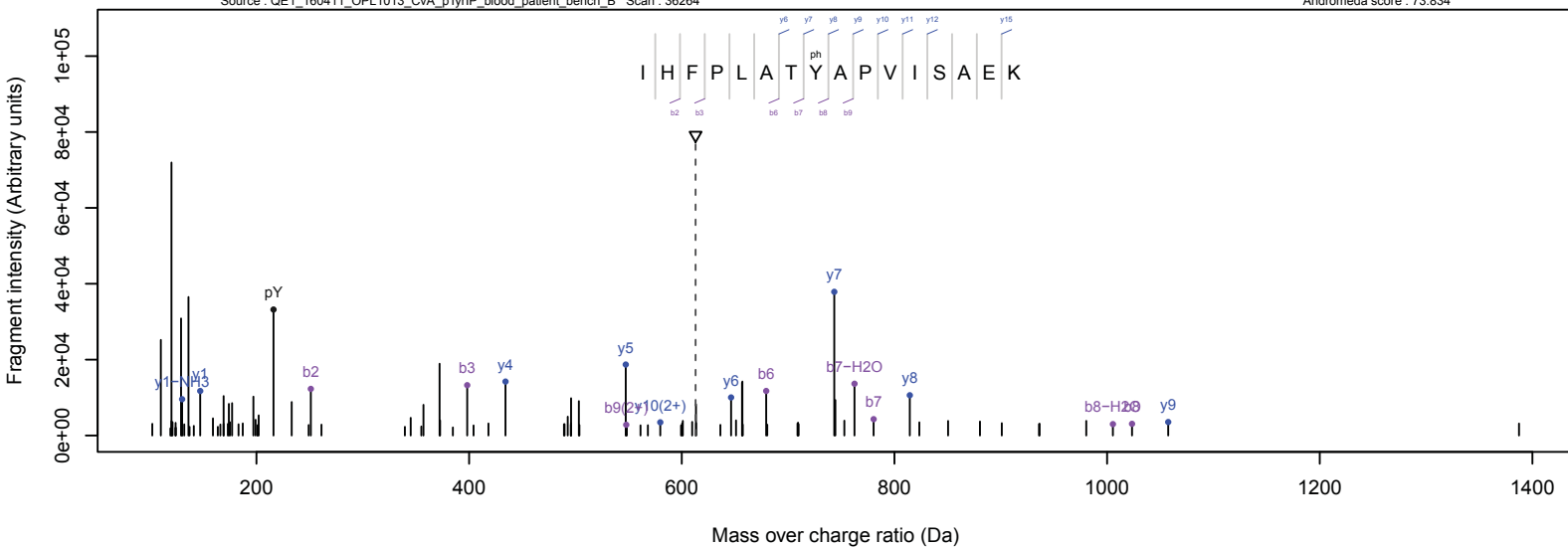
Andromeda score : 73.877



IHFPLATY(ph)APVISA EK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 36264

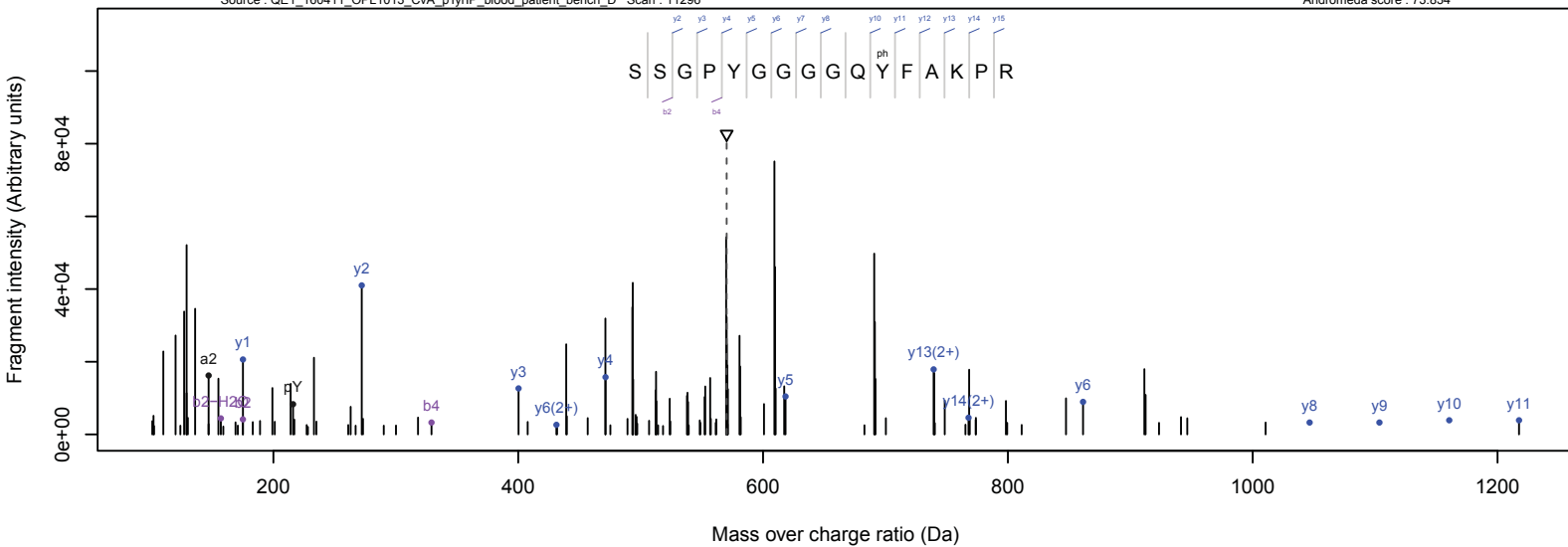
Andromeda score : 73.834



SSGPYGGGQY(ph)FAKPR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11296

Andromeda score : 73.834

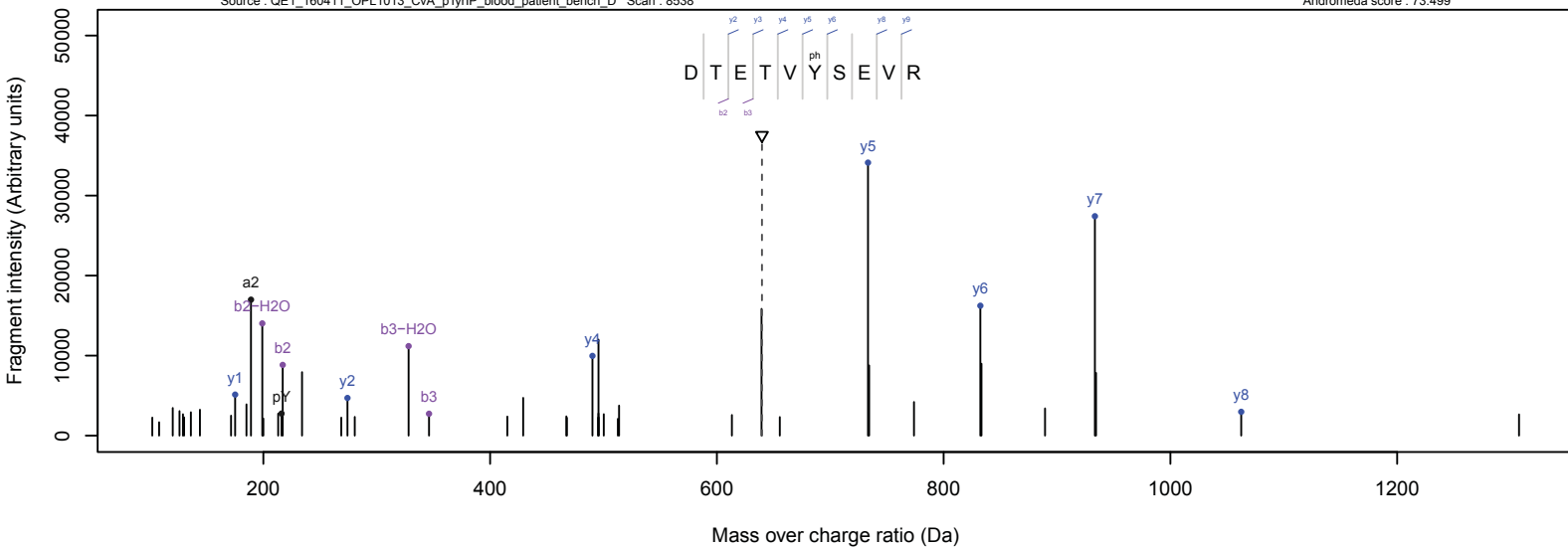




## DTETVY(ph)SEVR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 8538

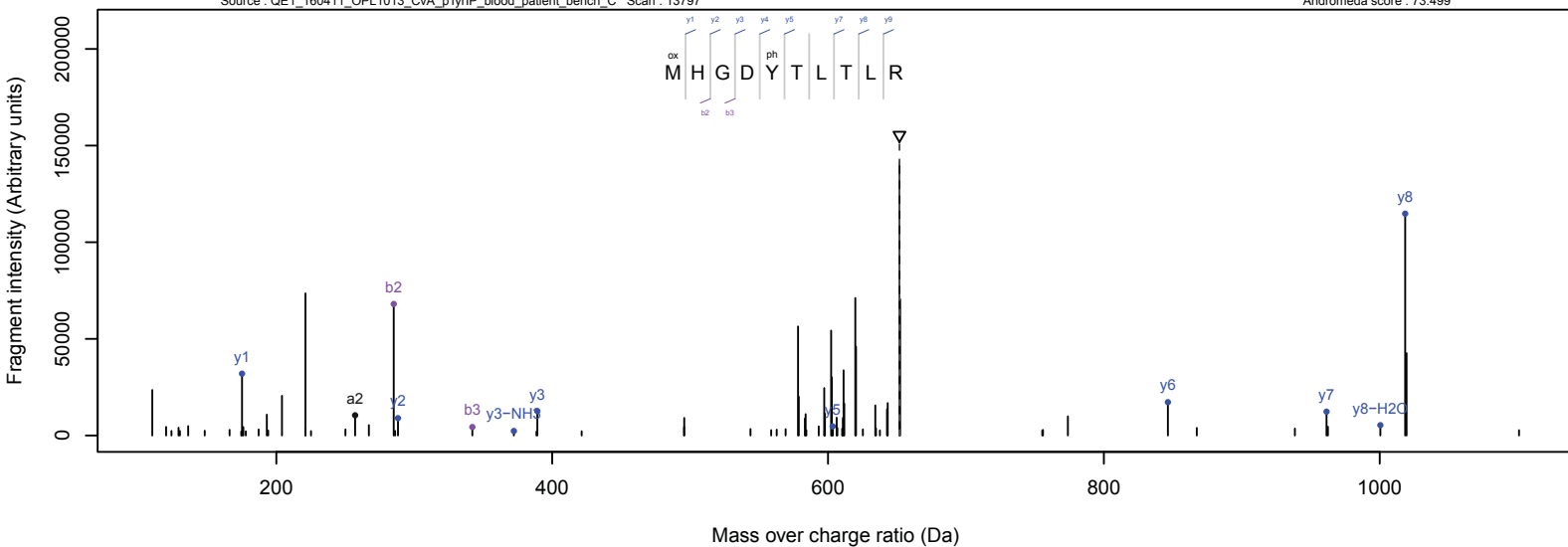
Andromeda score: 73.499



## M(ox)HGDY(ph)TLTLR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 13797

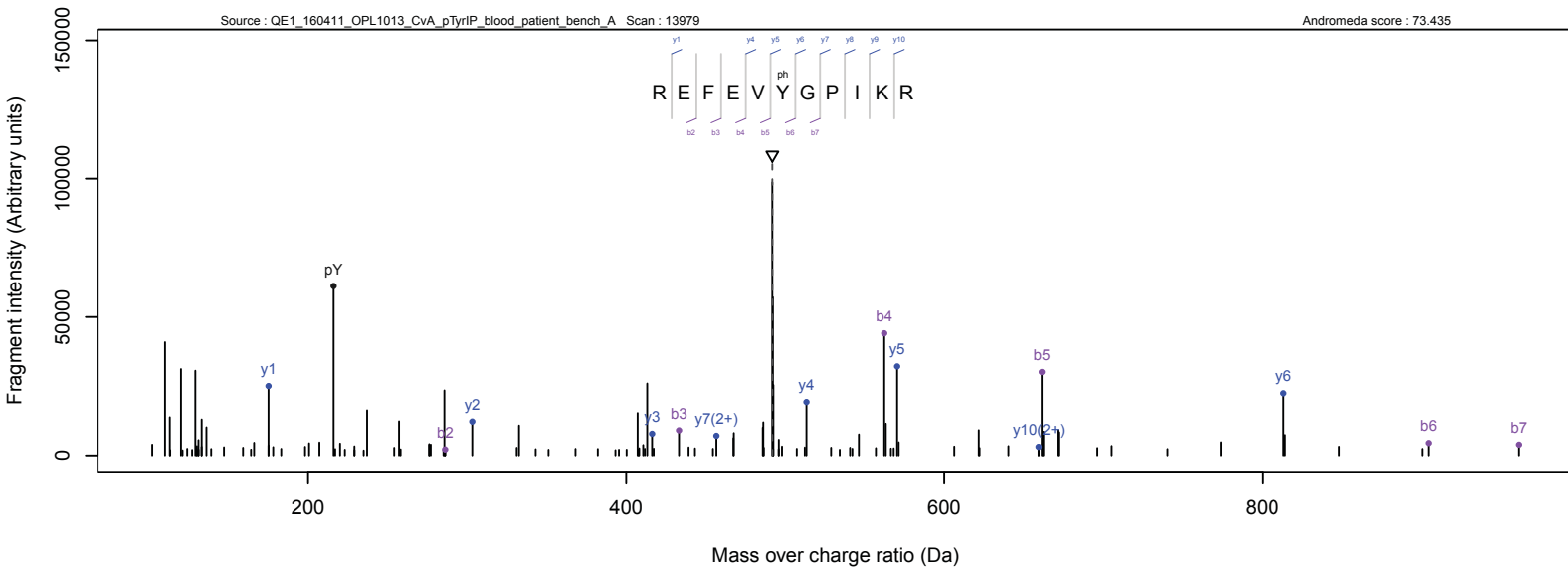
Andromeda score: 73.499



## REFEVY(ph)GPIKR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 13979

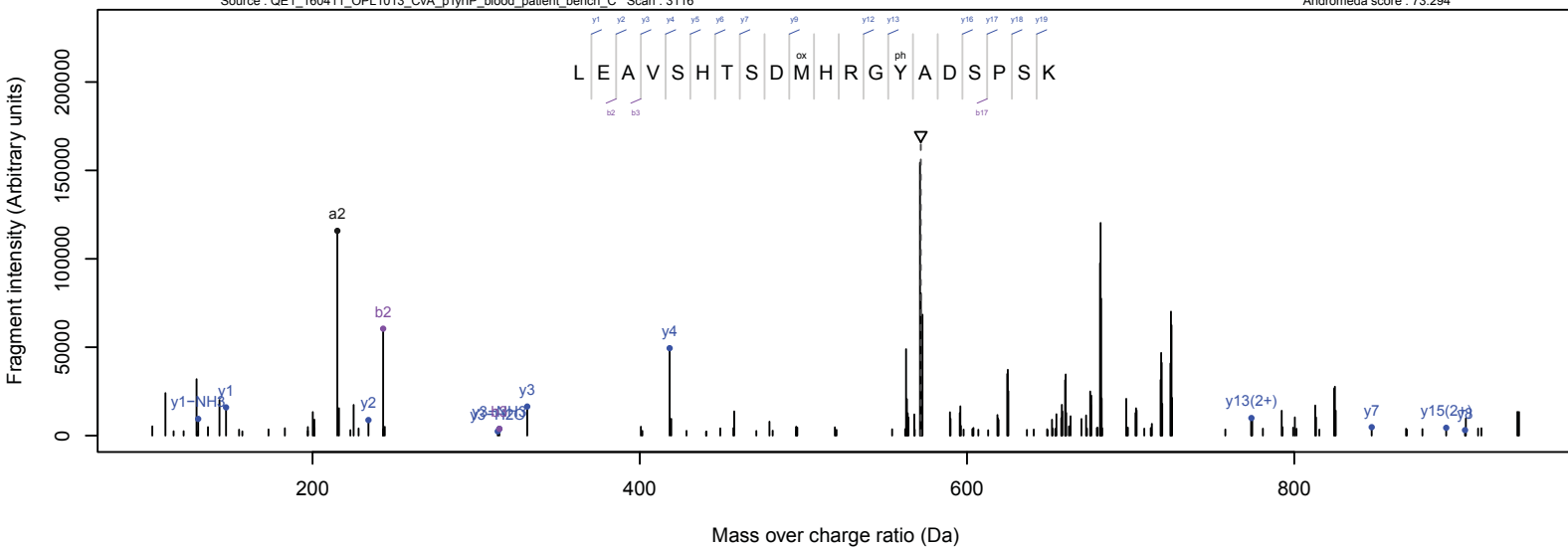
Andromeda score: 73.435



LEAVSHTSDM(ox)HRGY(ph)ADSPSK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3116

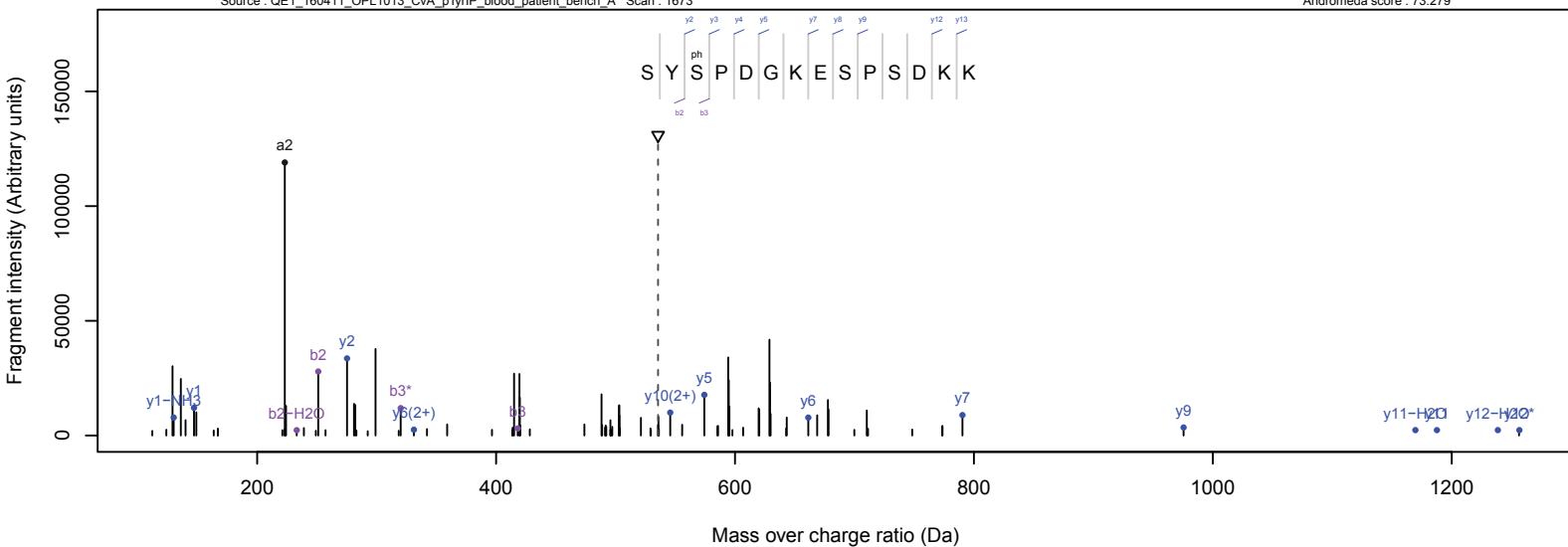
Andromeda score : 73.294



SYS(ph)PDGKESPSDKK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 1673

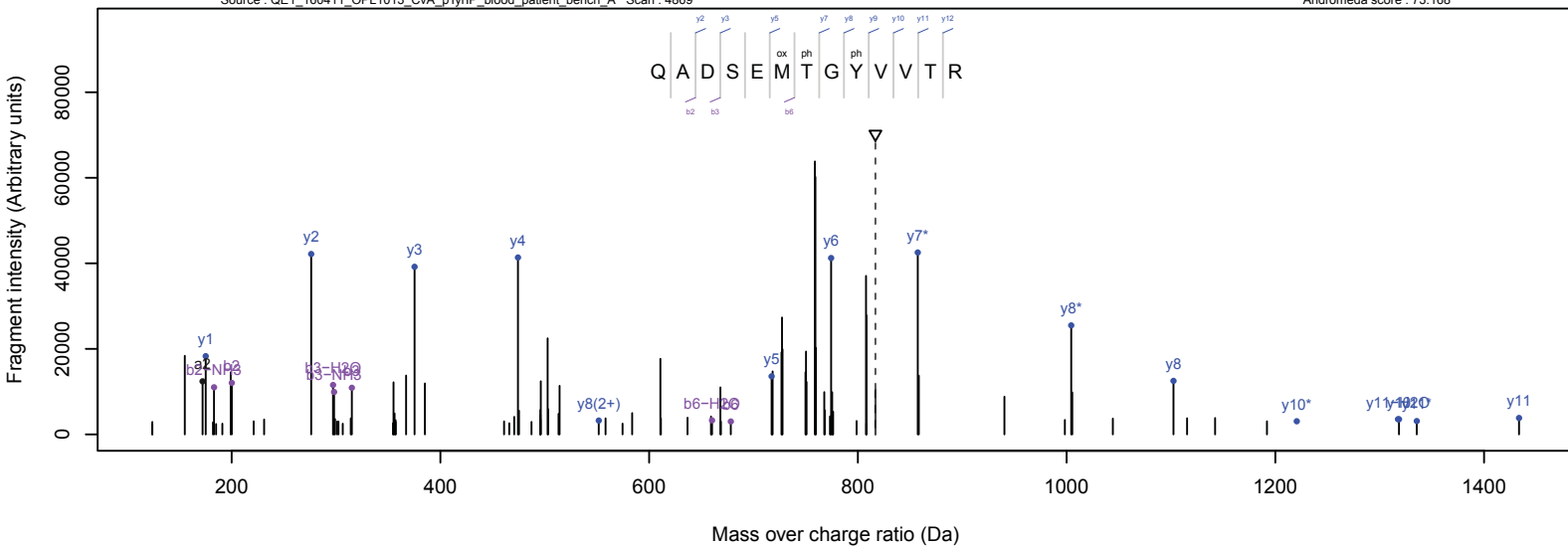
Andromeda score : 73.279

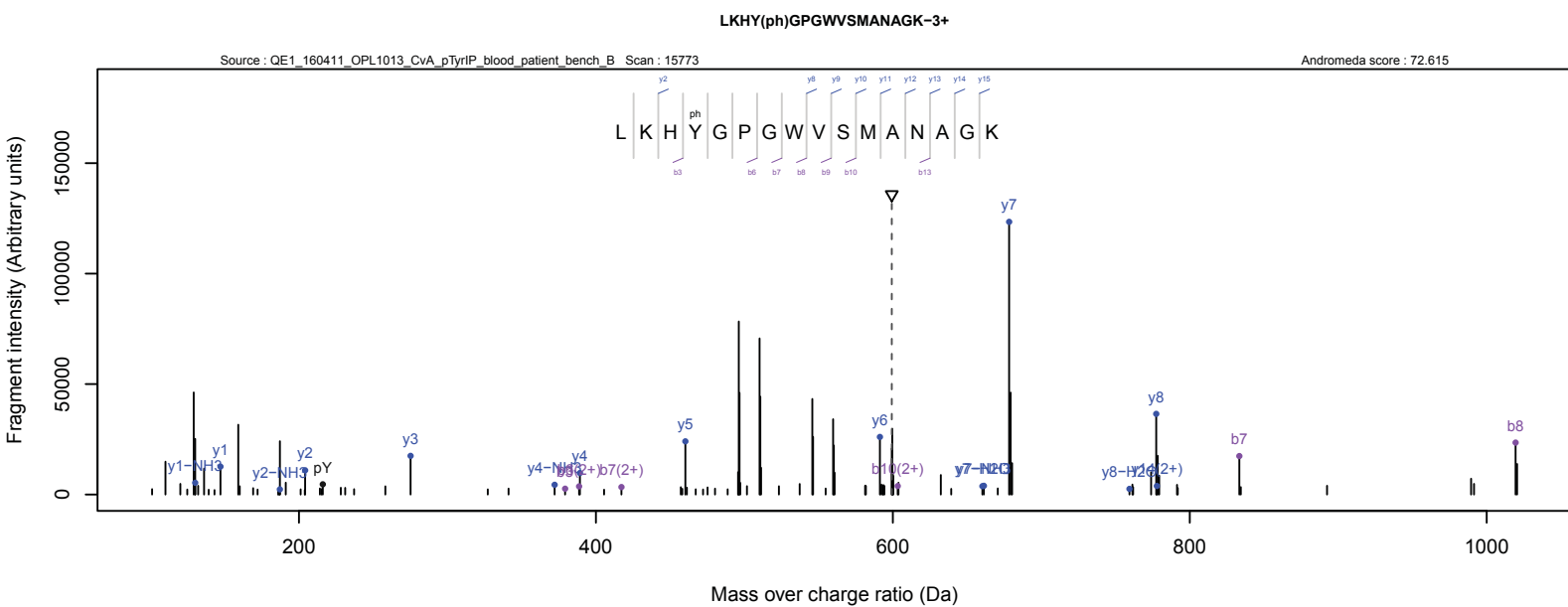
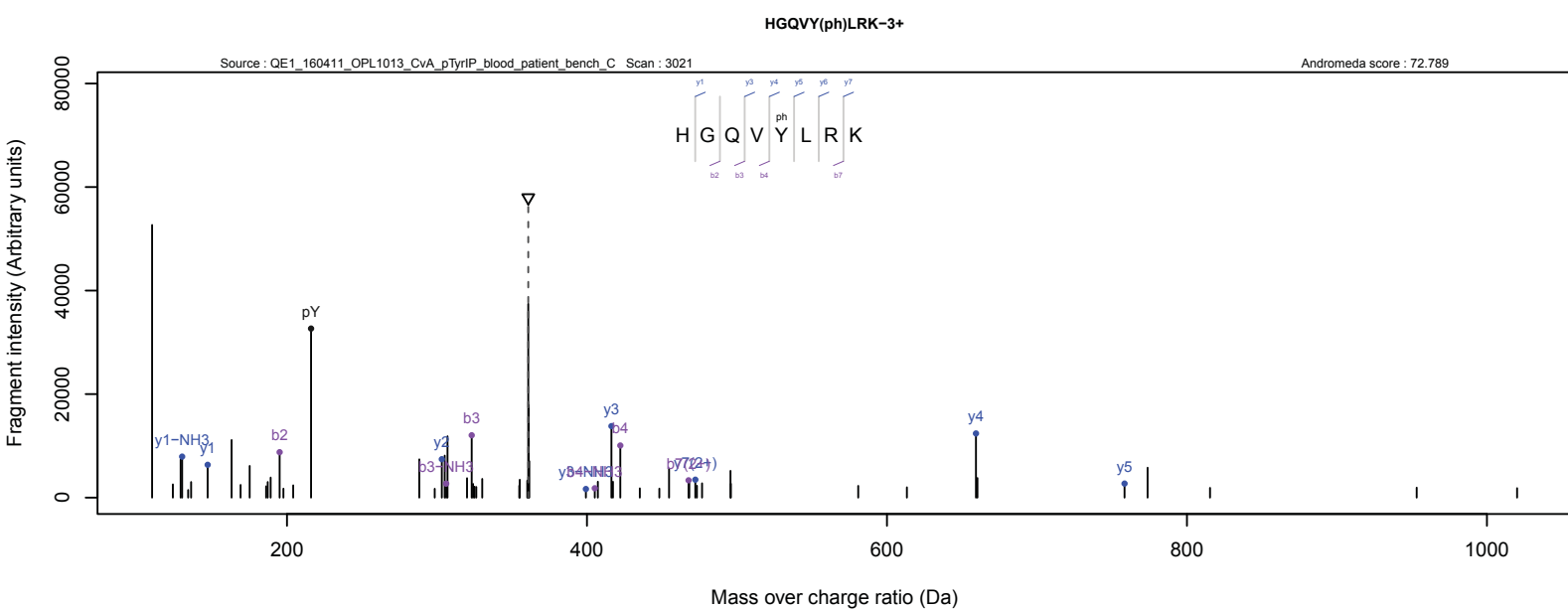
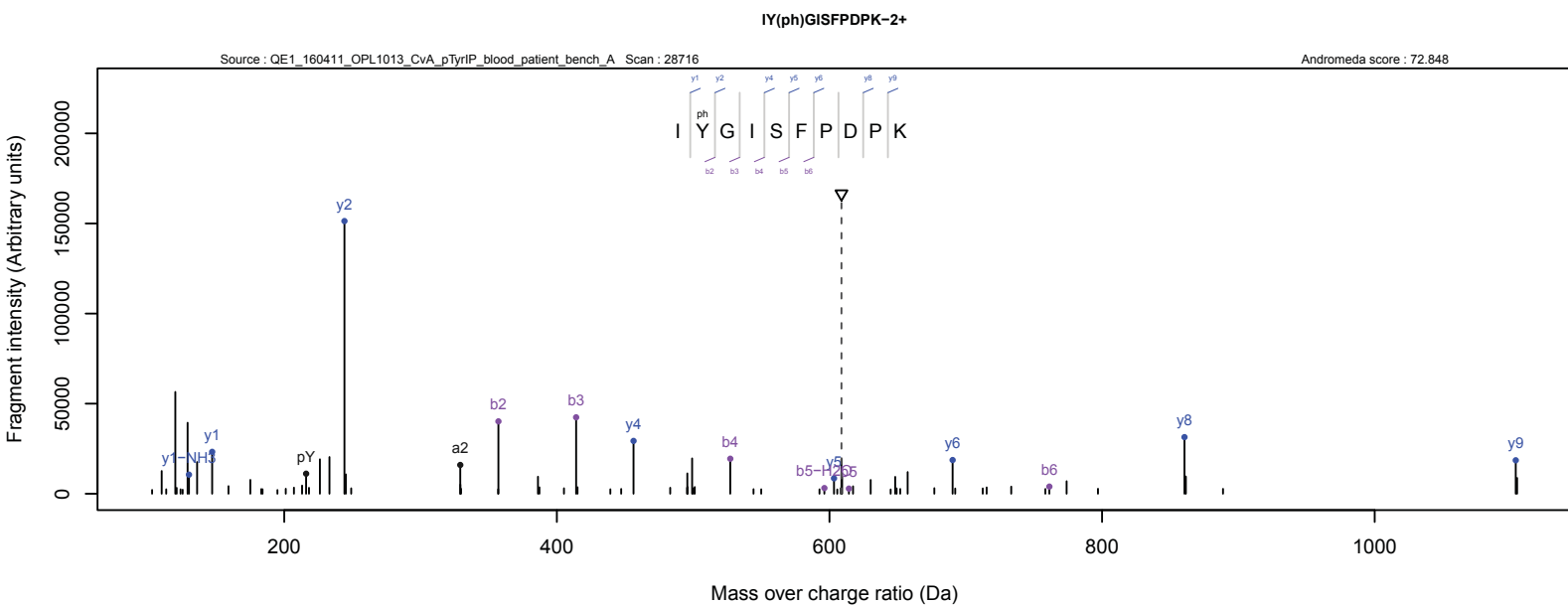


QADSEM(ox)T(ph)GY(ph)VVTR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 4869

Andromeda score : 73.168

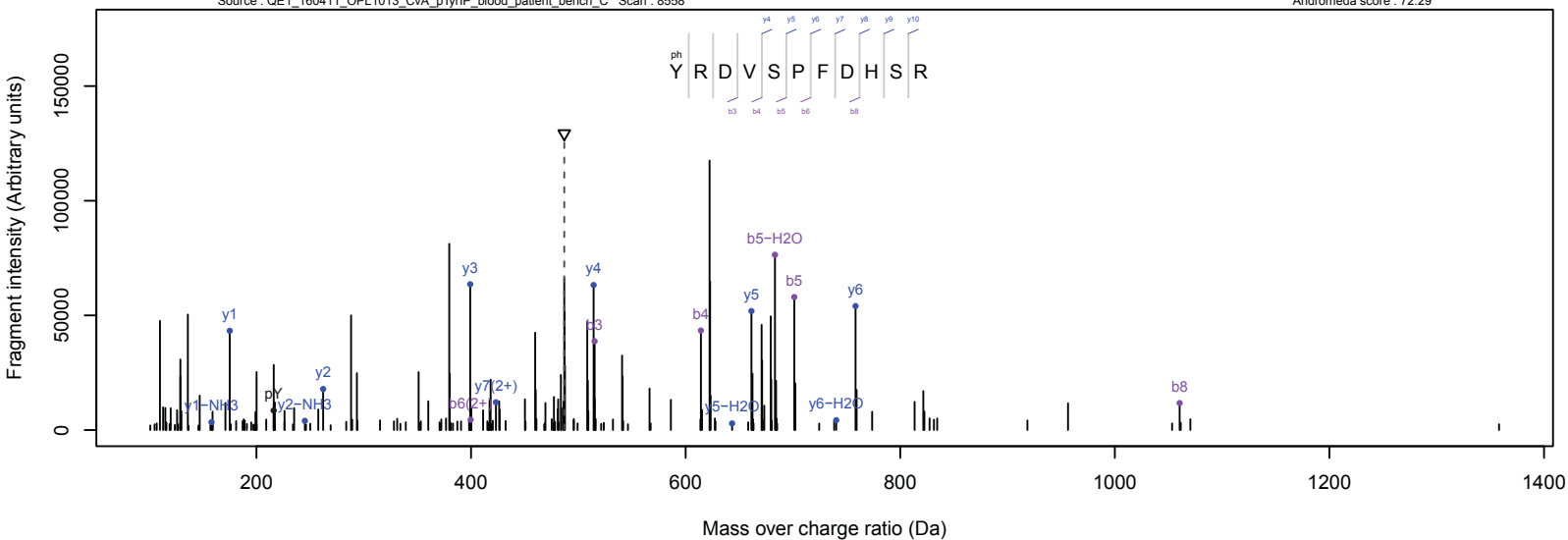




Y(ph)RDVSPFDHSR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 8558

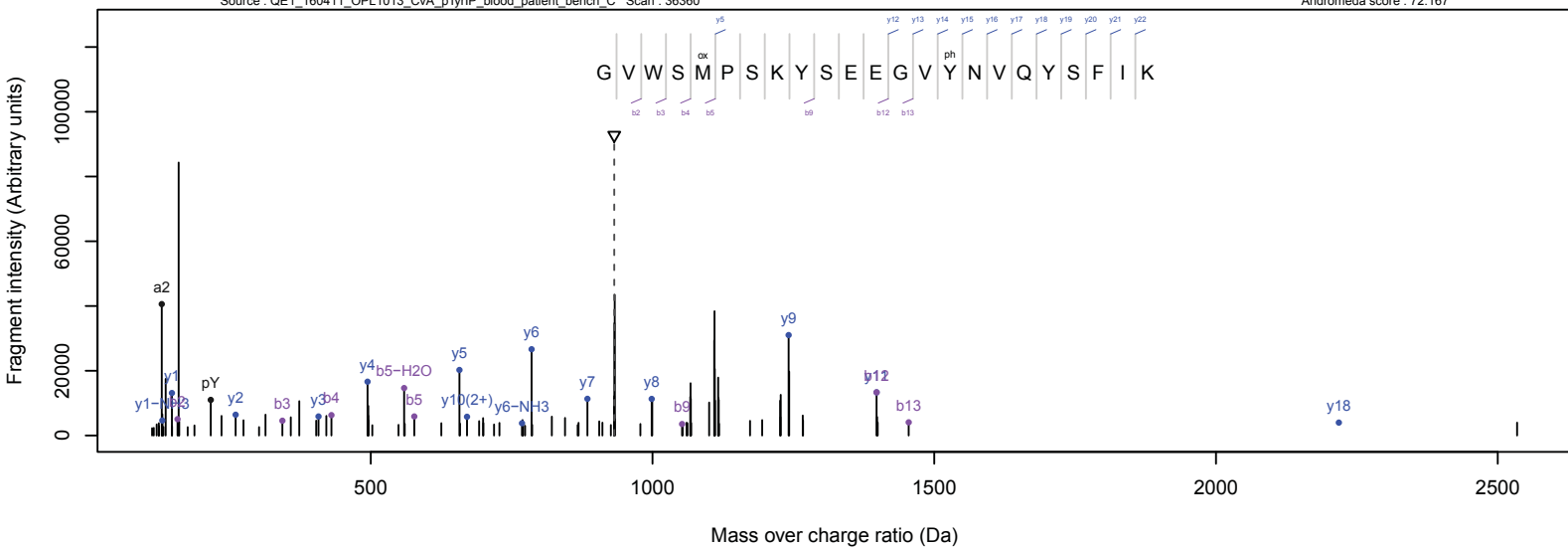
Andromeda score: 72.29



GVWSM(ox)PSKYSEEGV(ph)NVQYSFIK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 36360

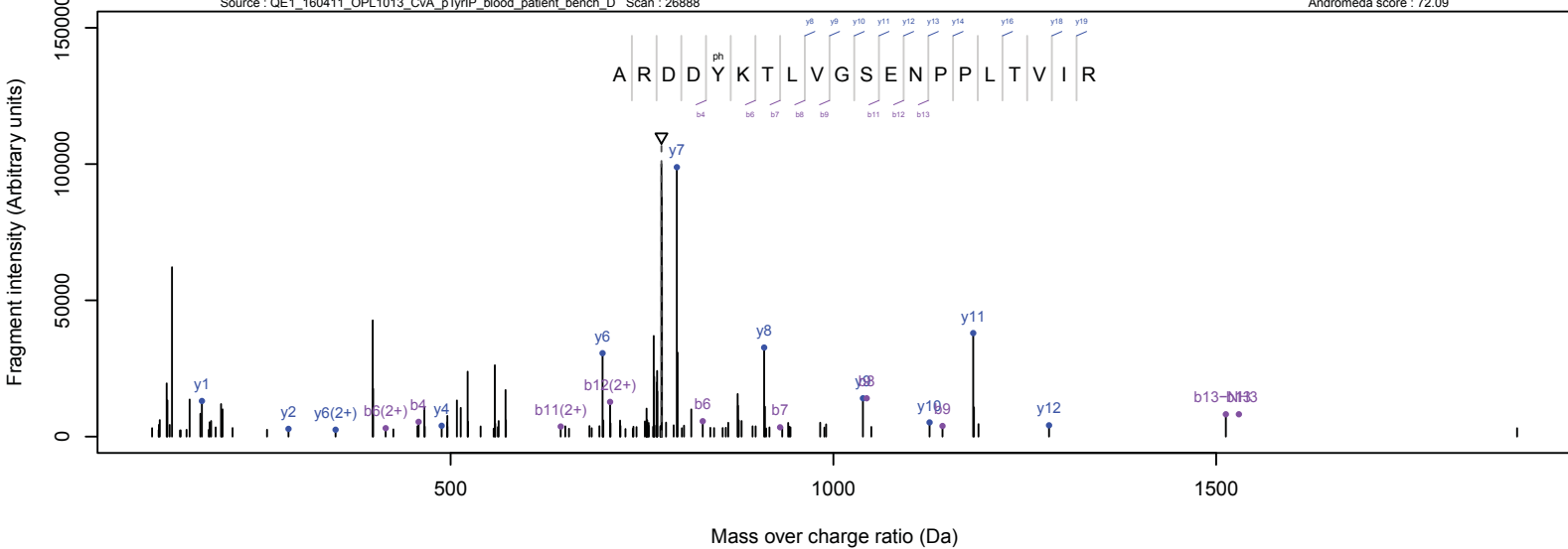
Andromeda score: 72.167



ARDDY(ph)KTLVGSENPPLTVIR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 26888

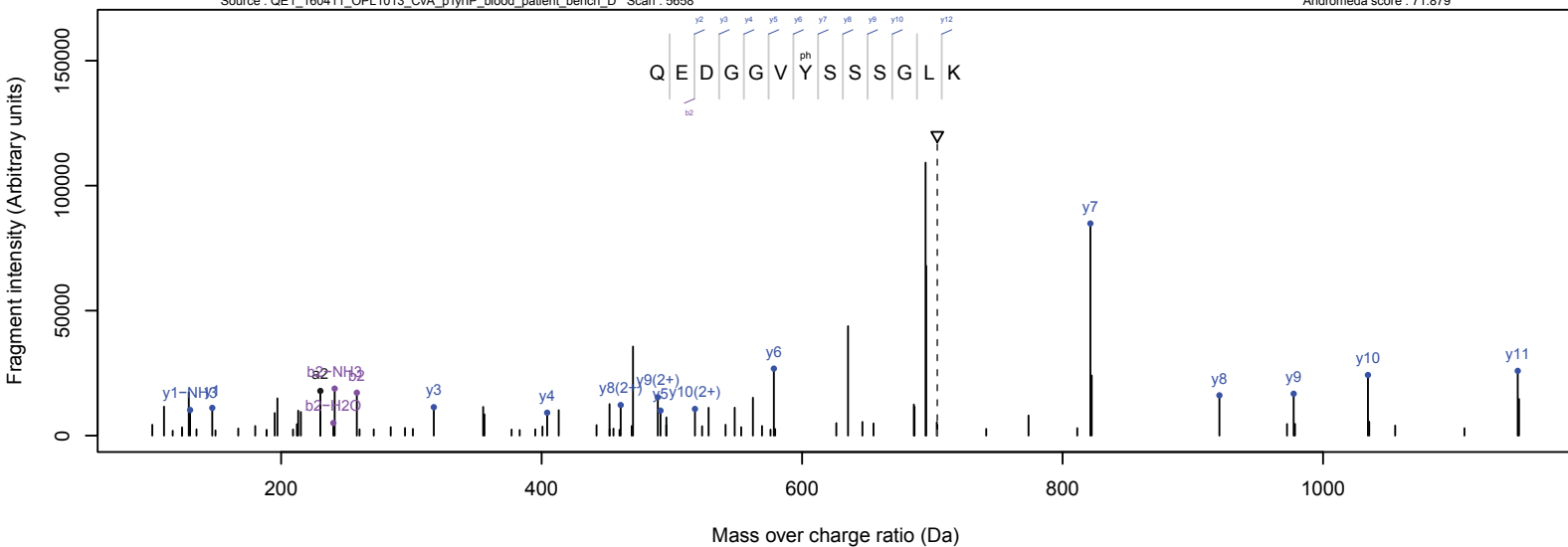
Andromeda score: 72.09



## QEDGGV(ph)SSGLK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5658

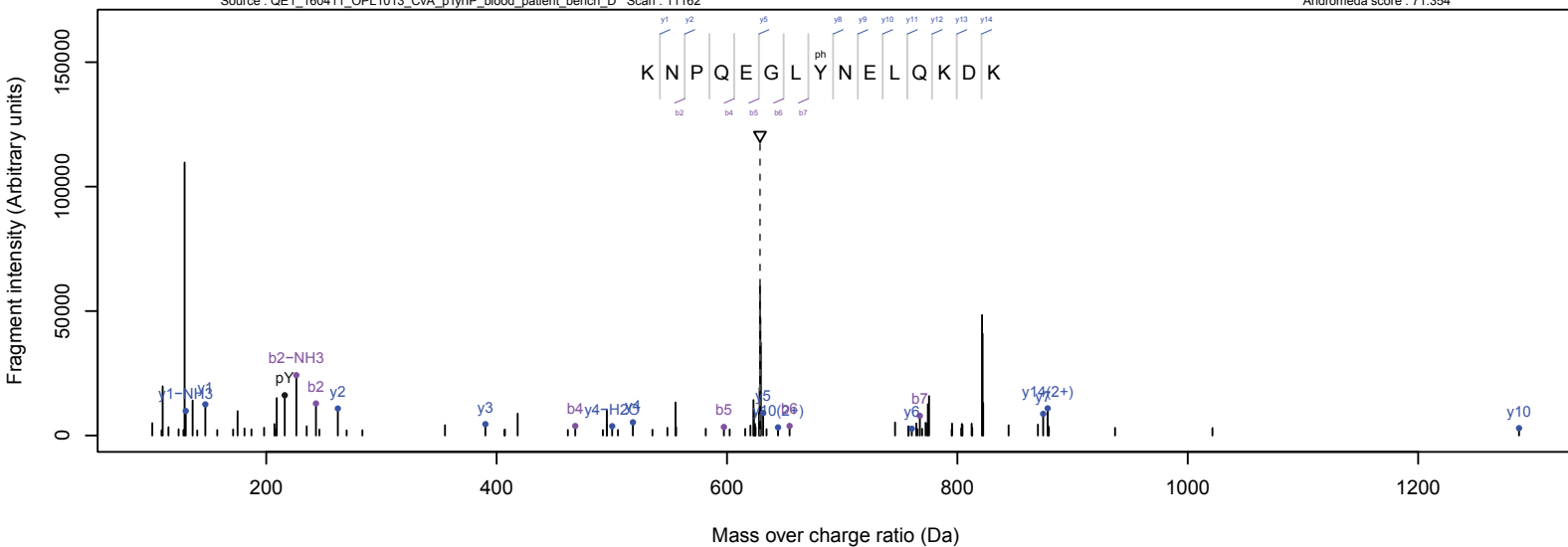
Andromeda score : 71.879



## KNPQEGLY(ph)NELQDK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11162

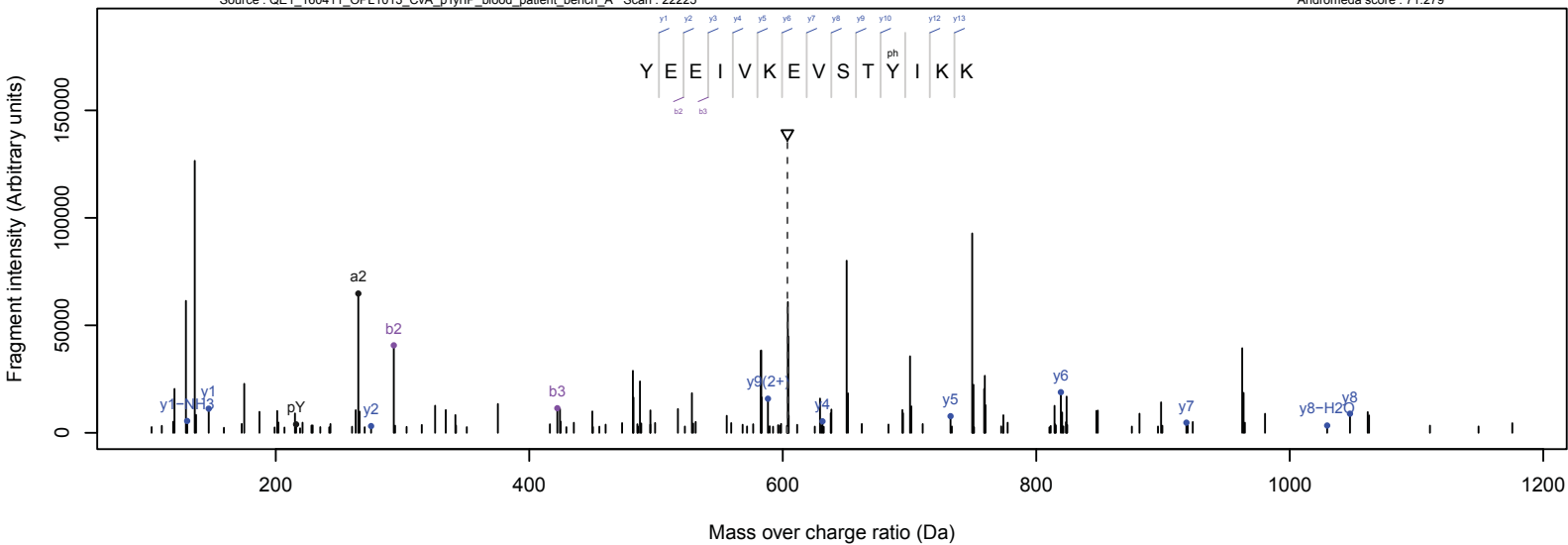
Andromeda score : 71.354



## YEEIVKEVSTY(ph)IKK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 22225

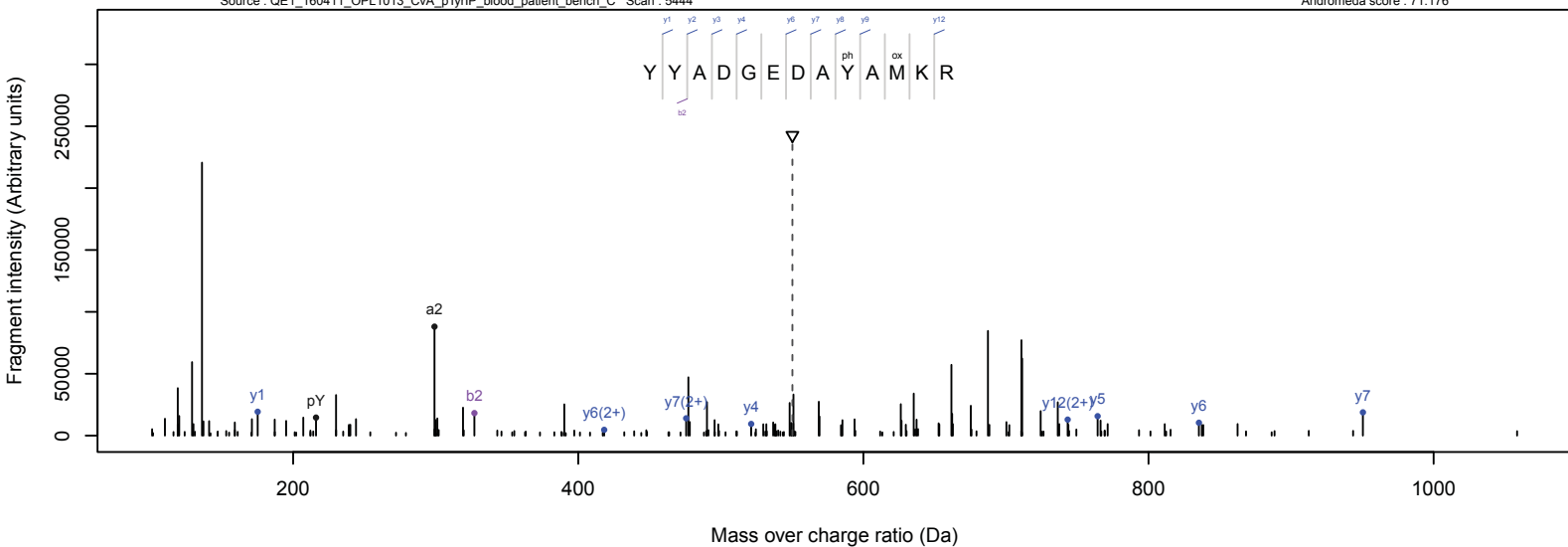
Andromeda score : 71.279



YYADGEDAYAM<sup>ph</sup>KR-3+

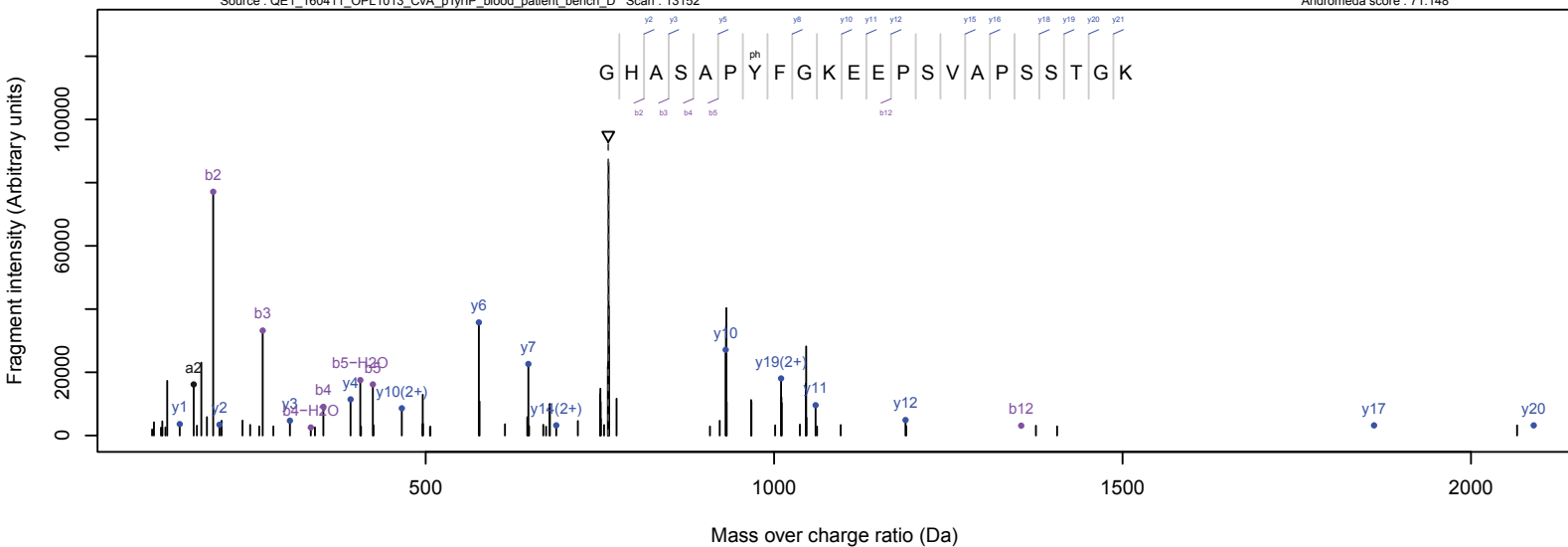
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 5444

Andromeda score: 71.176

GHASAPY<sup>ph</sup>FGKEEPSVAPSSSTGK-3+

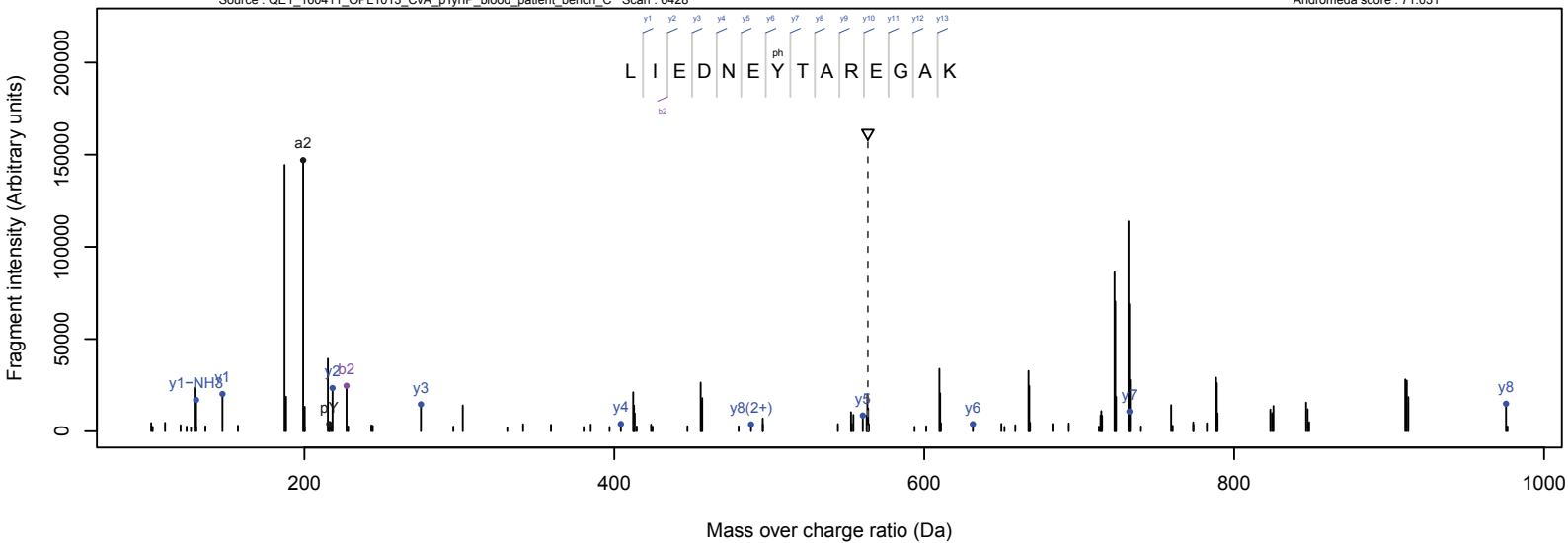
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 13152

Andromeda score: 71.148

LIEDNEY<sup>ph</sup>TAREGAK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 6428

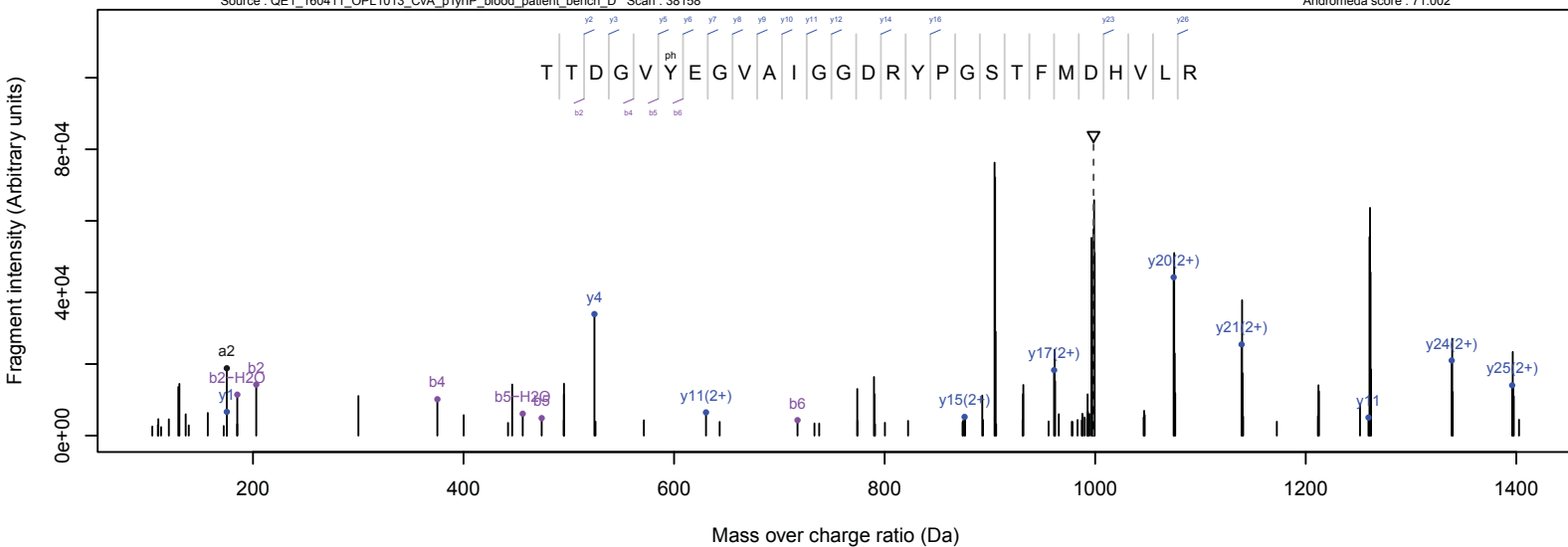
Andromeda score: 71.031



TTDGVY(ph)EGVAIGGDRYPGSTFMDHVL R-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 38158

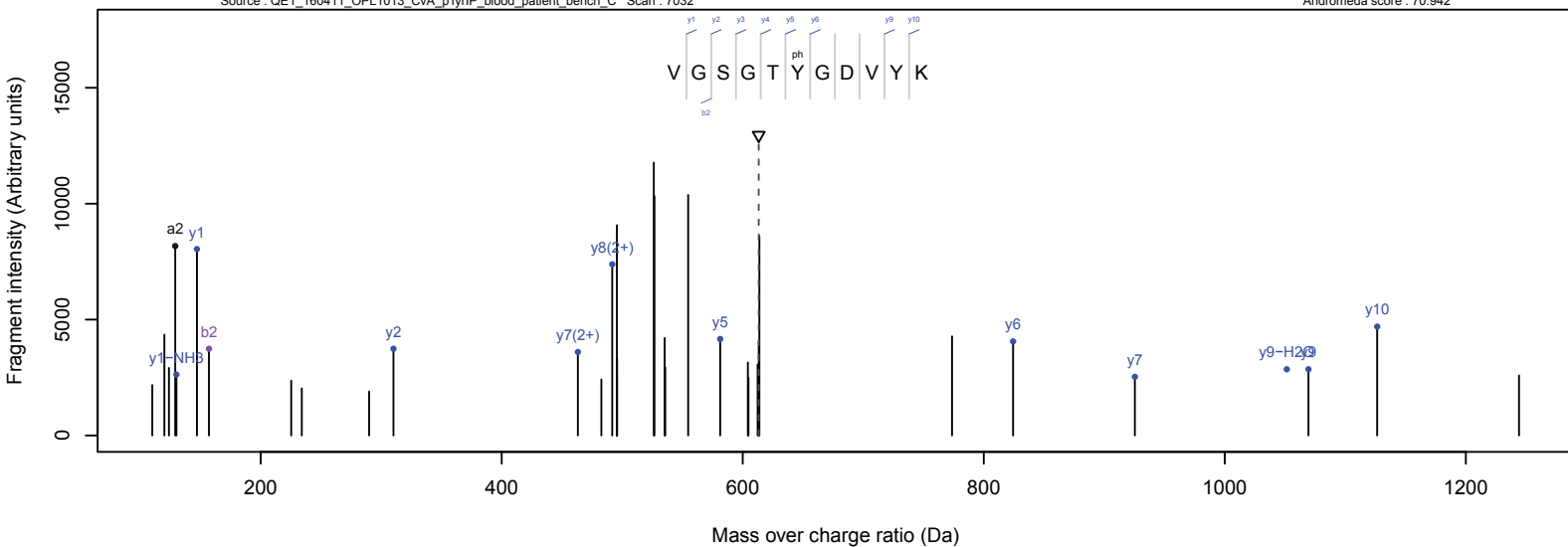
Andromeda score : 71.002



VGSGTY(ph)GDVYK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 7032

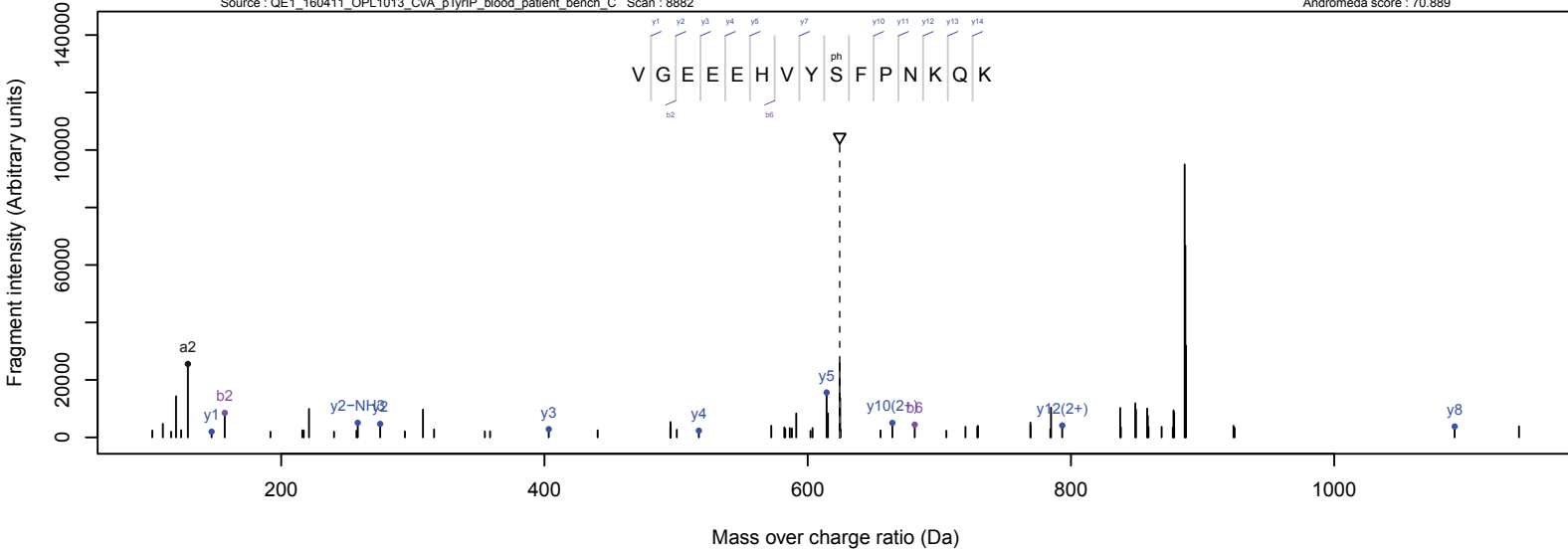
Andromeda score : 70.942



VGEEHVYS(ph)FPNKQK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 8882

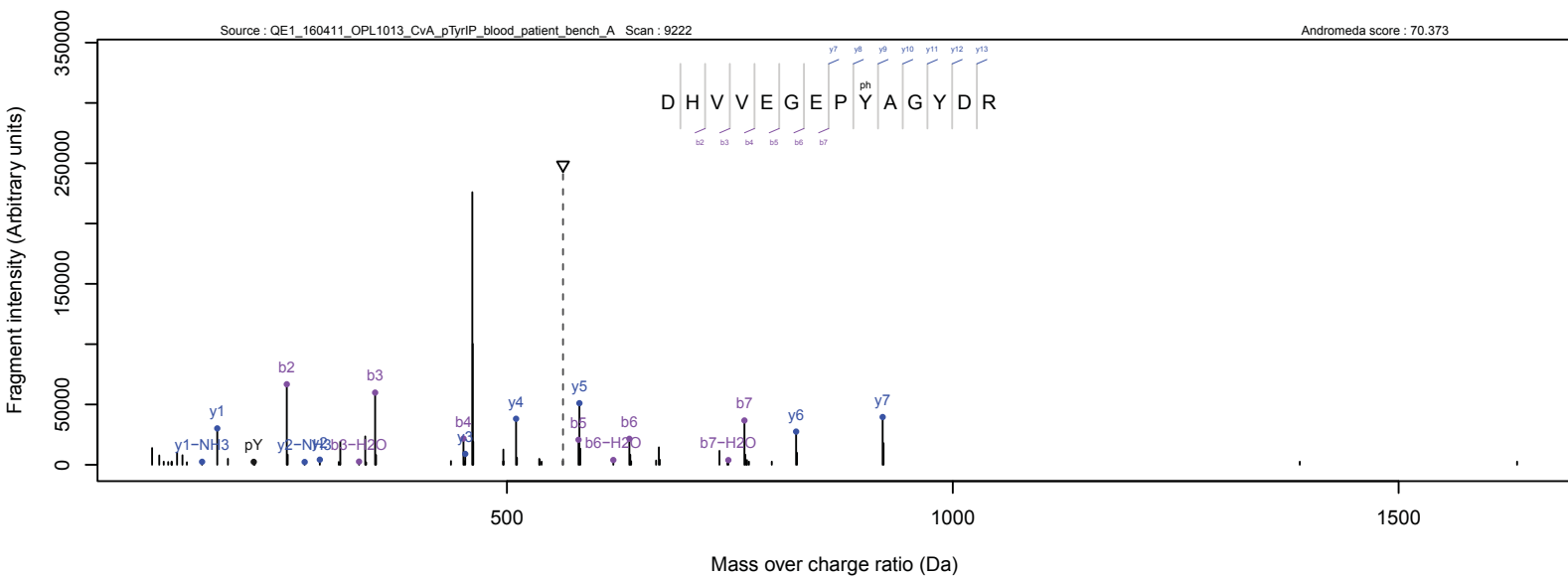
Andromeda score : 70.889



DHVVEGEPY(ph)AGYDR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 9222

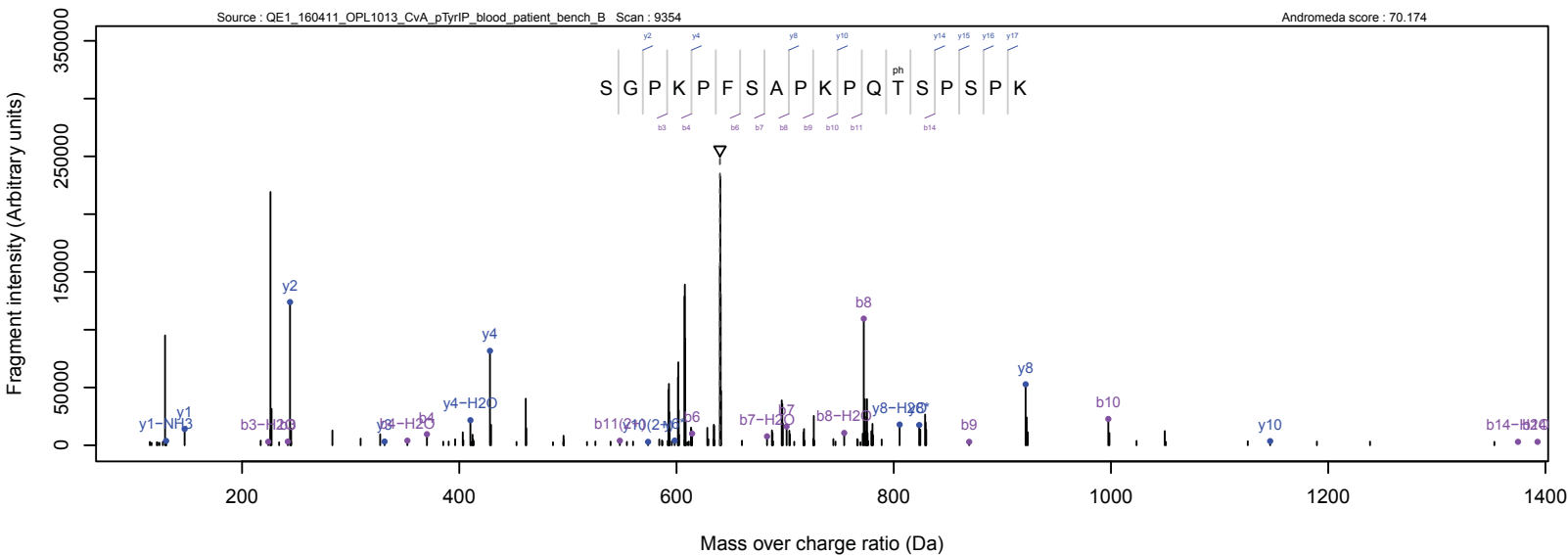
Andromeda score : 70.373



SGPKPFSAPKQT(ph)SPSPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 9354

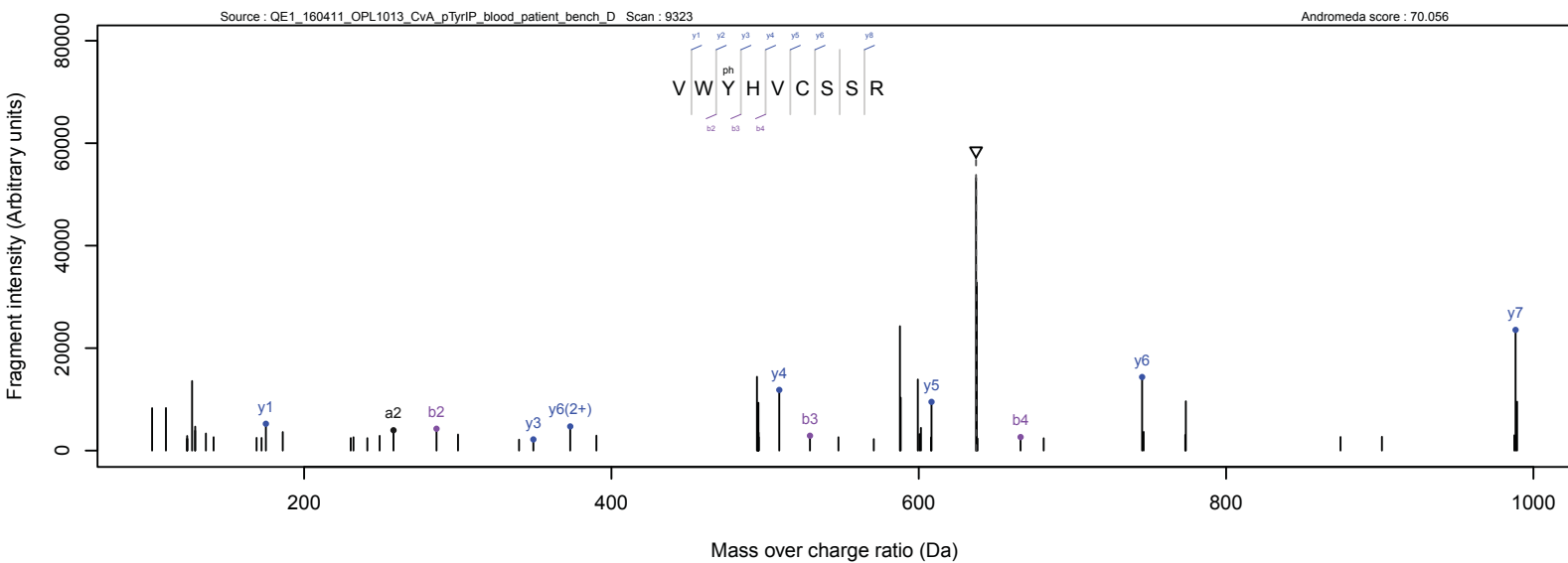
Andromeda score : 70.174



VWY(ph)HVCSSR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 9323

Andromeda score : 70.056

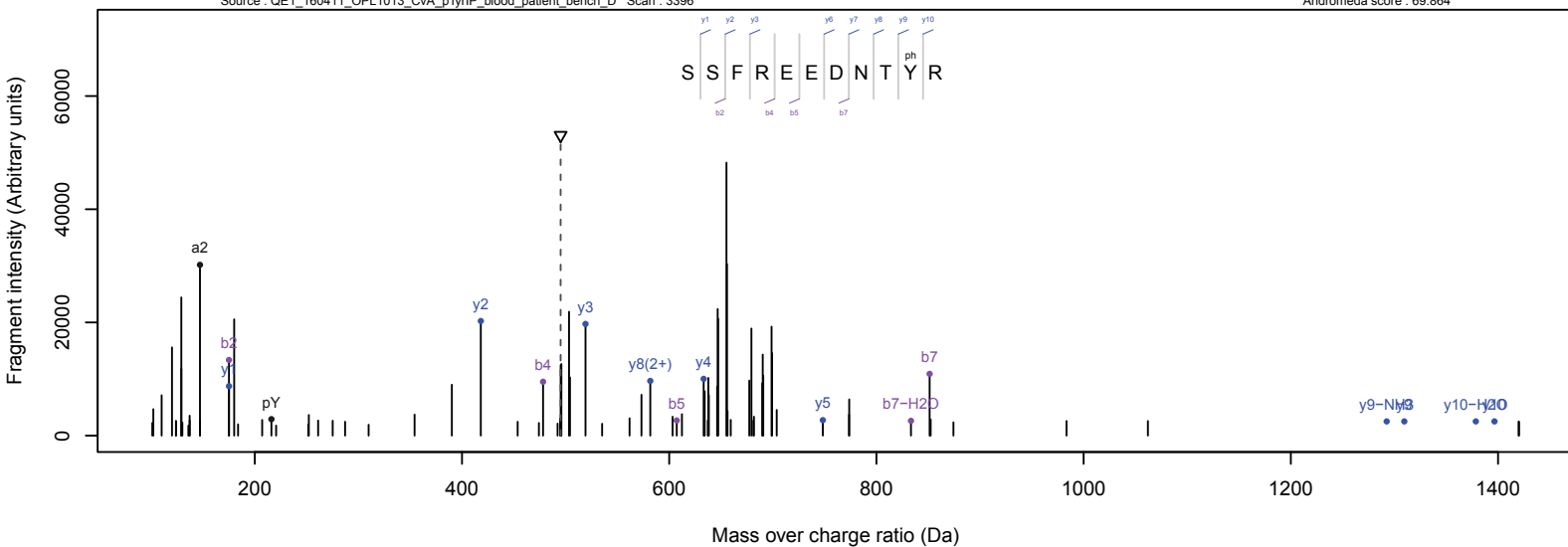




## SSFREEDNTY(ph)R-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 3396

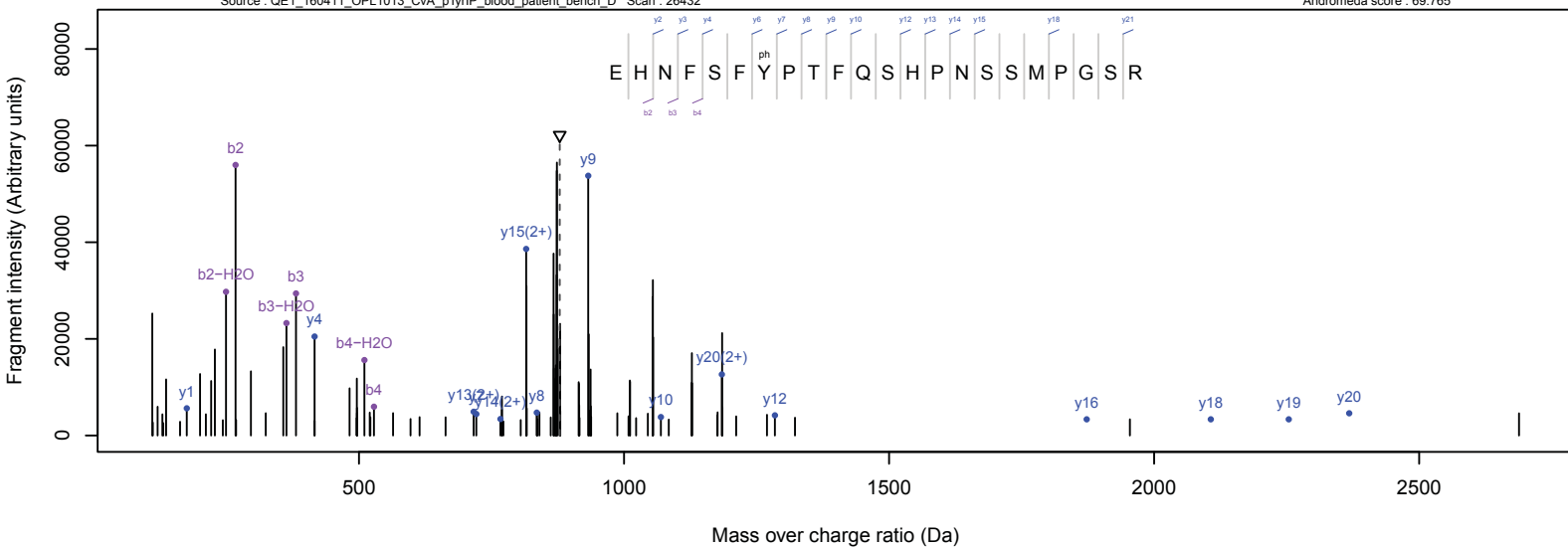
Andromeda score: 69.864



## EHNFSFY(ph)PTFQSHPNSSMPGSR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 26432

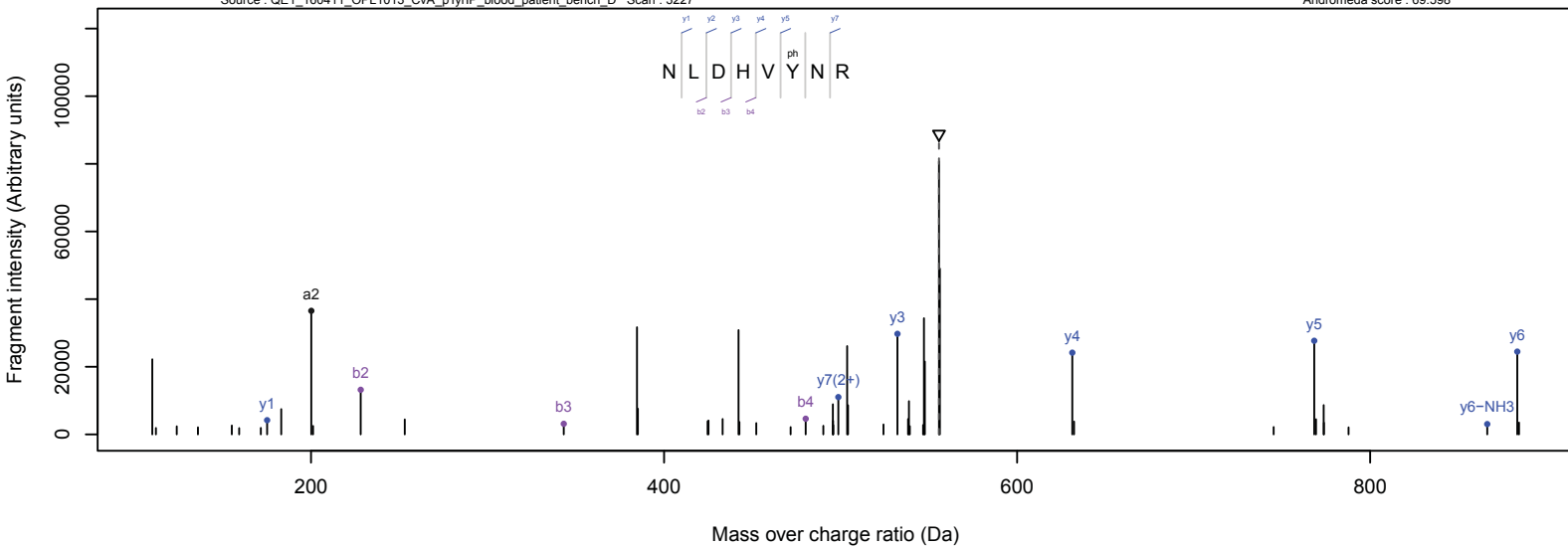
Andromeda score: 69.765



## NLDHVY(ph)NR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 3227

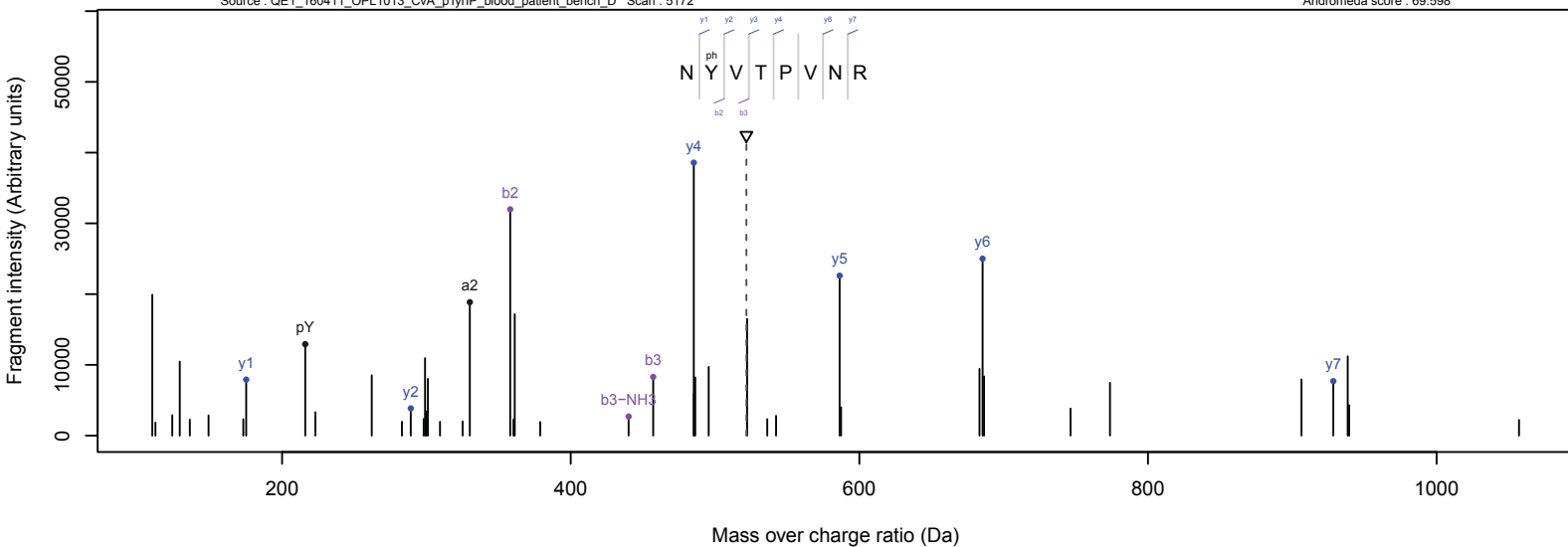
Andromeda score: 69.598



NY(ph)VTPVNR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5172

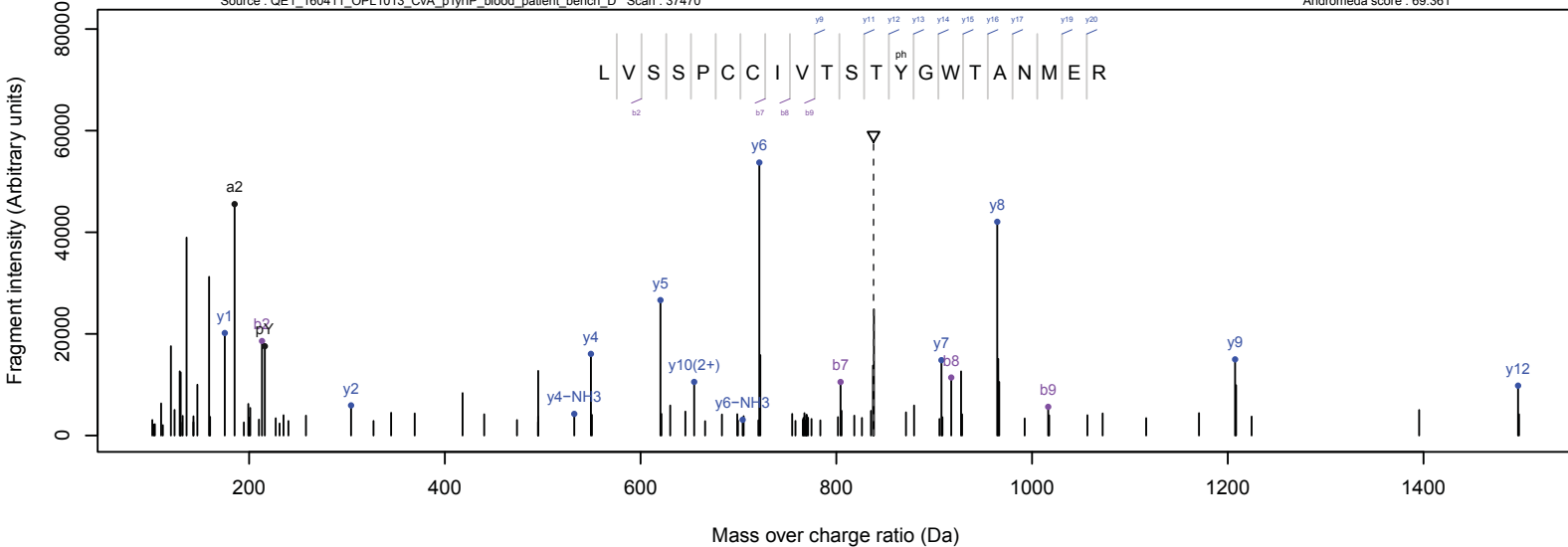
Andromeda score : 69.598



LVSSPCCIVTSTY(ph)GWTANMER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 37470

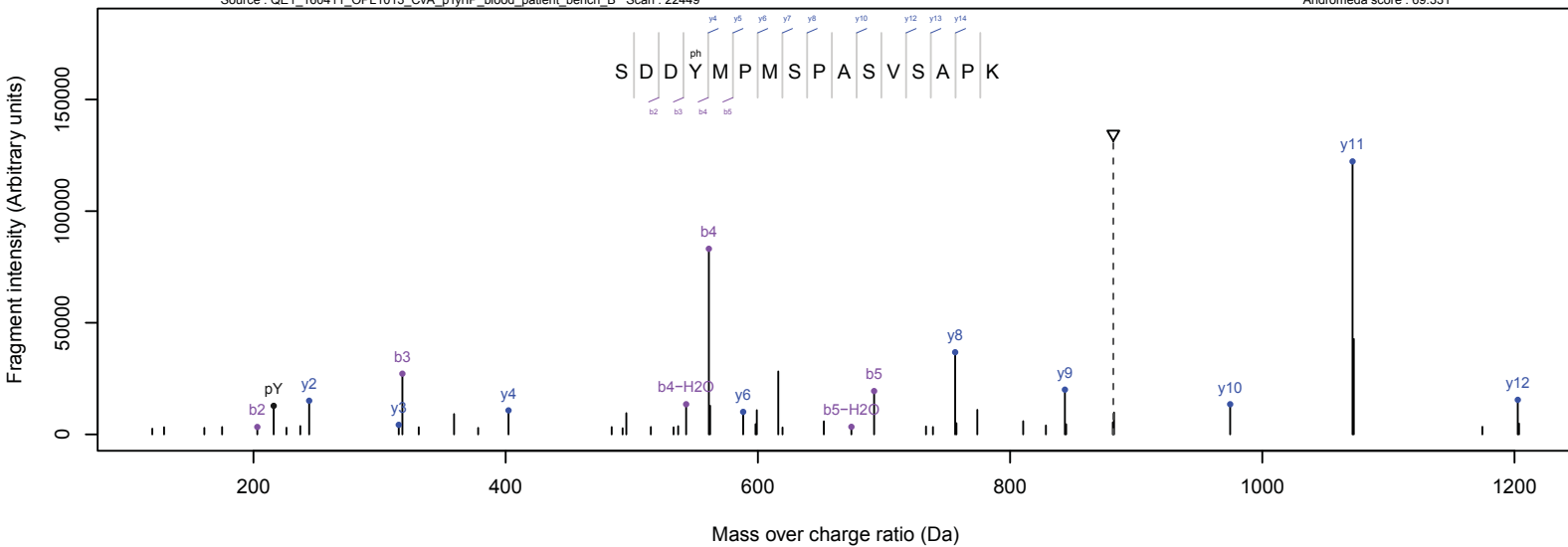
Andromeda score : 69.361



SDDY(ph)MPMPASVSAPK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 22449

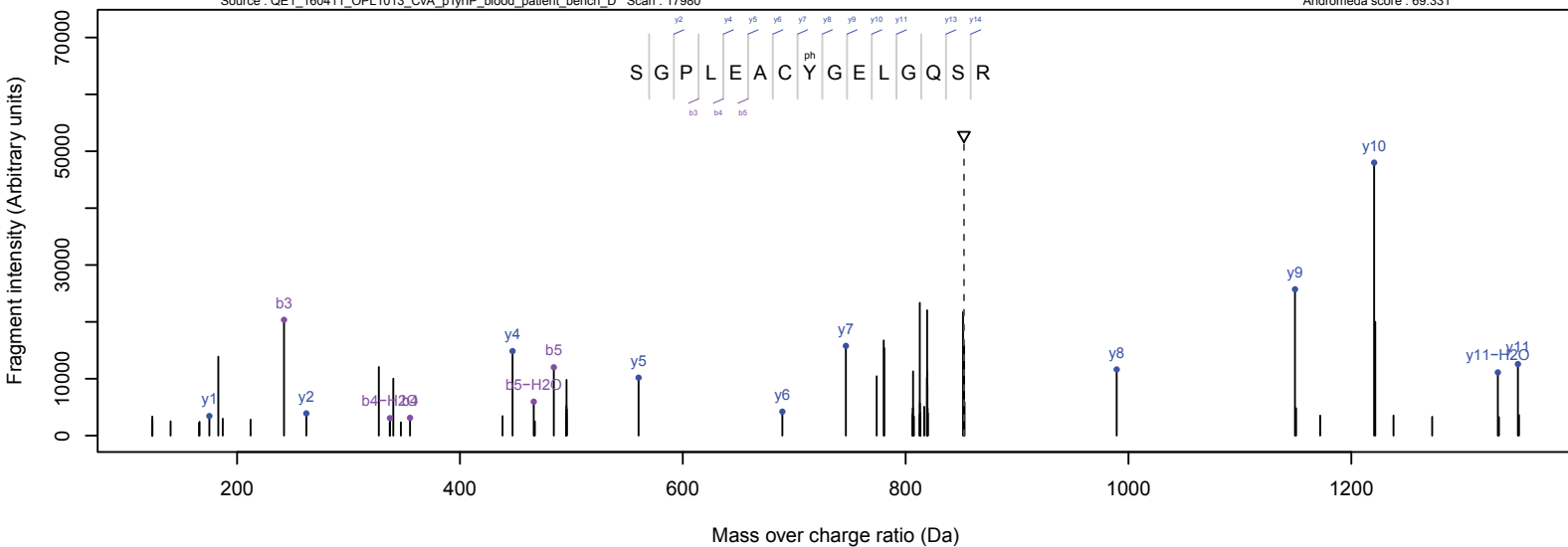
Andromeda score : 69.331



## SGPLEACY(ph)GELGQSR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17980

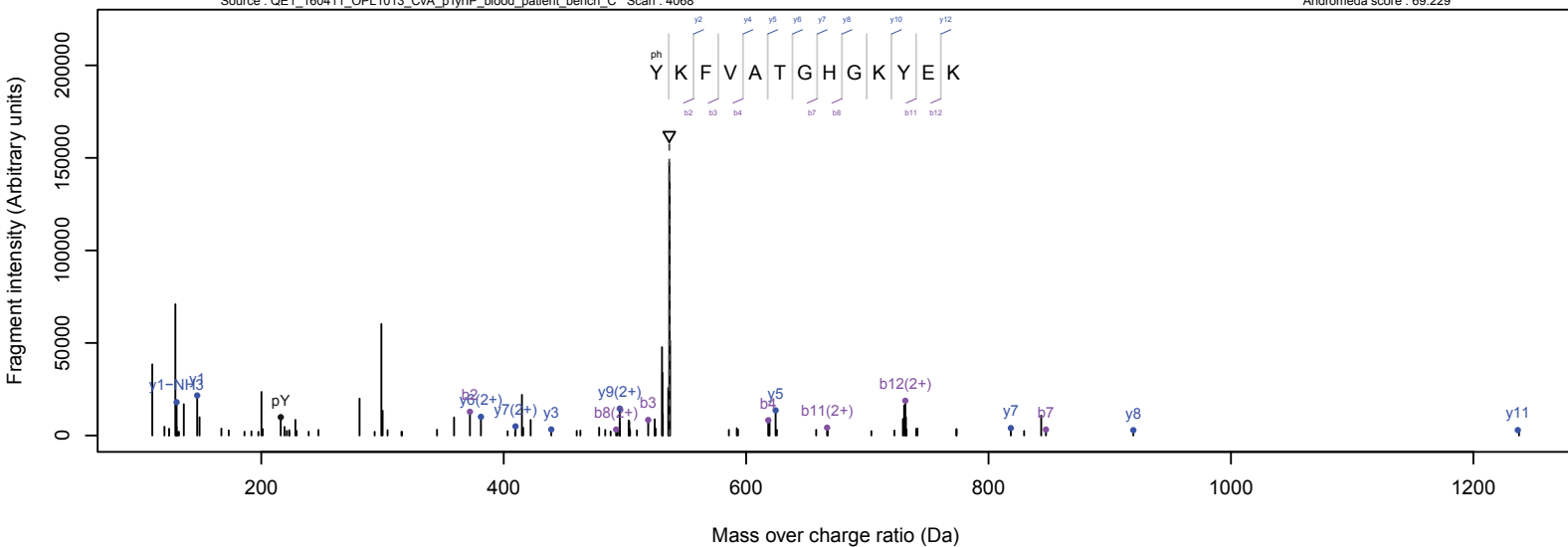
Andromeda score : 69.331



## Y(ph)KFVATGHGKYEK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 4068

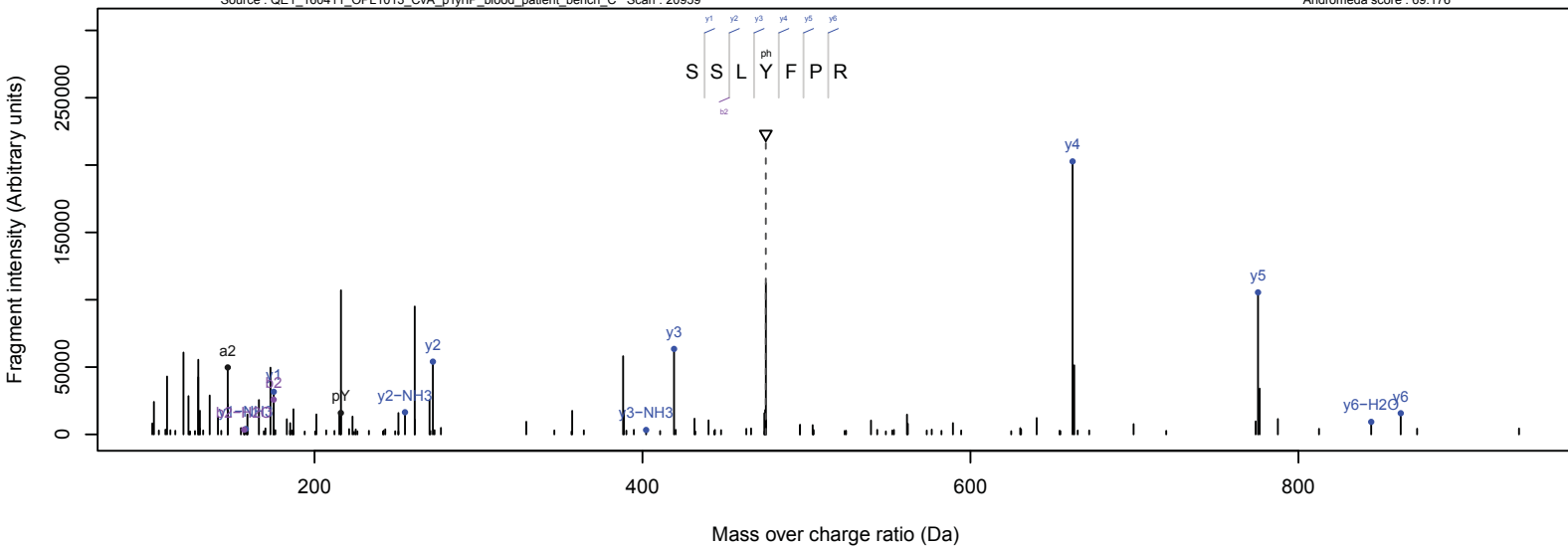
Andromeda score : 69.229



## SSLY(ph)FPR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 20959

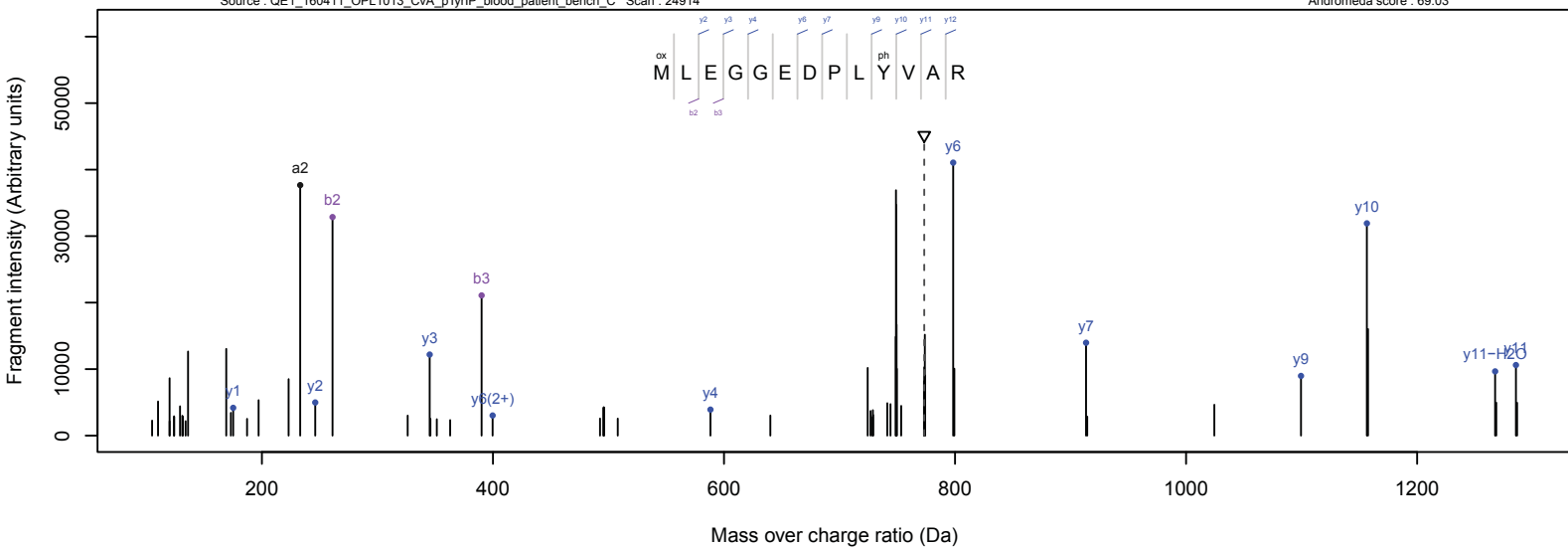
Andromeda score : 69.176



## M(ox)LEGGEDPLY(ph)VAR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 24914

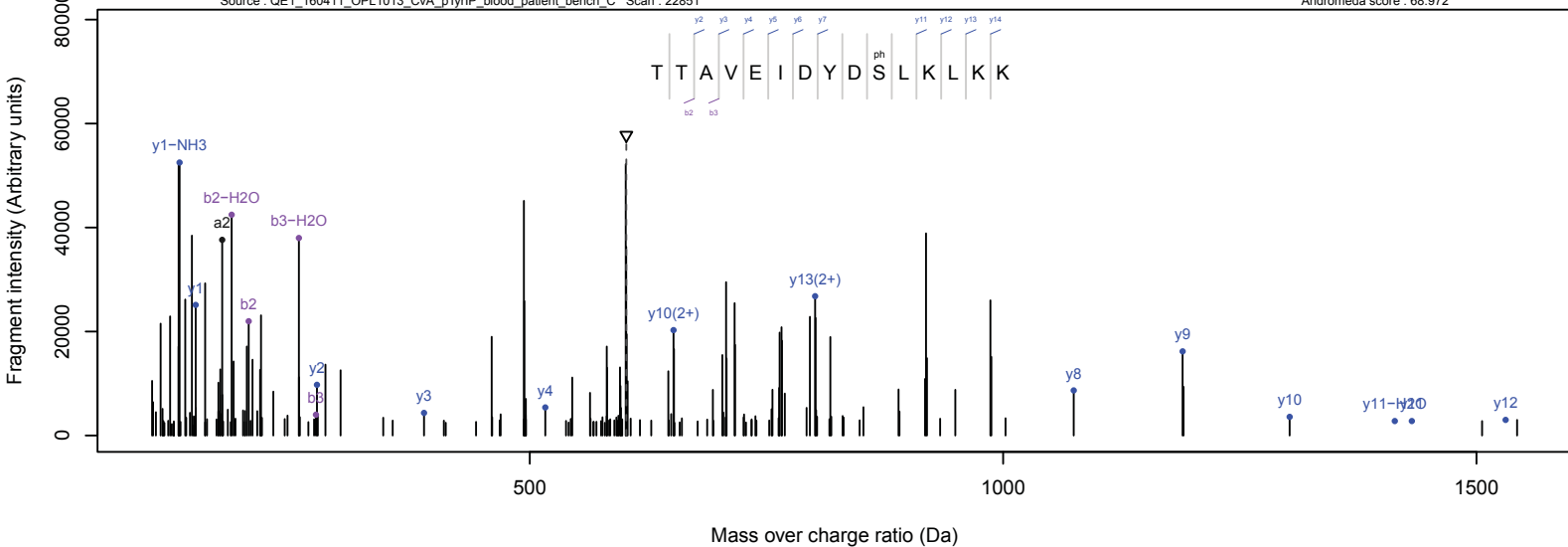
Andromeda score : 69.03



## TTAVEIDYDS(ph)LK LK K-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 22851

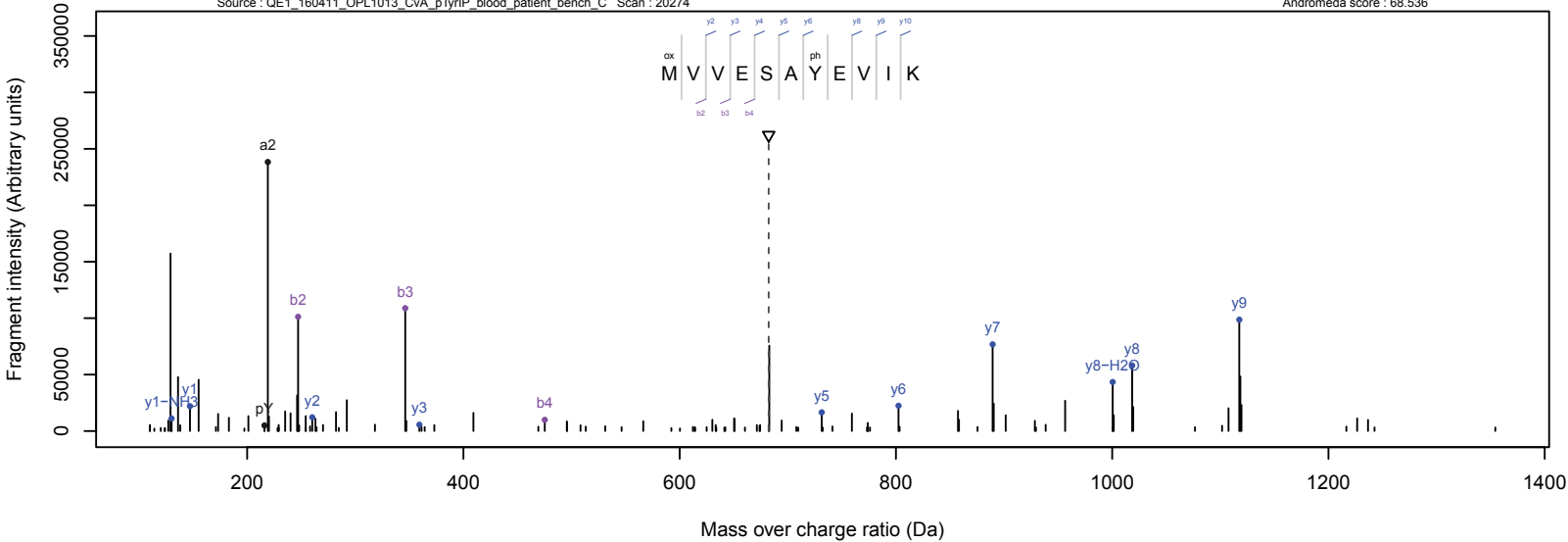
Andromeda score : 68.972



## M(ox)VVESAY(ph)EVIK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 20274

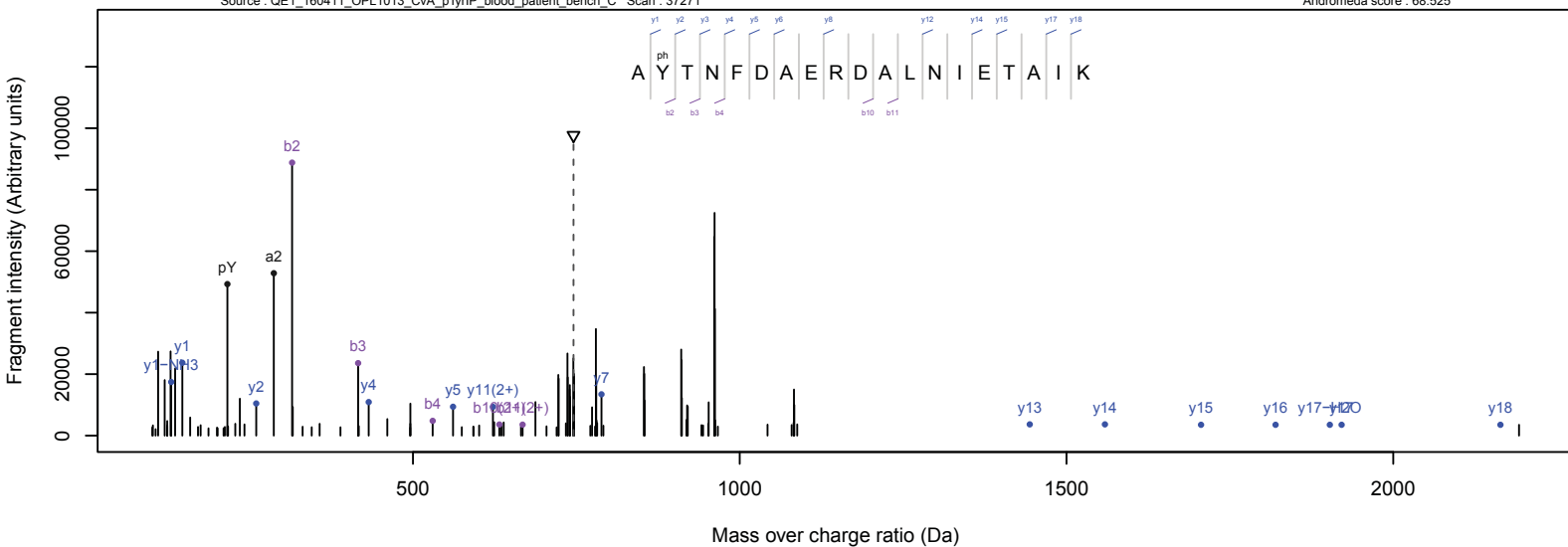
Andromeda score : 68.536



**AY(ph)TNFDAERDALNIETAIK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 37271

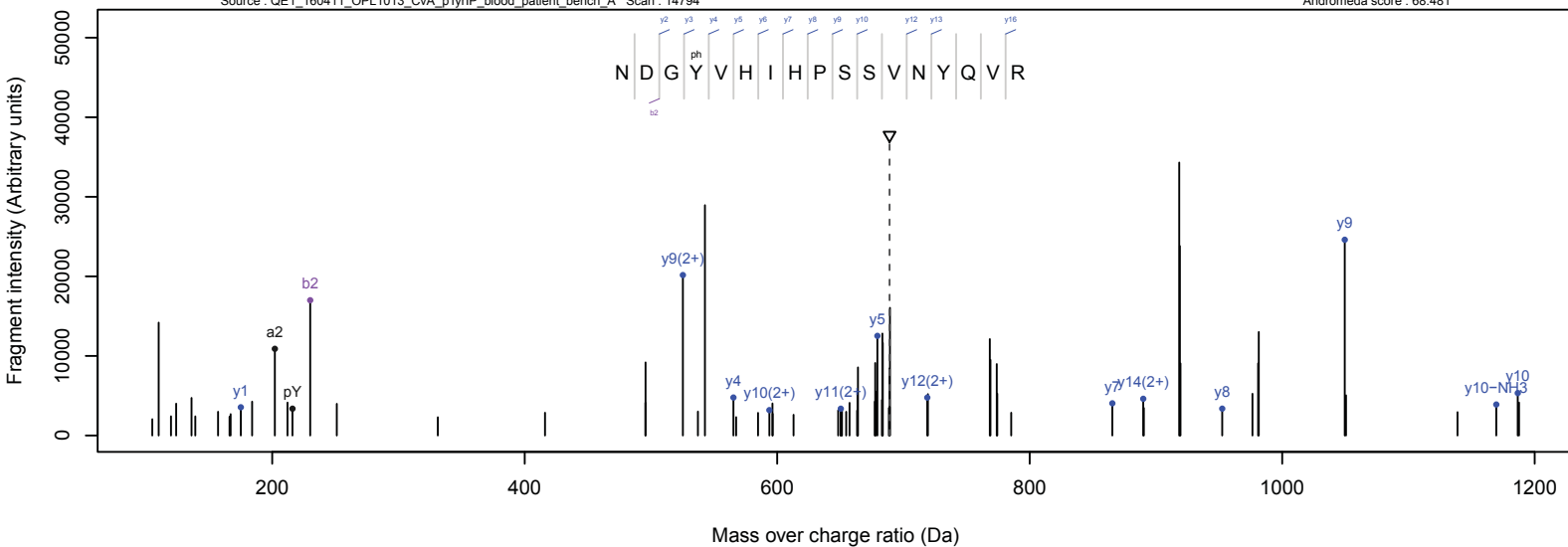
Andromeda score : 68.525



**NDGY(ph)VHIHPSSVNYQVR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 14794

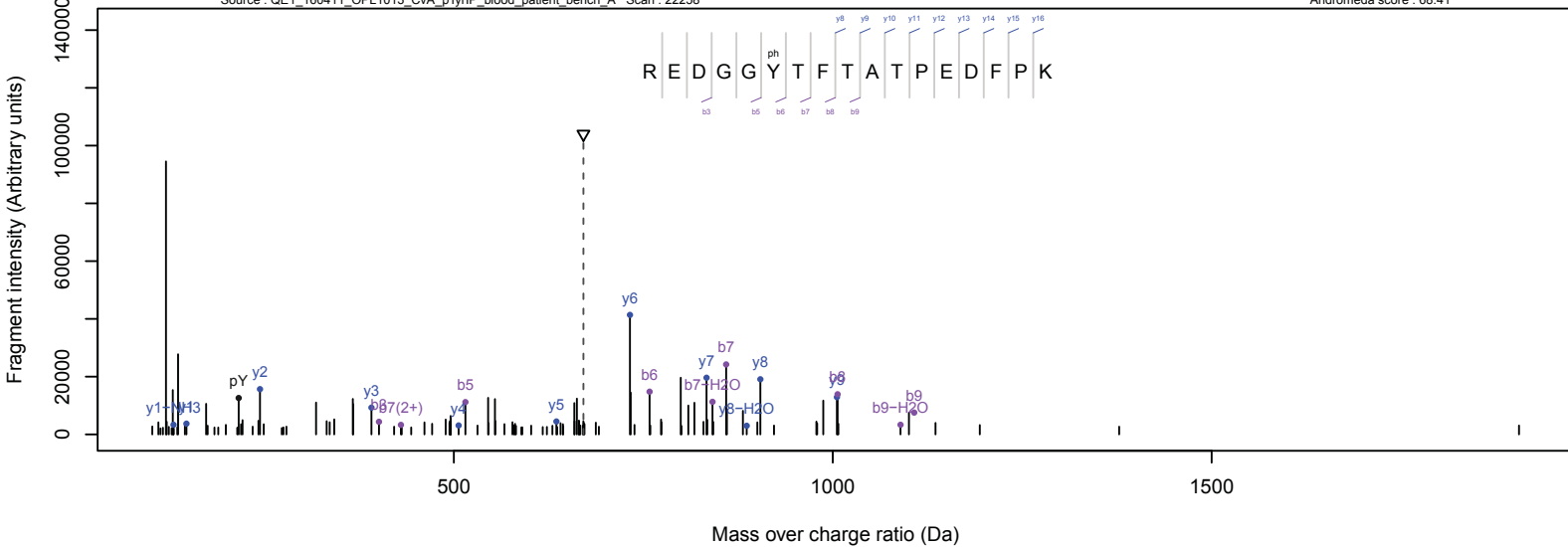
Andromeda score : 68.481



**REDGGY(ph)TFTATPEDFPK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 22258

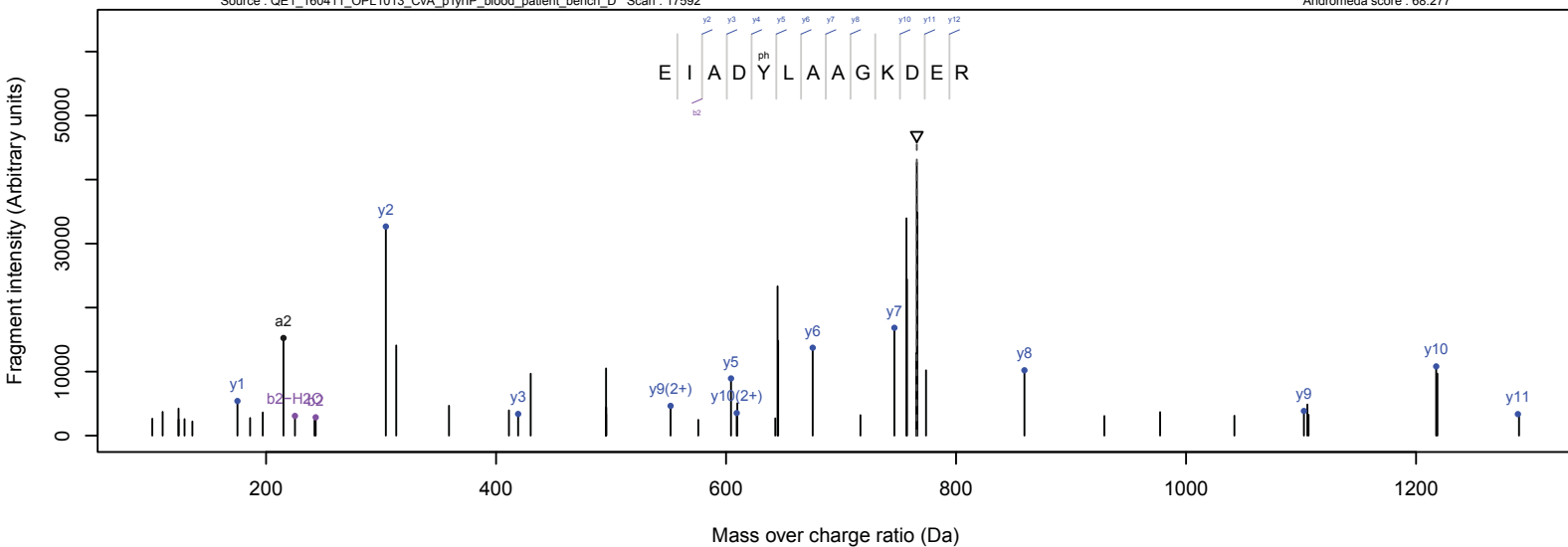
Andromeda score : 68.41



EIADY(ph)LAAGKDER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17592

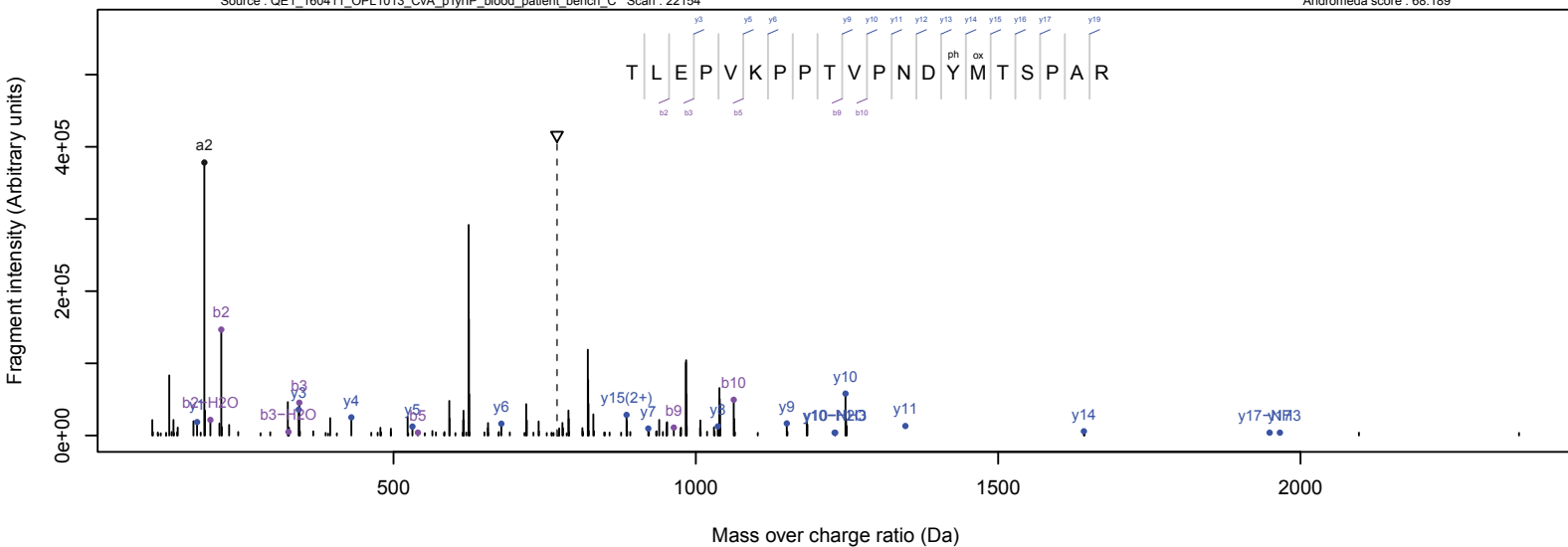
Andromeda score : 68.277



TLEPVKPTVPNDY(ph)M(ox)TSPAR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 22154

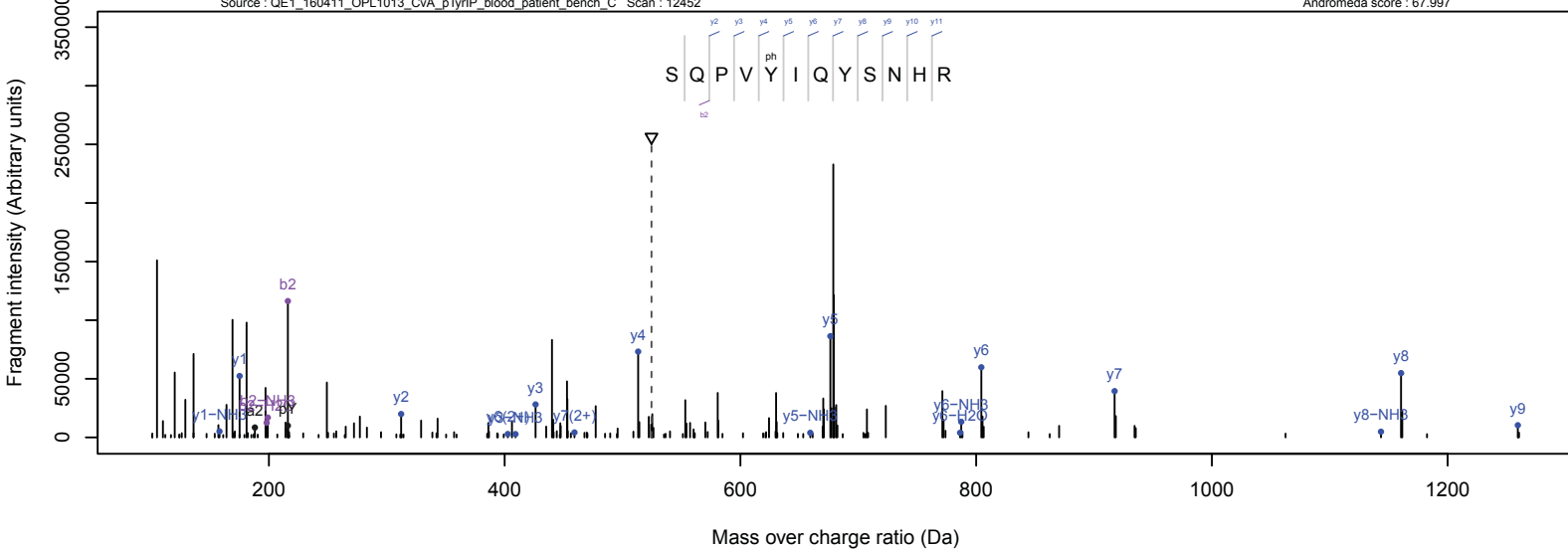
Andromeda score : 68.189



SQPVYI QYSNHR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 12452

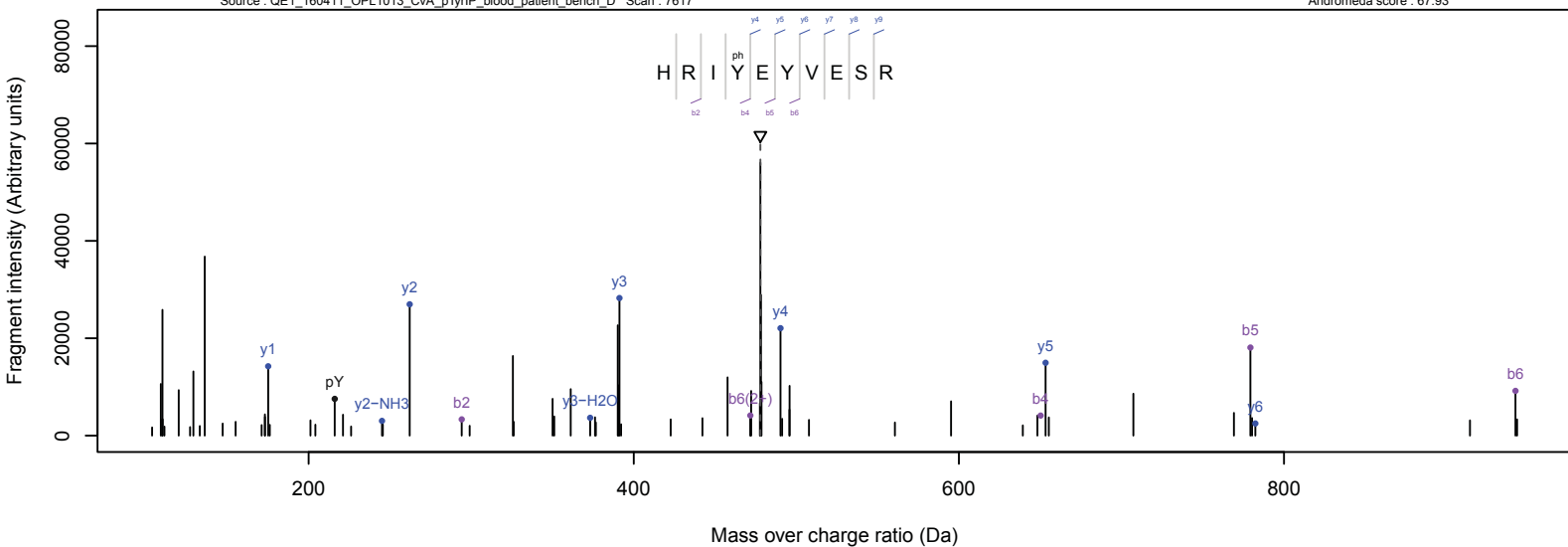
Andromeda score : 67.997



HRIY(ph)EYVESR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 7617

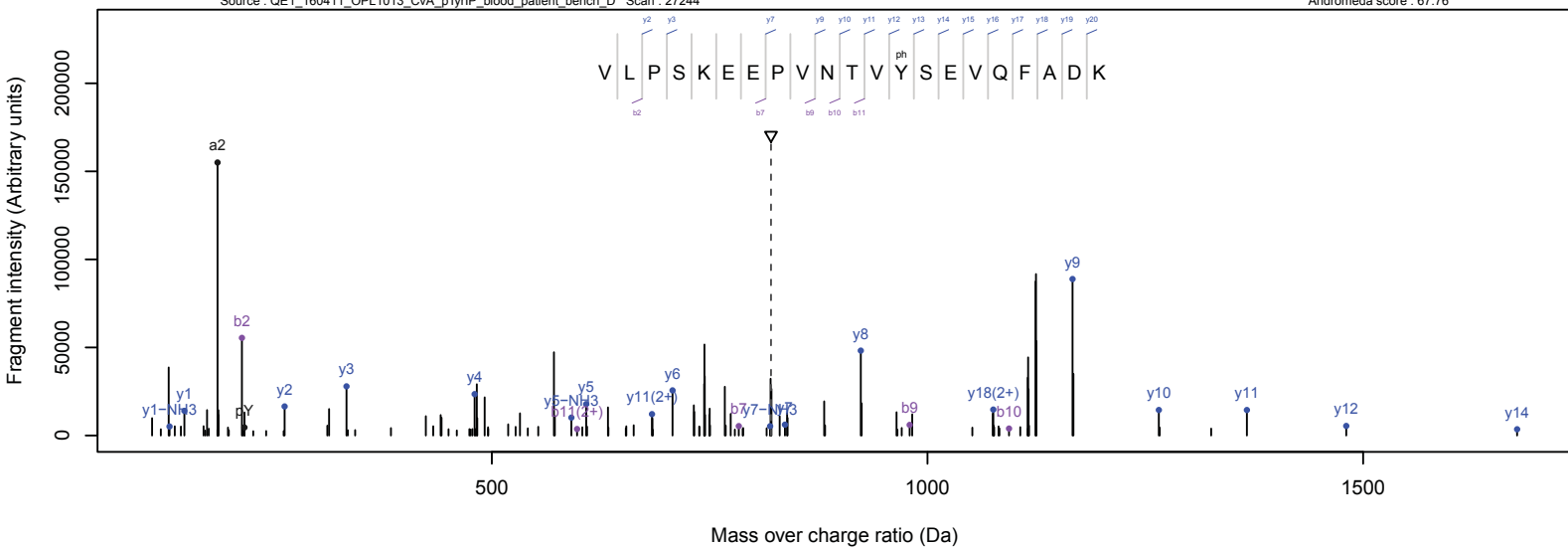
Andromeda score: 67.93



VLPKEEPVNTVY(ph)SEVQFADK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 27244

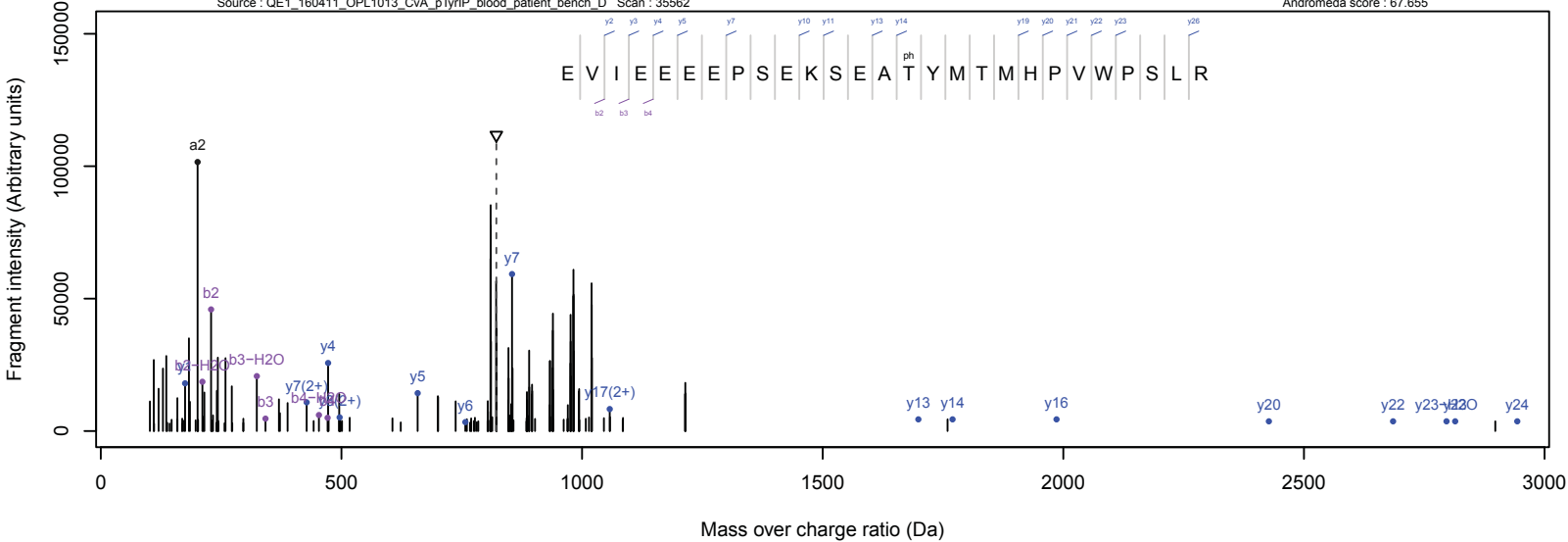
Andromeda score: 67.76



EVIEEPEKSEAT(ph)YMTMHPVWPSLR-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 35562

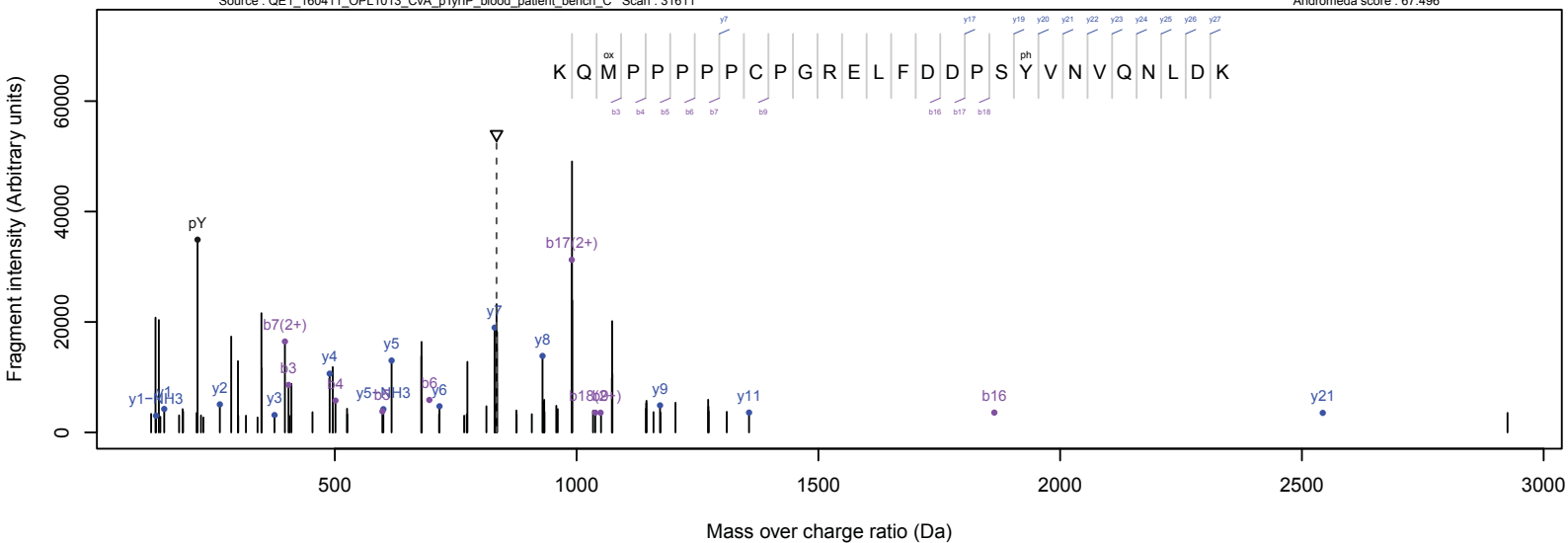
Andromeda score: 67.655



**KQM(ox)PPPPPCGRELFDDPSY(ph)VNVQNLDK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 31611

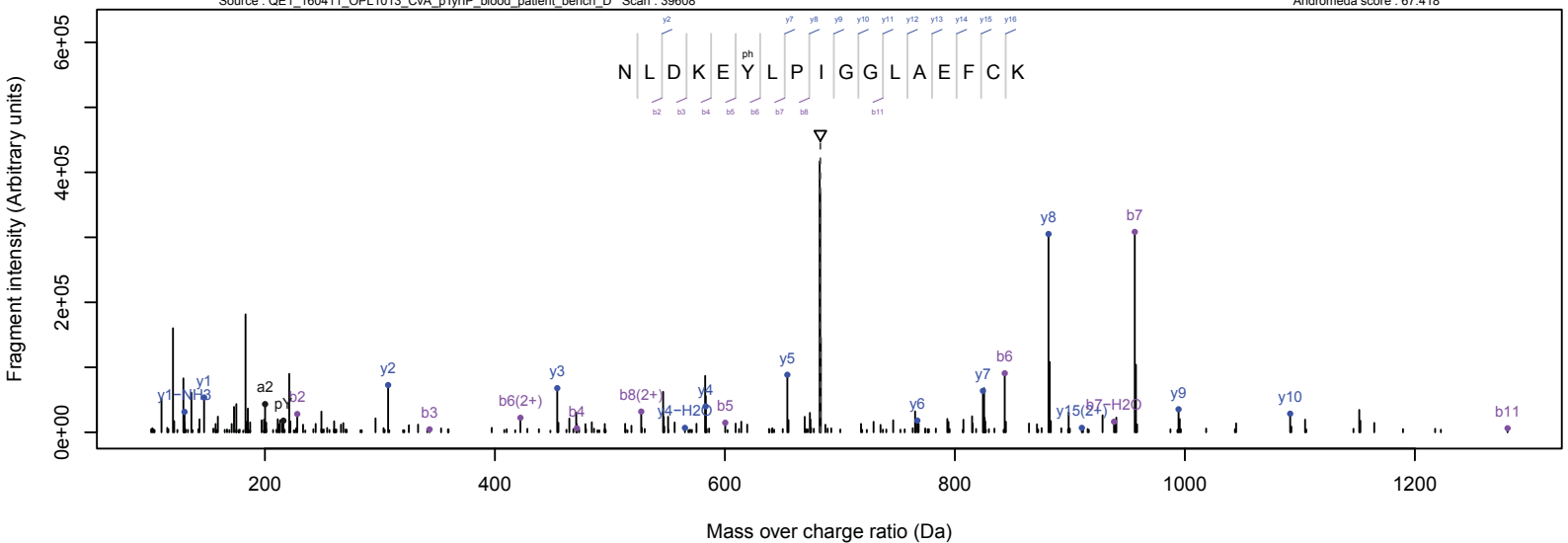
Andromeda score : 67.496



**NLDKEY(ph)LPIGGLAEFCK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 39608

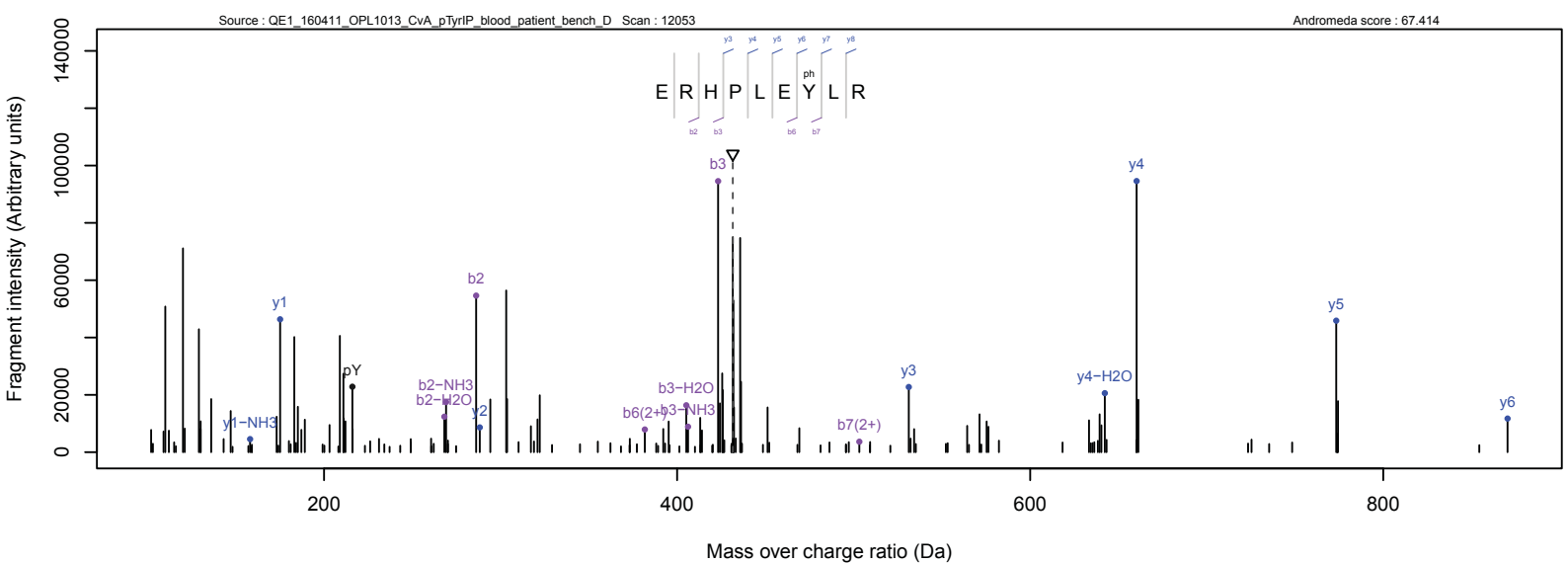
Andromeda score : 67.418



**ERHPLEY(ph)LR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12053

Andromeda score : 67.414

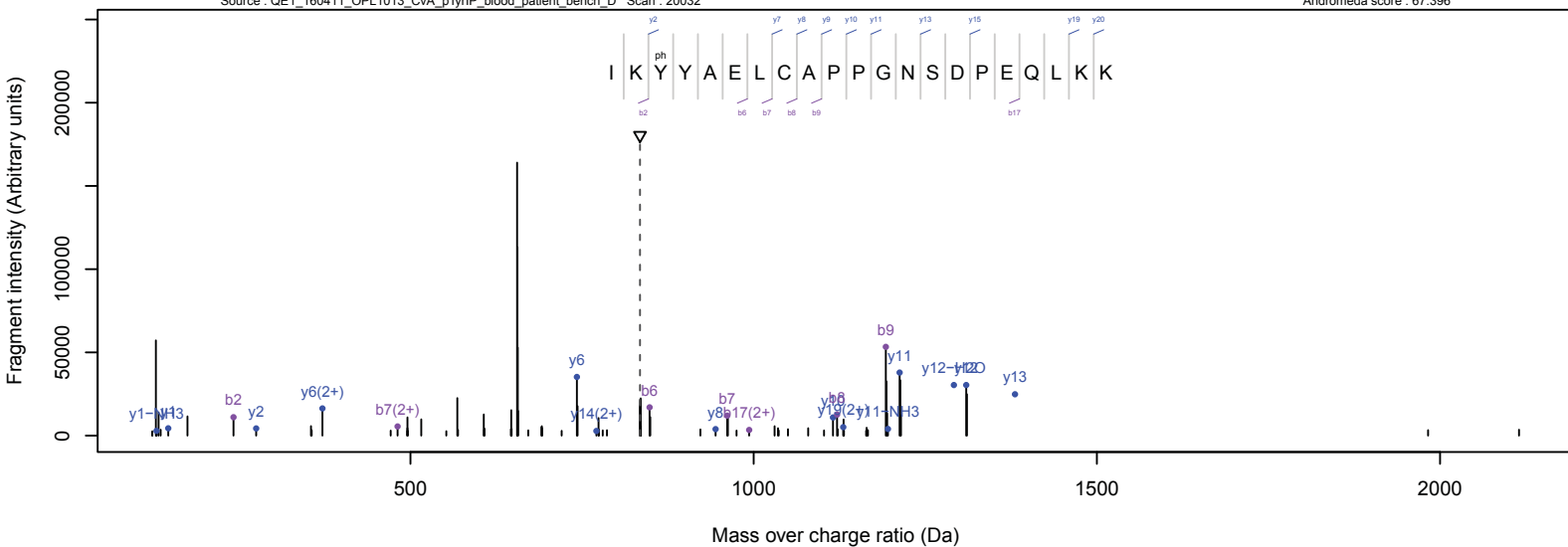




**IKY(ph)YAELCAPPGNSDPEQLKK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 20032

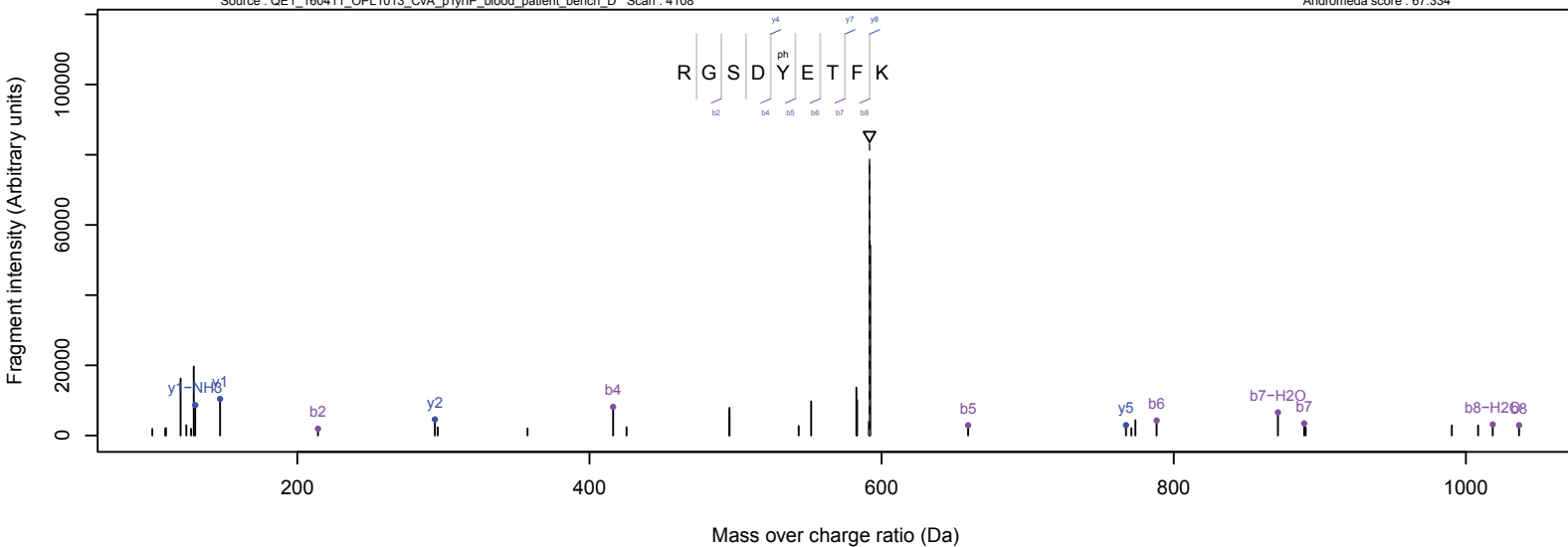
Andromeda score : 67.396



**RGSDY(ph)ETFK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 4108

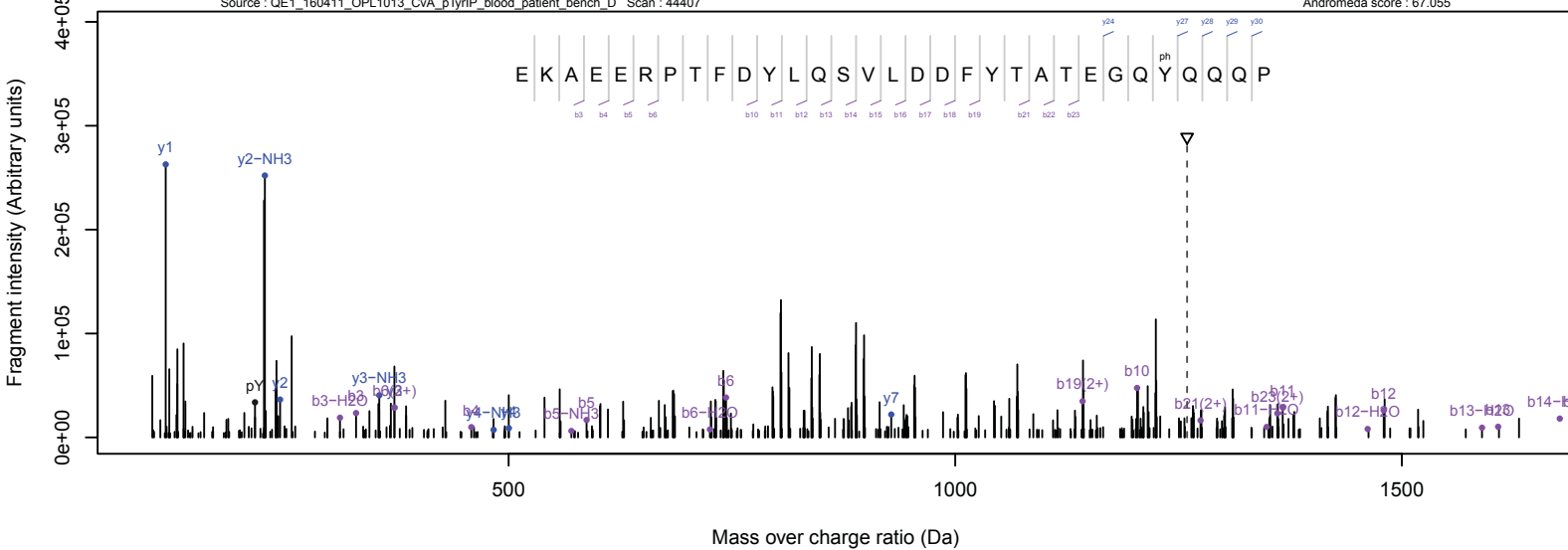
Andromeda score : 67.334



**EKAEERPTFDYLSVLDLDFYATGQY(ph)QQQP-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 44407

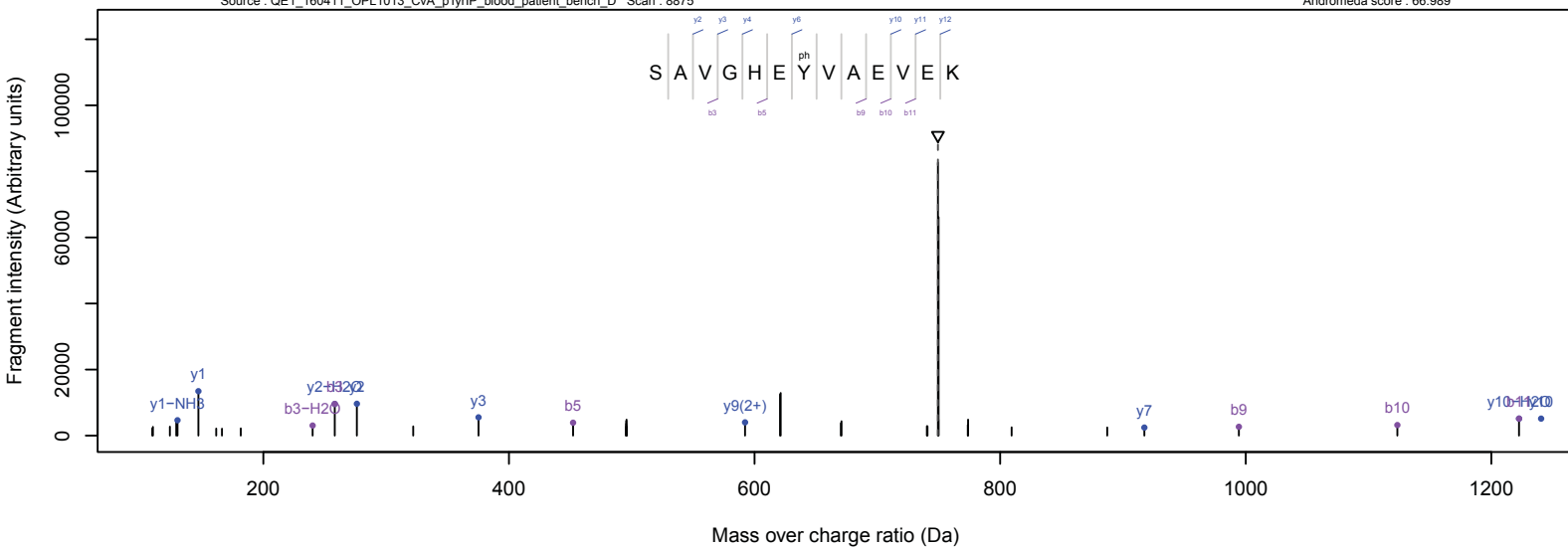
Andromeda score : 67.055



SAVGHEY(ph)VAEVEK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 8875

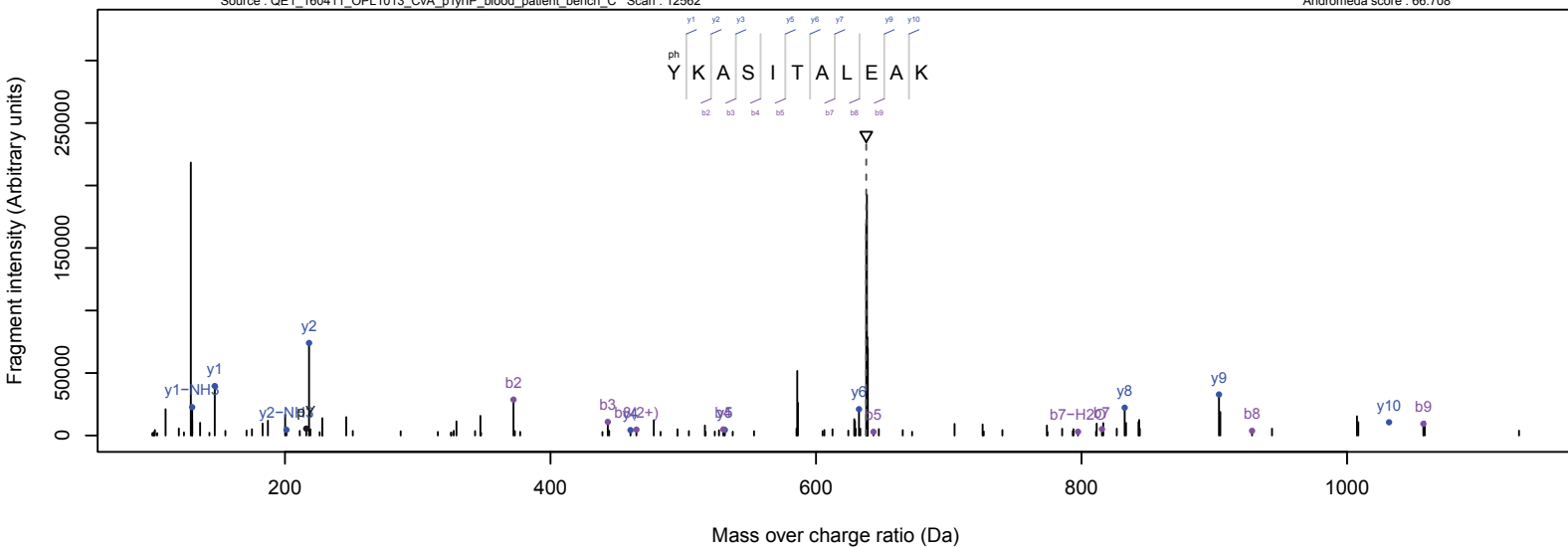
Andromeda score : 66.989



Y(ph)KASITALEAK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 12562

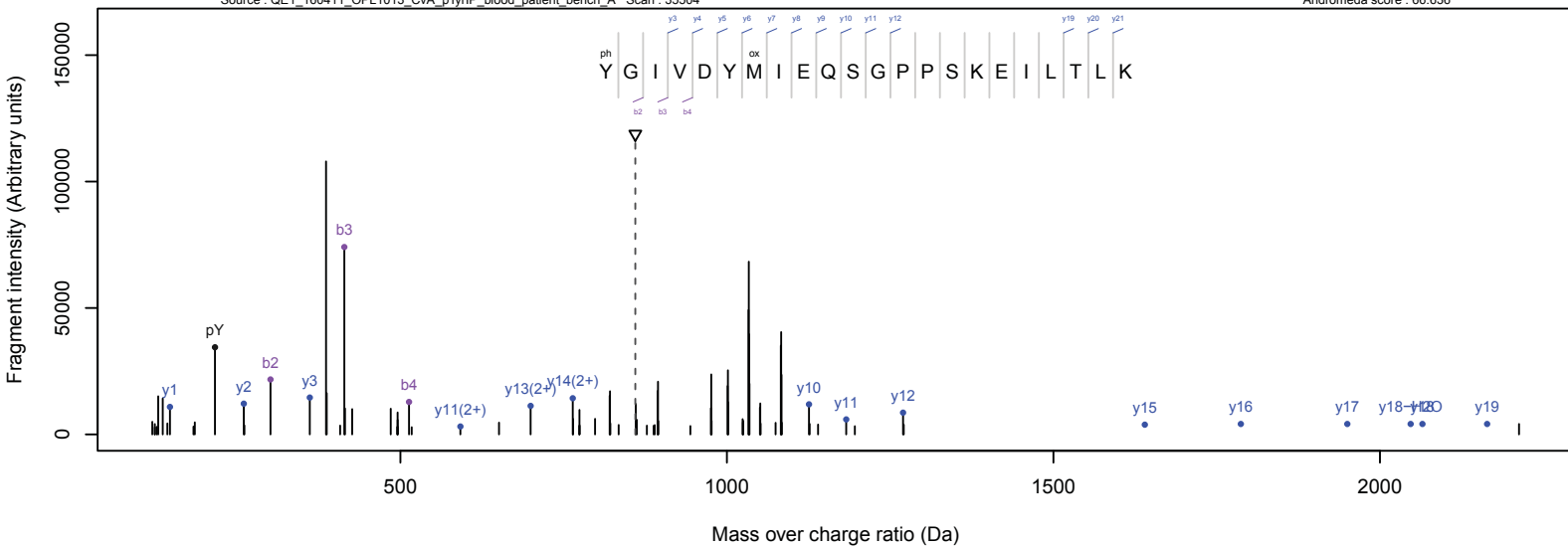
Andromeda score : 66.708



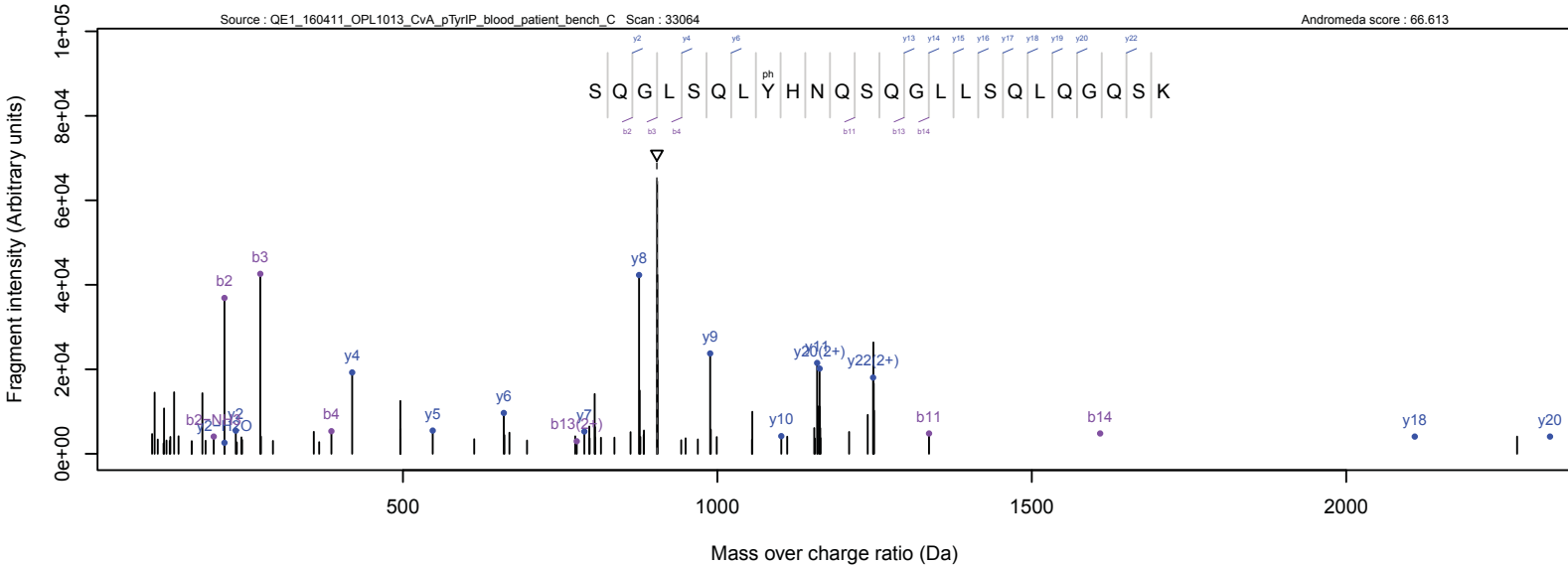
Y(ph)GIVDYM(ox)IEQSGPPSKEILTK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 35504

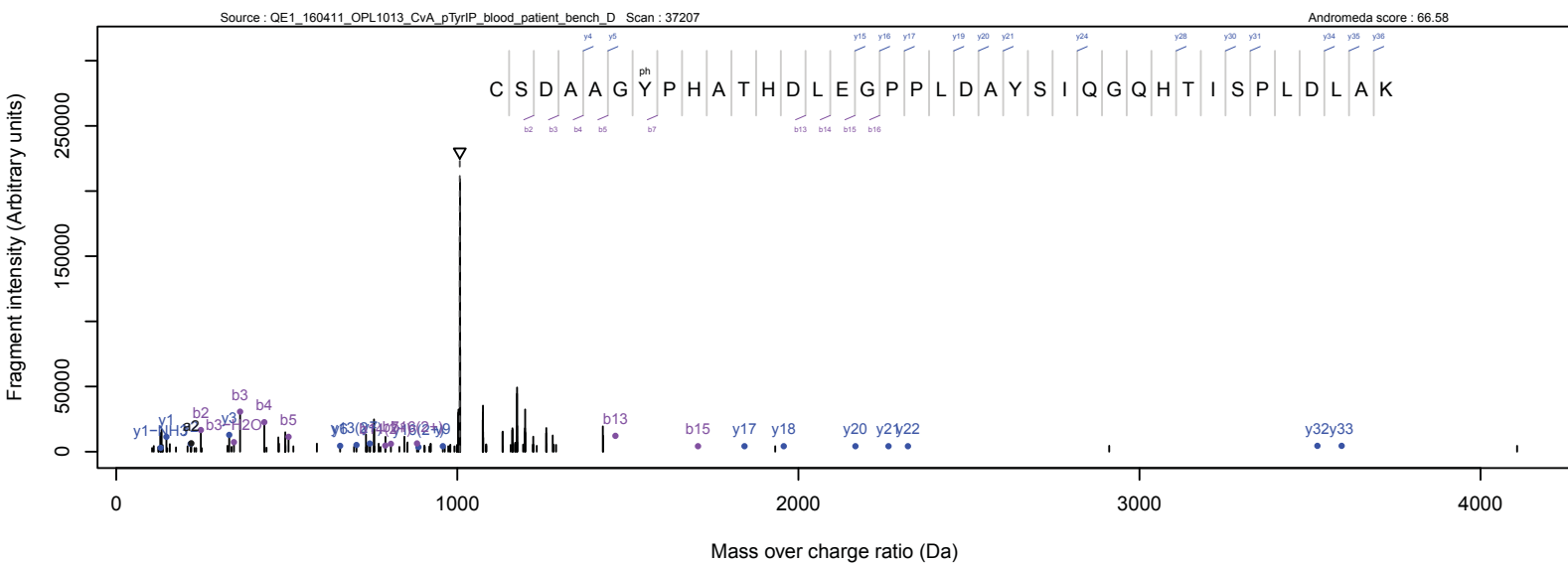
Andromeda score : 66.636



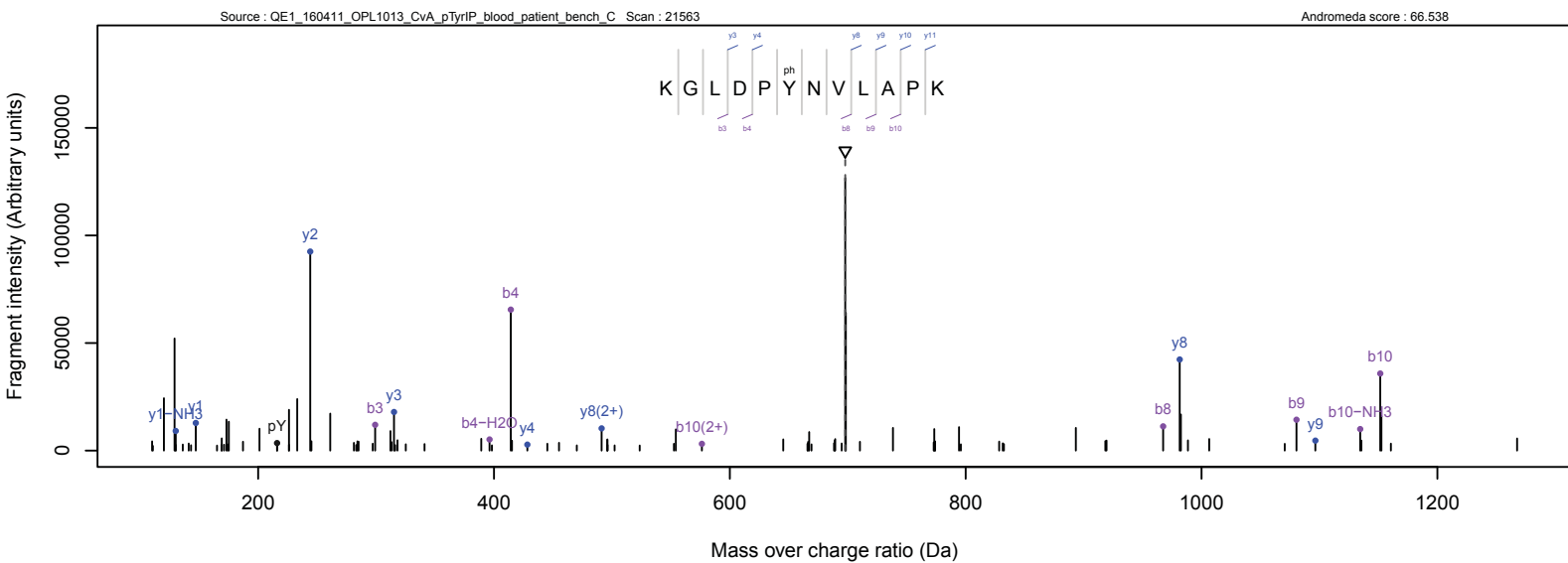
**SQGLS<sup>(ph)</sup>QLYHNQSQGLLSQLQGQSK-3+**



**CSDAAGY<sup>(ph)</sup>PHATHDLEGPPLDAYSIQGQHTISPLDLAK-4+**



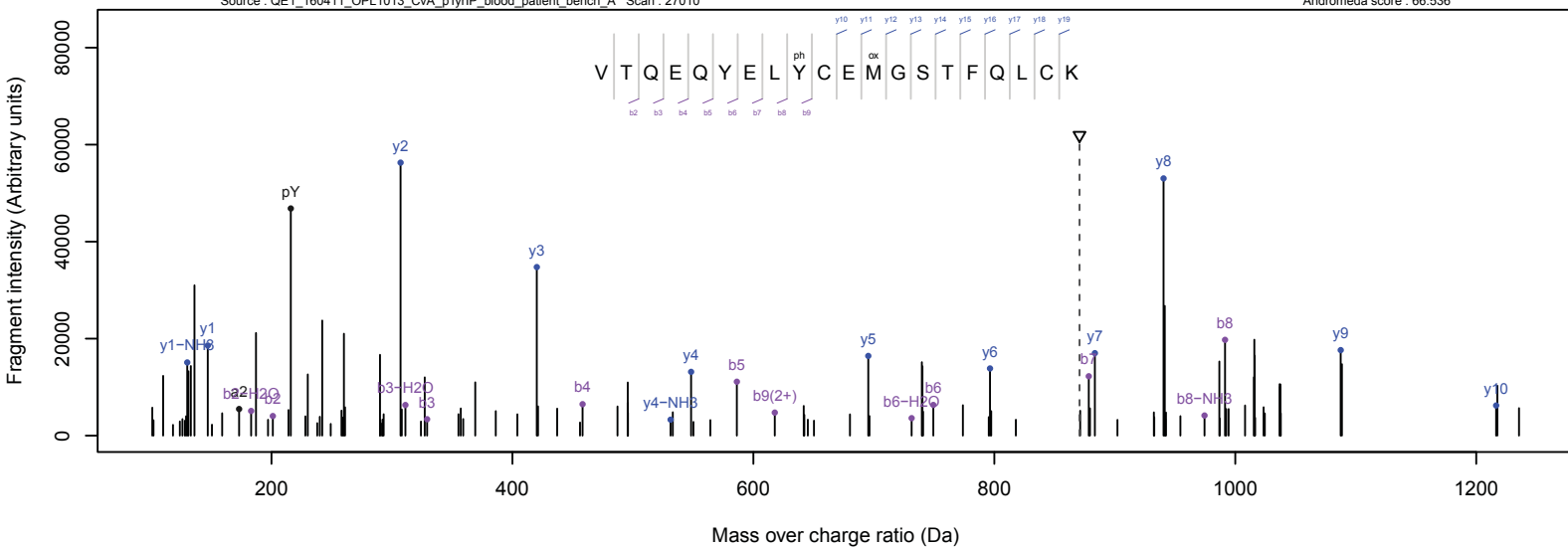
**KGLDPY<sup>(ph)</sup>NVLAPK-2+**



## VTQEYELY(ph)CEM(ox)GSTFLCK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 27010

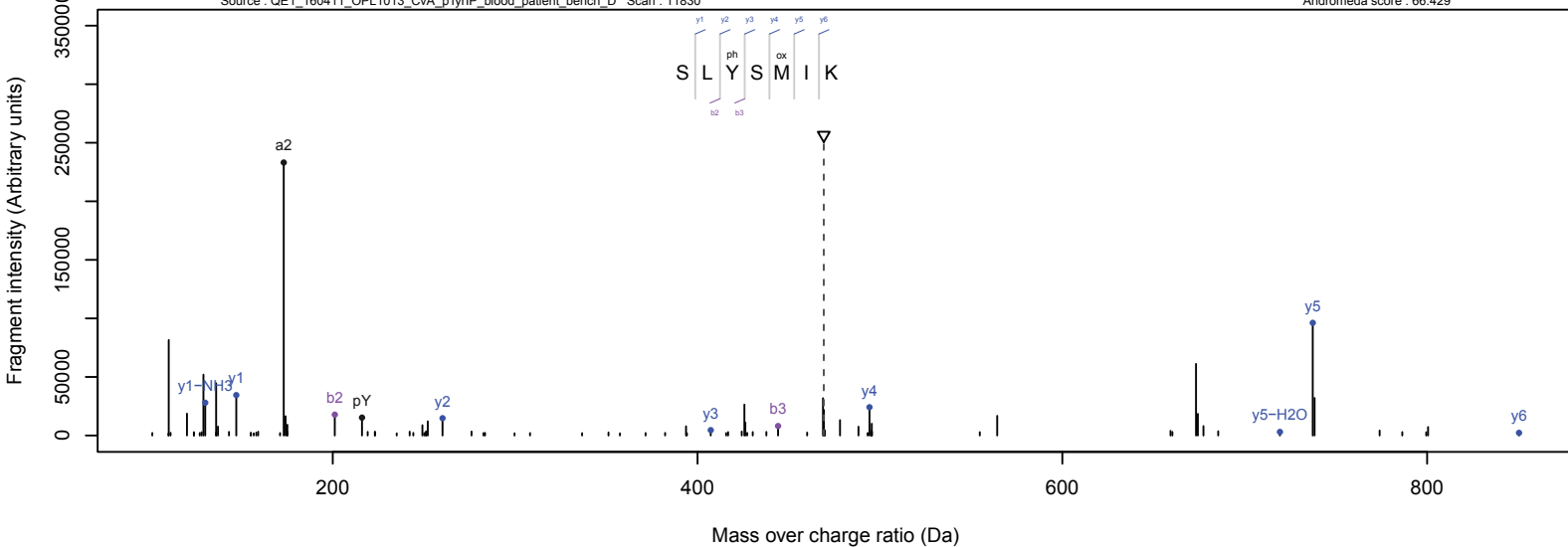
Andromeda score: 66.536



## SLY(ph)SM(ox)IK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 11830

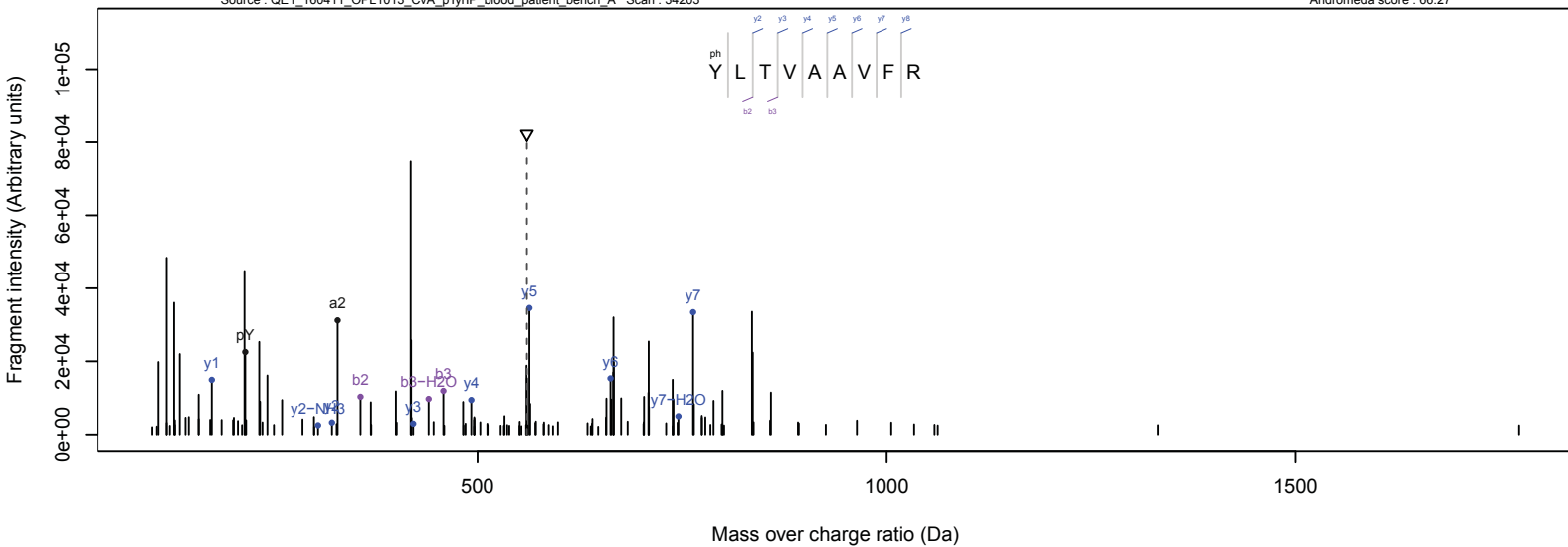
Andromeda score: 66.429



## Y(ph)LTVAAVFR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 34203

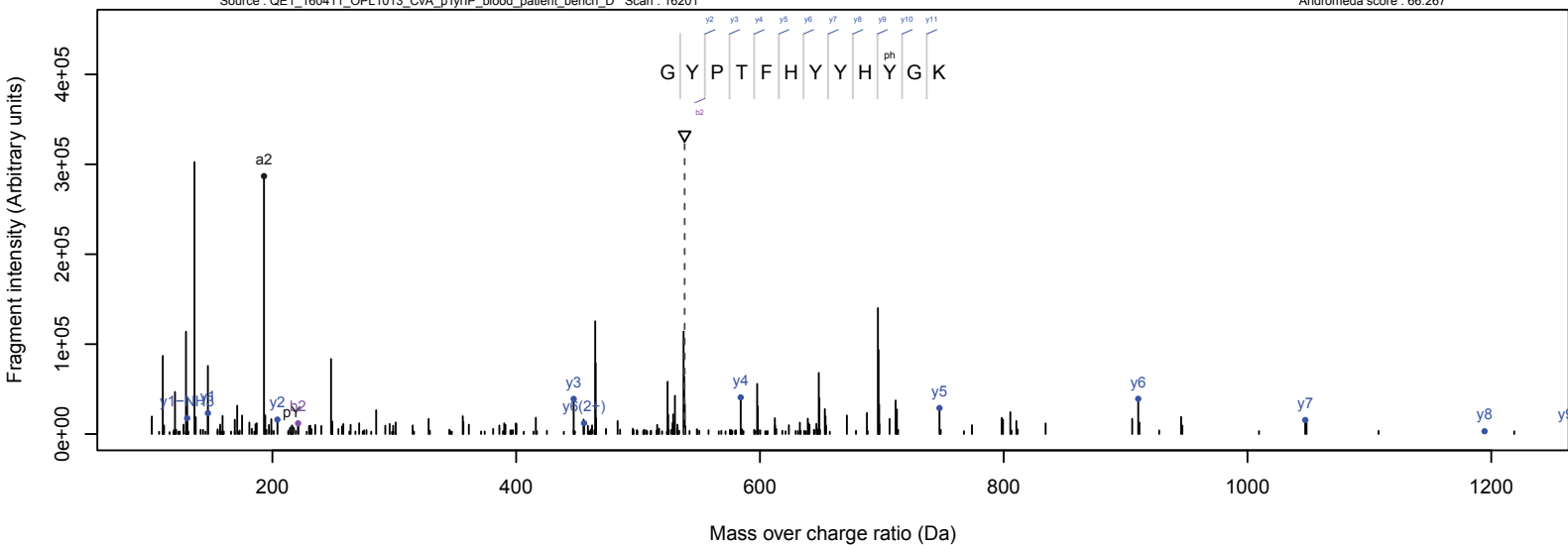
Andromeda score: 66.27



## GYPTFHYHY(ph)GK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 16201

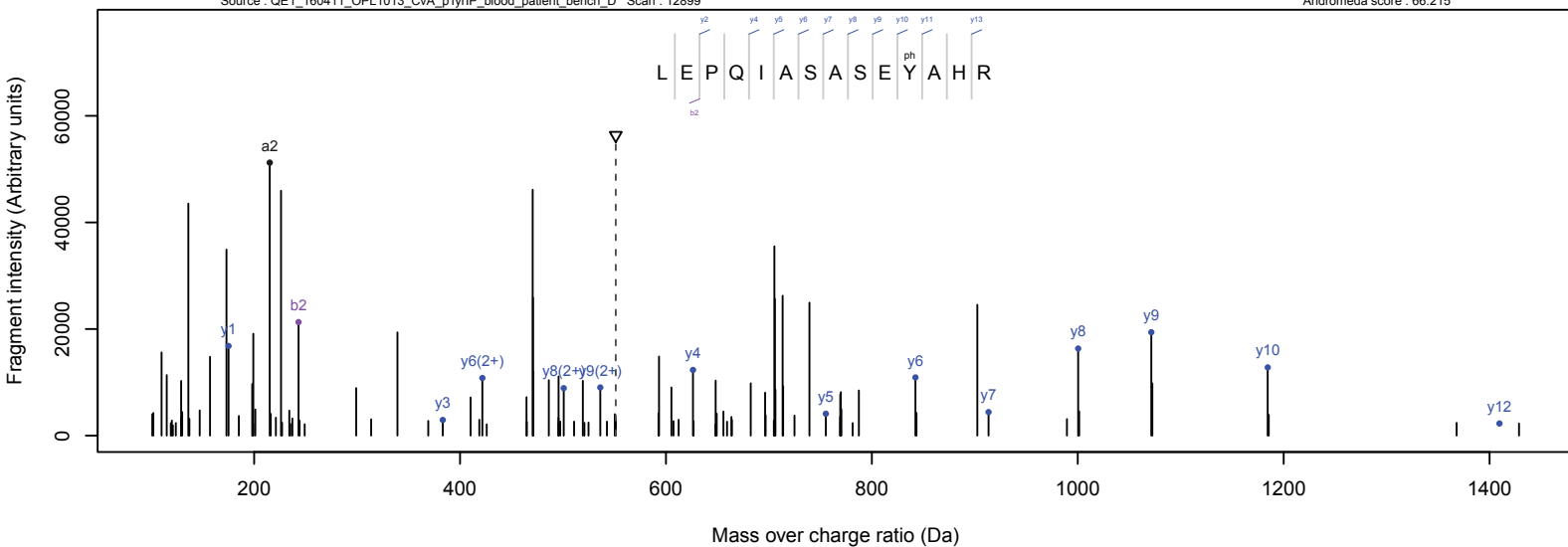
Andromeda score: 66.267



LEPQIASASEY(ph)AHR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 12899

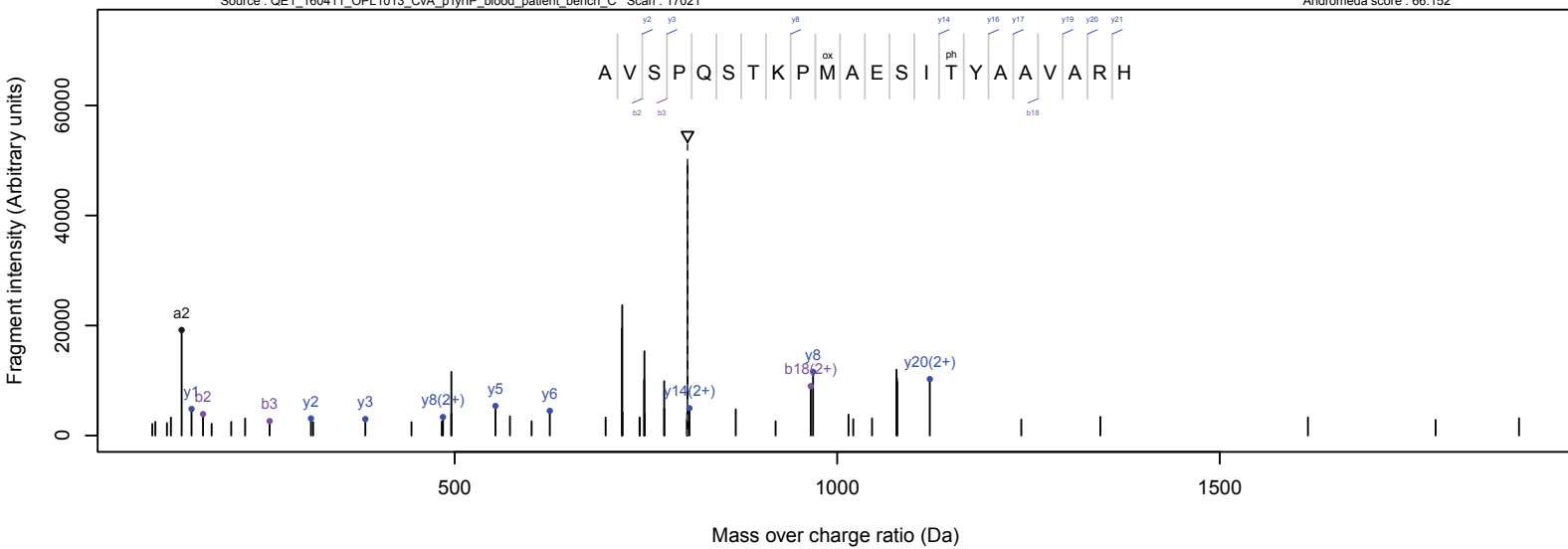
Andromeda score: 66.215



AVSPQSTKPM(ox)AESIT(ph)YAAVARH-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 17021

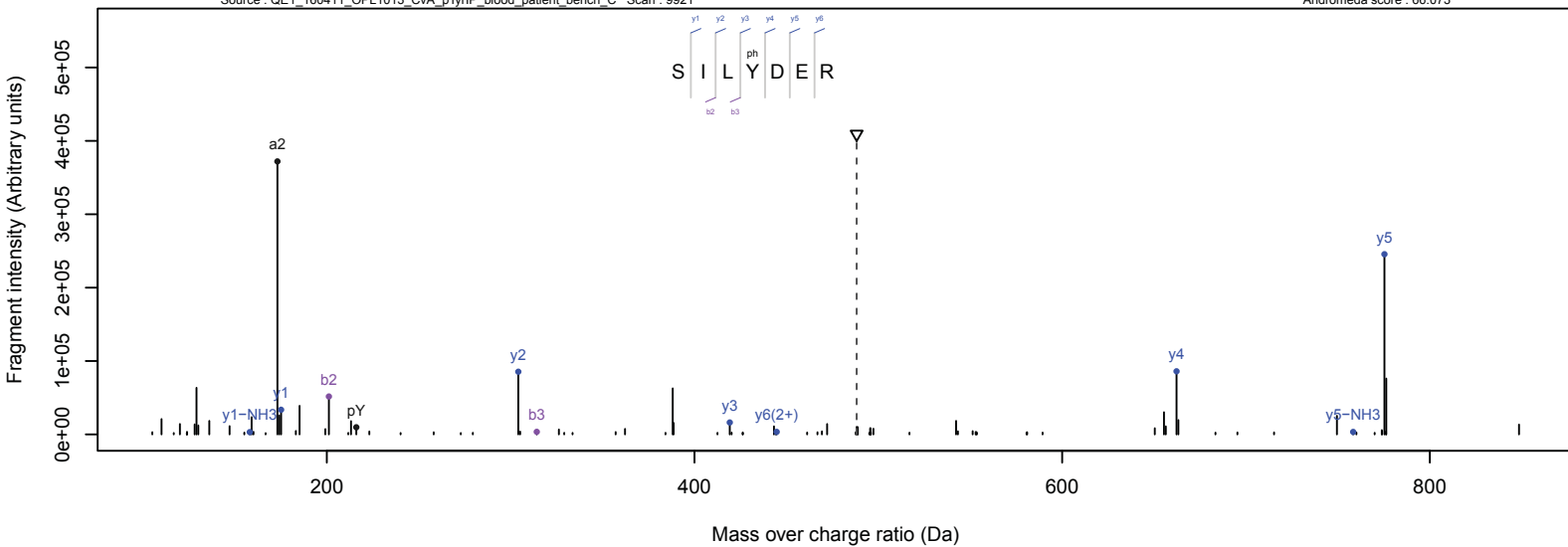
Andromeda score: 66.152



SILY(ph)DER-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 9921

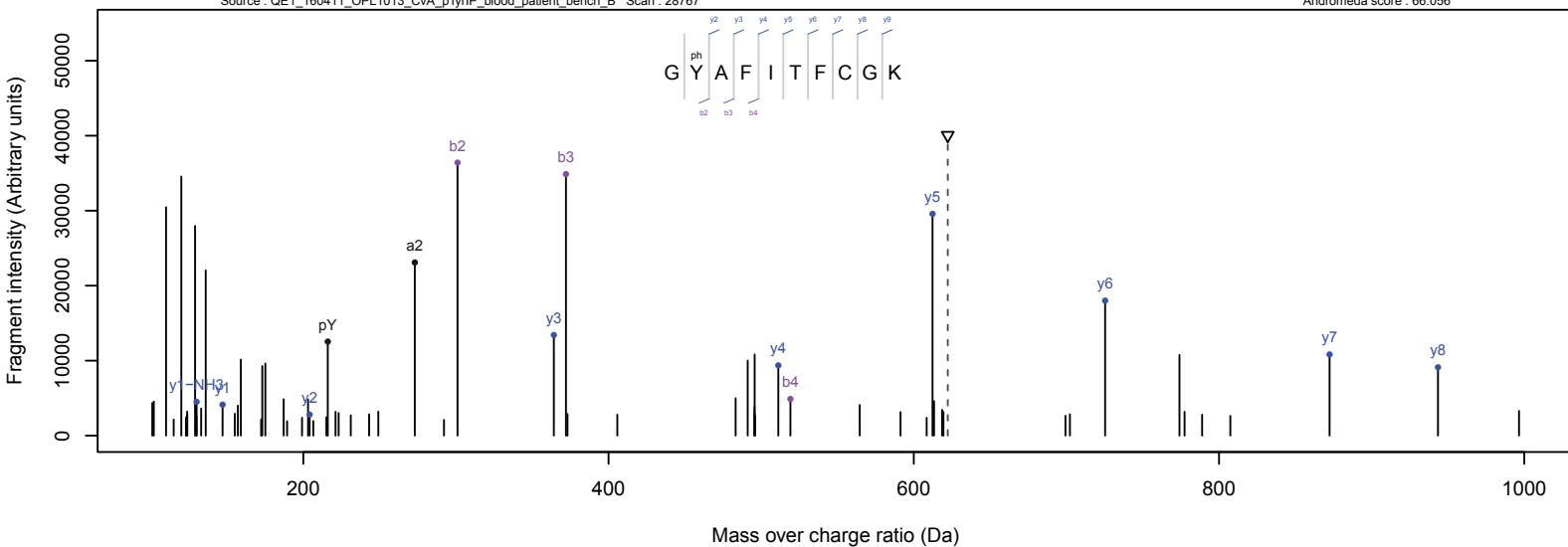
Andromeda score: 66.073



## GY(ph)AFITFCGK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 28767

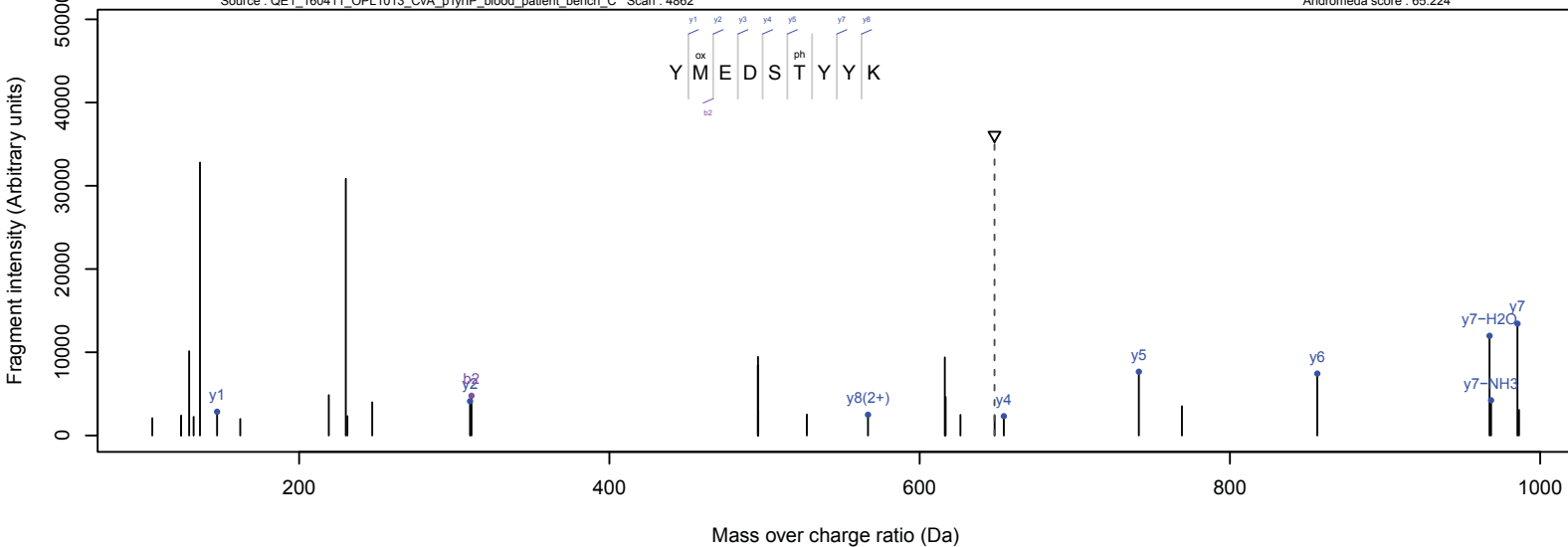
Andromeda score : 66.056



## YM(ox)EDST(ph)YYK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 4862

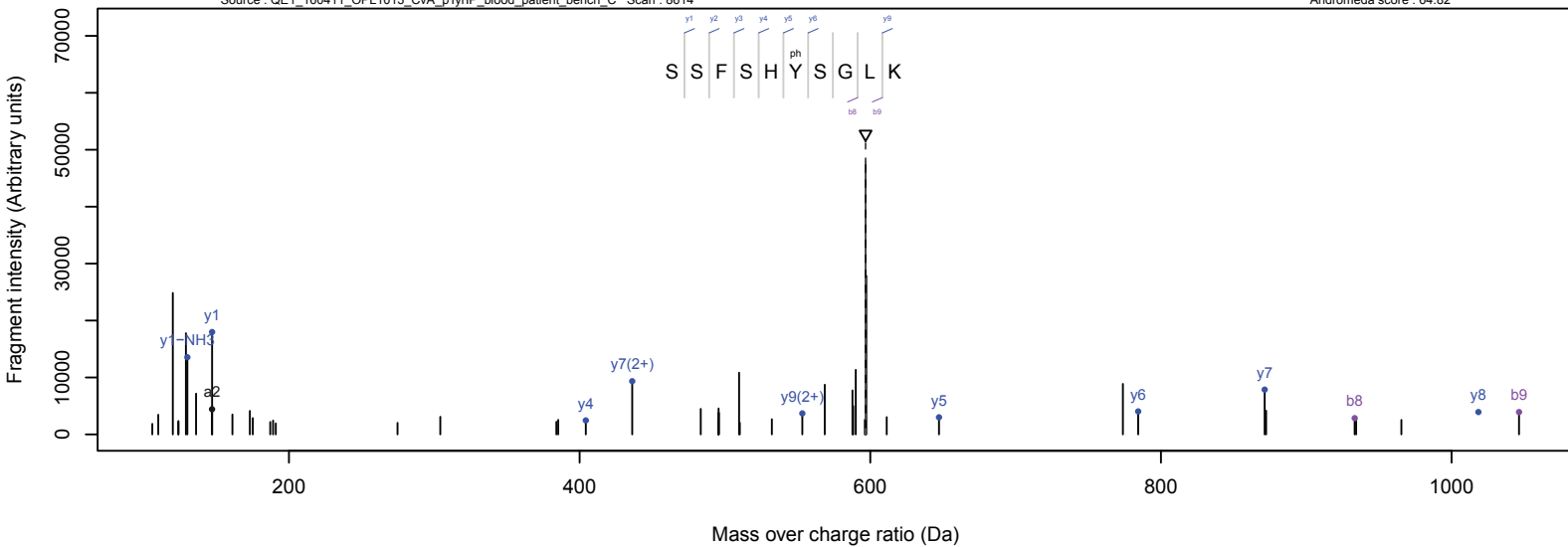
Andromeda score : 65.224



## SSFSHY(ph)SGLK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 8614

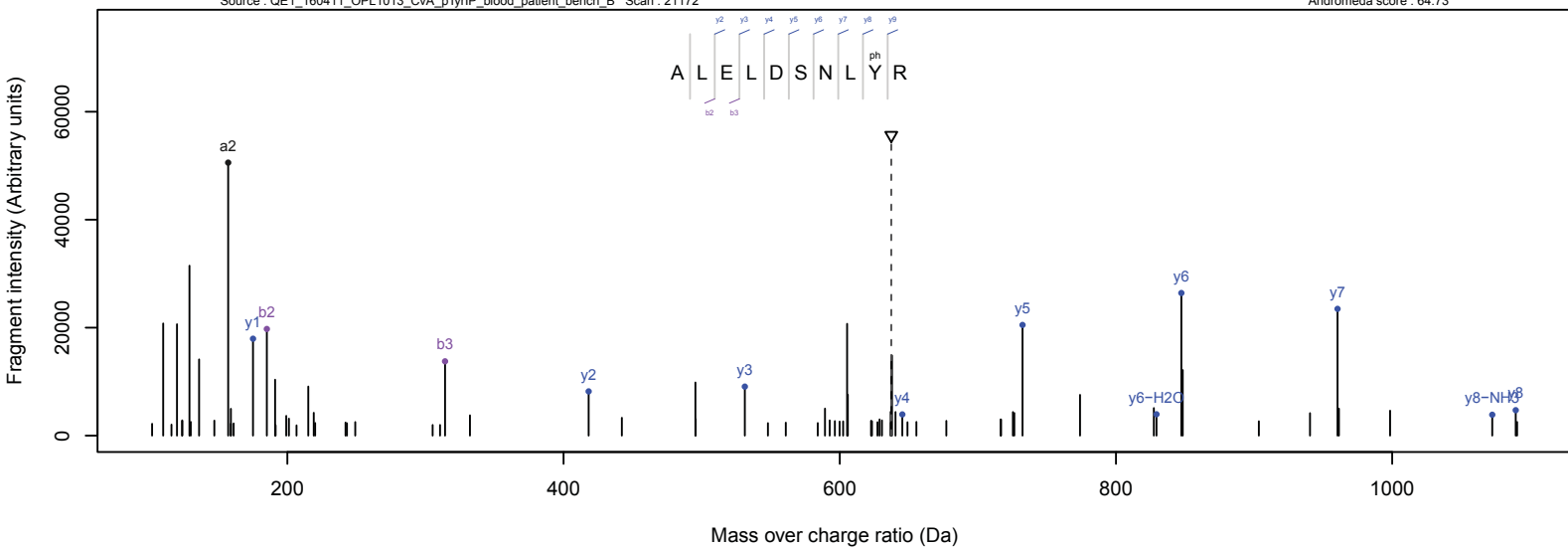
Andromeda score : 64.82



**ALELDSNLY(ph)R-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 21172

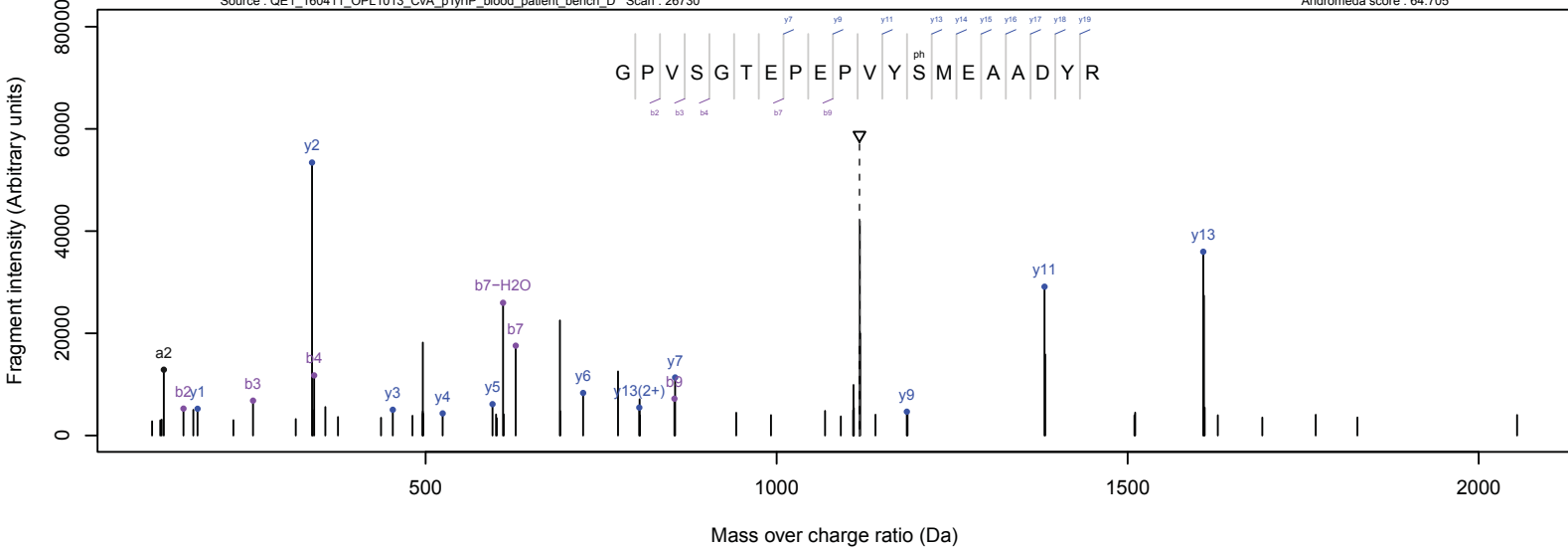
Andromeda score: 64.73



**GPVSGTEPEP VY S MEAADYR-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 26730

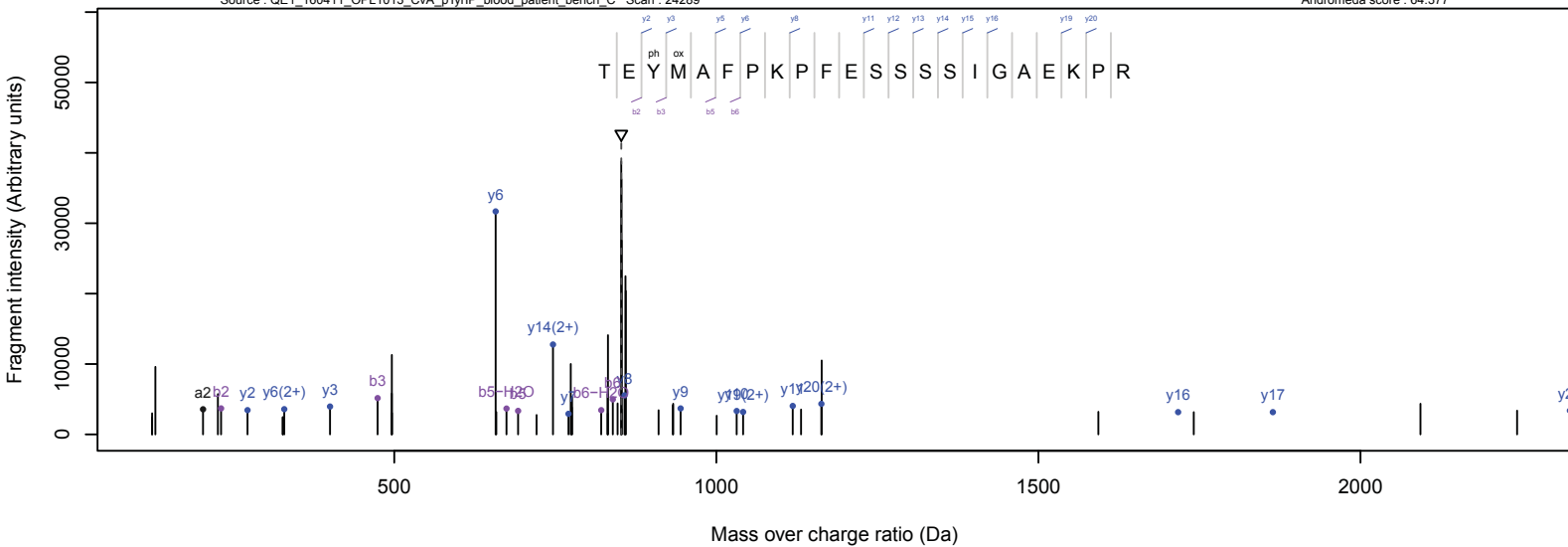
Andromeda score: 64.705



**TEY M(ox)AFP KPFESSSSIGAEKPR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 24289

Andromeda score: 64.377

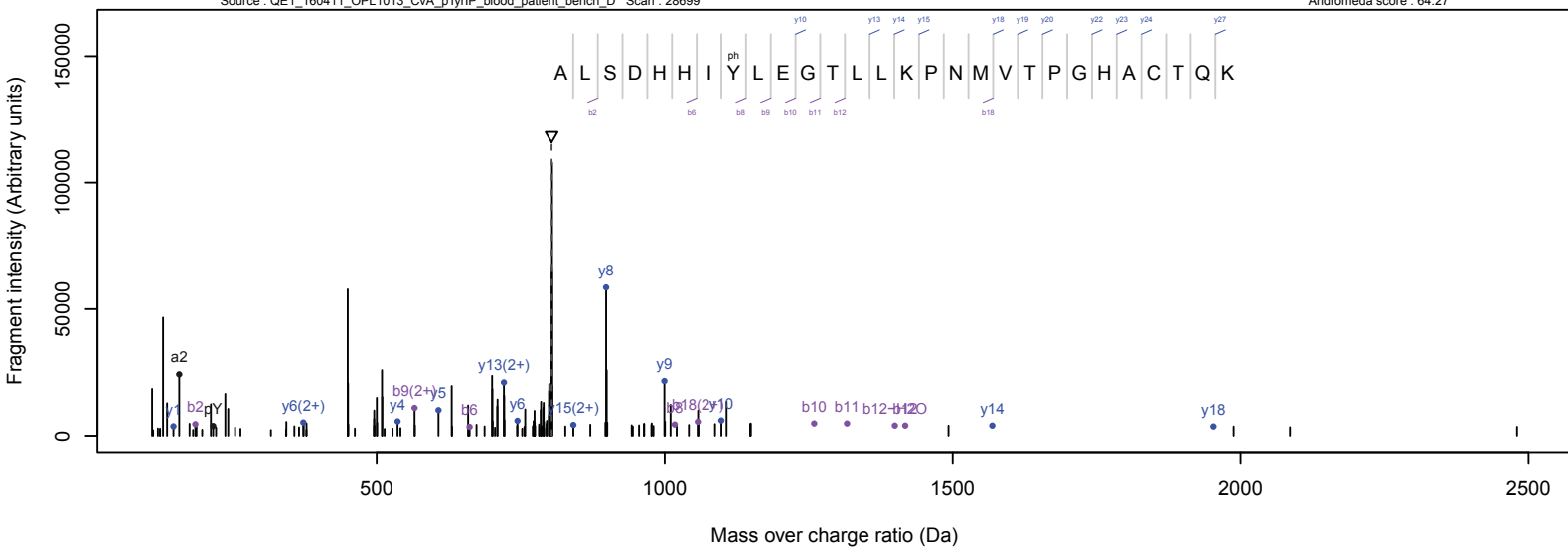




**ALSDHHIY(ph)LEGTL LKPNMVT P GHACTQK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28699

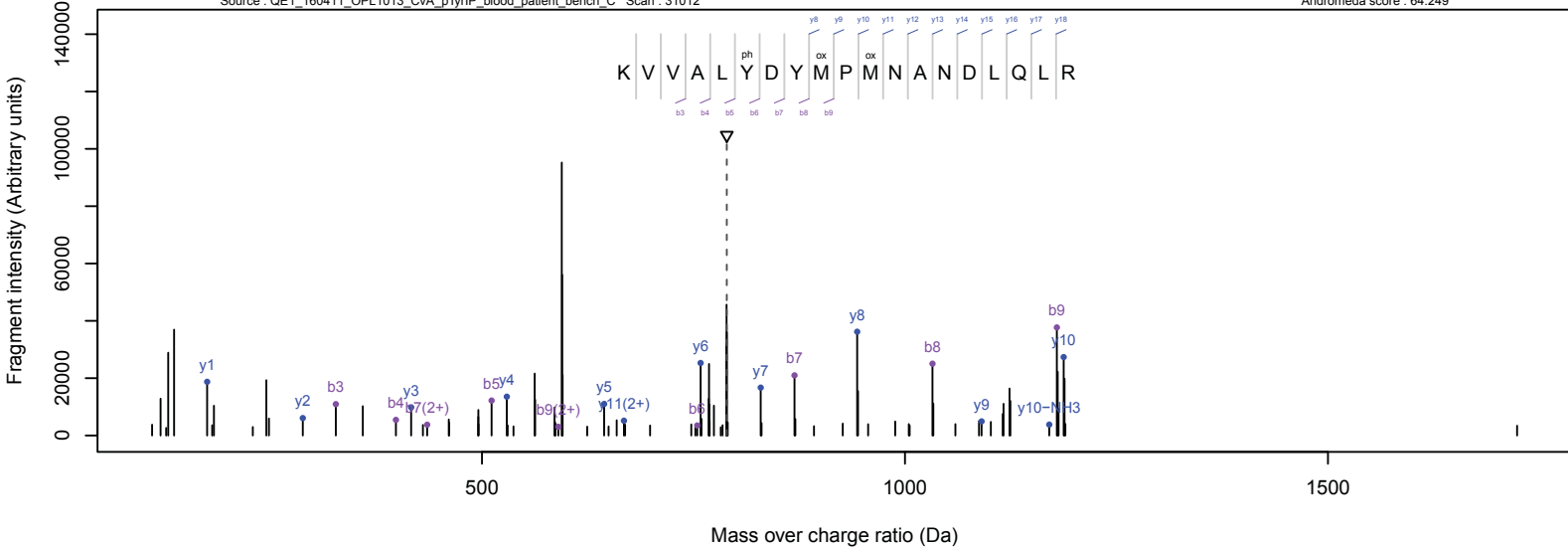
Andromeda score : 64.27



**KVVALY(ph)DYM(ox)PM(ox)NANDLQLR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 31012

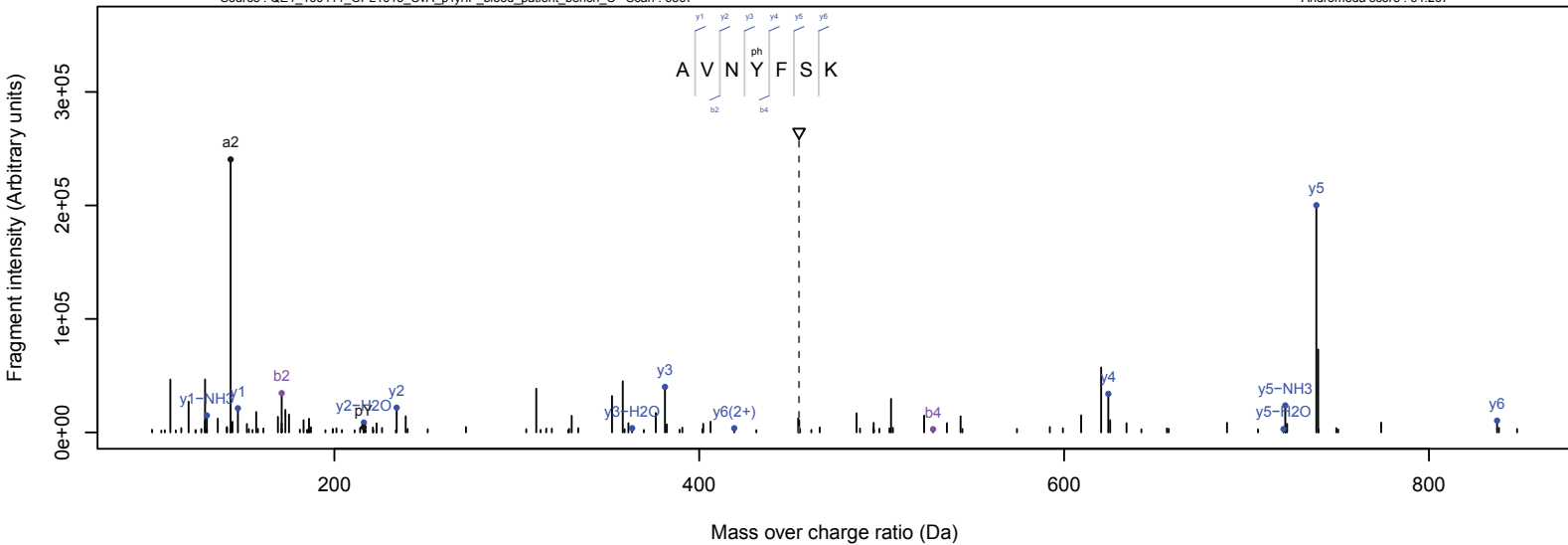
Andromeda score : 64.249



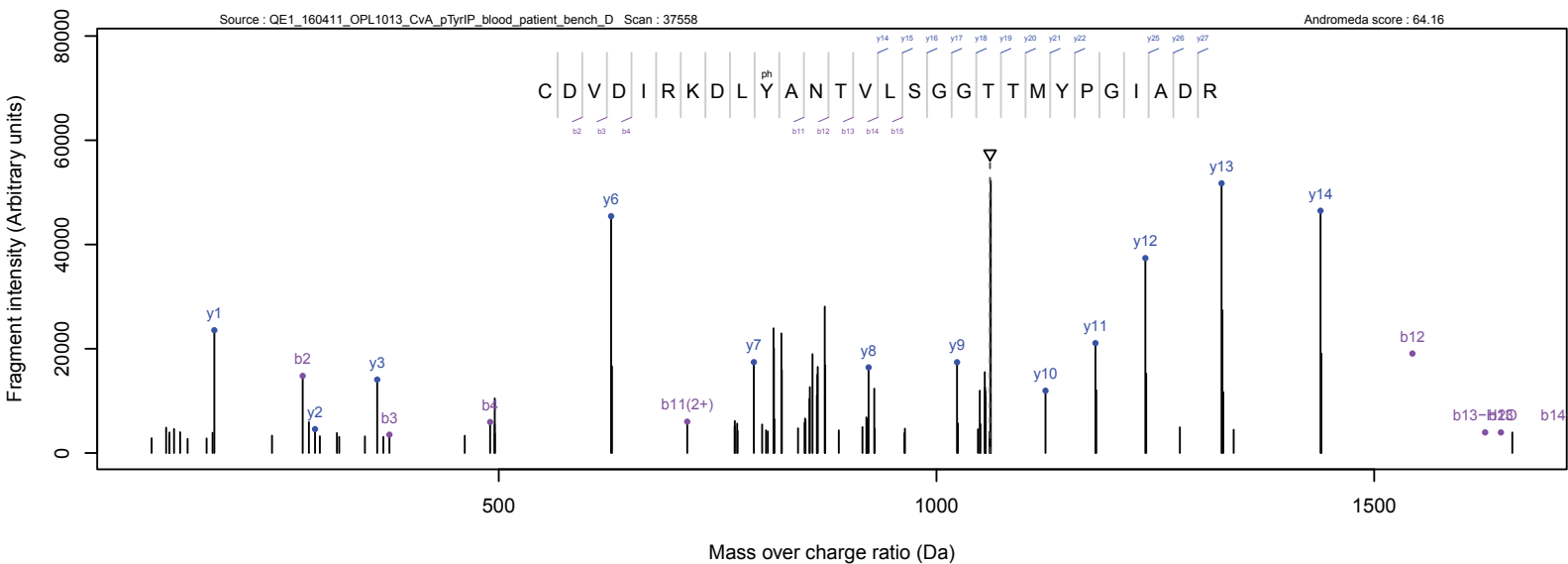
**AVNY(ph)FSK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 6567

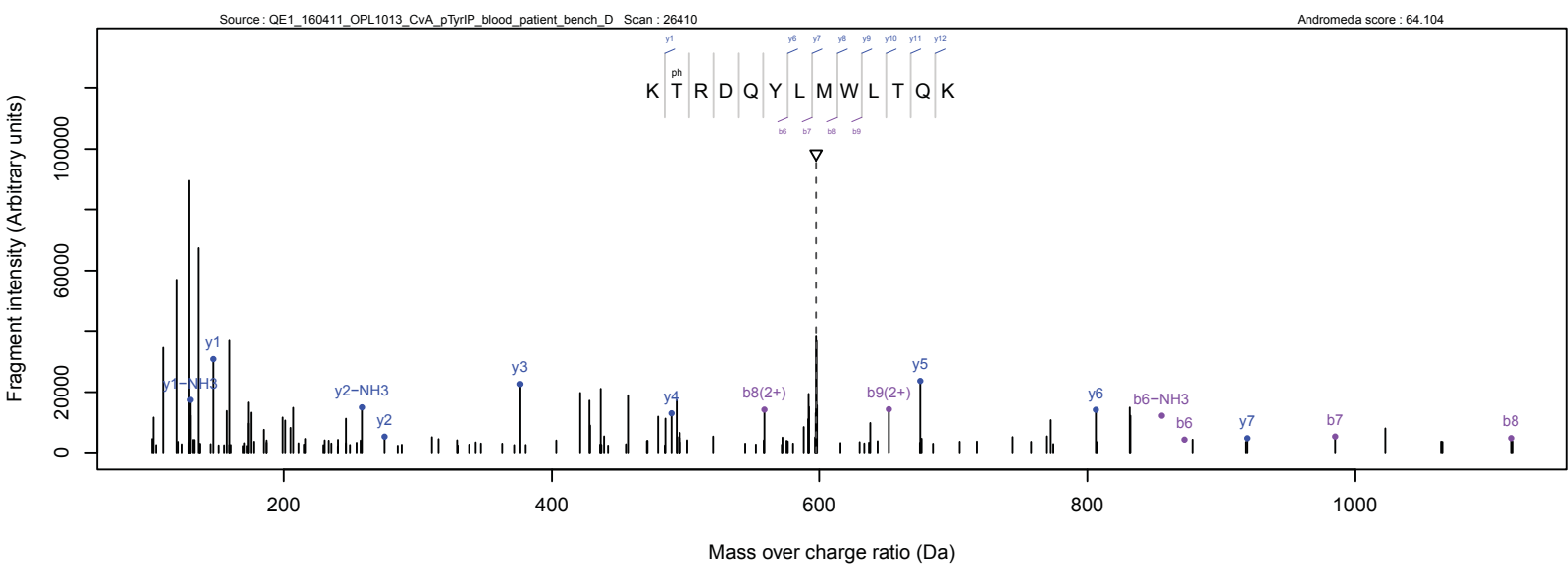
Andromeda score : 64.207



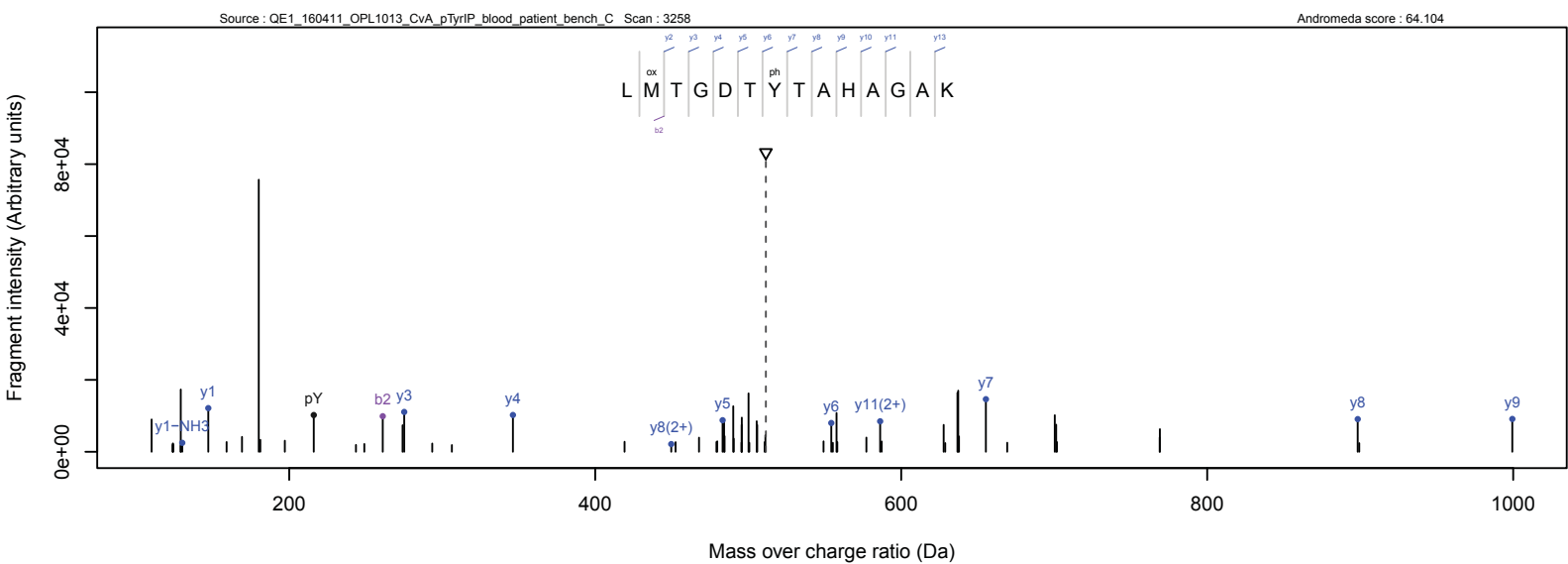
## CDVDIRKDL(ph)ANTVLSGGTTMYPGIADR-3+



## KT(ph)RDQYLMWLTQK-3+



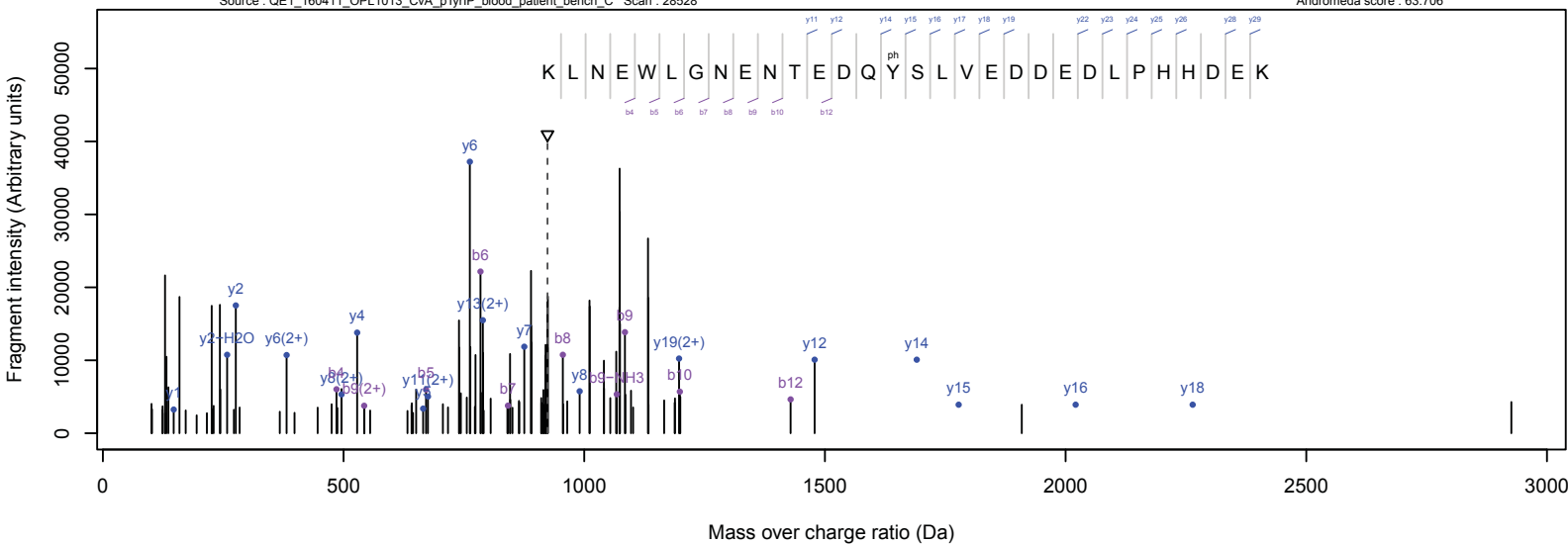
## LM(ox)TGDTY(ph)TAHAGAK-3+



**KLNEWLGNENTEDQ(ph)SLVEDDEDLPHHDEK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28528

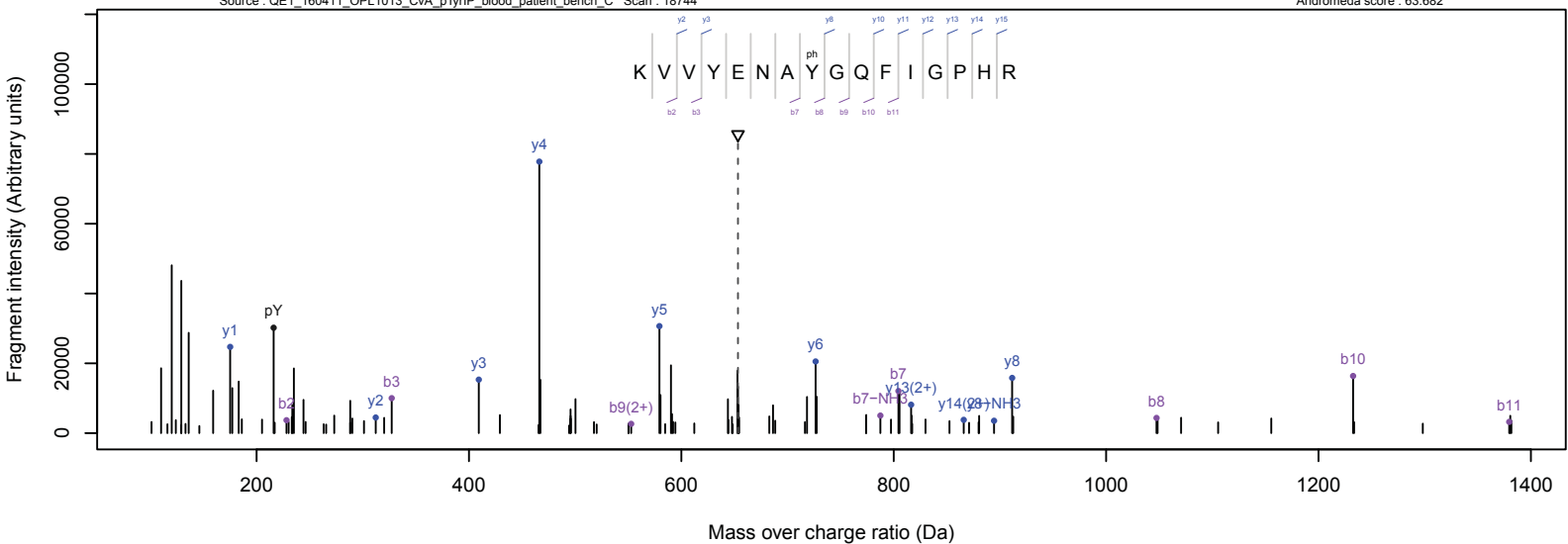
Andromeda score : 63.706



**KVYENAY(ph)GQFIGPHR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18744

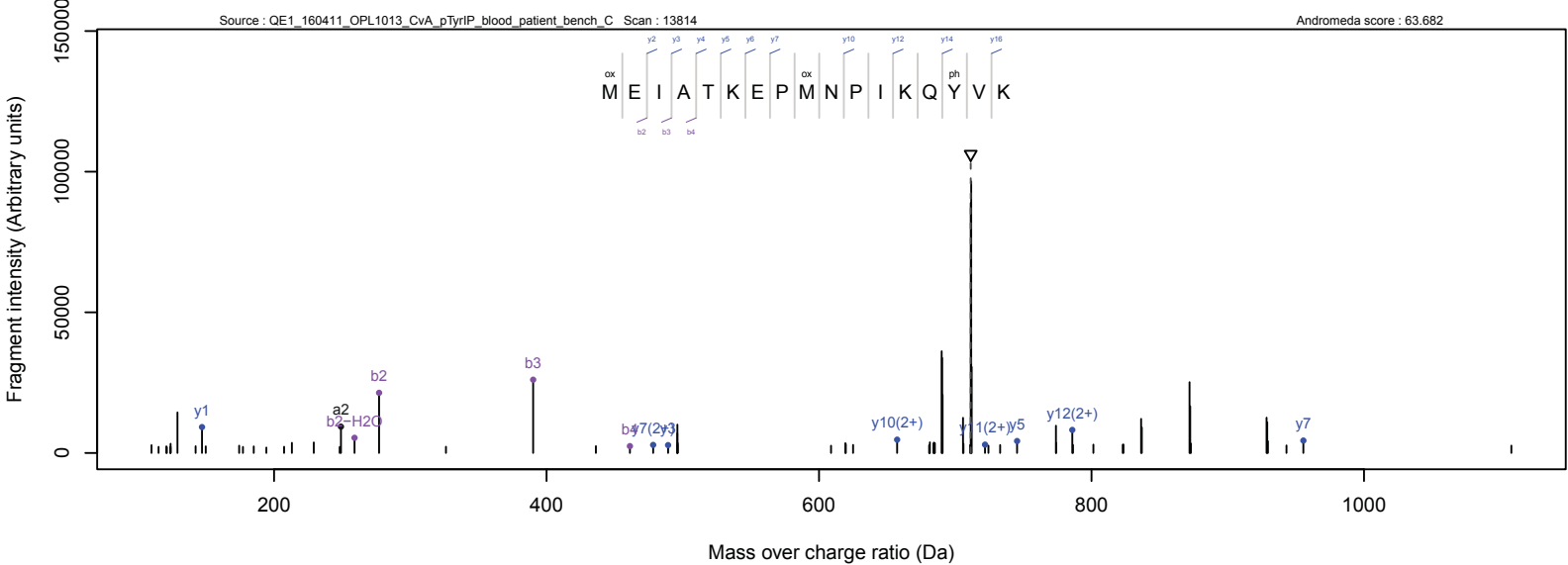
Andromeda score : 63.682



**M(ox)EIAATKEP(ox)NPIKQ(ph)YVK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 13814

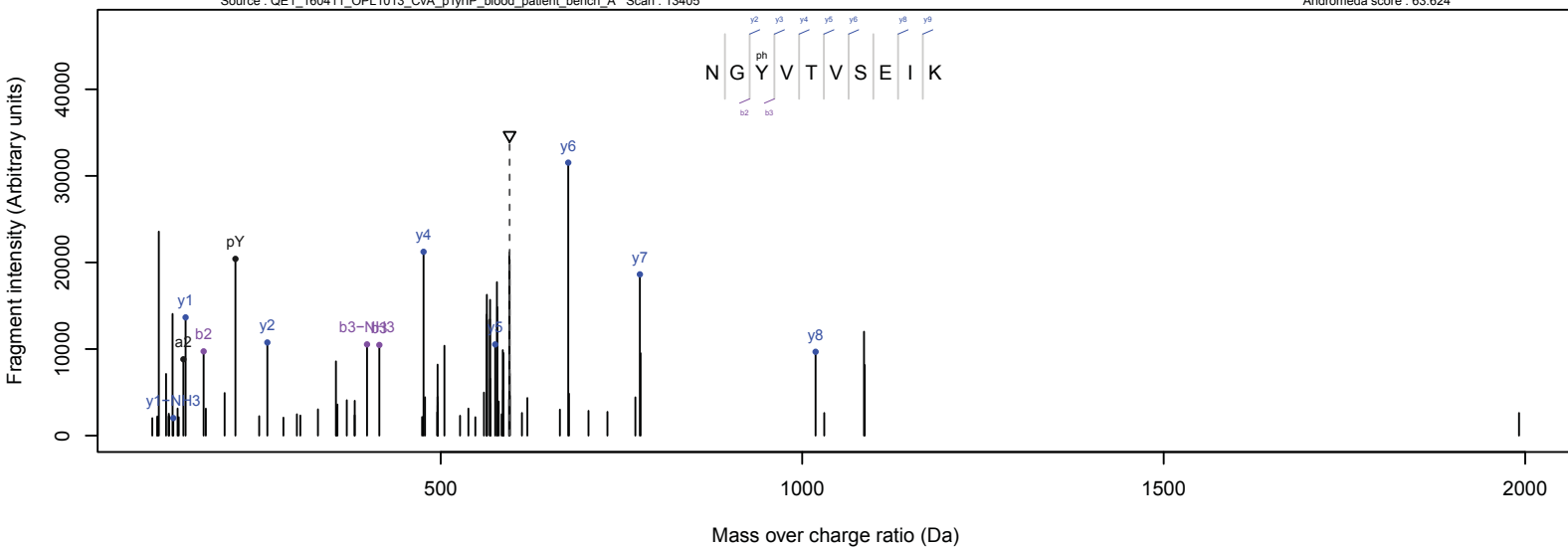
Andromeda score : 63.682



## NGY(ph)VTVEIK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 13405

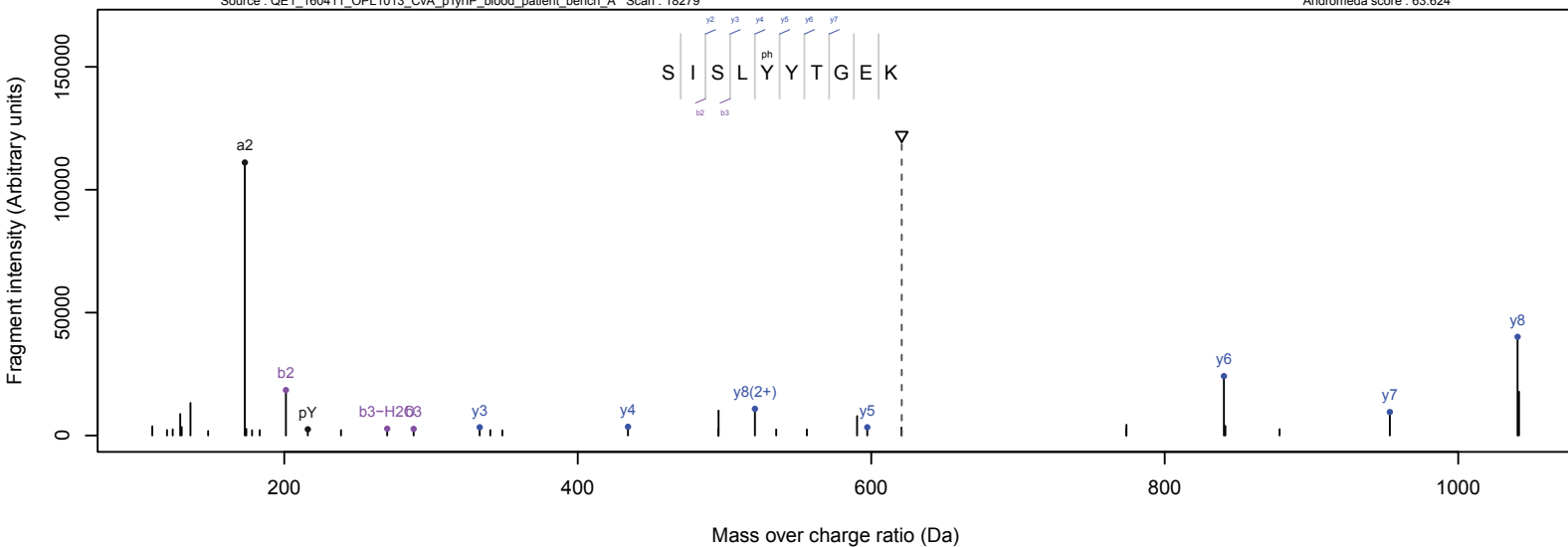
Andromeda score : 63.624



## SISLY(ph)YTGEK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 18279

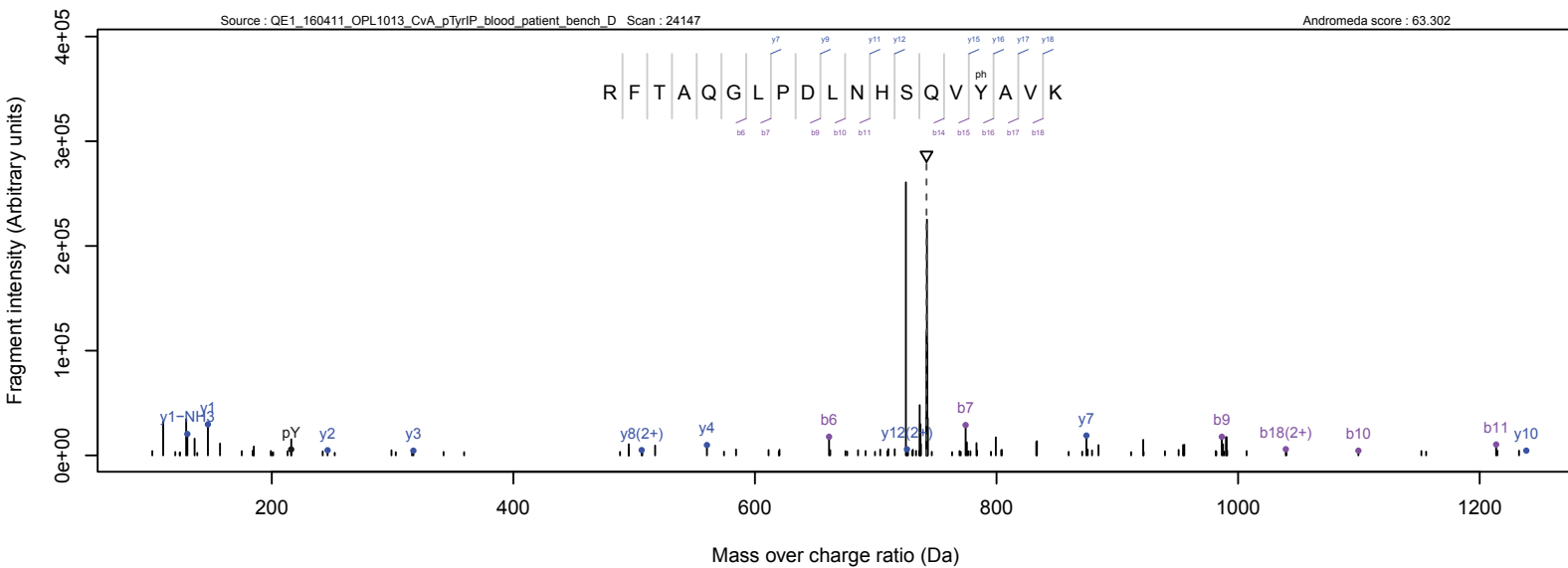
Andromeda score : 63.624



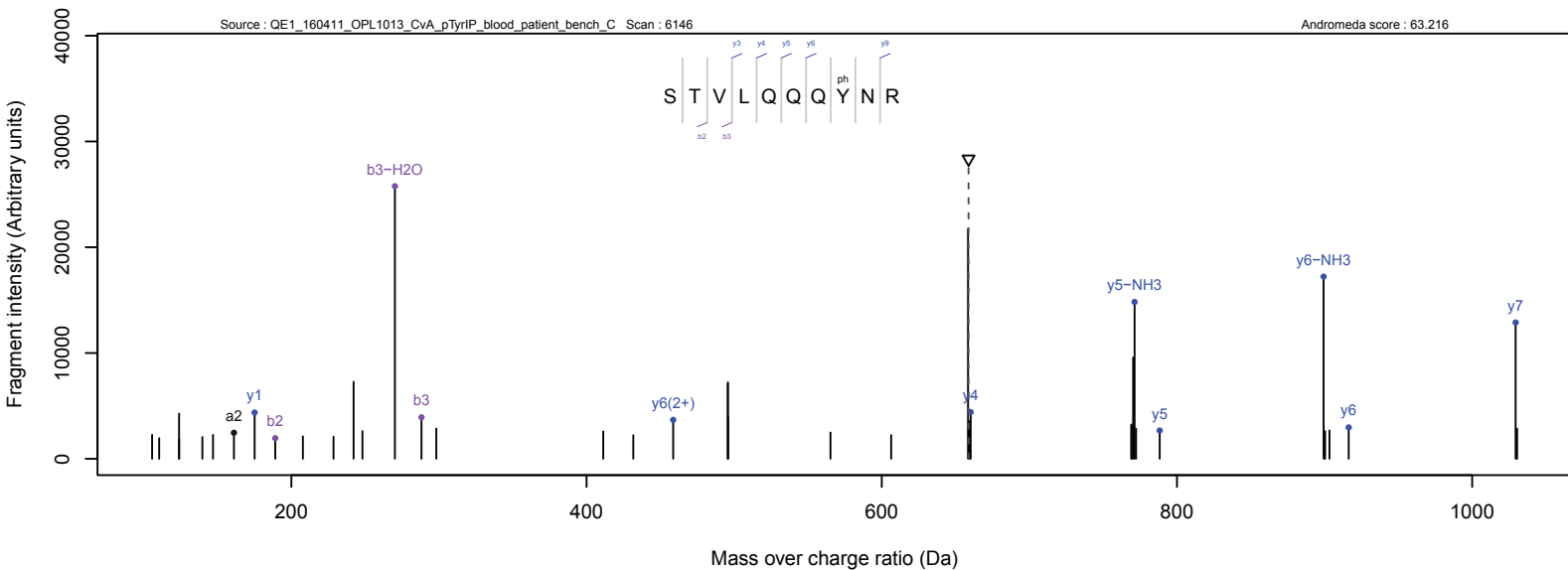
## RFTAQGLPDLNHSQVY(ph)AVK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24147

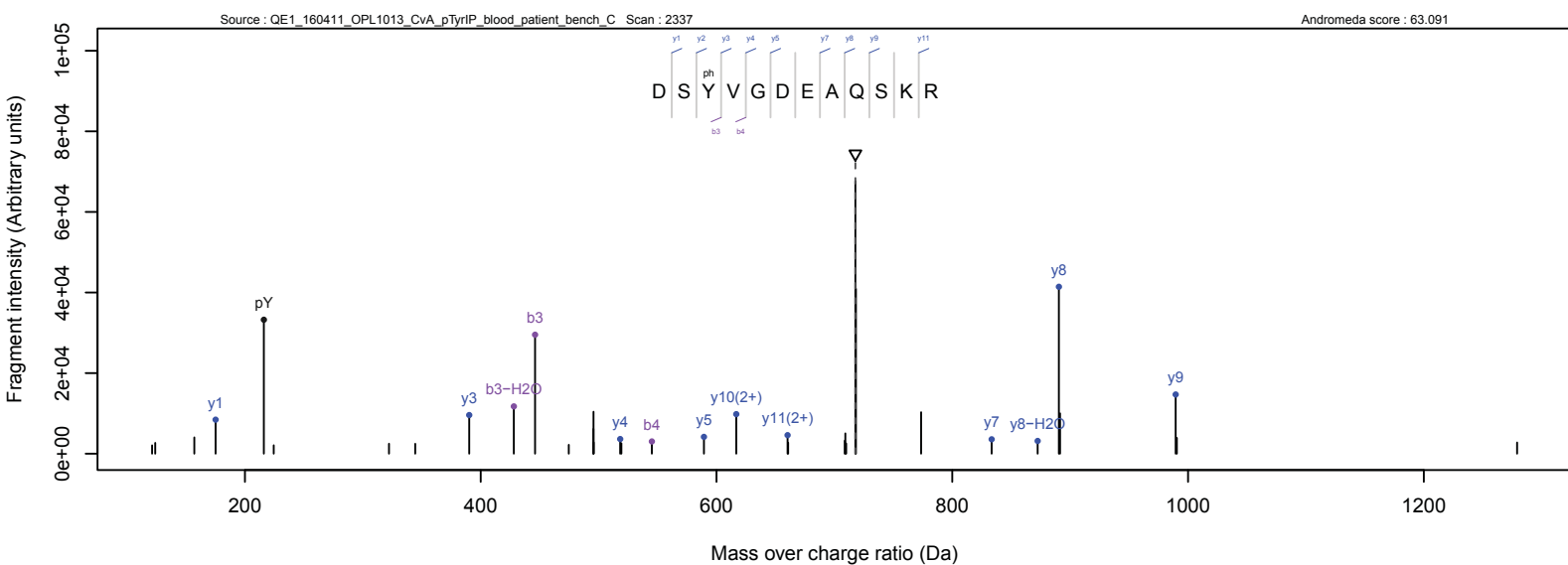
Andromeda score : 63.302



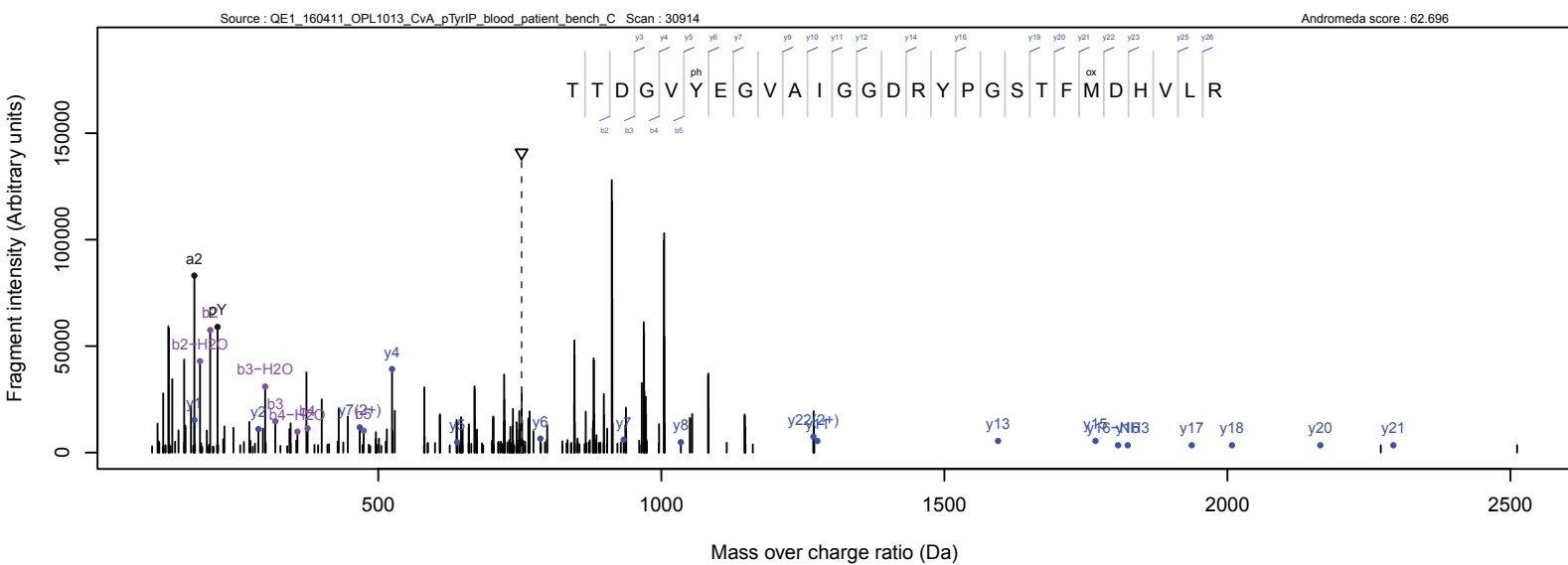
**STVLQQY(ph)NR-2+**



**DSY(ph)VGDEAQSQR-2+**



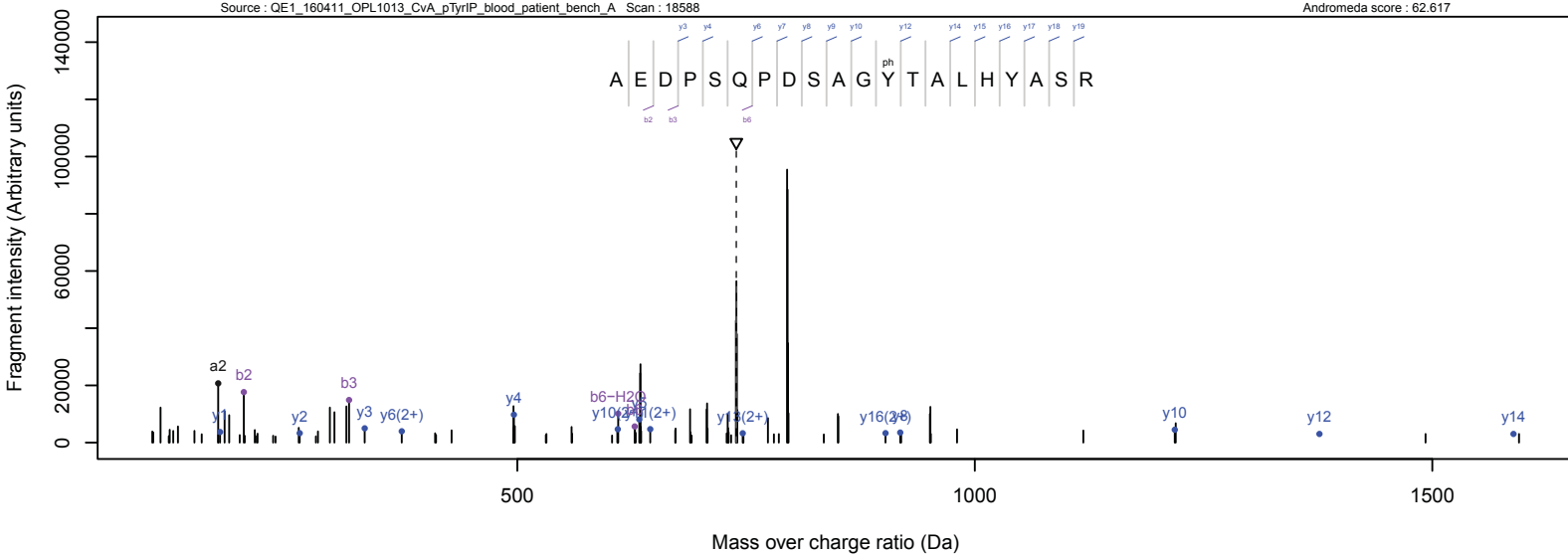
**TTDGVY(ph)EGVAIGDRYPGSTFMDHVL R-4+**



**AEDPSQPSAGY(ph)TALHYASR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 18588

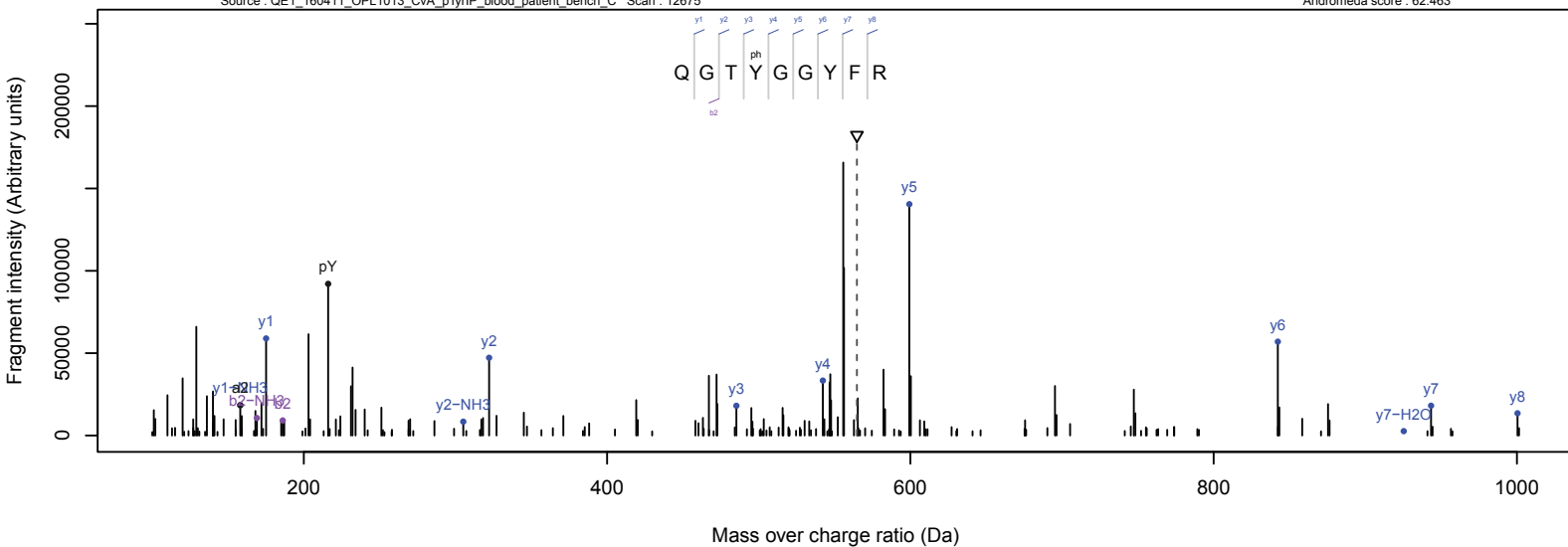
Andromeda score : 62.617



**QGTY(ph)GGYFR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 12675

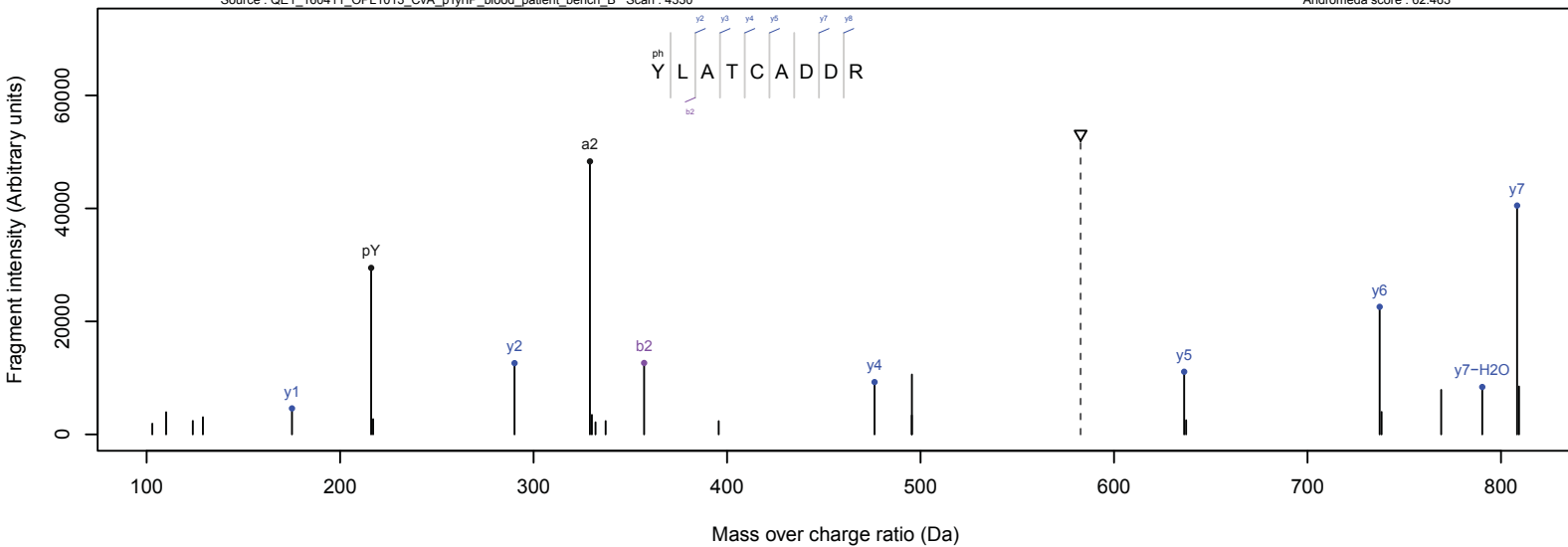
Andromeda score : 62.463



**Y(ph)LATCADDR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 4330

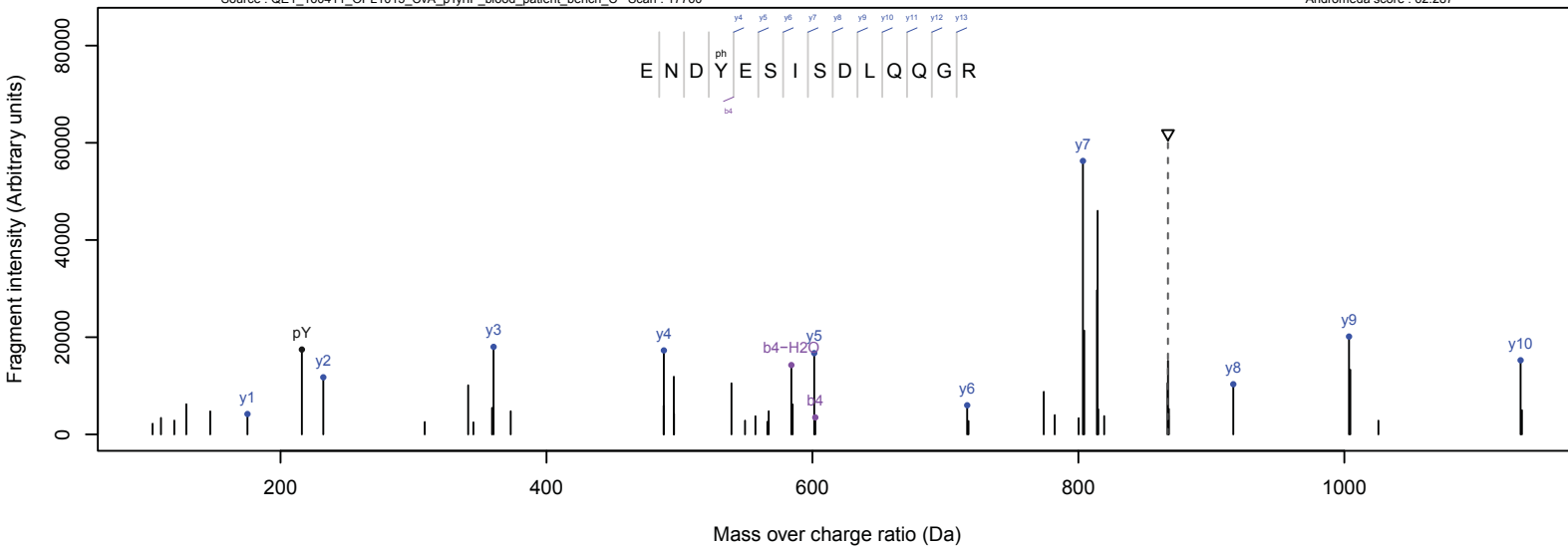
Andromeda score : 62.463



## ENDY(ph)ESISDLQQGR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 17760

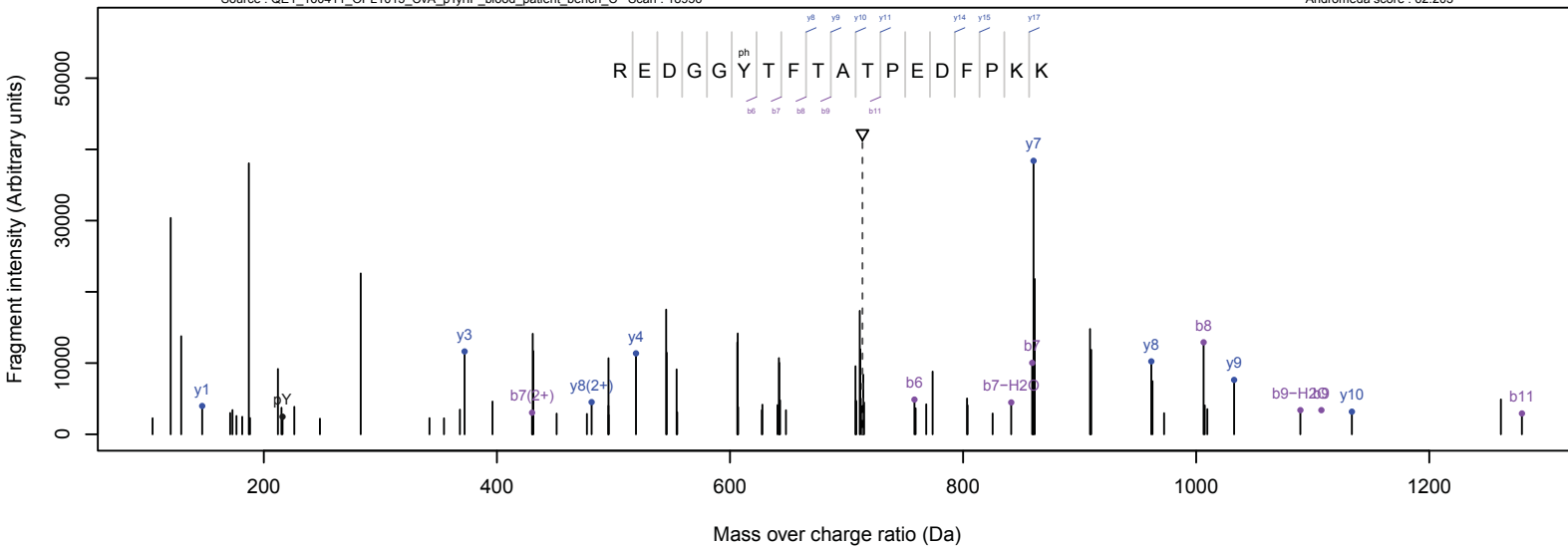
Andromeda score : 62.287



## REDGGY(ph)TFTATPEDFPKK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18956

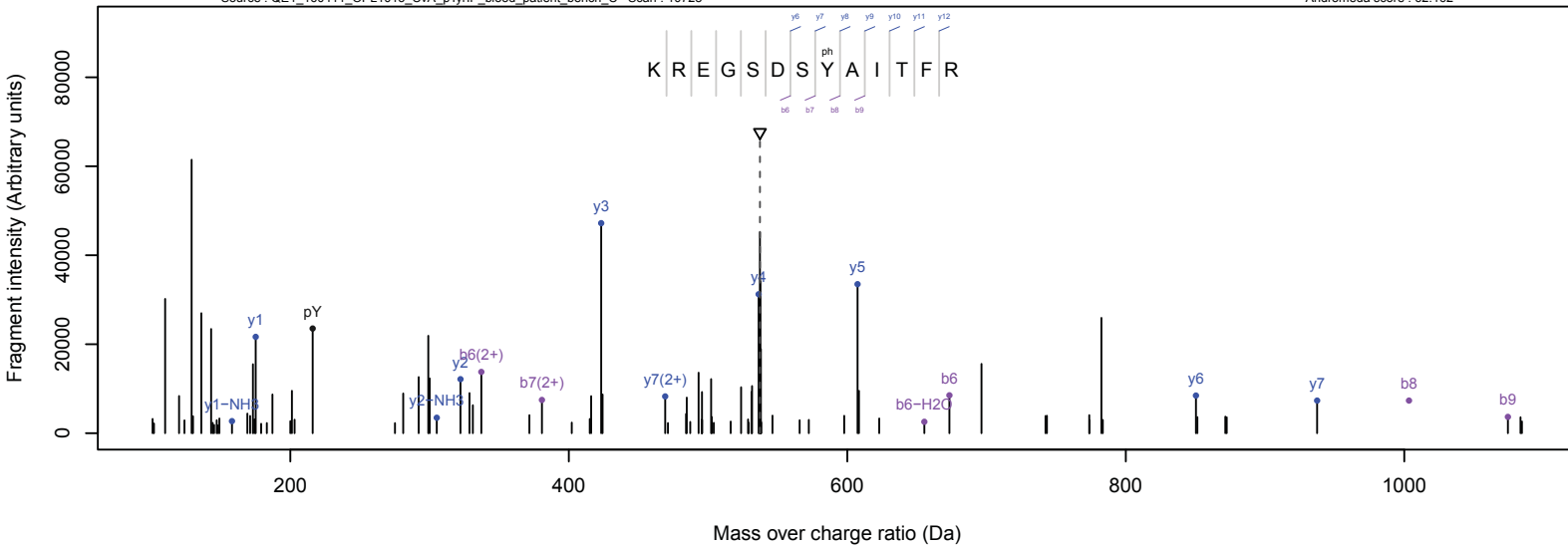
Andromeda score : 62.203



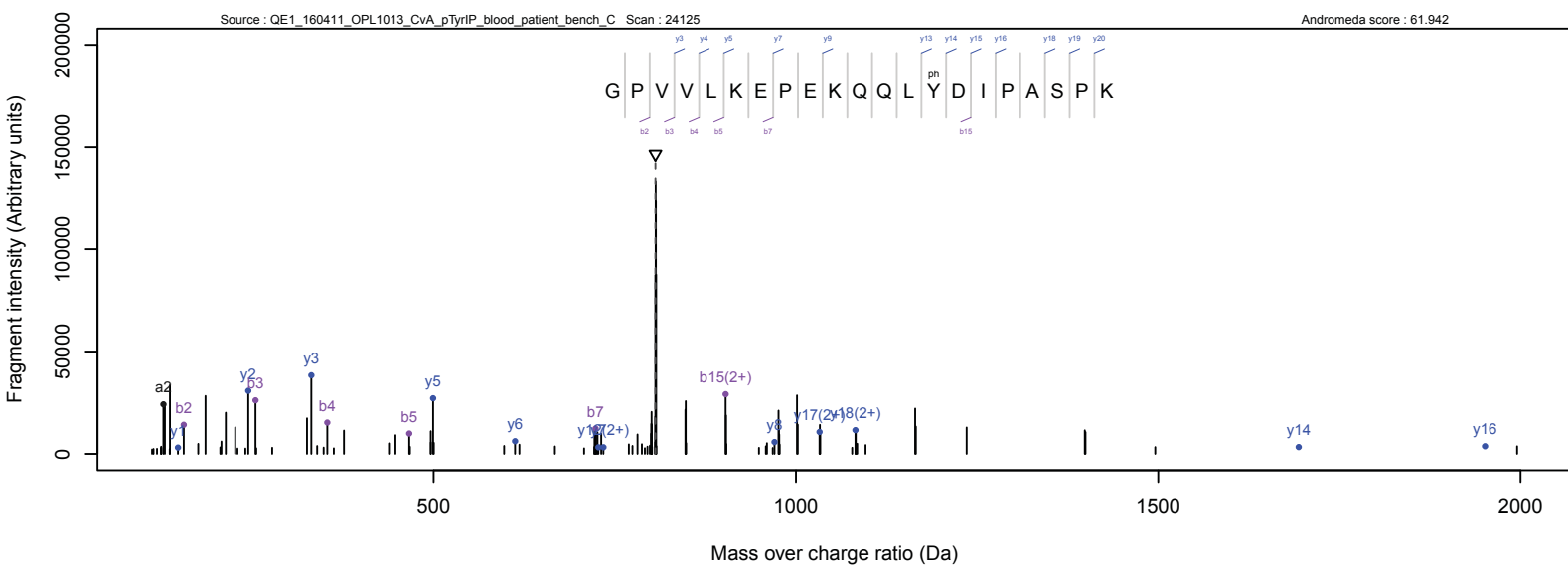
## KREGSDSY(ph)AITFR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 10728

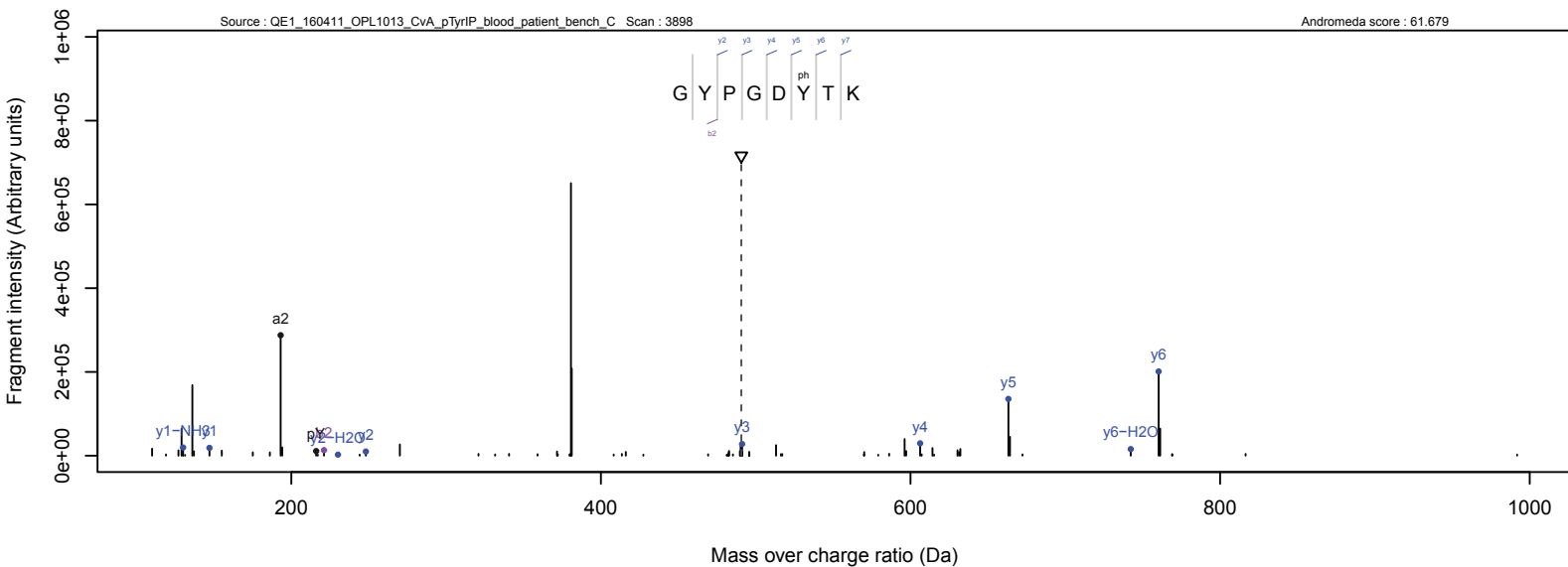
Andromeda score : 62.162



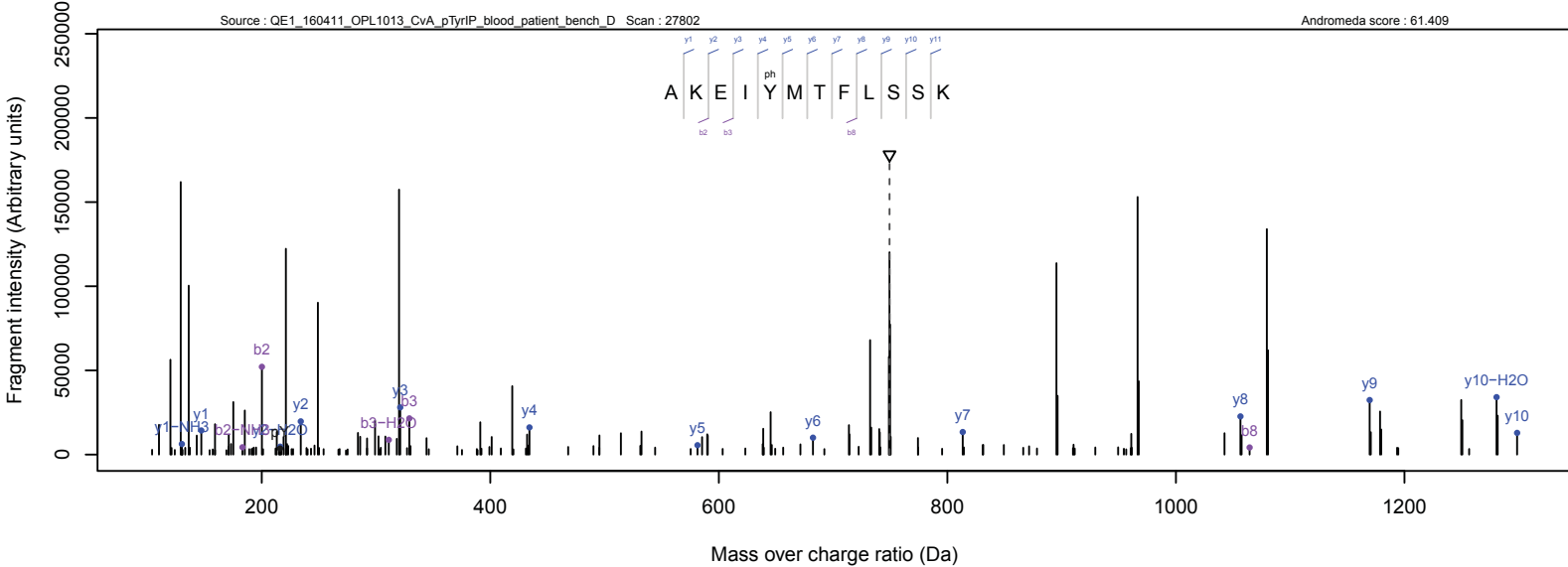
GPVVLKEPEKQLYDIPASPK-3+



GYPGDYTK-2+



AKEIYMTFLSSK-2+

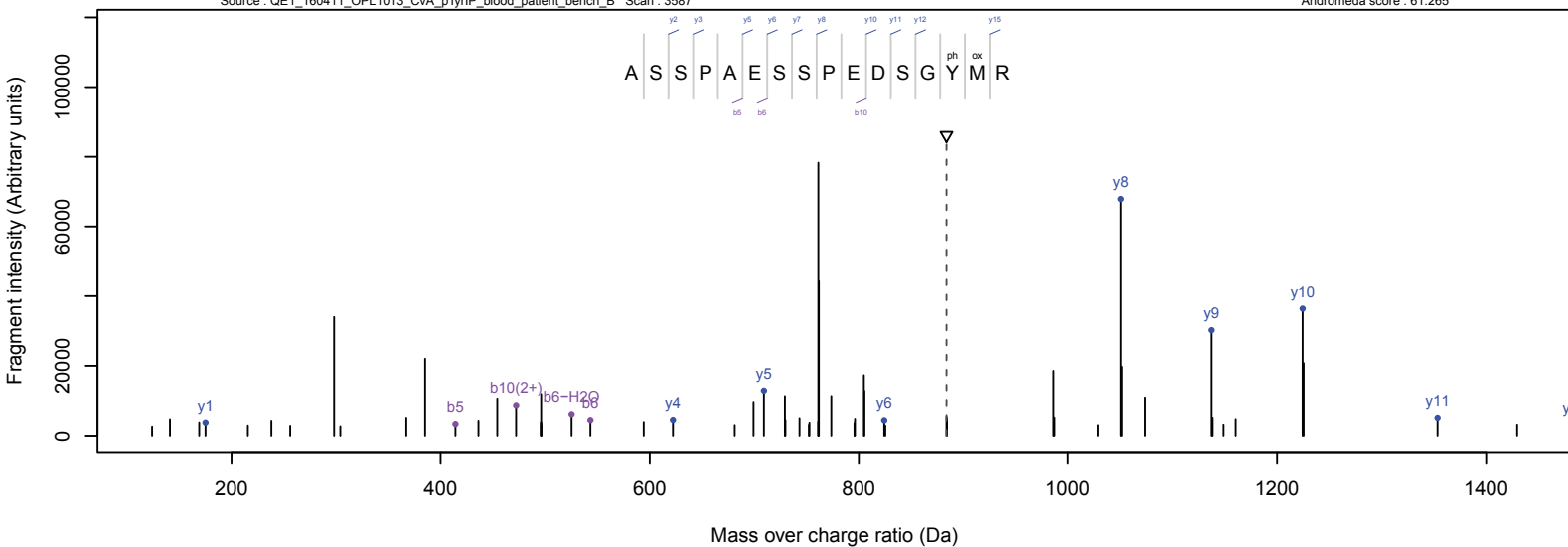




ASSPAESSPESGYM(ox)R-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 3587

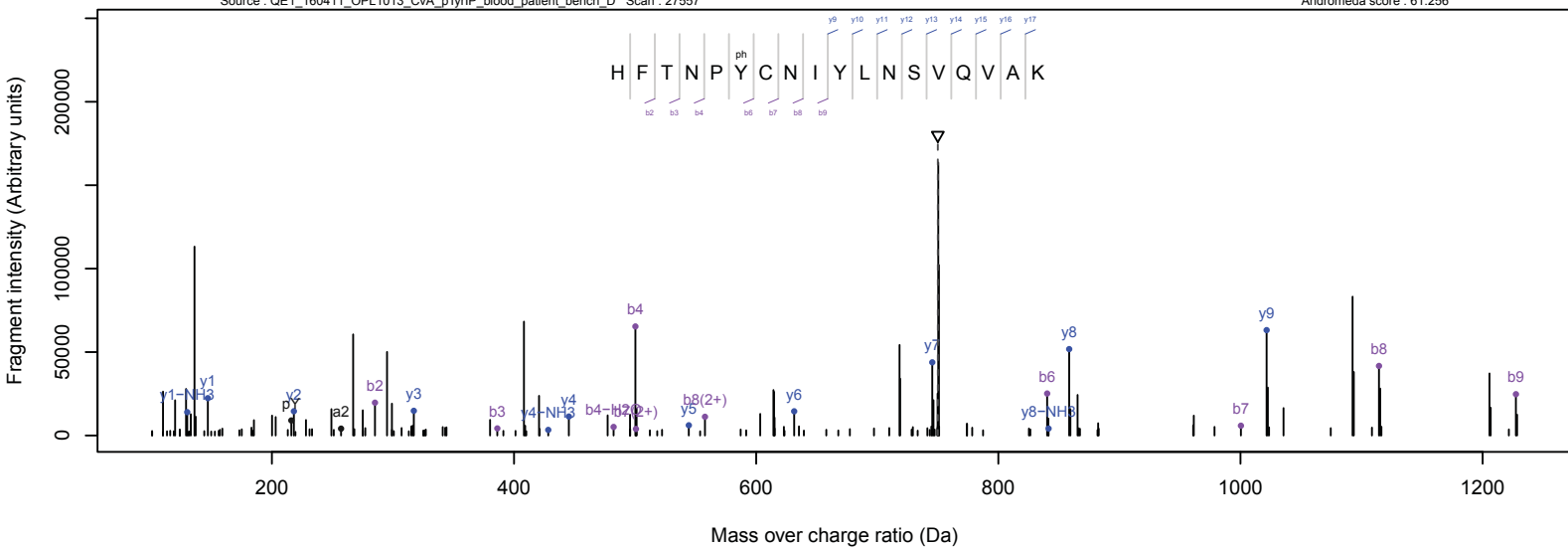
Andromeda score : 61.265



HFTNPY(ph)CNIYLSVQVAK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 27557

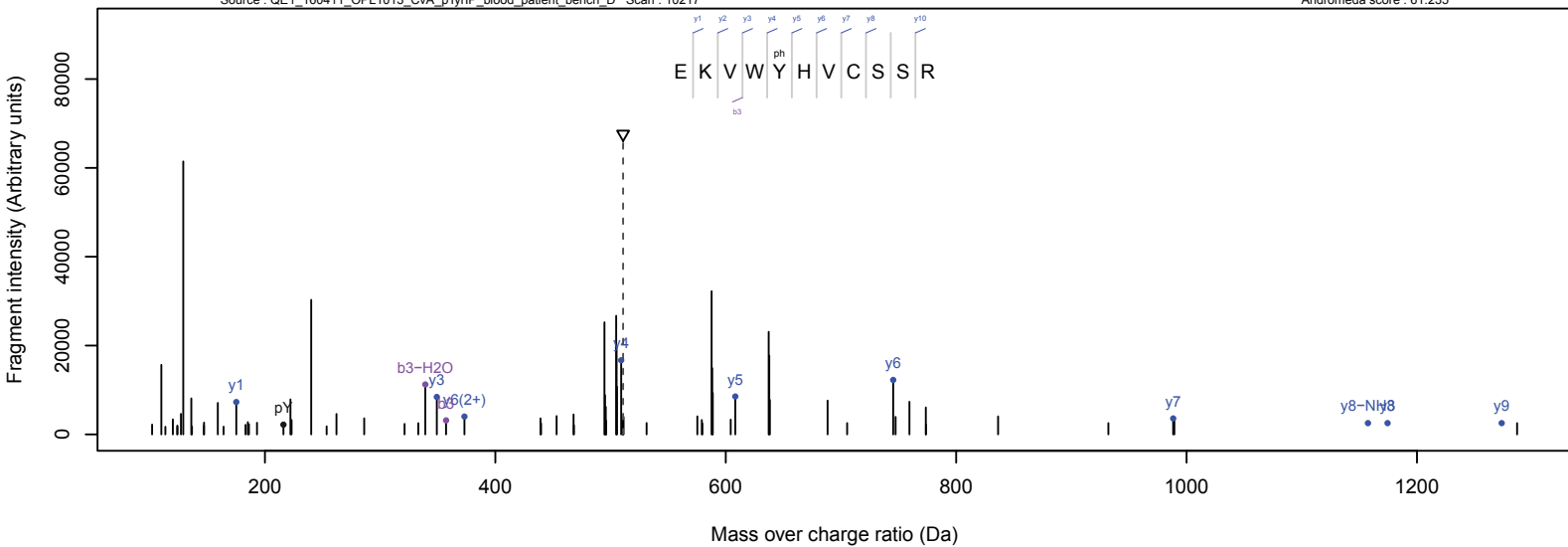
Andromeda score : 61.256



EKVWY(ph)HVCSSR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 10217

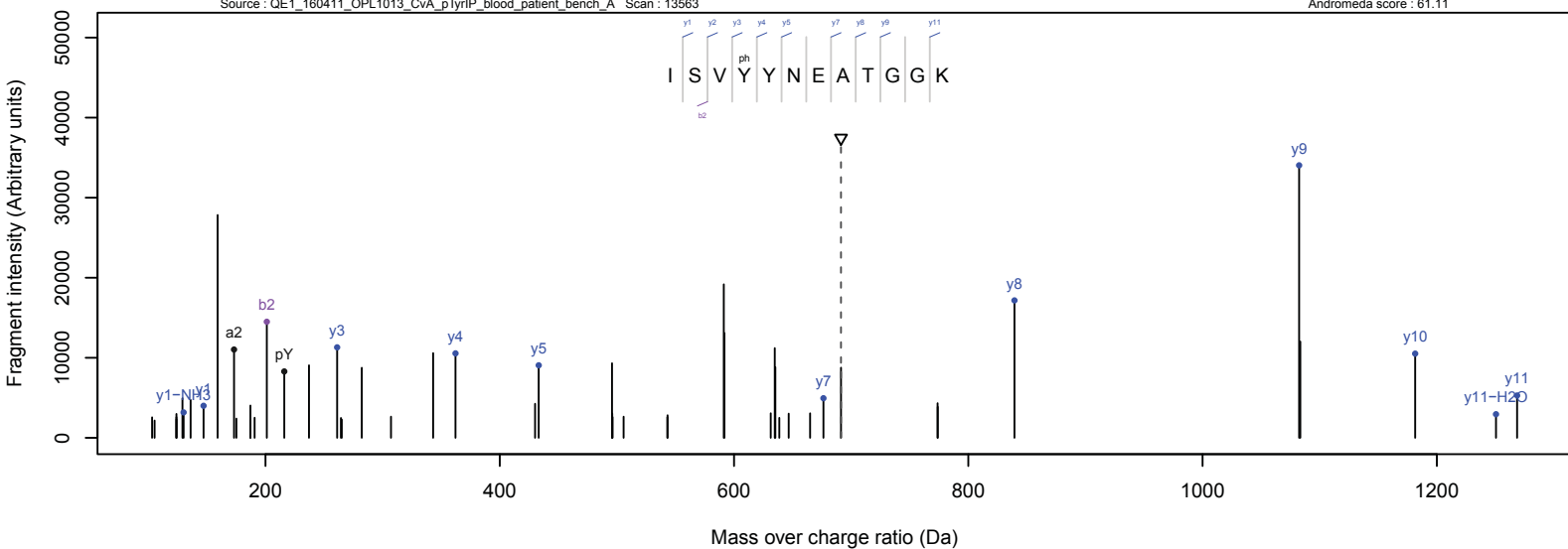
Andromeda score : 61.235



ISVY(ph)YNEATGGK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 13563

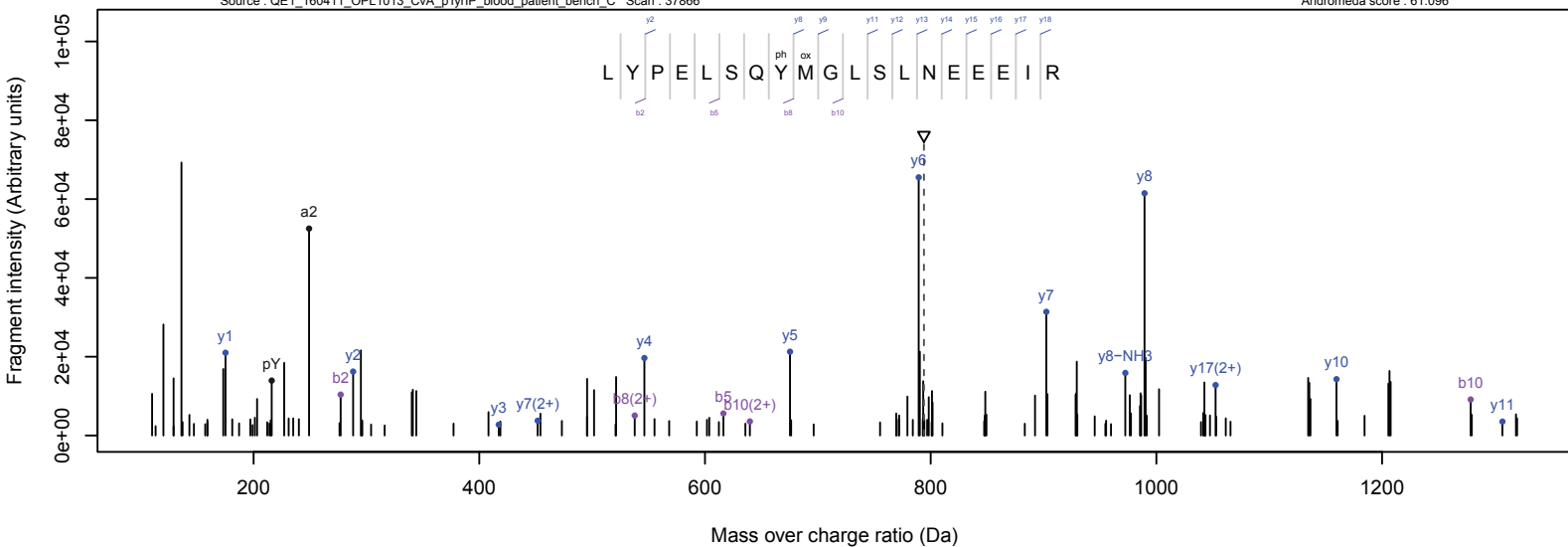
Andromeda score : 61.11



LYPELSQY(ph)M(ox)GLSNEEEIR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 37866

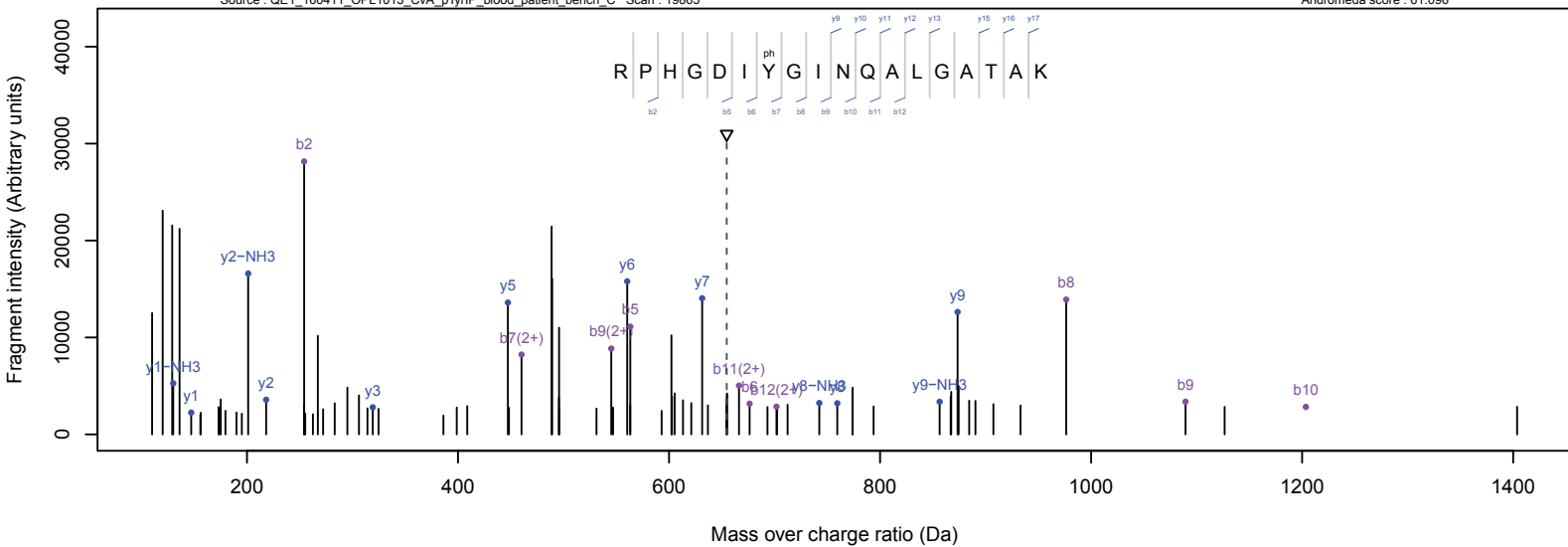
Andromeda score : 61.096



RPHGDIY(ph)GINQALGATAK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 19865

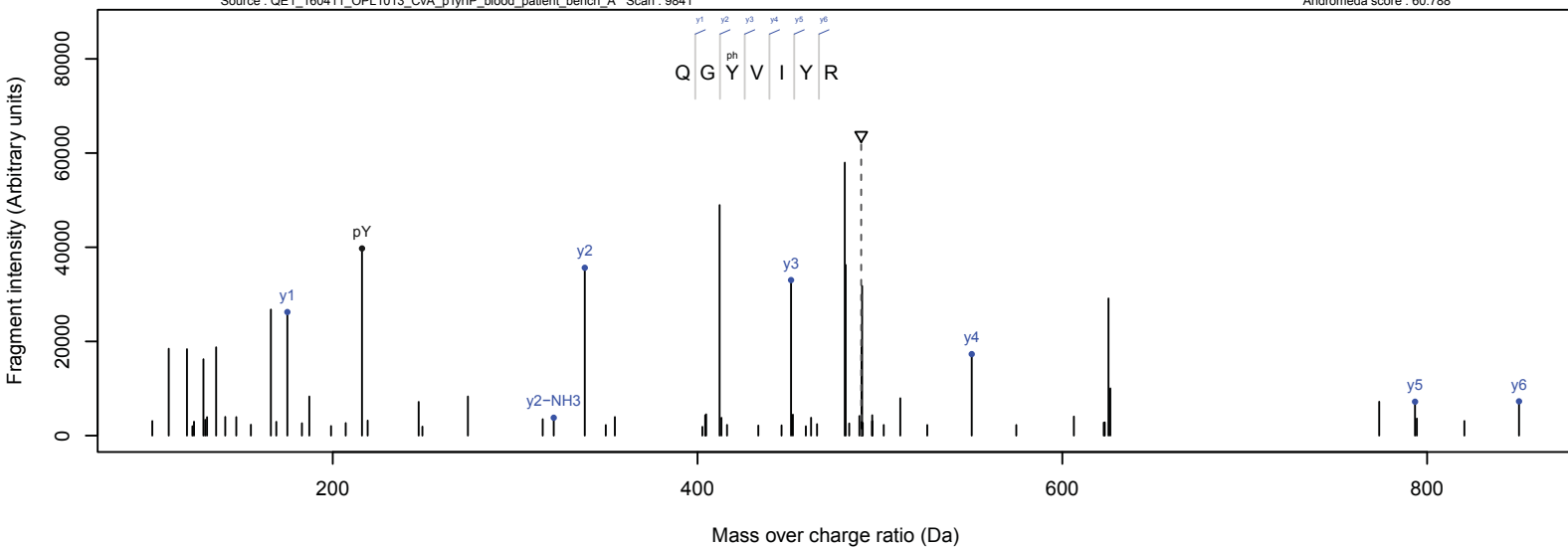
Andromeda score : 61.096



**QGY(ph)VIYR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 9841

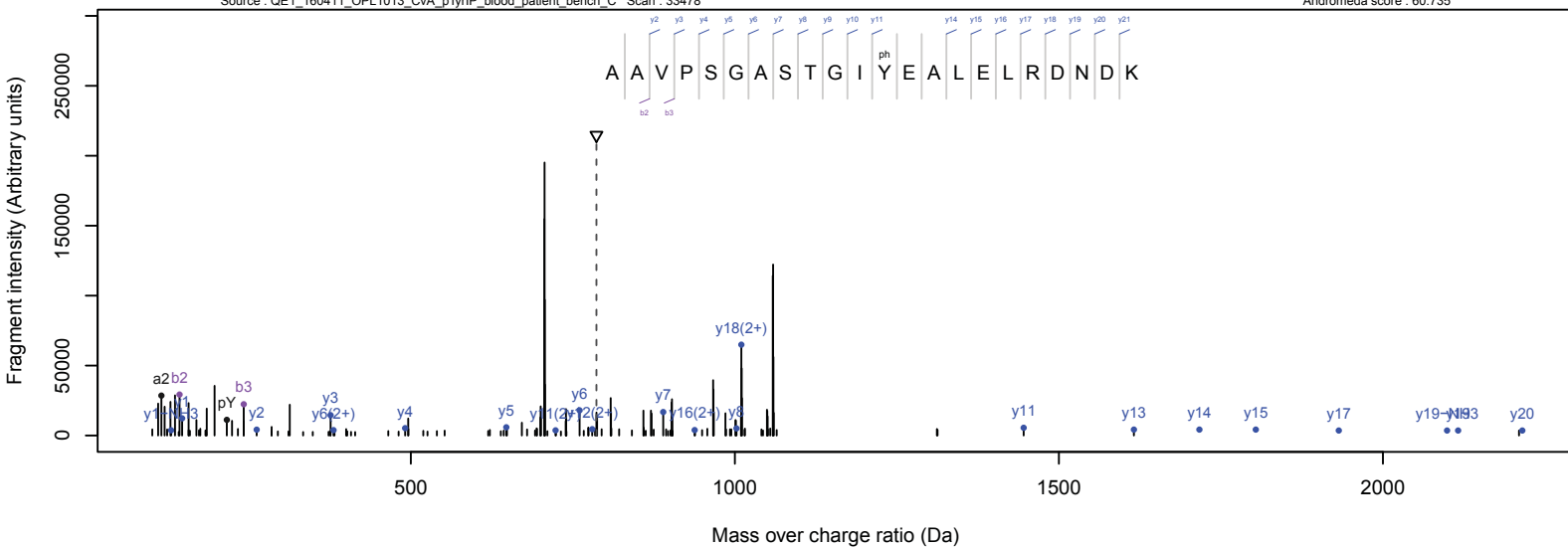
Andromeda score : 60.788



**AAVPSGASTGIY(ph)EALERDNDK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 33478

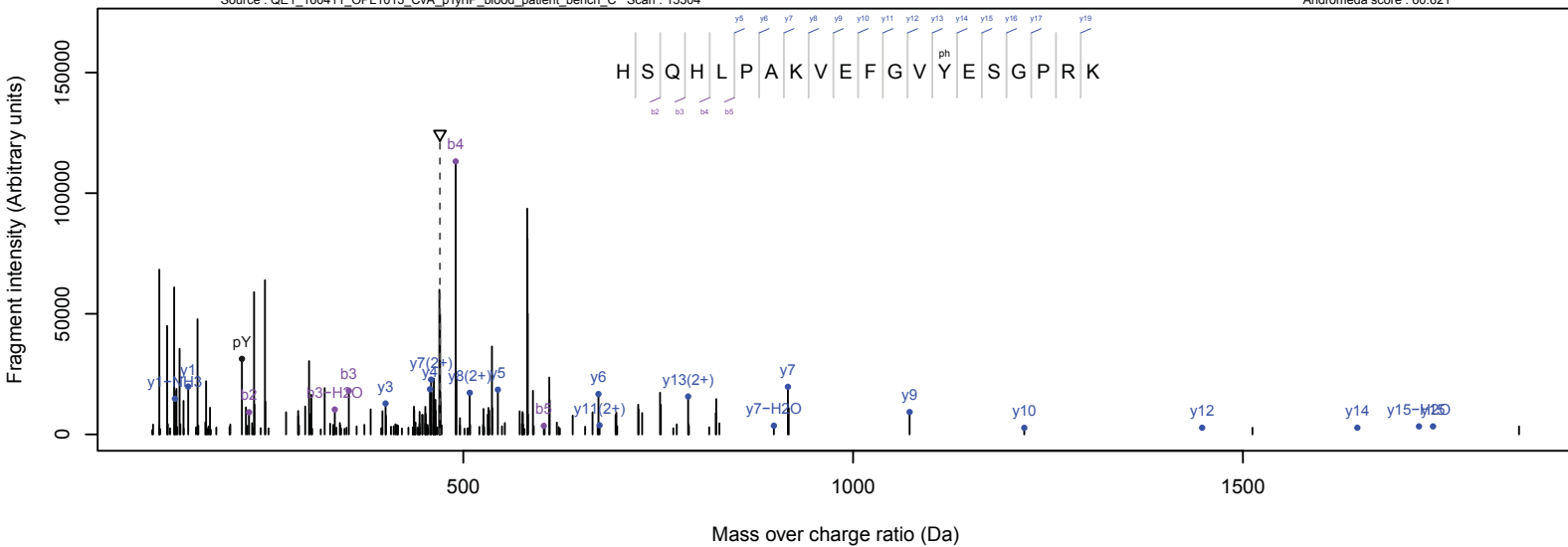
Andromeda score : 60.735



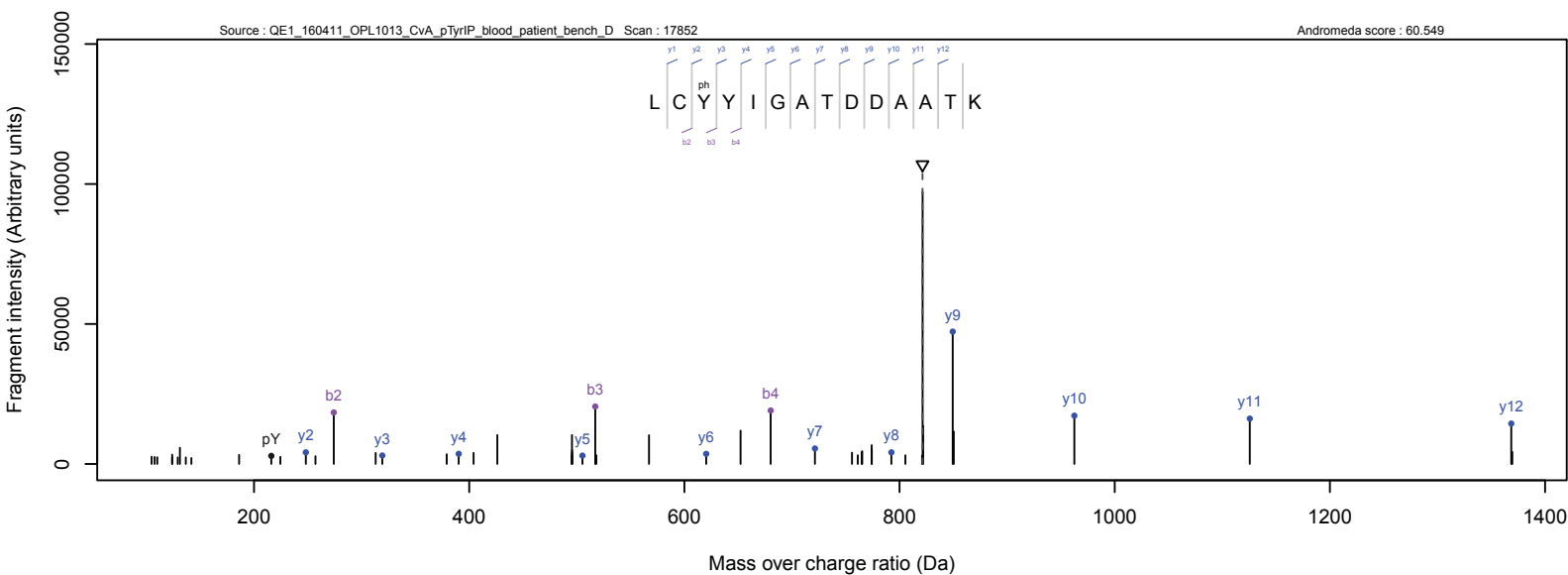
**HSQHLPKVEFGVY(ph)ESGPRK-5+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 15304

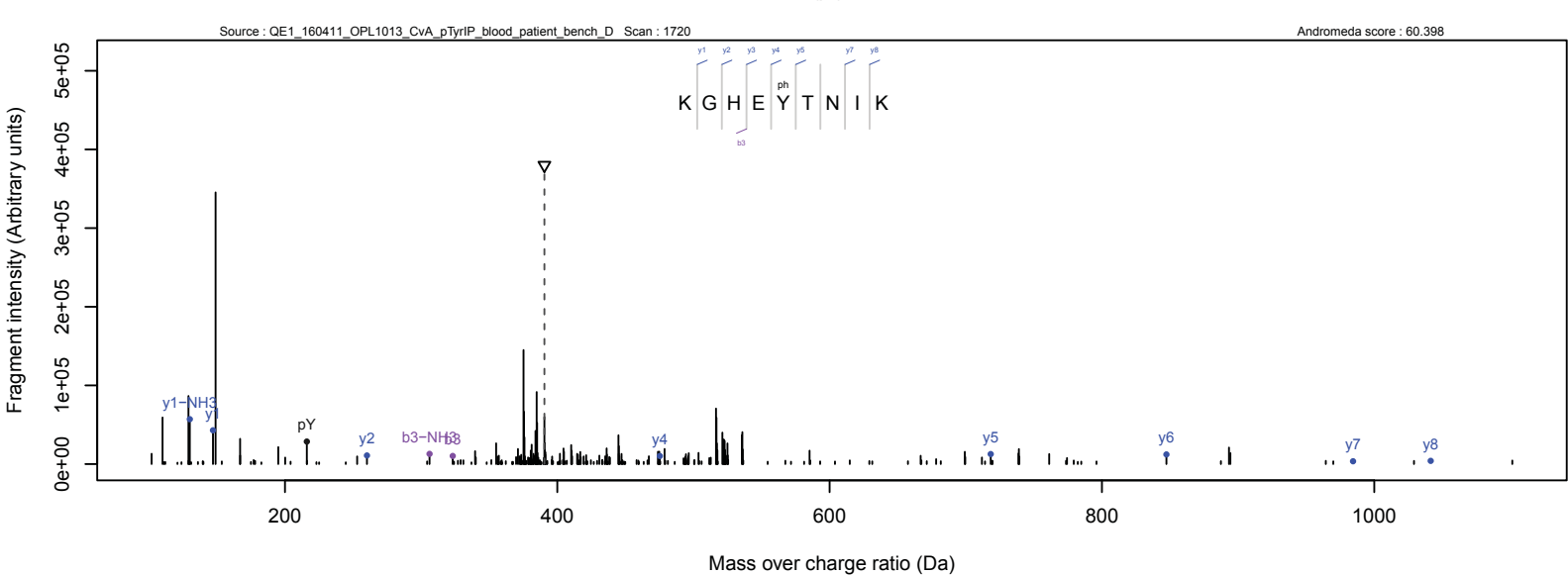
Andromeda score : 60.621



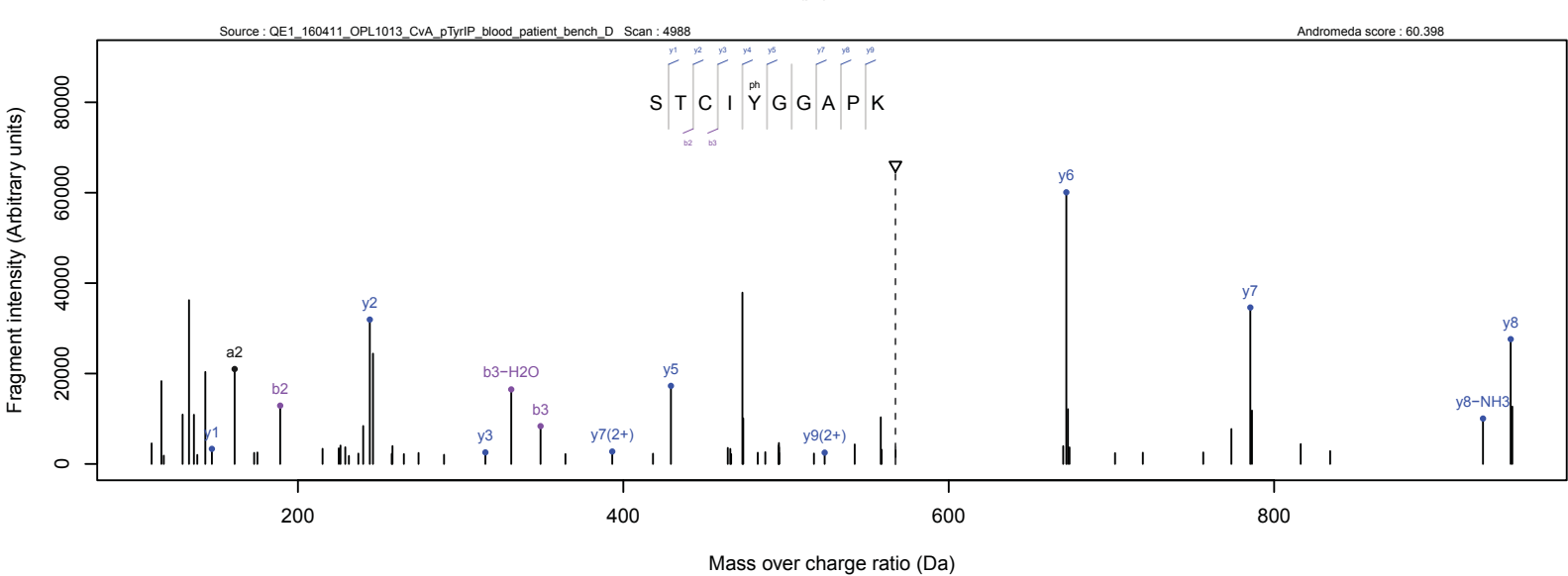
## LCY(ph)YIGATDDAATK-2+



## KGHEY(ph)TNIK-3+



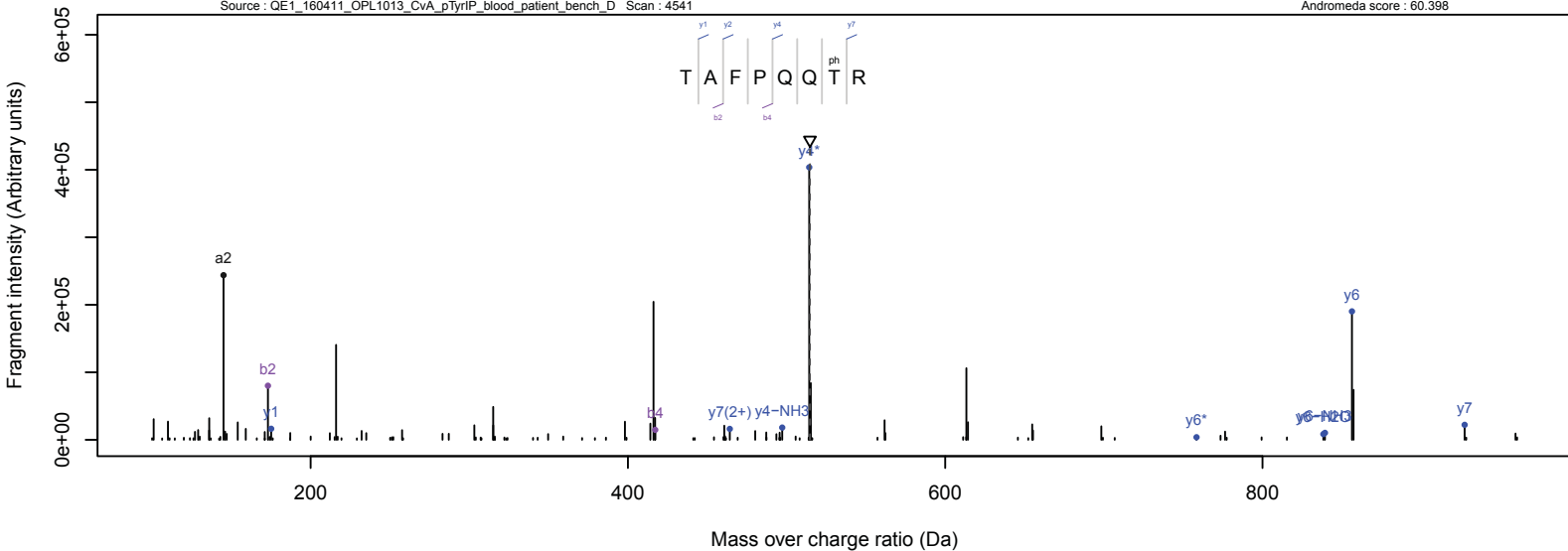
## STCIY(ph)GGAPK-2+



TAFPQQ(ph)R-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 4541

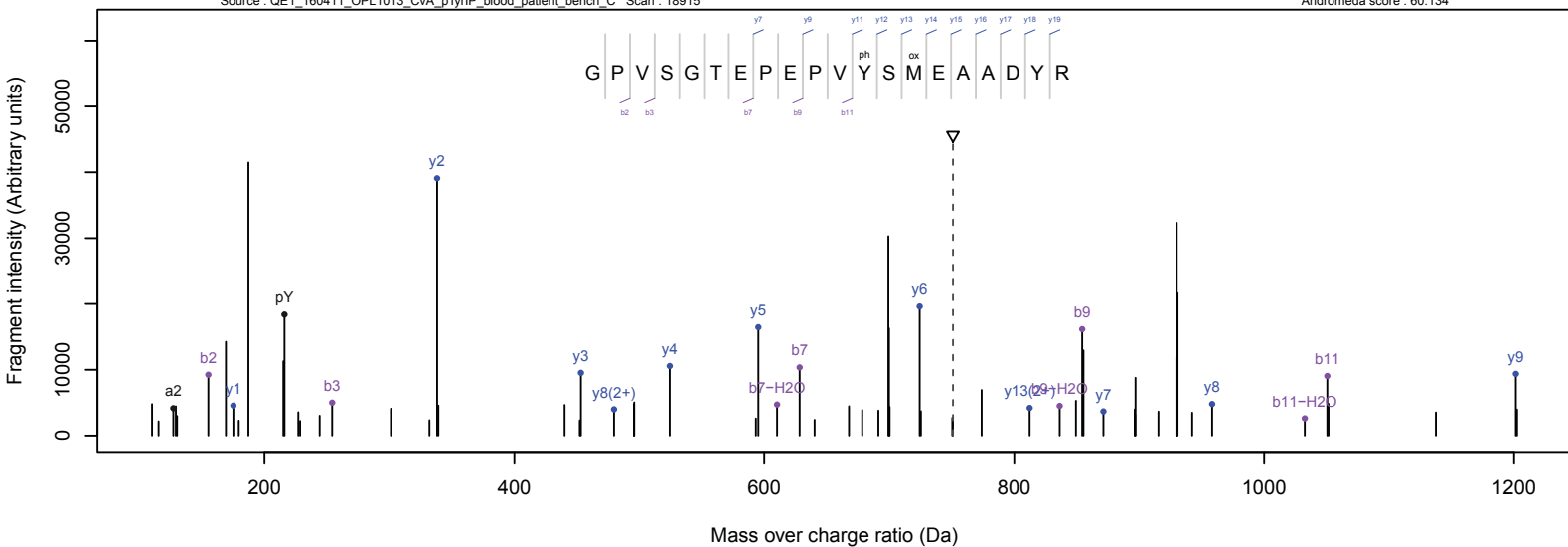
Andromeda score : 60.398



GPVSGTEPEPV(ph)SM(ox)EAADYR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18915

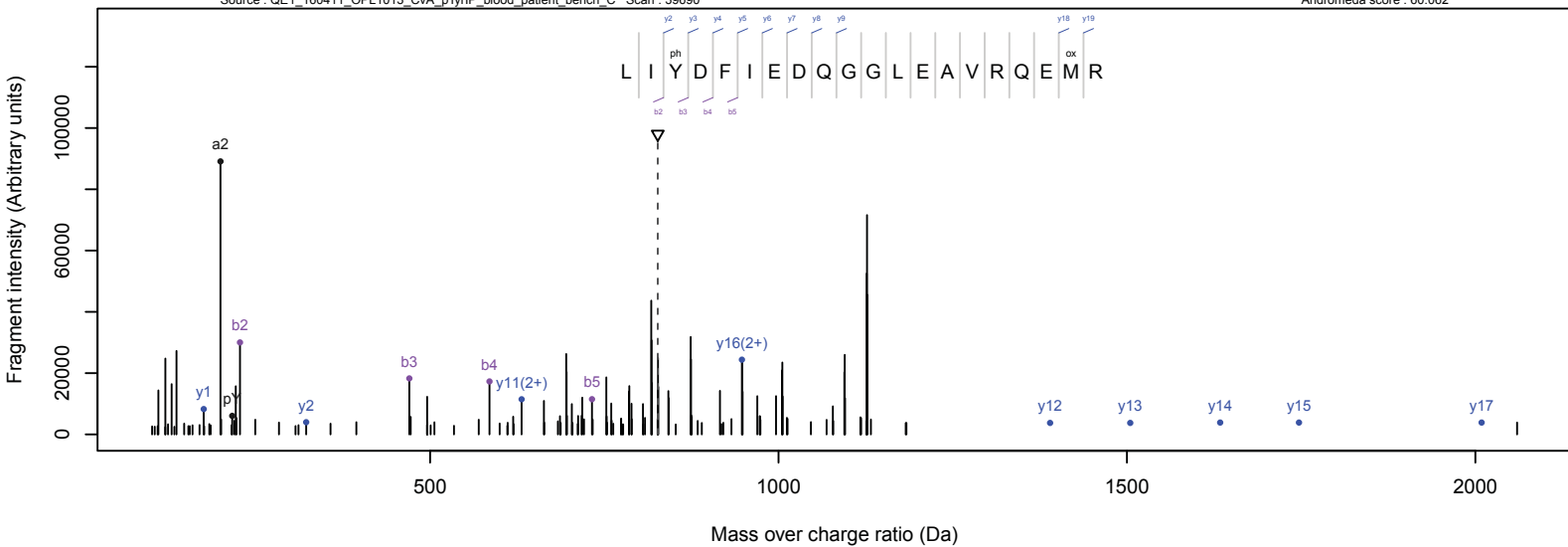
Andromeda score : 60.134



LIY(ph)DFIEDQGGLEAVRQEM(ox)R-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 39690

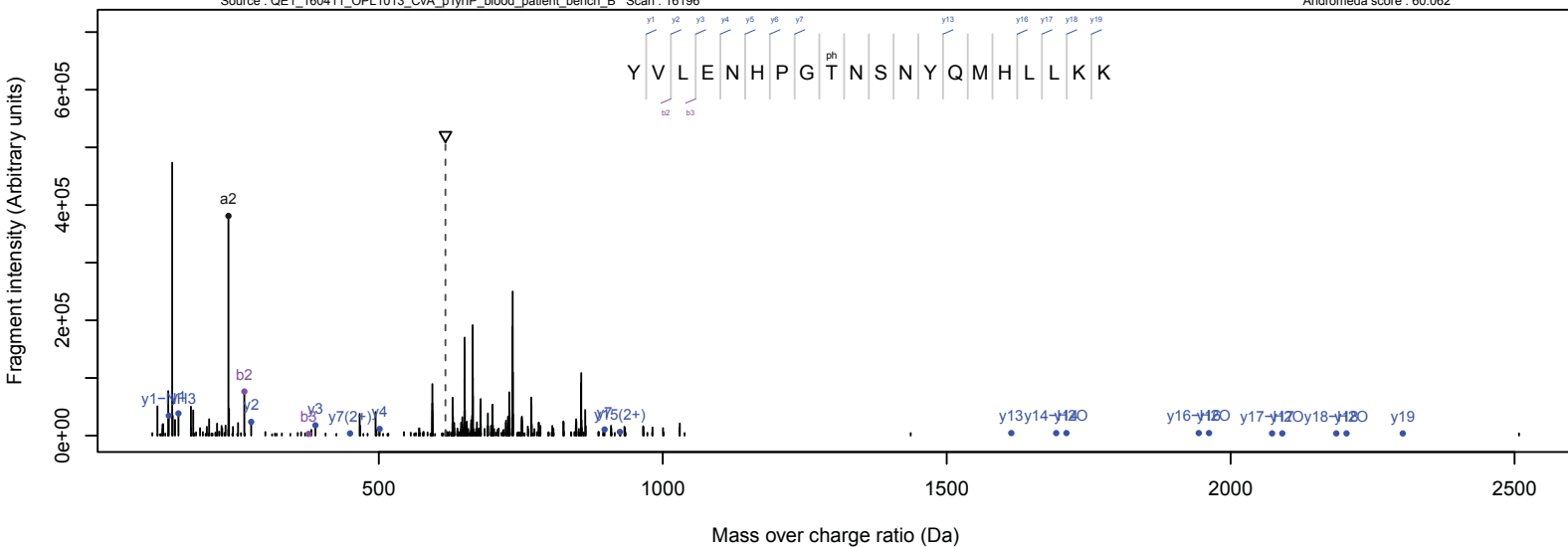
Andromeda score : 60.062



**YVLENHPG(ph)NSNYQMHLKK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 16196

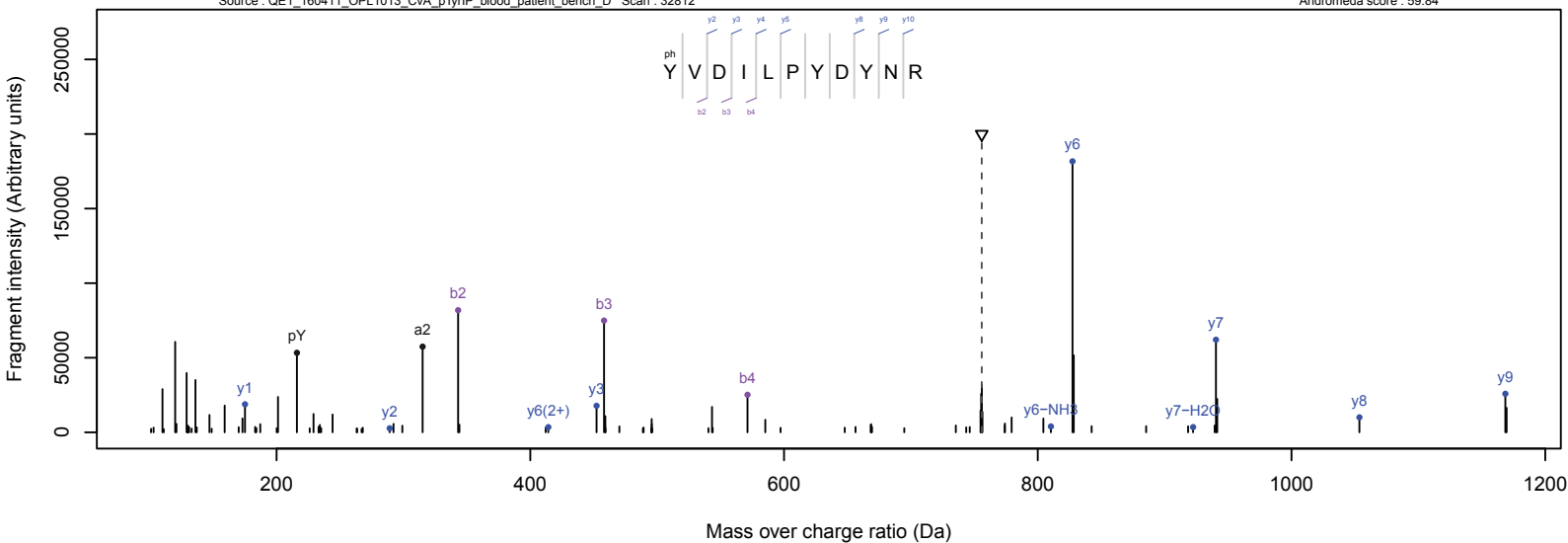
Andromeda score : 60.062



**Y(ph)VDILPYDYNR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 32812

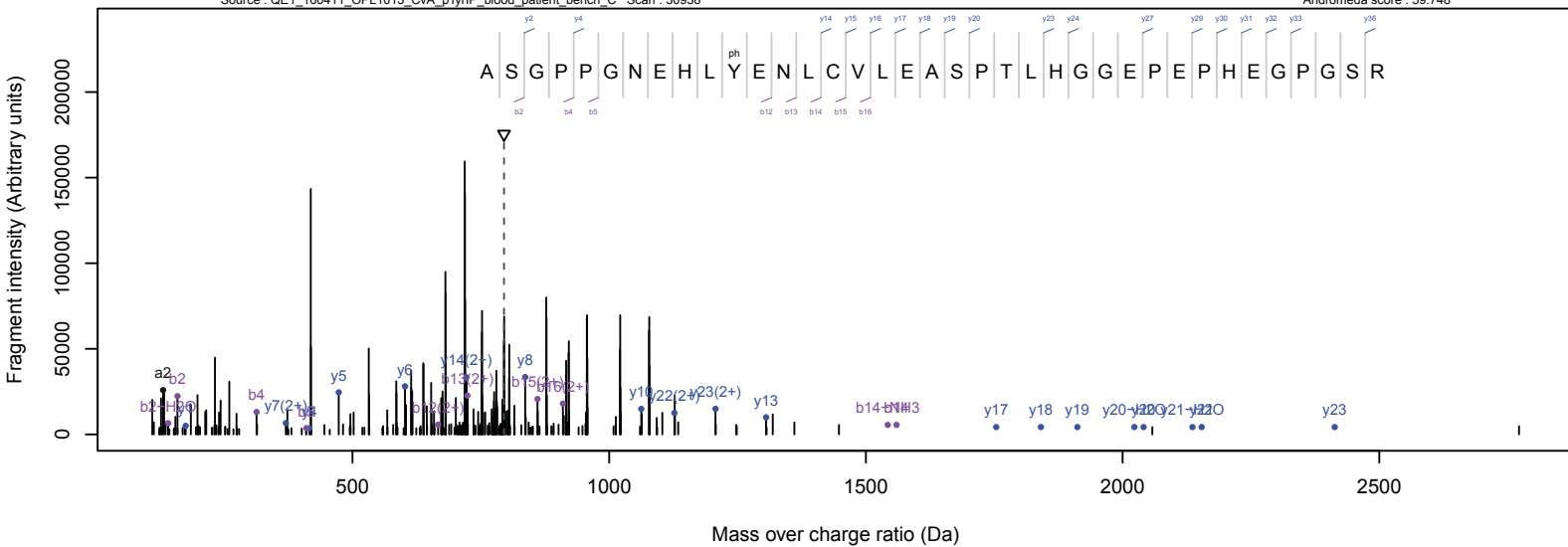
Andromeda score : 59.84



**ASGPPGNEHLY(ph)ENLCVLEASPTLHGGEPEPEHGGPSR-5+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 30938

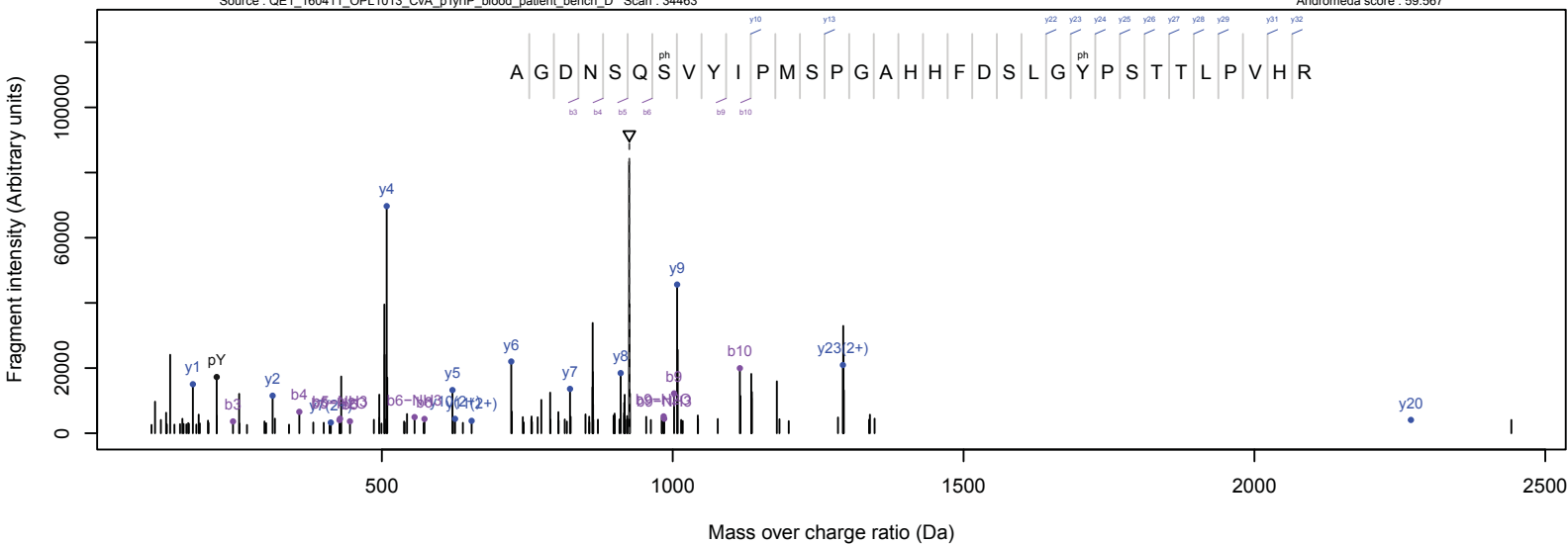
Andromeda score : 59.748



AGDNSQS(ph)VYIPMSPGAHHFDSLGY(ph)PSTTLPVHR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 34463

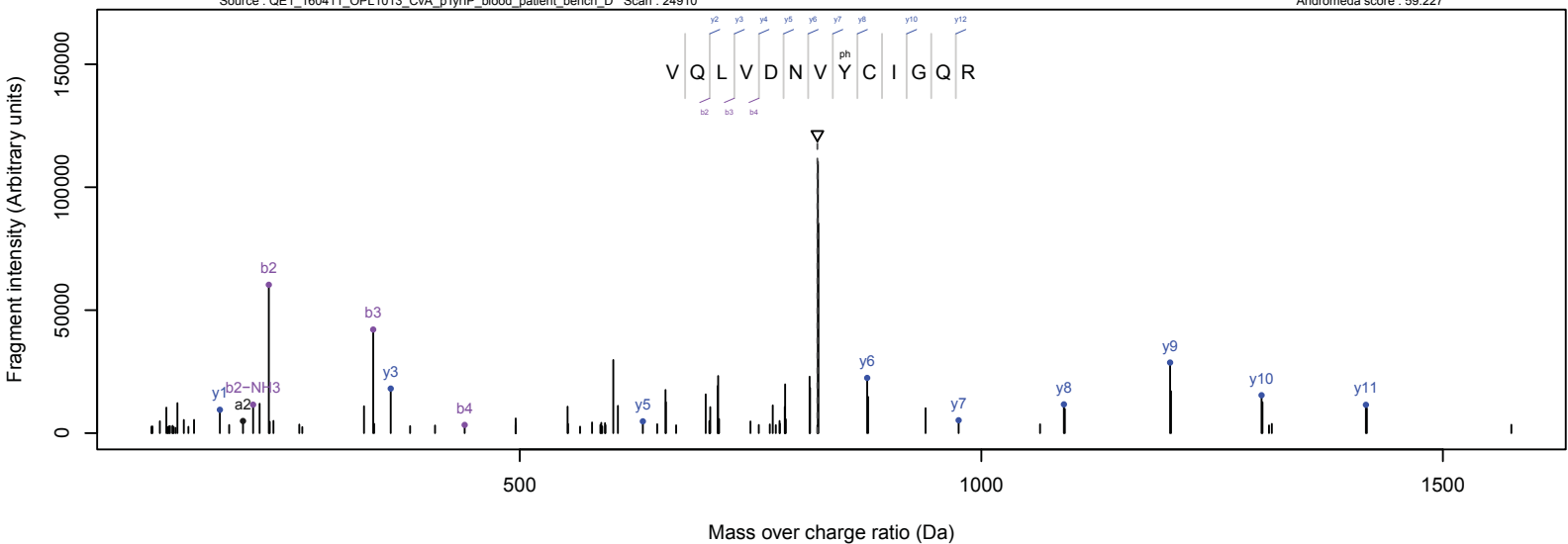
Andromeda score : 59.567



VQLVDNVY(ph)CIGQR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24910

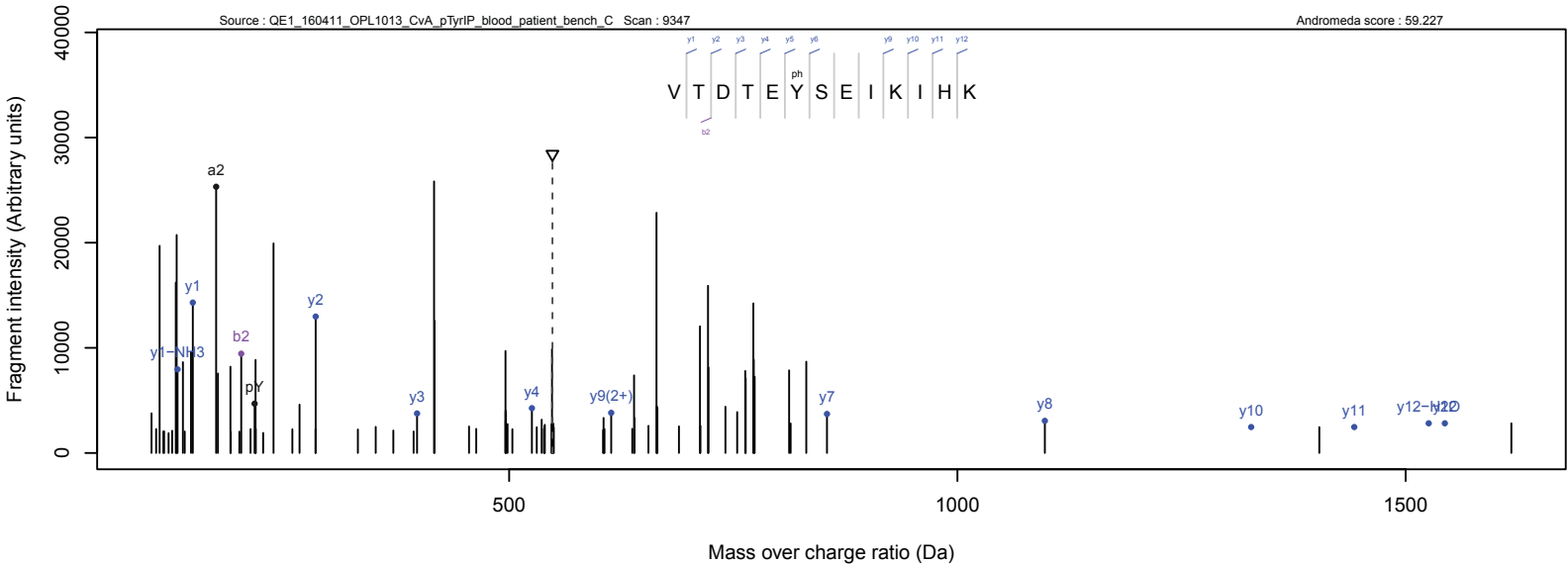
Andromeda score : 59.227



VTDEY(ph)SEIKHK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 9347

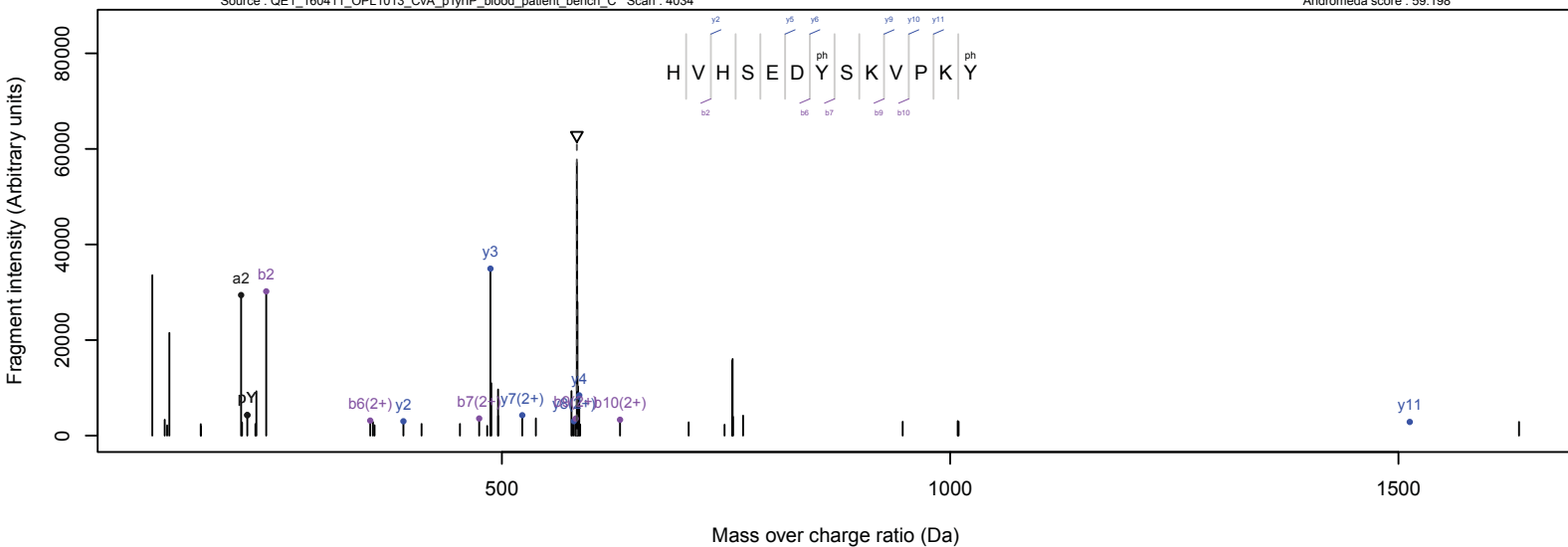
Andromeda score : 59.227



## HVHSEDY(ph)SKVPKY(ph)-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 4034

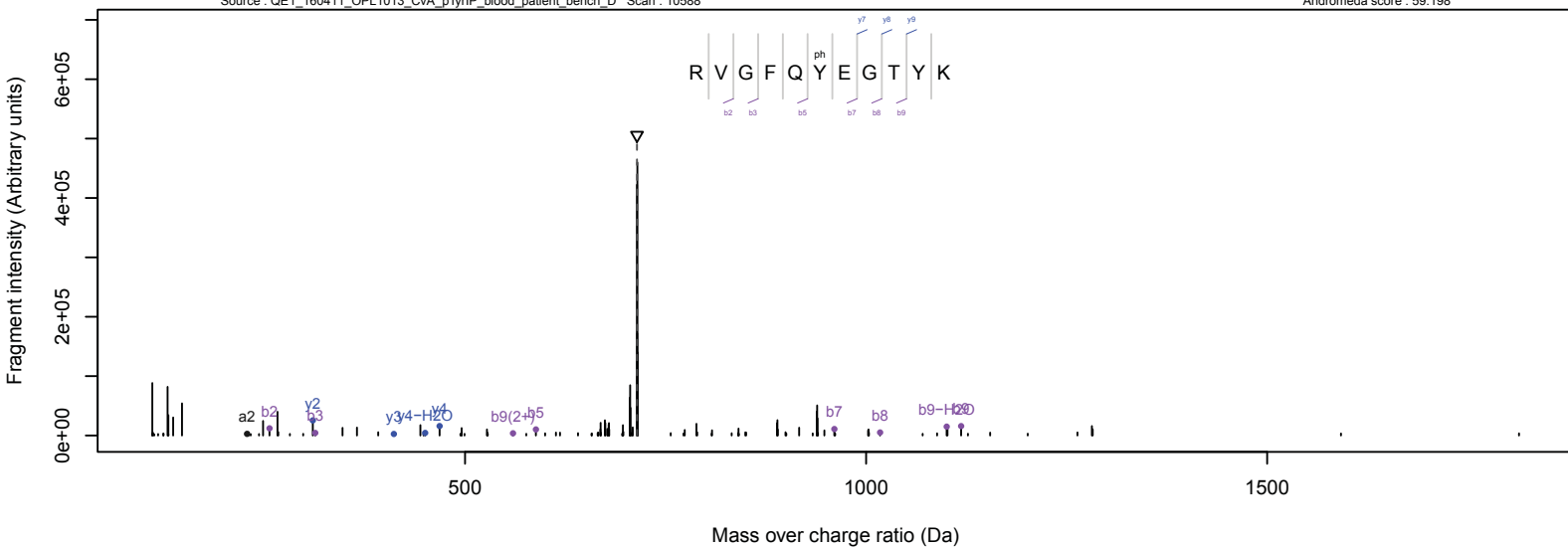
Andromeda score : 59.198



## RVGFQY(ph)EGTYK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 10588

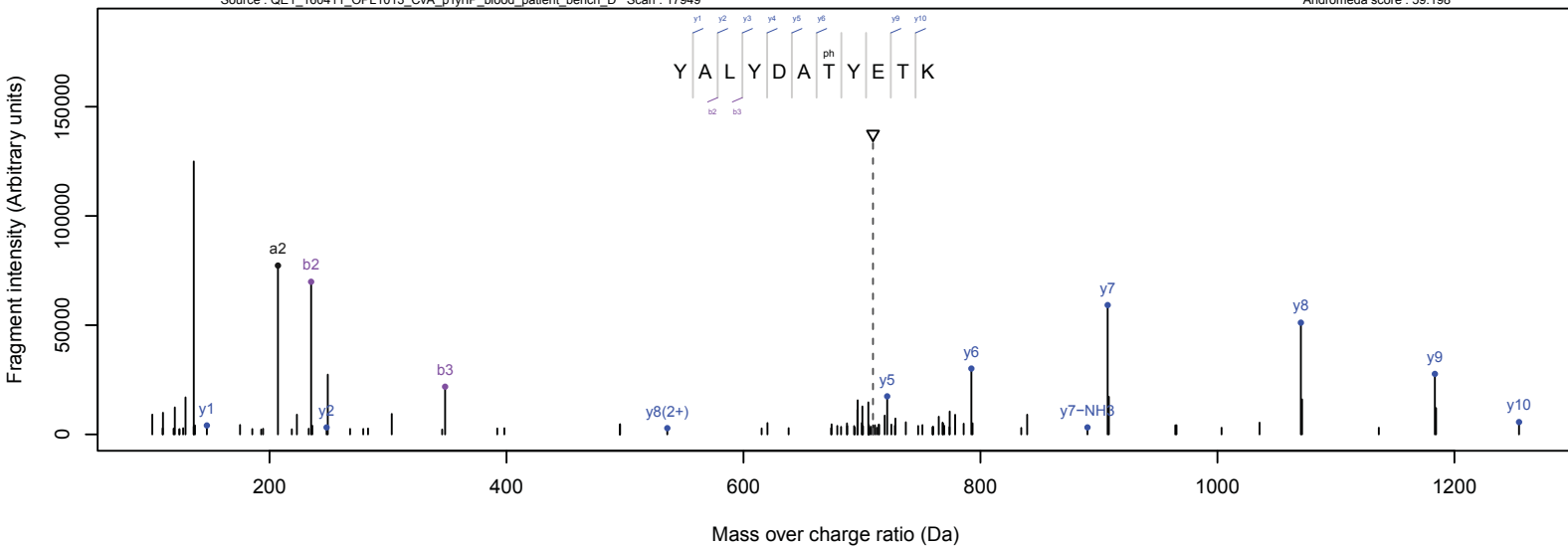
Andromeda score : 59.198



## YALYDAT(ph)YETK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17949

Andromeda score : 59.198

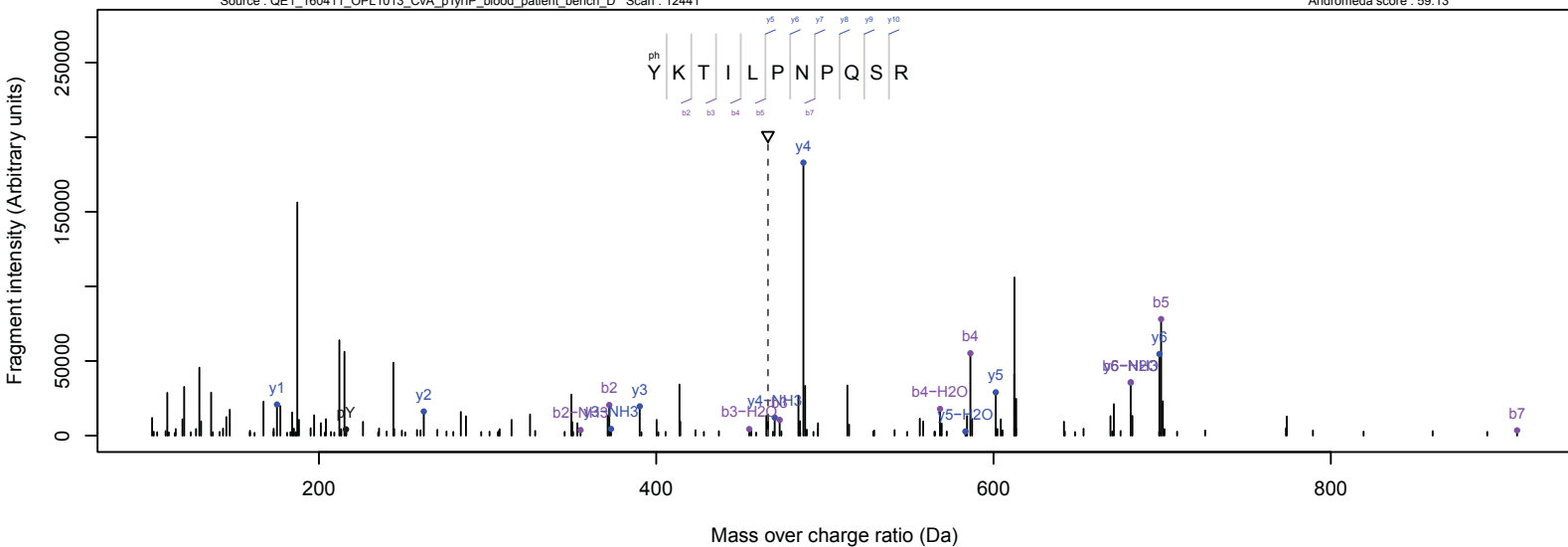




Y(ph)KTILPNPQSR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12441

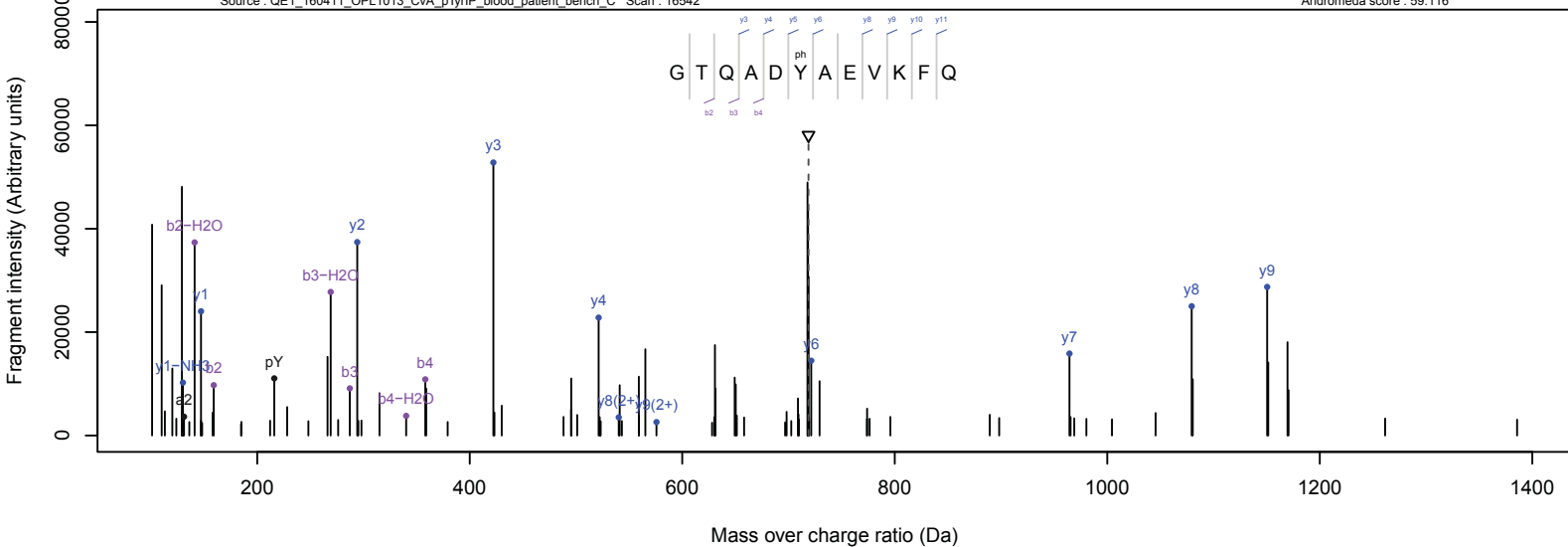
Andromeda score : 59.13



GTQADYAEVKFQ-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16542

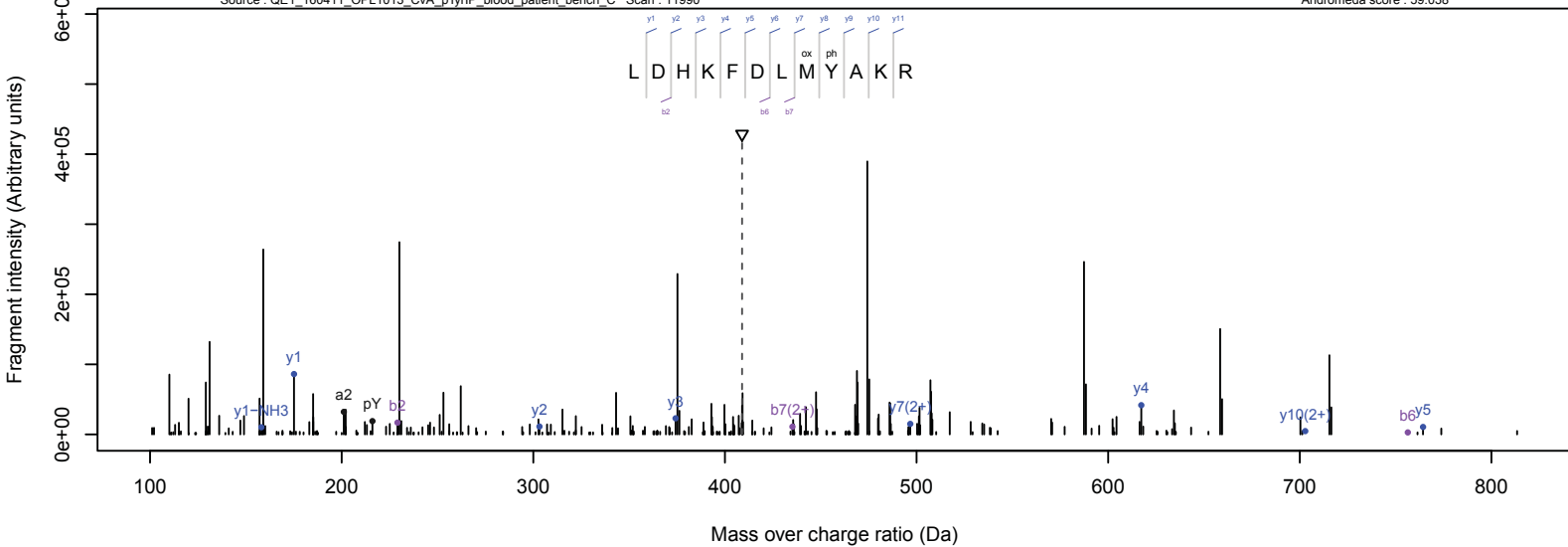
Andromeda score : 59.116



LDHKFDLM(ox)Y(ph)AKR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 11990

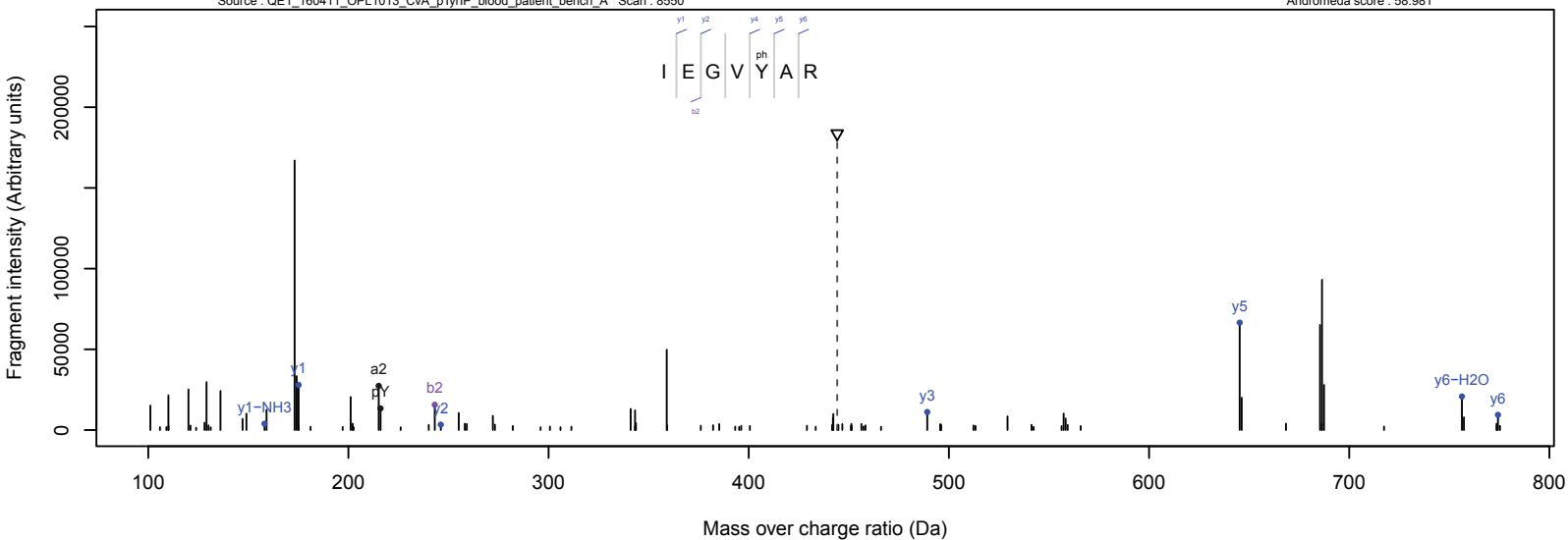
Andromeda score : 59.038



## IEGVY(ph)AR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 8550

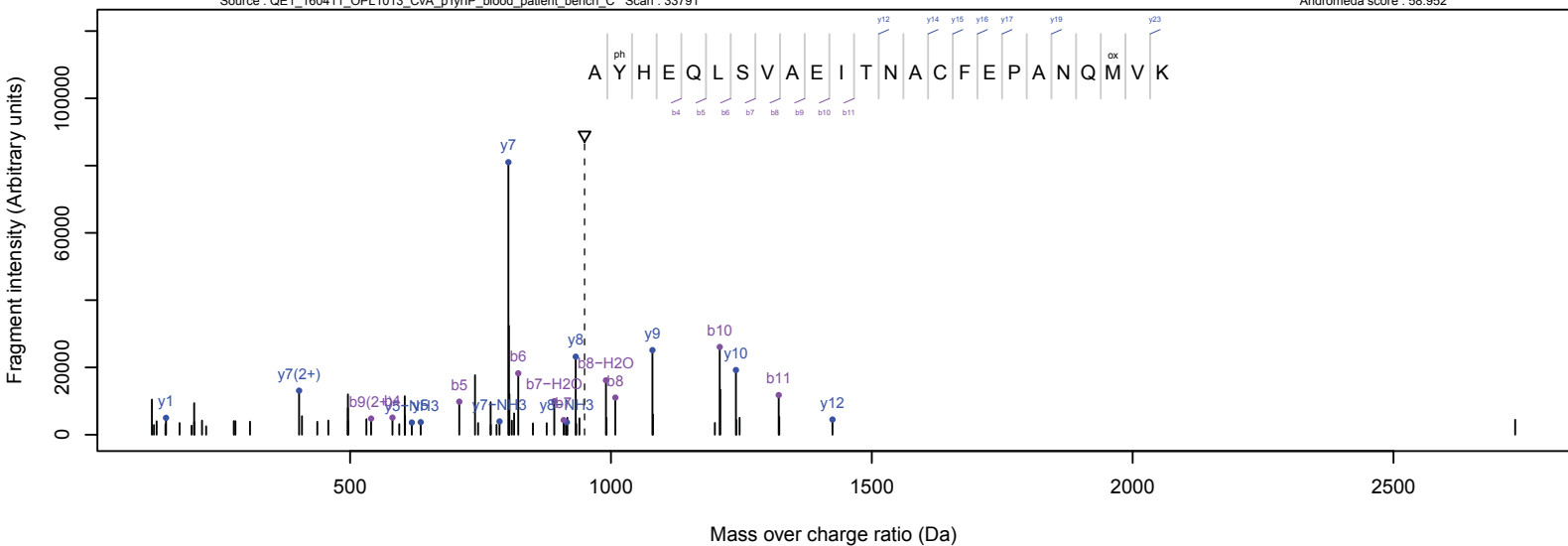
Andromeda score: 58.981



## AY(ph)HEQLSVAEITNACFEPANQM(ox)VK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 33791

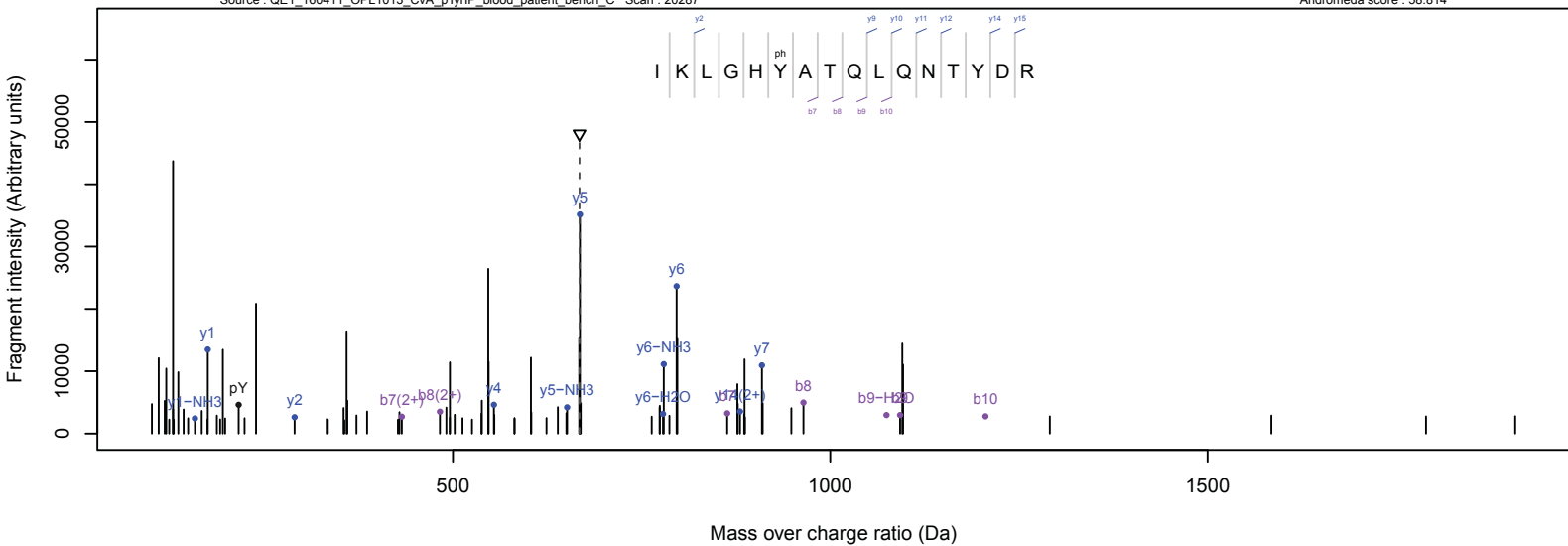
Andromeda score: 58.952



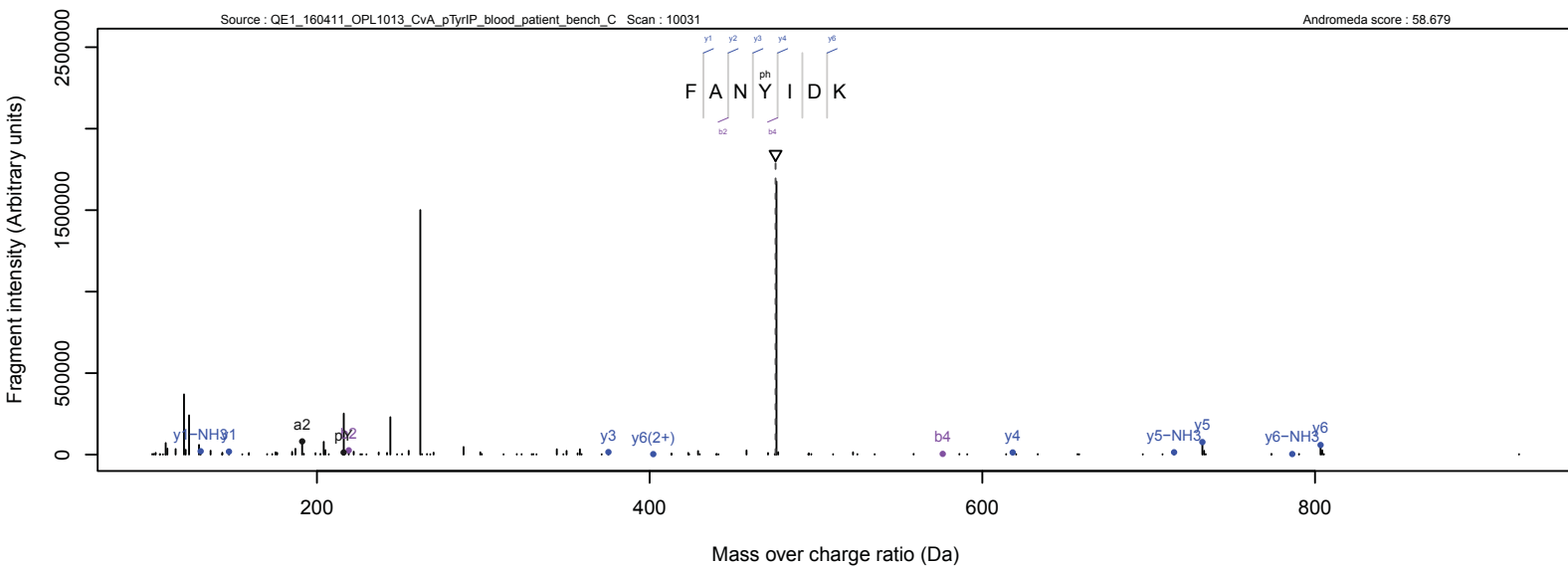
## IKLGHY(ph)ATQLQNTYDR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 20287

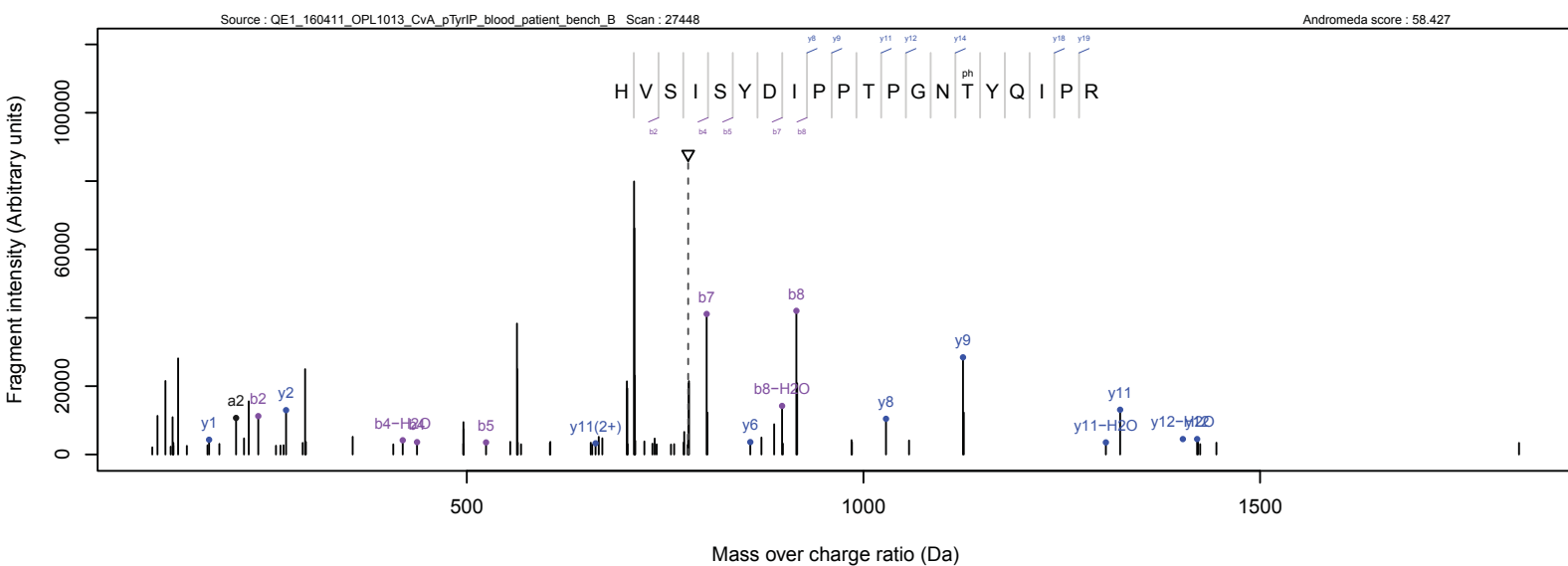
Andromeda score: 58.814



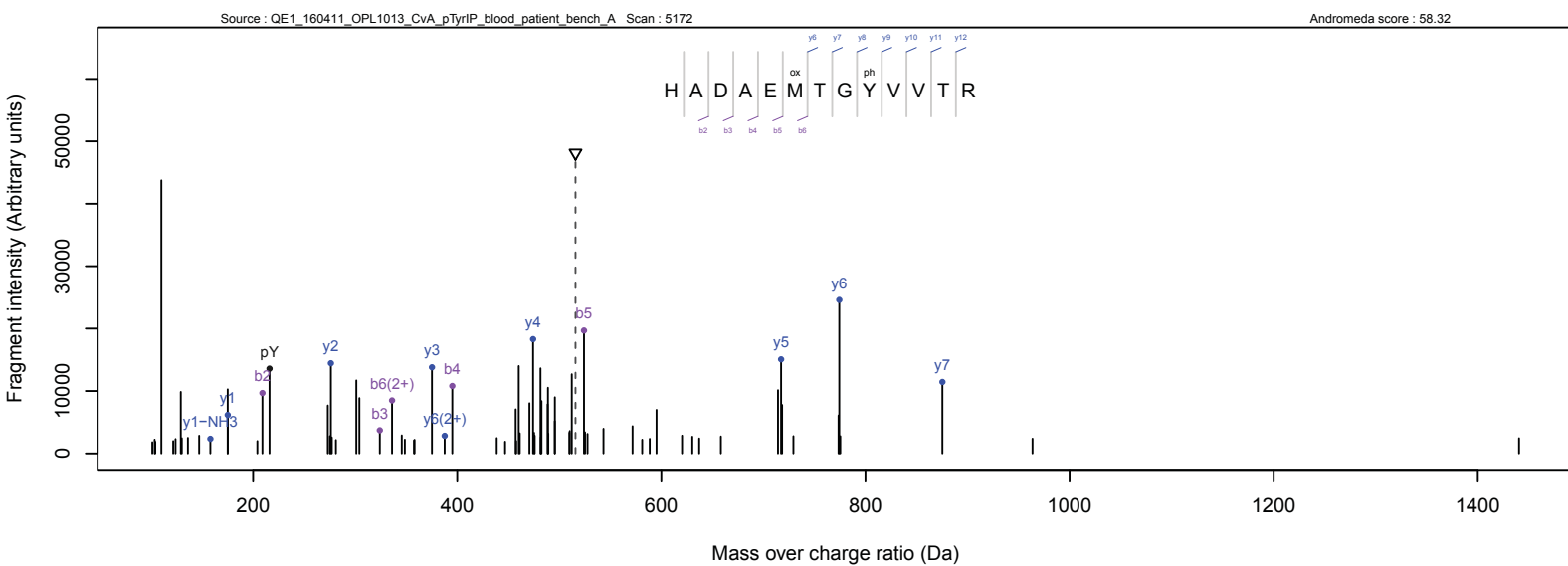
FANY(ph)IDK-2+



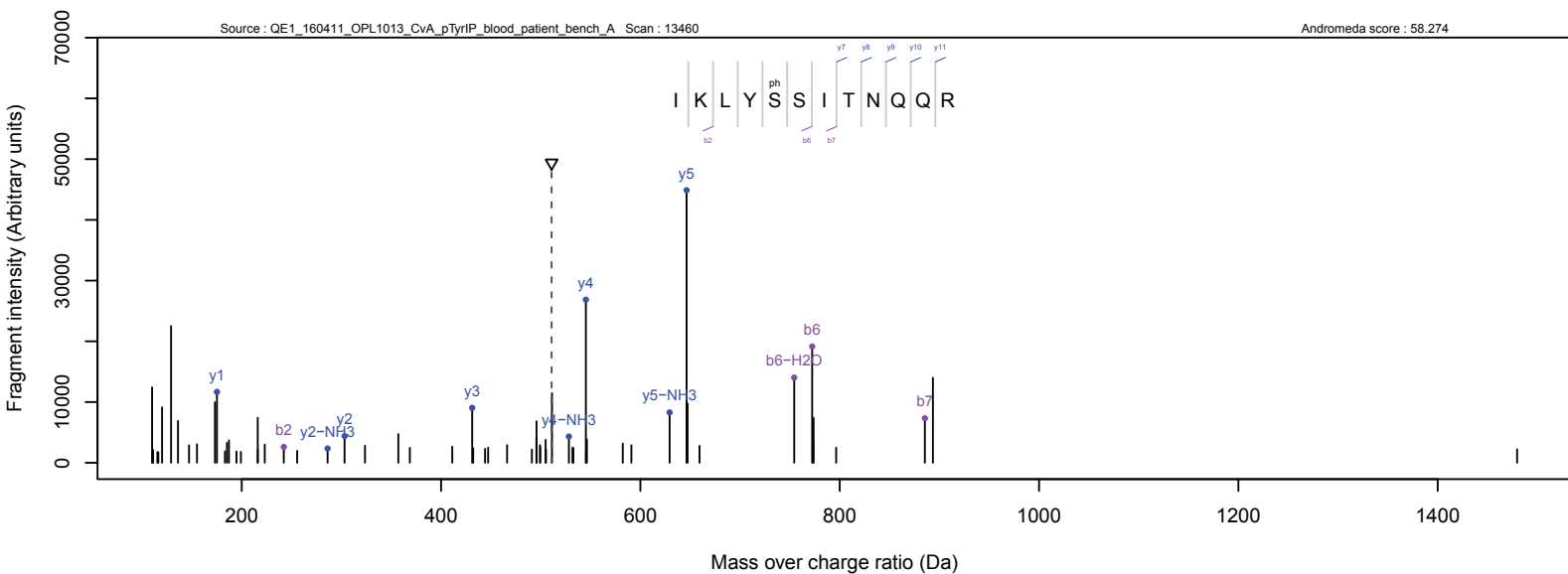
HVSISYDIPPTGNT(ph)YQIPR-3+



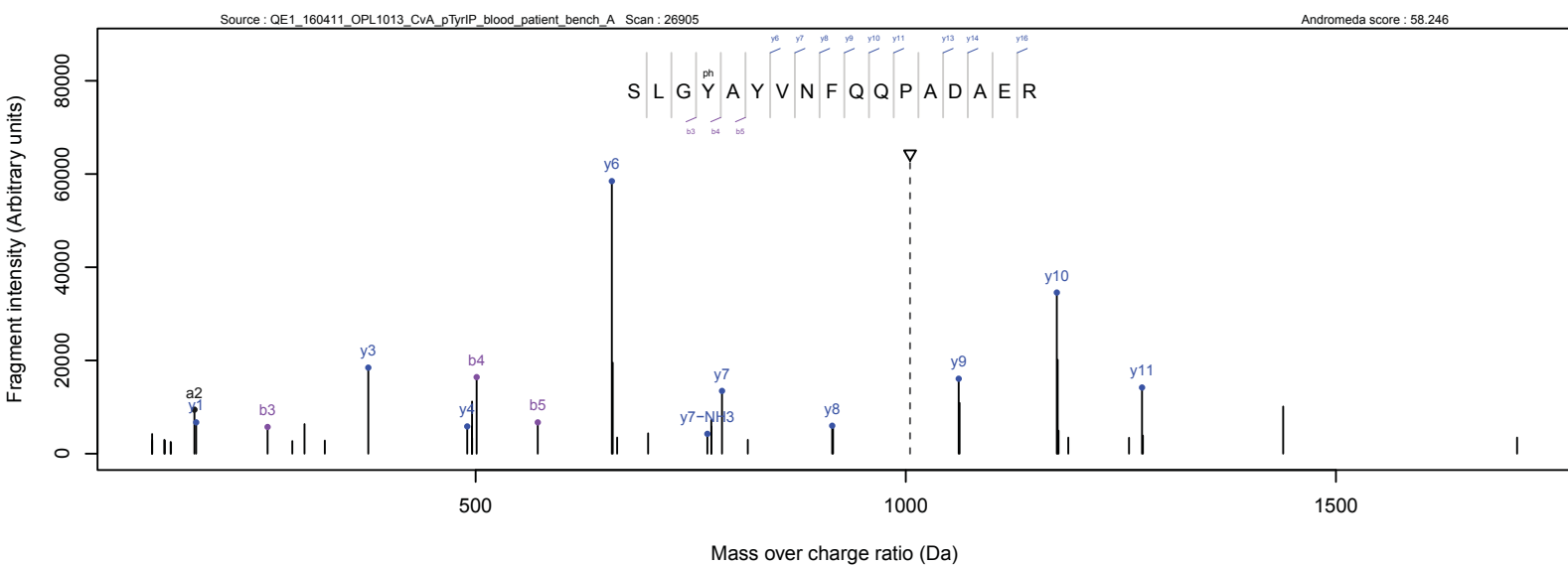
HADAEM(ox)TGY(ph)VVTR-3+



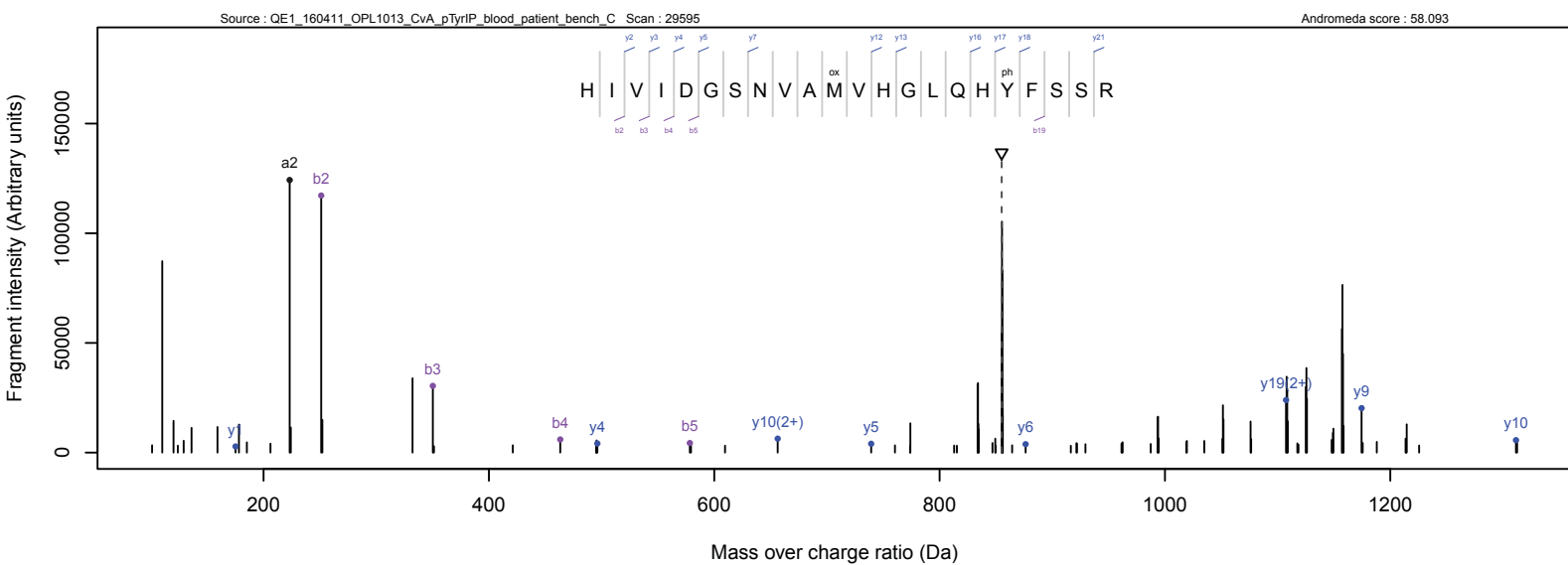
## IKLYS(ph)SITNQQR-3+

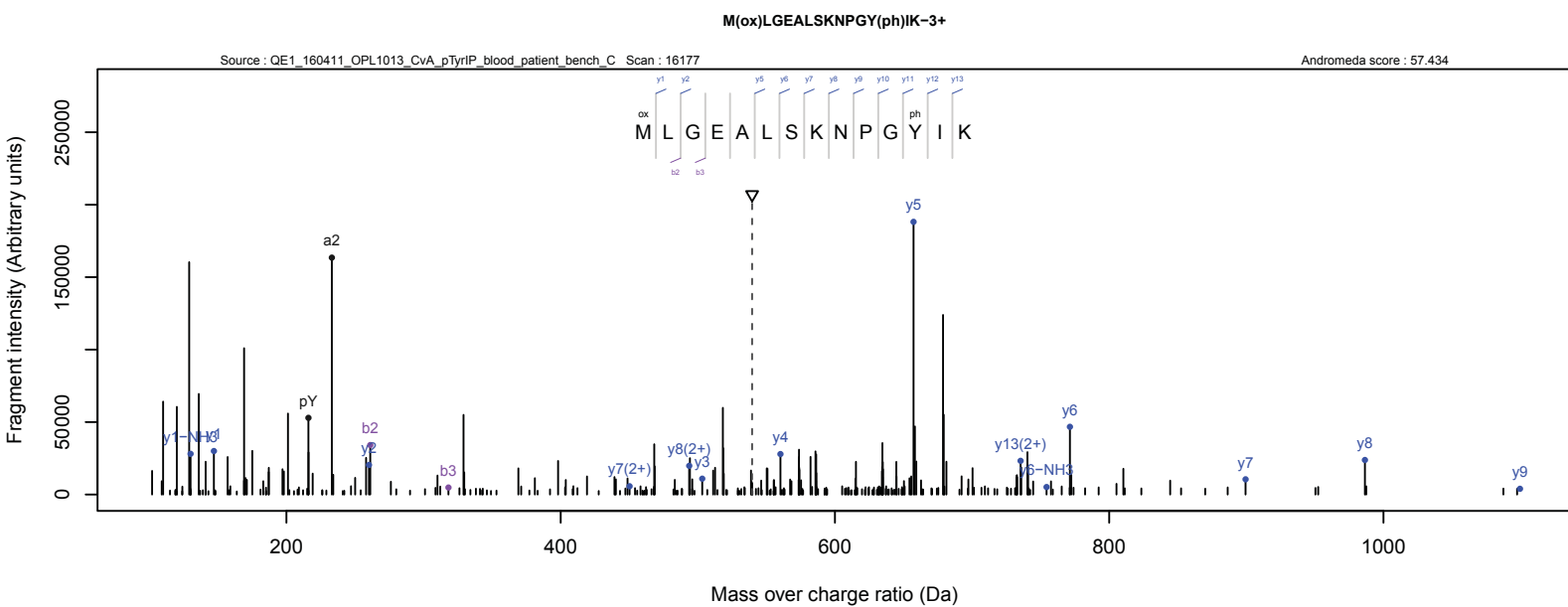
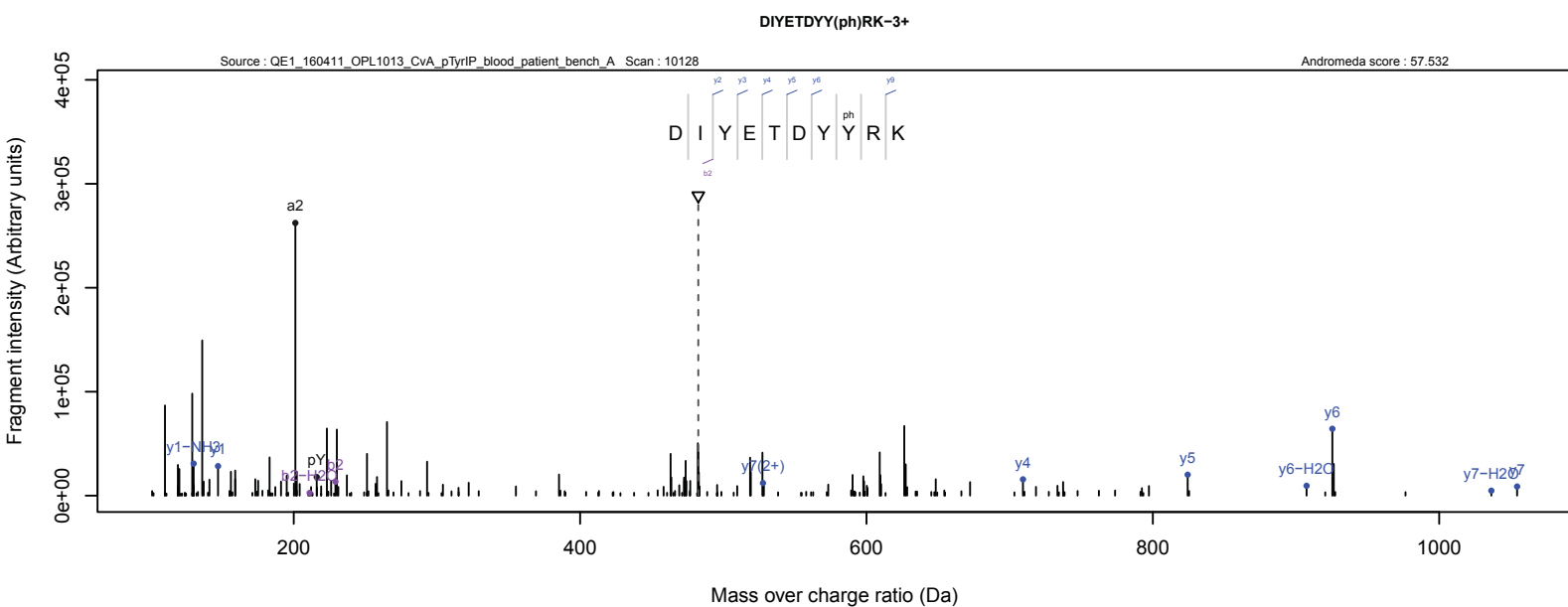
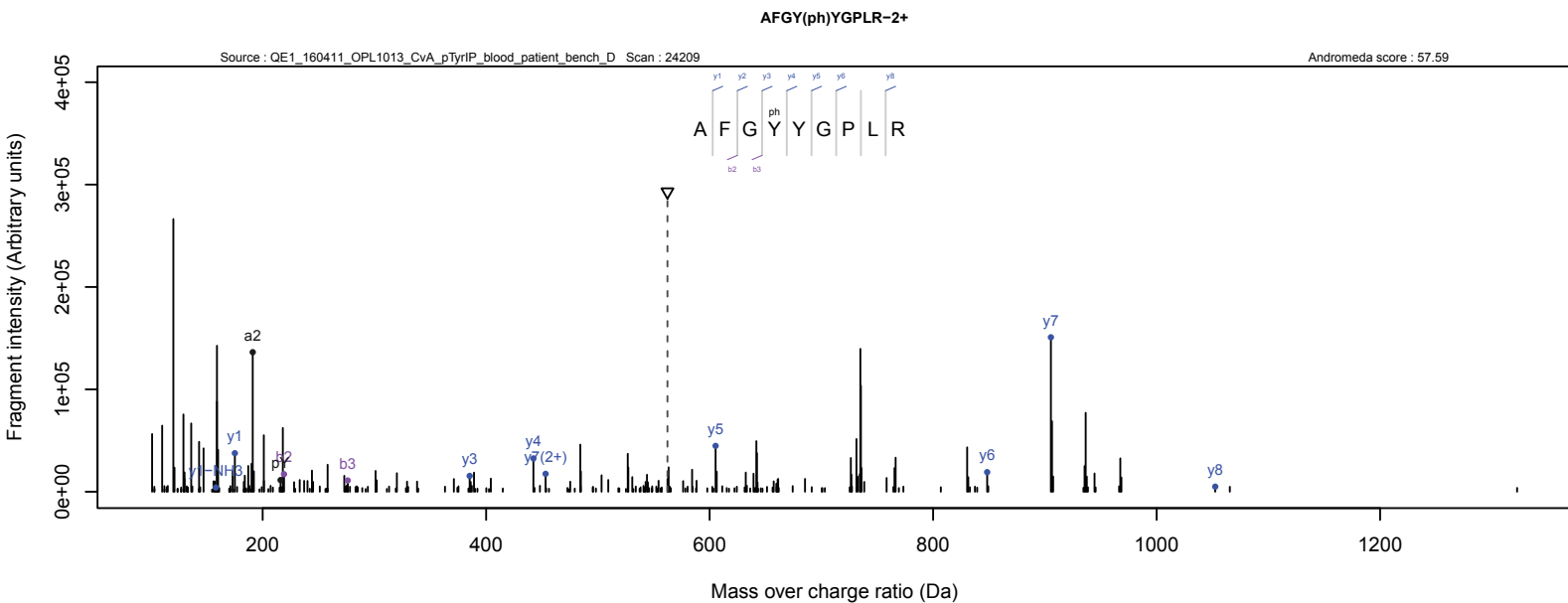


## SLGY(ph)AYVNFQQPADAER-2+



## HIVIDGSNVA(ox)VHGLQHY(ph)FSSR-3+

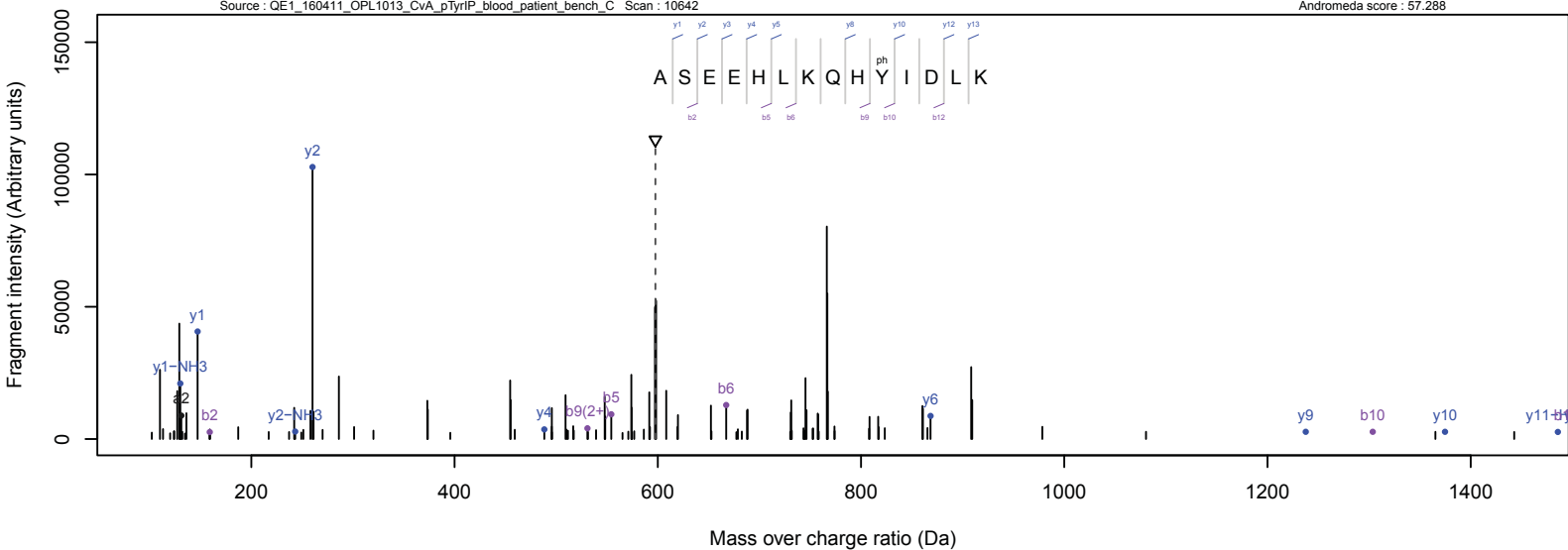




ASEEHLKQHY(ph)IDLK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 10642

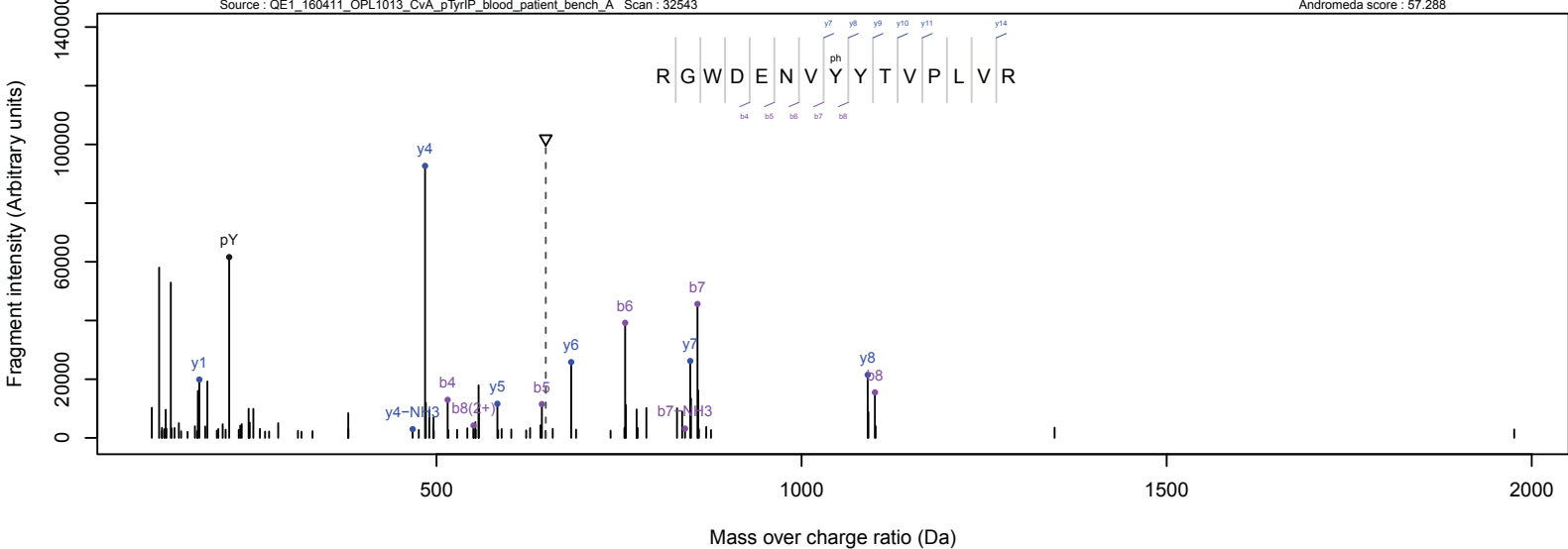
Andromeda score : 57.288



RGWDENVY(ph)YTVPLVR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 32543

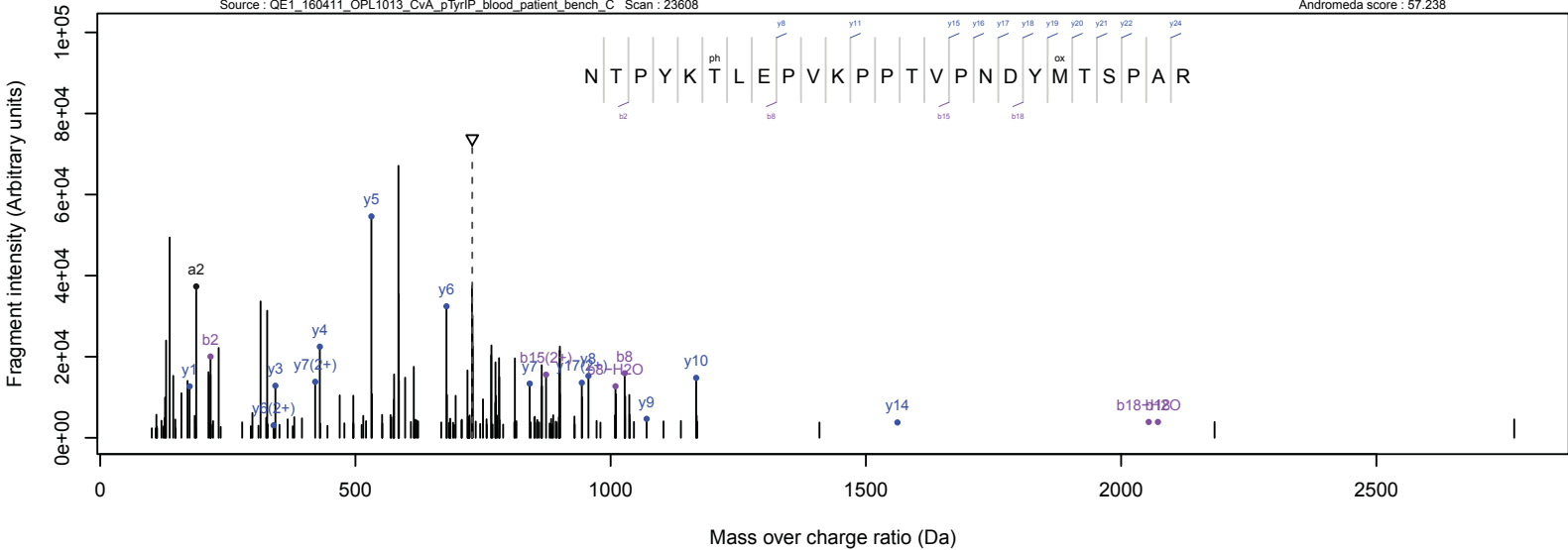
Andromeda score : 57.288



NTPYKT(ph)LEPVKPPPTVPNDYMTSPAR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 23608

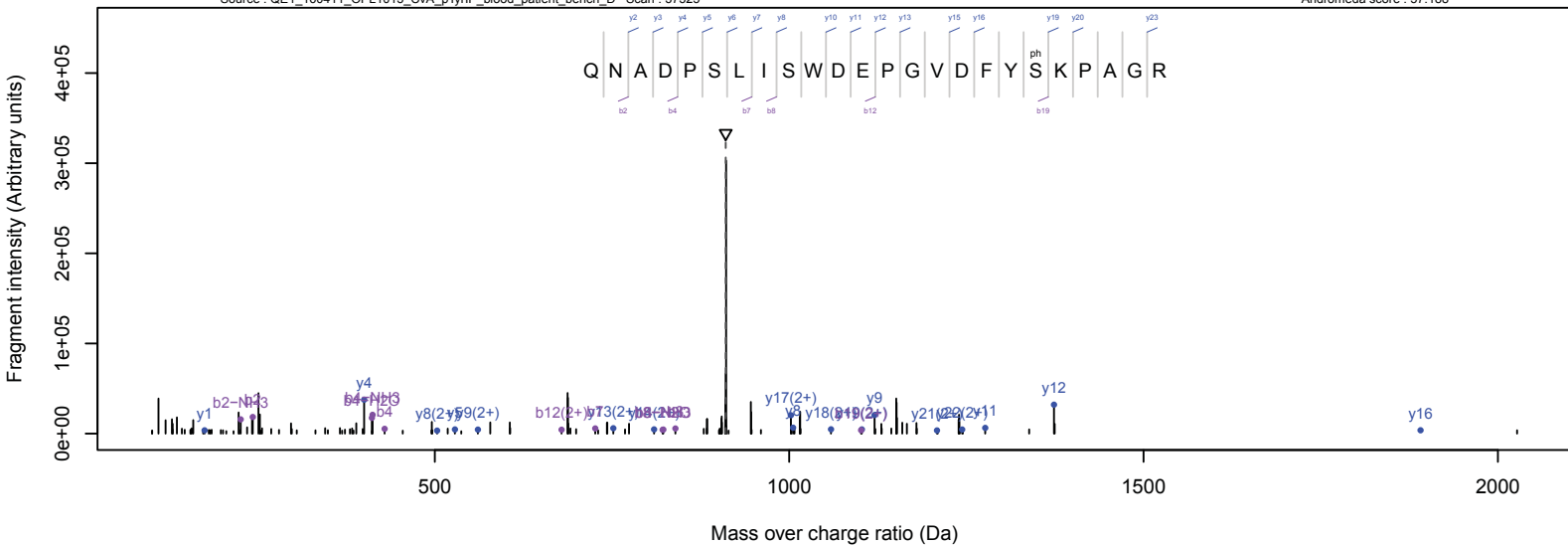
Andromeda score : 57.238



**QNADPSLISWDEPGLVDFY(ph)KPAIR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 37325

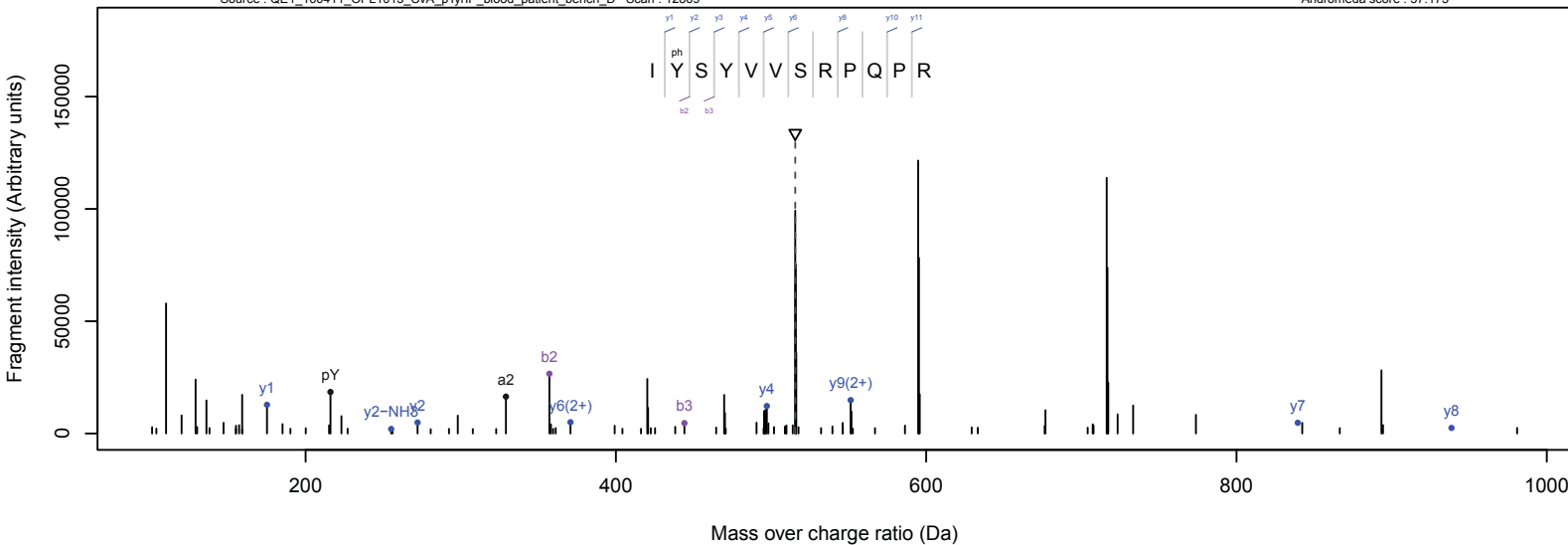
Andromeda score : 57.188



**IY(ph)SYVSRPQPR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12869

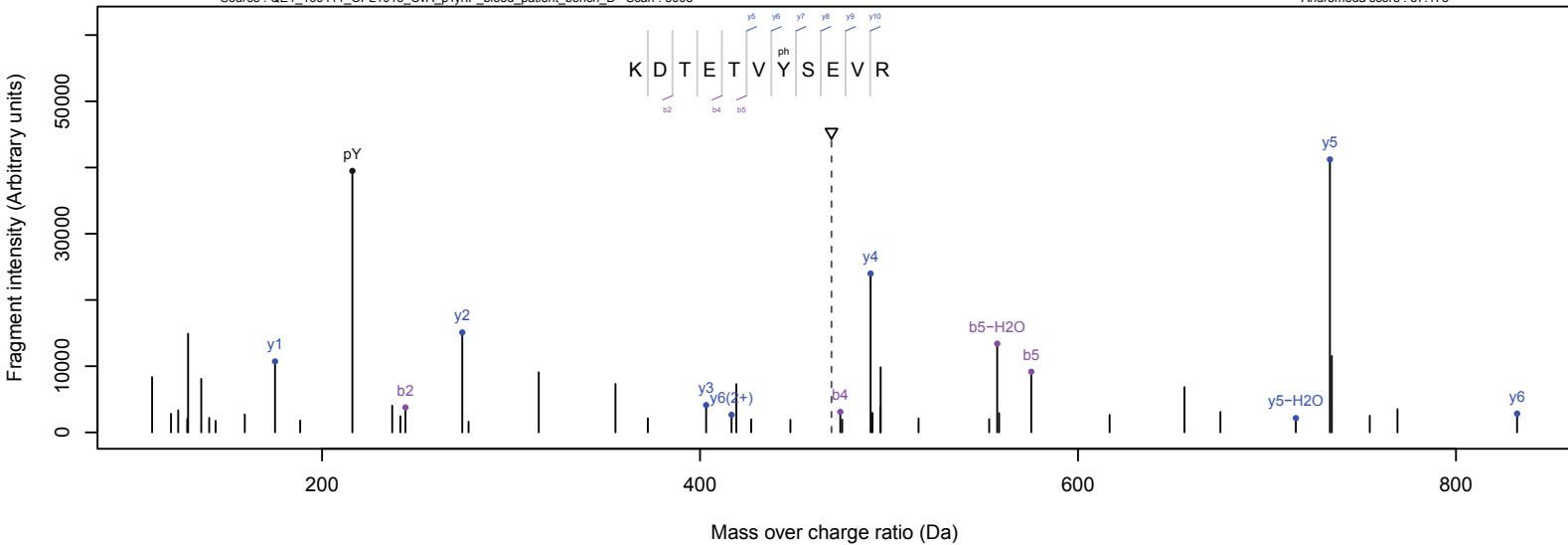
Andromeda score : 57.175



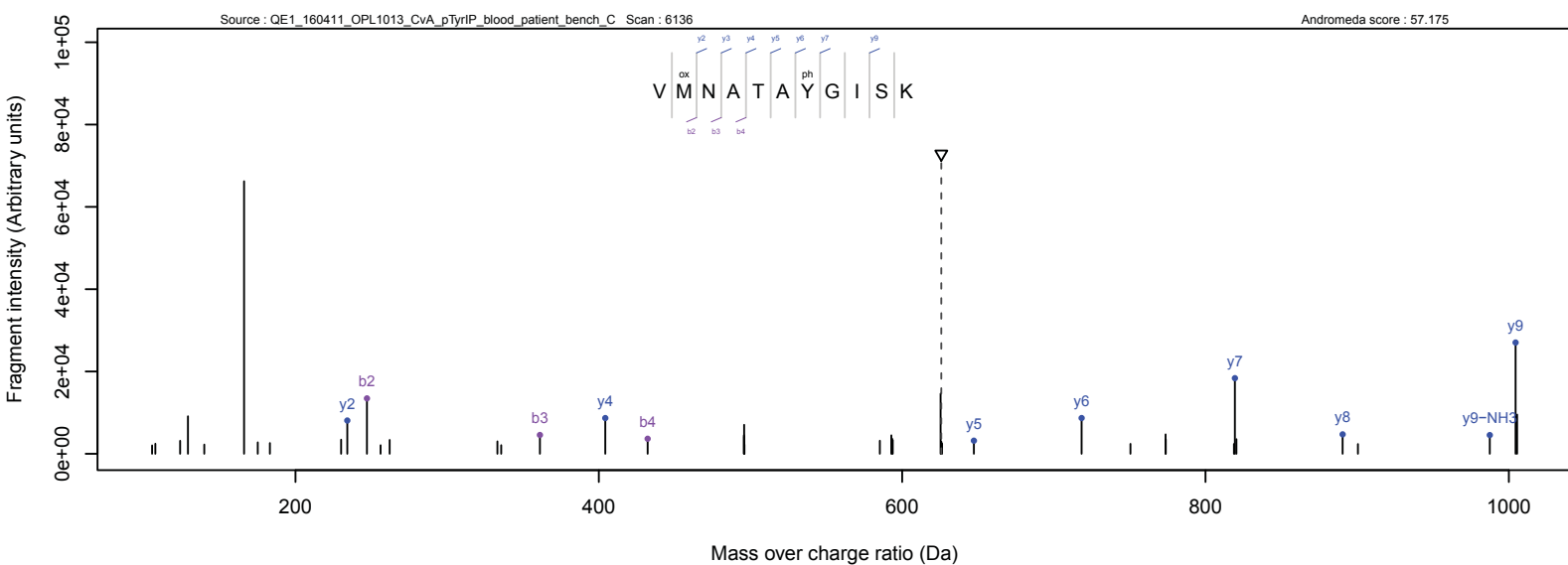
**KDETIVY(ph)SEVR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5008

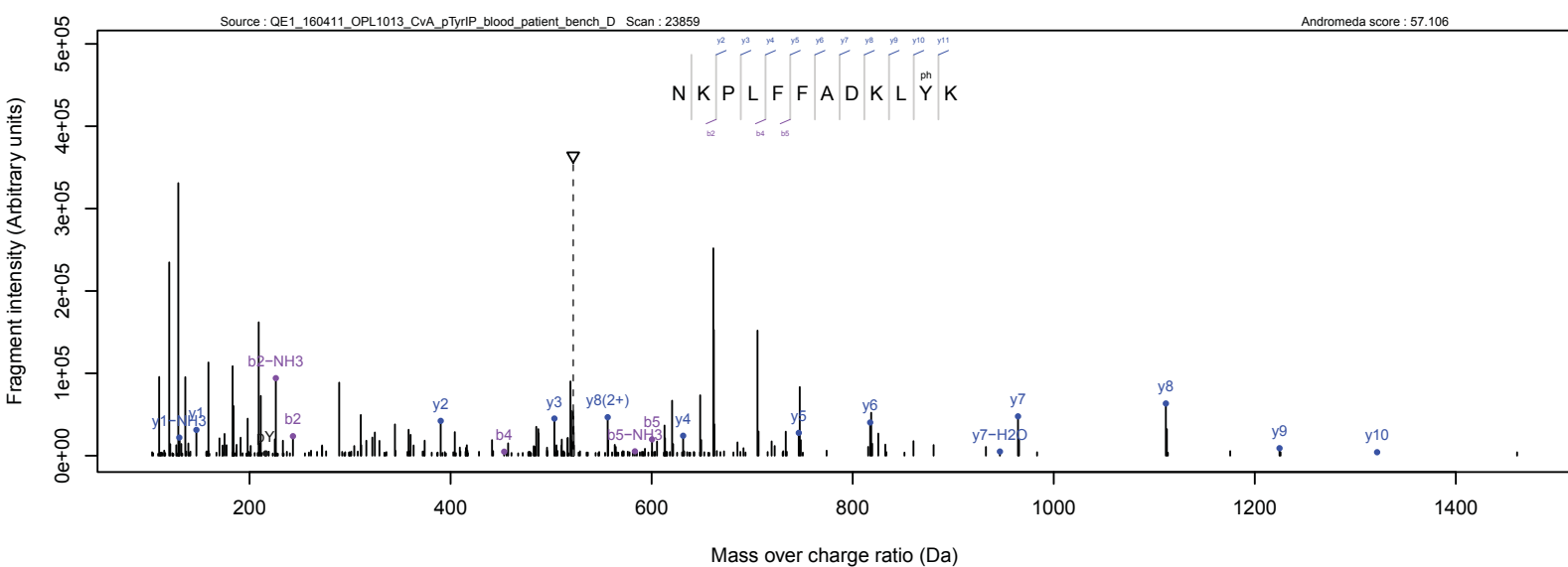
Andromeda score : 57.175



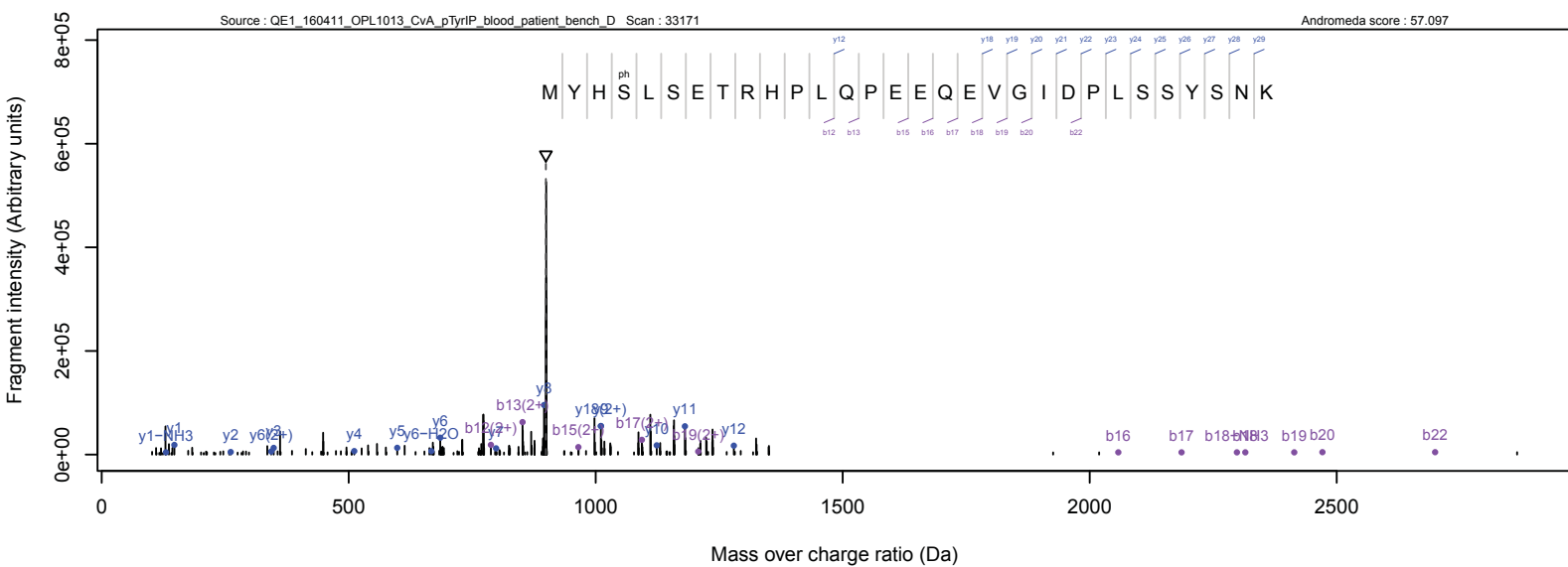
## VM(ox)NATAY(ph)GISK-2+



## NKPLFFADKLYK-3+



## (ac)MYHS(ph)LSETRHPLQPEEQEVGIDPLSSYSNK-4+



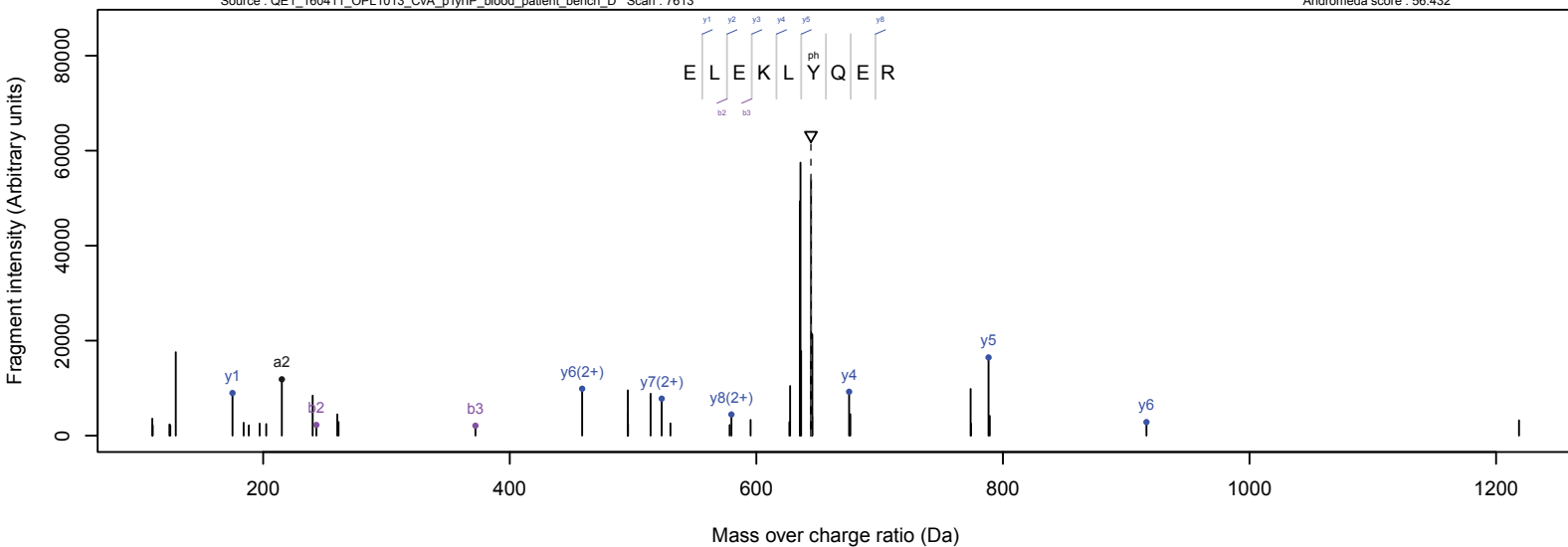




## ELEKLY(ph)QER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7613

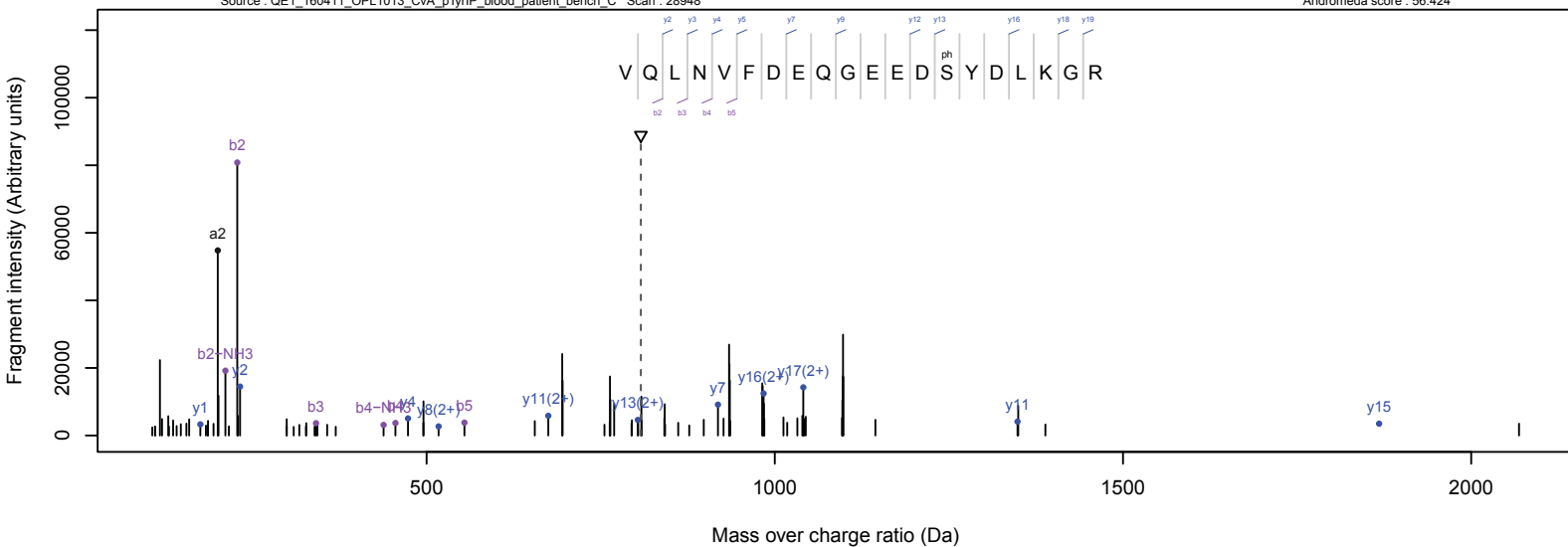
Andromeda score : 56.432



## VQLNVFDEQGEEDS(ph)YDLKGR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28948

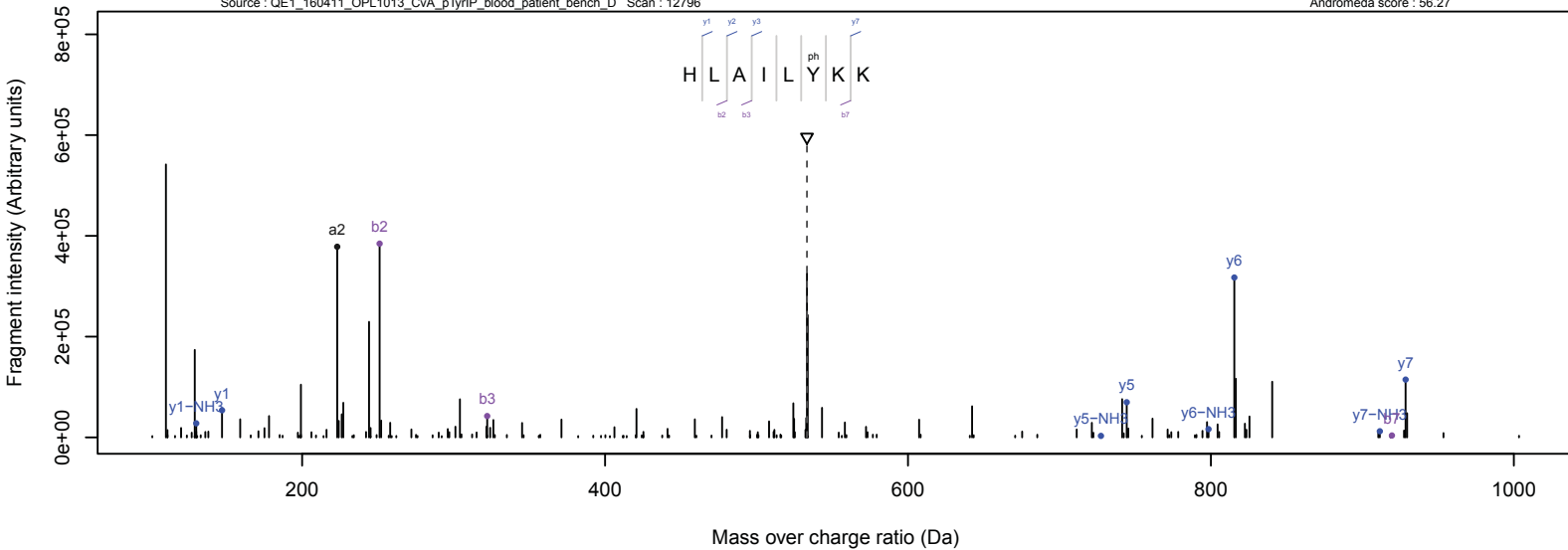
Andromeda score : 56.424

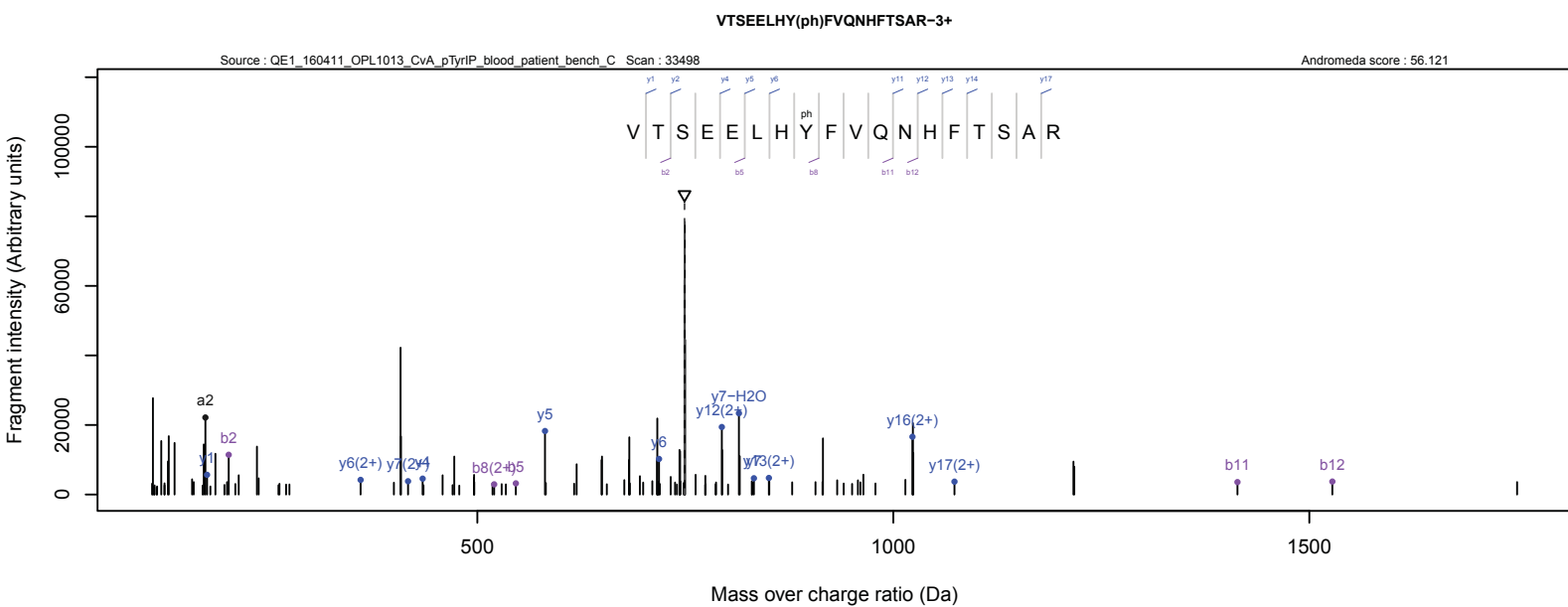
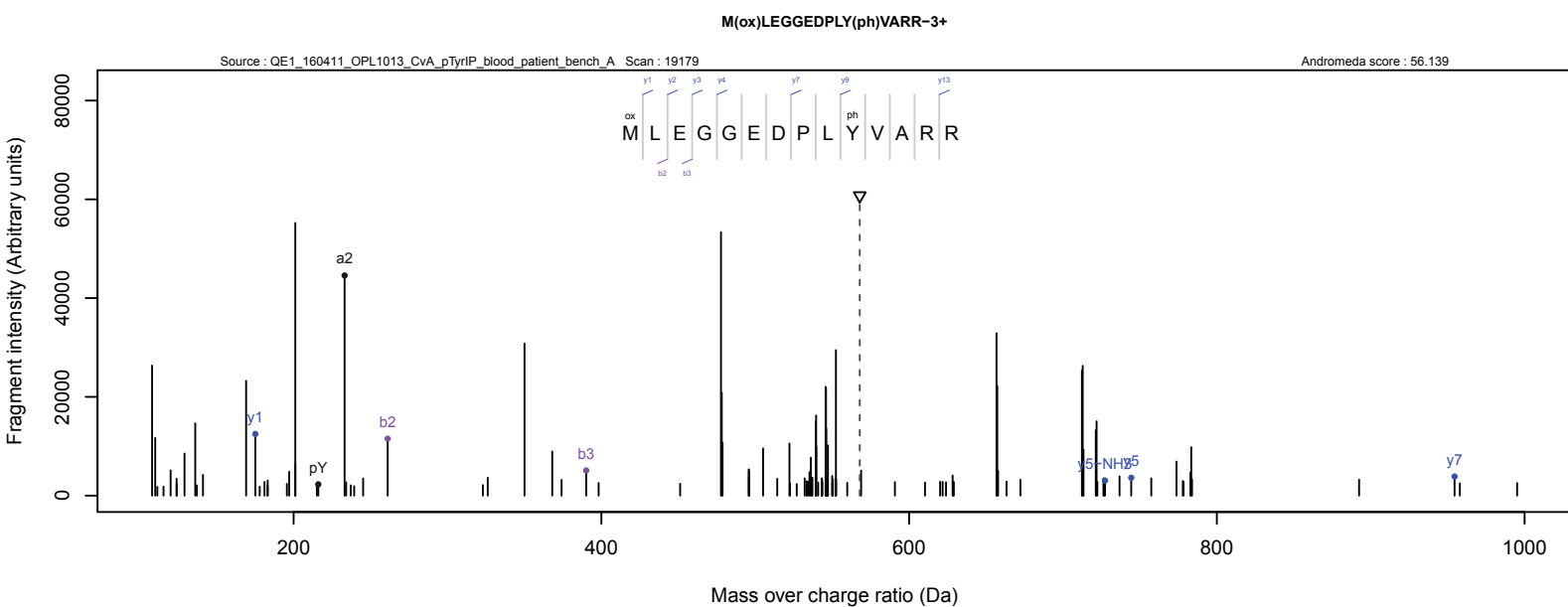
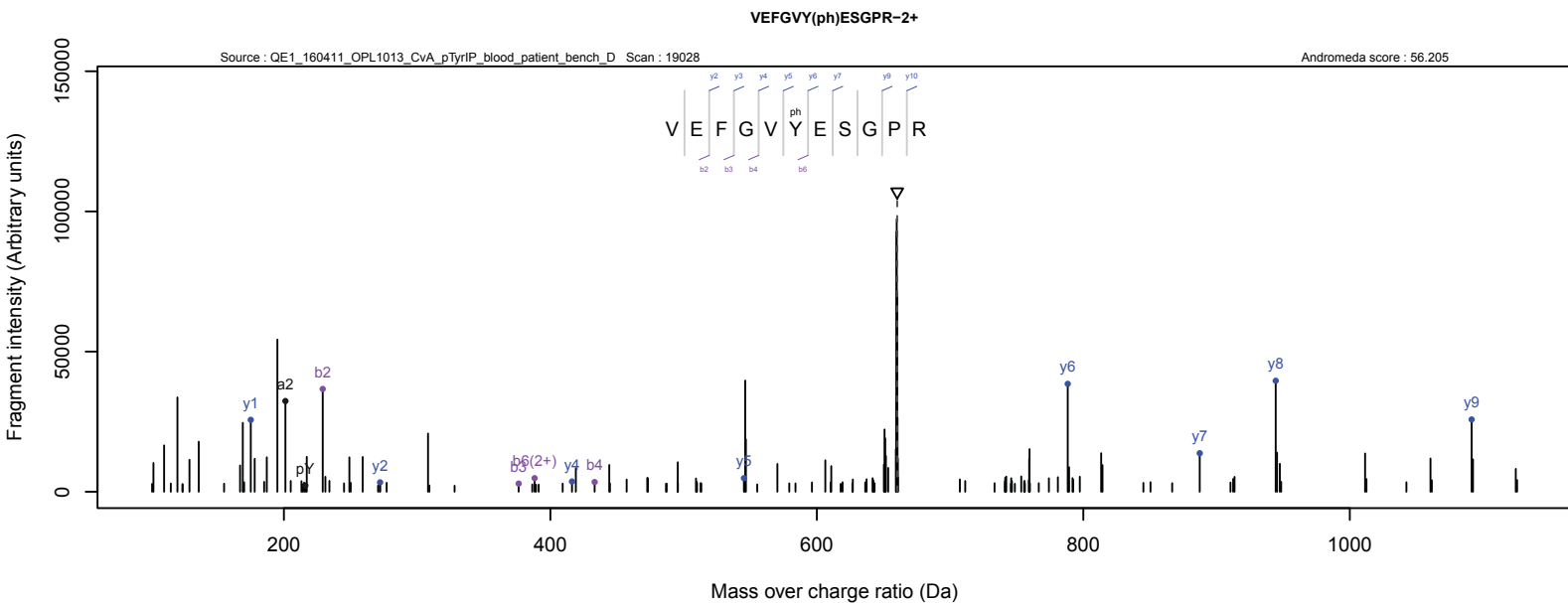


## HLAILY(ph)KK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12796

Andromeda score : 56.27

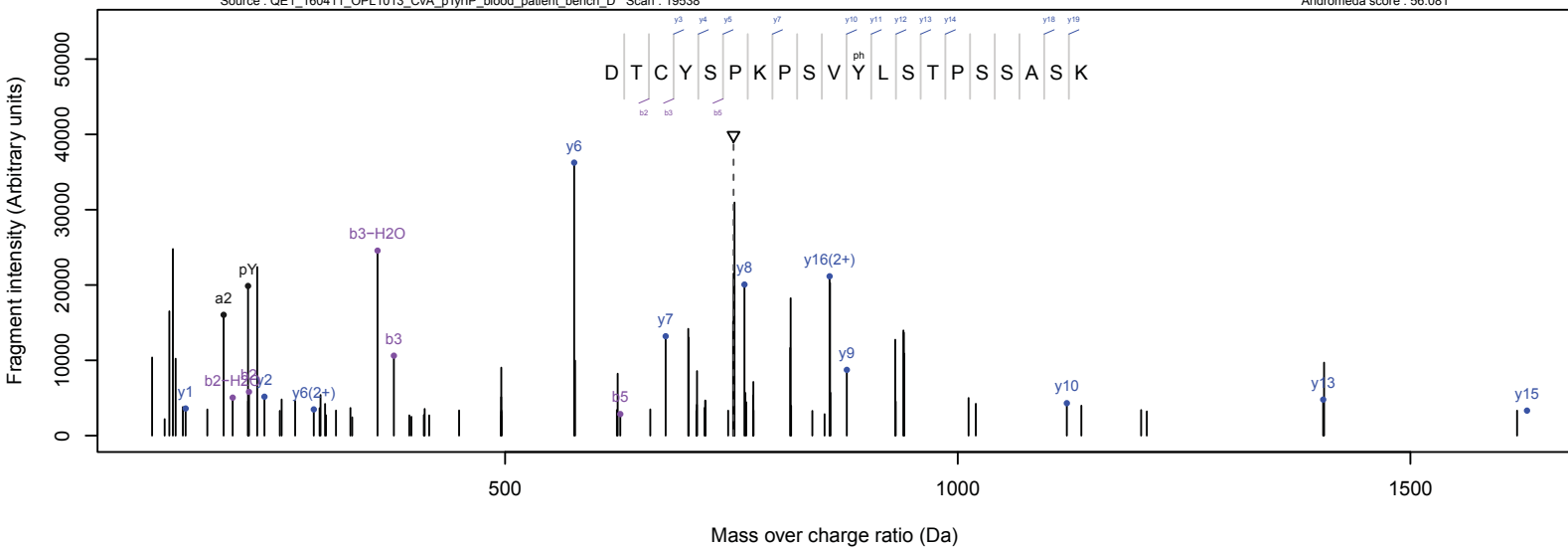




## DTCYSPKPSVY(ph)LSTPSSASK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 19538

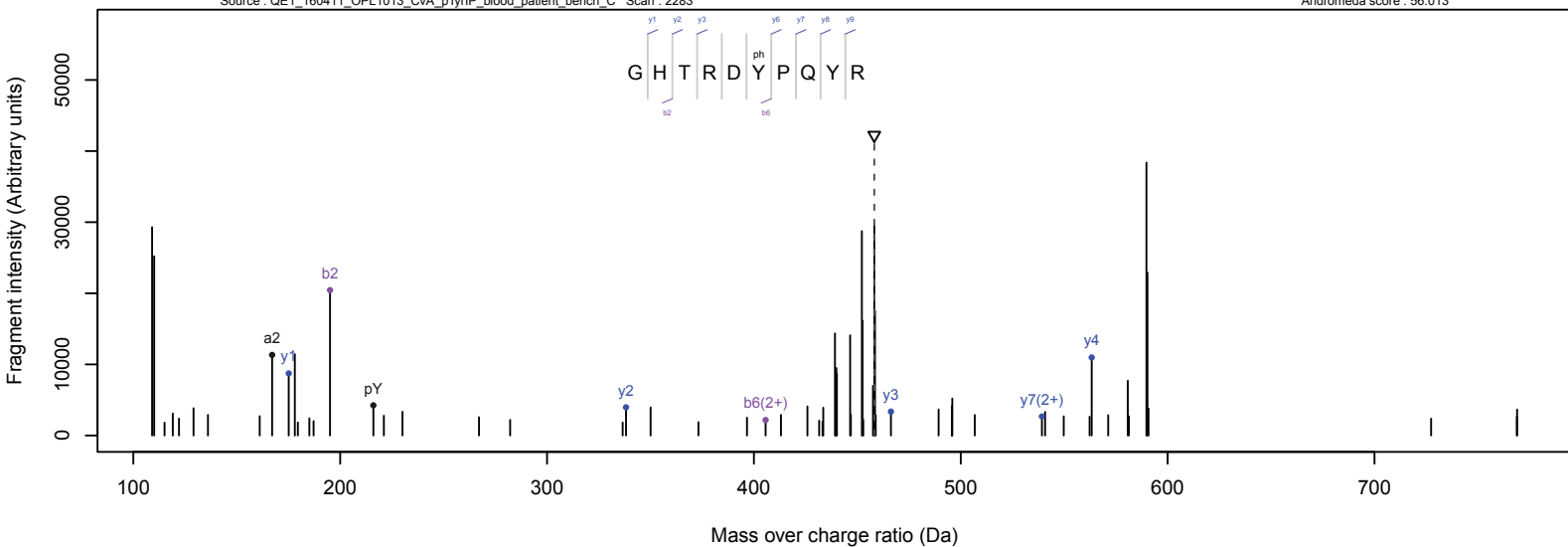
Andromeda score : 56.081



## GHTRDY(ph)PQYR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 2283

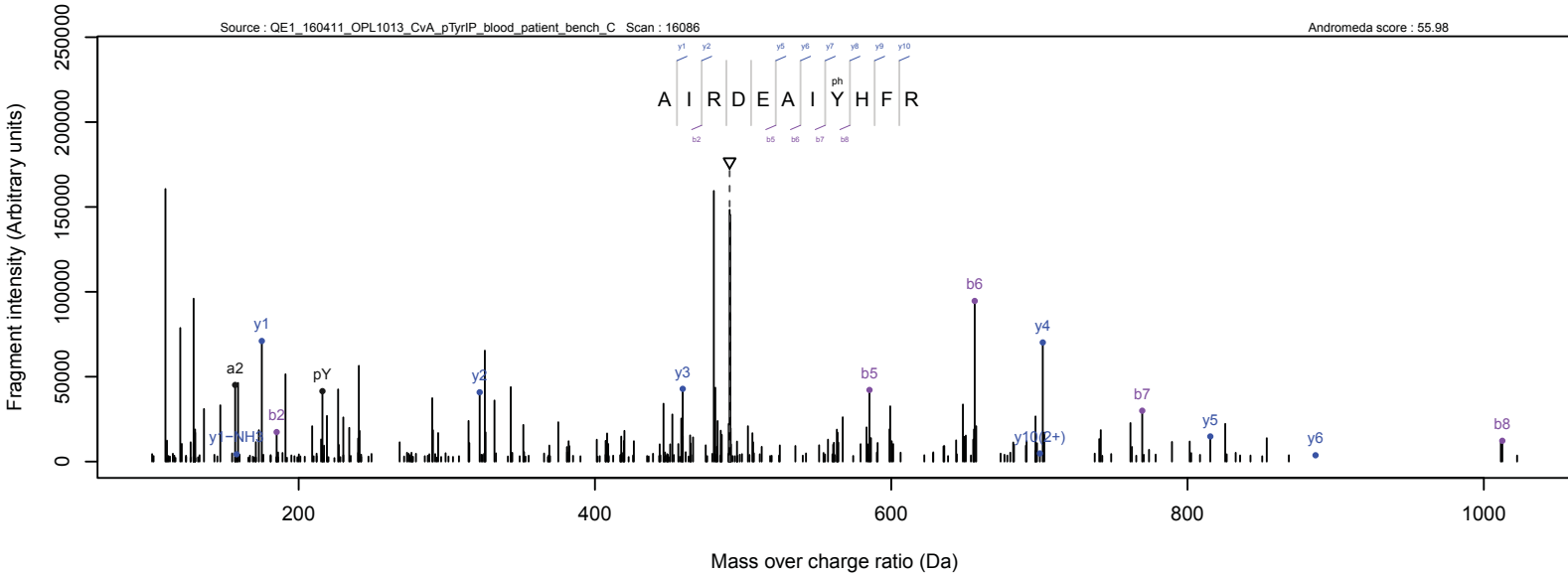
Andromeda score : 56.013



## AIRDEAIY(ph)HFR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16086

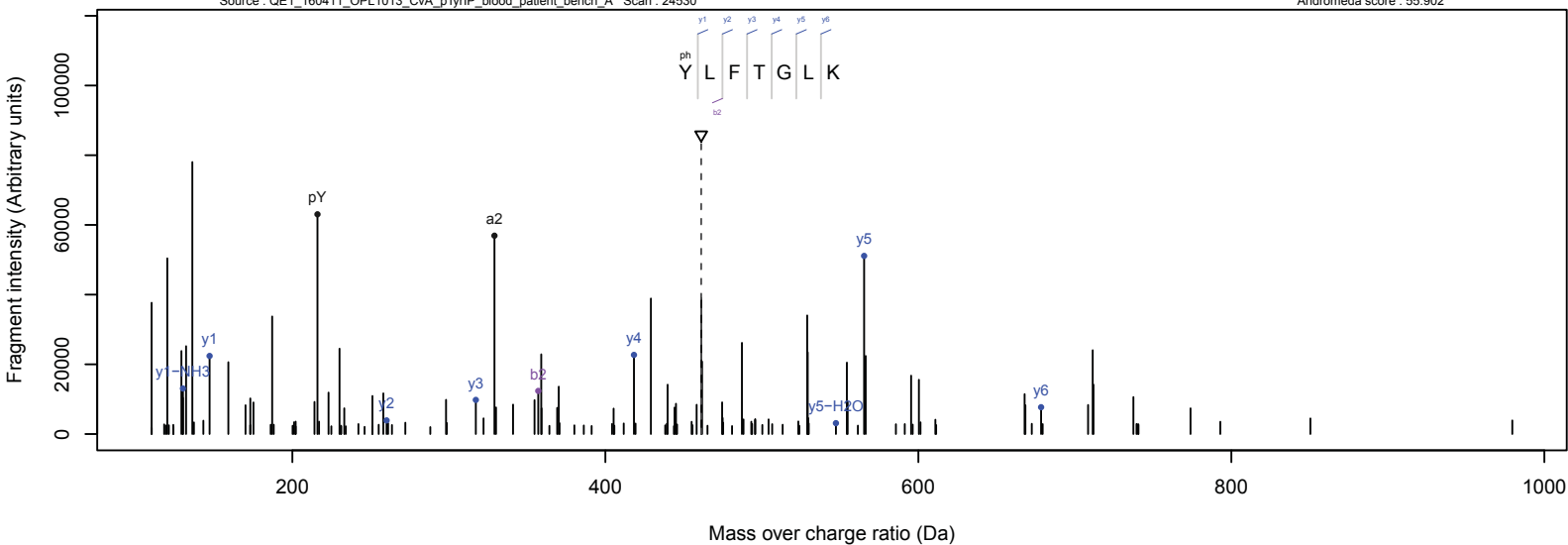
Andromeda score : 55.98



## Y(ph)LFTGLK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 24530

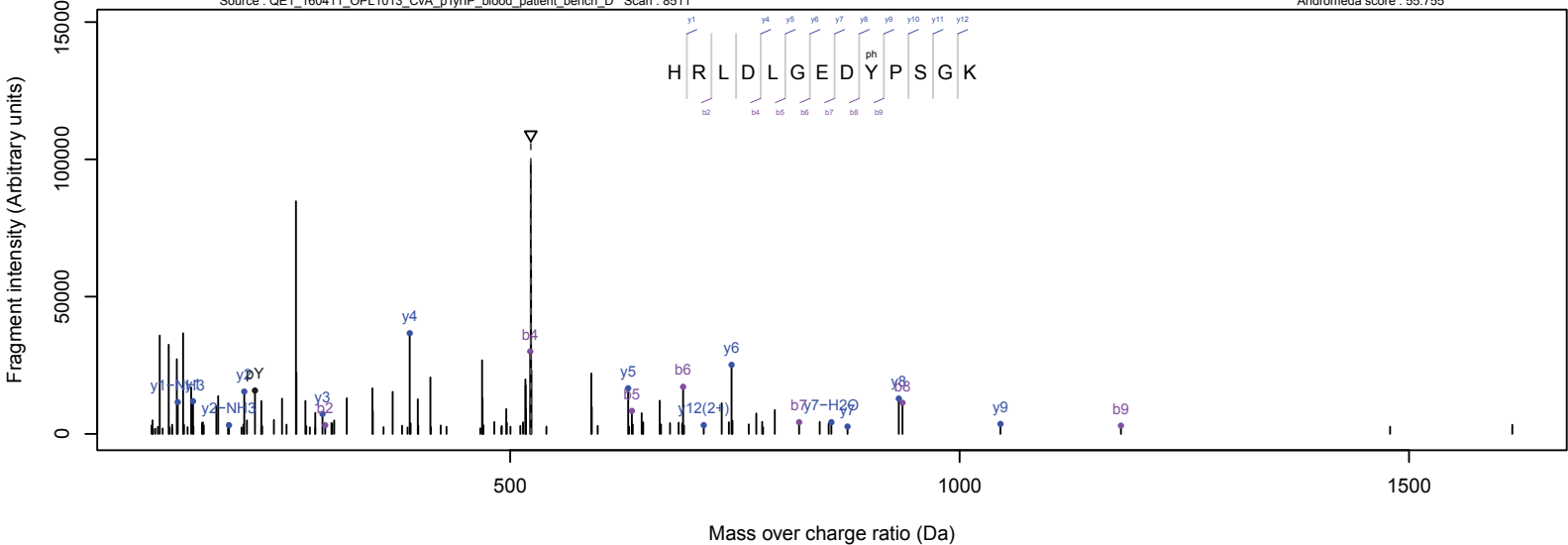
Andromeda score: 55.902



## HRLDLGEDY(ph)PSGK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 8511

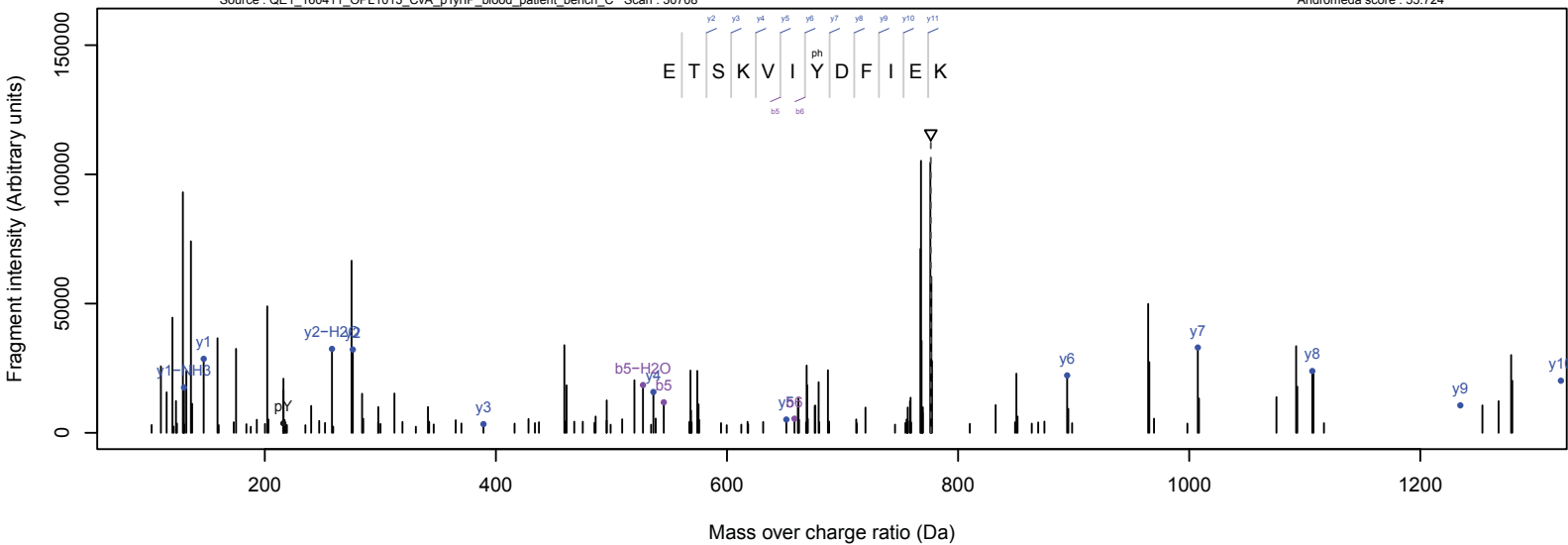
Andromeda score: 55.755



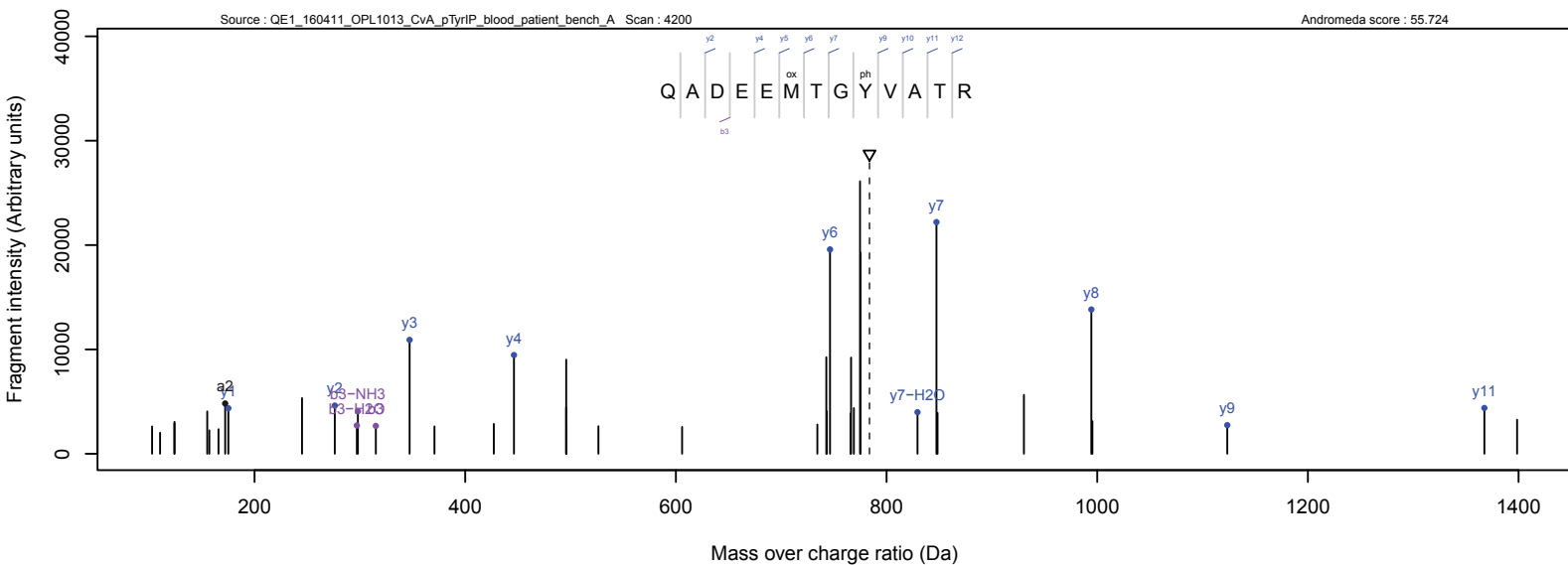
## ETSKVIY(ph)DFIEK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 30708

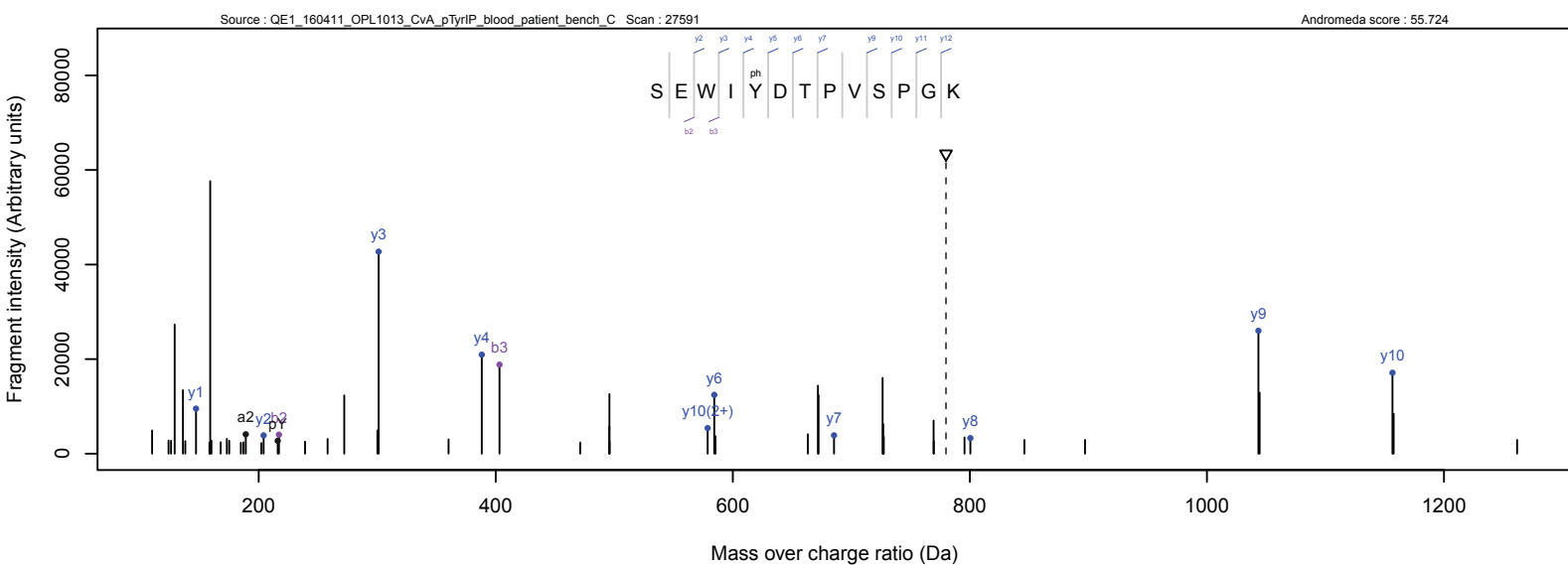
Andromeda score: 55.724



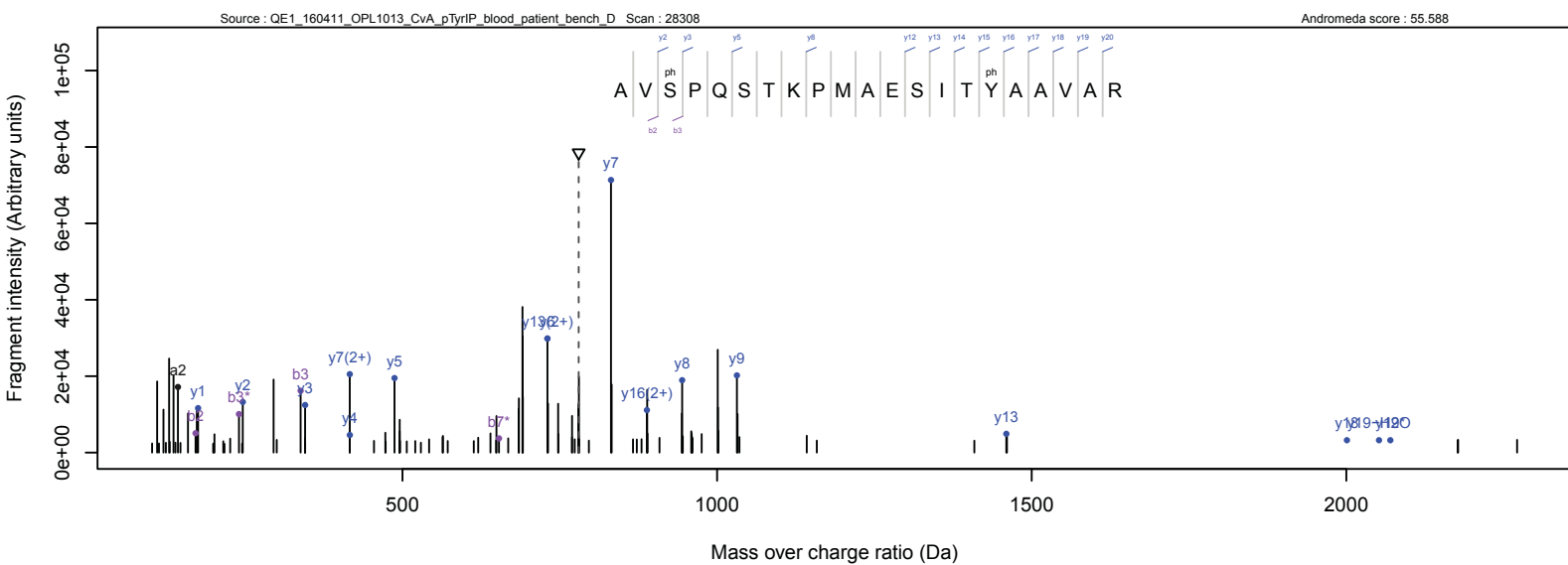
**QADEEM(ox)TGY(ph)VATR-2+**



**SEWIY(ph)DTPVSPGK-2+**



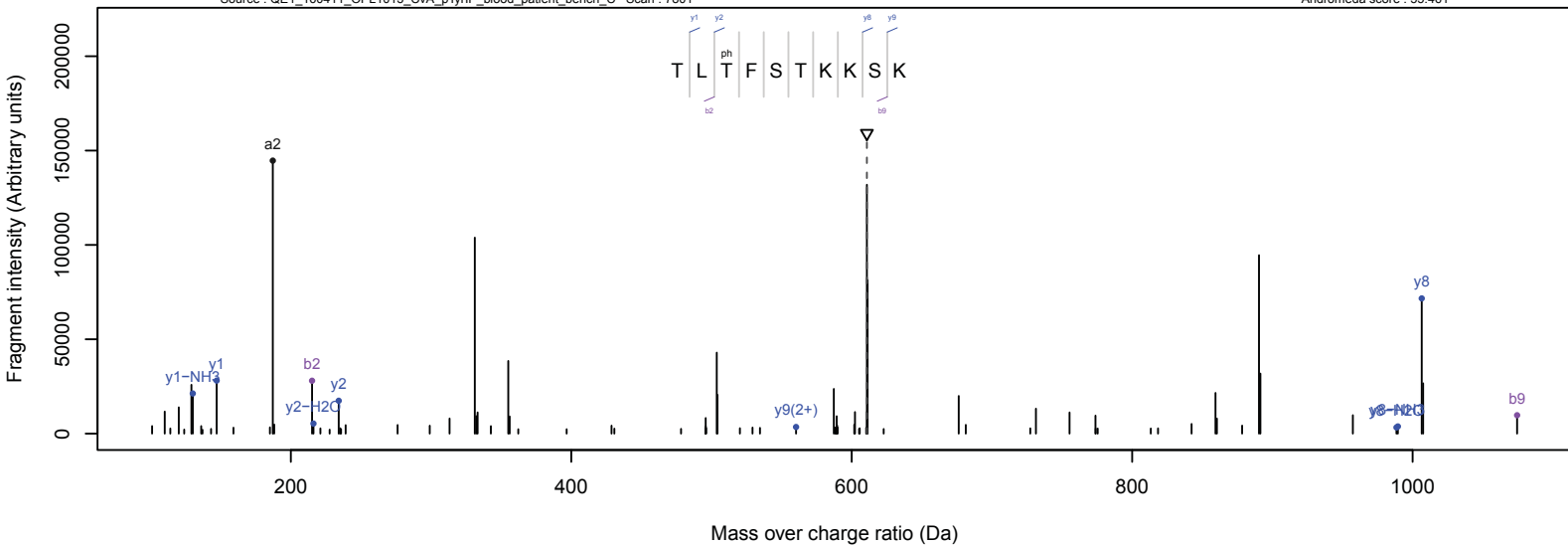
**AVS(ph)PQSTKPMAESIT(ph)YAAVAR-3+**



## TLT(ph)FSTKKS-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 7801

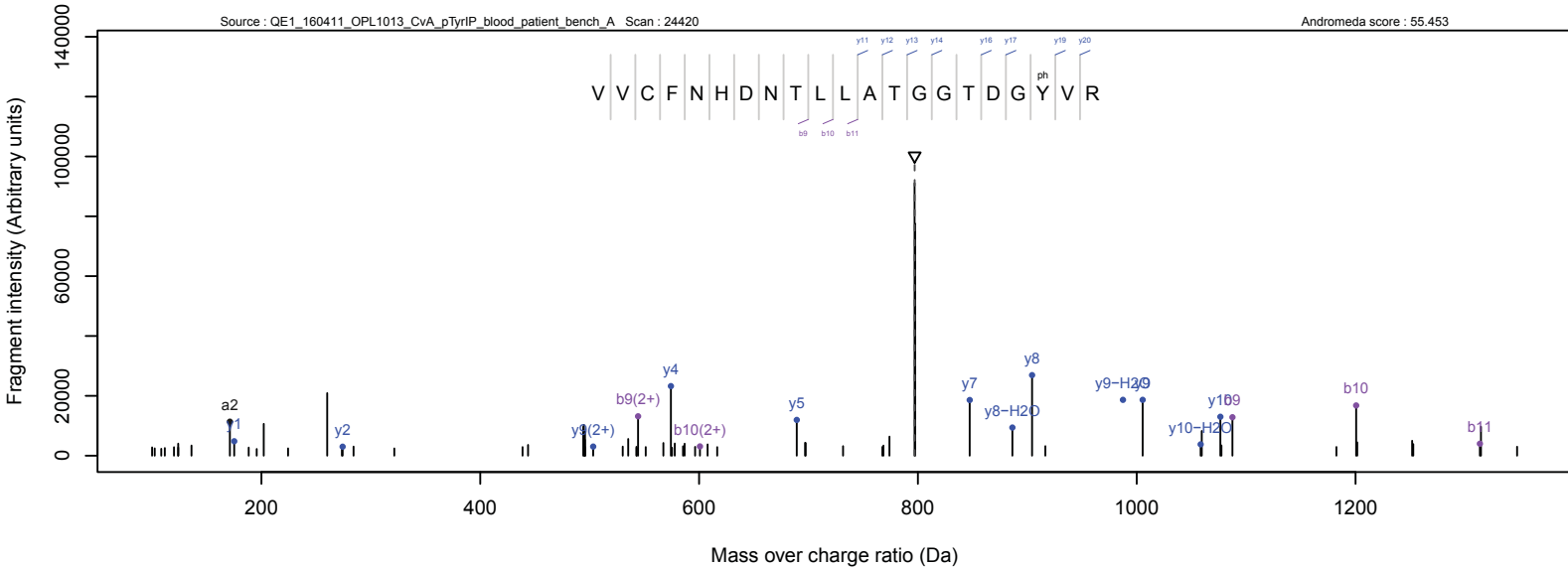
Andromeda score: 55.461



## VVCFNHDNTLLATGGTDGY(ph)VR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 24420

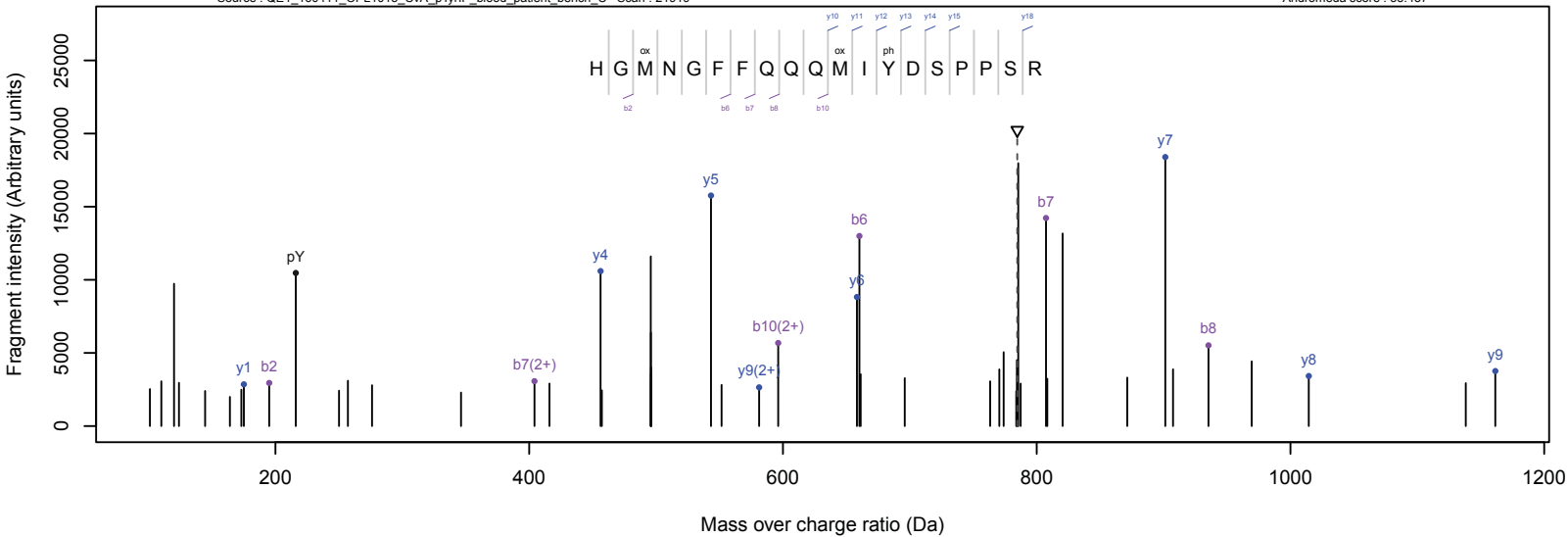
Andromeda score: 55.453



## HGM(ox)NGFFQQM(ox)IY(ph)DSPPSR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 21919

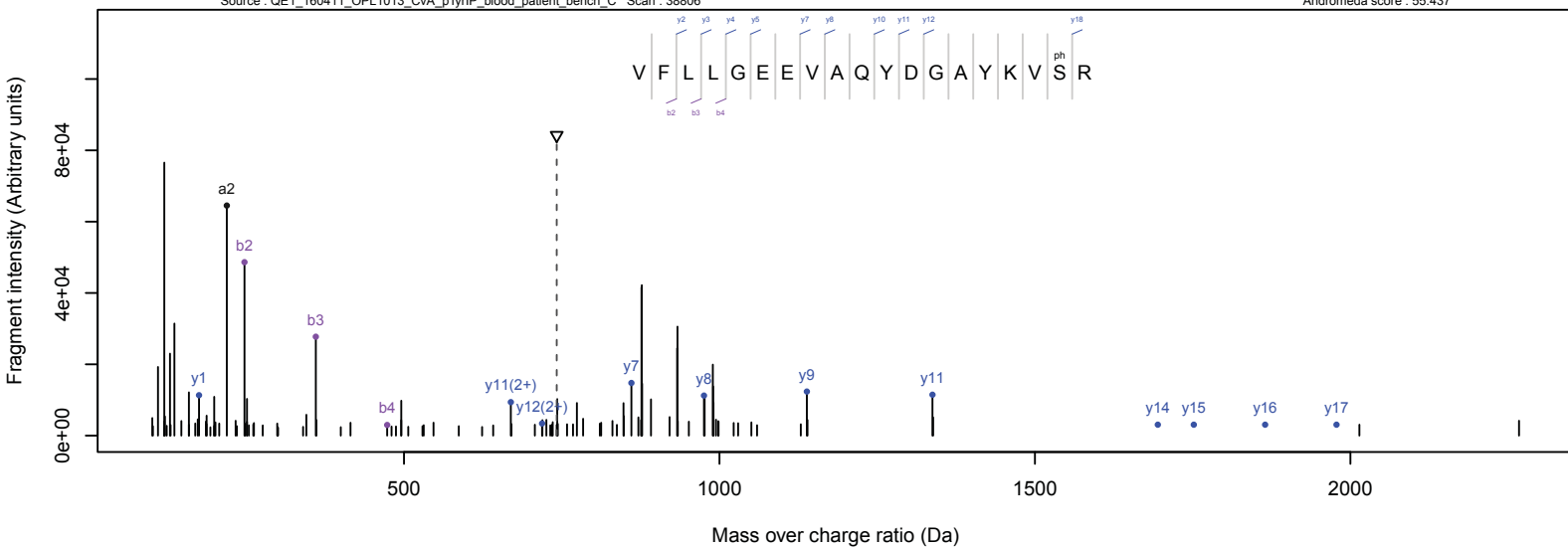
Andromeda score: 55.437



VLLGEEVAQYDGAYKVS(ph)R-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 38806

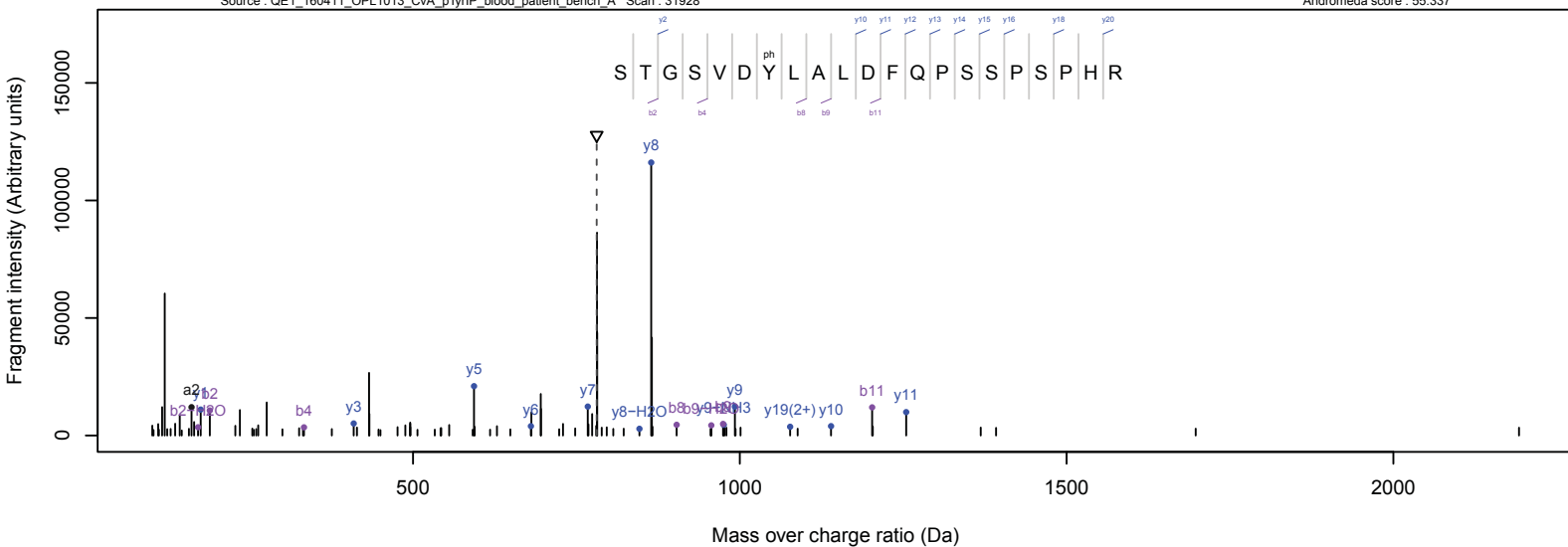
Andromeda score : 55.437



STGSVDY(ph)LALDFQPSSPSHR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 31928

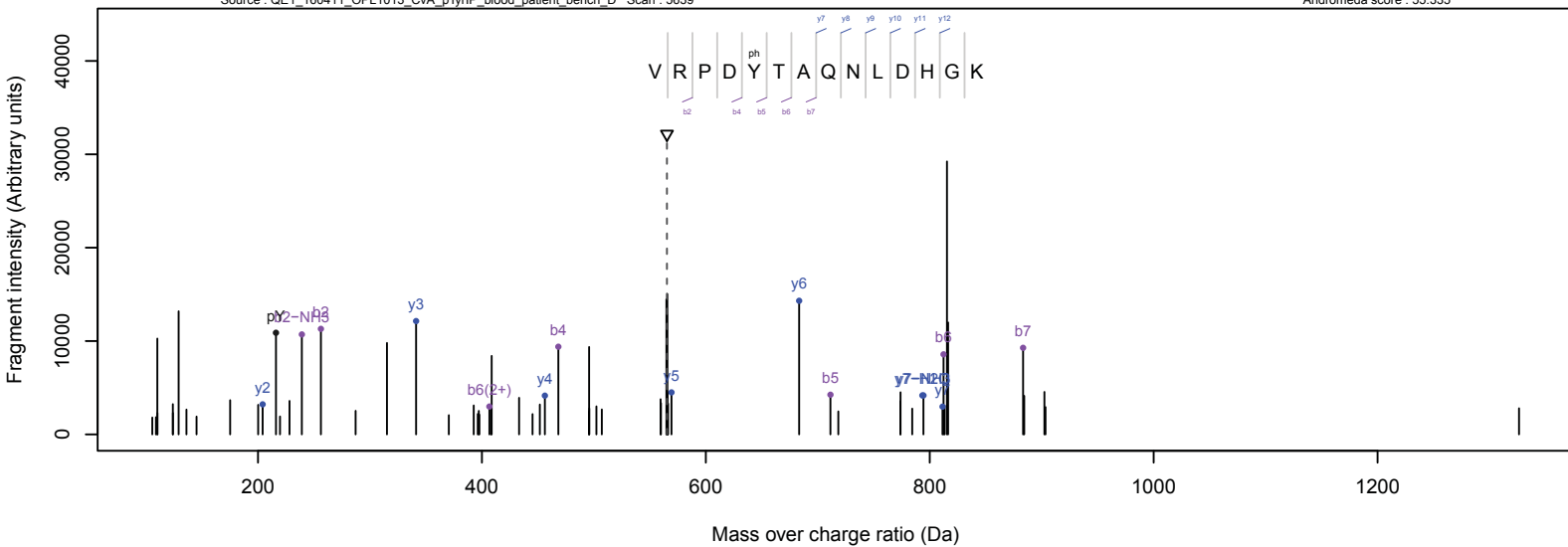
Andromeda score : 55.337



VRPDY(ph)TAQNLDHGK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 5639

Andromeda score : 55.335

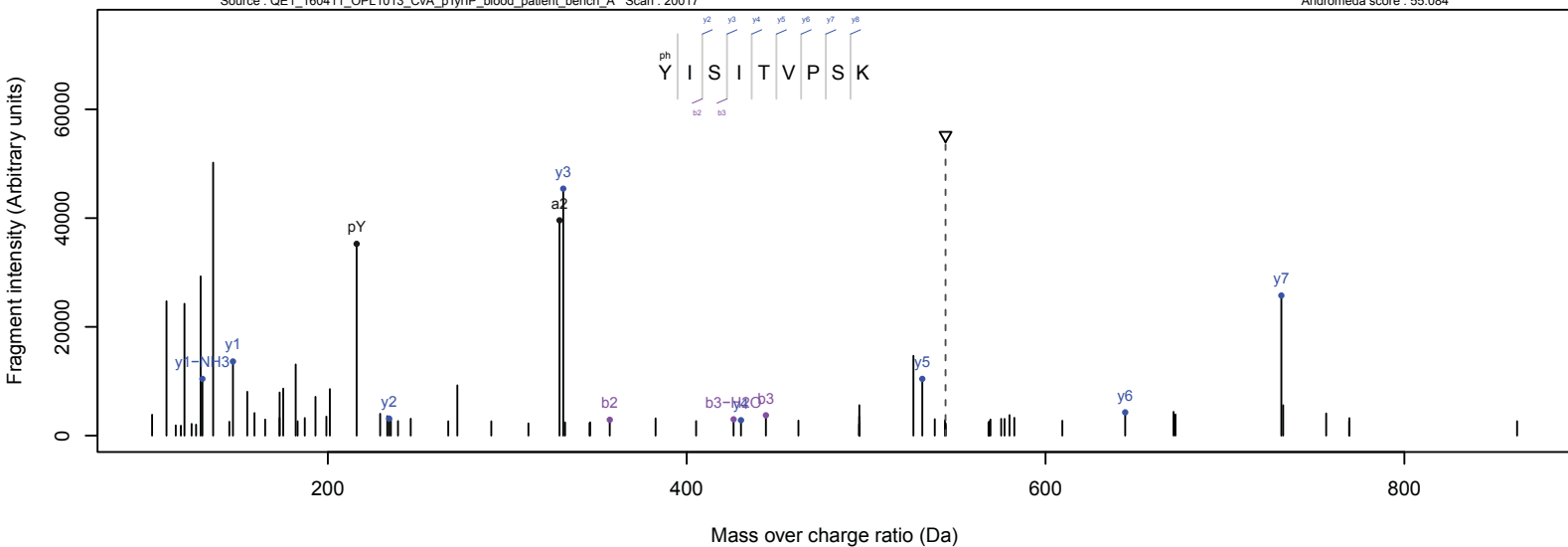




Y(ph)ISITVPSK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 20017

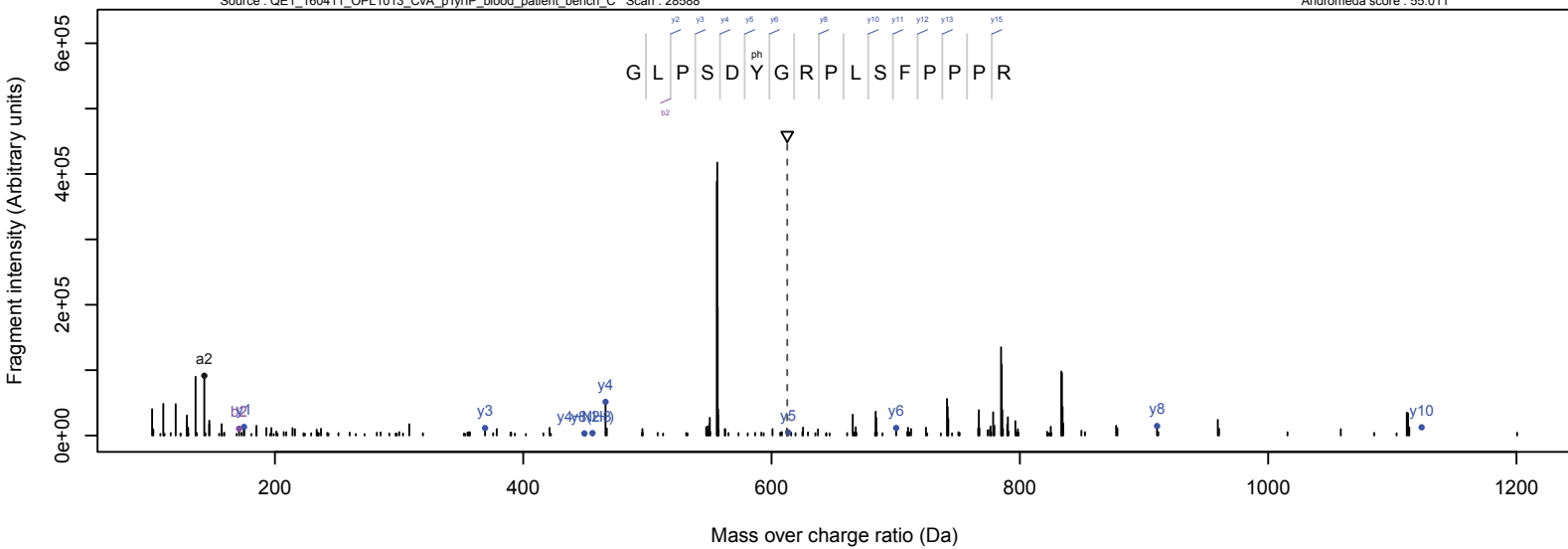
Andromeda score : 55.084



GLPSDY(ph)GRPLSFPPR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28588

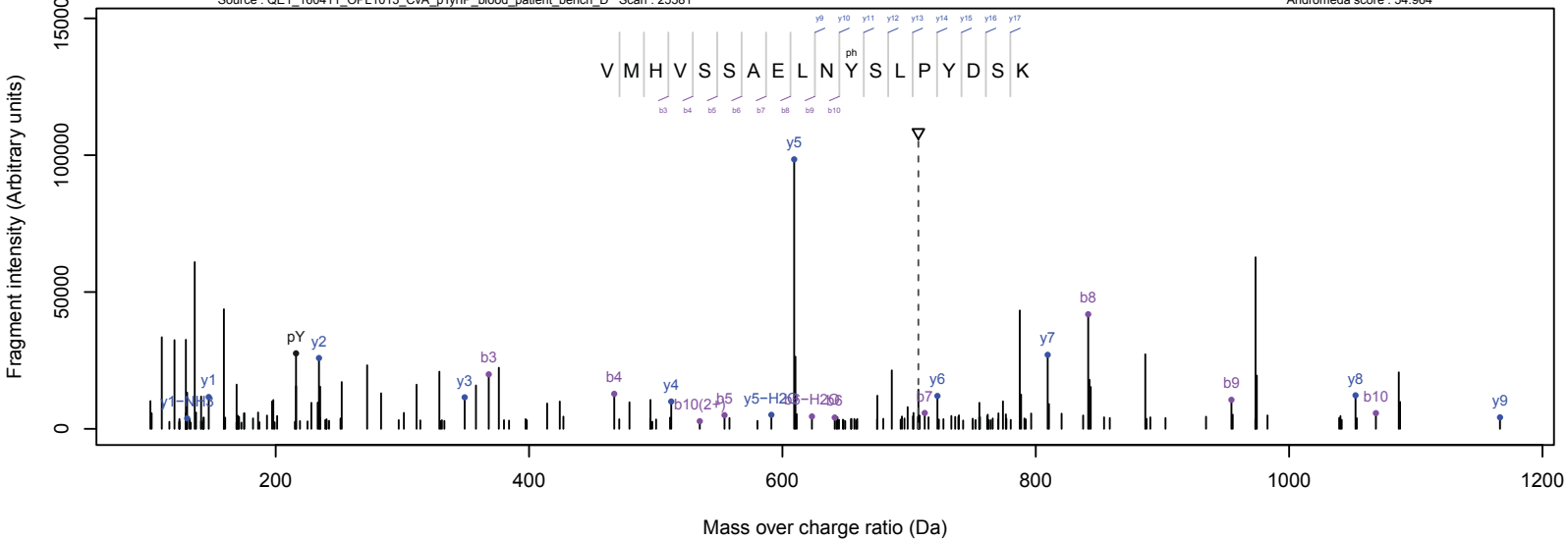
Andromeda score : 55.011



VMHVSSAELNY(ph)SLPYDSK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 25581

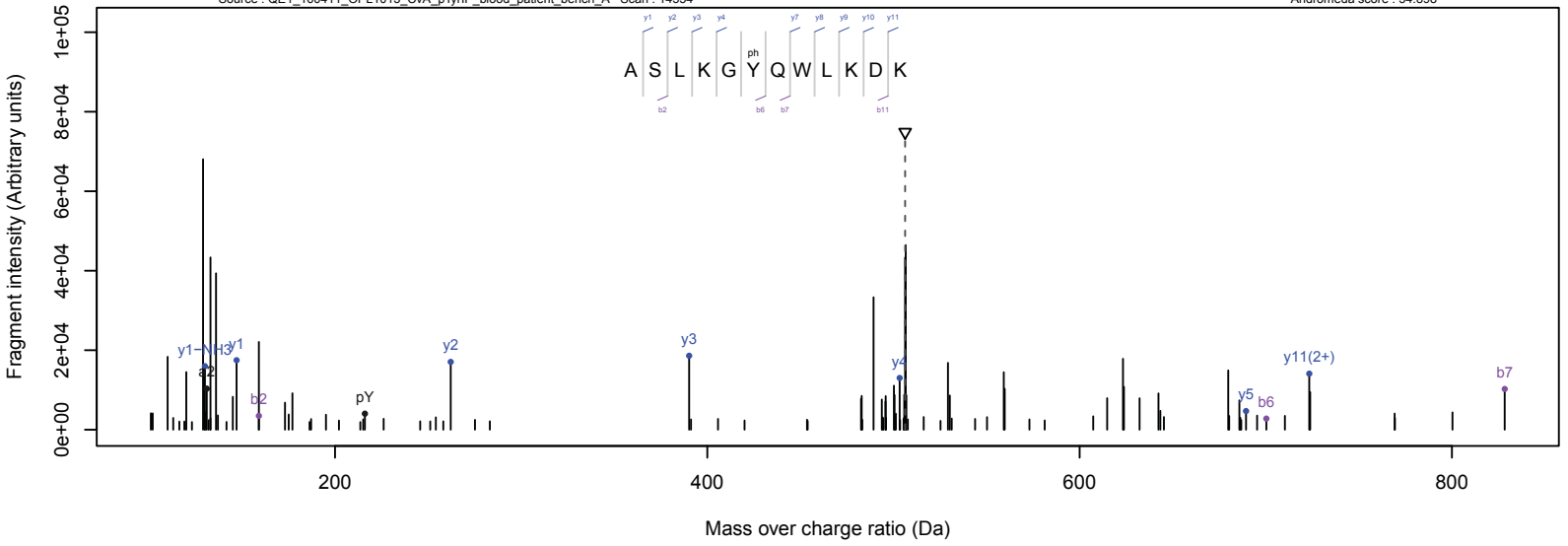
Andromeda score : 54.964



**ASLKG<sup>(ph)</sup>QWLKDK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 14554

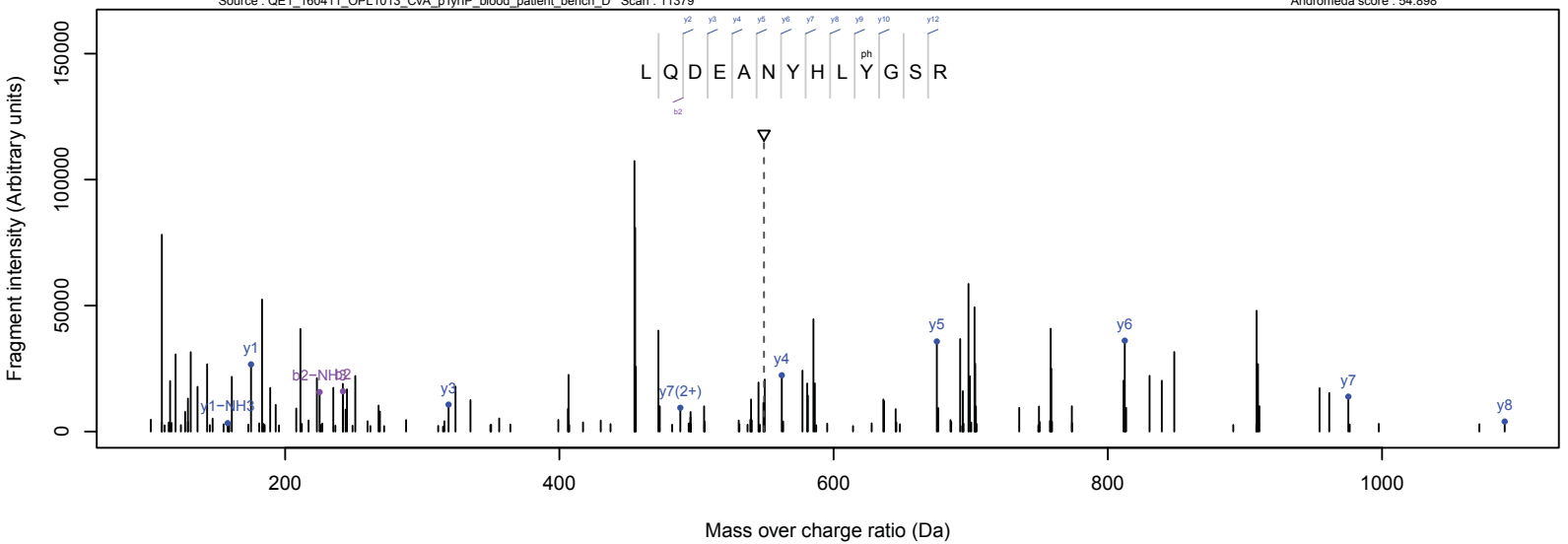
Andromeda score : 54.898



**LQDEANYHLY<sup>(ph)</sup>GSR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 11379

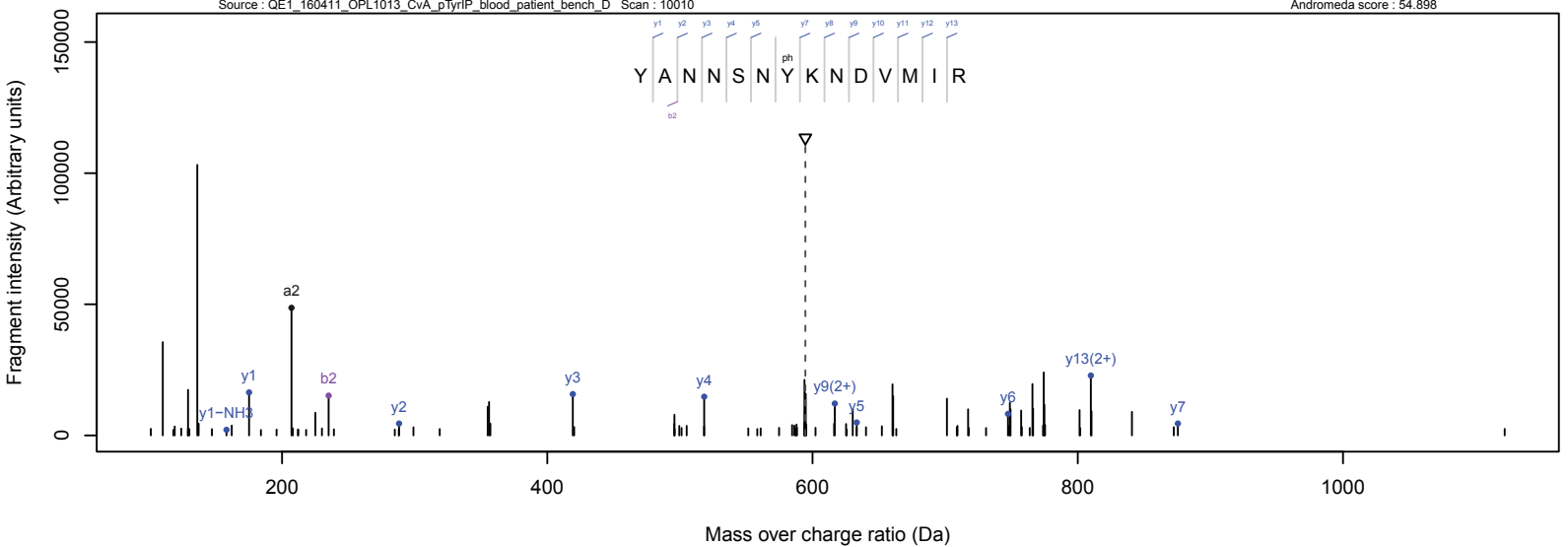
Andromeda score : 54.898



**YANNS<sup>(ph)</sup>KNVDMIR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 10010

Andromeda score : 54.898

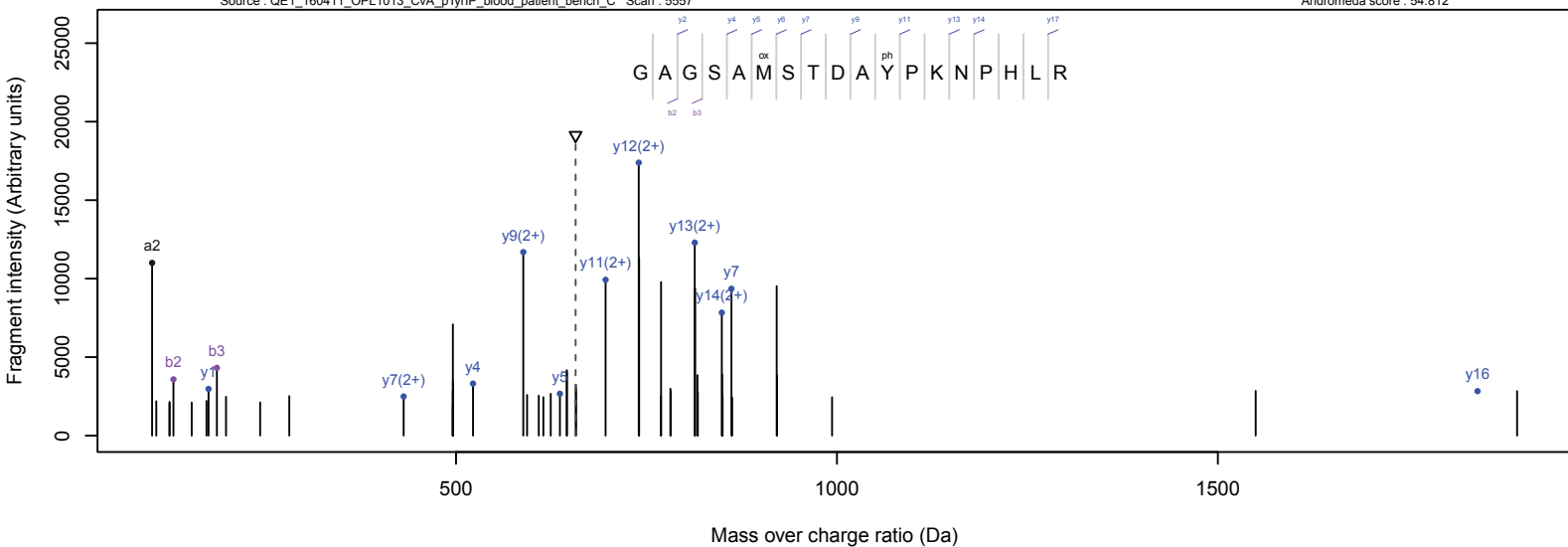




GAGSAM(ox)STDAY(ph)PKNPHLR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 5557

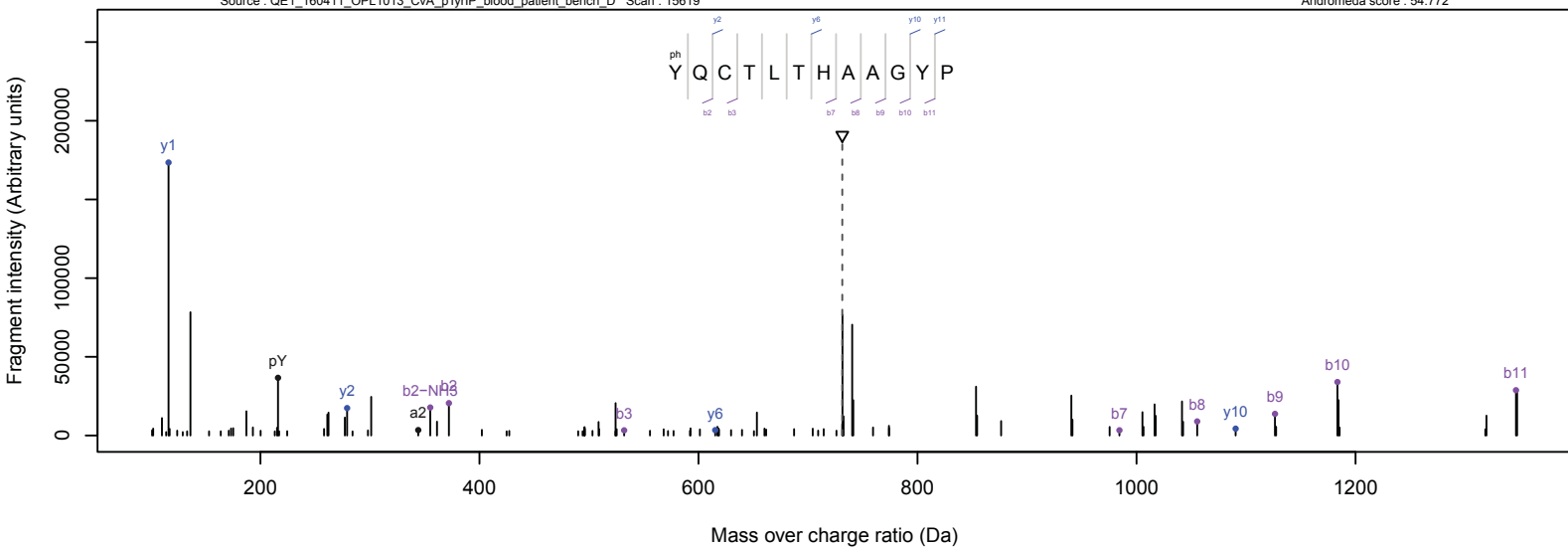
Andromeda score : 54.812



Y(ph)QCTLTHAAGYP-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 15619

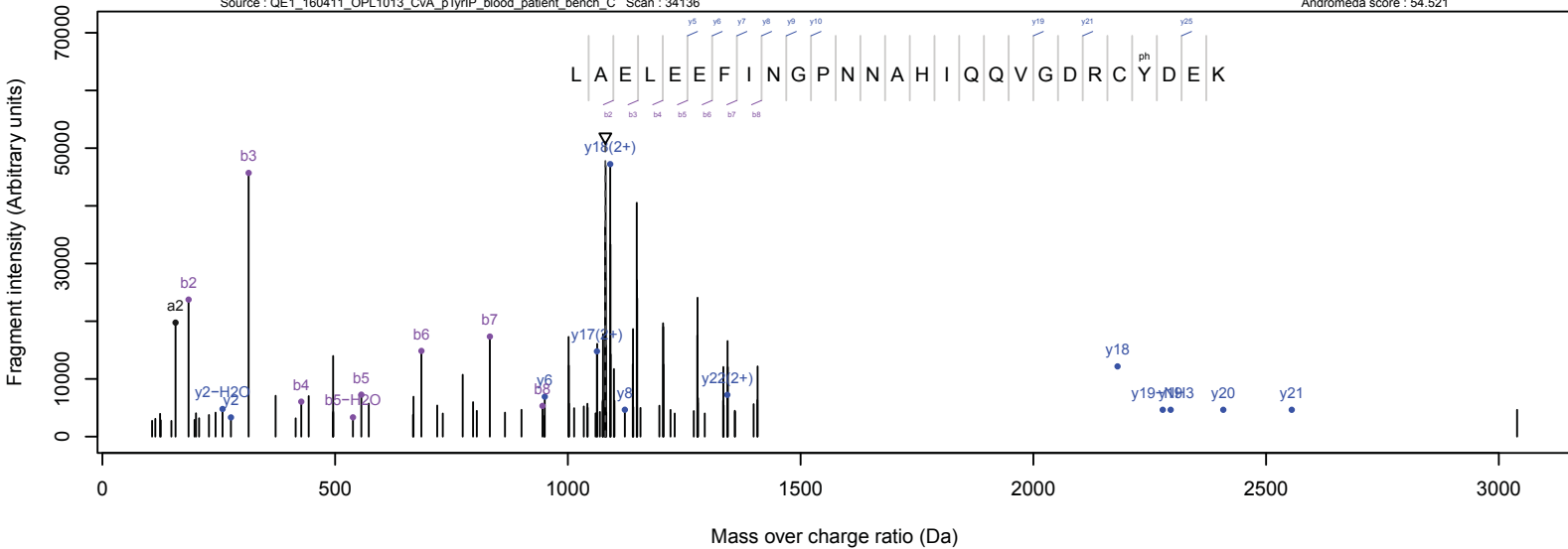
Andromeda score : 54.772



LAELEEFINGPNNNAHIQQVGDRCY(ph)DEK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 34136

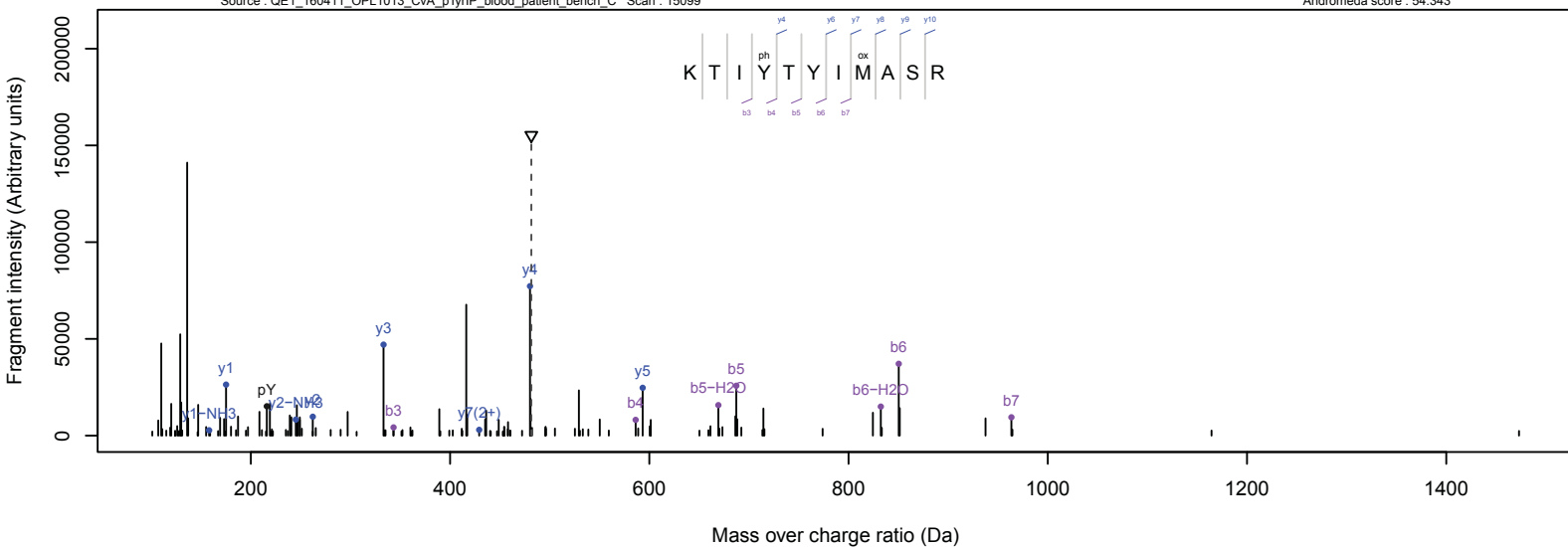
Andromeda score : 54.521



**KTIY(ph)TYIM(ox)ASR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 15099

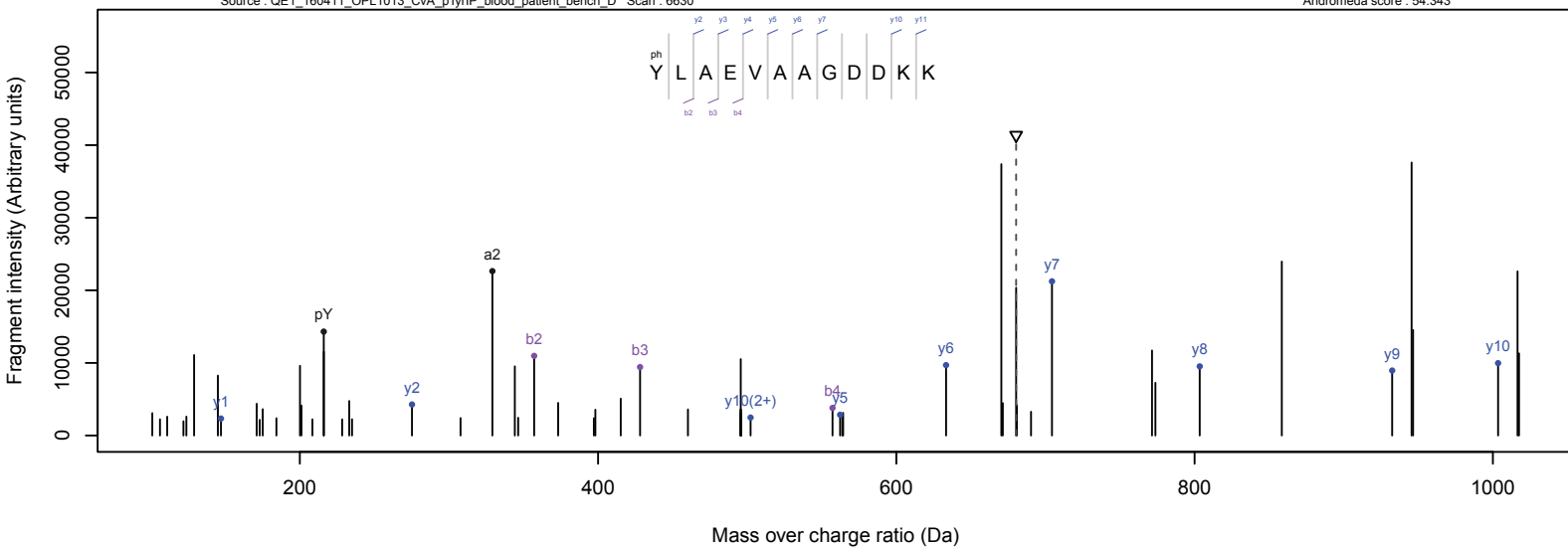
Andromeda score: 54.343



**Y(ph)LAEVAAGDDK-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 6630

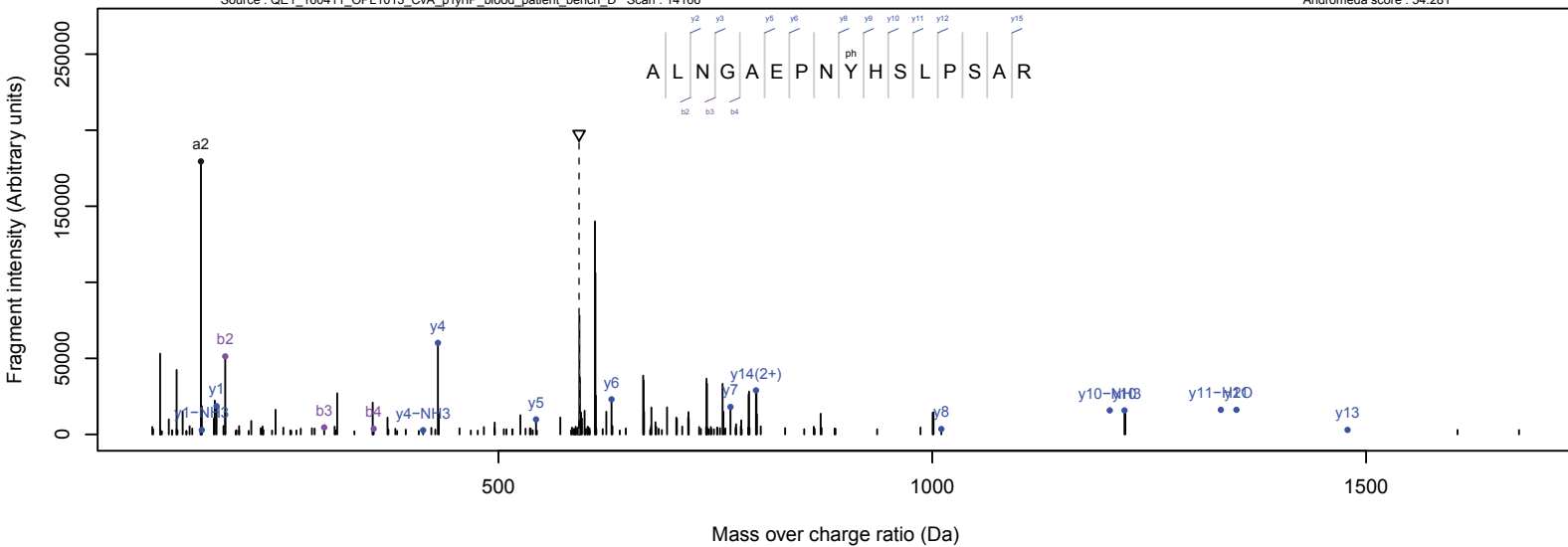
Andromeda score: 54.343



**ALNGAEPNY(ph)HSLPSAR-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 14166

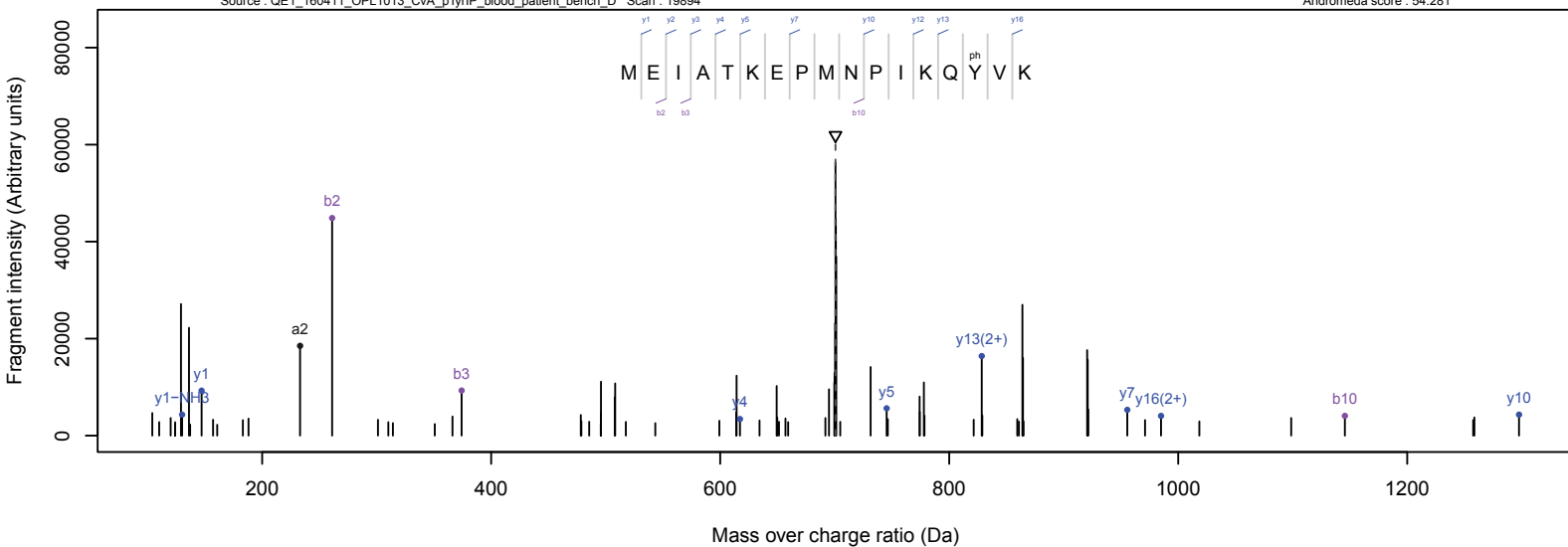
Andromeda score: 54.281



MEIATKEPMNPIK<sup>ph</sup>VK-3+

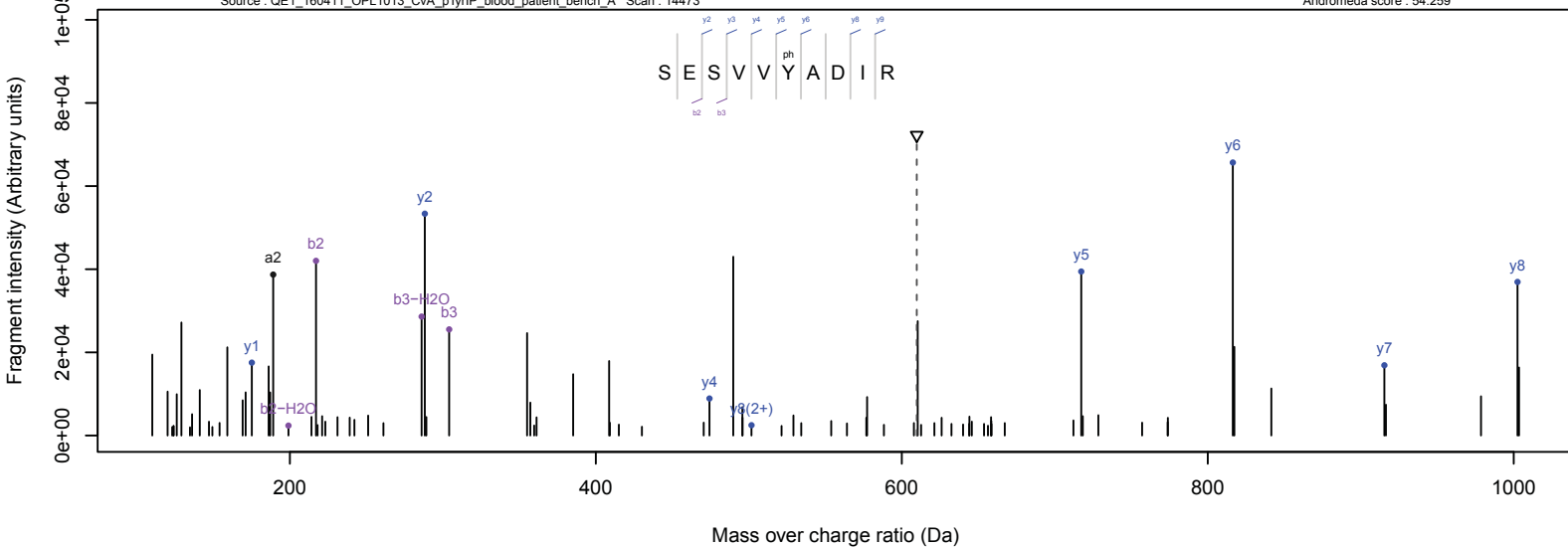
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 19894

Andromeda score: 54.281

SESVVY<sup>ph</sup>ADIR-2+

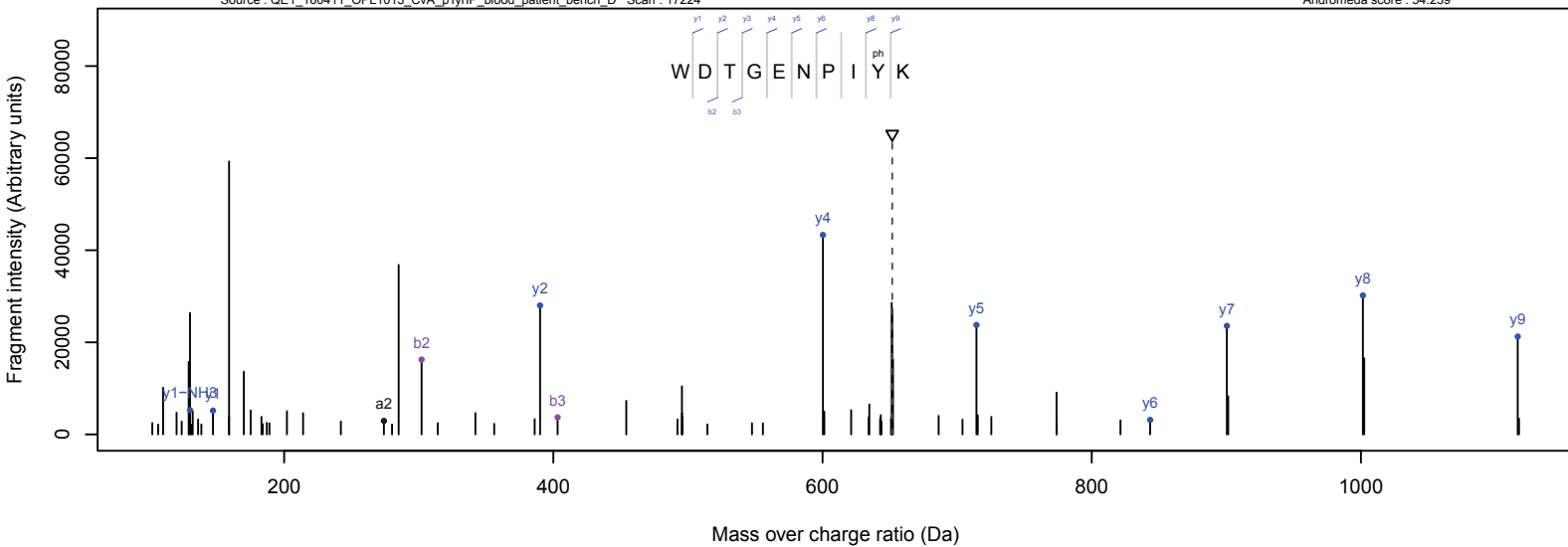
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 14473

Andromeda score: 54.259

WDTGENPIY<sup>ph</sup>K-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 17224

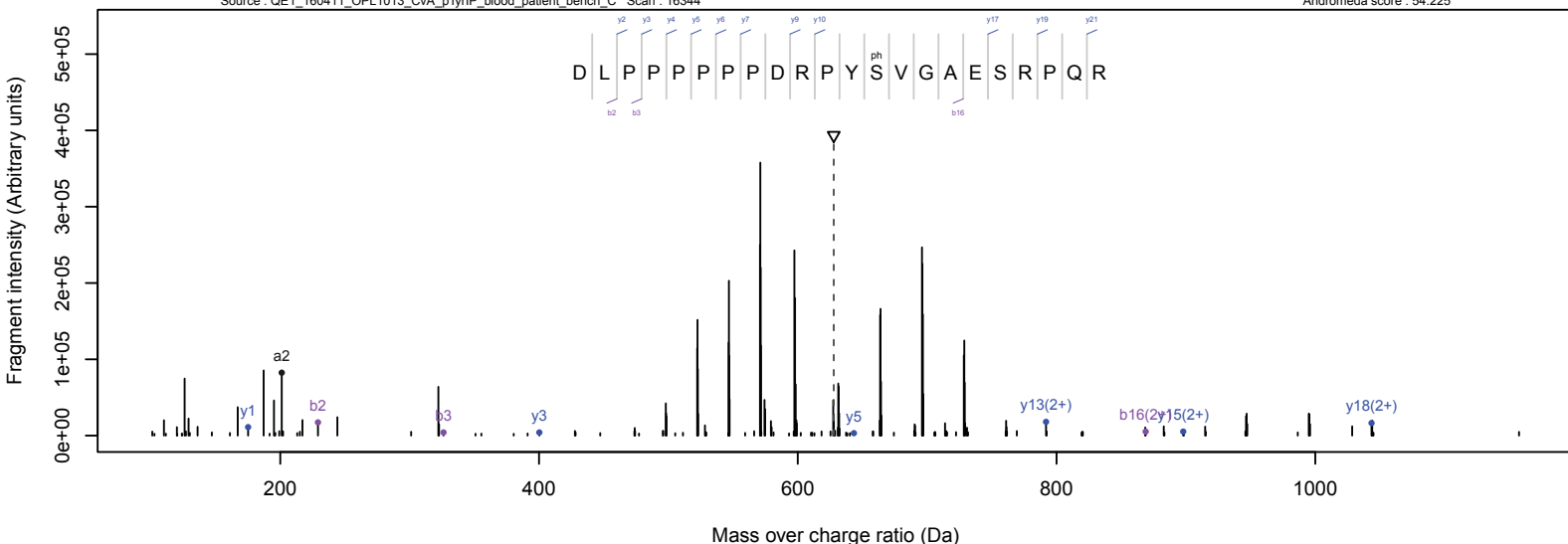
Andromeda score: 54.259



**DLPPPPDRPYS(ph)VGAESRPQR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16344

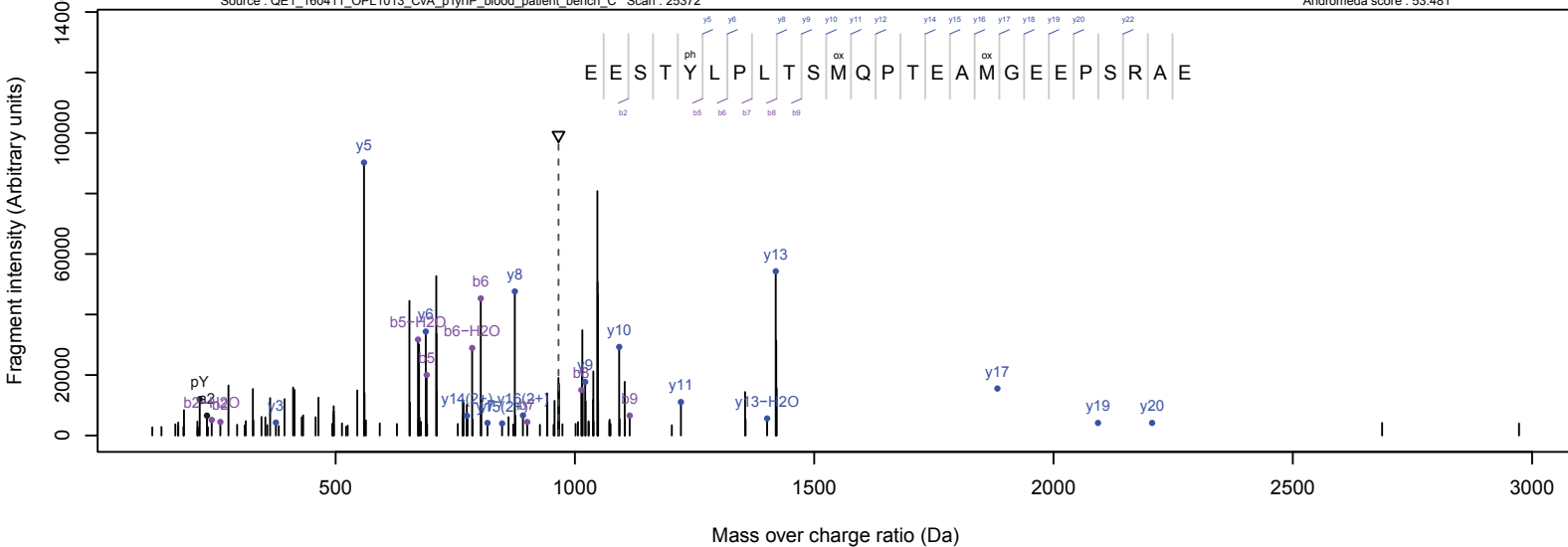
Andromeda score : 54.225



**EESTY(ph)LPLTSM(ox)QPTEAM(ox)GEEPSRAE-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 25372

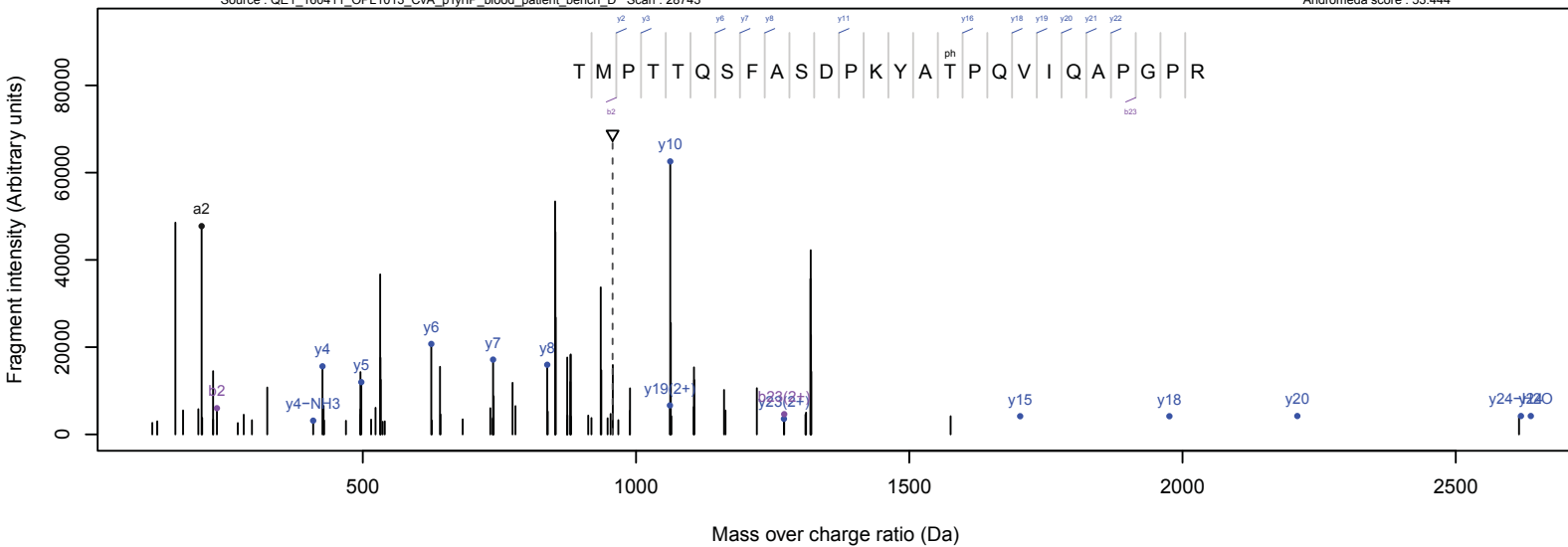
Andromeda score : 53.481



**TMPTTQSFASDPKYAT(ph)PQVIQAPGPR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28743

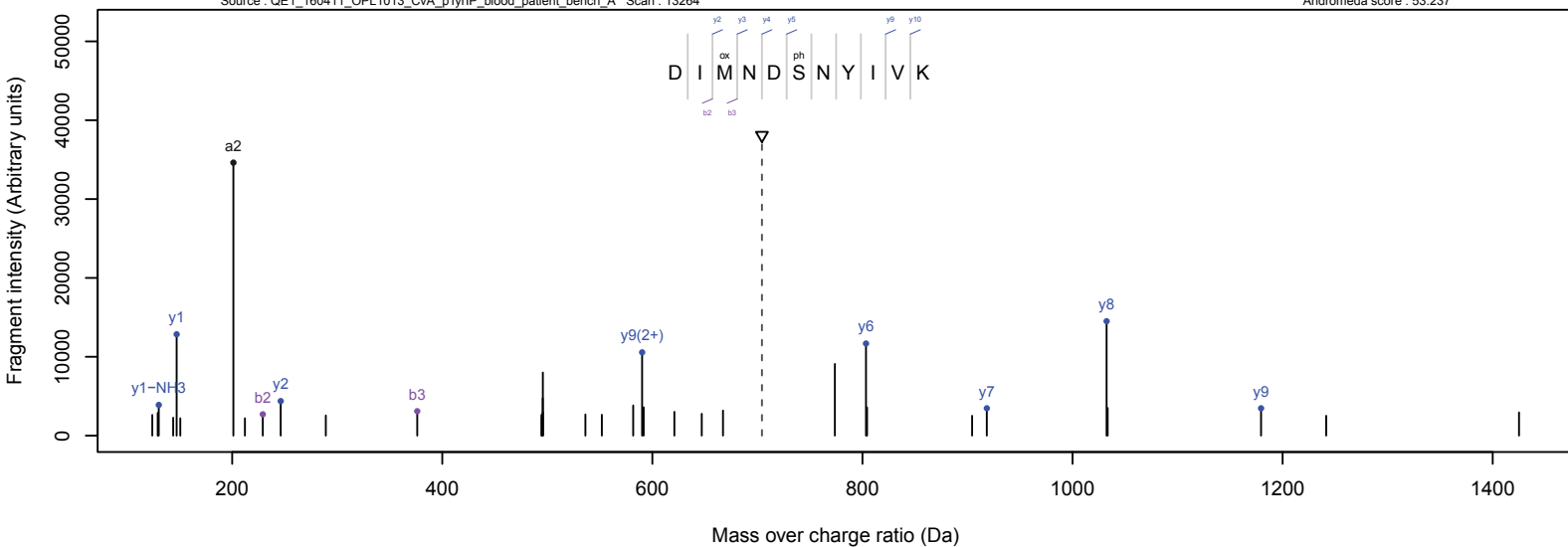
Andromeda score : 53.444



**DIM(ox)NDS(ph)NYIVK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 13284

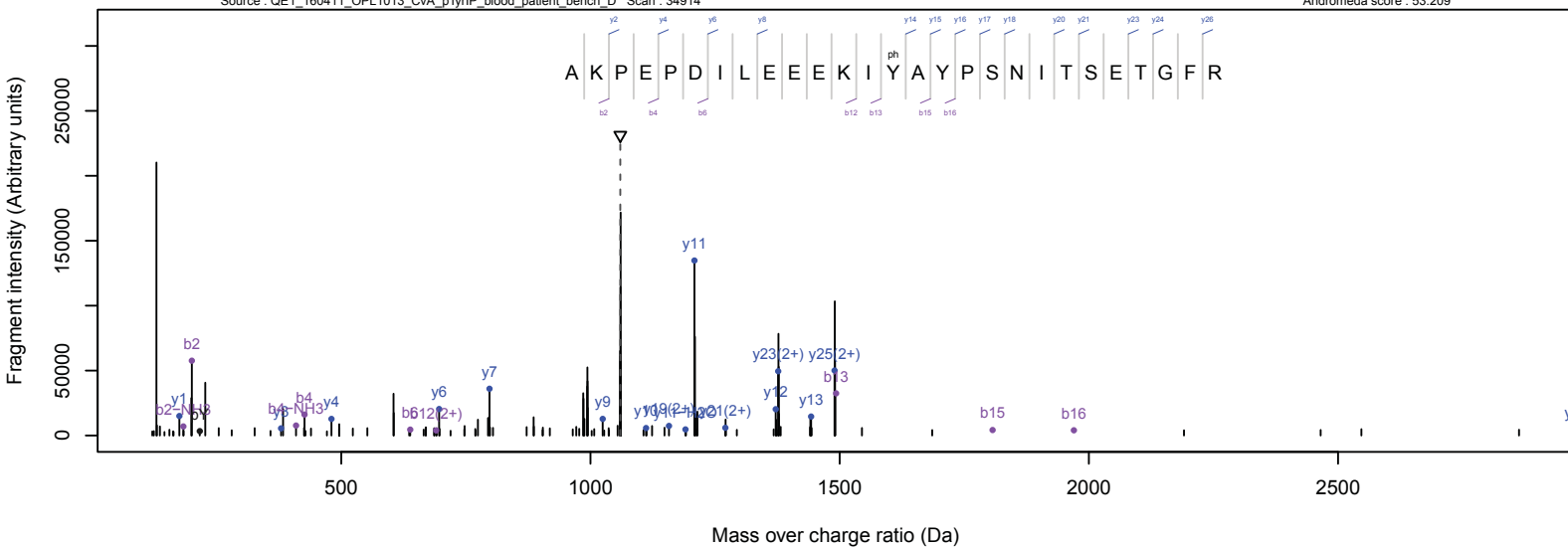
Andromeda score : 53.237



**AKPEPDILEEEKIYAYPSNITSETGFR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 34914

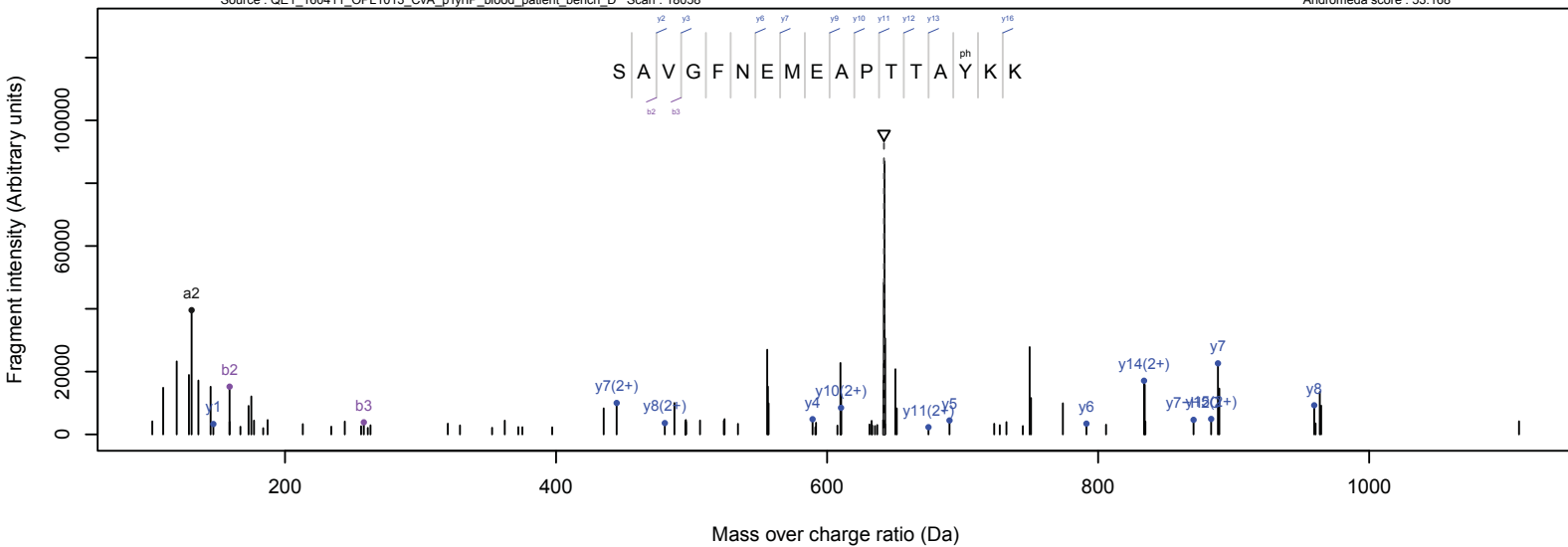
Andromeda score : 53.209



**SAVGFNEMEAPTAYK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 18058

Andromeda score : 53.168

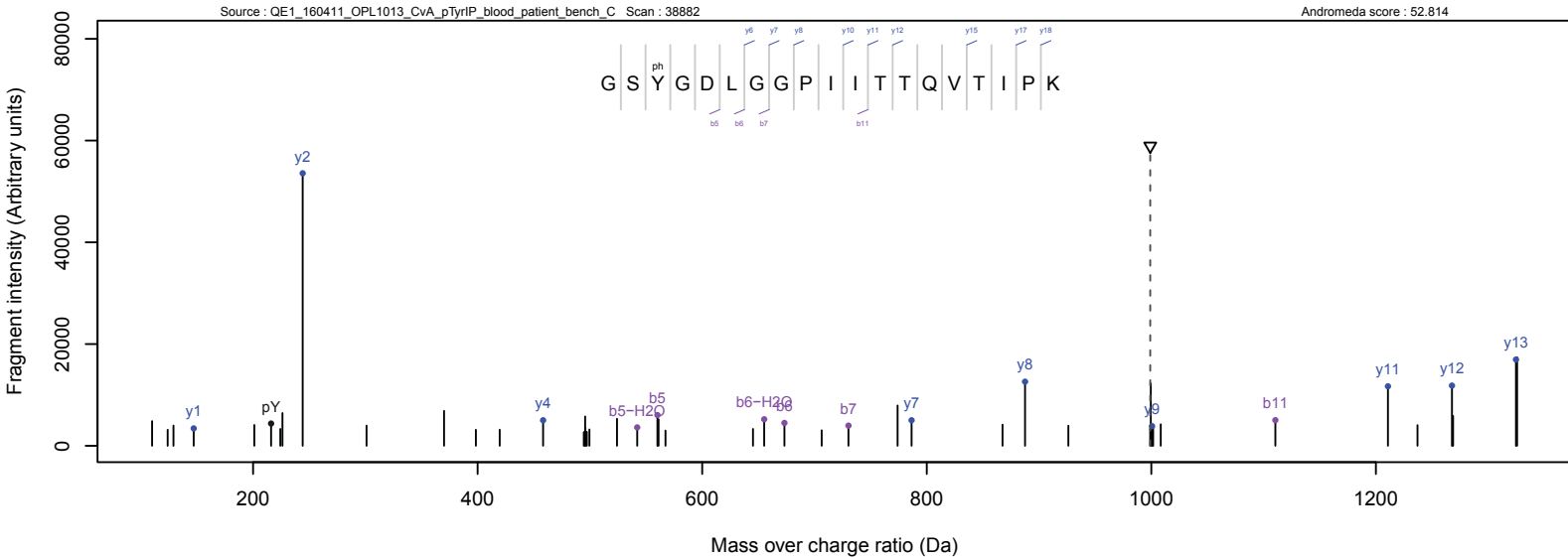




**GSY(ph)GDLGGPIITTQVTIPK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 38882

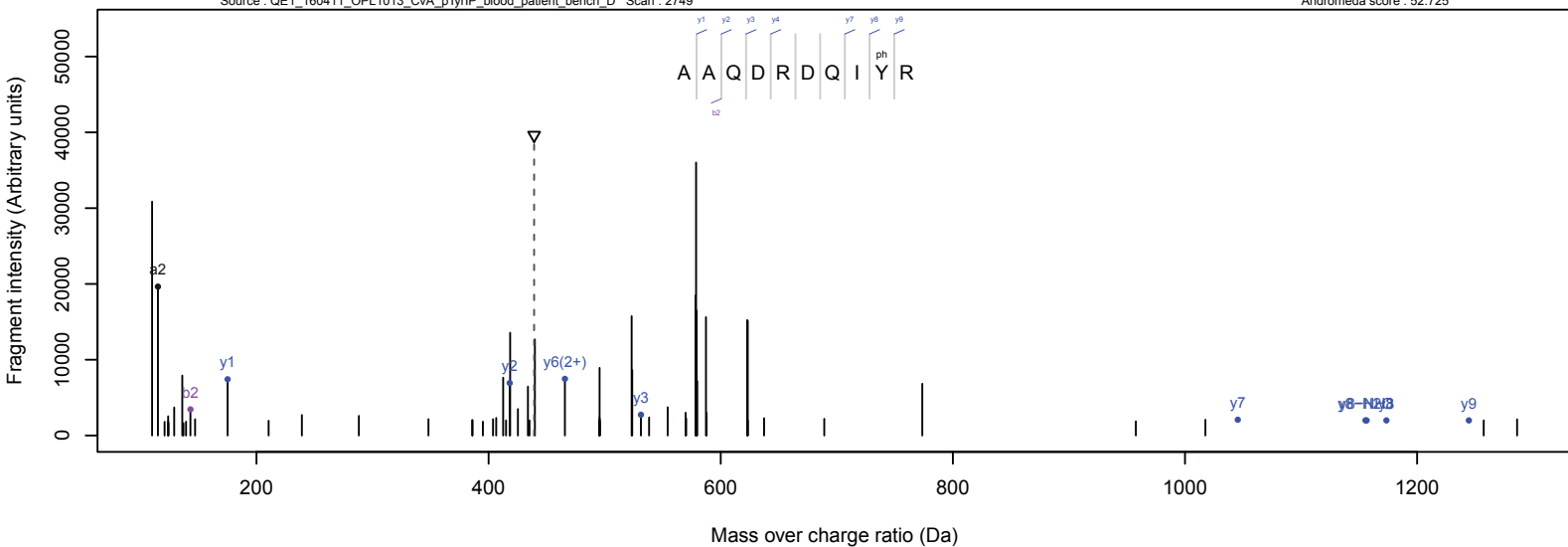
Andromeda score : 52.814



**AAQDRDQIY(ph)R-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 2749

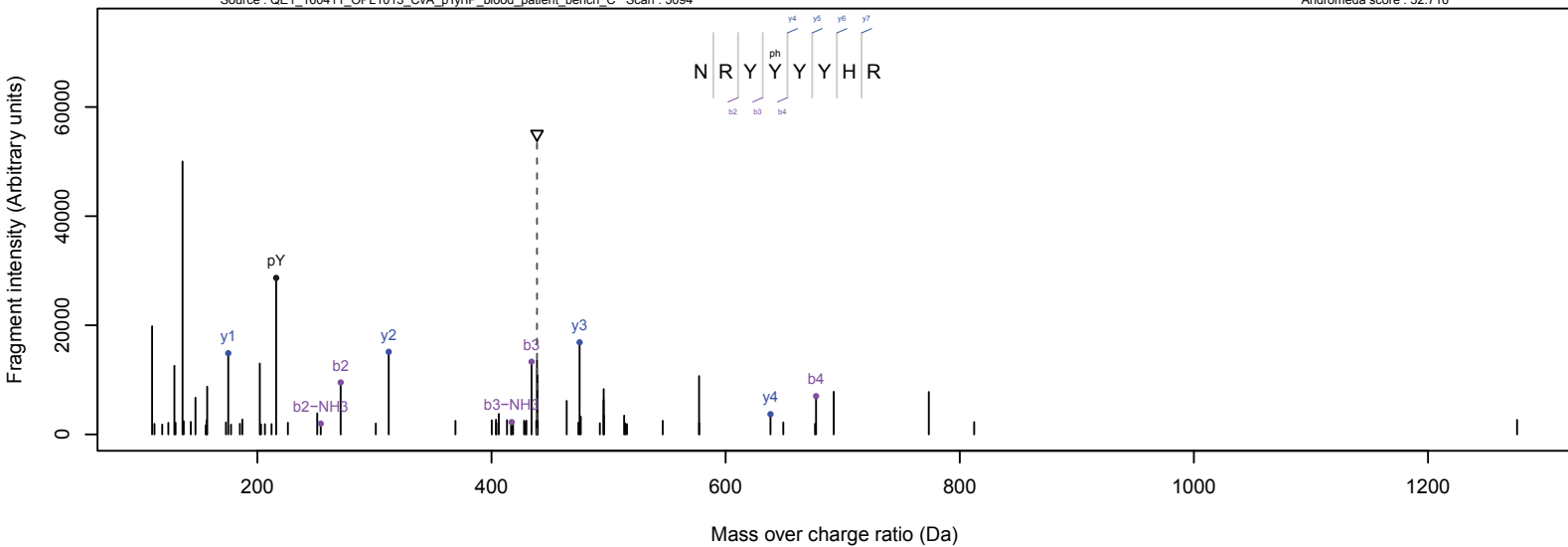
Andromeda score : 52.725



**NRYY(ph)YYHR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 5094

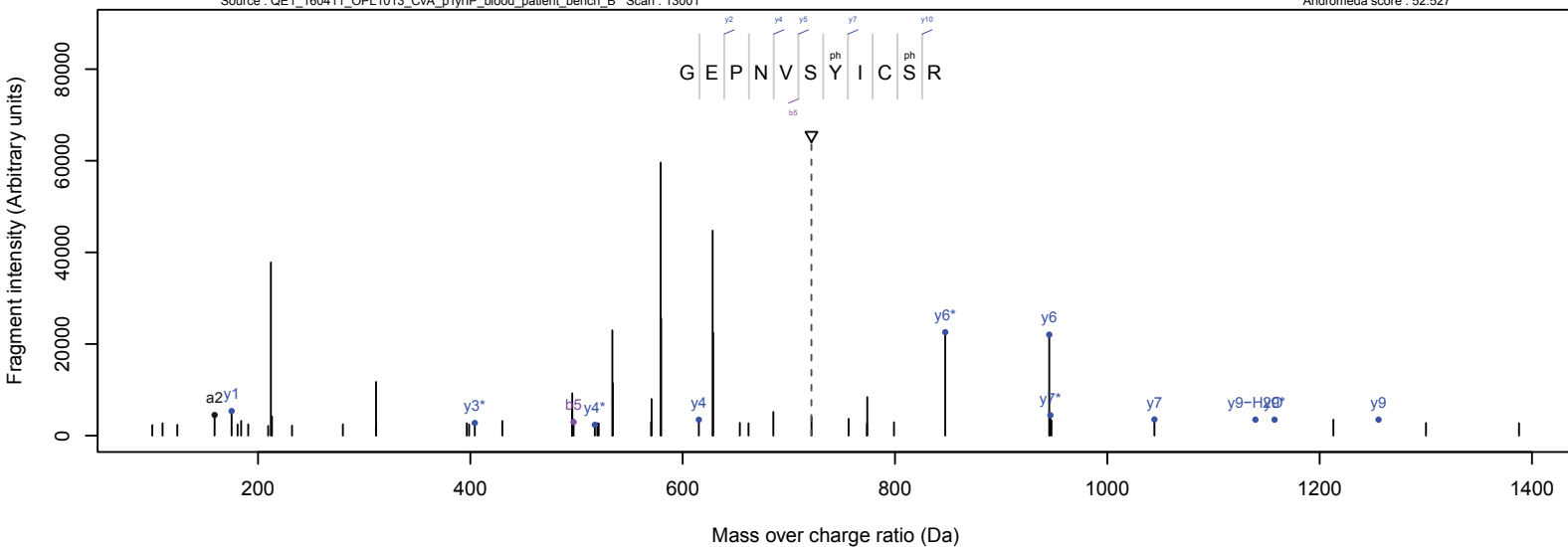
Andromeda score : 52.716



GEPNVS(ph)ICS(ph)R-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 13001

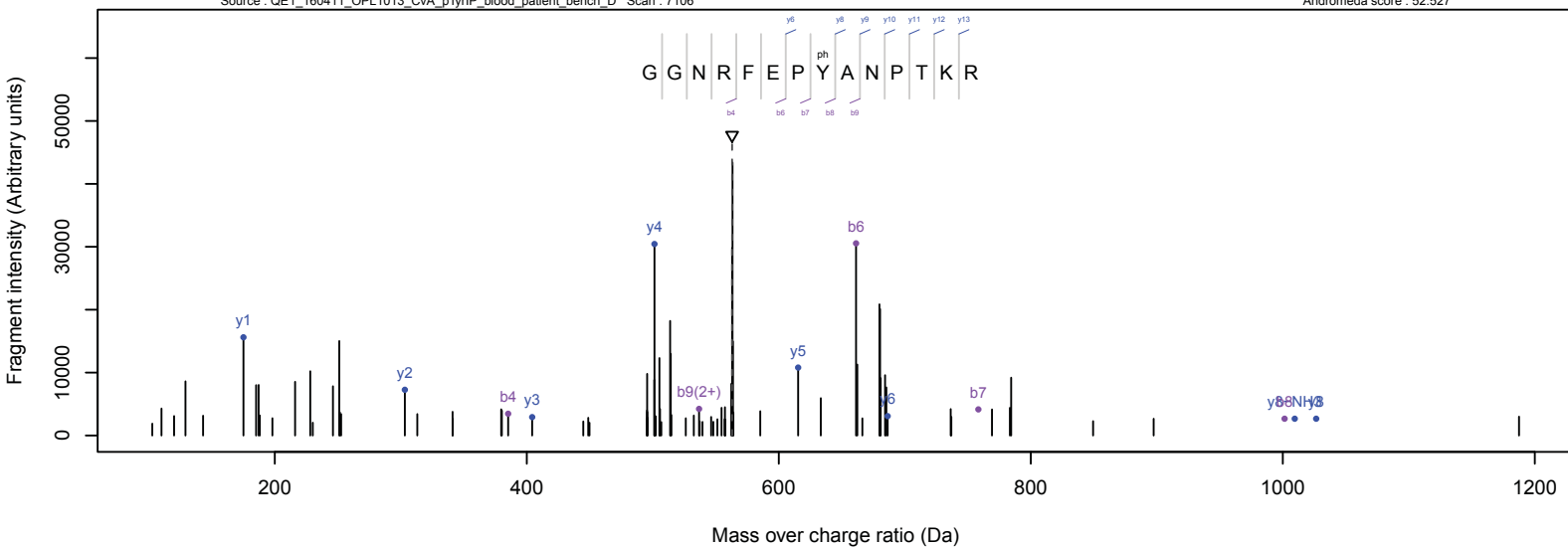
Andromeda score : 52.527



GGNRFEPY(ph)ANPTKR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7106

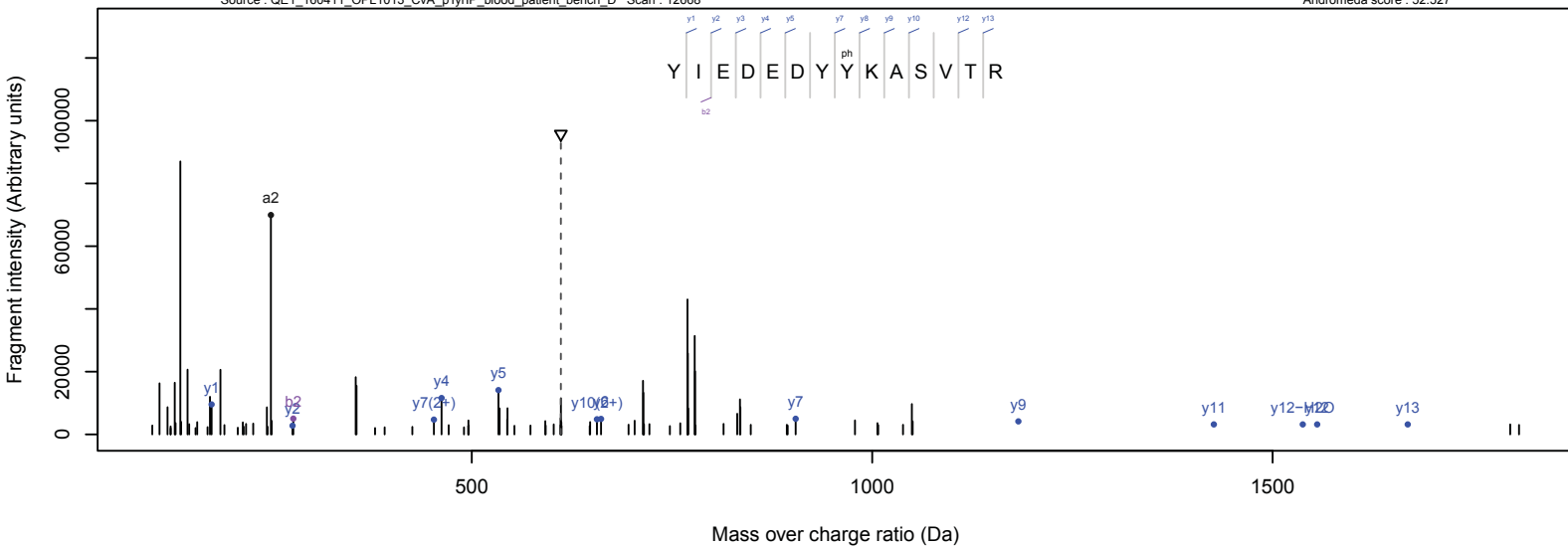
Andromeda score : 52.527



YIEDEDY(ph)KASVTR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12668

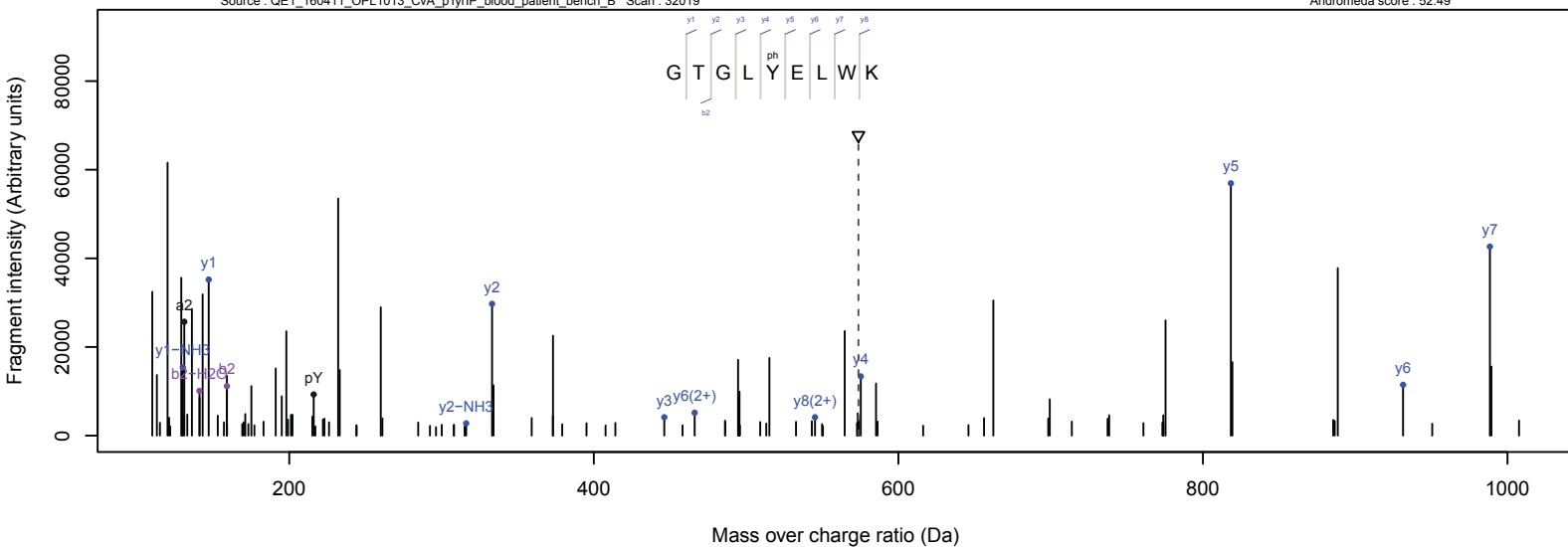
Andromeda score : 52.527



GTGLY(ph)ELWK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 32019

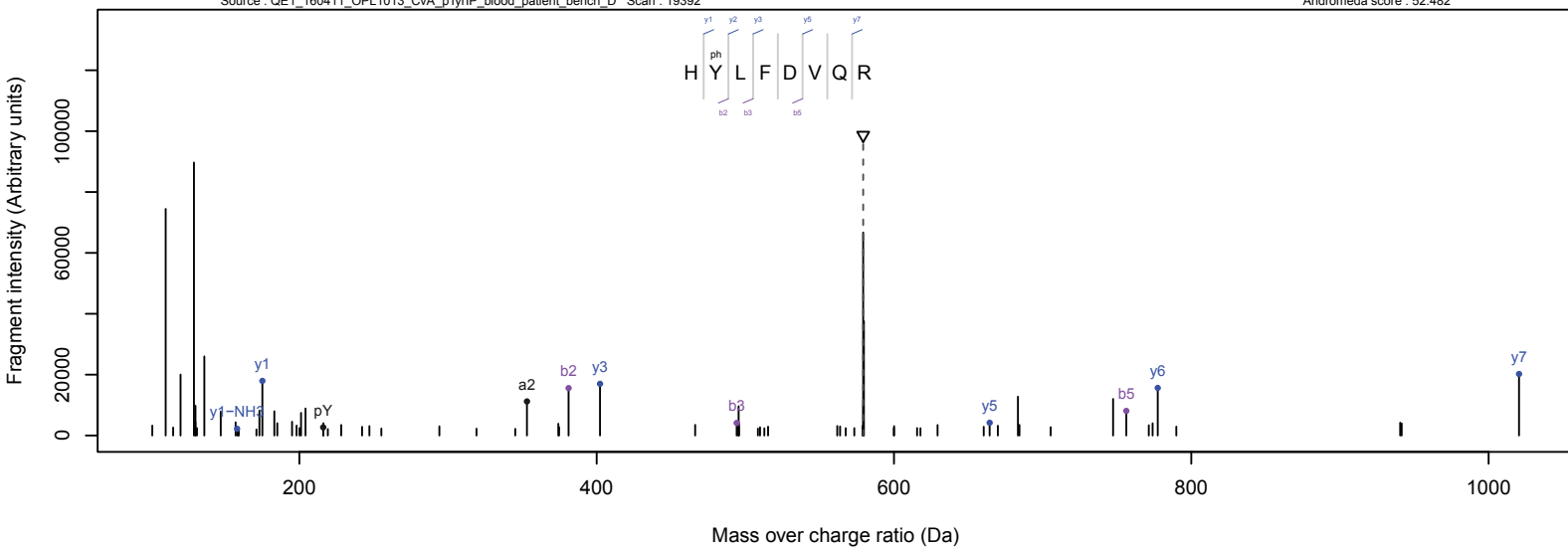
Andromeda score : 52.49



HY(ph)LFDVQR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 19392

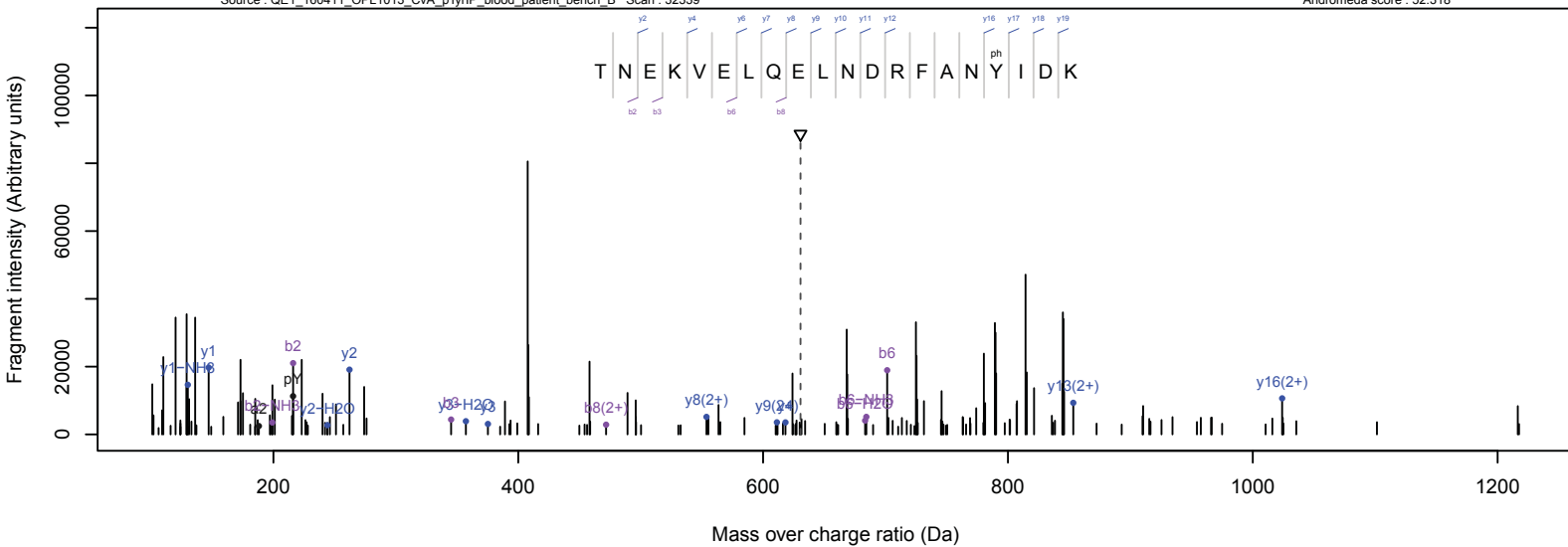
Andromeda score : 52.482



TNEKVELQELNDRFANY(ph)IDK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 32339

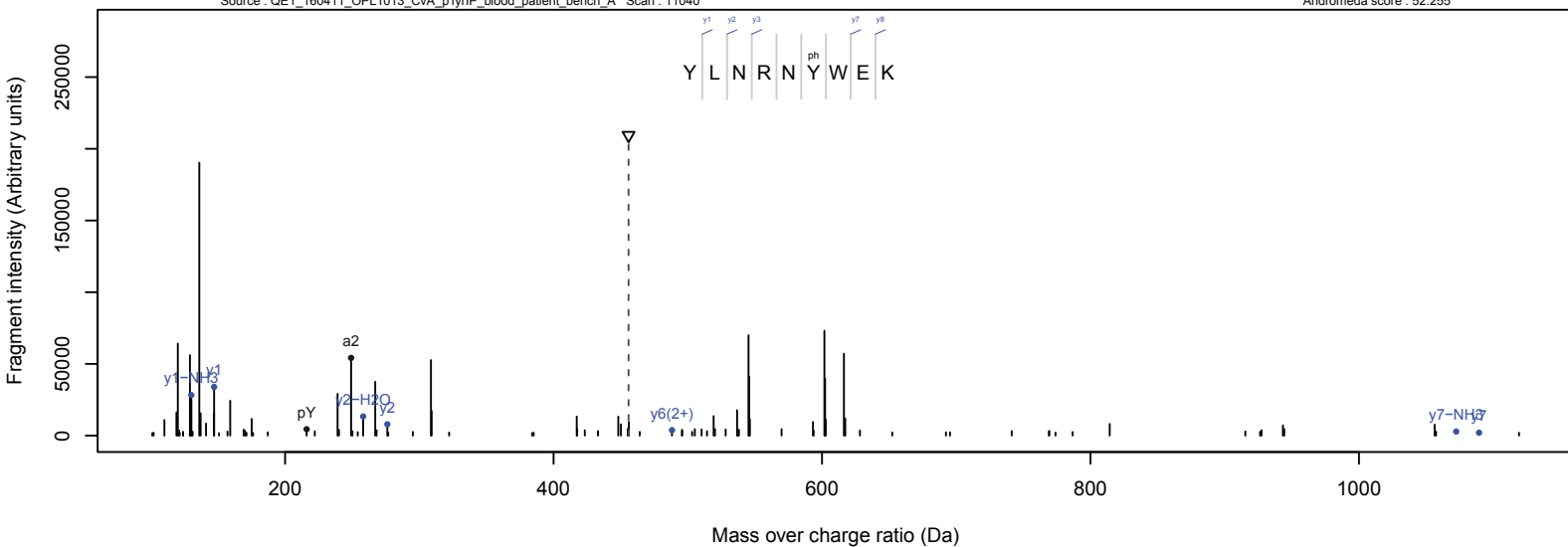
Andromeda score : 52.318



YLNRNY(ph)WEK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 11040

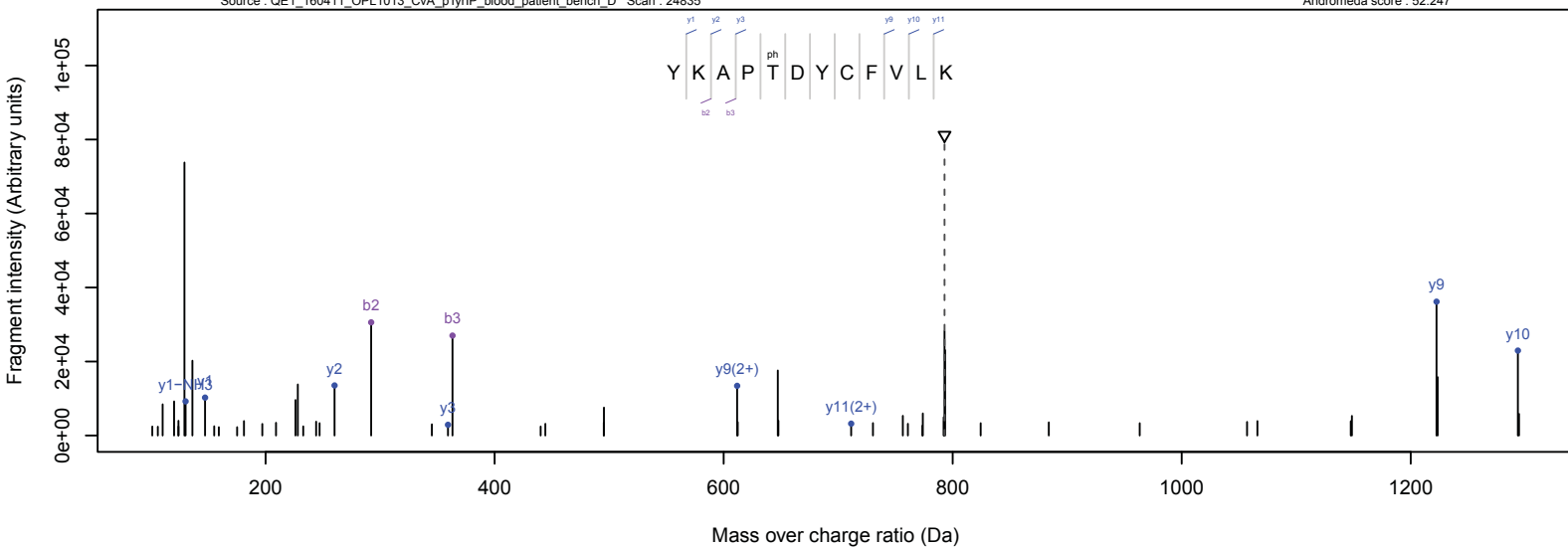
Andromeda score : 52.255



YKAPT(ph)DYCFVLK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24835

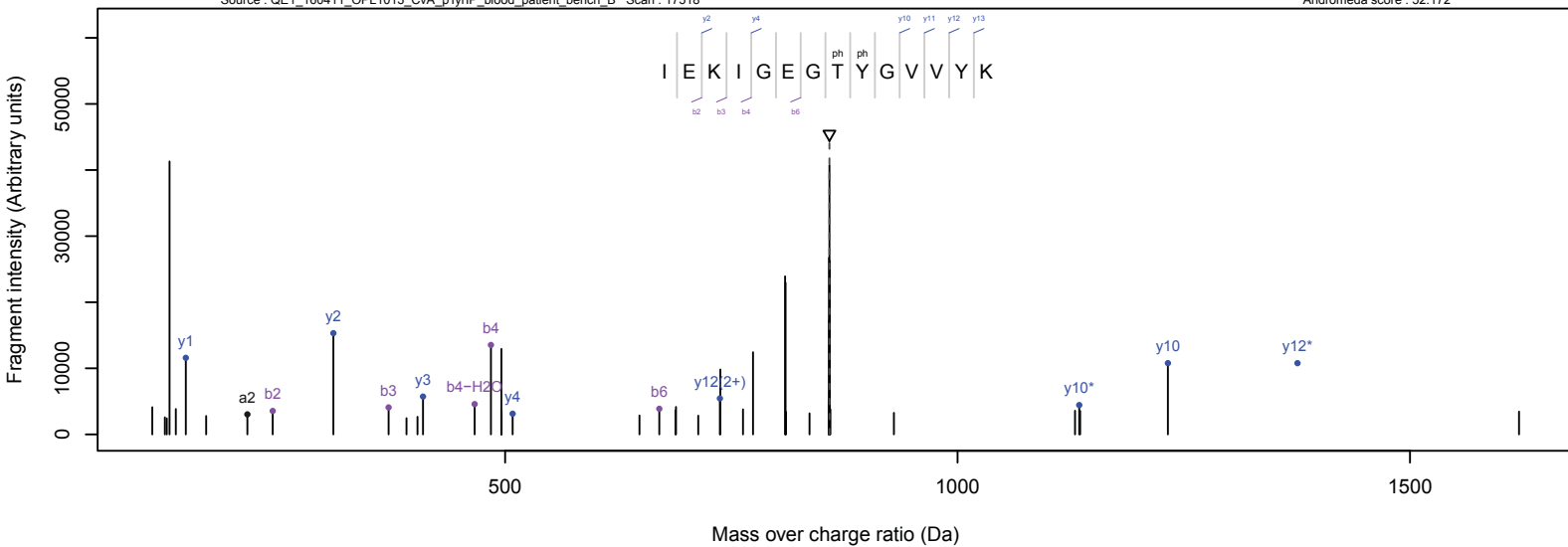
Andromeda score : 52.247



IEKIGEGT(ph)Y(ph)GVVYK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 17518

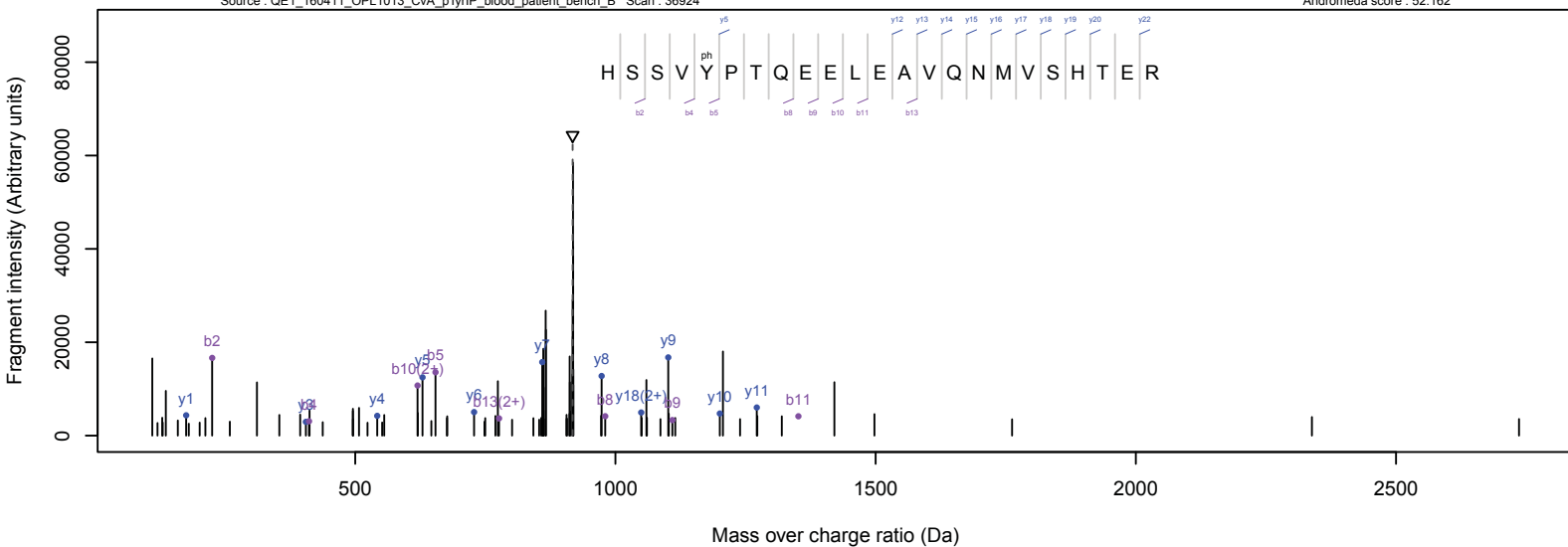
Andromeda score : 52.172



## HSSVY(ph)PTQEELEAVQNMVSHTER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 36924

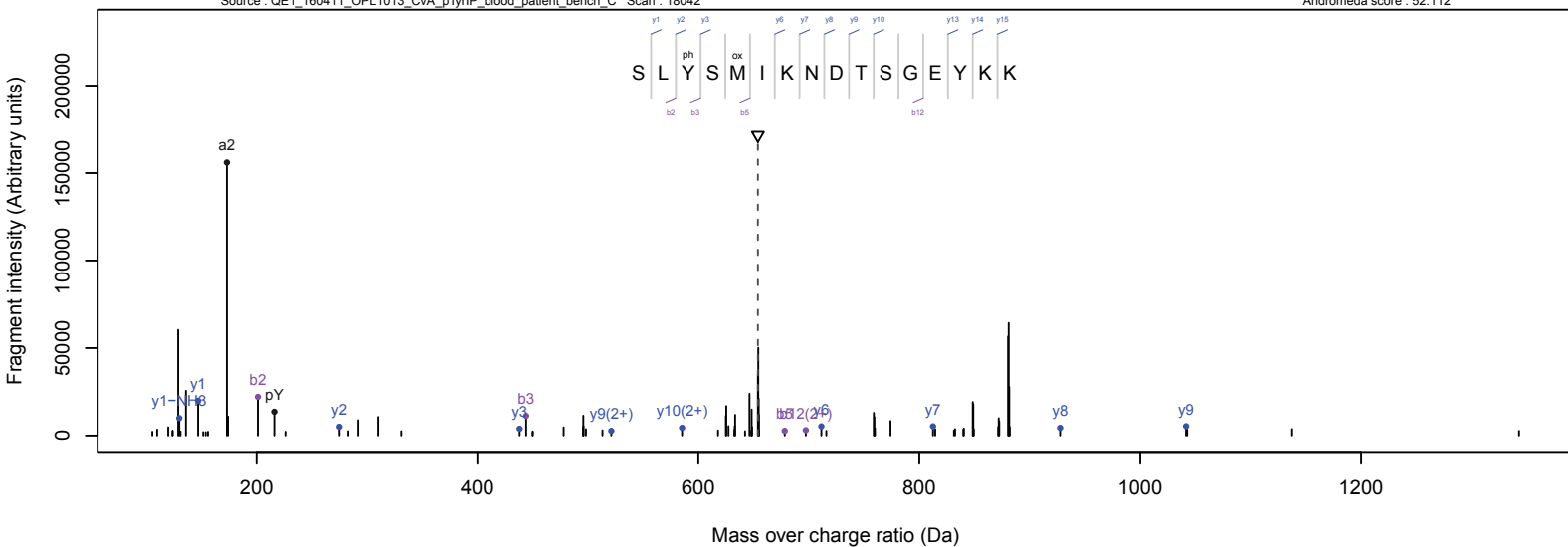
Andromeda score : 52.162



## SLY(ph)SM(ox)IKNDTSGEYKK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18042

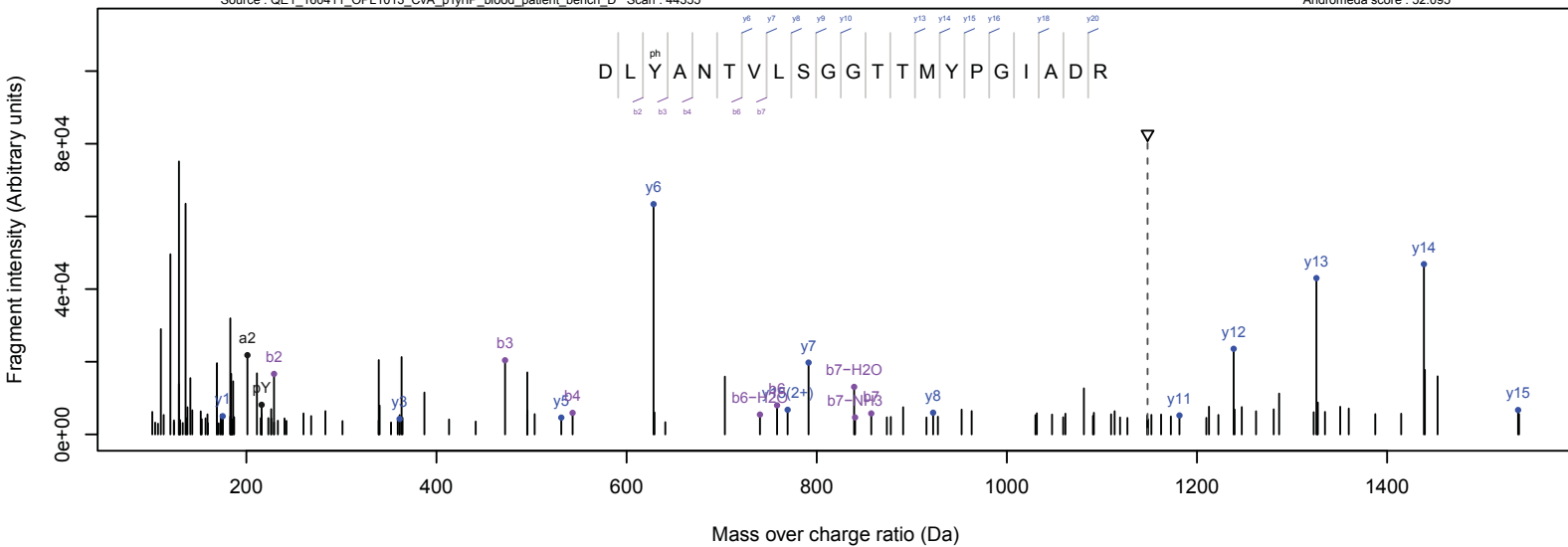
Andromeda score : 52.112



## DLY(ph)ANTVLSGGTTMYPGIADR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 44355

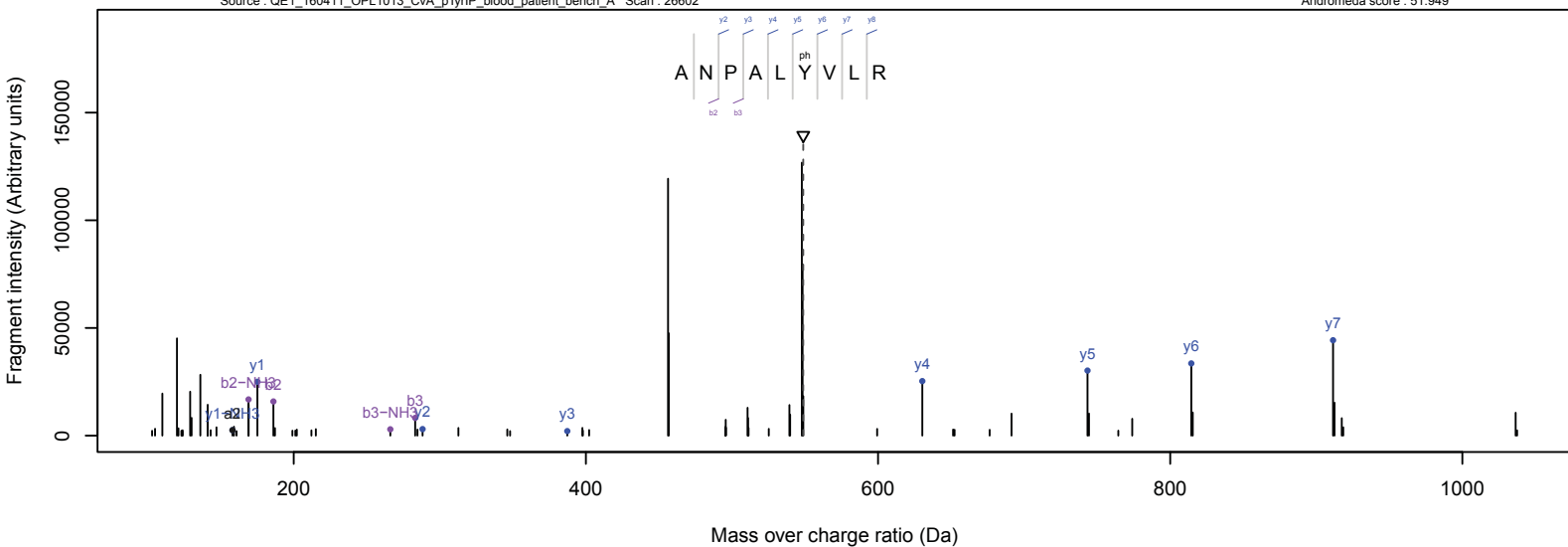
Andromeda score : 52.095



ANPALY(ph)VLR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 26602

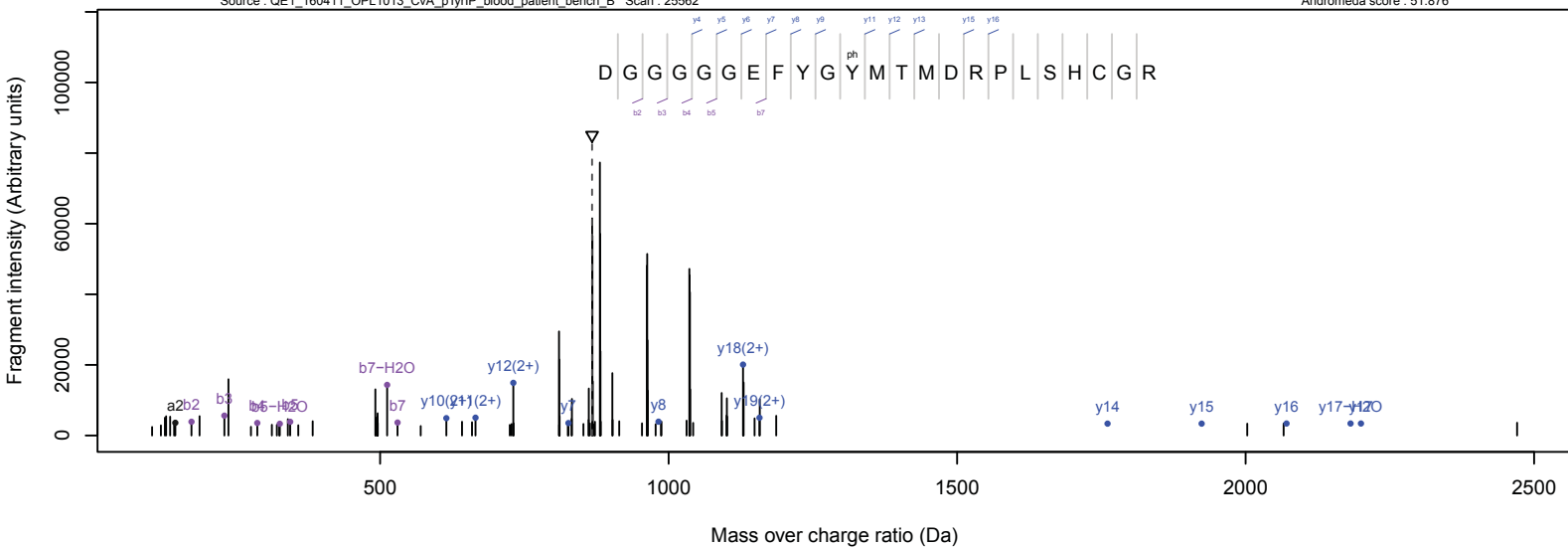
Andromeda score : 51.949



DGGGGGEFYGY(ph)MTMDRPLSHCGR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 25562

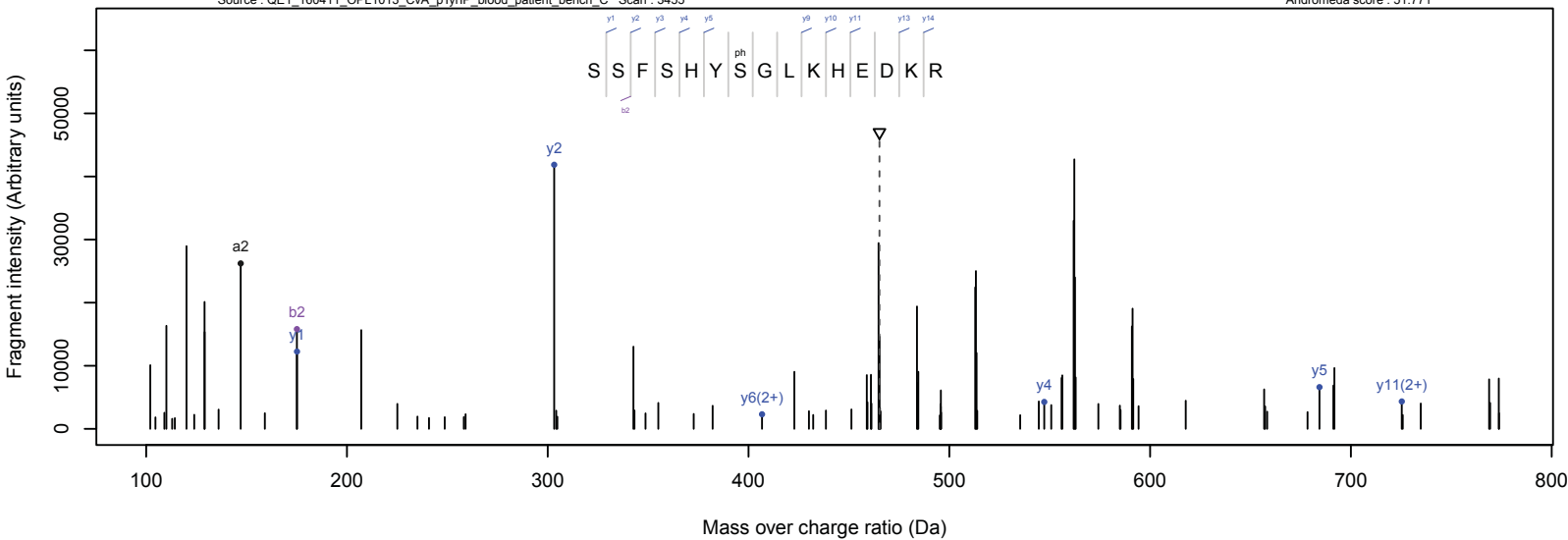
Andromeda score : 51.876



SSFHYS(ph)GLKHEDKR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3455

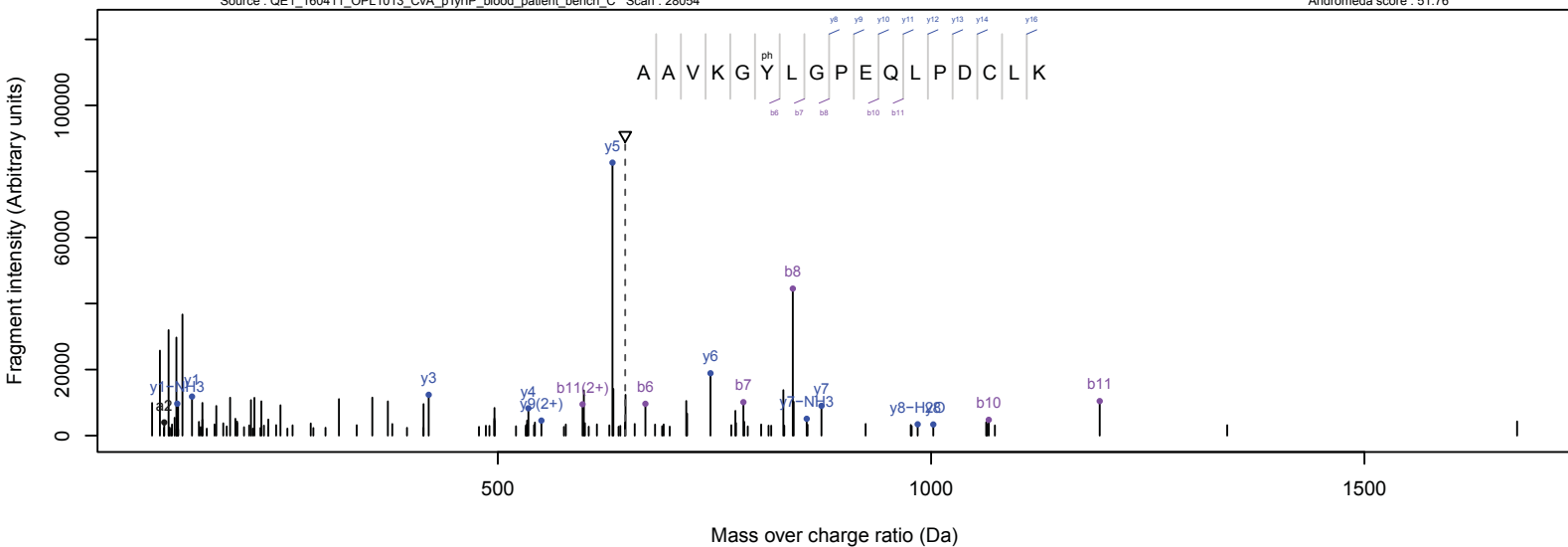
Andromeda score : 51.771



## AAVKGY(ph)LGPEQLPDCLK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28054

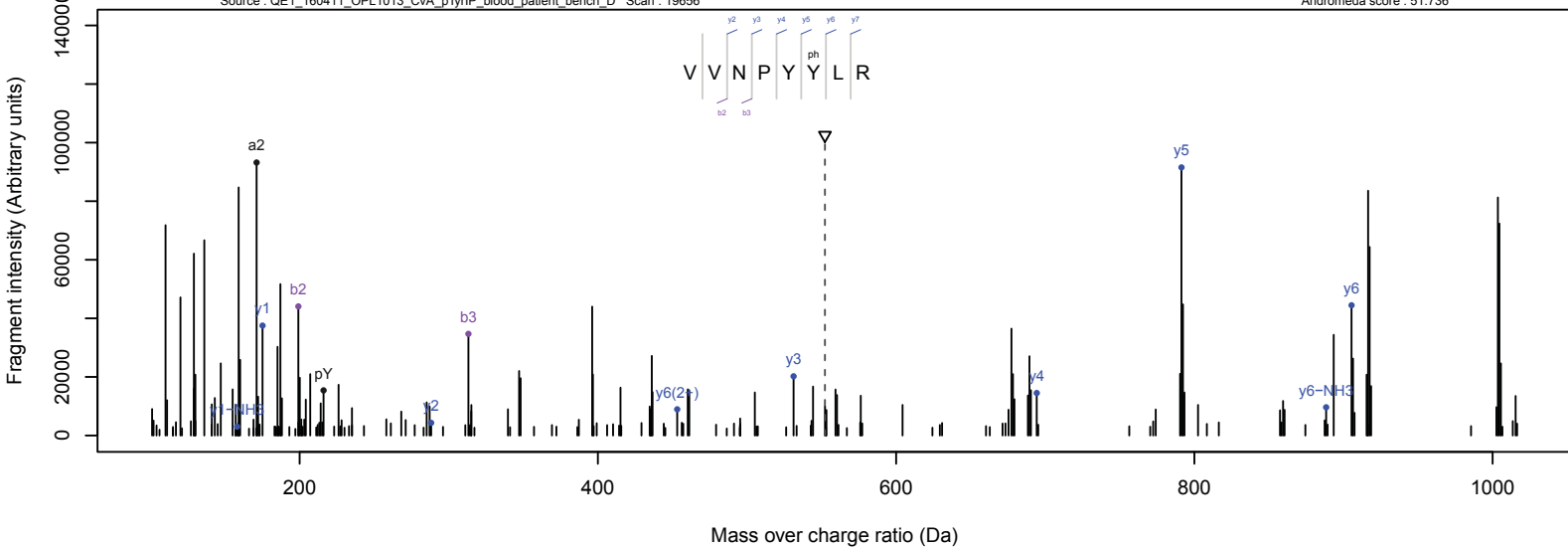
Andromeda score : 51.76



## VVNPYY(ph)LR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 19656

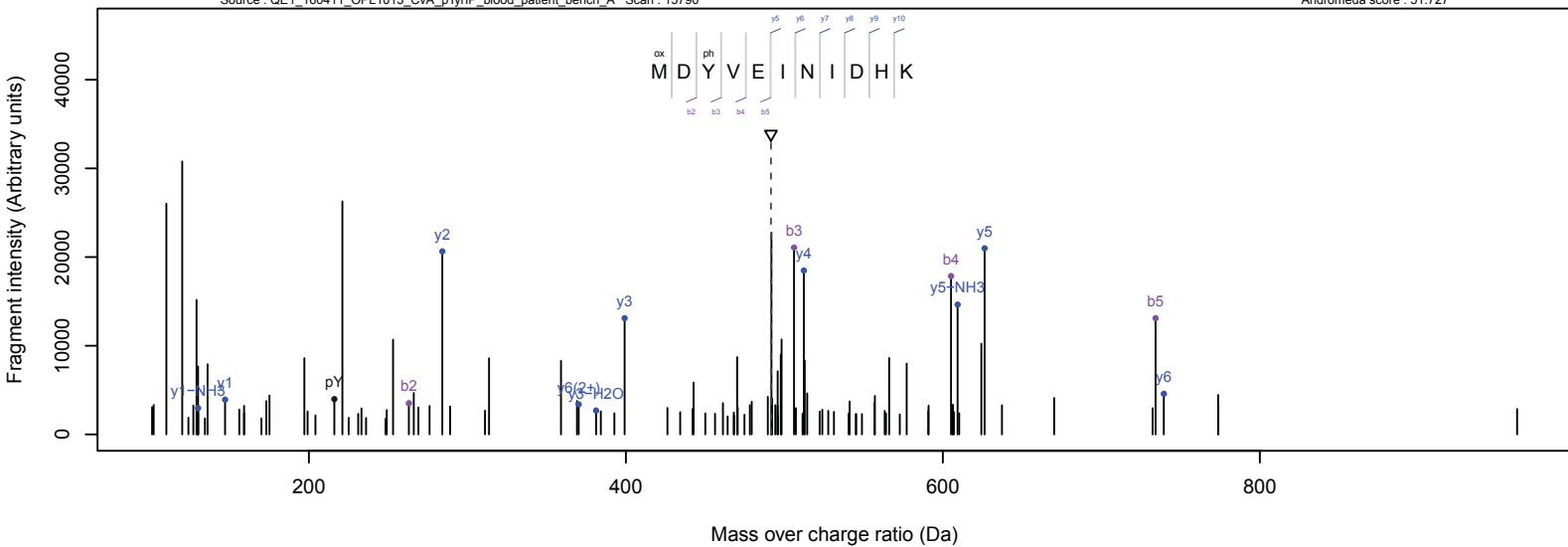
Andromeda score : 51.736



## M(ox)DY(ph)VEINIDHK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 15790

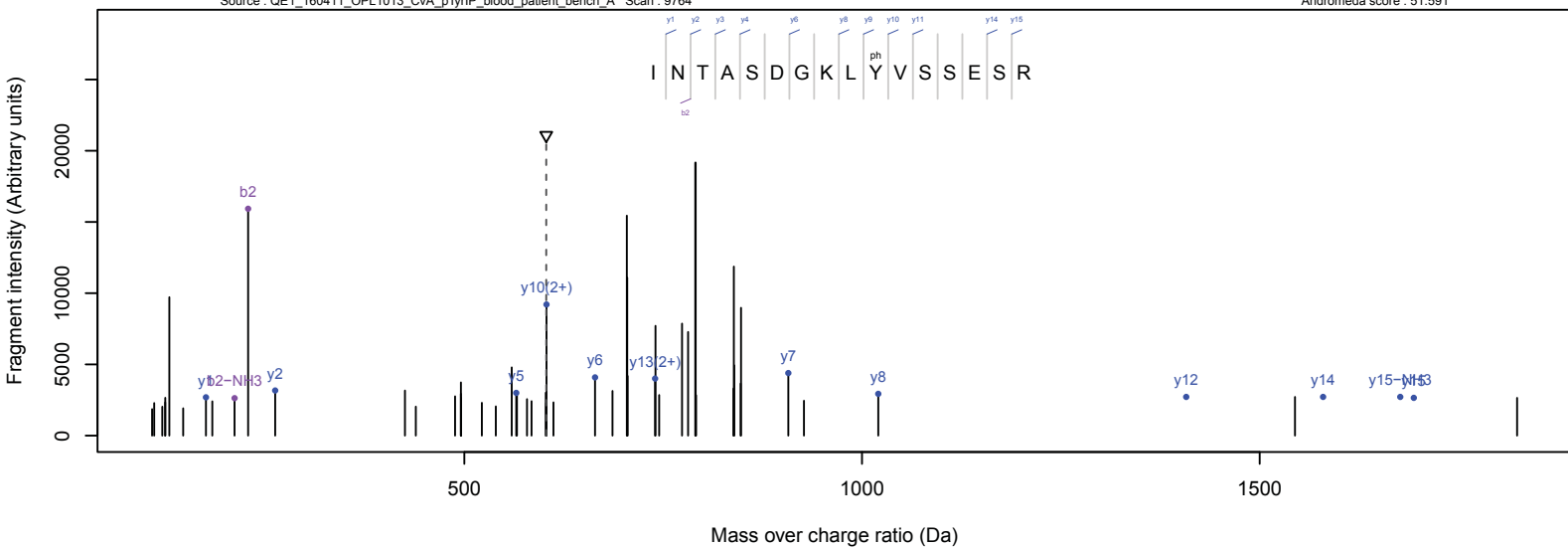
Andromeda score : 51.727



## INTASDGKLY(ph)VSESR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 9764

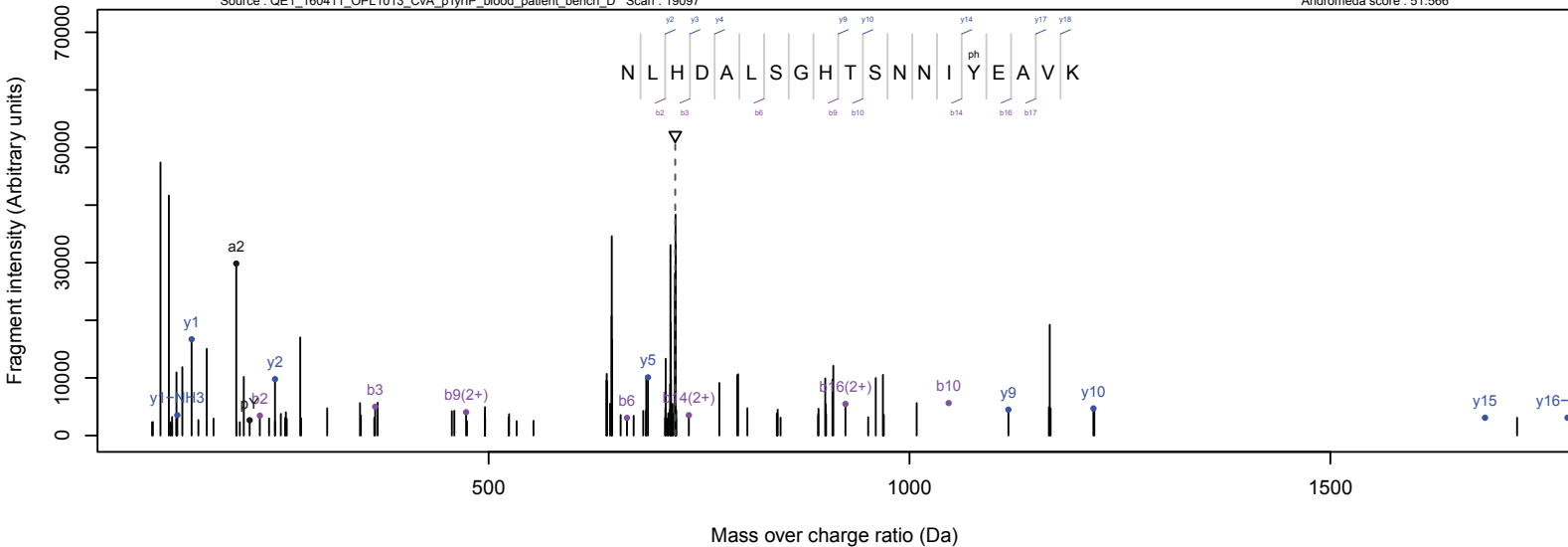
Andromeda score: 51.591



## NLHDALSGHTSNII(ph)EAVK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 19097

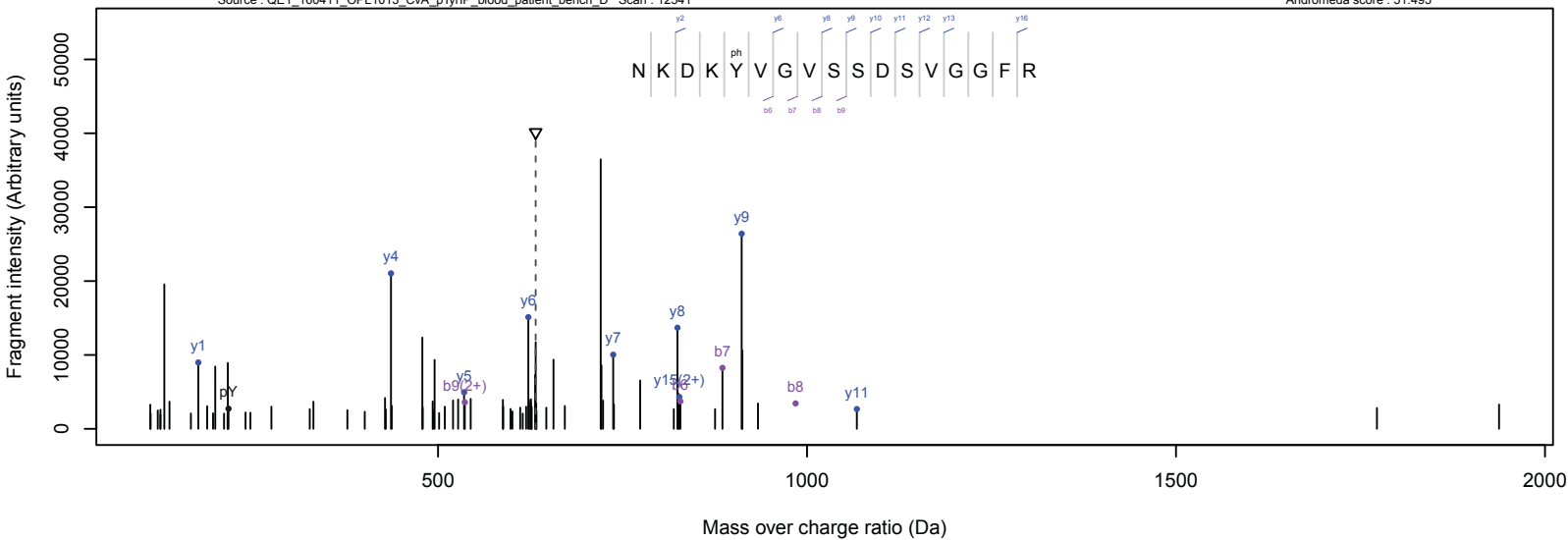
Andromeda score: 51.566



## NKDKY(ph)VGSSDSVGGFR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 12541

Andromeda score: 51.495

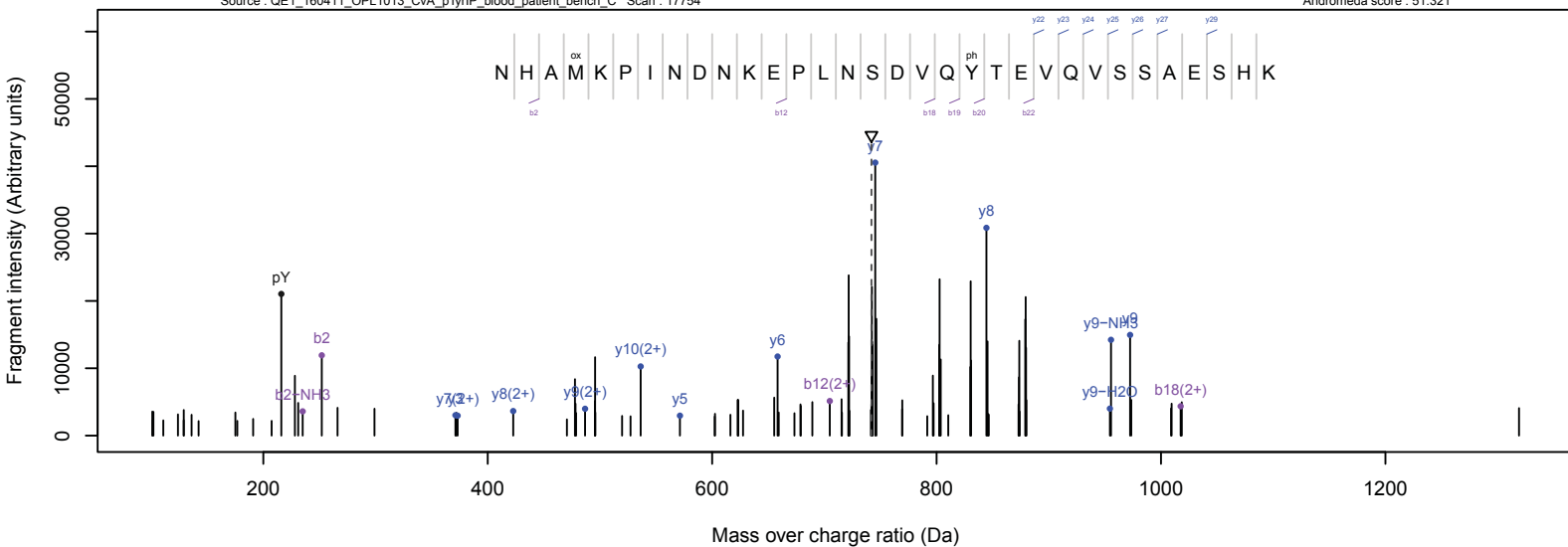




NHAM(ox)KPINDNKEPLNSDVQYTEVQVSSAESHK-5+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 17754

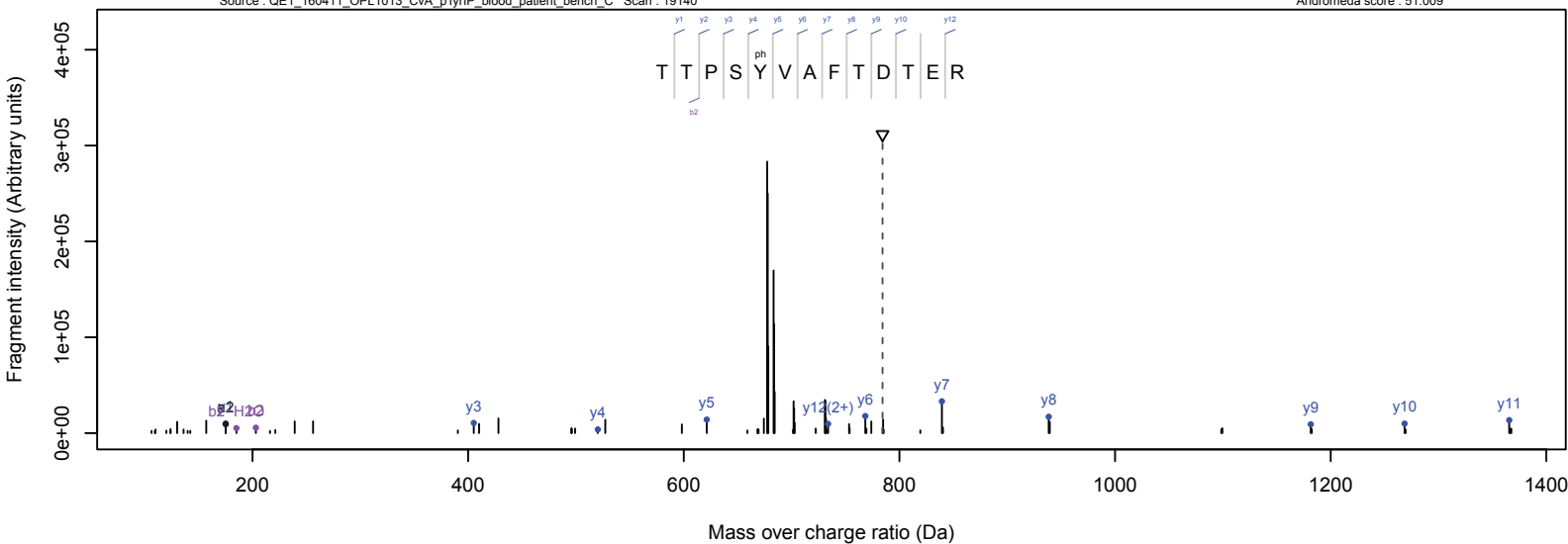
Andromeda score : 51.321



TTPSY(ph)VAFTDTER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 19140

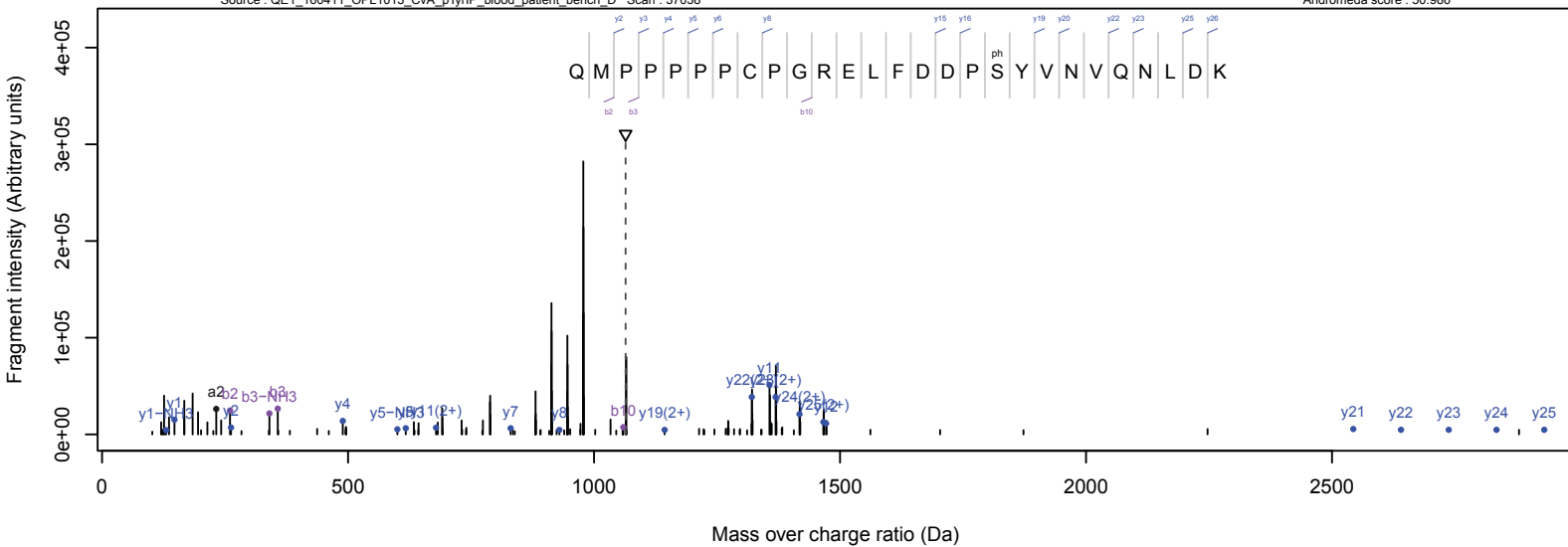
Andromeda score : 51.009



QMPPPPCPGRELFD DP SYVNVQNLDK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 37038

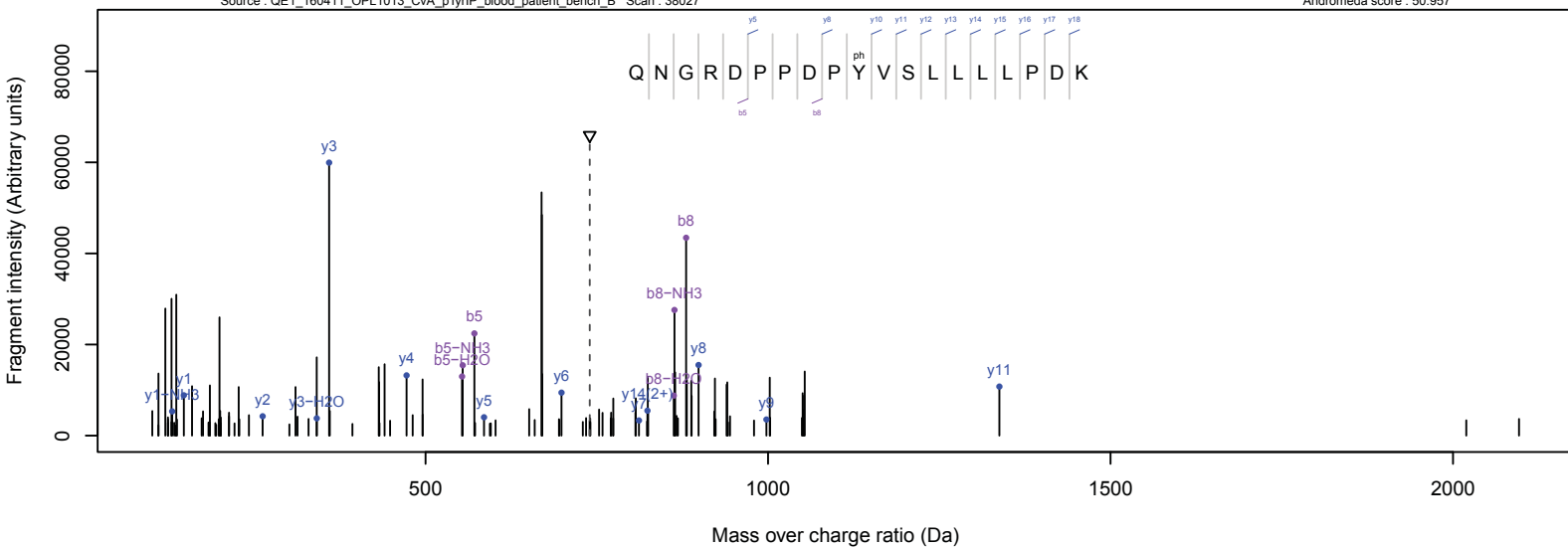
Andromeda score : 50.986



QNGRDPPD<sup>ph</sup>YVSLLLPDK-3+

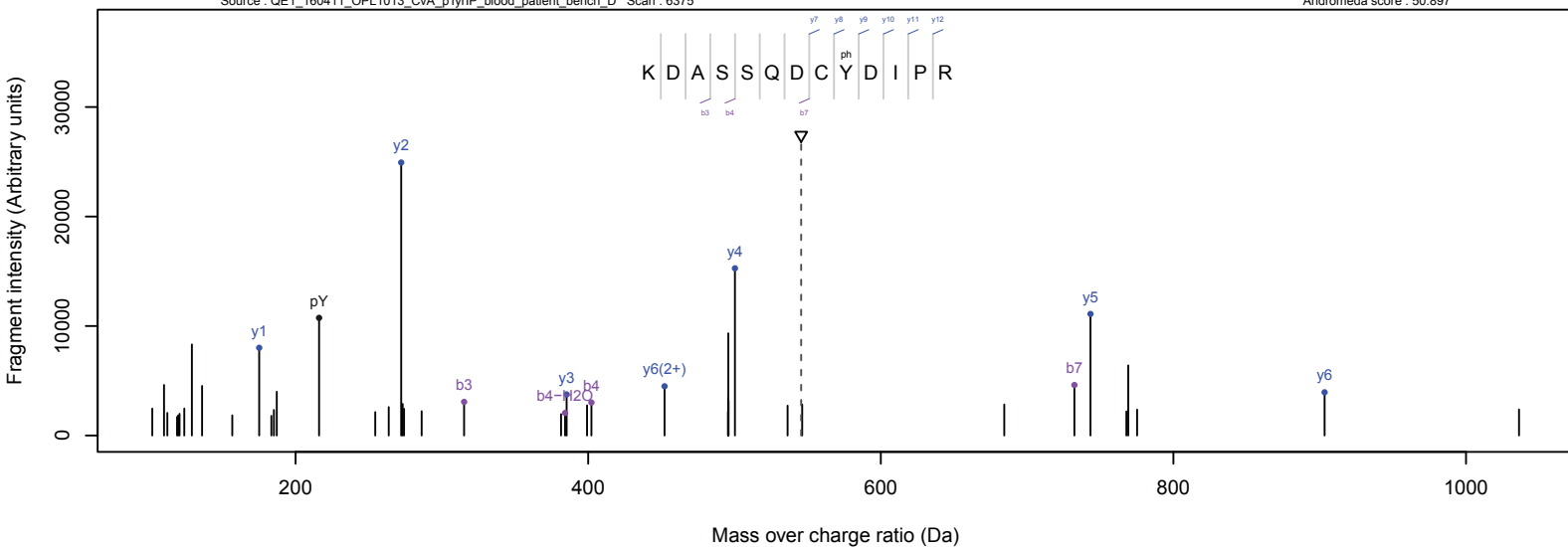
Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 38027

Andromeda score : 50.957

KDASSQDC<sup>ph</sup>YDIPR-3+

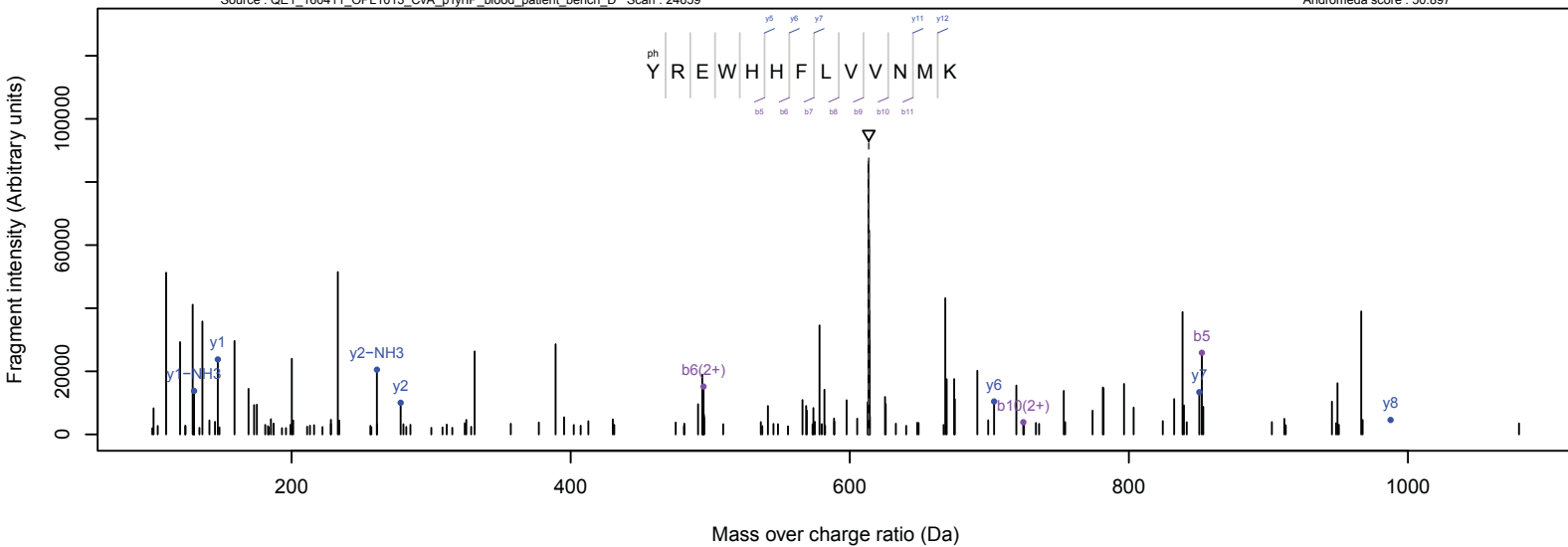
Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 6375

Andromeda score : 50.897

Y<sup>ph</sup>REWHHFLVNMK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24659

Andromeda score : 50.897

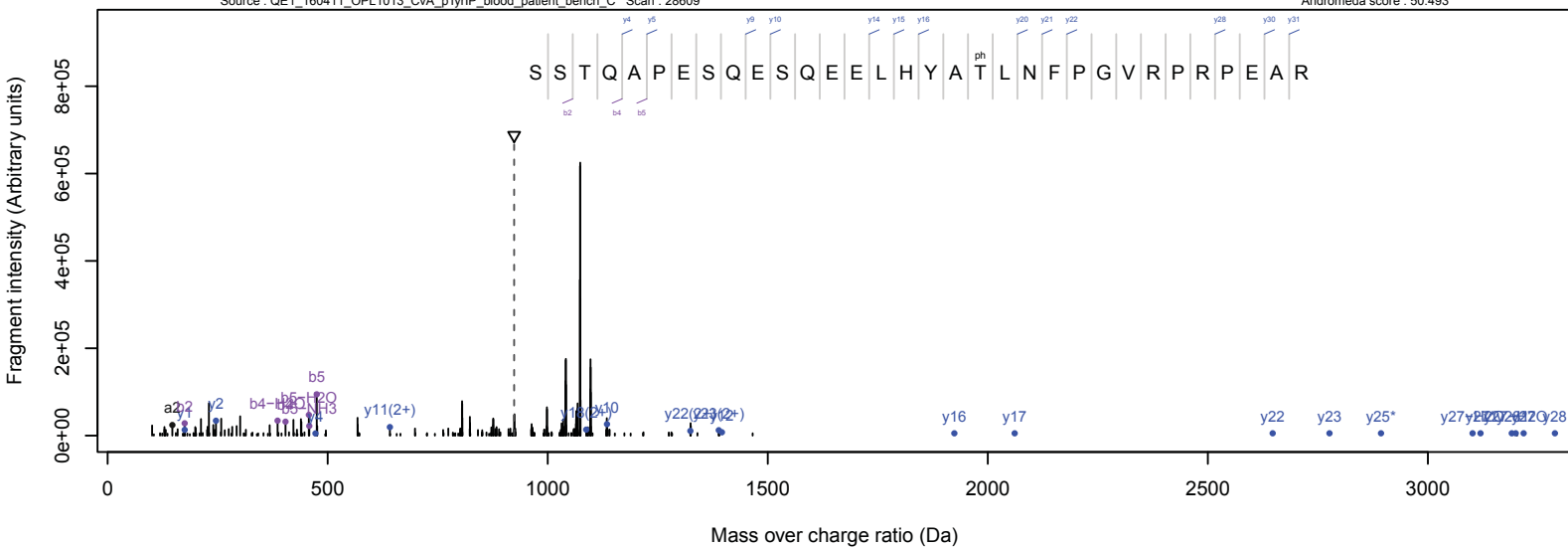




SSTQAPESQESQEELHYAT(ph)LNFGVVRPPEAR-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28609

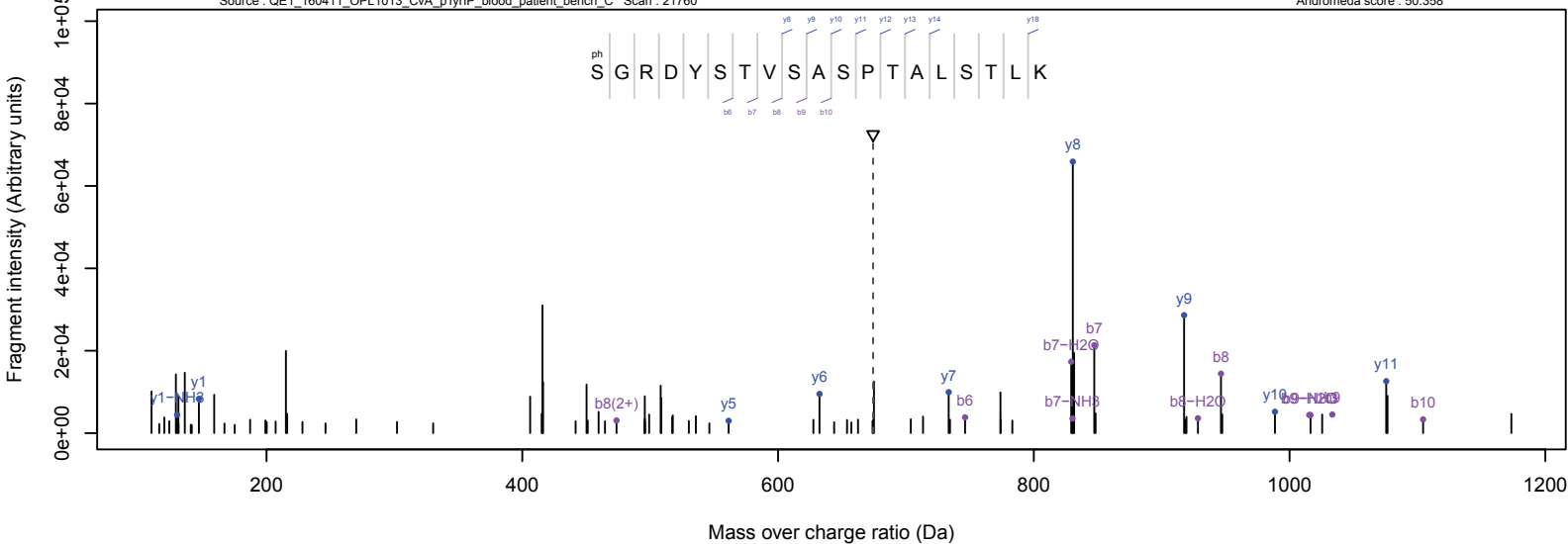
Andromeda score : 50.493



S(ph)GRDYSTVSASPTALSTLK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 21760

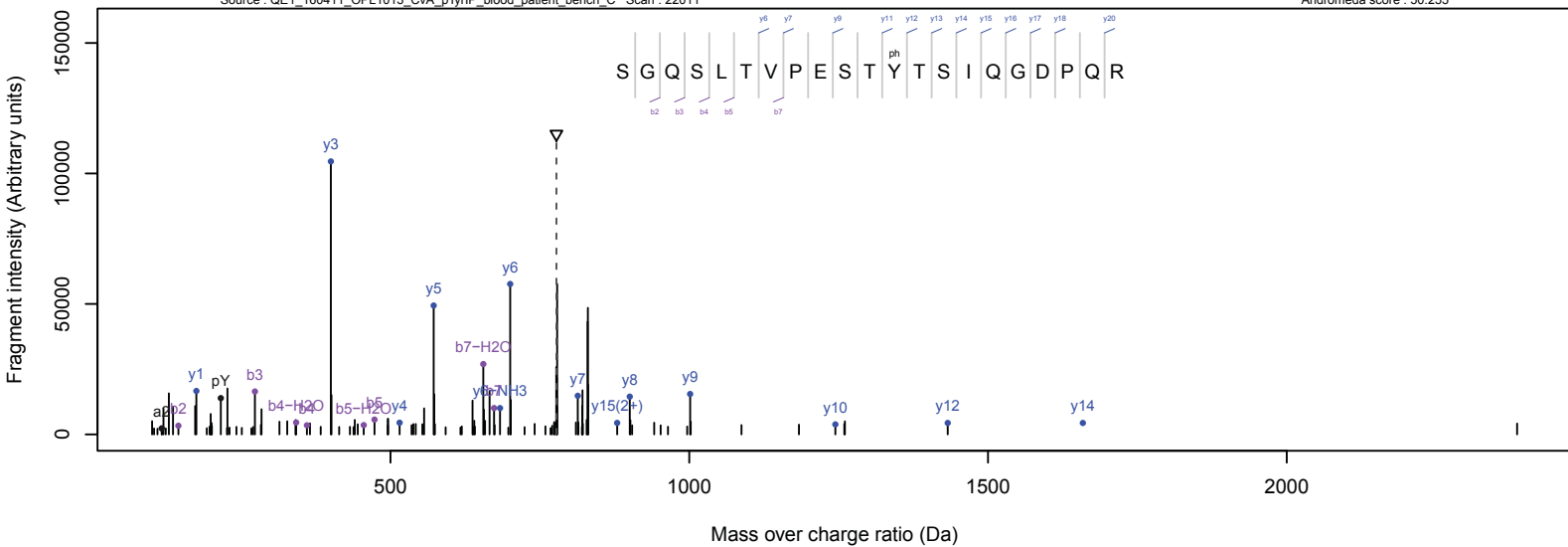
Andromeda score : 50.358



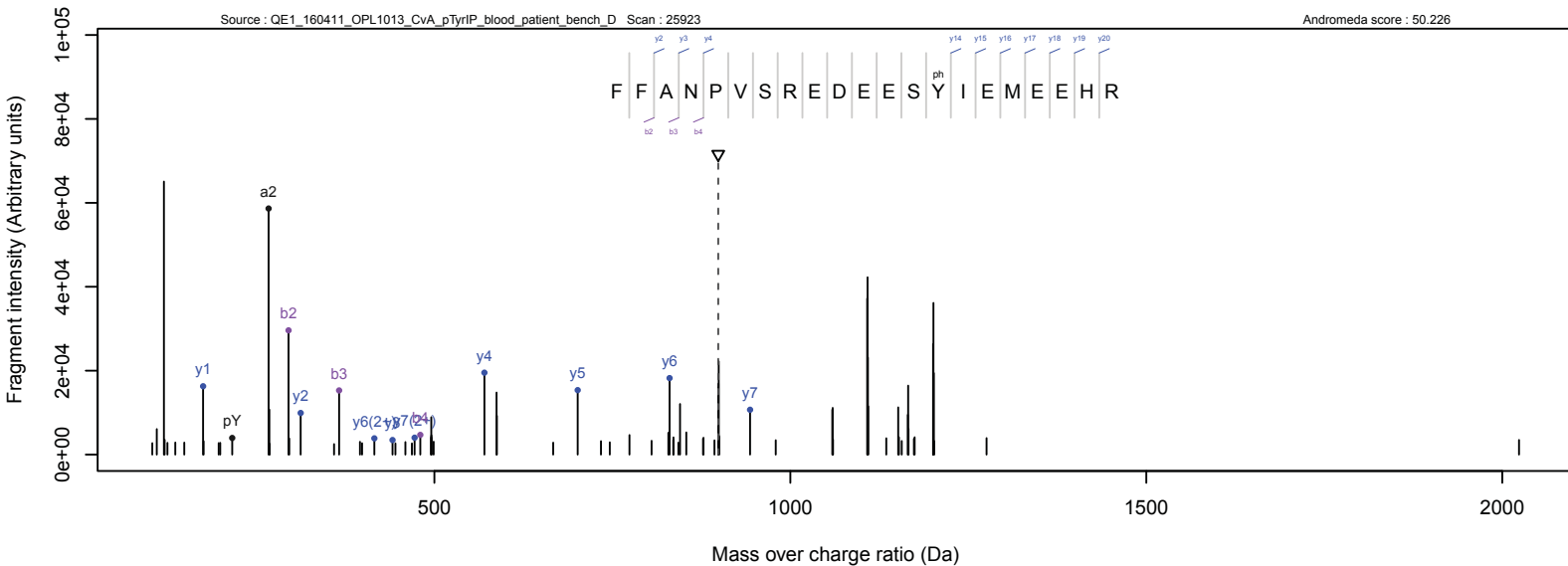
SGQLTVPESTY(ph)TSIQGDPQR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 22011

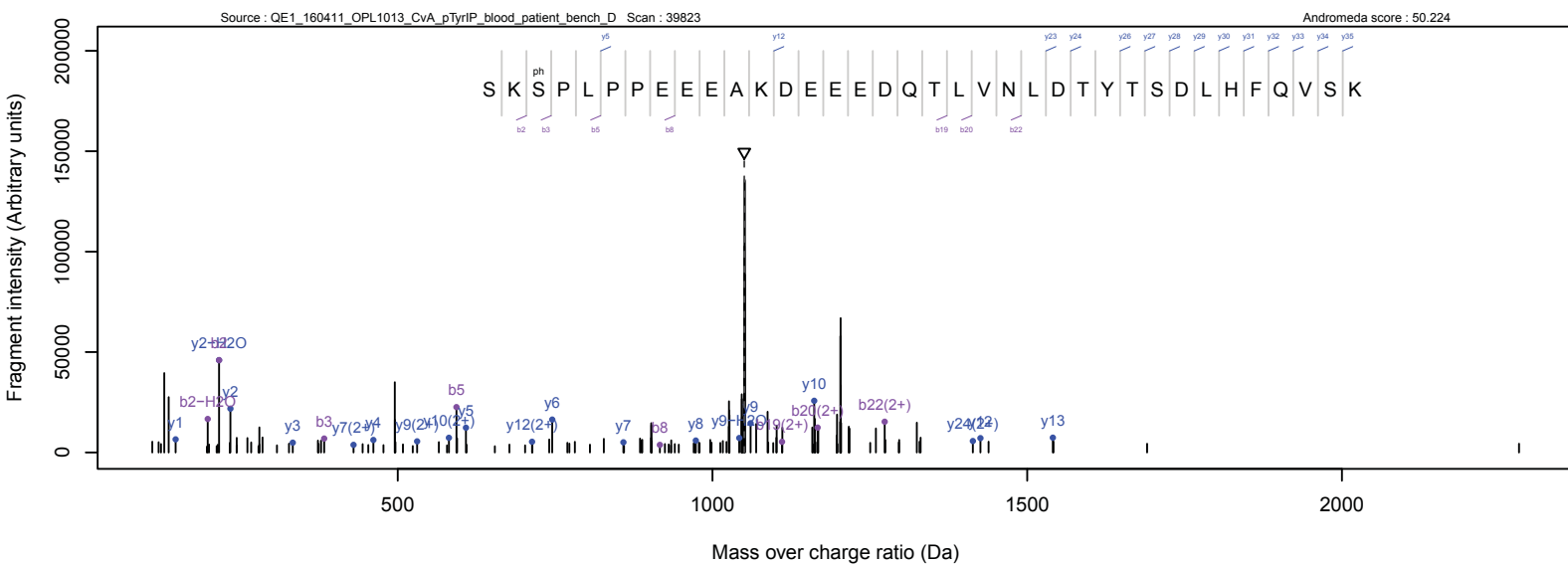
Andromeda score : 50.255



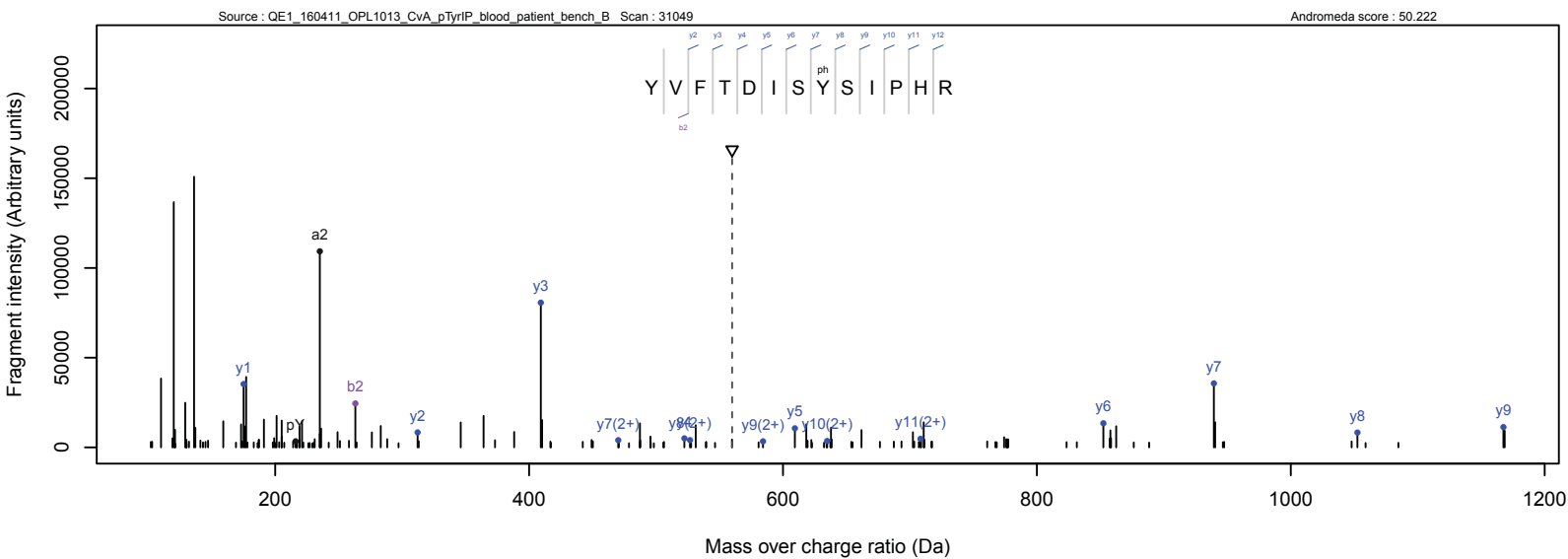
**FFANPVSREDEESY(ph)IEMEEHR-3+**



**SKS(ph)PLPPEEEAKDEEEDQTLVNLDTYTSDDLHFQVSK-4+**



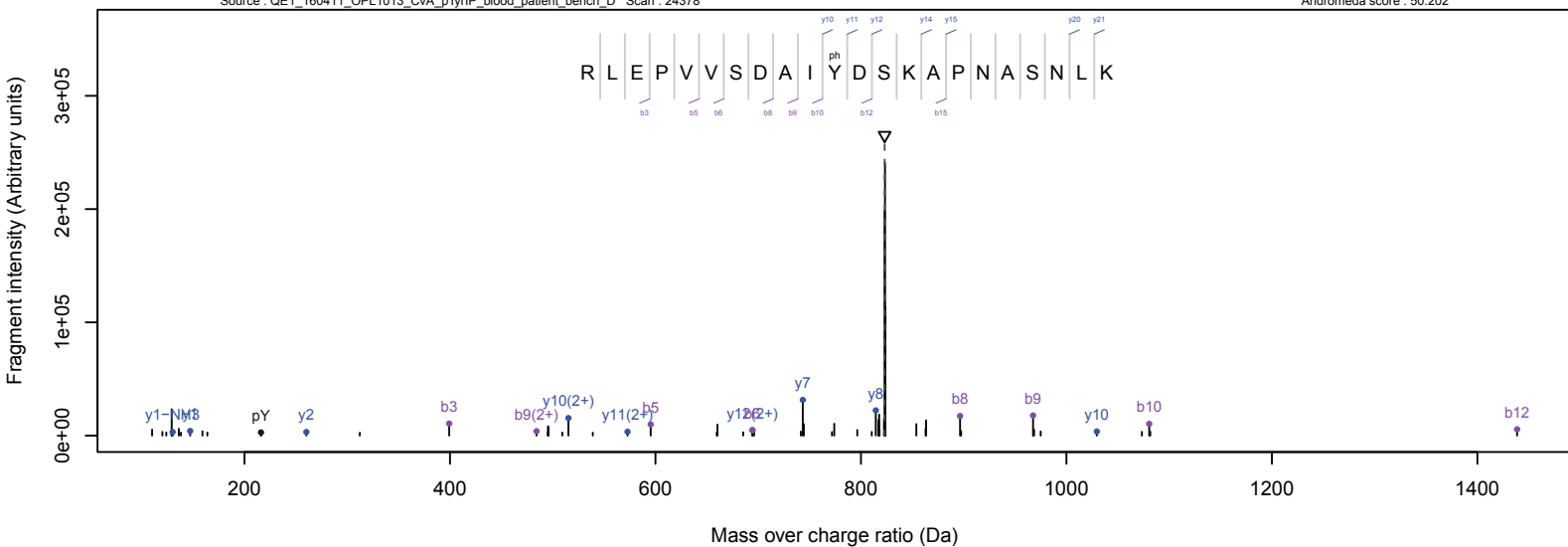
**YVFTDISY(ph)SIPHR-3+**



## RLEPVVSDAIY(ph)DSKAPNASNLK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 24378

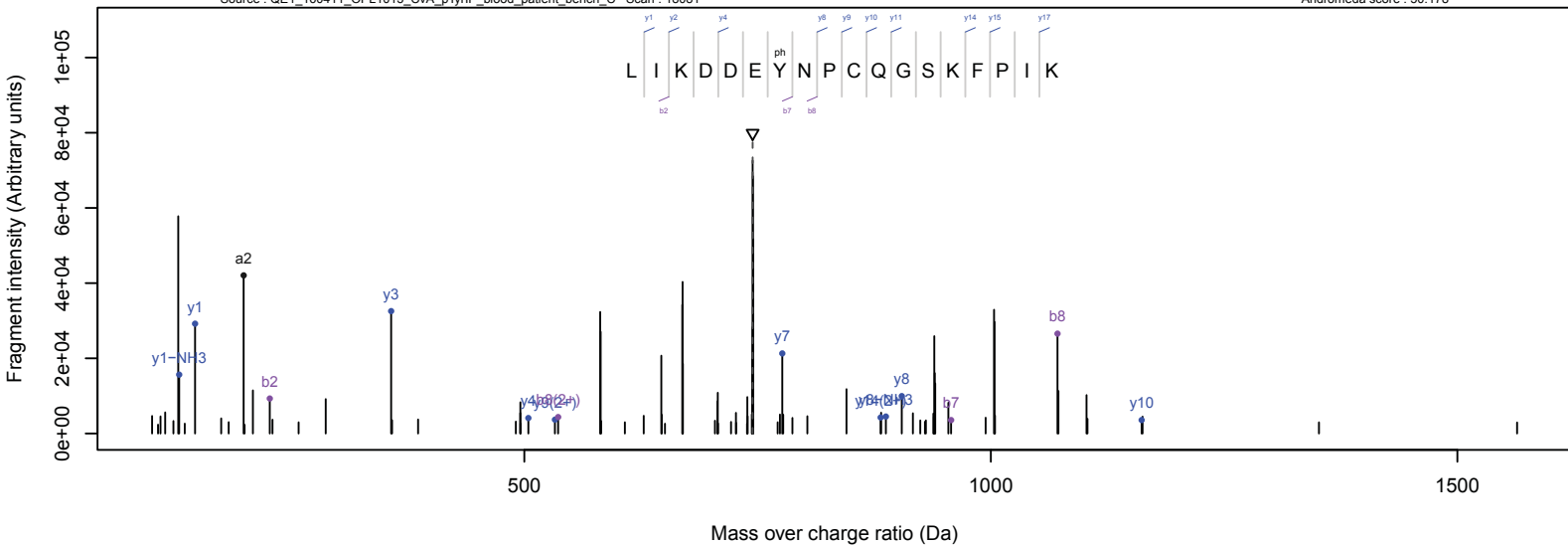
Andromeda score : 50.202



## LIKDDEY(ph)NPCQGSKFPIK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18081

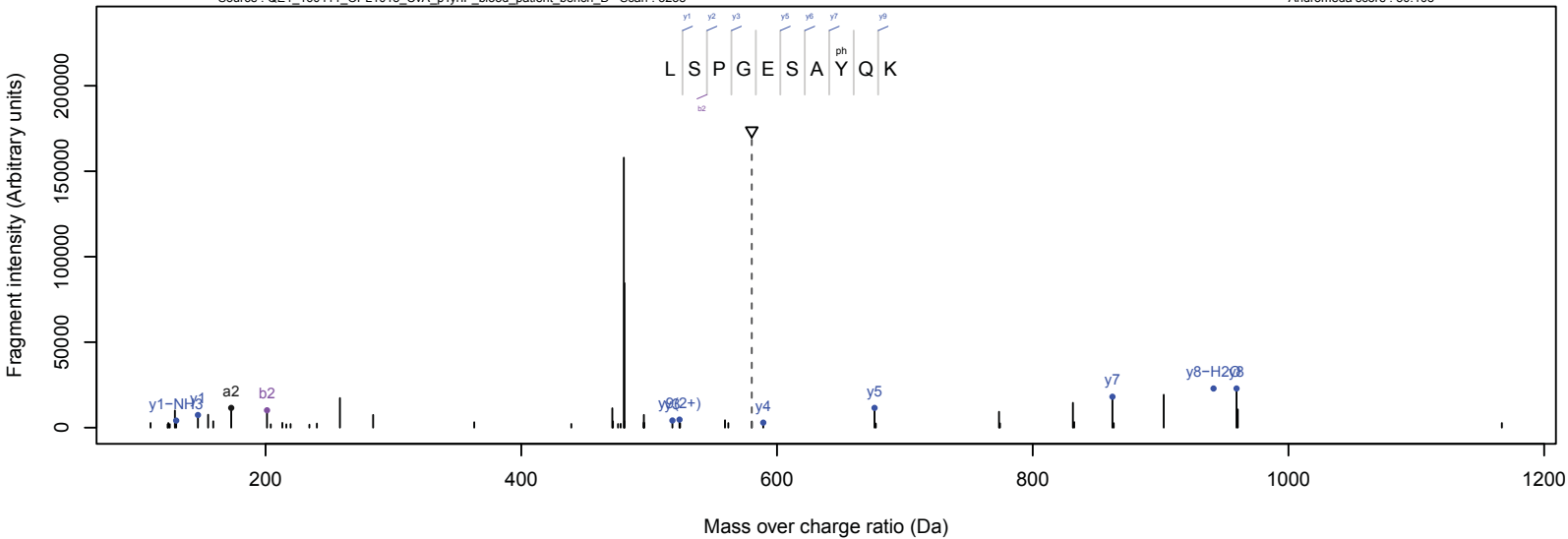
Andromeda score : 50.178



## LSPGESAY(ph)QK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 3238

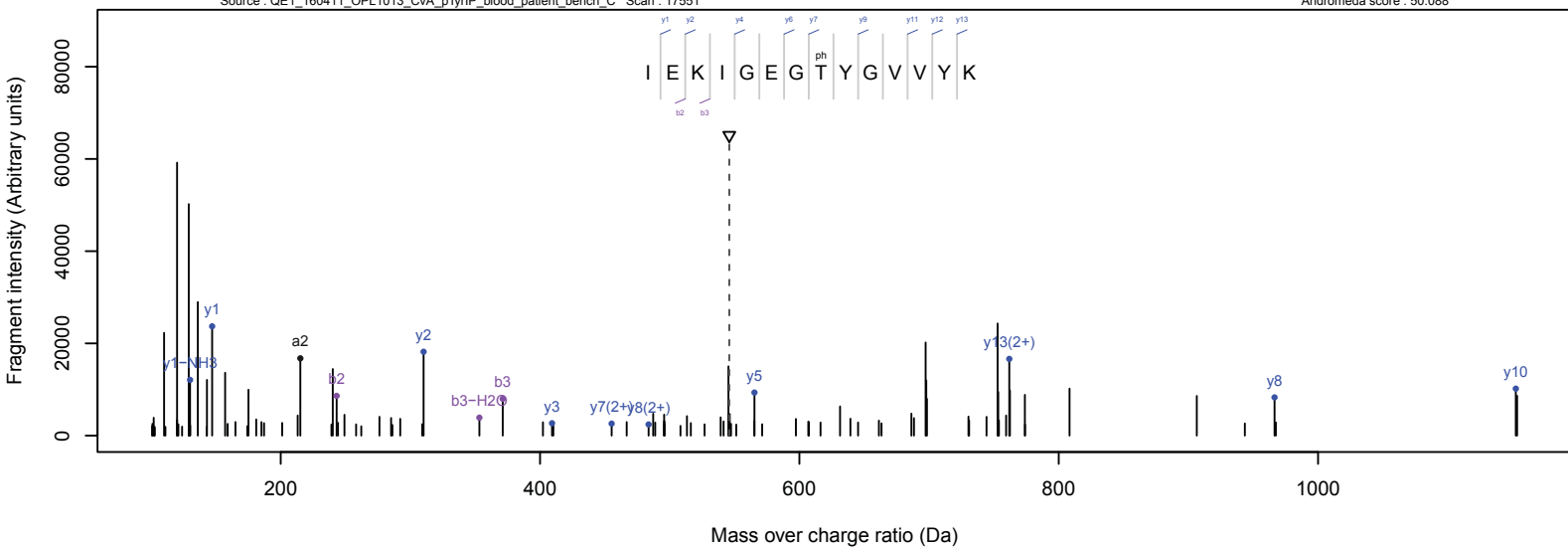
Andromeda score : 50.108



## IEKIGEGT(ph)YGVVYK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 17551

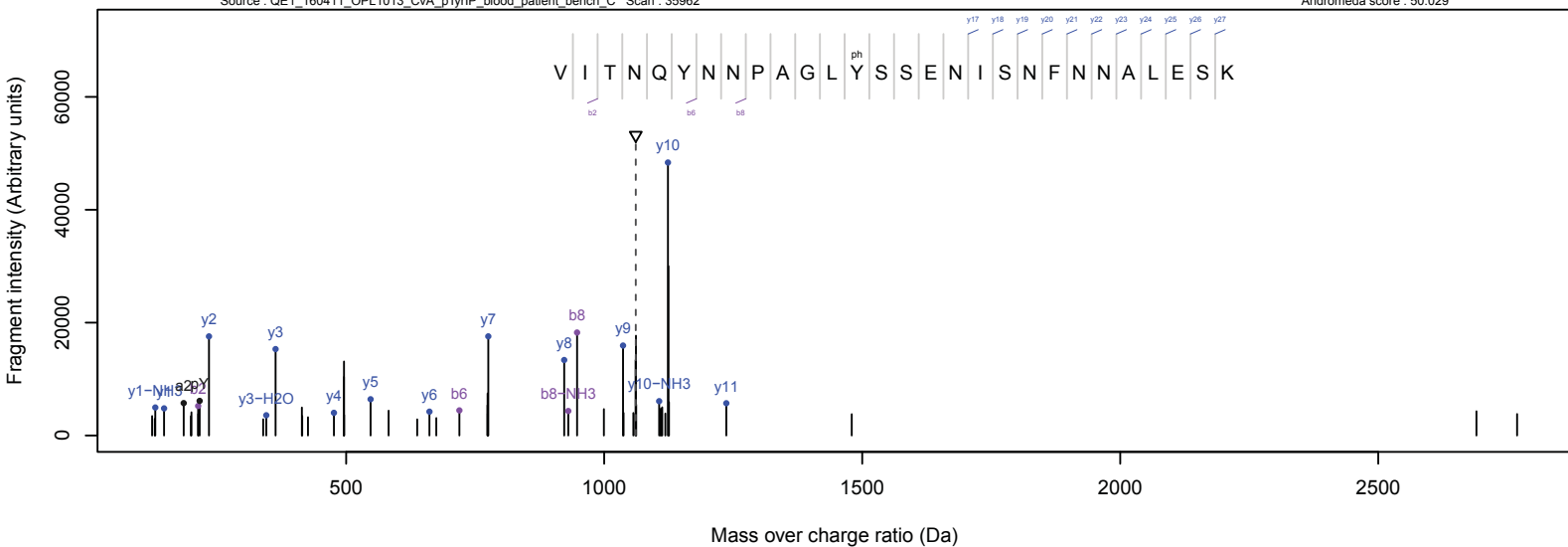
Andromeda score : 50.088



## VITNQYNNPAGLY(ph)SSENI SNFNNALESK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 35962

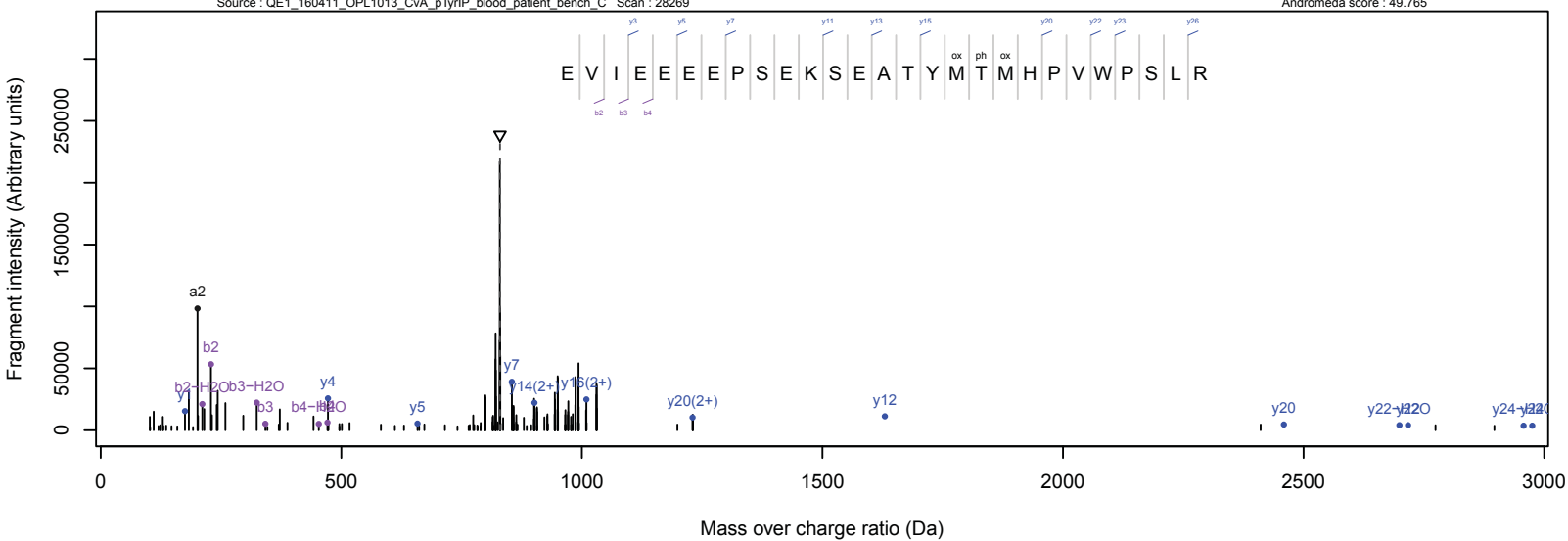
Andromeda score : 50.029



## EVIEEEPSEKSEATY M T M H P V W P S L R-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 28269

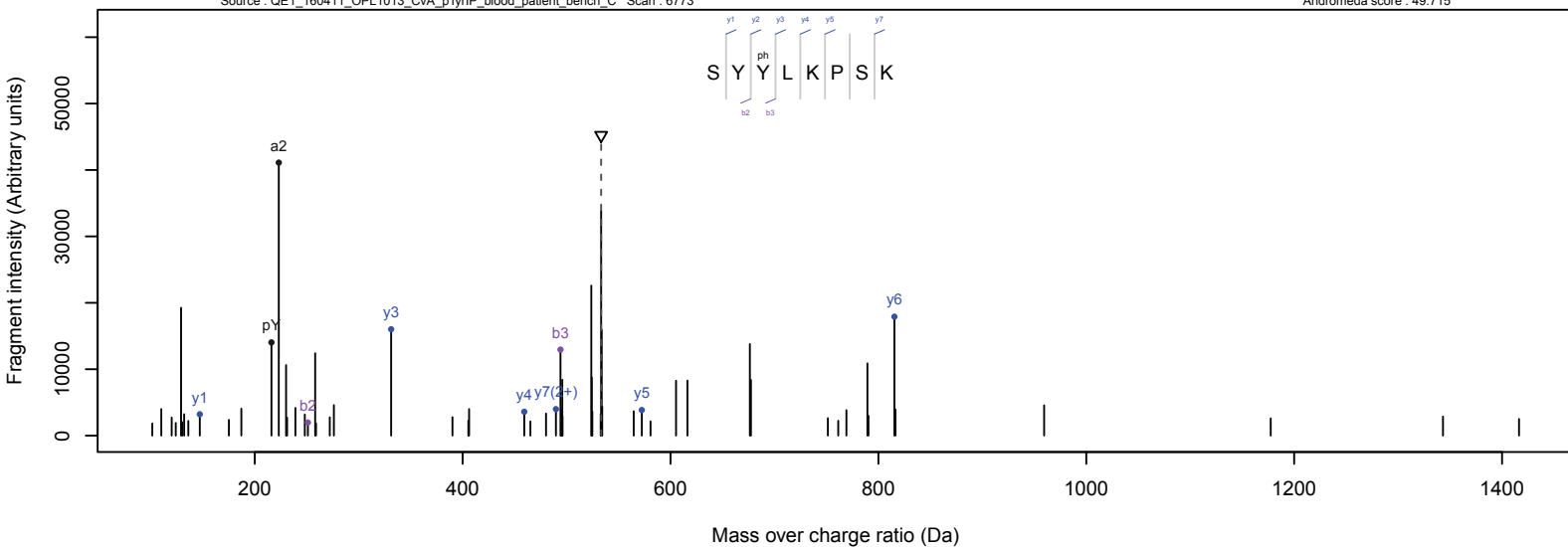
Andromeda score : 49.765



SY<sup>(ph)</sup>Y<sup>(ph)</sup>LKPSK-2+

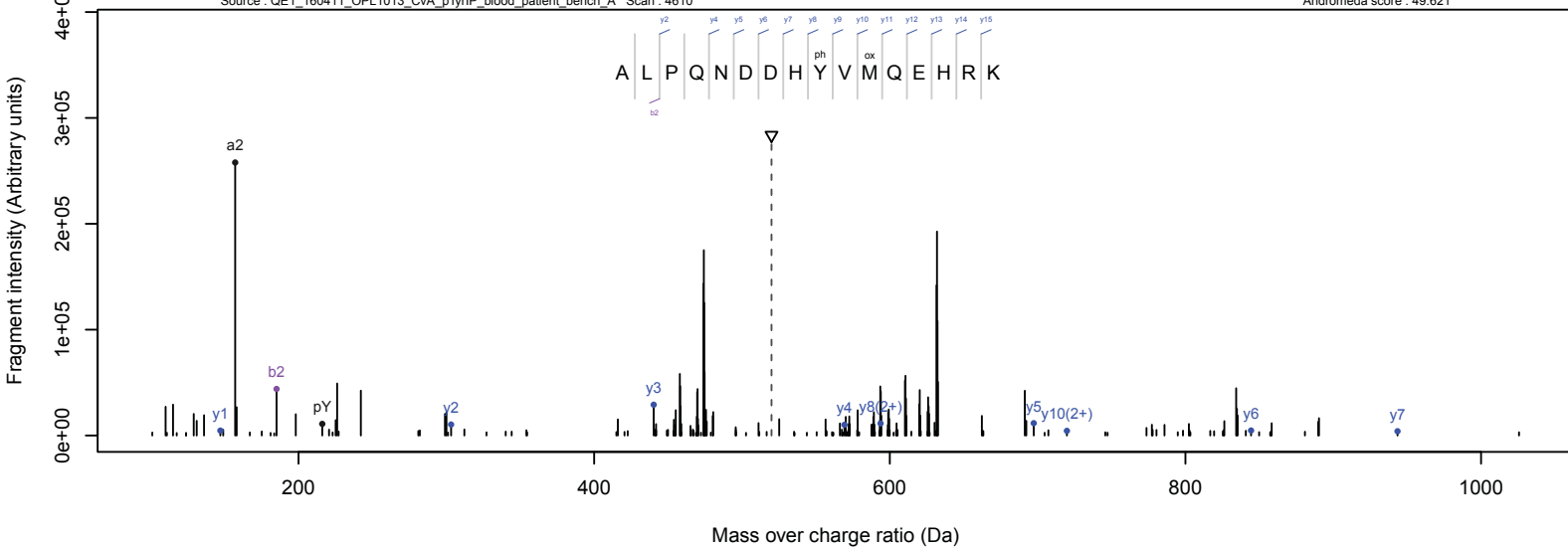
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 6773

Andromeda score: 49.715

ALPQNDDHY<sup>(ph)</sup>VM<sup>(ox)</sup>QEHRK-4+

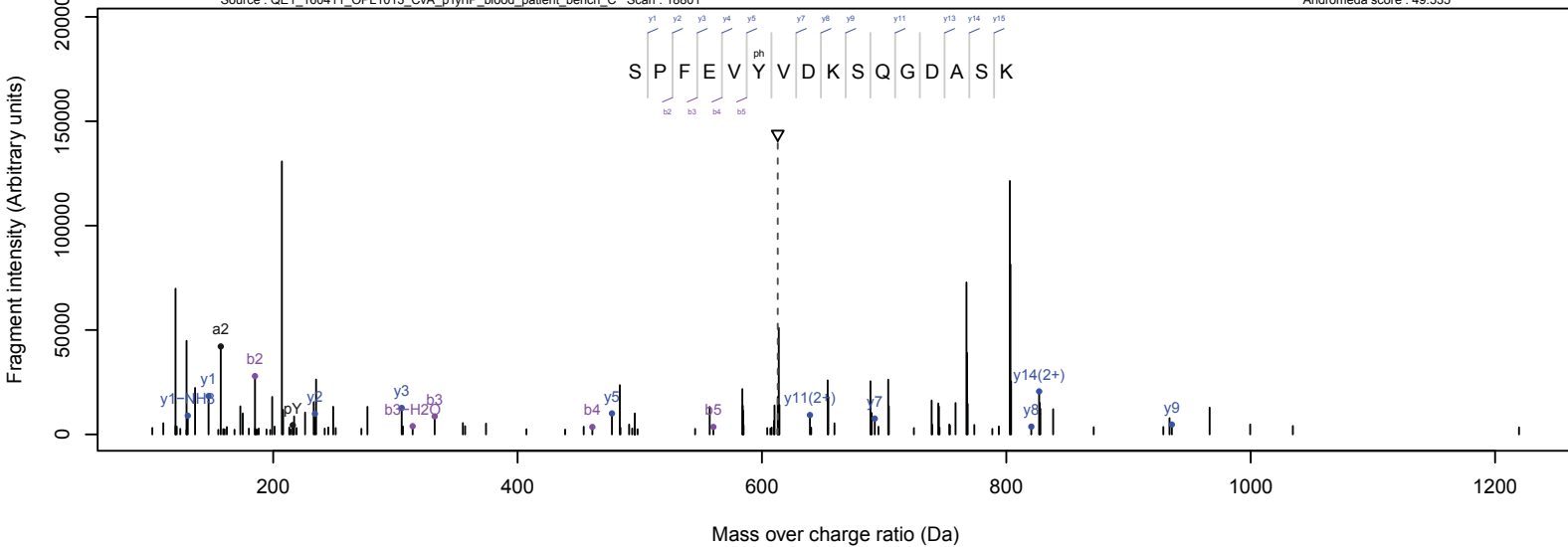
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 4610

Andromeda score: 49.621

SPFEVY<sup>(ph)</sup>VDKSQGDASK-3+

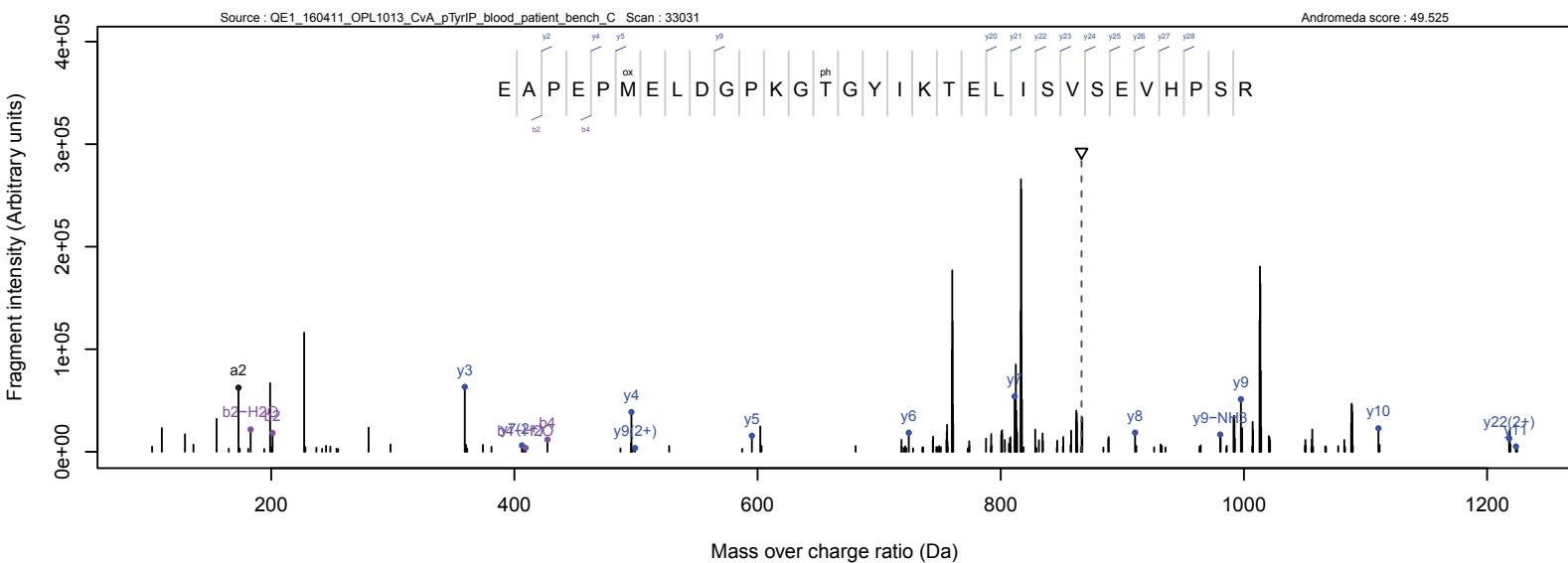
Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 18801

Andromeda score: 49.535

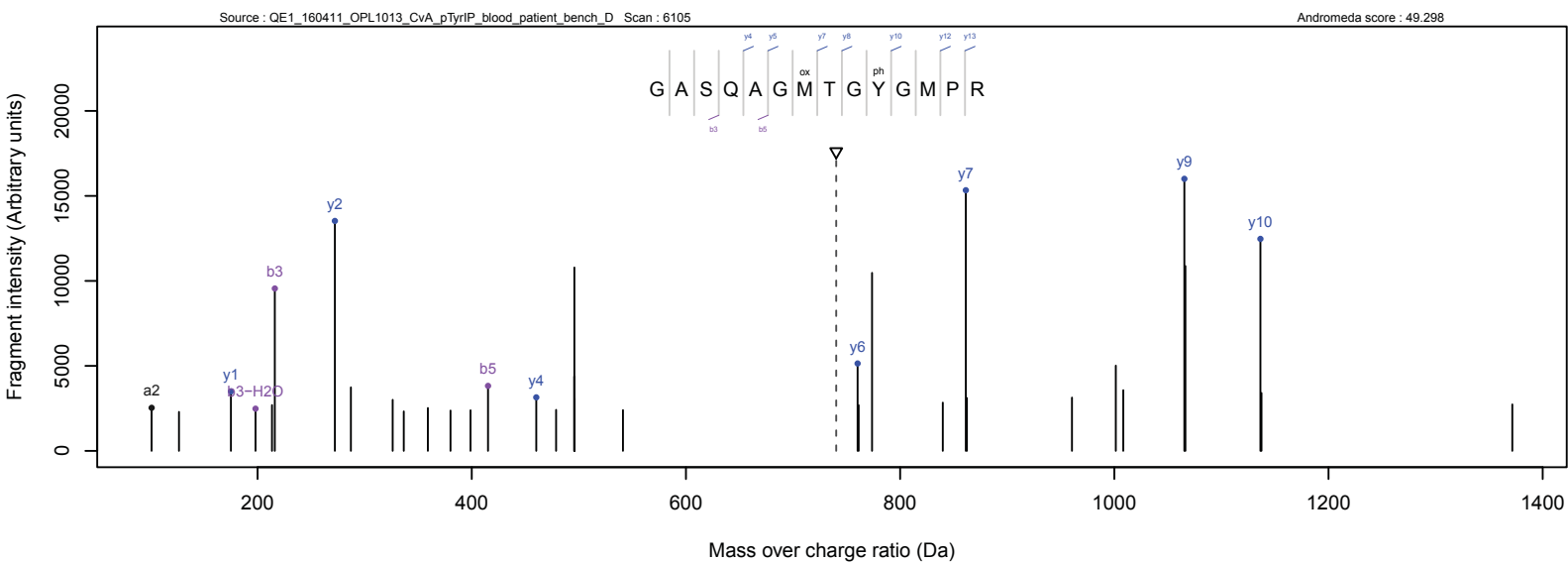




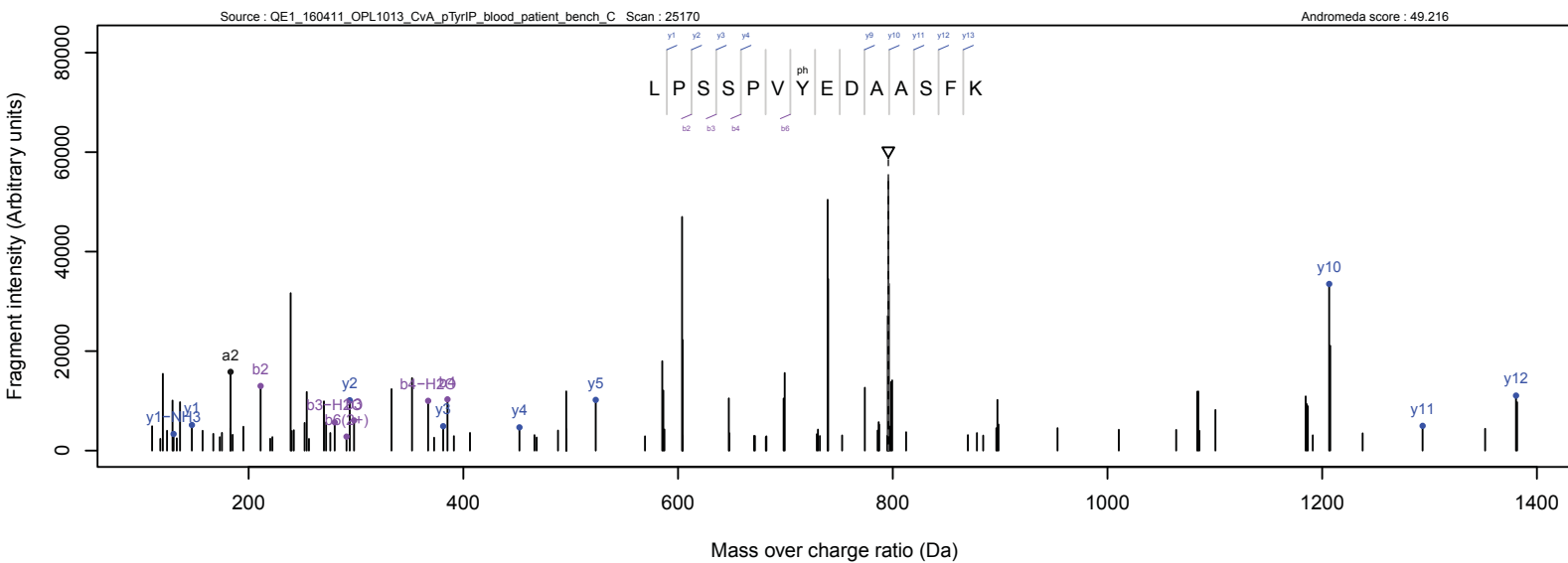
**EAPEPM(ox)ELDGPKG(ph)GYIKTELISVSEVHPSR-4+**



**GASQAGM(ox)TGY(ph)GMPR-2+**



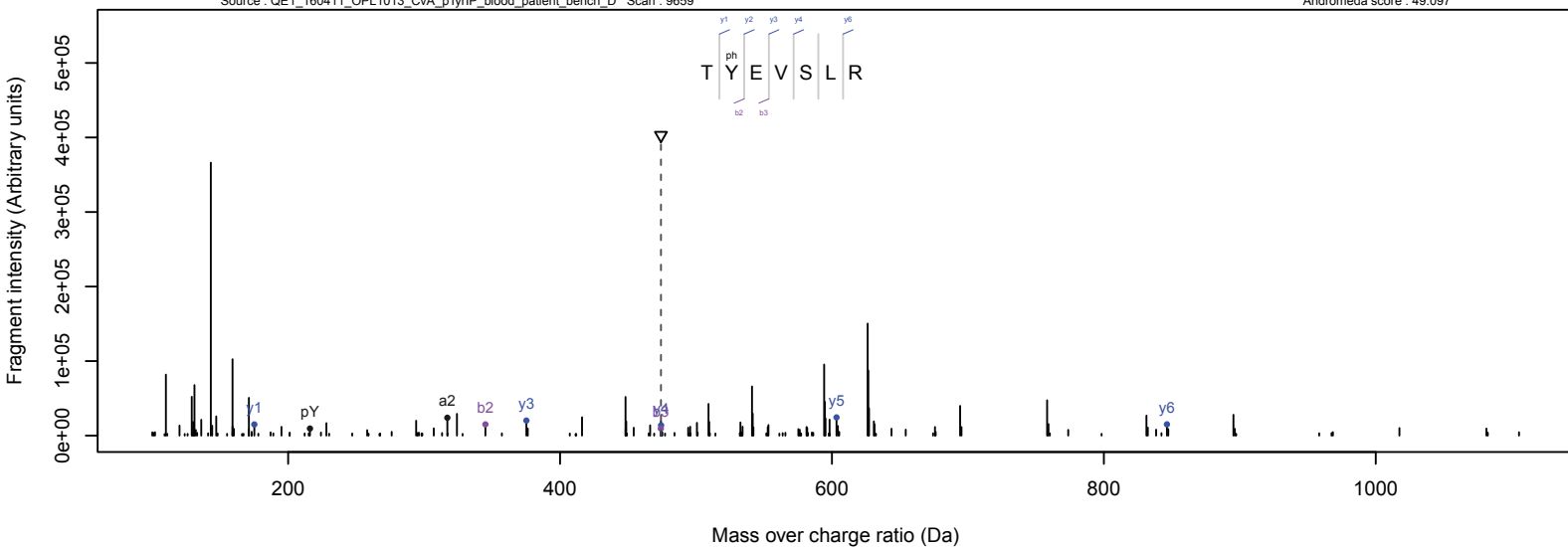
**LPSSPVY(ph)EDAASFK-2+**



## TY(ph)EVSLR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 9659

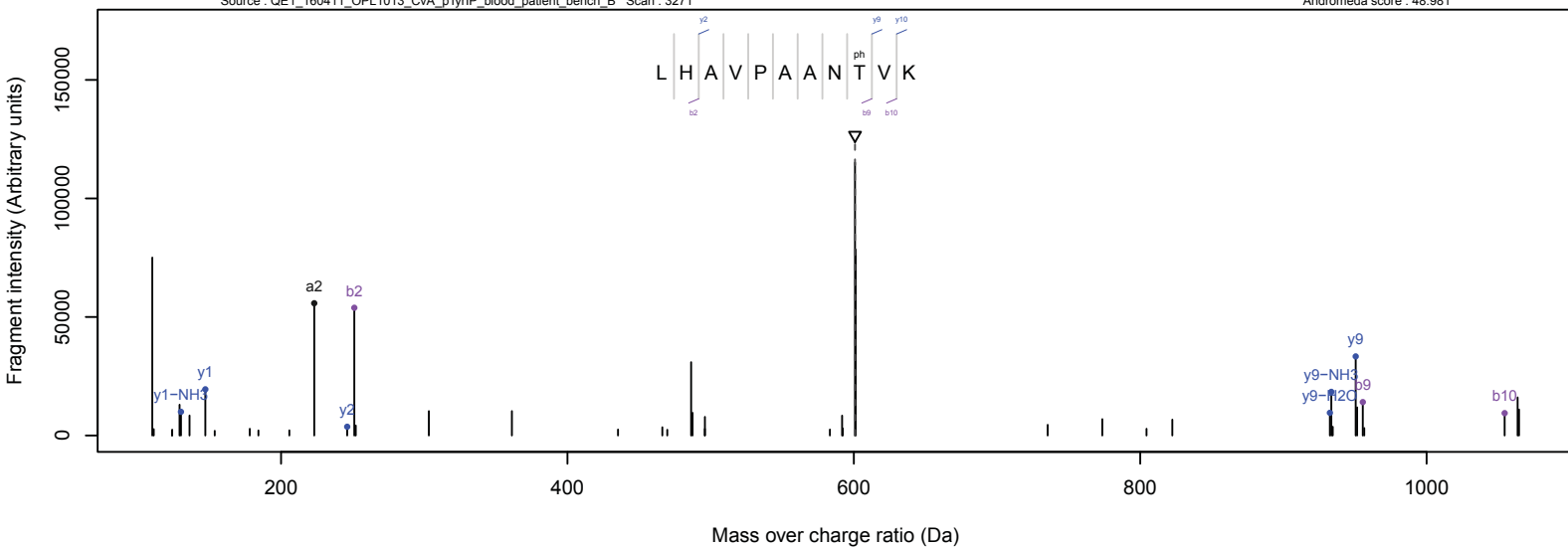
Andromeda score : 49.097



## LHAVPAANT(ph)VK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 3271

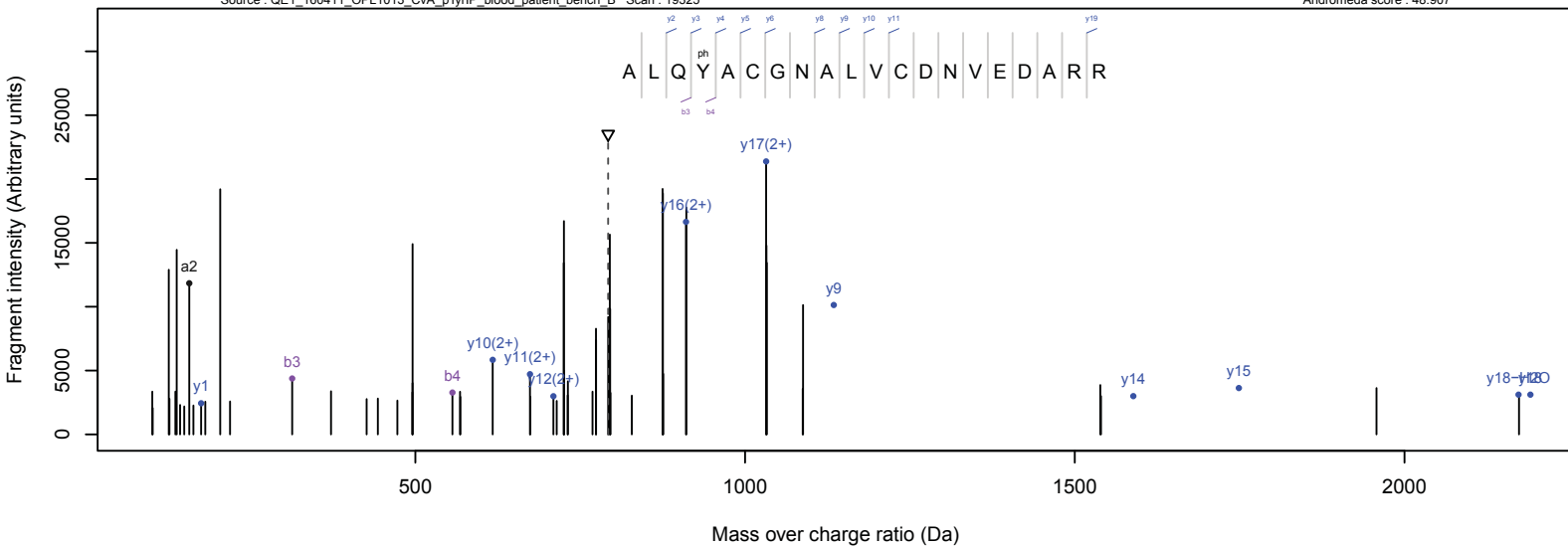
Andromeda score : 48.981



## ALQY(ph)ACGNALVCDNVEDARR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 19325

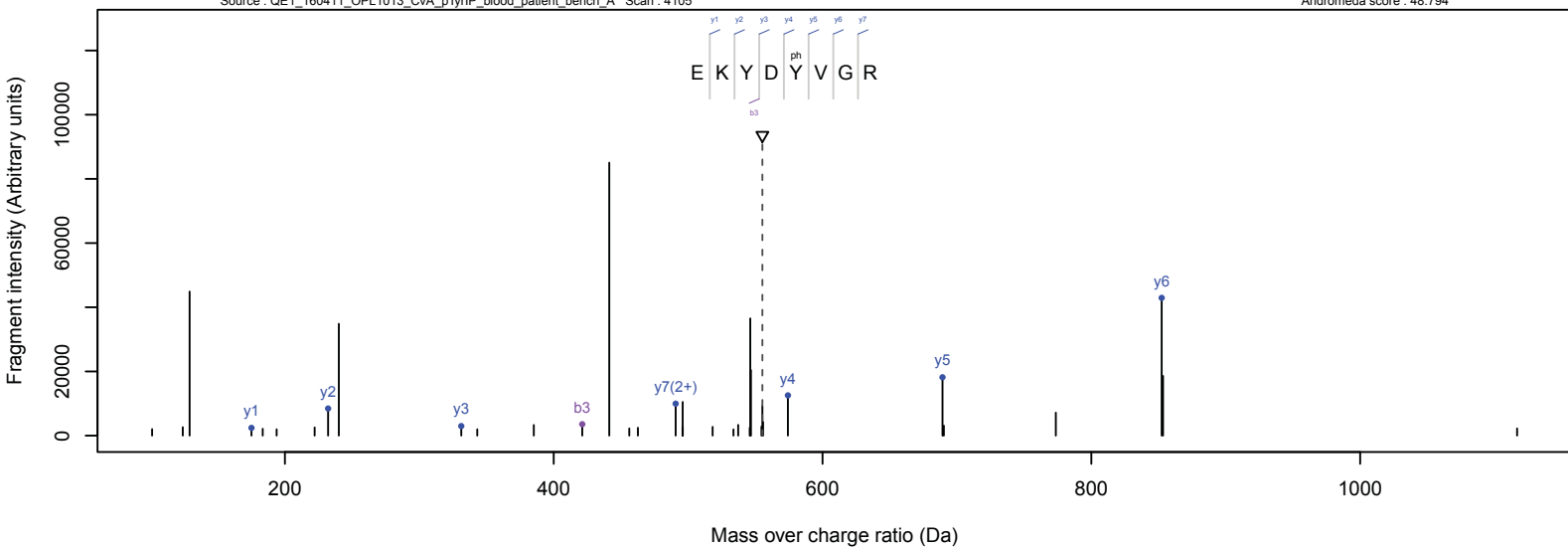
Andromeda score : 48.907



**EKYDY(ph)VGR-2+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 4105

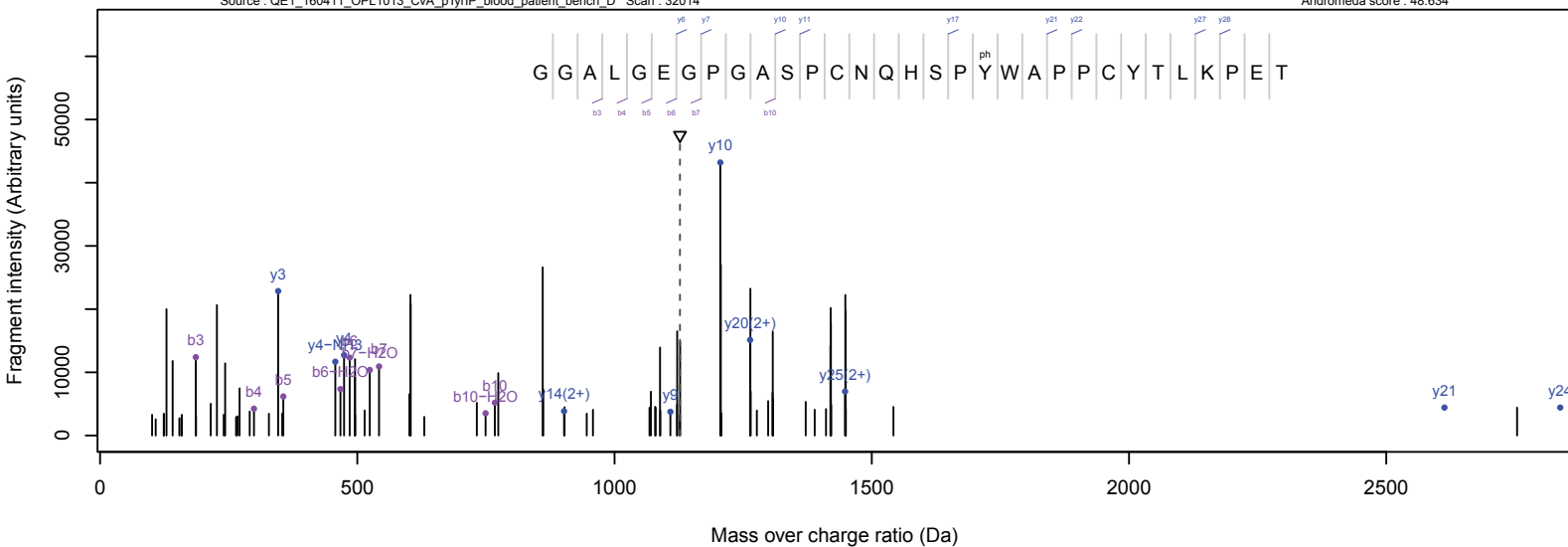
Andromeda score: 48.794



**GGALGEGPGASPCNQHSY(ph)WAPPCYTLKPET-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 32014

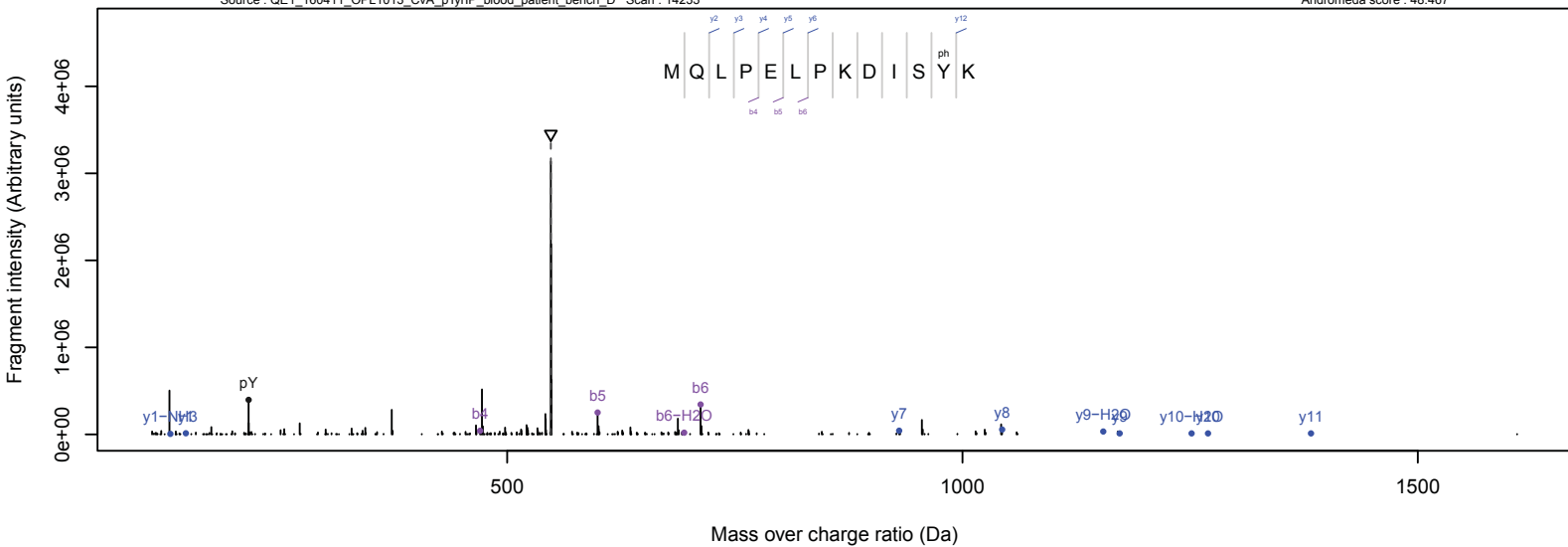
Andromeda score: 48.634



**MQLPELPKDISYK-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 14233

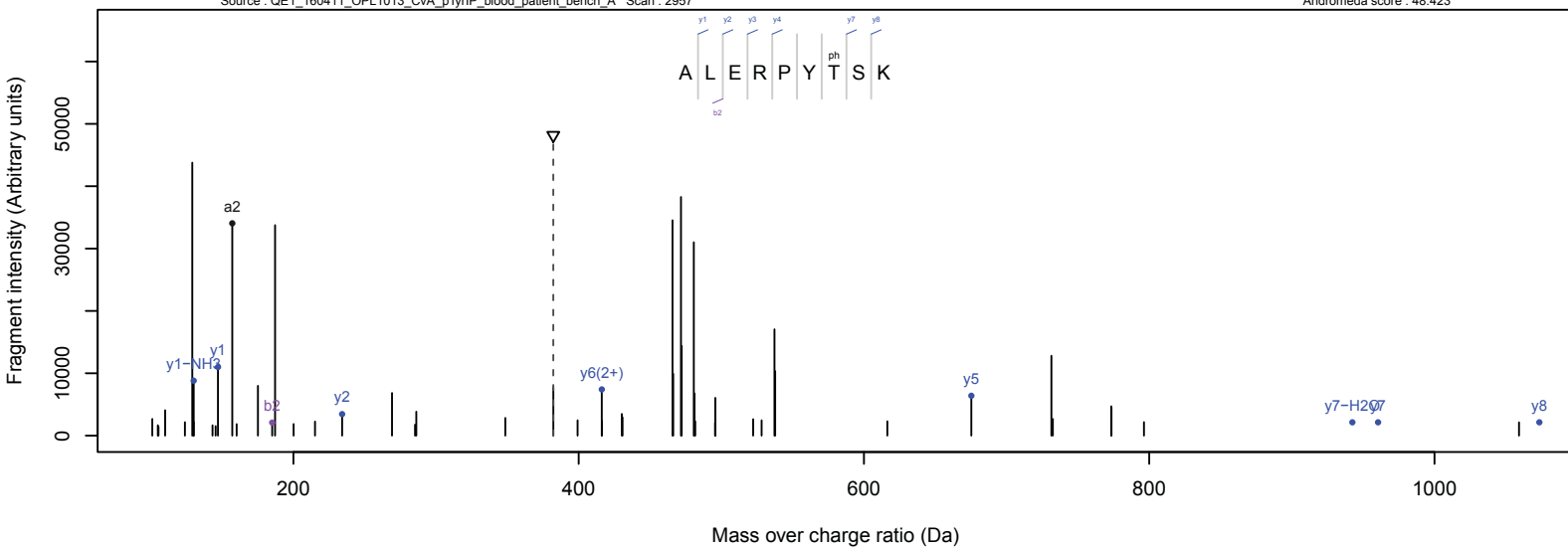
Andromeda score: 48.467



ALERPYP(ph)SK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 2957

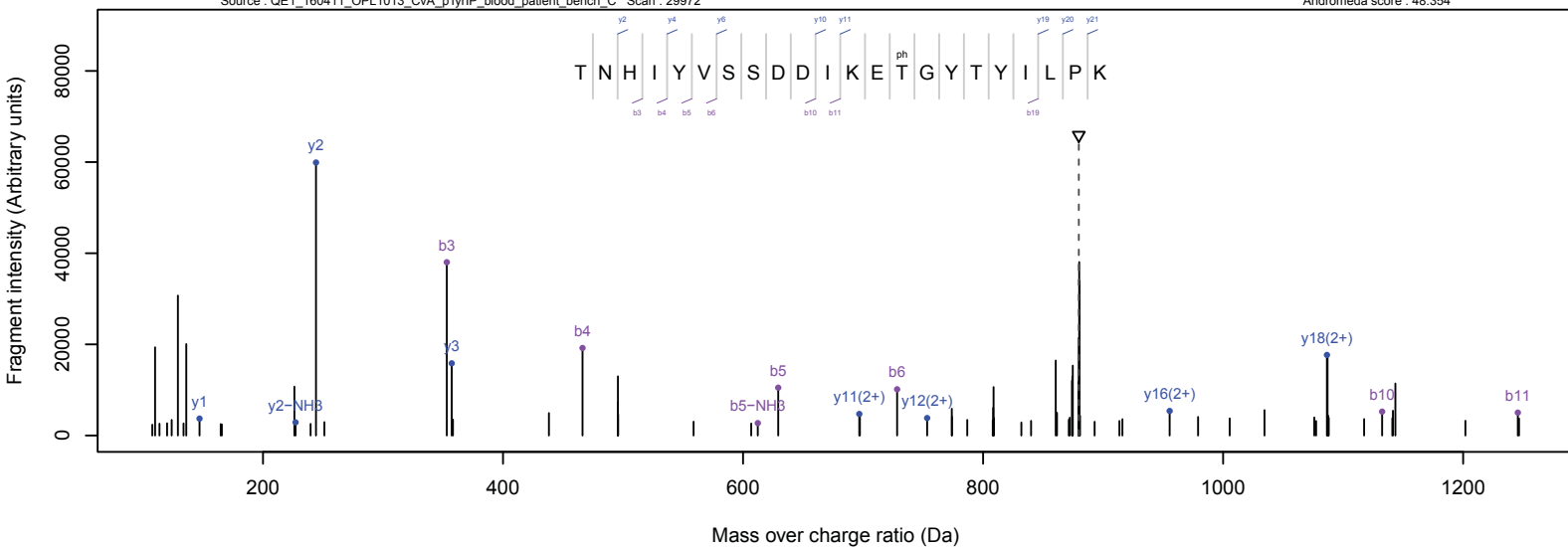
Andromeda score : 48.423



TNHIYVSSDDIKET(ph)GYTYILPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 29972

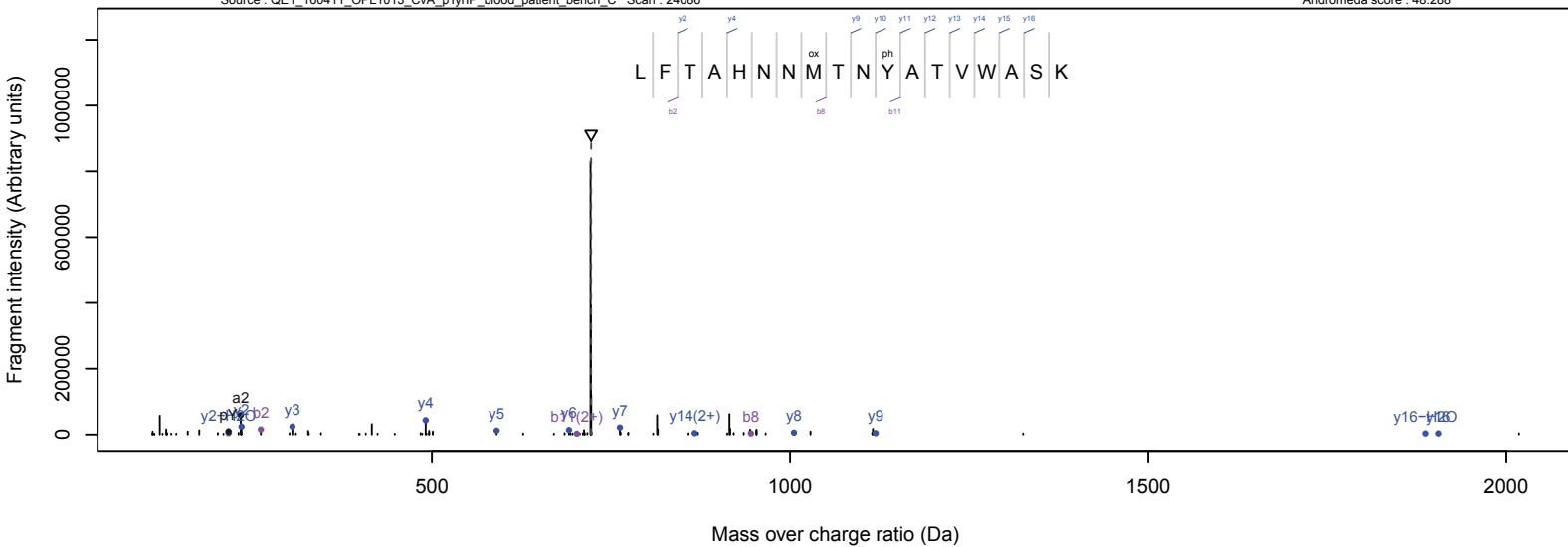
Andromeda score : 48.354



LFTAHHNNM(ph)ATVWASK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 24086

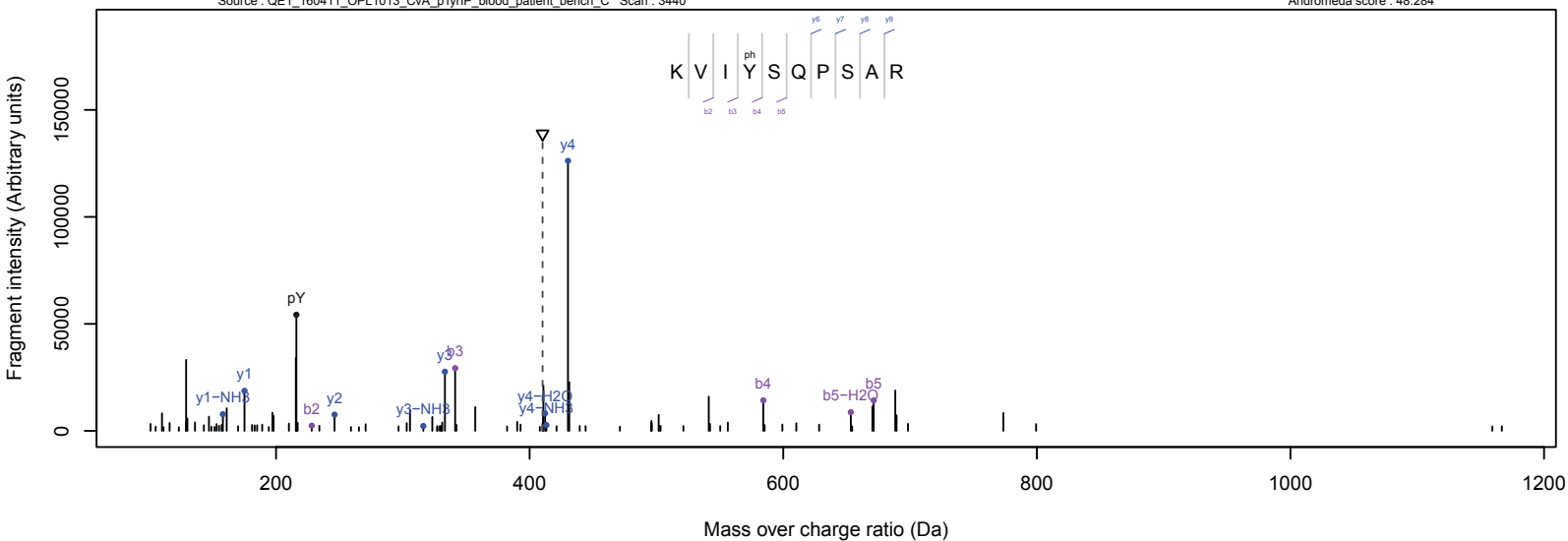
Andromeda score : 48.288



## KVIV(ph)SQPSAR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 3440

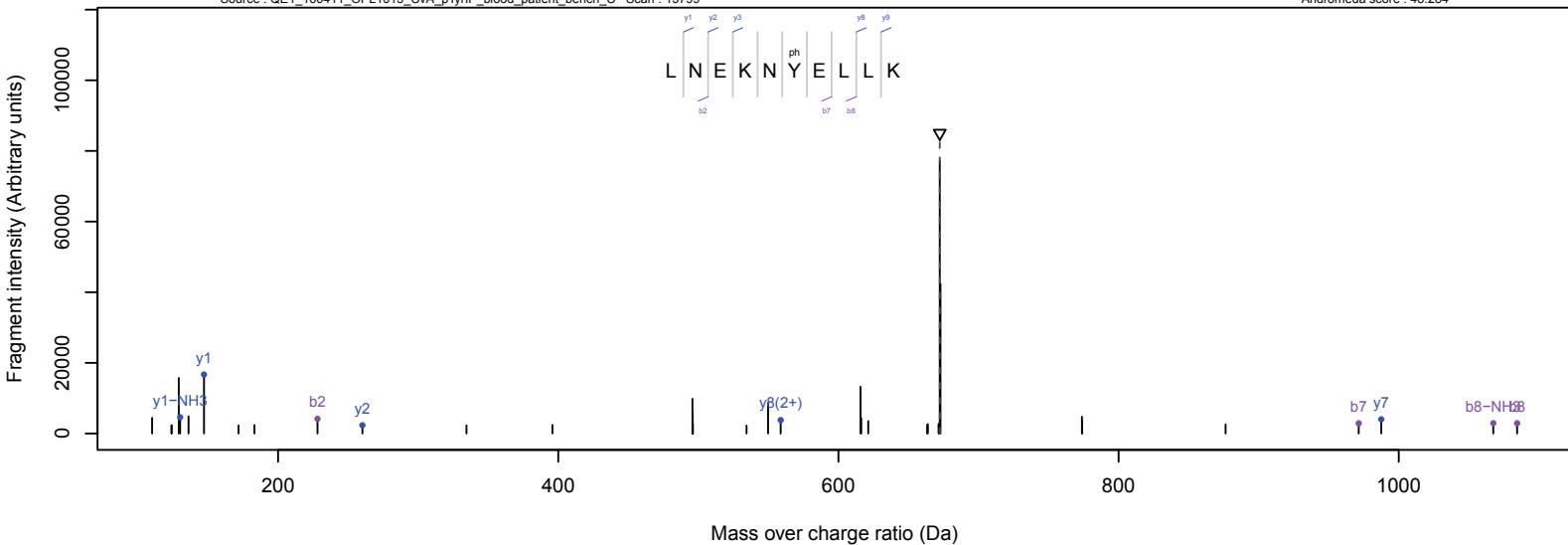
Andromeda score: 48.284



## LNEKNY(ph)ELLK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 13799

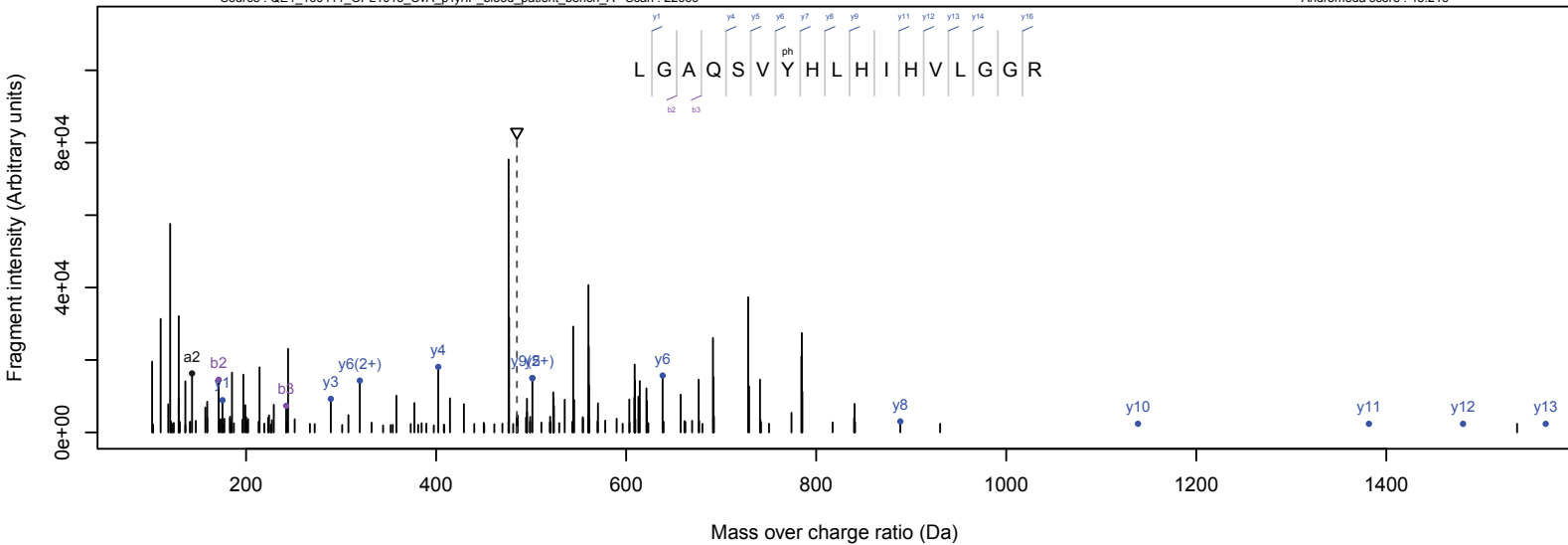
Andromeda score: 48.284



## LGAQSVY(ph)HLIHVLGGR-4+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan: 22669

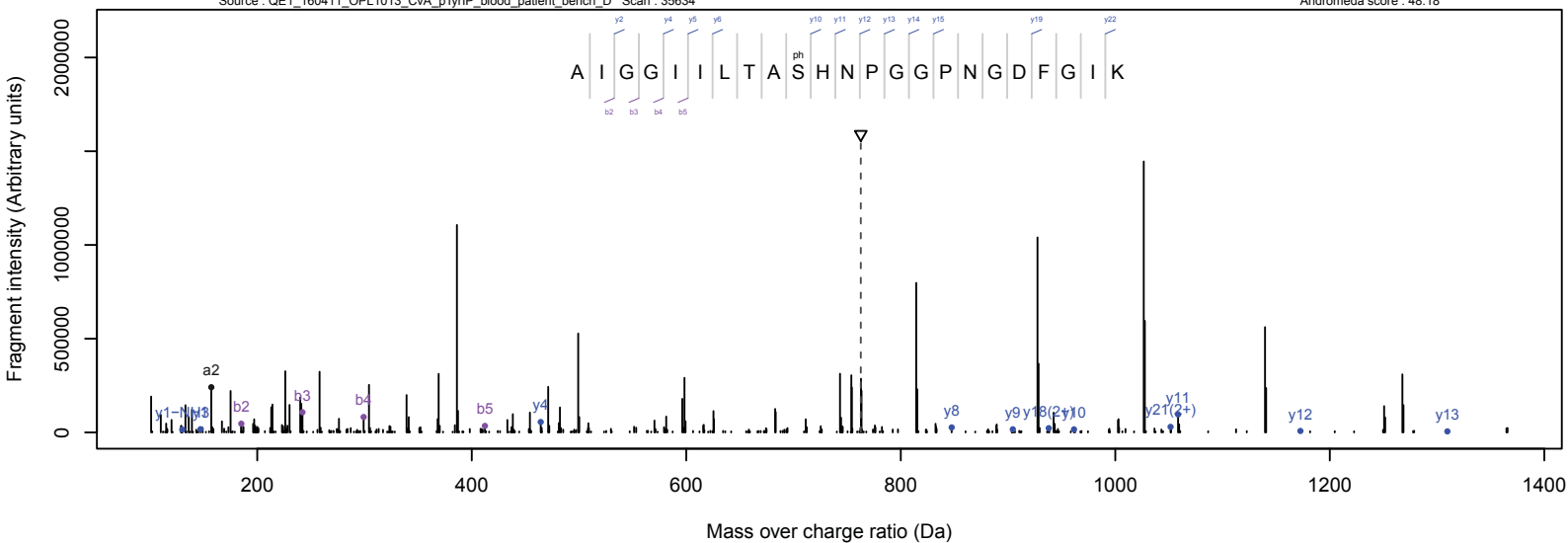
Andromeda score: 48.216



## AIGGILTAS(ph)HNPGGPNGDFGIK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 35634

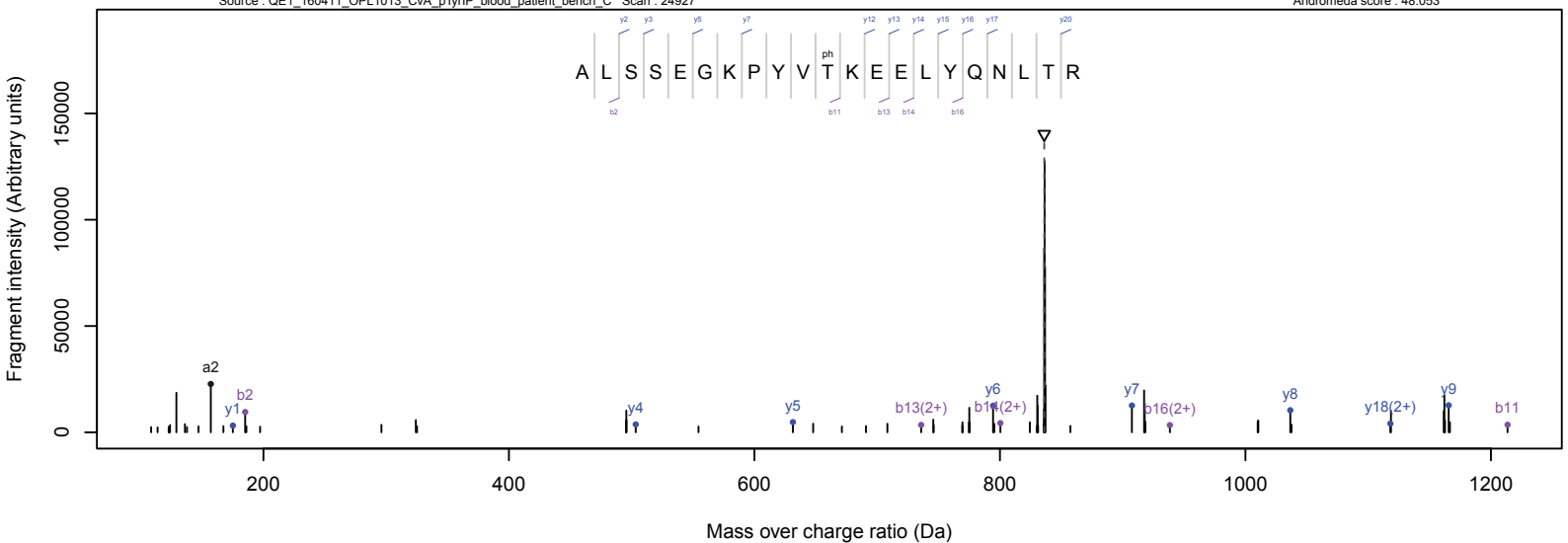
Andromeda score : 48.18



## ALSSEGKPYVT(ph)KEELYQLNLR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 24927

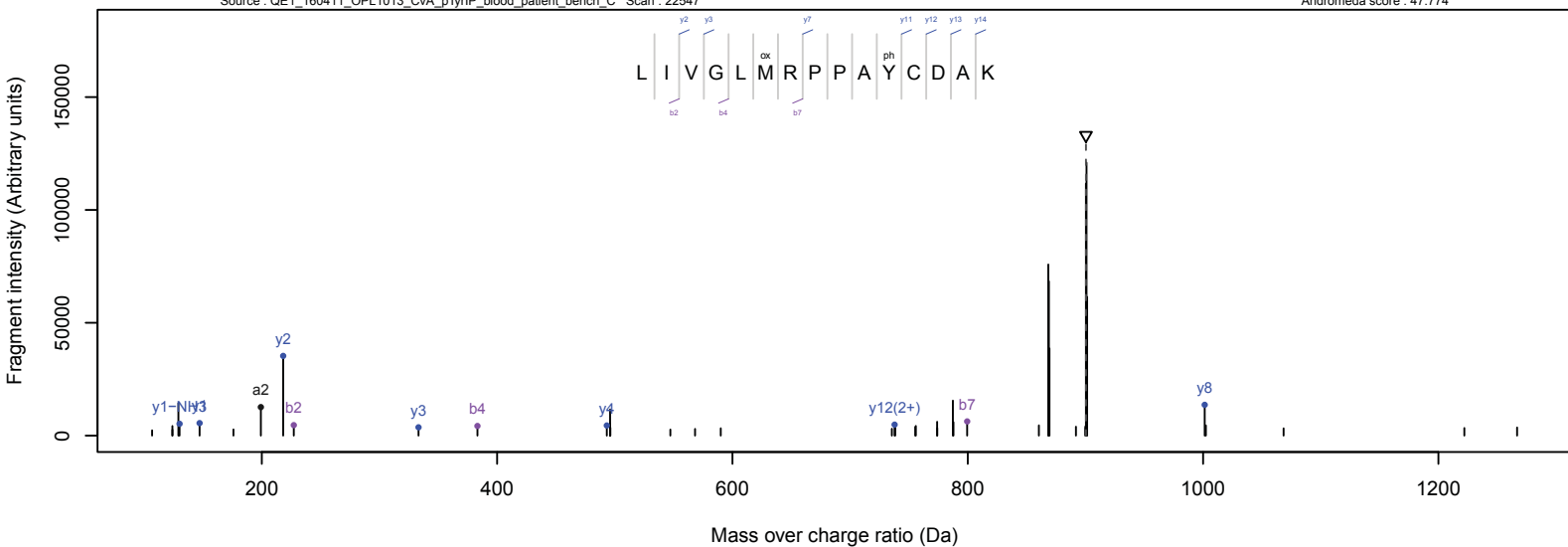
Andromeda score : 48.053



LIVGLM(ox)RPPAY(ph)CDAK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 22547

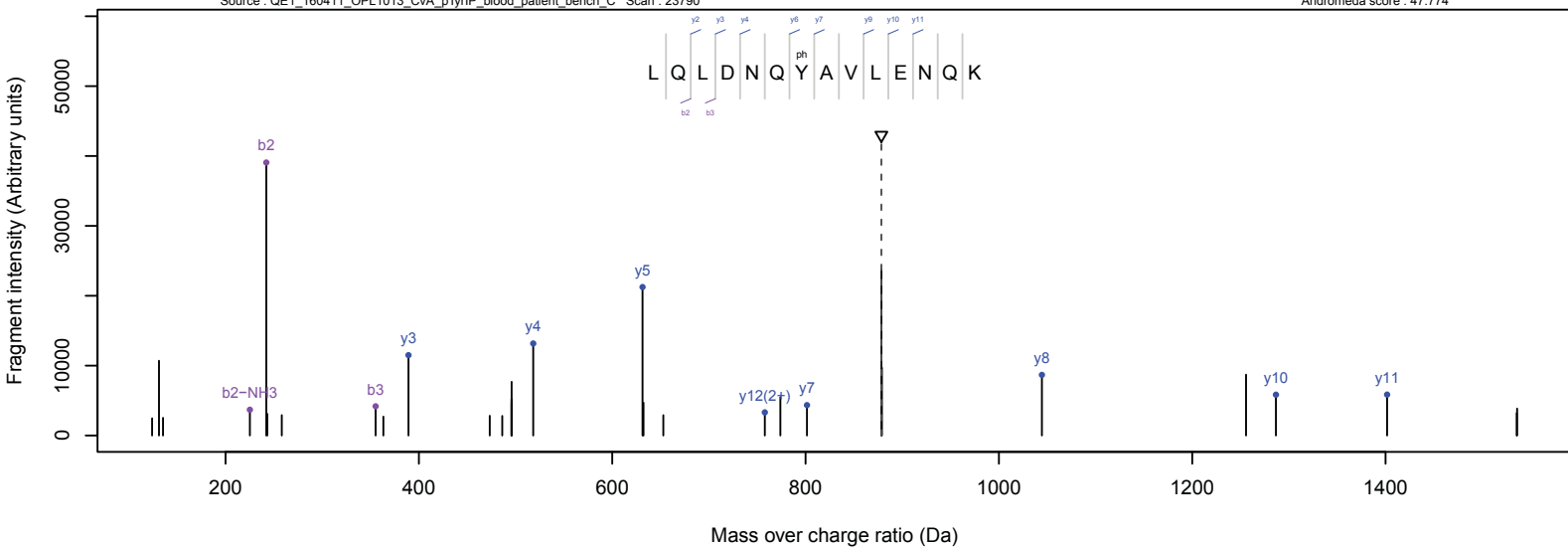
Andromeda score : 47.774



LQLDNQY(ph)AVLENQK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 23790

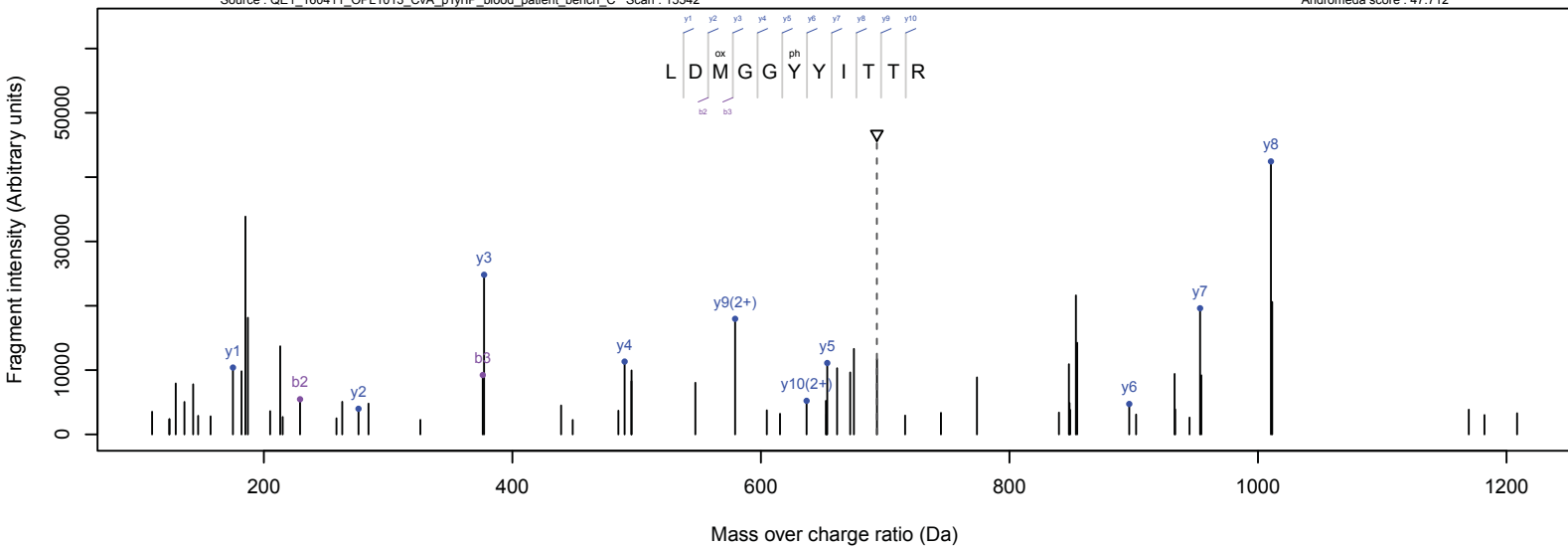
Andromeda score : 47.774



LDM(ox)GGY(ph)YITTR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 15542

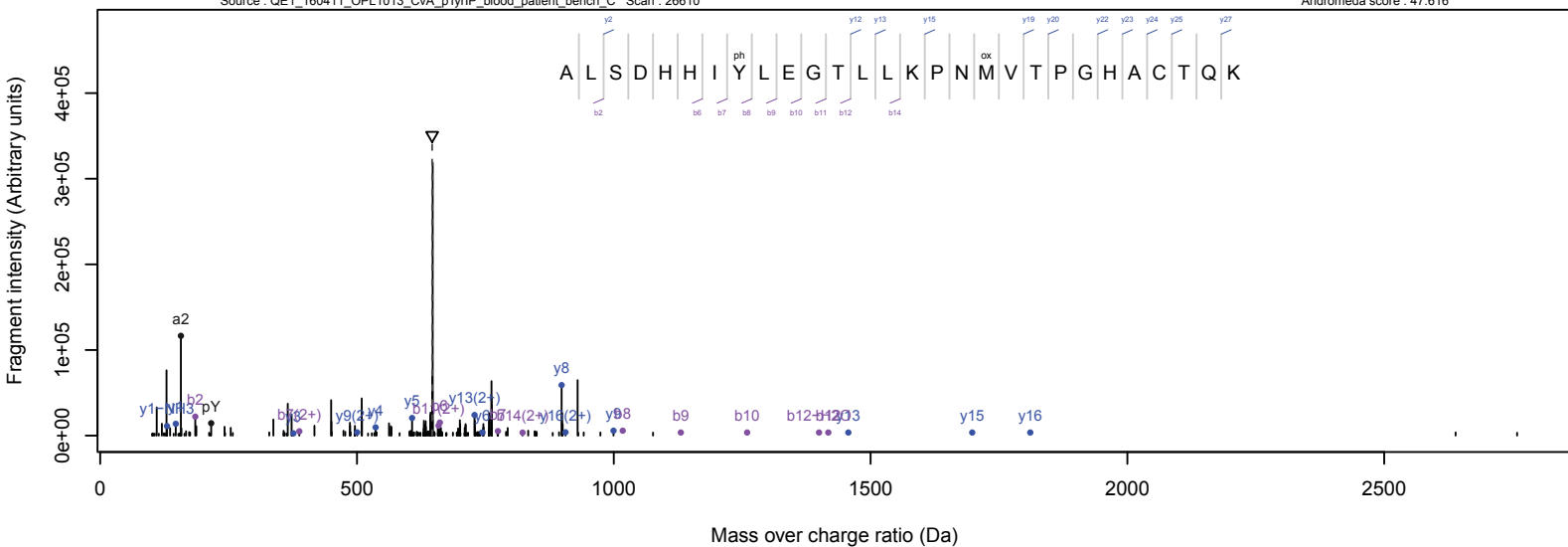
Andromeda score : 47.712



ALSDHIIY(ph)LEGTLKPNM(ox)VTPGHACTQK-5+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 26610

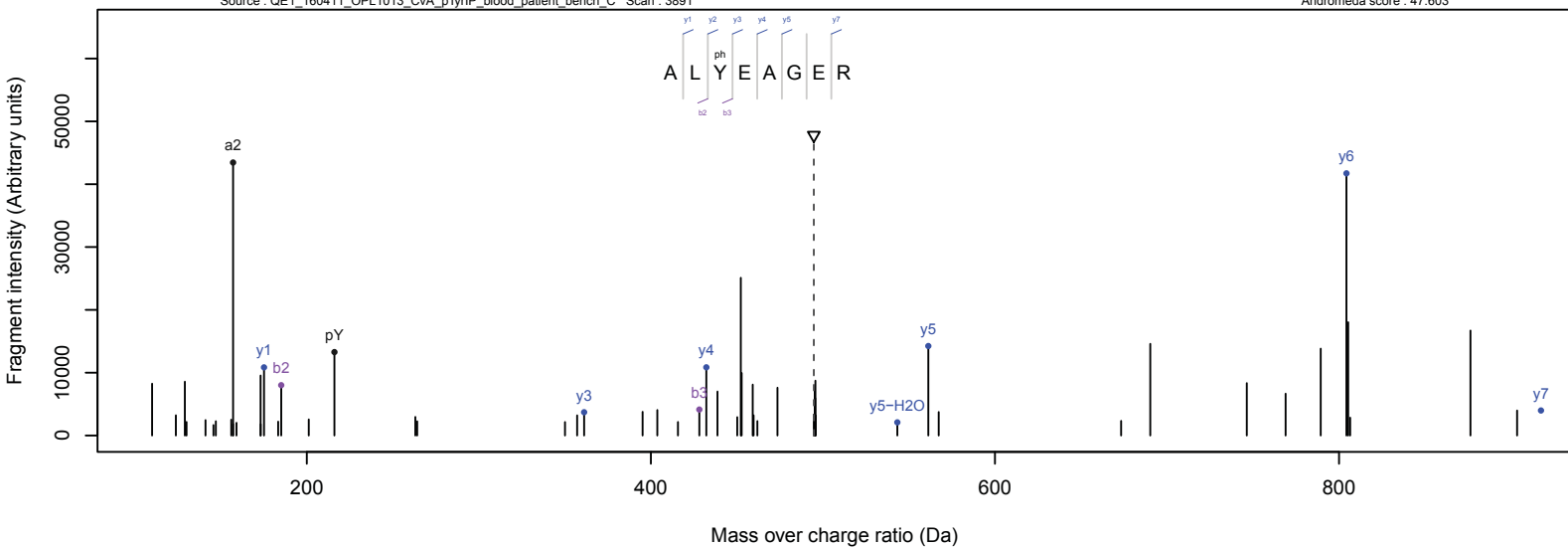
Andromeda score : 47.616



ALY(ph)EAGER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3891

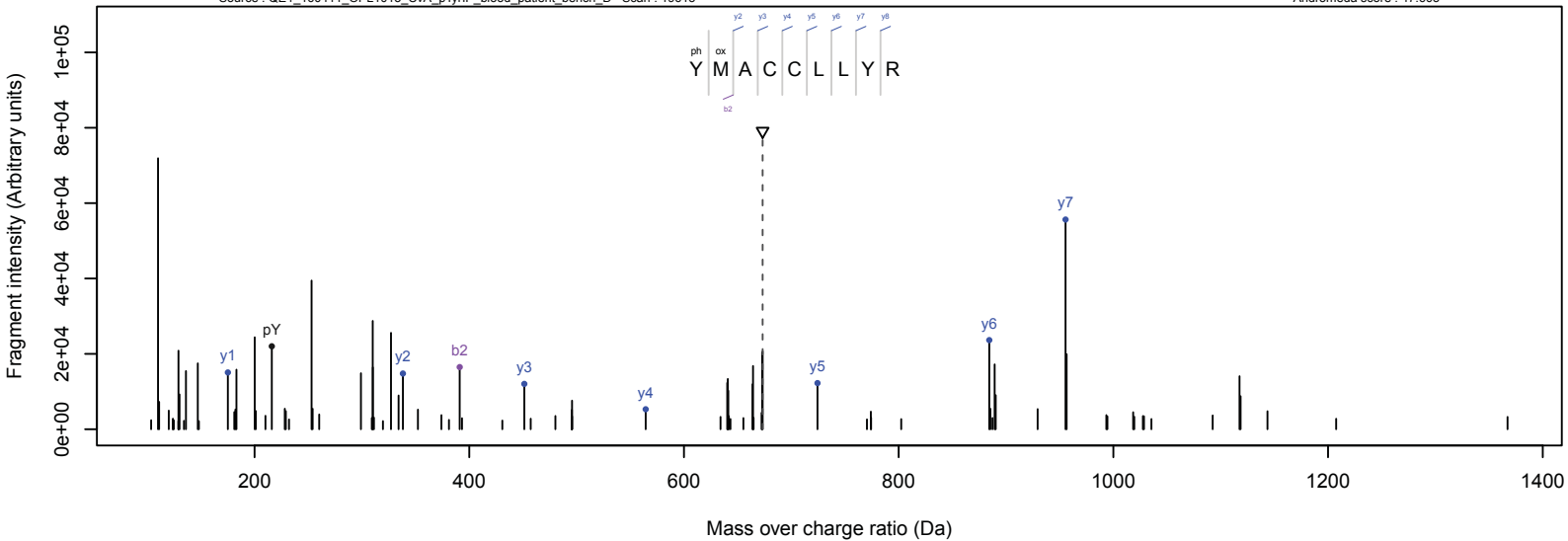
Andromeda score : 47.603



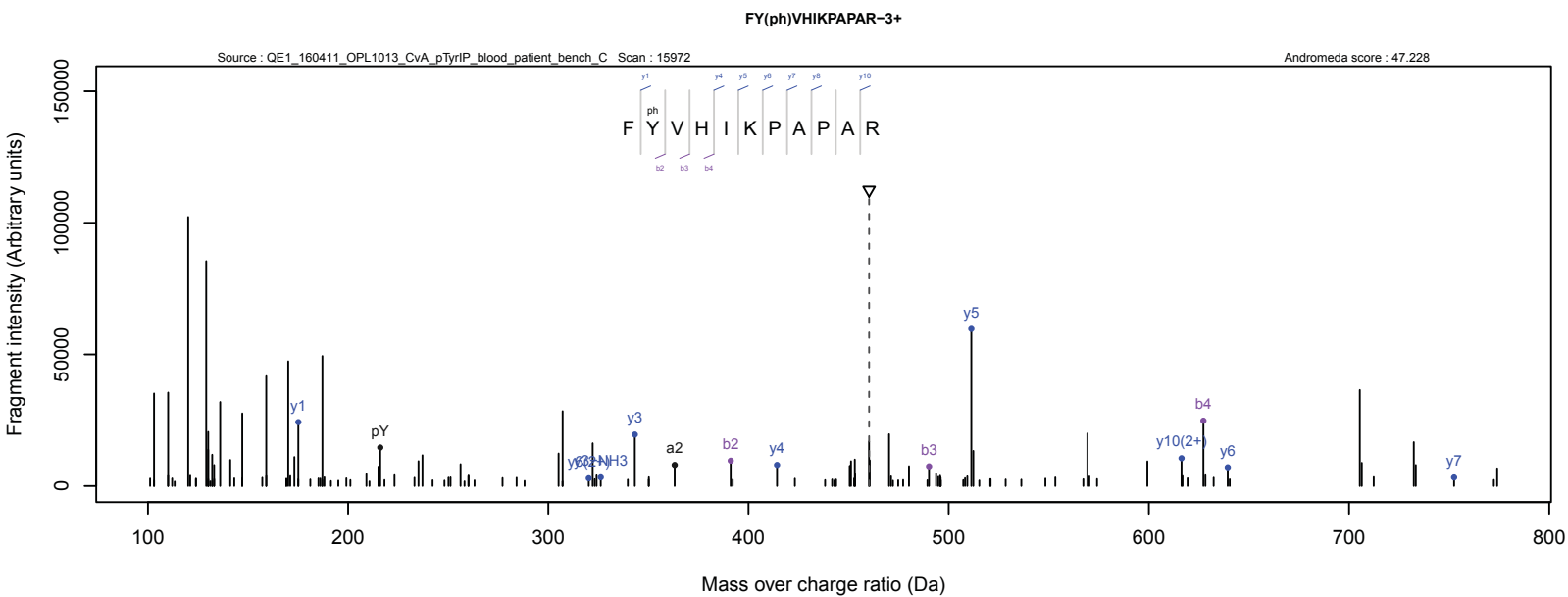
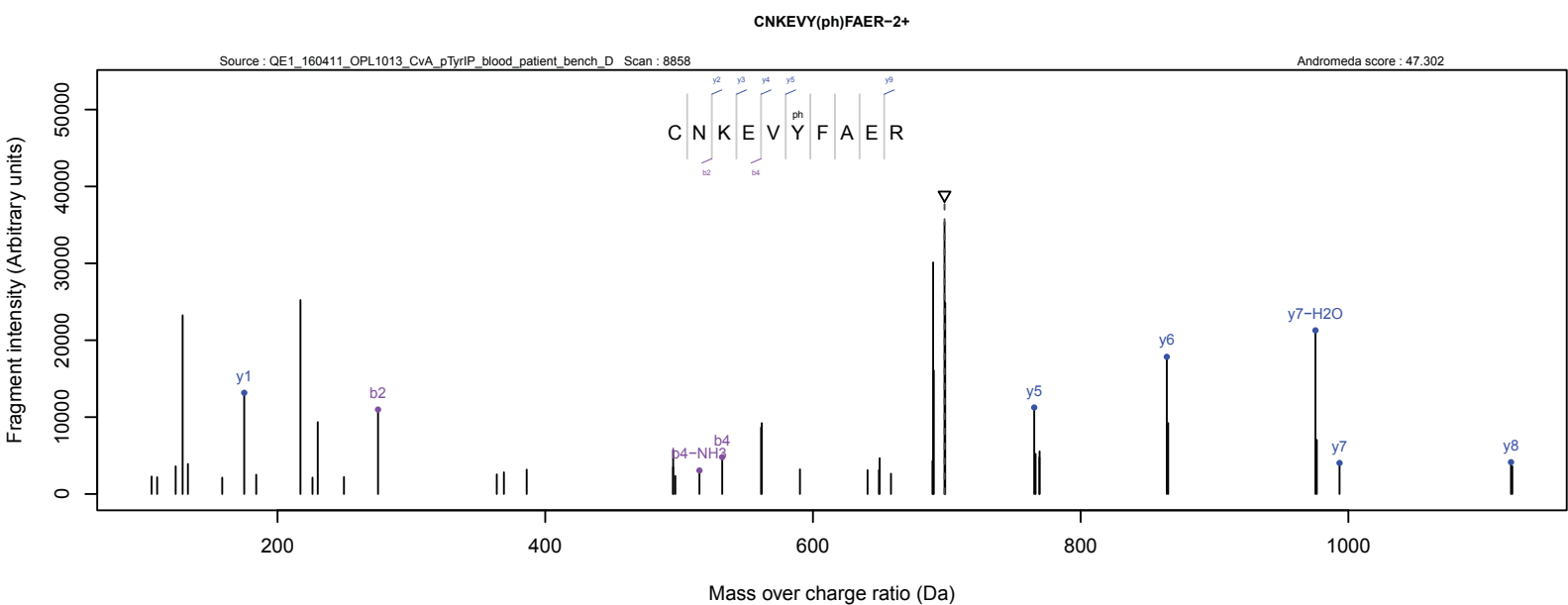
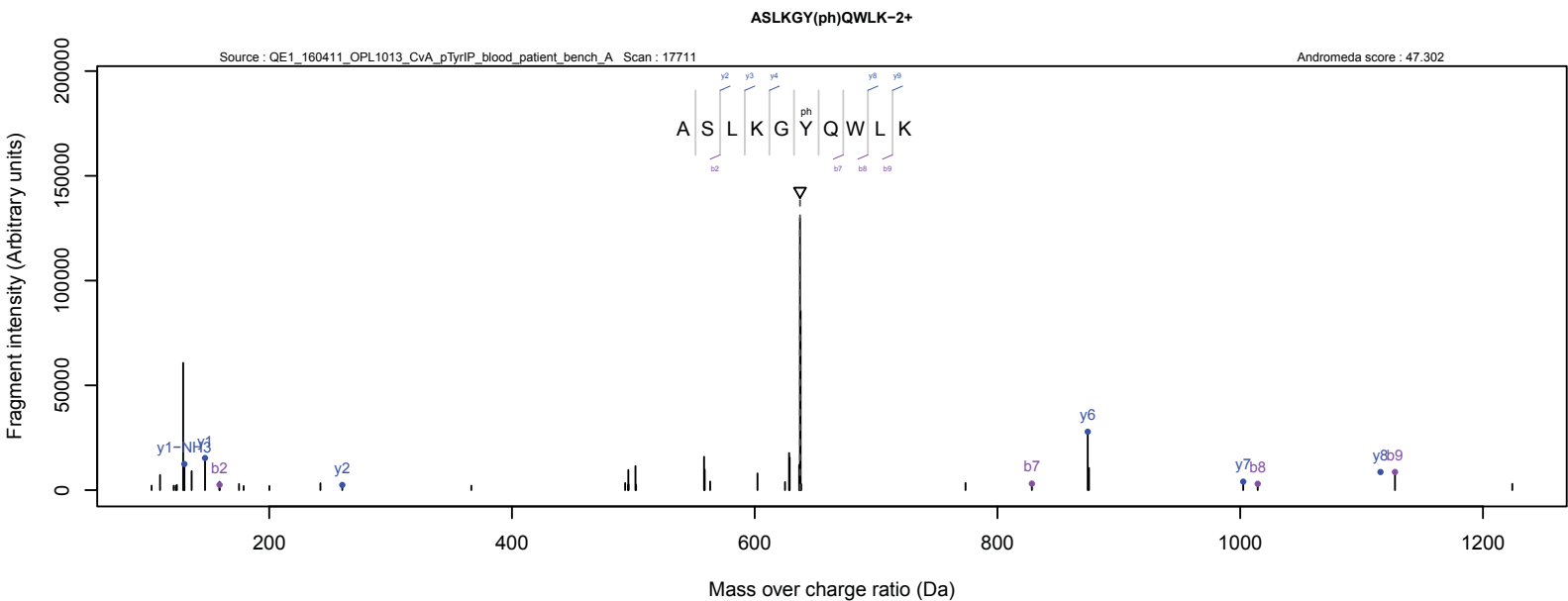
Y(ph)M(ox)ACLLYR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 19615

Andromeda score : 47.603



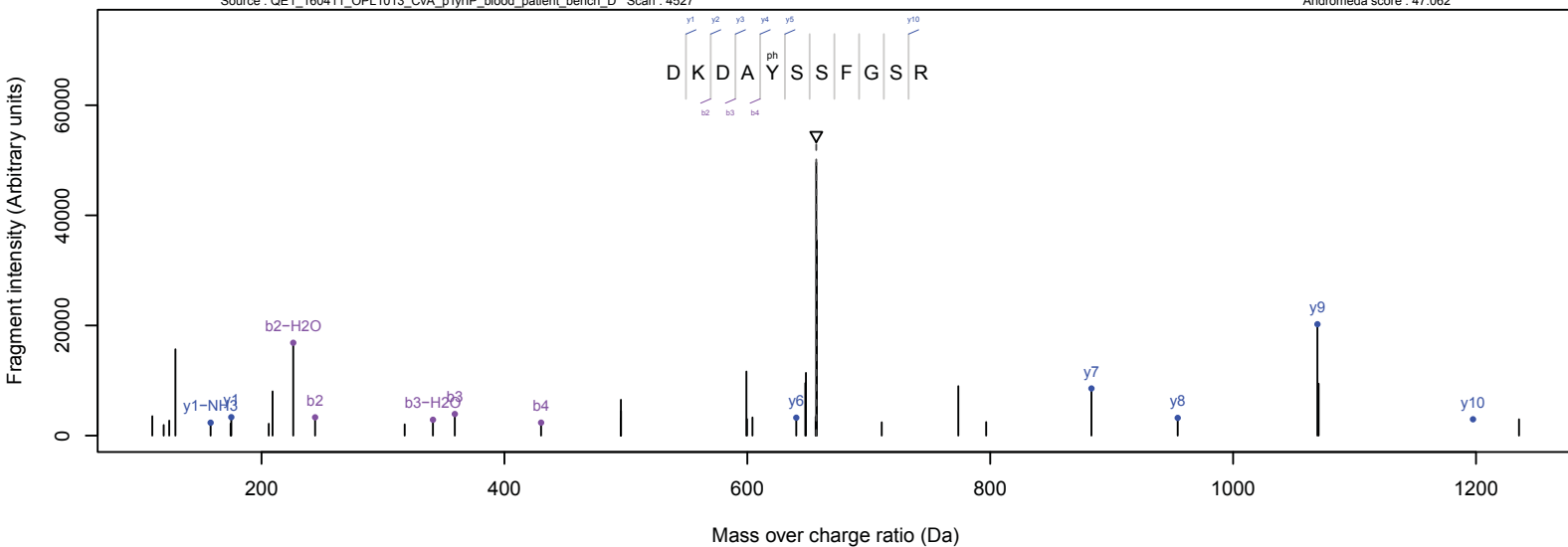




**DKDAY(ph)SSFGSR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 4527

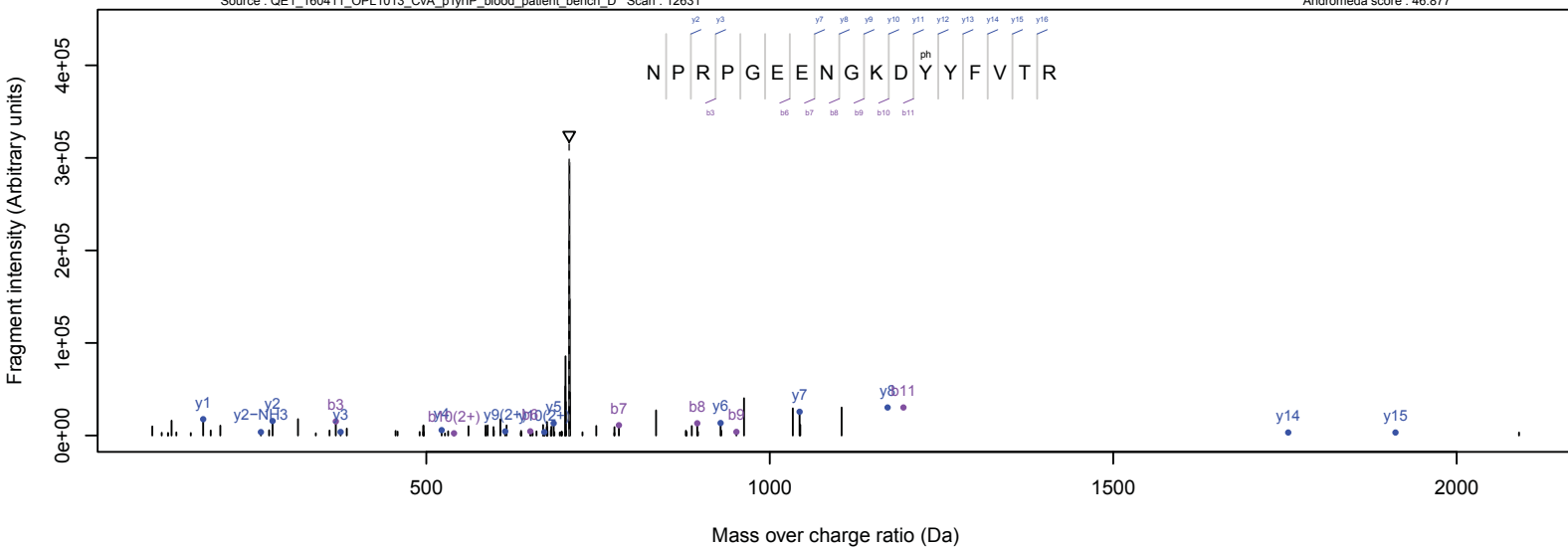
Andromeda score : 47.062



**NPRPGEENGKDY(ph)YFVTR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12631

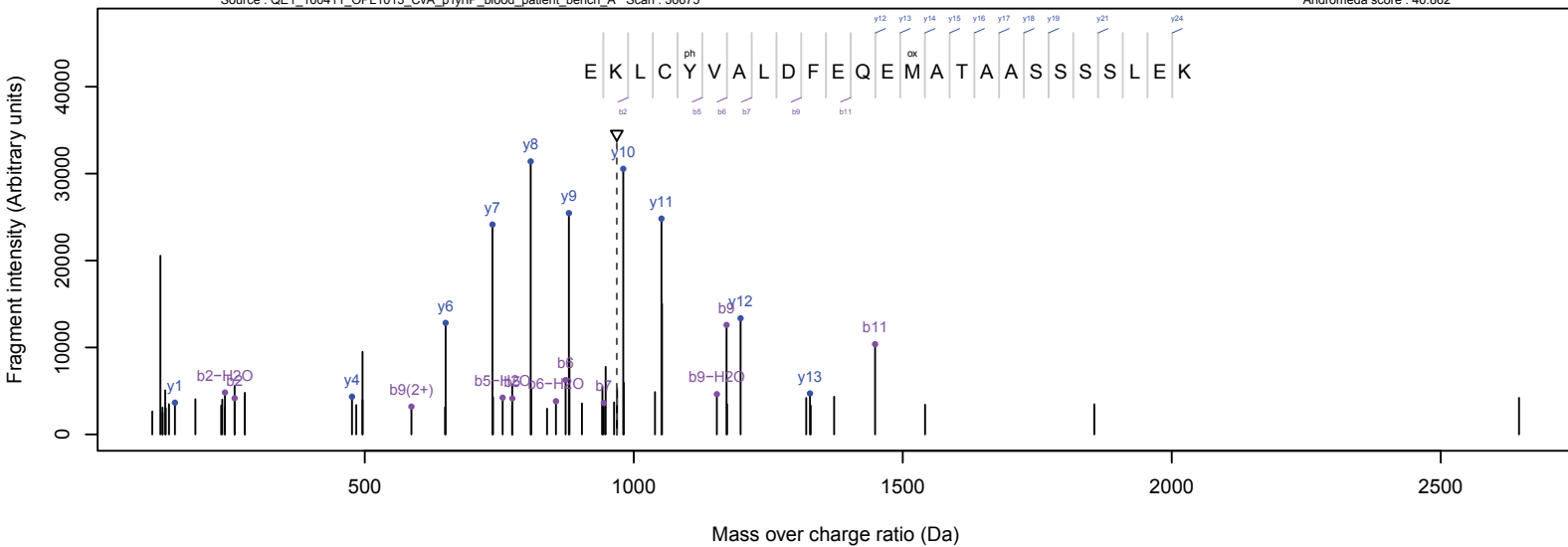
Andromeda score : 46.877



**EKLCY(ph)VALDFEQEM(ox)ATAASSSSLEK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 36675

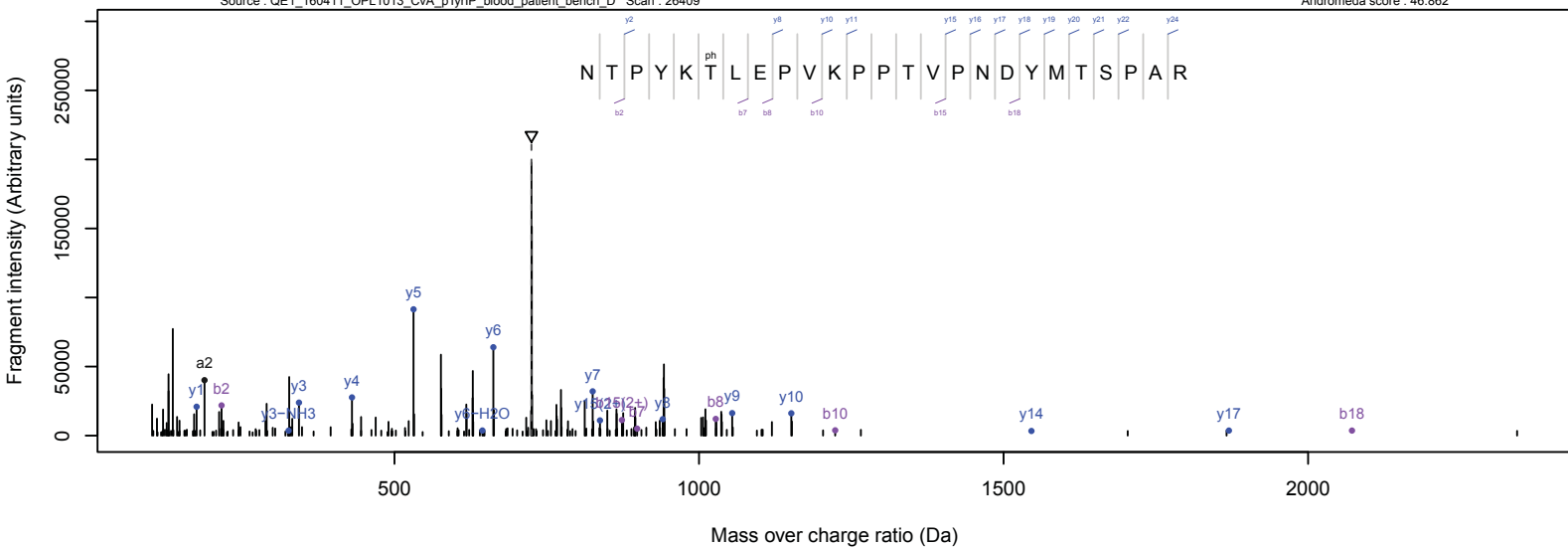
Andromeda score : 46.862



**NTPYK(ph)LEPVKPTVPNDYMTSPAR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 26409

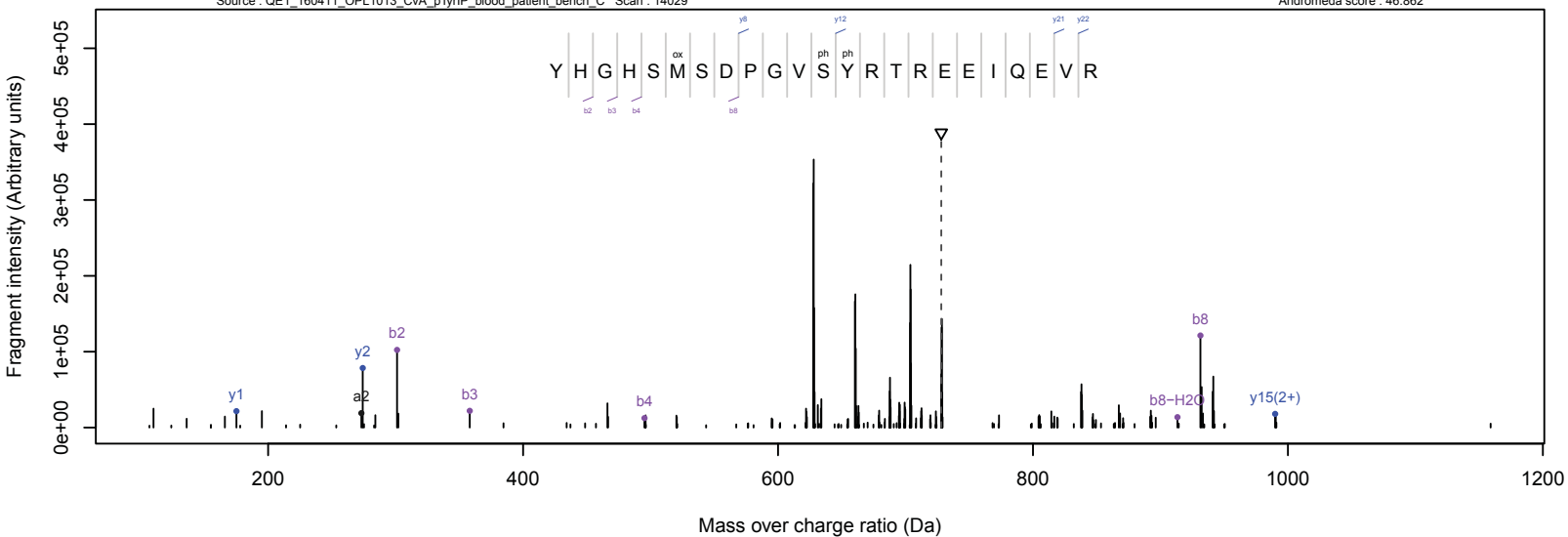
Andromeda score : 46.862



**YHGHSM(ox)SDPGVS(ph)Y(ph)RTREEIQEVR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 14029

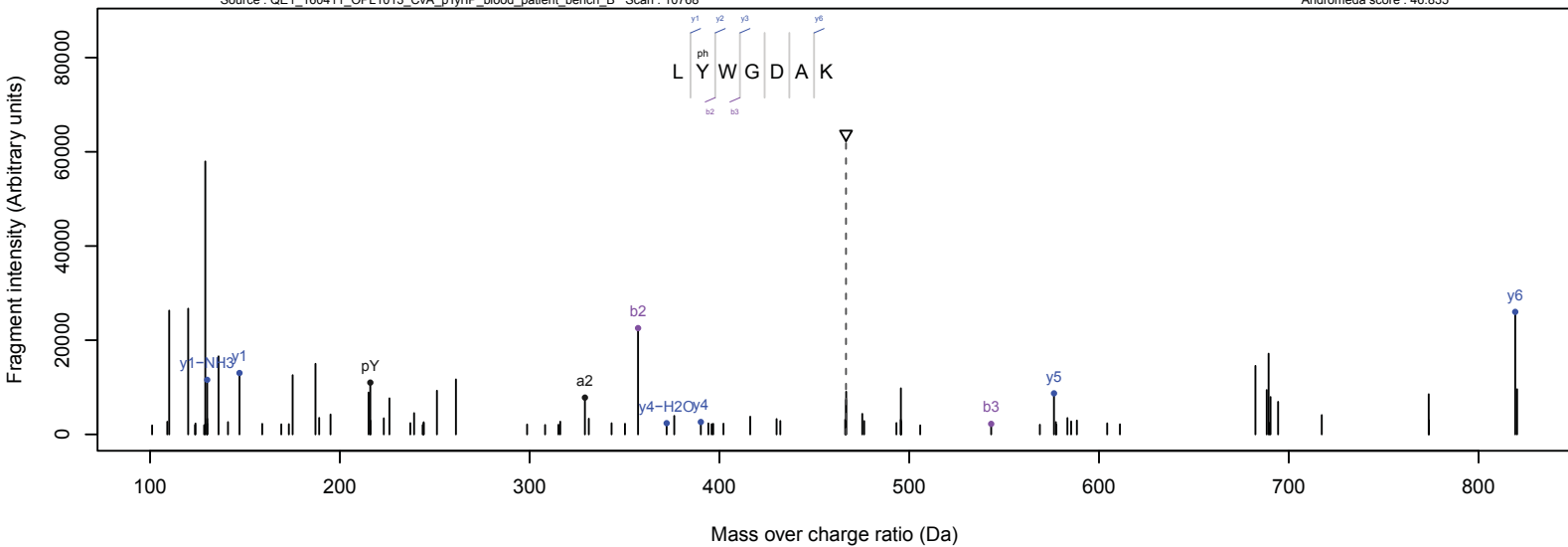
Andromeda score : 46.862



**LY(ph)WGDAK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 10768

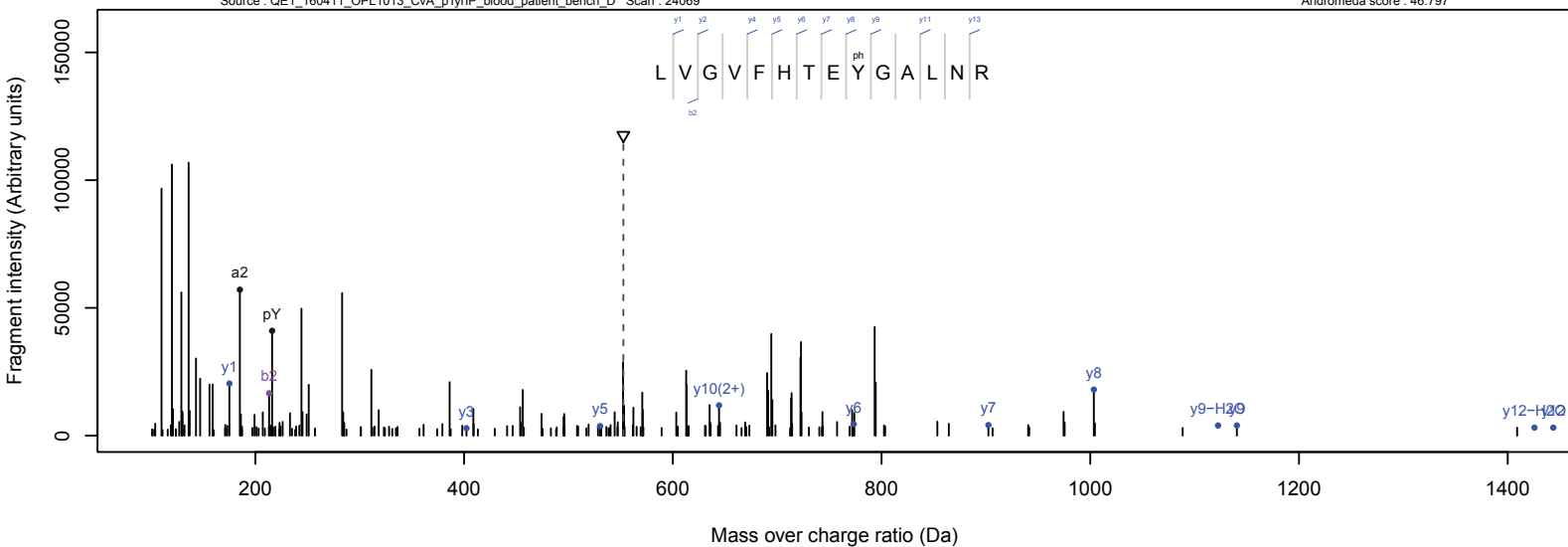
Andromeda score : 46.835



## LVGVFHTYEY(ph)GALNR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 24069

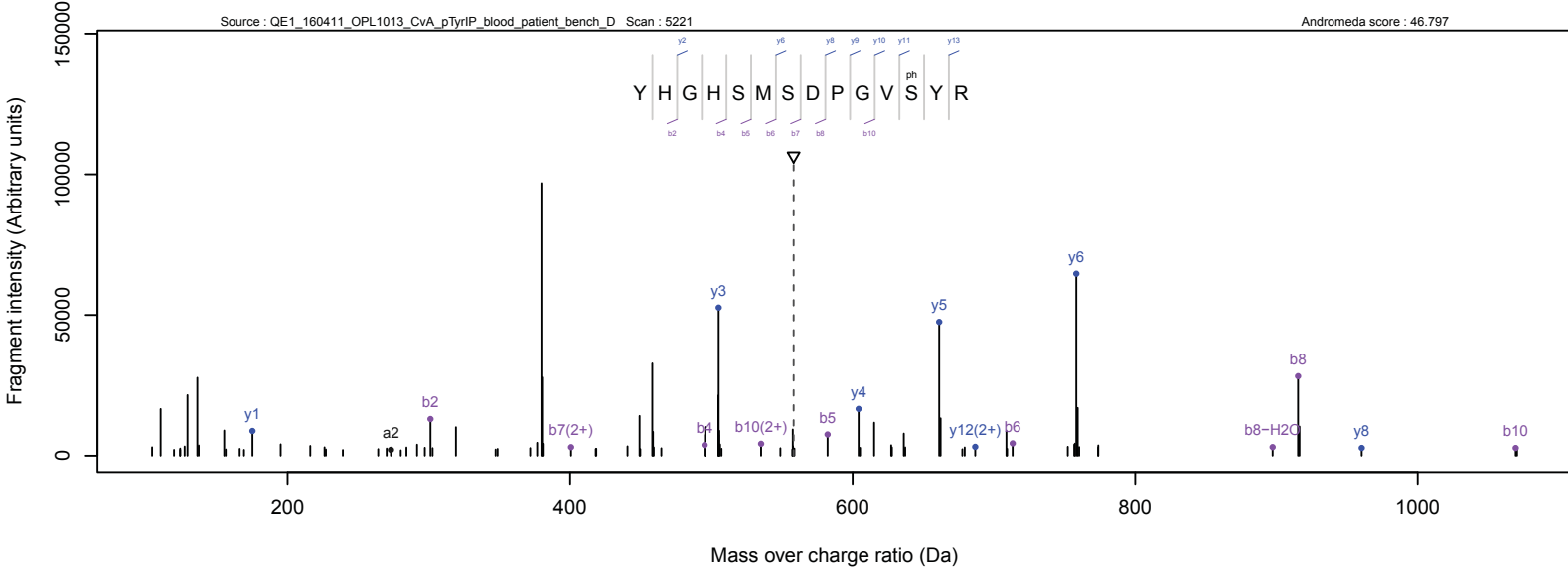
Andromeda score: 46.797



## YHGHSMDPGVS(ph)YR-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 5221

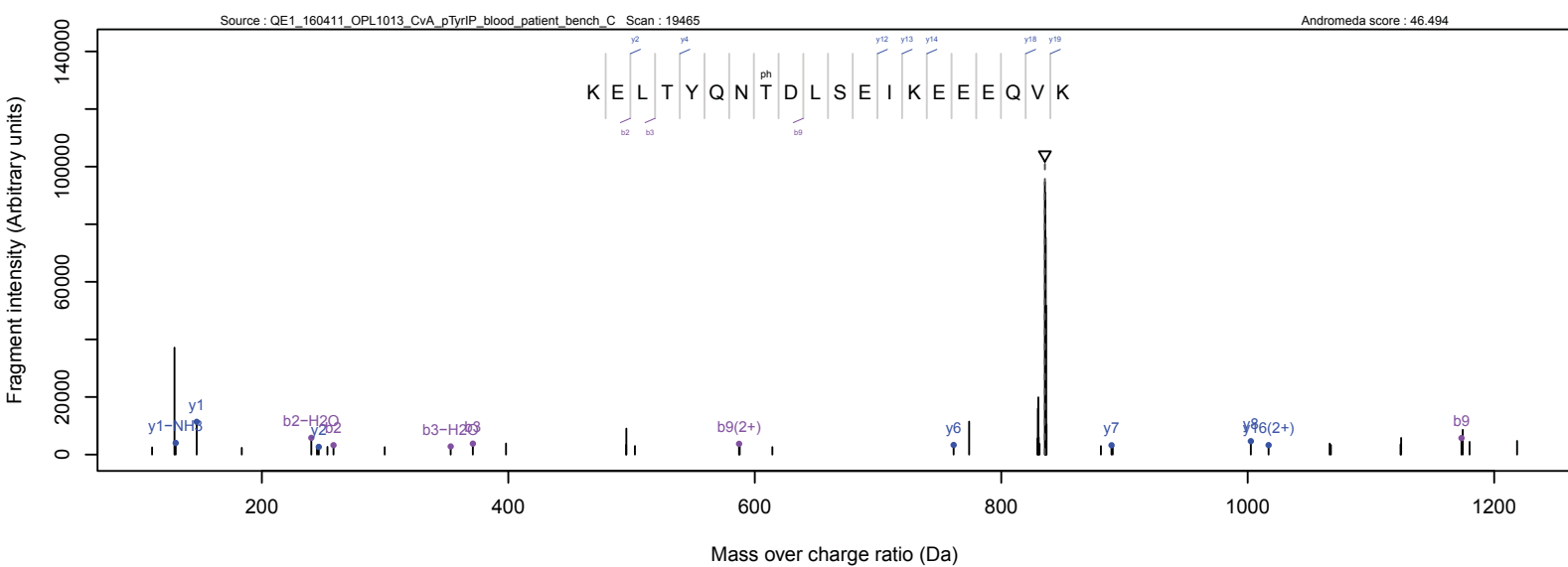
Andromeda score: 46.797



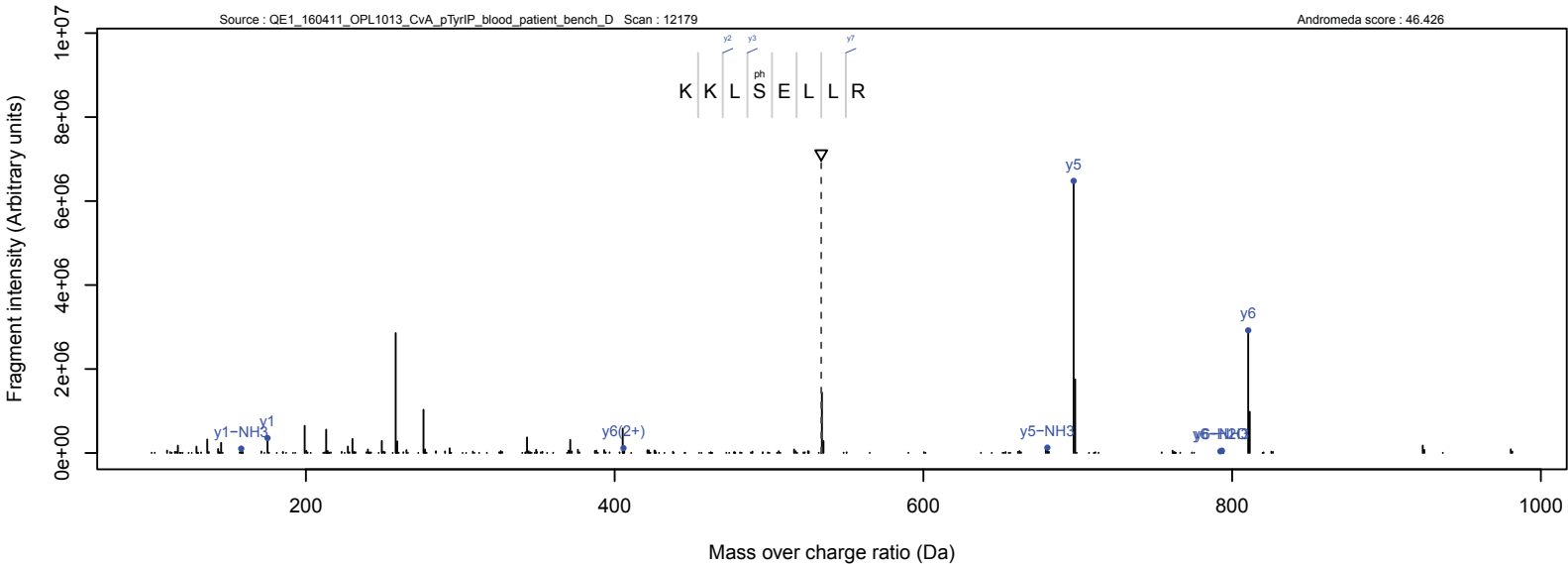
## KELTYQNT(ph)DLSEIKEEEQVK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 19465

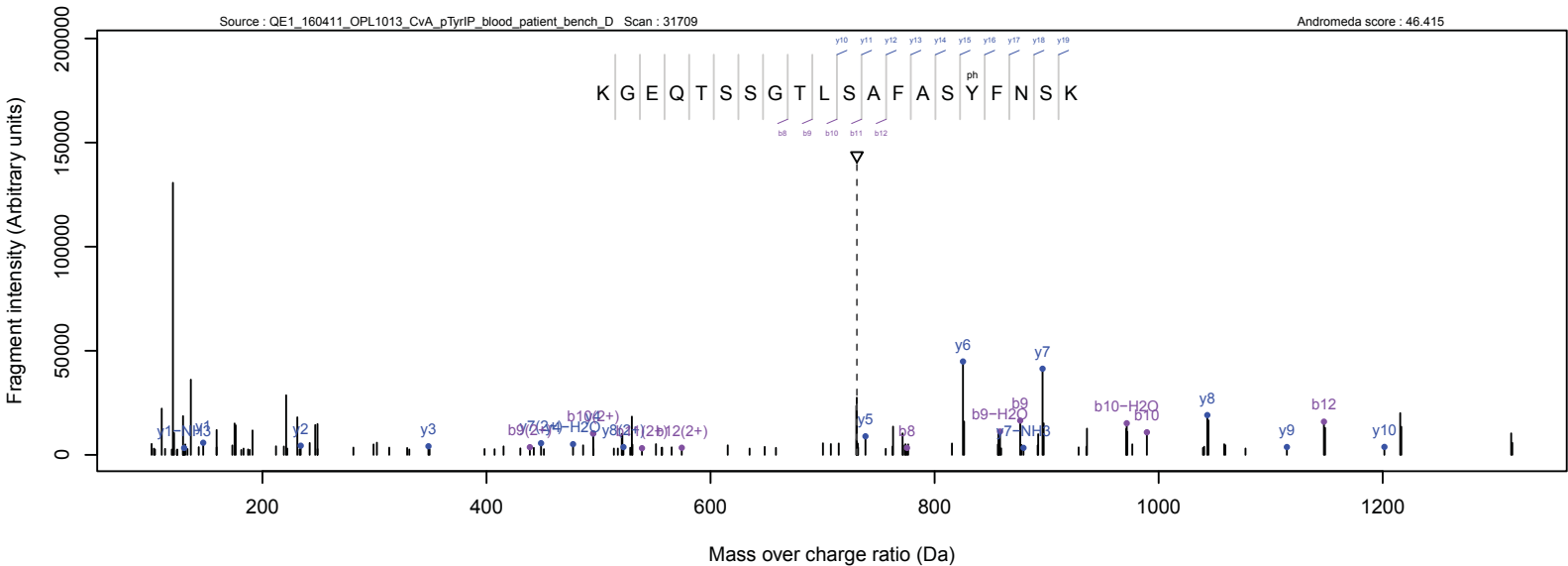
Andromeda score: 46.494



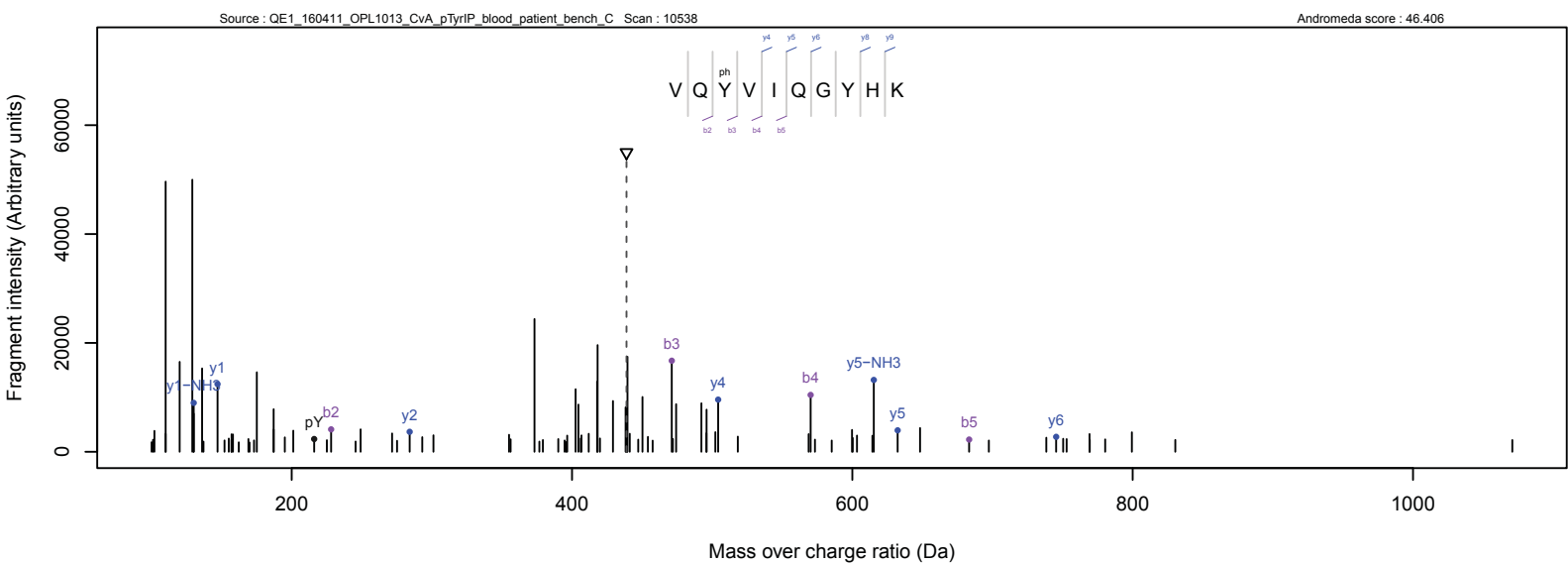
**KKLS(ph)ELLR-2+**



**KGEQTSSGTLSAFAS(ph)YFNSK-3+**



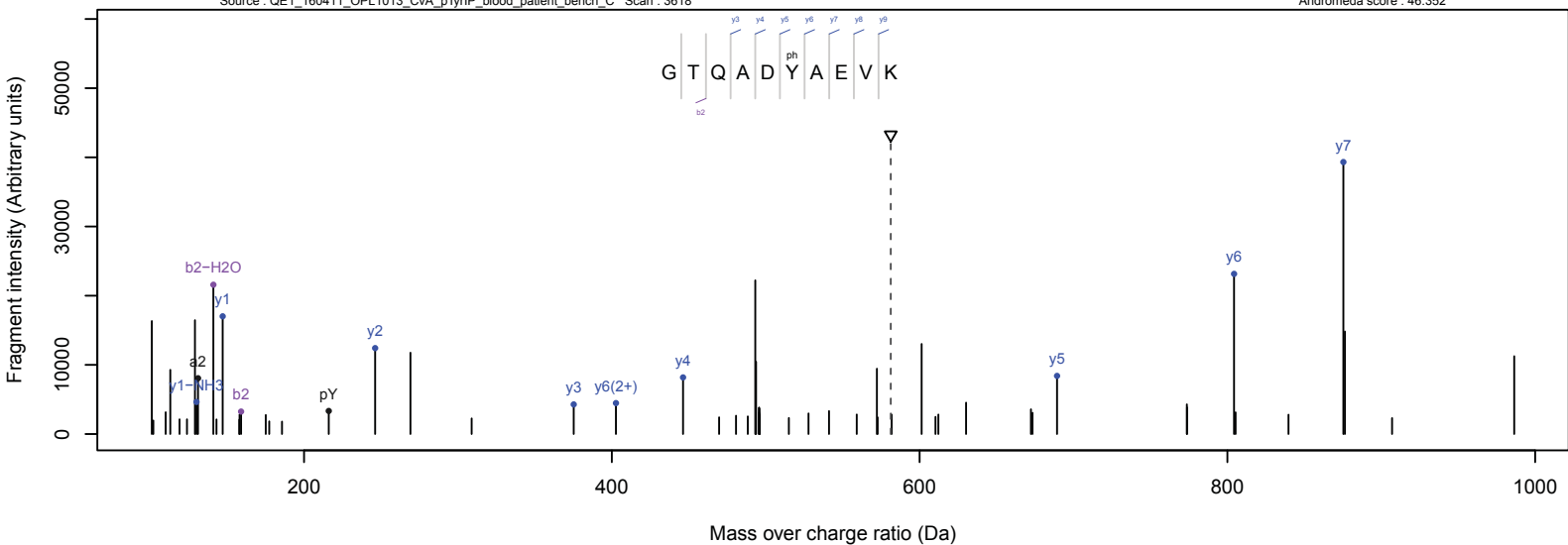
**VQY(ph)VIQGYHK-3+**



GTQADY(ph)AEVK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3618

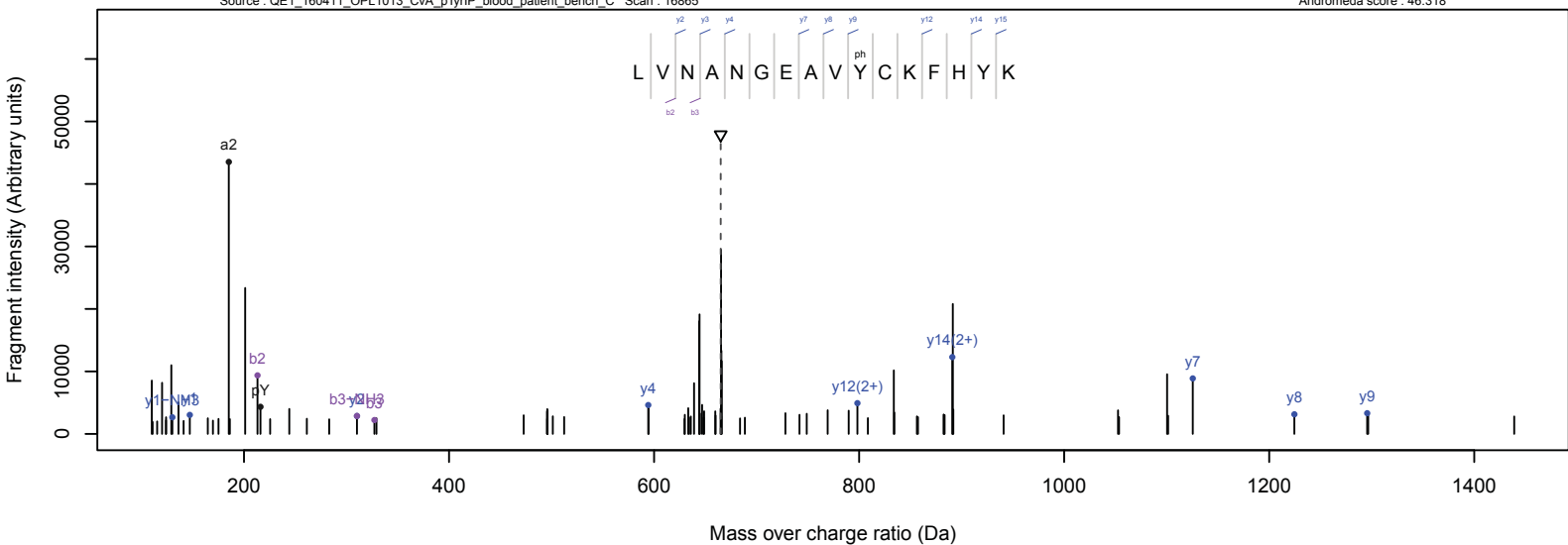
Andromeda score : 46.352



LVNANGEAVY(ph)CKFHYK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16865

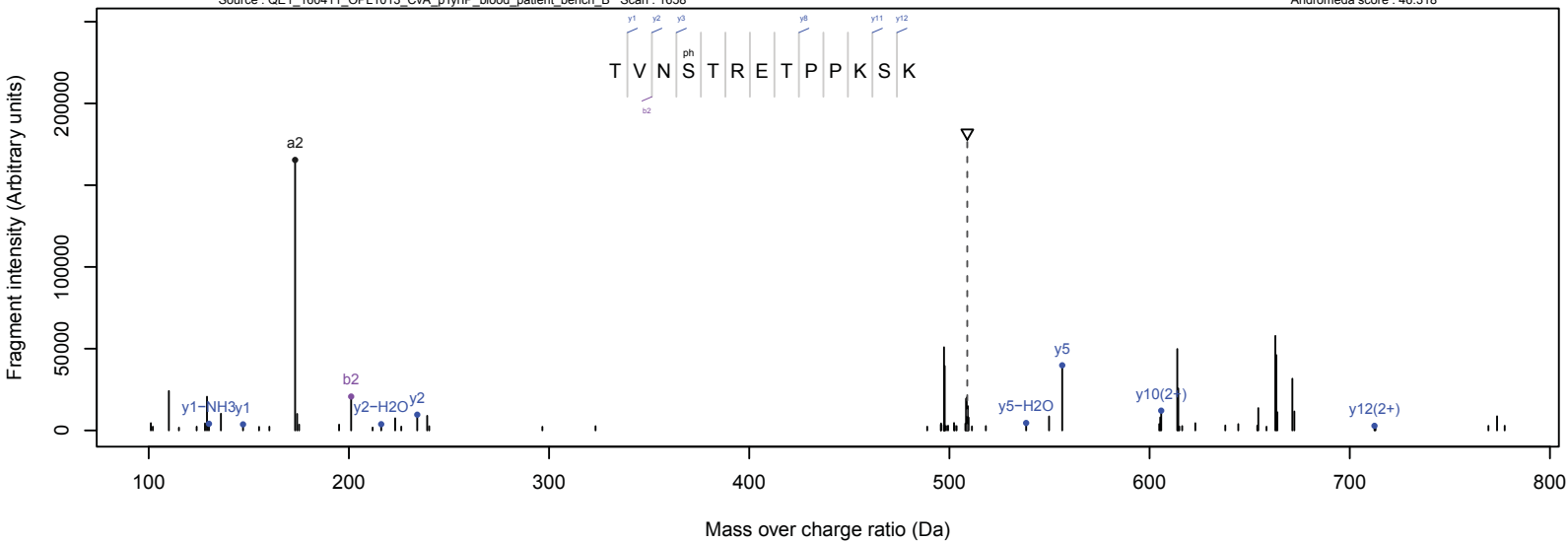
Andromeda score : 46.318



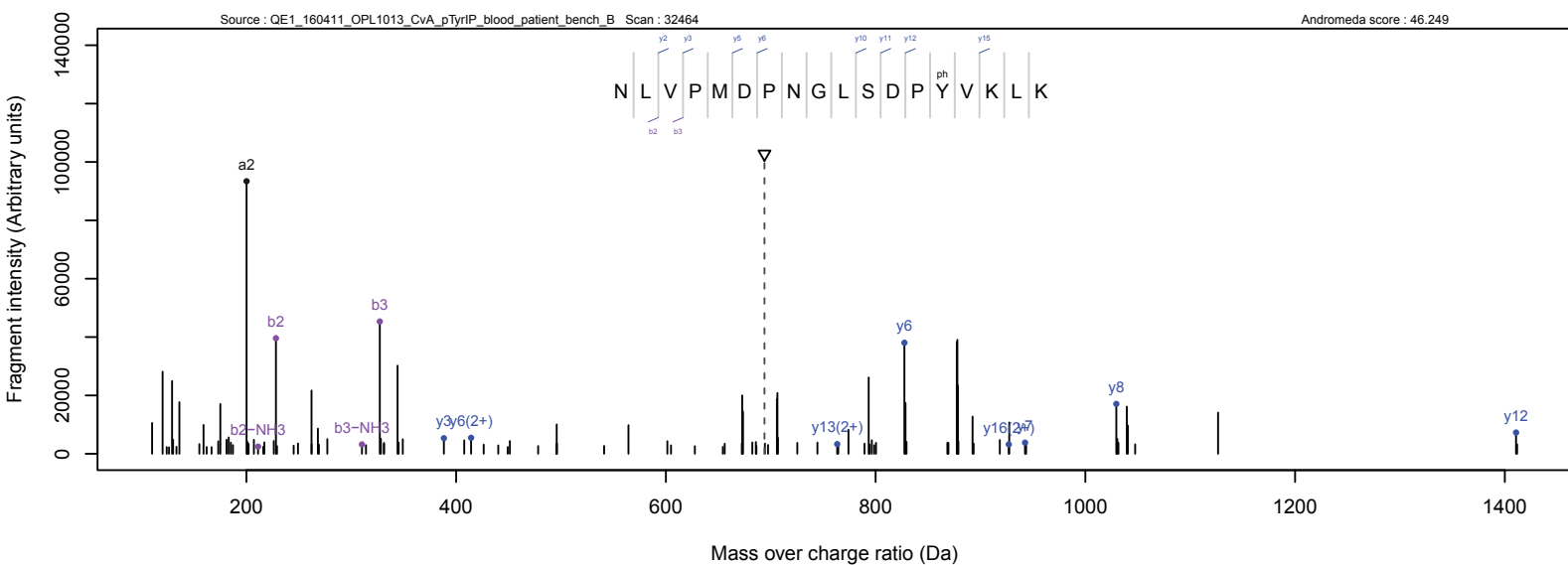
TVNS(ph)TRETTPKSK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 1658

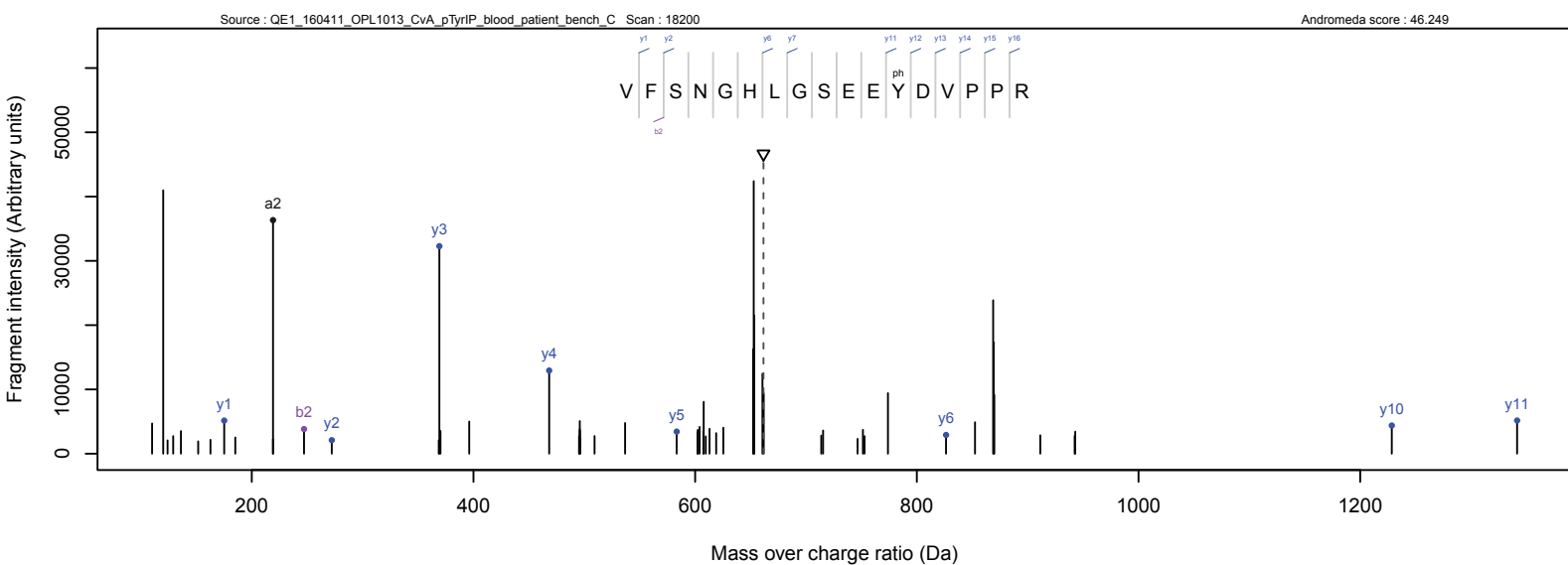
Andromeda score : 46.318



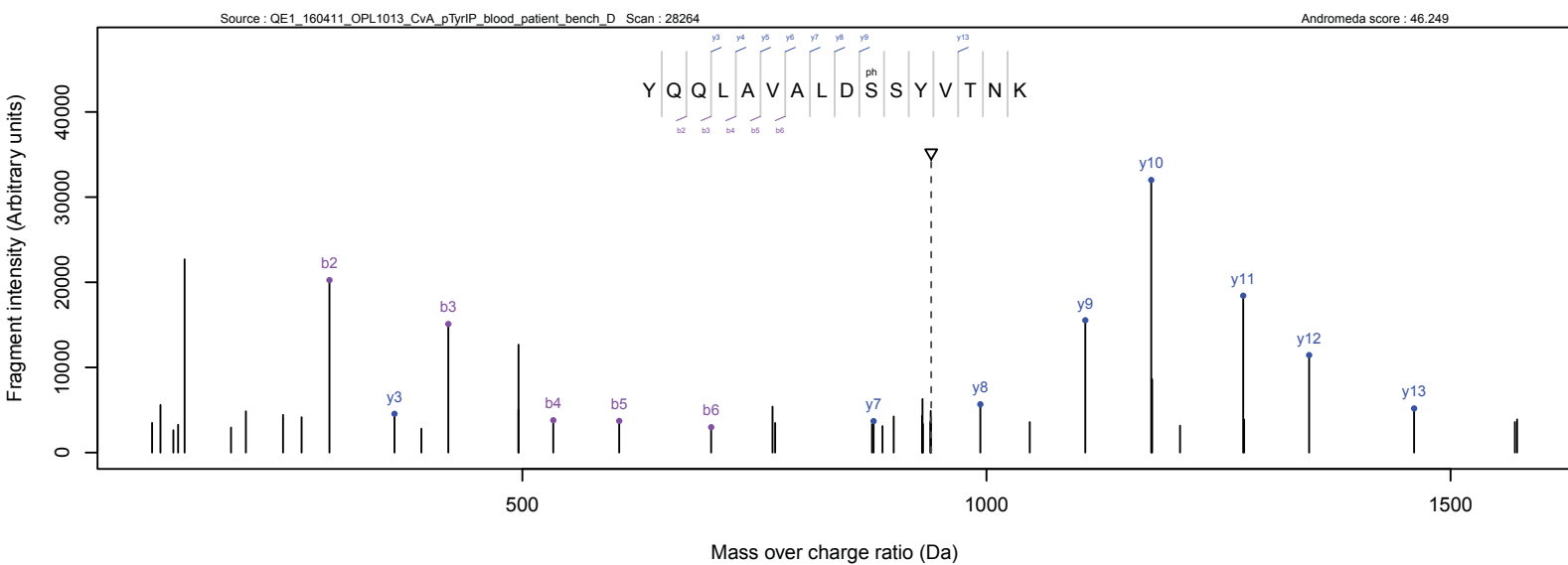
## NLVPMDPNGLSDPY(ph)VKLK-3+



## VFSNGLGSEEY(ph)DVPPR-3+



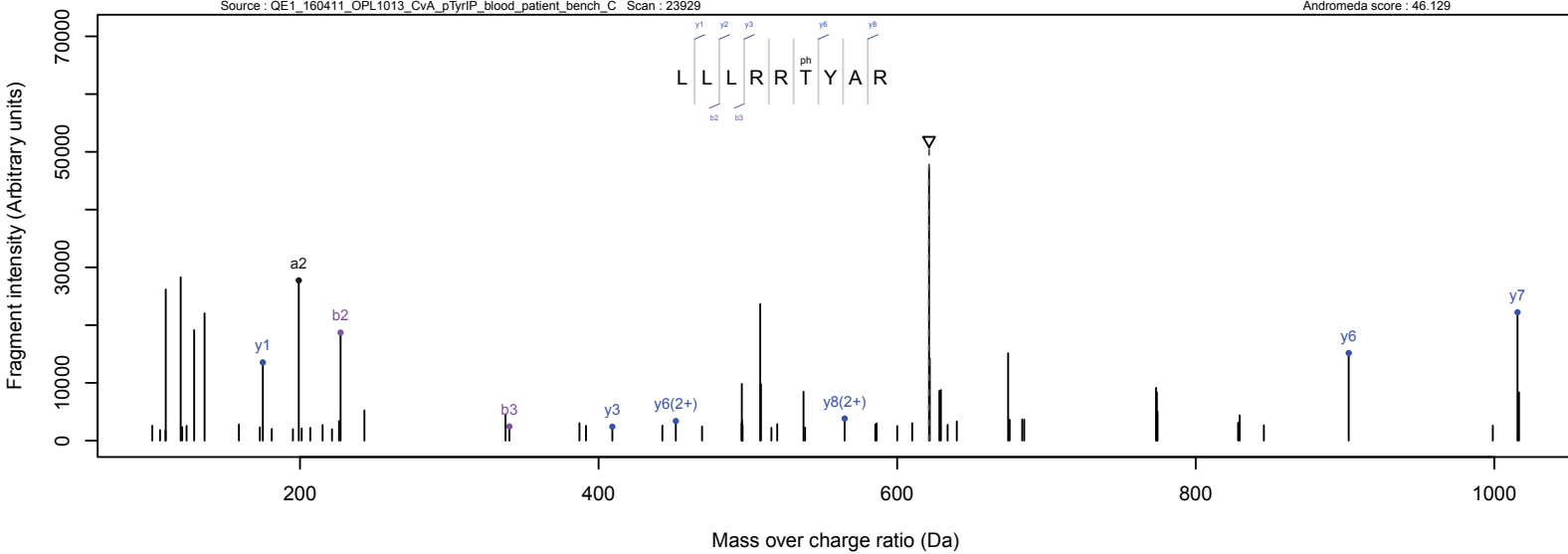
## YQLAVALDS(ph)SYVTNK-2+



## LLLRRT(ph)YAR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 23929

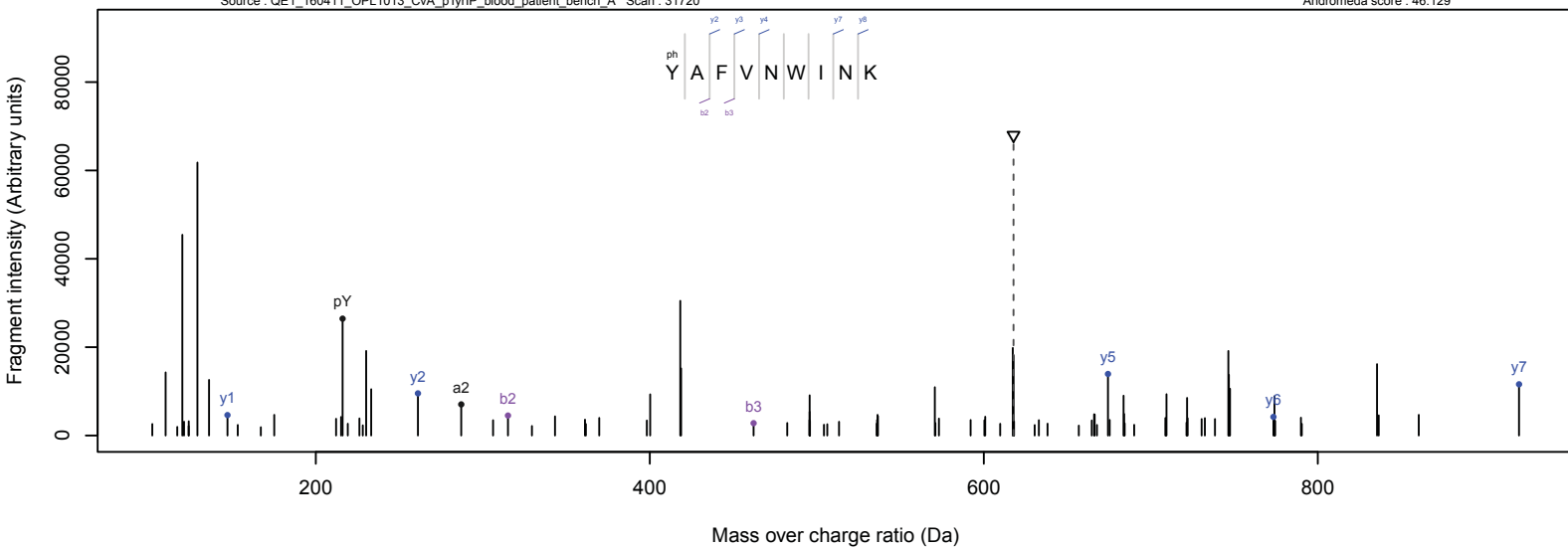
Andromeda score : 46.129



## Y(ph)AFVNWINK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 31720

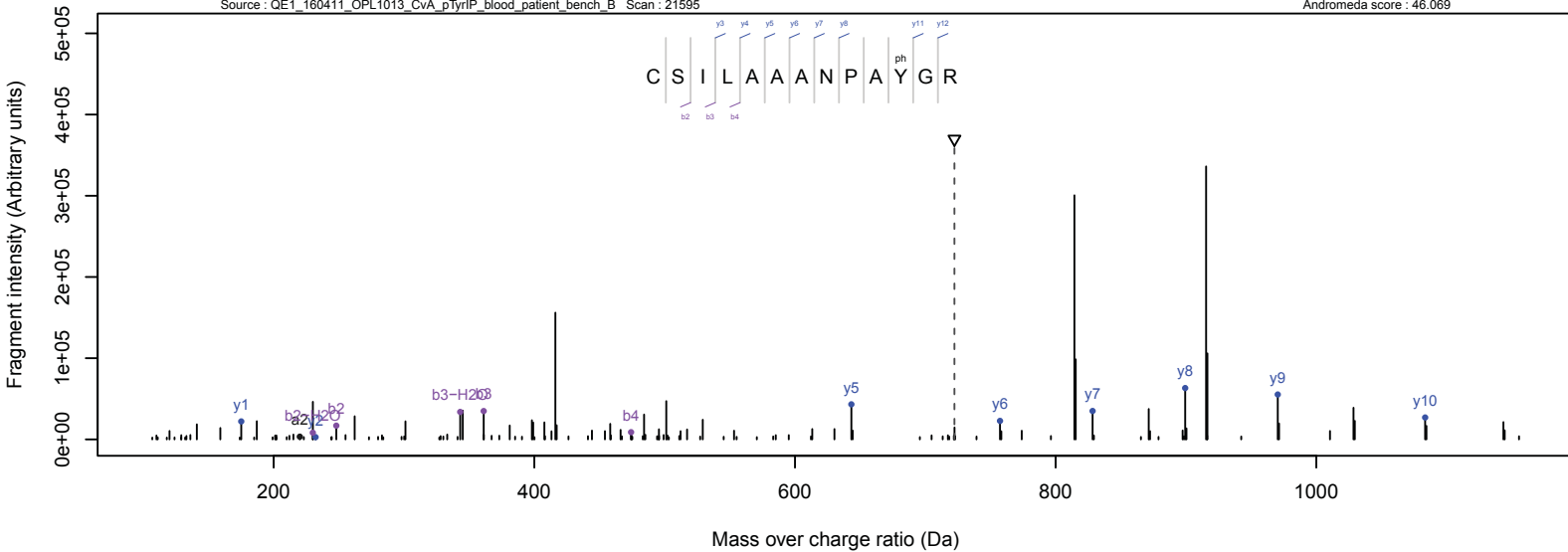
Andromeda score : 46.129



## CSILAAANPAY(ph)GR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 21595

Andromeda score : 46.069

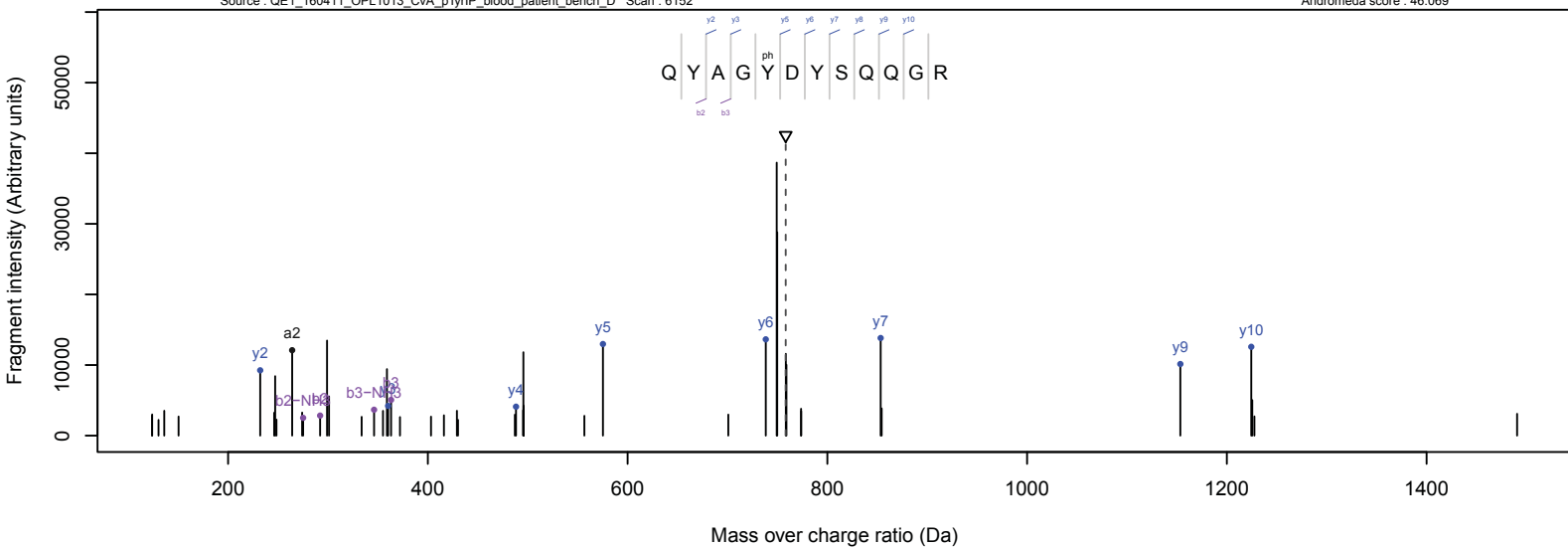




## QYAGY(ph)DYSQQGR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 6152

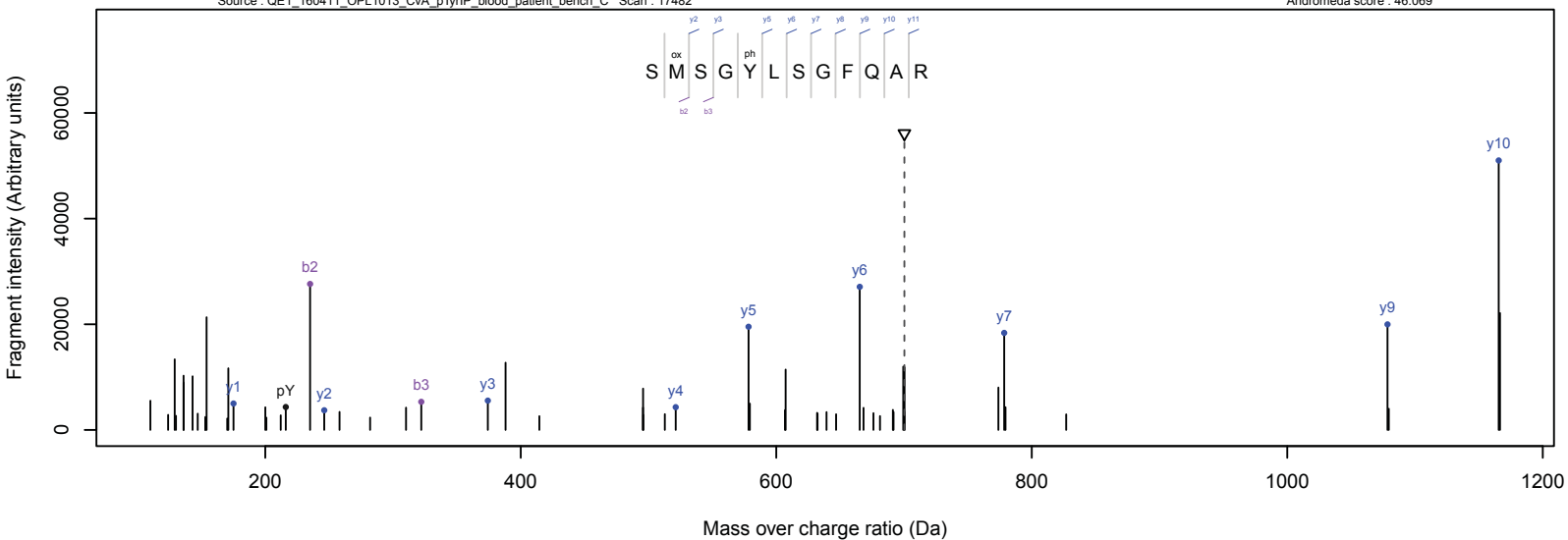
Andromeda score : 46.069



## SM(ox)SGY(ph)LSGFQAR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 17482

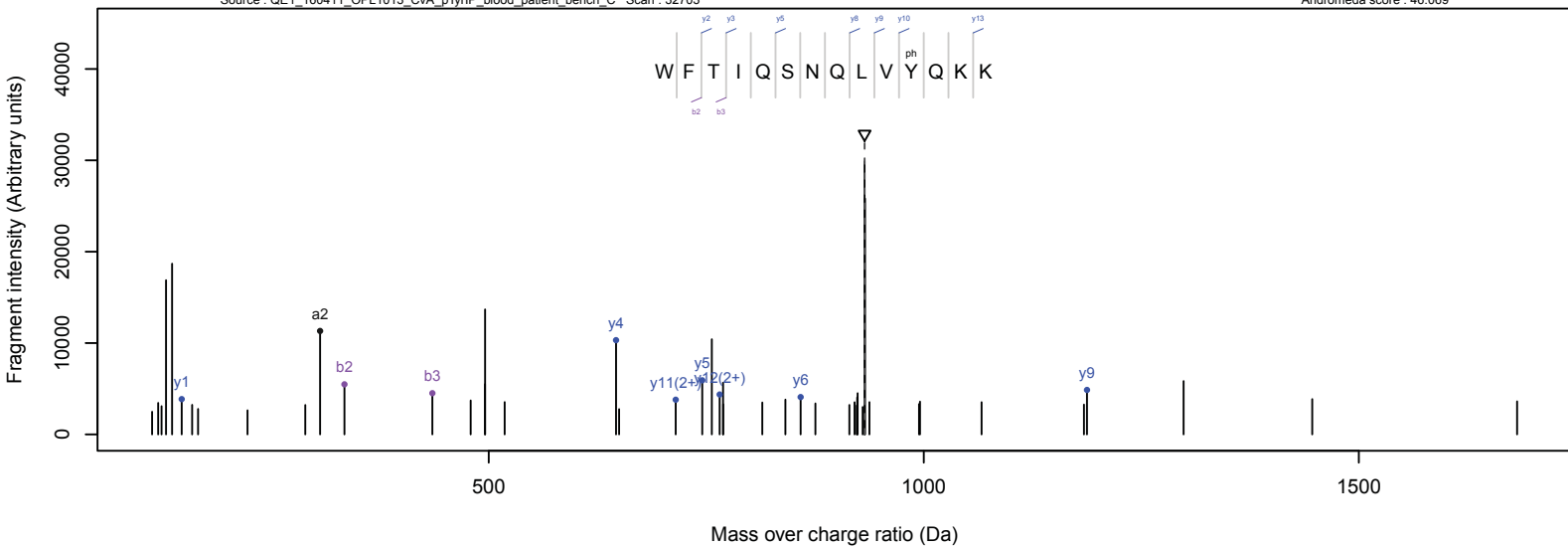
Andromeda score : 46.069



## WFTIQSNQLVY(ph)QKK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 32703

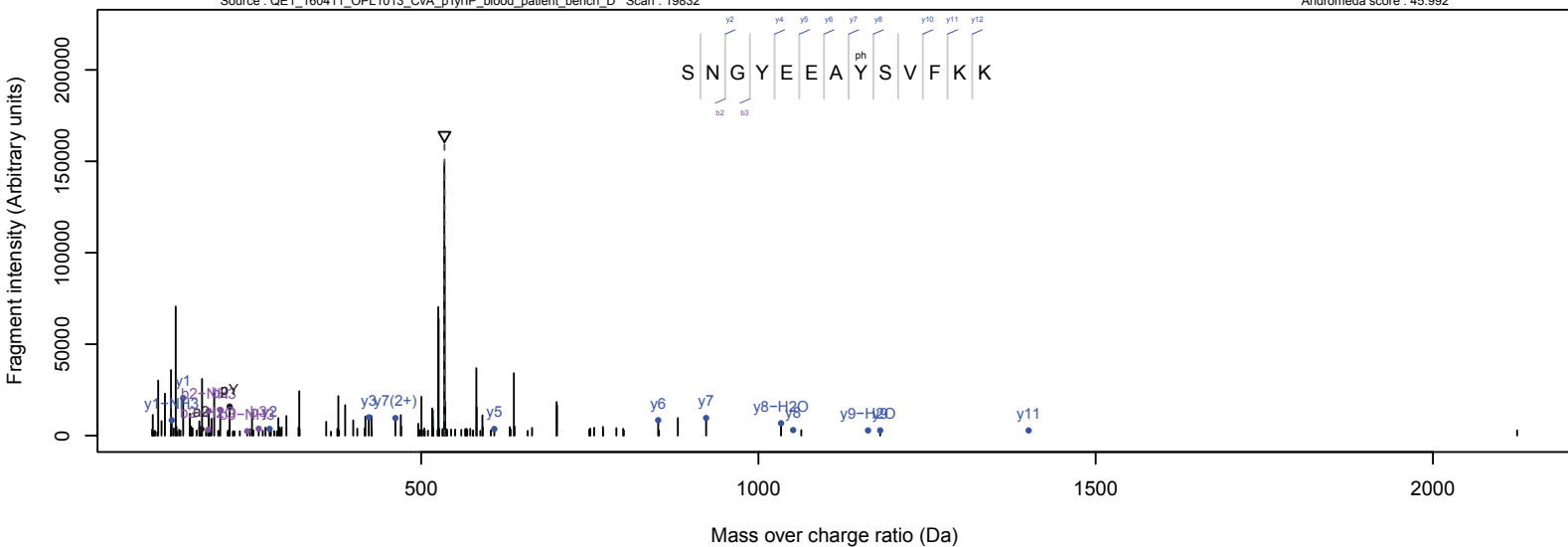
Andromeda score : 46.069



**SNGYEEAY(ph)SVFKK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 19832

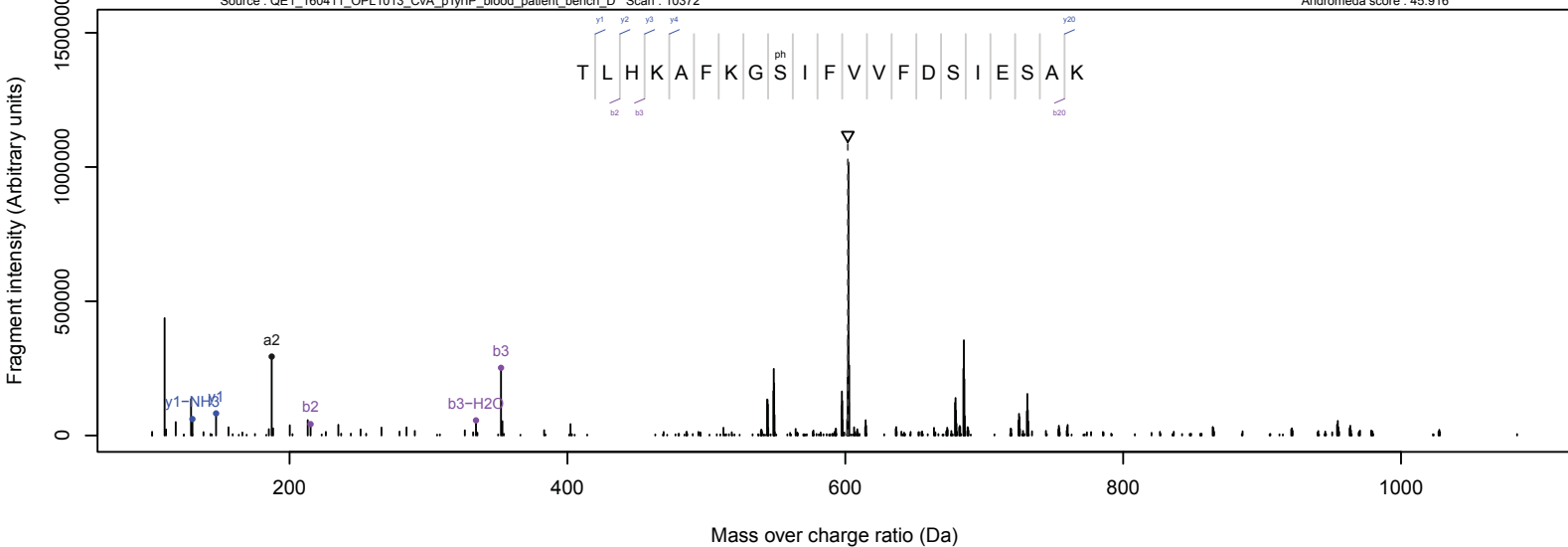
Andromeda score : 45.992



**TLHKAFKGS(ph)IFVVFDSIESAK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 10372

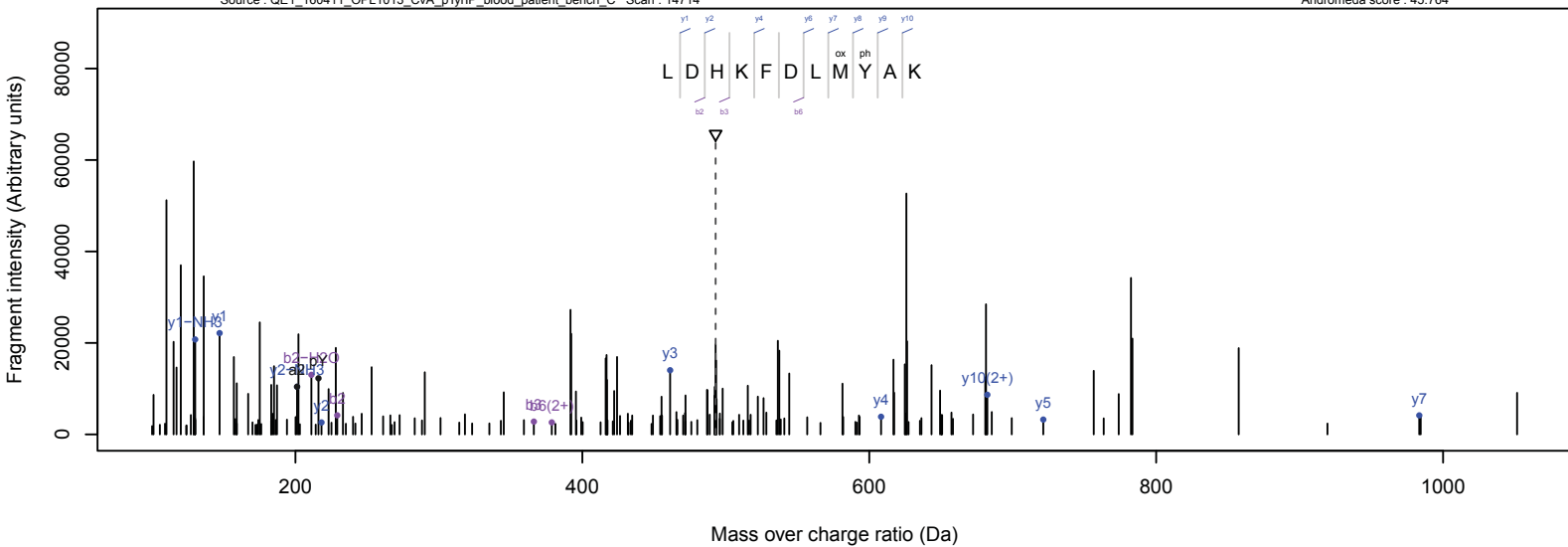
Andromeda score : 45.916



**LDHKFDLM(ox)Y(ph)AK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 14714

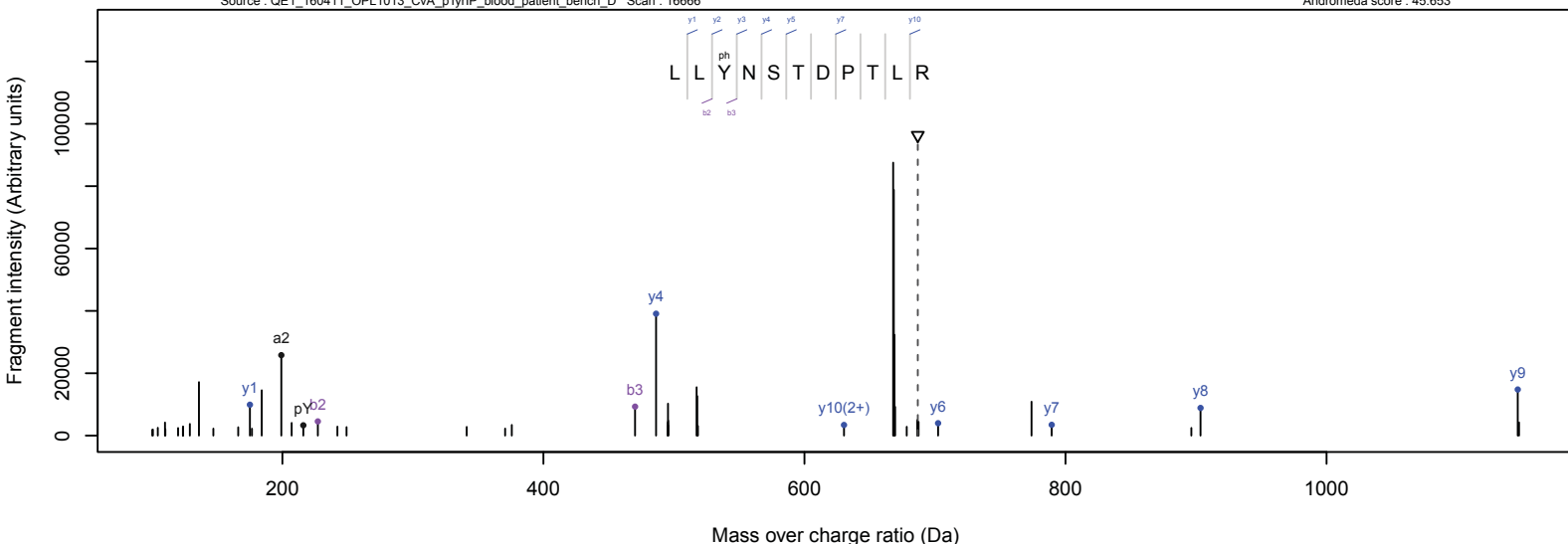
Andromeda score : 45.764



LLY(ph)NSTDPTLR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 16666

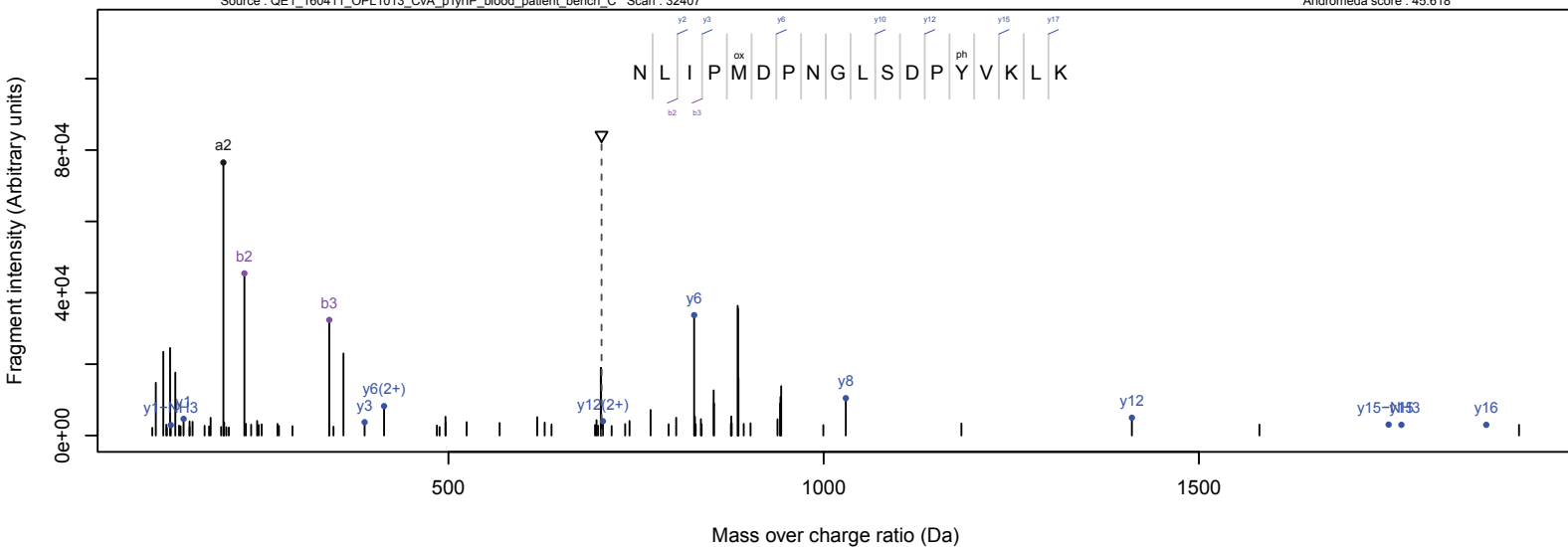
Andromeda score : 45.653



NLIPM(ox)DPNGLSDPY(ph)VKLK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 32407

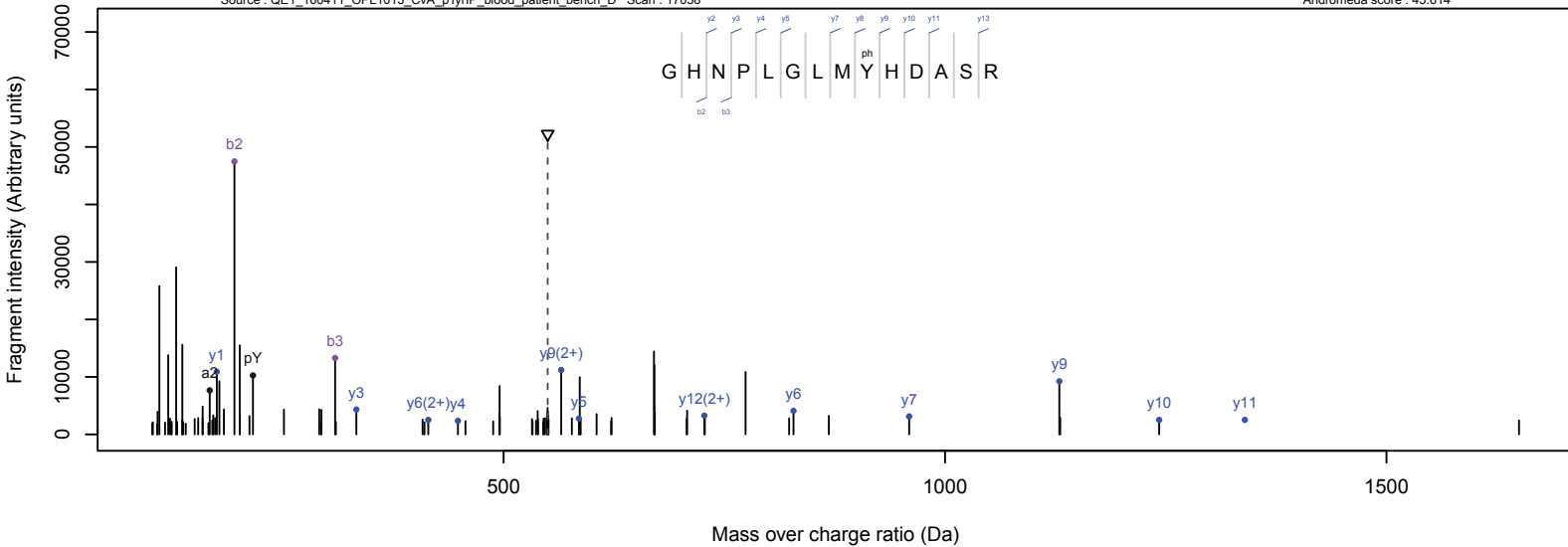
Andromeda score : 45.618



GHNPLGLMY(ph)HDASR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17058

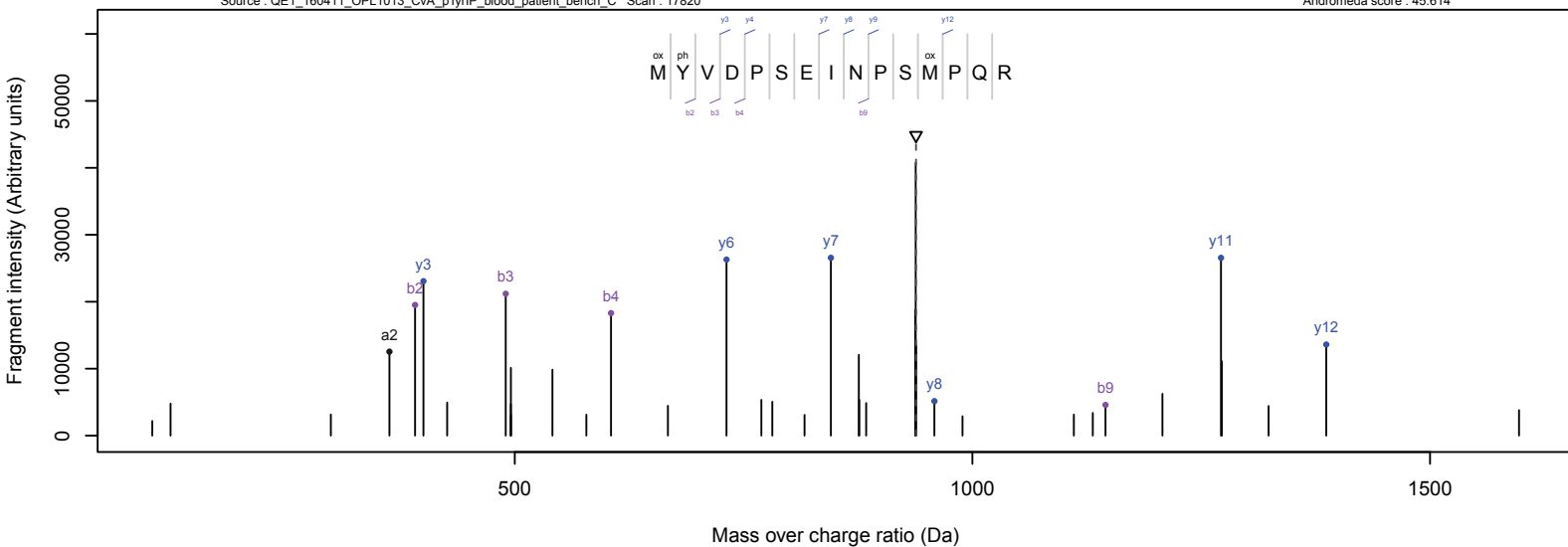
Andromeda score : 45.614



**M(ox)Y(ph)VDPSEINP<sup>ox</sup>SMPQR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 17820

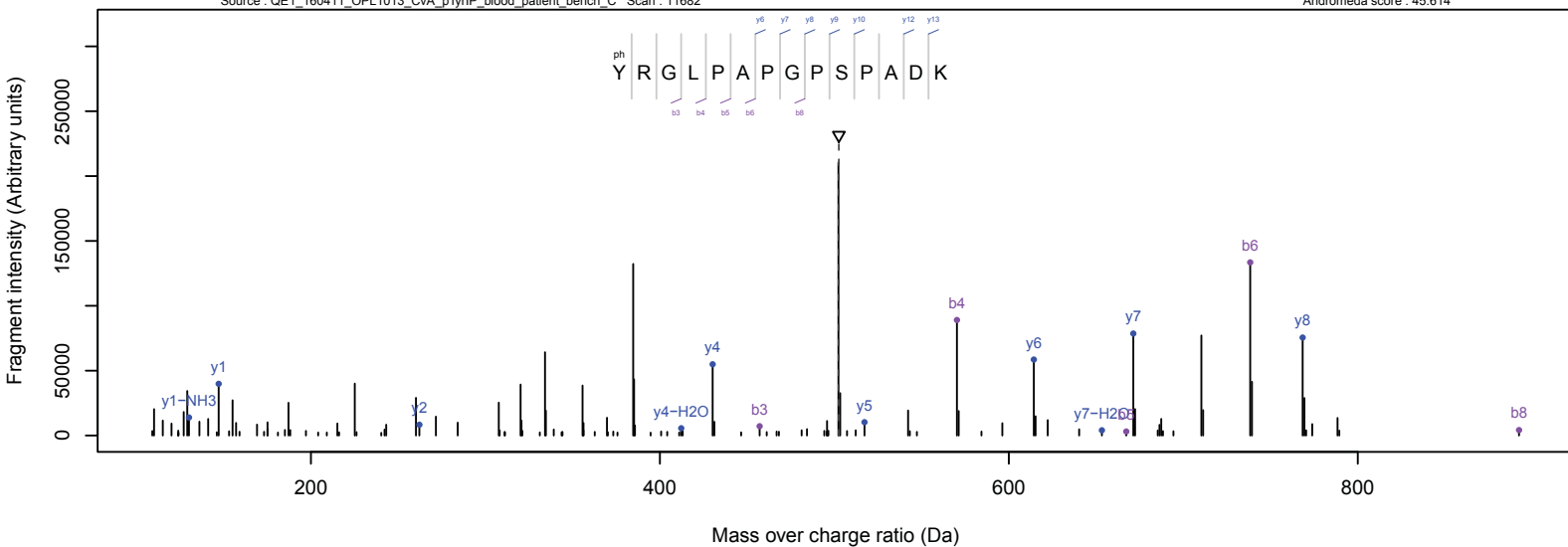
Andromeda score : 45.614



**Y(ph)RGLPAPGPSADK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 11682

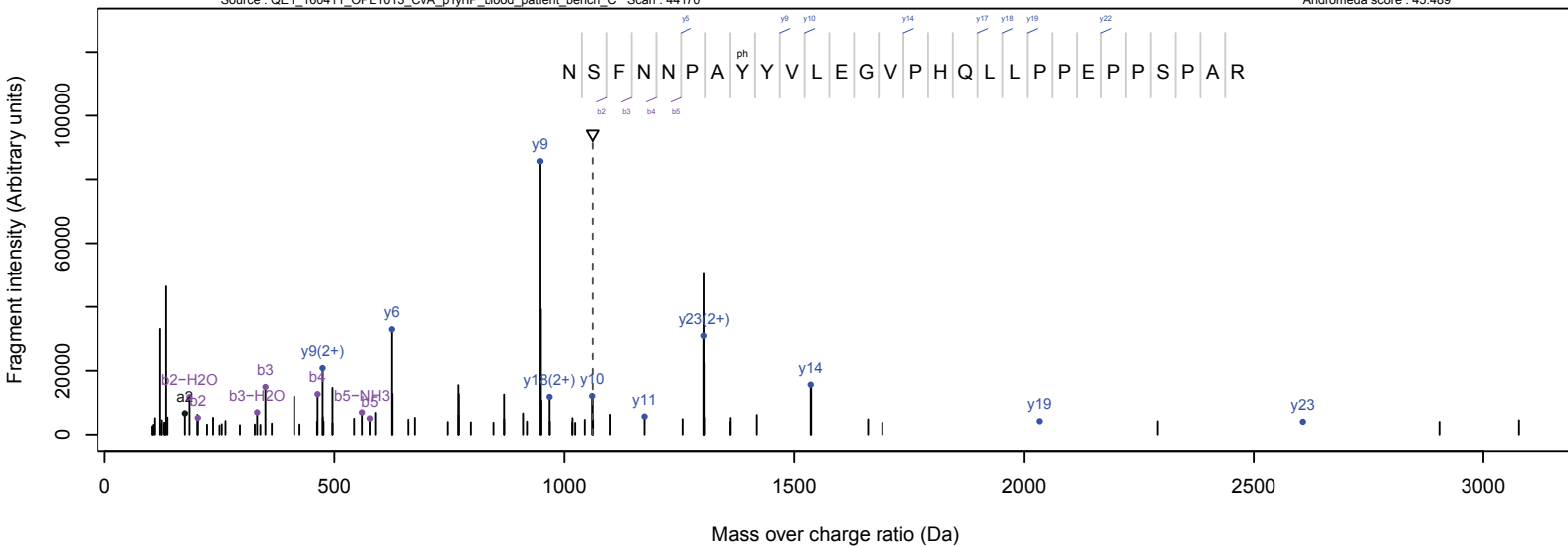
Andromeda score : 45.614



**NSFNNPAY(ph)YVLEGVPHQLLPPEPPSPAR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 44170

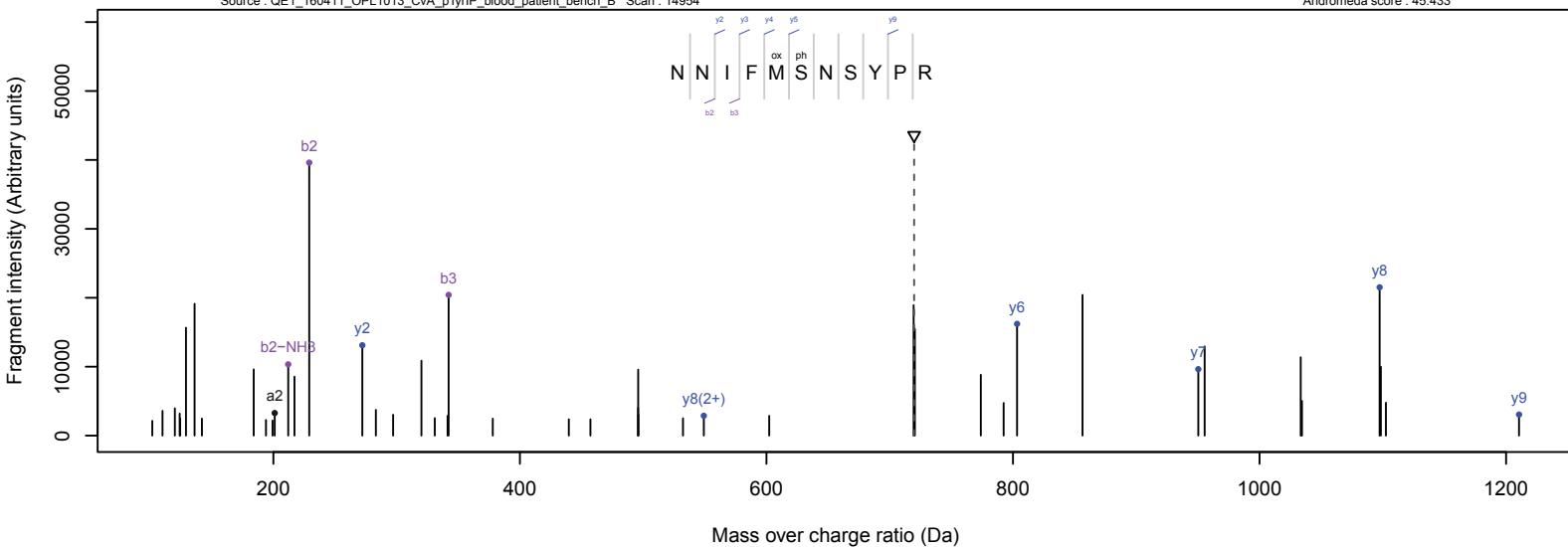
Andromeda score : 45.489



**NNIFM(ox)S(ph)NSYPR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 14954

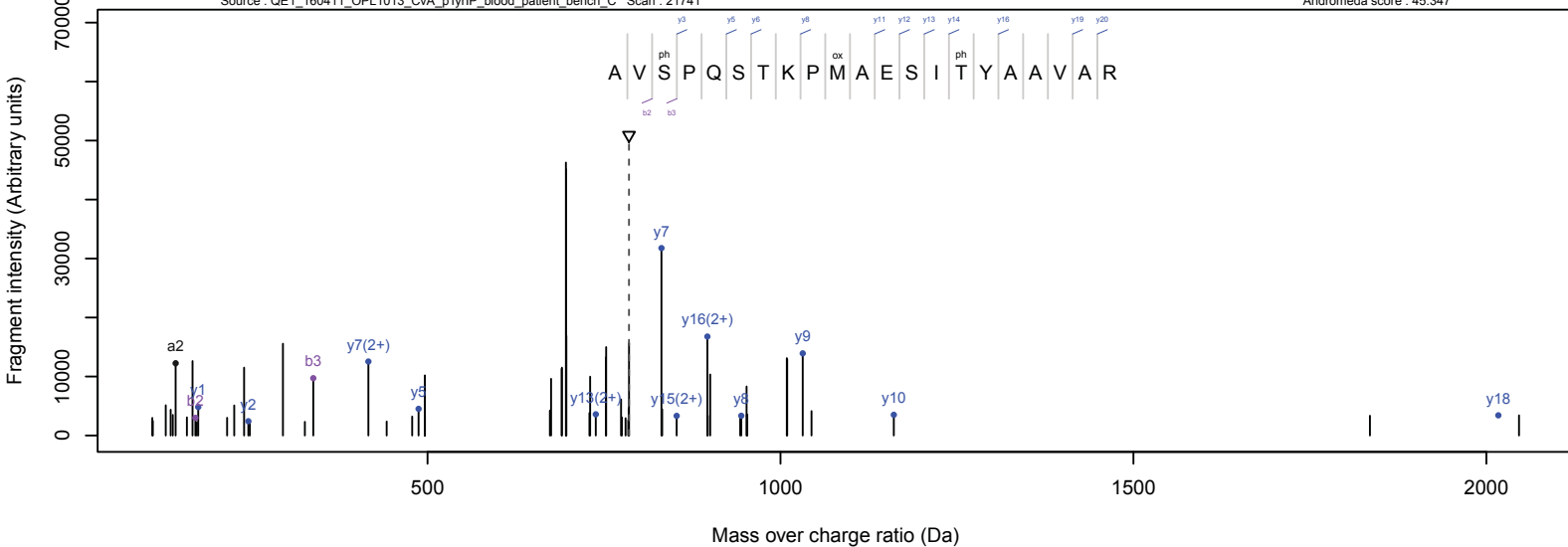
Andromeda score : 45.433



**AVS(ph)PQSTKPM(ox)AESIT(ph)YAAVAR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 21741

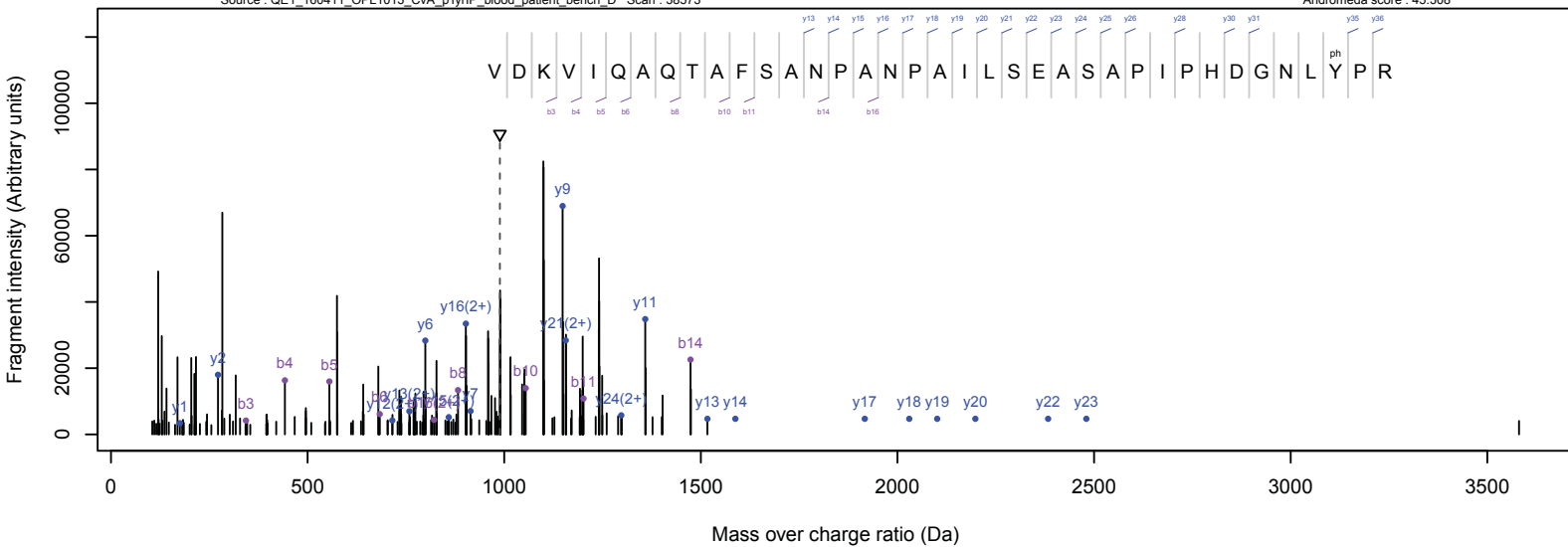
Andromeda score : 45.347



**VDKVIQAQTAFSANPANPAI LSEASAPIPHDGNLY(ph)PR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 38573

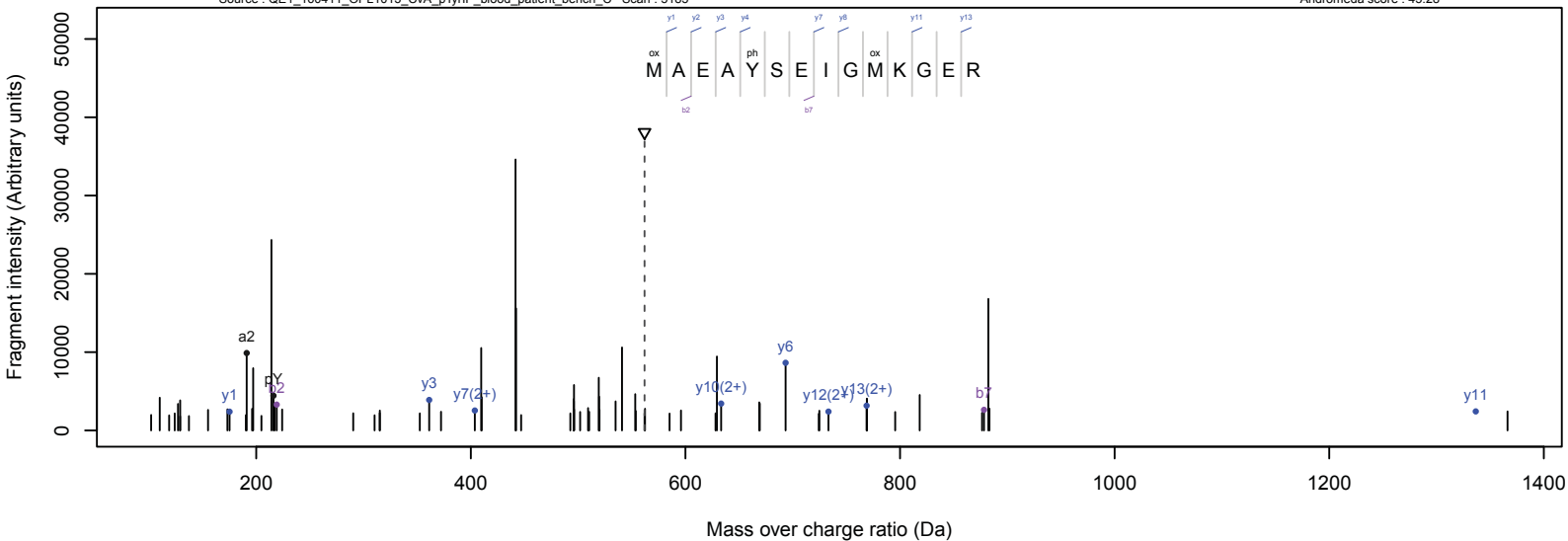
Andromeda score : 45.308



M(ox)AEAY(ph)SEIGMKGER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 5185

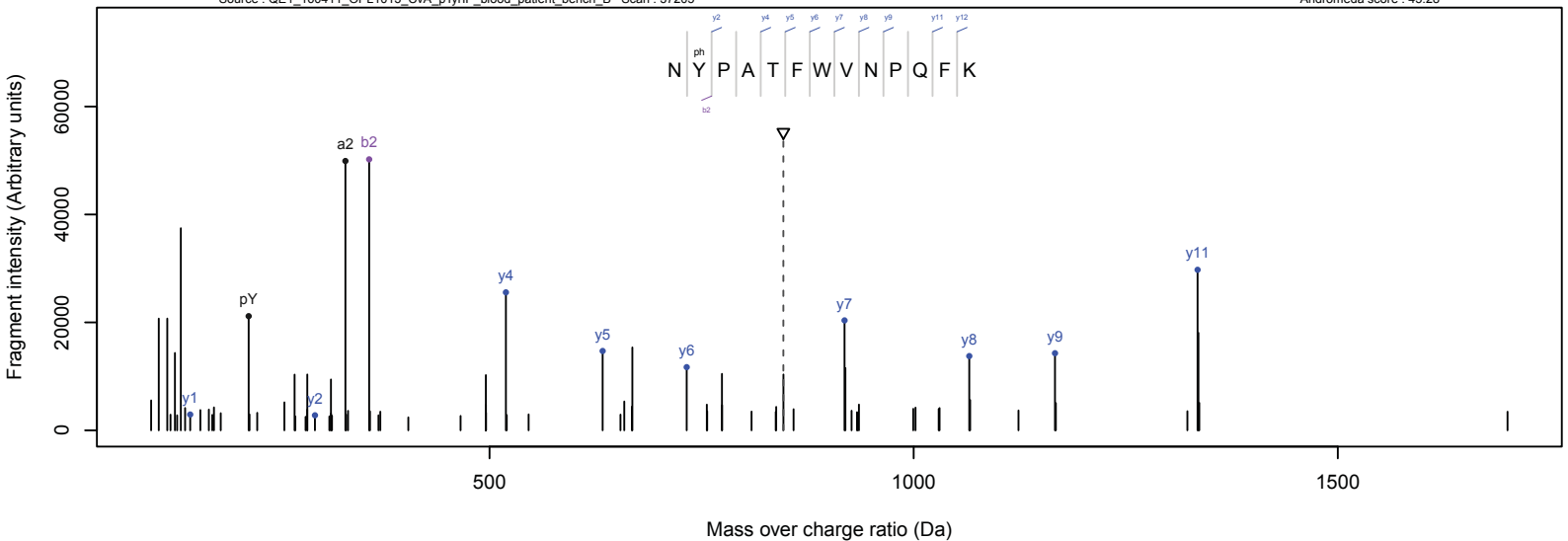
Andromeda score : 45.28



NY(ph)PATFWVNPQFK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 37205

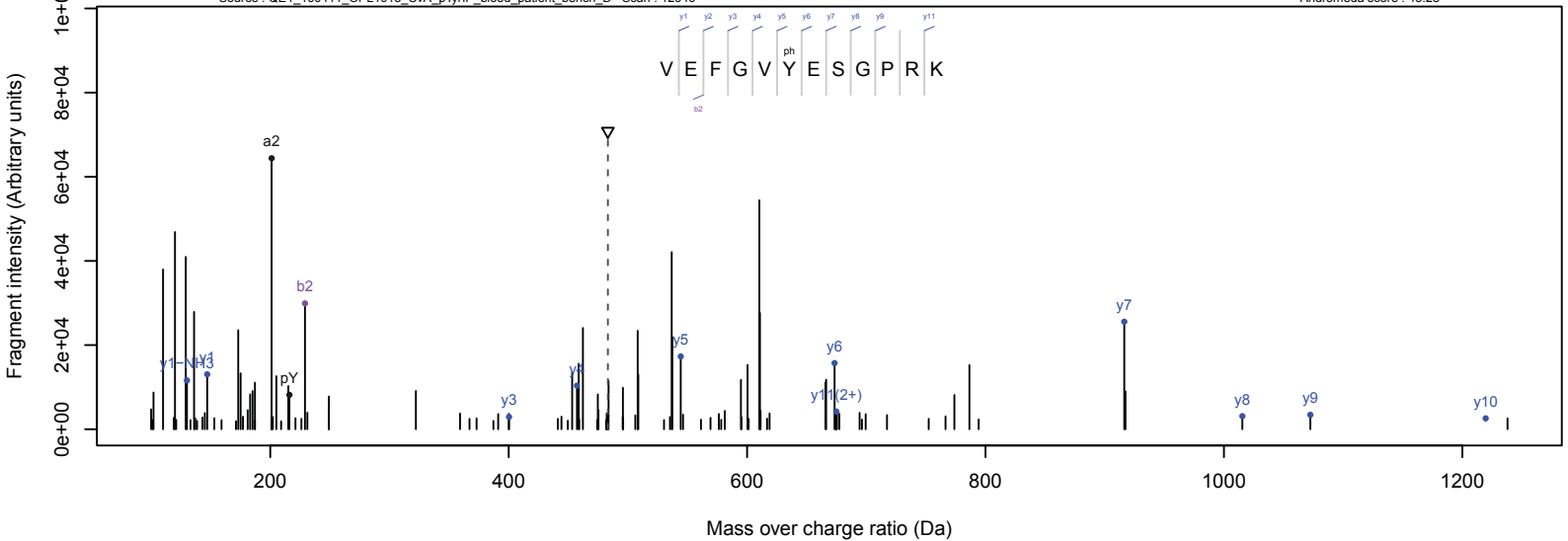
Andromeda score : 45.28



VEFGVY(ph)ESGPRK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12946

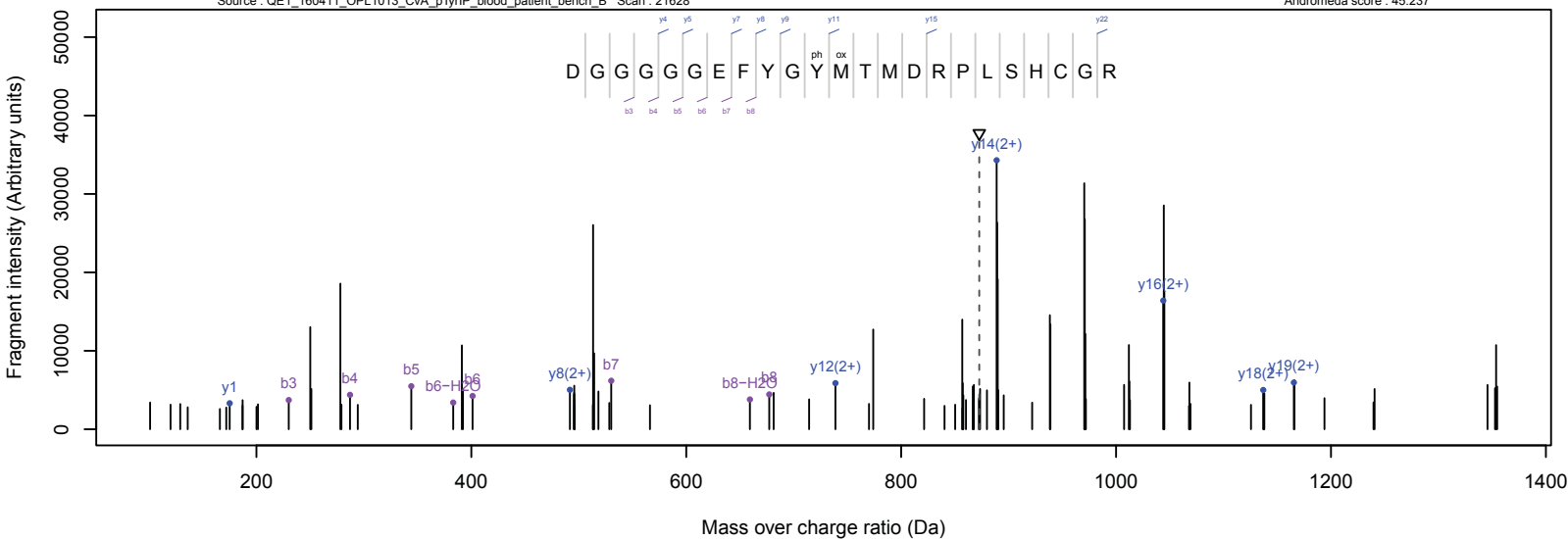
Andromeda score : 45.28



DGGGGGEFYGYM(ox)TMDRPLSHCGR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 21628

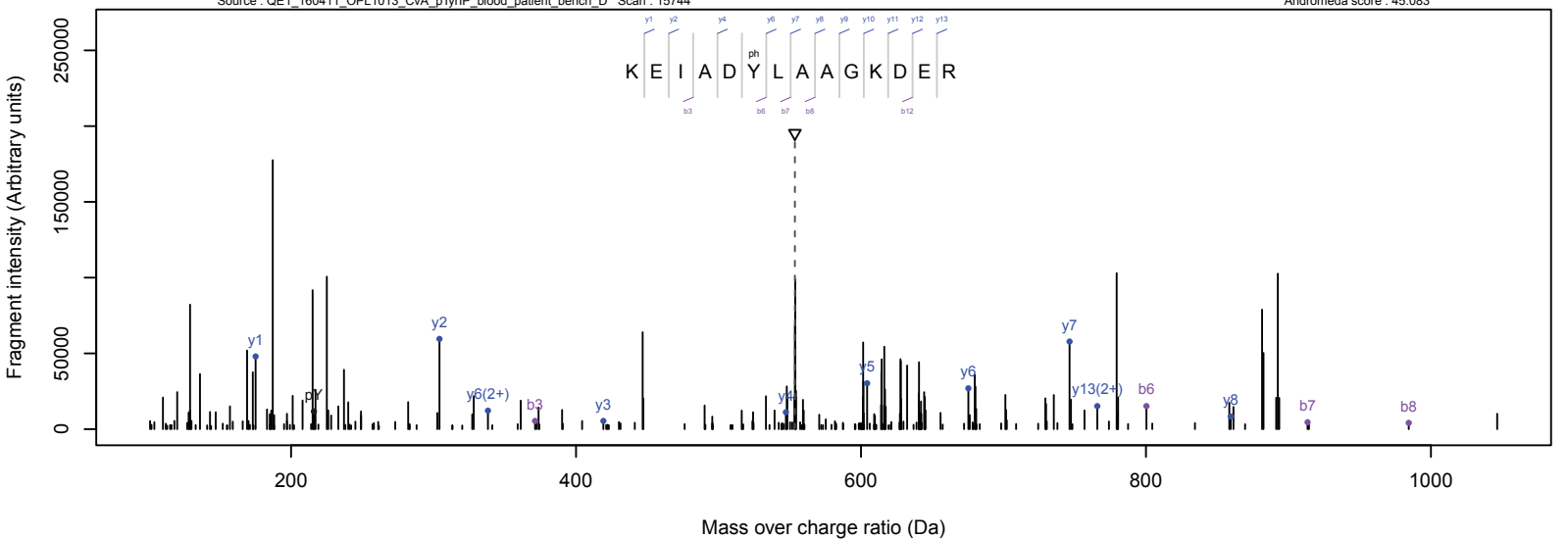
Andromeda score : 45.237



KEIADYLAAGKDER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 15744

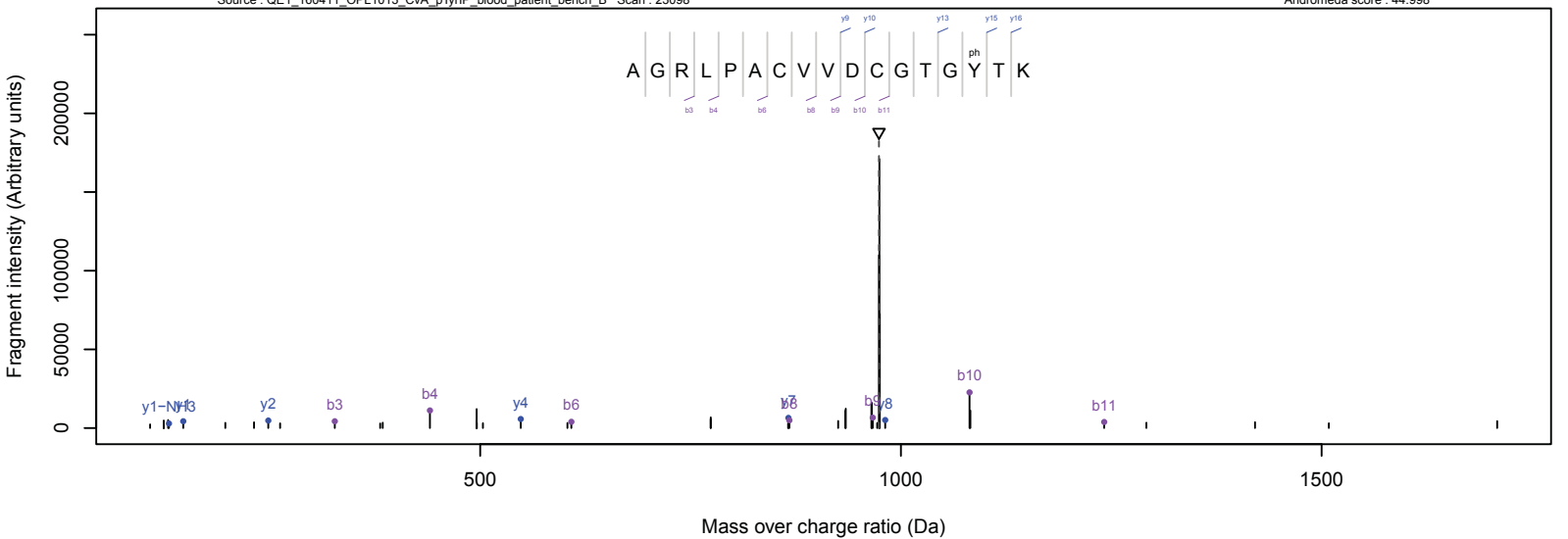
Andromeda score : 45.083



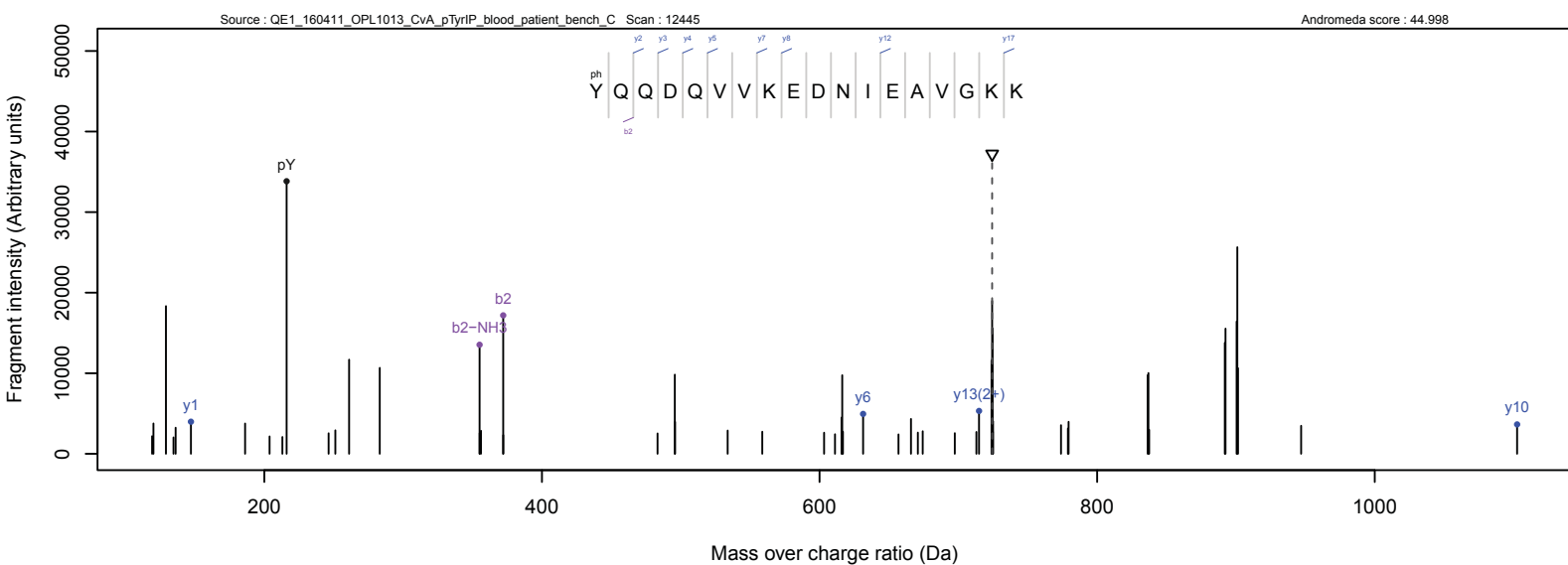
(ac)AGRLPACVVDCGTGYTK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 23098

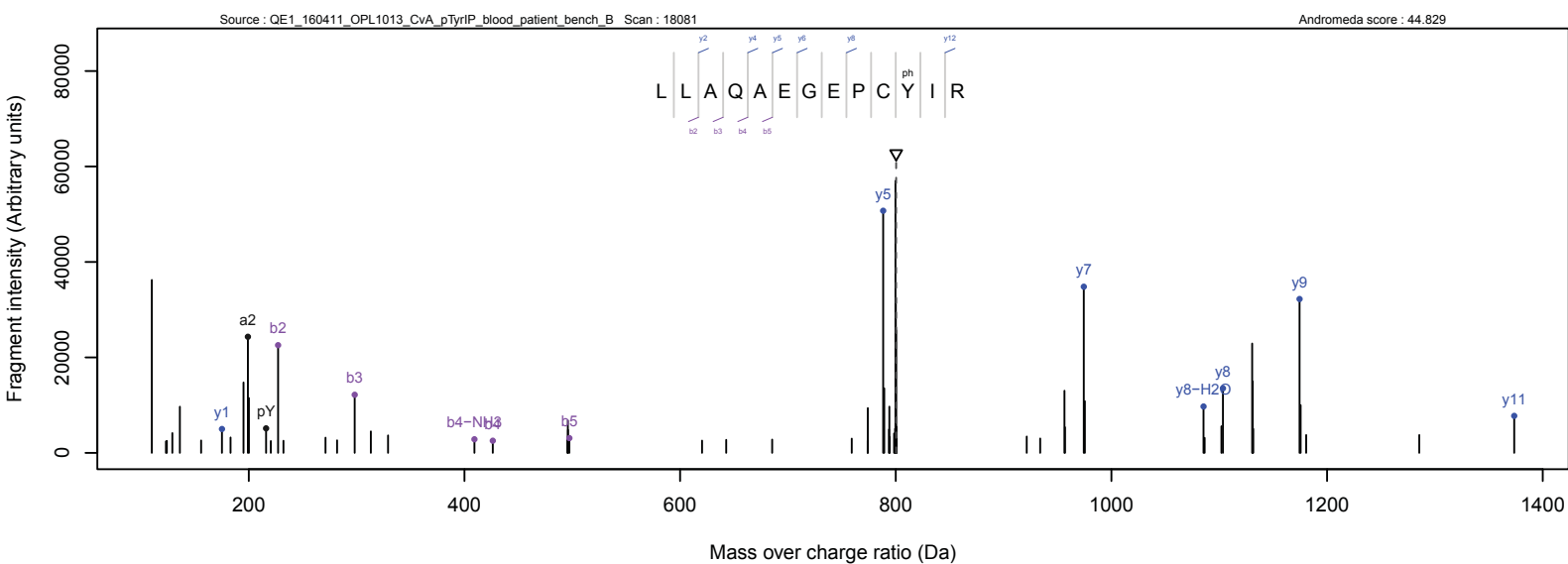
Andromeda score : 44.998



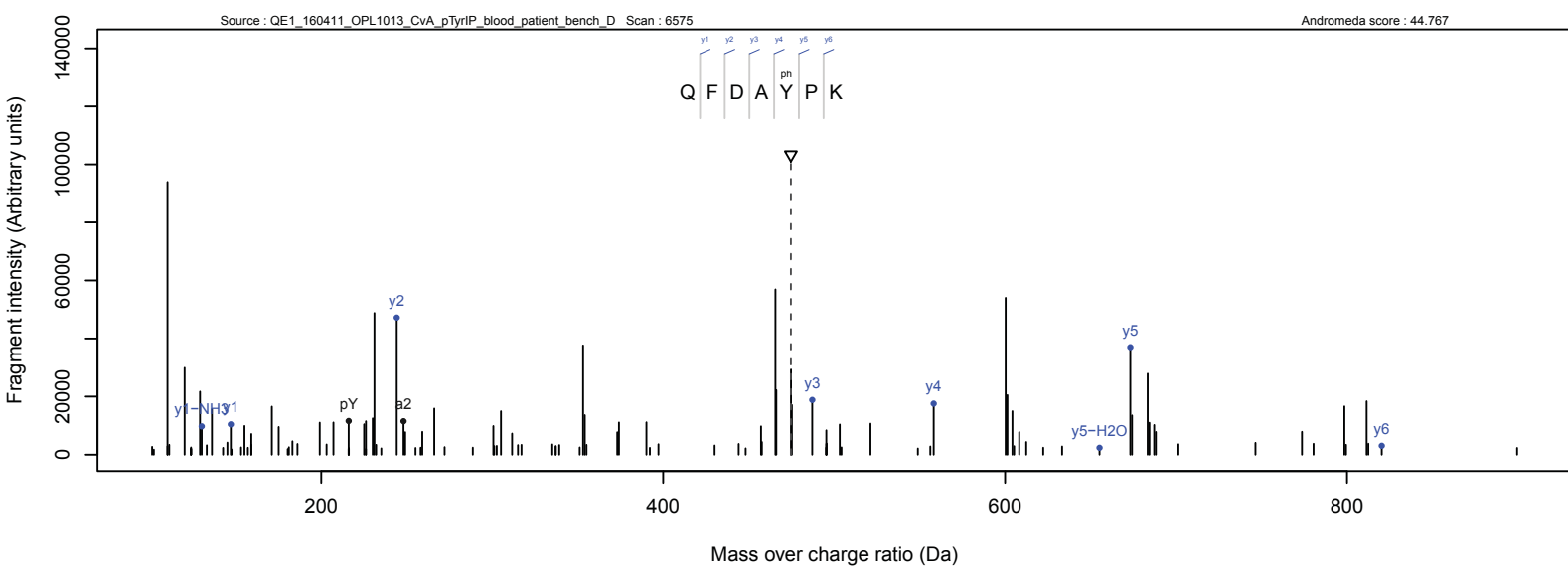
## Y(ph)QQDQVVKEDNIEAVGKK-3+



## LLAQAEGPCY(ph)IR-2+



## QFDAY(ph)PK-2+

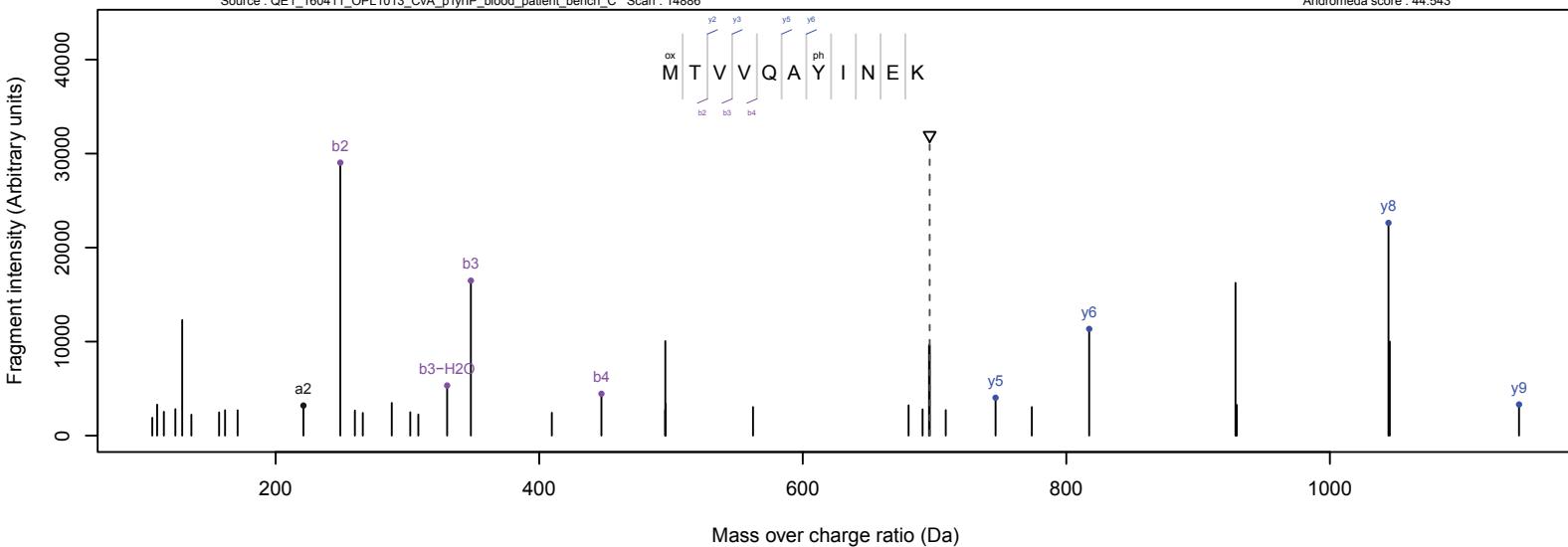




## M(ox)TVVQAY(ph)INEK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 14886

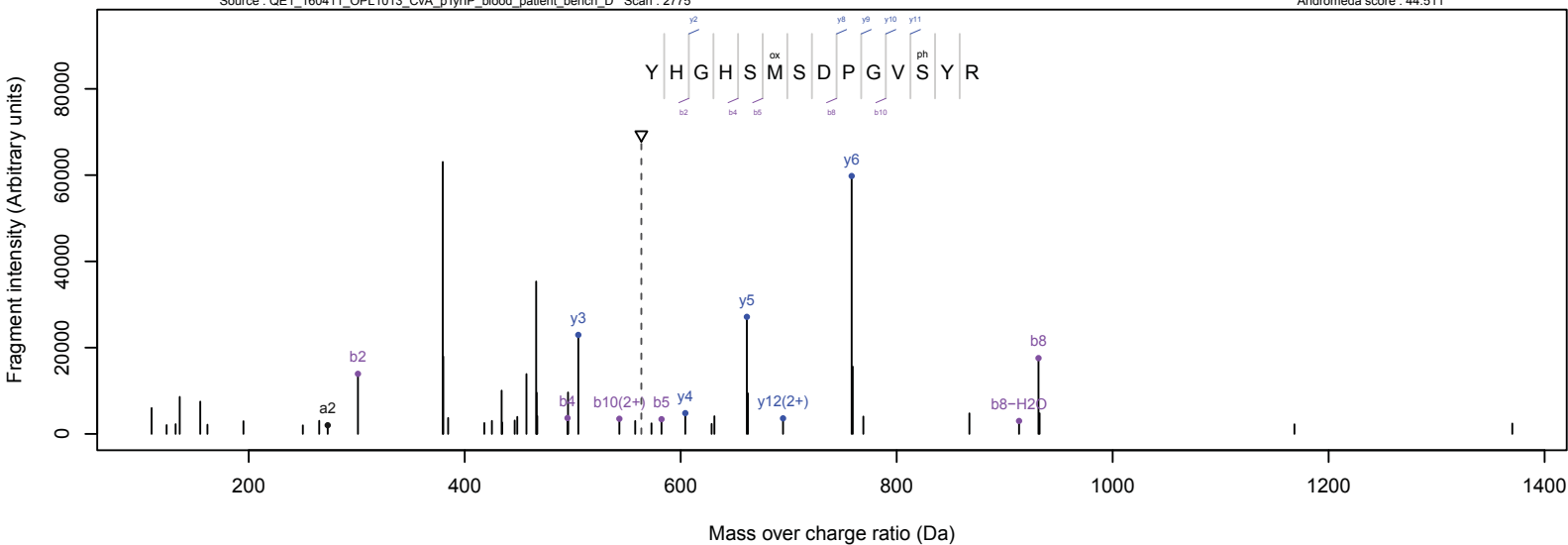
Andromeda score : 44.543



## YHGHSM(ox)SDPGVS(ph)YR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 2775

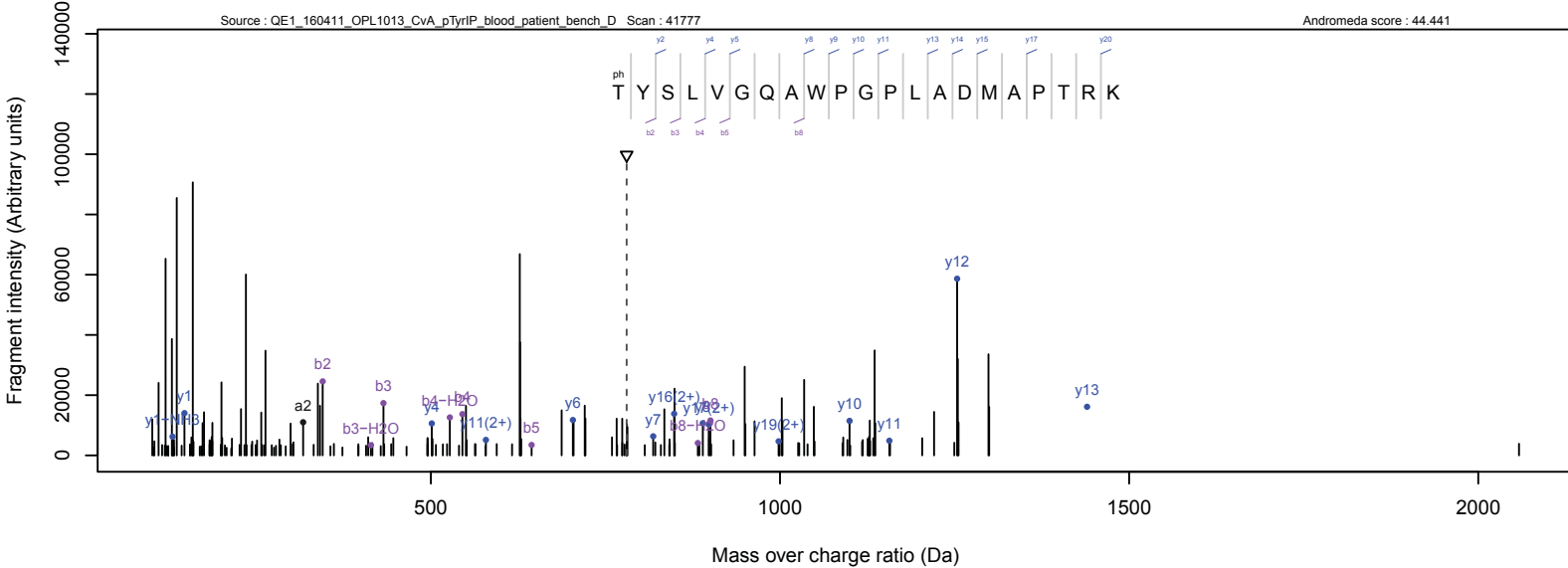
Andromeda score : 44.511



## T(ph)YSLVGAWPGPLADMAPTRK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 41777

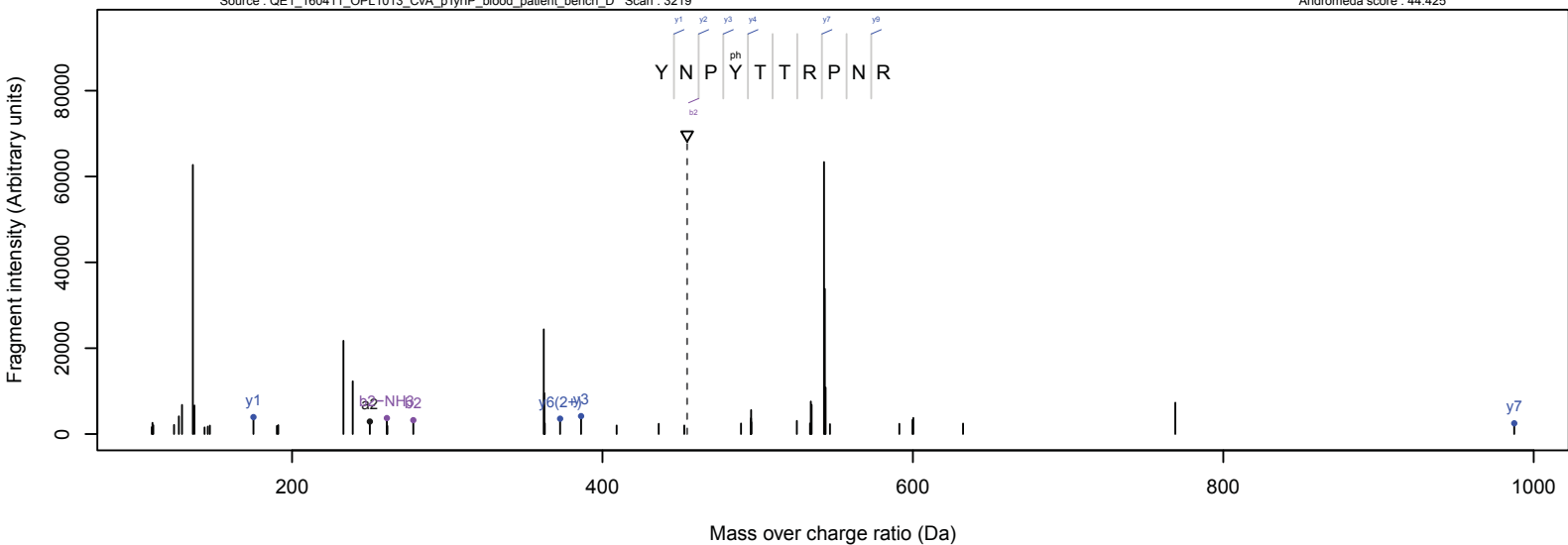
Andromeda score : 44.441



YNPY(ph)TTRPNR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 3219

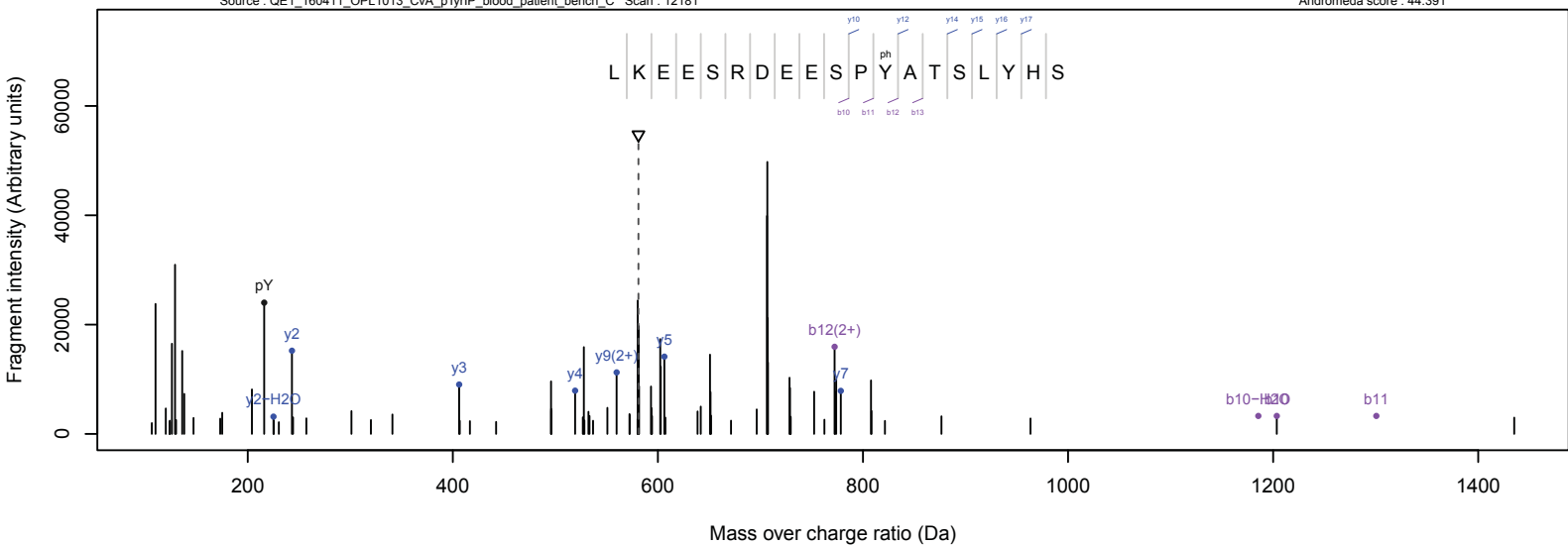
Andromeda score : 44.425



LKEESRDEESPY(ph)ATSLYHS-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 12181

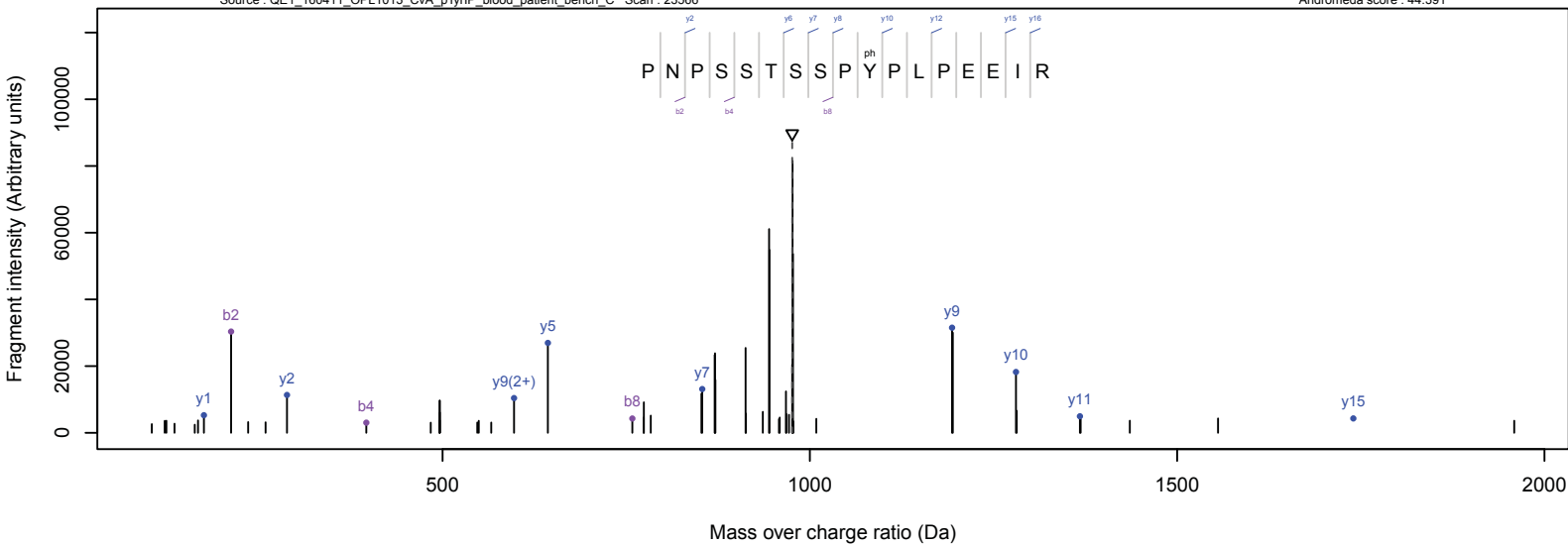
Andromeda score : 44.391



PNPSTSSPY(ph)PLPEEIR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 23566

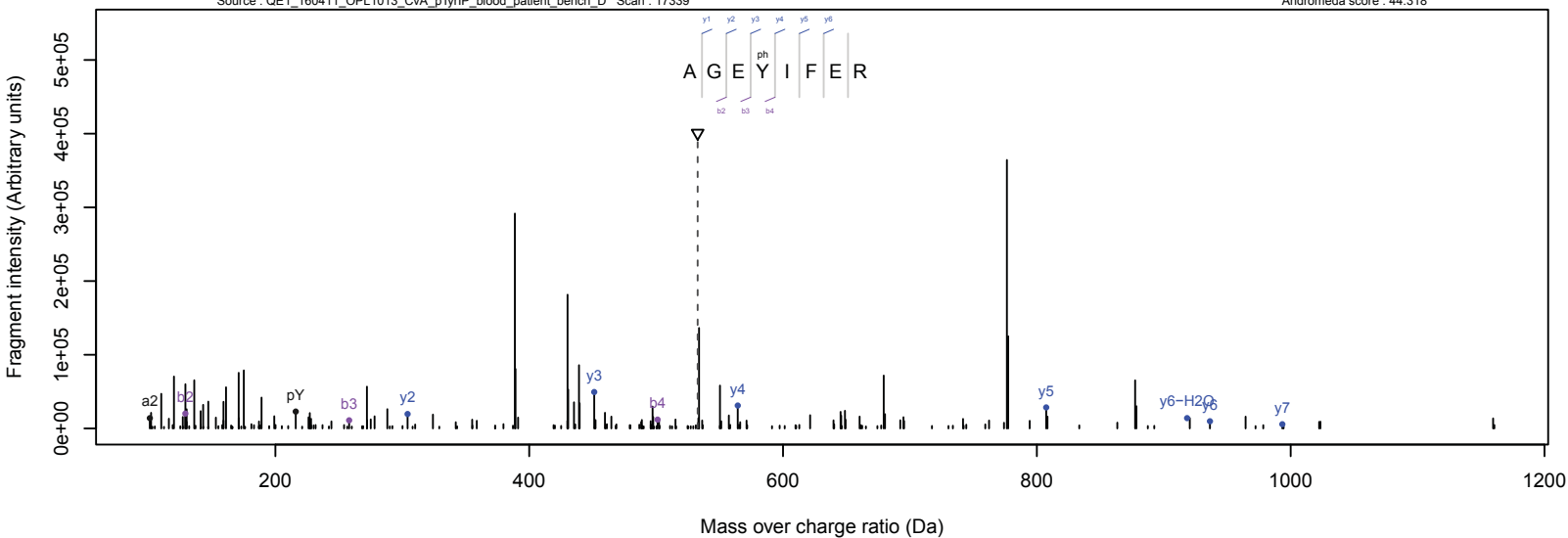
Andromeda score : 44.391



AGEY(ph)IFER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 17339

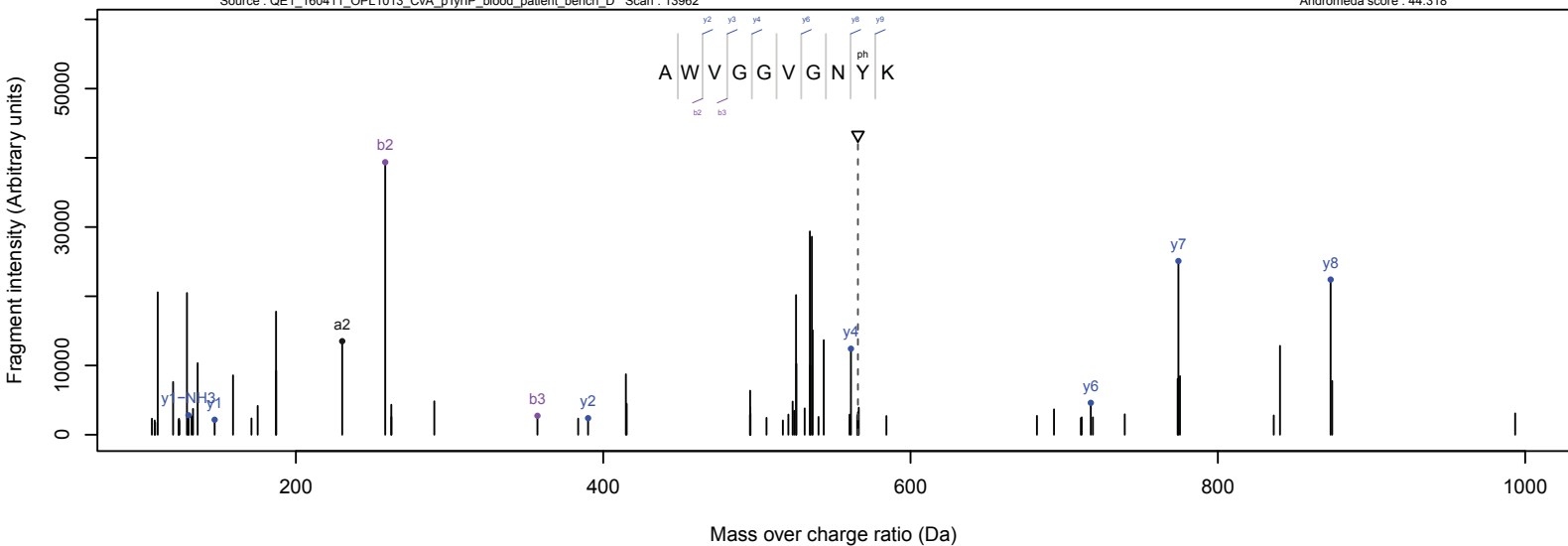
Andromeda score : 44.318



AWVGGVGN(ph)K-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 13962

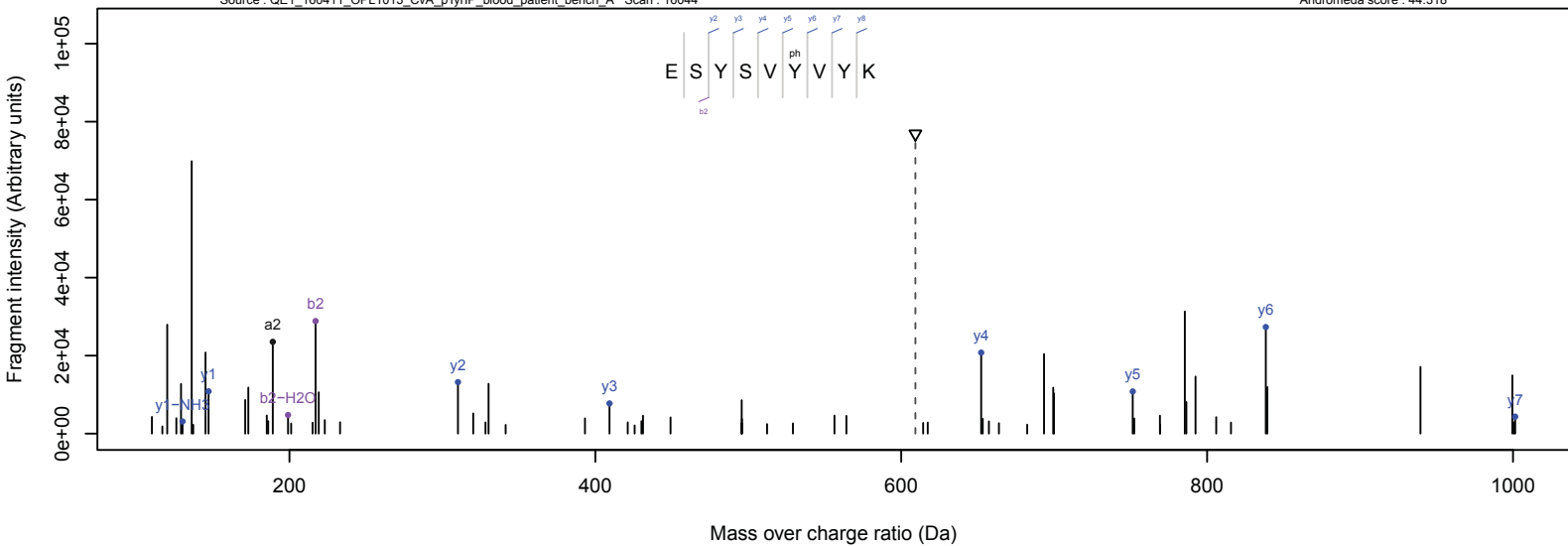
Andromeda score : 44.318



ESYSVY(ph)VYK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 16044

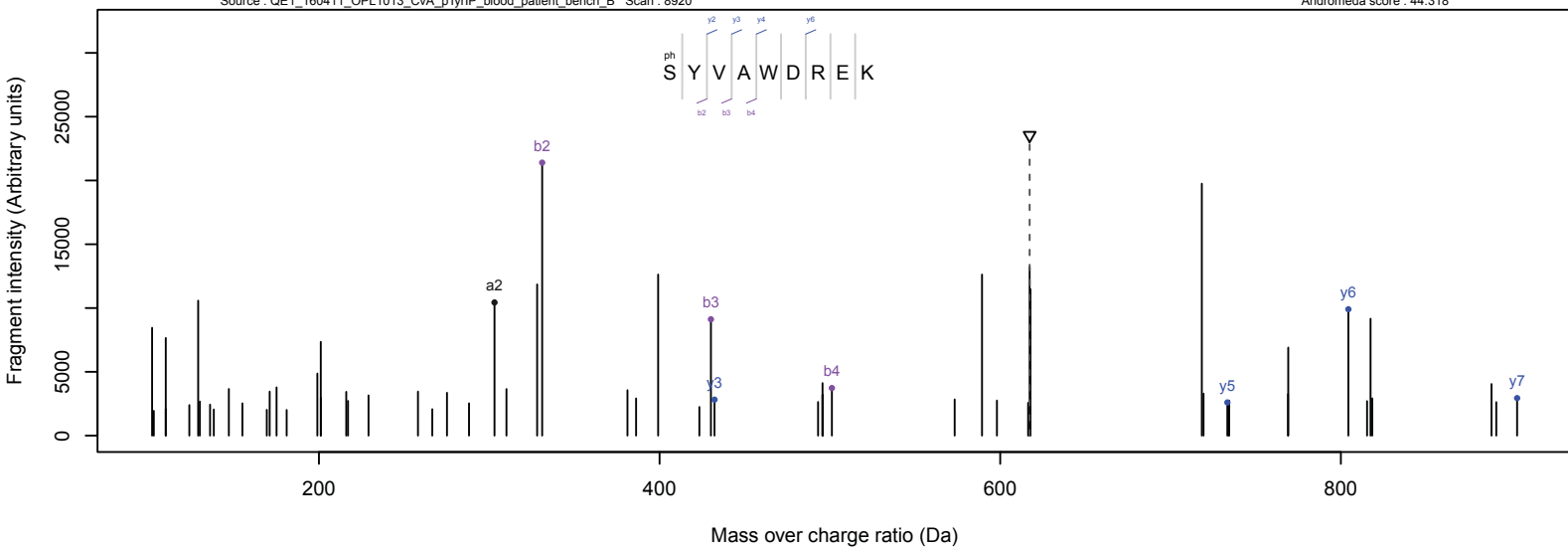
Andromeda score : 44.318



**S(ph)YVAWDREK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 8920

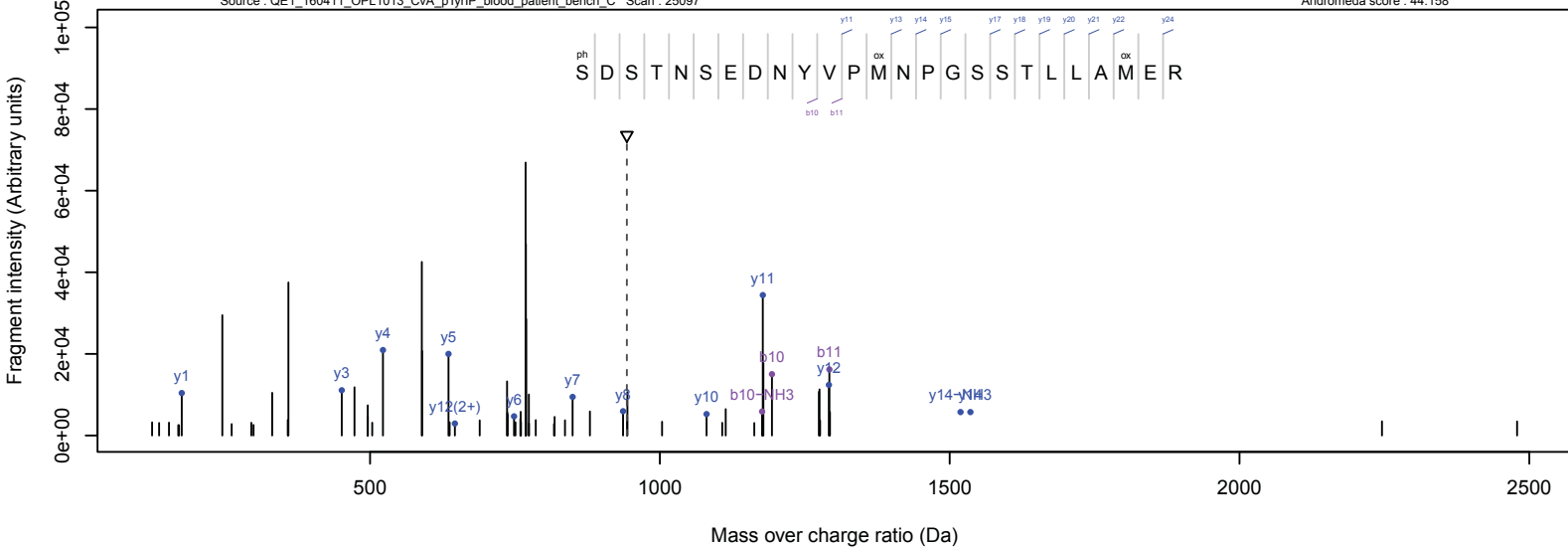
Andromeda score : 44.318



**S(ph)DSTNSEDNYPVPMNPGSSSTLLAMER-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 25097

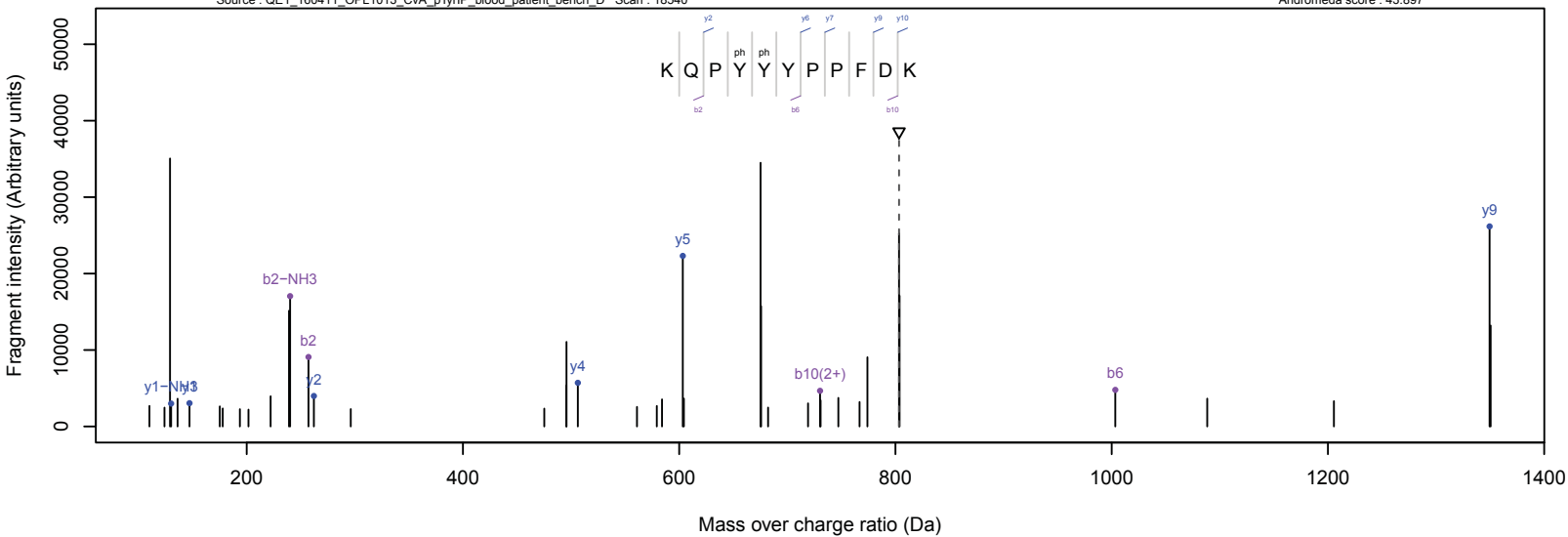
Andromeda score : 44.158



**KQPY(ph)YPPFDK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 18540

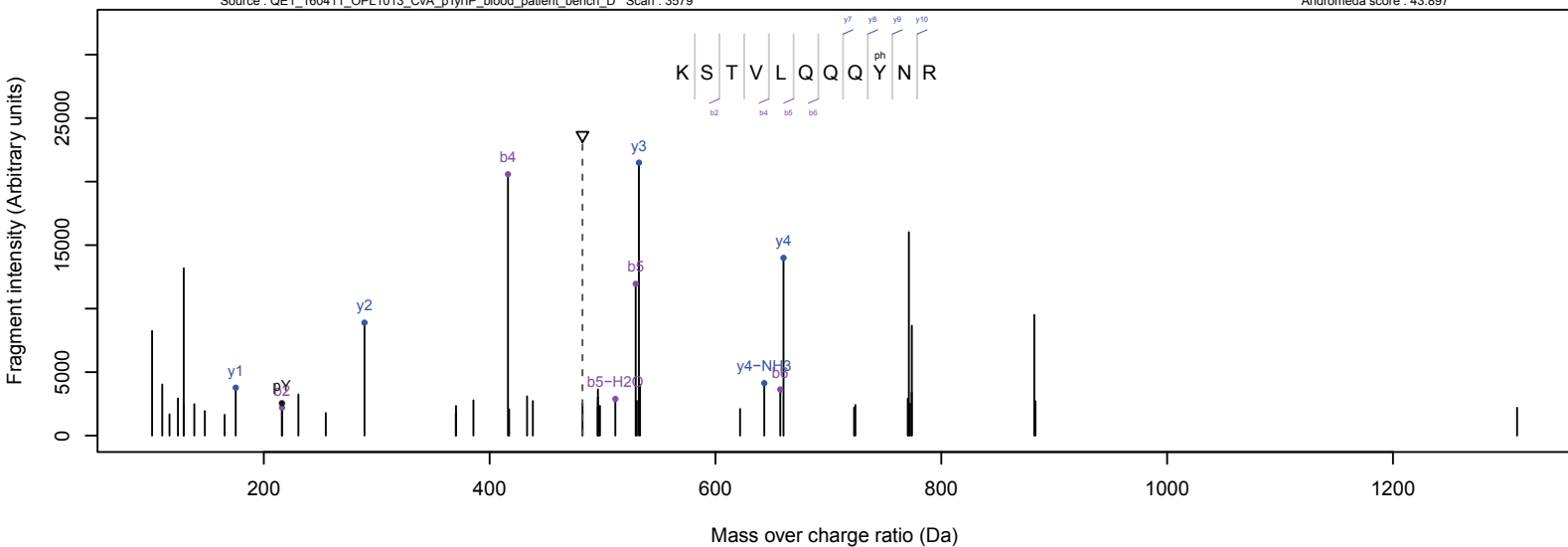
Andromeda score : 43.897



**KSTVLQQQY(ph)NR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 3579

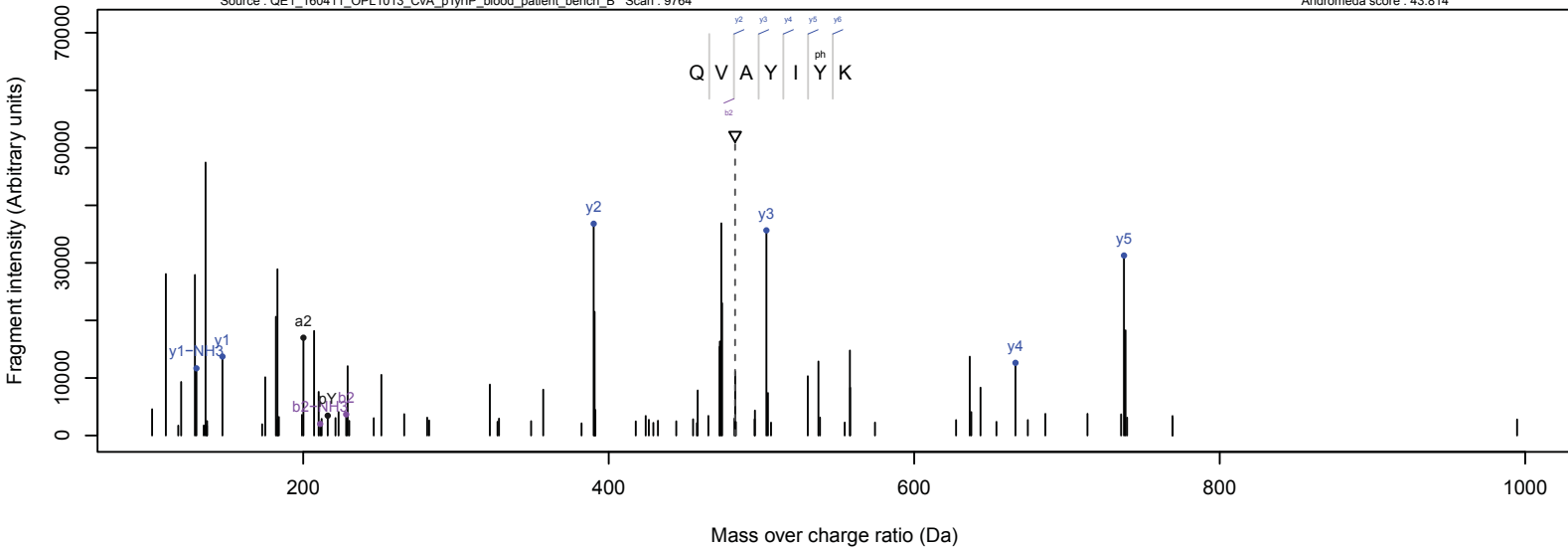
Andromeda score : 43.897



**QVAYIY(ph)K-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 9764

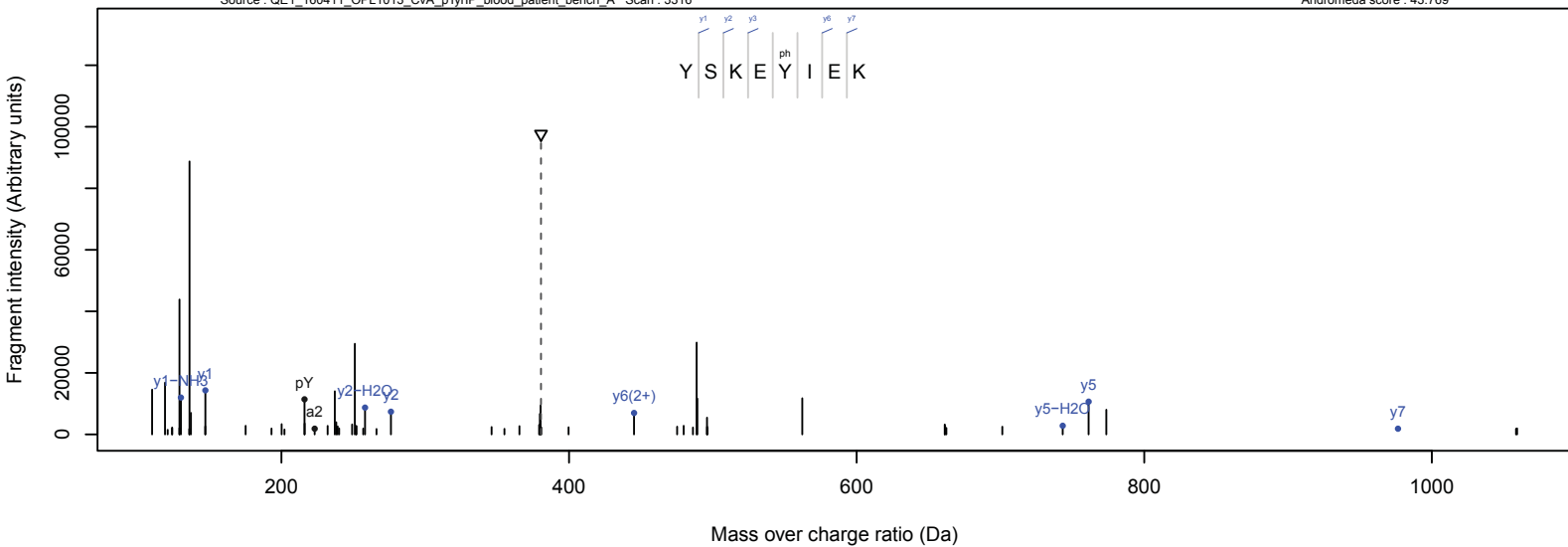
Andromeda score : 43.814



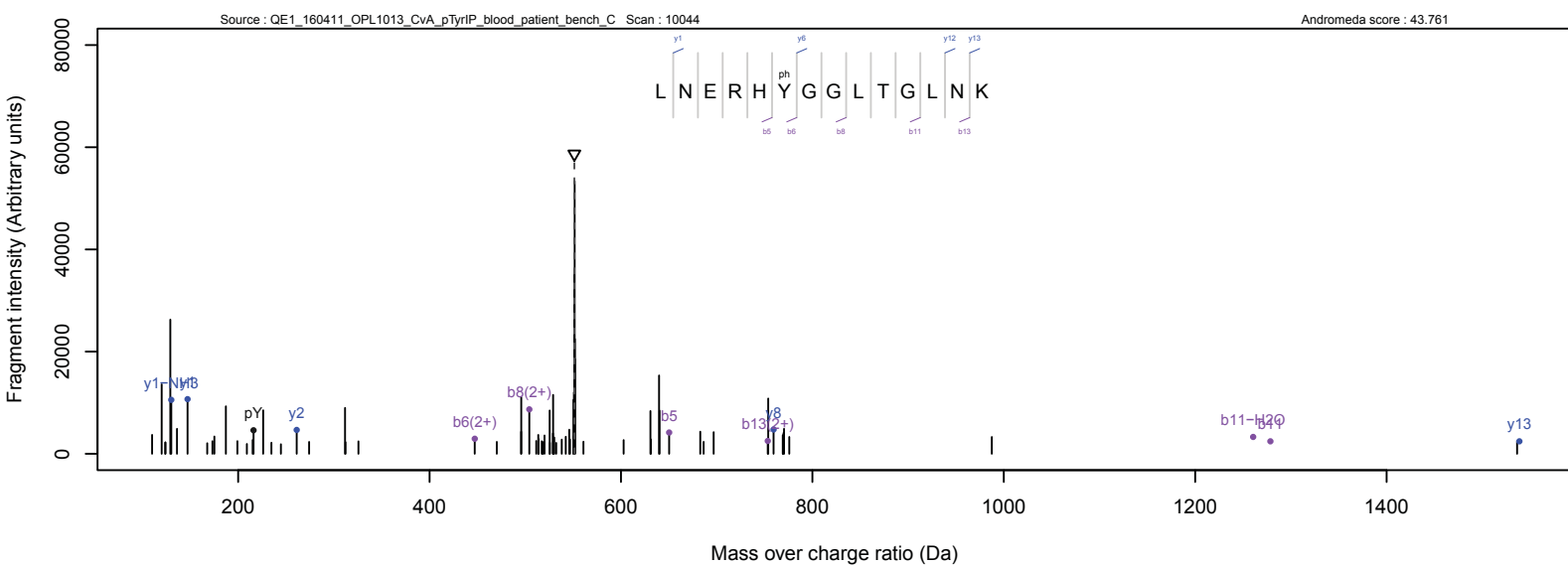
**YSKEY(ph)IEK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 3316

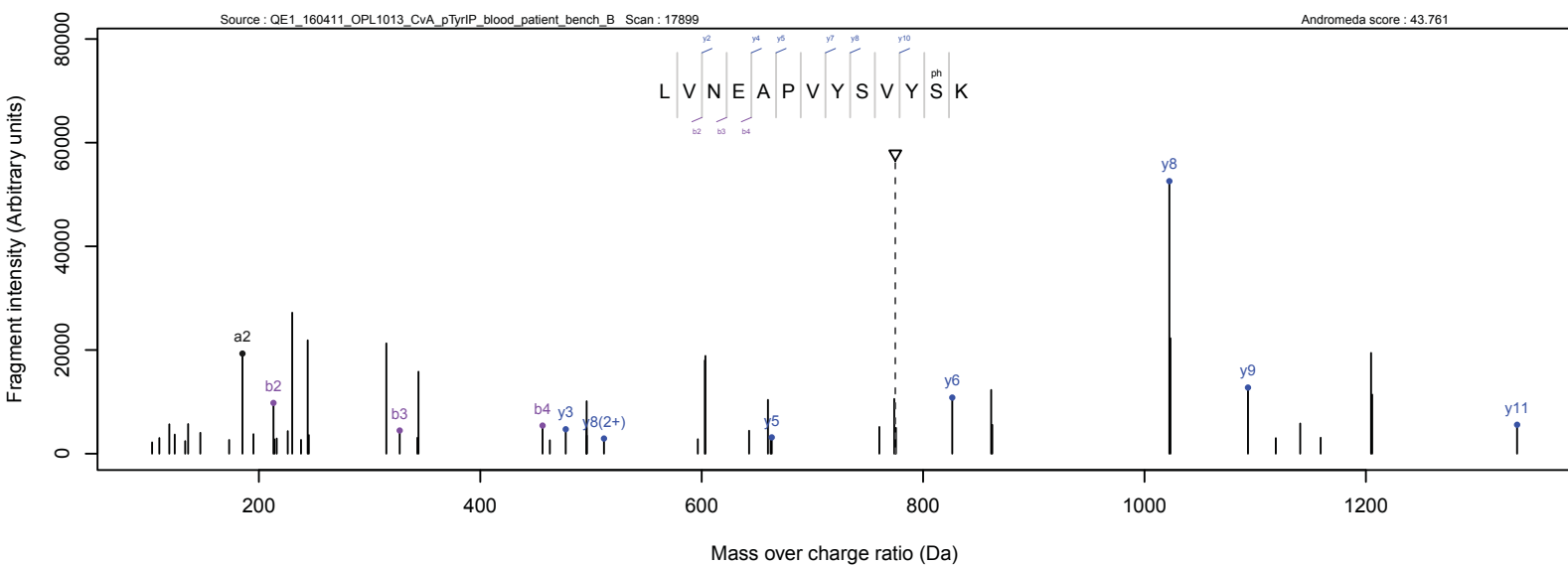
Andromeda score : 43.769



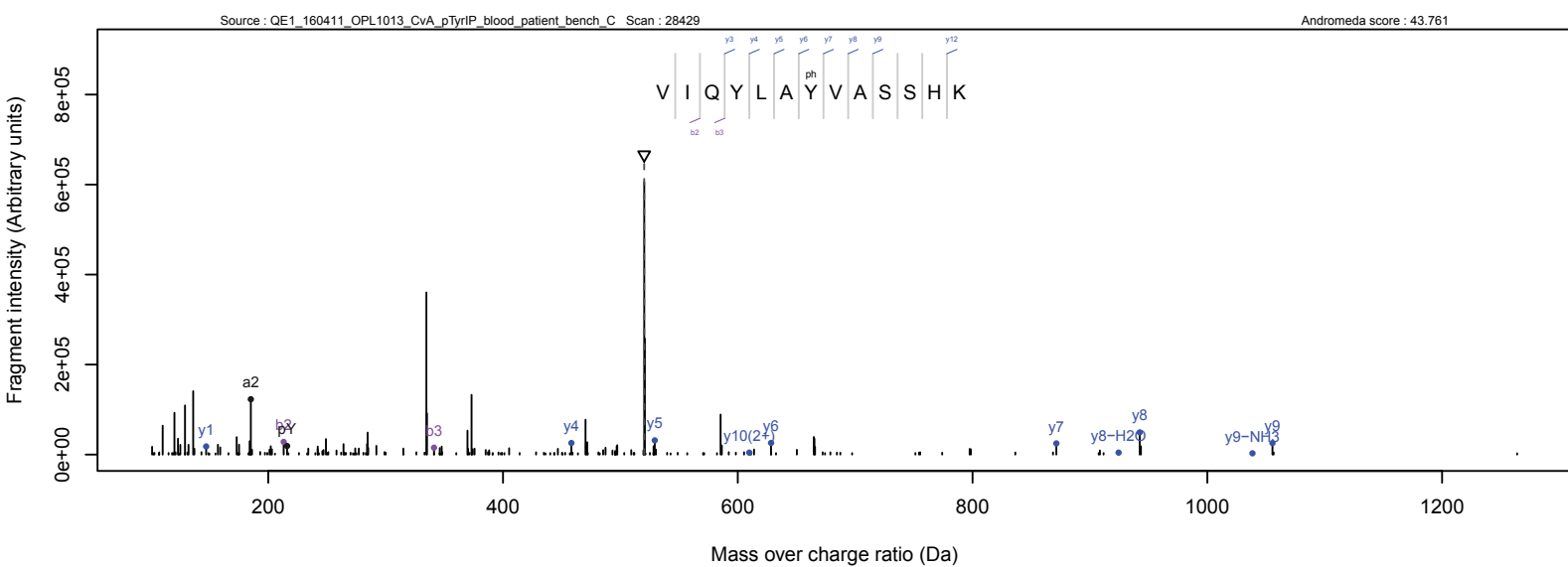
## LNERHY(ph)GGLTGLNK-3+



## LVNEAPVYSVYS(ph)K-2+



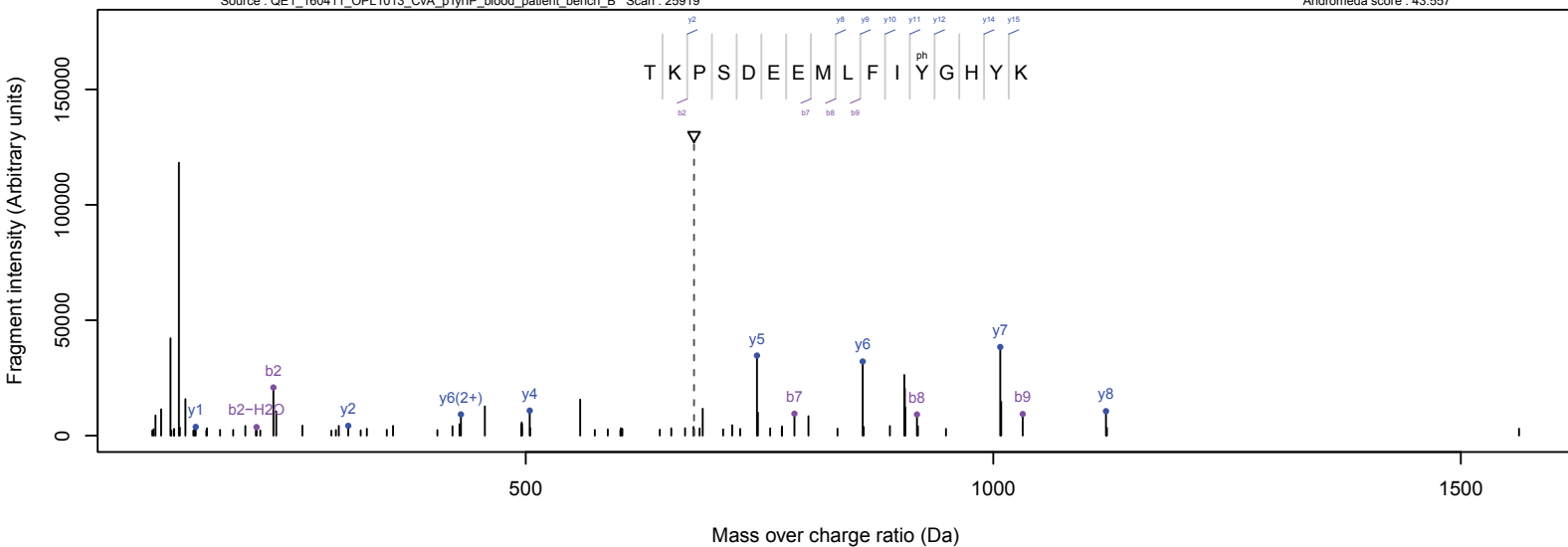
## VIQYLAYVASSHK-3+



**TKPSDEEMLFY(ph)GHYK-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan: 25919

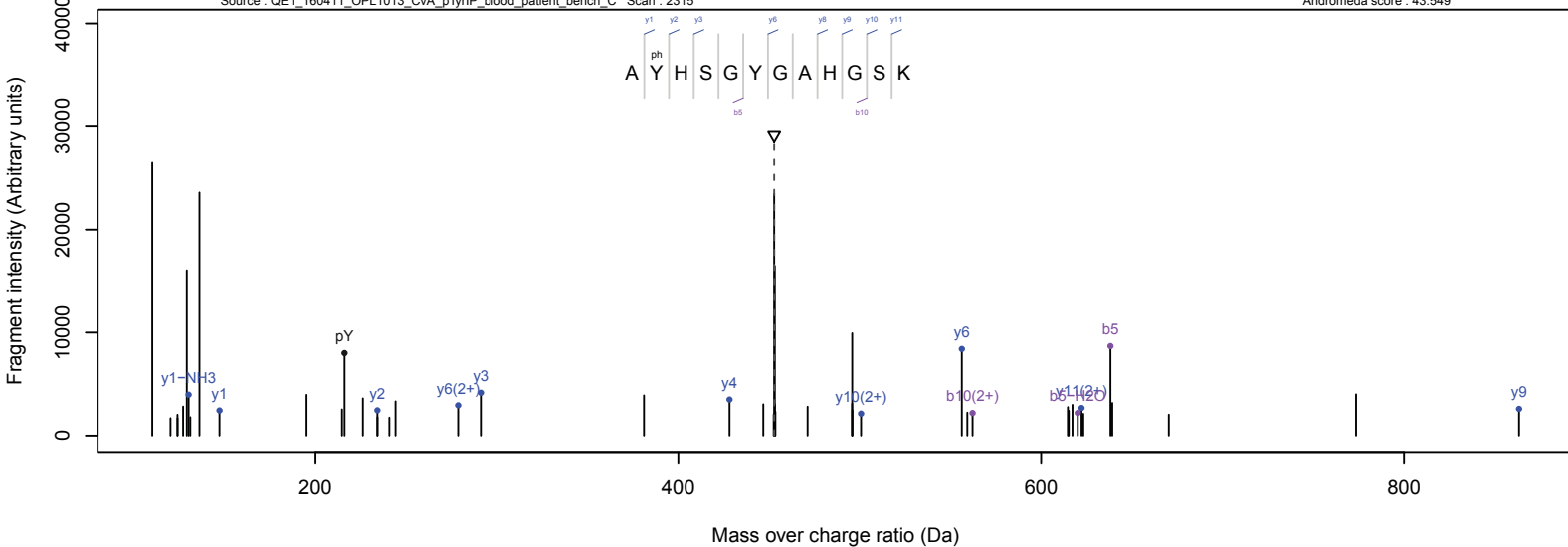
Andromeda score: 43.557



**(ac)AYHSGYGAHGSK-3+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 2315

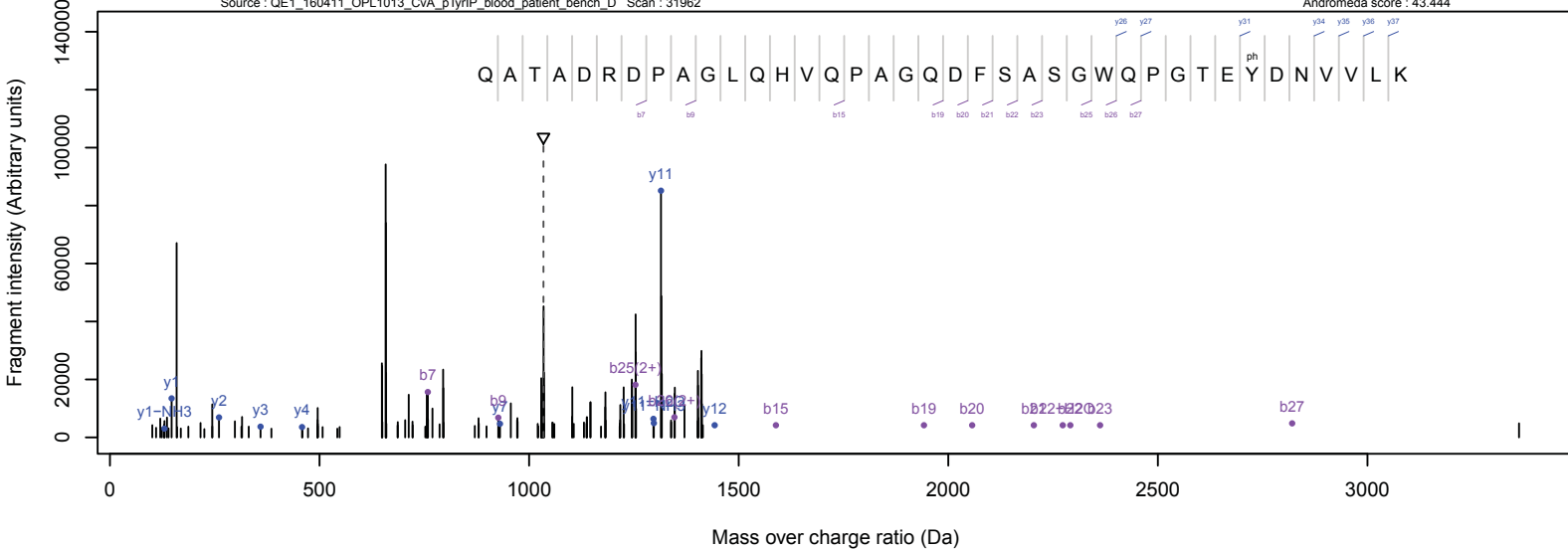
Andromeda score: 43.549



**QATADRDPAQLQHVQPAGQDFASGWPQTEY(ph)DNVVLK-4+**

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 31962

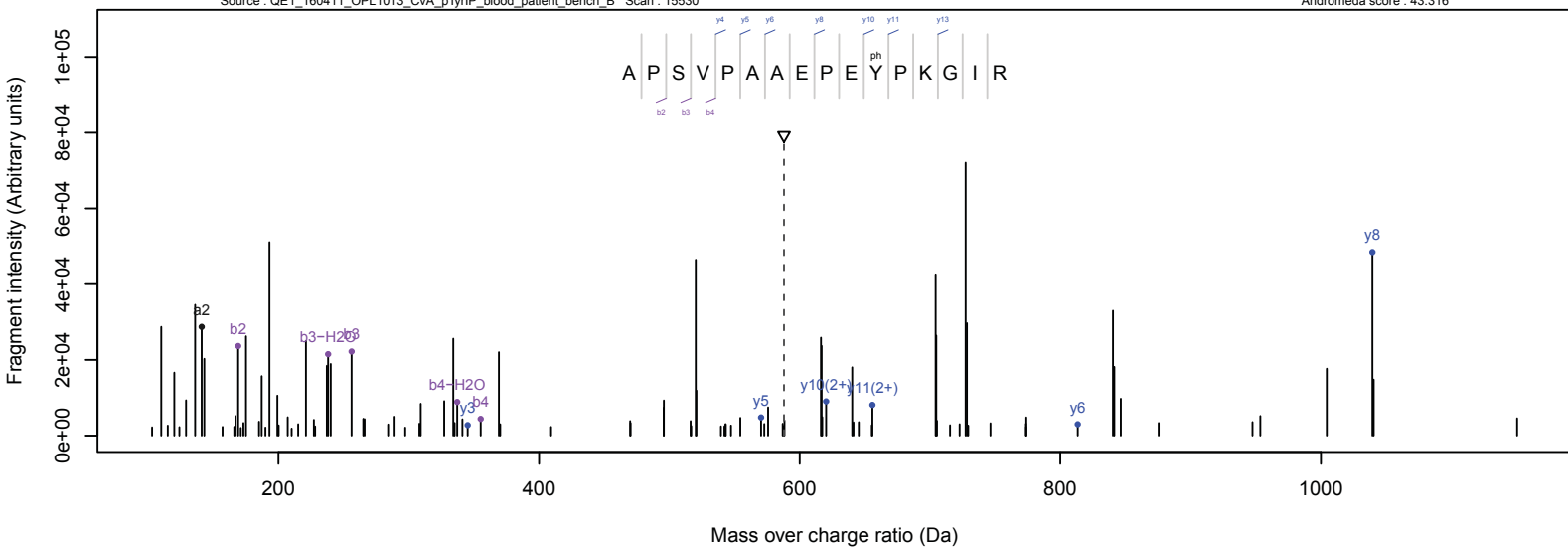
Andromeda score: 43.444



APSVPAEPEY(ph)PKGIR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 15530

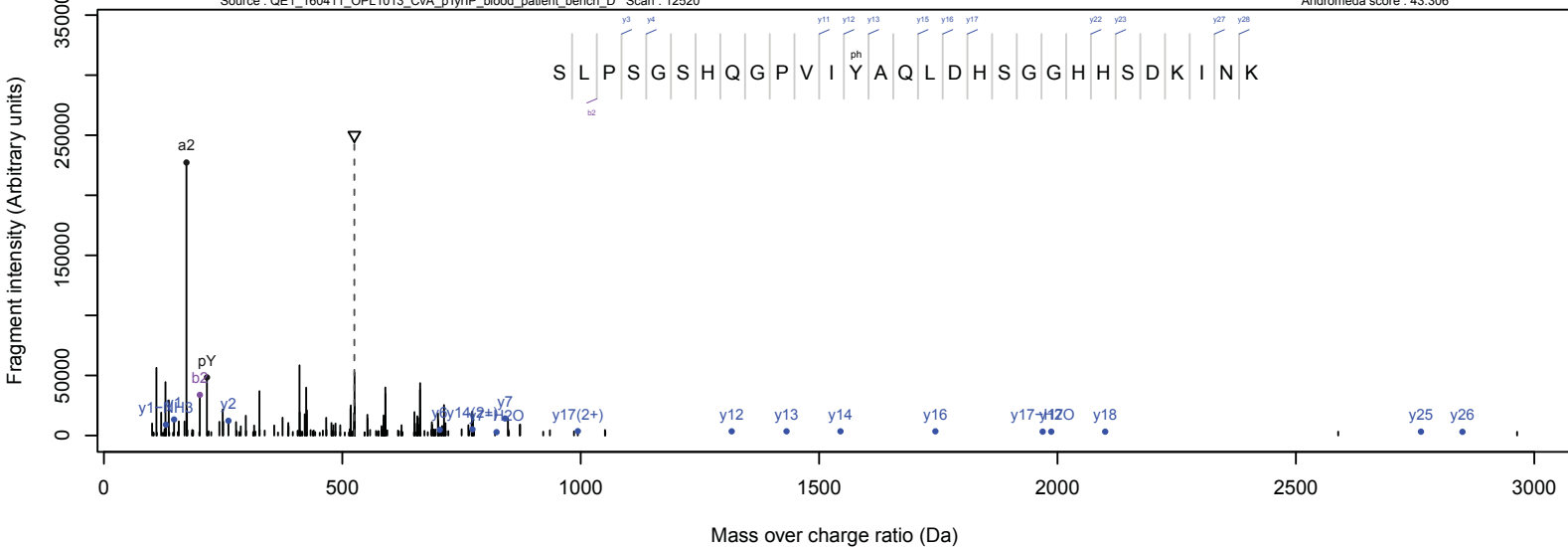
Andromeda score : 43.316



SLPSGSHQGPVIY(ph)AQLDHSGHHSDKINK-6+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 12520

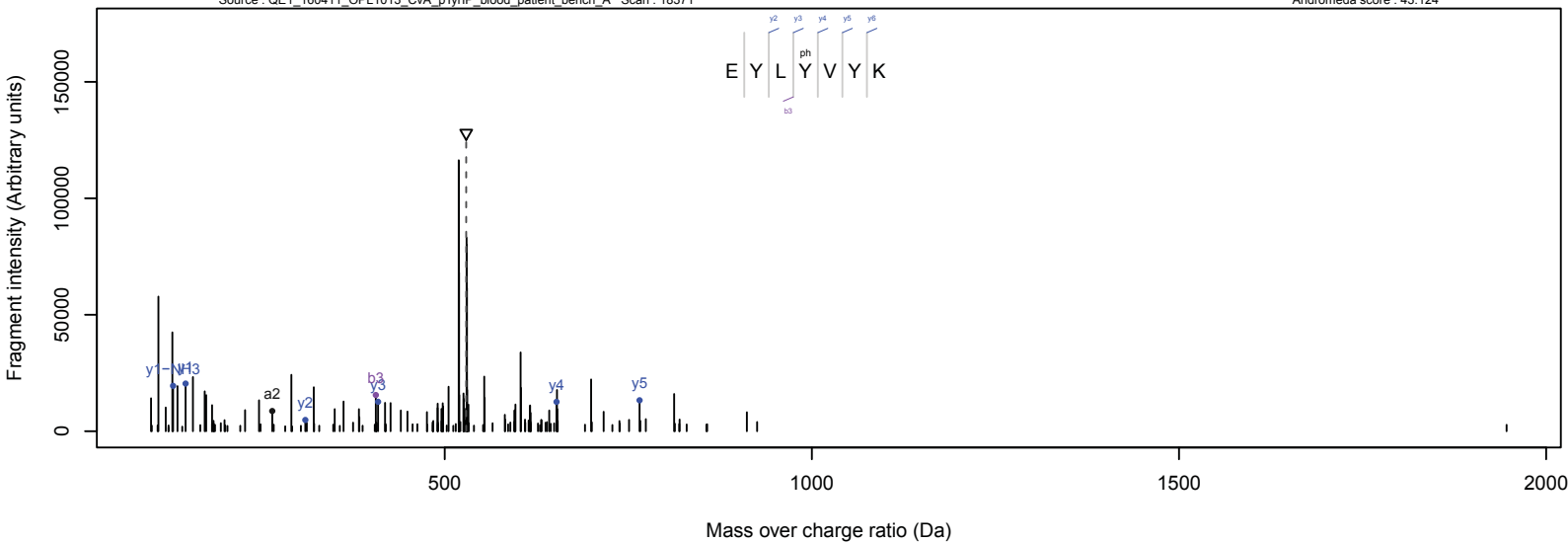
Andromeda score : 43.306



EYLY(ph)VYK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 18371

Andromeda score : 43.124

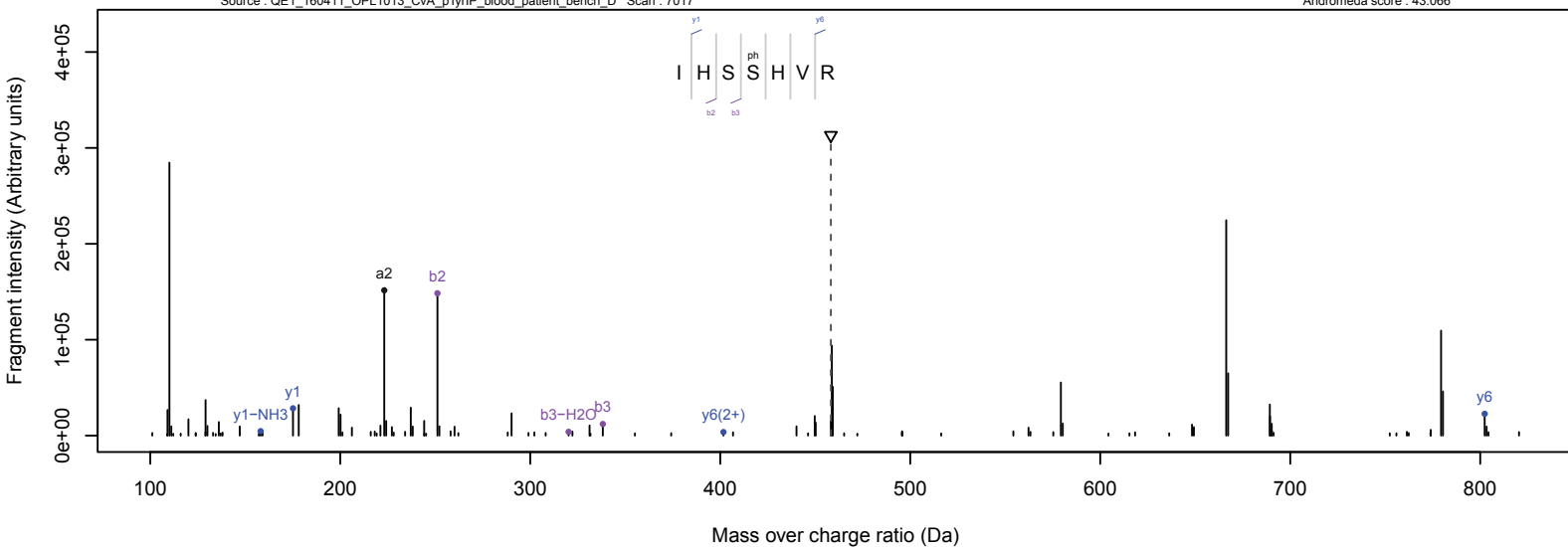




IHSS(ph)HVR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7017

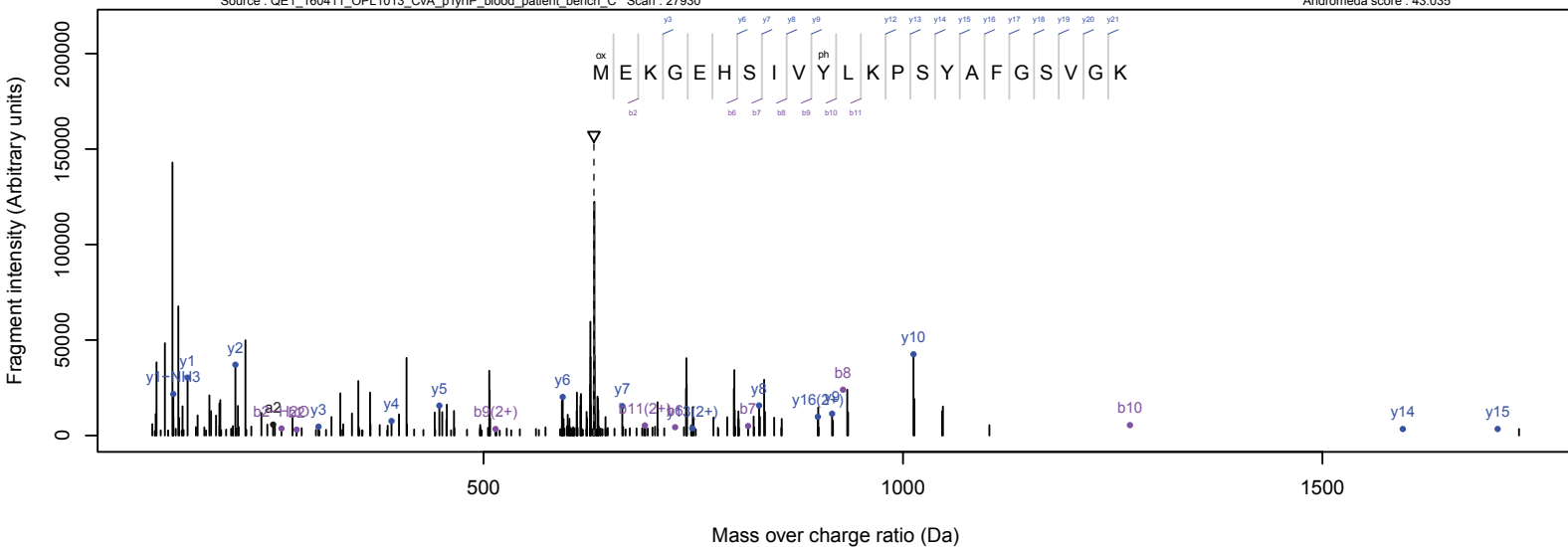
Andromeda score : 43.066



M(ox)EKGEHSIVY(ph)LKPSYAFGSV GK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 27930

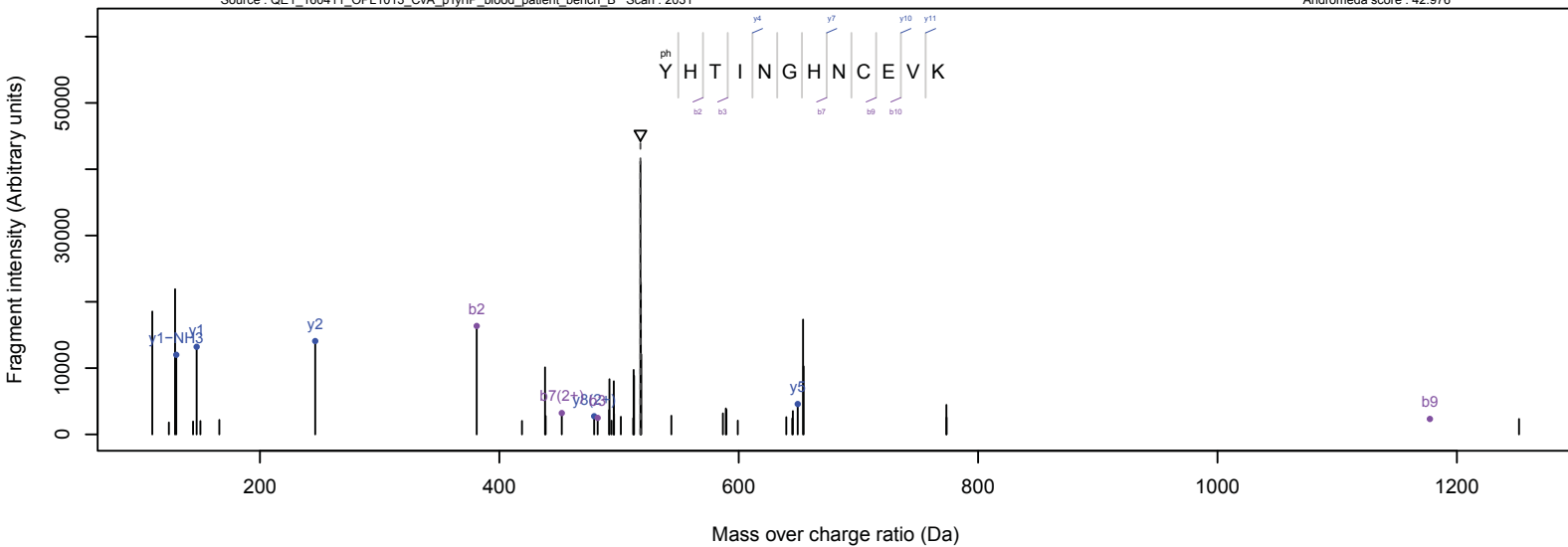
Andromeda score : 43.035



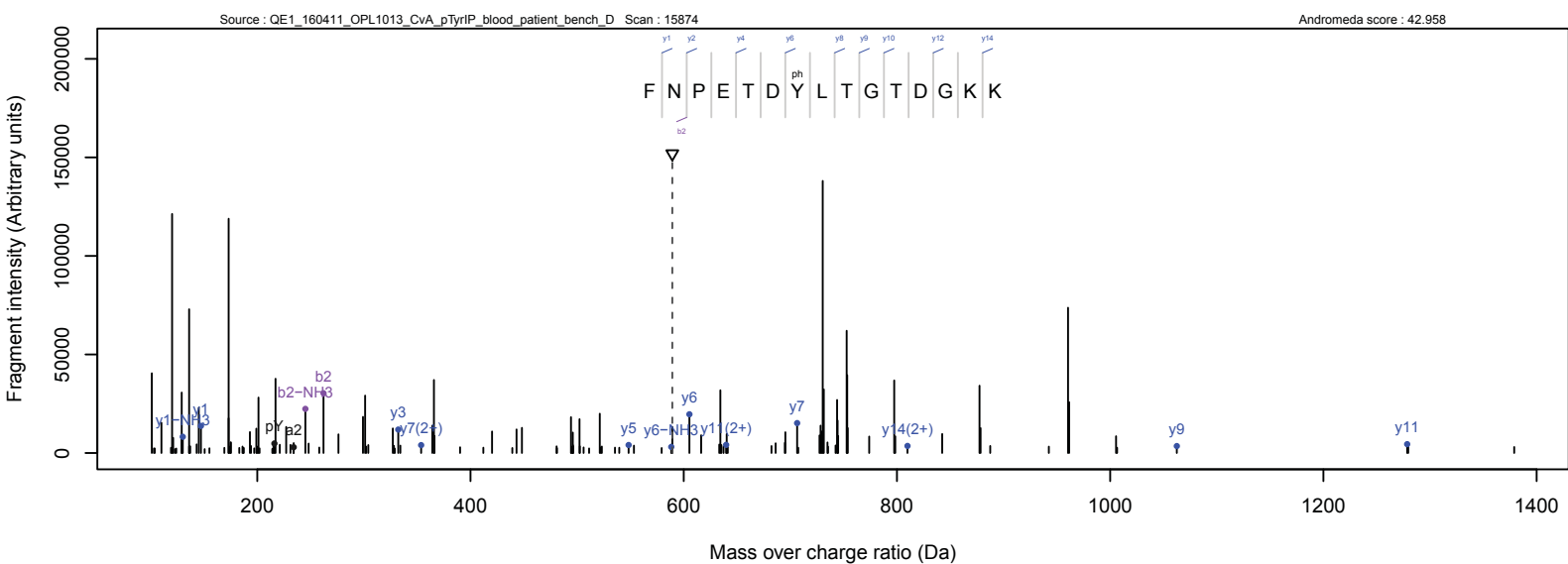
Y(ph)HTINGHNC EVK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 2031

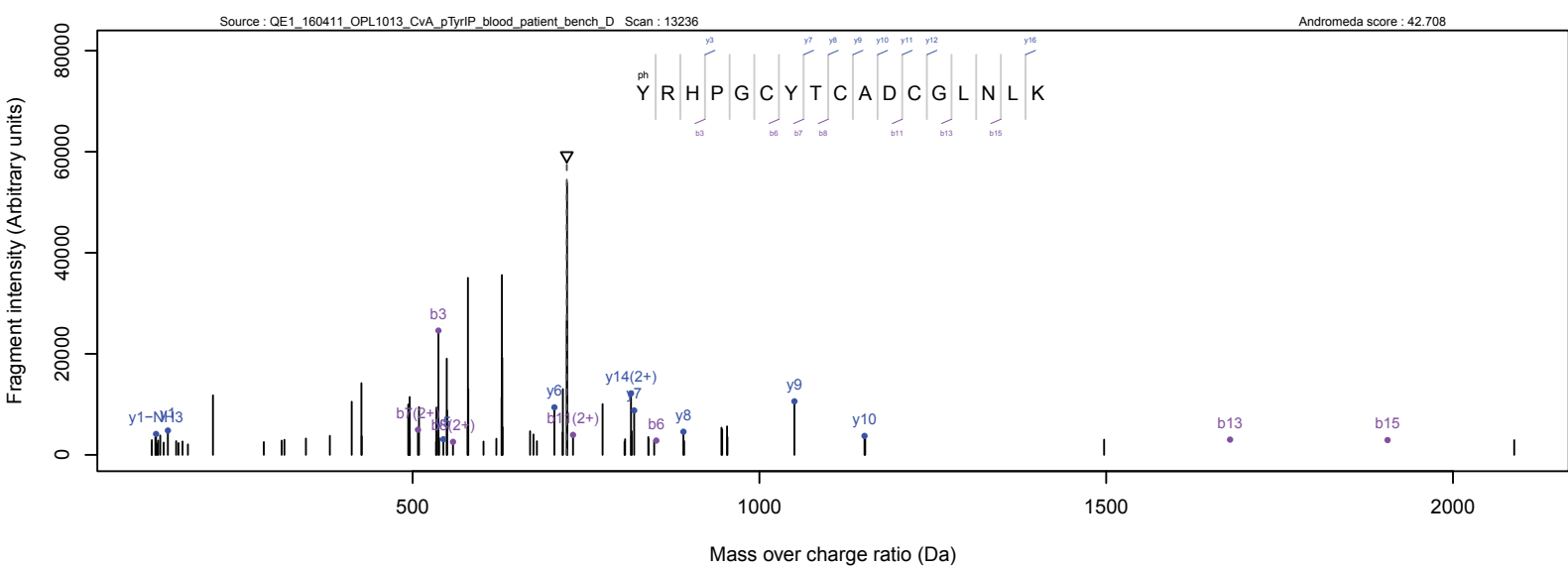
Andromeda score : 42.976



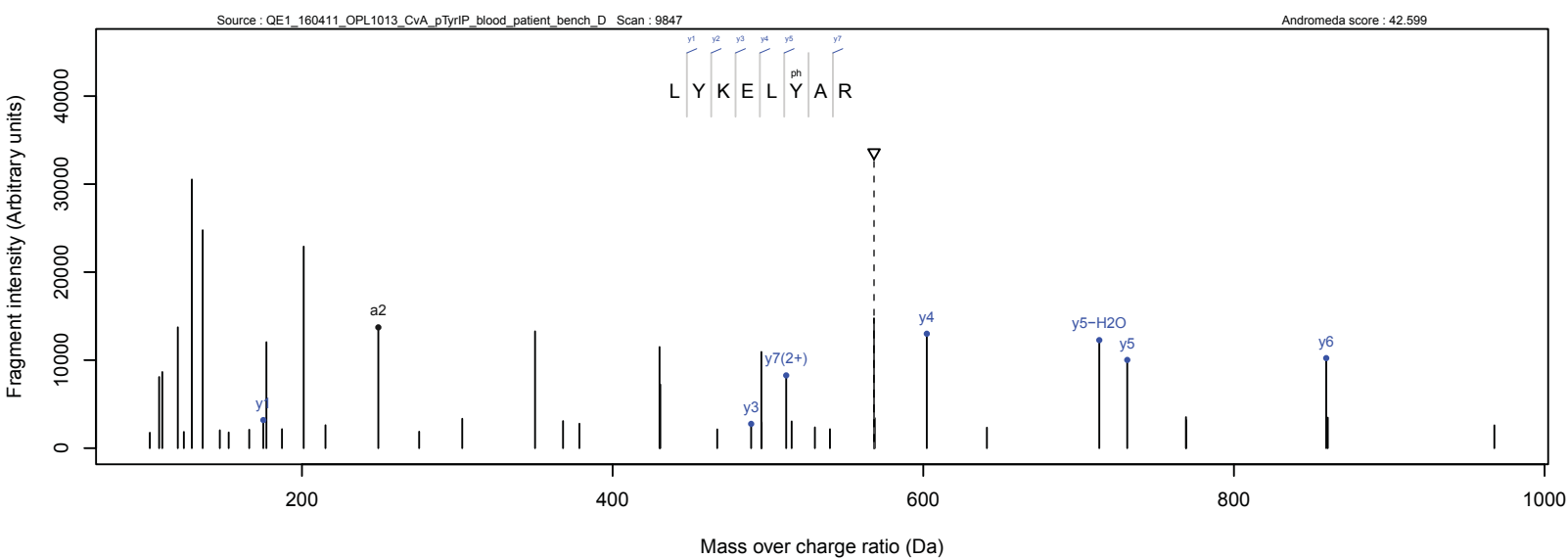
## FNPETDY(ph)LTGTDGKK-3+



## Y(ph)RHPGCYTCADCGLNLLK-3+



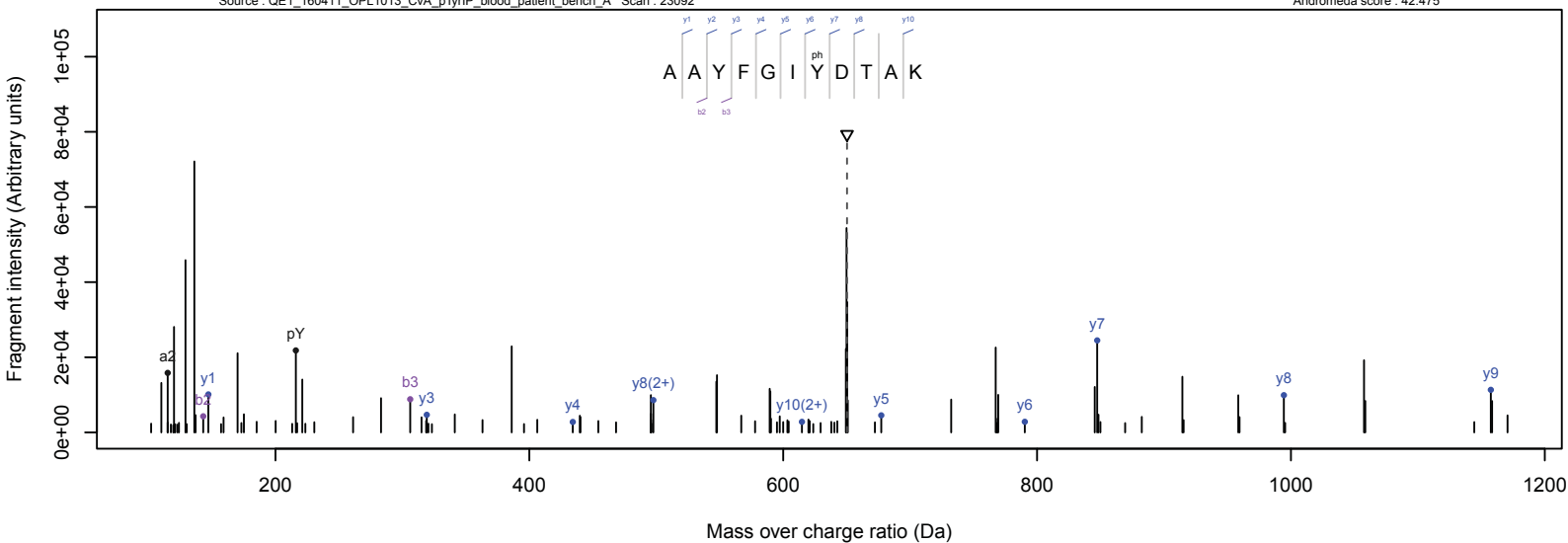
## LYKELY(ph)AR-2+



## AAFYGIY(ph)DTAK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 23092

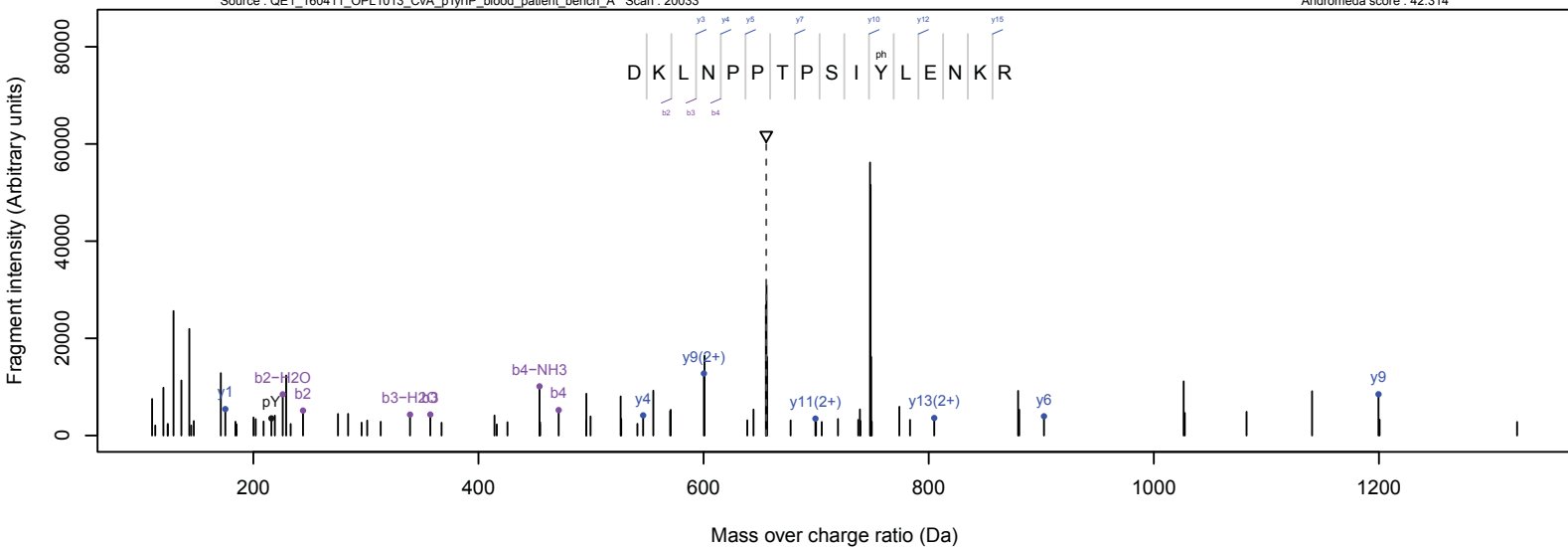
Andromeda score : 42.475



## DKLNPTPSIY(ph)LENKR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 20033

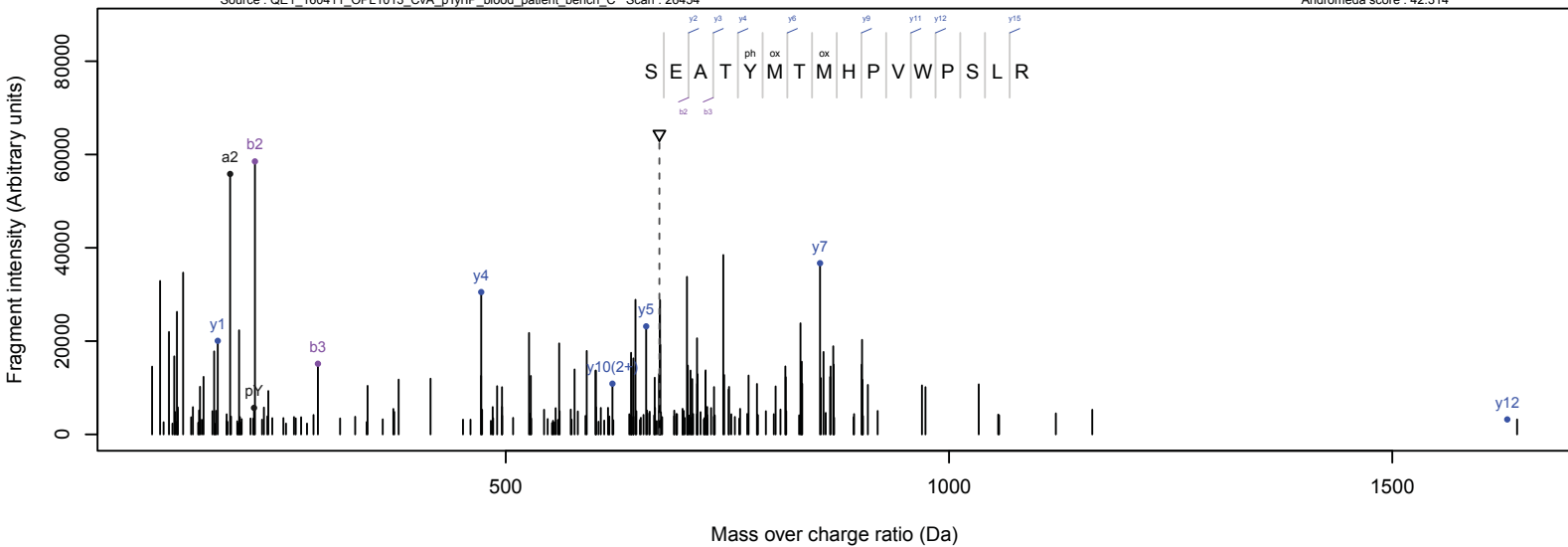
Andromeda score : 42.314



## SEATY(ph)M(ox)TM(ox)HPVWPSLR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 26454

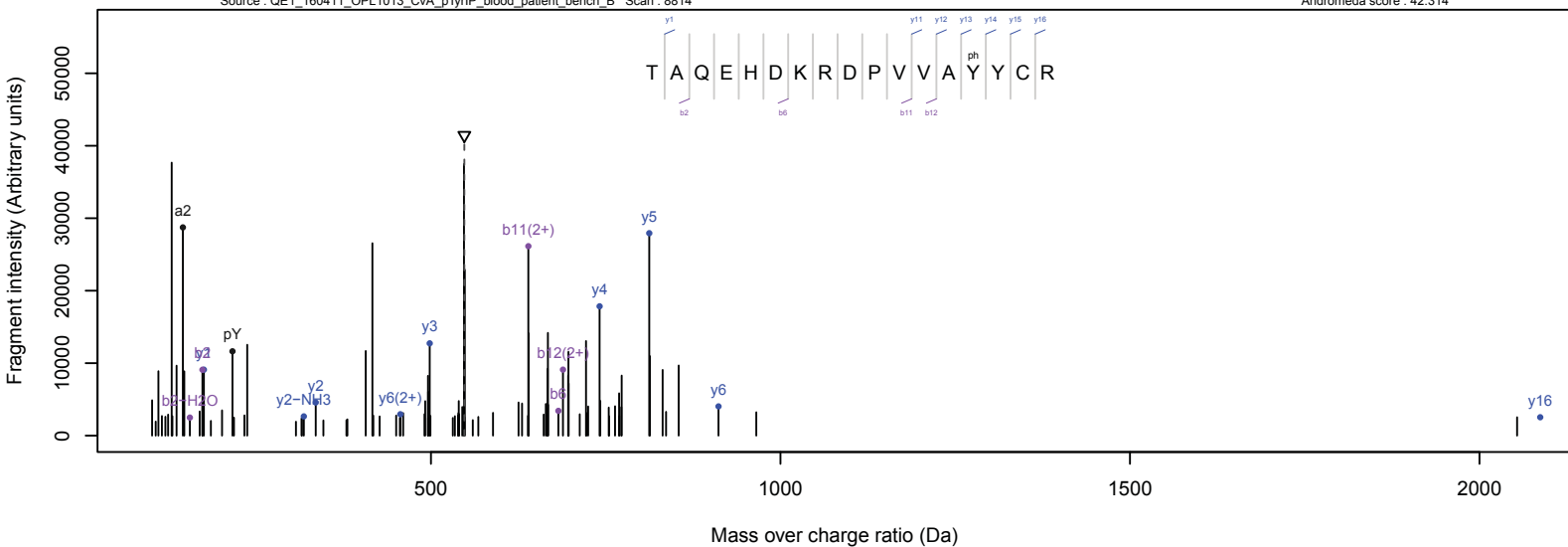
Andromeda score : 42.314



**TAQEHDKRDVPVAVYCR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 8814

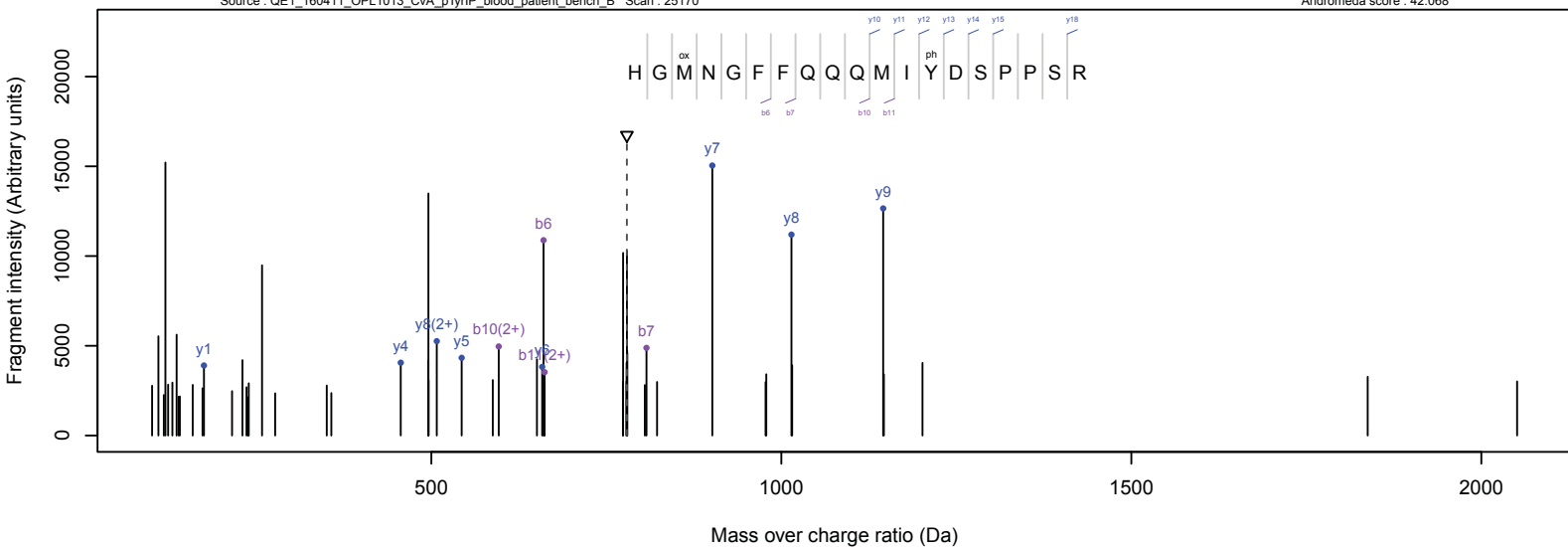
Andromeda score : 42.314



**HGM(ox)NGFFQQMIYDSSPPSR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 25170

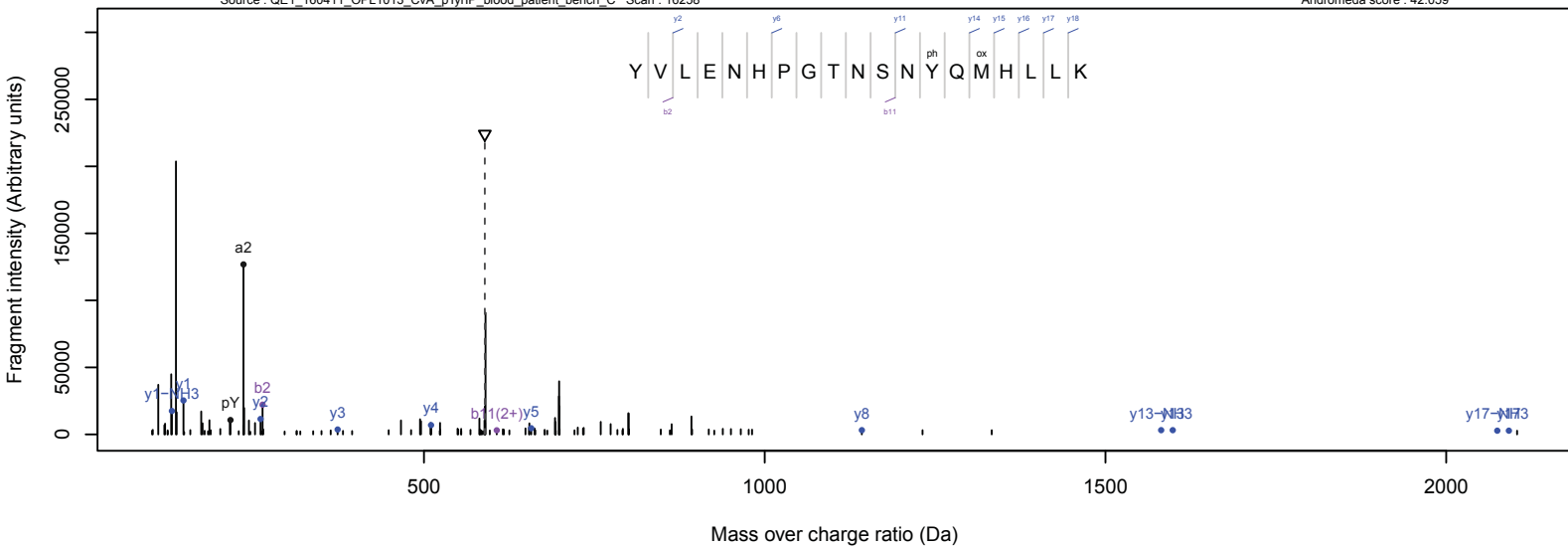
Andromeda score : 42.068



**YVLENHPGTNSNYQM(ox)HLLK-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 16258

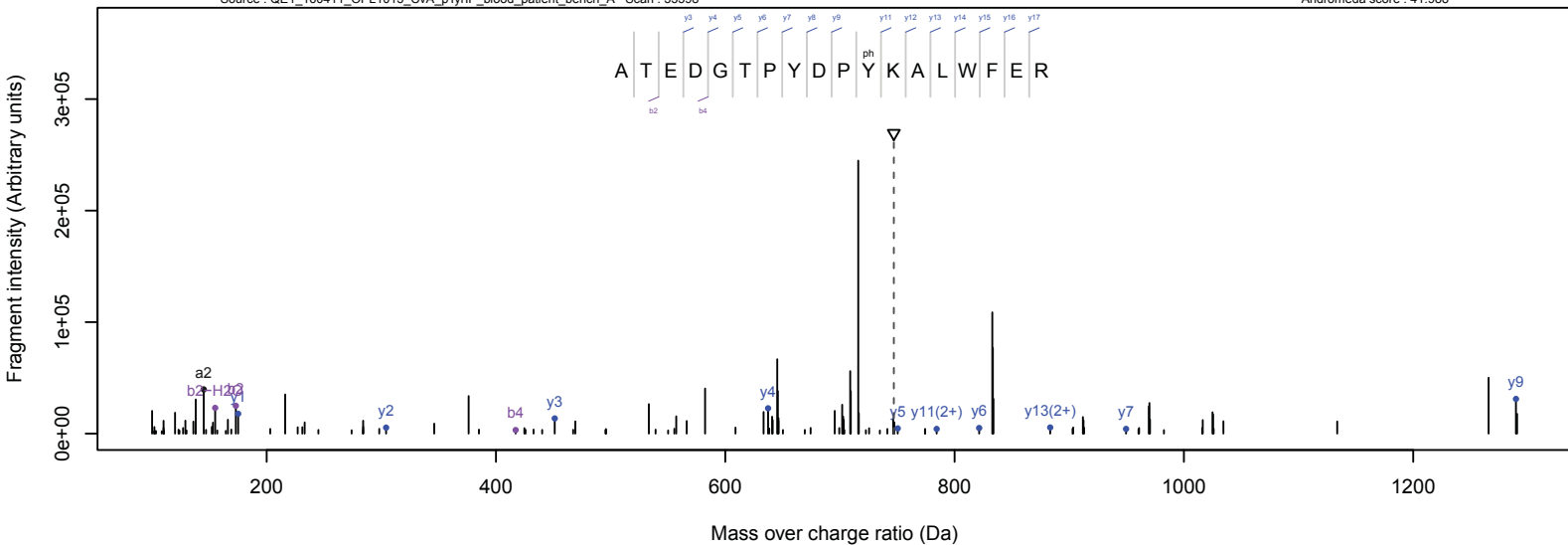
Andromeda score : 42.059



ATEDGTPYDPY(ph)KALWFER-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 33398

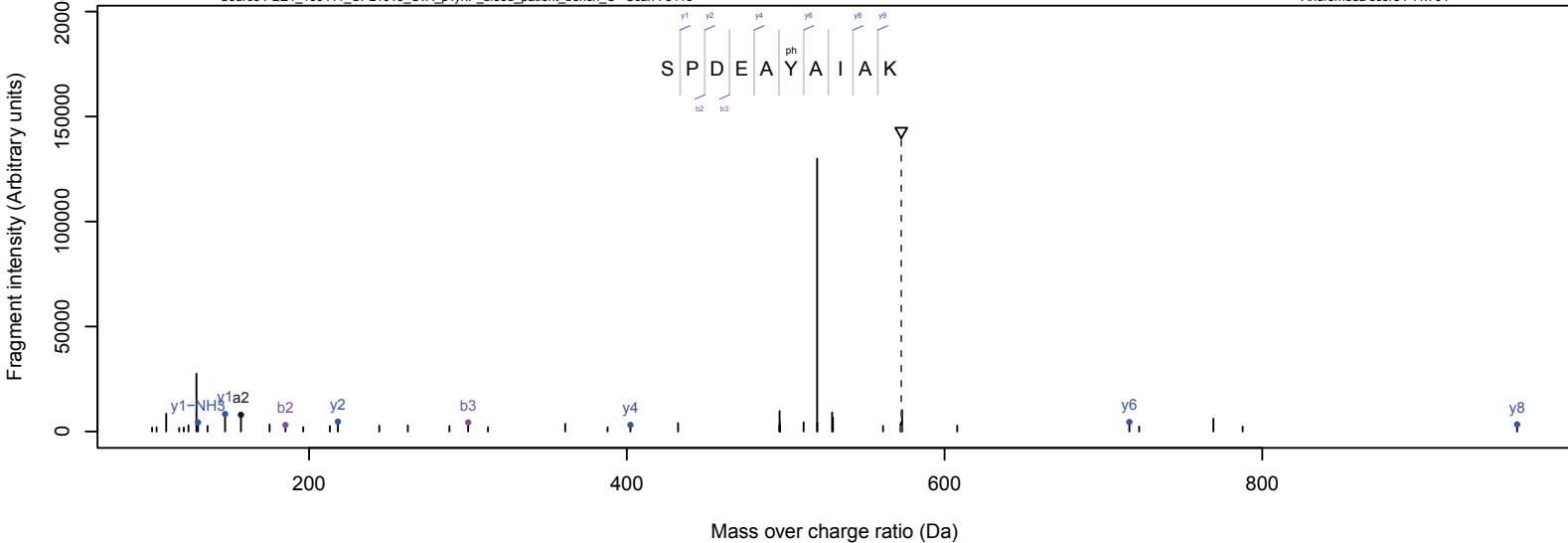
Andromeda score : 41.988



SPDEAY(ph)AIK-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 8418

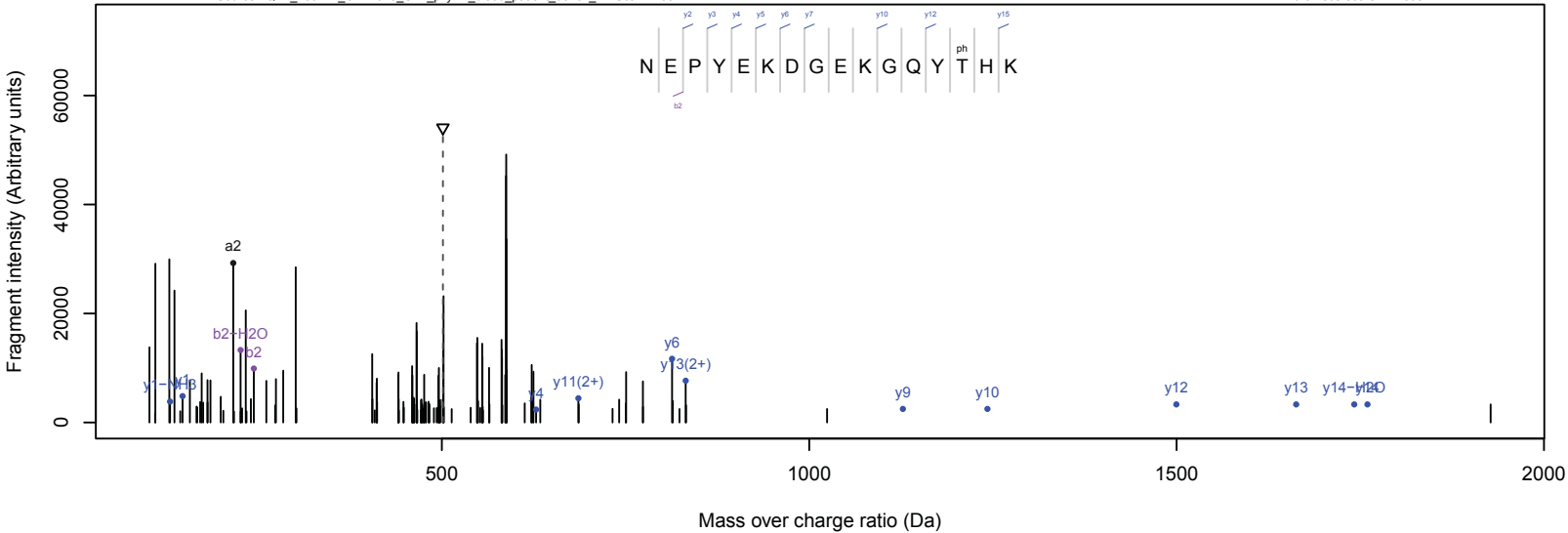
Andromeda score : 41.704



NEPYEKDGEKQY(ph)HK-4+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 1667

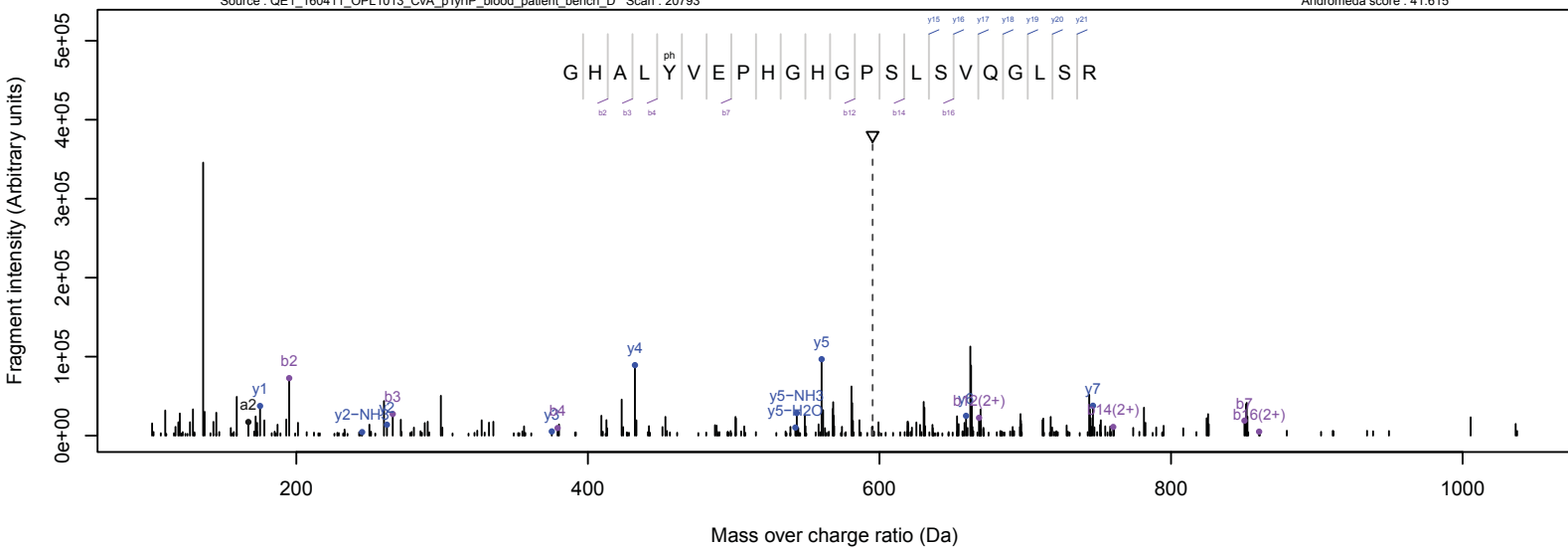
Andromeda score : 41.695



**GHALY(ph)VEPHGHGPSLSVQGLSR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 20793

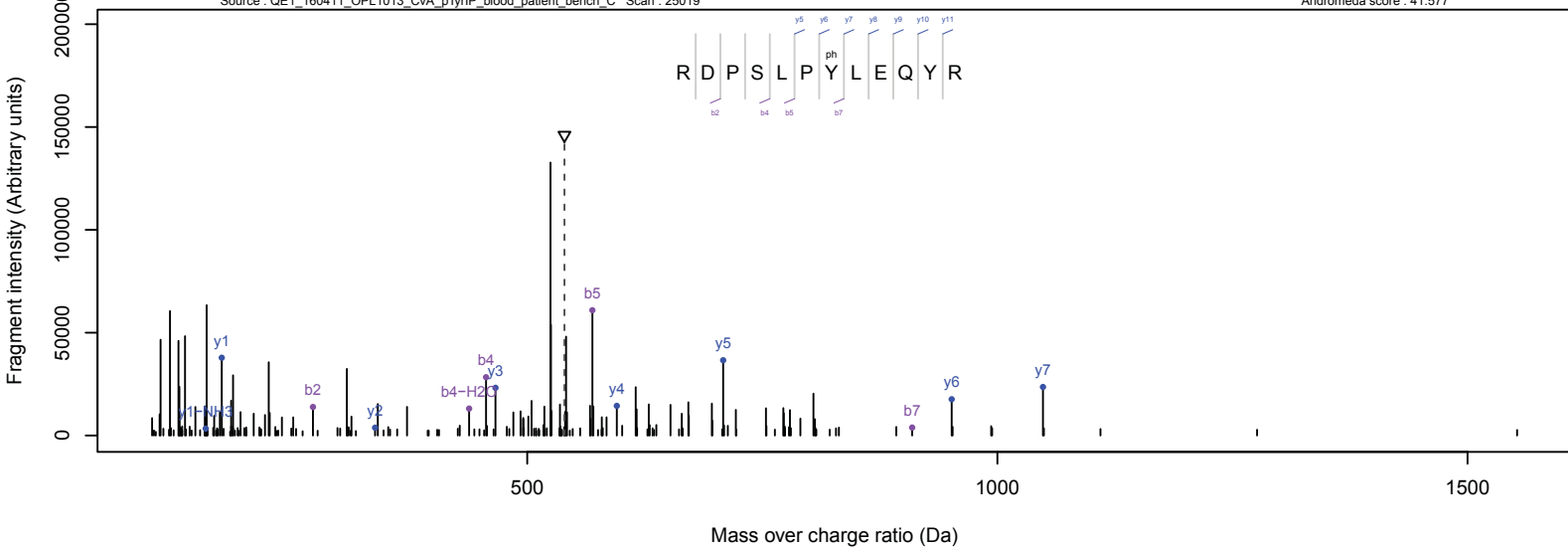
Andromeda score : 41.615



**RDPSLPY(ph)LEQYR-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 25019

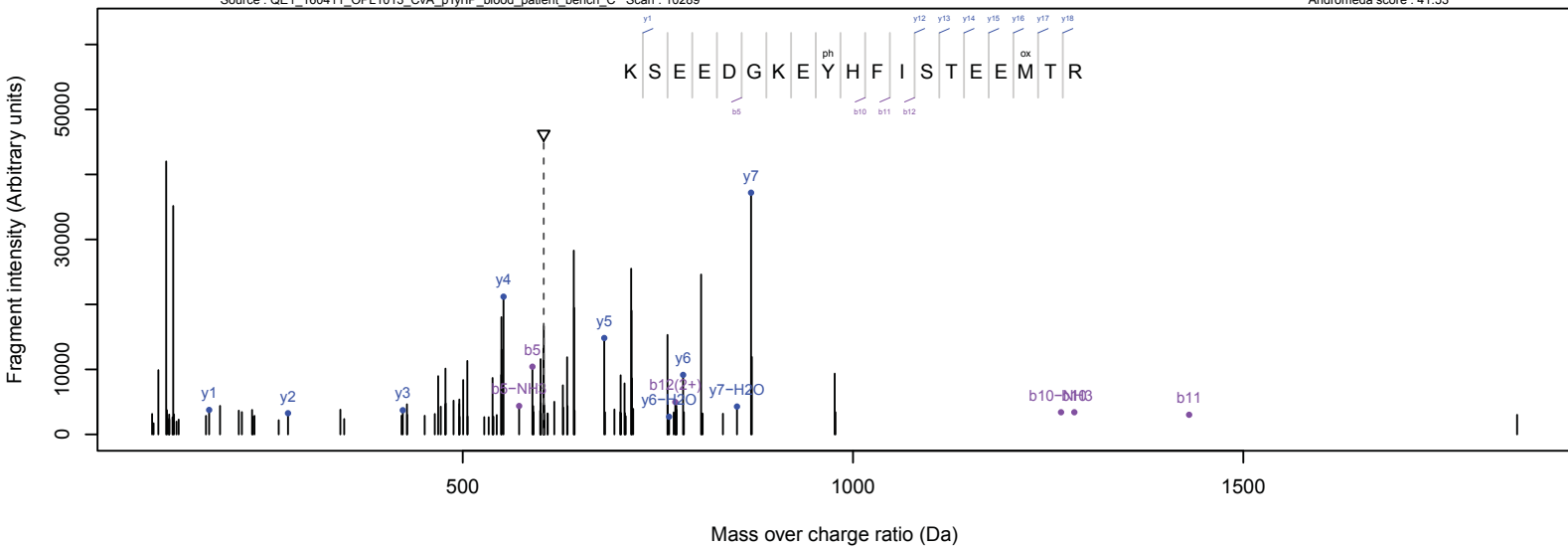
Andromeda score : 41.577



**KSEEDGKEY(ph)HFISTEEM(ox)TR-4+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 10289

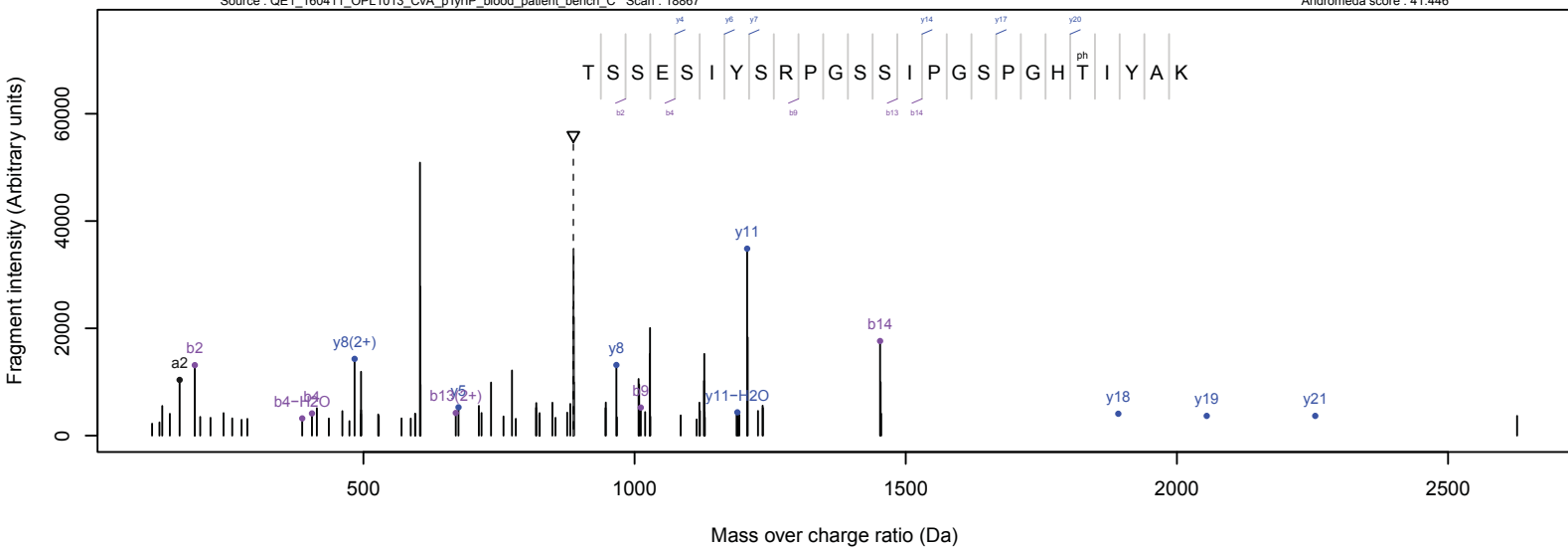
Andromeda score : 41.53



**TSESIYRPGSSIPGSPGHT(ph)IYAK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 18867

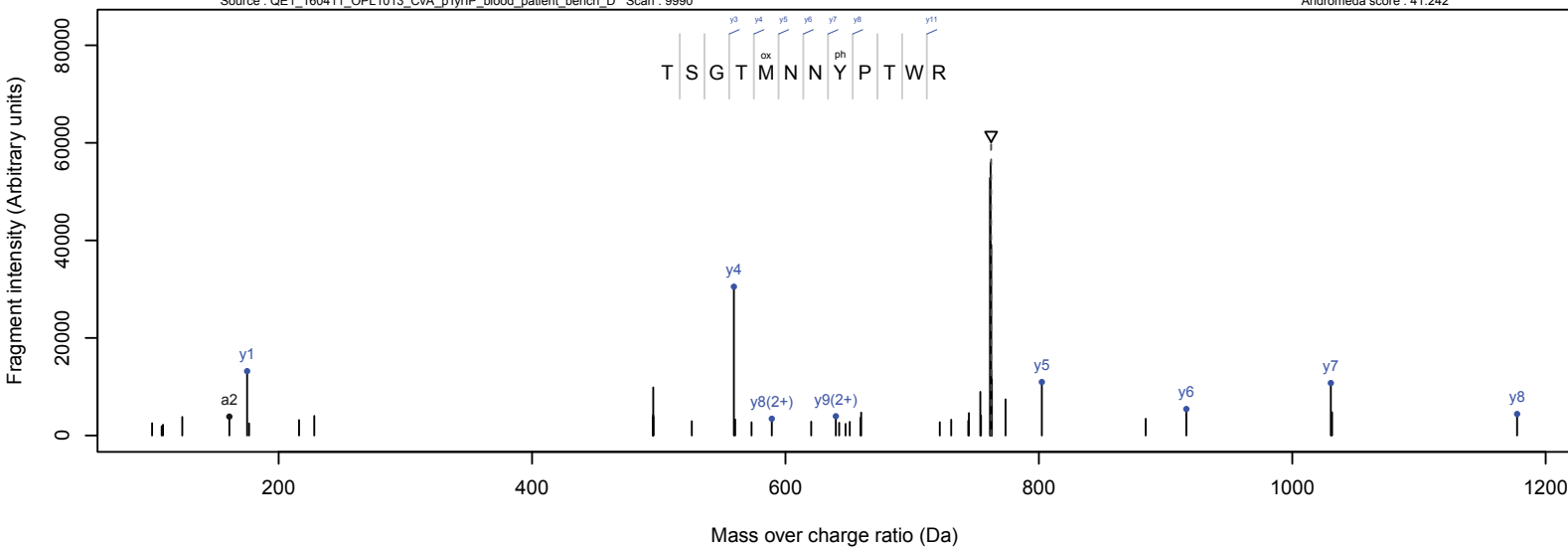
Andromeda score : 41.446



**TSGTM(ox)NNY(ph)PTWR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 9990

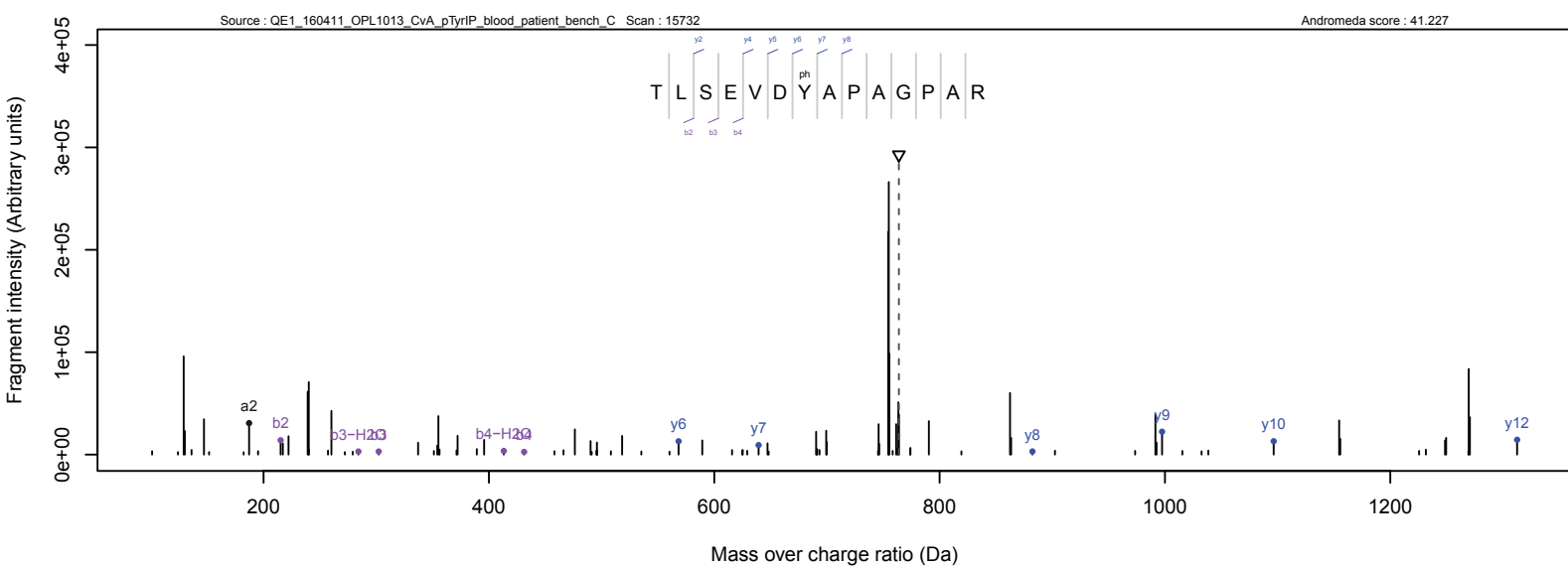
Andromeda score : 41.242



**TLSEVDY(ph)APAGPAR-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 15732

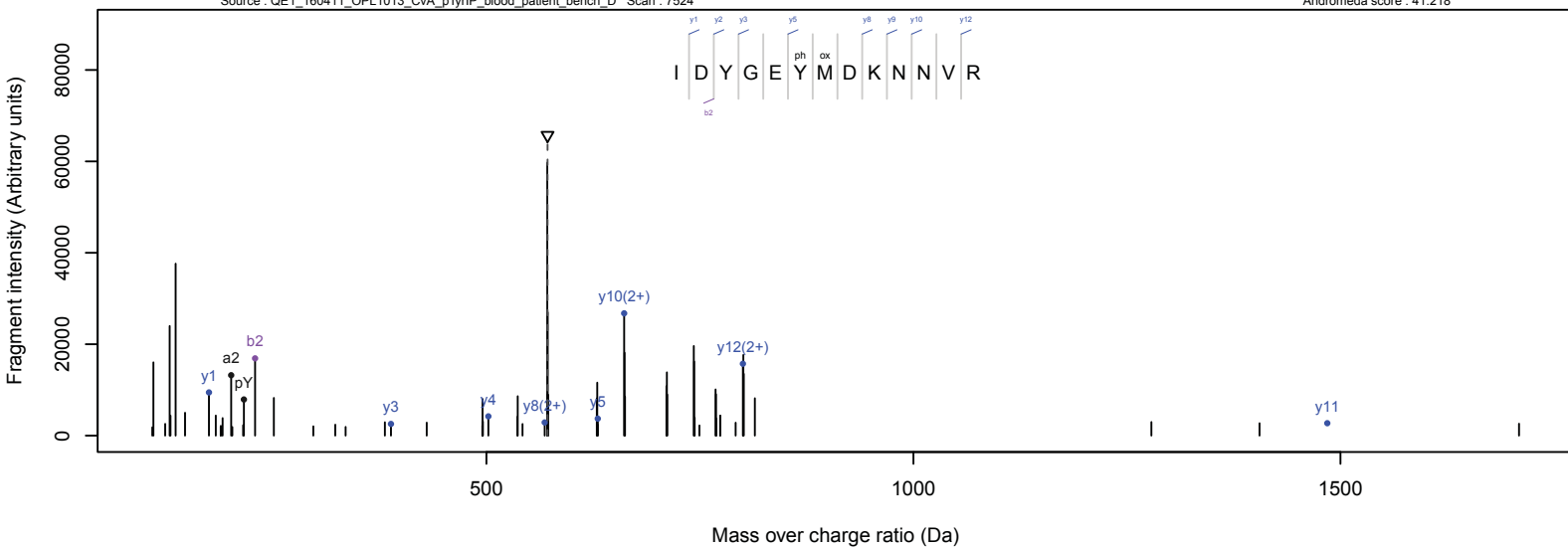
Andromeda score : 41.227



## IDYGEY(ph)M(ox)DKNNVR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 7524

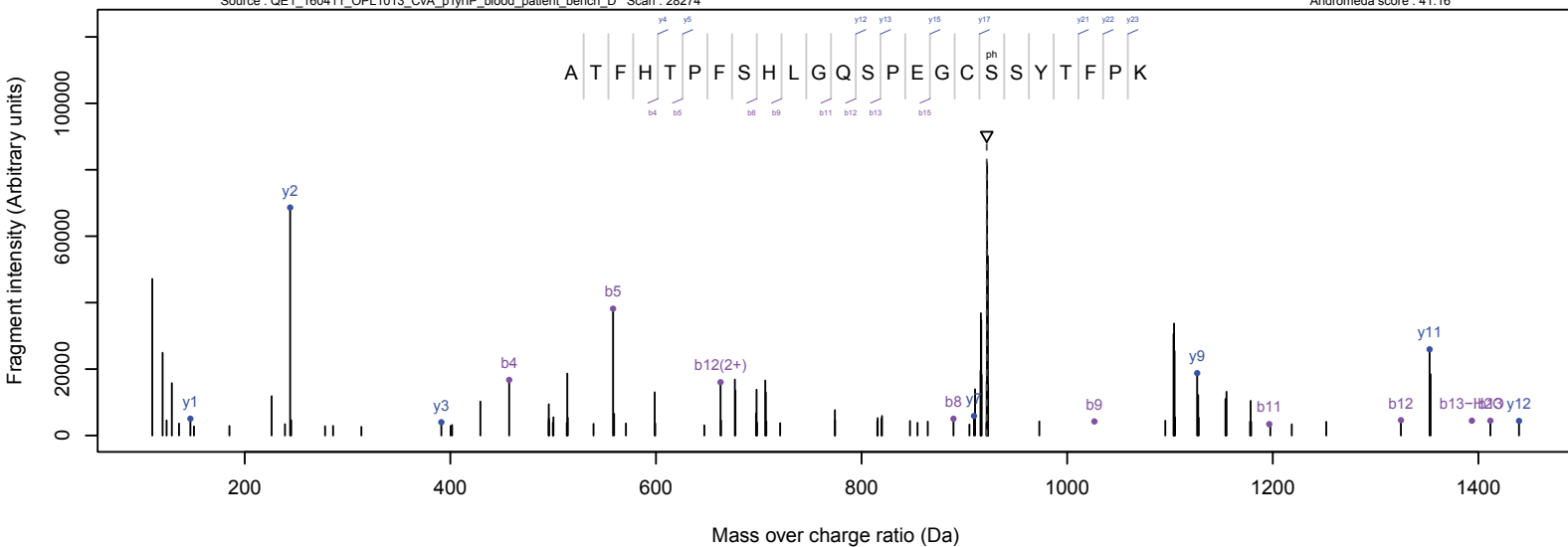
Andromeda score : 41.218



## ATFHTPFSLGQSPGCS(ph)SYTFPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28274

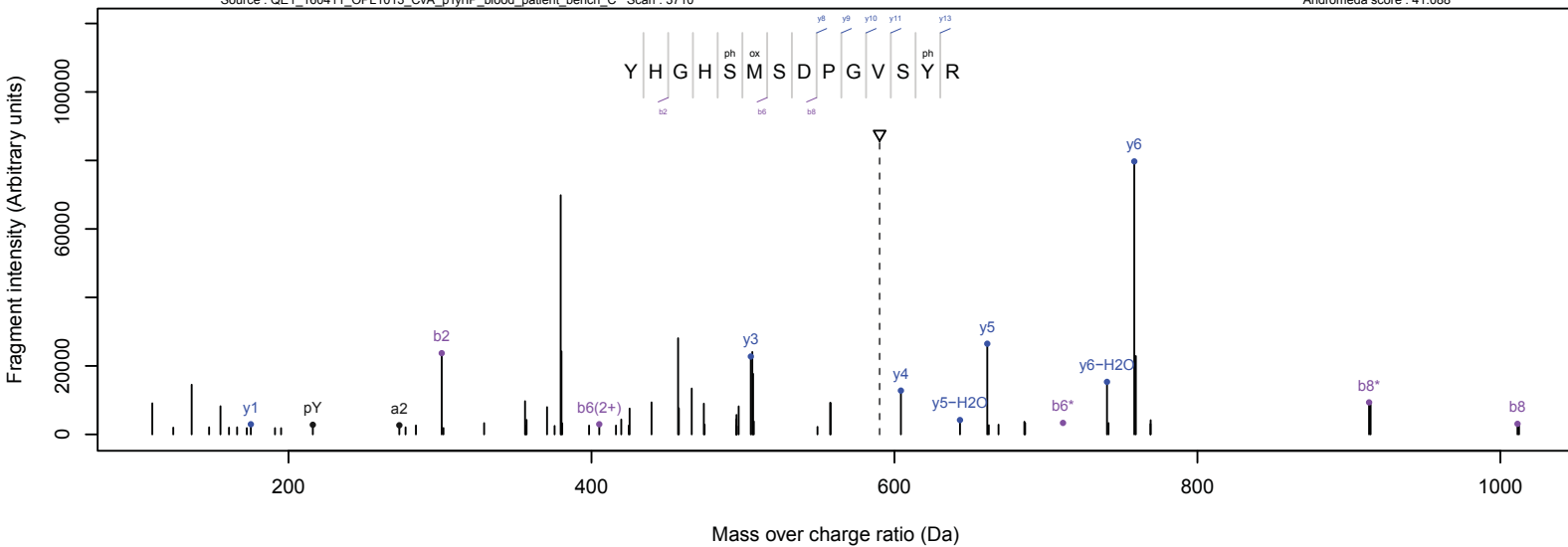
Andromeda score : 41.16



## YHGHS(ph)M(ox)SDPGVSY(ph)R-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 3710

Andromeda score : 41.088

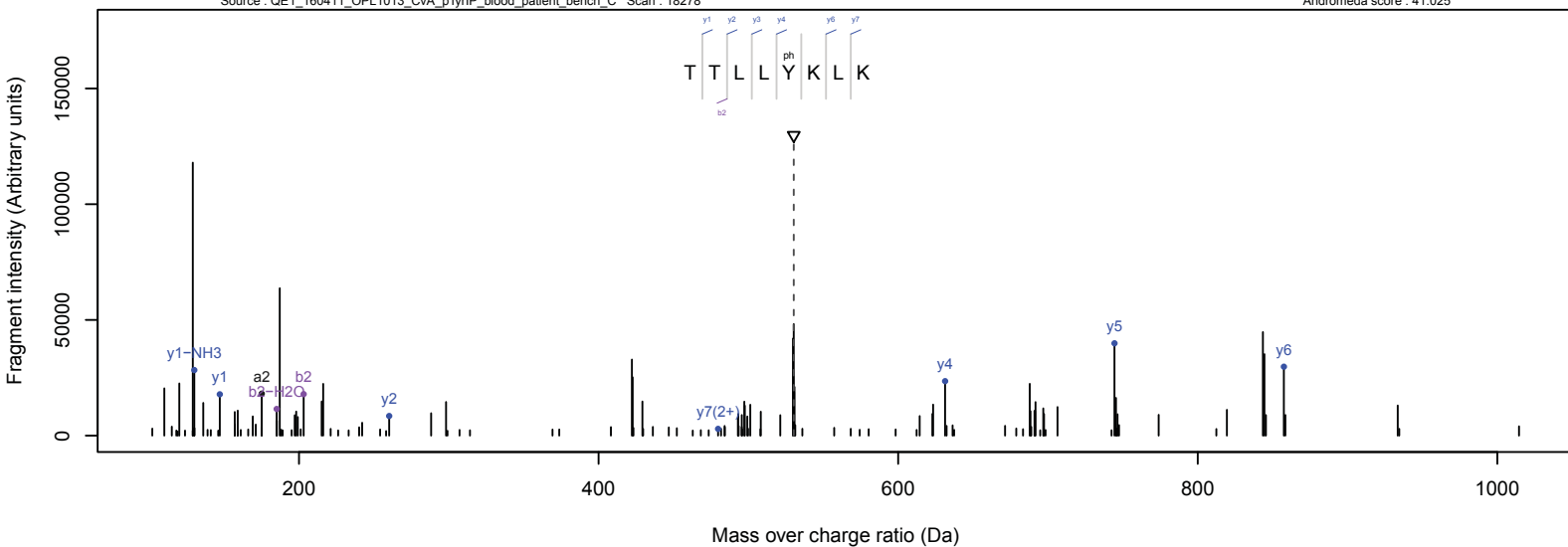




## TTLLY(ph)KLK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 18278

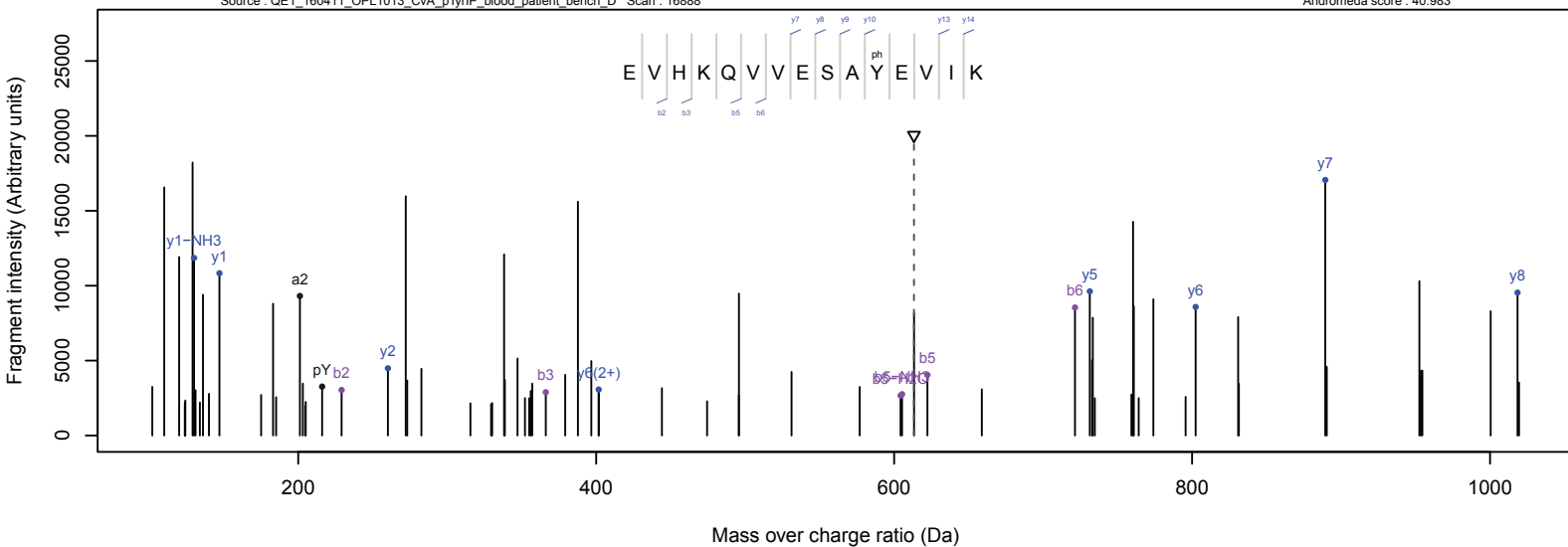
Andromeda score: 41.025



## EVHKQVSESAY(ph)EVIK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 16888

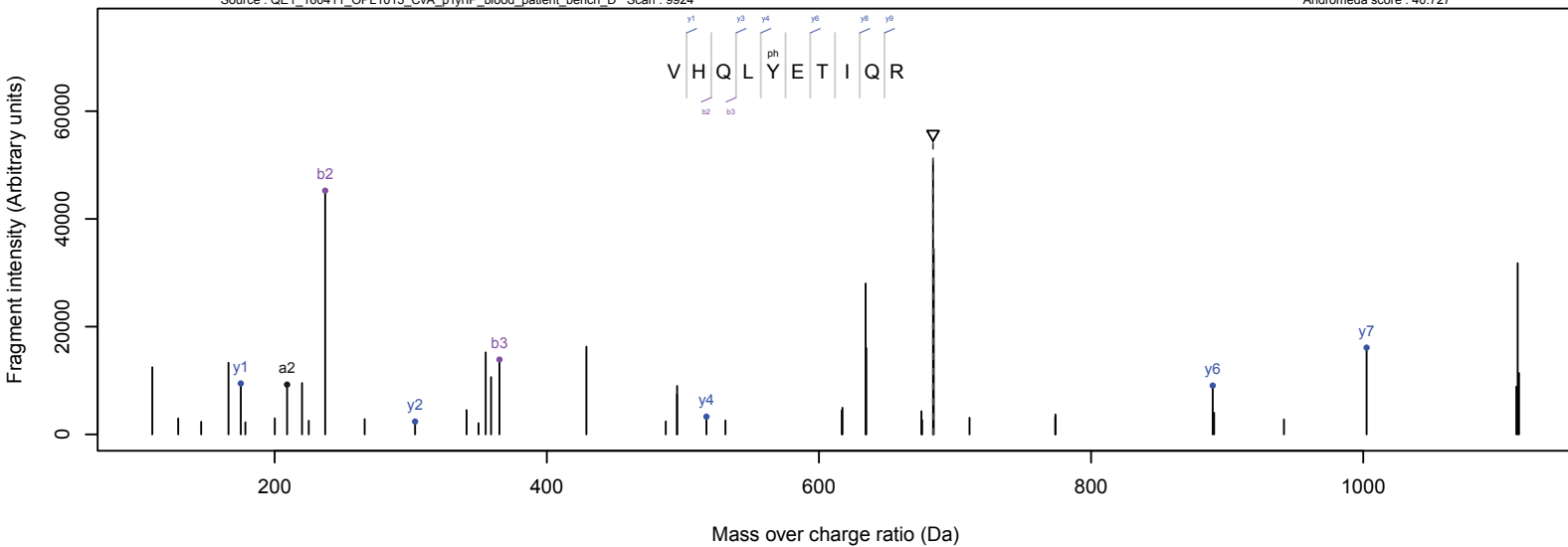
Andromeda score: 40.983



## VHQLY(ph)ETIQR-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 9924

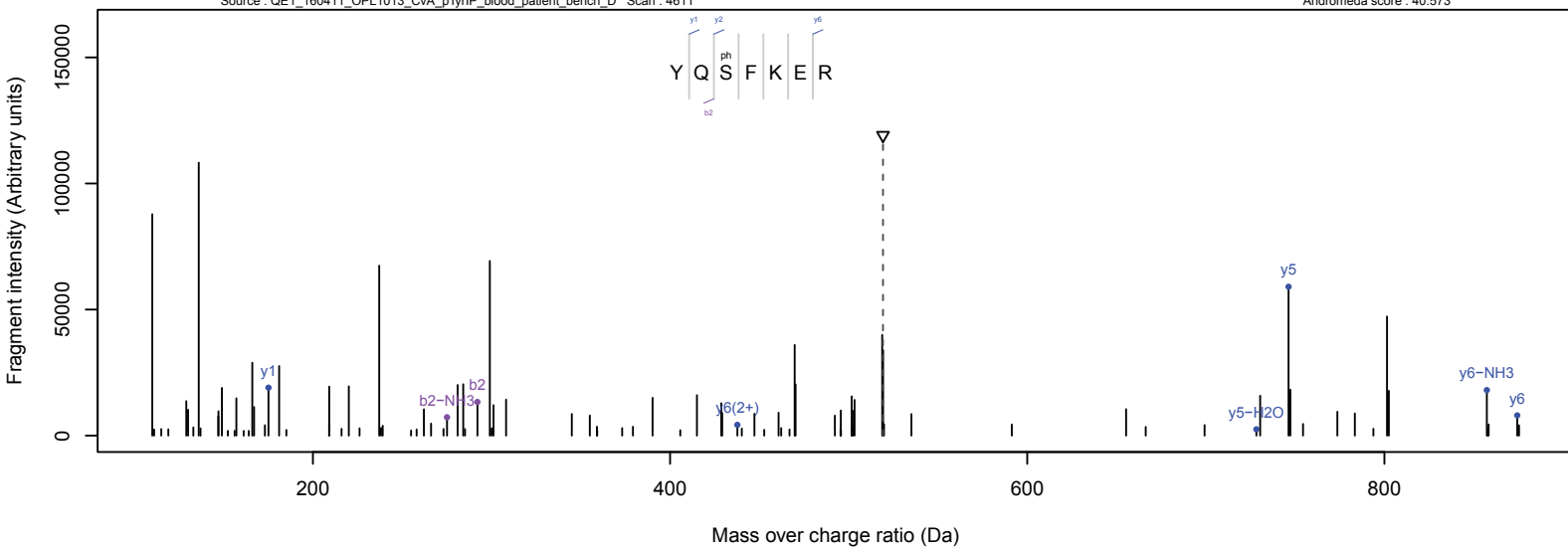
Andromeda score: 40.727



YQS(ph)FKER-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 4611

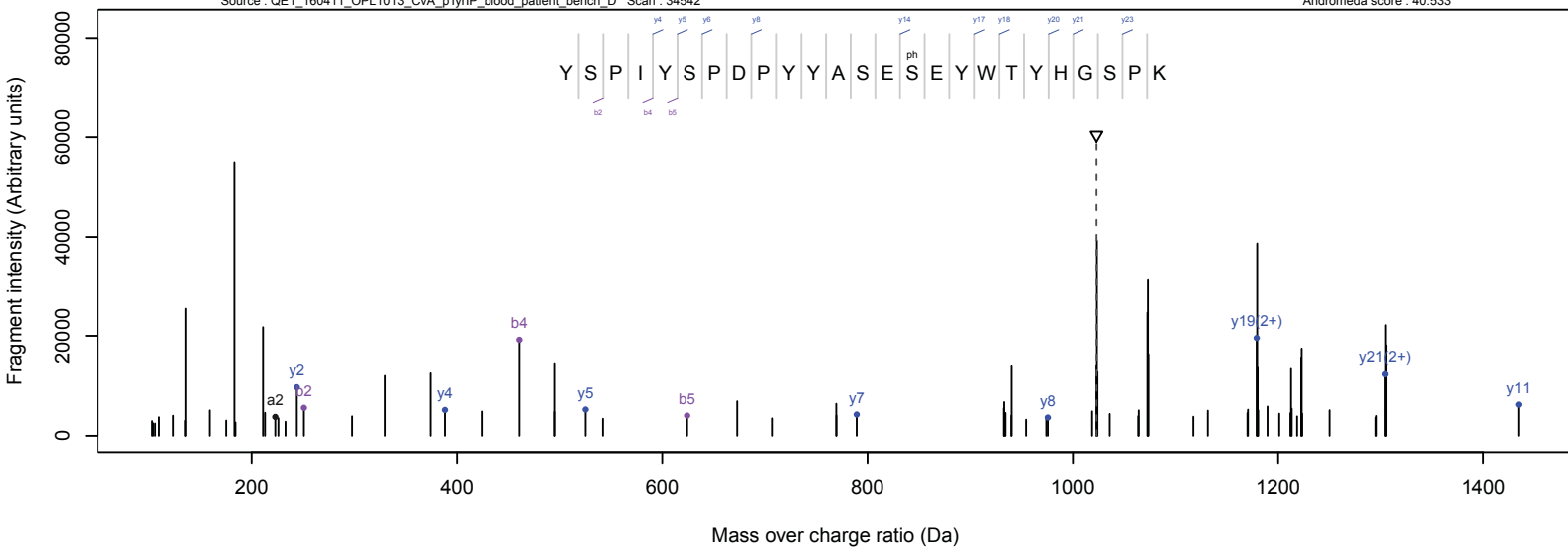
Andromeda score : 40.573



YSPIYSPDPYYASES(ph)EYWTYHGSPK-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 34542

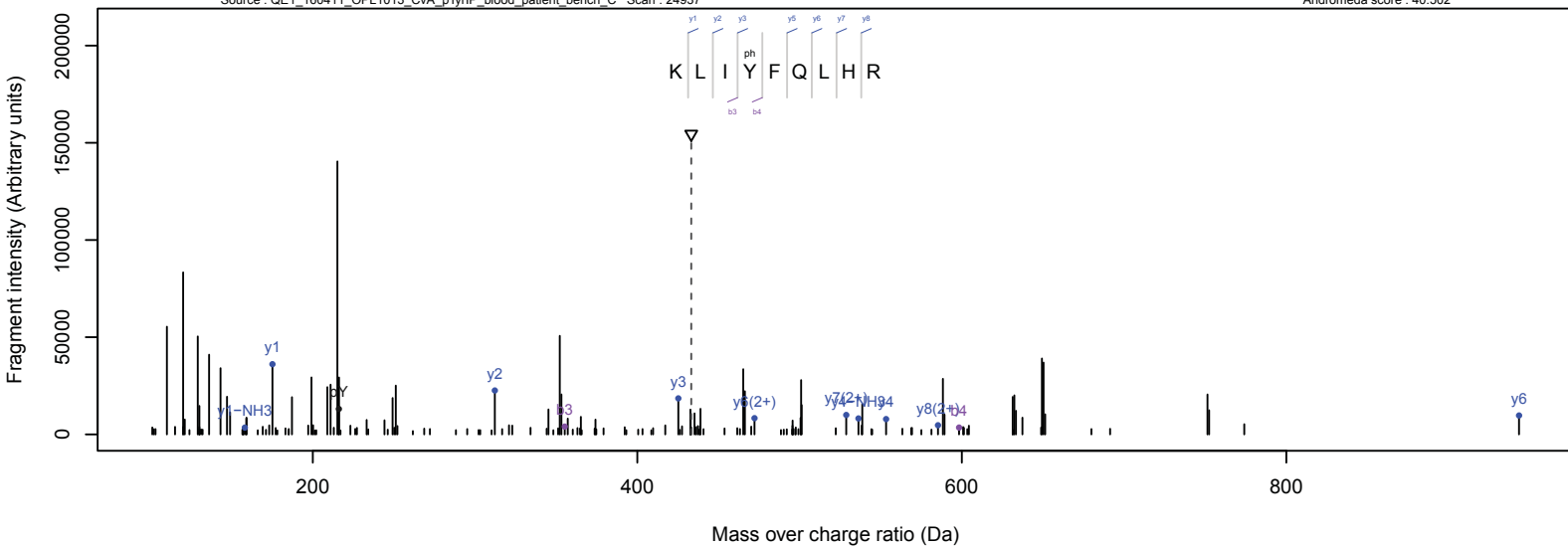
Andromeda score : 40.533



KLIY(ph)FQLHR-3+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 24937

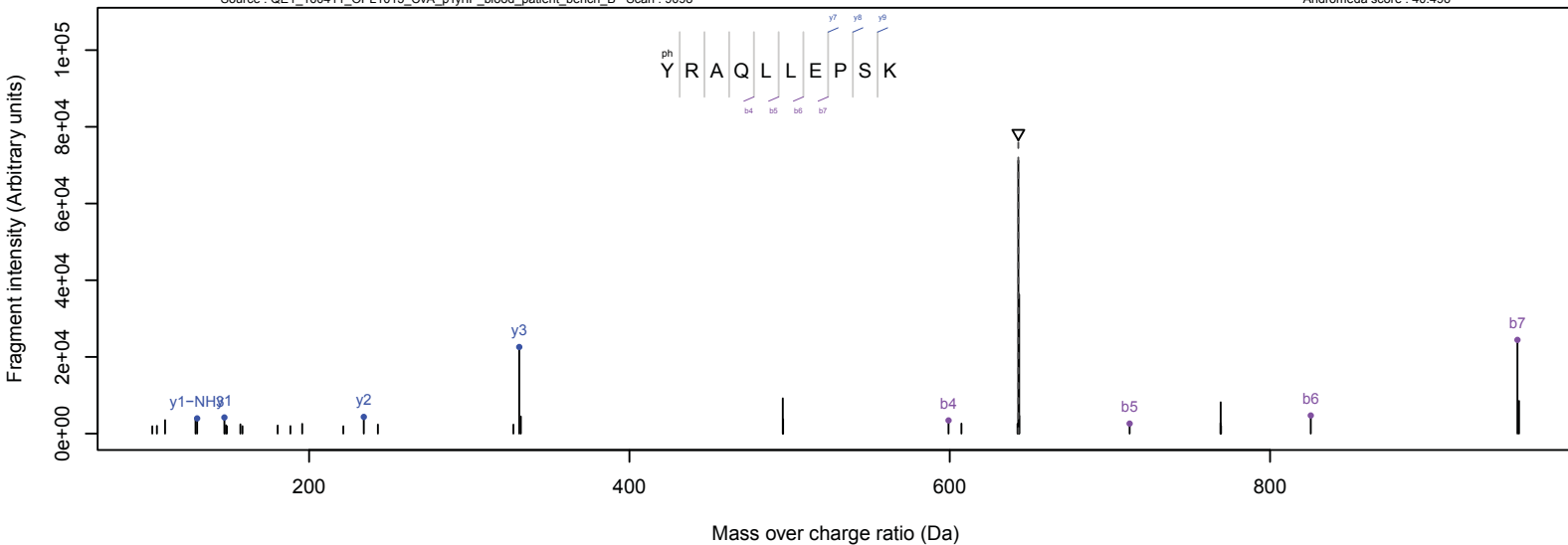
Andromeda score : 40.502



**Y(ph)RAQLLEPSK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_B Scan : 9098

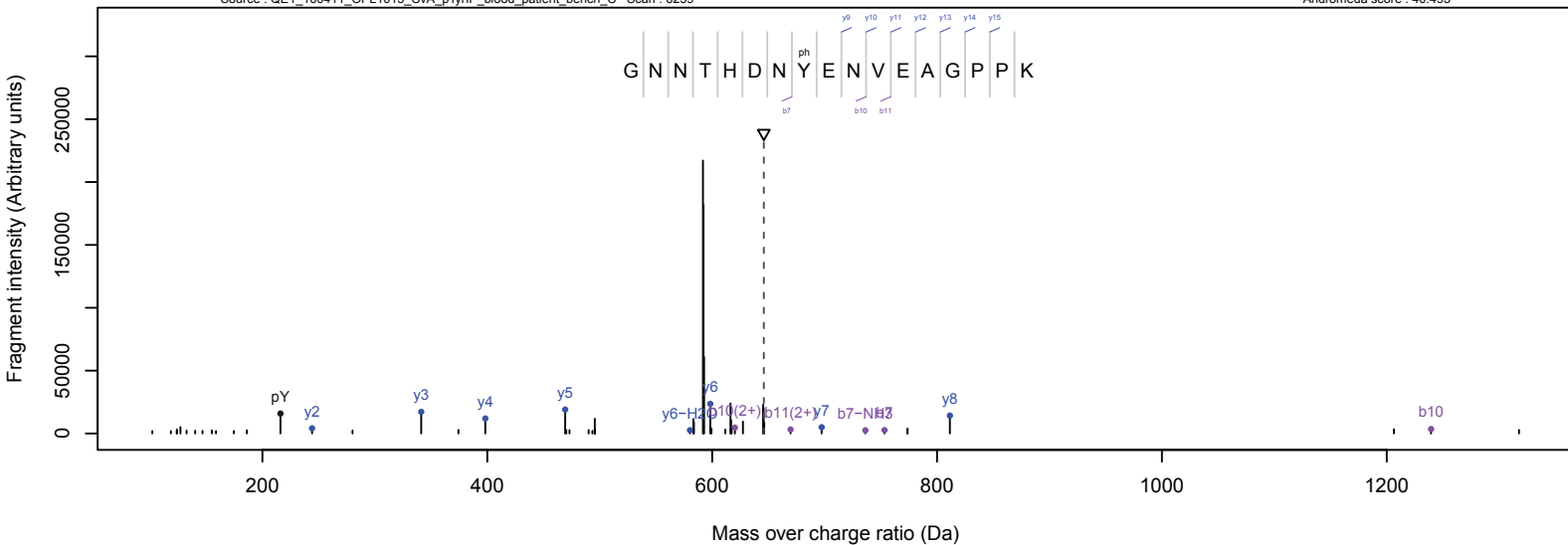
Andromeda score : 40.496



**GNNTHDNY(ph)ENVEAGPPK-3+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan : 6259

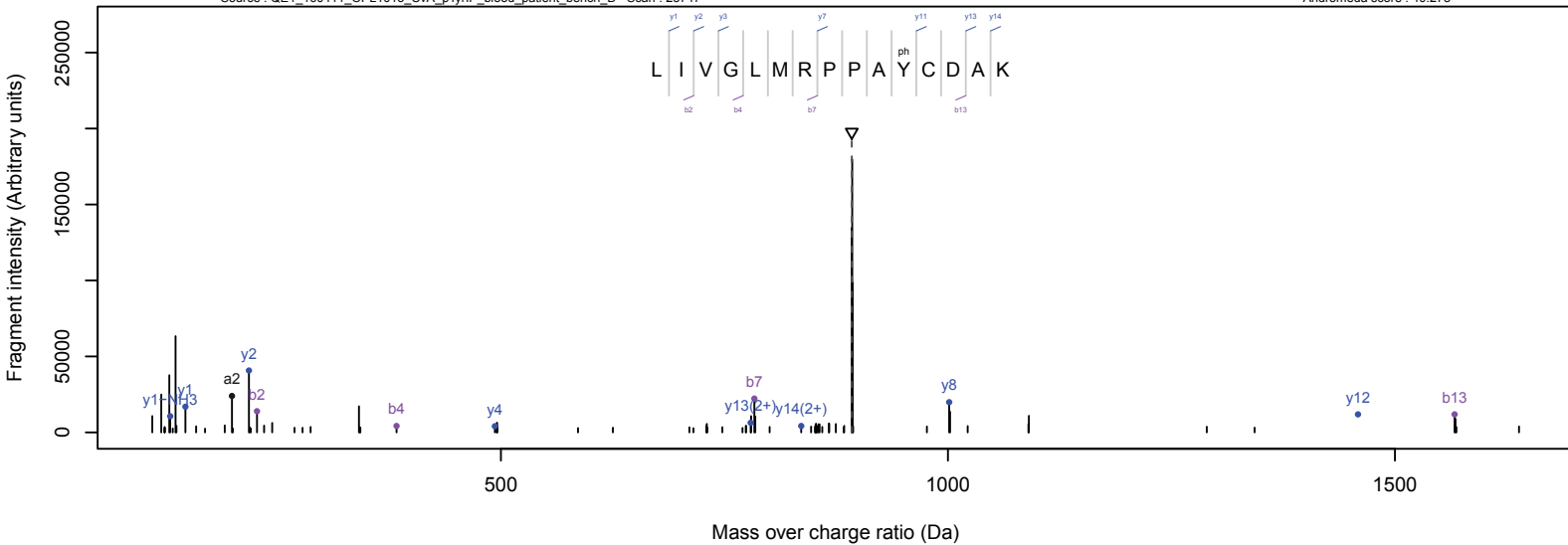
Andromeda score : 40.493



**LIVGLMRPPA(ph)CDAK-2+**

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan : 28747

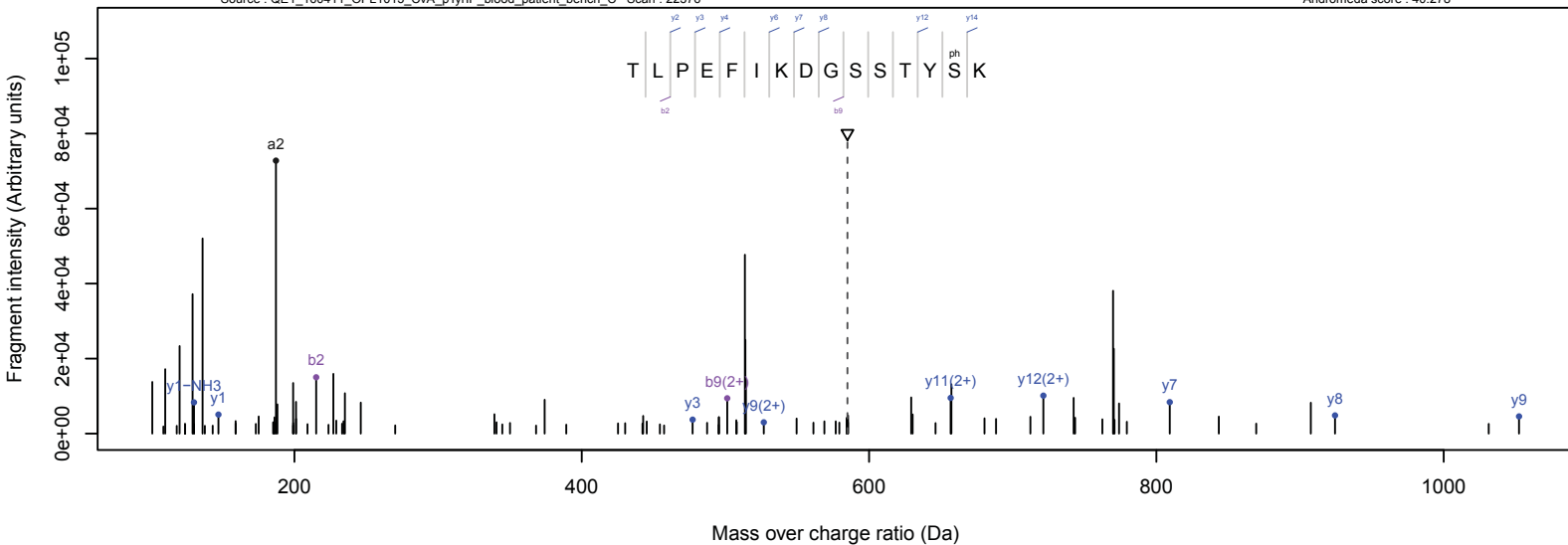
Andromeda score : 40.278



TLPEFIKGSSTYS(ph)K-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_C Scan: 22376

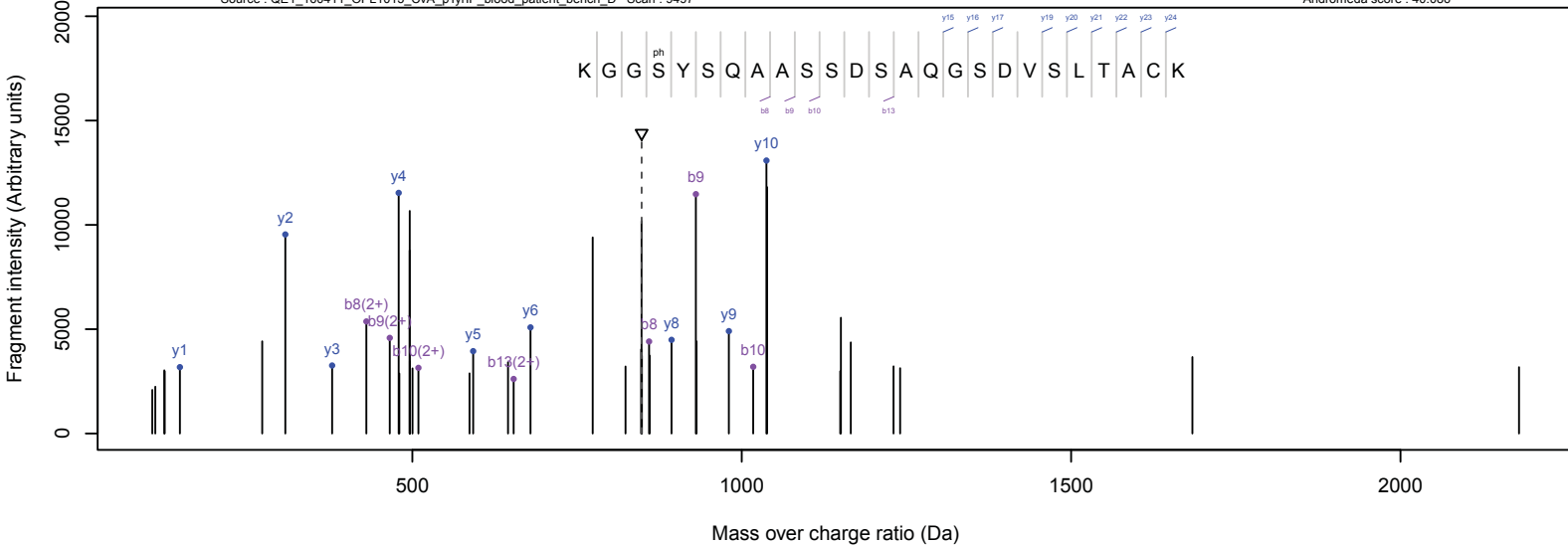
Andromeda score: 40.278



KGGS(ph)YSQAASSDSAQGS D VSLTACK-3+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 9497

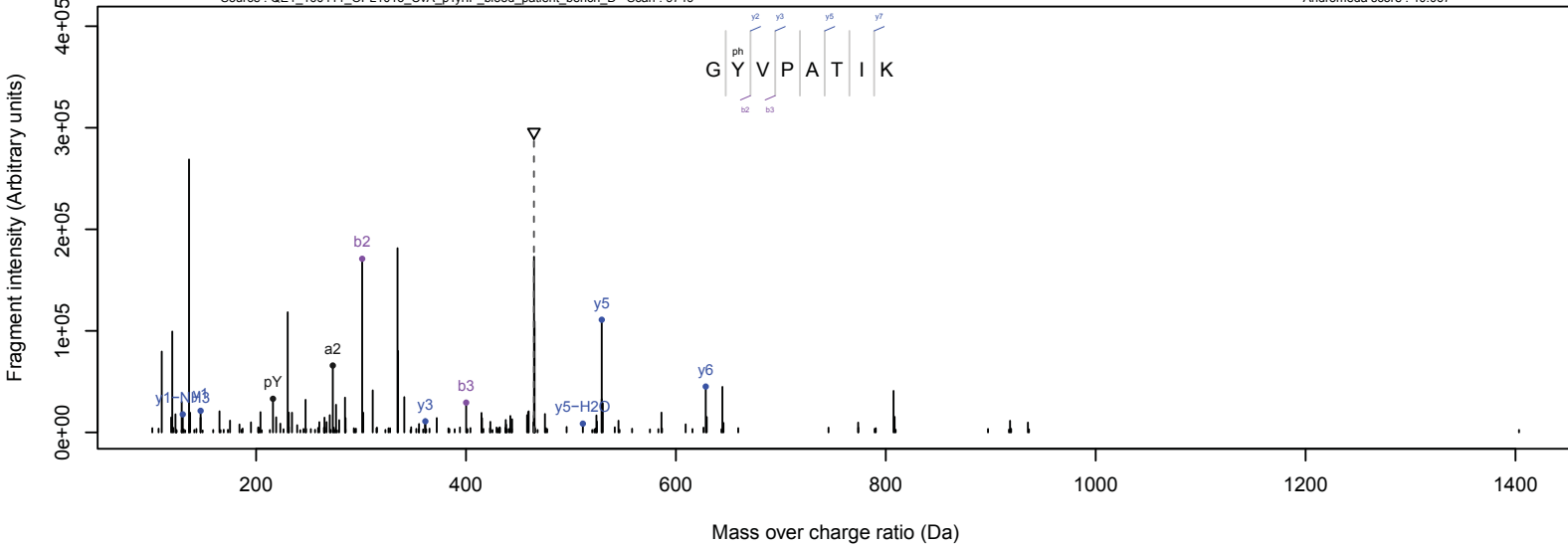
Andromeda score: 40.086



GY(ph)VPATIK-2+

Source: QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_D Scan: 9749

Andromeda score: 40.067



## SY(ph)VAWDR-2+

Source : QE1\_160411\_OPL1013\_CvA\_pTyrIP\_blood\_patient\_bench\_A Scan : 11808

Andromeda score : 40.067

