

Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

6102

Method

TOF; CID

Score

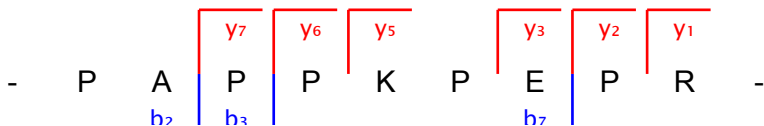
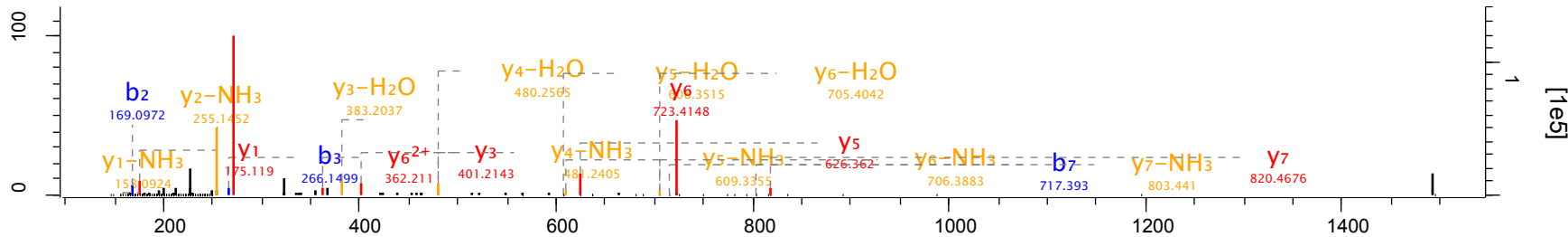
107.69

m/z

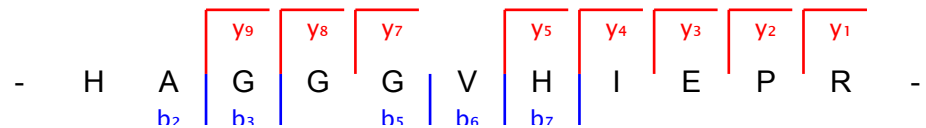
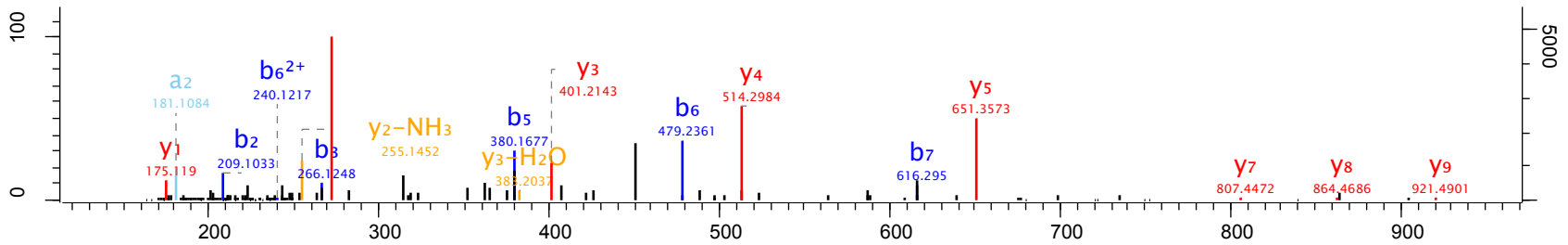
330.19

Gene names

HMGN4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract1_dyn_5ul_E1_01_580	6441	TOF; CID	90.05	377.2	NDUFB2



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

8834

Method

TOF; CID

Score

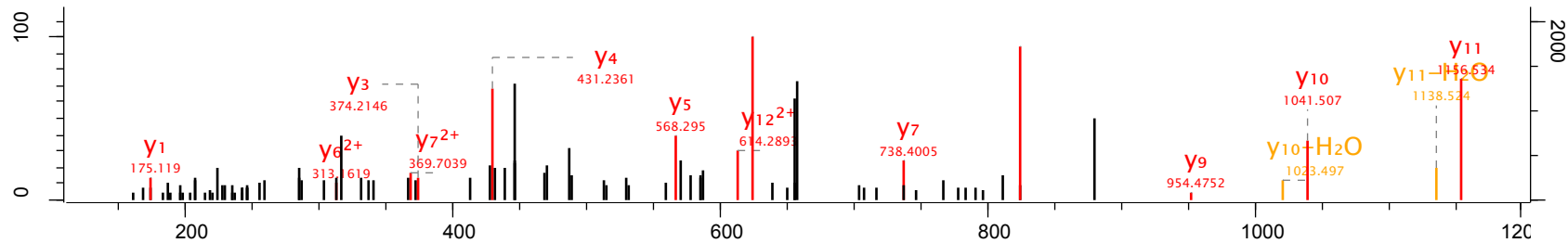
70.09

m/z

447.55

Gene names

BTBD7



- L A D S E S L G H G A Q R -

Fragmentation sites (y-ions) are indicated by red brackets above the amino acid sequence:

- y12²⁺ (between L and A)
- y11 (between A and D)
- y10 (between D and S)
- y9 (between S and E)
- y8 (between E and S)
- y7 (between S and L)
- y6 (between L and G)
- y5 (between G and H)
- y4 (between H and G)
- y3 (between G and A)
- y1 (between A and Q)

Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

9299

Method

TOF; CID

Score

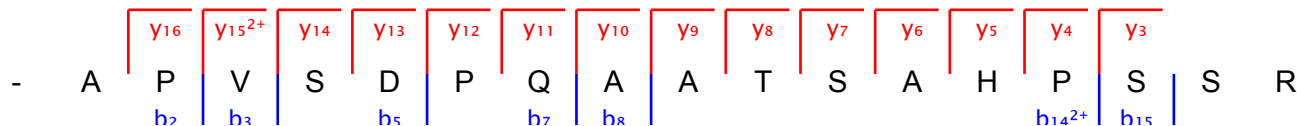
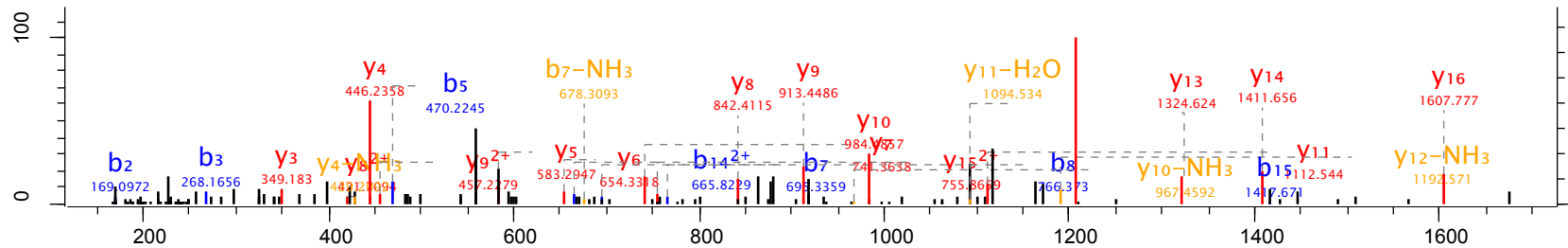
66.55

m/z

560.28

Gene names

SPSB3



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

13546

Method

TOF; CID

Score

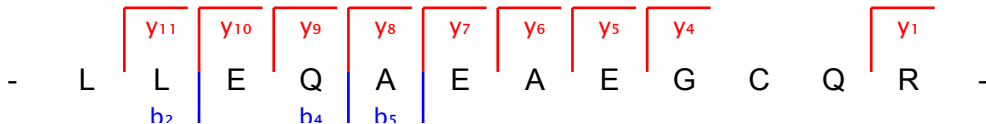
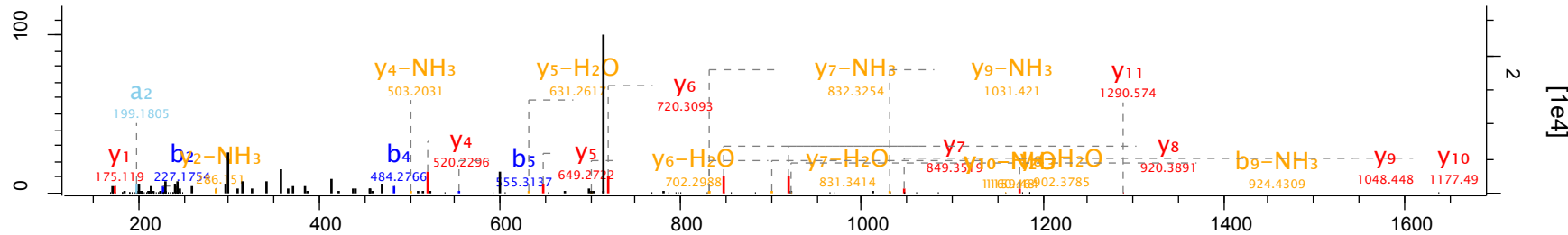
96.74

m/z

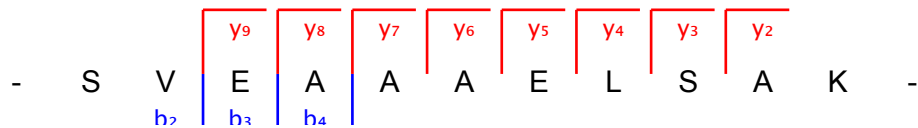
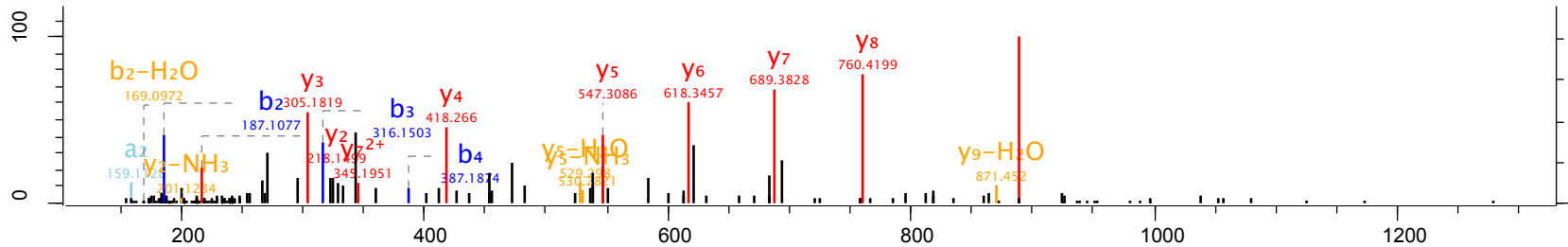
702.33

Gene names

TNFRSF10D



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract1_dyn_5ul_E1_01_580	14845	TOF; CID	65.18	538.28	PTMS



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

21407

Method

TOF; CID

Score

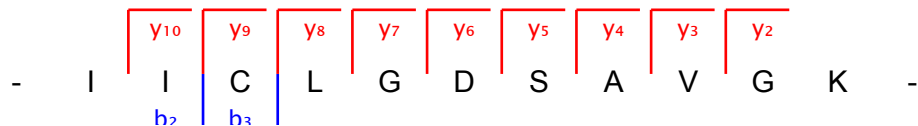
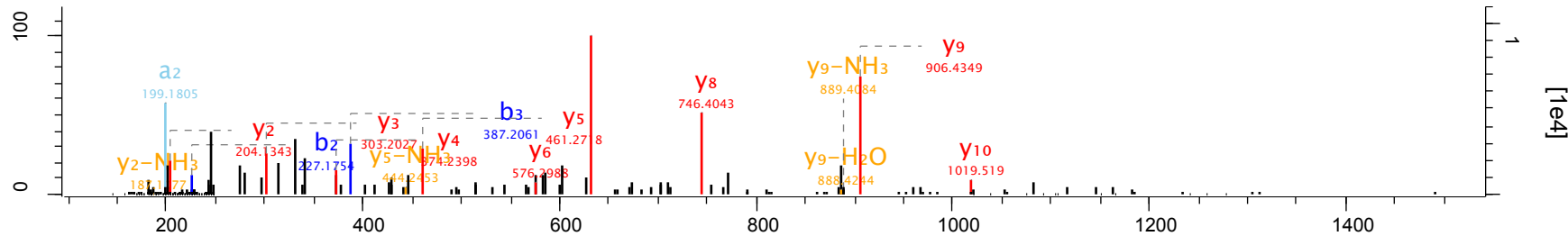
67.08

m/z

566.8

Gene names

RABL2B;RABL2A



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

22474

Method

TOF; CID

Score

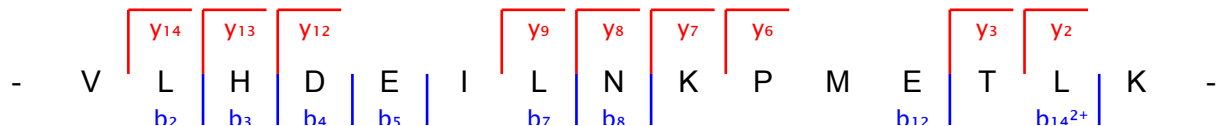
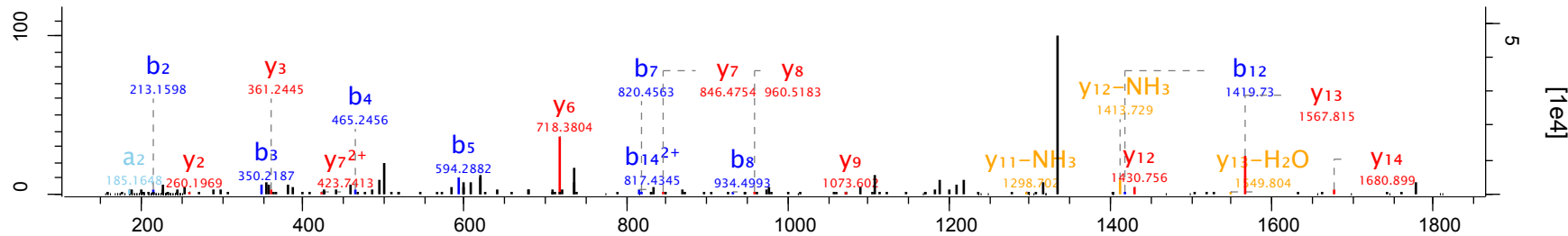
69.43

m/z

593.99

Gene names

SLC35A3



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

25904

Method

TOF; CID

Score

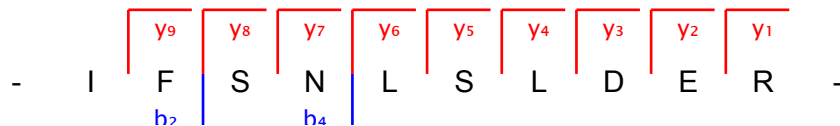
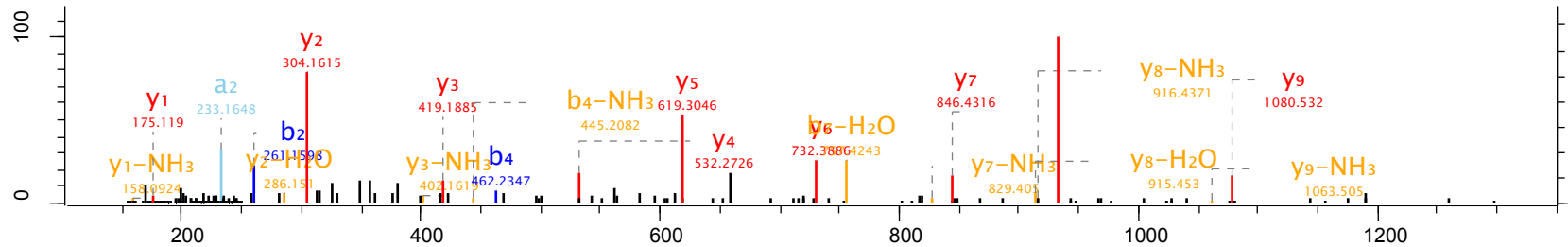
68.54

m/z

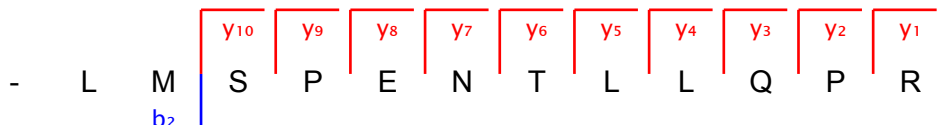
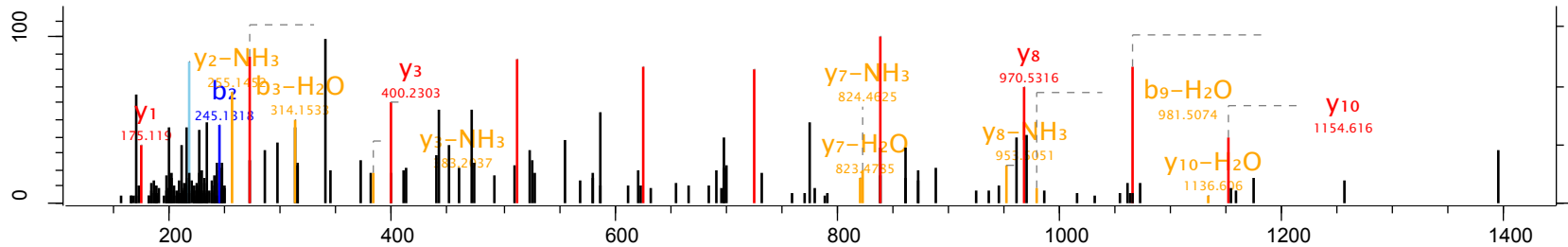
597.31

Gene names

FBXL17



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract1_dyn_5ul_E1_01_580	27070	TOF; CID	80.24	699.87	DEPTOR



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

27783

Method

TOF; CID

Score

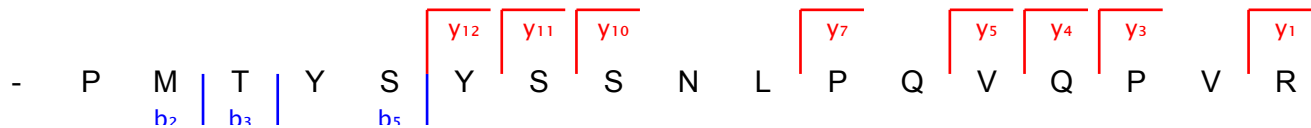
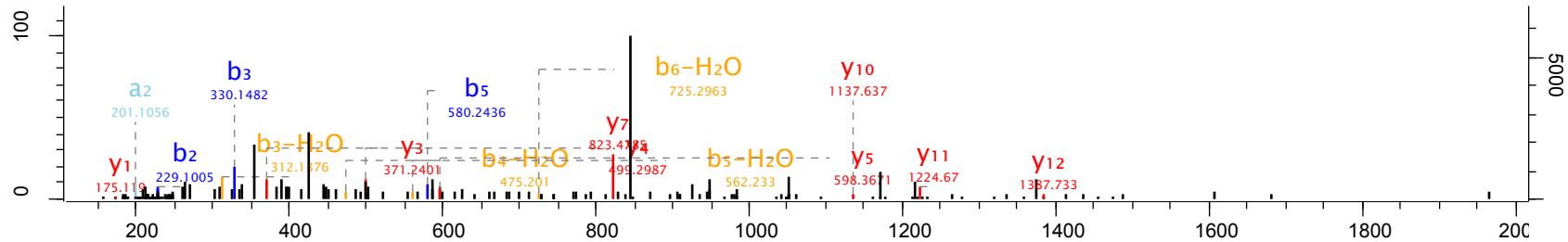
48.9

m/z

984.49

Gene names

HOXA11



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

33597

Method

TOF; CID

Score

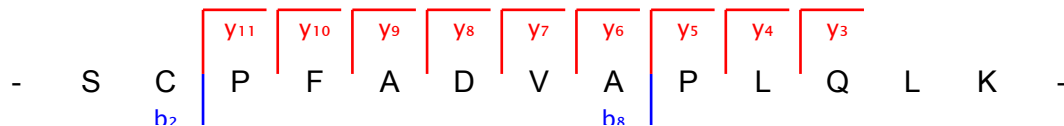
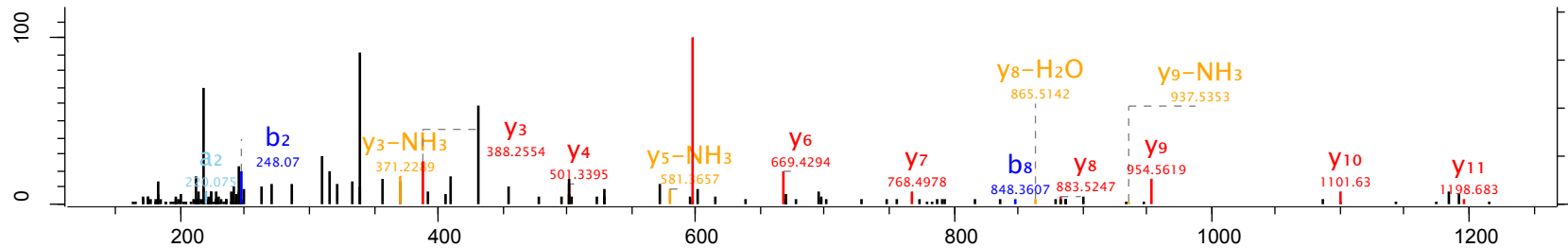
51.73

m/z

723.38

Gene names

SIVA1;GRM6



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

34618

Method

TOF; CID

Score

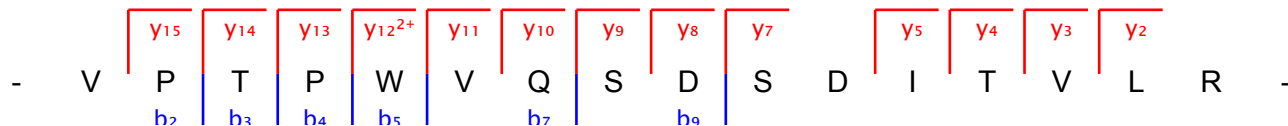
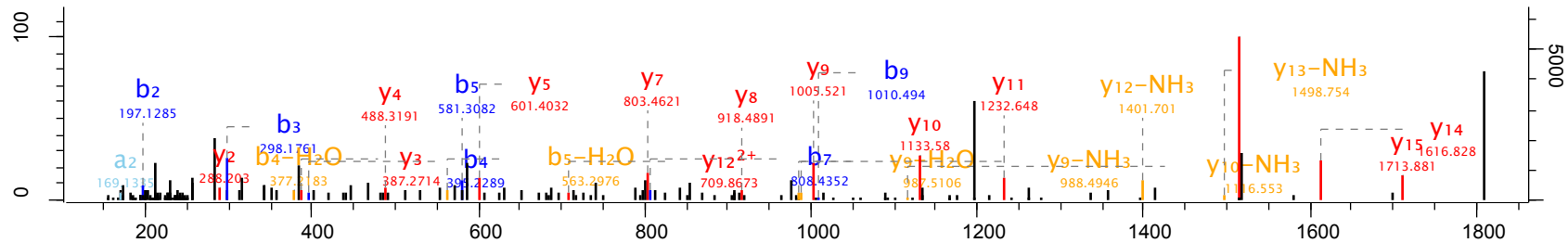
78.6

m/z

906.98

Gene names

RIMKLB



Raw file

20141014_fract1_dyn_5ul_E1_01_580

Scan

35665

Method

TOF; CID

Score

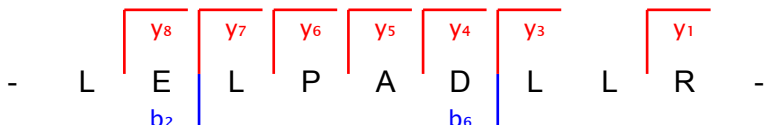
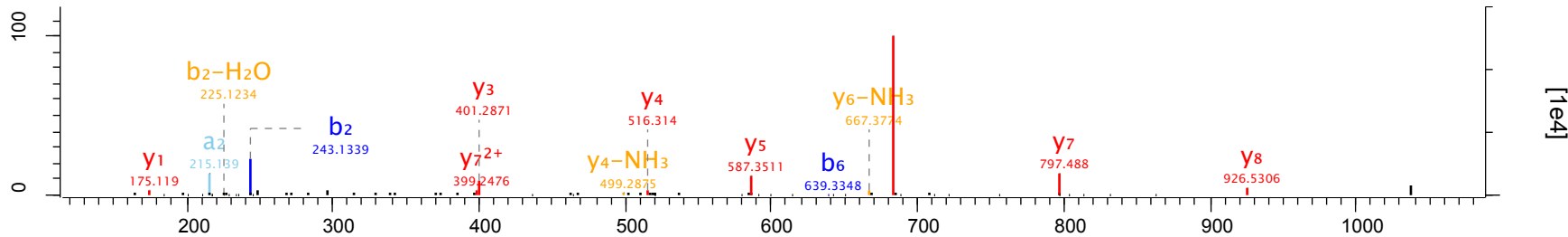
80.69

m/z

520.31

Gene names

SAMD11



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

3355

Method

TOF; CID

Score

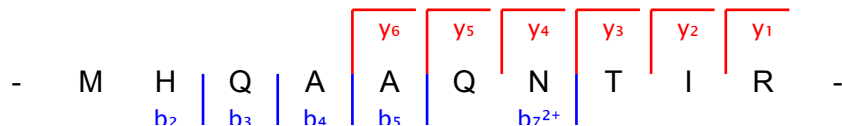
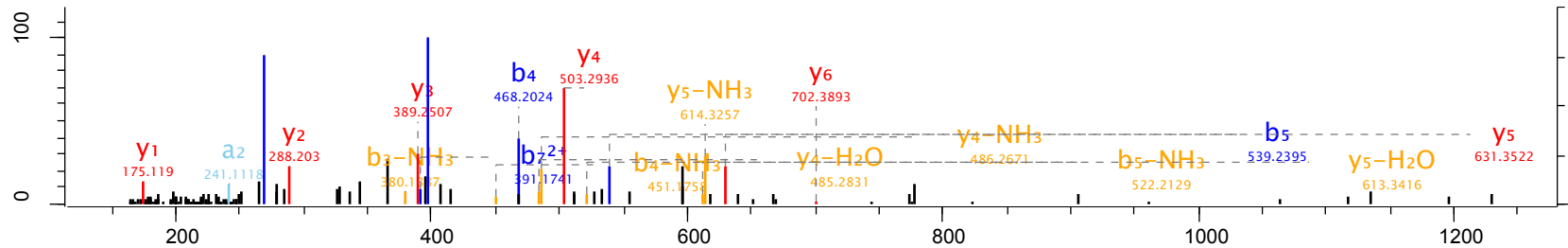
81.1

m/z

390.53

Gene names

BBIP1



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

5353

Method

TOF; CID

Score

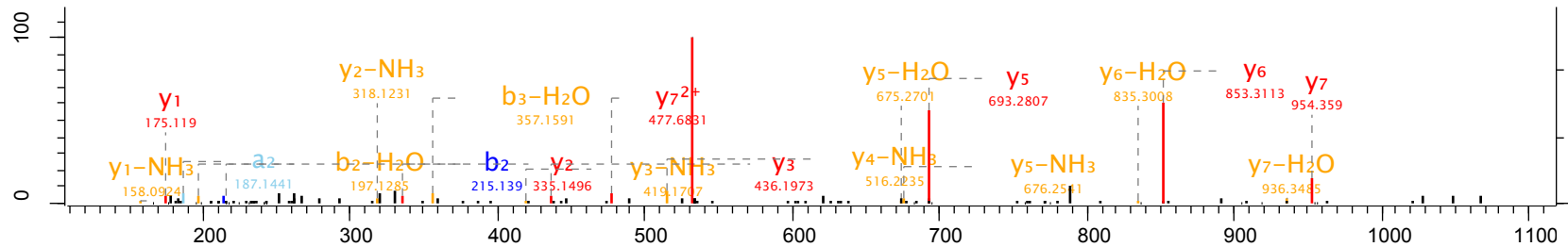
104.44

m/z

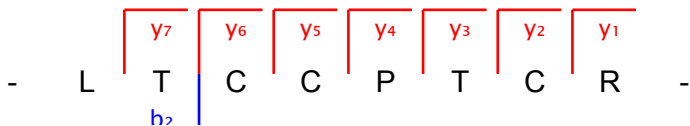
534.72

Gene names

SIAH1



[1e4]



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

5967

Method

TOF; CID

Score

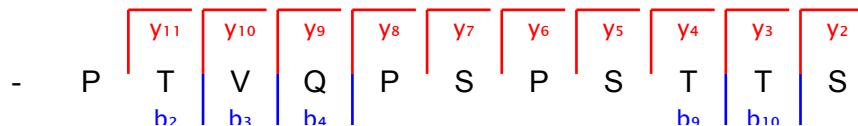
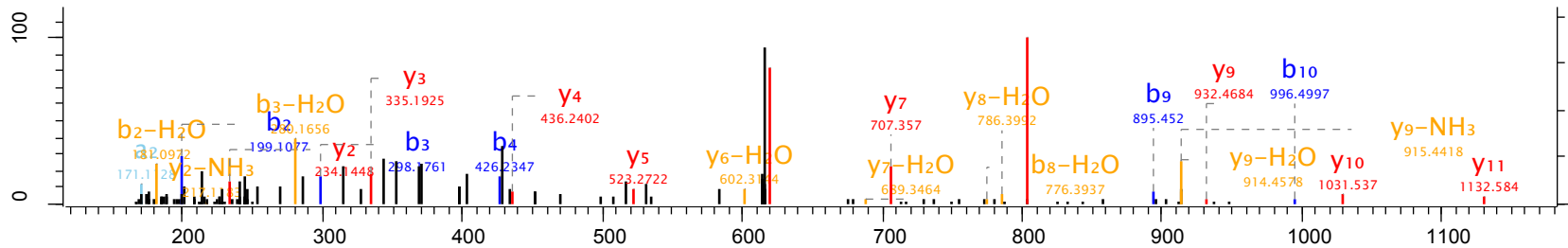
86.5

m/z

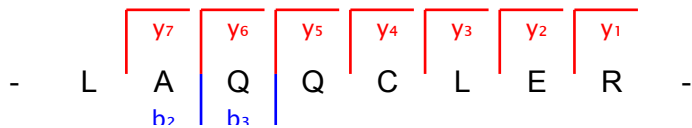
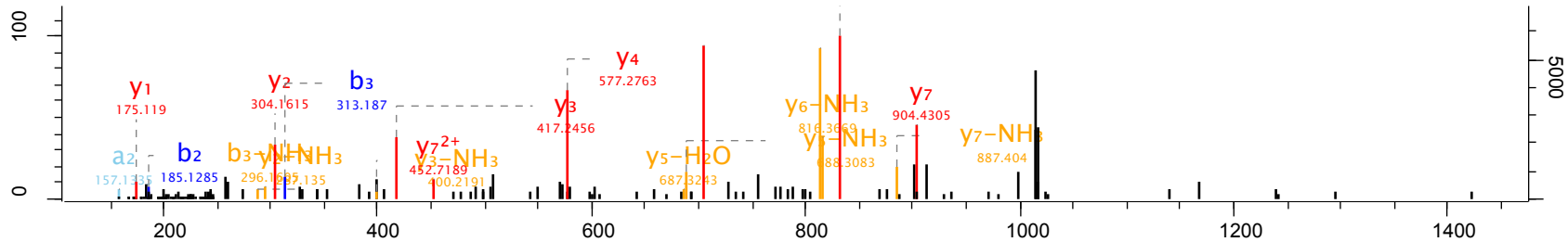
615.32

Gene names

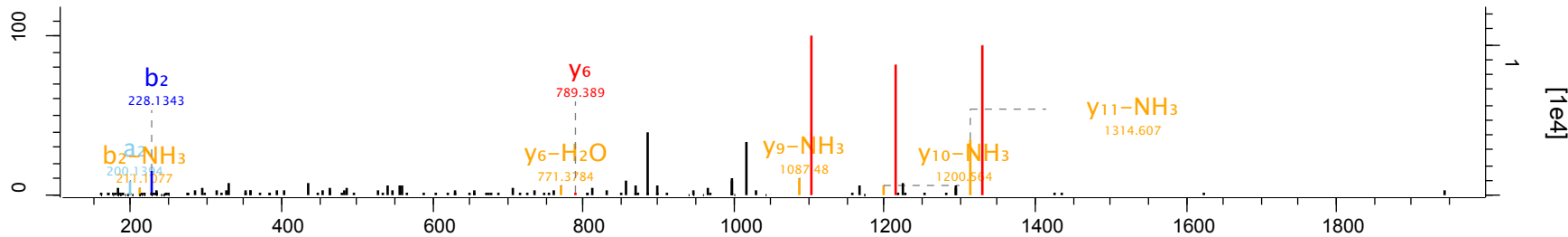
CD164



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	7794	TOF; CID	127.46	509.26	VPS9D1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	8678	TOF; CID	52.25	482.24	ZNF165



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

10721

Method

TOF; CID

Score

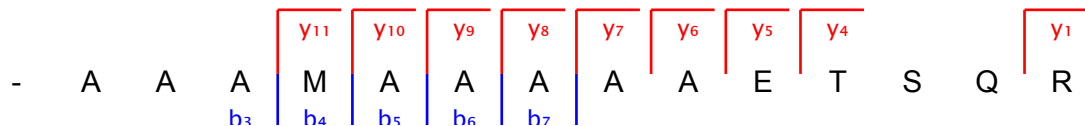
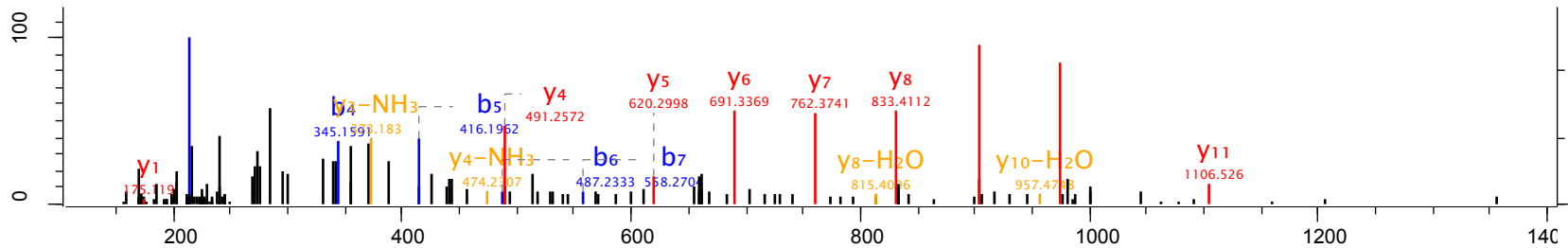
54.9

m/z

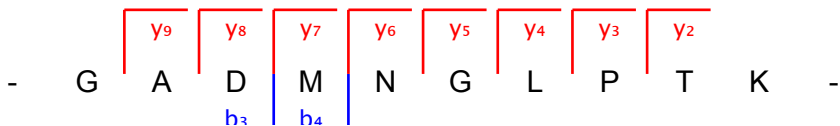
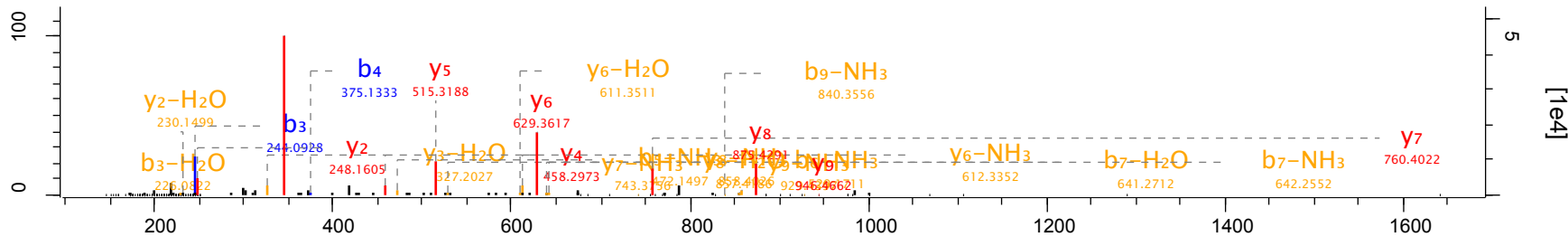
660.32

Gene names

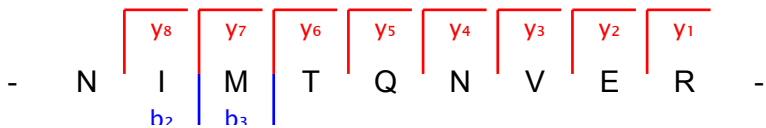
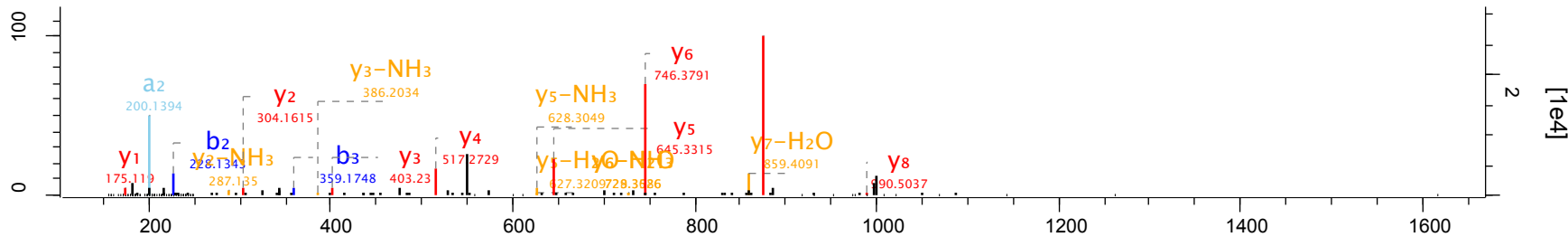
NDFIP2



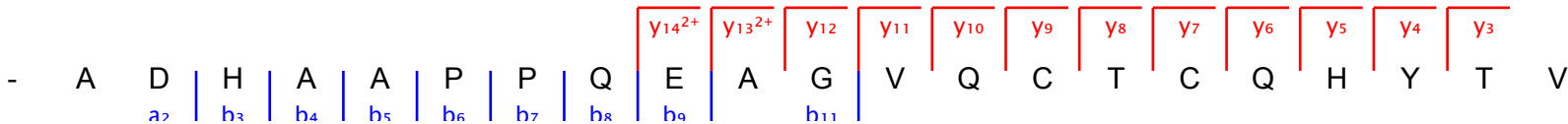
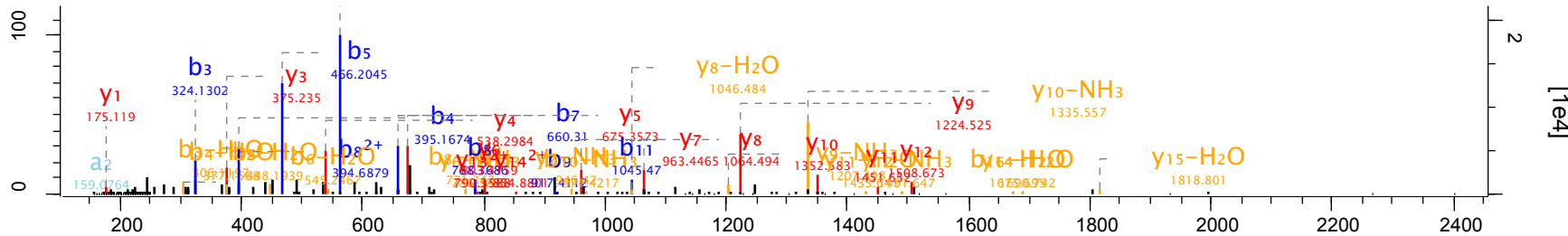
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	11931	TOF; CID	87.18	502.25	C18orf32



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	12068	TOF; CID	95.52	552.78	VAMP8



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	12220	TOF; CID	113.51	625.03	C11orf48



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

12521

Method

TOF; CID

Score

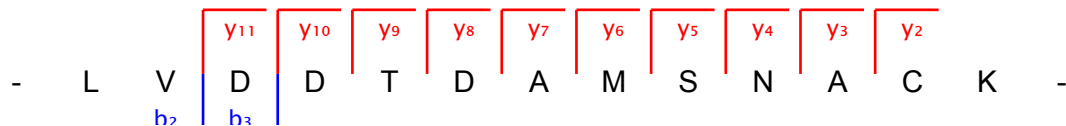
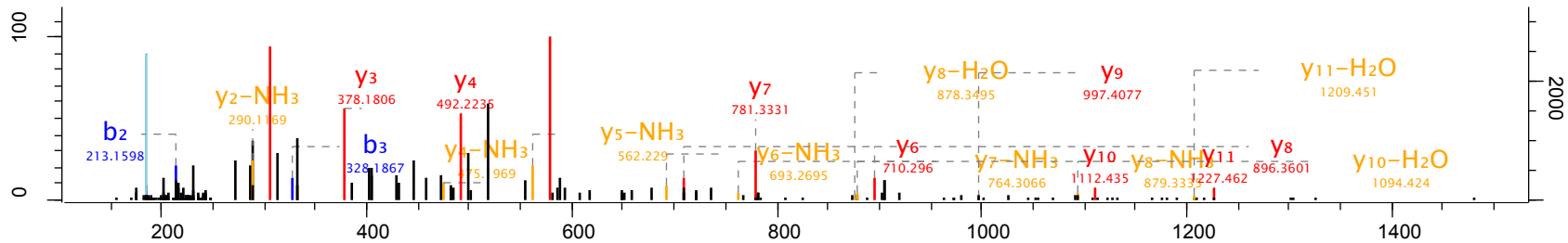
94.12

m/z

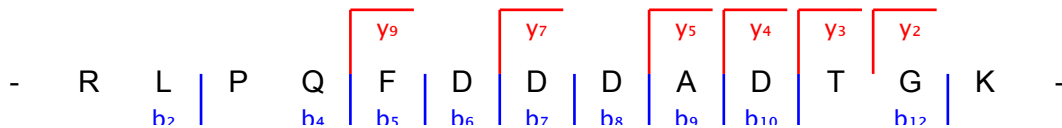
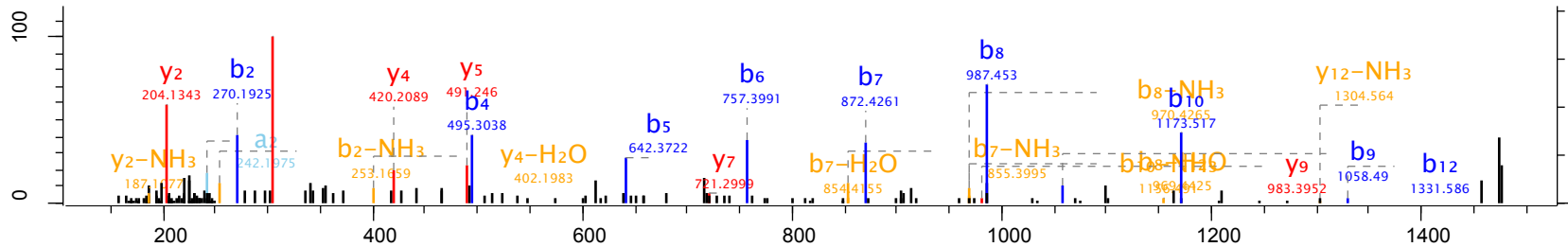
720.31

Gene names

LEPROTL1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	14642	TOF; CID	116.19	739.35	ZNHIT1



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

18284

Method

TOF; CID

Score

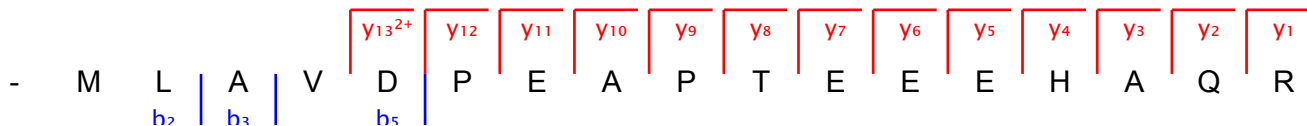
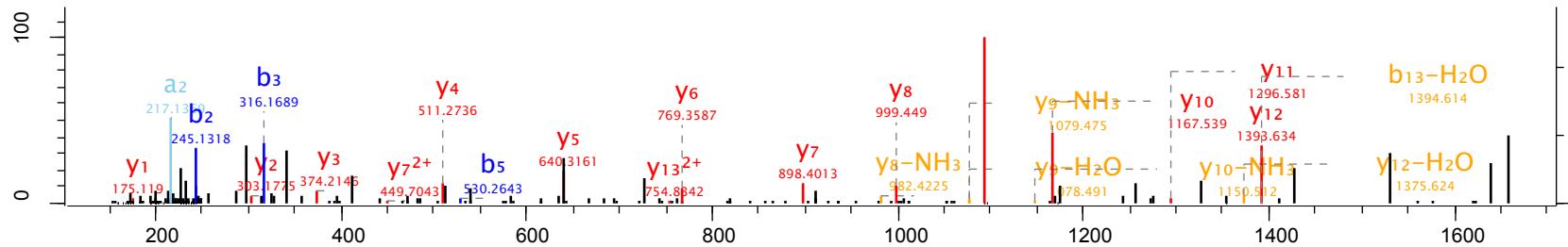
64.54

m/z

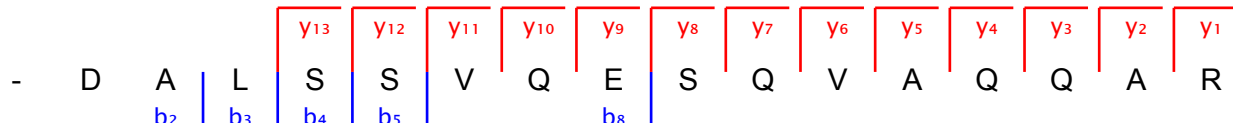
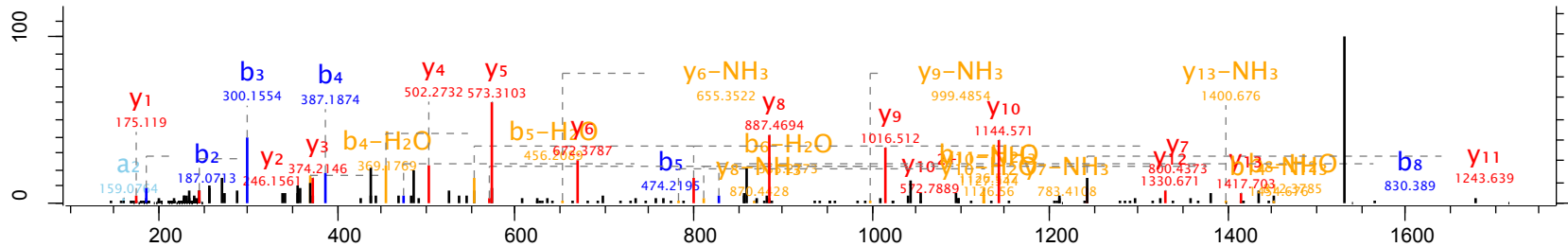
641.97

Gene names

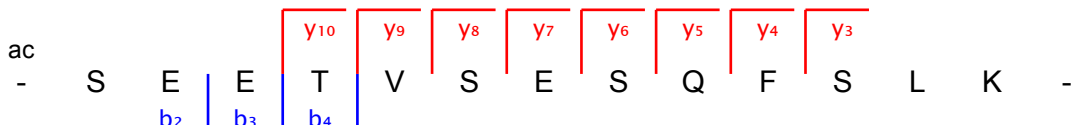
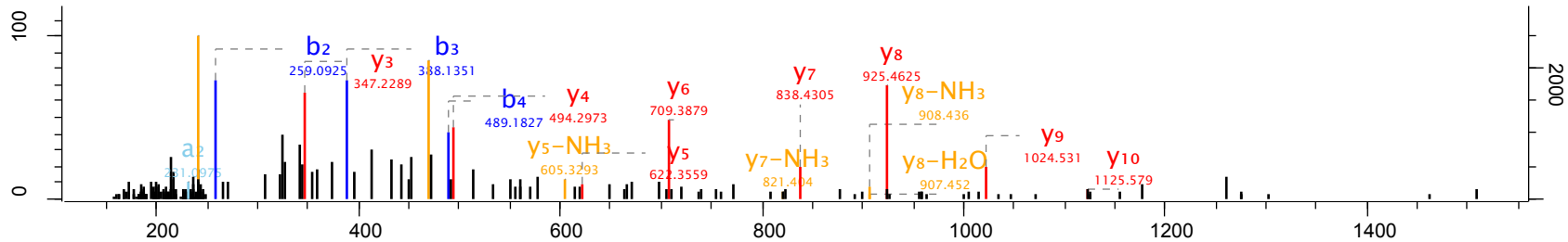
ITPKA



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	18570	TOF; CID	105.63	859.93	APOC3



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	26560	TOF; CID	73.93	756.86	FAM73A



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

26737

Method

TOF; CID

Score

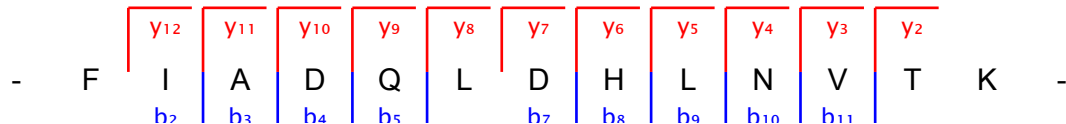
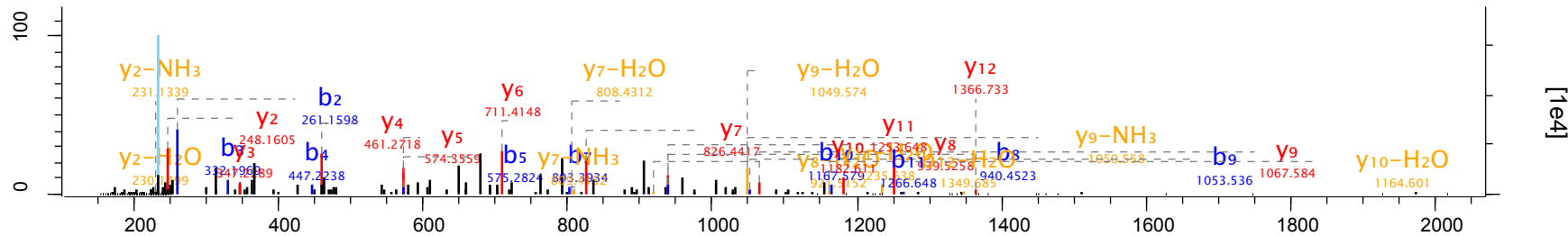
133.48

m/z

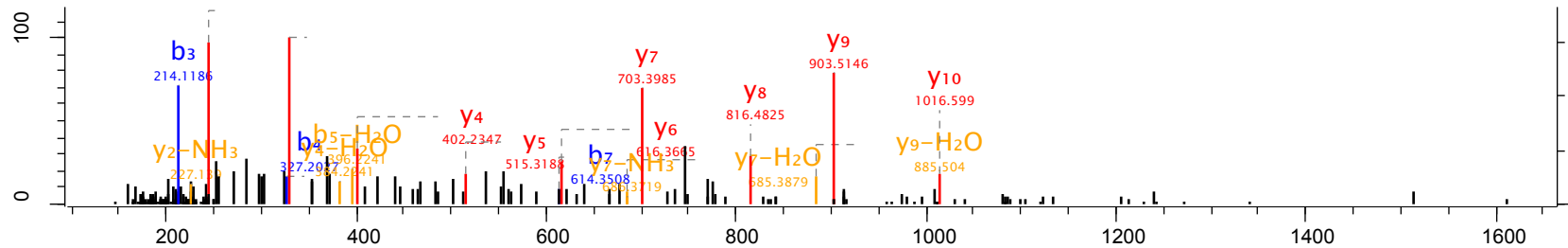
757.4

Gene names

MRPL57



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	27754	TOF; CID	49.3	615.36	EFNB1



Sequence: - A A A L S L S T L A S P K -

Fragmentation sites (b and y ions) are indicated by brackets below the sequence:

- b₃ (blue) is between the 3rd and 4th residues (A and L).
- b₄ (blue) is between the 4th and 5th residues (L and S).
- b₇ (blue) is between the 7th and 8th residues (S and T).
- y₂ (red) is between the 1st and 2nd residues (A and A).
- y₃ (red) is between the 2nd and 3rd residues (A and A).
- y₄ (red) is between the 3rd and 4th residues (A and L).
- y₅ (red) is between the 4th and 5th residues (L and S).
- y₆ (red) is between the 5th and 6th residues (S and T).
- y₇ (red) is between the 6th and 7th residues (T and L).
- y₈ (red) is between the 7th and 8th residues (L and A).
- y₉ (red) is between the 8th and 9th residues (A and S).
- y₁₀ (red) is between the 9th and 10th residues (S and P).

Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

28473

Method

TOF; CID

Score

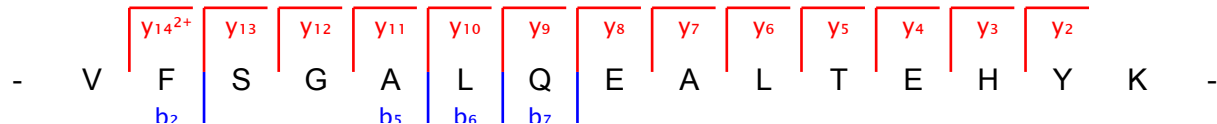
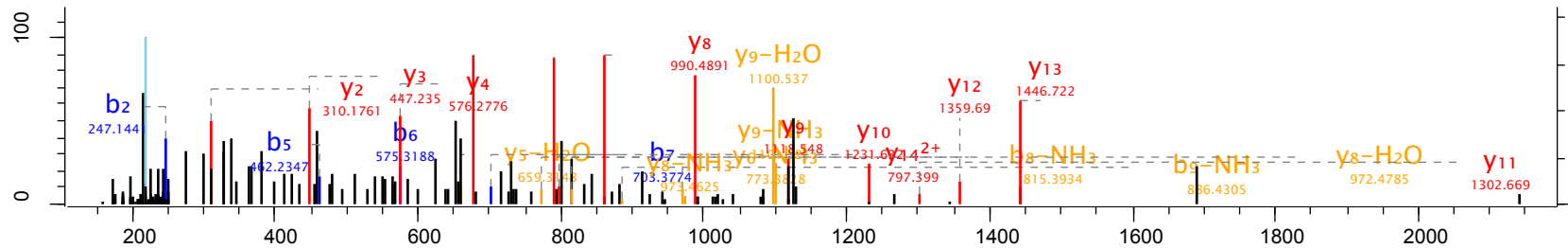
94.49

m/z

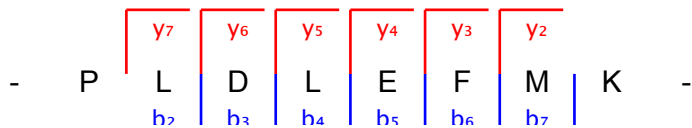
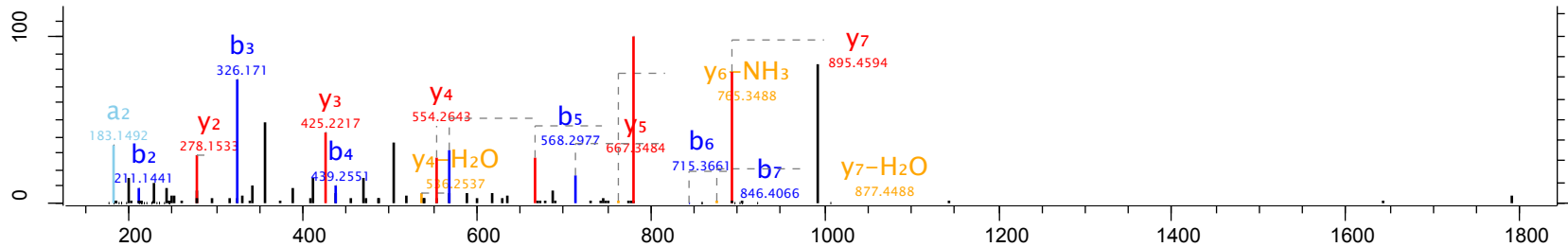
565.29

Gene names

BTG2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	30966	TOF; CID	109.11	496.76	SEPT3



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

33792

Method

TOF; CID

Score

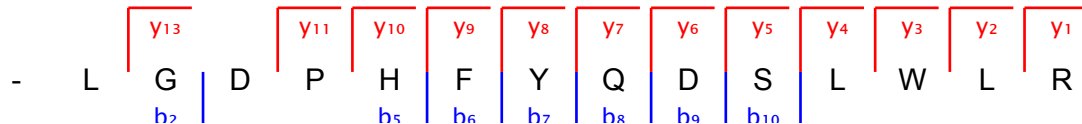
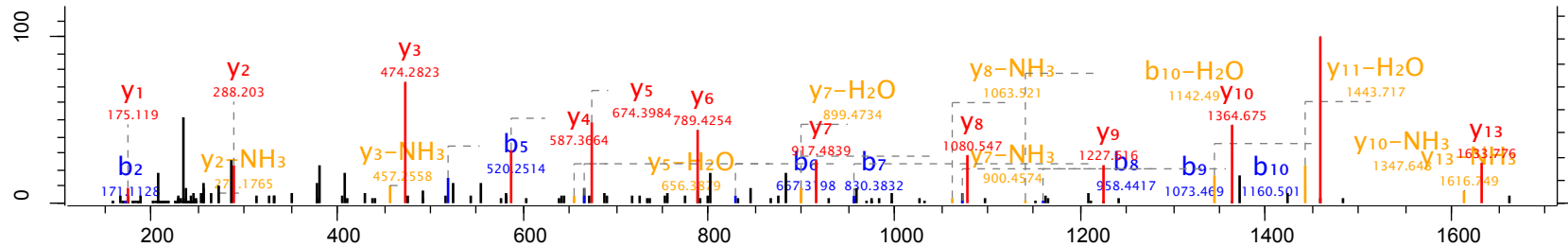
136.83

m/z

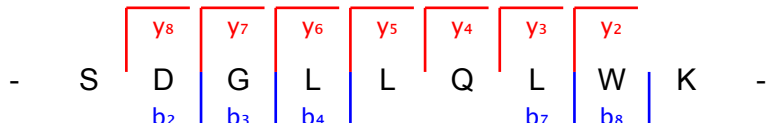
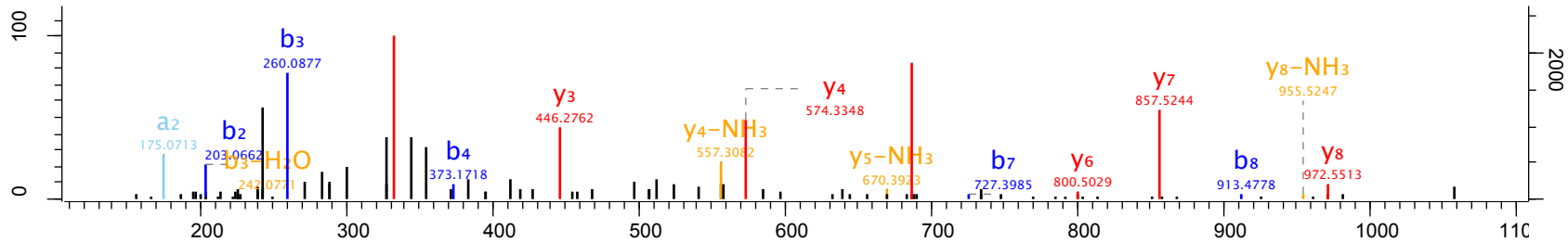
582.95

Gene names

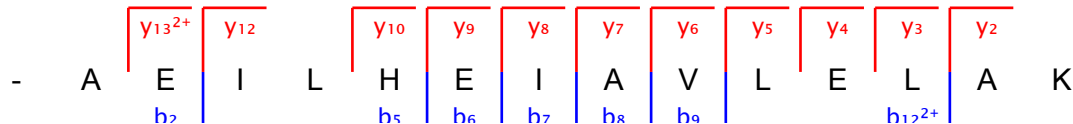
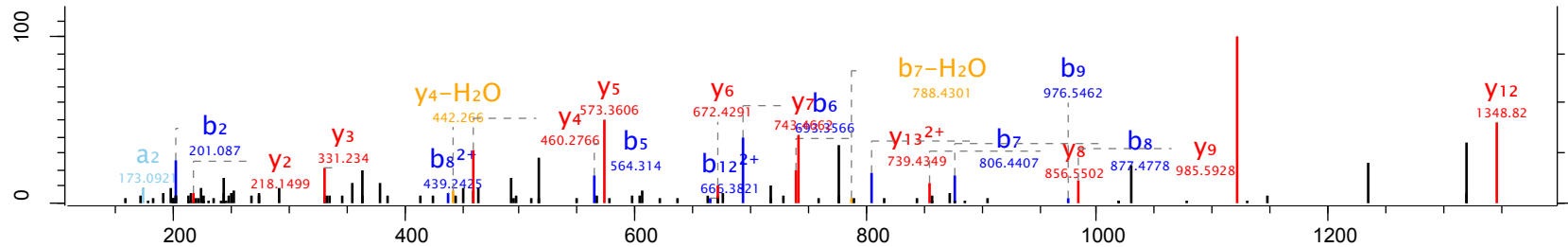
TMEM138



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	35148	TOF; CID	90.15	530.29	PHLDA1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract2_dyn_5ul_E2_01_581	37360	TOF; CID	82.65	516.97	STK17B



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

38373

Method

TOF; CID

Score

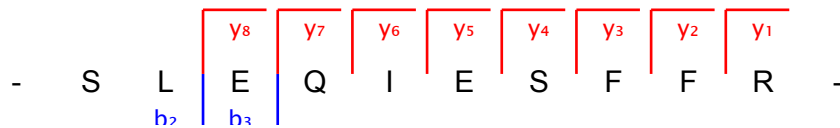
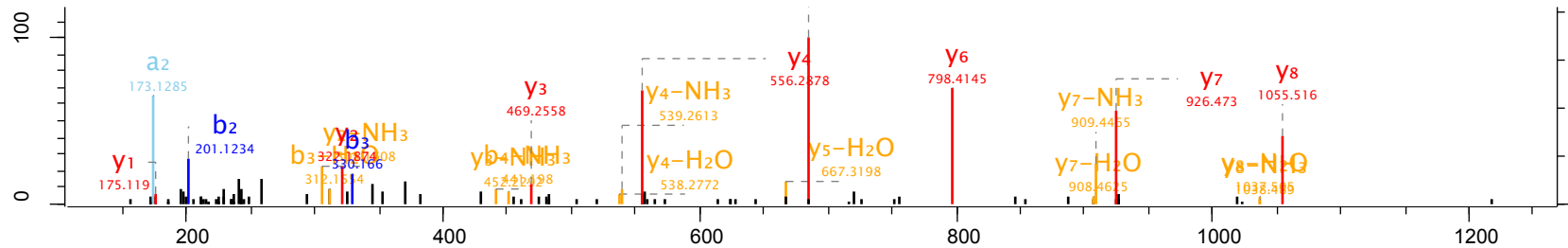
142.43

m/z

628.32

Gene names

SLC2A6



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

41523

Method

TOF; CID

Score

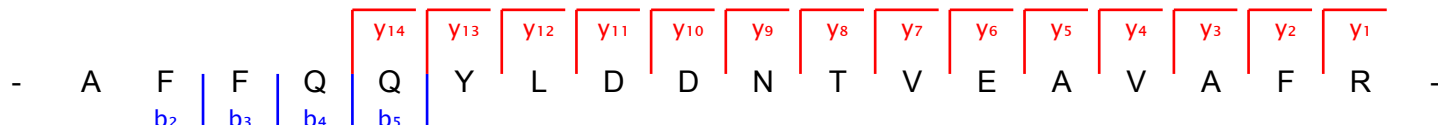
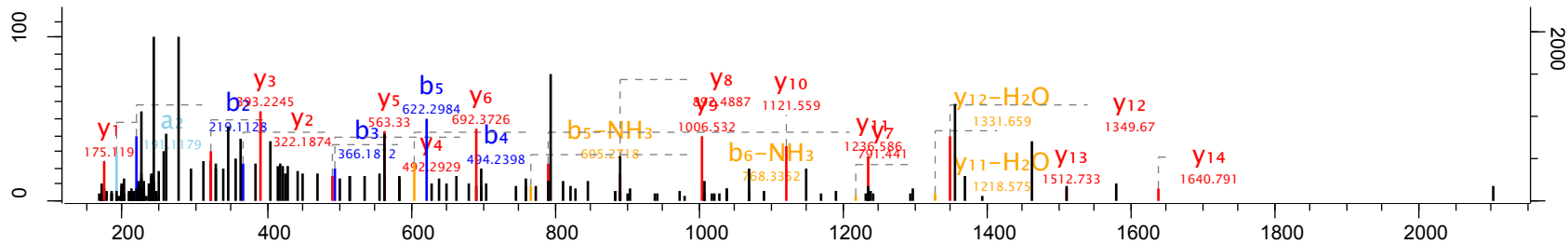
54.96

m/z

1068.02

Gene names

FKTN



Raw file

20141014_fract2_dyn_5ul_E2_01_581

Scan

41524

Method

TOF; CID

Score

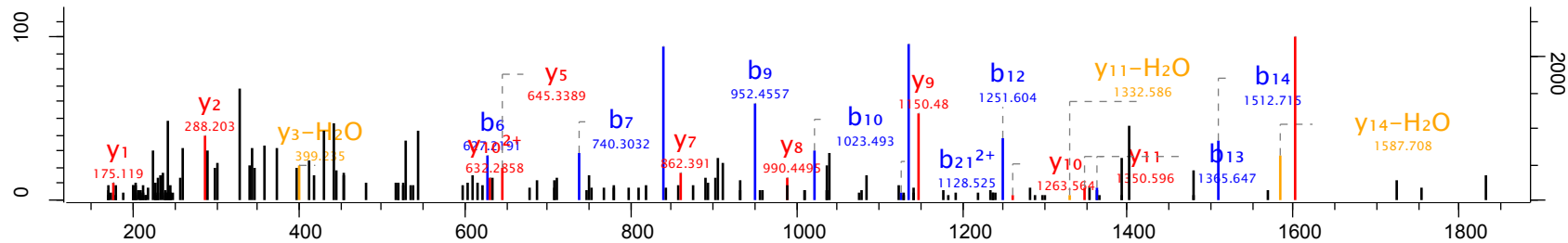
43.73

m/z

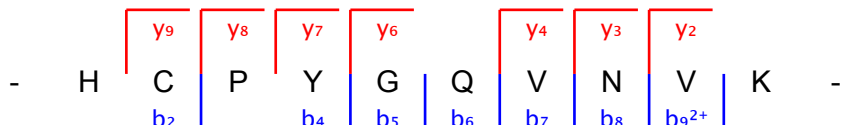
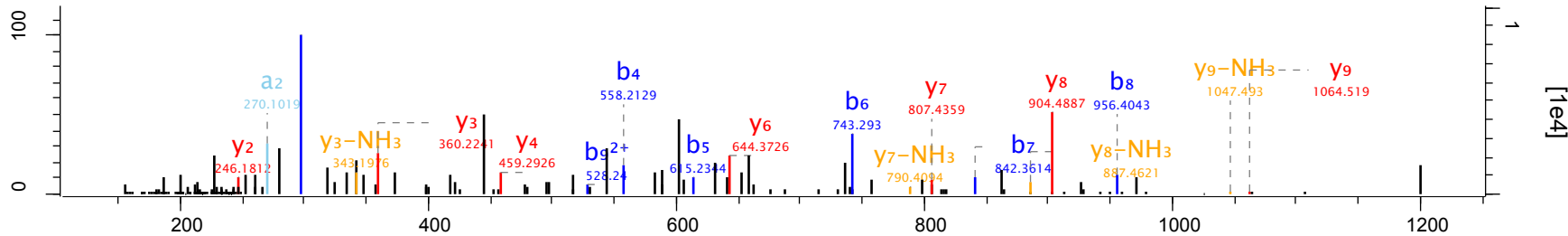
1040.48

Gene names

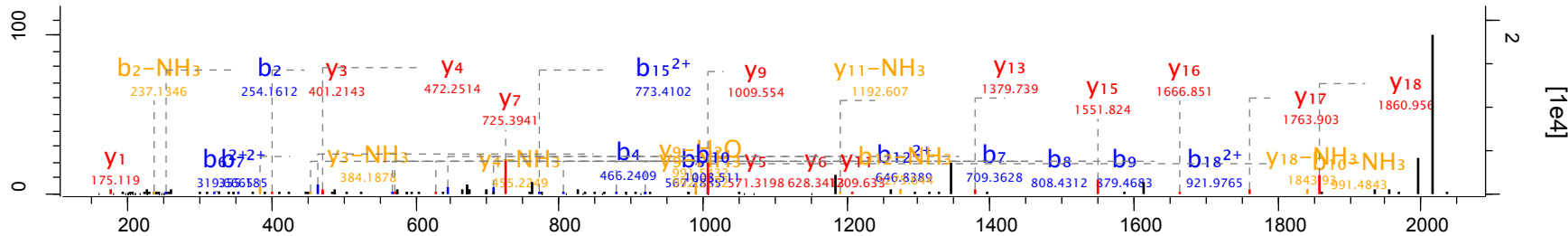
FUOM



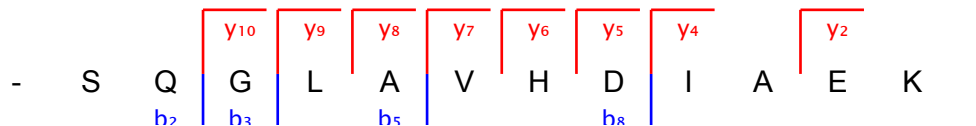
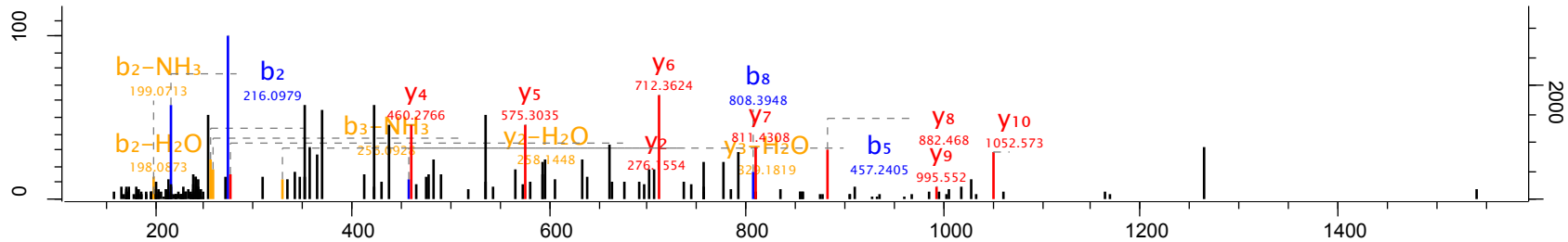
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	7727	TOF; CID	104.01	601.79	C1orf53



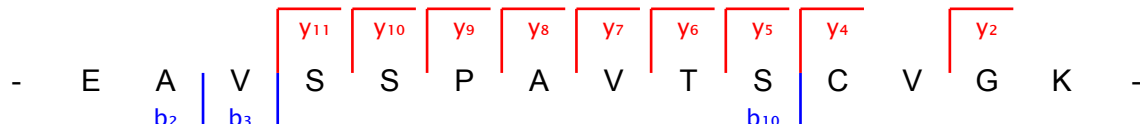
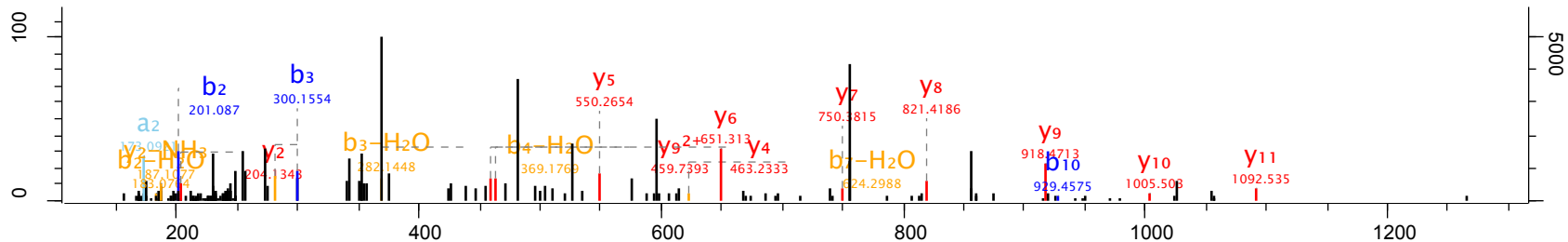
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	11647	TOF; CID	84.16	673.36	ZNF784



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	12784	TOF; CID	55.72	634.33	ADCK5



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	13752	TOF; CID	49.22	696.34	KLHL24



Raw file

20141014_fract3_dyn_5ul_E3_01_582

Scan

13989

Method

TOF; CID

Score

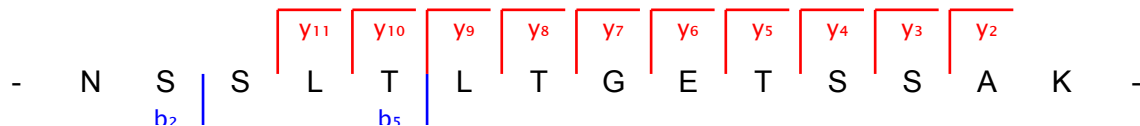
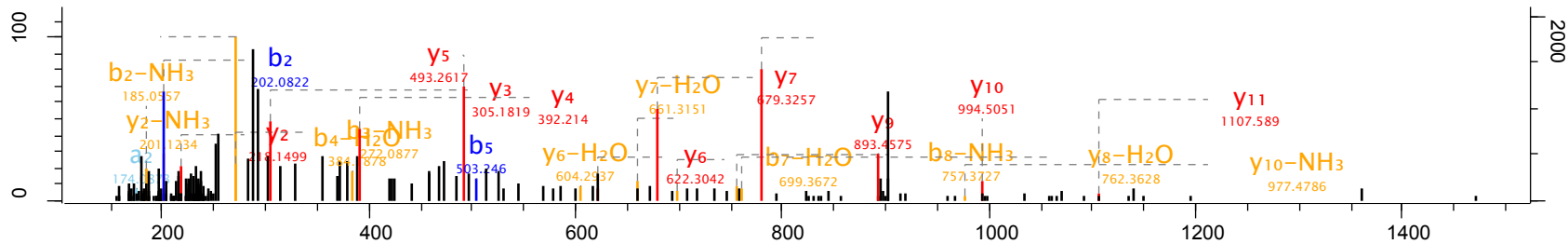
54.81

m/z

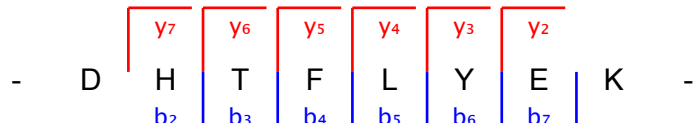
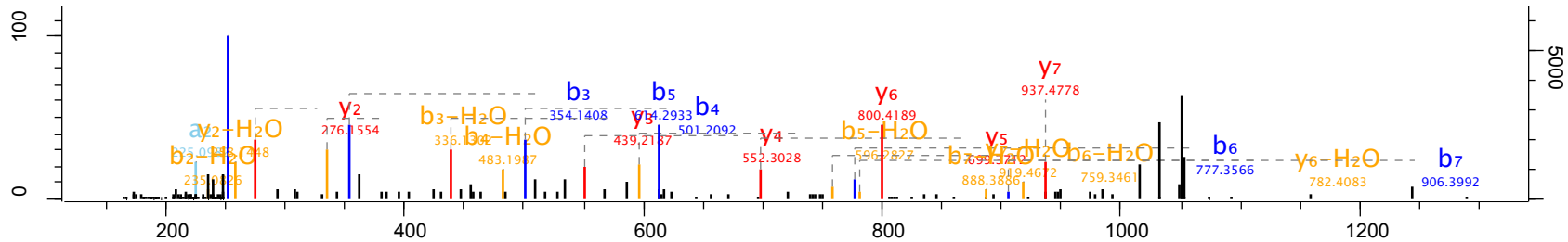
698.35

Gene names

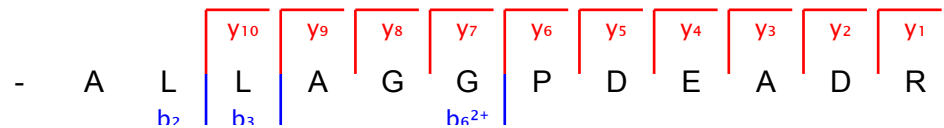
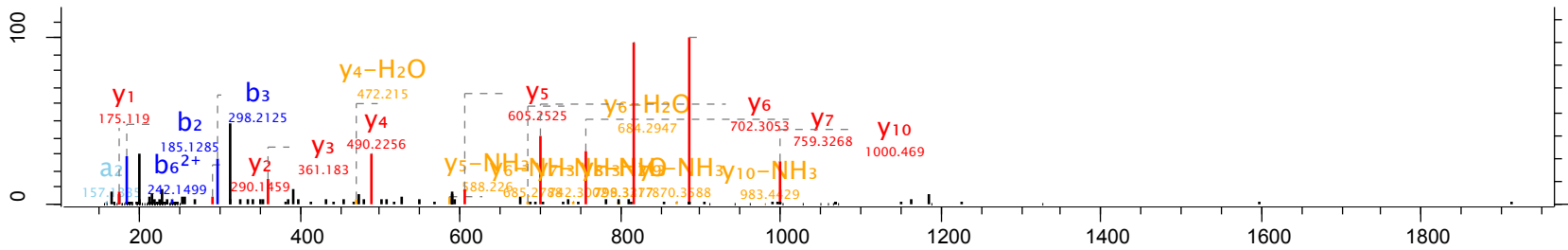
IFT43



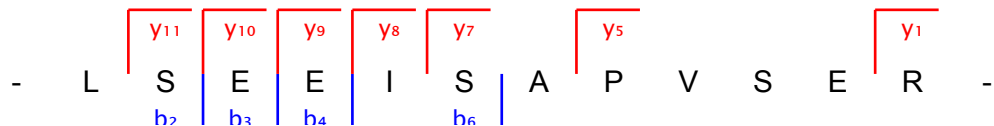
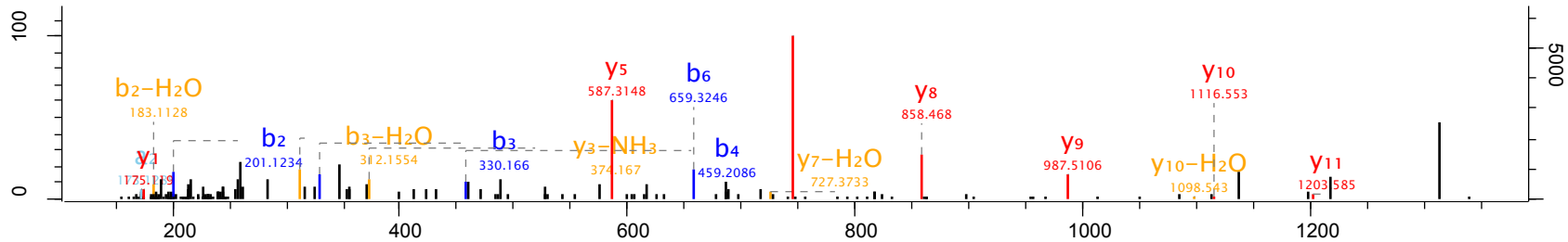
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	14124	TOF; CID	183.65	526.75	C14orf1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	15692	TOF; CID	90.7	592.8	MFSD1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	17790	TOF; CID	71.18	658.84	ZMYM6NB



Raw file

20141014_fract3_dyn_5ul_E3_01_582

Scan

21322

Method

TOF; CID

Score

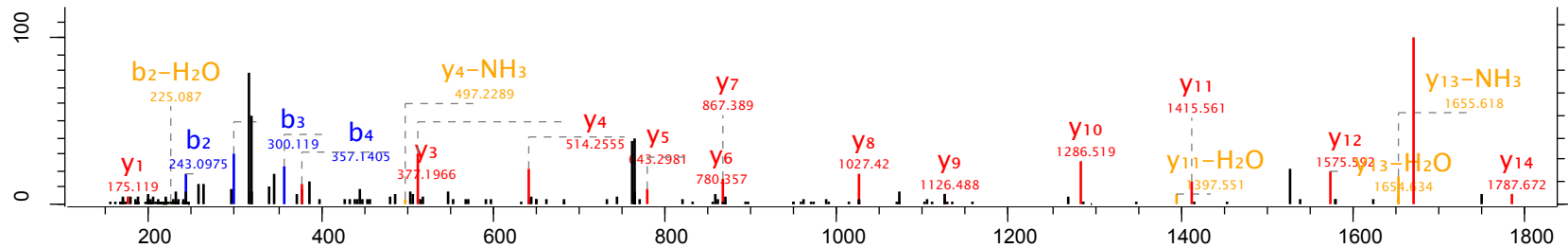
79.49

m/z

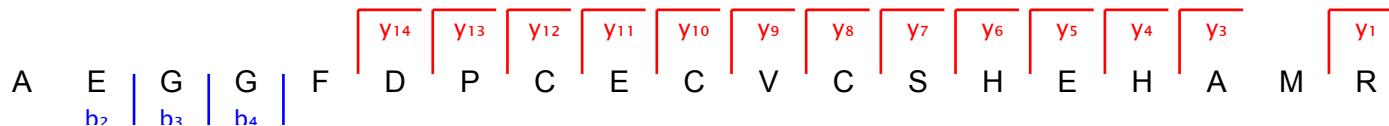
764.63

Gene names

SMIM14



ac



Raw file

20141014_fract3_dyn_5ul_E3_01_582

Scan

22524

Method

TOF; CID

Score

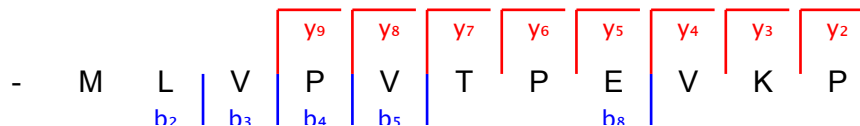
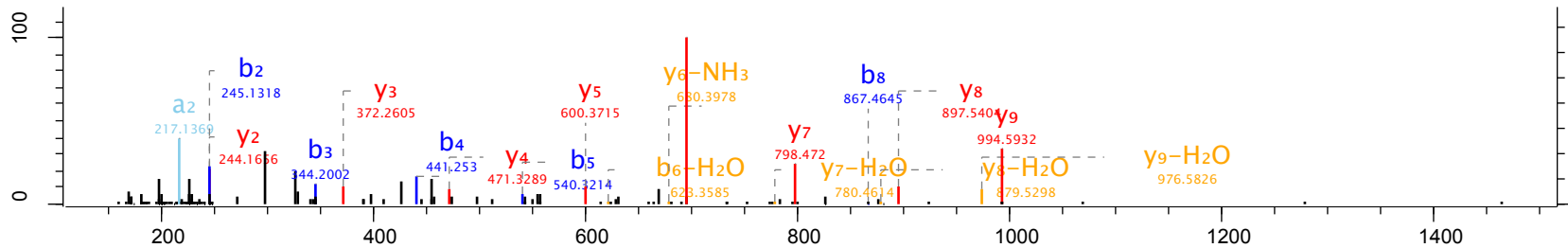
81.62

m/z

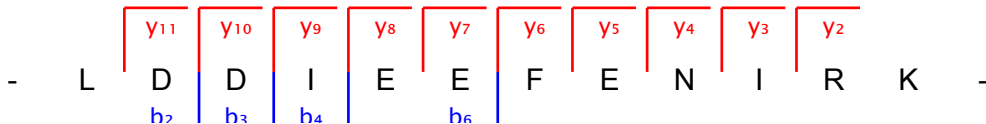
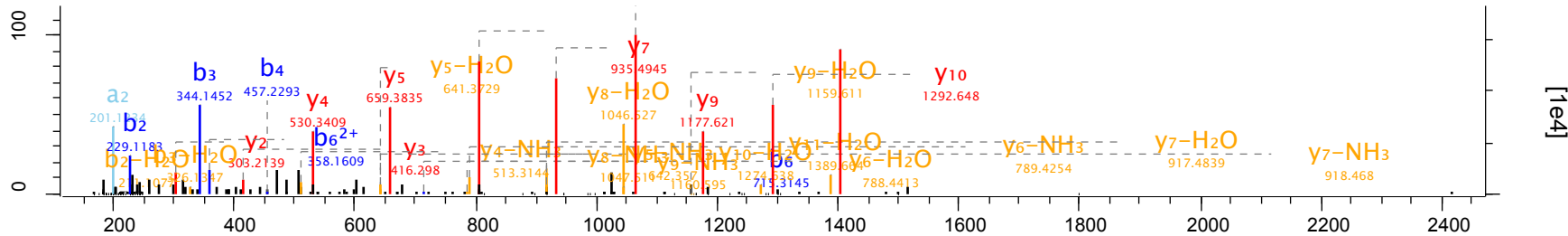
446.6

Gene names

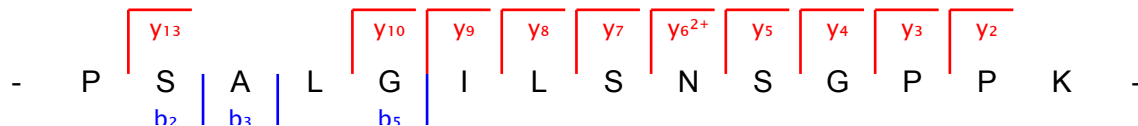
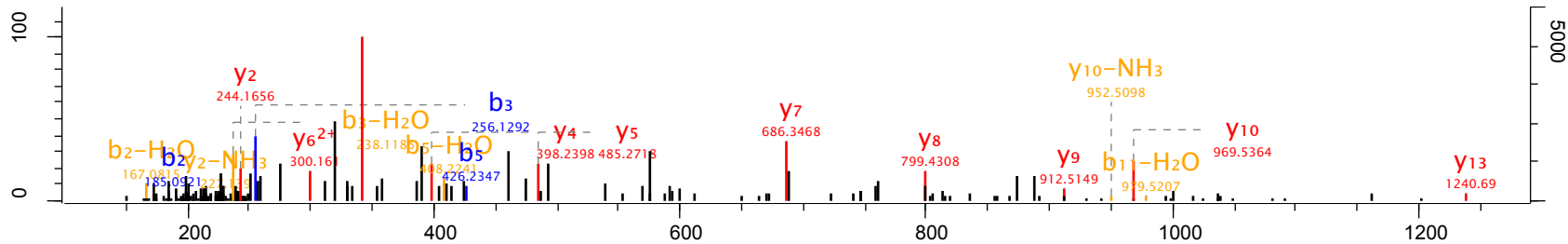
HMGN5



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	24106	TOF; CID	186.81	507.59	CDC26



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	25926	TOF; CID	51.64	669.37	GREB1



Raw file

20141014_fract3_dyn_5ul_E3_01_582

Scan

27220

Method

TOF; CID

Score

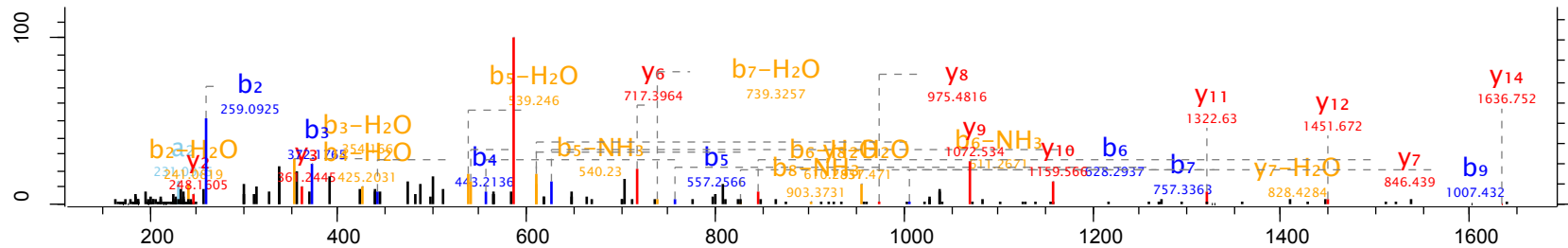
81.64

m/z

1040.48

Gene names

TCTEX1D2

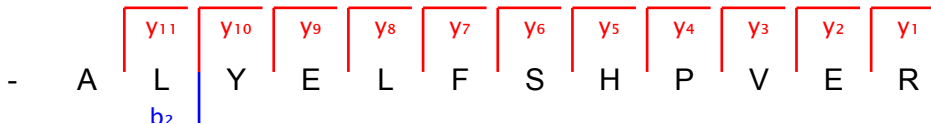
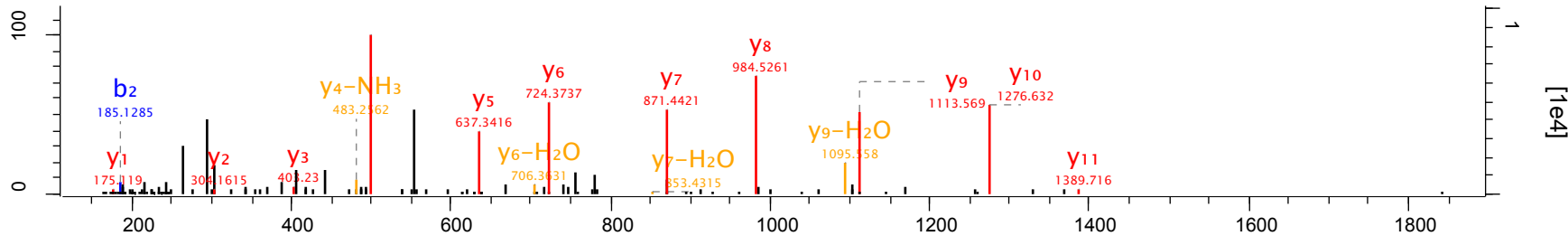


- E E L A N A E Y S P E E M P Q L T K -

b₂ | b₃ | b₄ | b₅ | b₆ | b₇ | b₉

y₁₄ | y₁₂ | y₁₁ | y₁₀ | y₉ | y₈ | y₇ | y₆ | y₅ | y₃ | y₂

Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	28058	TOF; CID	58.32	487.59	TMEM231



Raw file

20141014_fract3_dyn_5ul_E3_01_582

Scan

28977

Method

TOF; CID

Score

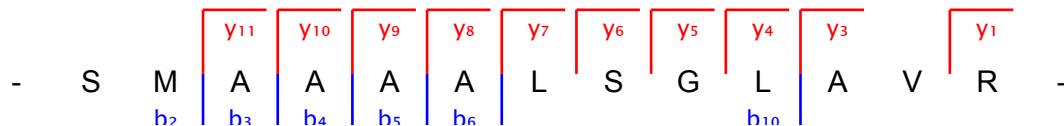
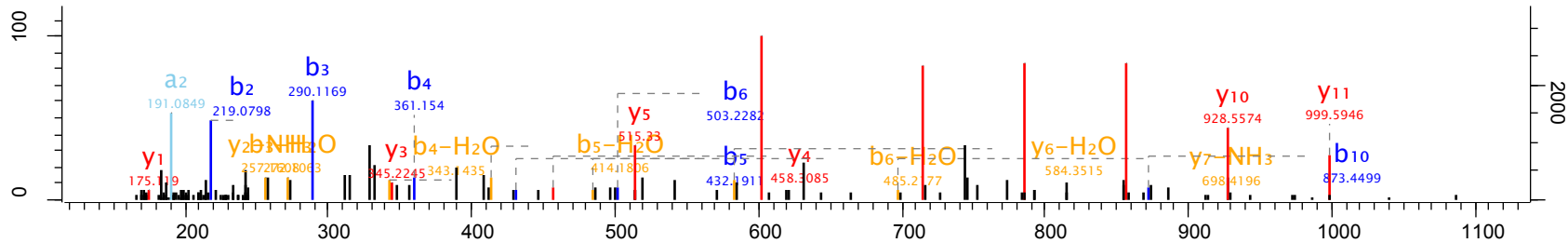
72.68

m/z

609.33

Gene names

SMIM10



Raw file

20141014_fract3_dyn_5ul_E3_01_582

Scan

30250

Method

TOF; CID

Score

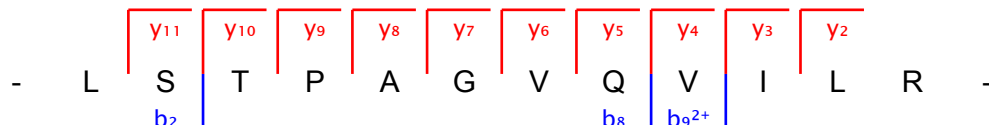
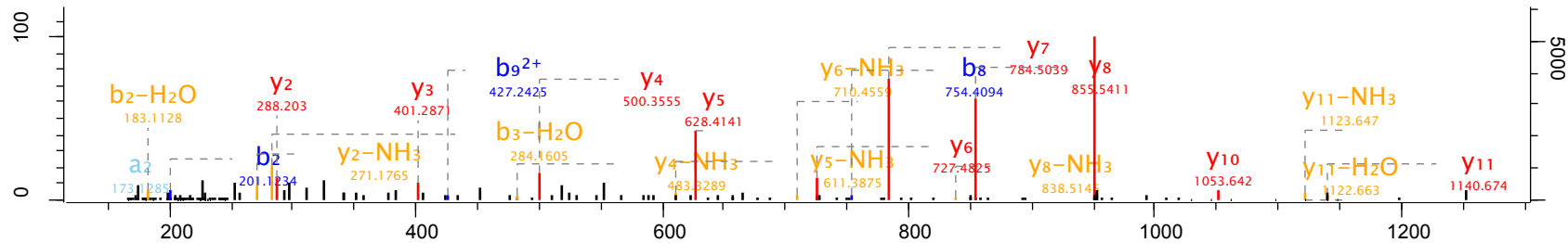
97.16

m/z

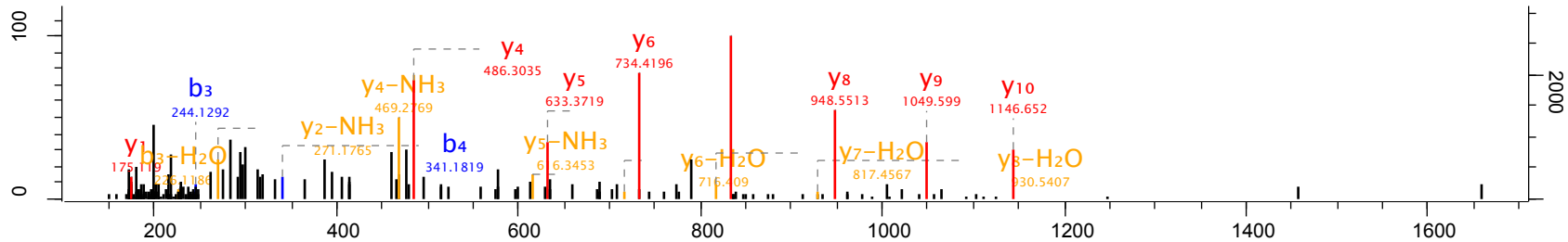
627.38

Gene names

MRPL34



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract3_dyn_5ul_E3_01_582	30675	TOF; CID	50.09	695.39	CCDC71L



- A A T P T L T T F P T I R -

b₃ (under T)
b₄ (under P)
y₁₀ (over P)
y₉ (over T)
y₈ (over L)
y₇ (over T)
y₆ (over T)
y₅ (over F)
y₄ (over P)
y₁ (over R)

Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract3_dyn_5ul_E3_01_582

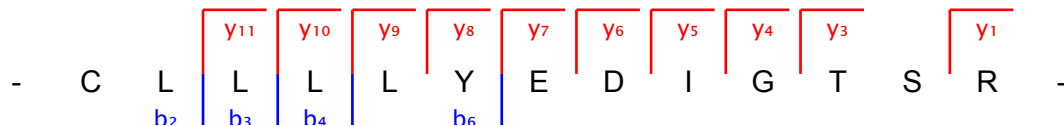
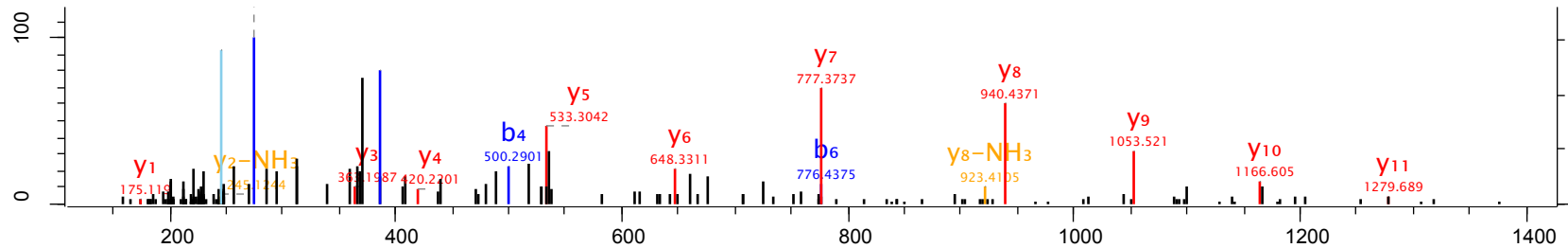
38024

TOF; CID

65.04

777.4

TPRA1



Raw file

20141014_fract3_dyn_5ul_E3_01_582

Scan

38068

Method

TOF; CID

Score

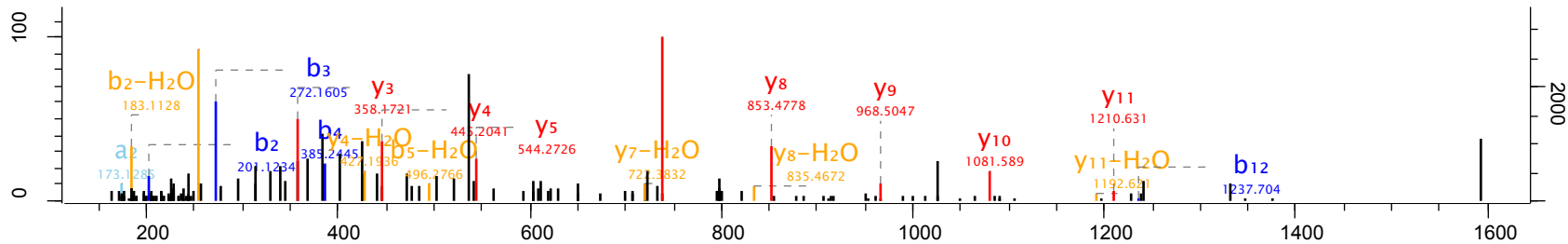
86.8

m/z

798.44

Gene names

CLDN12



- L S A I E I D I P V V S H T T -
 b₂ | b₃ | b₄ | y₁₁ | y₁₀ | y₉ | y₈ | y₇ | y₅ | y₄ | y₃
 b₁₂

Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

3789

Method

TOF; CID

Score

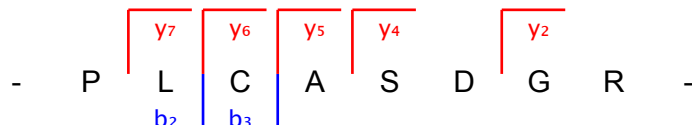
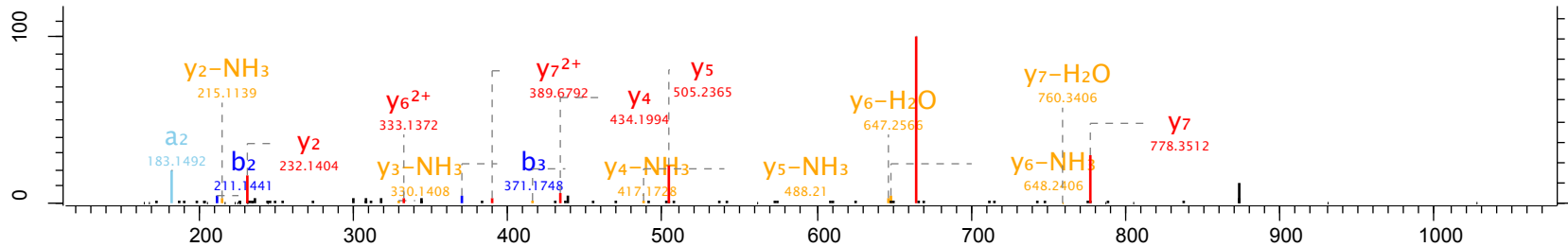
111.01

m/z

438.2

Gene names

SMOC2



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

4789

Method

TOF; CID

Score

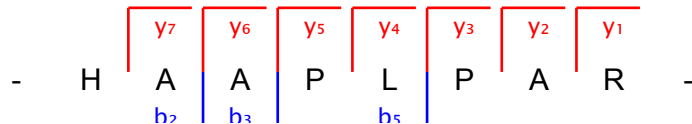
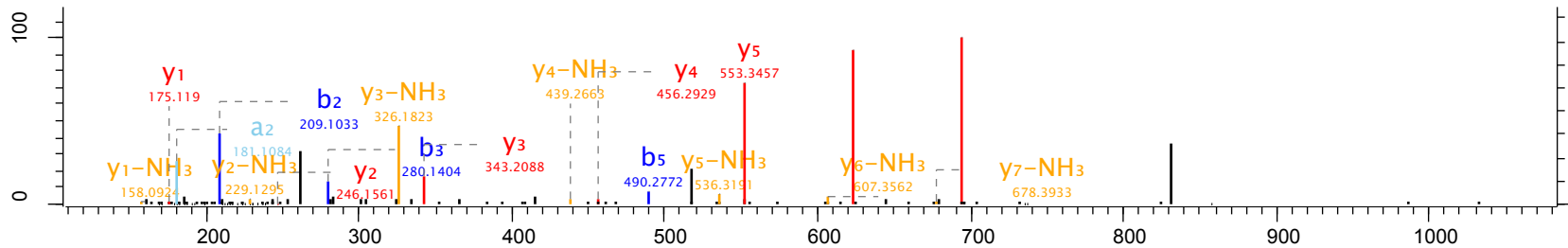
132.09

m/z

416.74

Gene names

TP53INP2



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

8664

Method

TOF; CID

Score

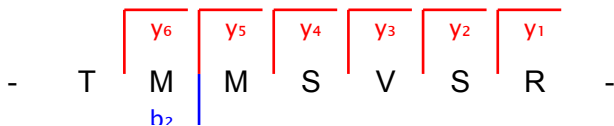
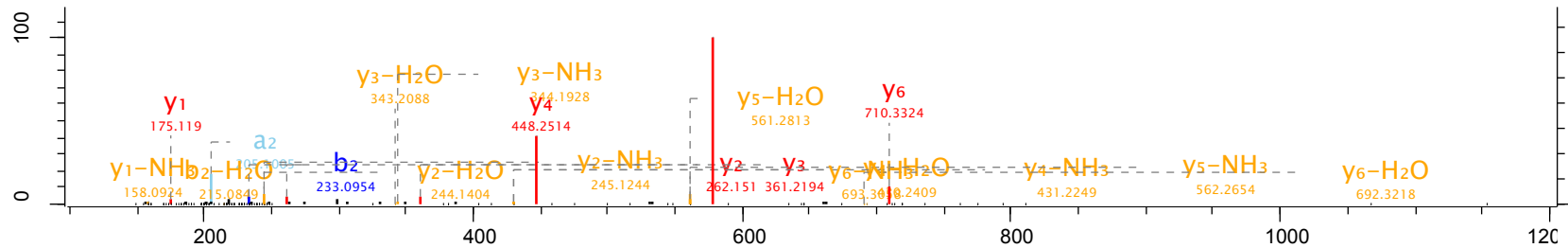
130.77

m/z

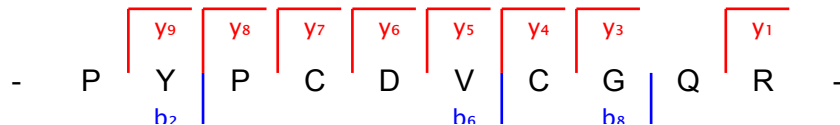
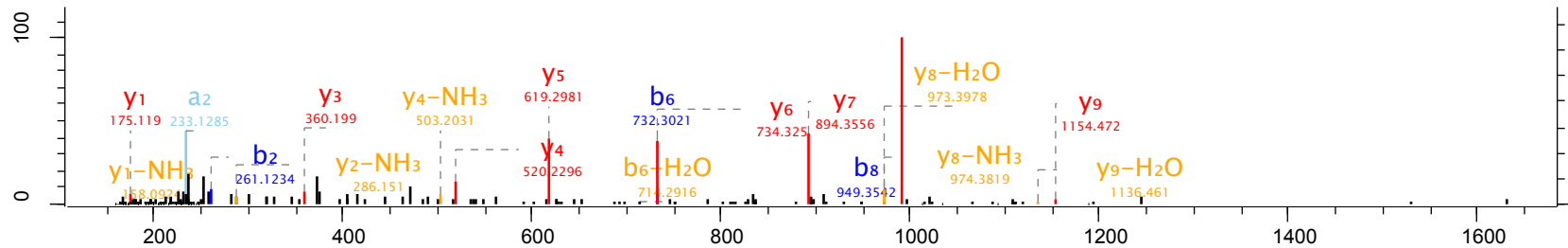
406.19

Gene names

CID



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract4_dyn_5ul_E4_01_583	9921	TOF; CID	85.35	626.26	ZNF652;ZBTB47



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

17515

Method

TOF; CID

Score

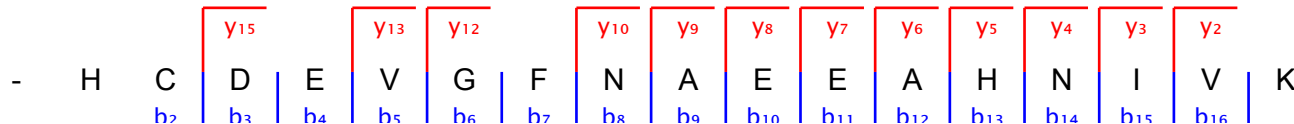
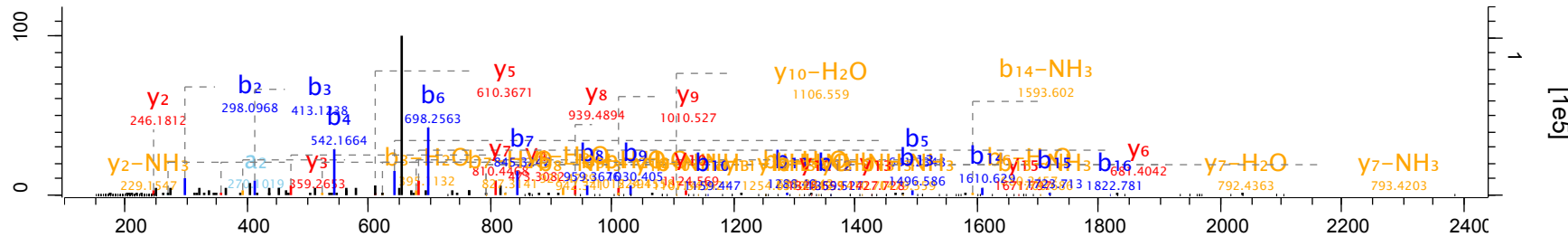
160.81

m/z

657.3

Gene names

DYNLT3



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

19196

Method

TOF; CID

Score

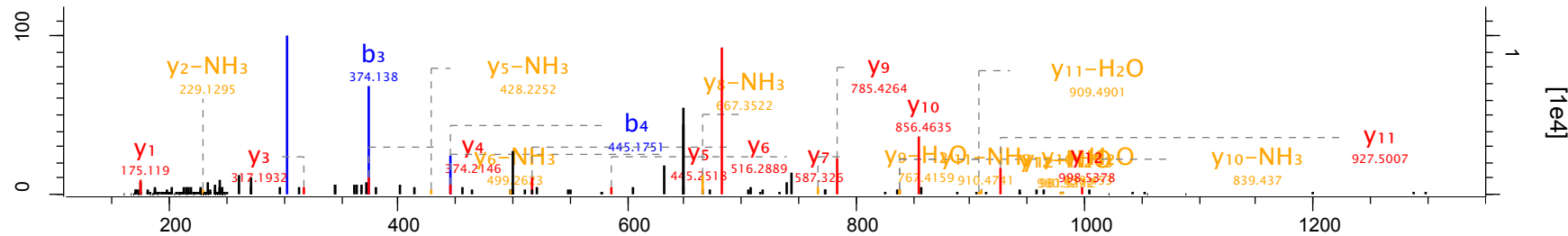
91.28

m/z

650.82

Gene names

SLC36A4



ac

M E A A A T P A A A G A A R -

b2

b3

b4

Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

20269

Method

TOF; CID

Score

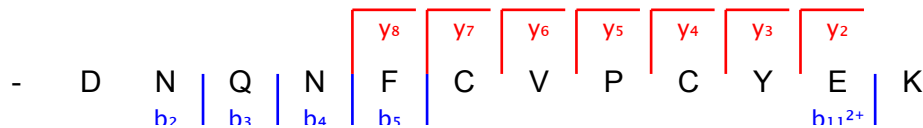
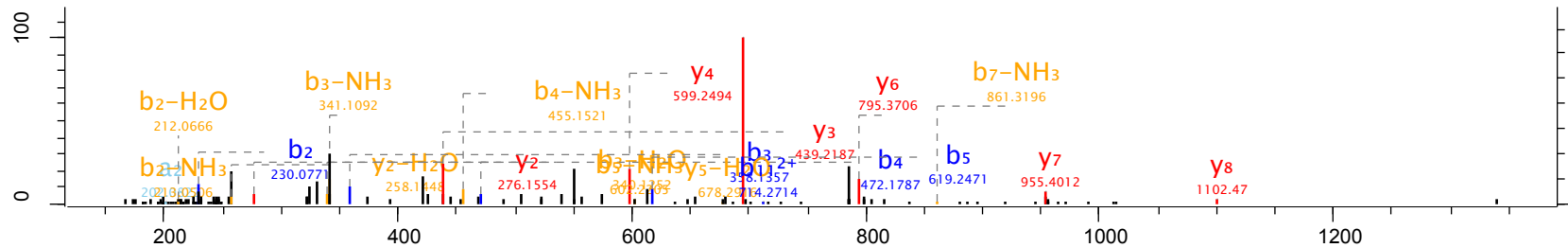
57.43

m/z

787.32

Gene names

FHL2



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

20370

Method

TOF; CID

Score

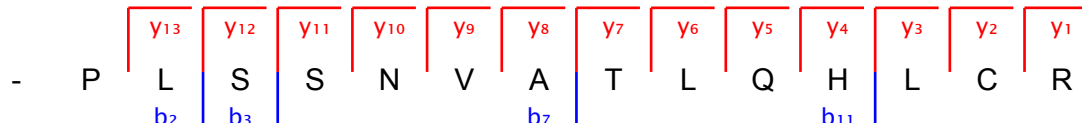
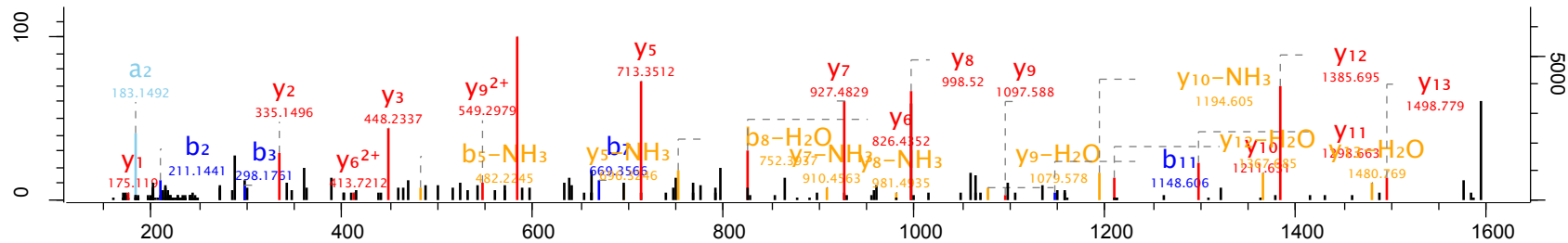
109.83

m/z

532.61

Gene names

SOCS3



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

20560

Method

TOF; CID

Score

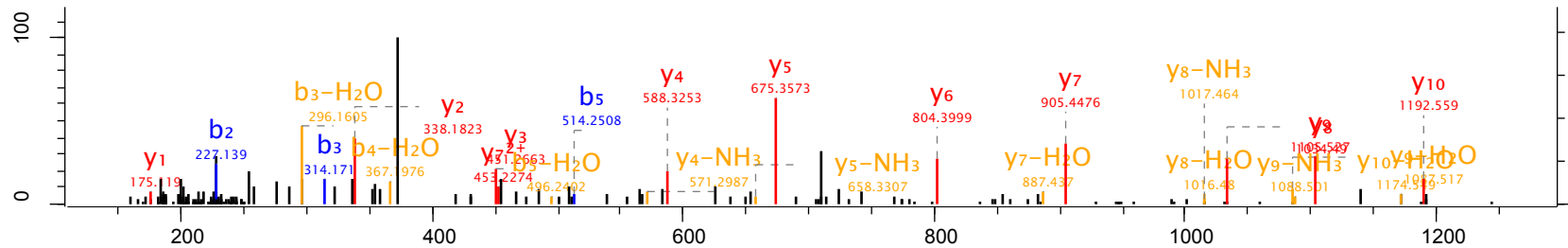
111.52

m/z

709.85

Gene names

CYB561D2



ac

- A L S A E T E S H I Y R -

 b₂ b₃ b₅

 y₁₀ y₉ y₈ y₇ y₆ y₅ y₄ y₃ y₂ y₁

Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

24861

Method

TOF; CID

Score

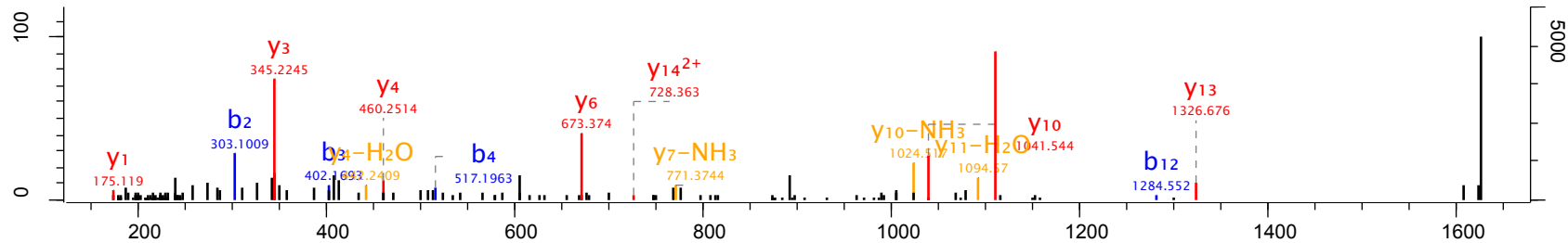
64.24

m/z

814.89

Gene names

C4orf3



ac

M E V D A P G V D G R D G L R -
 b₂ b₃ b₄ b₁₂

y₁₄²⁺ y₁₃ y₁₁ y₁₀ y₆ y₄ y₃ y₁

Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

25014

Method

TOF; CID

Score

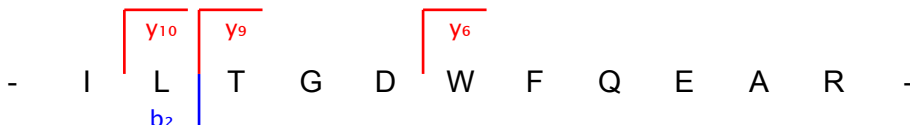
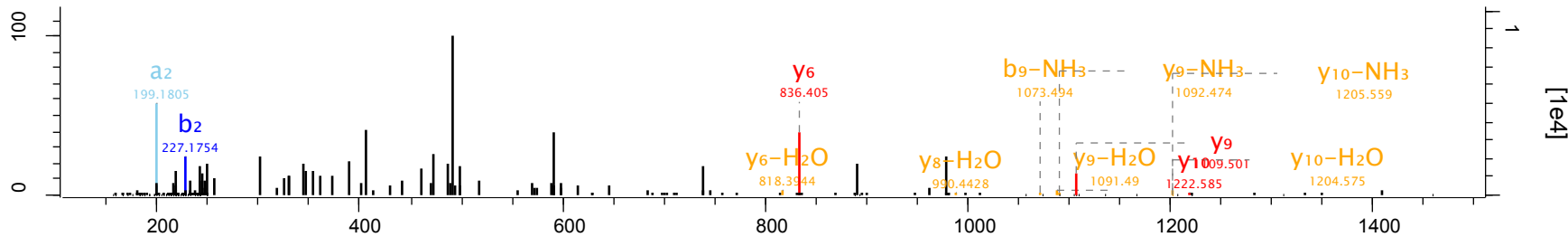
59.91

m/z

668.35

Gene names

SYTL1



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

25540

Method

TOF; CID

Score

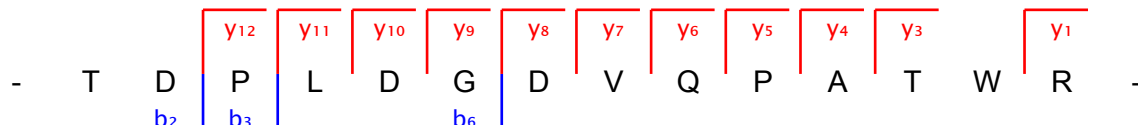
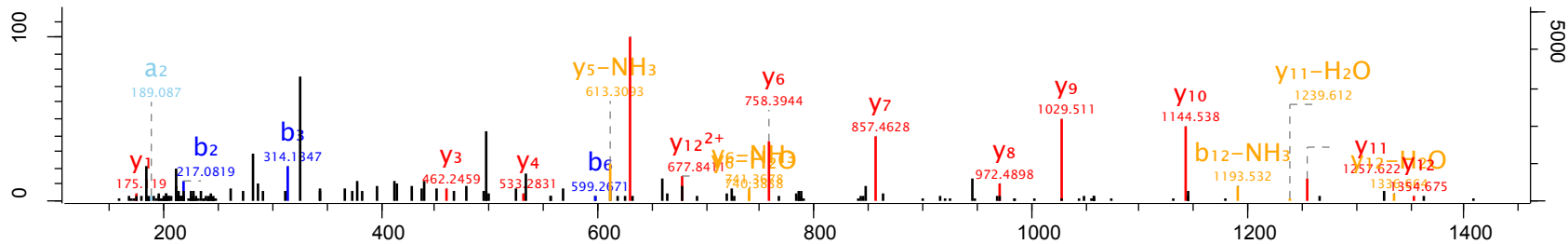
65.37

m/z

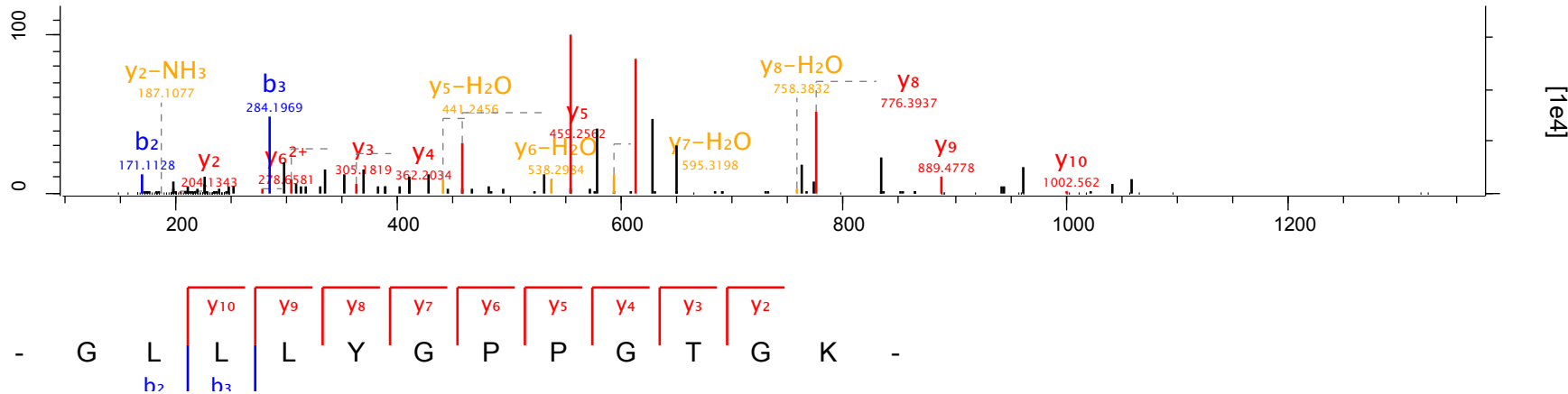
785.87

Gene names

HGSNAT



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract4_dyn_5ul_E4_01_583	26021	TOF; CID	60.16	586.84	KATNAL2



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

26163

Method

TOF; CID

Score

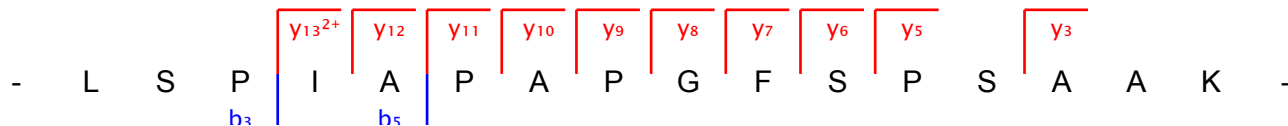
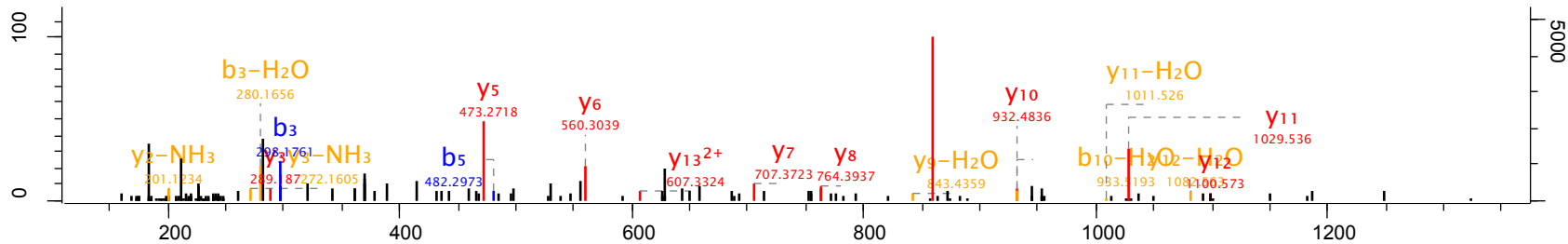
57.97

m/z

755.91

Gene names

KLF10



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

26226

Method

TOF; CID

Score

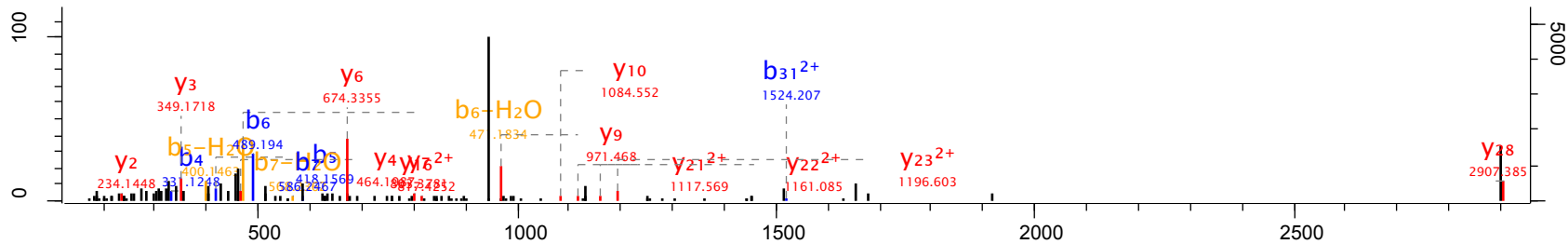
27.53

m/z

1132.86

Gene names

TECPR1



- G S G E | S | A | P | S D T D A S S E V E R P G P

h₄ | h₅ | h₆ | h₇

y₂₈

y₂₃²⁺ | y₂₂²⁺ | y₂₁²⁺

y₁₆²⁺

Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

26734

Method

TOF; CID

Score

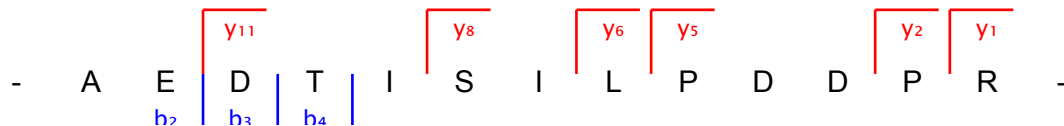
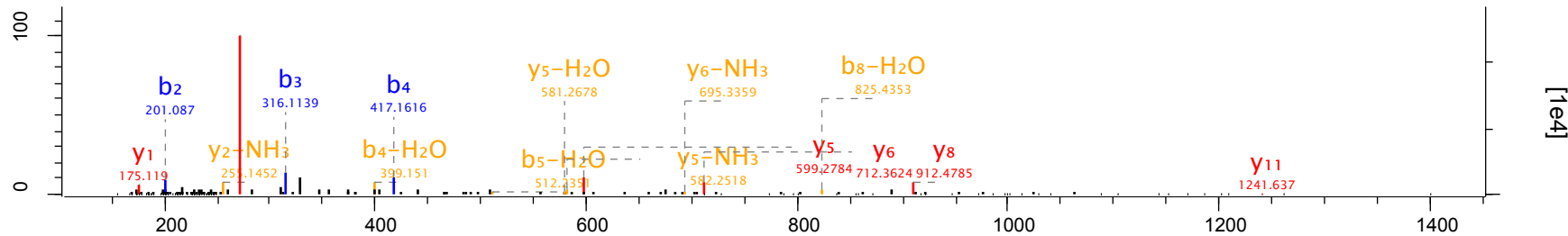
78.32

m/z

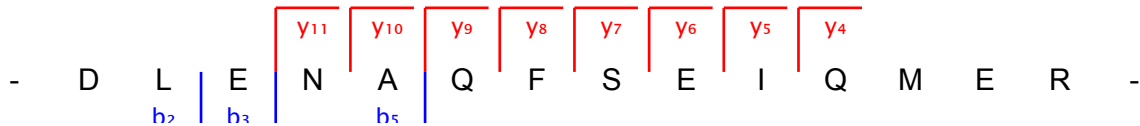
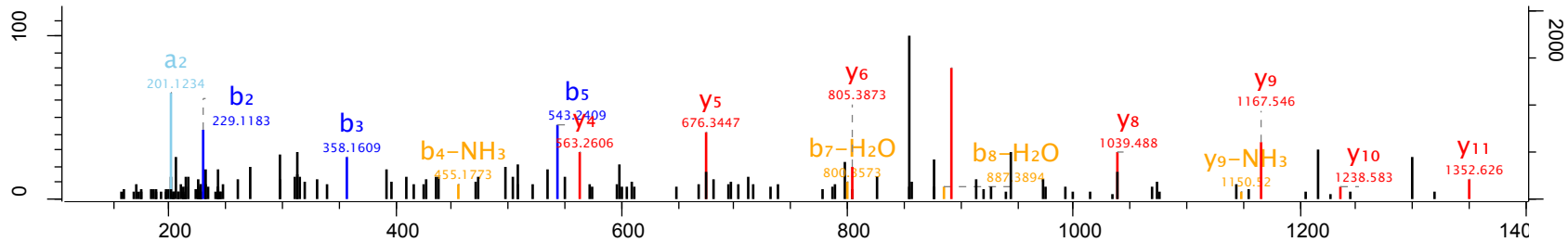
721.36

Gene names

CHURC1-FNTB;CHURC1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract4_dyn_5ul_E4_01_583	26770	TOF; CID	52.53	855.39	METTL7B



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

27431

Method

TOF; CID

Score

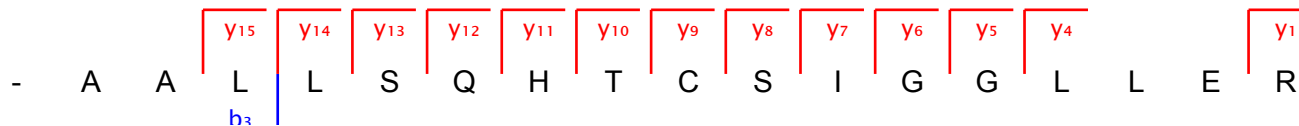
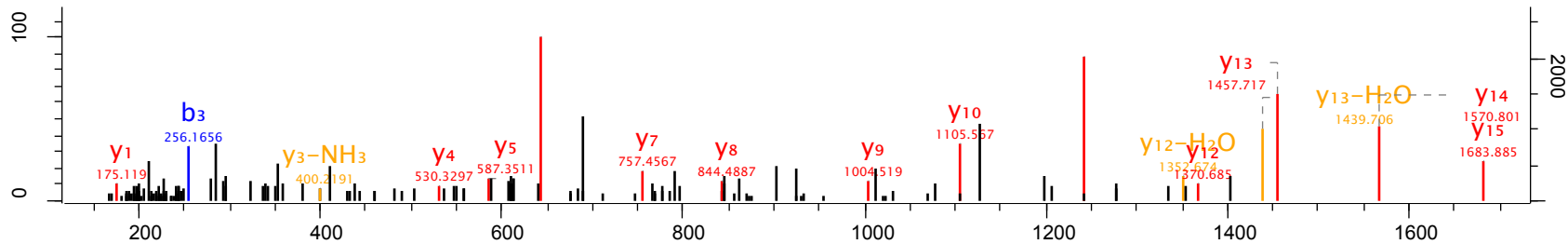
51.09

m/z

609.32

Gene names

SETD6



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract4_dyn_5ul_E4_01_583

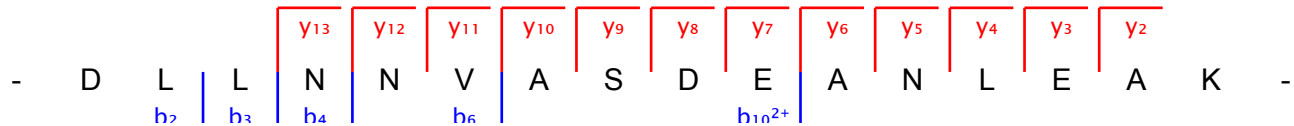
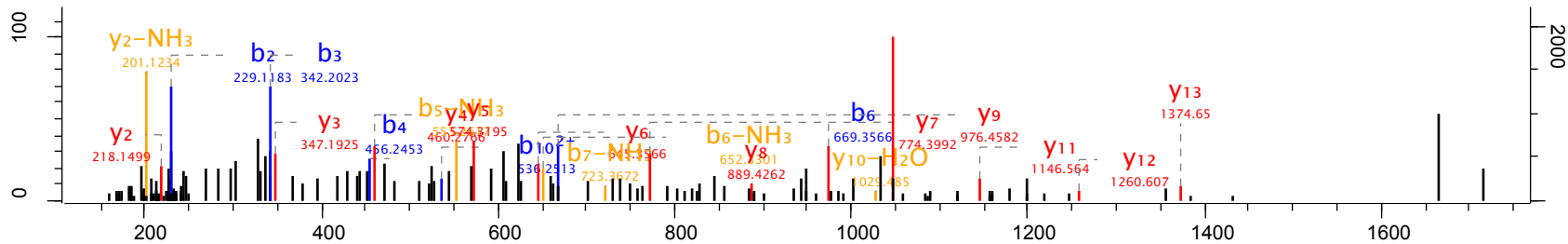
28844

TOF; CID

87.43

858.42

CLUAP1



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract4_dyn_5ul_E4_01_583

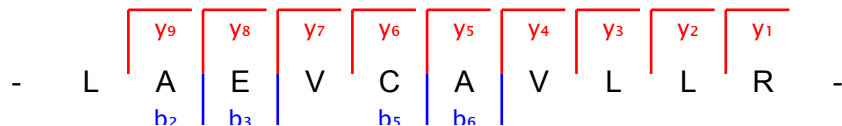
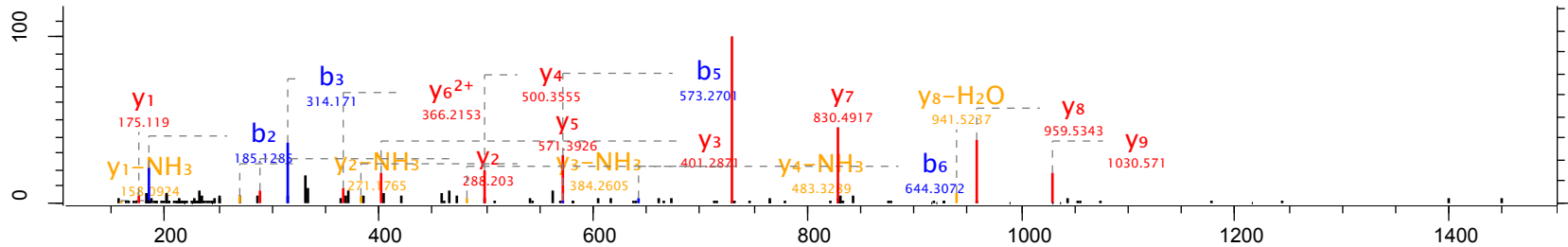
29058

TOF; CID

86.18

572.33

BOK



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

29096

Method

TOF; CID

Score

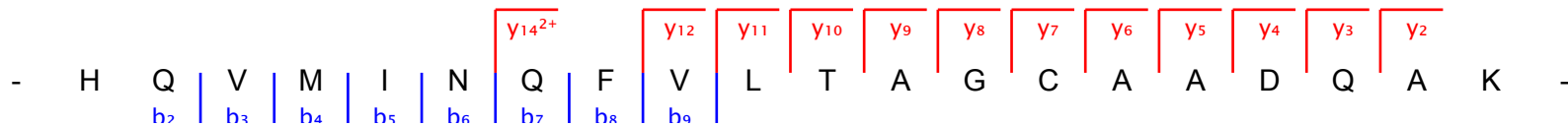
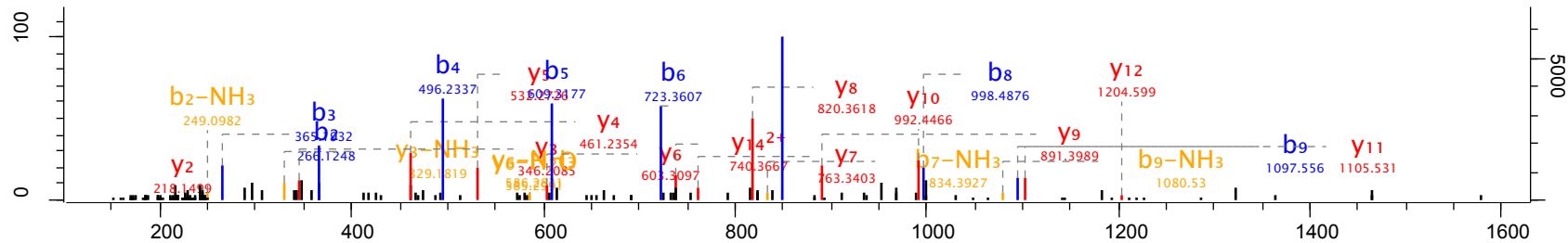
99.09

m/z

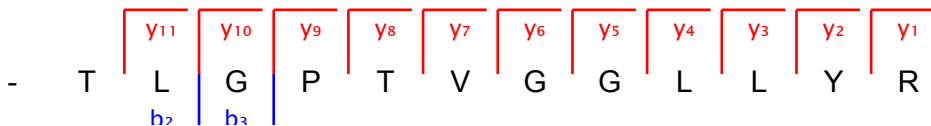
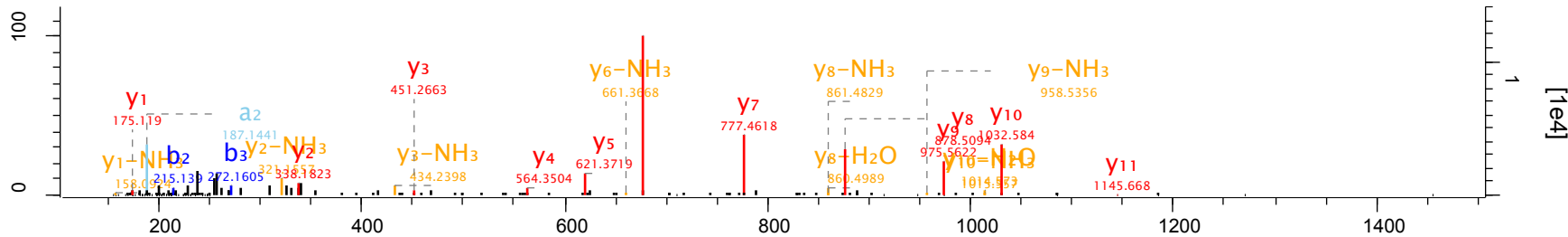
735.03

Gene names

UBALD1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract4_dyn_5ul_E4_01_583	29929	TOF; CID	103.43	623.86	SLC22A18



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

30589

Method

TOF; CID

Score

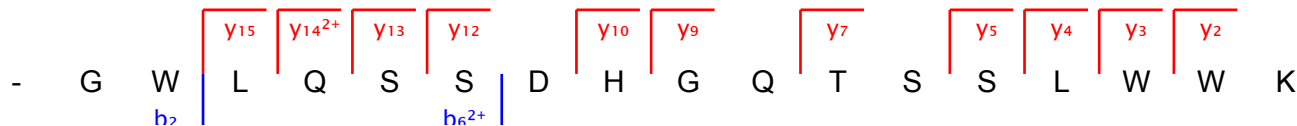
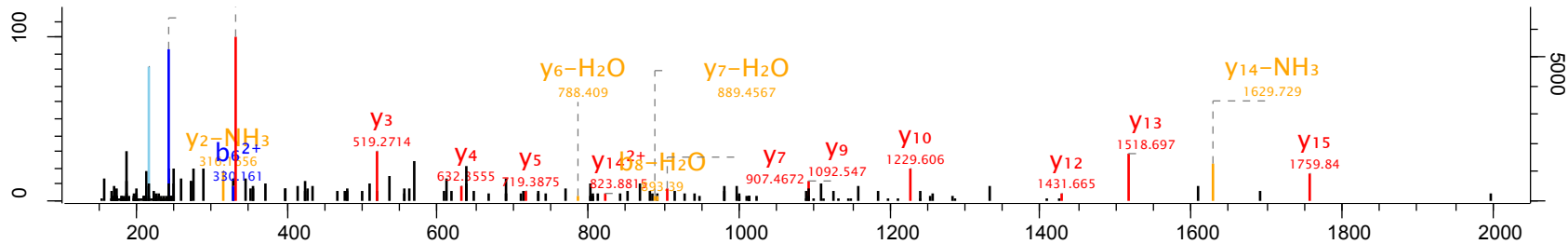
53.56

m/z

668.31

Gene names

PERP



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

32830

Method

TOF; CID

Score

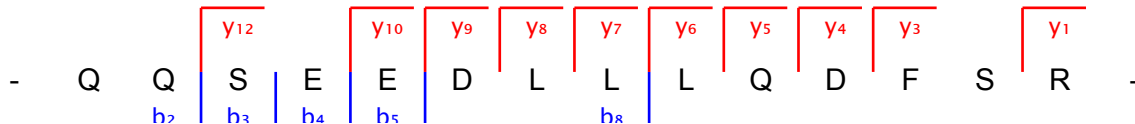
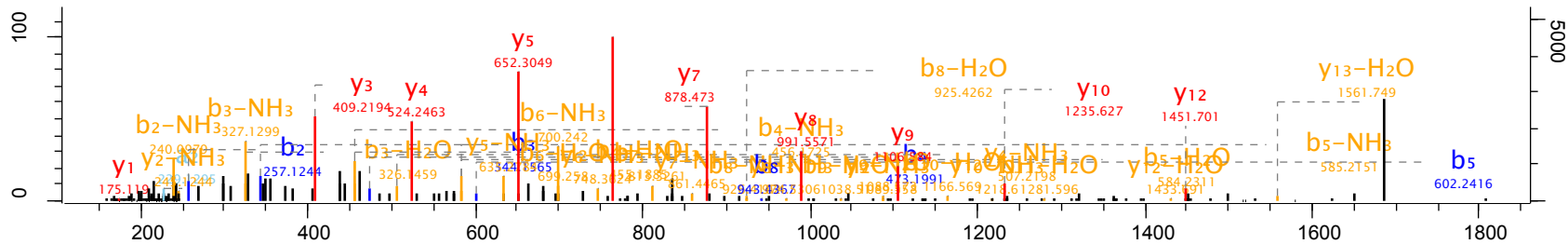
146.37

m/z

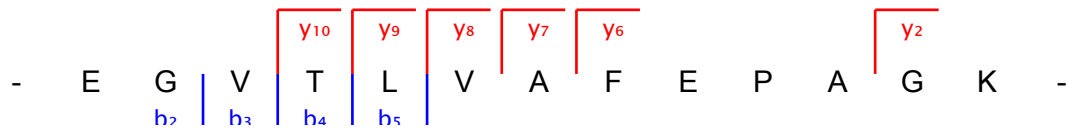
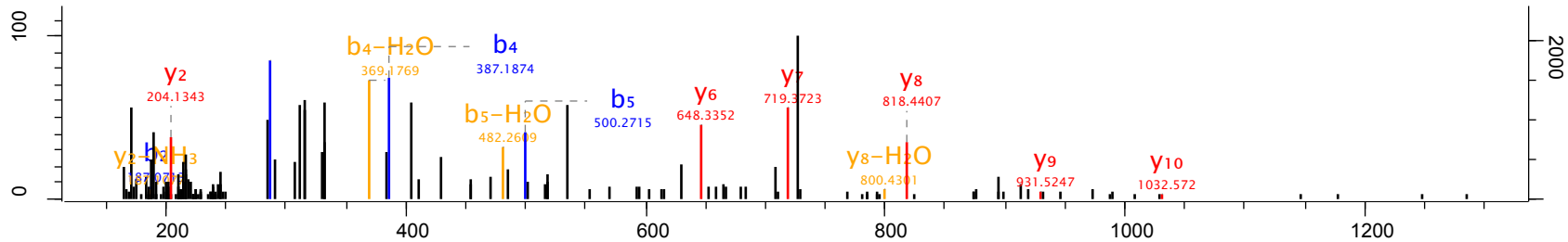
854.41

Gene names

SSR3



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract4_dyn_5ul_E4_01_583	33651	TOF; CID	51.73	659.36	CDC14B



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

34503

Method

TOF; CID

Score

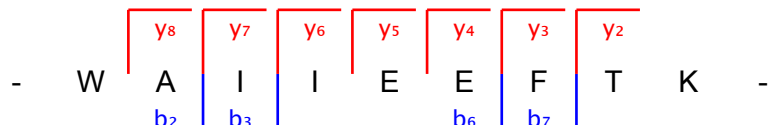
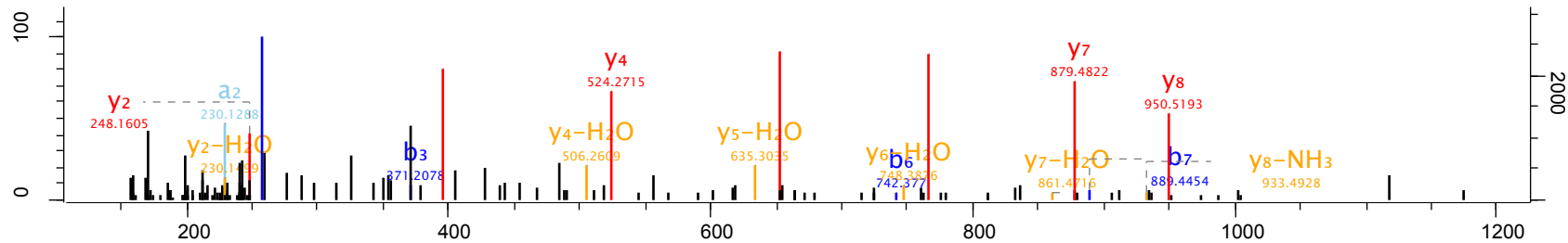
77.66

m/z

568.8

Gene names

MT-ND2



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

38279

Method

TOF; CID

Score

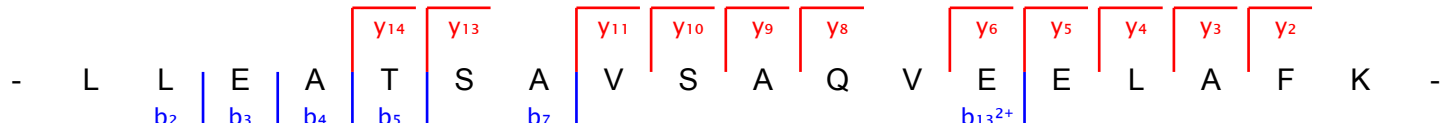
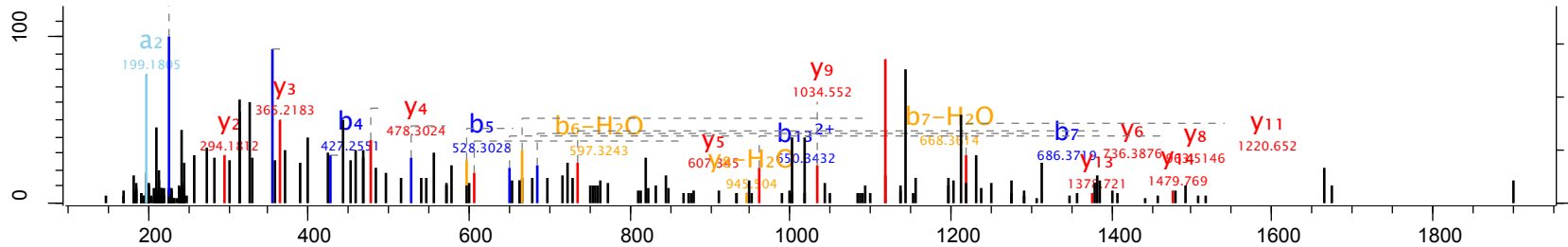
59.25

m/z

953.5

Gene names

C4orf46



Raw file

20141014_fract4_dyn_5ul_E4_01_583

Scan

38846

Method

TOF; CID

Score

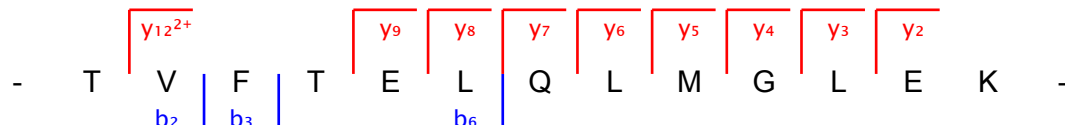
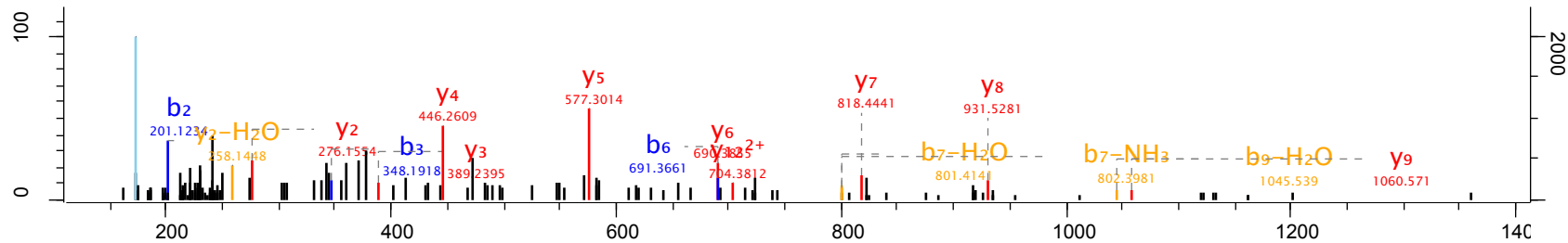
73.93

m/z

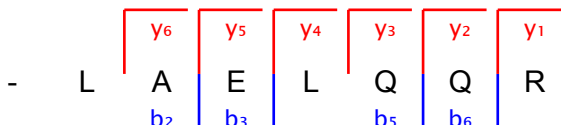
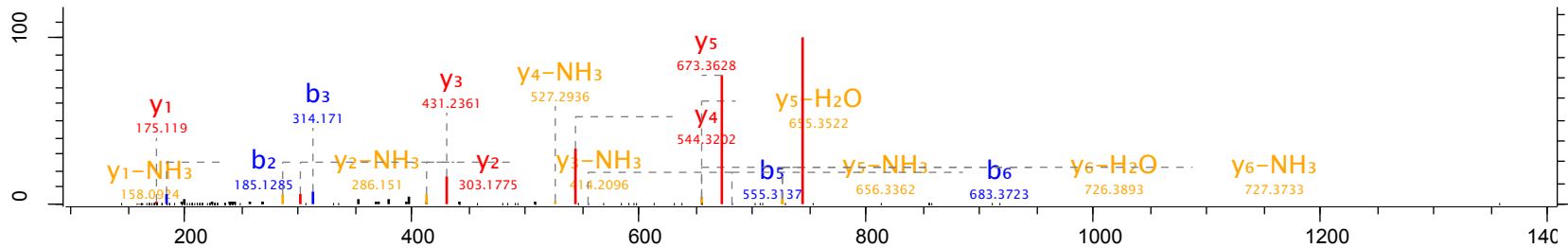
754.9

Gene names

BARX1

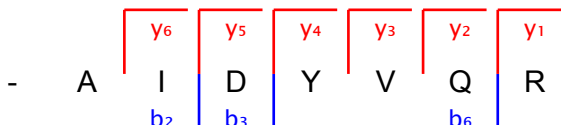
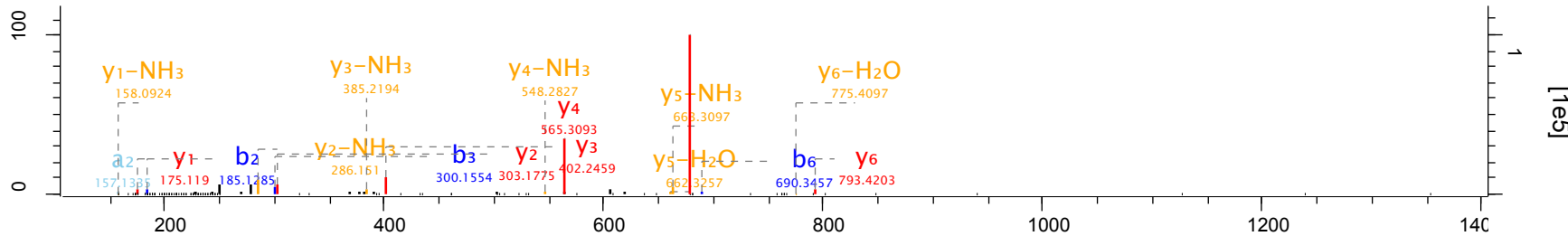


Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	7476	TOF; CID	139.28	429.24	VAMP5



[1e4]

Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	9936	TOF; CID	134.66	432.73	TSPAN3



Raw file

20141014_fract5_dyn_5ul_E5_01_584

Scan

12423

Method

TOF; CID

Score

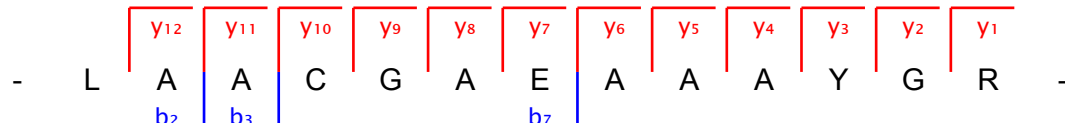
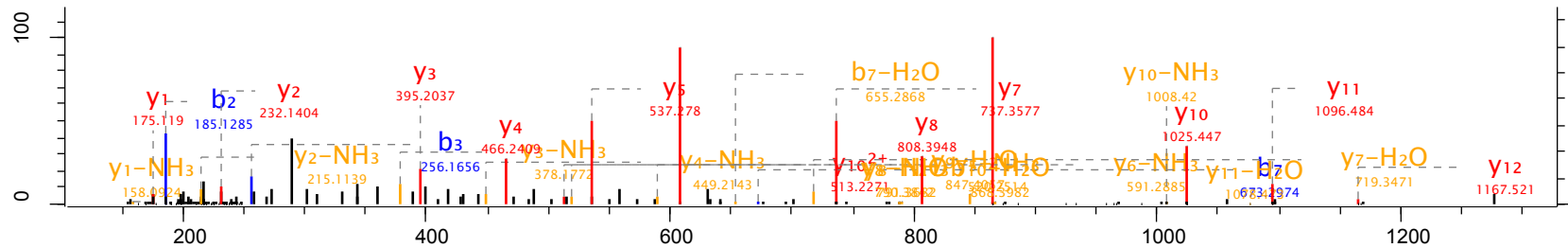
135.1

m/z

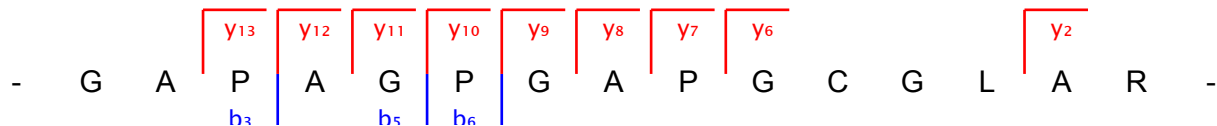
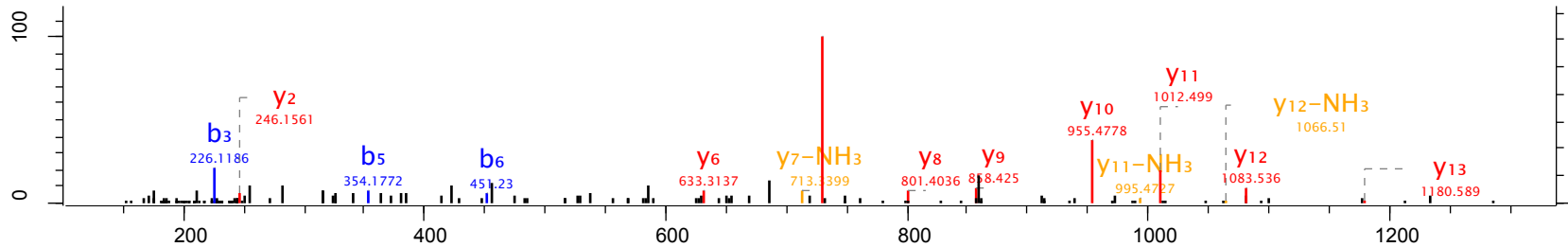
640.8

Gene names

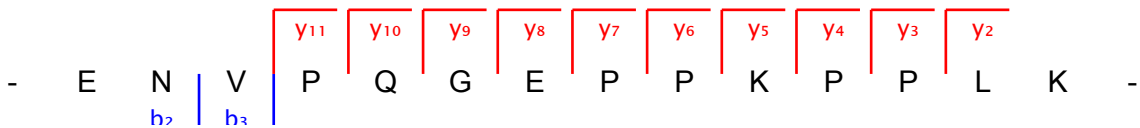
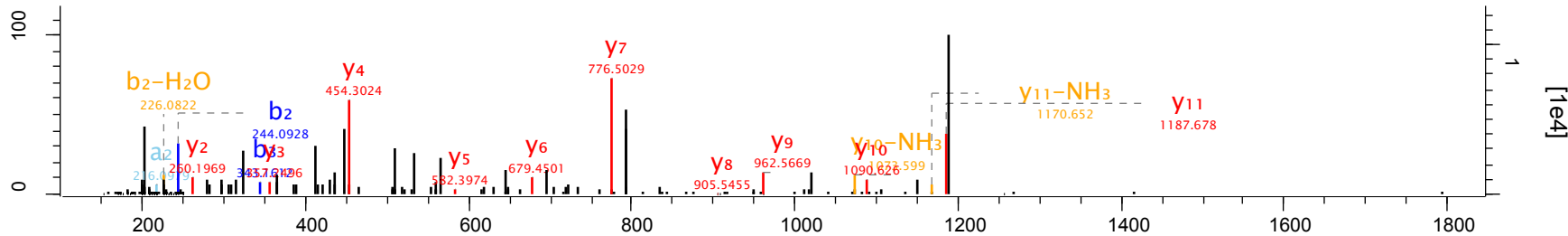
C17orf89



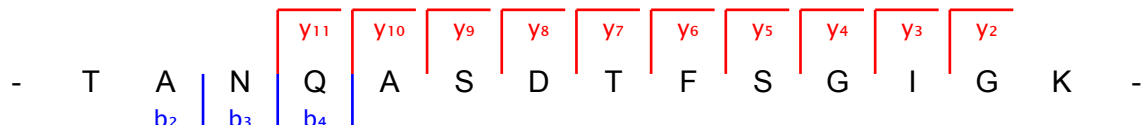
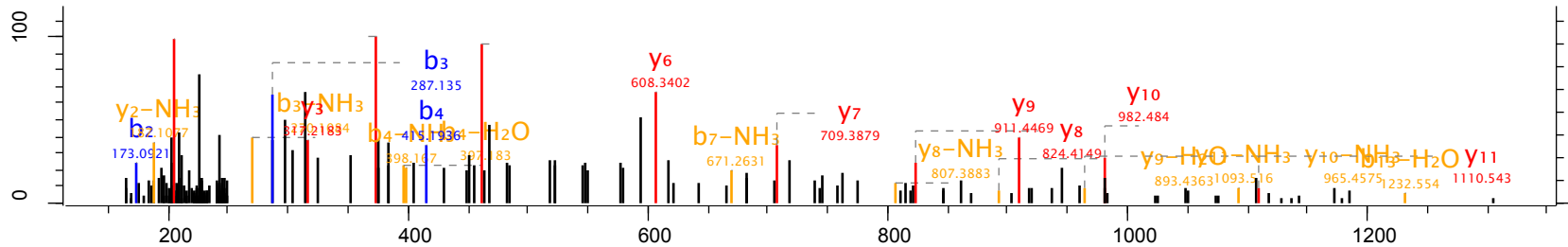
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	12753	TOF; CID	66.99	654.82	MORC4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	13734	TOF; CID	48.61	511.28	IFI27L2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	15192	TOF; CID	63.29	698.84	ADIRF



Raw file

20141014_fract5_dyn_5ul_E5_01_584

Scan

21272

Method

TOF; CID

Score

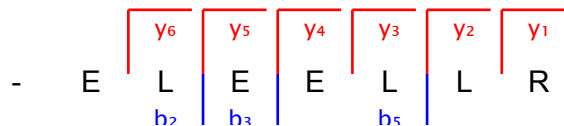
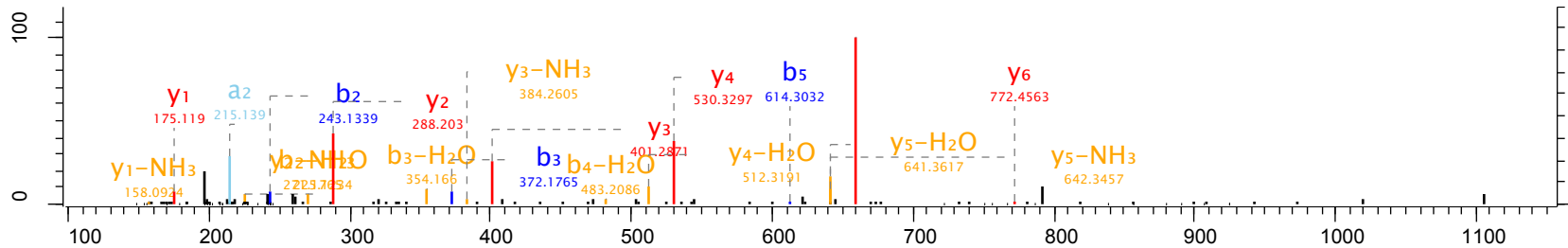
143

m/z

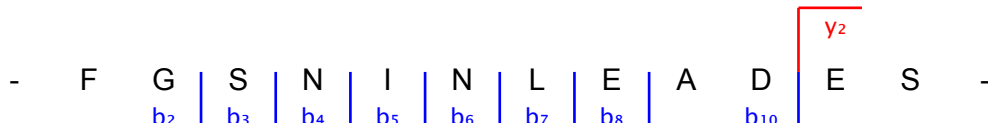
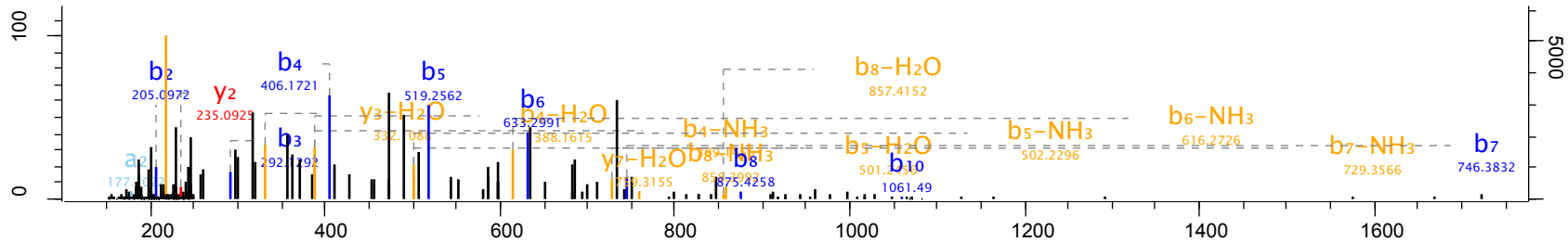
451.25

Gene names

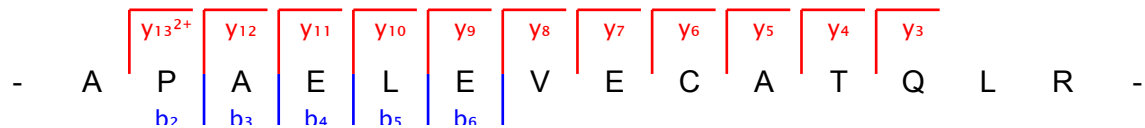
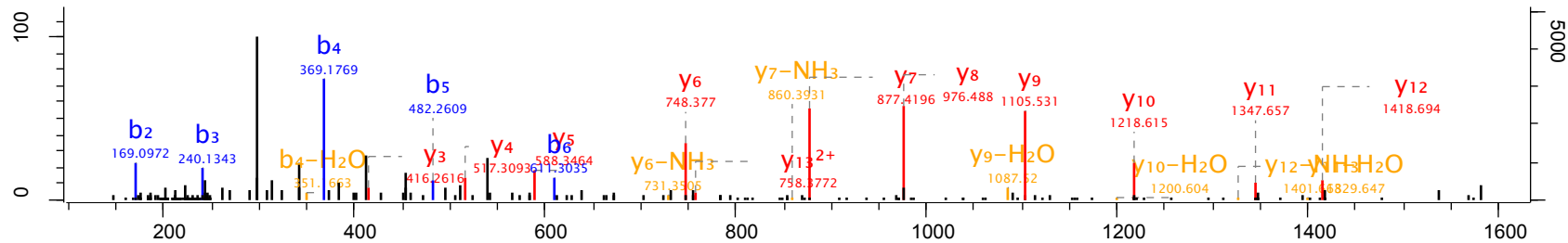
BTBD16



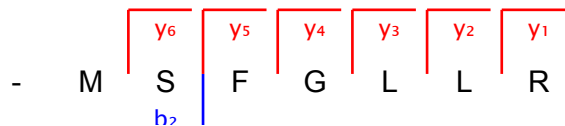
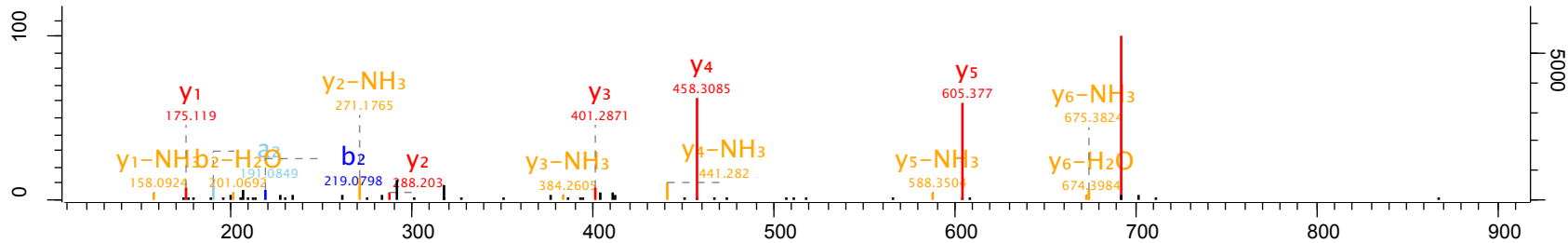
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	23325	TOF; CID	72.34	648.29	PFDN4



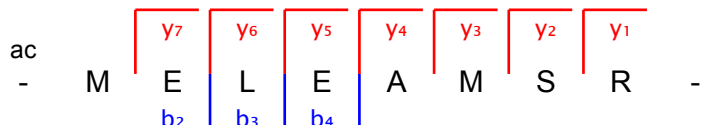
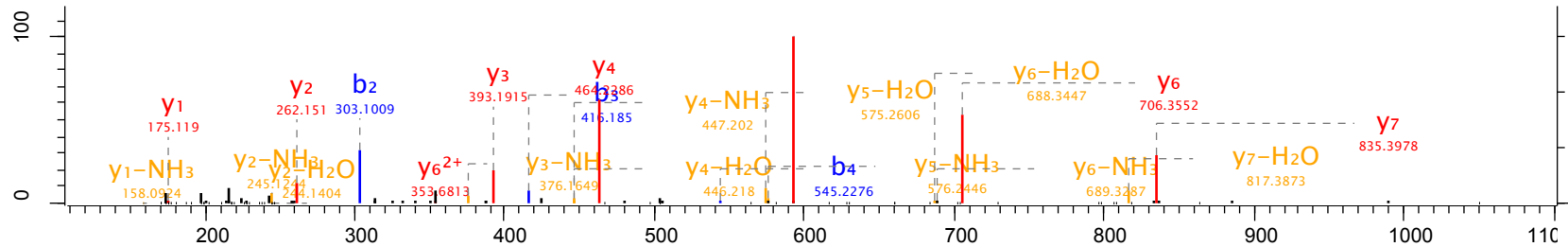
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	23615	TOF; CID	97.27	793.89	PMAIP1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	27722	TOF; CID	148.28	412.22	SMDT1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	28067	TOF; CID	143.28	504.72	TMEM258



Raw file

20141014_fract5_dyn_5ul_E5_01_584

Scan

30004

Method

TOF; CID

Score

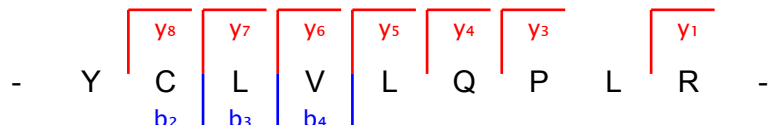
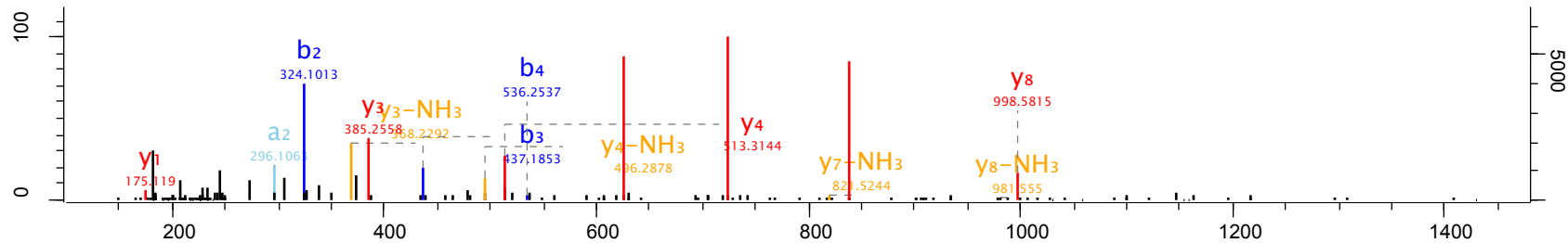
106.04

m/z

581.32

Gene names

ZDHHC12



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract5_dyn_5ul_E5_01_584

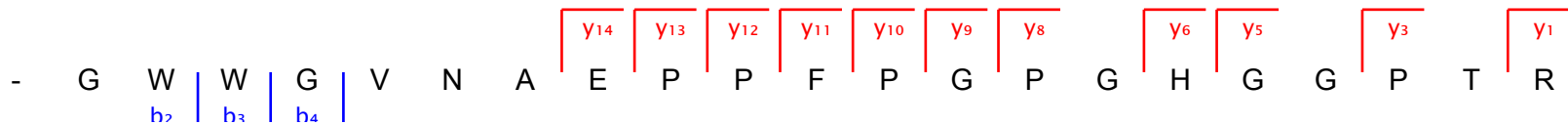
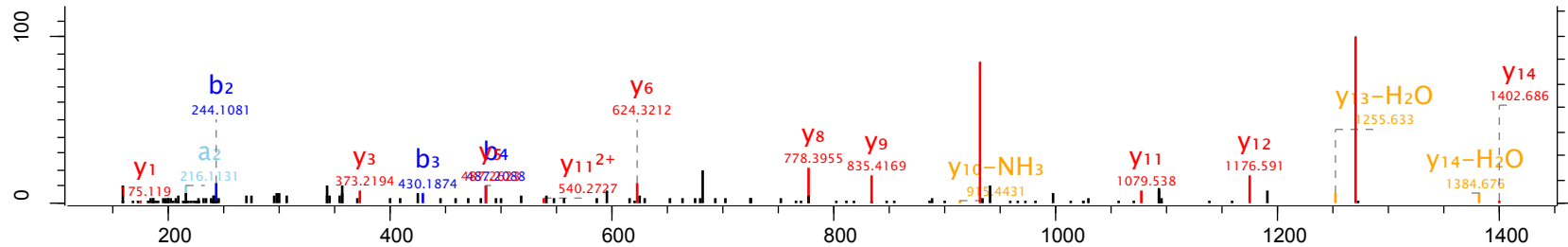
31391

TOF; CID

61.94

725.35

PRR3



Raw file

20141014_fract5_dyn_5ul_E5_01_584

Scan

32361

Method

TOF; CID

Score

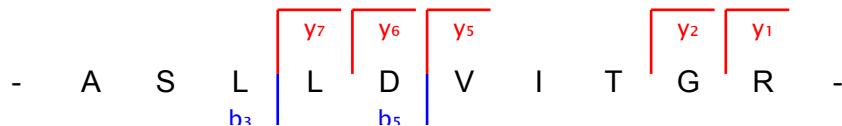
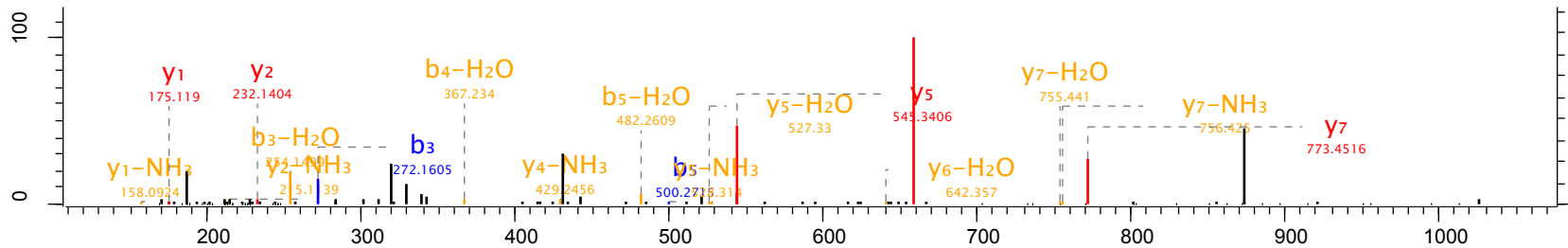
72.33

m/z

522.8

Gene names

ABCG8



Raw file

20141014_fract5_dyn_5ul_E5_01_584

Scan

33982

Method

TOF; CID

Score

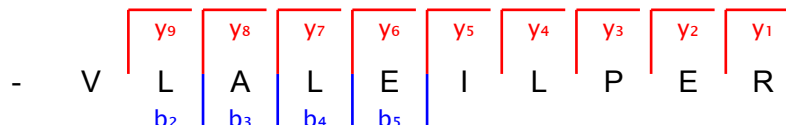
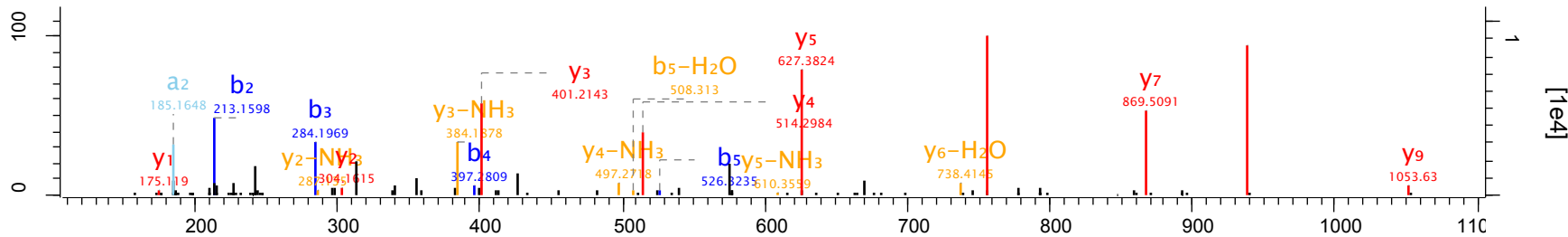
90.05

m/z

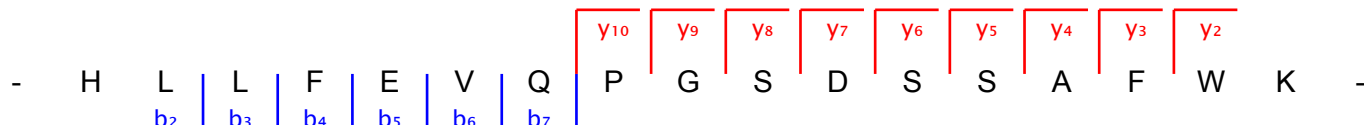
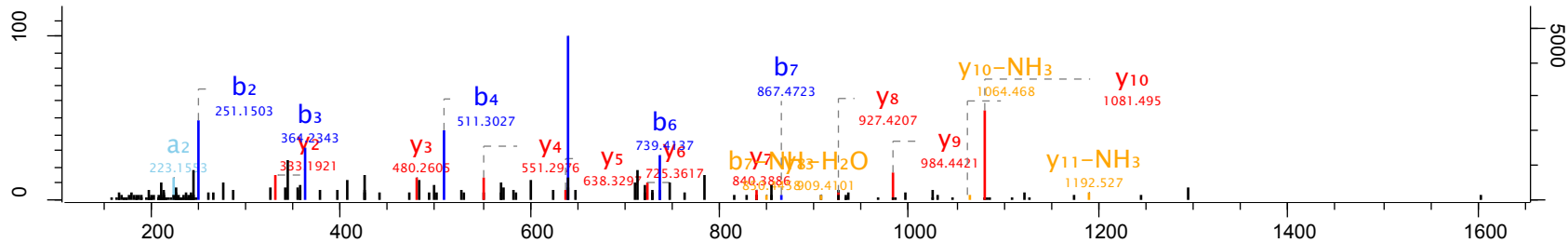
577.35

Gene names

SPIN4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	34491	TOF; CID	59.73	650.32	RNF141



Raw file

20141014_fract5_dyn_5ul_E5_01_584

Scan

34718

Method

TOF; CID

Score

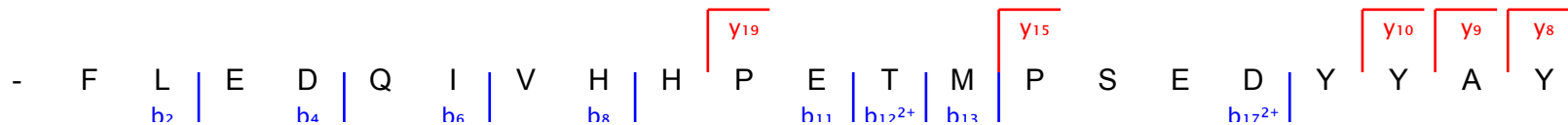
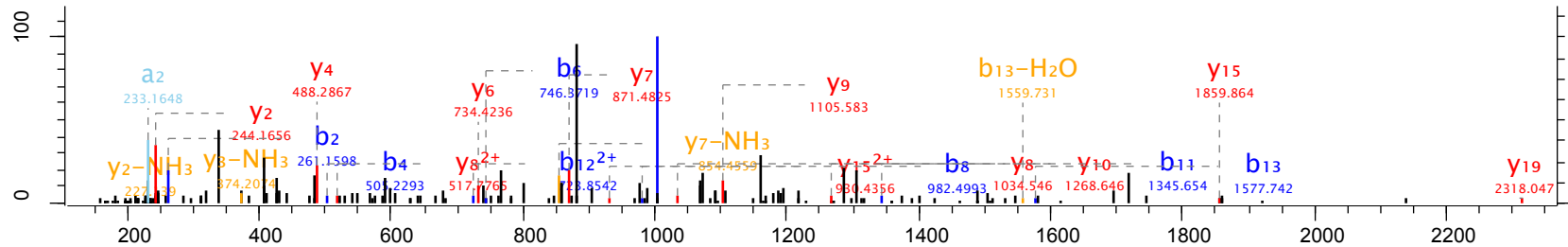
35.54

m/z

860.41

Gene names

ADPRM



Raw file

20141014_fract5_dyn_5ul_E5_01_584

Scan

36099

Method

TOF; CID

Score

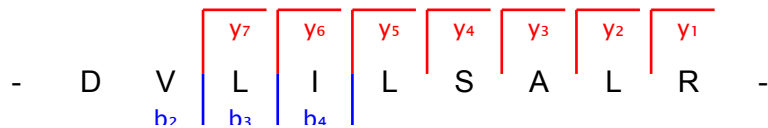
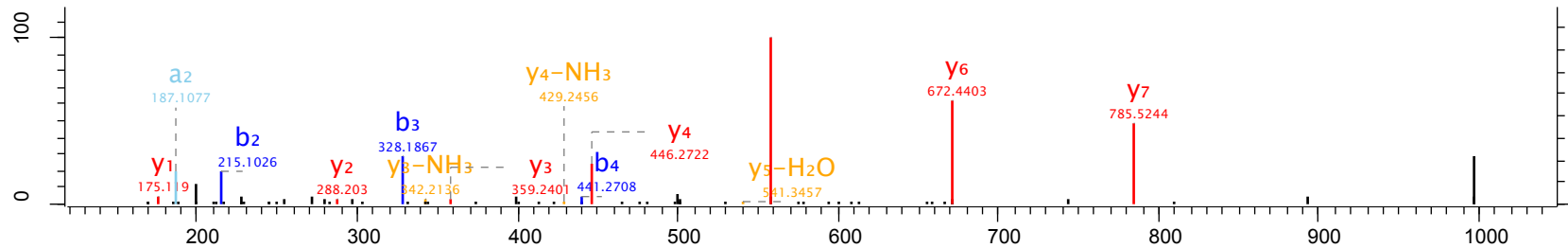
80.75

m/z

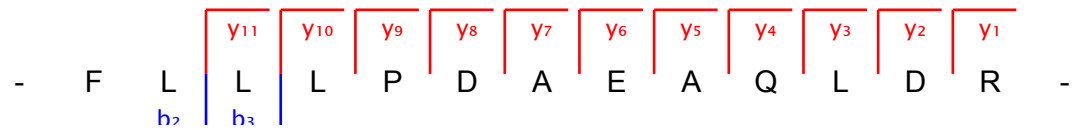
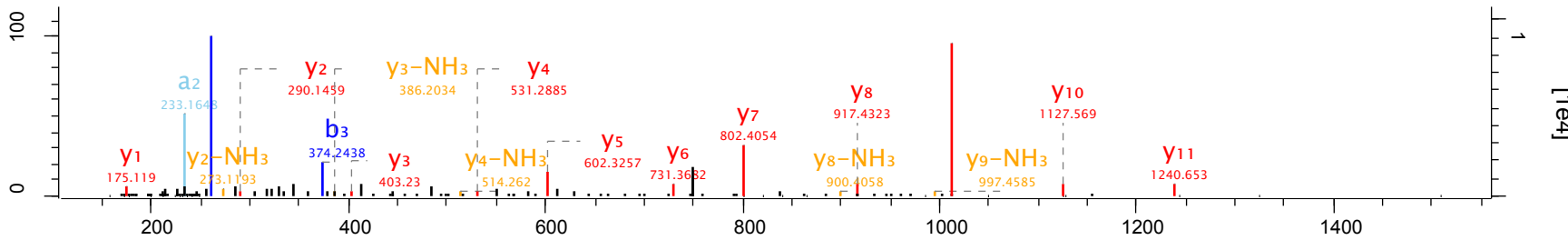
500.31

Gene names

IFI44



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract5_dyn_5ul_E5_01_584	36548	TOF; CID	90.41	750.9	TMEM128



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

4782

Method

TOF; CID

Score

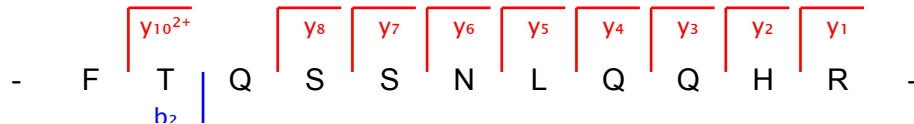
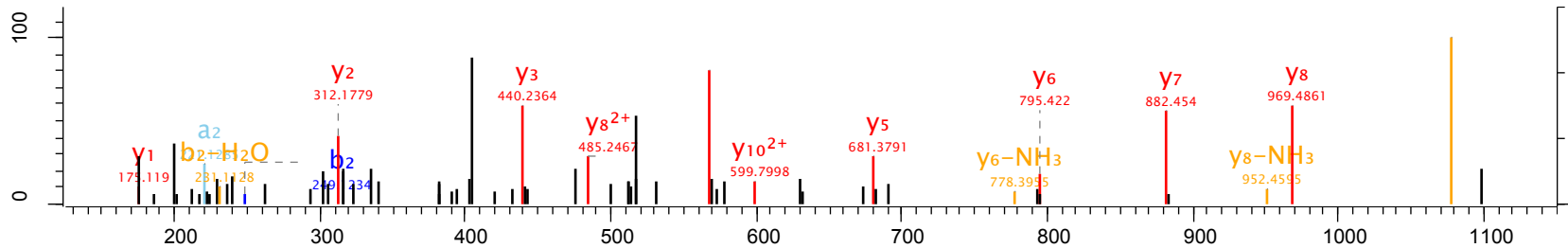
52.39

m/z

449.22

Gene names

ZNF526



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

5304

Method

TOF; CID

Score

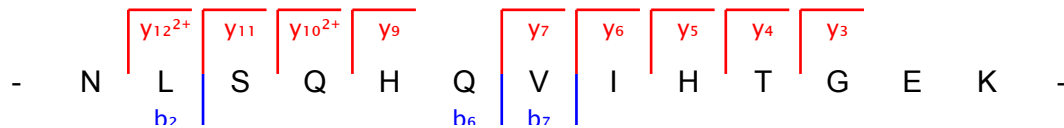
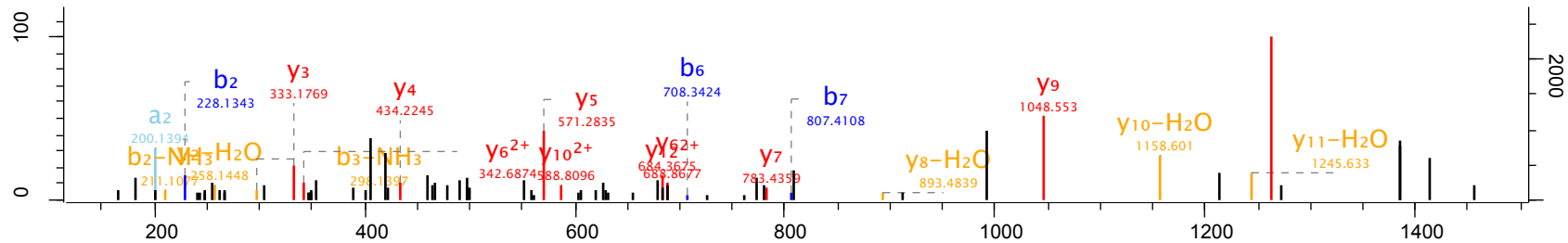
71.88

m/z

497.59

Gene names

ZNF689



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

5558

Method

TOF; CID

Score

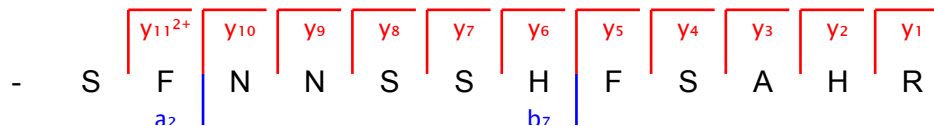
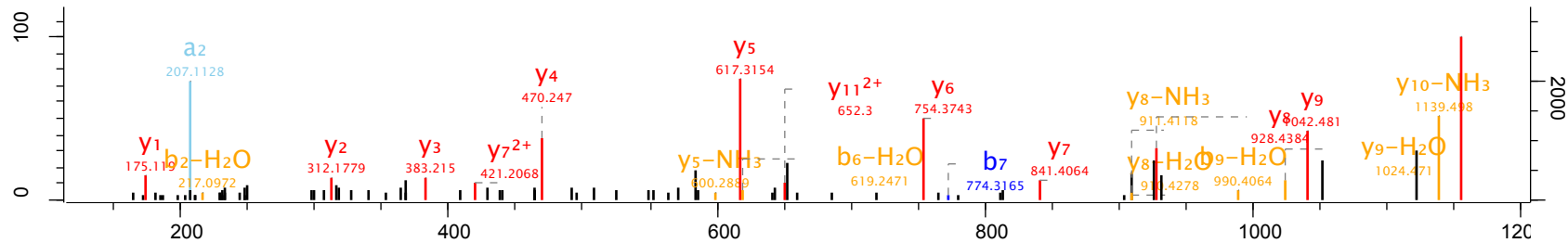
84.17

m/z

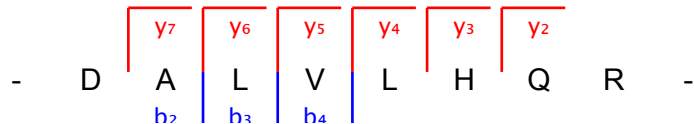
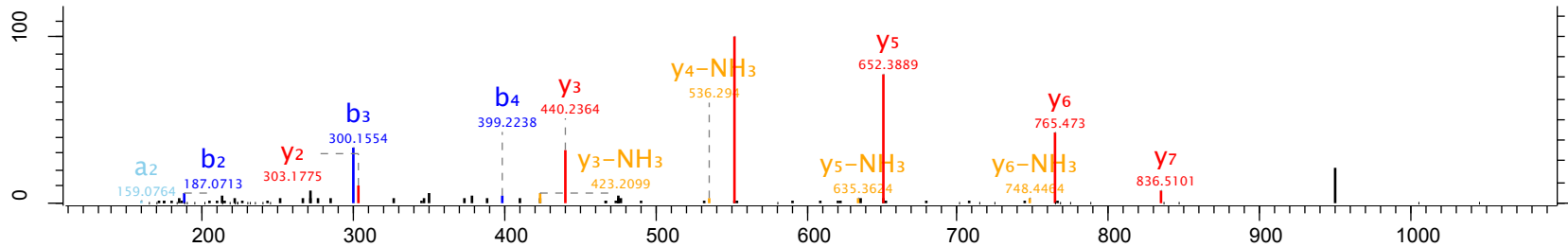
464.21

Gene names

ZSCAN29



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract6_dyn_5ul_E6_01_585	9632	TOF; CID	111.01	476.27	ZNF584



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

9932

Method

TOF; CID

Score

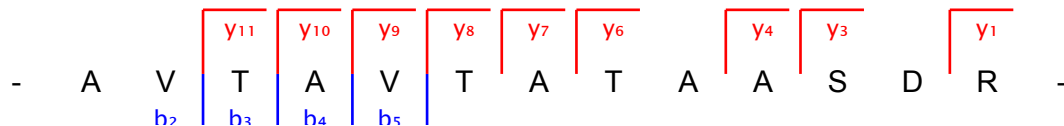
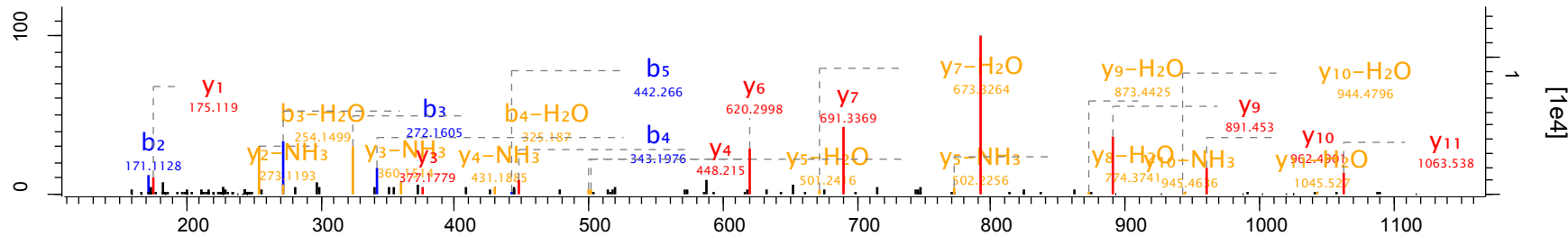
113.99

m/z

617.32

Gene names

HDGFRP2



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

11959

Method

TOF; CID

Score

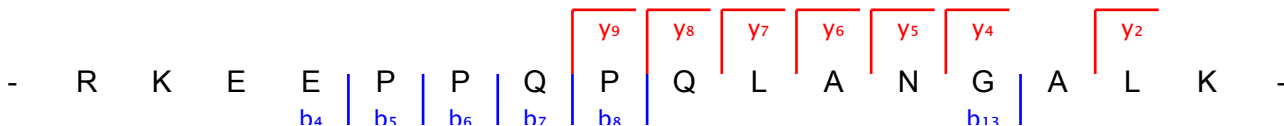
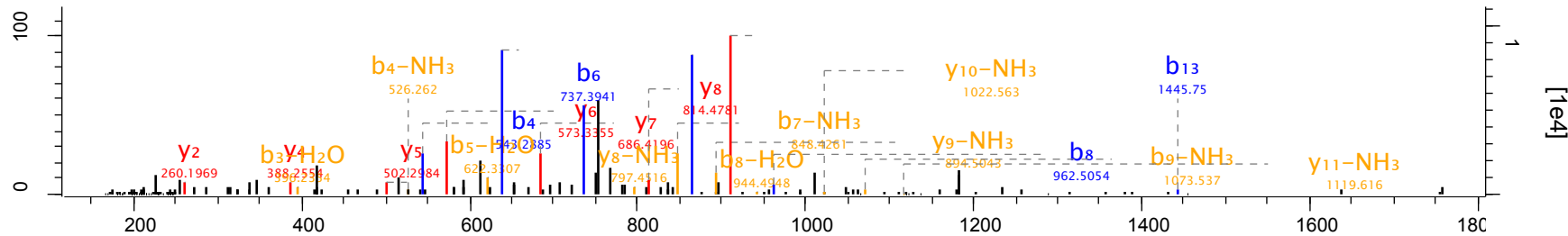
62.86

m/z

592.66

Gene names

C20orf24



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract6_dyn_5ul_E6_01_585

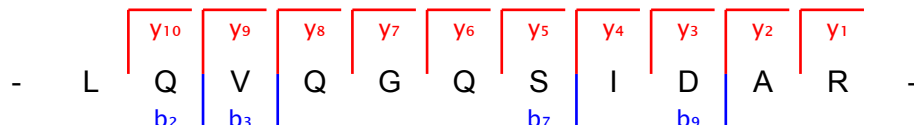
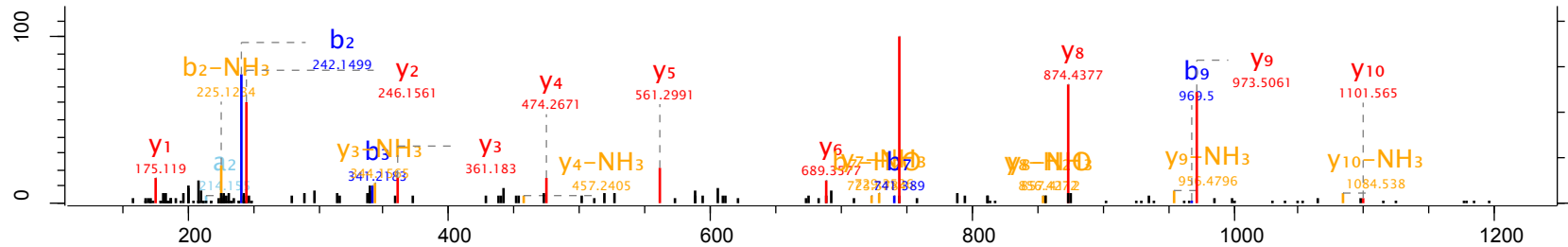
12811

TOF; CID

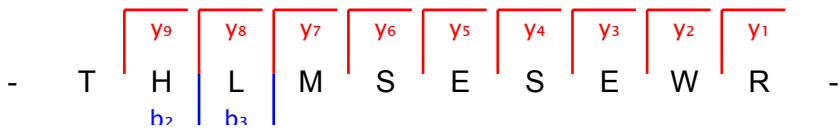
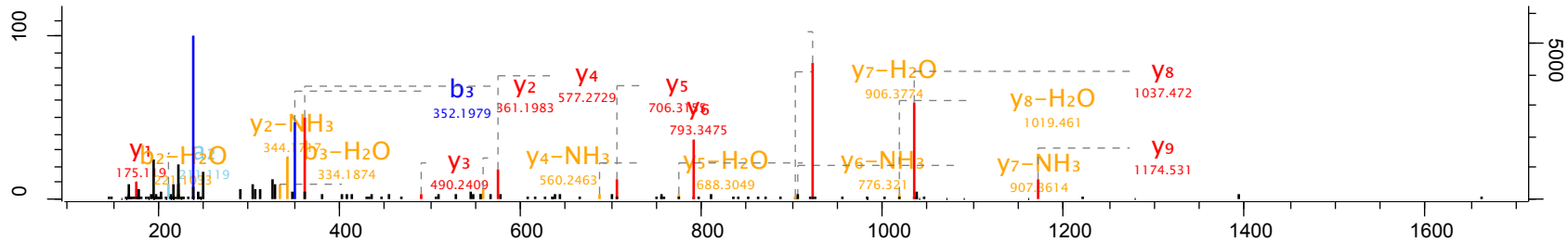
96.74

607.83

SLC25A14



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract6_dyn_5ul_E6_01_585	13708	TOF; CID	123.79	638.29	CKS1B



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

18036

Method

TOF; CID

Score

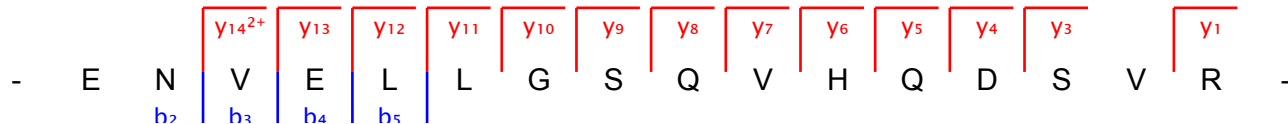
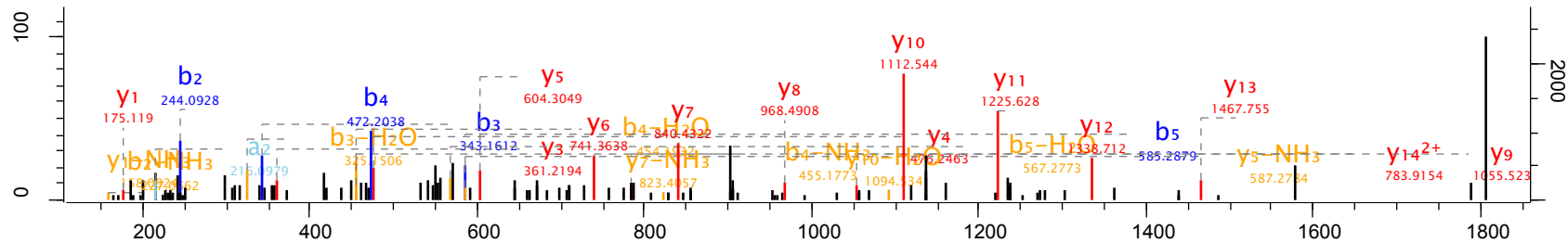
77.22

m/z

905.46

Gene names

TMEM44



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

26348

Method

TOF; CID

Score

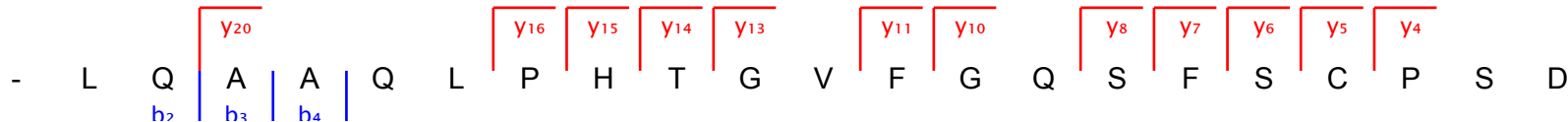
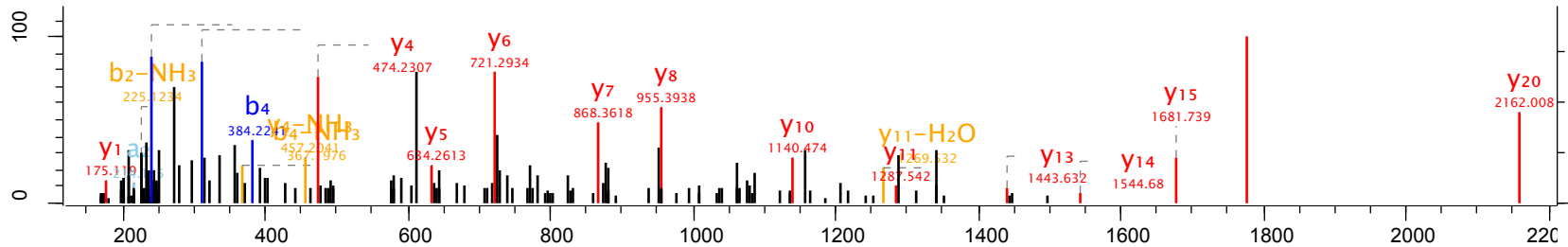
61.96

m/z

802.06

Gene names

CD68



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

33574

Method

TOF; CID

Score

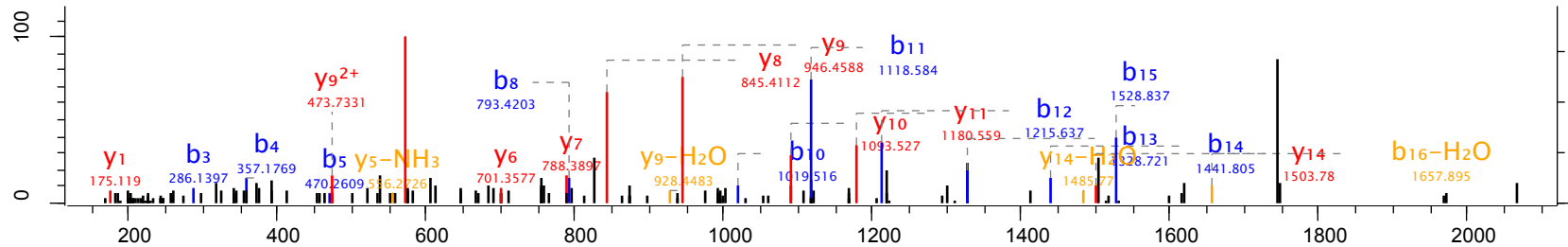
91.24

m/z

874.79

Gene names

ALDH8A1



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

33729

Method

TOF; CID

Score

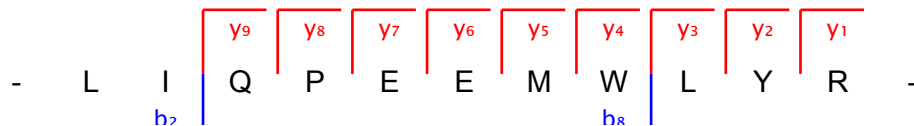
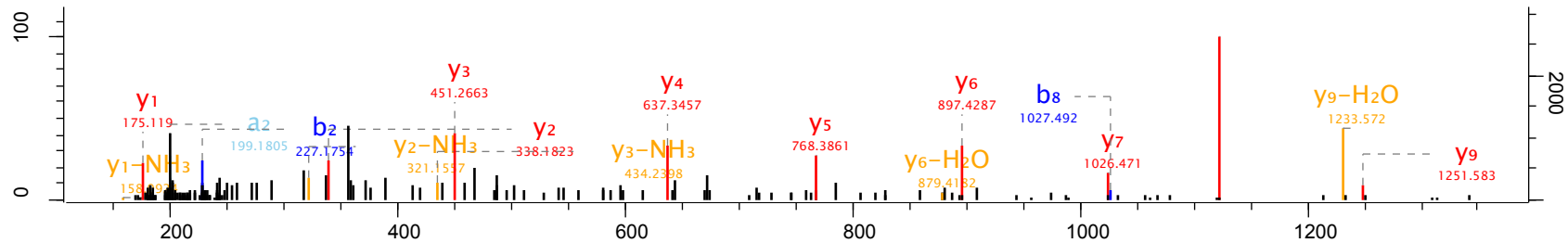
88.6

m/z

739.38

Gene names

PPAPDC1B



Raw file

20141014_fract6_dyn_5ul_E6_01_585

Scan

34378

Method

TOF; CID

Score

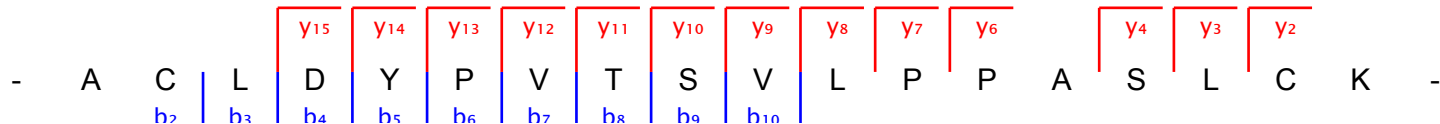
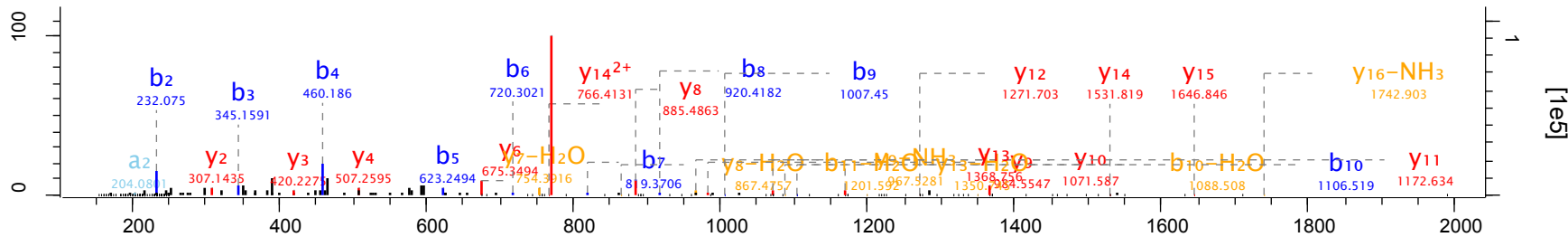
111.01

m/z

996.5

Gene names

PTTG1IP



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

4502

Method

TOF; CID

Score

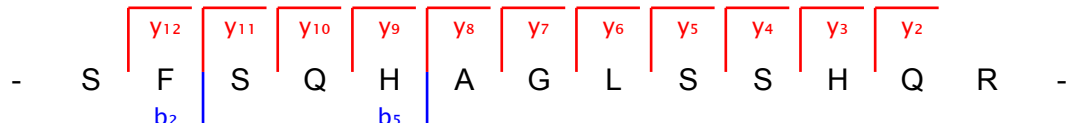
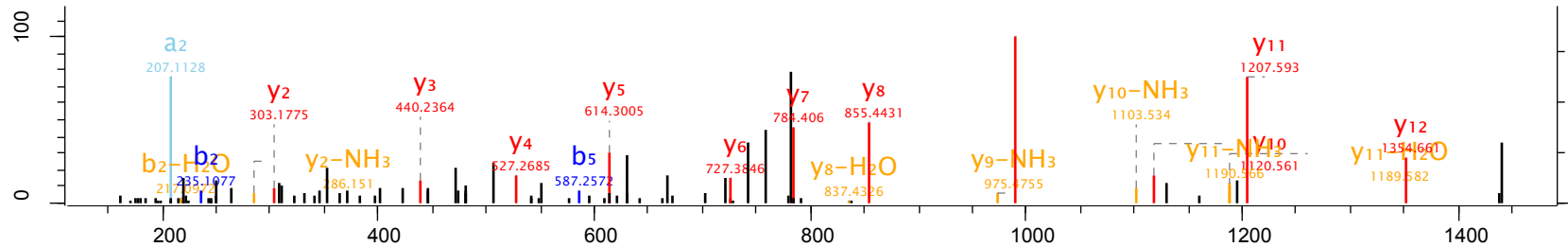
74.48

m/z

481.23

Gene names

ZSCAN21



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

4604

Method

TOF; CID

Score

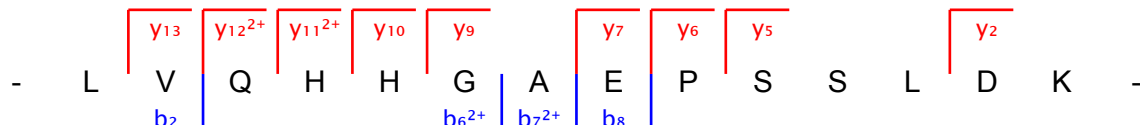
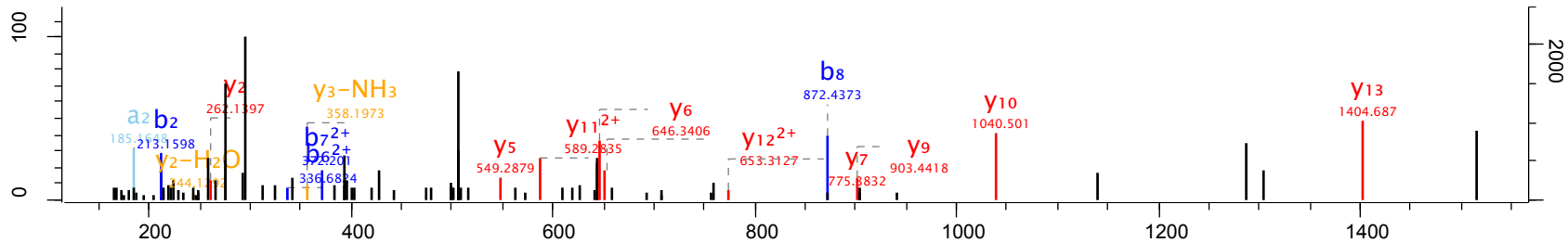
55.72

m/z

506.59

Gene names

STOX2;GDAP1



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

7247

Method

TOF; CID

Score

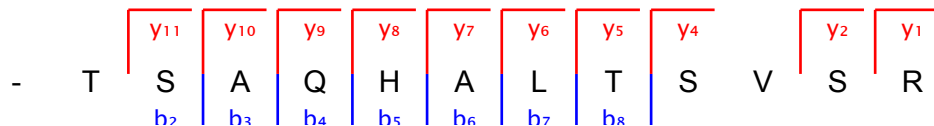
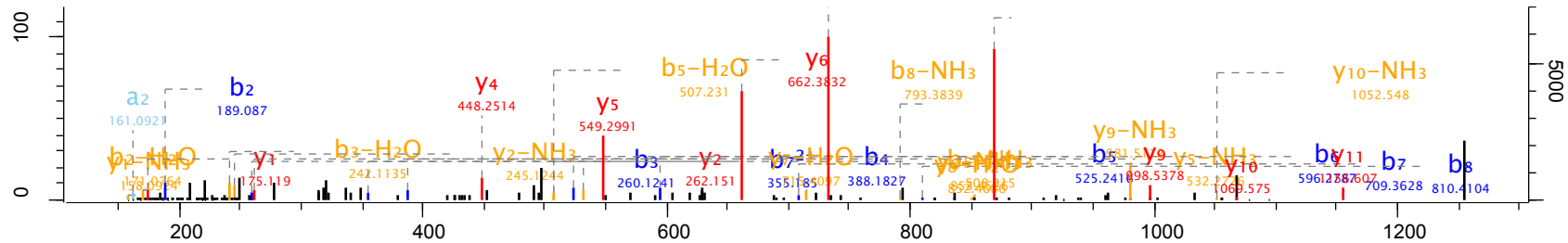
121.86

m/z

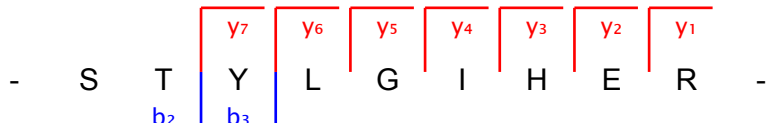
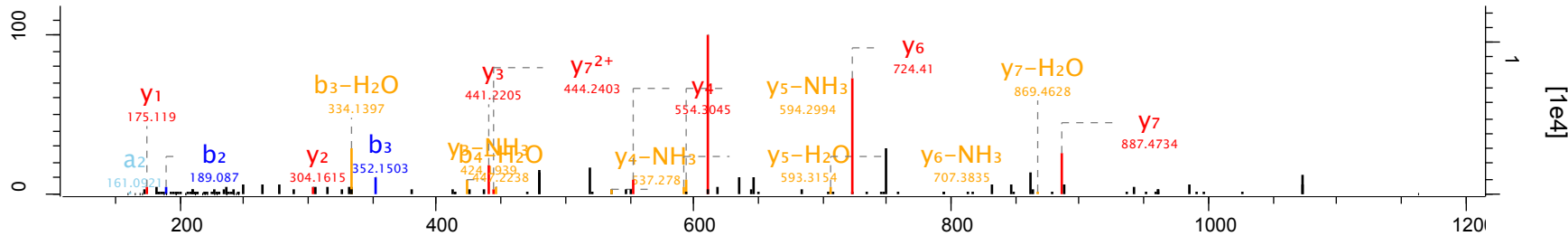
629.33

Gene names

CXCR4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract7_dyn_5ul_E7_01_586	10964	TOF; CID	78.94	538.28	ZNF670



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

12817

Method

TOF; CID

Score

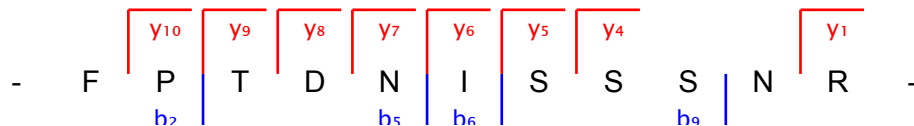
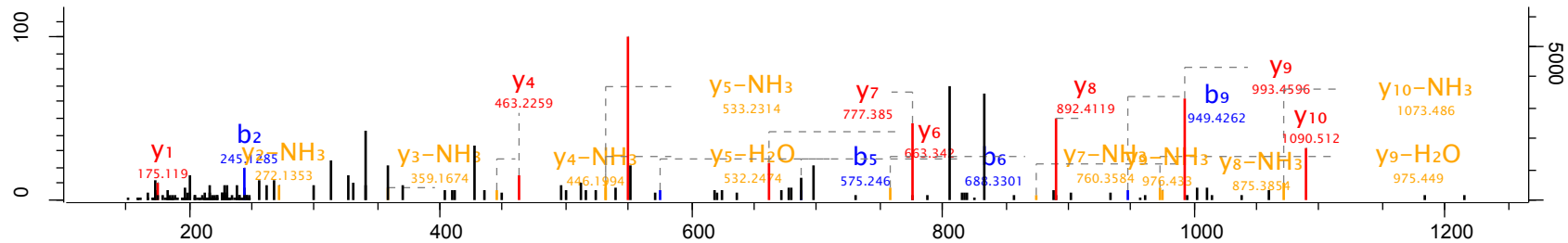
77.6

m/z

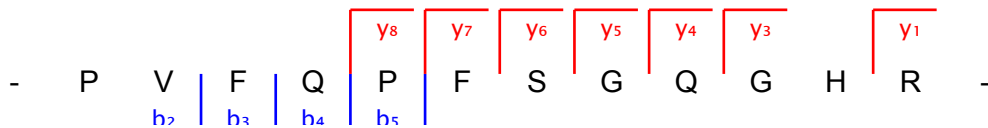
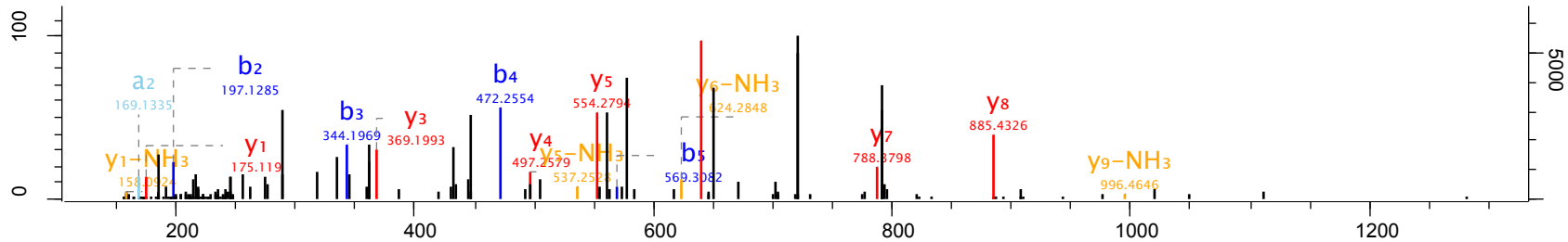
619.29

Gene names

USP53



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract7_dyn_5ul_E7_01_586	17754	TOF; CID	49.63	453.23	UBXN2A



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

18292

Method

TOF; CID

Score

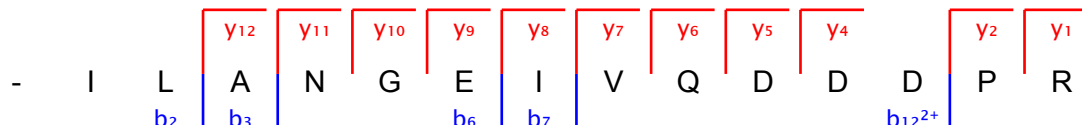
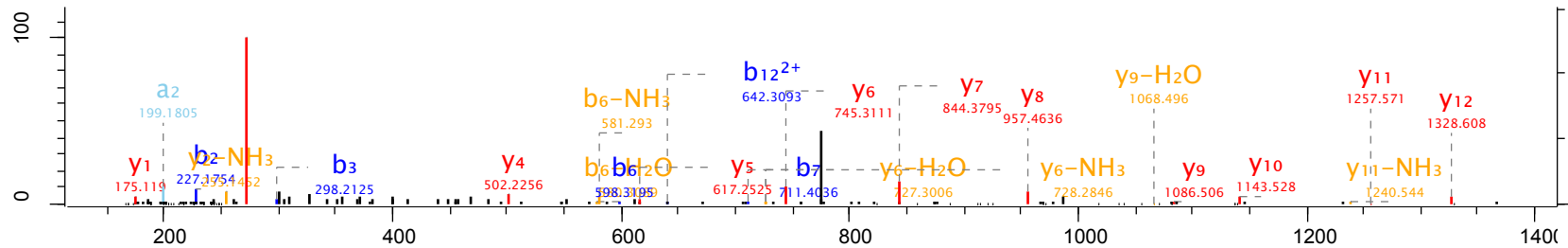
103.13

m/z

777.89

Gene names

C10orf35



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

20843

Method

TOF; CID

Score

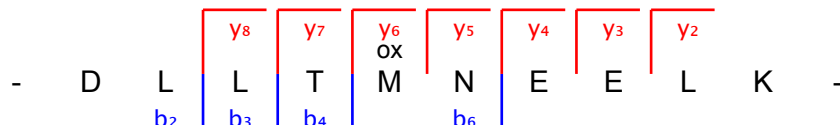
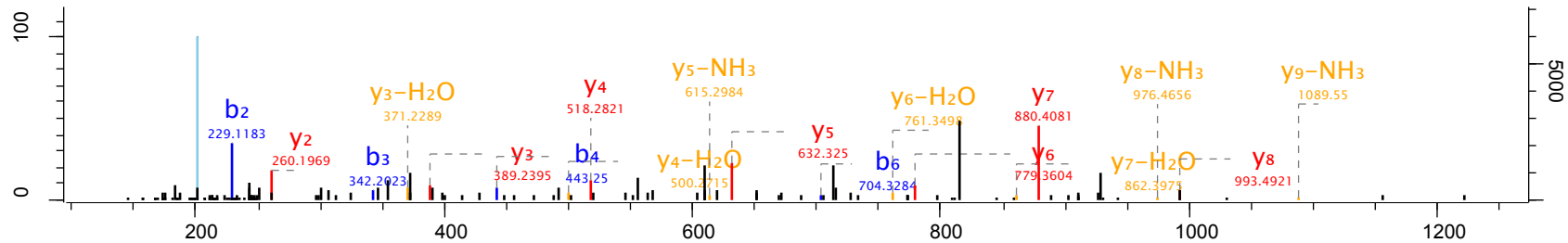
67.93

m/z

611.8

Gene names

TAF13



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

20969

Method

TOF; CID

Score

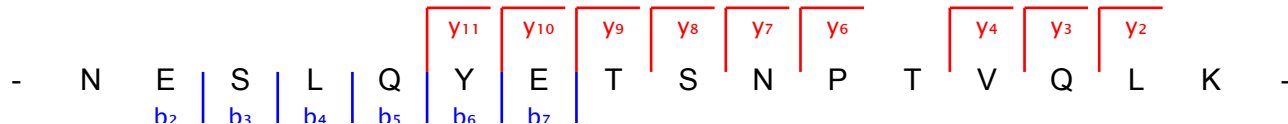
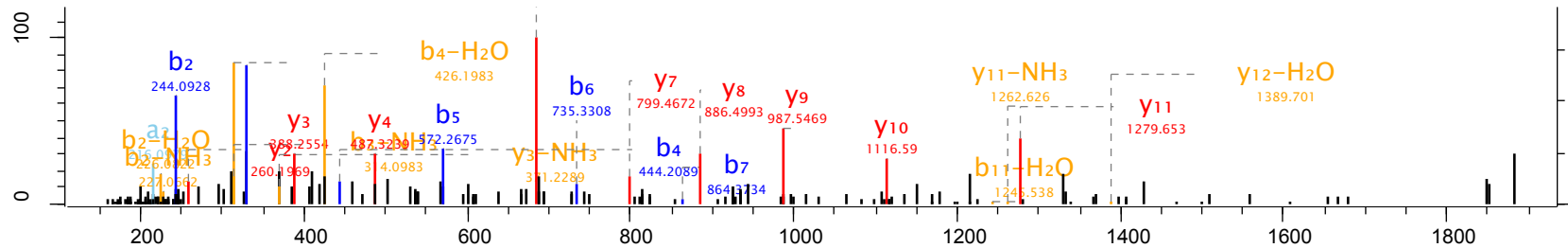
80.74

m/z

926.46

Gene names

LCORL



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

24972

Method

TOF; CID

Score

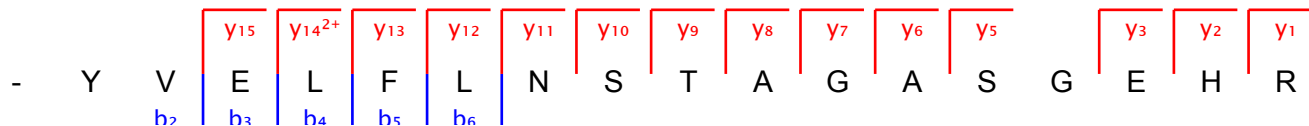
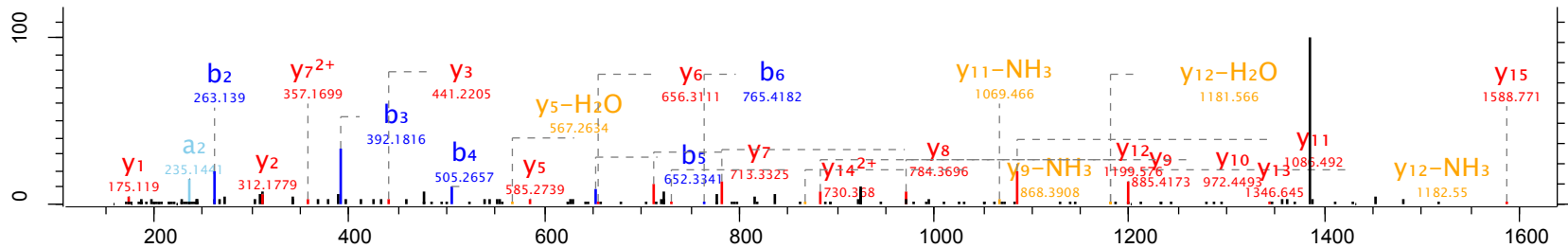
85.36

m/z

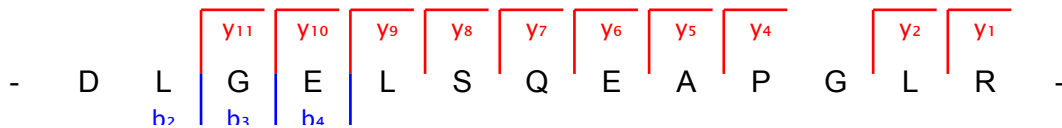
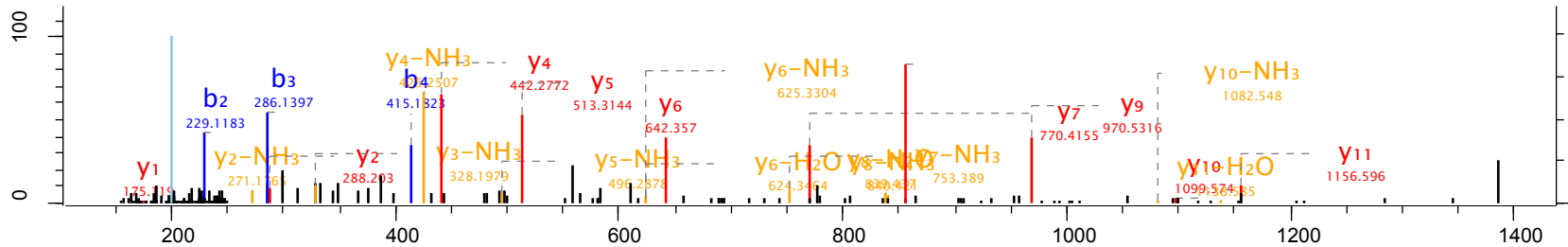
926.46

Gene names

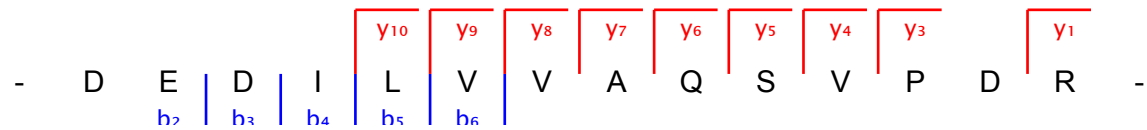
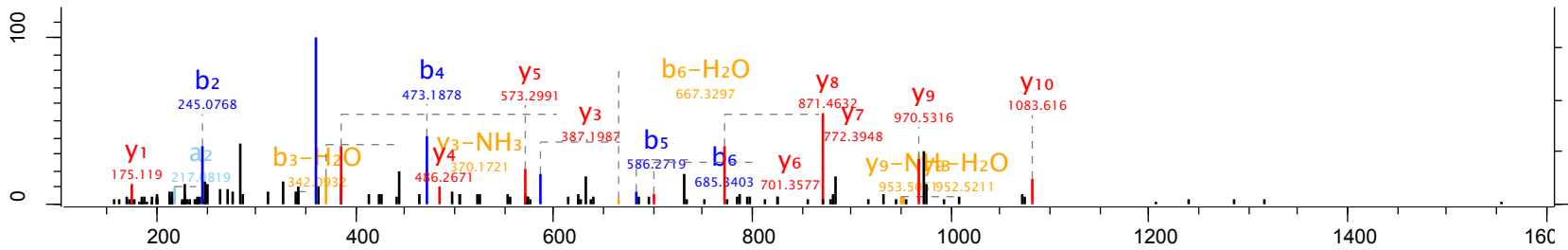
HNRNPH1



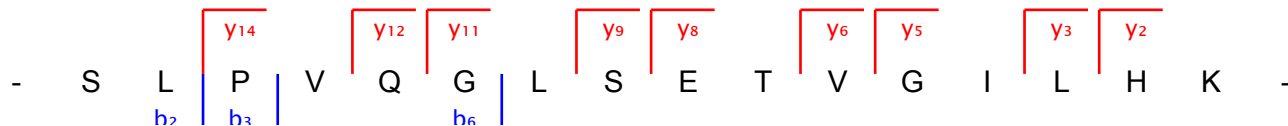
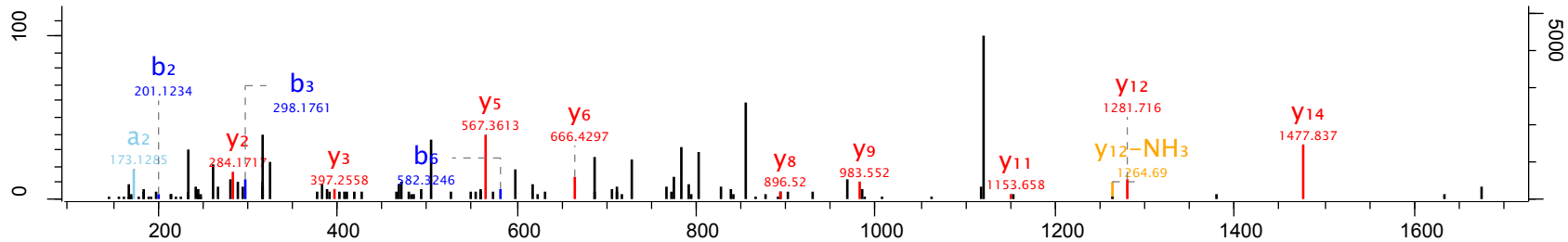
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract7_dyn_5ul_E7_01_586	25098	TOF; CID	92.86	692.86	PLEKHO2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract7_dyn_5ul_E7_01_586	29994	TOF; CID	63.29	778.4	TBC1D16



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract7_dyn_5ul_E7_01_586	31114	TOF; CID	39.34	560.32	GRAMD3



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

34957

Method

TOF; CID

Score

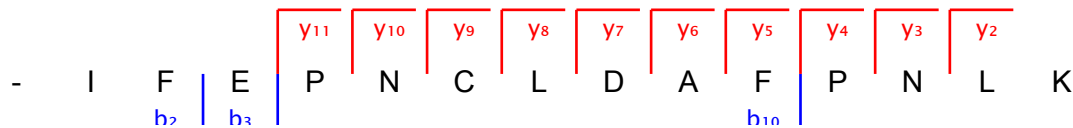
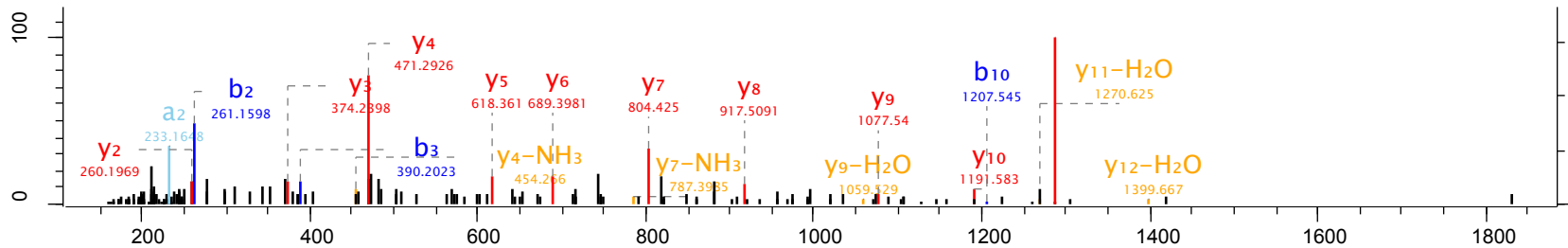
63.16

m/z

839.42

Gene names

GSTM4



Raw file

20141014_fract7_dyn_5ul_E7_01_586

Scan

37190

Method

TOF; CID

Score

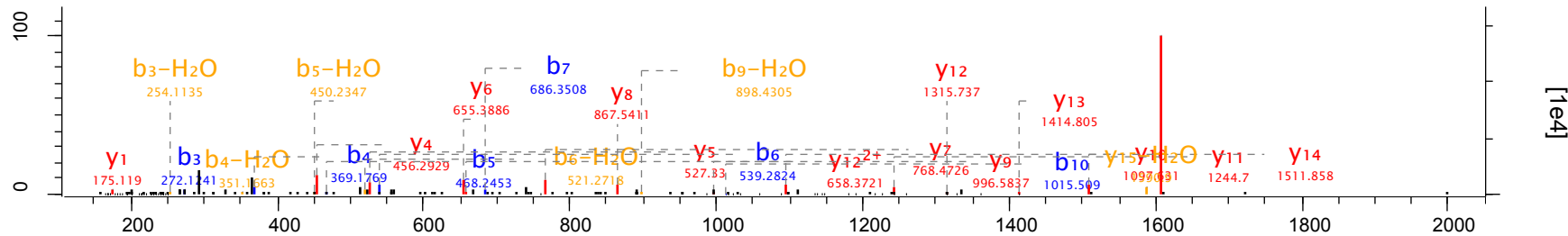
93.23

m/z

892.49

Gene names

PPP1R3F



- S S P P V A F T E V L Q A P A I R -

b₃ b₄ b₅ b₆ b₇ b₁₀

y₁₅ y₁₄ y₁₃ y₁₂ y₁₁ y₁₀ y₉ y₈ y₇ y₆ y₅ y₄ y₁

Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

9717

Method

TOF; CID

Score

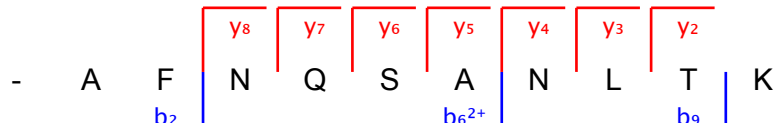
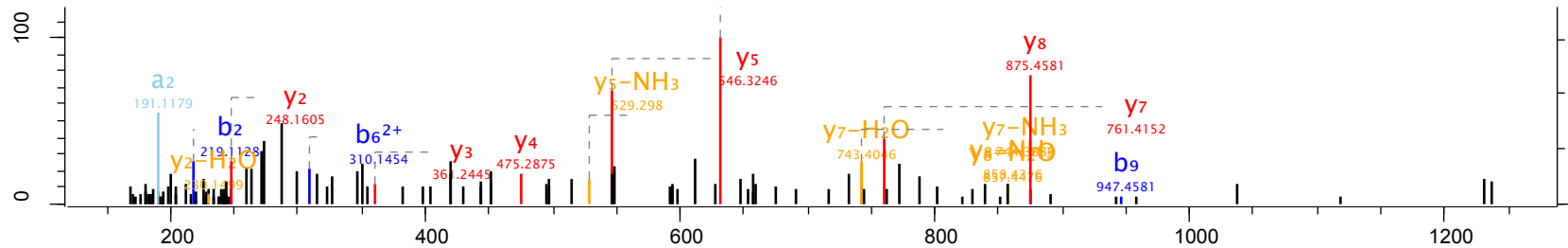
72.2

m/z

547.28

Gene names

ZNF675



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

9944

Method

TOF; CID

Score

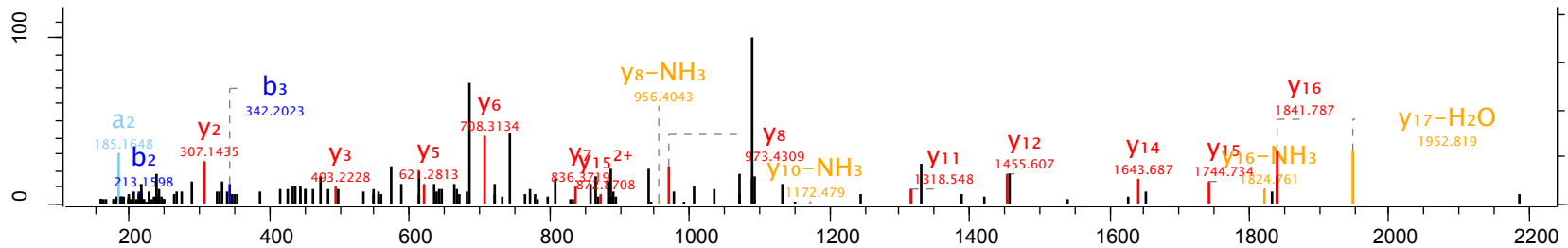
49.04

m/z

546.75

Gene names

LPHN1



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

13387

Method

TOF; CID

Score

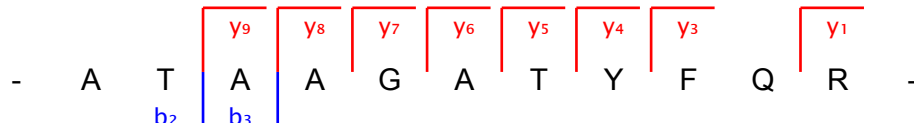
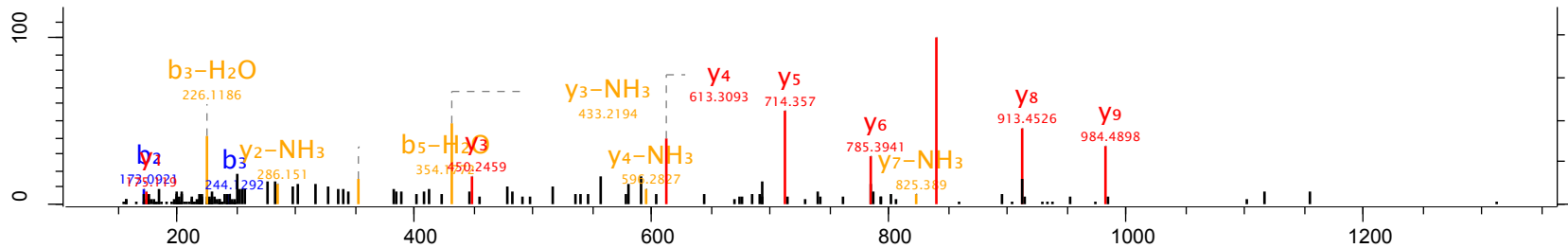
59.2

m/z

578.79

Gene names

TMEM254



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

13435

Method

TOF; CID

Score

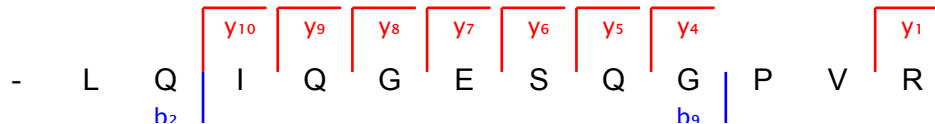
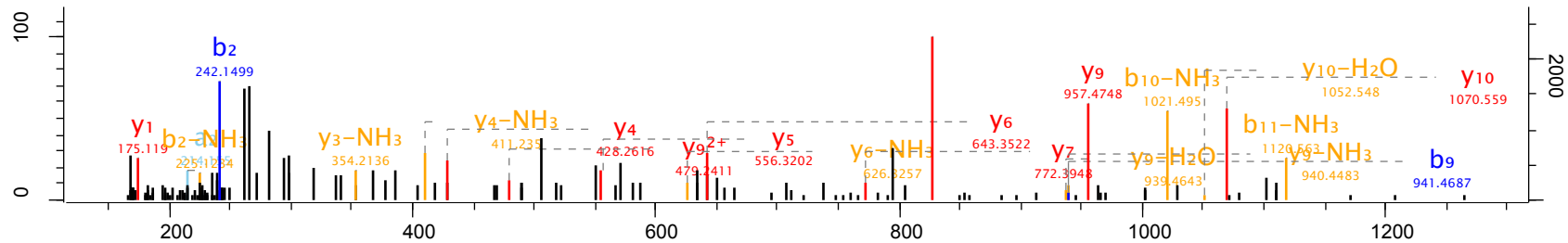
68.52

m/z

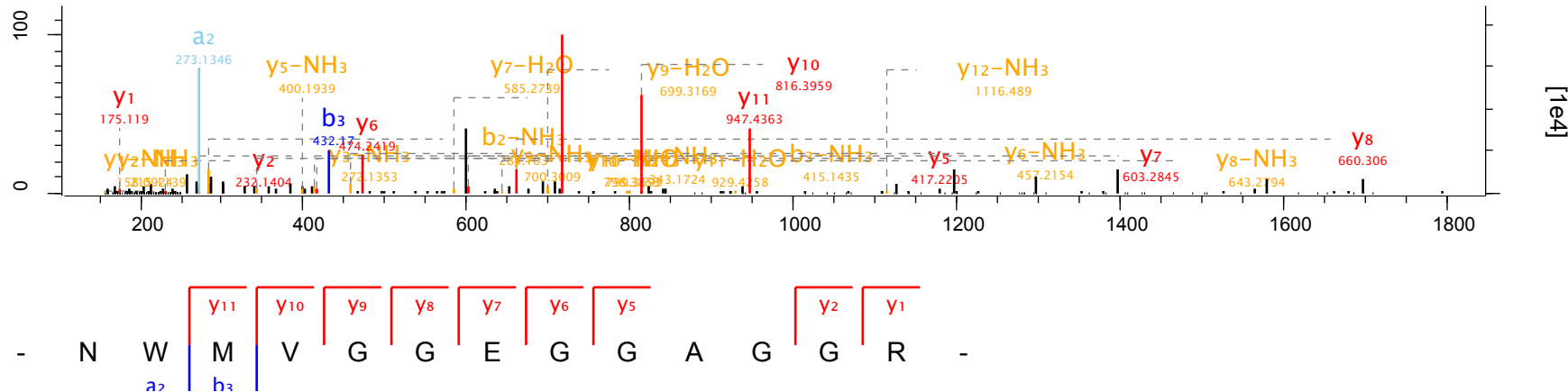
656.35

Gene names

UCP2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract8_dyn_5ul_E8_01_587	16168	TOF; CID	63.61	624.78	C11orf83



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

17376

Method

TOF; CID

Score

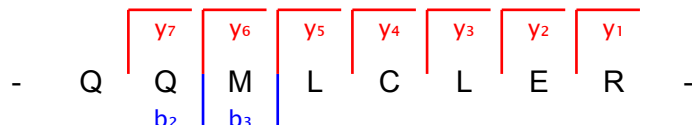
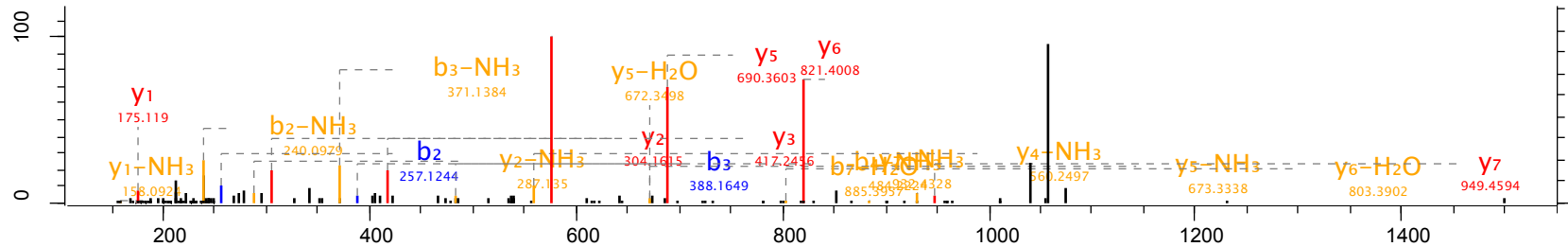
110.41

m/z

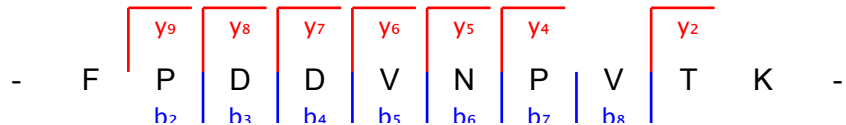
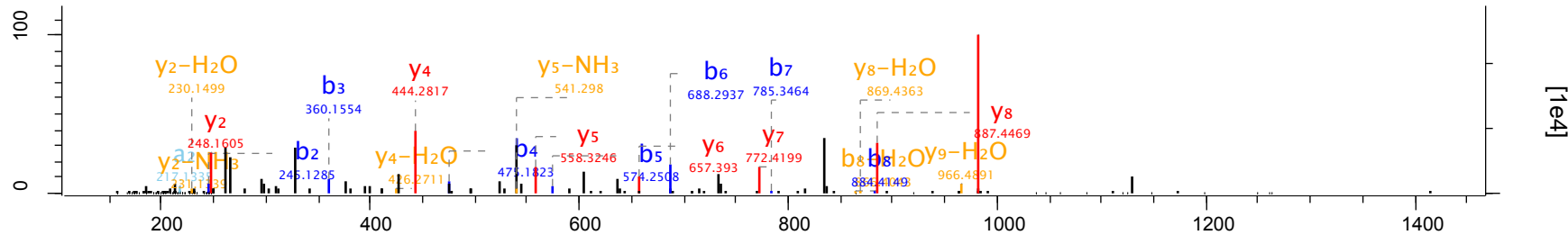
539.26

Gene names

NPDC1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract8_dyn_5ul_E8_01_587	17469	TOF; CID	94.82	566.29	C6orf57



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

23332

Method

TOF; CID

Score

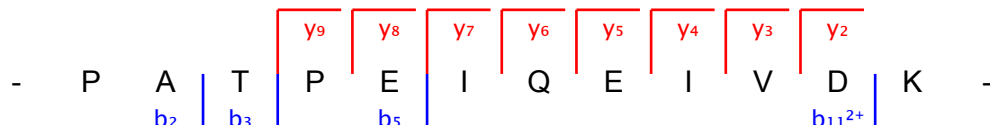
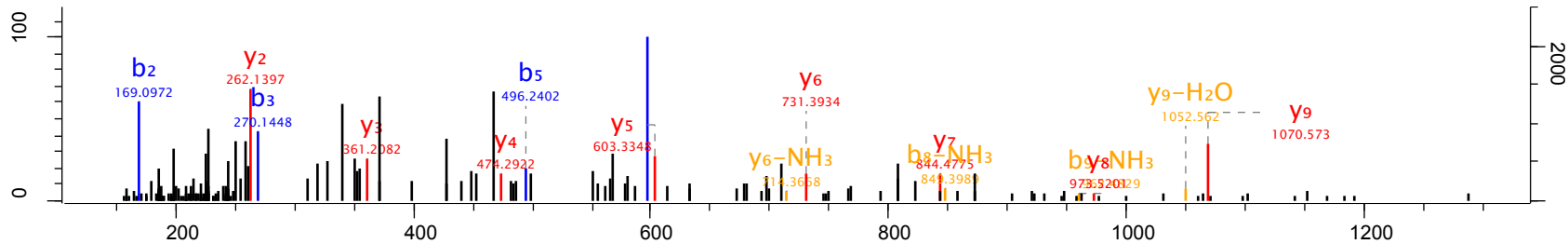
61.11

m/z

670.36

Gene names

CSTA



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

24831

Method

TOF; CID

Score

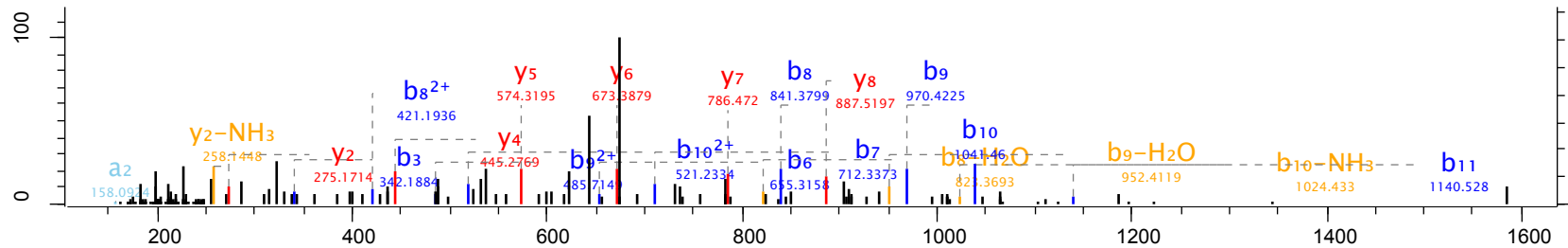
60.33

m/z

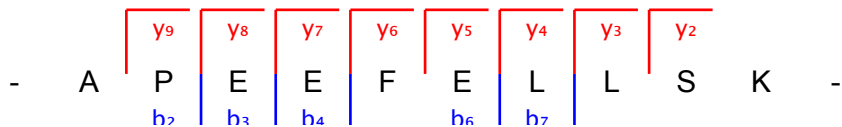
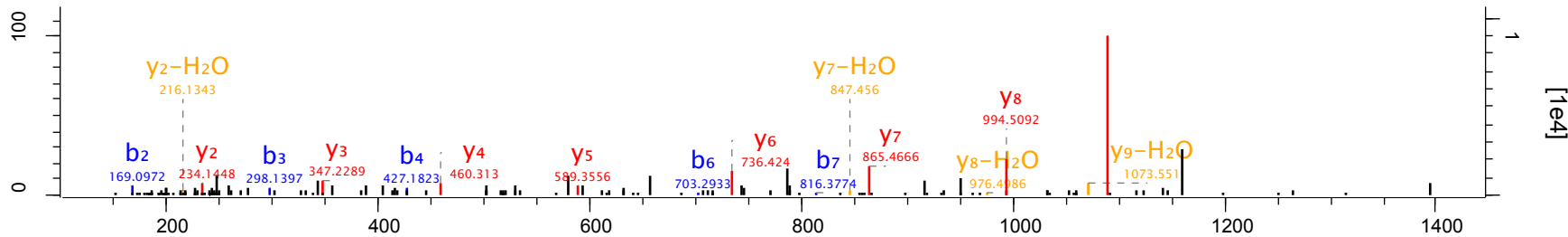
676.35

Gene names

ZNF202



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract8_dyn_5ul_E8_01_587	25255	TOF; CID	67.38	582.29	TMLHE



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

25841

Method

TOF; CID

Score

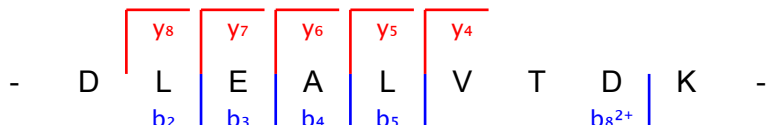
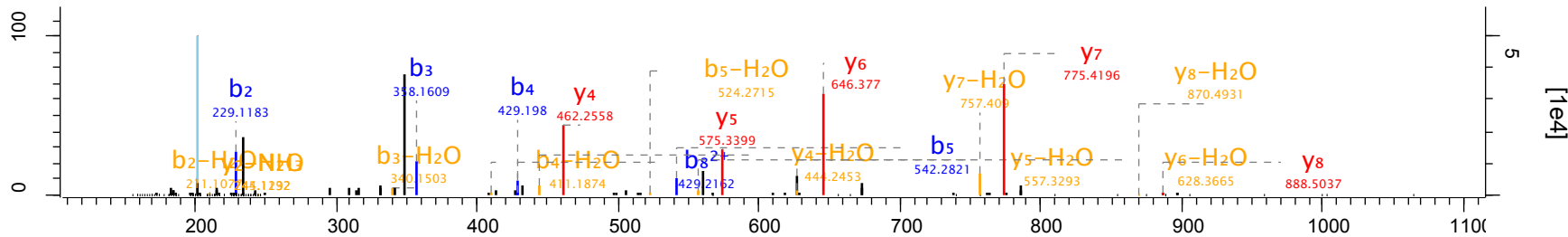
102.07

m/z

502.27

Gene names

PGM5



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

26052

Method

TOF; CID

Score

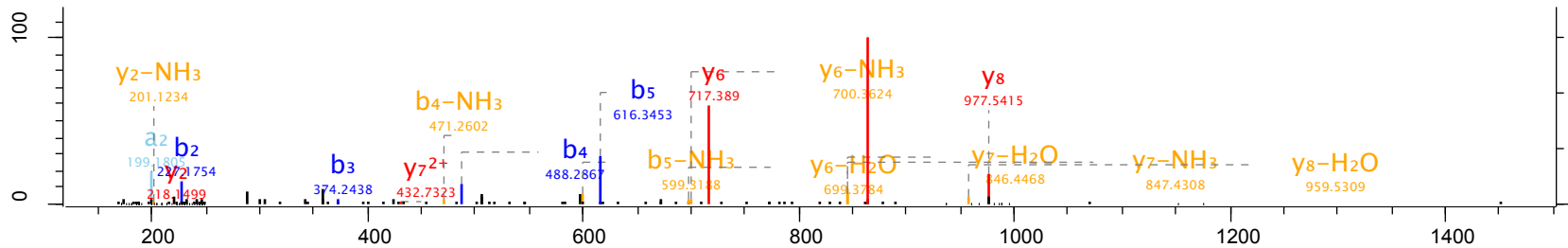
114.28

m/z

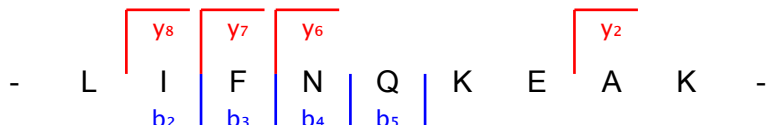
546.32

Gene names

BHLHB9



[1e4]



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

29898

Method

TOF; CID

Score

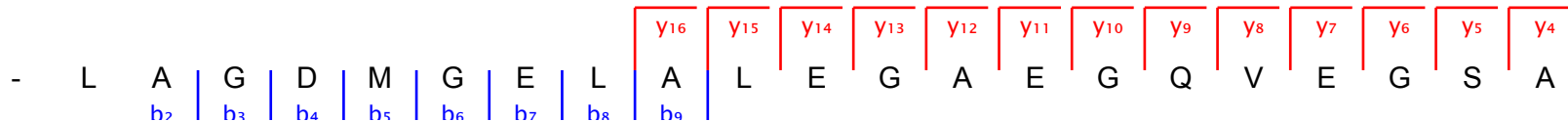
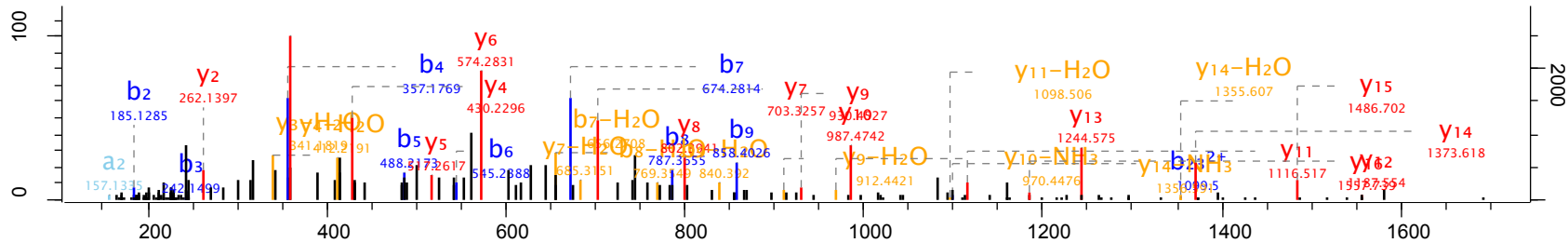
113.42

m/z

1173.05

Gene names

PKIG



Raw file

20141014_fract8_dyn_5ul_E8_01_587

Scan

31118

Method

TOF; CID

Score

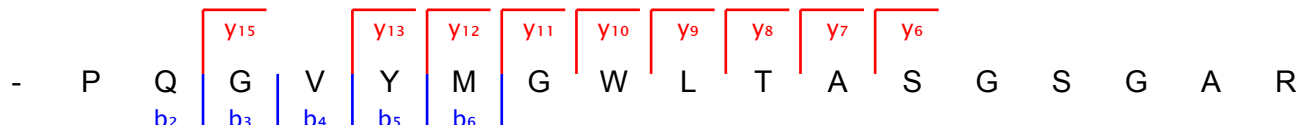
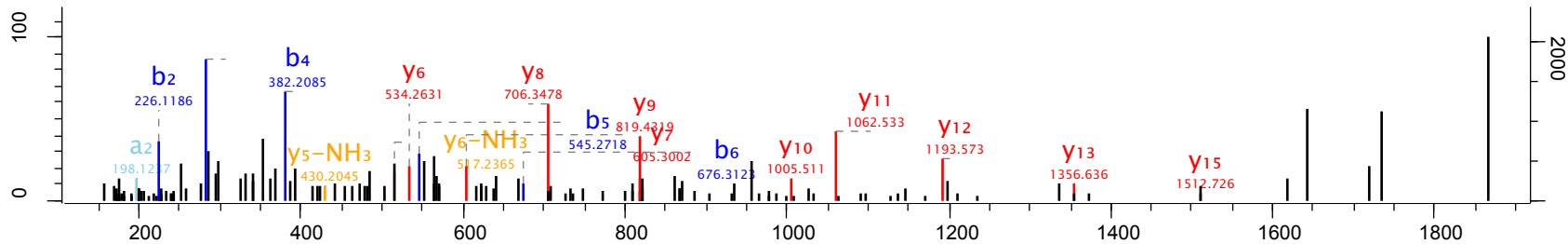
49.54

m/z

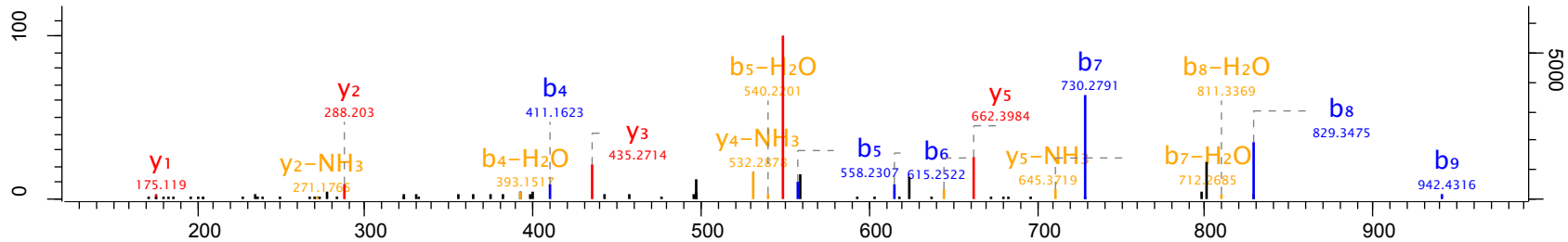
869.42

Gene names

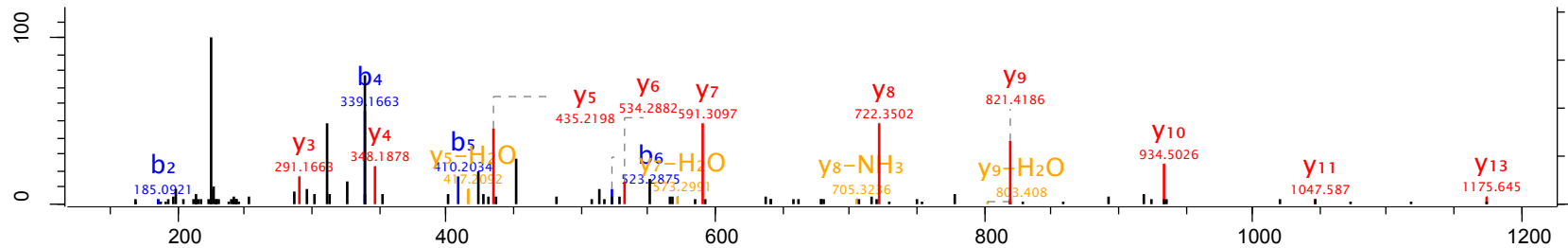
MFSD8



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract8_dyn_5ul_E8_01_587	36842	TOF; CID	80.24	497.58	KCTD7



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract8_dyn_5ul_E8_01_587	38691	TOF; CID	54.28	729.39	IDNK



ac

- A A | P G | A L | L V | M G | V S | G S G K -

b₂ | b₄ | b₅ | b₆

y₁₃ | y₁₁ | y₁₀ | y₉ | y₈ | y₇ | y₆ | y₅ | y₄ | y₃

Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

4674

Method

TOF; CID

Score

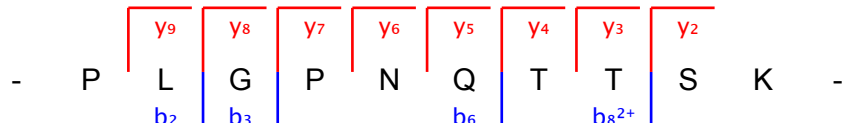
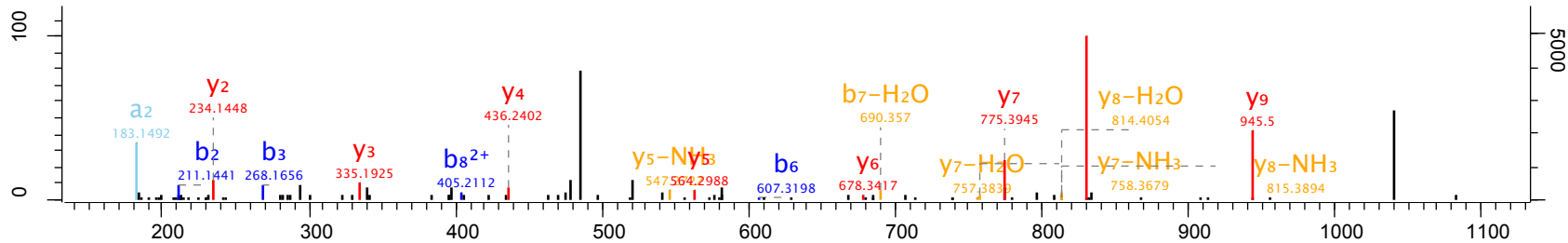
91.55

m/z

521.78

Gene names

SLC16A7



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

Method

Score

m/z

Gene names

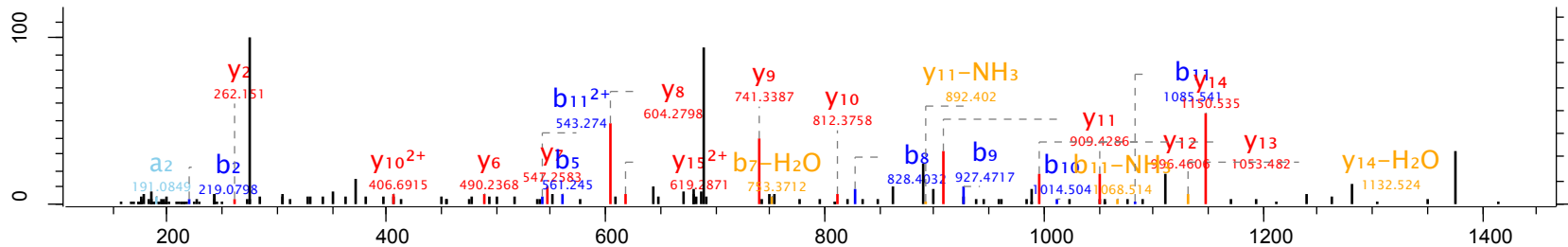
9754

TOF; CID

37.46

690.32

PPAPDC2



- S M | E G R | P L G | V | S | A | S S S S S | S P G S P

h₂ | h₅ | h₈ | h₉ | h₁₀ | h₁₁

y₁₅²⁺ y₁₄ y₁₃ y₁₂ y₁₁

Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

11317

Method

TOF; CID

Score

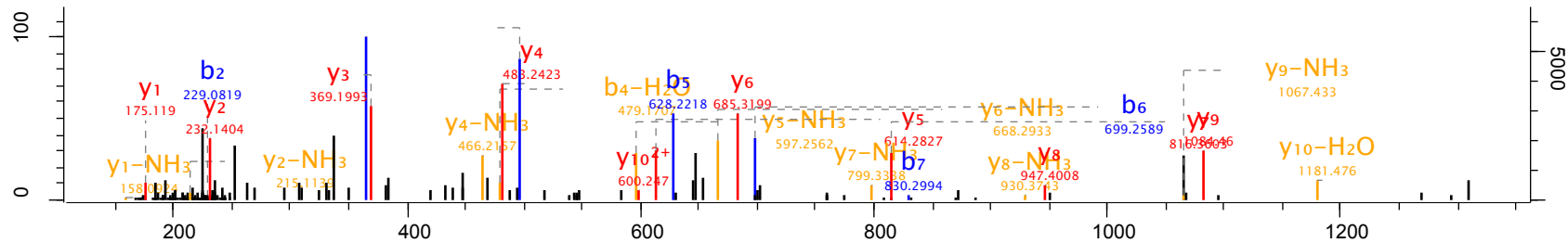
133.81

m/z

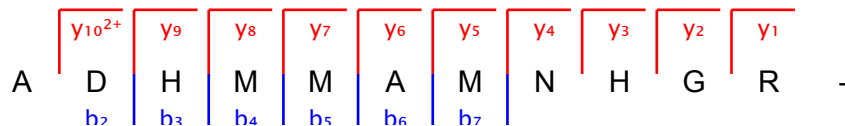
438.18

Gene names

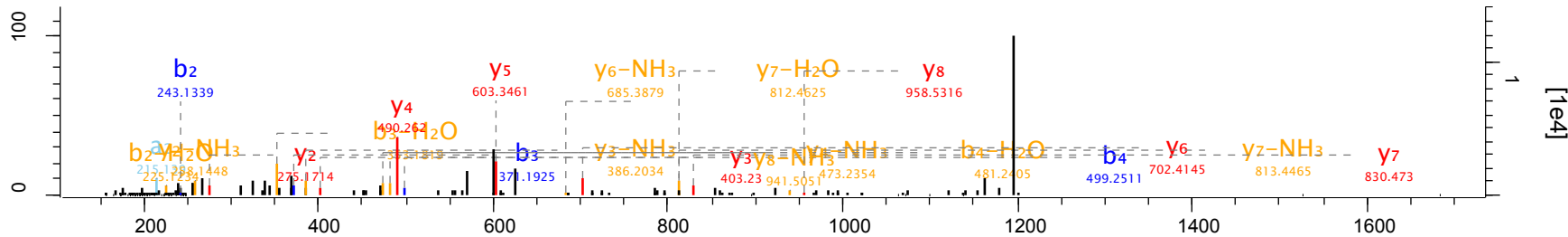
CITED2



ac



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract9_dyn_5ul_F1_01_588	12656	TOF; CID	90.14	600.83	SPATA24



Protein sequence: - E L Q Q V I S Q Q K -

Fragmentation sites (b and y ions) are indicated by brackets above the sequence:

- b₂ (blue) is between L and Q.
- b₃ (blue) is between Q and Q.
- b₄ (blue) is between V and I.
- y₂ (red) is between Q and Q.
- y₃ (red) is between Q and Q.
- y₄ (red) is between S and Q.
- y₅ (red) is between I and S.
- y₆ (red) is between V and I.
- y₇ (red) is between Q and Q.
- y₈ (red) is between L and Q.

Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

13990

Method

TOF; CID

Score

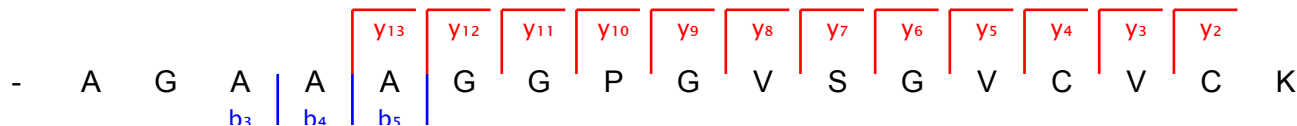
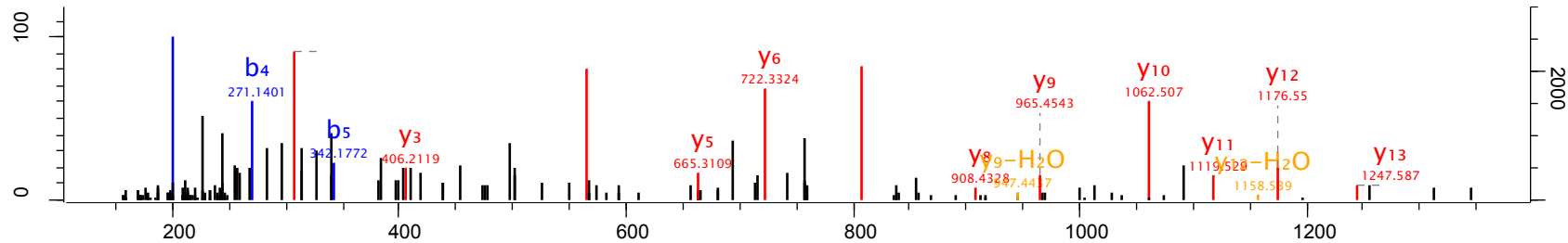
59.73

m/z

759.36

Gene names

IGFBP7



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

15090

Method

TOF; CID

Score

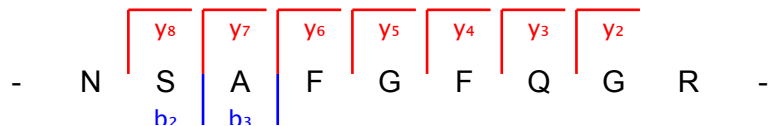
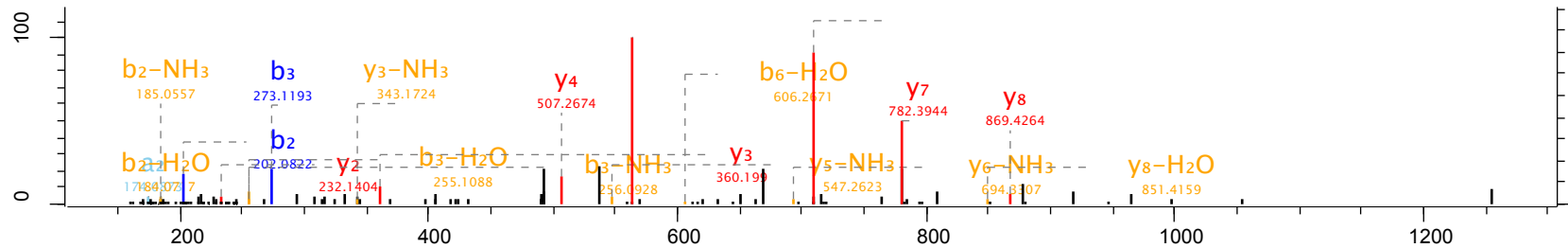
83.62

m/z

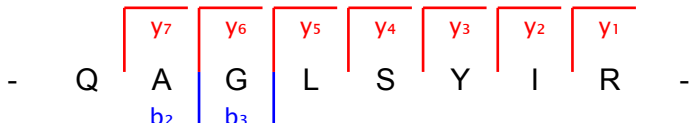
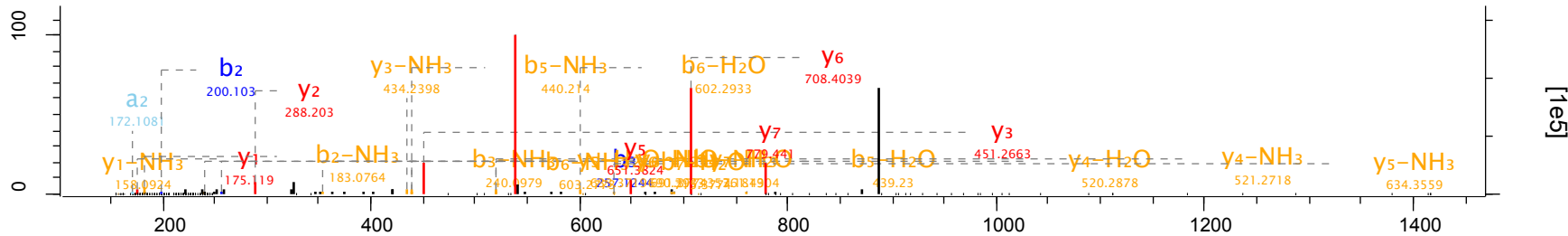
492.24

Gene names

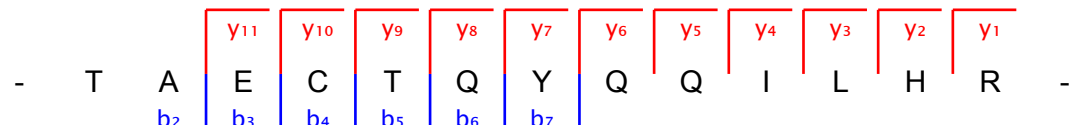
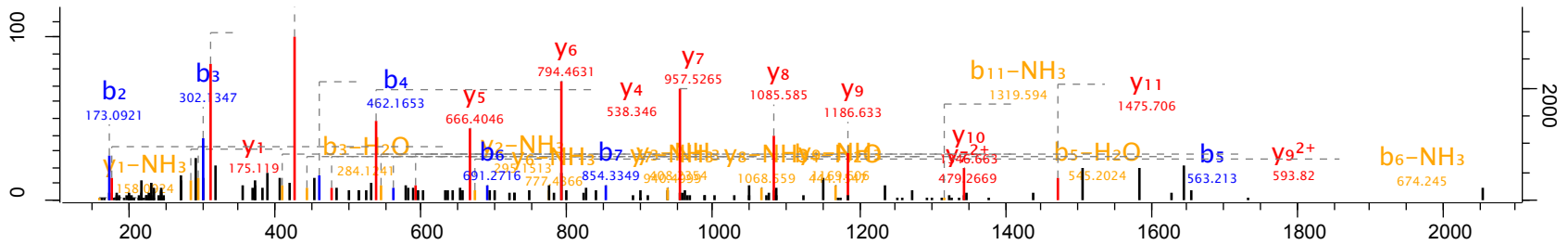
TNFSF9



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract9_dyn_5ul_F1_01_588	15195	TOF; CID	102.46	454.25	ATP5E;ATP5EP2



Raw file: 20141014_fract9_dyn_5ul_F1_01_588
 Scan: 16089
 Method: TOF; CID
 Score: 143.42
 m/z: 824.4
 Gene names: SS18L1



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

18962

Method

TOF; CID

Score

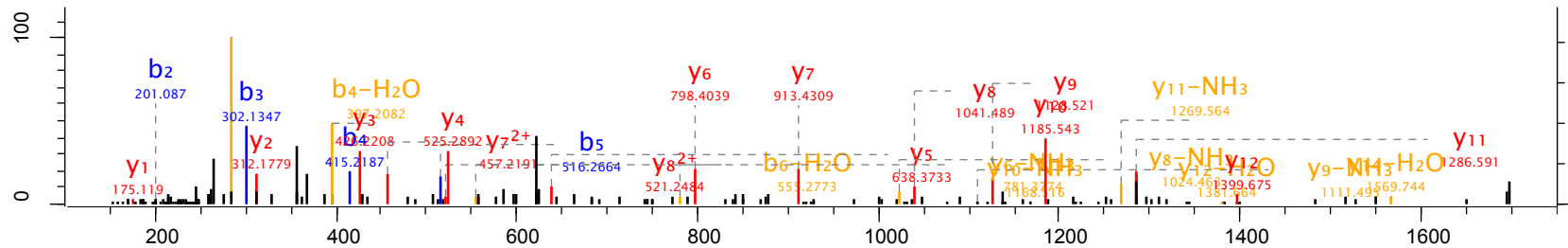
110.44

m/z

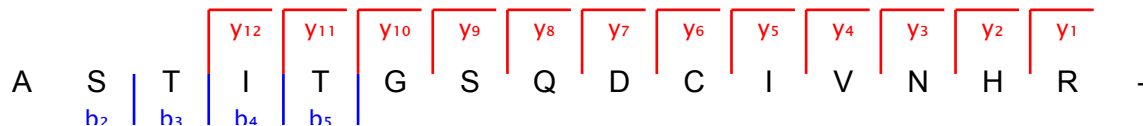
850.9

Gene names

ZNF239



ac



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

23263

Method

TOF; CID

Score

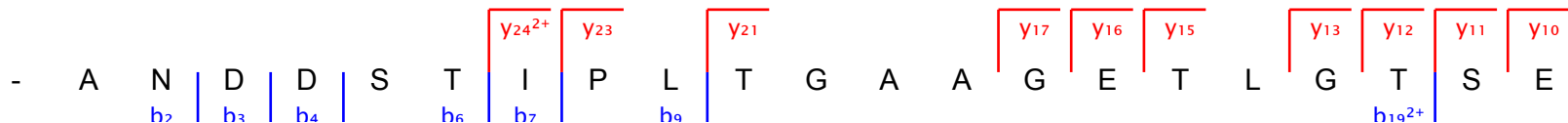
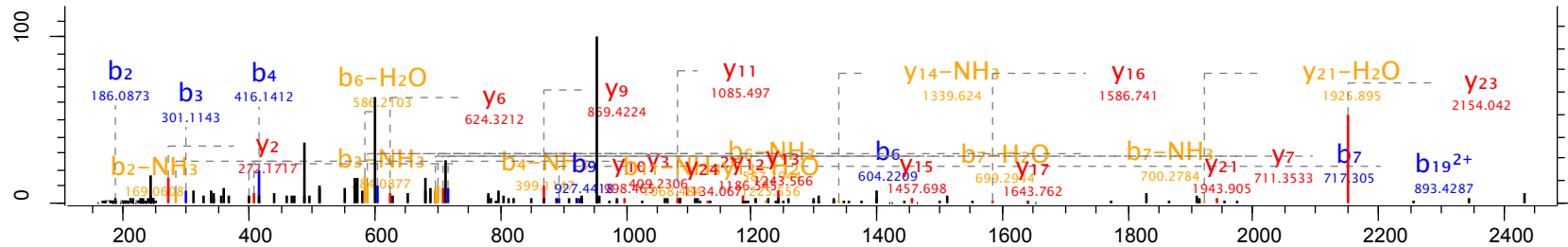
41.36

m/z

957.78

Gene names

SLC20A2



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

23673

Method

TOF; CID

Score

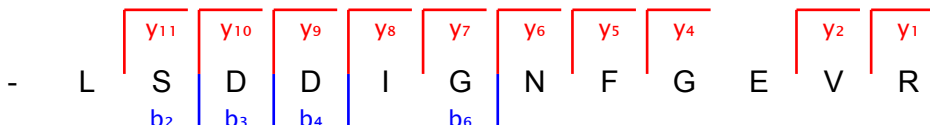
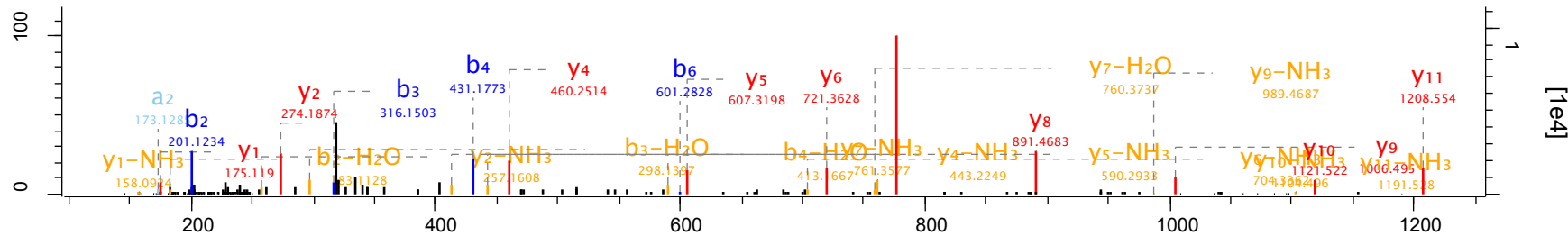
140.14

m/z

661.32

Gene names

SLC6A9



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

25104

Method

TOF; CID

Score

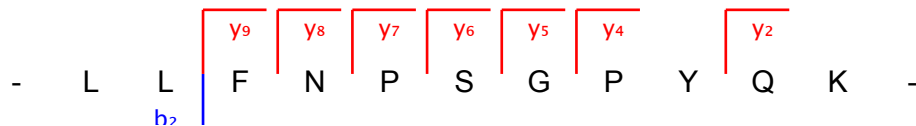
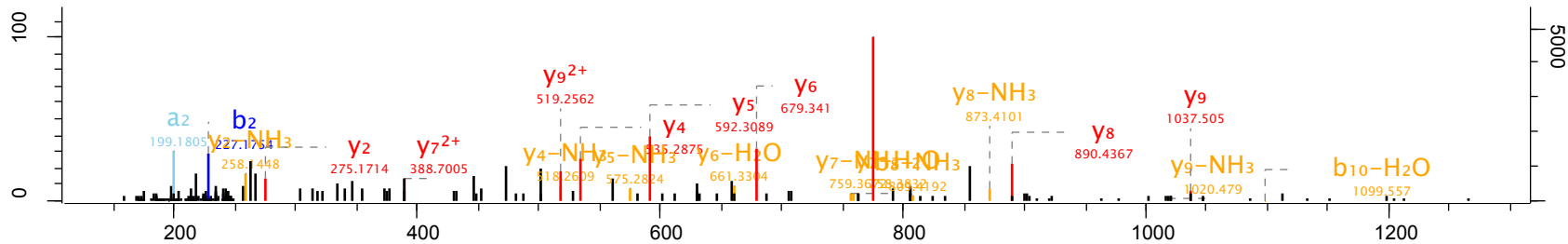
93.32

m/z

632.34

Gene names

CKLF



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

29640

Method

TOF; CID

Score

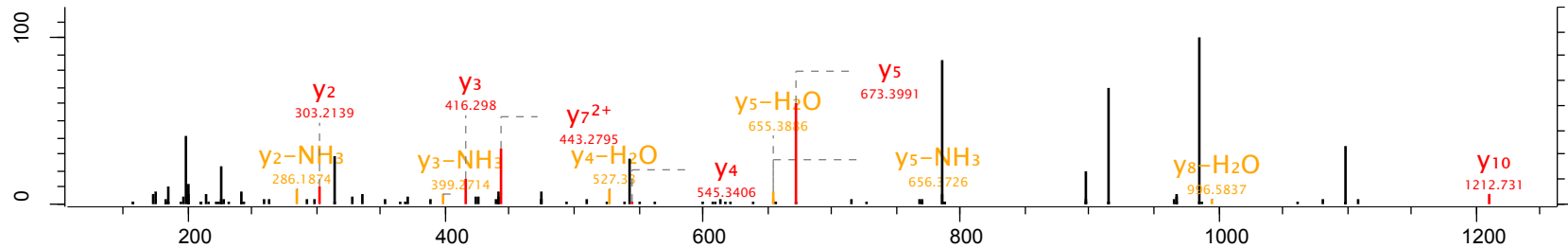
57.53

m/z

442.61

Gene names

FAM150B



- L V V E L V Q E L R K -

Red brackets above the sequence indicate the following y-series: y₁₀ (above V), y_{7²⁺} (above L), y₅ (above Q), y₄ (above E), y₃ (above L), and y₂ (above R).

Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

29908

Method

TOF; CID

Score

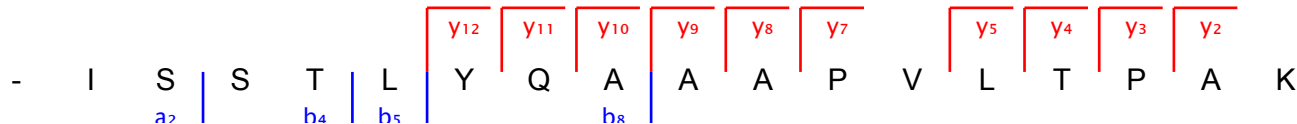
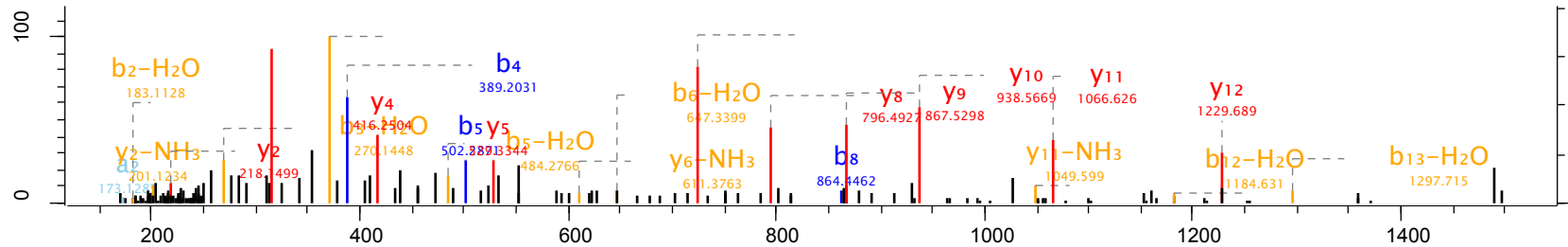
64.45

m/z

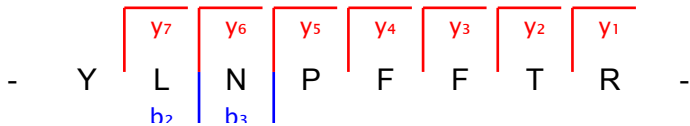
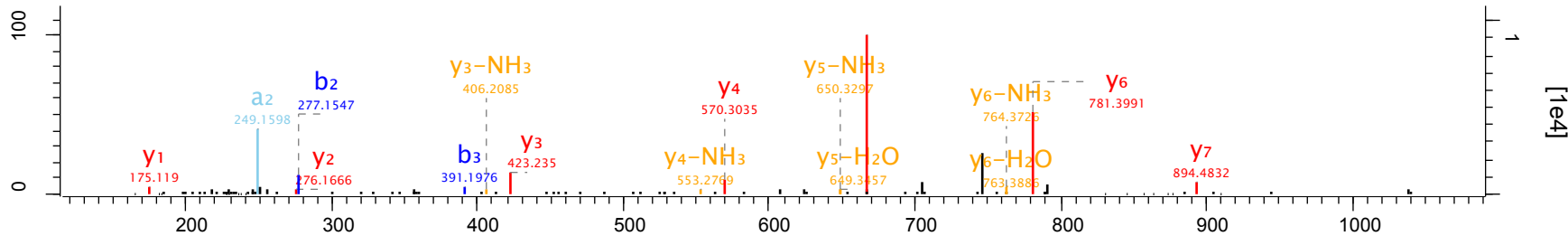
865.99

Gene names

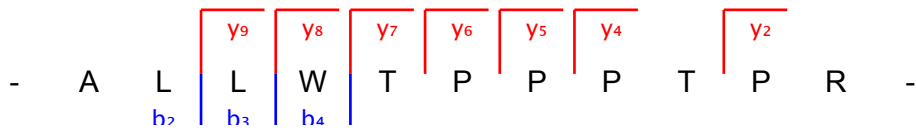
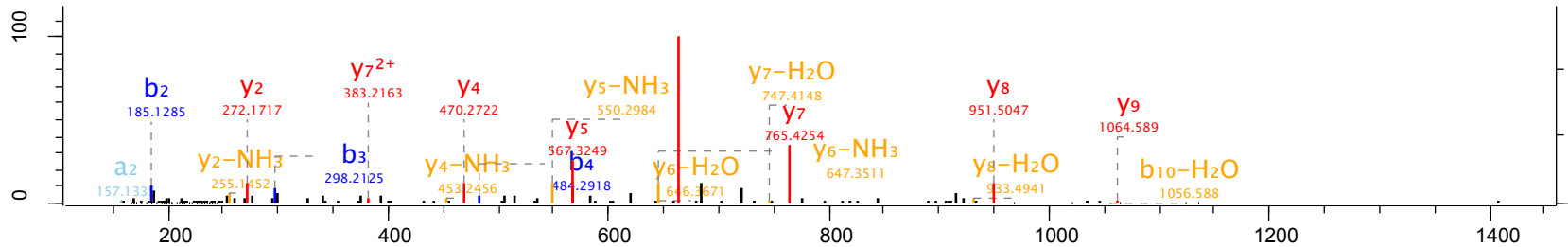
KRTCAP2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract9_dyn_5ul_F1_01_588	30140	TOF; CID	111.95	529.28	SLC9A8



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract9_dyn_5ul_F1_01_588	30511	TOF; CID	108.43	624.86	RITA1



[1e4]

Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

31463

Method

TOF; CID

Score

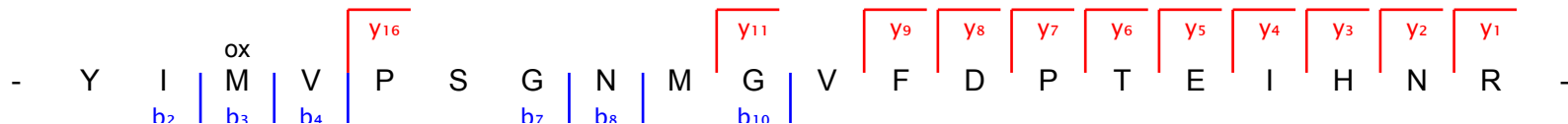
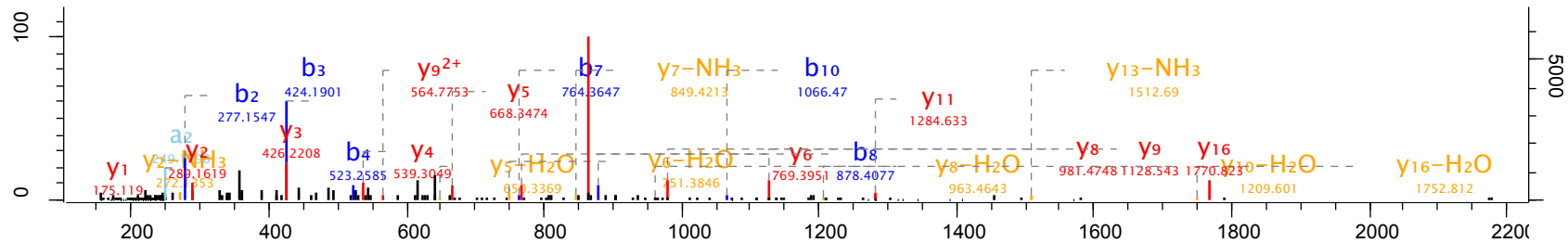
86.8

m/z

765.36

Gene names

CNIH4



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

31846

Method

TOF; CID

Score

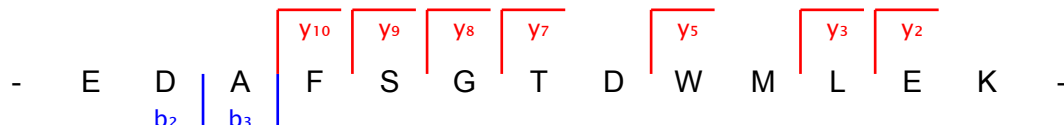
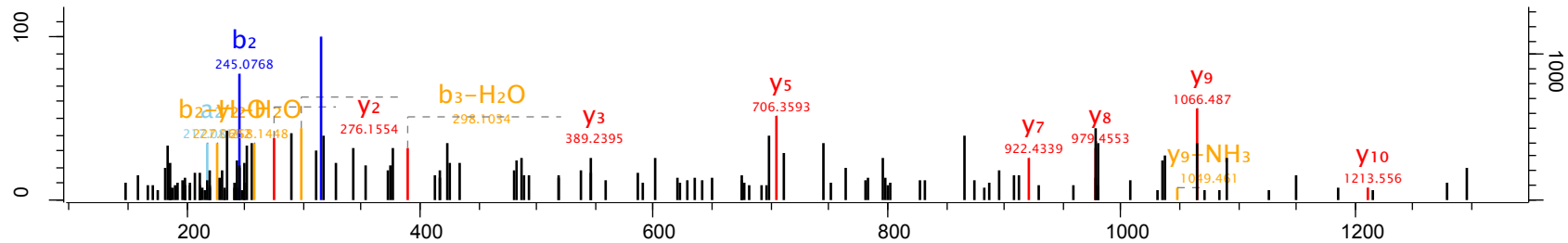
50.9

m/z

764.84

Gene names

ATF4



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

32555

Method

TOF; CID

Score

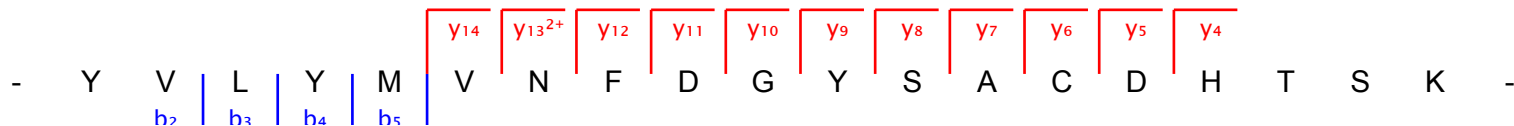
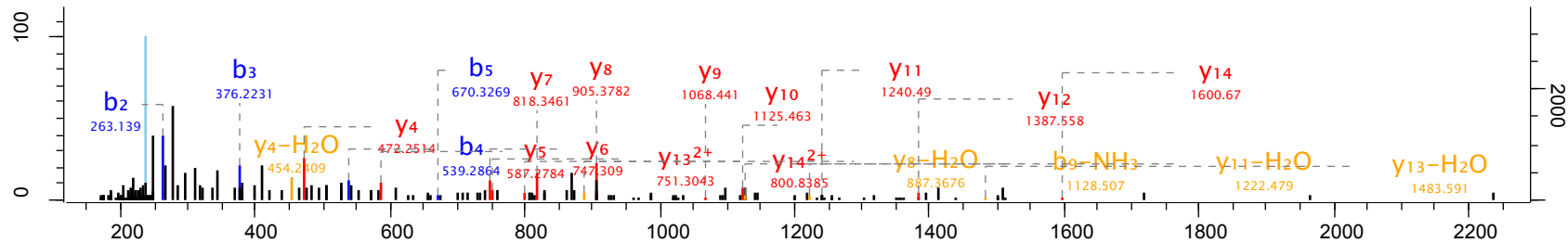
53.45

m/z

757.67

Gene names

EFNA5



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

36625

Method

TOF; CID

Score

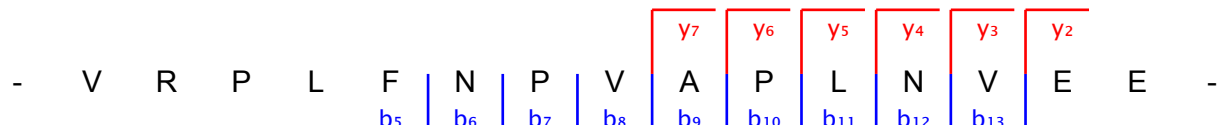
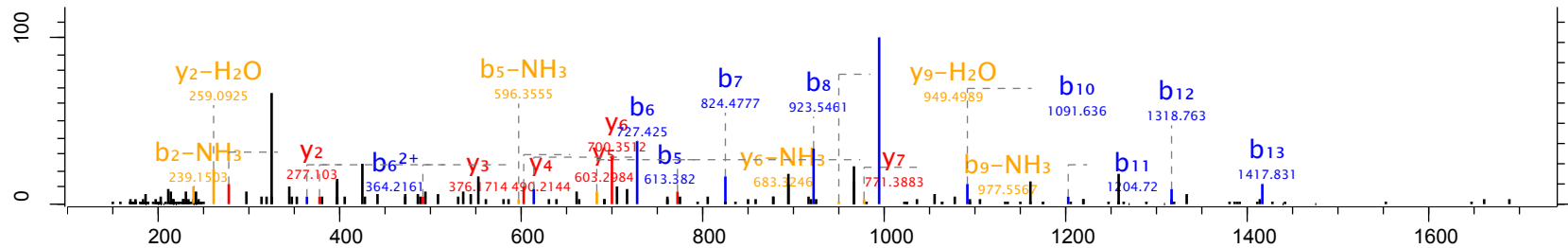
70.26

m/z

847.47

Gene names

MTERF2



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

37674

Method

TOF; CID

Score

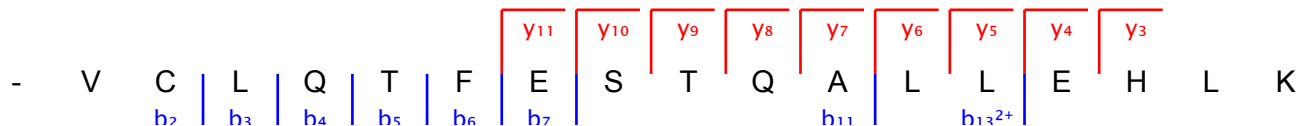
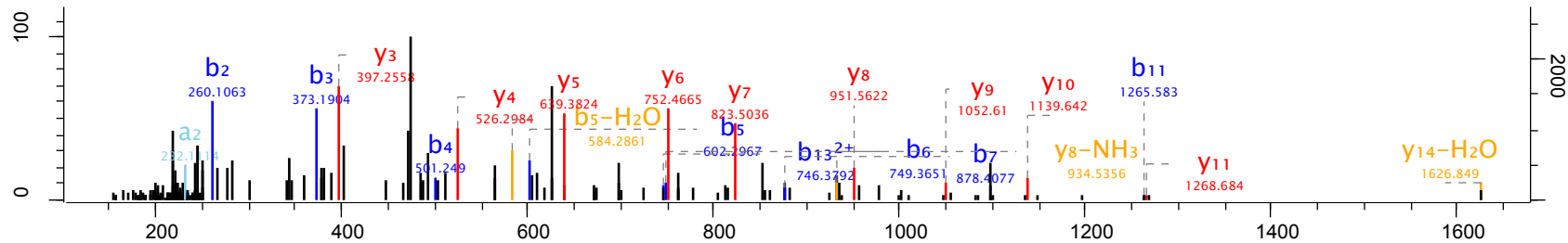
57.17

m/z

673.35

Gene names

PLAGL2



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

38231

Method

TOF; CID

Score

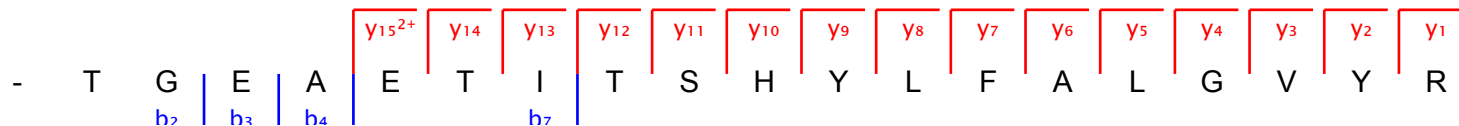
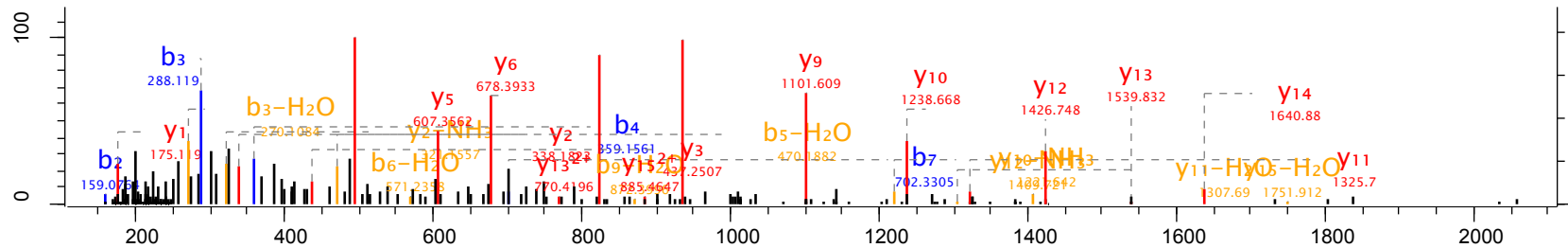
84.94

m/z

710.36

Gene names

KDELR1



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

38402

Method

TOF; CID

Score

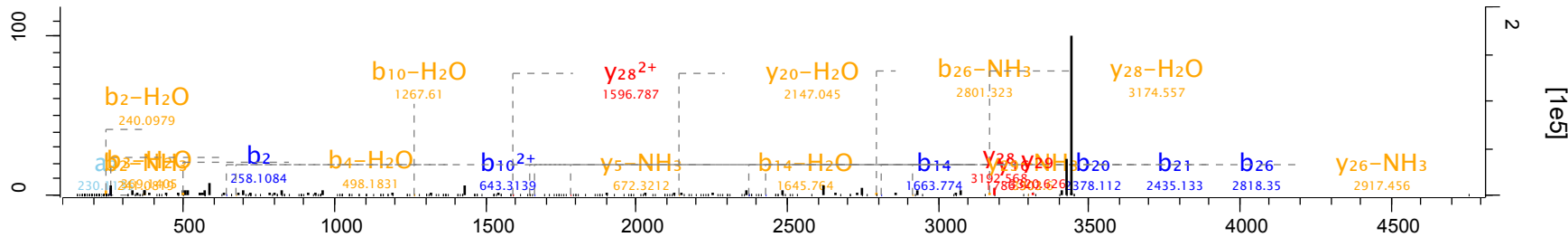
34.04

m/z

1150.89

Gene names

ZAN



Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

38564

Method

TOF; CID

Score

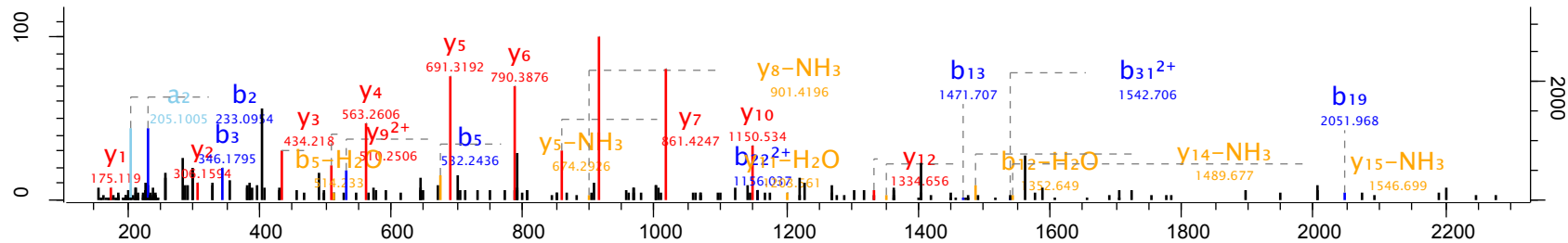
31.29

m/z

969.7

Gene names

CDC42SE1



- T M I G E P M N F V H L T H I G S G E M G

b₂ | b₃ | b₅ | b₁₃ | b₁₉

Raw file

20141014_fract9_dyn_5ul_F1_01_588

Scan

39519

Method

TOF; CID

Score

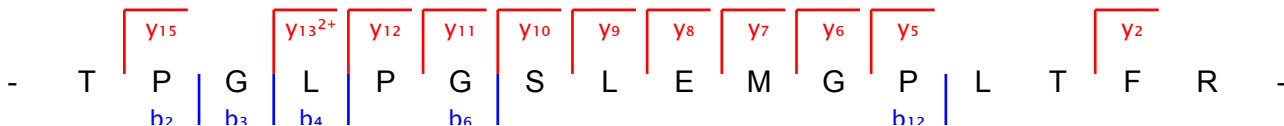
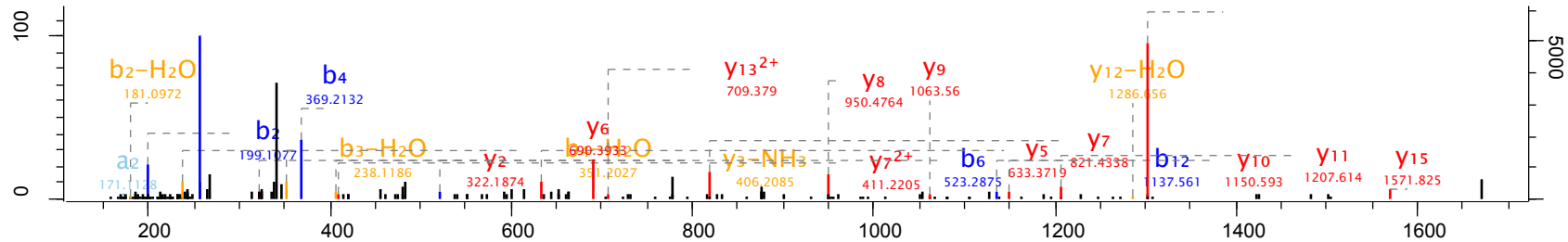
75.02

m/z

836.94

Gene names

ZNF273



Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

4168

Method

TOF; CID

Score

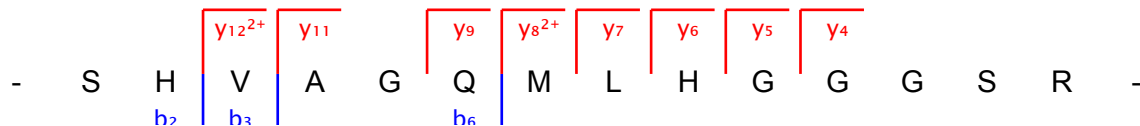
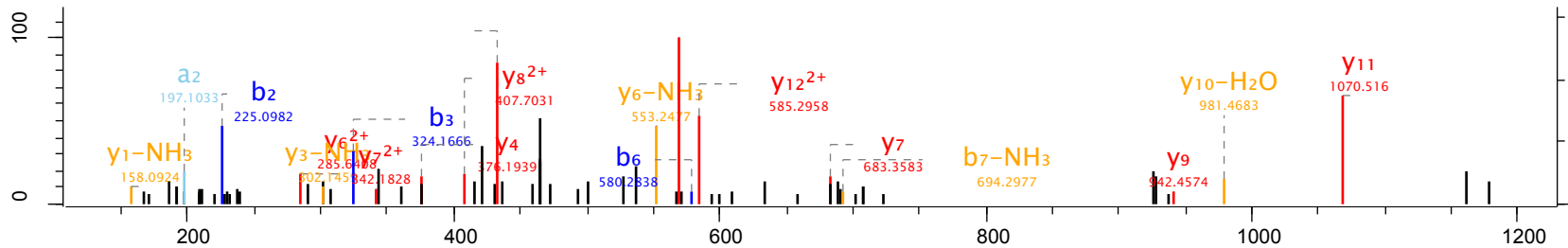
64.1

m/z

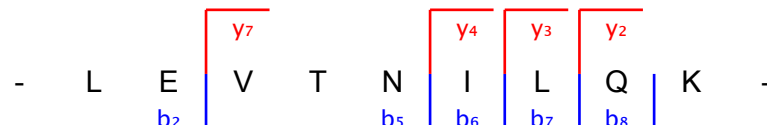
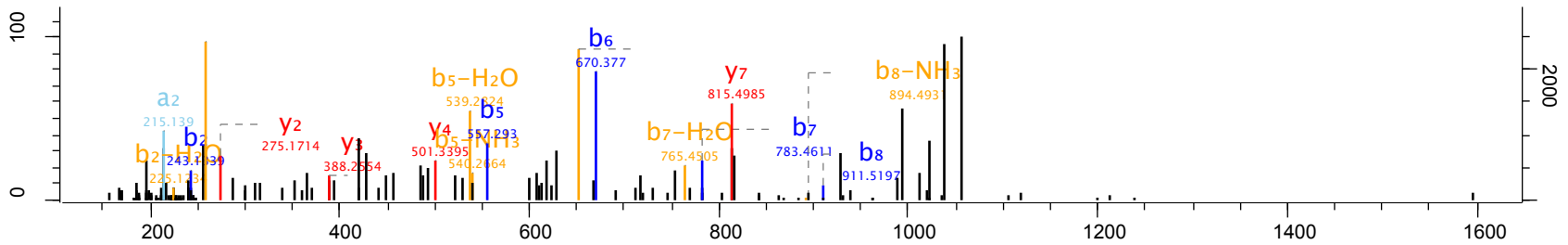
465.23

Gene names

CRAT



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract10_dyn_5ul_F2_01_589	11667	TOF; CID	91.09	529.31	RPGRIP1



Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

16187

Method

TOF; CID

Score

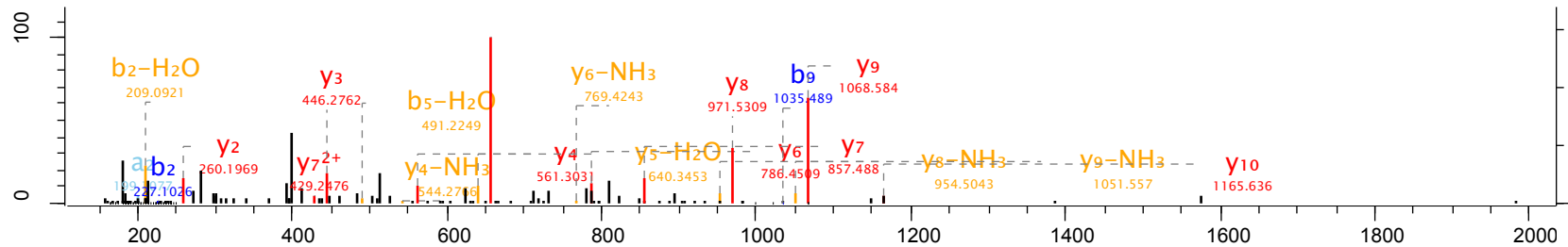
86.46

m/z

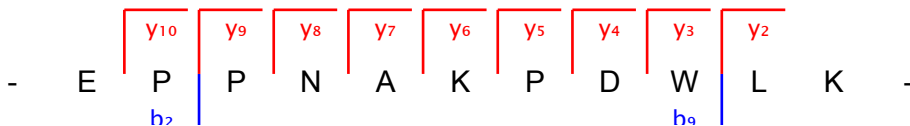
432.9

Gene names

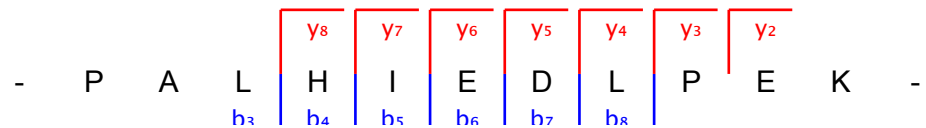
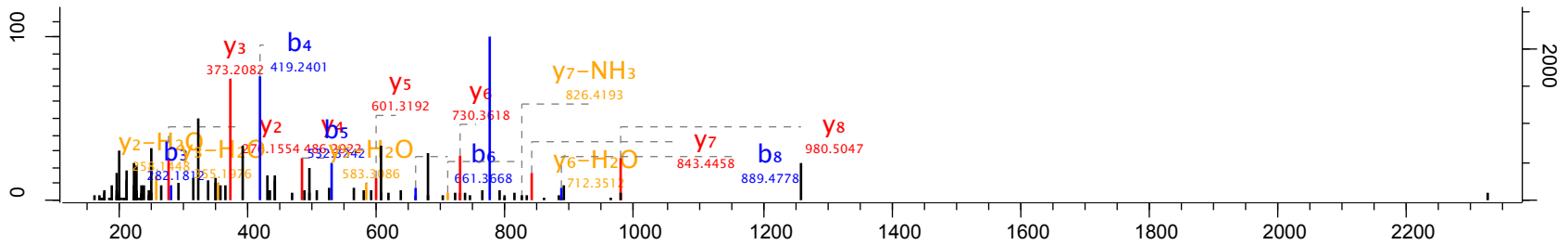
NDUFC1



[1e4]



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract10_dyn_5ul_F2_01_589	19145	TOF; CID	75.82	421.23	GNG11



Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

22968

Method

TOF; CID

Score

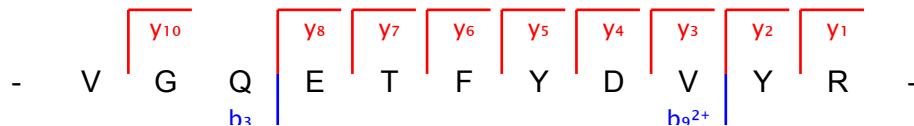
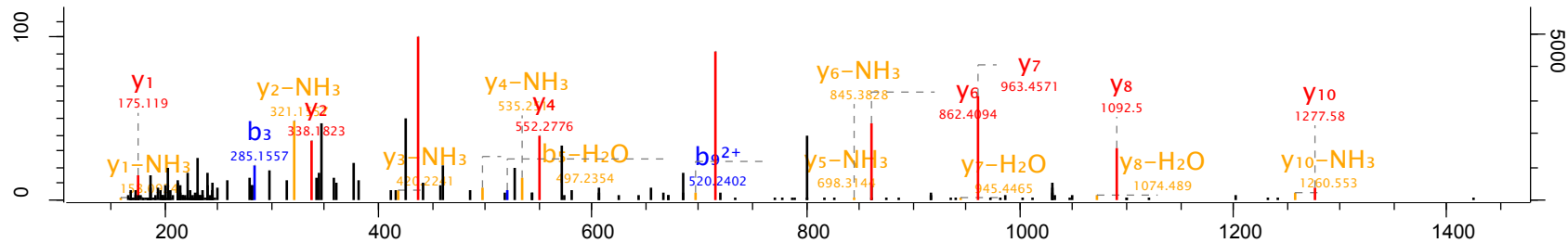
91.7

m/z

688.83

Gene names

SMIM4



Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

23140

Method

TOF; CID

Score

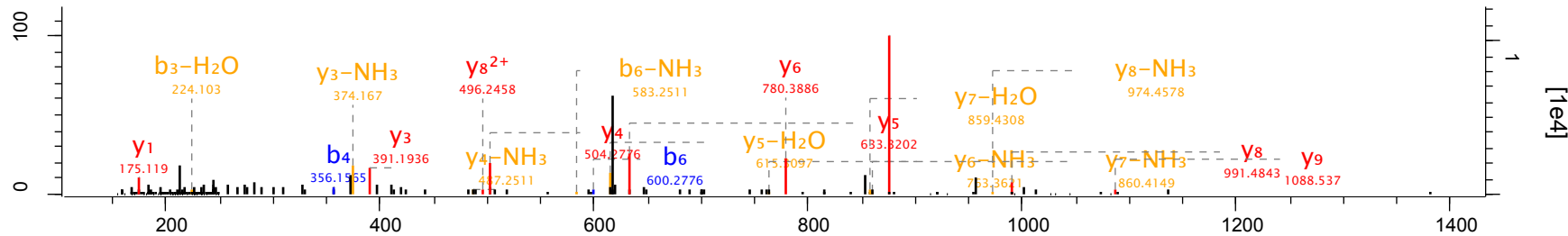
71.08

m/z

617.3

Gene names

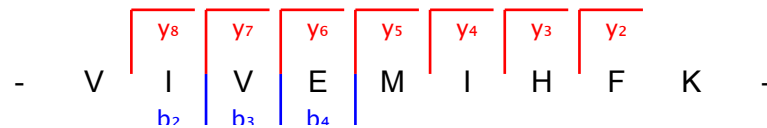
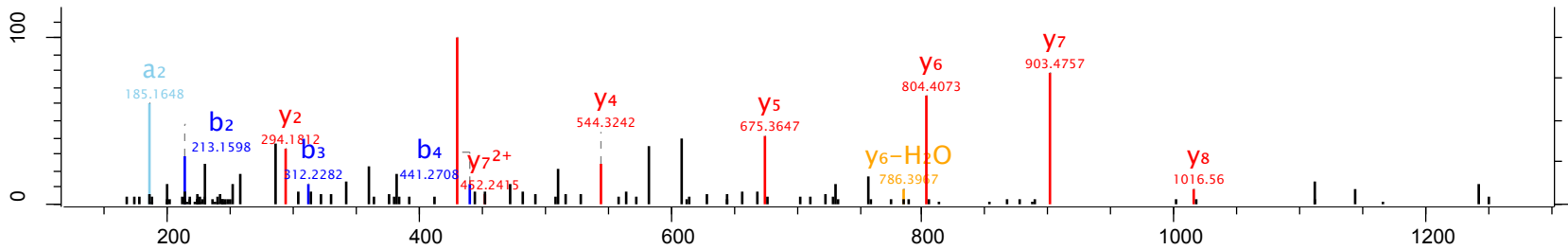
TMEM104



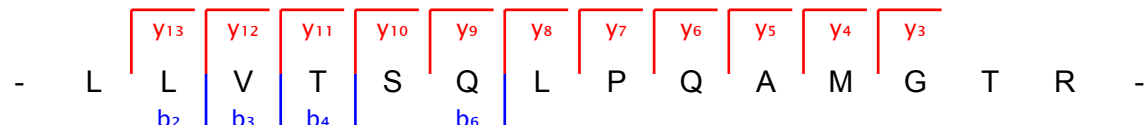
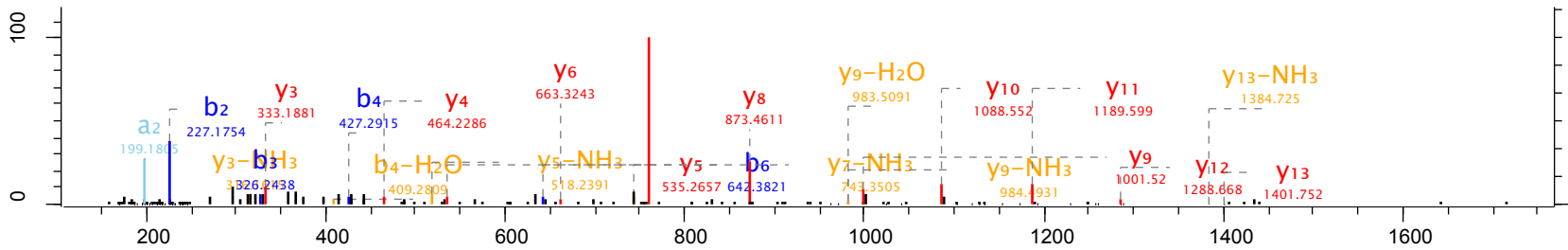
- G S P N P F E I T D R -

Fragmentation sites are indicated by brackets above the sequence, labeled y1 through y9. The b4 and b6 labels are positioned below the N and F residues, respectively.

Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract10_dyn_5ul_F2_01_589	26607	TOF; CID	74.92	558.32	STC2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract10_dyn_5ul_F2_01_589	27475	TOF; CID	80.51	757.92	C9orf91



Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

27642

Method

TOF; CID

Score

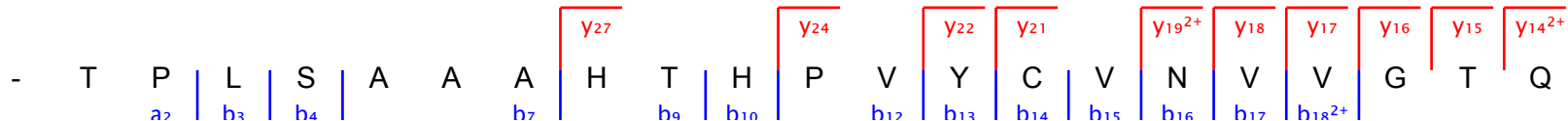
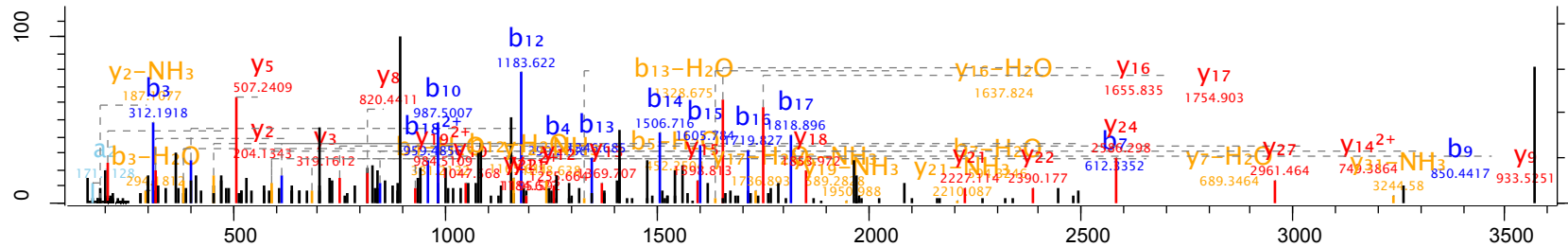
118.22

m/z

894.45

Gene names

DYNC111



Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

32218

Method

TOF; CID

Score

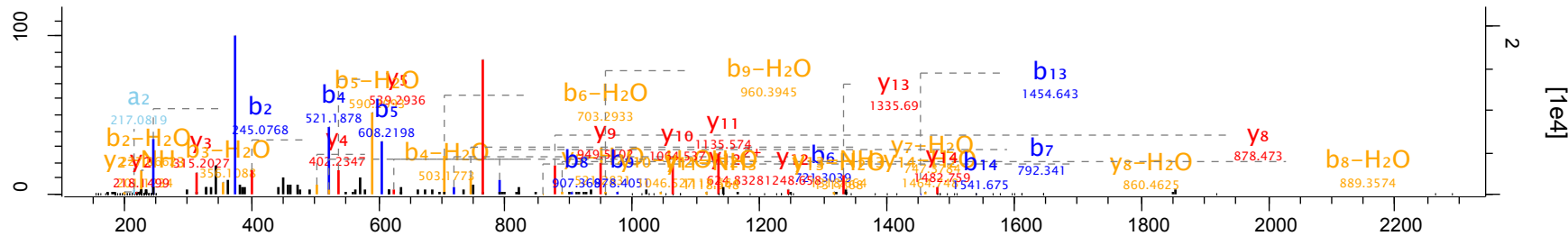
170.77

m/z

928.44

Gene names

MAPK1IP1L



ac

S

D

E

F

S

L

A

D

A

L

P

E

H

S

P

A

K

-

b₂b₃b₄b₅b₆b₇b₈b₉b₁₃b₁₄y₁₄y₁₃y₁₂y₁₁y₁₀y₉y₈y₇y₅y₄y₃y₂

Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

33291

Method

TOF; CID

Score

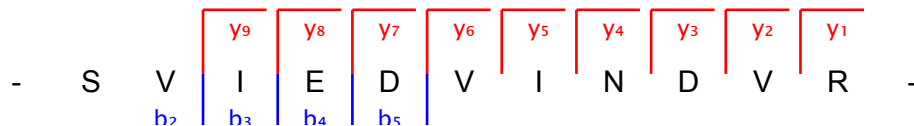
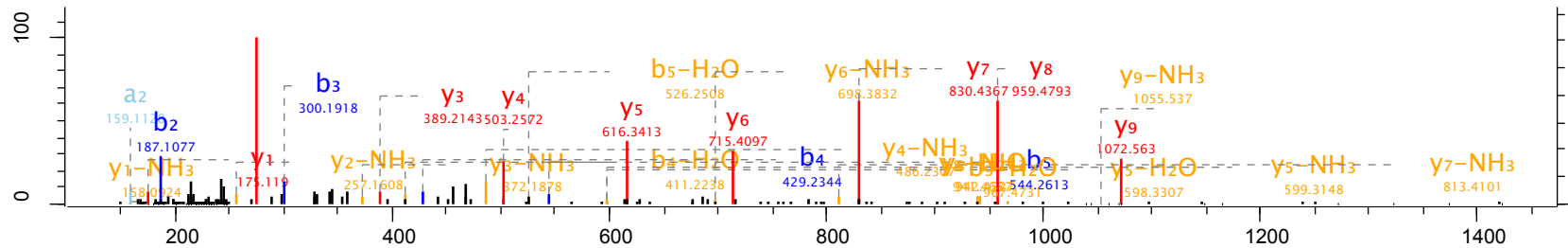
140.48

m/z

629.84

Gene names

GTF2A1; HIST1H4F



Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

35403

Method

TOF; CID

Score

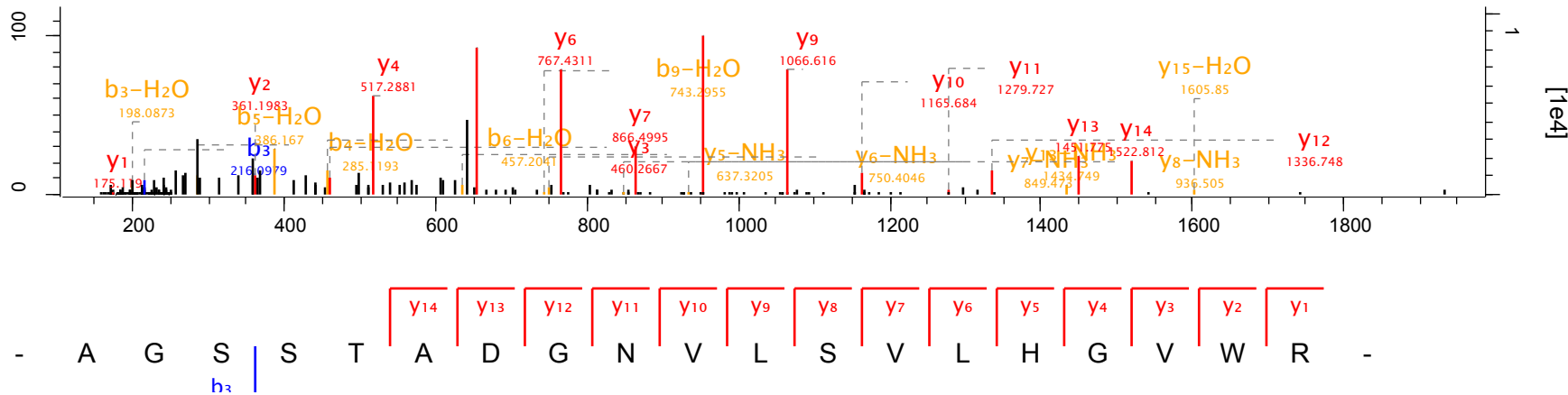
68.24

m/z

642.67

Gene names

SLC25A26



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract10_dyn_5ul_F2_01_589

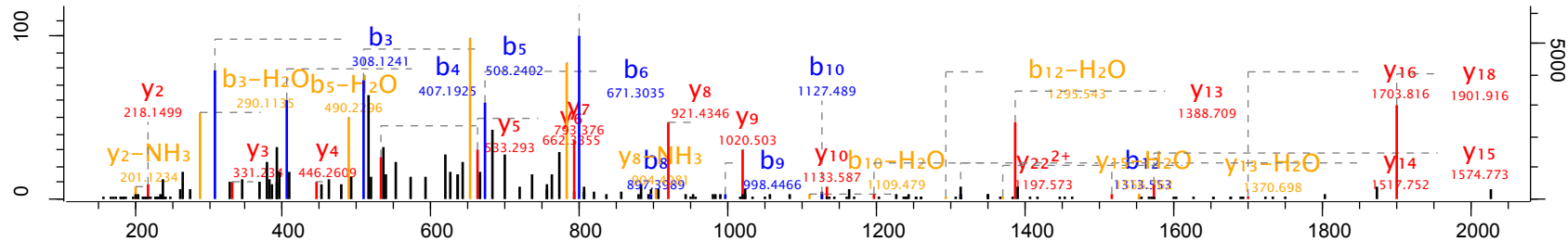
37241

TOF; CID

118.37

1351.63

SMAGP



- G S Y V T Y E P T E G E P S A I V Q M E S
 b₃ b₄ b₅ b₆ b₇ b₈ b₉ b₁₀ b₁₂ y₂₂²⁺ y₁₈ y₁₆ y₁₅ y₁₄ y₁₃ y₁₀ y₉ y₈ y₇ y₆ y₅

Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

40719

Method

TOF; CID

Score

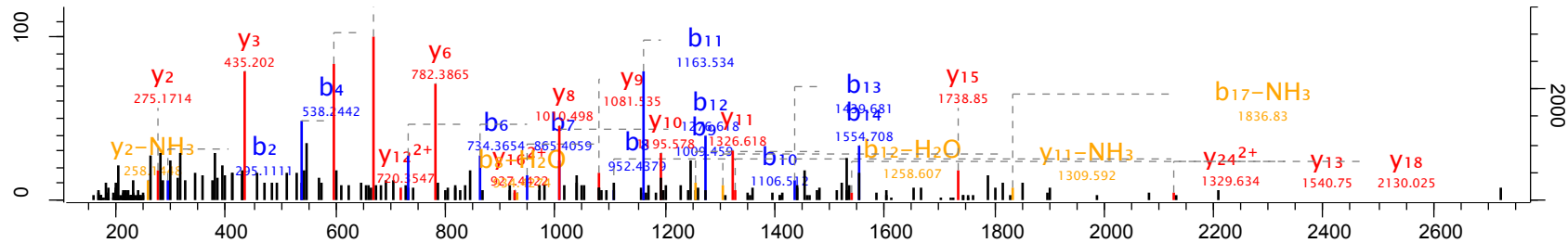
71.67

m/z

1098.52

Gene names

CNIH1



- Y M | S R | P V | M | S | G | P | G L | Y D | P T T | I M | N A
 b₂ | b₄ | b₆ | b₇ | b₈ | b₉ | b₁₀ | b₁₁ | b₁₂ | b₁₃ | b₁₄

y₂₄²⁺ y₁₈ y₁₆²⁺ y₁₅ y₁₃ y₁₂²⁺ y₁₁ y₁₀ y₉

Raw file

20141014_fract10_dyn_5ul_F2_01_589

Scan

41167

Method

TOF; CID

Score

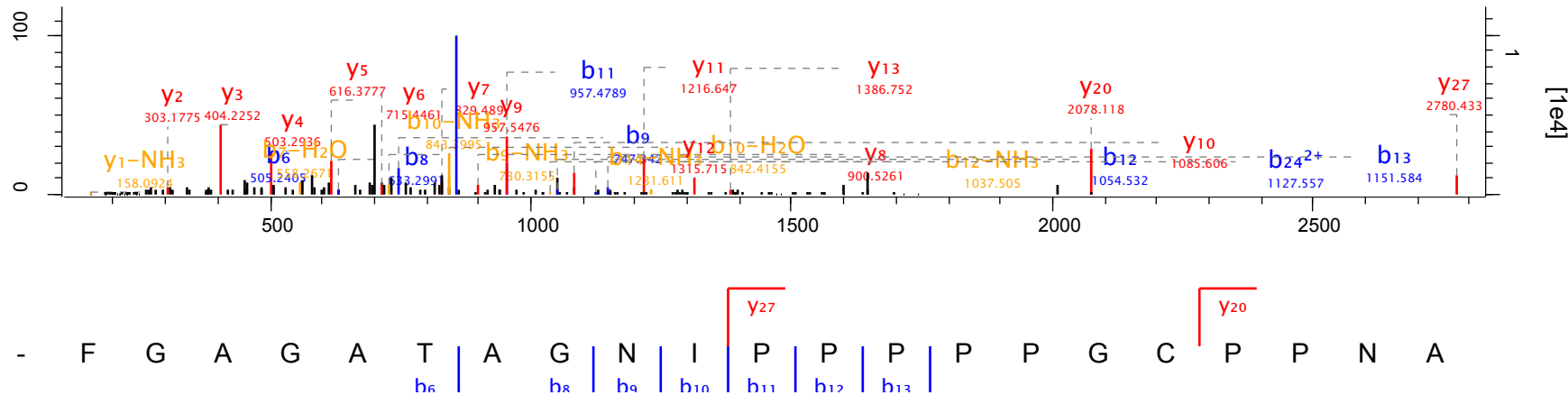
36.55

m/z

1214.62

Gene names

DAZAP2



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

6157

Method

TOF; CID

Score

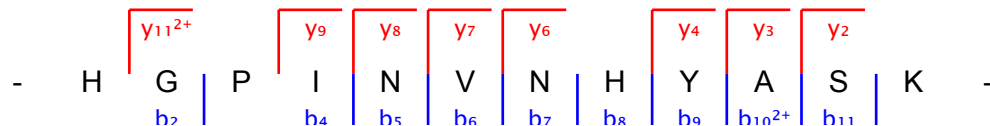
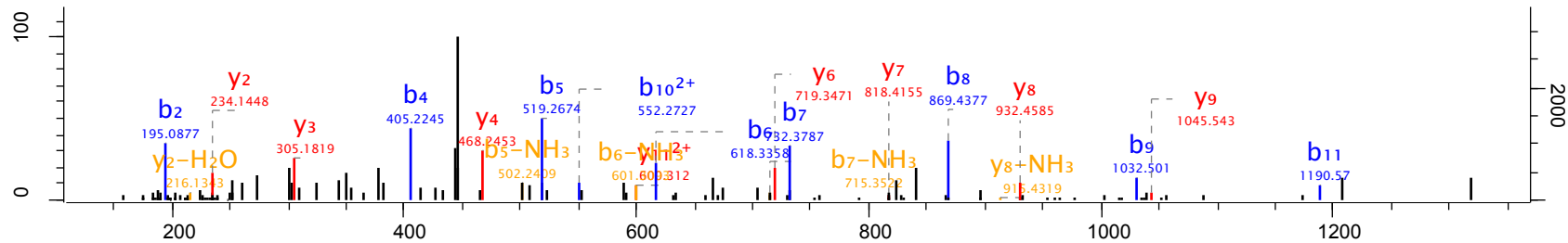
101.56

m/z

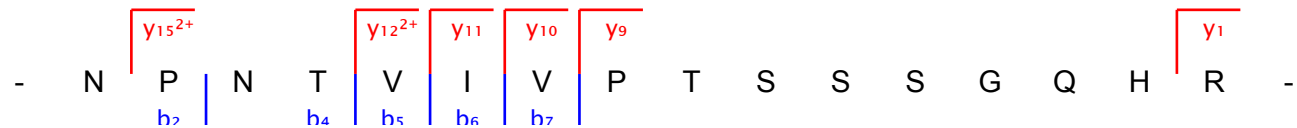
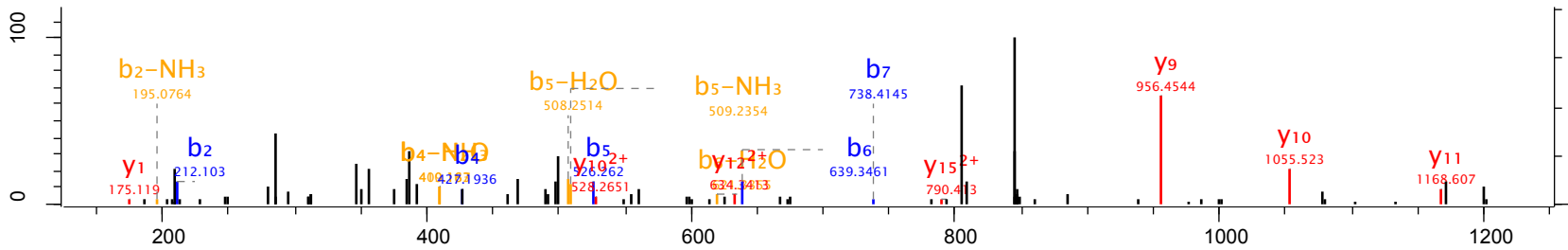
446.23

Gene names

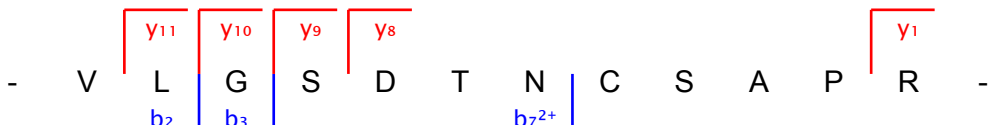
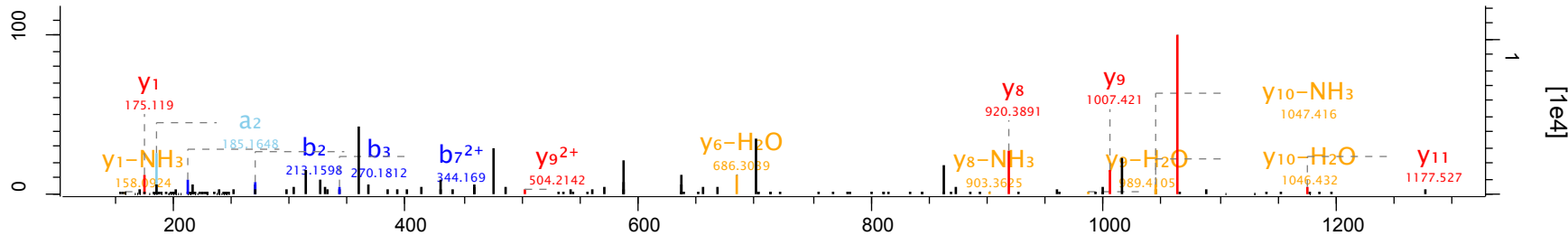
NINJ1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	10827	TOF; CID	43.04	847.43	C11orf49



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	11461	TOF; CID	54.34	638.8	PNMA3



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

12591

Method

TOF; CID

Score

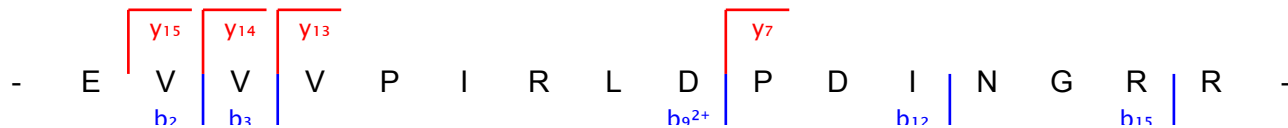
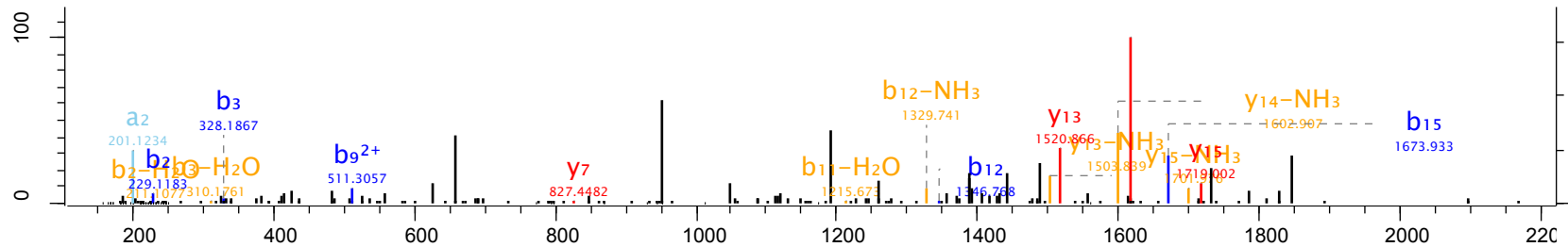
58.89

m/z

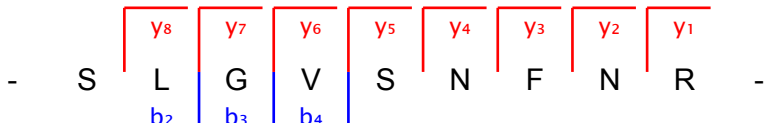
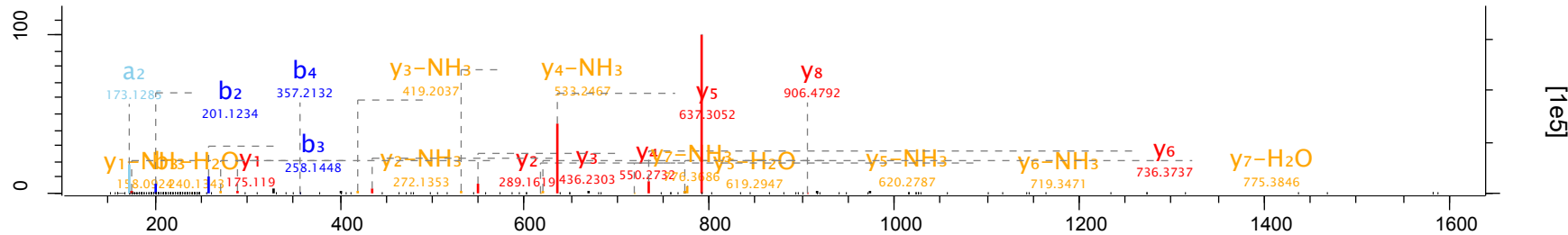
463.02

Gene names

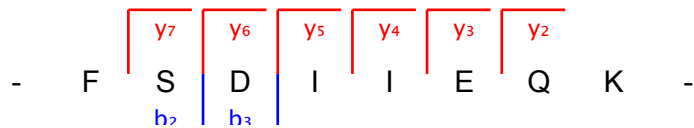
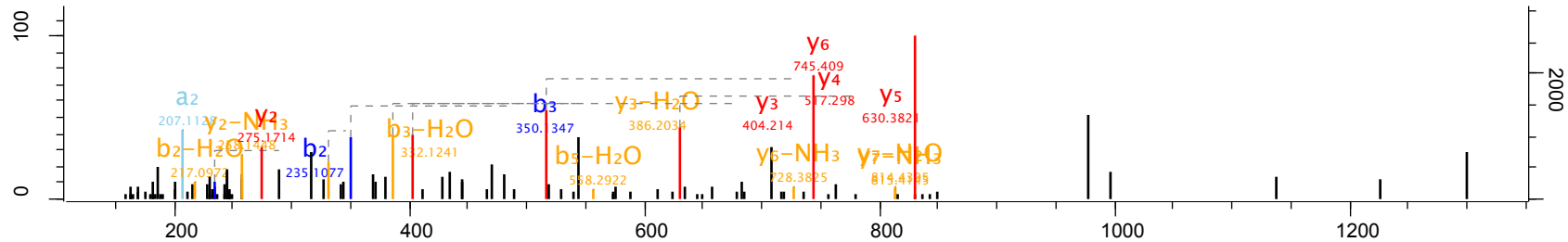
ADAMTS15



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	14078	TOF; CID	152.34	497.26	AKR1D1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	15252	TOF; CID	101.43	490.26	ZBED8



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

15492

Method

TOF; CID

Score

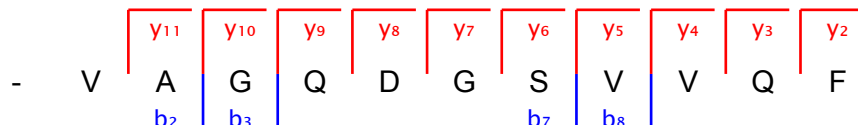
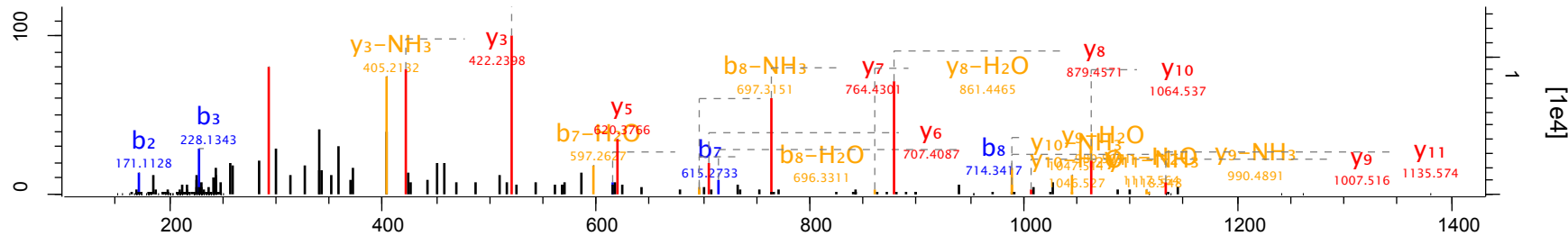
122.33

m/z

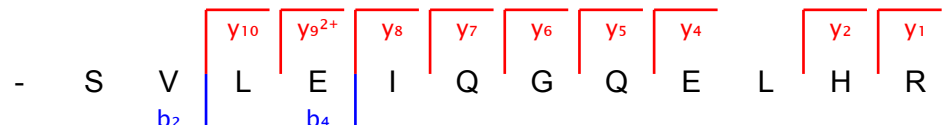
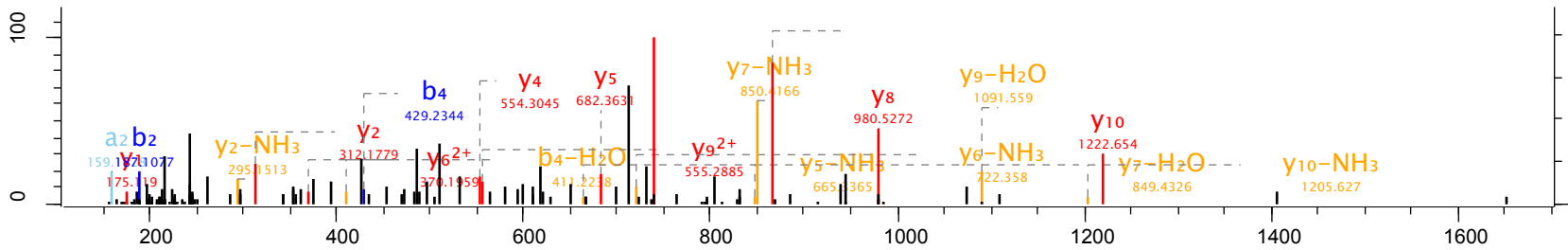
617.82

Gene names

SUMO2;SUMO3;SUMO4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	17495	TOF; CID	72.29	470.25	CCDC88B



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

17959

Method

TOF; CID

Score

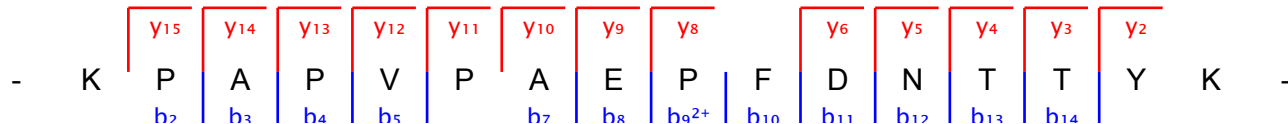
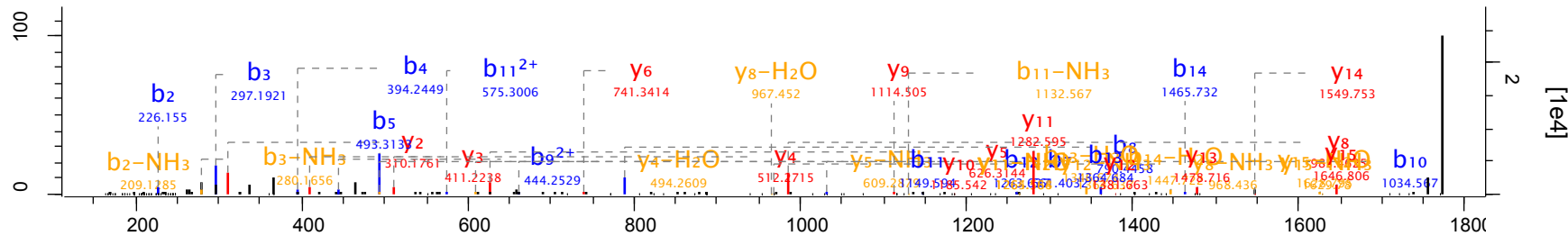
145.82

m/z

888.45

Gene names

NDUFV3



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

19980

Method

TOF; CID

Score

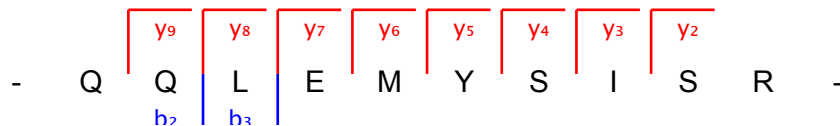
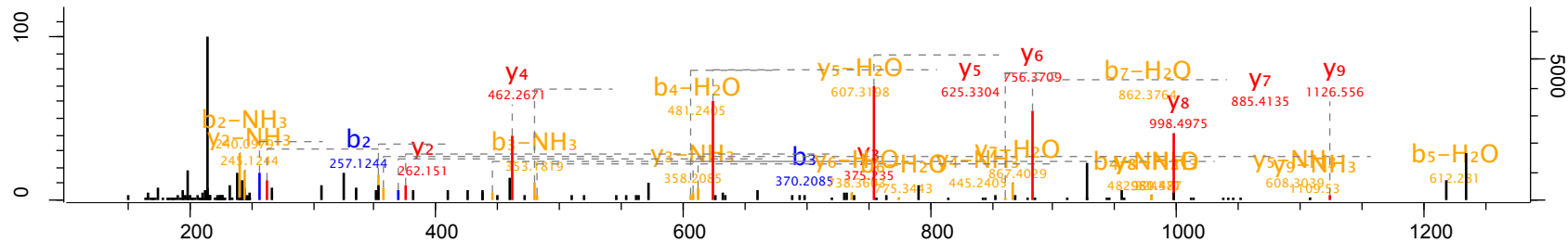
108.47

m/z

627.81

Gene names

SMIM19



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

21302

Method

TOF; CID

Score

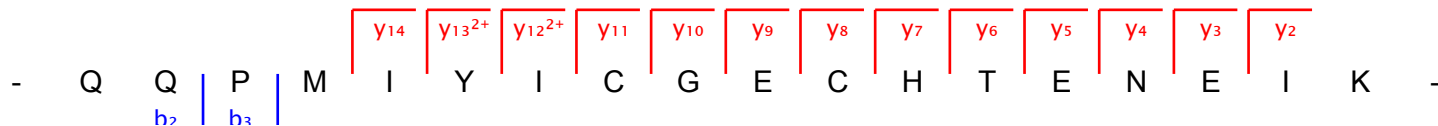
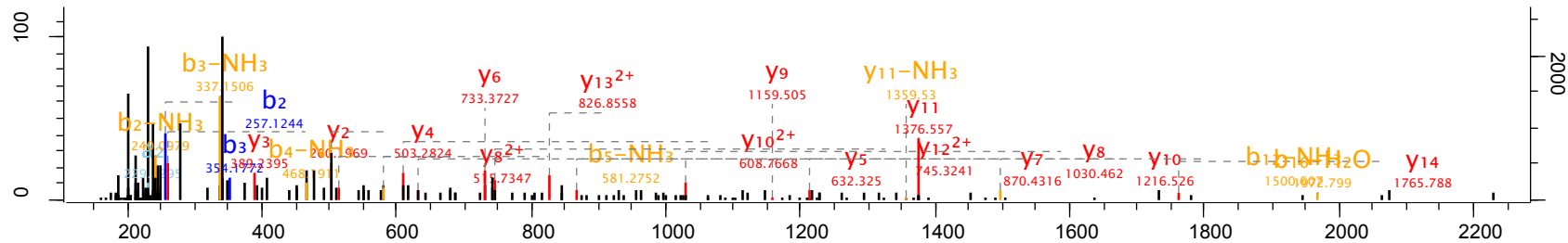
49.92

m/z

751

Gene names

POLR2K



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

24147

Method

TOF; CID

Score

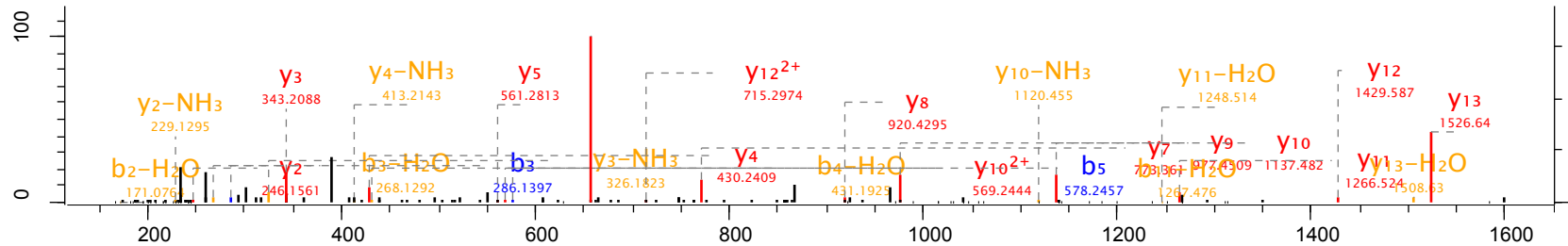
78.51

m/z

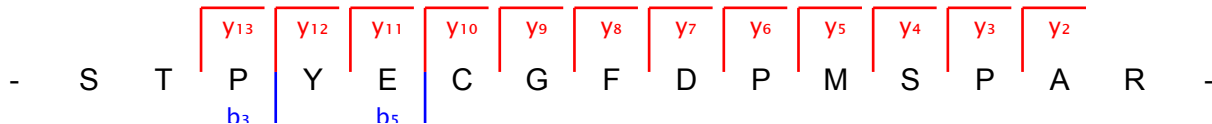
858.37

Gene names

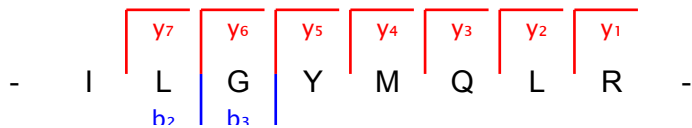
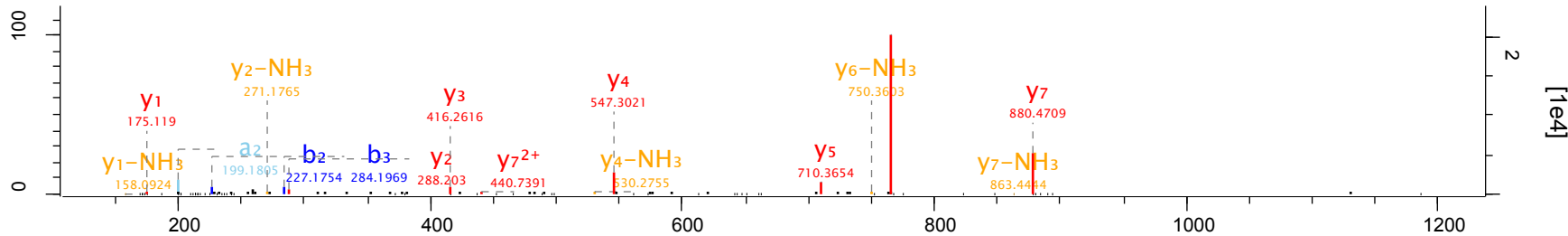
MT-ND3



[1e4]



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	24934	TOF; CID	107.32	497.28	MT-ND1



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

25502

Method

TOF; CID

Score

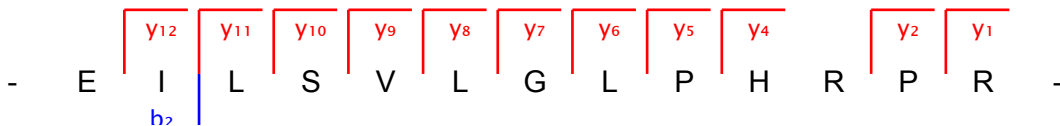
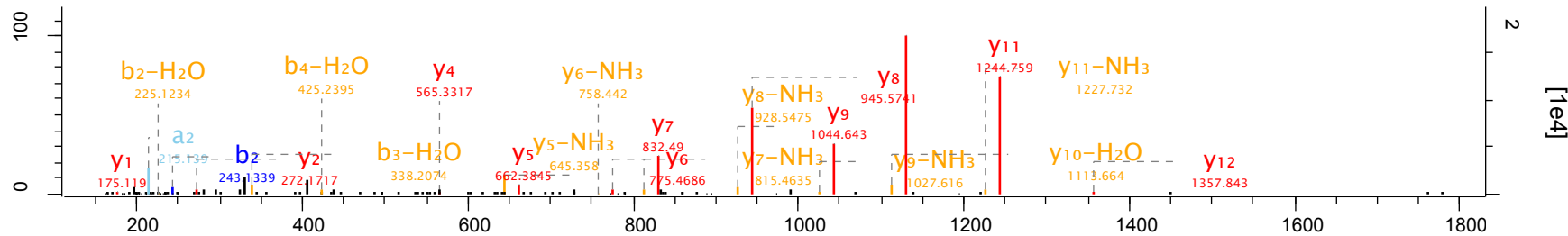
98.1

m/z

496.3

Gene names

BMP6



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

27185

Method

TOF; CID

Score

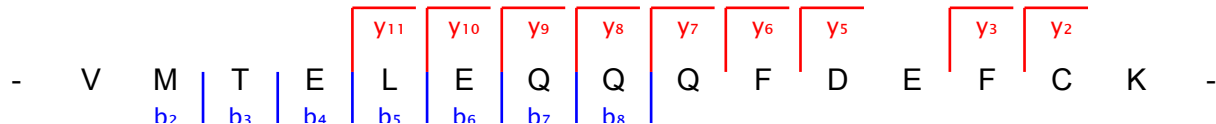
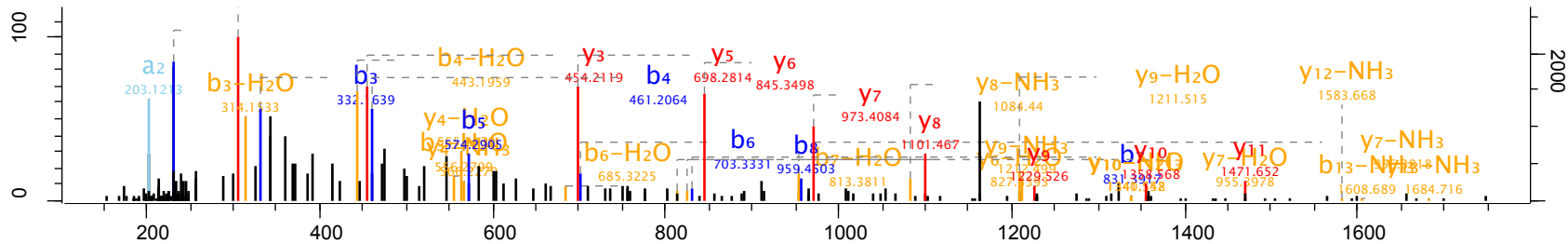
157.86

m/z

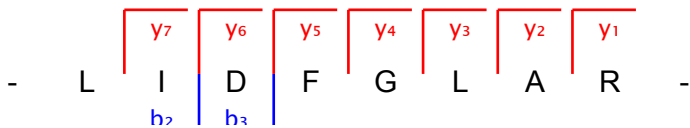
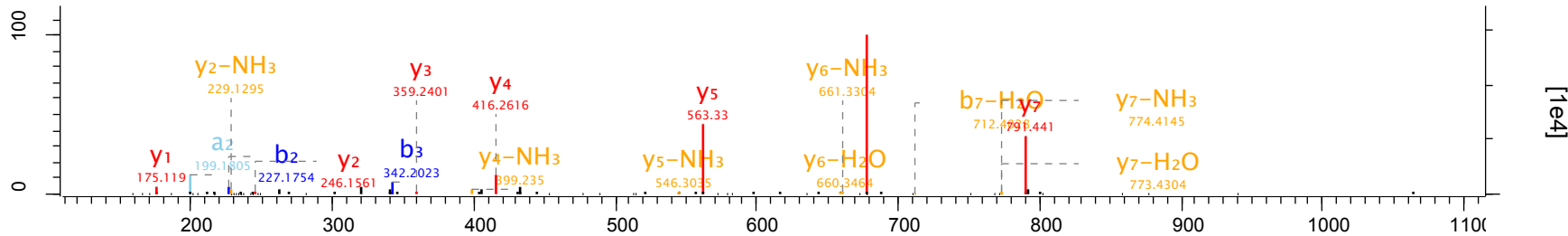
966.93

Gene names

CCDC23



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	28502	TOF; CID	120.46	452.77	MYLK



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

30360

Method

TOF; CID

Score

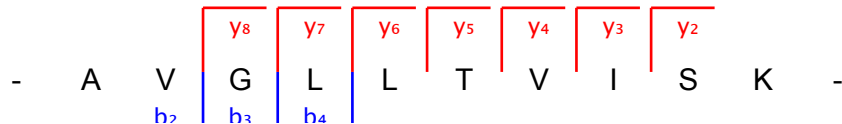
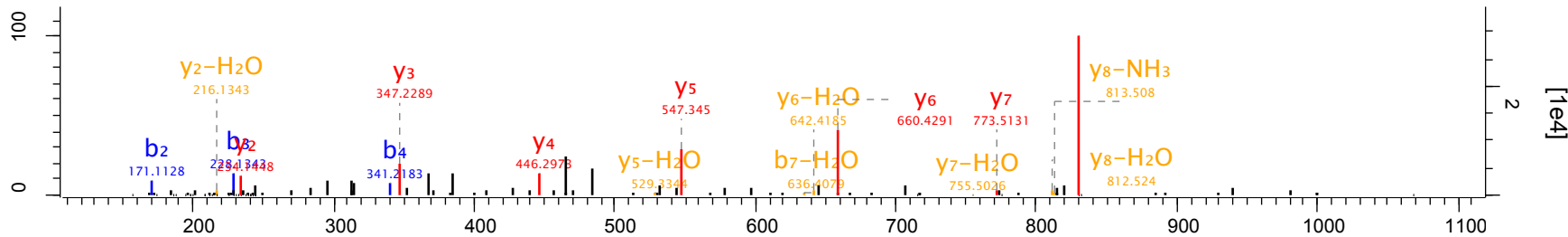
66.92

m/z

501.83

Gene names

PSCA



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

30862

Method

TOF; CID

Score

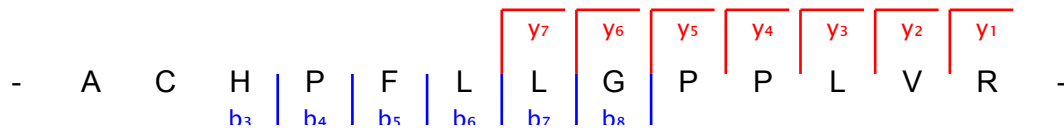
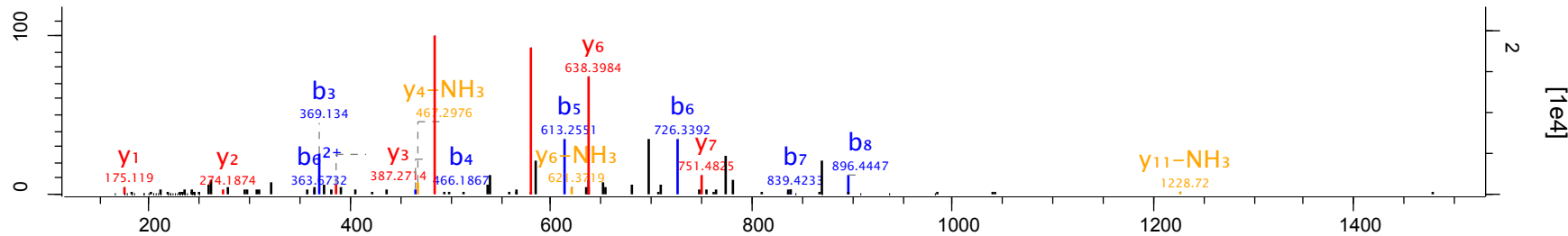
68.54

m/z

493.61

Gene names

TOR1AIP2



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

31303

Method

TOF; CID

Score

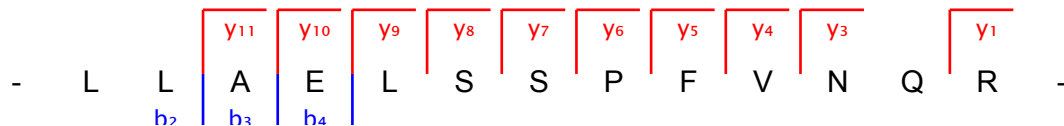
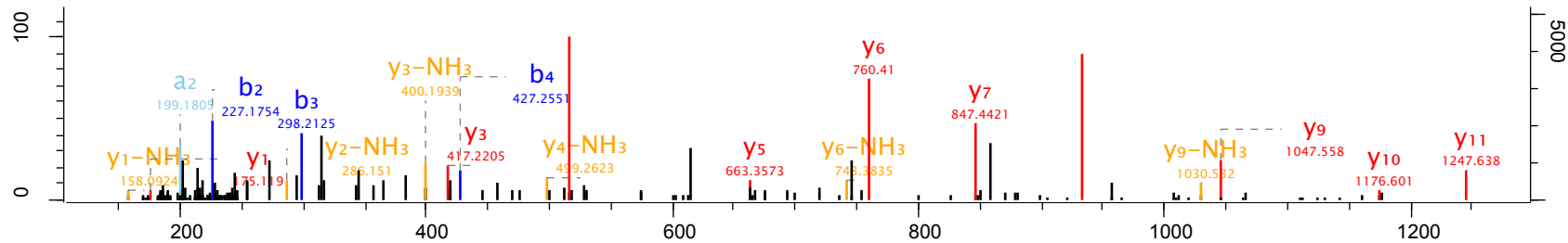
107.09

m/z

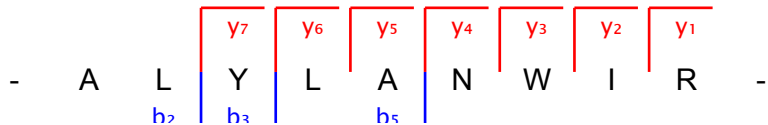
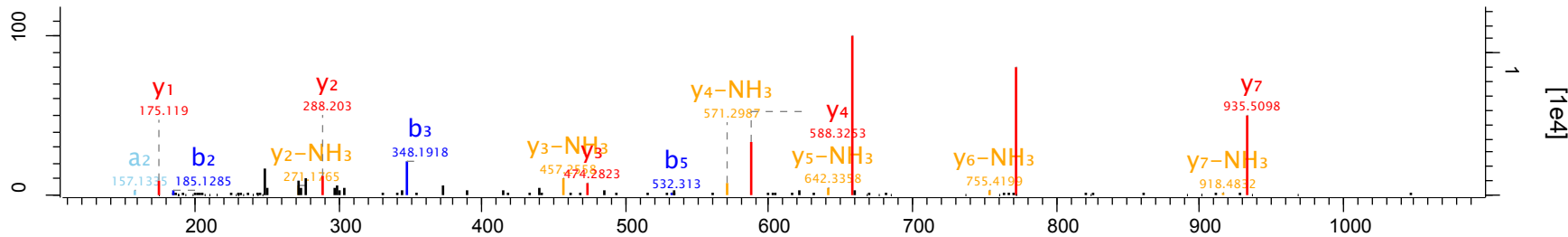
738.41

Gene names

TMEM56;TMEM56-RWDD3



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract11_dyn_5ul_F3_01_590	35180	TOF; CID	120.46	560.32	KDEL3



Raw file

20141014_fract11_dyn_5ul_F3_01_590

Scan

36573

Method

TOF; CID

Score

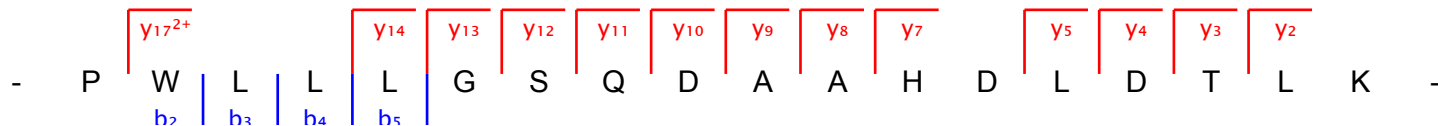
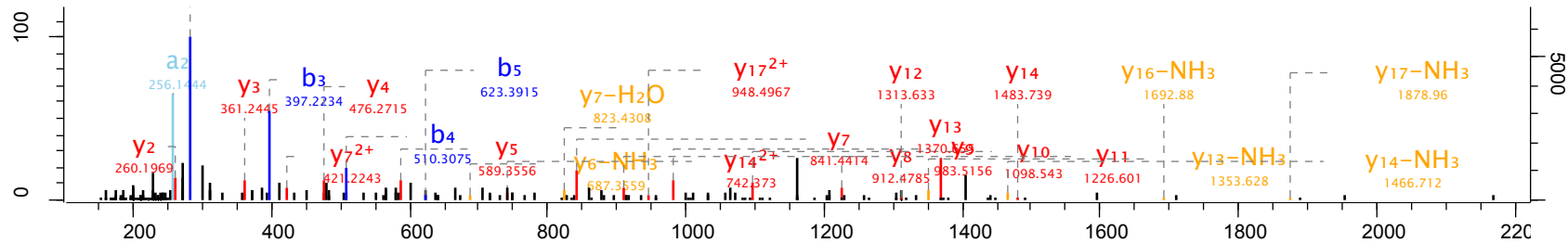
78.44

m/z

665.35

Gene names

DUSP19



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

5855

Method

TOF; CID

Score

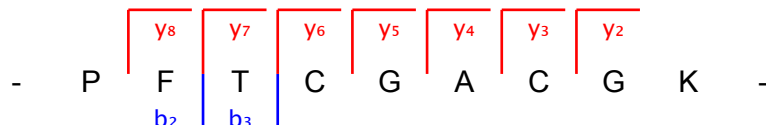
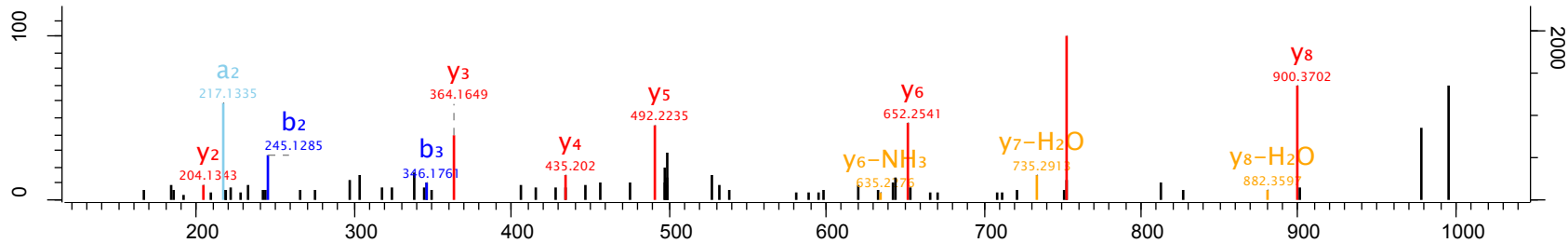
76.06

m/z

499.21

Gene names

ZNF580



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

7555

Method

TOF; CID

Score

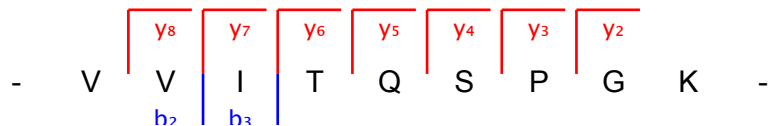
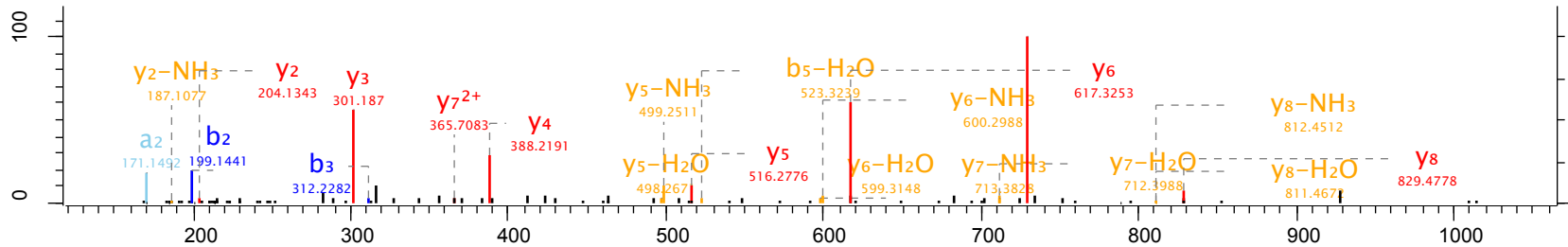
95.36

m/z

464.78

Gene names

TMEM185B;TMEM185A



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

7804

Method

TOF; CID

Score

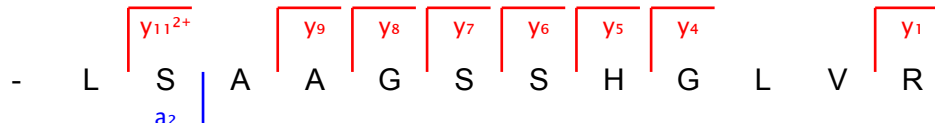
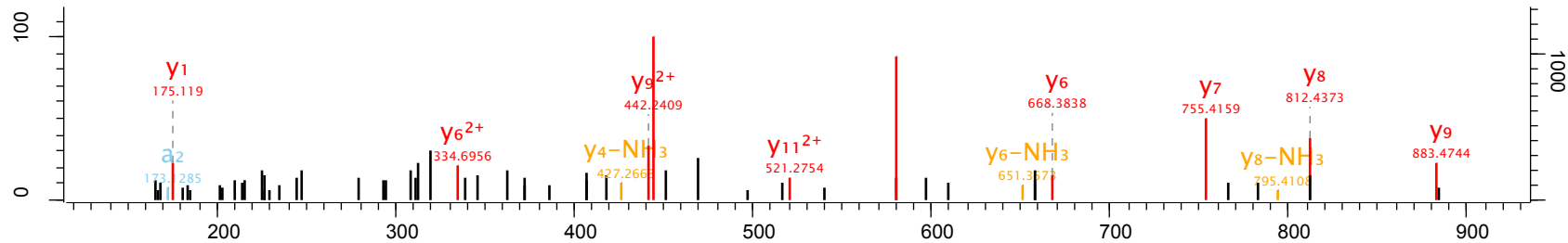
50.79

m/z

385.55

Gene names

TNRC6C



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract12_dyn_5ul_F4_01_591

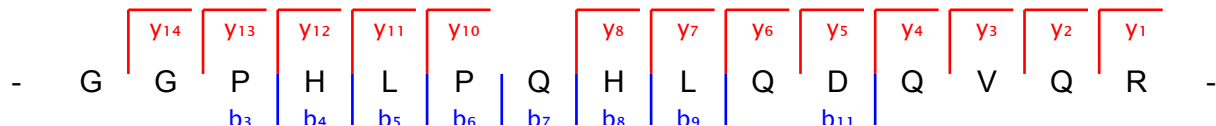
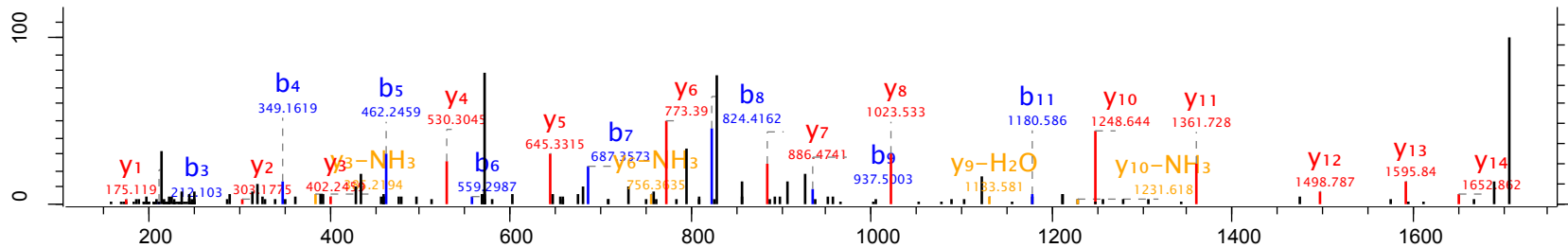
12205

TOF; CID

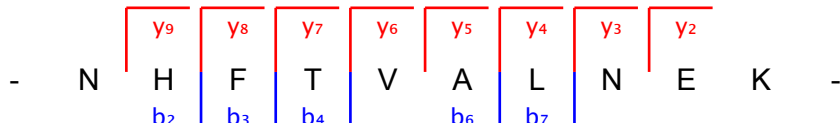
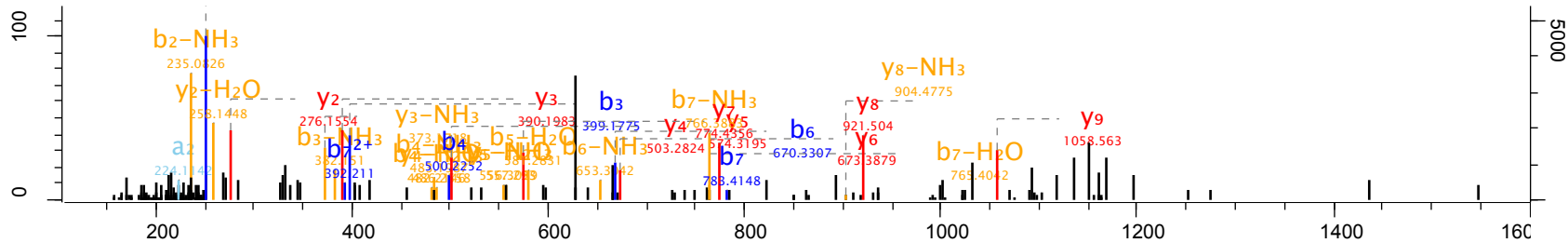
75.91

570.63

HPS1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	12895	TOF; CID	151.07	586.8	SULT1E1



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

14368

Method

TOF; CID

Score

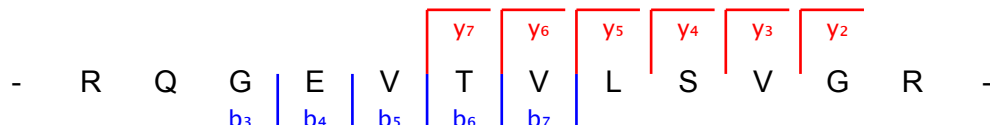
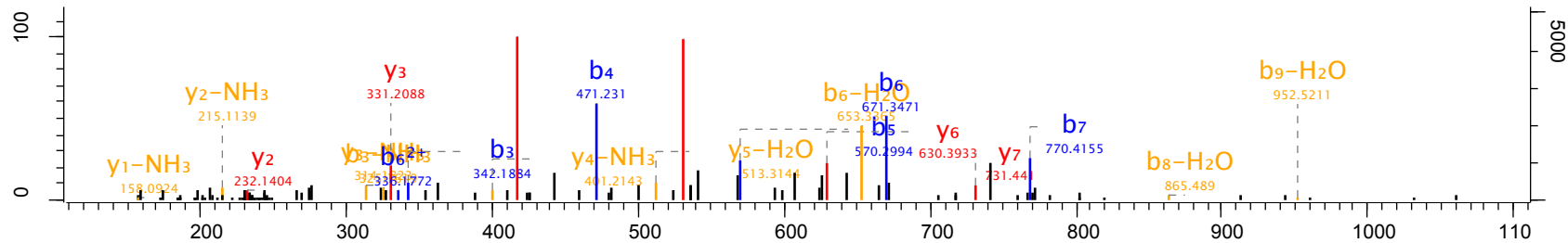
69.03

m/z

434.25

Gene names

RELL1



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

14370

Method

TOF; CID

Score

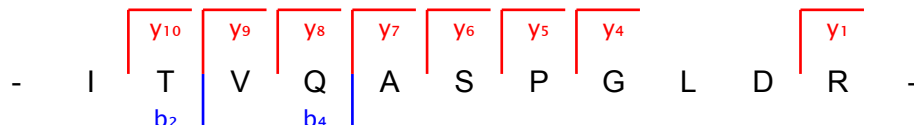
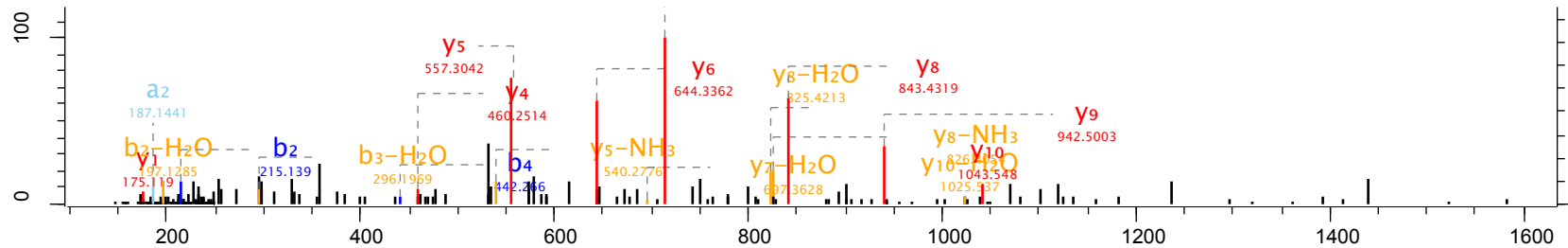
74.61

m/z

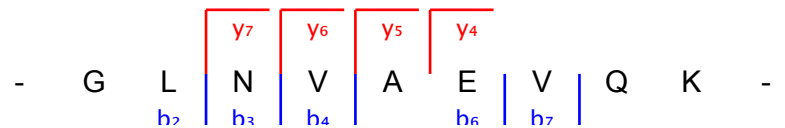
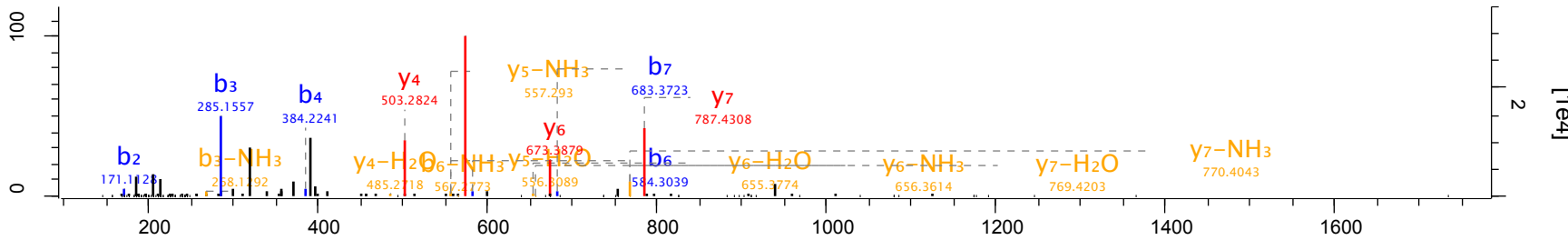
578.82

Gene names

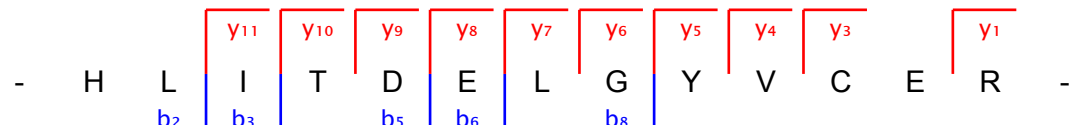
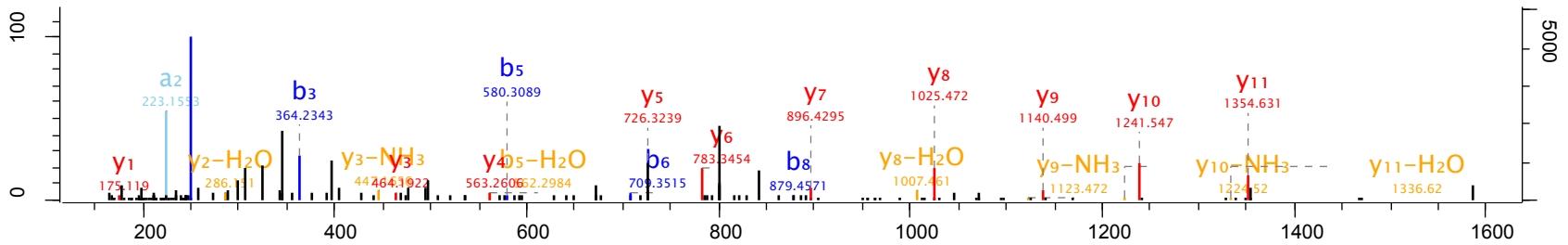
MAP3K11



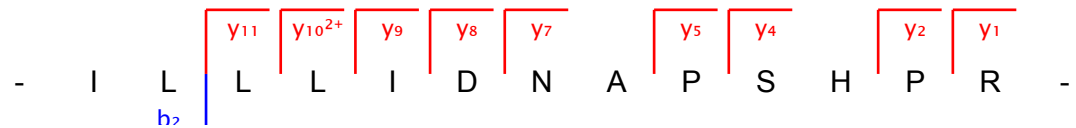
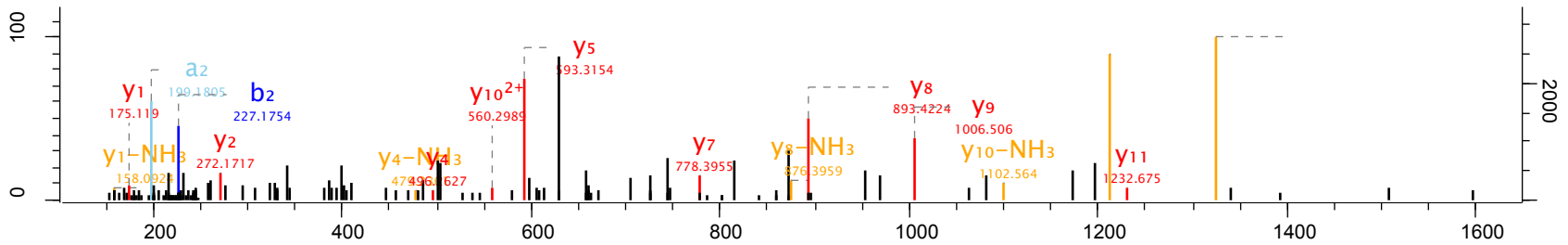
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	15999	TOF; CID	93.6	479.27	SLIT1



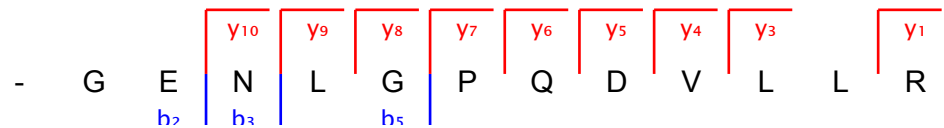
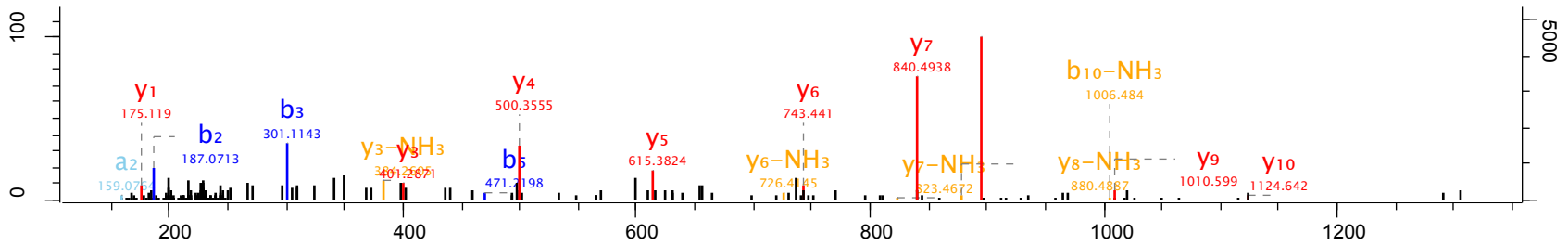
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	23595	TOF; CID	89.23	802.89	C12orf49



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	25261	TOF; CID	43.59	486.95	TIGD1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	25732	TOF; CID	59.23	655.86	USP40



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

28389

Method

TOF; CID

Score

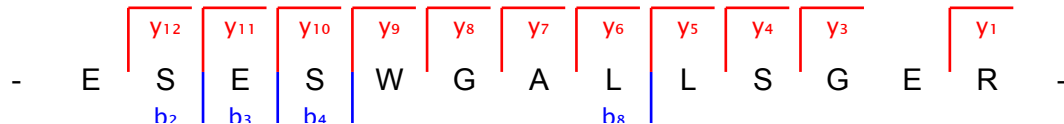
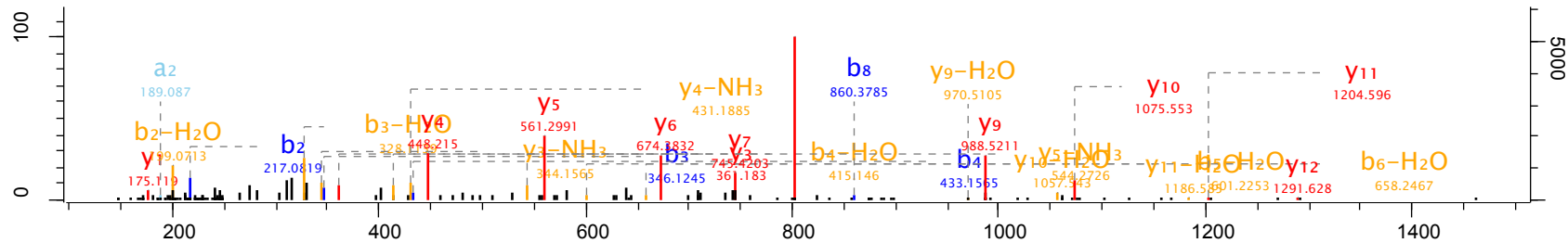
121.56

m/z

710.84

Gene names

SLC39A13



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

28819

Method

TOF; CID

Score

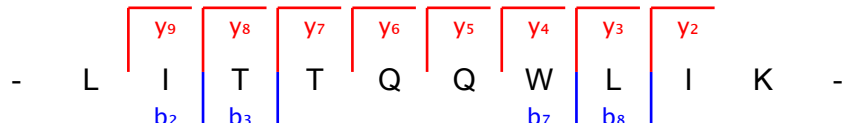
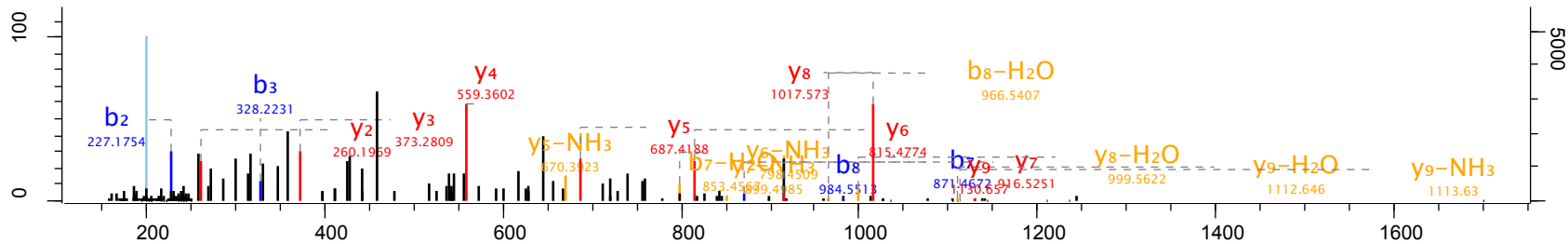
112.36

m/z

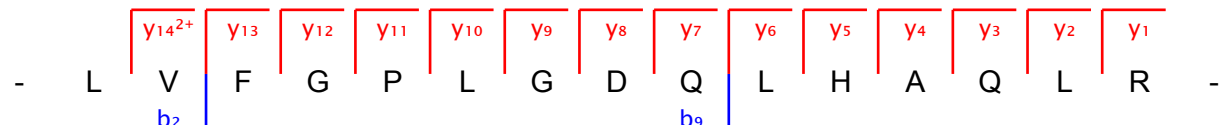
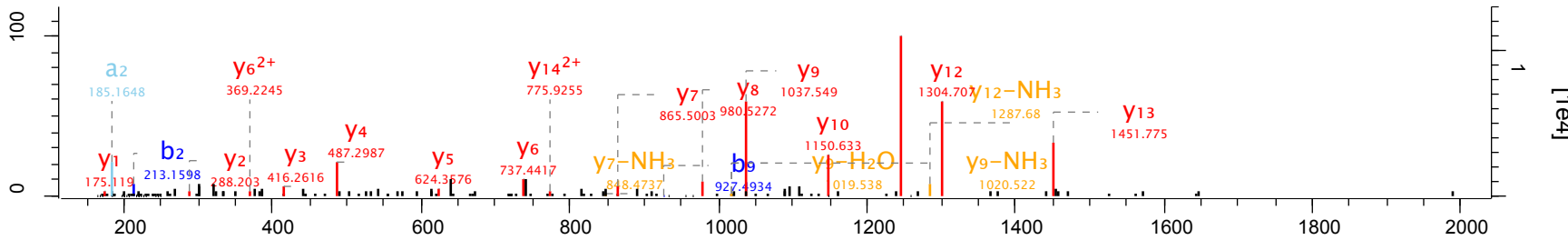
622.37

Gene names

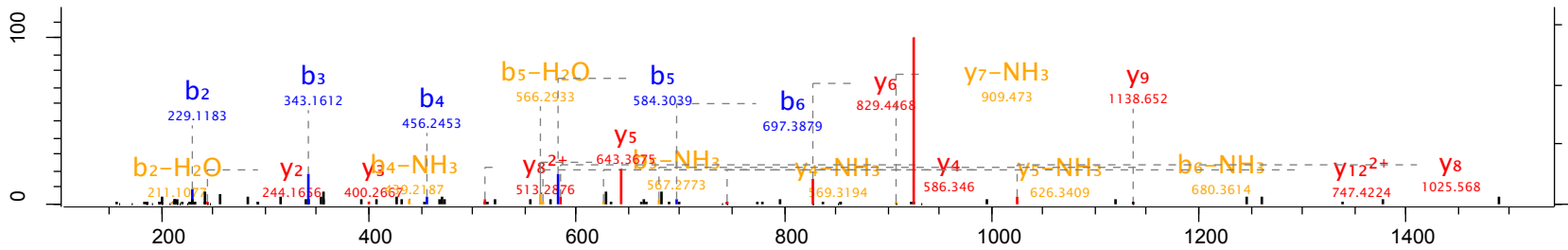
MT-ATP6



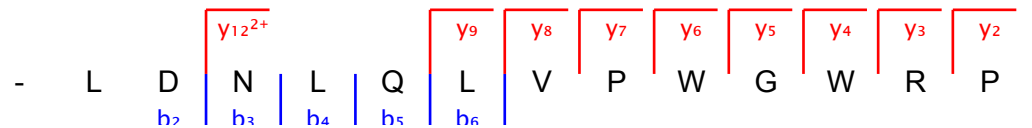
Raw file 20141014_fract12_dyn_5ul_F4_01_591 Scan 30571 Method TOF; CID Score 69.88 m/z 555.31 Gene names ELF3



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	34630	TOF; CID	77.22	574.99	ZMYND19



[1e4]



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

34915

Method

TOF; CID

Score

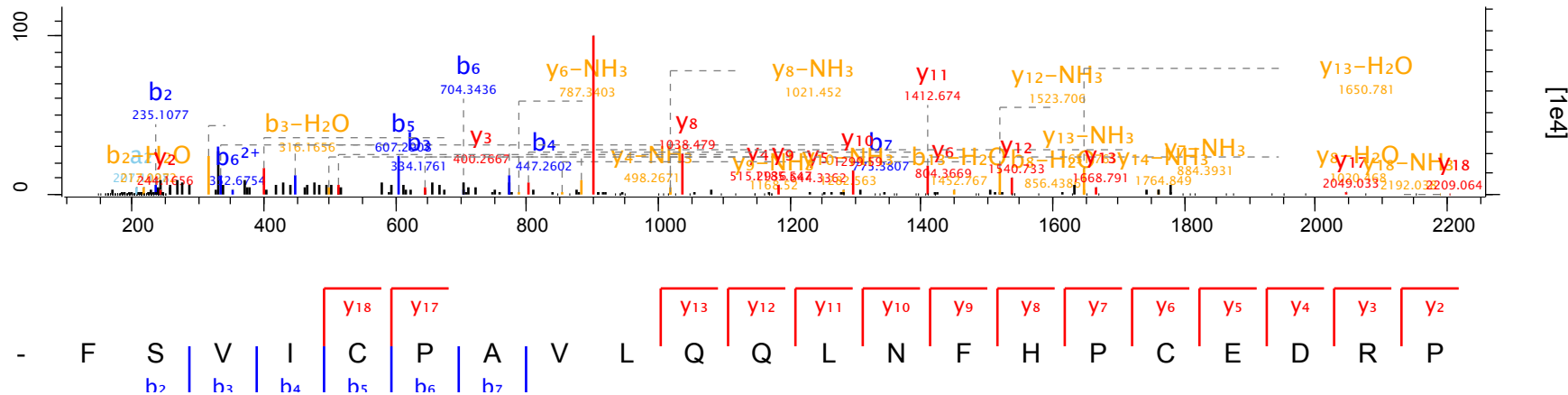
140.2

m/z

666.09

Gene names

SLC39A8



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

35587

Method

TOF; CID

Score

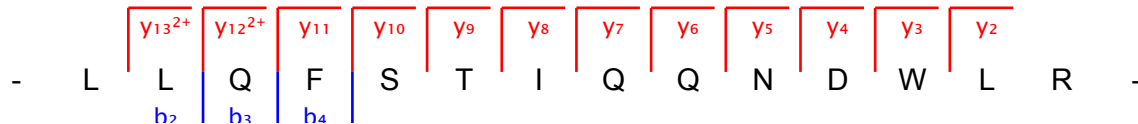
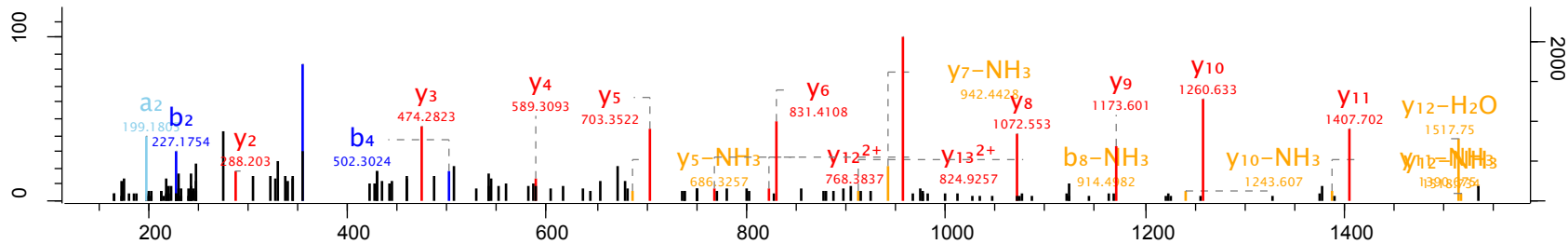
88.28

m/z

881.47

Gene names

C5orf28



Raw file

20141014_fract12_dyn_5ul_F4_01_591

Scan

37009

Method

TOF; CID

Score

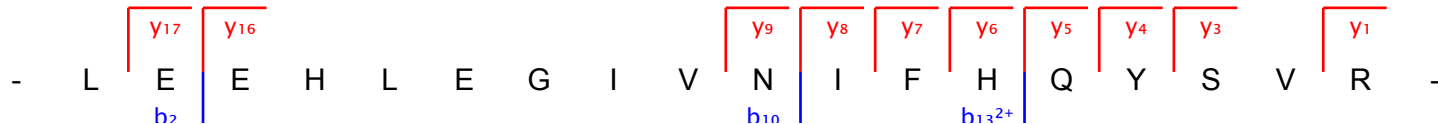
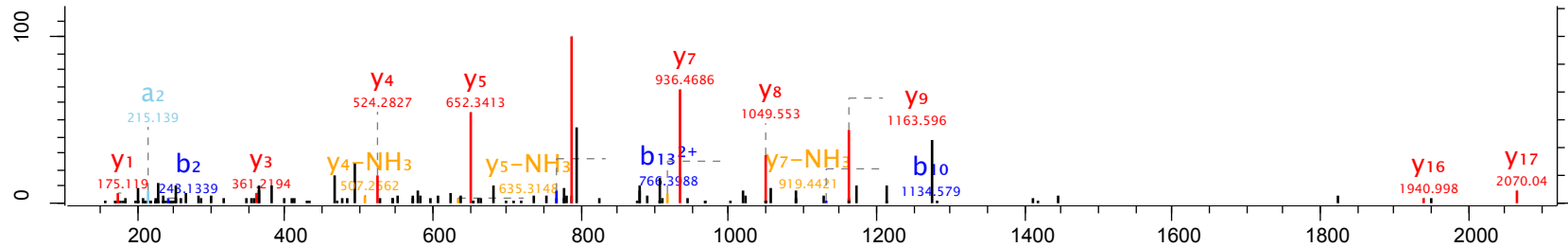
41.26

m/z

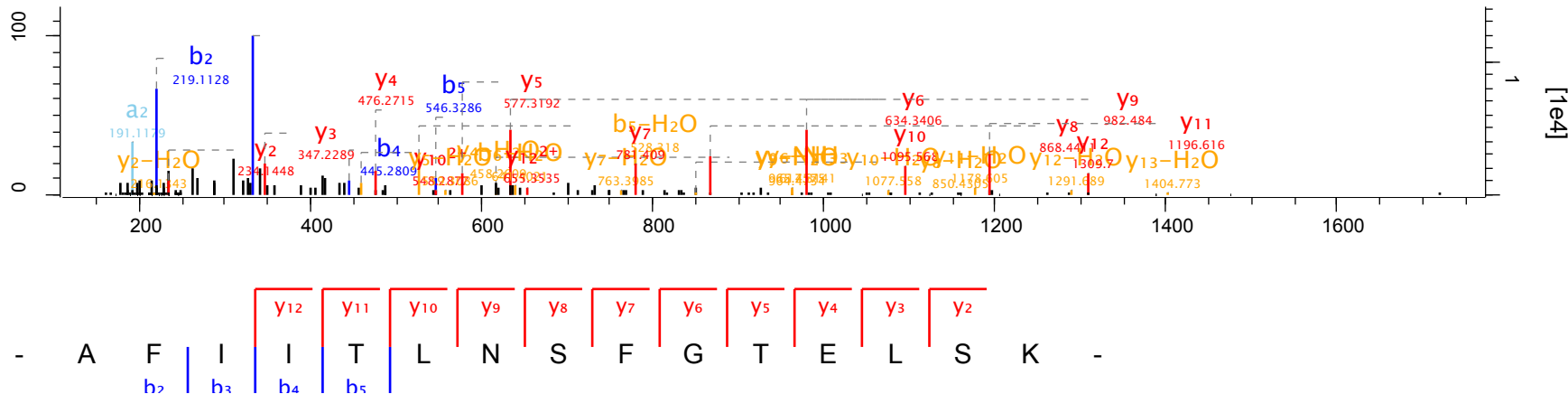
546.79

Gene names

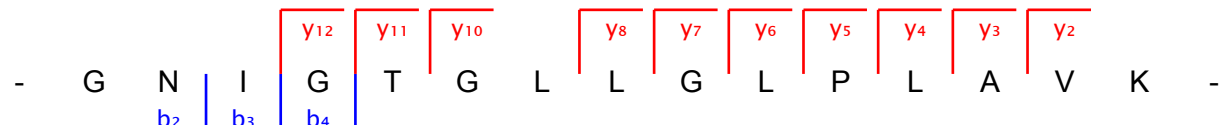
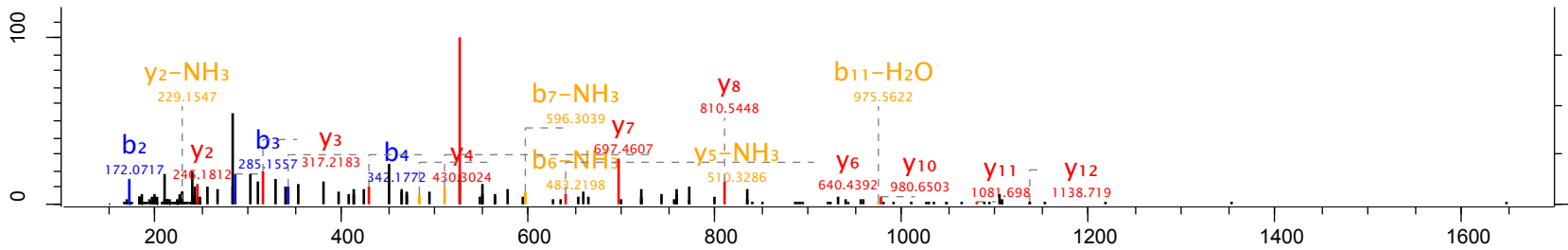
S100A12



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	38365	TOF; CID	128.22	821.44	ATP6V0D2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract12_dyn_5ul_F4_01_591	38881	TOF; CID	53.3	712.44	SLC36A1



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

13502

Method

TOF; CID

Score

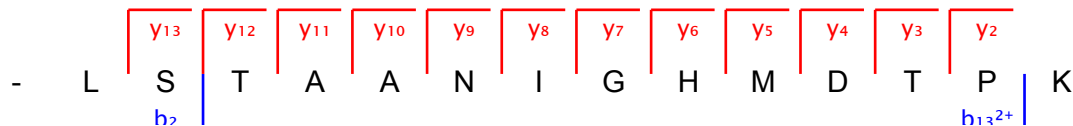
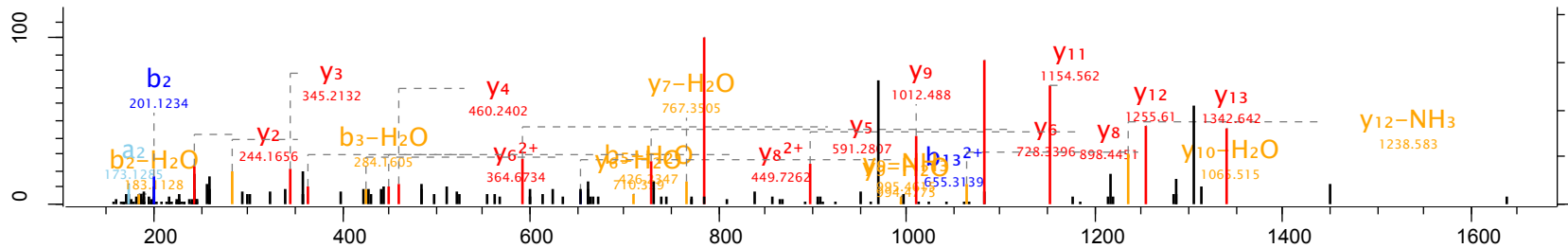
103.13

m/z

485.91

Gene names

SLC41A1



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

18052

Method

TOF; CID

Score

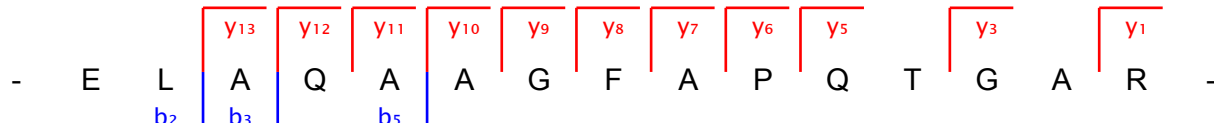
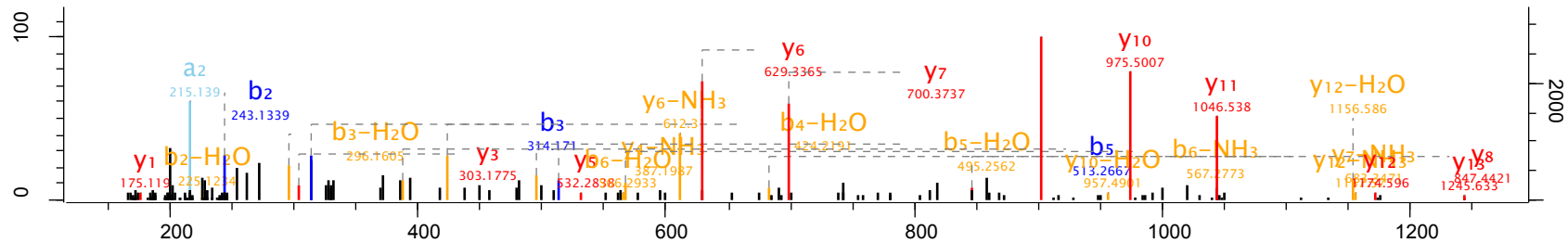
106.14

m/z

744.38

Gene names

DNAAF3



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

19254

Method

TOF; CID

Score

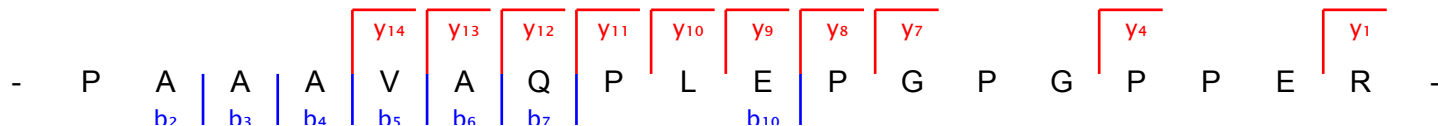
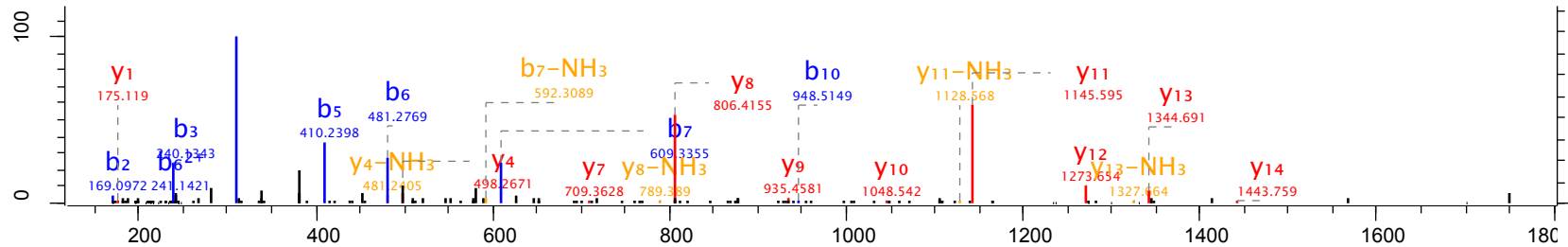
71.69

m/z

877.46

Gene names

KCTD2



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

19954

Method

TOF; CID

Score

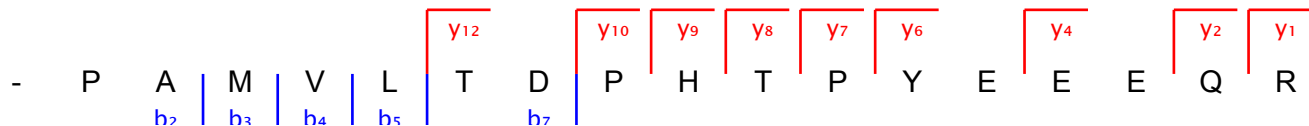
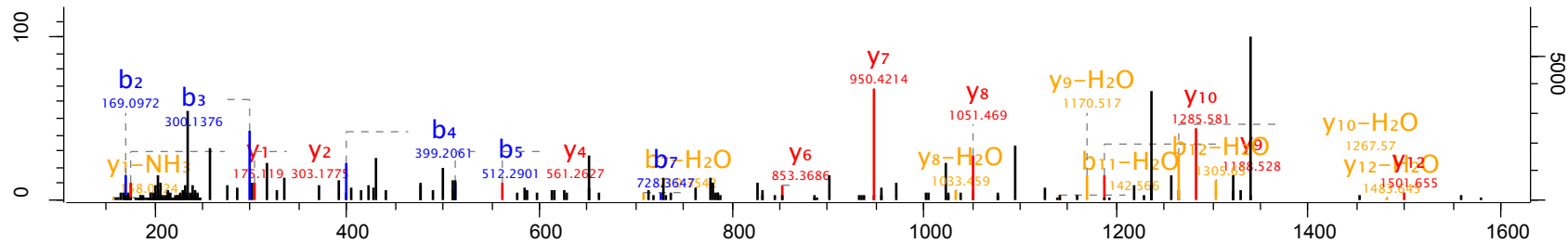
61.17

m/z

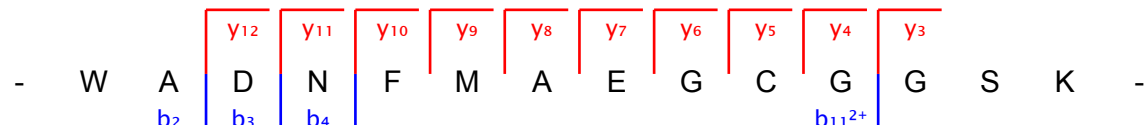
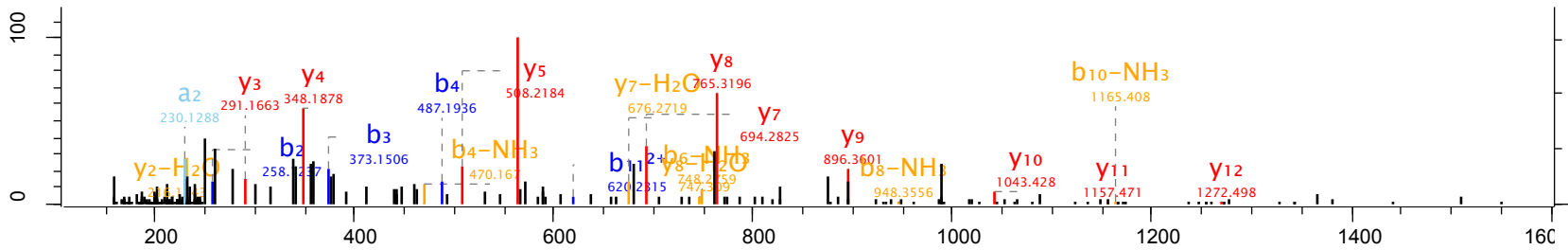
672.32

Gene names

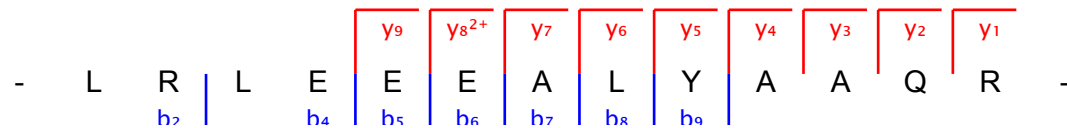
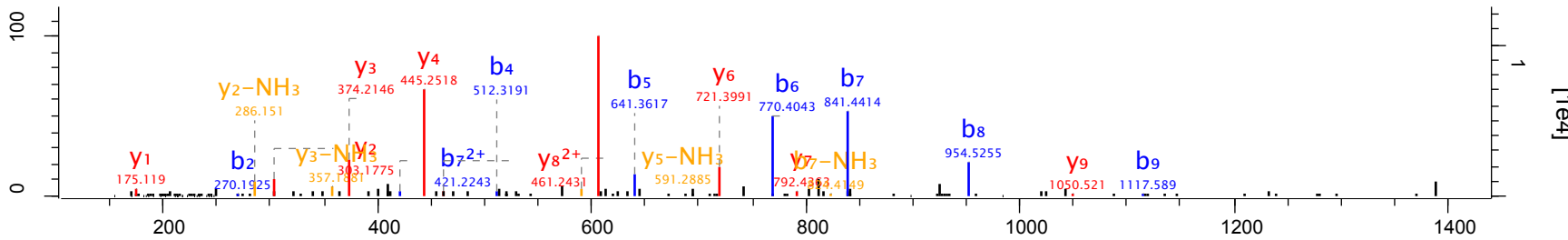
IL17RA



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	21431	TOF; CID	60.97	765.82	SLC35F6



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	22212	TOF; CID	62.29	521.28	APIAR



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

22541

Method

TOF; CID

Score

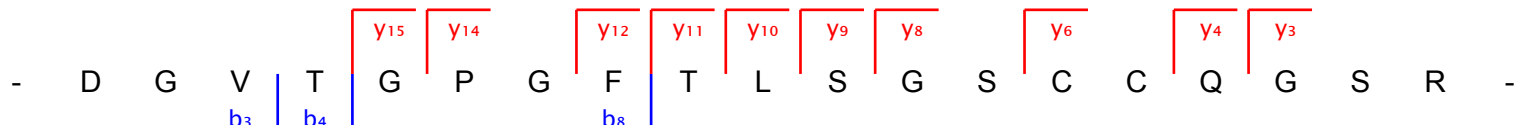
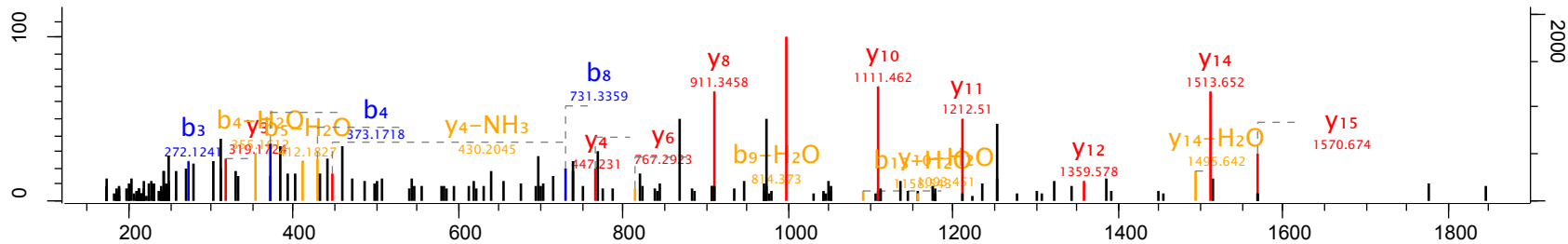
58.02

m/z

972.42

Gene names

LYPD3



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

23383

Method

TOF; CID

Score

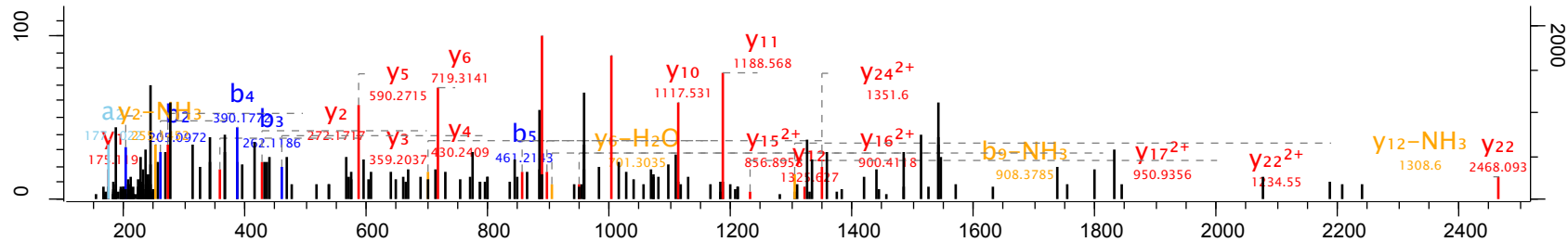
33.96

m/z

773.85

Gene names

TM2D2



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

24347

Method

TOF; CID

Score

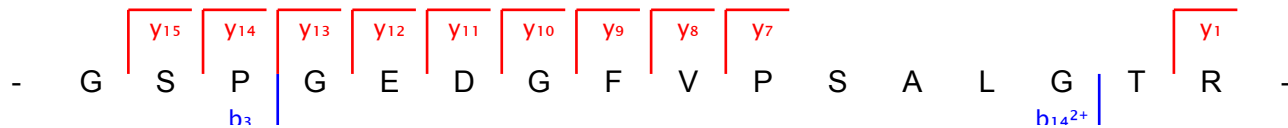
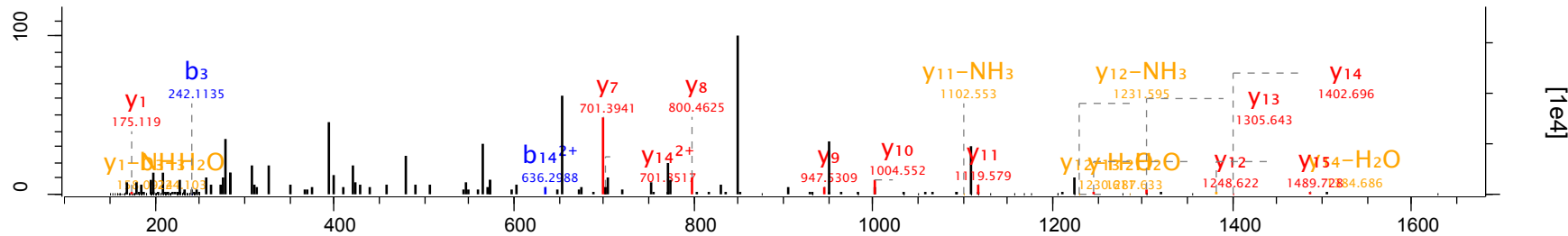
42.09

m/z

773.88

Gene names

METTL10



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

24524

Method

TOF; CID

Score

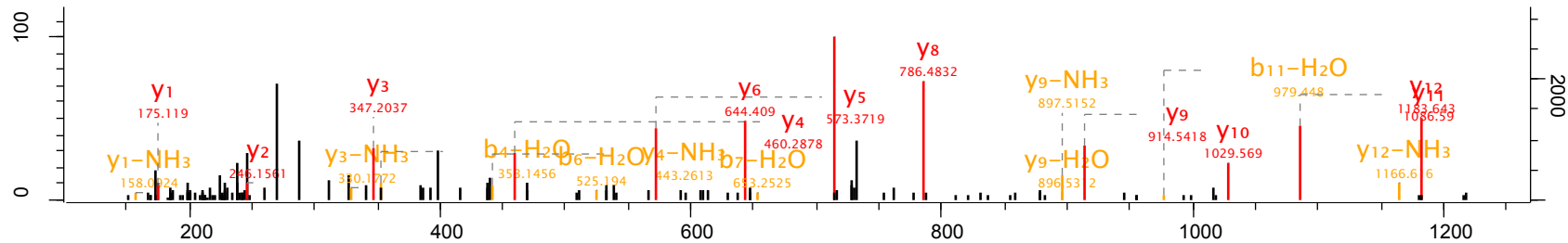
48.22

m/z

728.87

Gene names

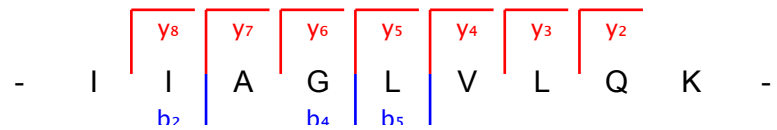
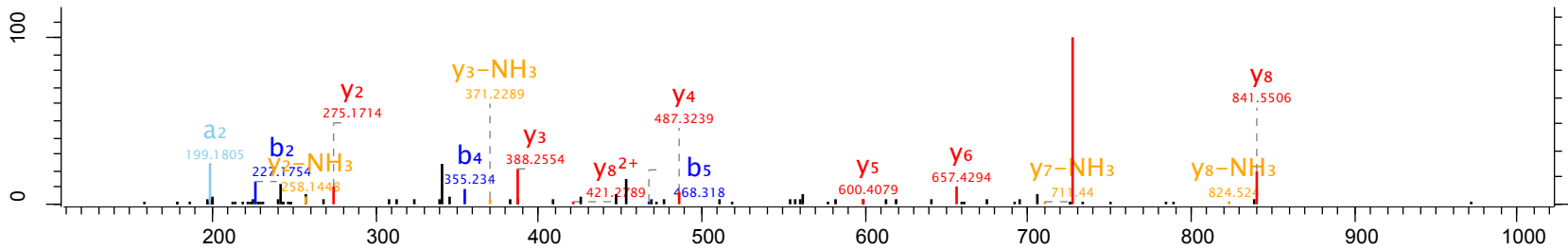
ZNF655



- E G S P G D Q A A A L L T A R -

Fragmentation diagram showing the sequence of amino acids: P, G, D, Q, A, A, A, L, L, T, A, R. Red brackets above the sequence indicate the fragmentation points for the y-ions (y1 to y12) shown in the mass spectrum above.

Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	24998	TOF; CID	81.3	477.82	BBS10



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

27038

Method

TOF; CID

Score

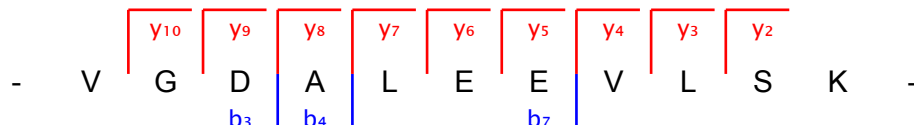
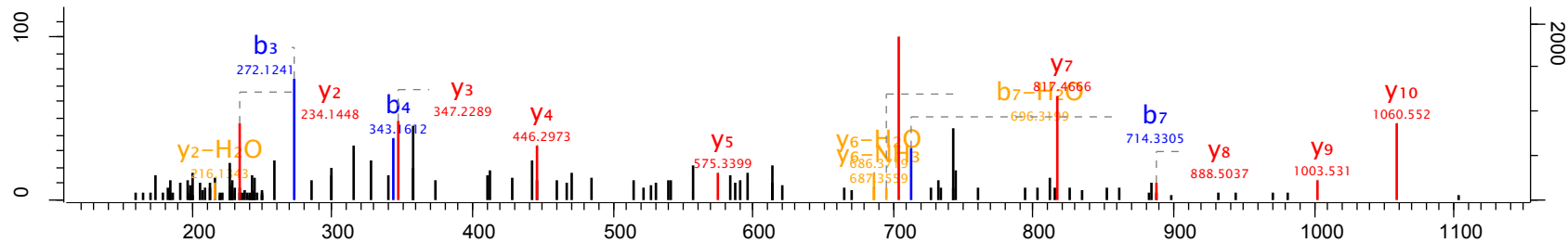
62.09

m/z

580.31

Gene names

GADD45A



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

27860

Method

TOF; CID

Score

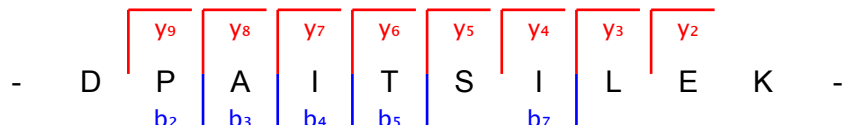
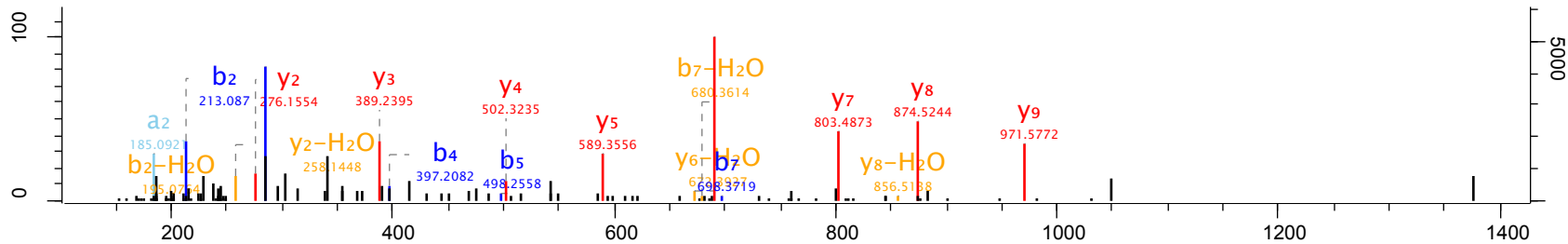
103.08

m/z

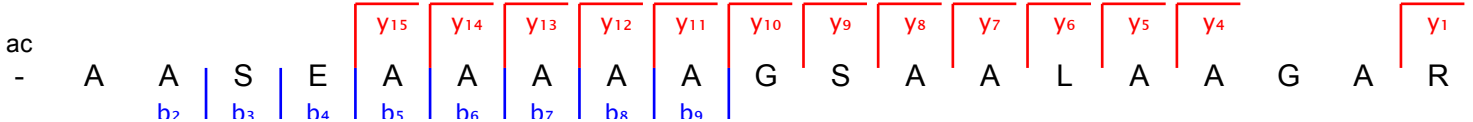
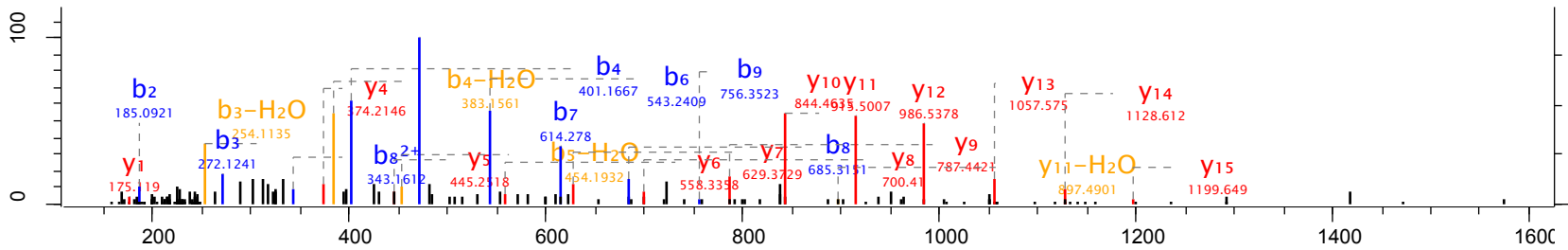
543.8

Gene names

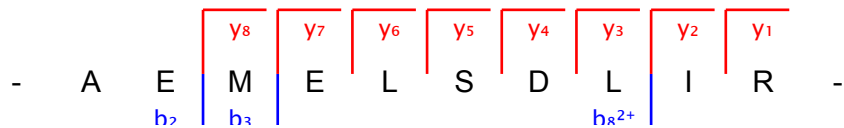
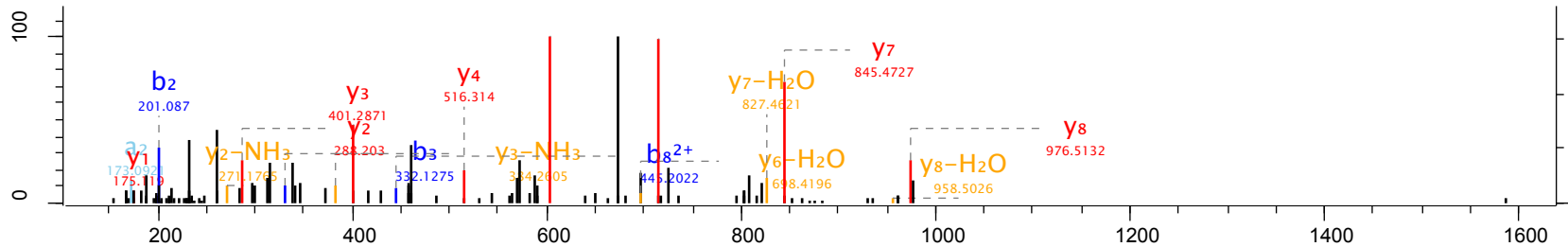
CPA4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	29425	TOF; CID	78.67	800.41	ATMIN



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	30641	TOF; CID	72.32	588.8	PKD2



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

31178

Method

TOF; CID

Score

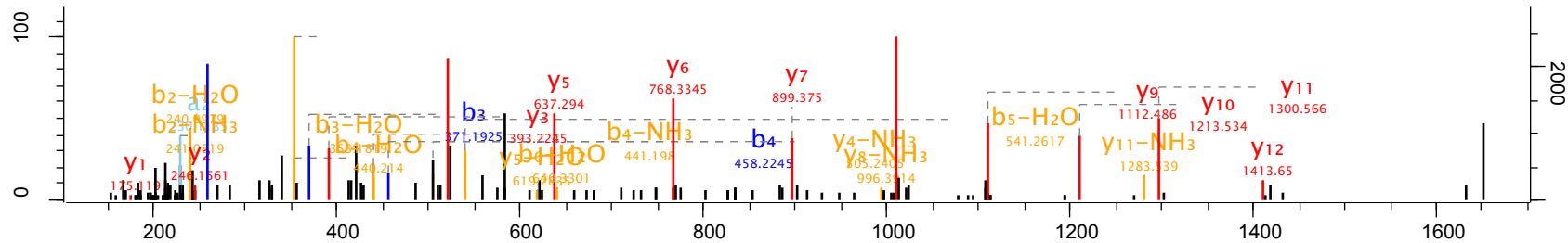
118.51

m/z

836.38

Gene names

WRB



- Q E L S T V N M M D E F A R -

b₂ b₃ b₄

y₁₂ y₁₁ y₁₀ y₉ y₈ y₇ y₆ y₅ y₄ y₃ y₂ y₁

Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

32116

Method

TOF; CID

Score

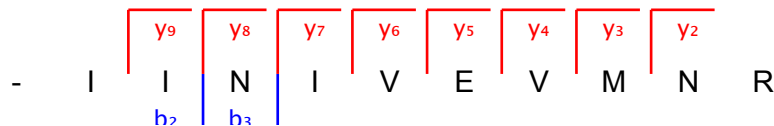
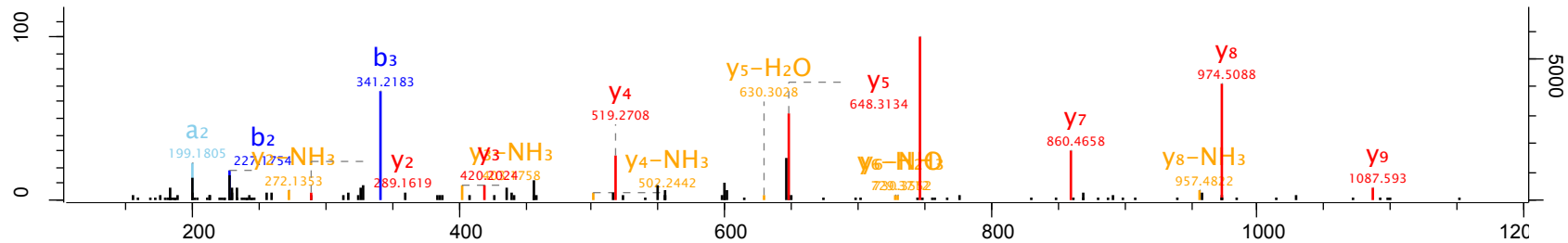
104.05

m/z

600.84

Gene names

ATG101



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

32472

Method

TOF; CID

Score

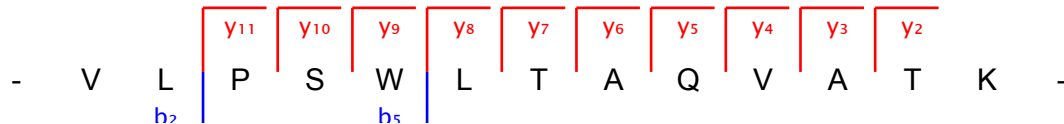
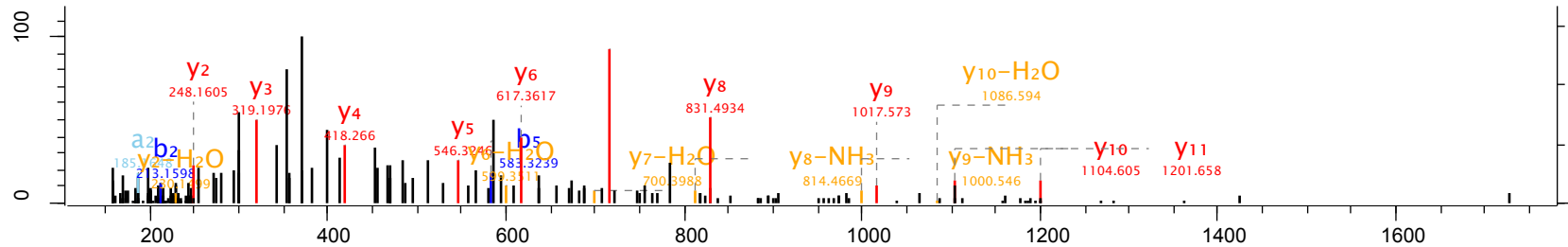
60.2

m/z

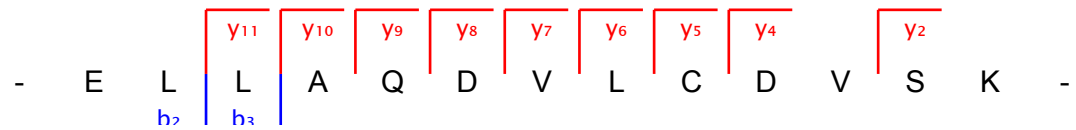
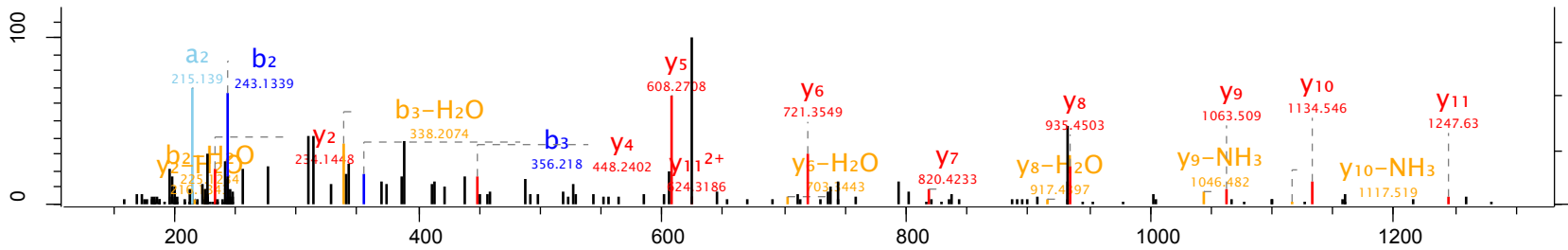
707.41

Gene names

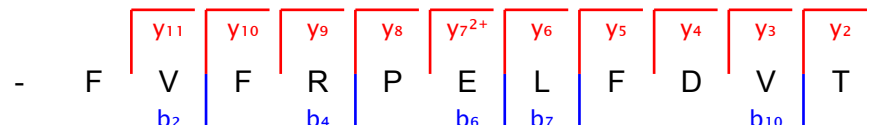
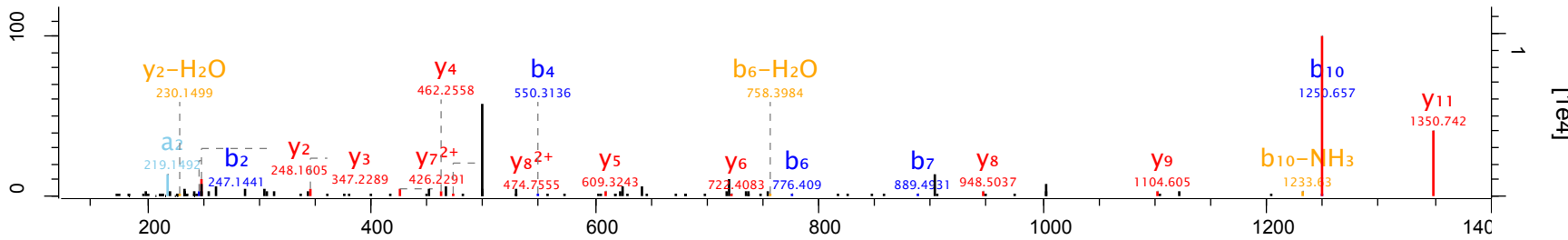
MRI



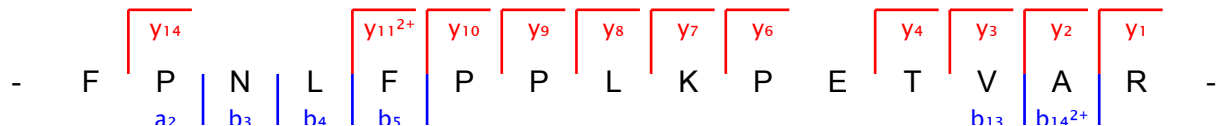
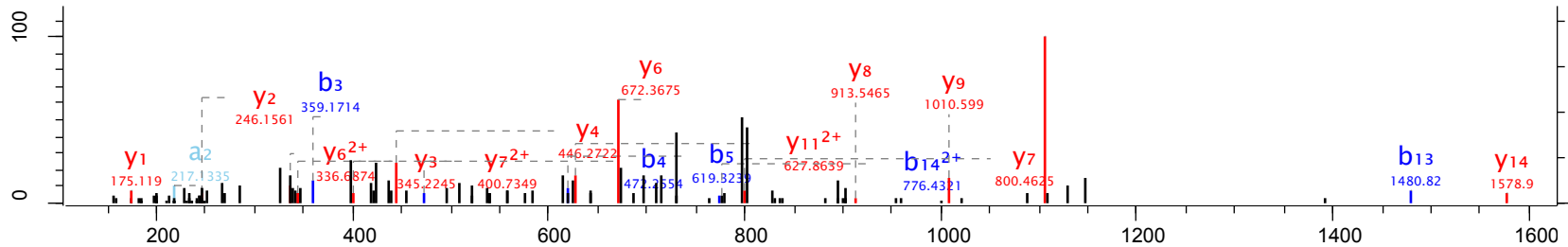
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	32774	TOF; CID	55.72	745.38	PPARGC1B



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	33810	TOF; CID	63.19	499.94	METTL4

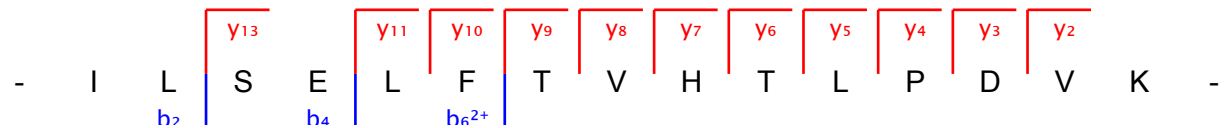
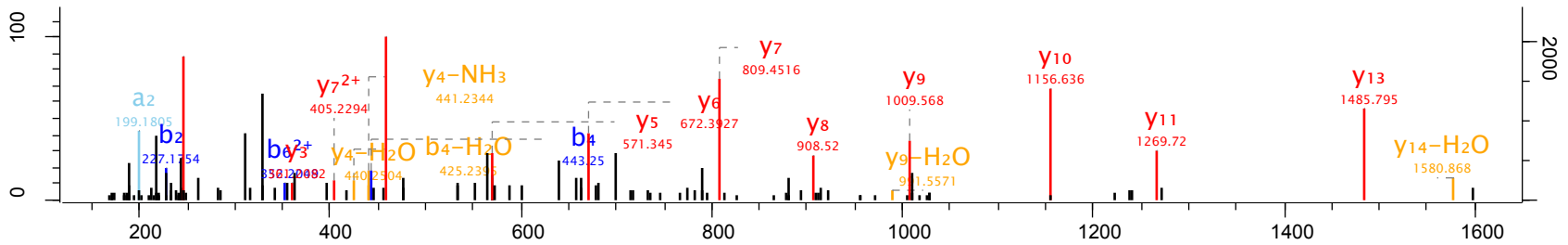


Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	34518	TOF; CID	45.44	576	DHRS3

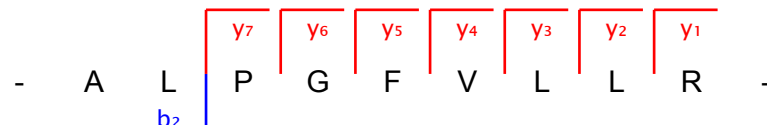
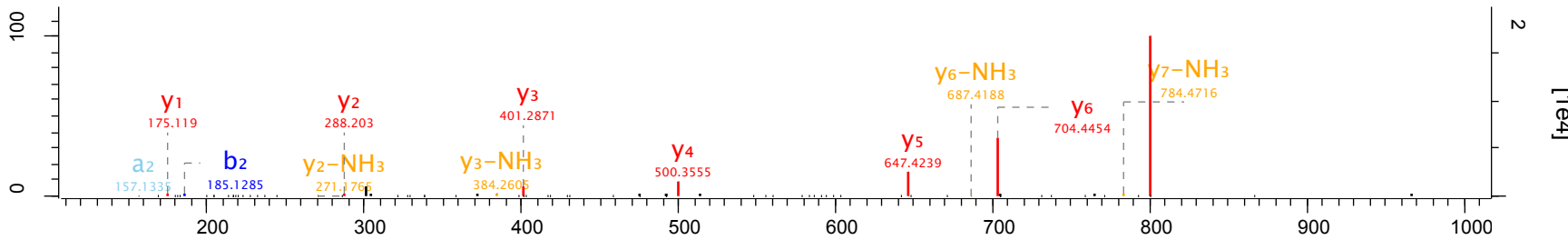


Raw file
 20141014_fract13_dyn_5ul_F5_01_592

Scan	Method	Score	m/z	Gene names
34790	TOF; CID	79.34	571.32	G2E3



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract13_dyn_5ul_F5_01_592	35837	TOF; CID	81.3	493.31	SLC29A3



Raw file

20141014_fract13_dyn_5ul_F5_01_592

Scan

37719

Method

TOF; CID

Score

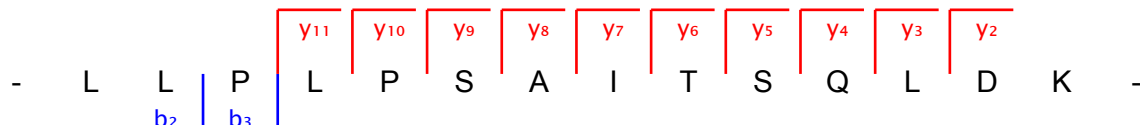
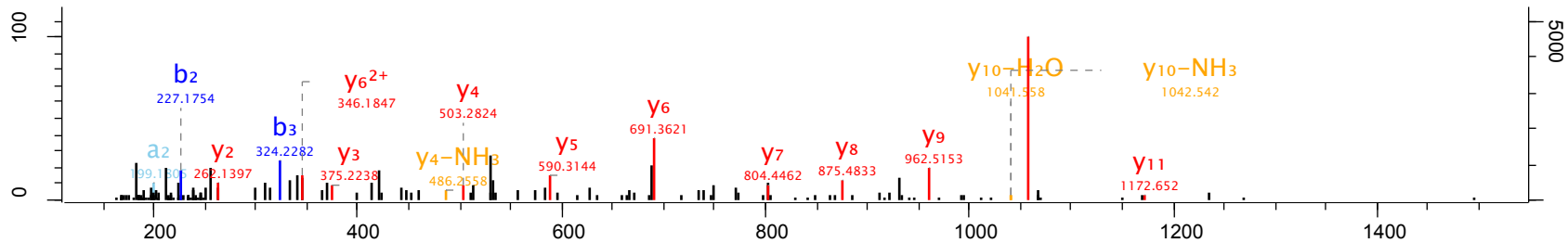
60.55

m/z

748.44

Gene names

SIM2;SIM1



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

10326

Method

TOF; CID

Score

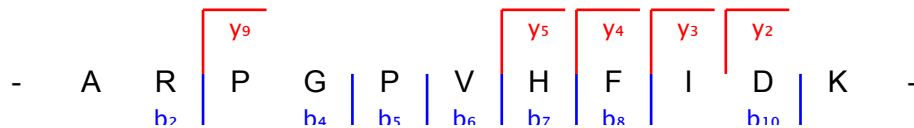
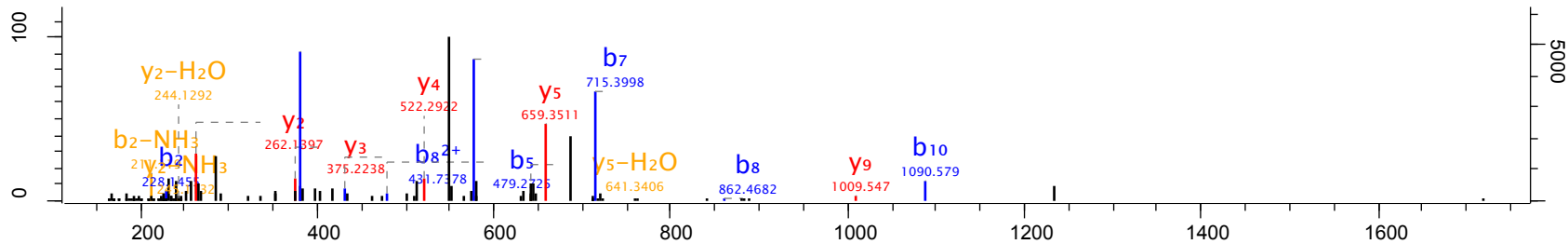
65.18

m/z

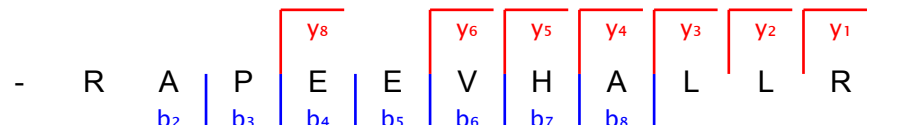
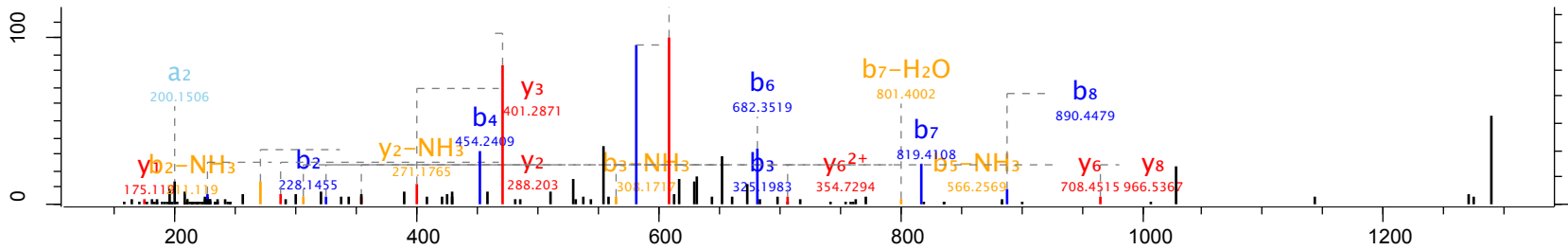
412.9

Gene names

ARRDC1-AS1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract14_dyn_5ul_F6_01_593	11009	TOF; CID	68.52	430.91	DPY19L3



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

11297

Method

TOF; CID

Score

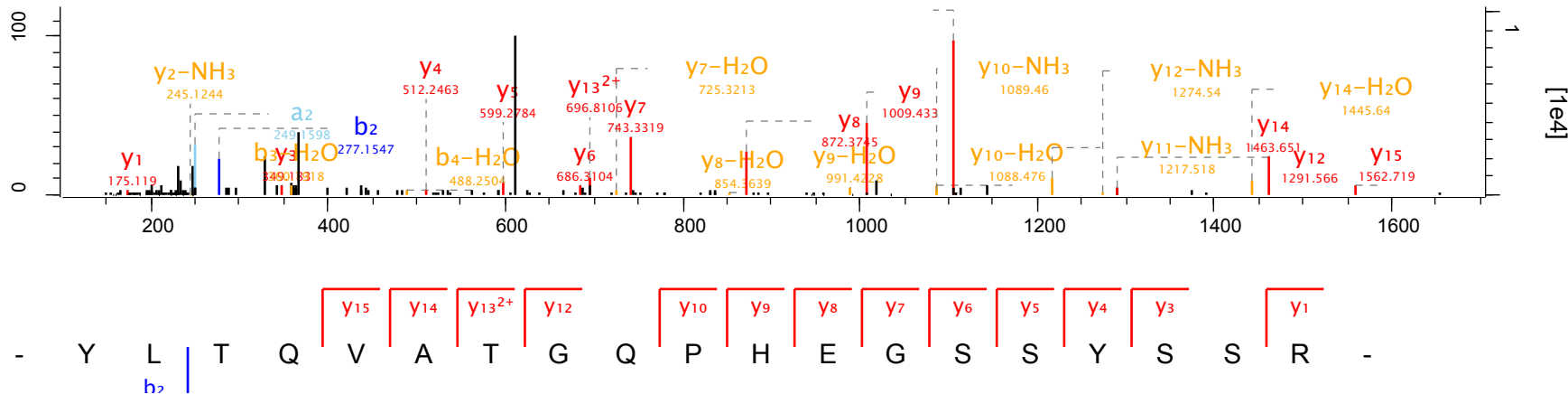
90.38

m/z

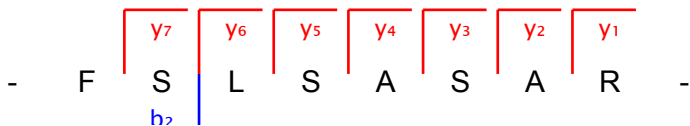
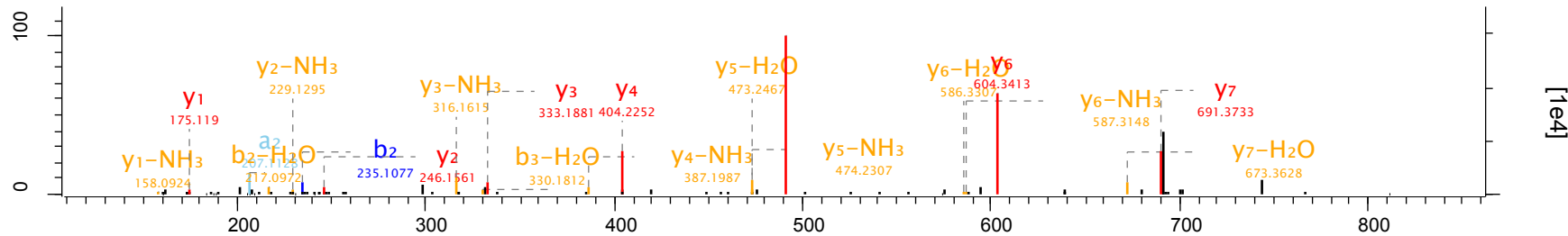
690.33

Gene names

MED11



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract14_dyn_5ul_F6_01_593	12997	TOF; CID	118.74	420.73	ARMC7



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

13885

Method

TOF; CID

Score

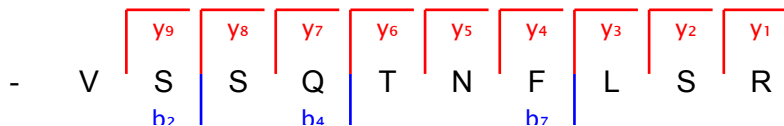
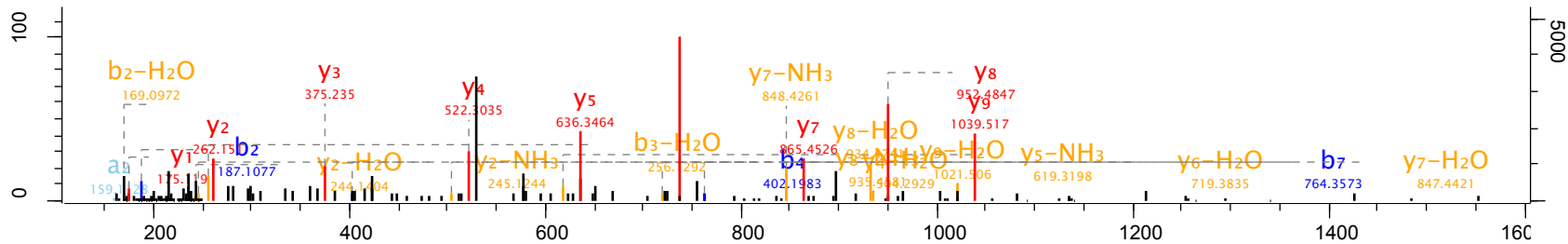
88.16

m/z

569.79

Gene names

SGMS2



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

18083

Method

TOF; CID

Score

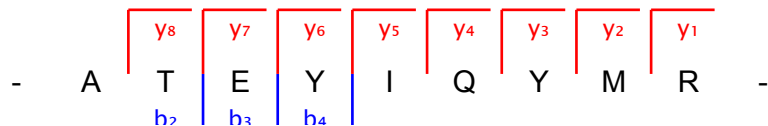
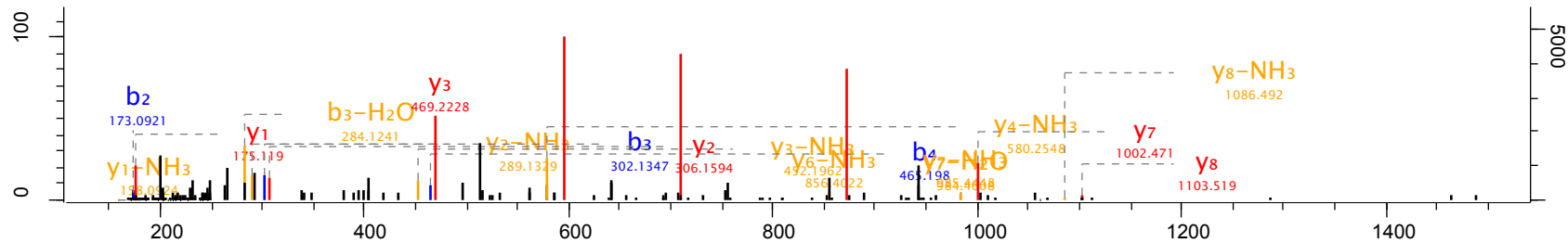
116.37

m/z

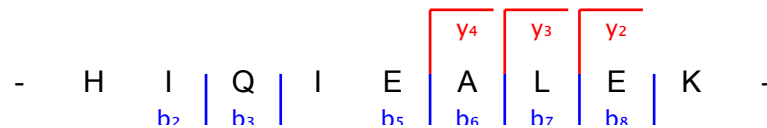
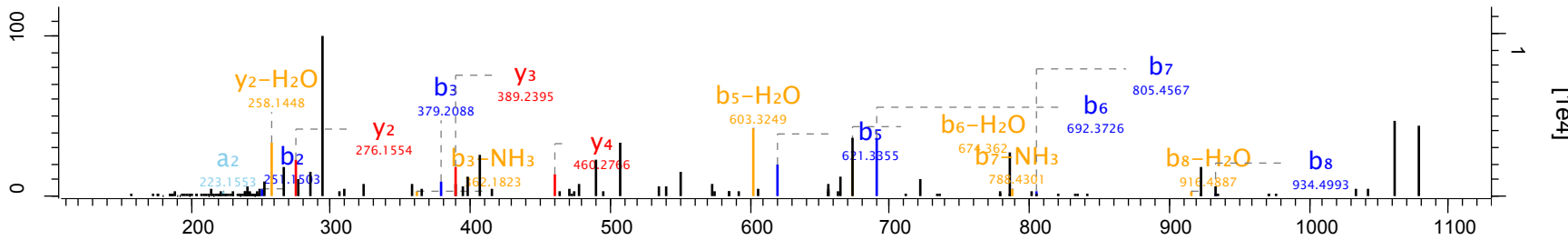
587.78

Gene names

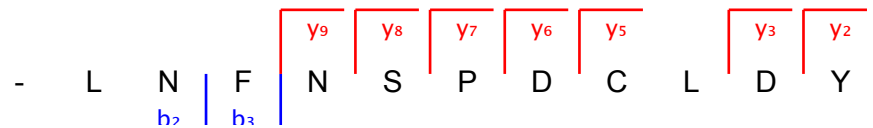
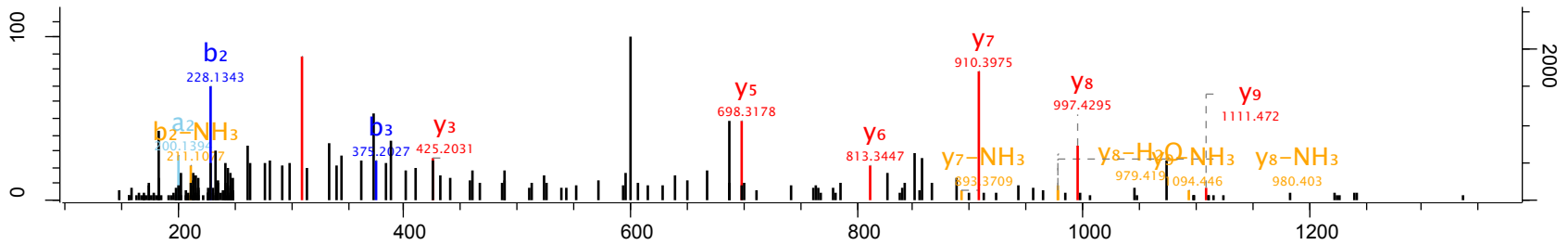
MAX



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract14_dyn_5ul_F6_01_593	20499	TOF; CID	79.15	540.81	ADAMTS14



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract14_dyn_5ul_F6_01_593	22717	TOF; CID	64.12	743.34	OTX1



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

24847

Method

TOF; CID

Score

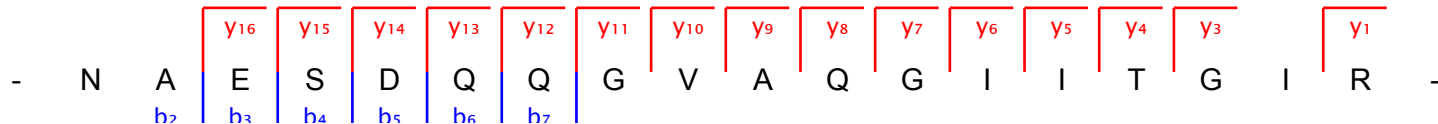
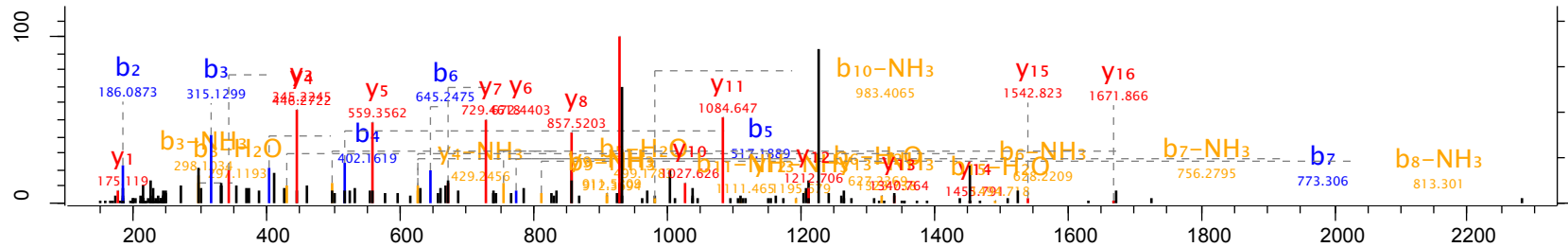
118.96

m/z

928.97

Gene names

HIATL1



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

30294

Method

TOF; CID

Score

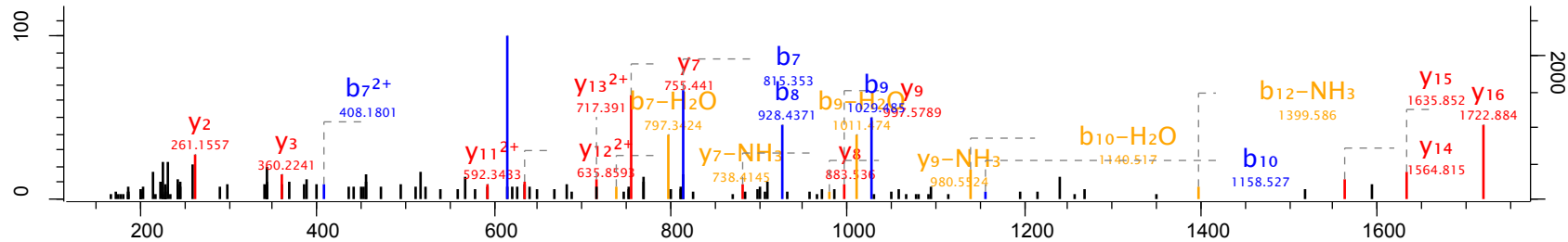
69.85

m/z

813.9

Gene names

PAG1

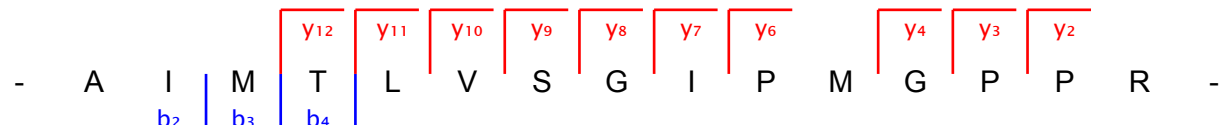
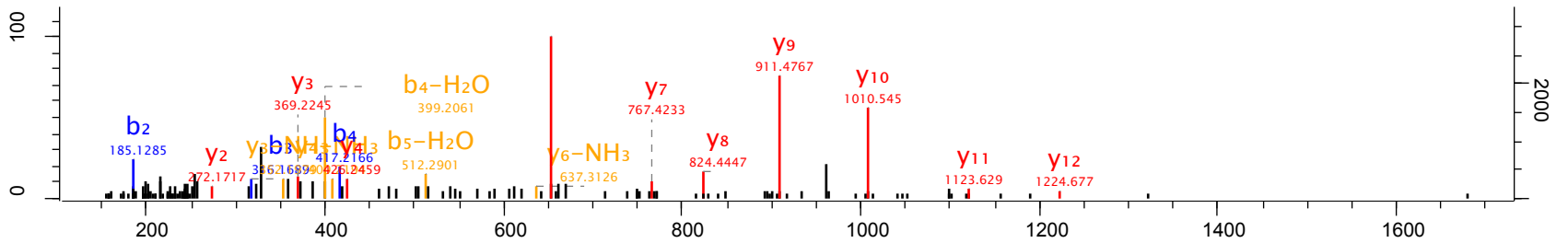


- S R E E D P T L T E E E I S A M Y S S V N

h₅ | h₇ | h₈ | h₉ | h₁₀

y₁₆ | y₁₅ | y₁₄ | y₁₃²⁺ | y₁₂²⁺ | y₁₁²⁺ | y₉

Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract14_dyn_5ul_F6_01_593	35876	TOF; CID	78.81	770.42	NPL



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

35994

Method

TOF; CID

Score

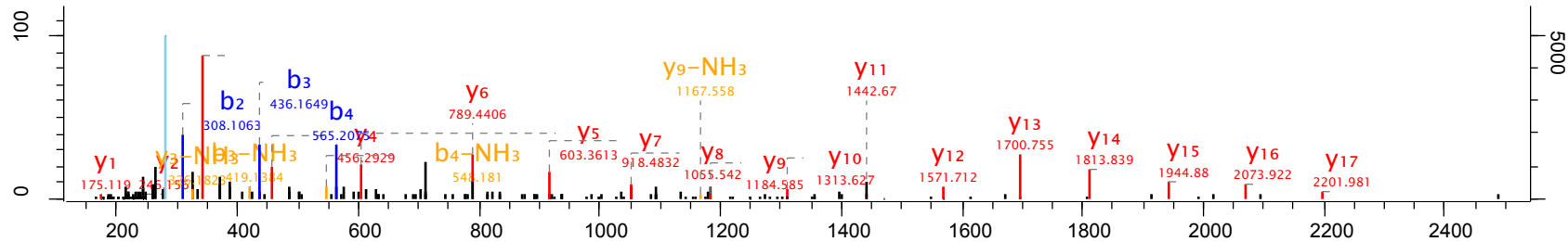
100.02

m/z

837.36

Gene names

PAIP2



- C F Q E M L E E E E H E W F I P A R -

Fragmentation sites are indicated by red brackets above the sequence, corresponding to the y ions labeled in the spectrum above. Blue brackets below the sequence correspond to the b ions labeled in the spectrum above.

Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

36542

Method

TOF; CID

Score

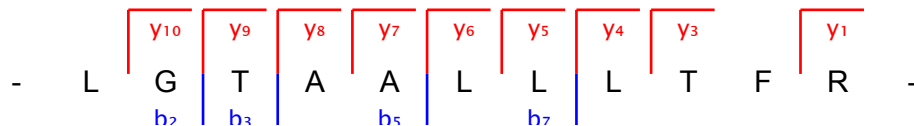
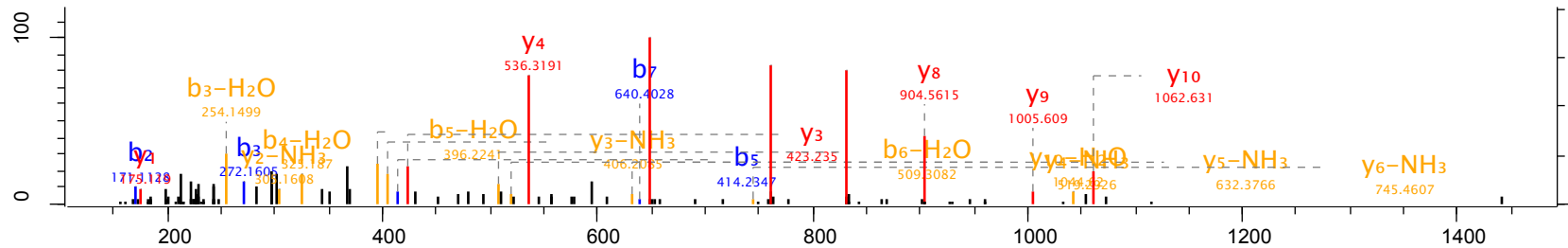
98.16

m/z

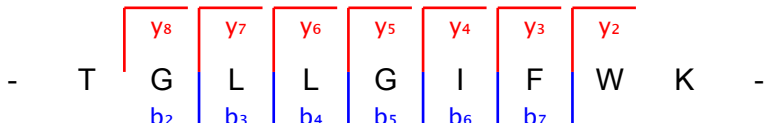
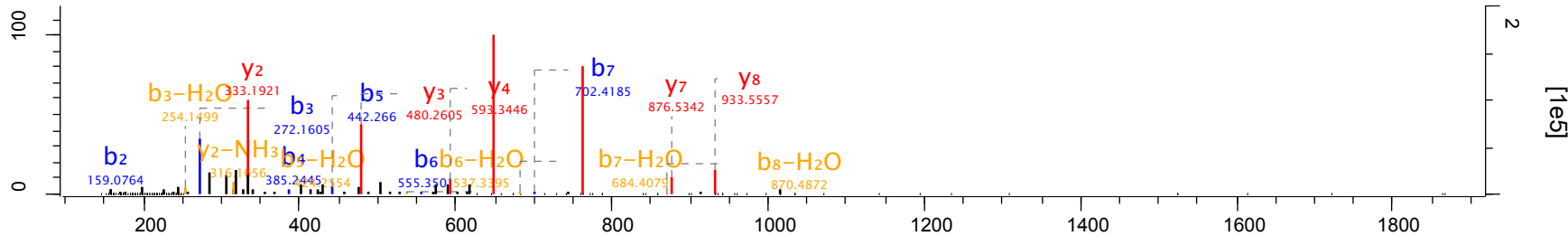
588.36

Gene names

KREMEN2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract14_dyn_5ul_F6_01_593	39070	TOF; CID	96.03	517.8	TMEM167A



Raw file

20141014_fract14_dyn_5ul_F6_01_593

Scan

39365

Method

TOF; CID

Score

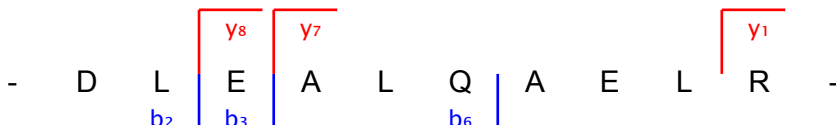
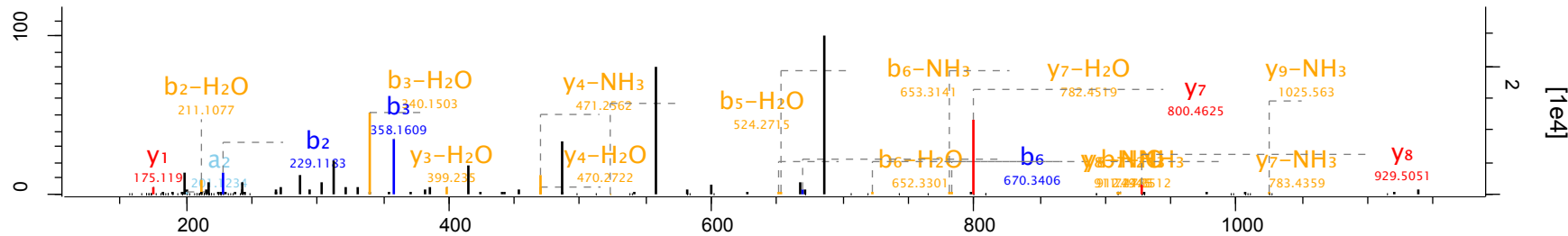
74.17

m/z

579.31

Gene names

FAM184B



Raw file

20141014_fract15_dyn_5ul_F7_01_594

Scan

13920

Method

TOF; CID

Score

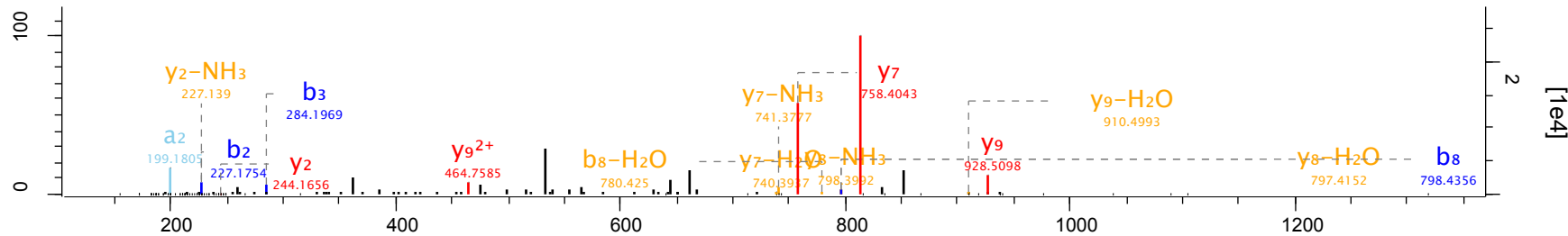
69.37

m/z

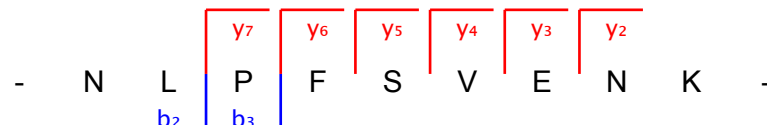
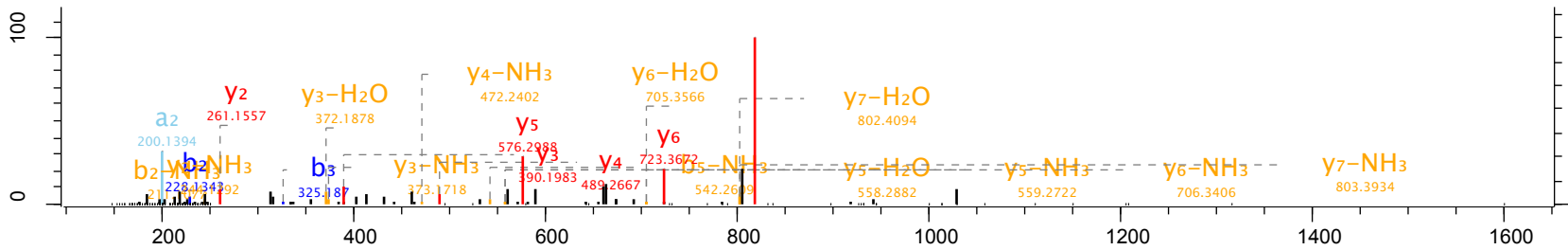
521.3

Gene names

CCDC17



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract15_dyn_5ul_F7_01_594	20708	TOF; CID	99.01	524.28	COX7C



[1e4]

Raw file

20141014_fract15_dyn_5ul_F7_01_594

Scan

24123

Method

TOF; CID

Score

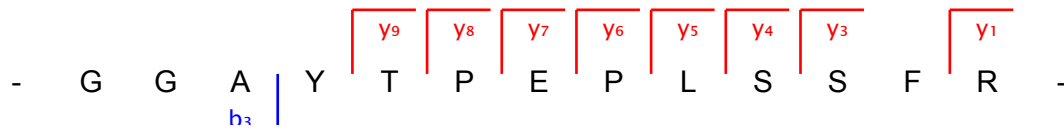
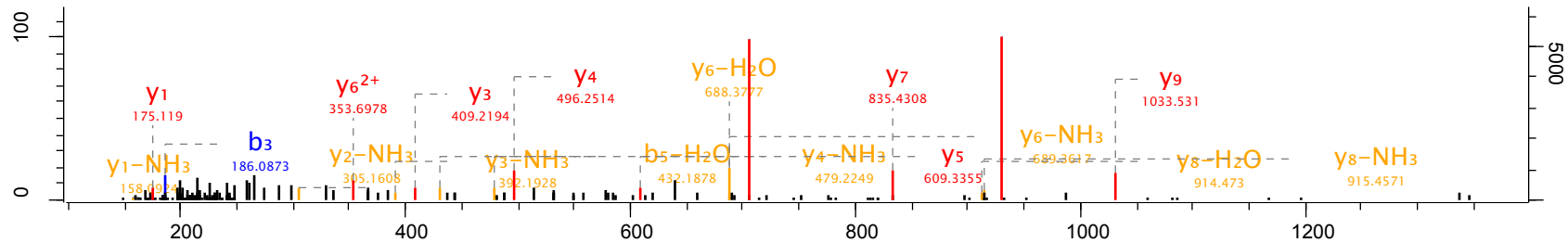
69.03

m/z

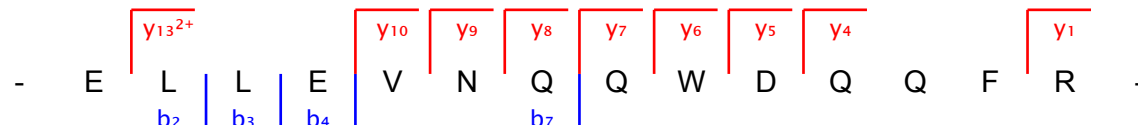
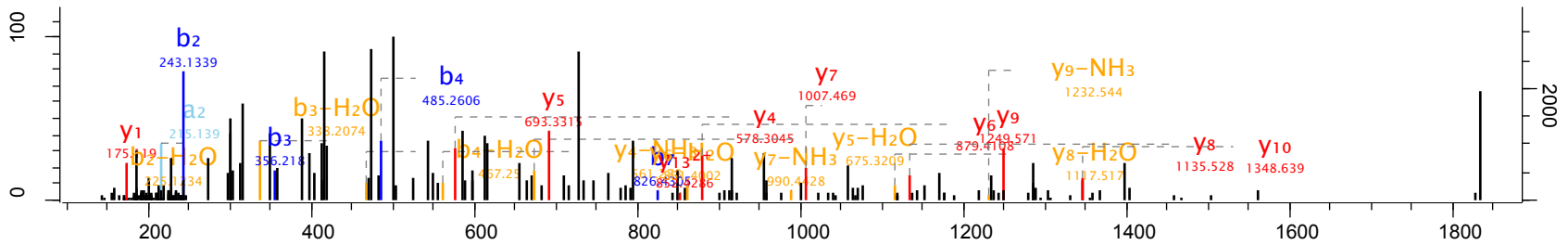
691.34

Gene names

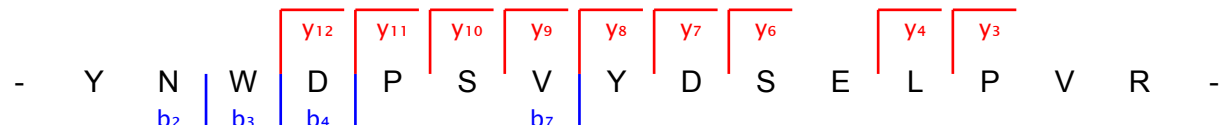
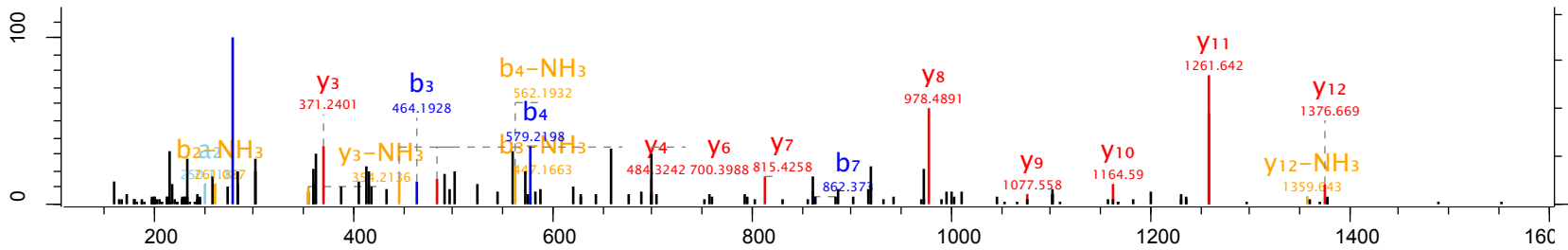
TMEM164;RP13-360B22.2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract15_dyn_5ul_F7_01_594	30946	TOF; CID	63.73	916.95	TNIP3



Raw file
 20141014_fract15_dyn_5ul_F7_01_594
 Scan 32768 Method TOF; CID Score 54.86 m/z 920.93 Gene names DEAF1



Raw file

20141014_fract15_dyn_5ul_F7_01_594

Scan

34288

Method

TOF; CID

Score

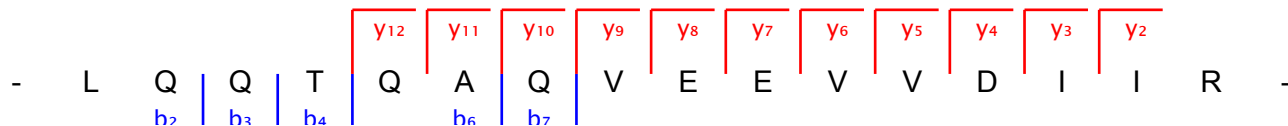
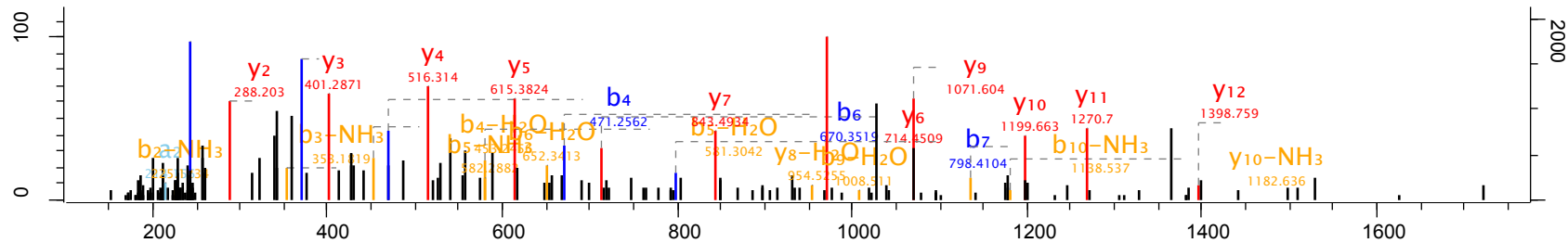
88.39

m/z

935.51

Gene names

VAMP1



Raw file

20141014_fract15_dyn_5ul_F7_01_594

Scan

34743

Method

TOF; CID

Score

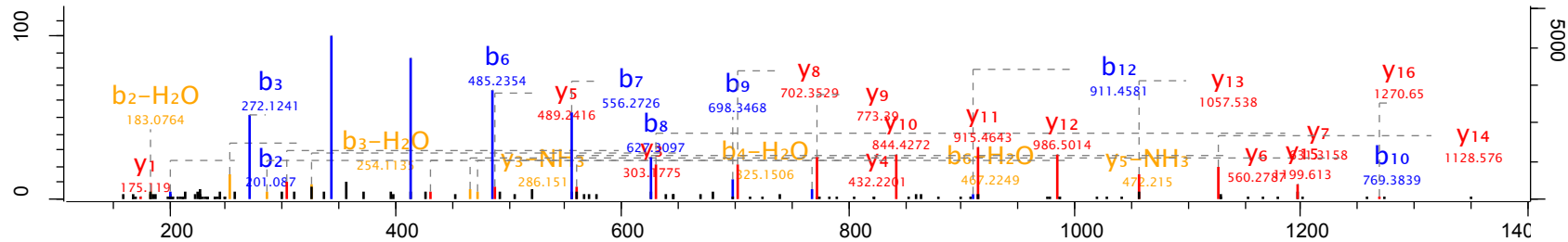
119.62

m/z

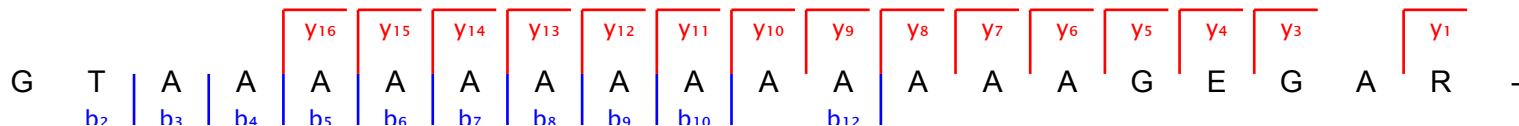
806.9

Gene names

NIPA1



ac



Raw file

20141014_fract15_dyn_5ul_F7_01_594

Scan

36642

Method

TOF; CID

Score

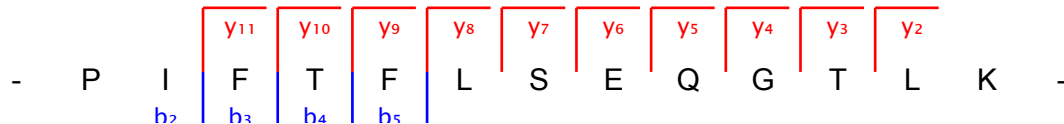
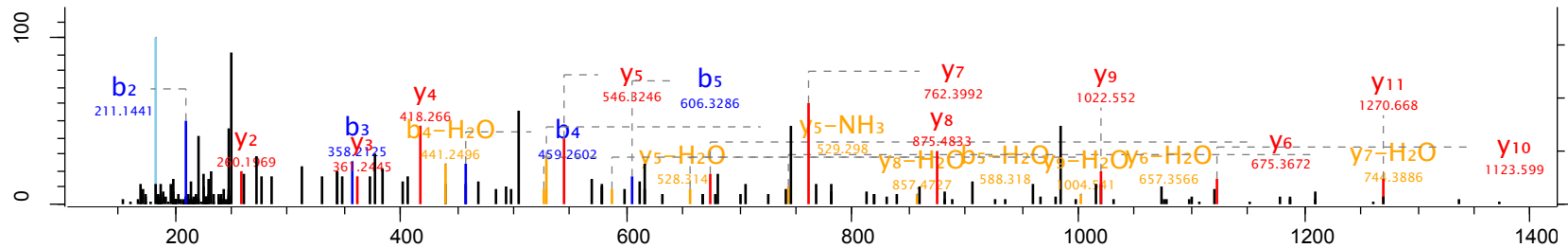
90.86

m/z

741.41

Gene names

VAV3



Raw file

20141014_fract15_dyn_5ul_F7_01_594

Scan

39052

Method

TOF; CID

Score

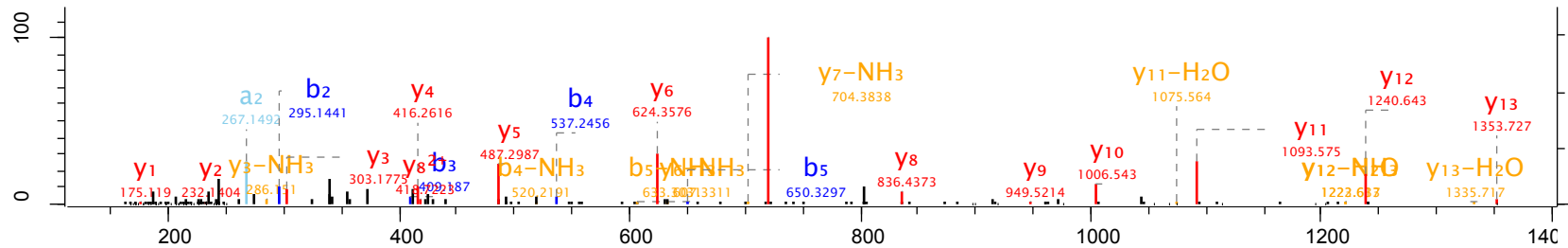
91.66

m/z

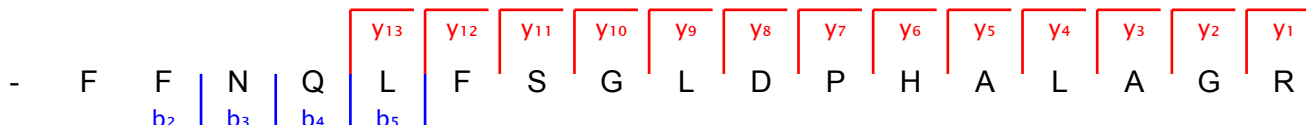
630.99

Gene names

CSAD



[1e4]



Raw file

20141014_fract15_dyn_5ul_F7_01_594

Scan

41392

Method

TOF; CID

Score

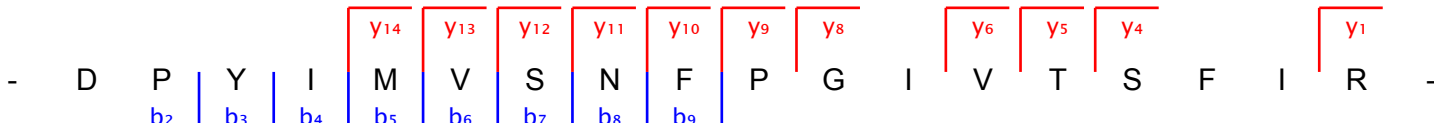
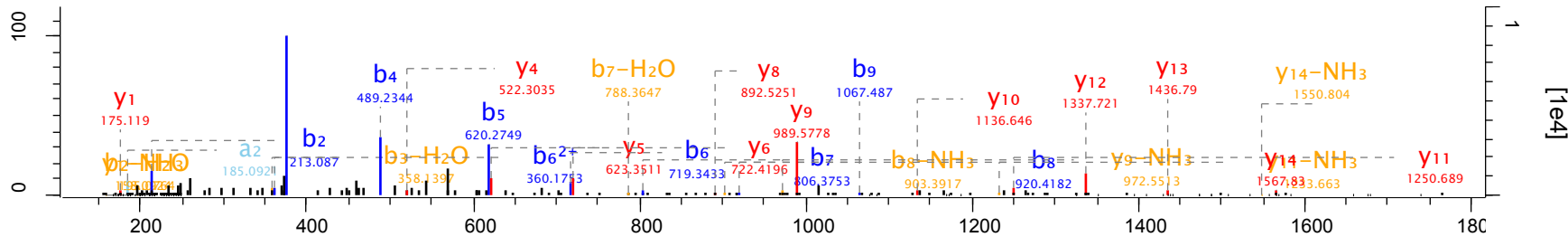
86.8

m/z

1029.03

Gene names

SLC50A1



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

8984

Method

TOF; CID

Score

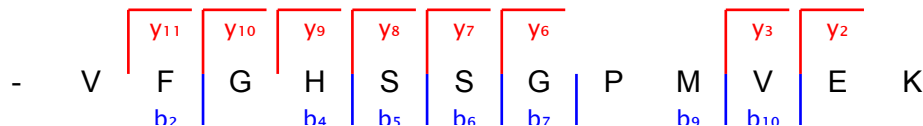
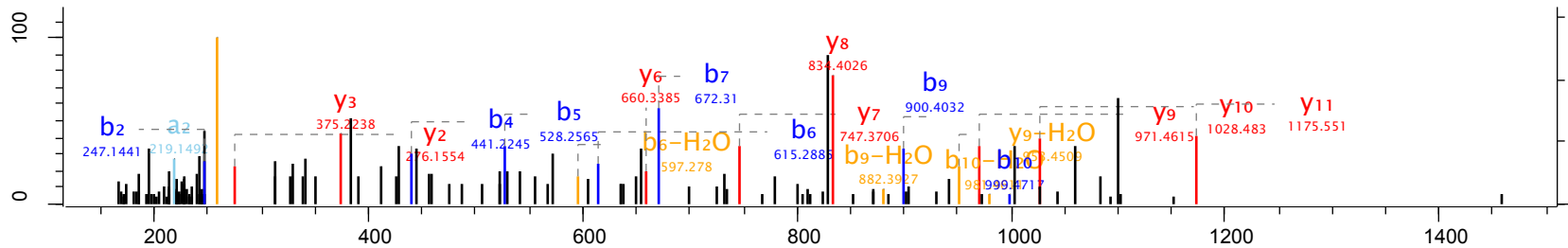
94.4

m/z

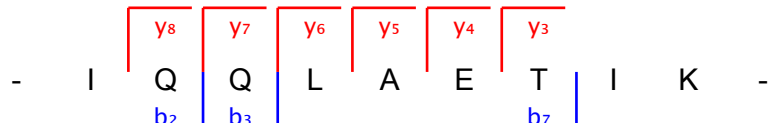
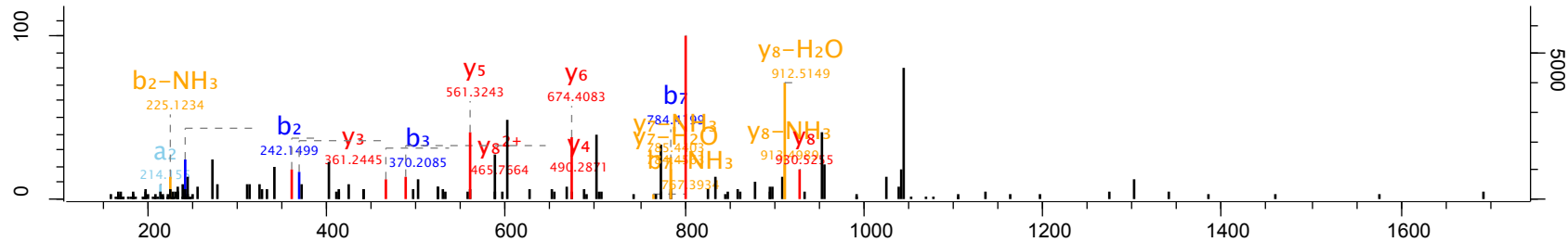
637.81

Gene names

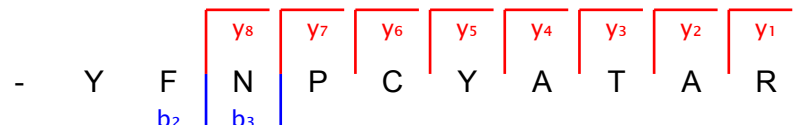
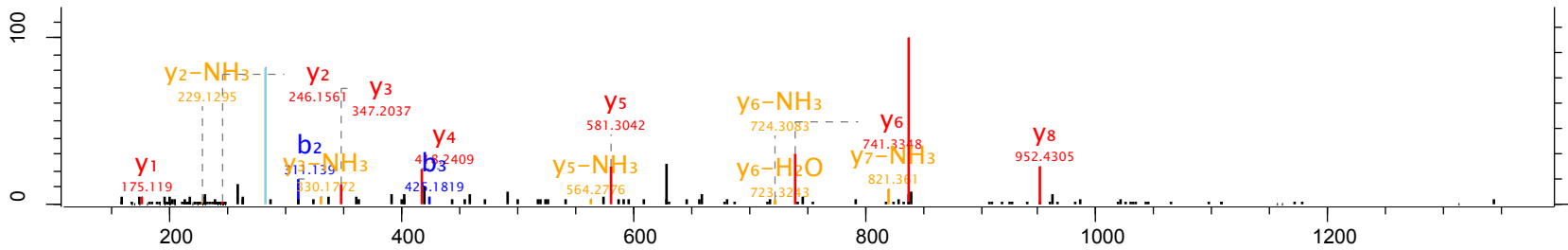
FUNDC1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract16_dyn_5ul_F8_01_595	13765	TOF; CID	88.5	522.8	TRAF5



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract16_dyn_5ul_F8_01_595	15101	TOF; CID	65.28	631.78	CRLS1



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

15536

Method

TOF; CID

Score

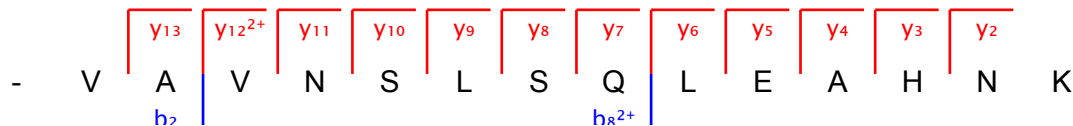
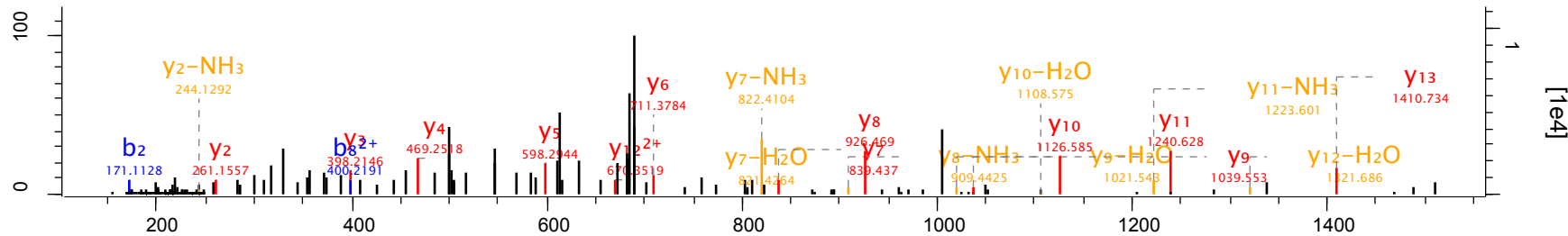
61.41

m/z

504.27

Gene names

ZNF385A;SOX6



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

15923

Method

TOF; CID

Score

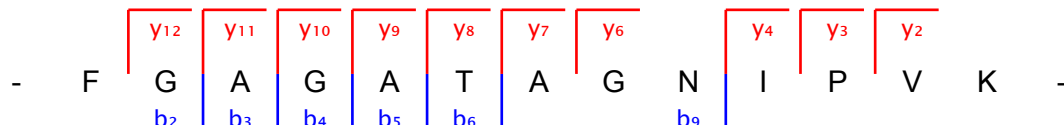
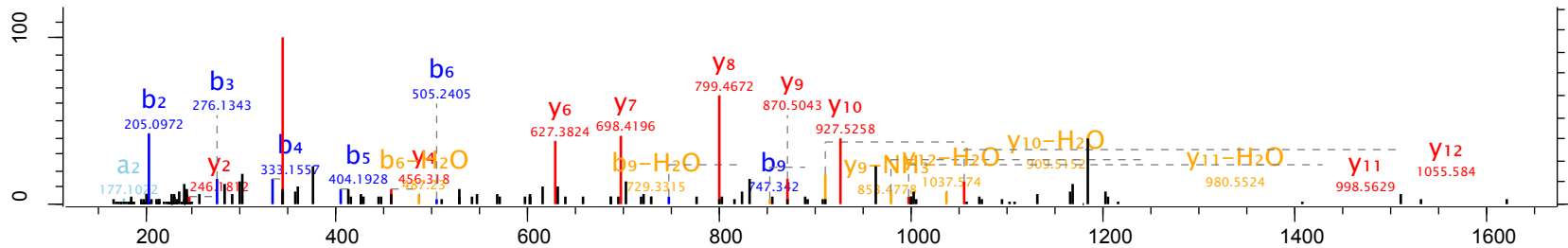
89.43

m/z

602.33

Gene names

DAZAP2



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

18528

Method

TOF; CID

Score

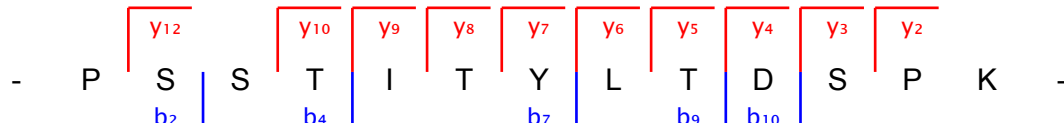
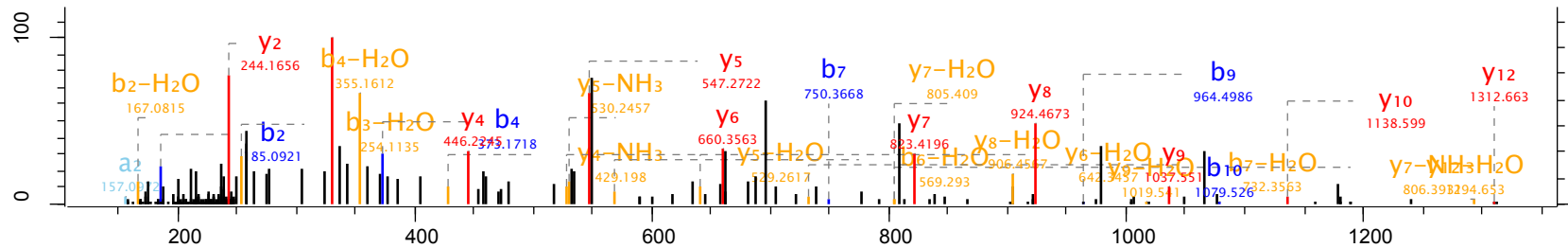
104.26

m/z

705.36

Gene names

TMEM69



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

19533

Method

TOF; CID

Score

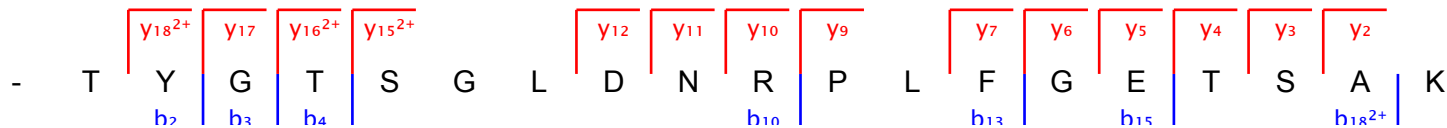
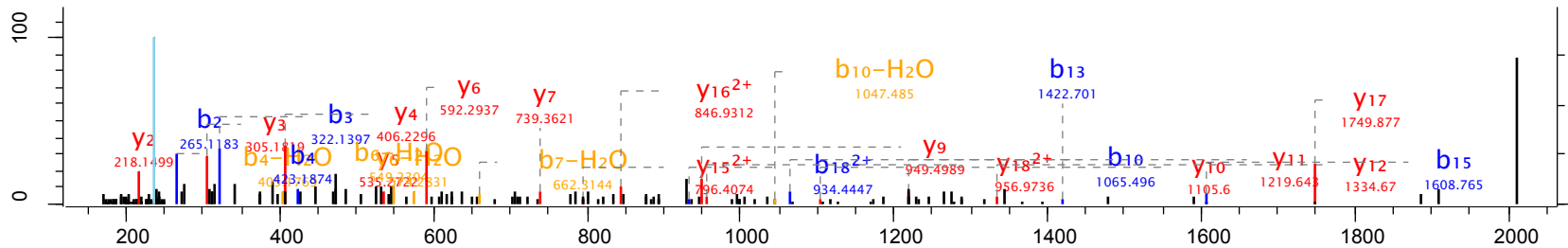
68.95

m/z

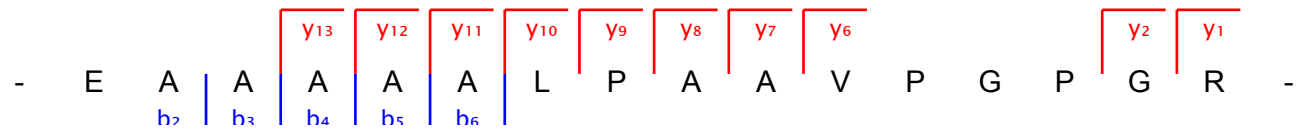
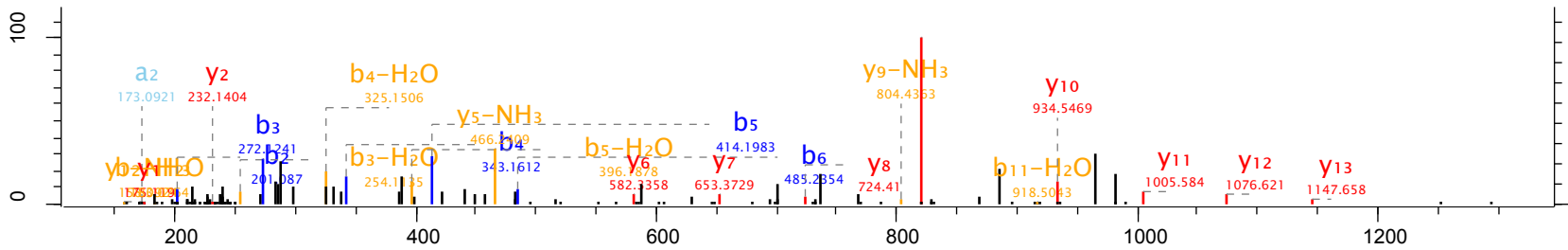
1007.49

Gene names

TMEM243



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract16_dyn_5ul_F8_01_595	20390	TOF; CID	78.51	709.89	DIRC2



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

25303

Method

TOF; CID

Score

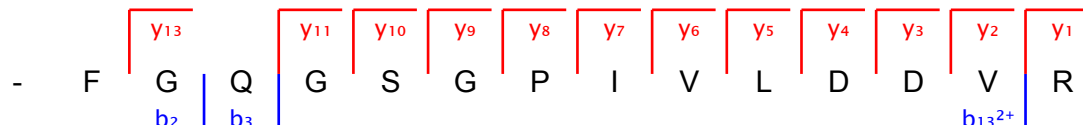
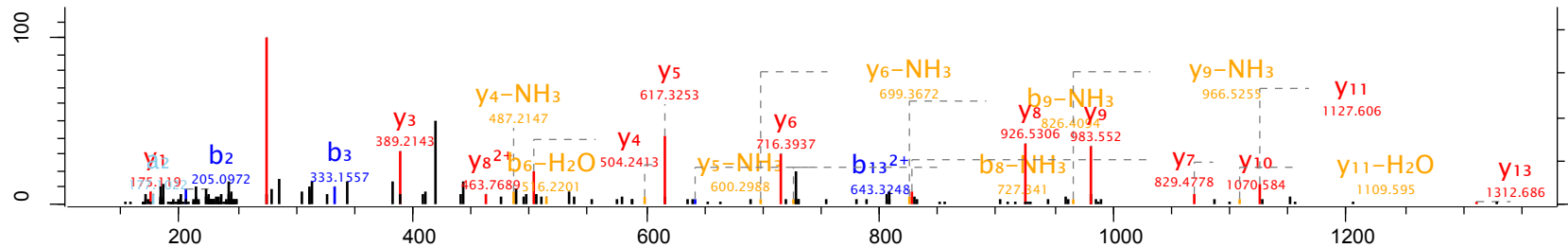
72.21

m/z

730.38

Gene names

DMBT1



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

28148

Method

TOF; CID

Score

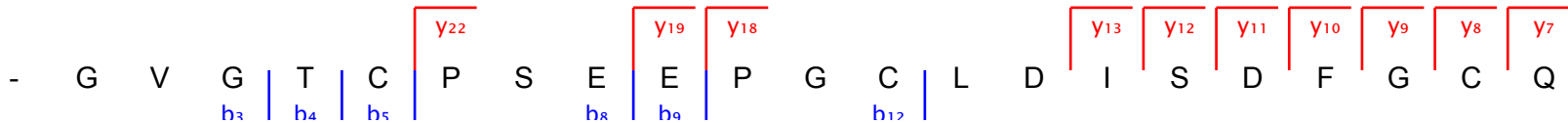
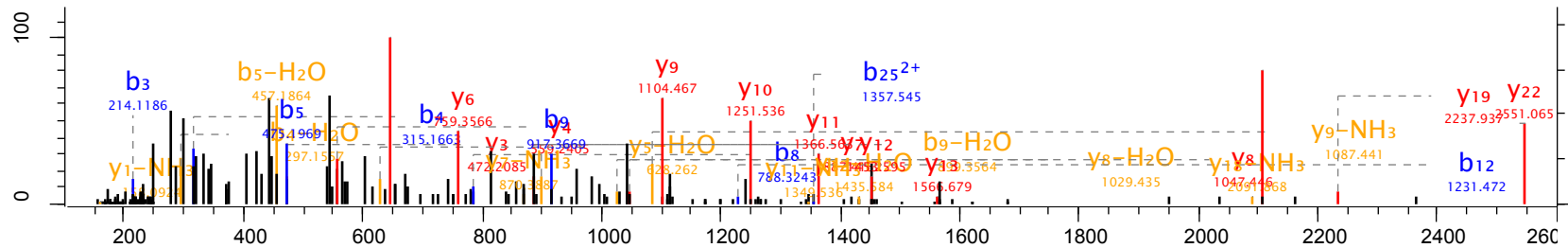
81.97

m/z

1010.09

Gene names

FAM199X



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

32575

Method

TOF; CID

Score

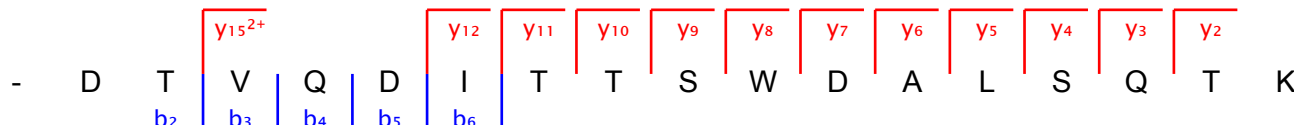
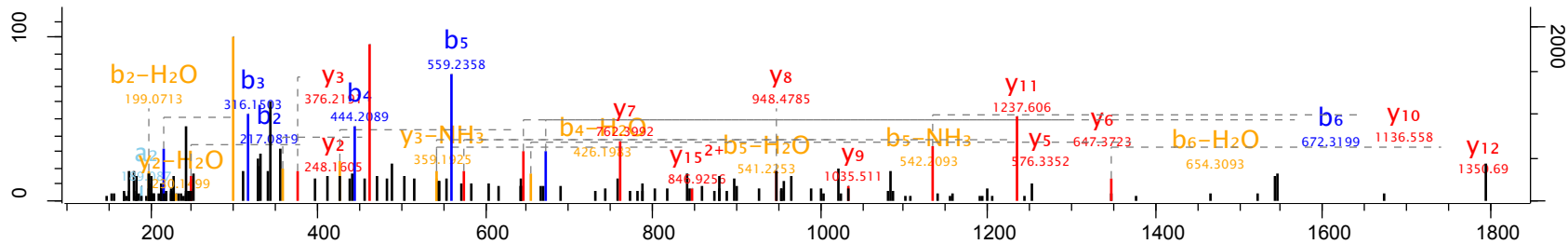
91.4

m/z

955.46

Gene names

CEP350



Raw file

20141014_fract16_dyn_5ul_F8_01_595

Scan

34265

Method

TOF; CID

Score

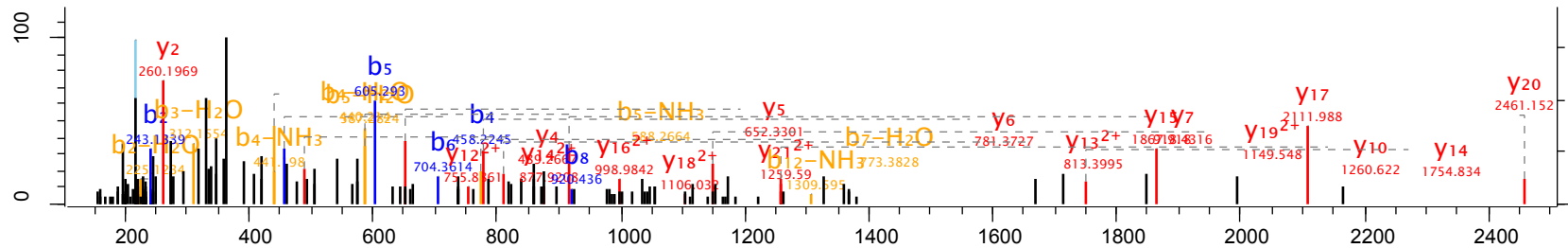
59.37

m/z

860.4

Gene names

SWI5

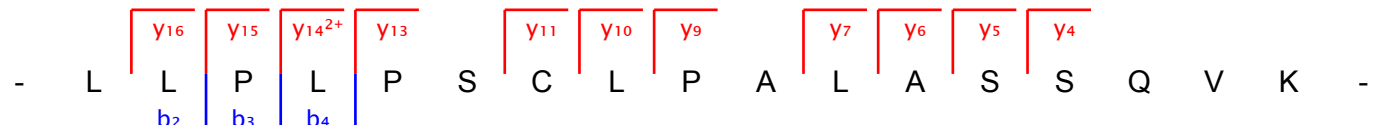
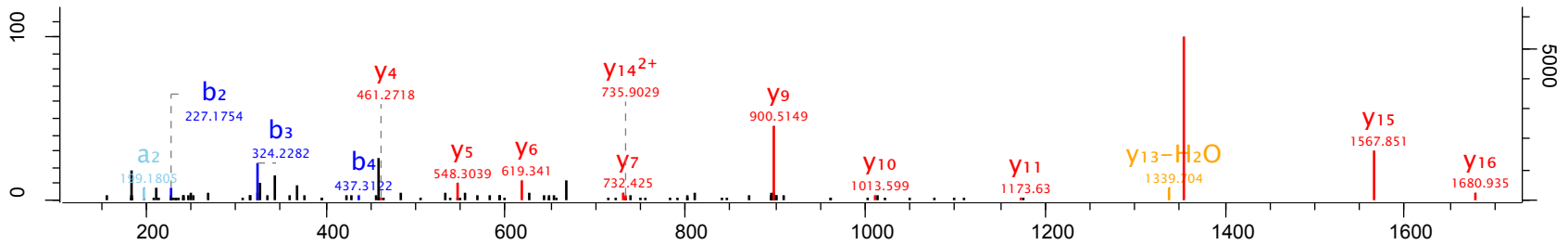


- E I S Q F V S E G Y S V D E L E D H I T Q

h₂ h₄ h₅ h₆ h₈

y₂₁²⁺ y₂₀ y₁₉²⁺ y₁₈²⁺ y₁₇ y₁₆²⁺ y₁₅ y₁₄ y₁₃²⁺ y₁₂²⁺ y₁₀

Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract16_dyn_5ul_F8_01_595	37267	TOF; CID	66.83	898.01	R3HDM4



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract17_dyn_5ul_G1_01_596

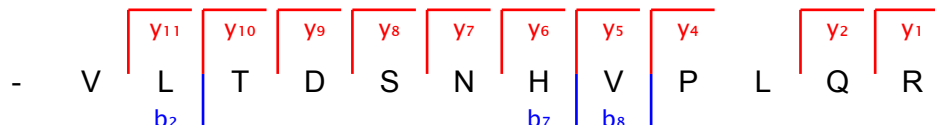
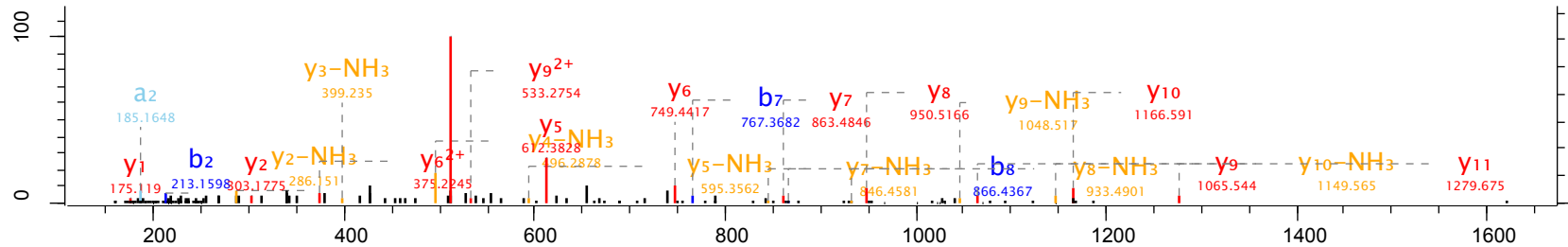
11851

TOF; CID

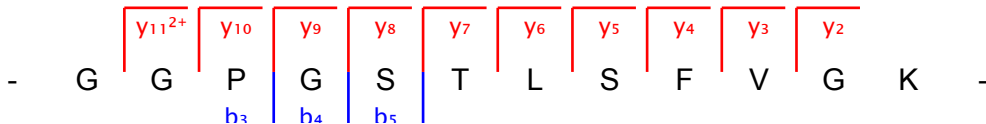
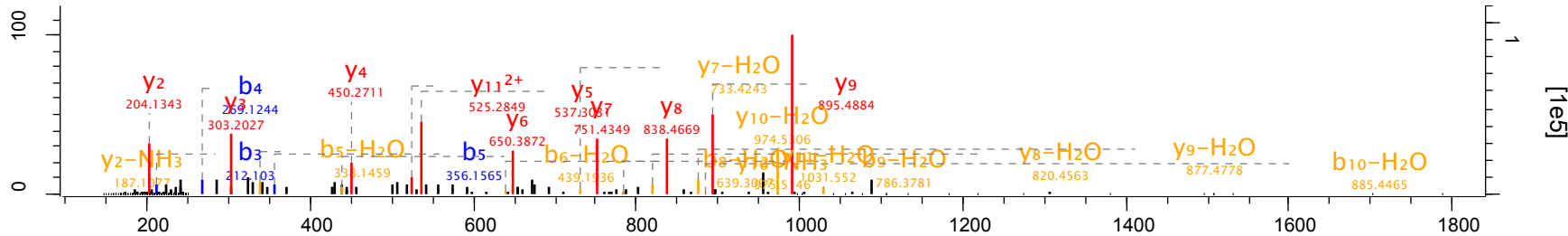
110.08

689.87

FBXO33



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract17_dyn_5ul_G1_01_596	19333	TOF; CID	91.96	553.79	C19orf43



Raw file

20141014_fract17_dyn_5ul_G1_01_596

Scan

20256

Method

TOF; CID

Score

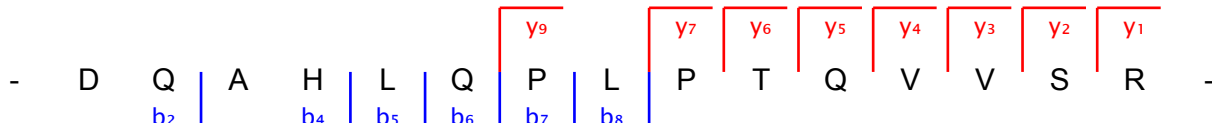
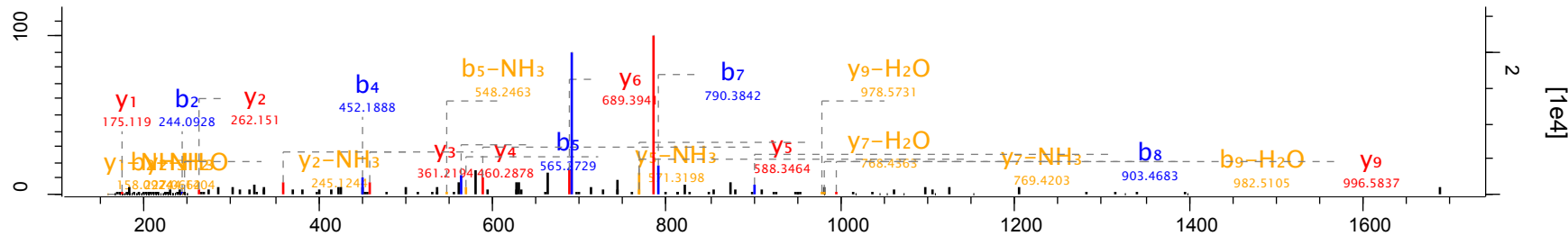
52.17

m/z

563.64

Gene names

ELF4



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract17_dyn_5ul_G1_01_596

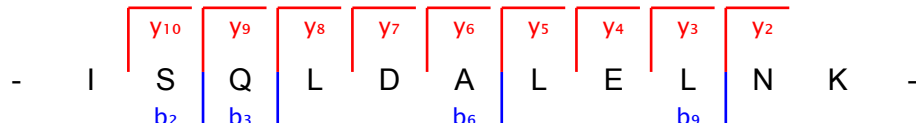
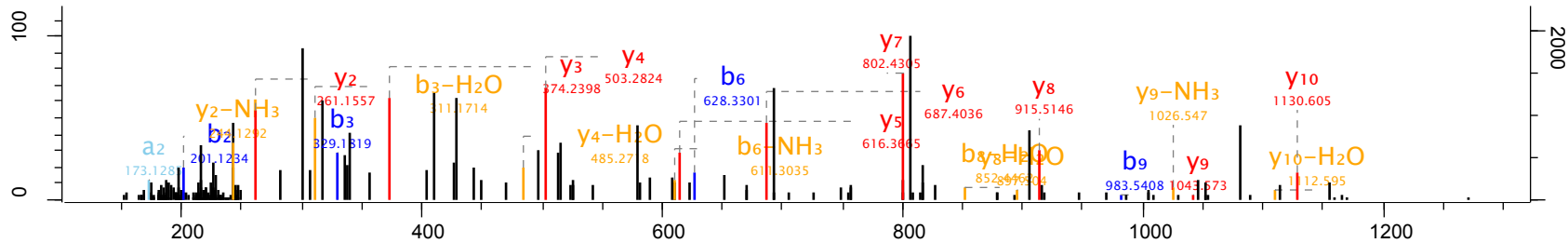
23450

TOF; CID

79.28

622.34

PEX2



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract17_dyn_5ul_G1_01_596

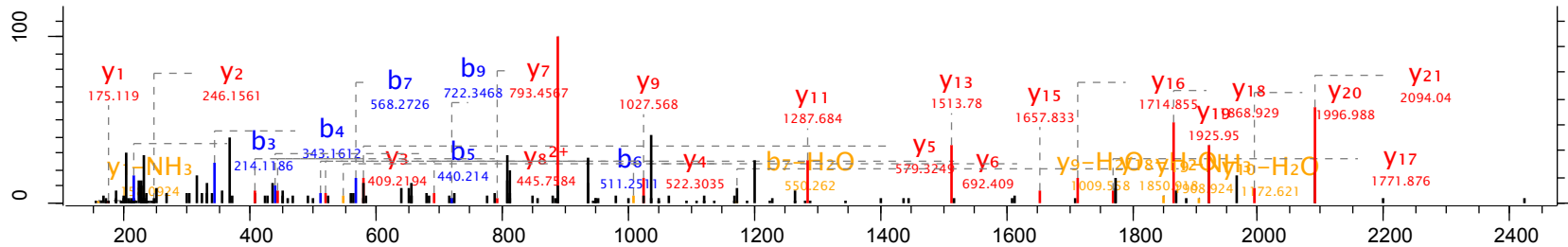
23528

TOF; CID

62.03

813.07

LHFPL2



- G G V E P A G P G G G S P E P Y H P T L G

h₃ h₄ h₅ h₆ h₇ h₉

y₂₁ y₂₀ y₁₉ y₁₈ y₁₇ y₁₆ y₁₅ y₁₃ y₁₁ y₉ y₈ y₇ y₆ y₅

Raw file

20141014_fract17_dyn_5ul_G1_01_596

Scan

23833

Method

TOF; CID

Score

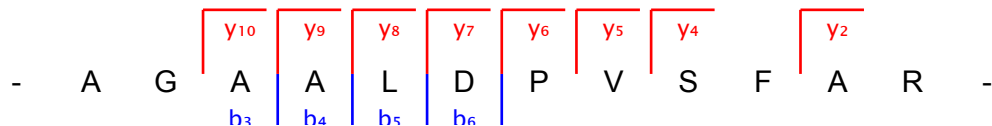
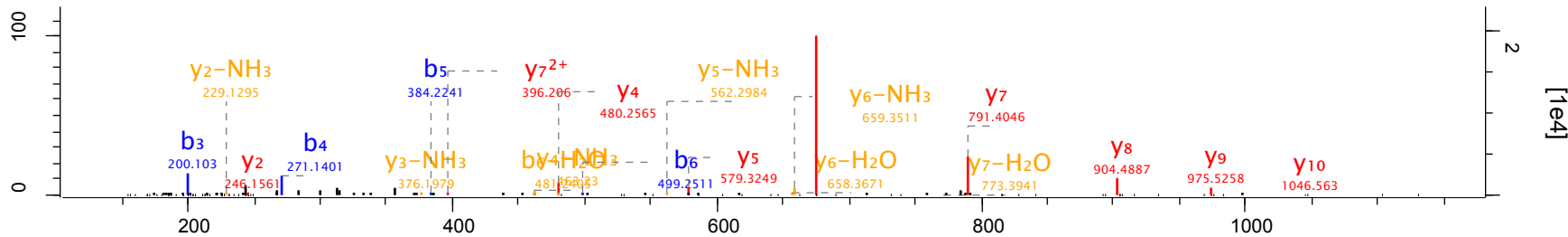
98.16

m/z

587.81

Gene names

SYNGR3



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract17_dyn_5ul_G1_01_596

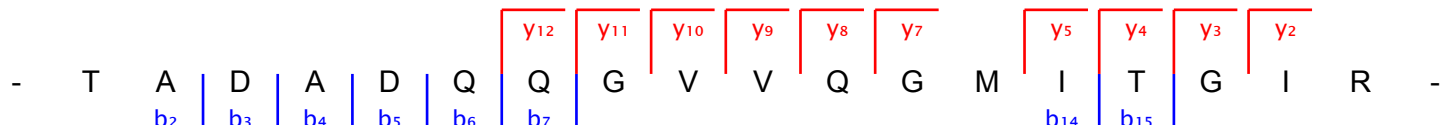
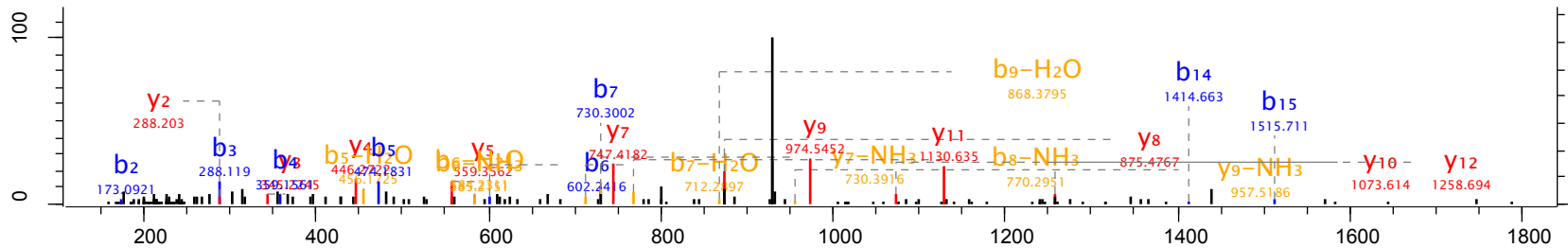
27158

TOF; CID

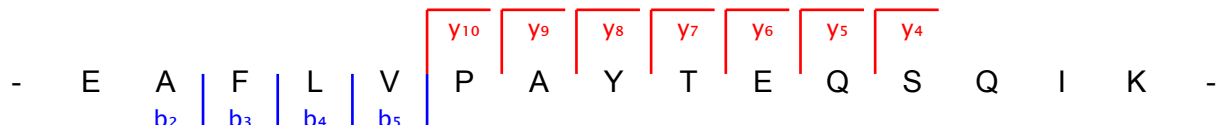
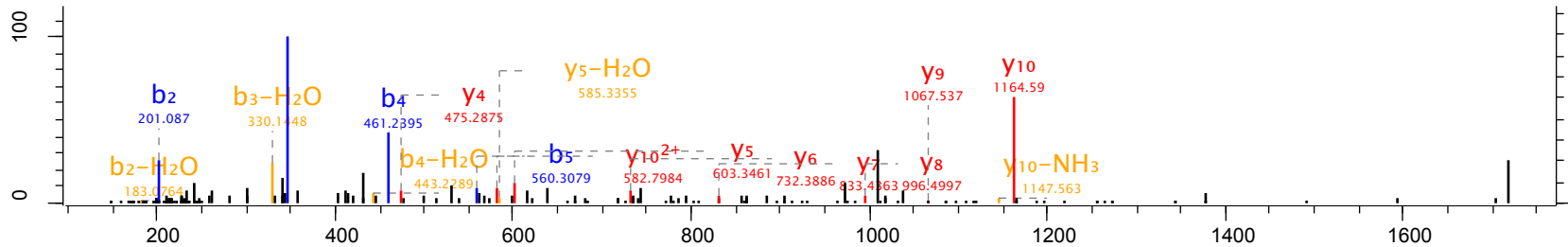
64

930.47

HIAT1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract17_dyn_5ul_G1_01_596	27778	TOF; CID	58.89	862.4	PSENN



Raw file

20141014_fract17_dyn_5ul_G1_01_596

Scan

29706

Method

TOF; CID

Score

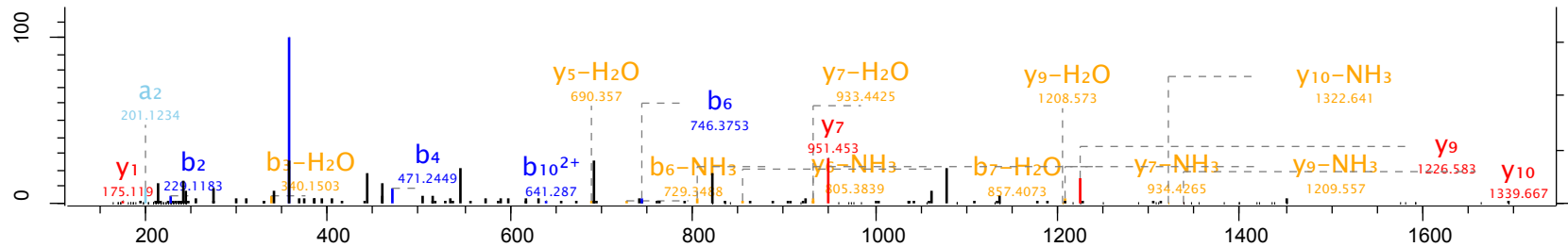
67.88

m/z

848.91

Gene names

WDR65



[1e4]



Raw file

20141014_fract17_dyn_5ul_G1_01_596

Scan

31561

Method

TOF; CID

Score

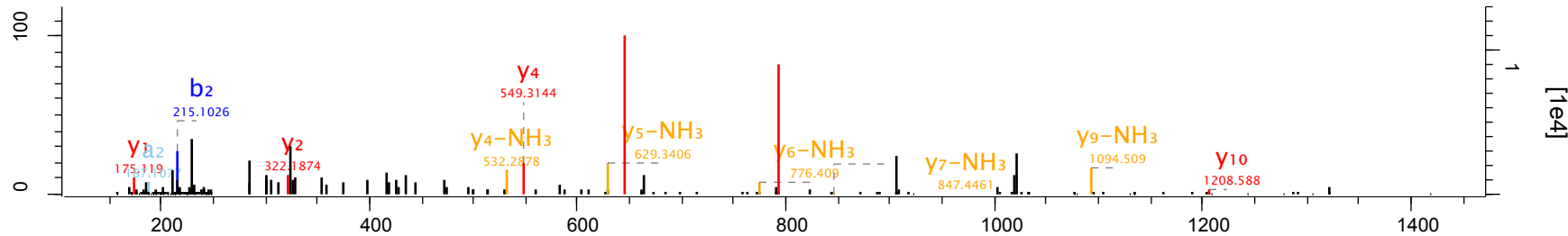
53.24

m/z

712.84

Gene names

DUSP18



ac

T A P S C A F P V Q F R -

b2

y10

y6

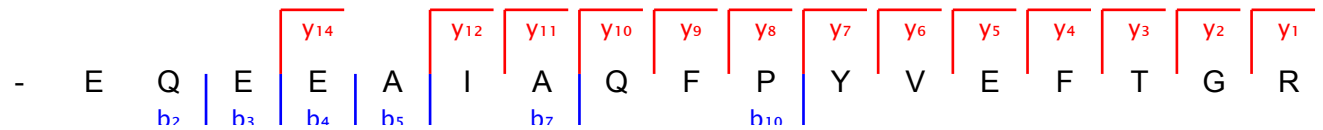
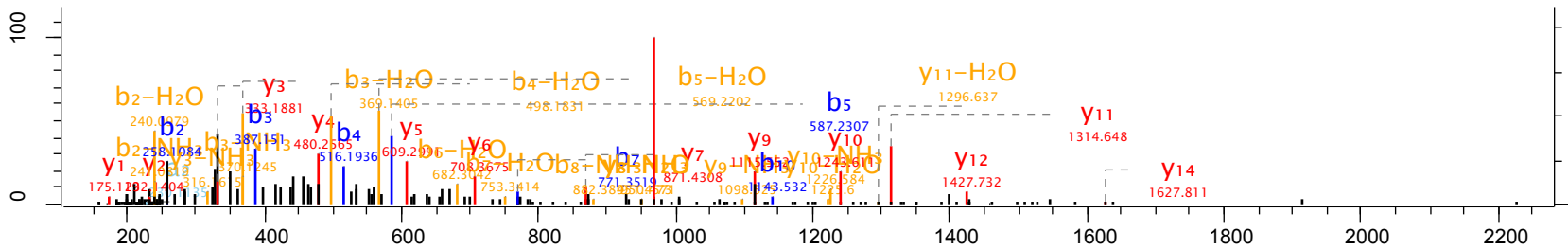
y5

y4

y2

y1

Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract17_dyn_5ul_G1_01_596	34721	TOF; CID	149.73	1007.98	TMEM106C



Raw file

20141014_fract17_dyn_5ul_G1_01_596

Scan

36308

Method

TOF; CID

Score

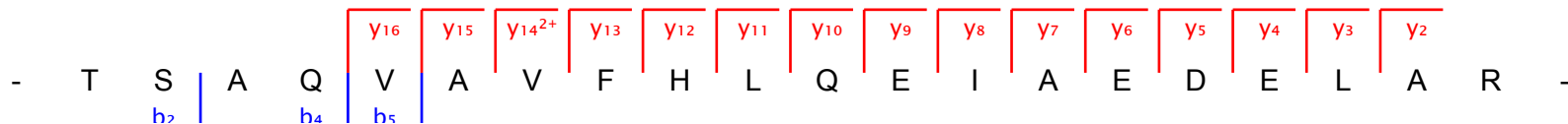
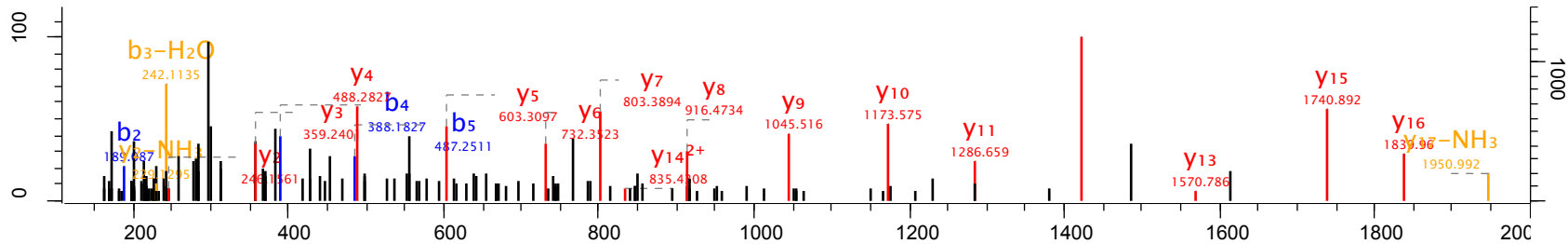
38.72

m/z

743.38

Gene names

ZNF512B



Raw file

20141014_fract17_dyn_5ul_G1_01_596

Scan

38062

Method

TOF; CID

Score

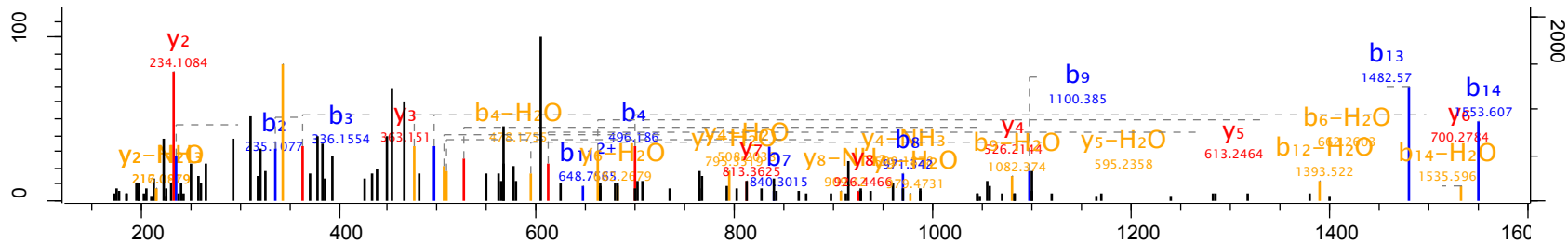
52.46

m/z

1240.52

Gene names

RNF11



- S F T C P S C M E P V D A A L L S S Y E T

b_2 | b_3 | b_4 | b_7 | b_8 | b_9 | b_{11}^{2+} | b_{13} | b_{14}

y_8 | y_7 | y_6 | y_5 | y_4 | y_3 | y_2

Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract18_dyn_5ul_G2_01_597

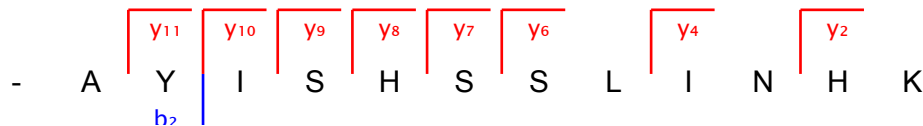
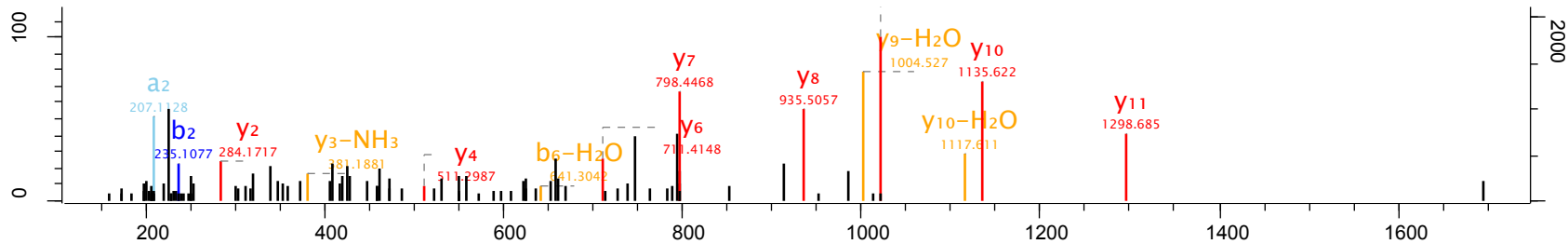
8120

TOF; CID

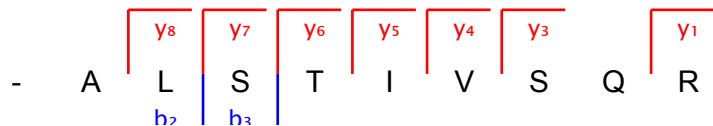
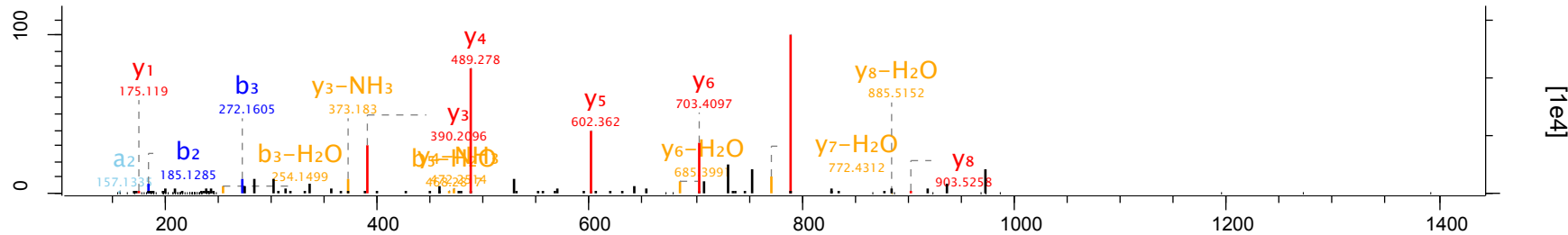
56.2

457.24

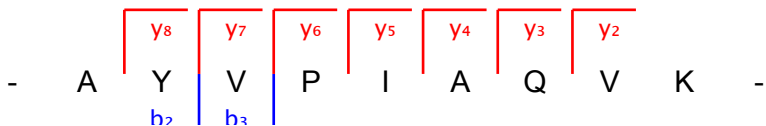
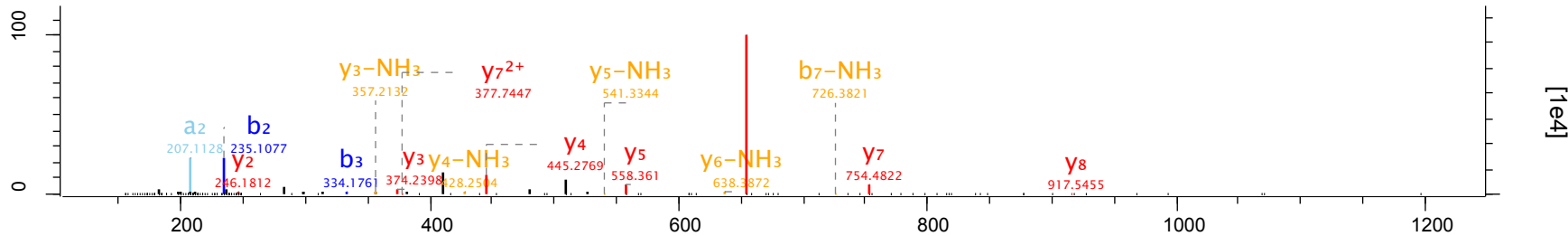
ZFP62



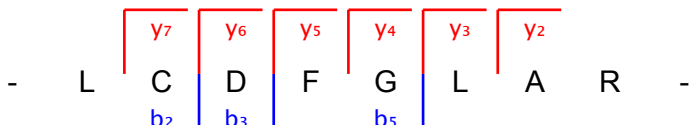
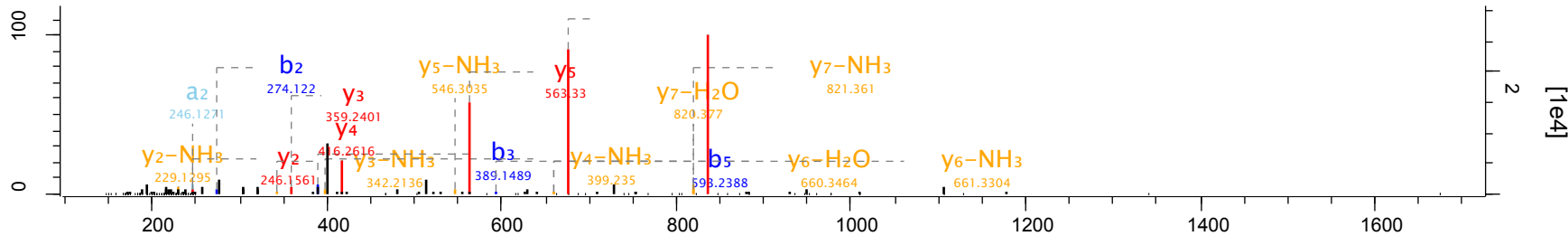
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract18_dyn_5ul_G2_01_597	12482	TOF; CID	80.23	487.78	CENPW



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract18_dyn_5ul_G2_01_597	16907	TOF; CID	106.04	494.79	GPNMB



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract18_dyn_5ul_G2_01_597	17983	TOF; CID	120.92	476.24	MAPK15



Raw file

20141014_fract18_dyn_5ul_G2_01_597

Scan

23835

Method

TOF; CID

Score

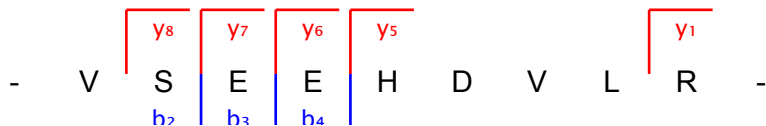
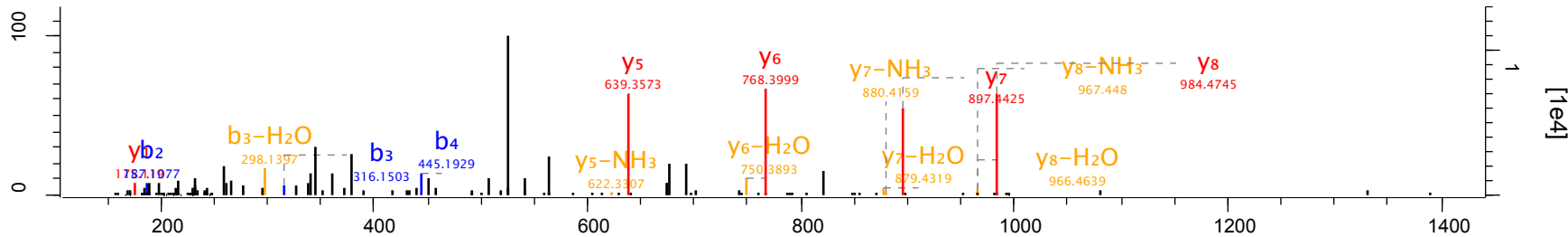
83.95

m/z

542.78

Gene names

HCLS1



Raw file

Scan

Method

Score

m/z

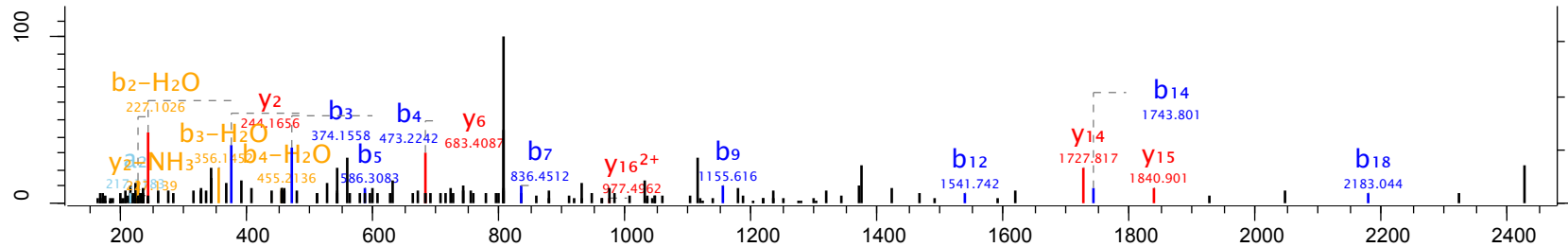
20141014_fract18_dyn_5ul_G2_01_597

26720

TOF; CID

50.36

809.4



ac

- T T E V I L H Y R P C E S D P T Q L P K -

a₂ b₃ b₄ b₅ b₇ b₉ b₁₂ b₁₄ b₁₈

y₁₆²⁺ y₁₅ y₁₄ y₆ y₂

Raw file

20141014_fract18_dyn_5ul_G2_01_597

Scan

27725

Method

TOF; CID

Score

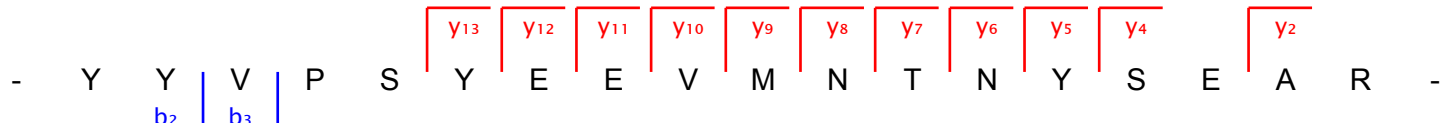
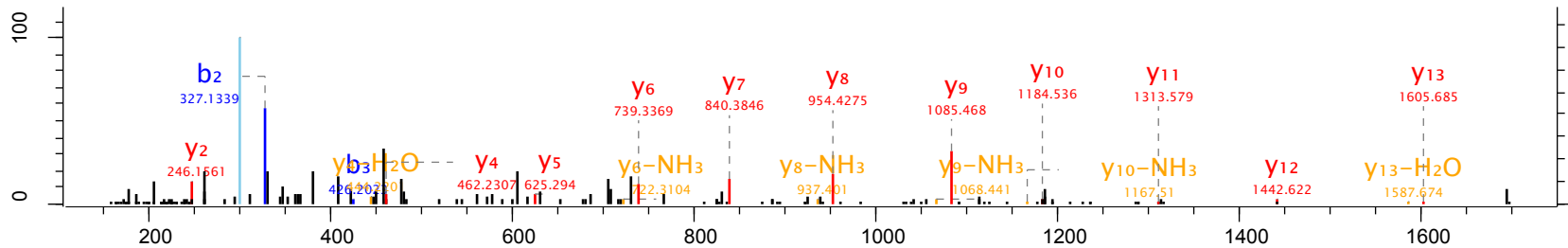
57.94

m/z

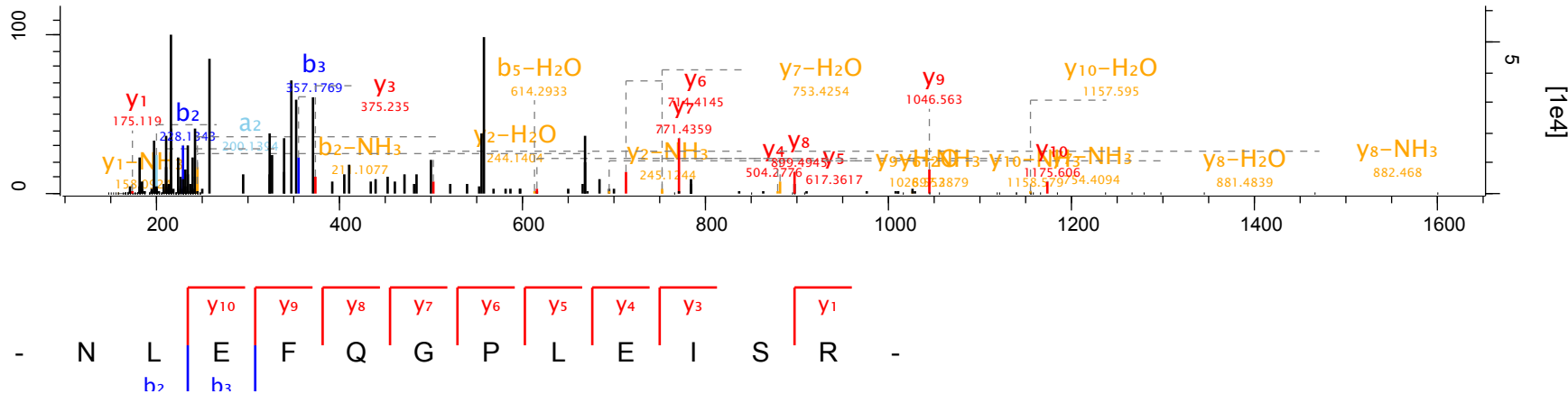
1108.99

Gene names

TMEM51



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract18_dyn_5ul_G2_01_597	28110	TOF; CID	59.8	702.37	VPS37C



Raw file

20141014_fract18_dyn_5ul_G2_01_597

Scan

28746

Method

TOF; CID

Score

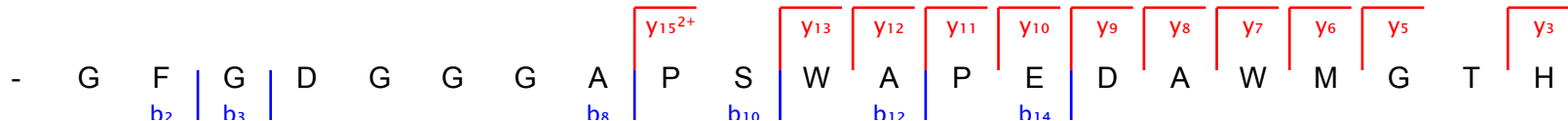
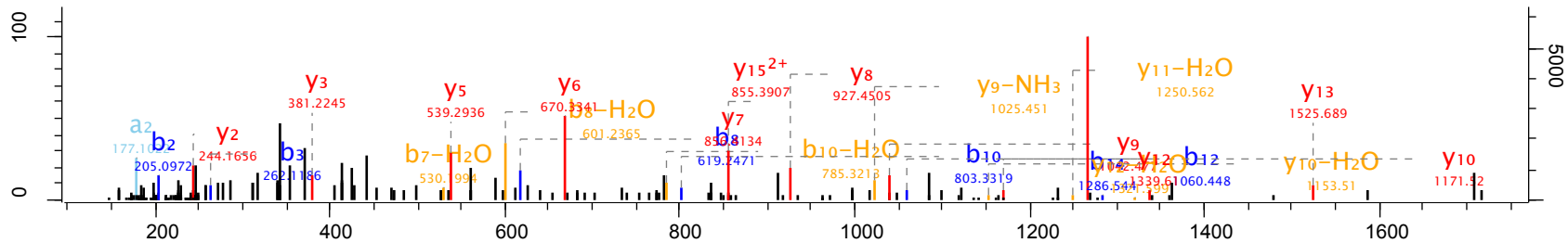
62.1

m/z

777.33

Gene names

TSEN15



Raw file

20141014_fract18_dyn_5ul_G2_01_597

Scan

29037

Method

TOF; CID

Score

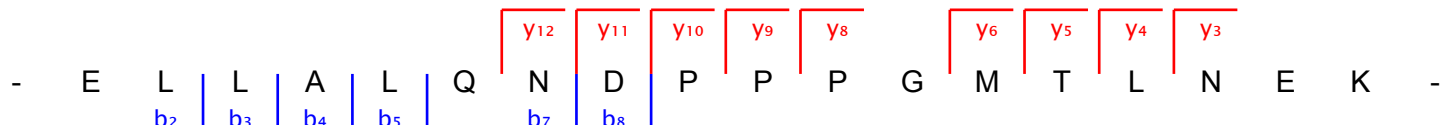
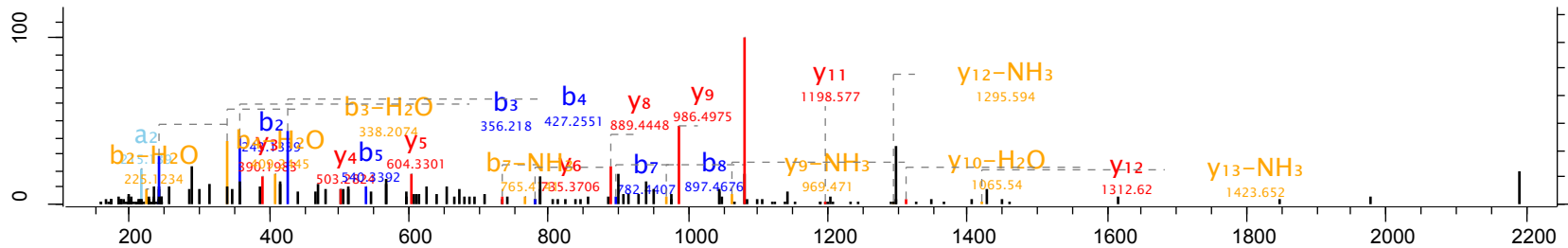
68.24

m/z

990.51

Gene names

UBE2W



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract18_dyn_5ul_G2_01_597

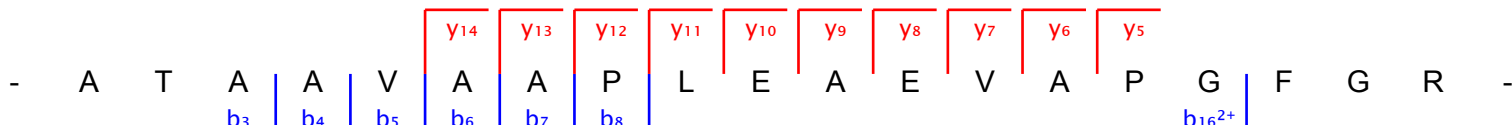
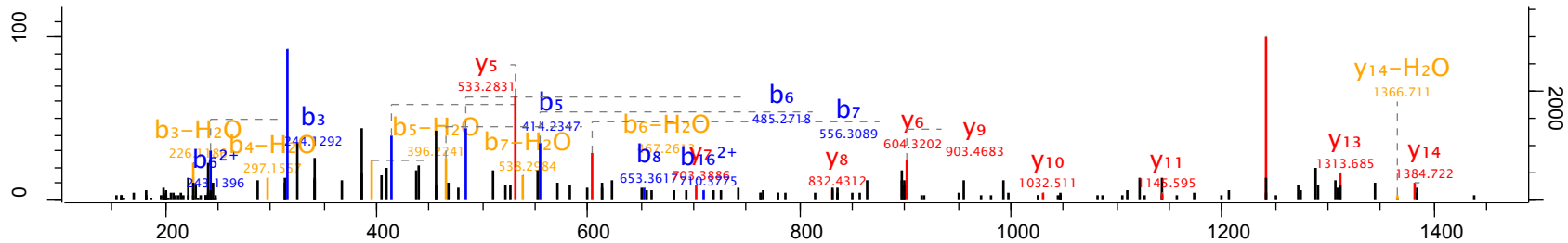
29448

TOF; CID

82.45

899.98

BEGAIN



Raw file

20141014_fract18_dyn_5ul_G2_01_597

Scan

30228

Method

TOF; CID

Score

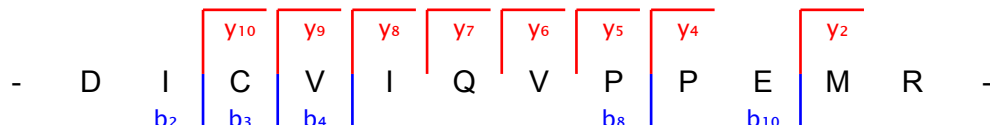
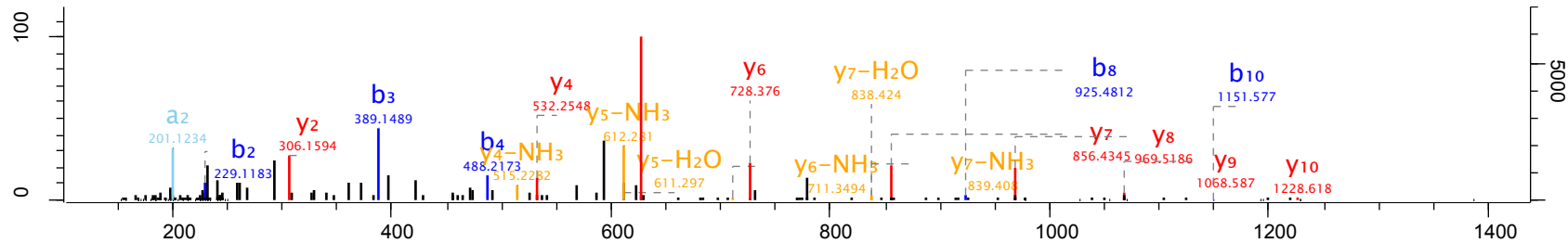
108.72

m/z

728.87

Gene names

MALSU1



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract18_dyn_5ul_G2_01_597

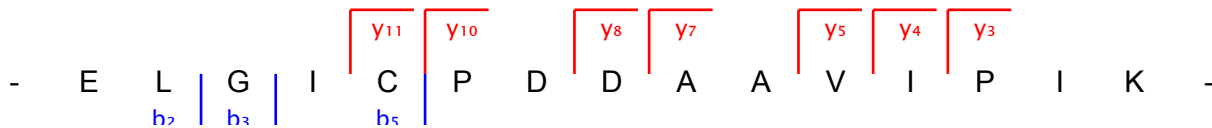
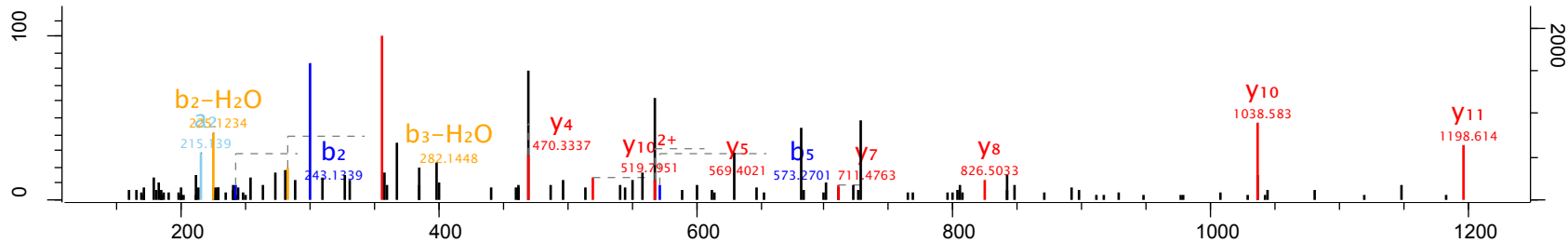
31765

TOF; CID

47.04

805.93

PIP



Raw file

20141014_fract18_dyn_5ul_G2_01_597

Scan

32329

Method

TOF; CID

Score

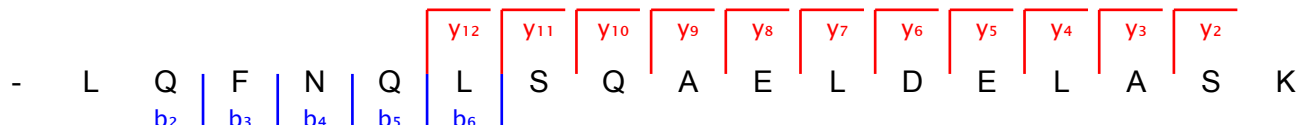
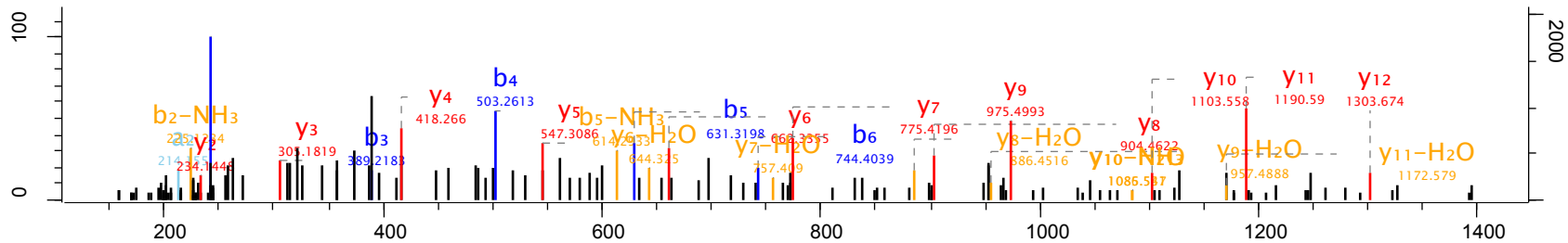
96.1

m/z

968

Gene names

EFHC1



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract18_dyn_5ul_G2_01_597

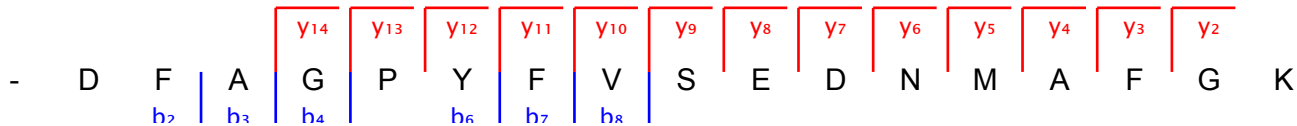
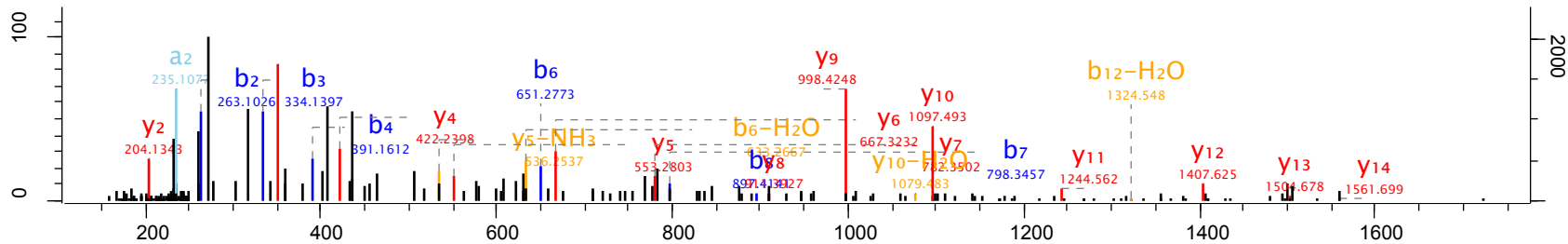
33912

TOF; CID

78.44

948.42

TMEM222



Raw file

20141014_fract18_dyn_5ul_G2_01_597

Scan

34439

Method

TOF; CID

Score

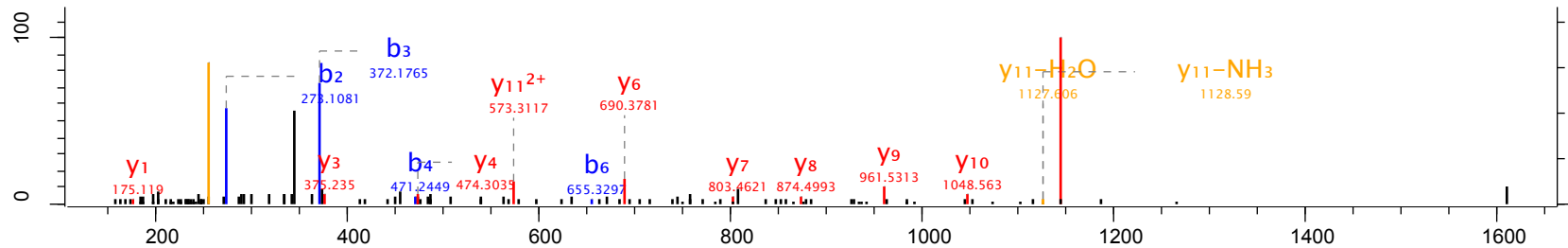
60.55

m/z

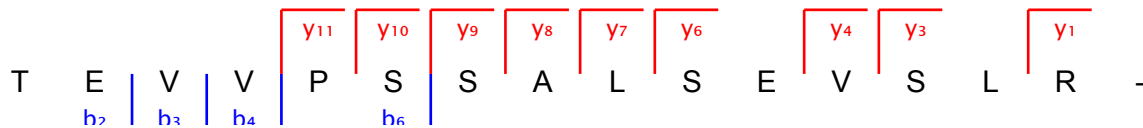
808.43

Gene names

NAA60



ac



Raw file

20141014_fract19_dyn_5ul_G3_01_598

Scan

10366

Method

TOF; CID

Score

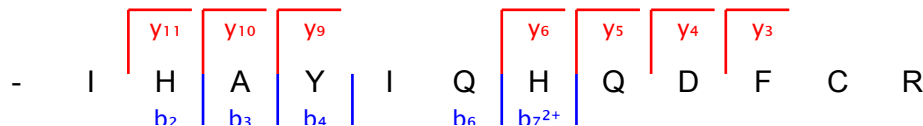
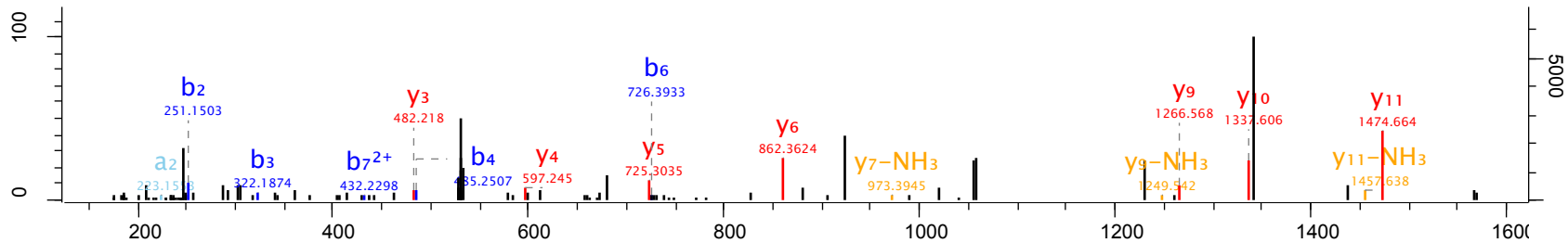
55.04

m/z

530.25

Gene names

MGAT5B



Raw file

20141014_fract19_dyn_5ul_G3_01_598

Scan

14294

Method

TOF; CID

Score

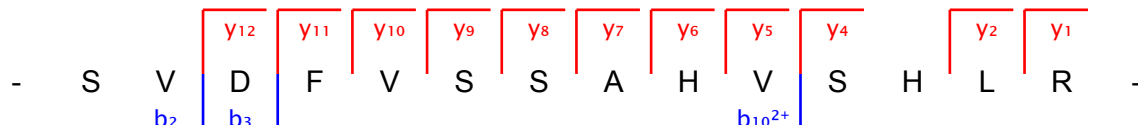
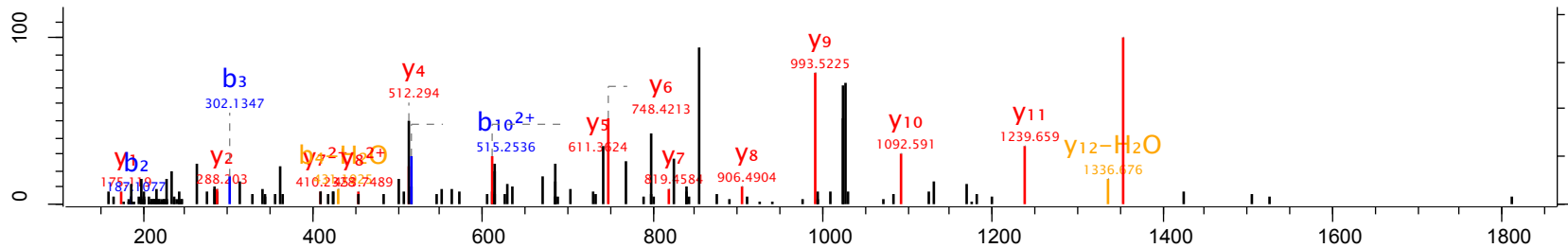
44.81

m/z

514.26

Gene names

TMEM53



Raw file

20141014_fract19_dyn_5ul_G3_01_598

Scan

18406

Method

TOF; CID

Score

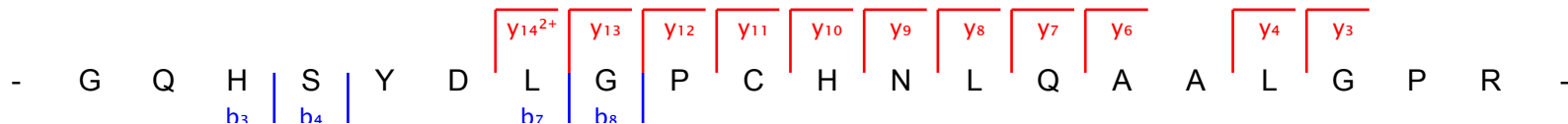
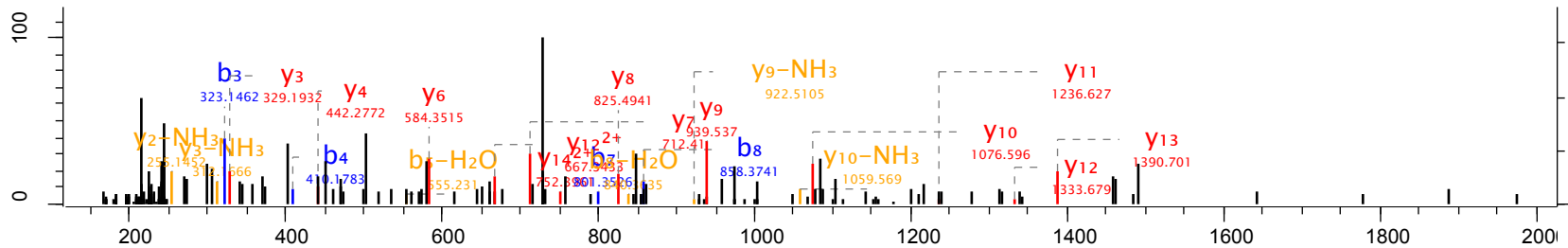
34.4

m/z

731.36

Gene names

ZDHHC24



Raw file

20141014_fract19_dyn_5ul_G3_01_598

Scan

22302

Method

TOF; CID

Score

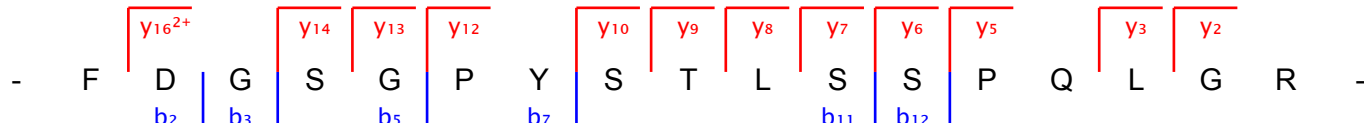
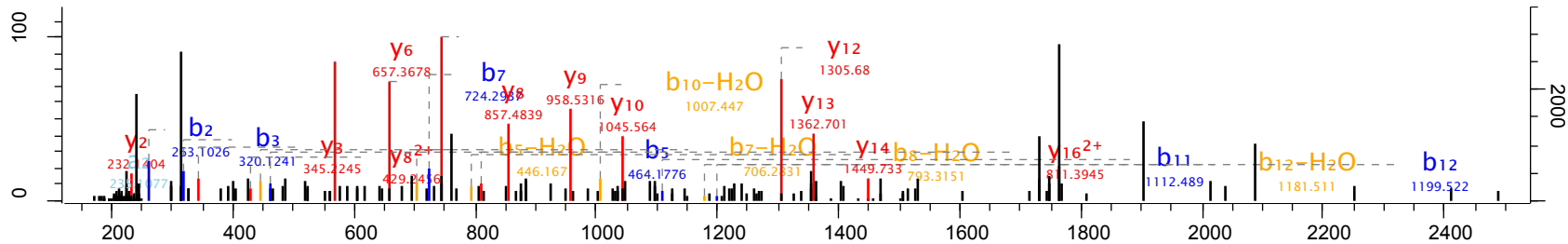
70.78

m/z

885.42

Gene names

DNASE1L1



Raw file

20141014_fract19_dyn_5ul_G3_01_598

Scan

22540

Method

TOF; CID

Score

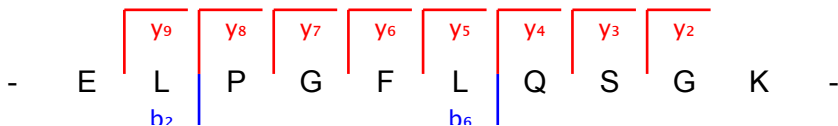
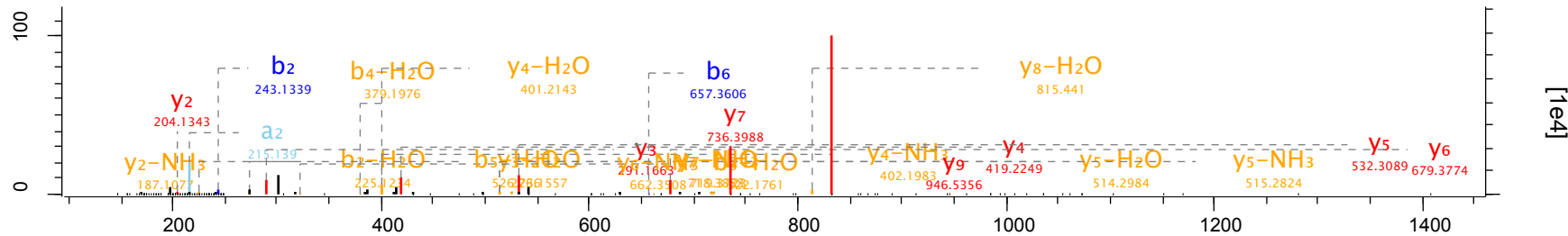
99.57

m/z

538.29

Gene names

S100P



Raw file

20141014_fract19_dyn_5ul_G3_01_598

Scan

27711

Method

TOF; CID

Score

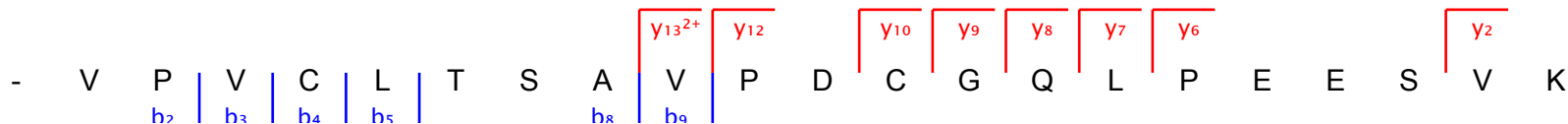
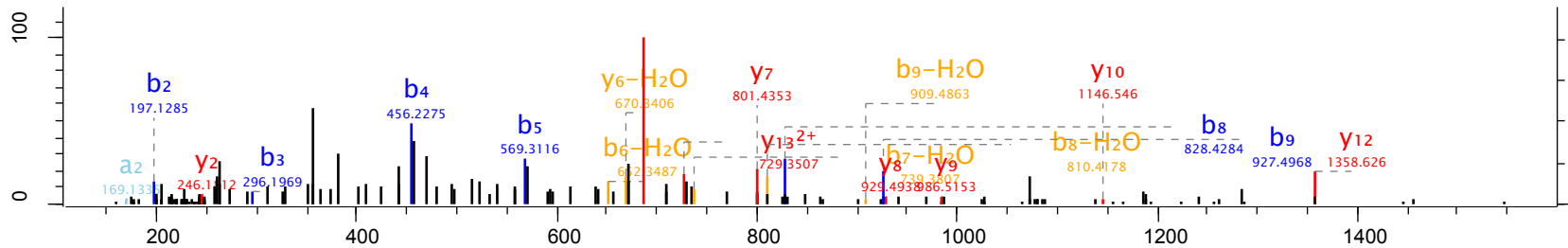
50.96

m/z

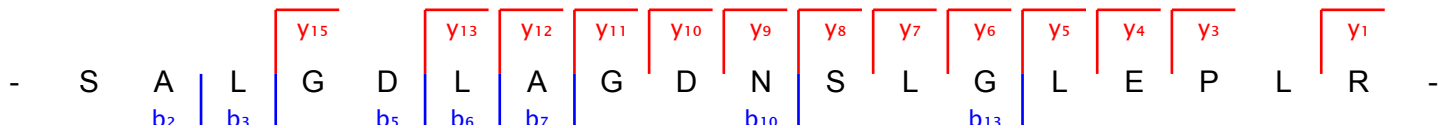
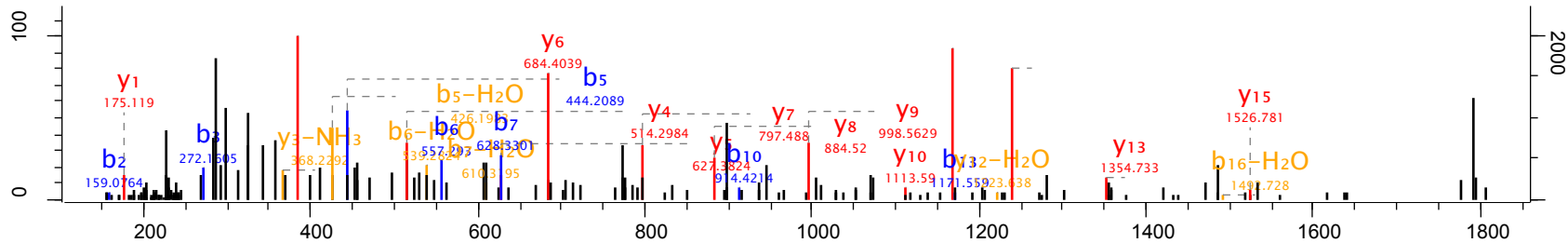
1143.56

Gene names

CDON



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract19_dyn_5ul_G3_01_598	32776	TOF; CID	69.75	899.47	RNF130



Raw file

20141014_fract19_dyn_5ul_G3_01_598

Scan

38338

Method

TOF; CID

Score

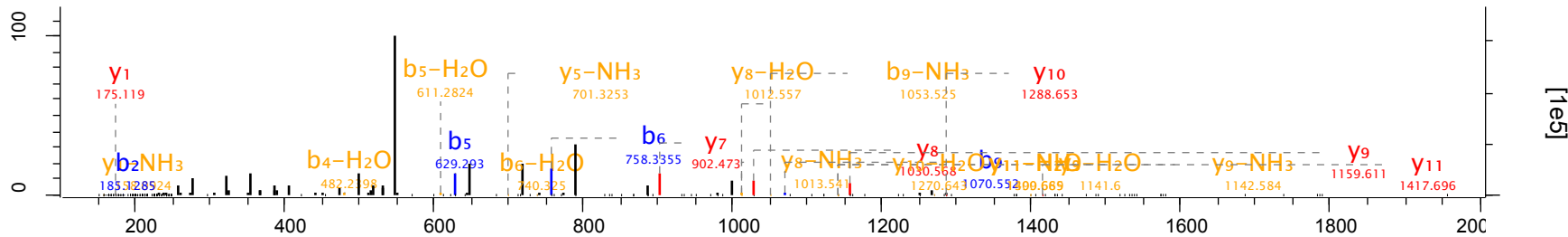
74.54

m/z

894.45

Gene names

CCDC70



- A L W E E E K A L W V E E R -

b₂ | b₅ | b₆ | b₉

y₁₁ | y₁₀ | y₉ | y₈ | y₇ | y₁

Raw file

20141014_fract20_dyn_5ul_G4_01_599

Scan

15417

Method

TOF; CID

Score

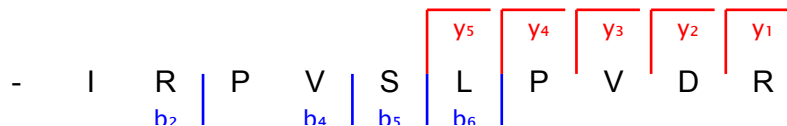
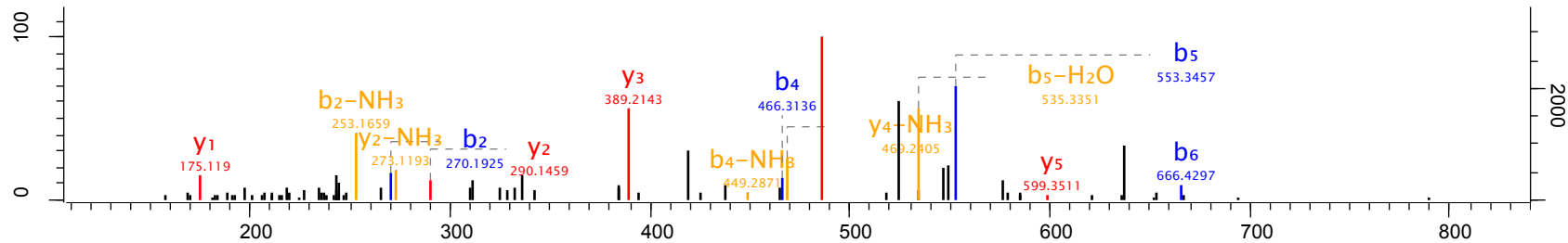
63.62

m/z

384.57

Gene names

ARHGAP29



Raw file

20141014_fract20_dyn_5ul_G4_01_599

Scan

16215

Method

TOF; CID

Score

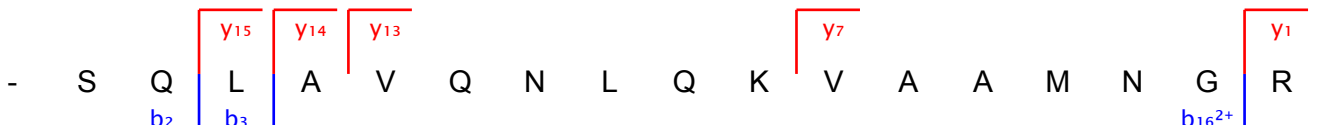
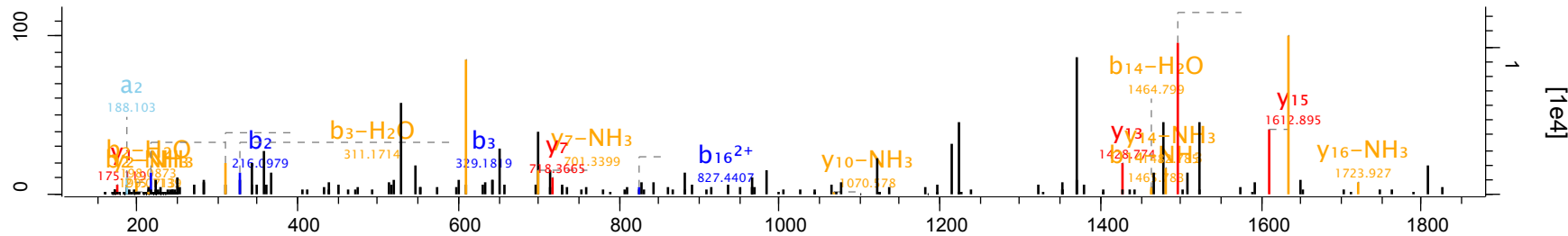
44.69

m/z

610.33

Gene names

SLC22A20



Raw file

20141014_fract20_dyn_5ul_G4_01_599

Scan

24655

Method

TOF; CID

Score

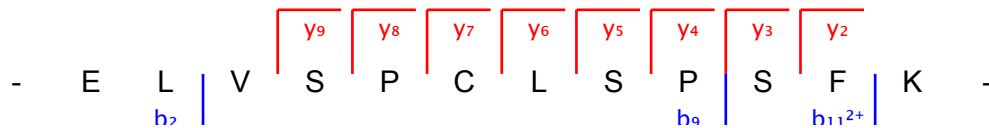
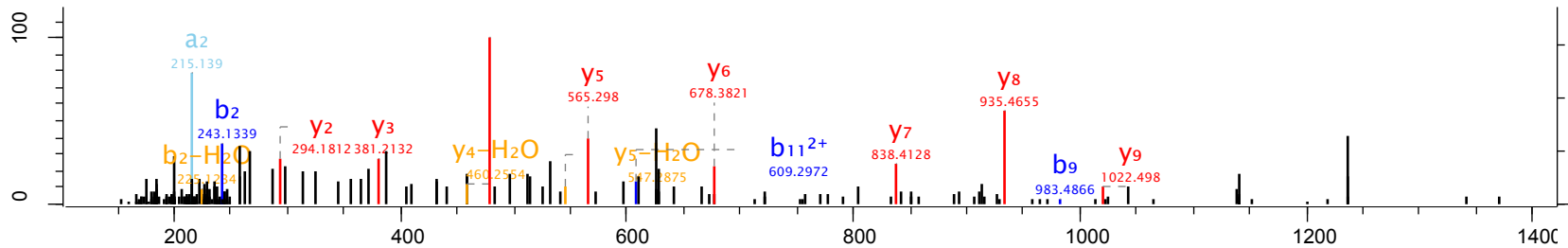
52.07

m/z

682.35

Gene names

ENTPD6



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract20_dyn_5ul_G4_01_599

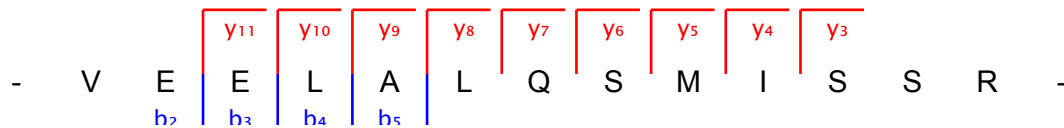
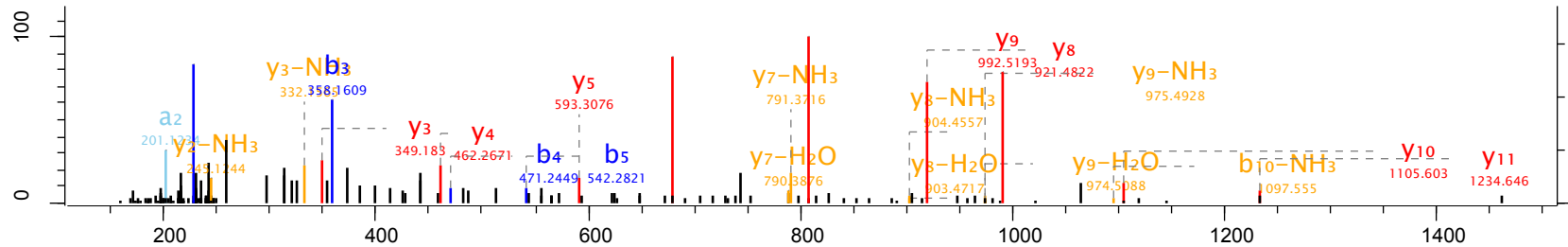
28236

TOF; CID

98.94

731.88

SNAPC5



Raw file

20141014_fract20_dyn_5ul_G4_01_599

Scan

28849

Method

TOF; CID

Score

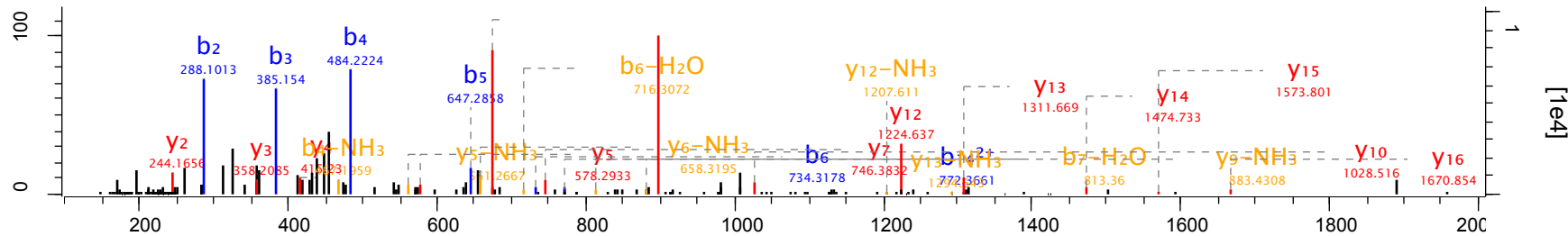
80.54

m/z

980.47

Gene names

FAM168A



ac

- M N P V Y S P V Q P G A P Y G N P K -
 b₂ b₃ b₄ b₅ b₆ b₁₄²⁺ y₁₆ y₁₅ y₁₄ y₁₃ y₁₂ y₁₀ y₉ y₇ y₆ y₅ y₄ y₃ y₂

Raw file

20141014_fract20_dyn_5ul_G4_01_599

Scan

29642

Method

TOF; CID

Score

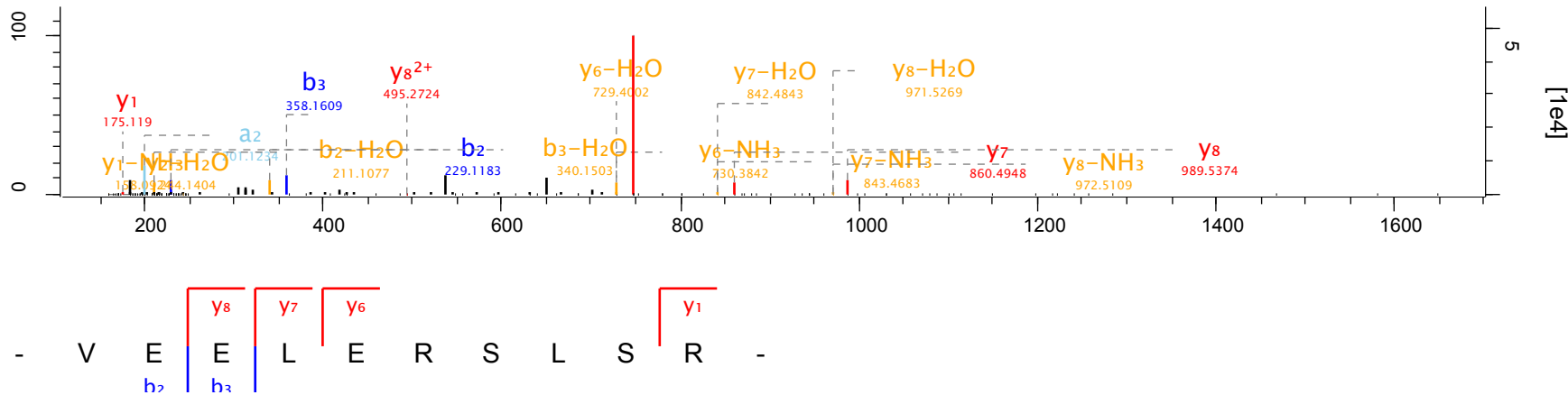
93.18

m/z

609.33

Gene names

GOLGA8A;GOLGA8B



Raw file

20141014_fract20_dyn_5ul_G4_01_599

Scan

31462

Method

TOF; CID

Score

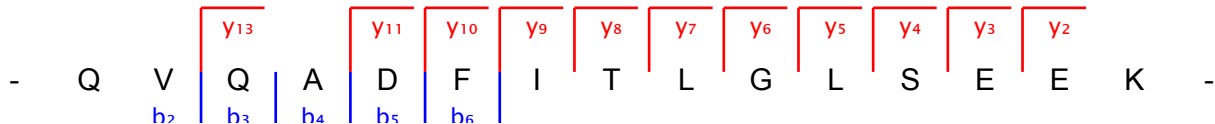
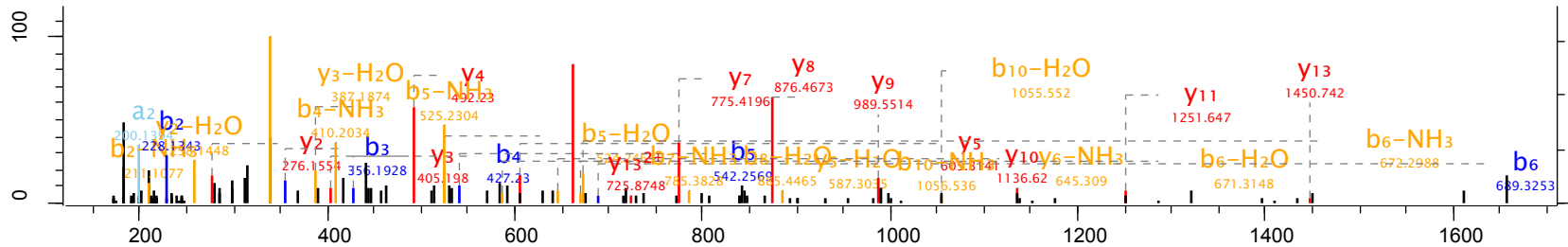
100.43

m/z

840.44

Gene names

COMMD7



Raw file

20141014_fract20_dyn_5ul_G4_01_599

Scan

33602

Method

TOF; CID

Score

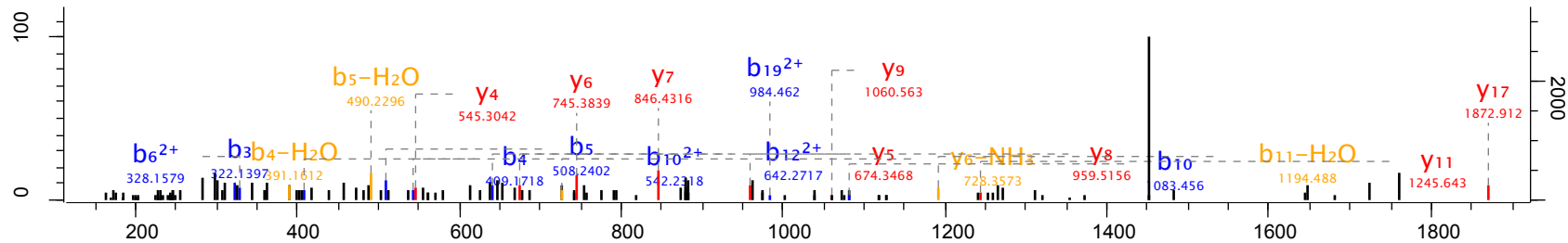
36.08

m/z

881.42

Gene names

SAYSD1



Raw file

20141014_fract21_dyn_5ul_G5_01_600

Scan

6457

Method

TOF; CID

Score

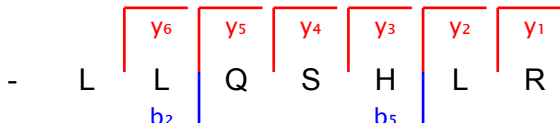
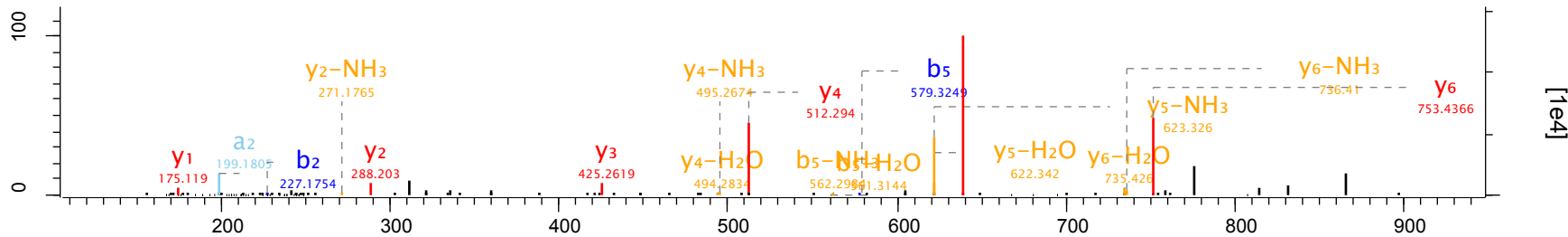
138.37

m/z

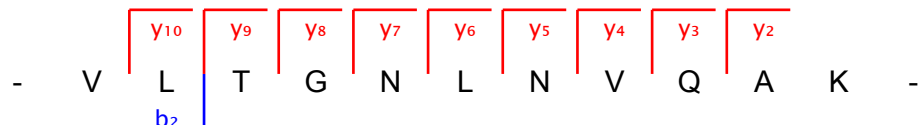
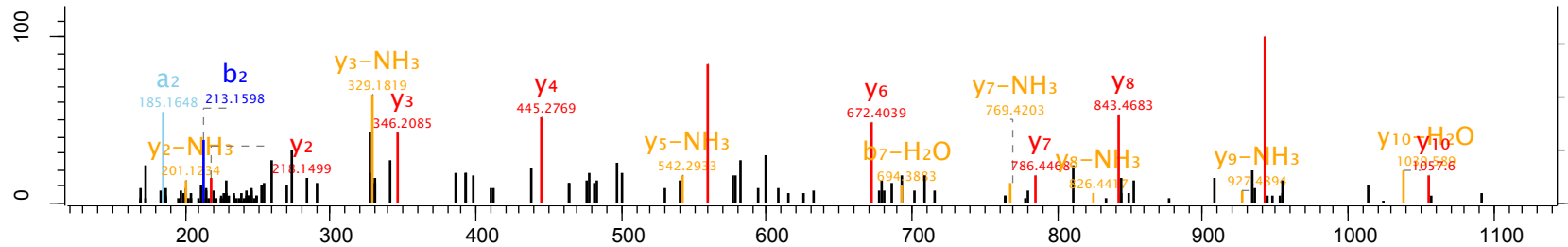
433.76

Gene names

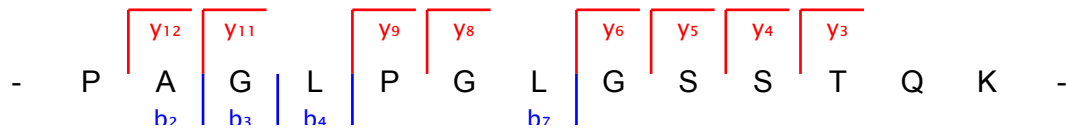
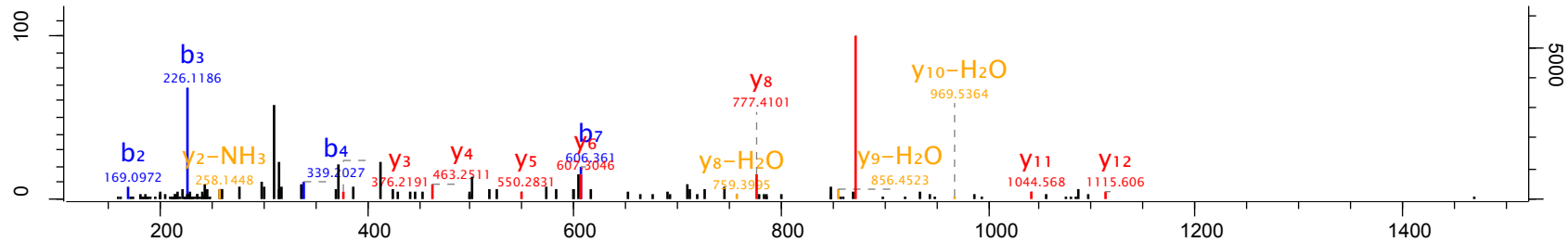
SLC39A1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract21_dyn_5ul_G5_01_600	13464	TOF; CID	91.66	578.84	KIAA1107



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract21_dyn_5ul_G5_01_600	15422	TOF; CID	50.9	606.83	AGBL5



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract21_dyn_5ul_G5_01_600

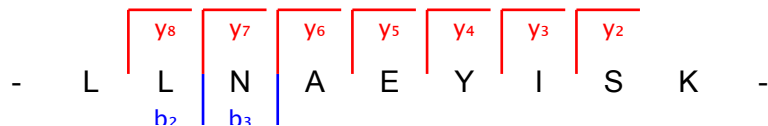
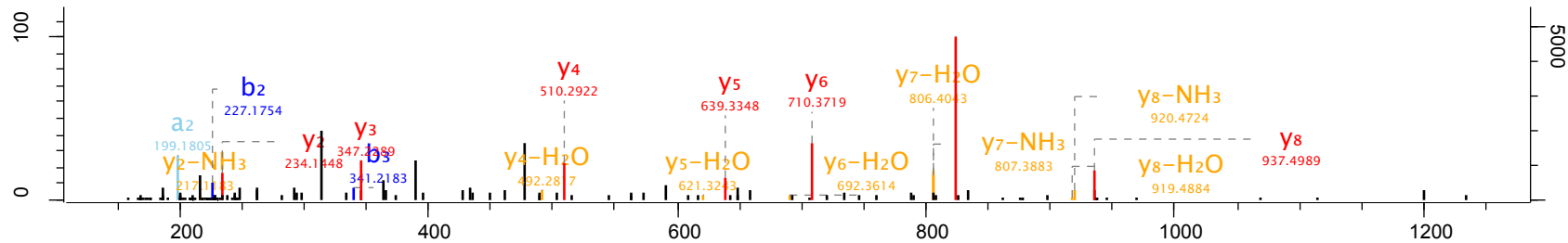
15509

TOF; CID

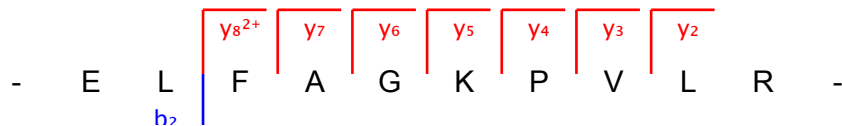
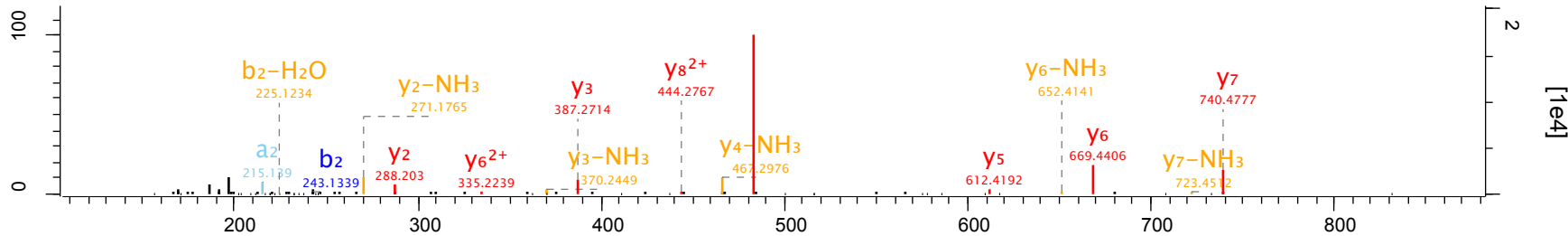
90.15

525.79

ABLIM2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract21_dyn_5ul_G5_01_600	16497	TOF; CID	76.22	377.23	NIPA2



Raw file

20141014_fract21_dyn_5ul_G5_01_600

Scan

17422

Method

TOF; CID

Score

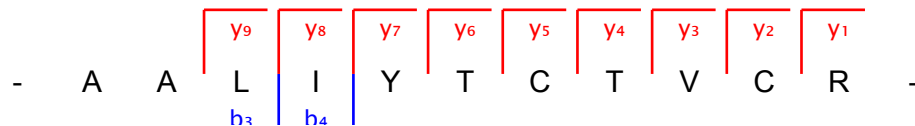
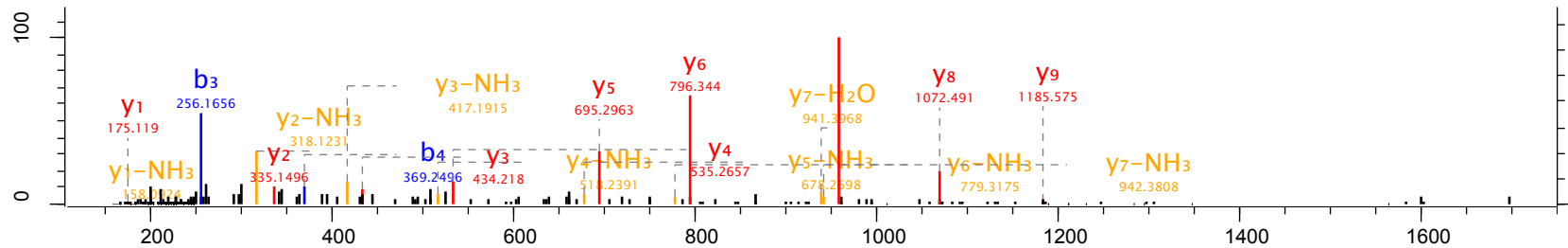
121.9

m/z

664.33

Gene names

ZNF706



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract21_dyn_5ul_G5_01_600

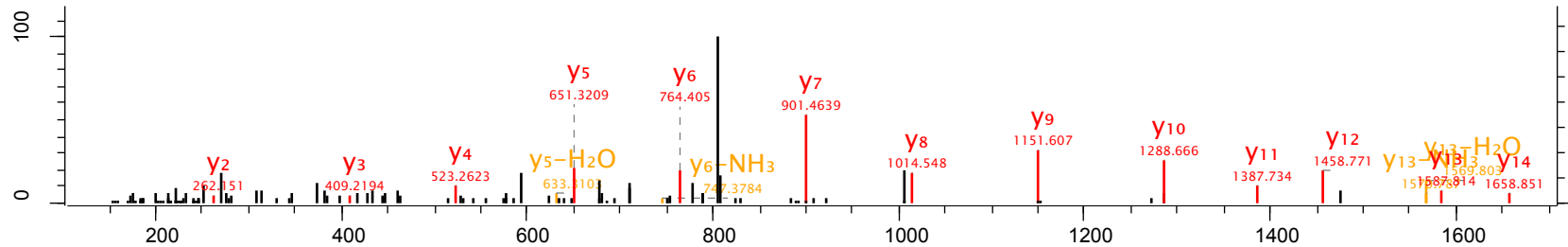
19806

TOF; CID

50.18

461.74

ZBTB45



ac

-

A

A

A

E

A

V

H

H

I

H

L

Q

N

F

S

R

-

y14

y13

y12

y11

y10

y9

y8

y7

y6

y5

y4

y3

y2

Raw file

20141014_fract21_dyn_5ul_G5_01_600

Scan

22193

Method

TOF; CID

Score

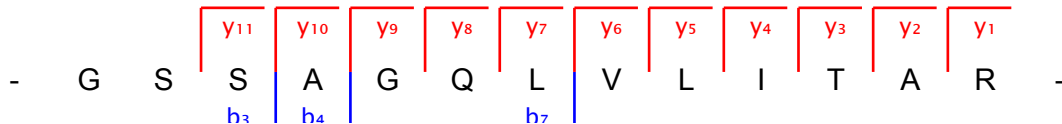
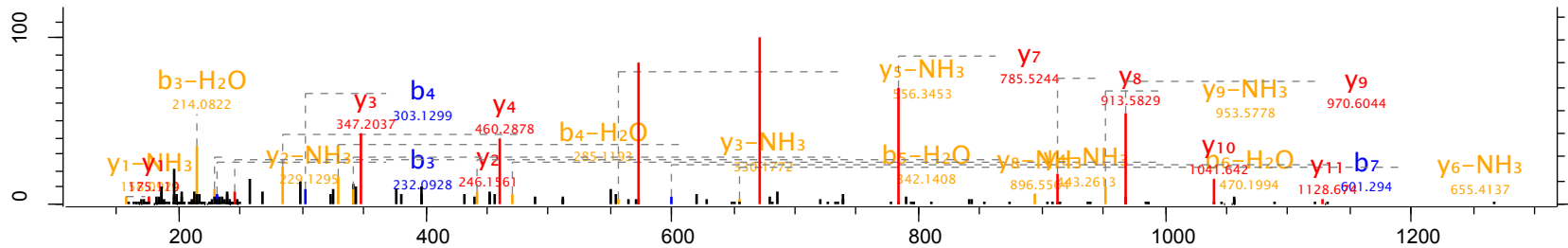
78.4

m/z

636.86

Gene names

TMEM219



Raw file

20141014_fract21_dyn_5ul_G5_01_600

Scan

22235

Method

TOF; CID

Score

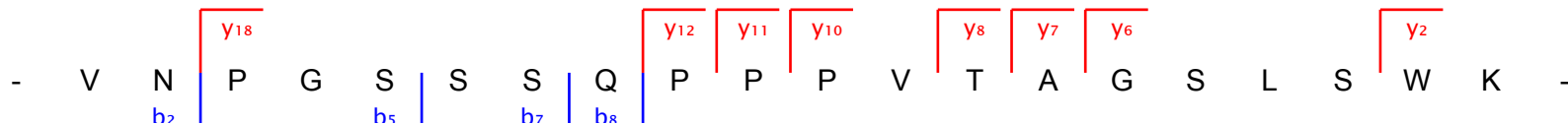
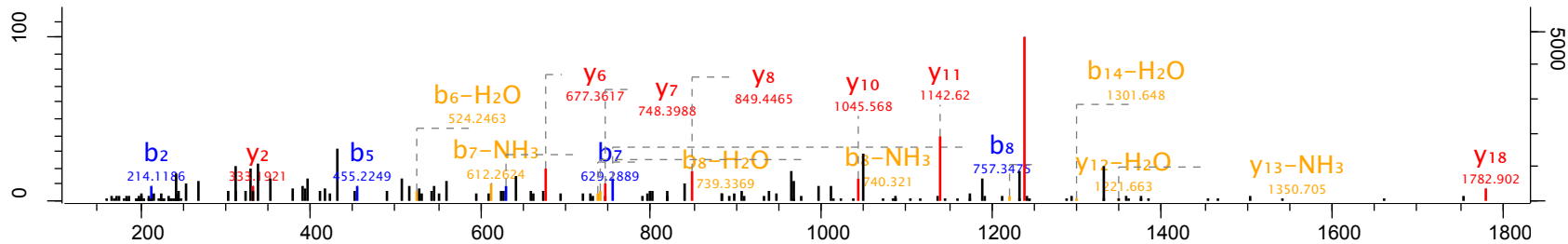
36.92

m/z

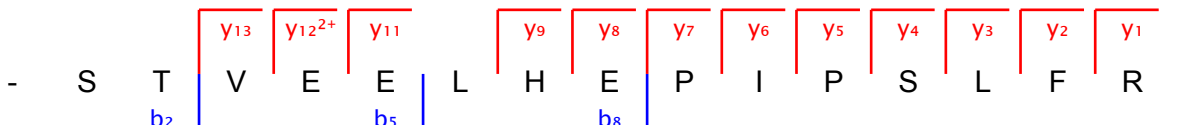
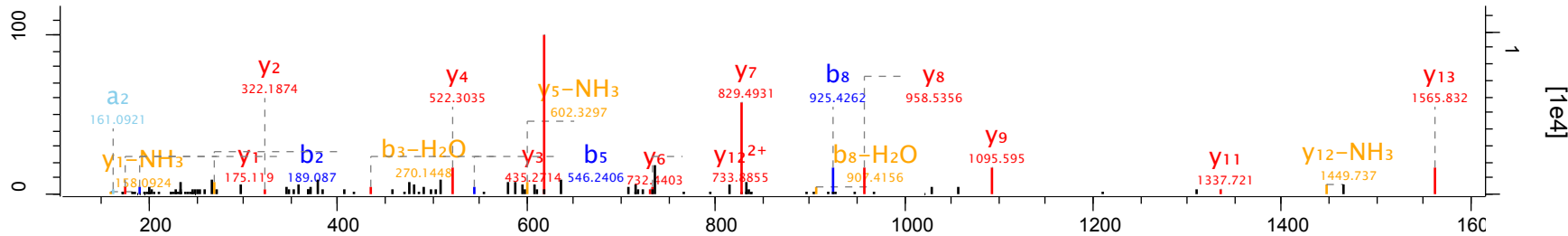
998.51

Gene names

LMO4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract21_dyn_5ul_G5_01_600	28402	TOF; CID	63.68	585.64	TWSG1



Raw file

20141014_fract22_dyn_5ul_G6_01_601

Scan

8661

Method

TOF; CID

Score

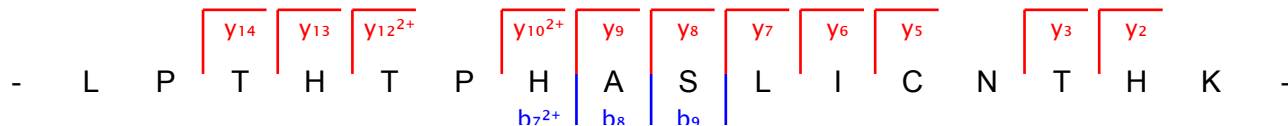
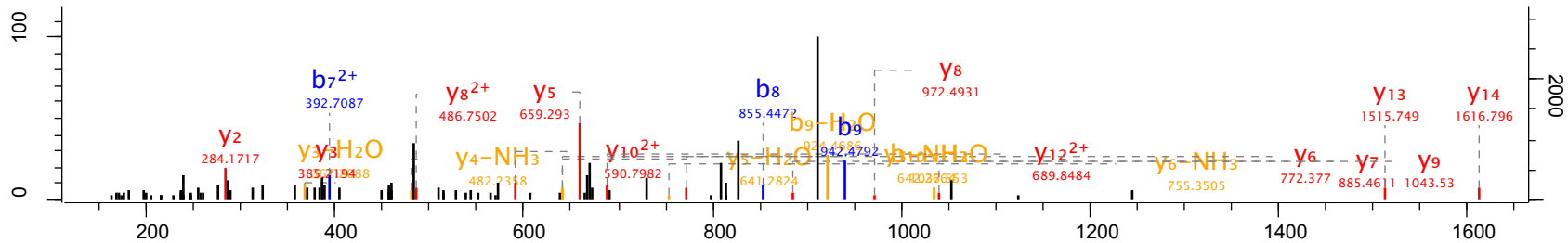
63.06

m/z

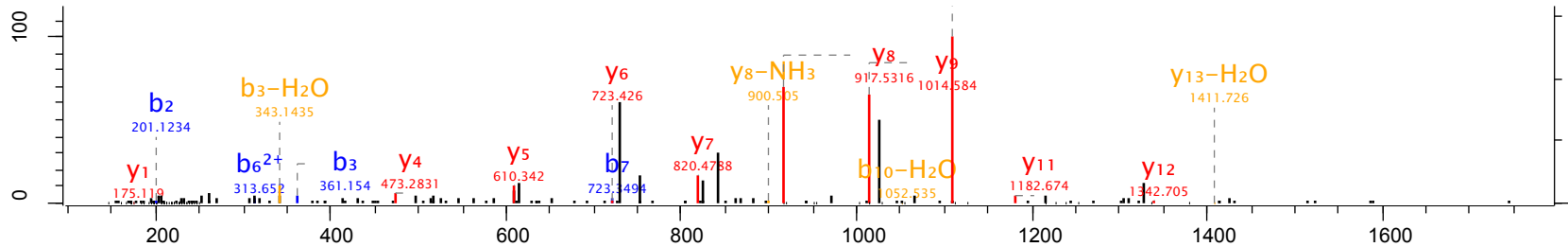
457.74

Gene names

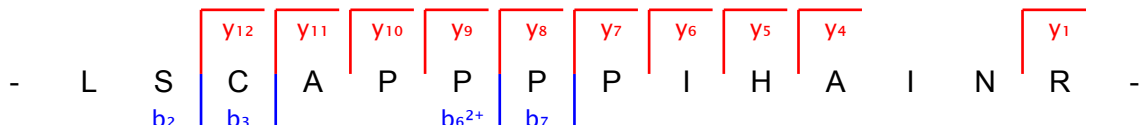
ZNF780A



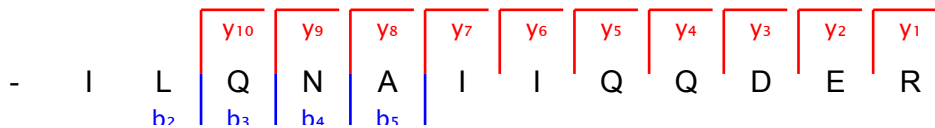
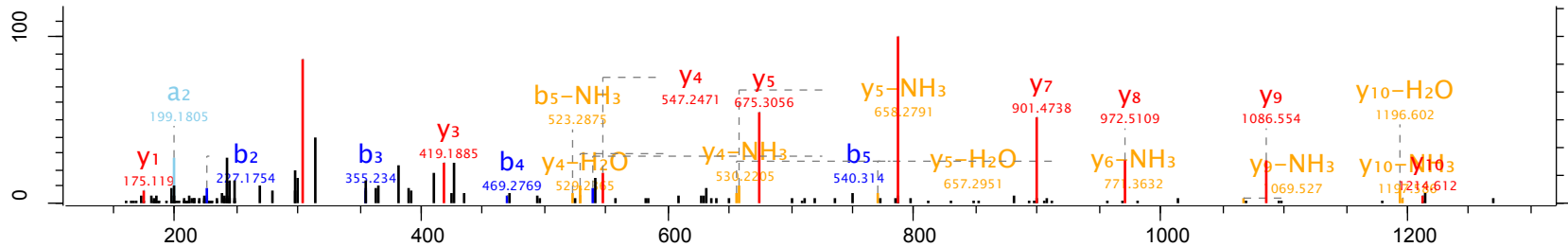
Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract22_dyn_5ul_G6_01_601	14883	TOF; CID	42.32	515.28	SLC23A2



[1e4]



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract22_dyn_5ul_G6_01_601	16056	TOF; CID	114.72	721.39	AKAP7



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract22_dyn_5ul_G6_01_601

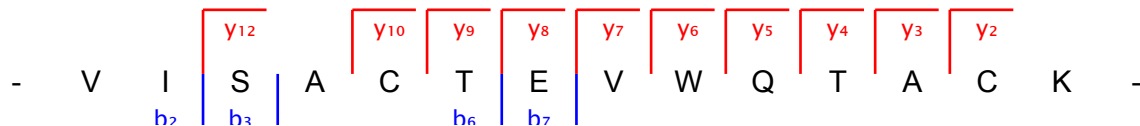
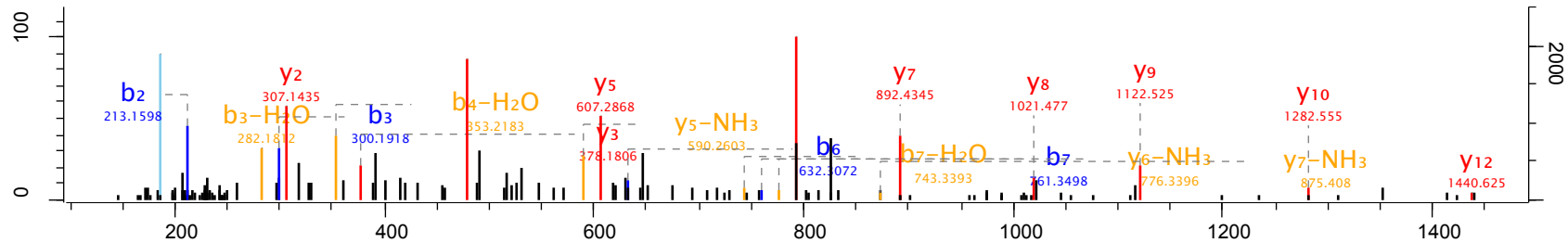
19737

TOF; CID

105.03

826.89

FBXW2



Raw file

20141014_fract22_dyn_5ul_G6_01_601

Scan

22553

Method

TOF; CID

Score

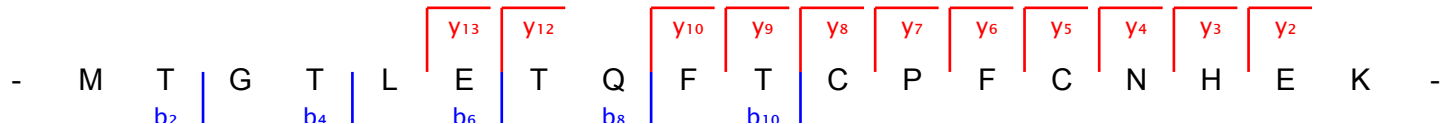
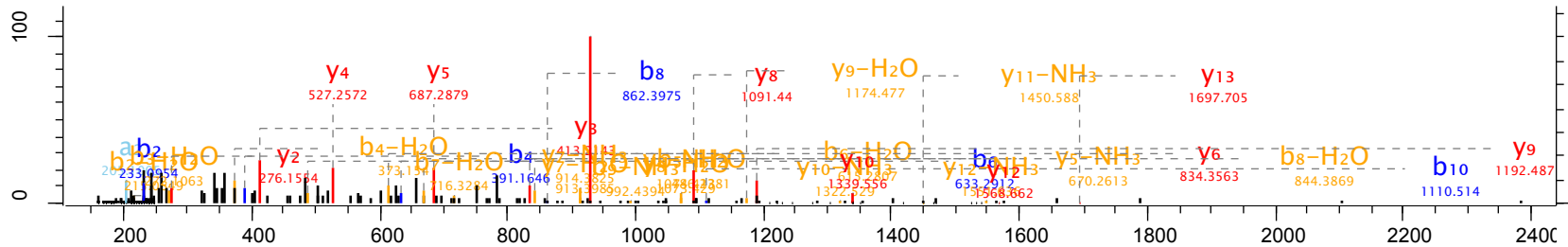
105.9

m/z

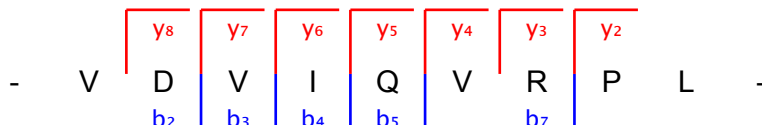
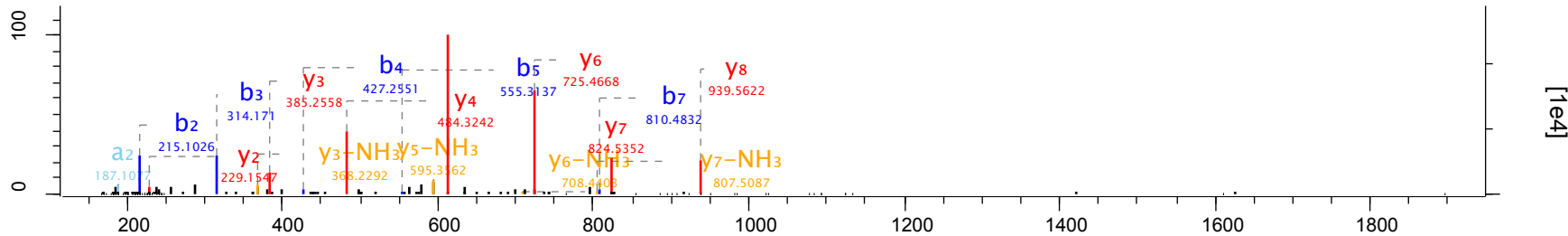
734.65

Gene names

ELOF1



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract22_dyn_5ul_G6_01_601	24438	TOF; CID	103.55	519.82	OPN3



Raw file

20141014_fract22_dyn_5ul_G6_01_601

Scan

32899

Method

TOF; CID

Score

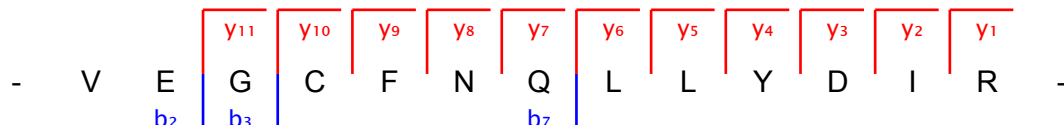
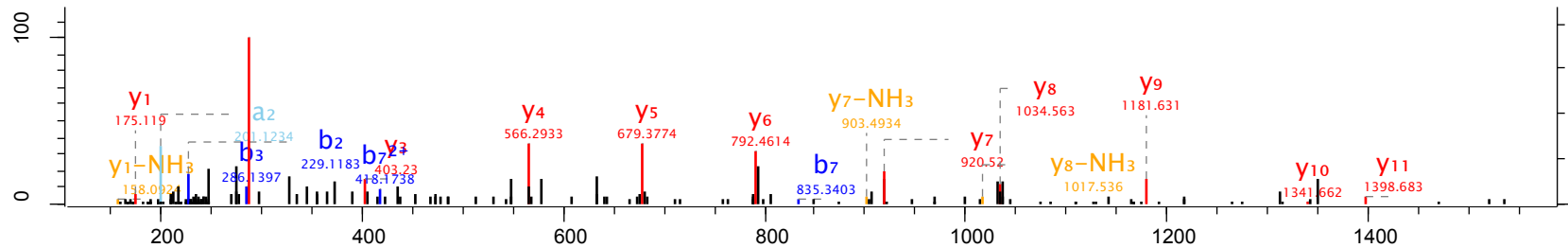
68.26

m/z

813.89

Gene names

TSPAN1



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract22_dyn_5ul_G6_01_601

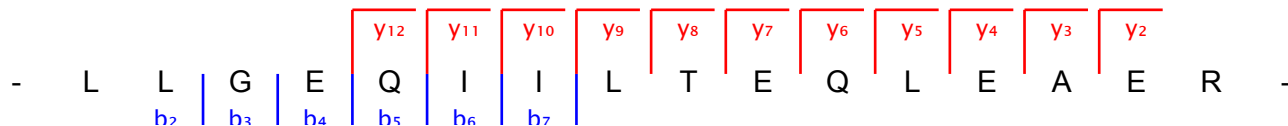
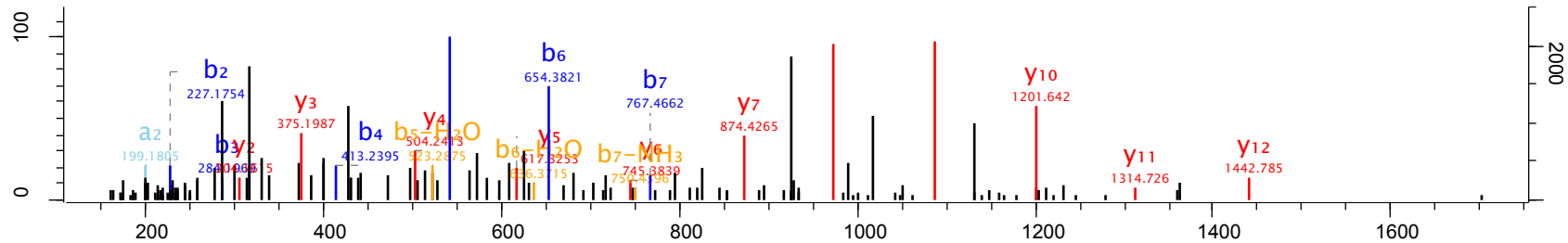
33398

TOF; CID

72.58

928.51

PET117



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

8390

Method

TOF; CID

Score

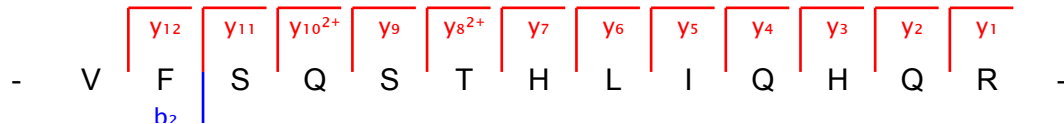
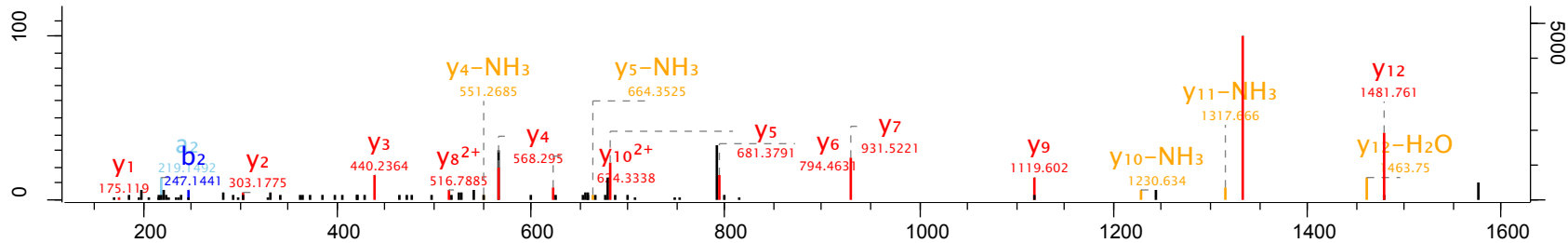
90.61

m/z

527.61

Gene names

ZNF34



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

10940

Method

TOF; CID

Score

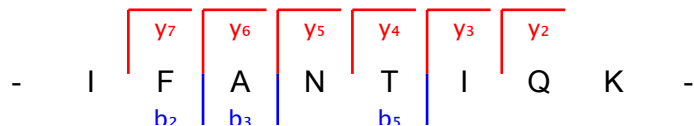
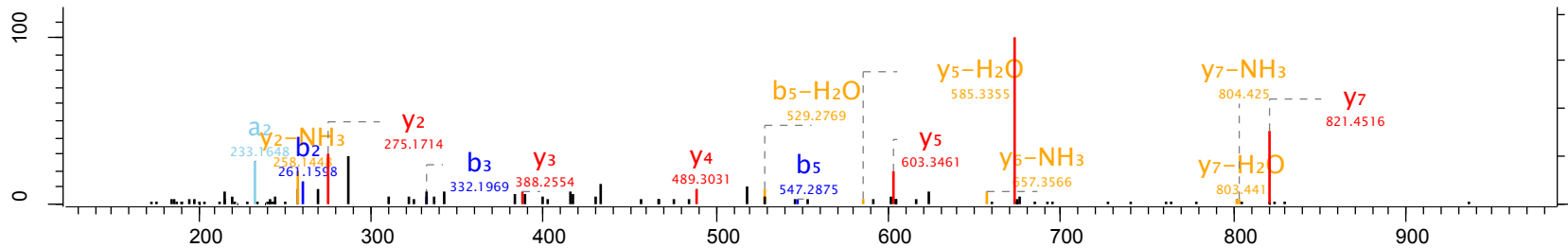
103.74

m/z

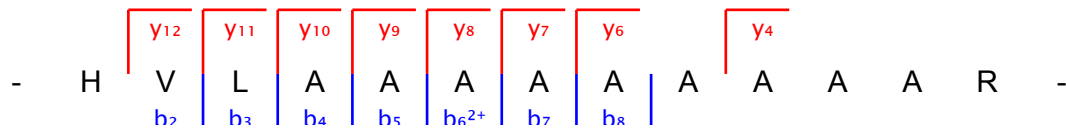
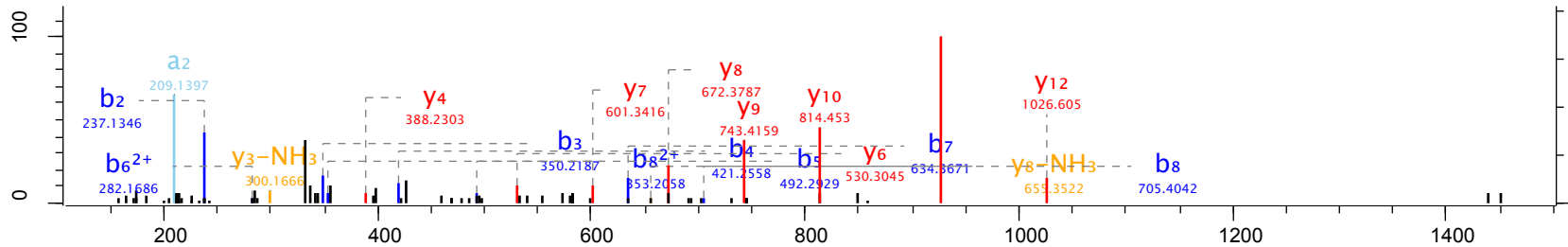
467.77

Gene names

ENTPD4



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract23_dyn_5ul_G7_01_602	11199	TOF; CID	87.18	582.33	MIDN



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

12421

Method

TOF; CID

Score

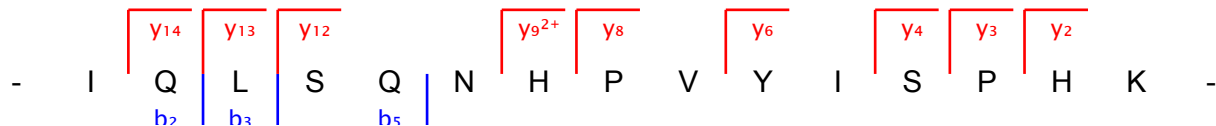
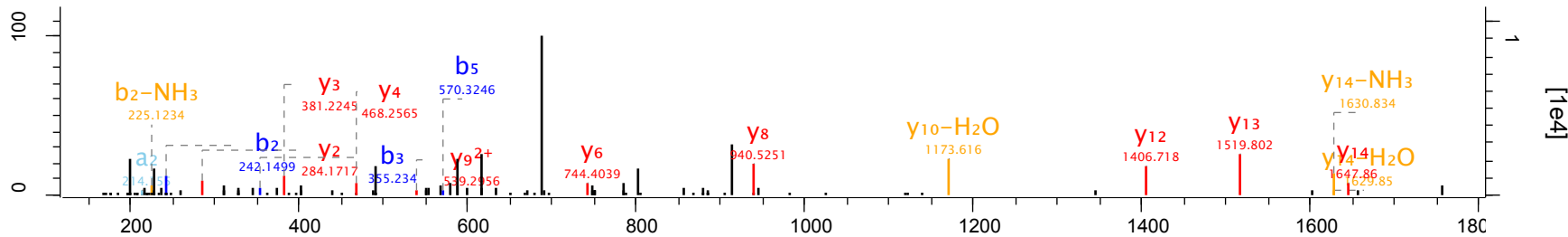
55.66

m/z

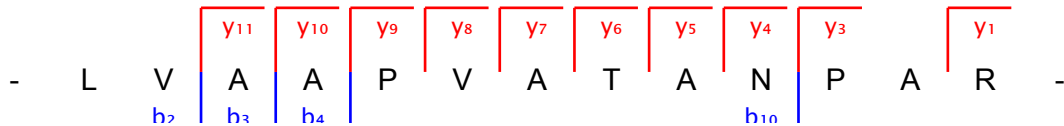
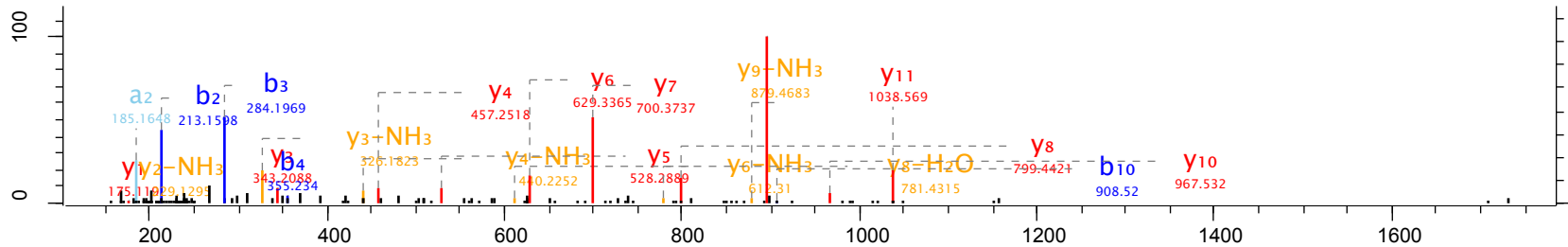
587.65

Gene names

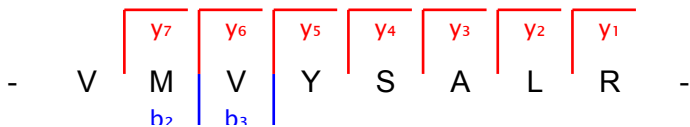
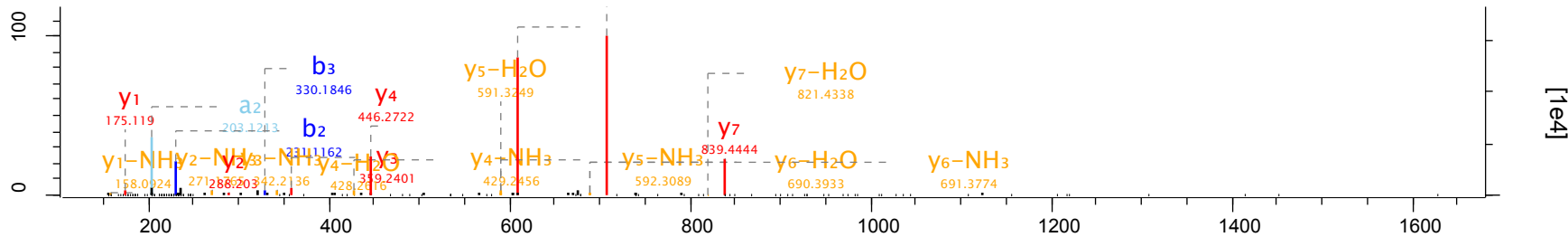
RBL2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract23_dyn_5ul_G7_01_602	13733	TOF; CID	94.12	625.86	FKRP



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract23_dyn_5ul_G7_01_602	16291	TOF; CID	117.1	469.76	APH1A



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract23_dyn_5ul_G7_01_602

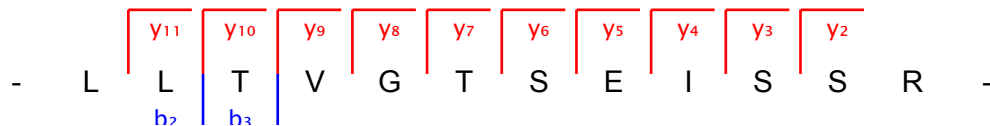
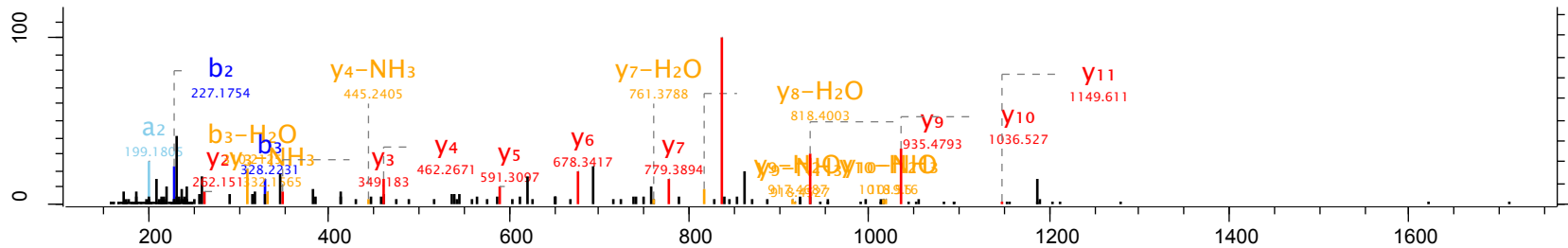
17175

TOF; CID

75.5

632.35

TTBK2



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

17490

Method

TOF; CID

Score

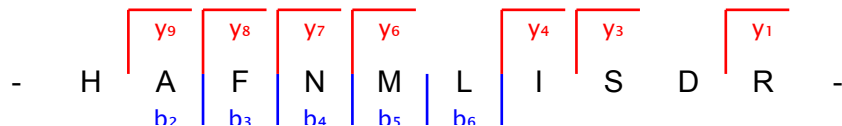
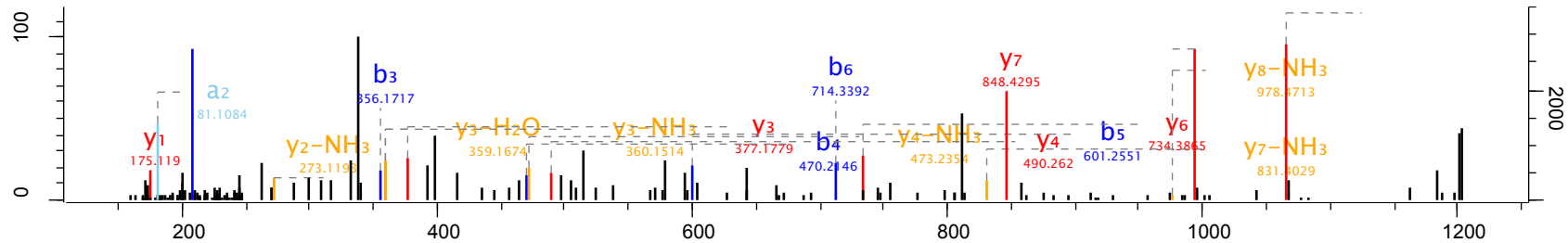
99.97

m/z

602.3

Gene names

GALNT11



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

22769

Method

TOF; CID

Score

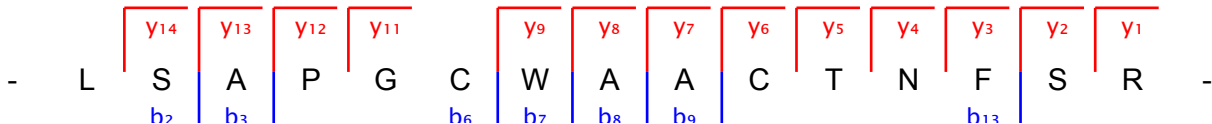
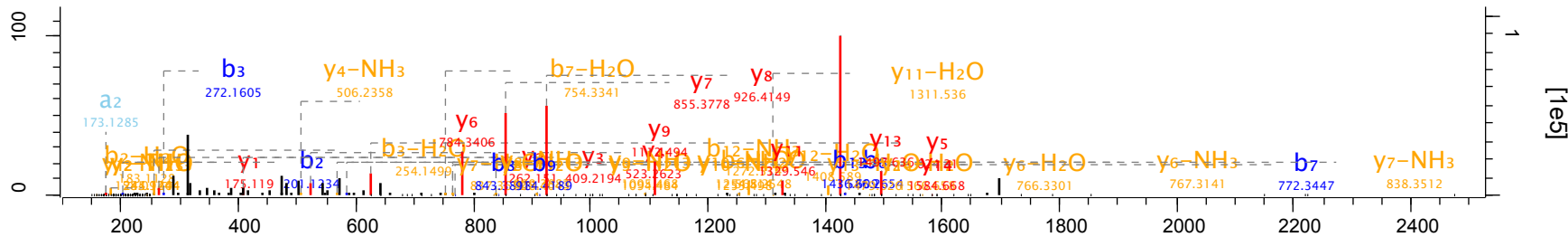
149.59

m/z

849.38

Gene names

PLP2



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

23436

Method

TOF; CID

Score

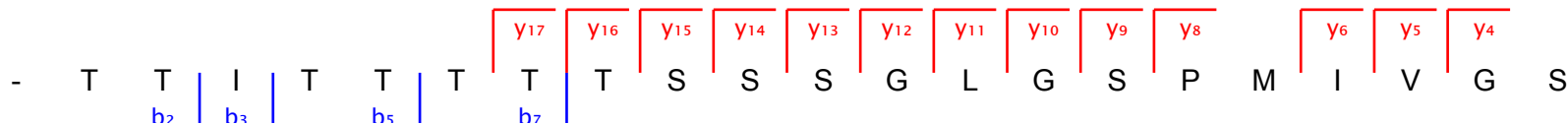
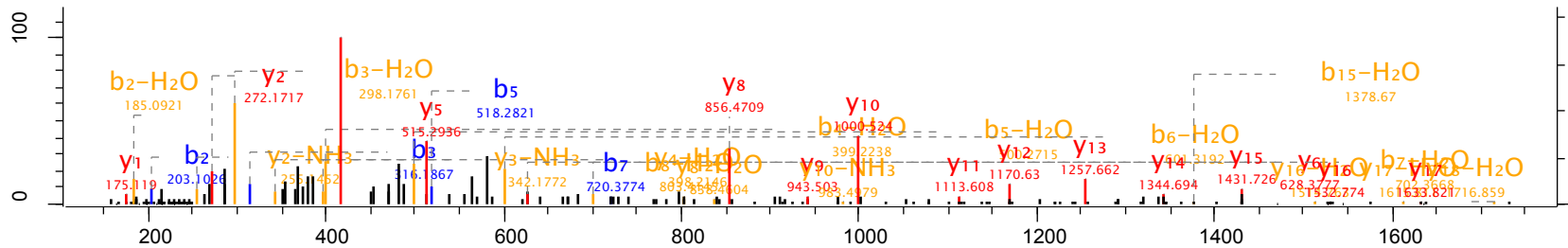
81.7

m/z

1127.07

Gene names

MYADM



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract23_dyn_5ul_G7_01_602

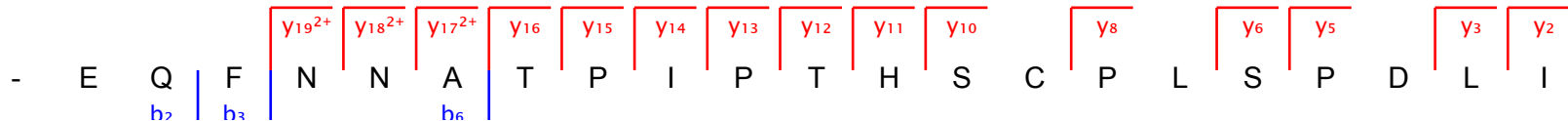
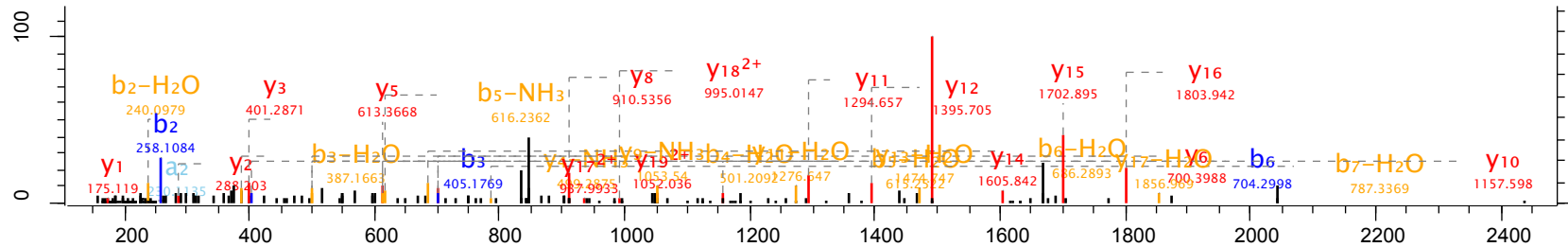
25922

TOF; CID

74.44

836.41

TMEM39A



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

30065

Method

TOF; CID

Score

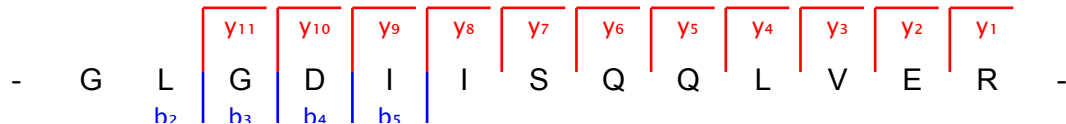
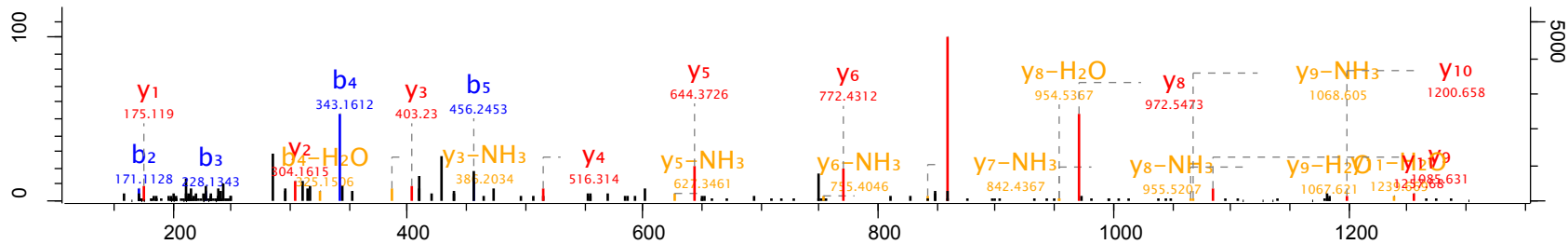
93.65

m/z

714.39

Gene names

MPV17



Raw file

20141014_fract23_dyn_5ul_G7_01_602

Scan

30090

Method

TOF; CID

Score

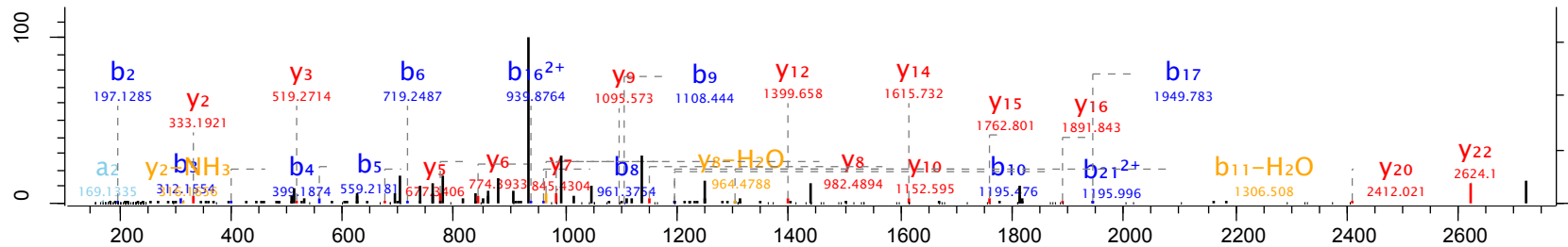
56.13

m/z

908.73

Gene names

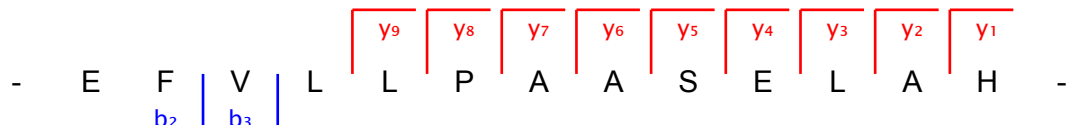
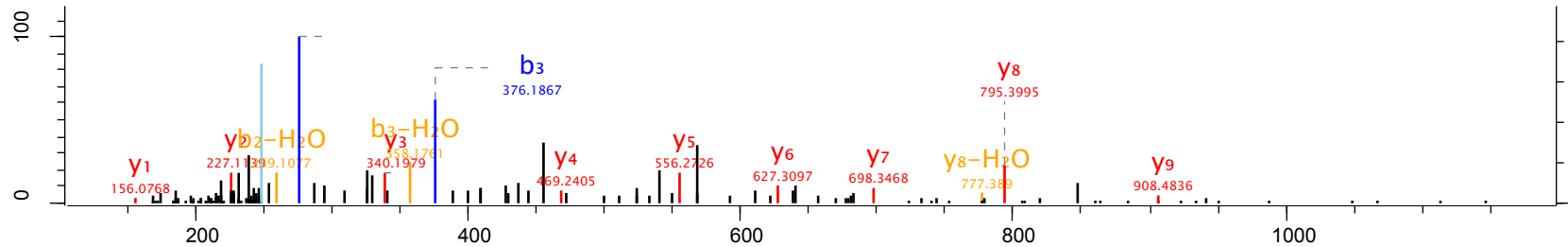
TSPAN4



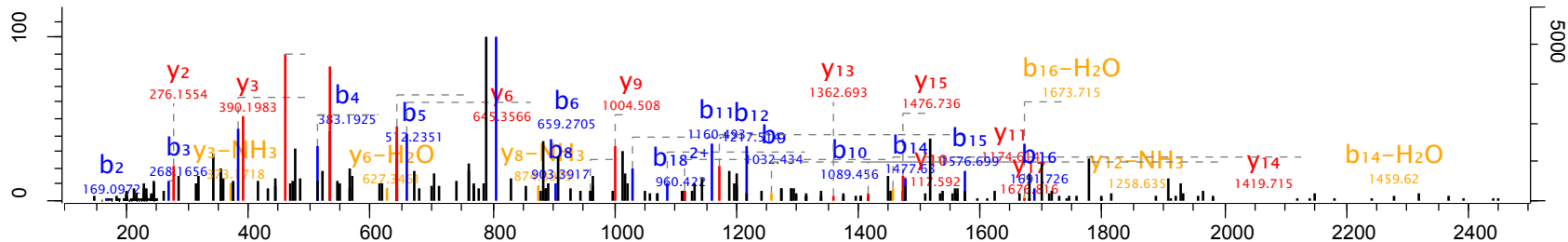
[1e4]



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract23_dyn_5ul_G7_01_602	35313	TOF; CID	65.84	698.87	PRKD2



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract23_dyn_5ul_G7_01_602	37784	TOF; CID	51.71	1199.88	MYEOV2



Raw file

20141014_fract24_dyn_5ul_G8_01_603

Scan

12552

Method

TOF; CID

Score

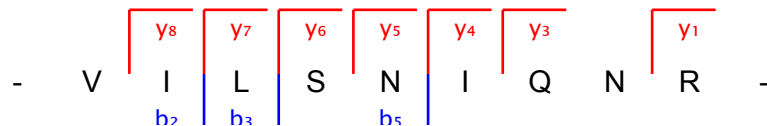
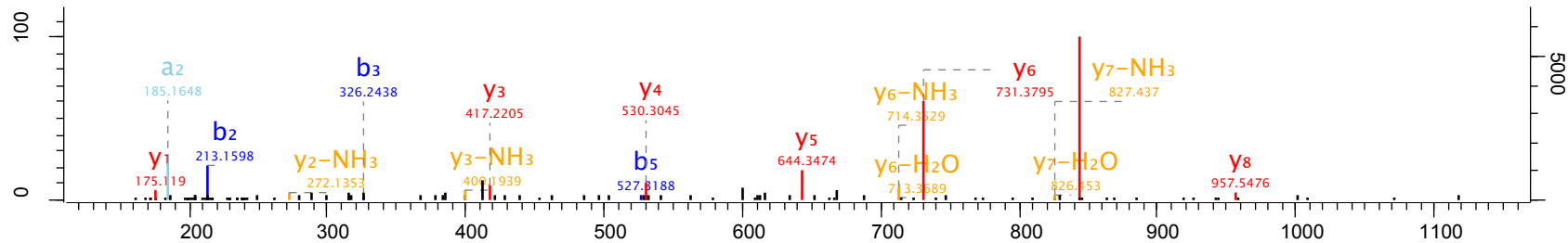
95.5

m/z

528.81

Gene names

TTC7B



Raw file

20141014_fract24_dyn_5ul_G8_01_603

Scan

16867

Method

TOF; CID

Score

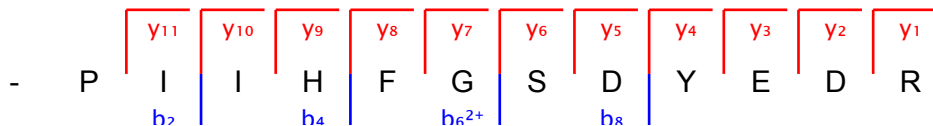
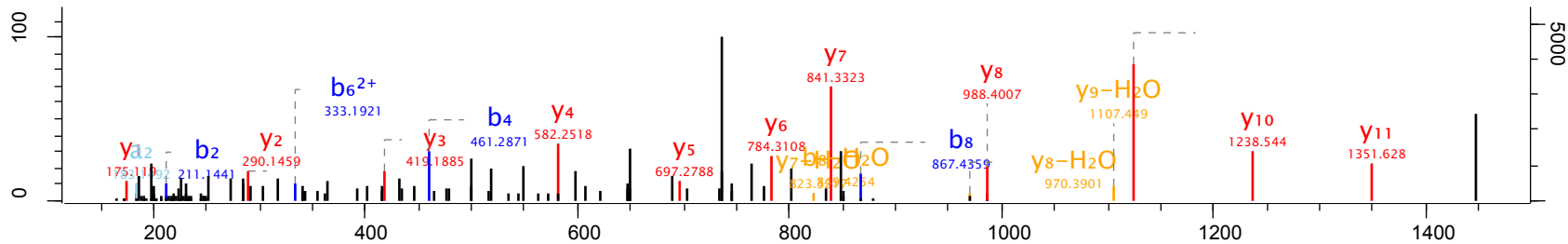
69.35

m/z

483.56

Gene names

PRNP



Raw file

20141014_fract24_dyn_5ul_G8_01_603

Scan

18665

Method

TOF; CID

Score

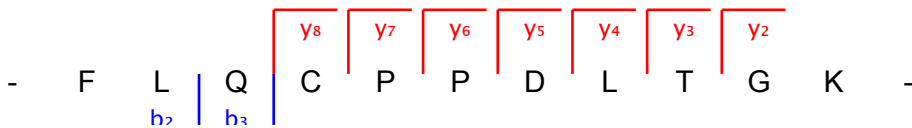
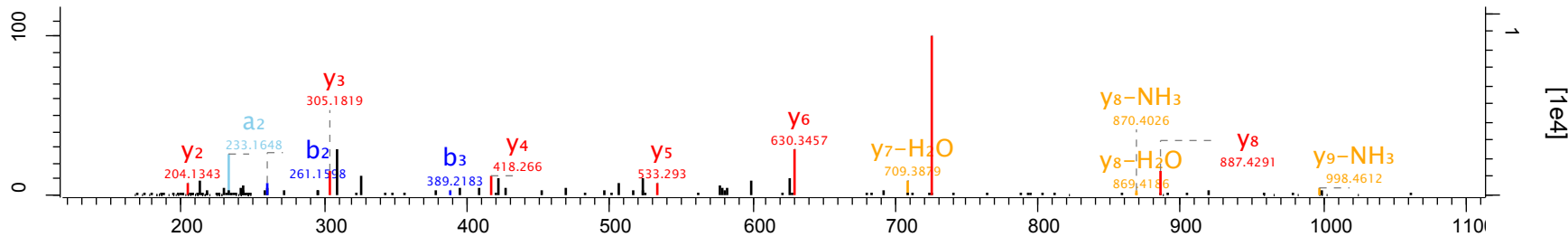
69.72

m/z

638.32

Gene names

PIGB



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract24_dyn_5ul_G8_01_603

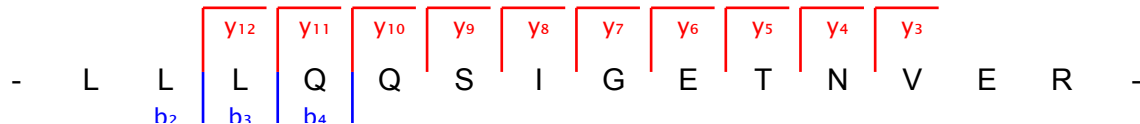
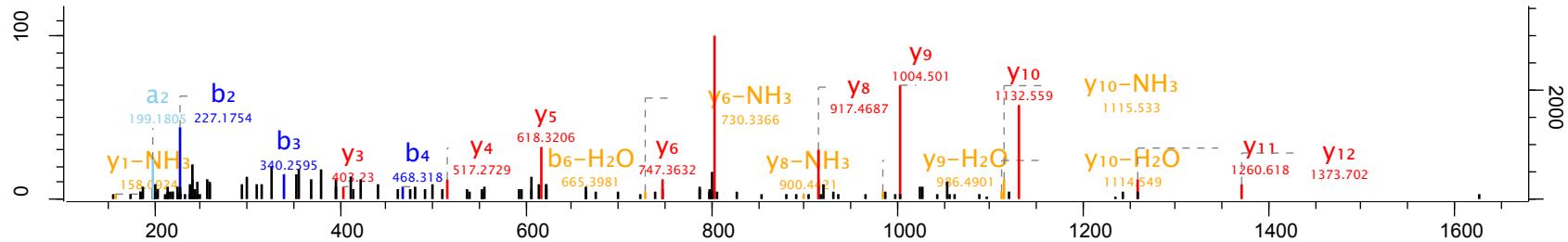
20741

TOF; CID

68.04

800.44

ZCCHC10



Raw file

20141014_fract24_dyn_5ul_G8_01_603

Scan

21101

Method

TOF; CID

Score

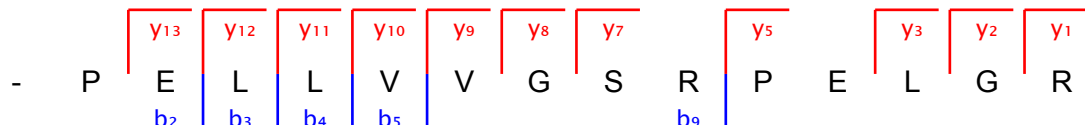
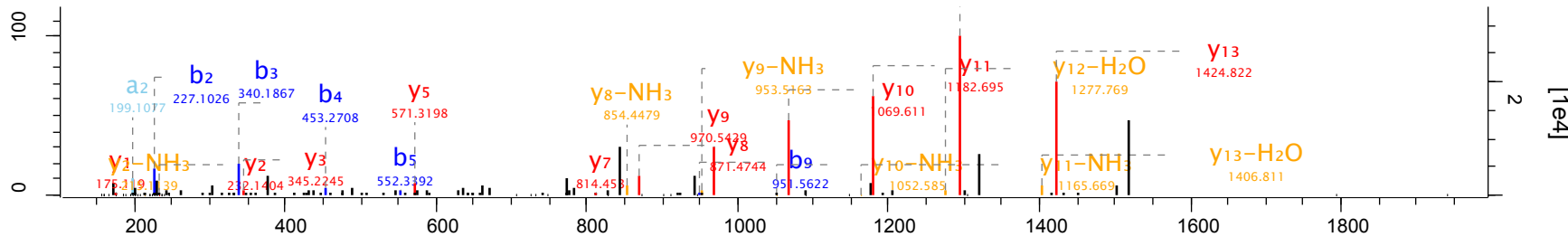
81.63

m/z

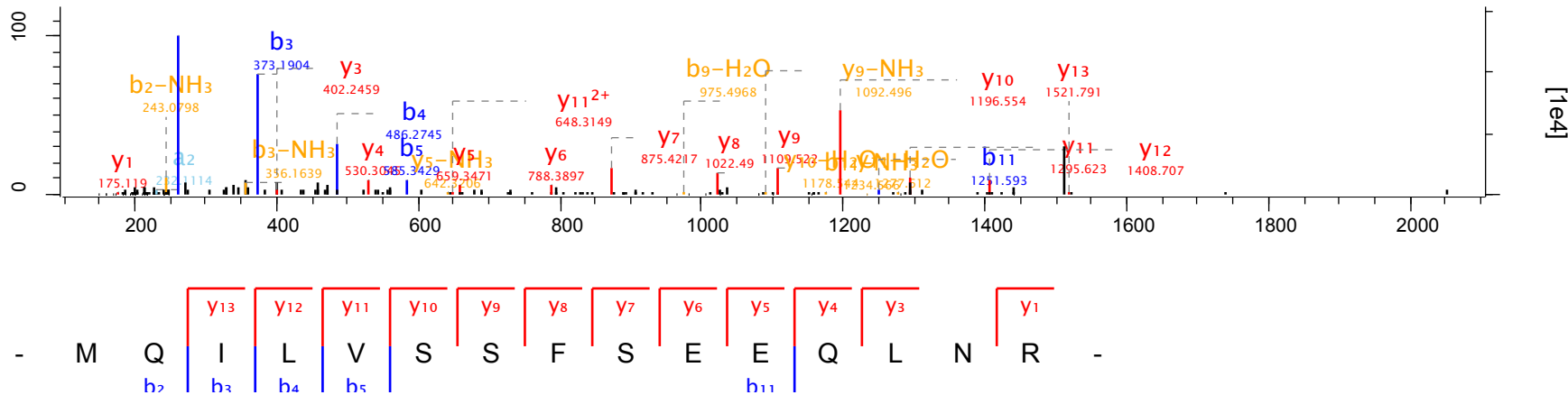
507.96

Gene names

EPM2A



Raw file	Scan	Method	Score	m/z	Gene names
20141014_fract24_dyn_5ul_G8_01_603	27723	TOF; CID	84.51	891.45	TAF11



Raw file

Scan

Method

Score

m/z

Gene names

20141014_fract24_dyn_5ul_G8_01_603

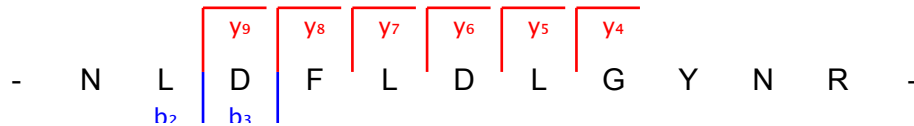
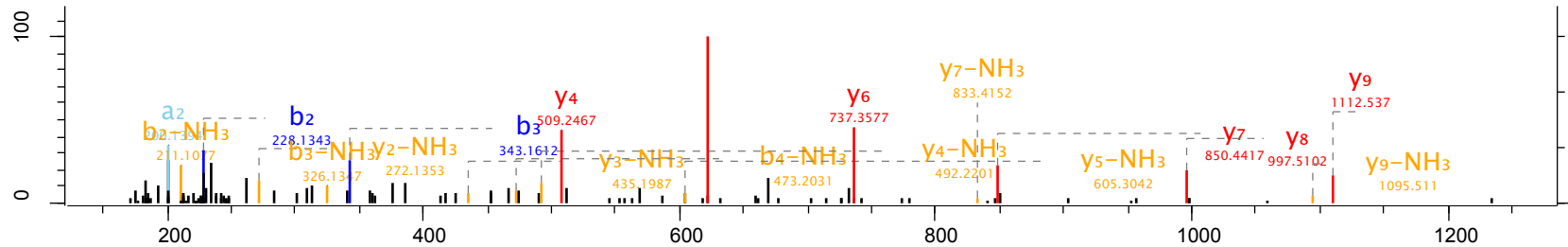
32521

TOF; CID

91.55

670.33

LRRTM4



Raw file

20141014_fract24_dyn_5ul_G8_01_603

Scan

32753

Method

TOF; CID

Score

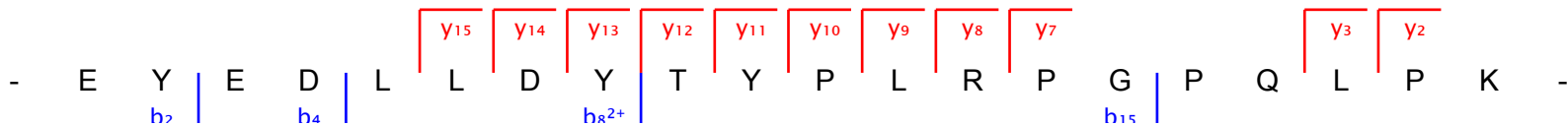
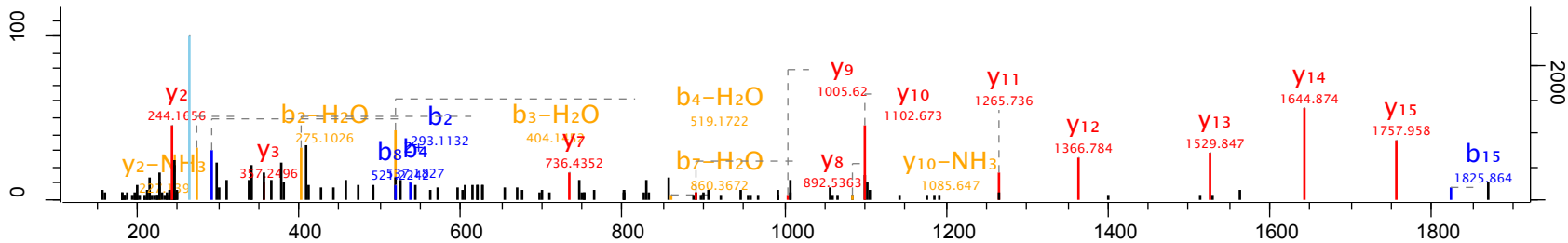
58.69

m/z

803.41

Gene names

CEP68



Raw file

20141014_fract24_dyn_5ul_G8_01_603

Scan

33196

Method

TOF; CID

Score

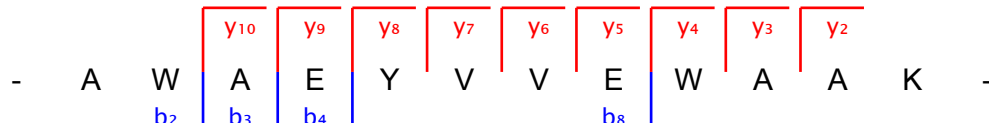
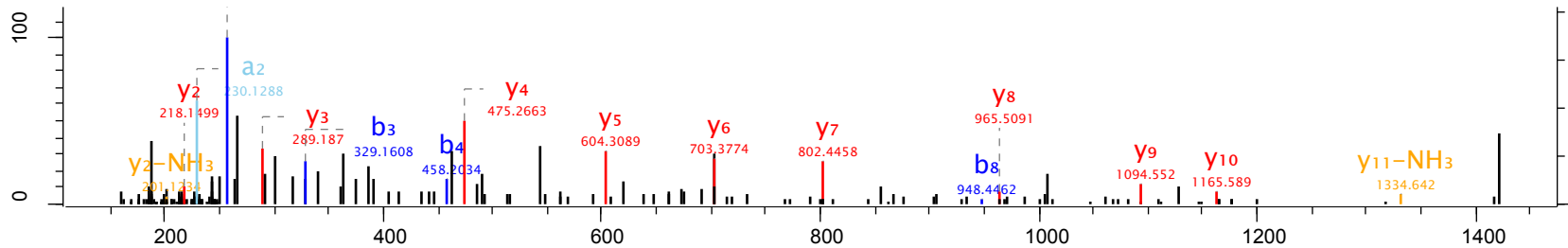
52.07

m/z

712.36

Gene names

SMIM15



Raw file

20141014_fract24_dyn_5ul_G8_01_603

Scan

34885

Method

TOF; CID

Score

103.83

m/z

814.95

Gene names

RCE1

