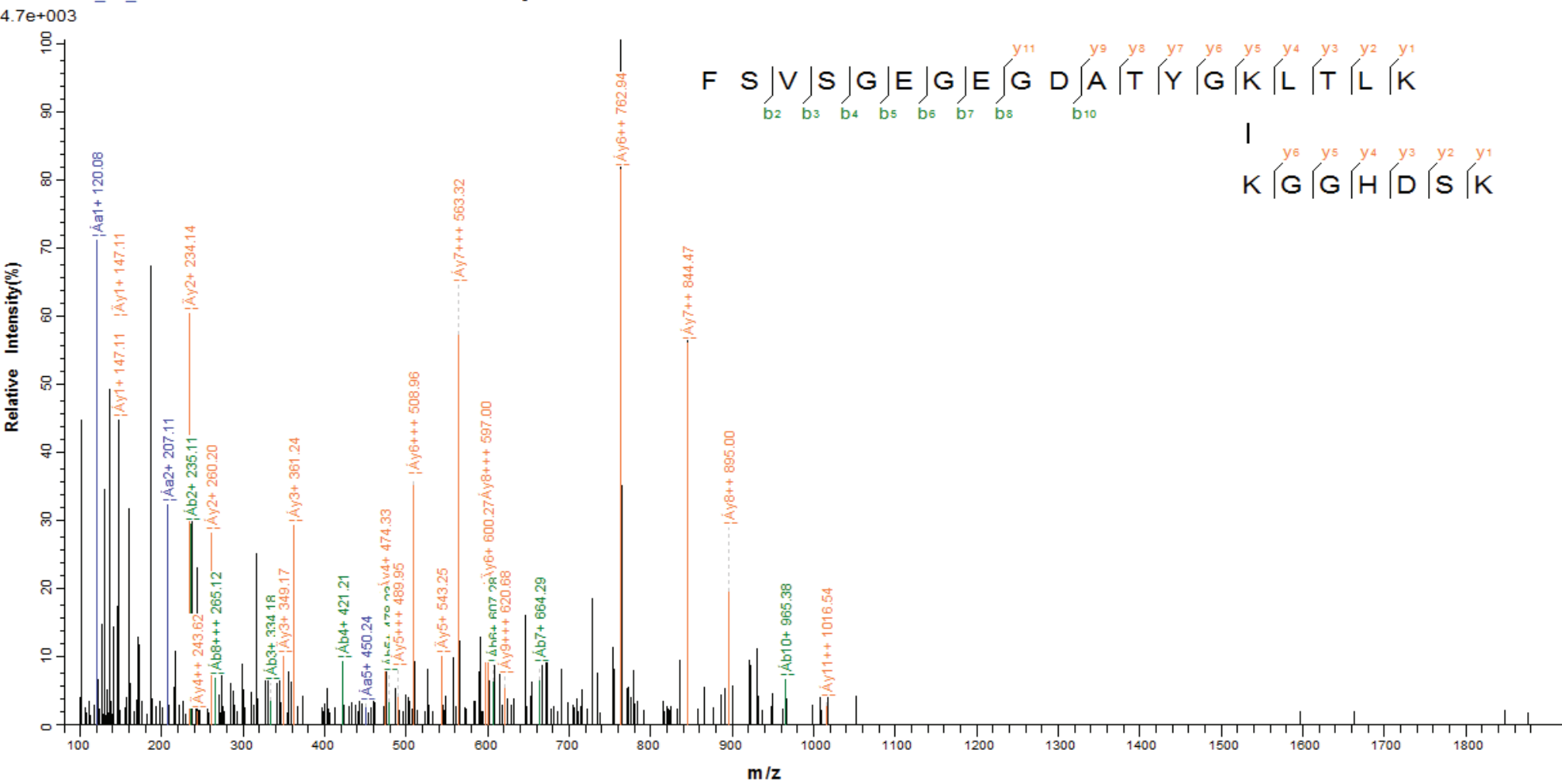
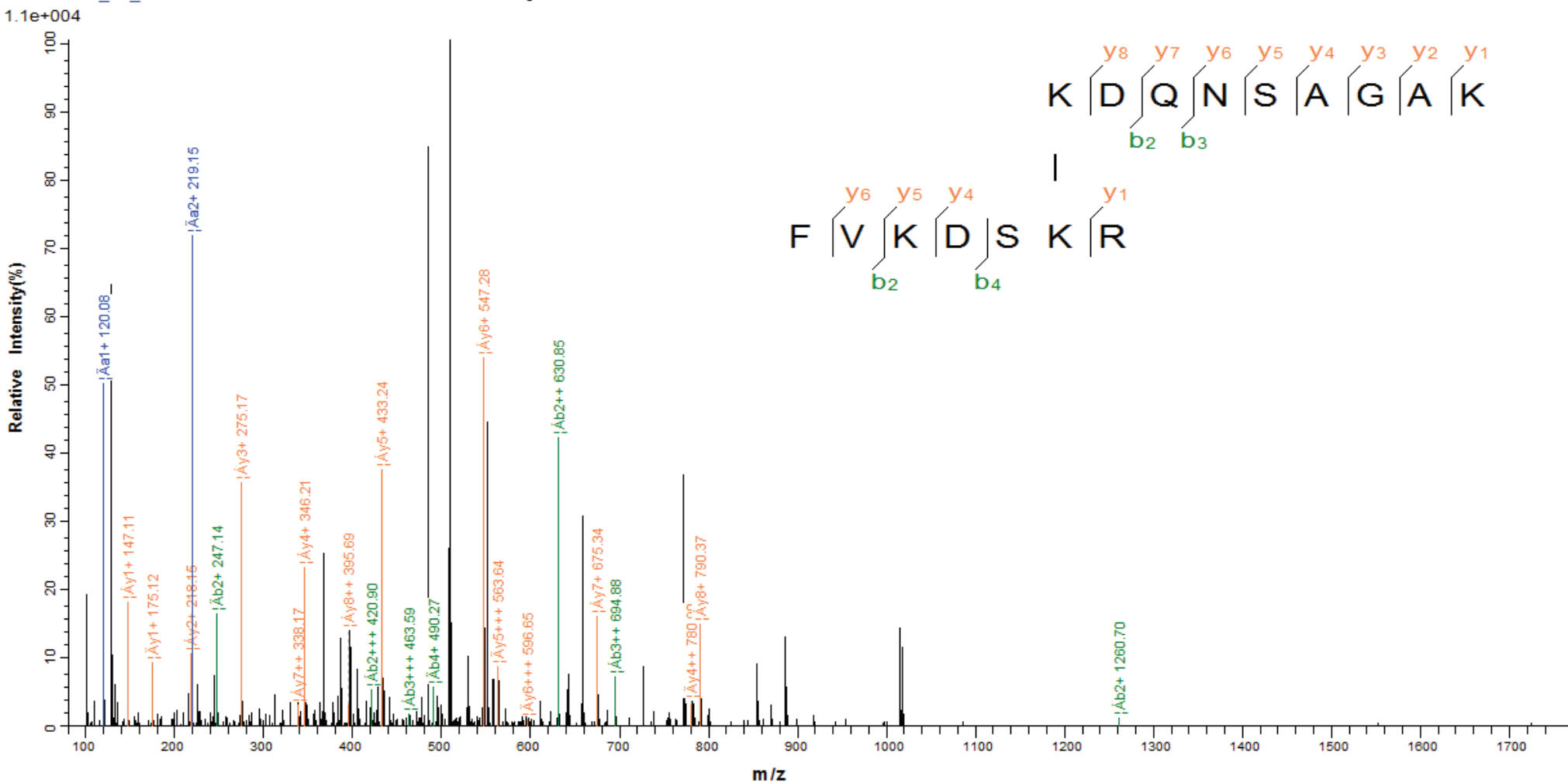
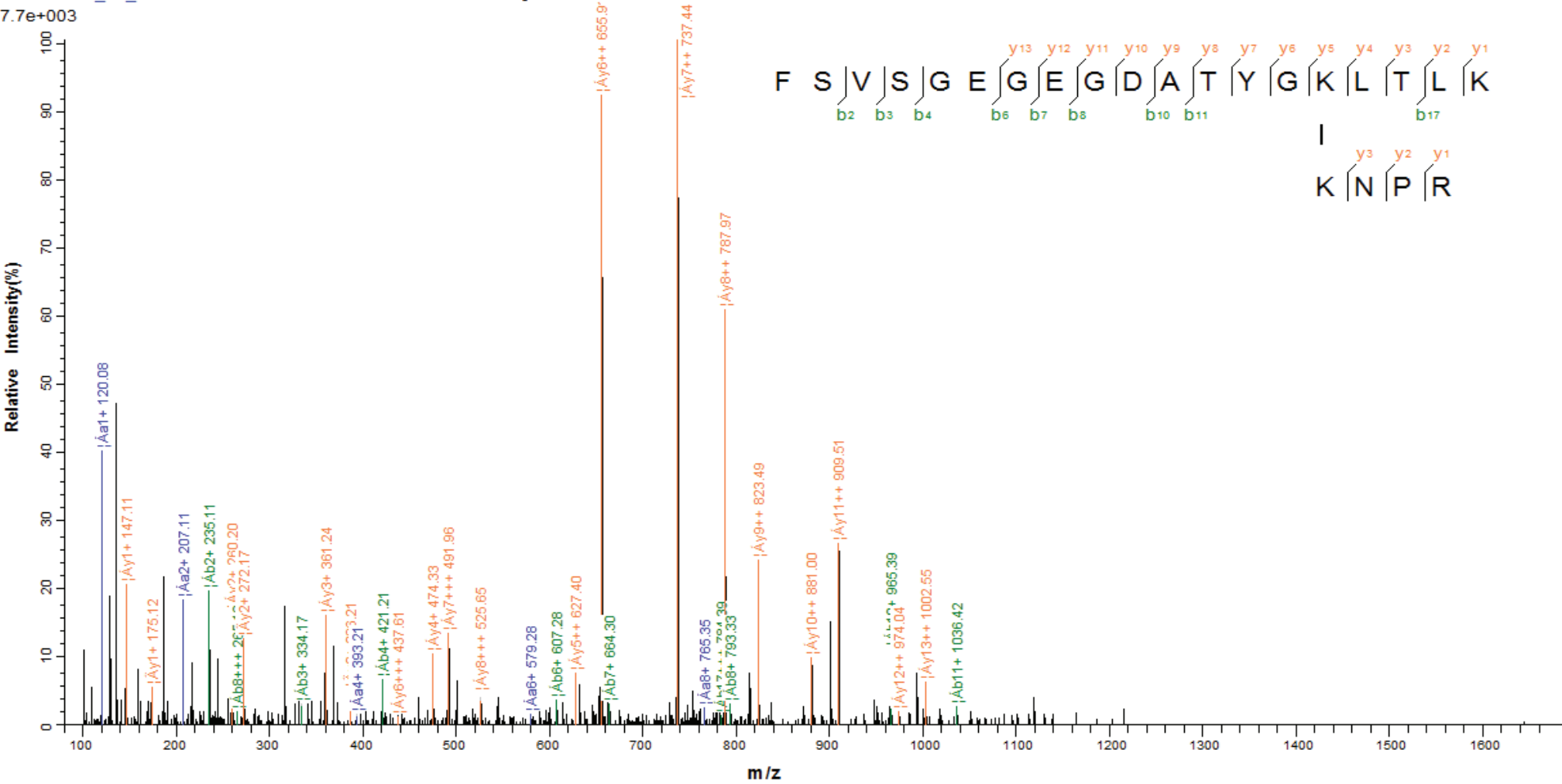


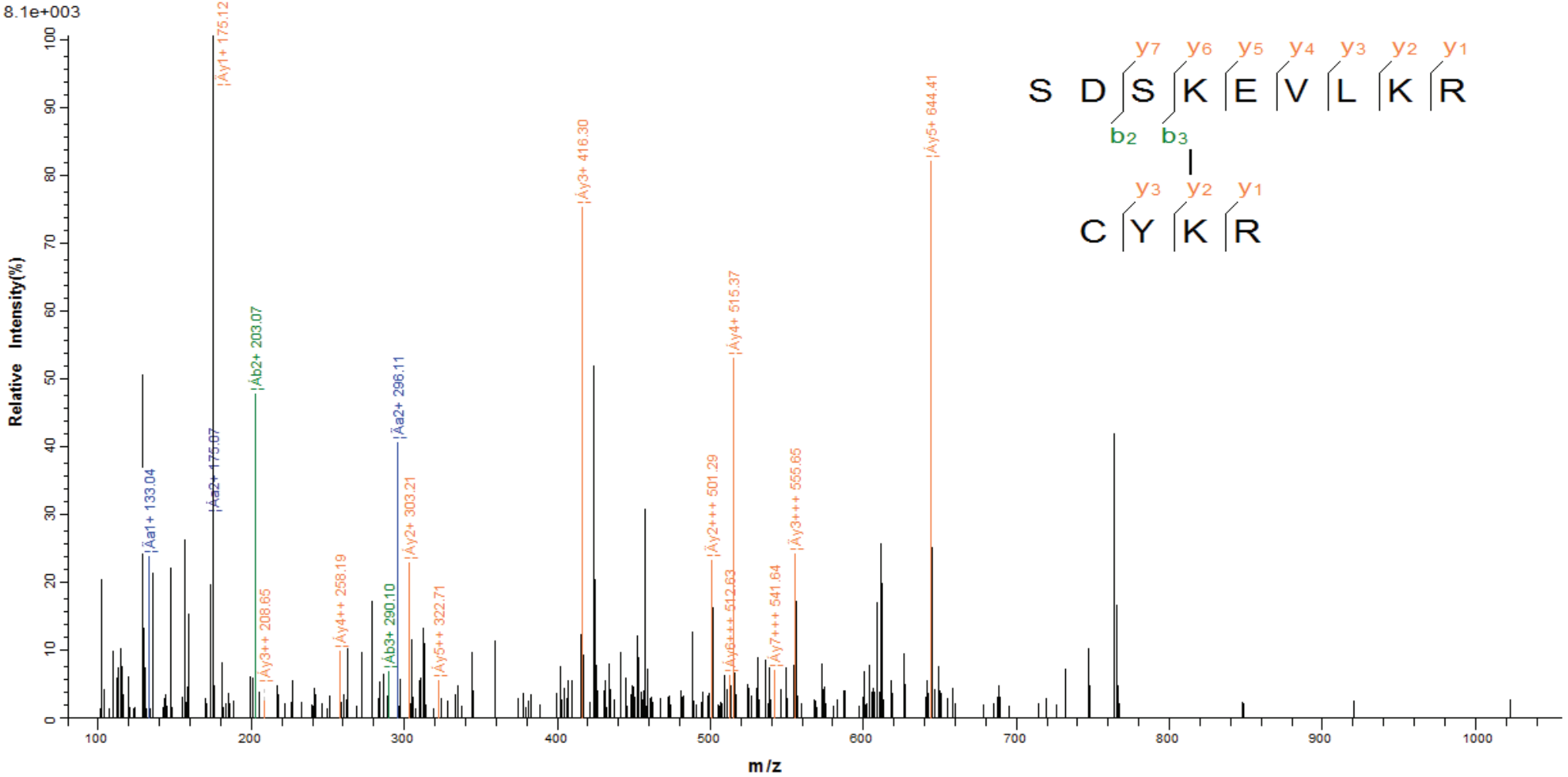
Fig. S16

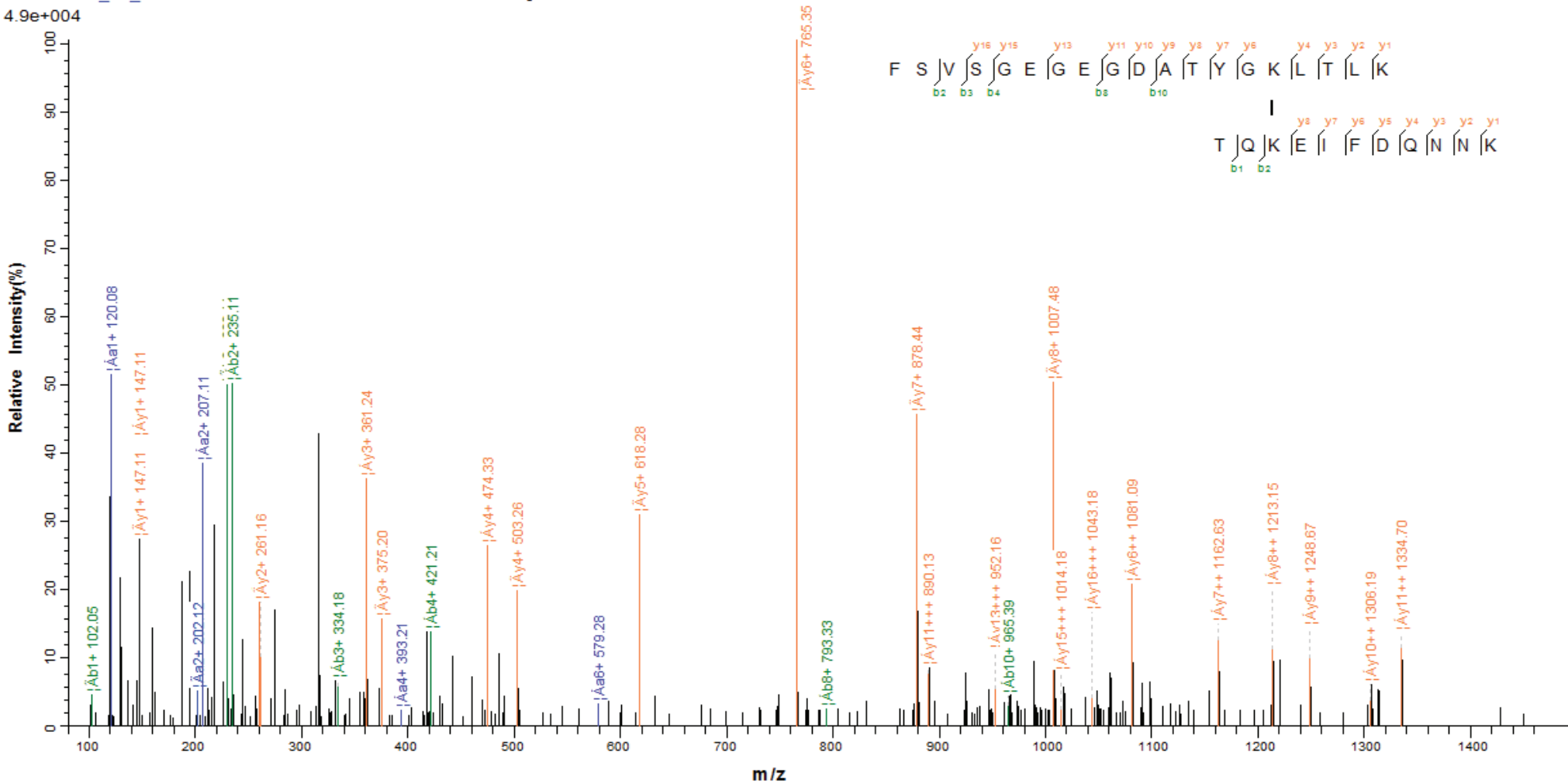
Annotated high-resolution HCD spectra
of crosslink peptides of APC/C complex

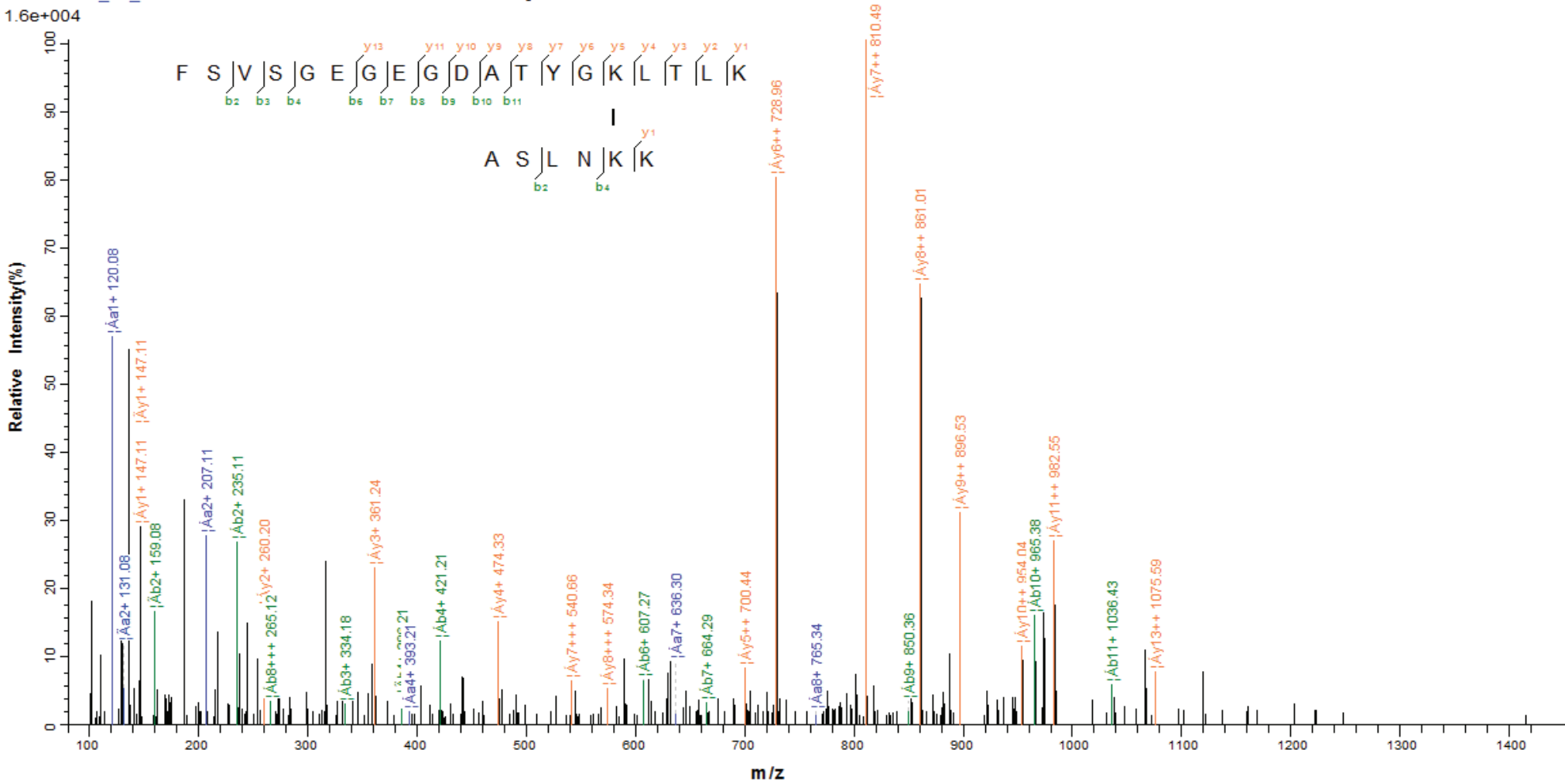






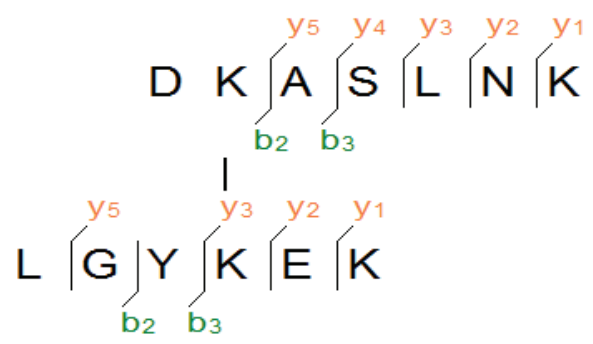
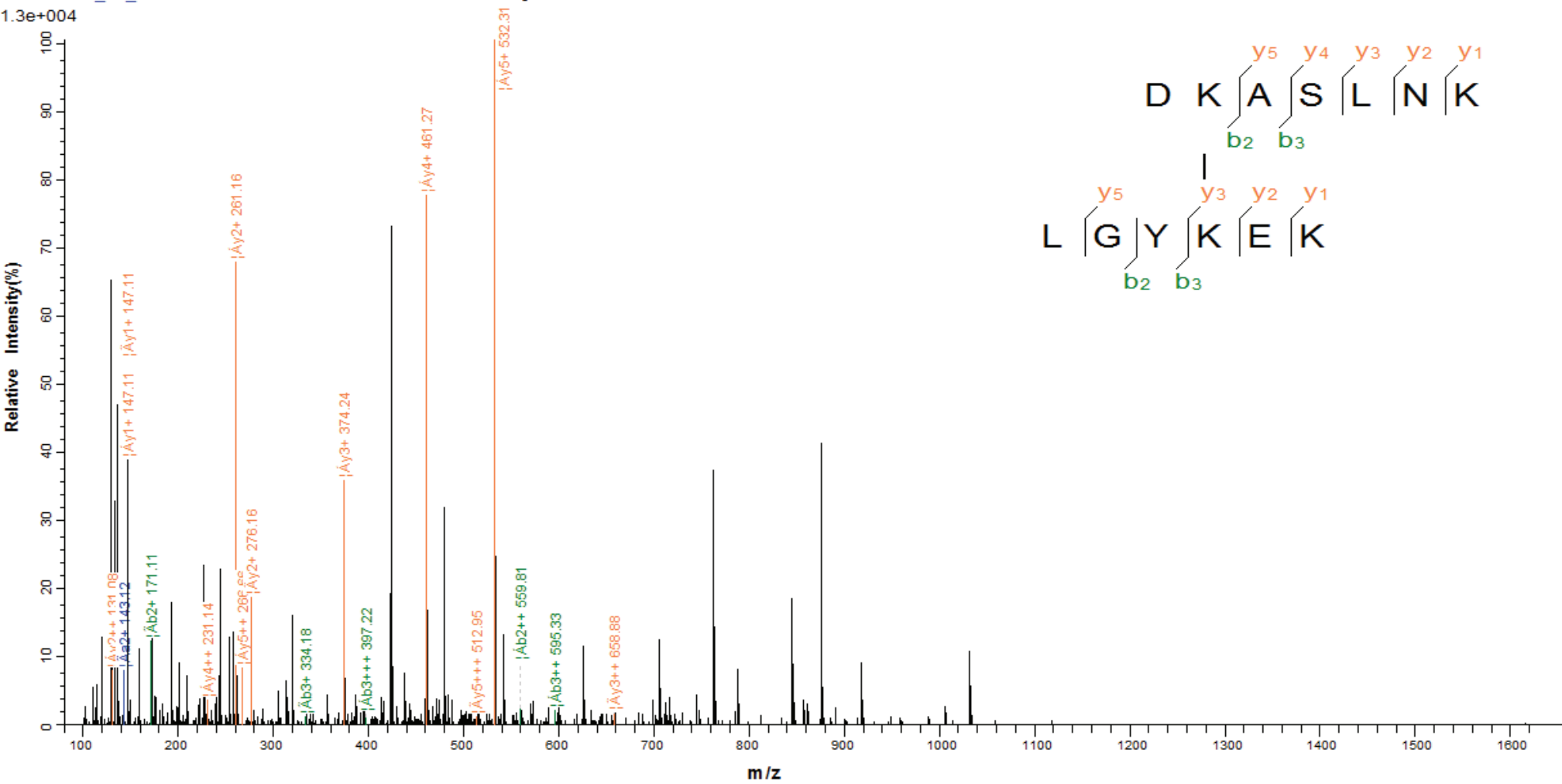




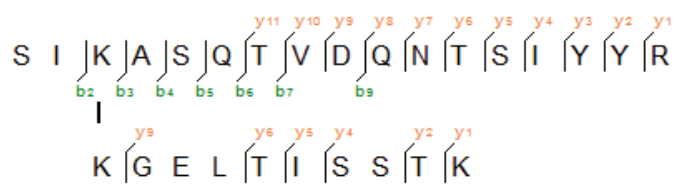
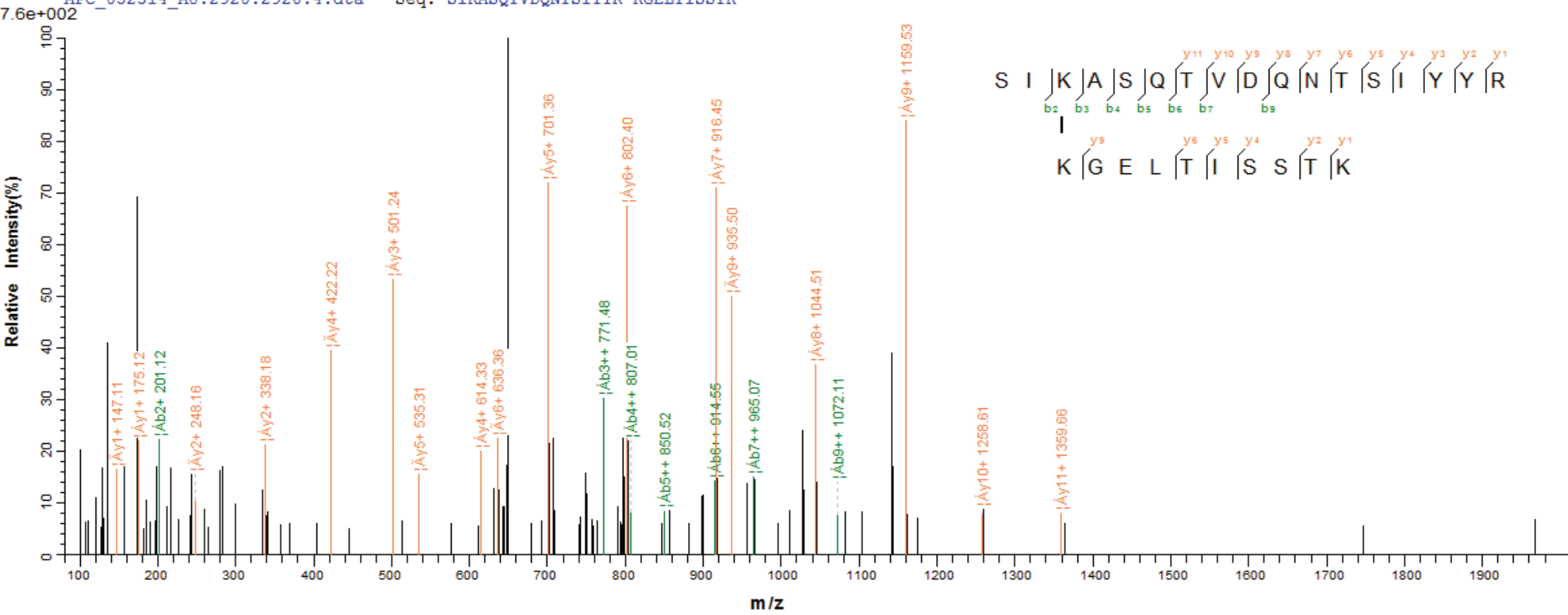


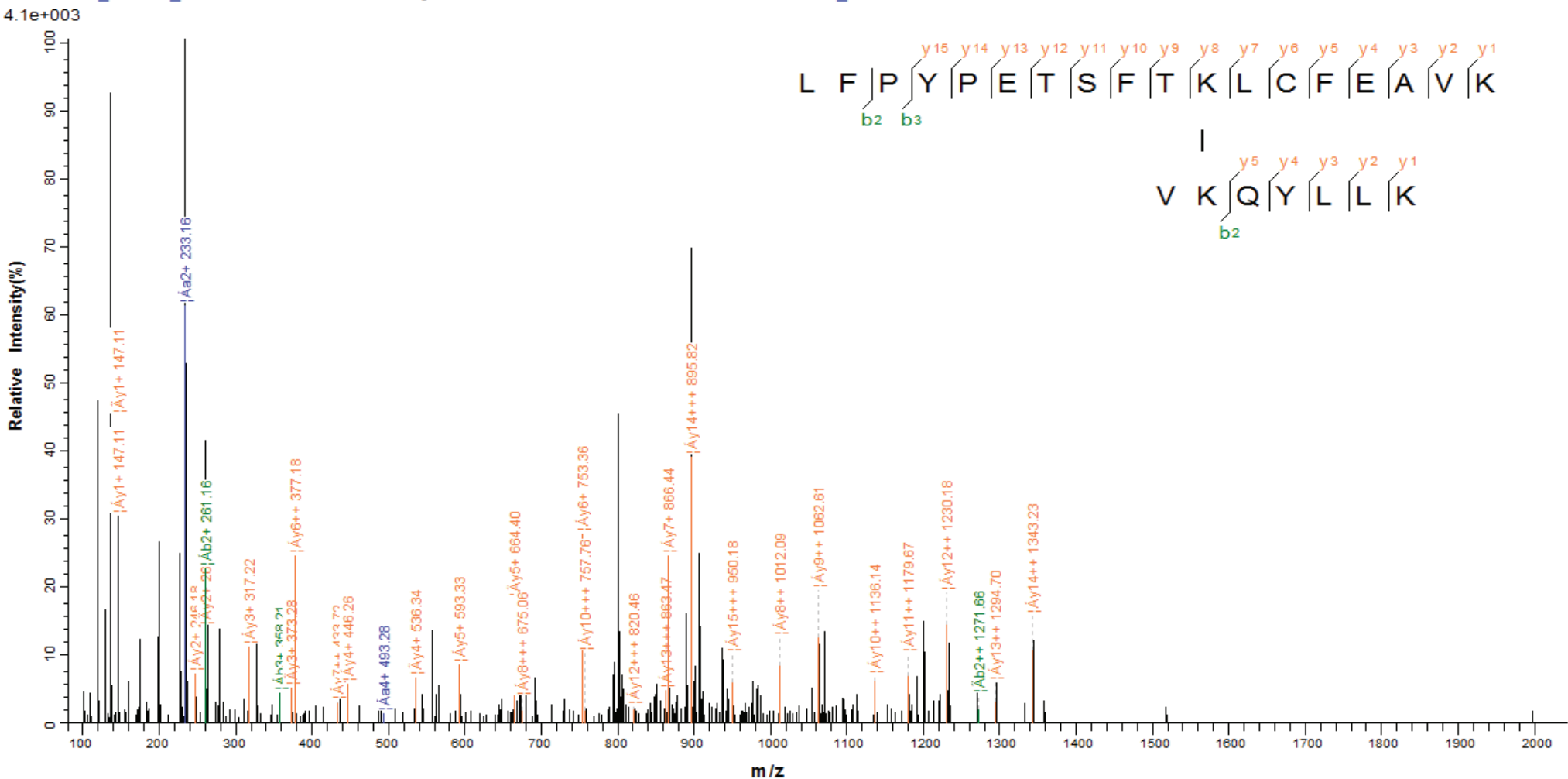
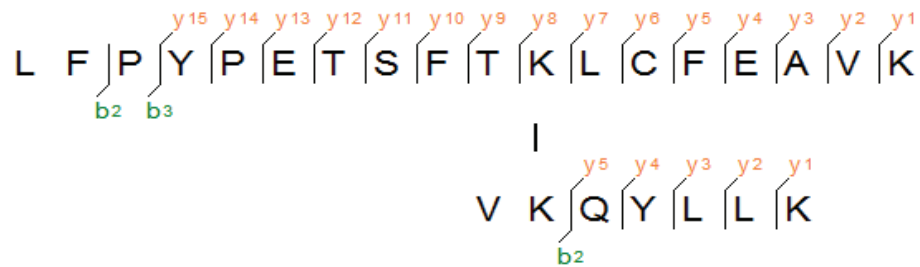
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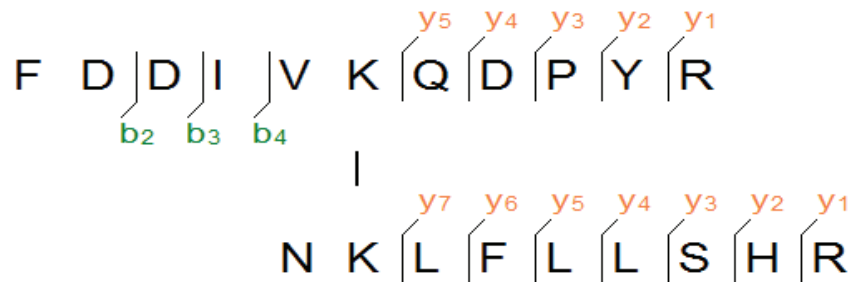
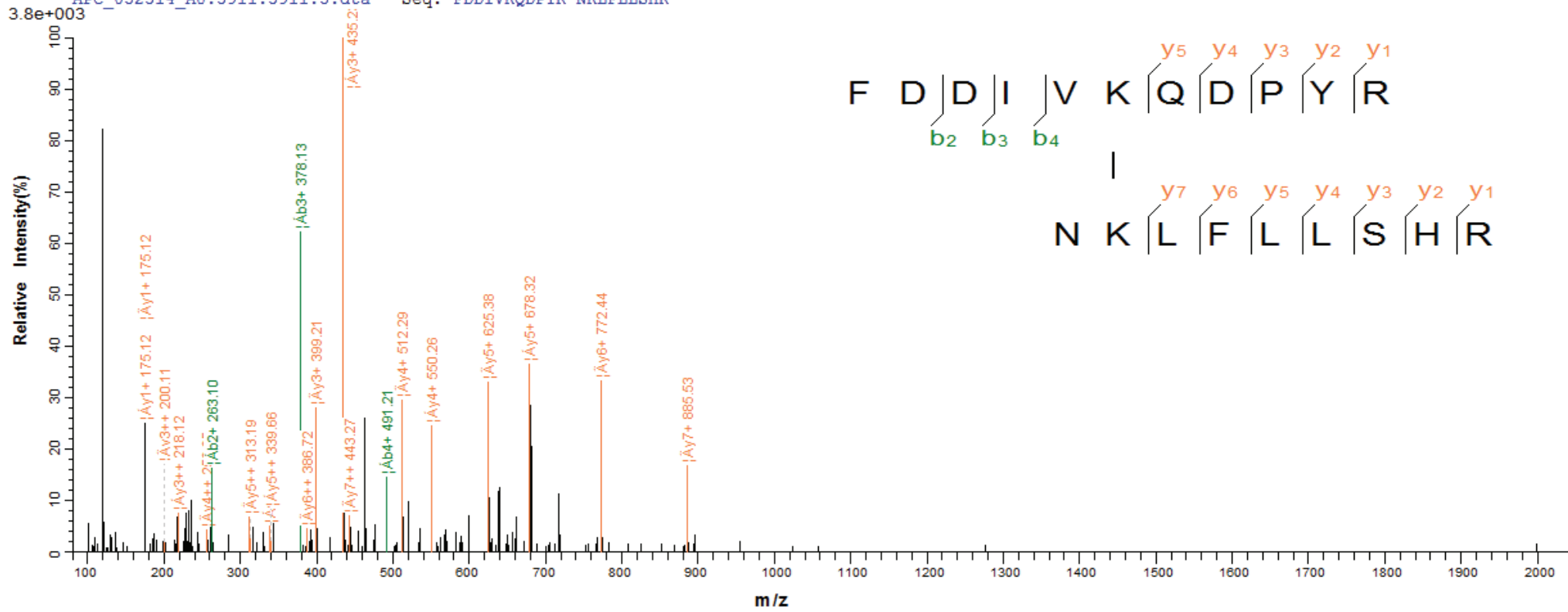
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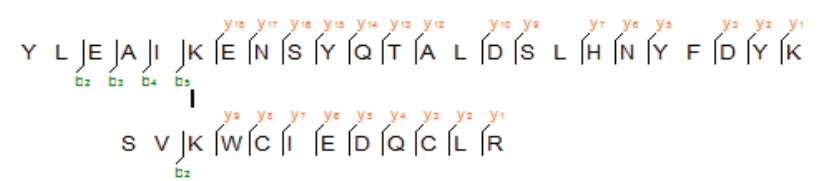
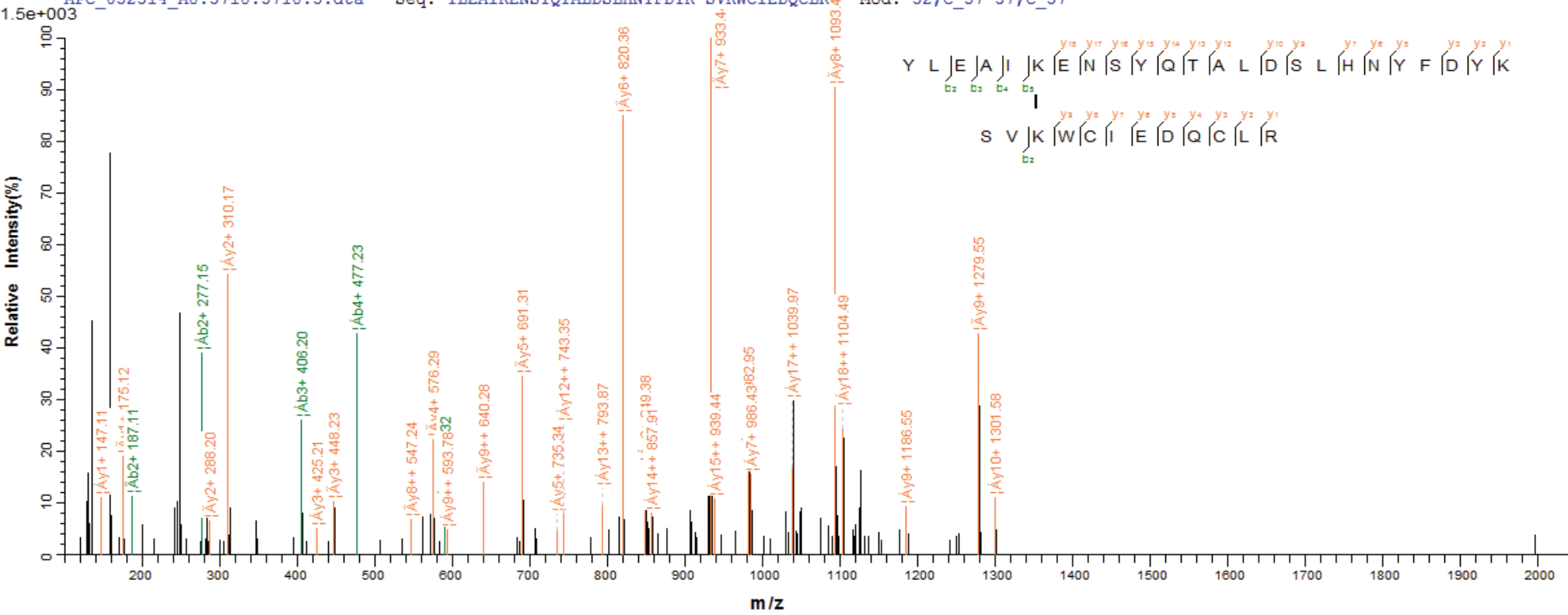
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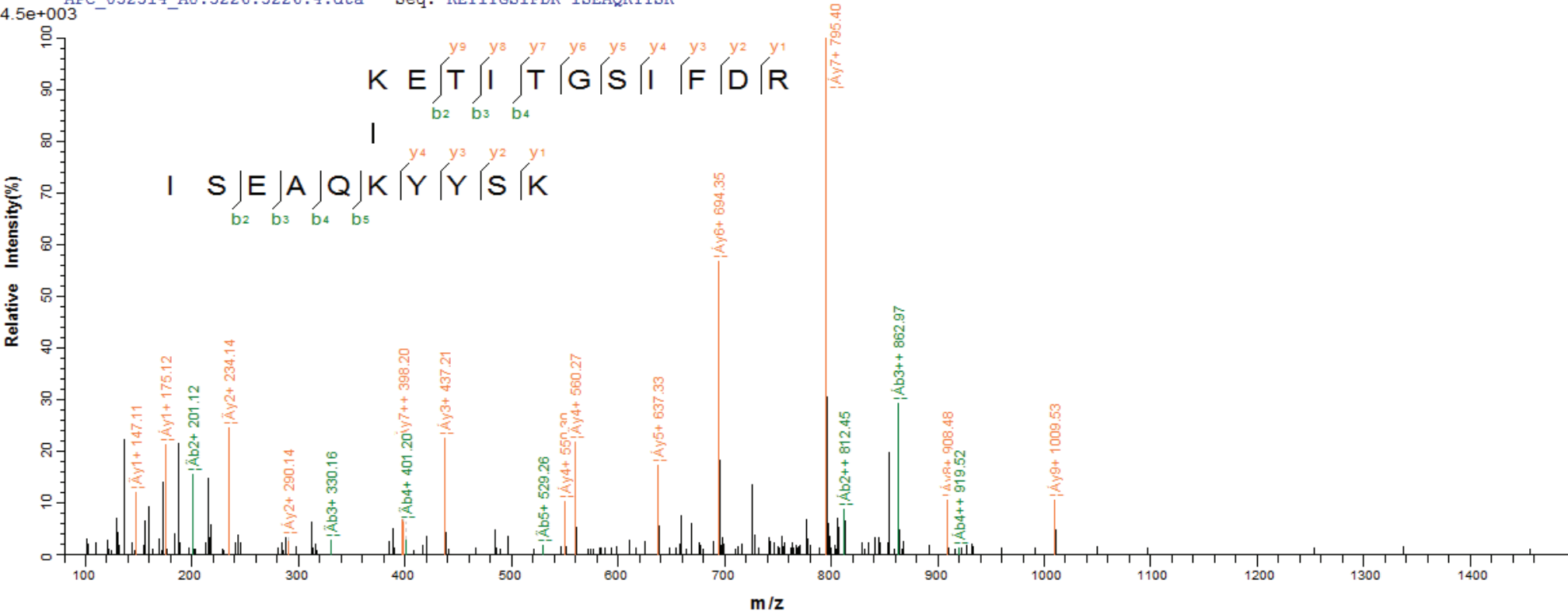


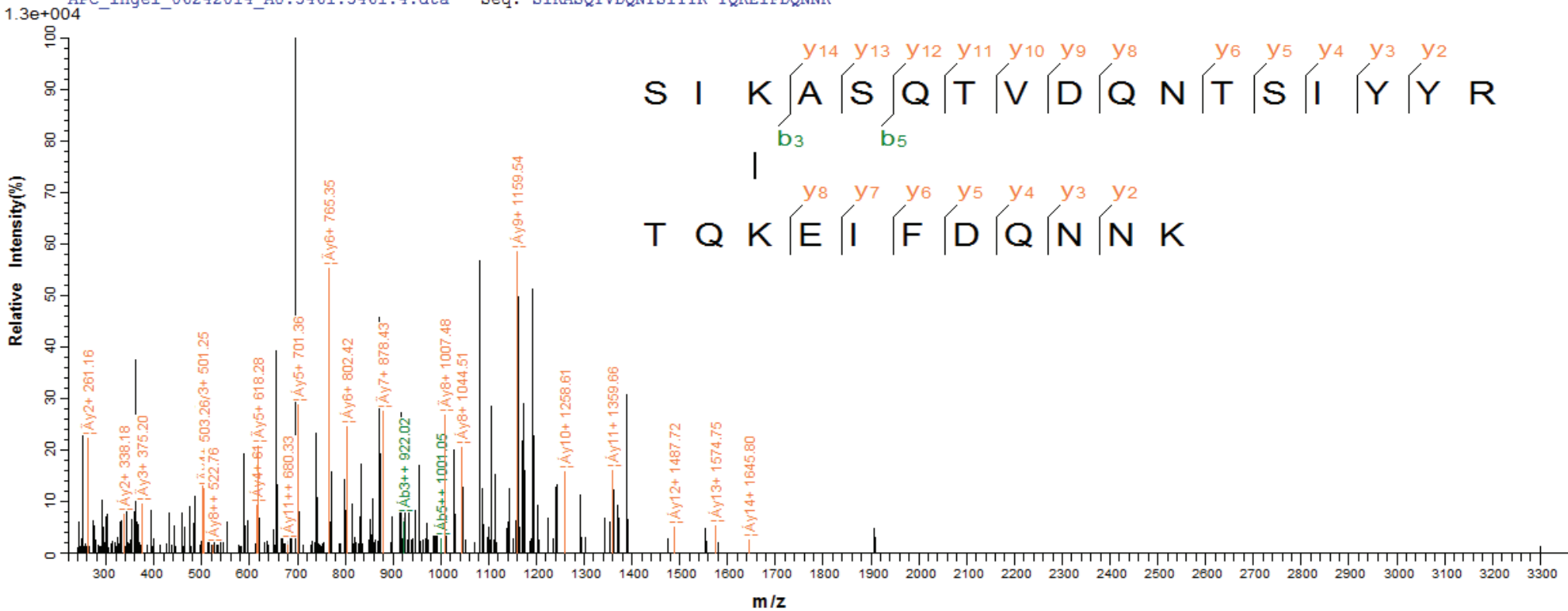
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Mod: 32,C_57 37,C_57





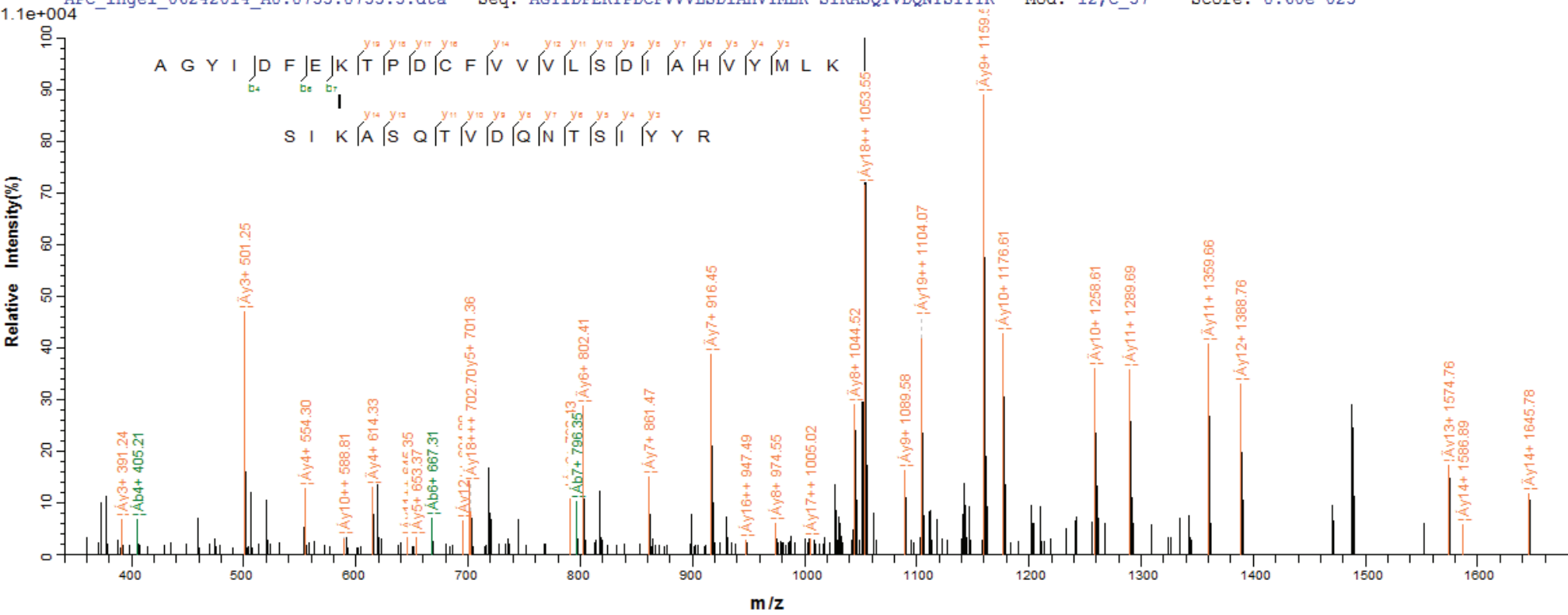


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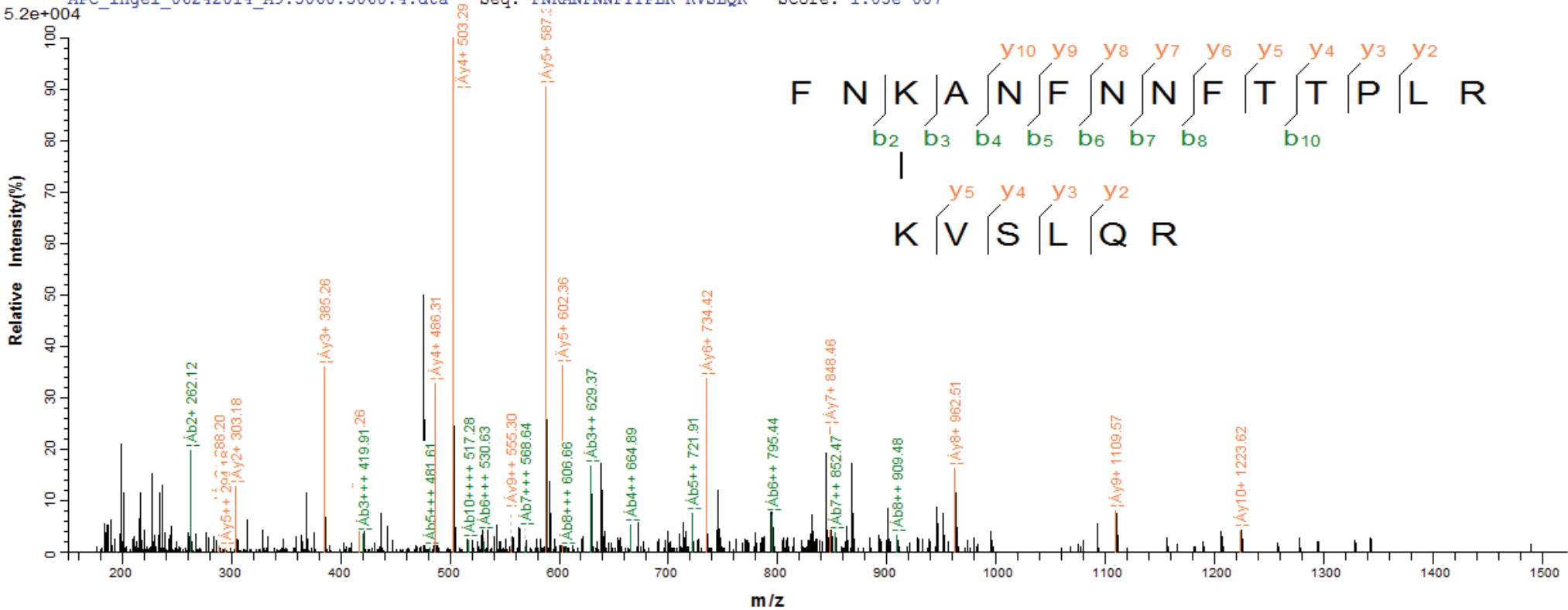
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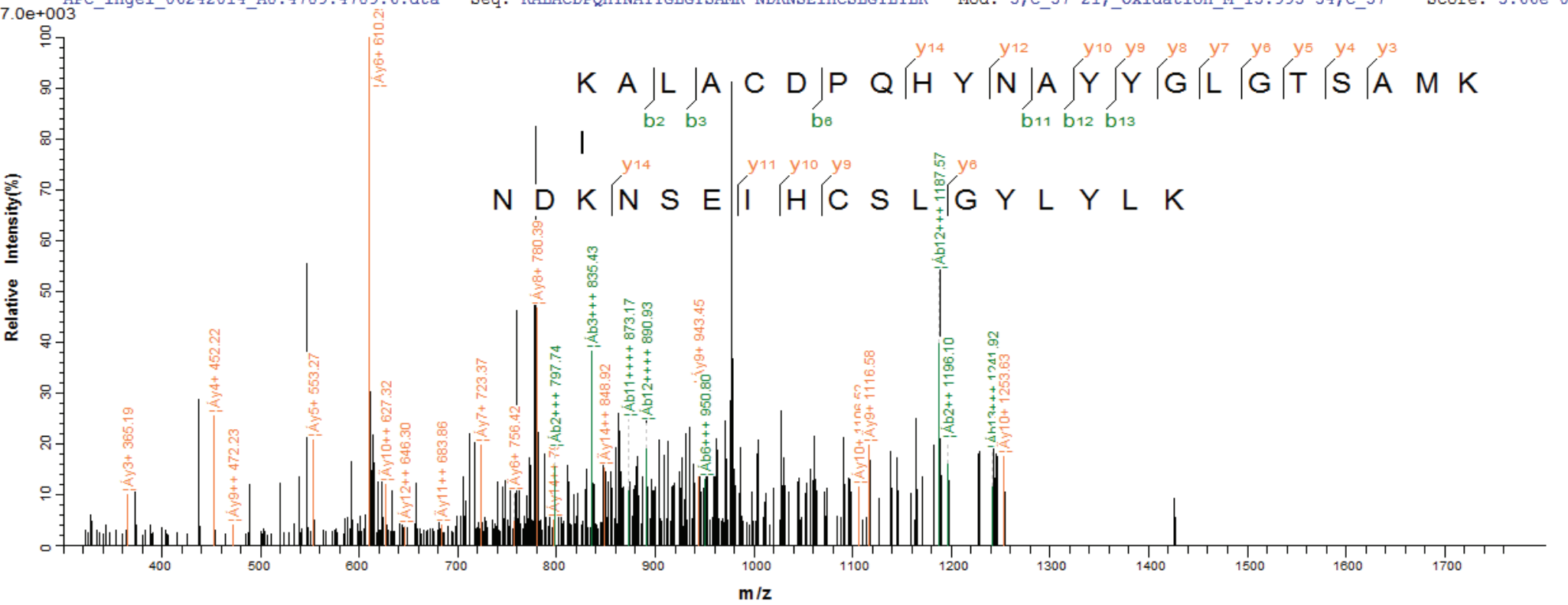
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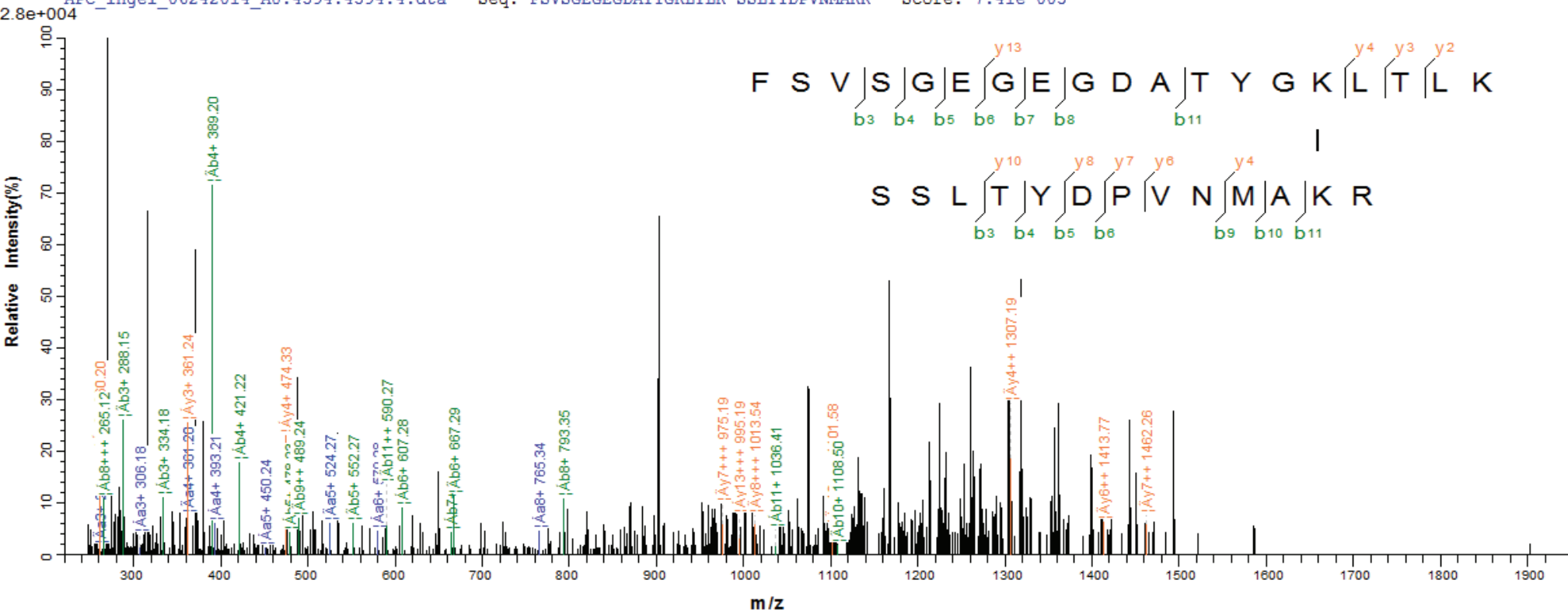
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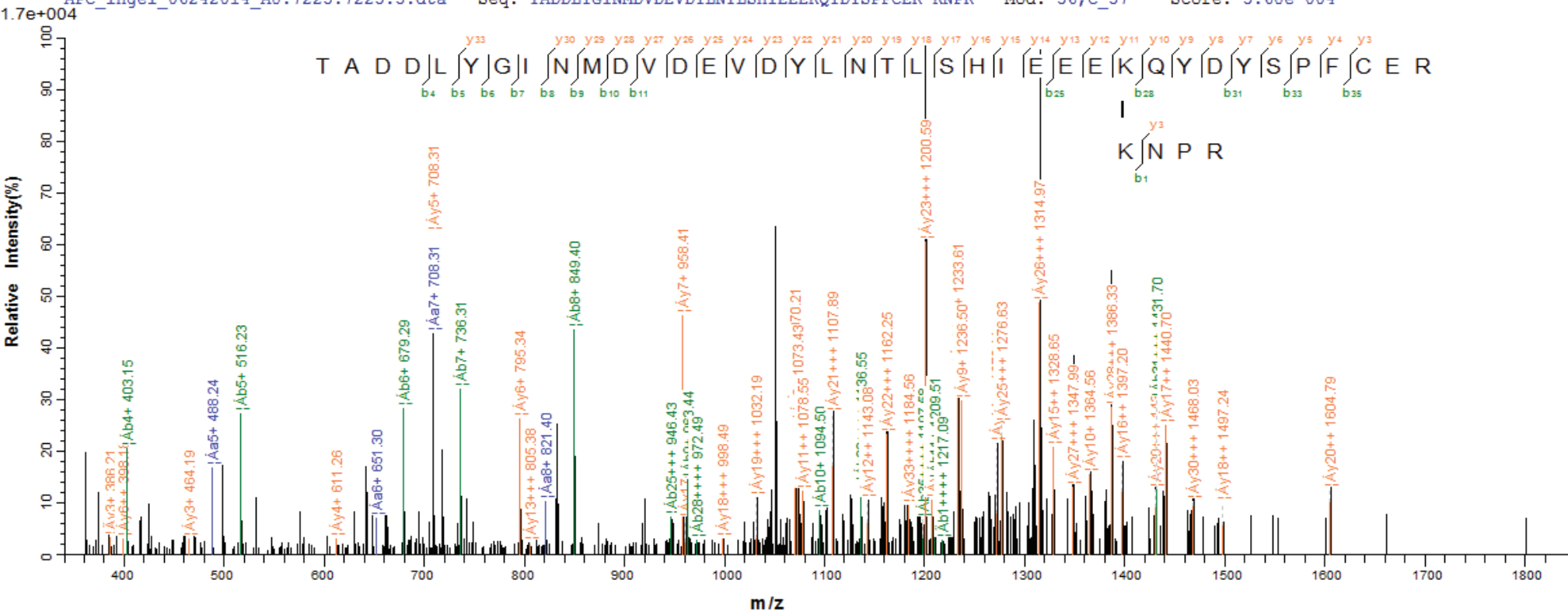


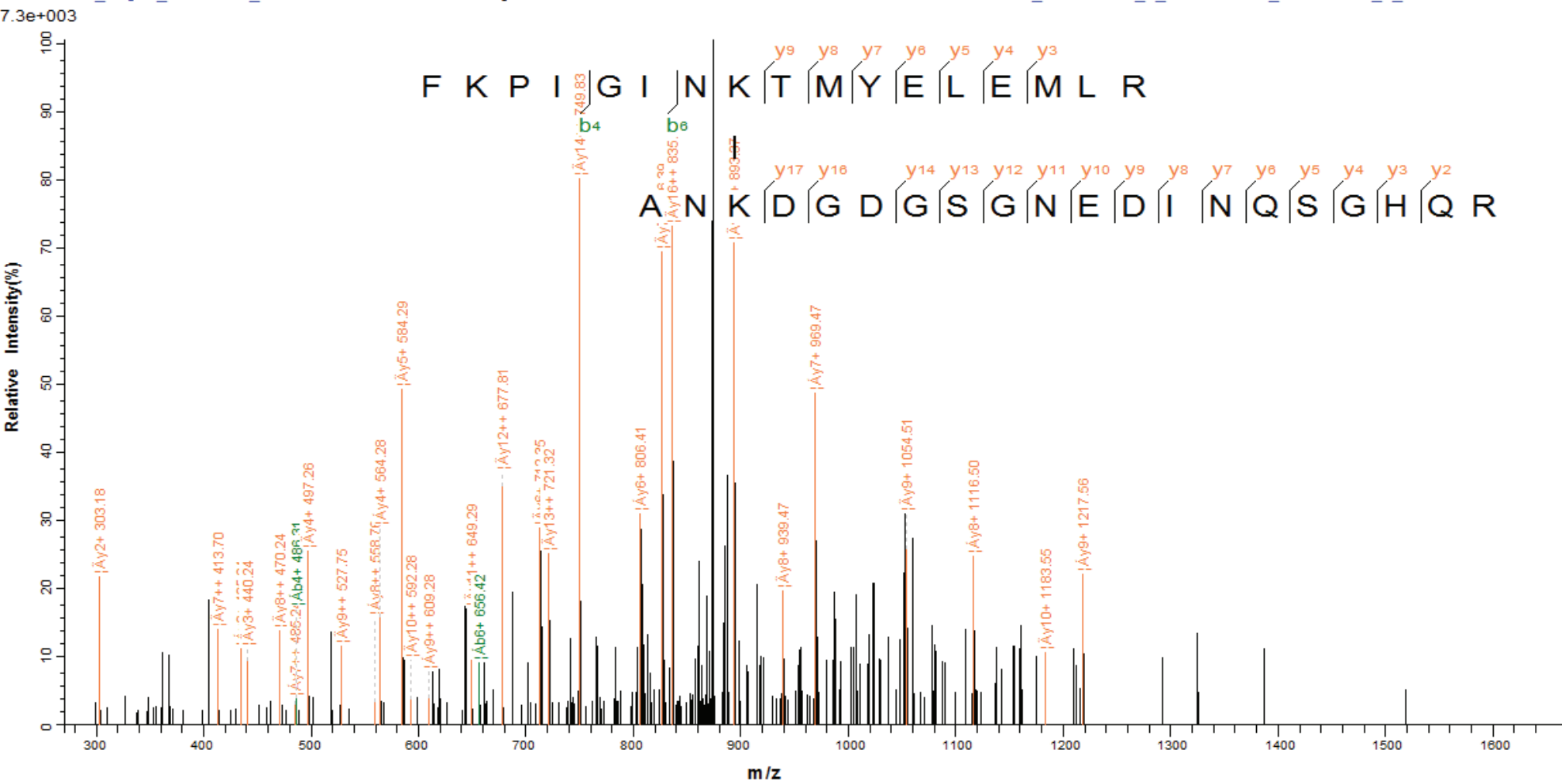
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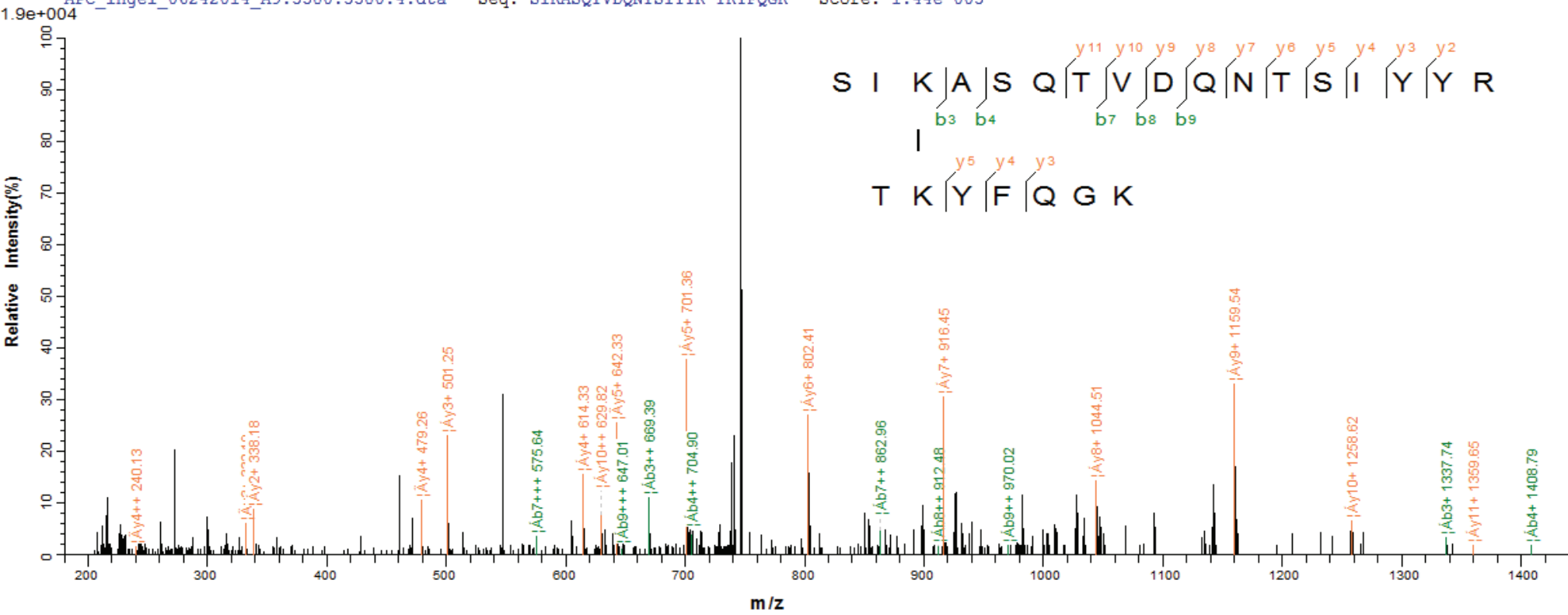




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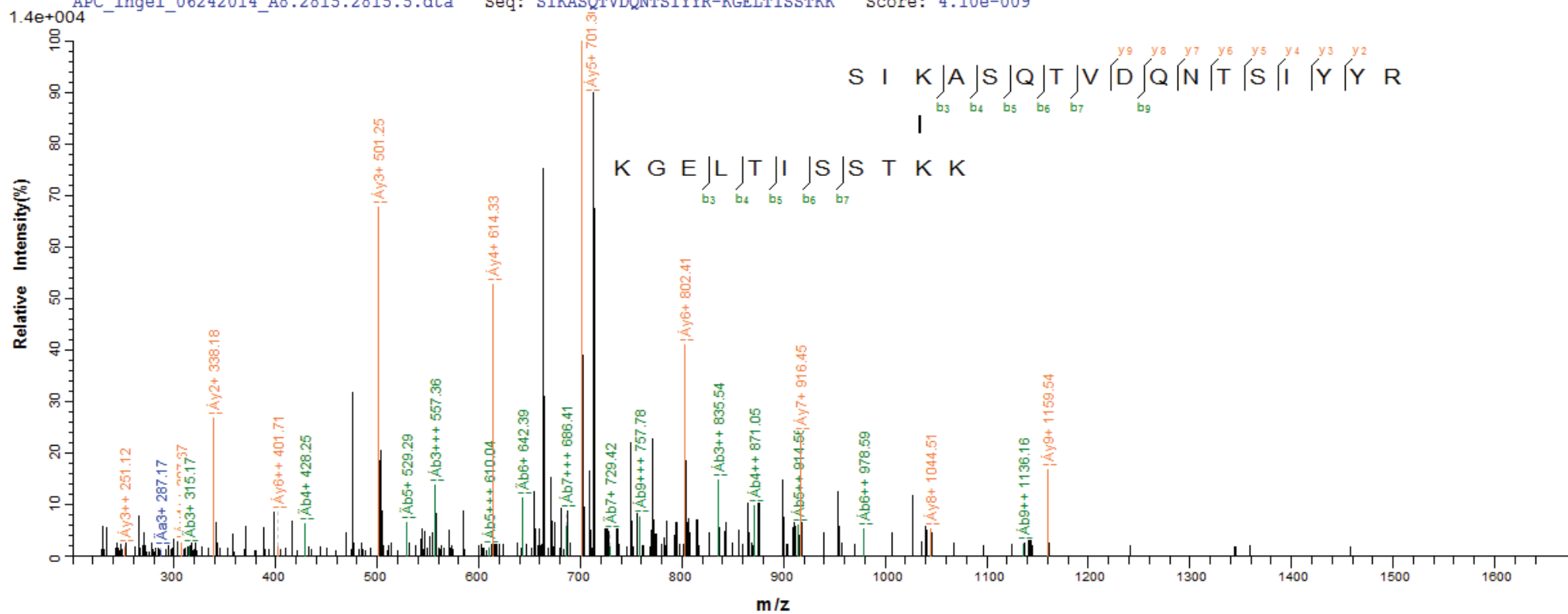
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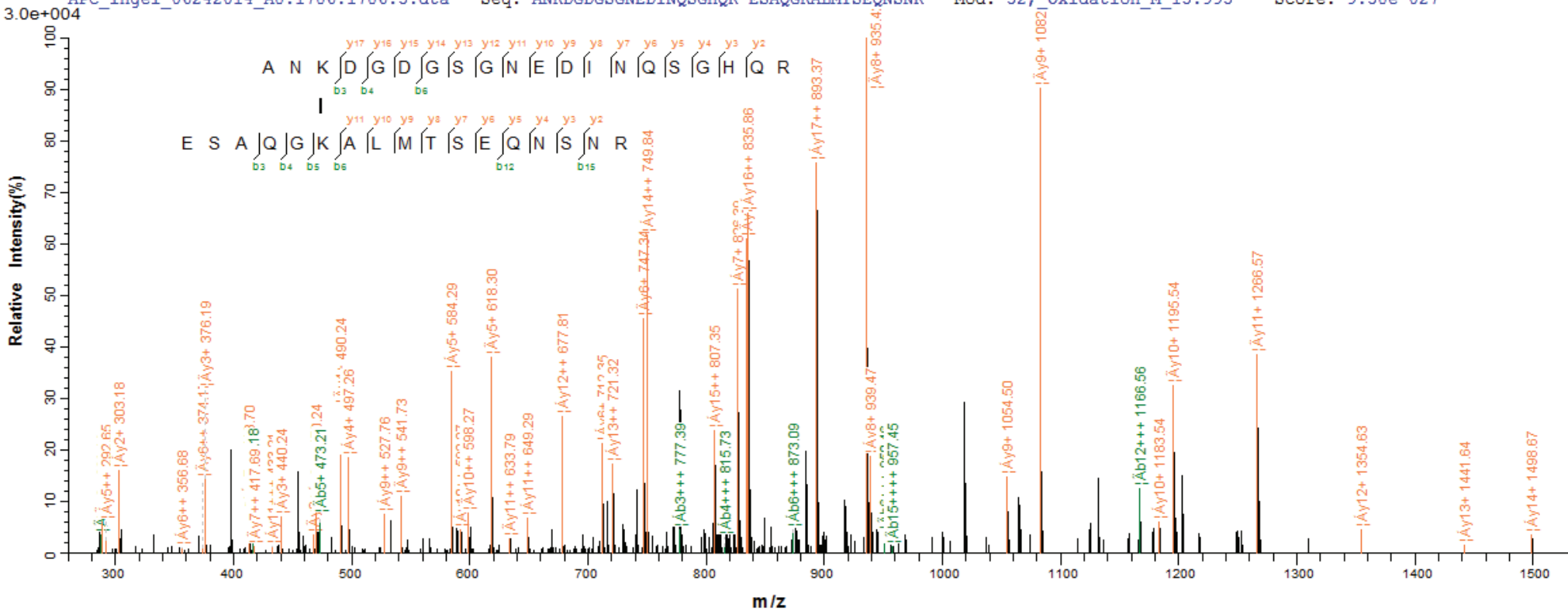


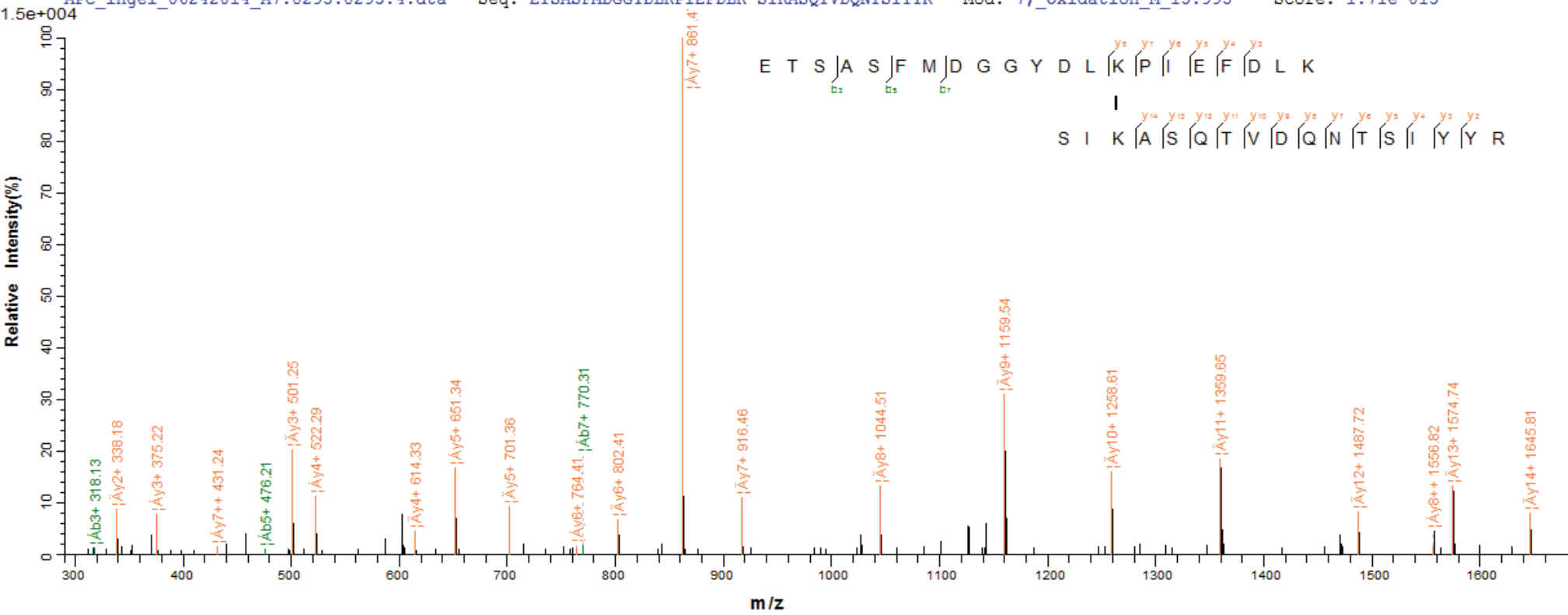
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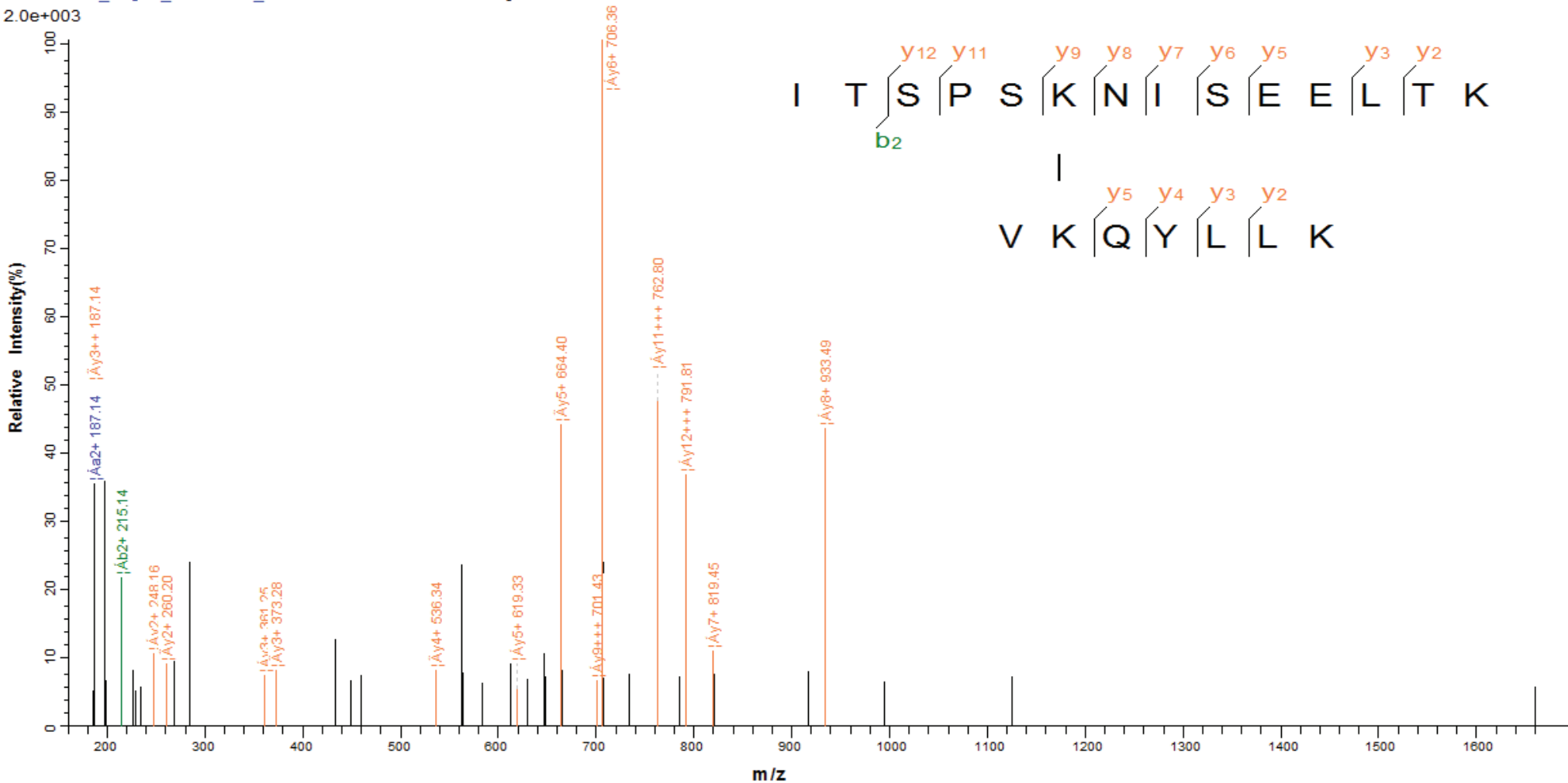
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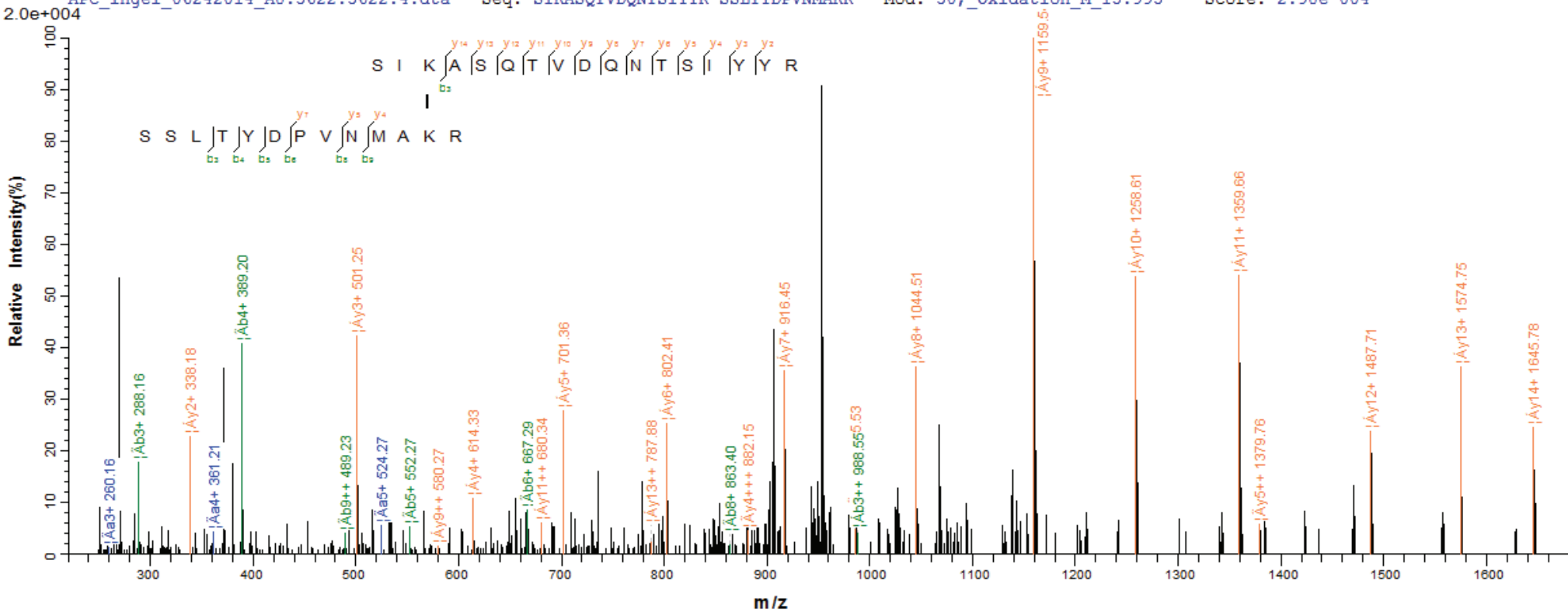
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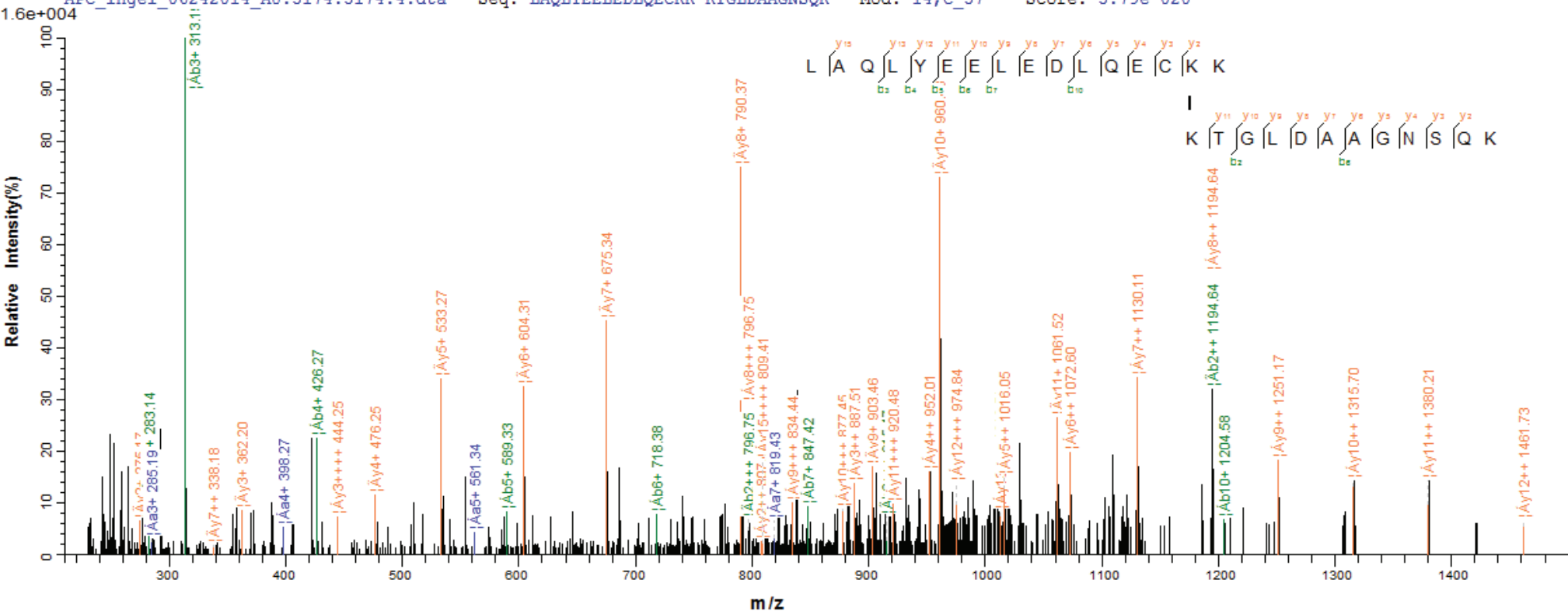




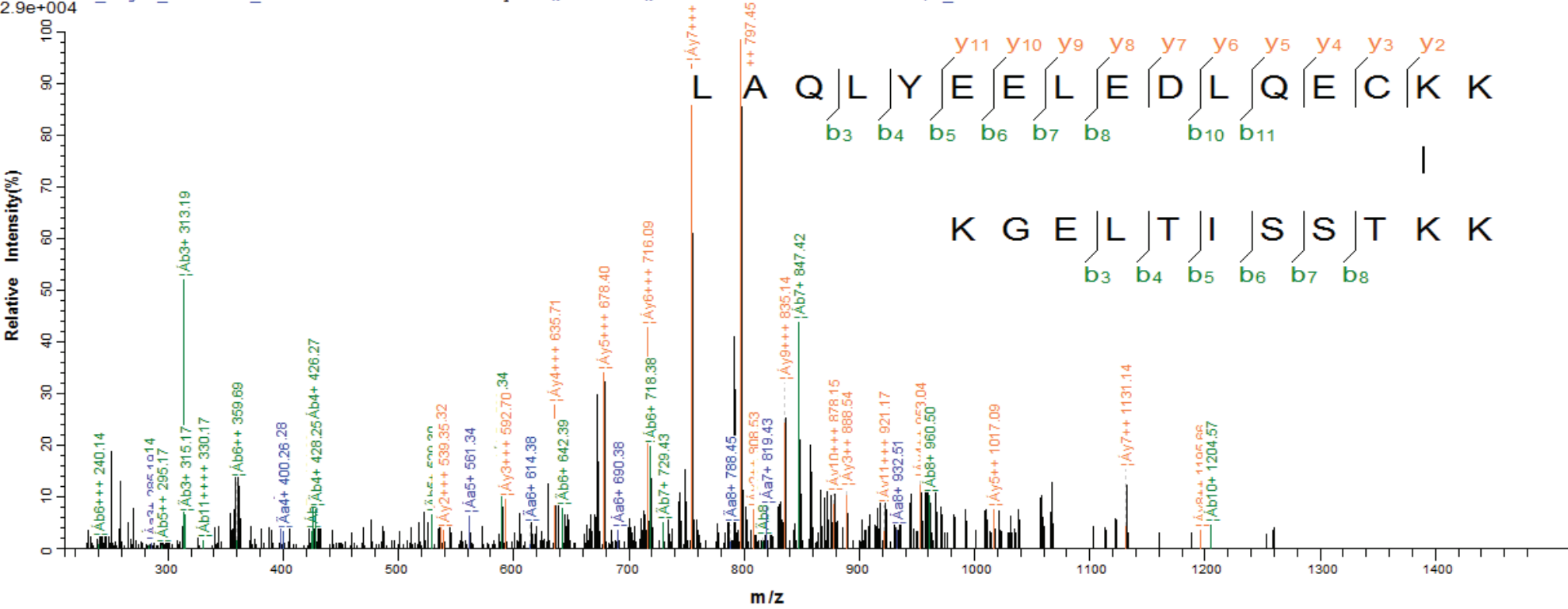




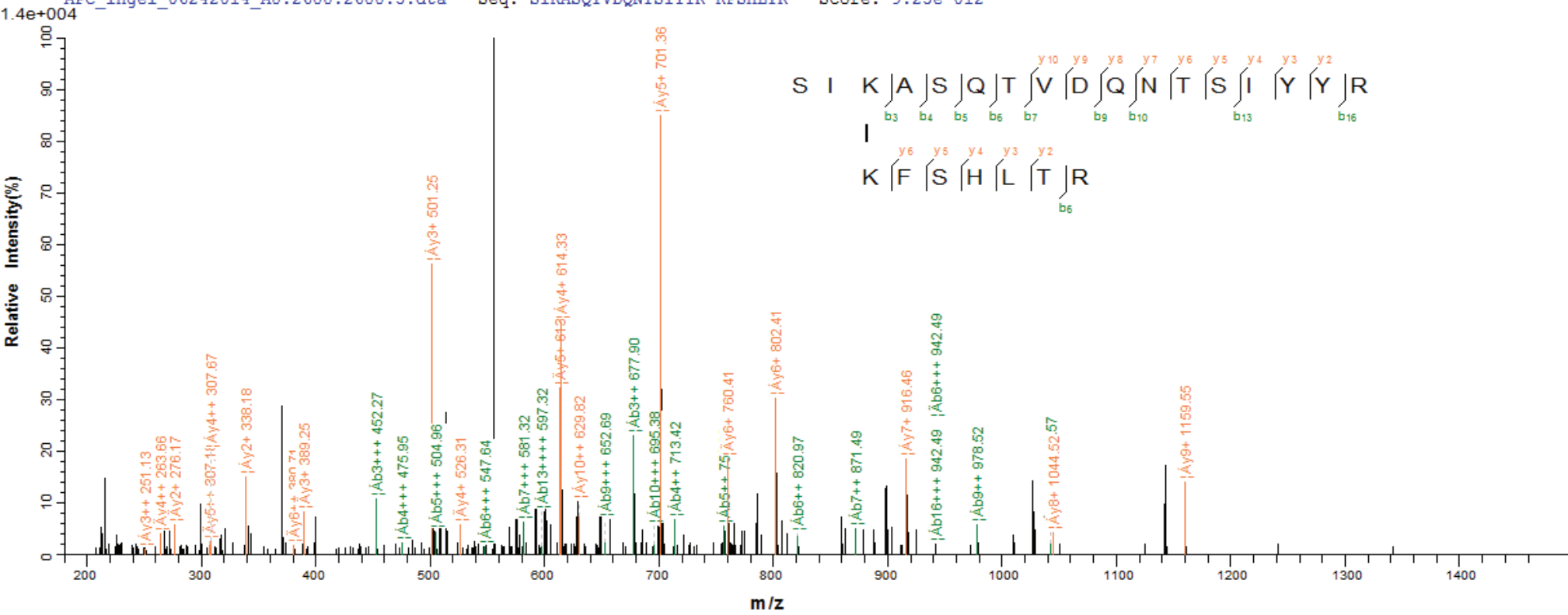
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APC_ingel_06242014_A8.4927.4927.5.dta Seq: LAQLYEELEDLQECKK-KGELTISSTKK Mod: 14,C_57 Score: 8.59e-010



L A Q L Y E E L E D L Q E C K K
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 K G E L T I S S T K K
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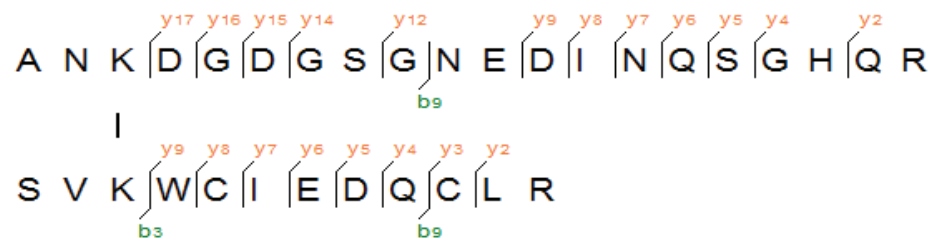
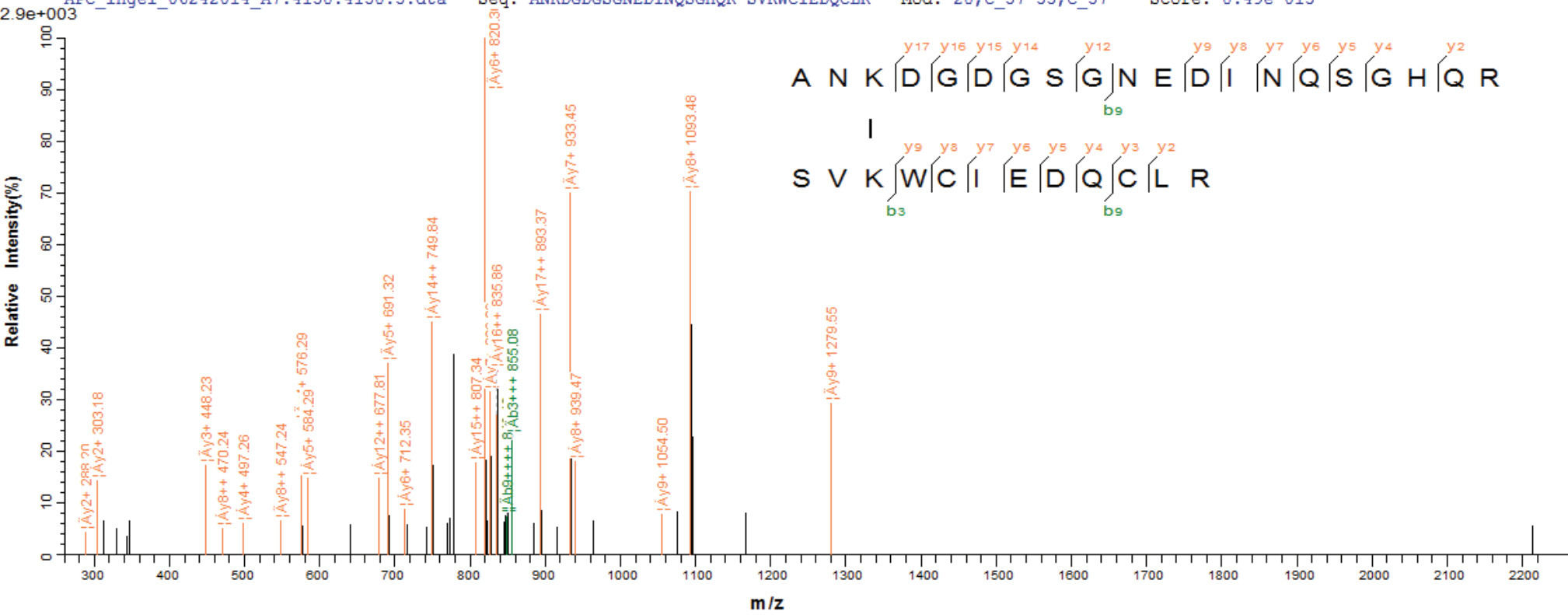


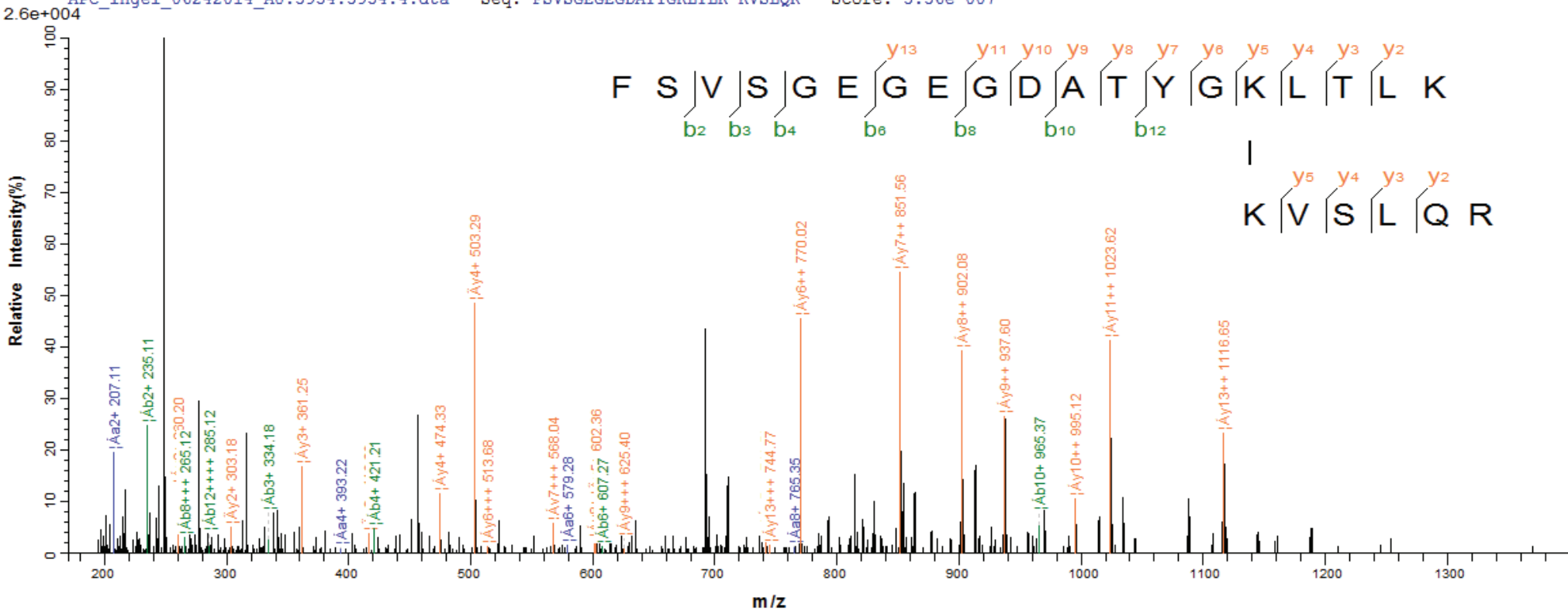
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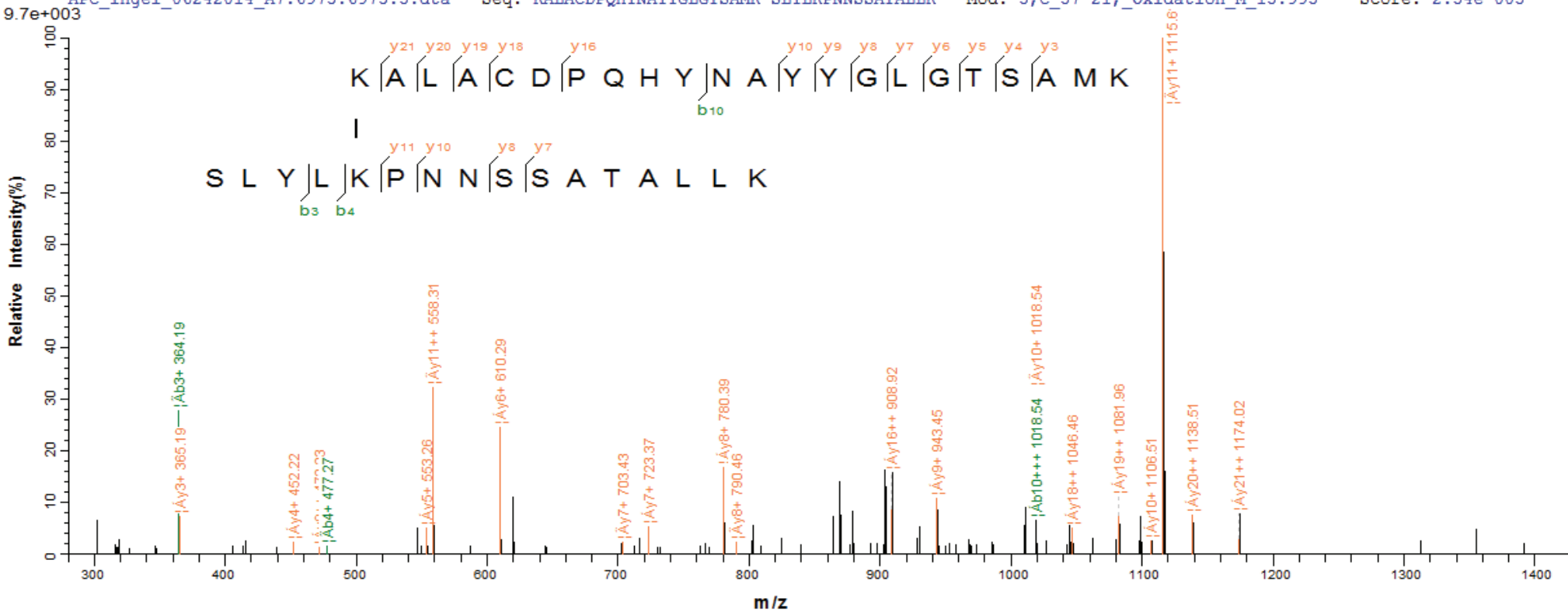
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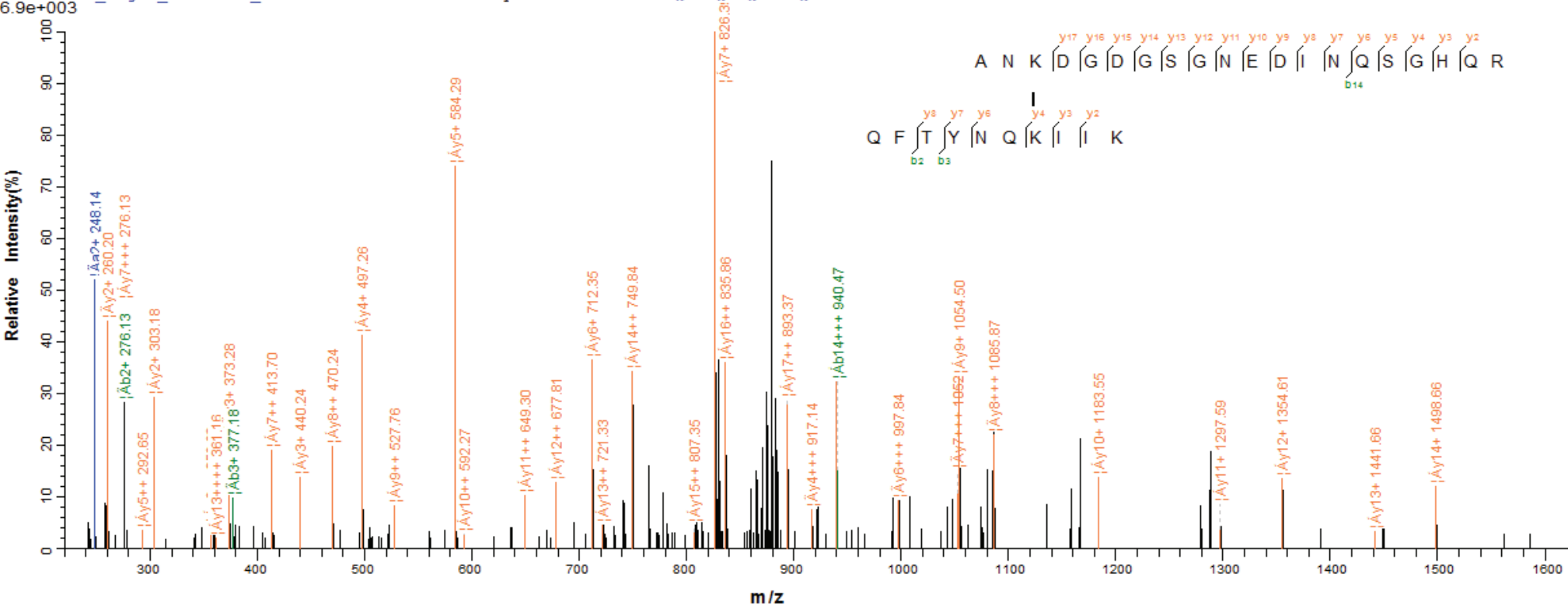
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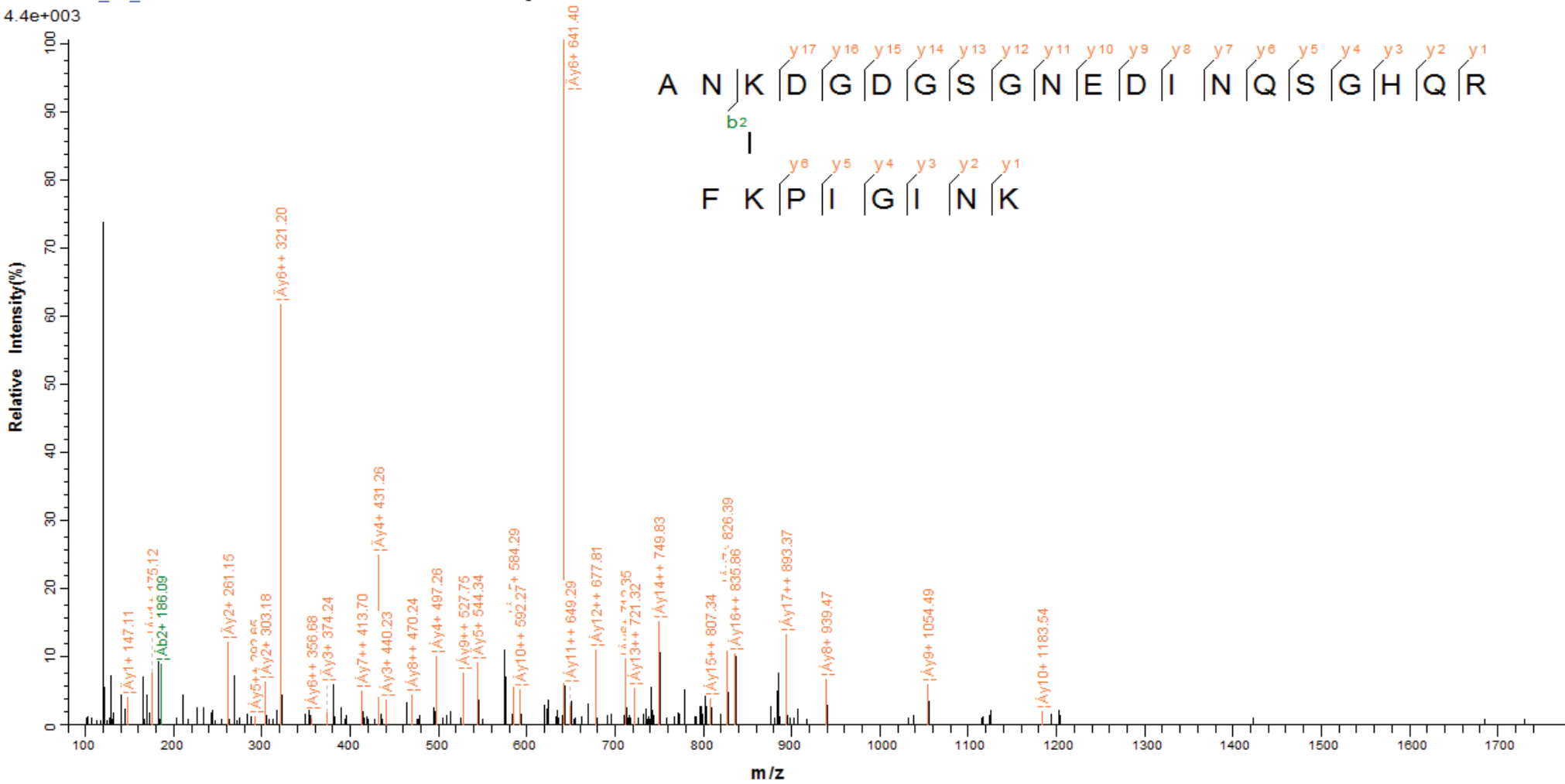
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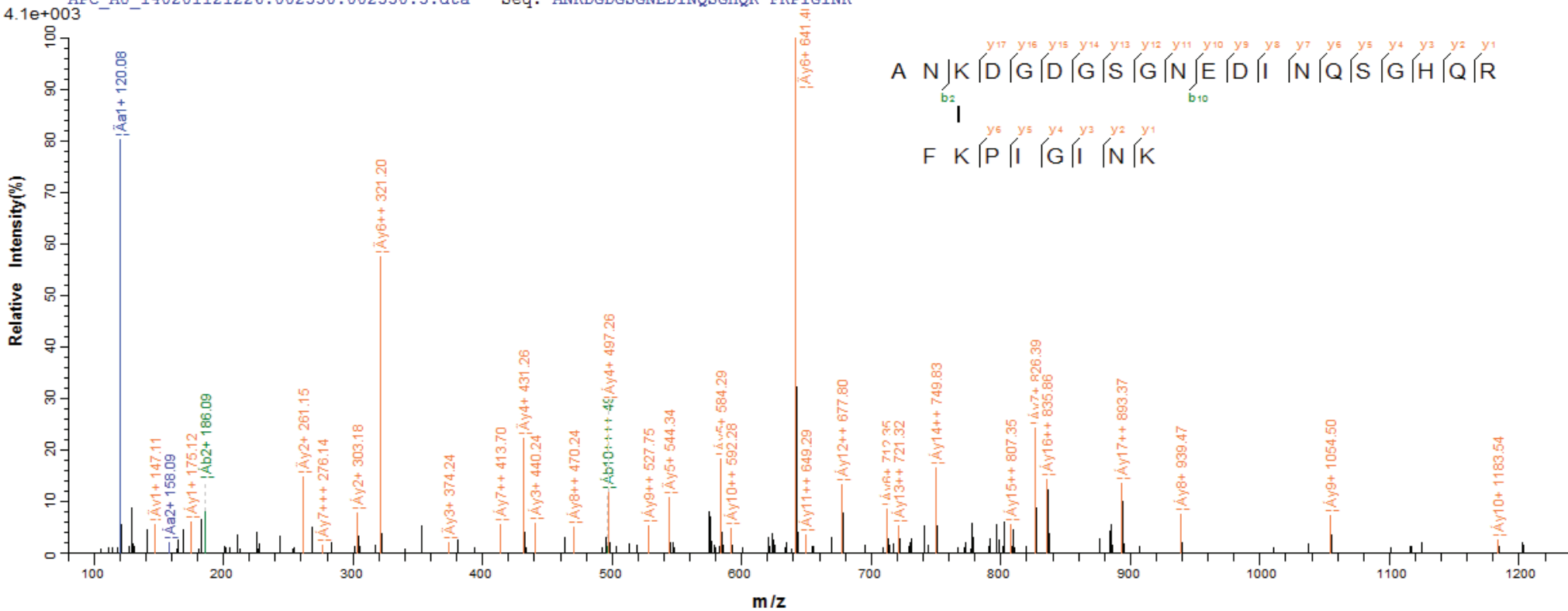


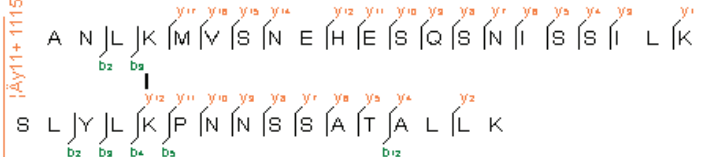
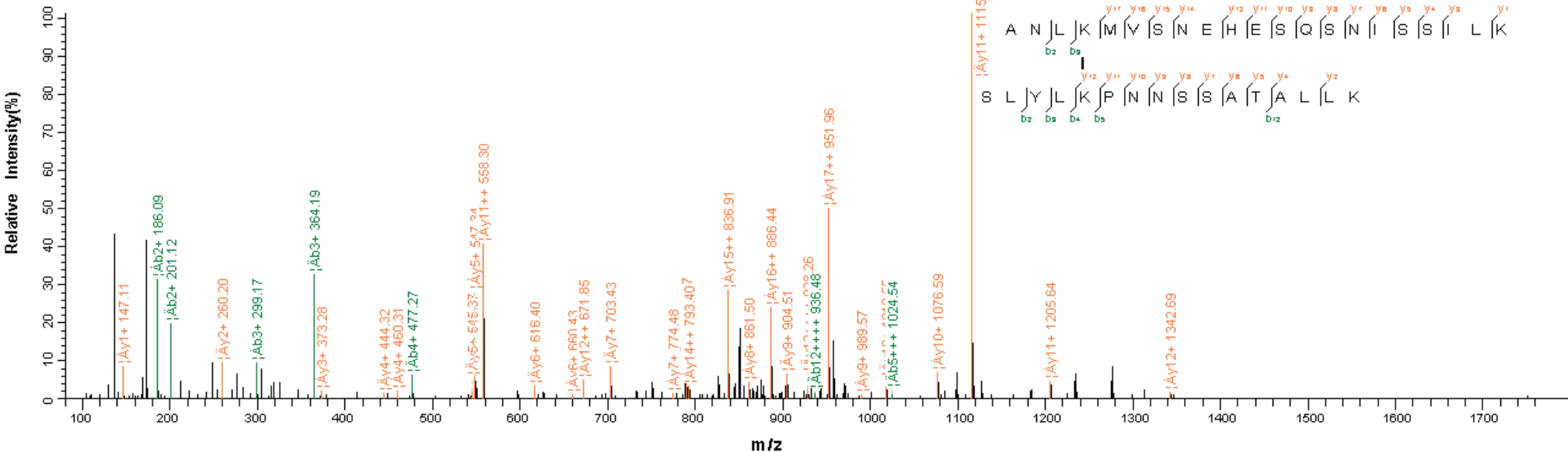


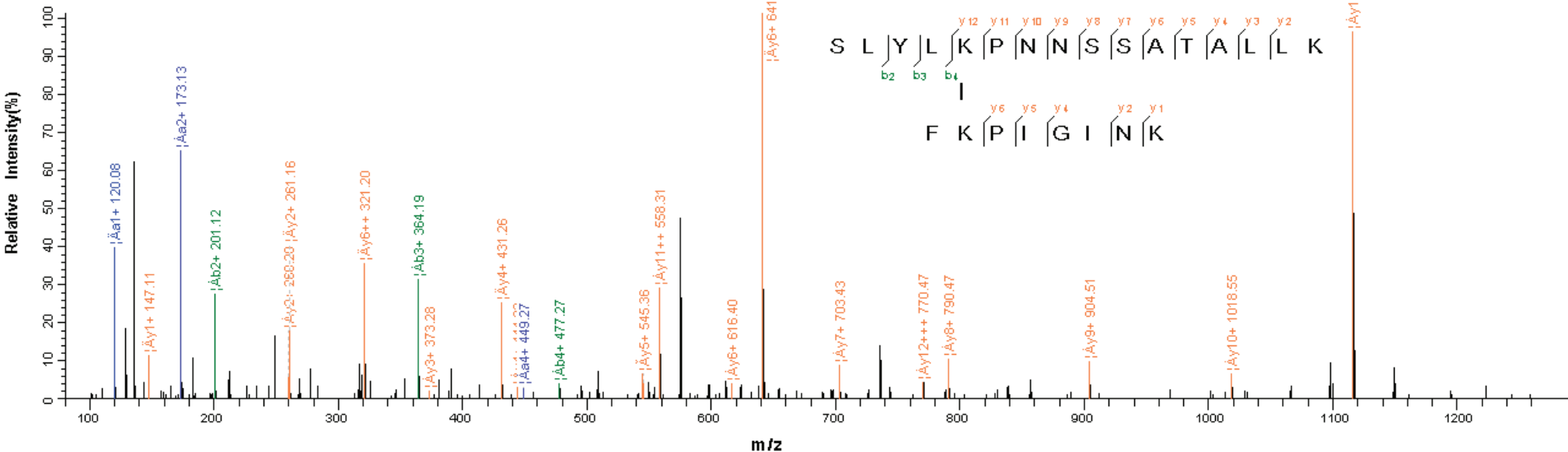




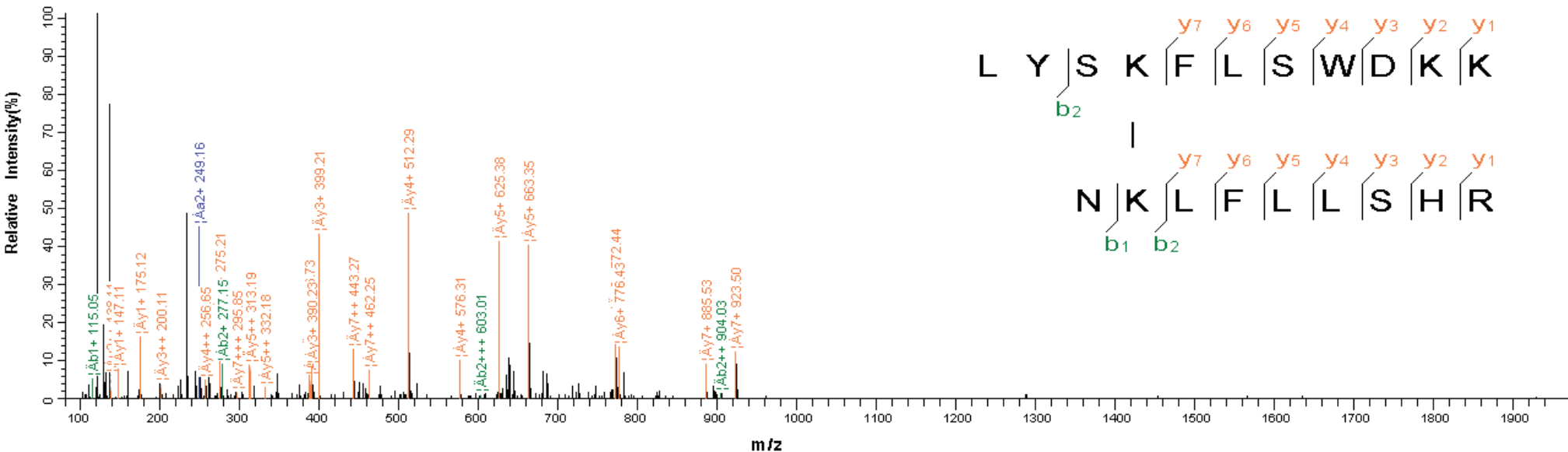




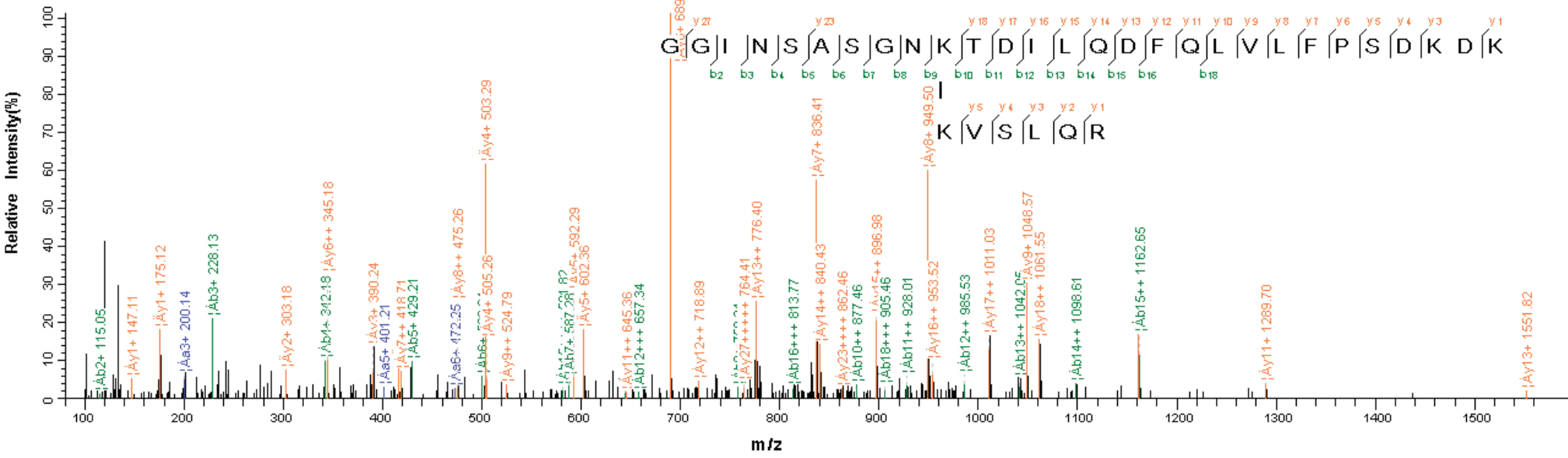




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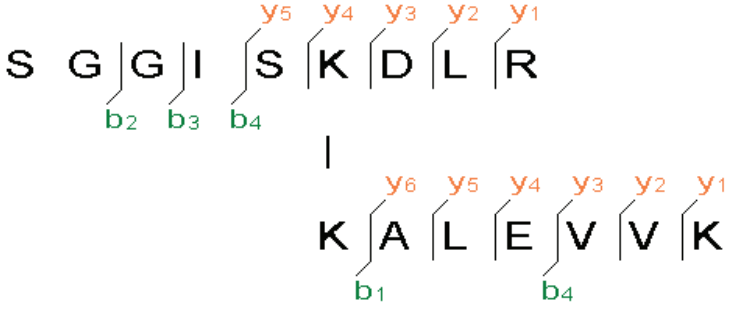
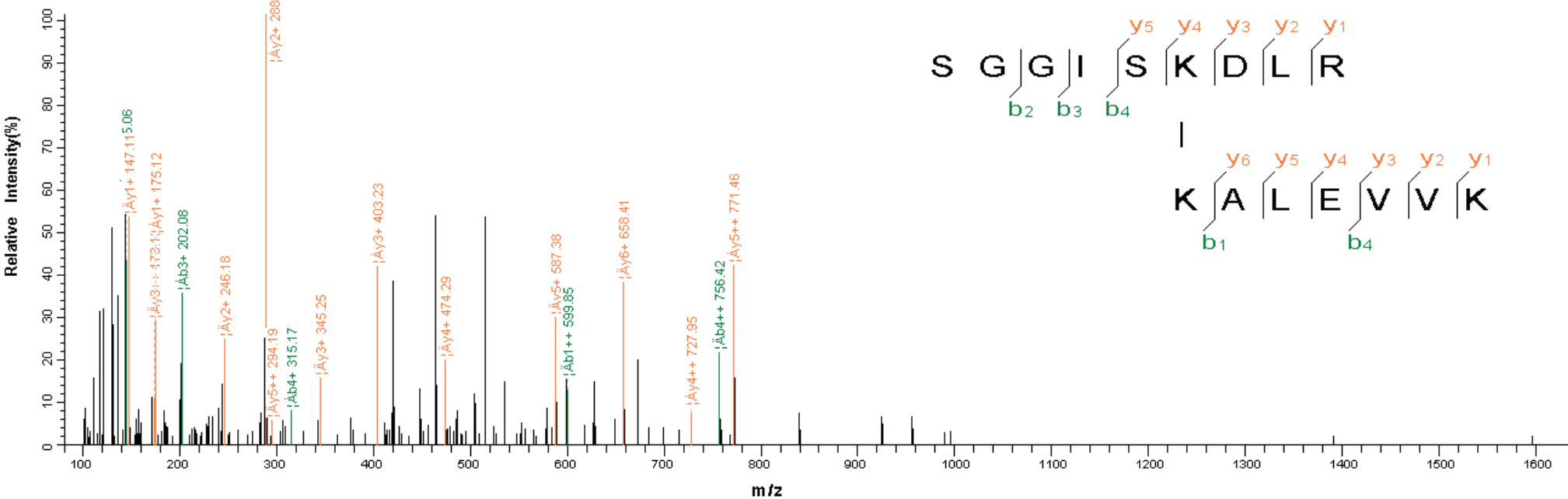


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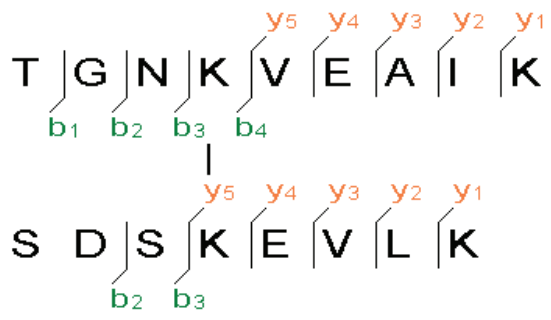
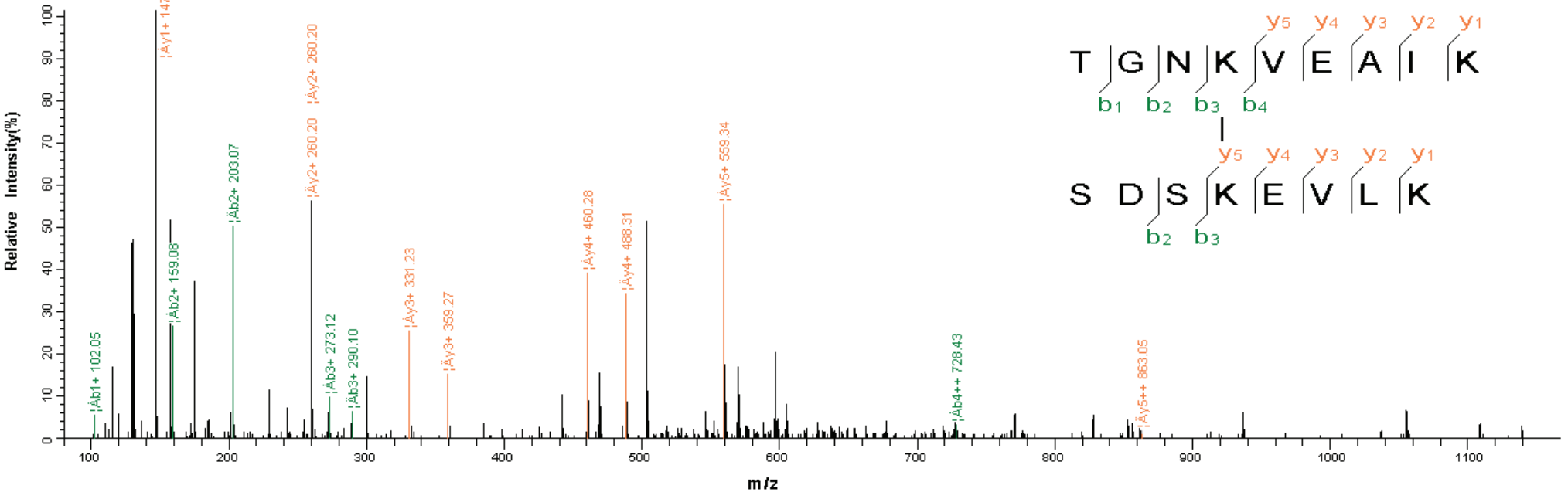


GGINSASGNKTDILQDFQLVLFPPSDKDK-KVSLQR
 b2 b3 b4 b5 b6 b7 b8 b9 b10 b11 b12 b13 b14 b15 b16 b18
 y1 y2 y3 y4 y5 y6 y7 y8 y9 y10 y11 y12 y13 y14 y15 y16 y17 y18 y23
 K V S L Q R

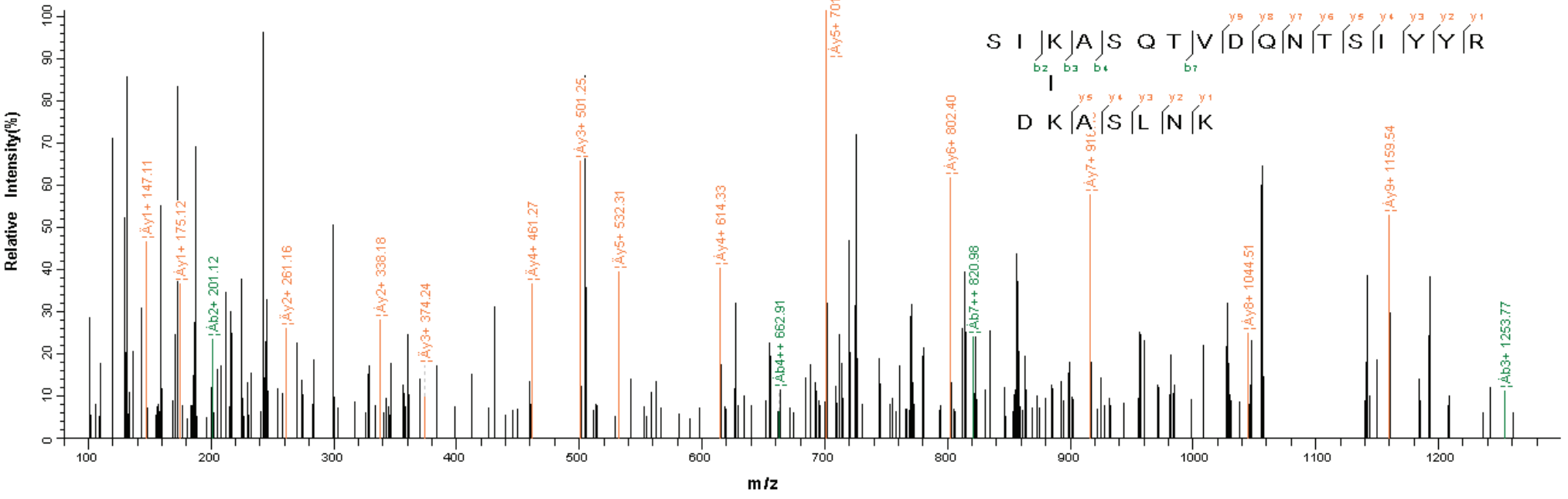
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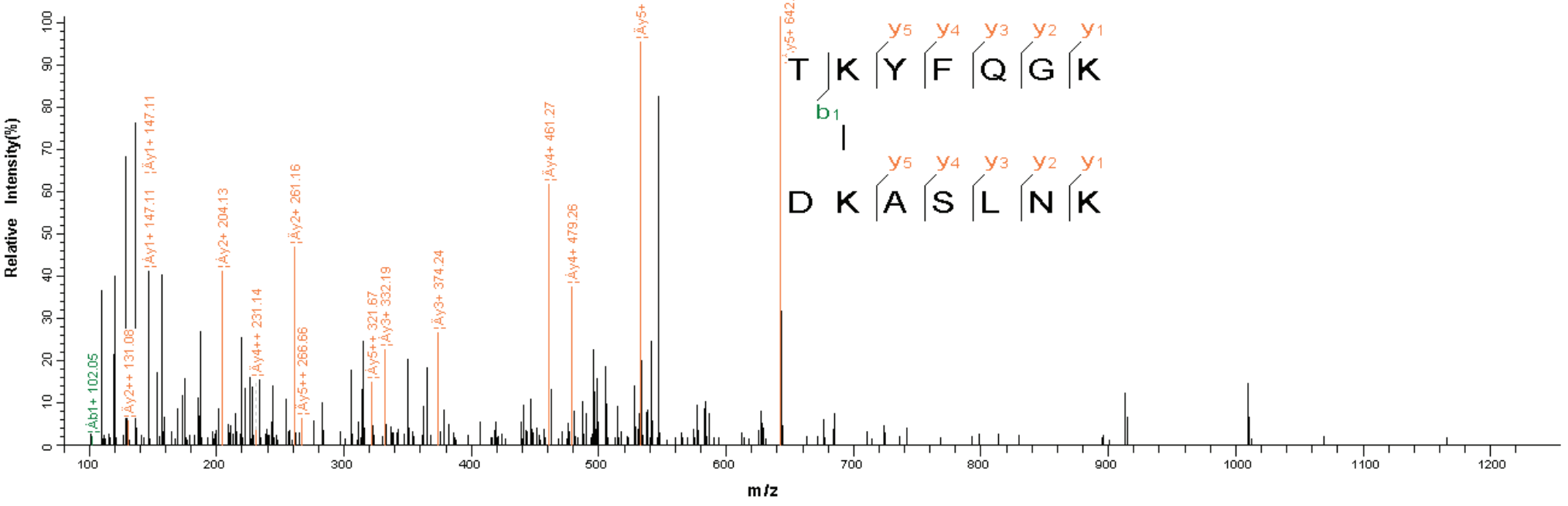


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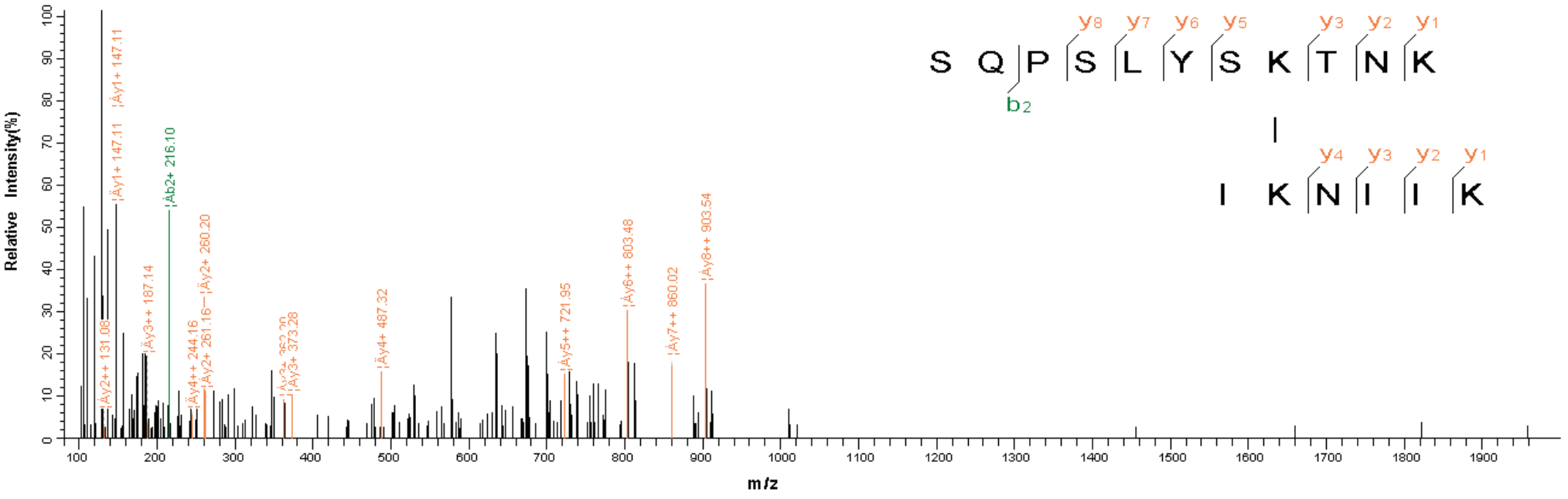


S I K A S Q T V D Q N T S I Y Y R
bz b3 b4 b7
y9 y8 y7 y6 y5 y4 y3 y2 y1
D K A S L N K
y5 y4 y3 y2 y1

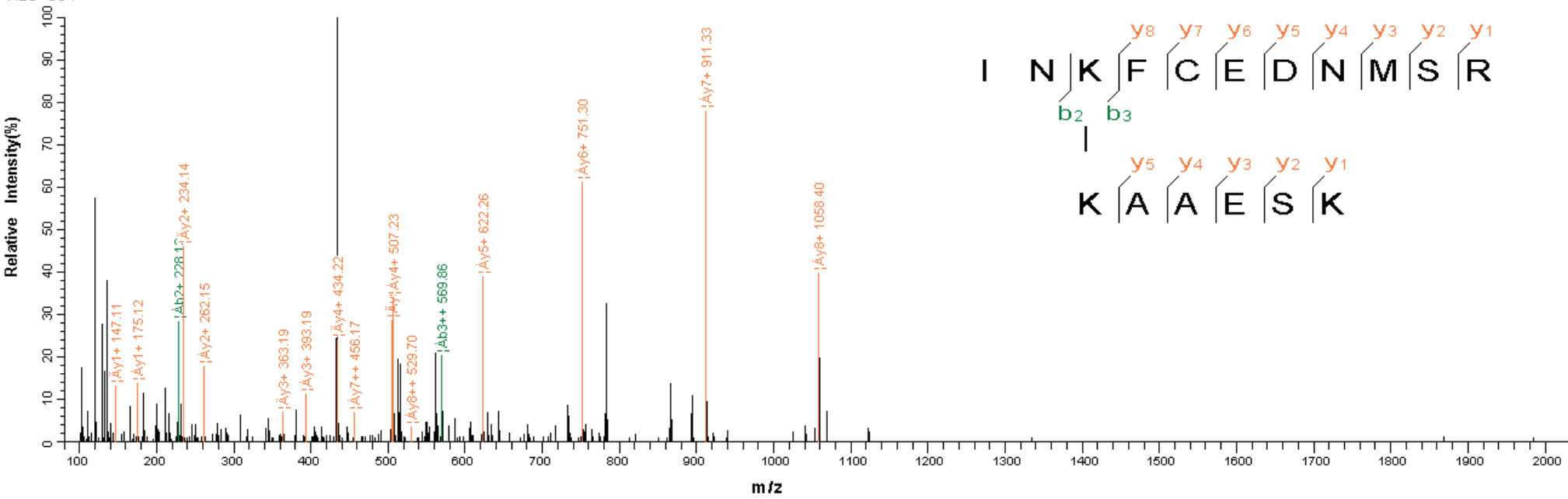
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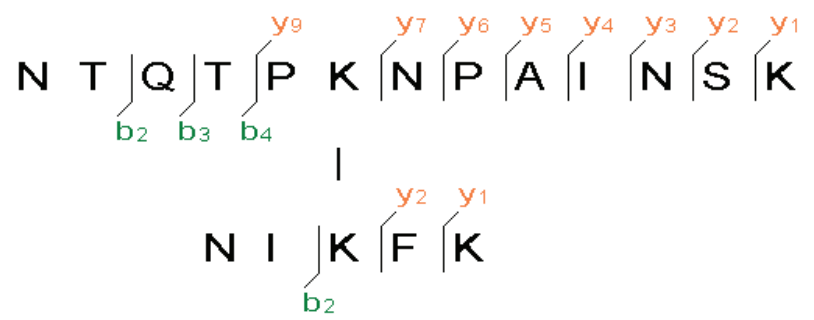
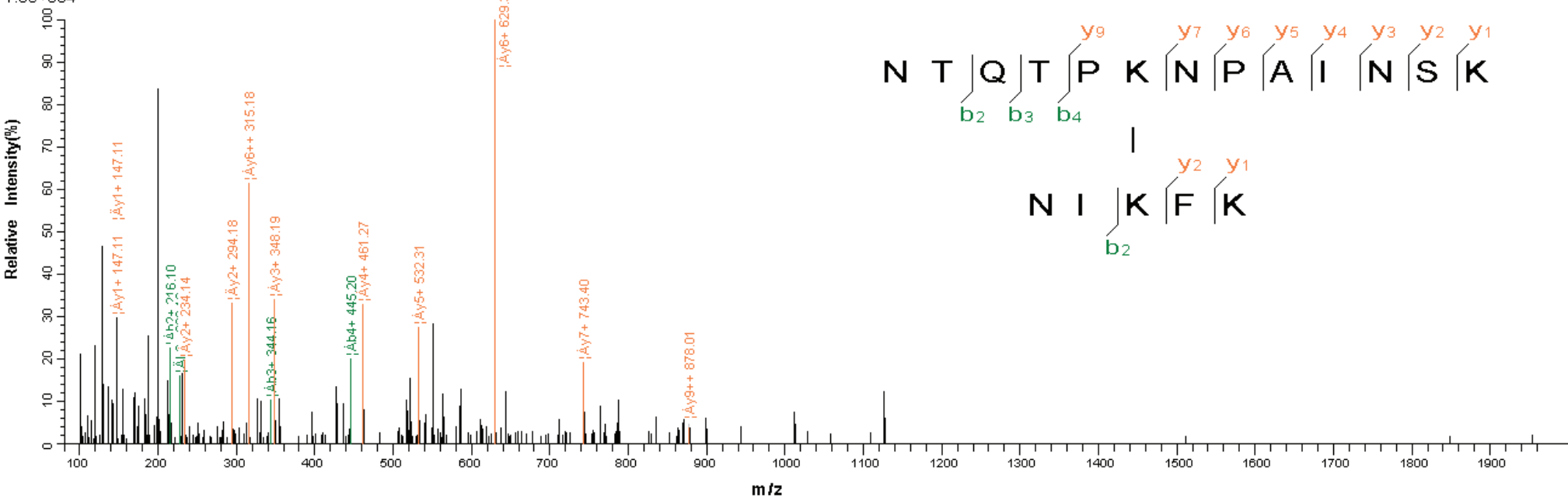
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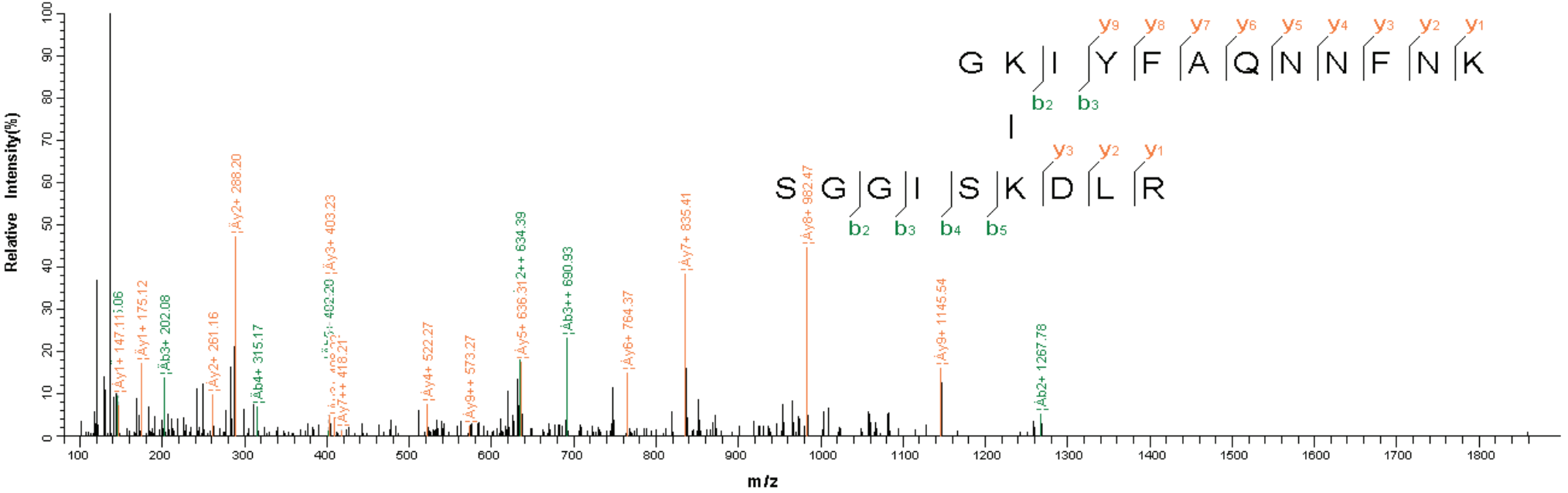
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PC_A9_140131205754.003758.003758.4.dta Seq: NTQTPKNPAINSK-NIRFK

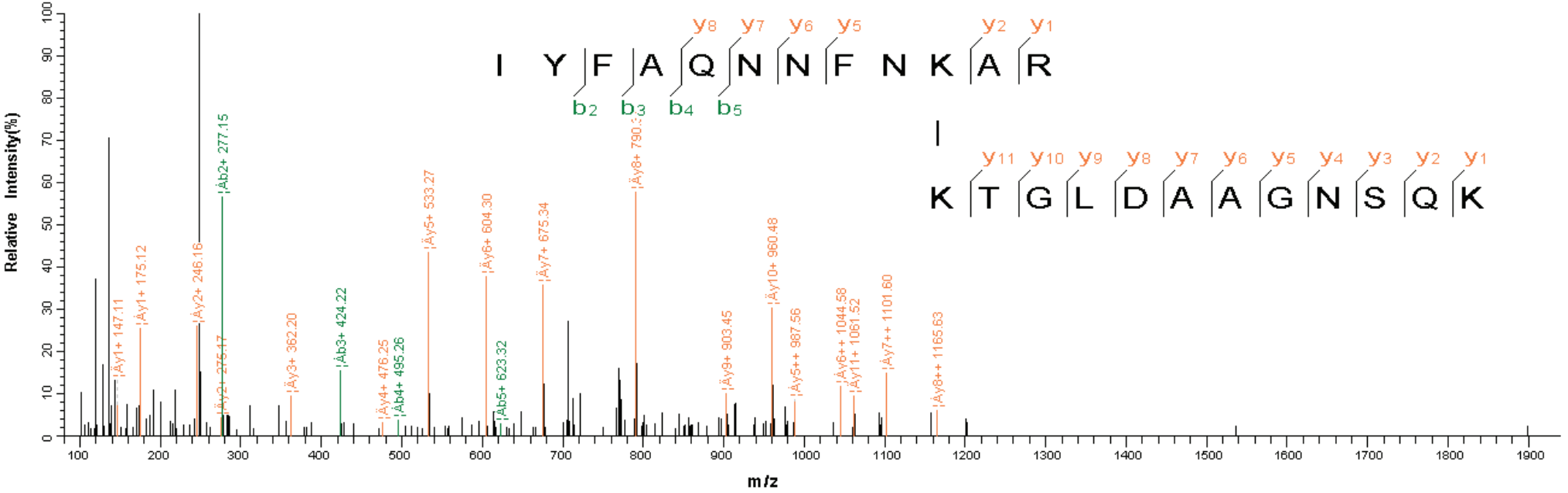


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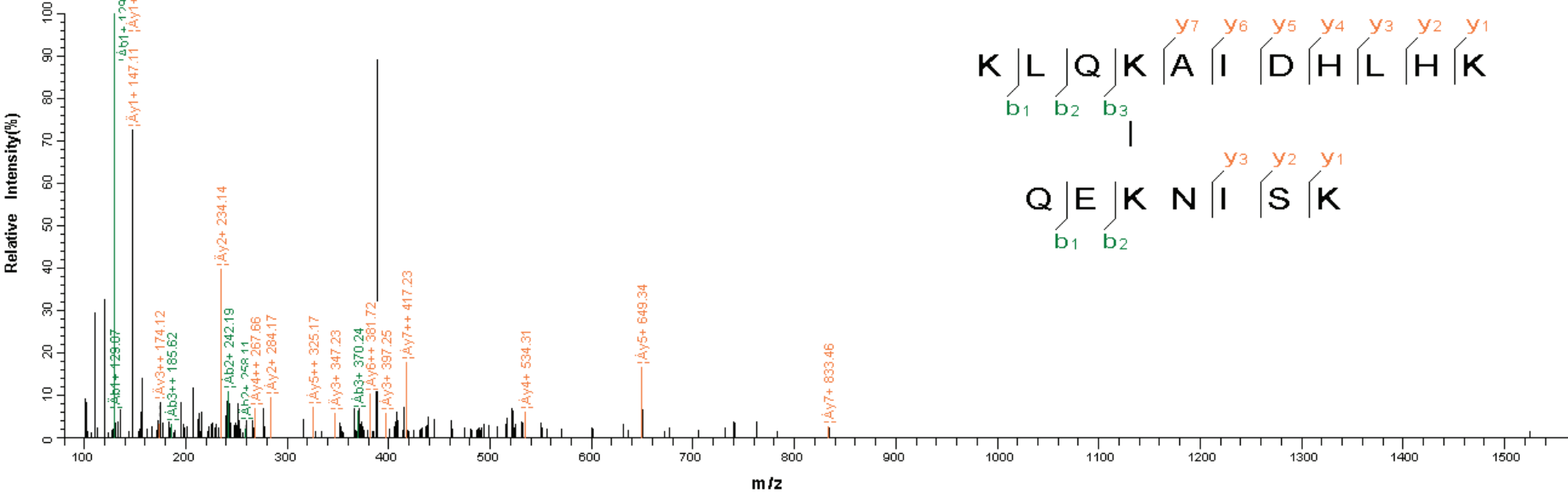


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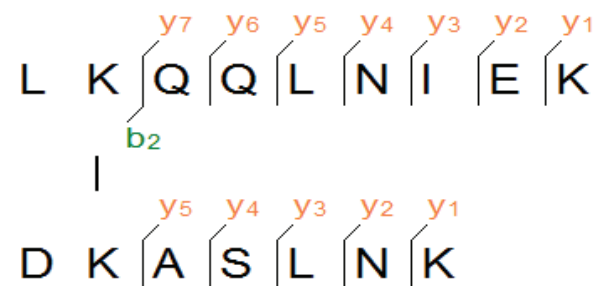
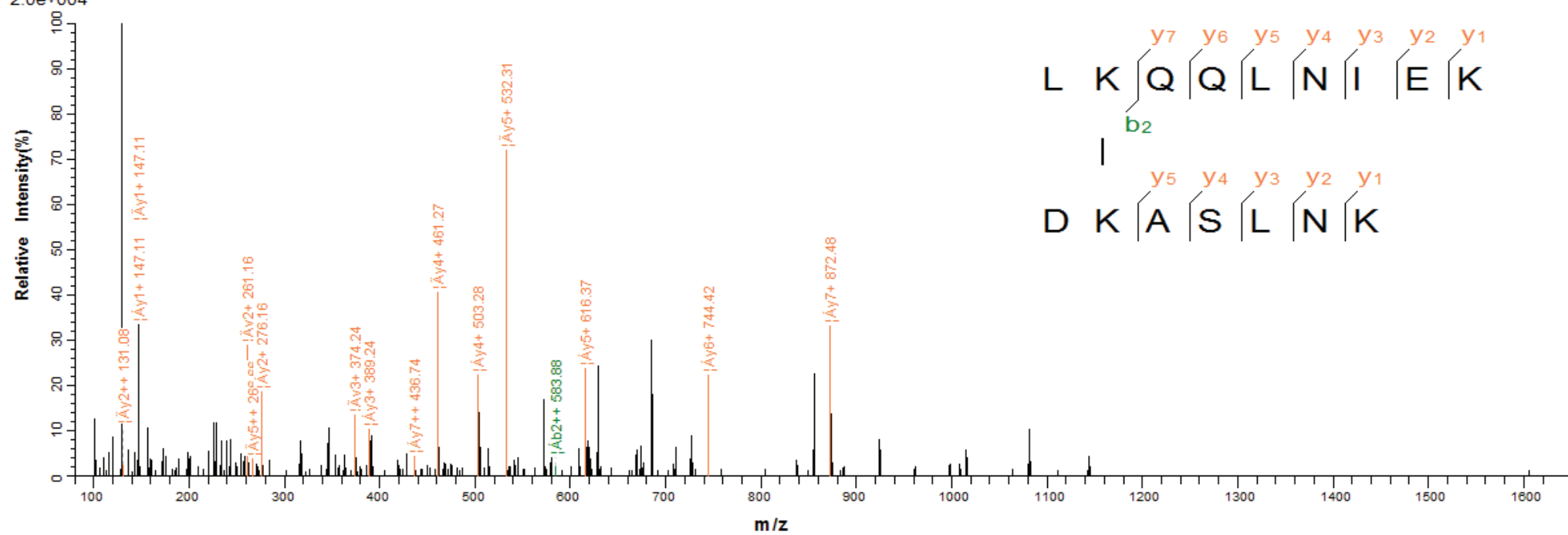
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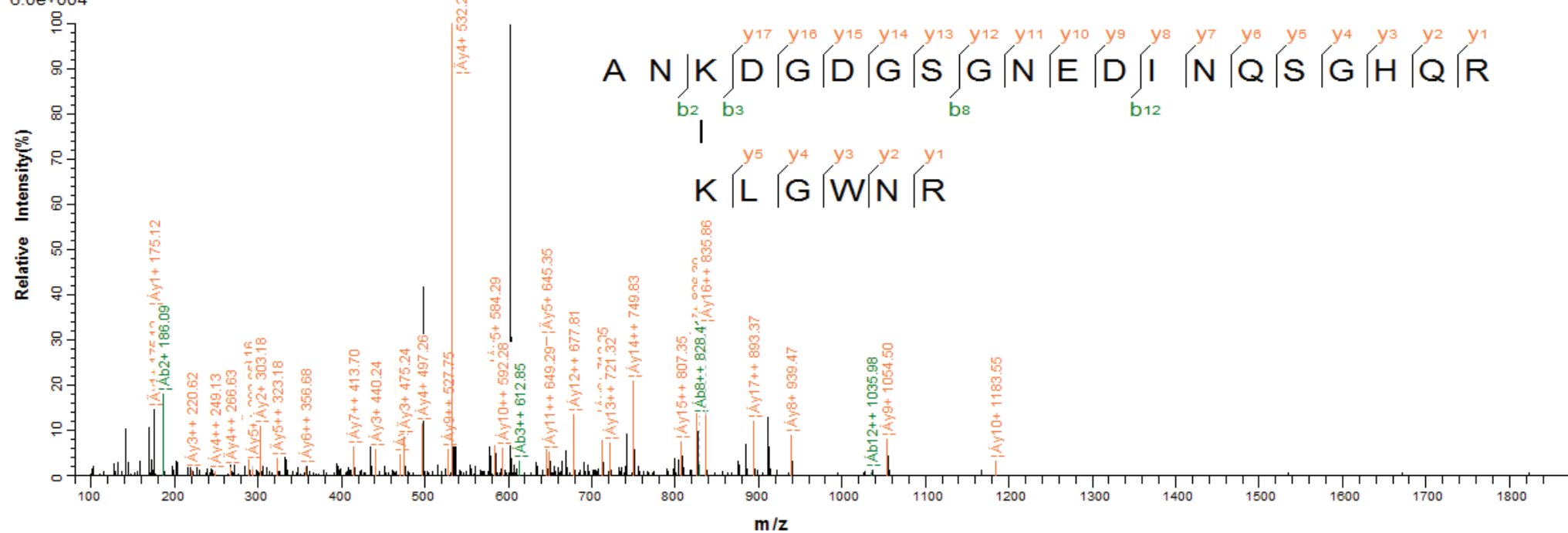
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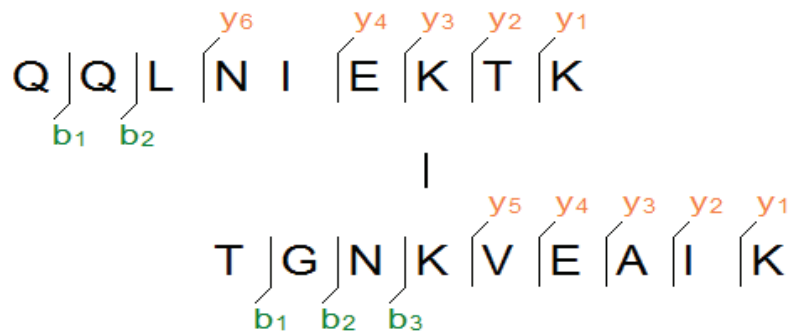
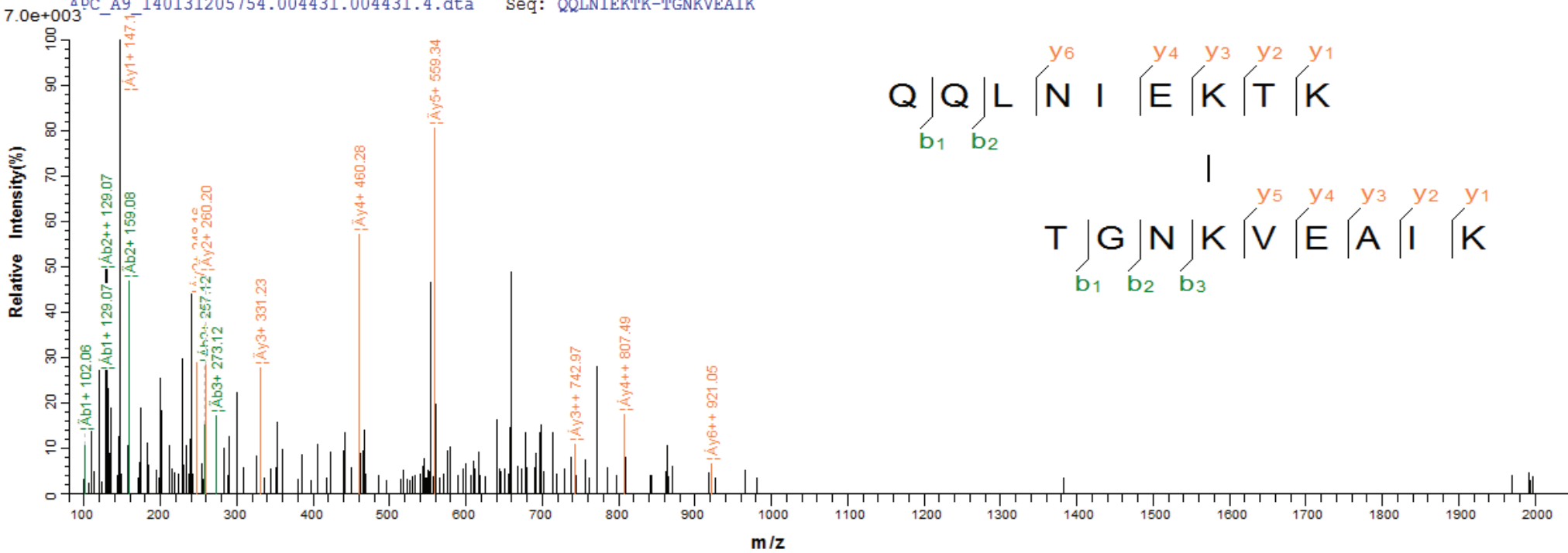
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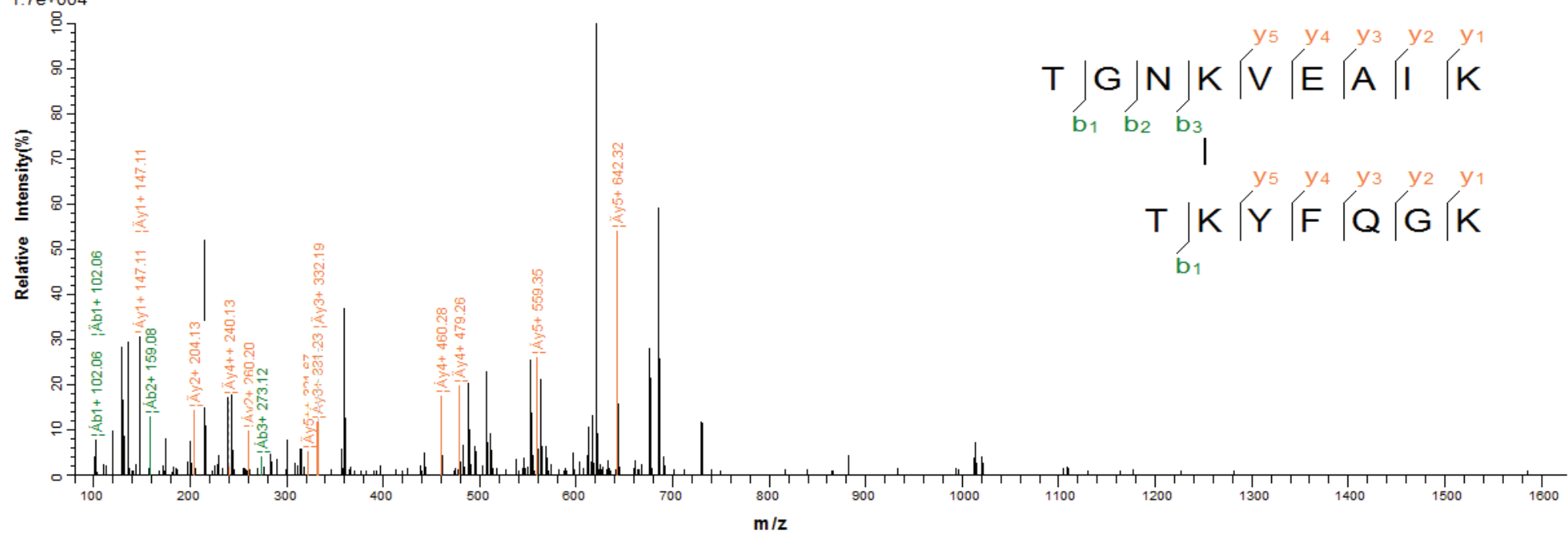
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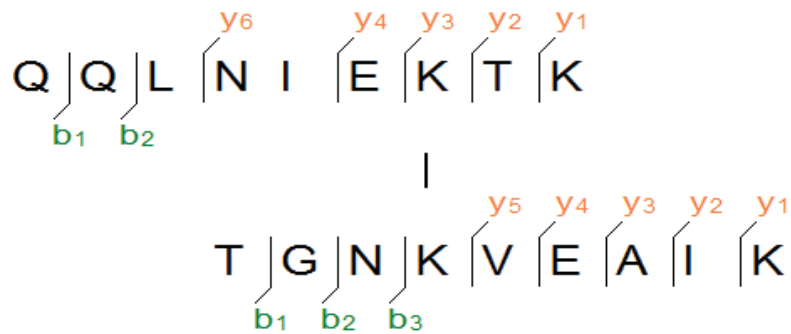
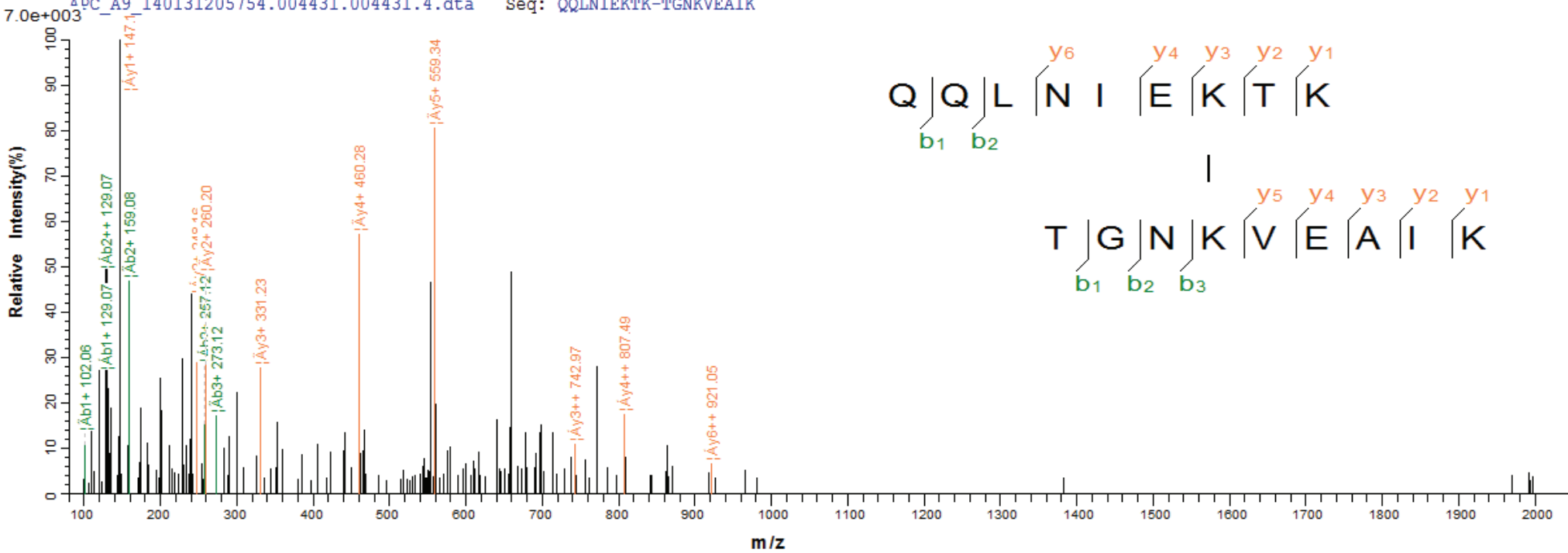
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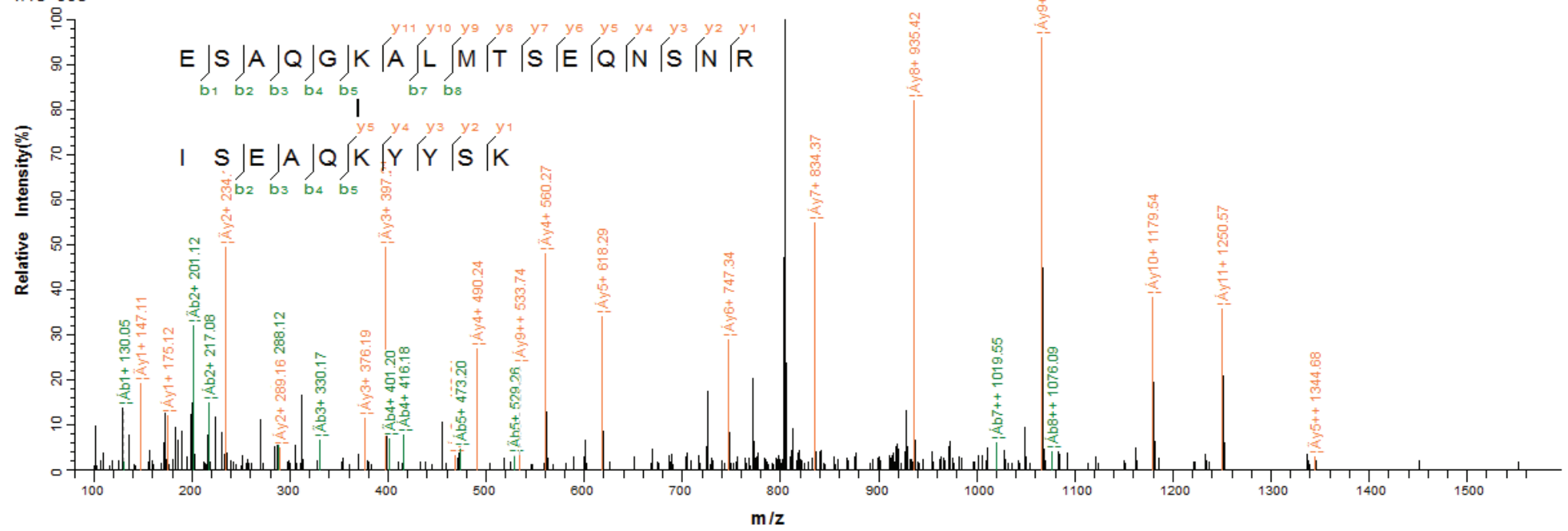
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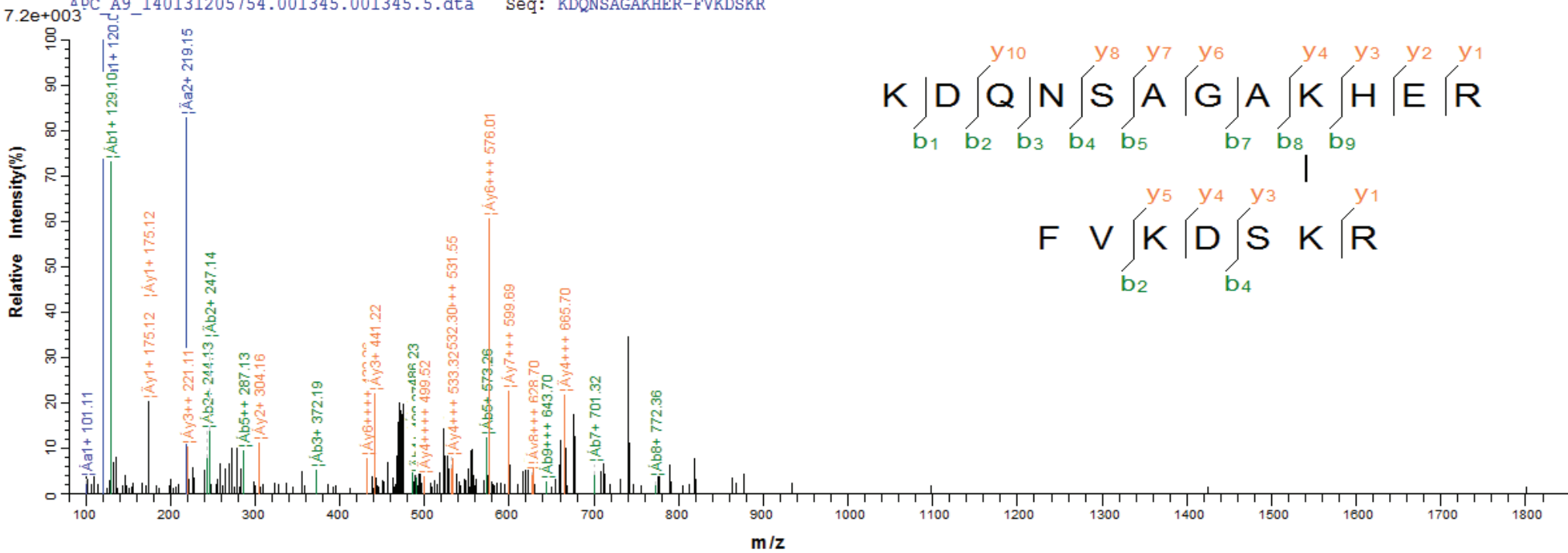
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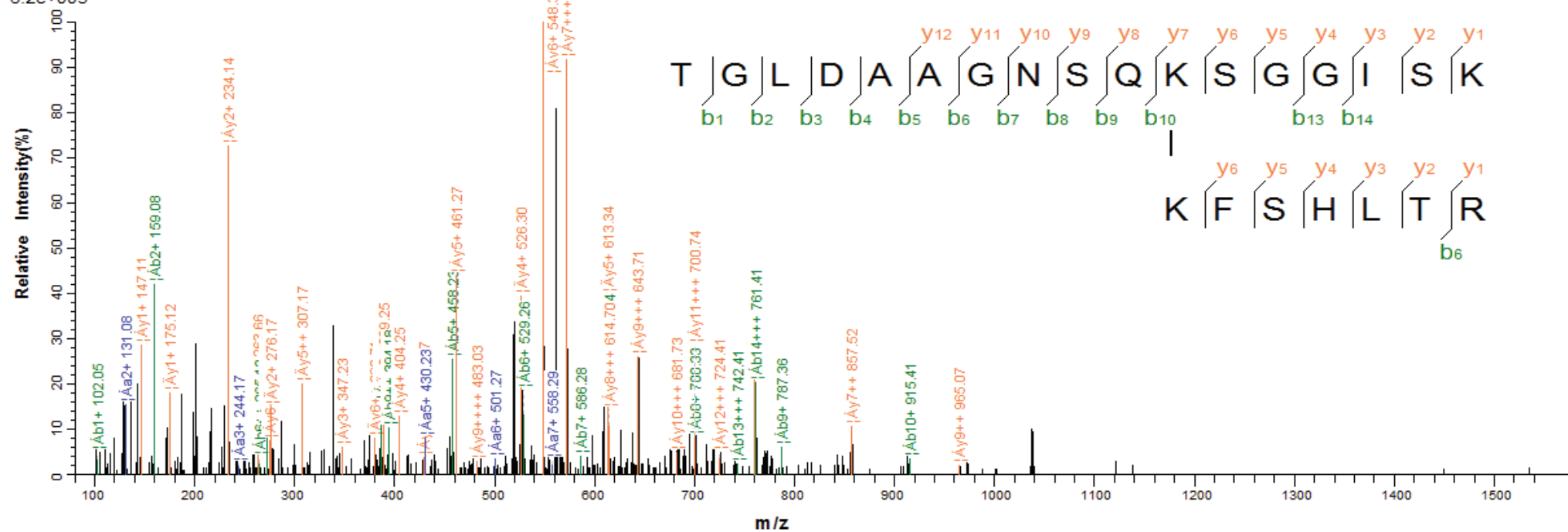


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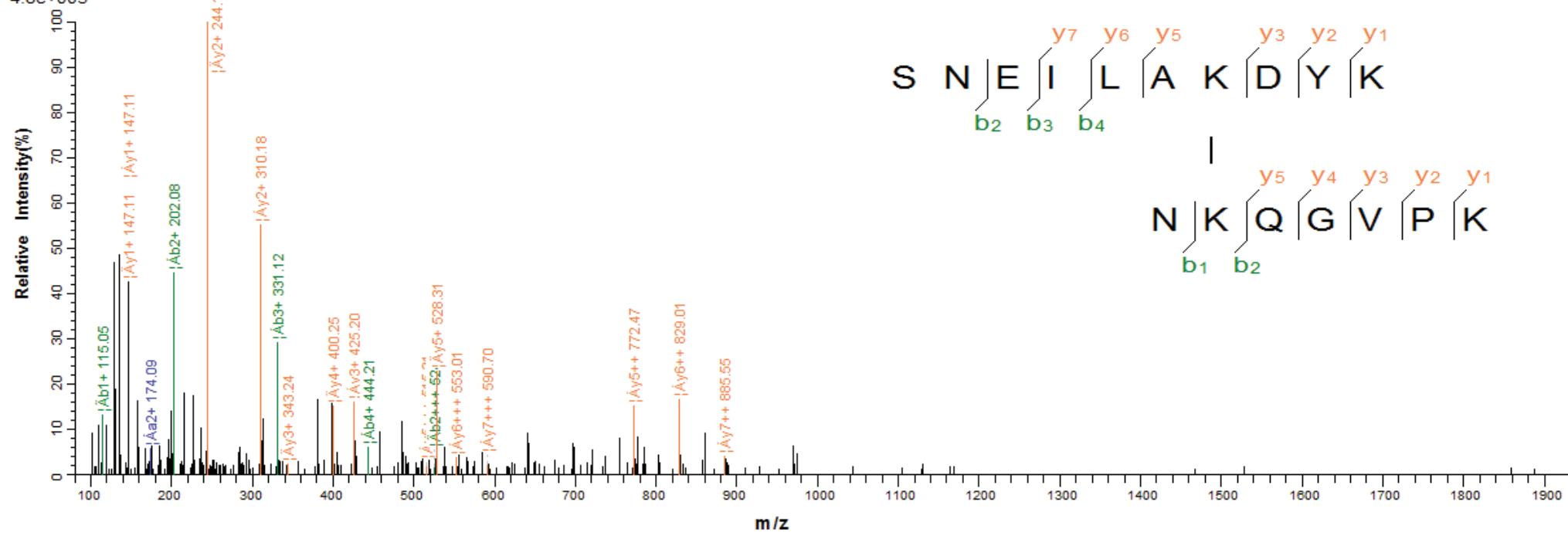


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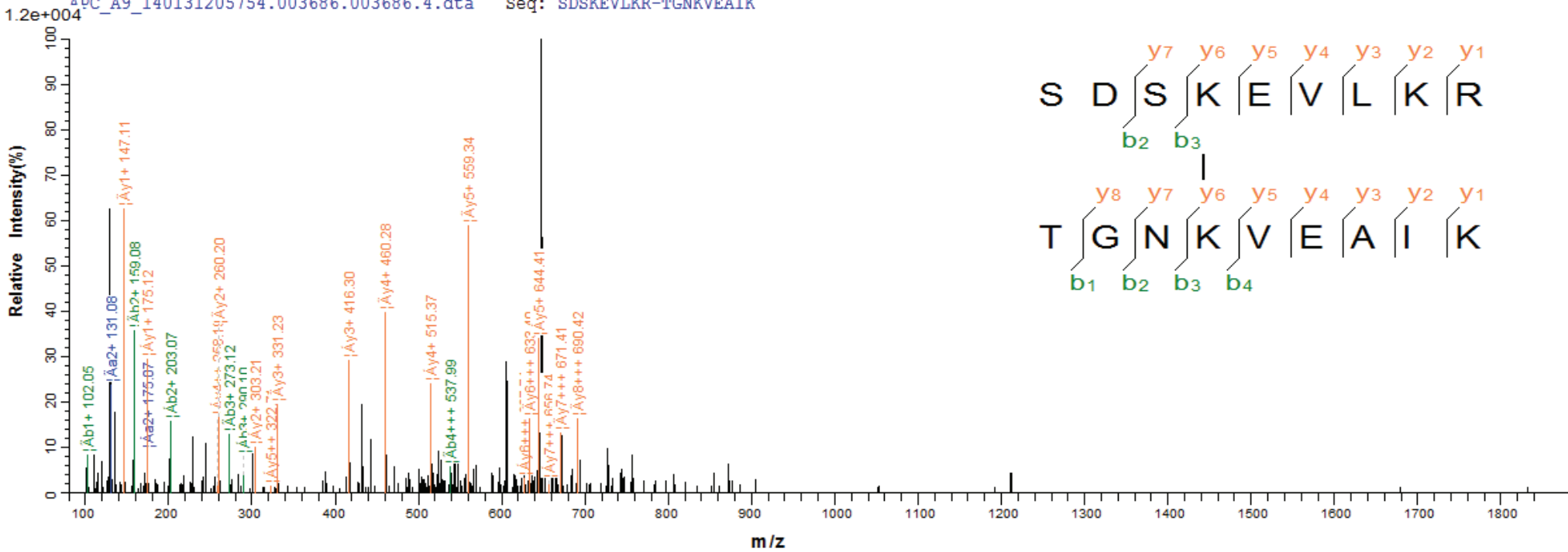
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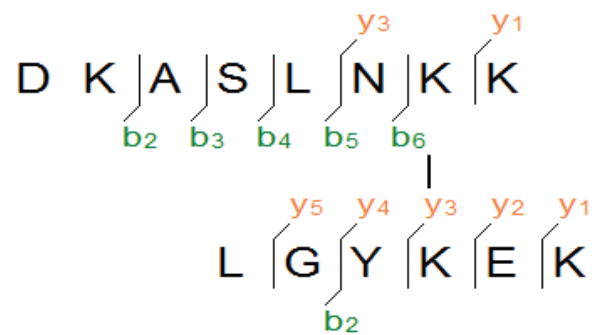
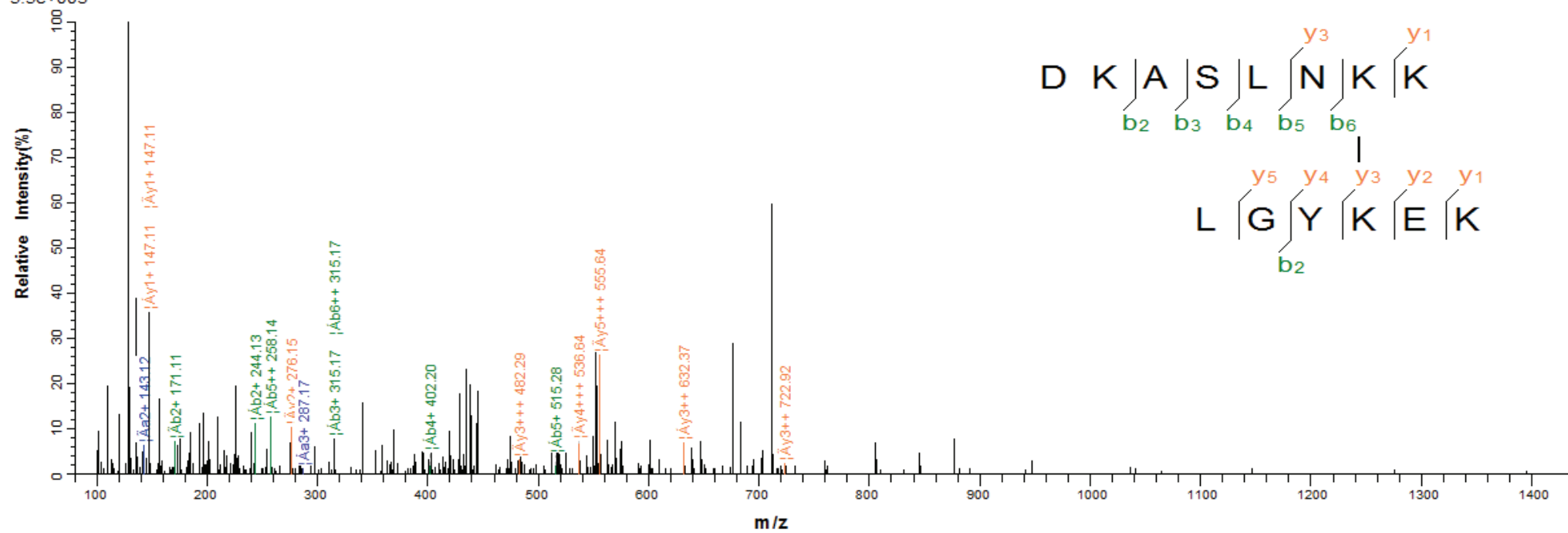
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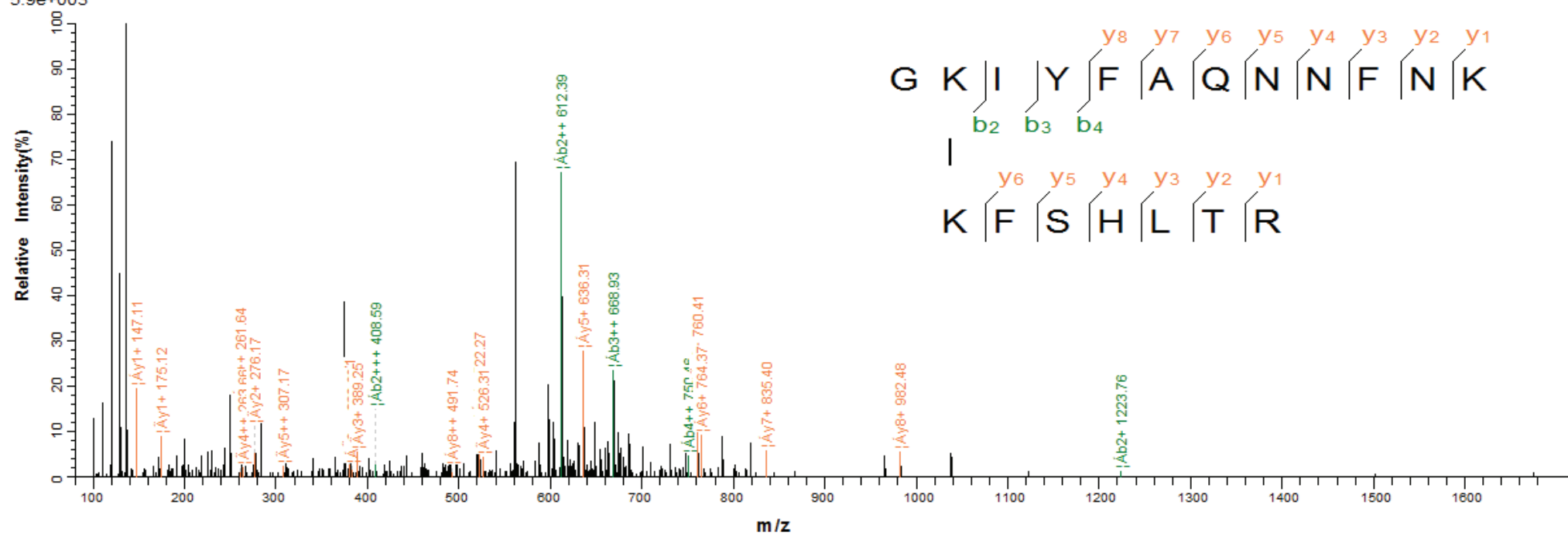
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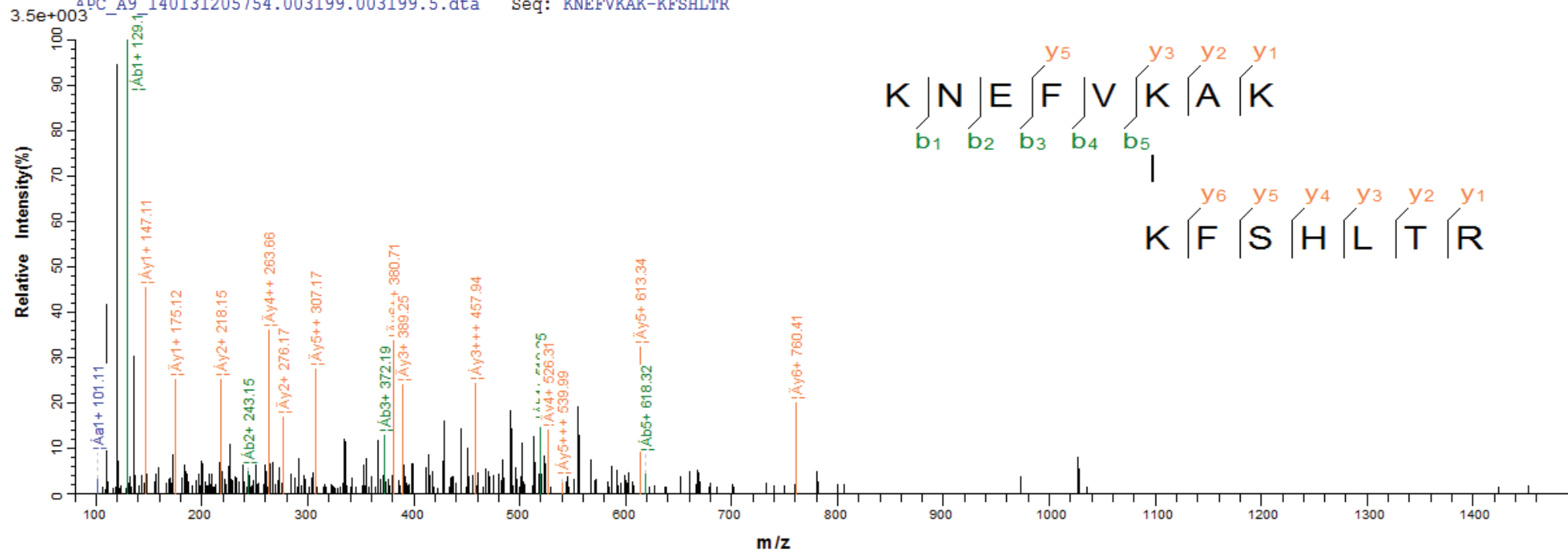
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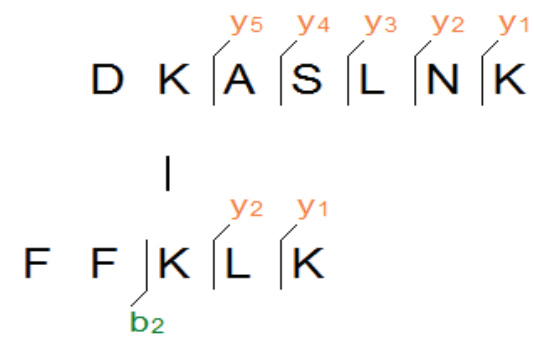
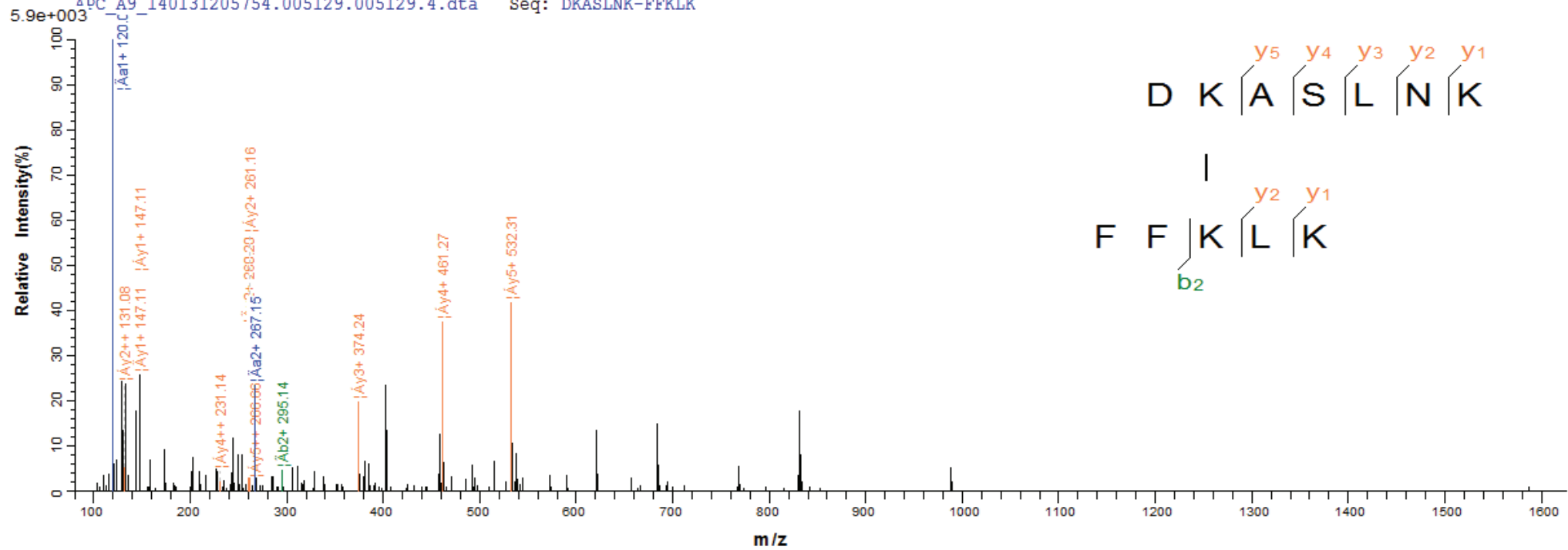
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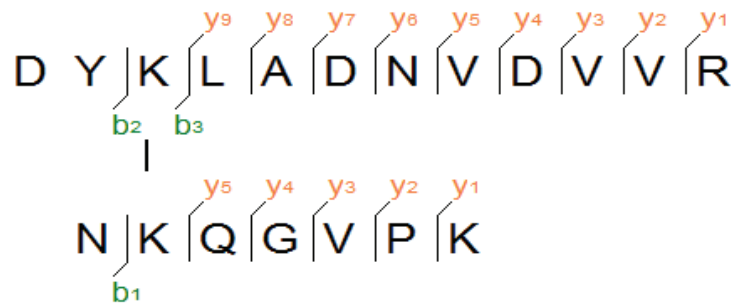
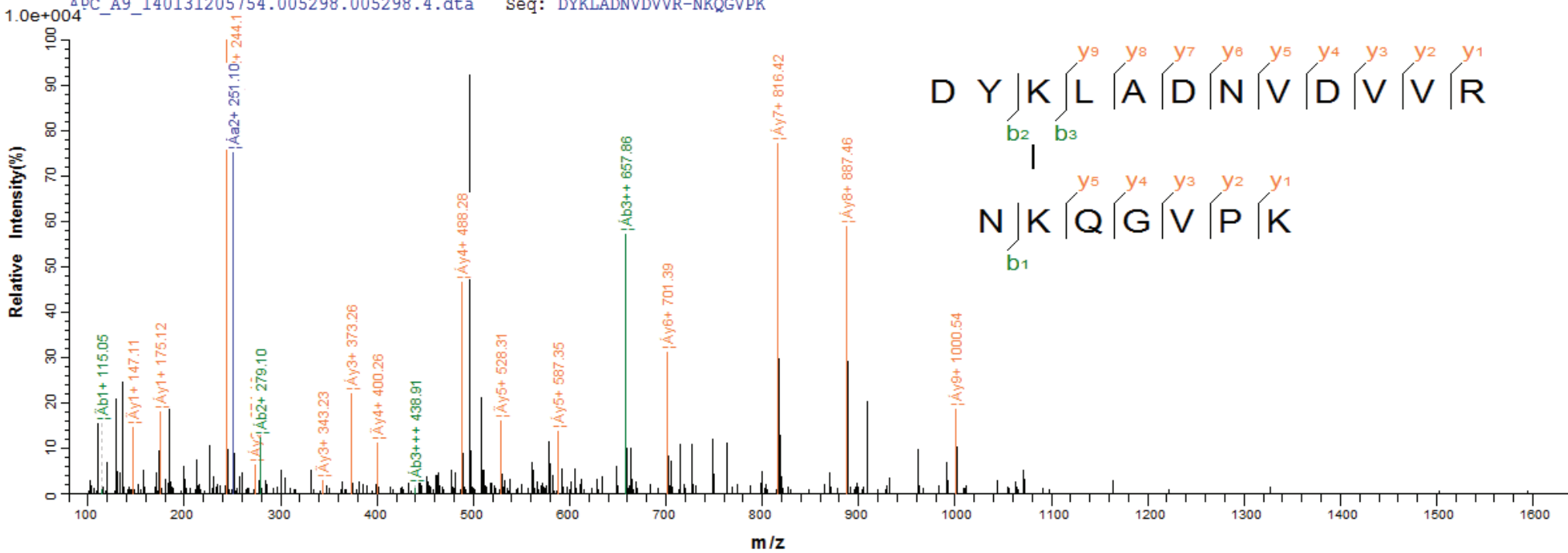
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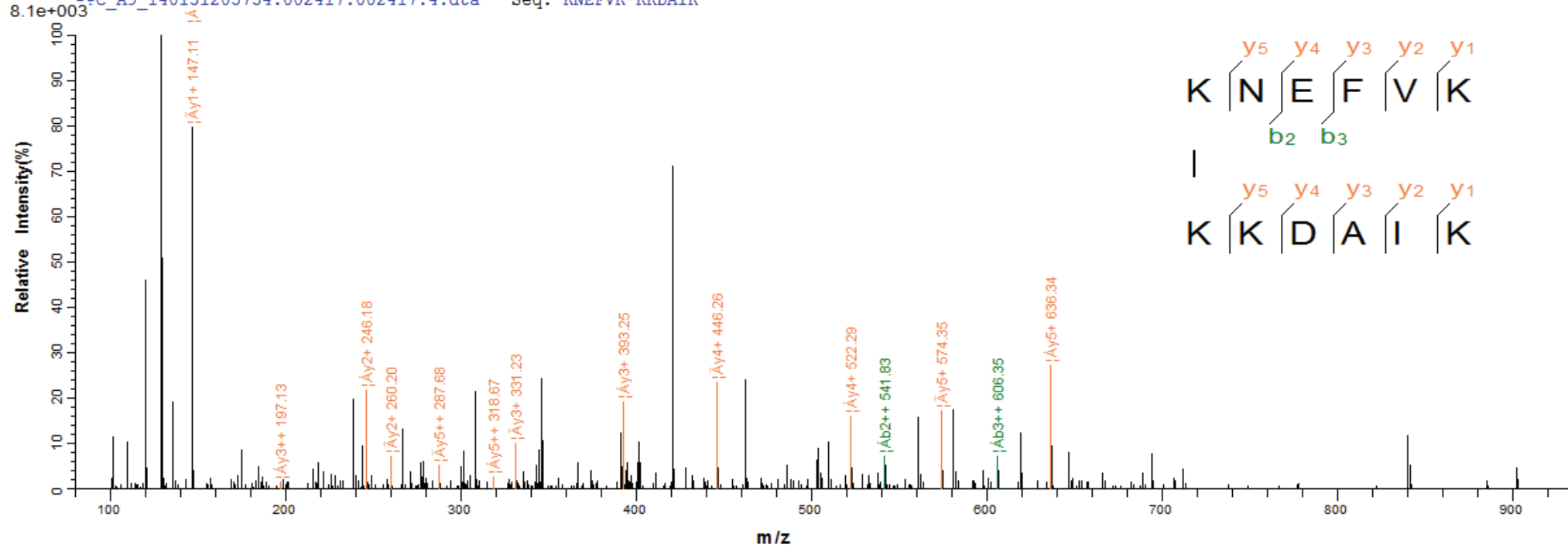
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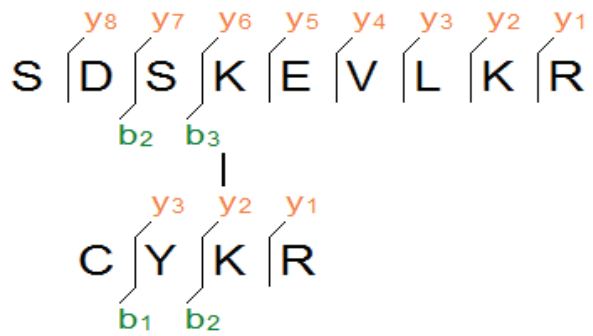
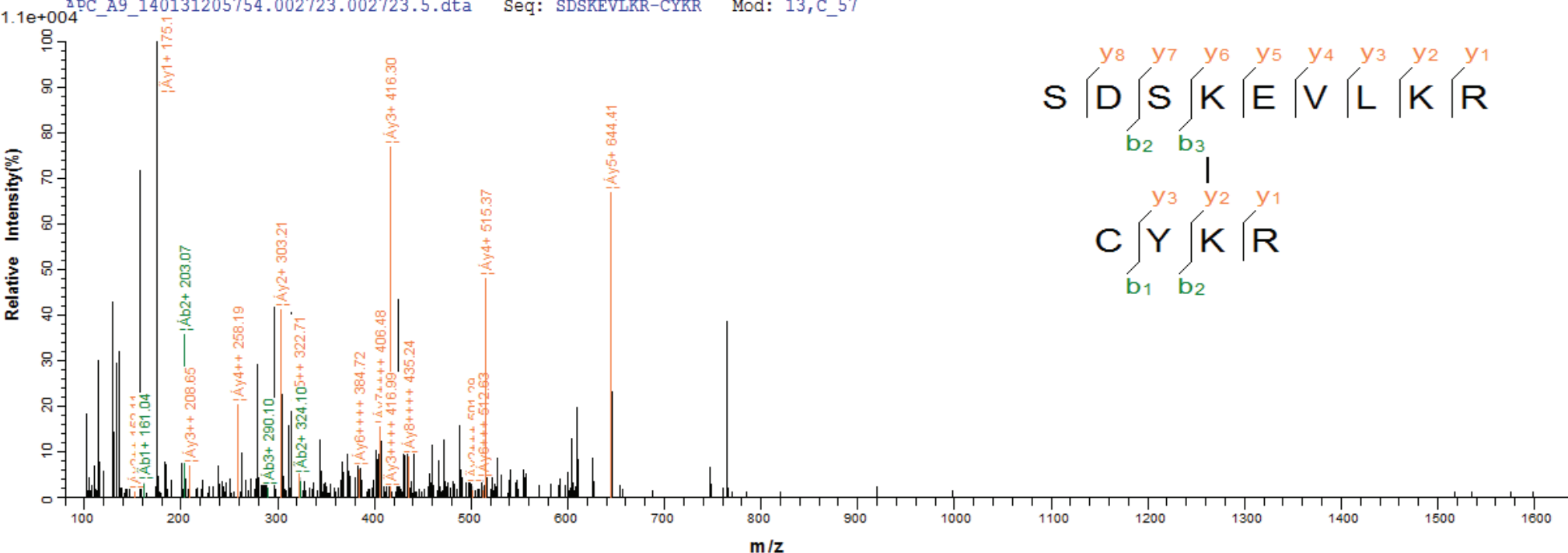
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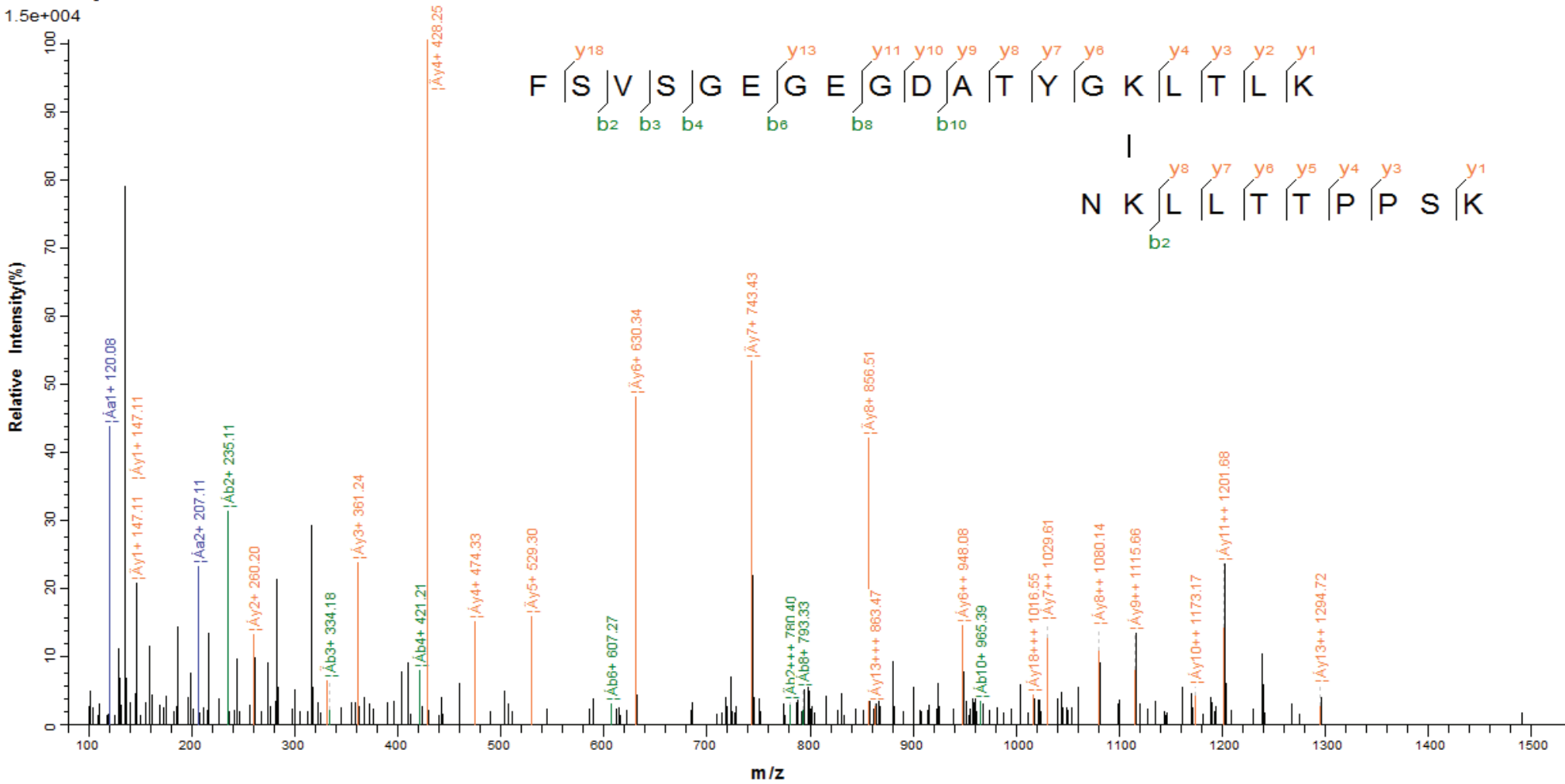
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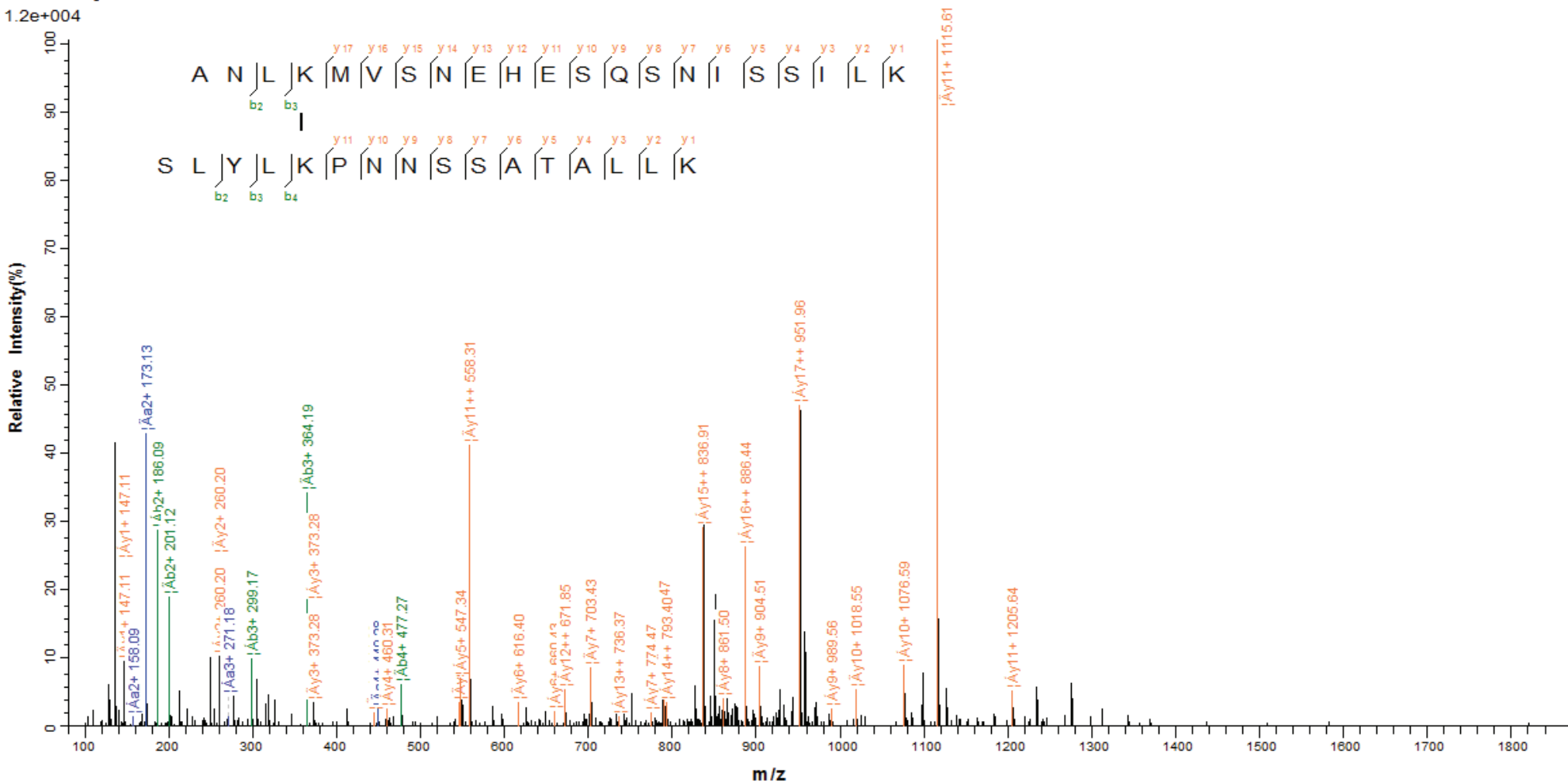
APC_A9_140131205754.002723.002723.5.dta Seq: SDSKEVLKR-CYKR Mod: 13,C_57

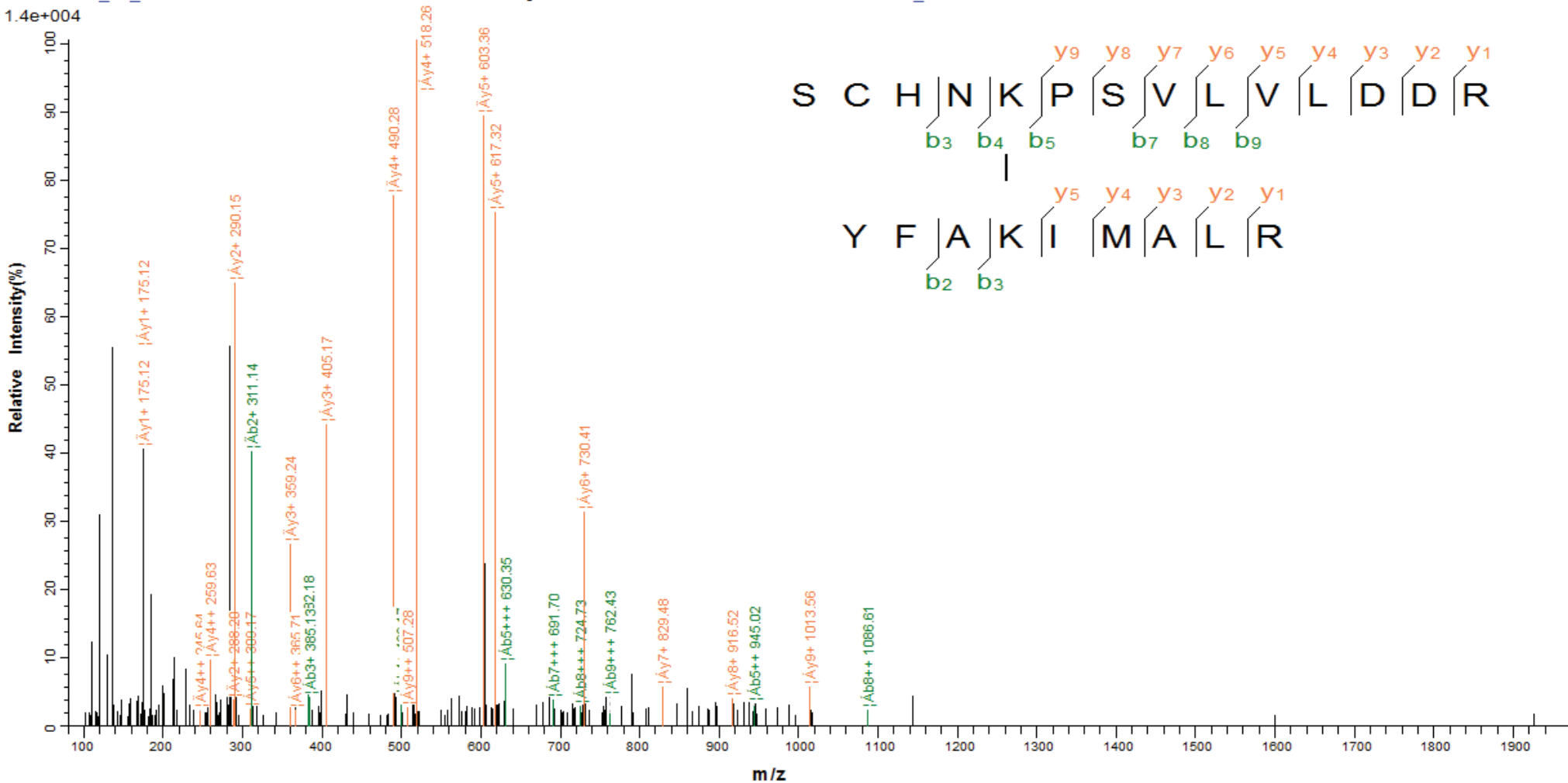


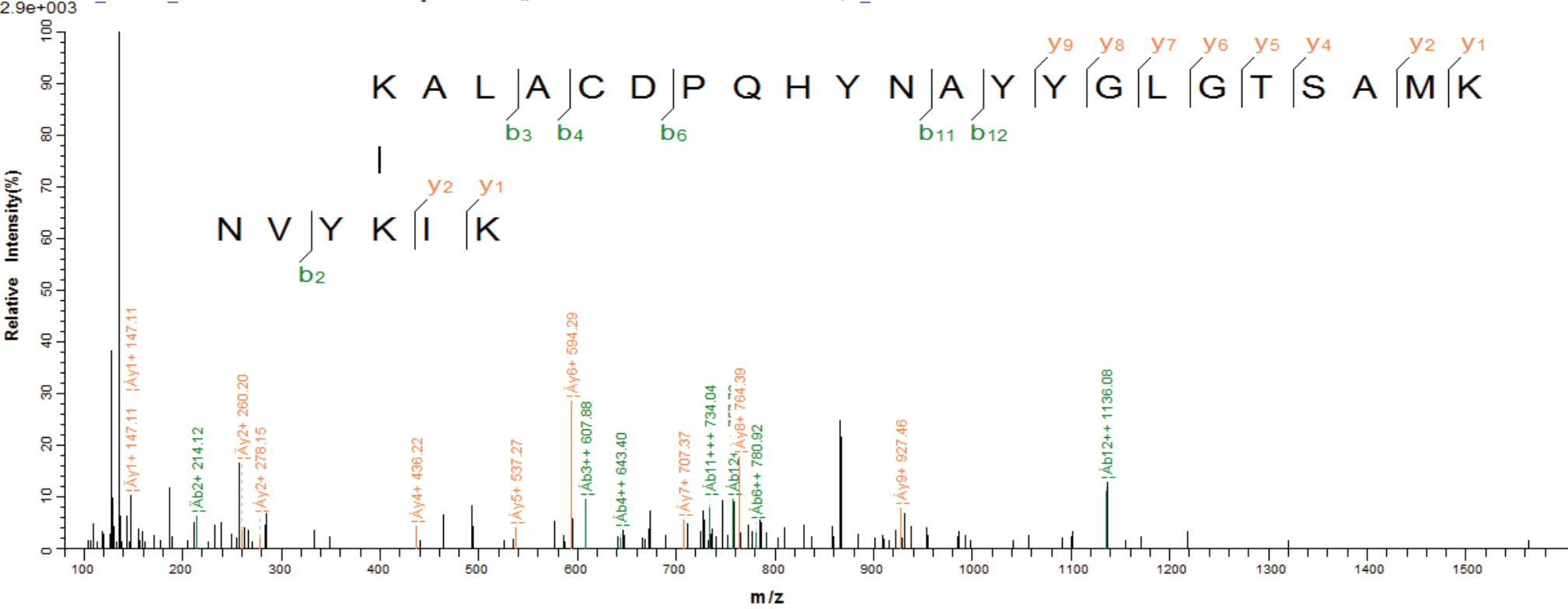
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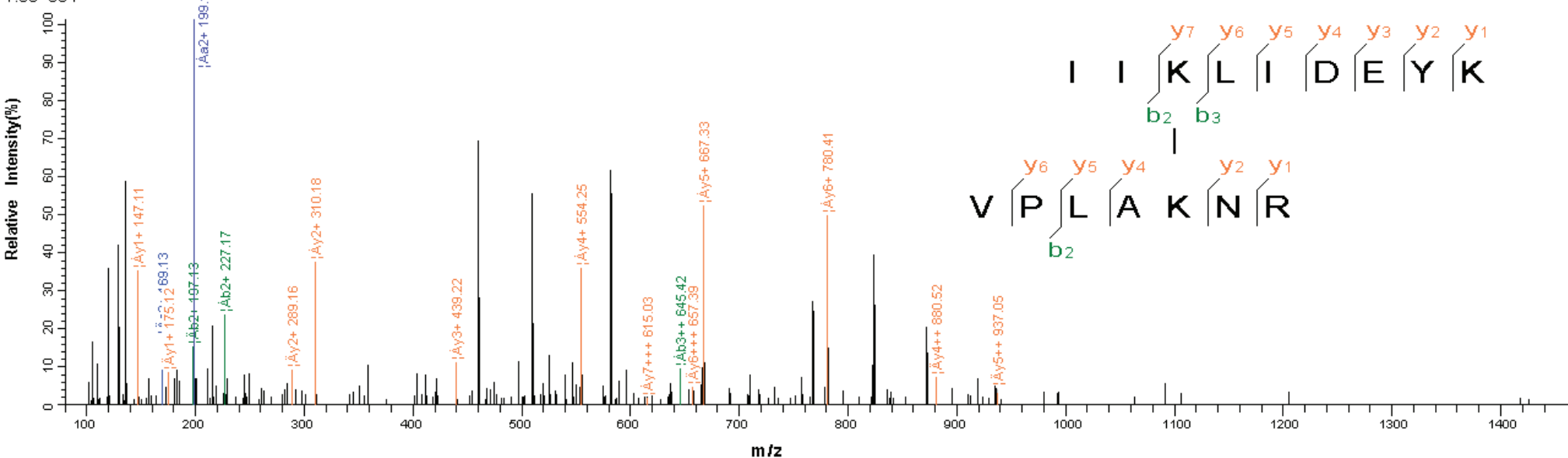
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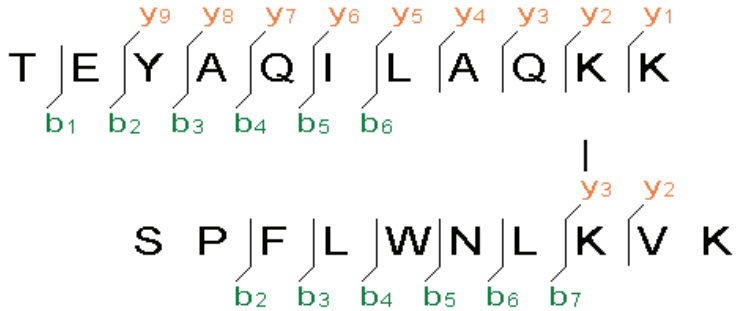
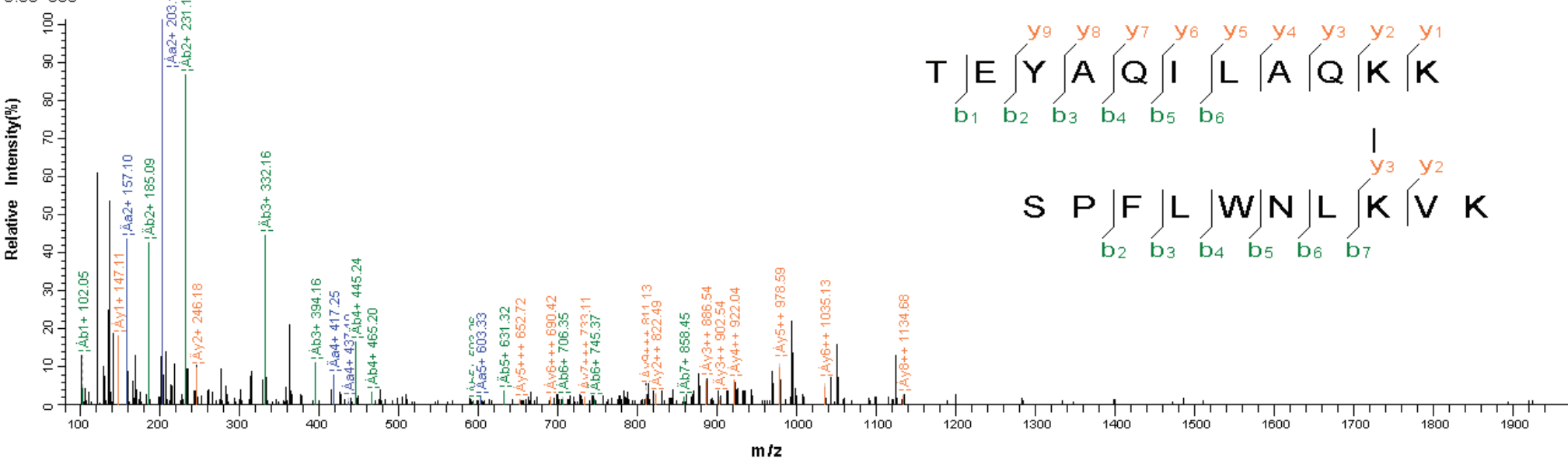


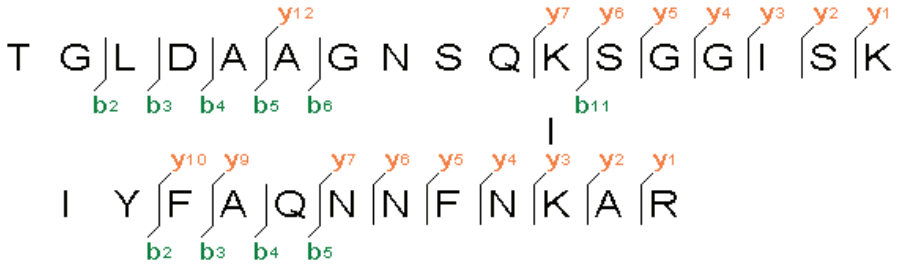
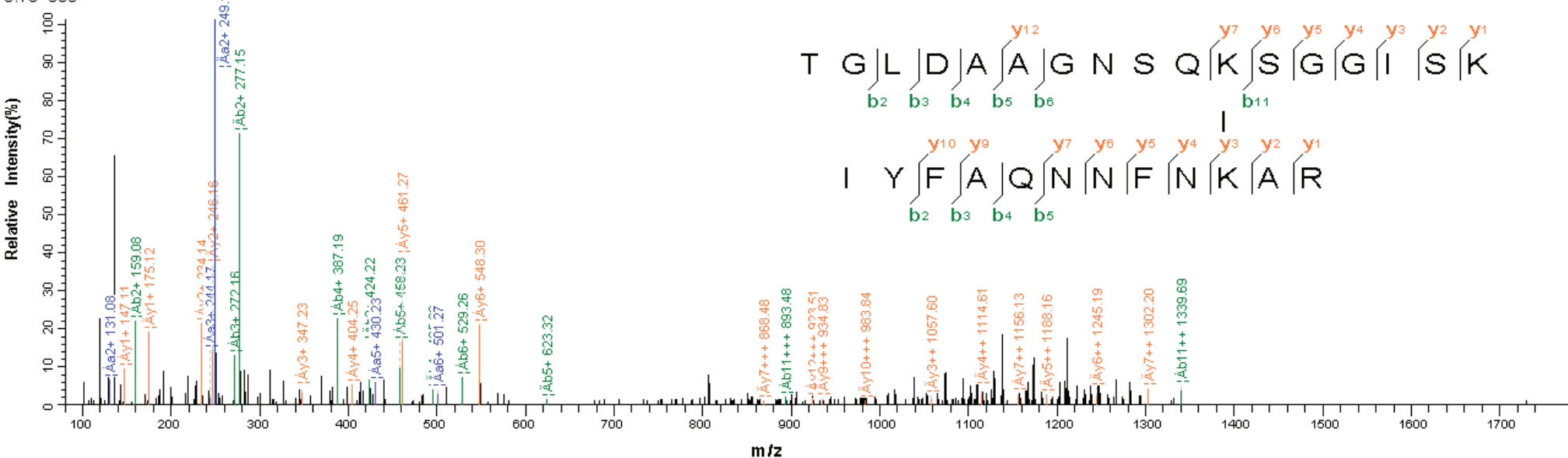


1.6e+004 PC_A8_140201121226.004228.004228.4.dta Seq: IIKLIDEYK-VPLAKNR Charge: 4

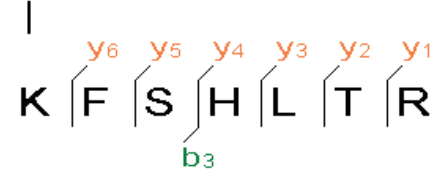
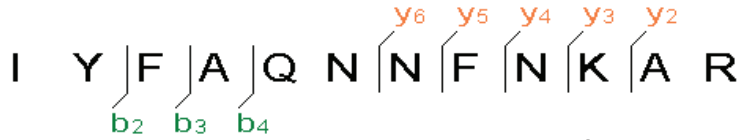
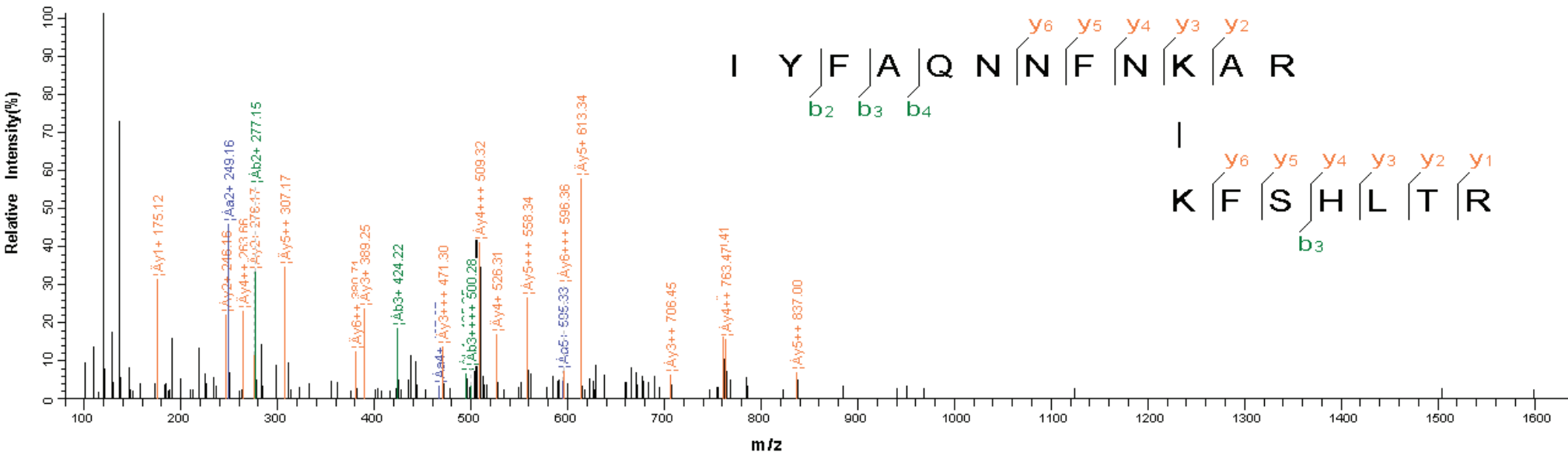


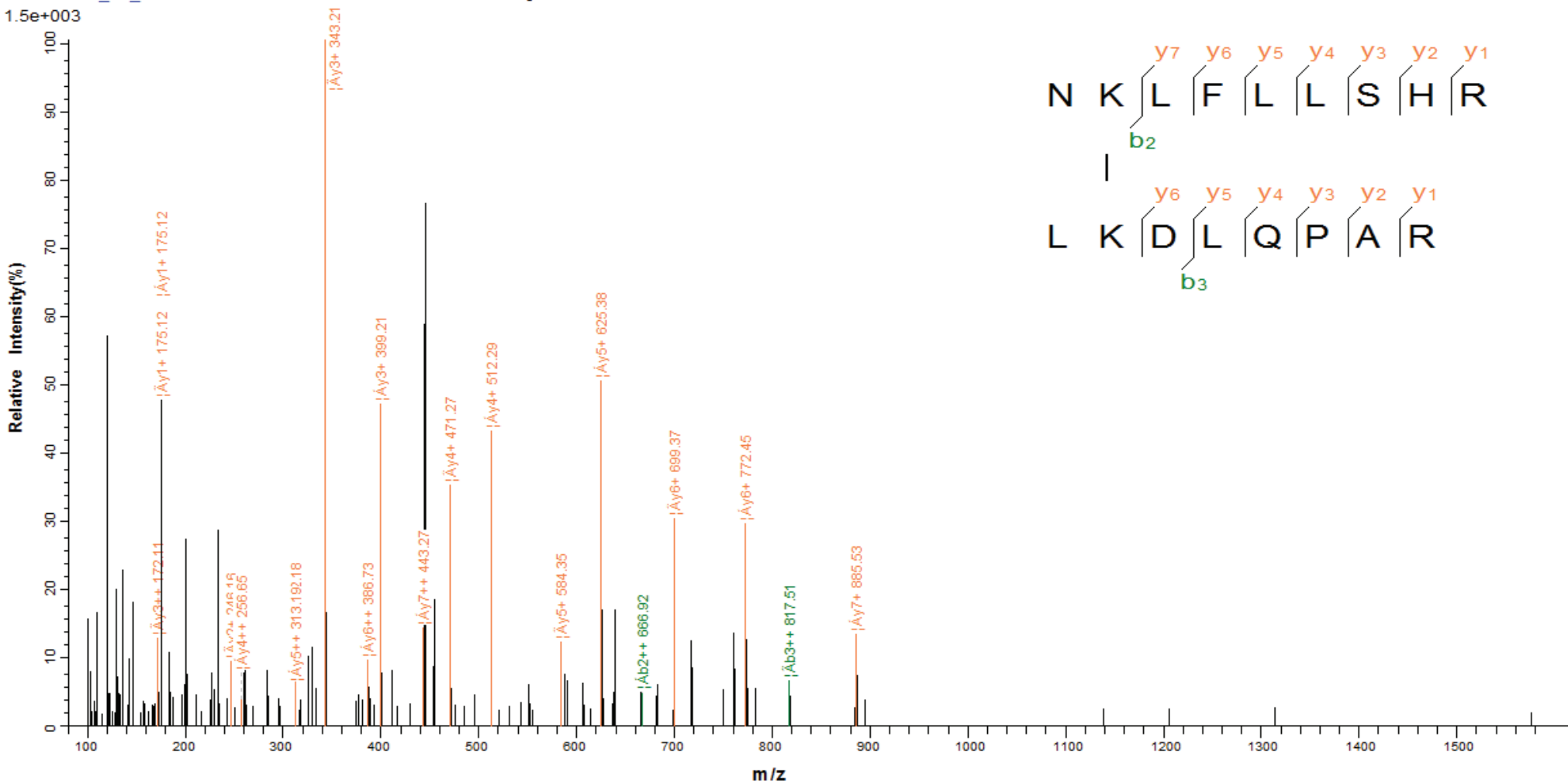
6.6e+003 PC_A8_140201121226.006843.006843.4.dta Seq: TEYAQLAQKK-SPFLWNLKVK Charge: 4



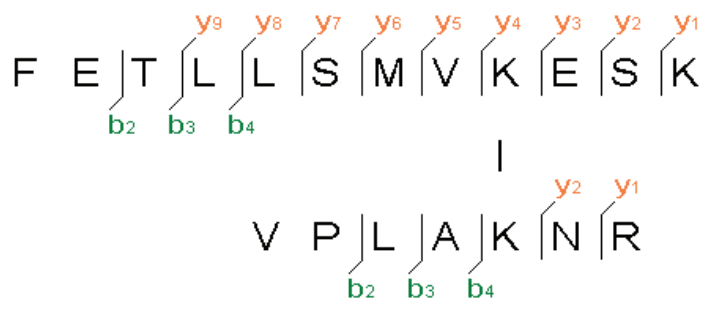
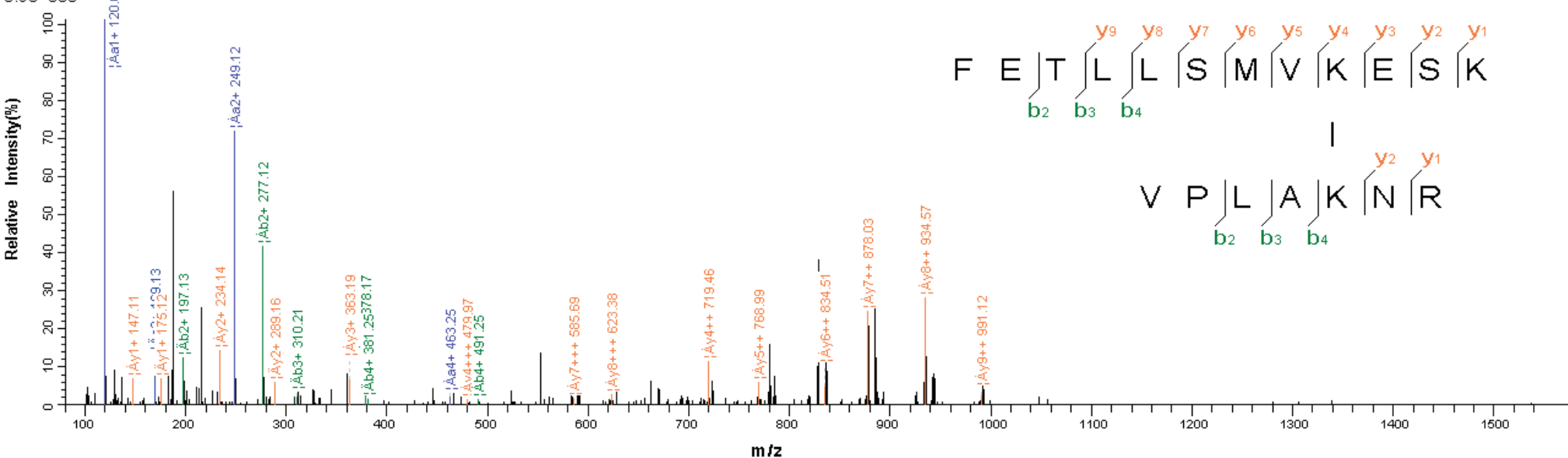


3.8e+003 PC_A8_140201121226.003490.003490.5.dta Seq: IYFAQNMFNFKAR-KPSHLTR Charge: 5





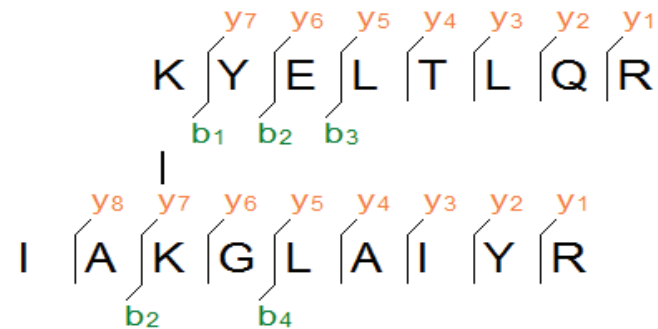
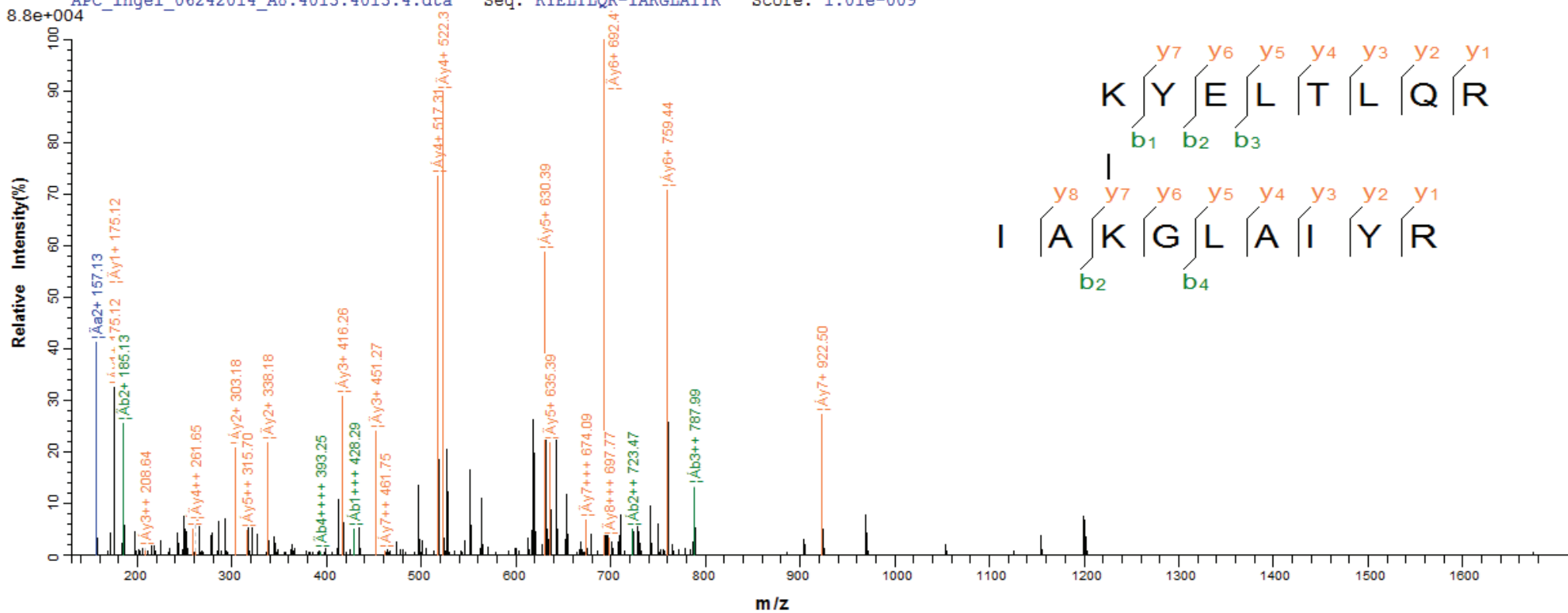
6.9e+003 PC_A8_140201121226.006331.006331.4.dta Seq: FETLLSMVKESK-VPLAKNR Charge: 4



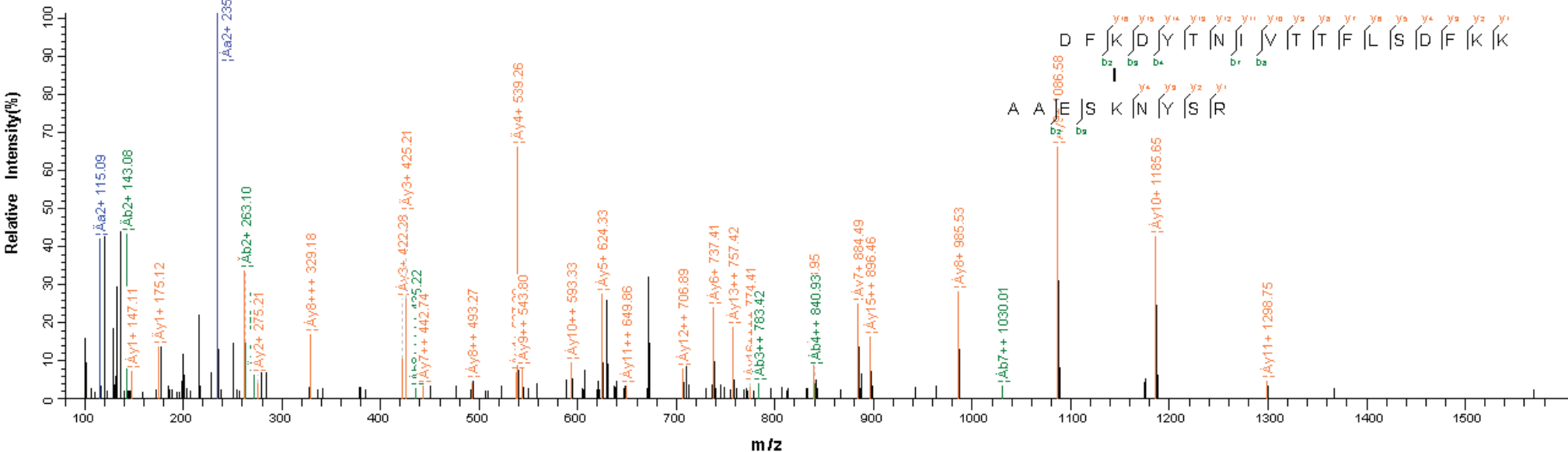
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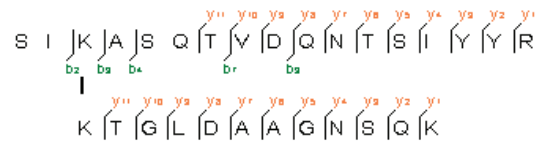
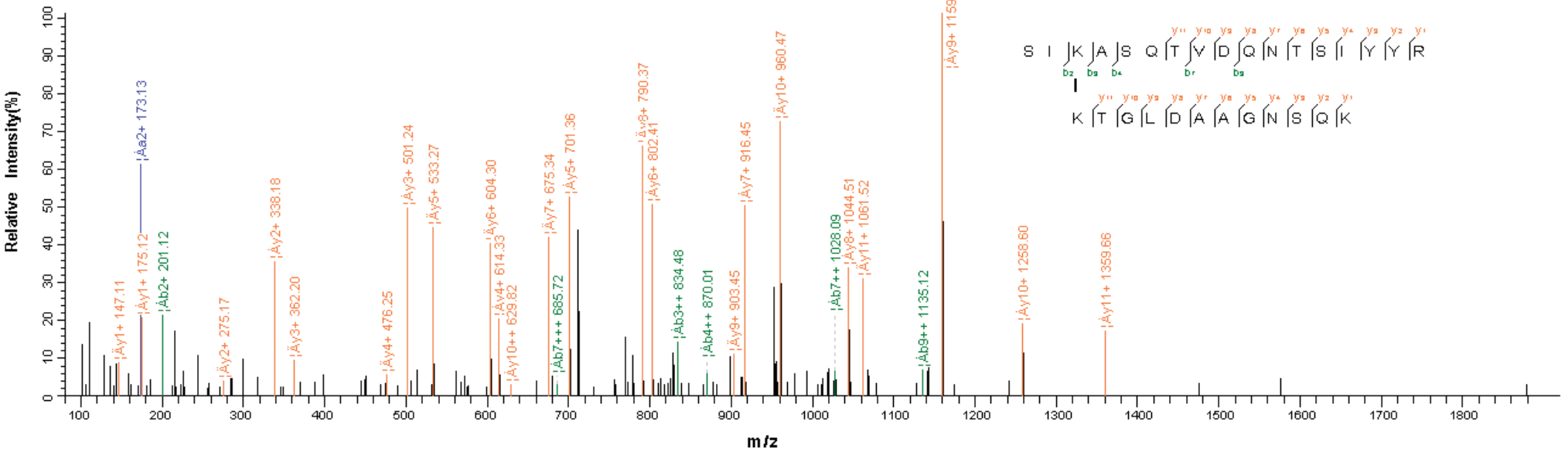
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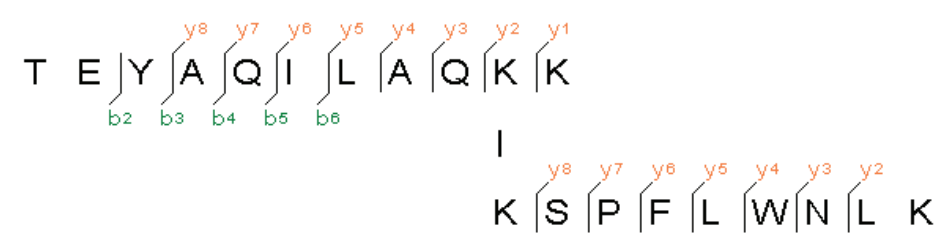
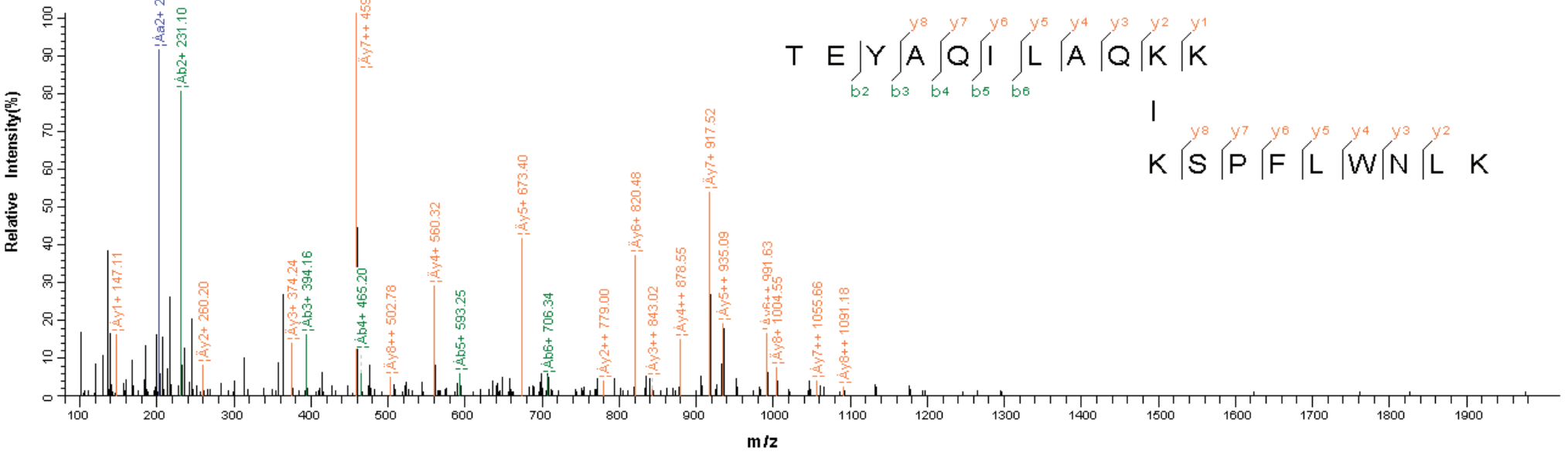


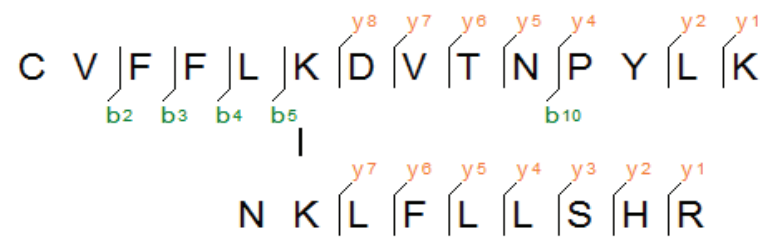
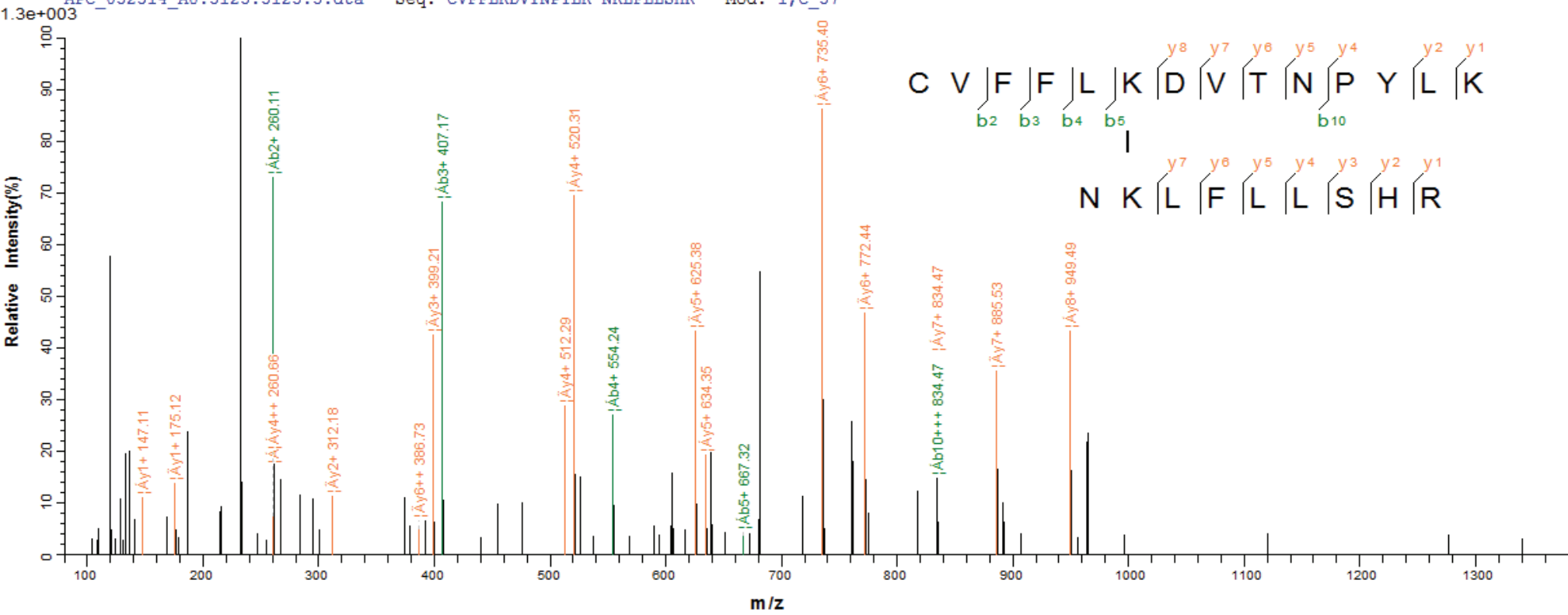
1.6e+003 PC_A8_140201121226.008860.008860.5.dta Seq: DFKDYTNIVTTFLLSDFKK-AAESKNYSR Charge: 5



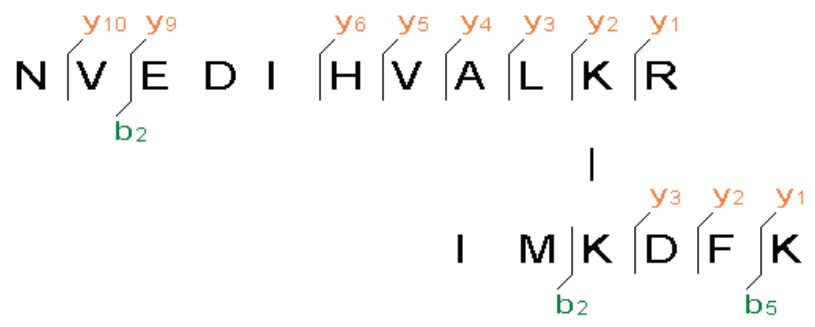
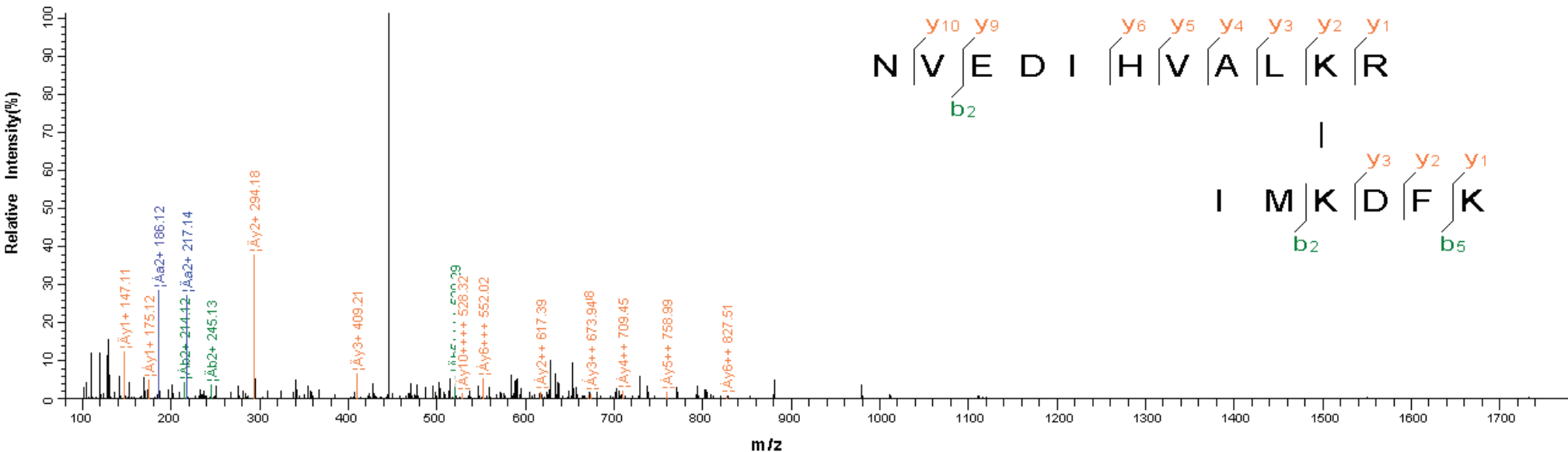


PC_A8_1402011226.006508.006508.4.dta Seq: TEYAQILAQKK-KSPFLWNLK Charge: 4

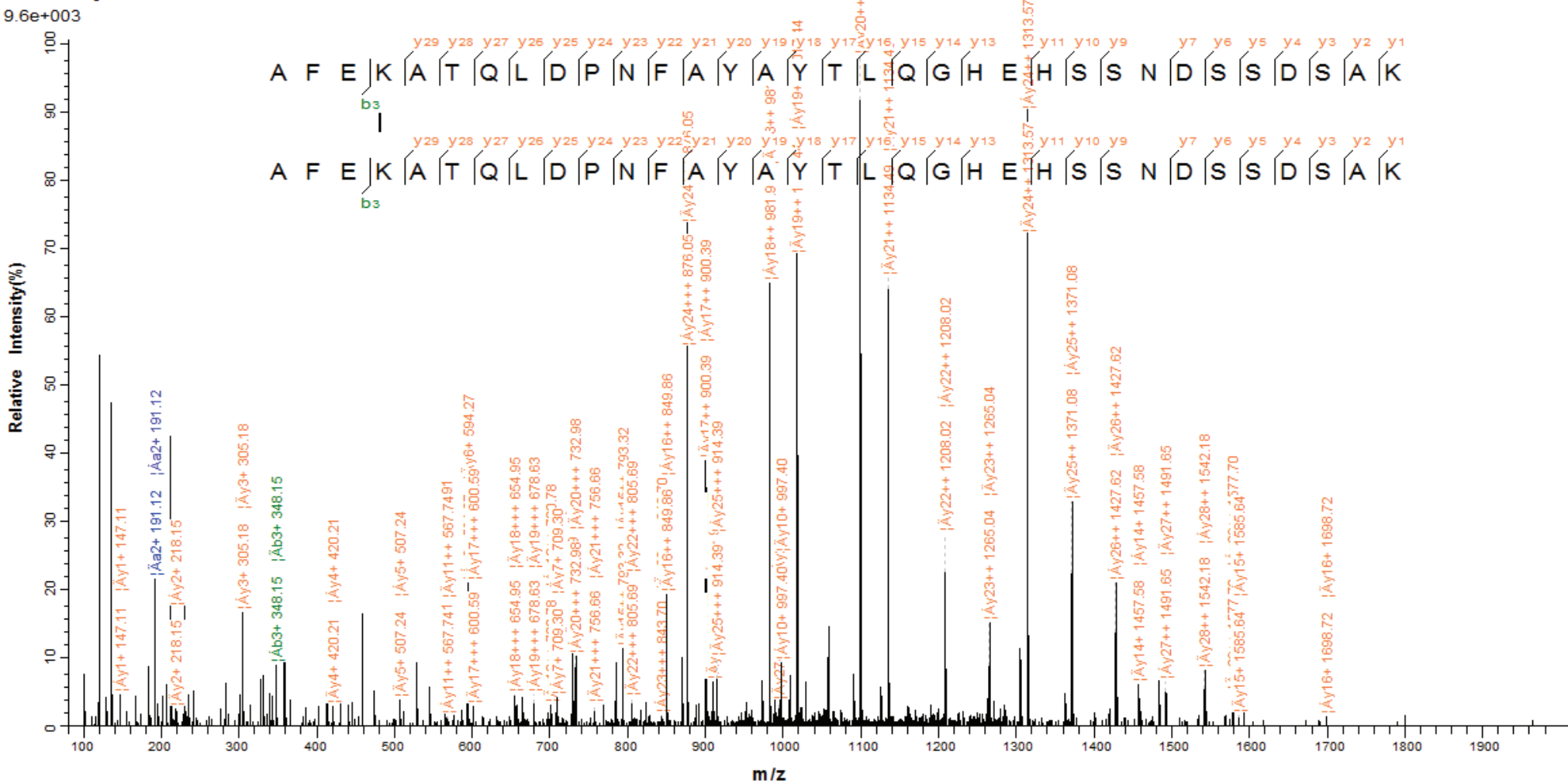


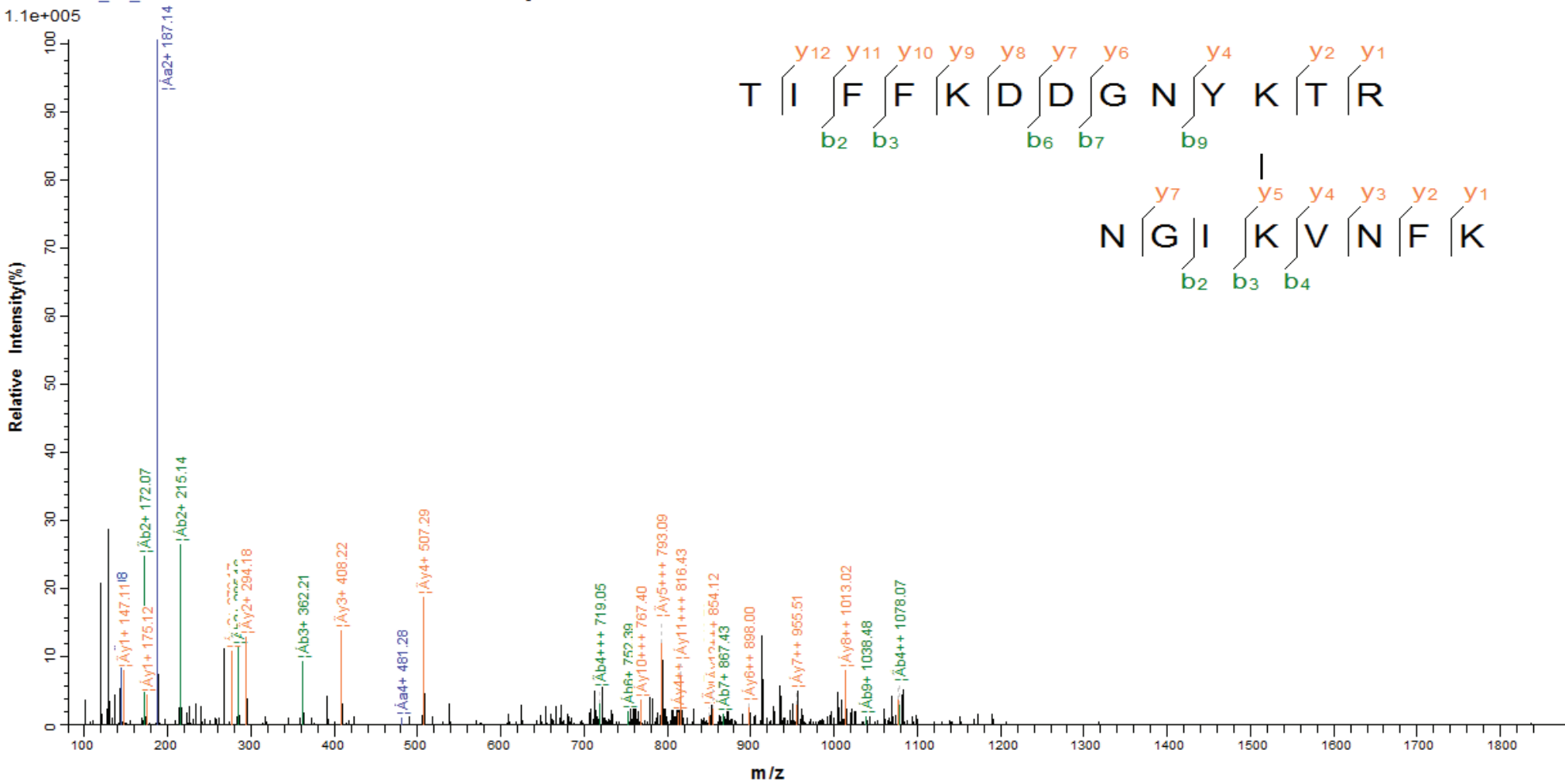


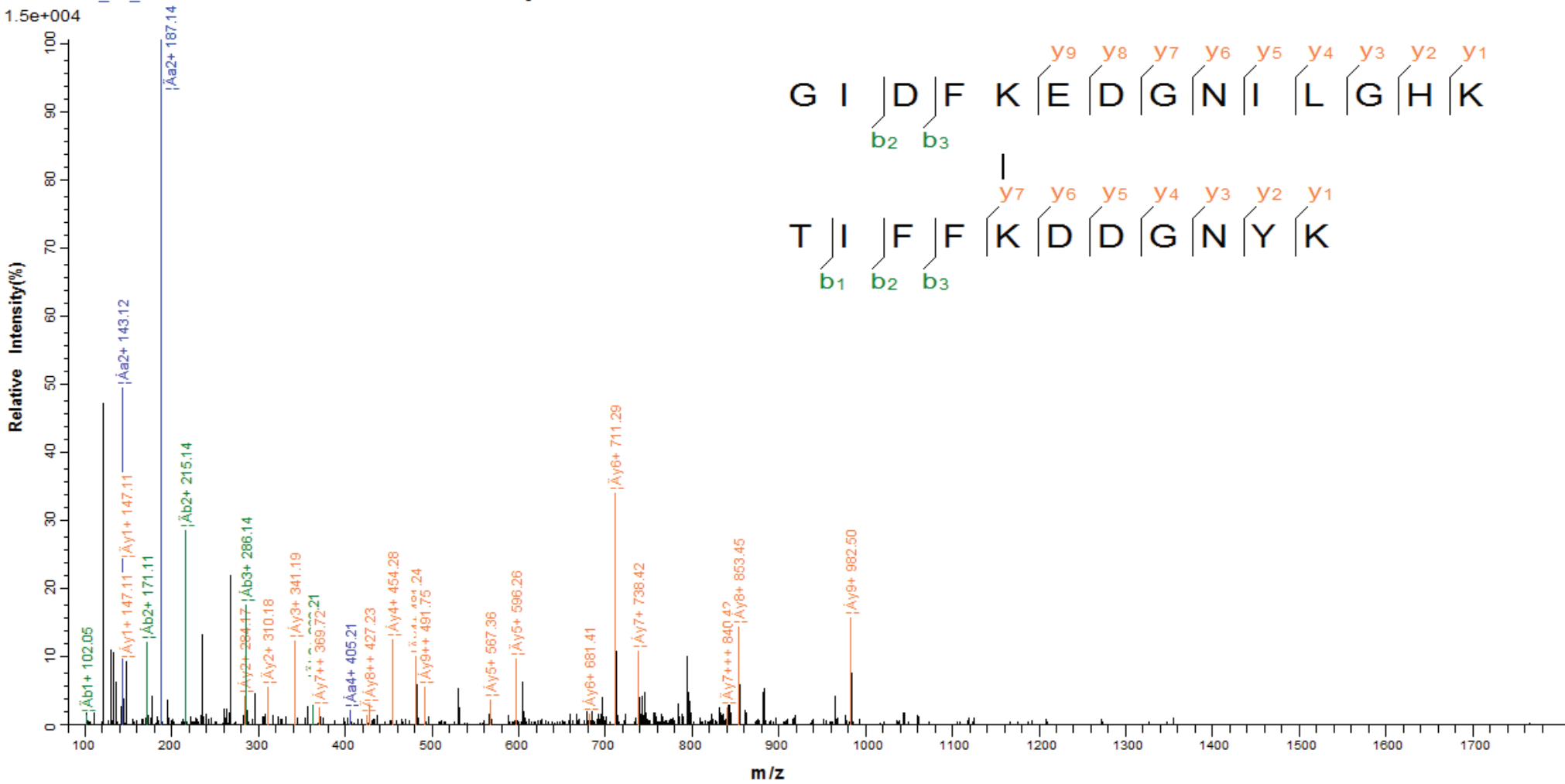
9.9e+003 PC_A8_140201121226.004040.004040.5.dta Seq: NVEDIHVALKR-IMKDFK Charge: 5

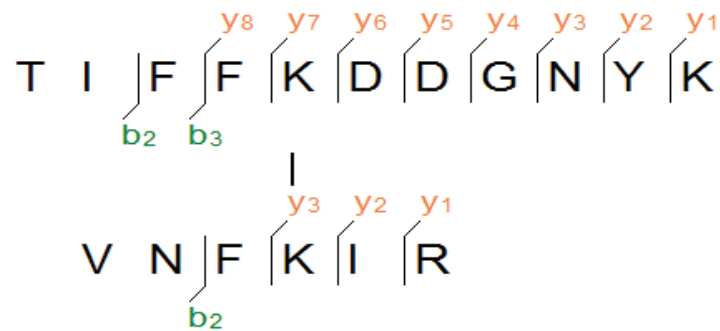
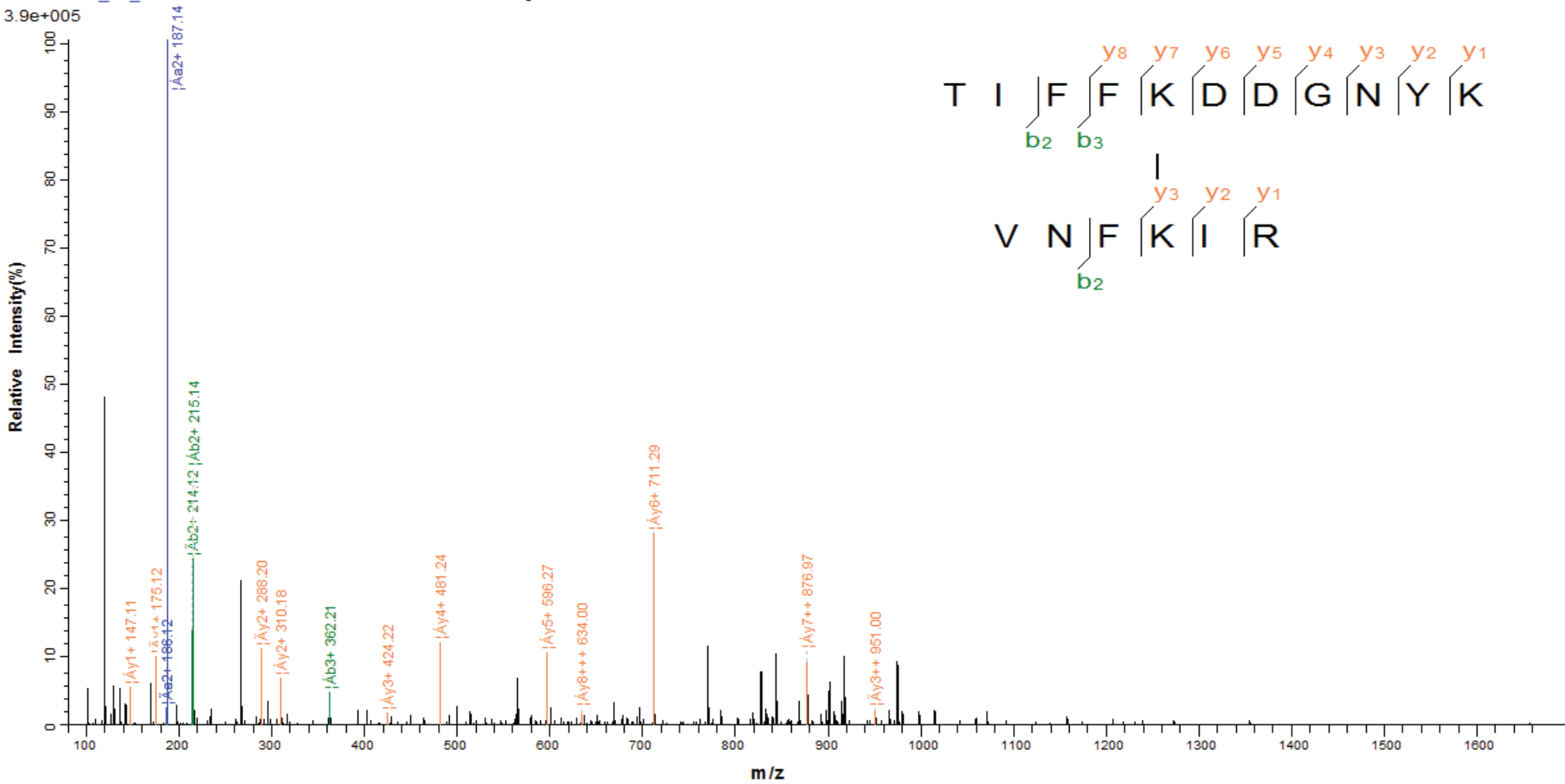


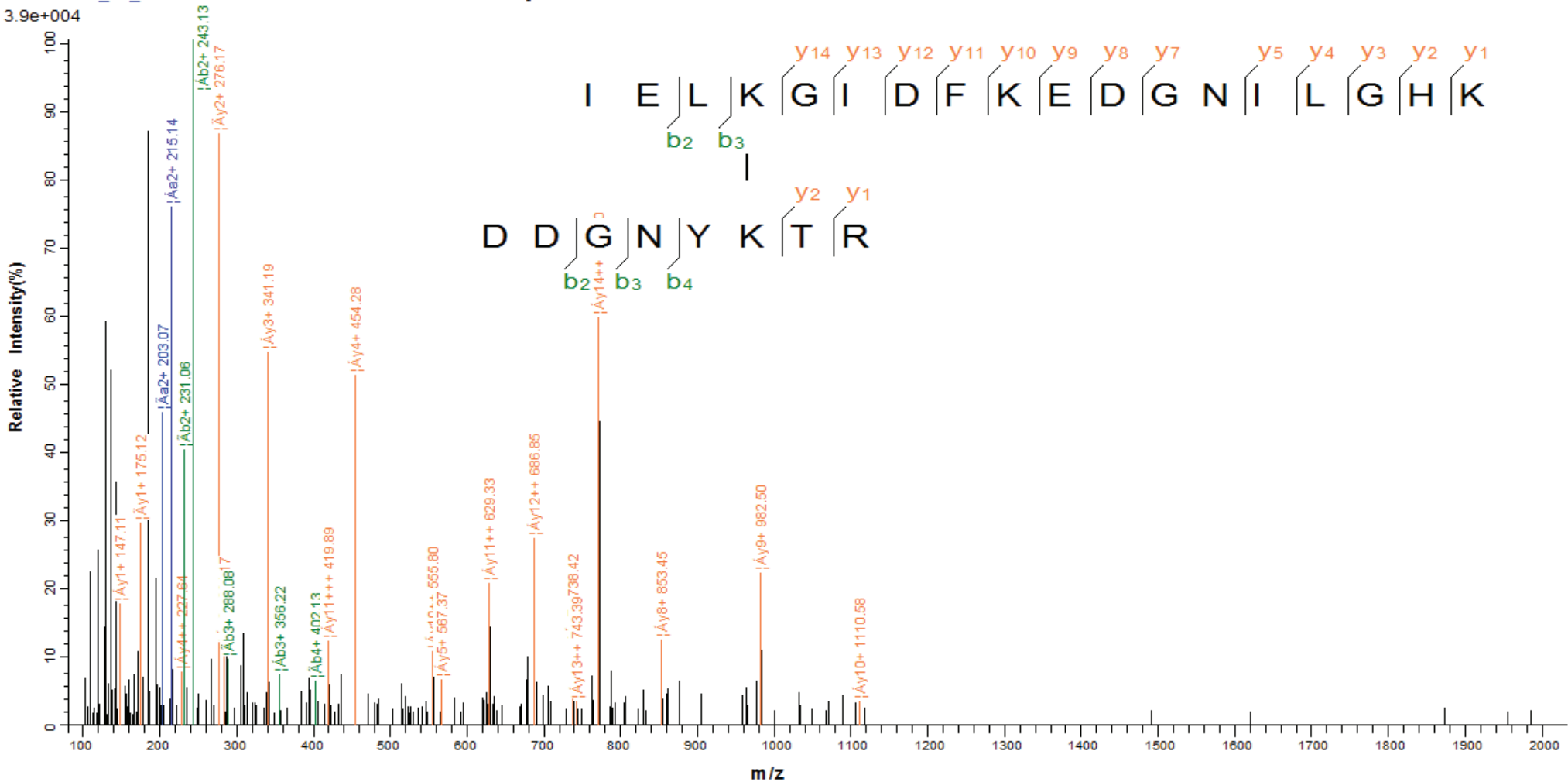
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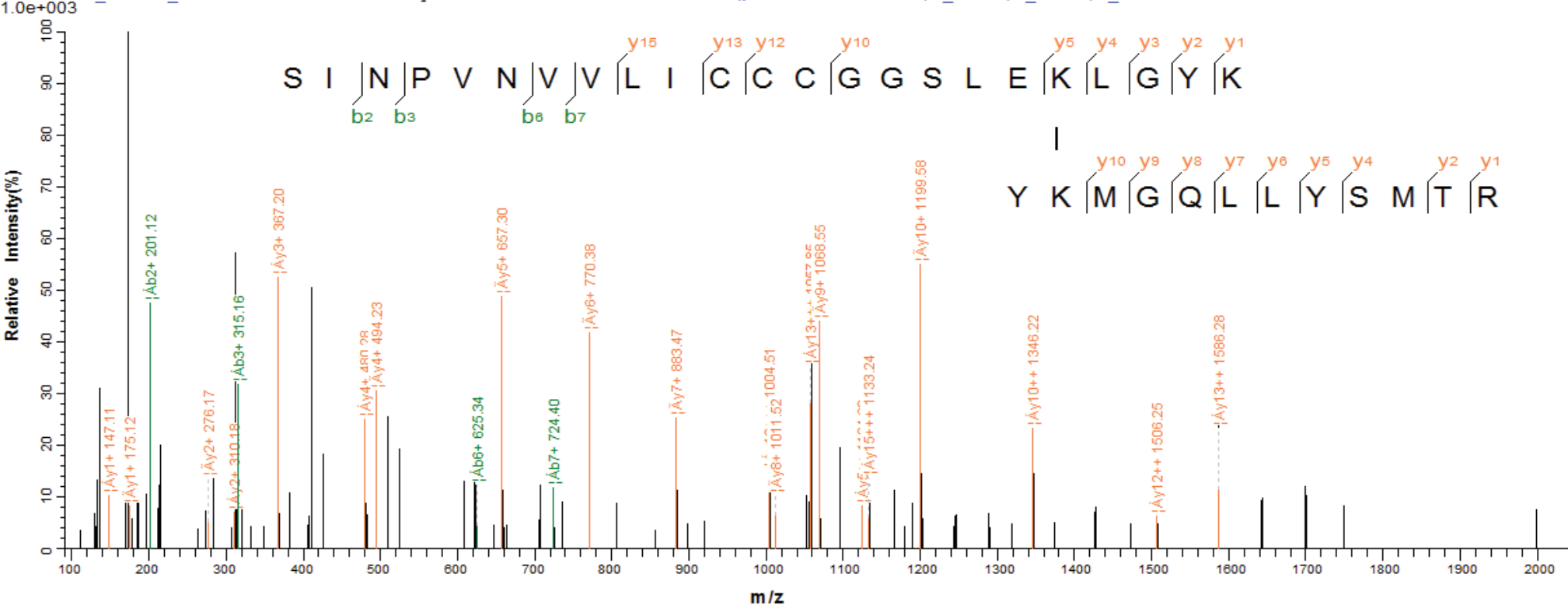




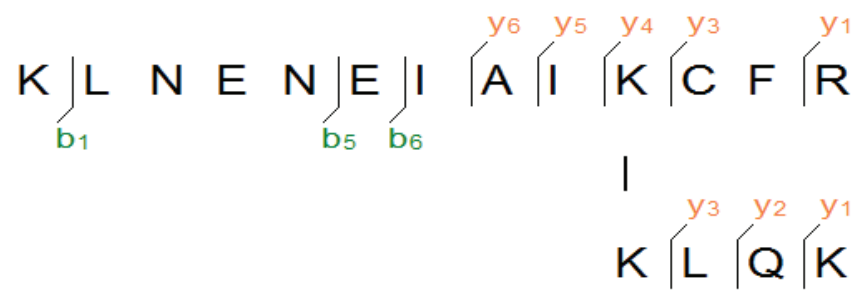
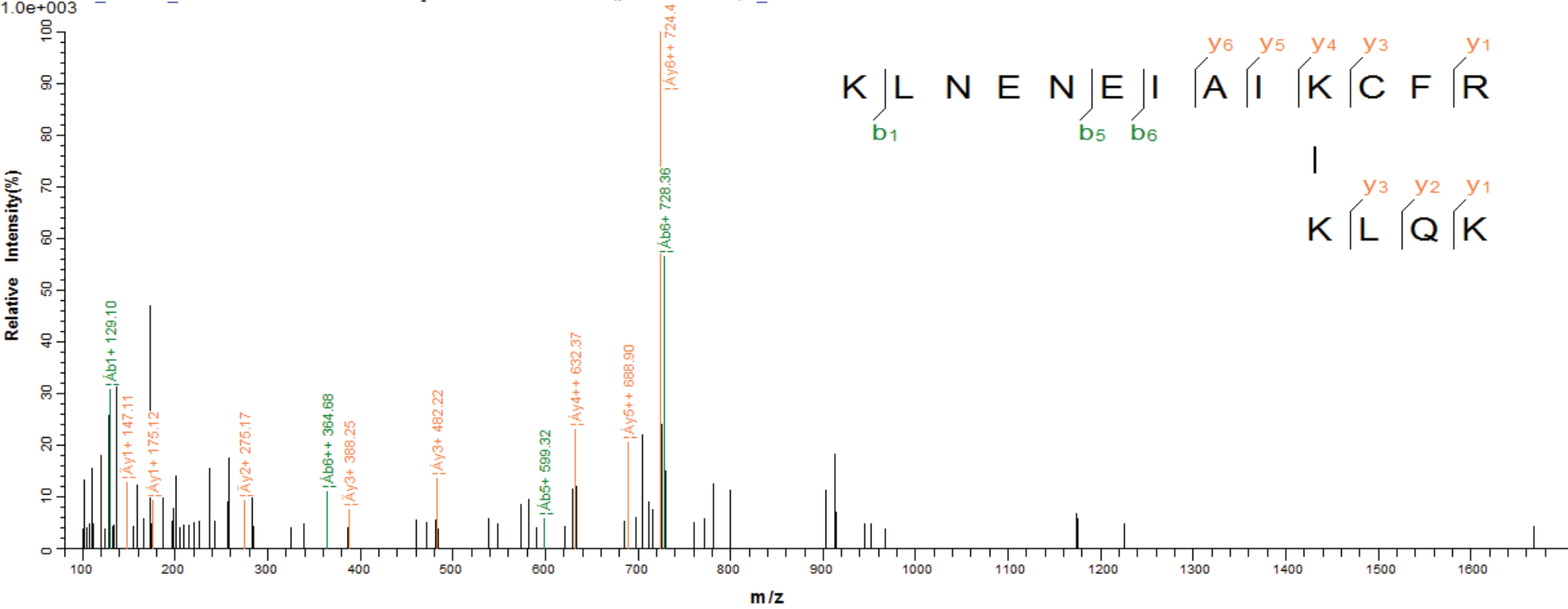


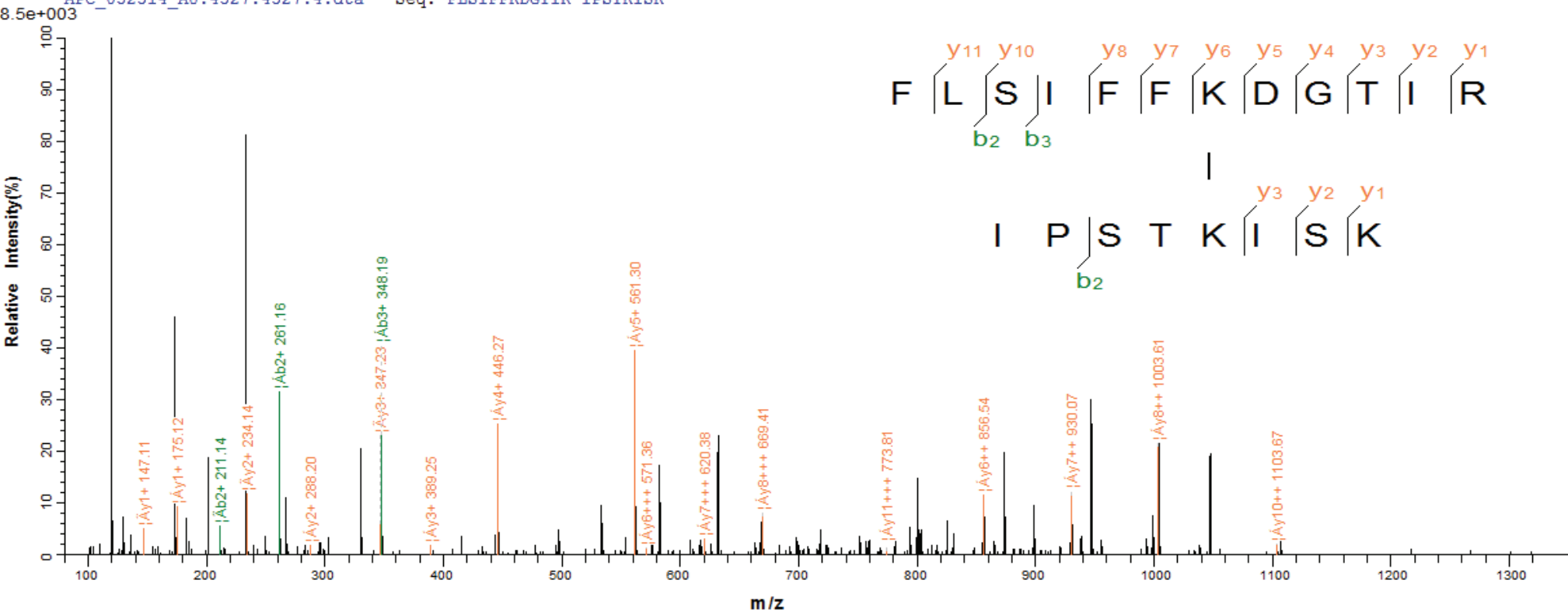




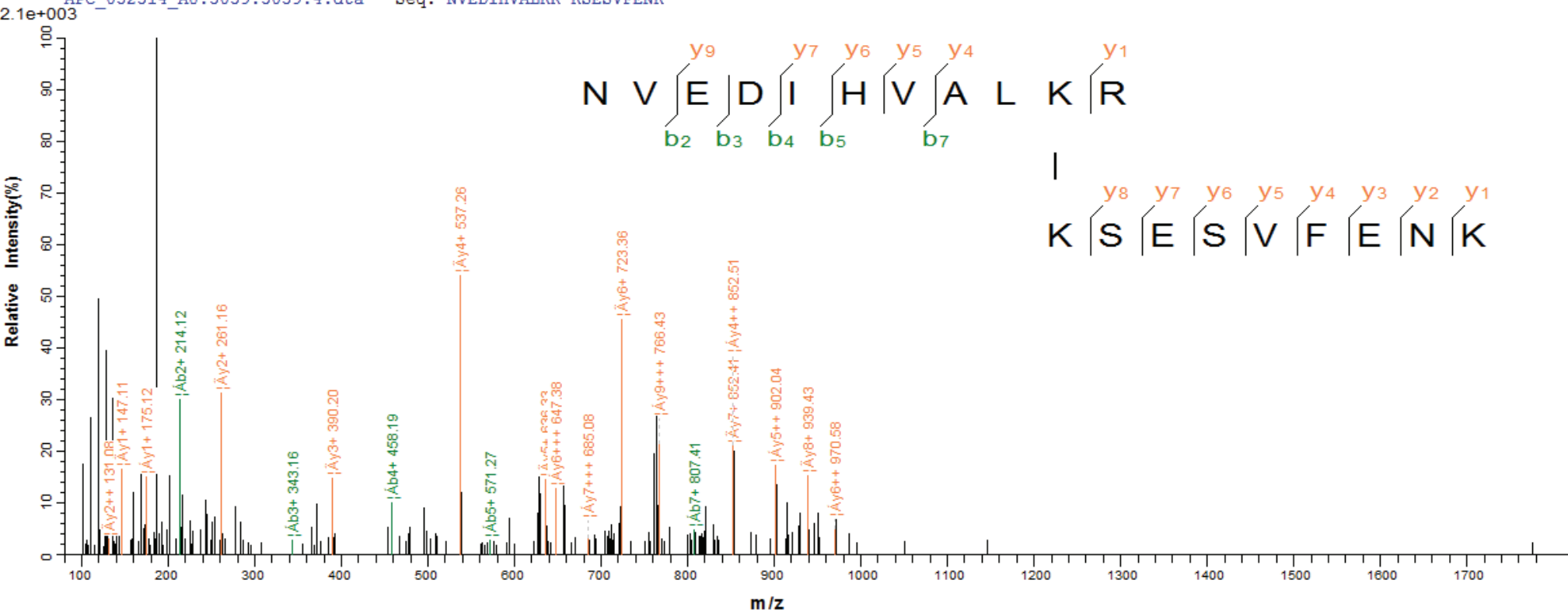


APC_052314_A8.2613.2613.4.dta Seq: KLNENEIAIKCFR-KLQK Mod: 11,C_57



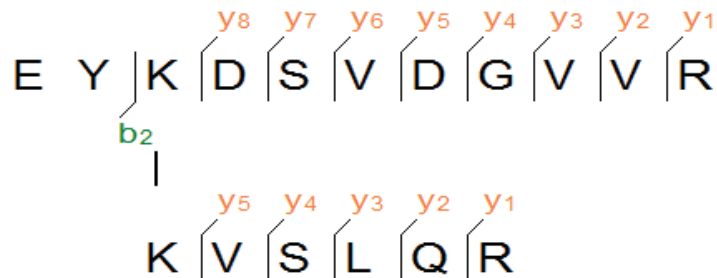
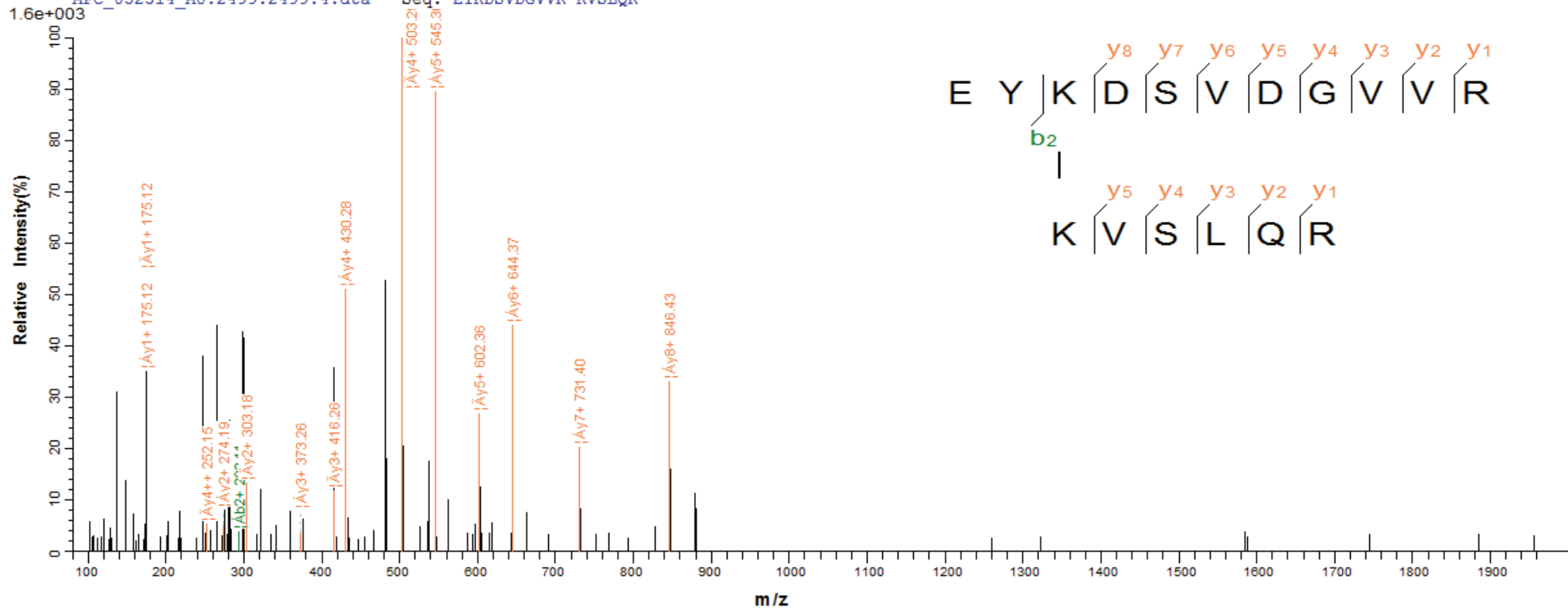


APC_052314_A8.3039.3039.4.dta Seq: NVEDIHVALKR-KSESVFENK



APC_052314_A8.2499.2499.4.dta

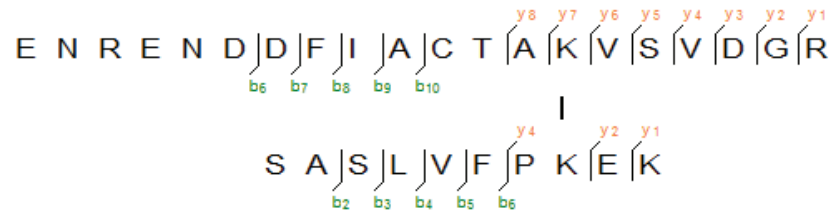
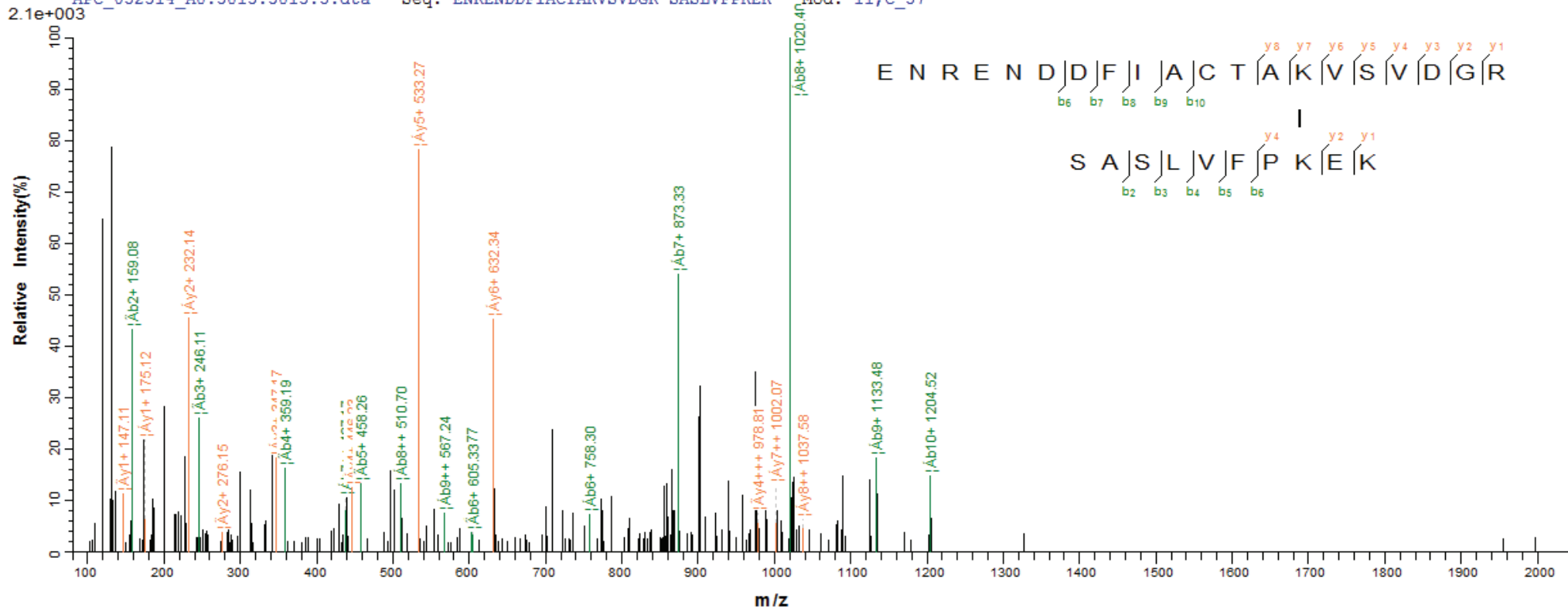
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APC_052314_A8.3813.3813.5.dta

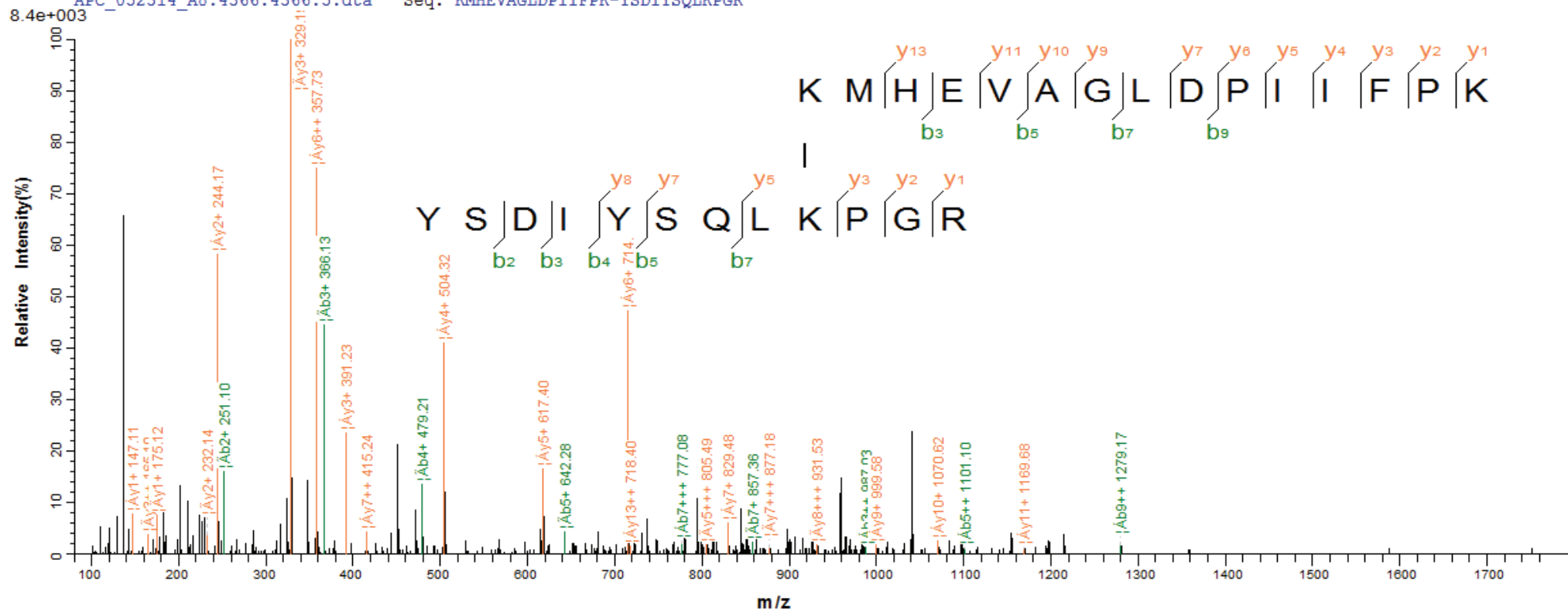
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Mod: 11, C_57

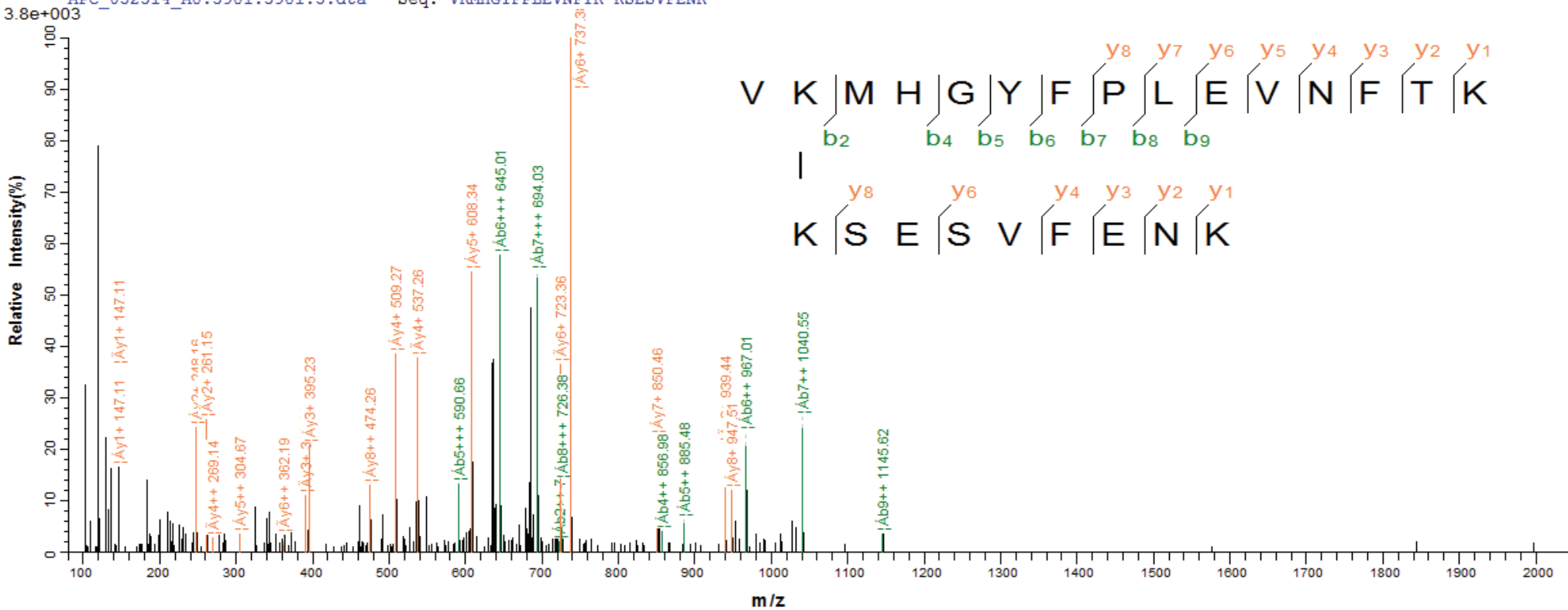


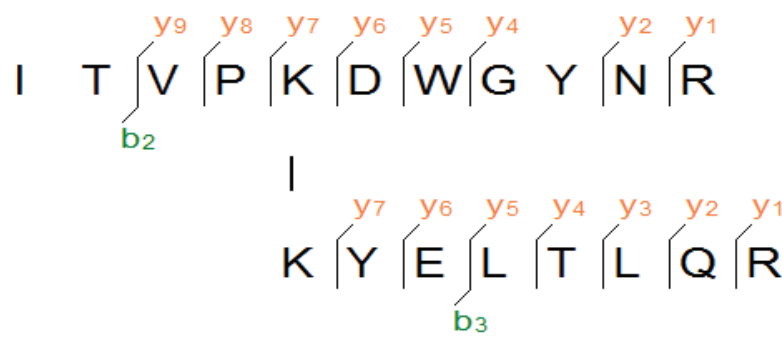
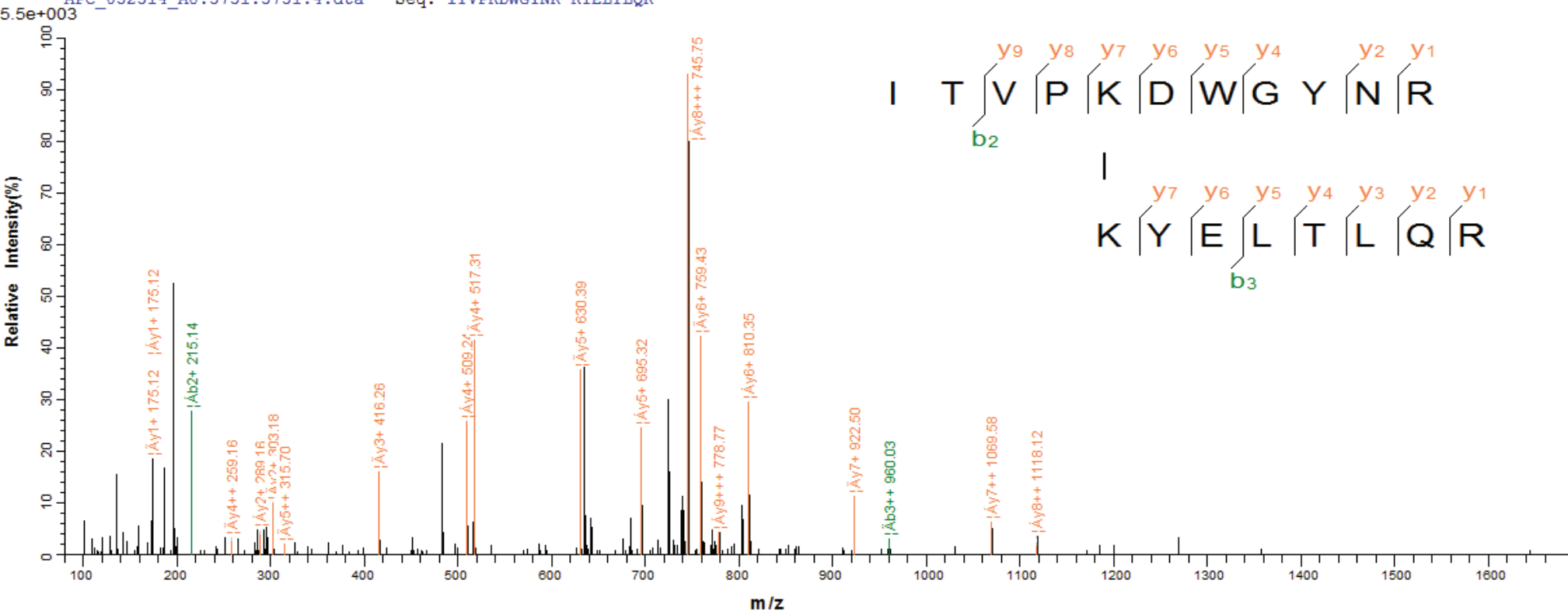
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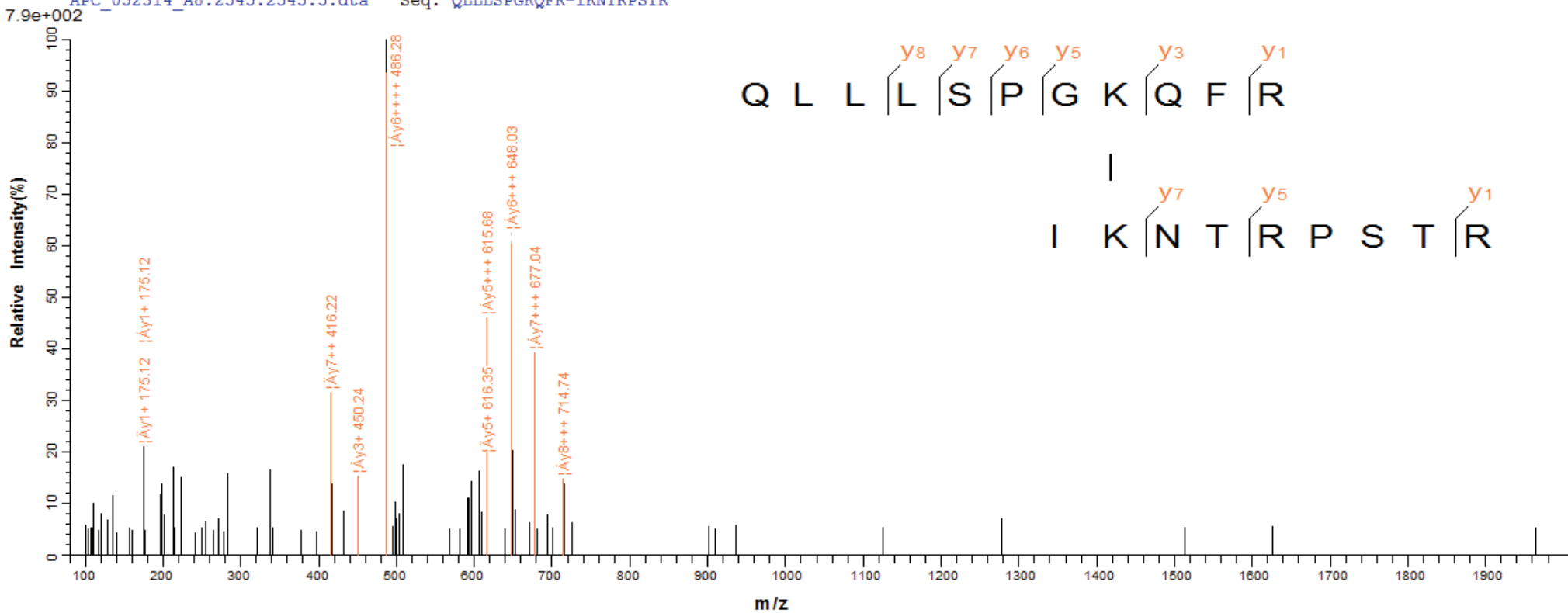


APC_052314_A8.3981.3981.5.dta Seq: VKMHGYFPLEVNF TK-KSESVFENK

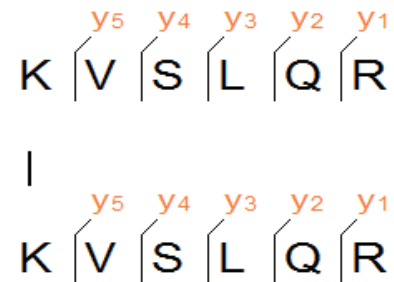
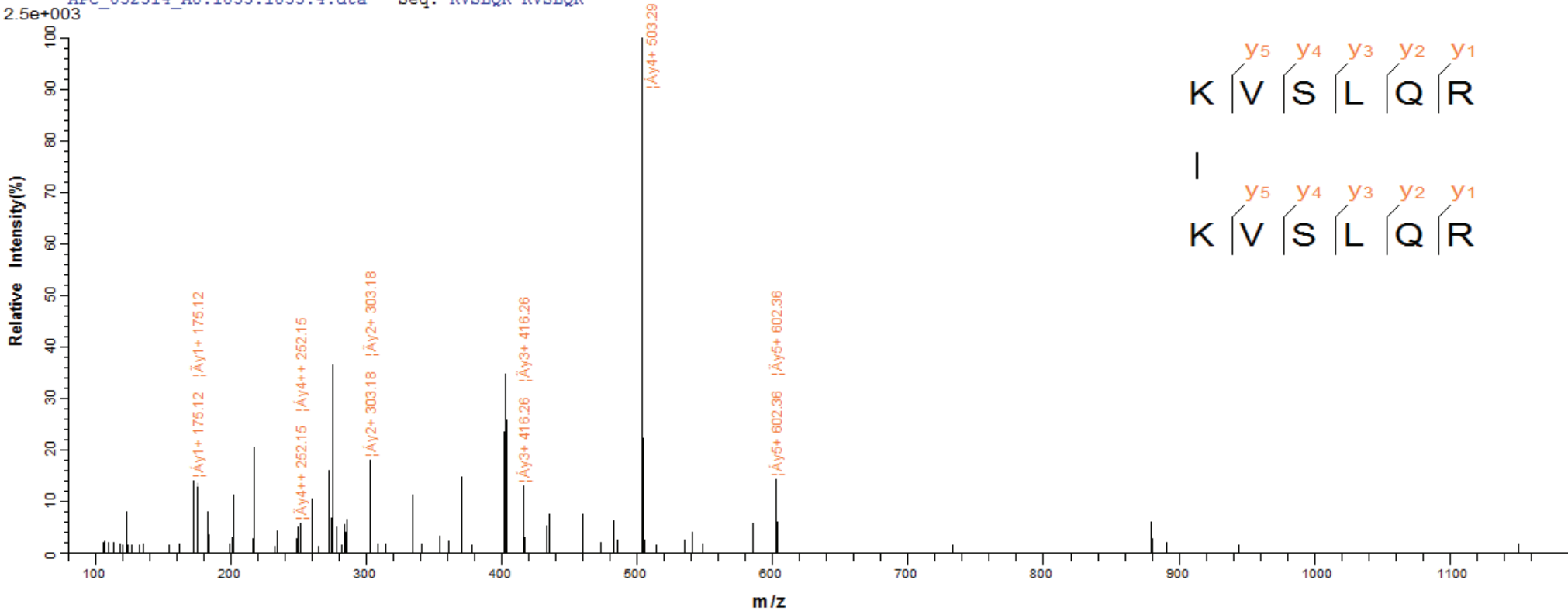




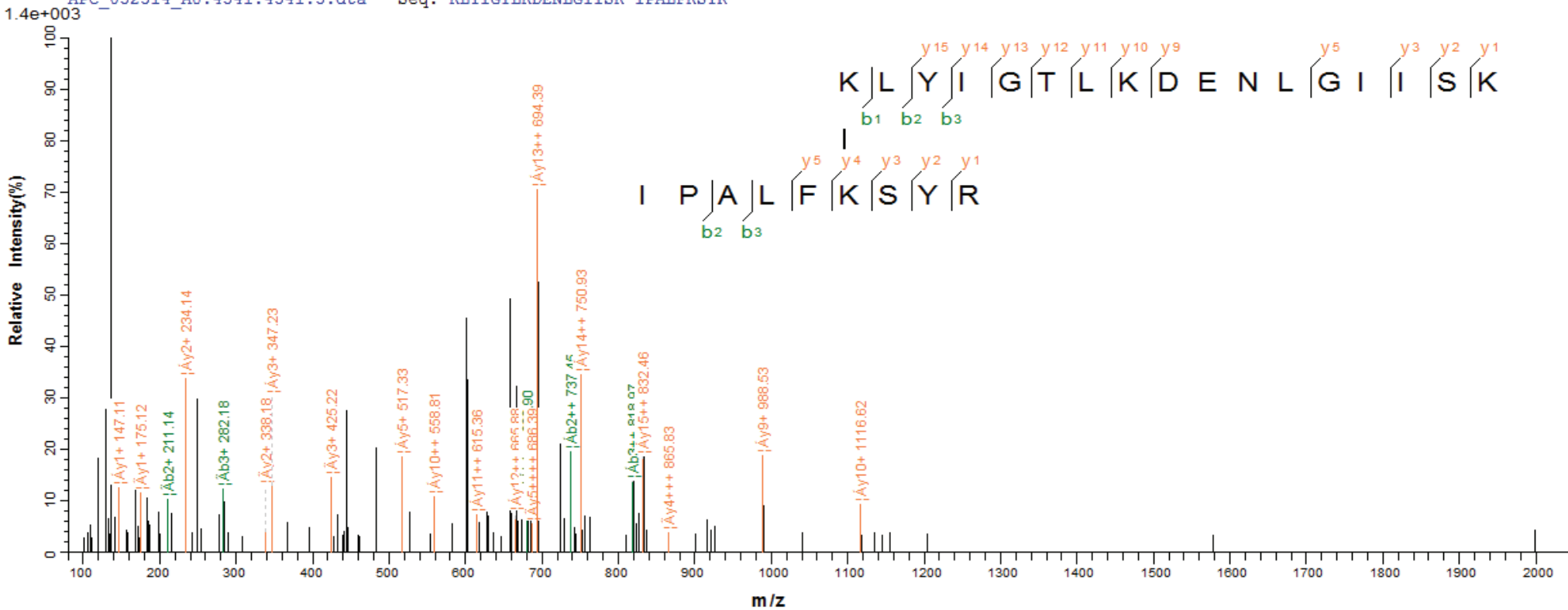
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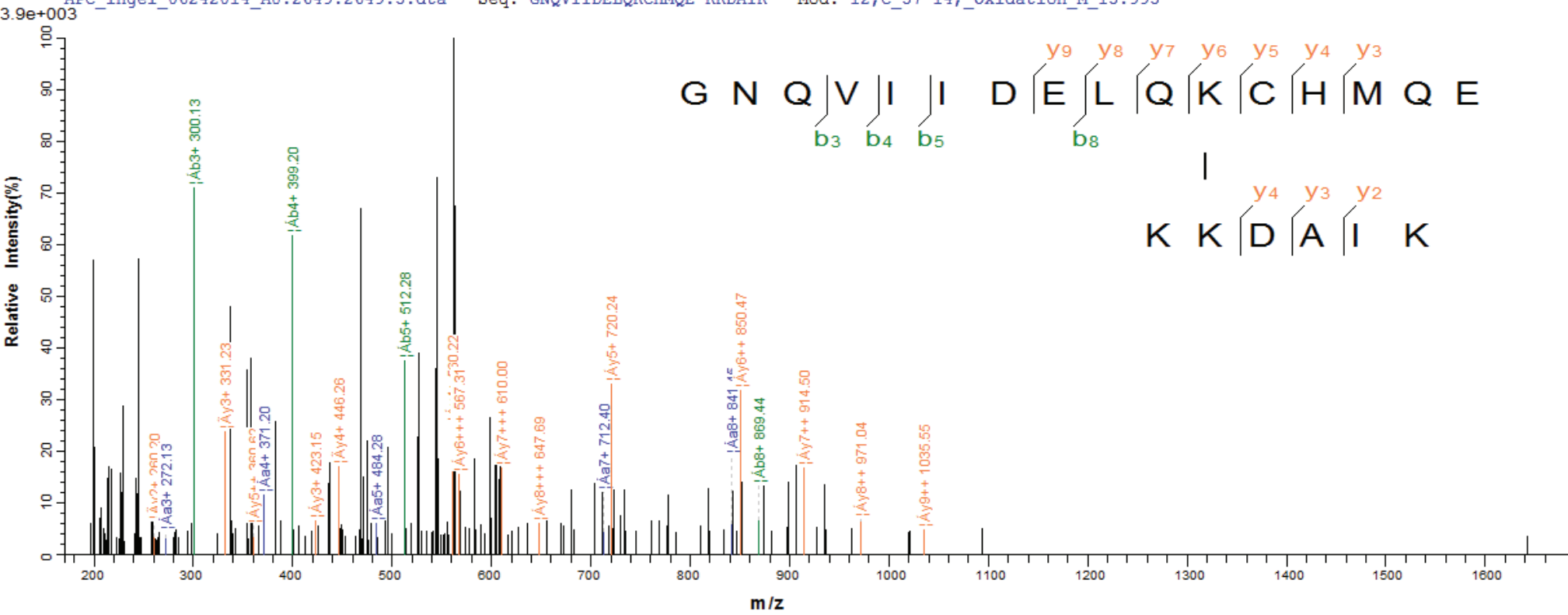


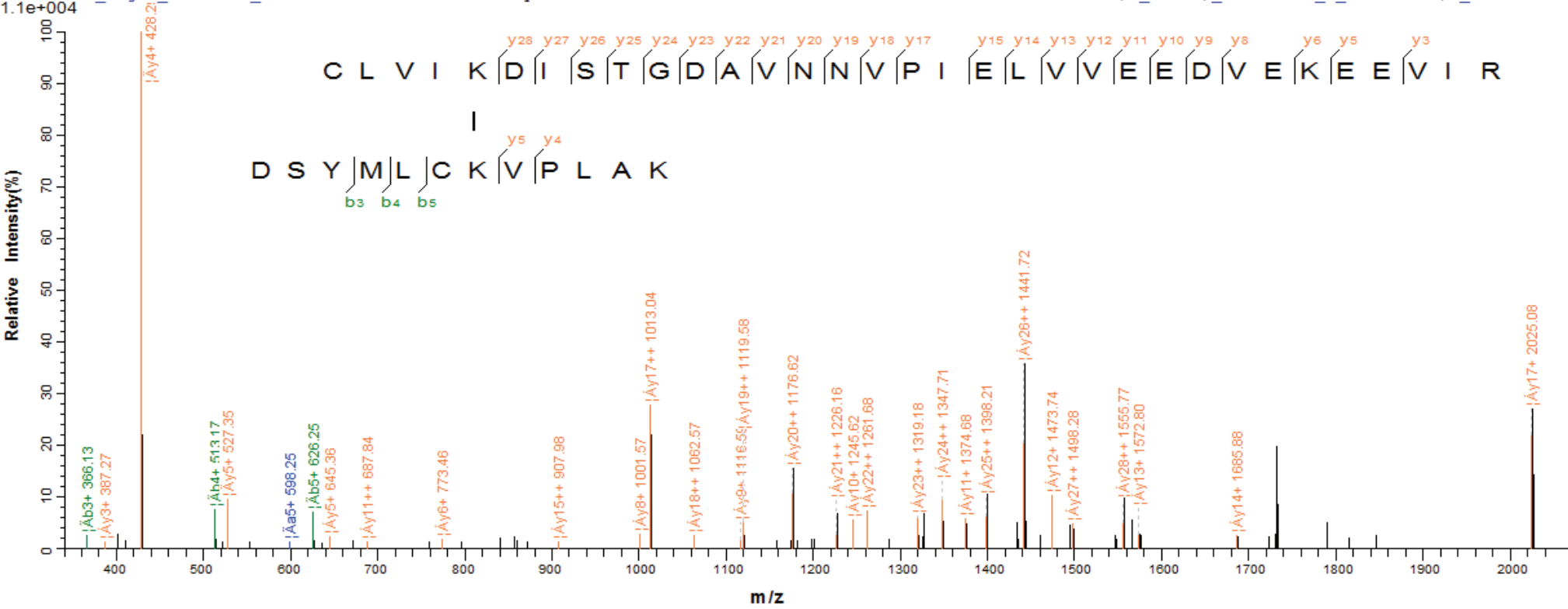
APC_052314_A8.1853.1853.4.dta Seq: KVSLQR-KVSLQR



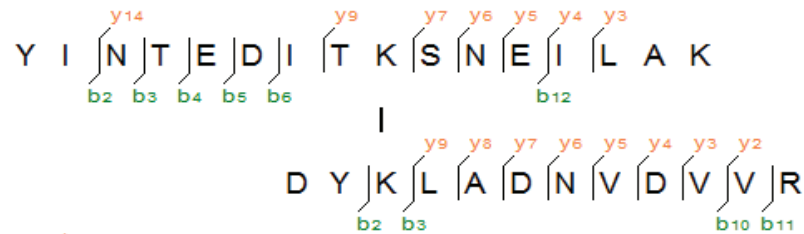
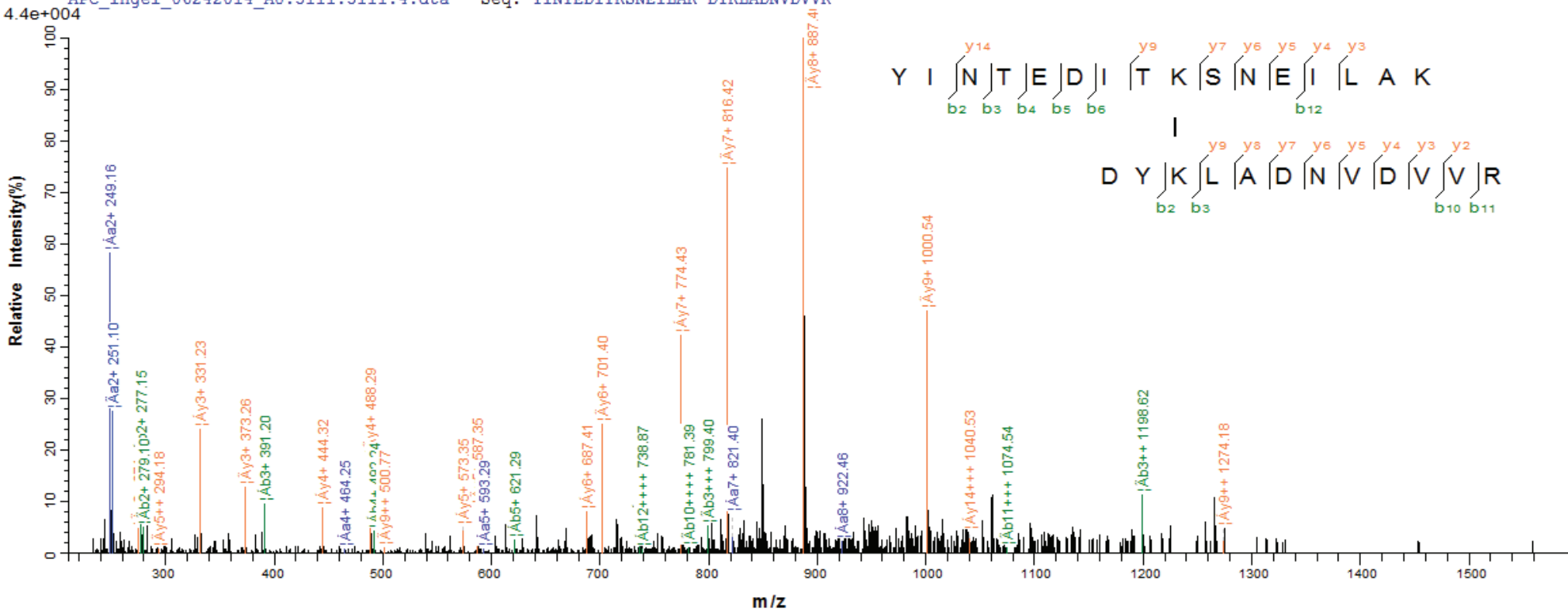
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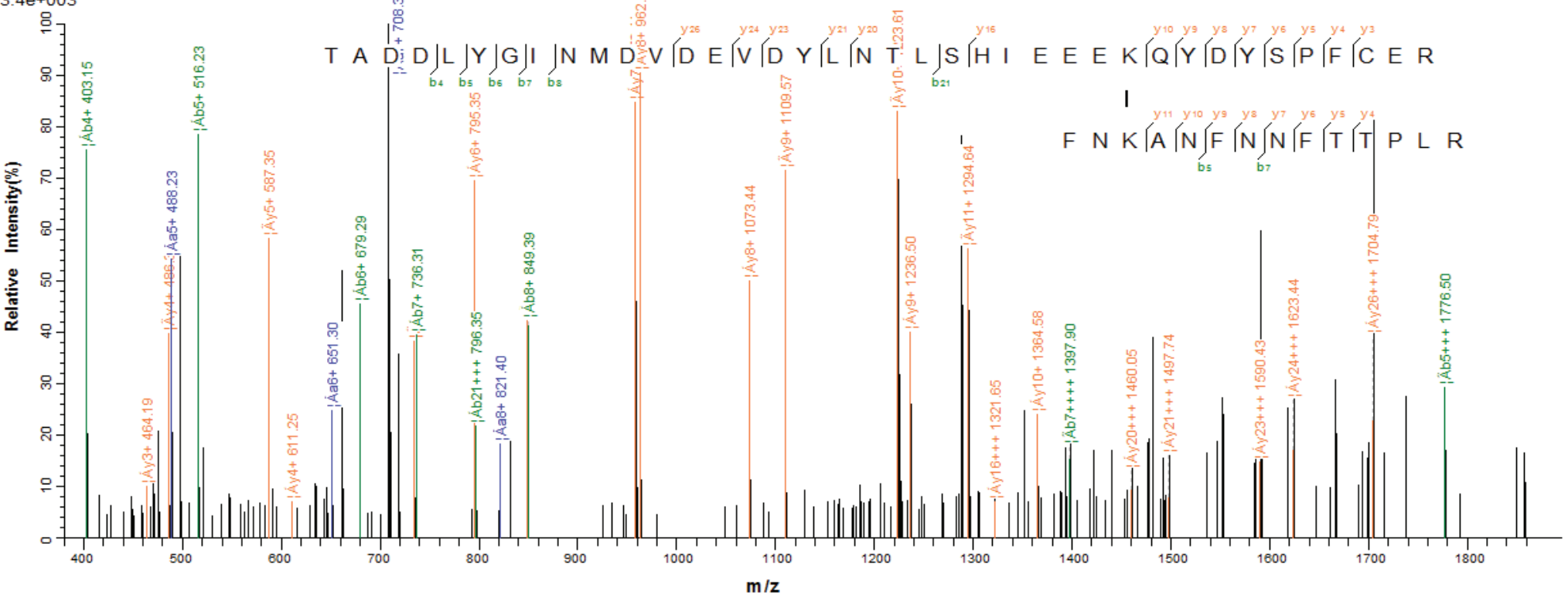


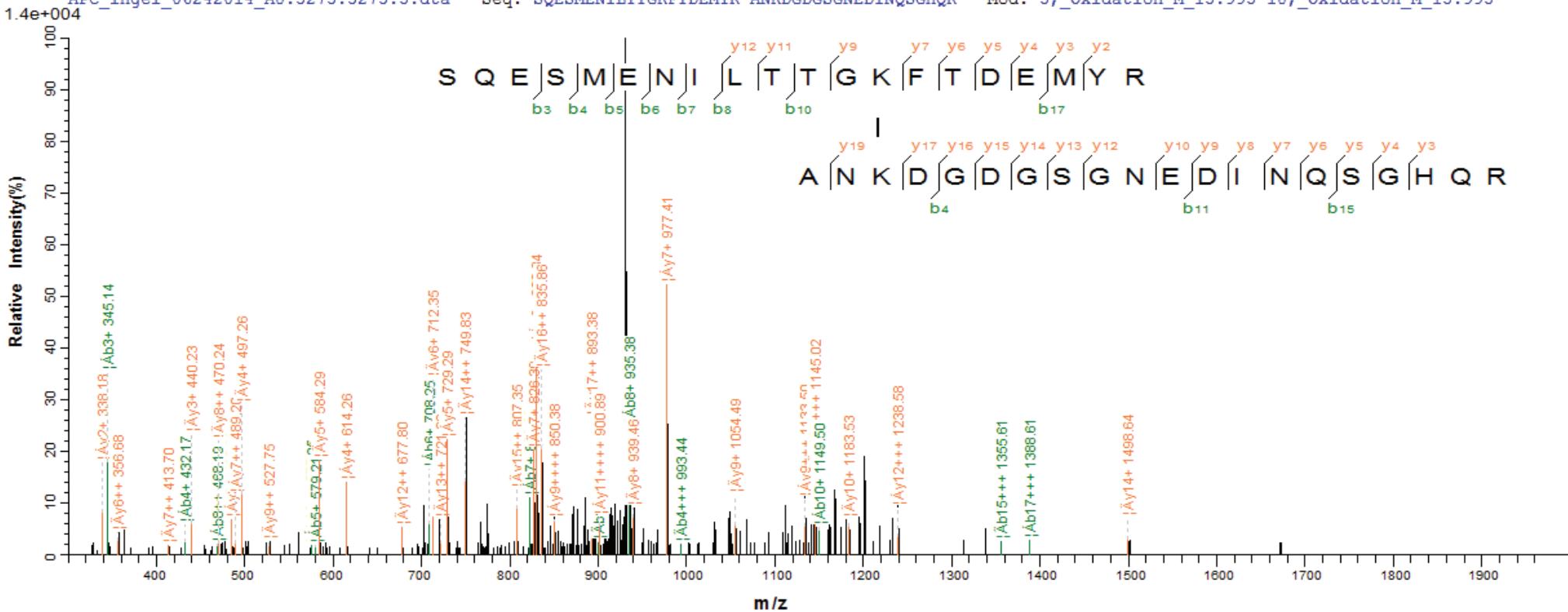


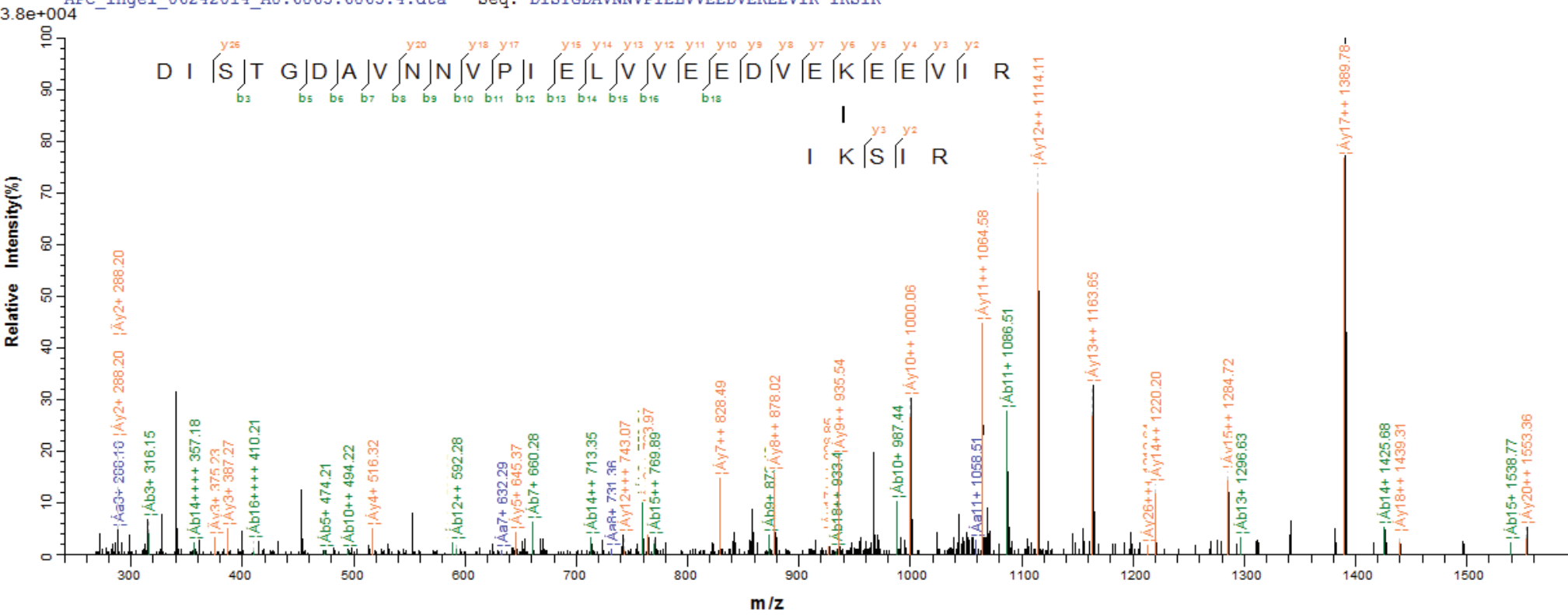


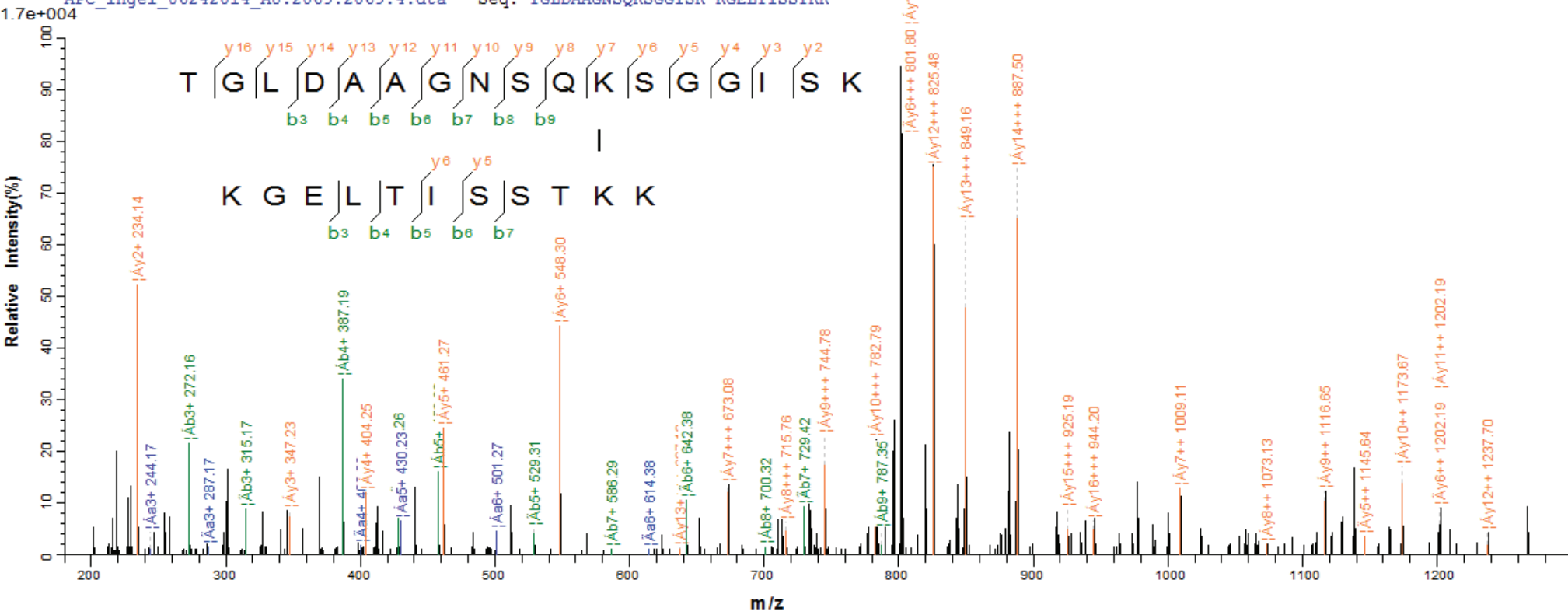
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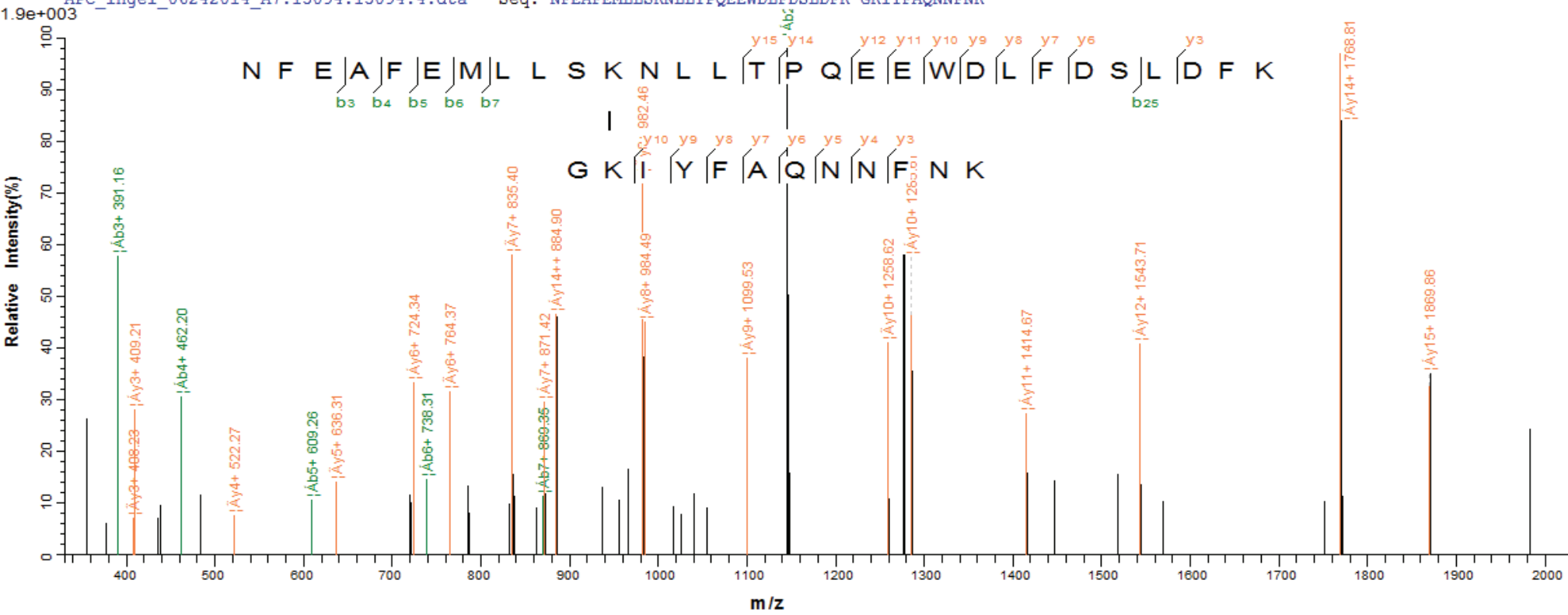




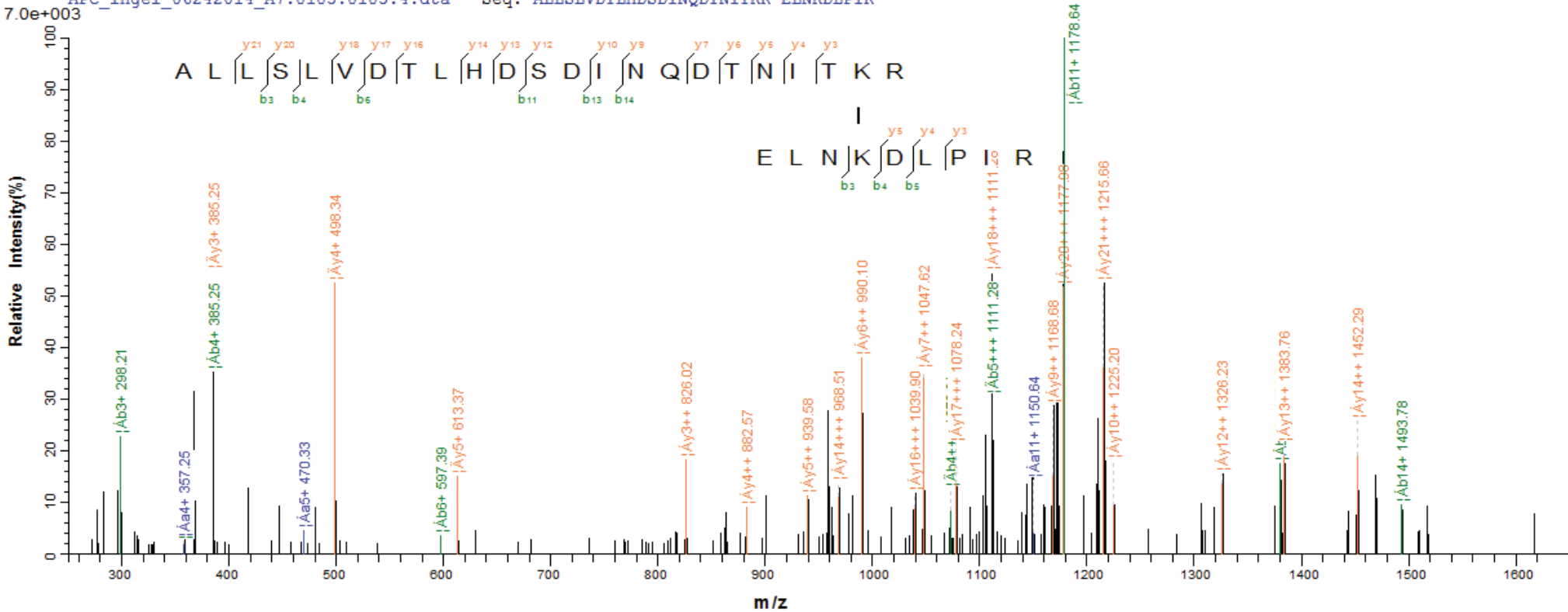


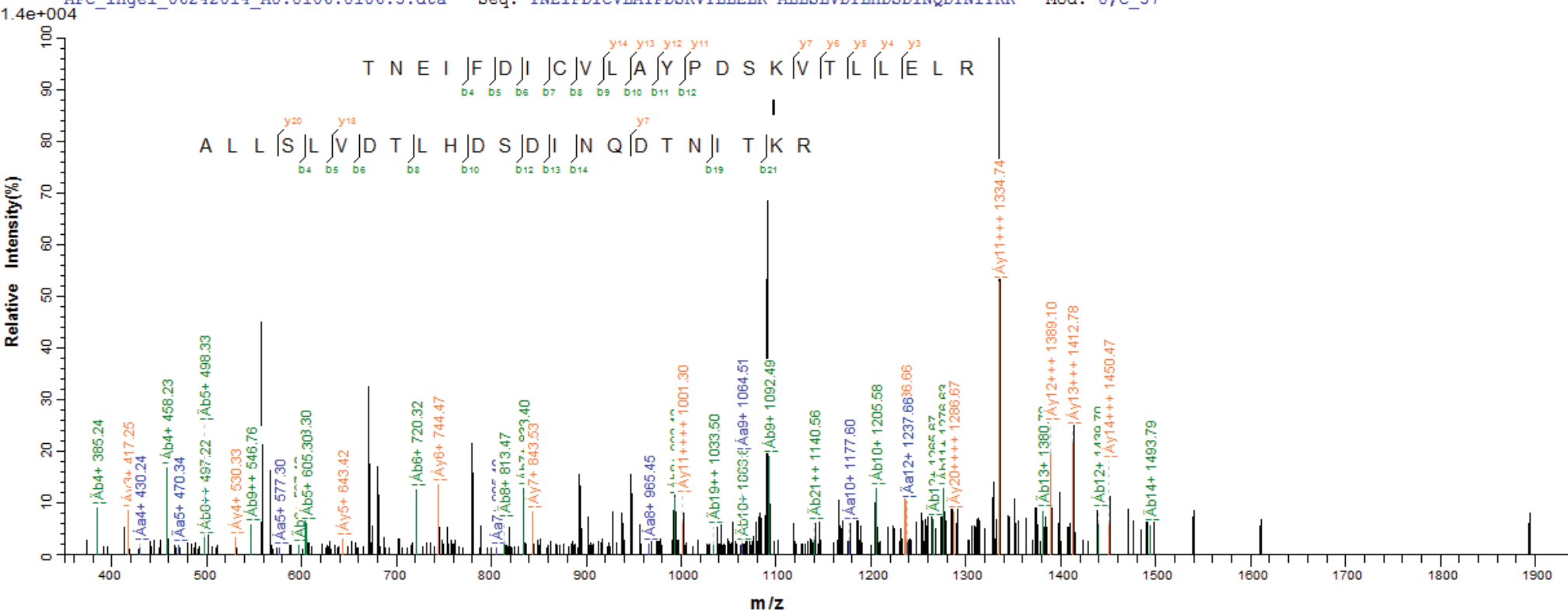


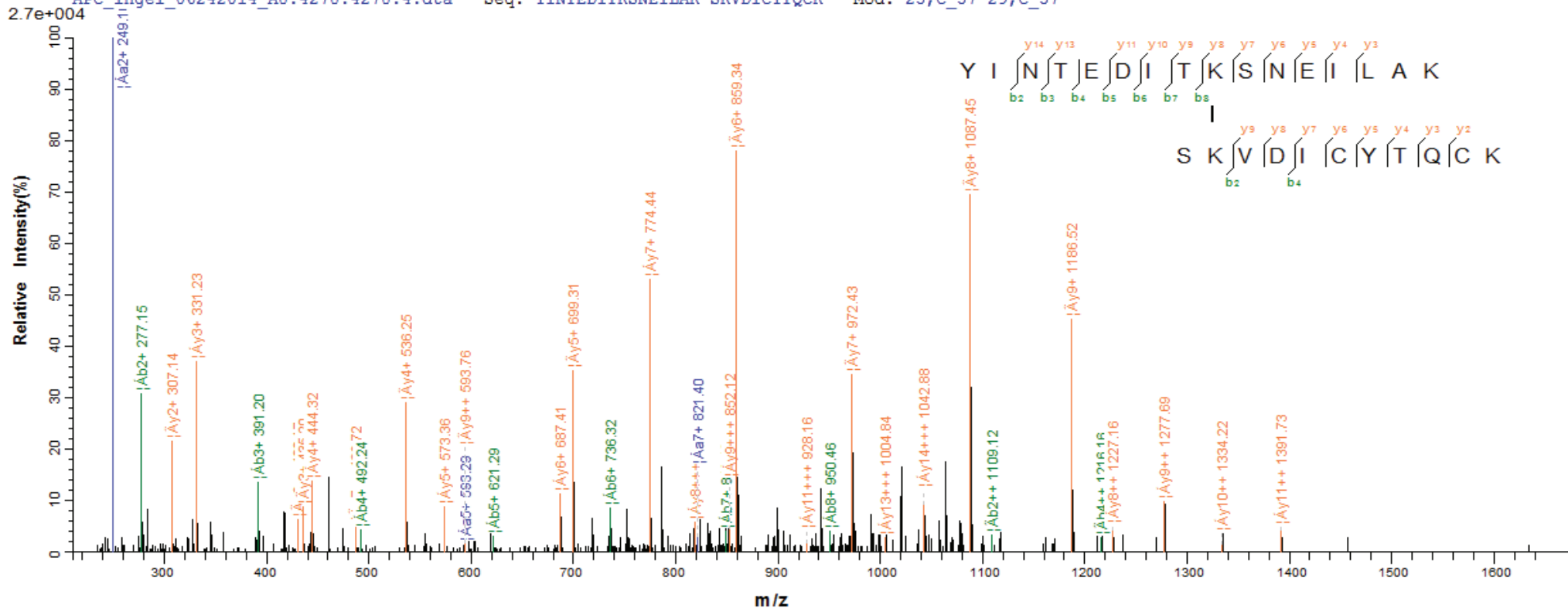




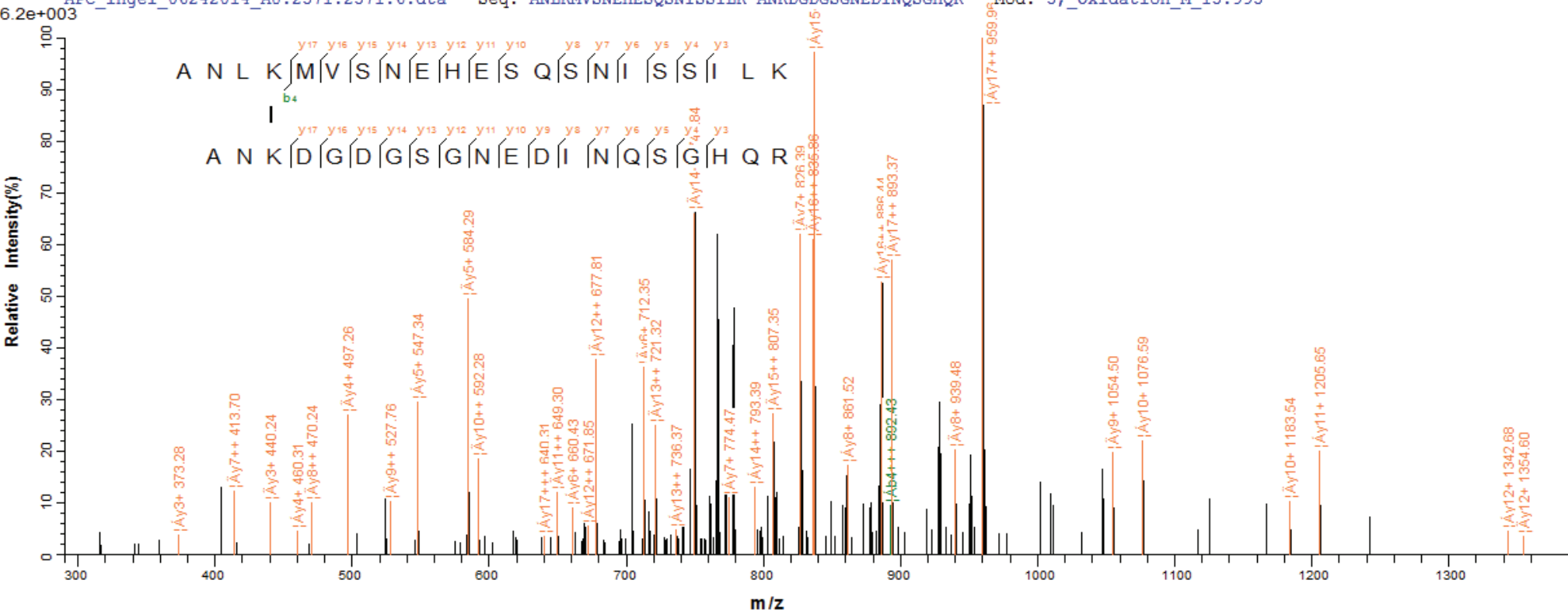
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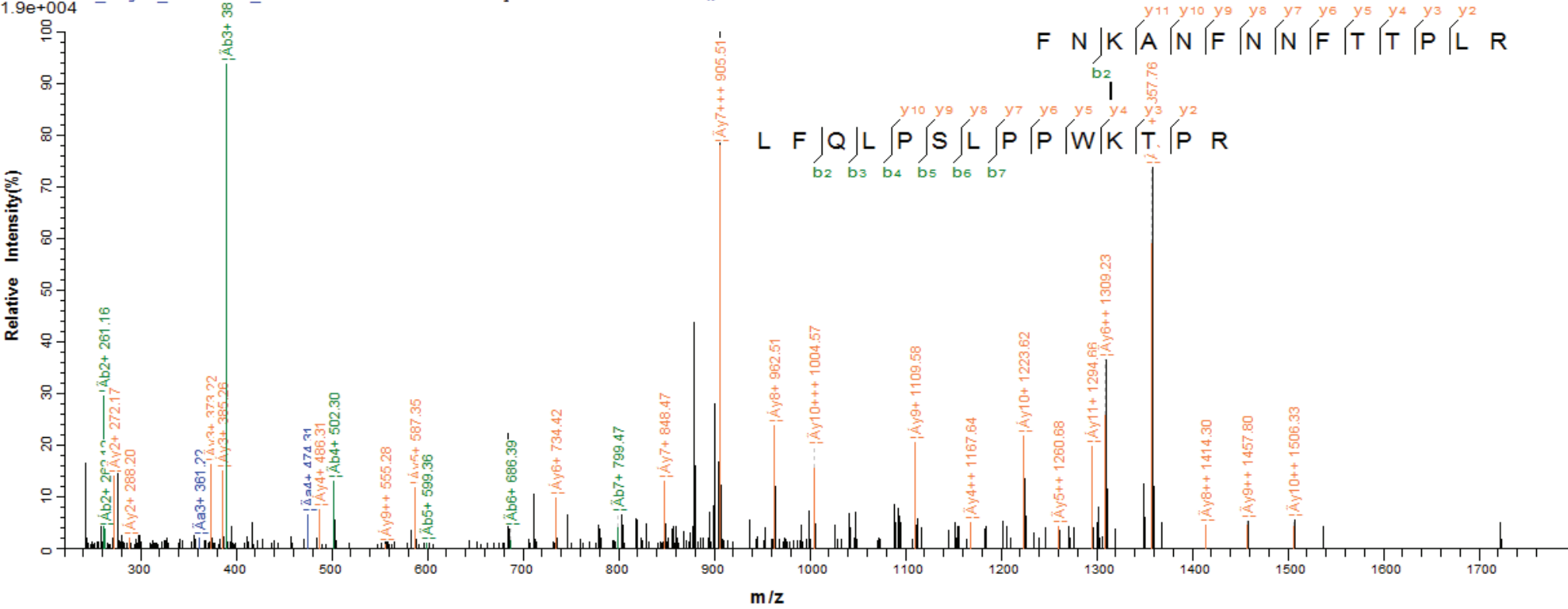


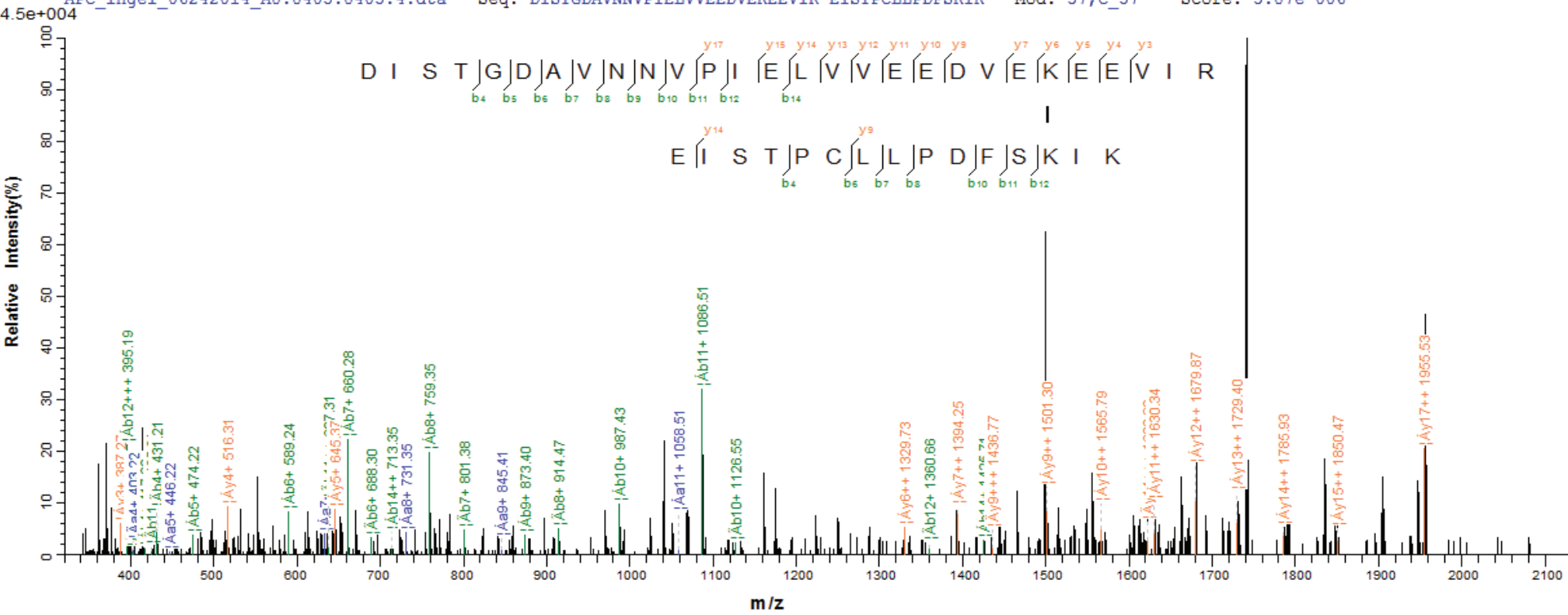




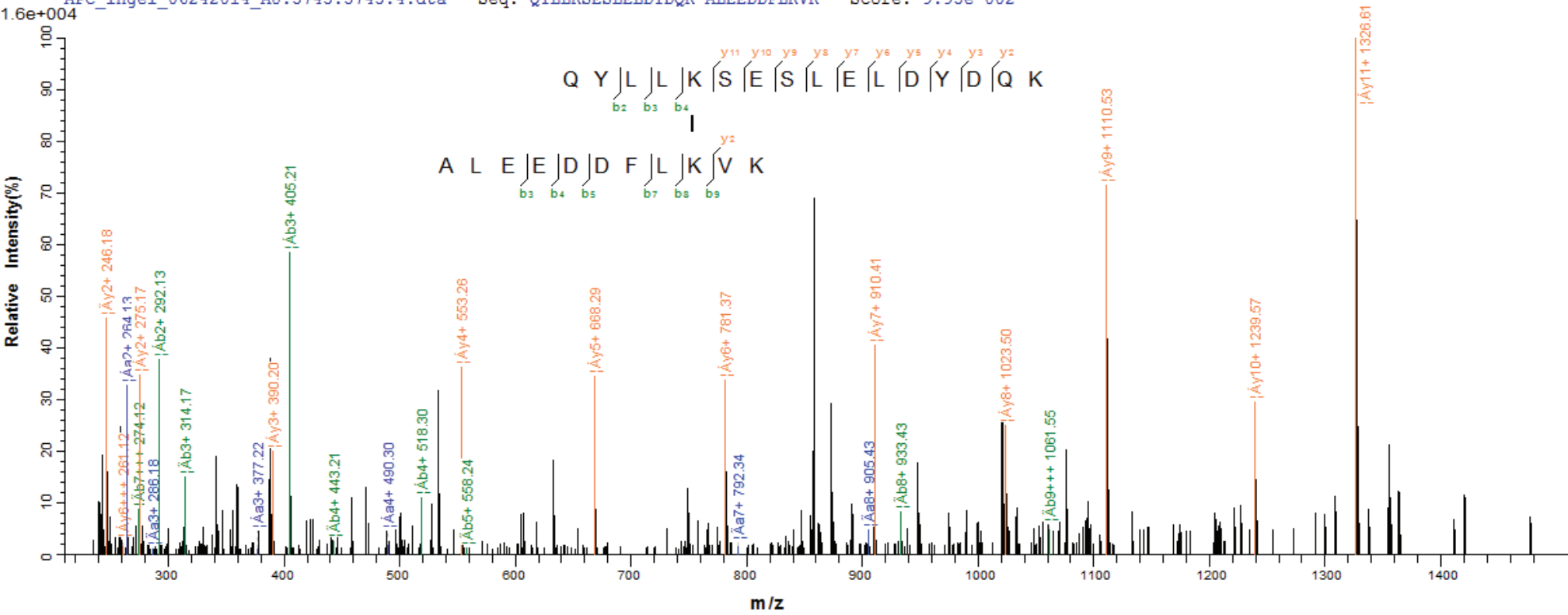
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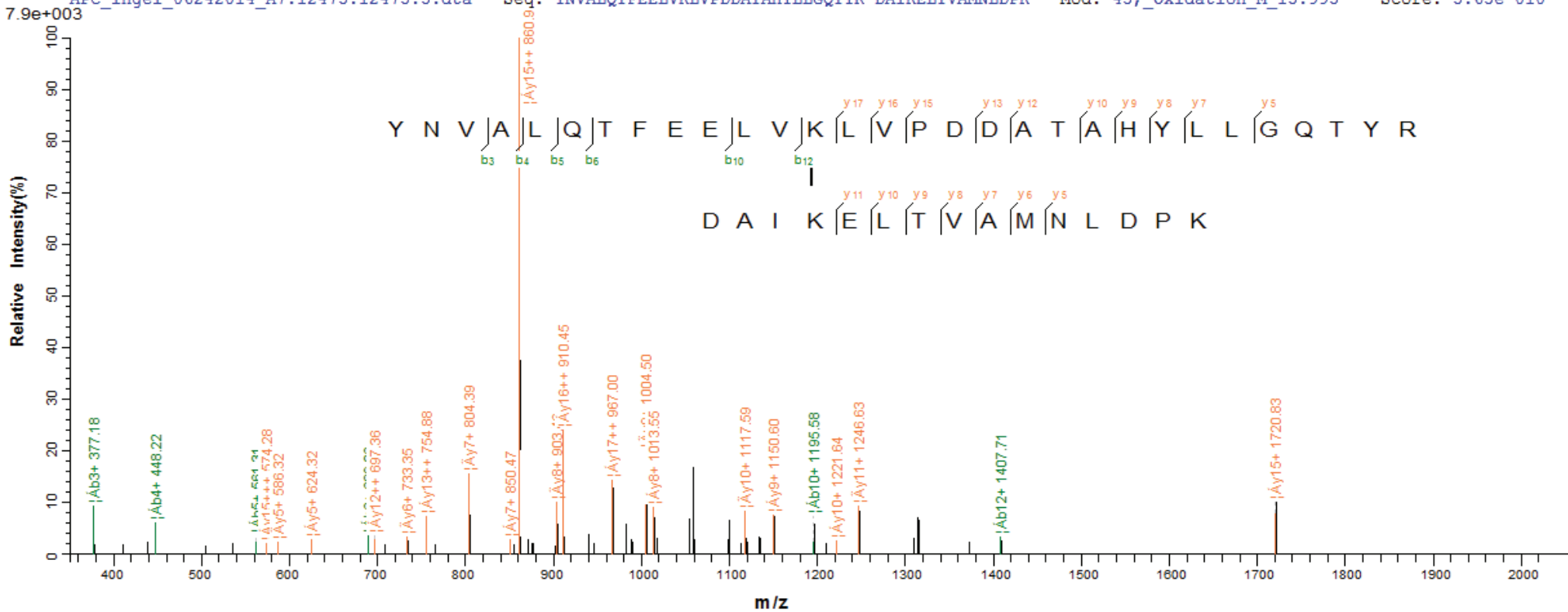




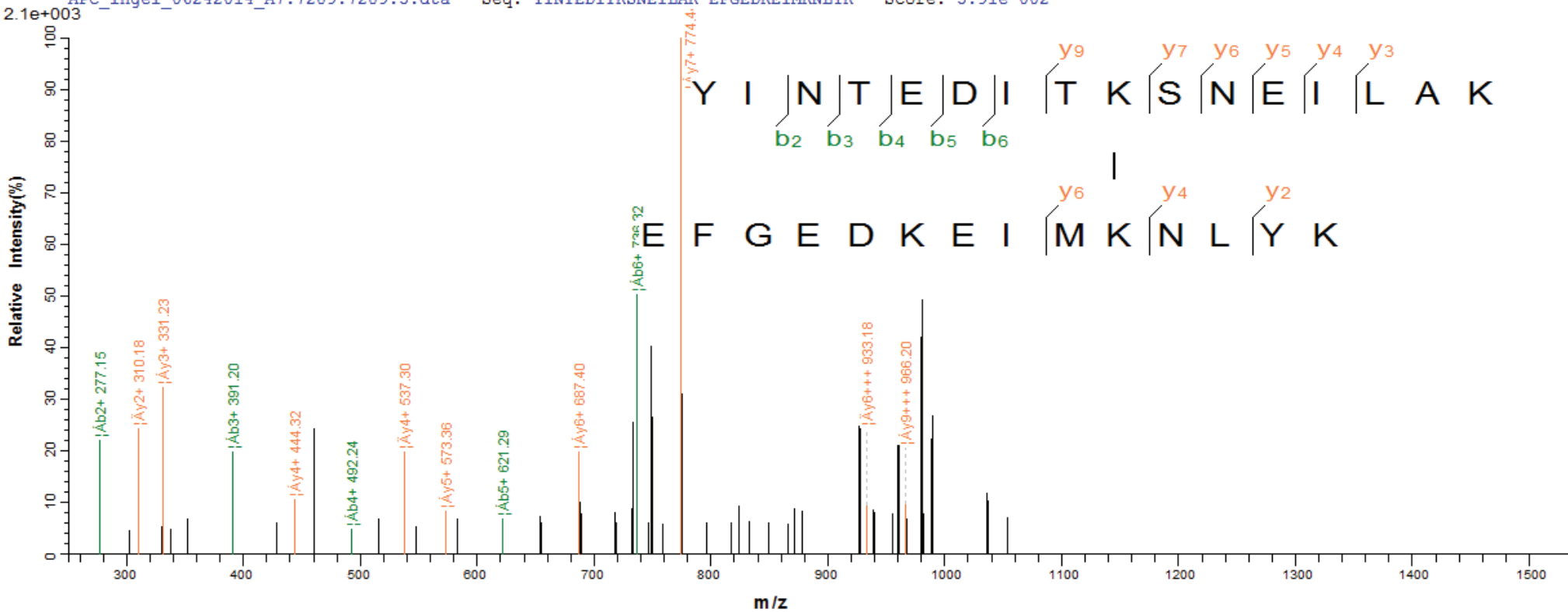


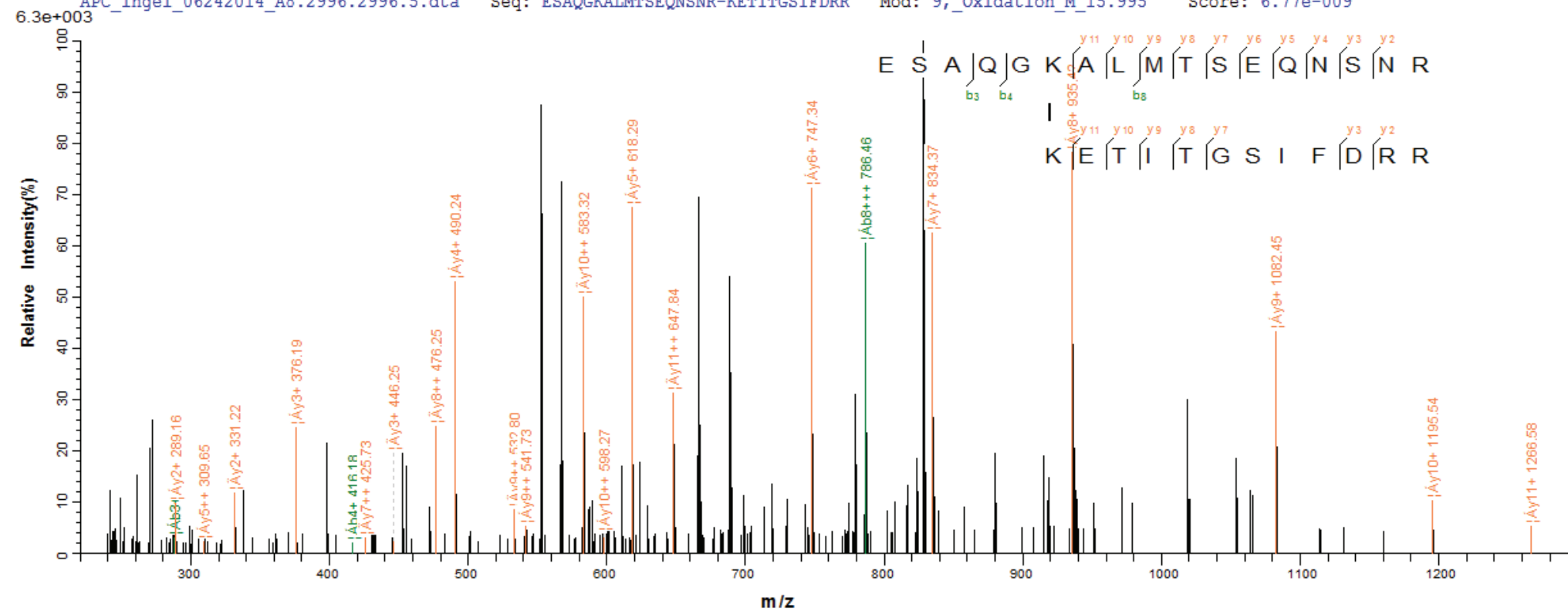
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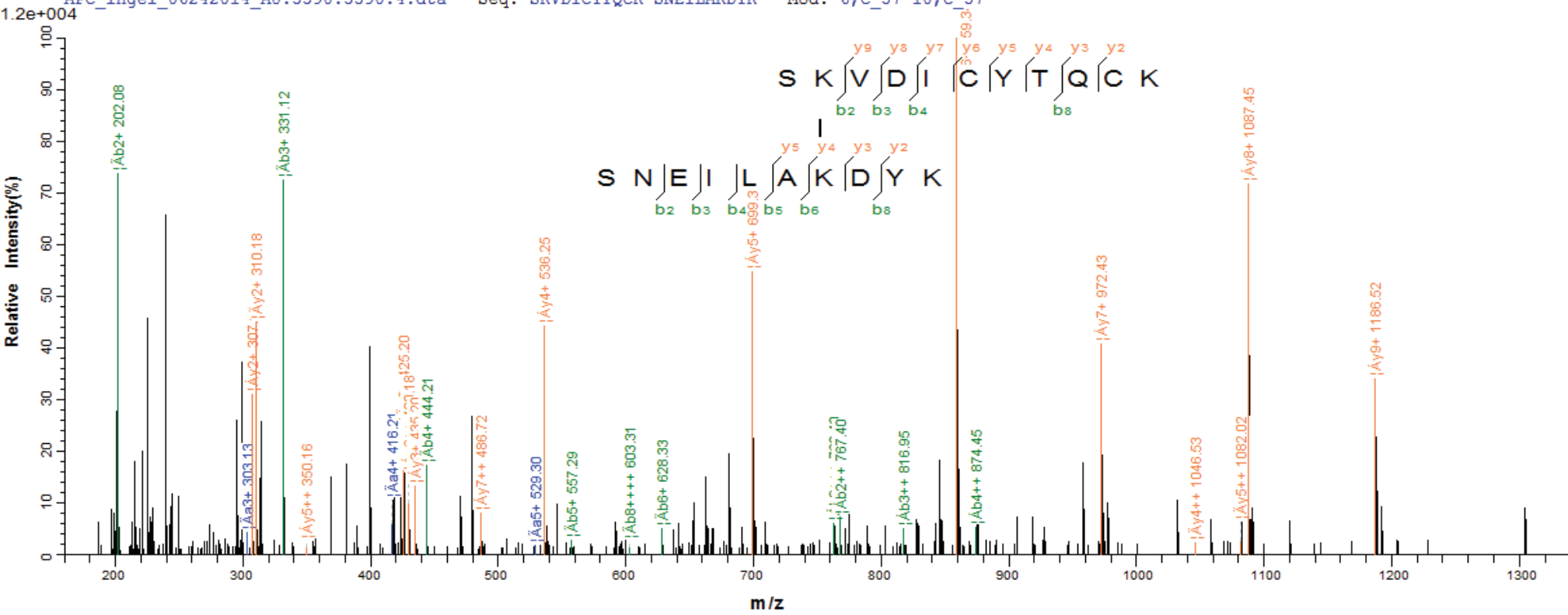




APC_ingel_06242014_A7.7289.7289.5.dta Seq: YINTEDITSNEILAK-EFGEDKEIMKNLYK Score: 5.91e-002

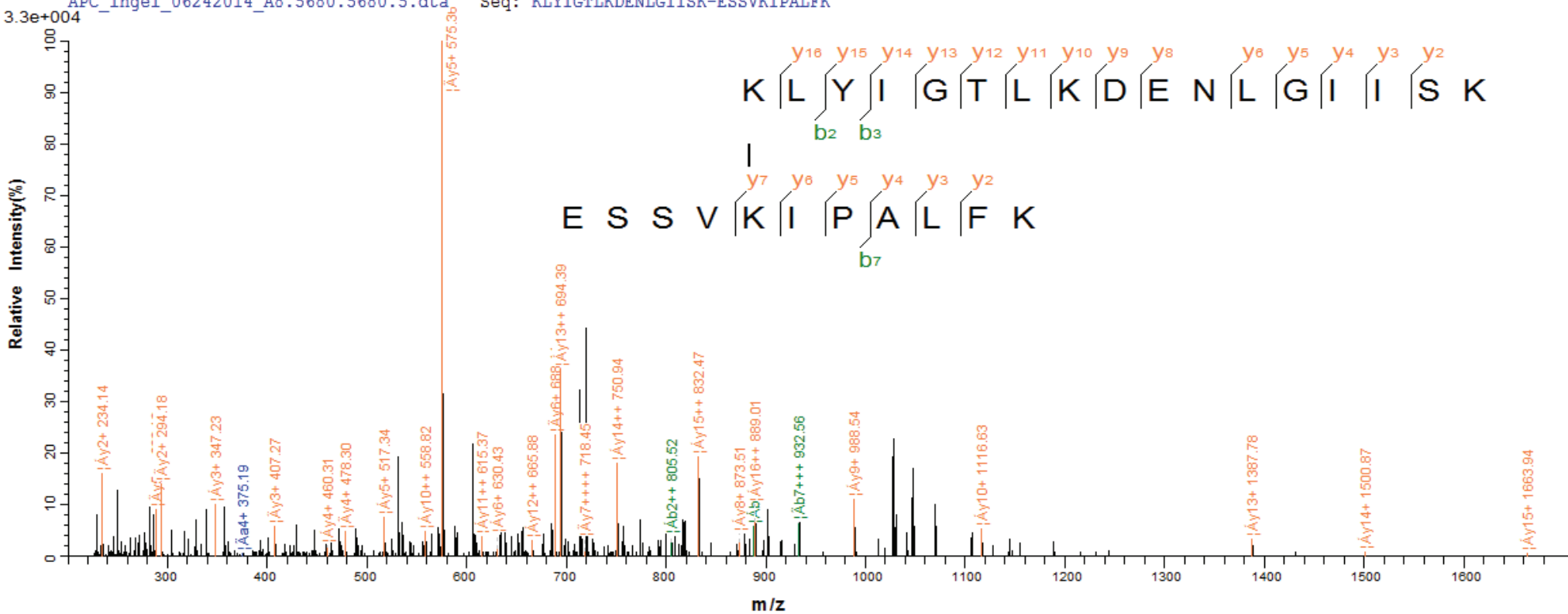






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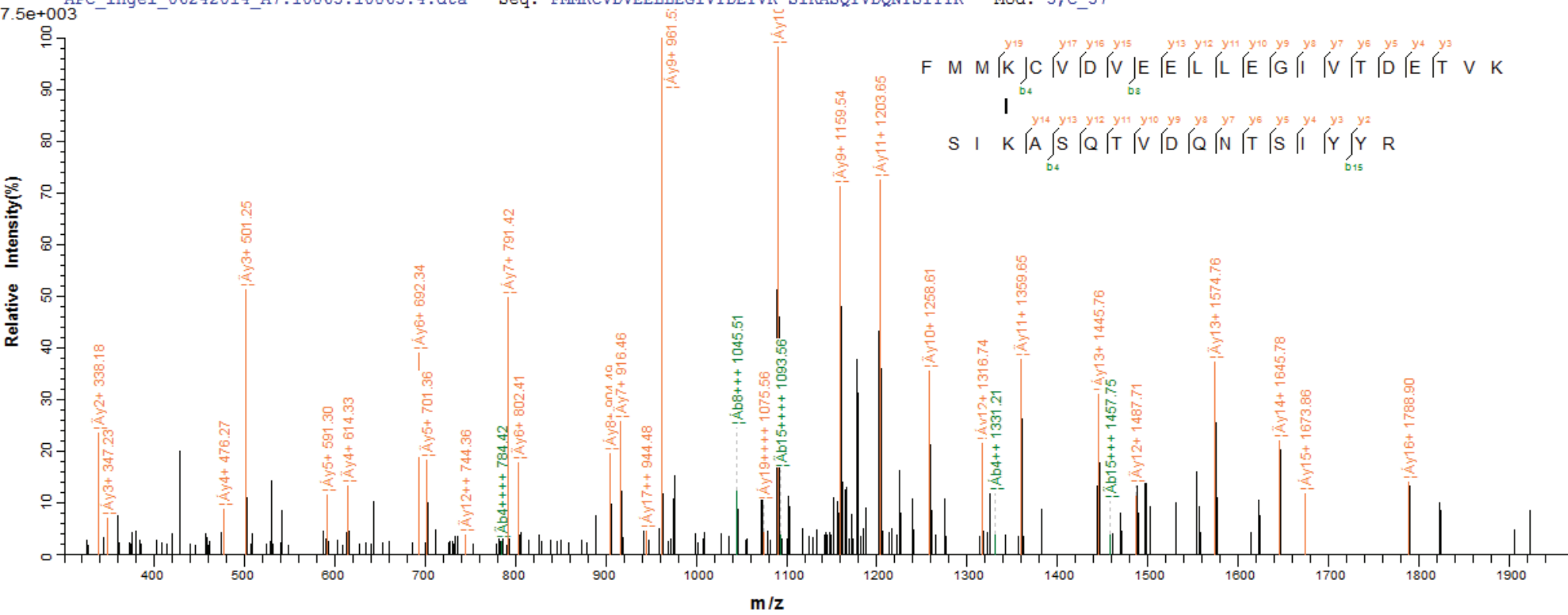
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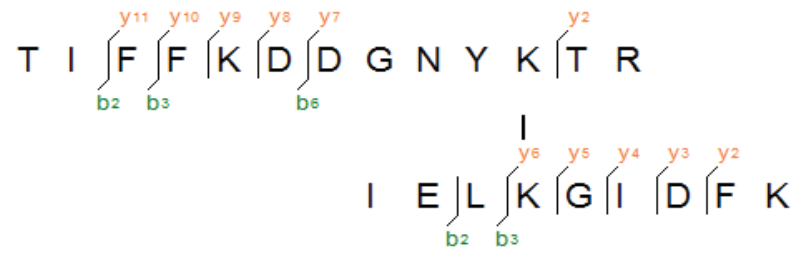
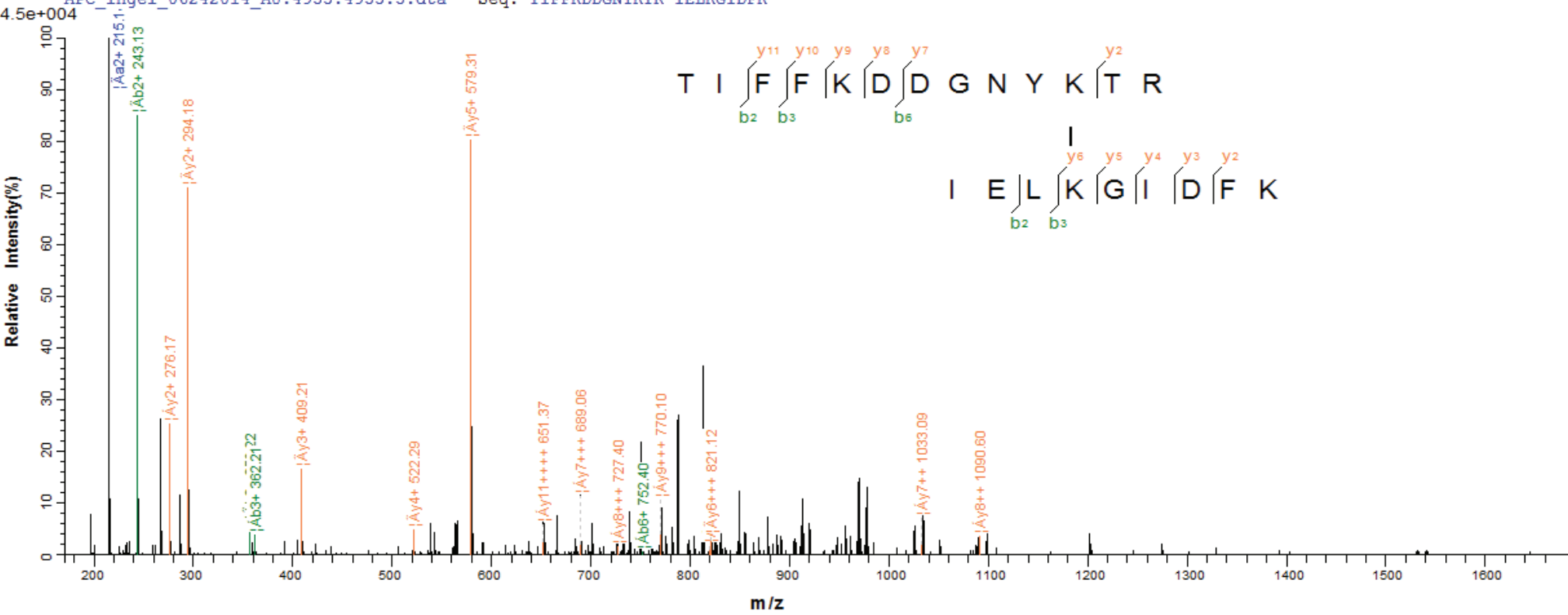
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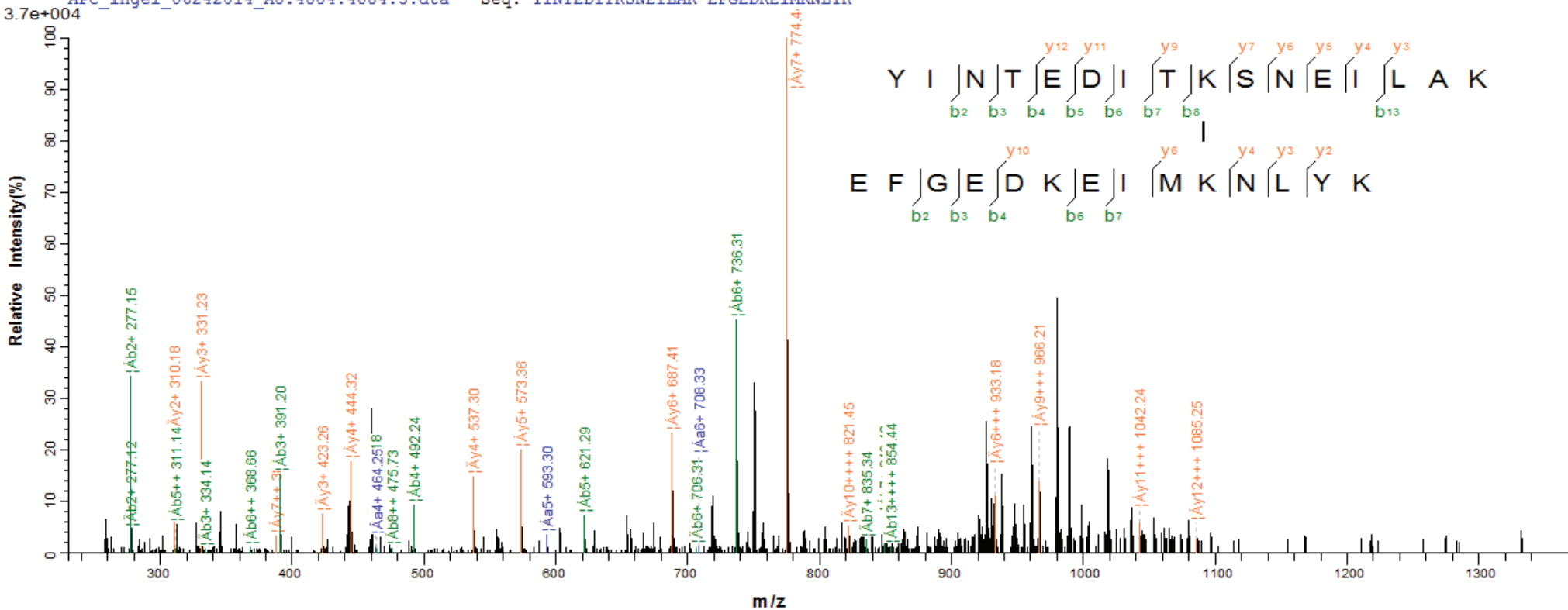
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Mod: 5,C_57

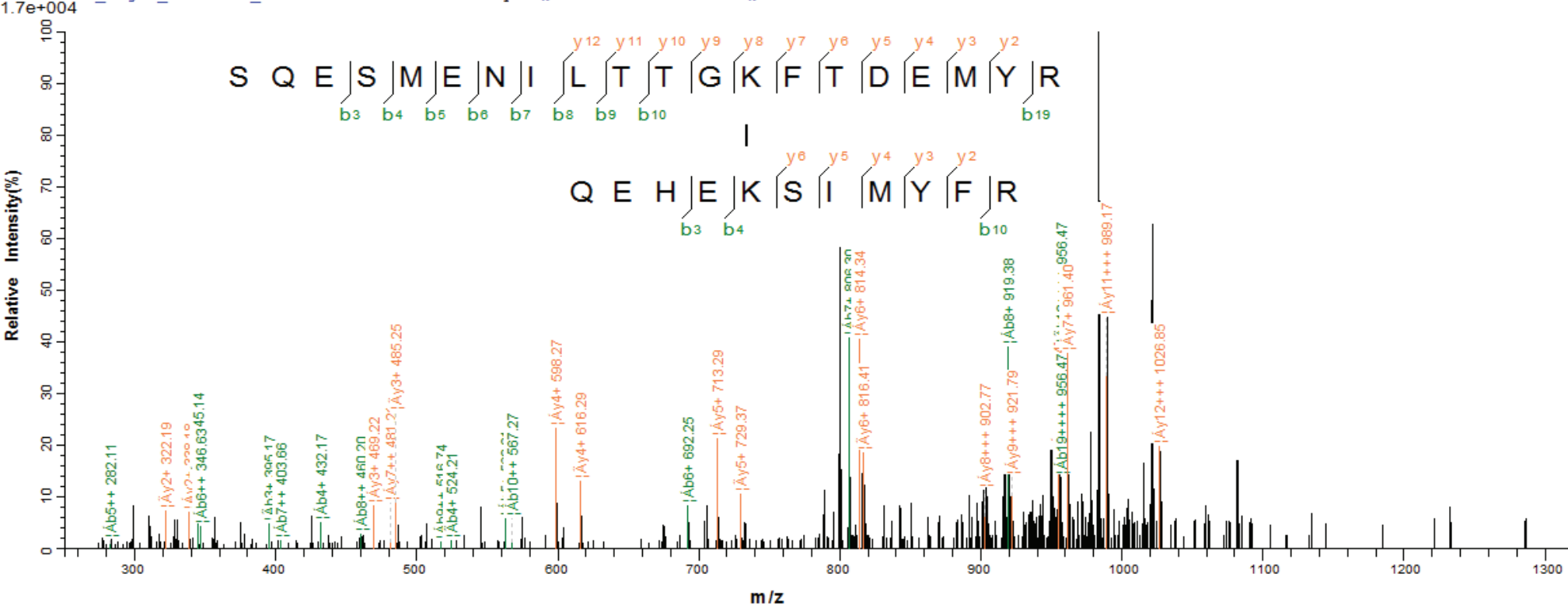


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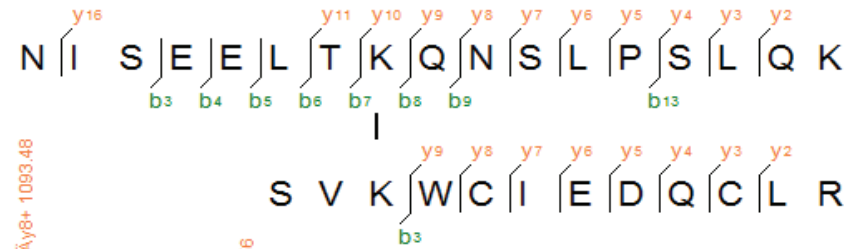
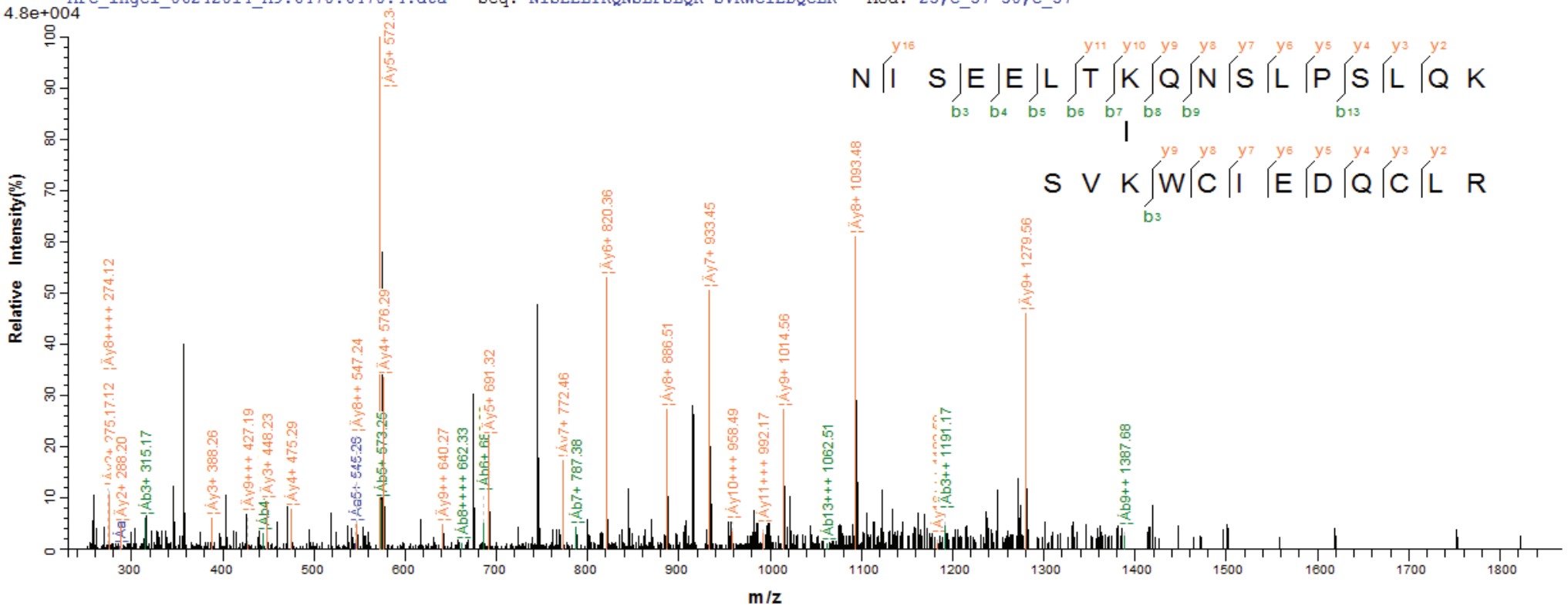
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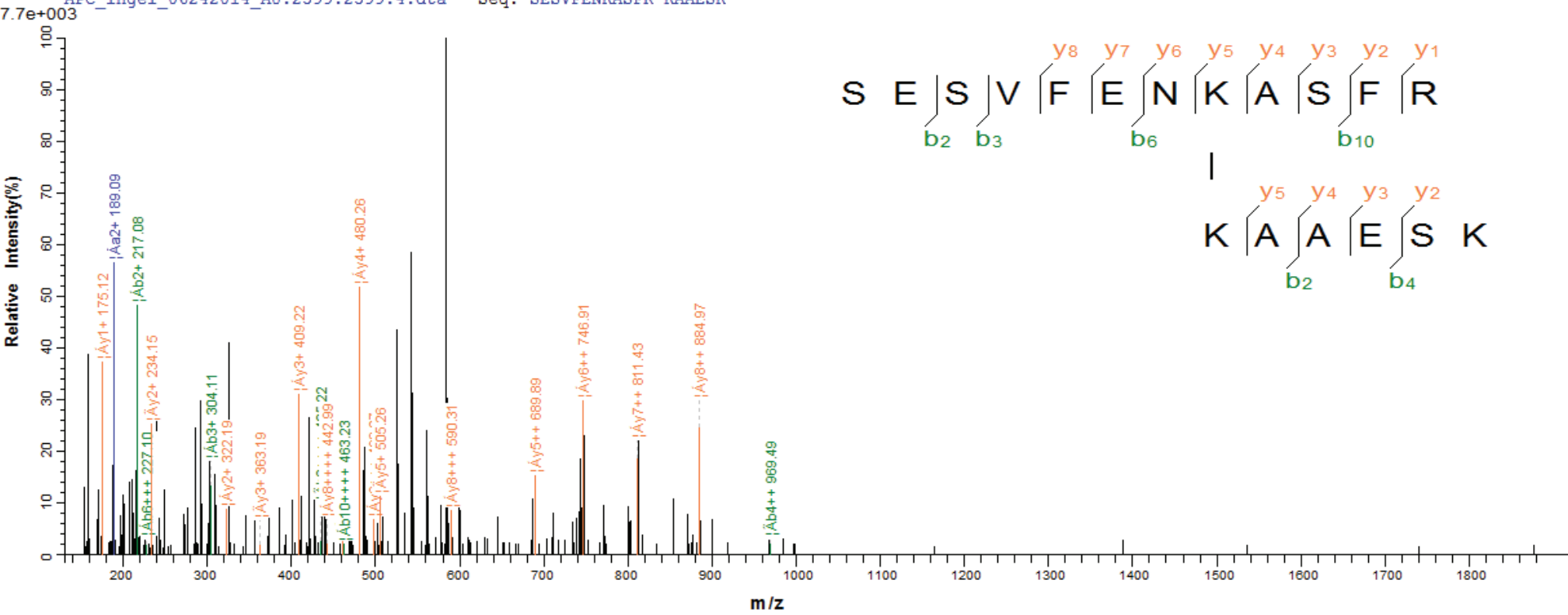


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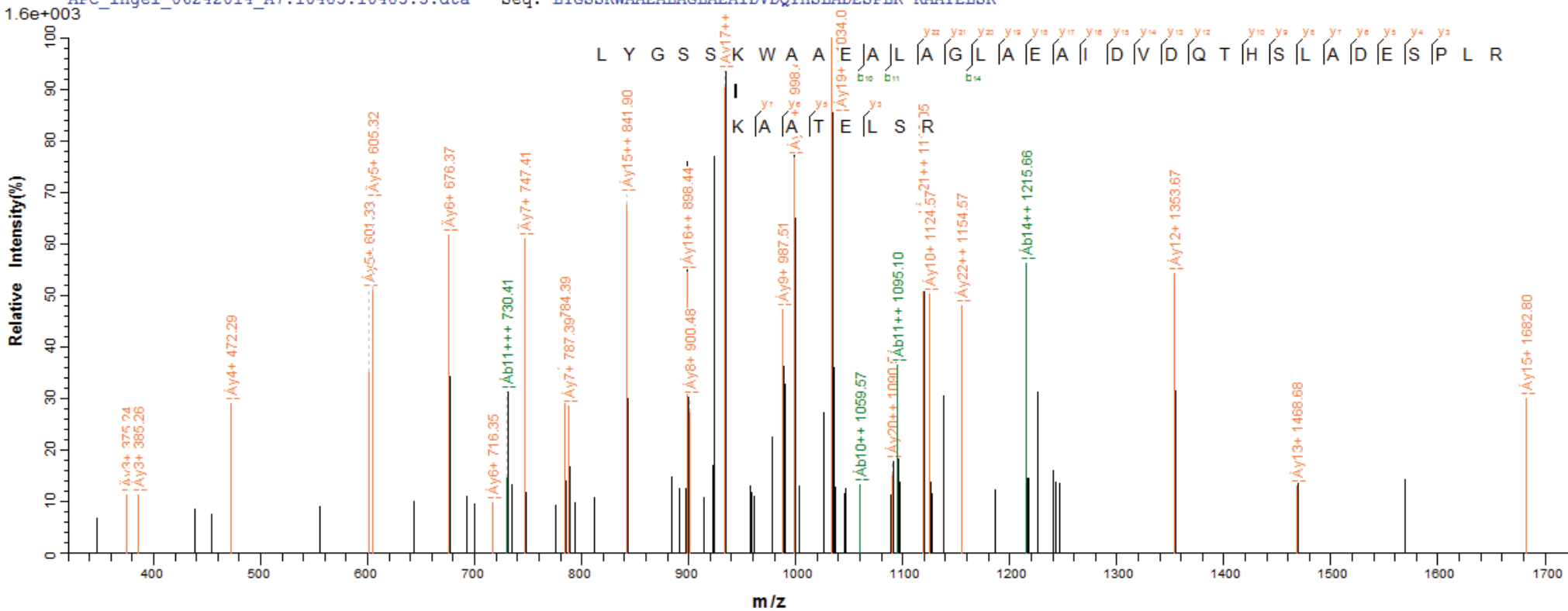
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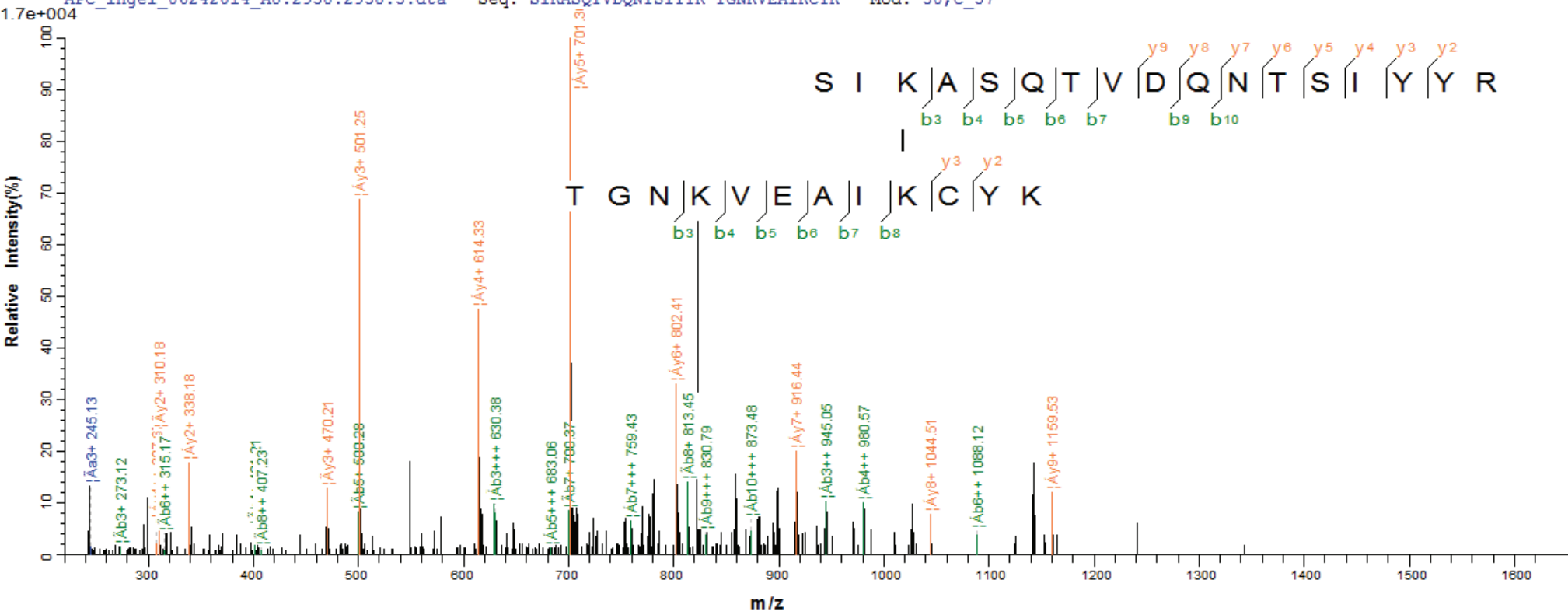




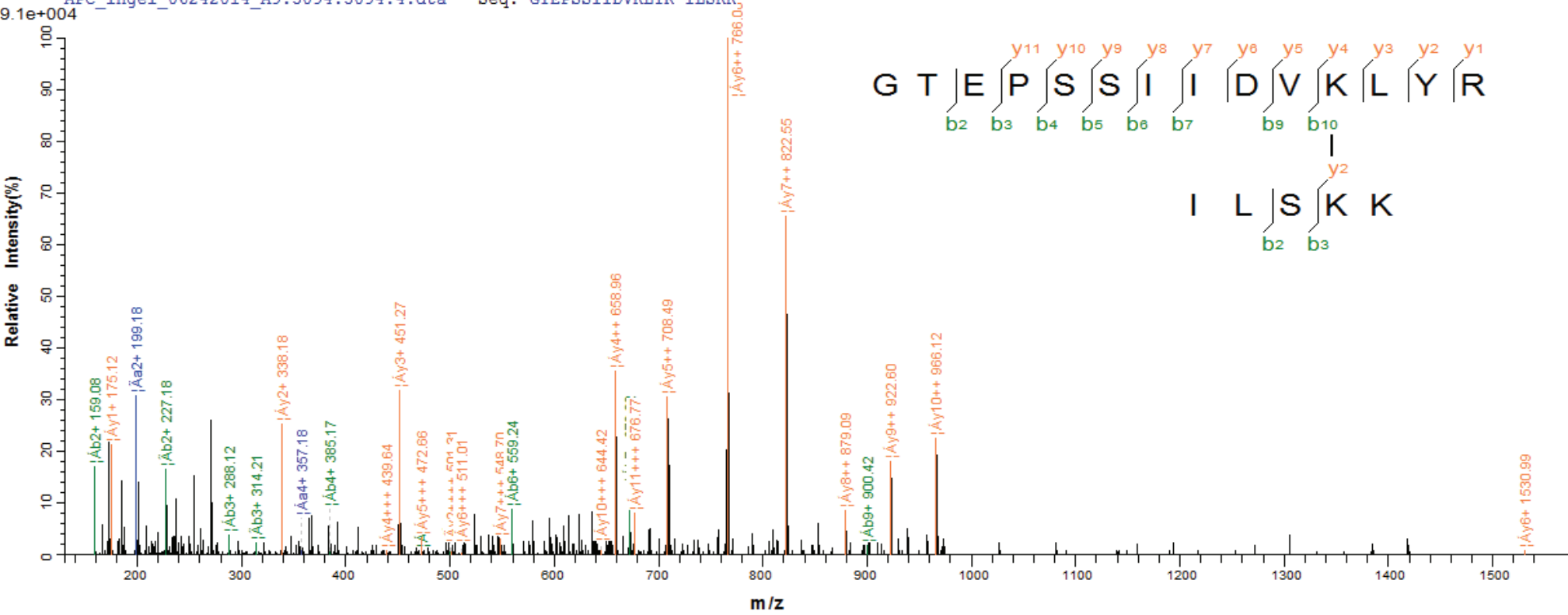
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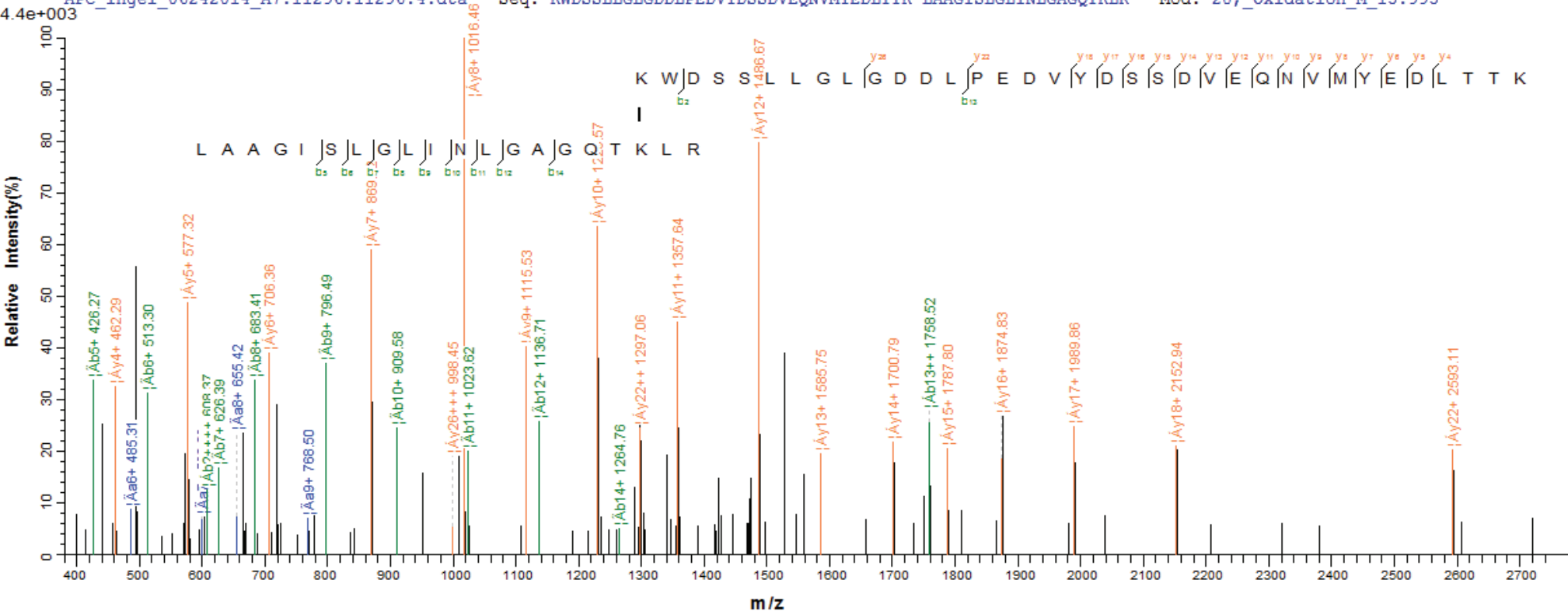
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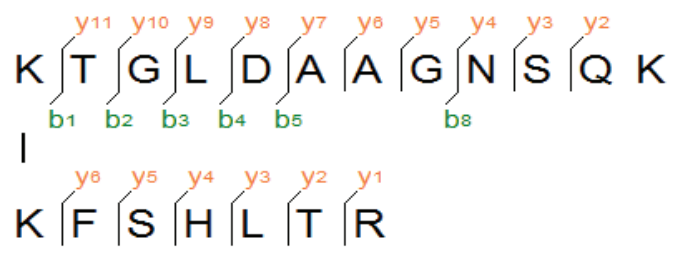
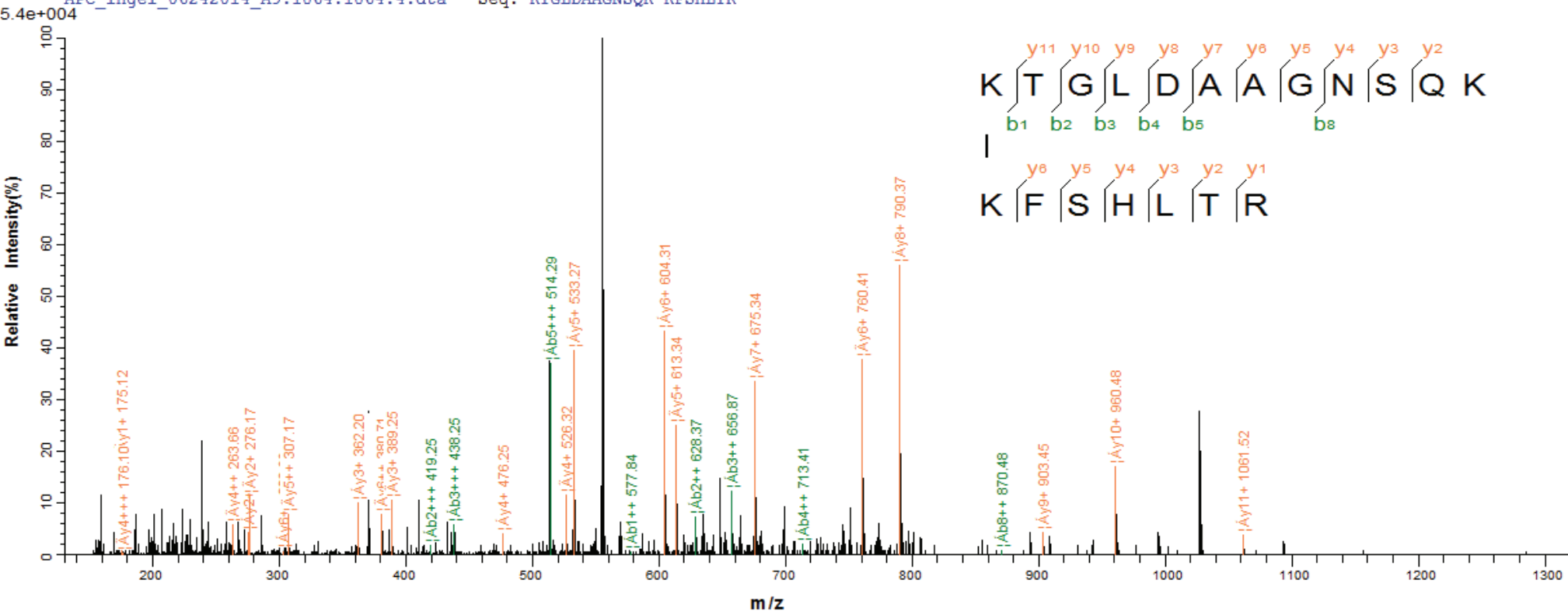


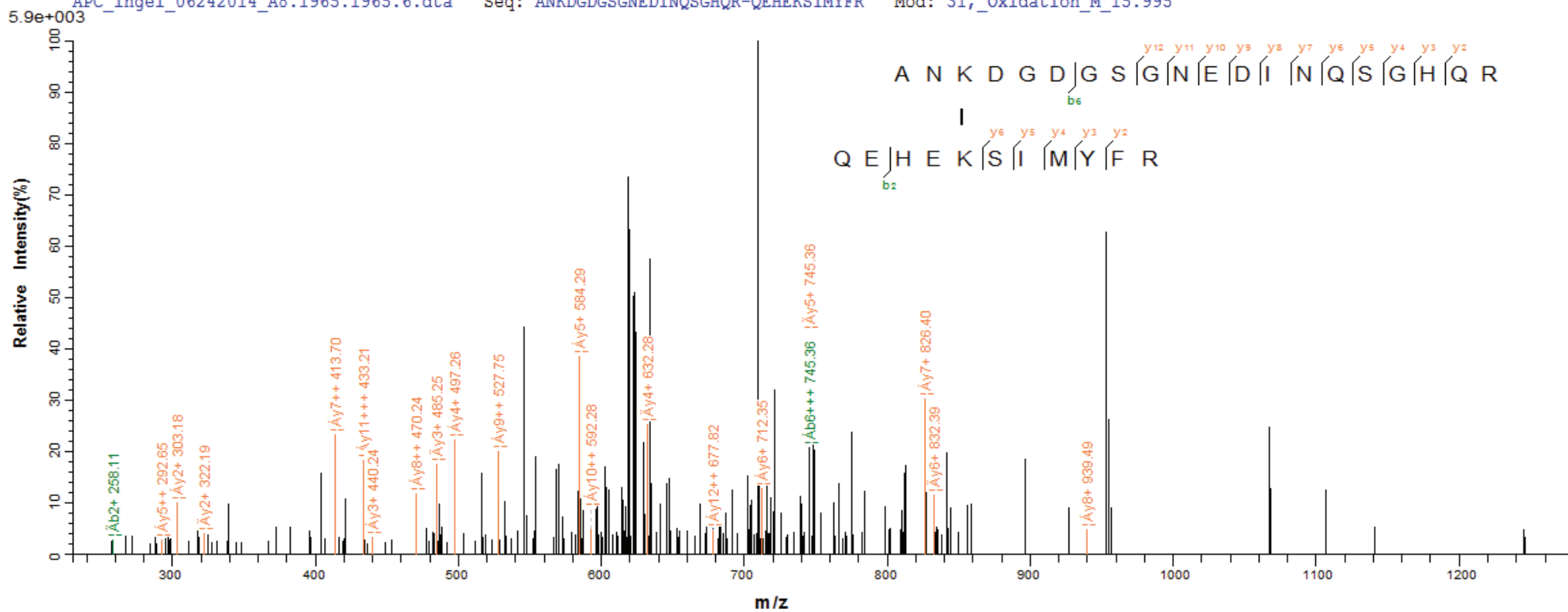
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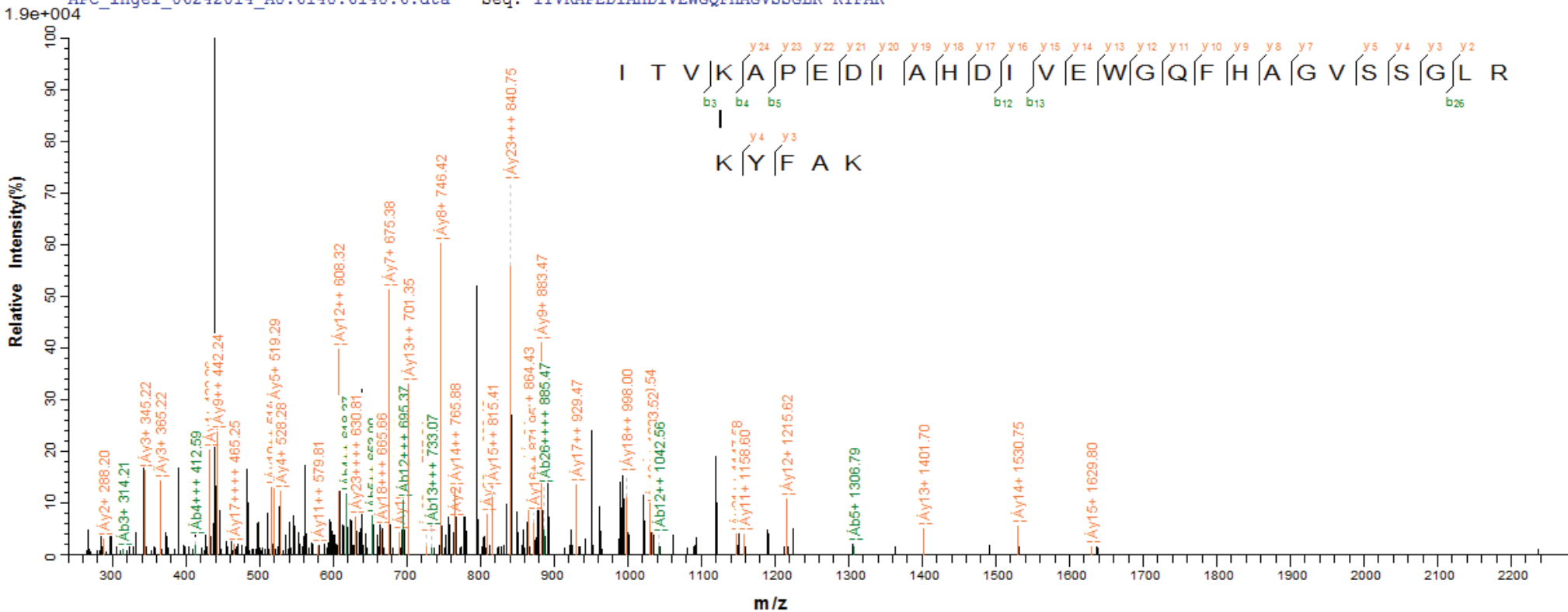






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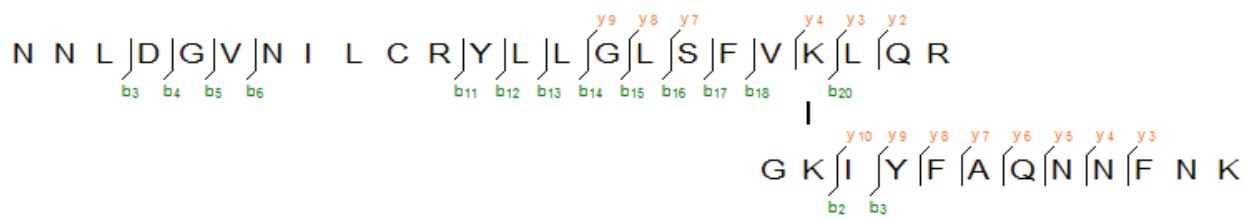
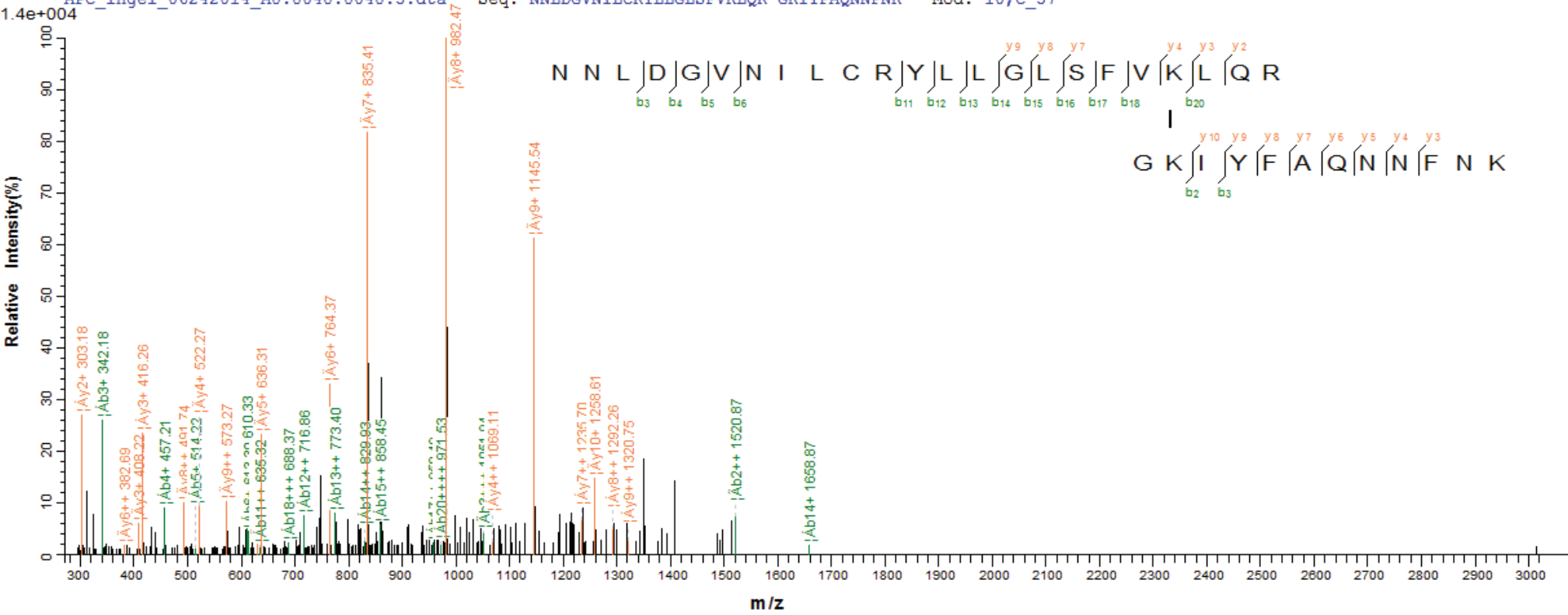
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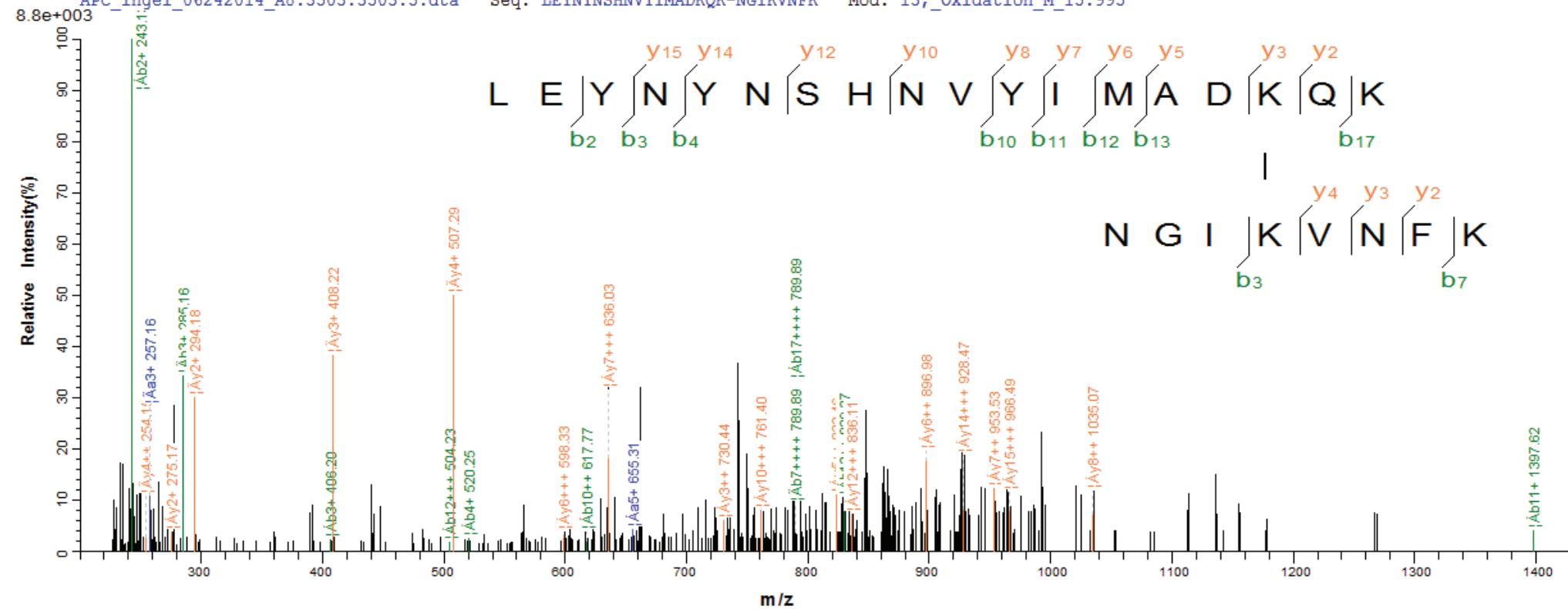
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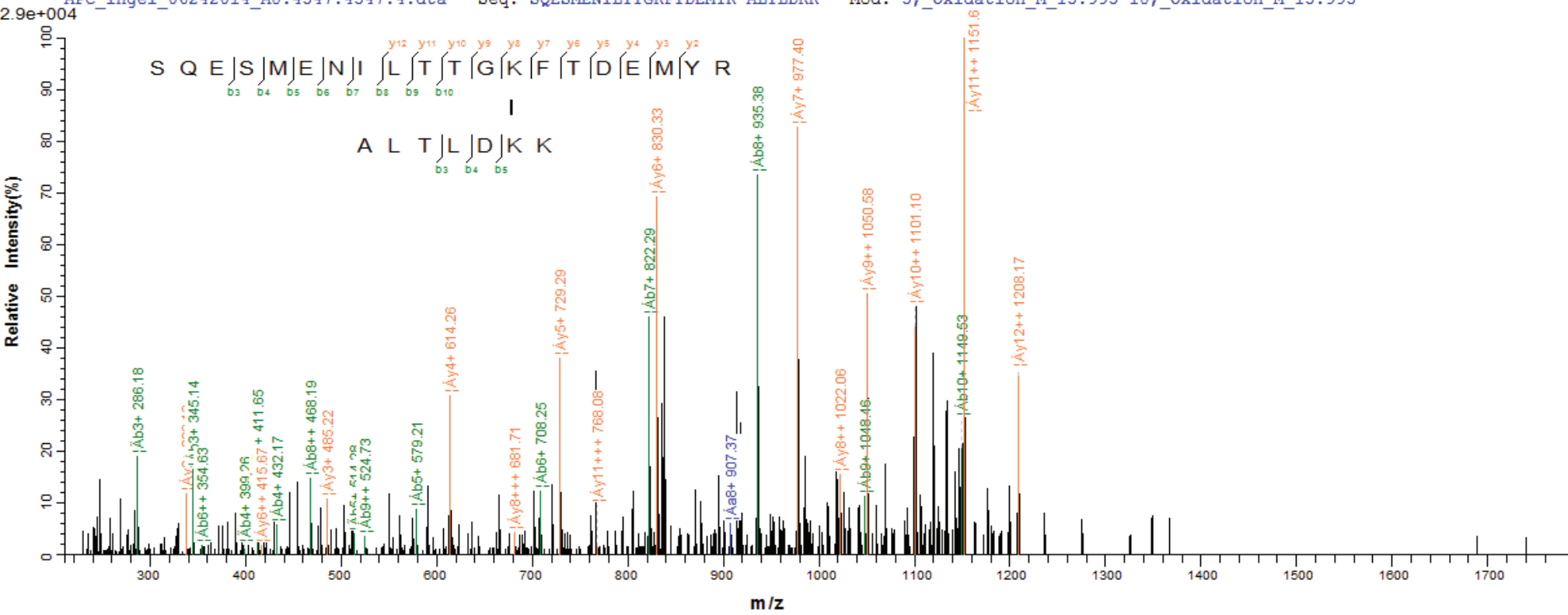


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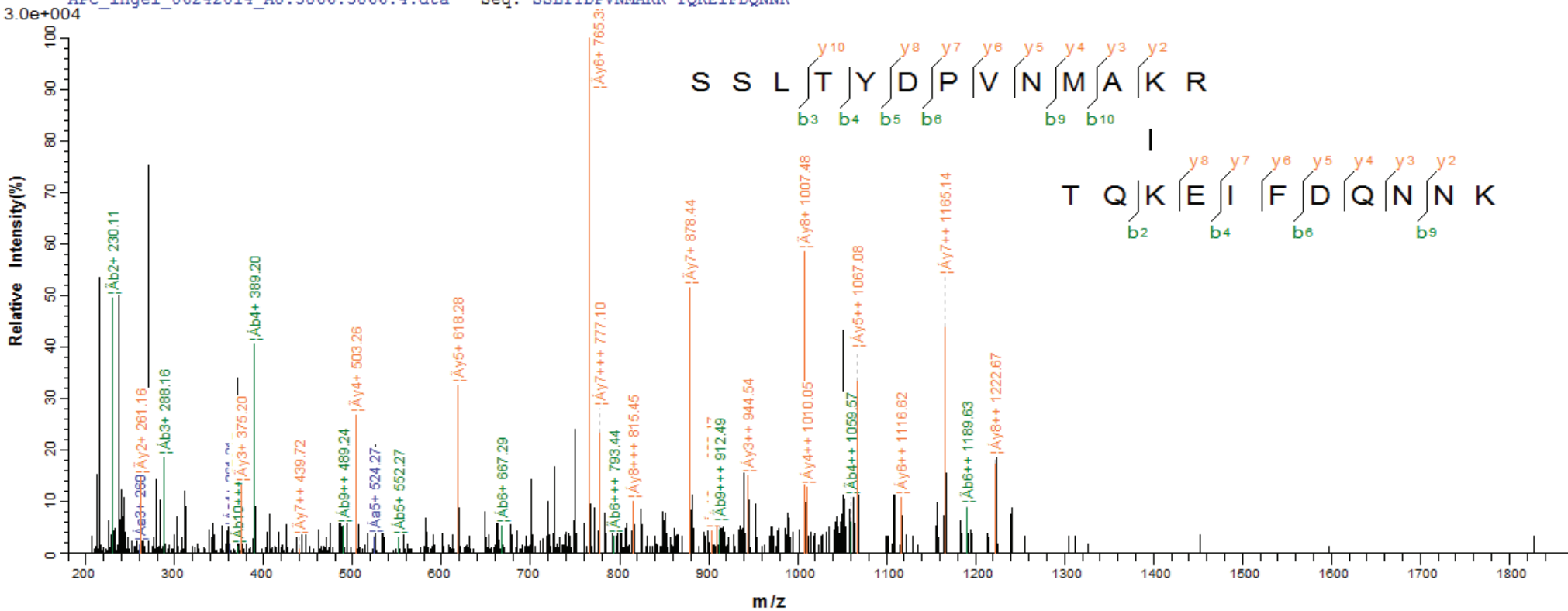
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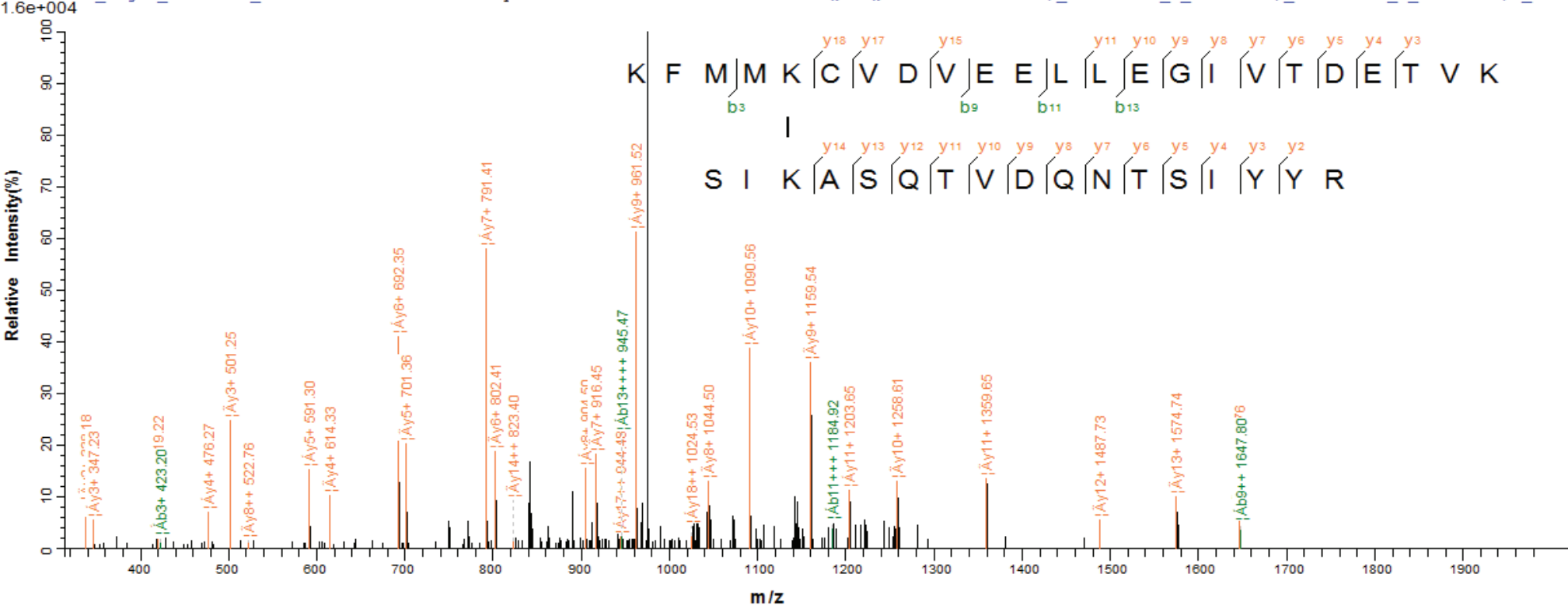
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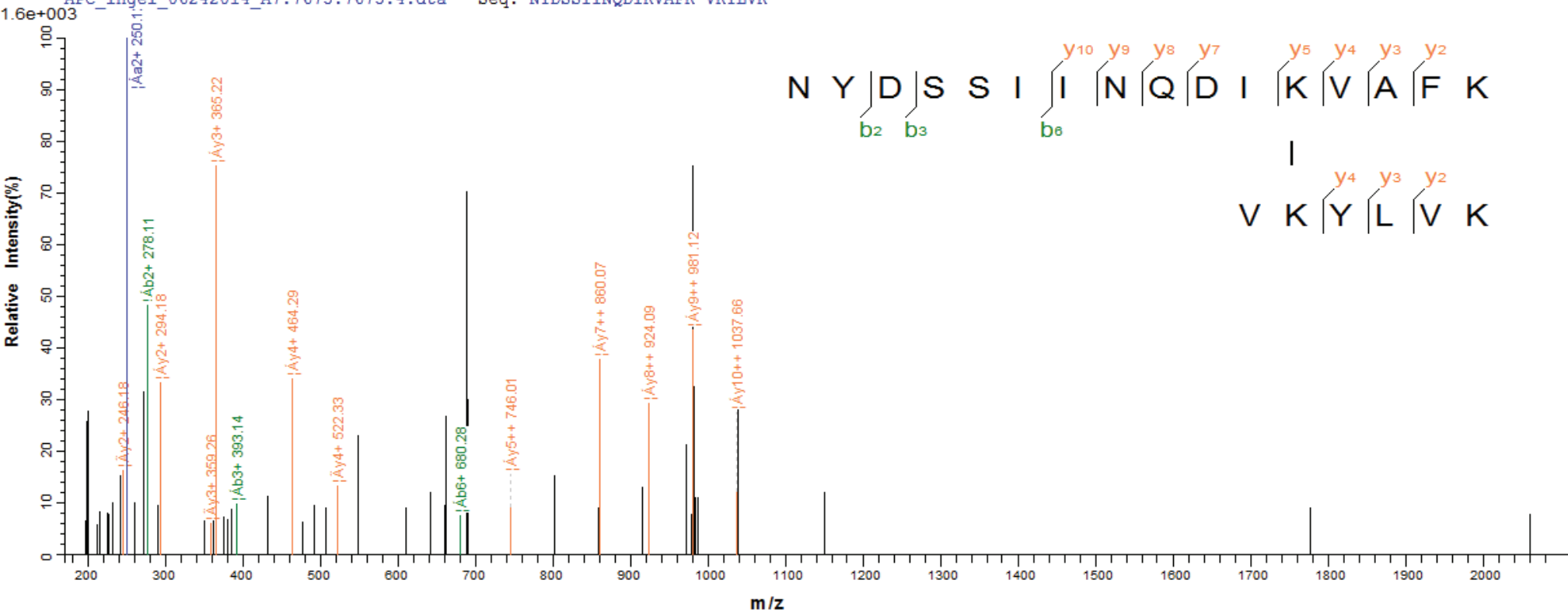




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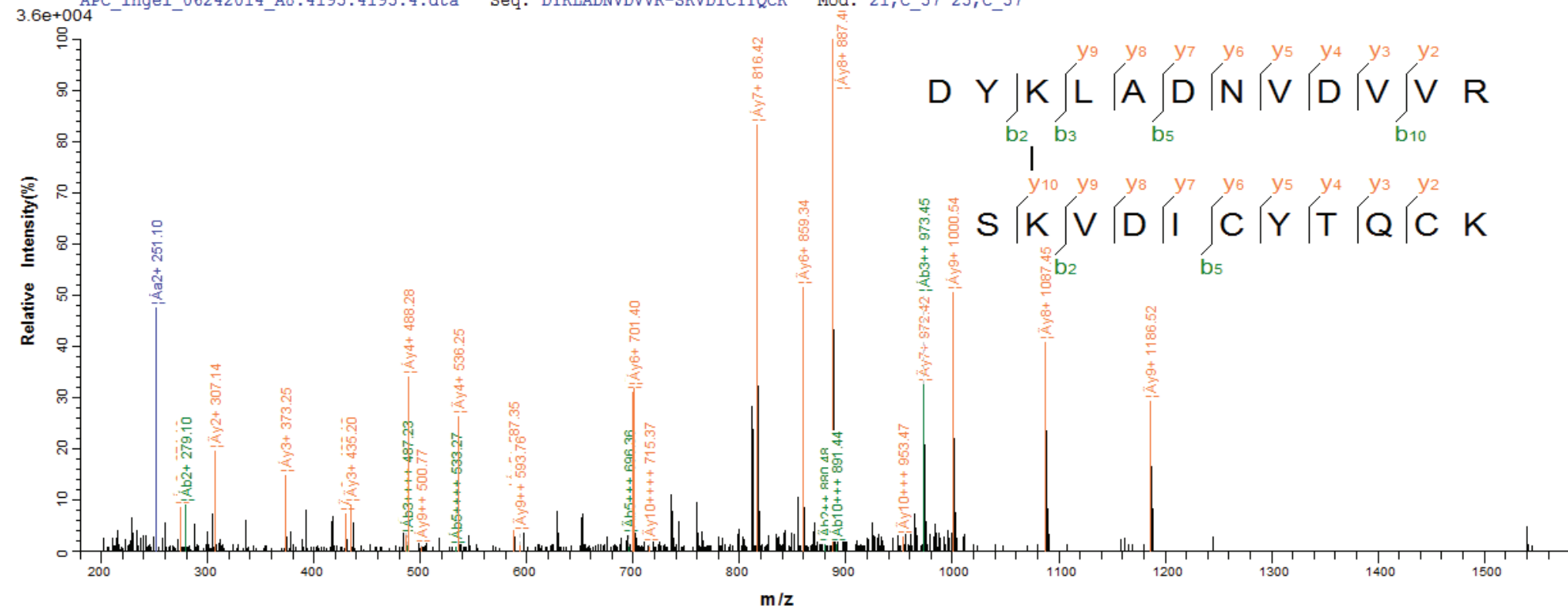




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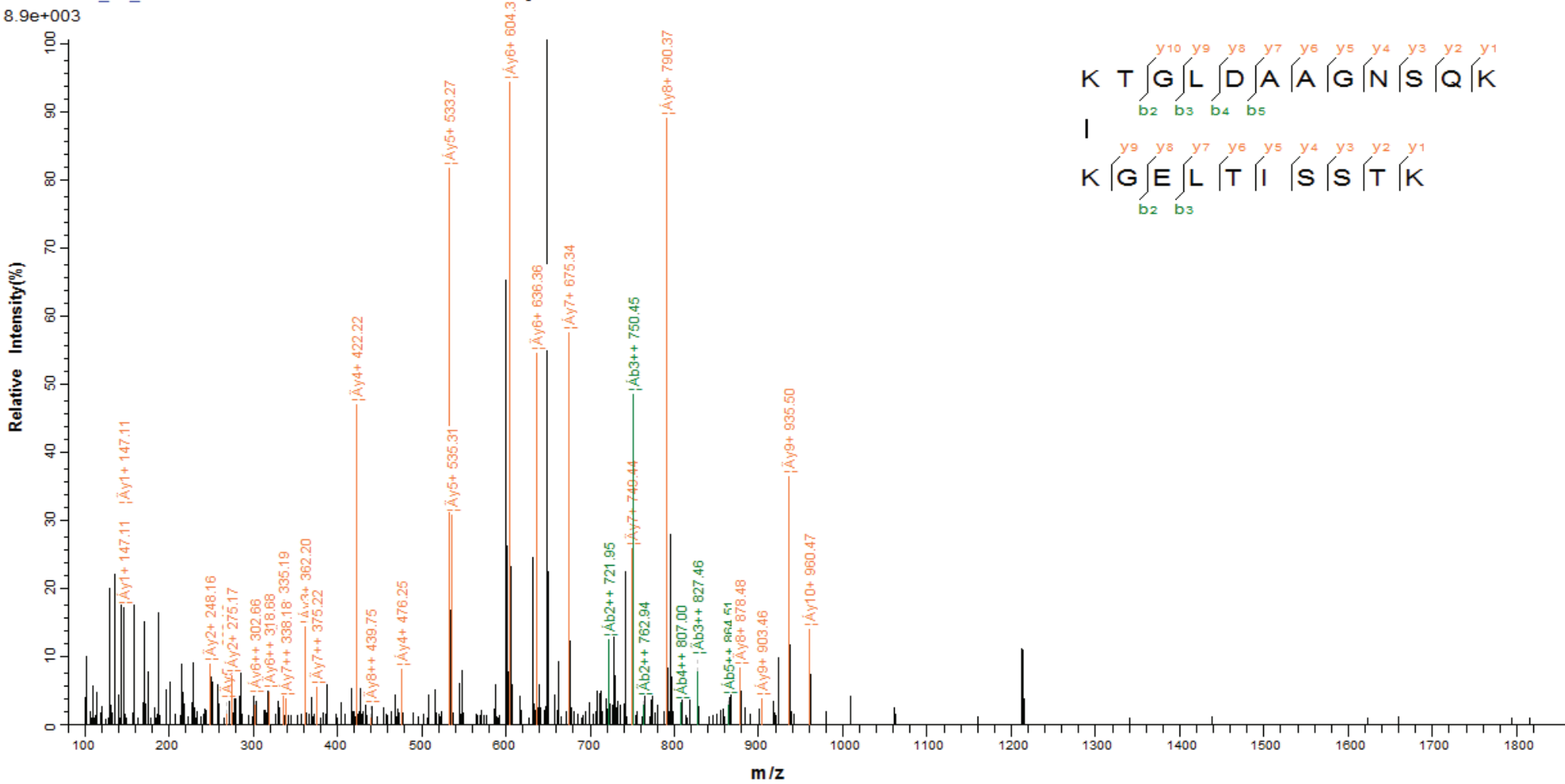
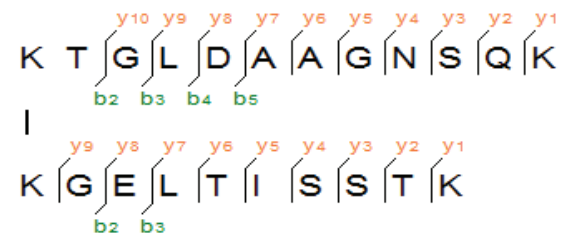
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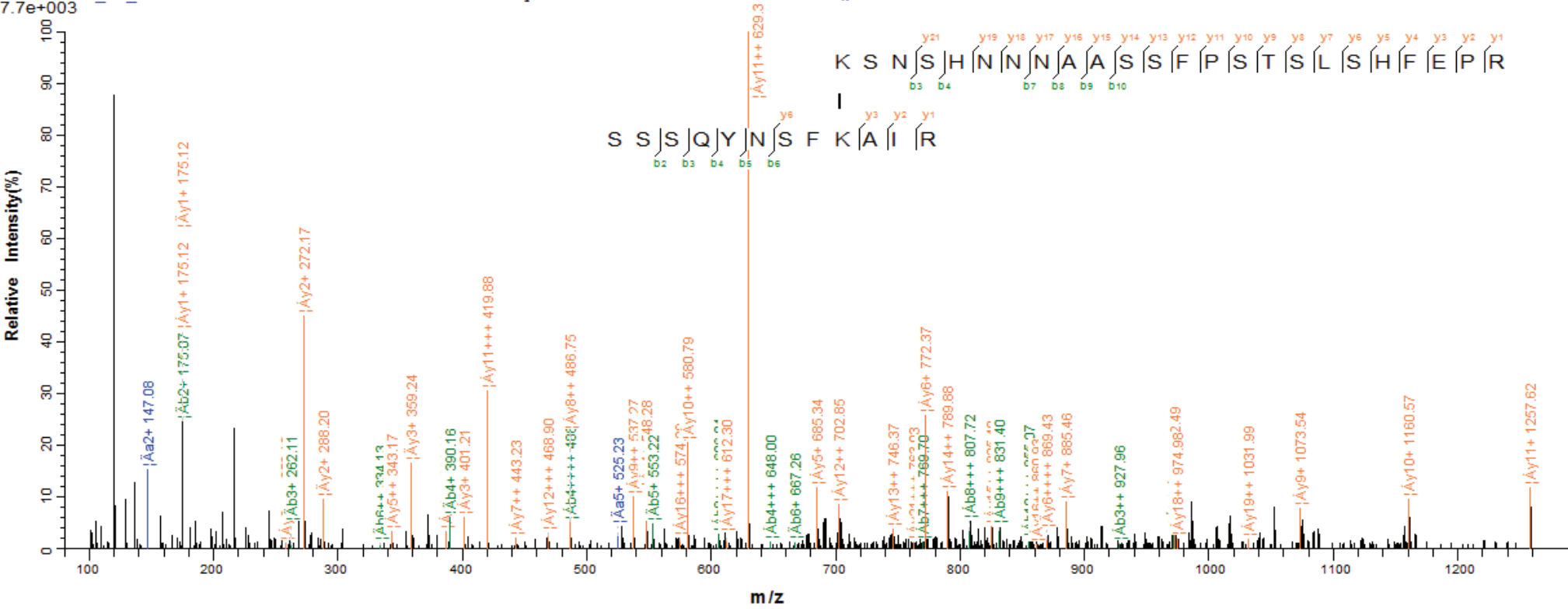
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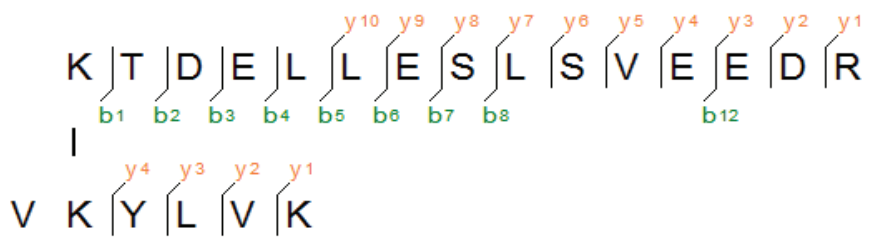
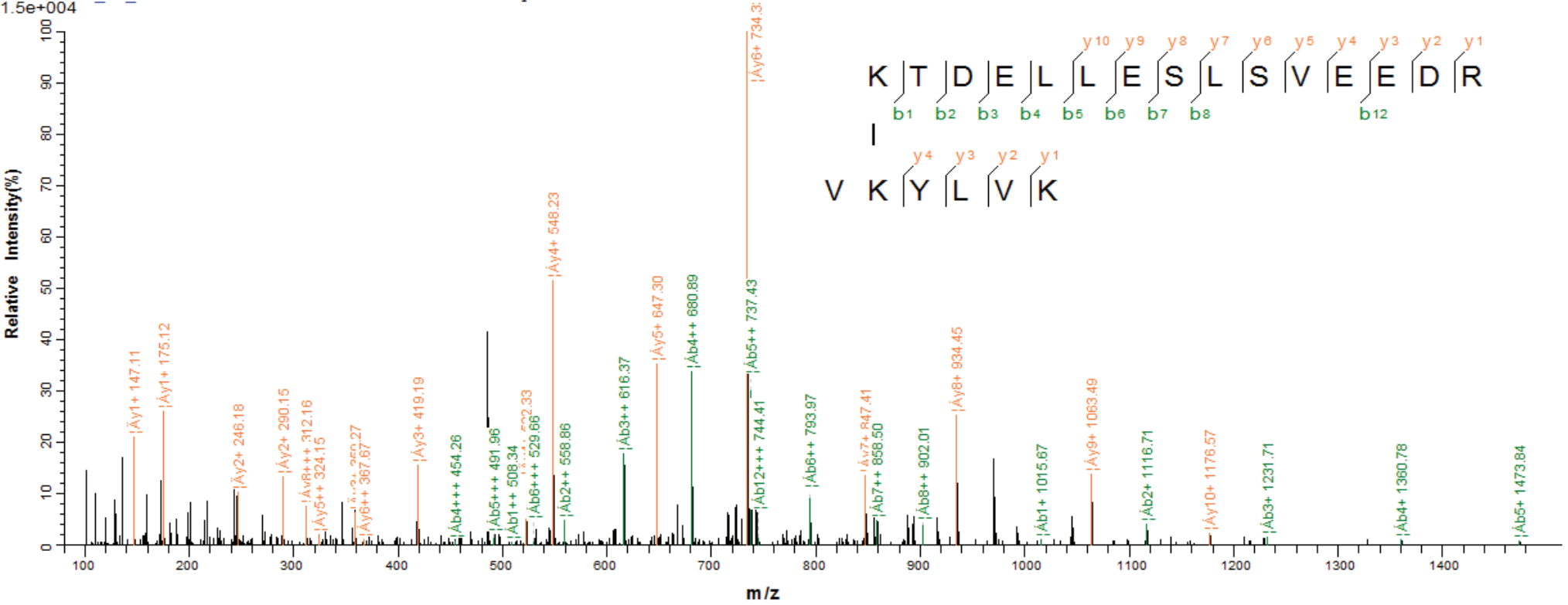


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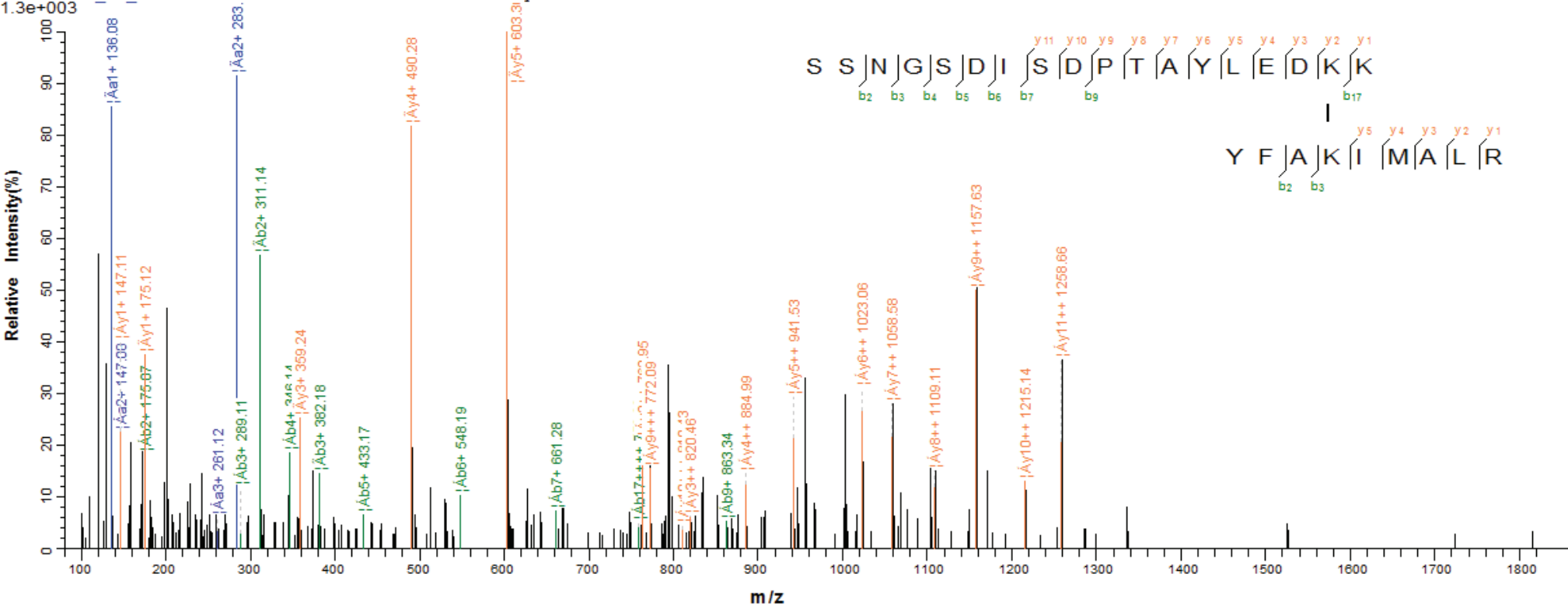






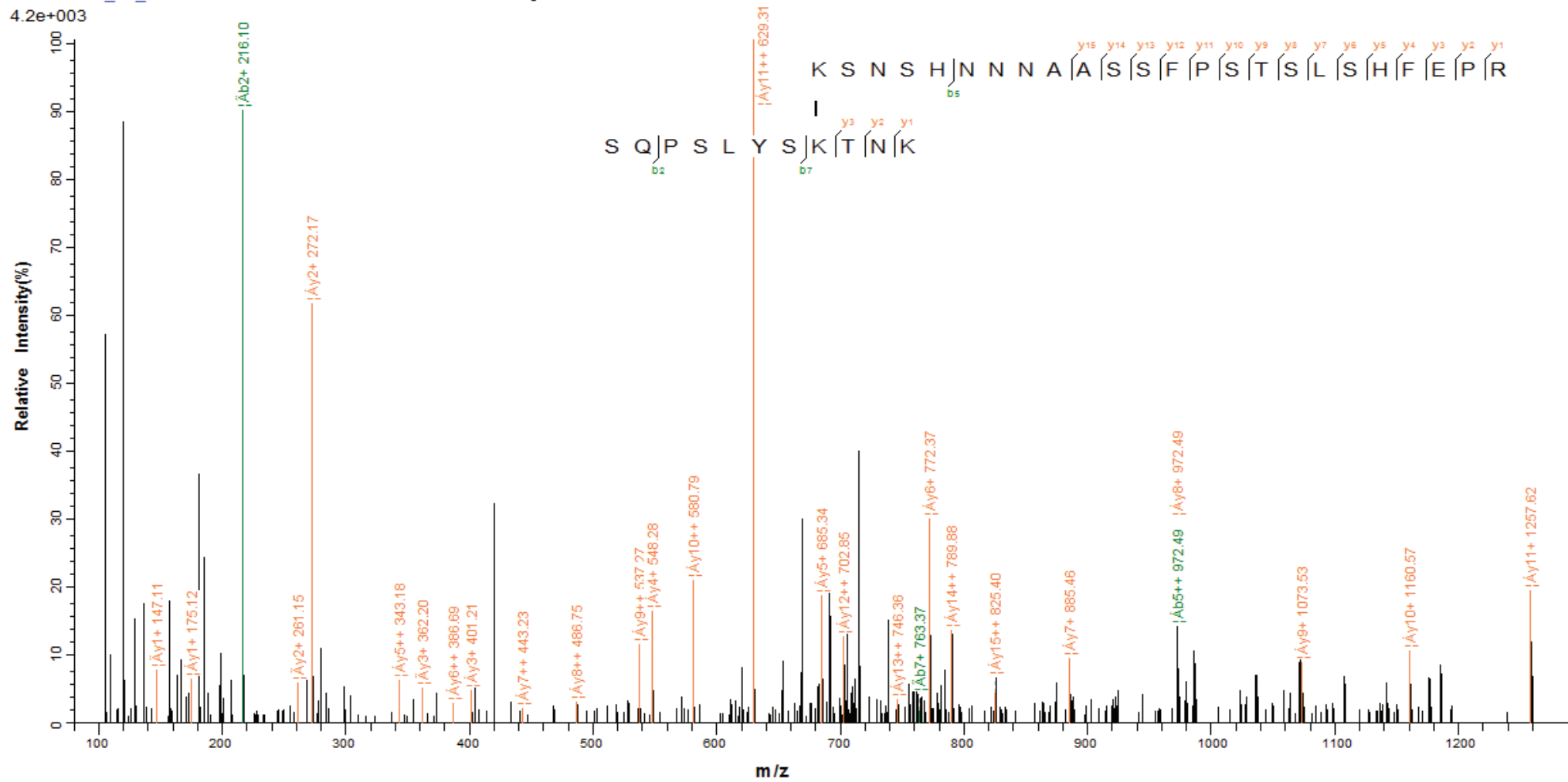
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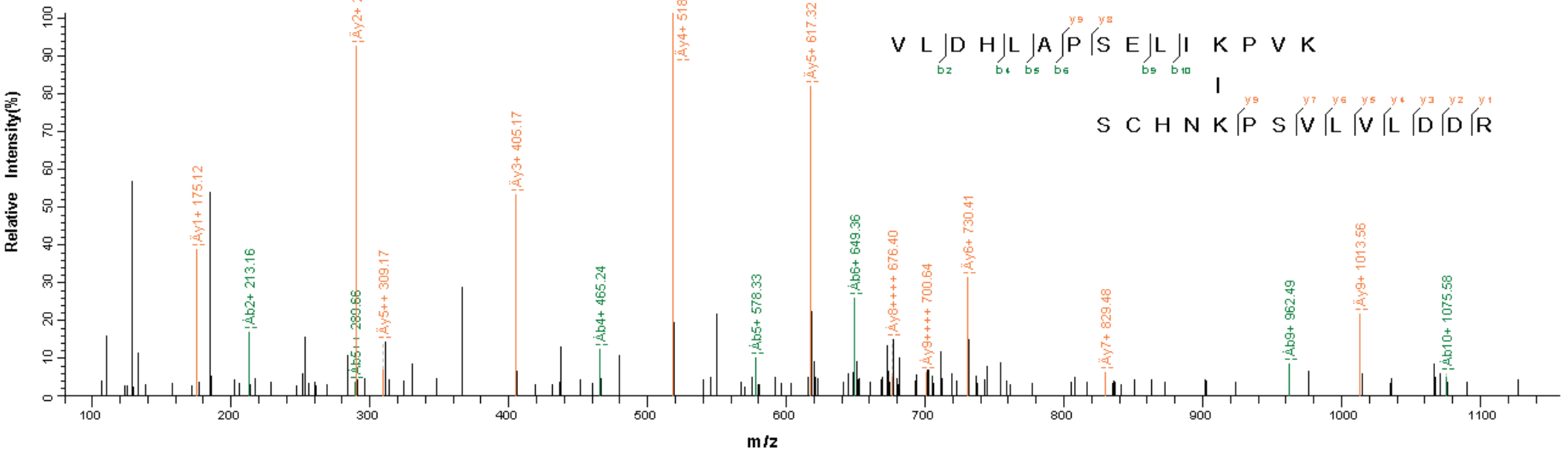
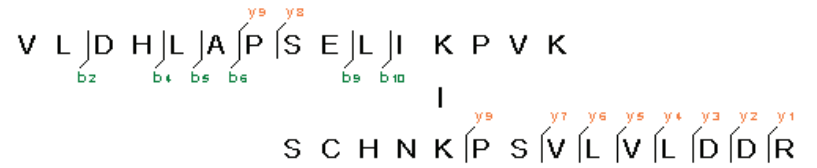


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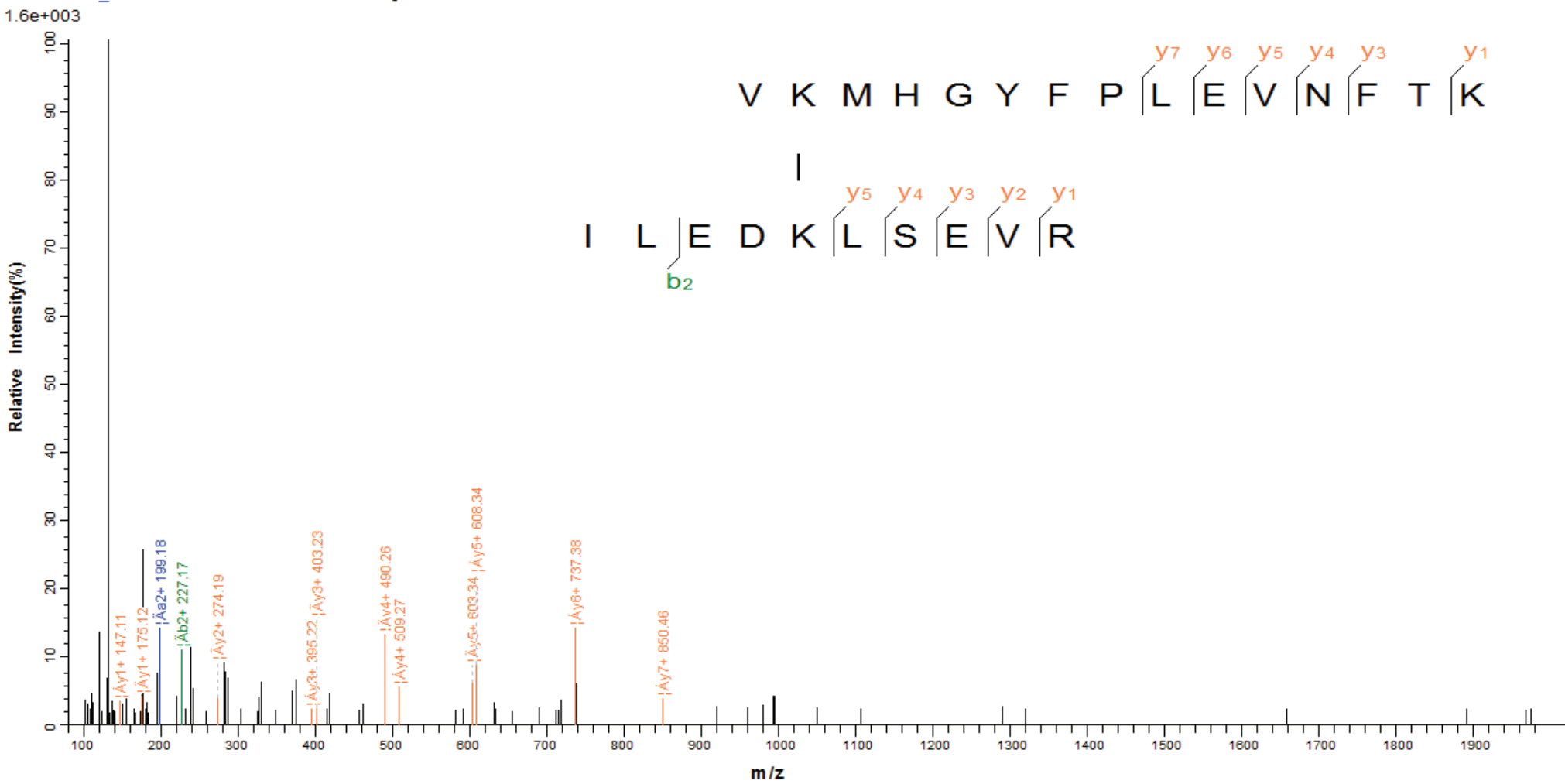
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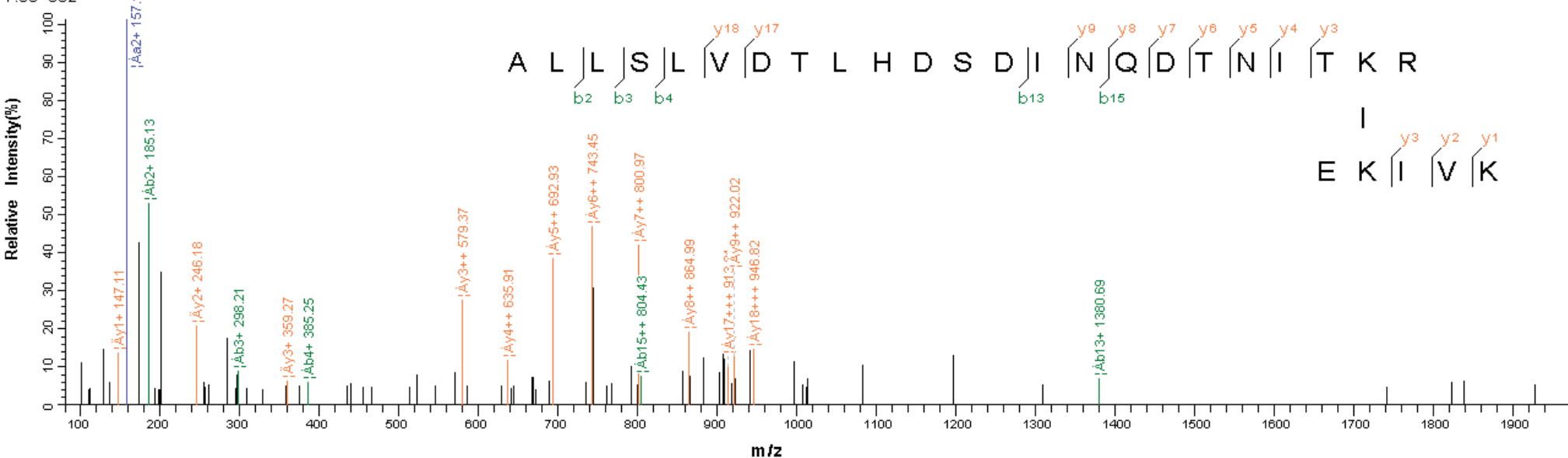
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Charge: 6 Score: 2.50e-007

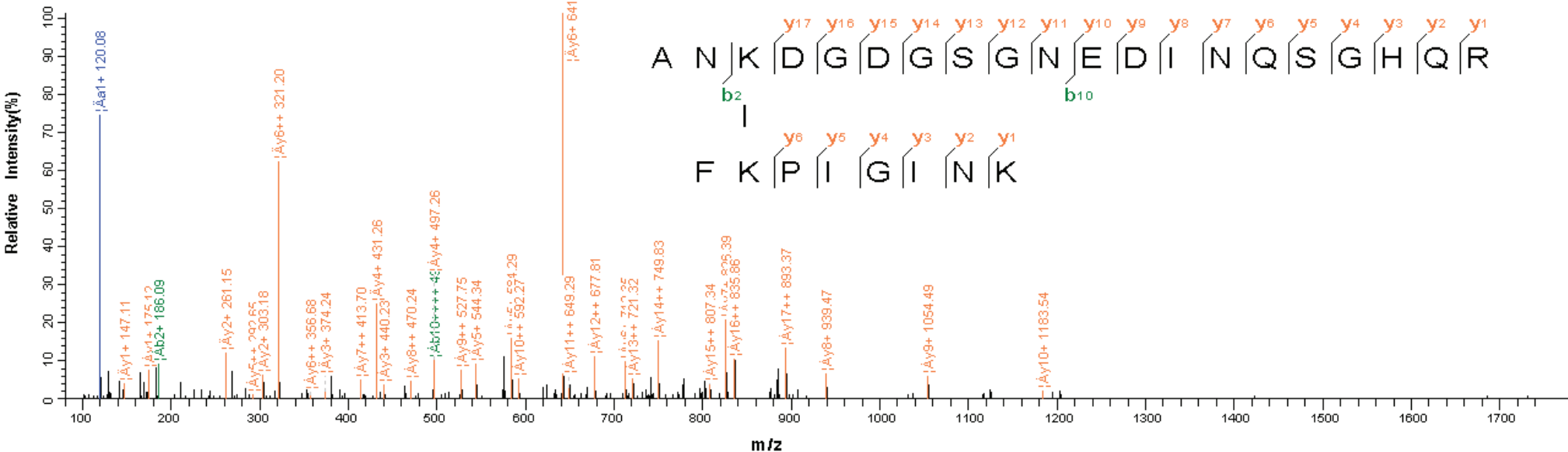


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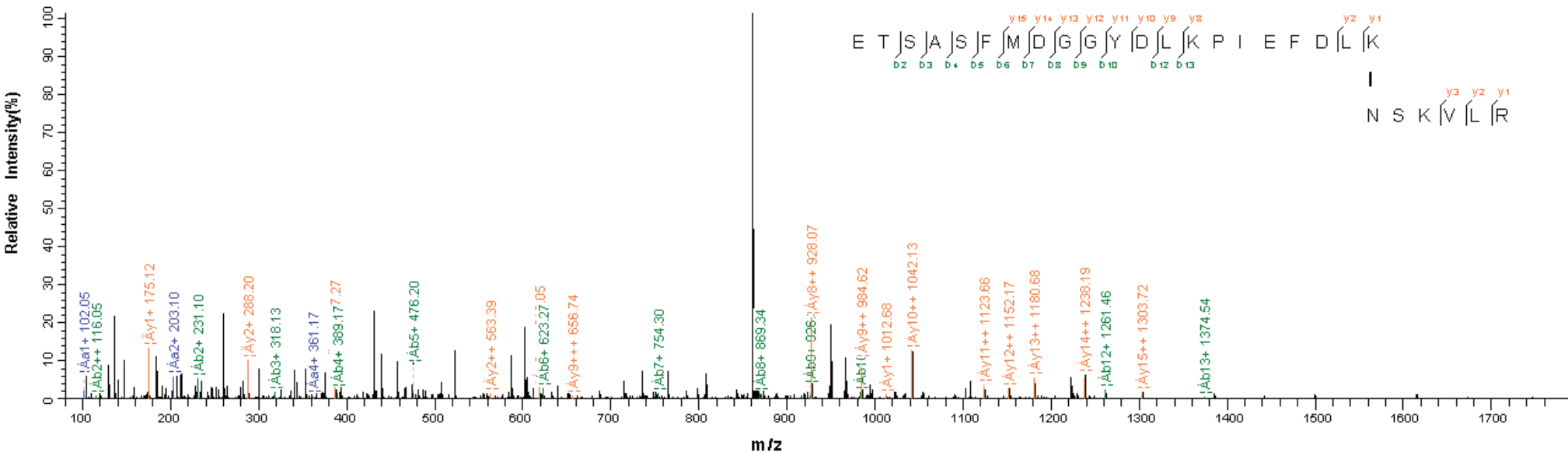




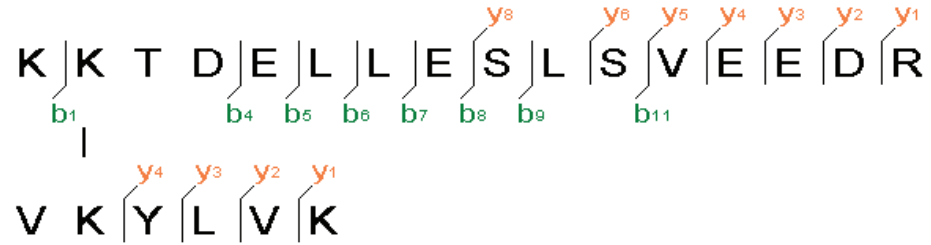
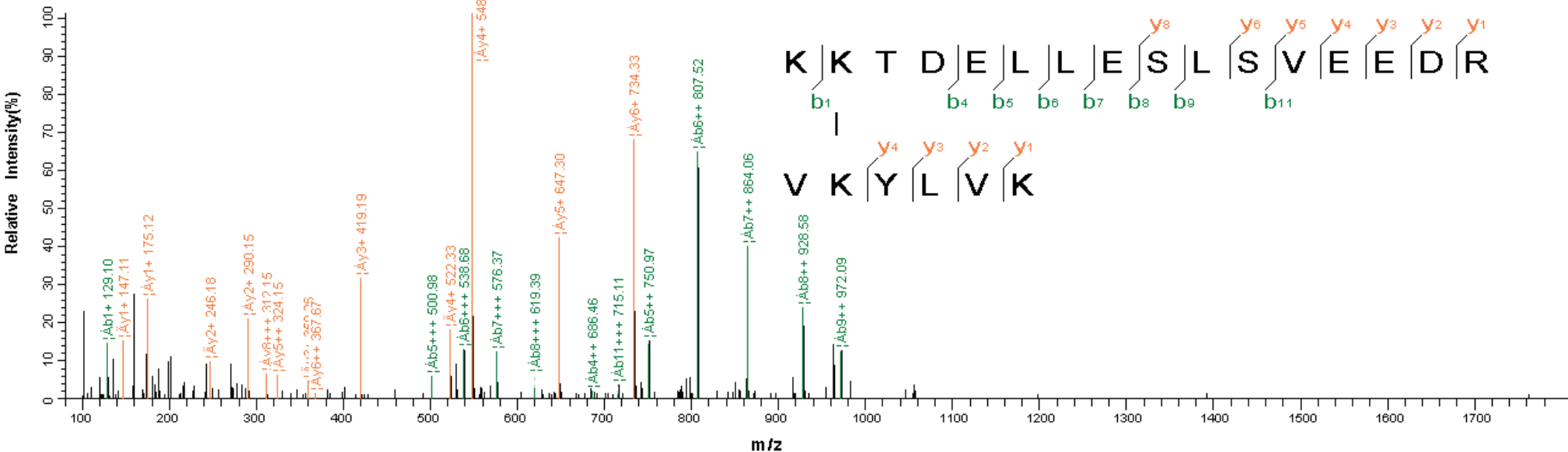
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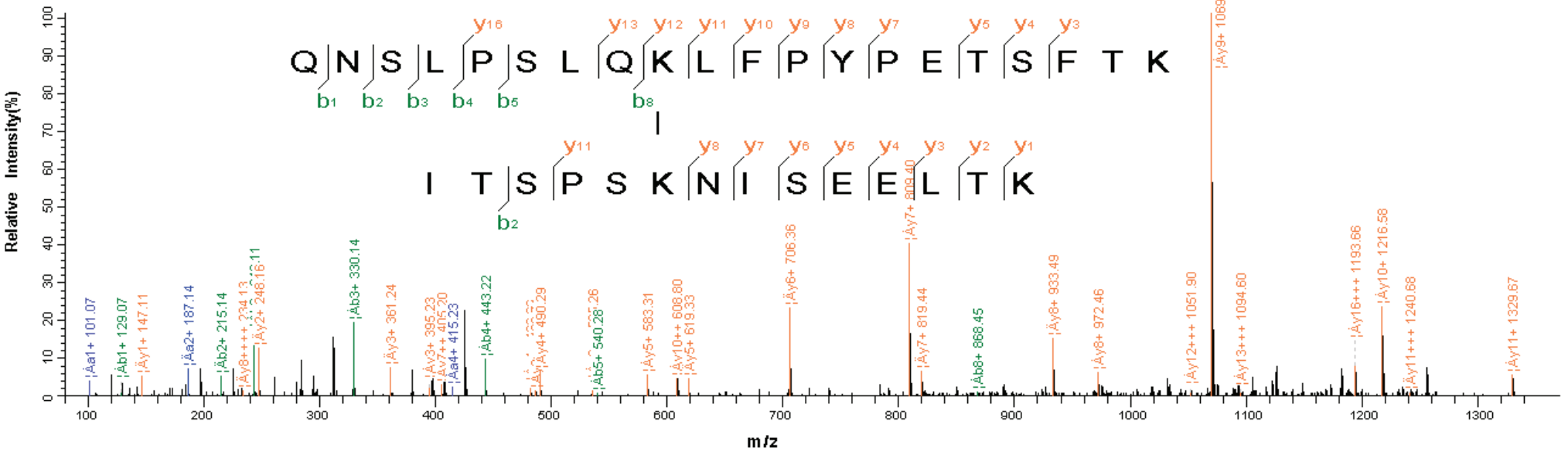
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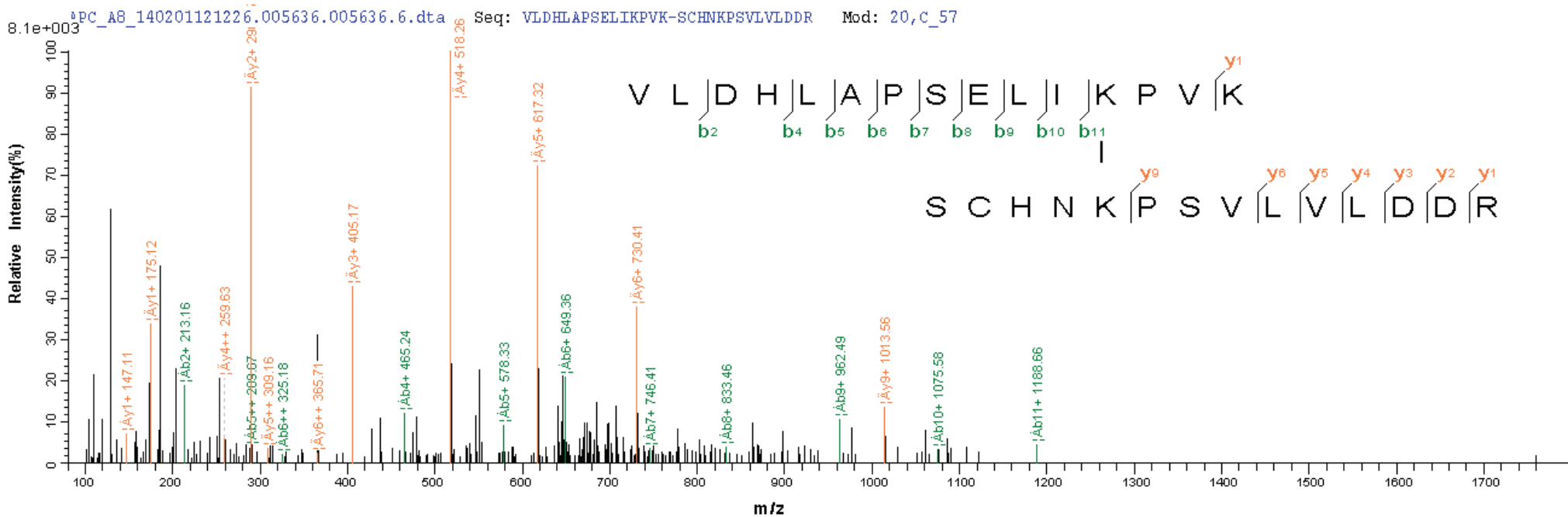
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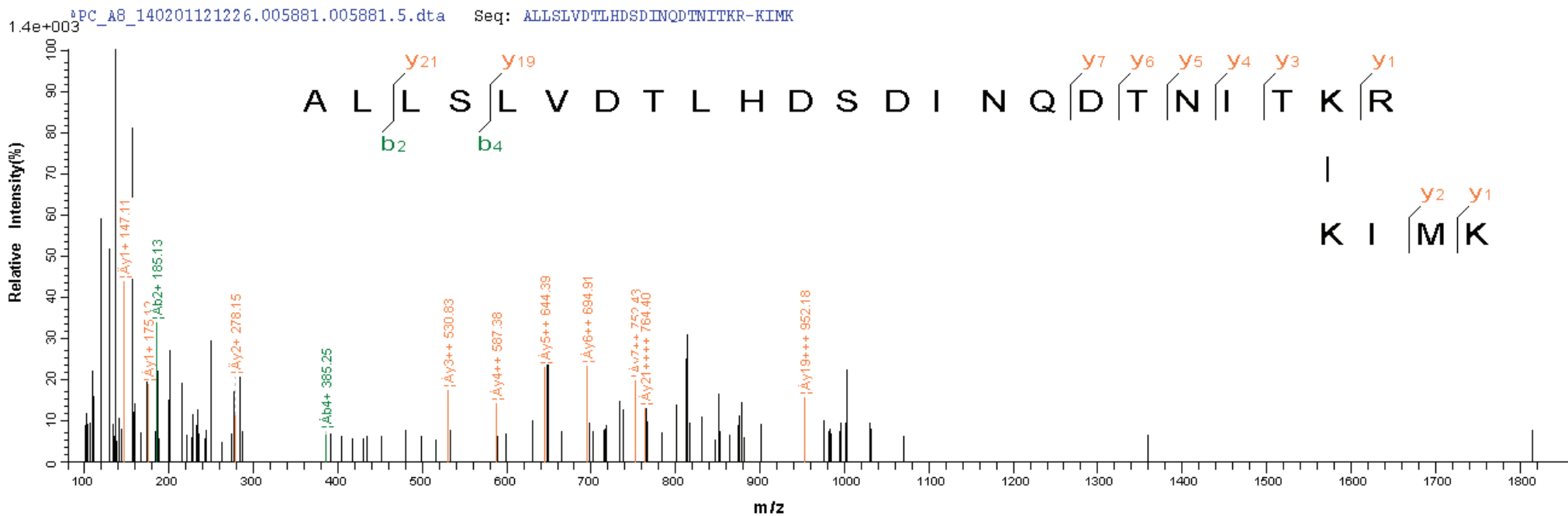
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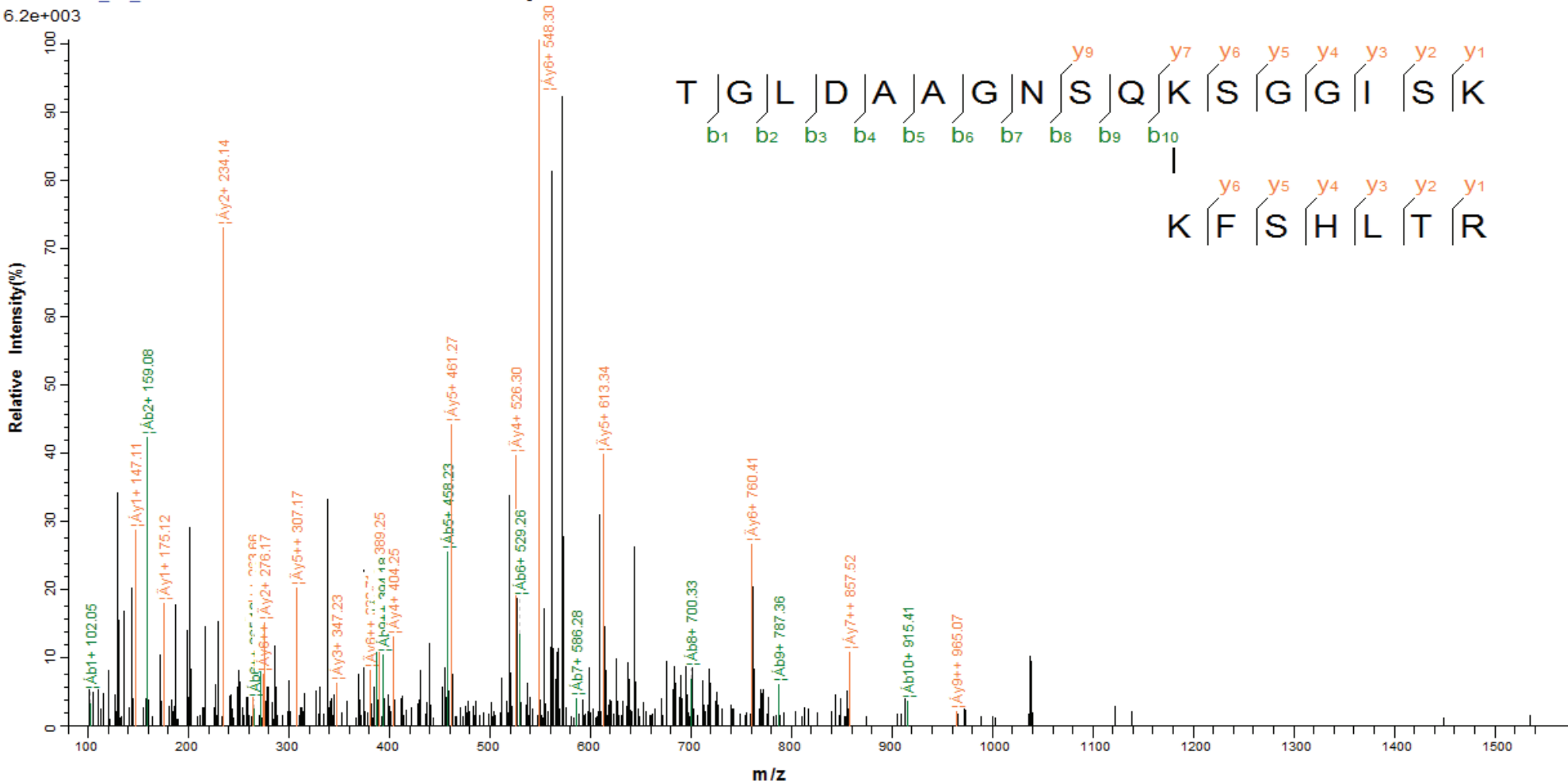
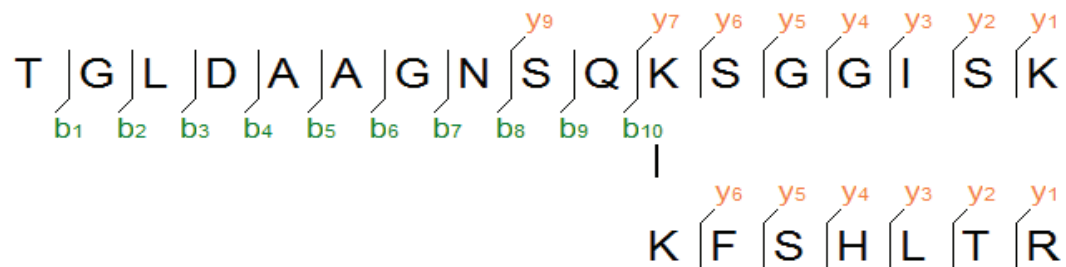


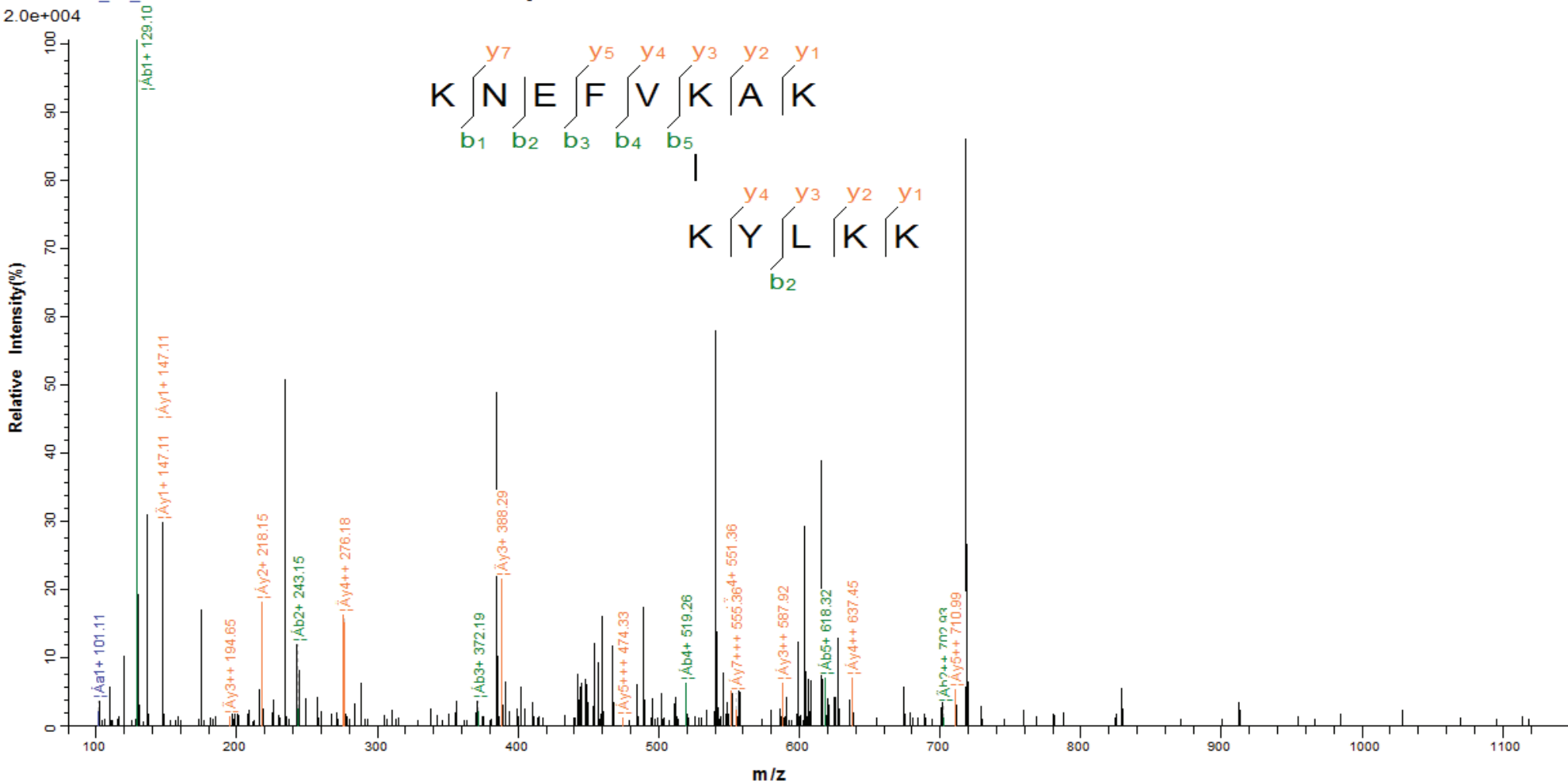
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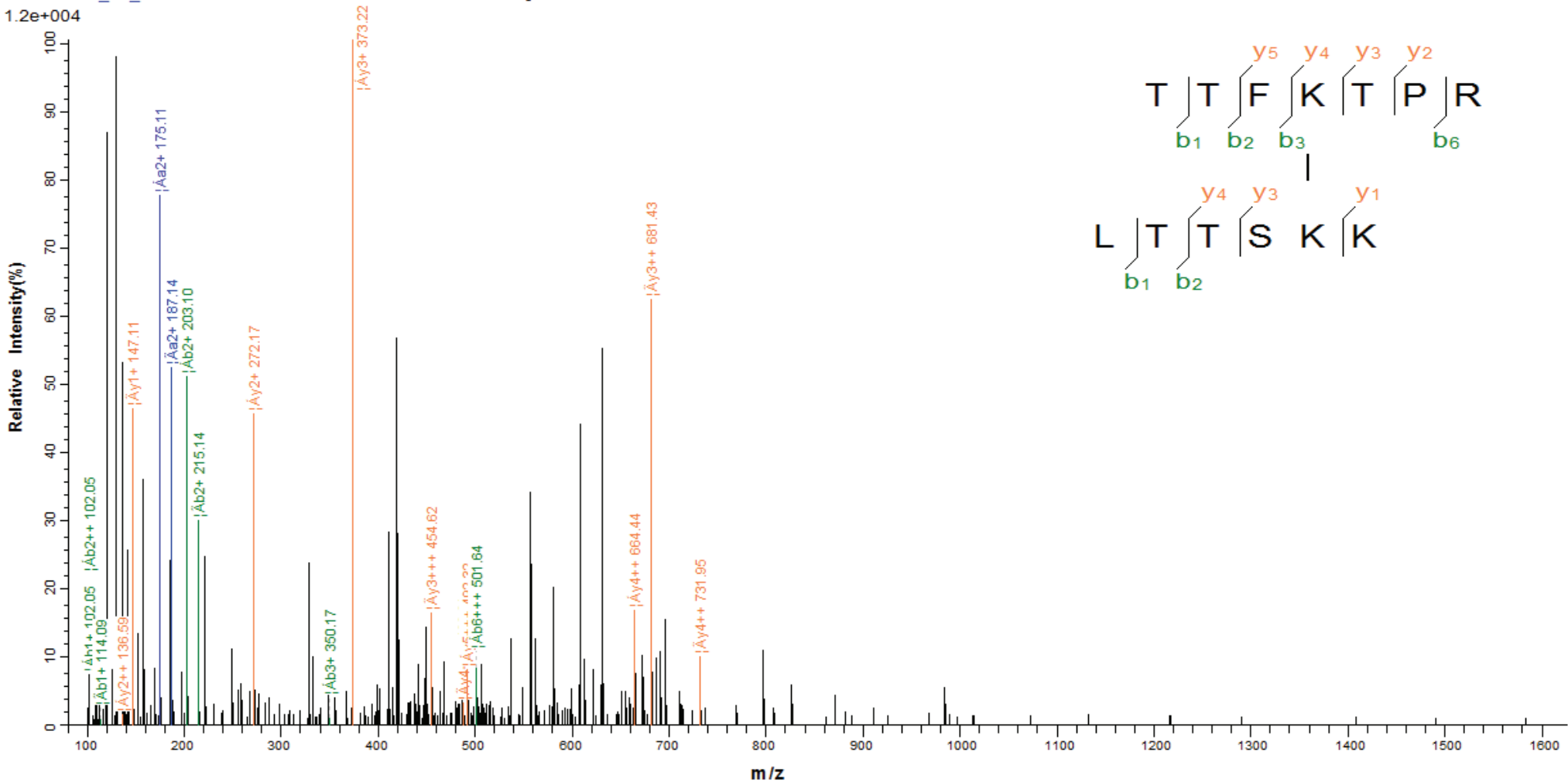


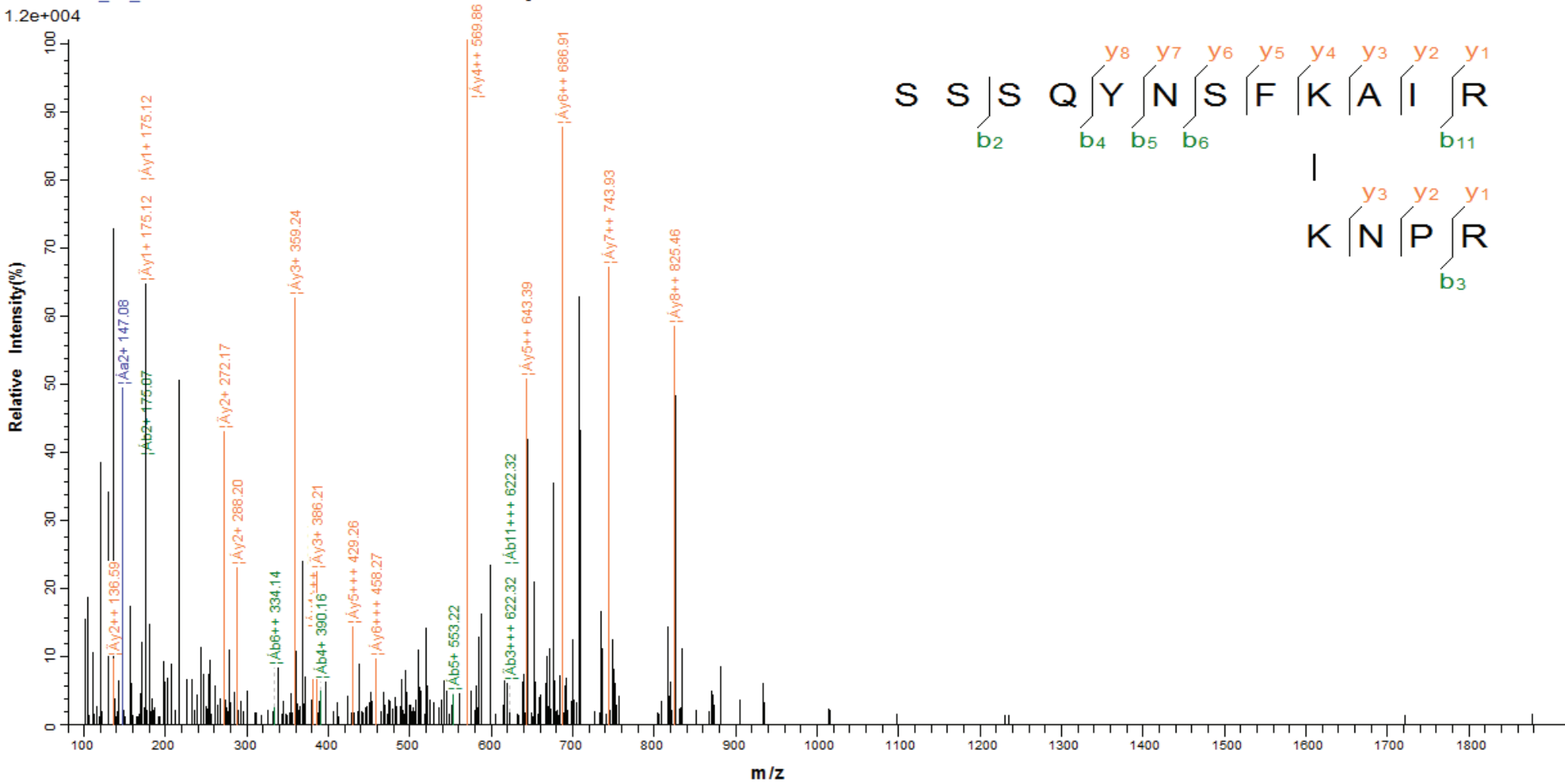
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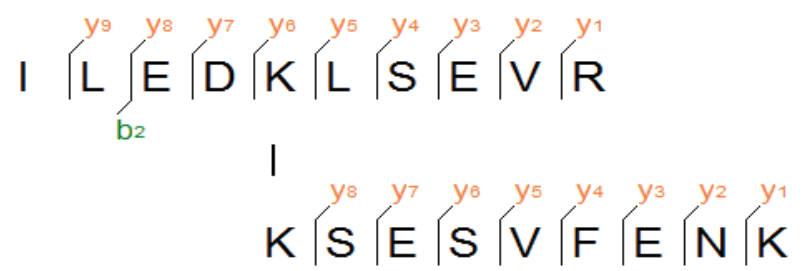
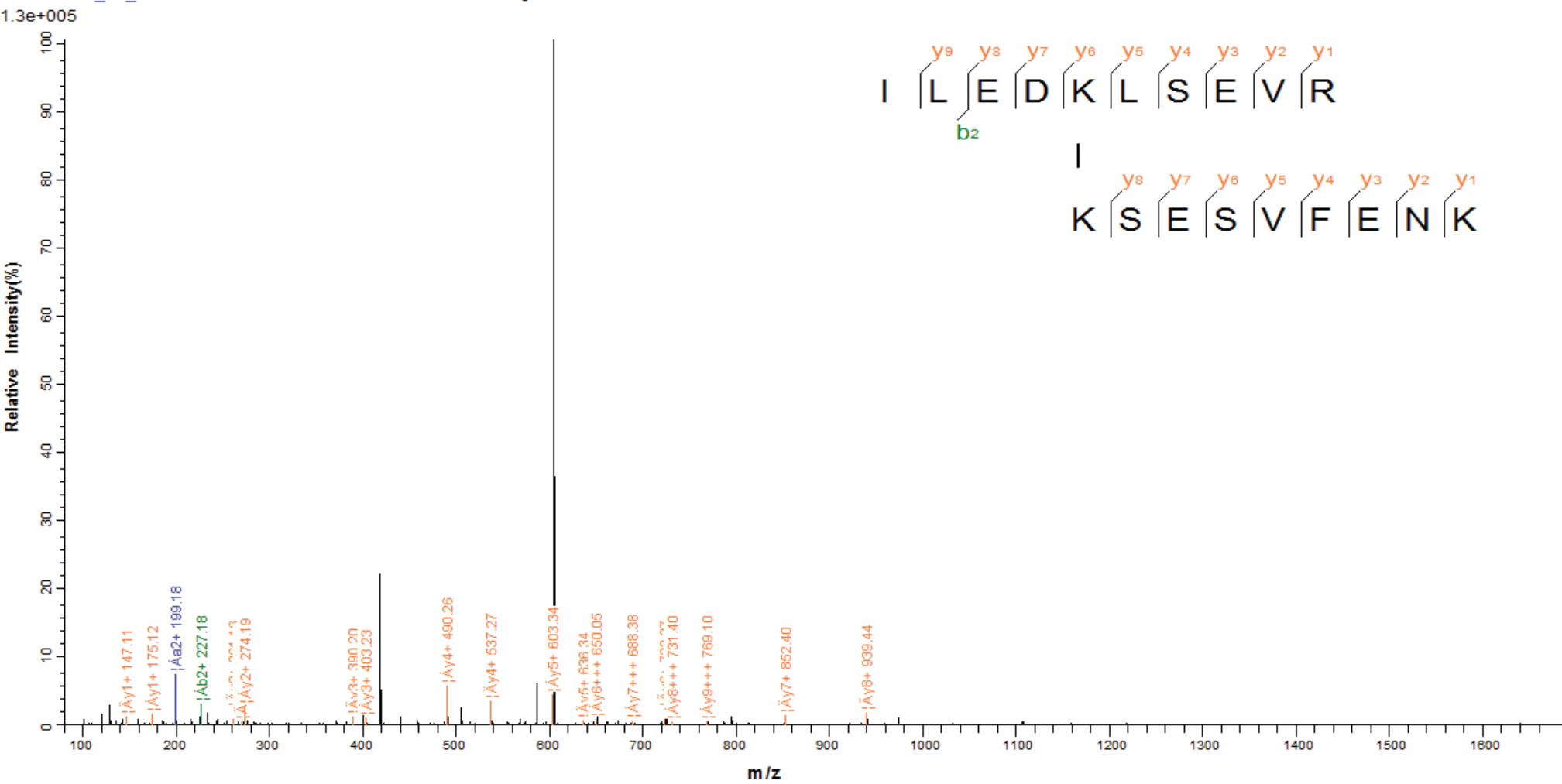
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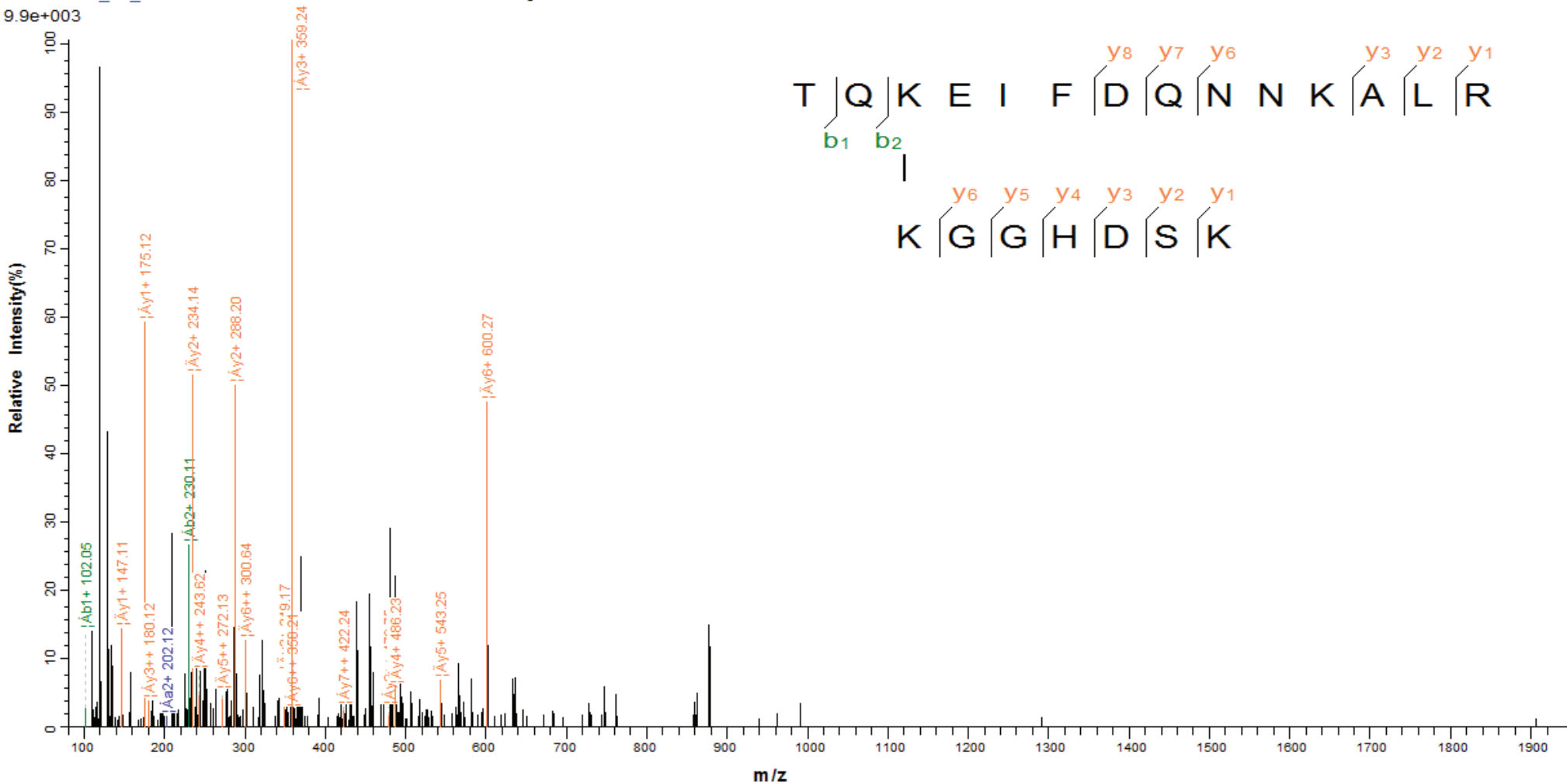


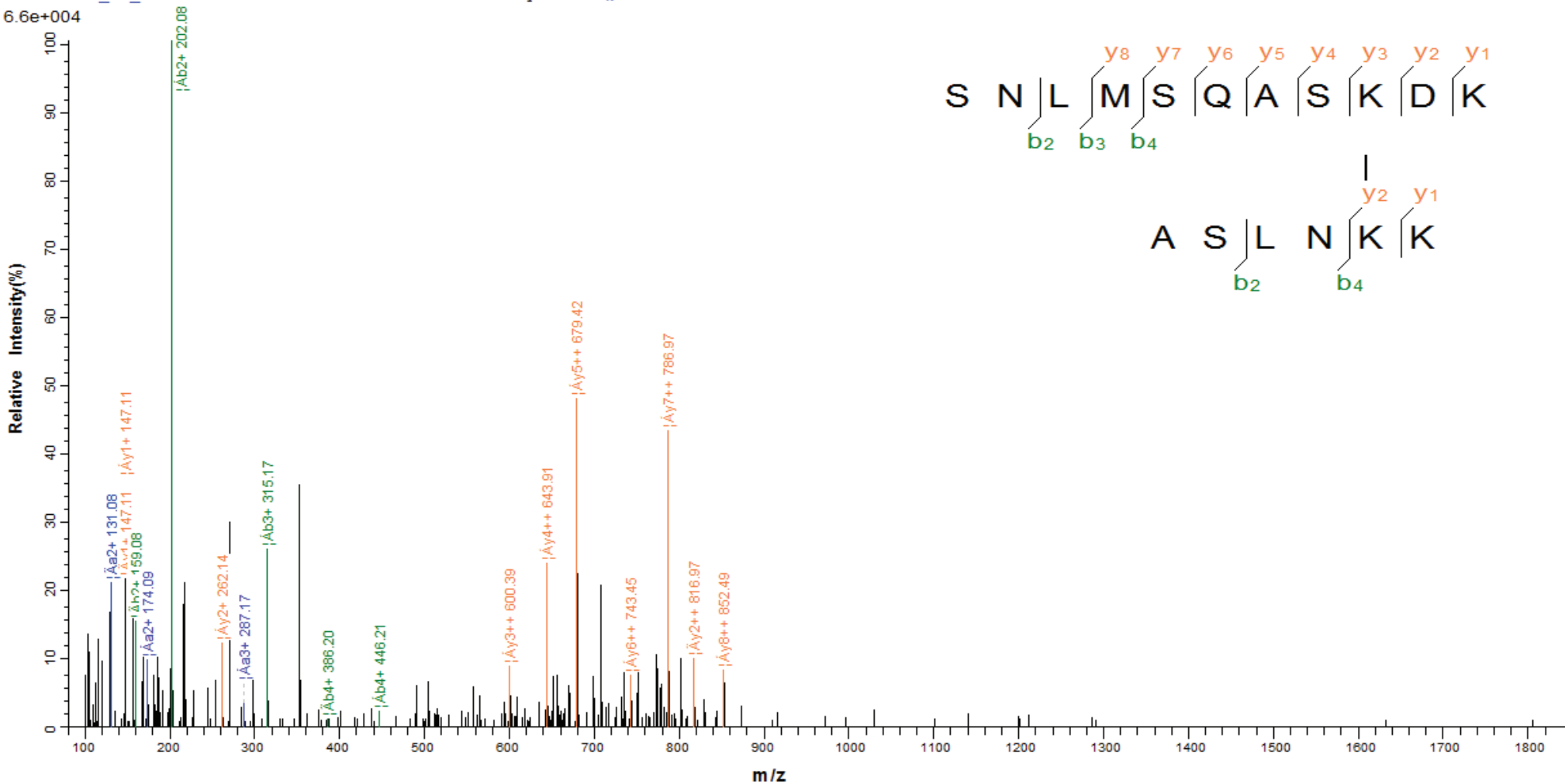


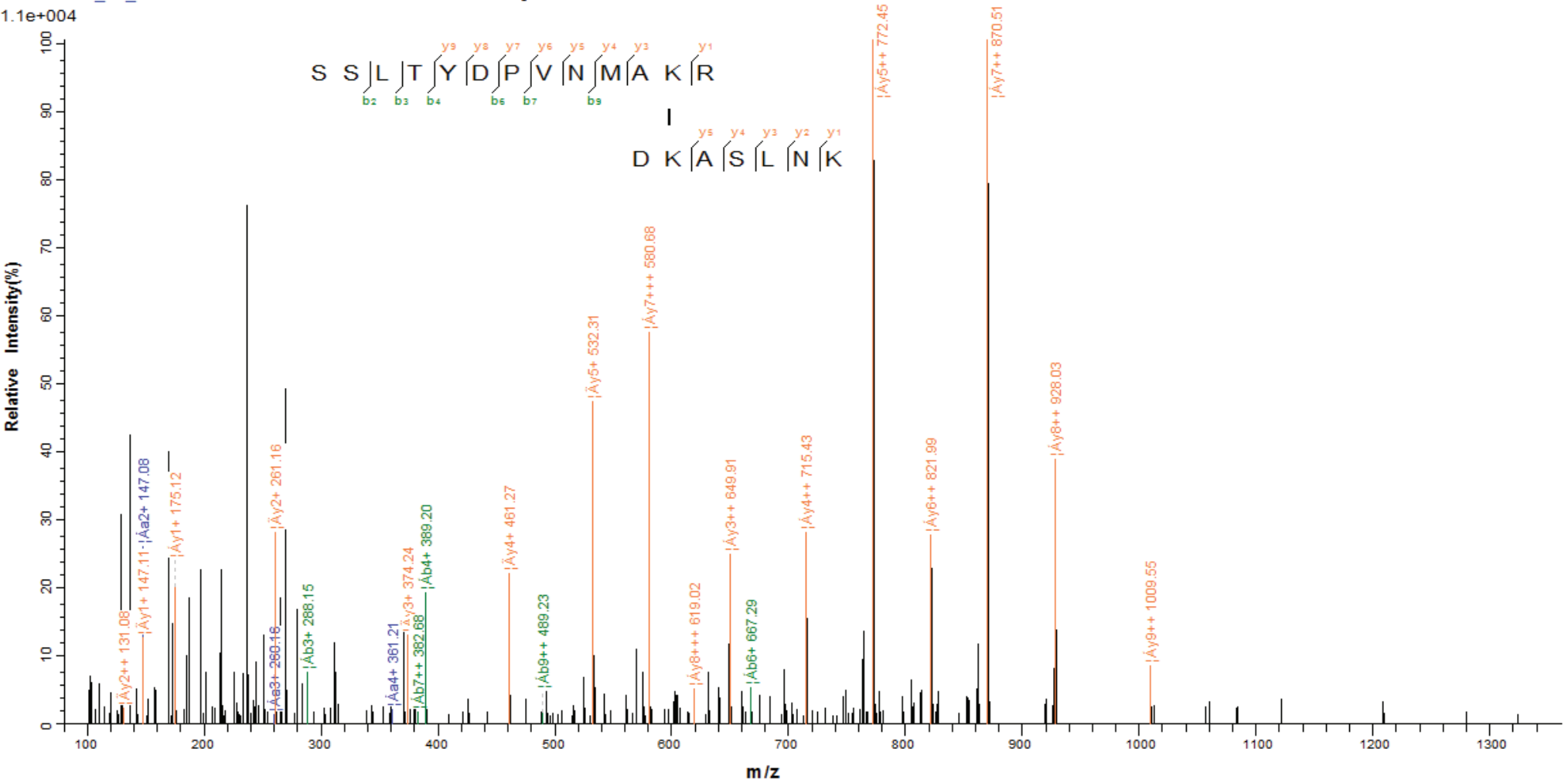
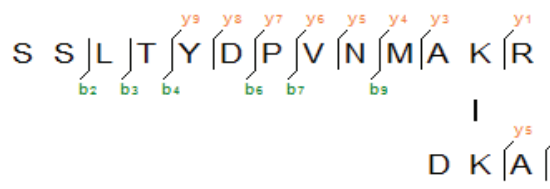


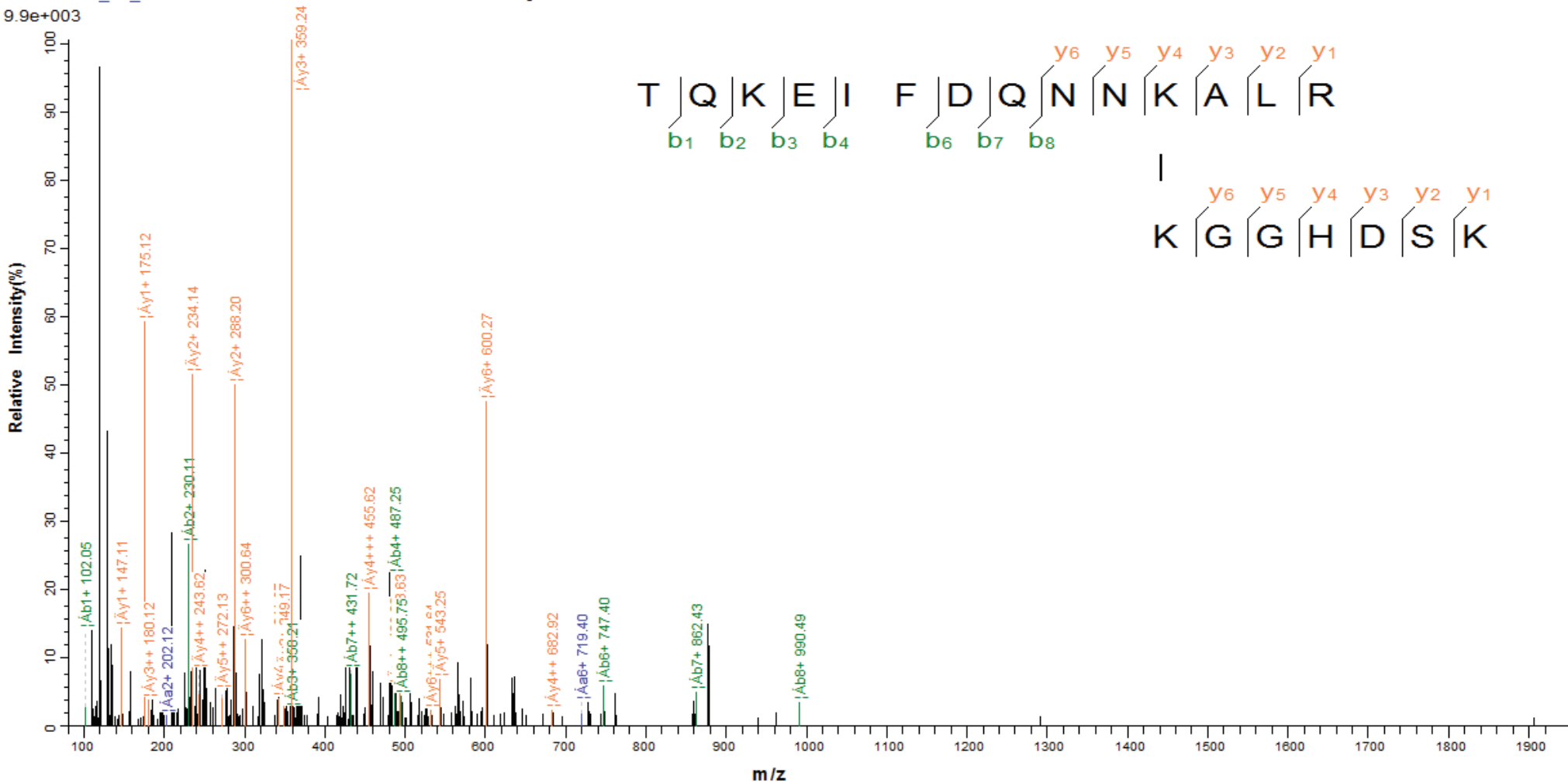


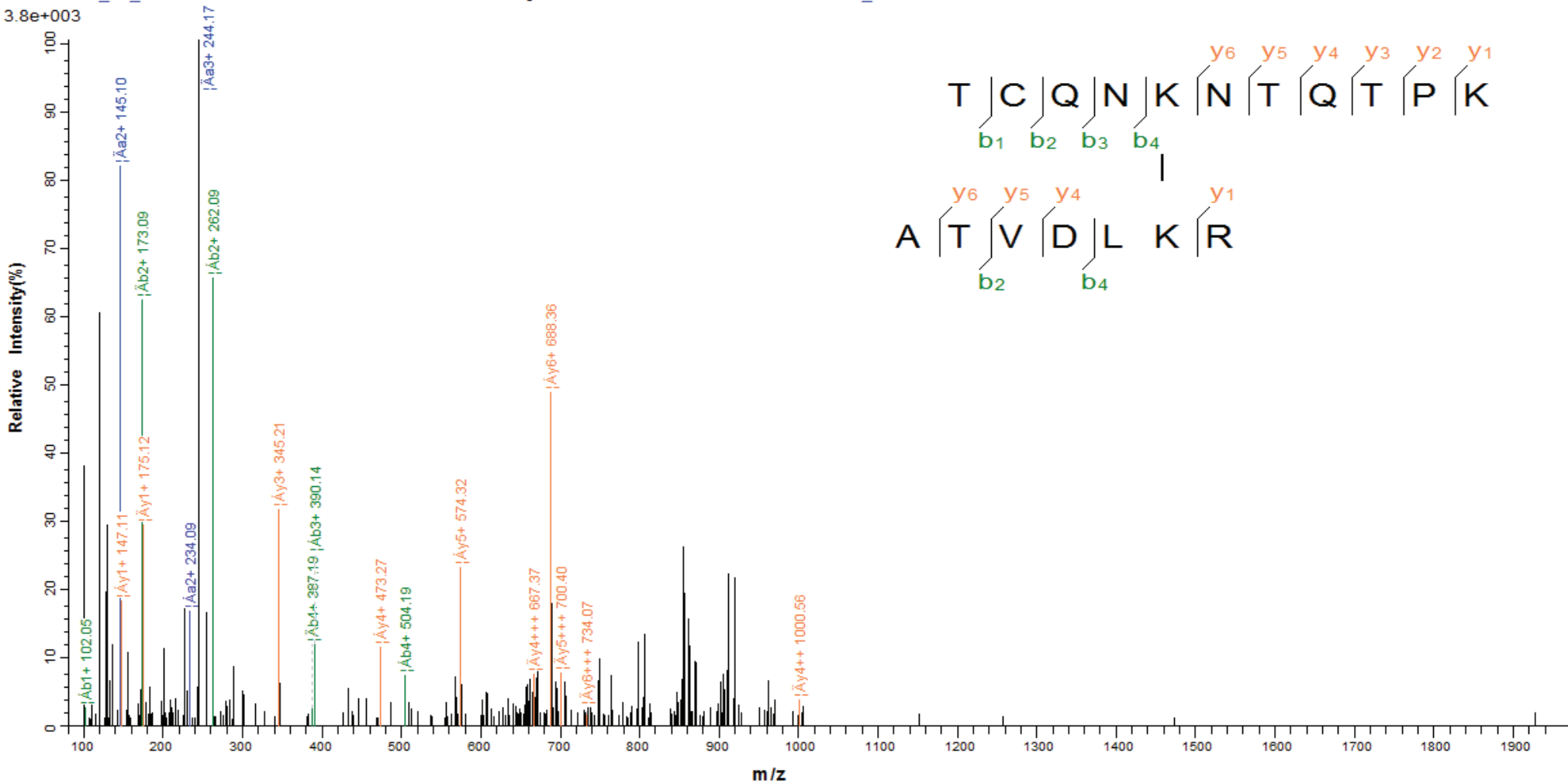
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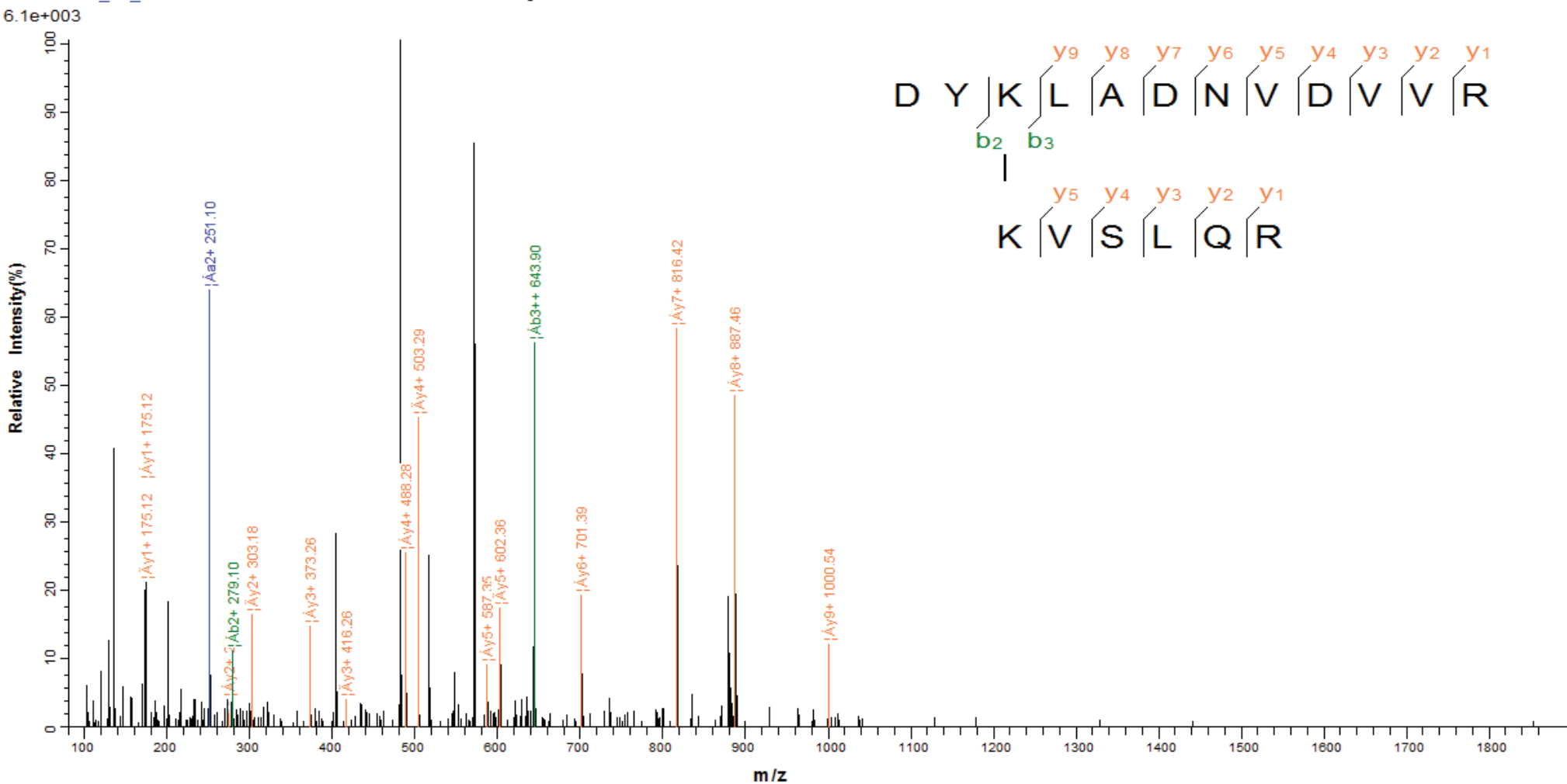


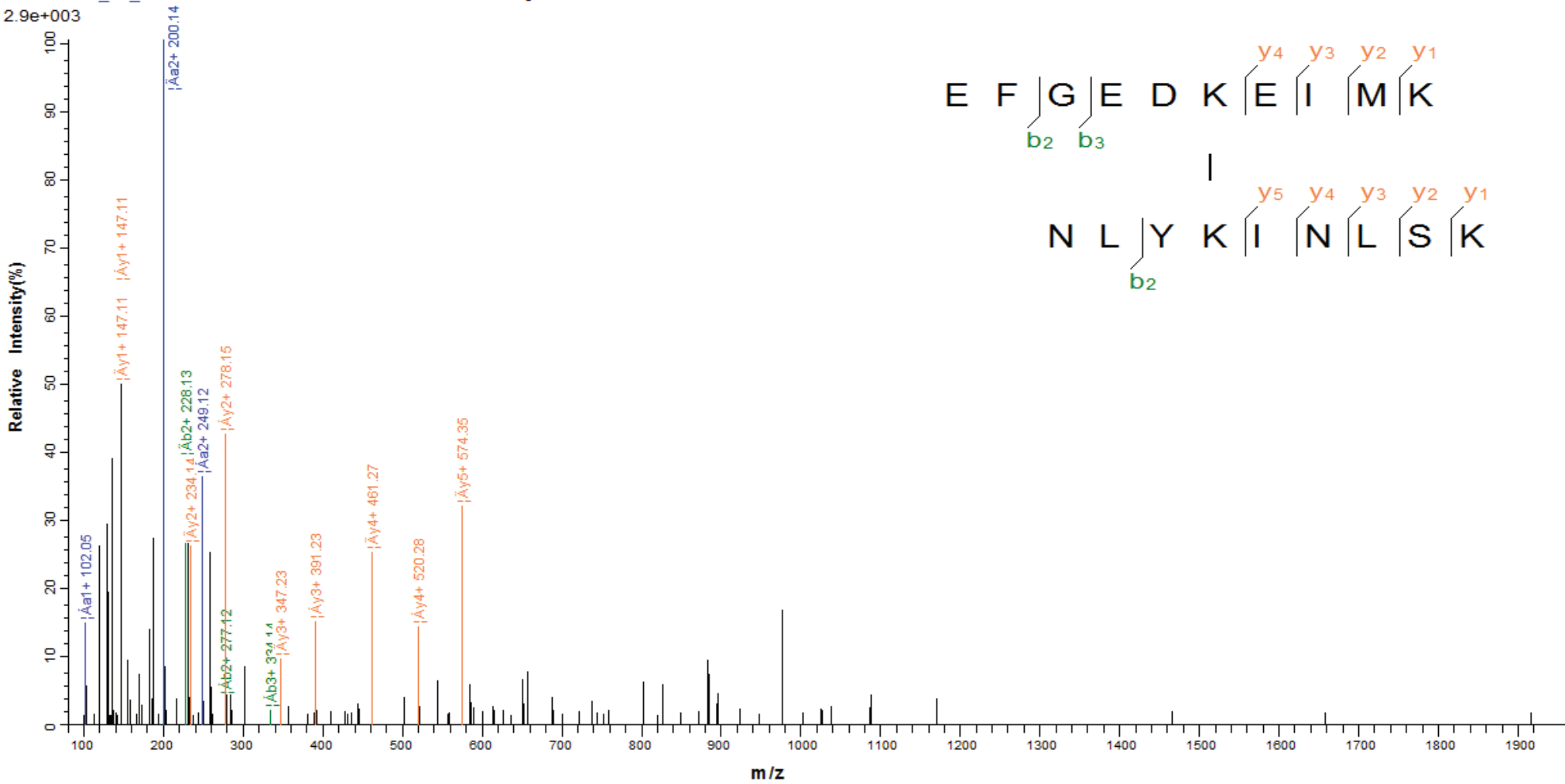


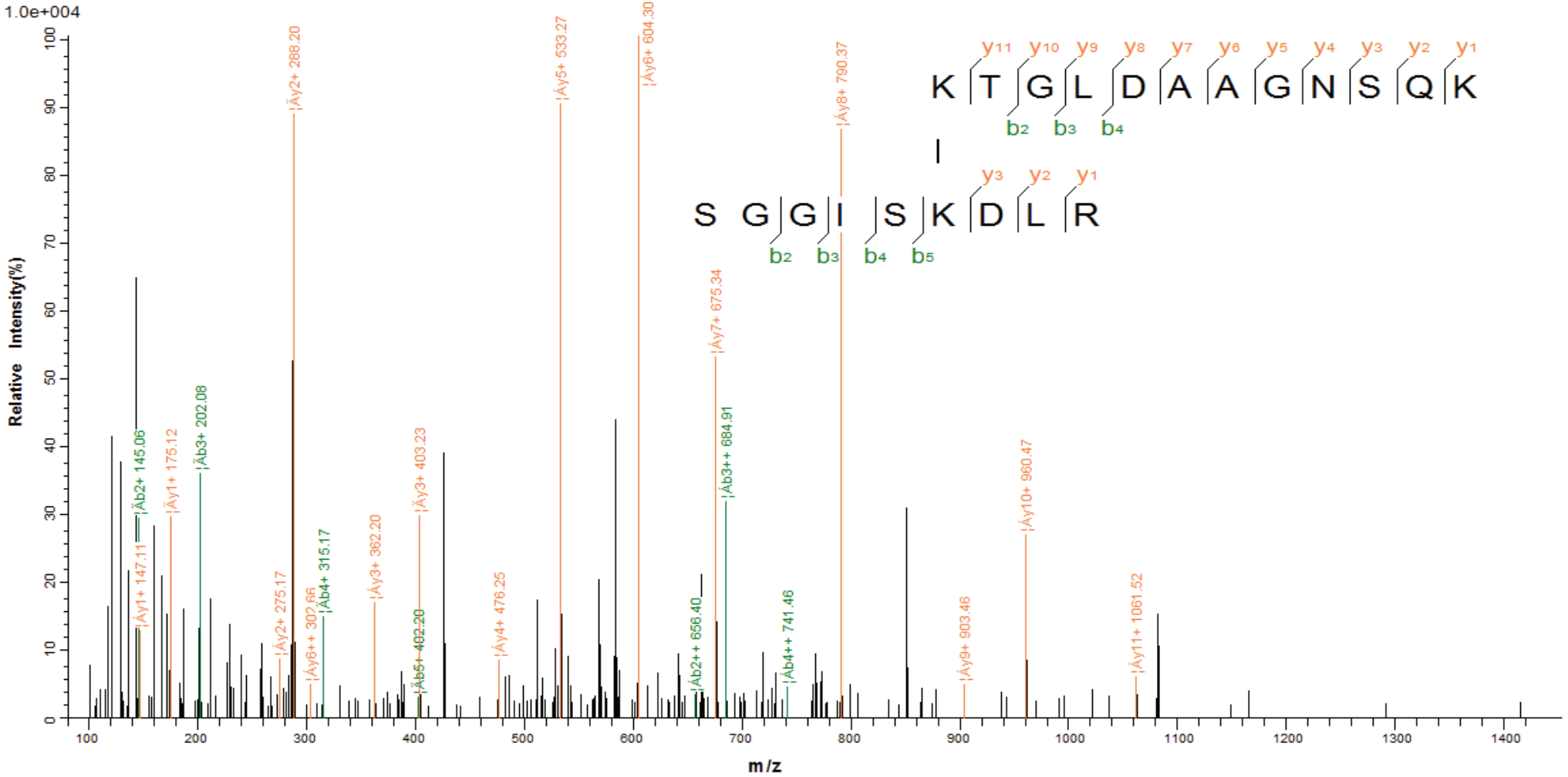


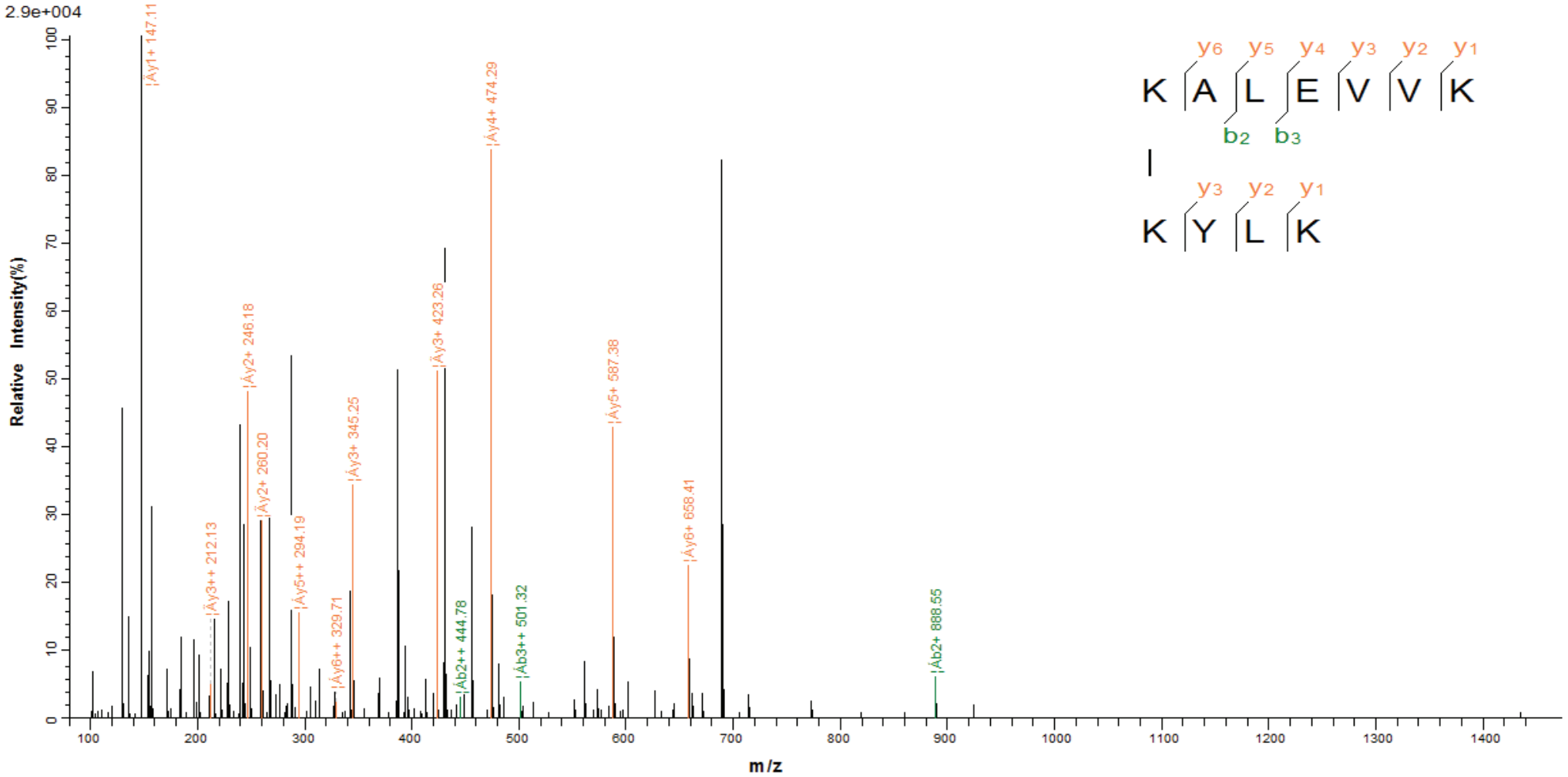


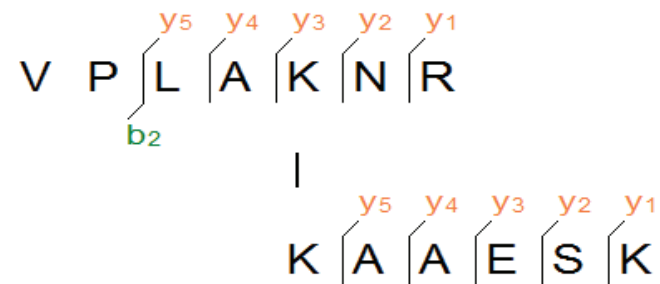
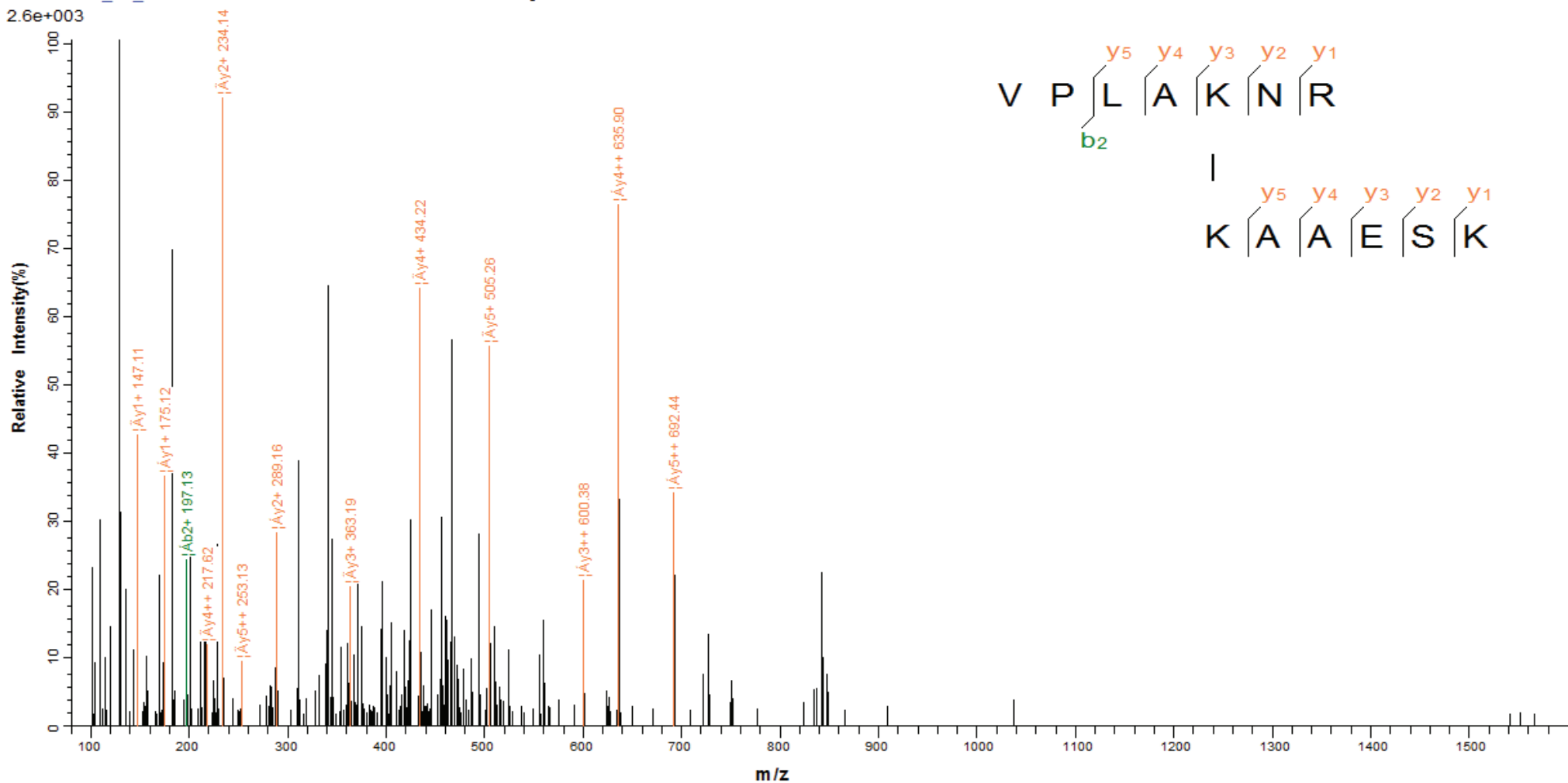


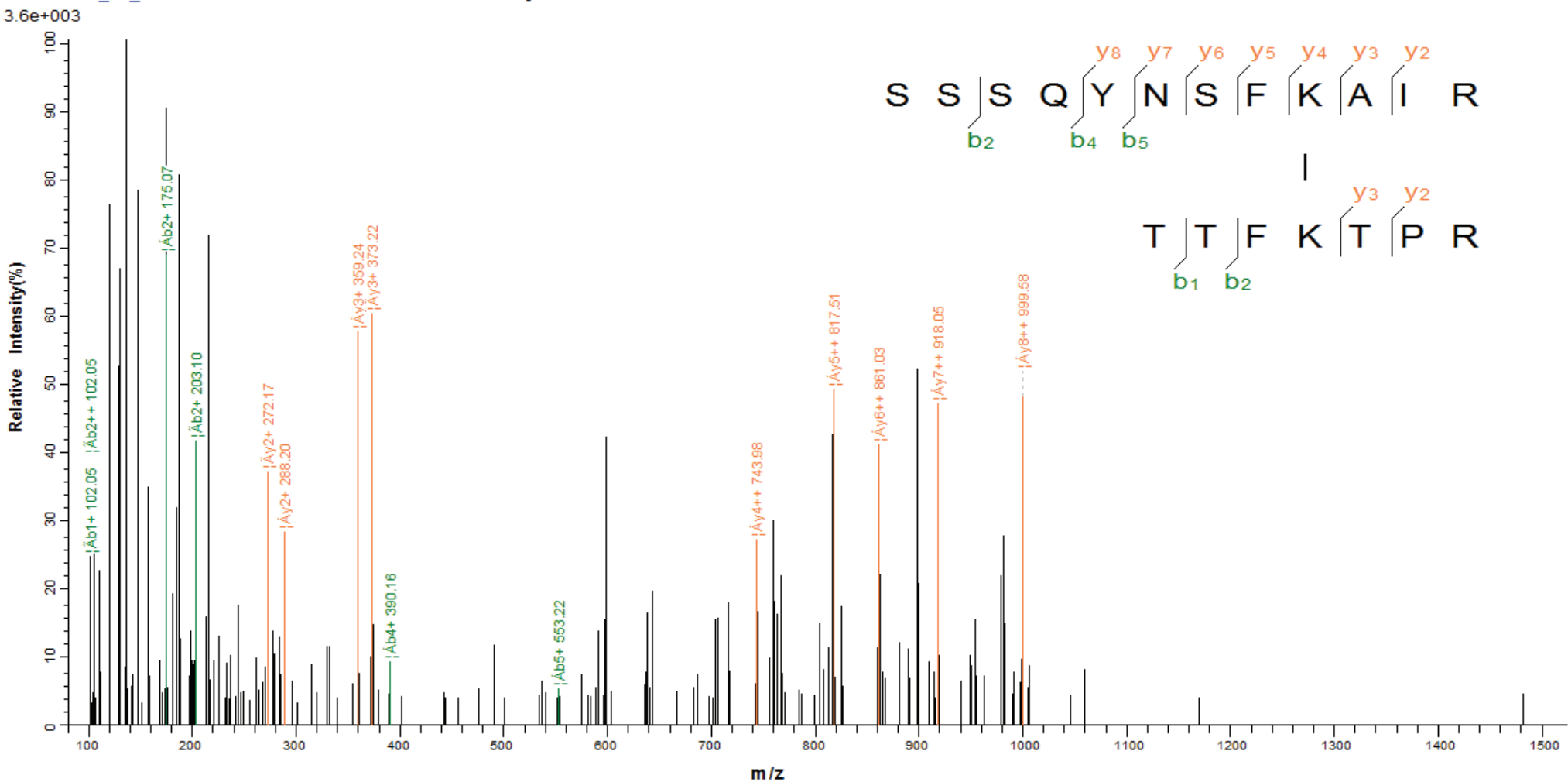


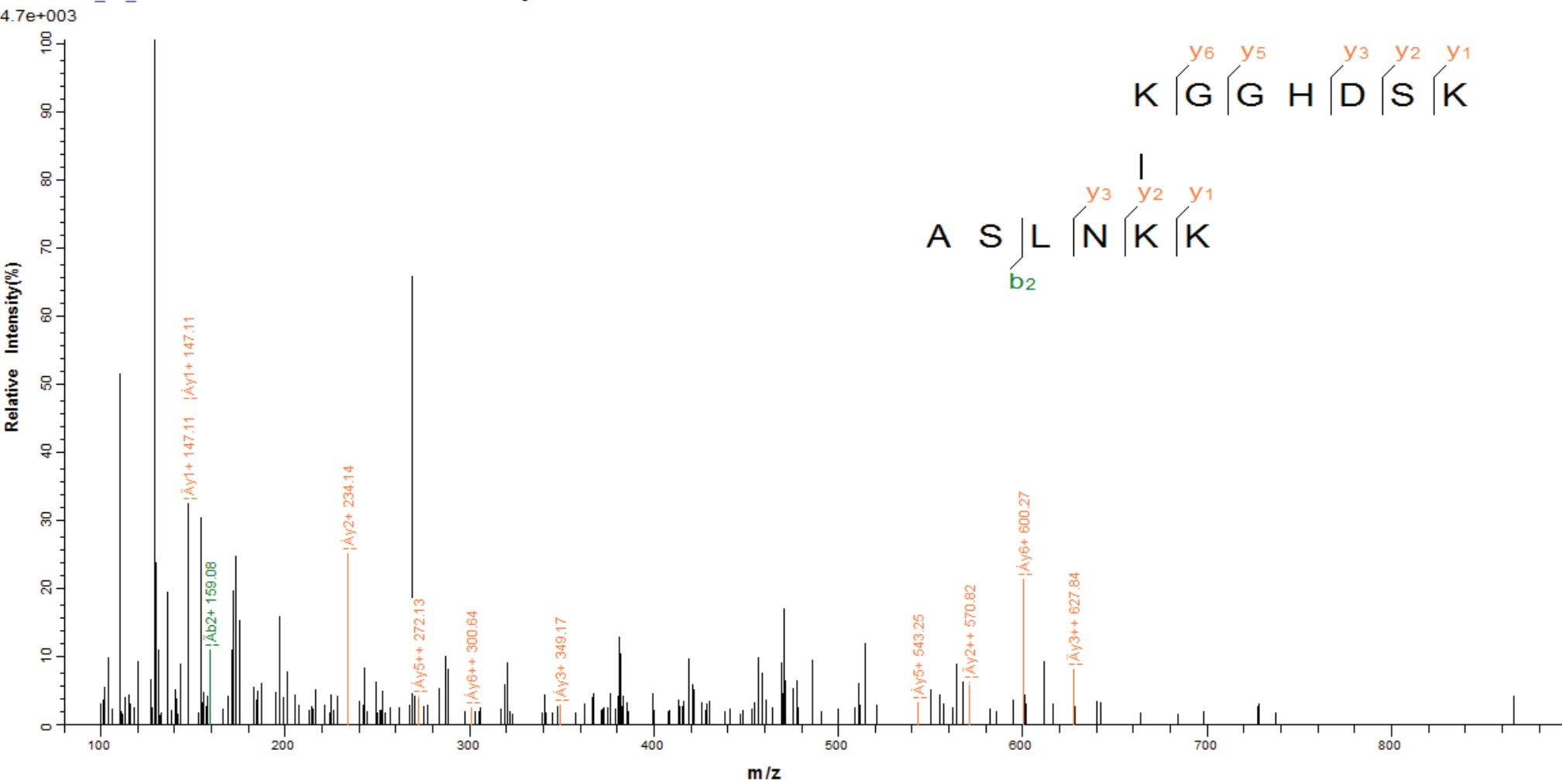


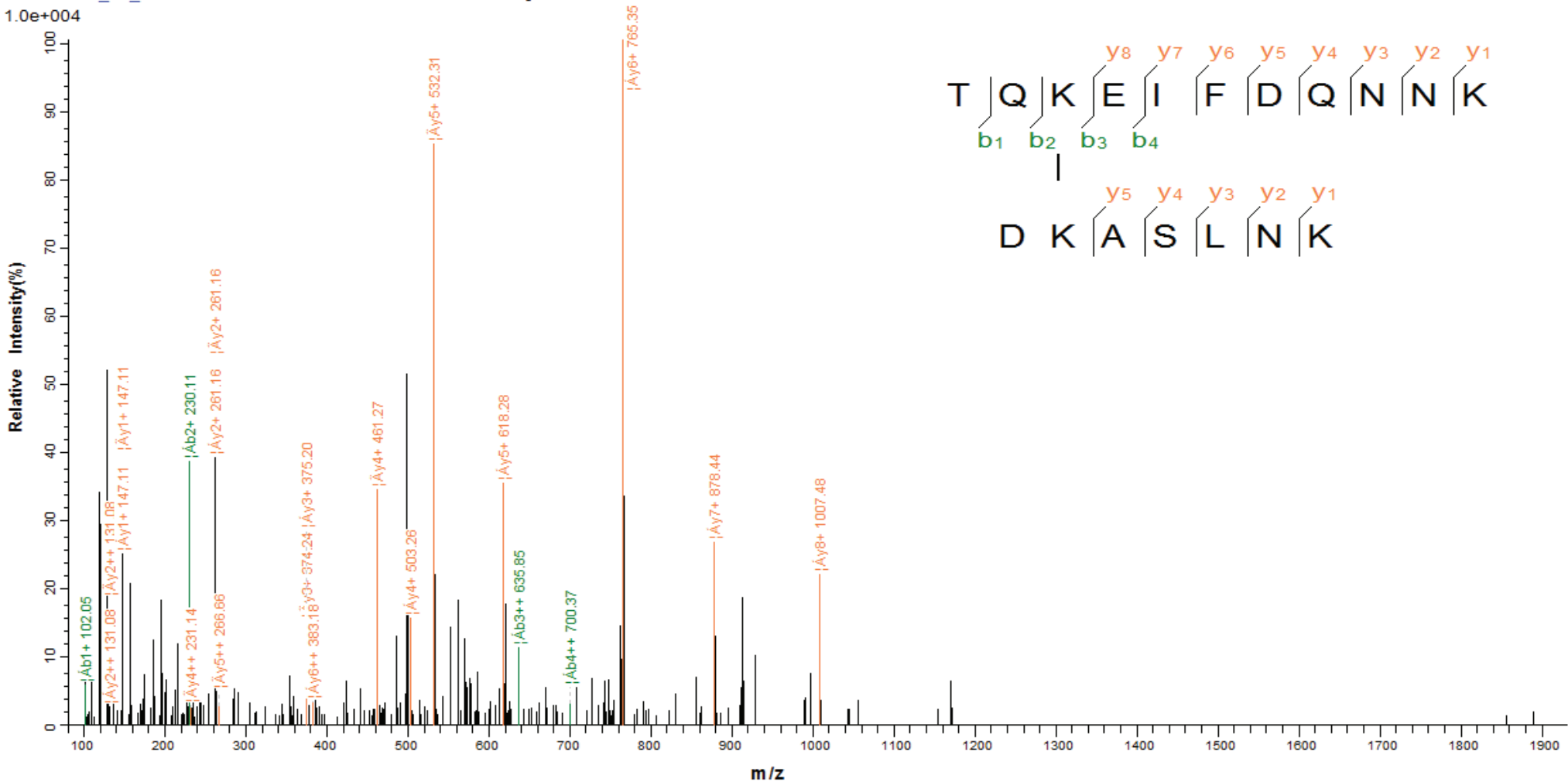


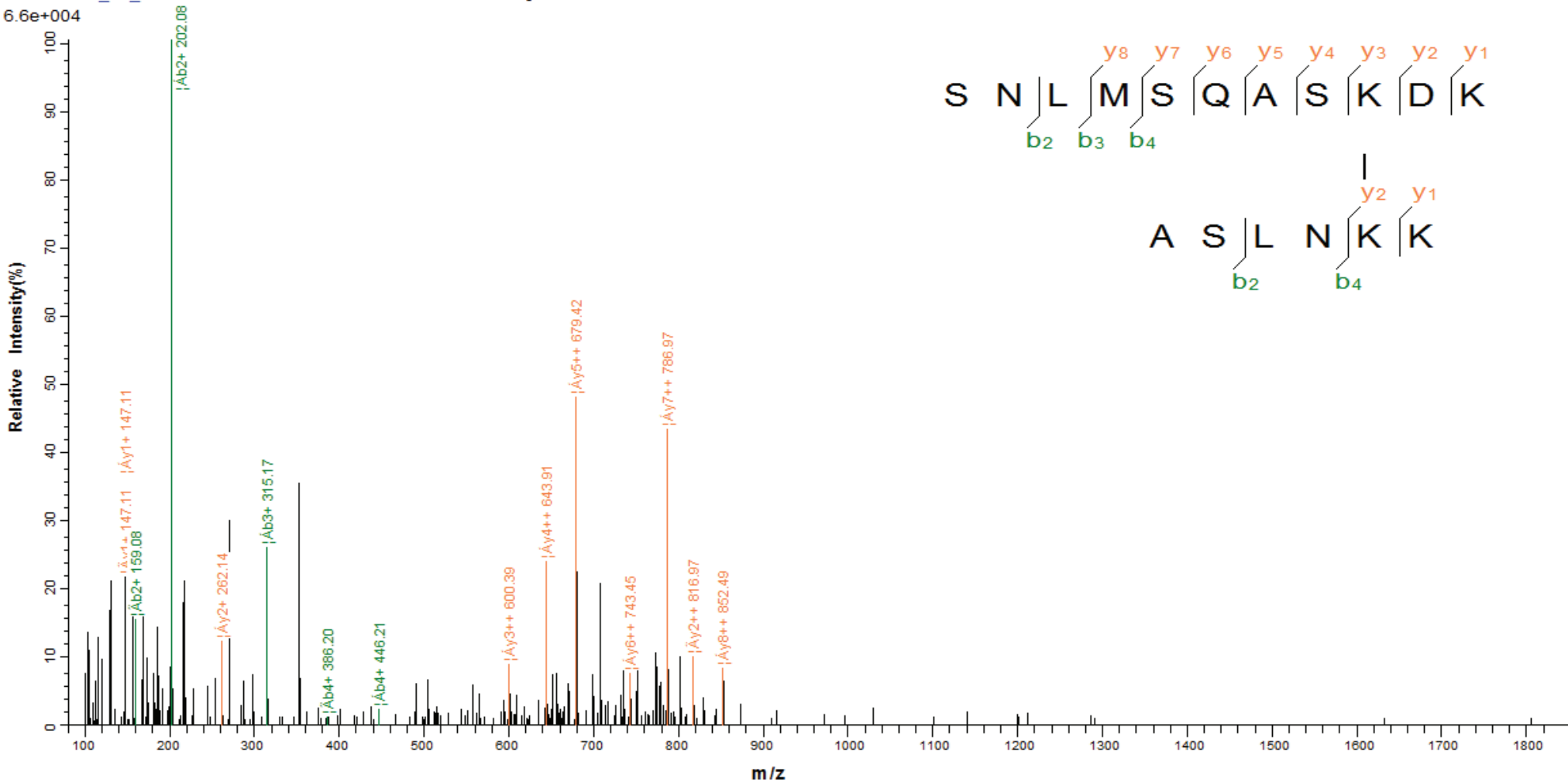


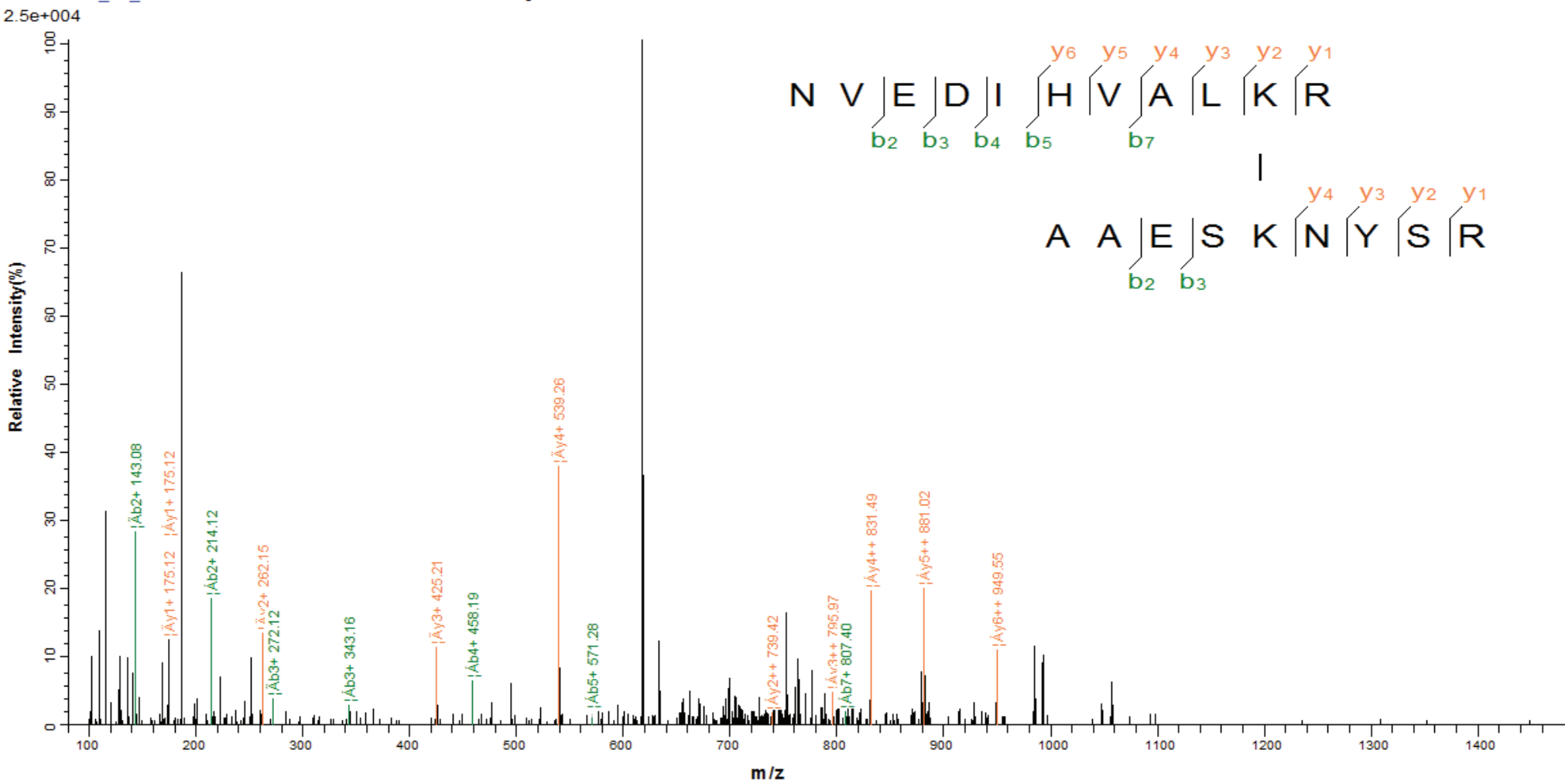


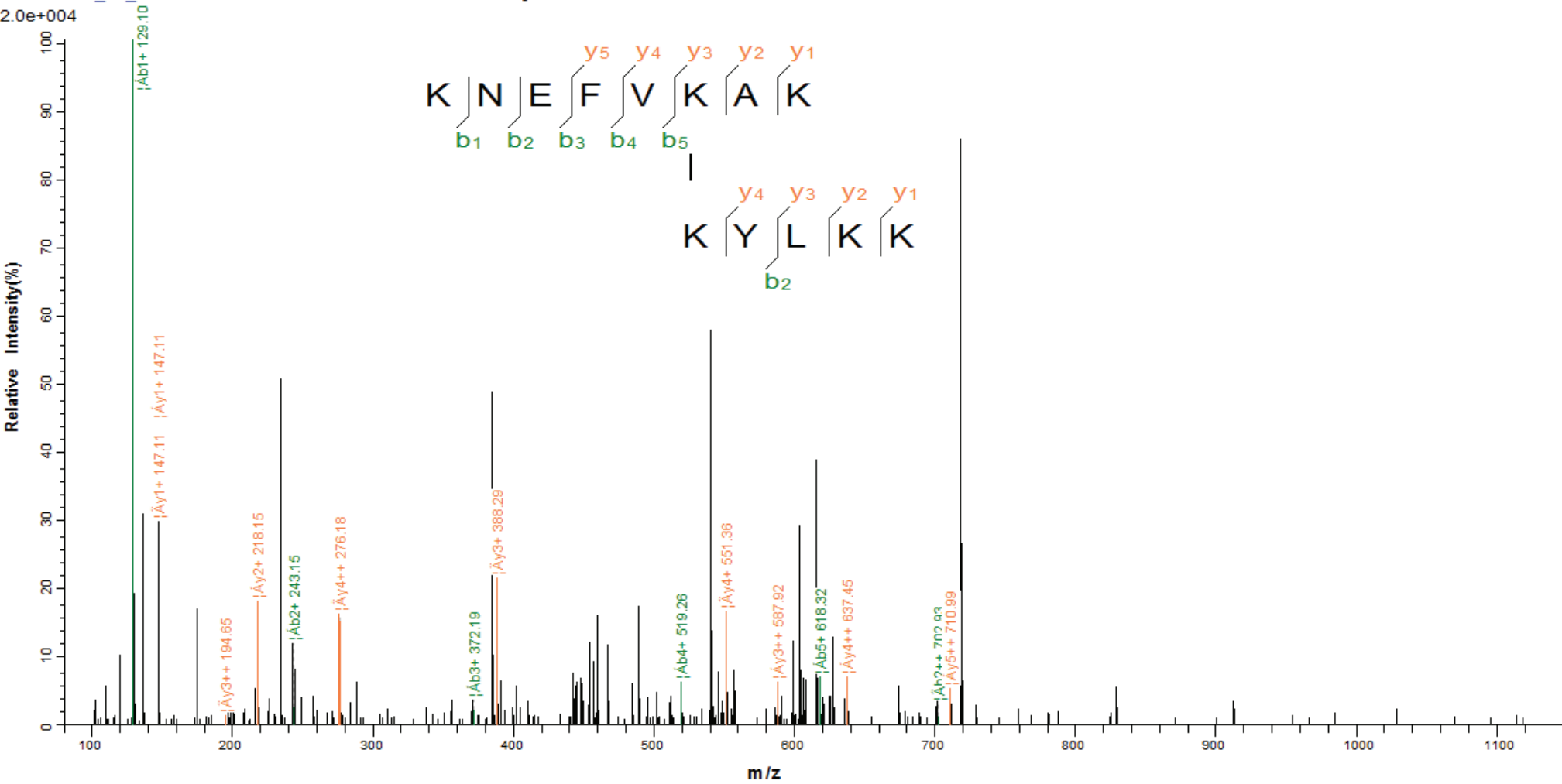


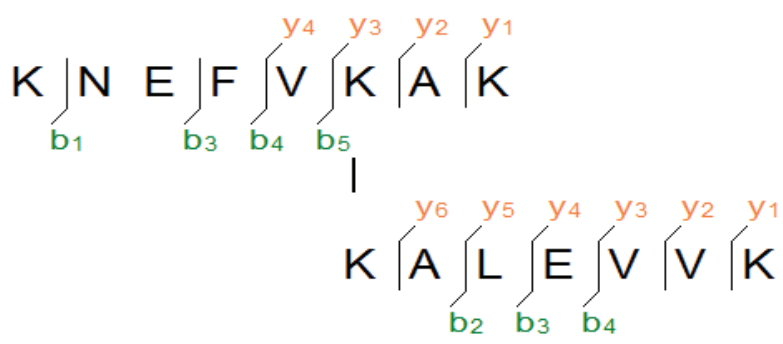
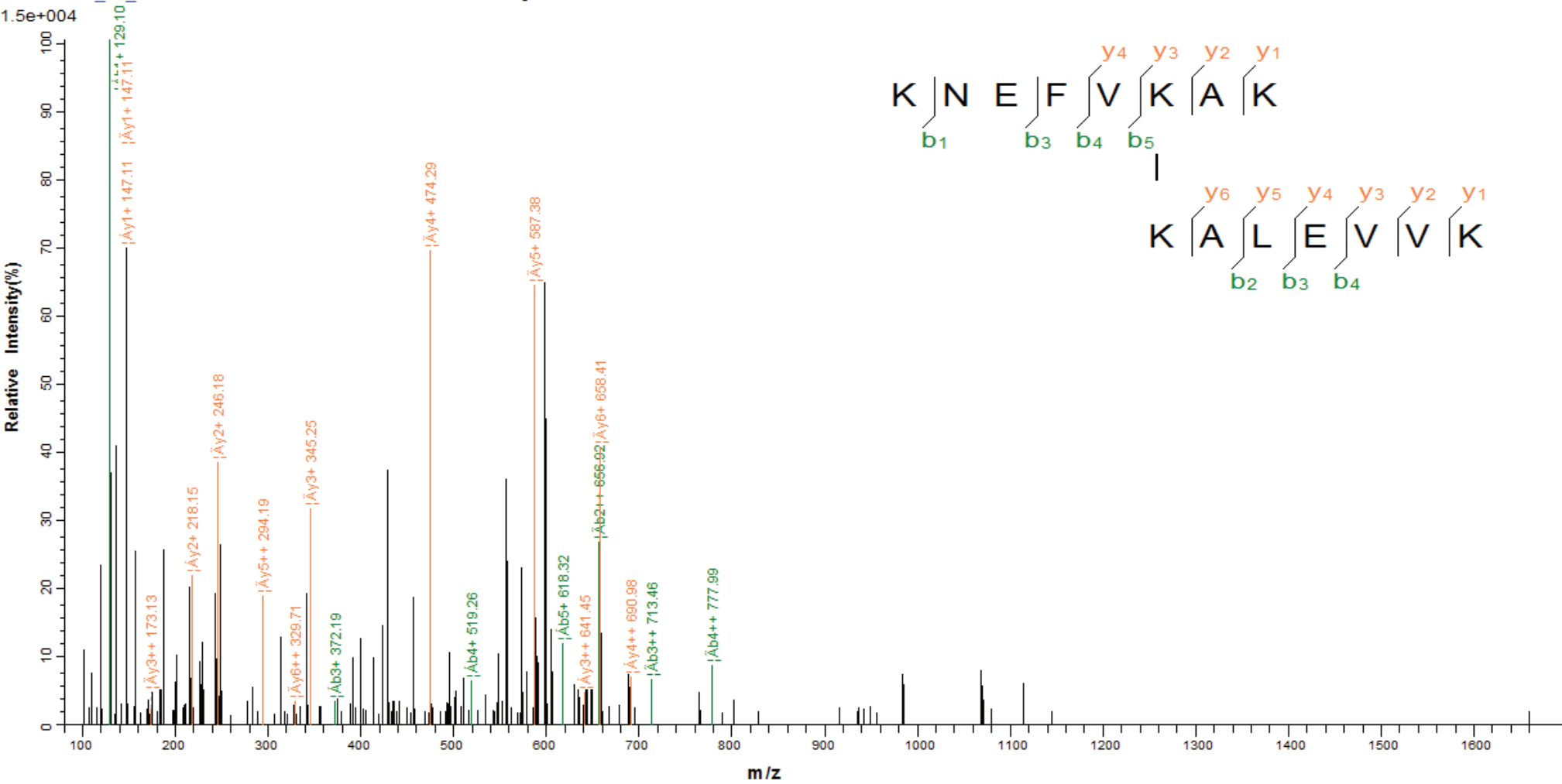


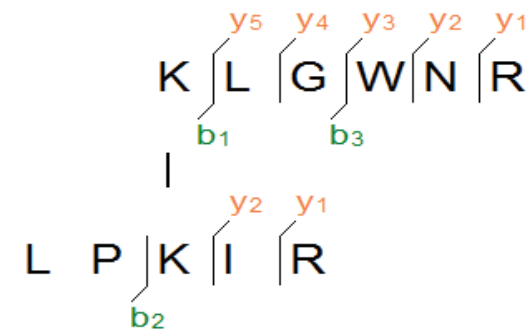
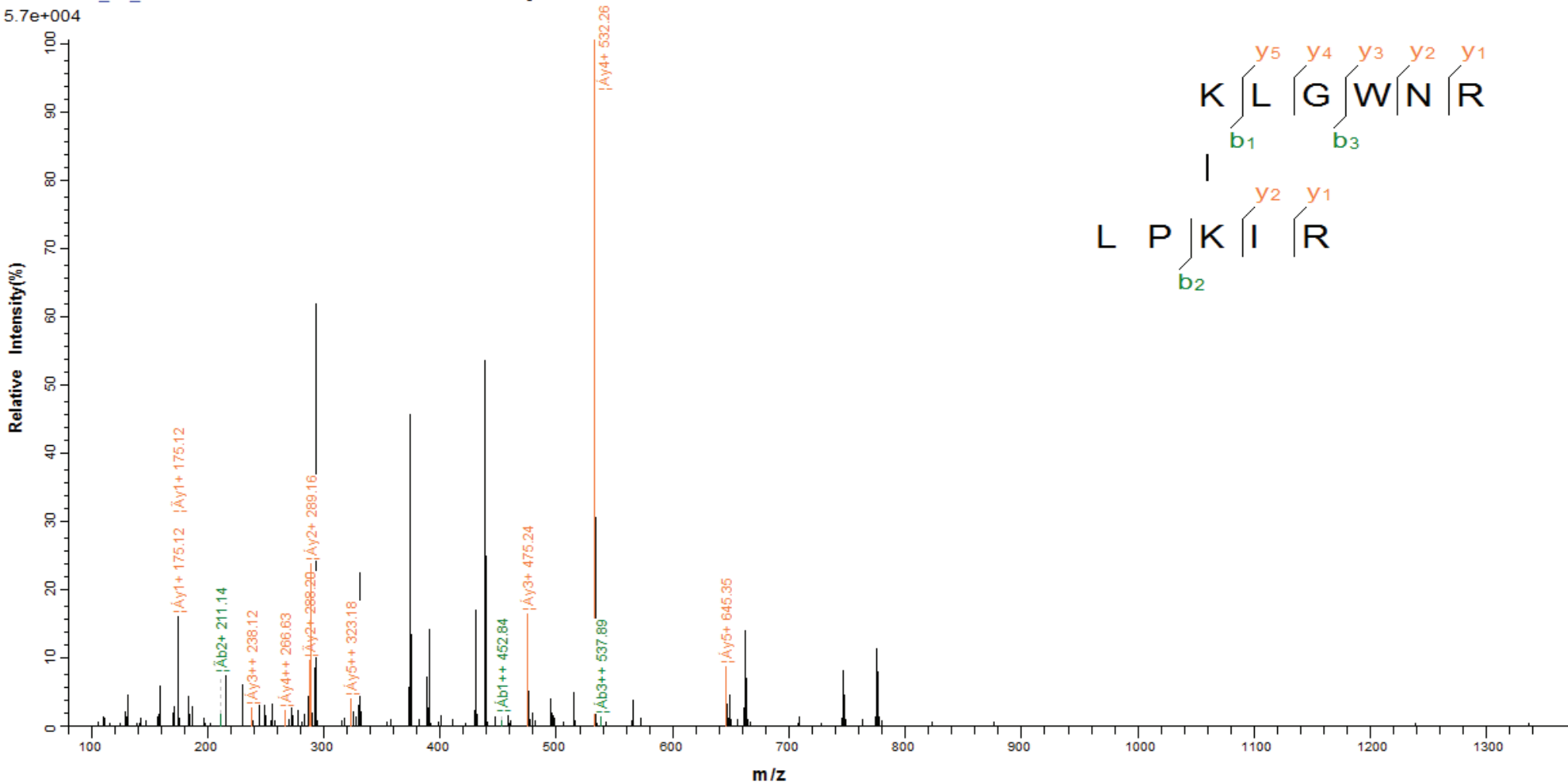


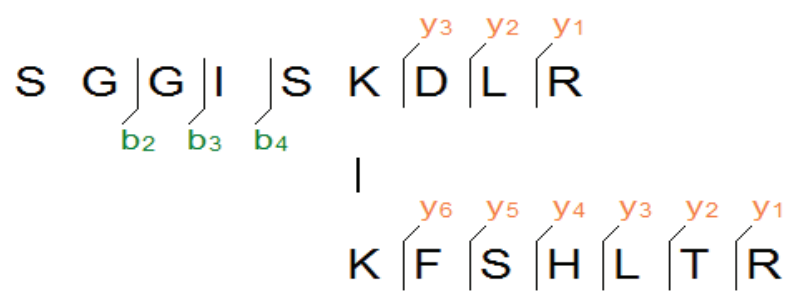
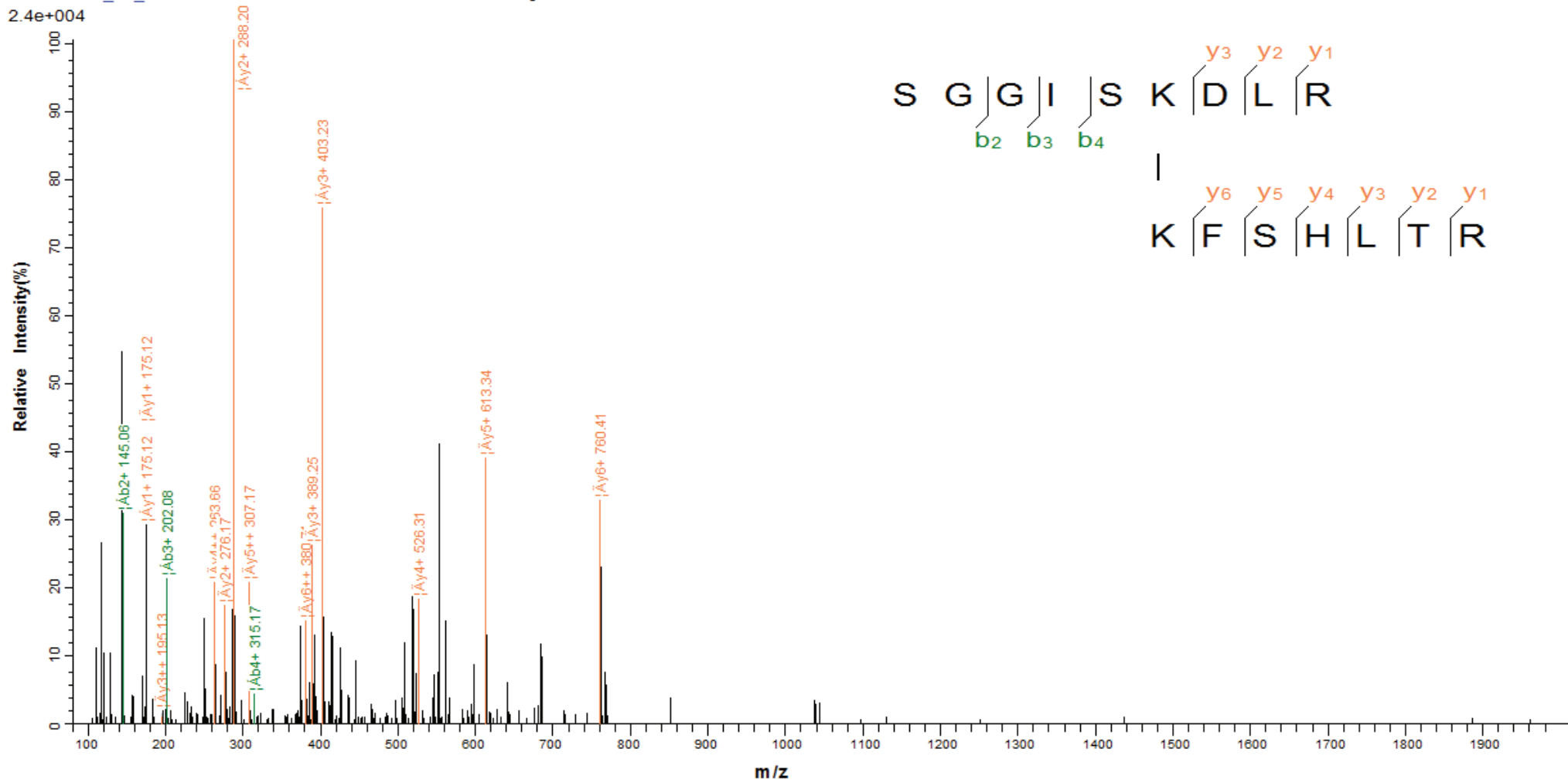




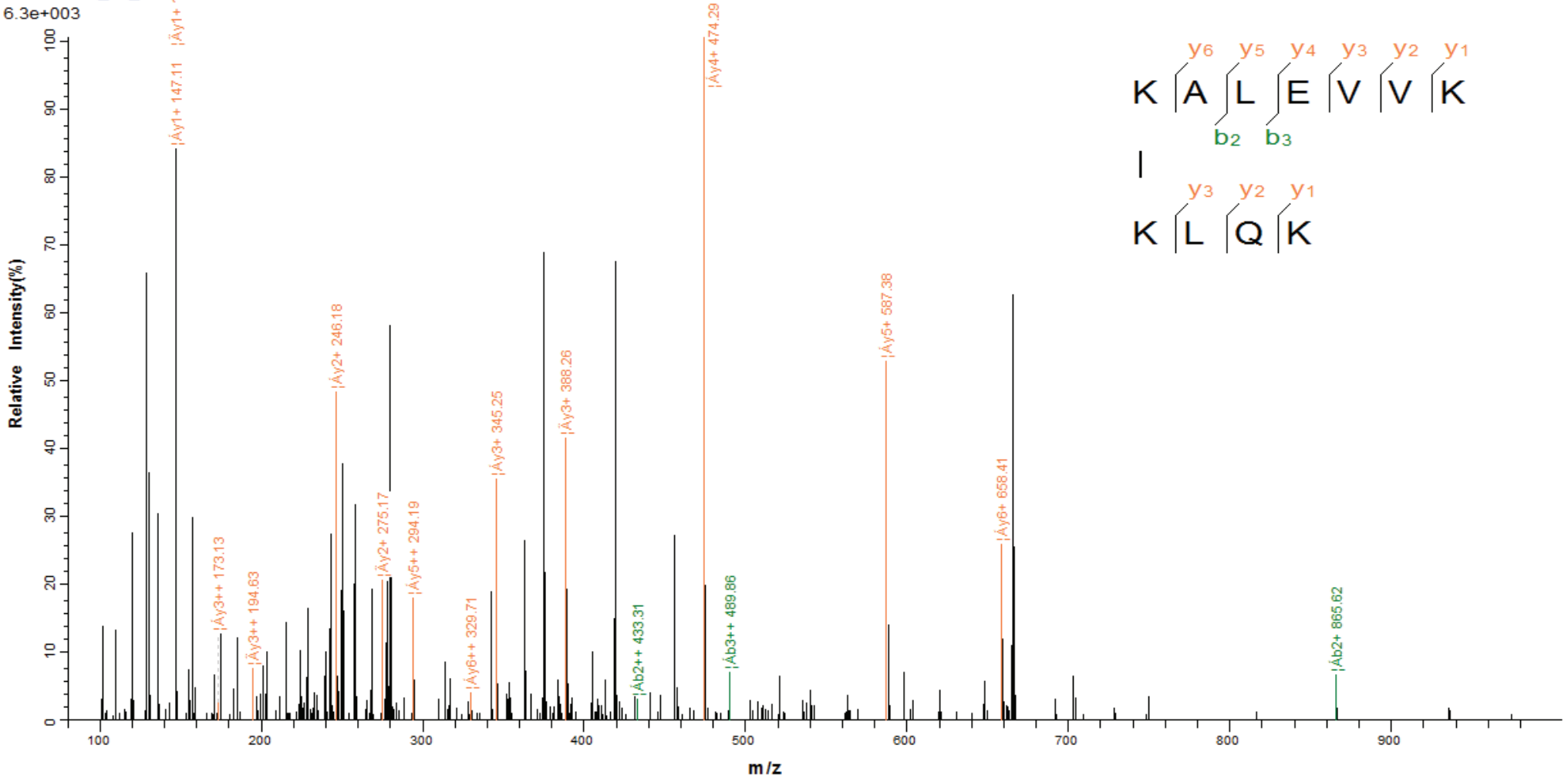






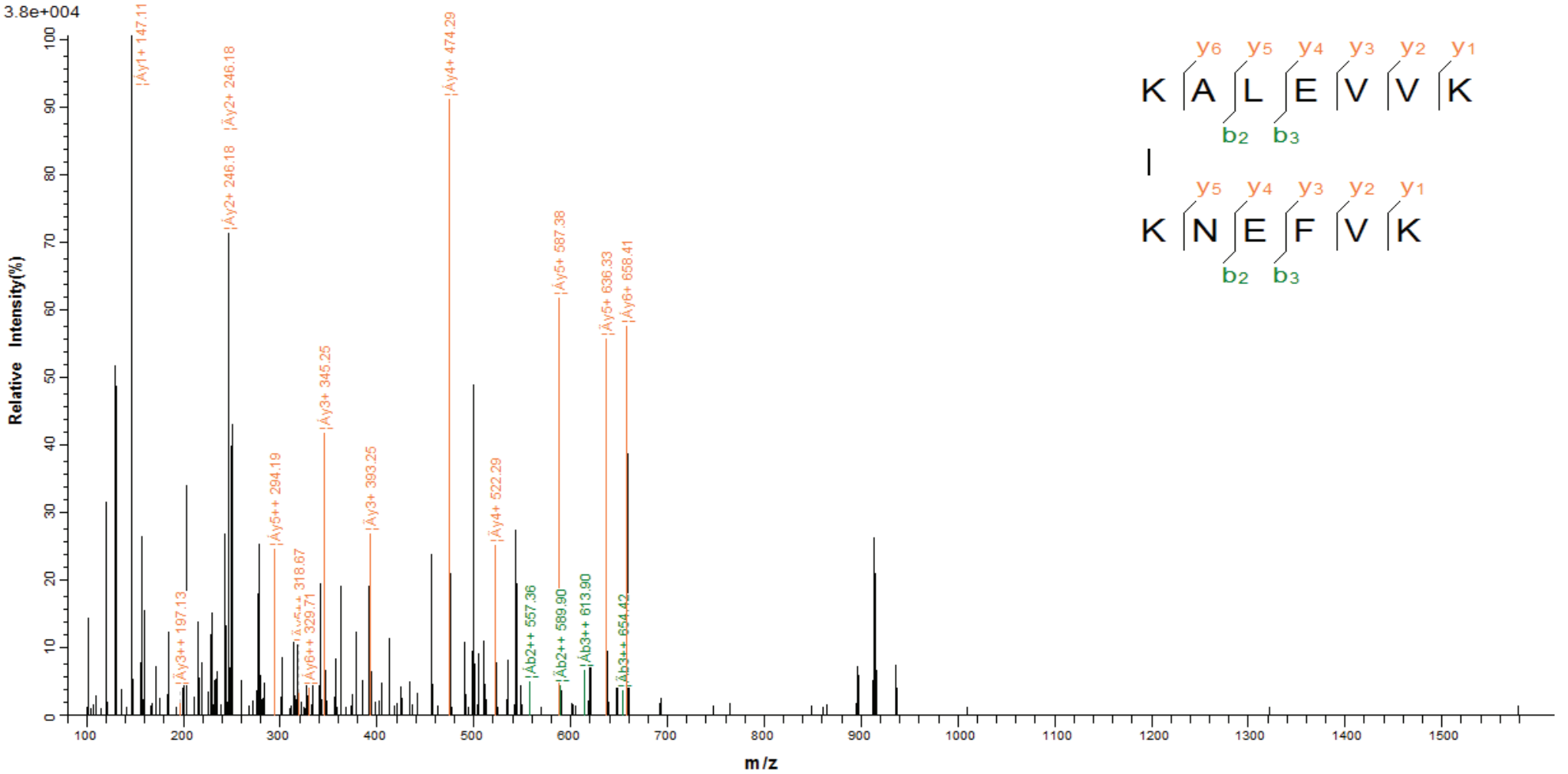
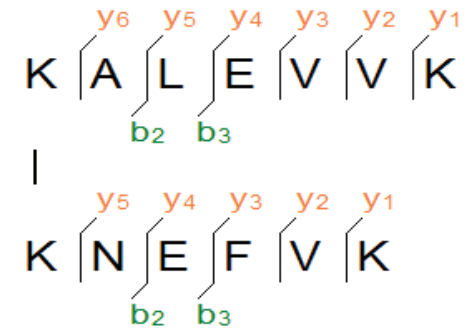


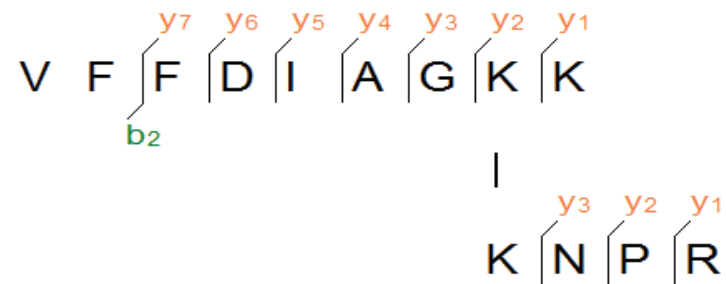
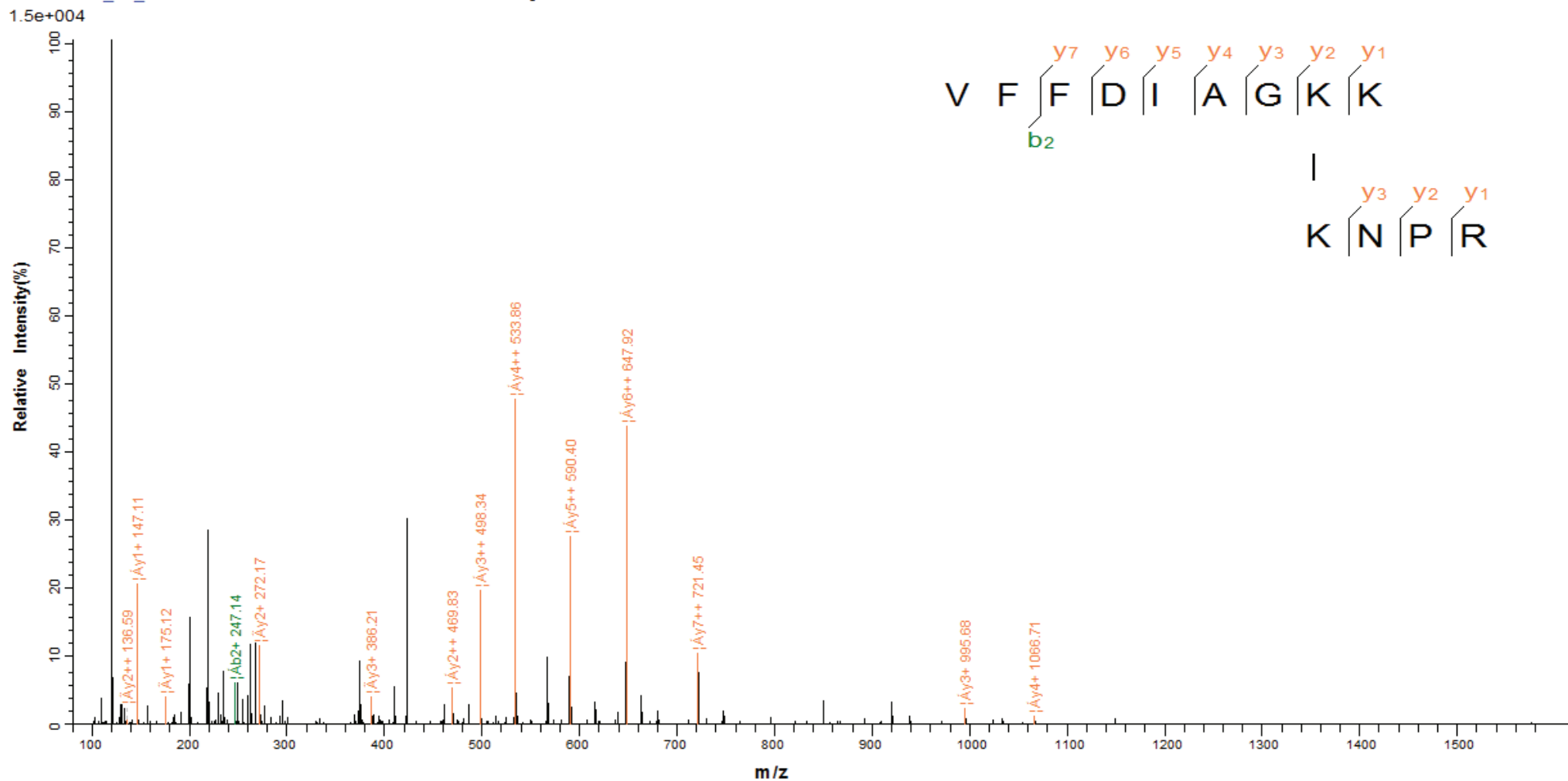
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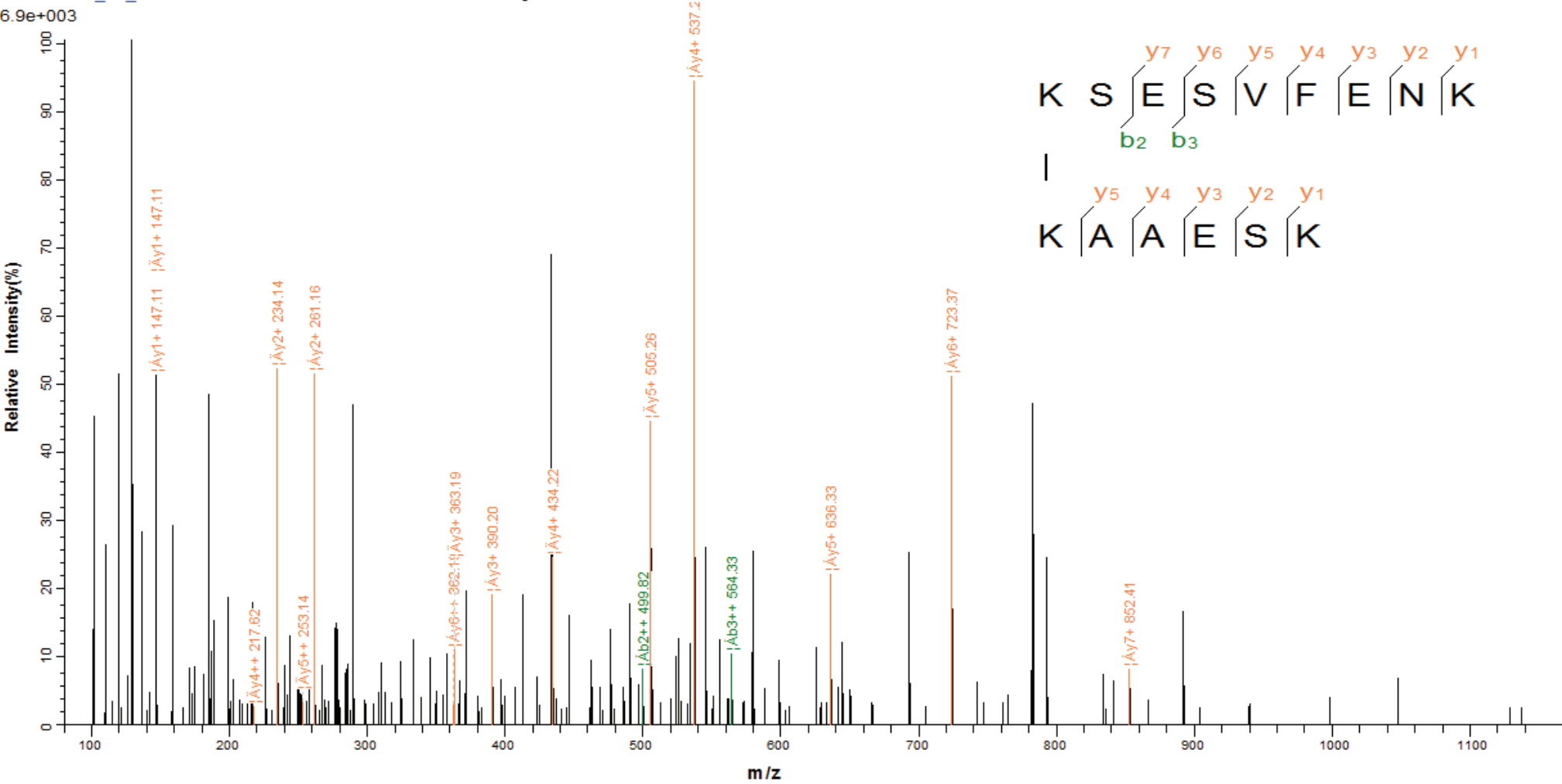


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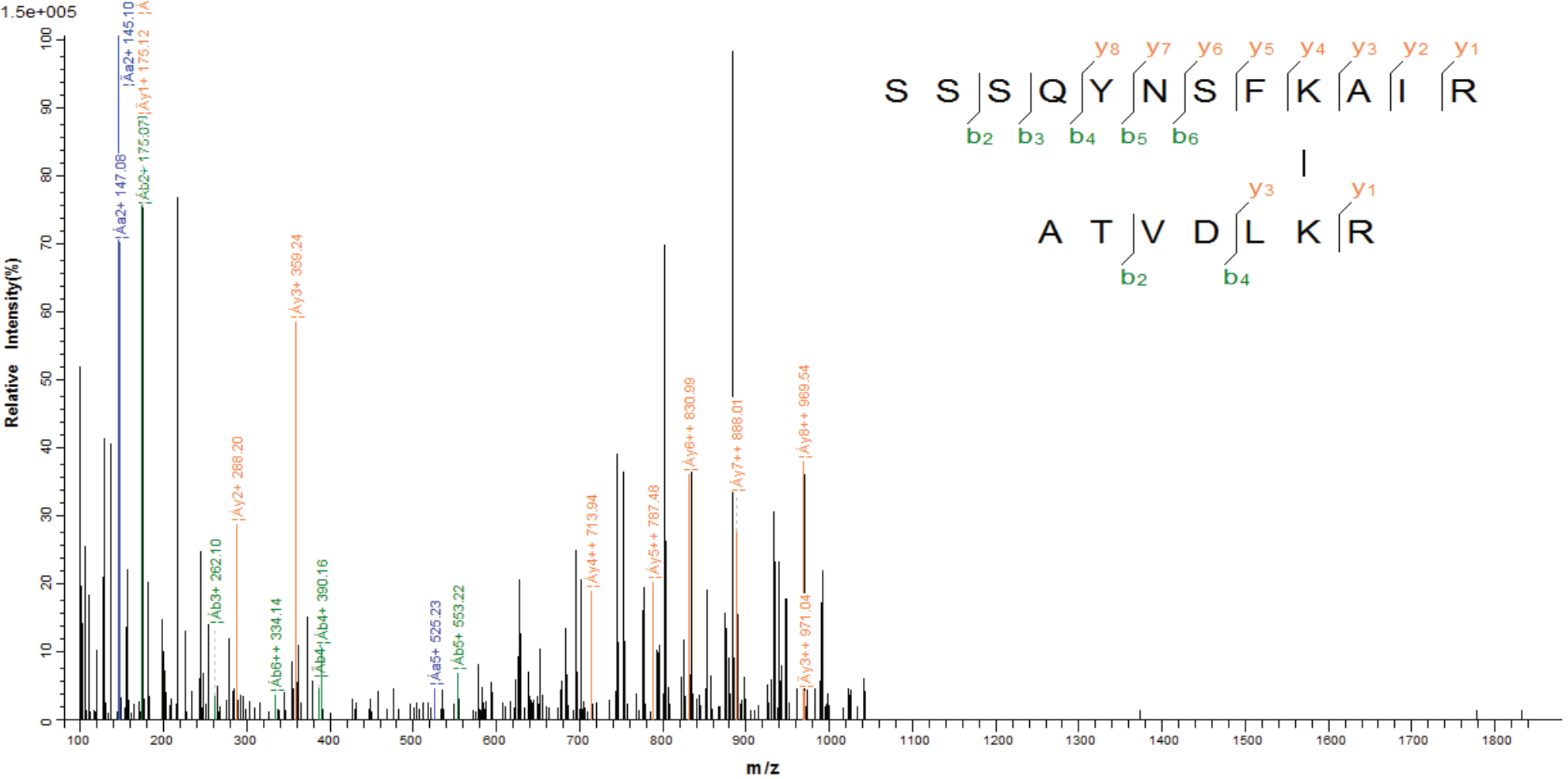
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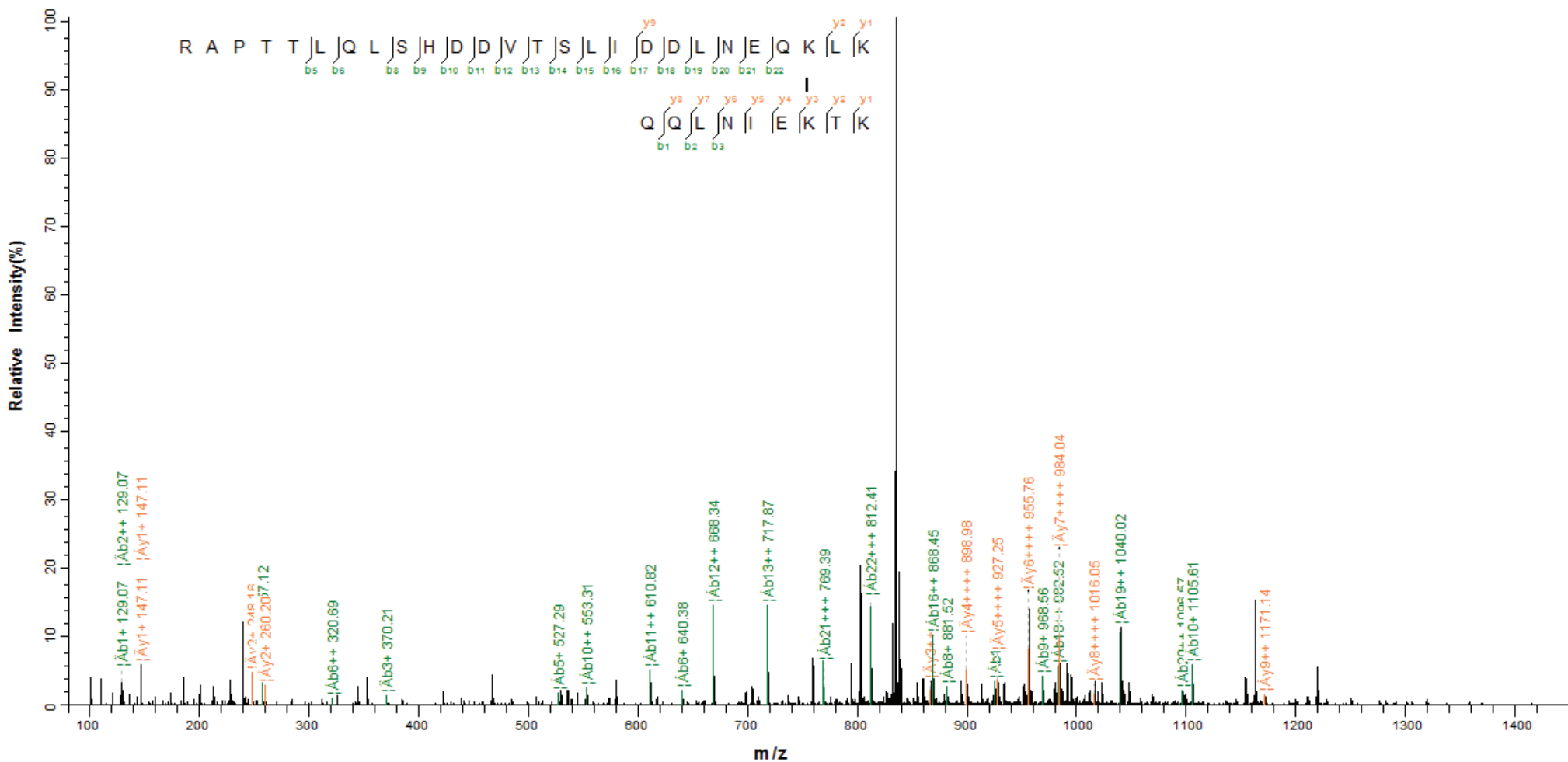


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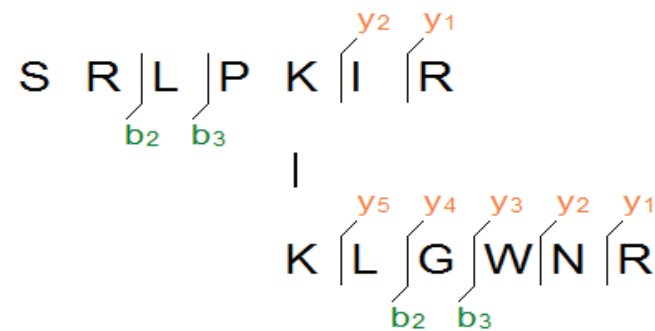
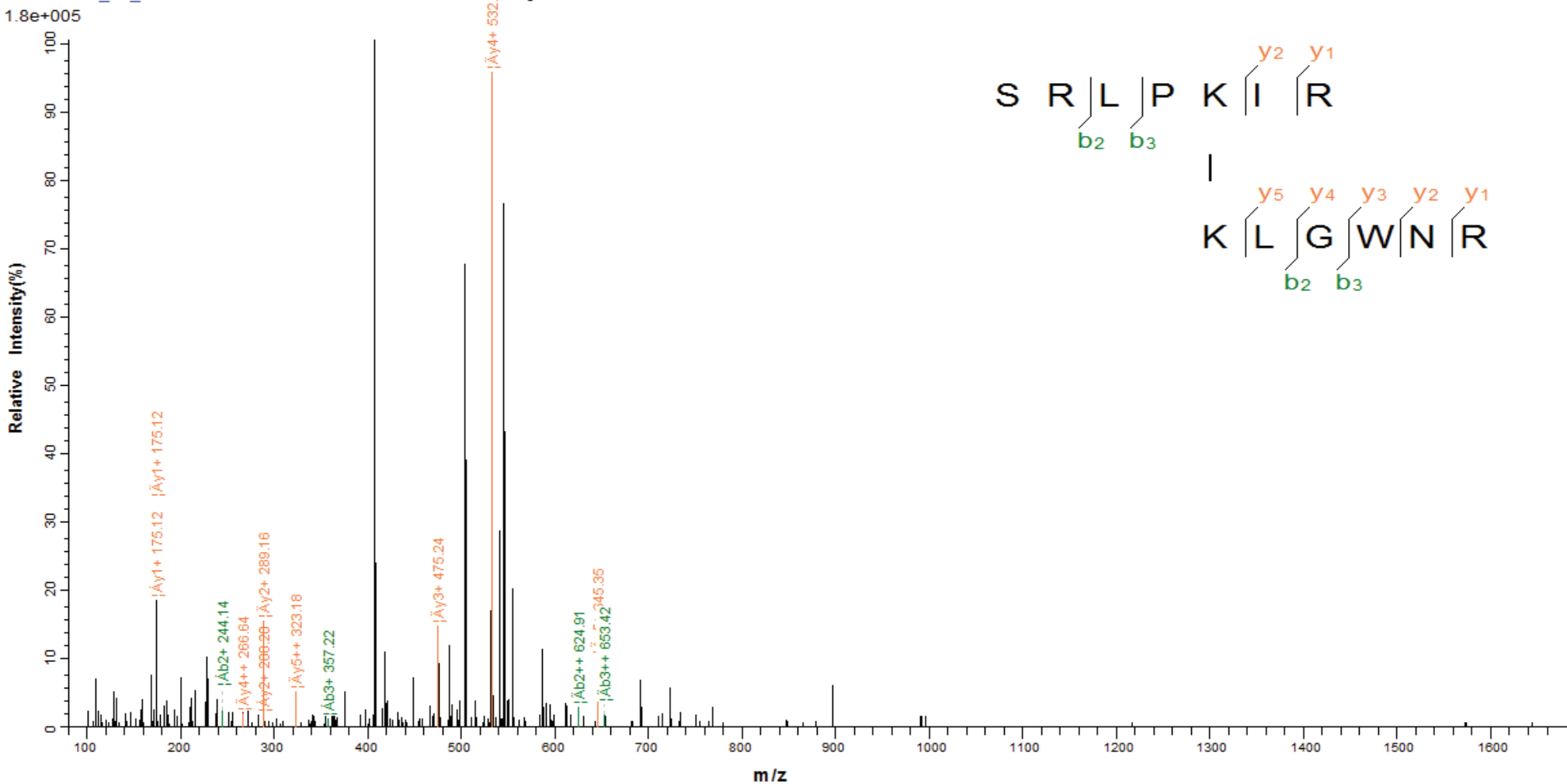


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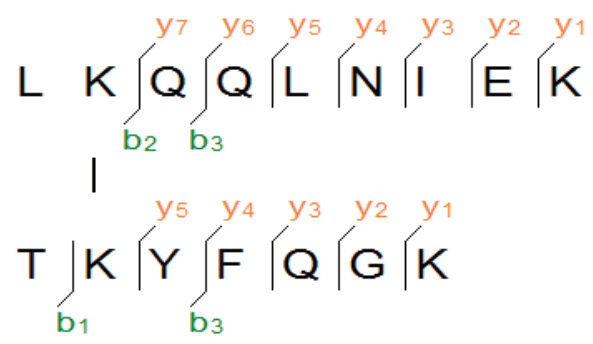
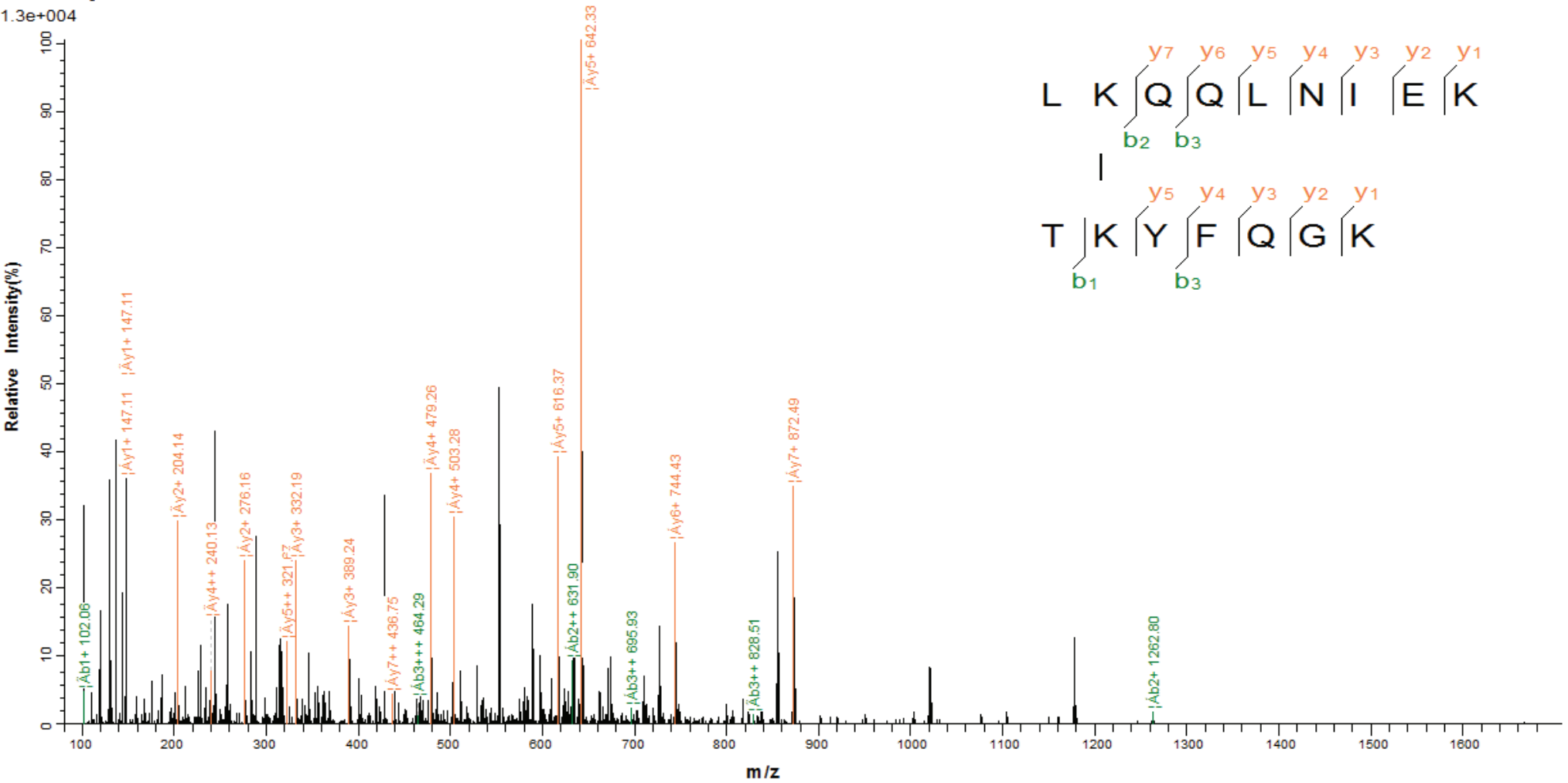
3.4e+004



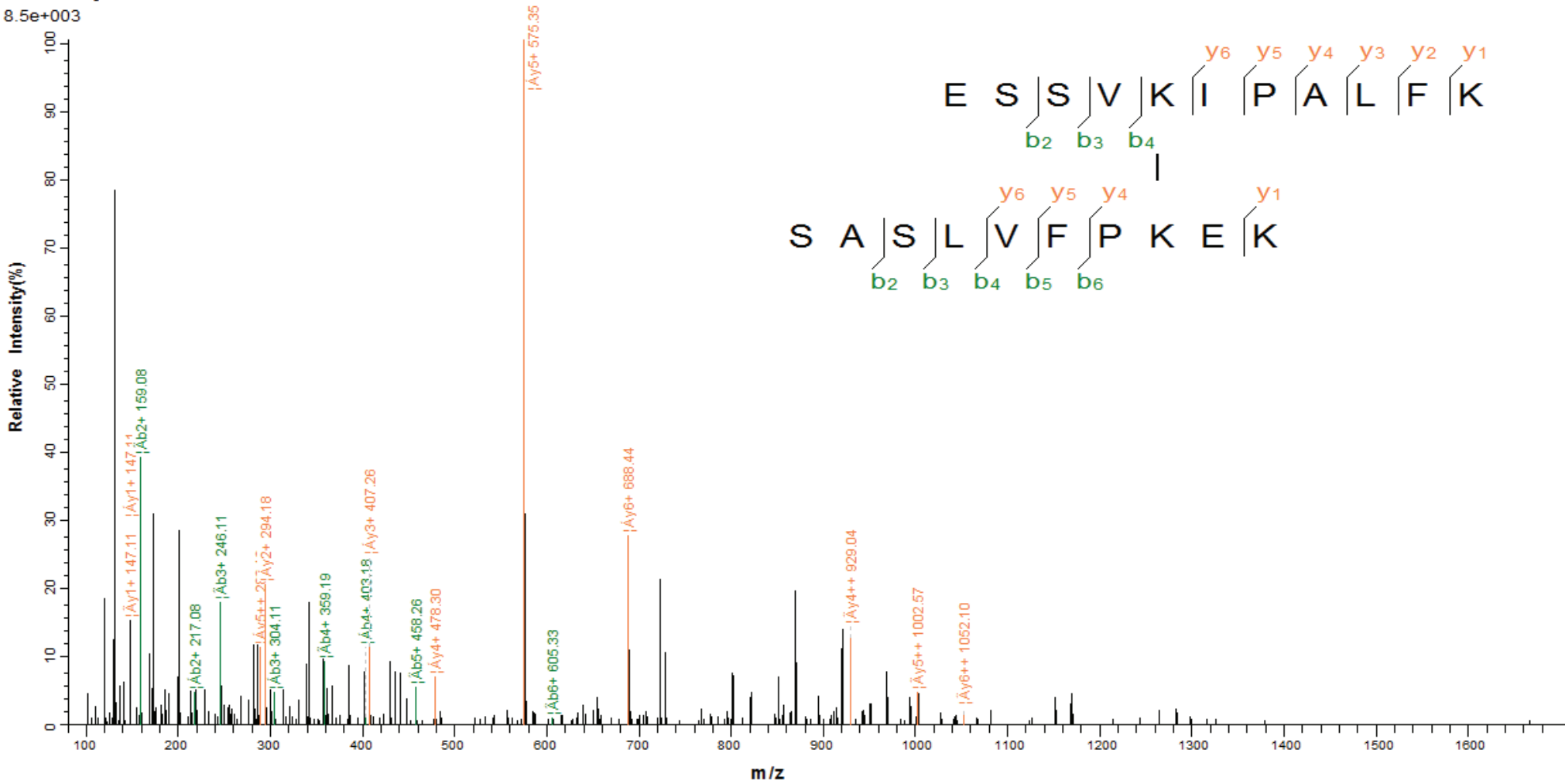
APC_A9_140131205754.004462.004462.4.dta Seq: SRLPKIR-KLGWNR



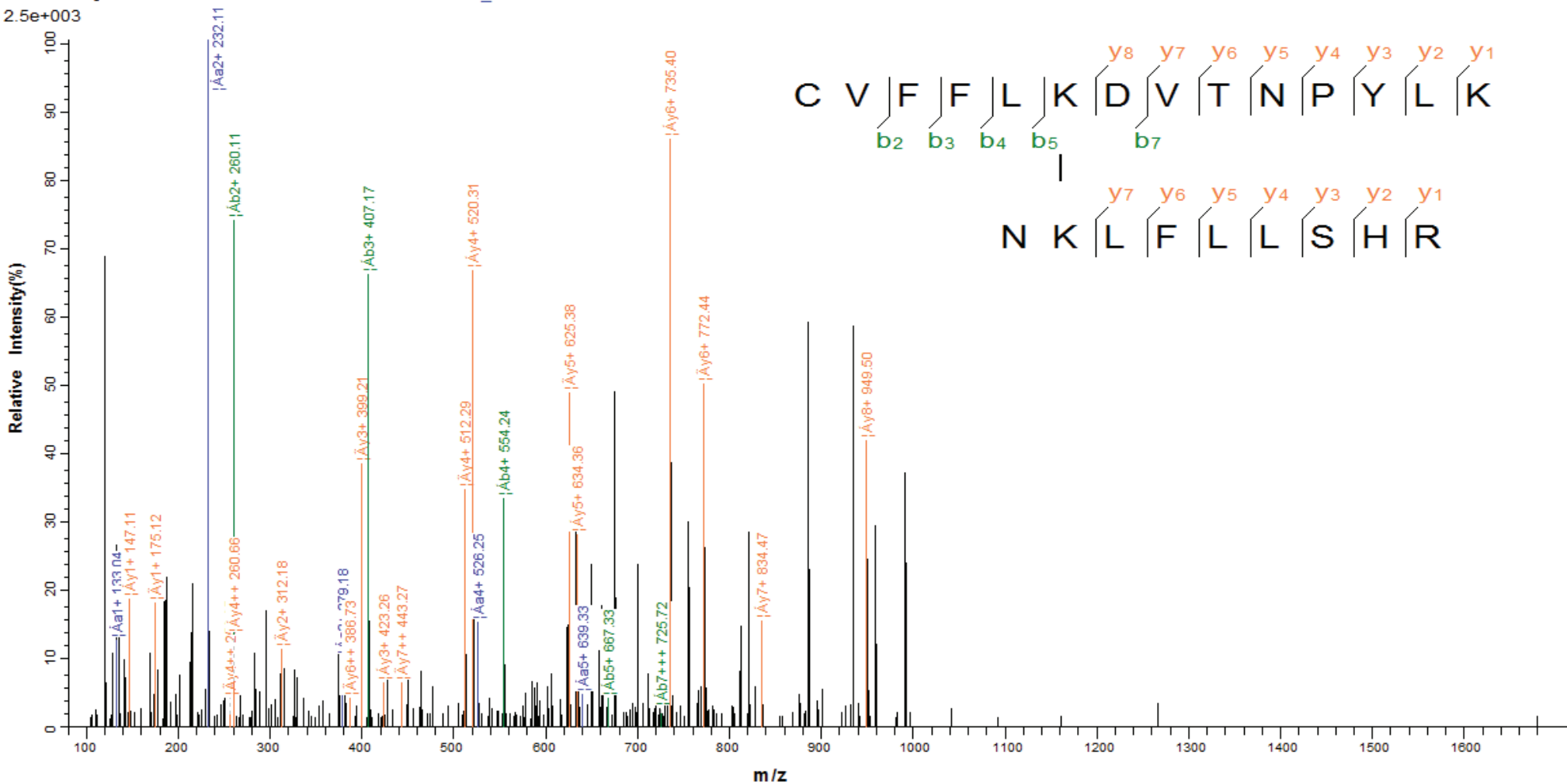
Seq: LKQQLNIEK-TRYFQ GK



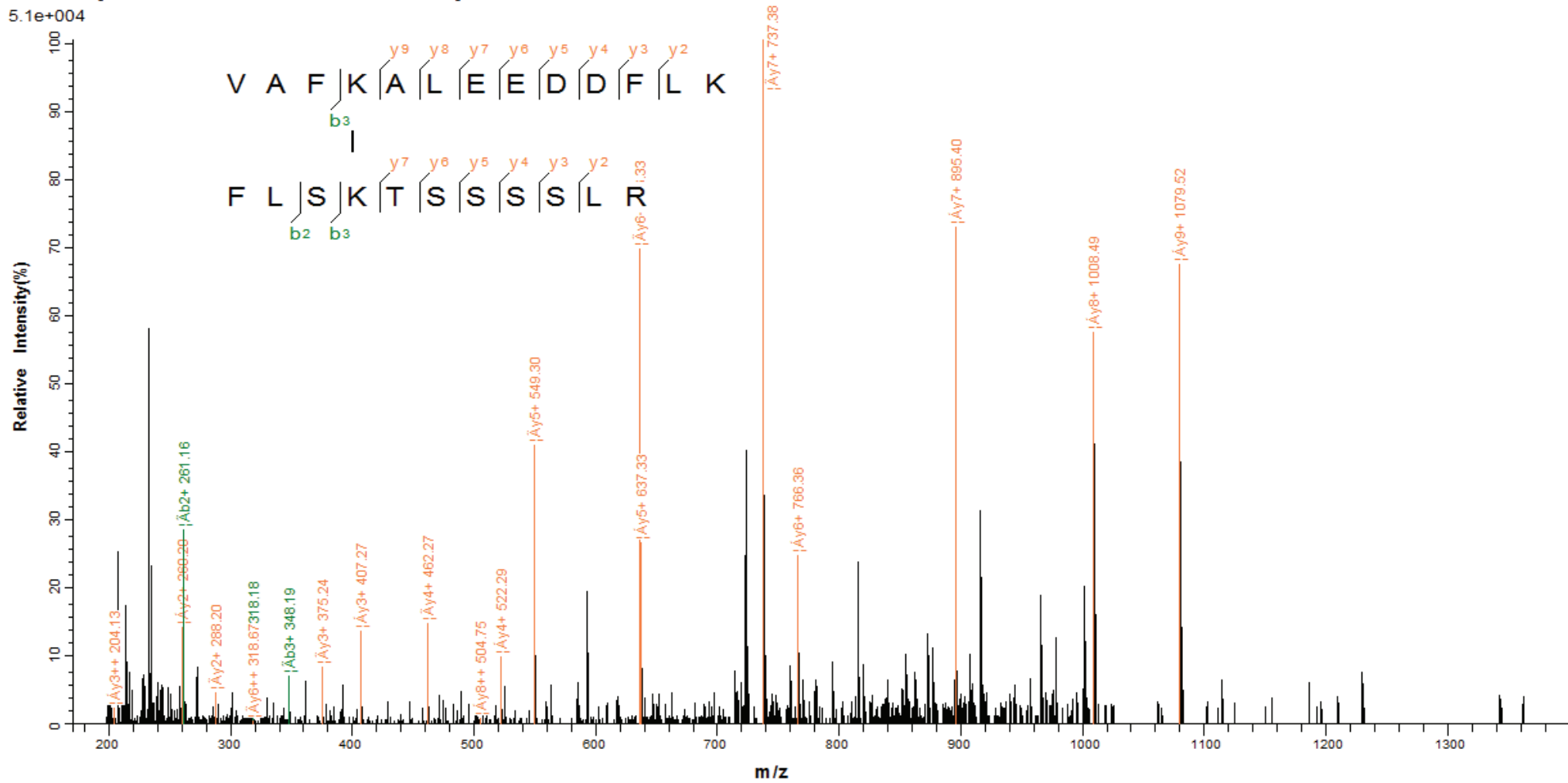
Seq: ESSVKIPALFK-SASLVFPKEK



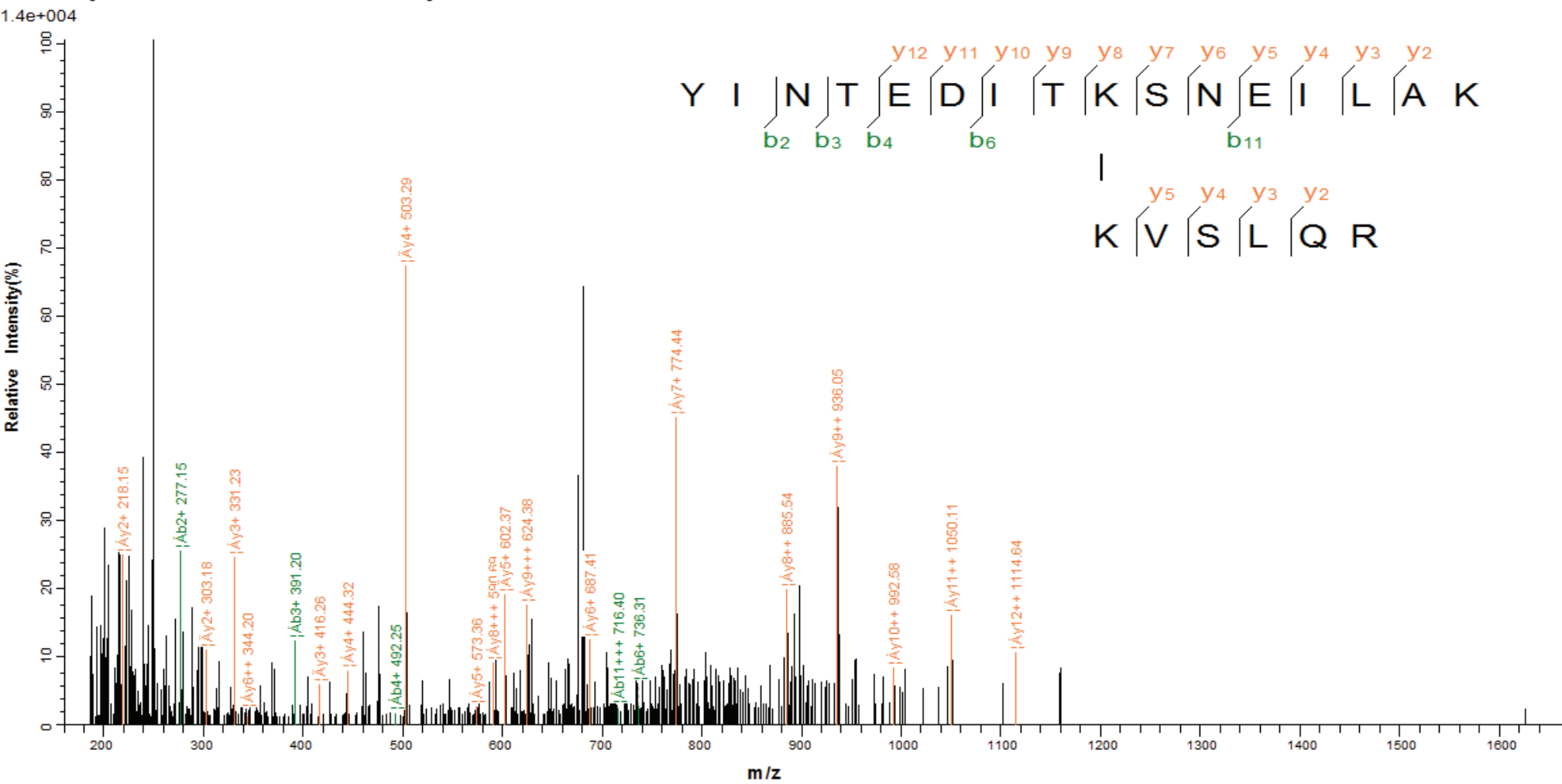
Seq: CVFFLKDVTFNPK-NKLFLLSHR Mod: 1,C_57



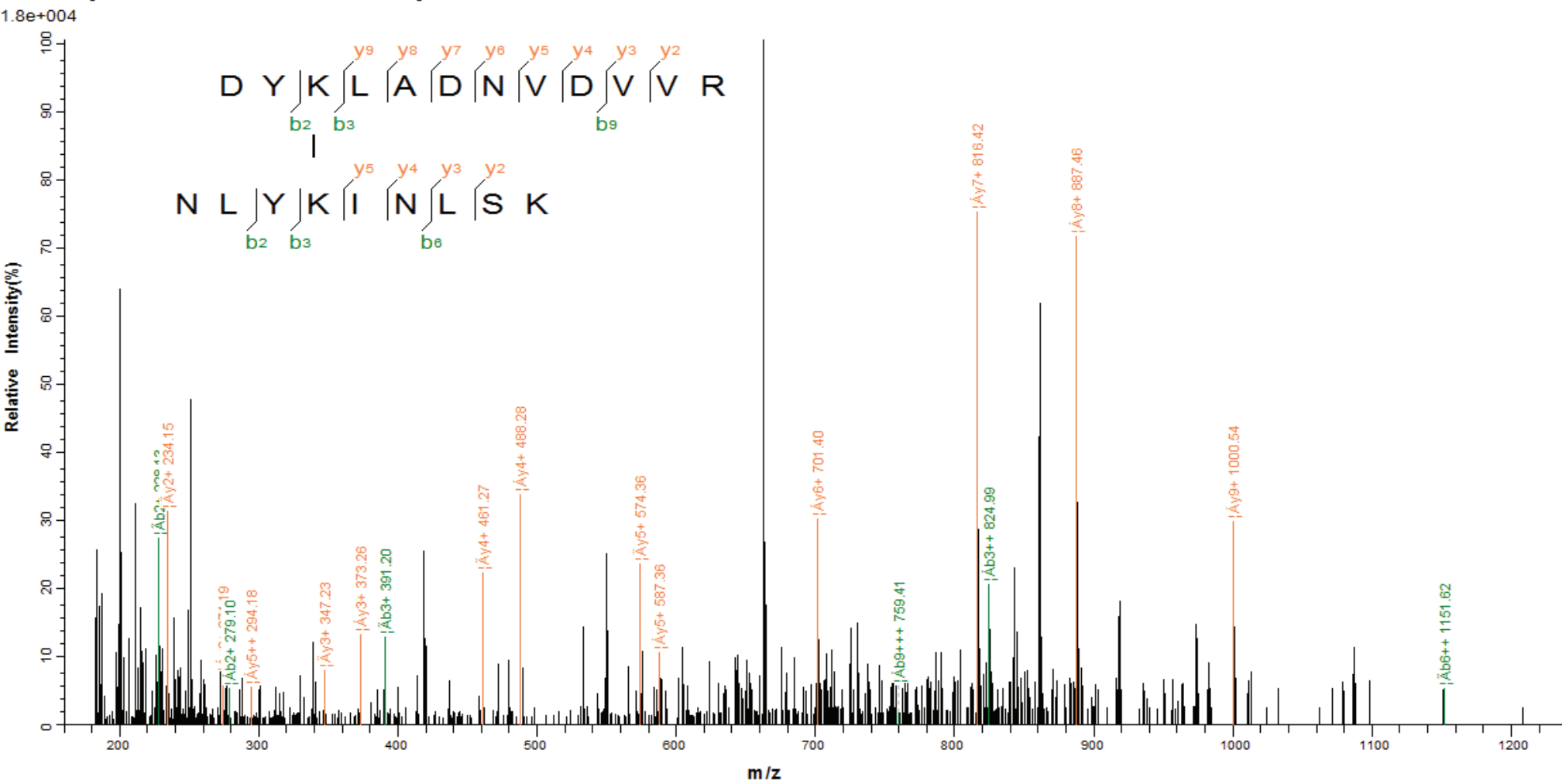
Seq: VAFKALEEDDFLK-FLSKTSSSSLR Charge: 4 MH+: 2887.5782



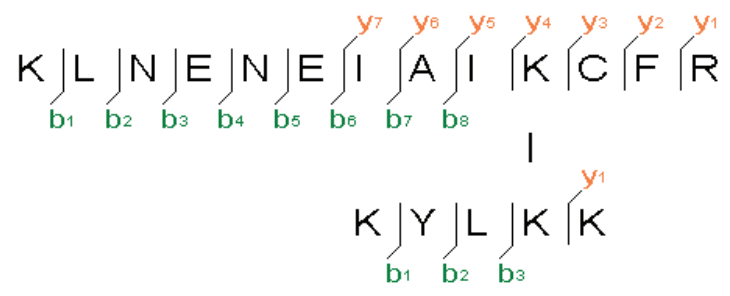
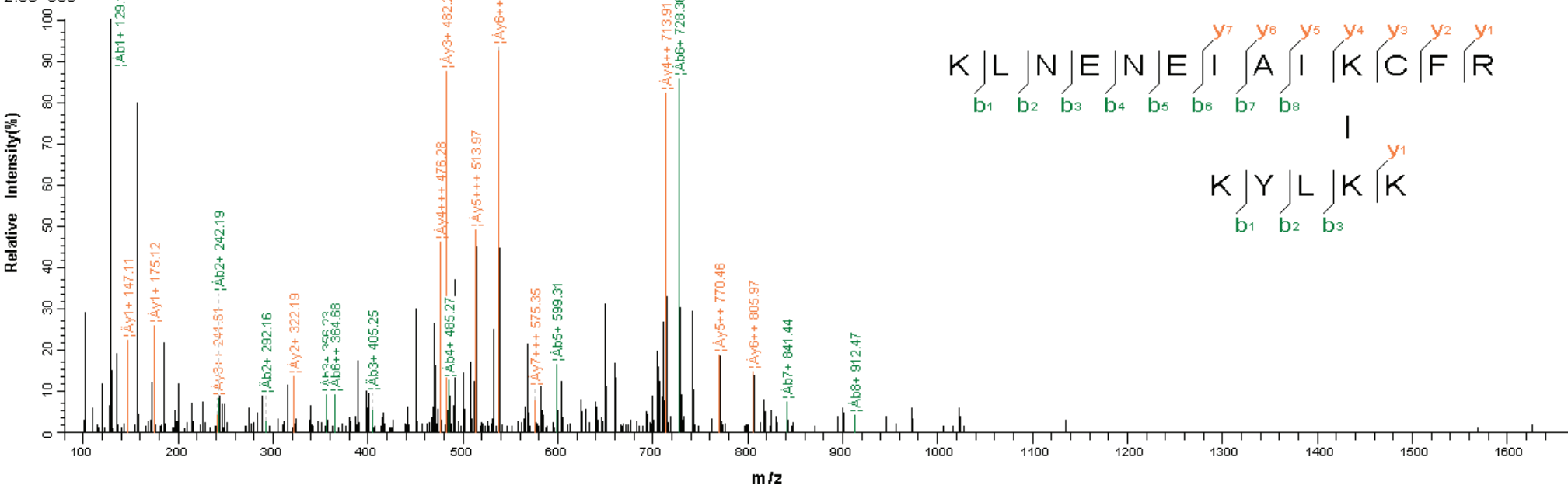
Seq: YINTEITKSNEILAK Charge: 4 MH+: 2720.4982



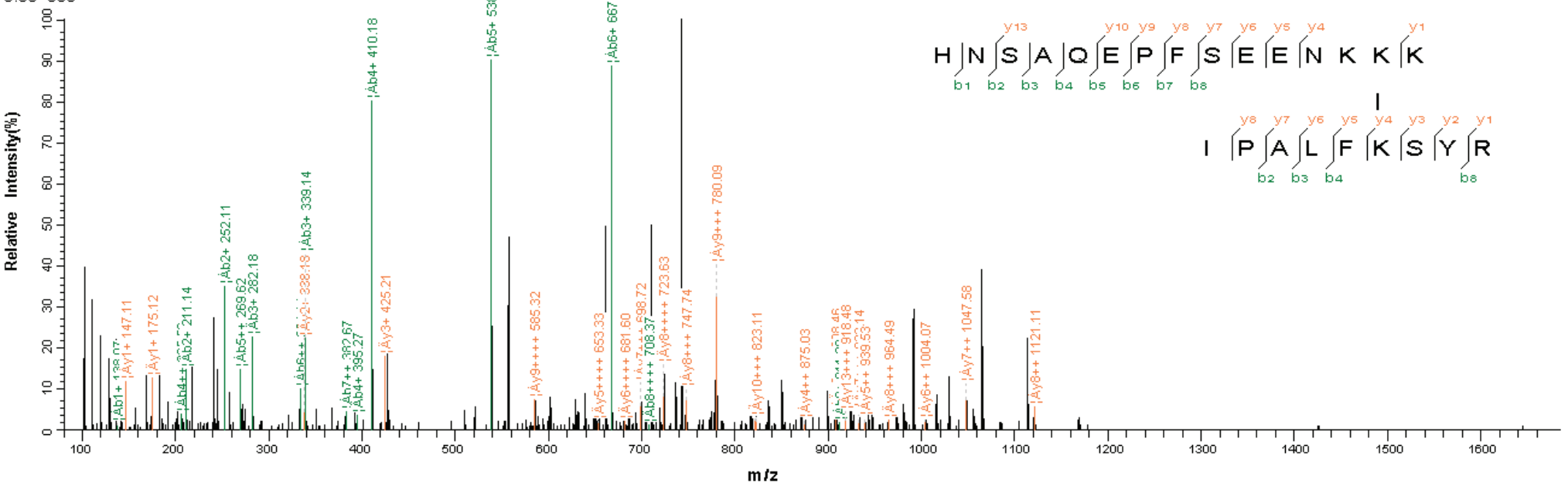
Seq: DYKLADNVDR-NLYKINLSK Charge: 4 MH+: 2649.4982



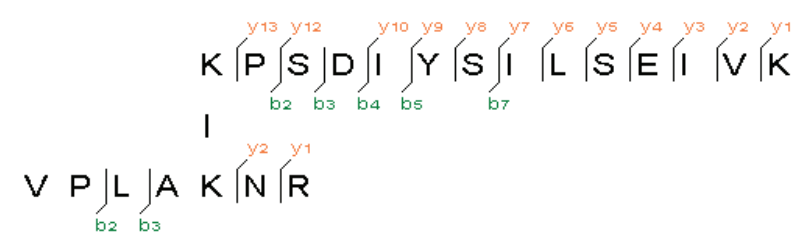
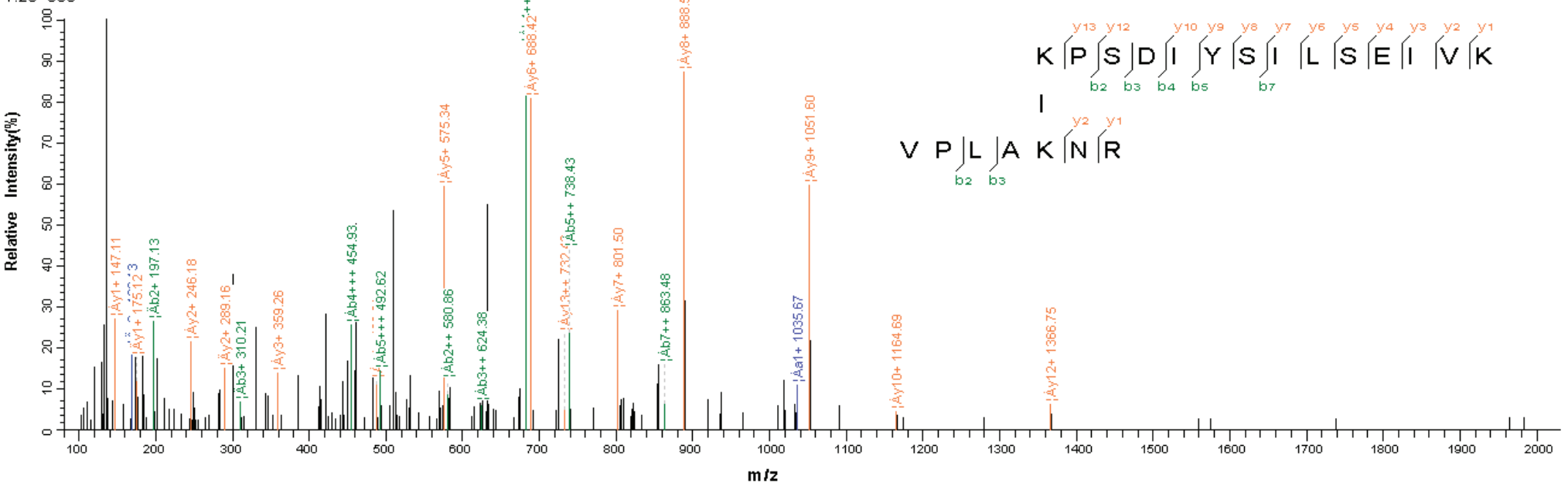
PC_A8_140201121226.003158.003158.5.dta Seq: KLNENEIAIKCFR-KYLKK Mod: 11,C_57



PC_A8_140201121226.003502.003502.5.dta Seq: HNSAQEPFSEENKKK-IPALFKYSR Score: 9.99e-006

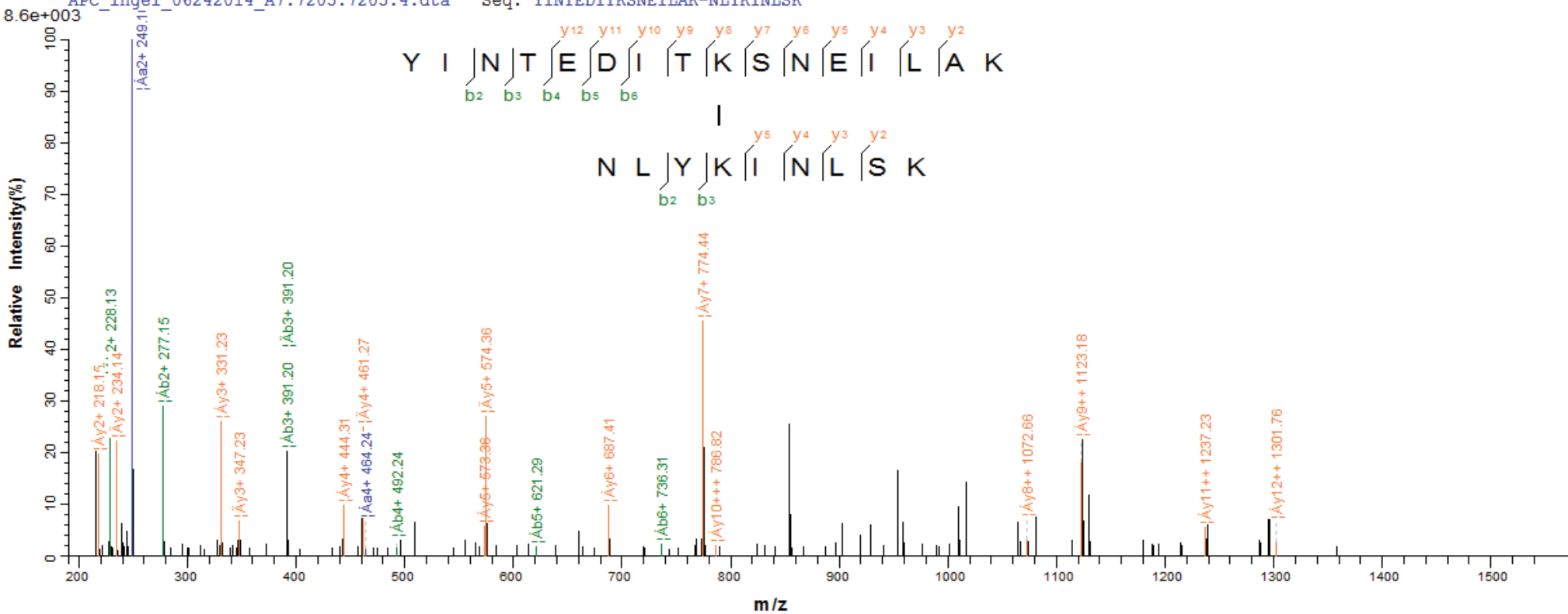
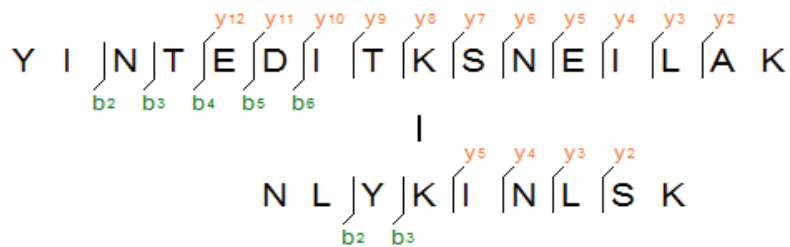


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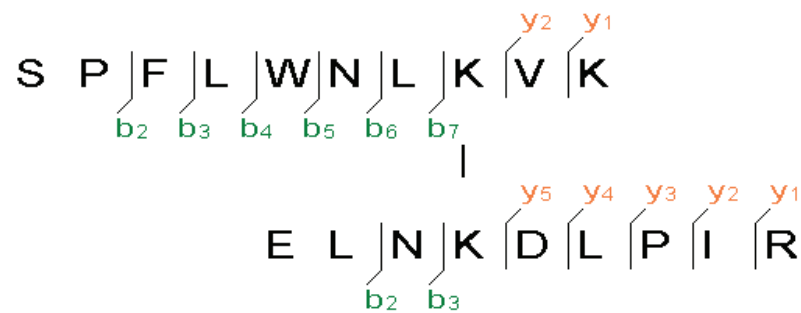
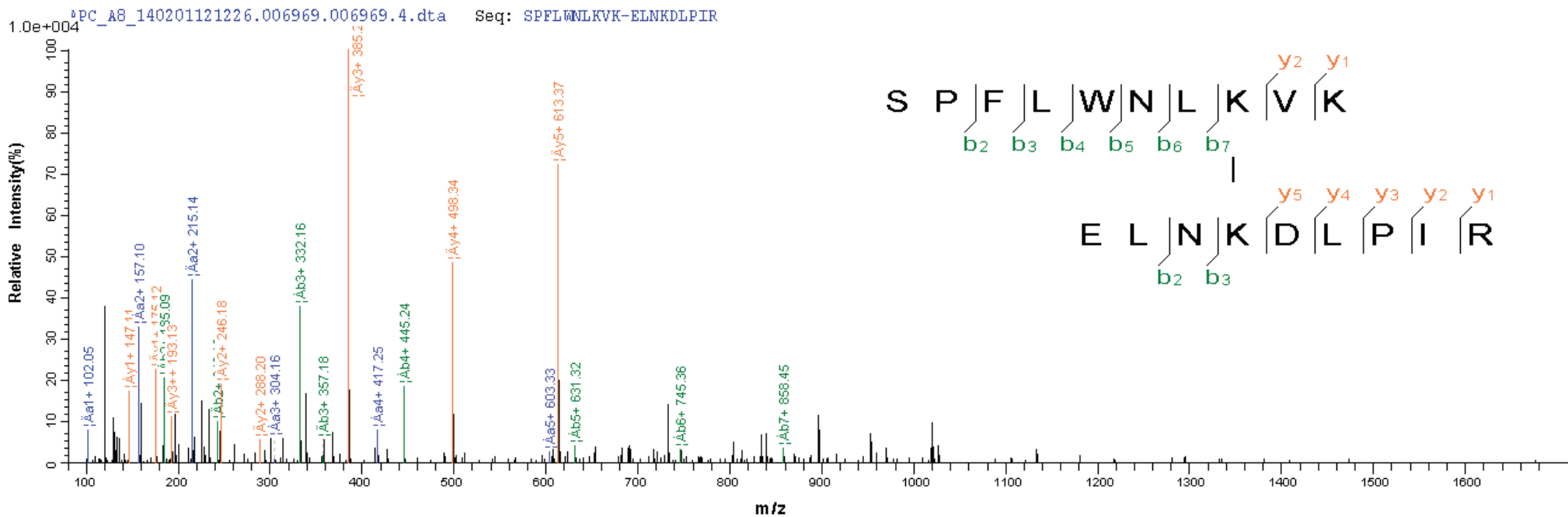


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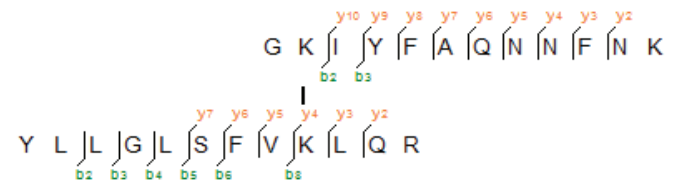
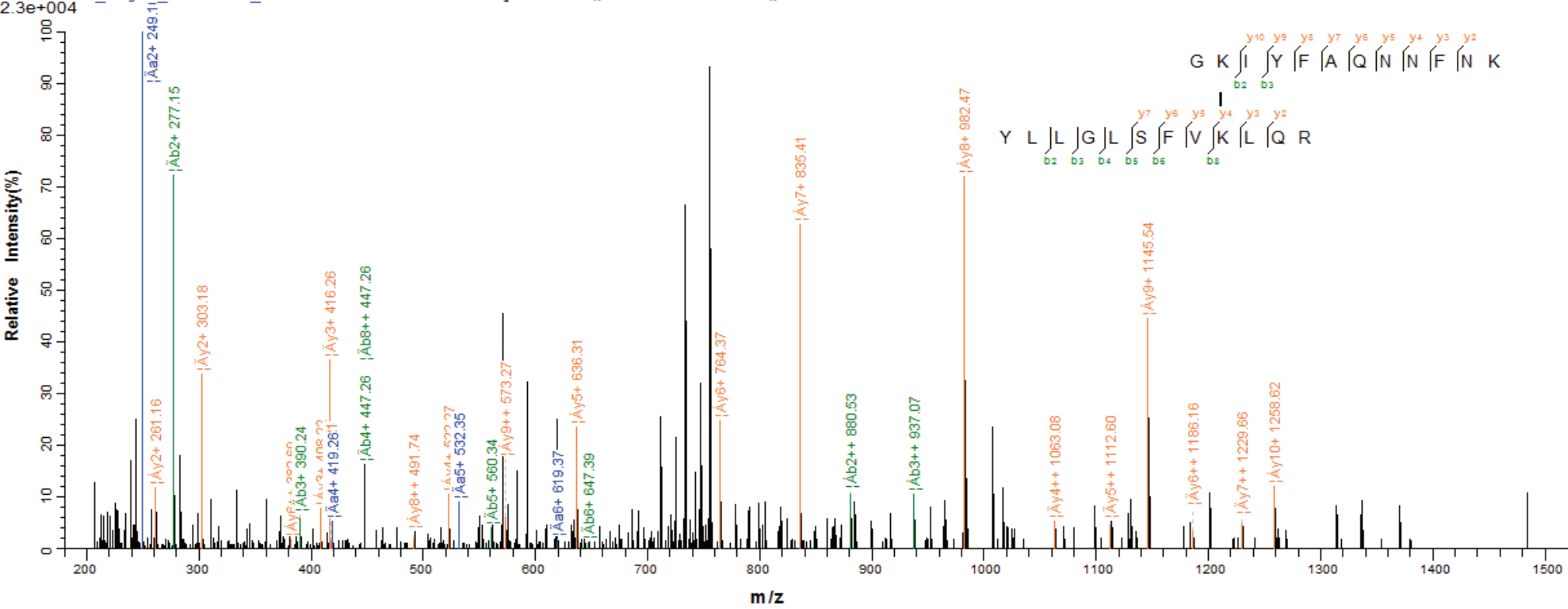
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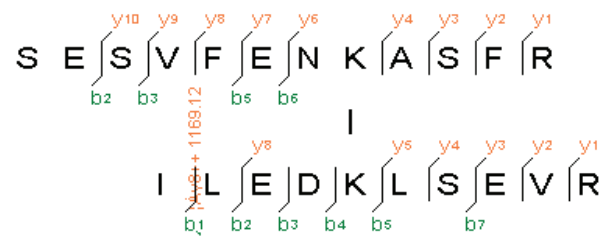
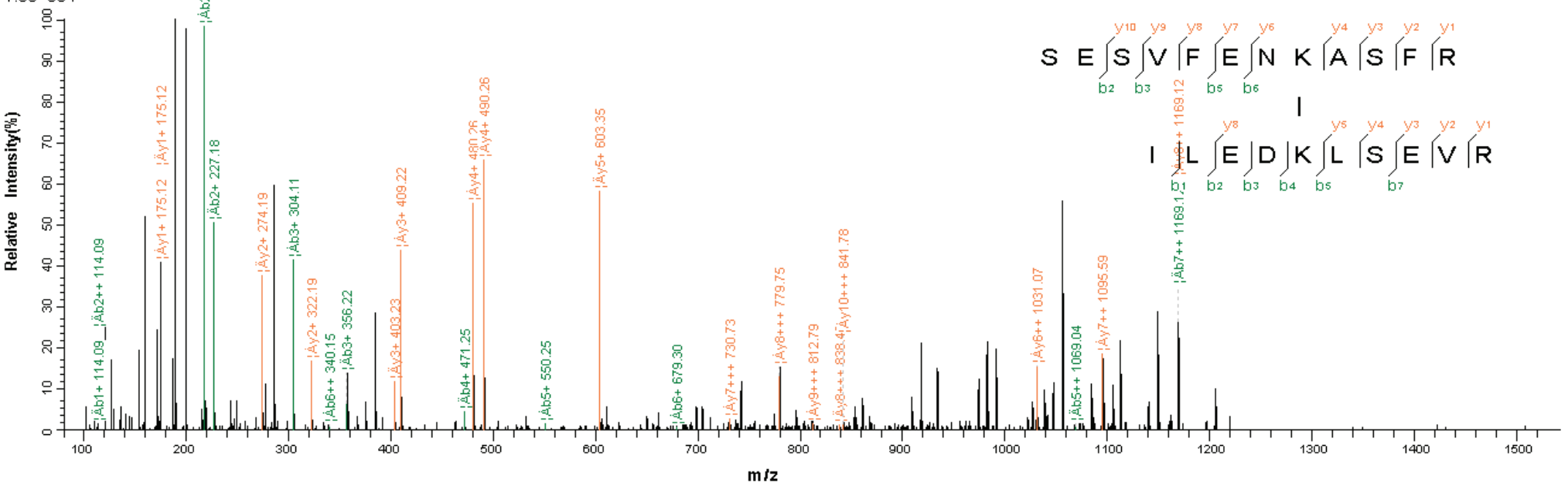
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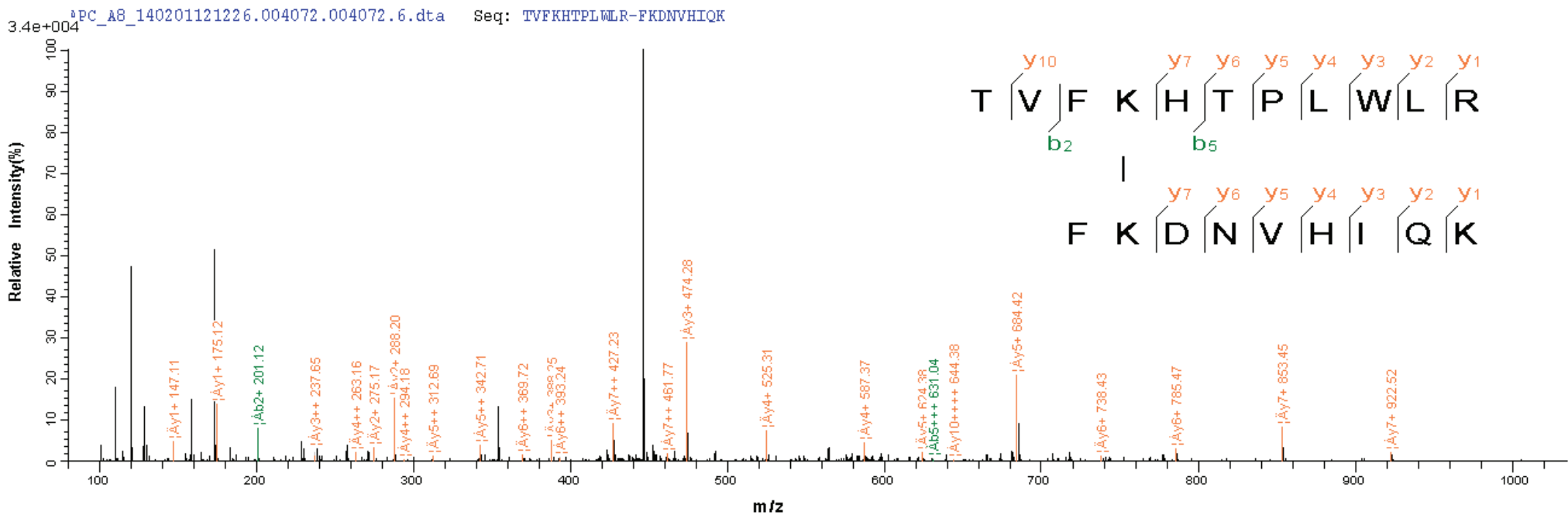
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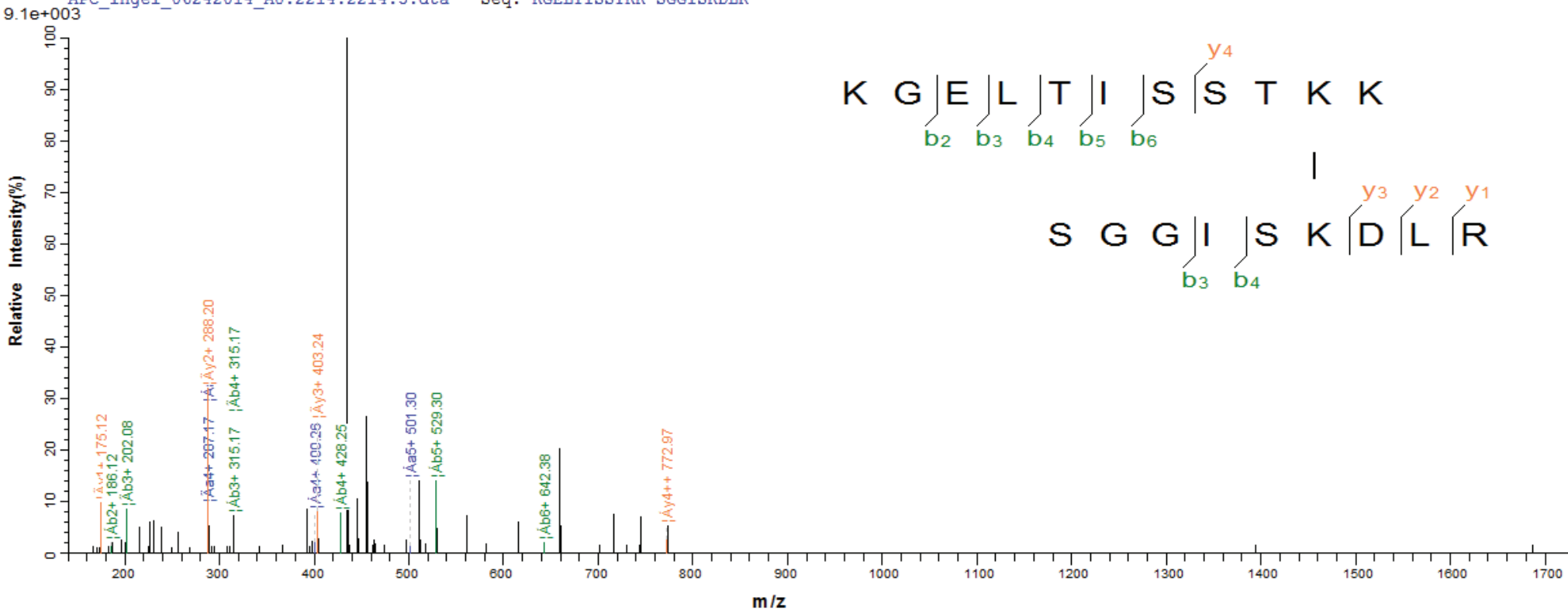


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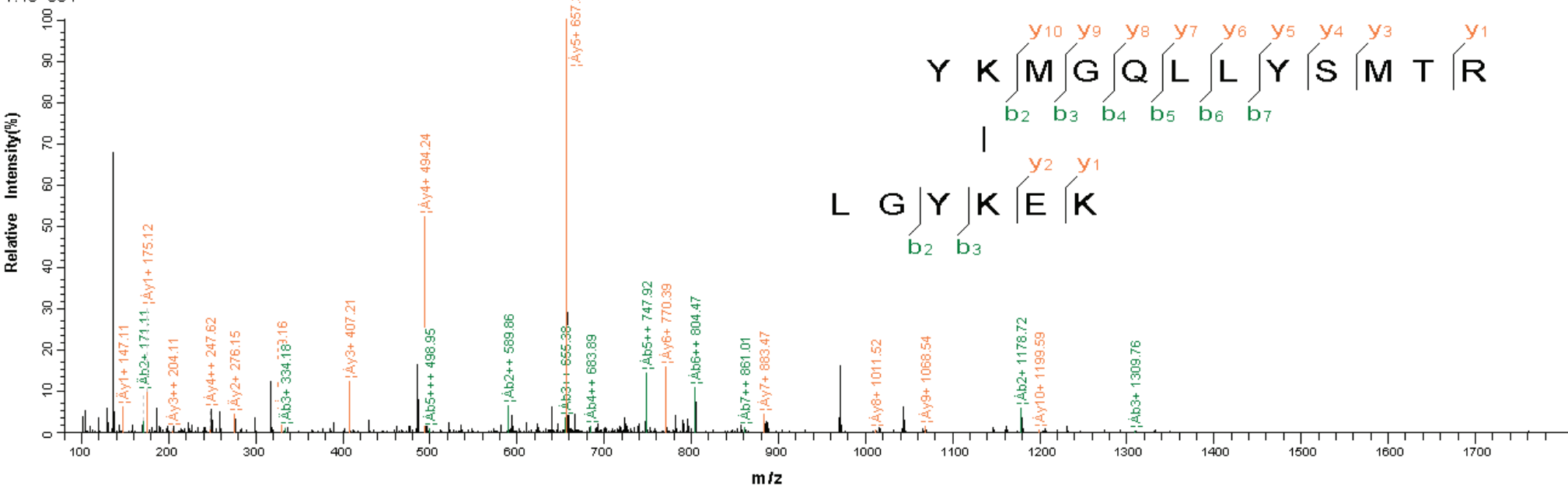


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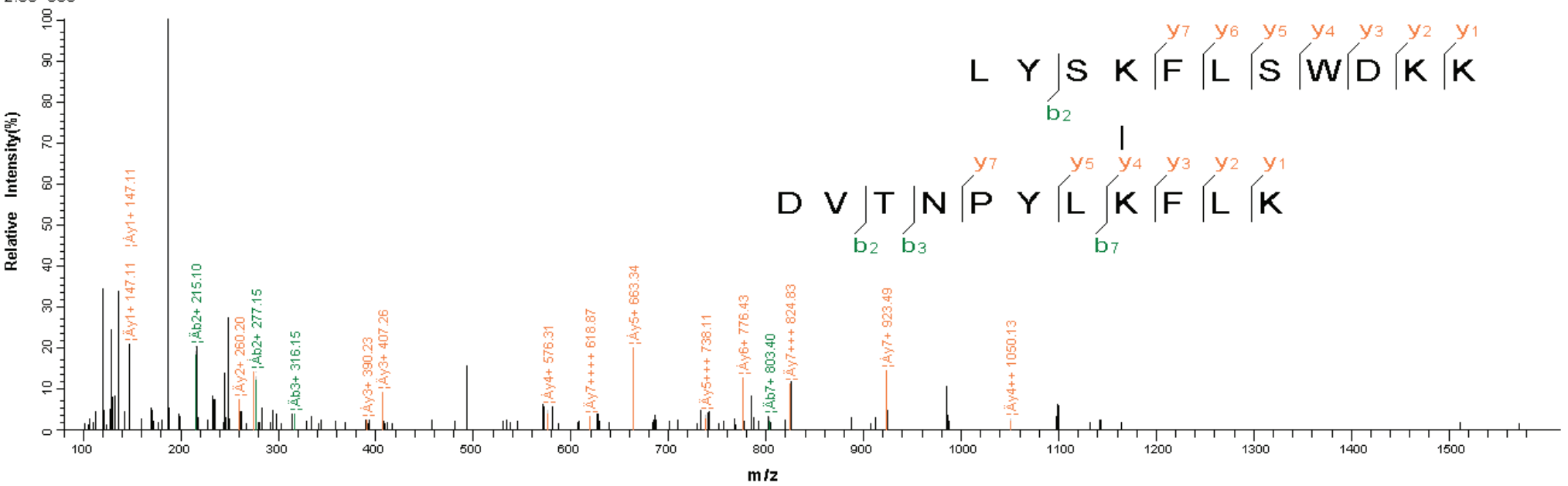




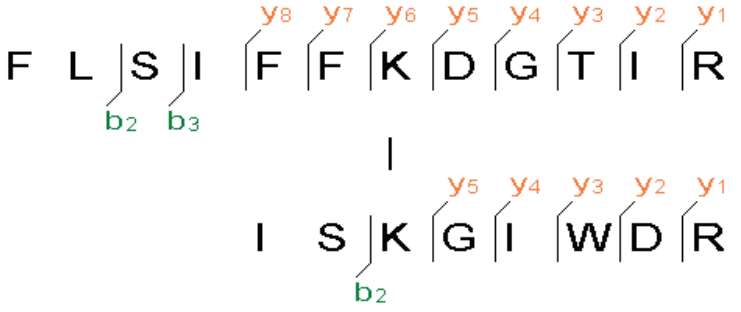
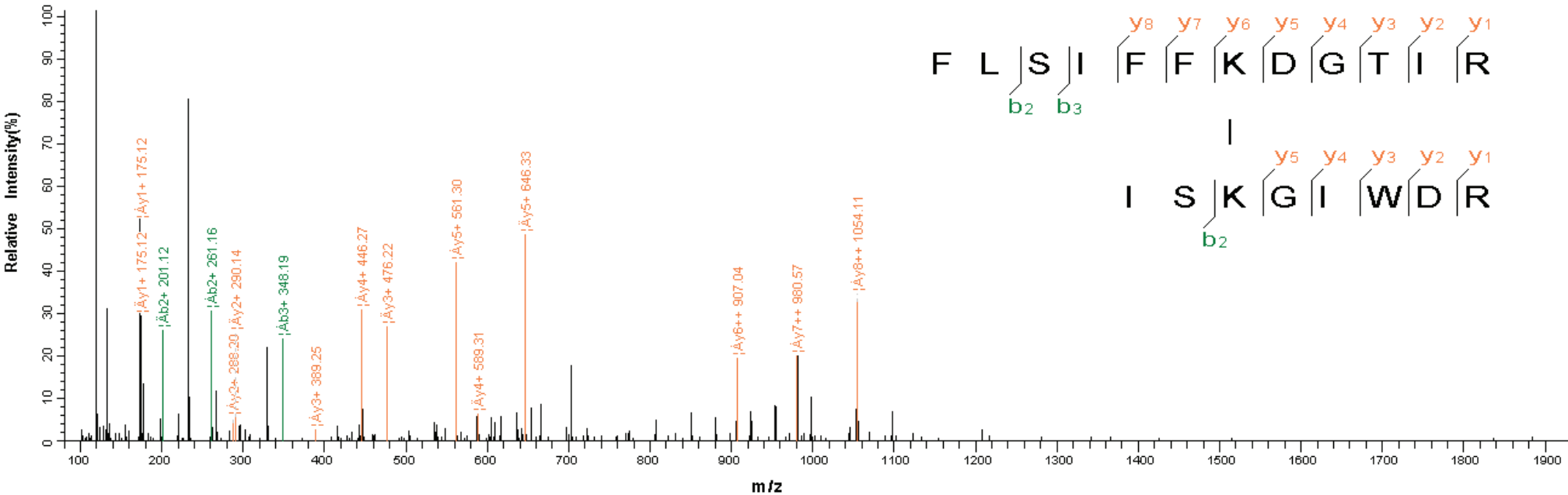
PC_A8_140201121226.005684.005684.4.dta Seq: YKMGQLLYSMTR-LGYKEK

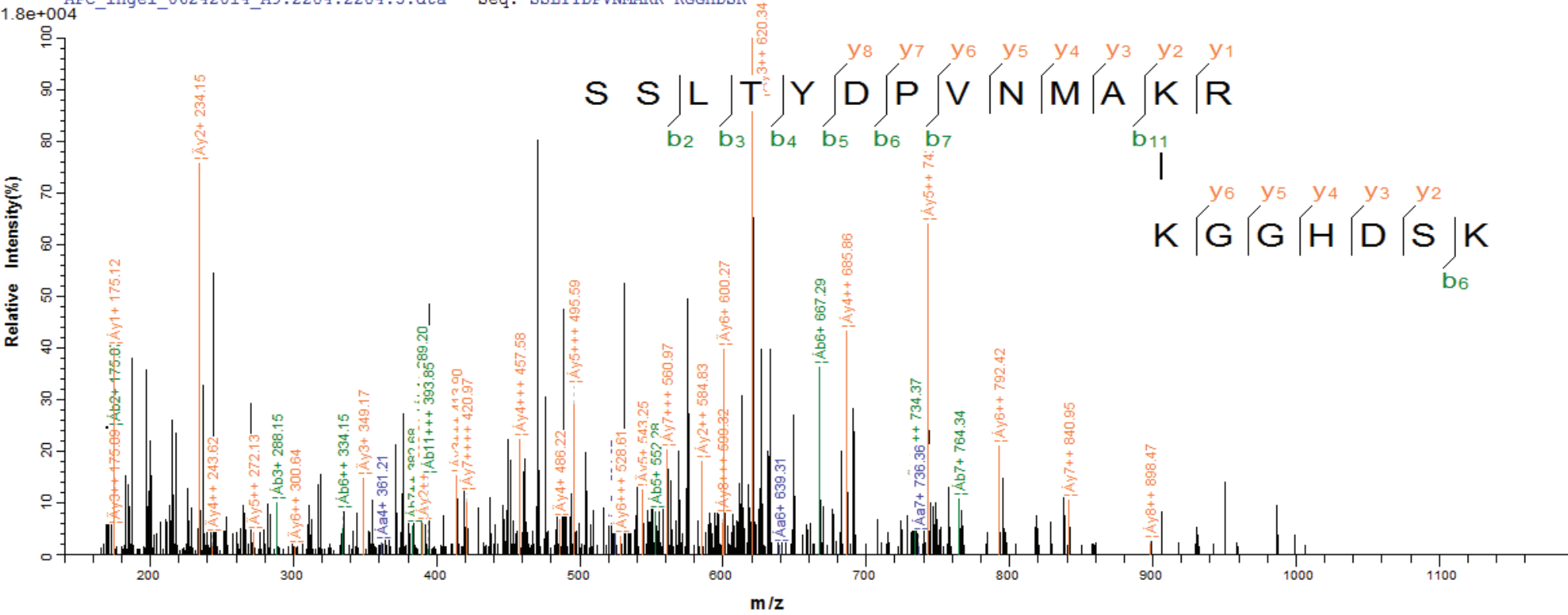


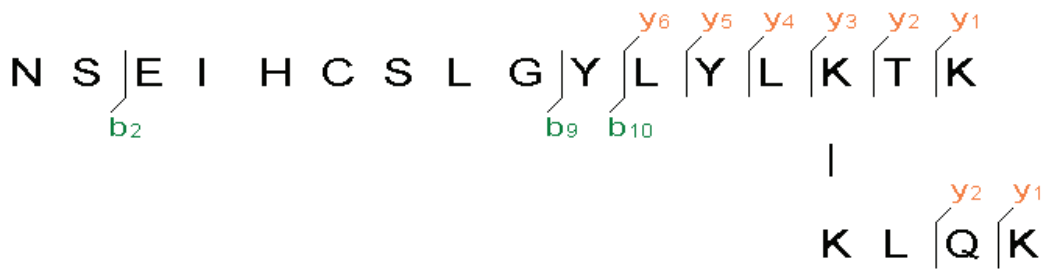
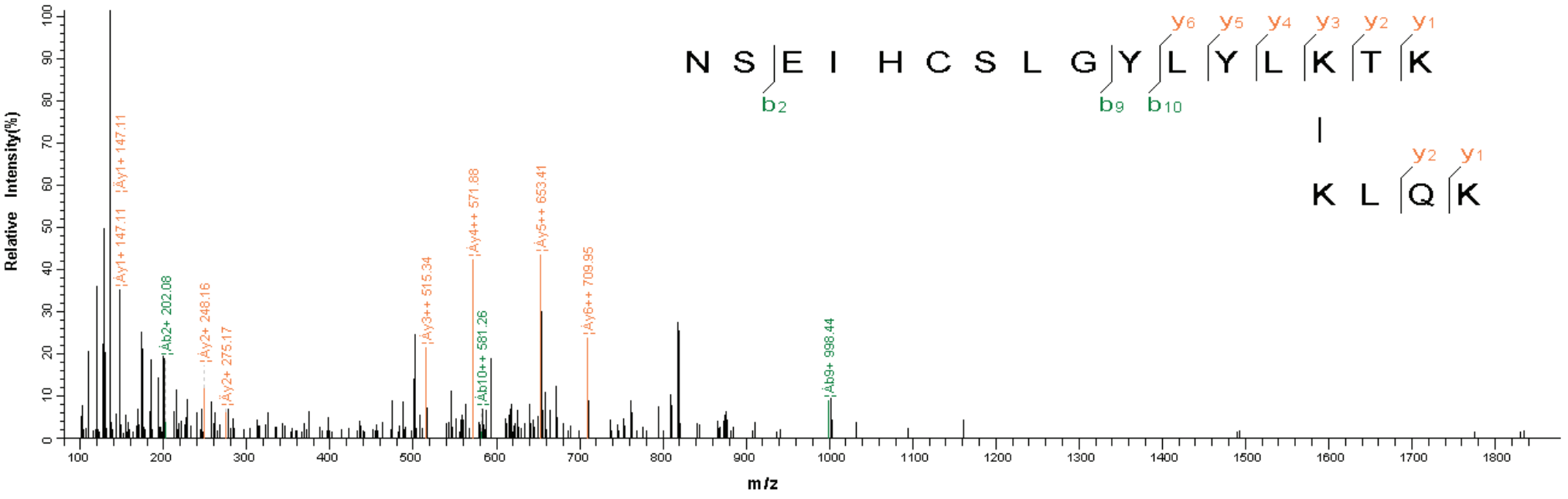
PC_A8_140201121226.007602.007602.5.dta Seq: LYSKFLSWDKK-DVTNPYLKFLK



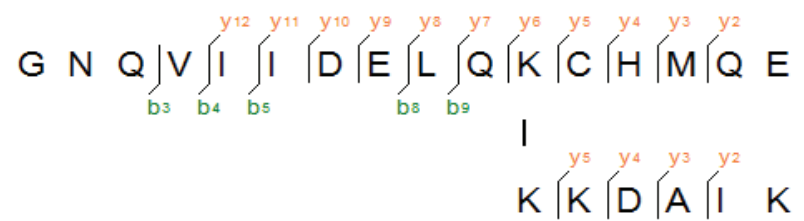
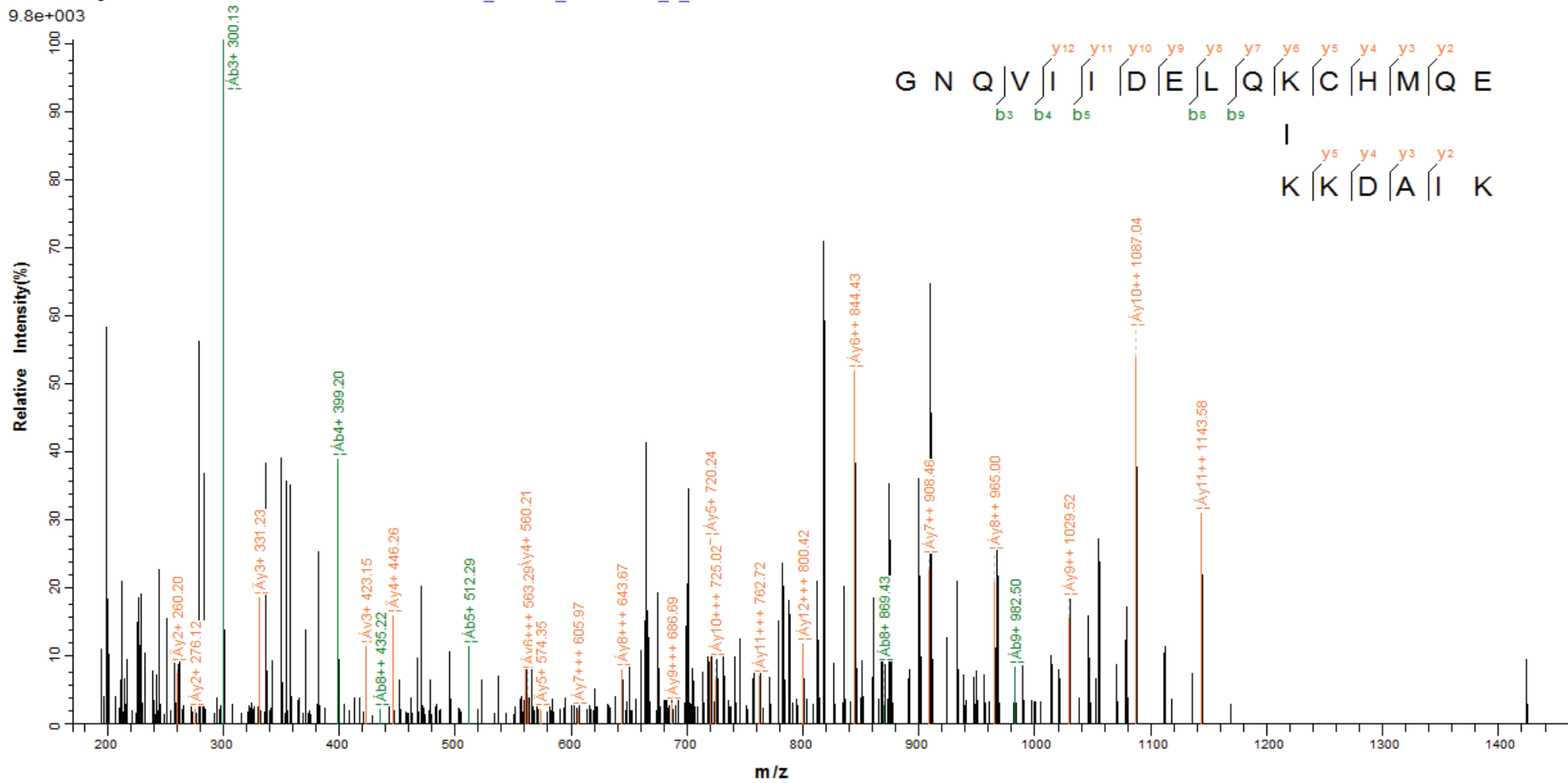
4.6e+003 PC_A8_140201121226.007697.007697.4.dta Seq: FLSIFFKDG TIR-ISKGIWDR



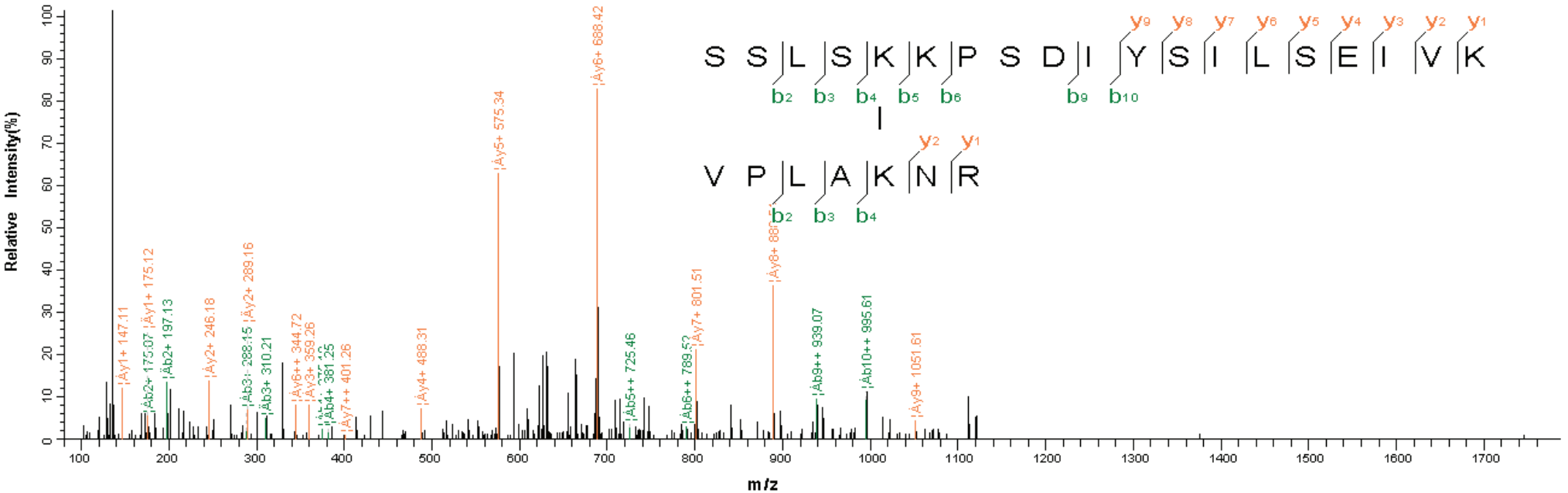


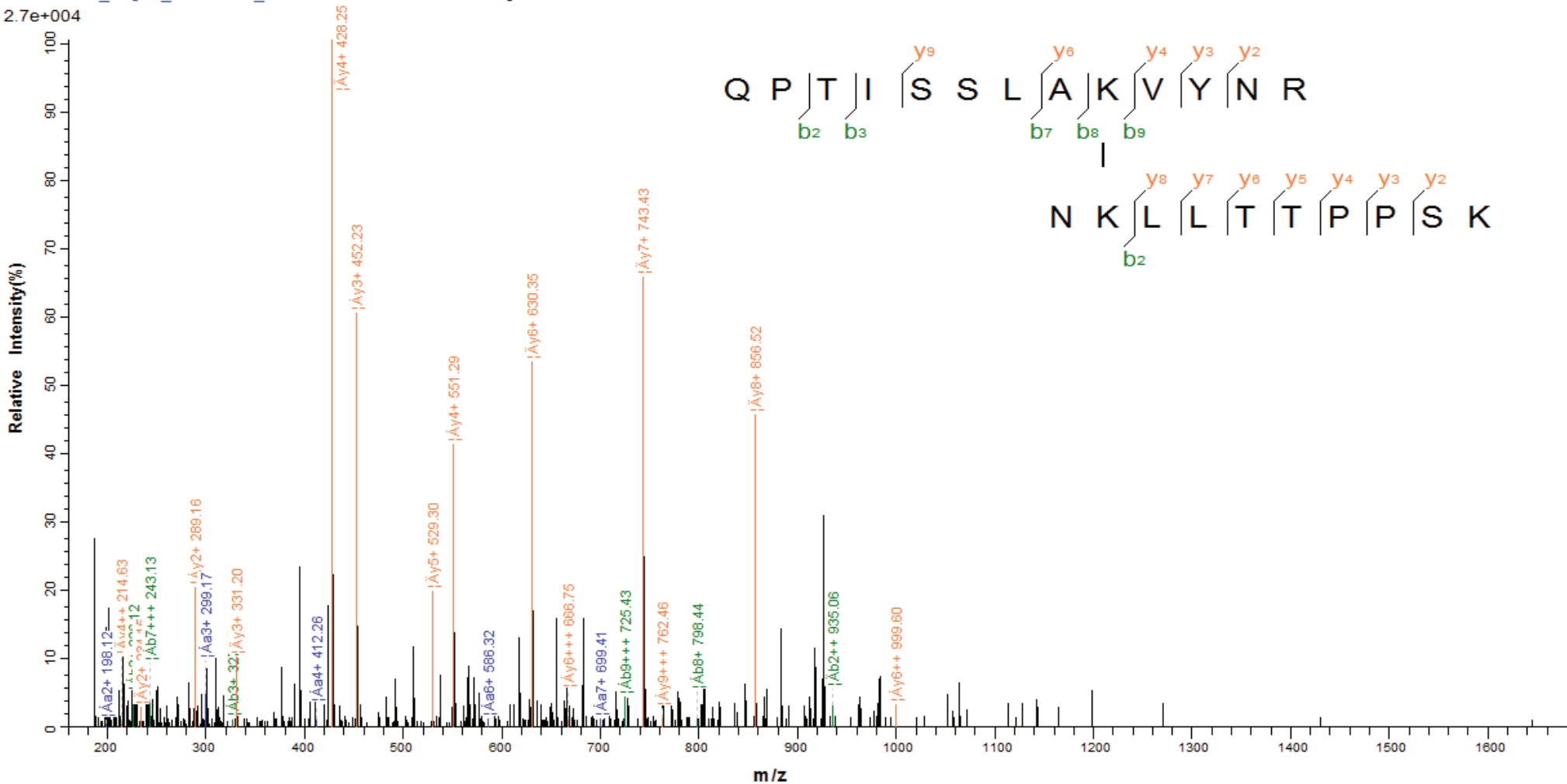


Seq: GNQVIIDELQKCHMQE-KKDAIK Mod: 12,C_57 14, Oxidation_M_15.995 MH+: 2798.4258



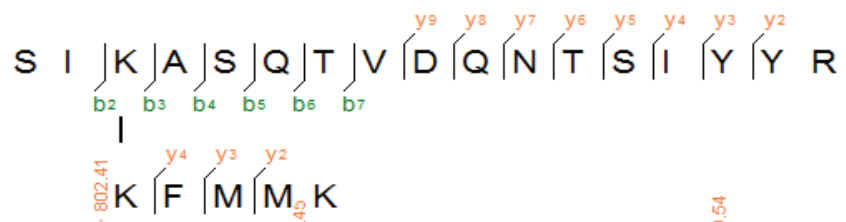
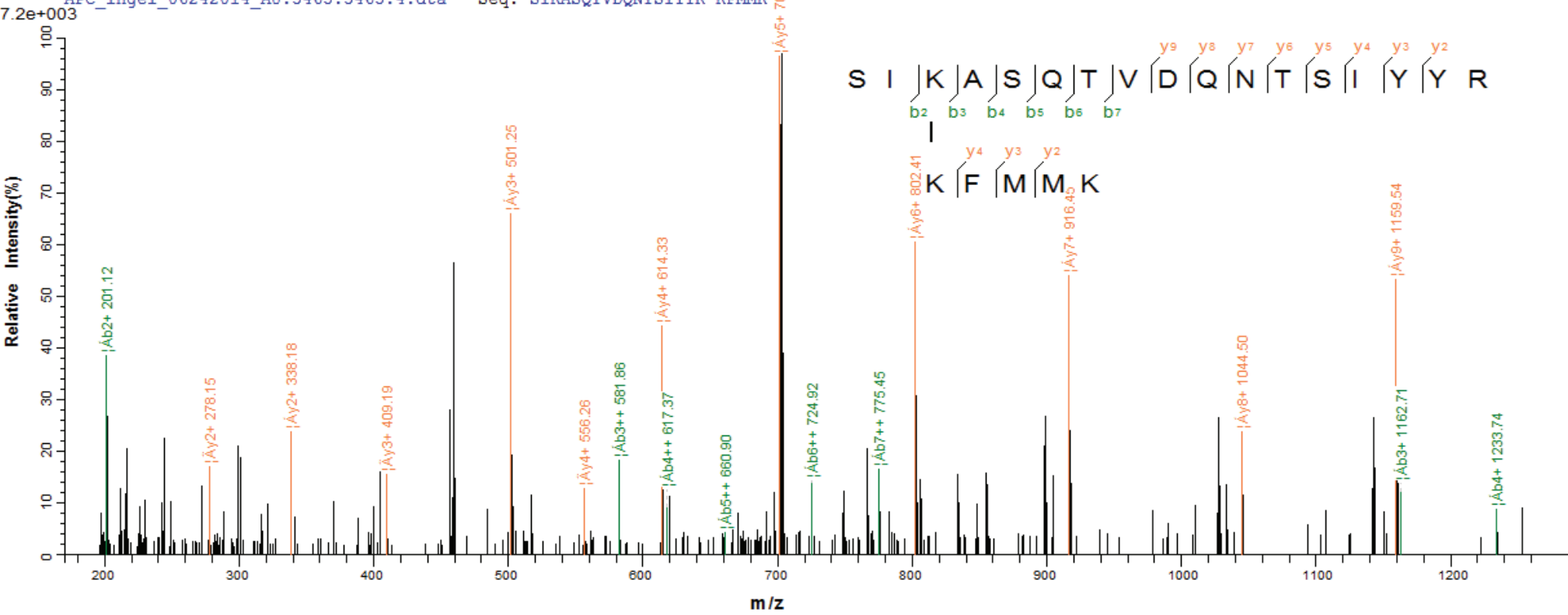
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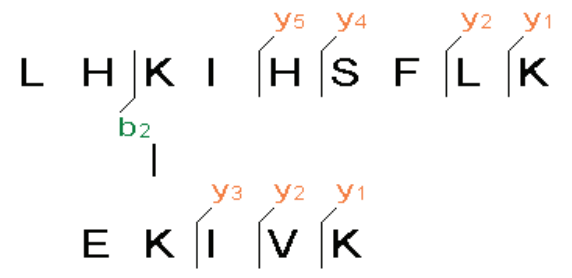
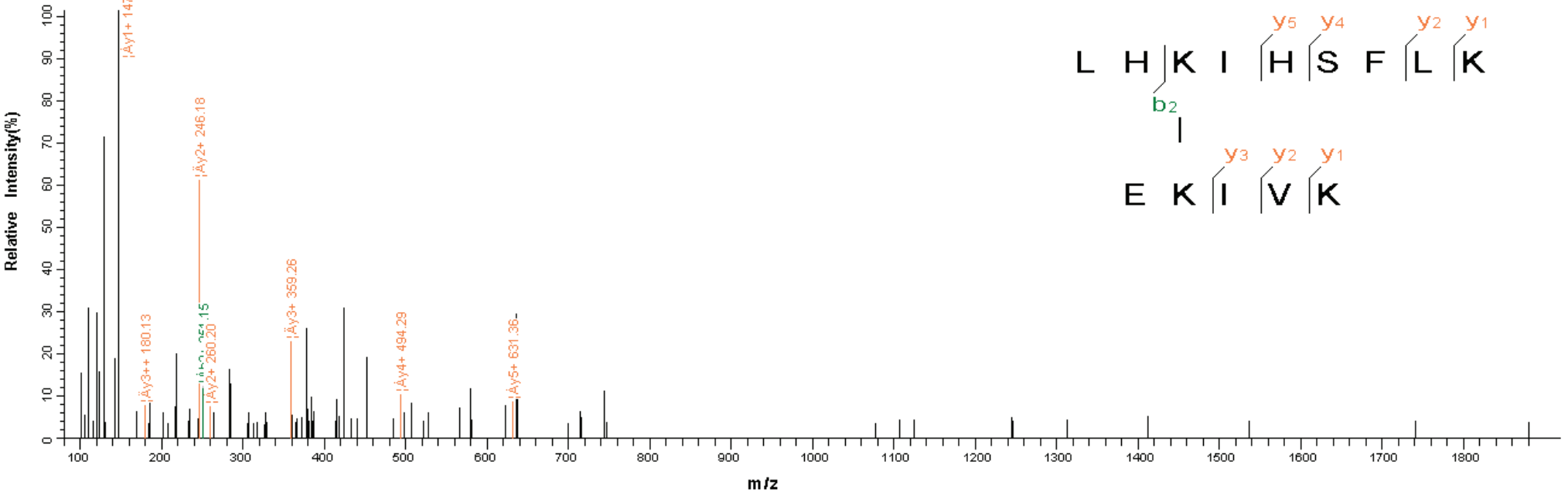


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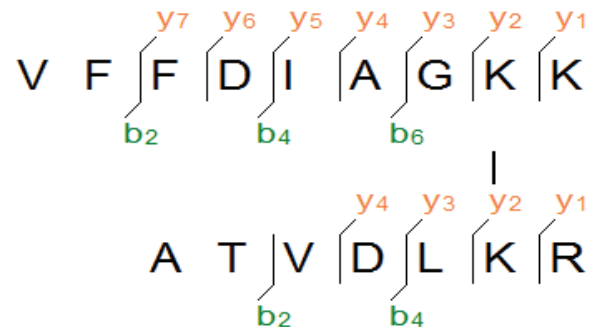
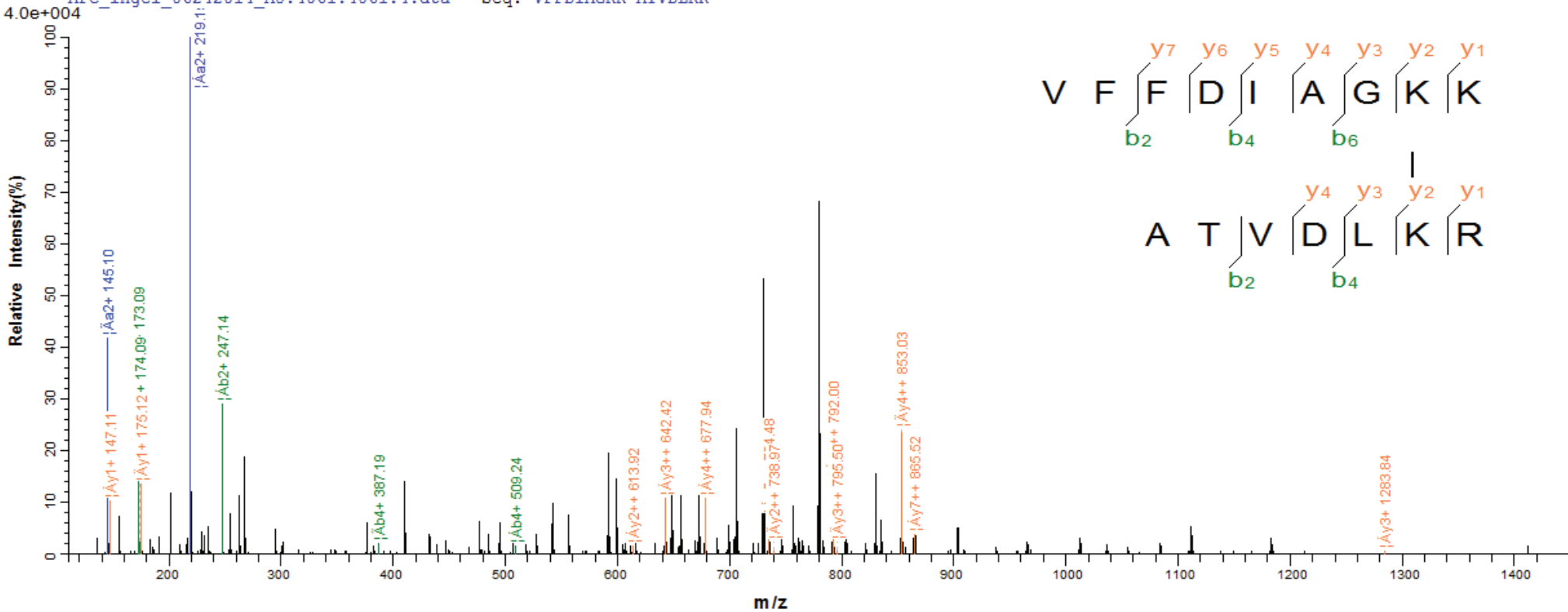
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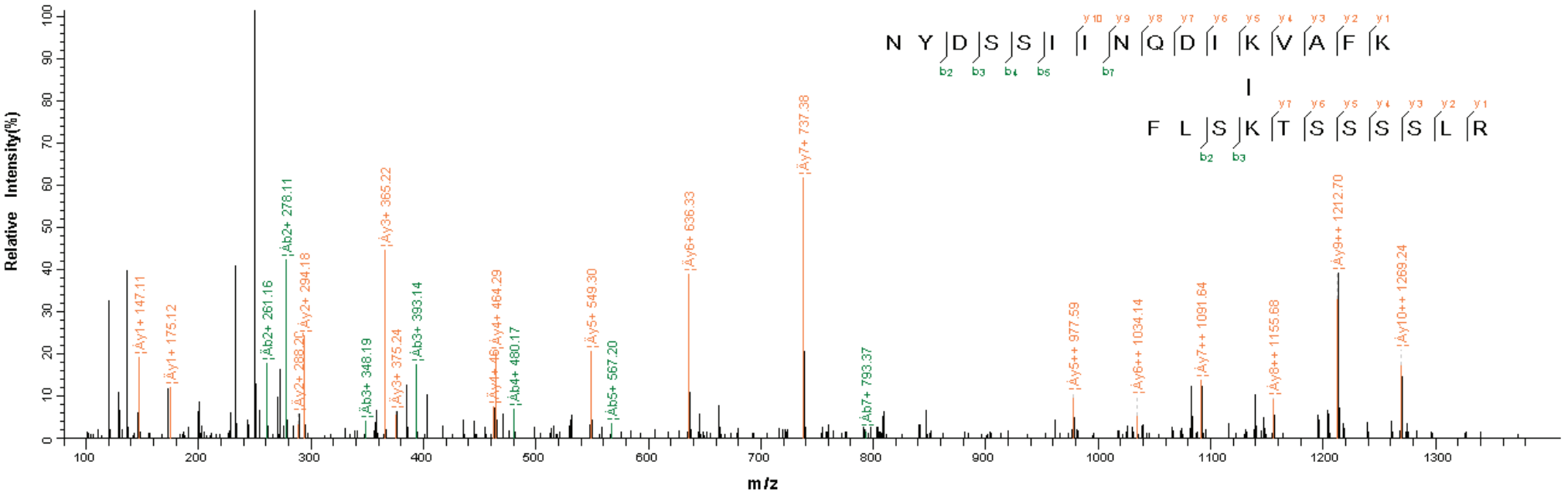
PC_A8 140201121226.002743.002743.5.dta Seq: LHKIHSFLK-EKIVK



APC_ingel_06242014_A8.4061.4061.4.dta Seq: VFFDIAGKK-ATVDLKR



1.0e+004 PC_A8_140201121226.006467.006467.4.dta Seq: NYDSSIINQDIKVAFK-FLSKTSSSSLR



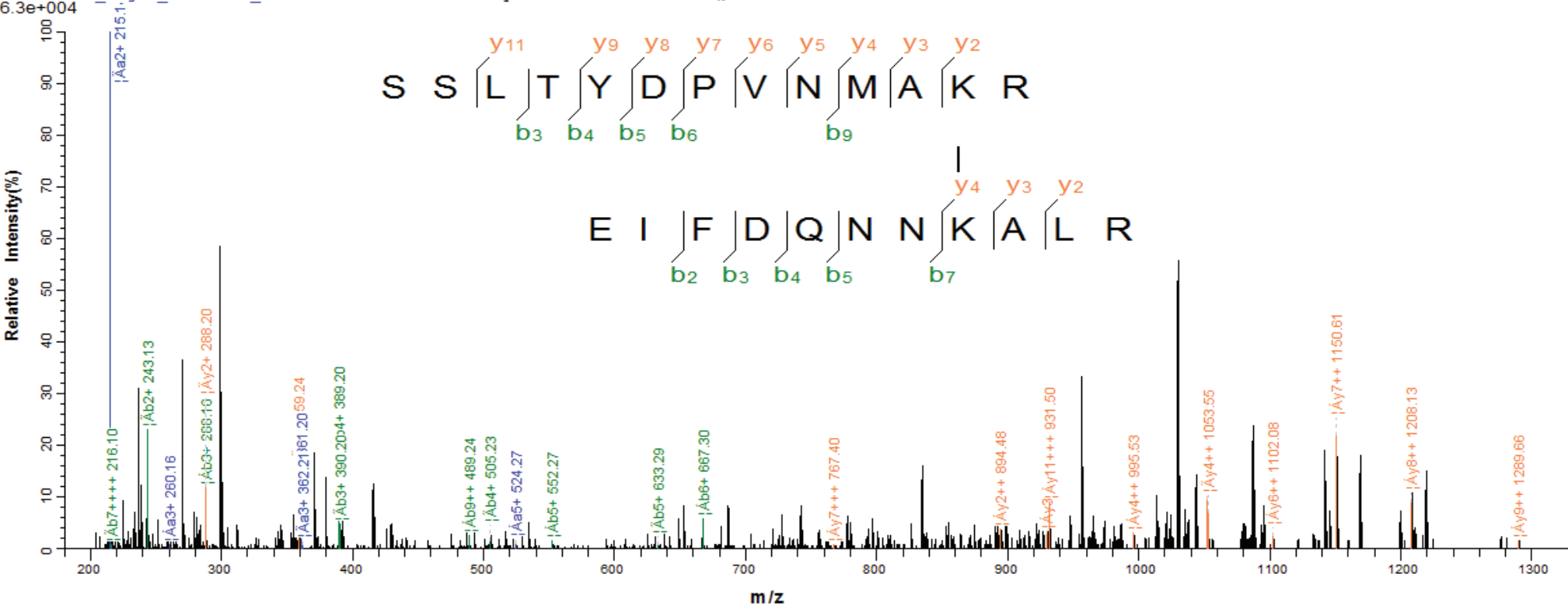
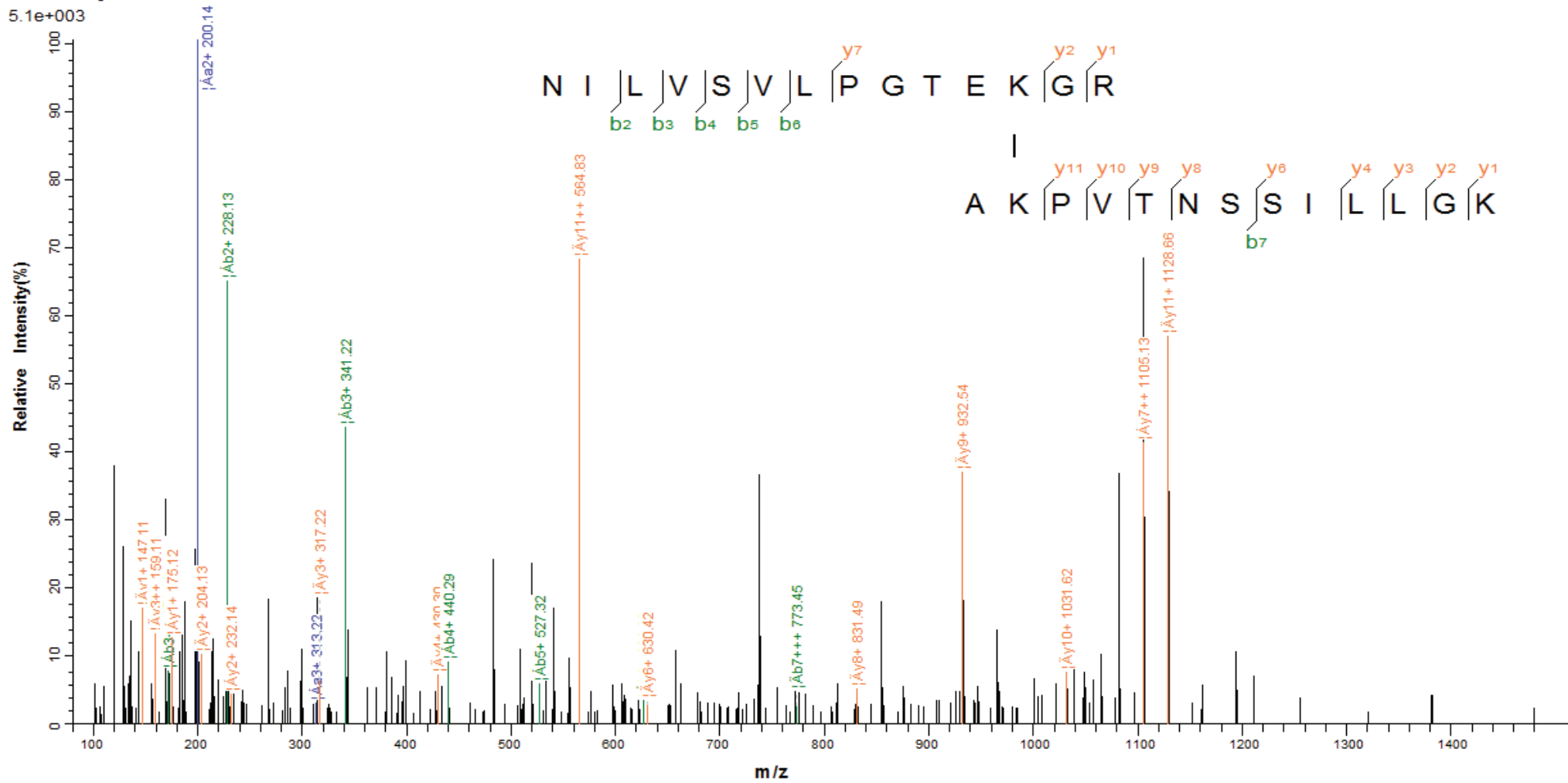


Fig. S17

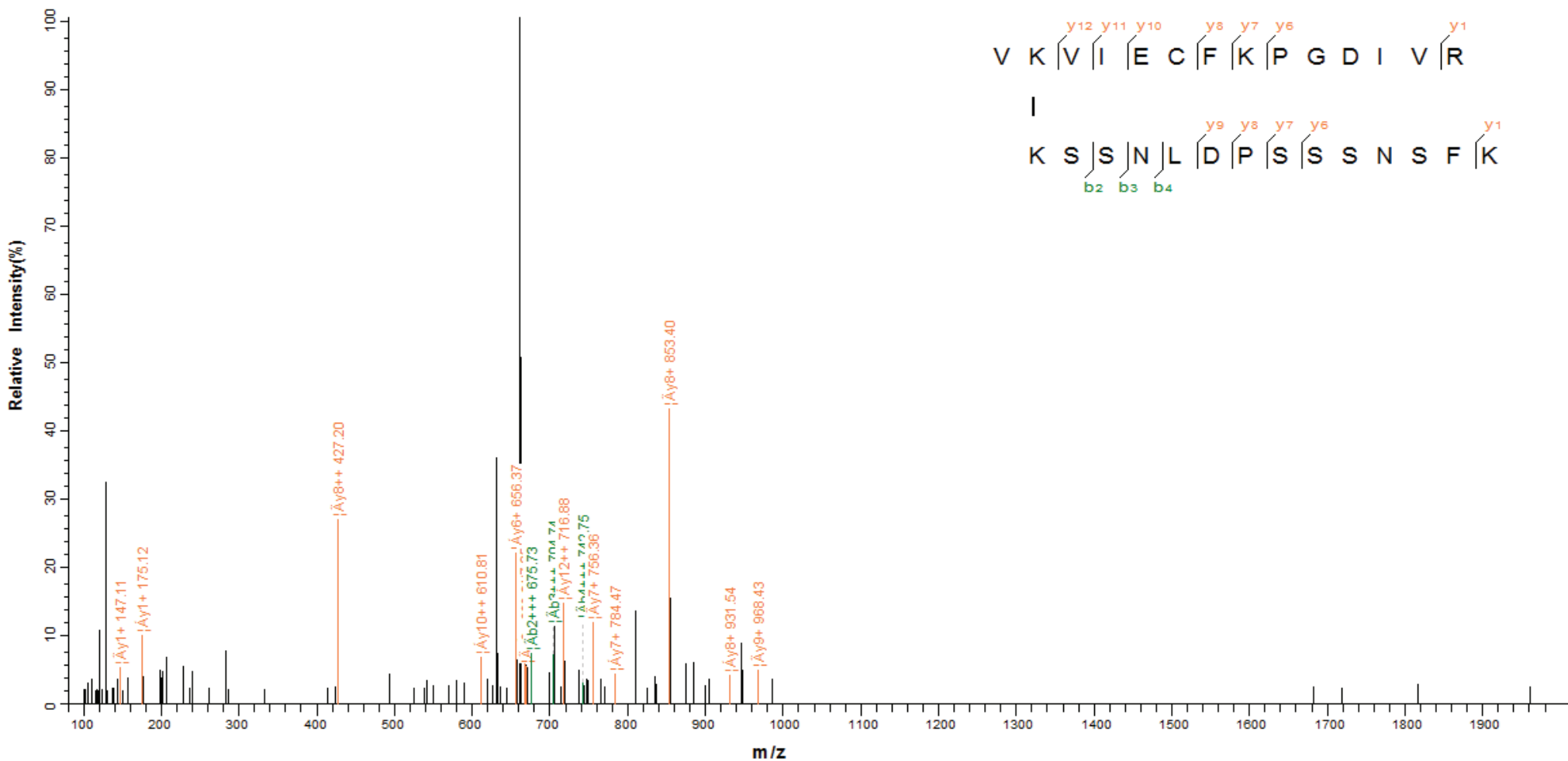
Annotated high-resolution HCD spectra of
crosslink peptides of exosome complex

Seq: NILVSVLPGTEKGR-AKPVTNSSILLGK



Seq: VKVIECFKPGDIVR-KSSNLDPSSSNSFK Mod: 6,C_57

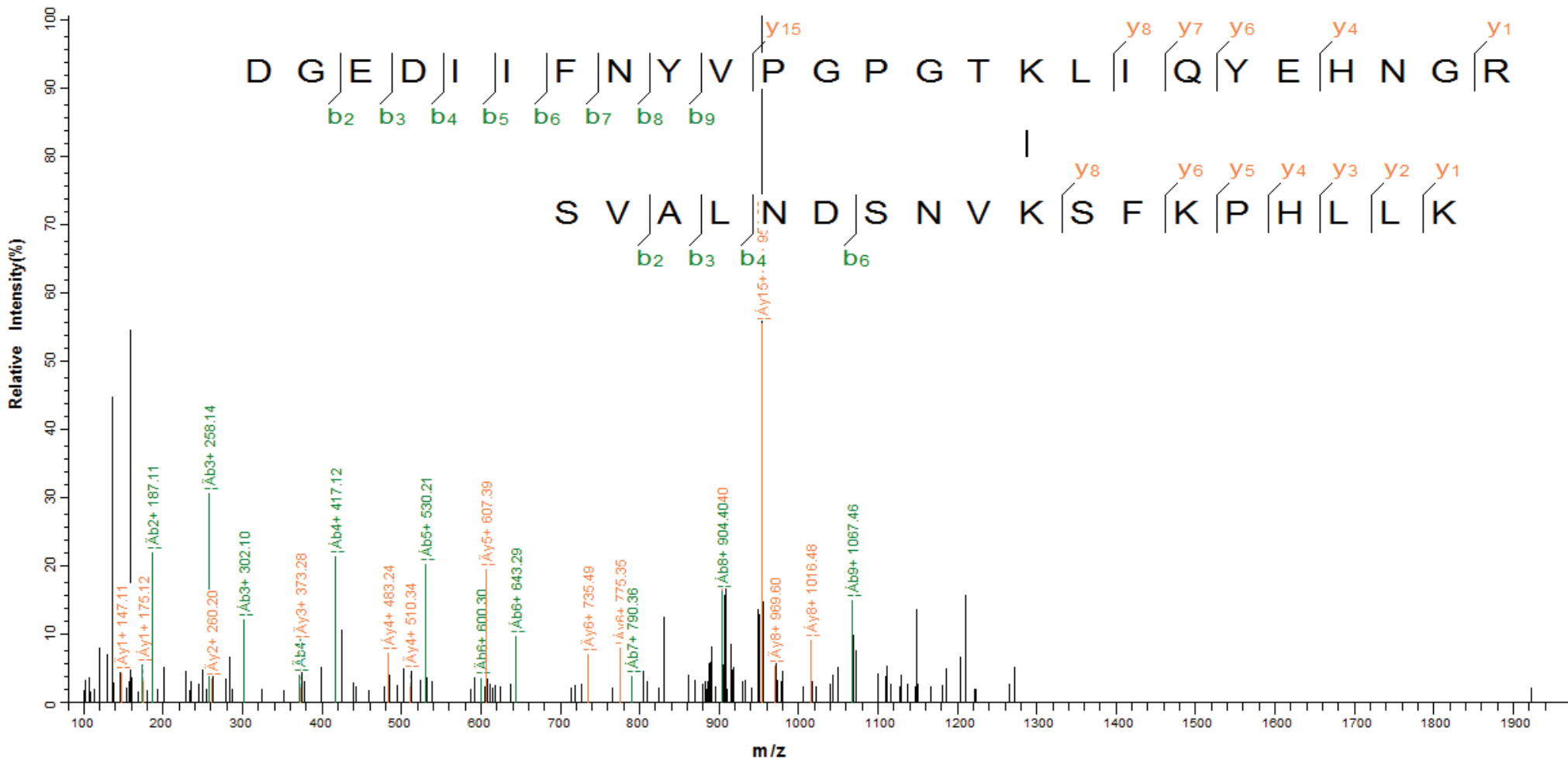
2.2e+003



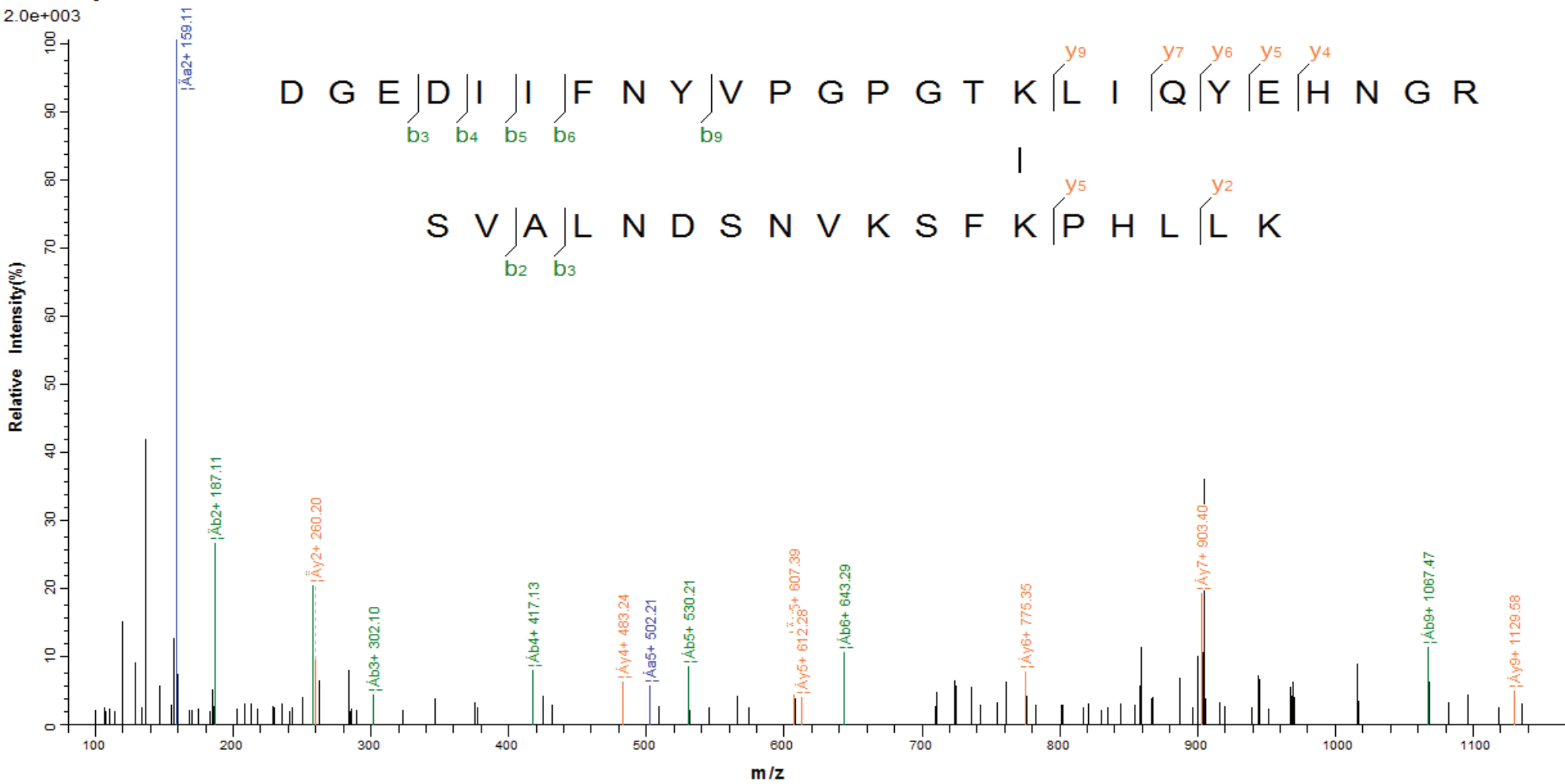
V K V I E C F K P G D I V R
I
K S S N L D P S S S N S F K
b2 b3 b4

Seq: DGEDIIFNYVPGPGTKLIQYEHNGR-SVALNDSNVKSFKPHLLK Score: 2.24e-012

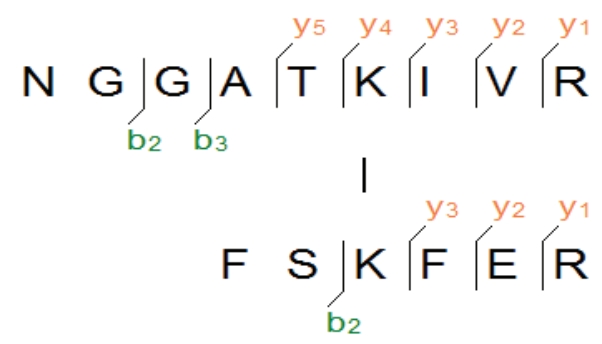
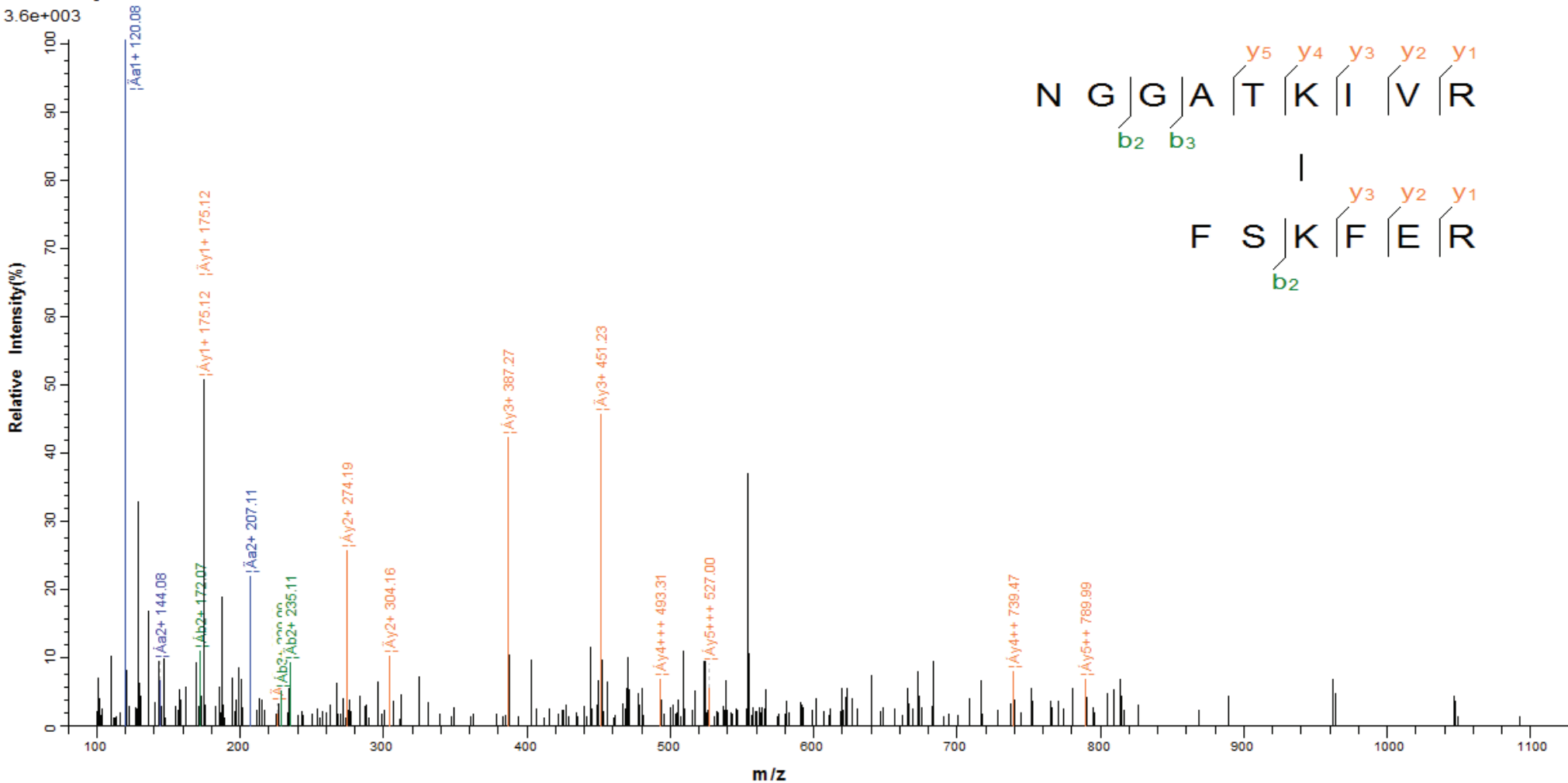
2.4e+003



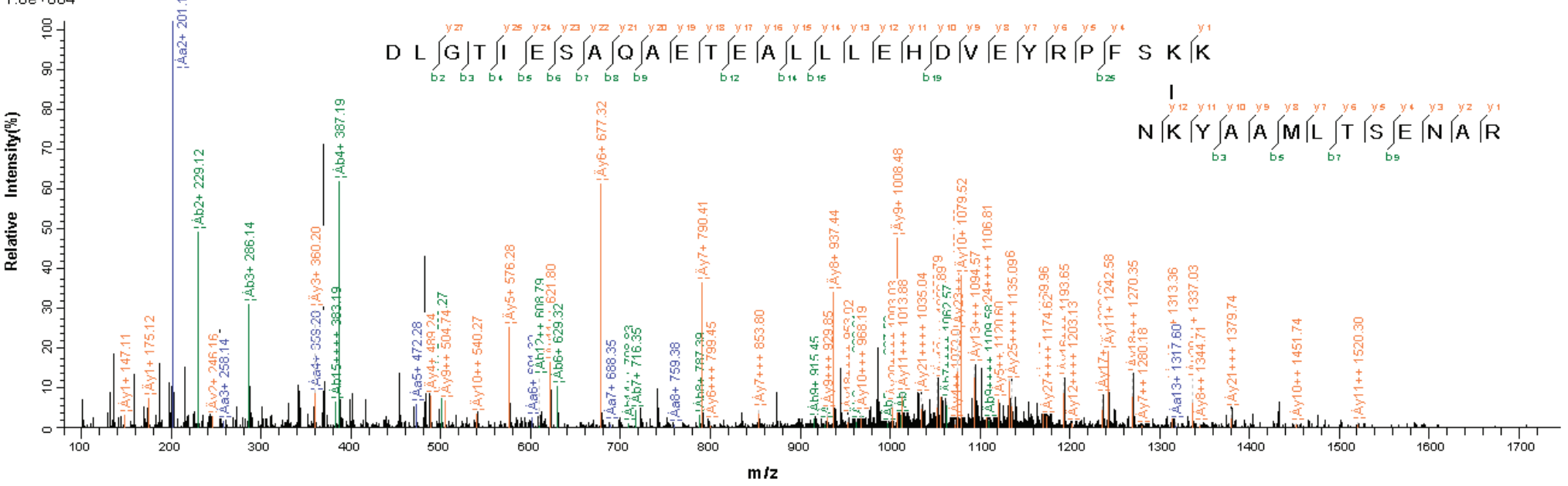
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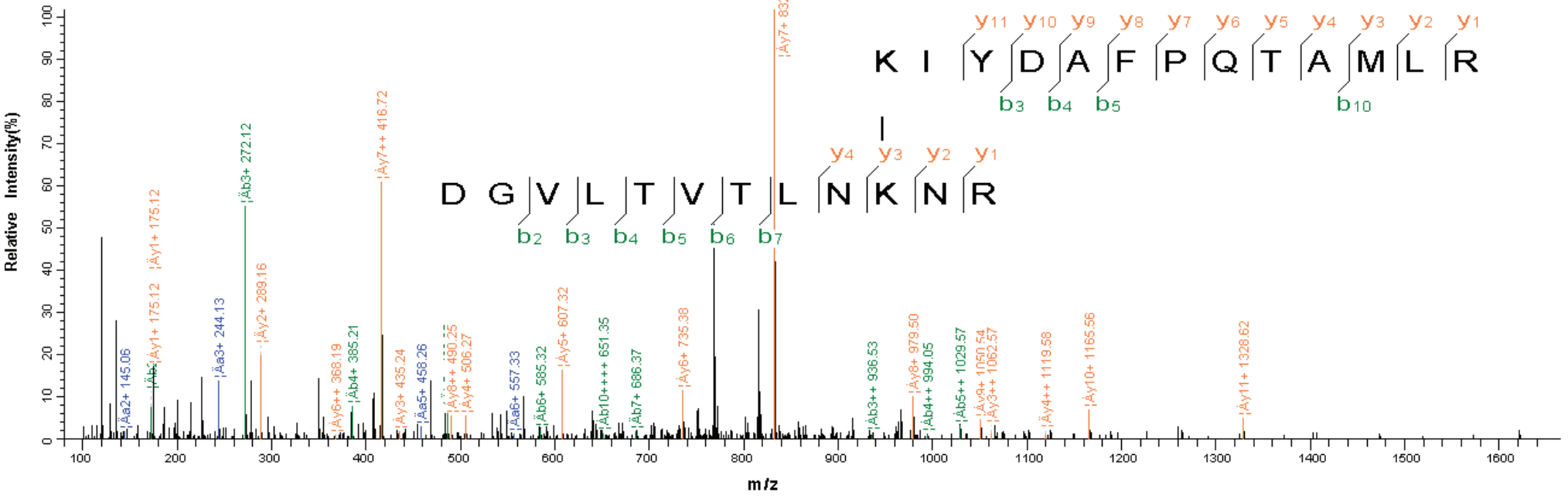
Seq: NGGATKIVR-FSKFER



Exosome_GFP_test_A78.003476.003476.5.dta Seq: DLGTTESAQAETEALHLDVYRPFSSKK-NKYAAMLTSENAF Mod: 38,_Oxidation_M_15.995 Score: 6.88e-007

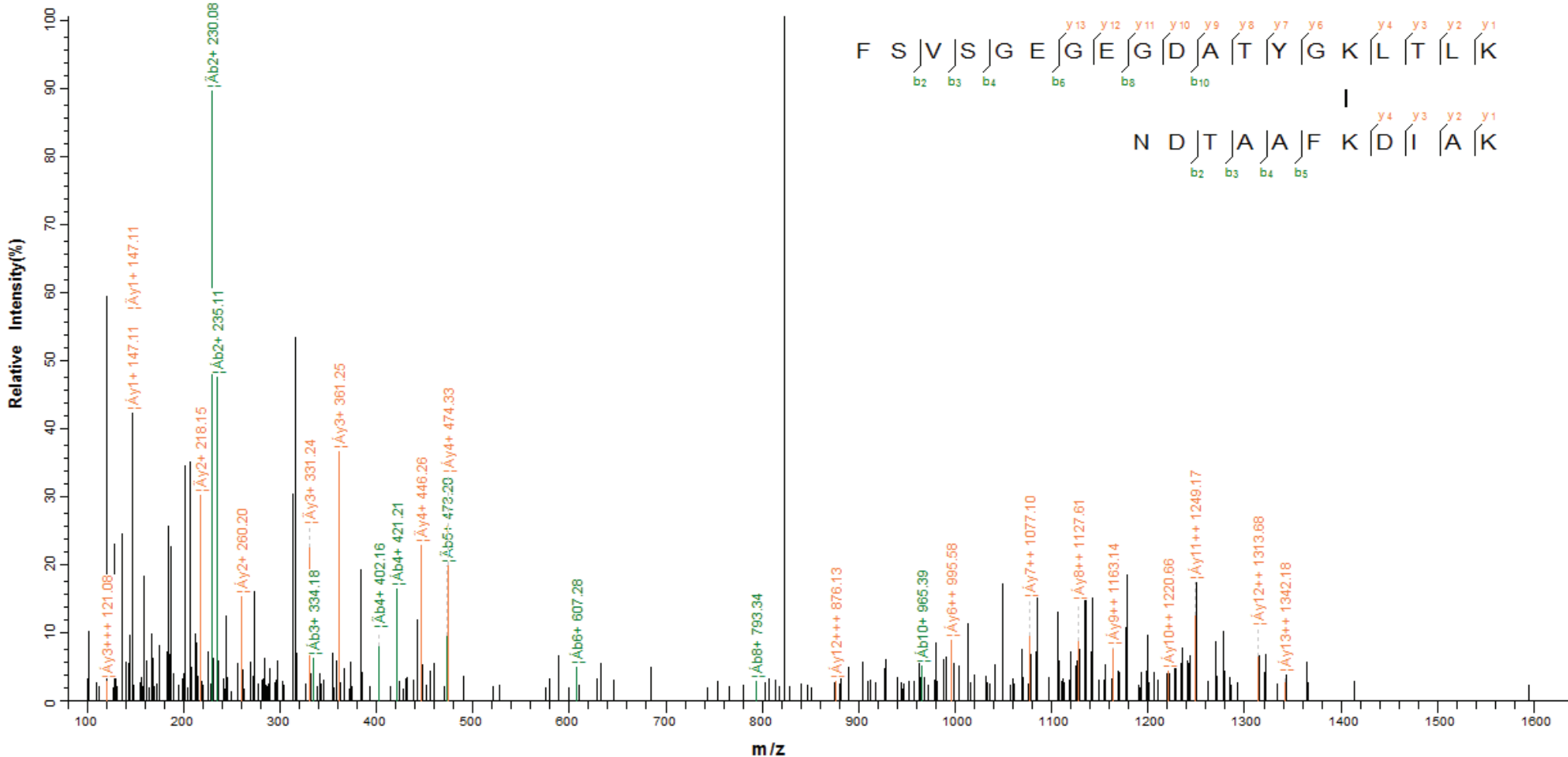


Seq: KIYDAFPQTAMLR-DGVLTVTLNKNR Mod: 11,_Oxidation_M_15.995

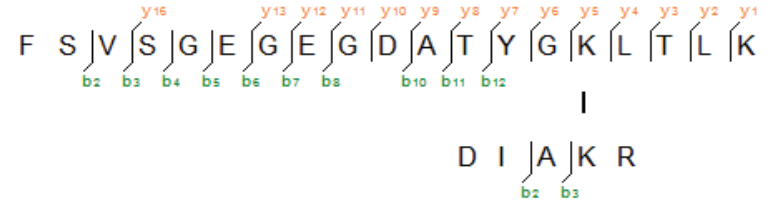
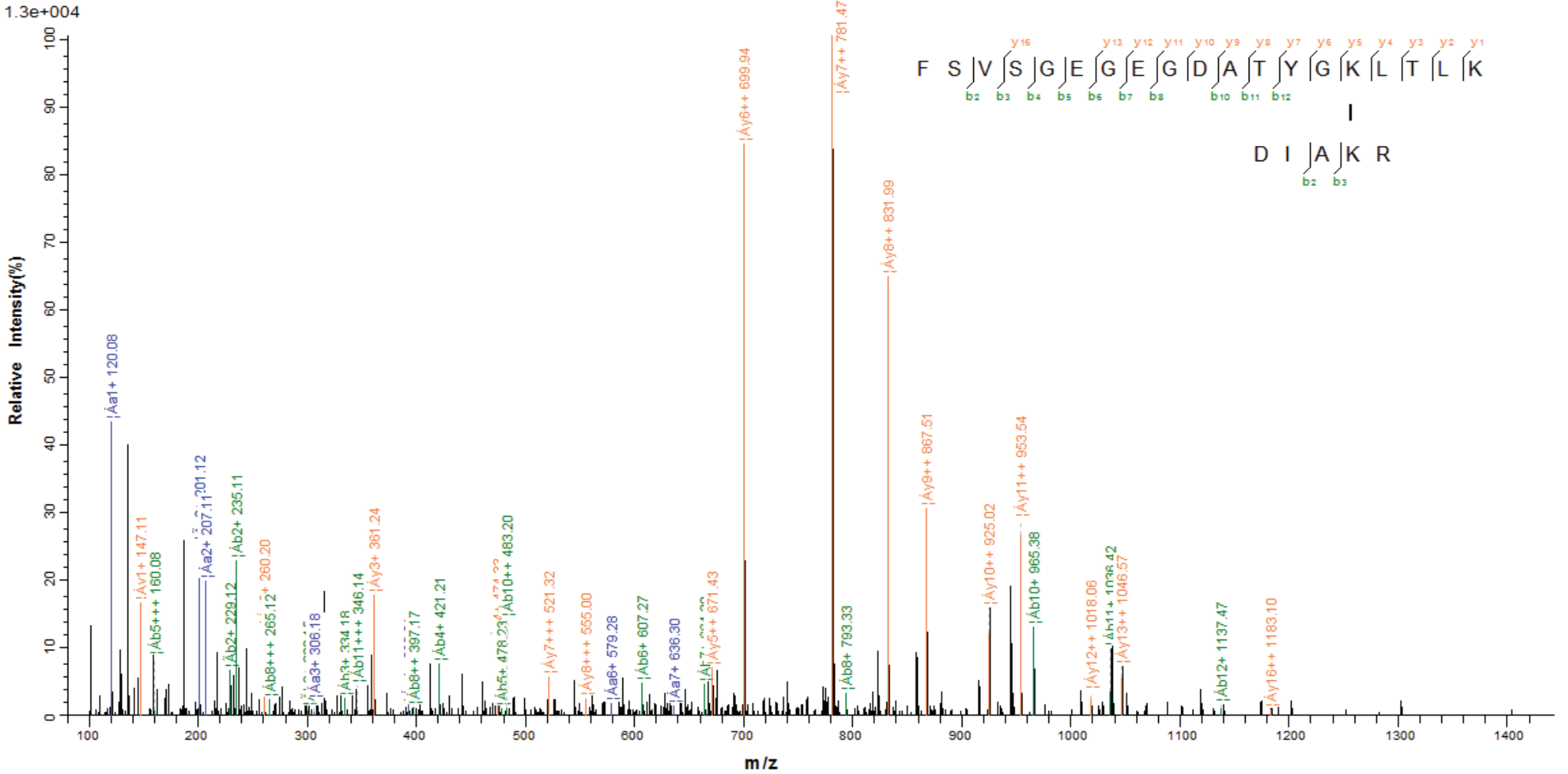


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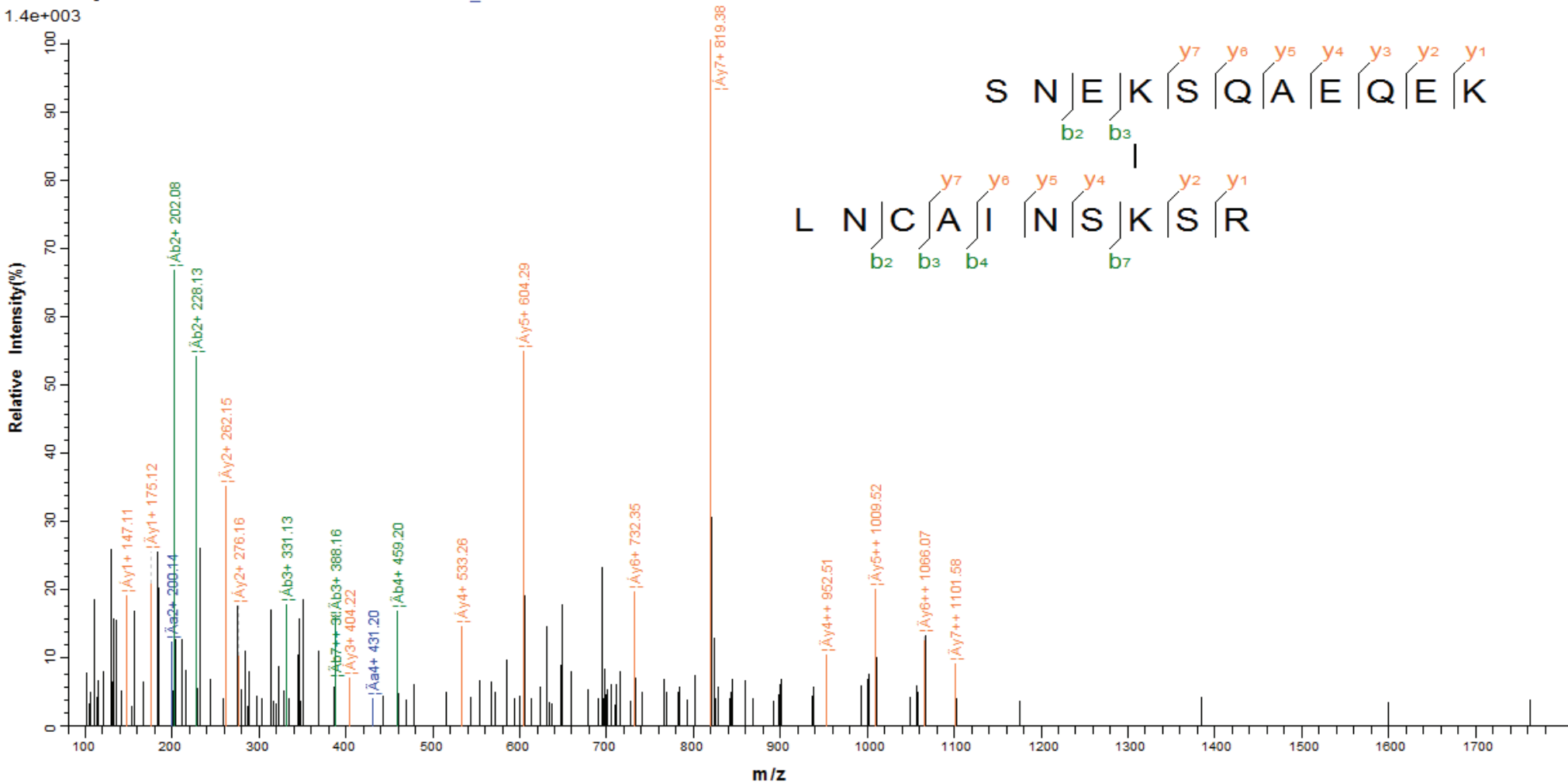
2.8e+003



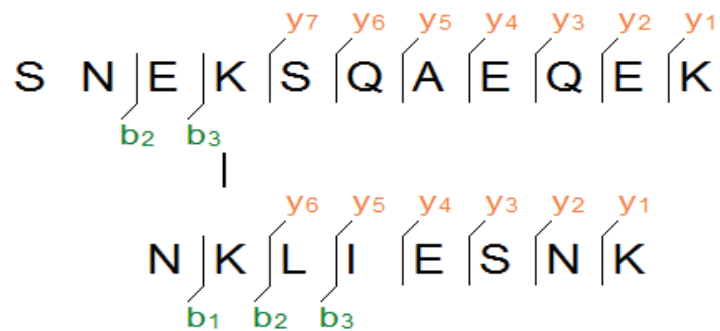
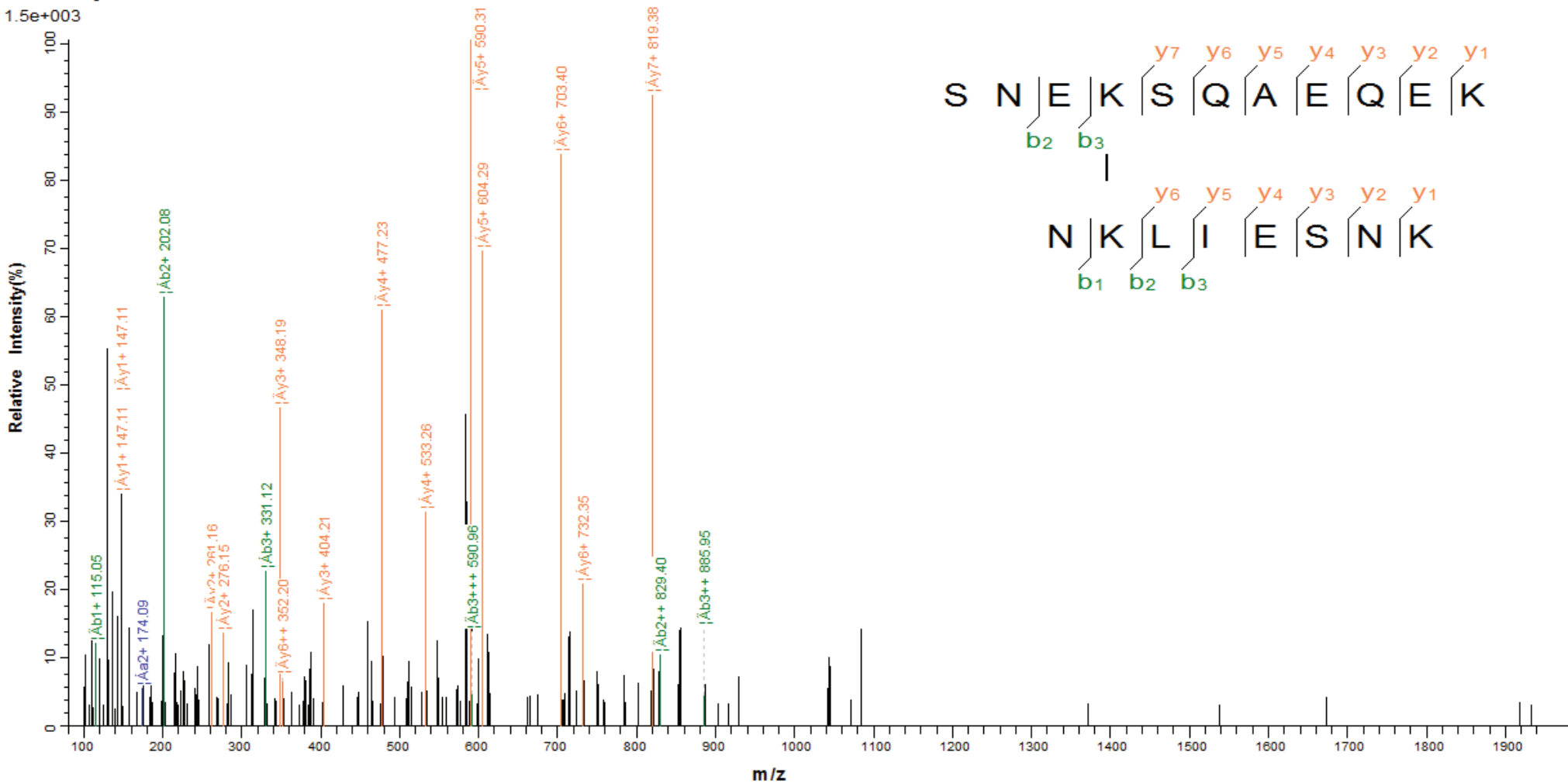
Seq: FSVS⁺GEGGDATY⁺GKLT⁺LK-DIAKR



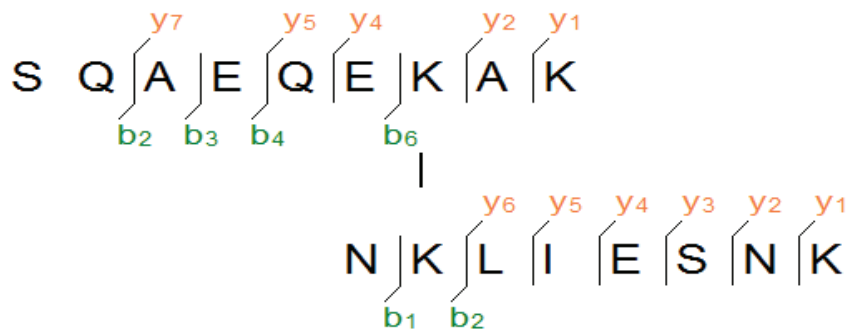
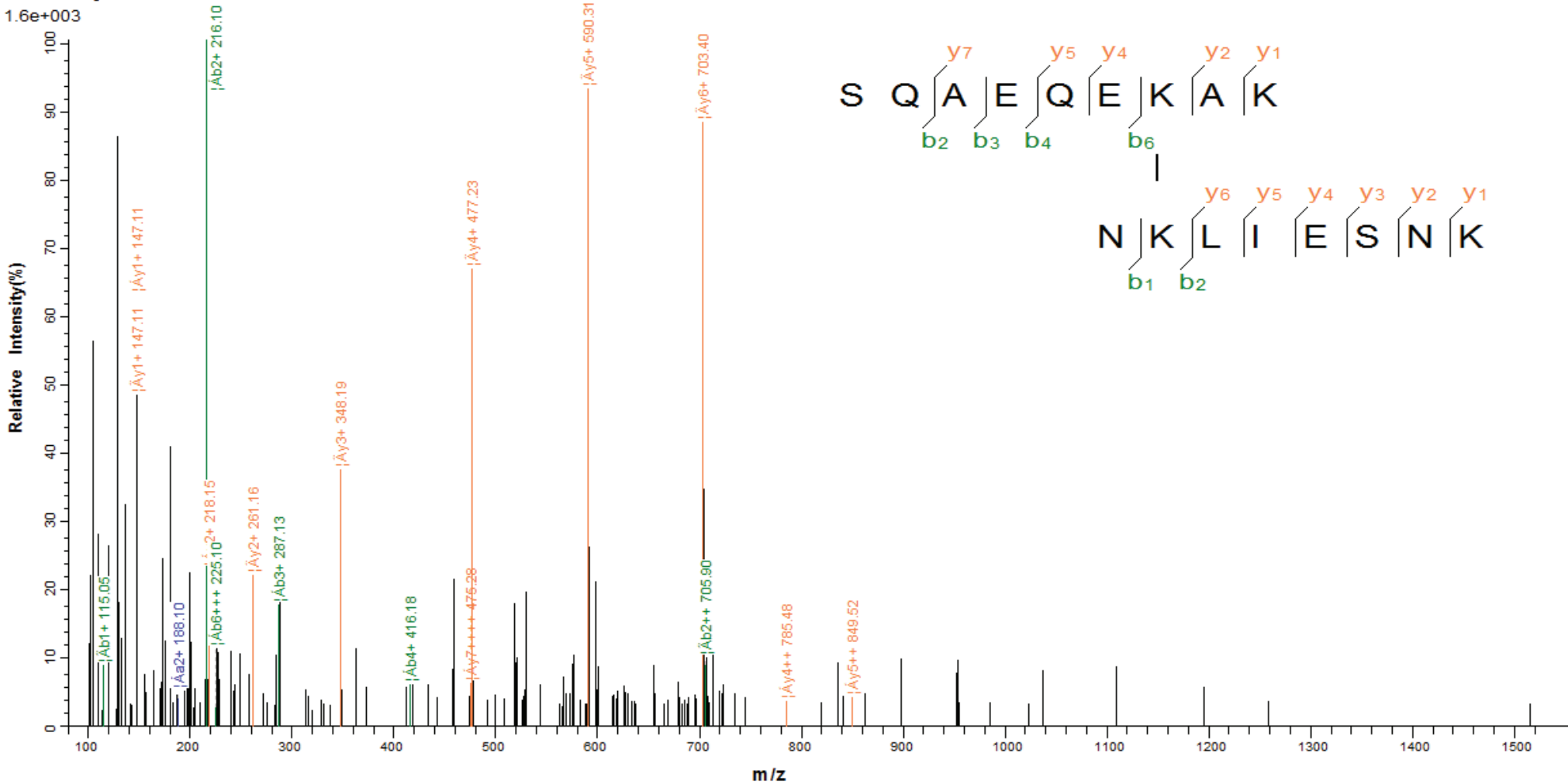
Seq: SNEKSQAEQEK-LNCAINSKR Mod: 17,C_57



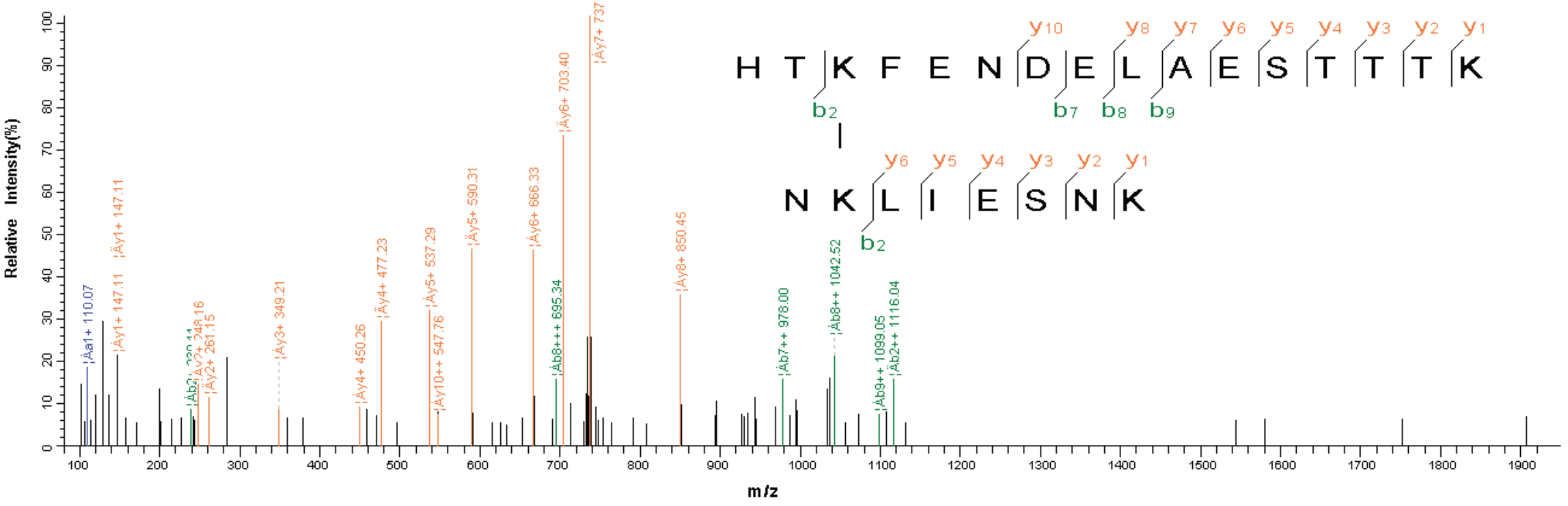
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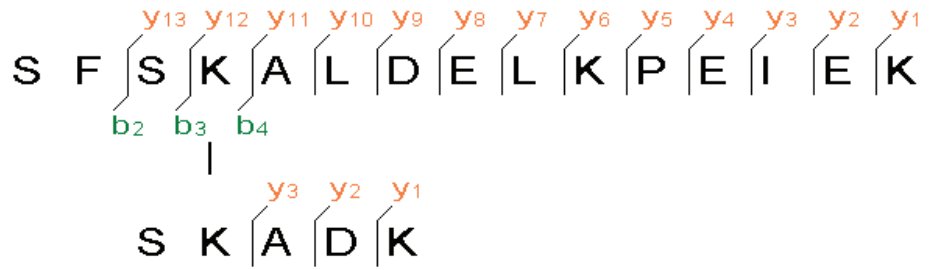
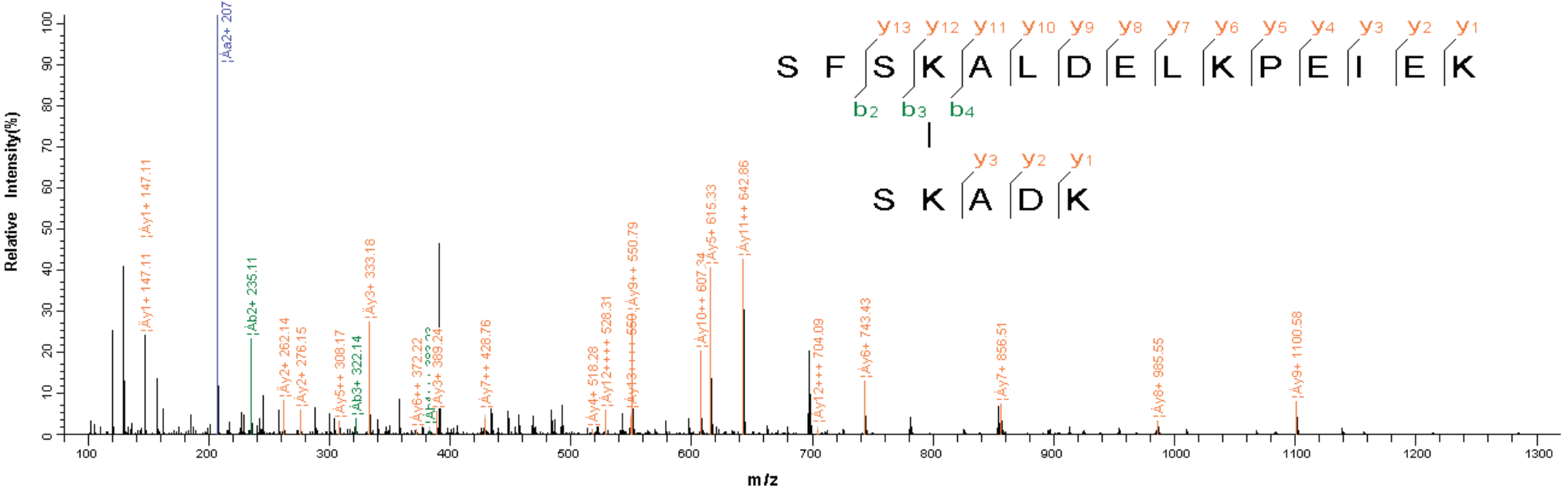
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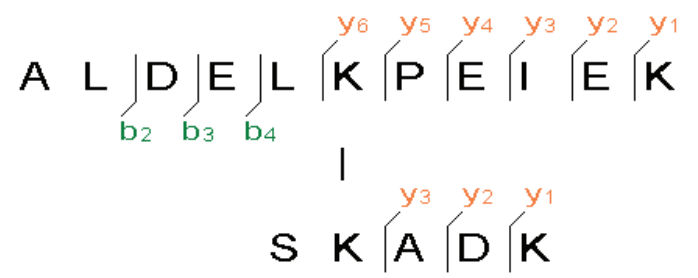
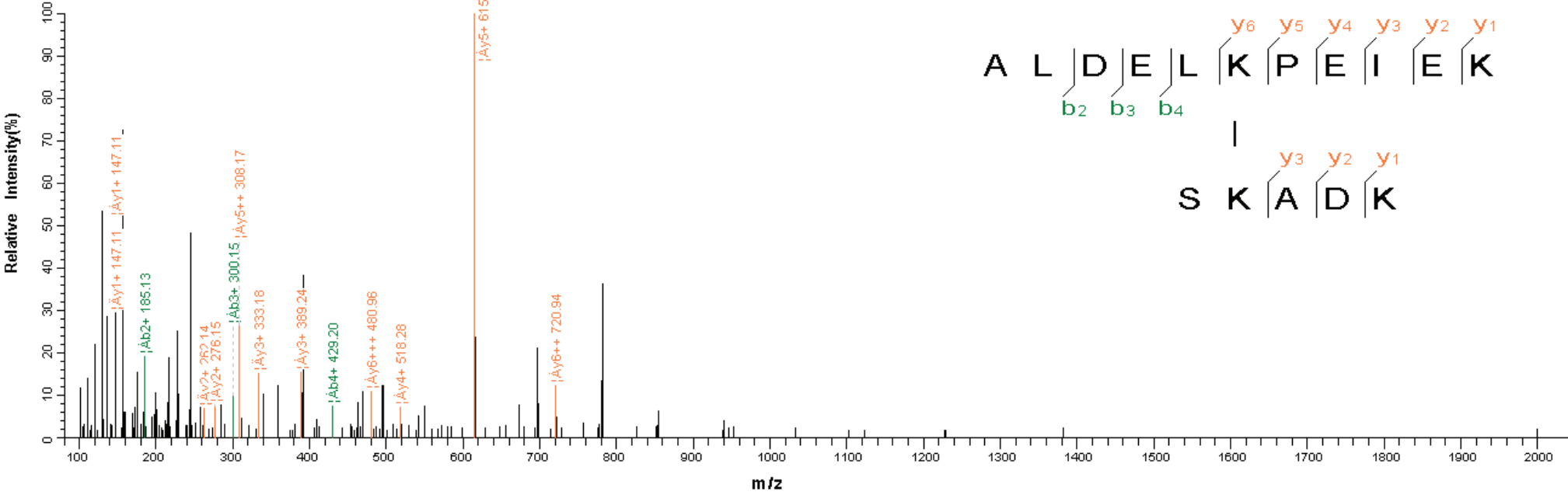
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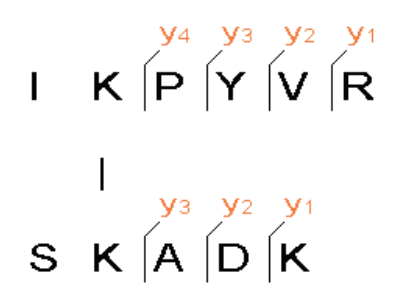
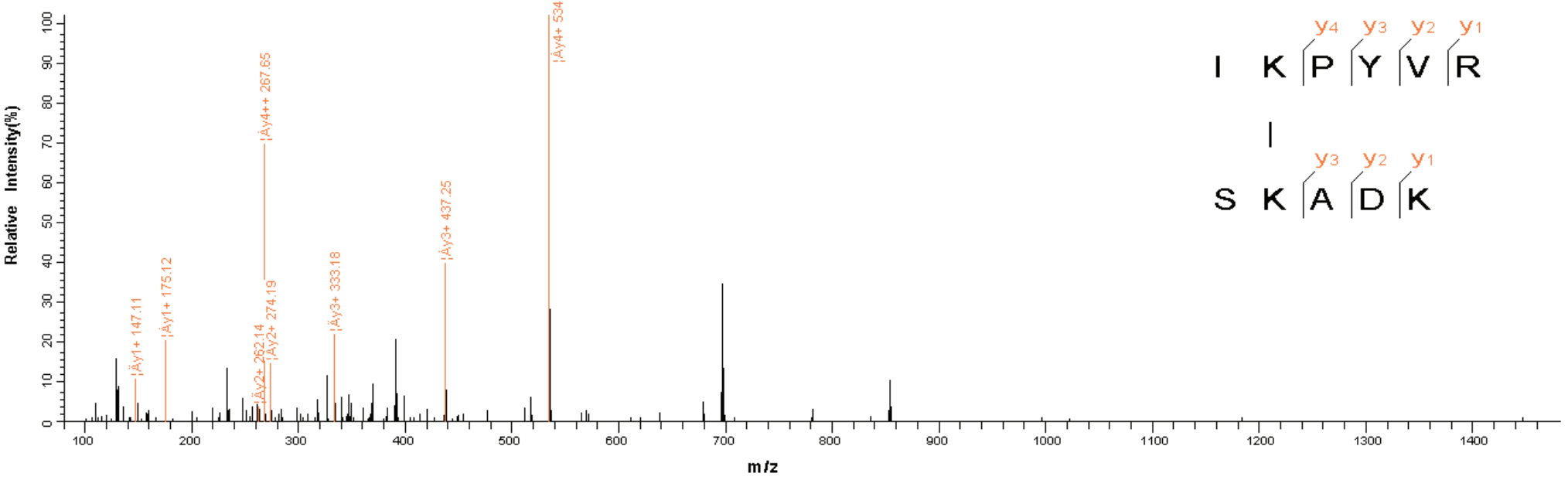
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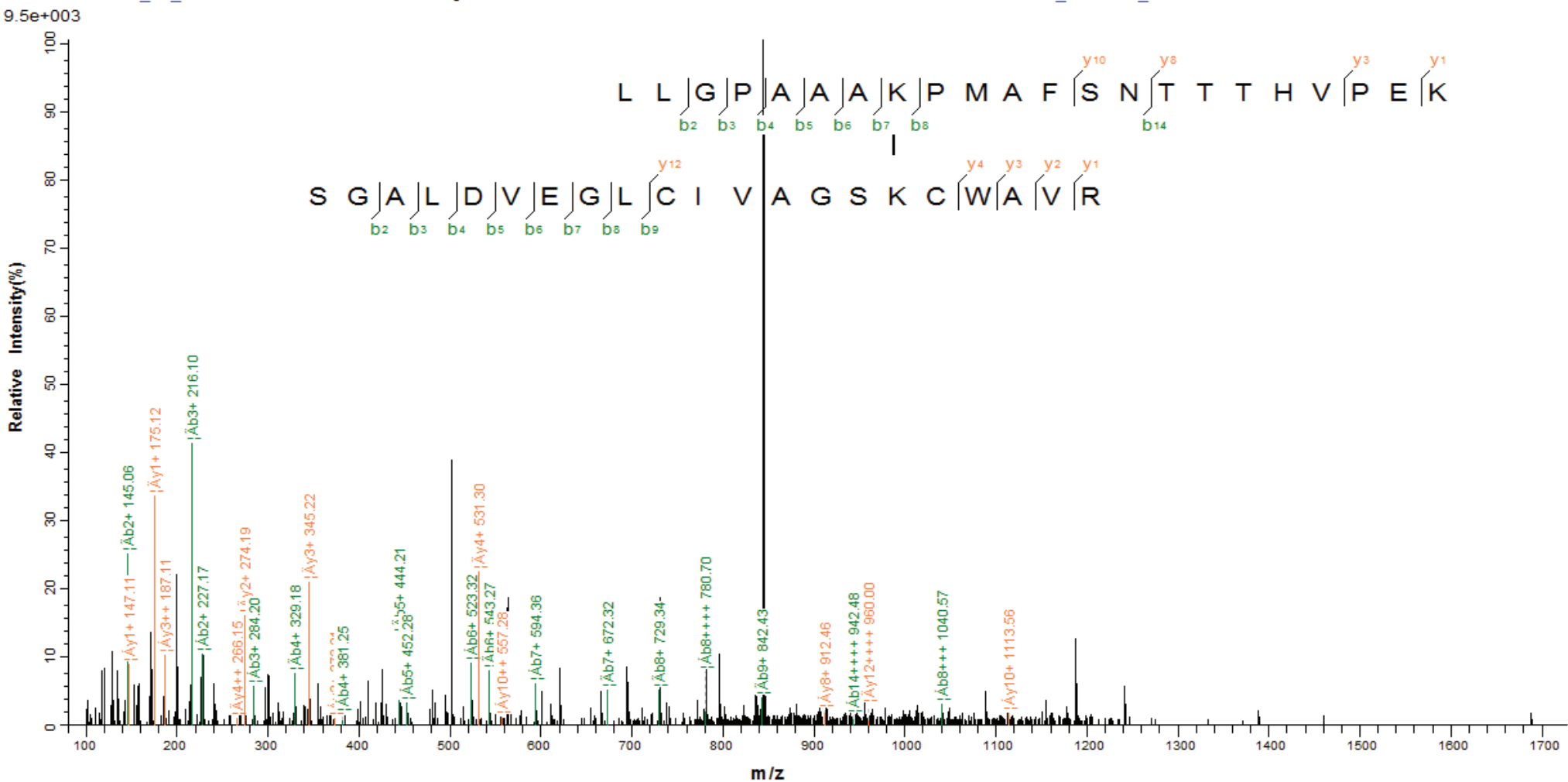


seq: ALDELKPEIEK-SKADK



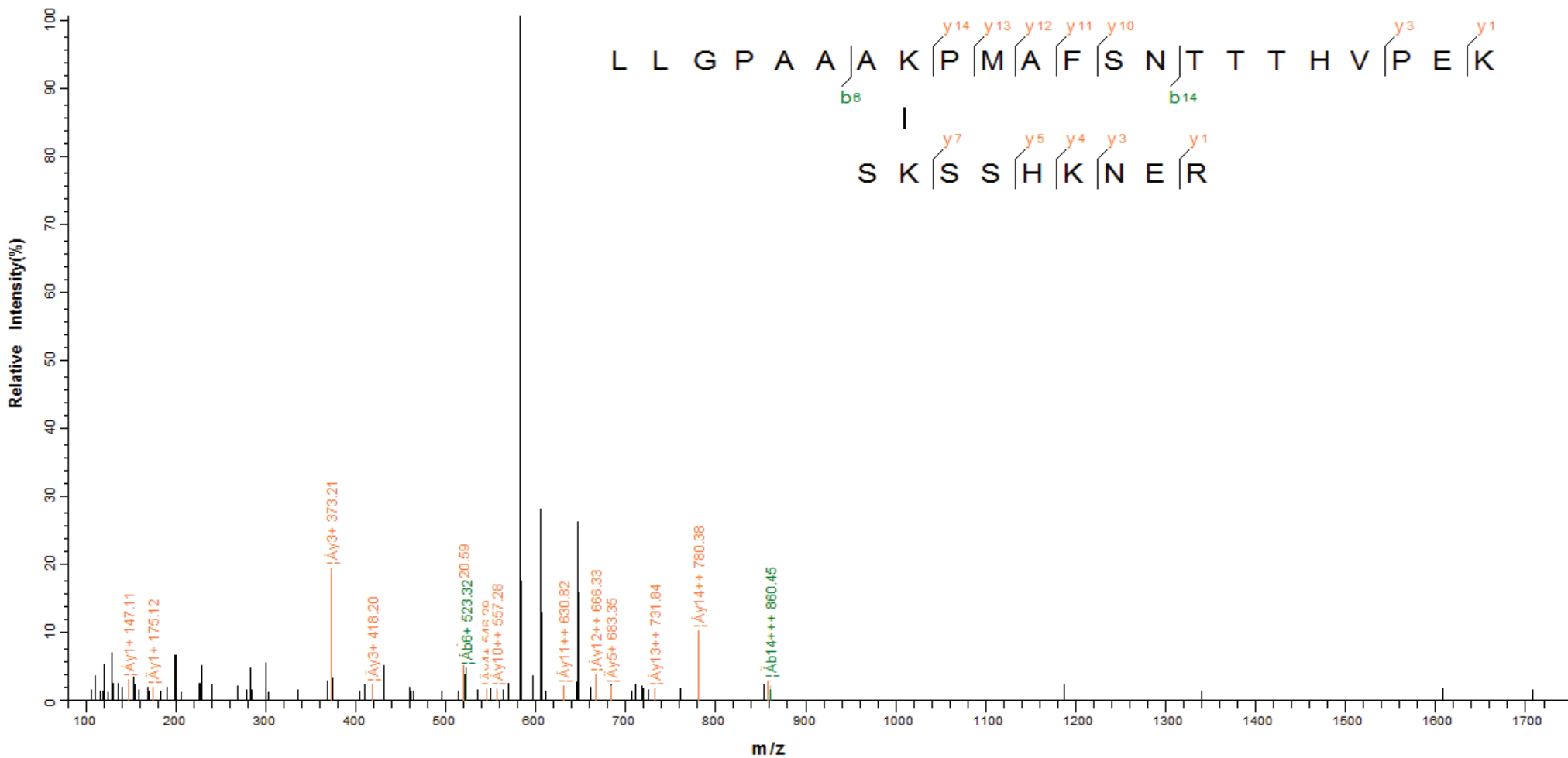
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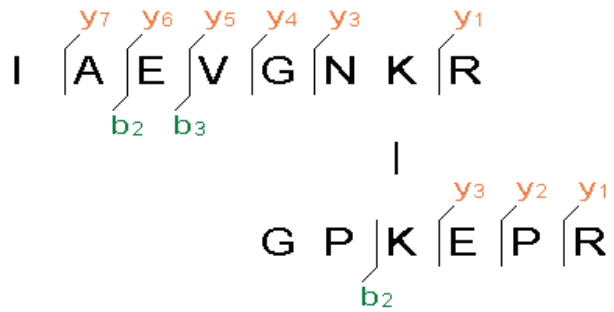
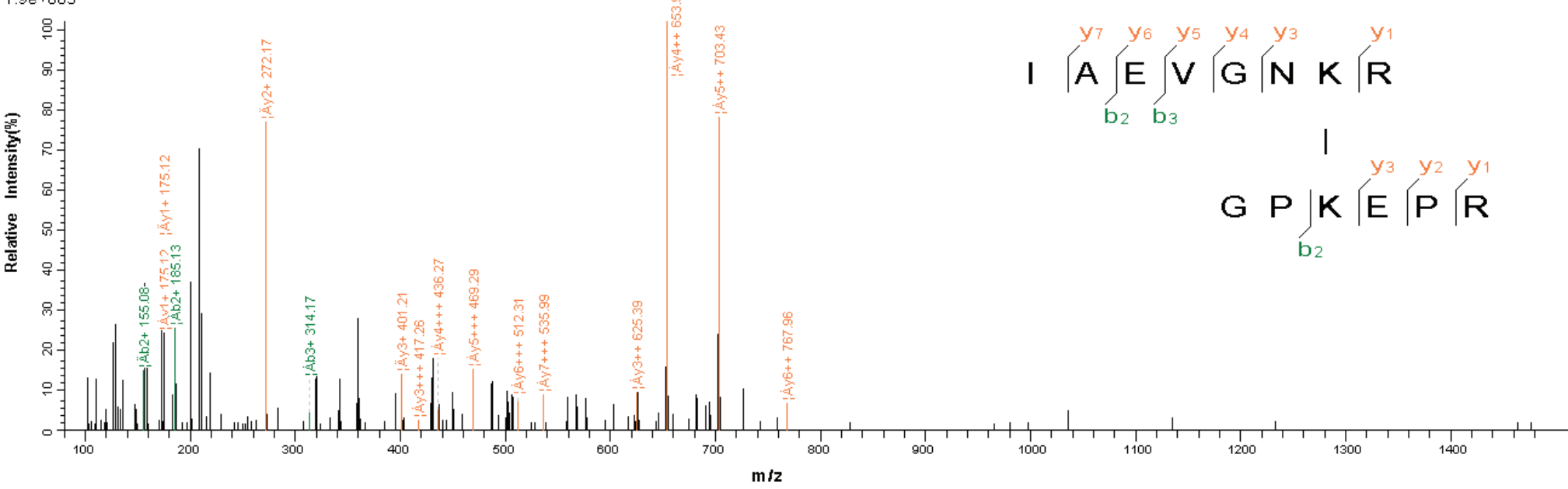


exosome_A8_1.003784.003784.6.dta Seq: LLGPAAAKPMAFSNTTTHVPEK-SKSSHKNER

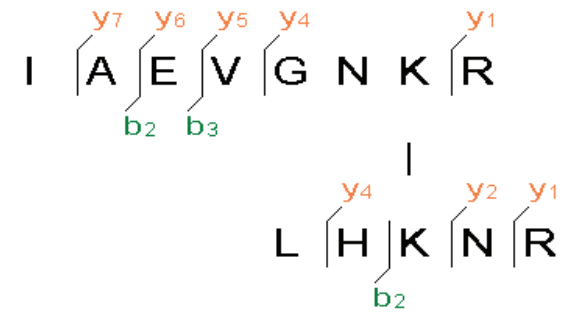
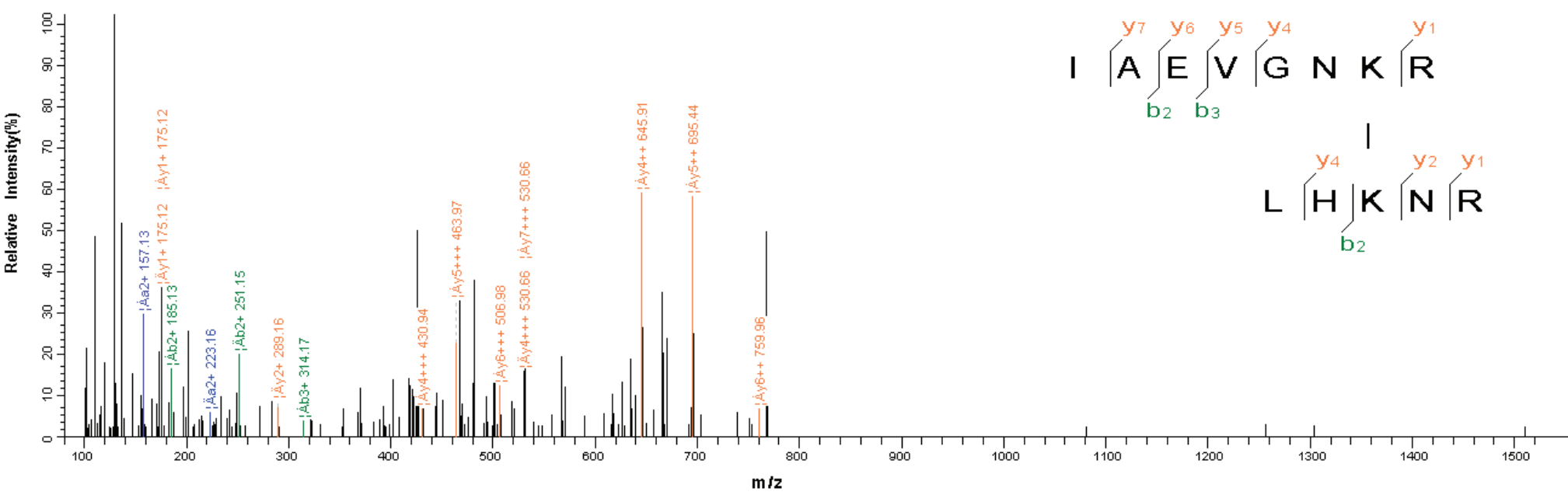
3.2e+003



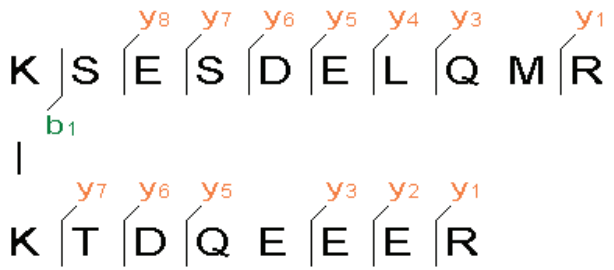
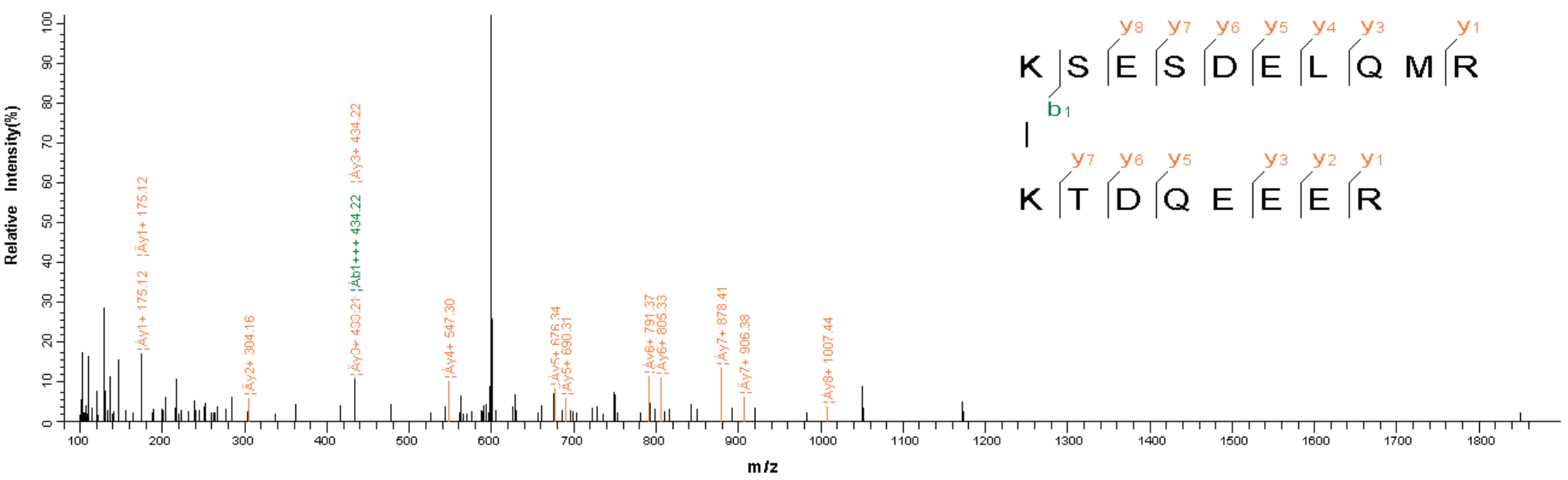
exosome_A9_140503145423.1521.1521.4.dta Seq: IAEVGNKR-GPKEPR



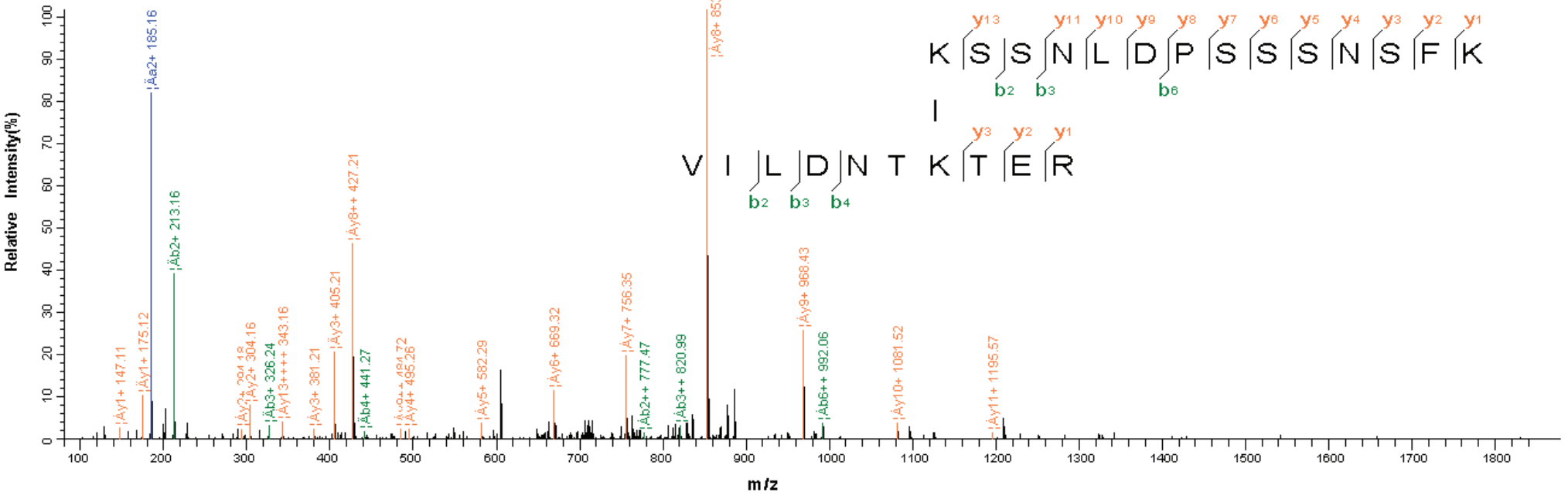
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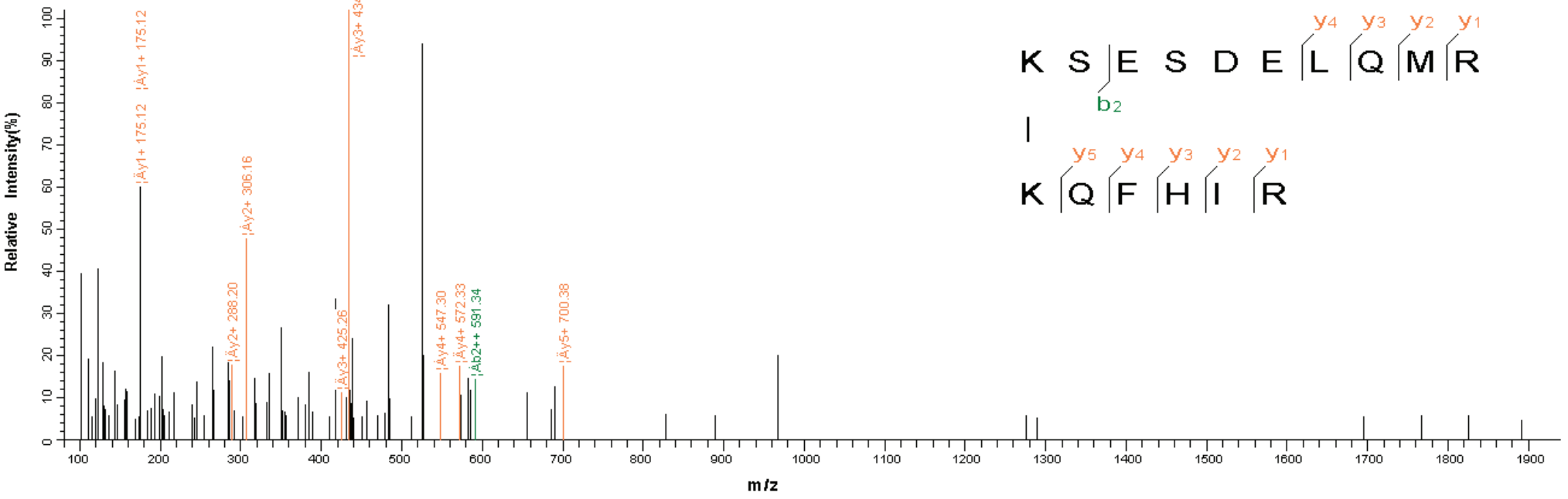
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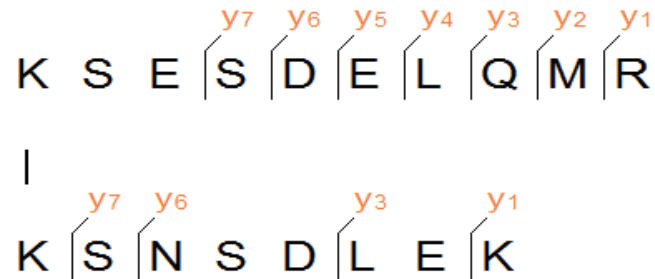
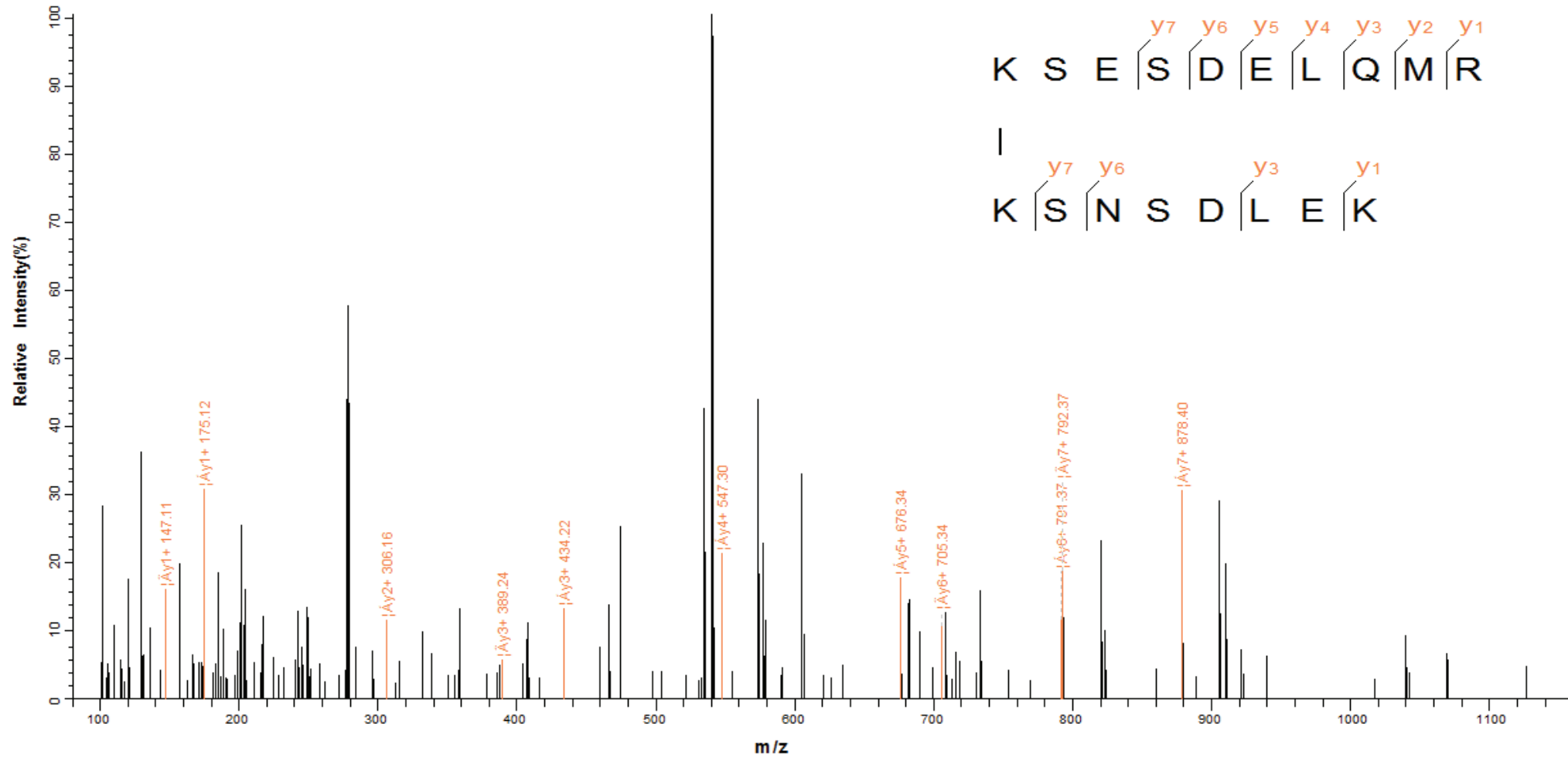


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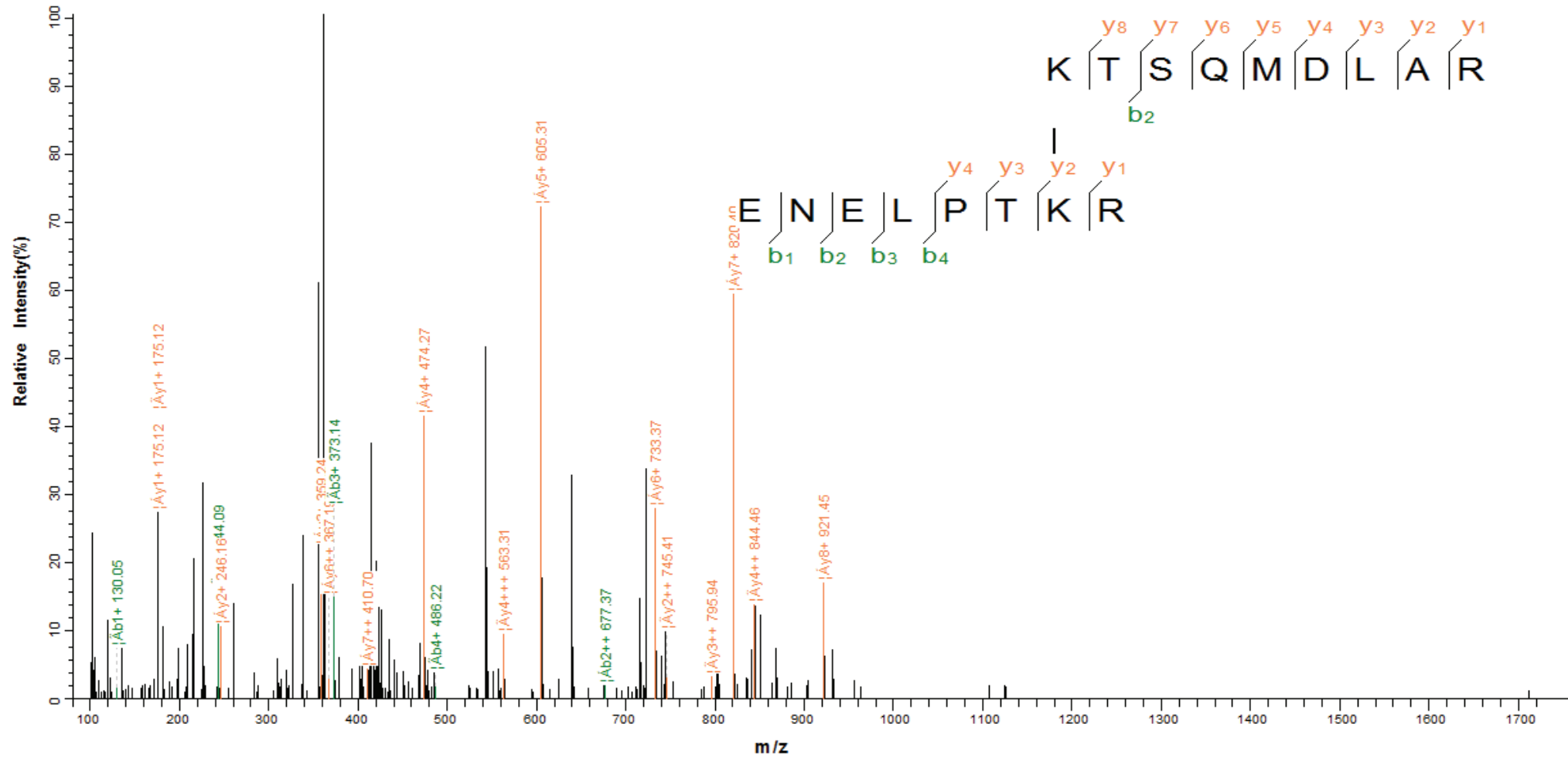
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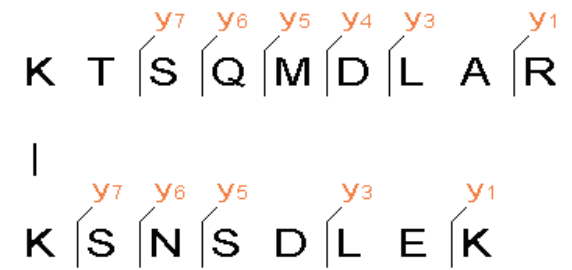
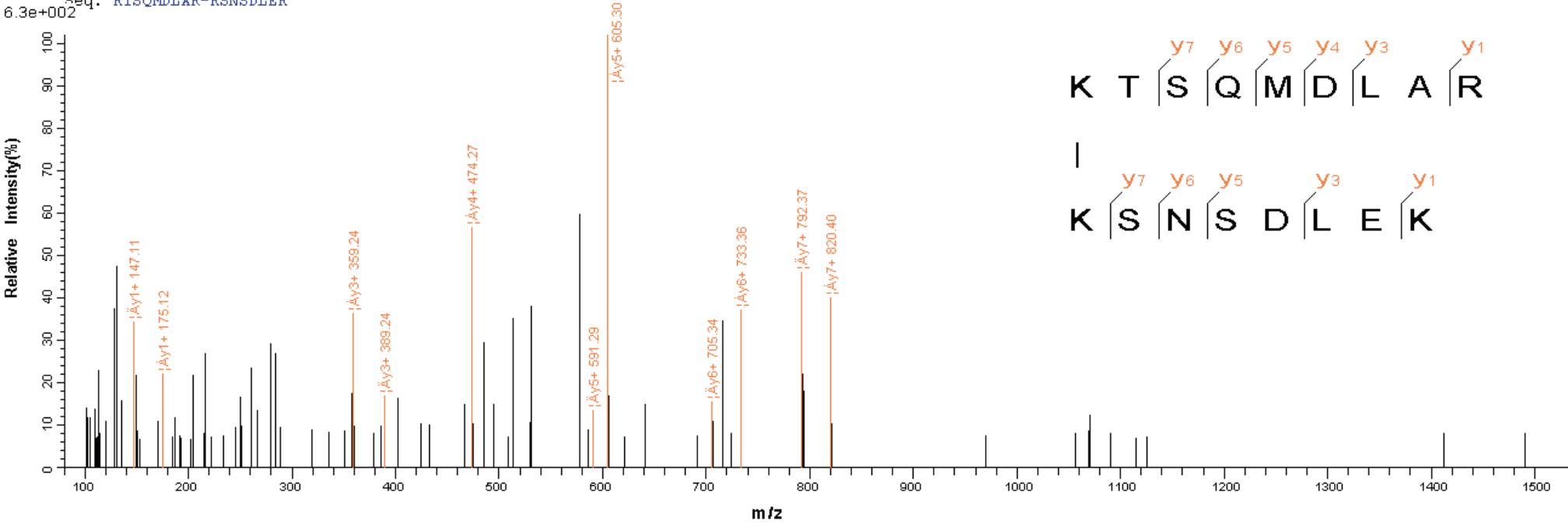


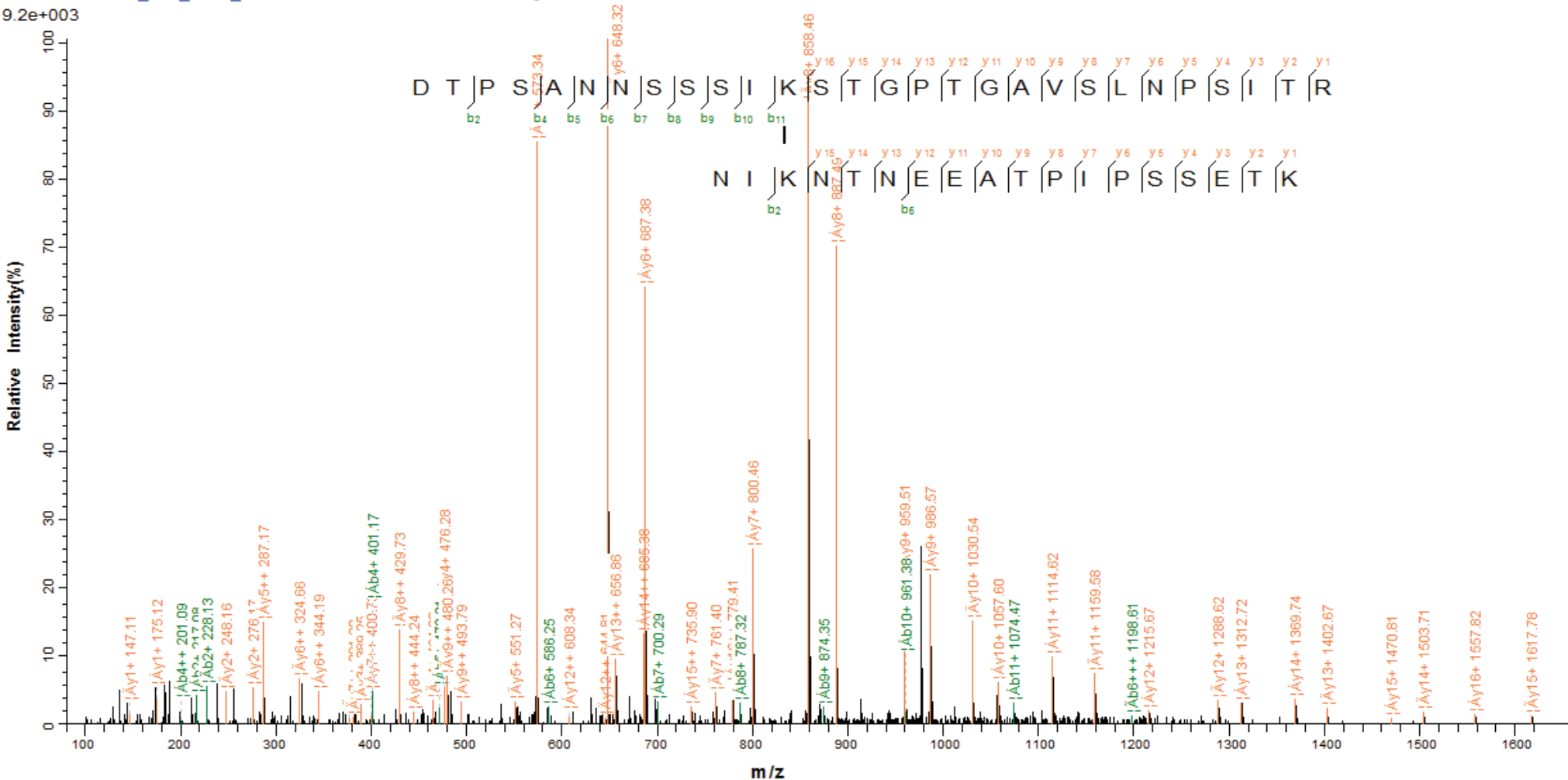
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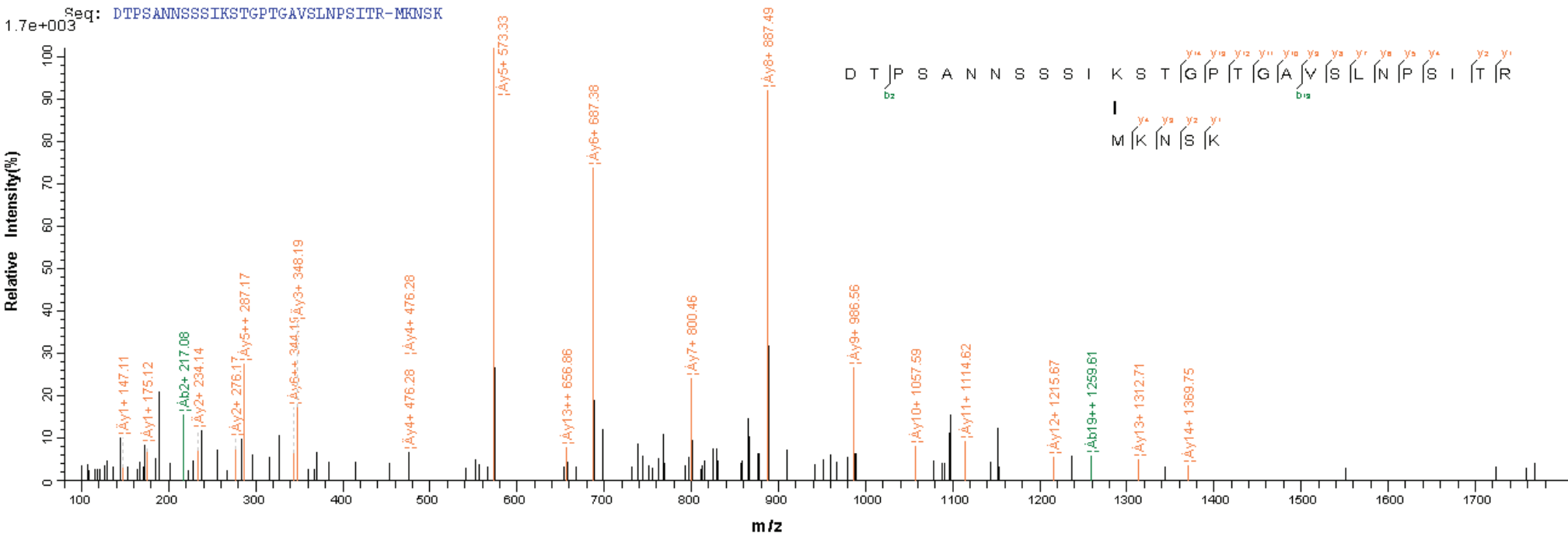


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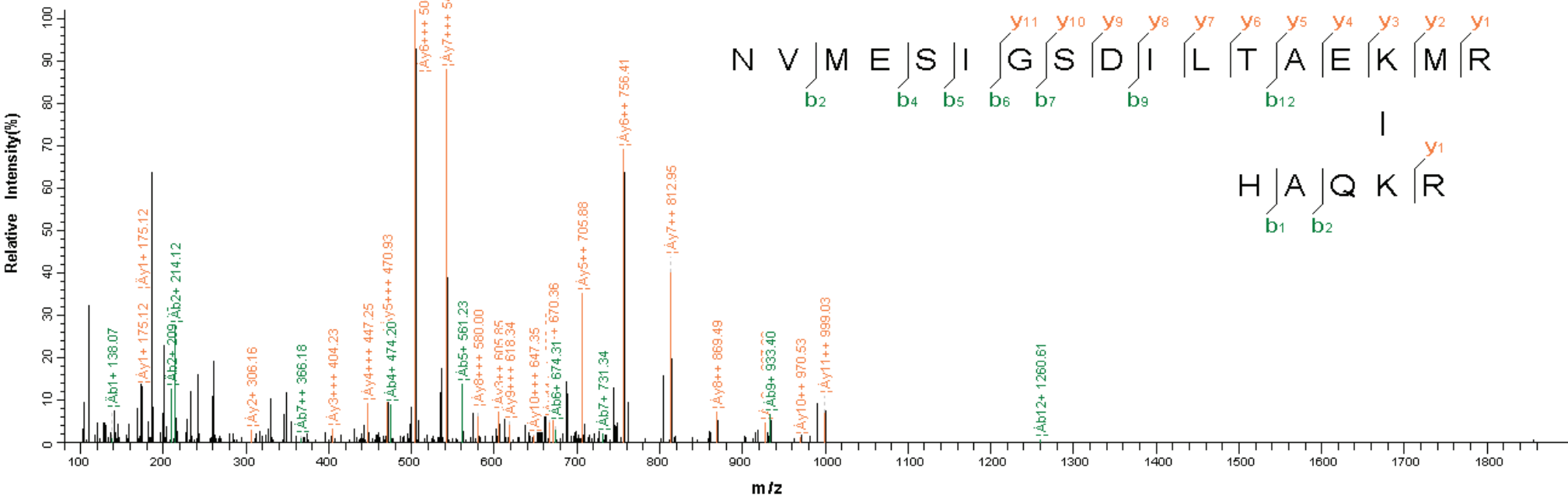




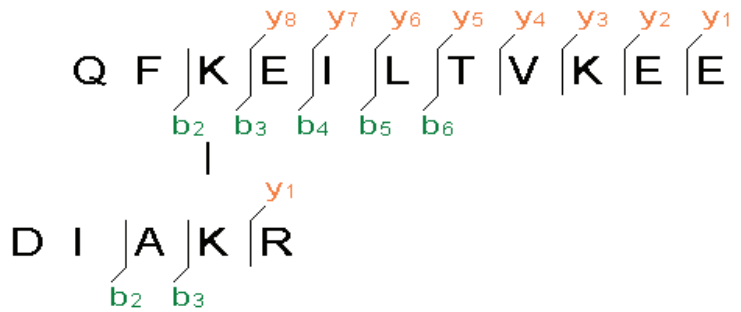
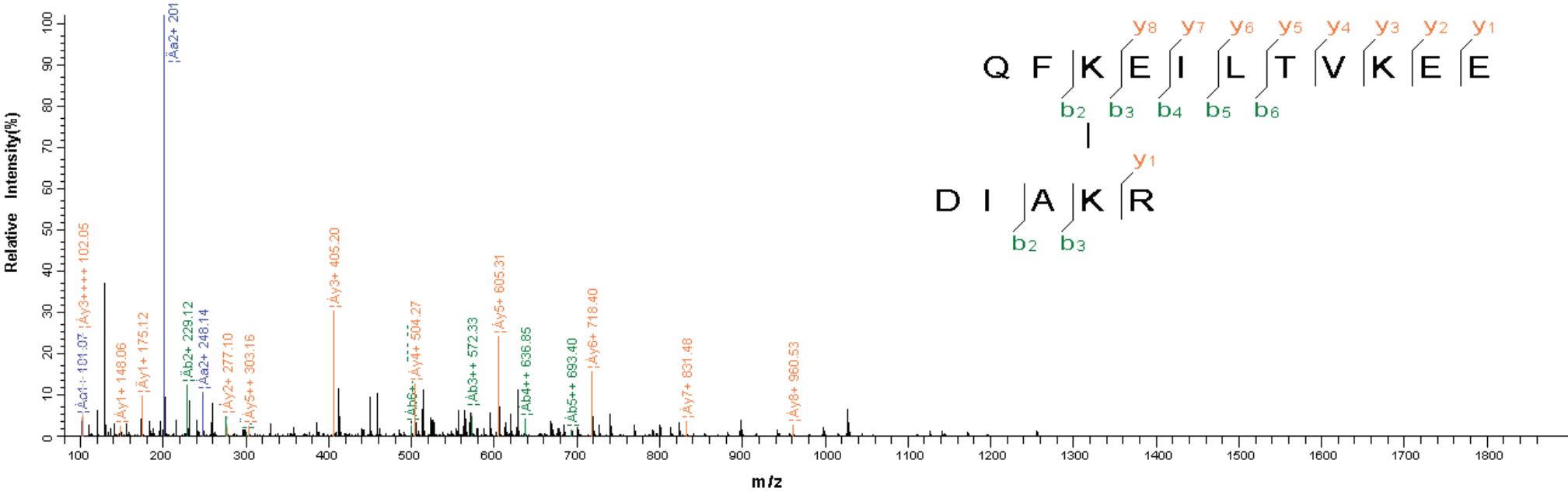
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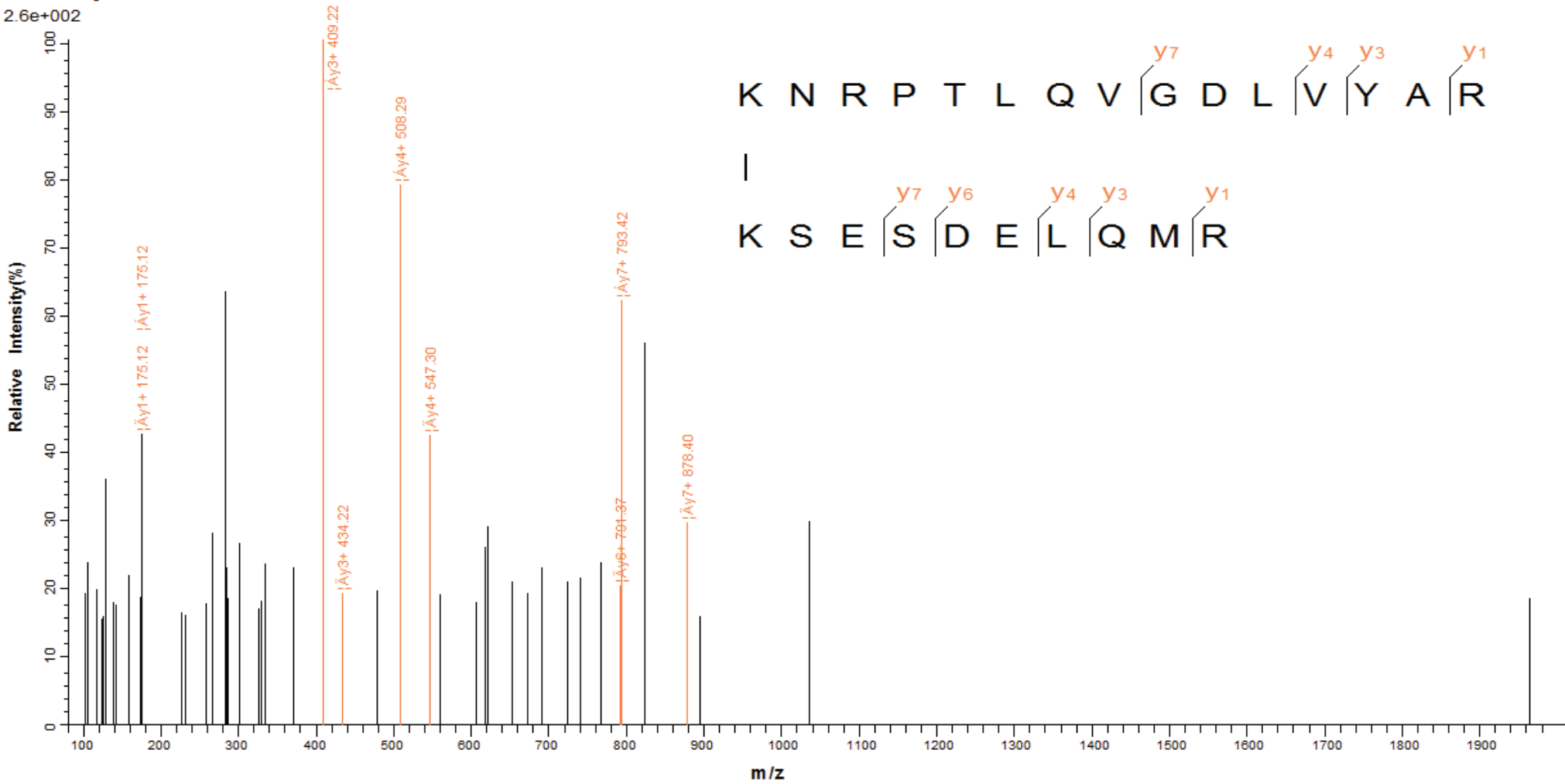
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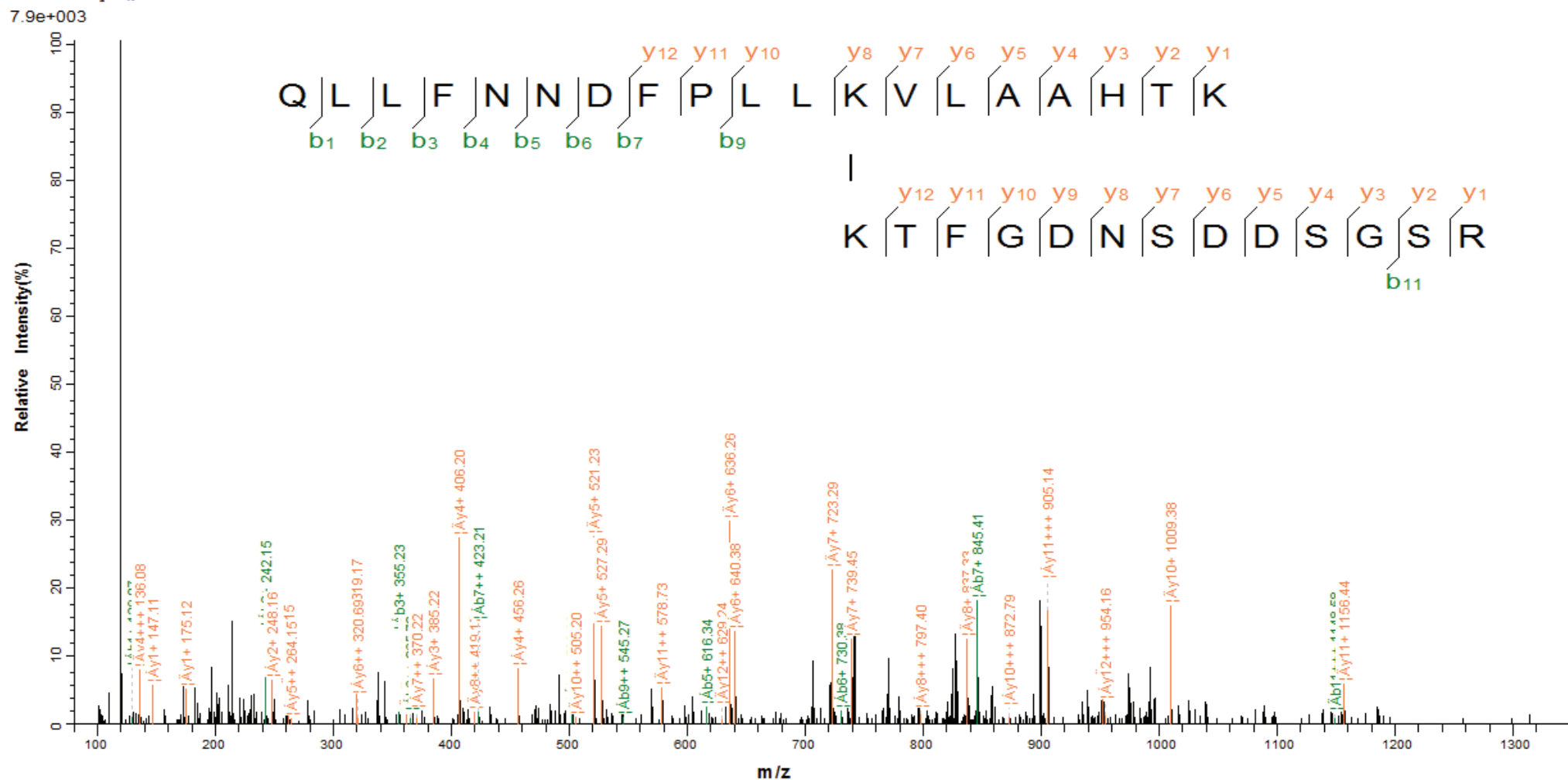
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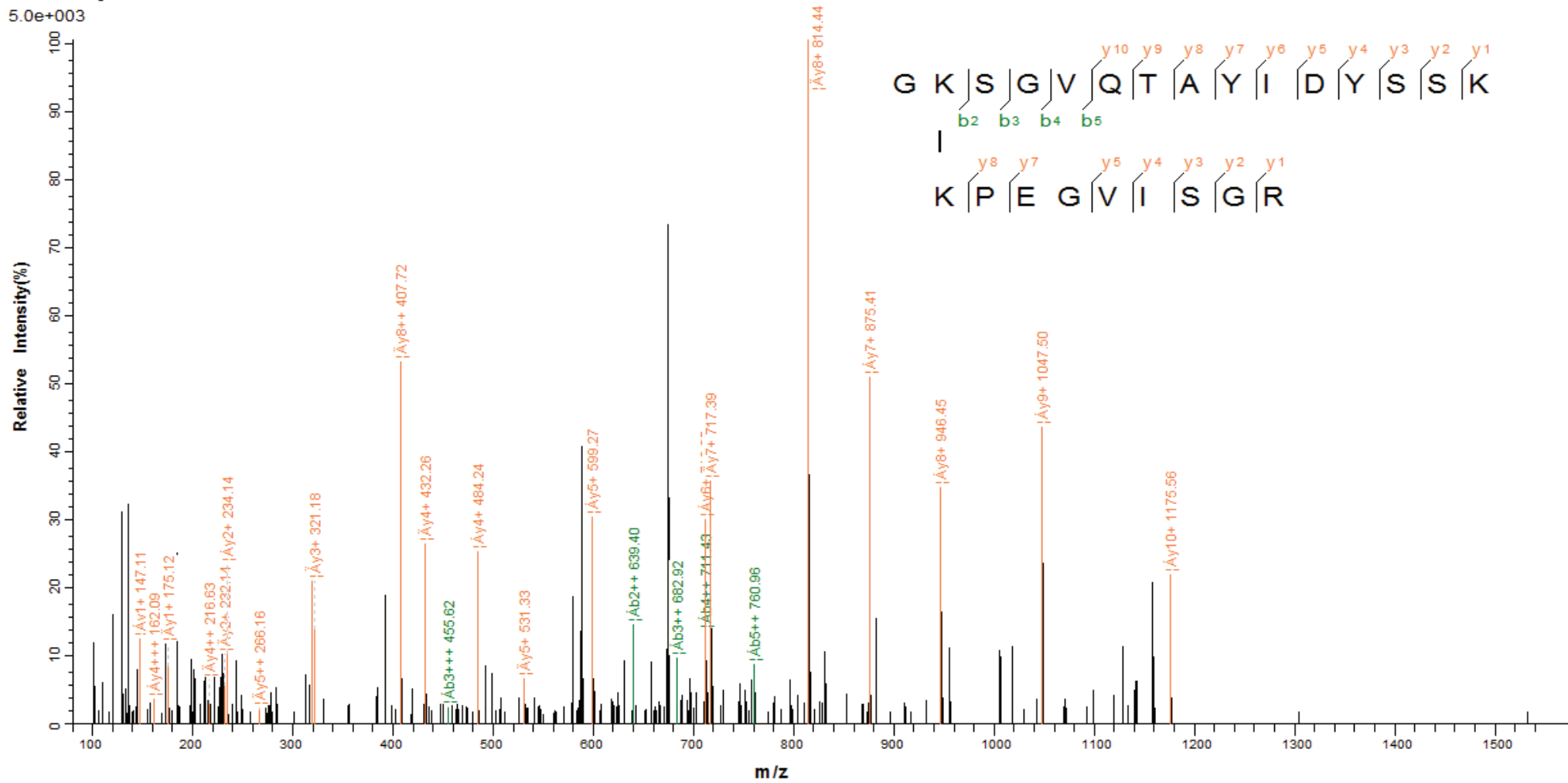
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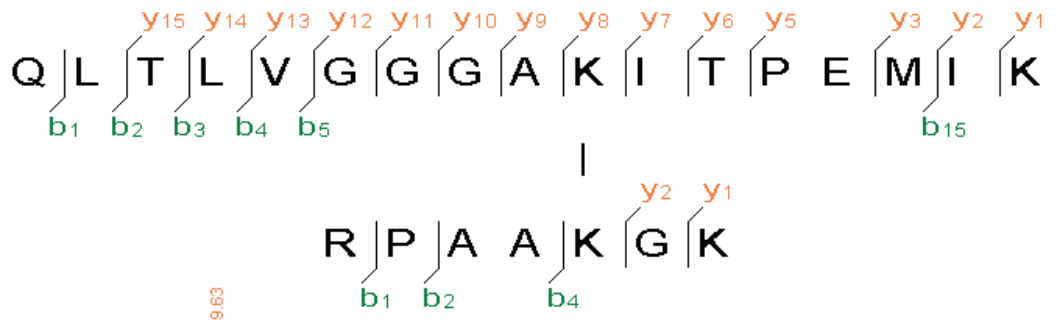
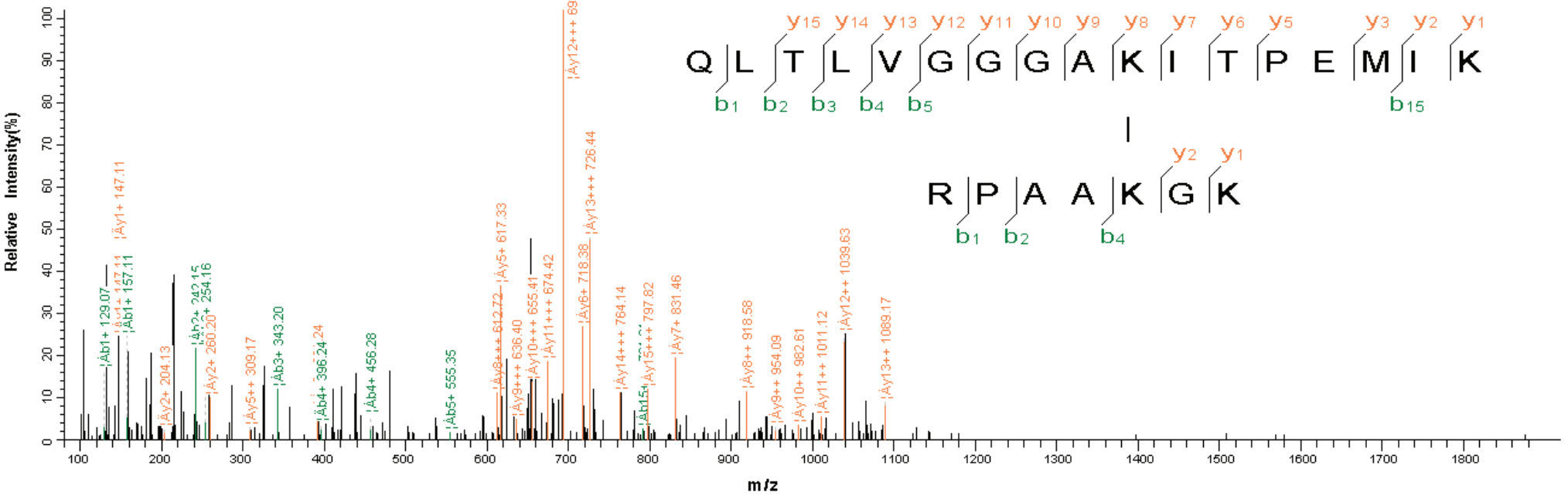
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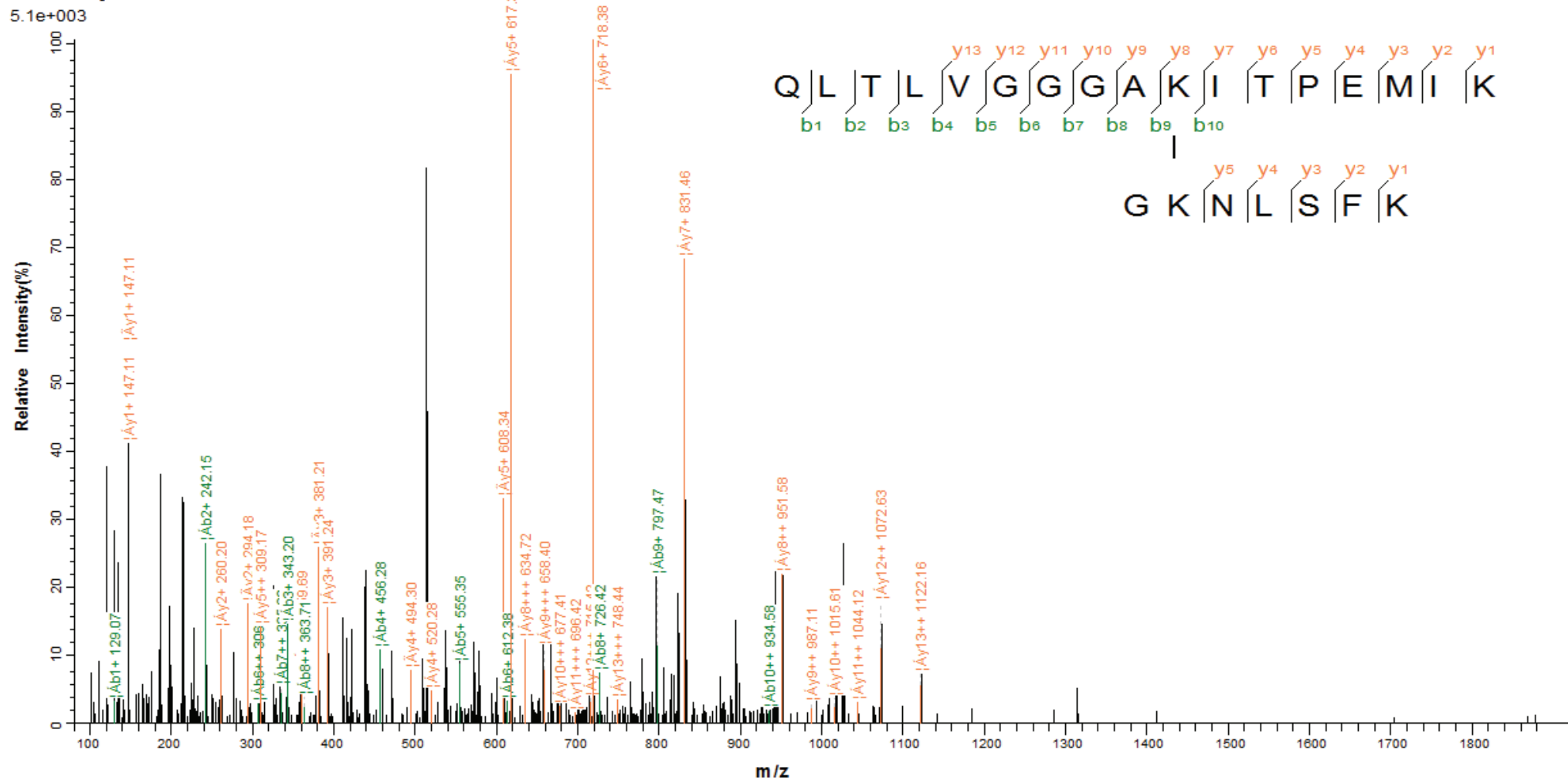
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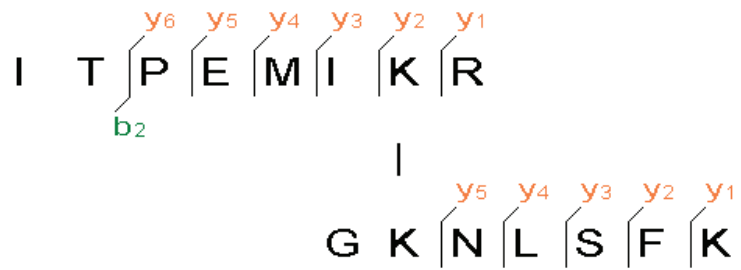
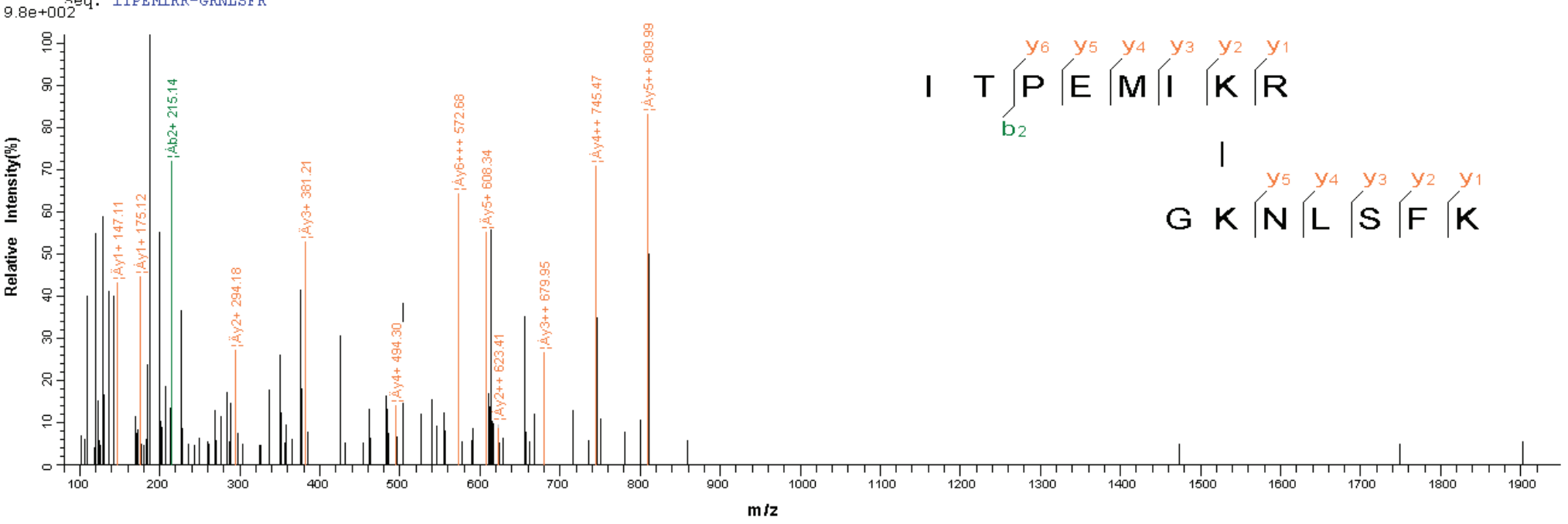
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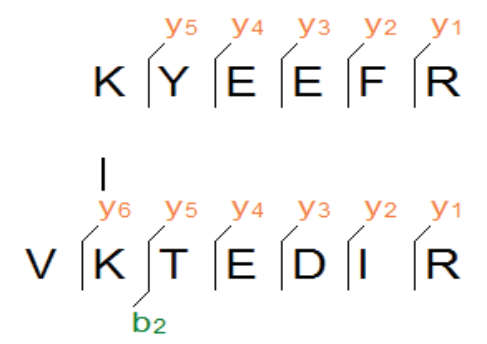
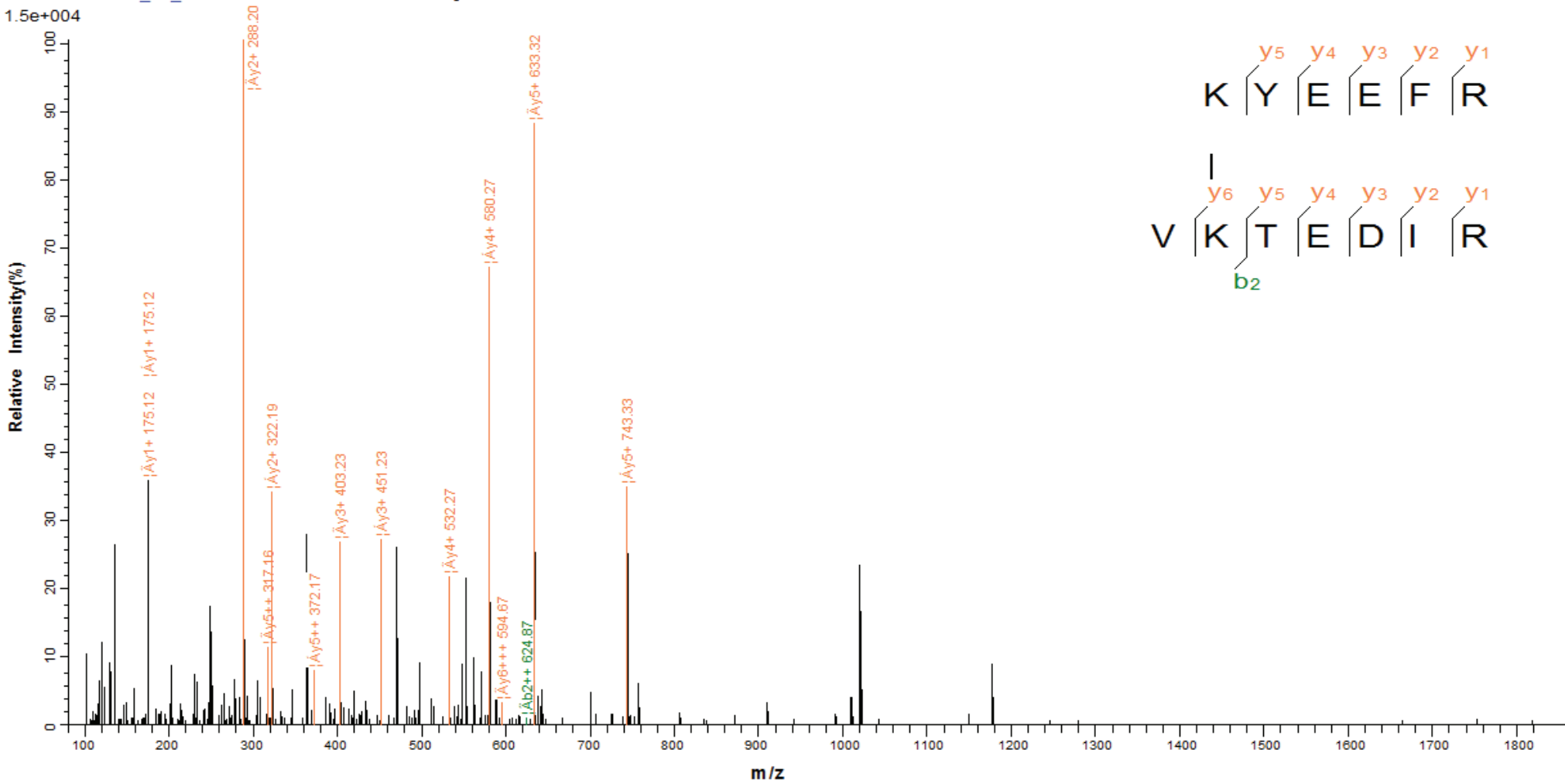
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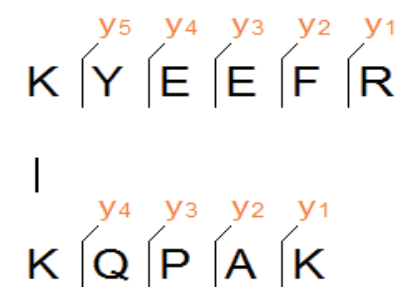
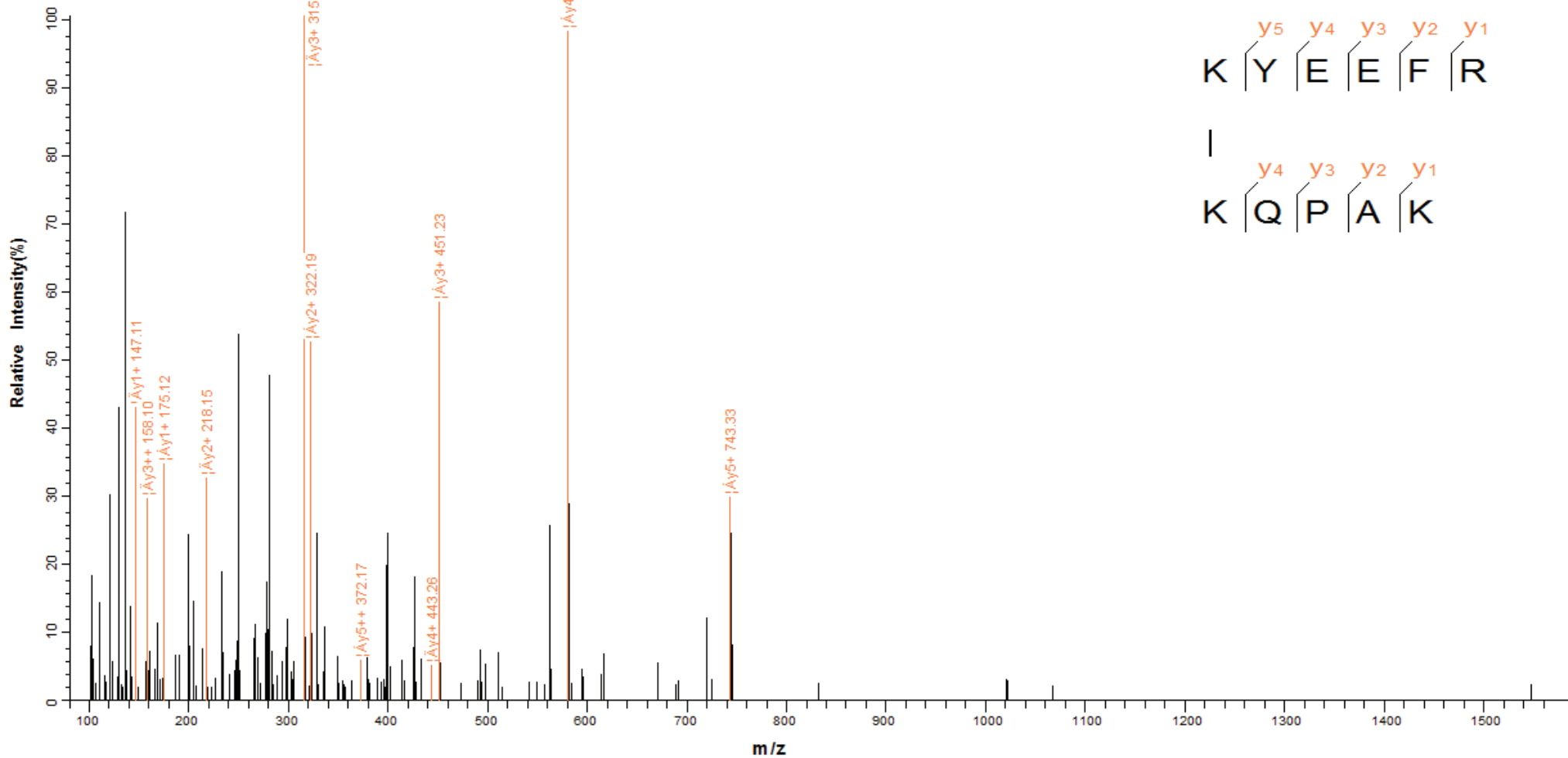


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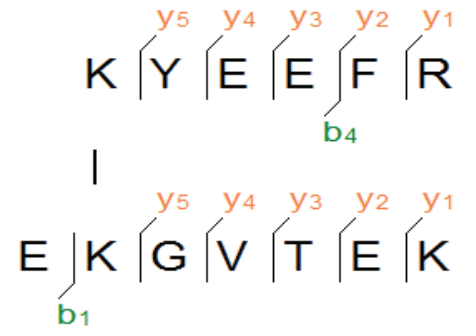
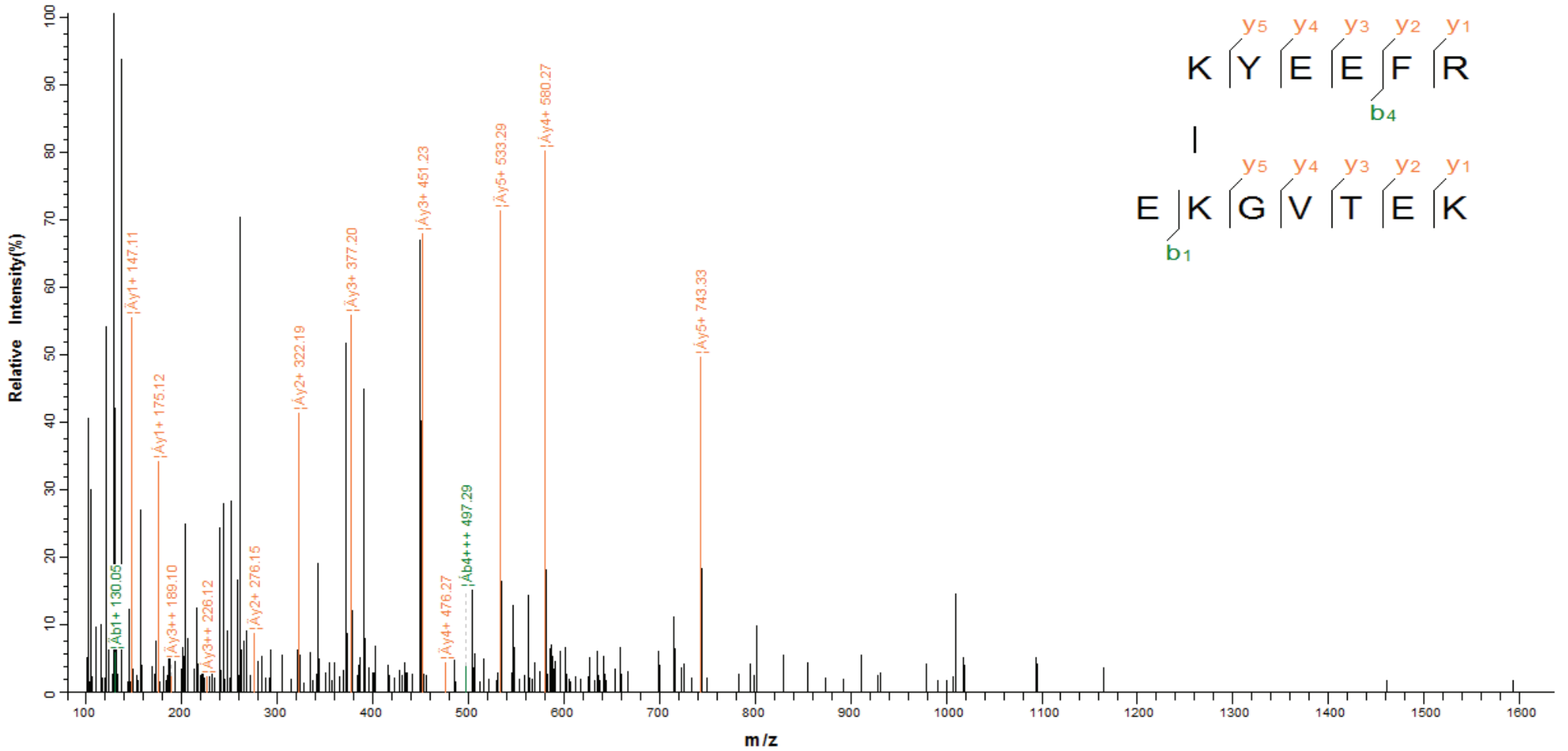
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2.3e+003



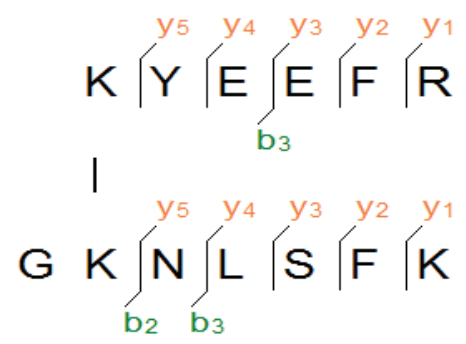
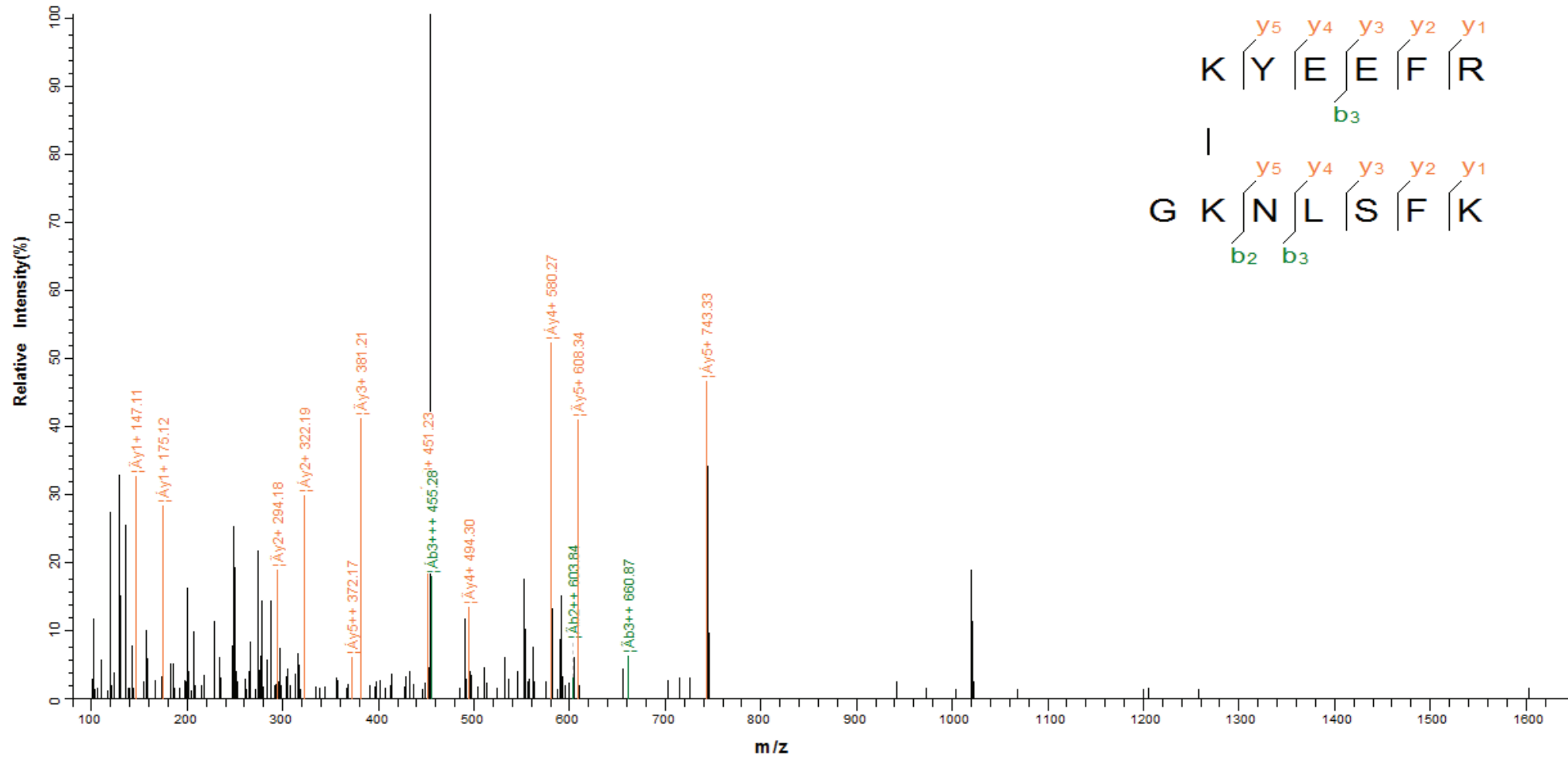
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3.0e+003

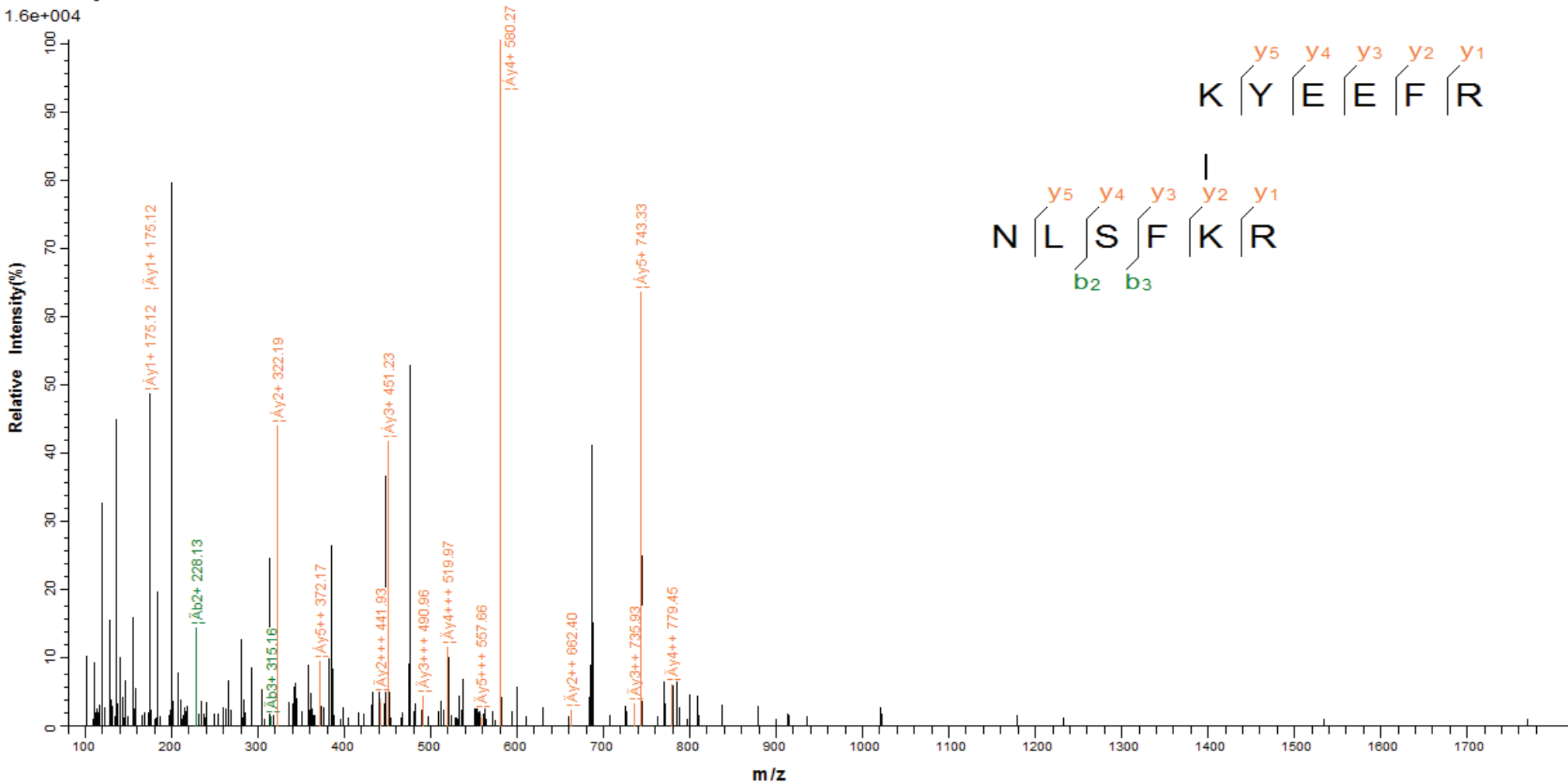


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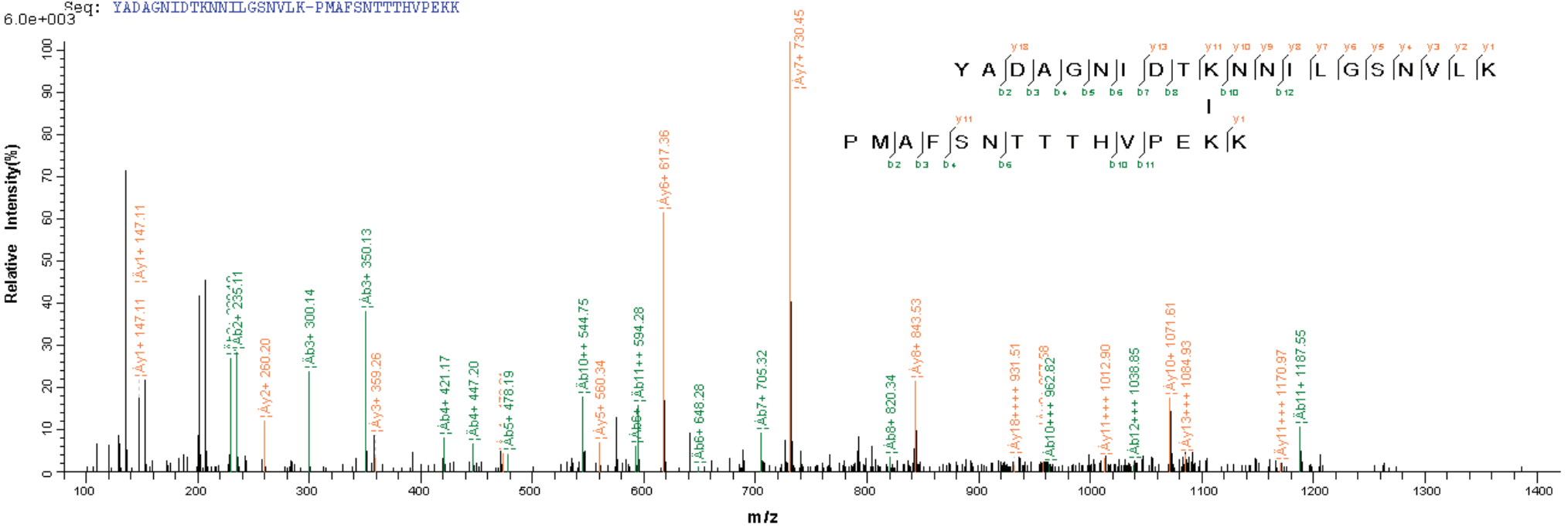
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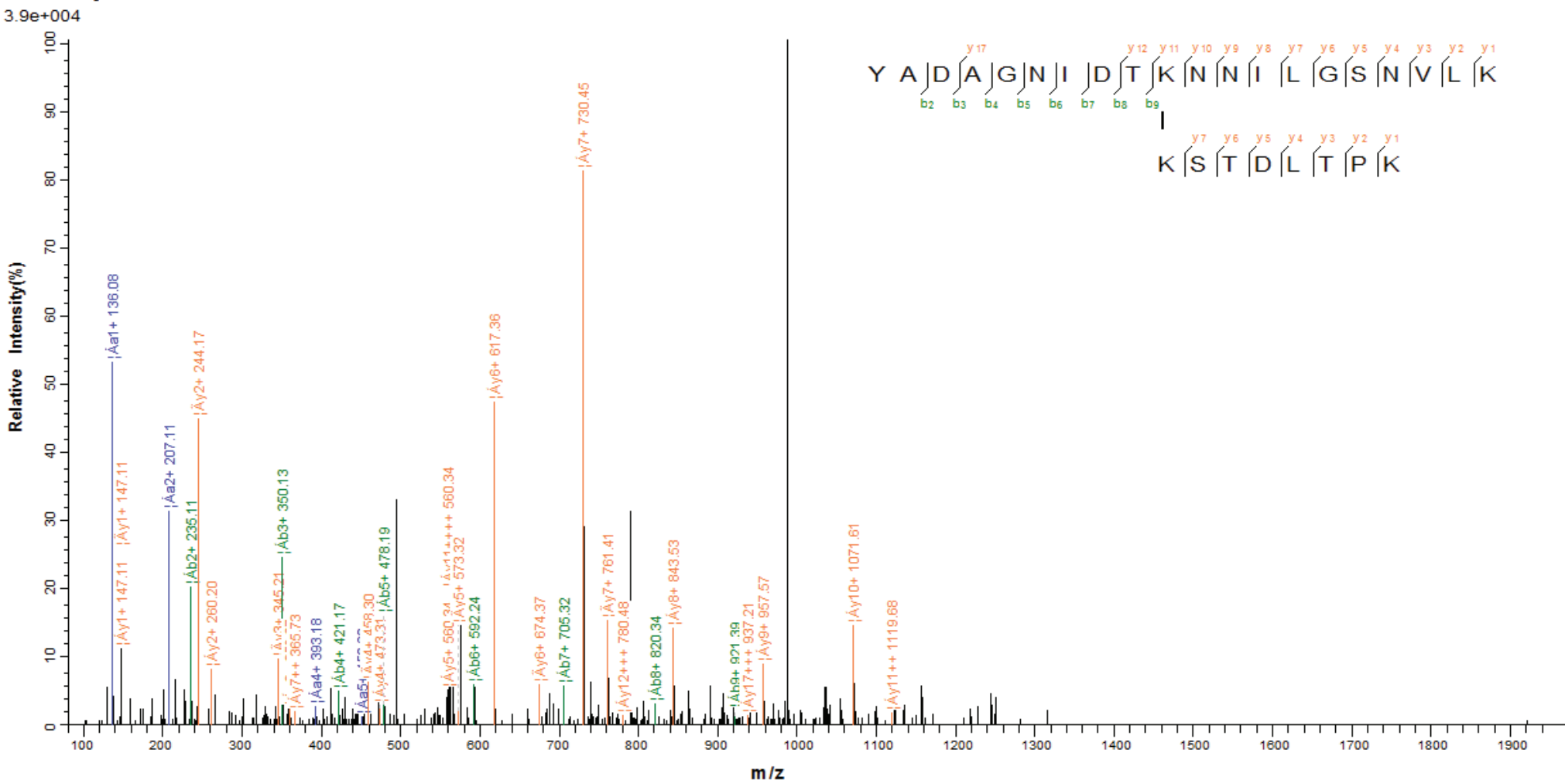
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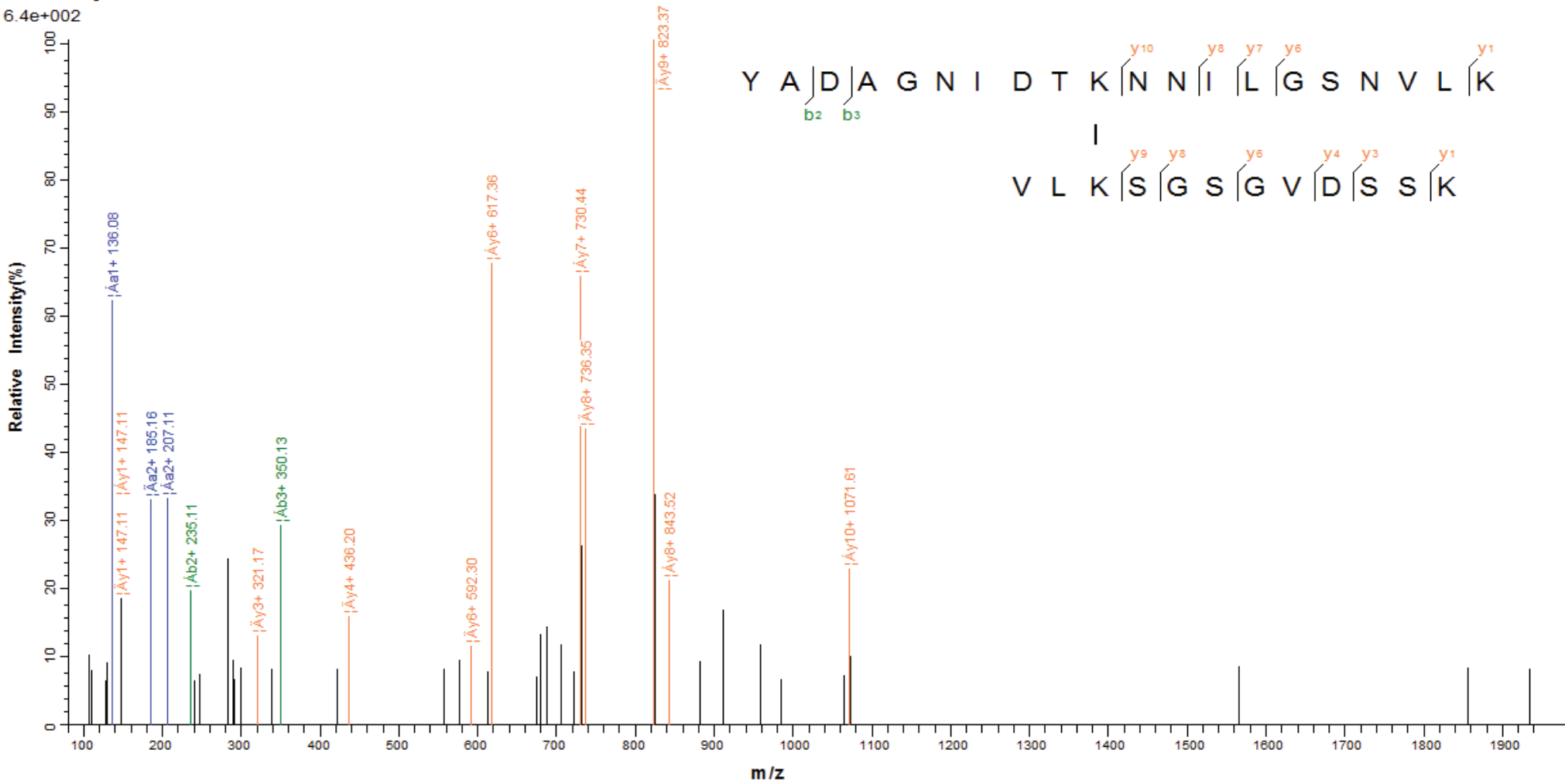
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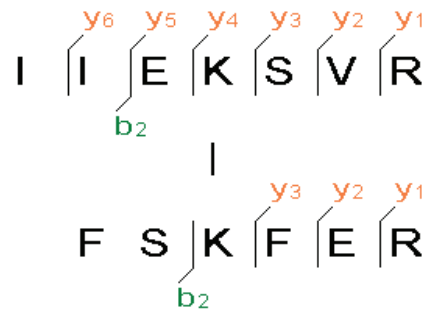
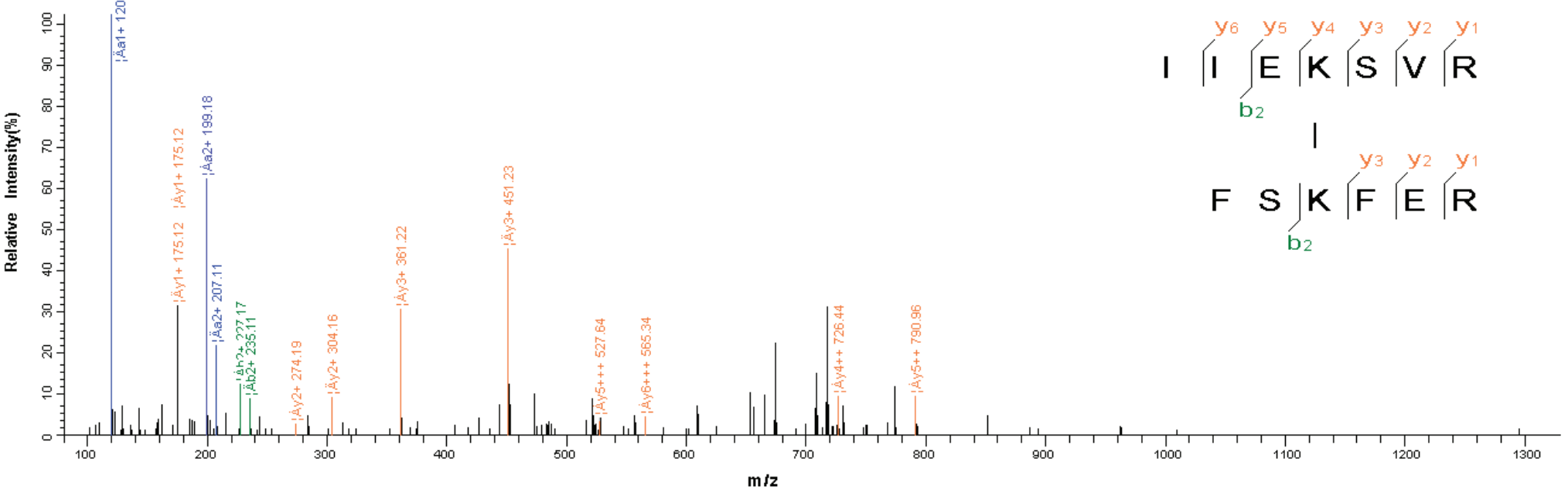
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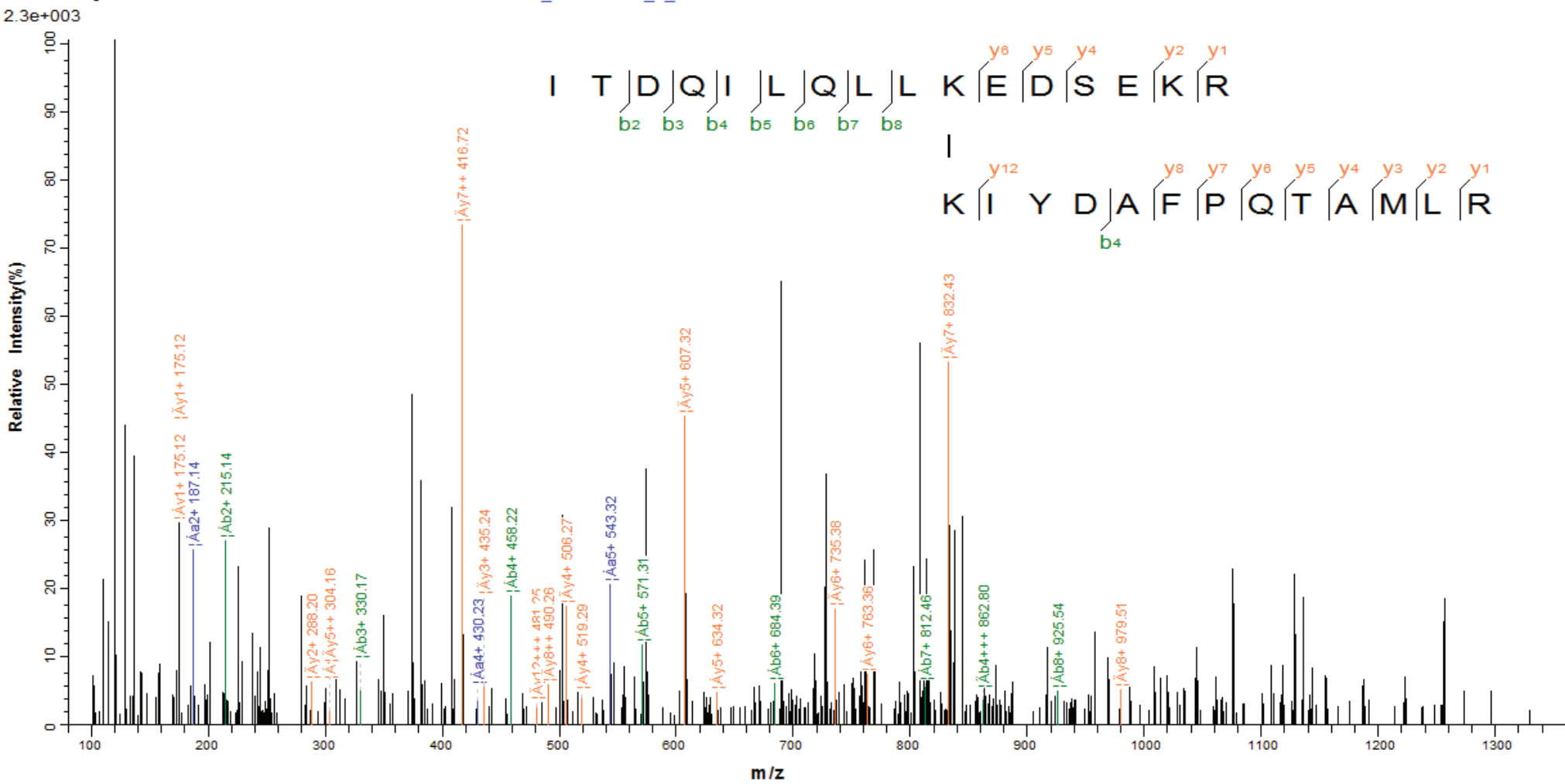
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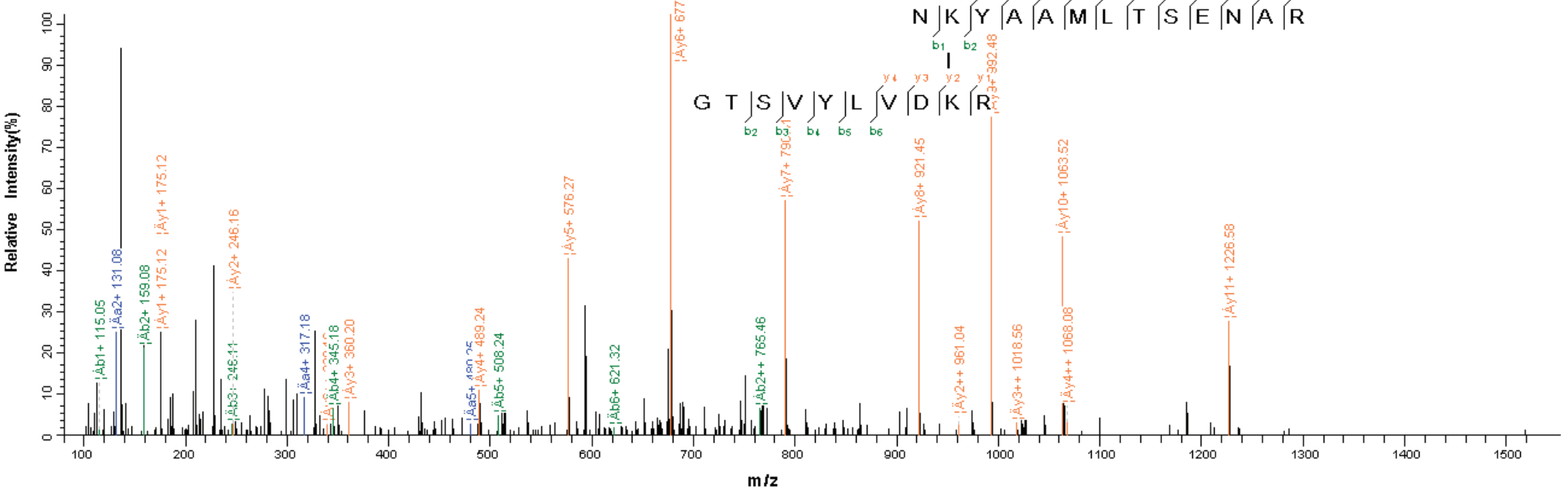
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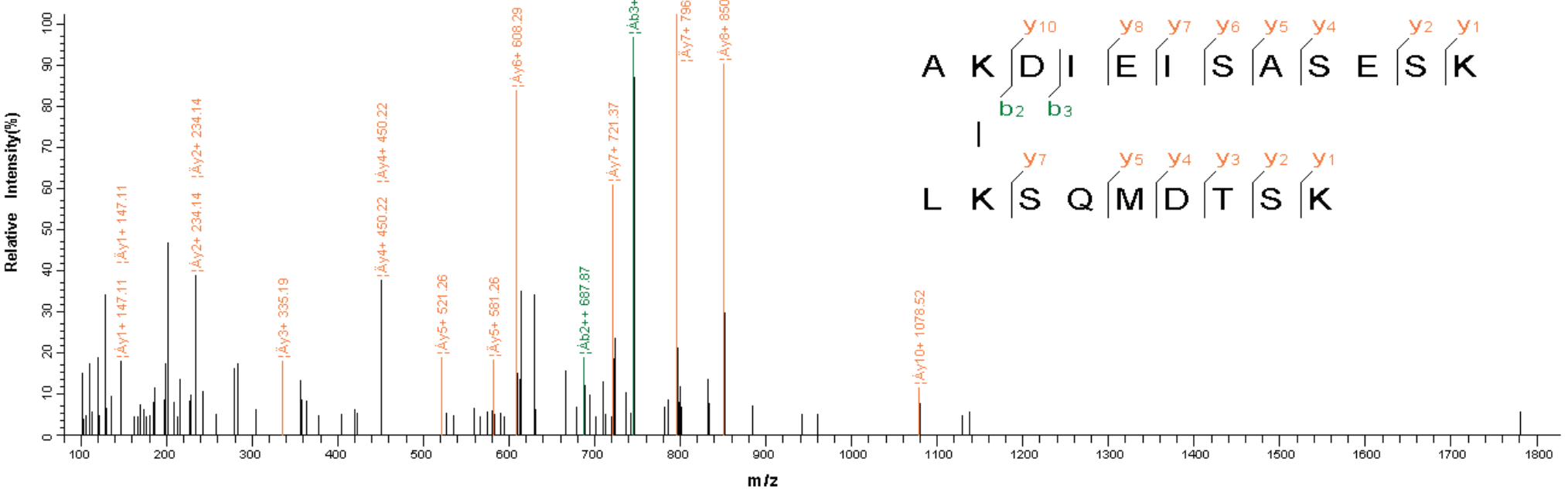
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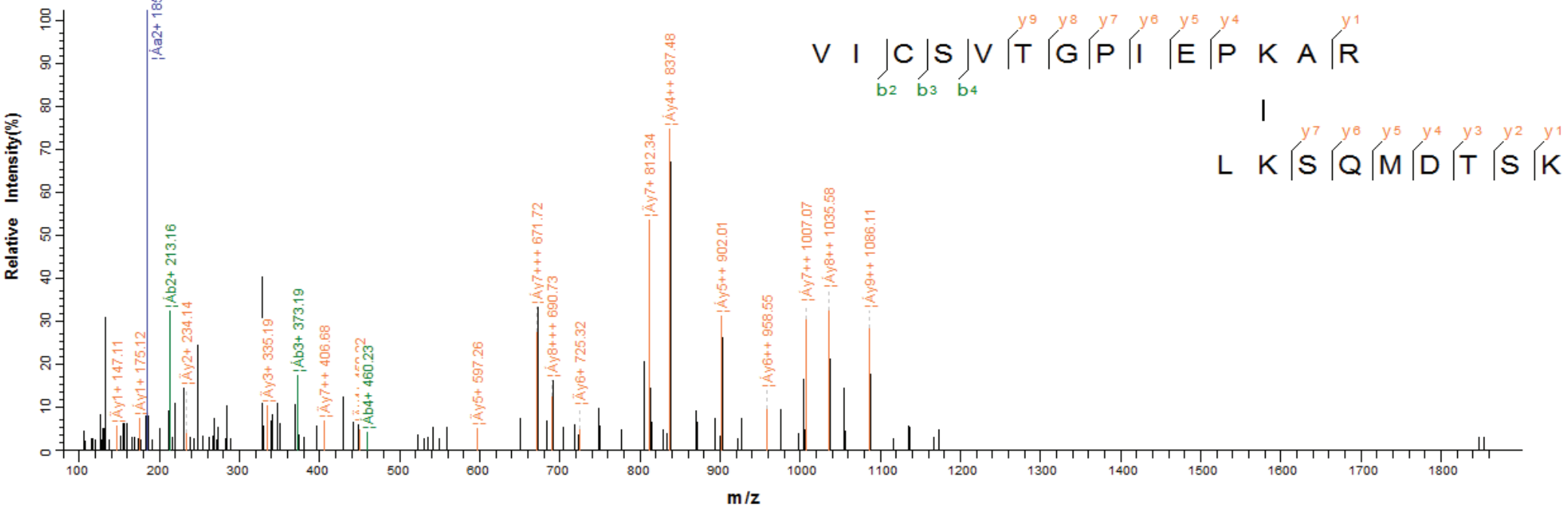
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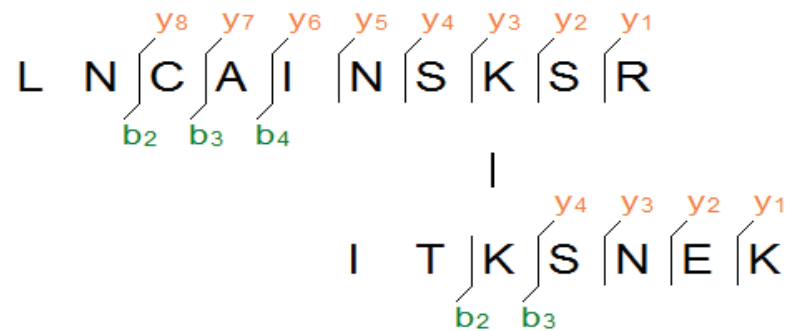
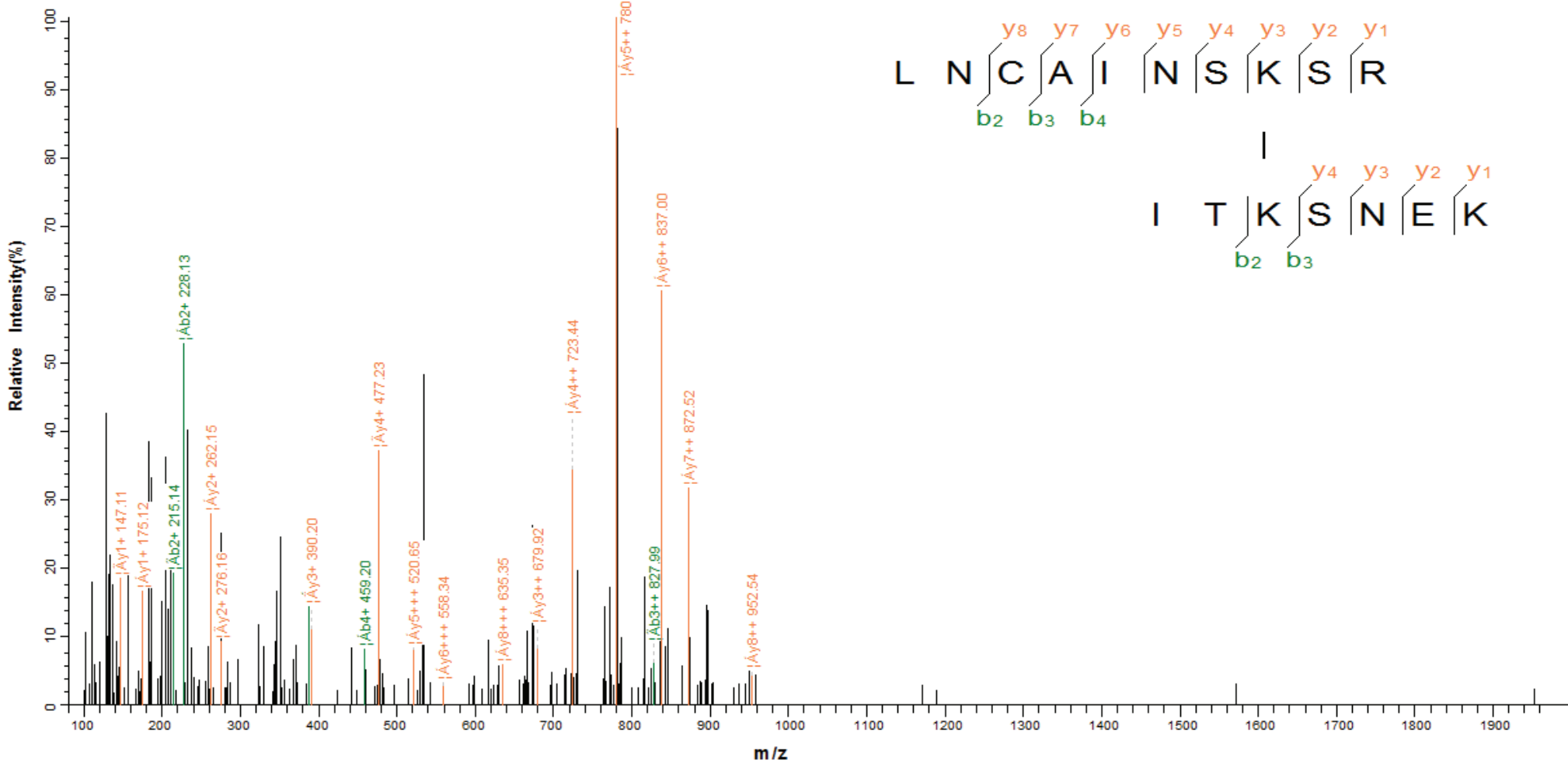


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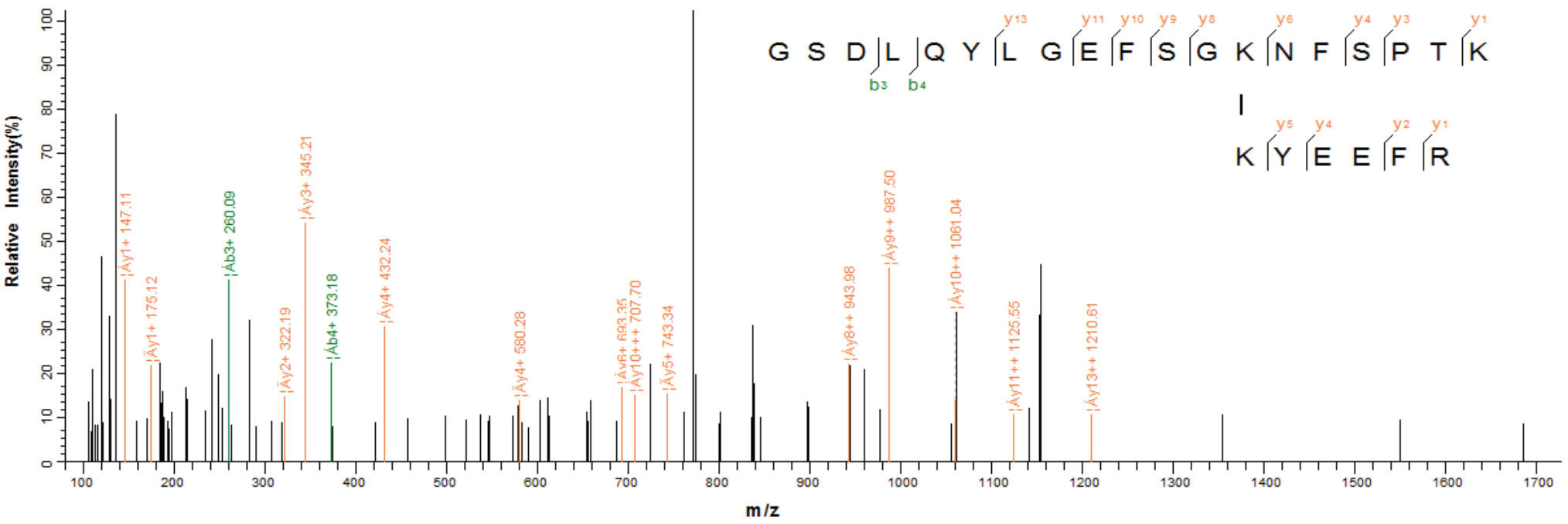


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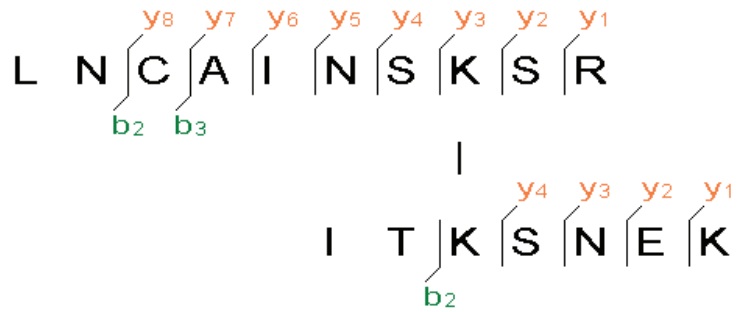
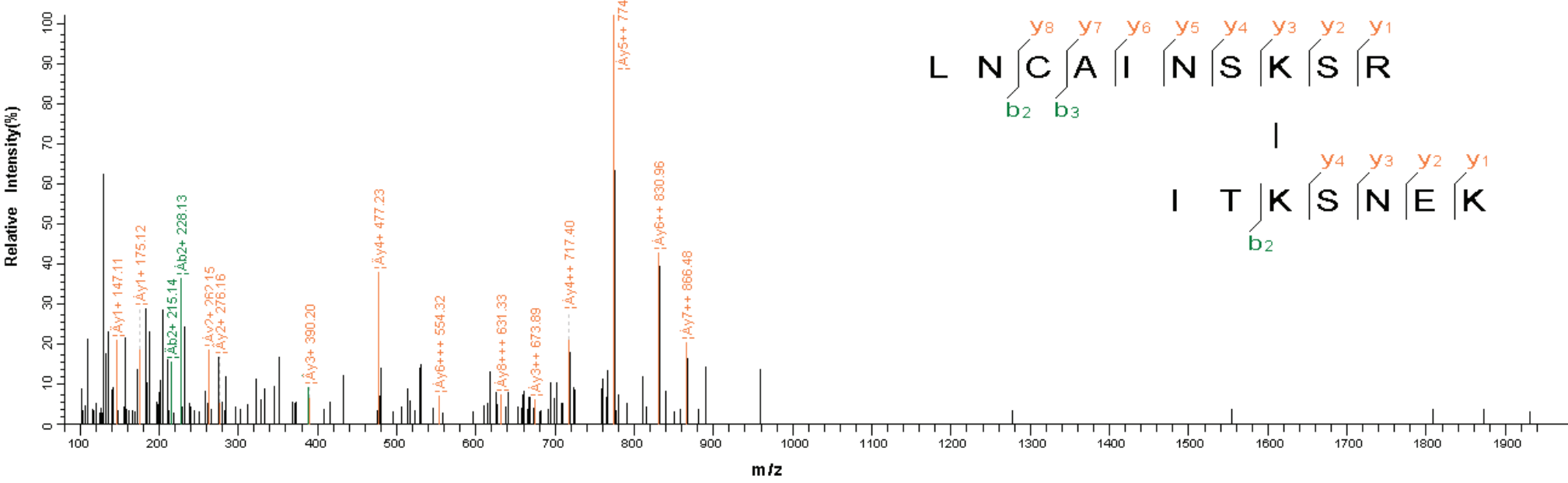
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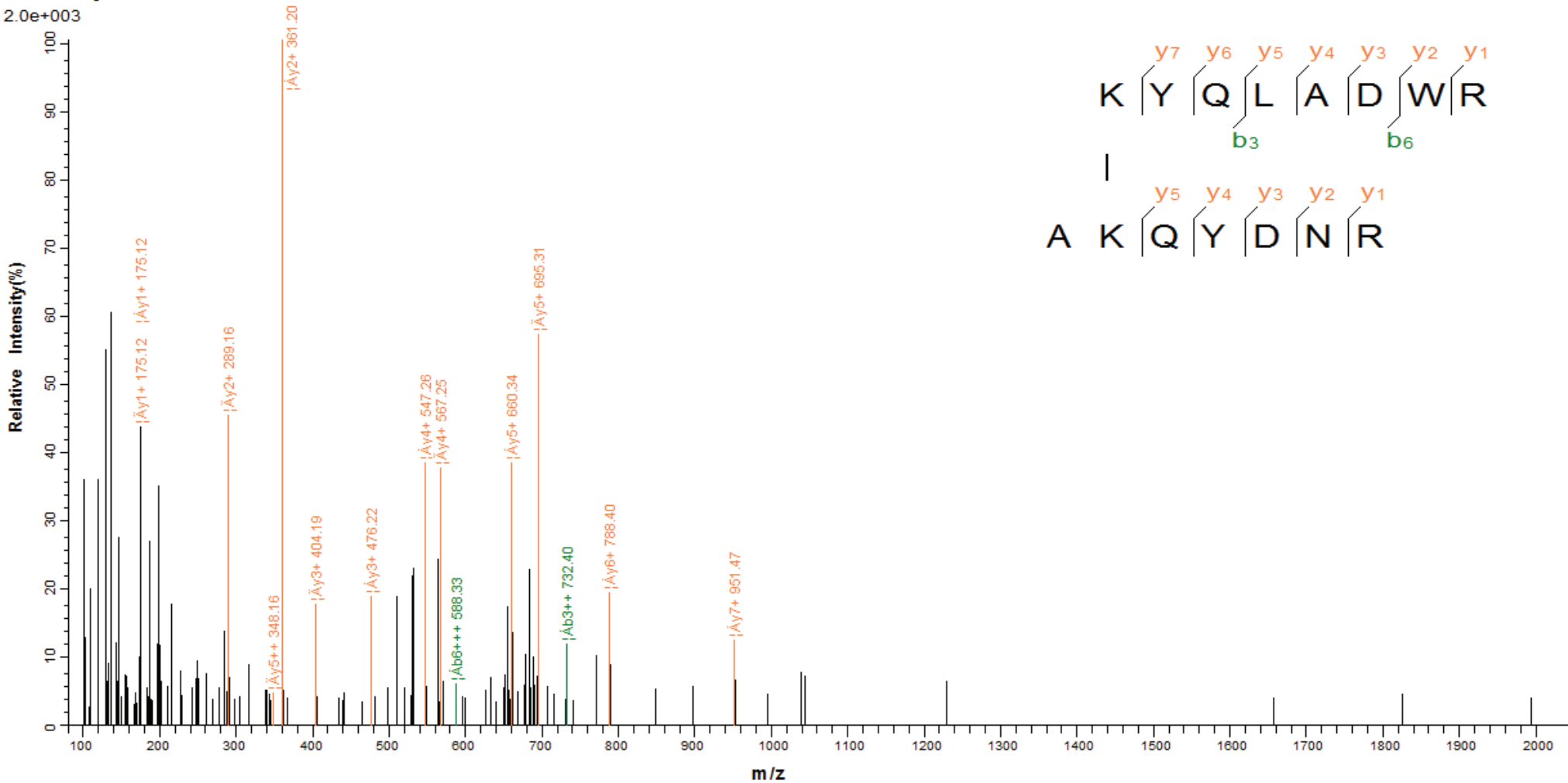
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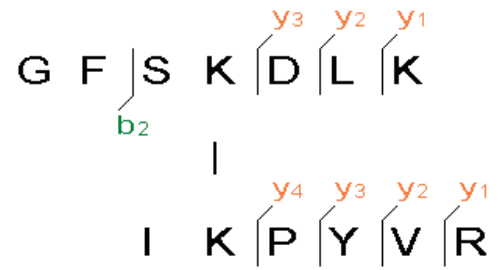
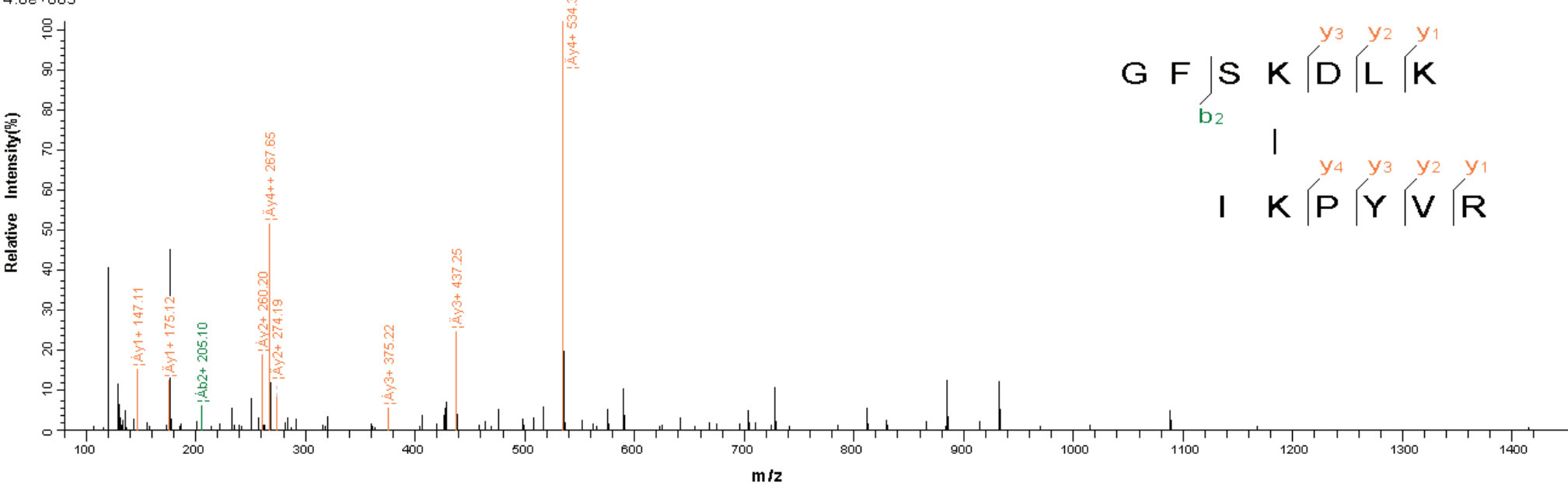
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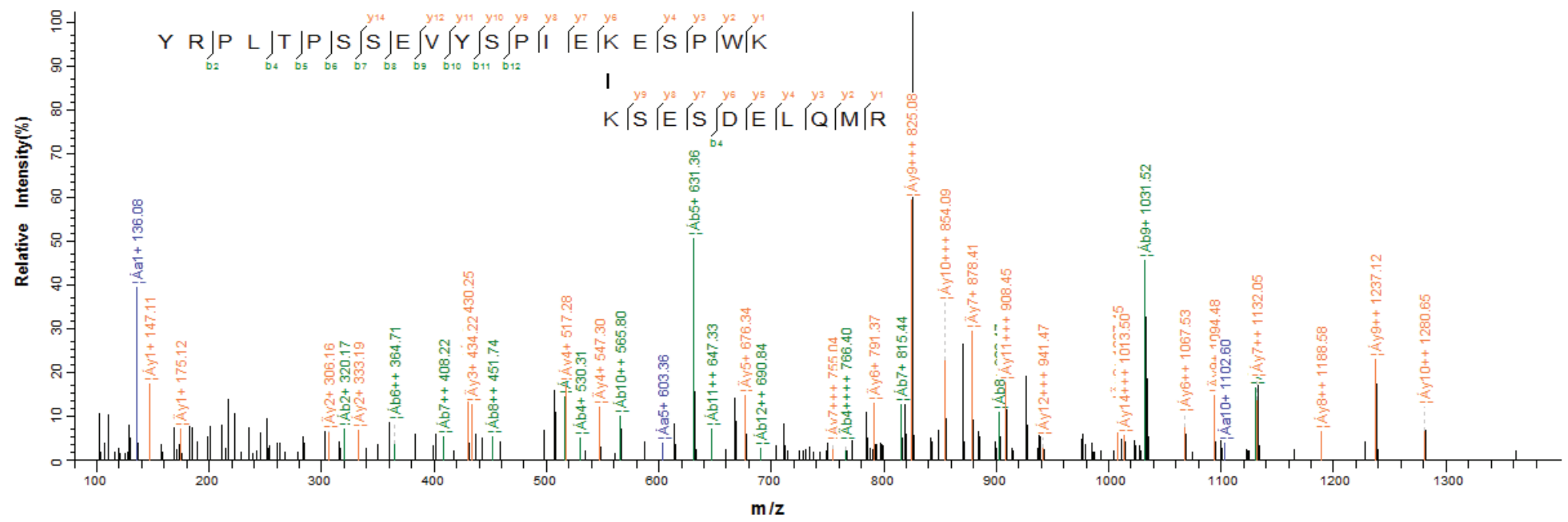
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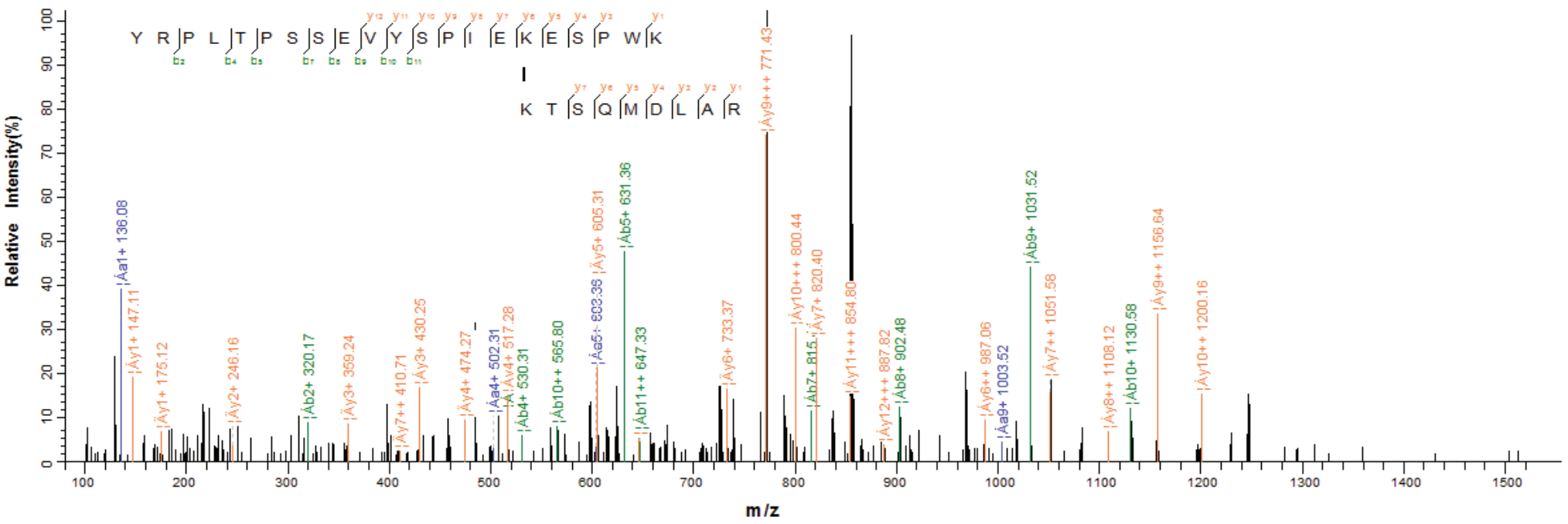
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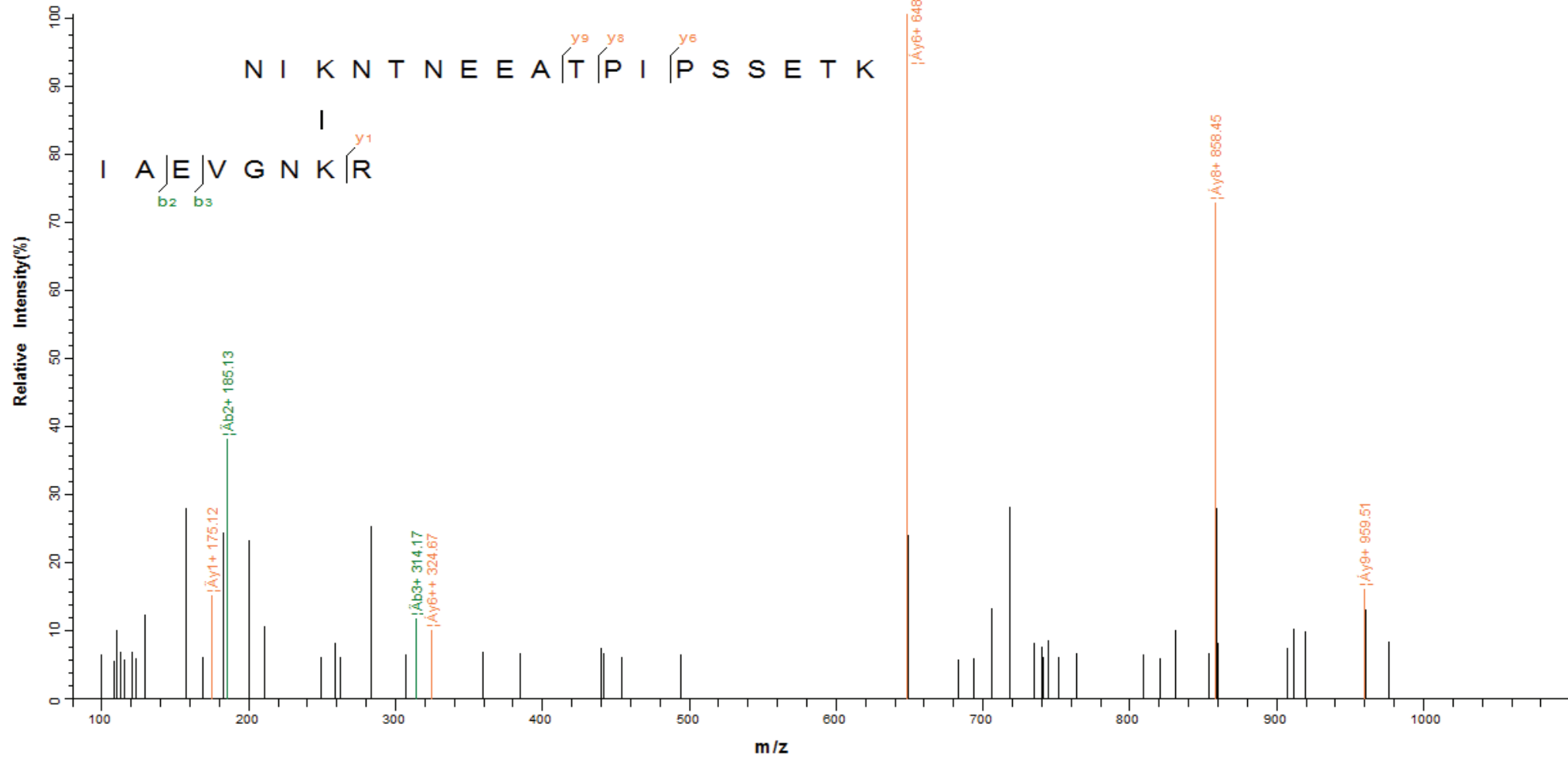


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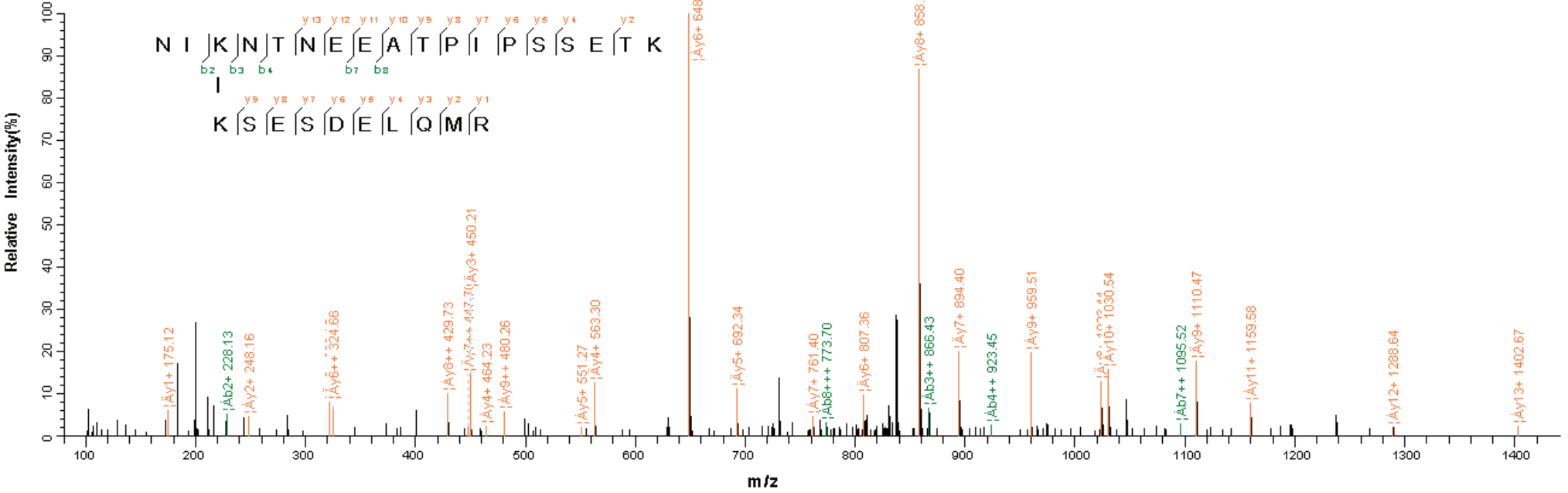


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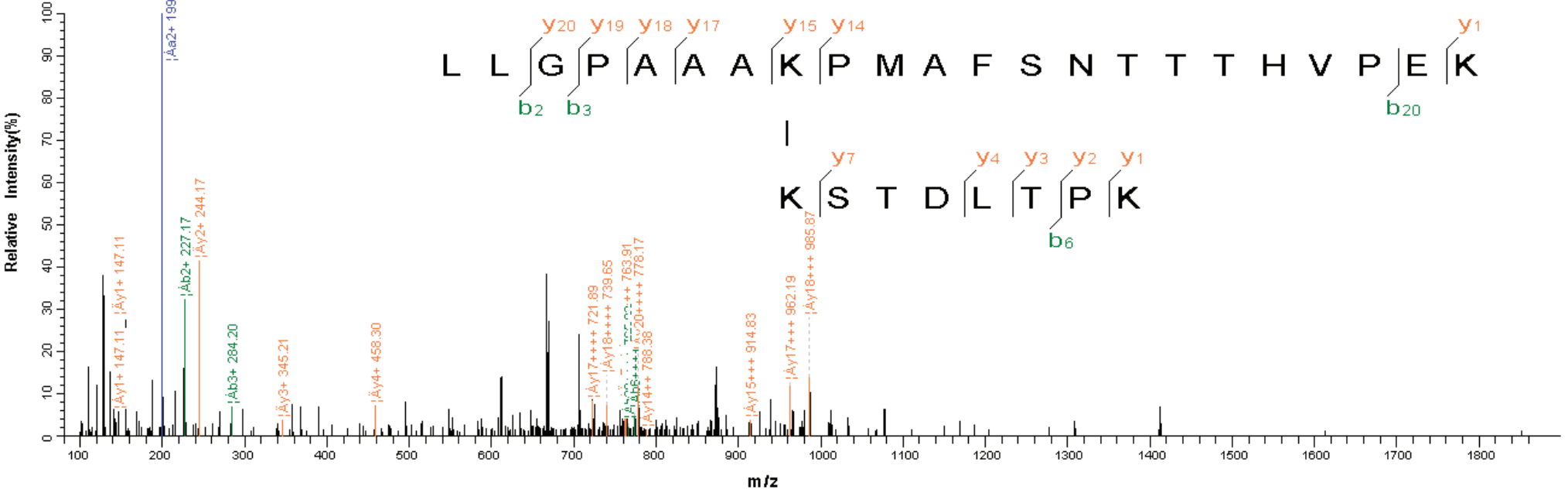
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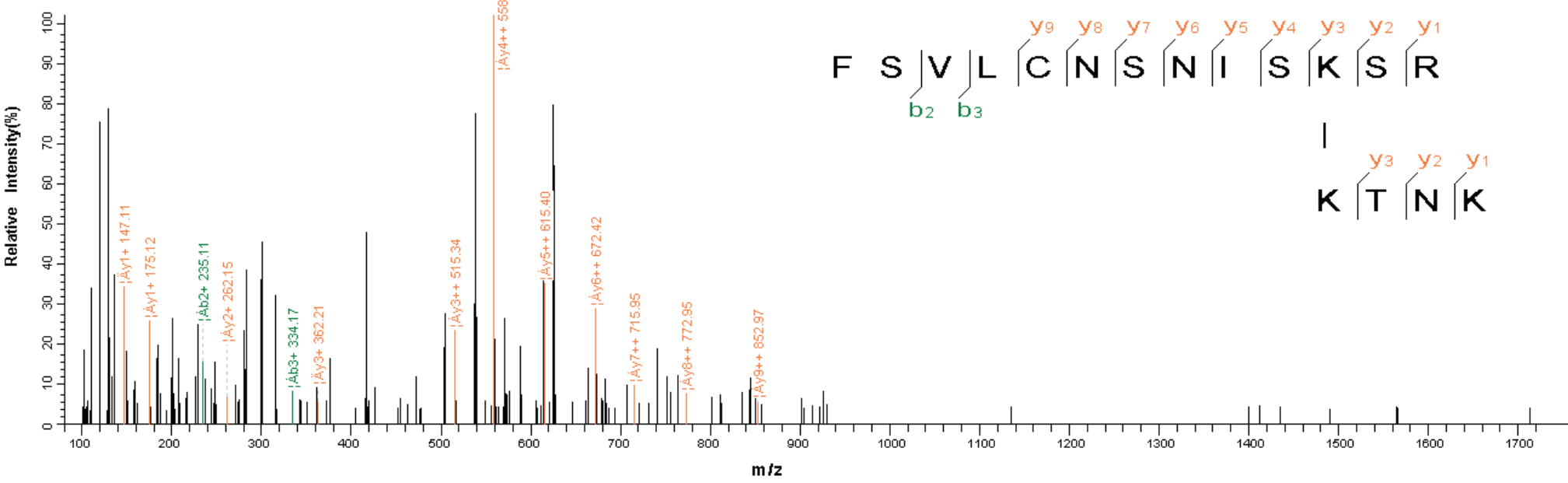
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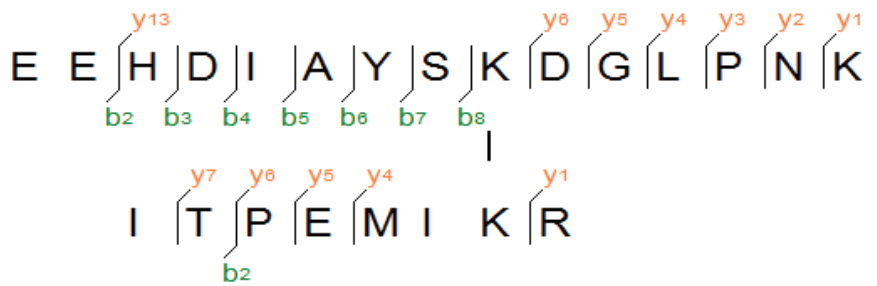
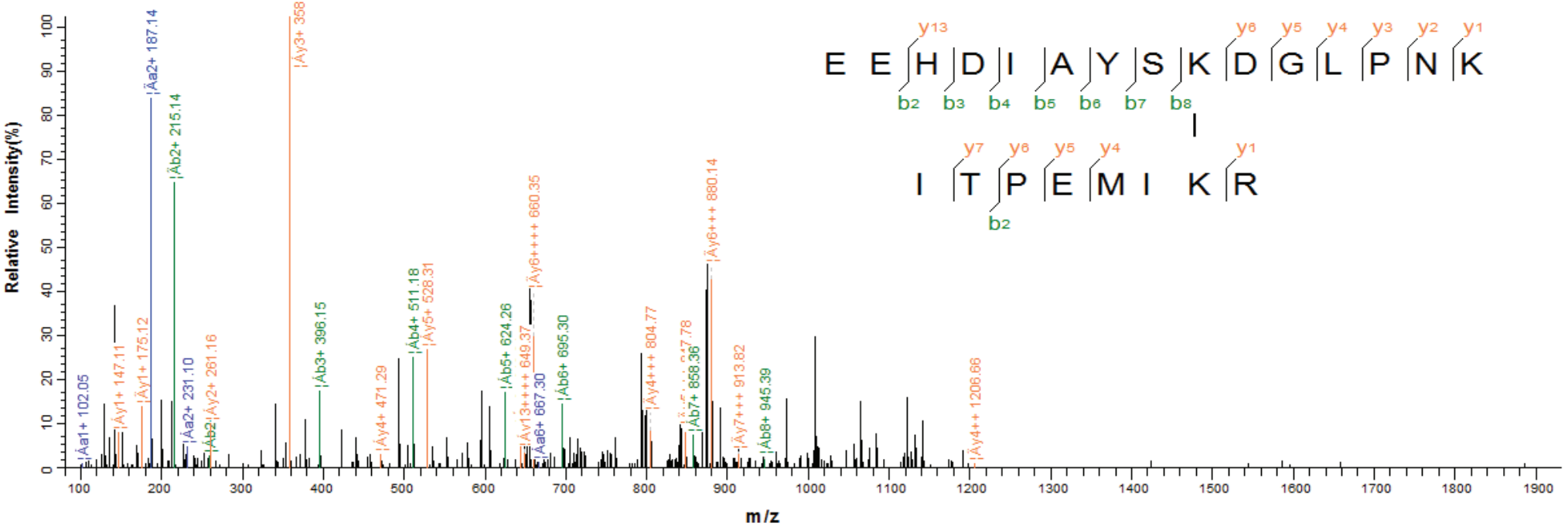
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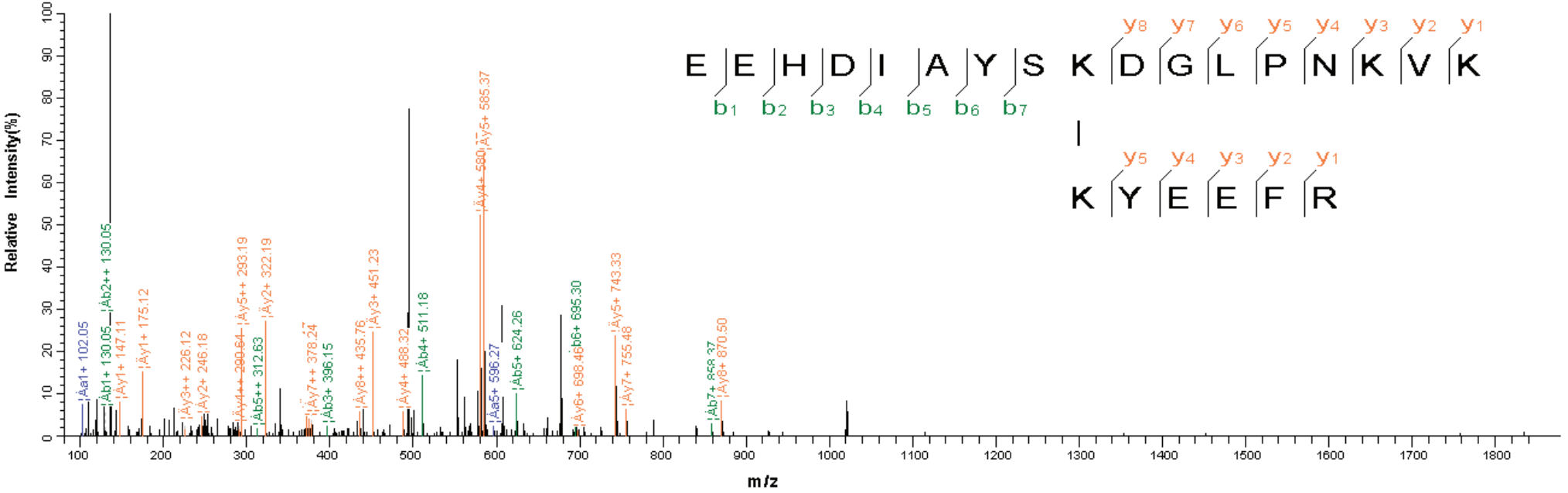
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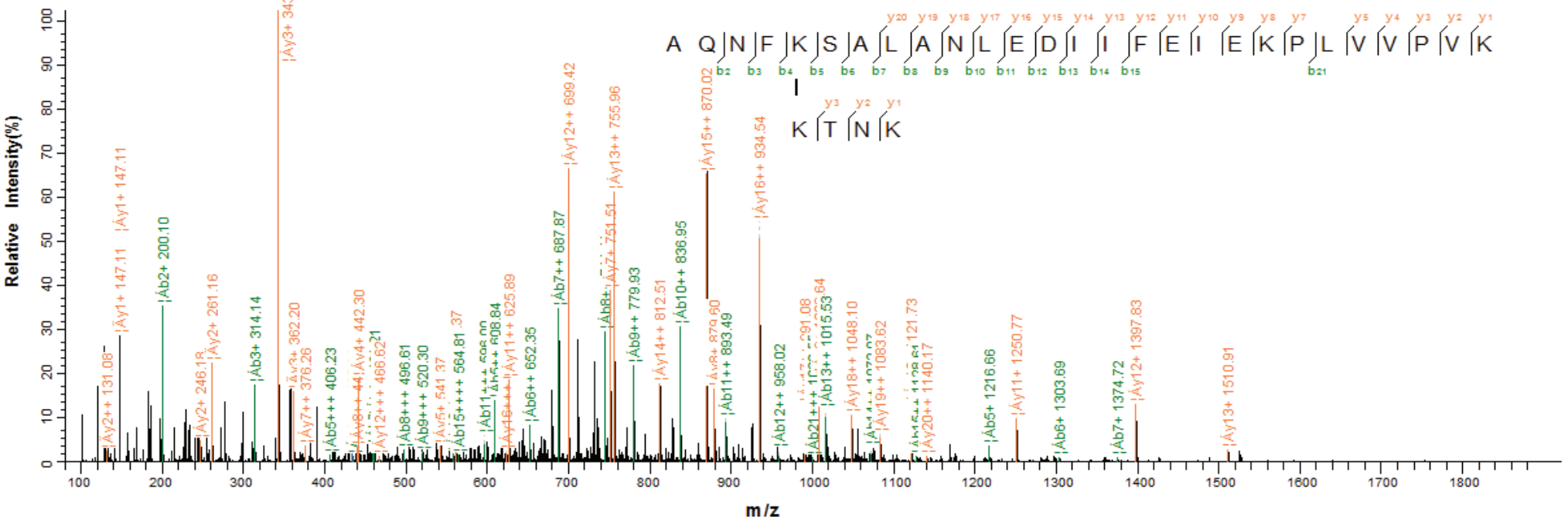
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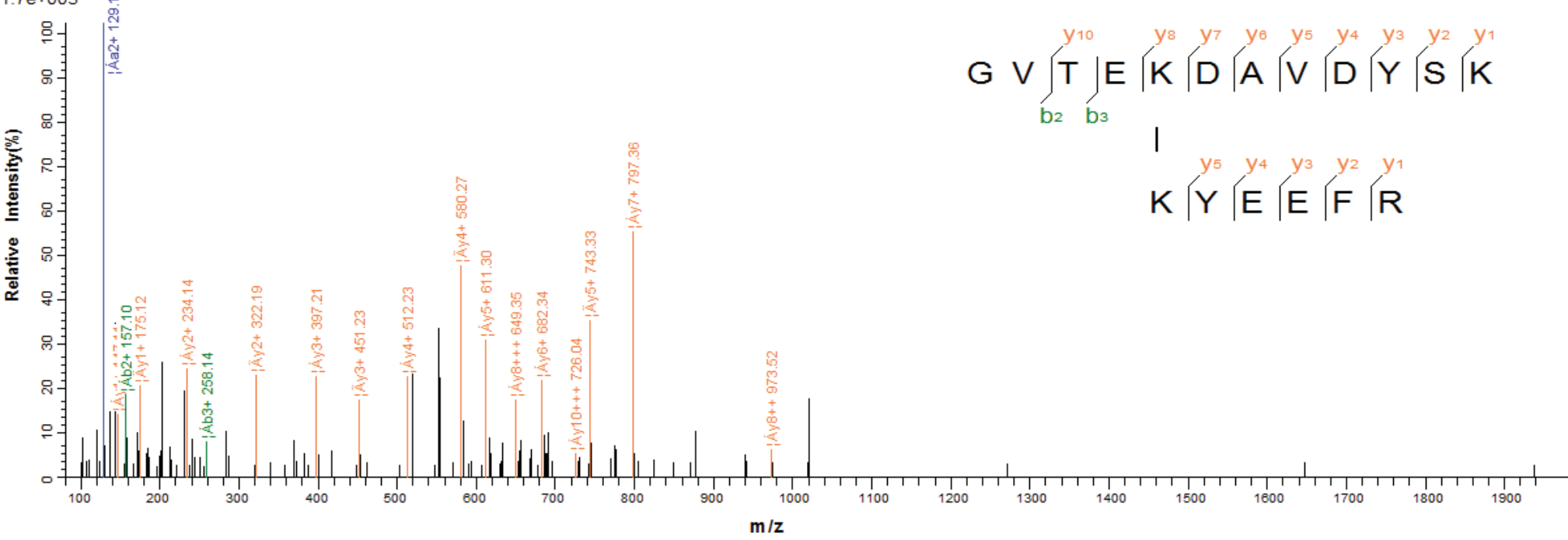
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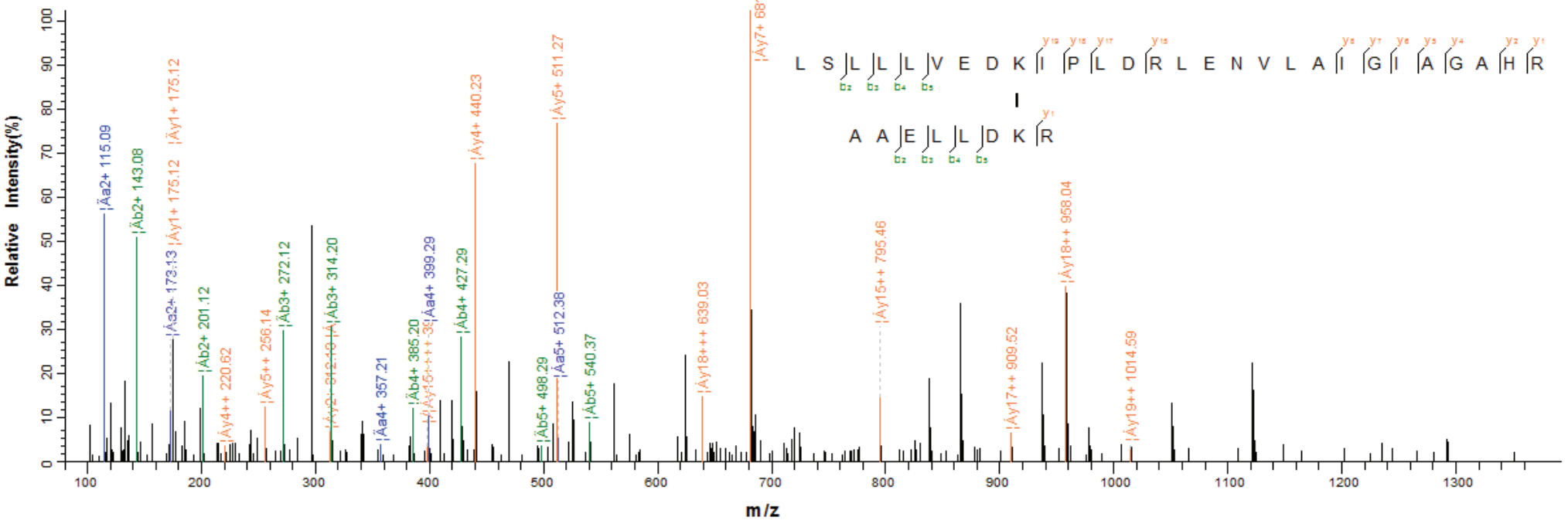
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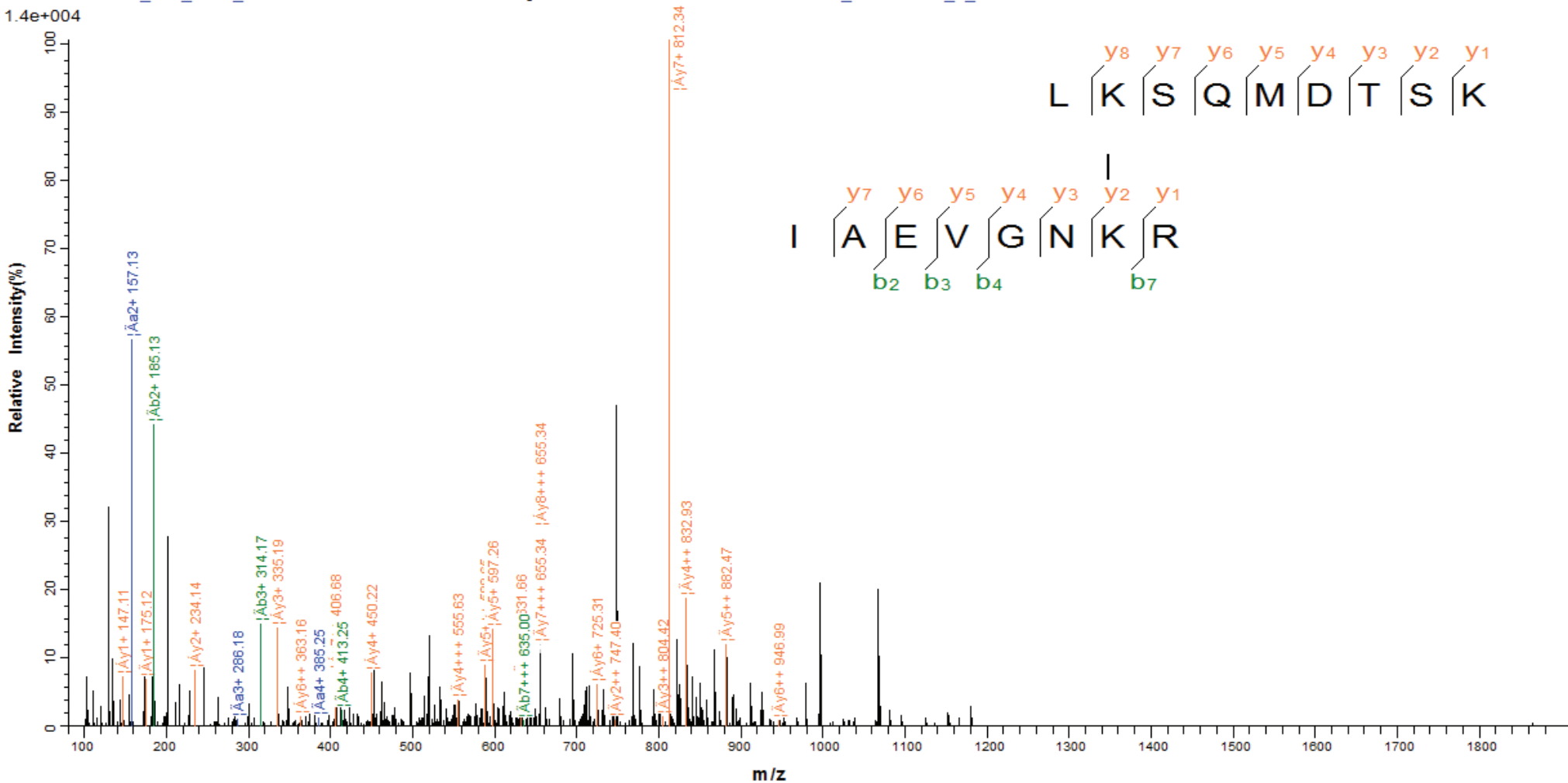


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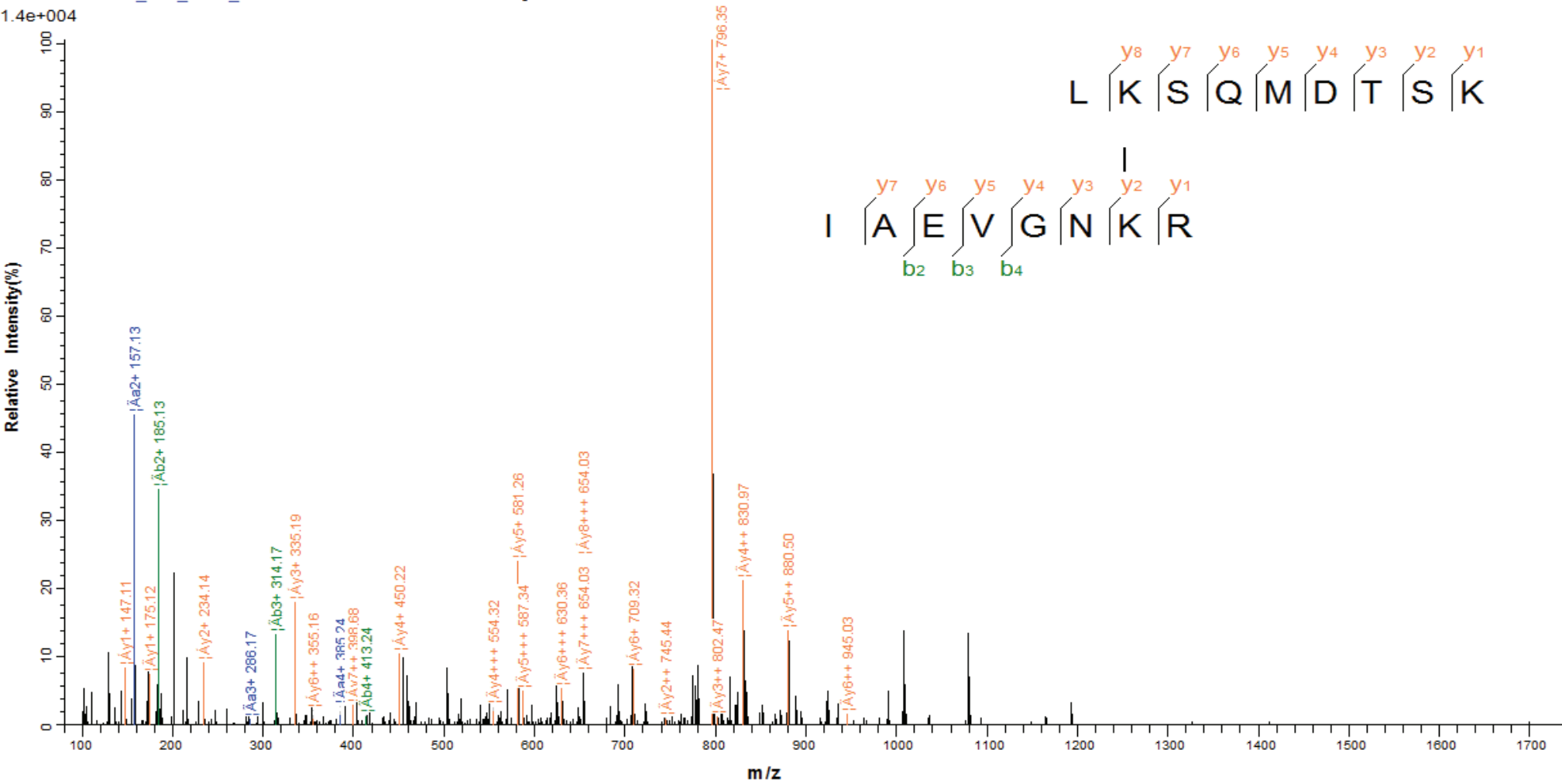


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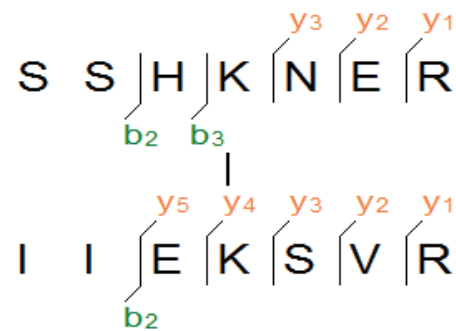
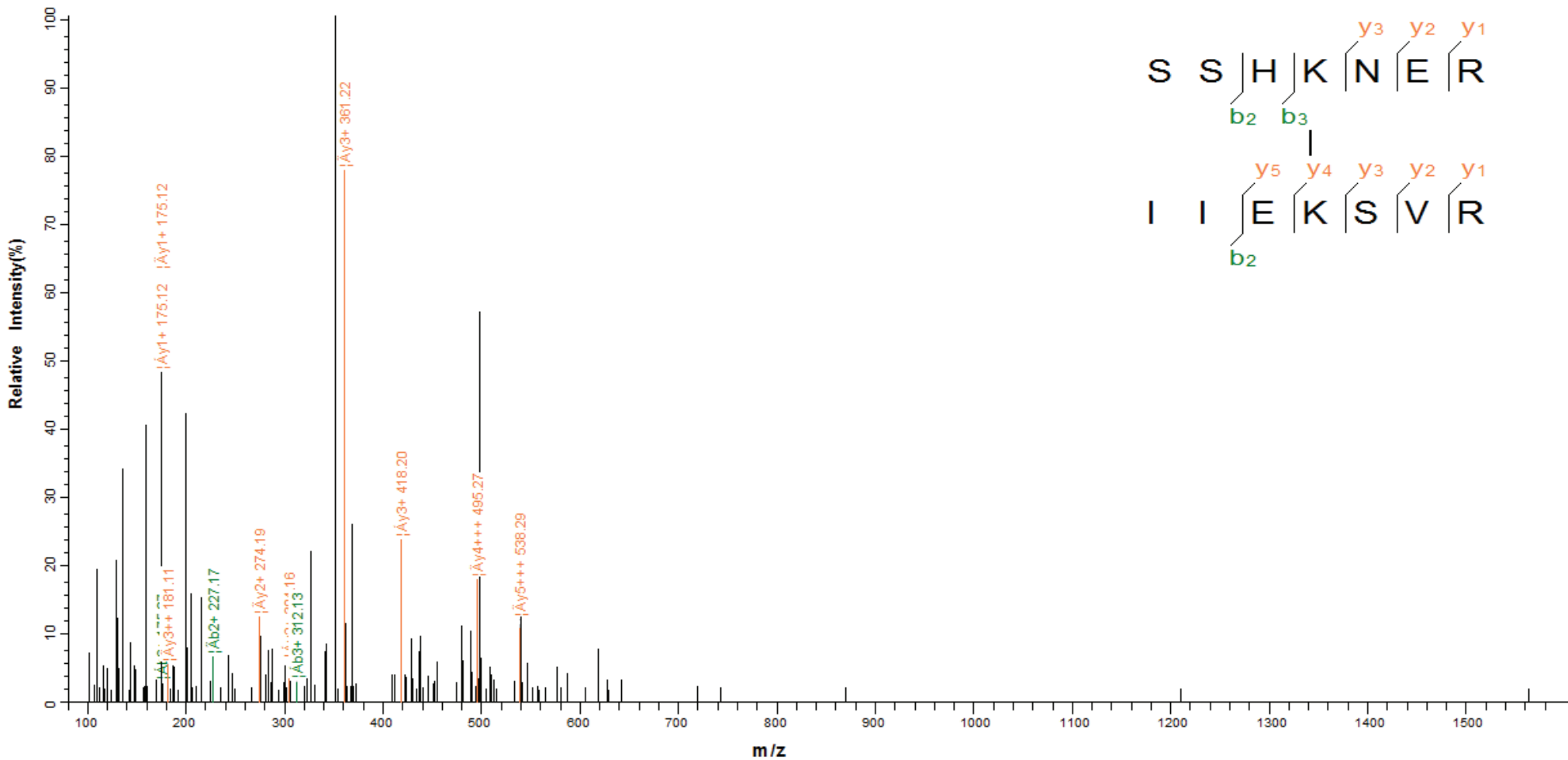


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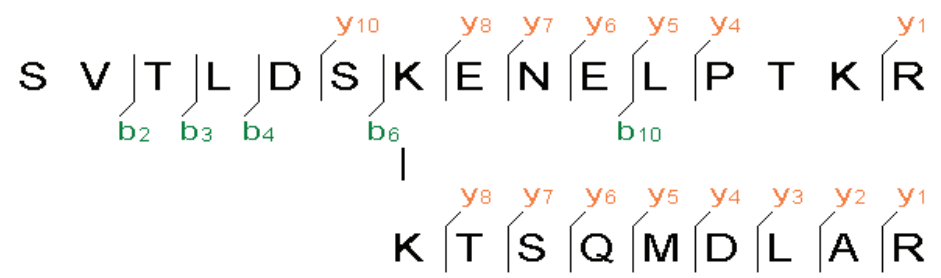
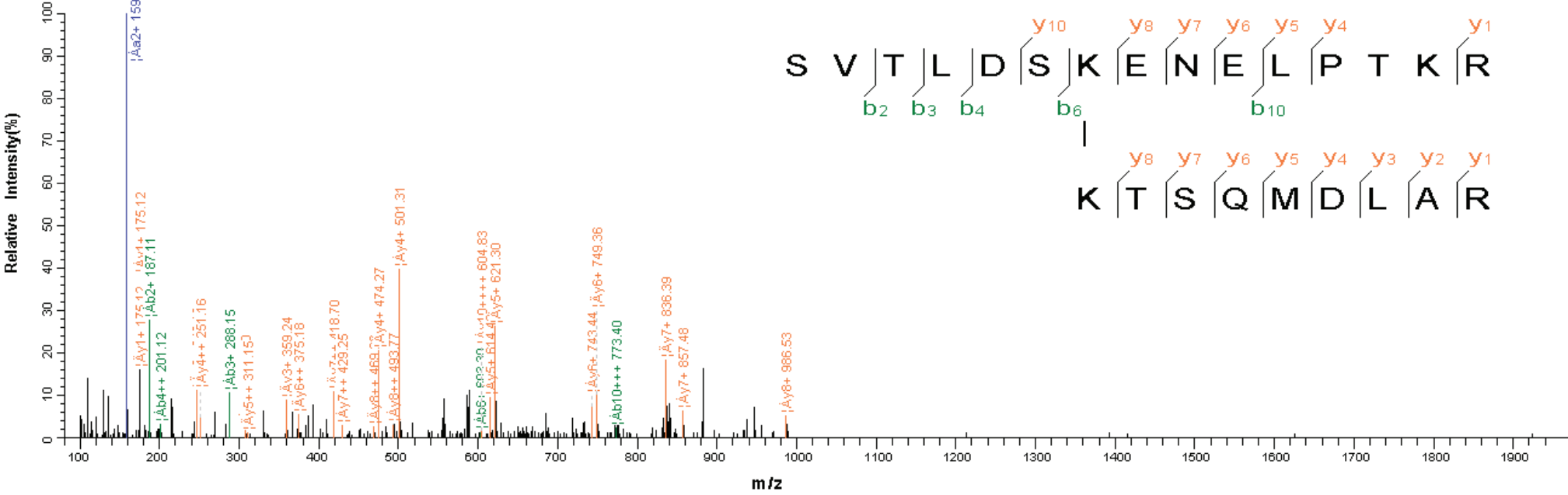


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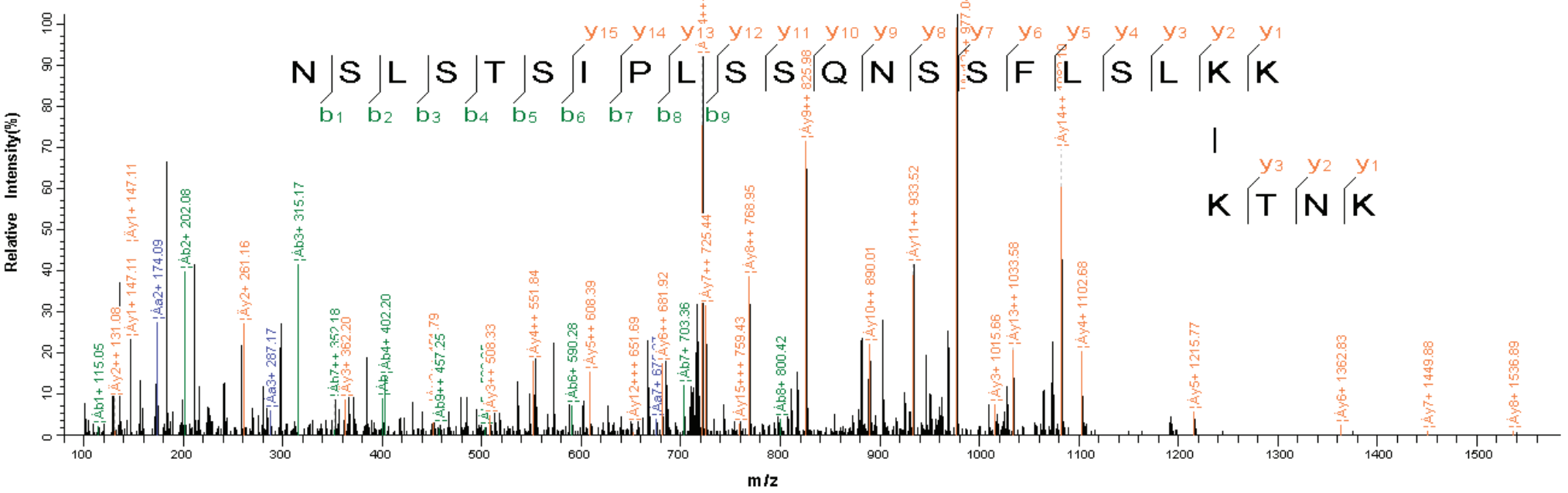
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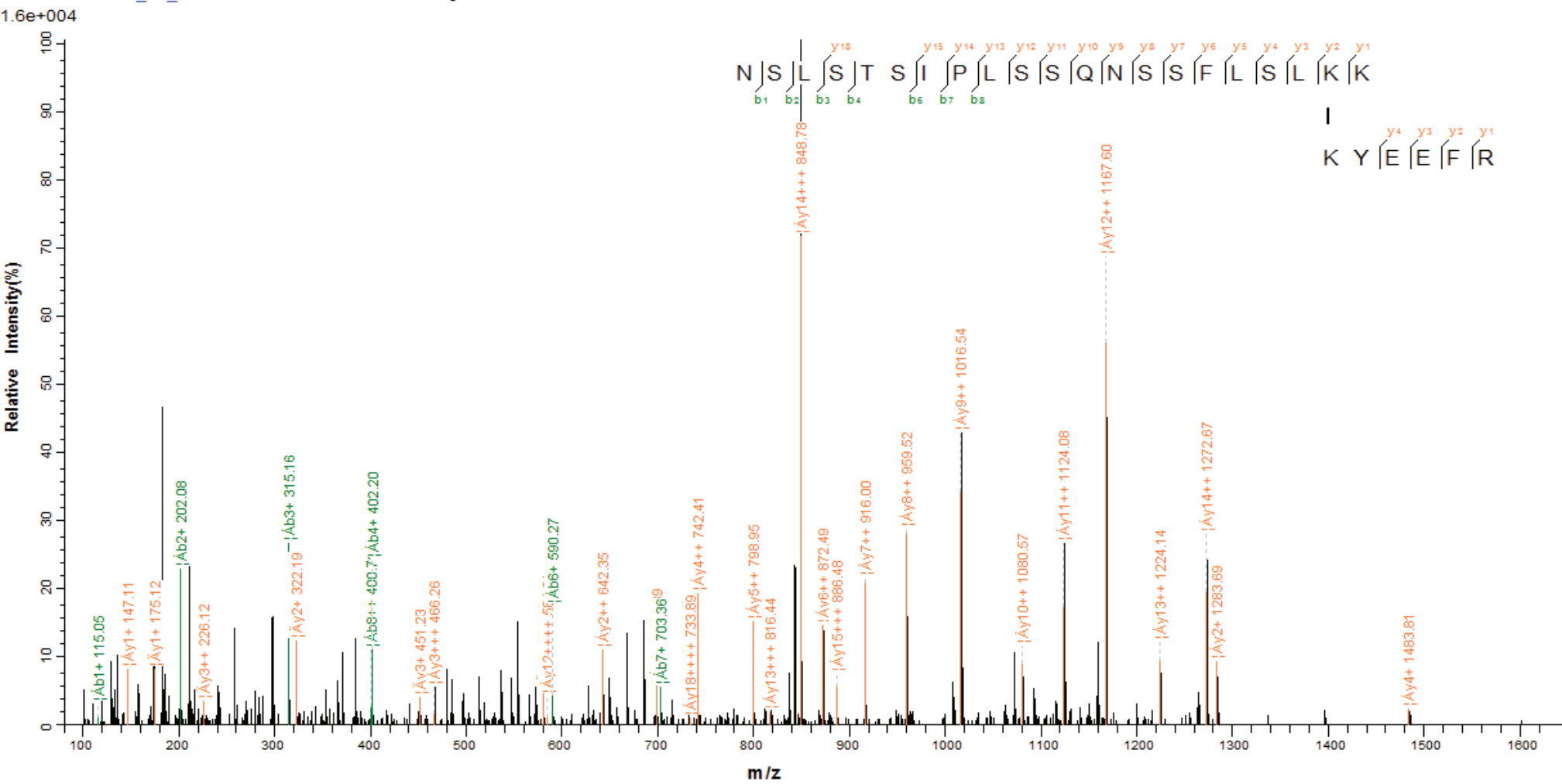
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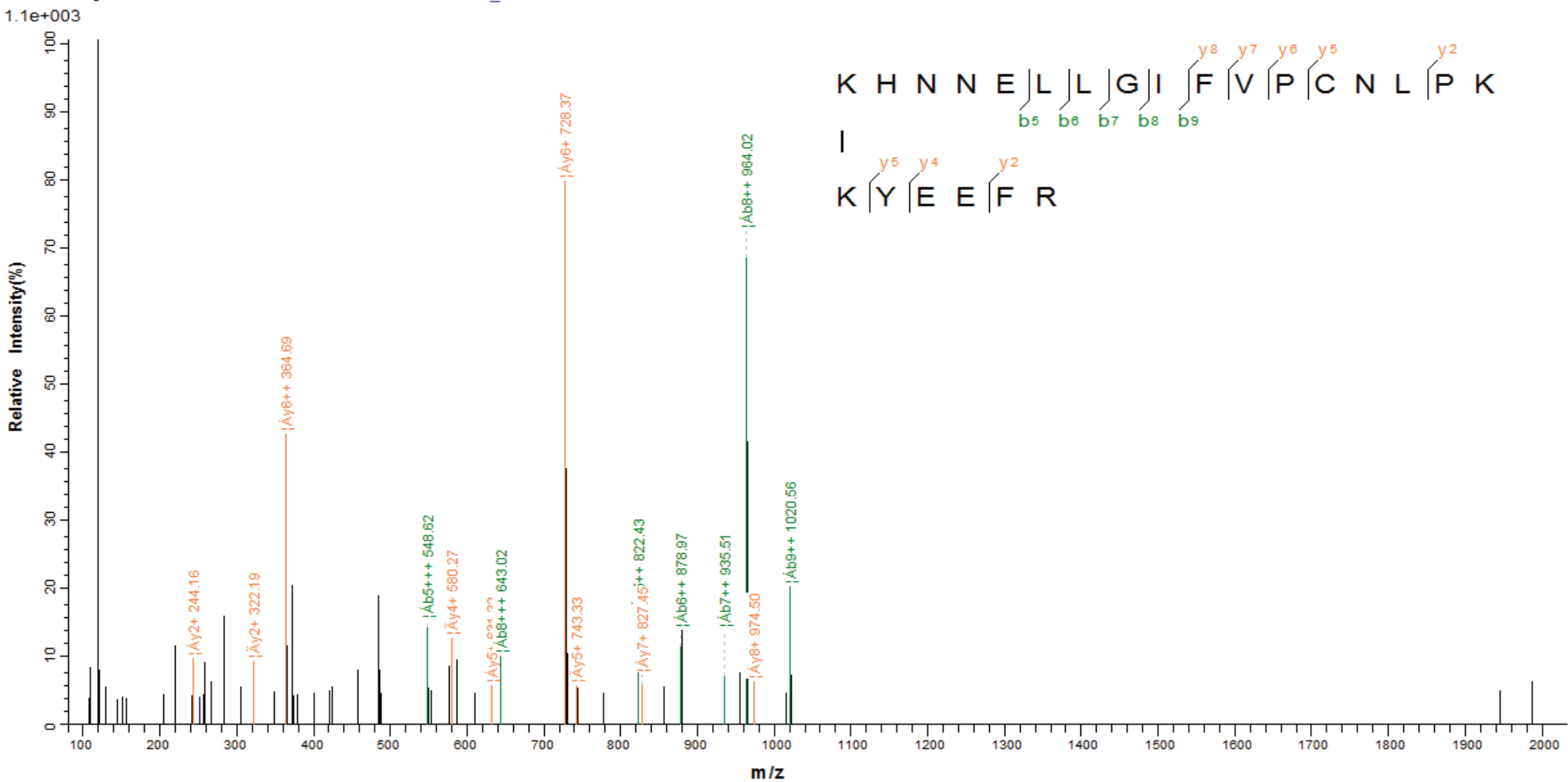
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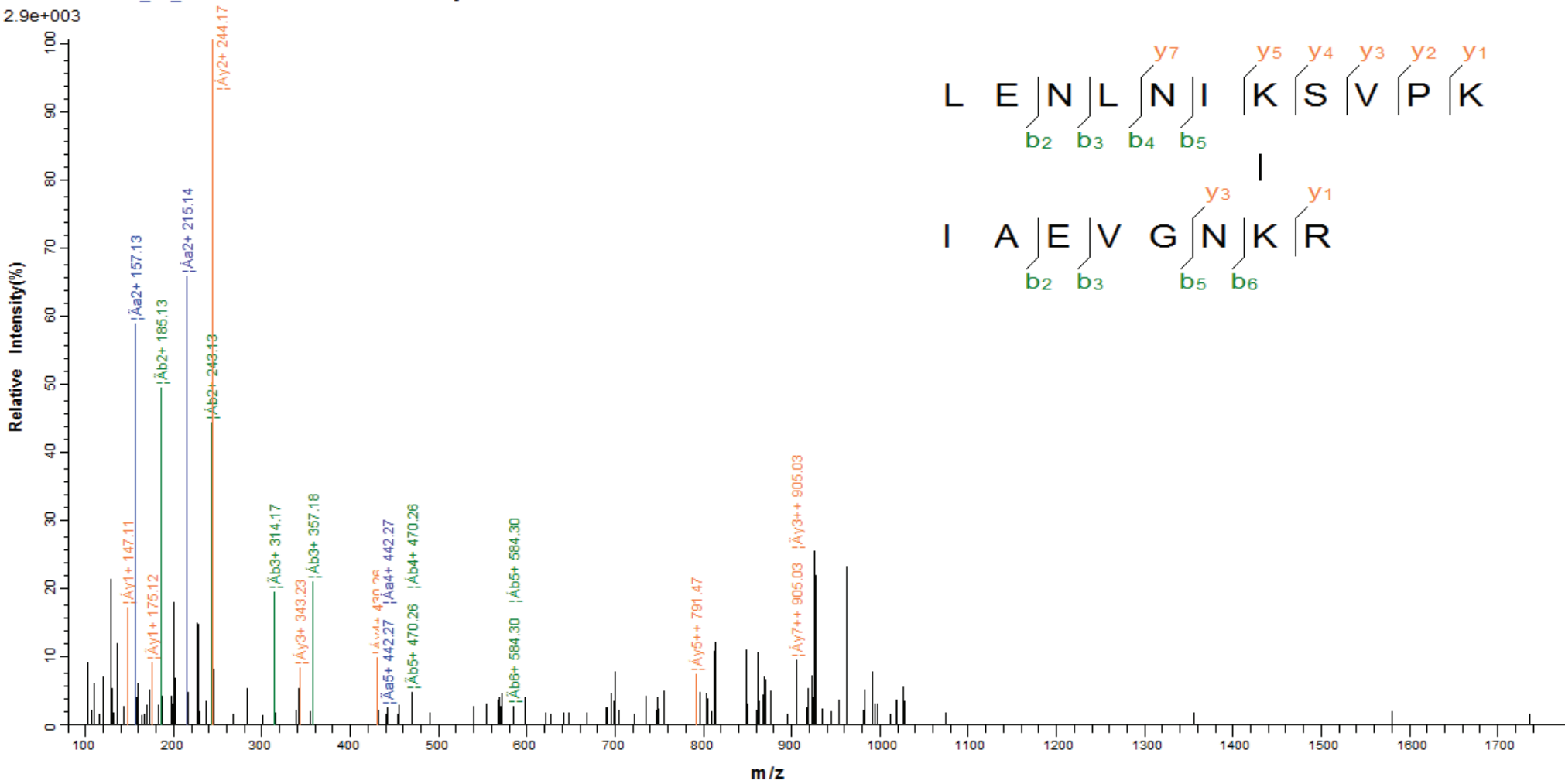
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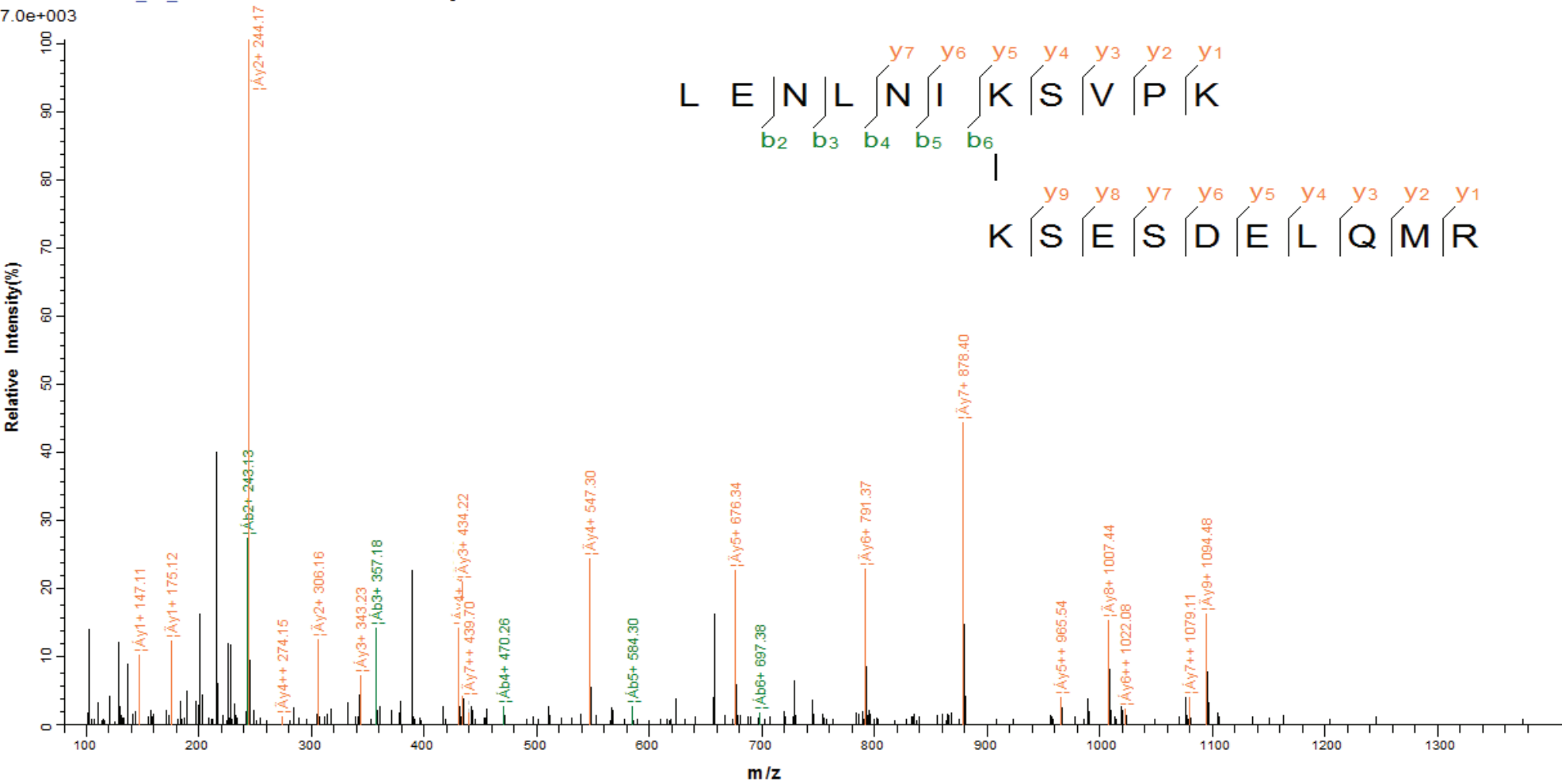


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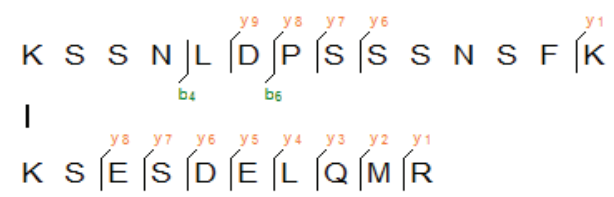
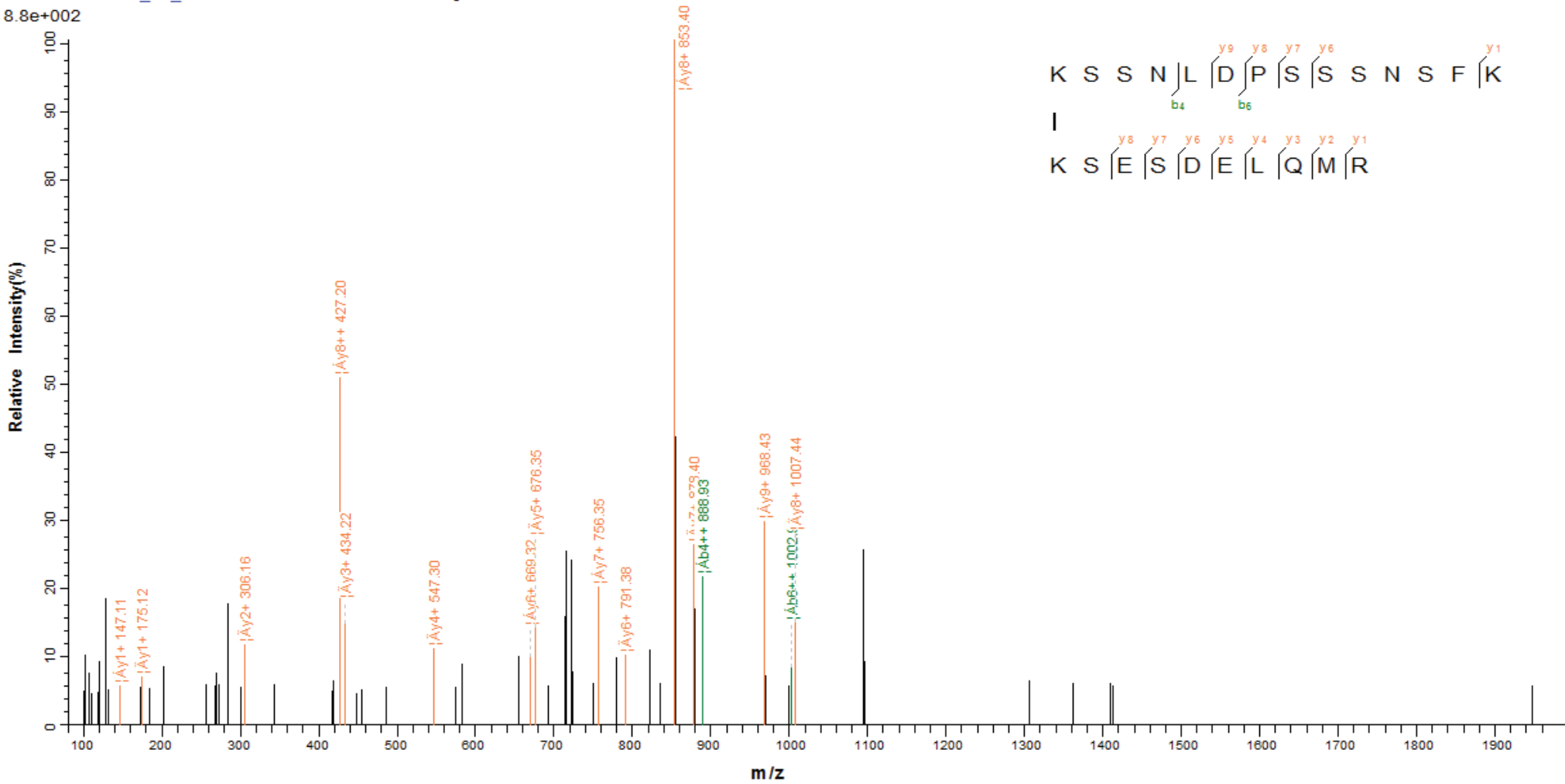


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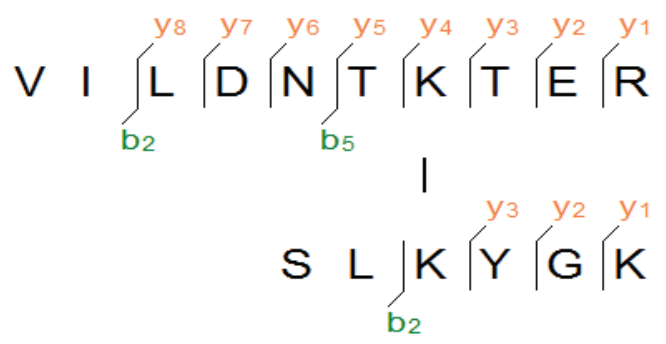
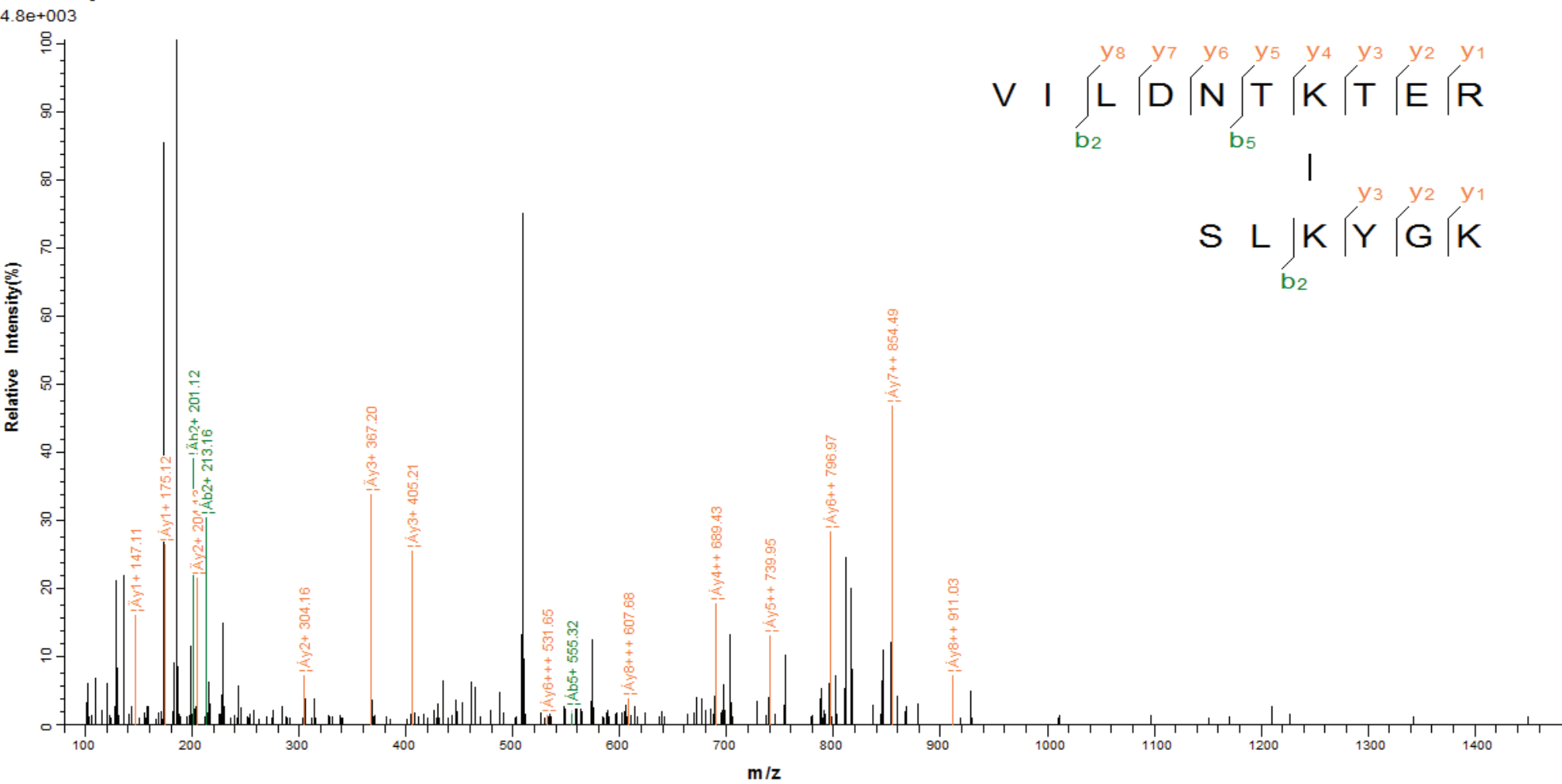




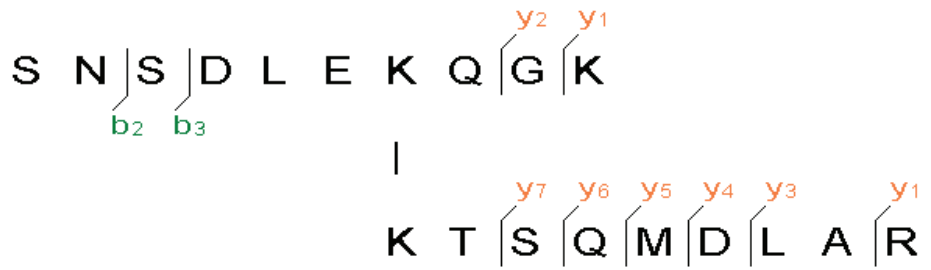
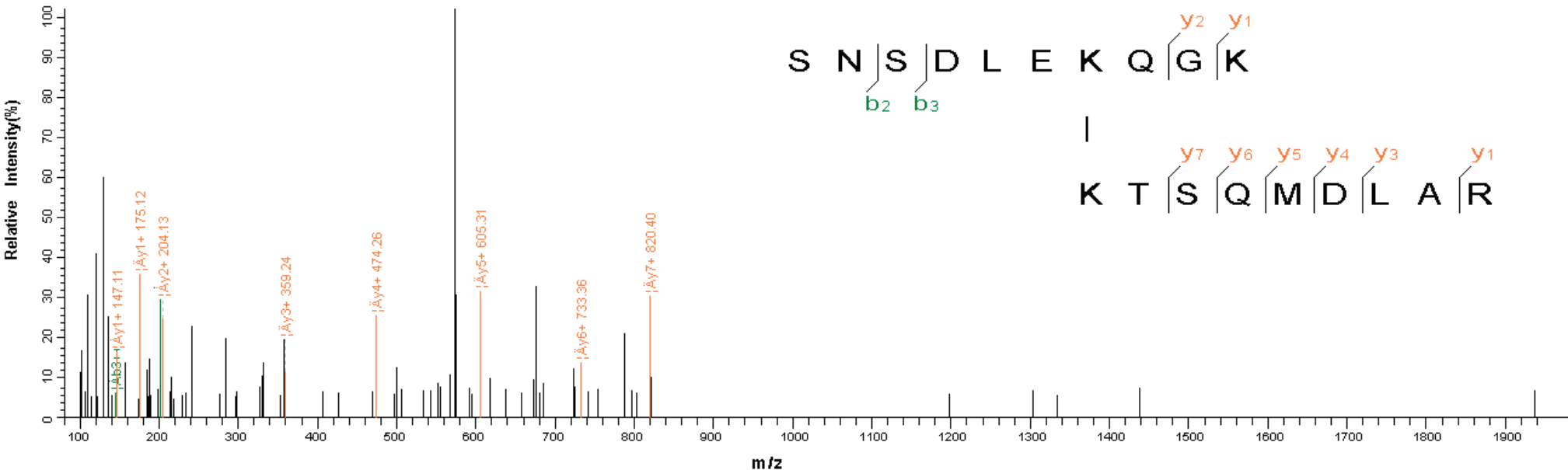
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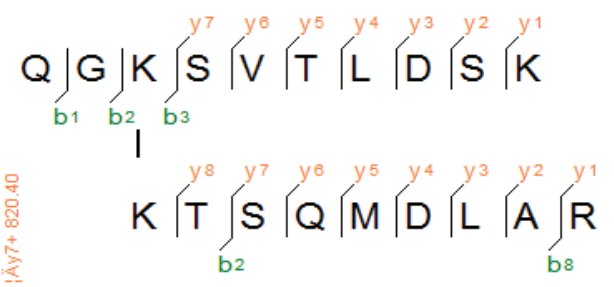
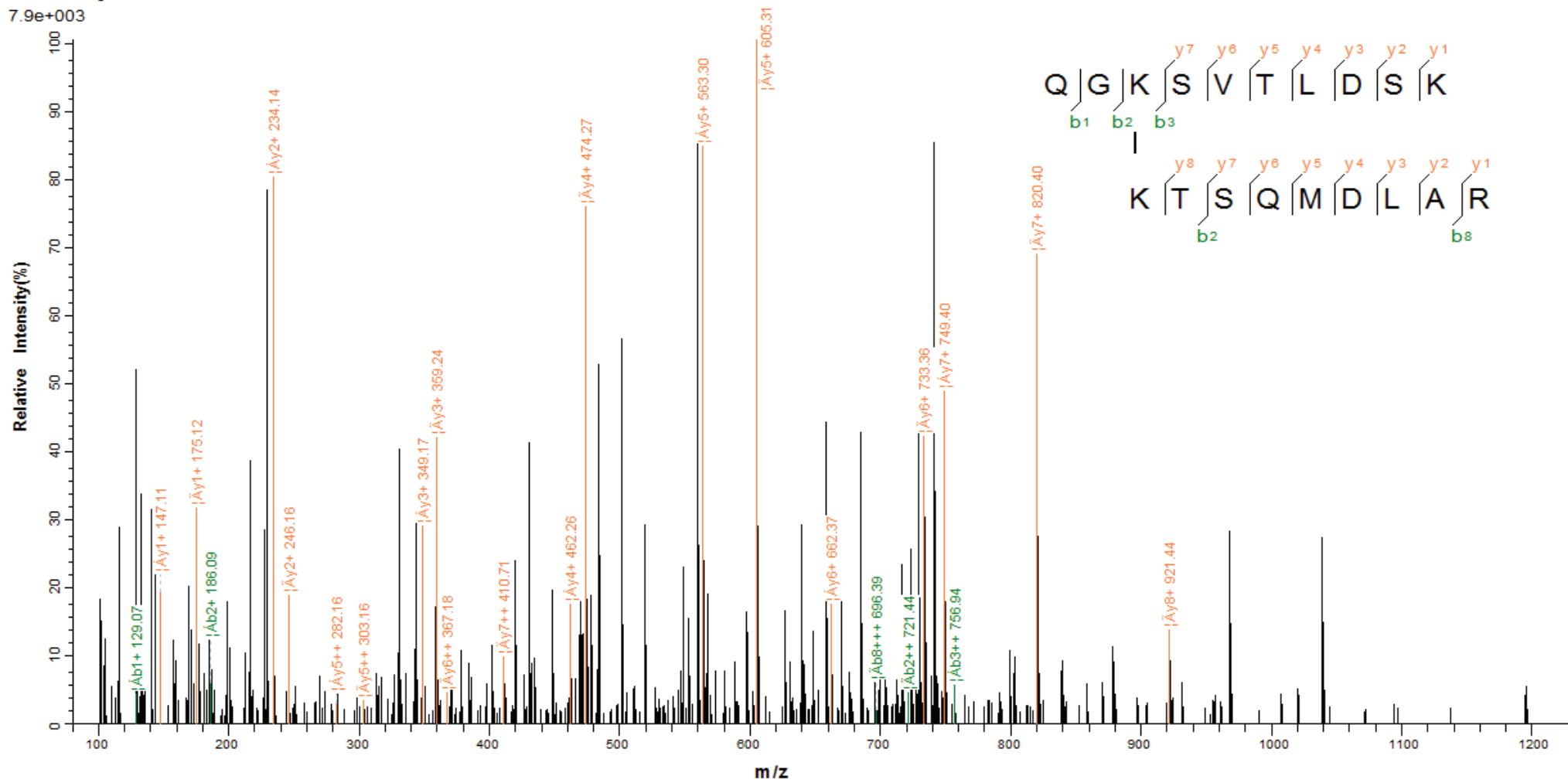
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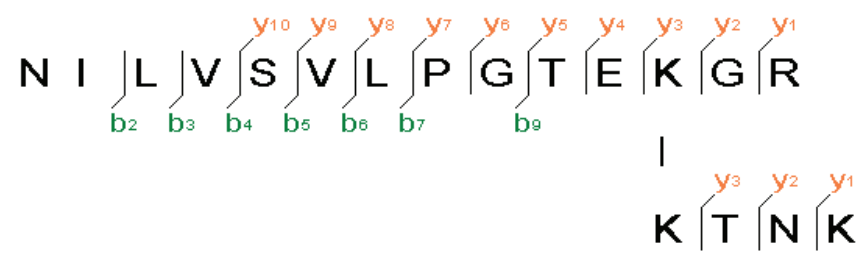
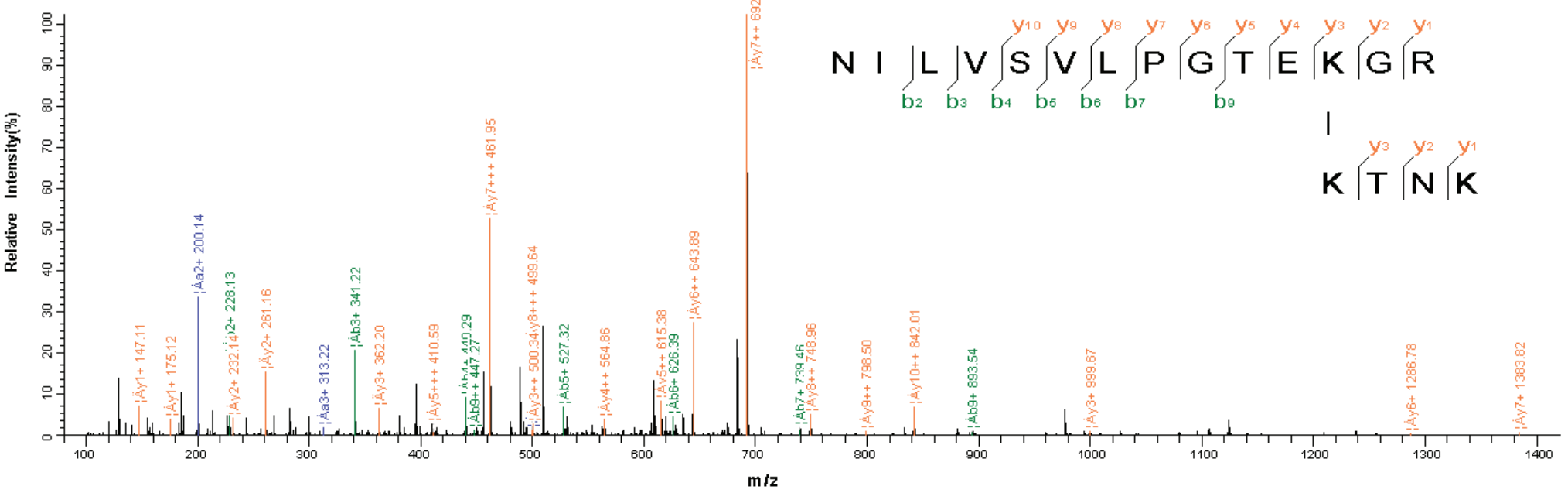
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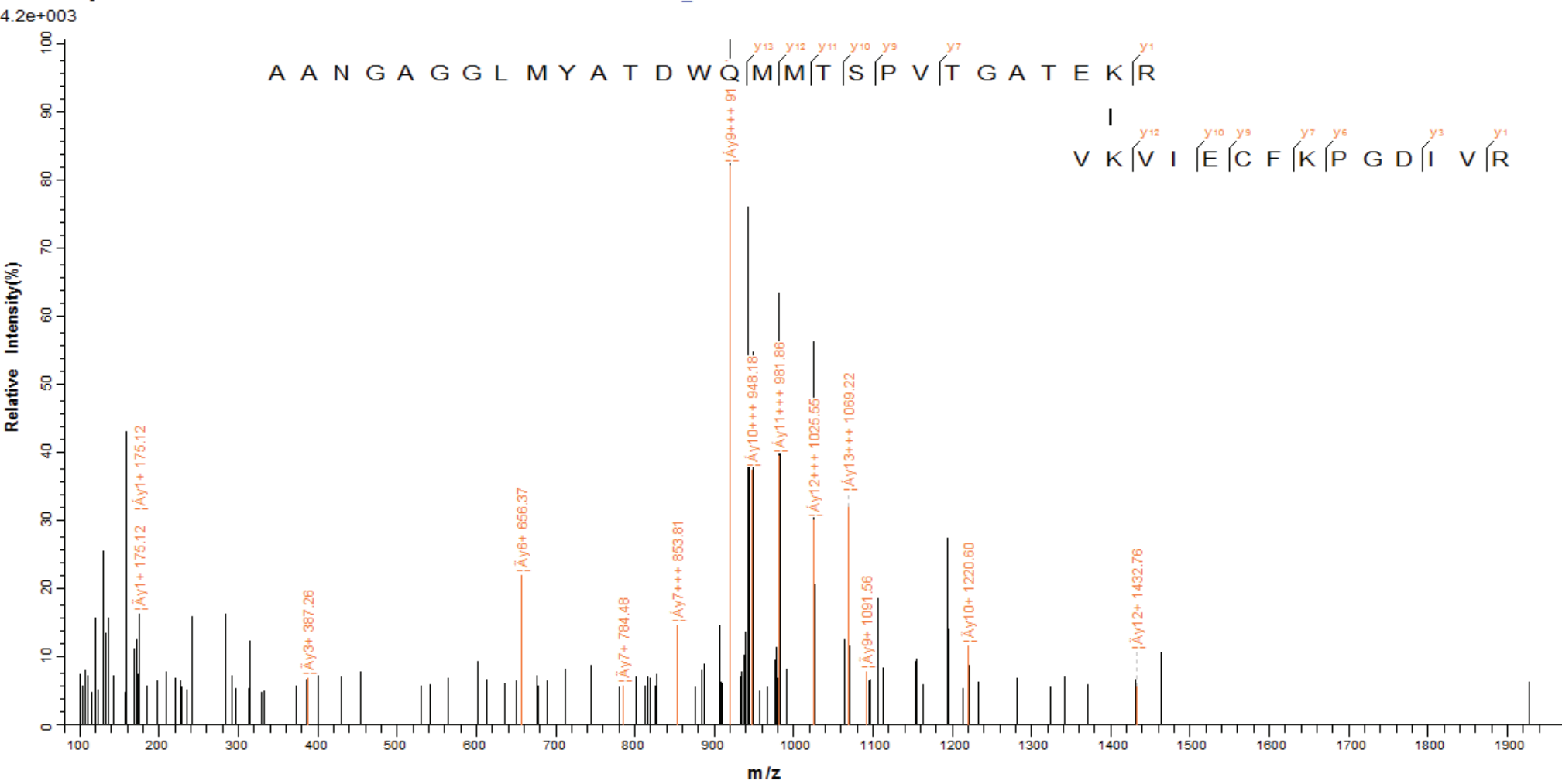
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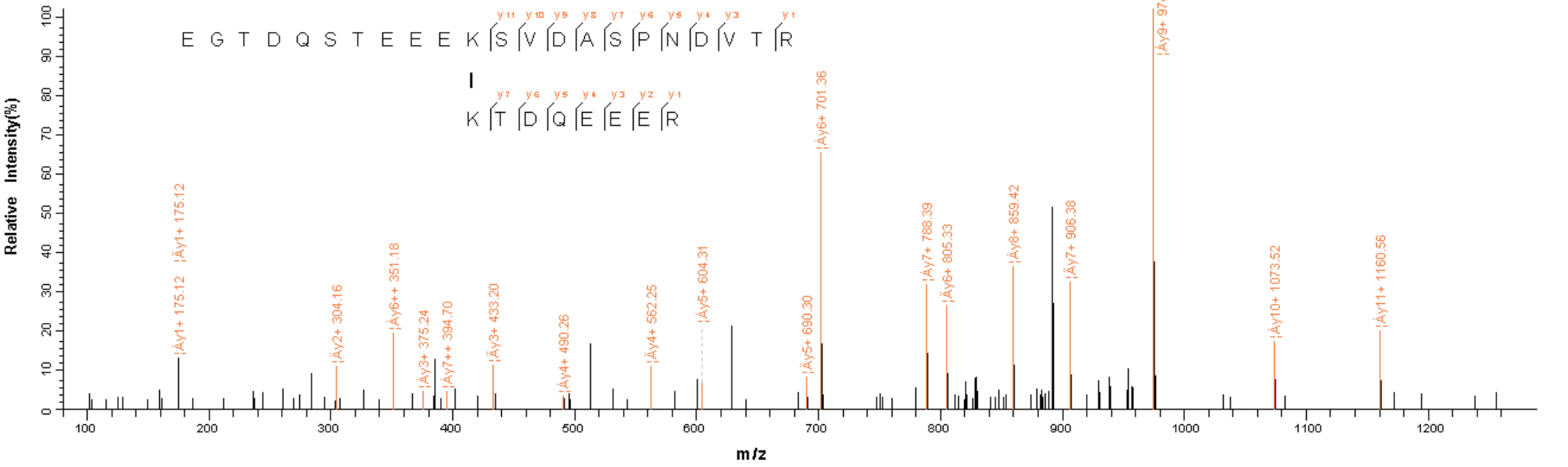
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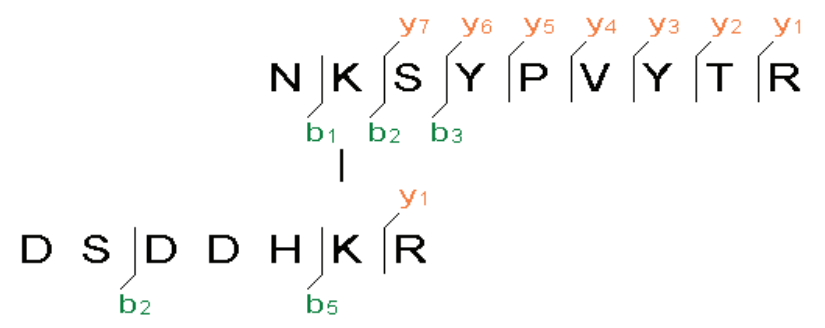
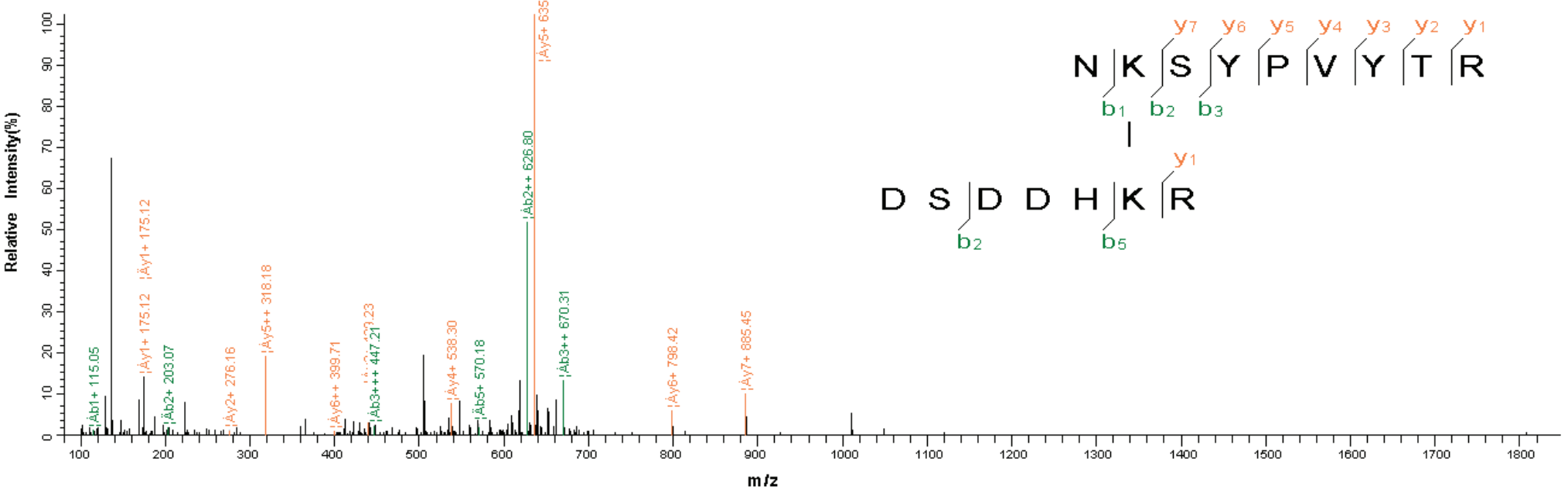
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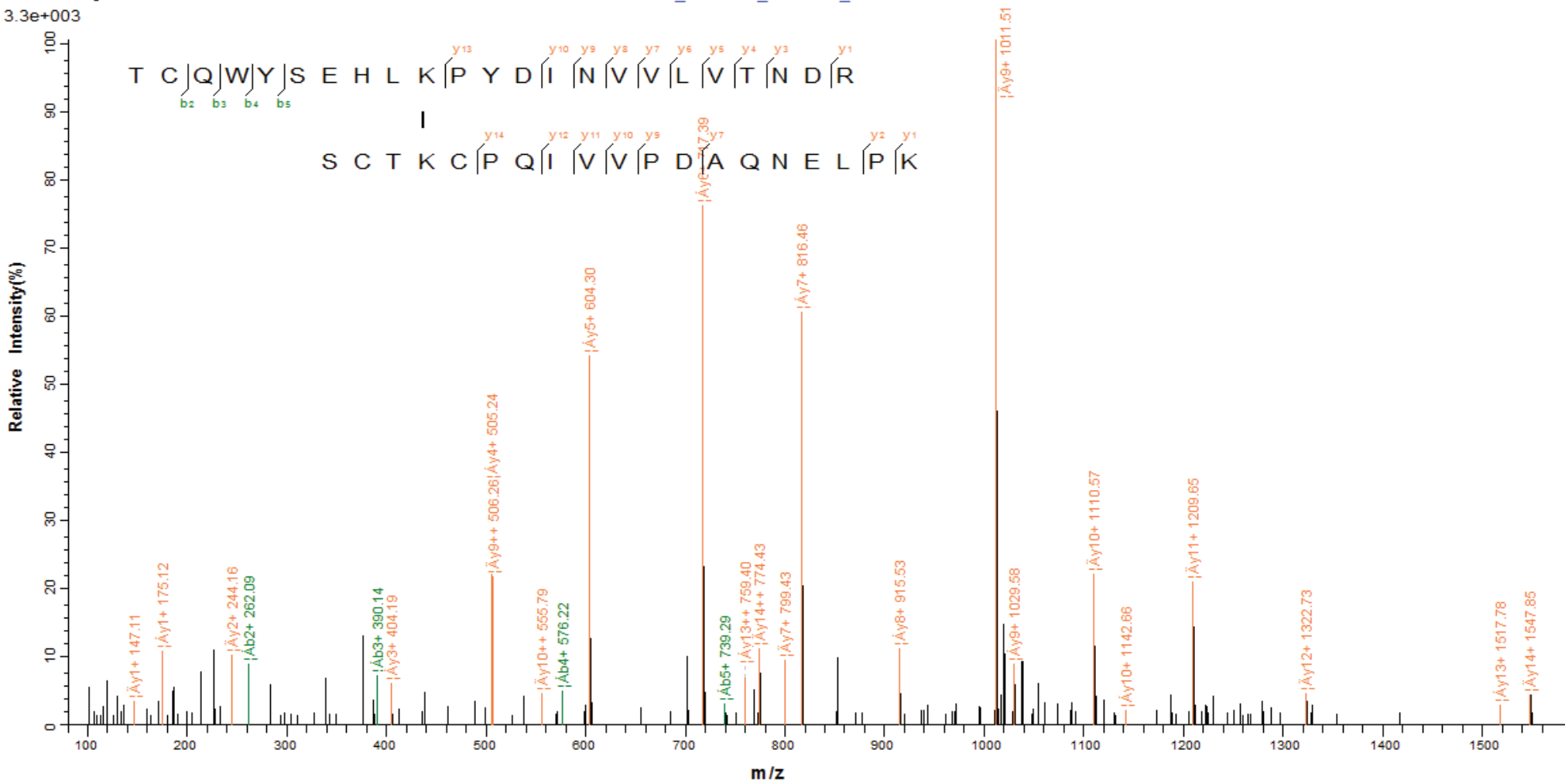
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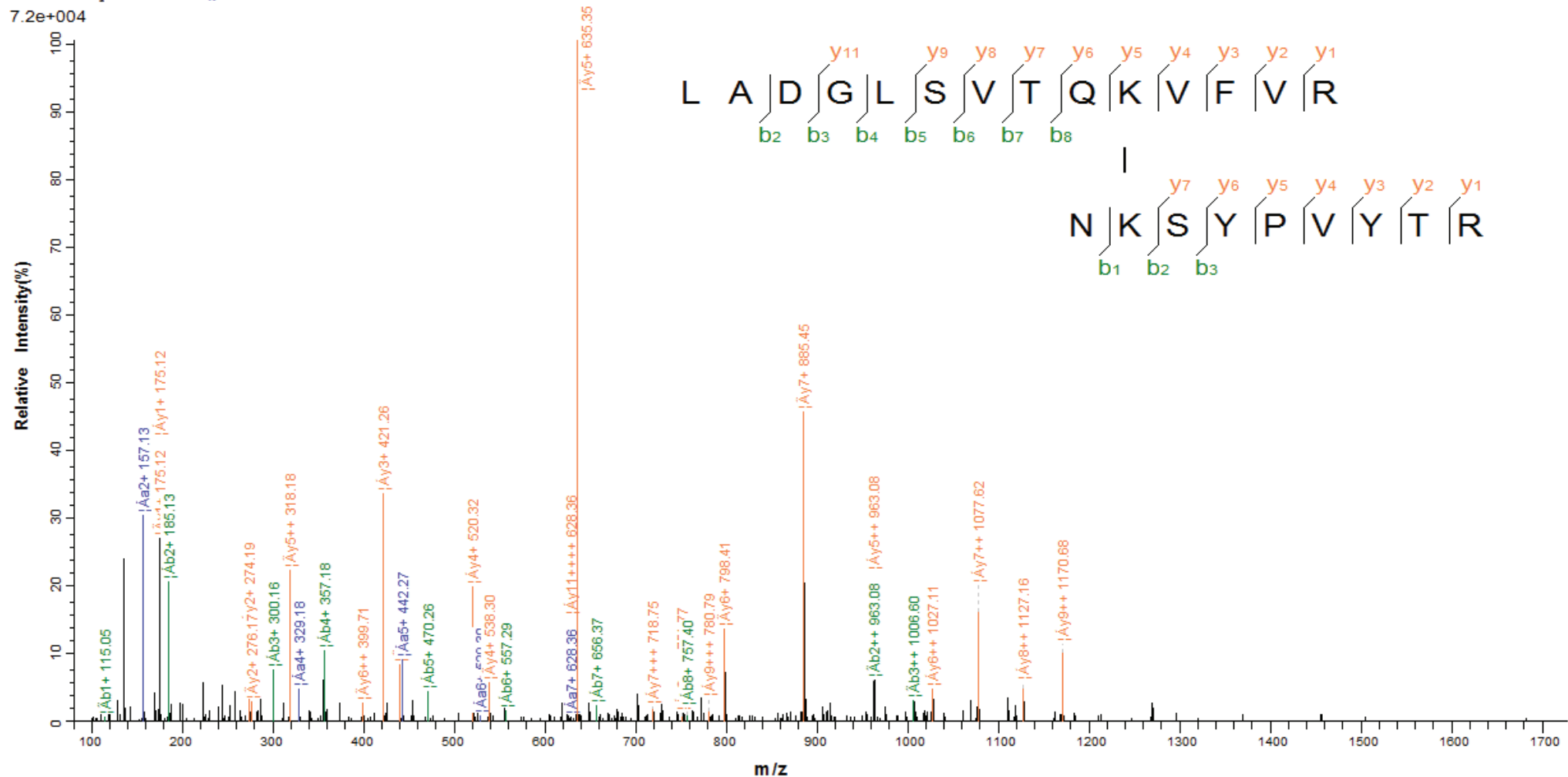
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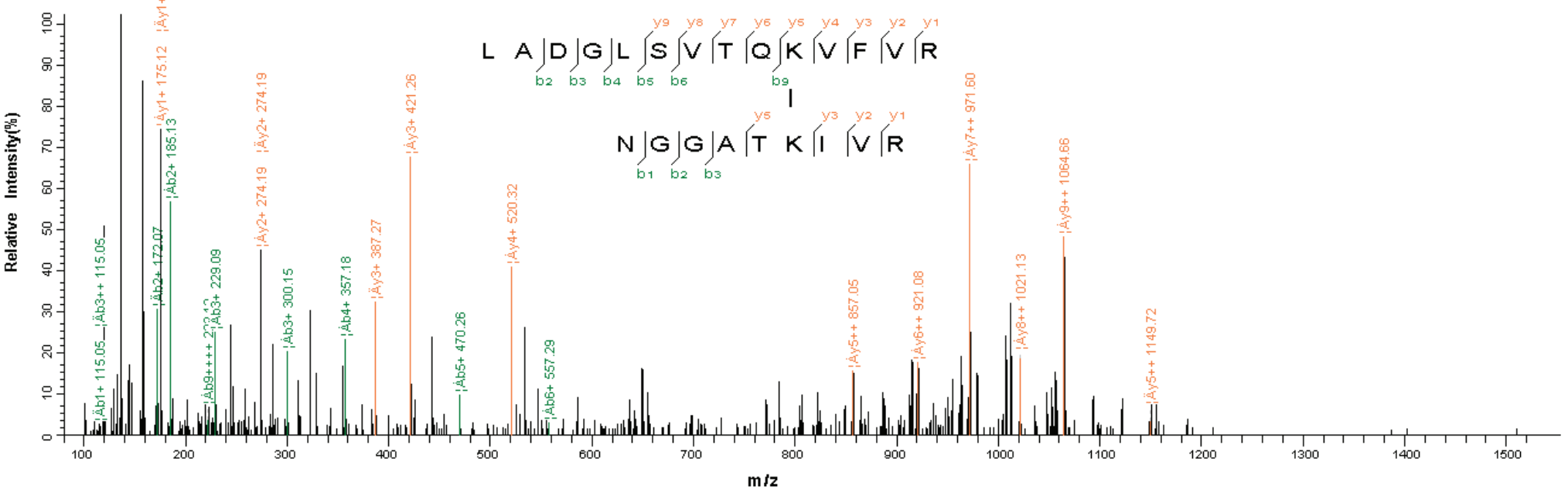
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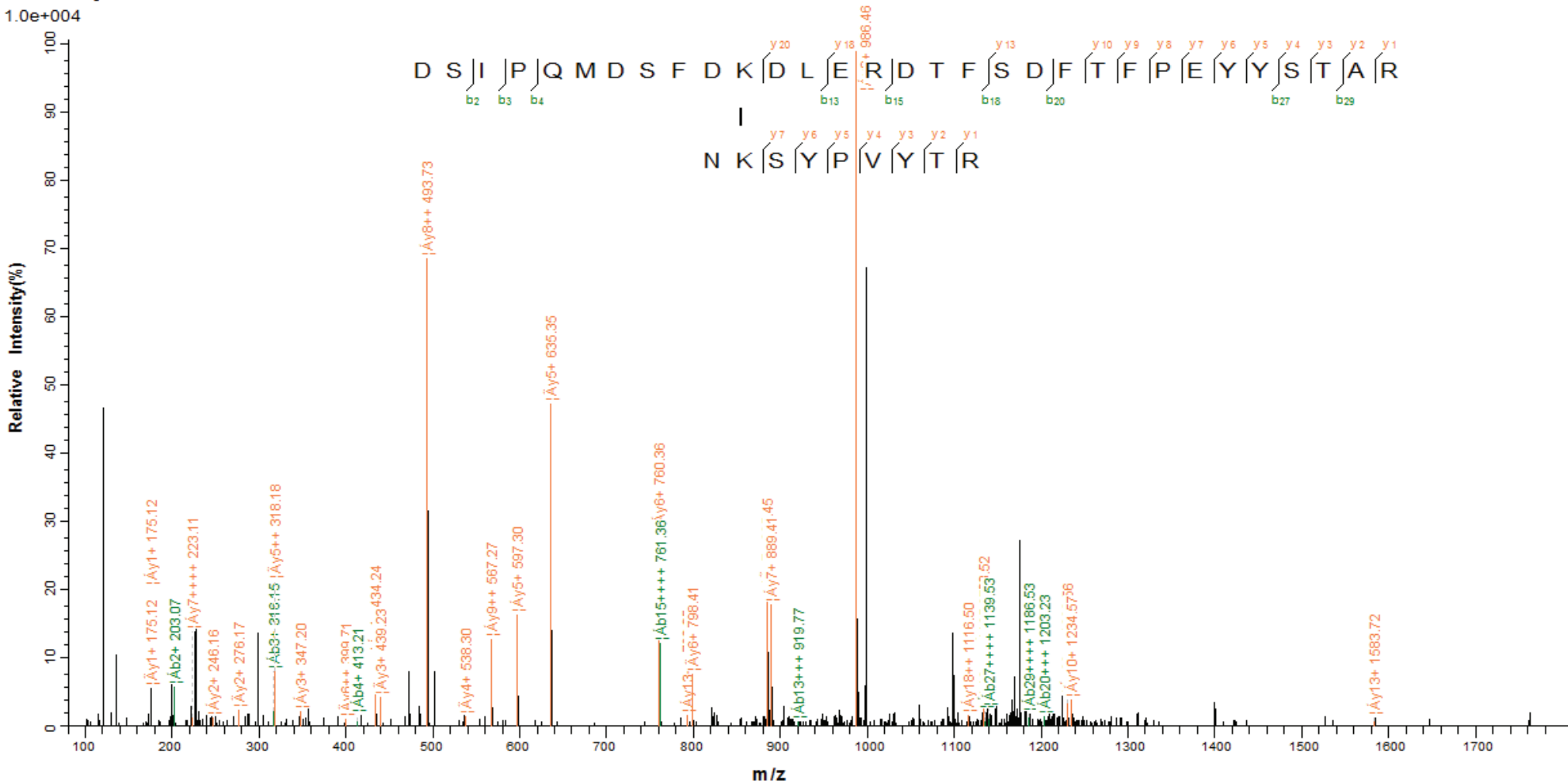
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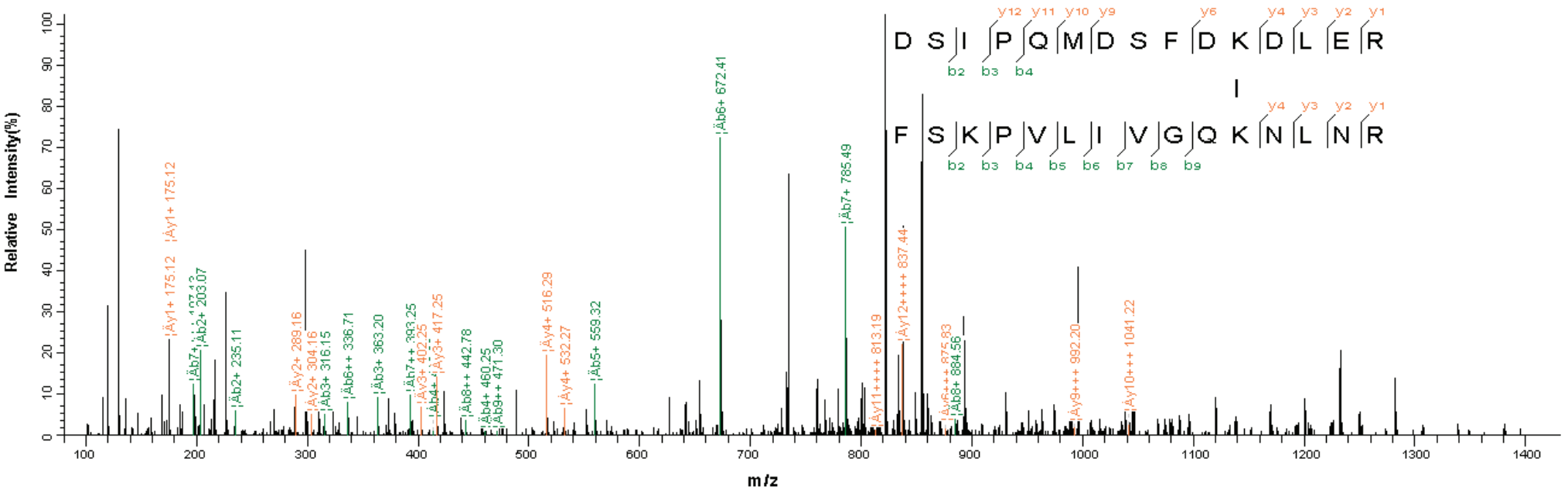


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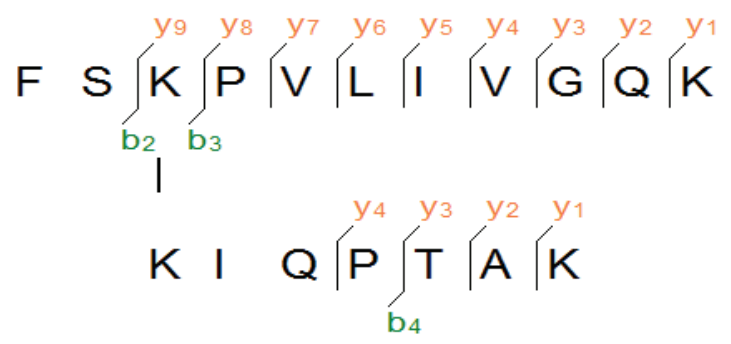
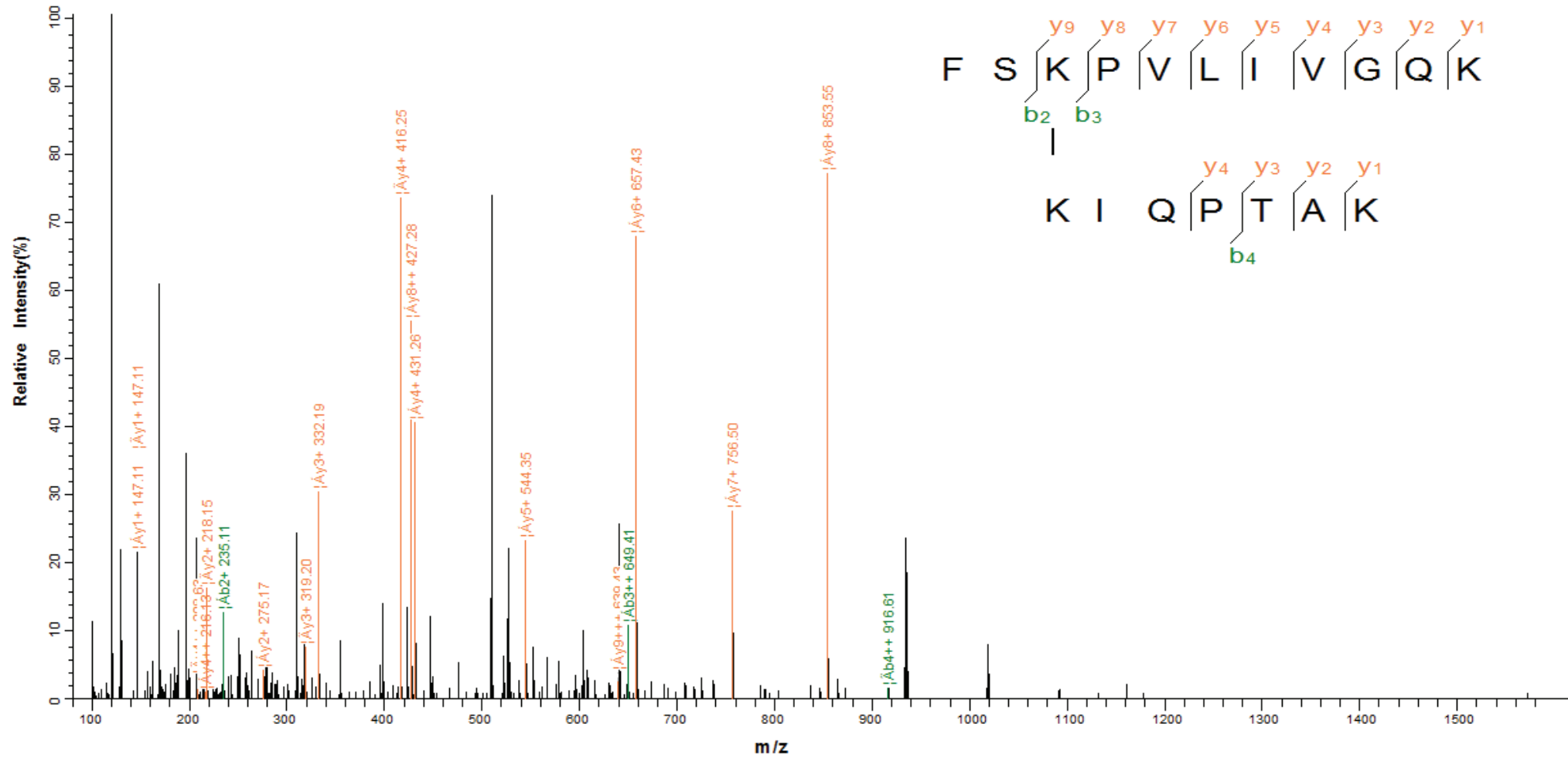
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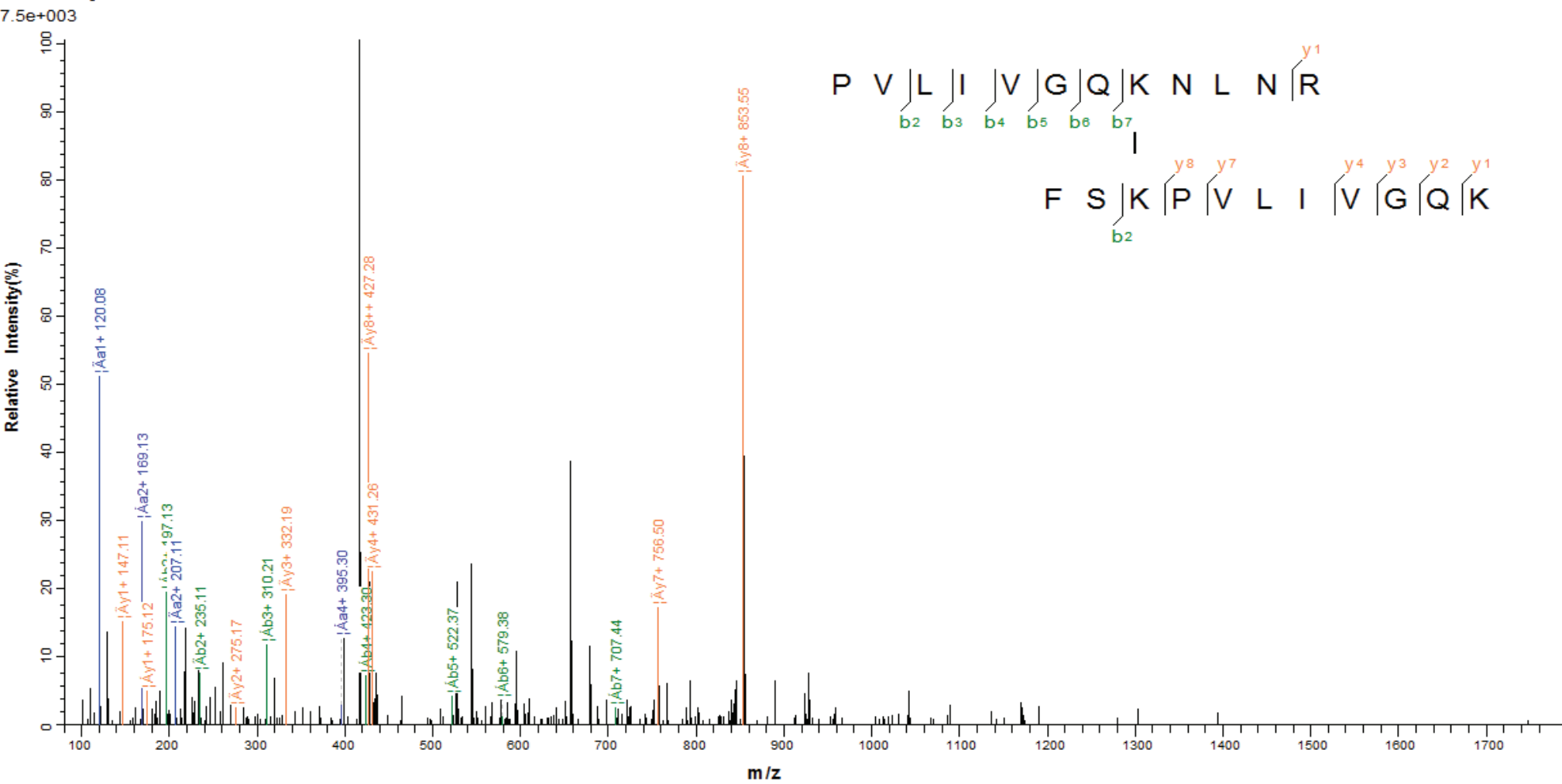


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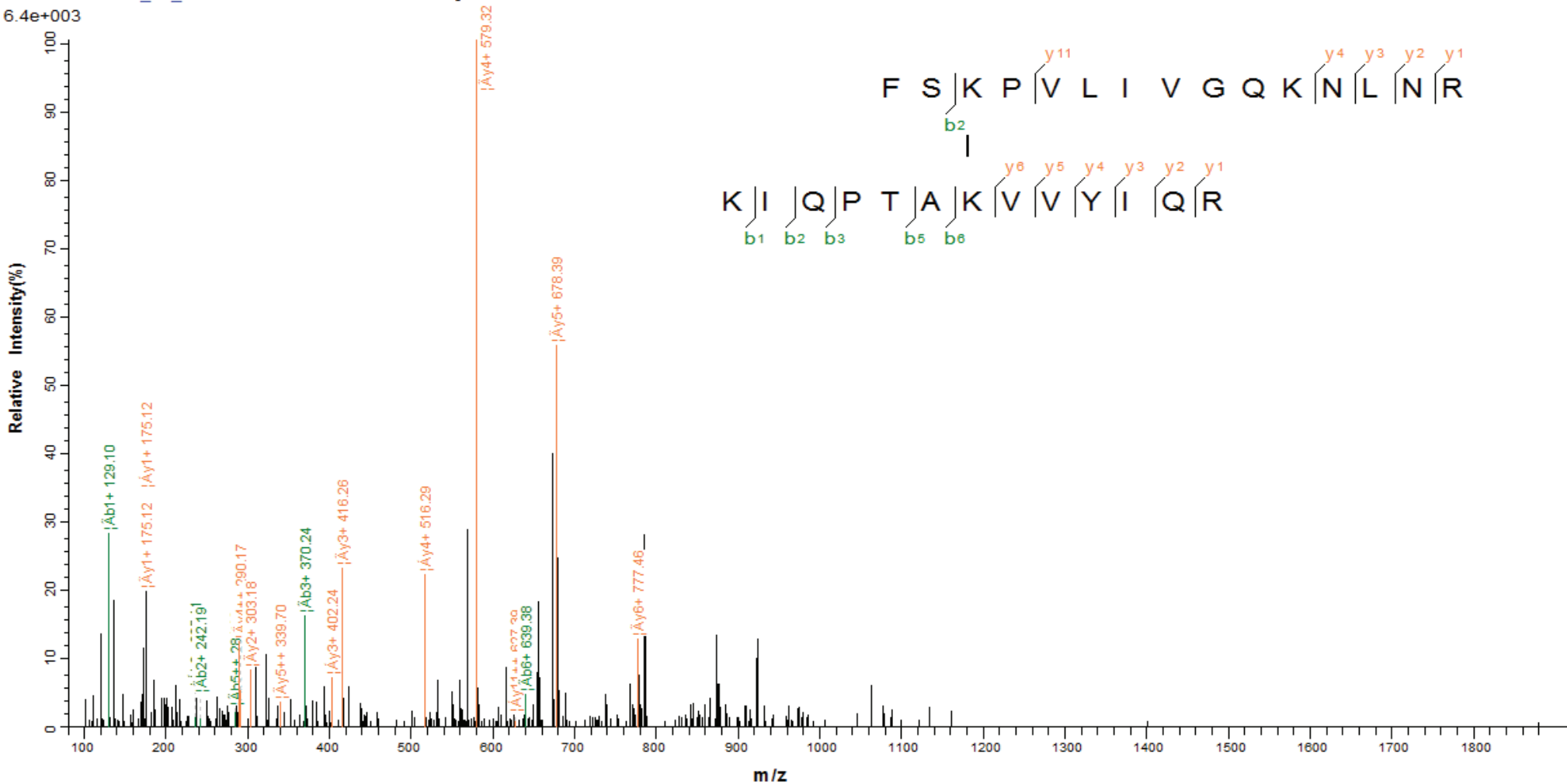
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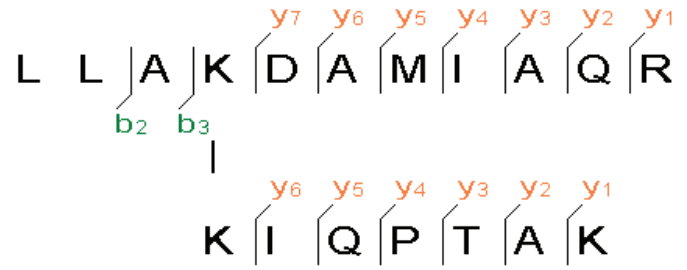
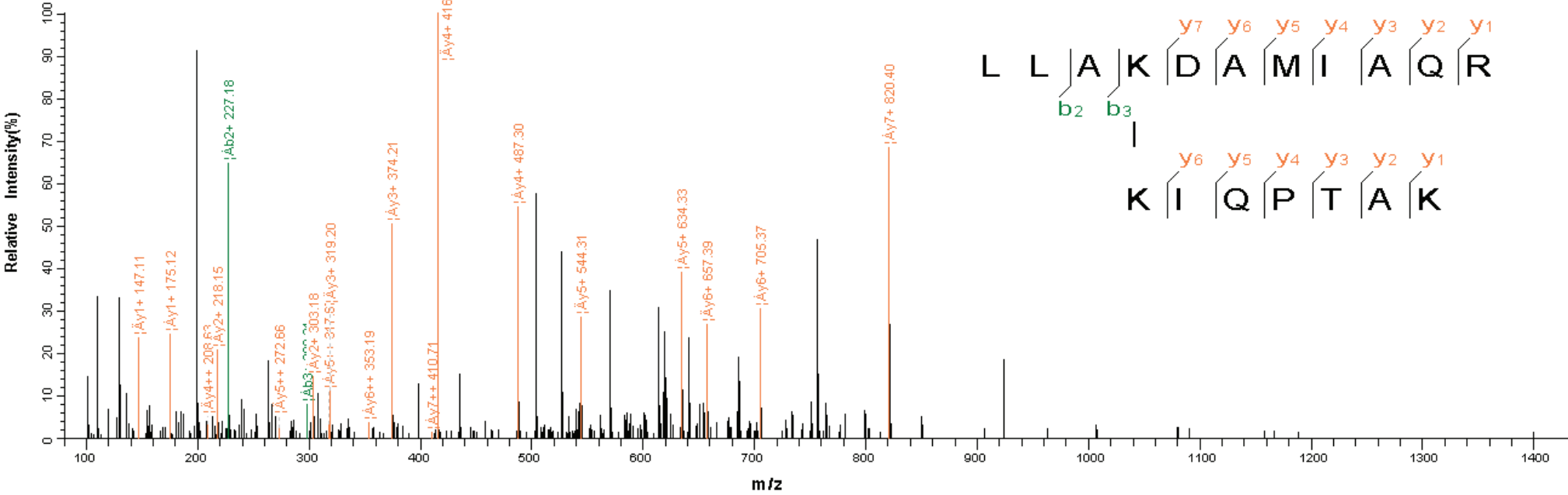
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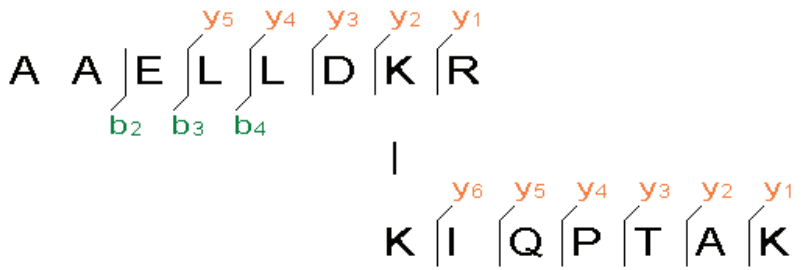
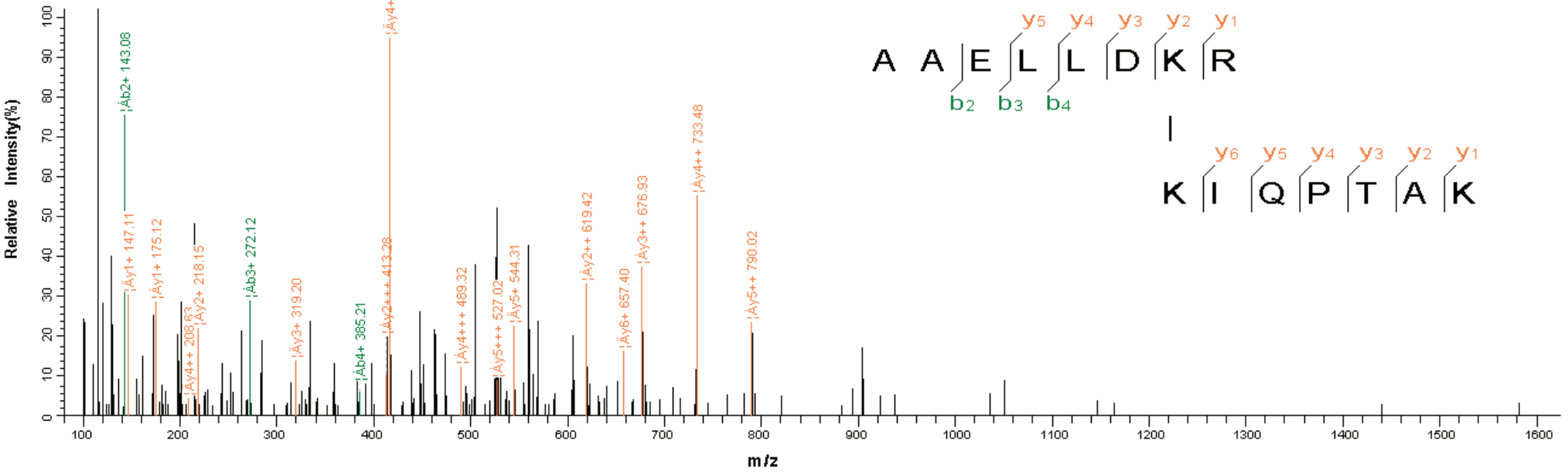
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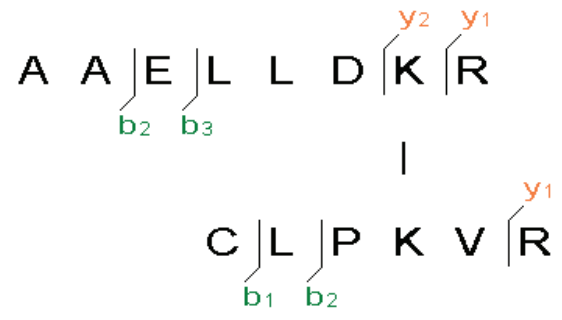
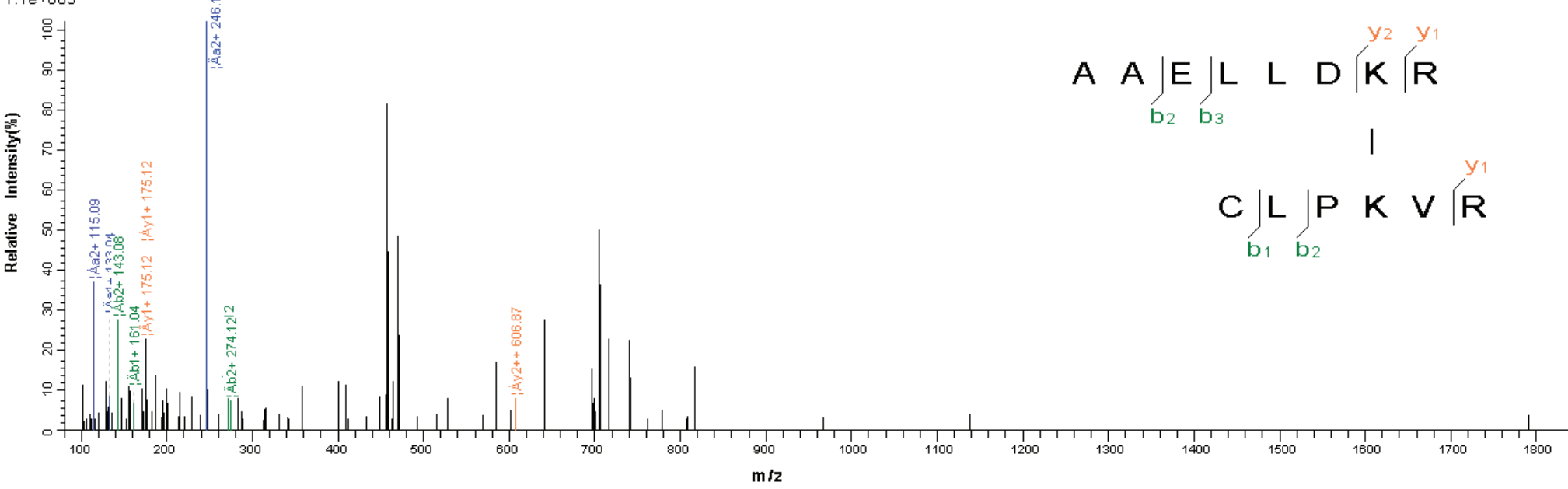
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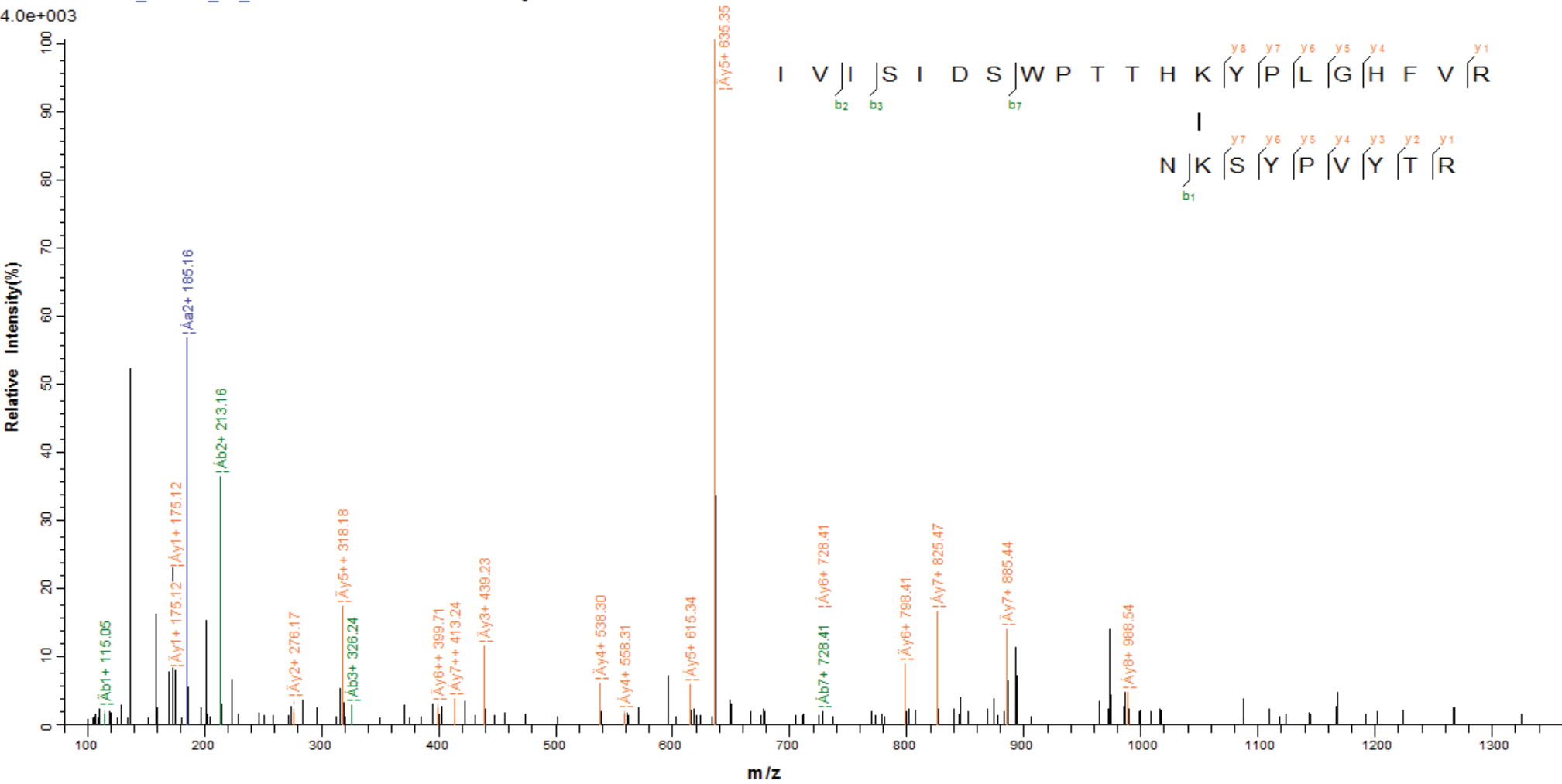


exosome_A9_140503145423.2671.2671.4.dta Seq: AAELLDKR-KIQPTAK

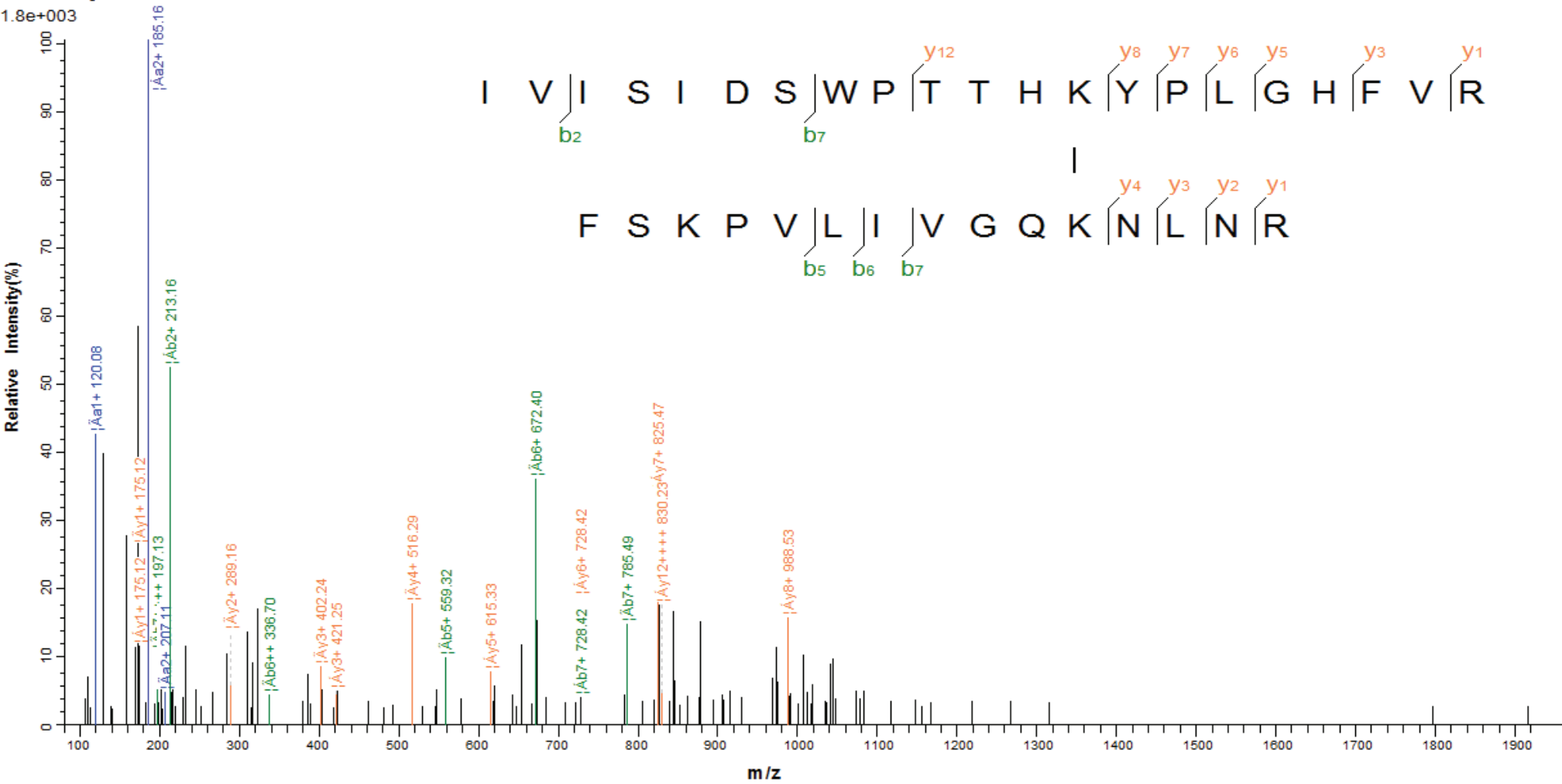


exosome_A9_140503145423.3261.3261.4.dta Seq: AAELLDKR-CLPKVR Mod: 12,C_57

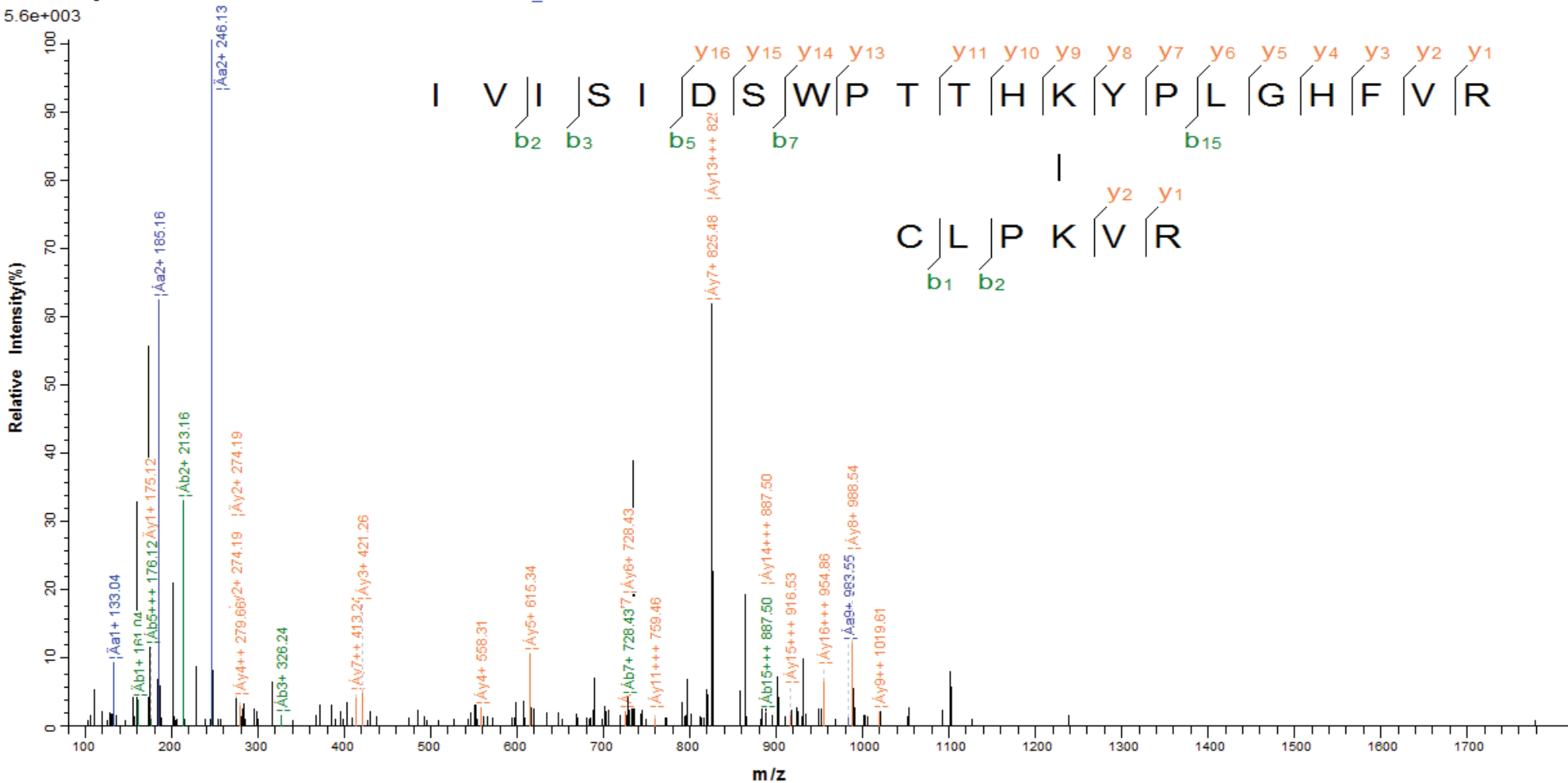




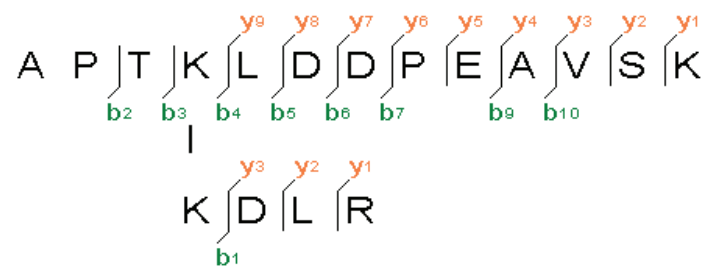
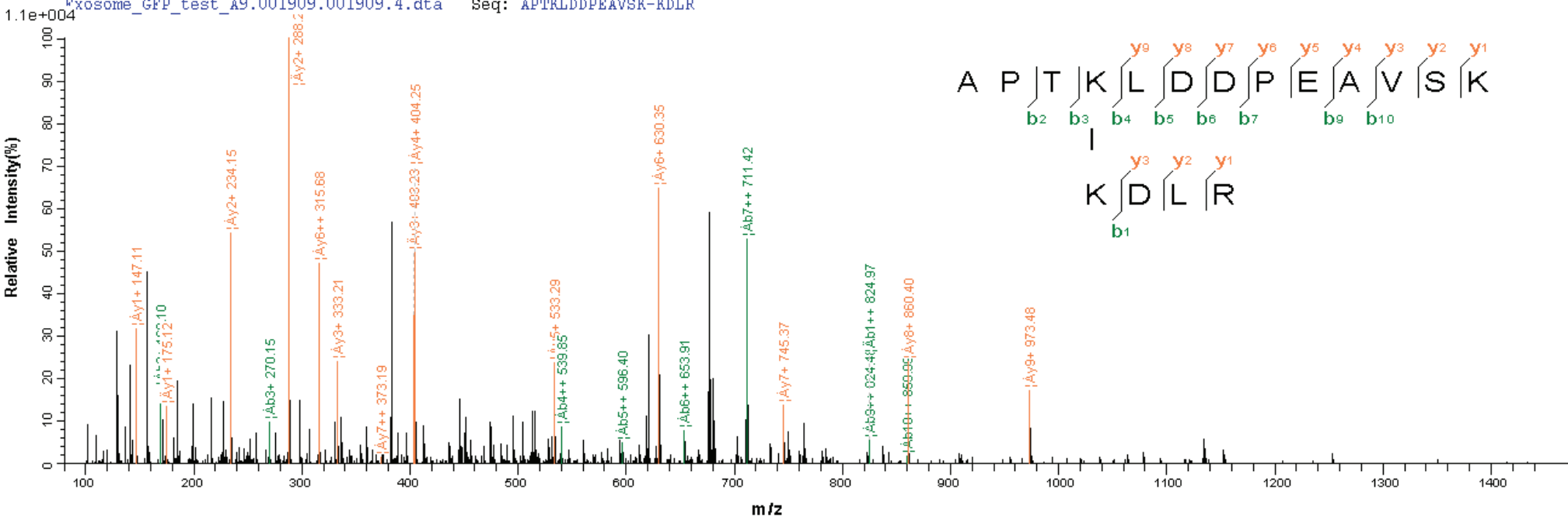
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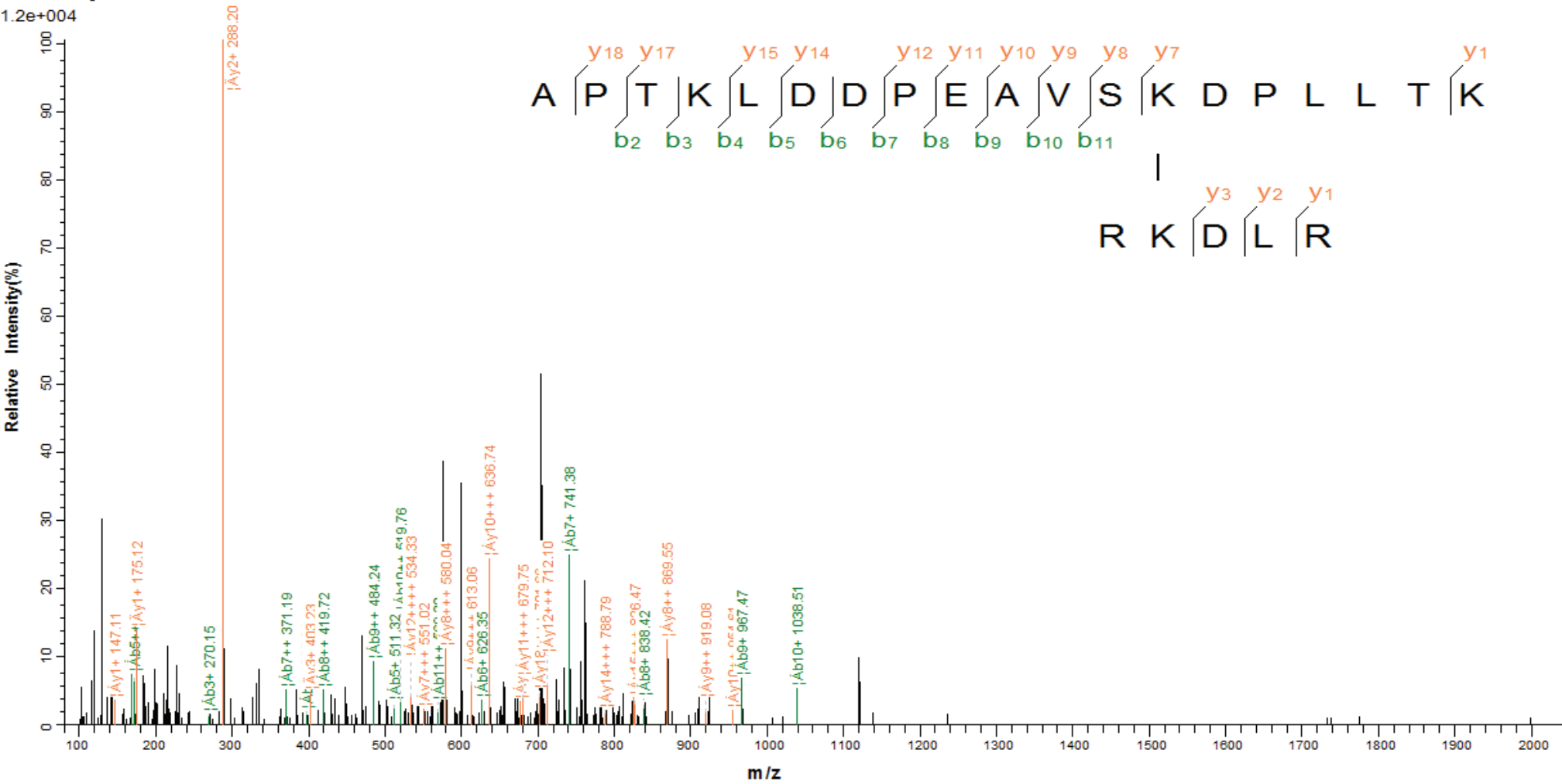
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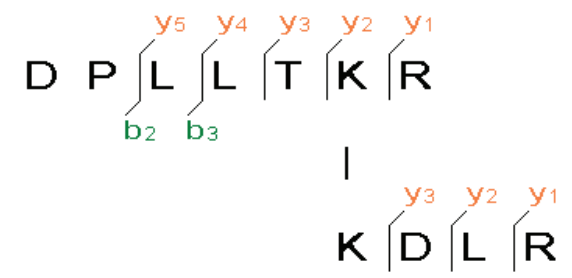
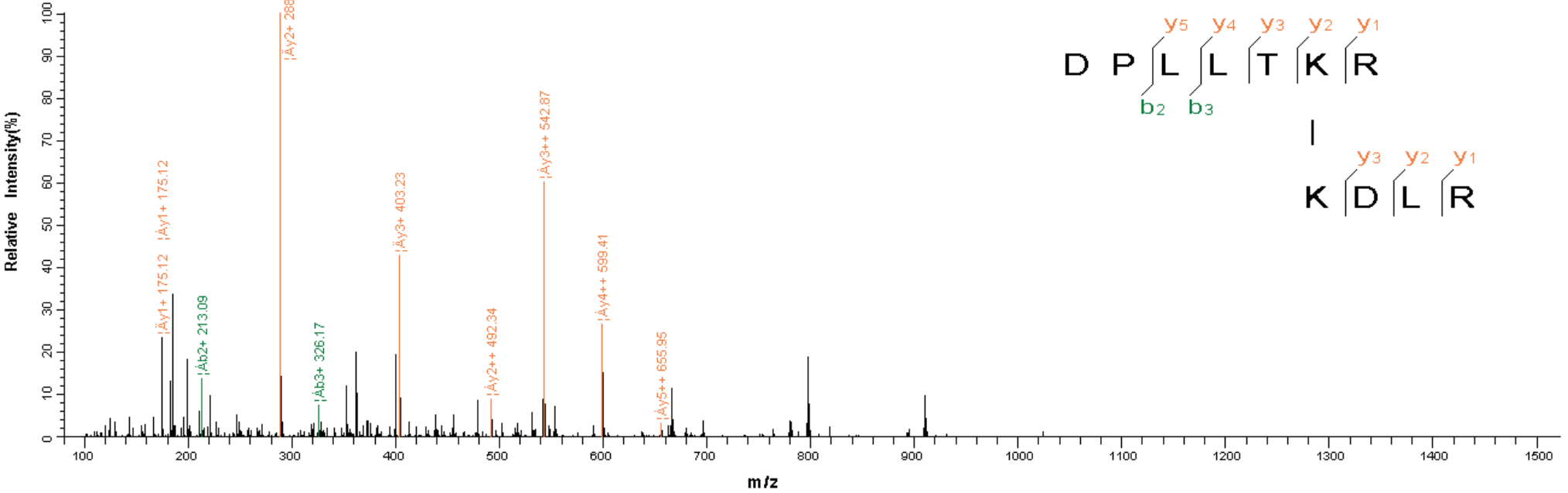
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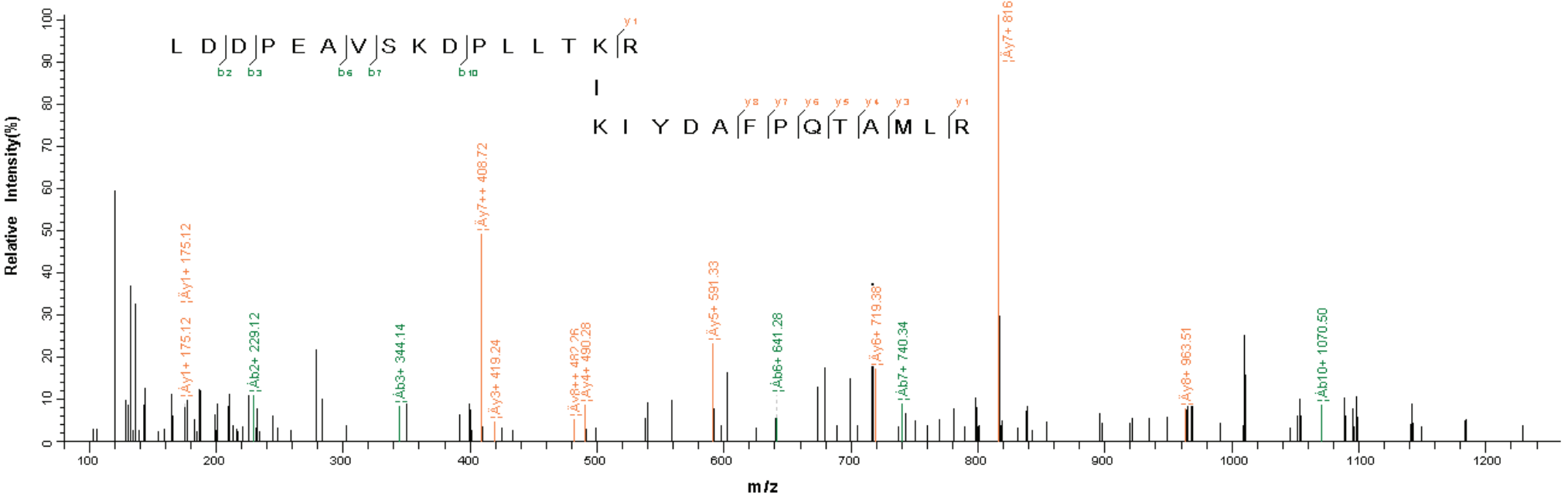
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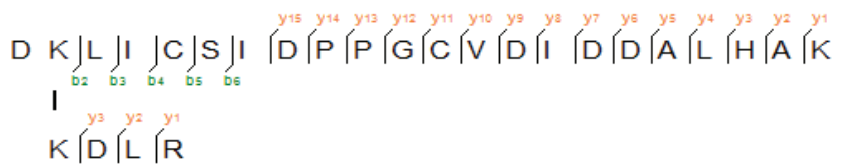
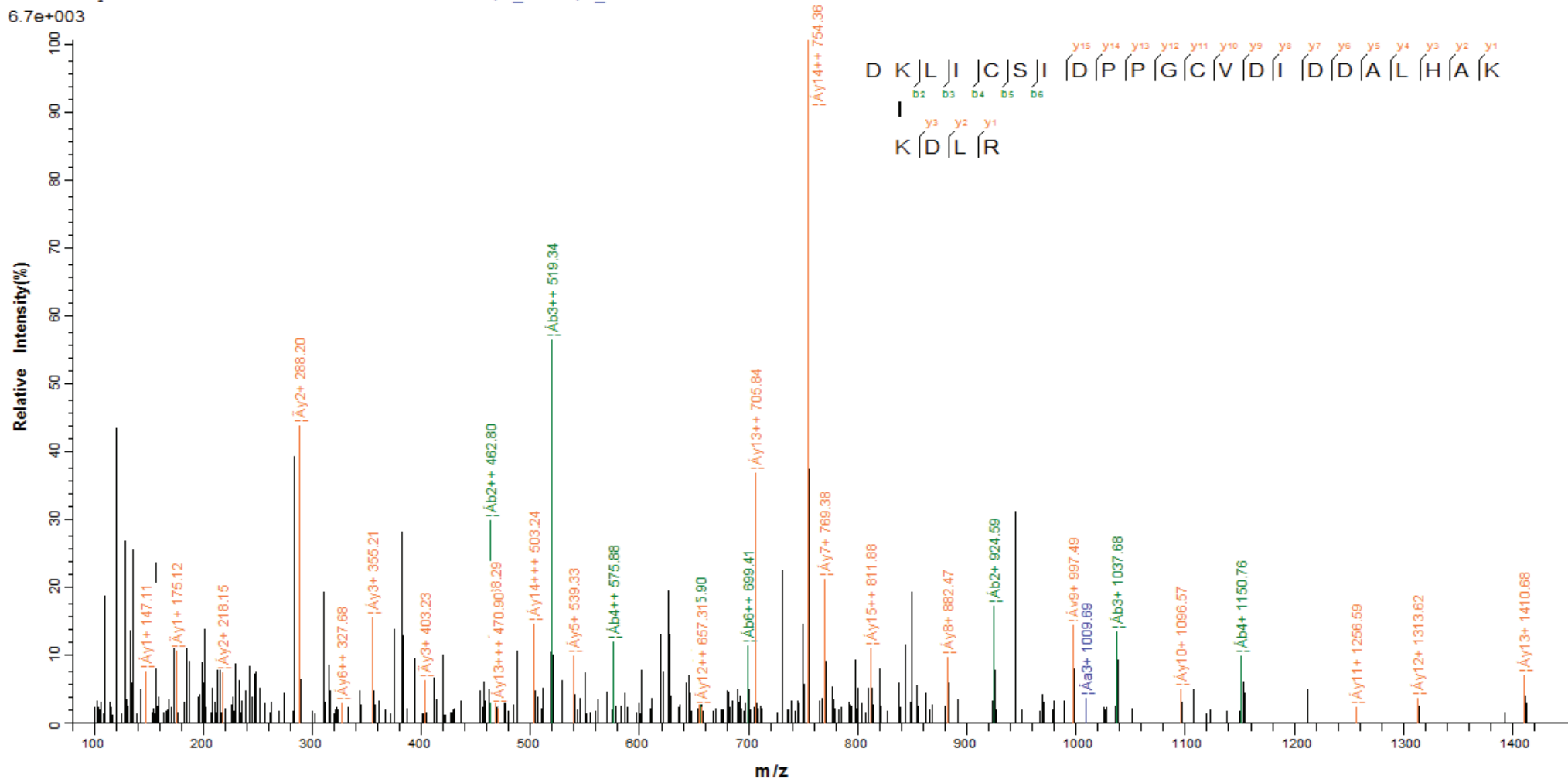
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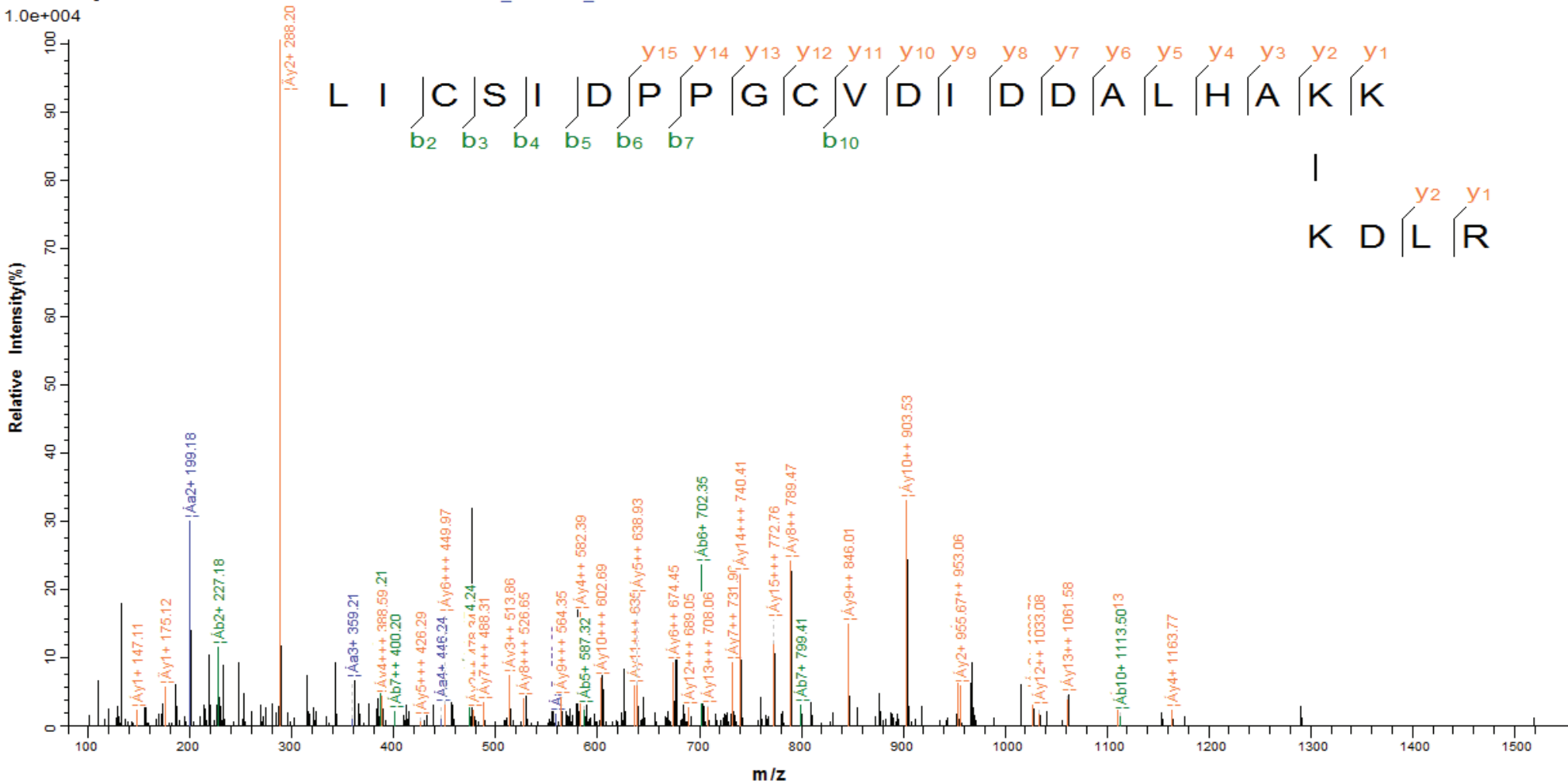
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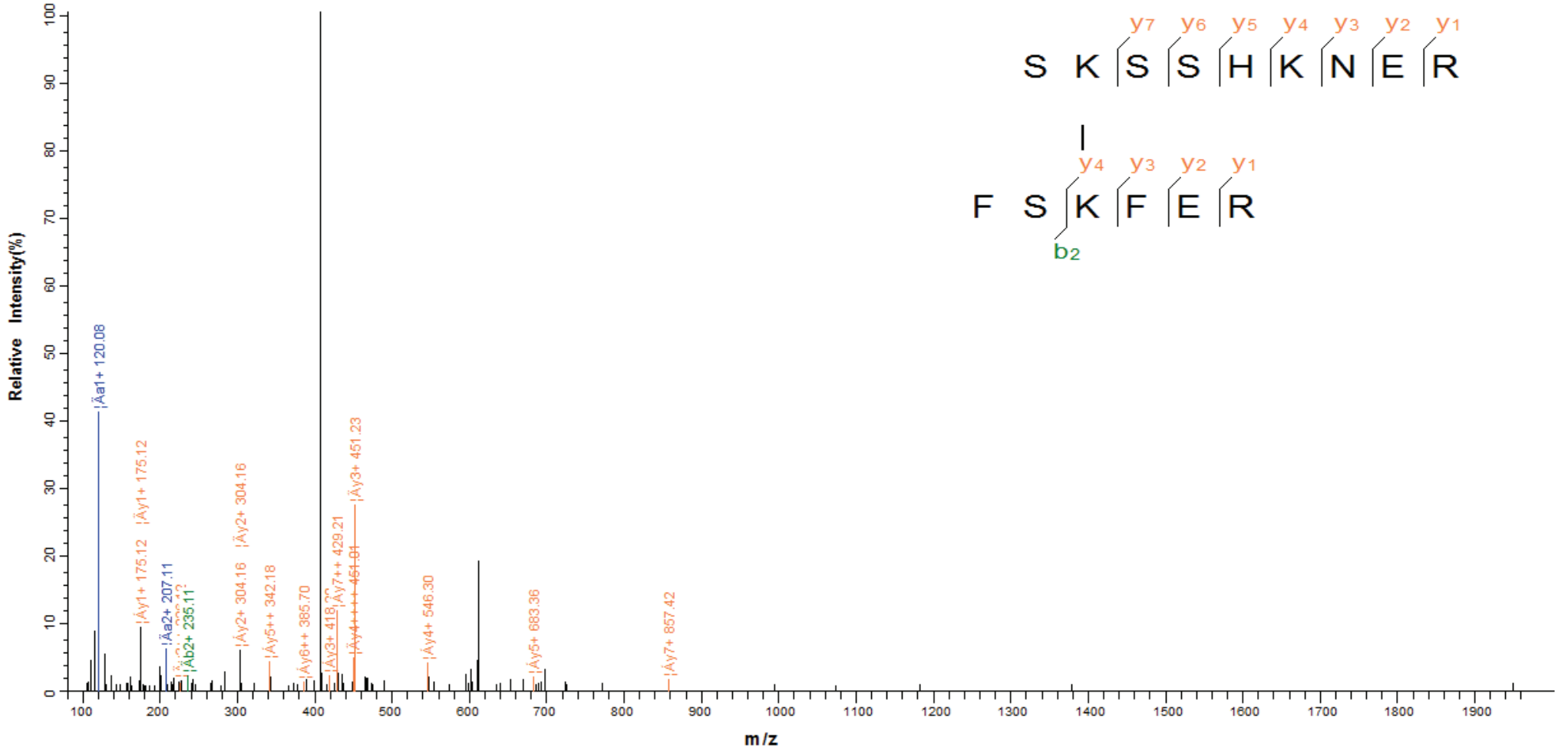


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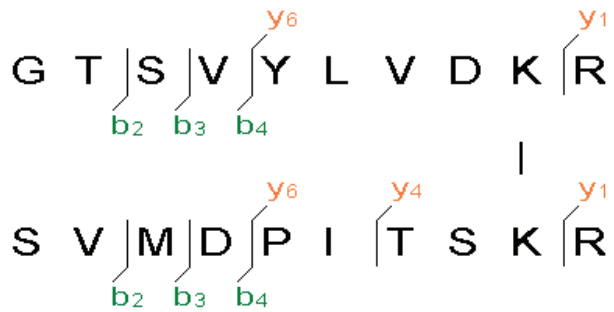
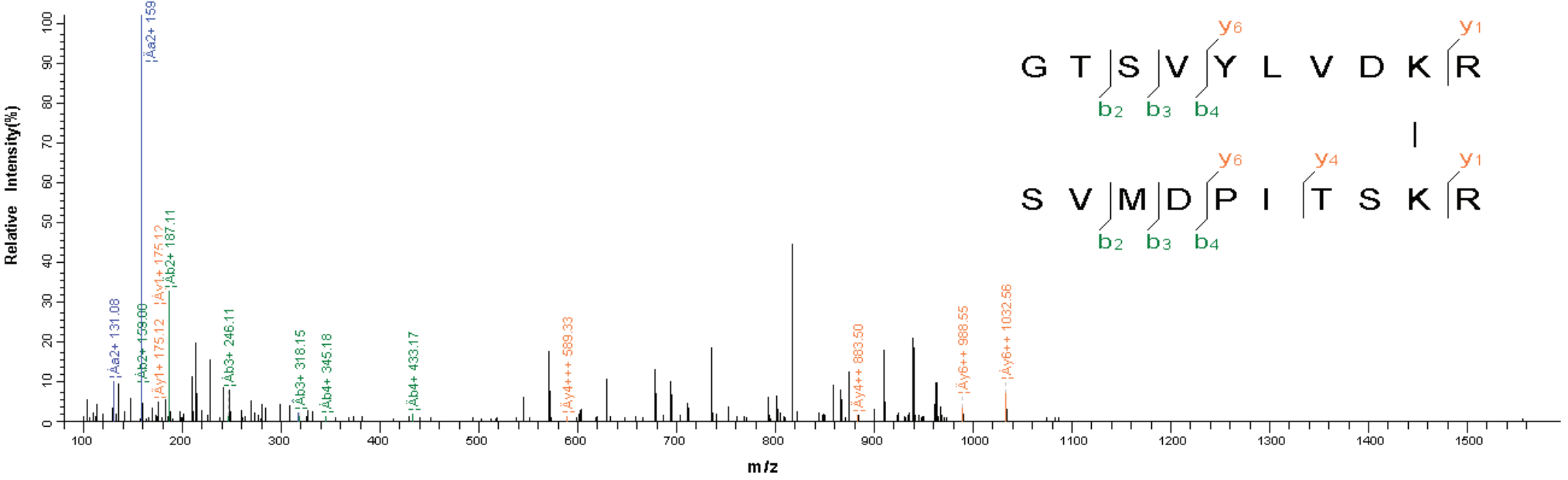


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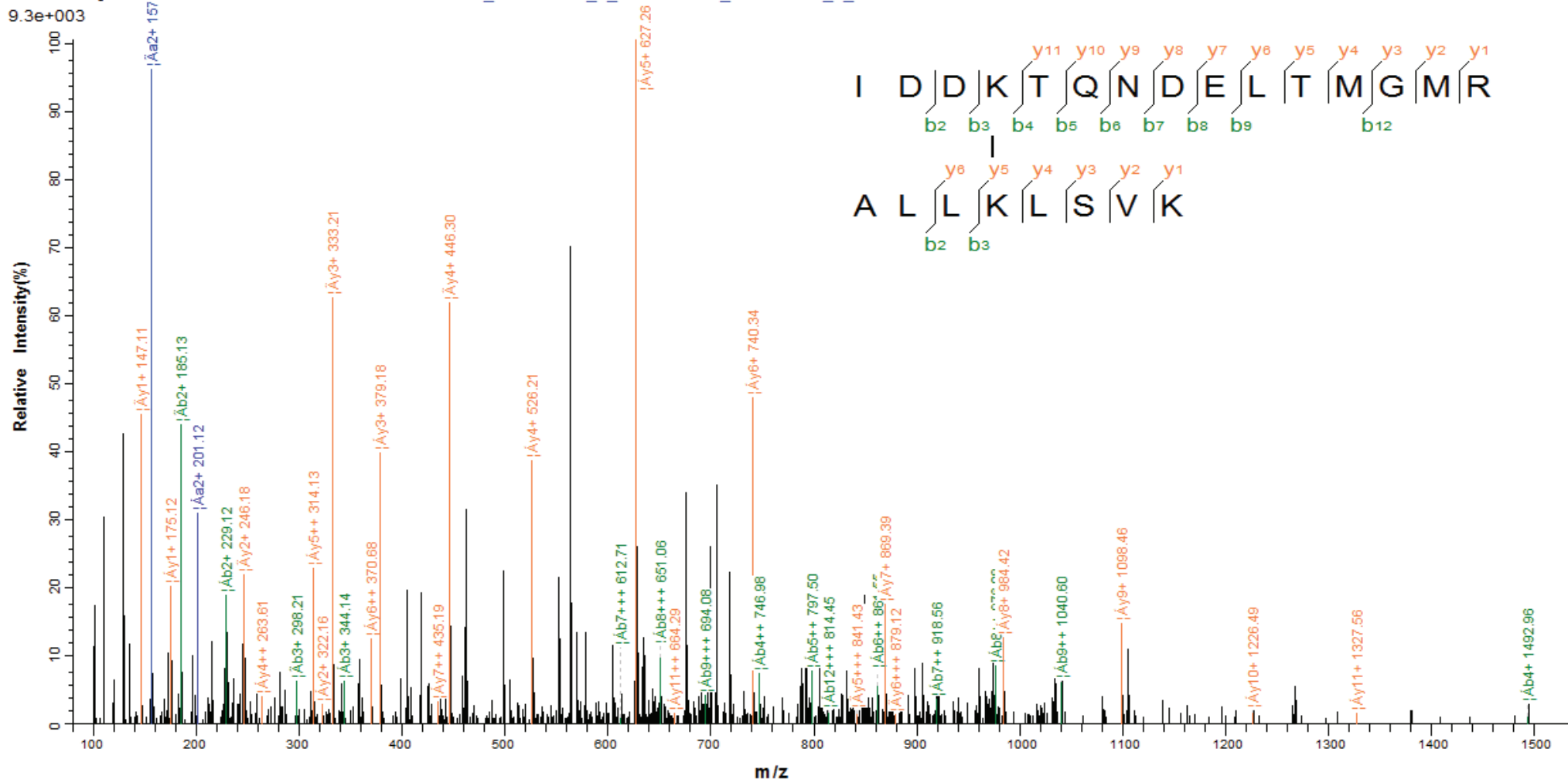
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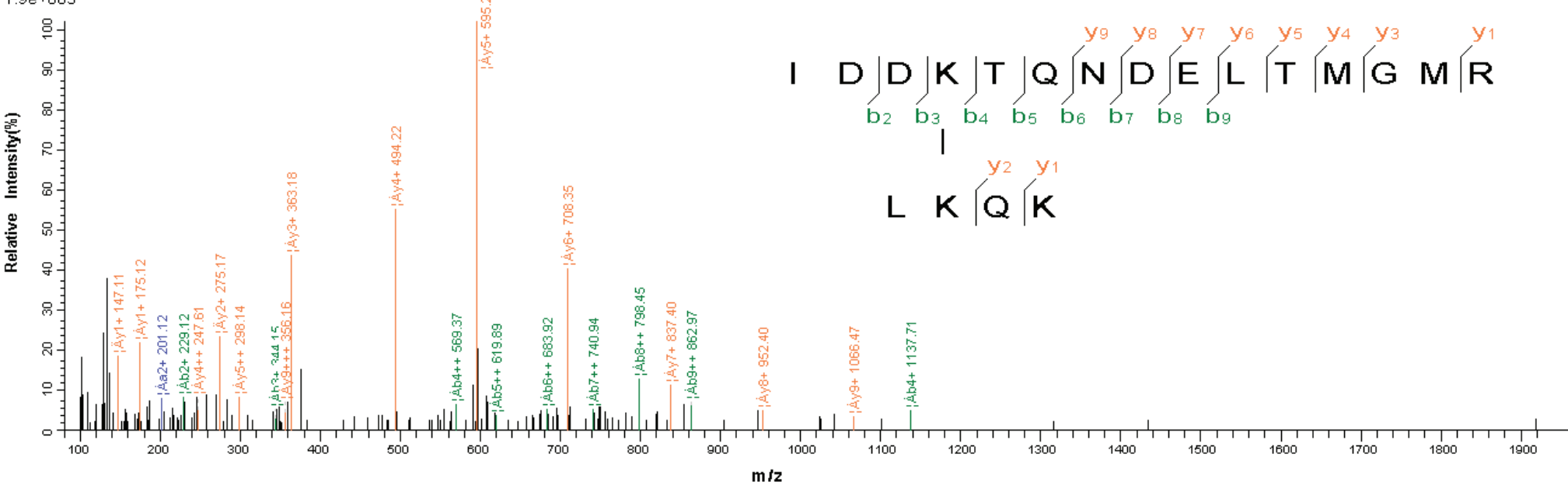
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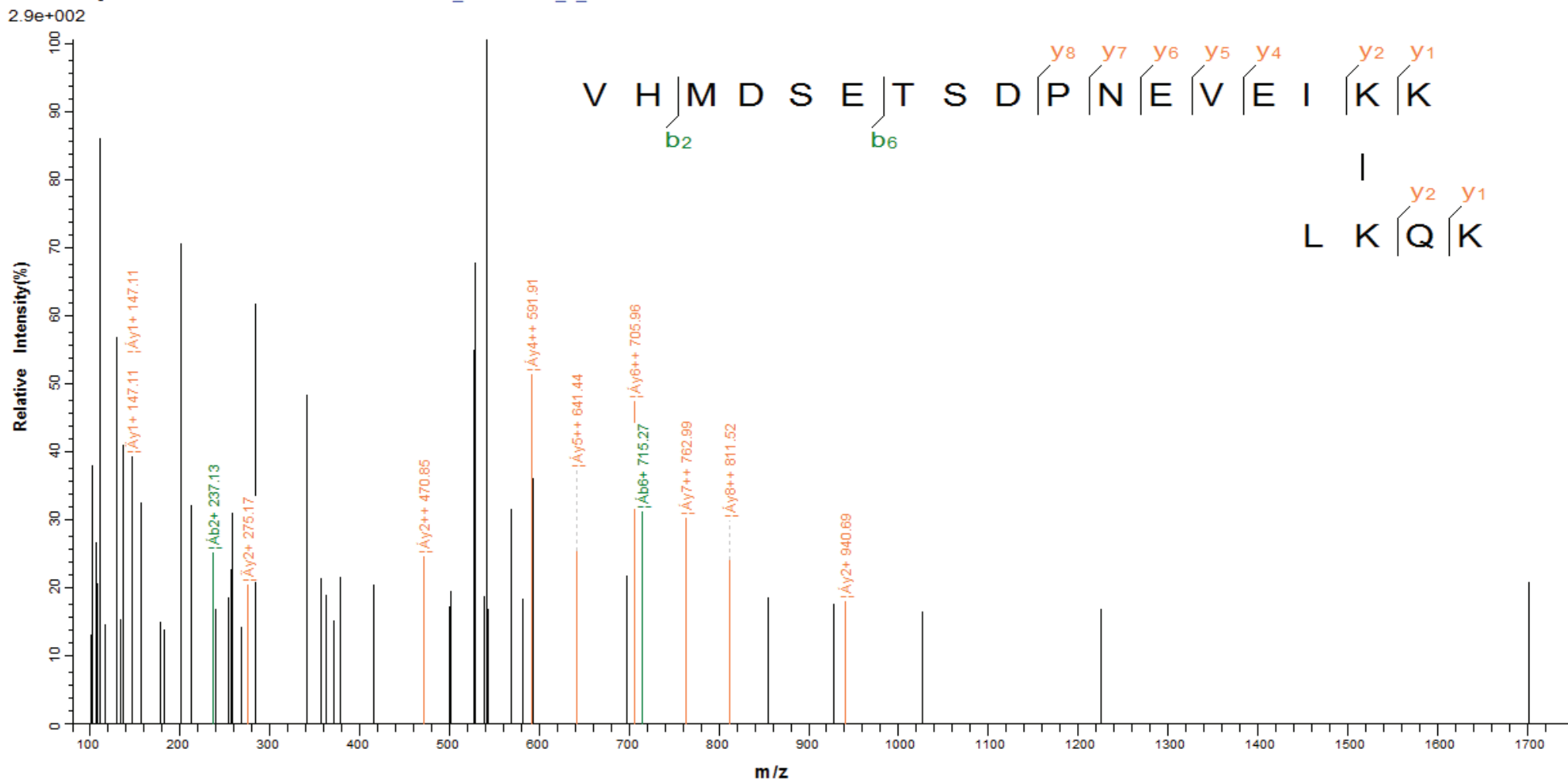
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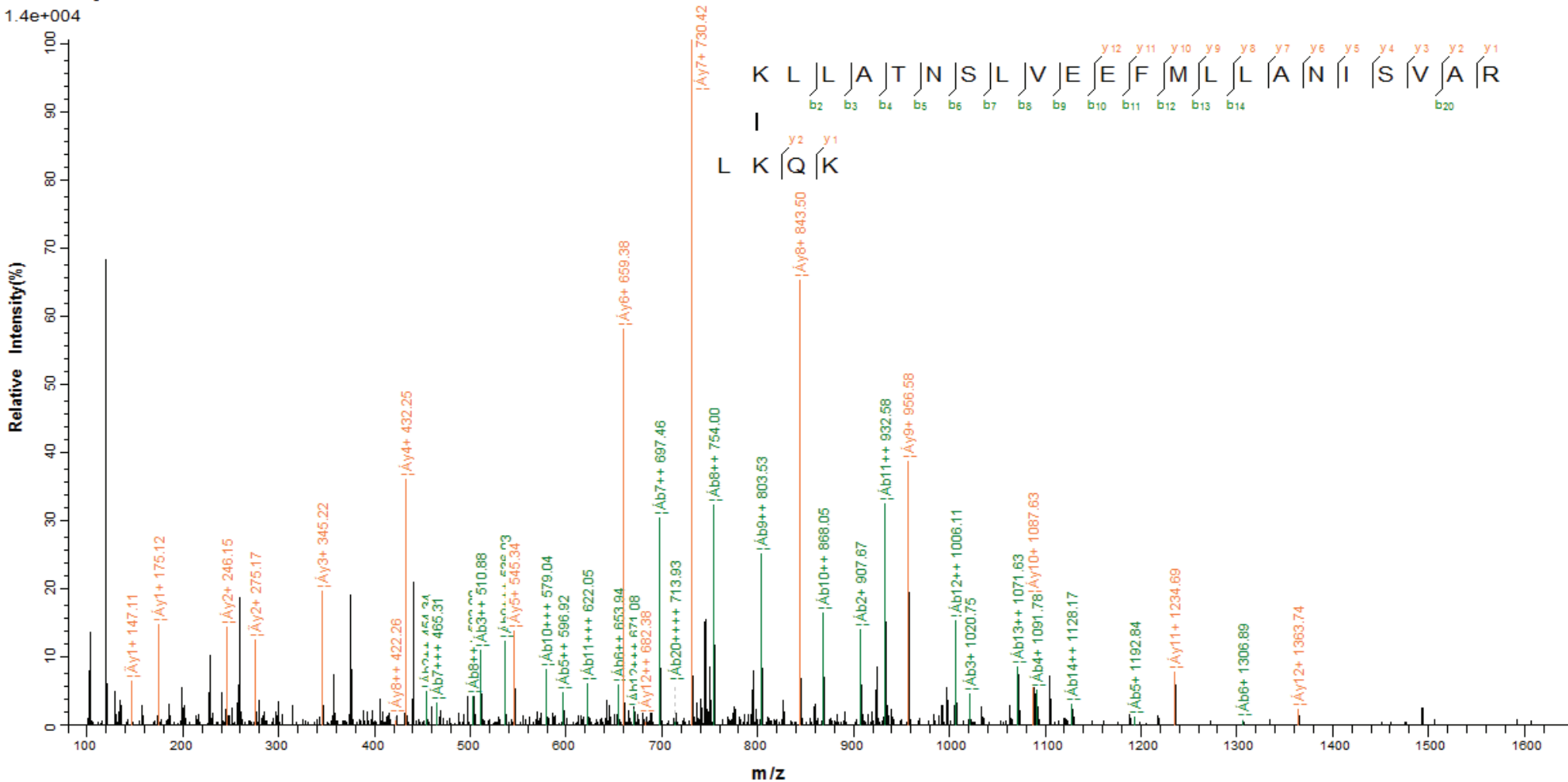
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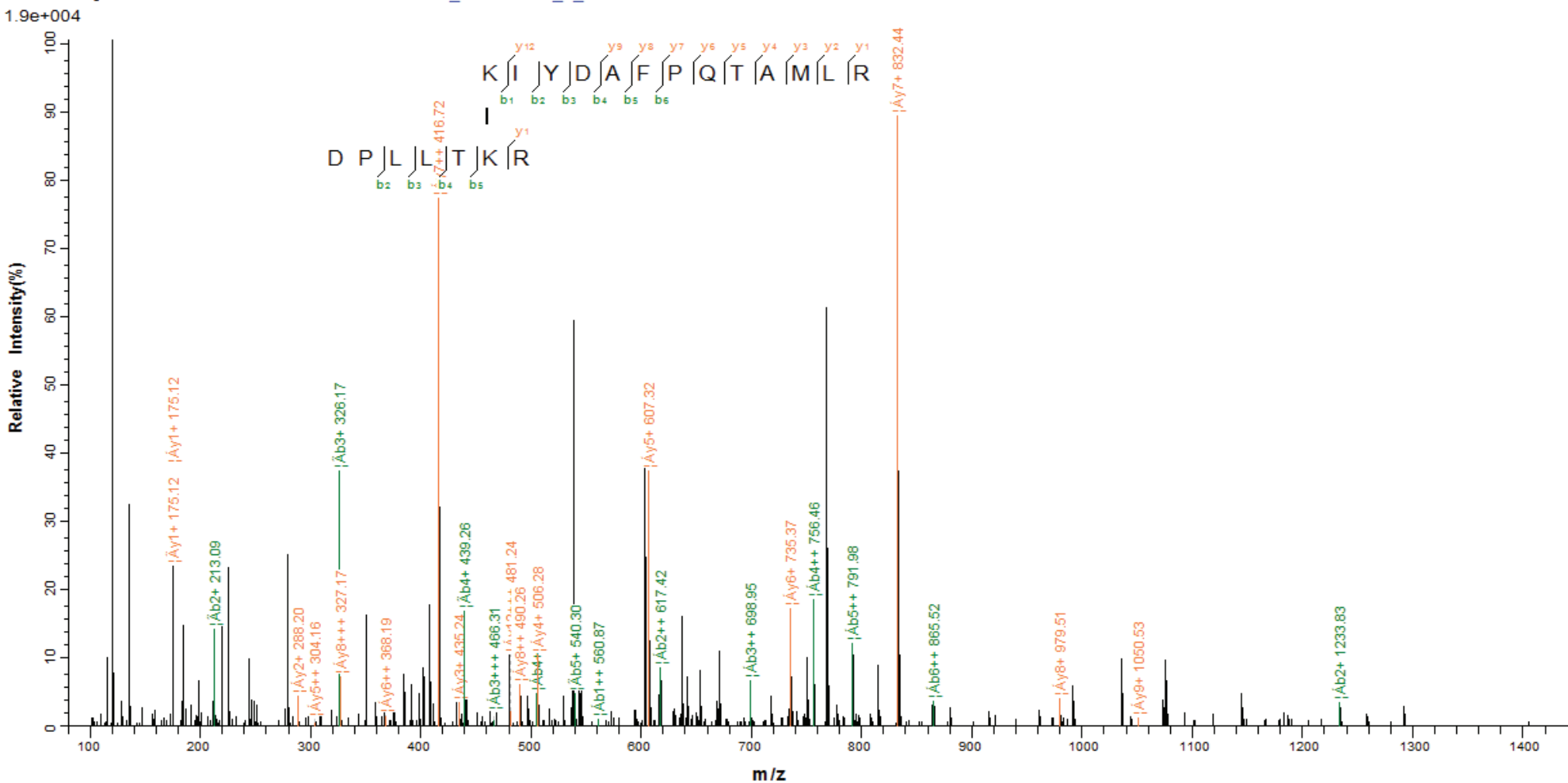
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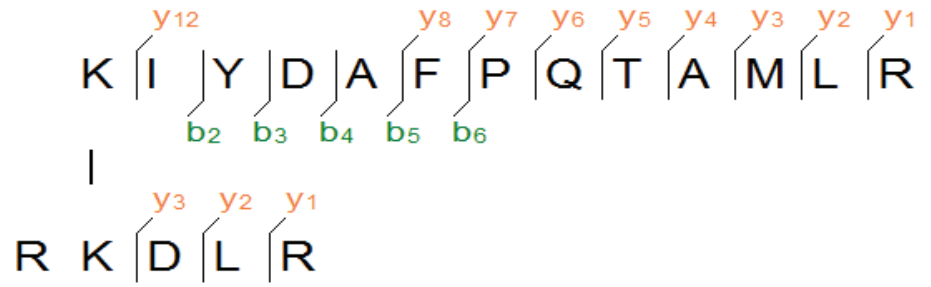
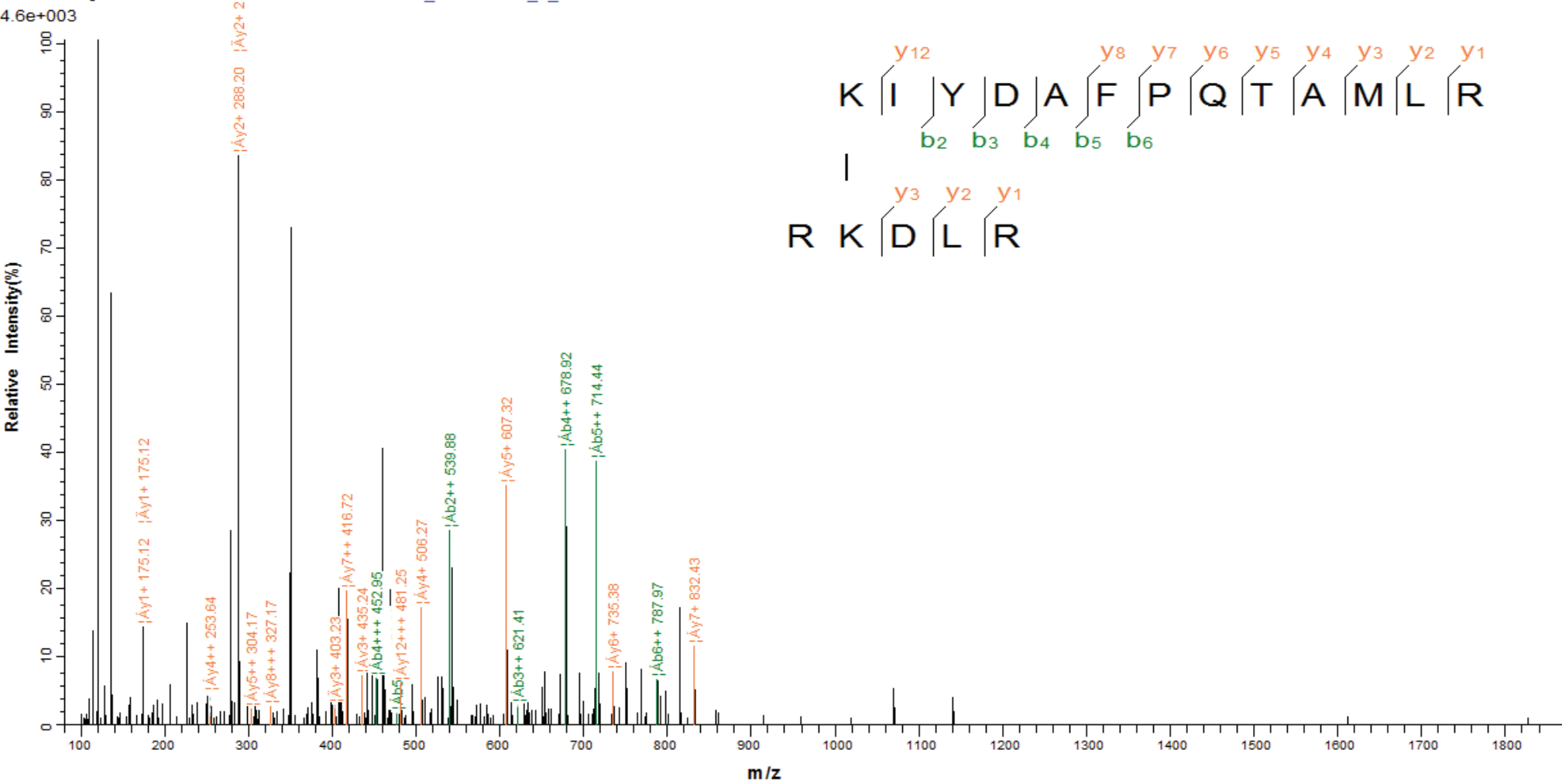
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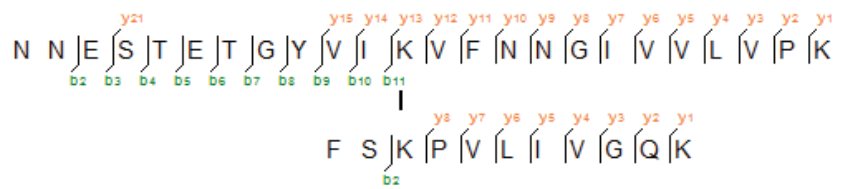
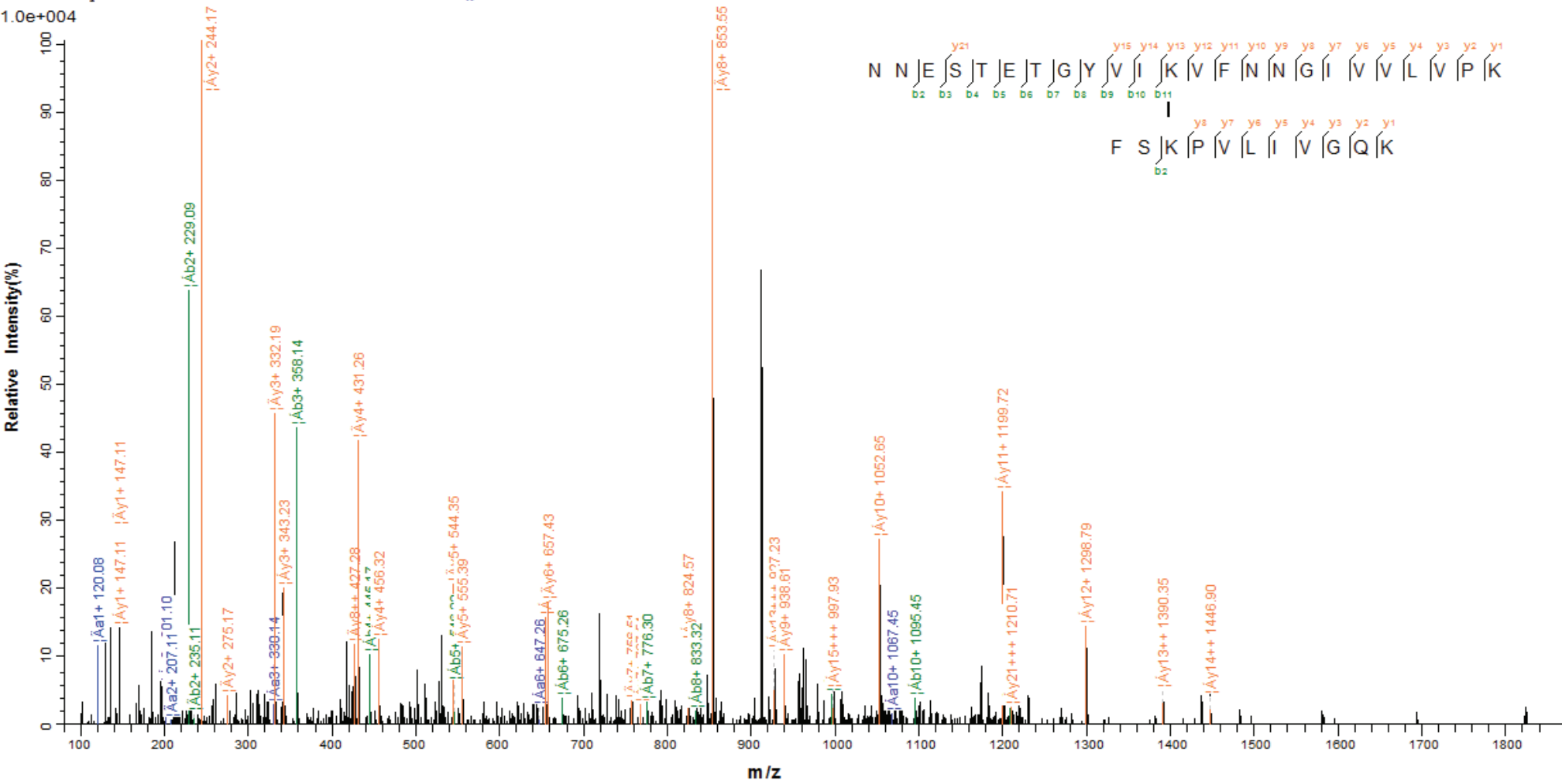
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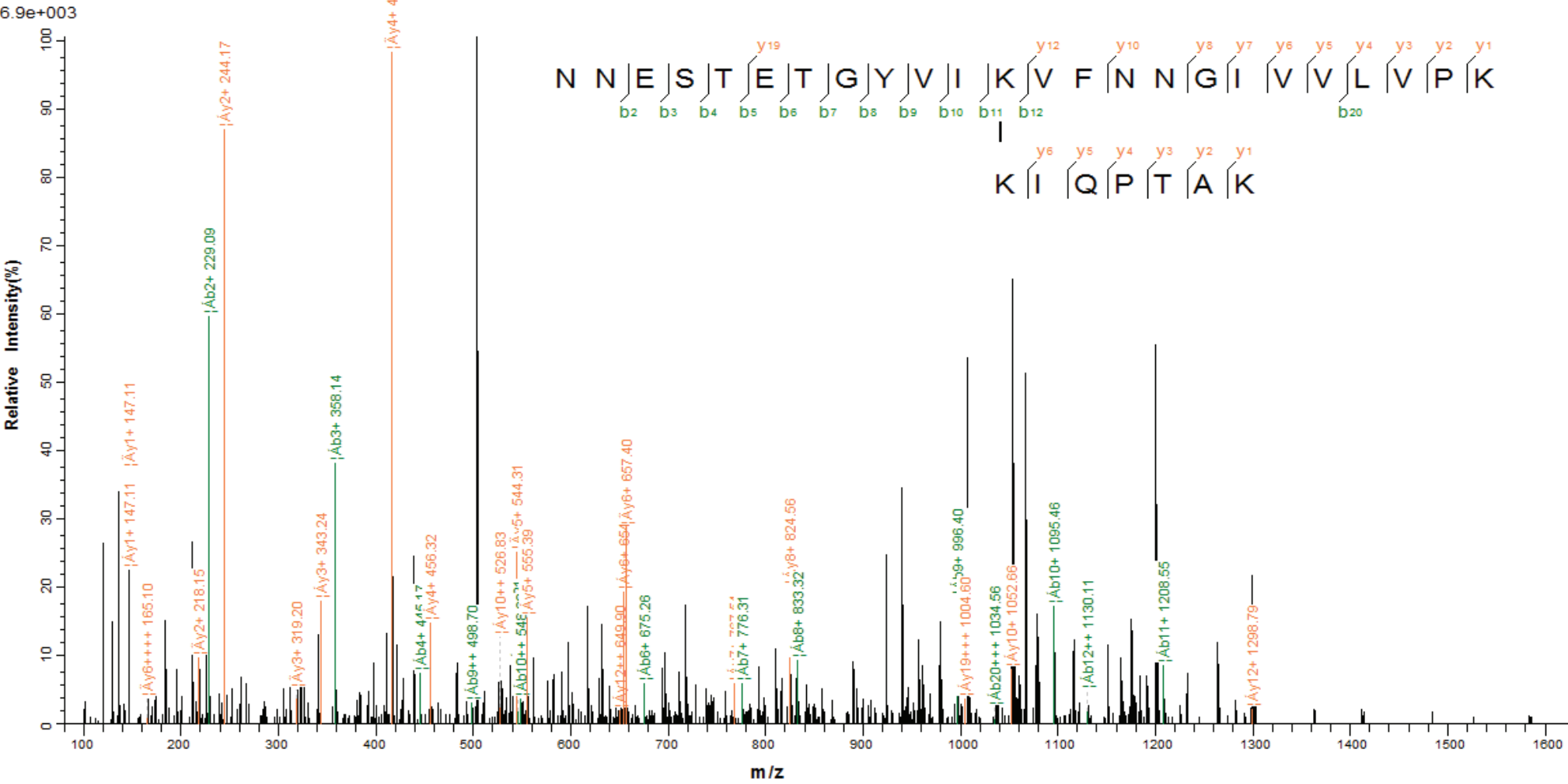
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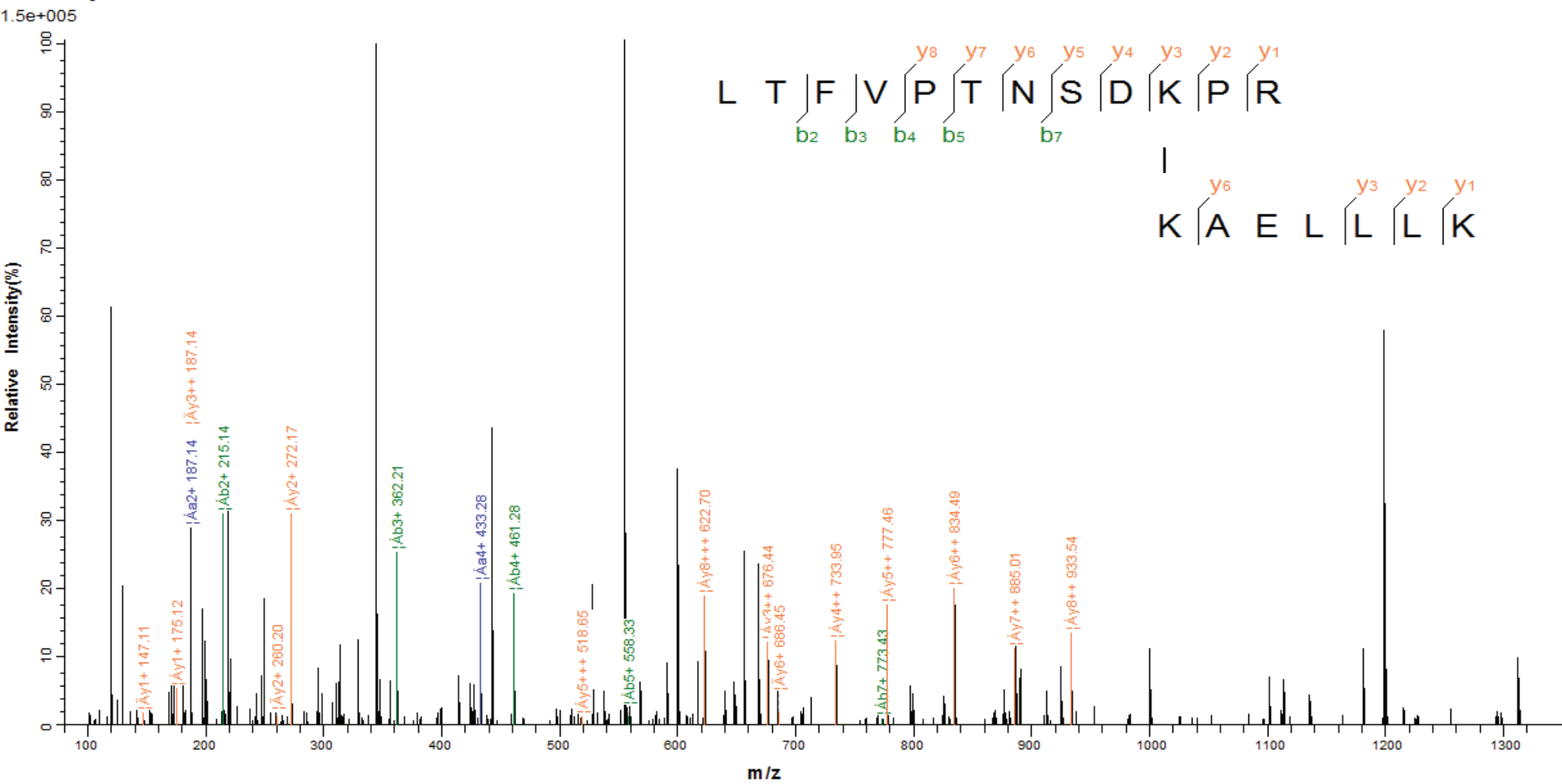
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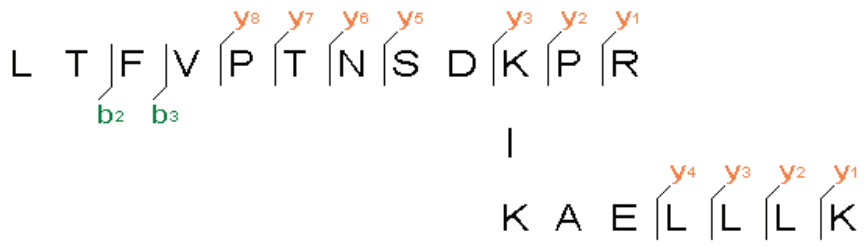
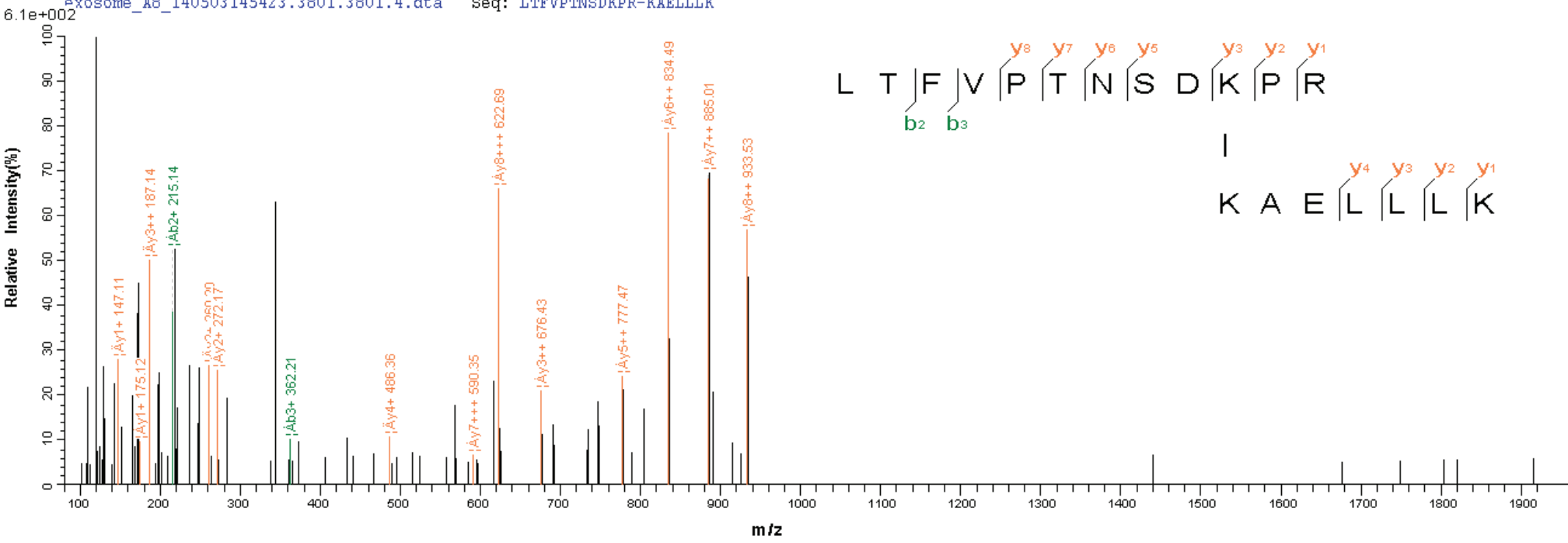
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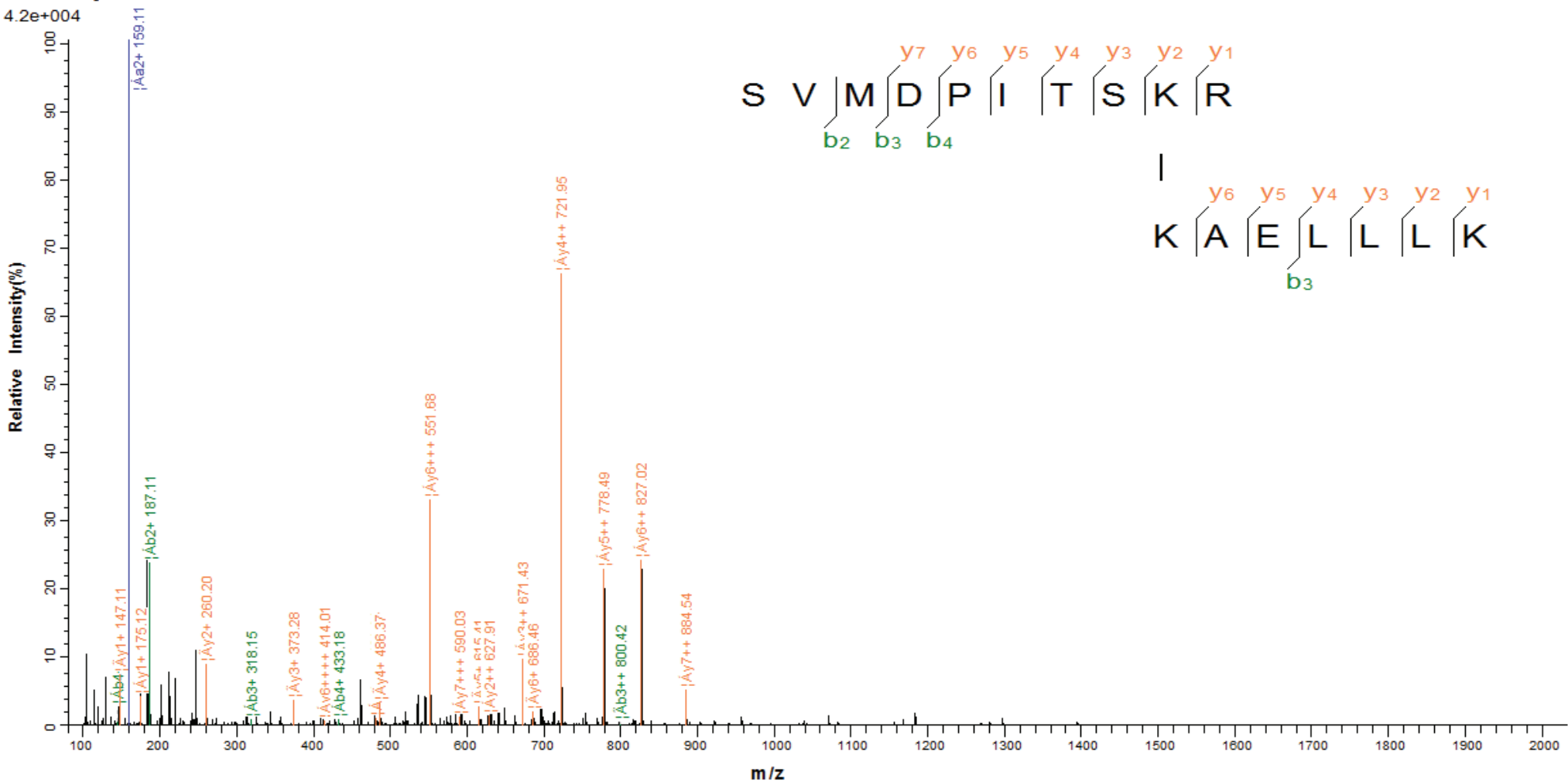
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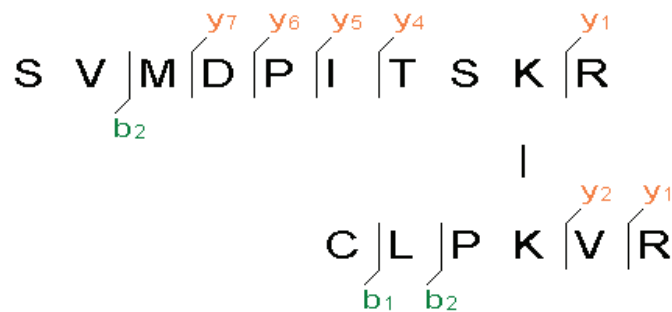
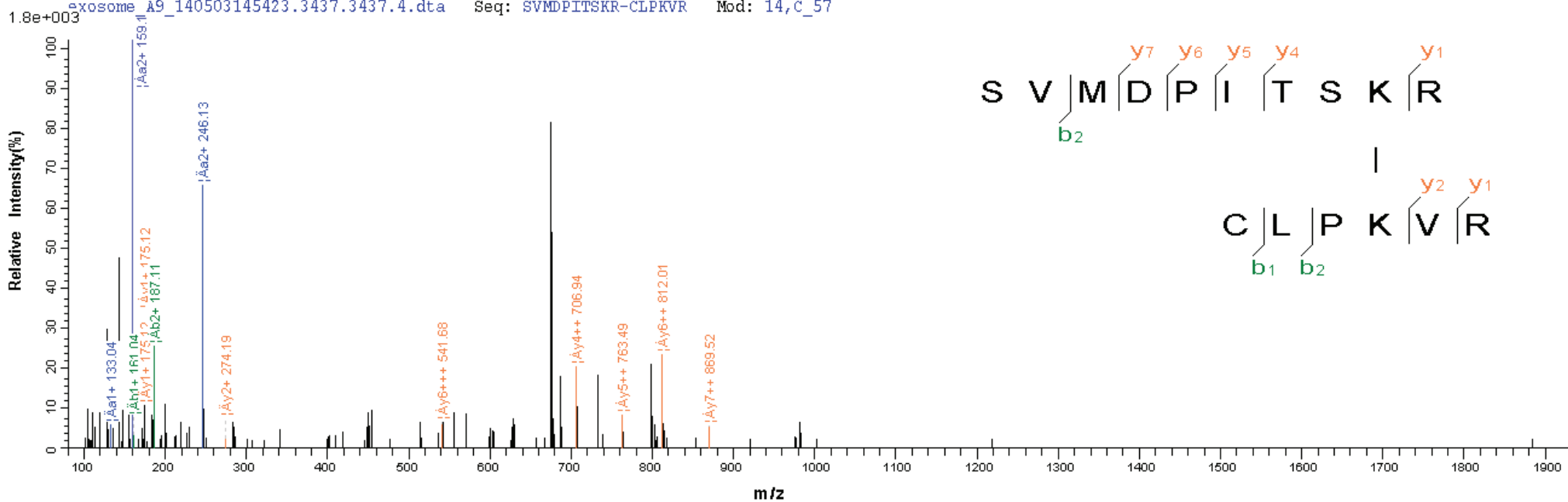
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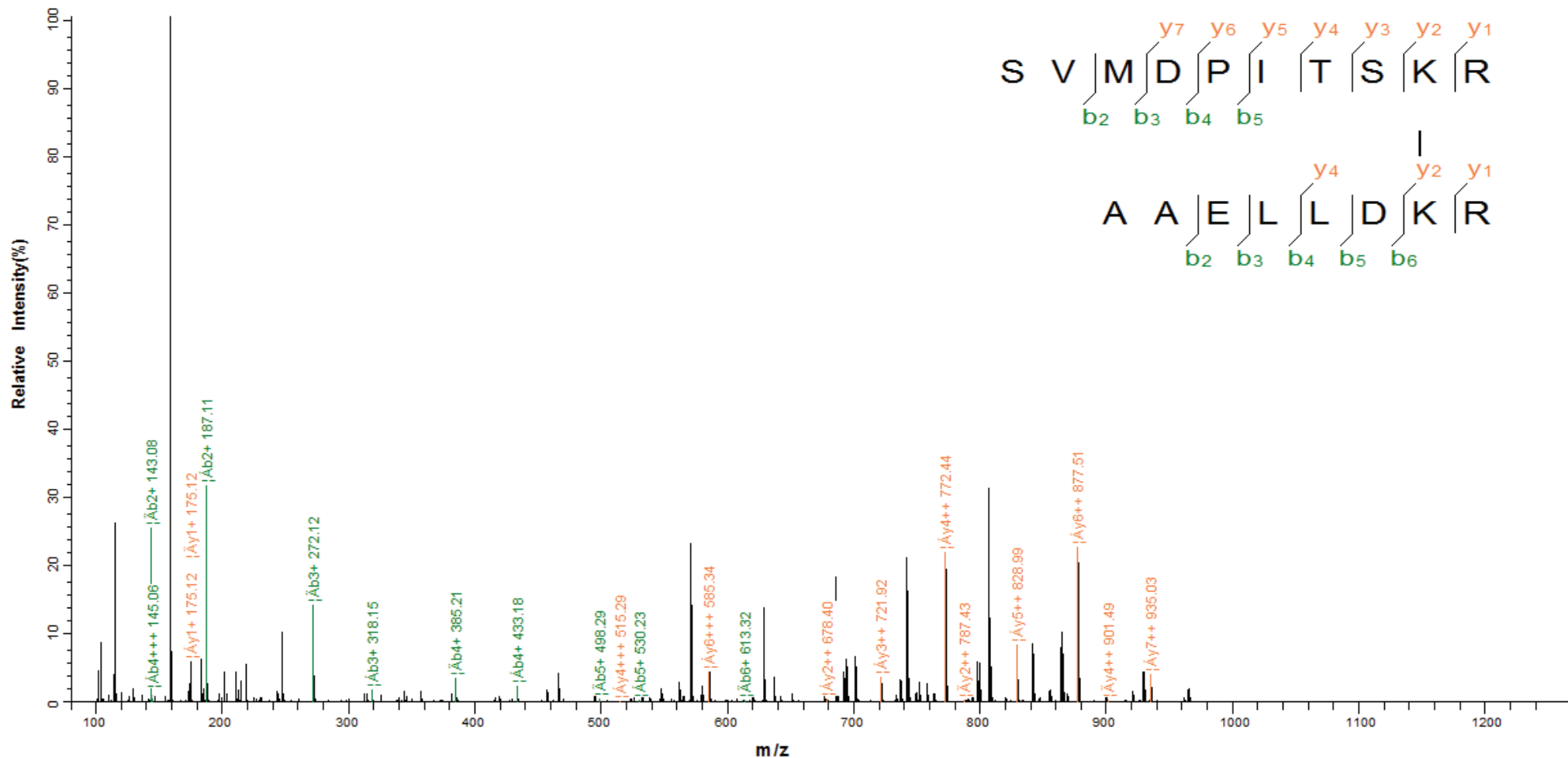


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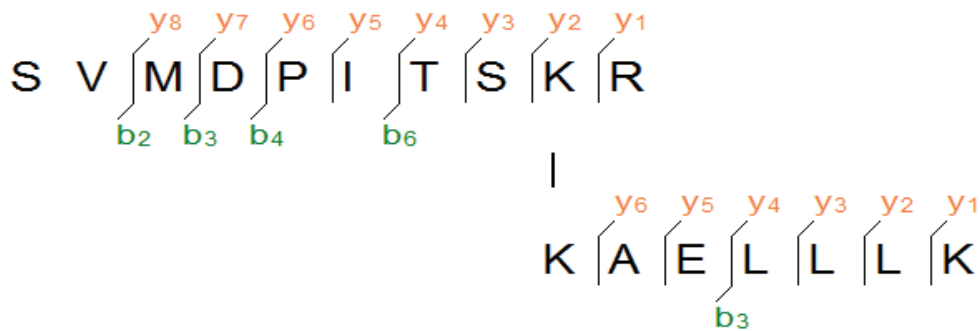
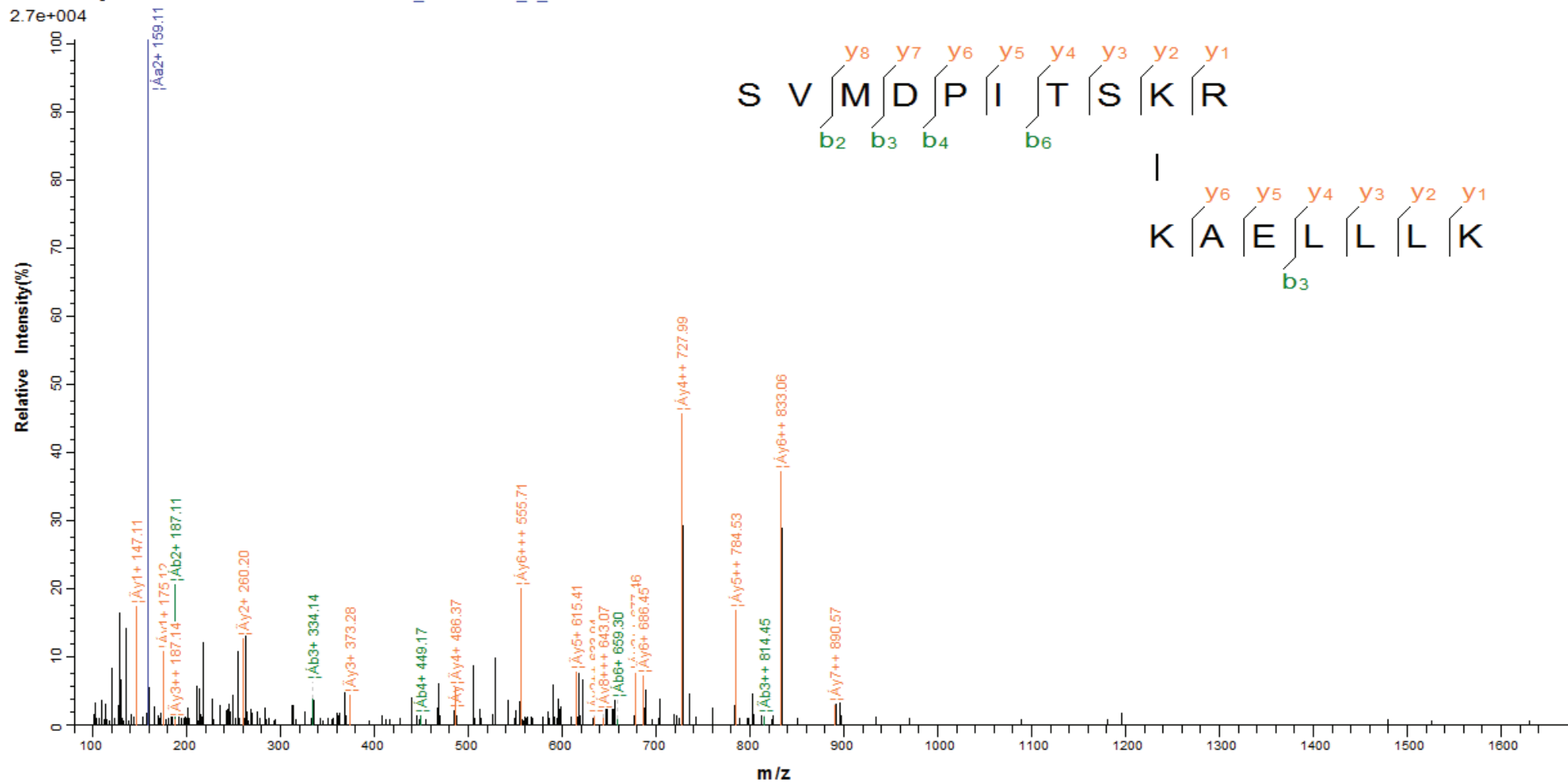


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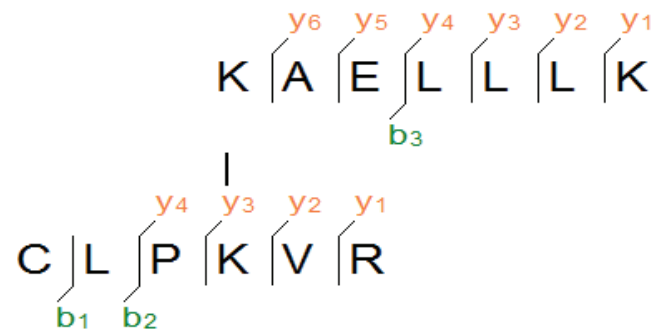
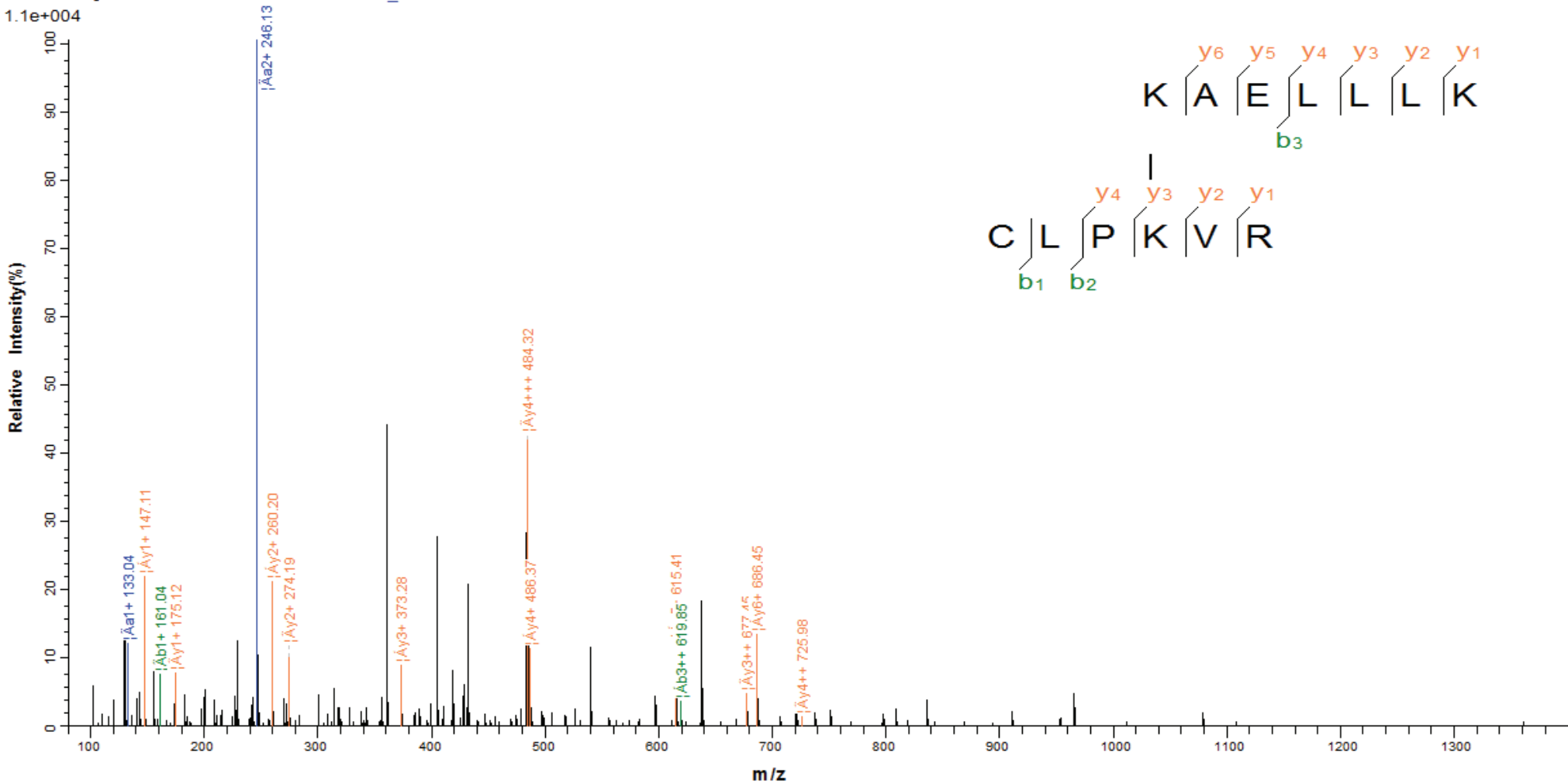
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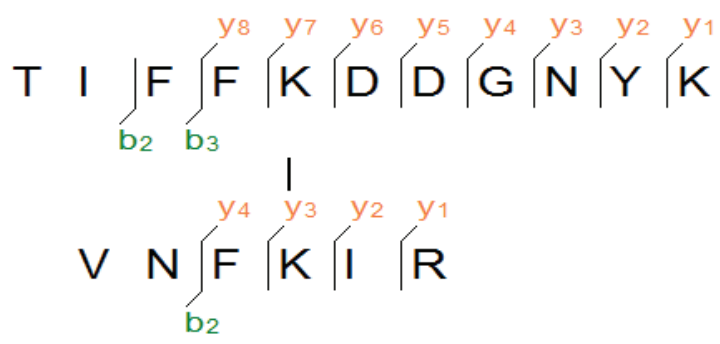
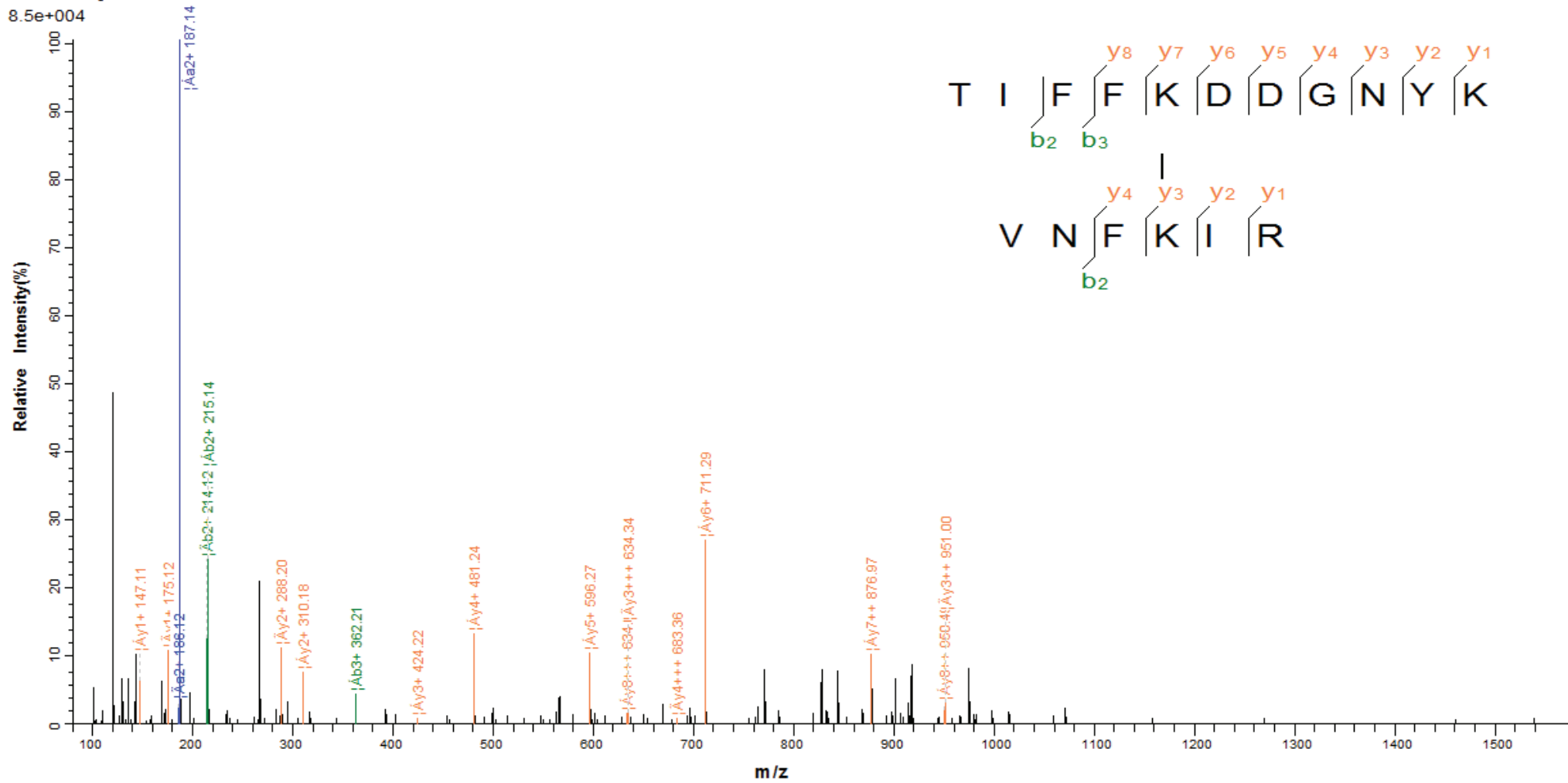
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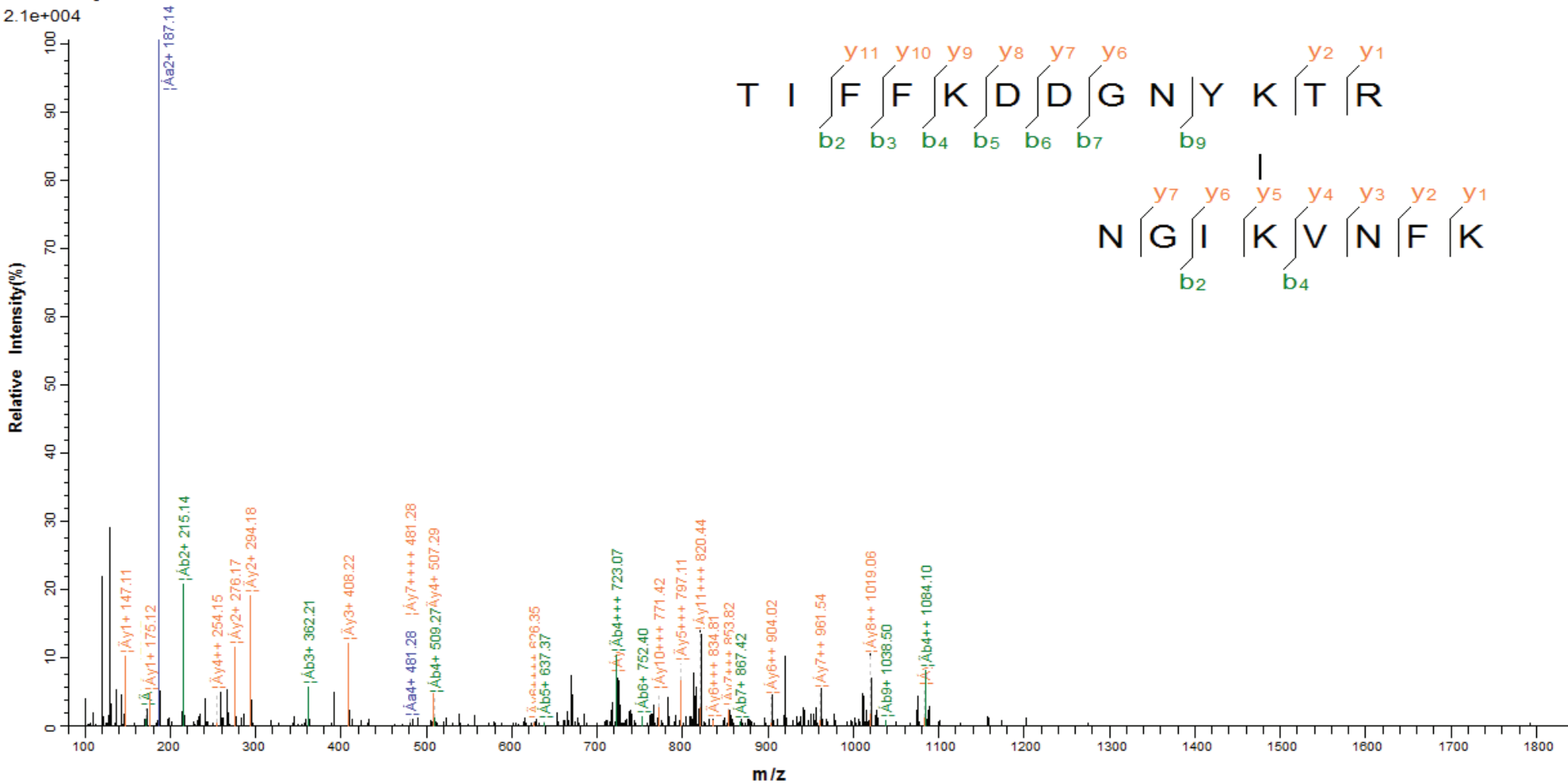
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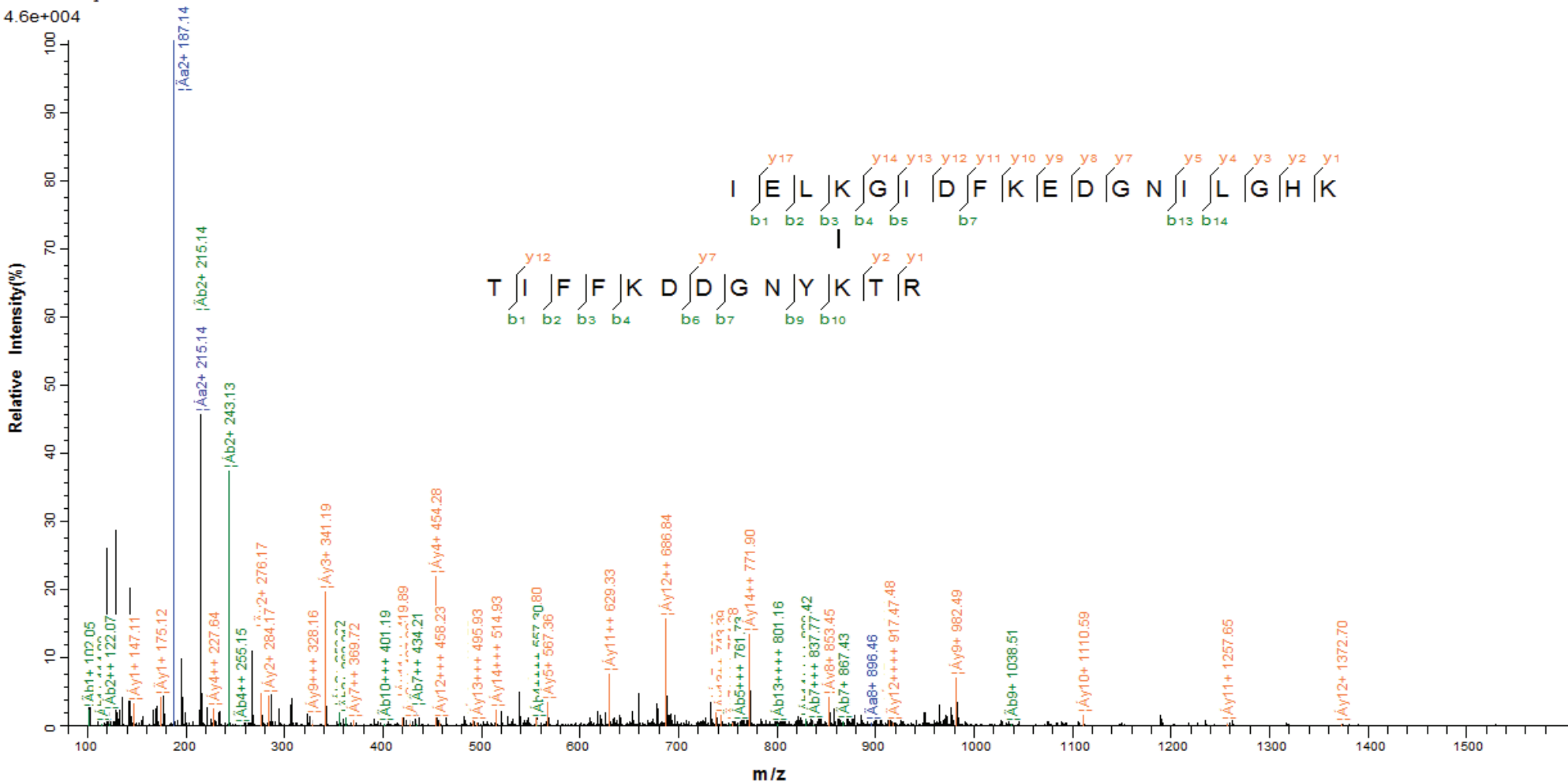
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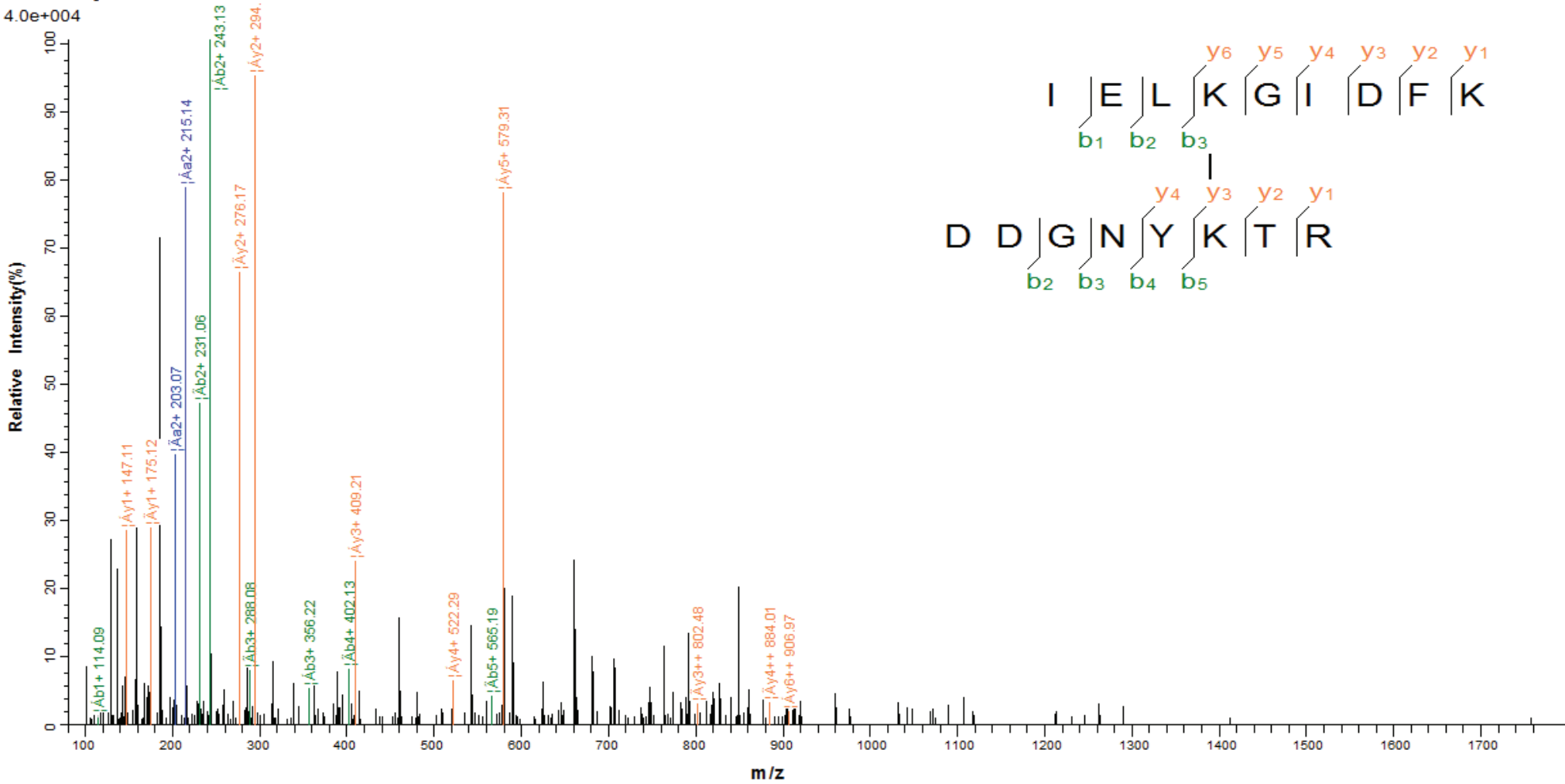
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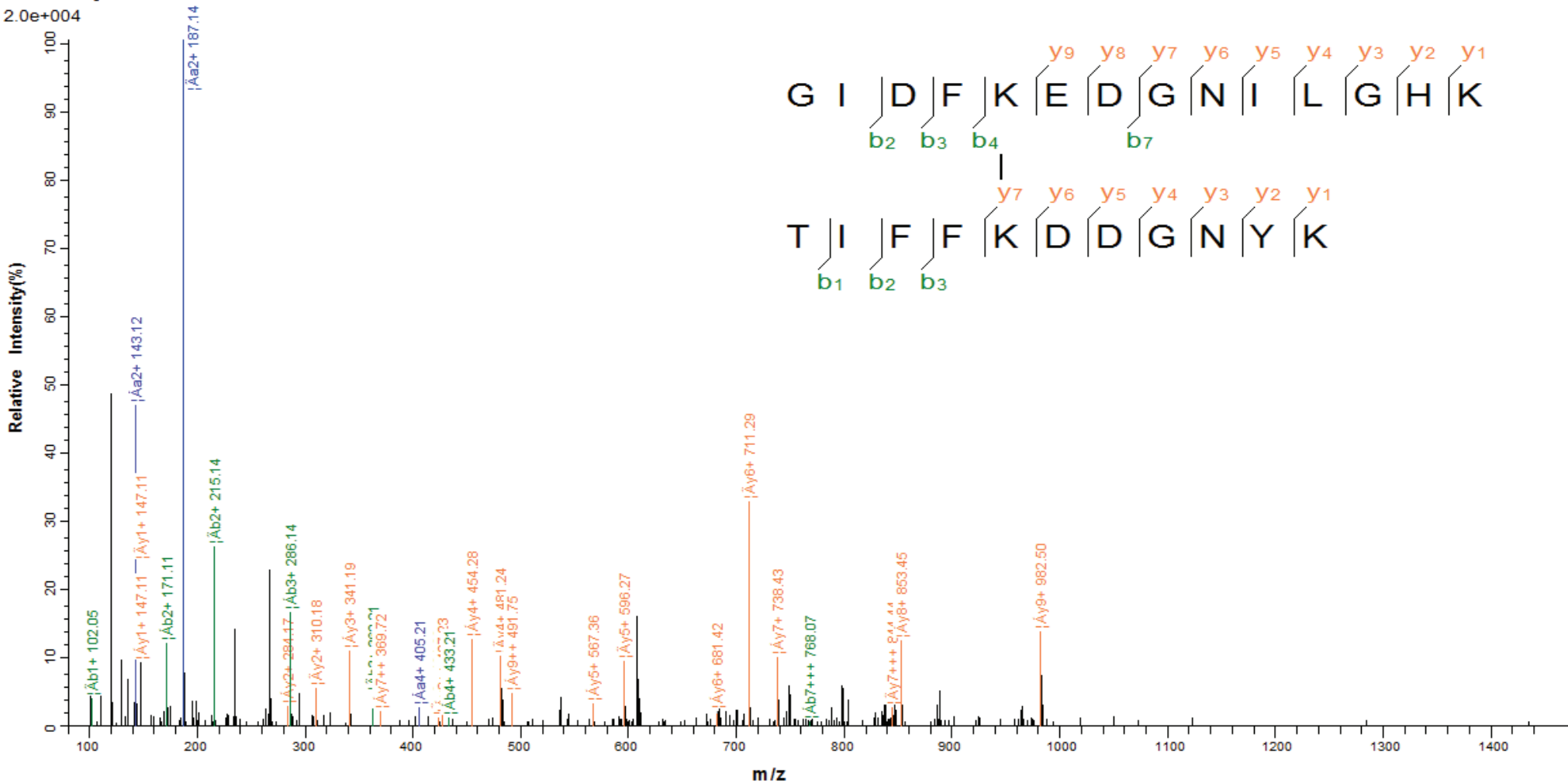
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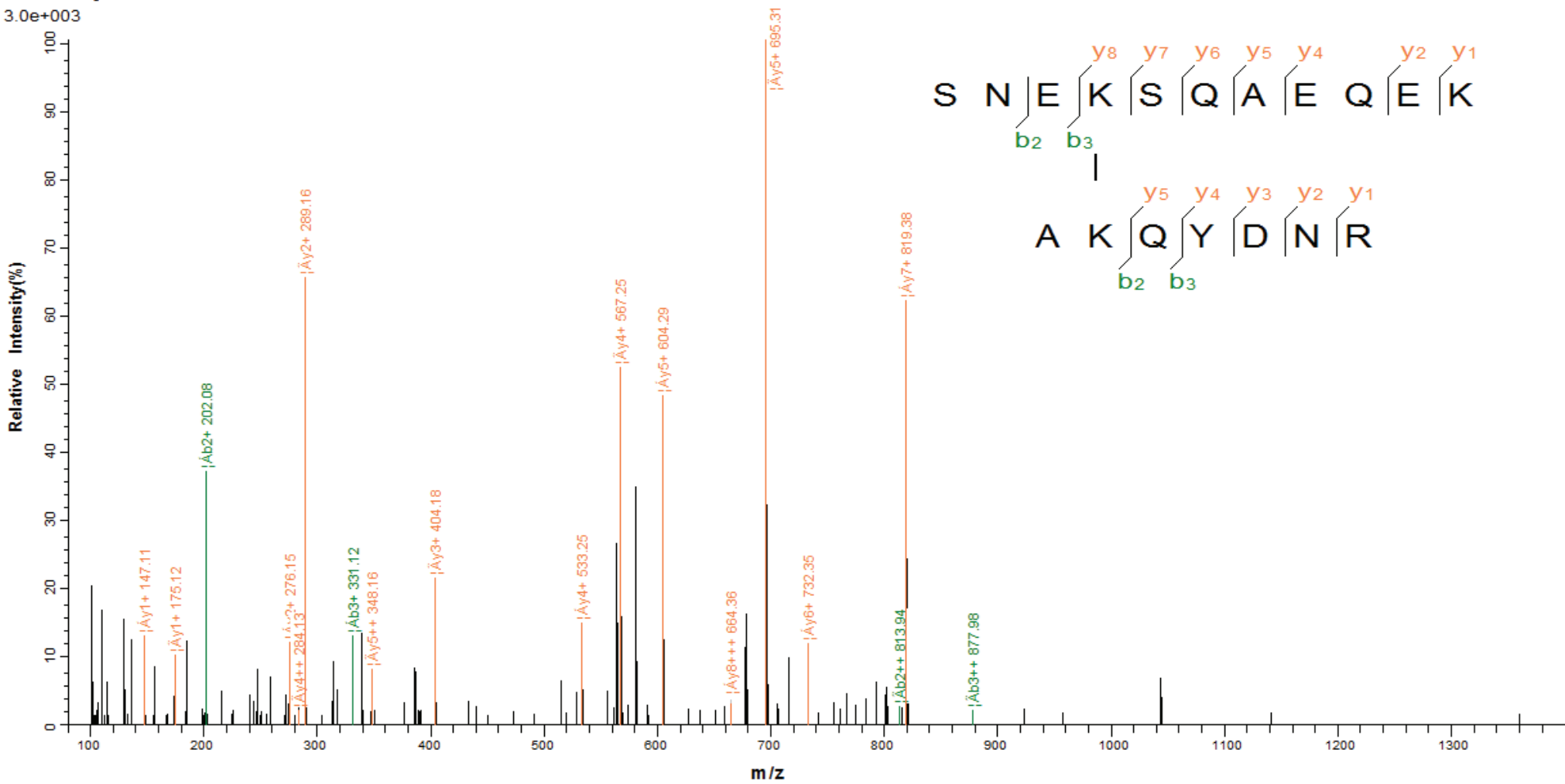
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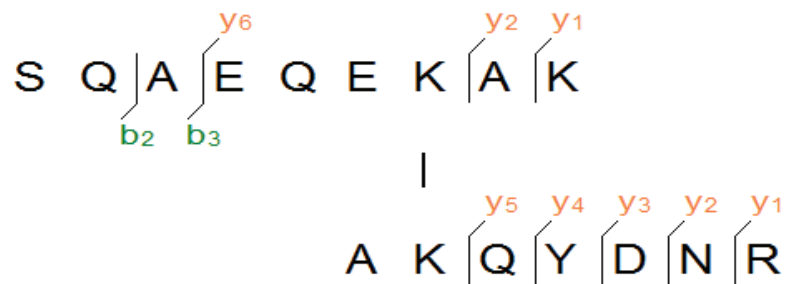
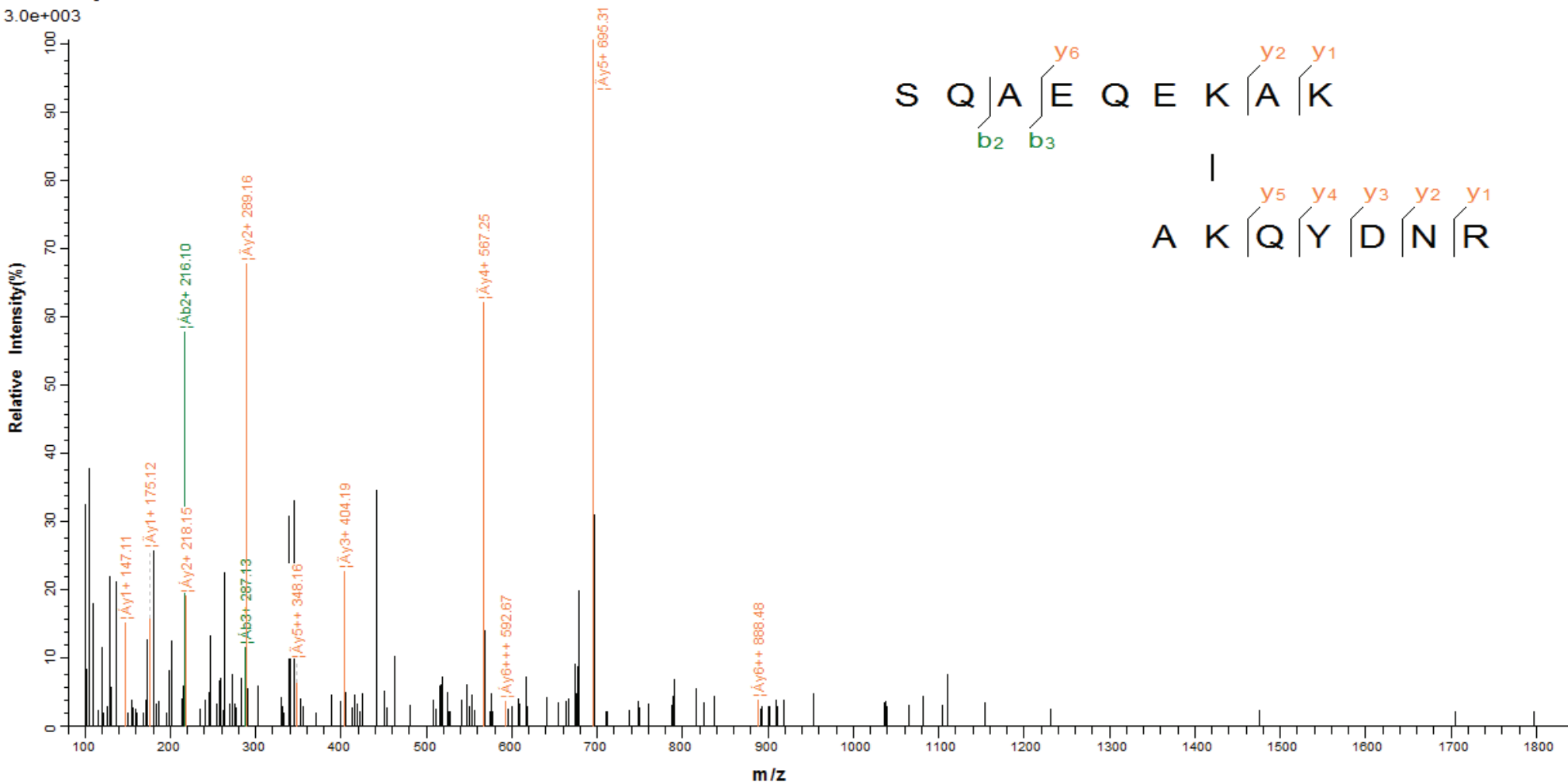
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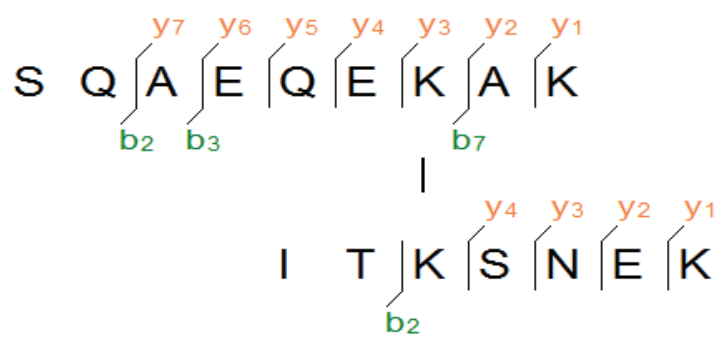
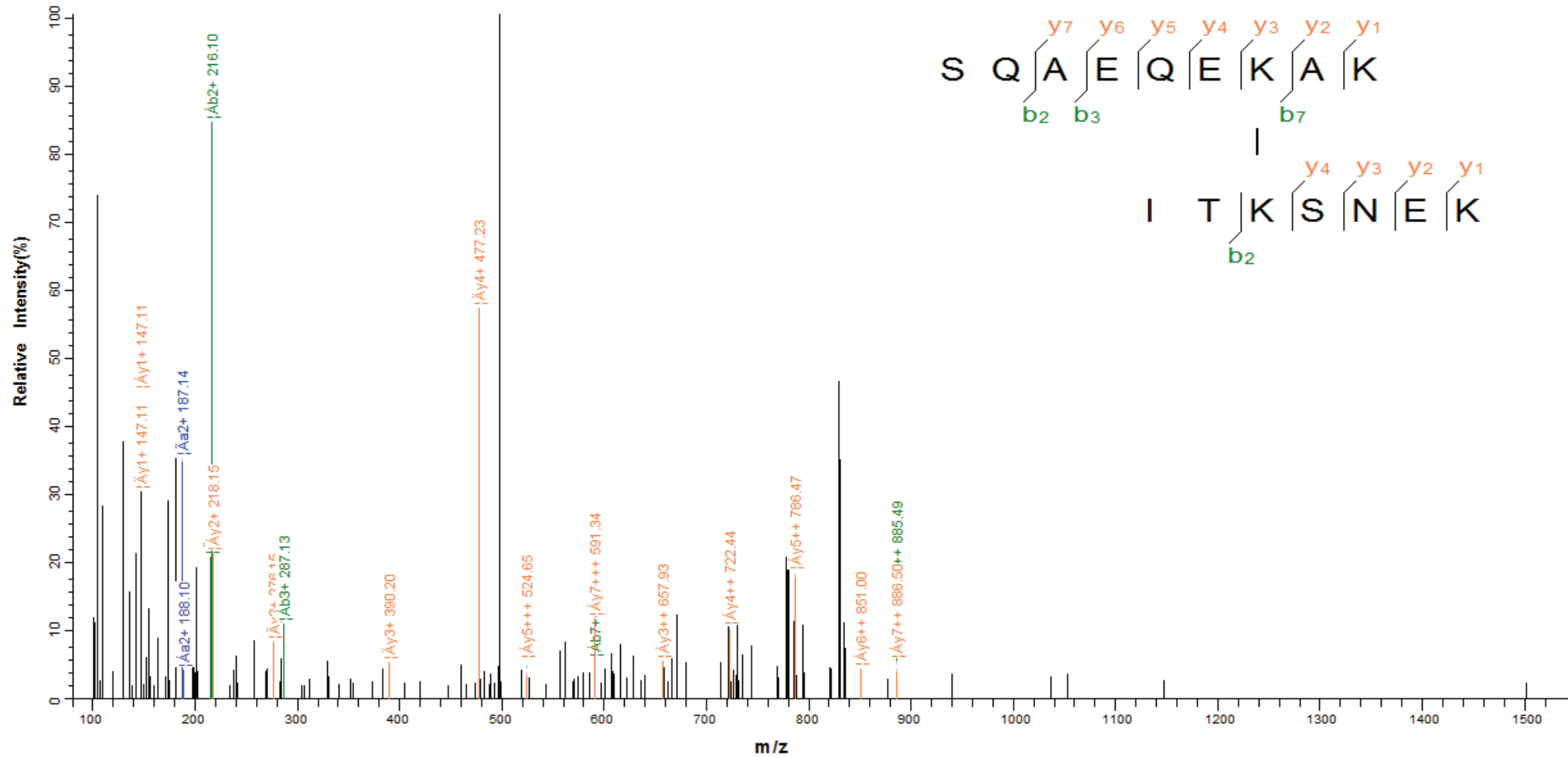


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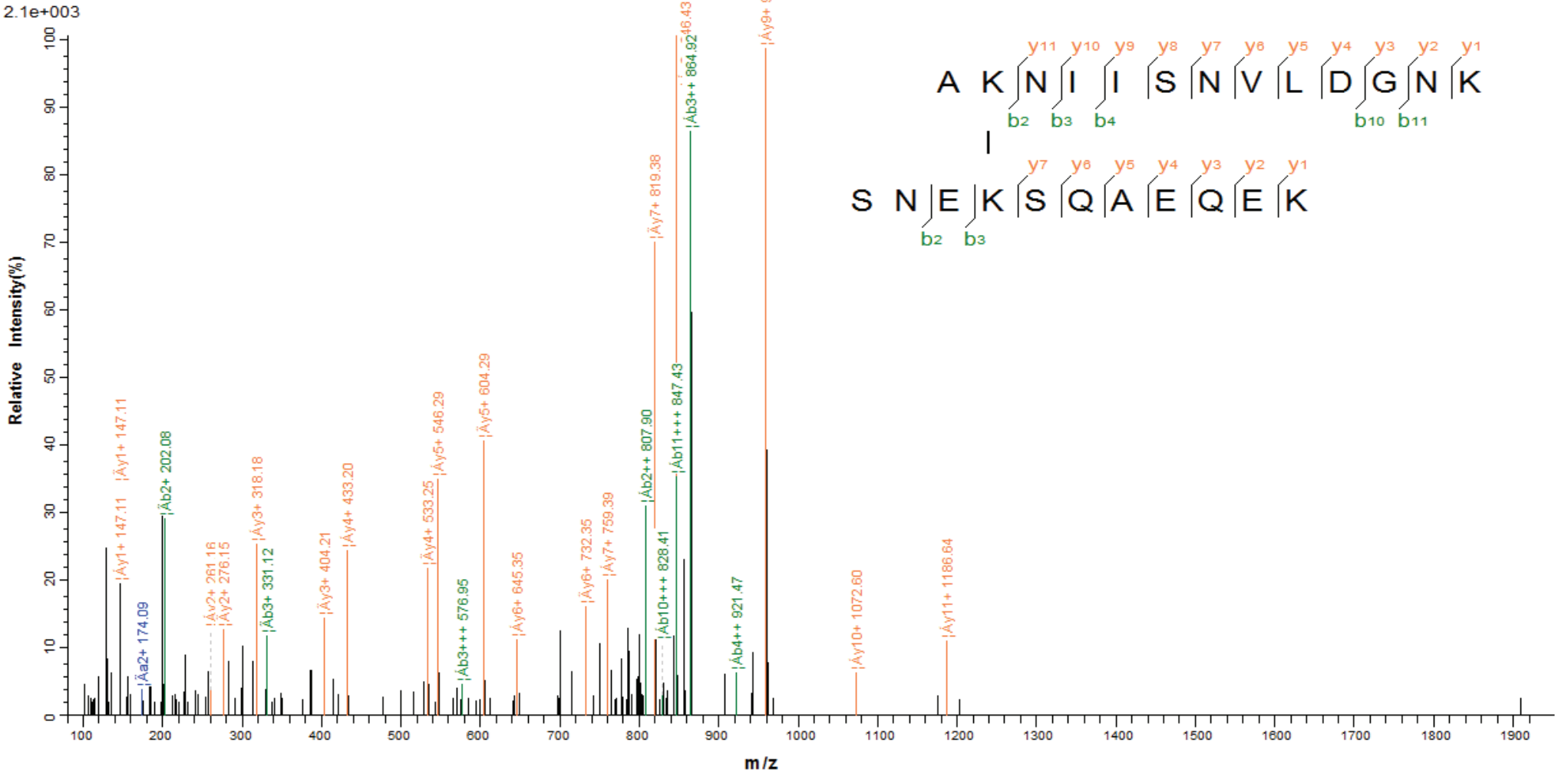


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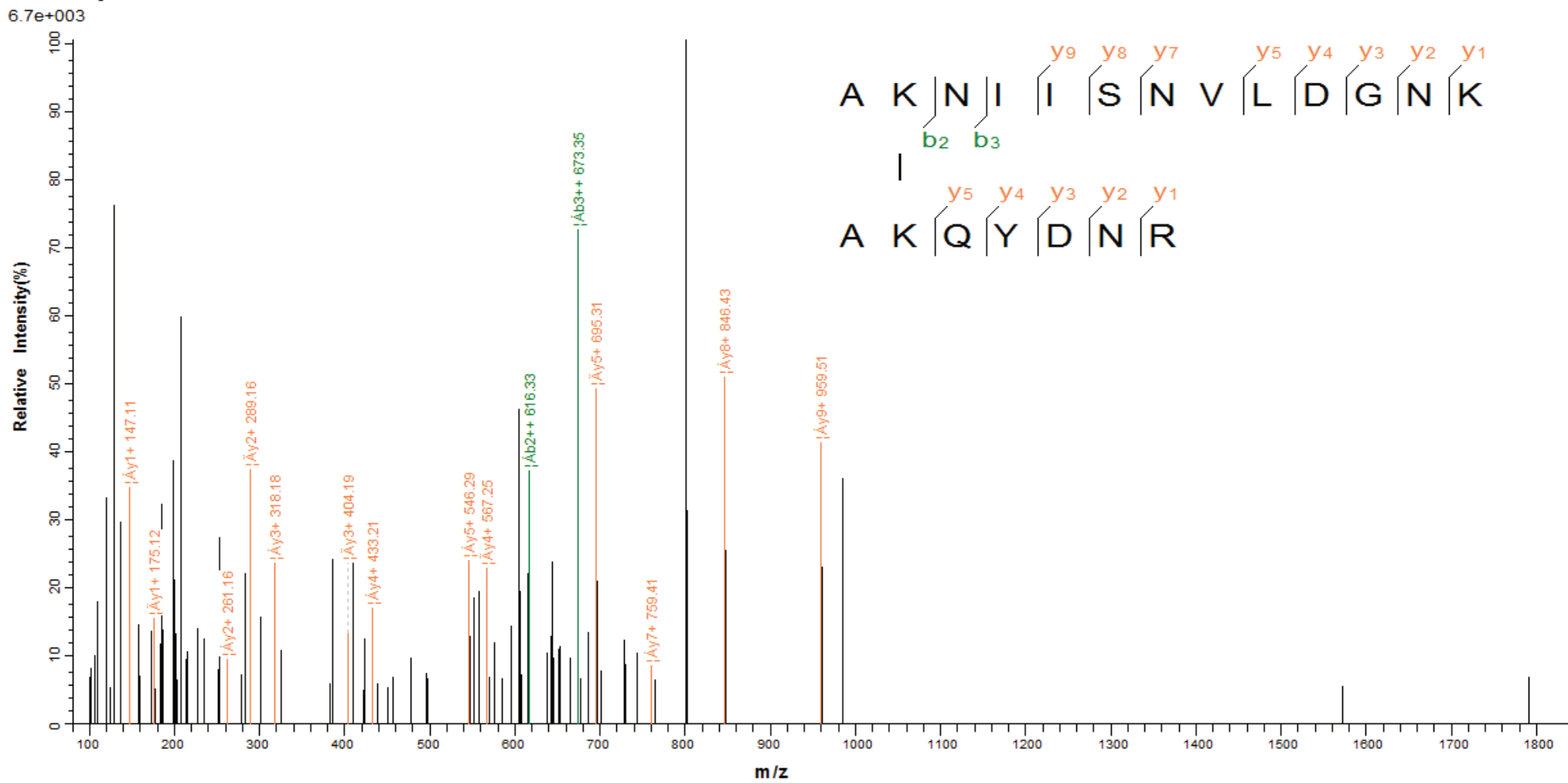
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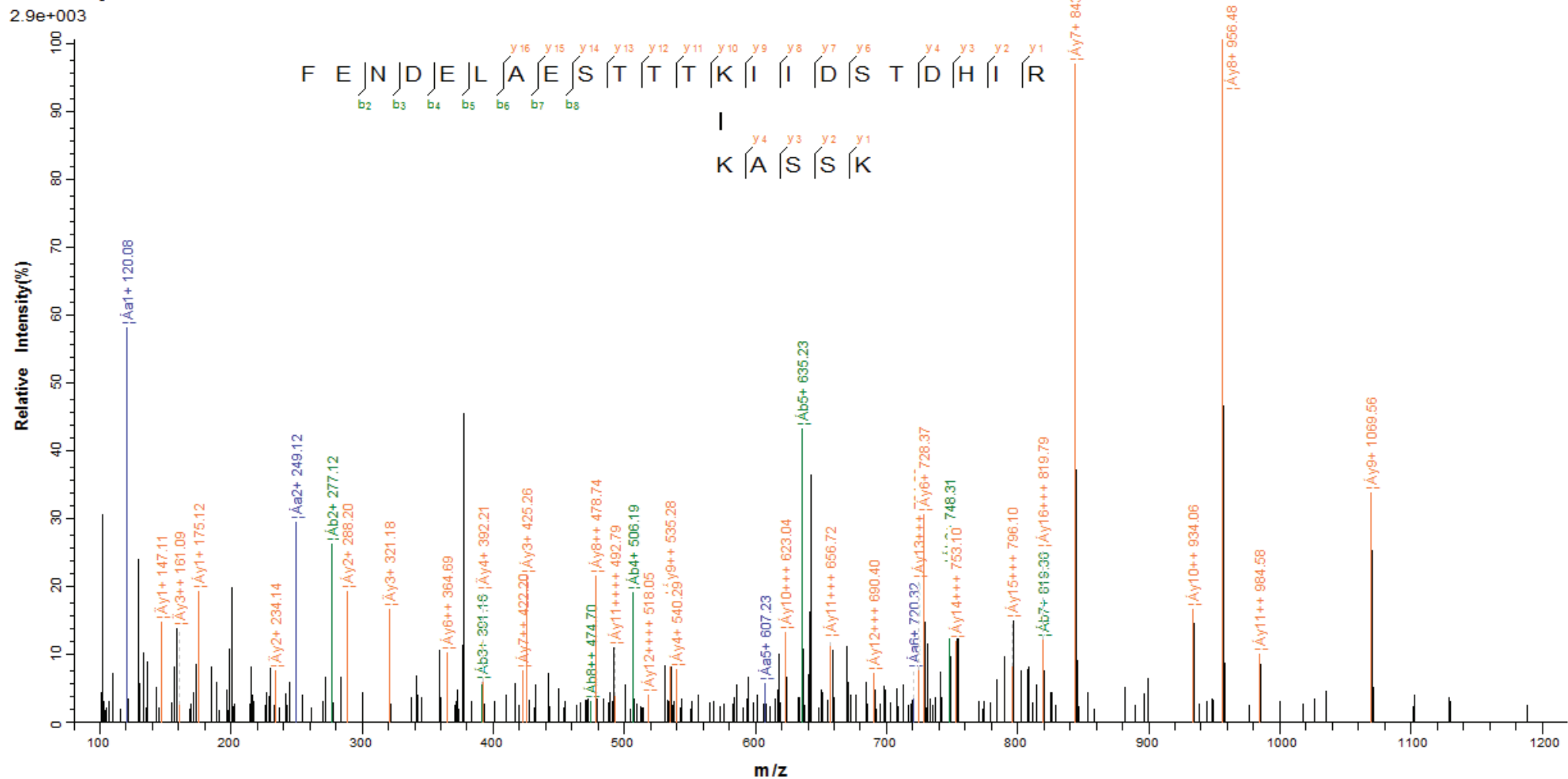
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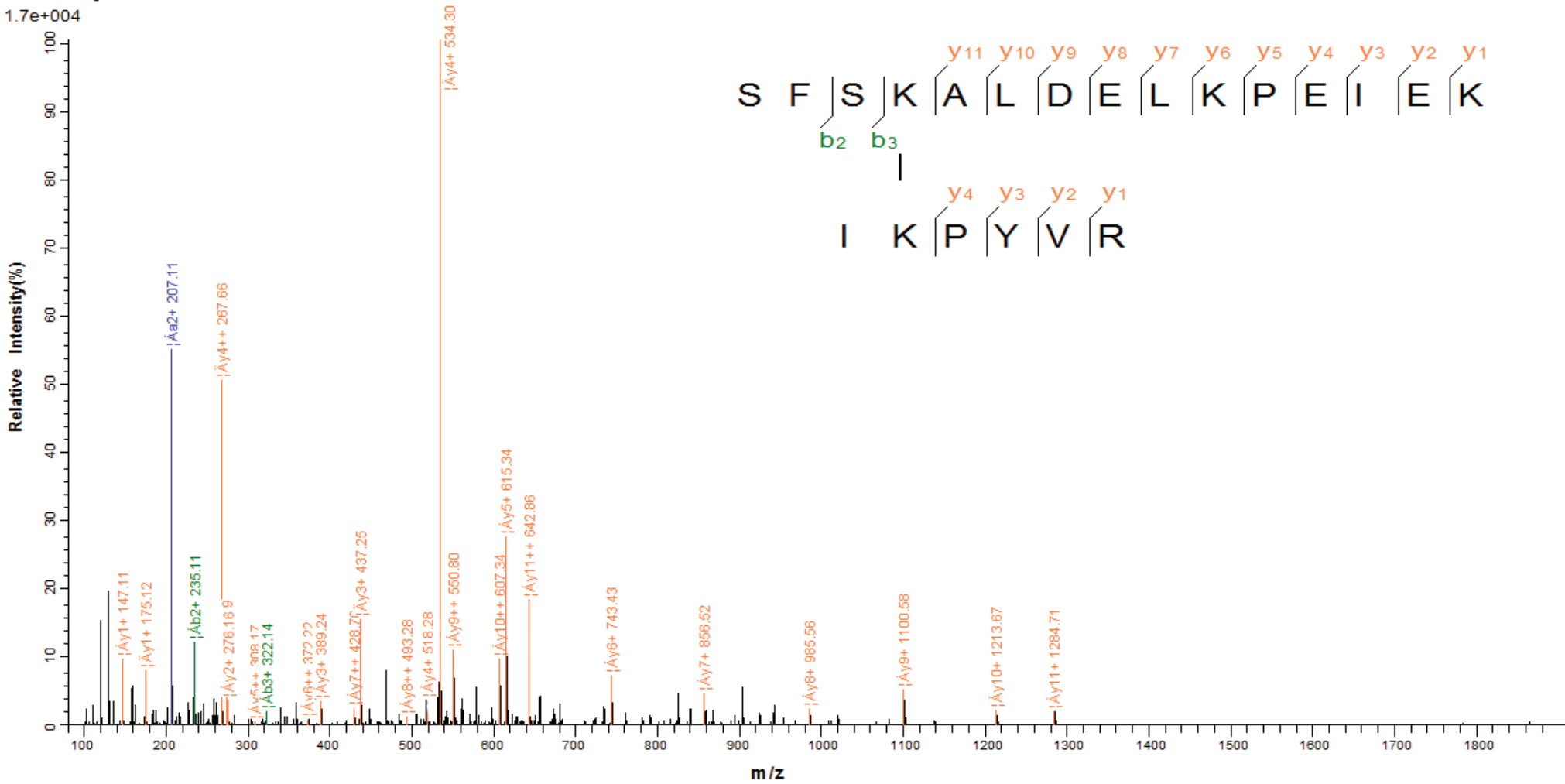
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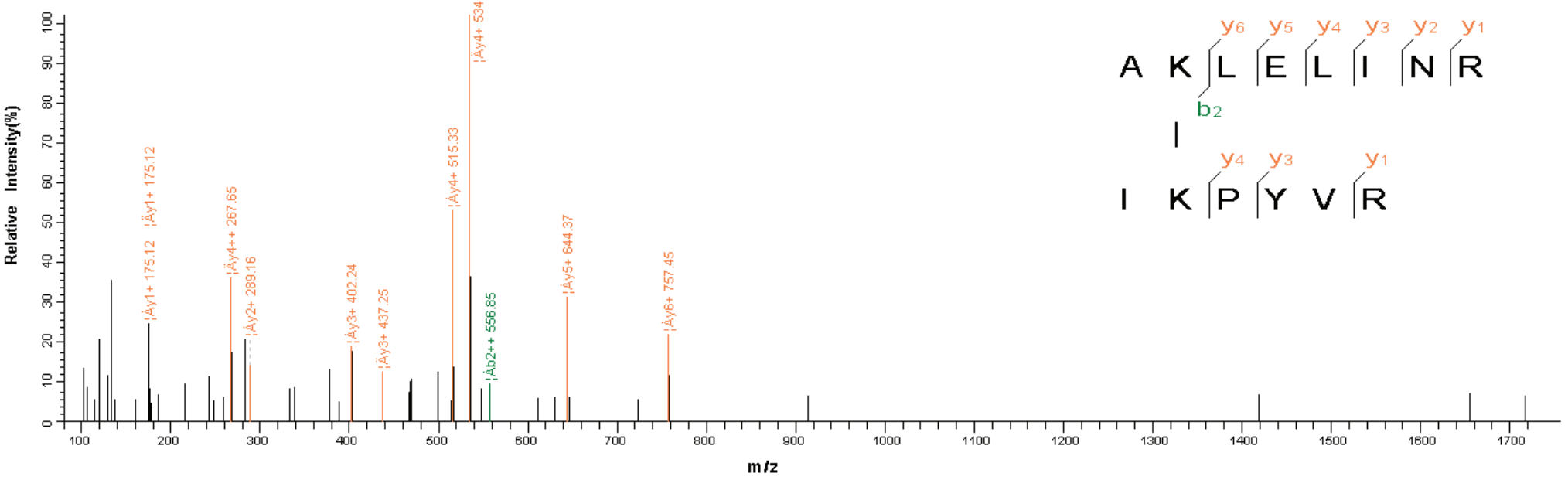
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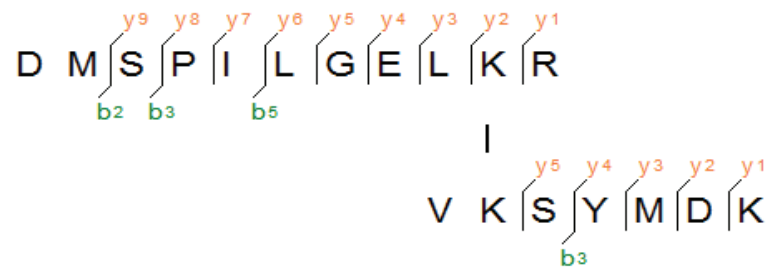
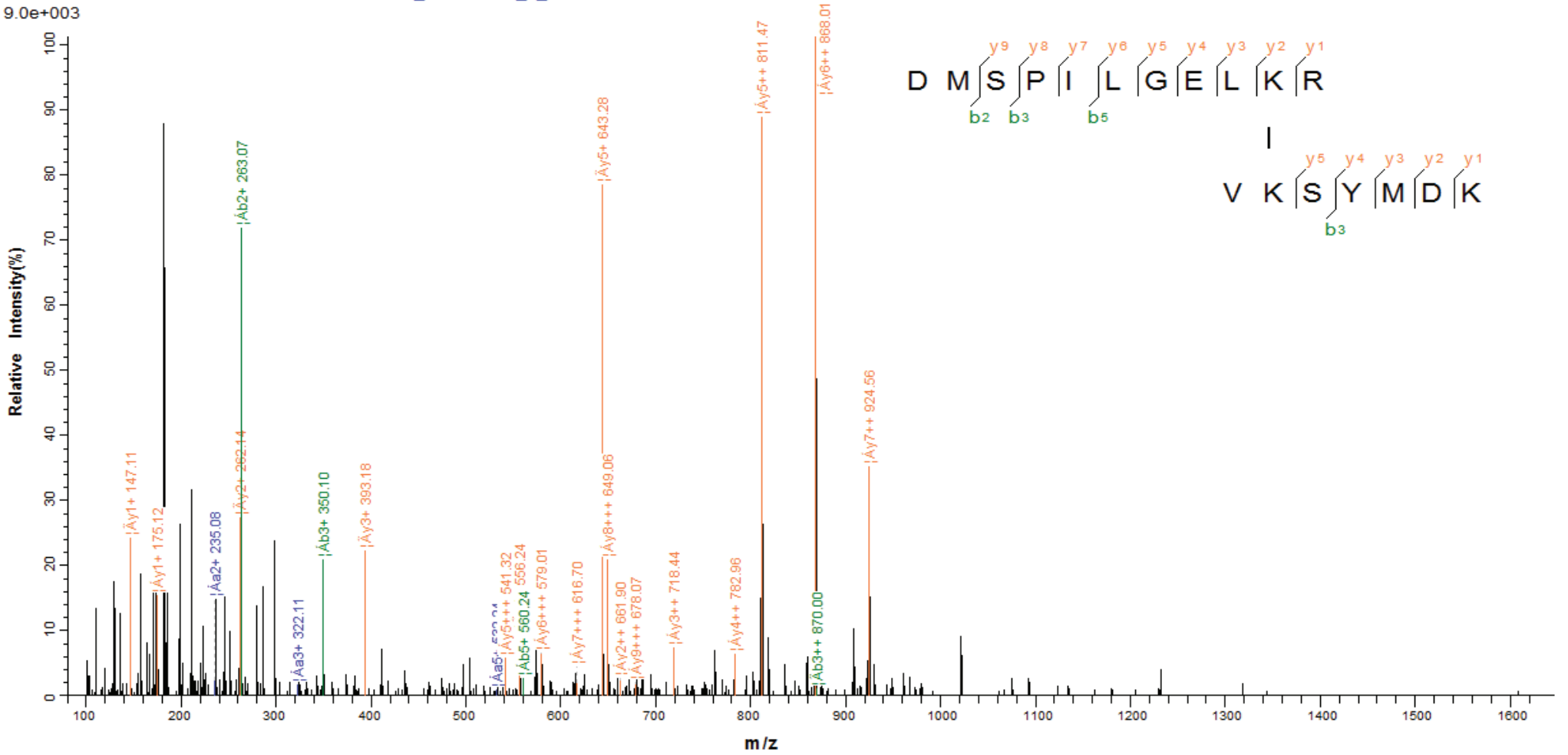
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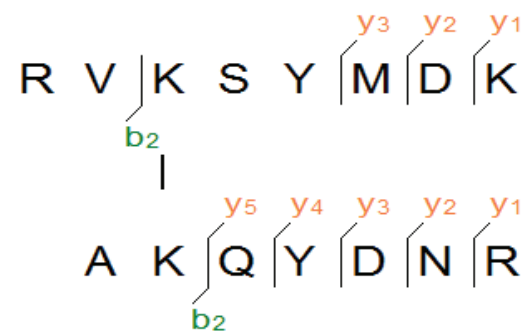
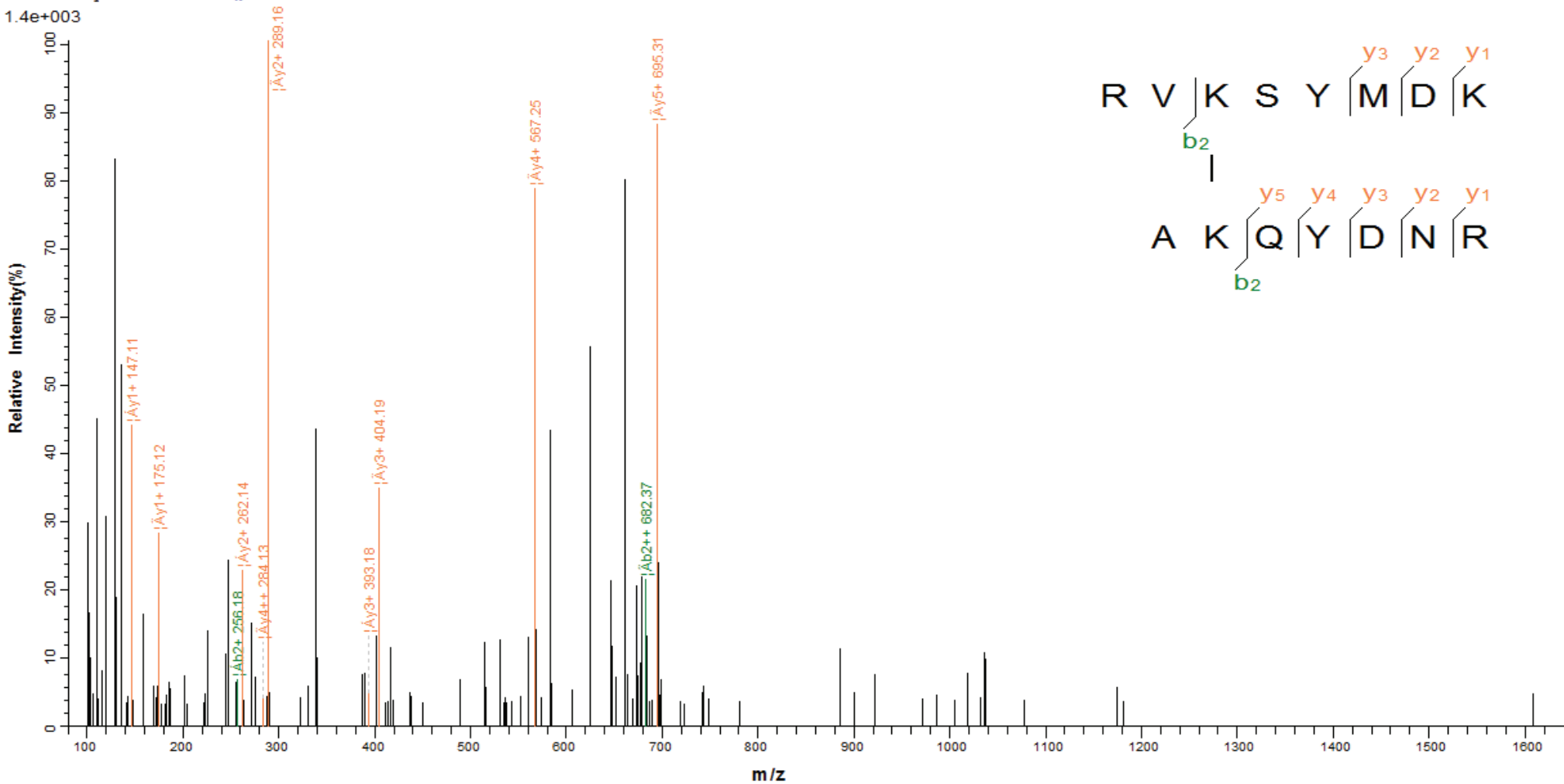
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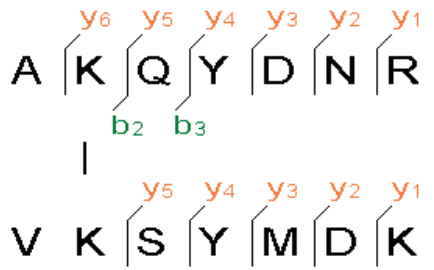
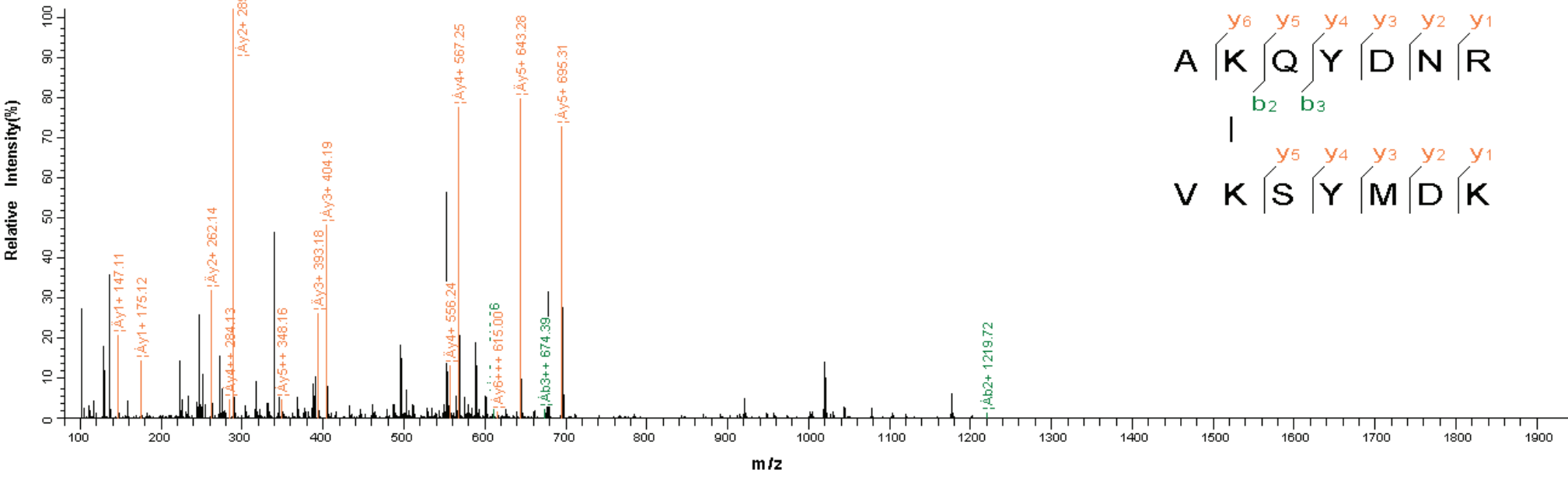
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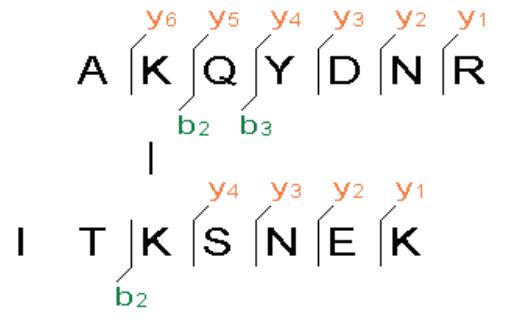
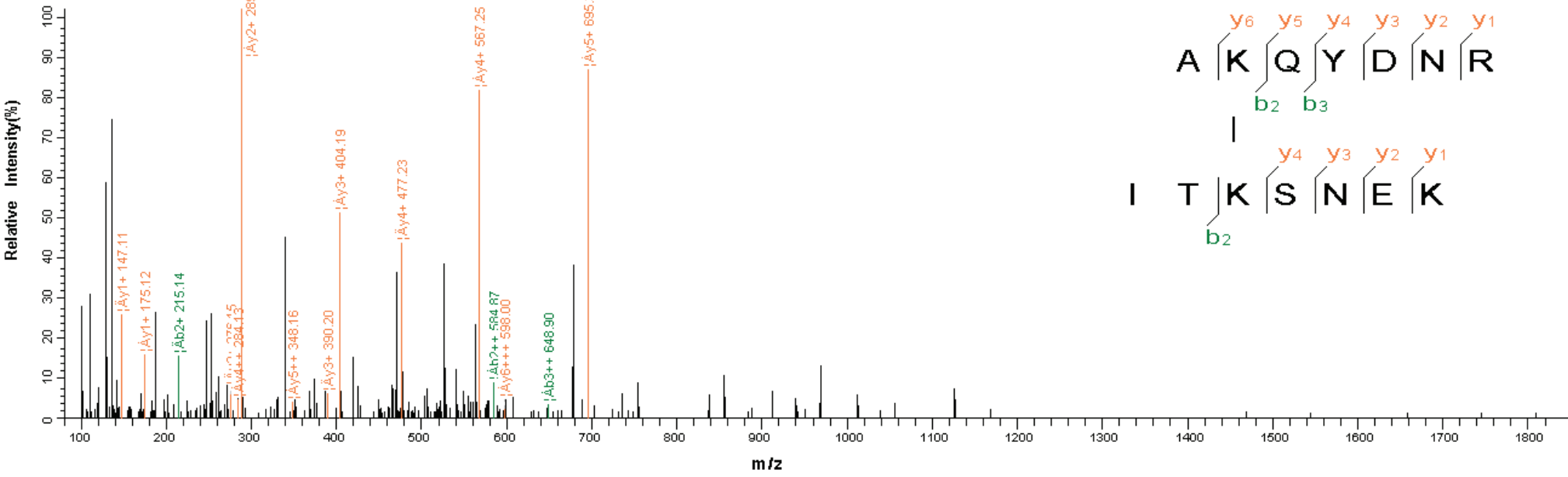
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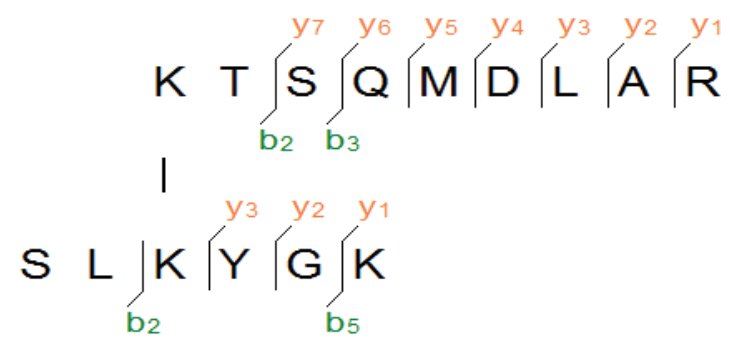
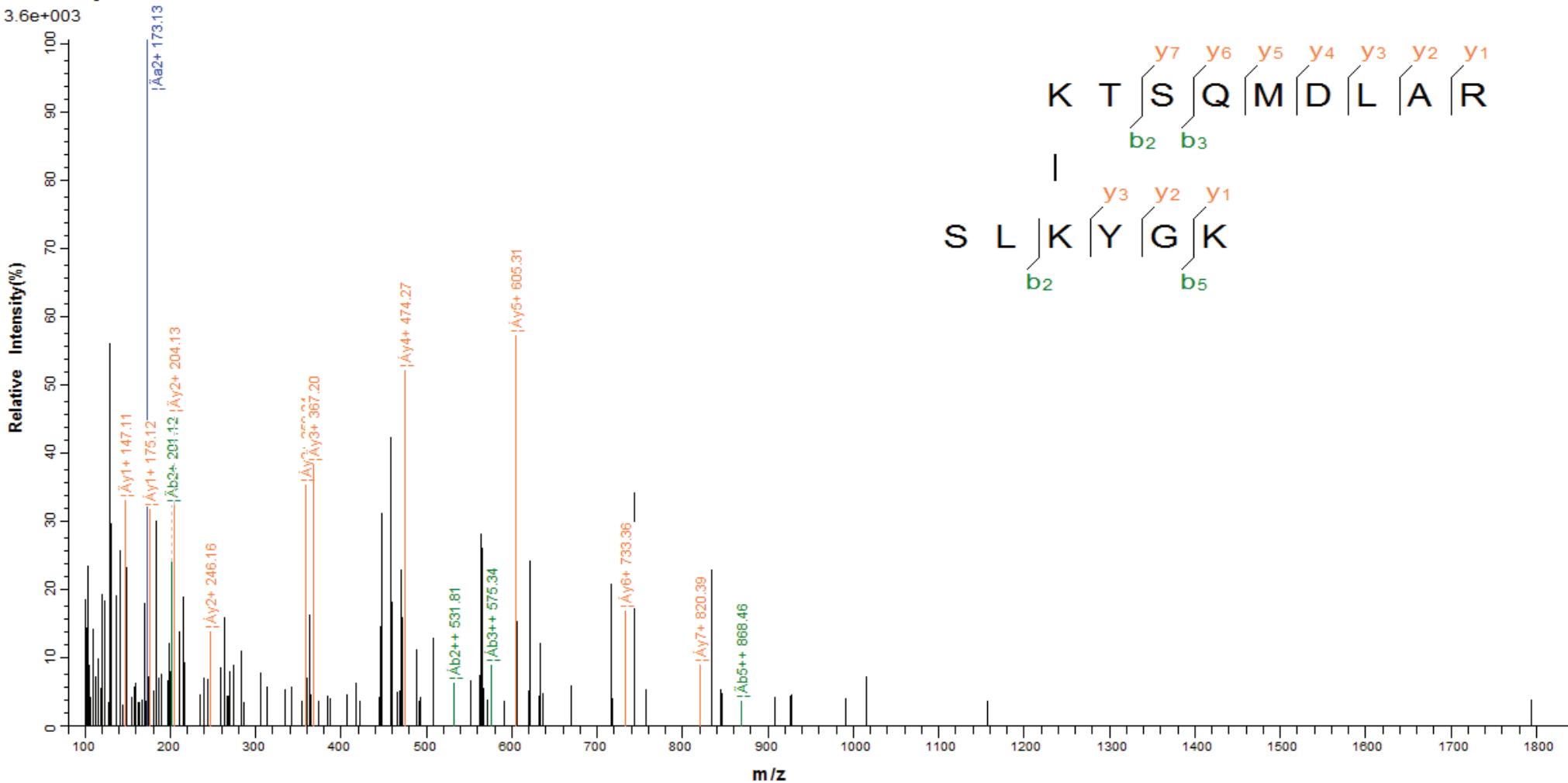
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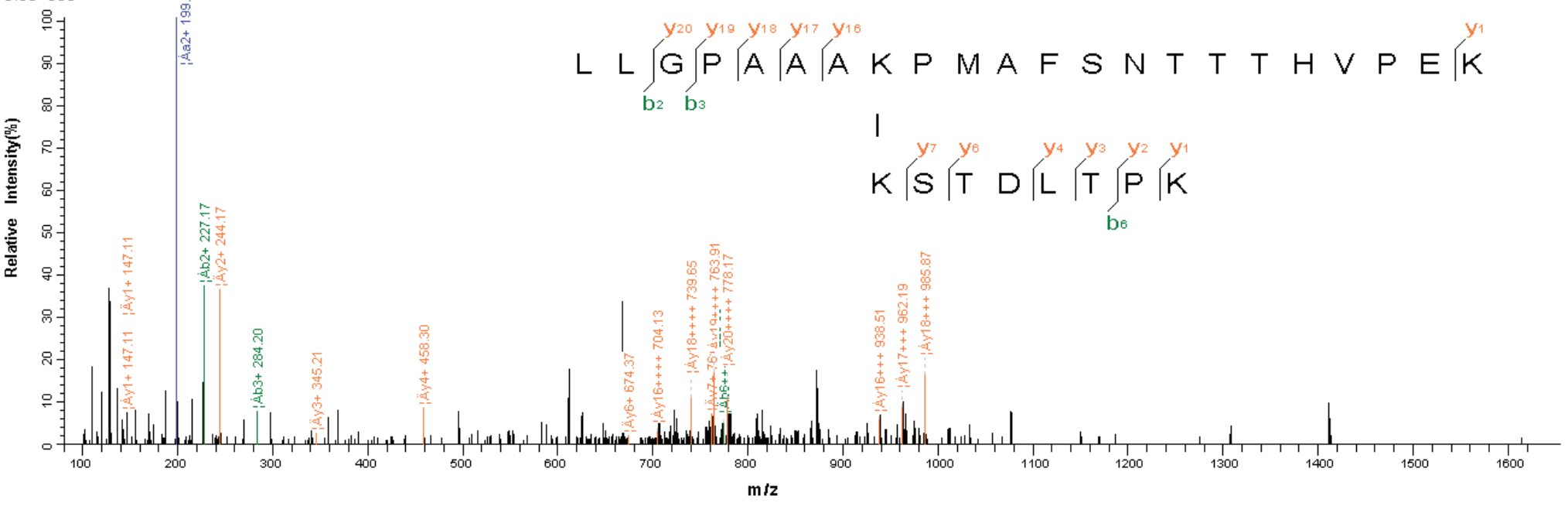
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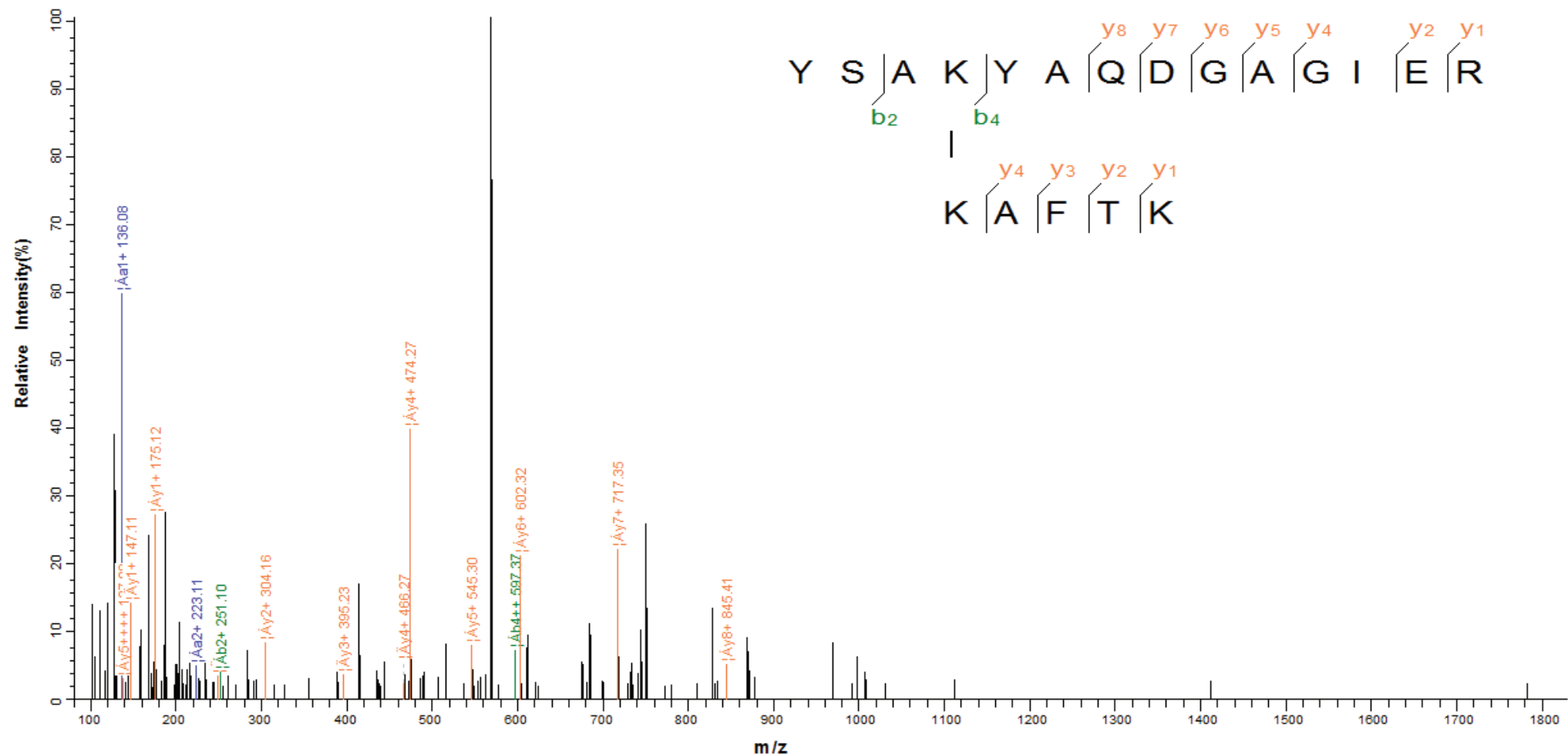


Exosome_GFP_test_A78.002101.002101.5.dta Seq: LLGPAAAKPMAFSNTTTHVPEK-KSTDLTPK Mod: 10,_Oxidation_M_15.995

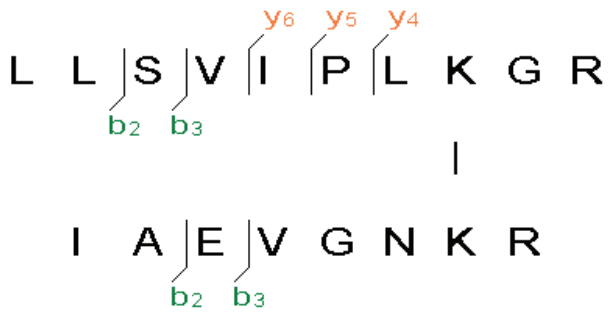
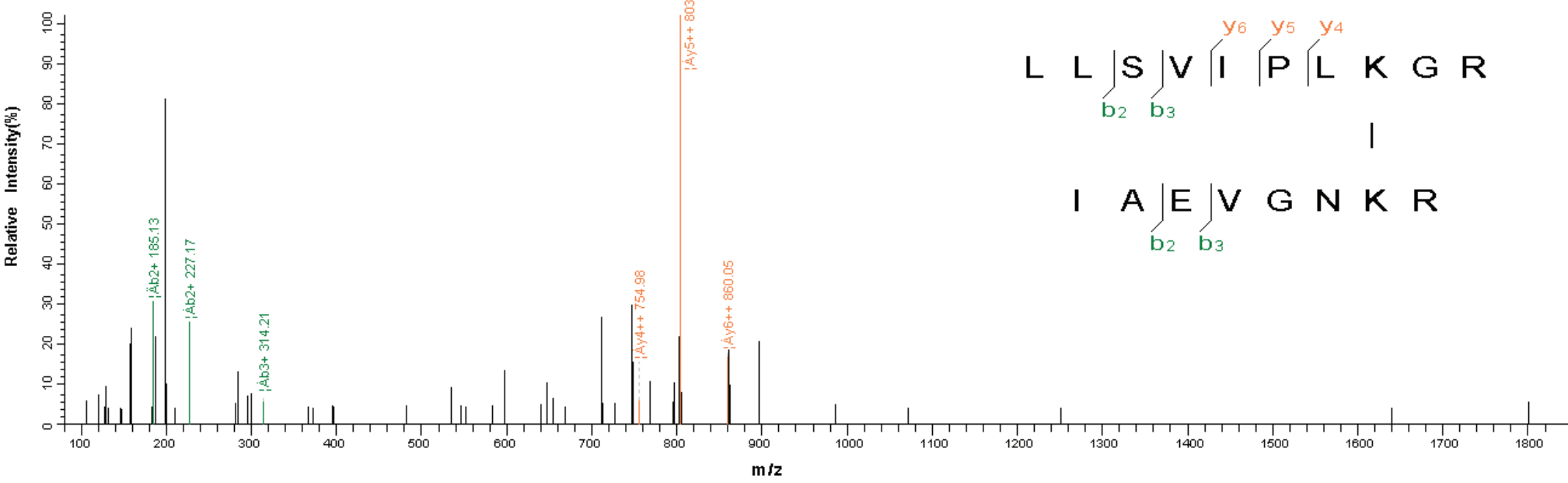


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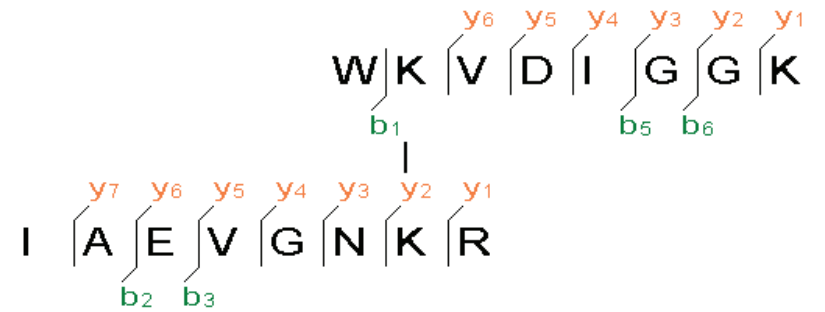
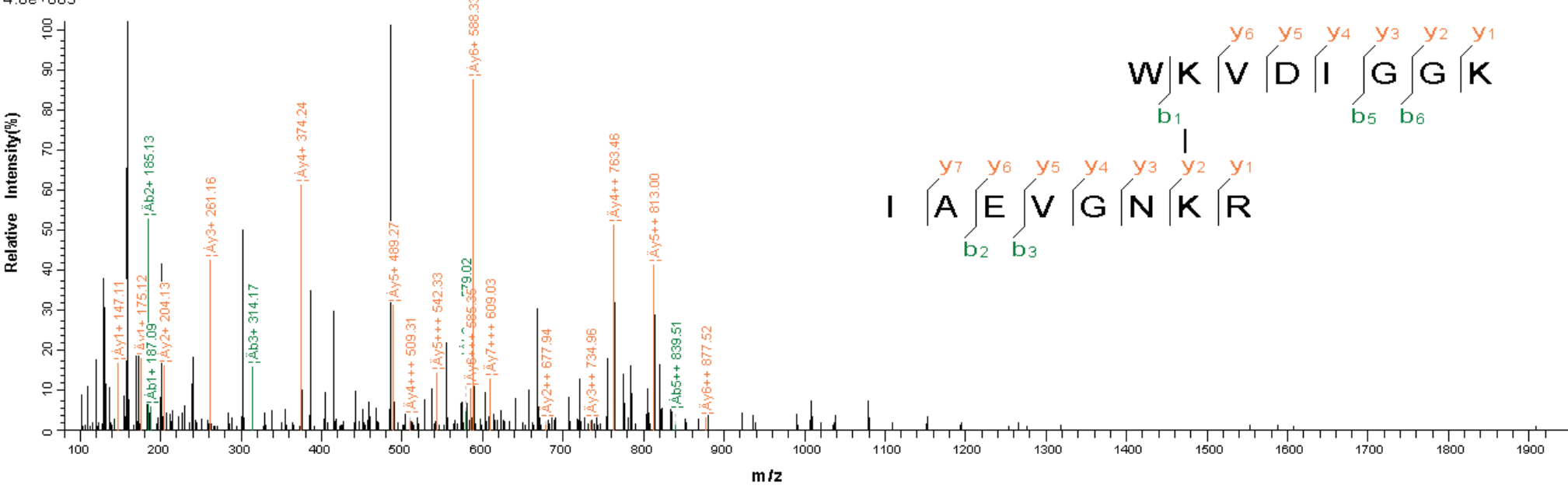
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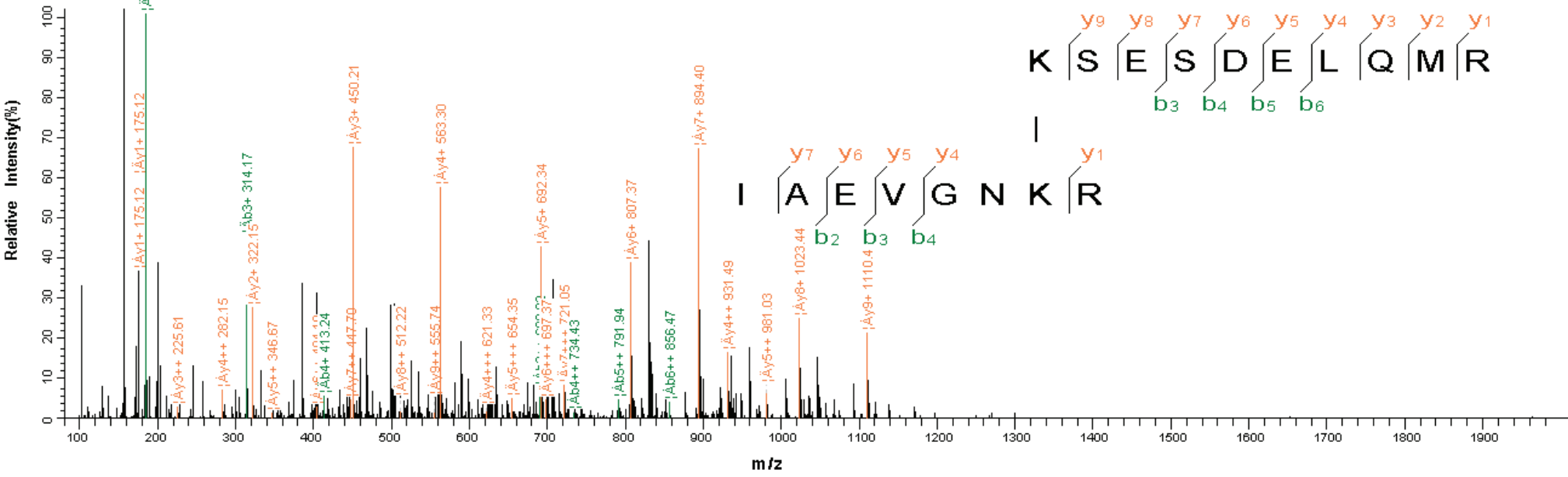
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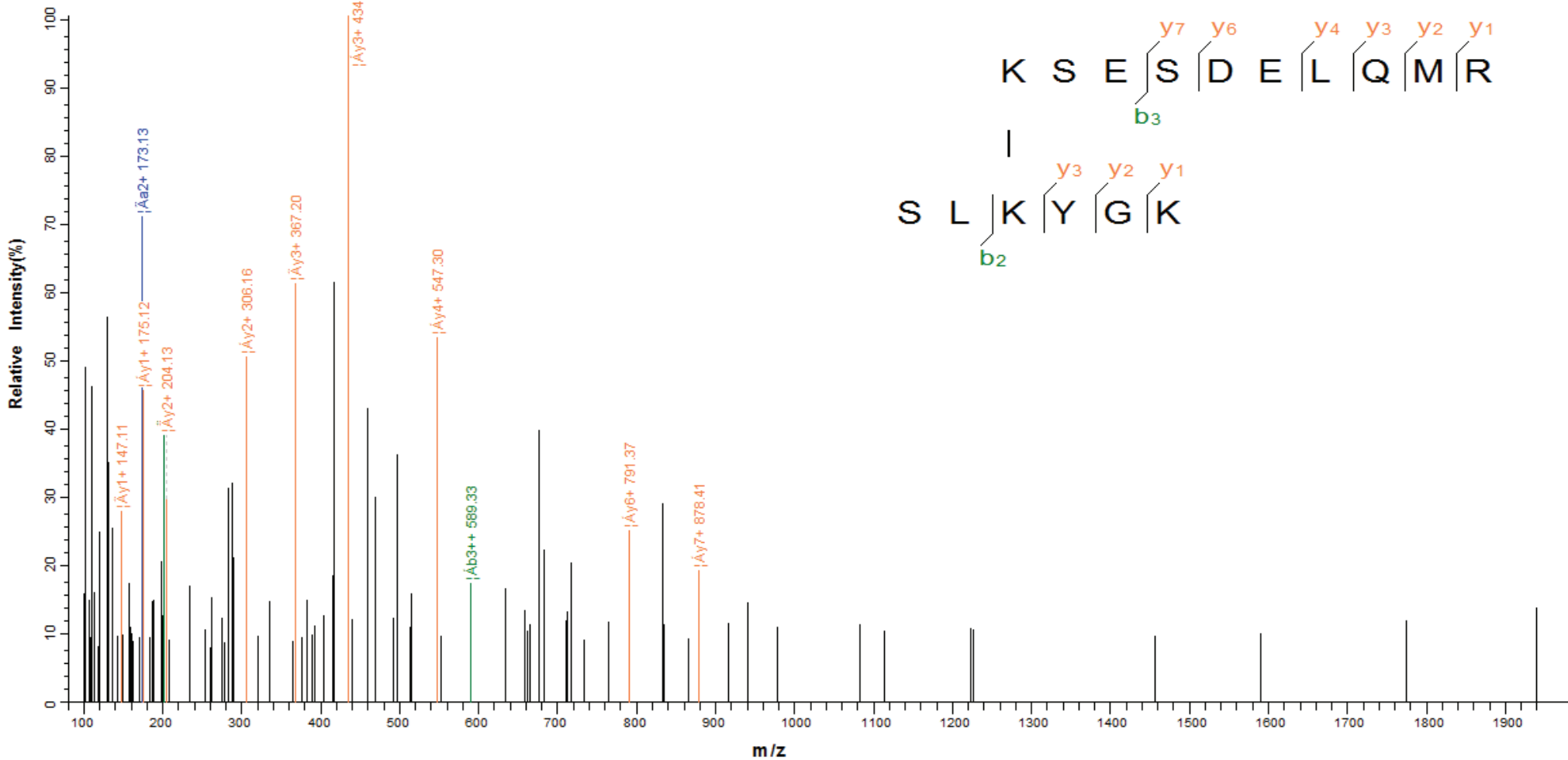


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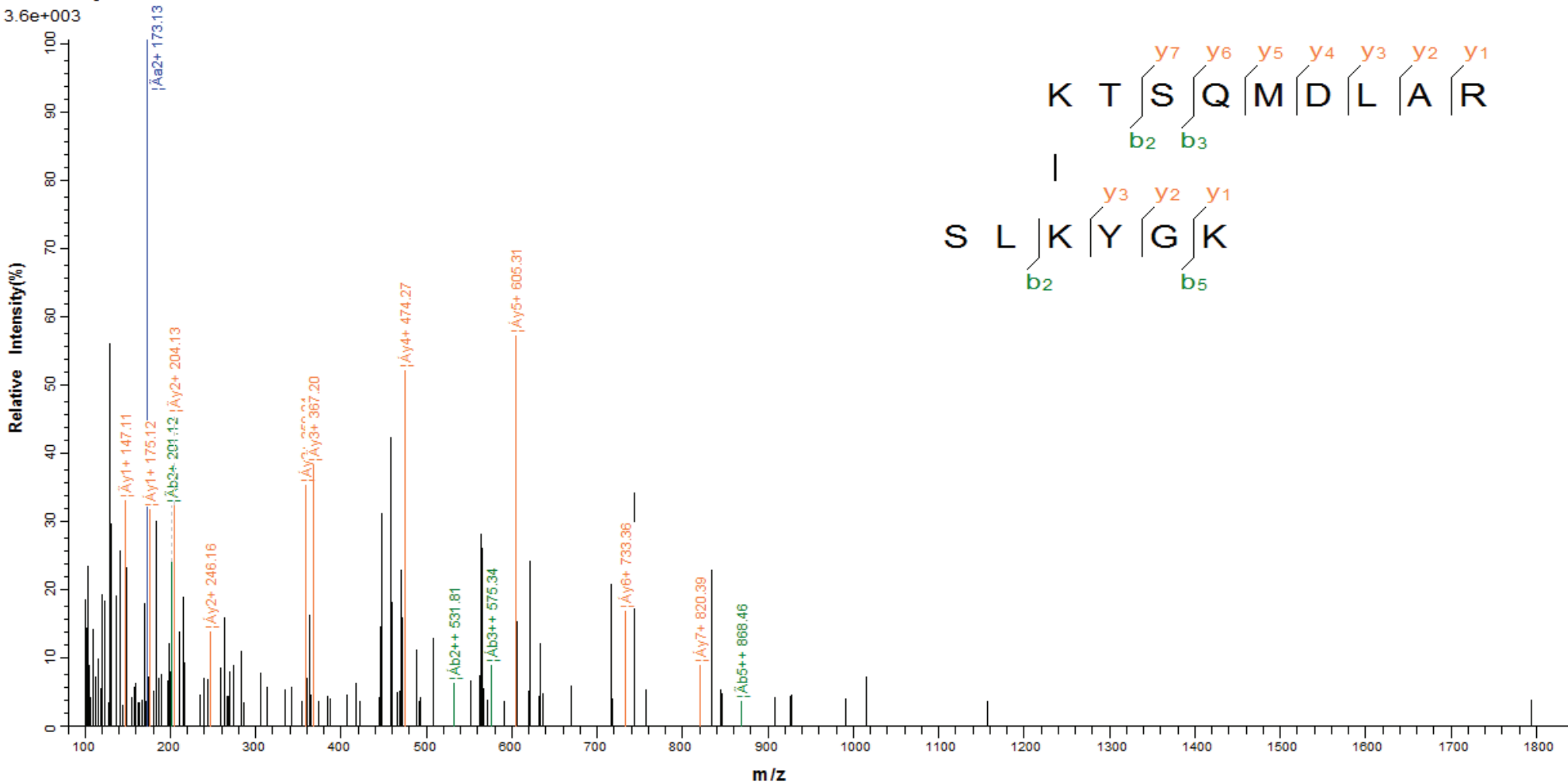


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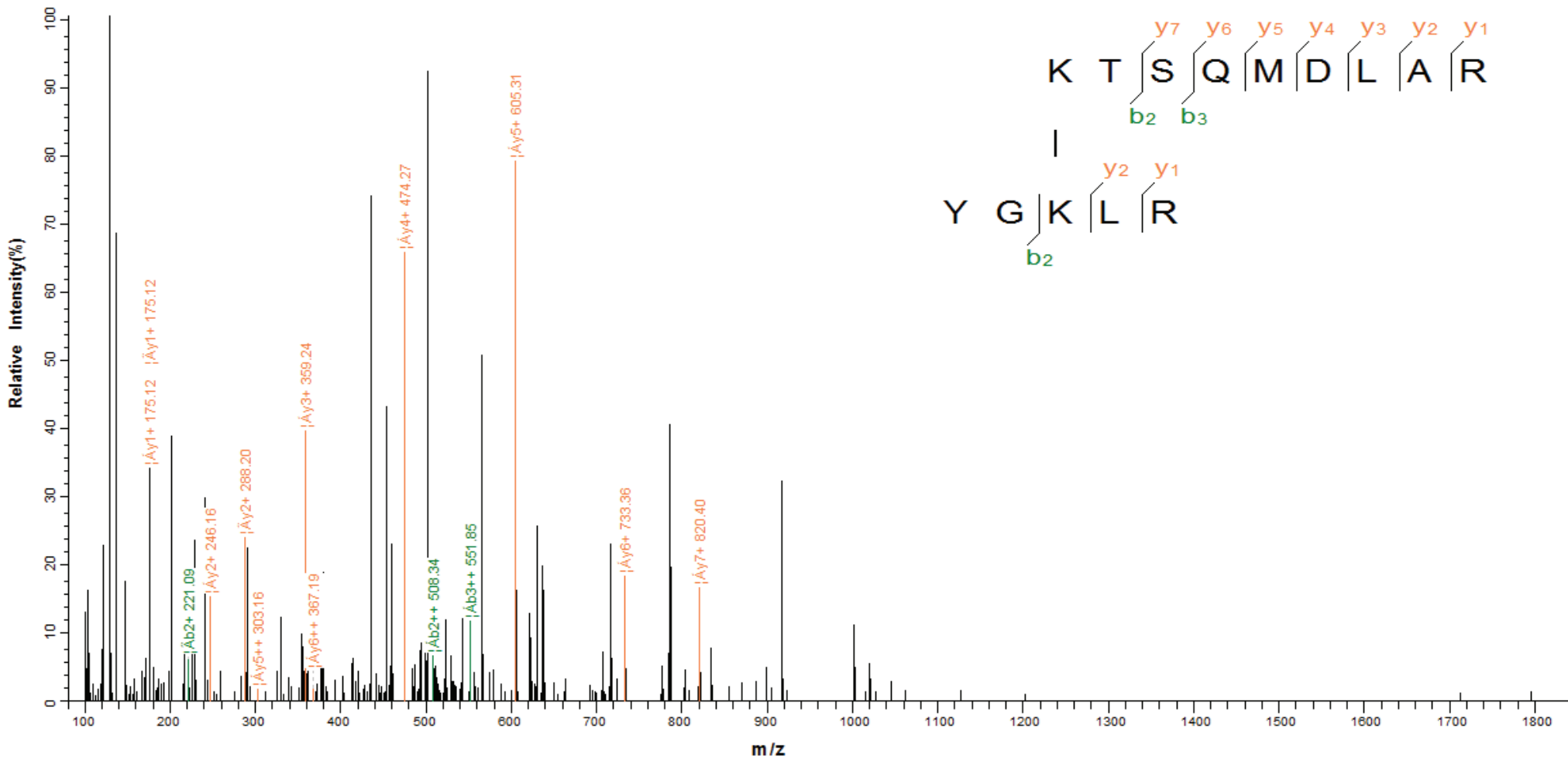


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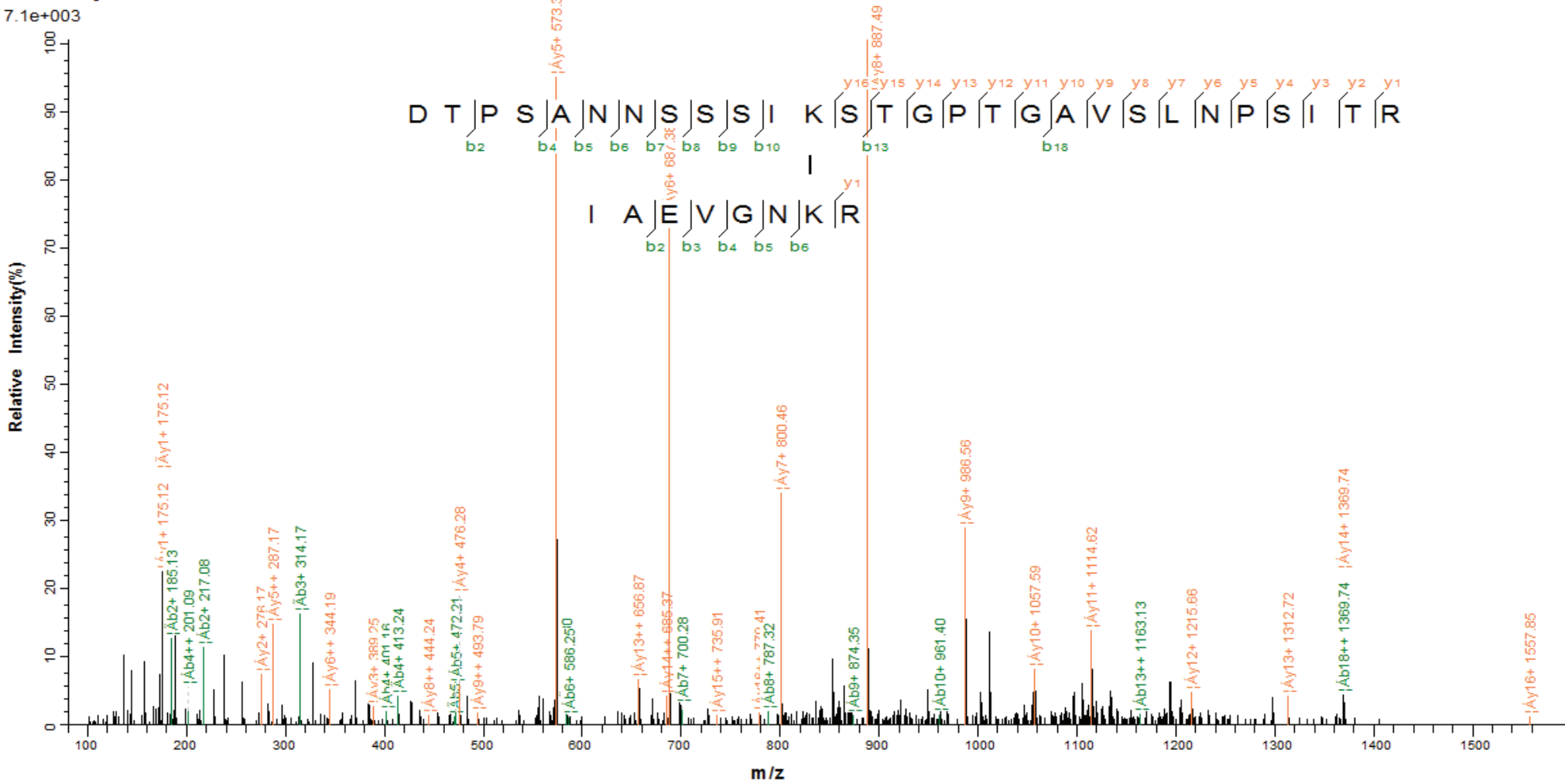


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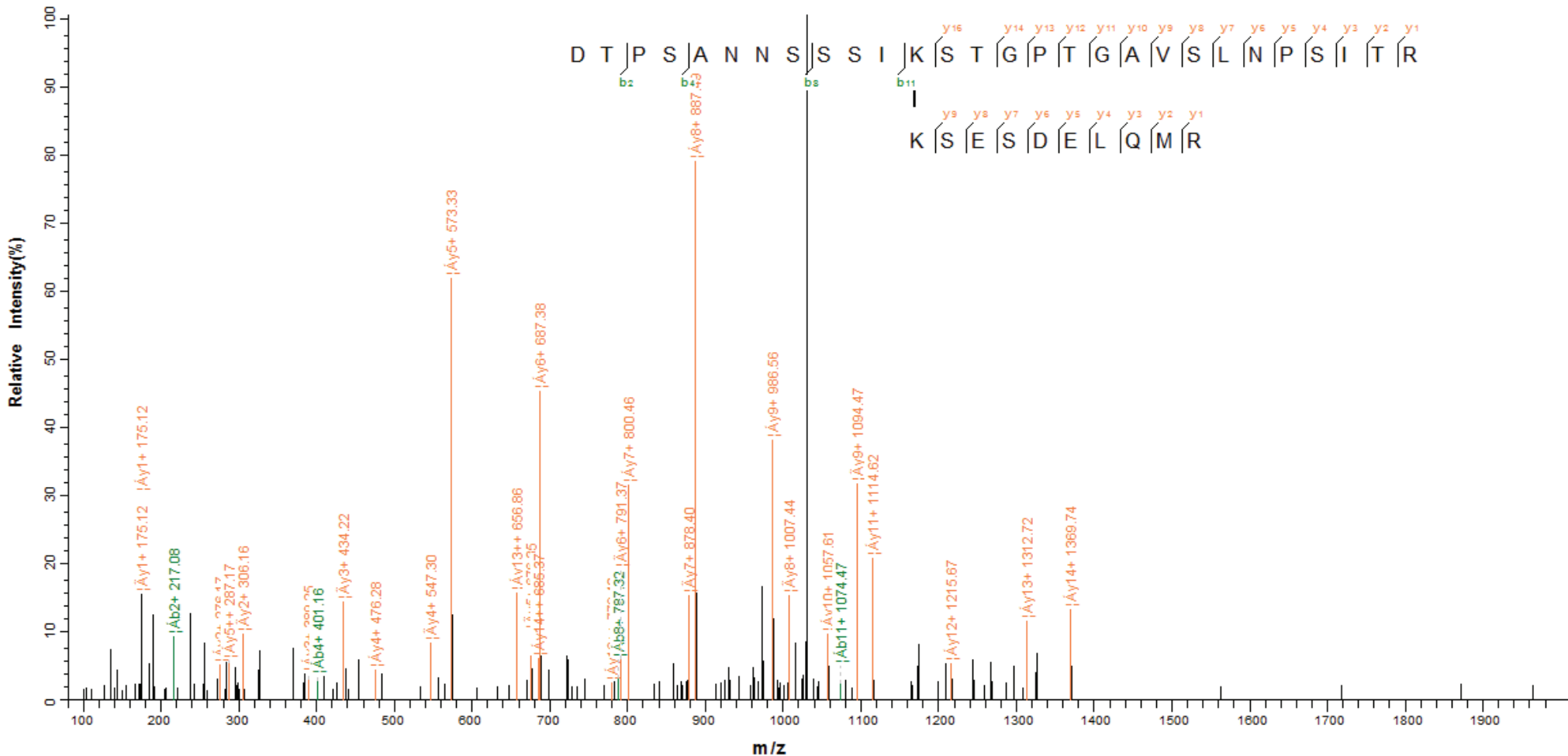


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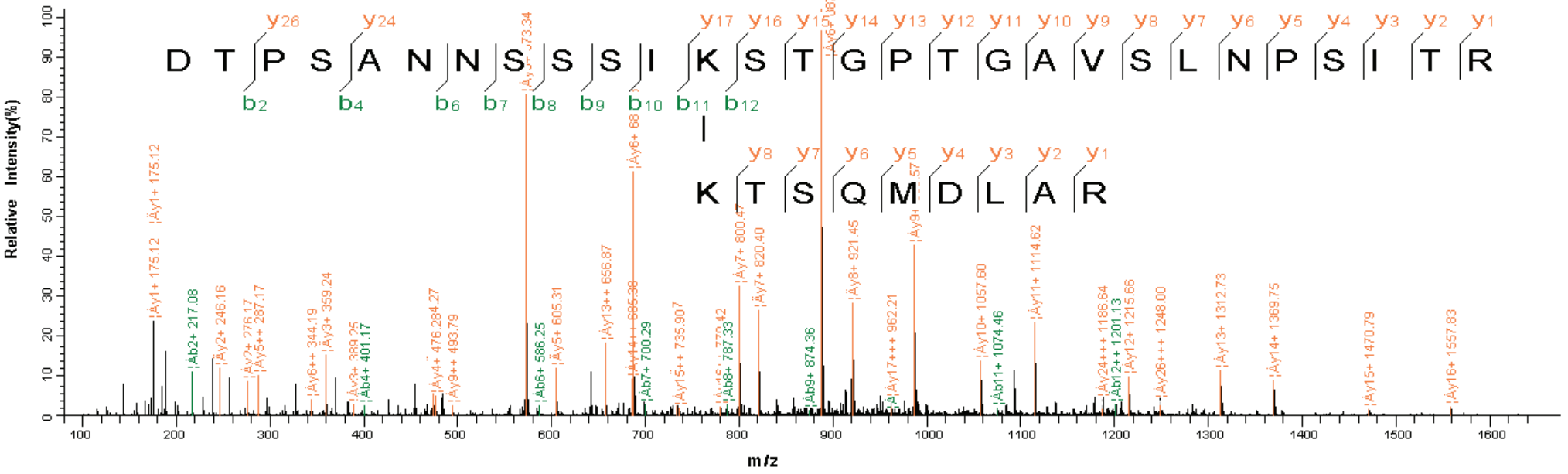


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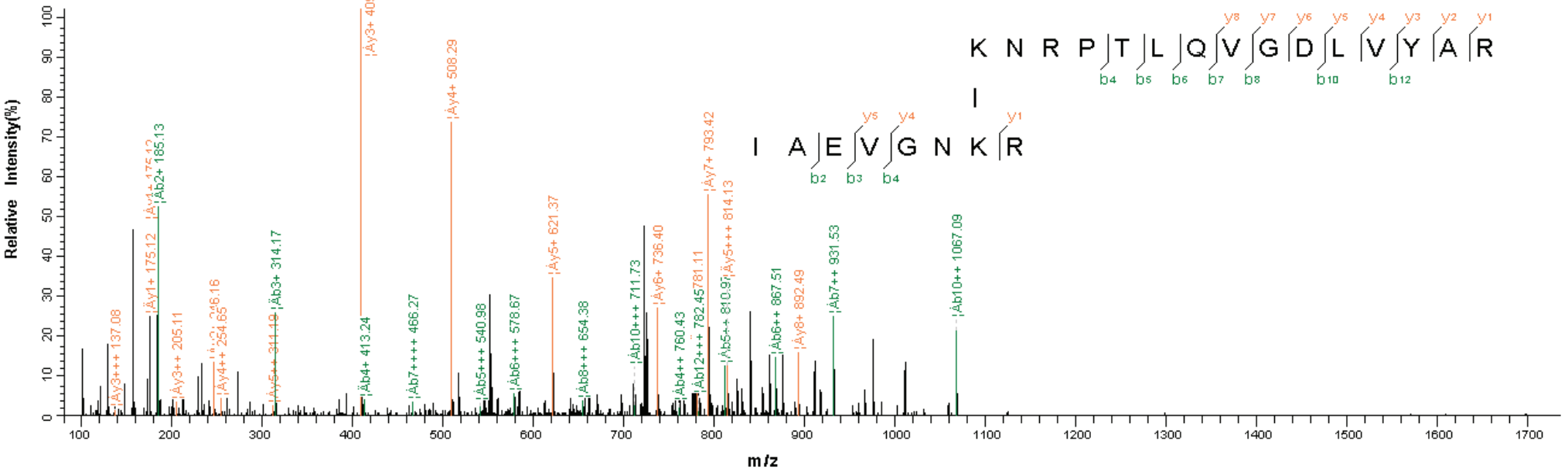
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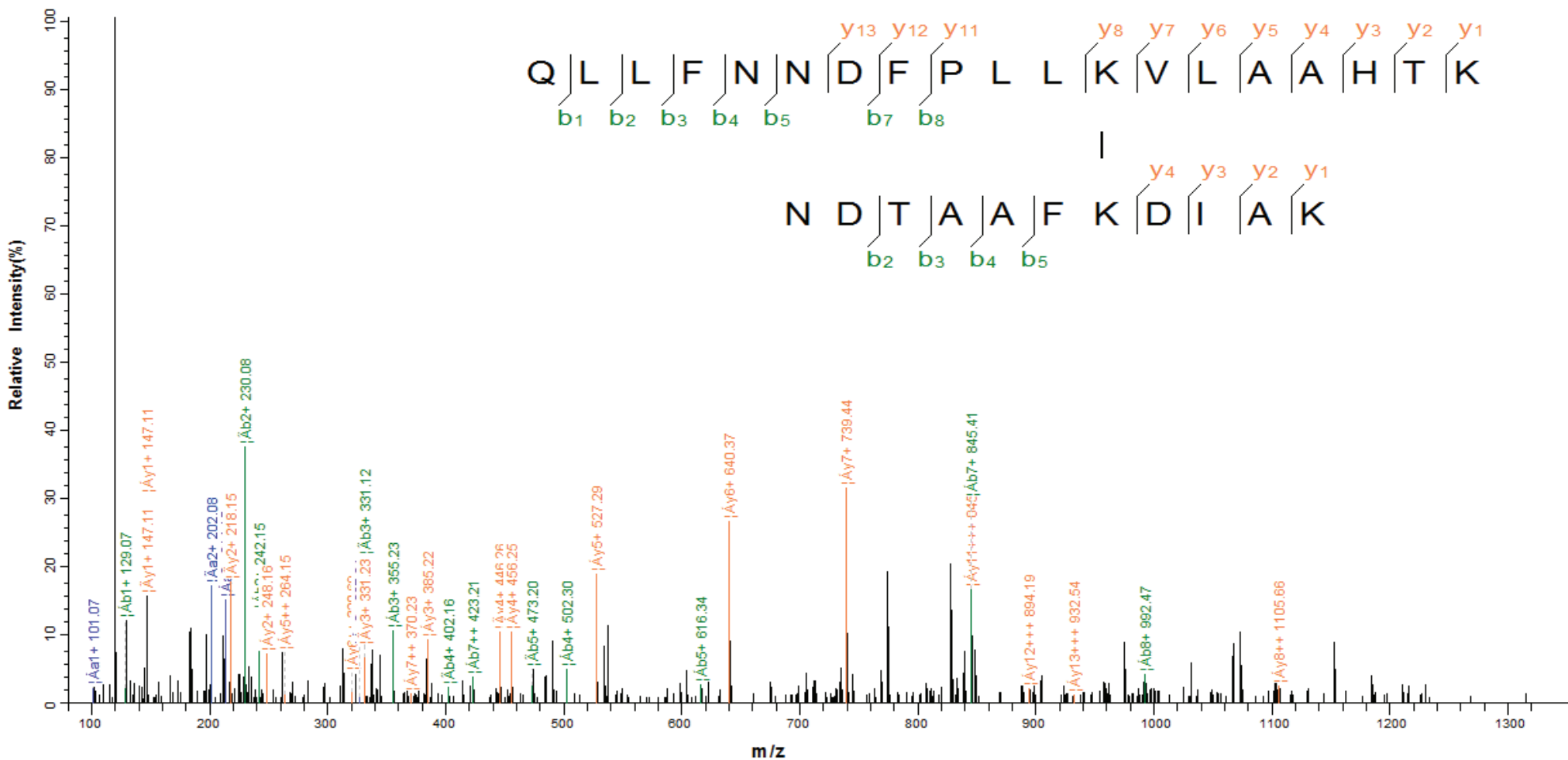


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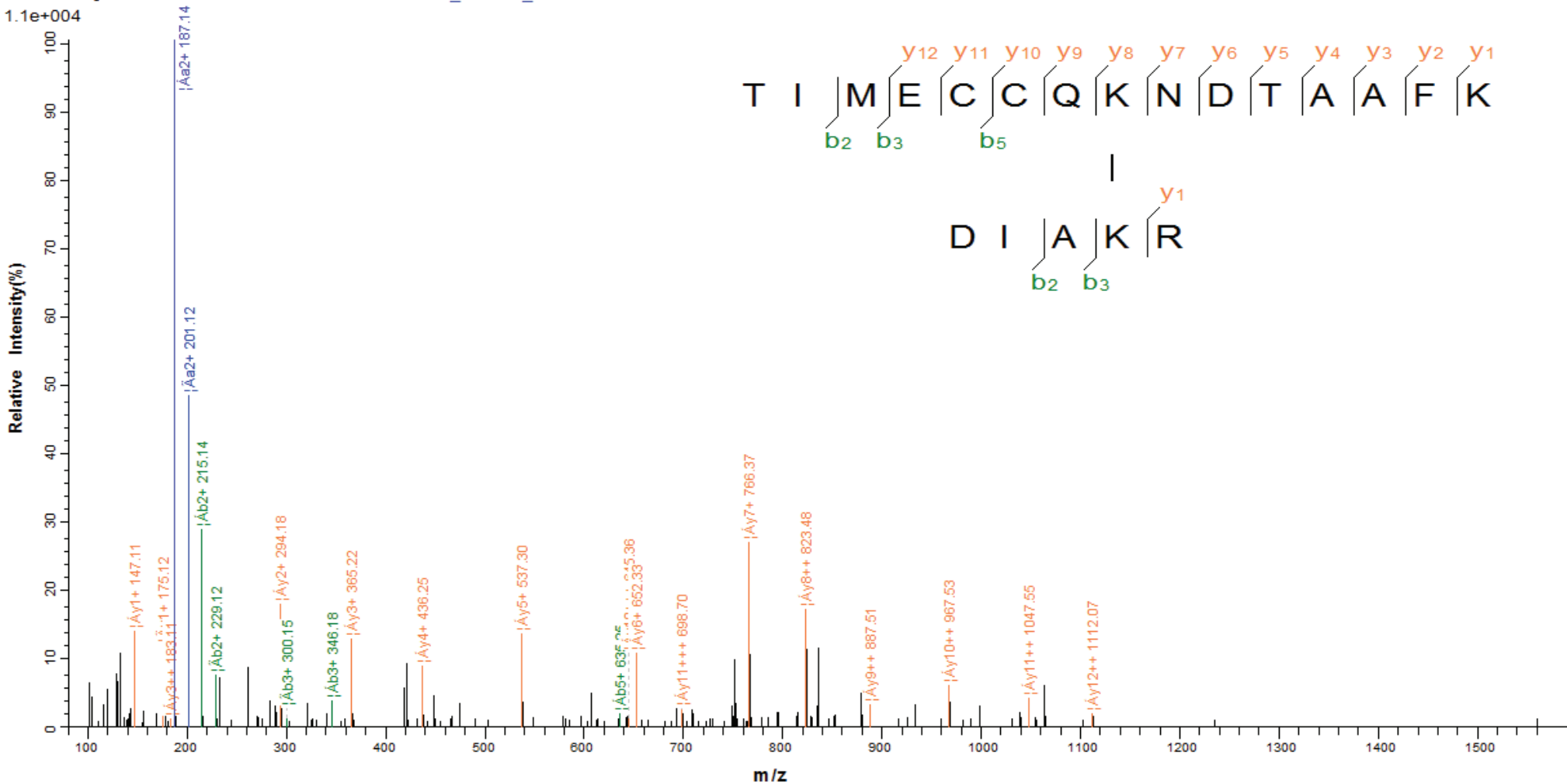


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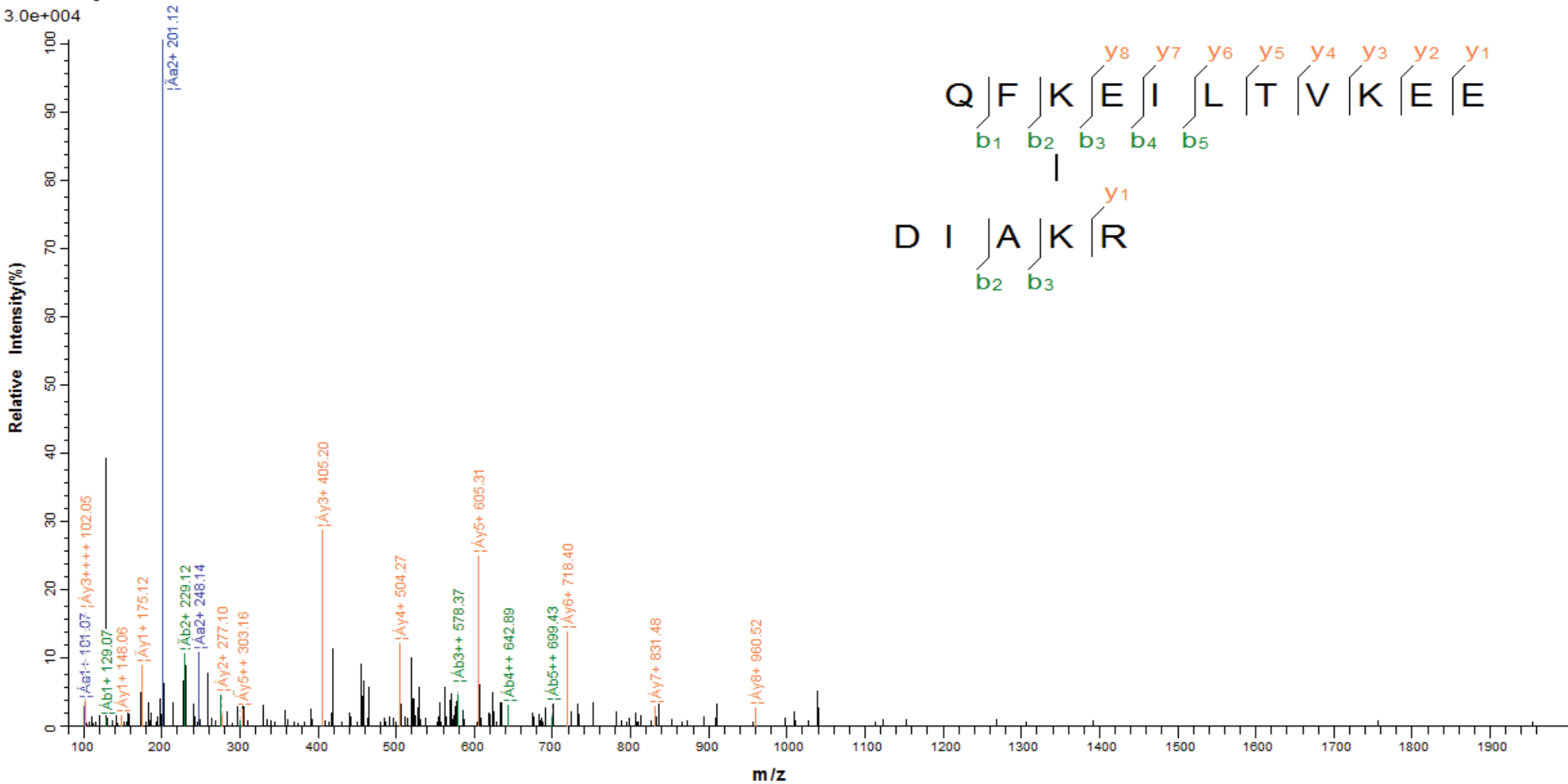
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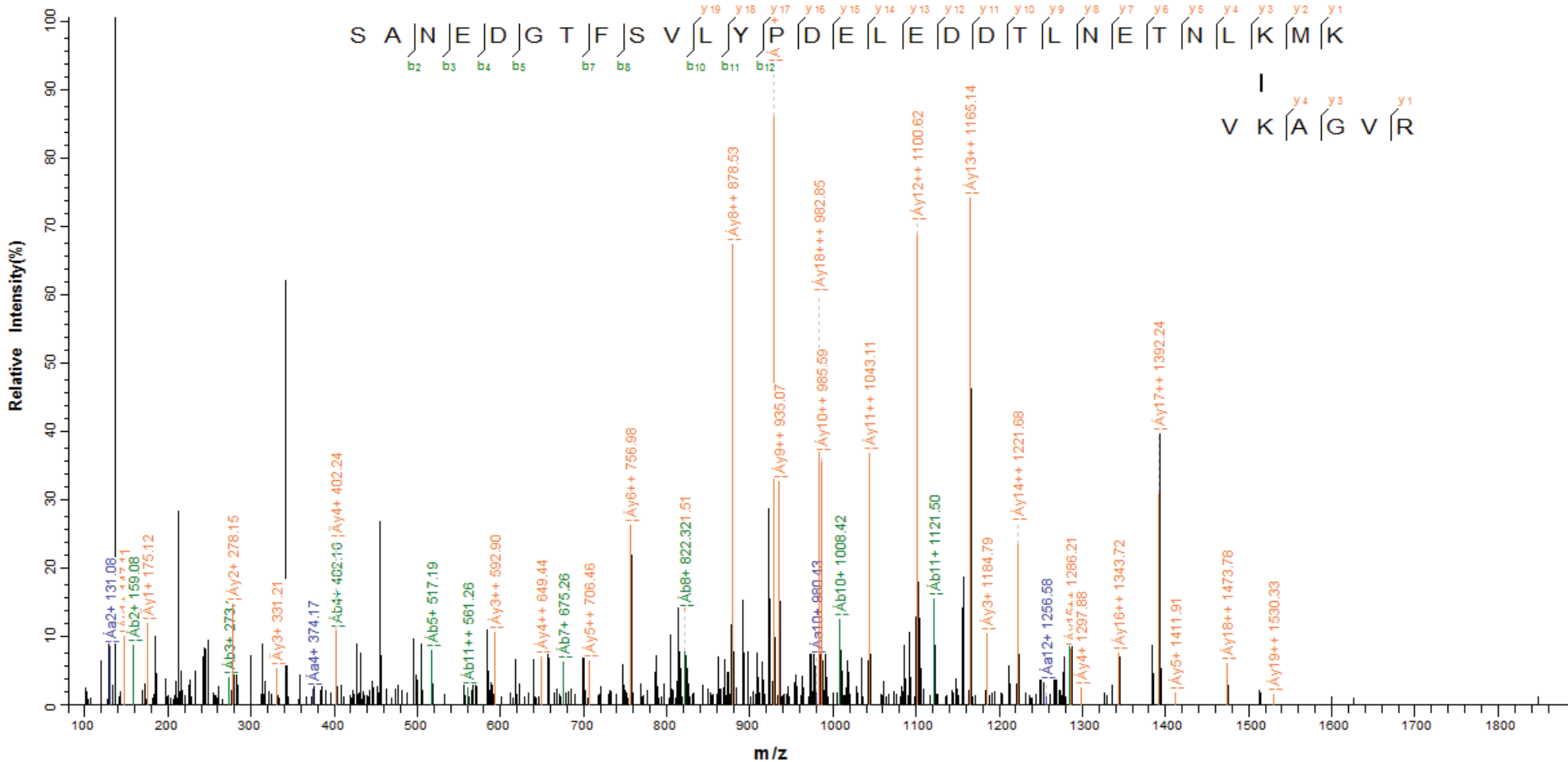


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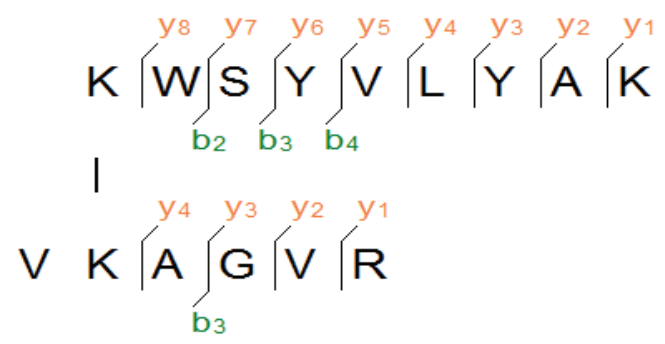
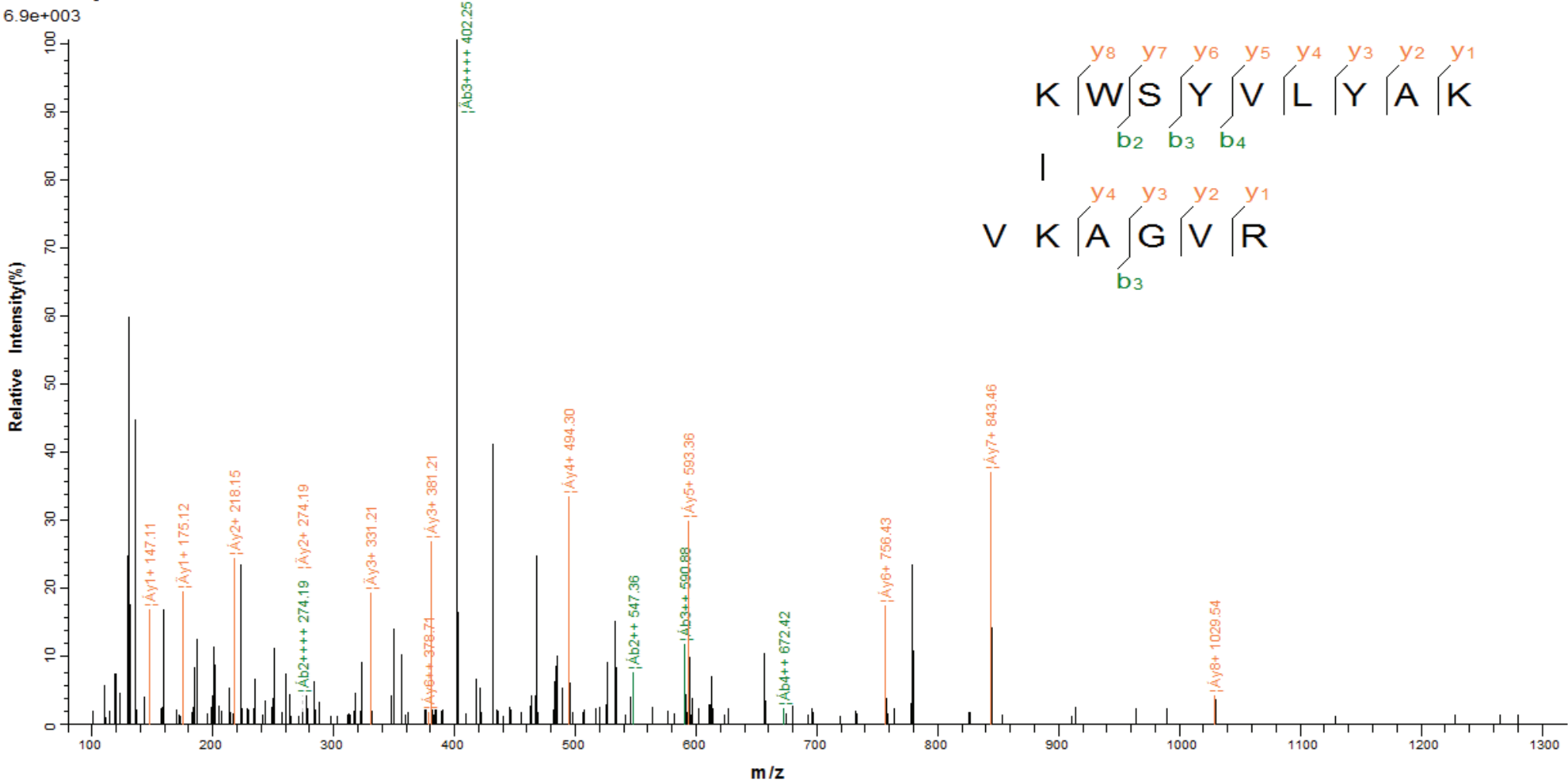


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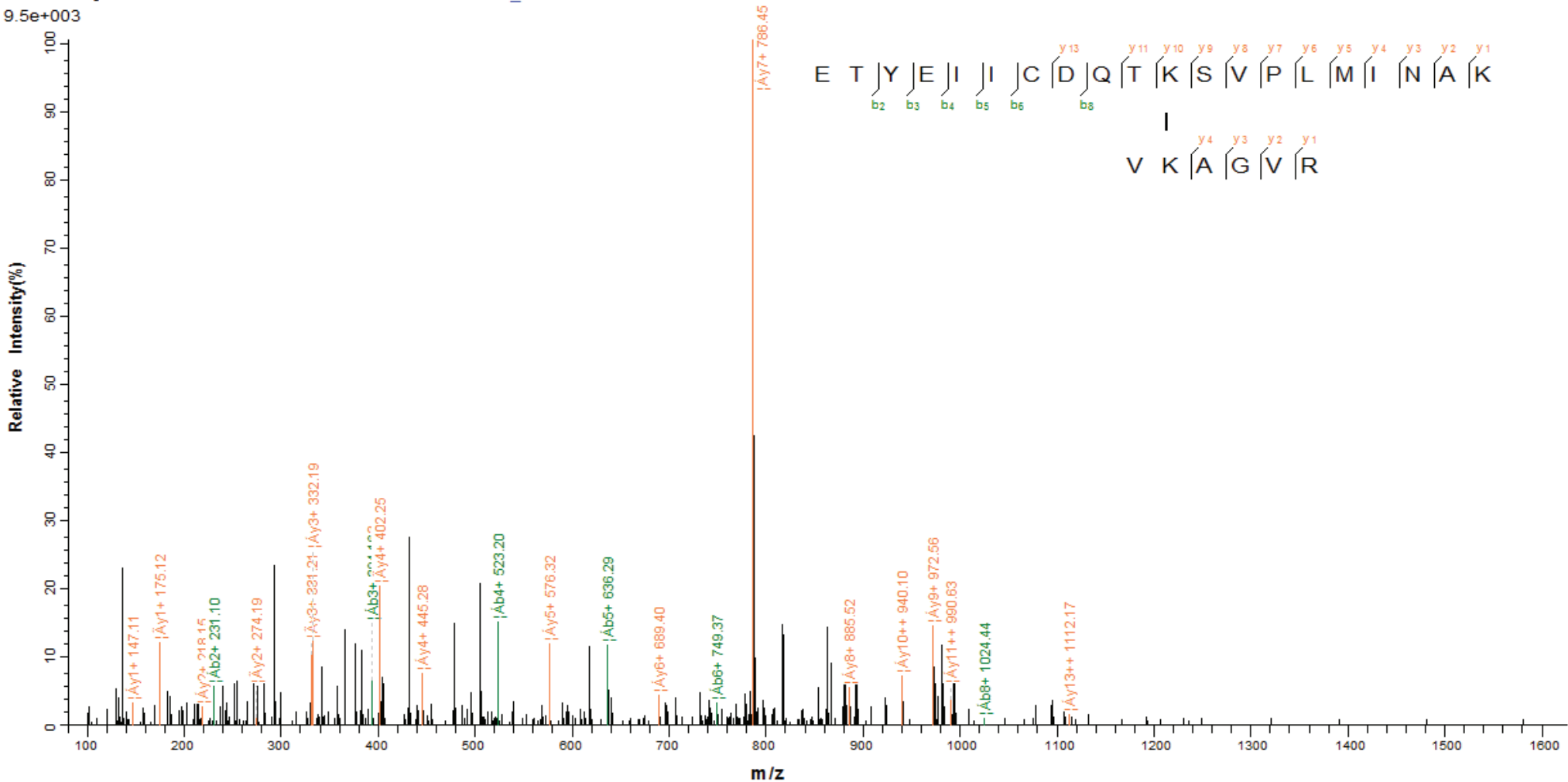
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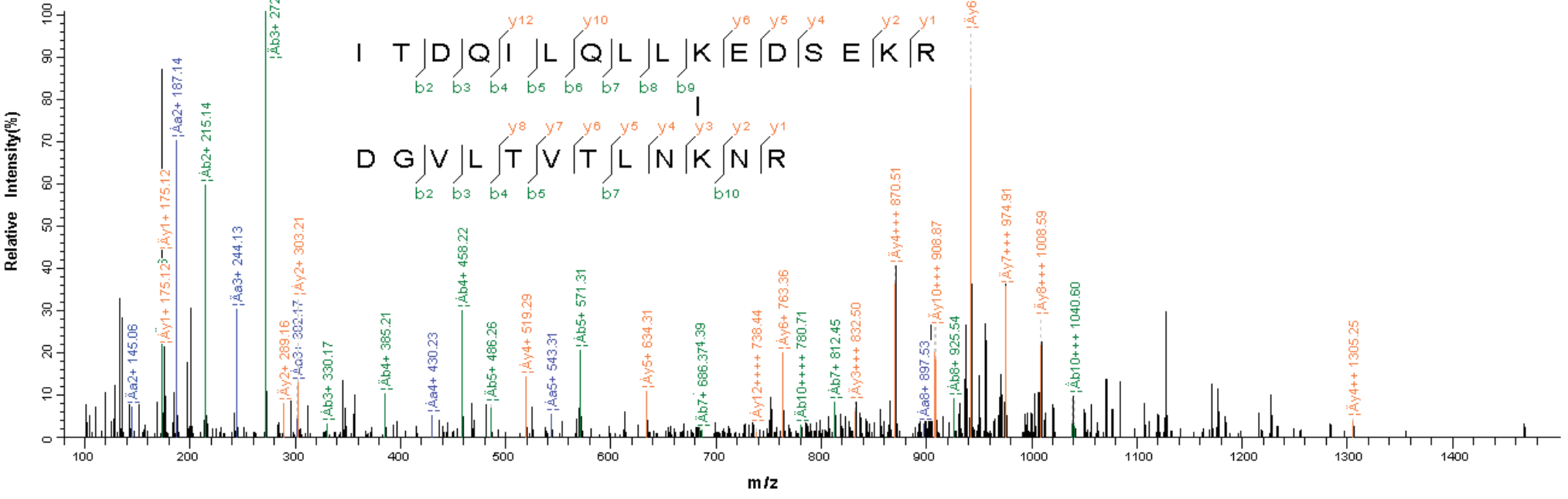
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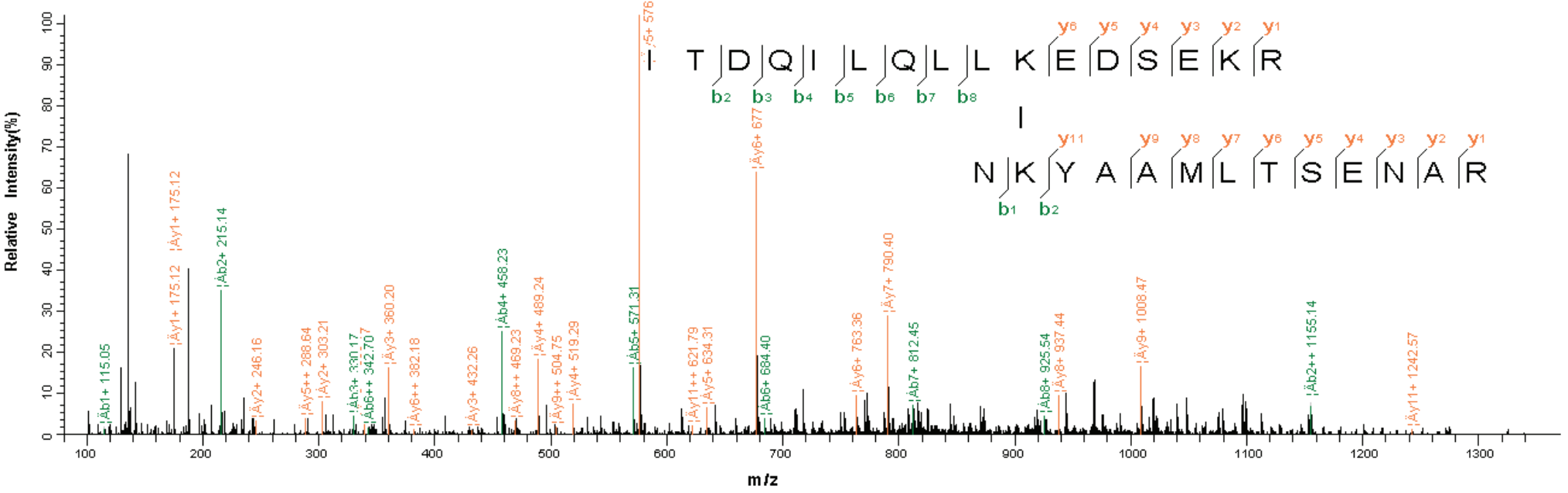
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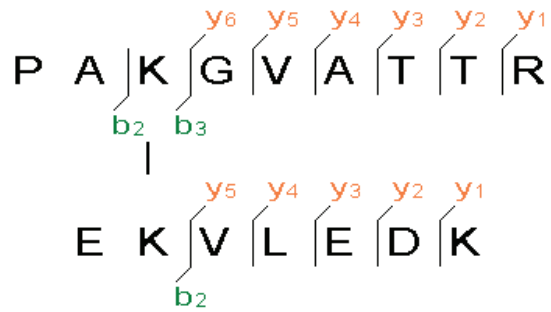
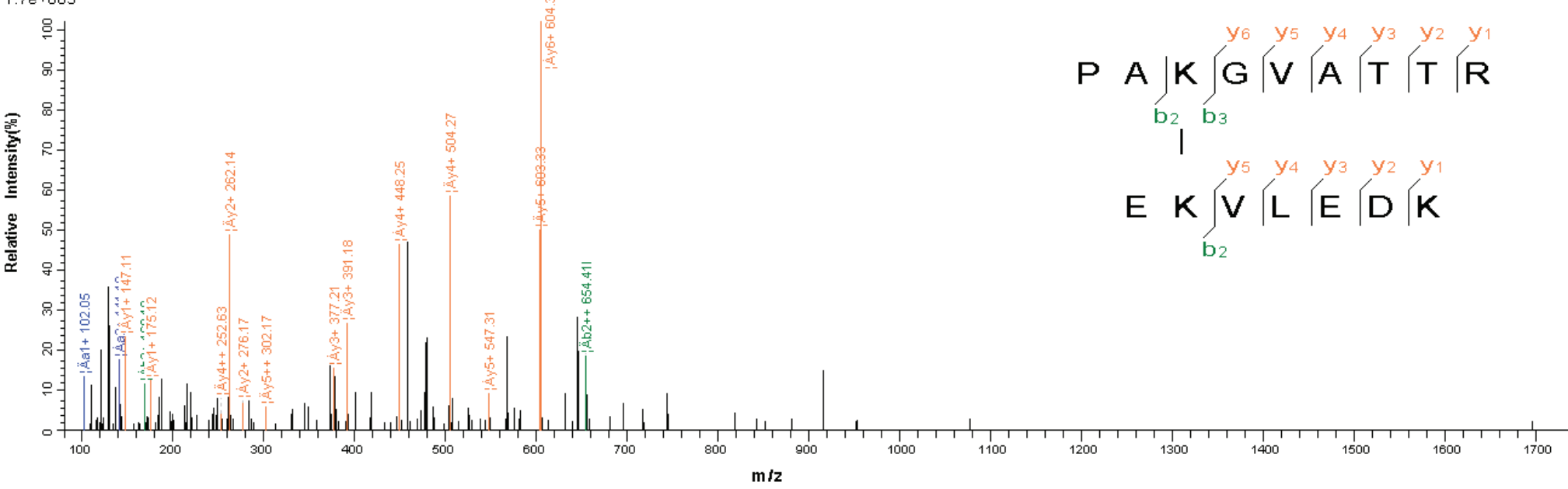
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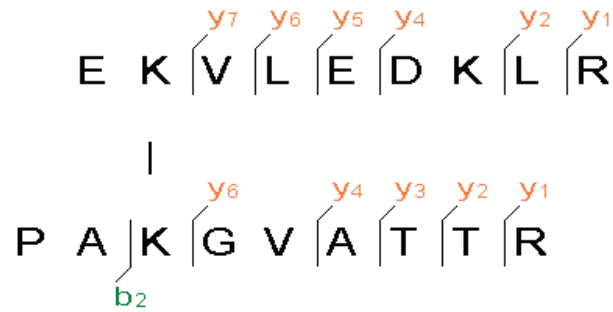
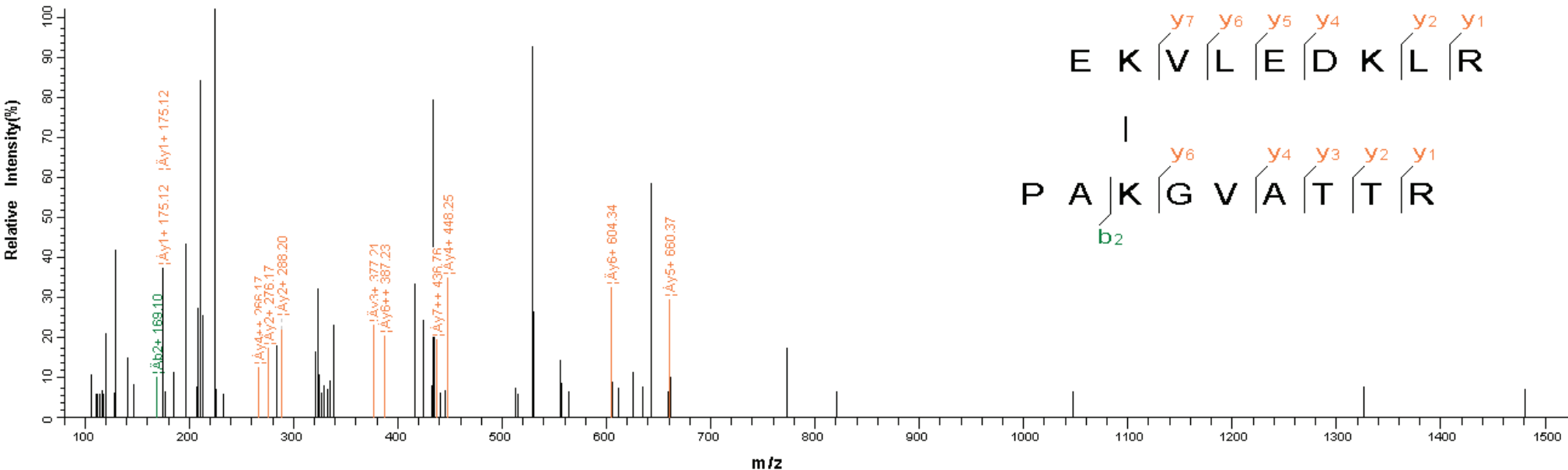
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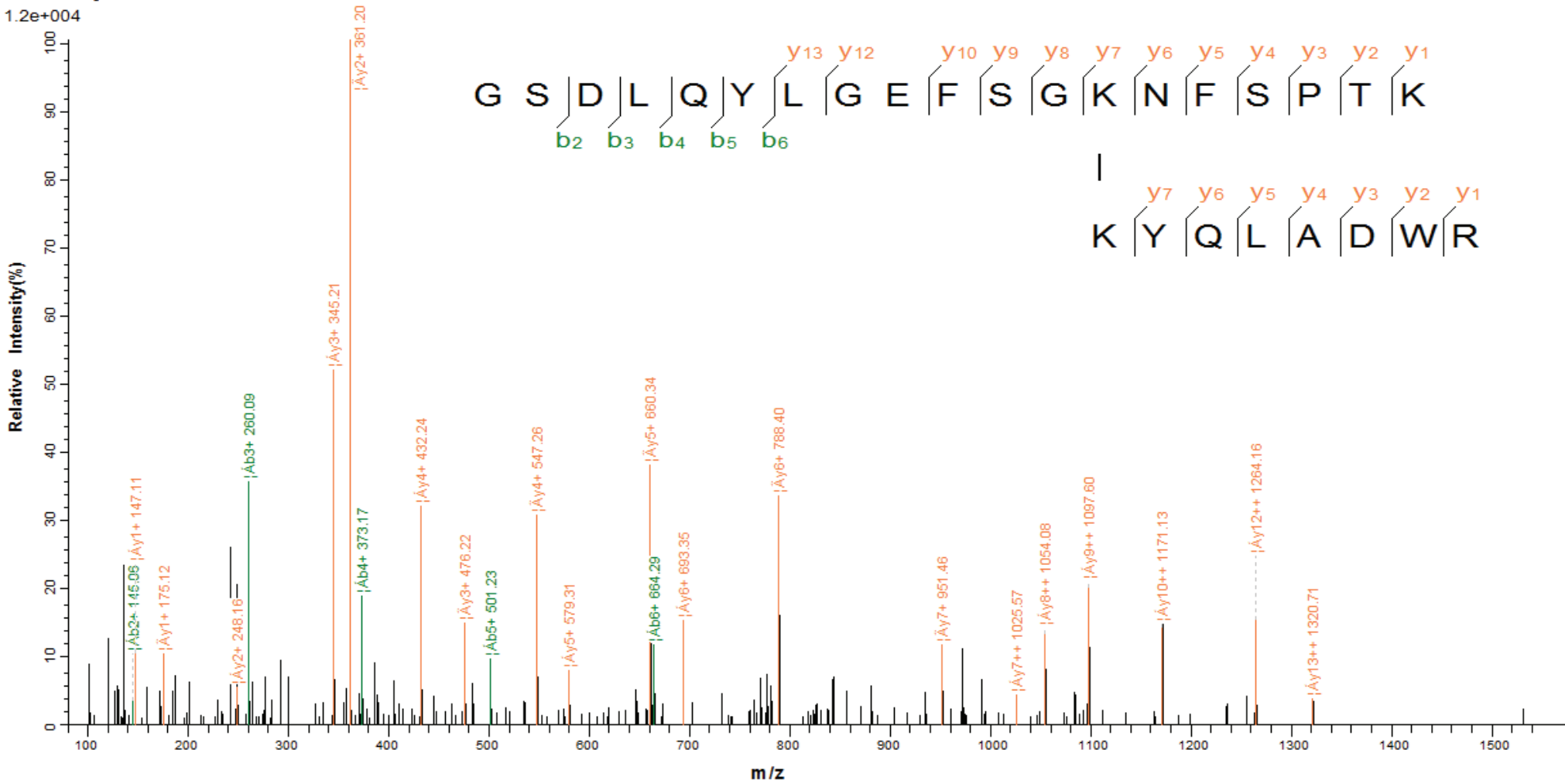
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5.2e+002 exosome_A8_140503145423.2332.2332.5.dta Seq: EKVLEDKLR-PAKGVATTR

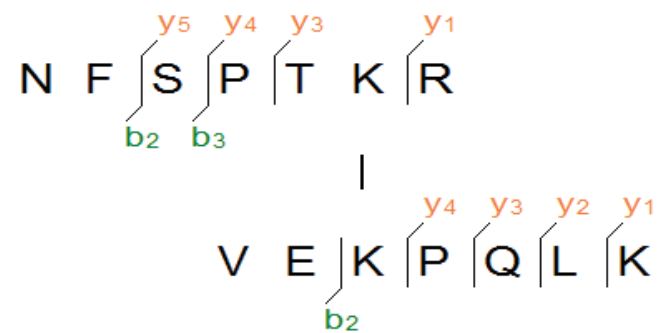
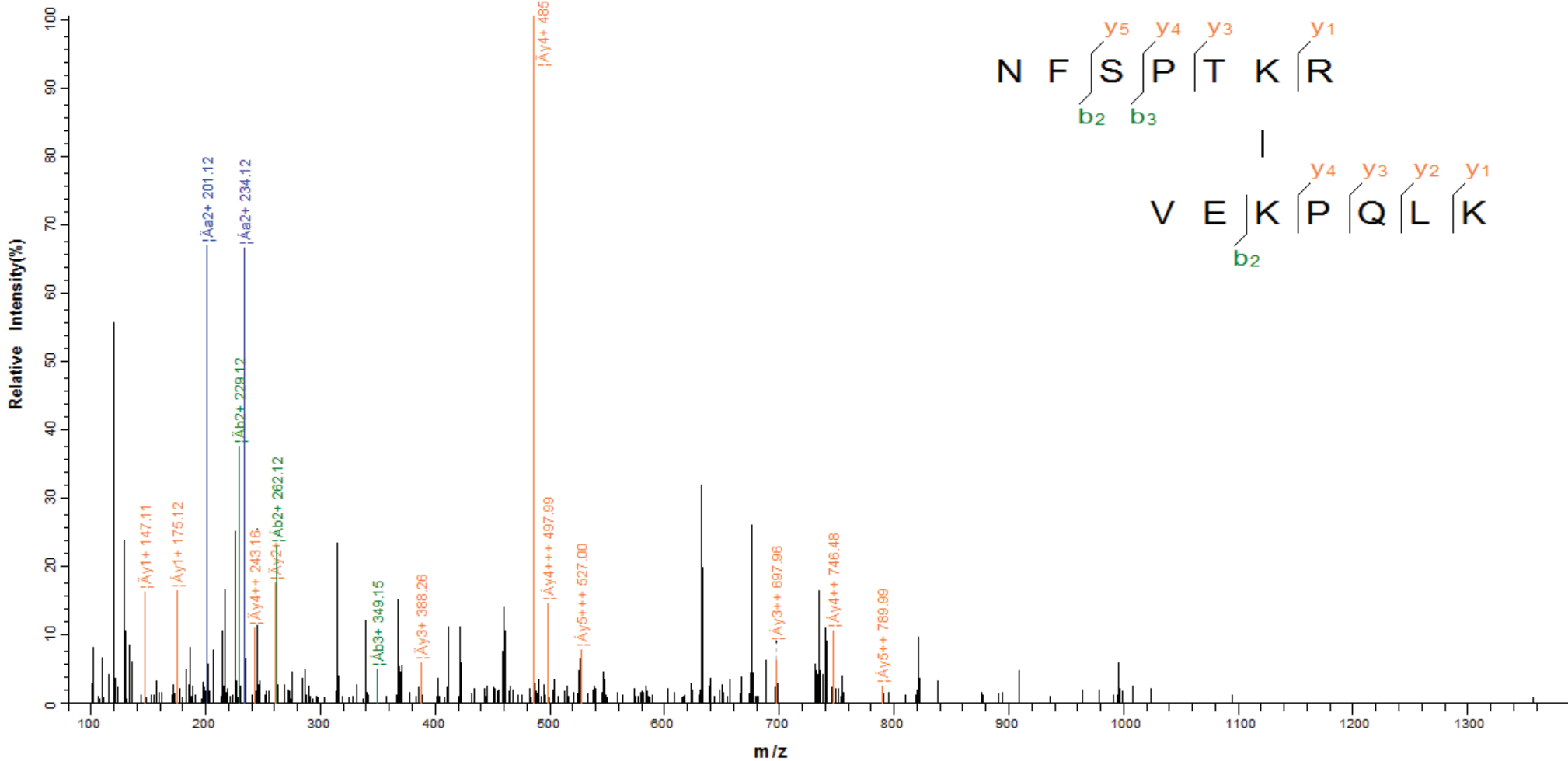


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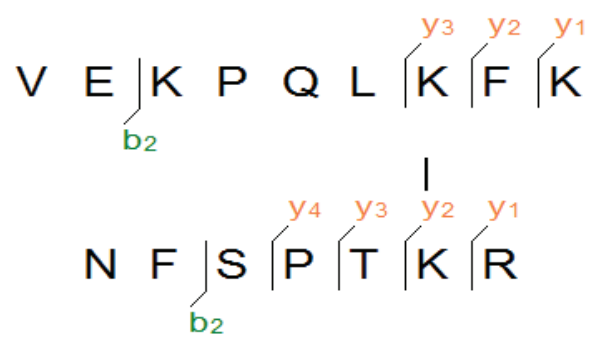
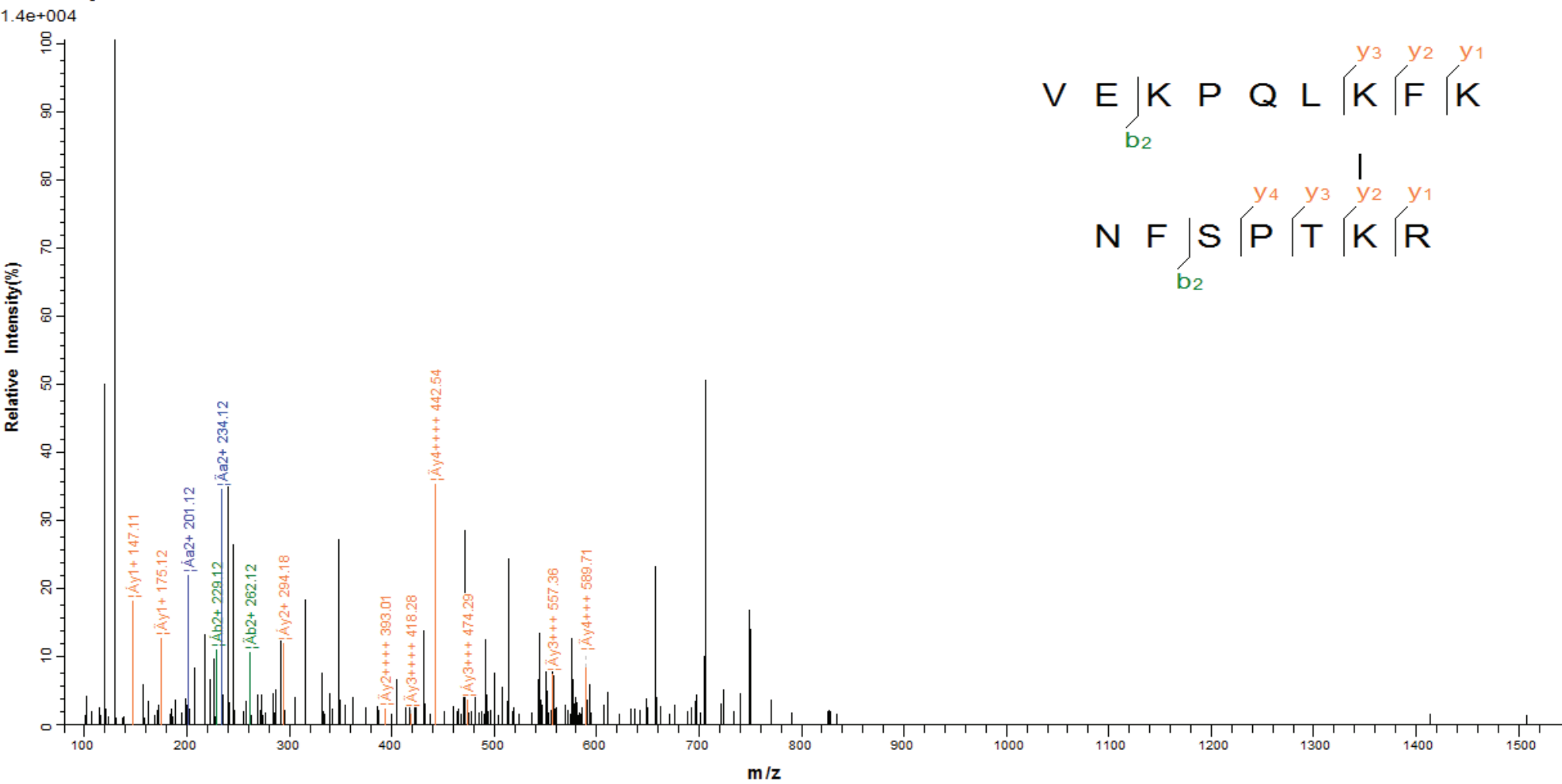


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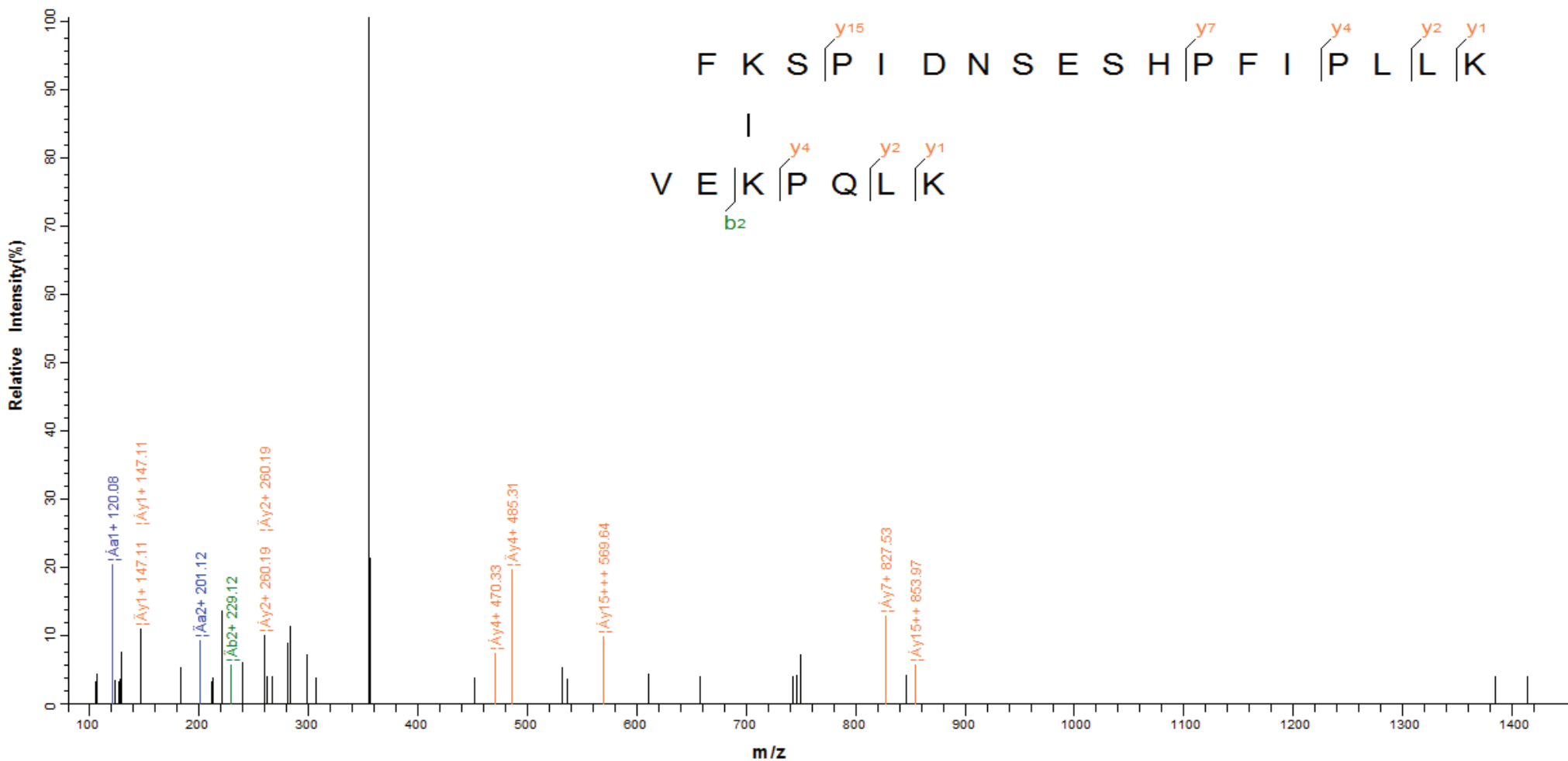


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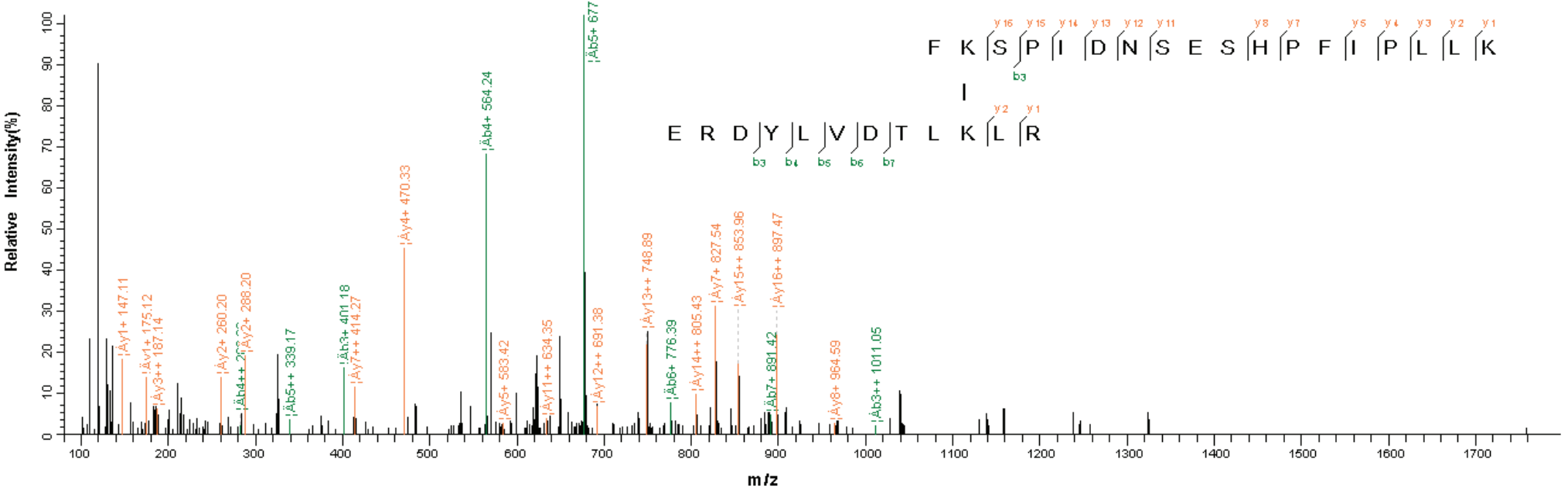


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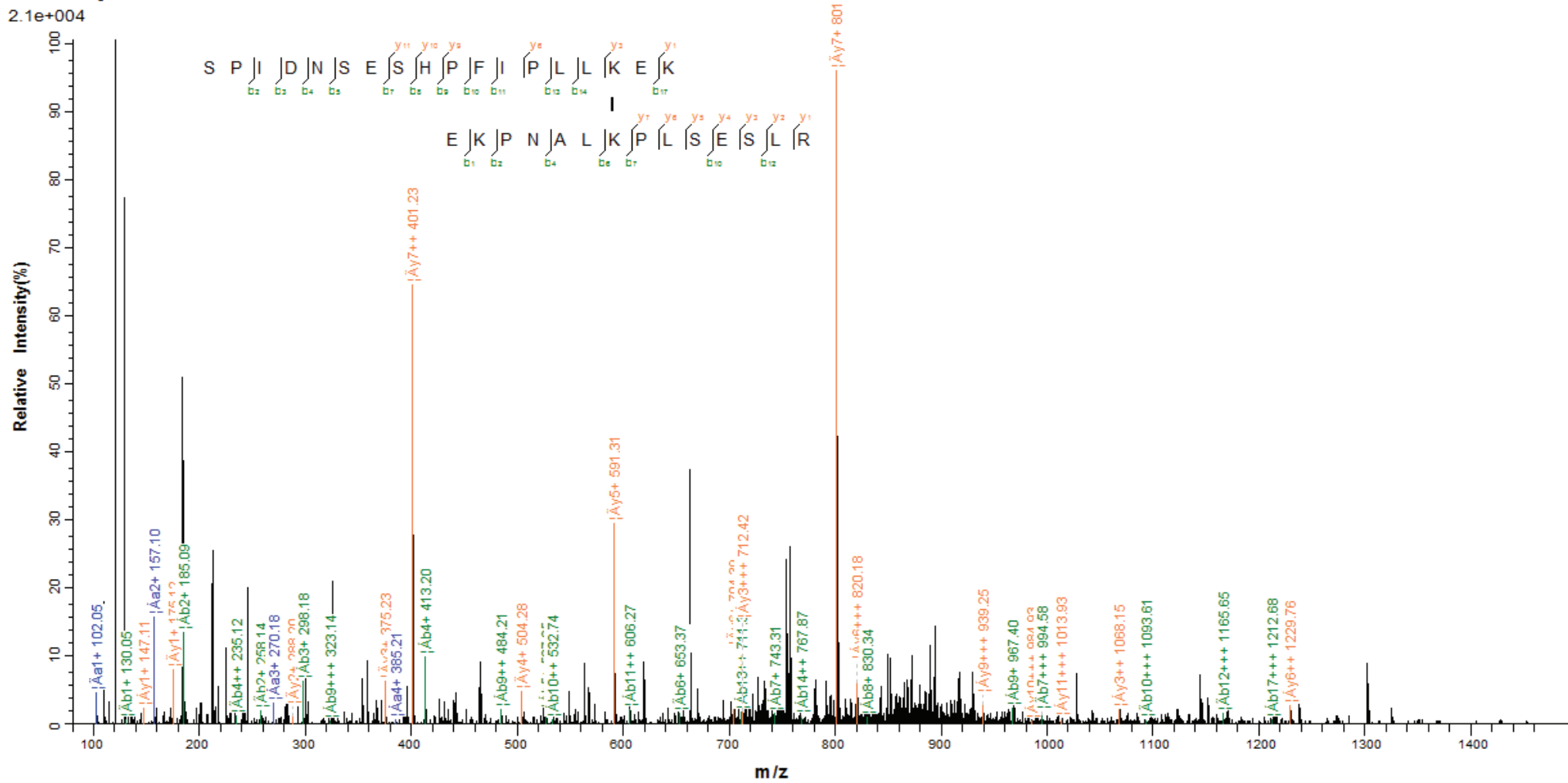


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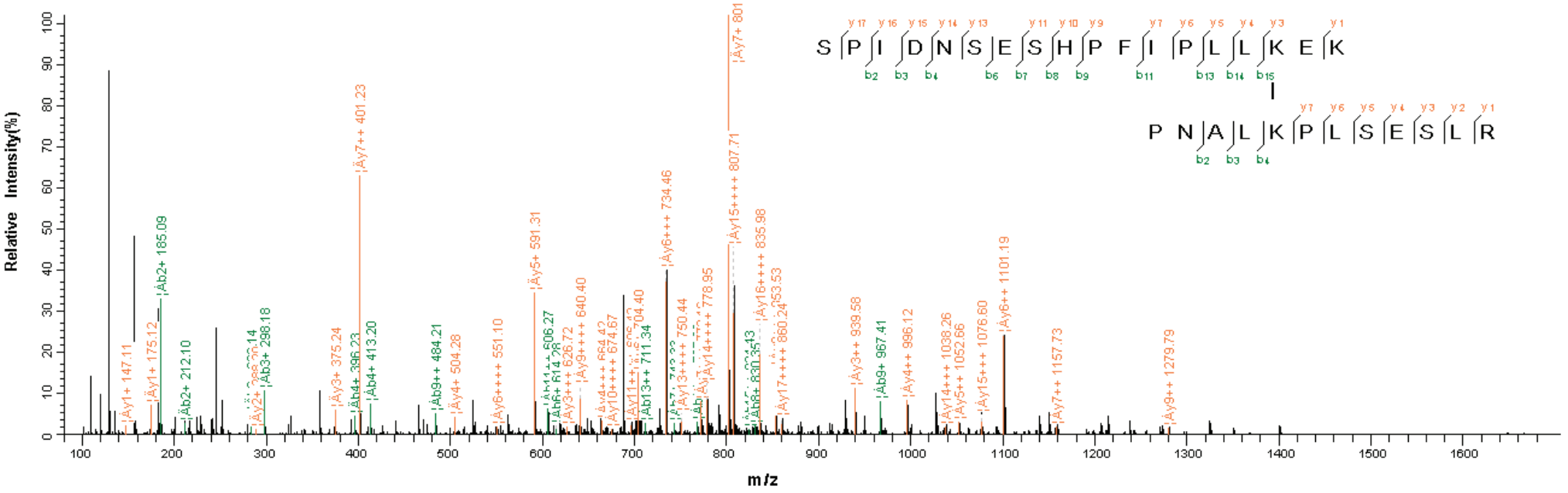


F K S P I D N S E S H P F I P L L K
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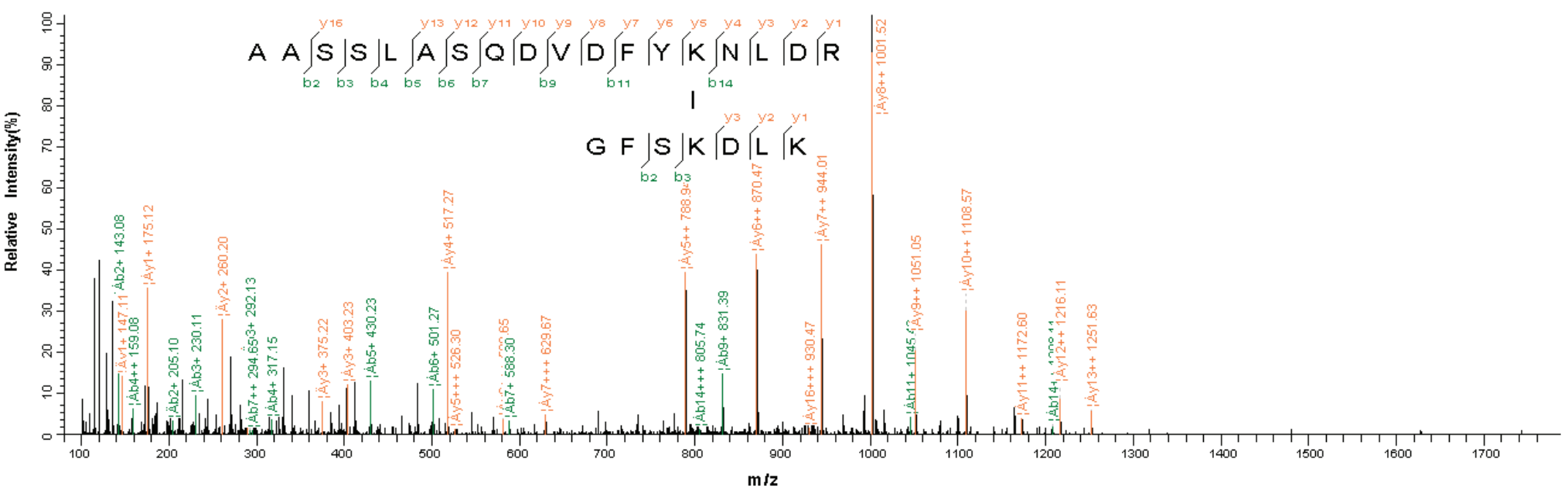
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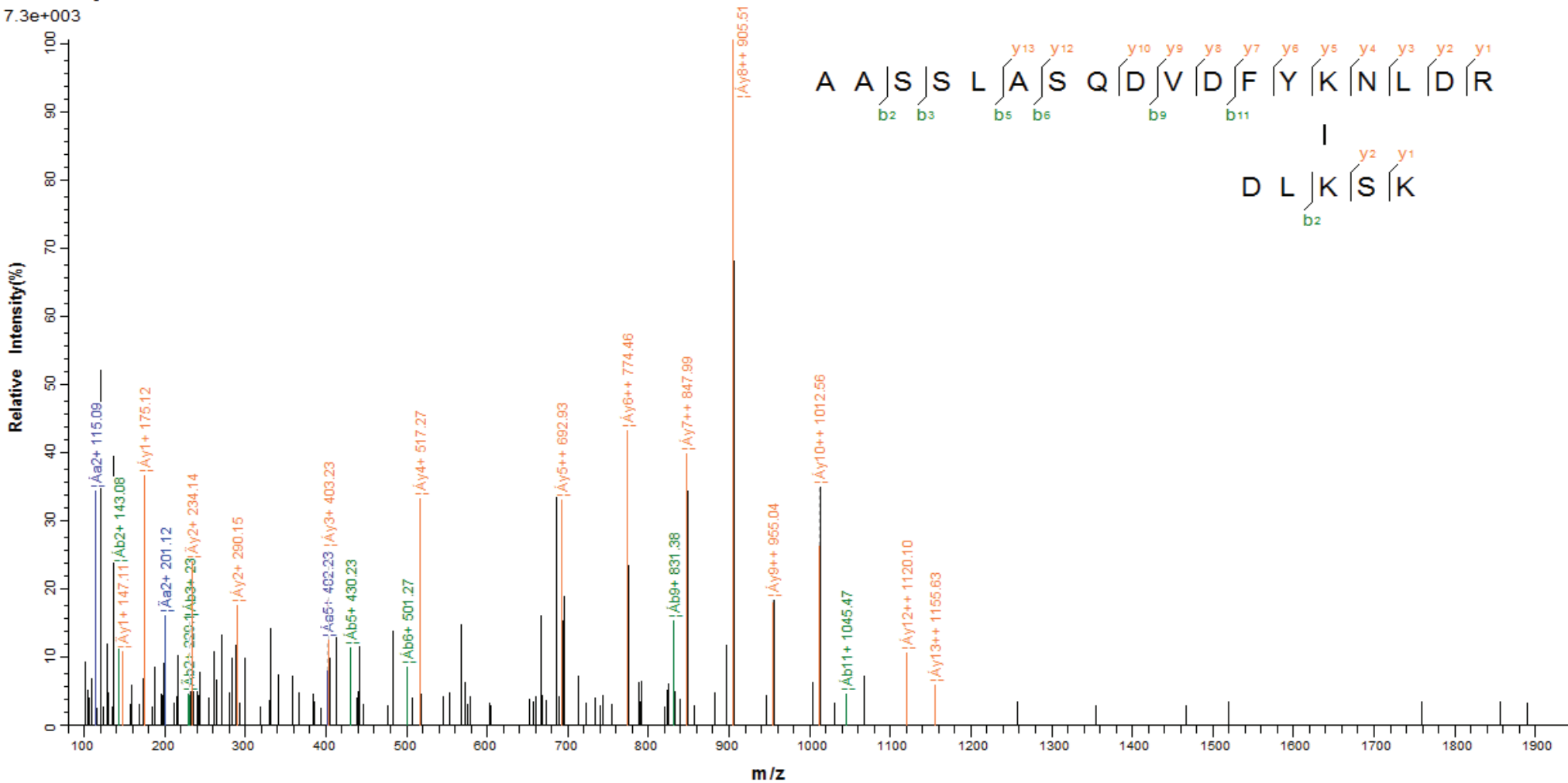
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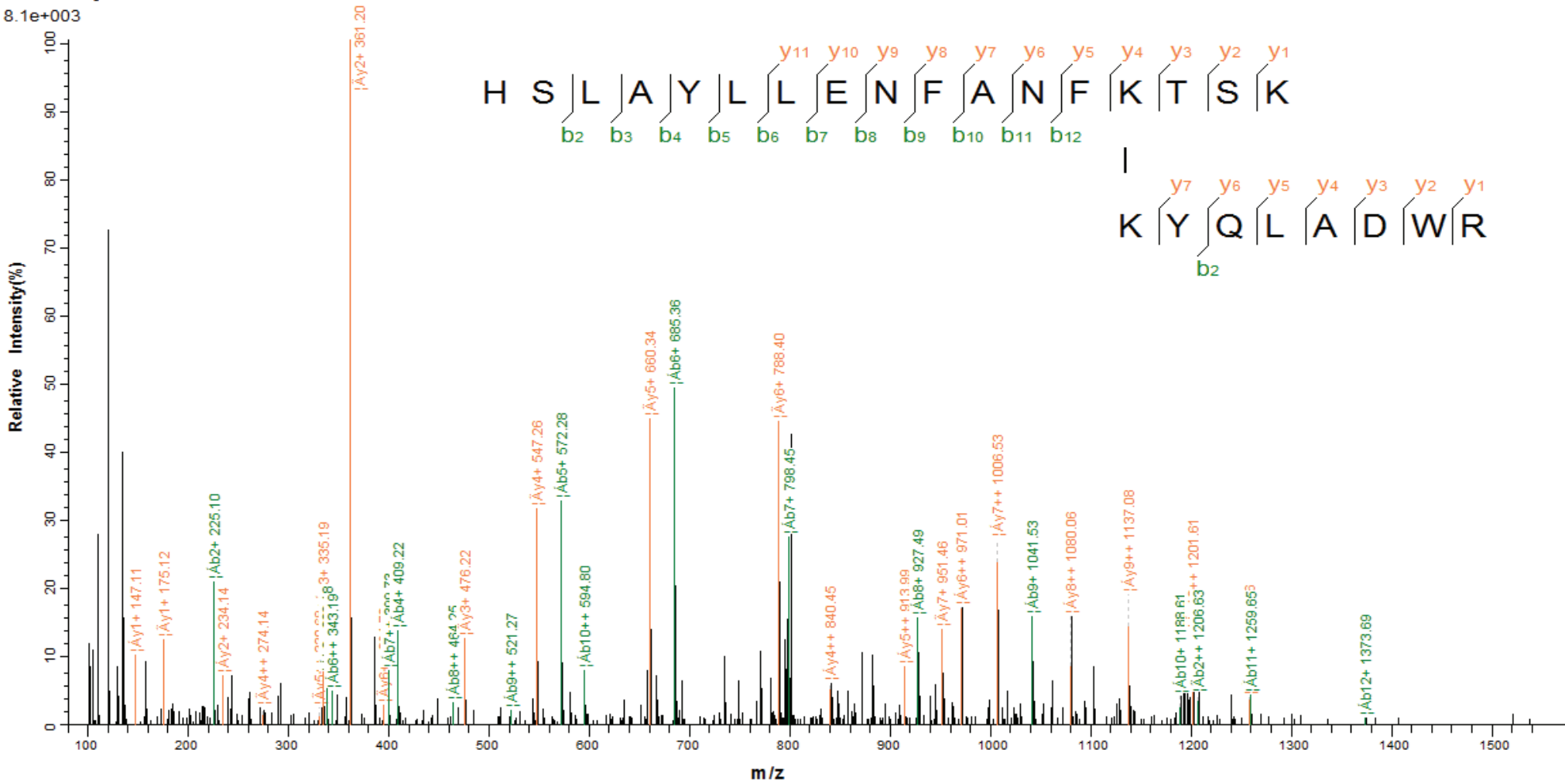
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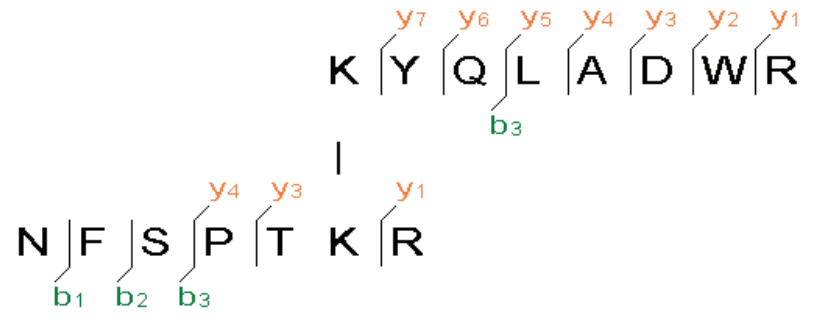
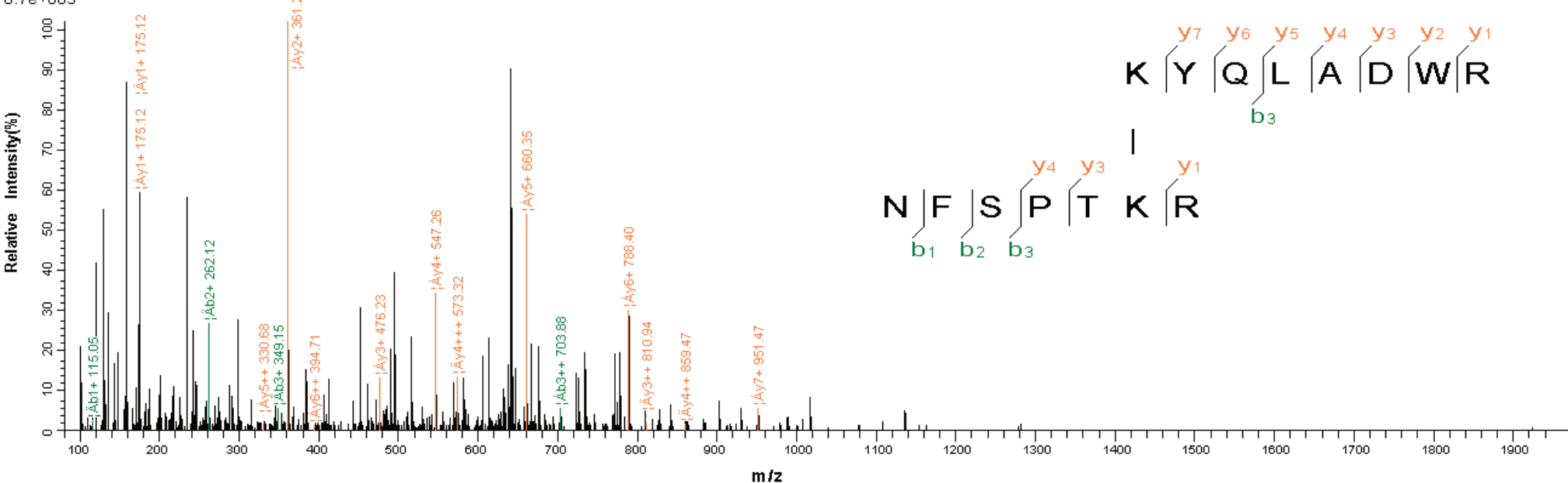
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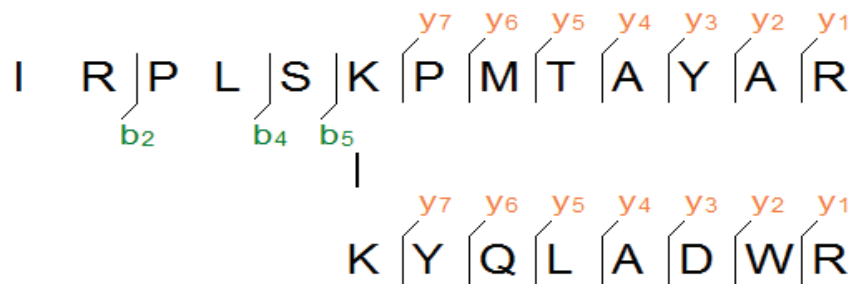
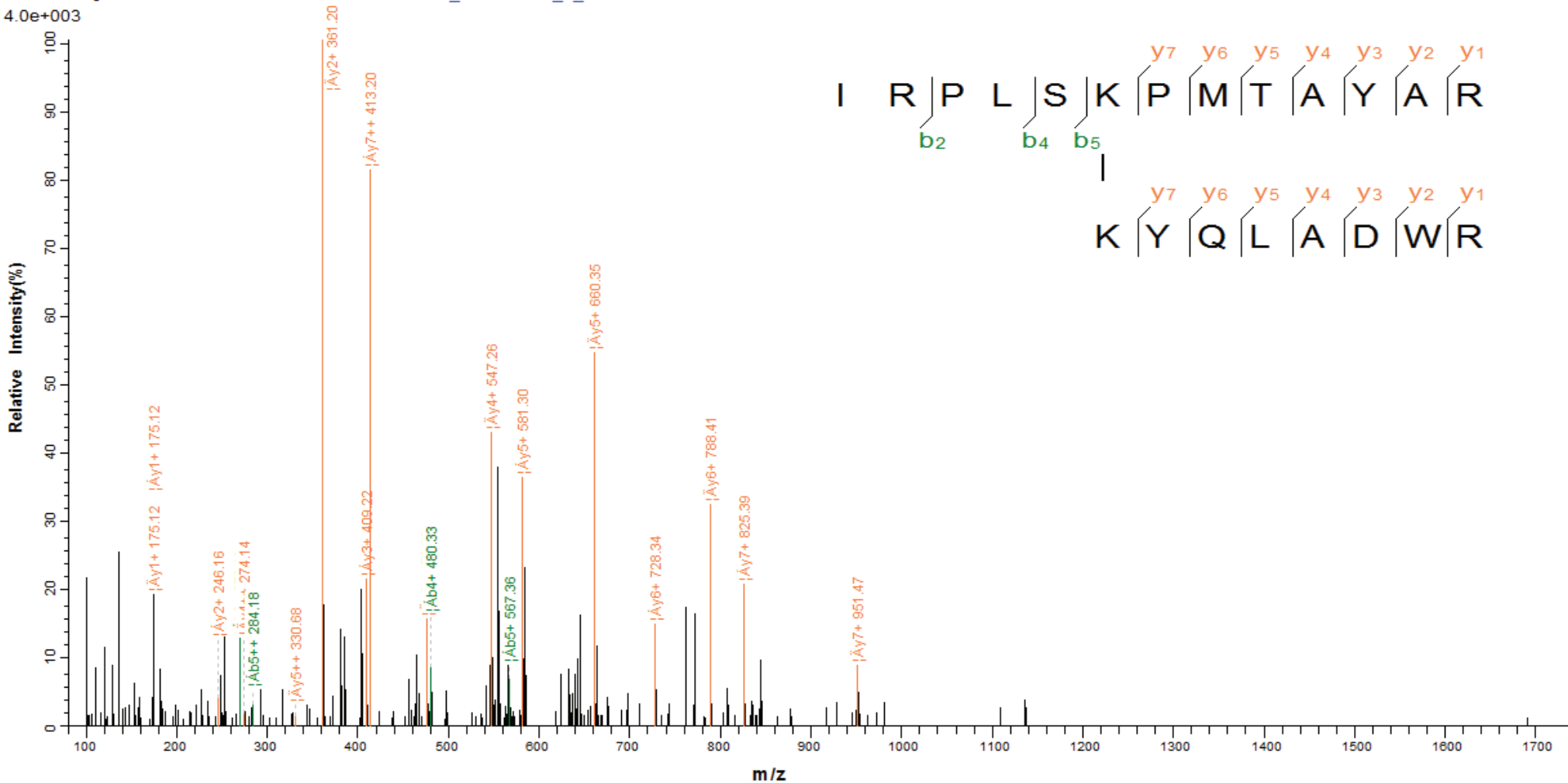
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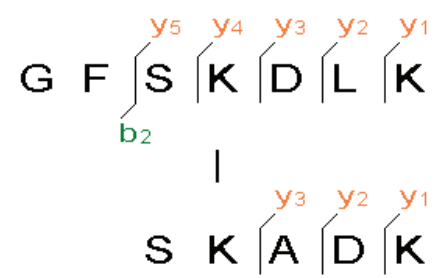
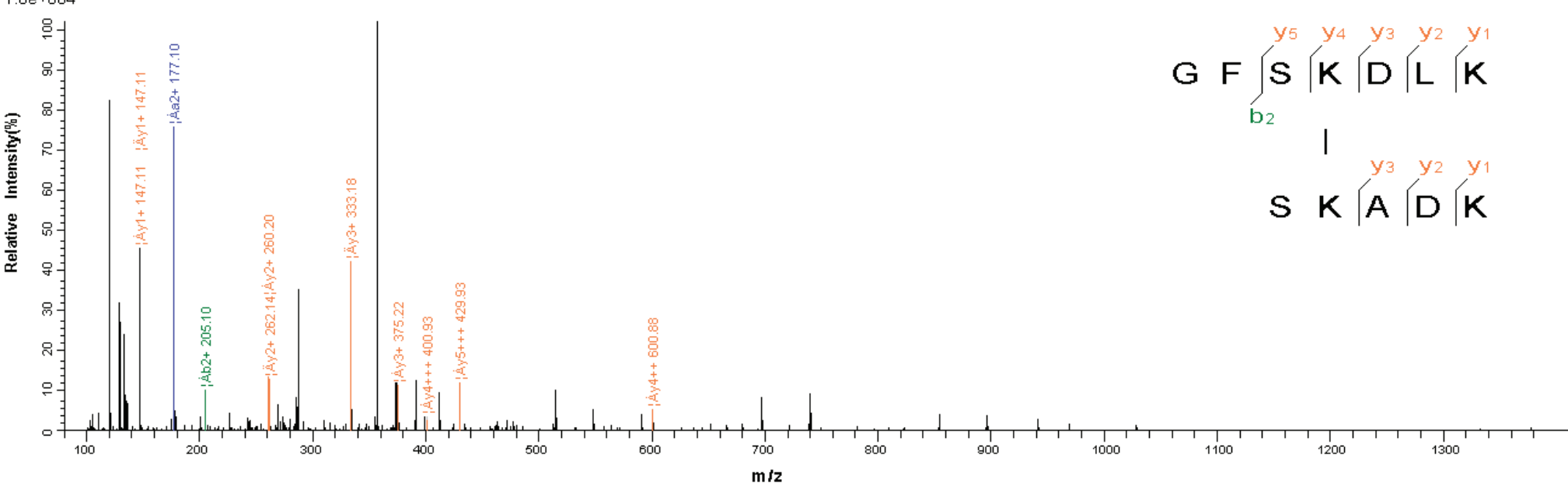
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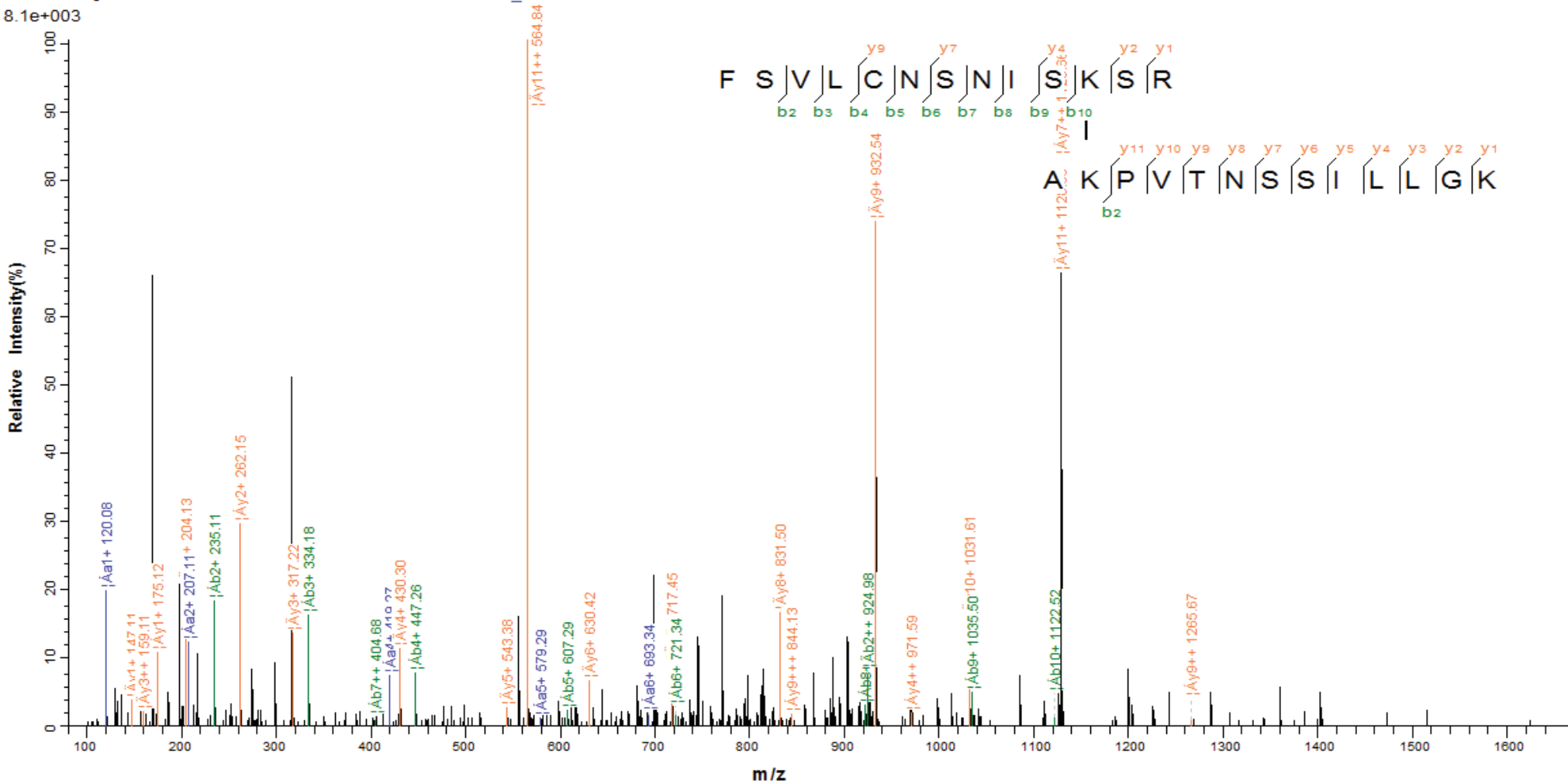
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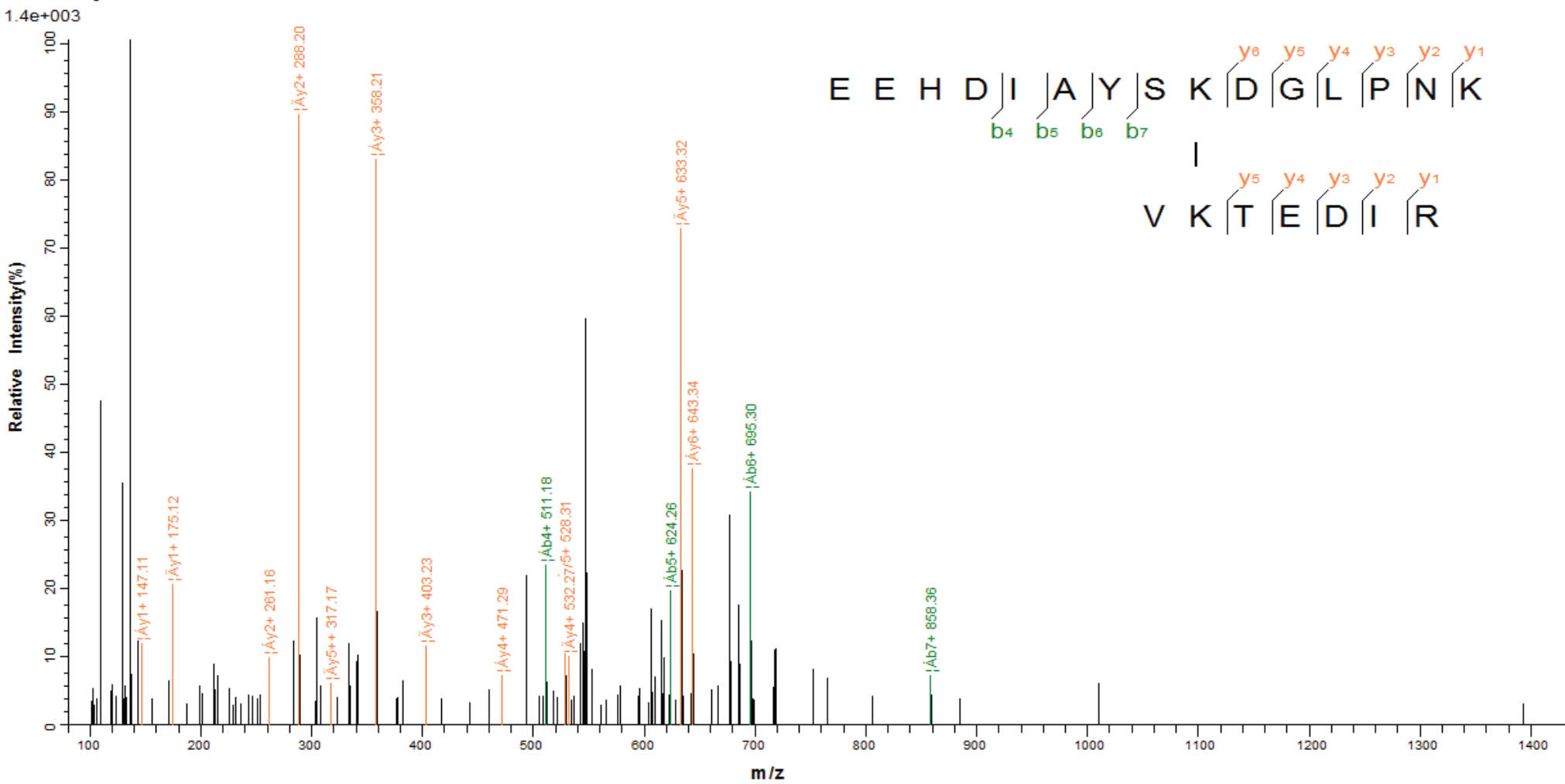
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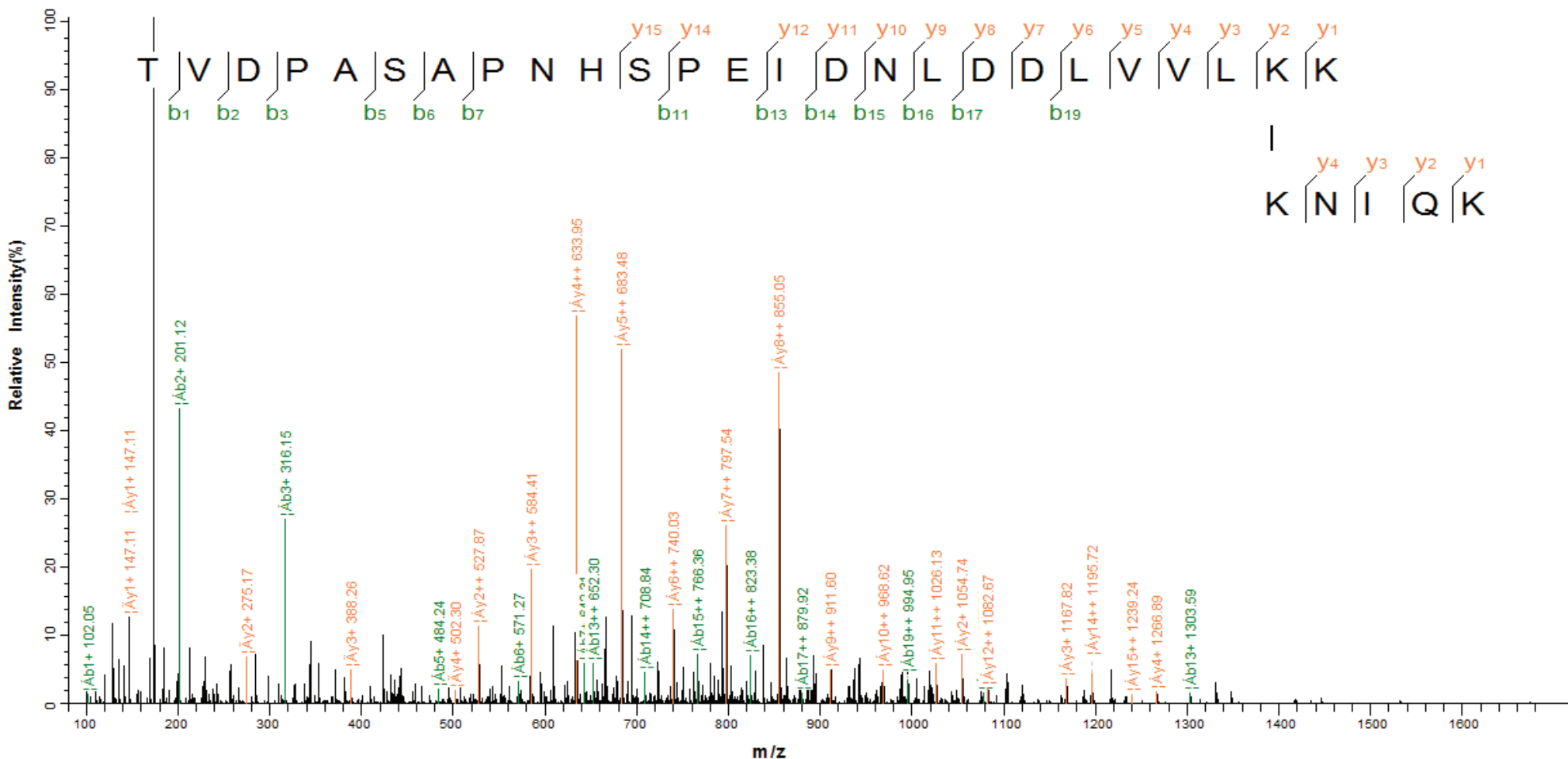


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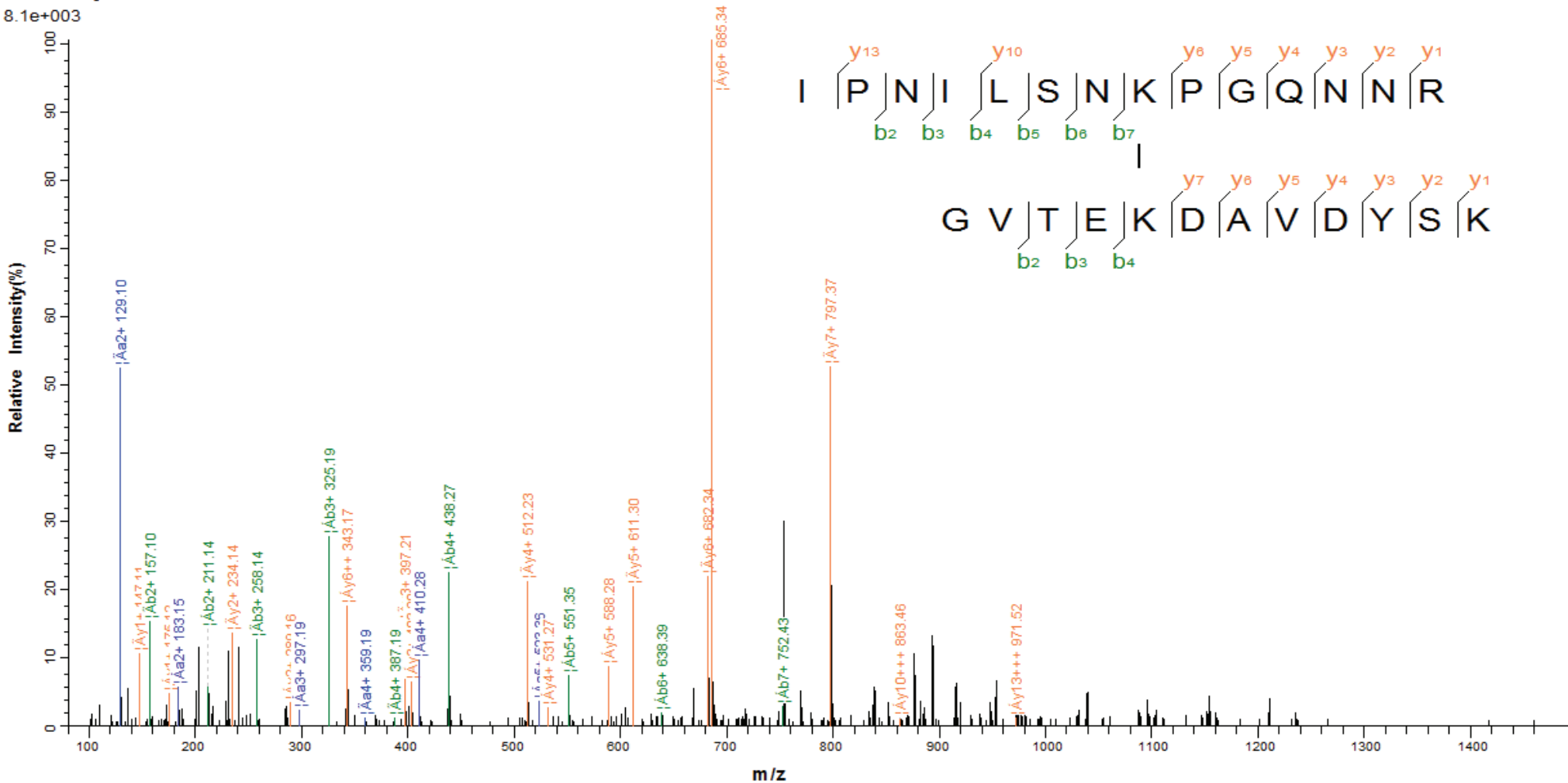


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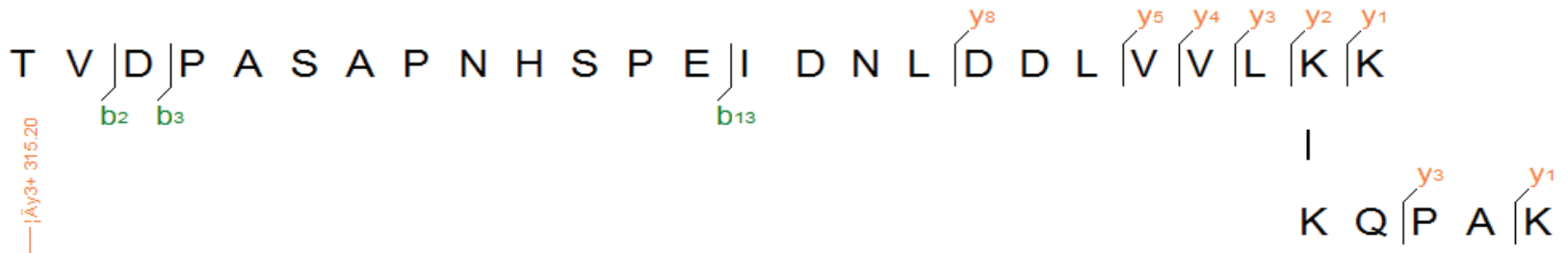
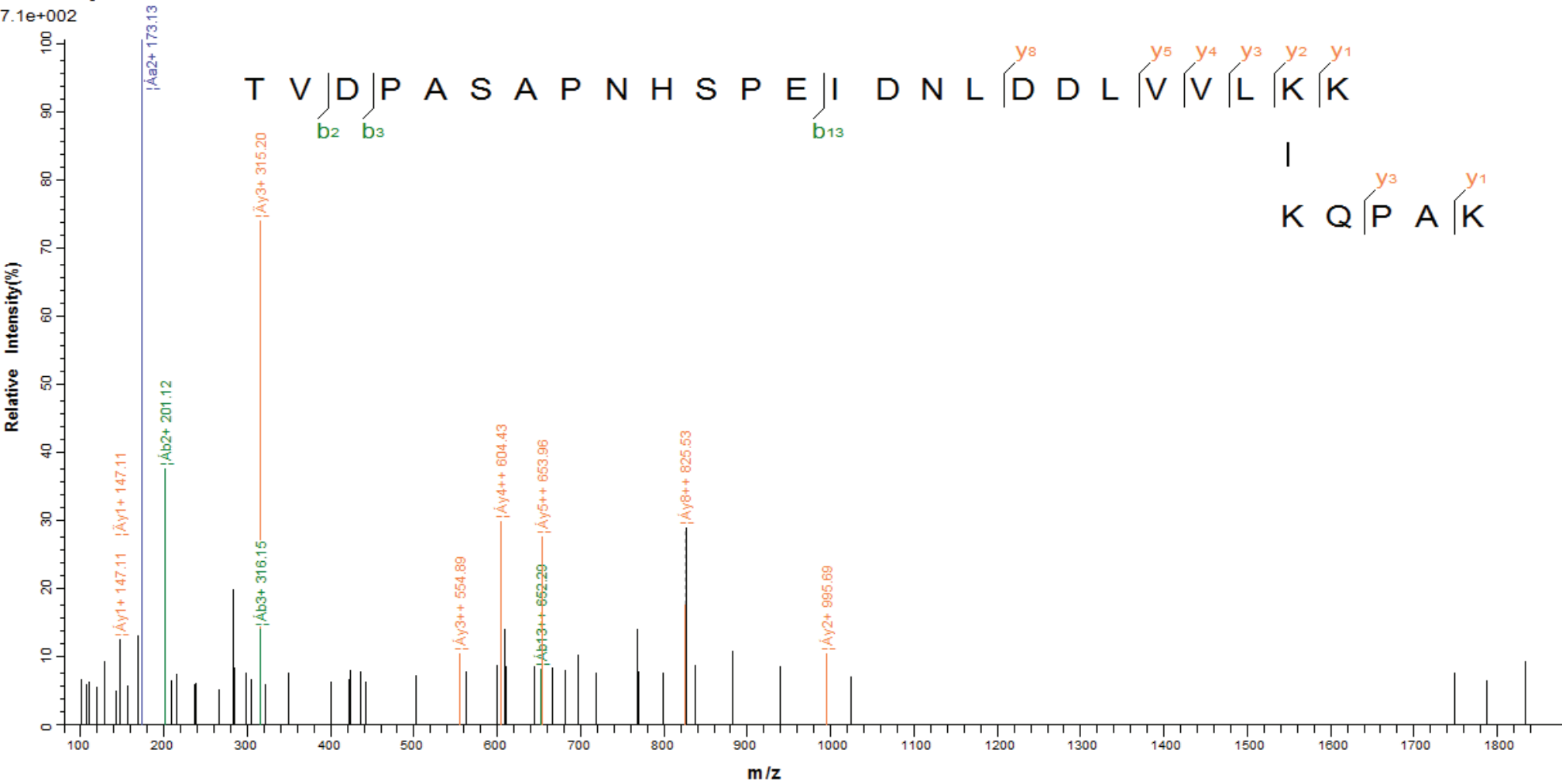
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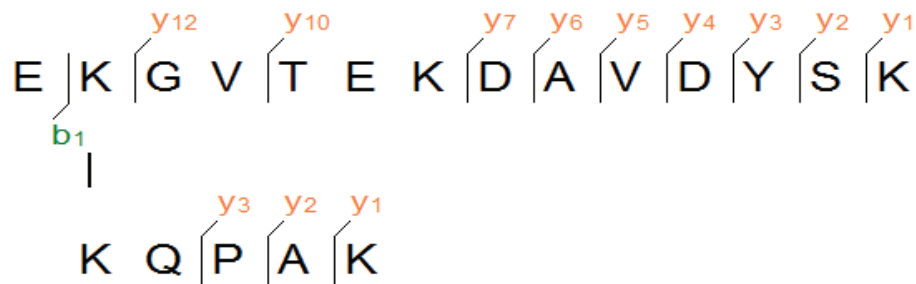
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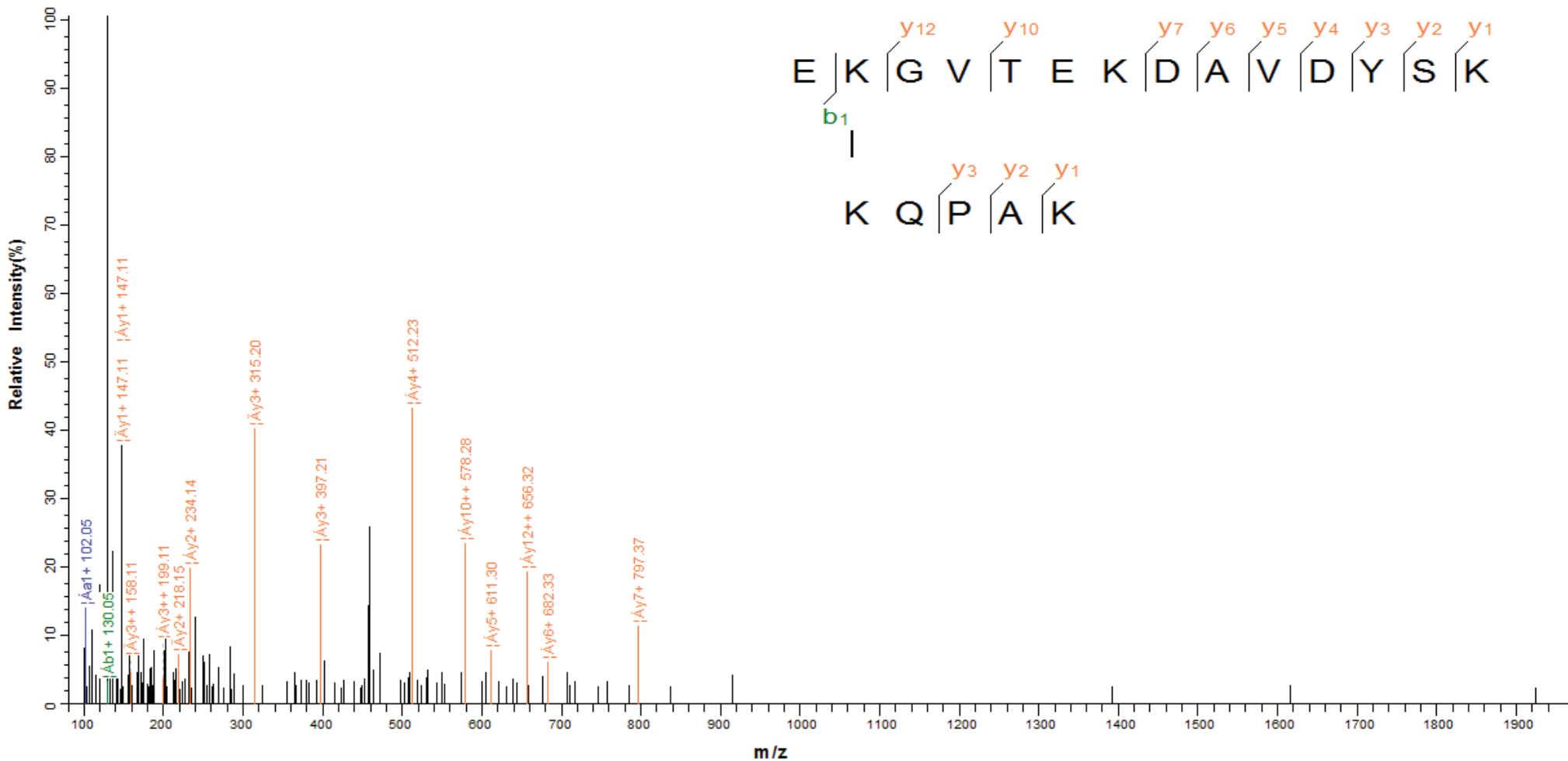
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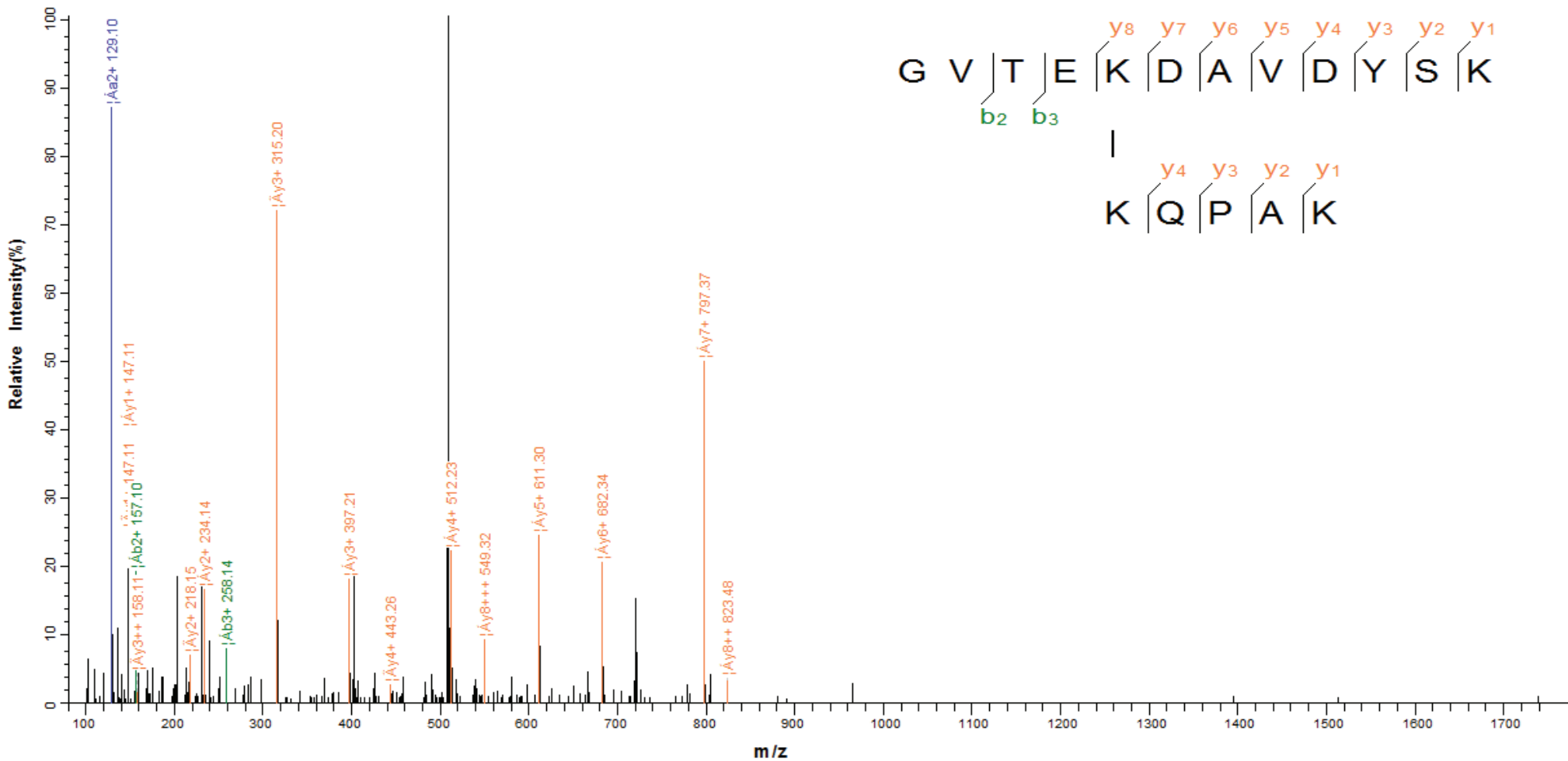


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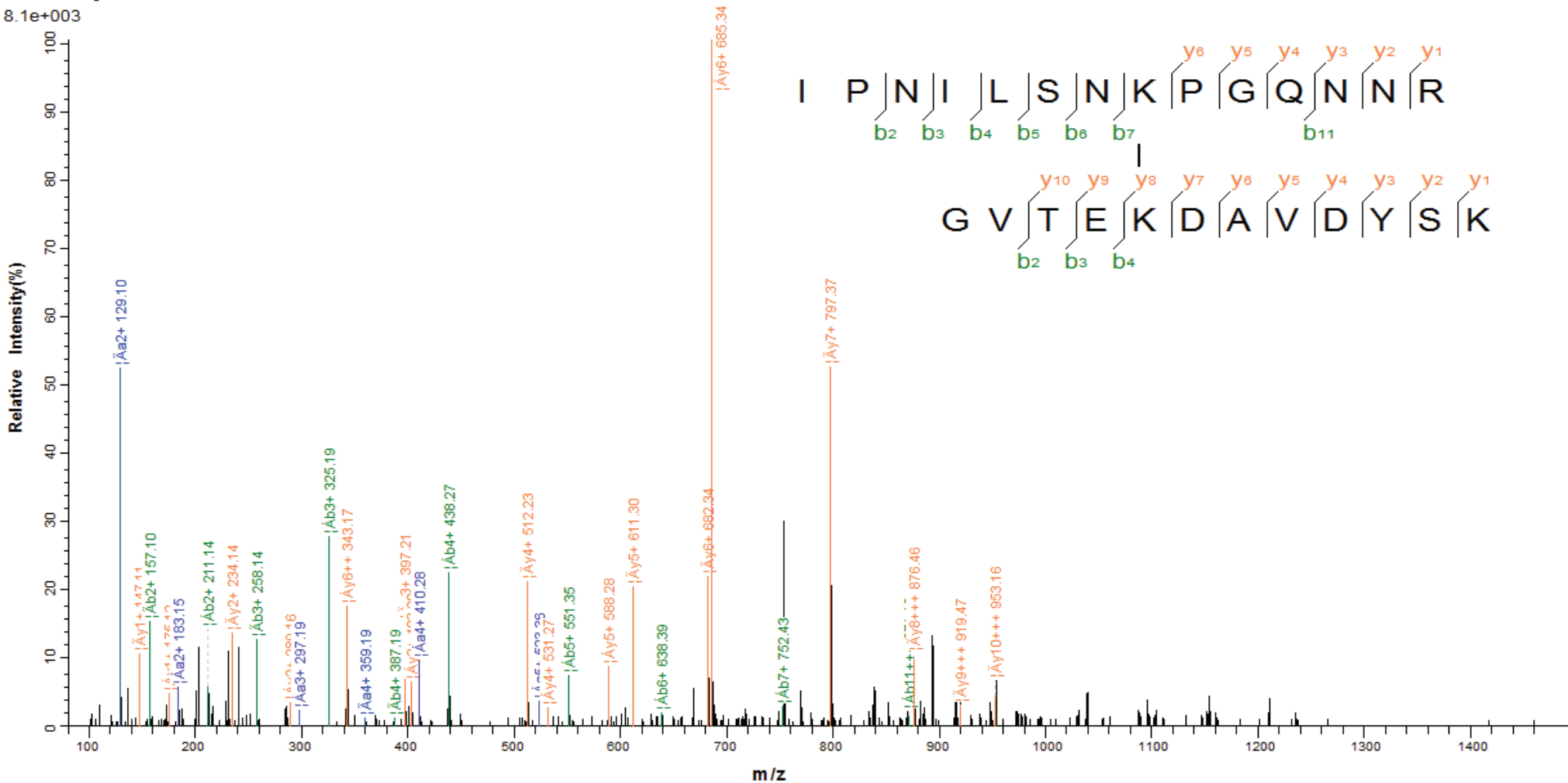


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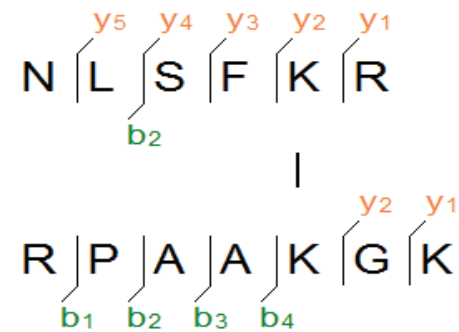
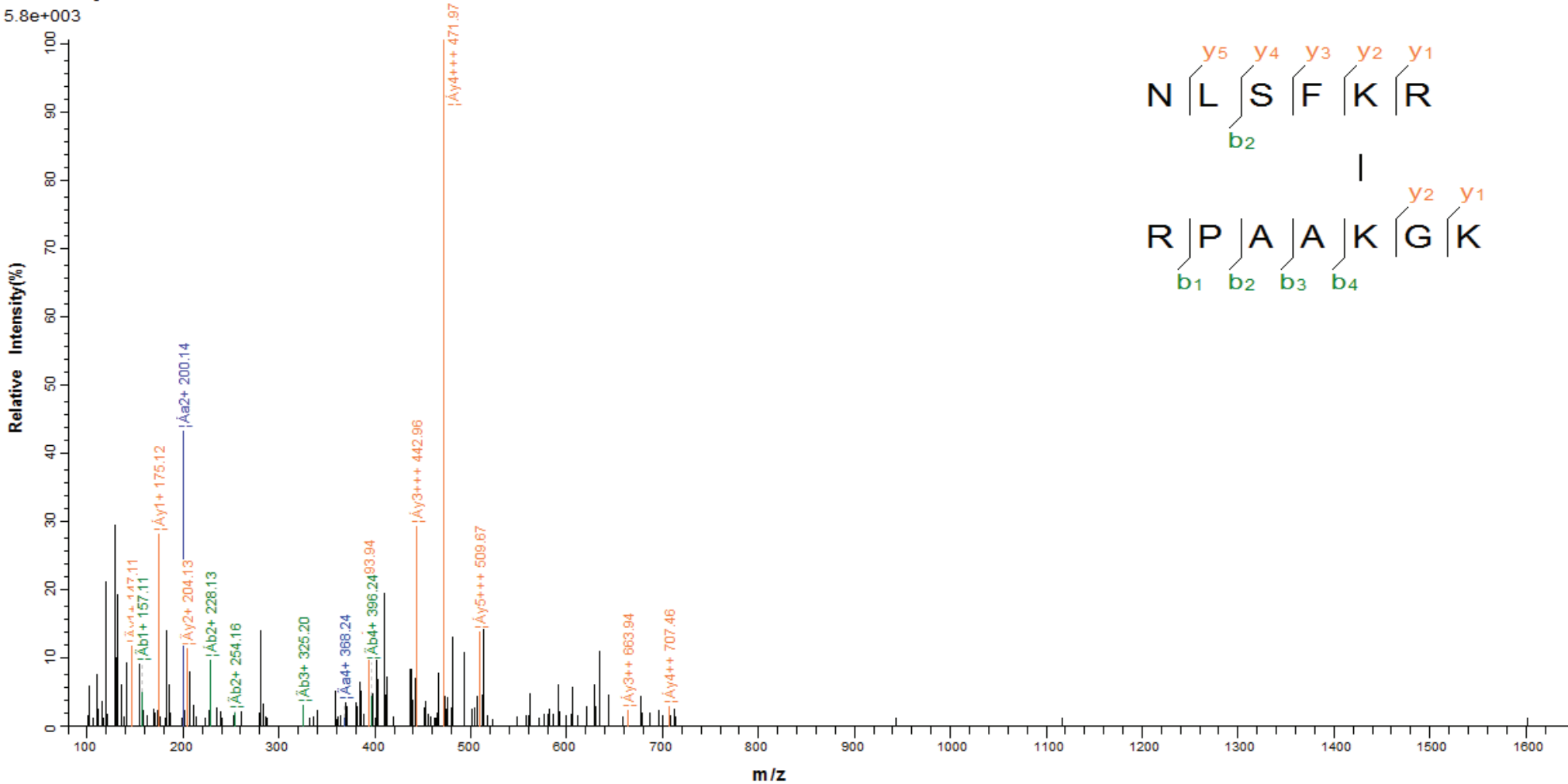
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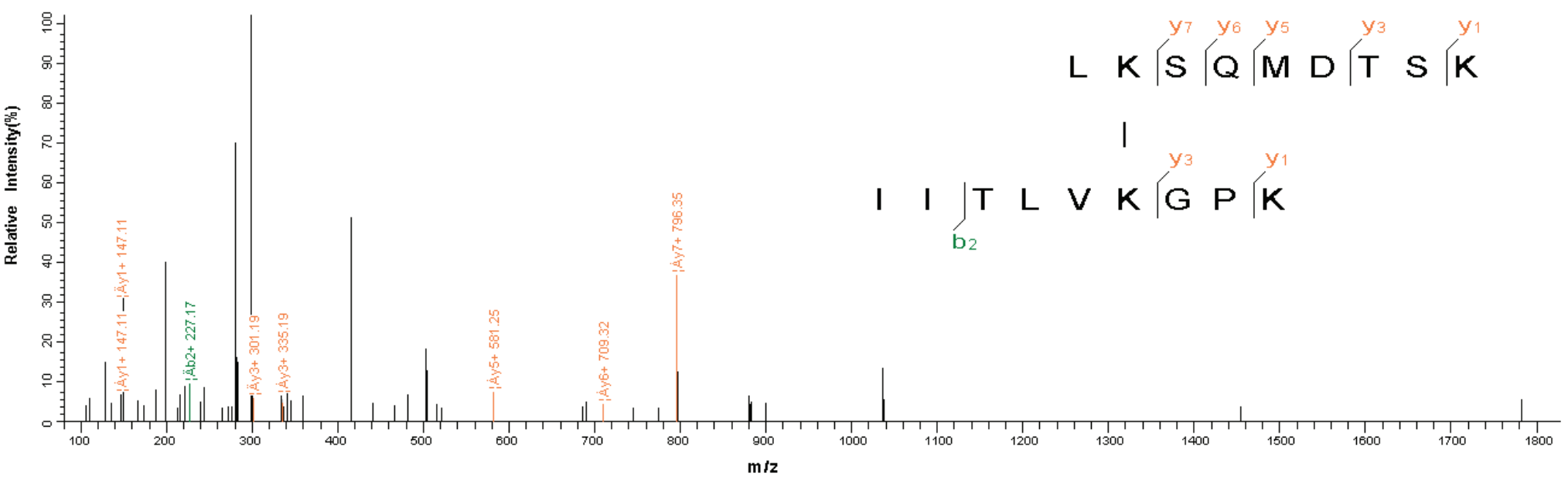
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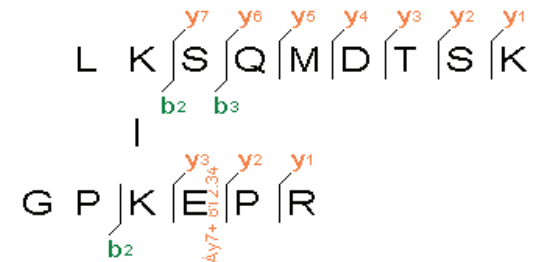
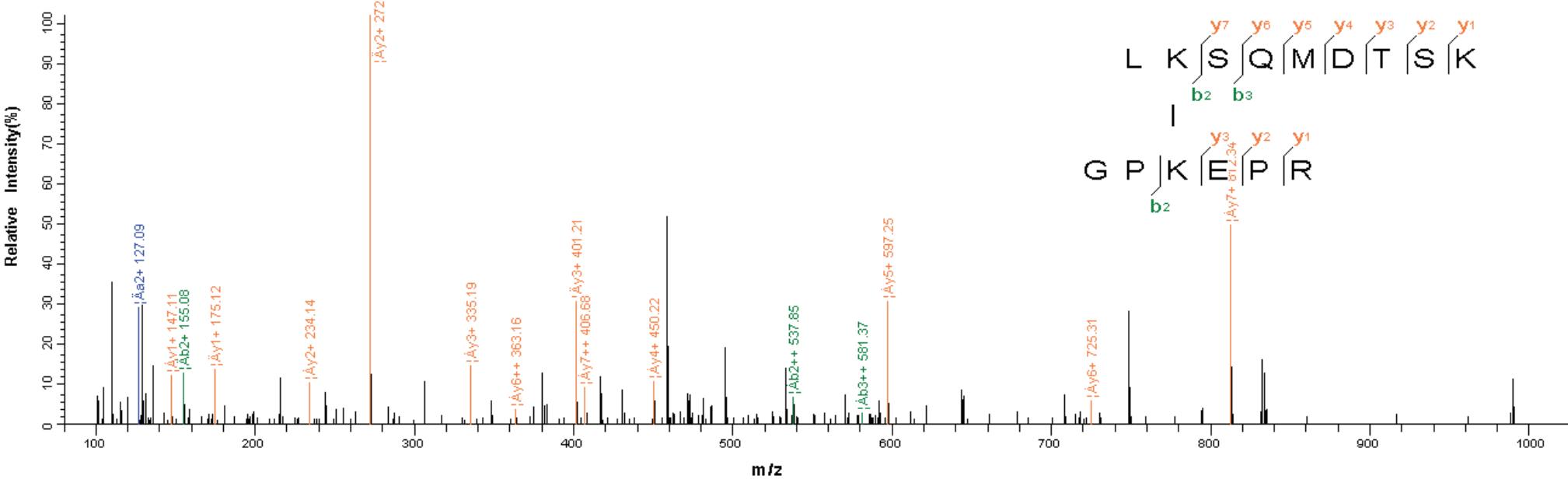
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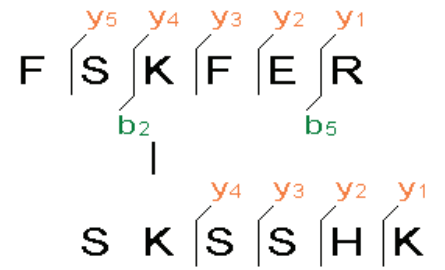
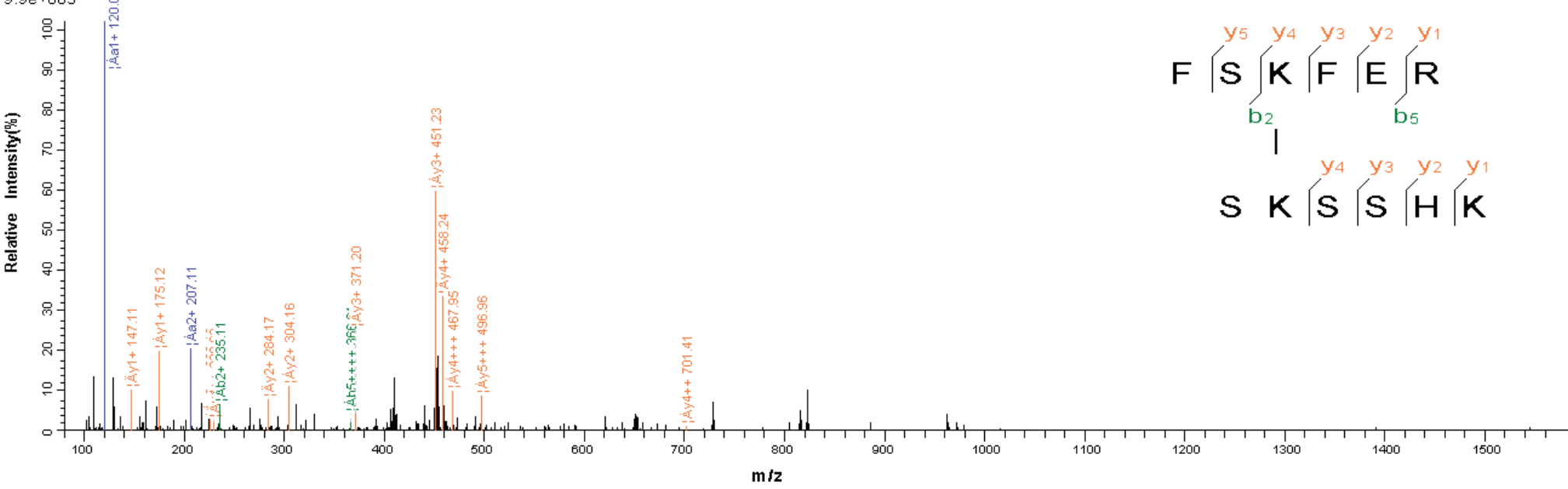
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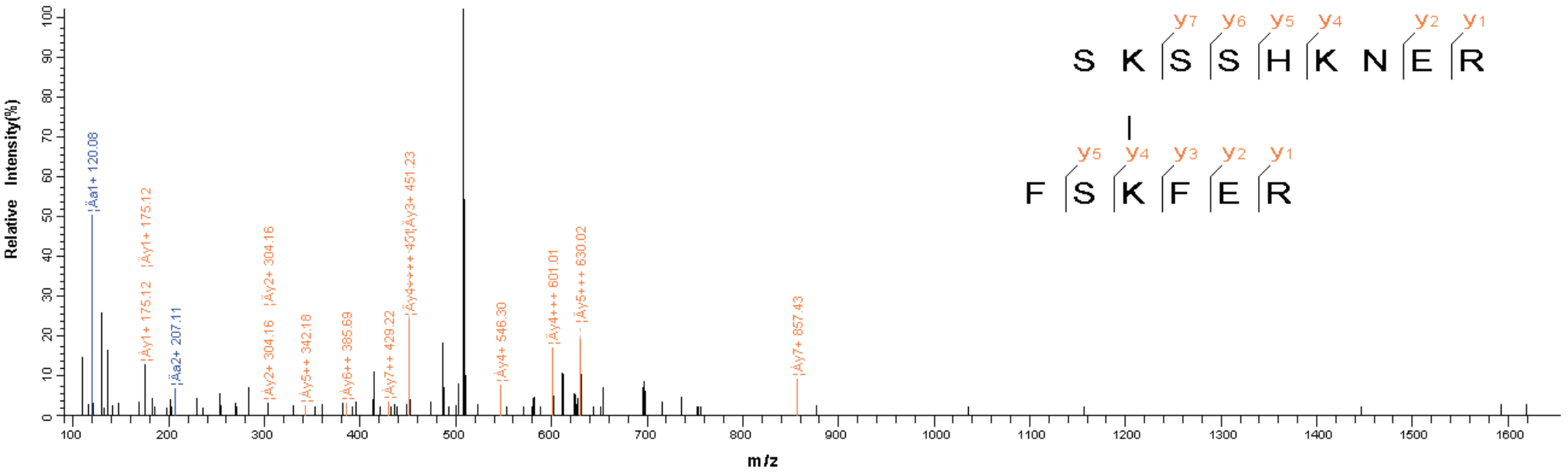
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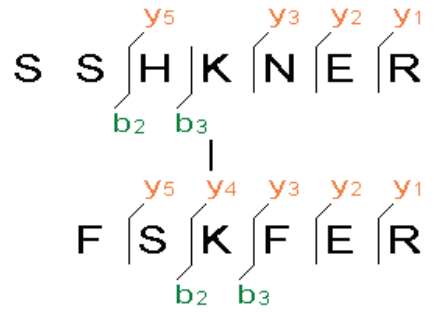
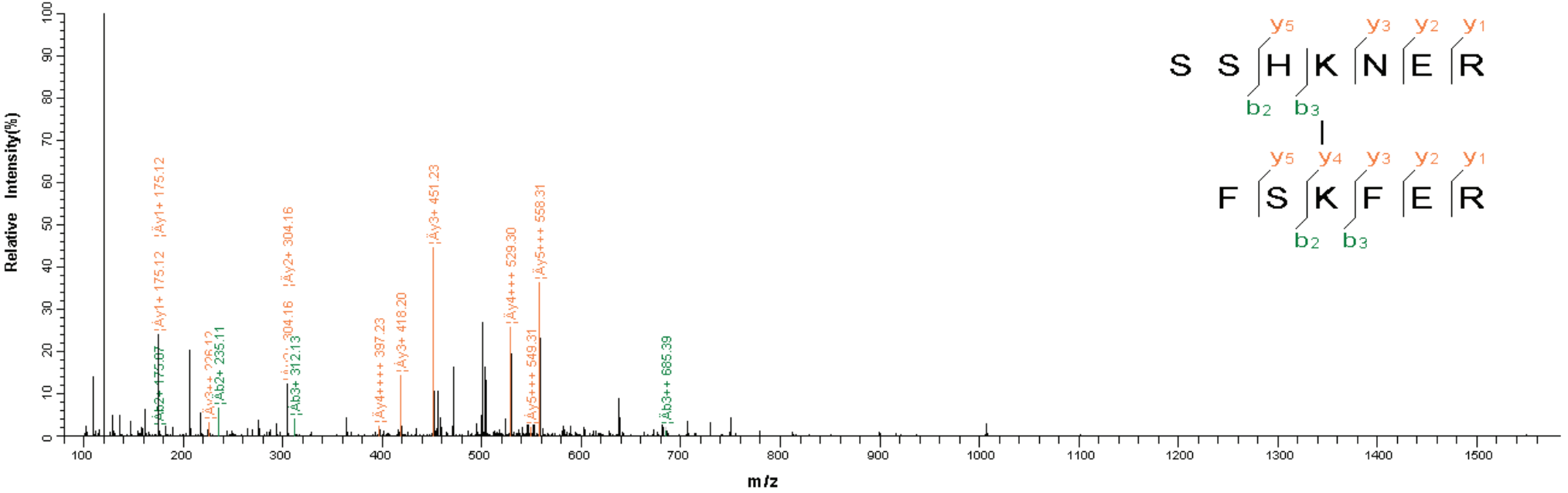
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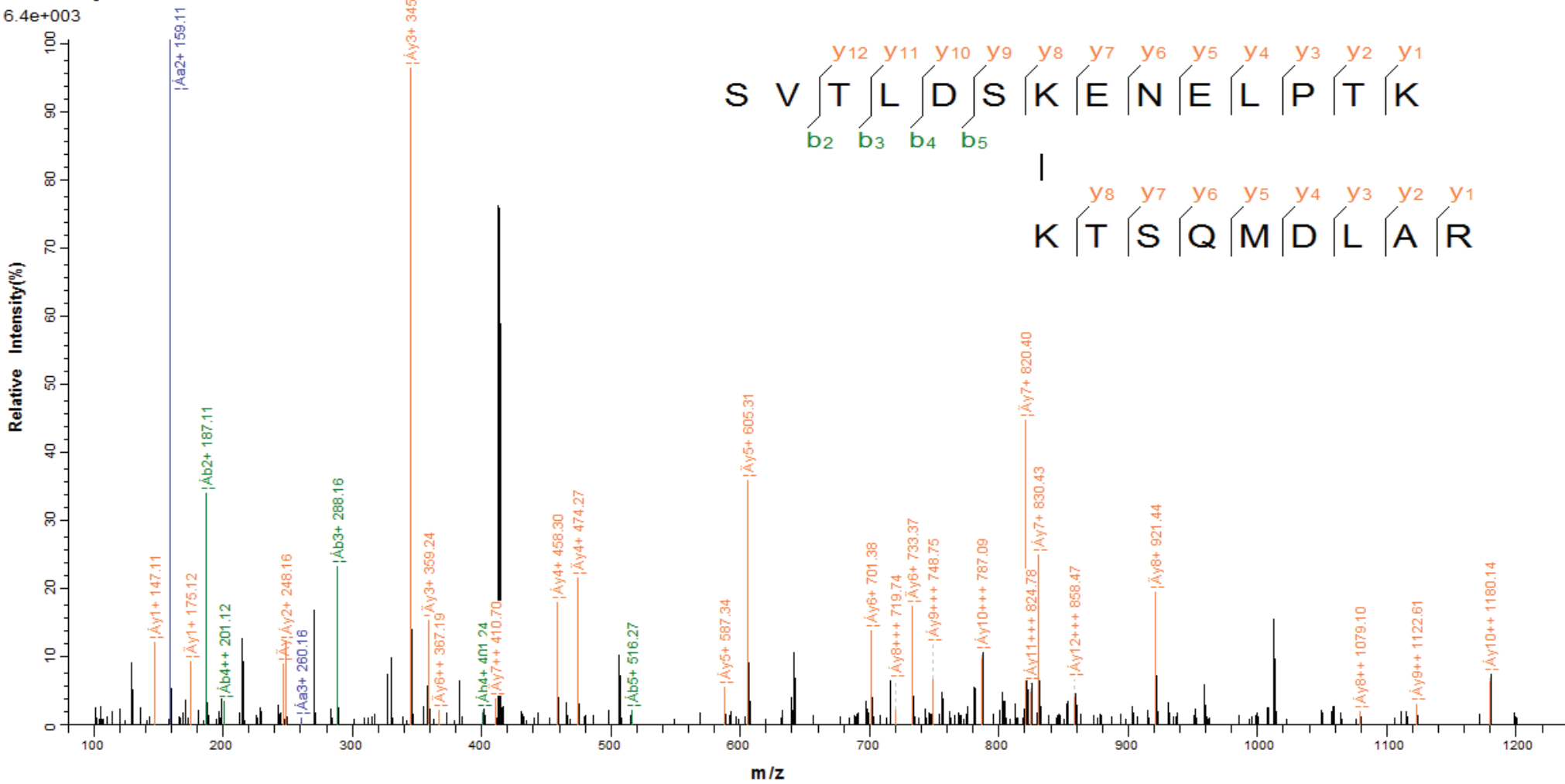
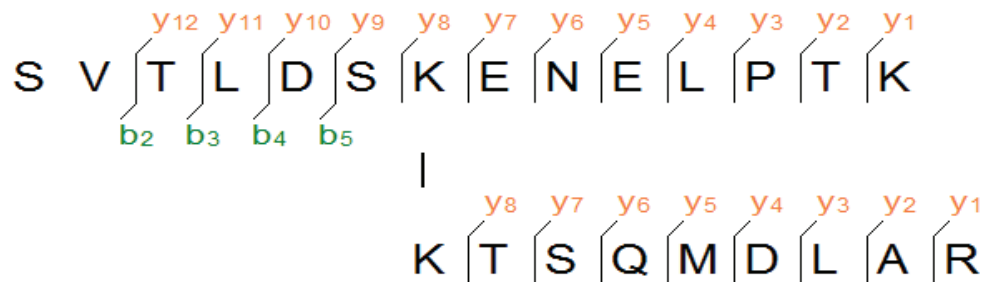
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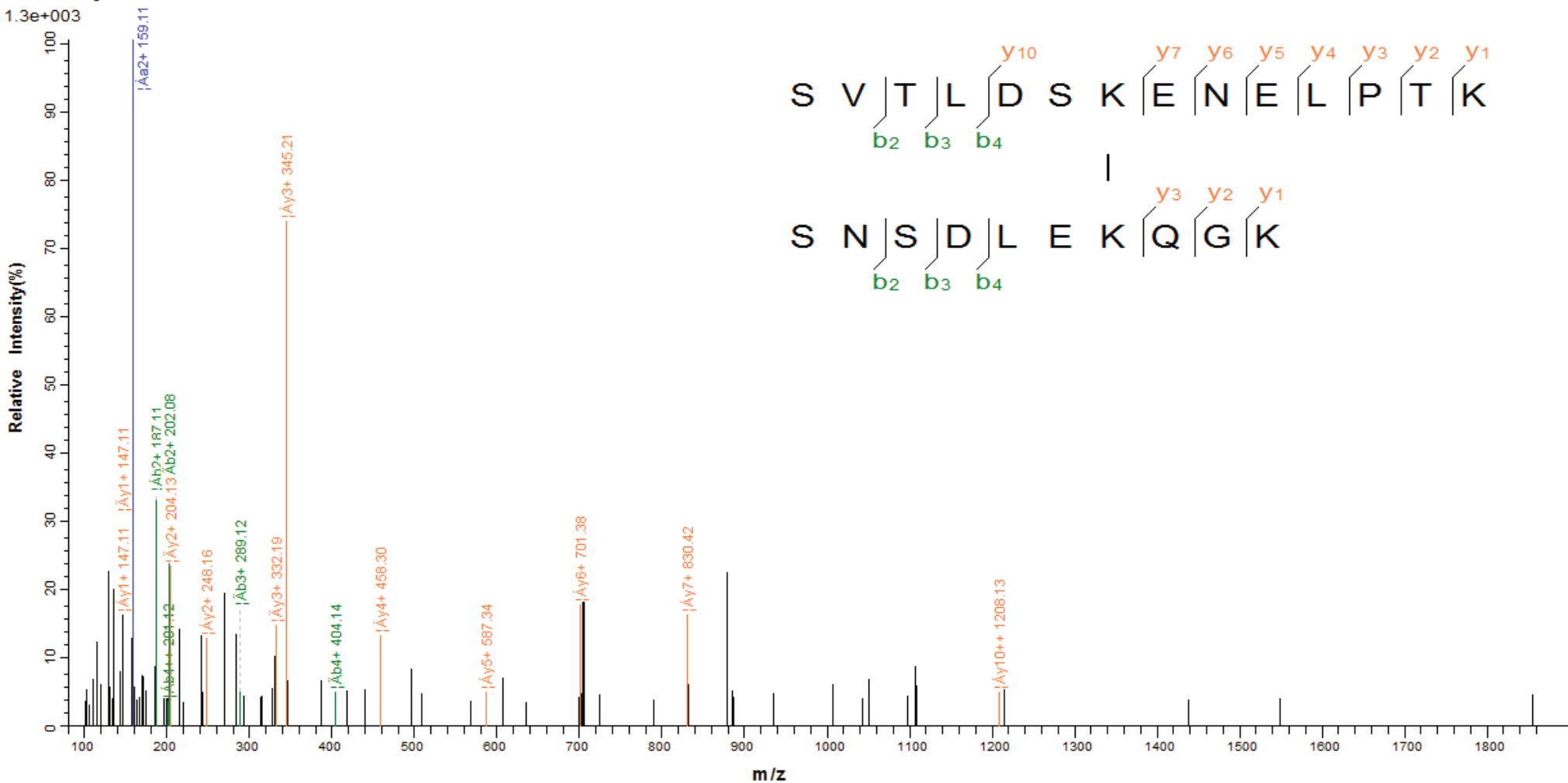
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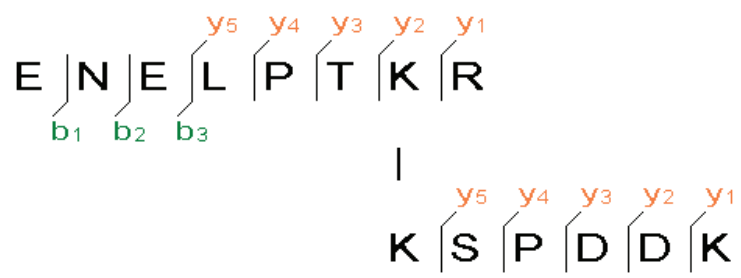
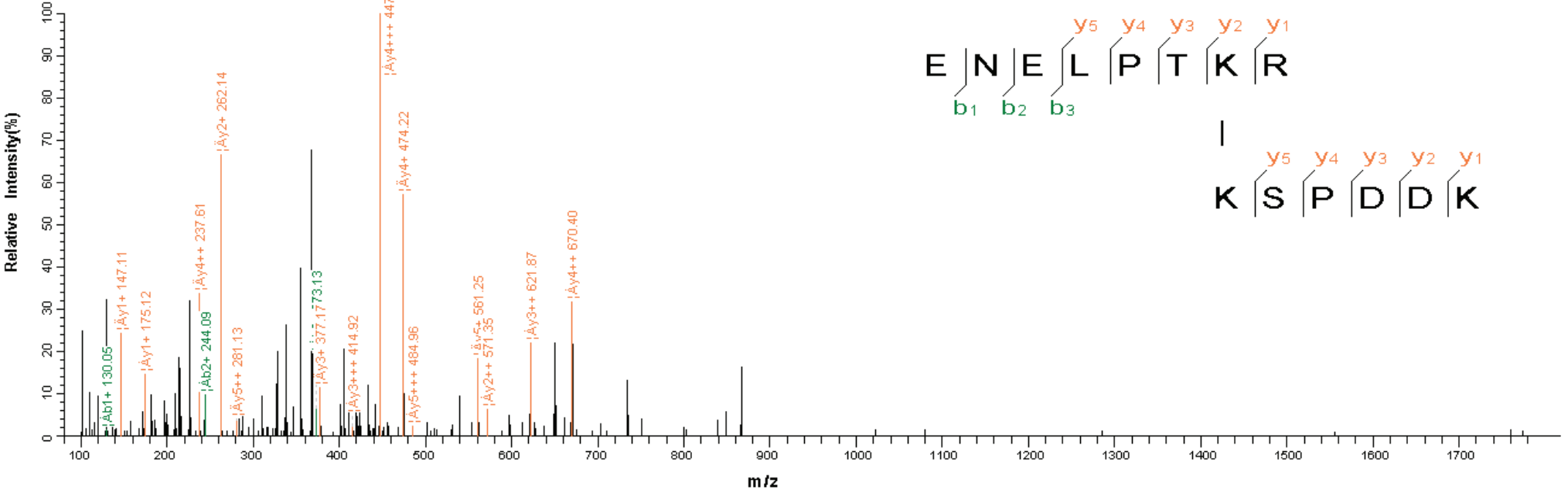
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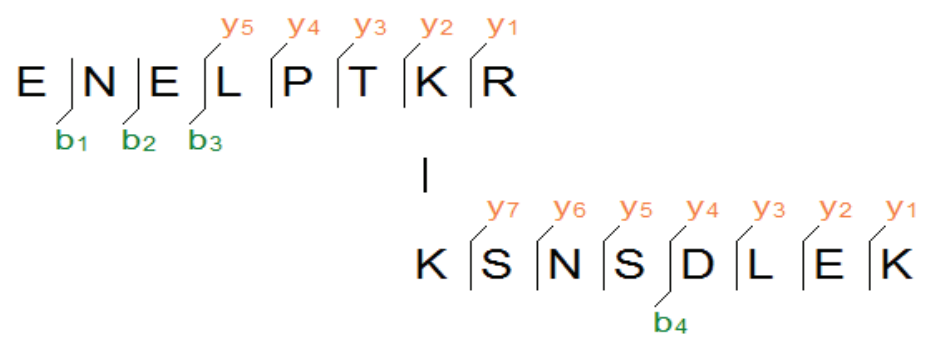
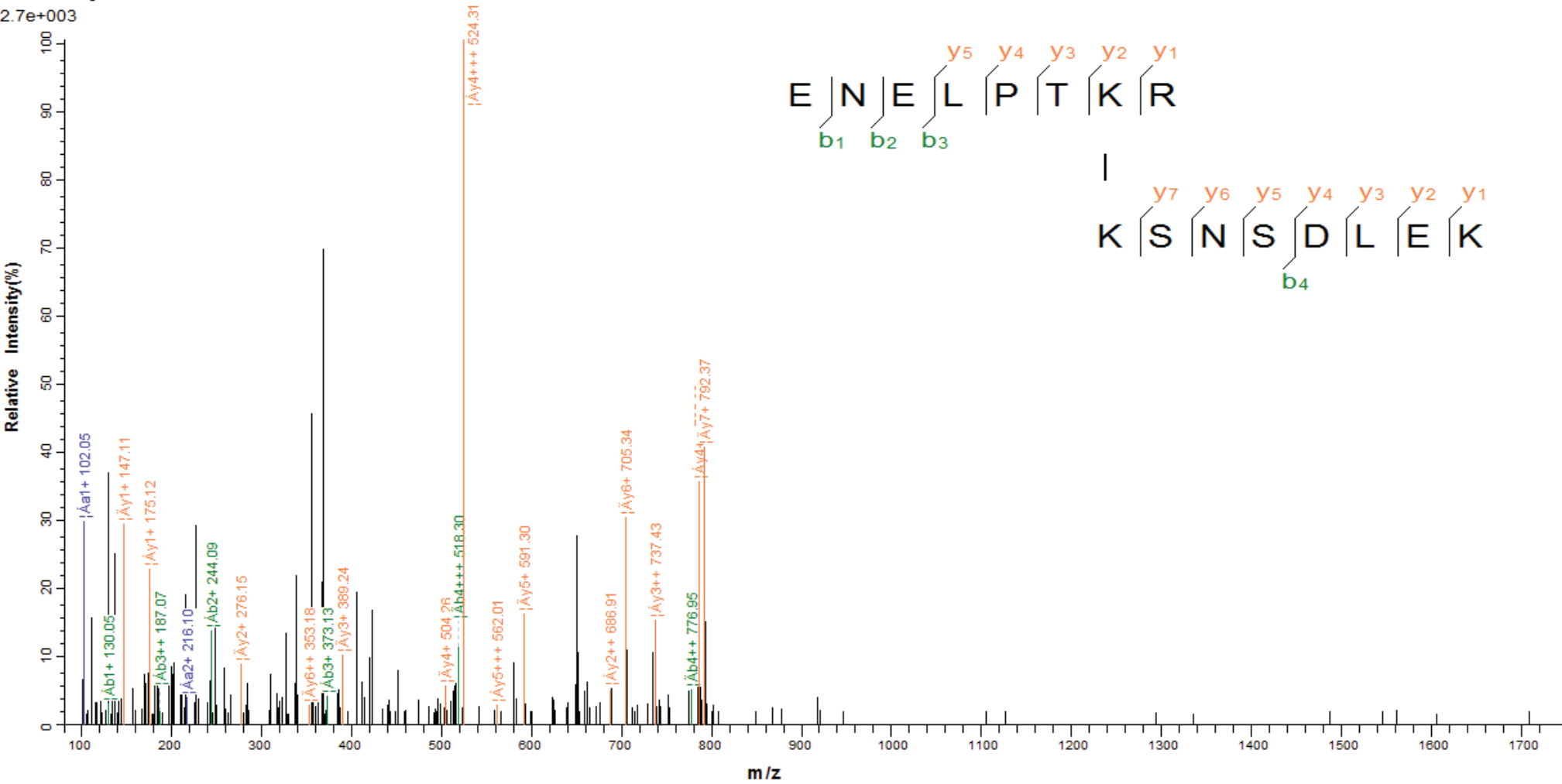
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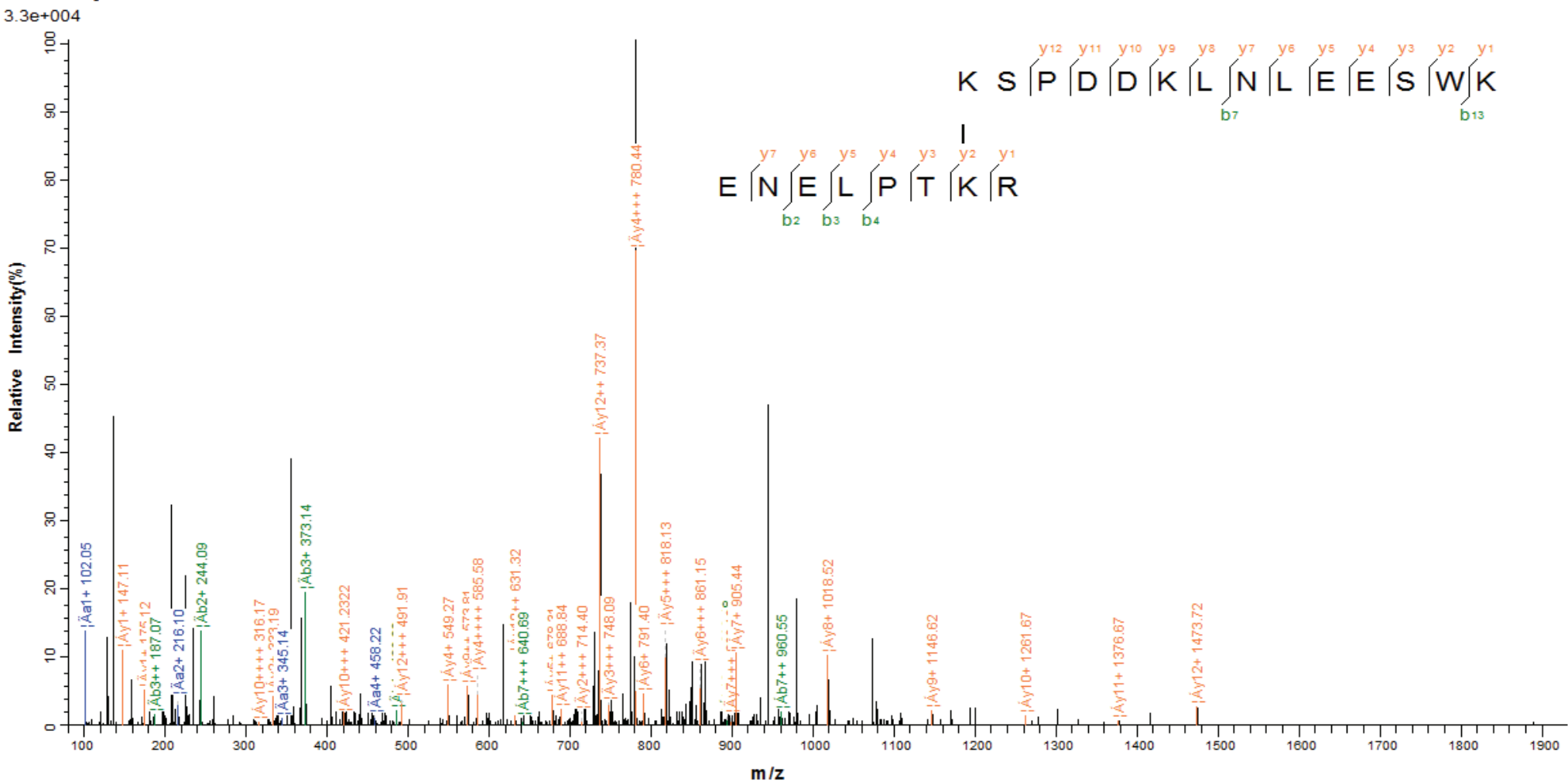
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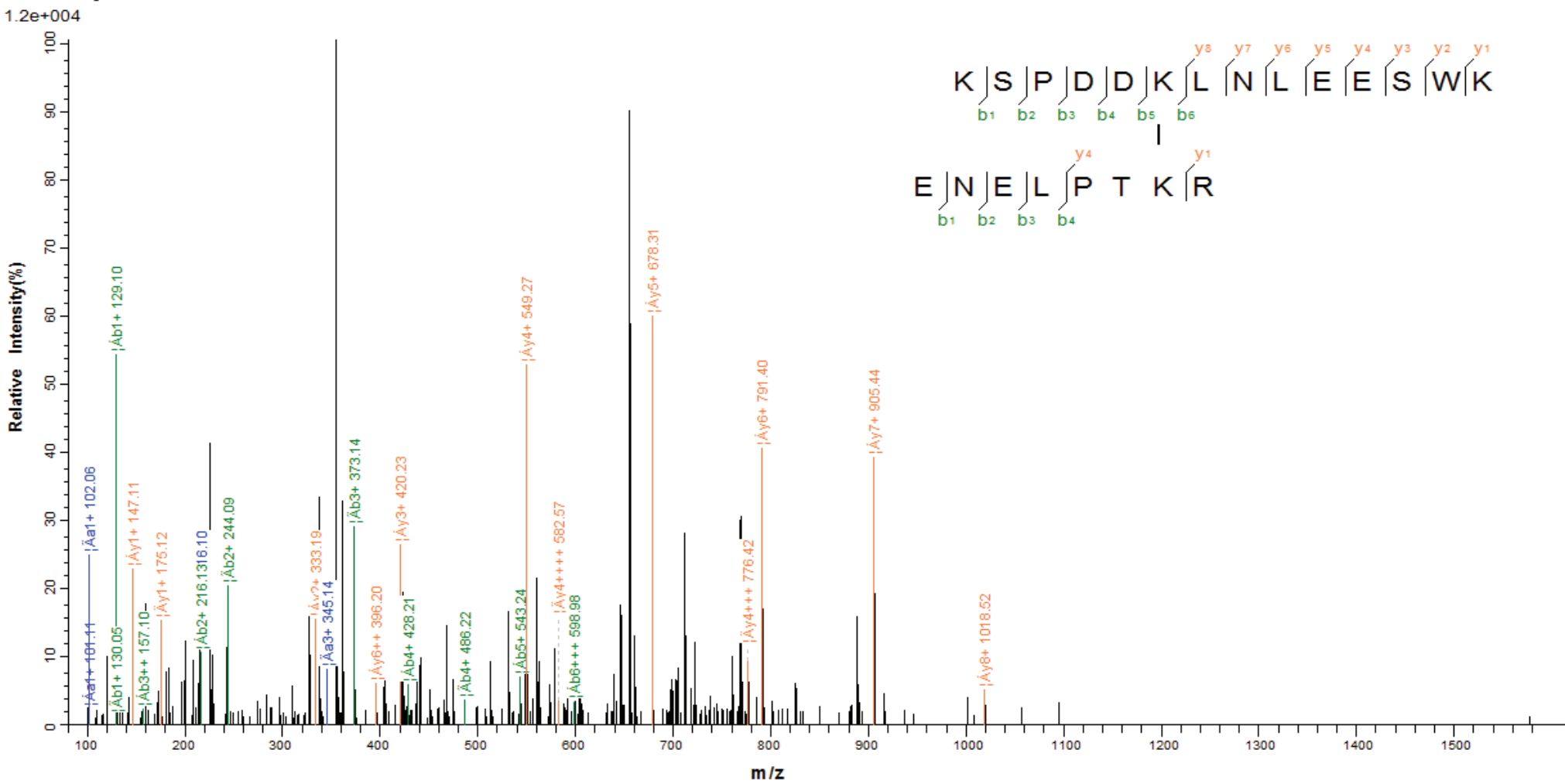
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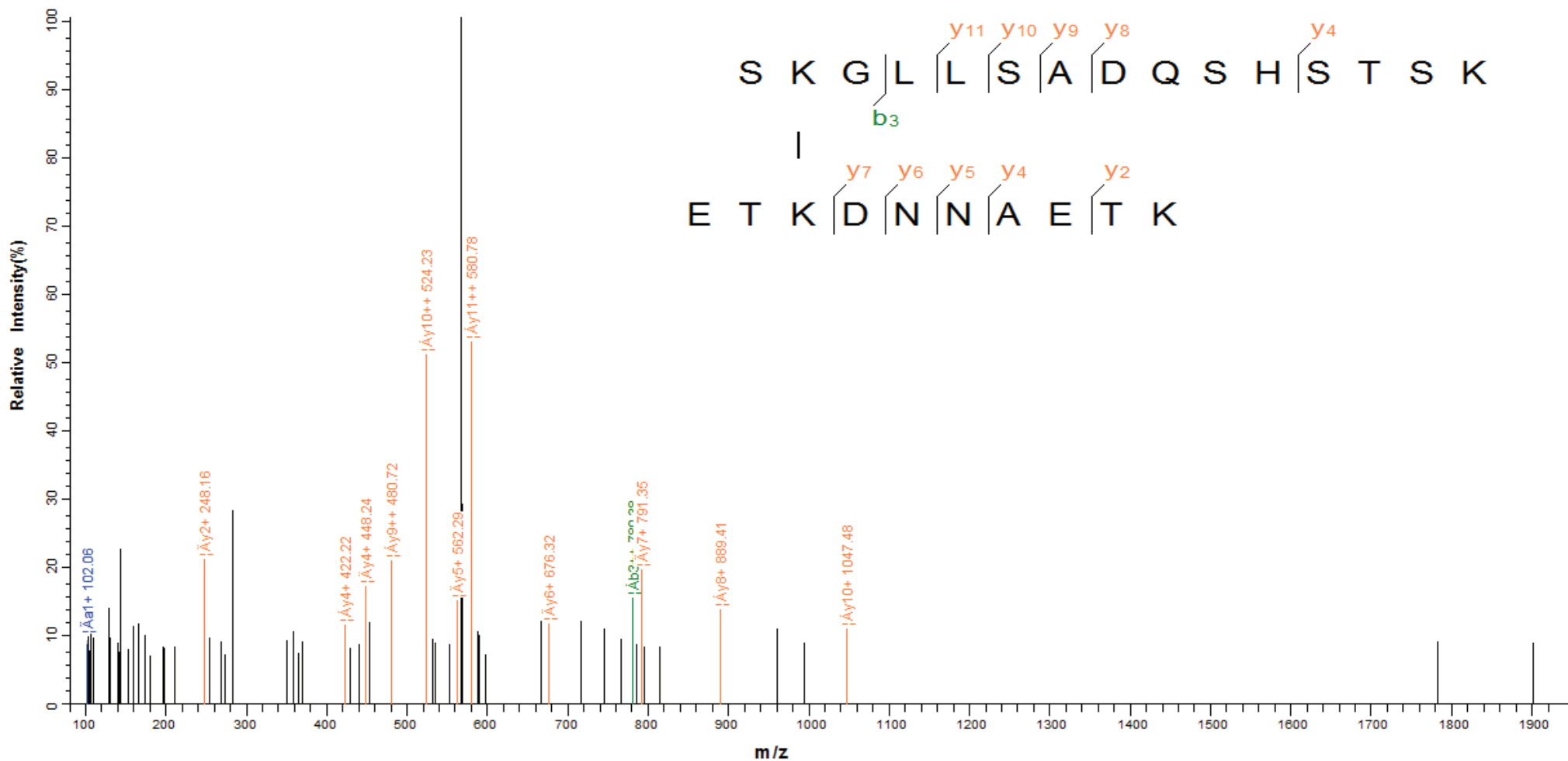


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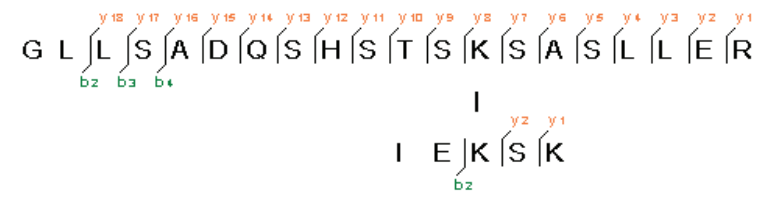
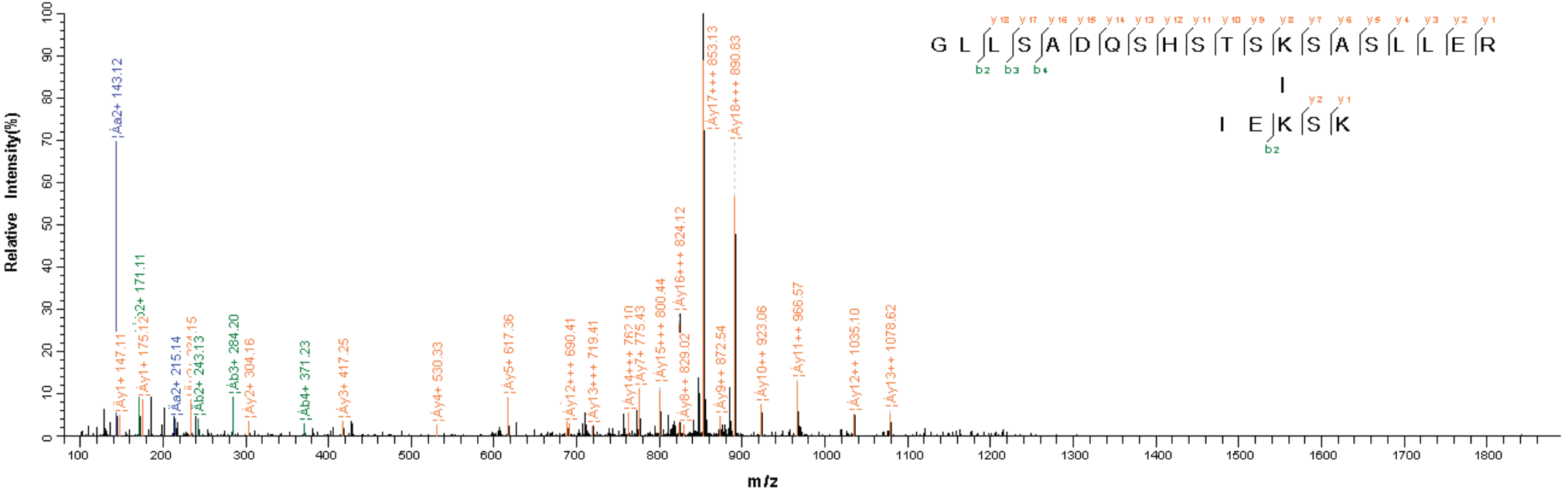


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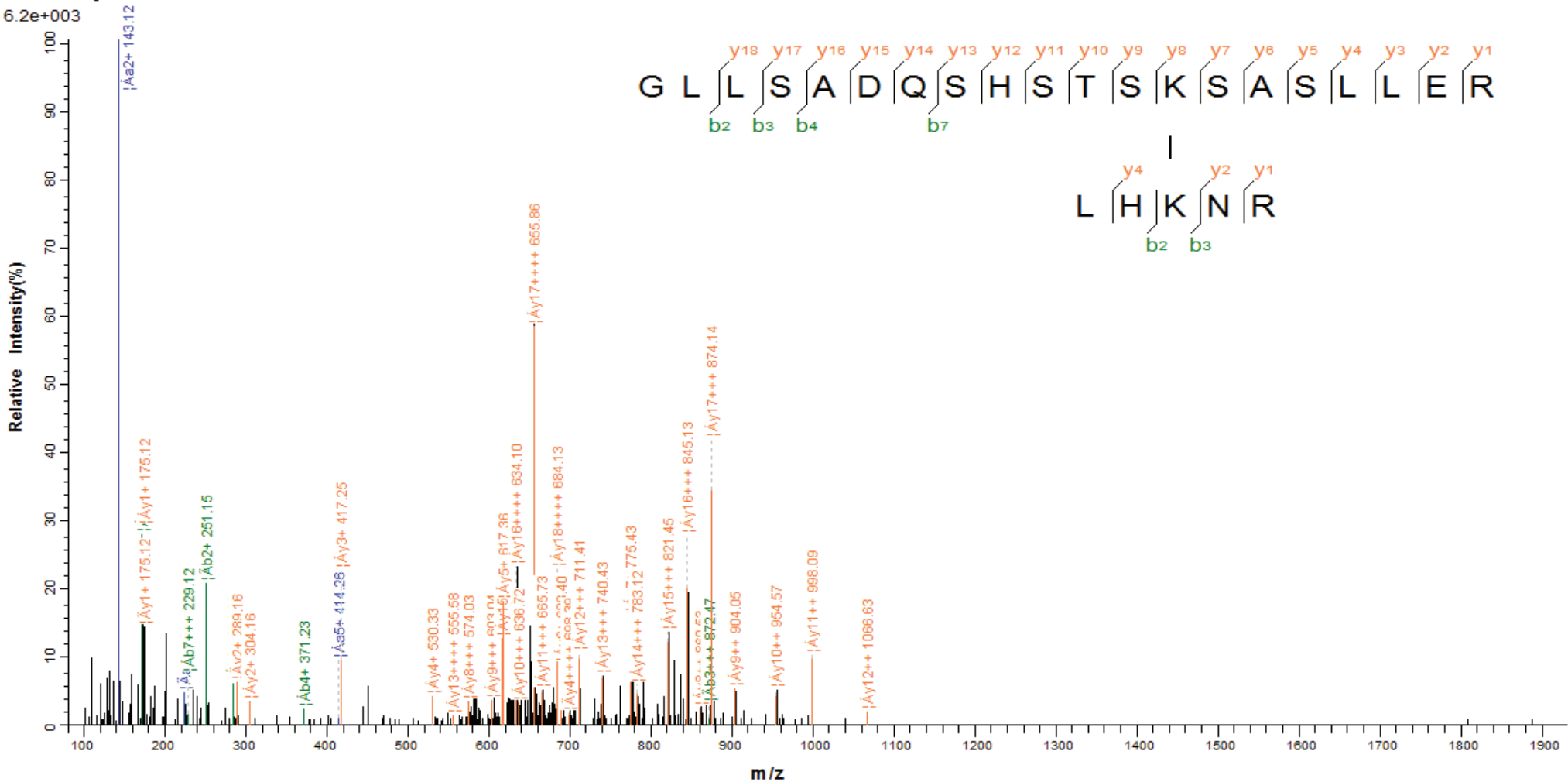
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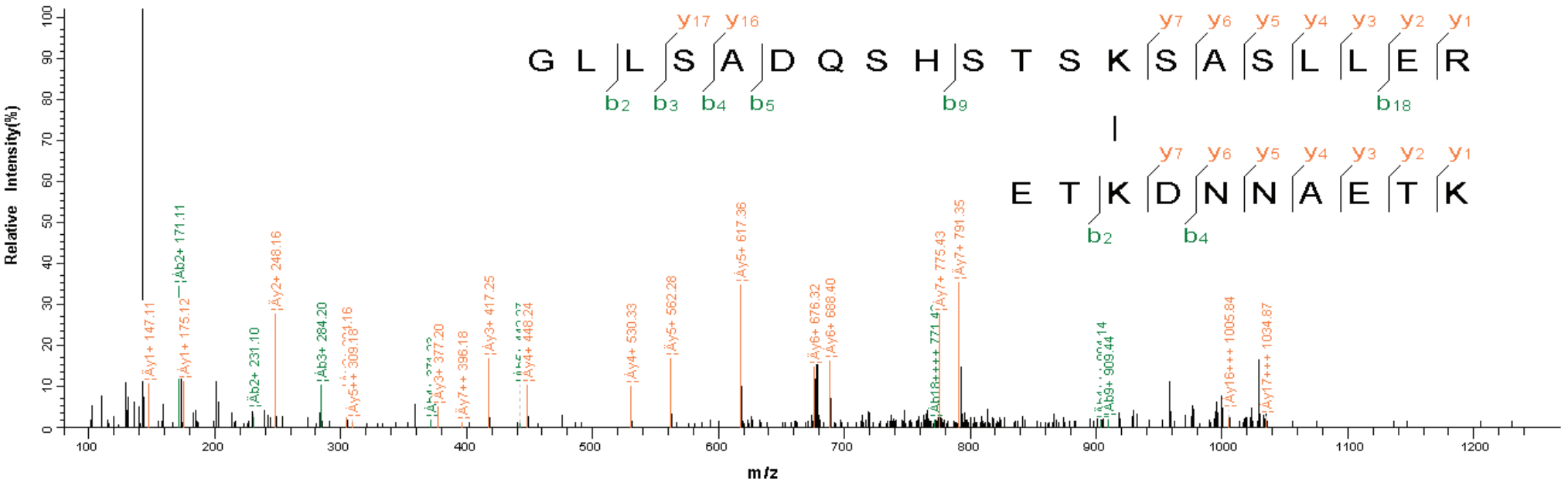
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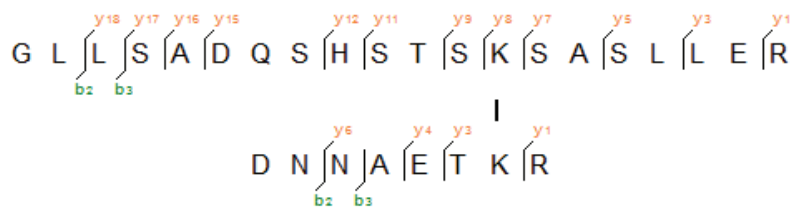
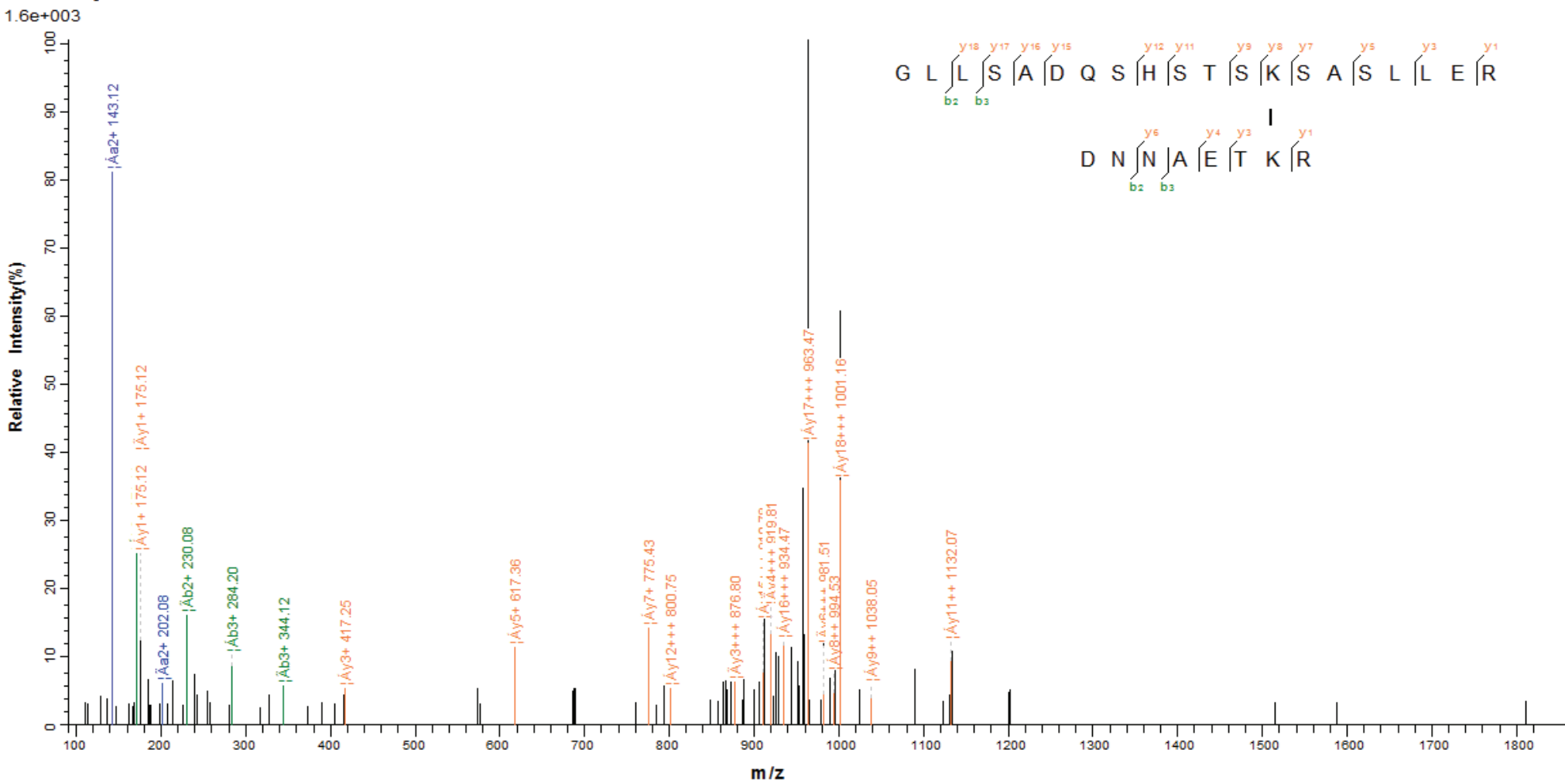
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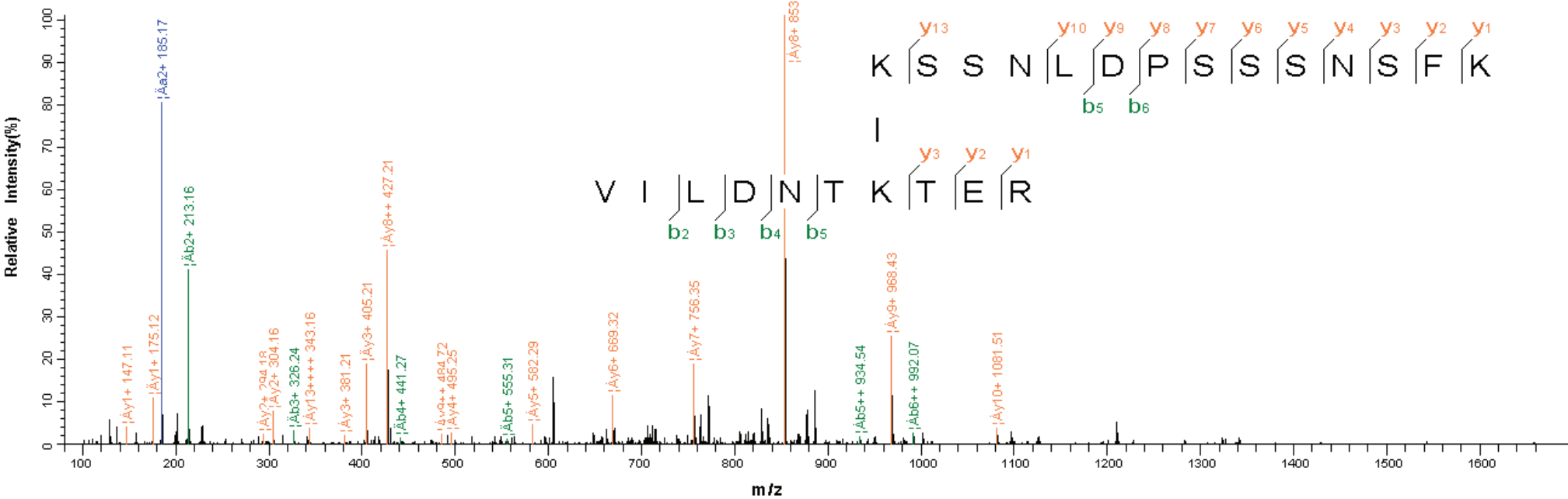
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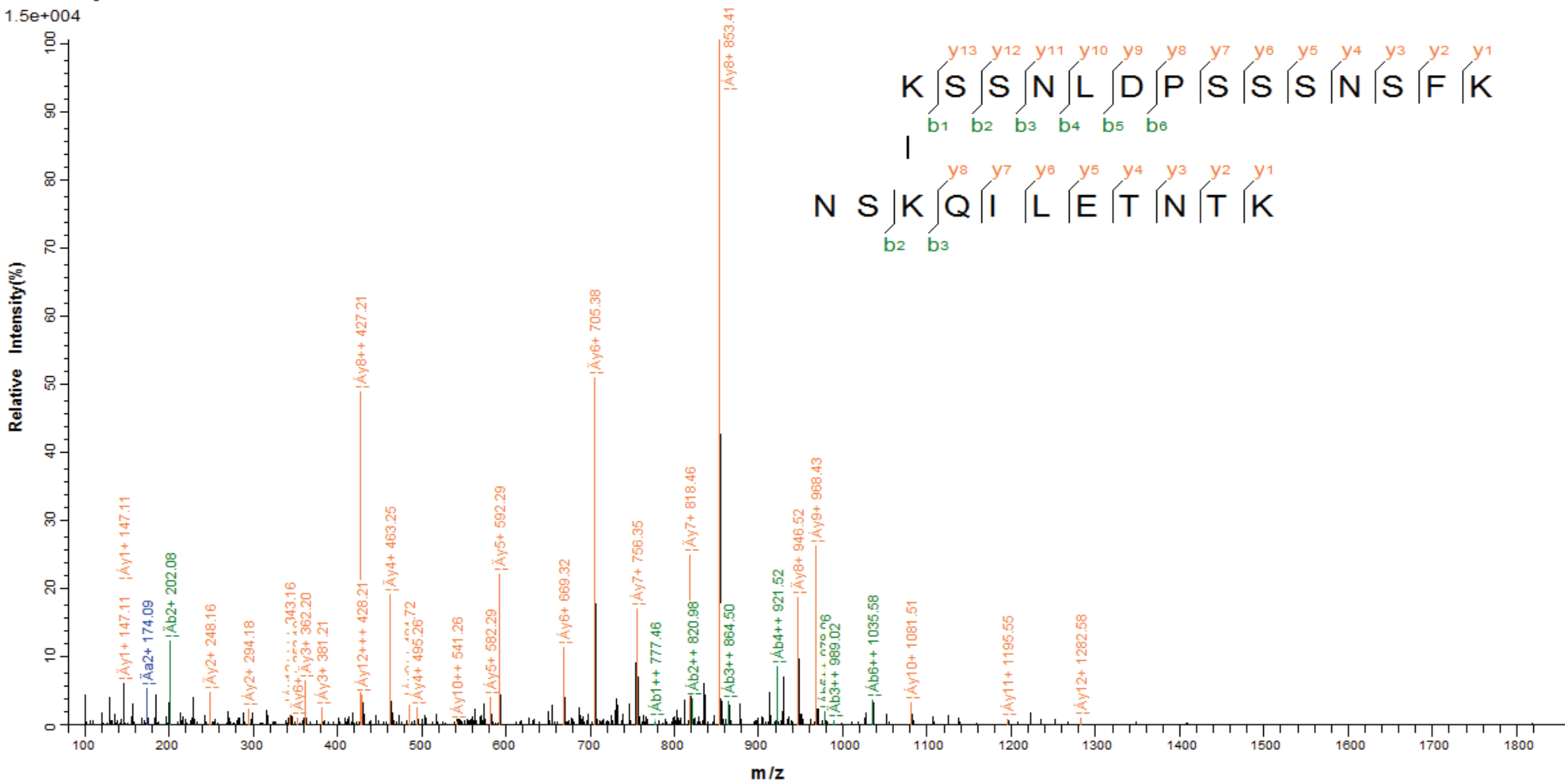
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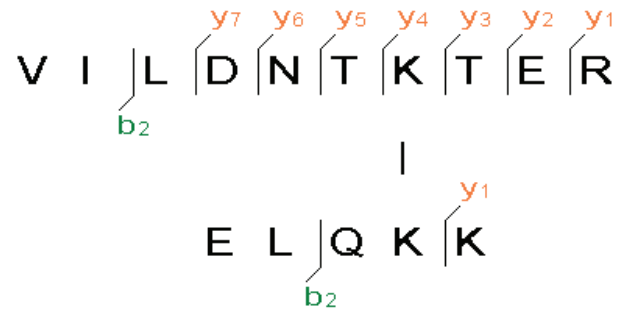
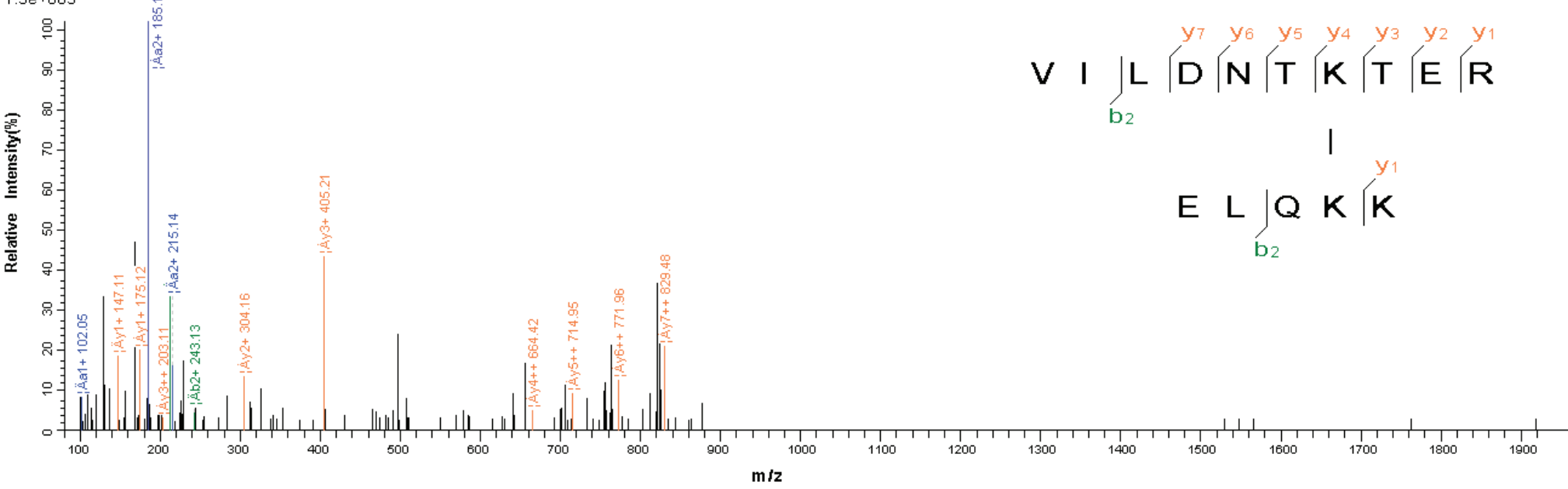
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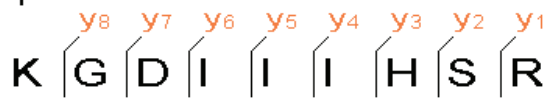
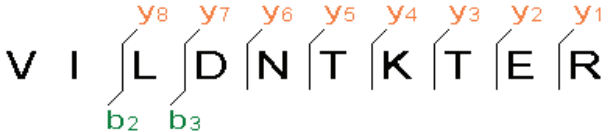
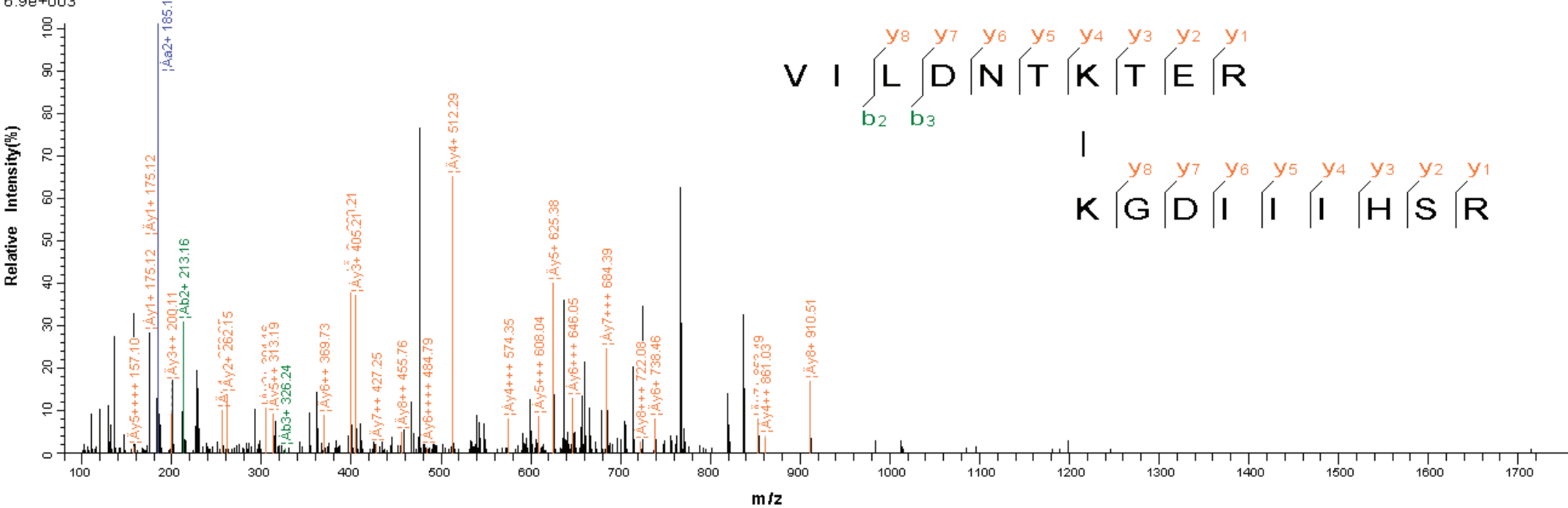
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exosome_A9_140503145423.2392.2392.4.dta Seq: VILDNTKTER-ELQKK

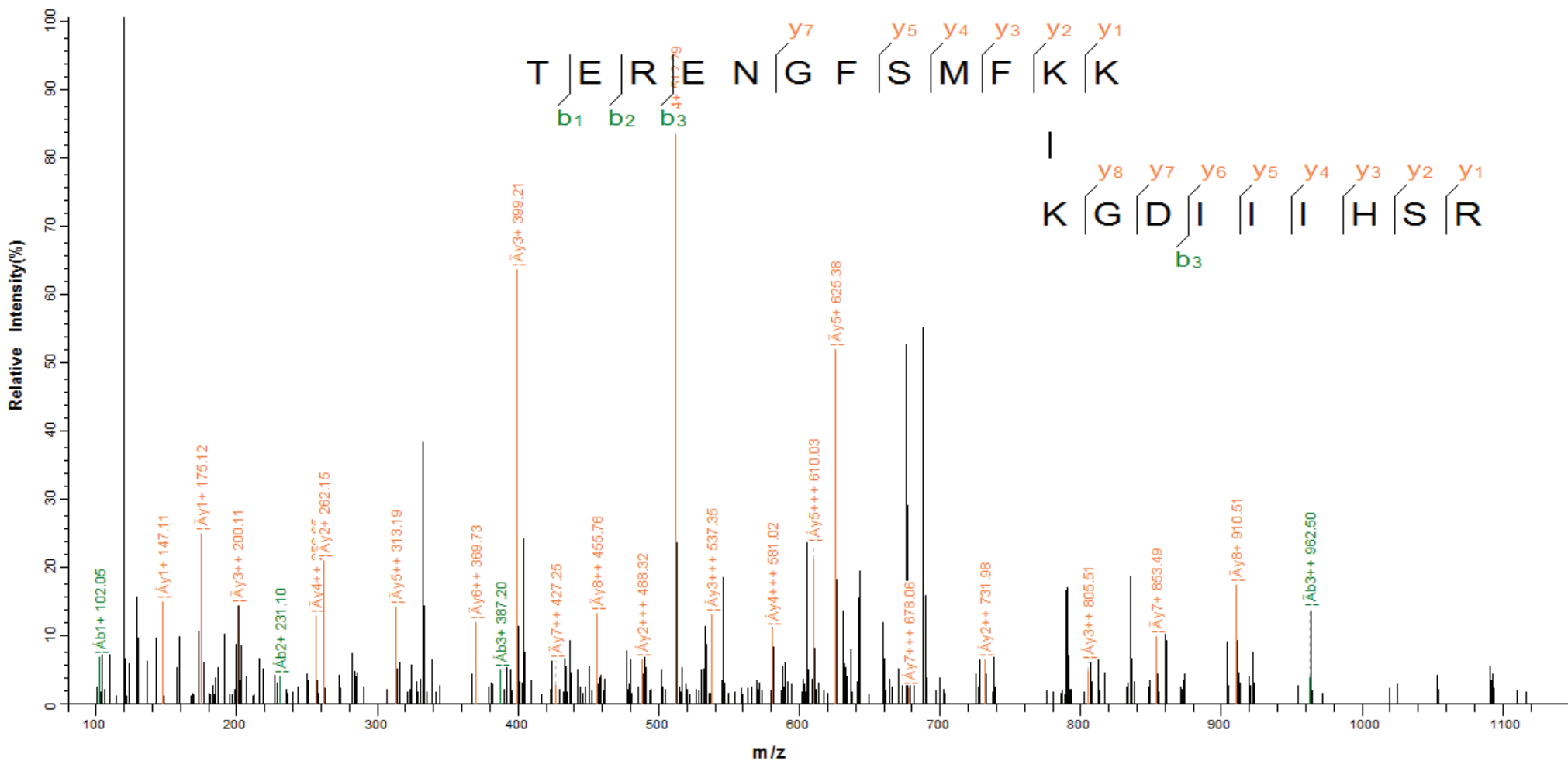


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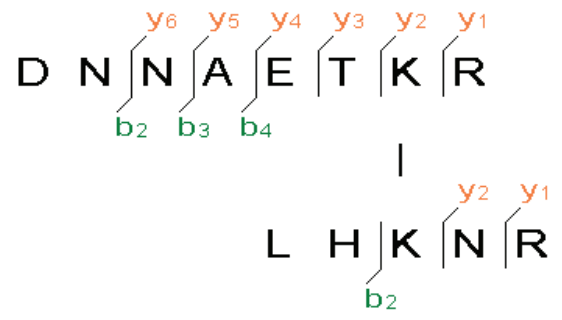
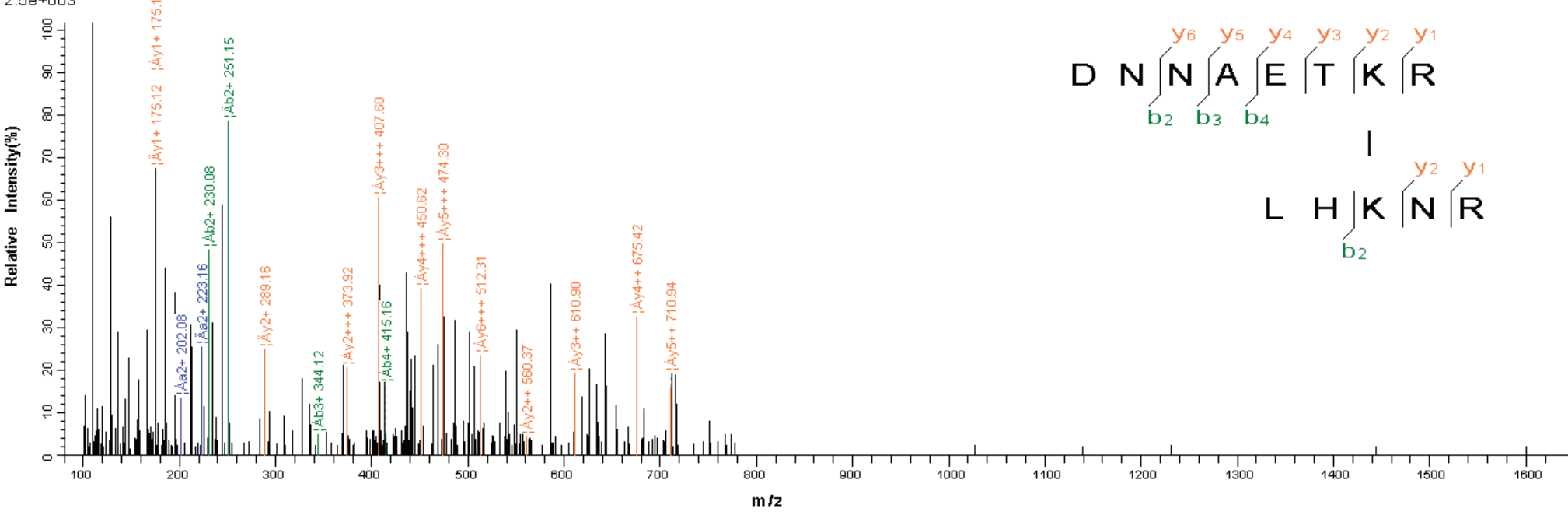


Seq: TERENGFNSMFKK

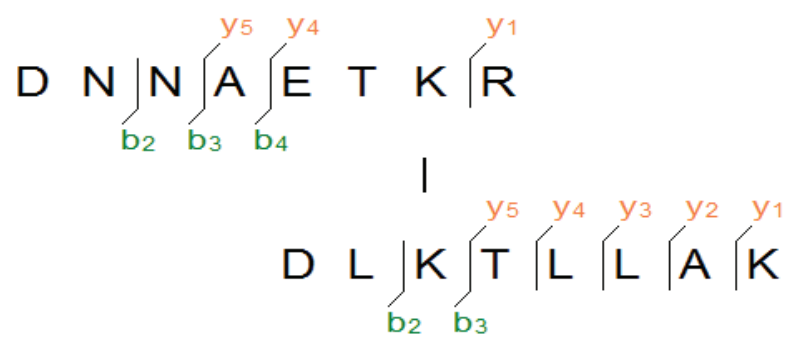
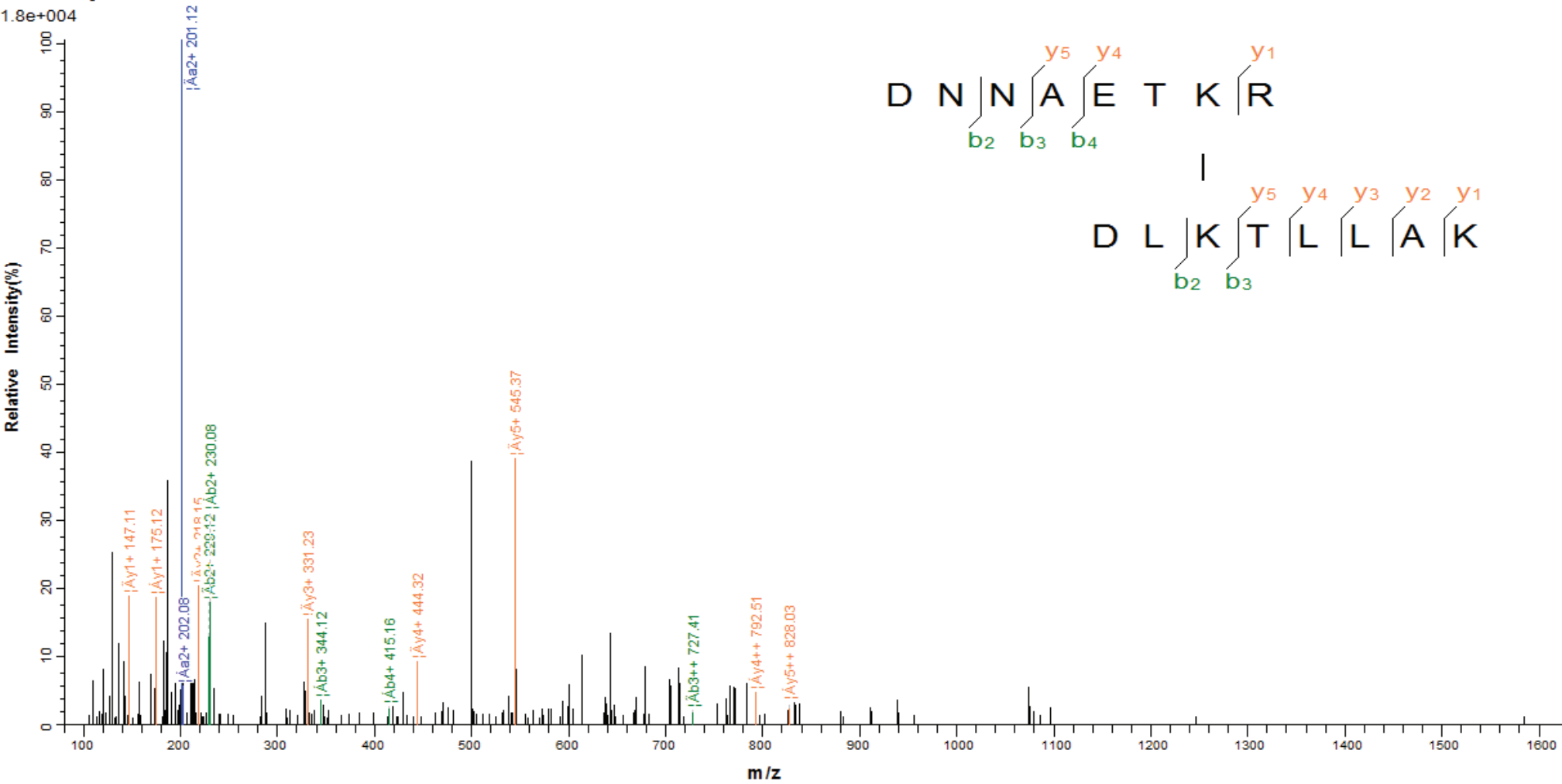
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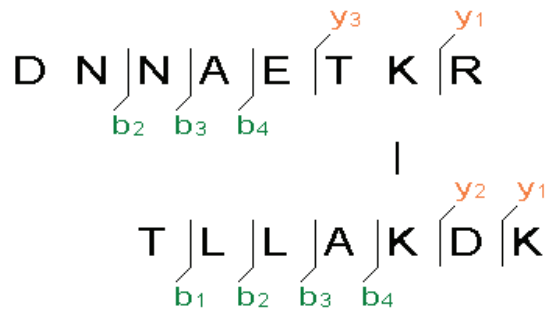
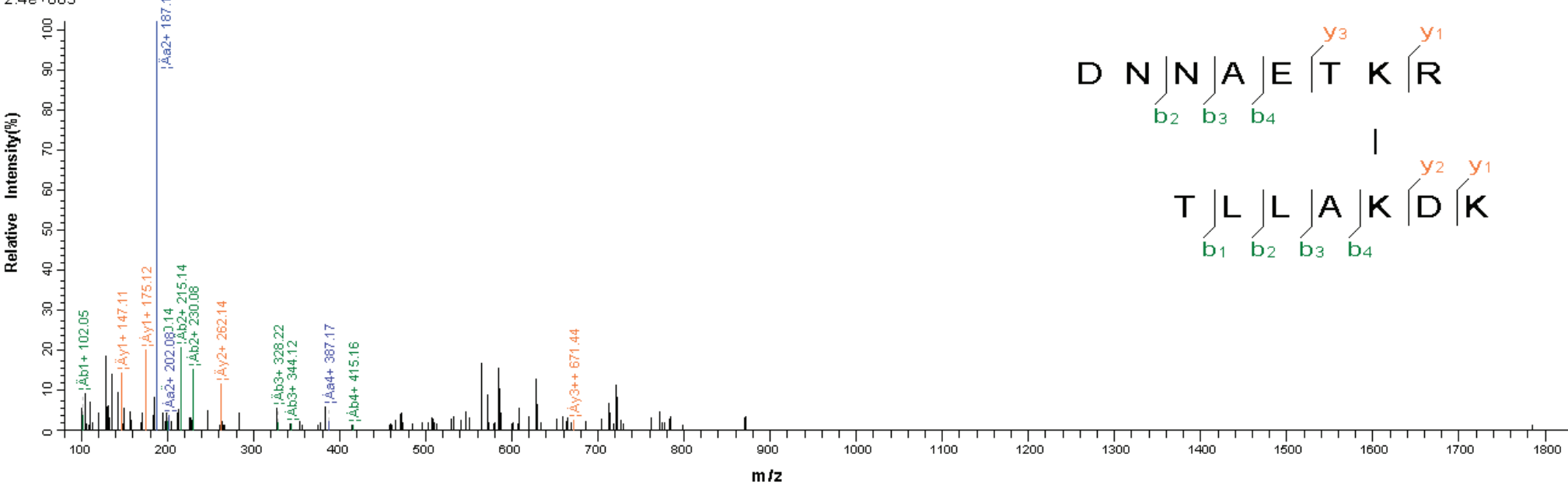
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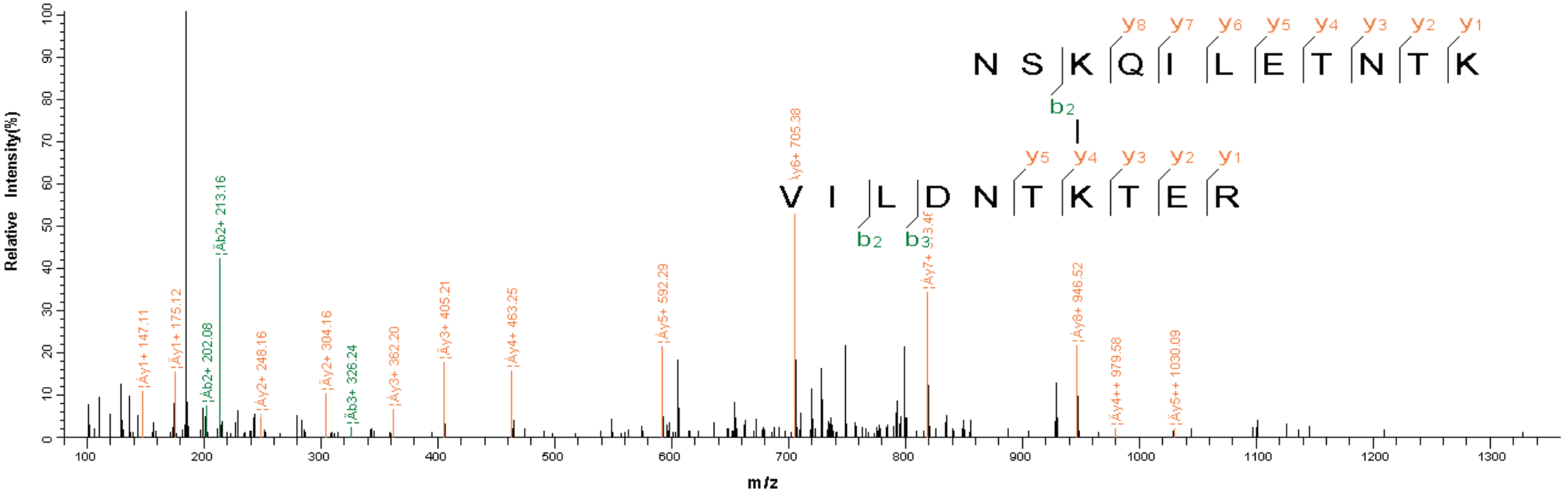
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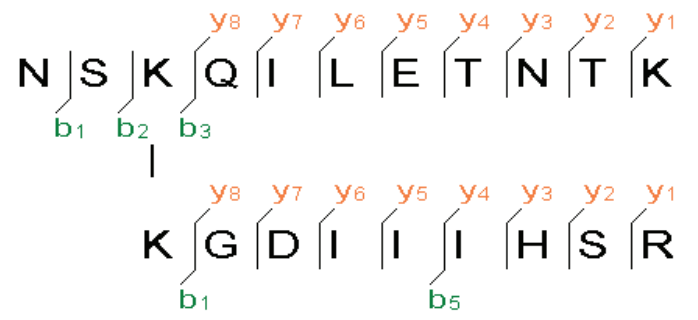
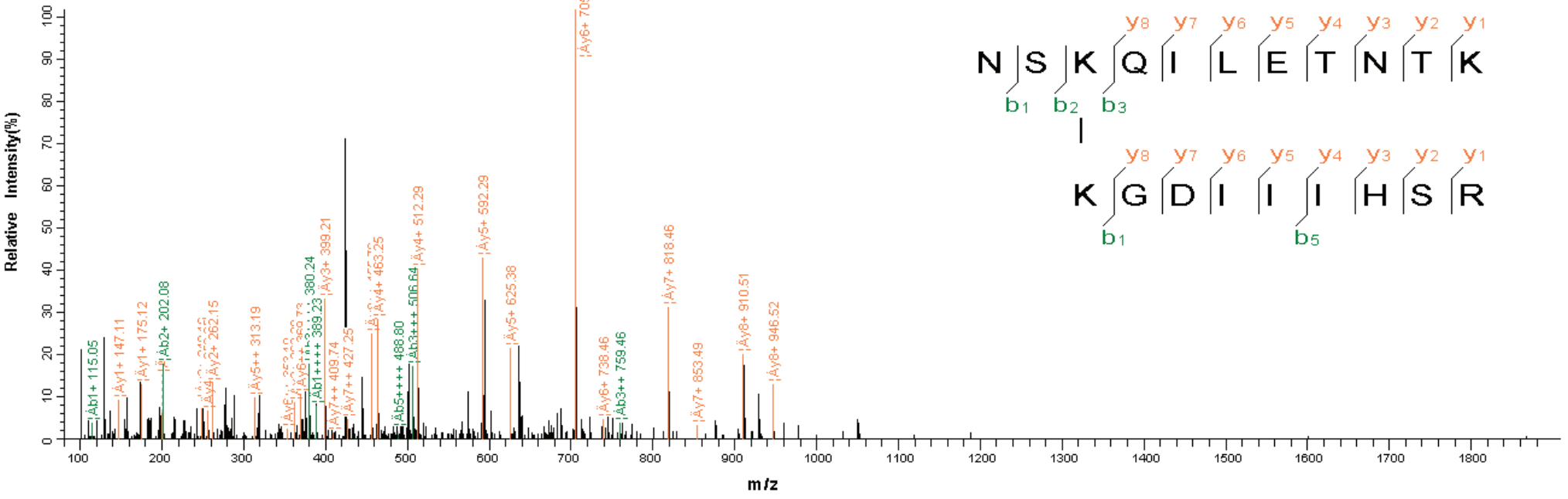
exosome_A9 140503145423.2027.2027.4.dta Seq: DNNAETKR-TLLAKDK



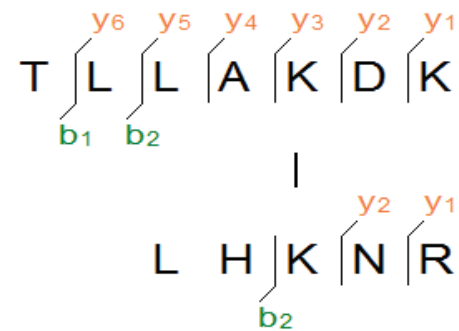
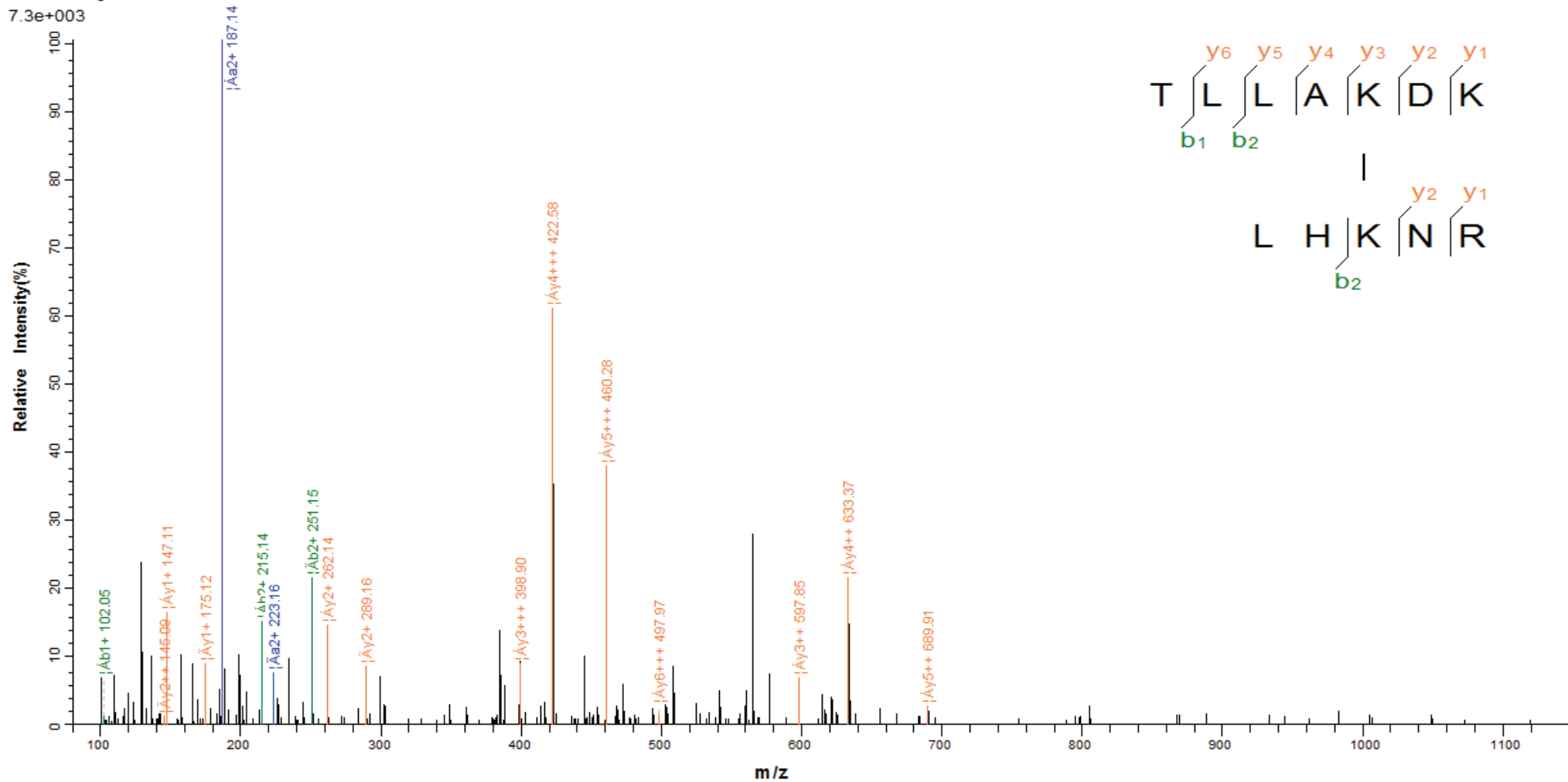
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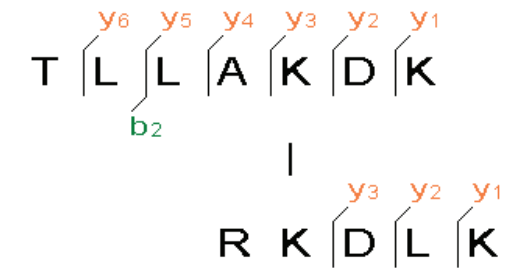
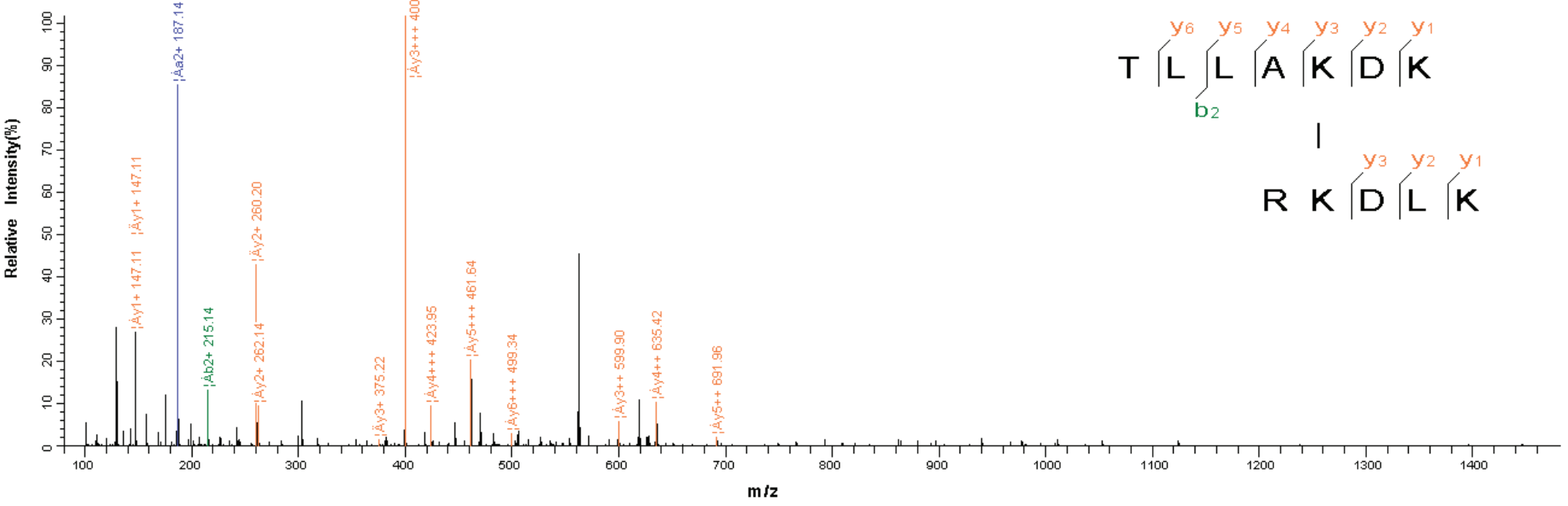
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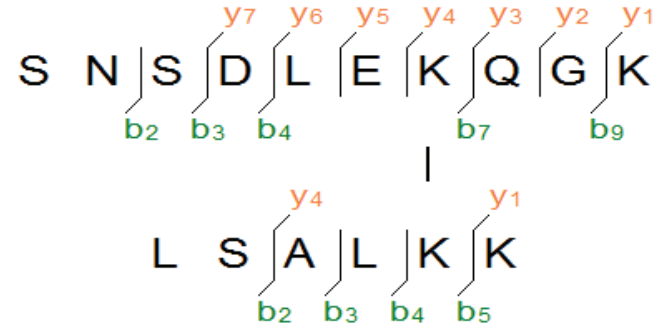
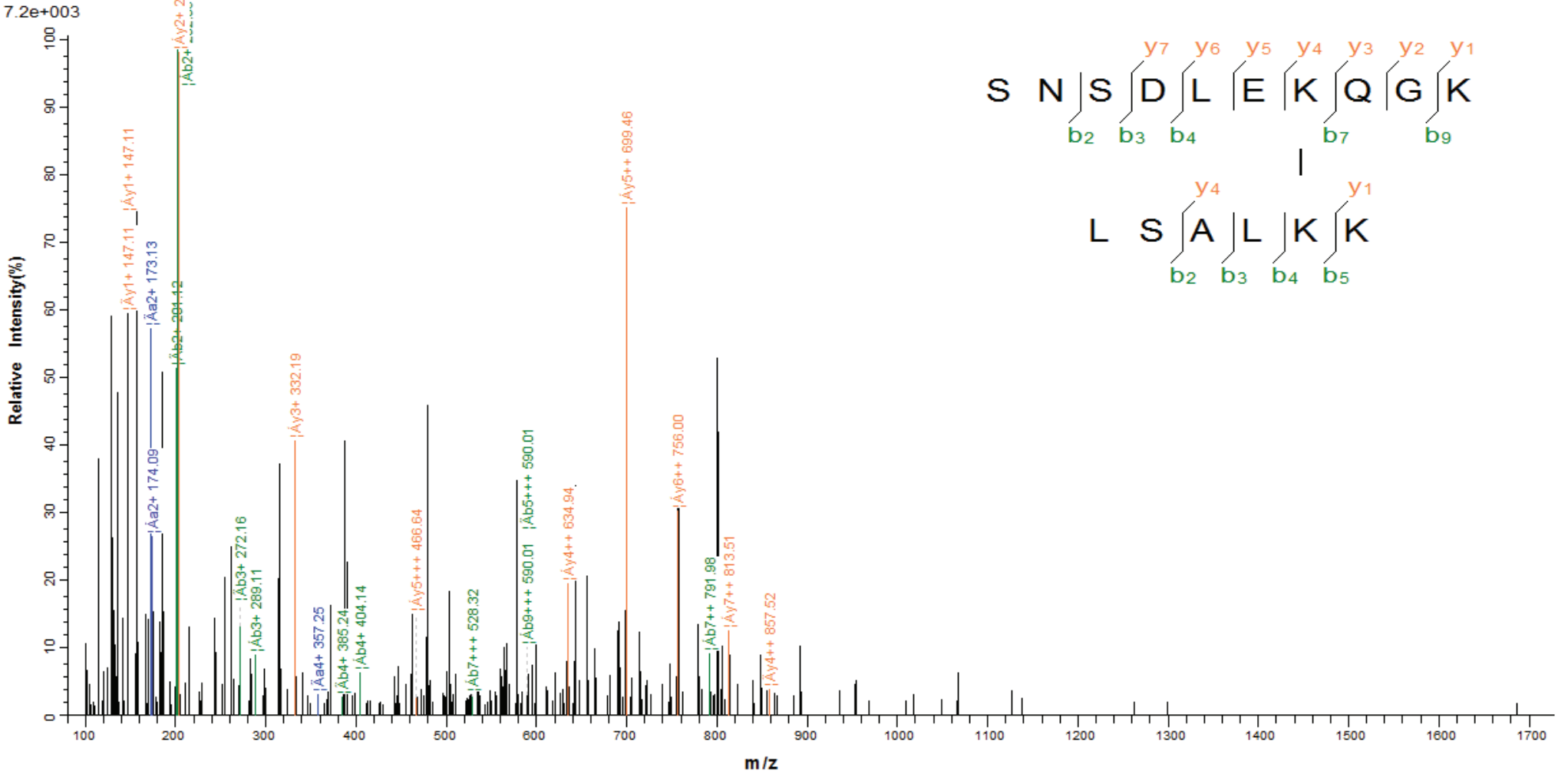
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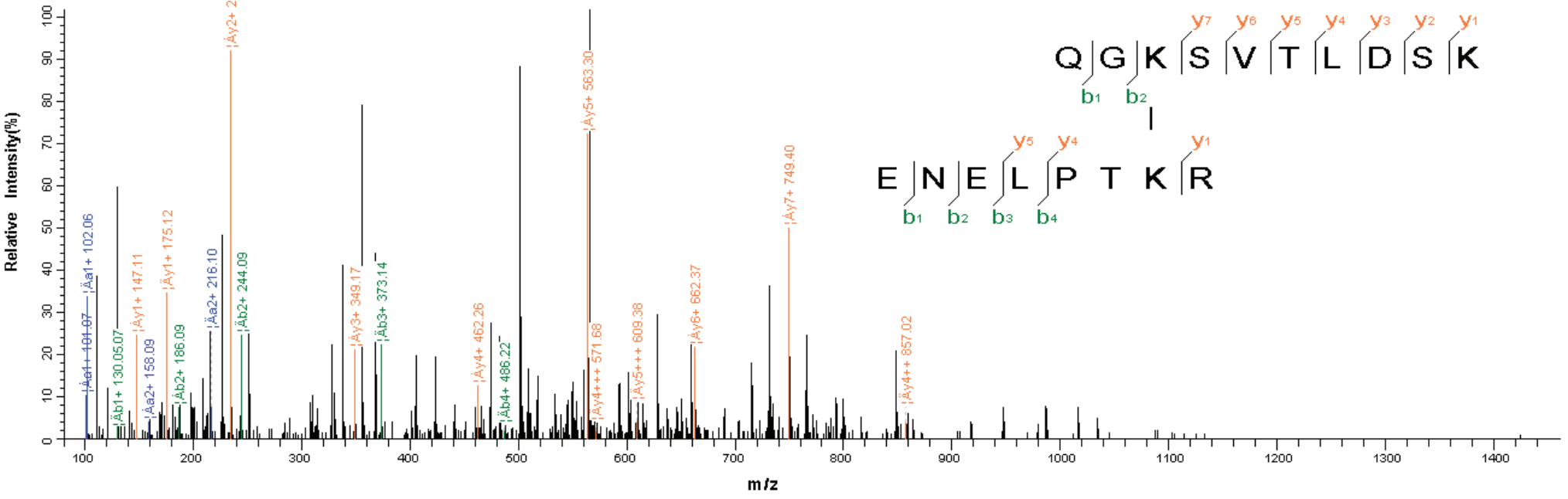
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Seq: SNSDLEKQK-LSALKK



Seq: QGKSVTLDSK-ENELPTRK



Seq: QGKSVTLDSK-KNSDLEK

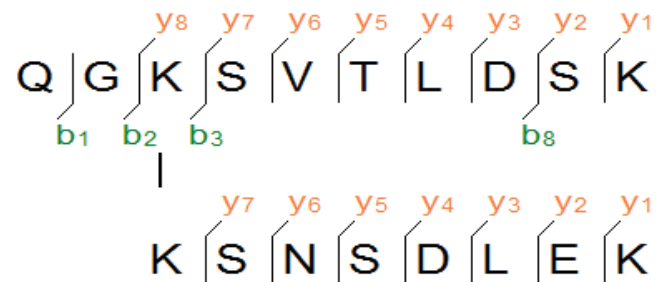
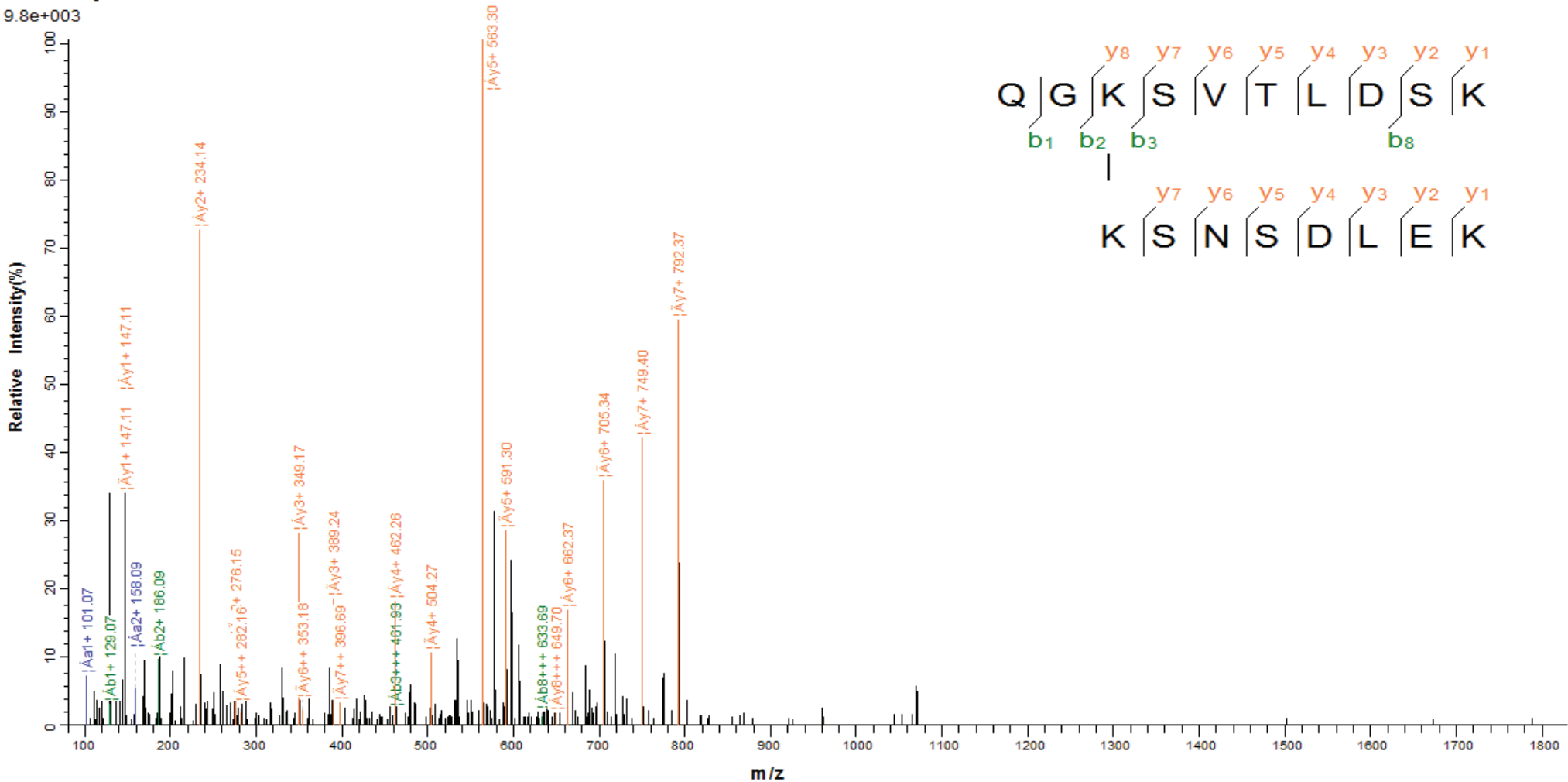


Fig. S18

Annotated high-resolution HCD spectra
of crosslink peptides of Beclin1 complex

