

protein: Y23H5A.5a
PEP: 0.00080664
Charge: 2
light
MODSEQ: _ALAPLTILAGR_

y1_1
b1_2

y2_8
b1_3

y1_3

y1_4

y1_5
b2_11

b1_7

b1_8

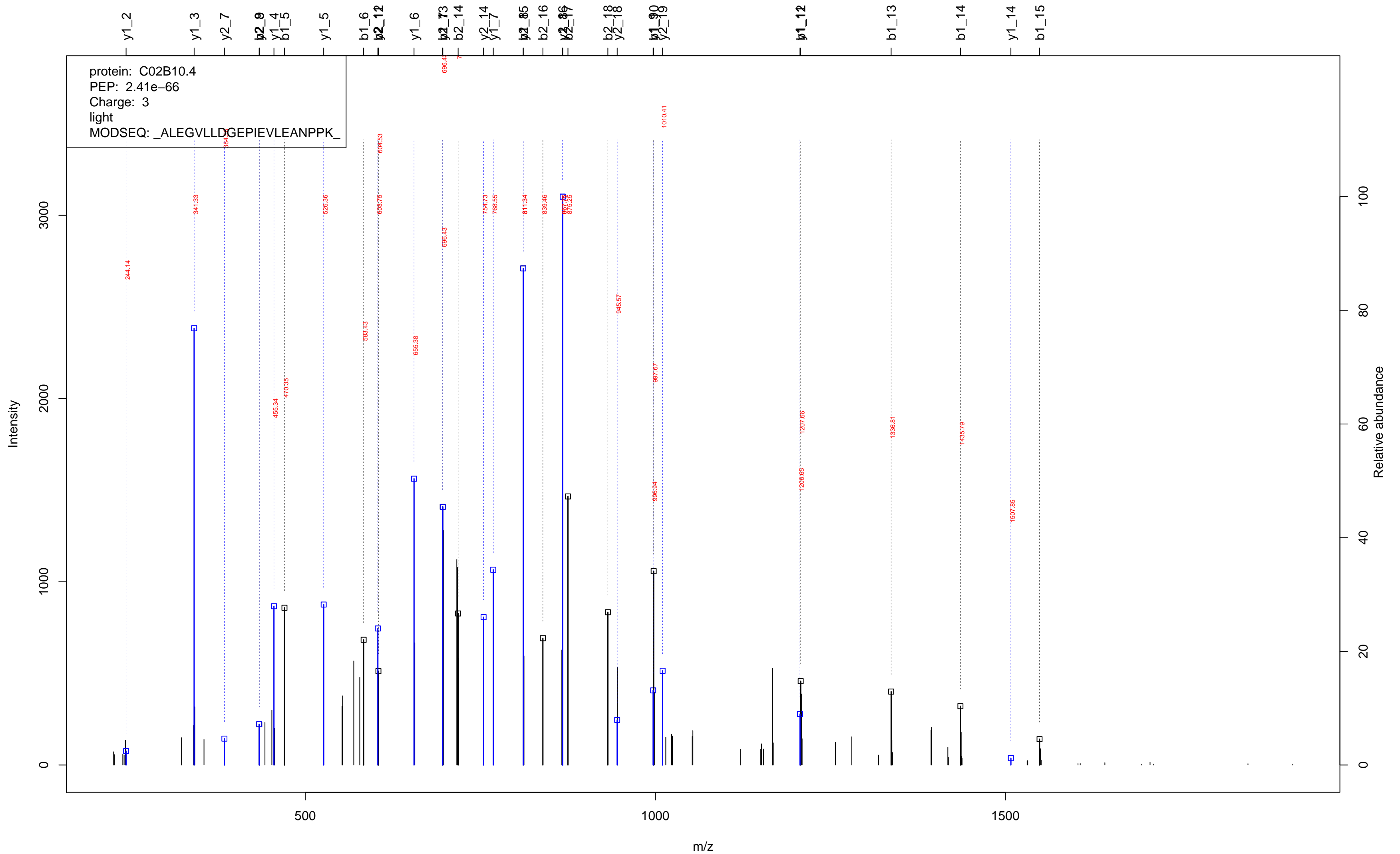
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b1_9

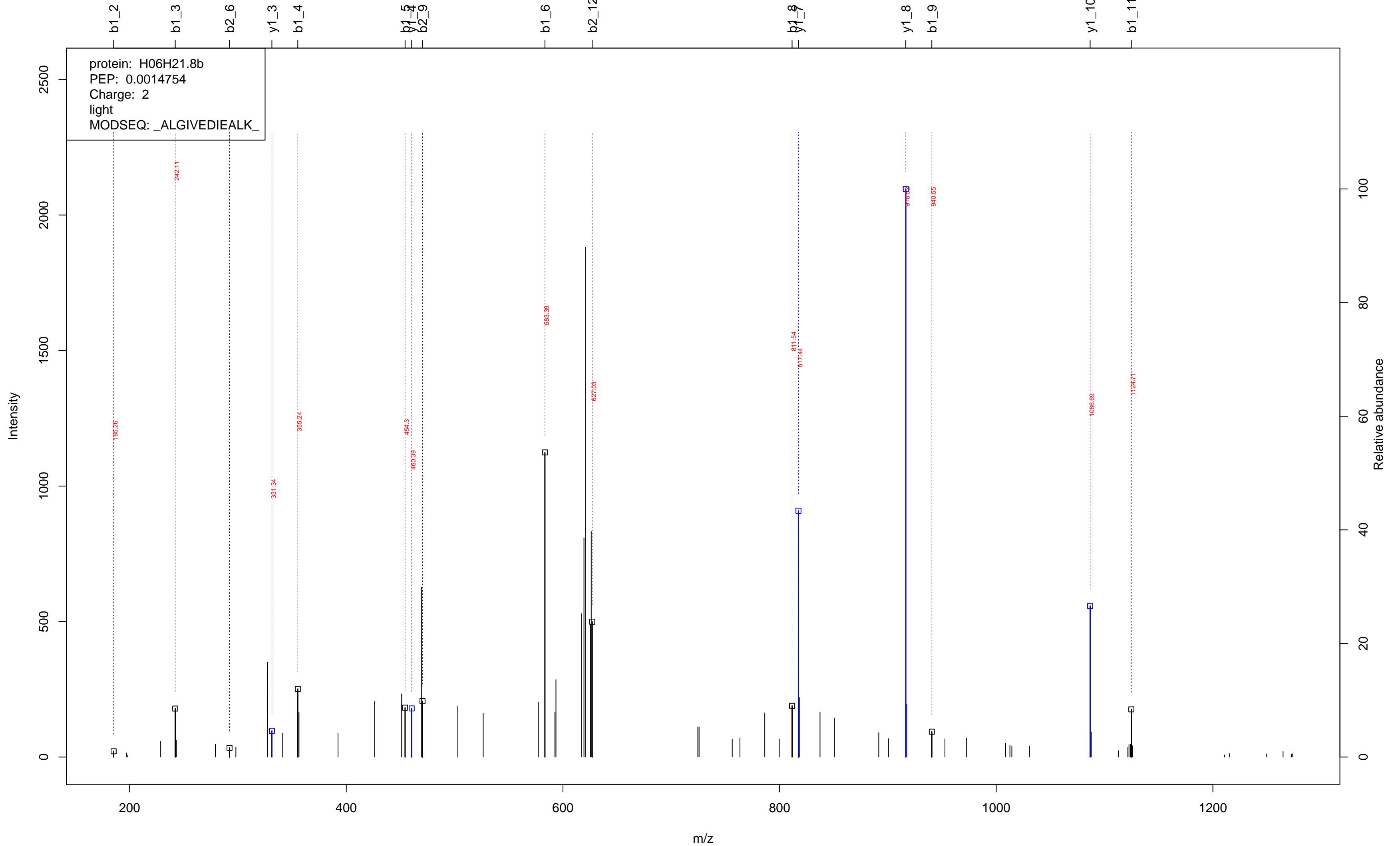
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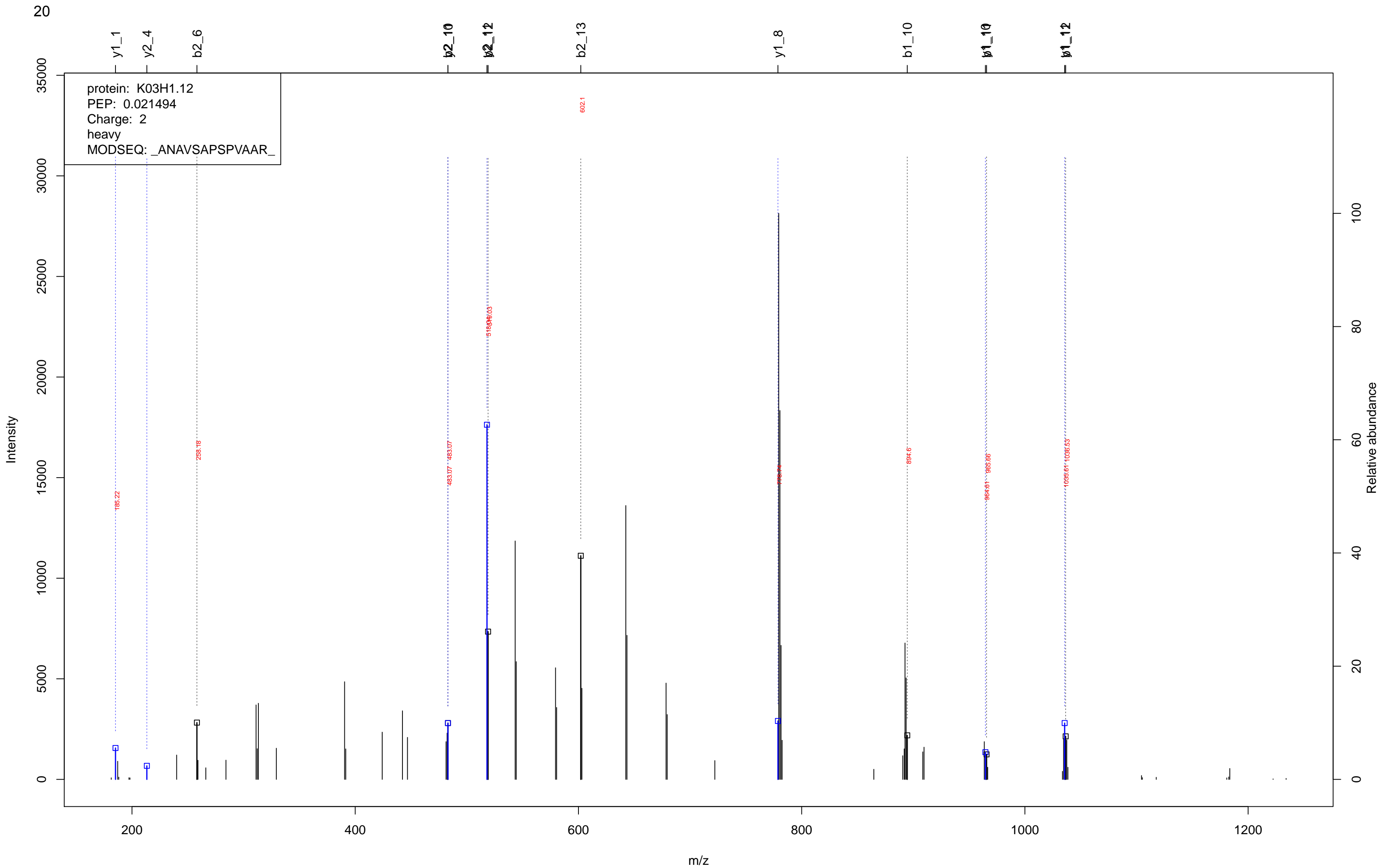
Intensity

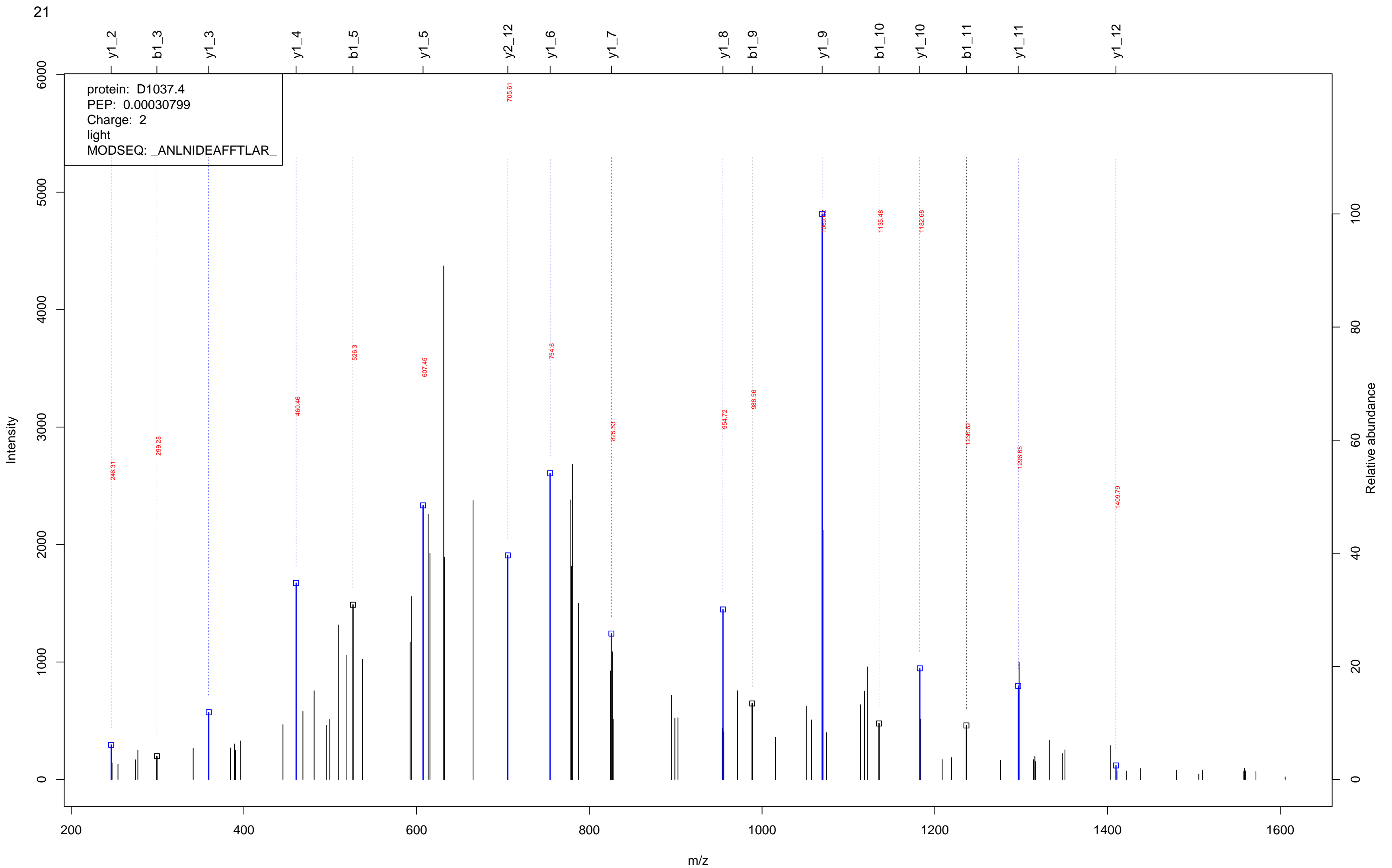
Relative abundance

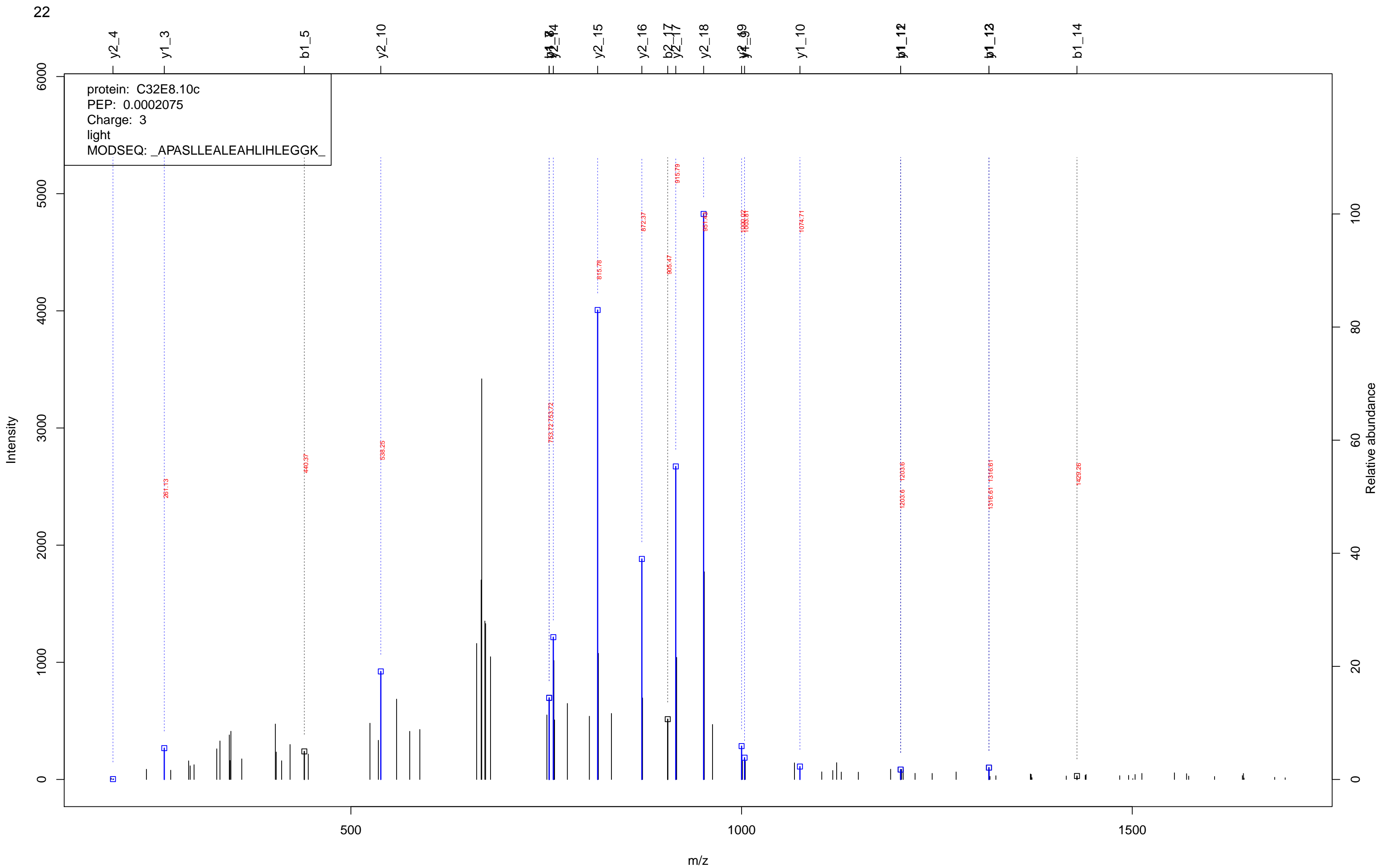
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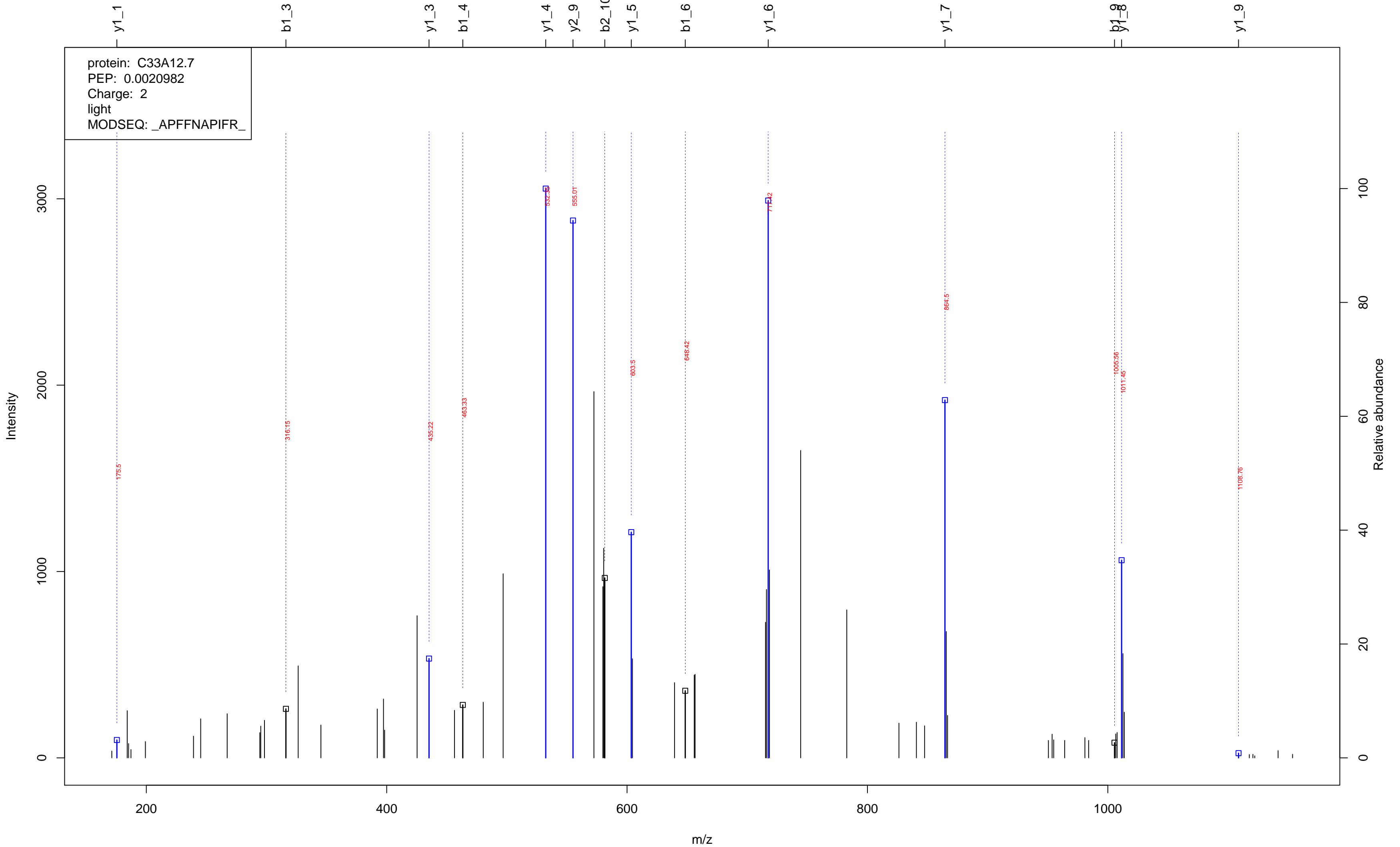


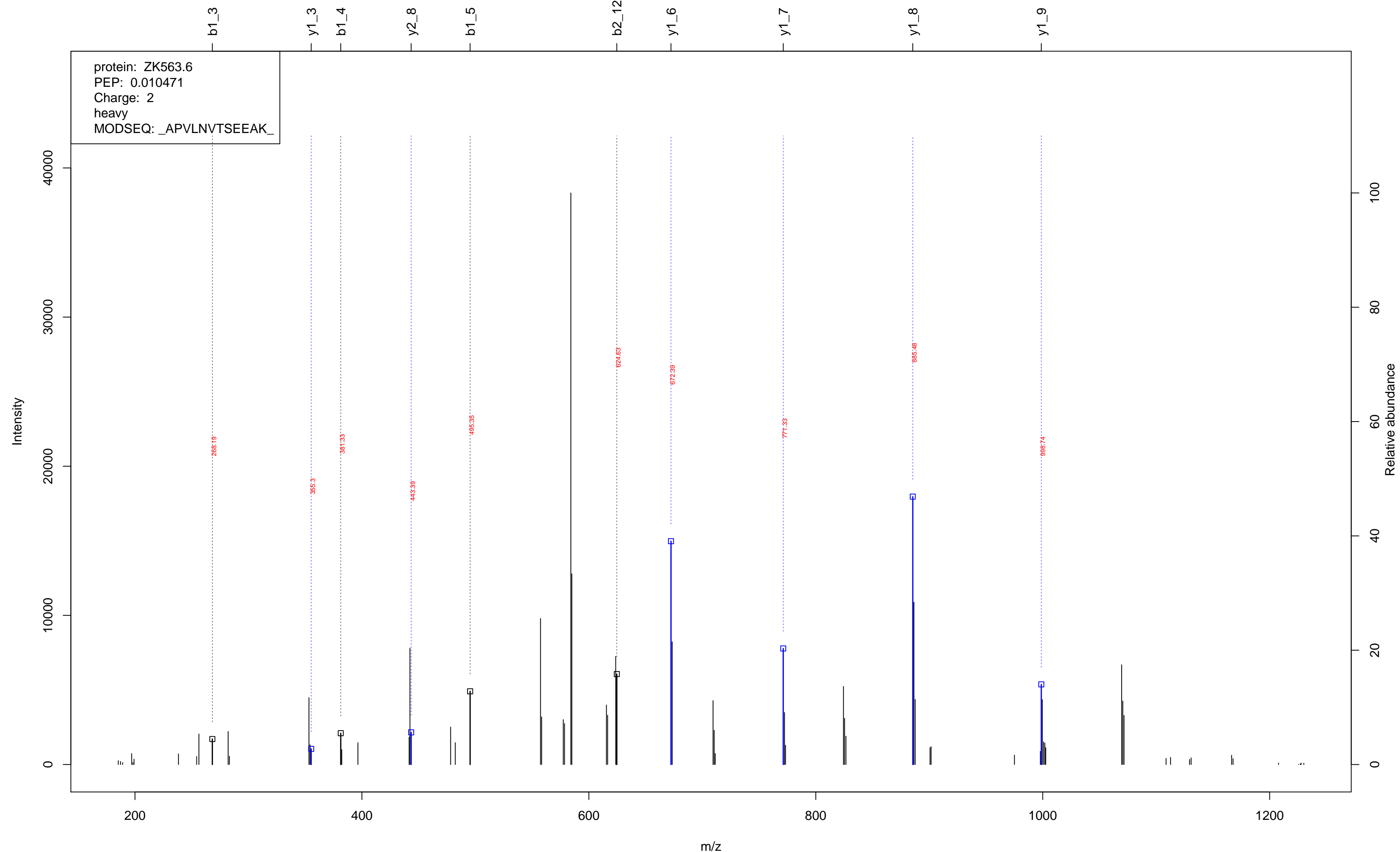


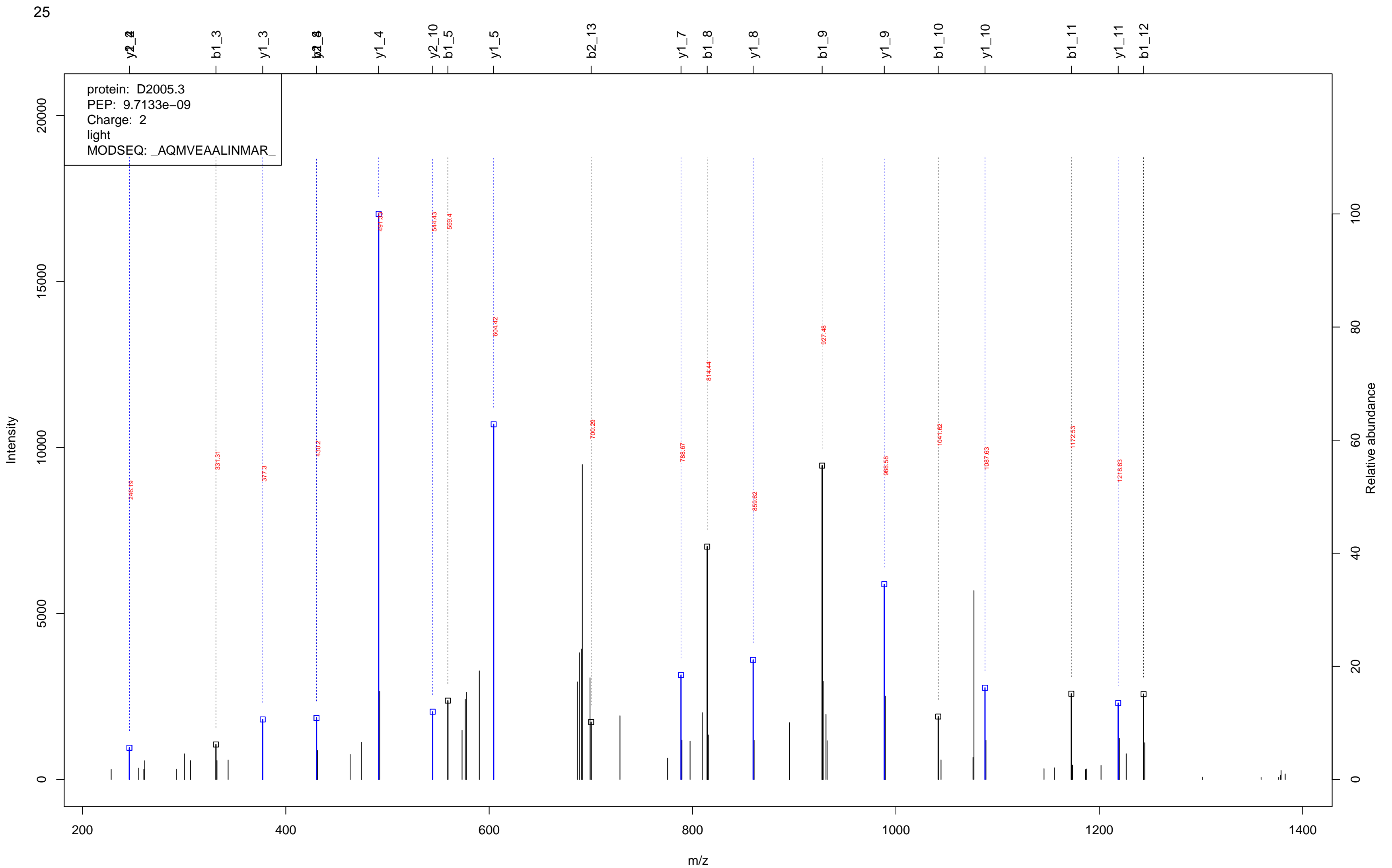


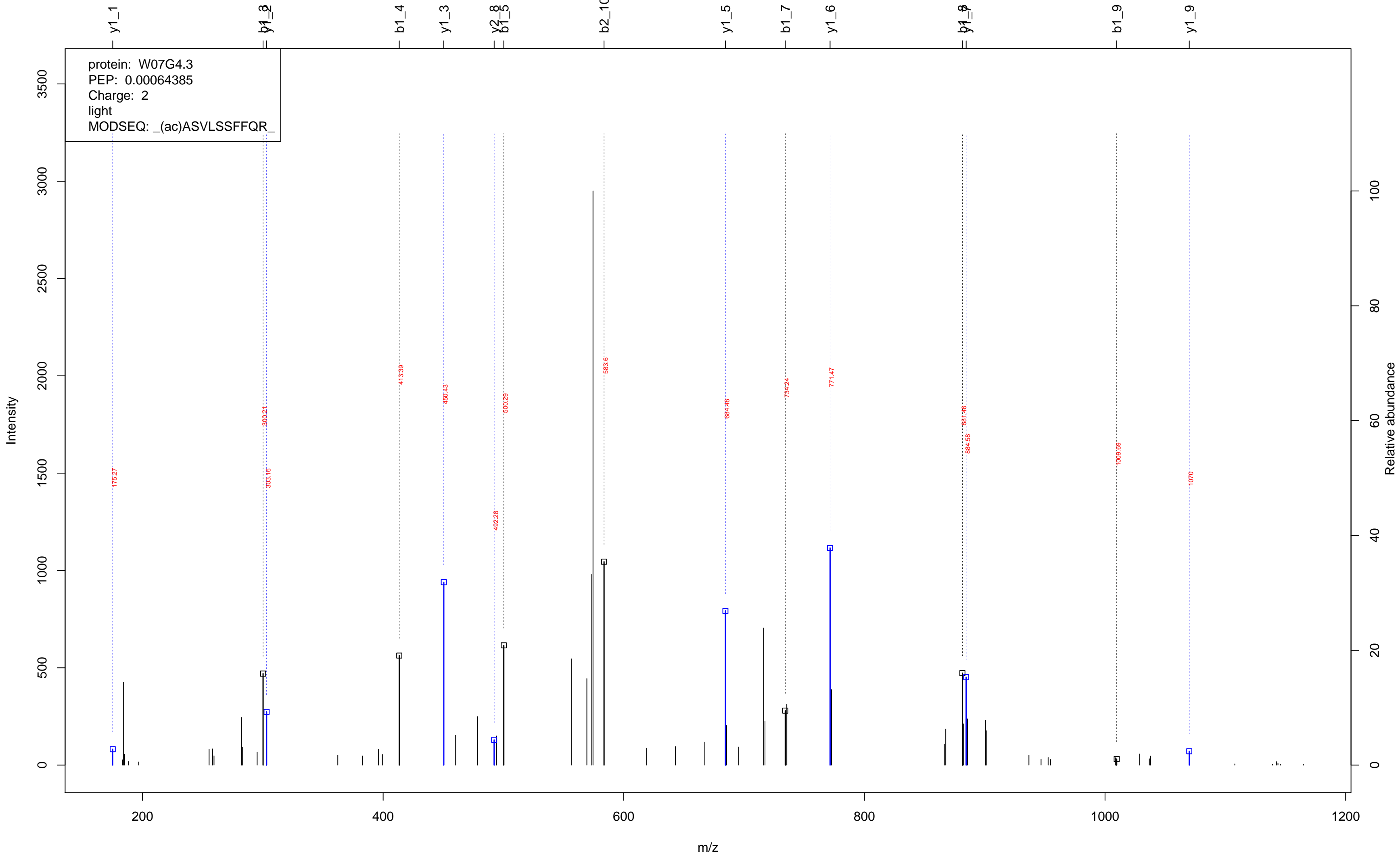


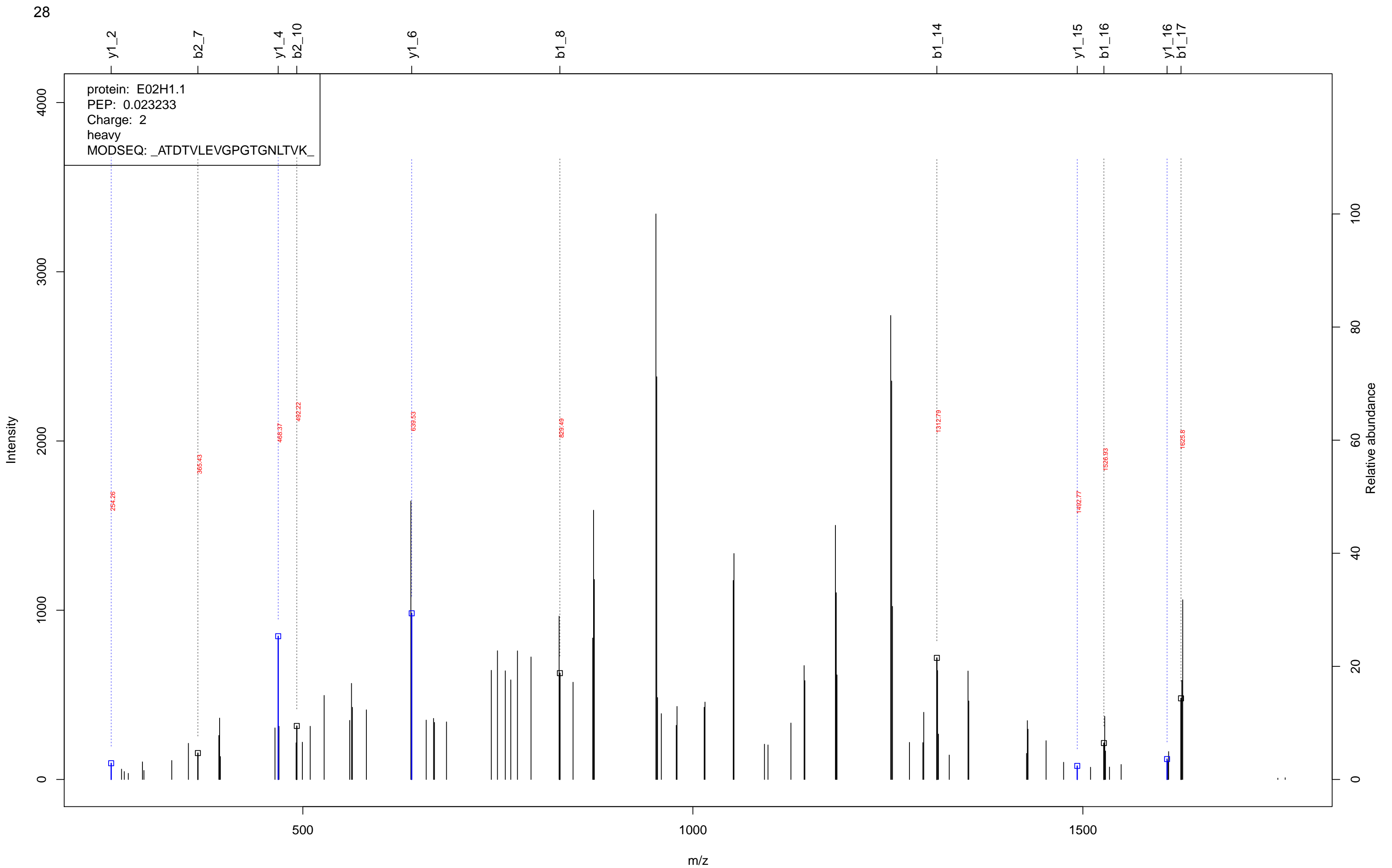


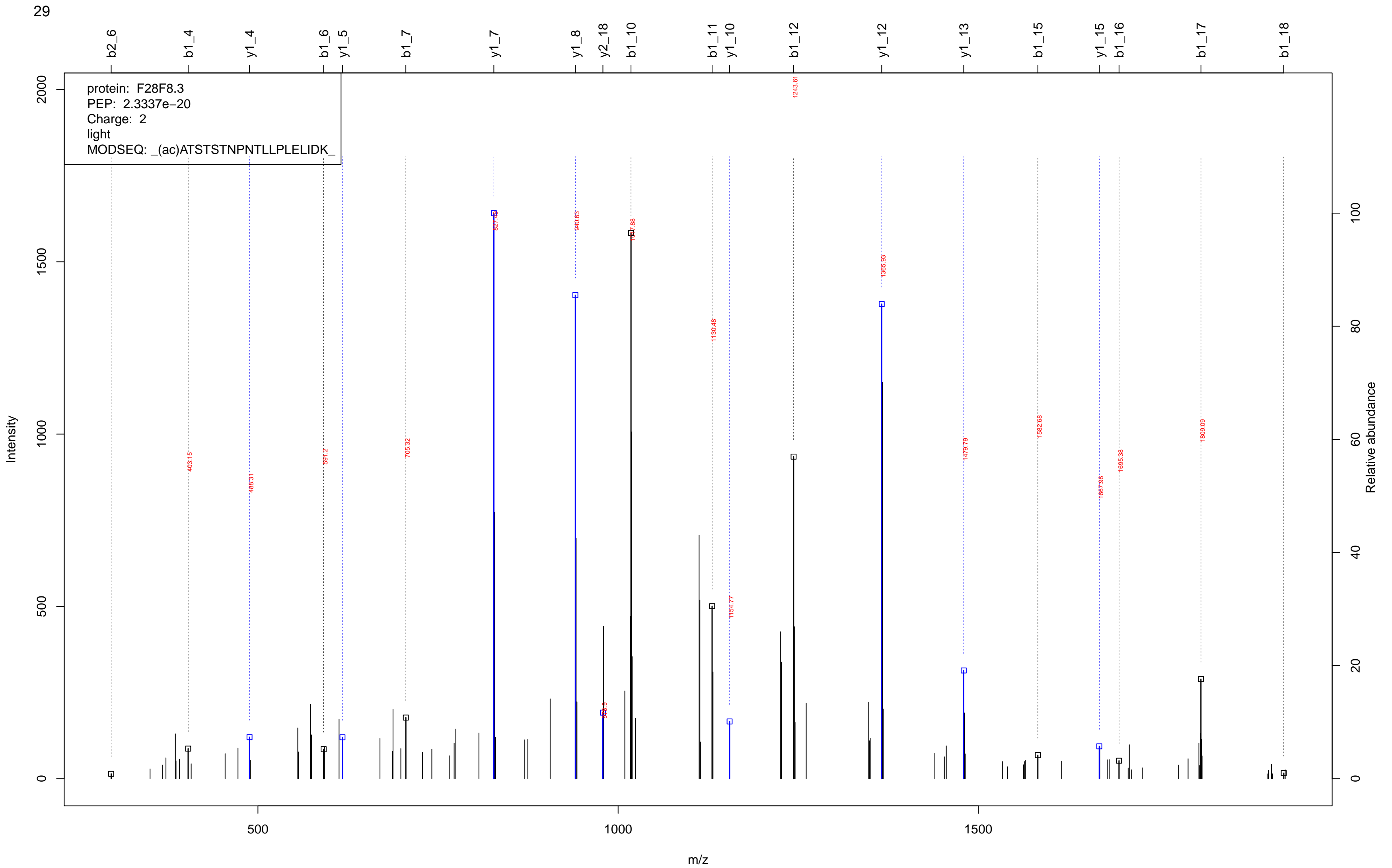


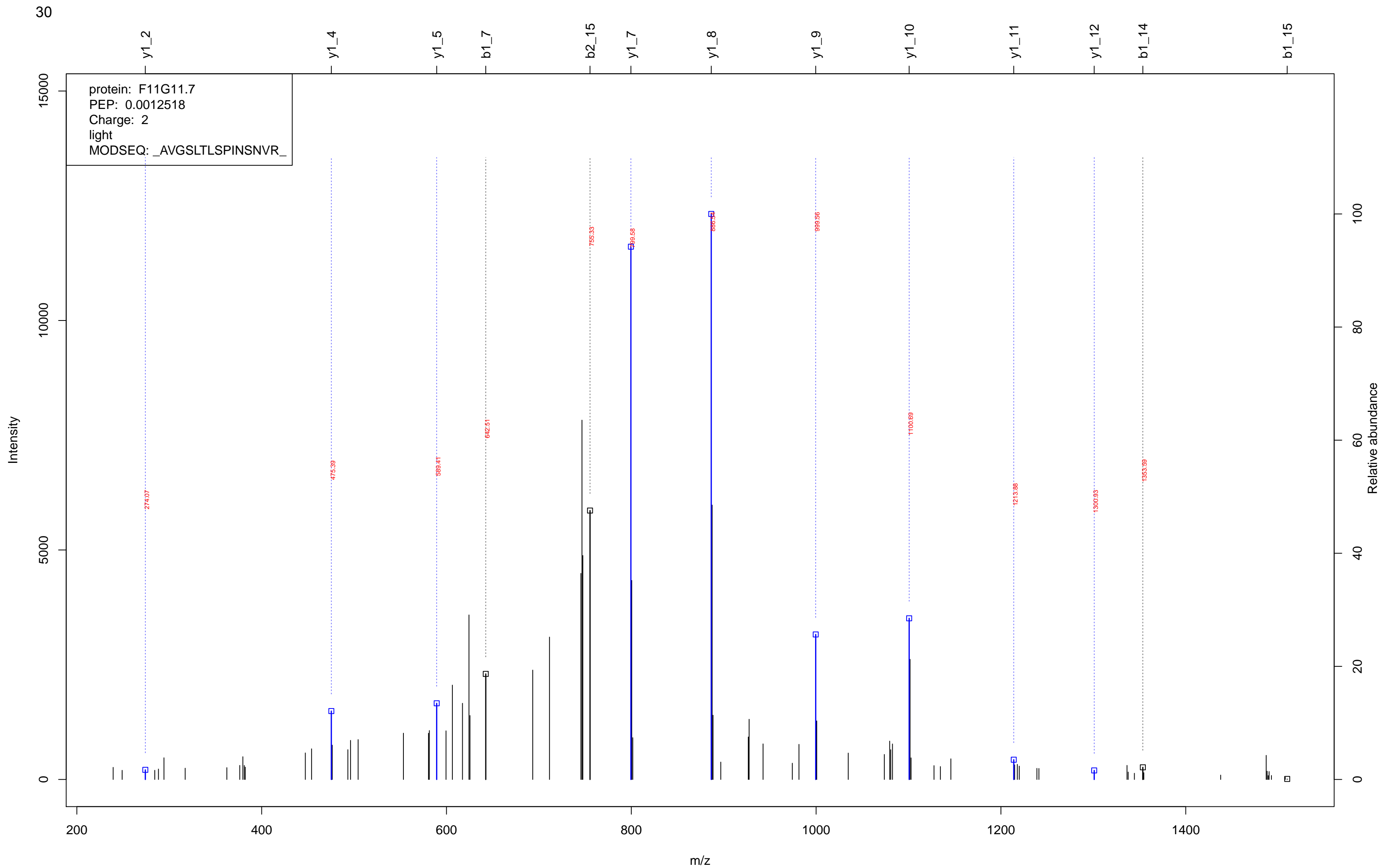


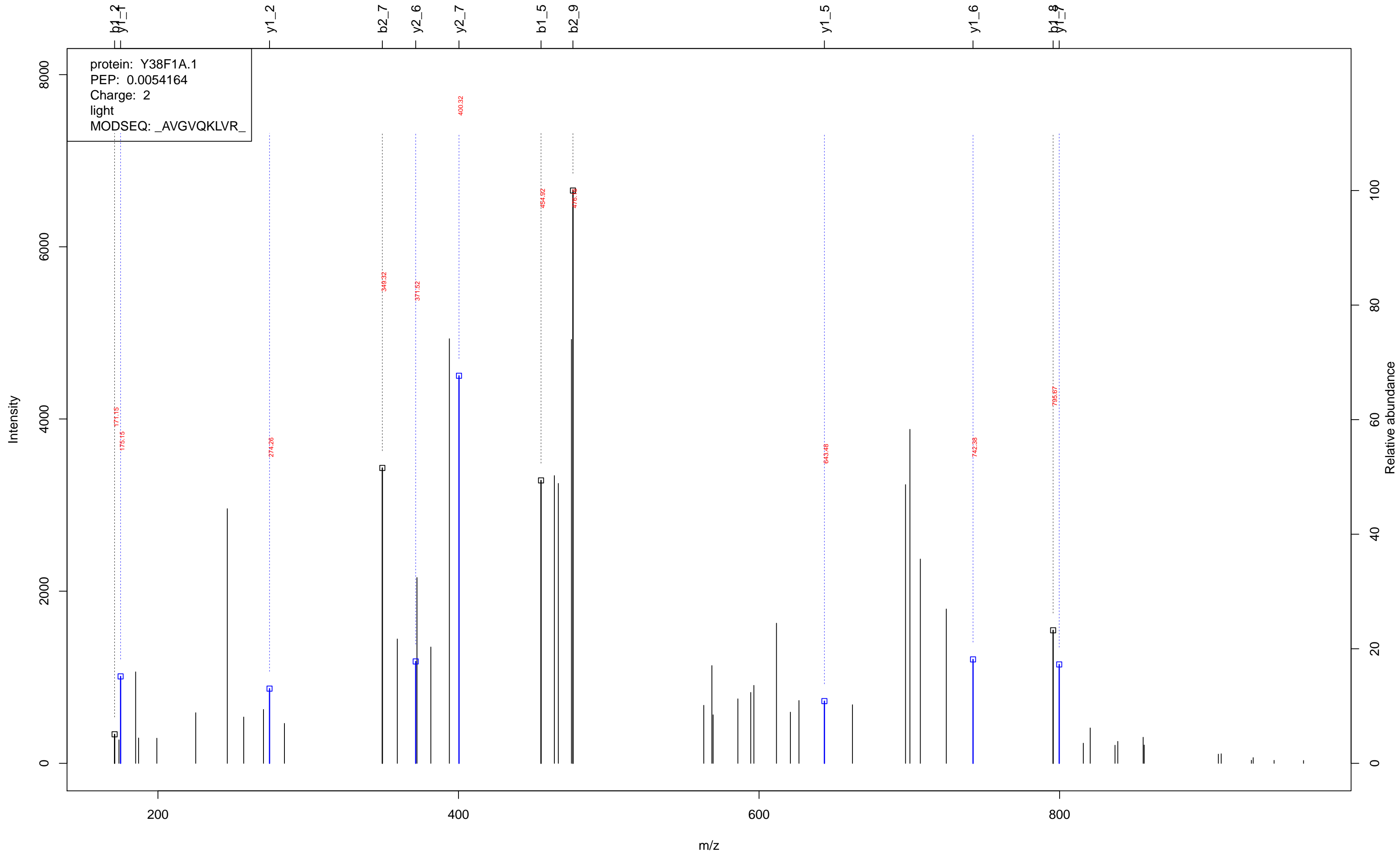


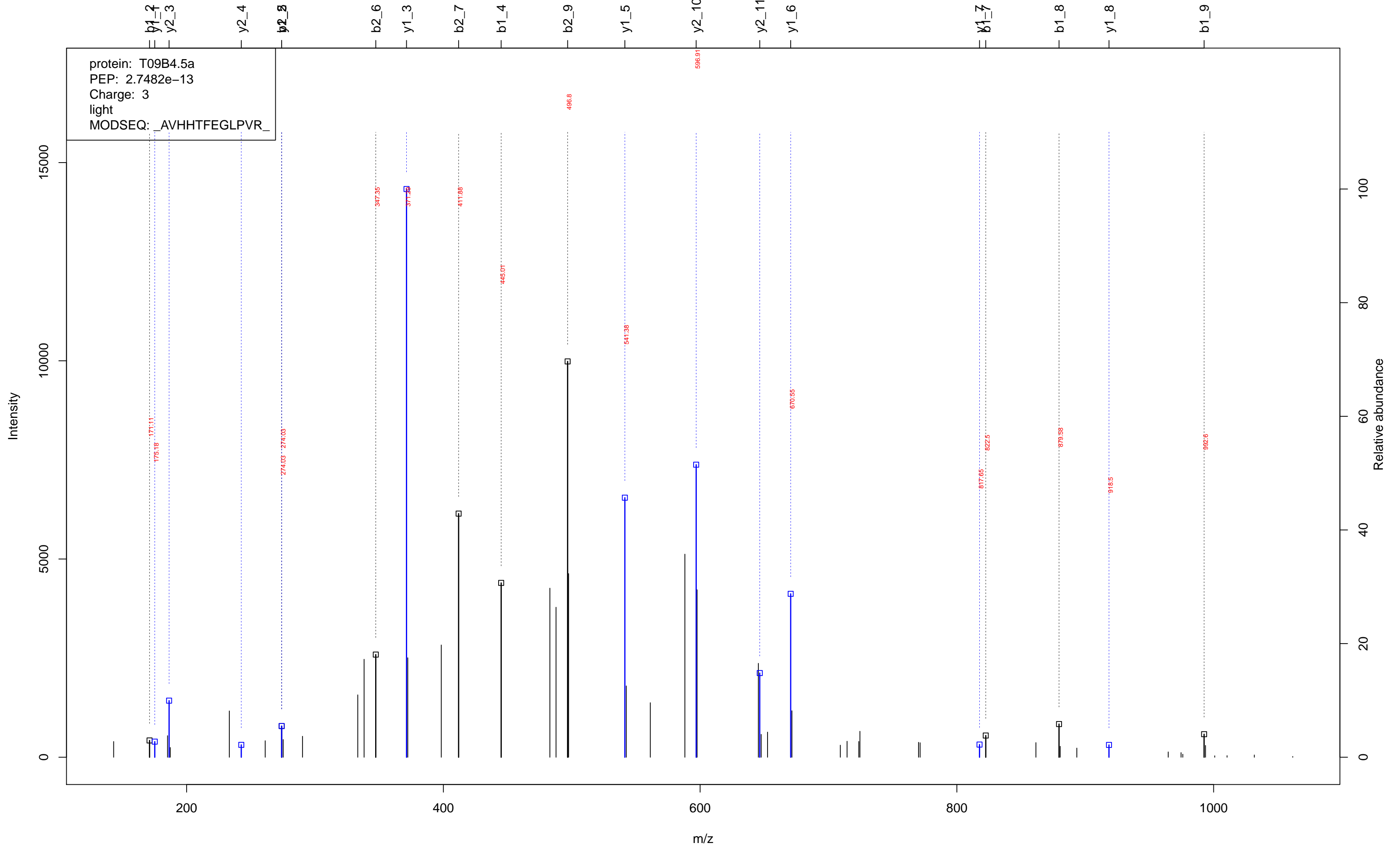


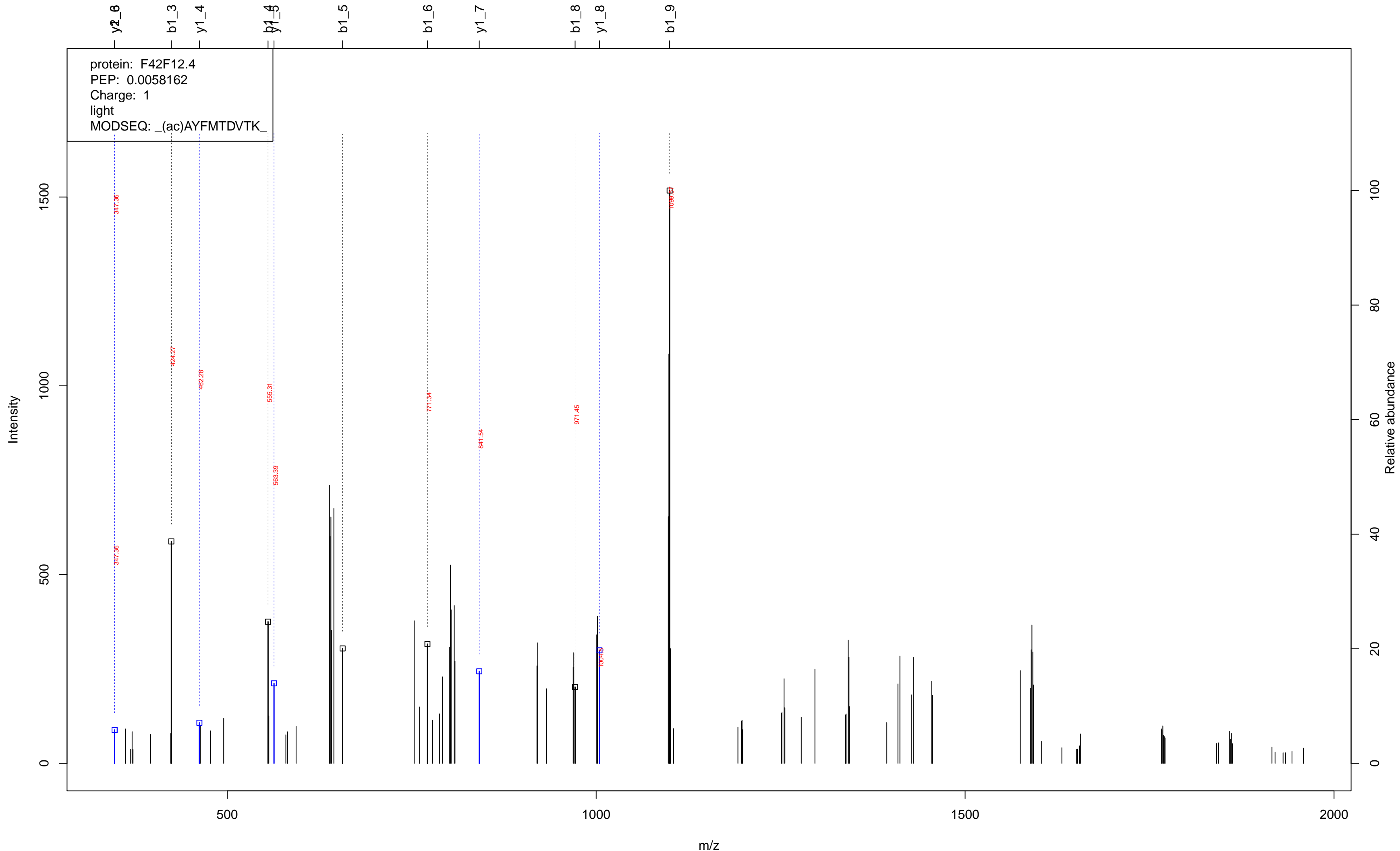


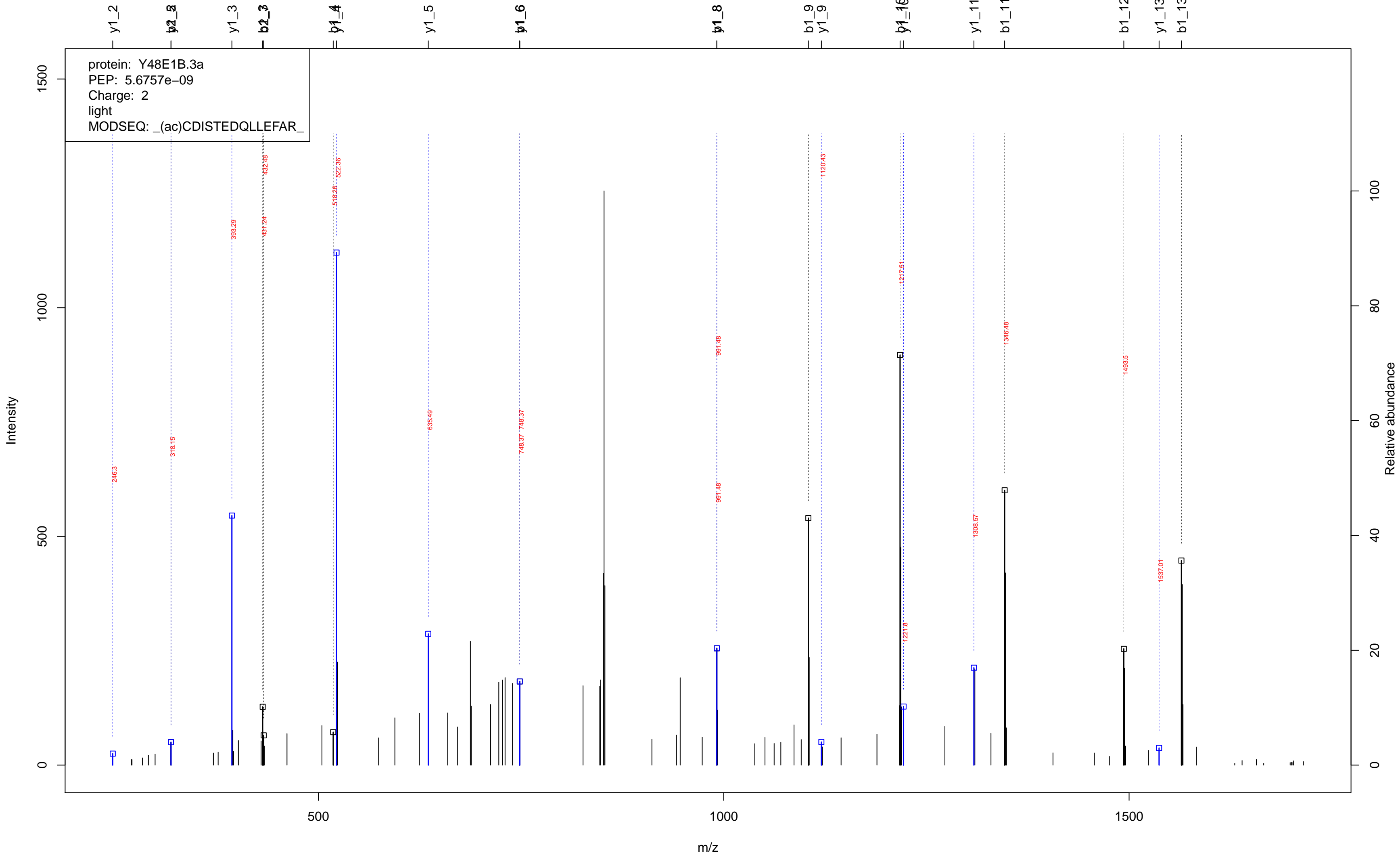


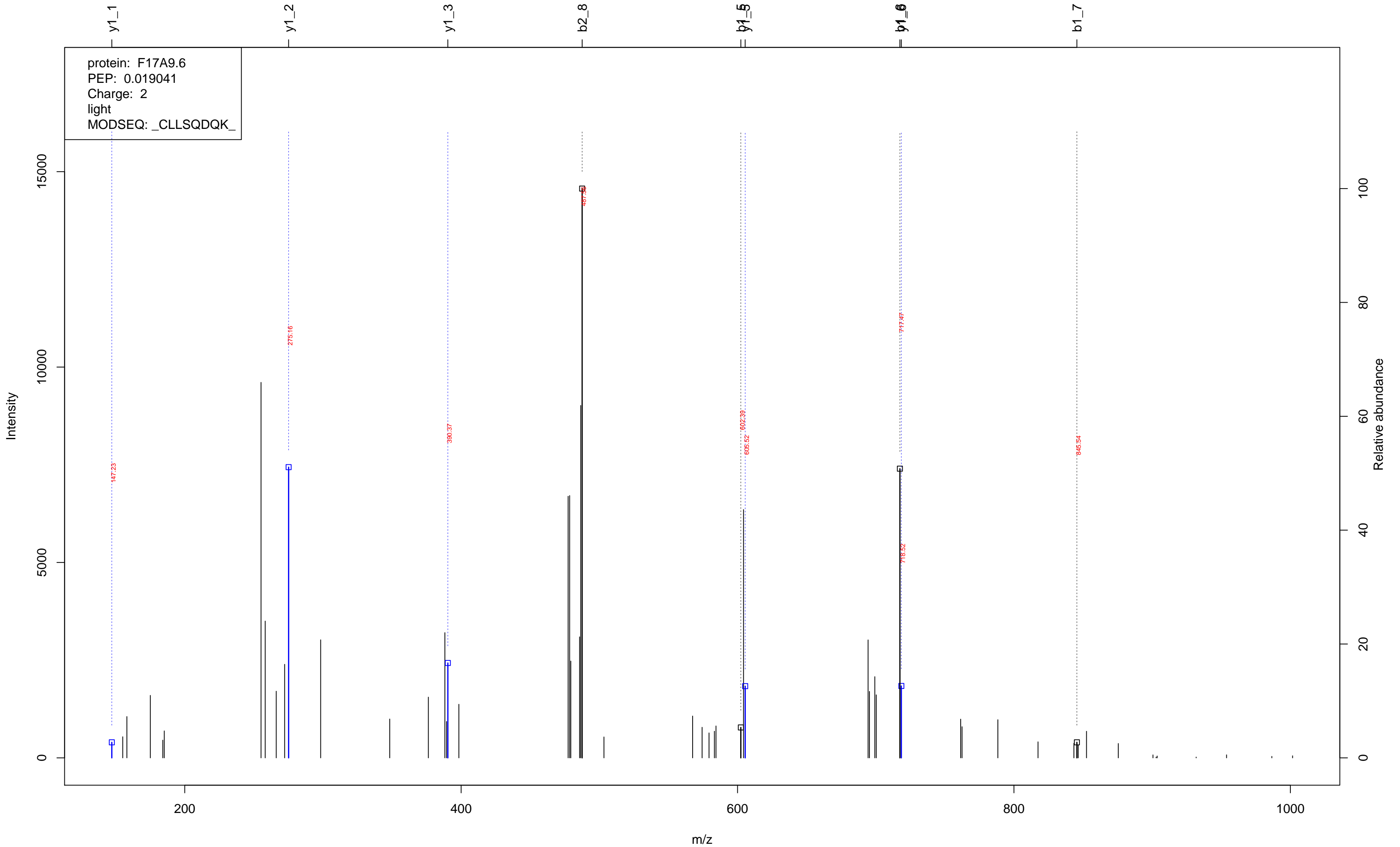


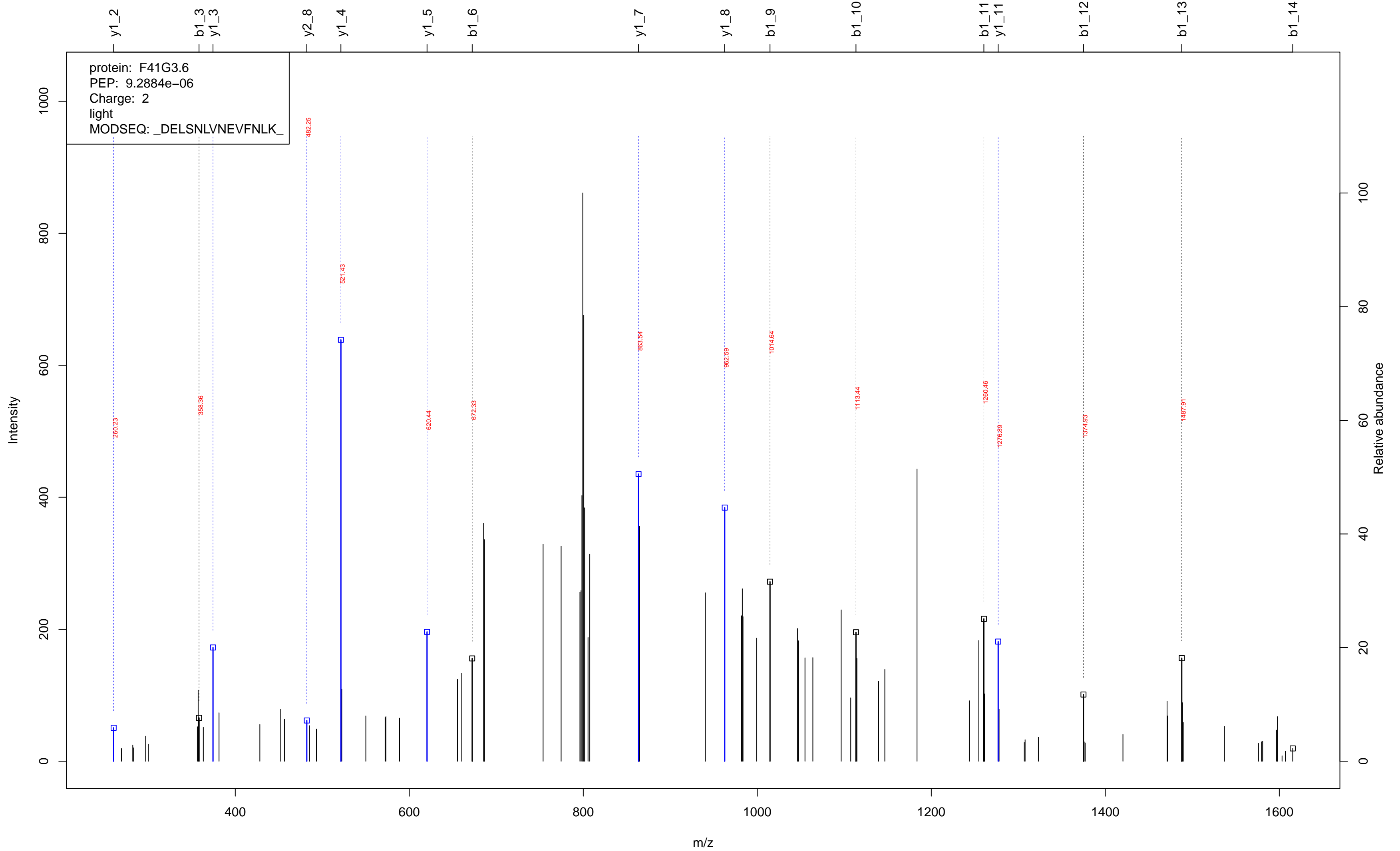


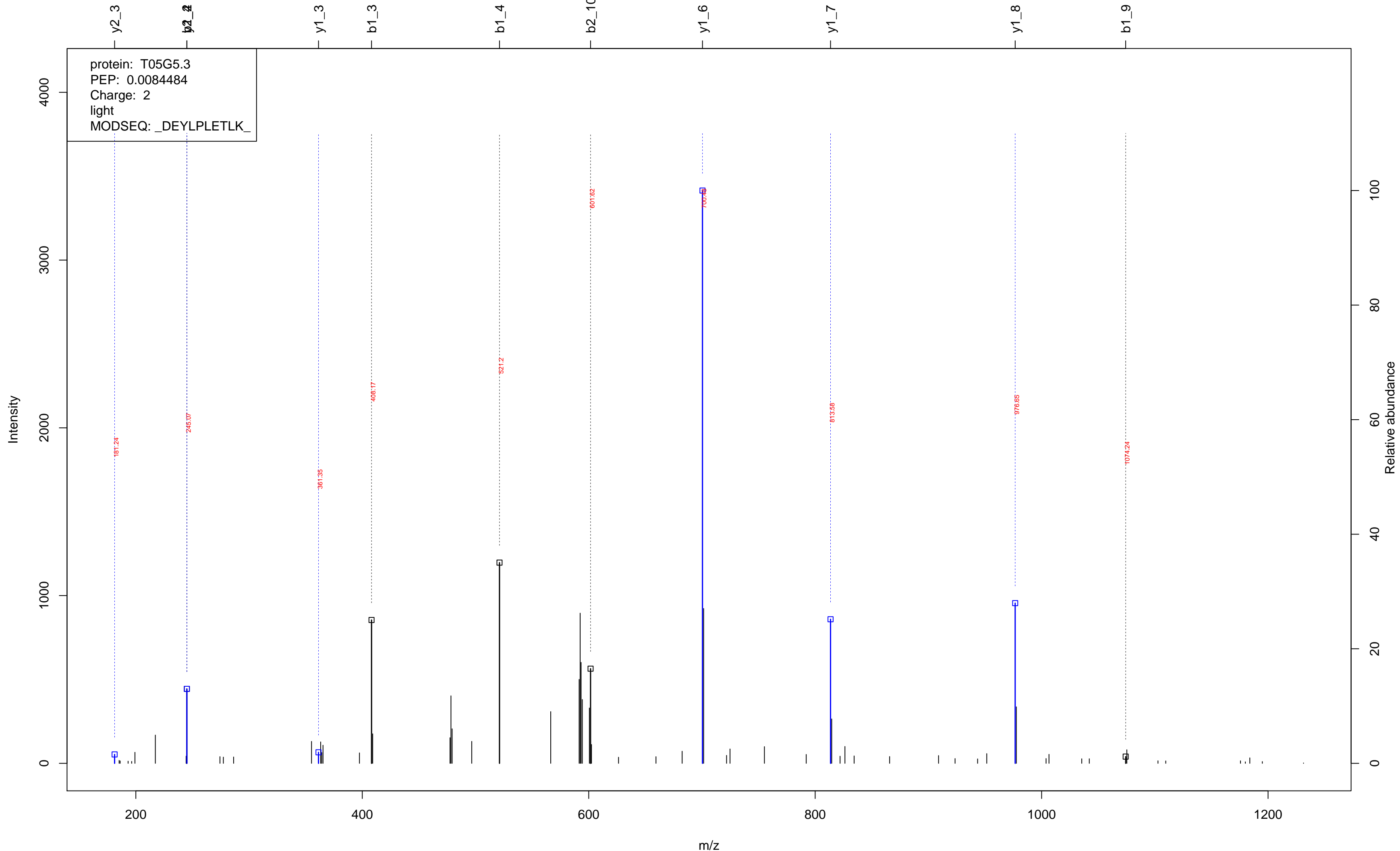


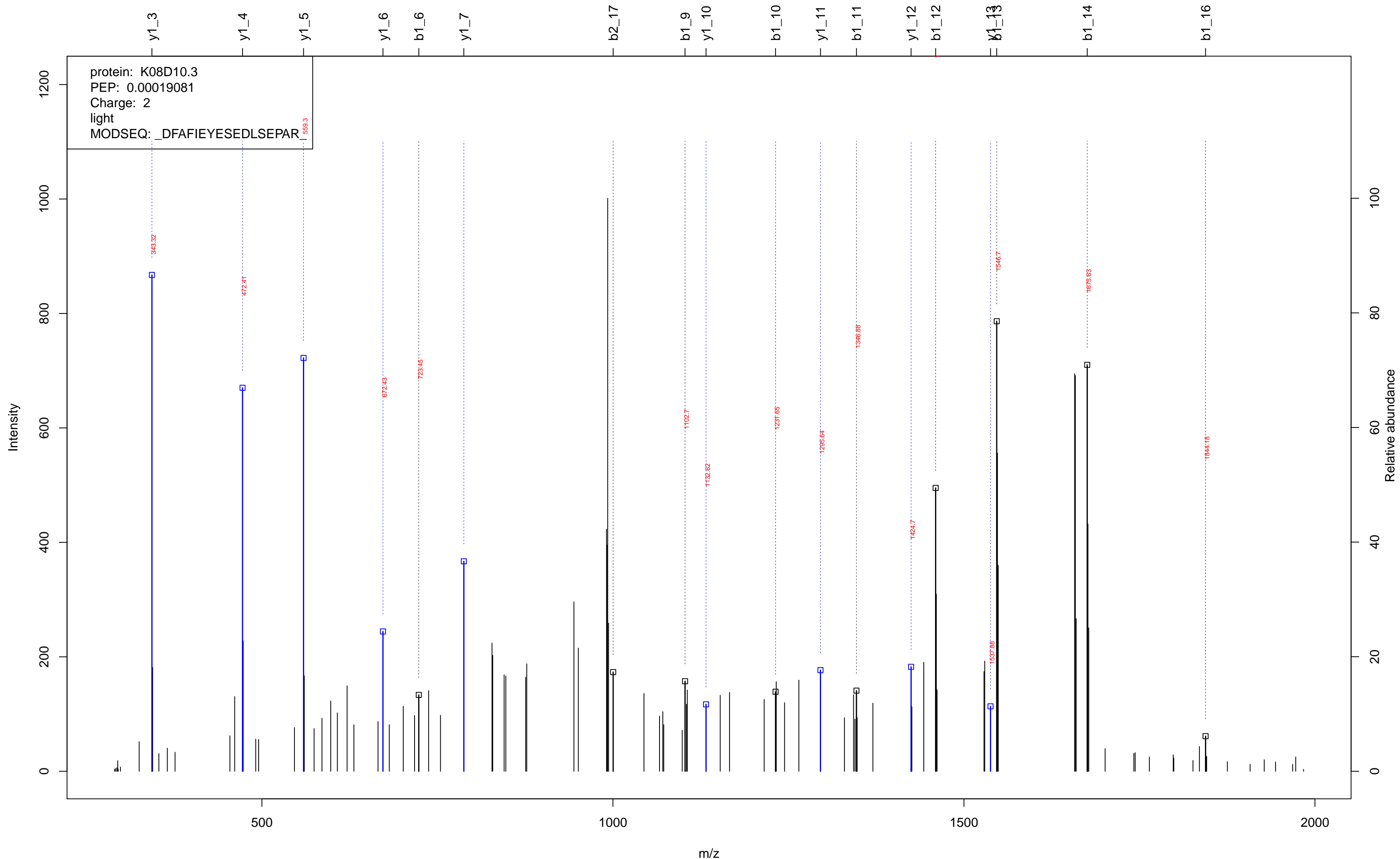


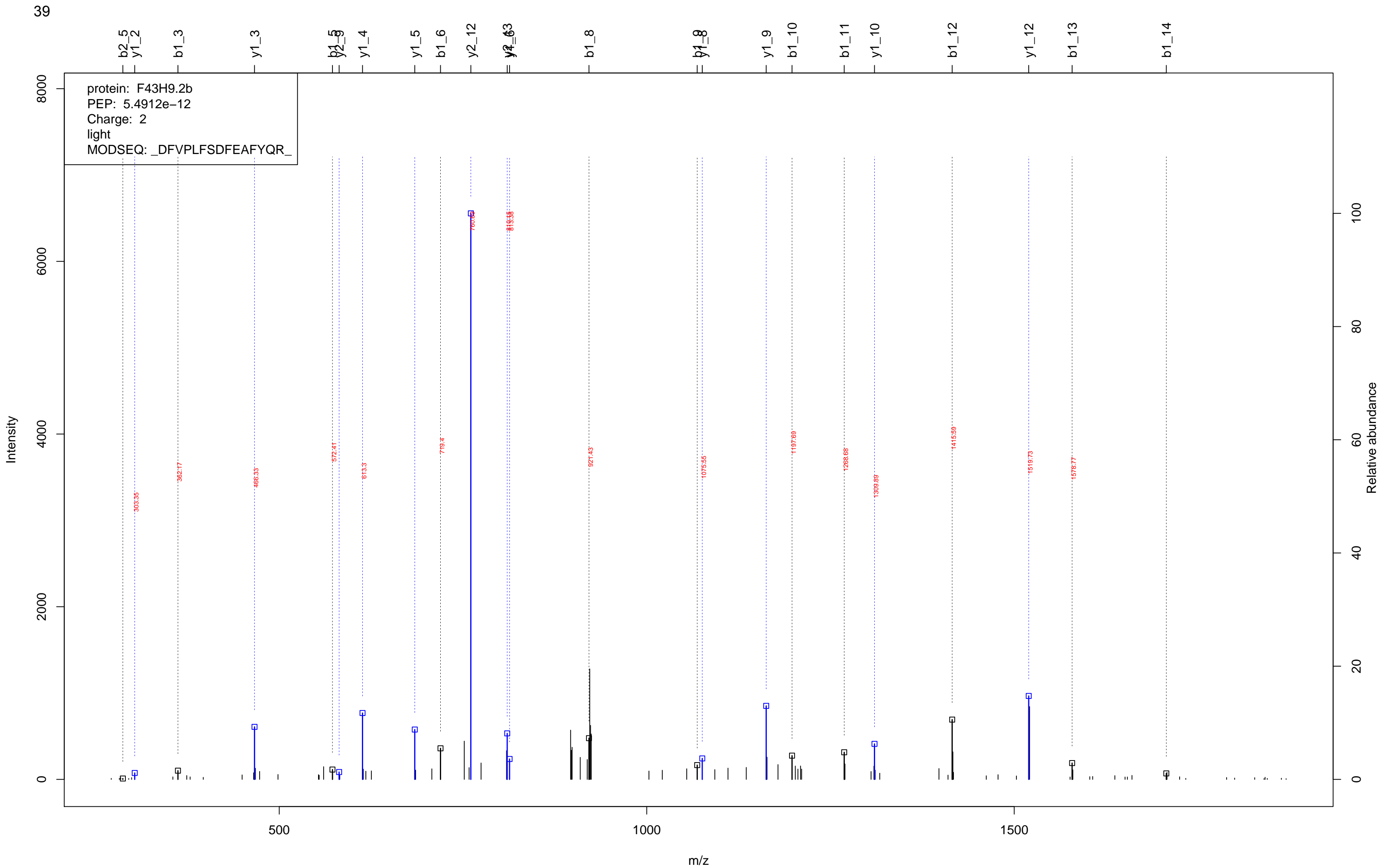


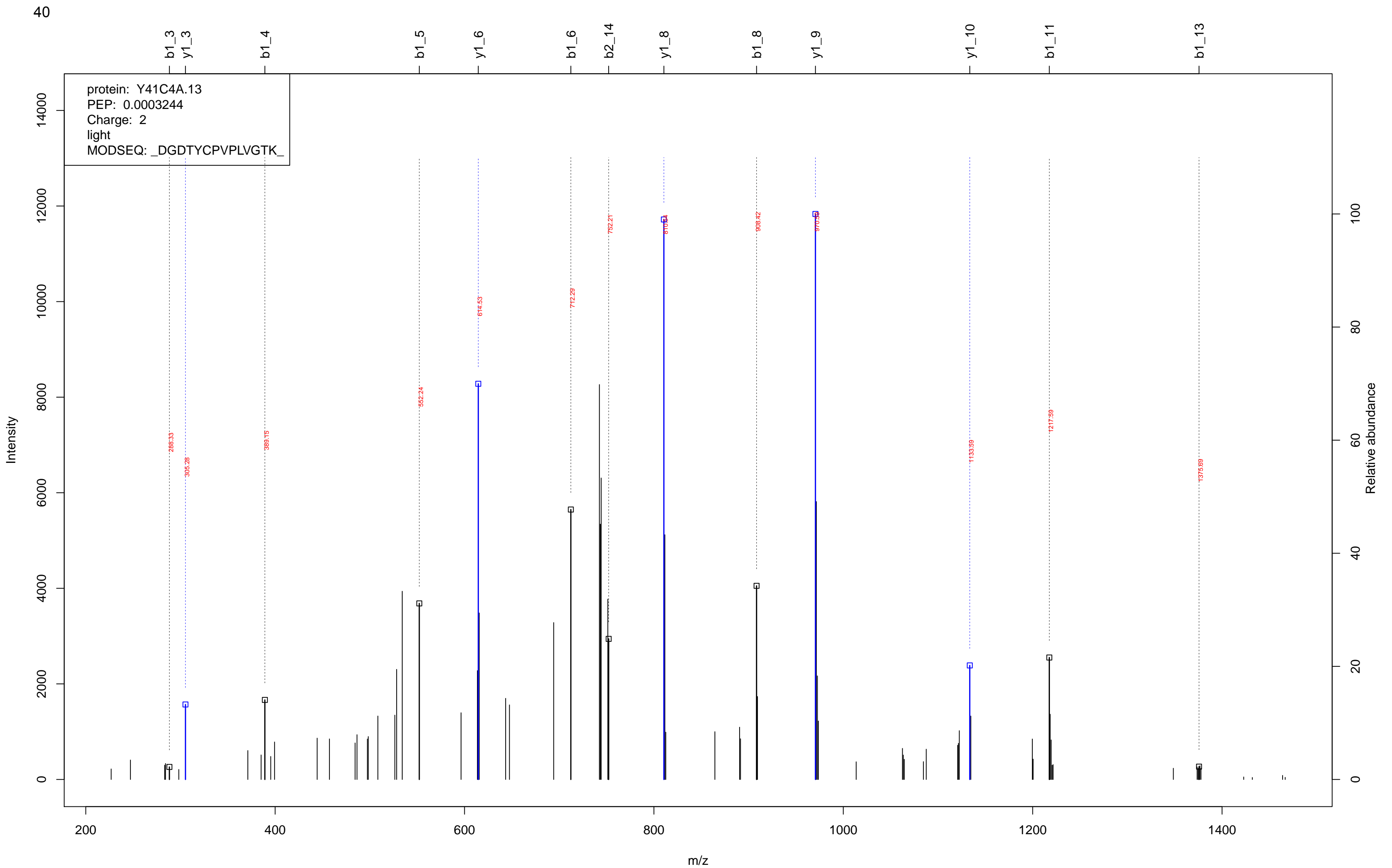


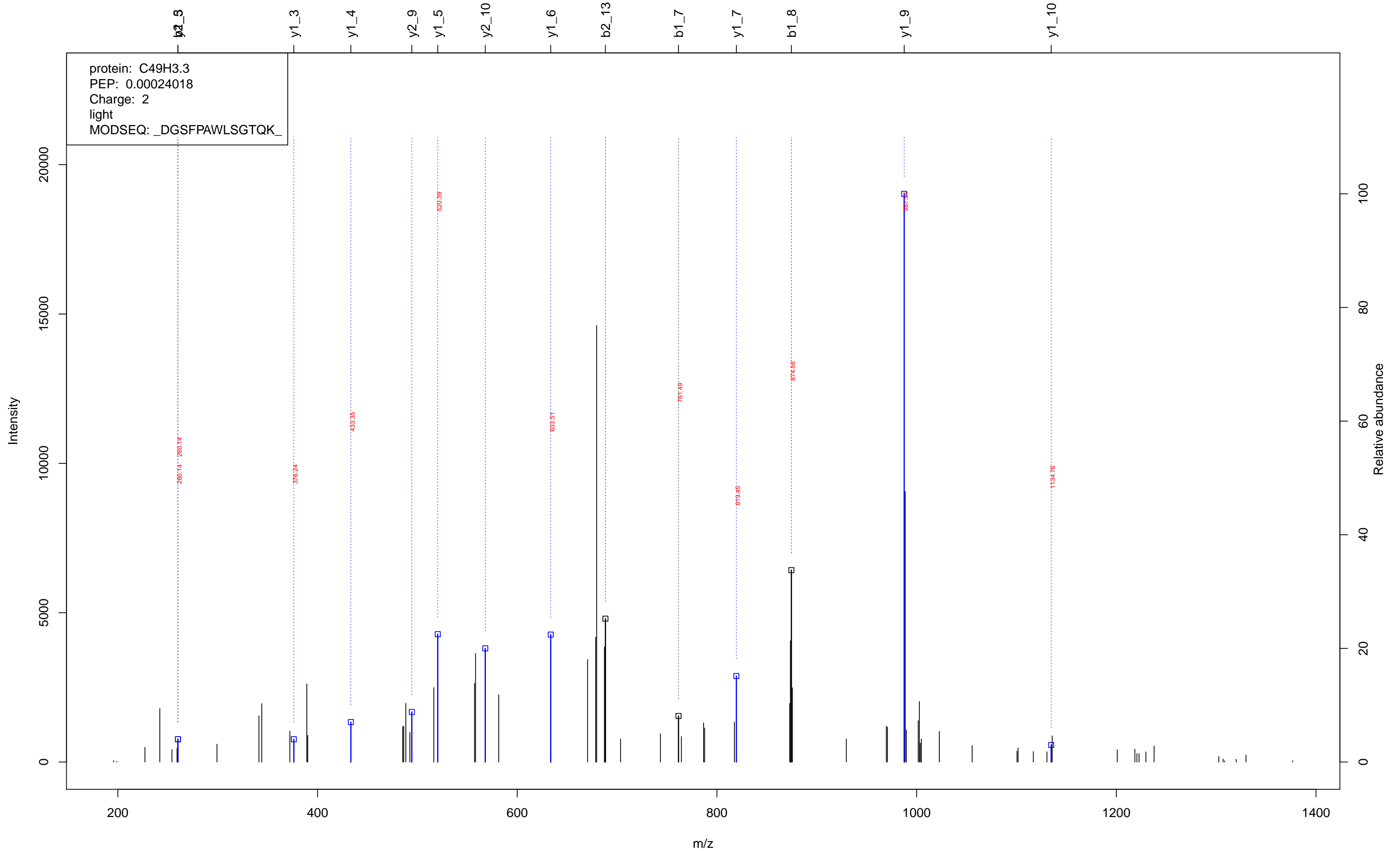


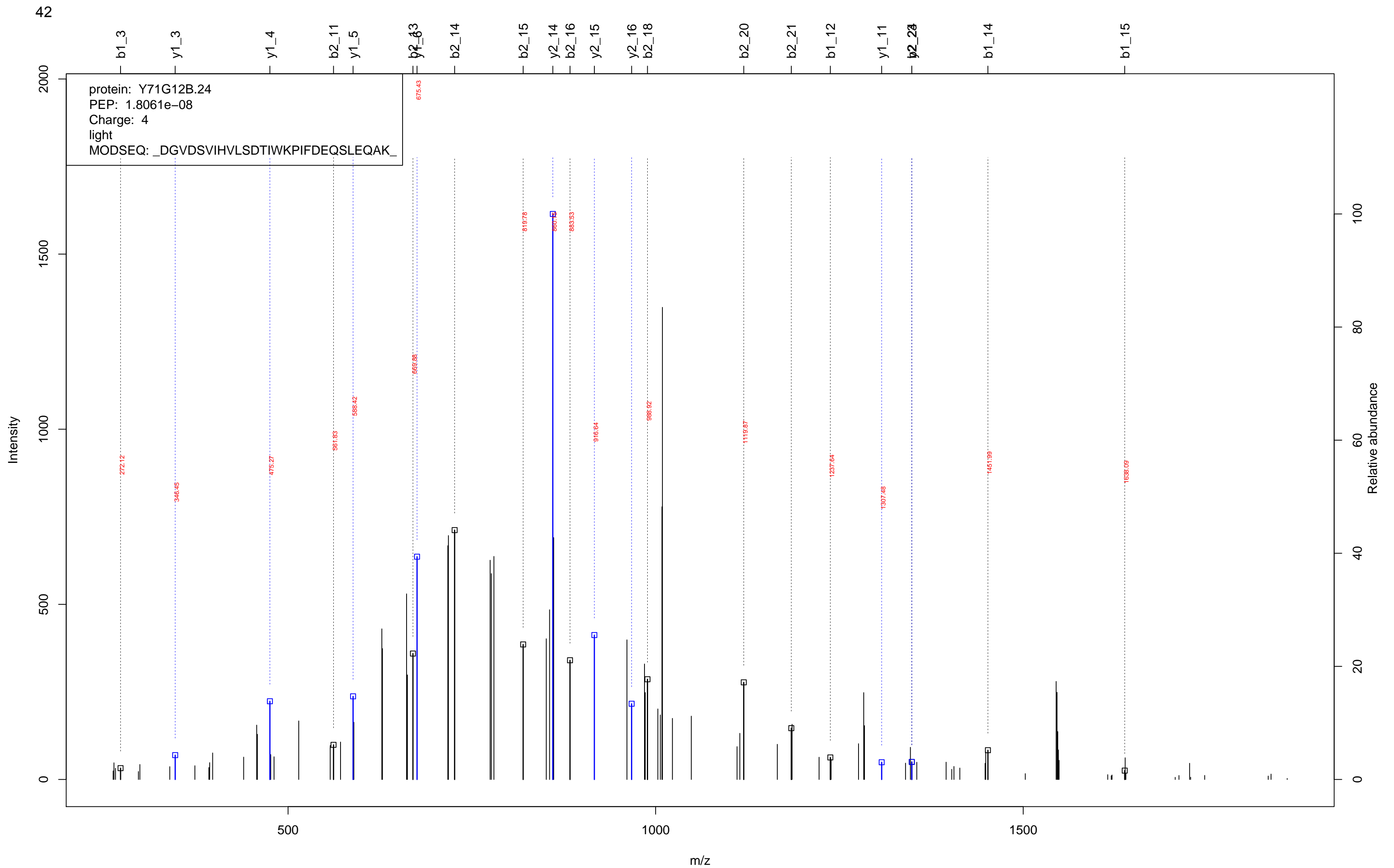


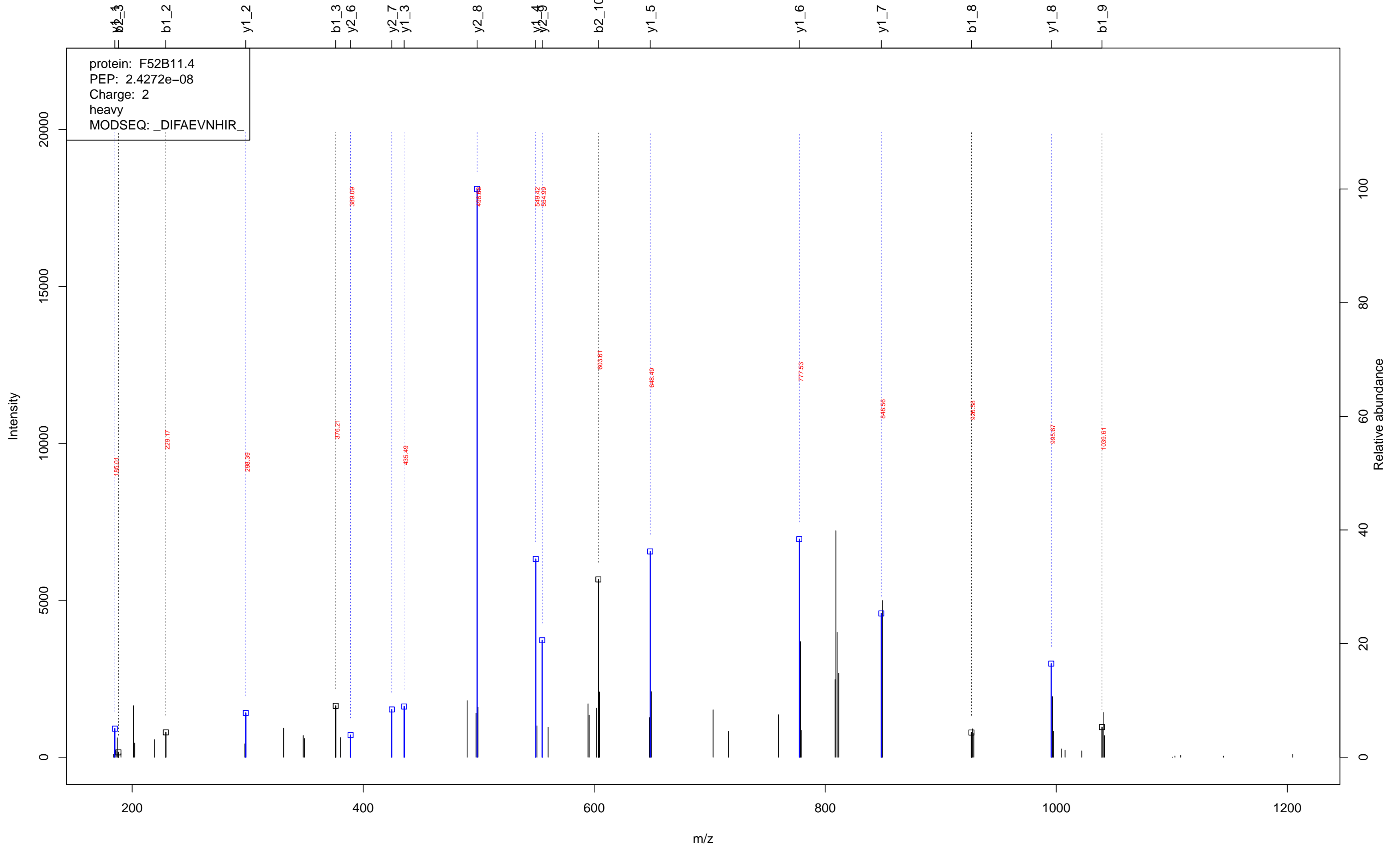


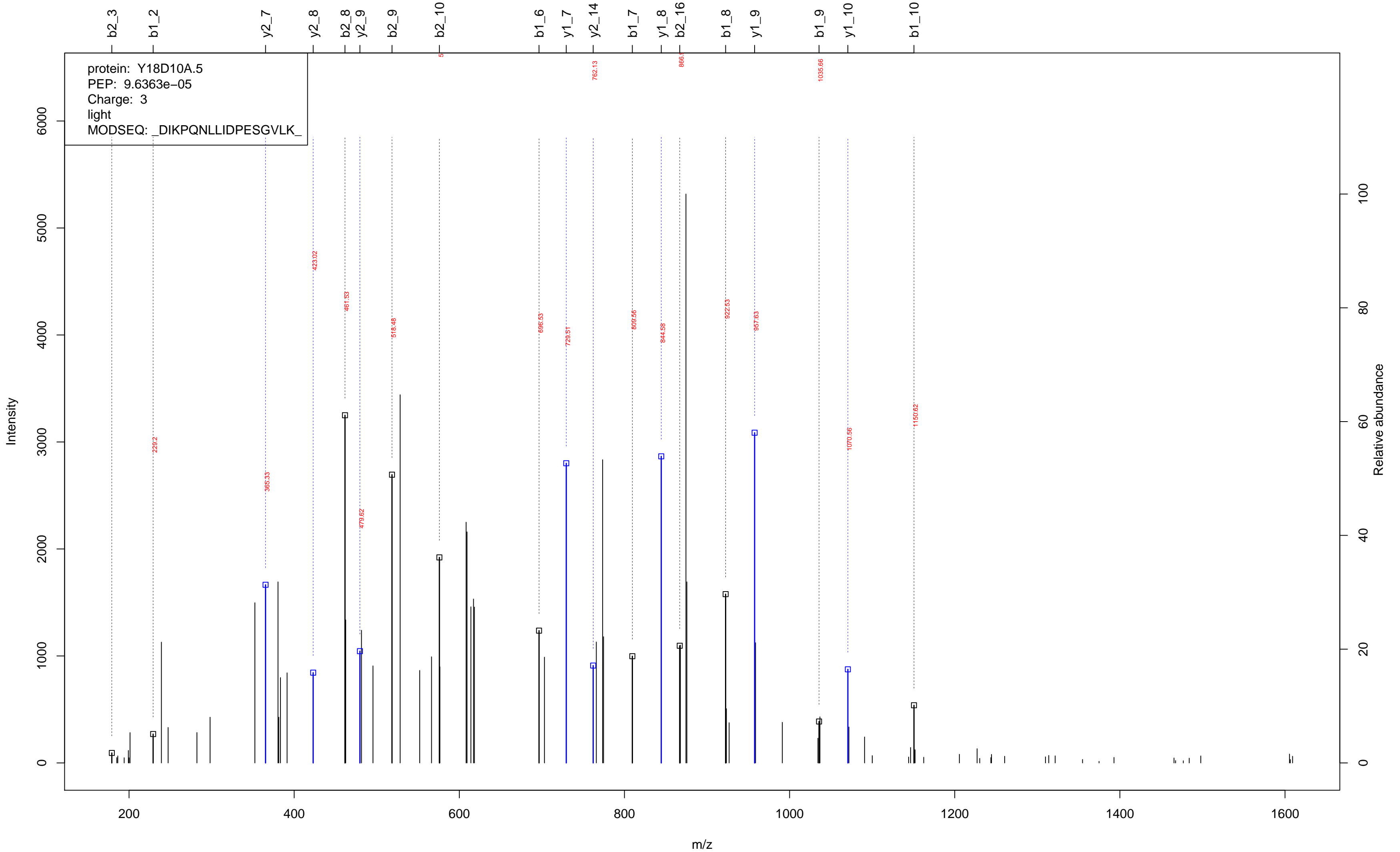


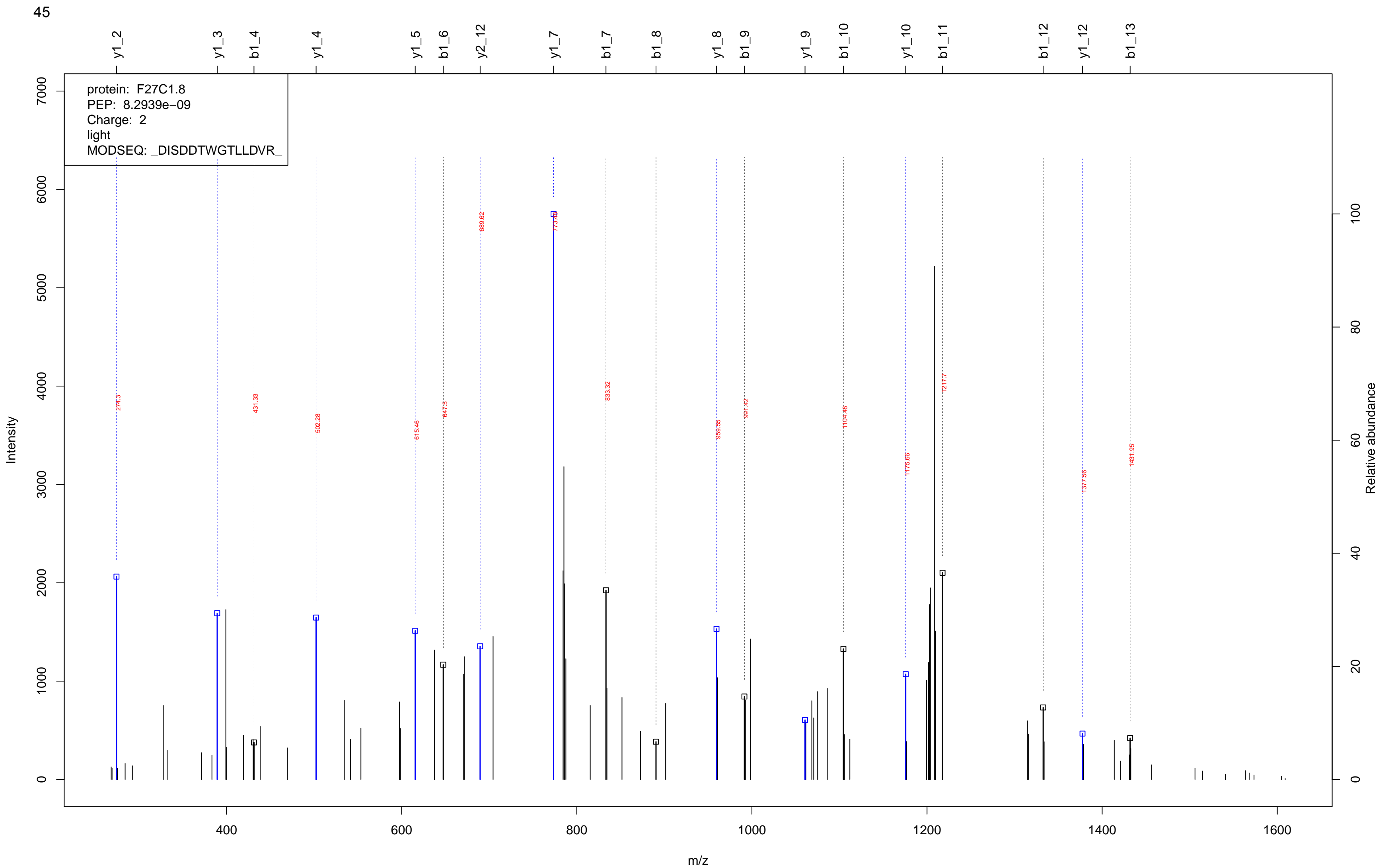


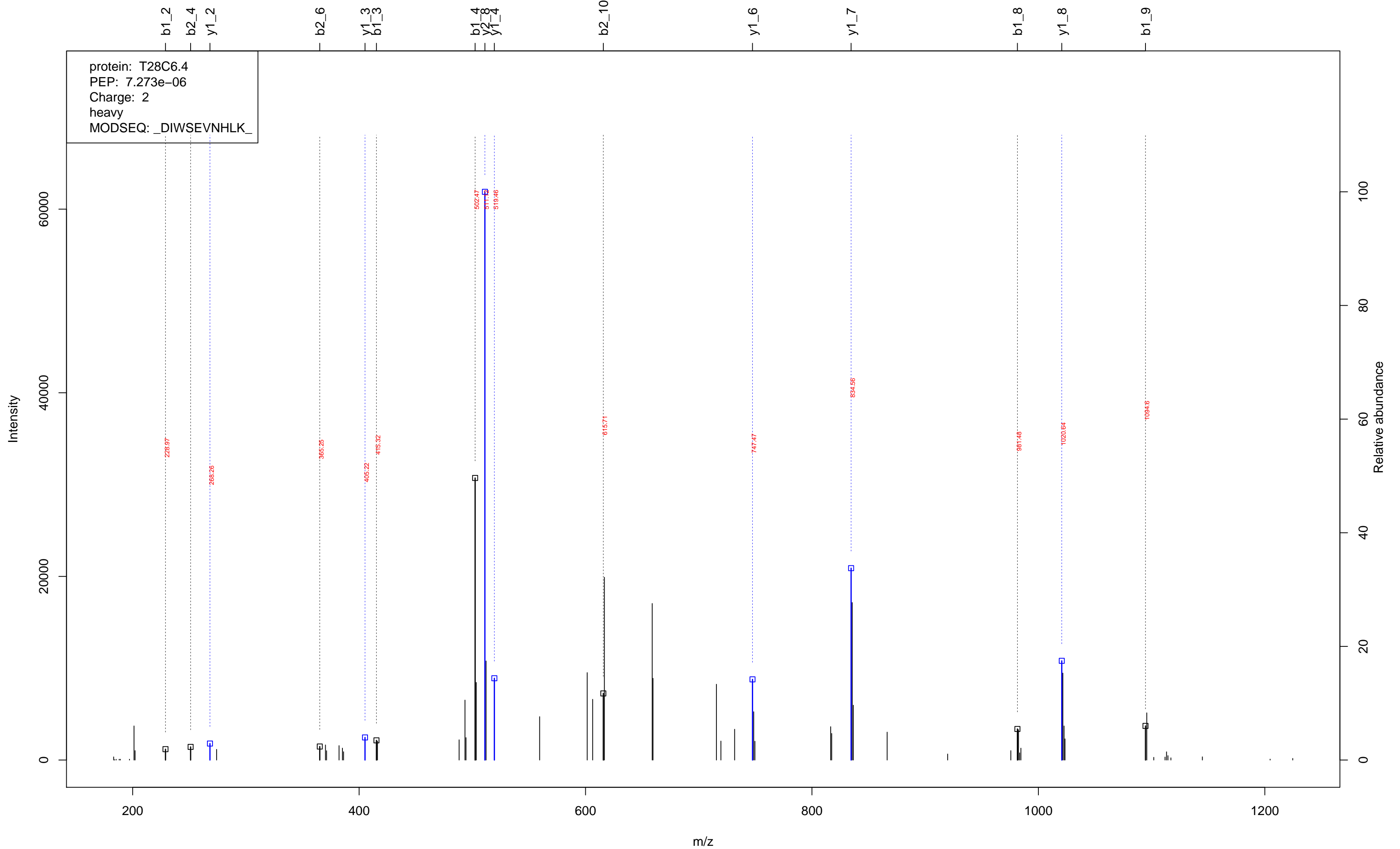


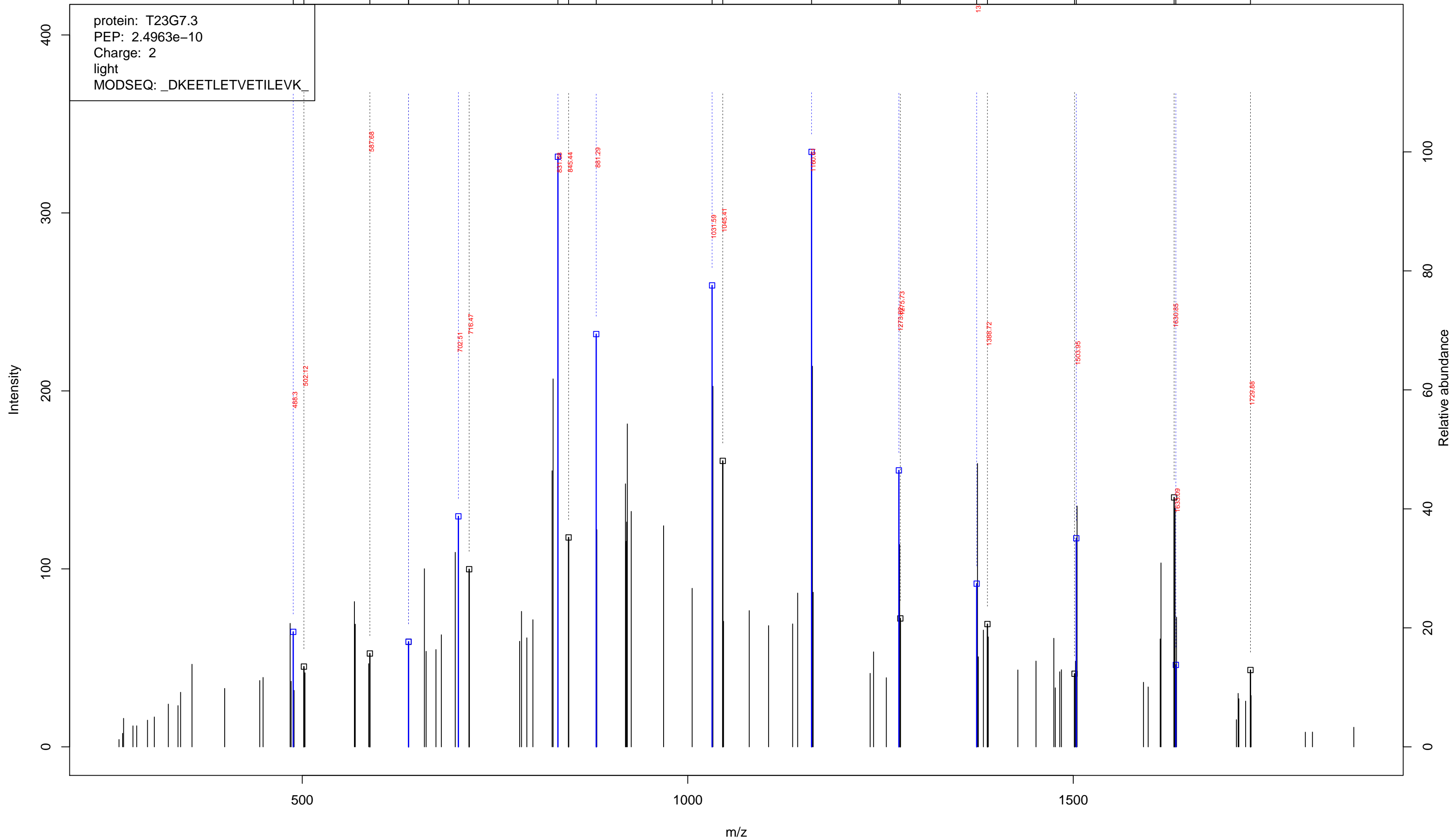


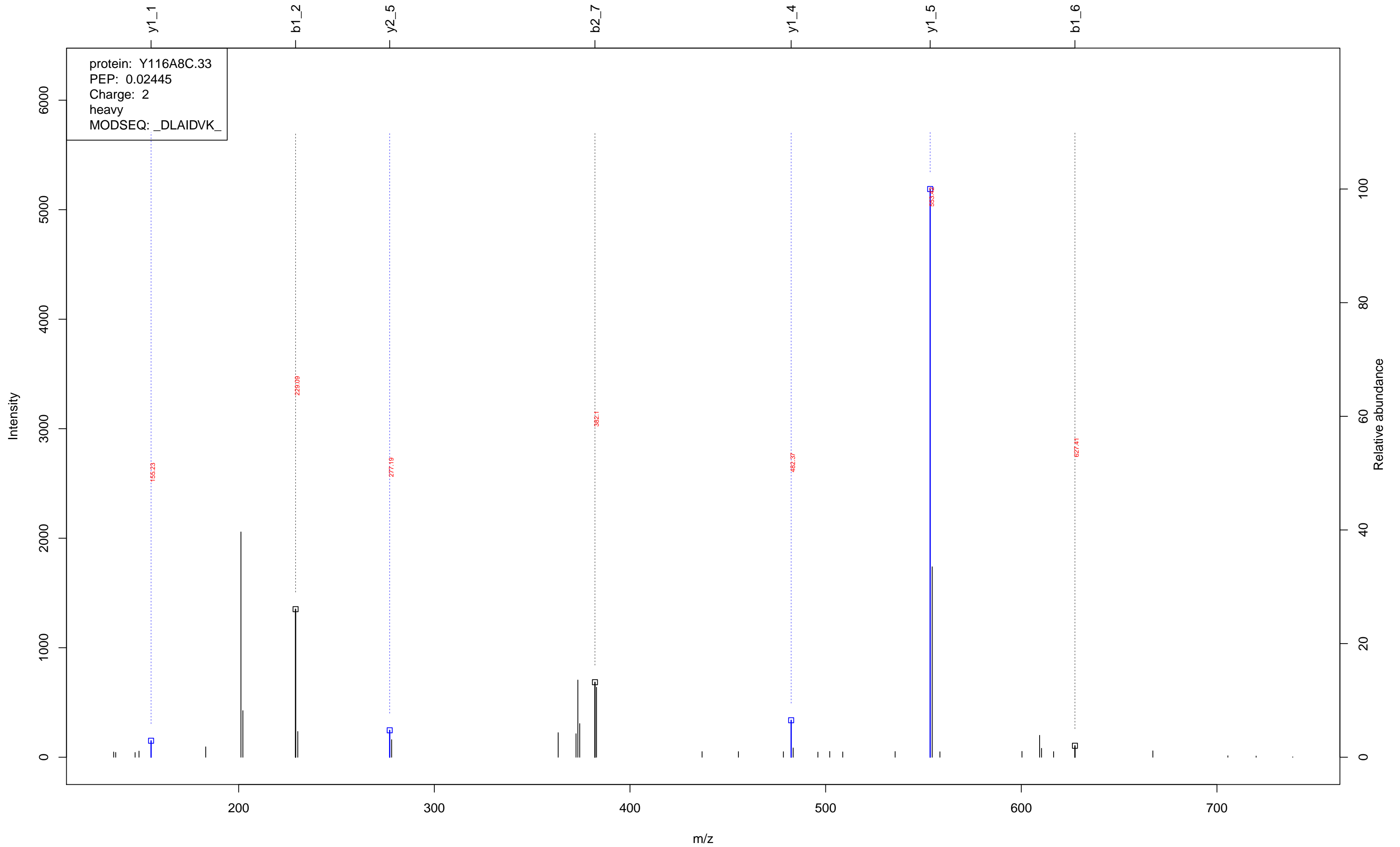


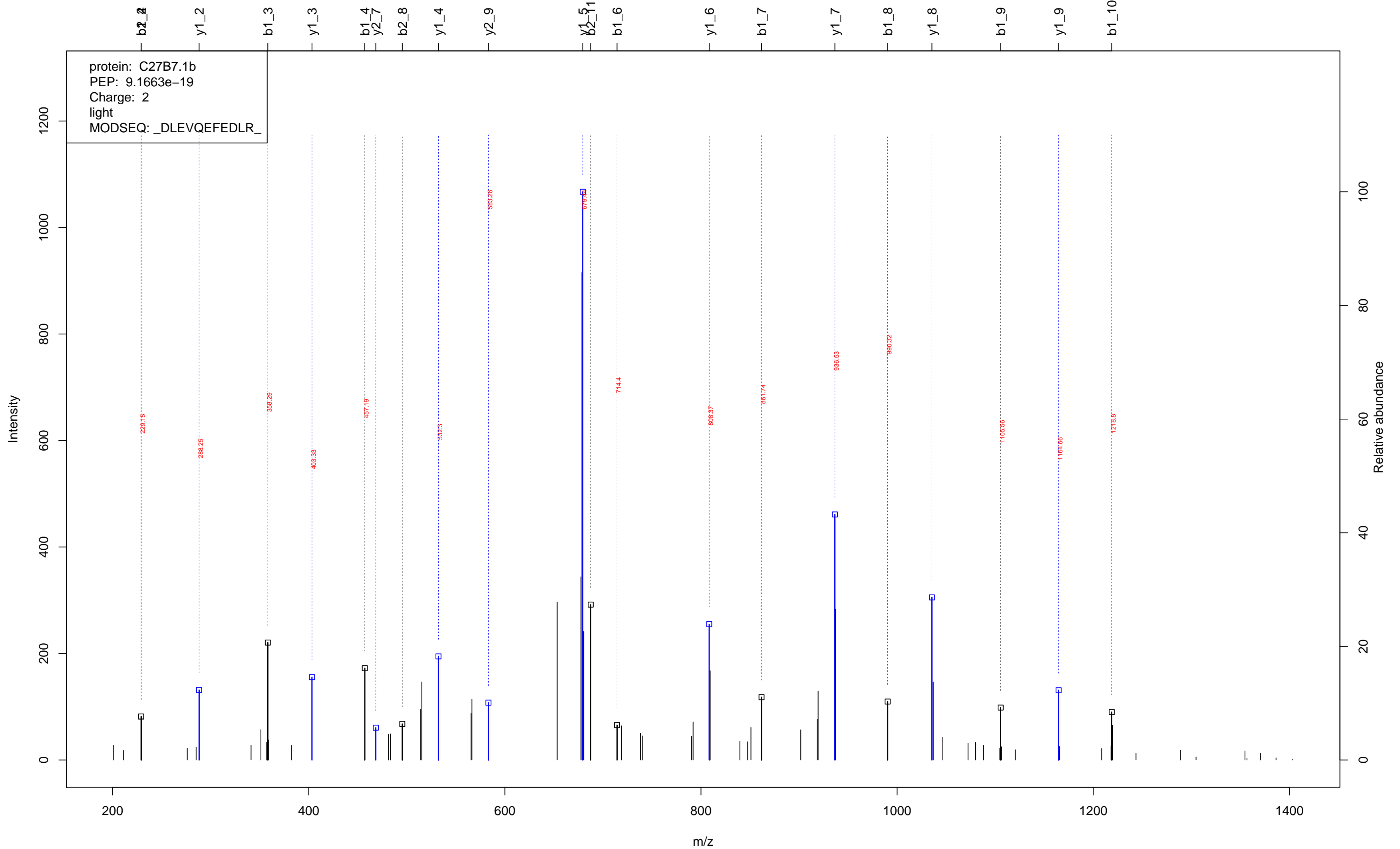


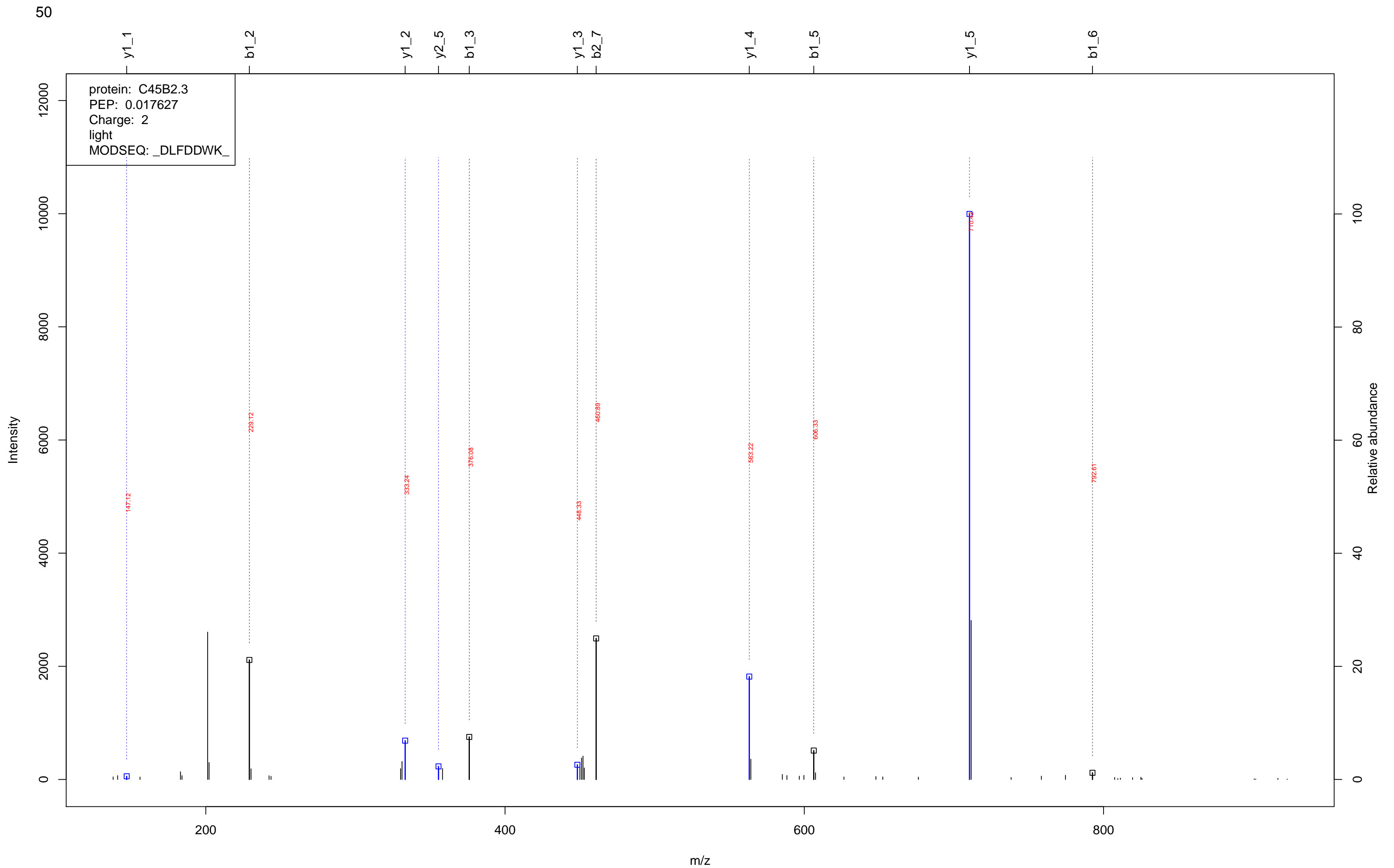


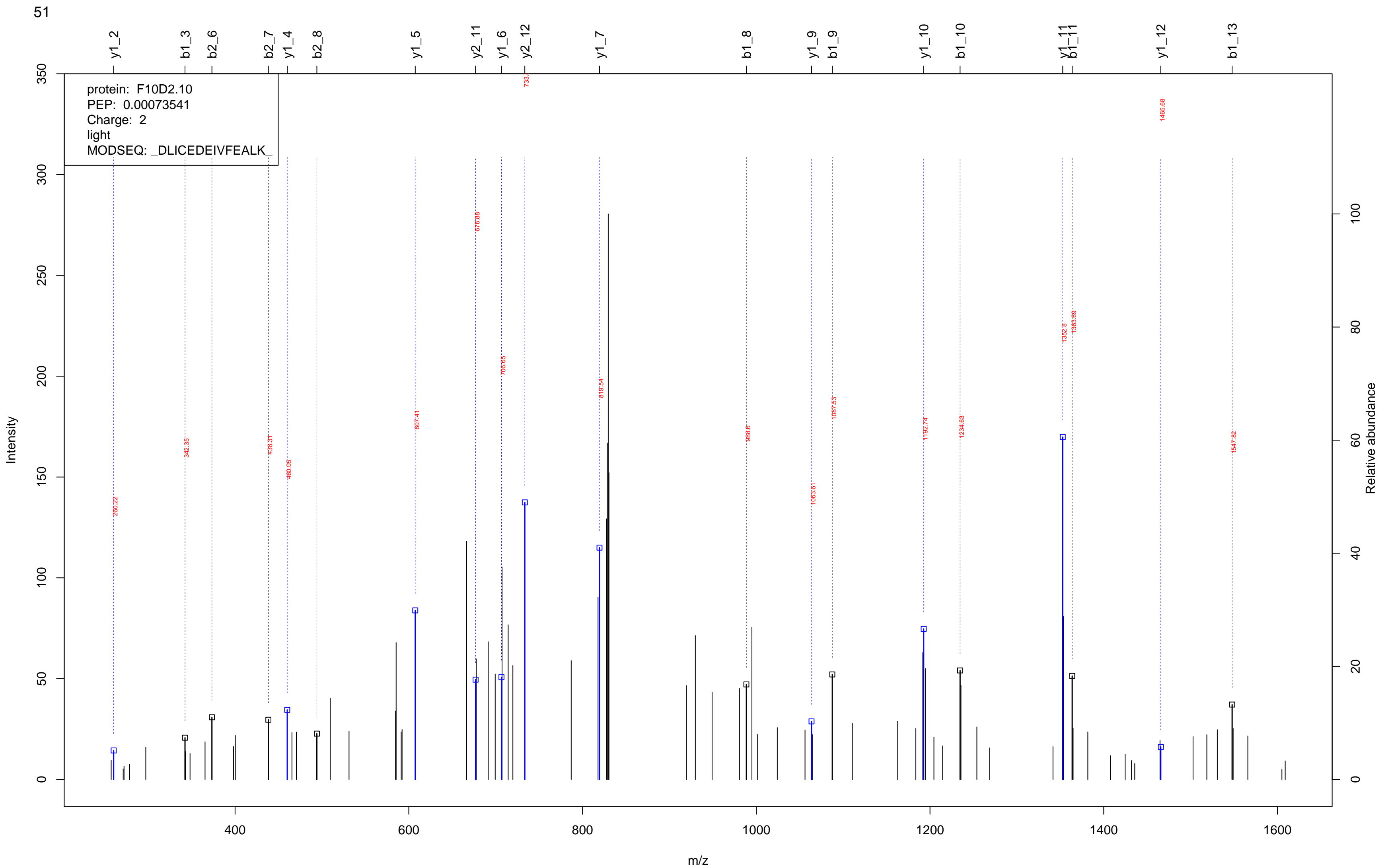


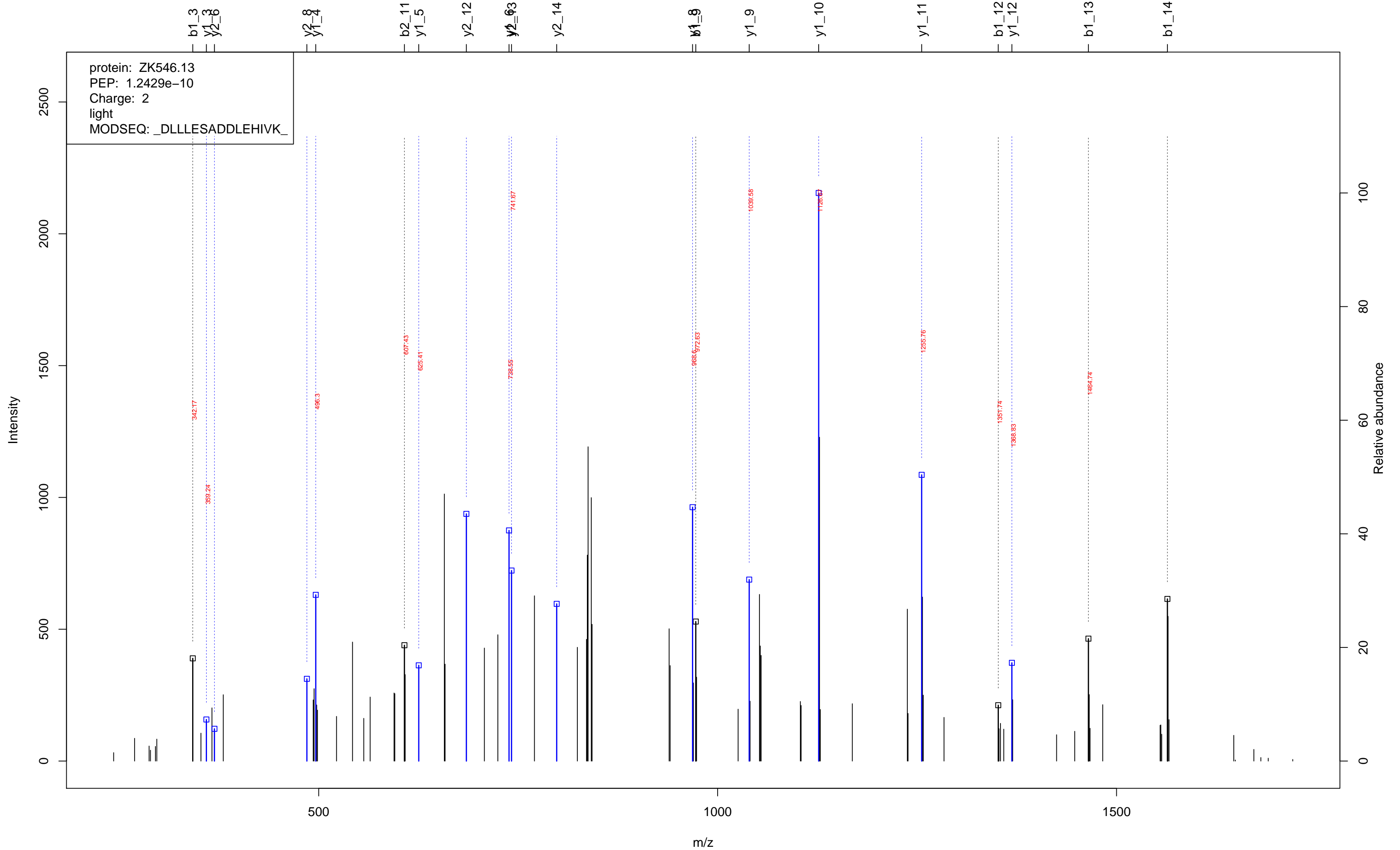


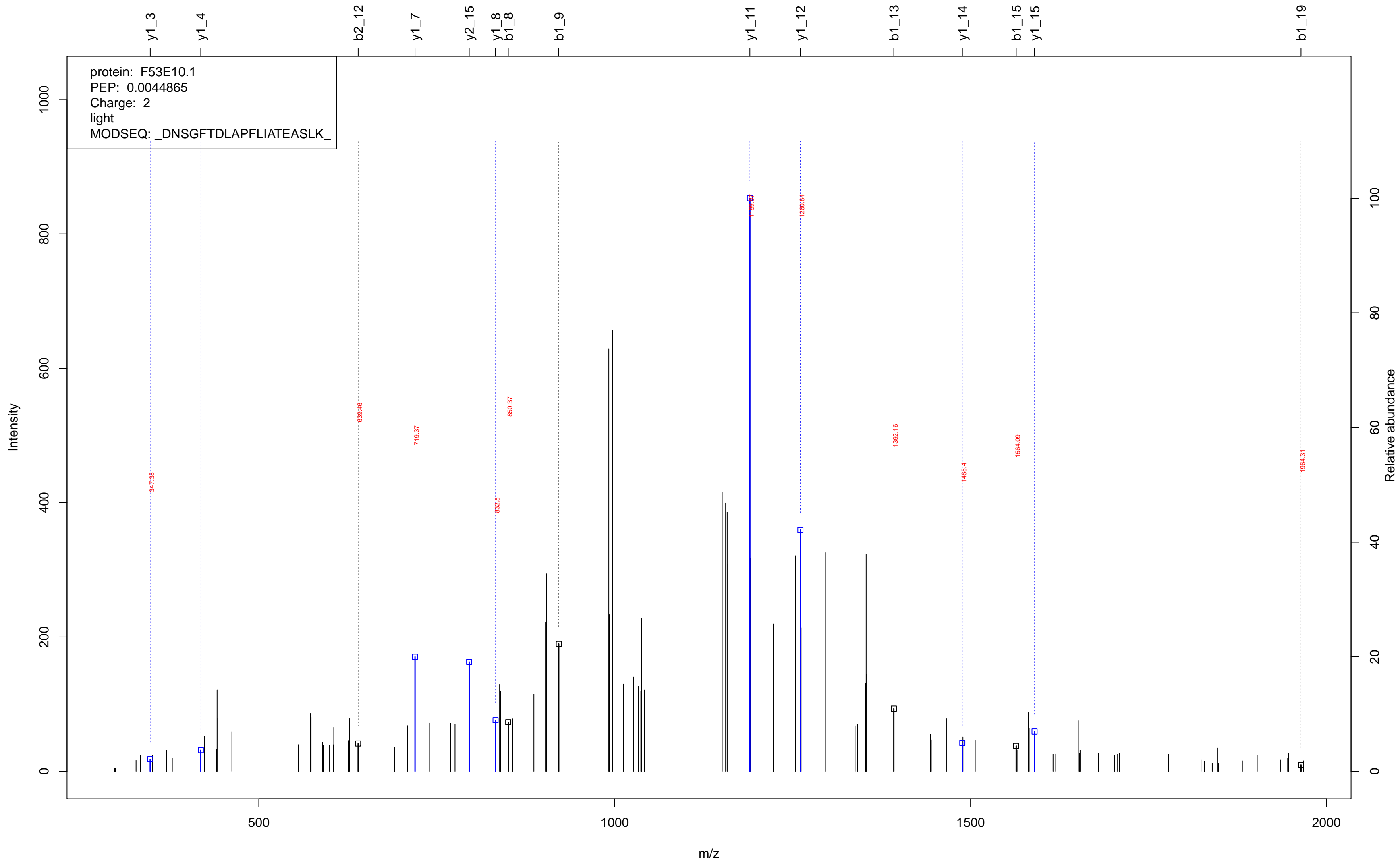


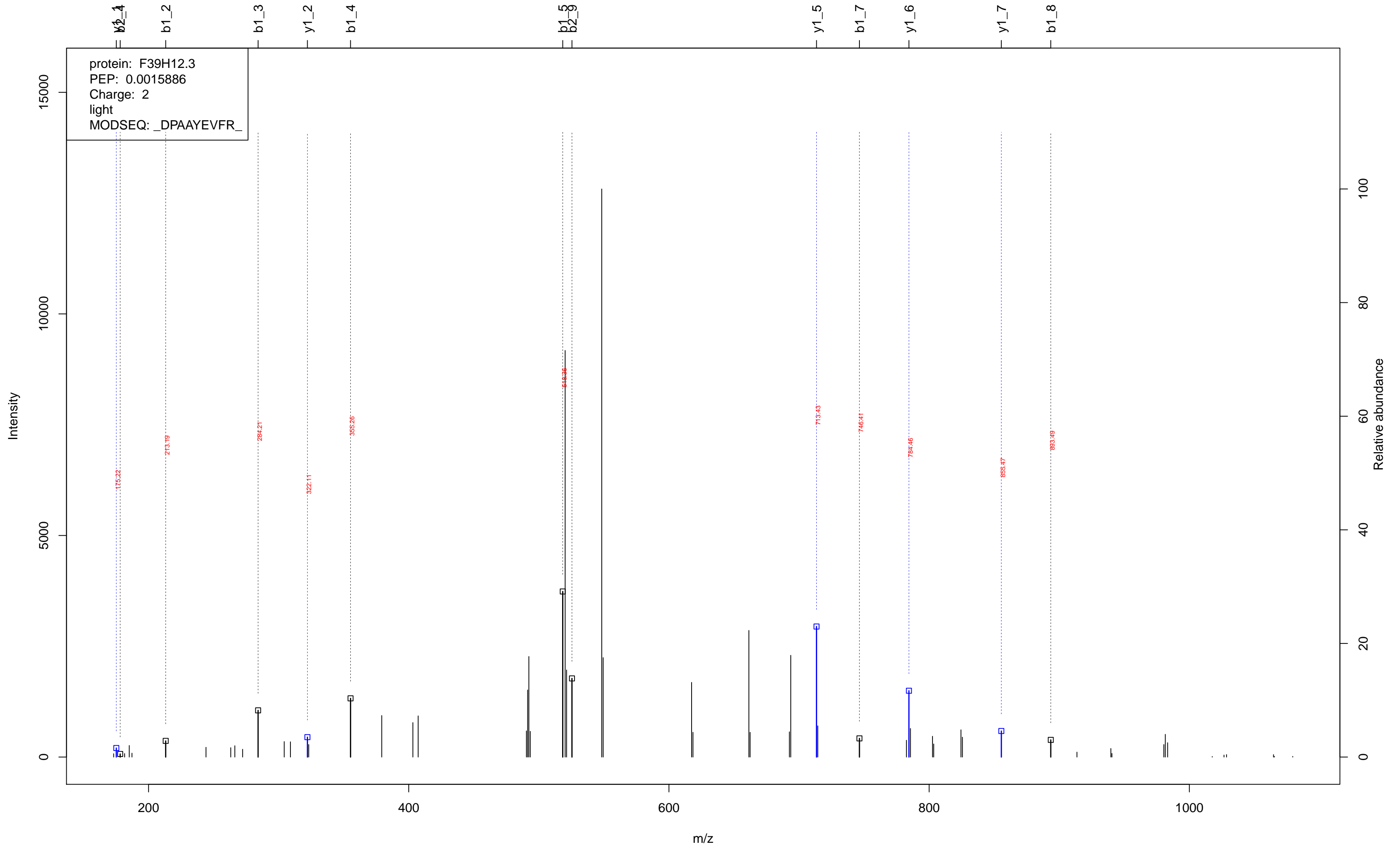


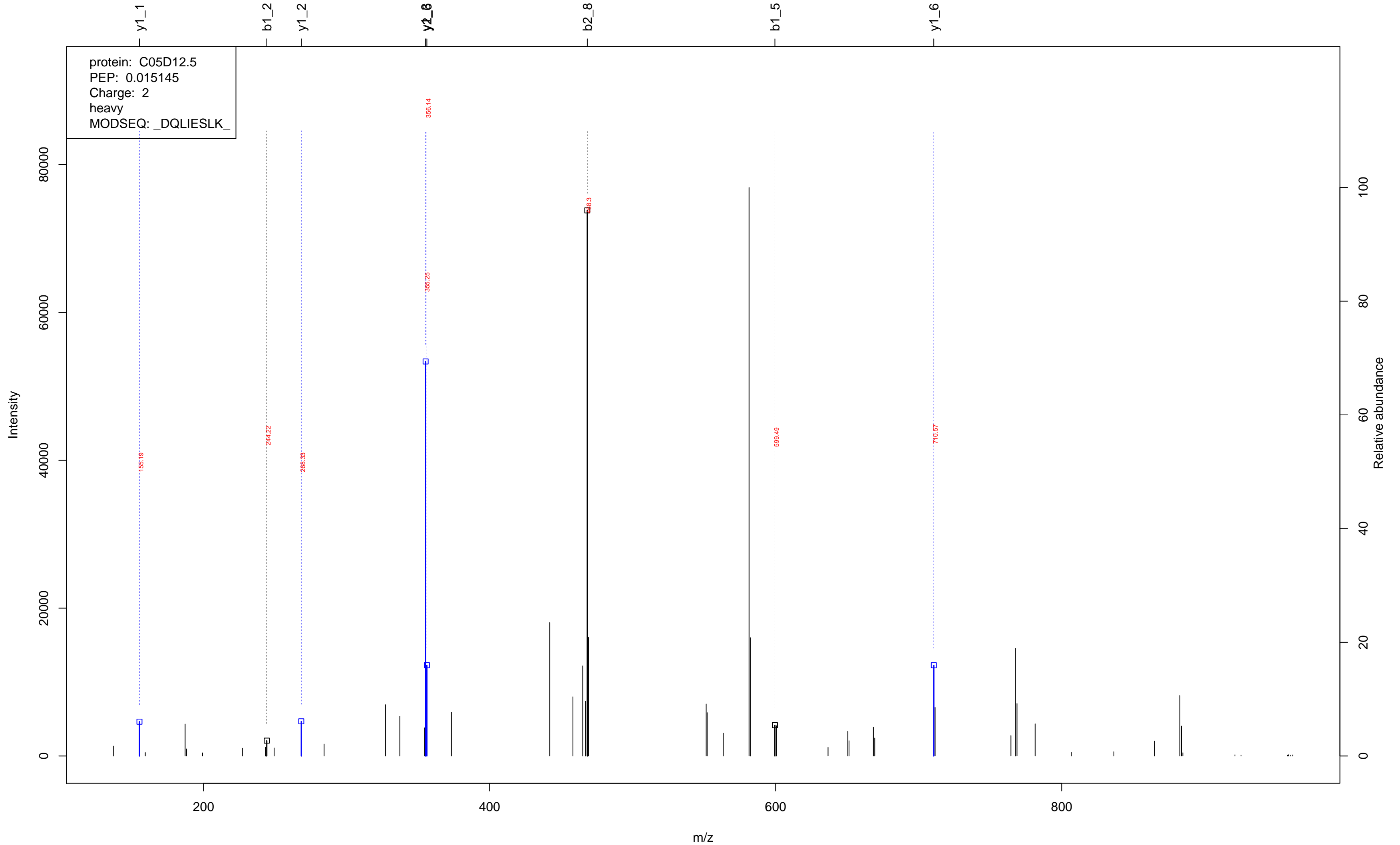


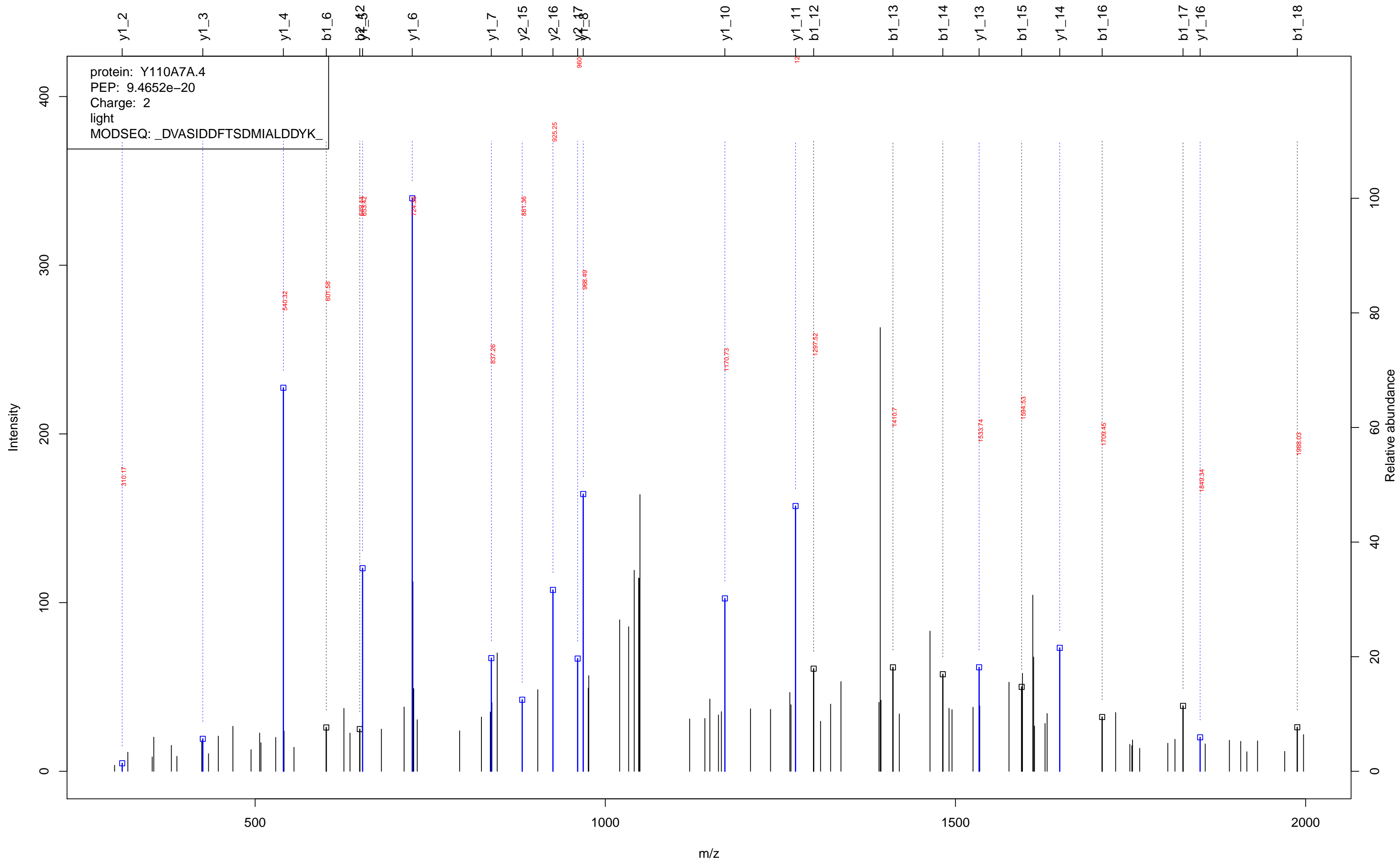


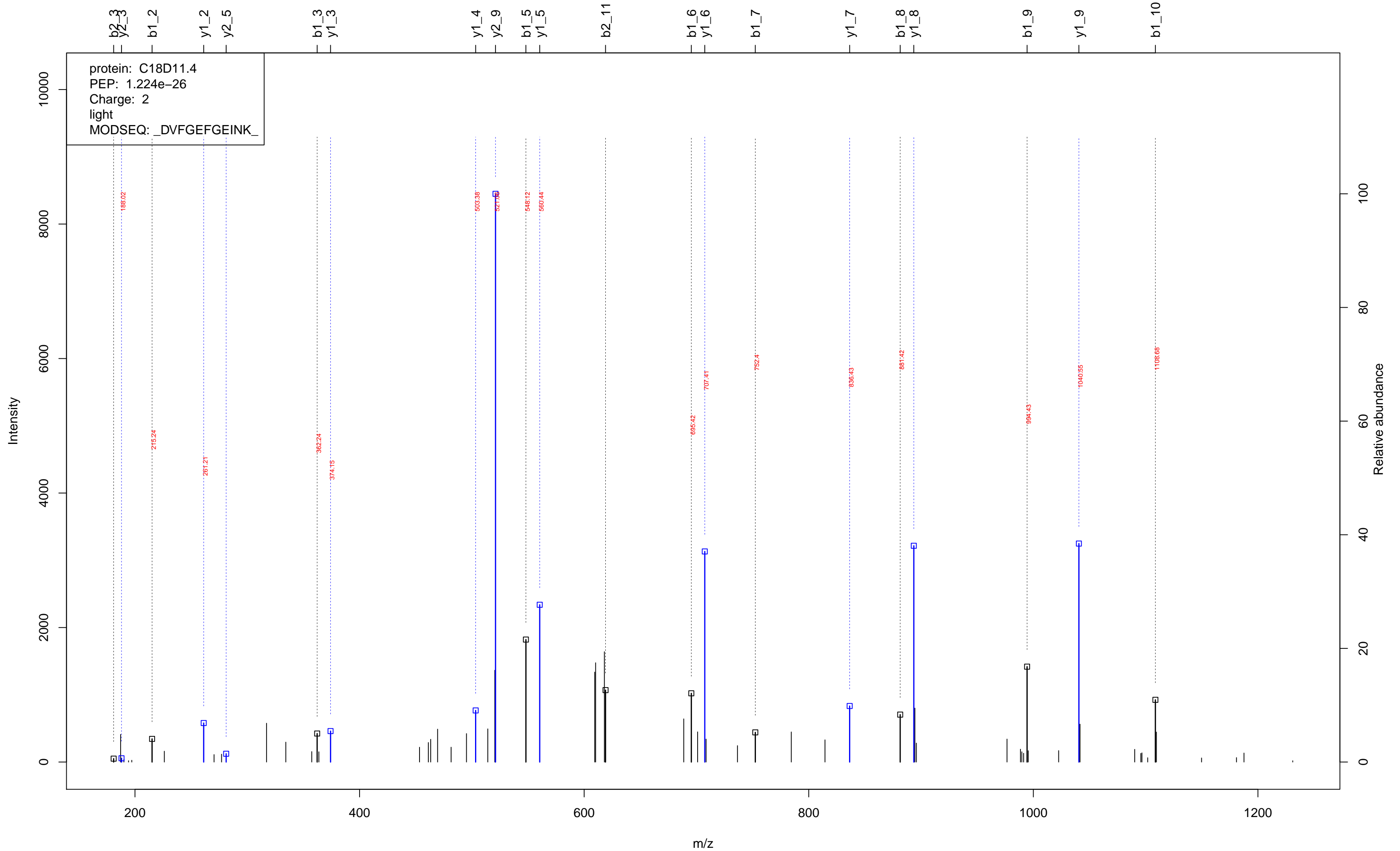


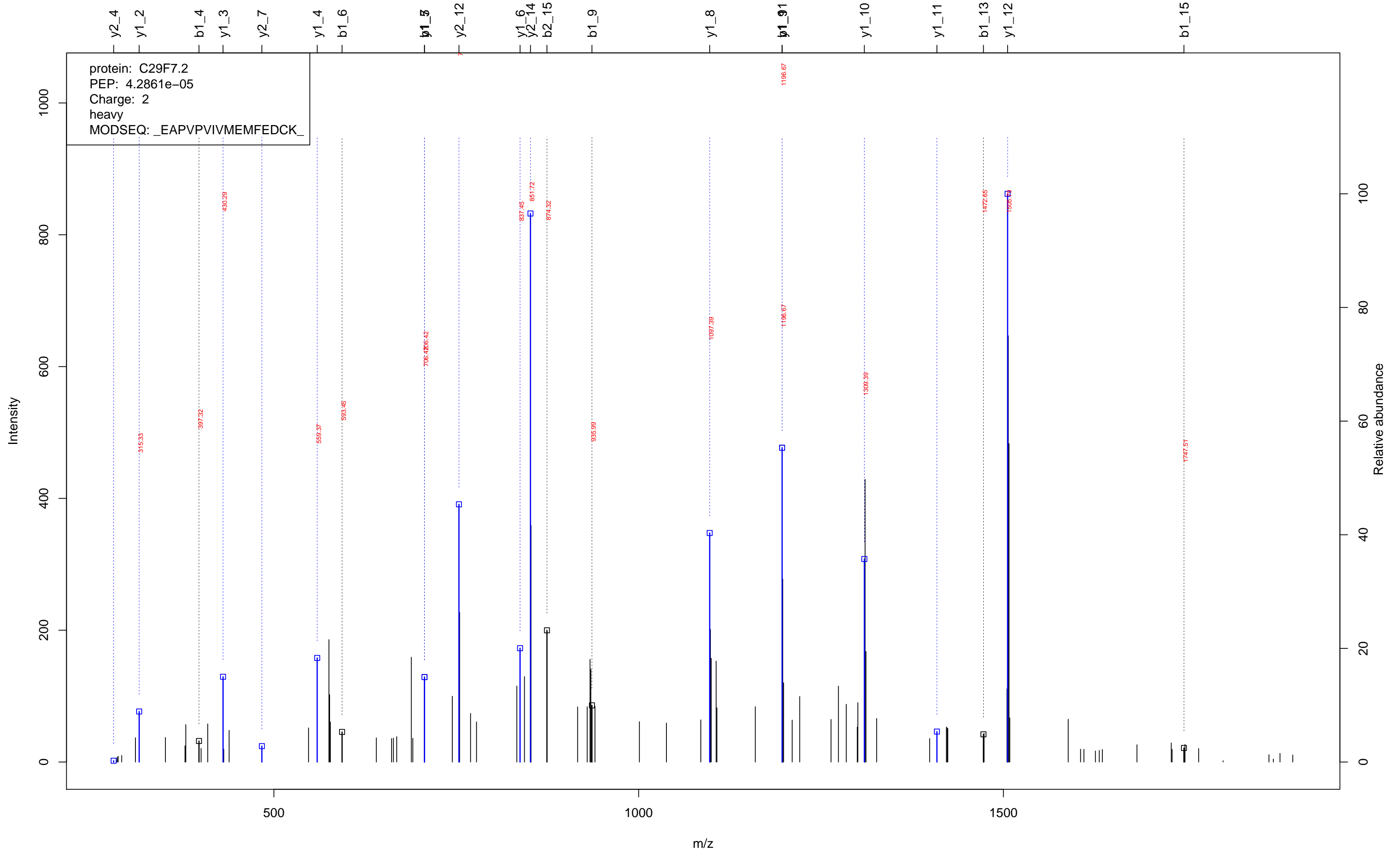


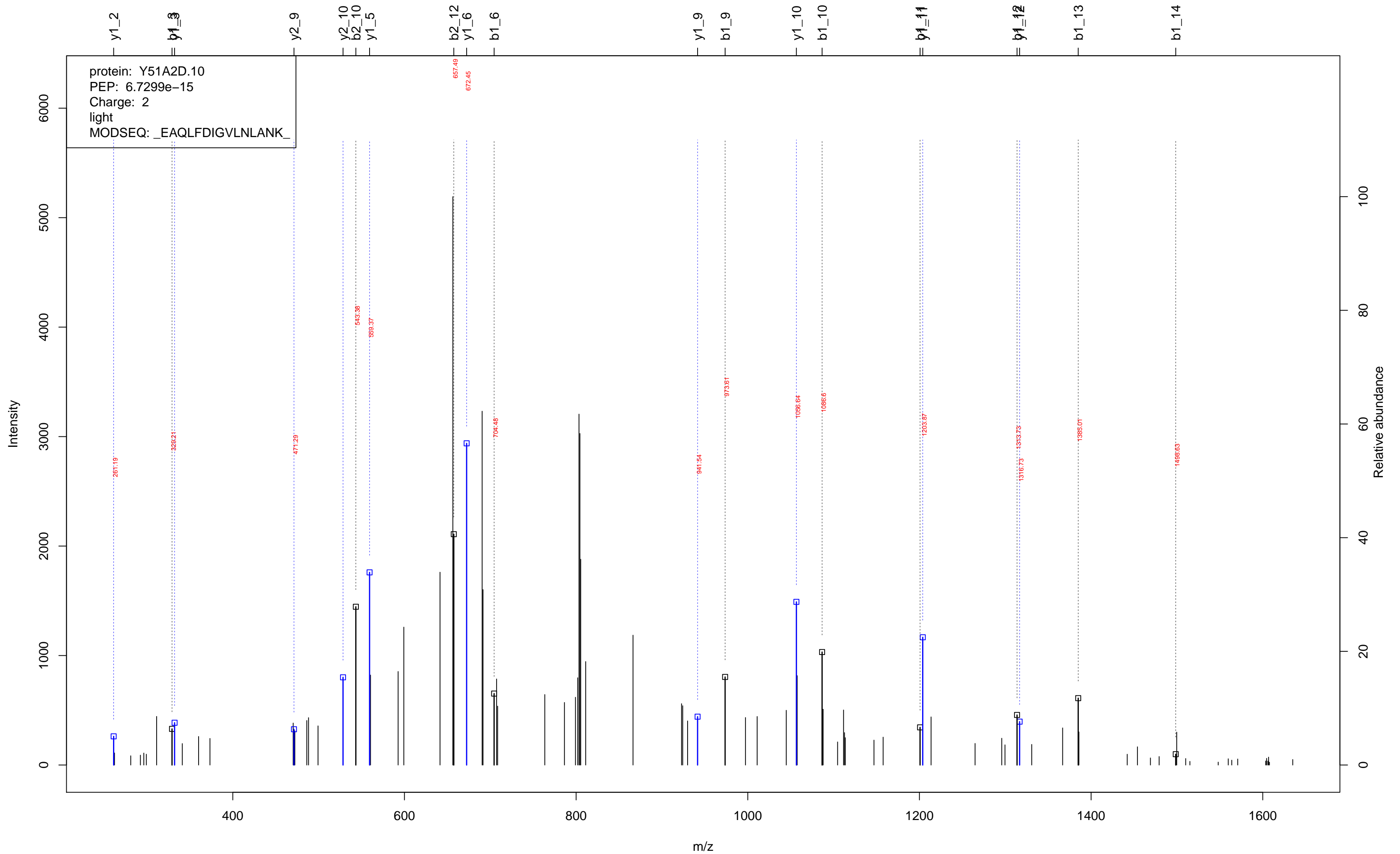


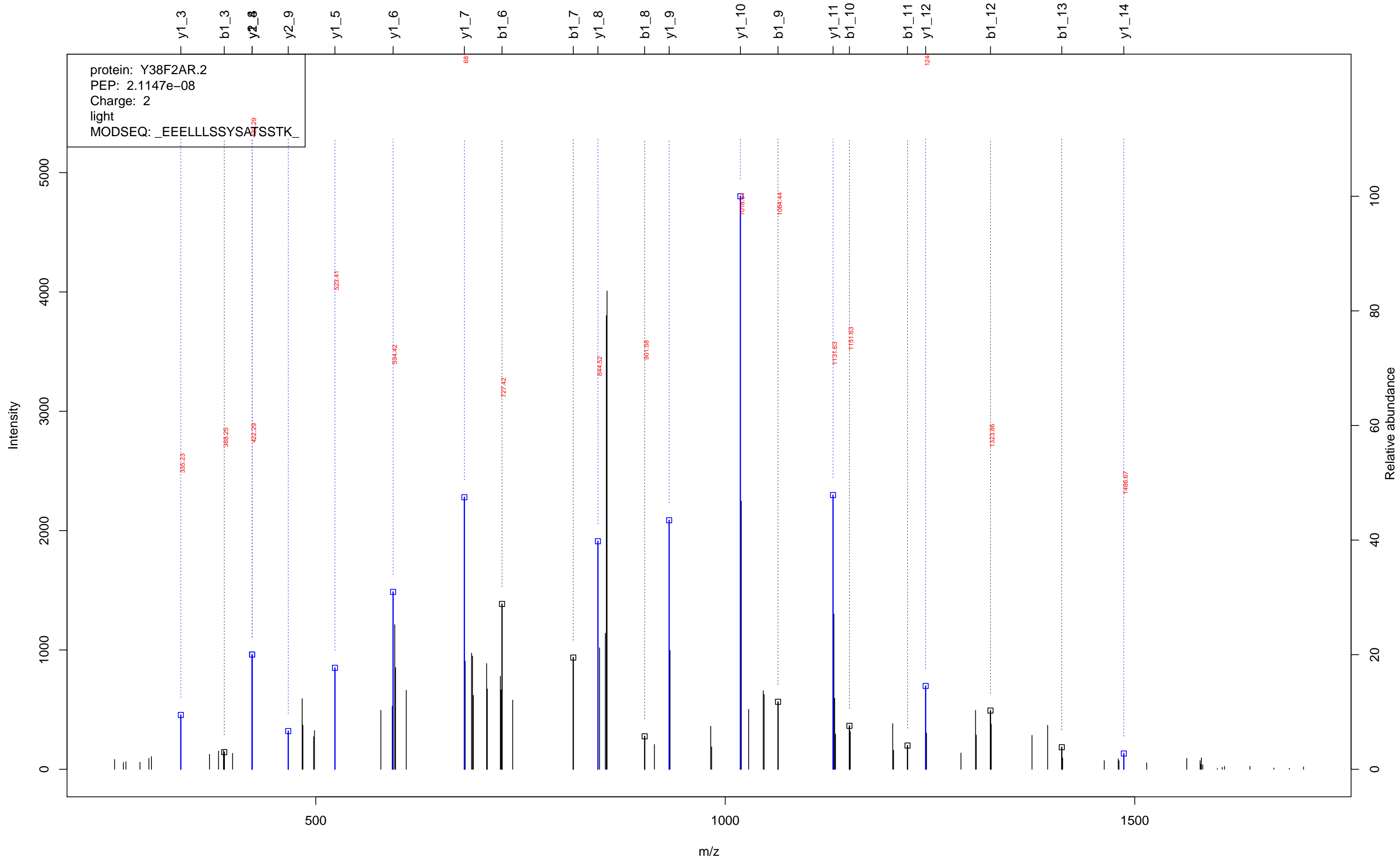


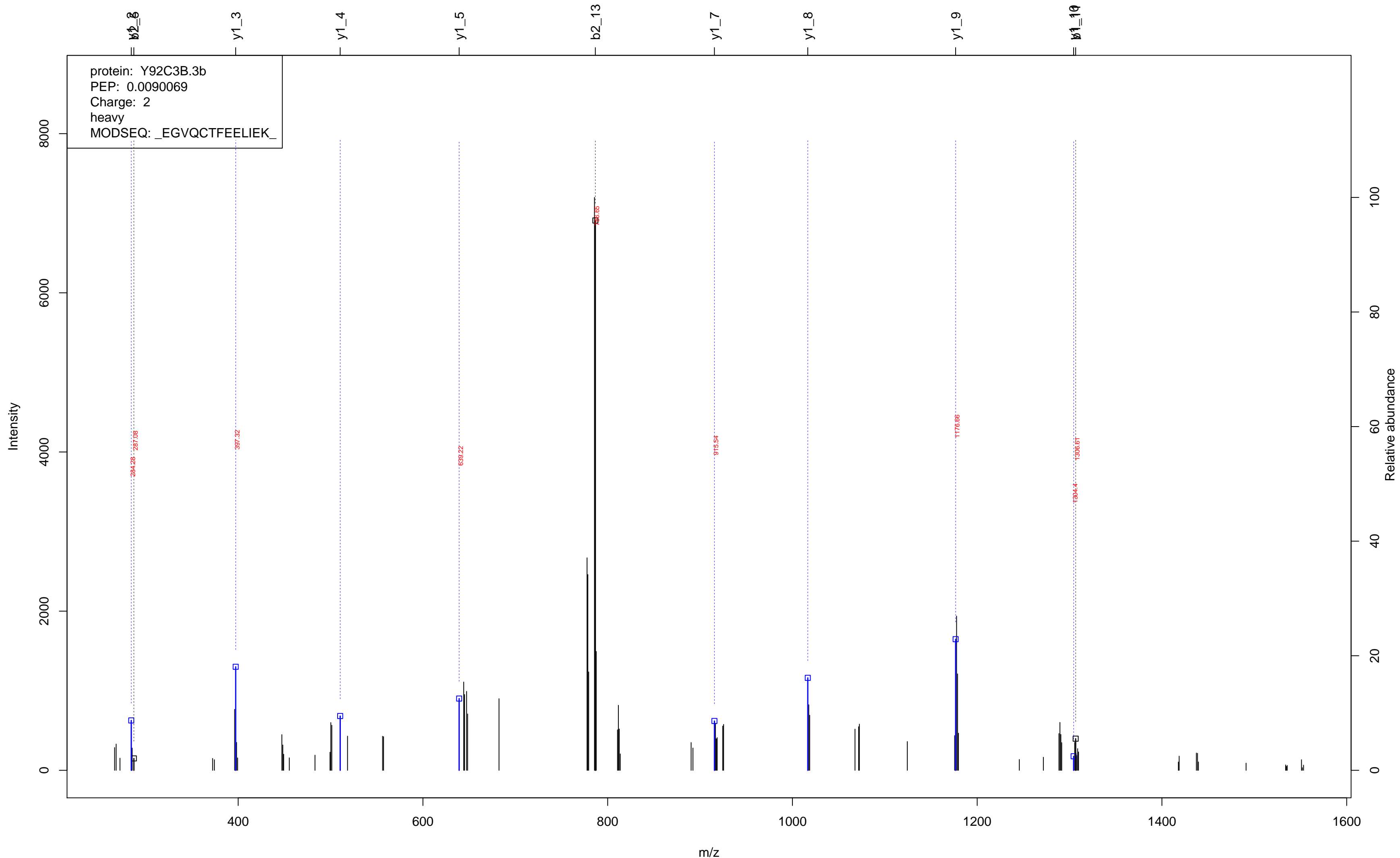


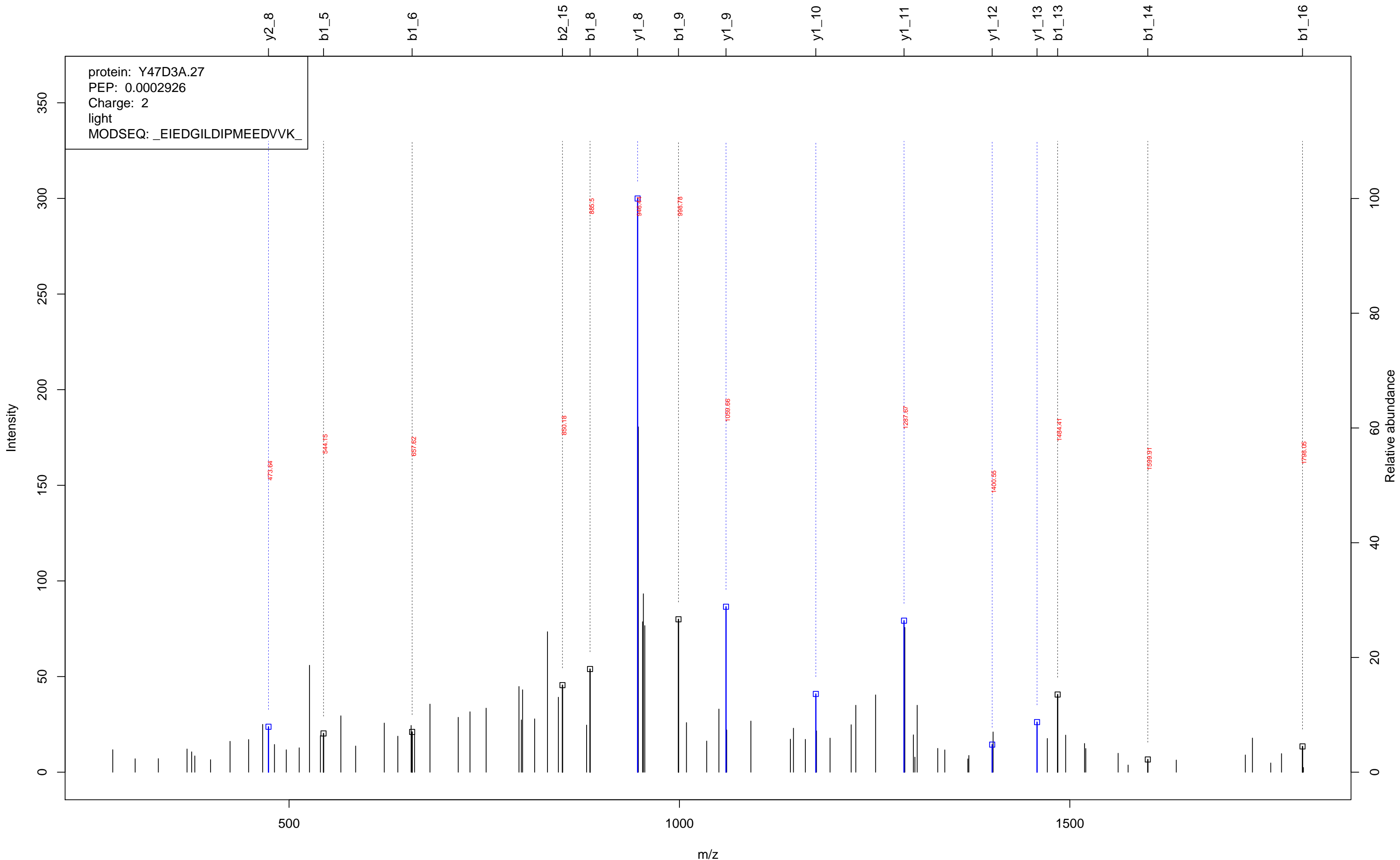






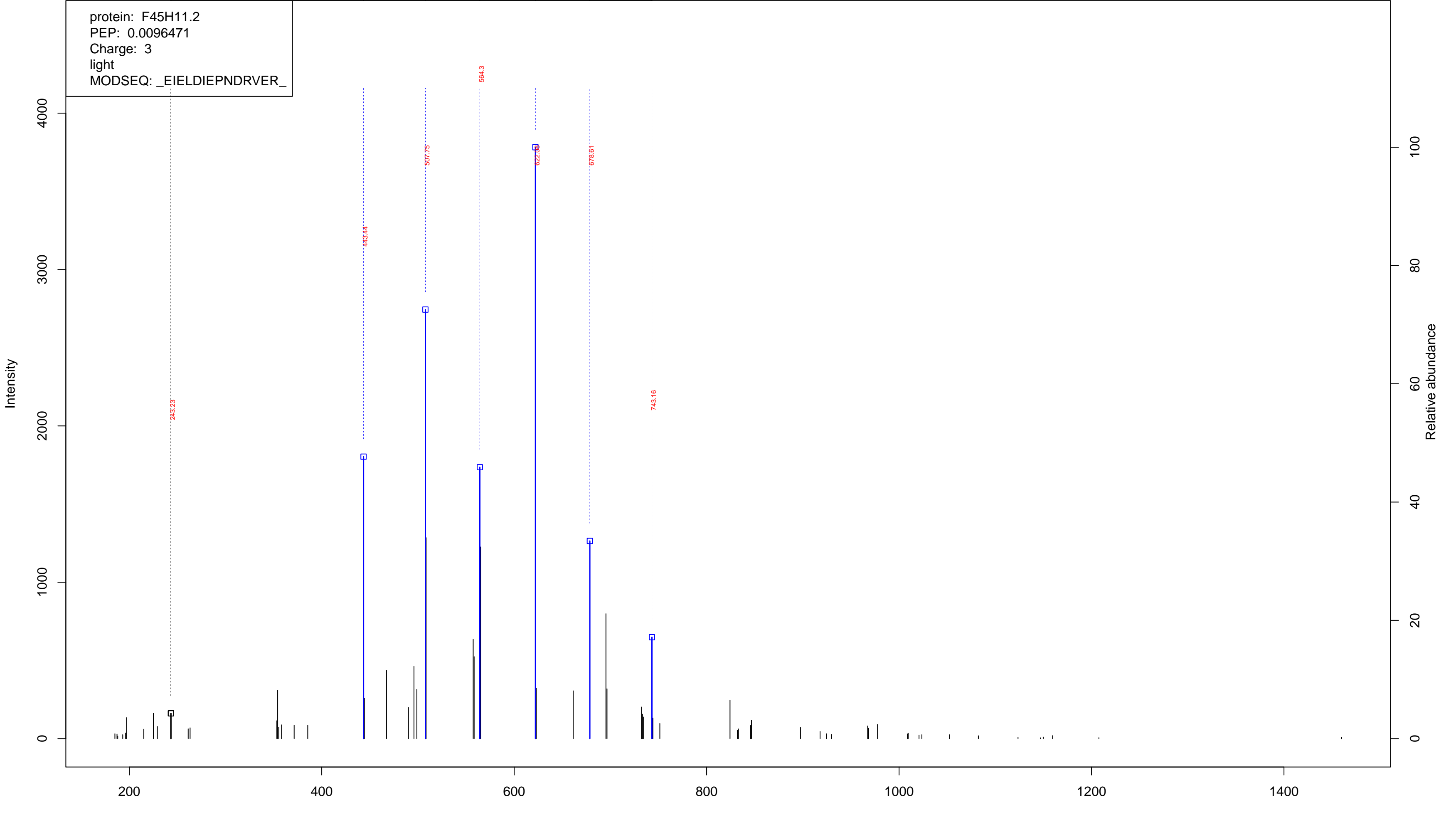


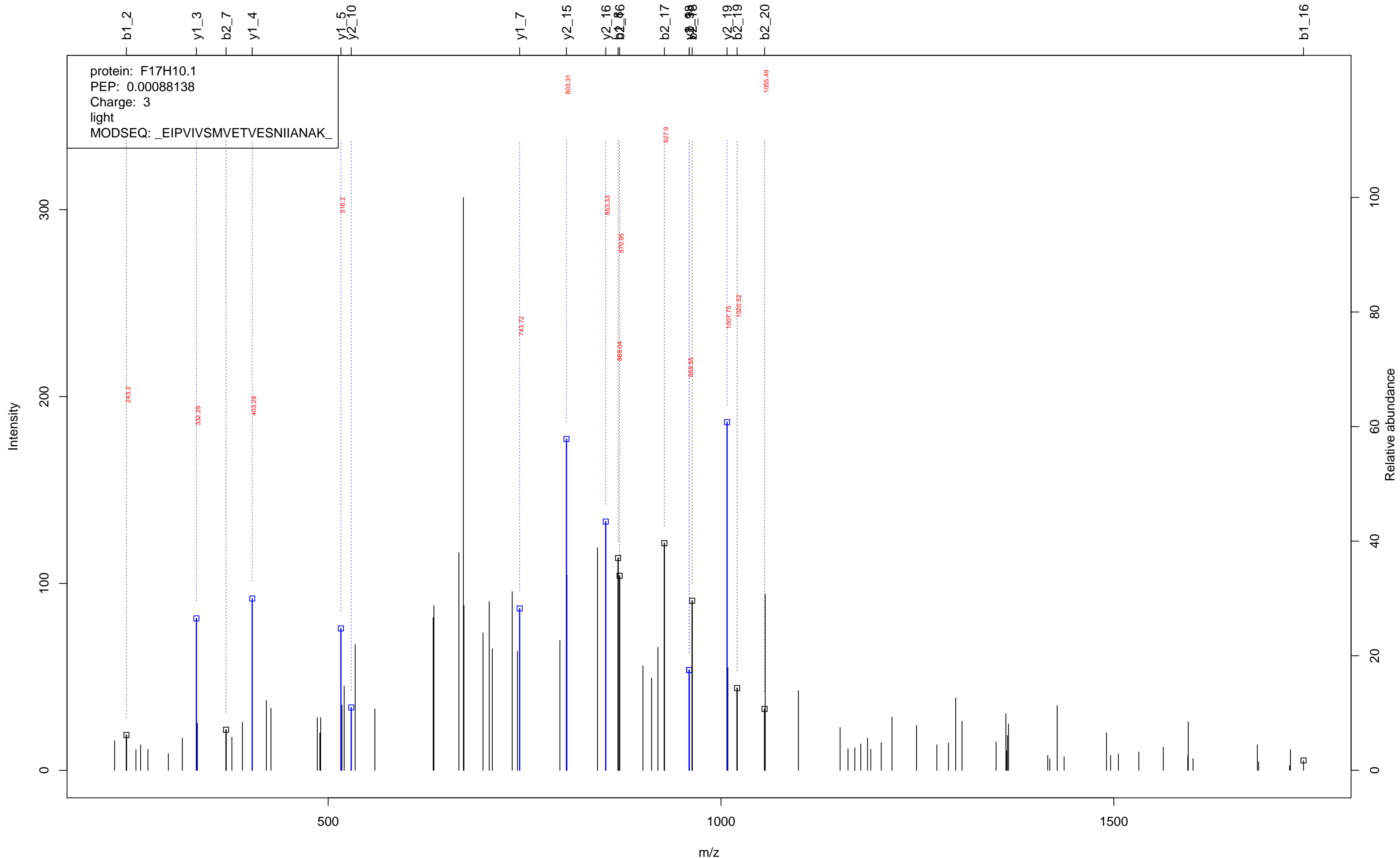


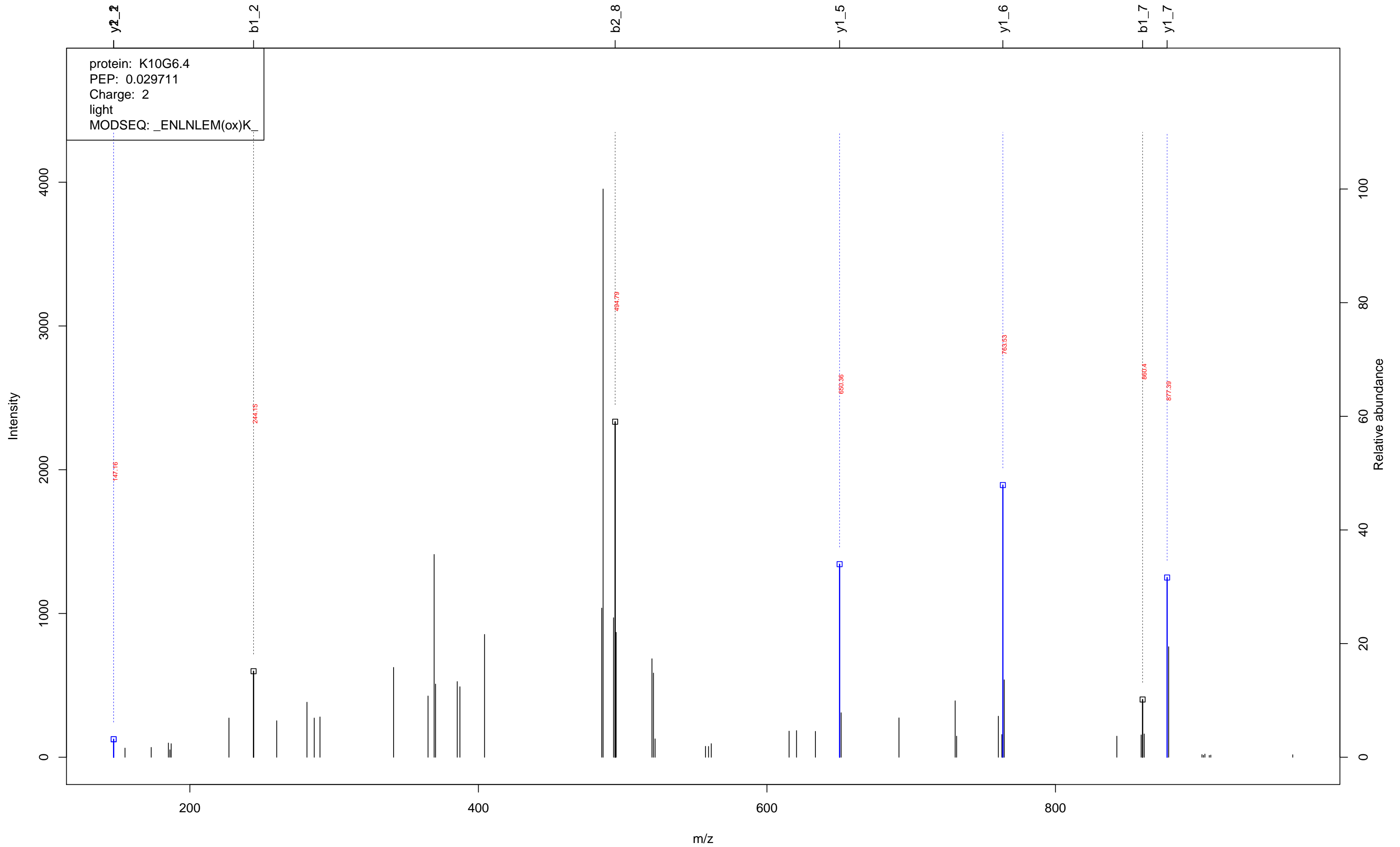


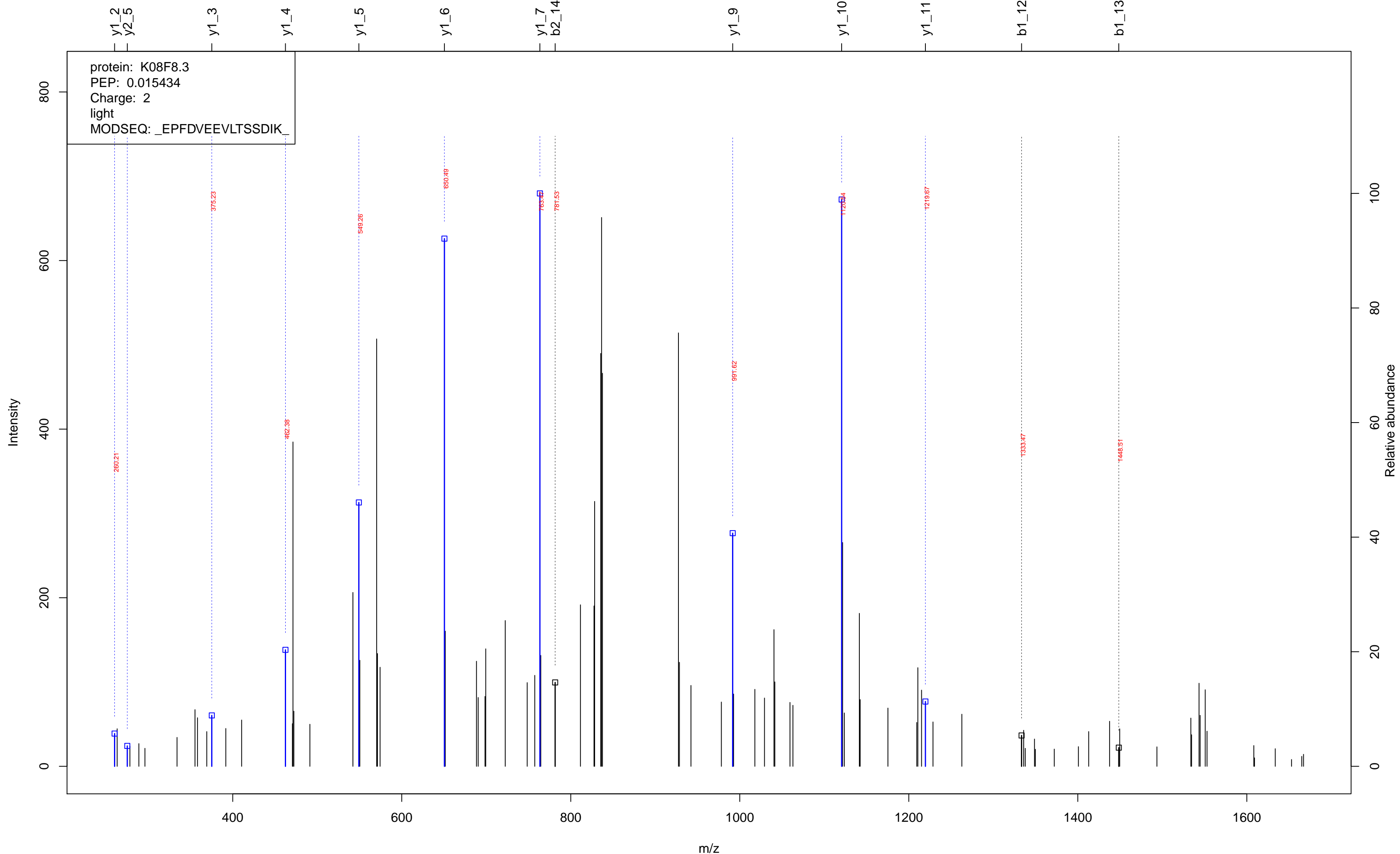
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y2_8
y2_9
y2_10
y2_11
y2_12

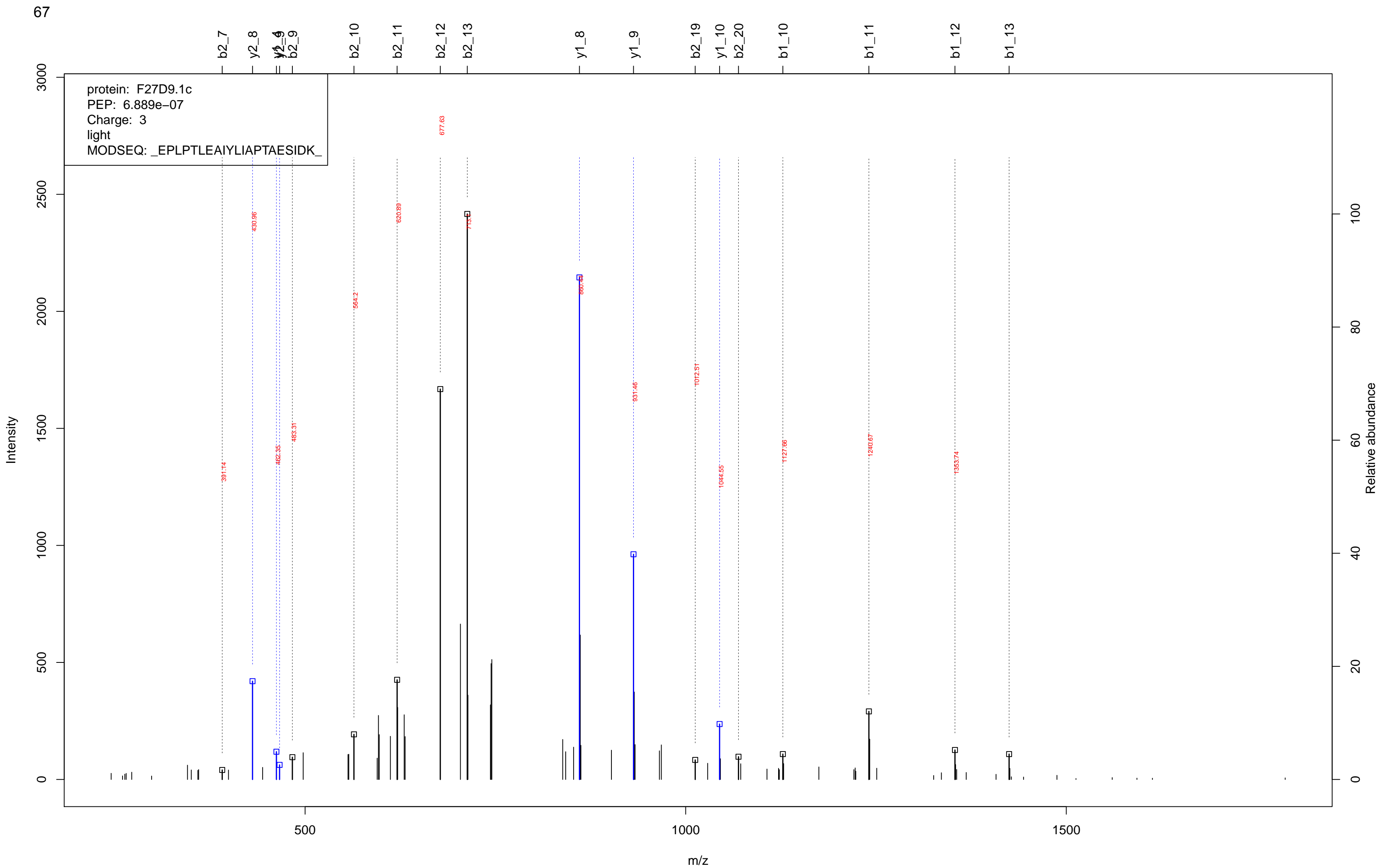
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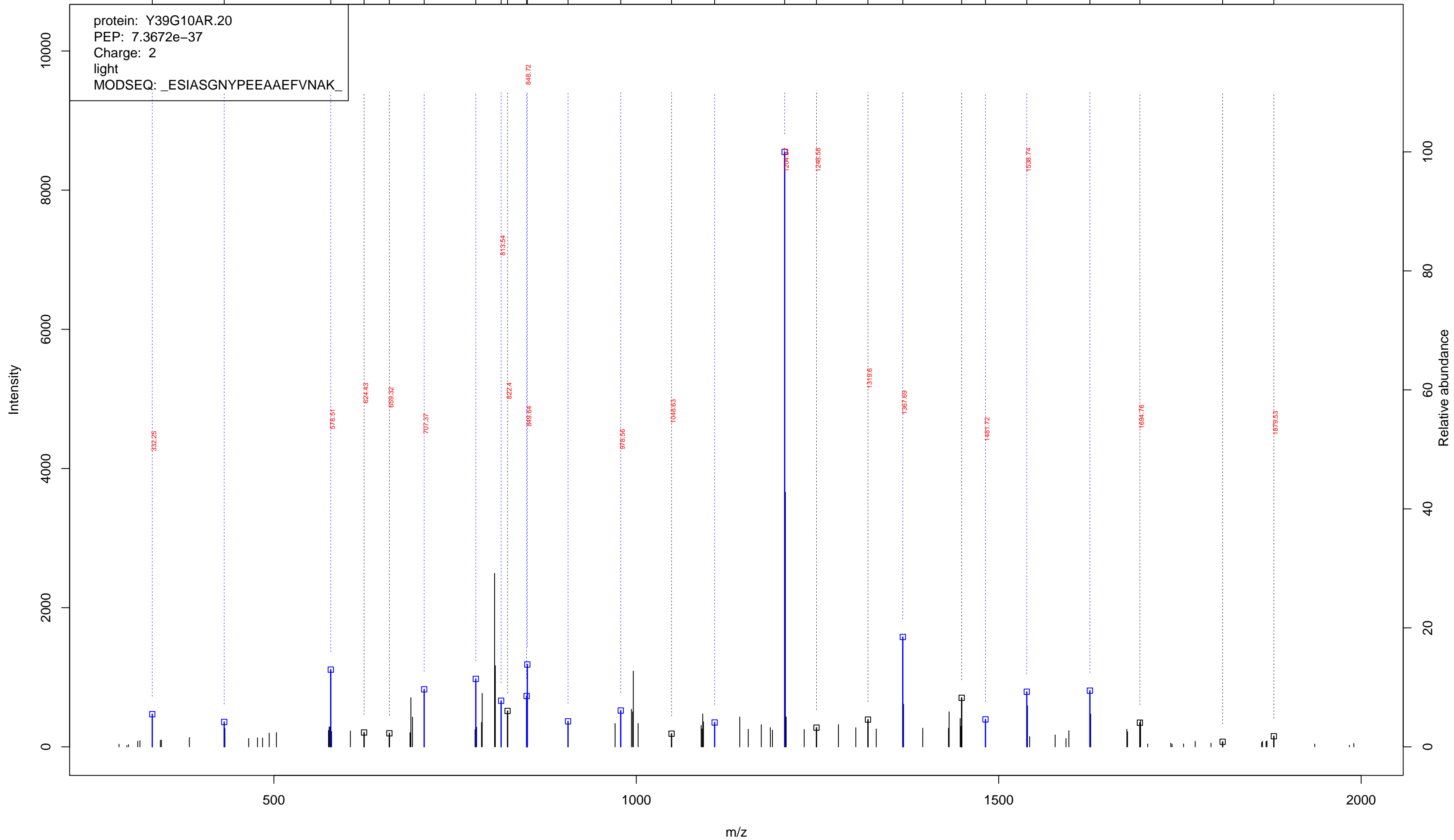






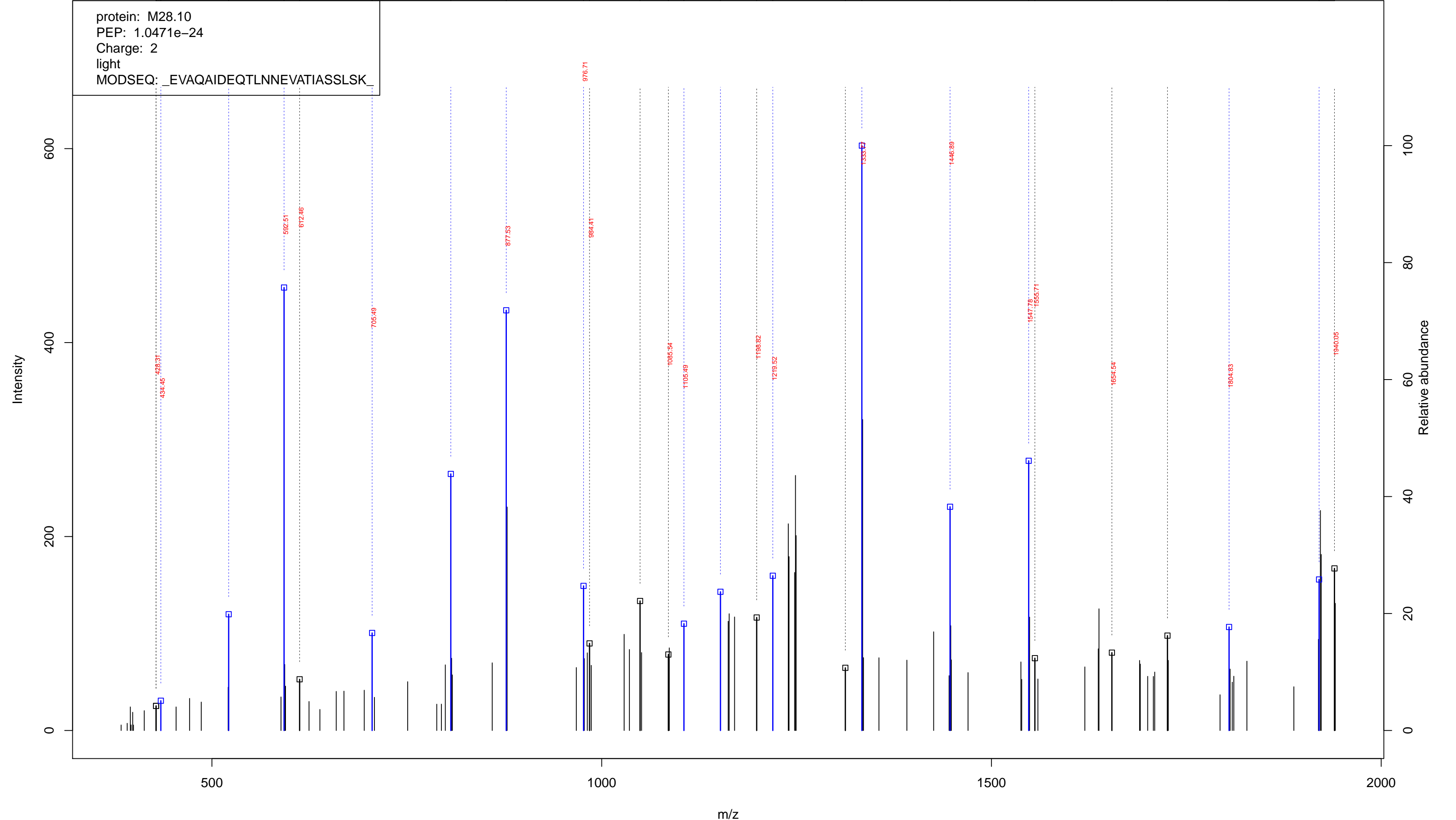


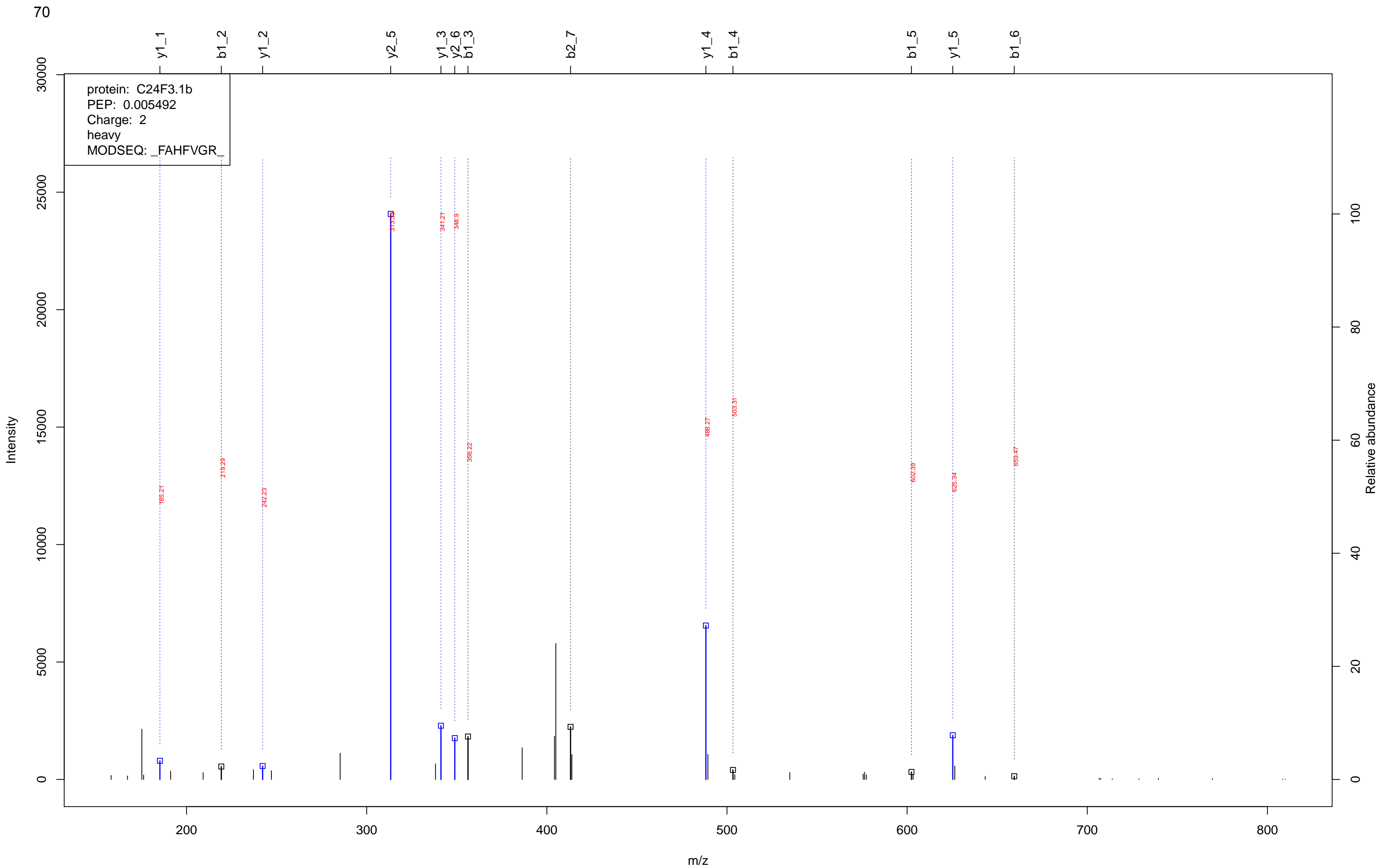


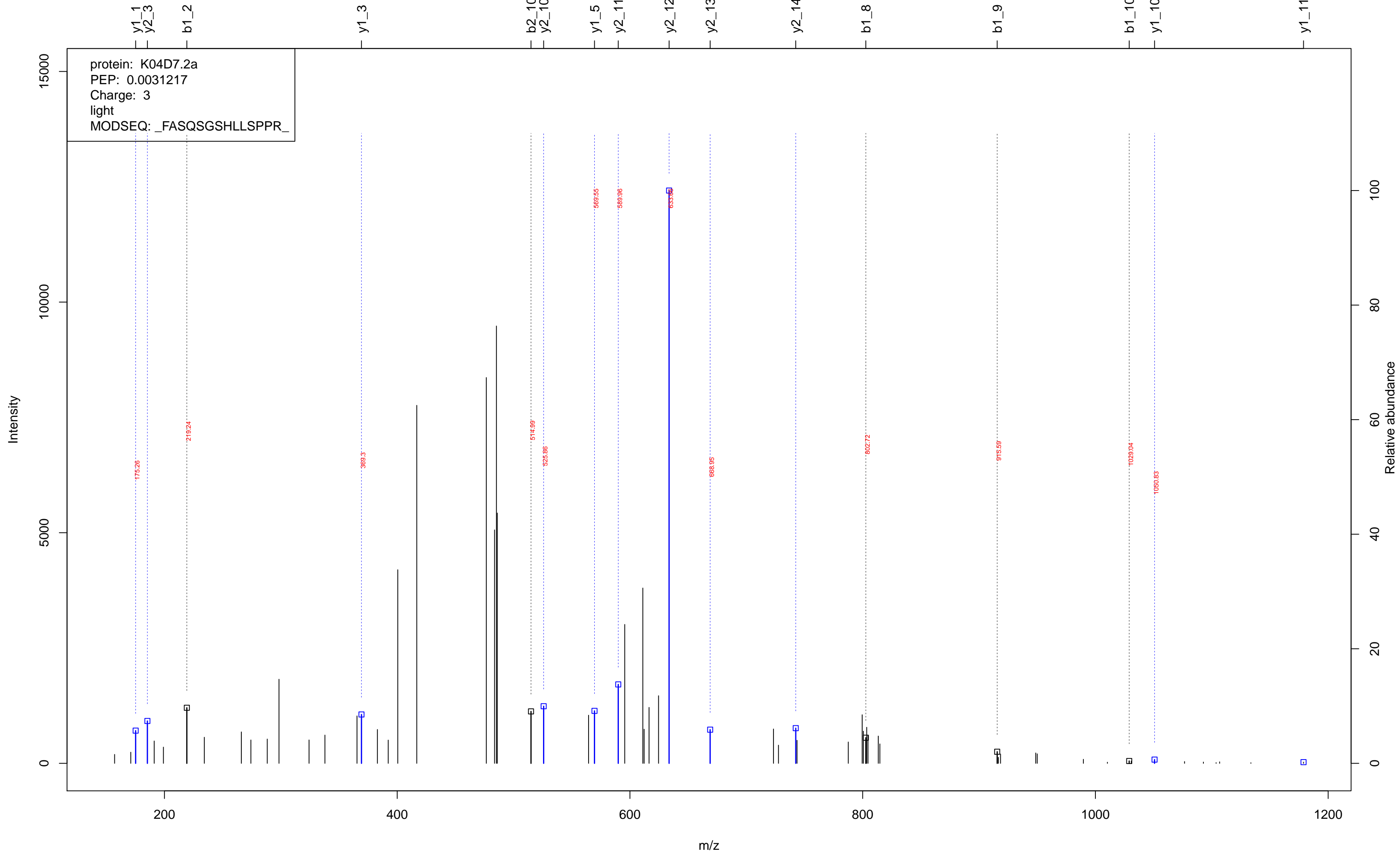


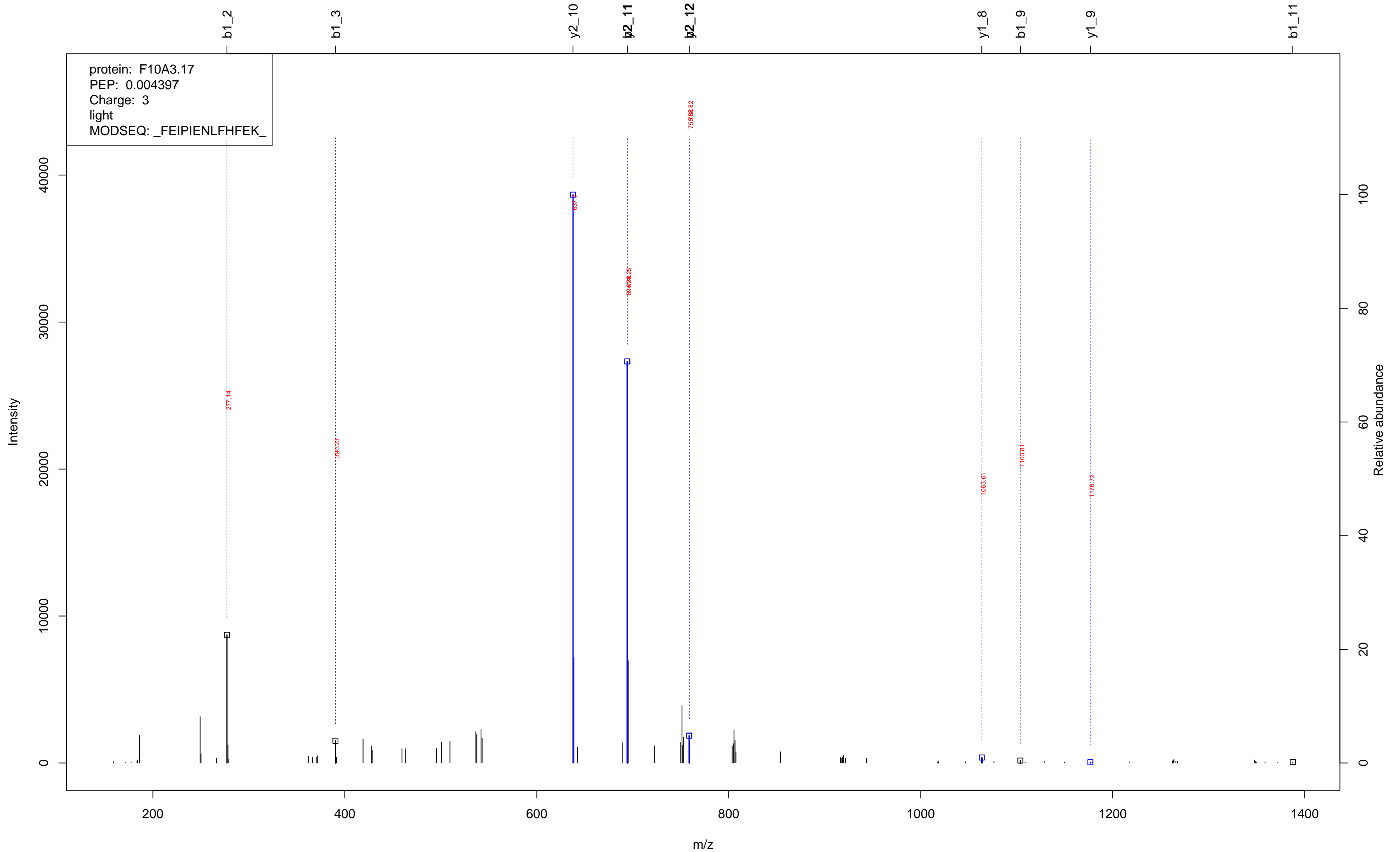
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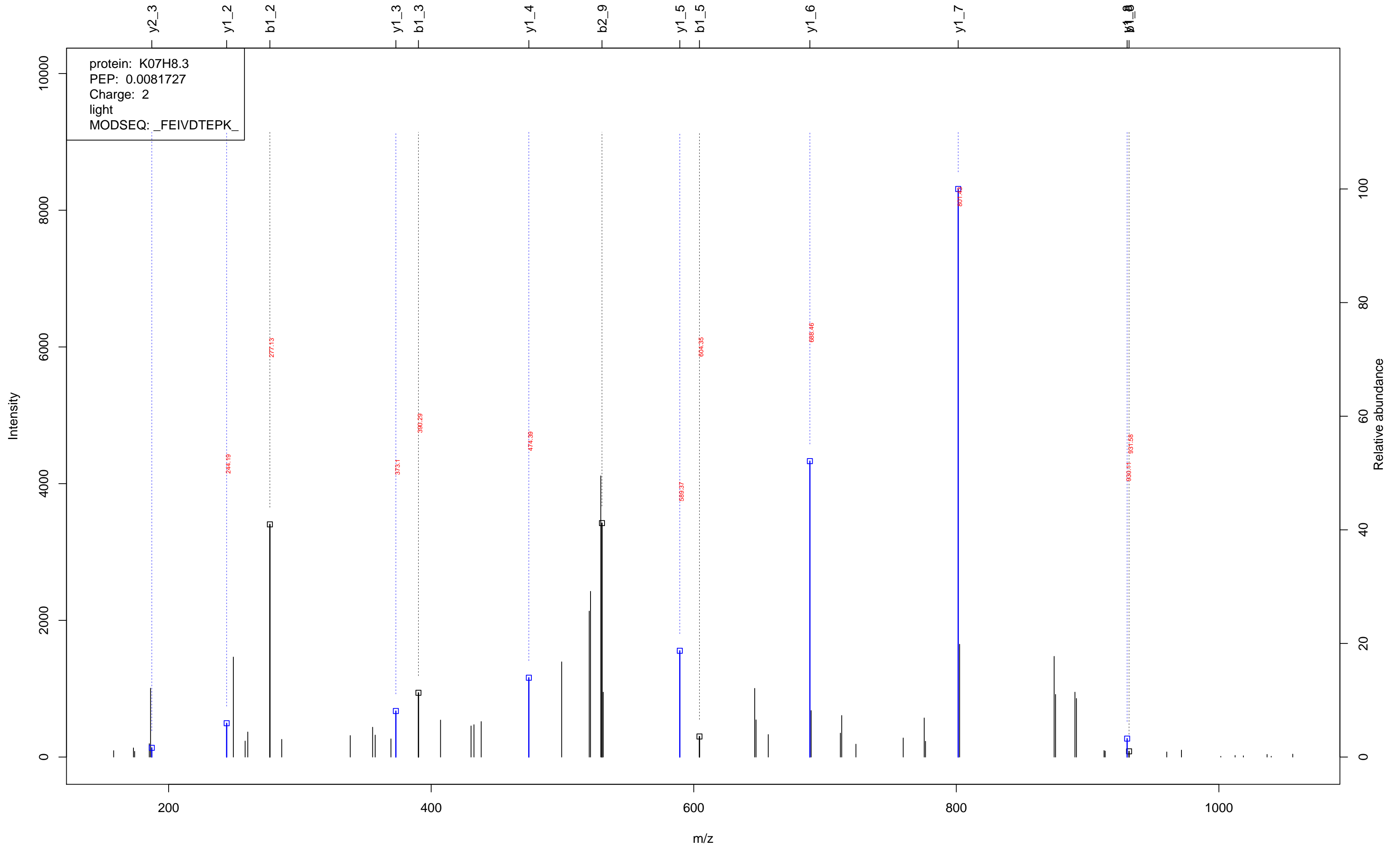
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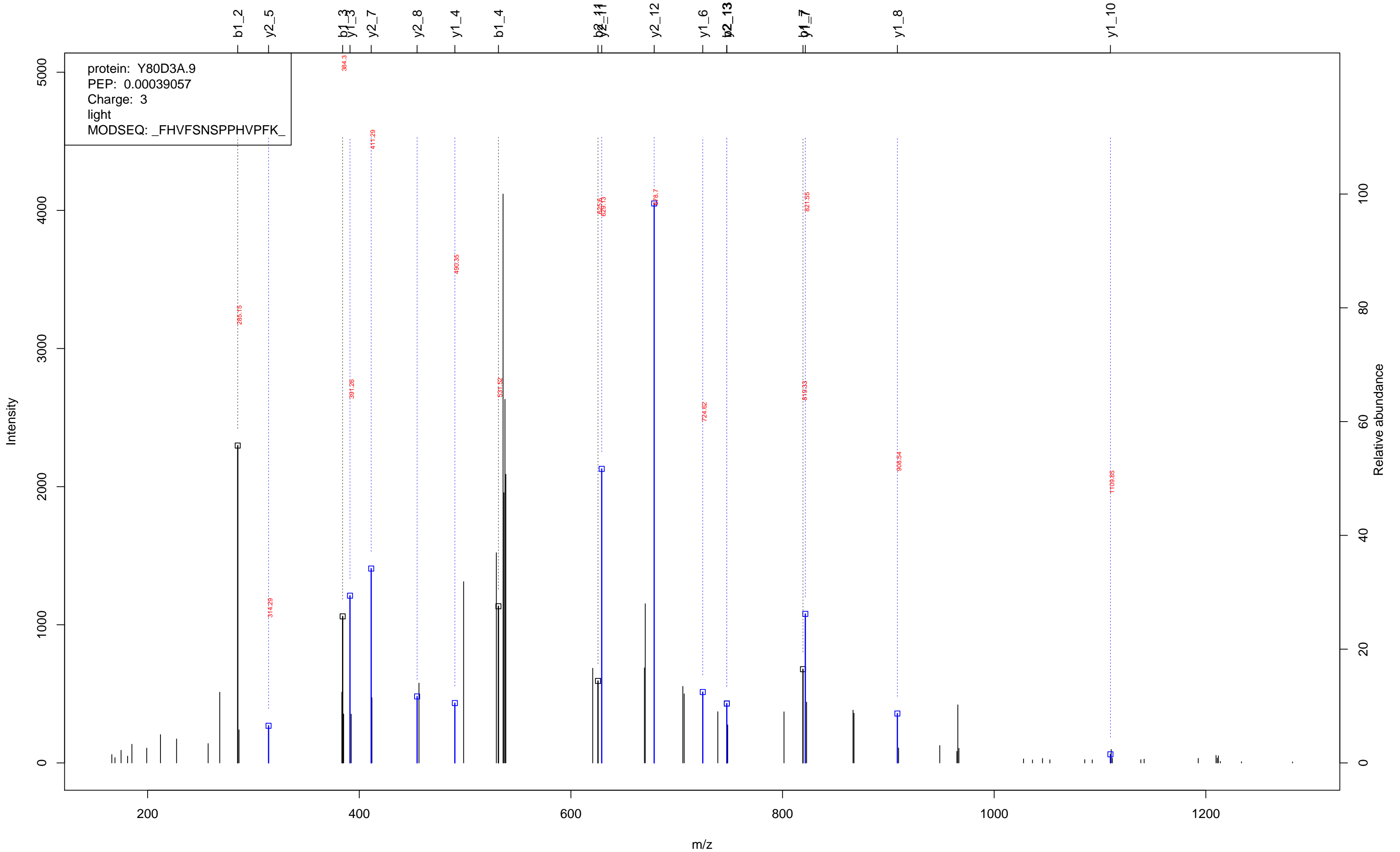


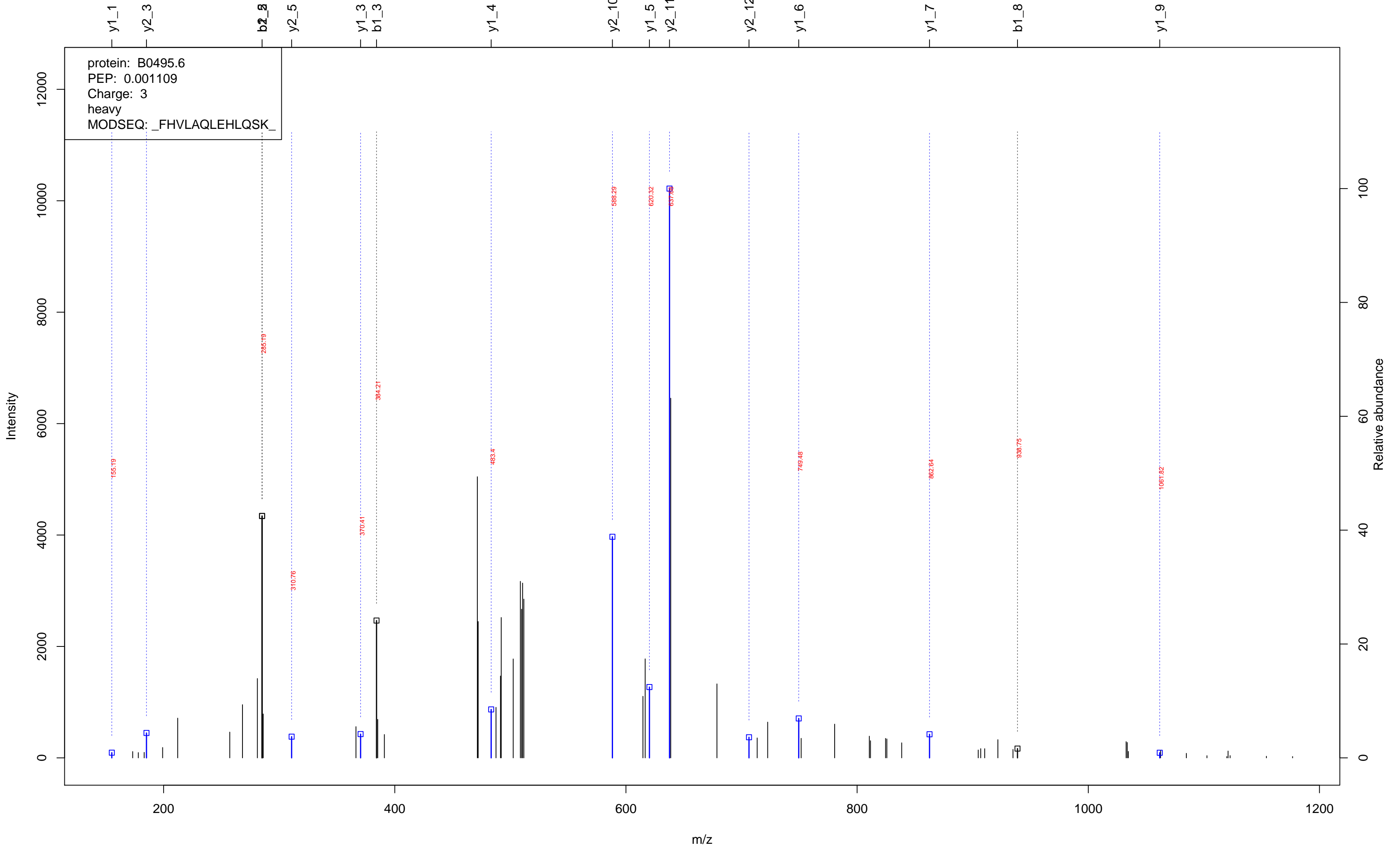


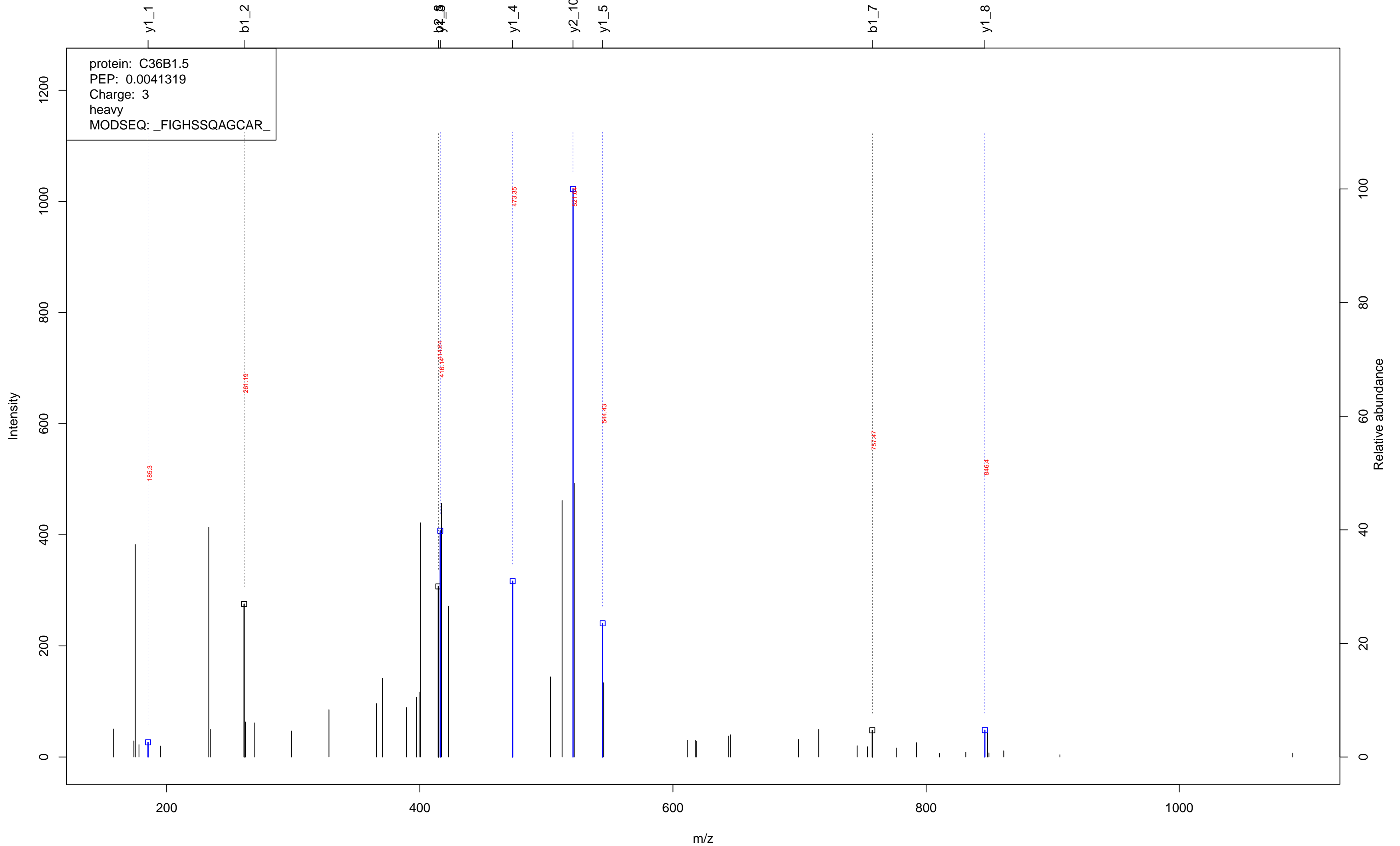


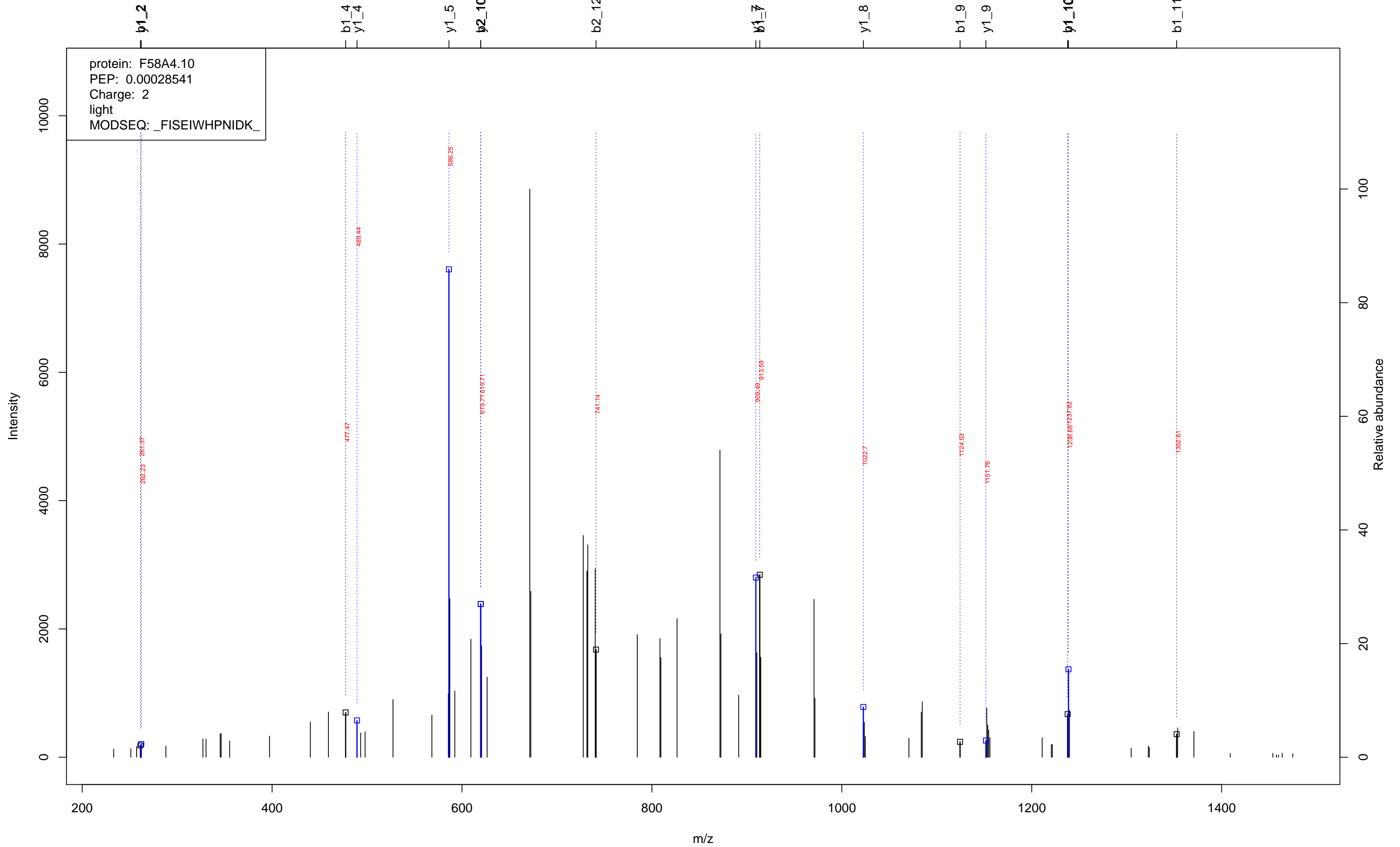












b2_3

b1_2

y1_2

b1_3

b2_8

y1_4

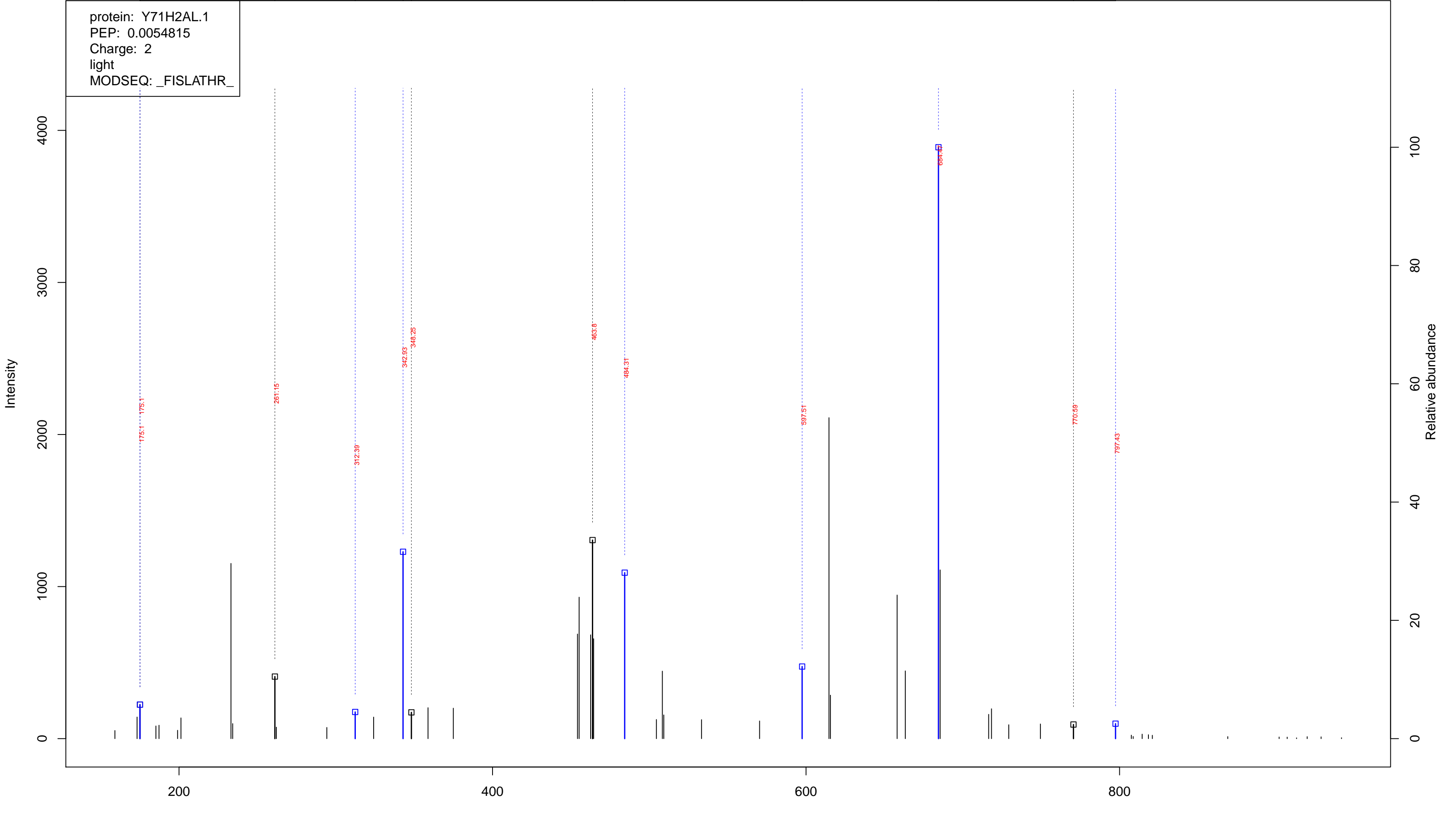
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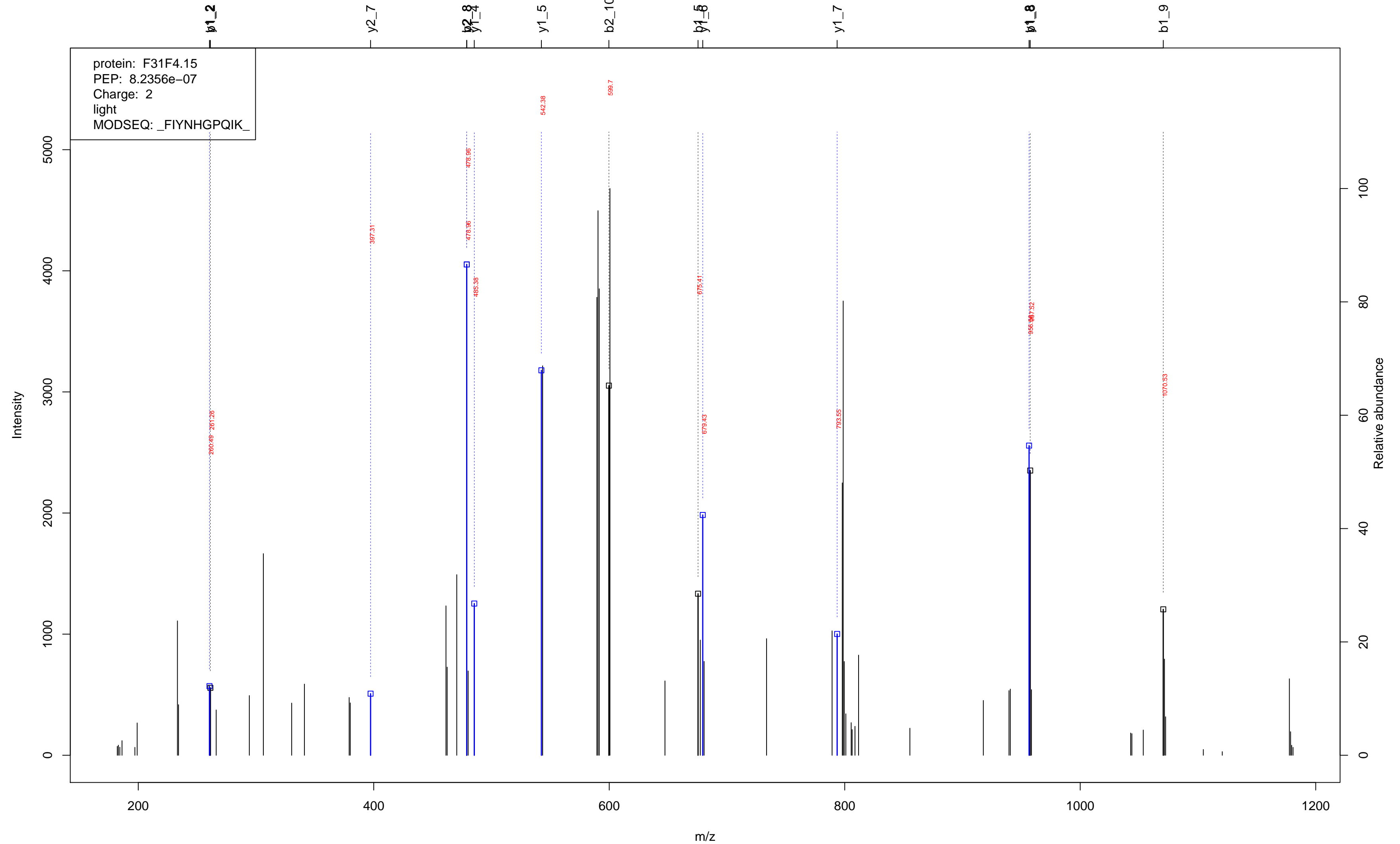
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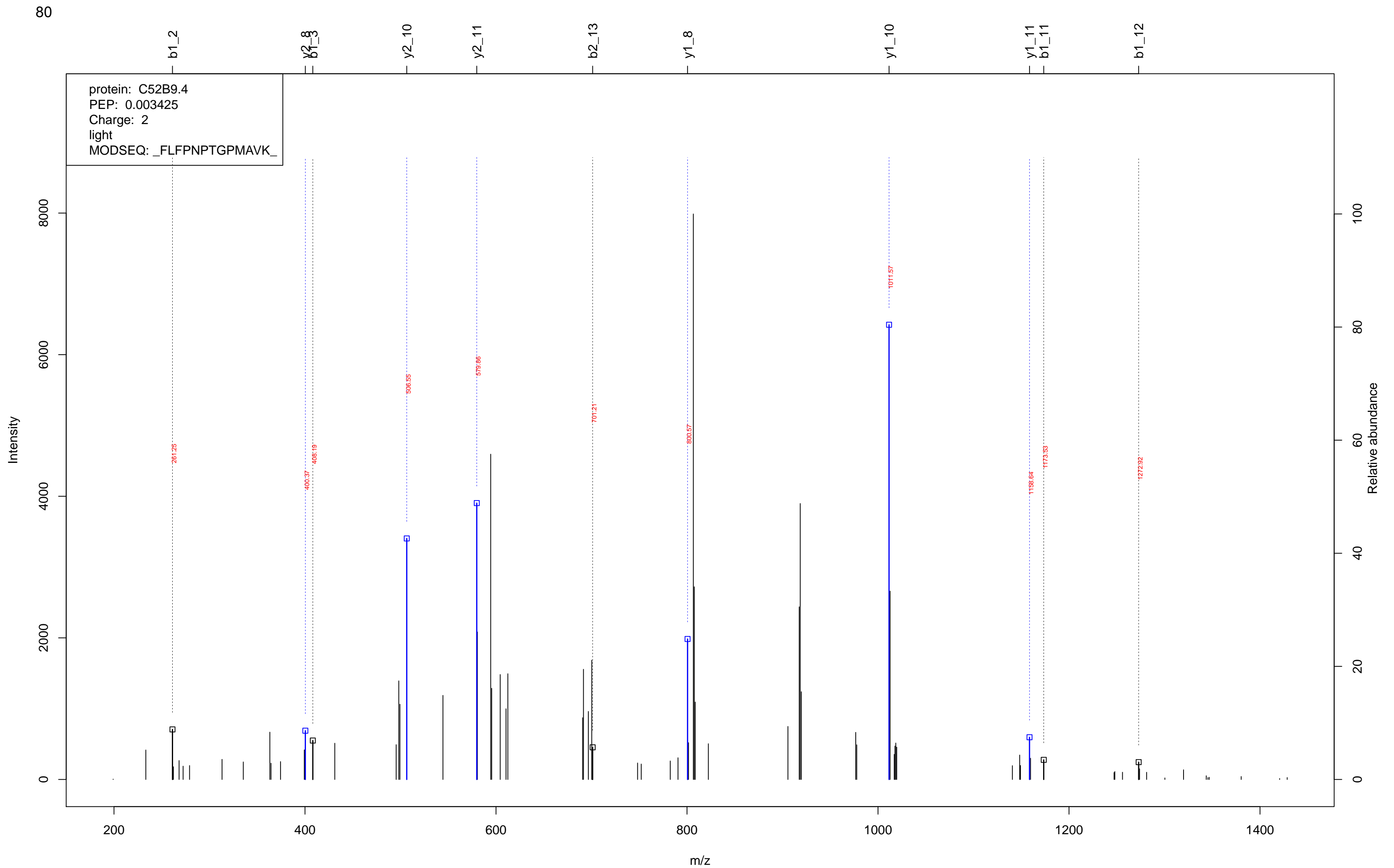
b1_7

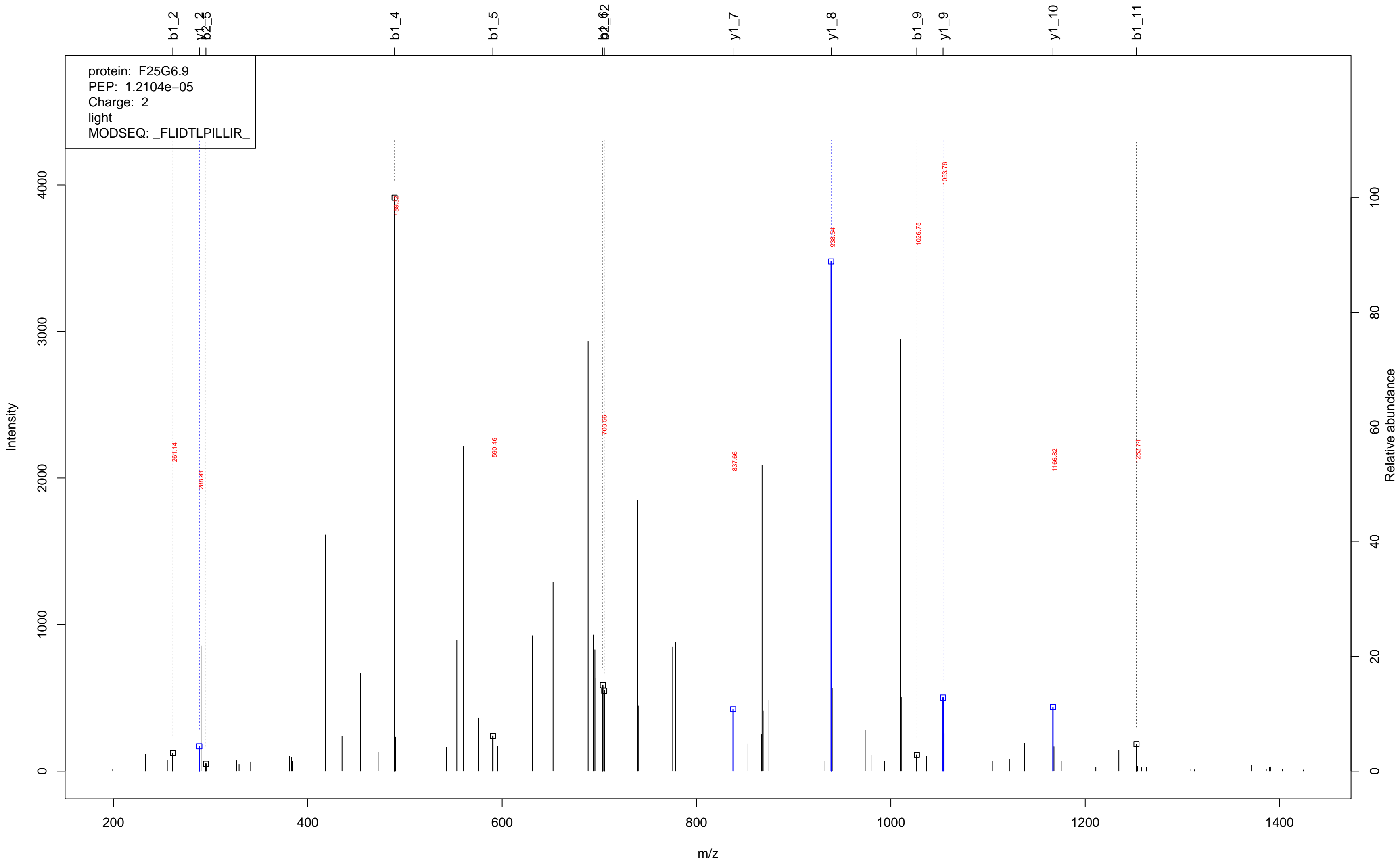
y1_7

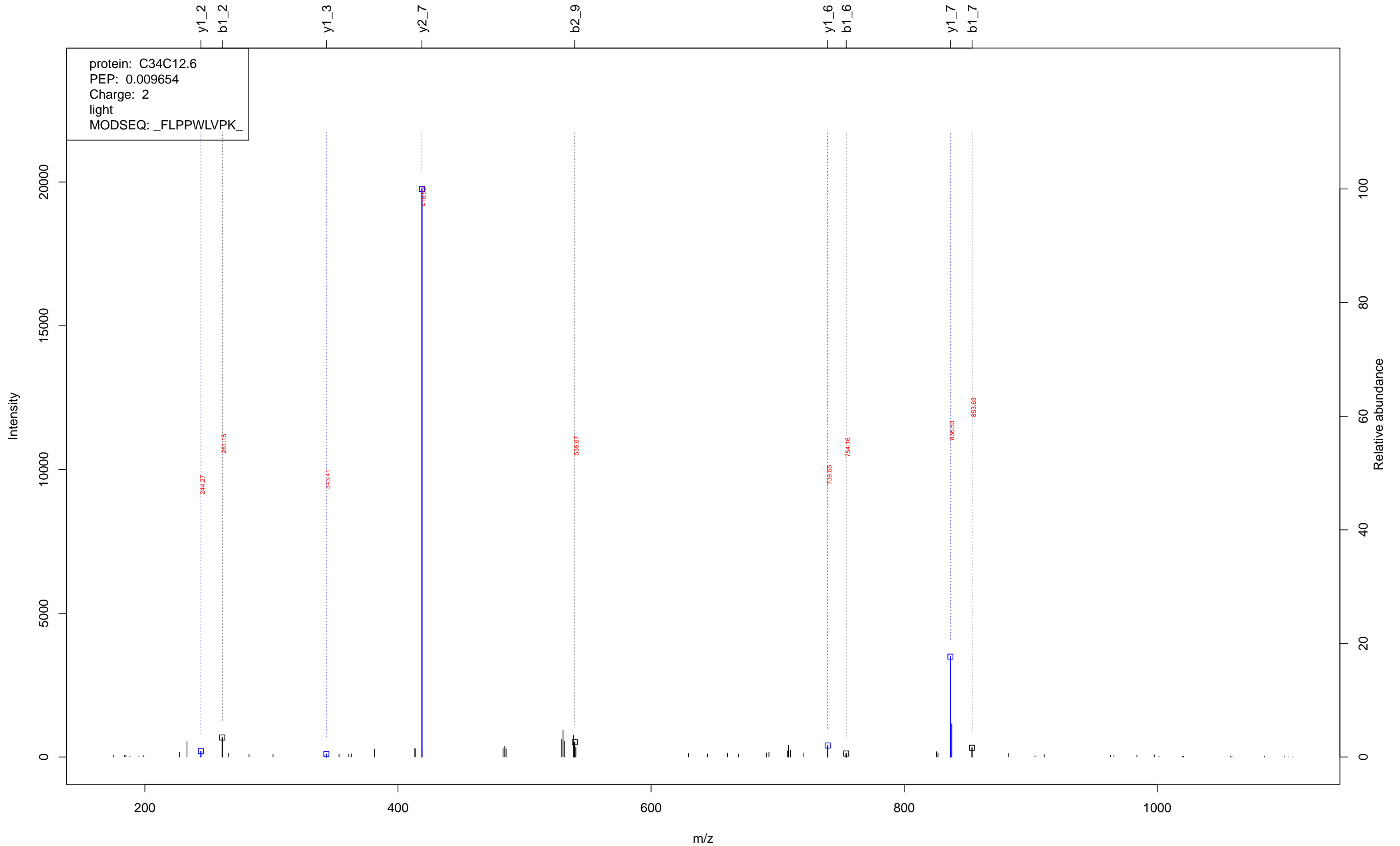
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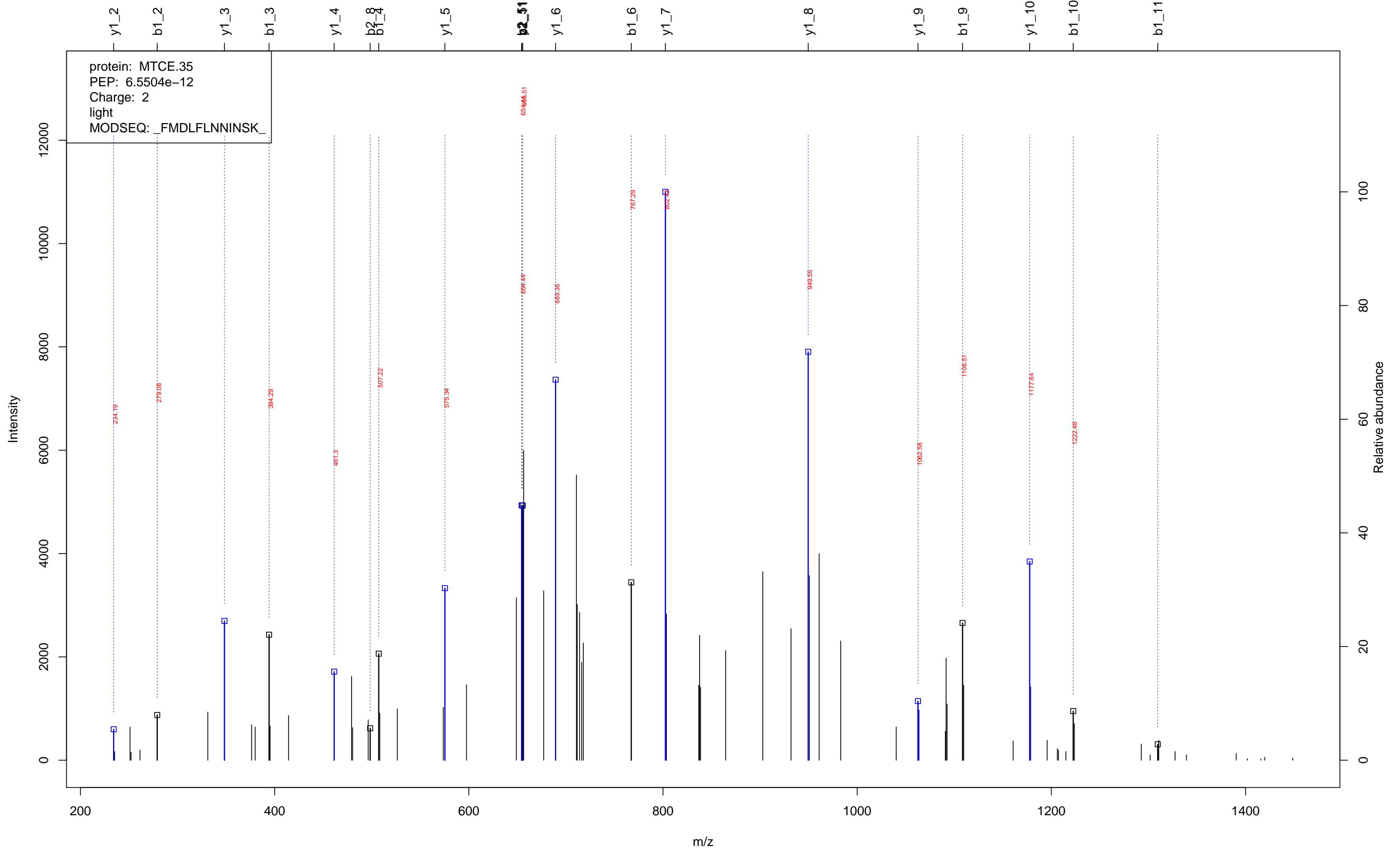


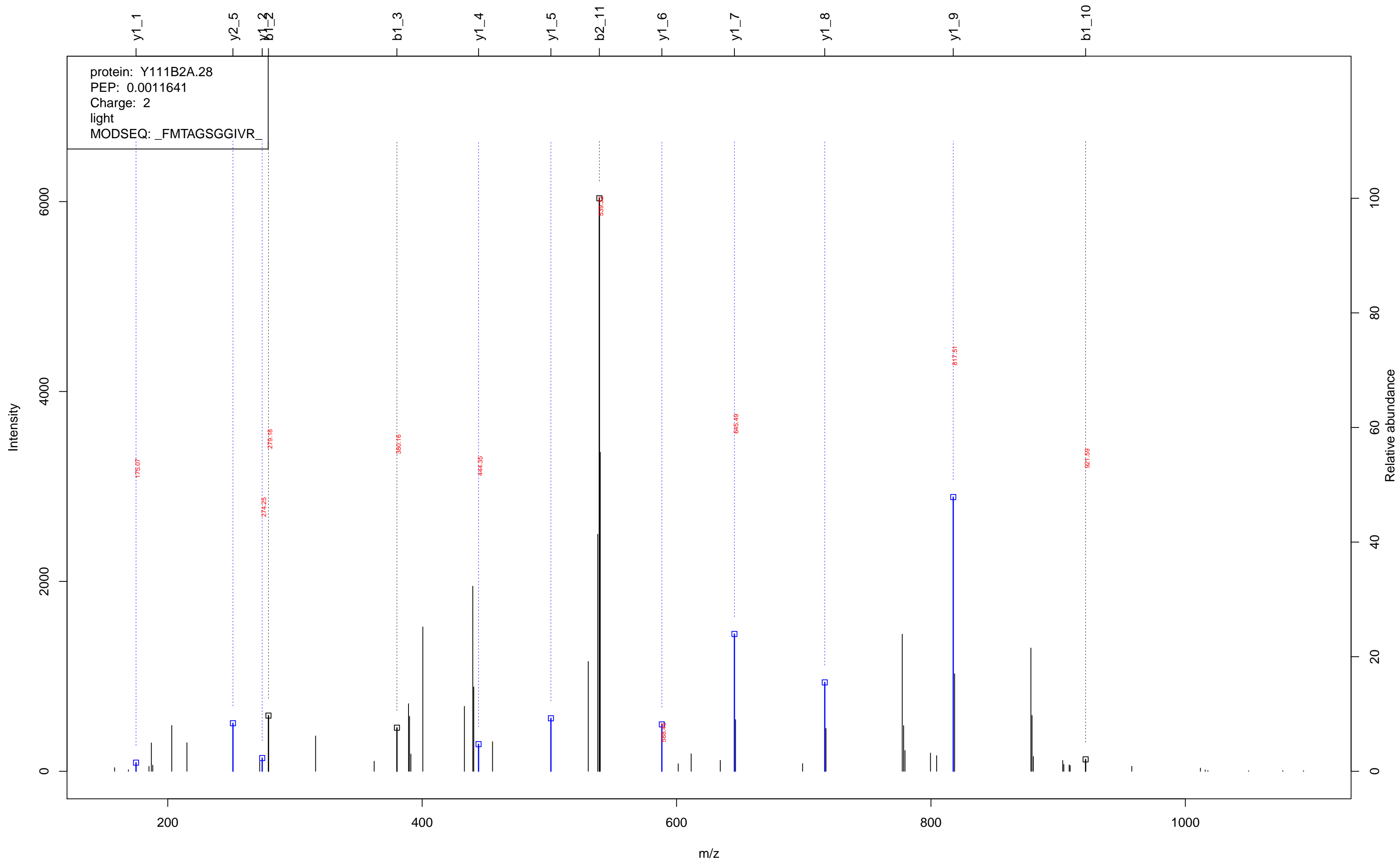


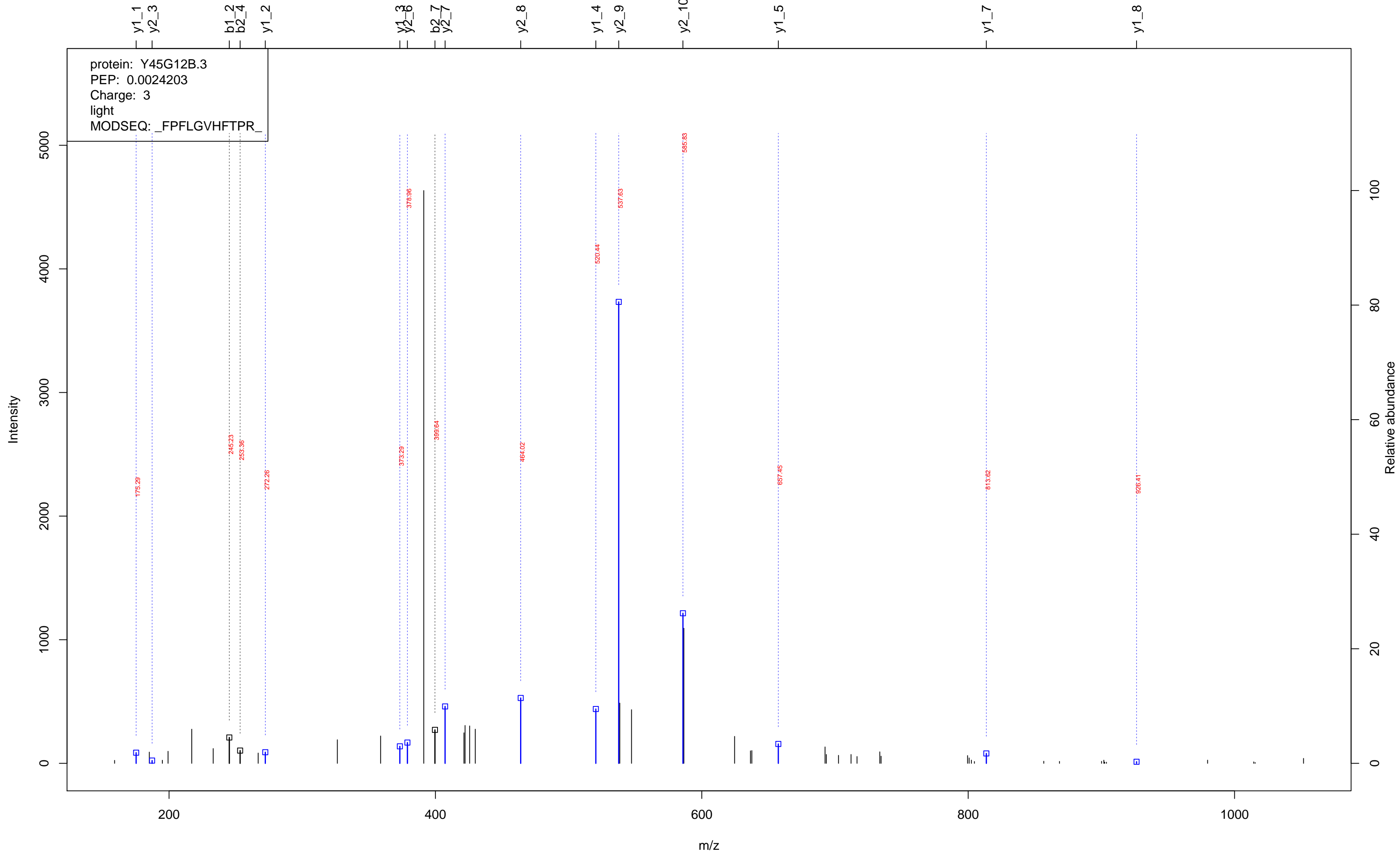


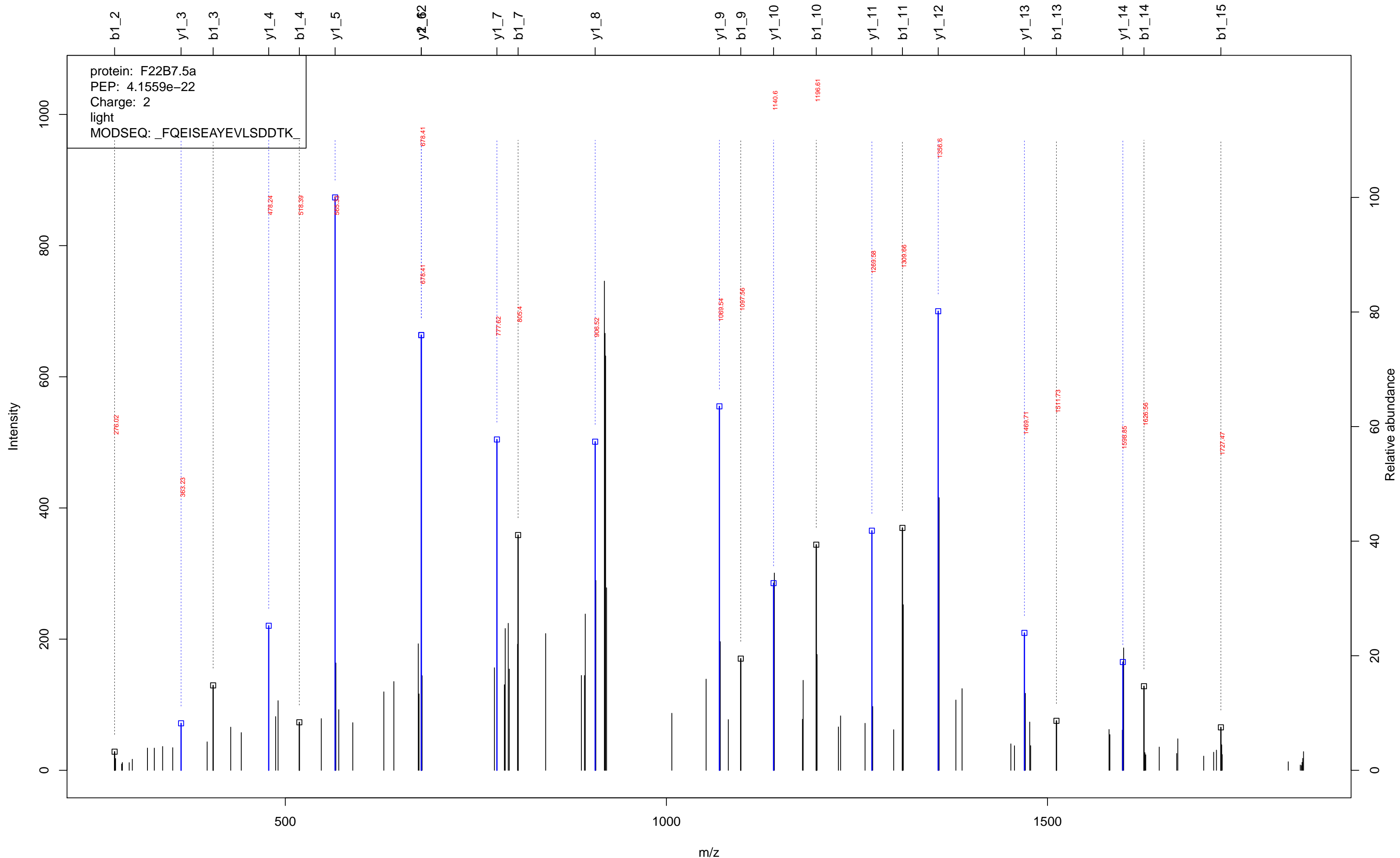




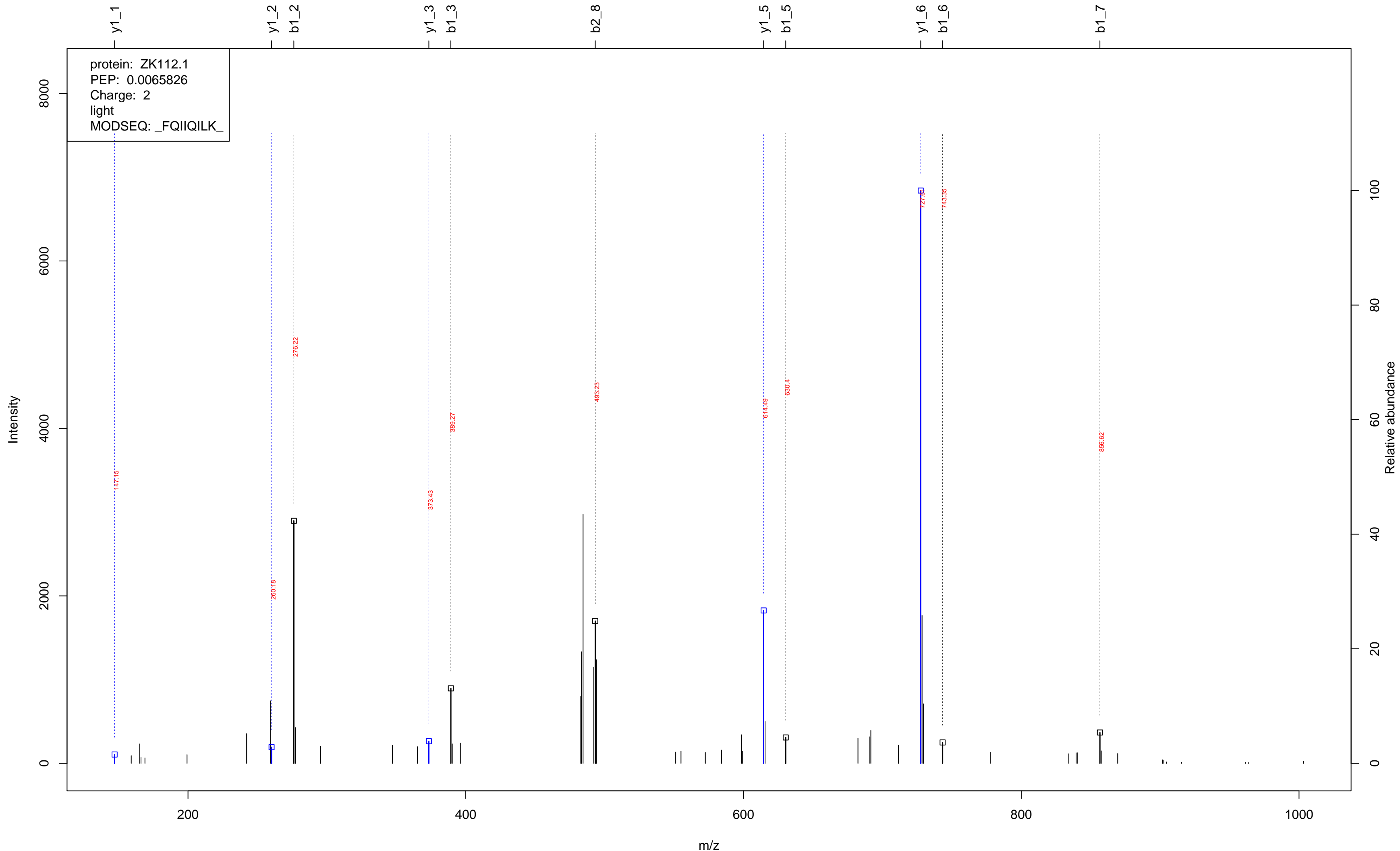


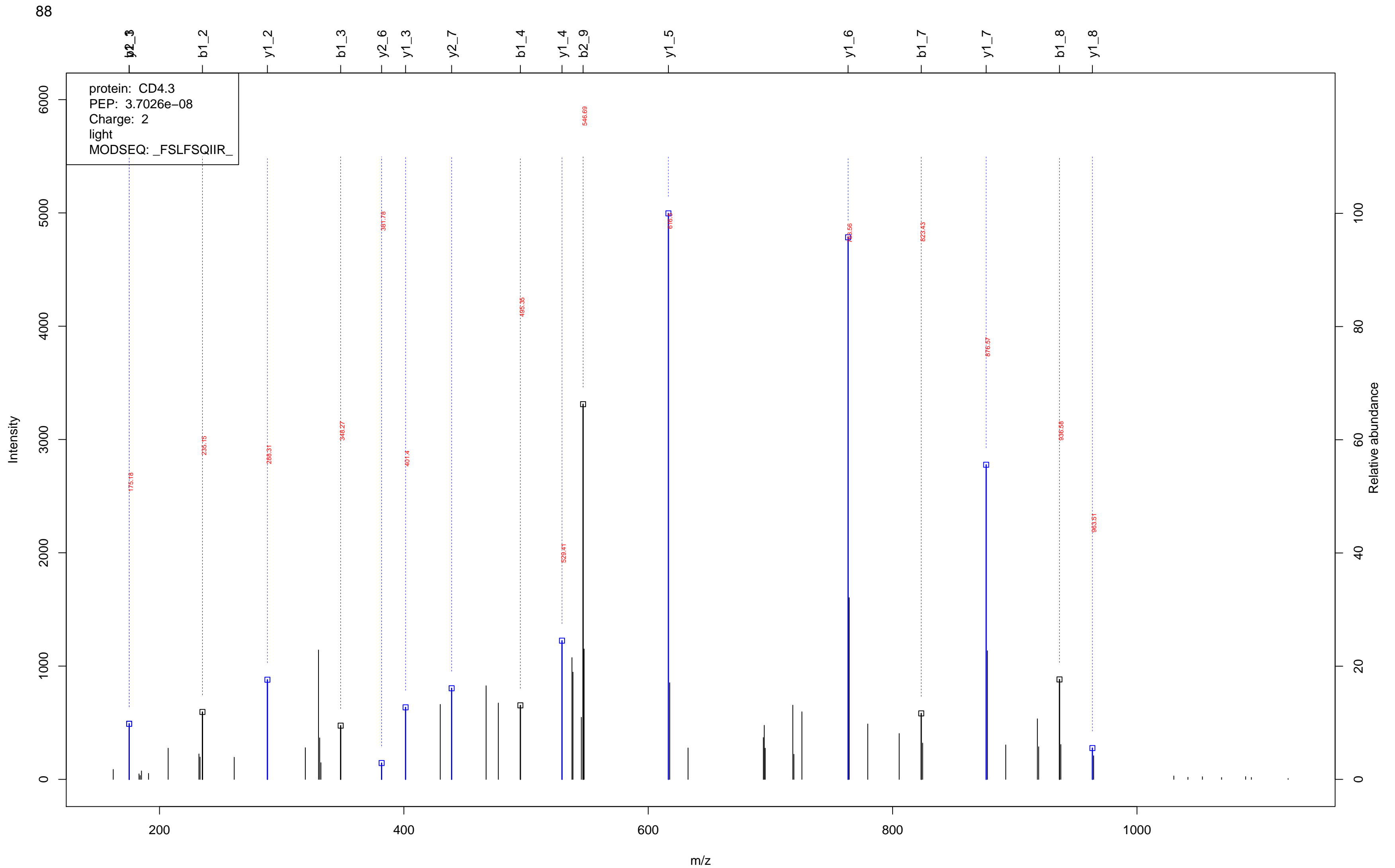


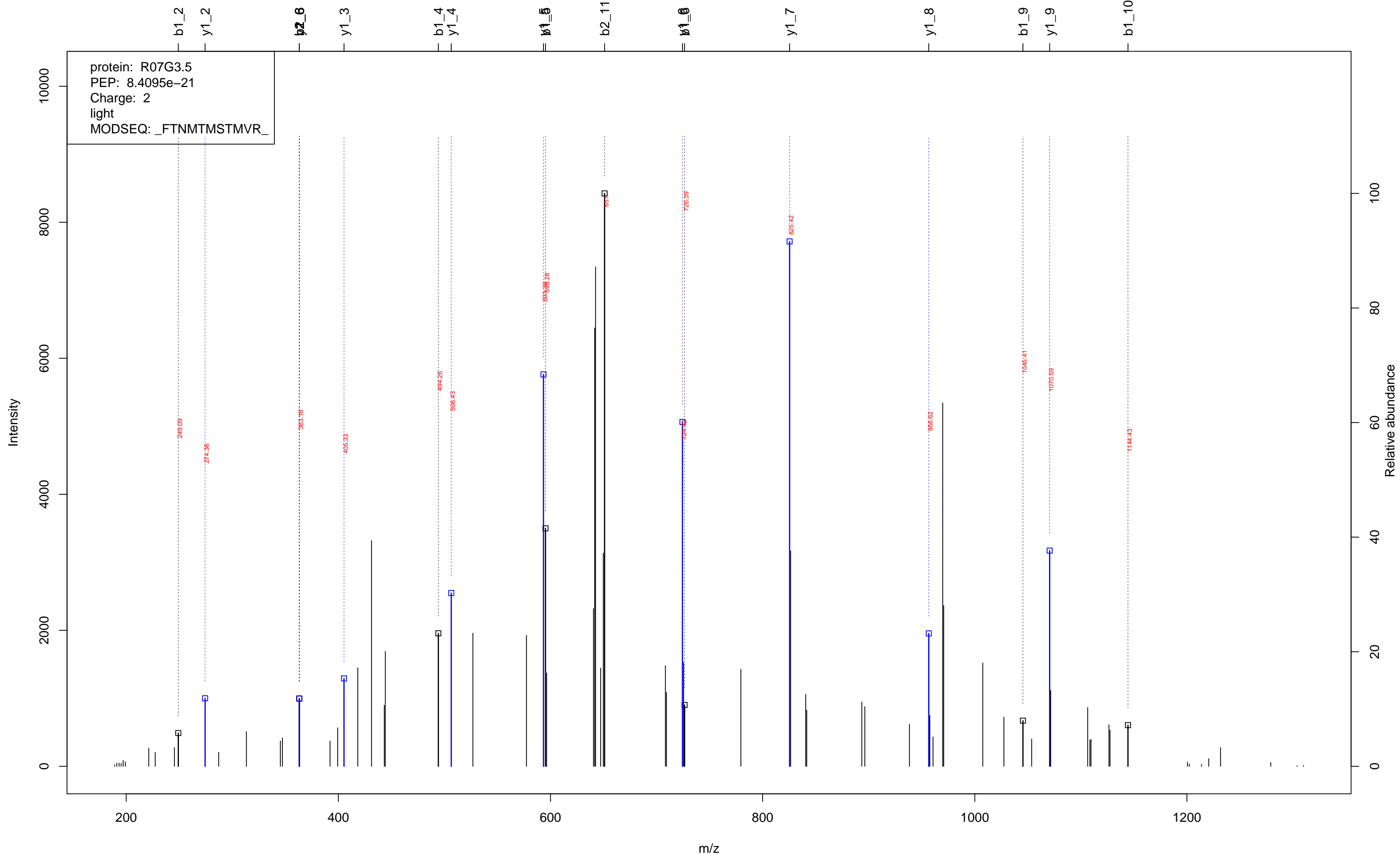


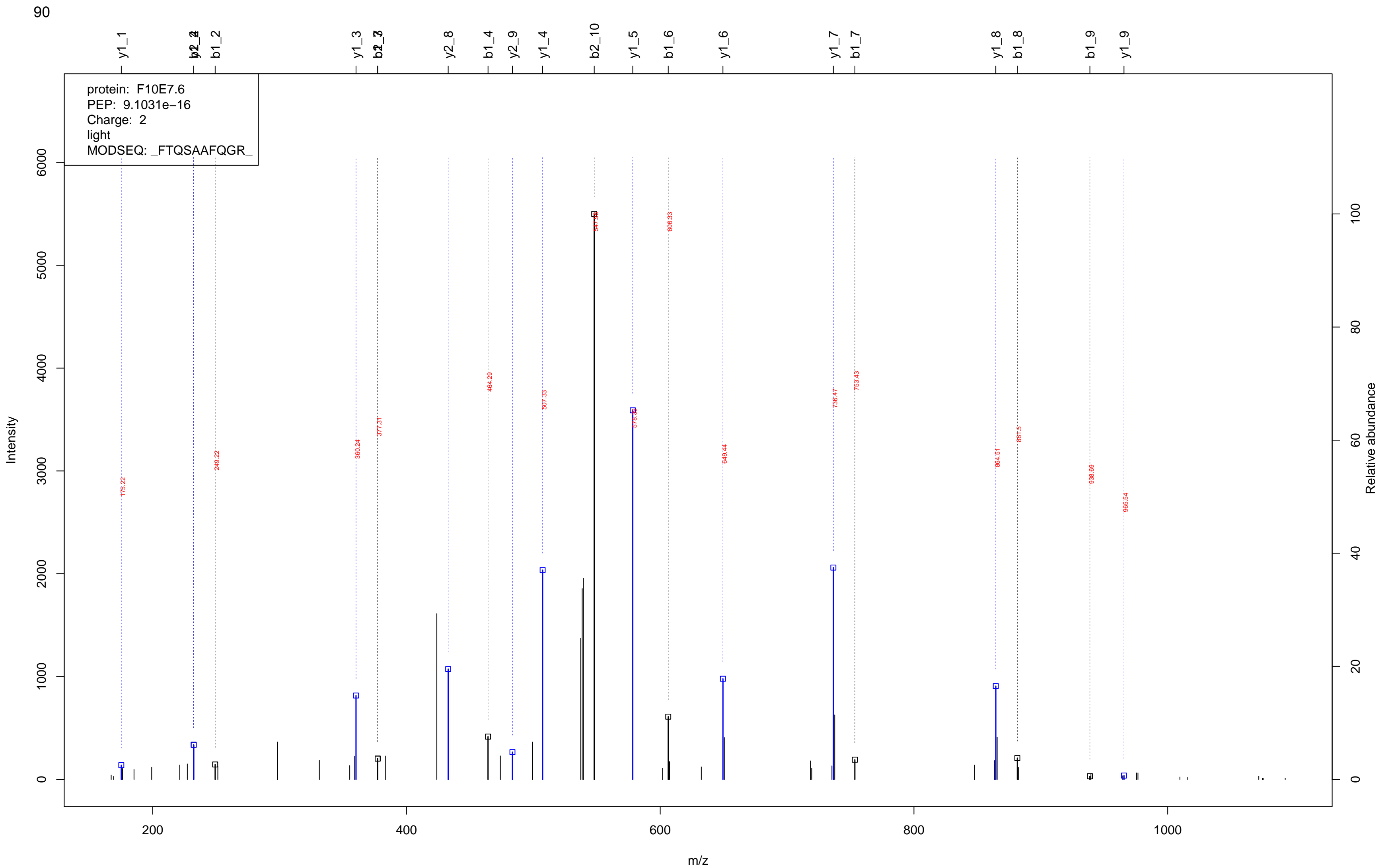


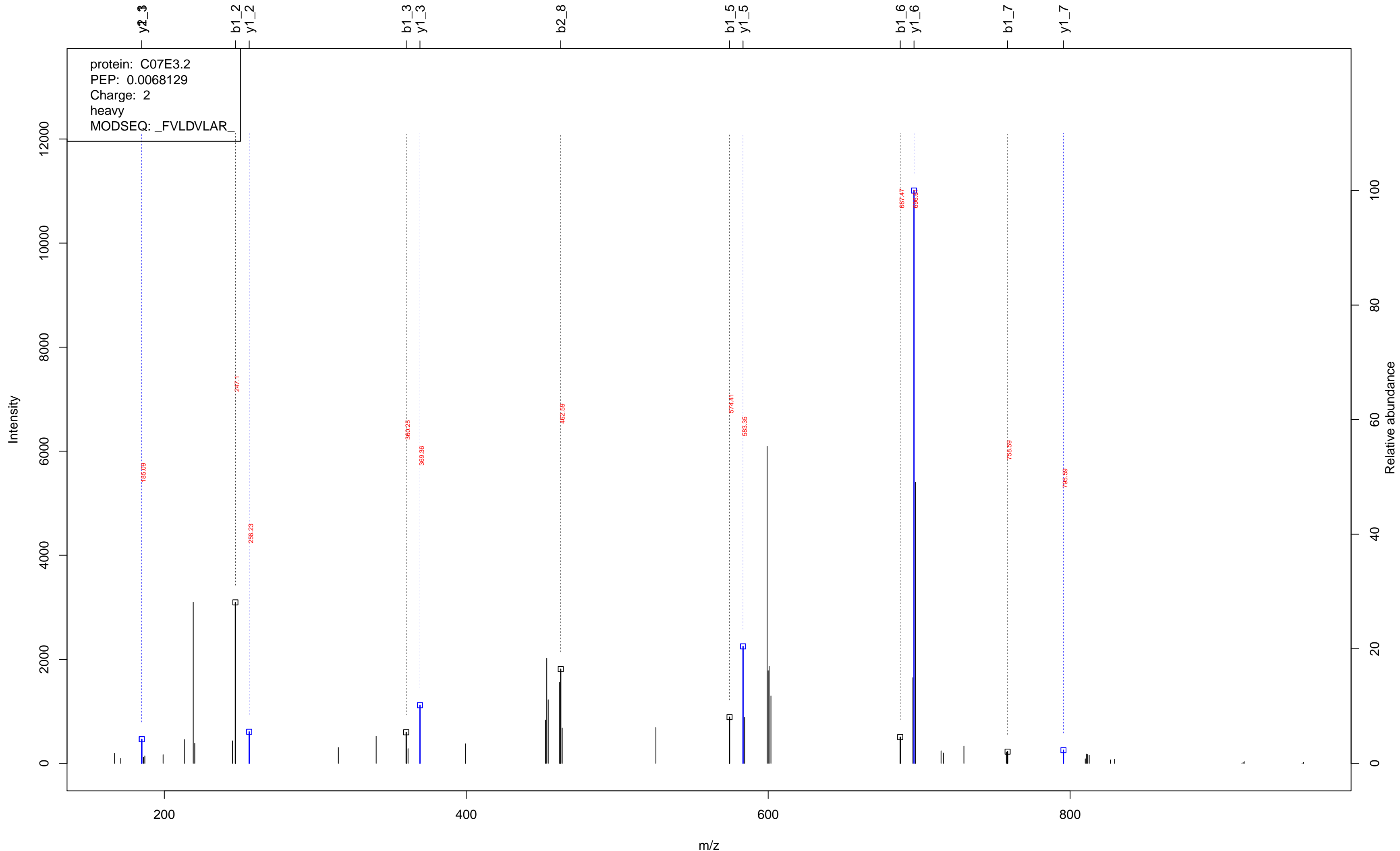
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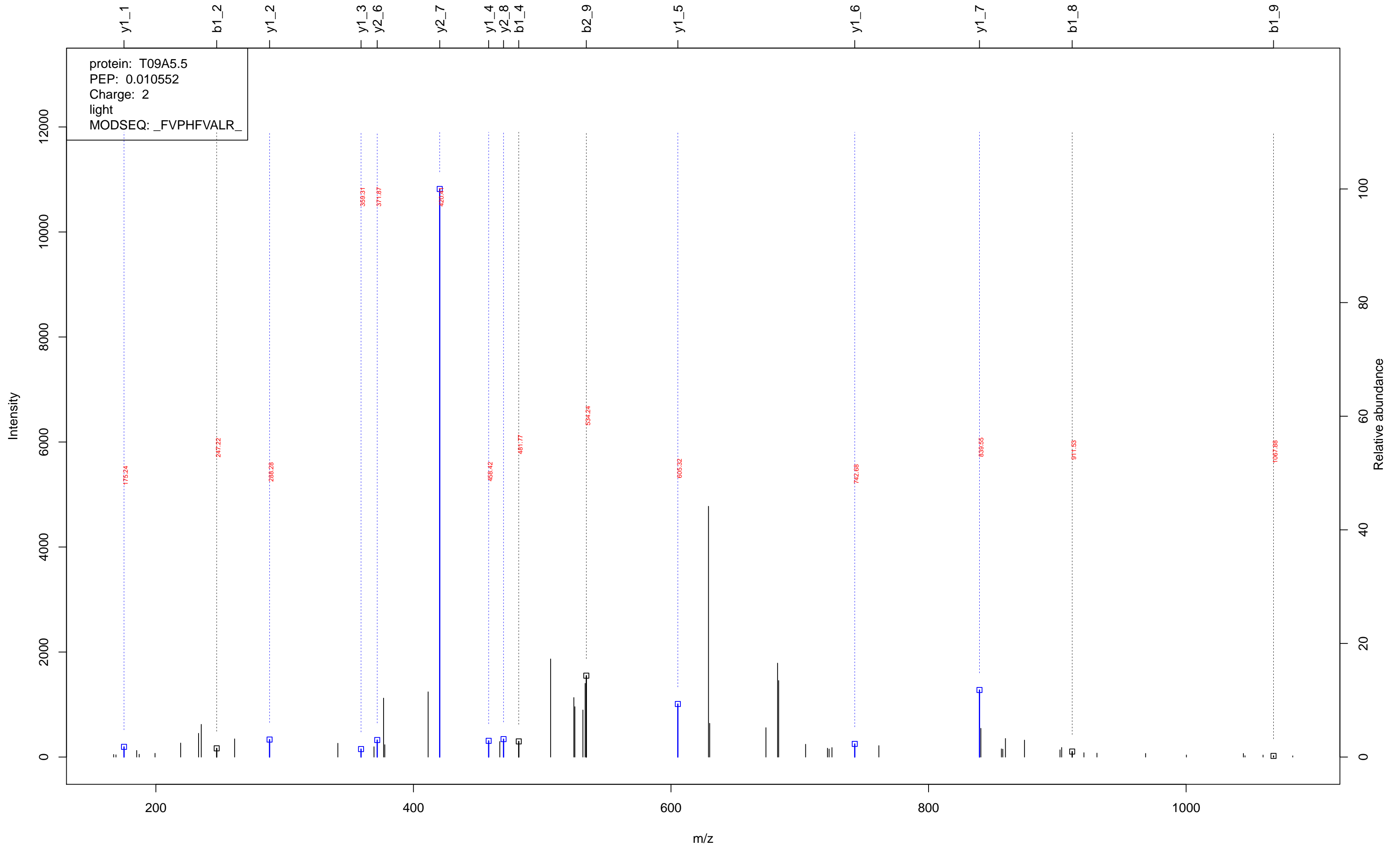


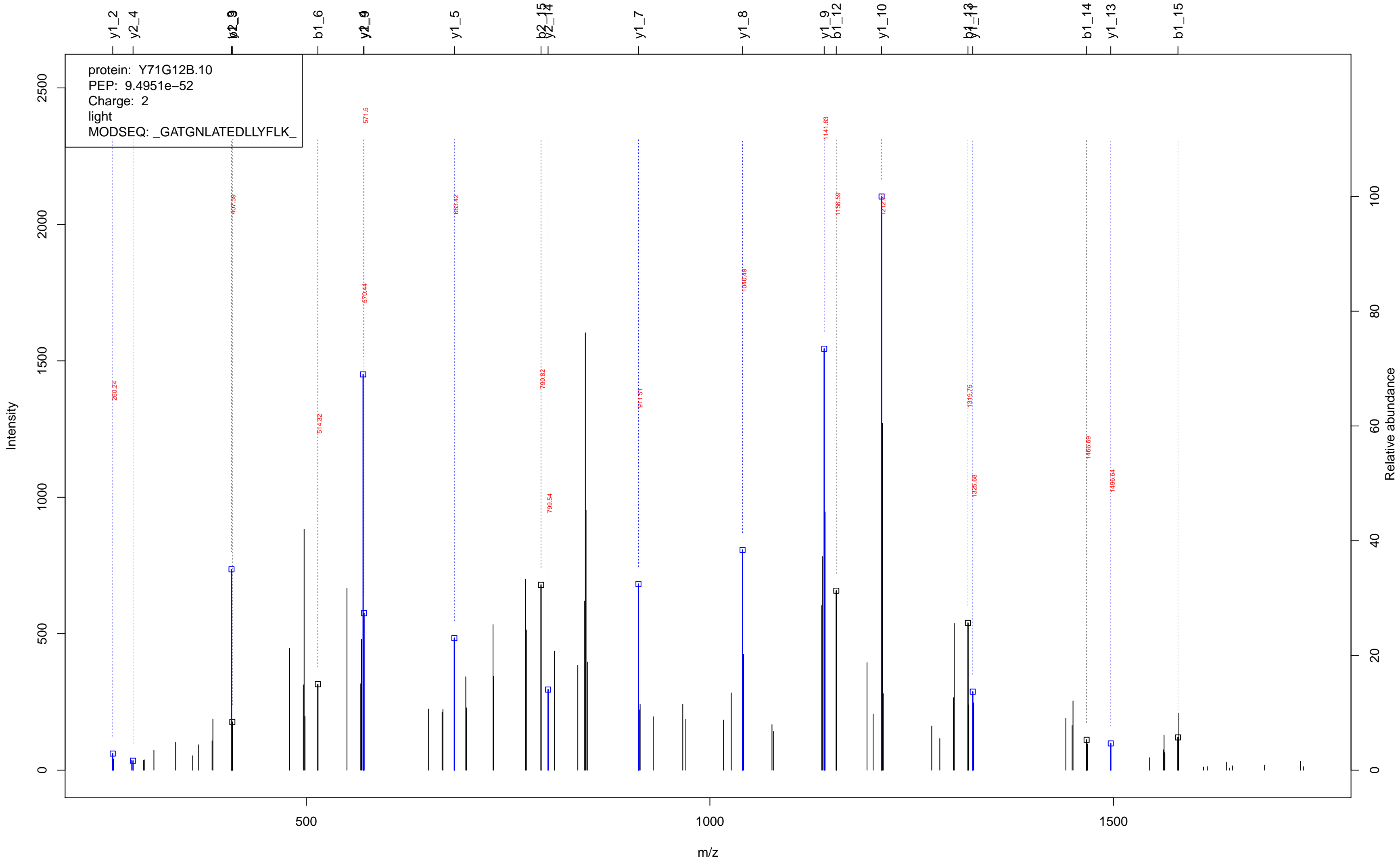


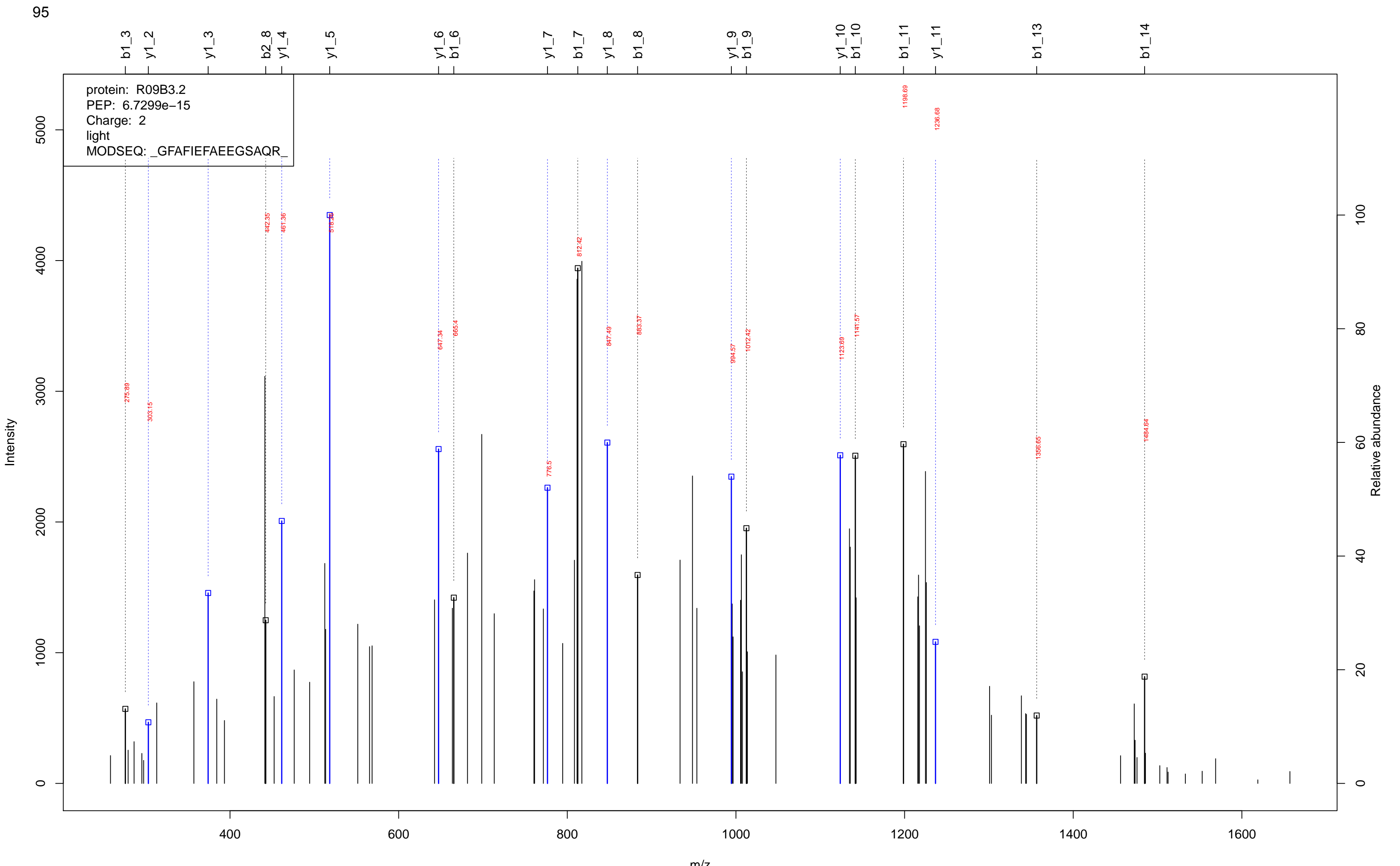


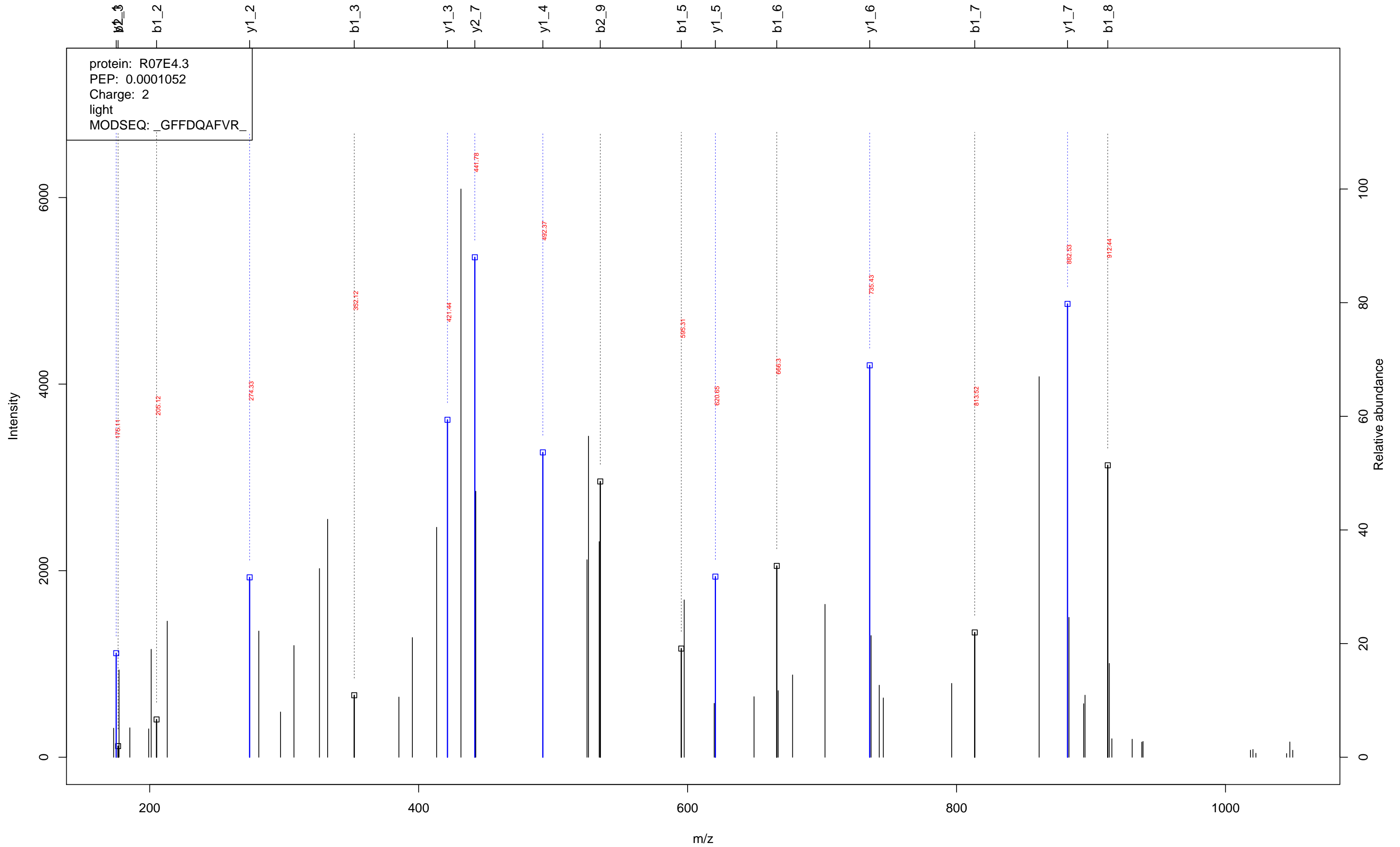


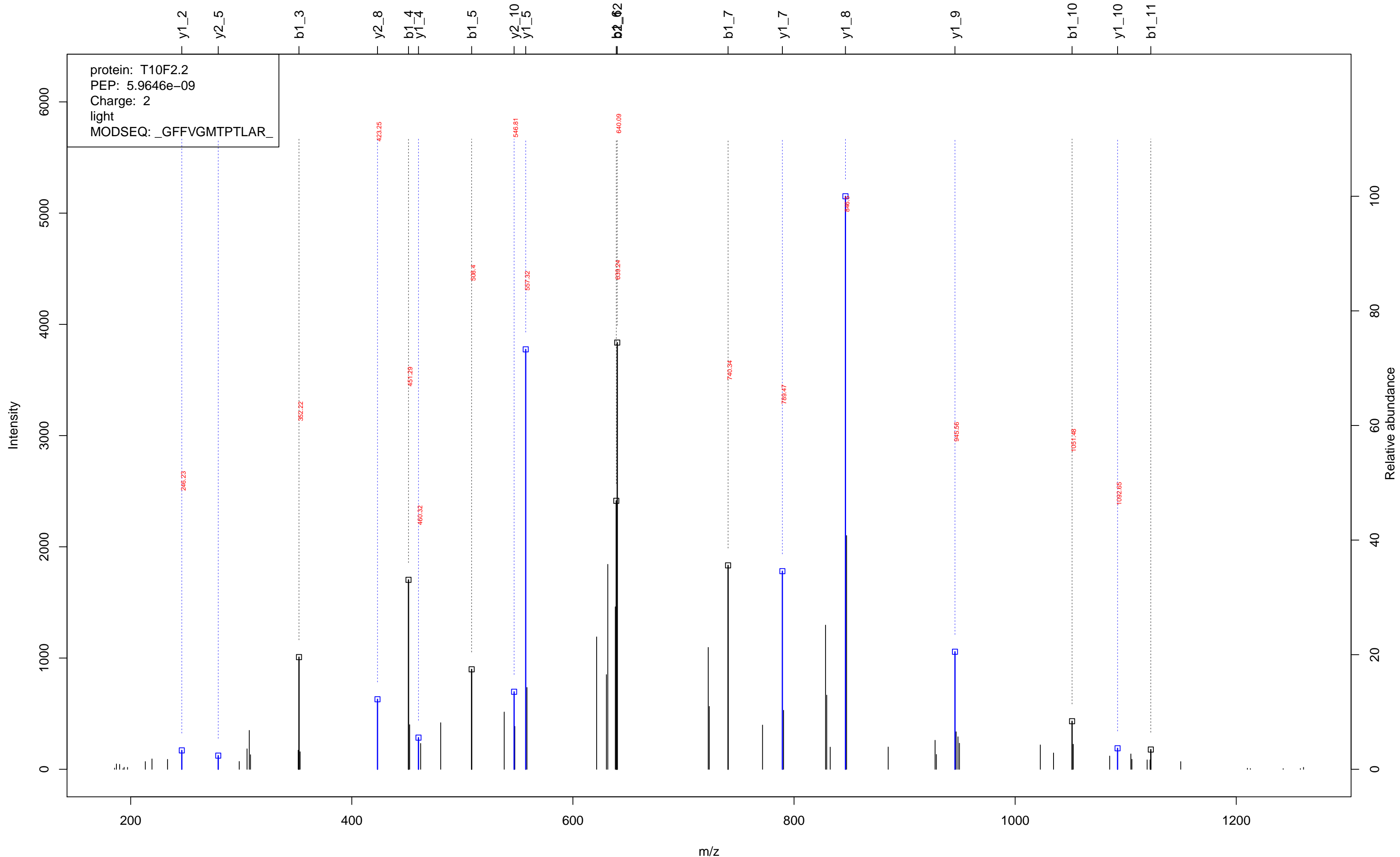


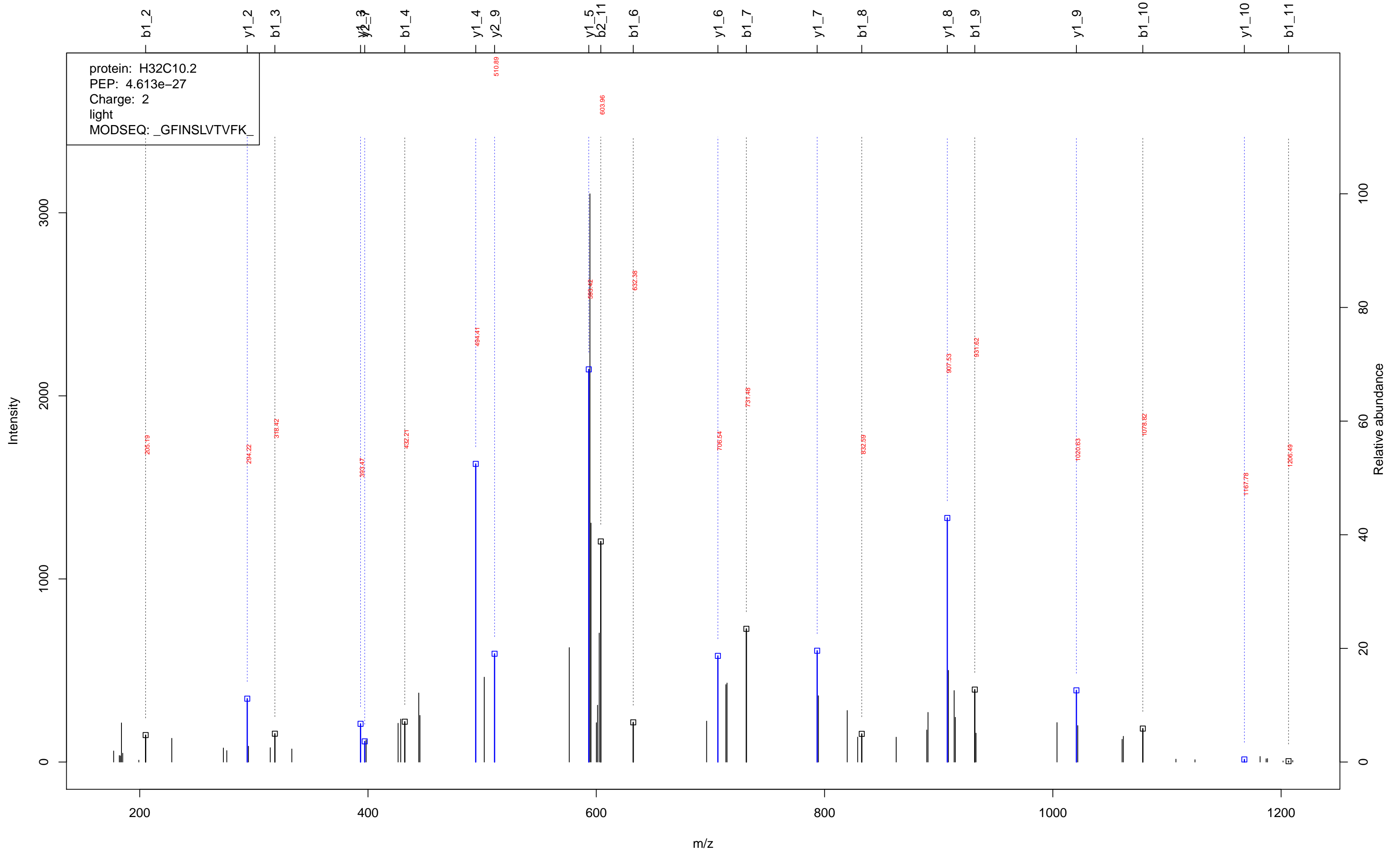


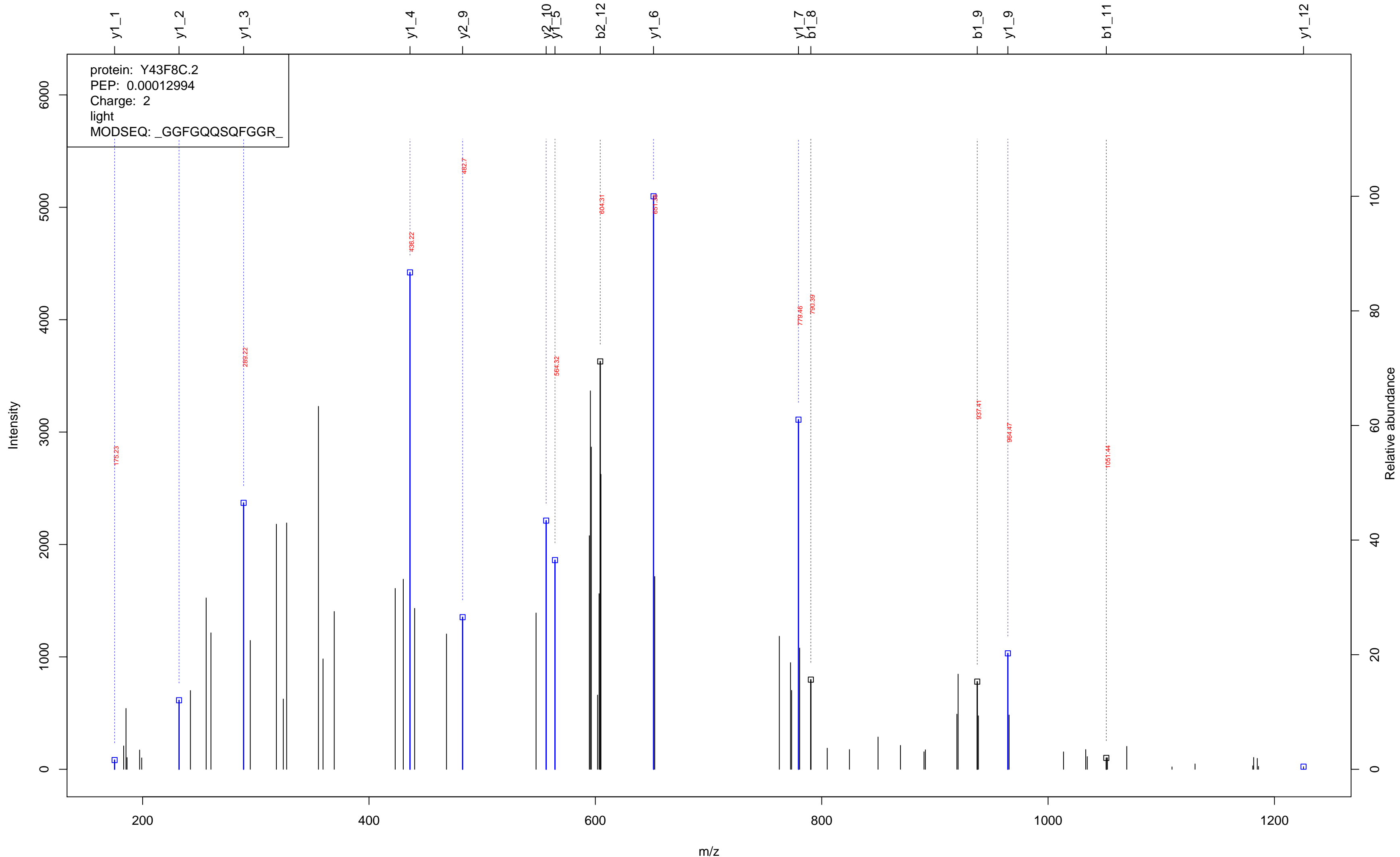


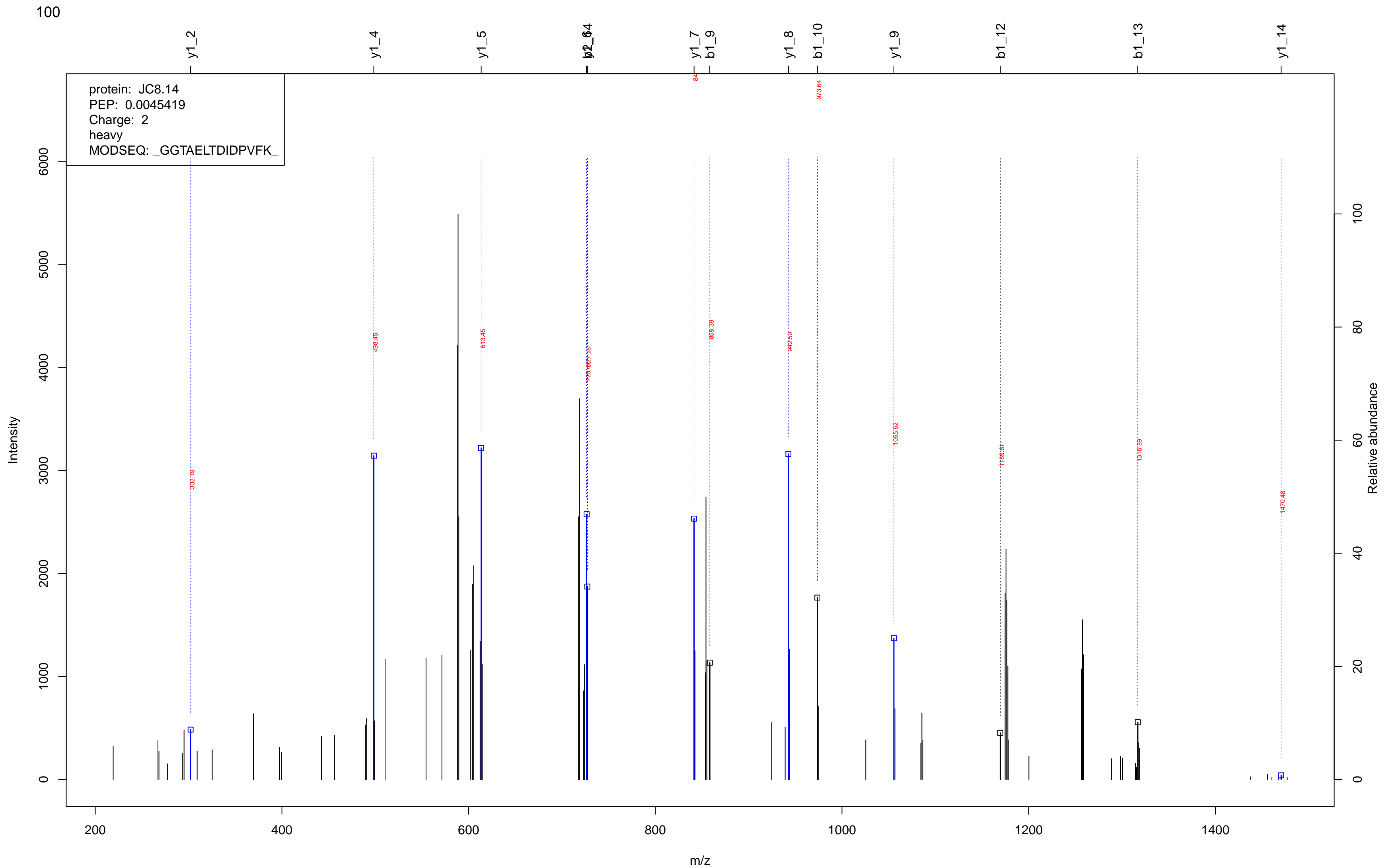


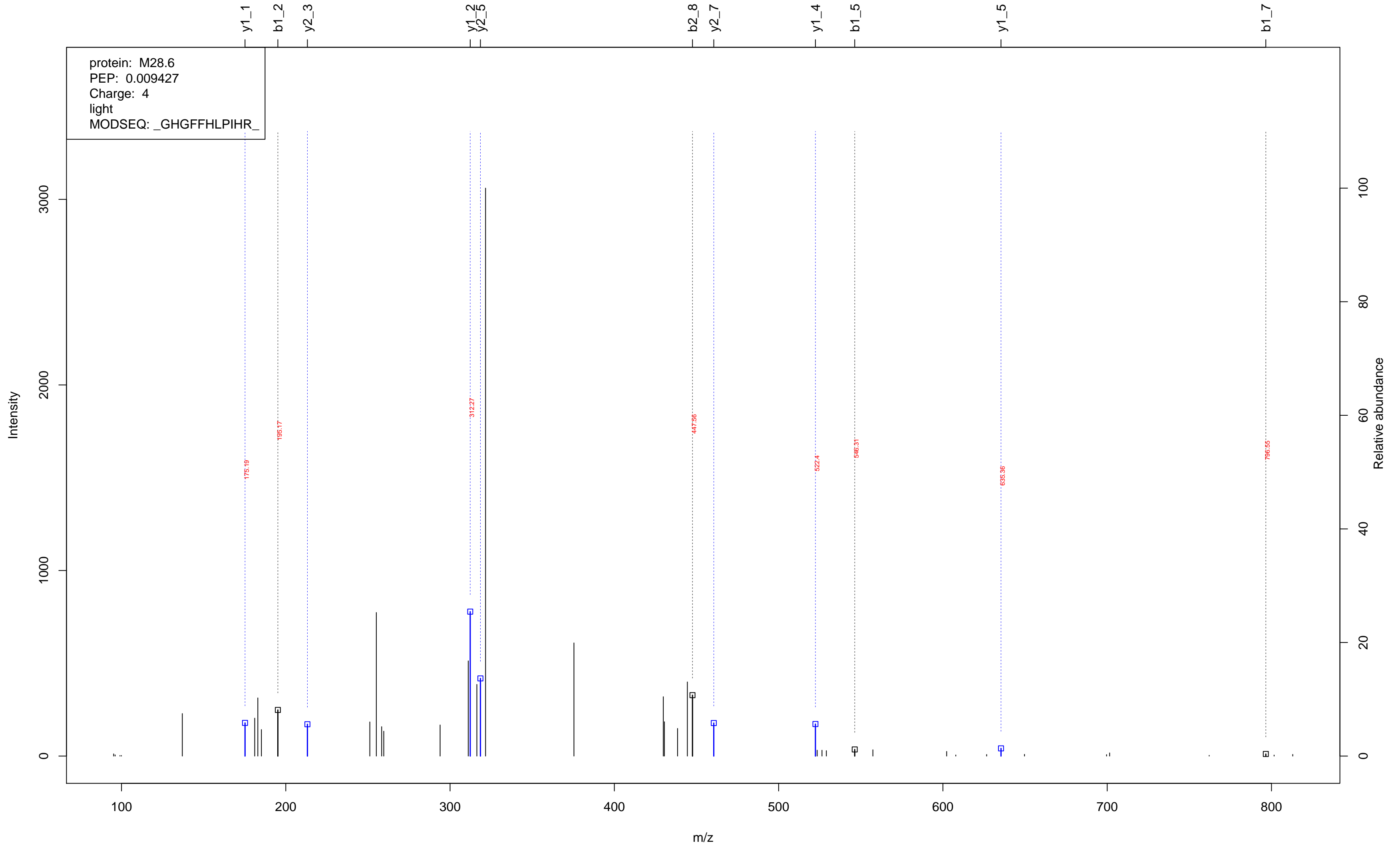


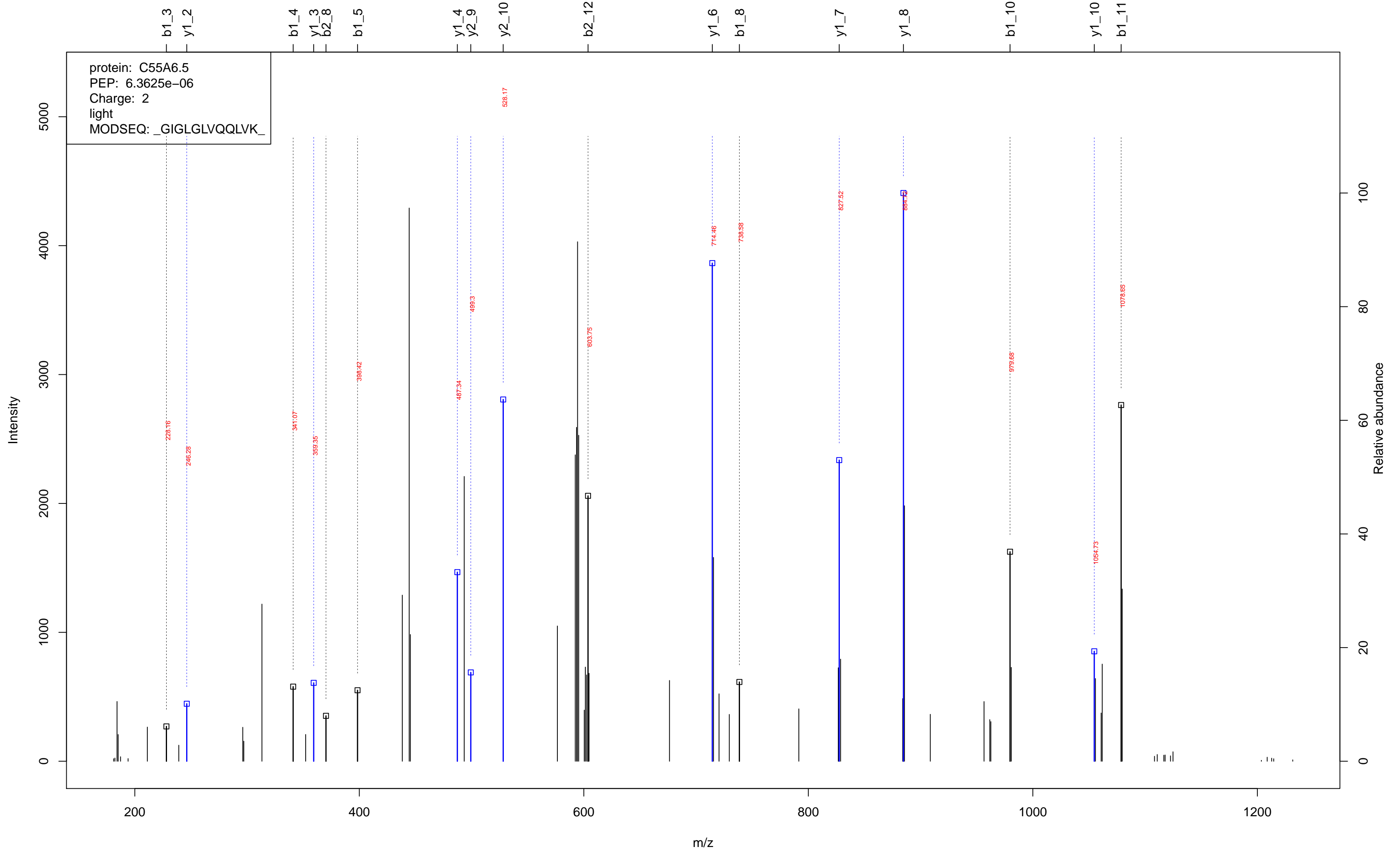


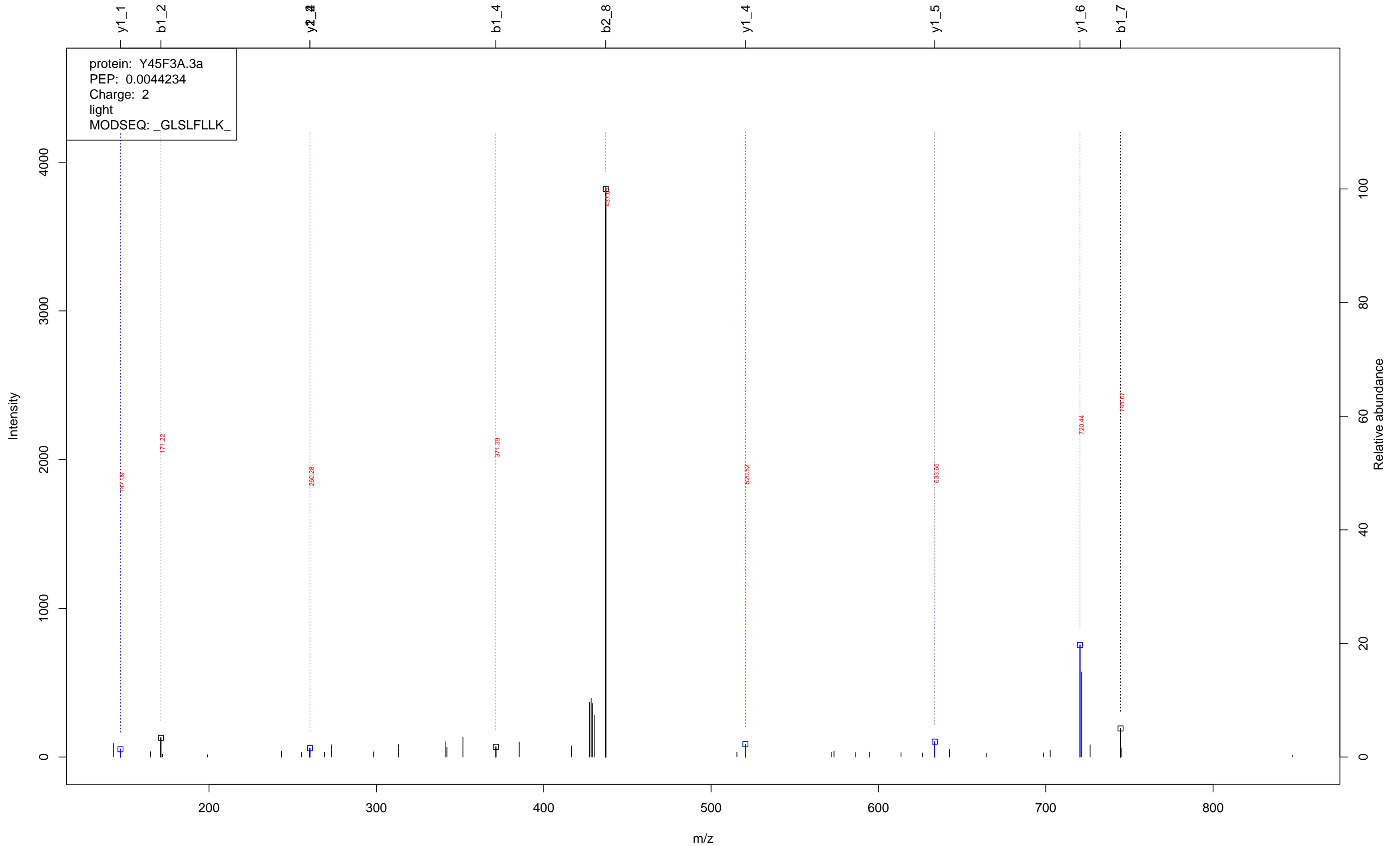


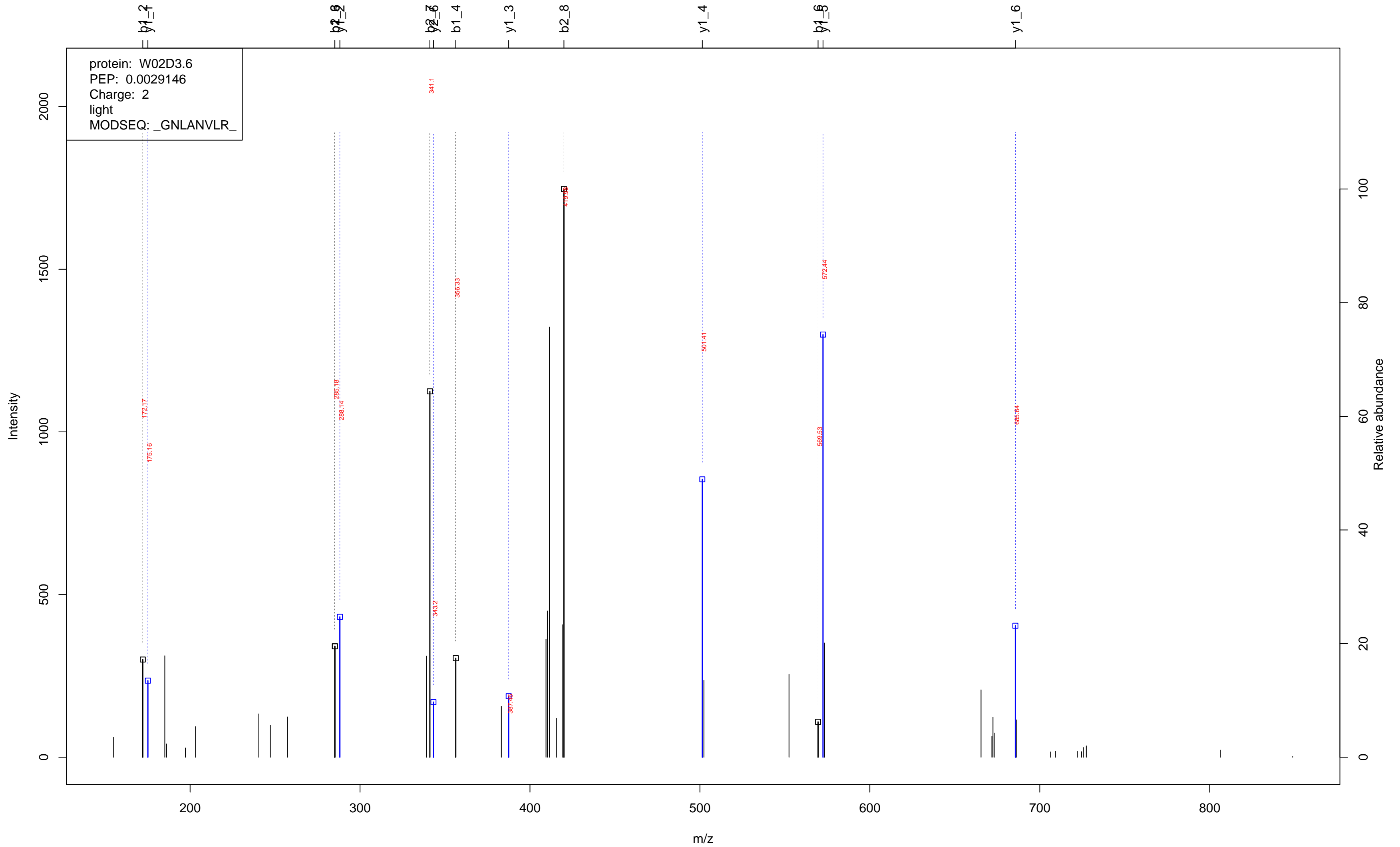


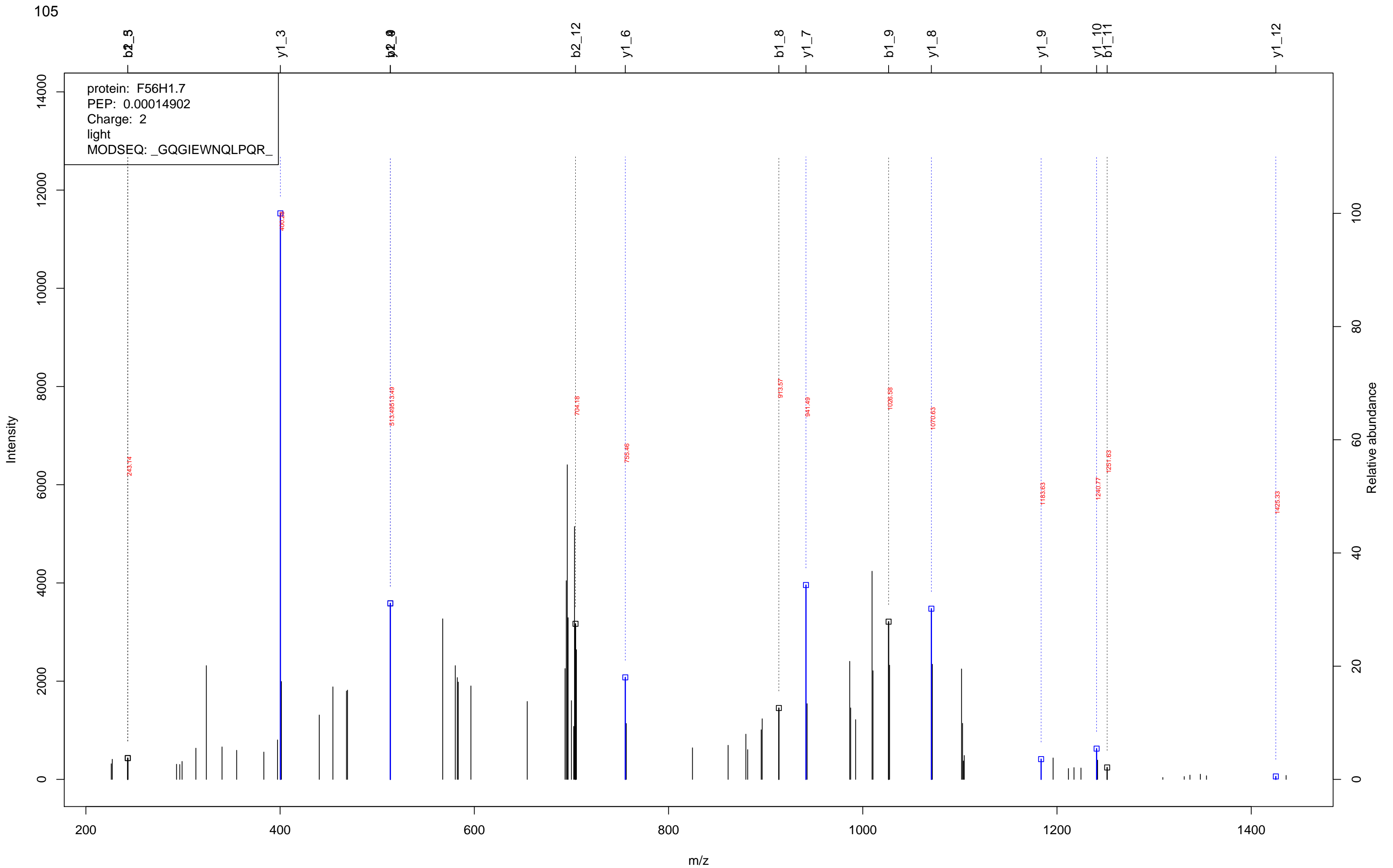


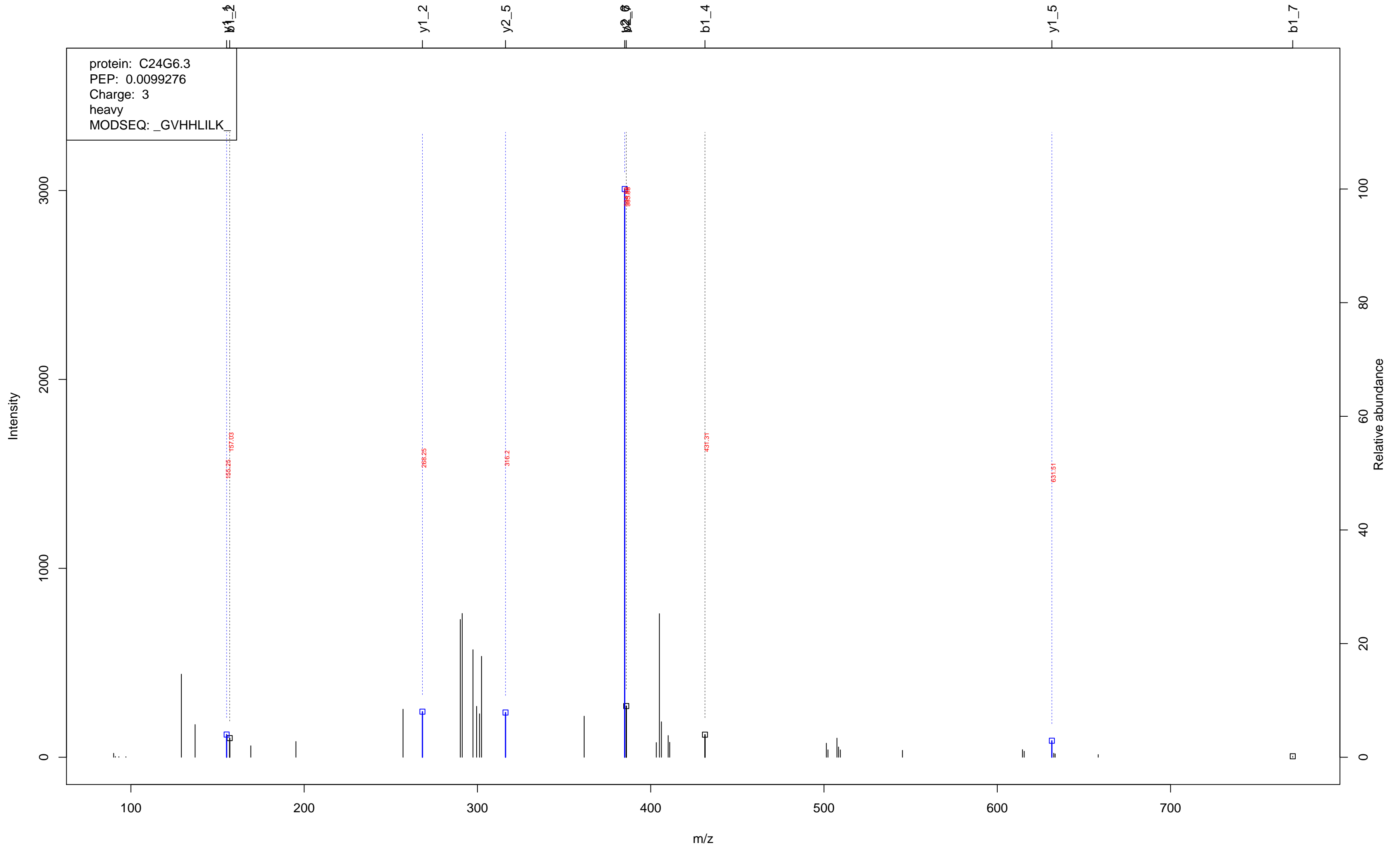


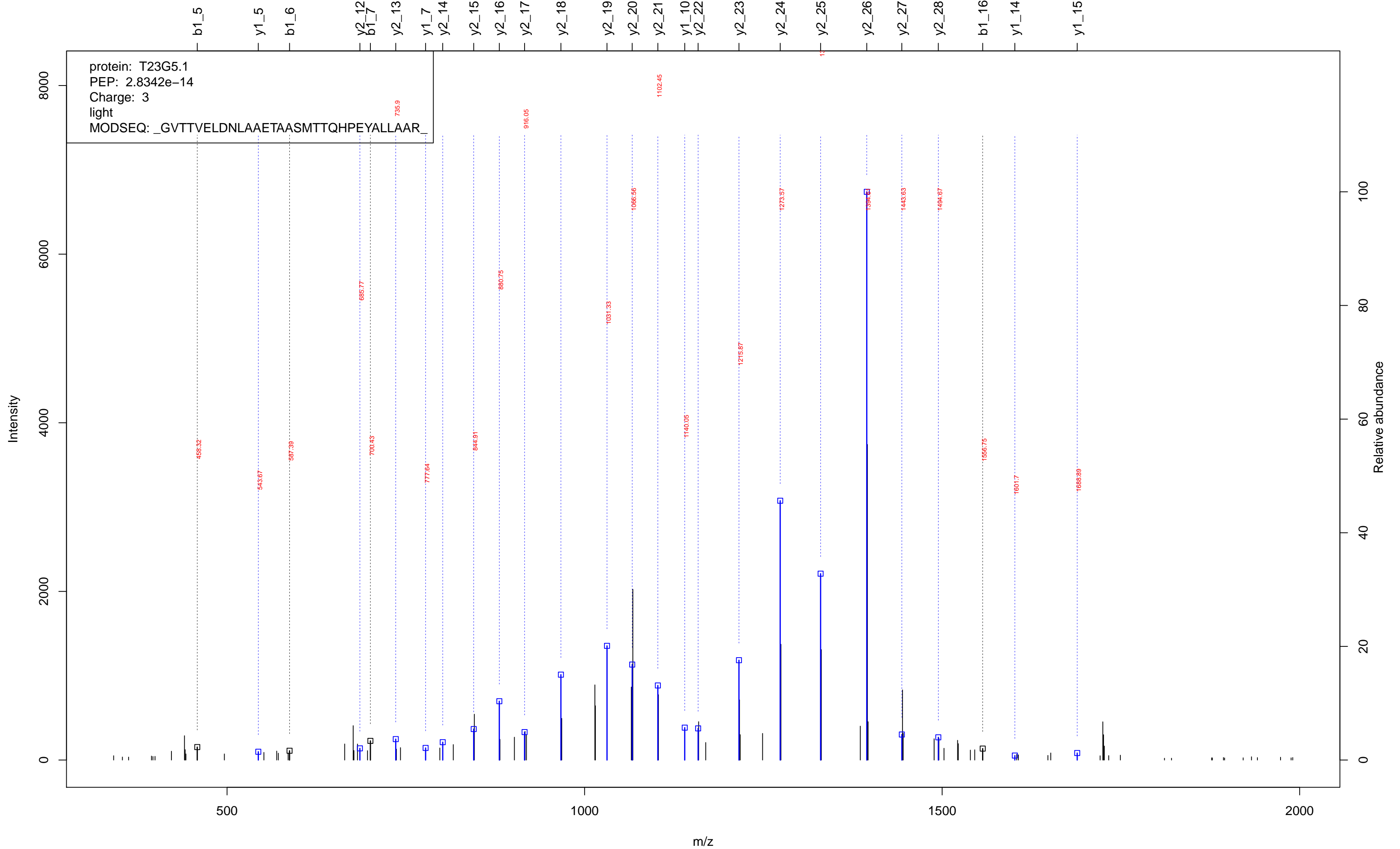


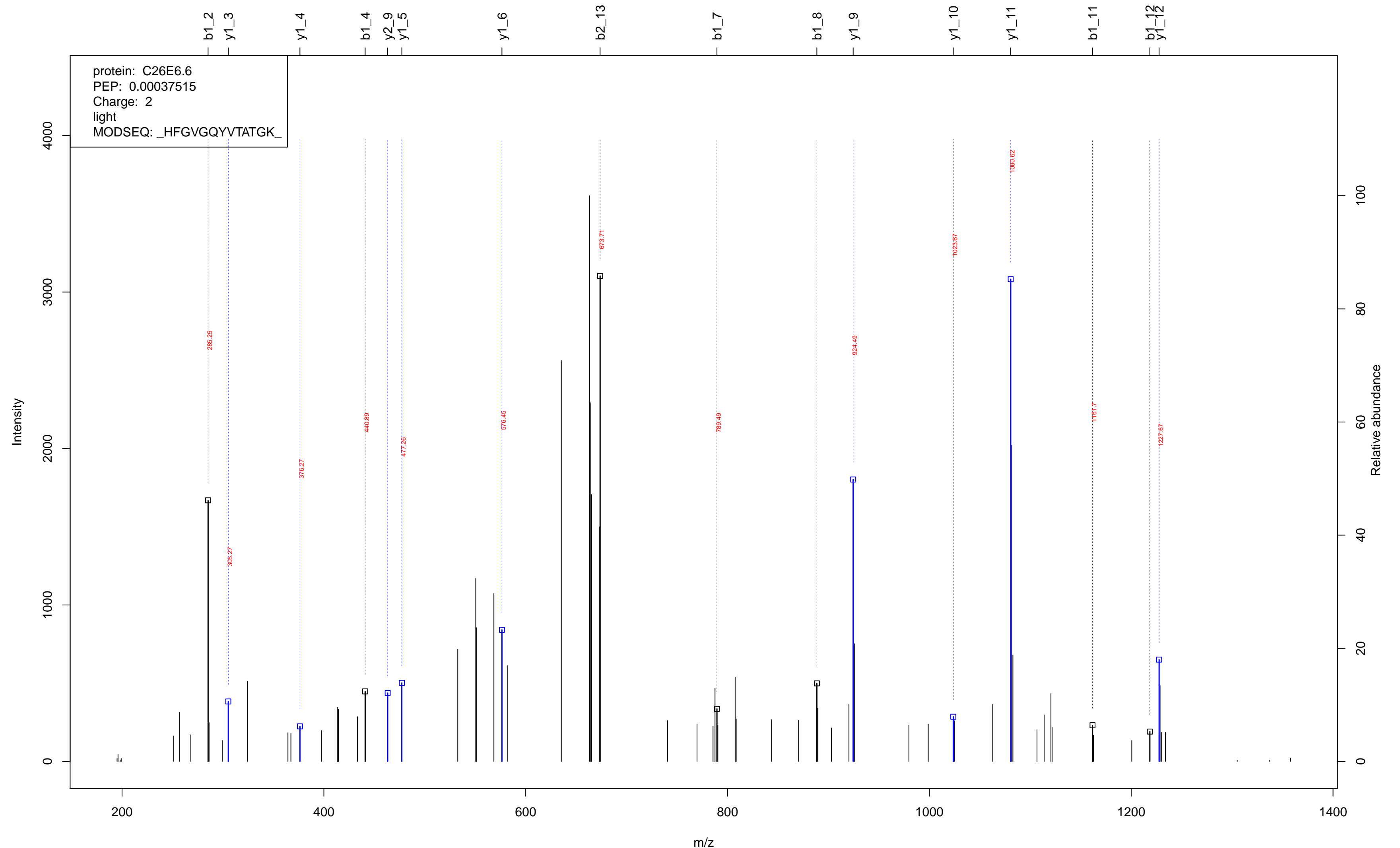


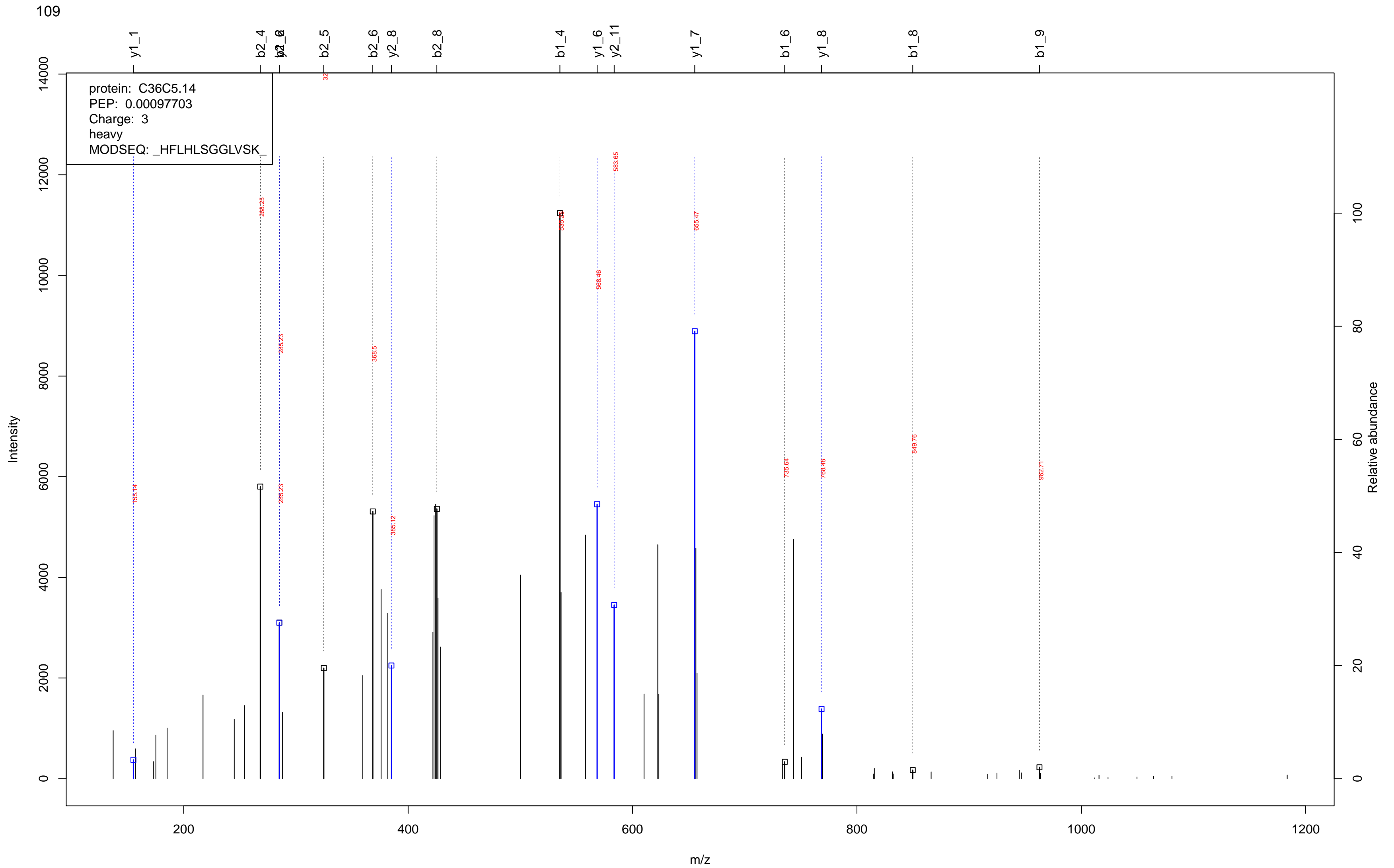


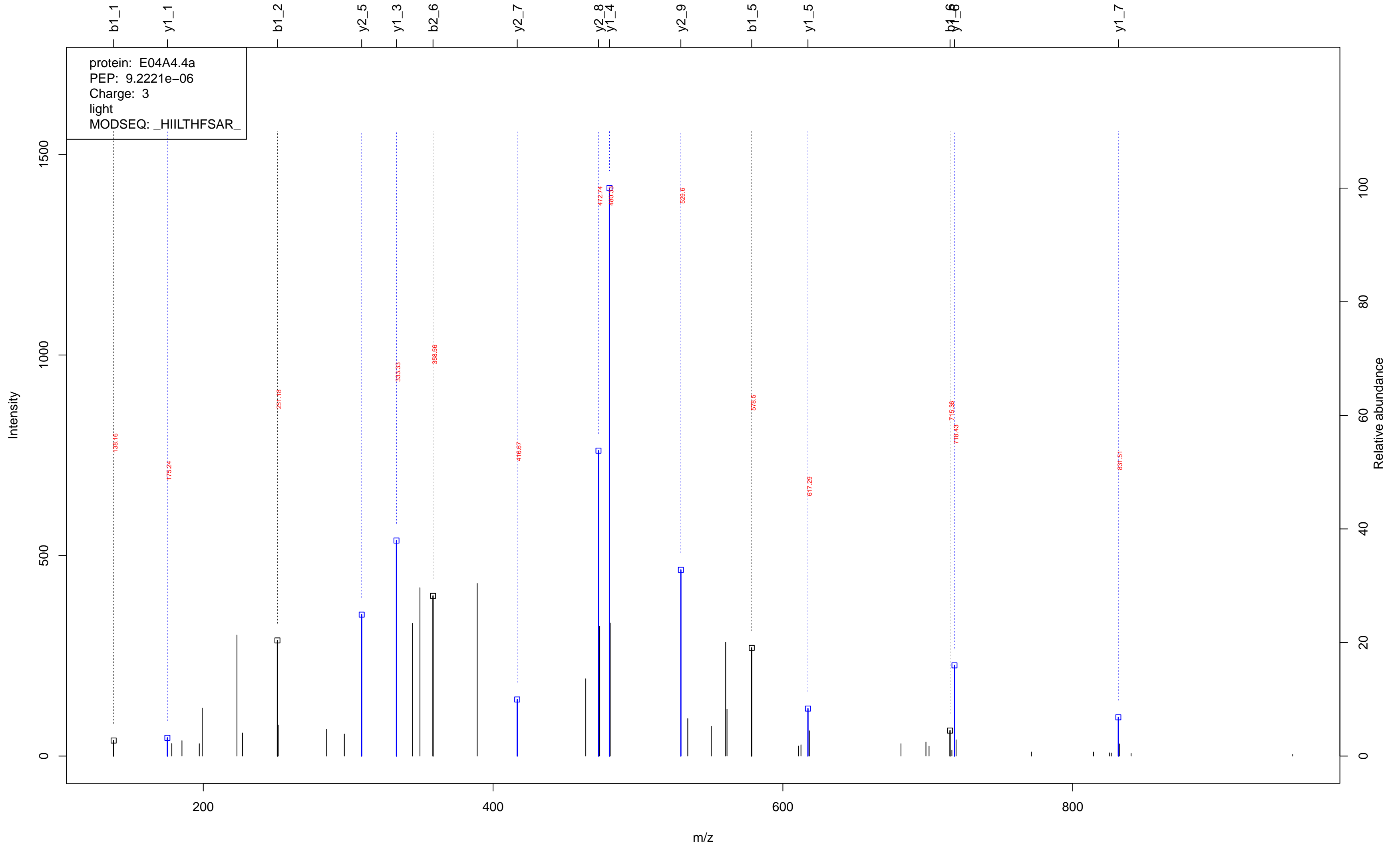


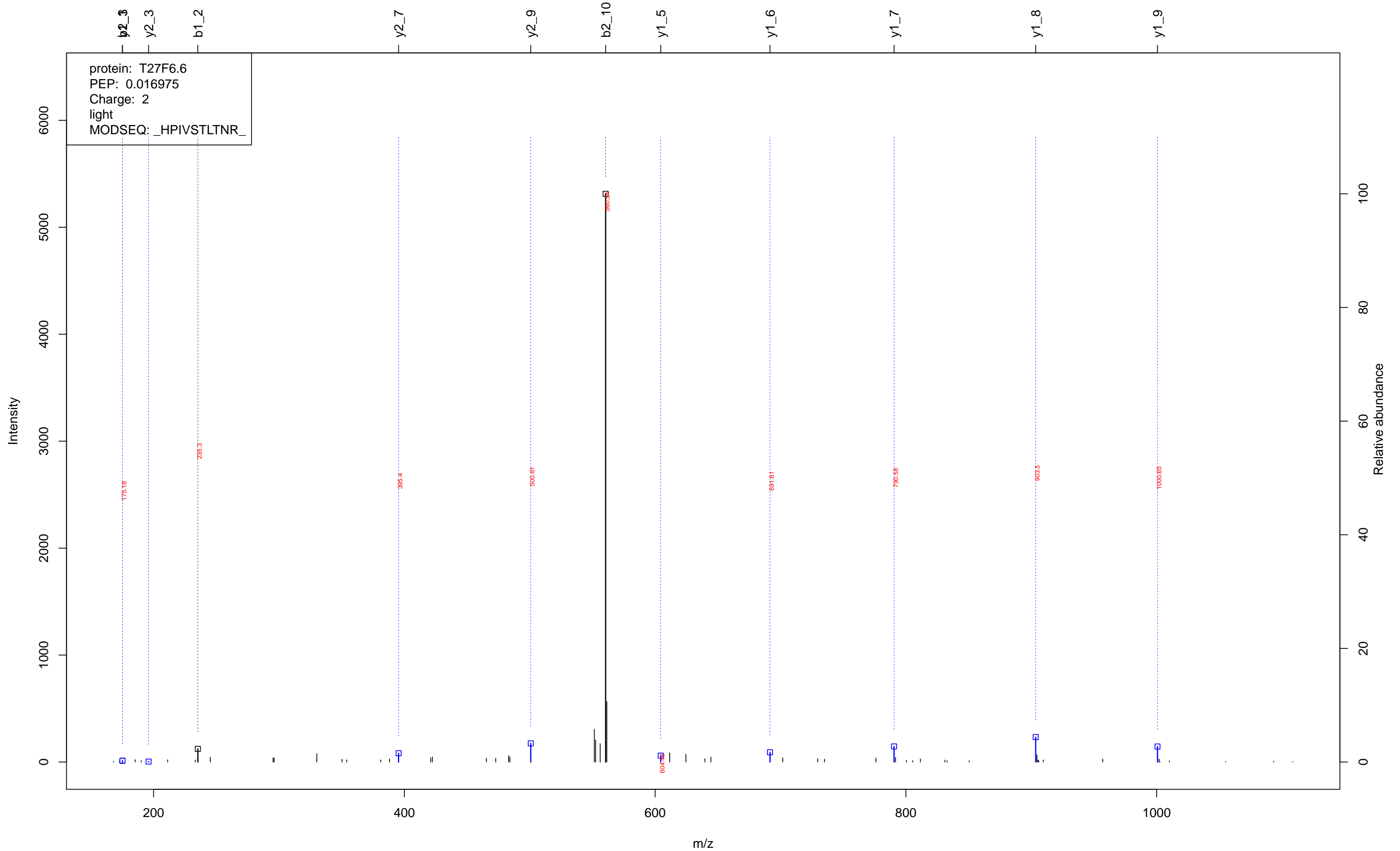


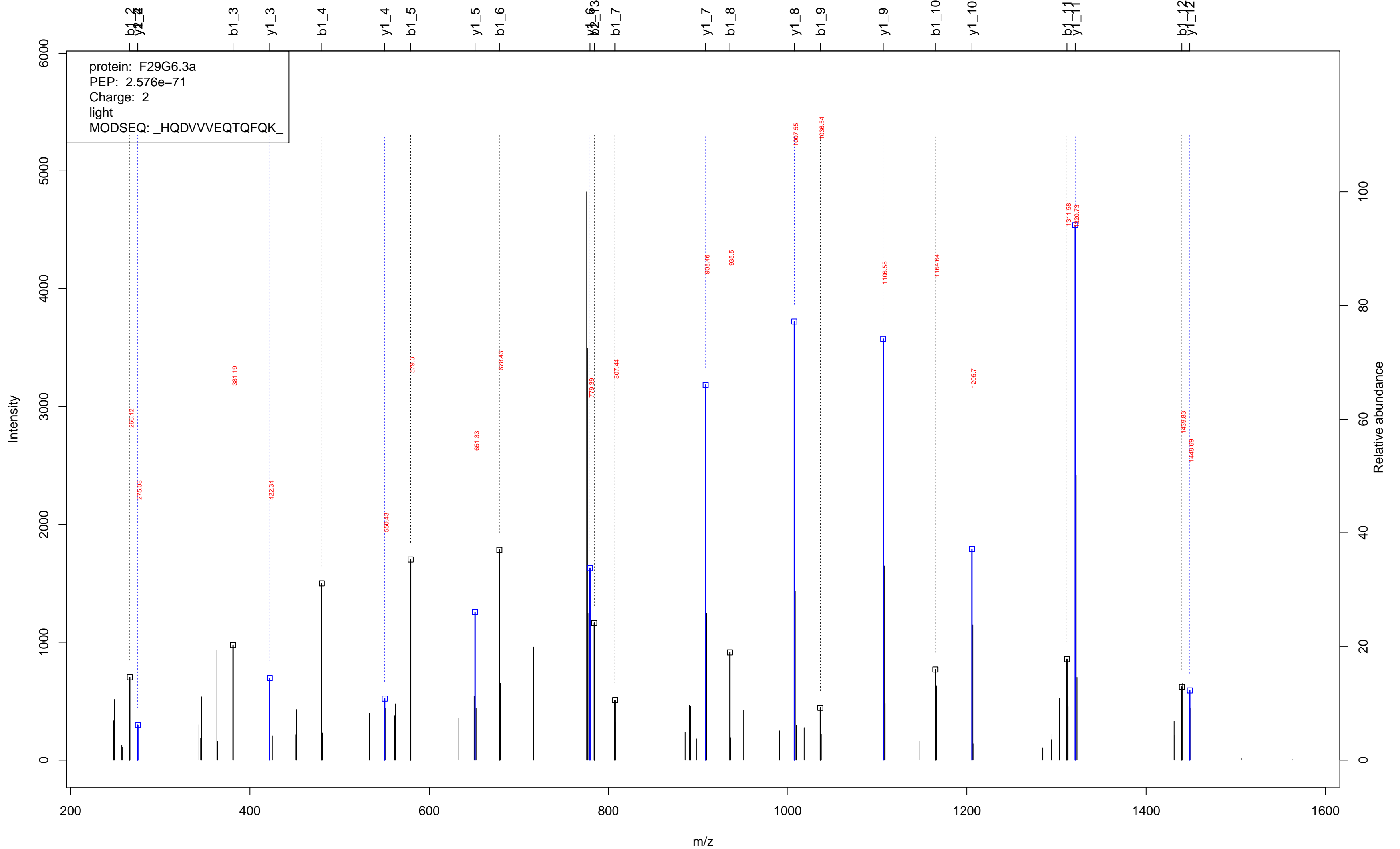




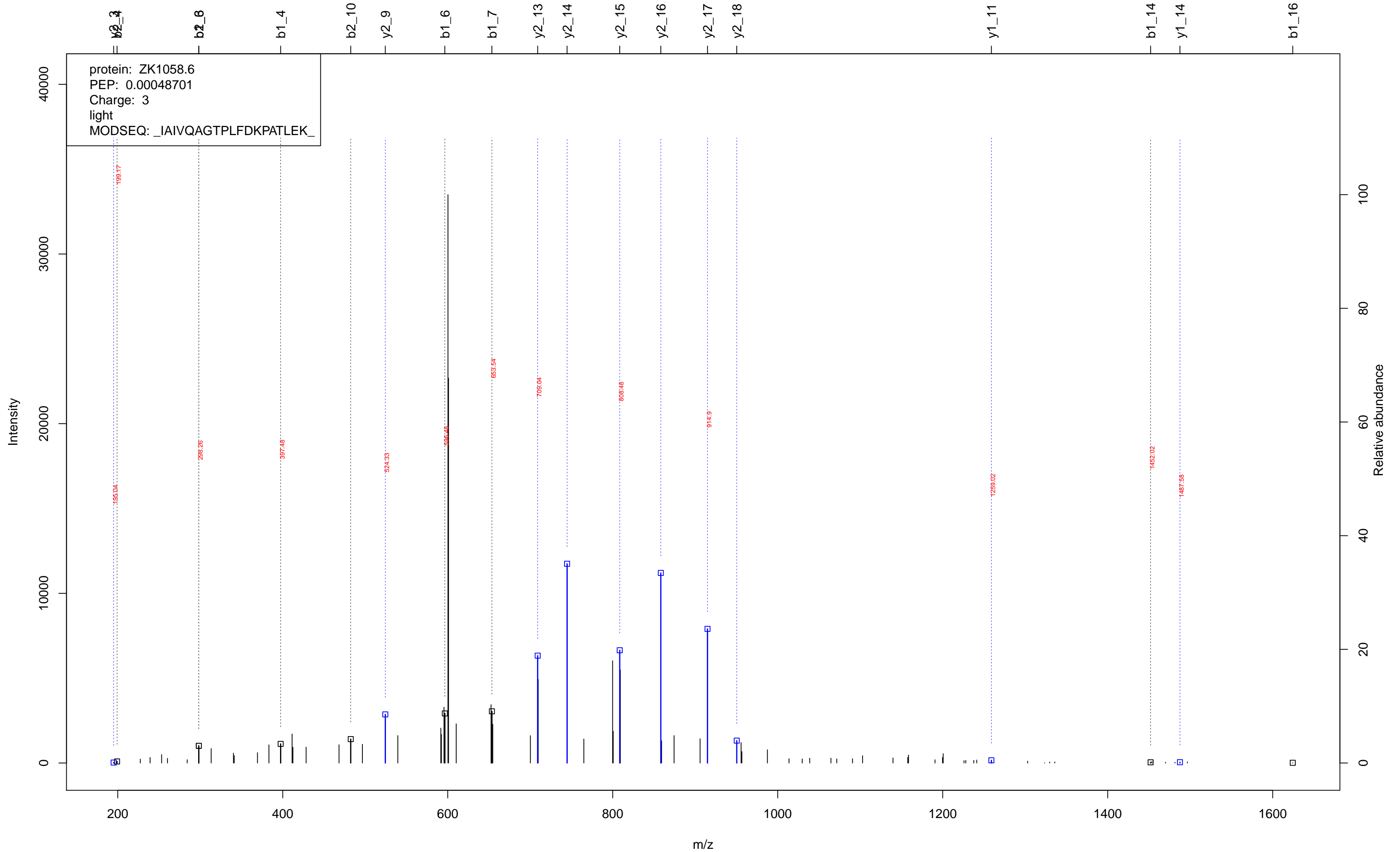


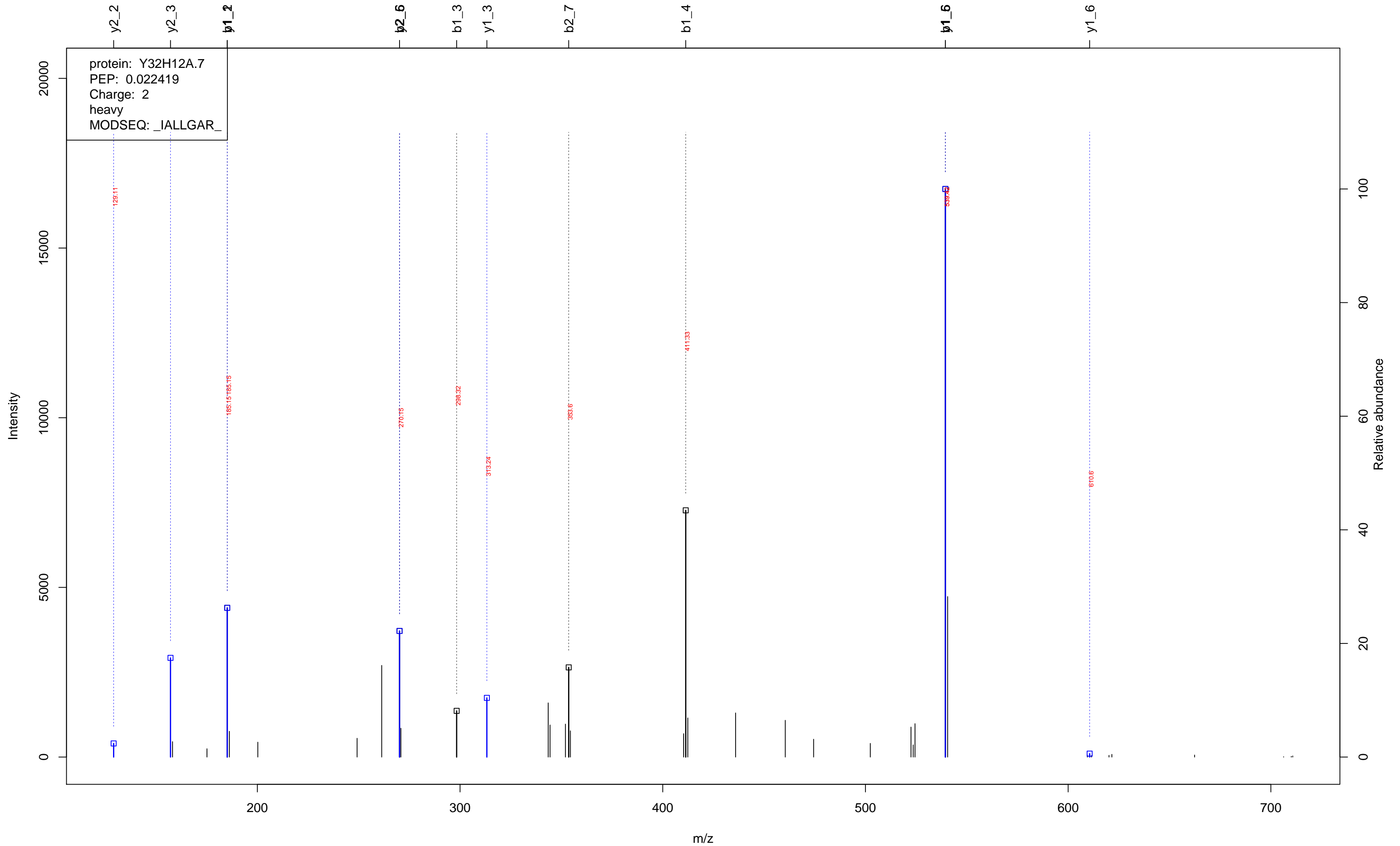


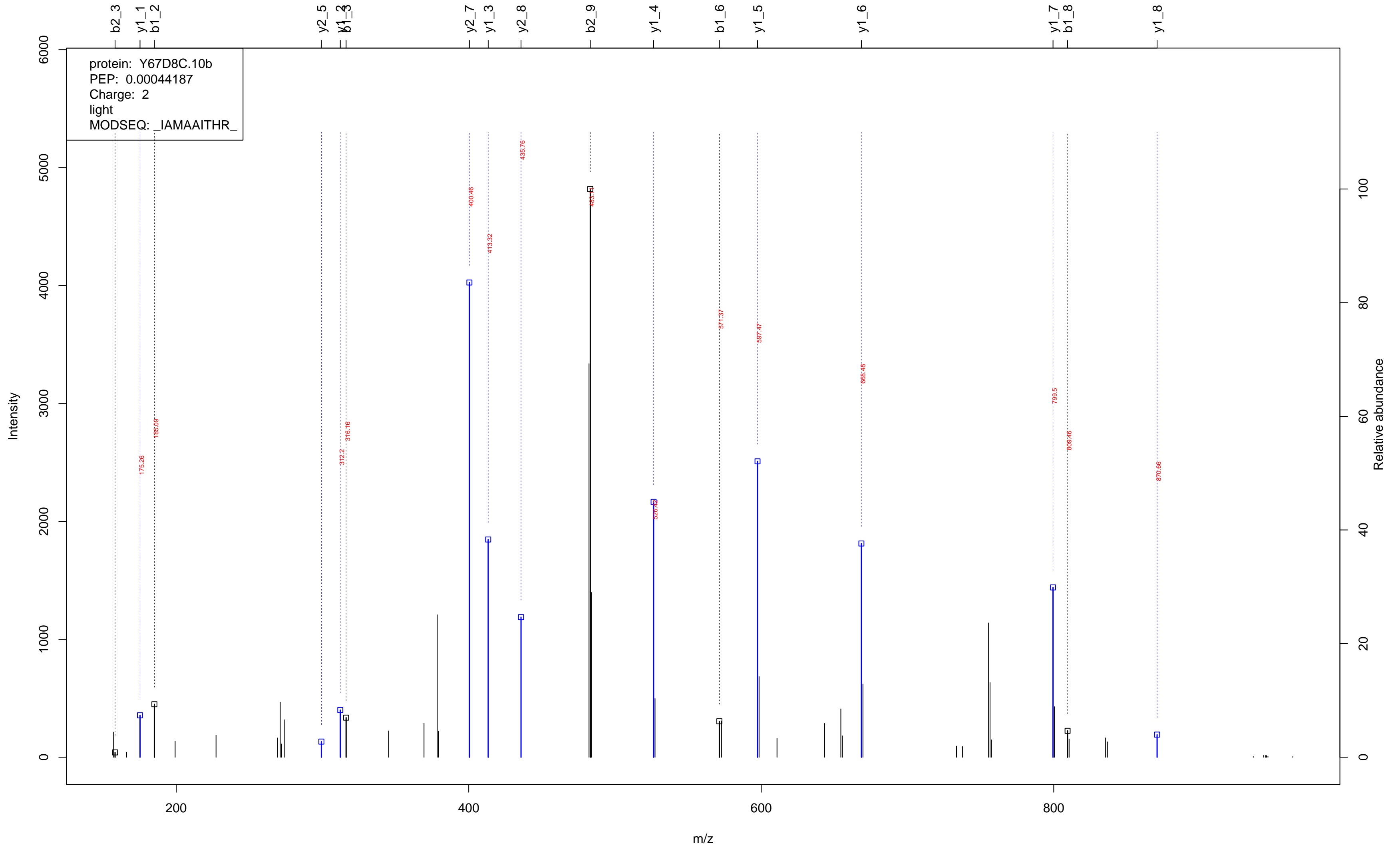


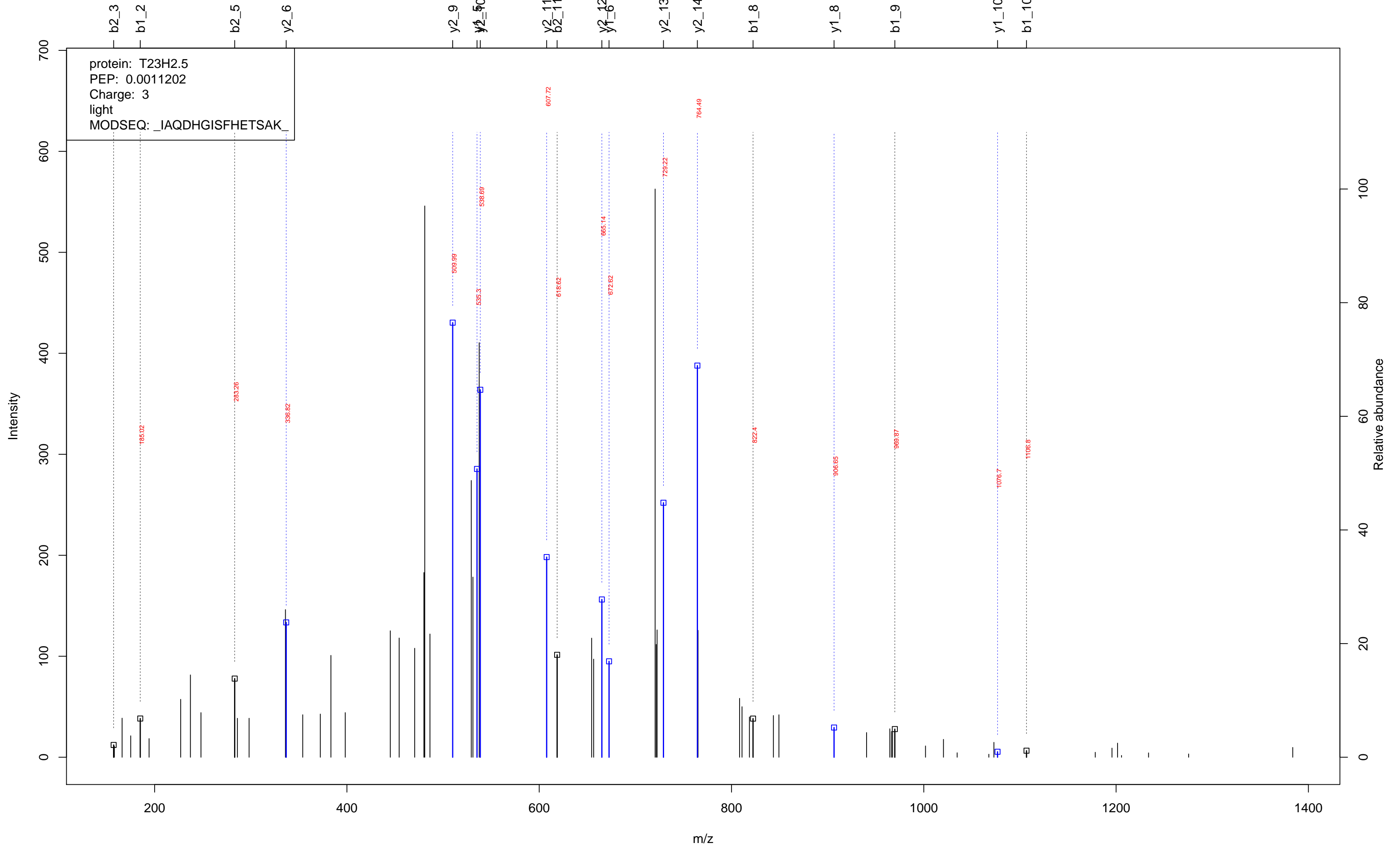


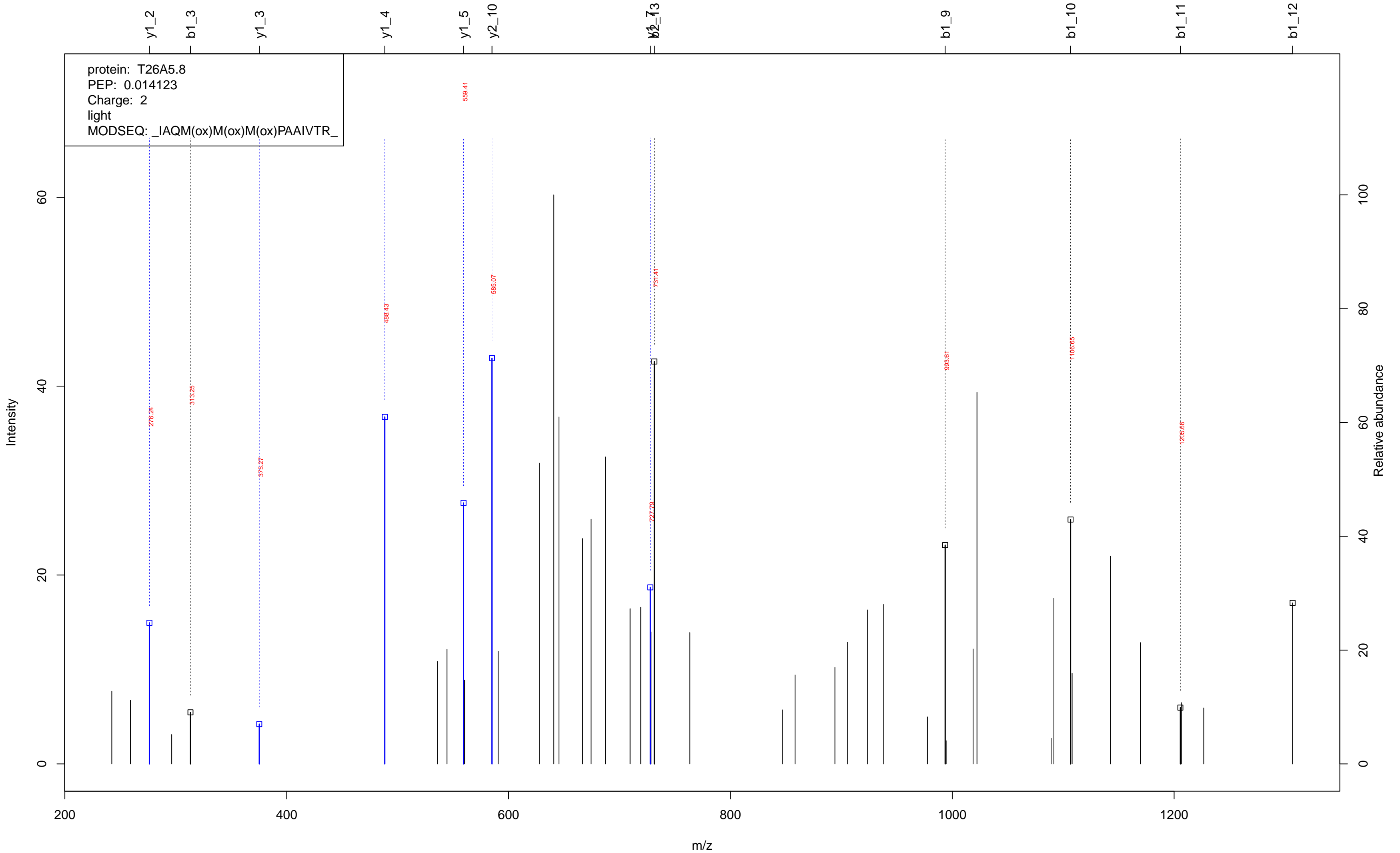
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Charge: 2
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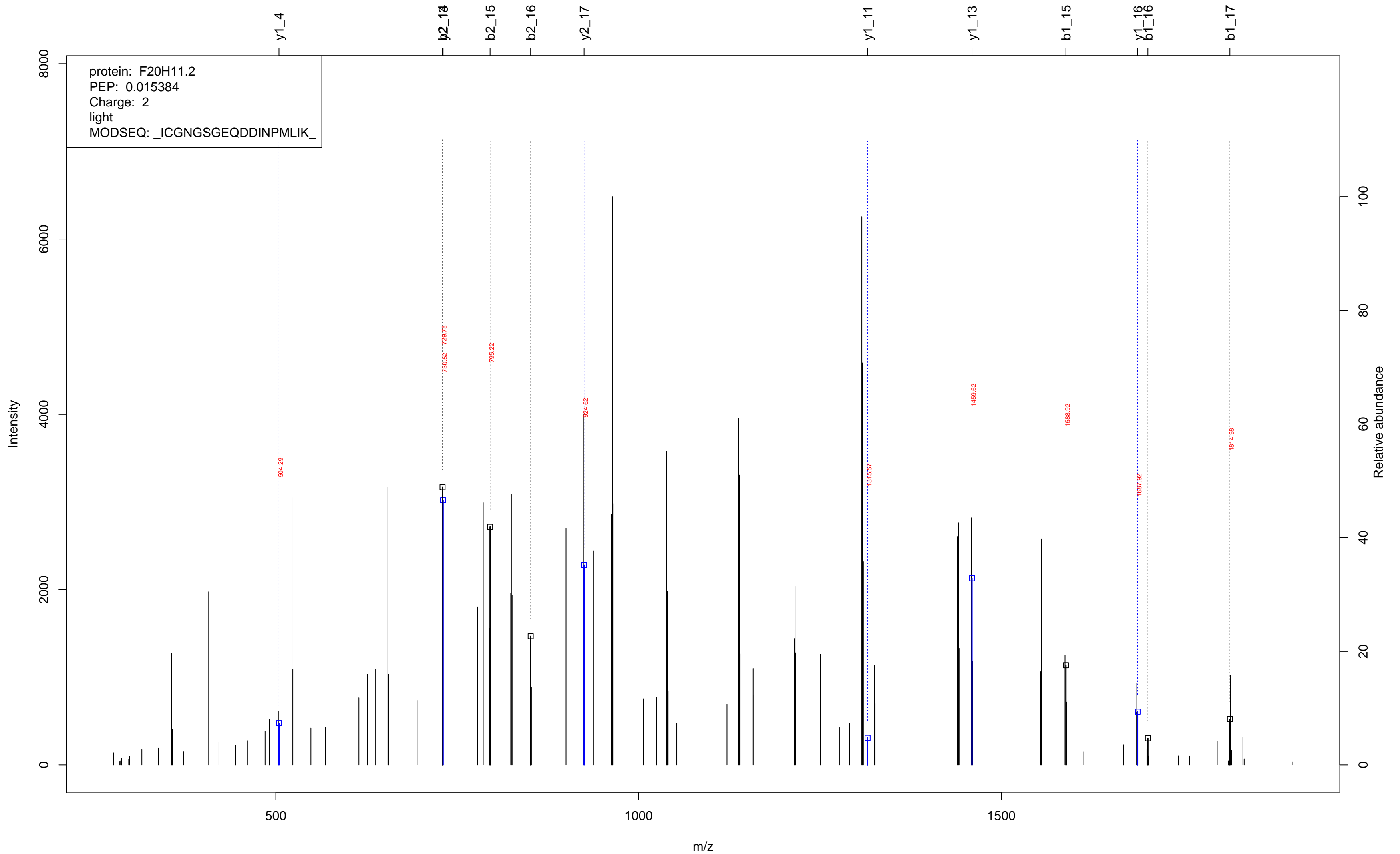


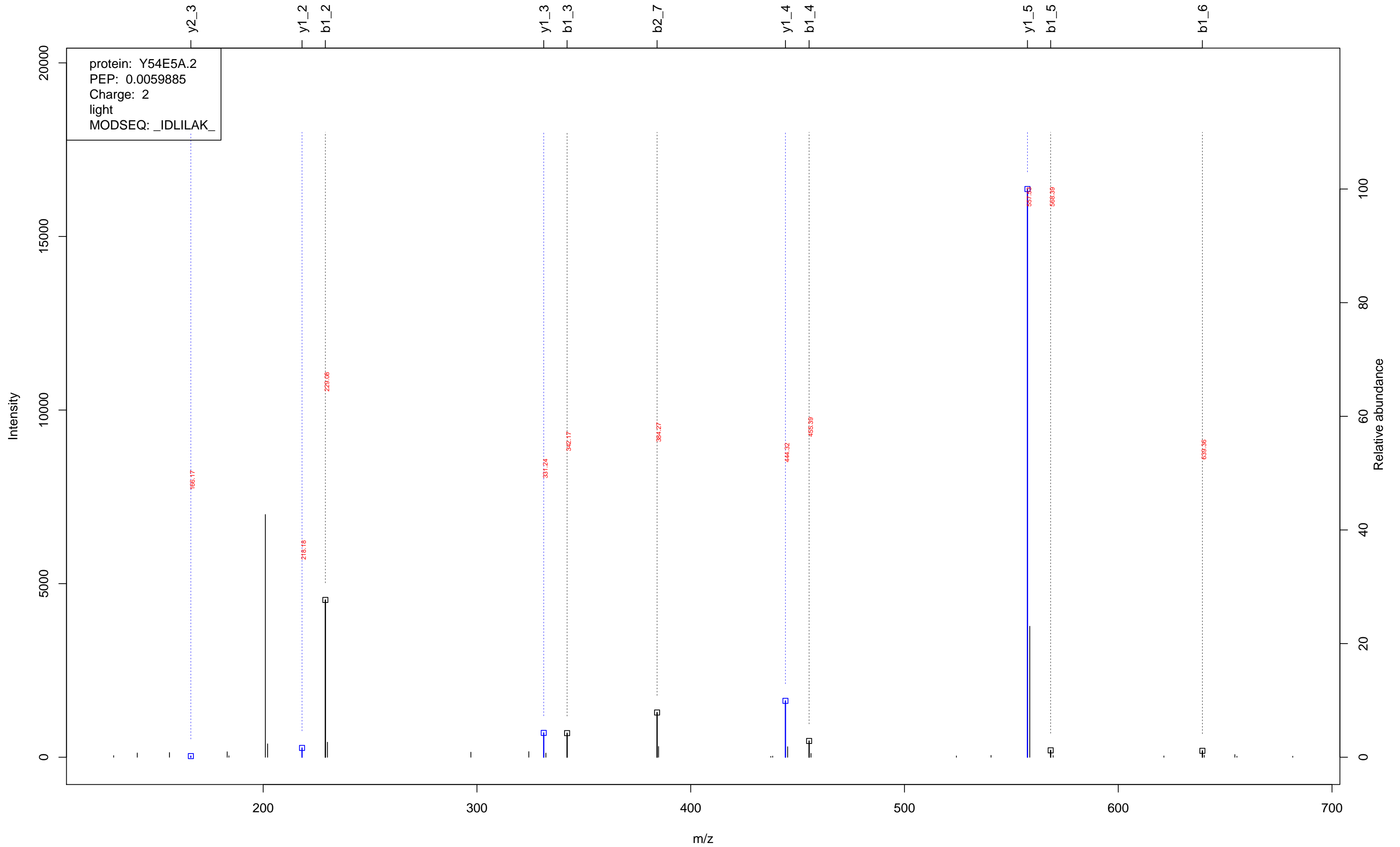


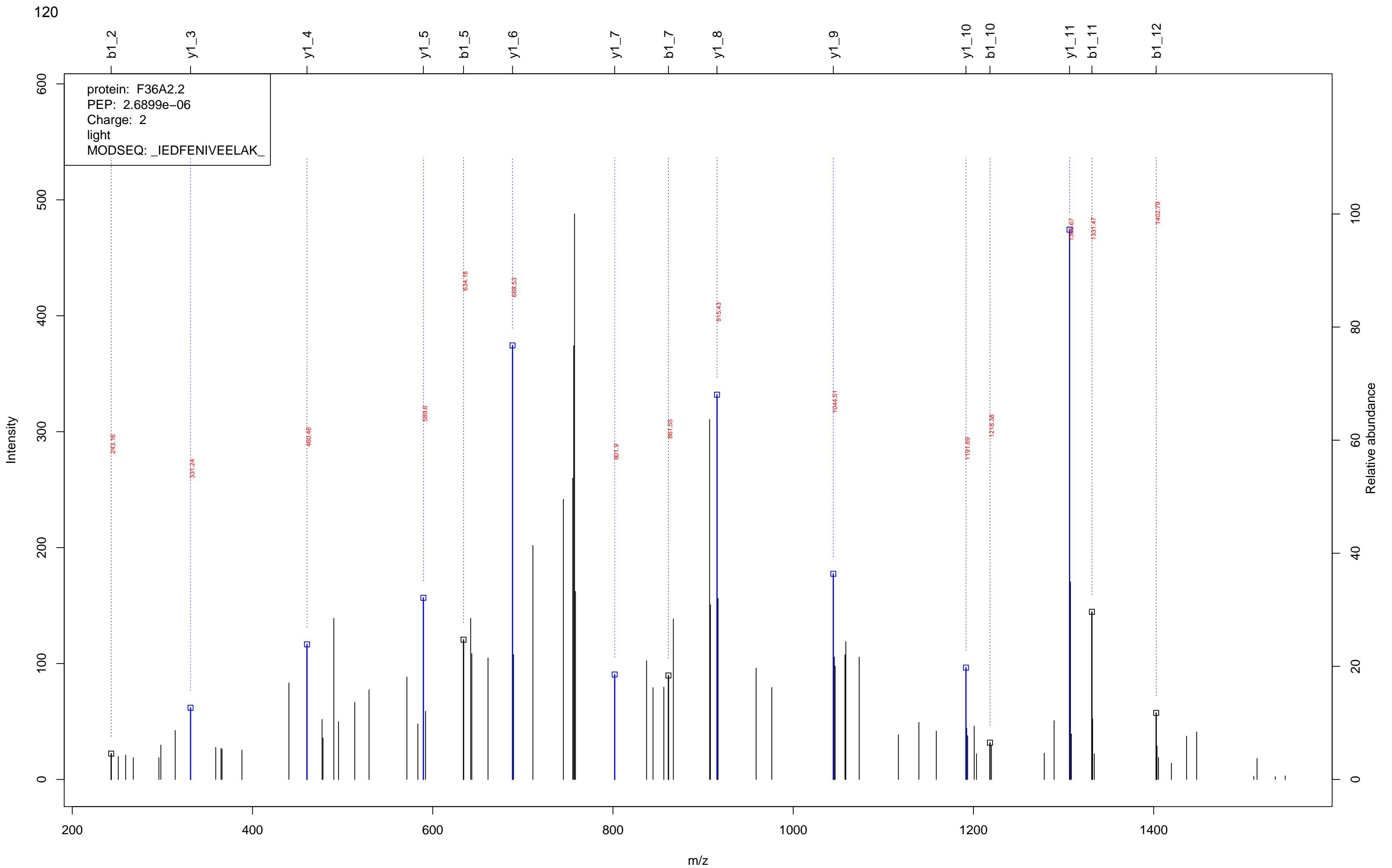


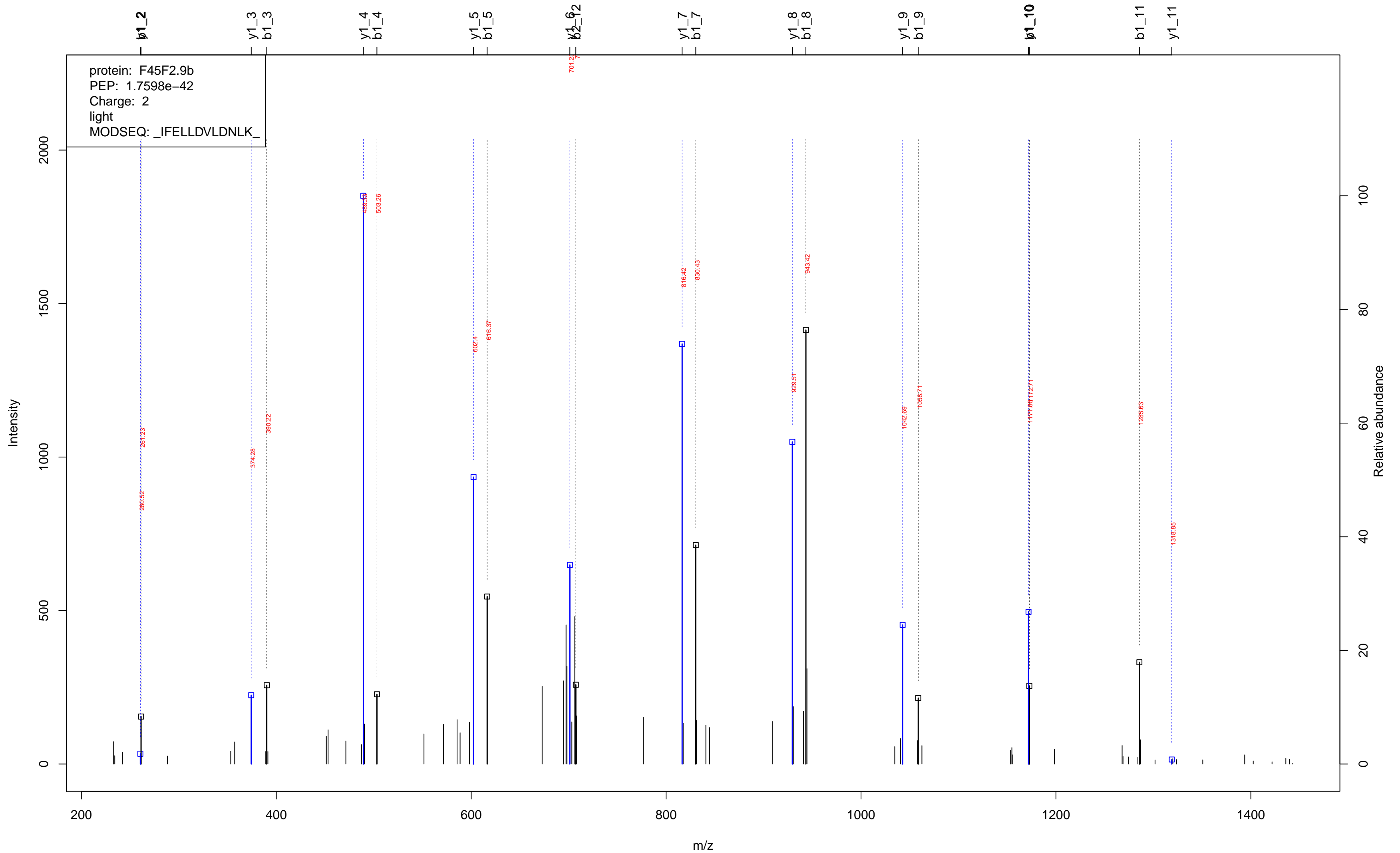


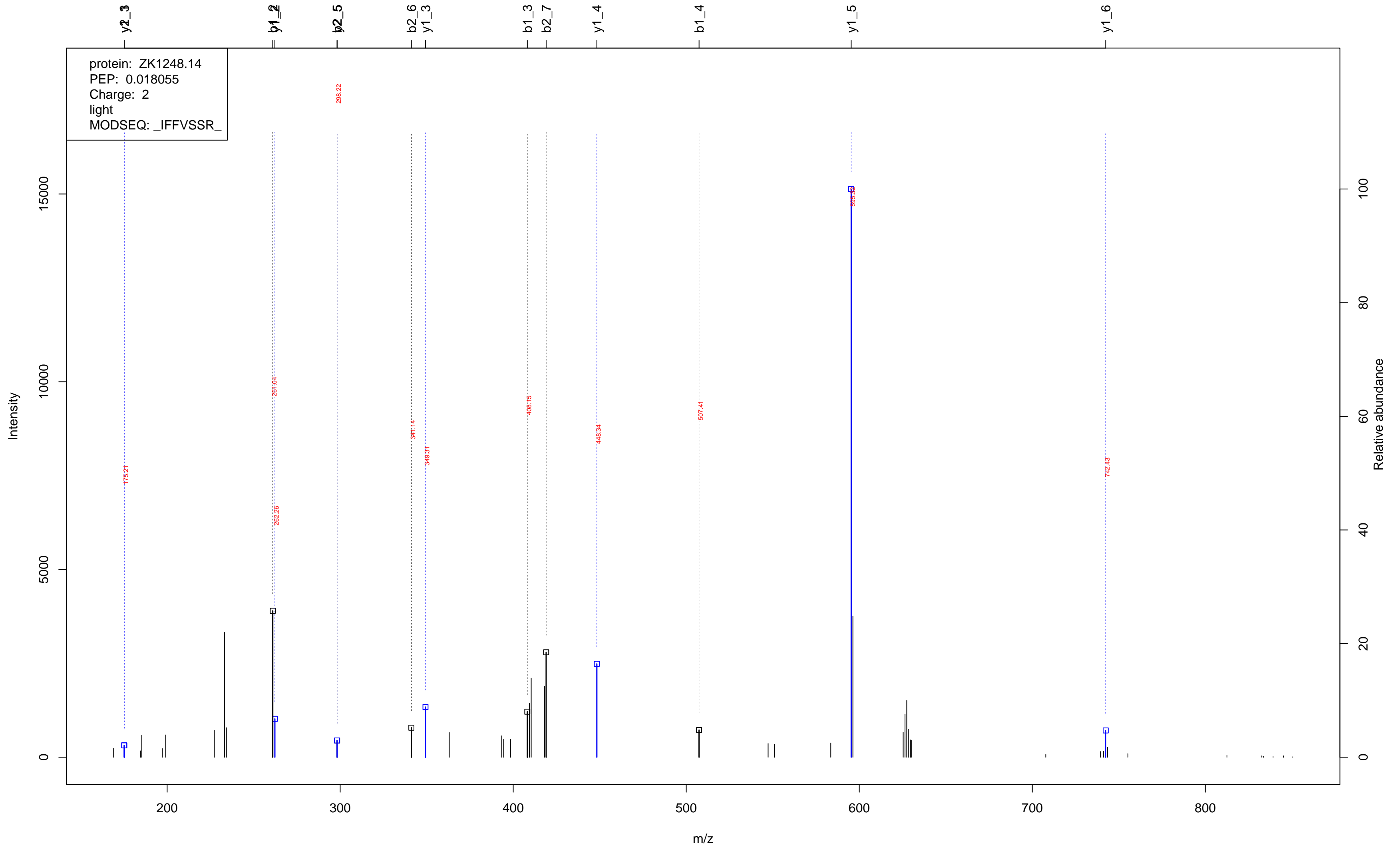


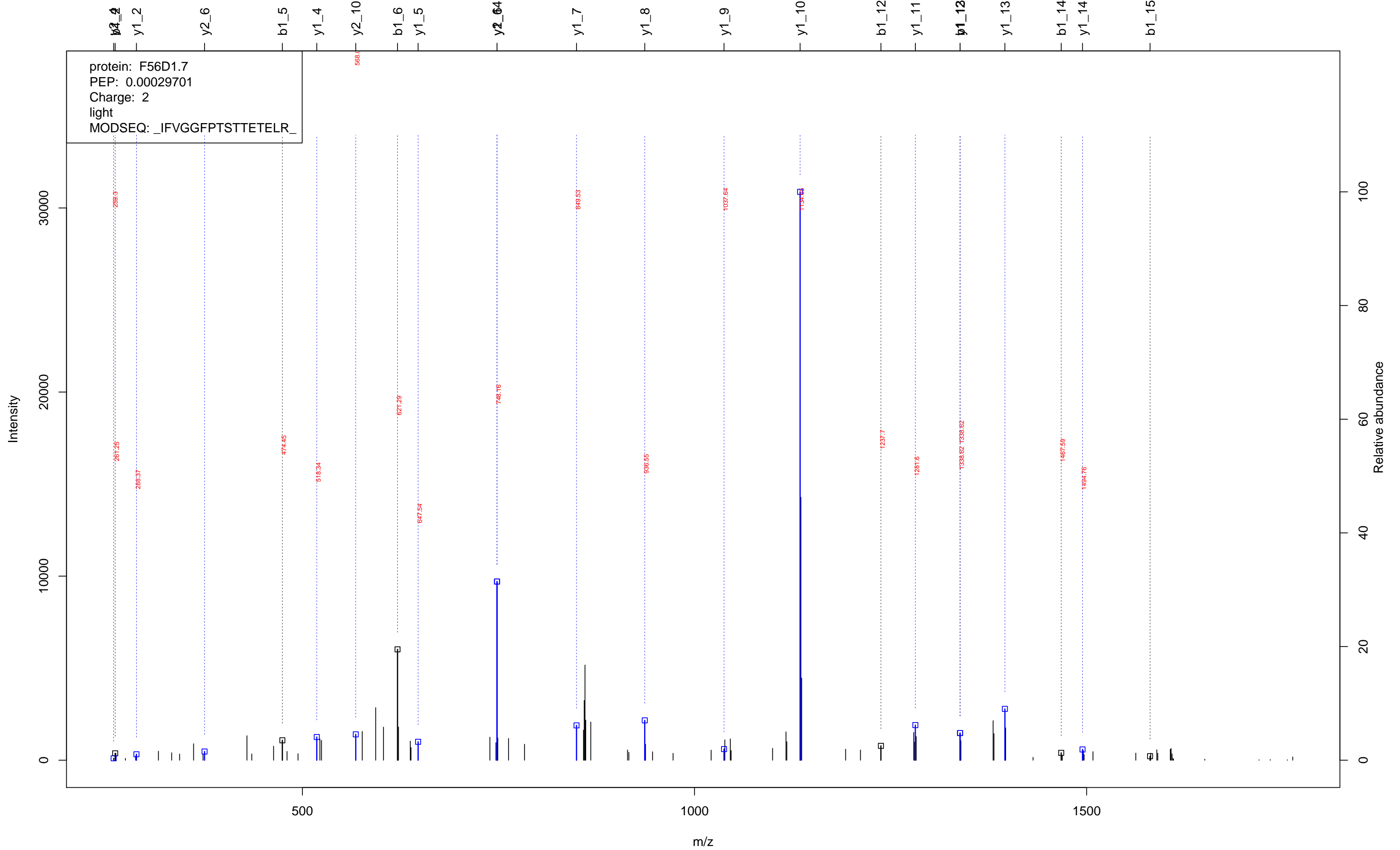


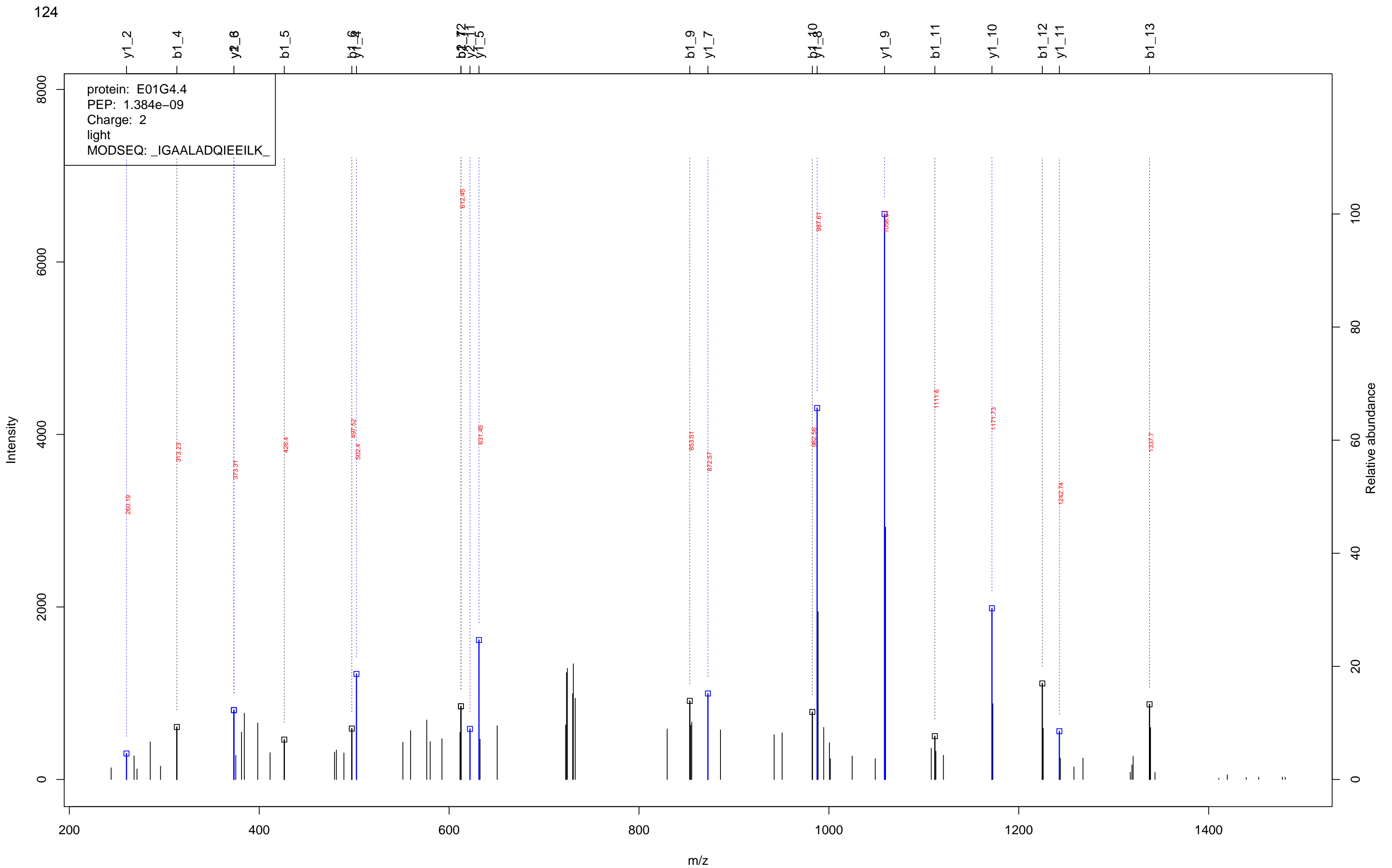




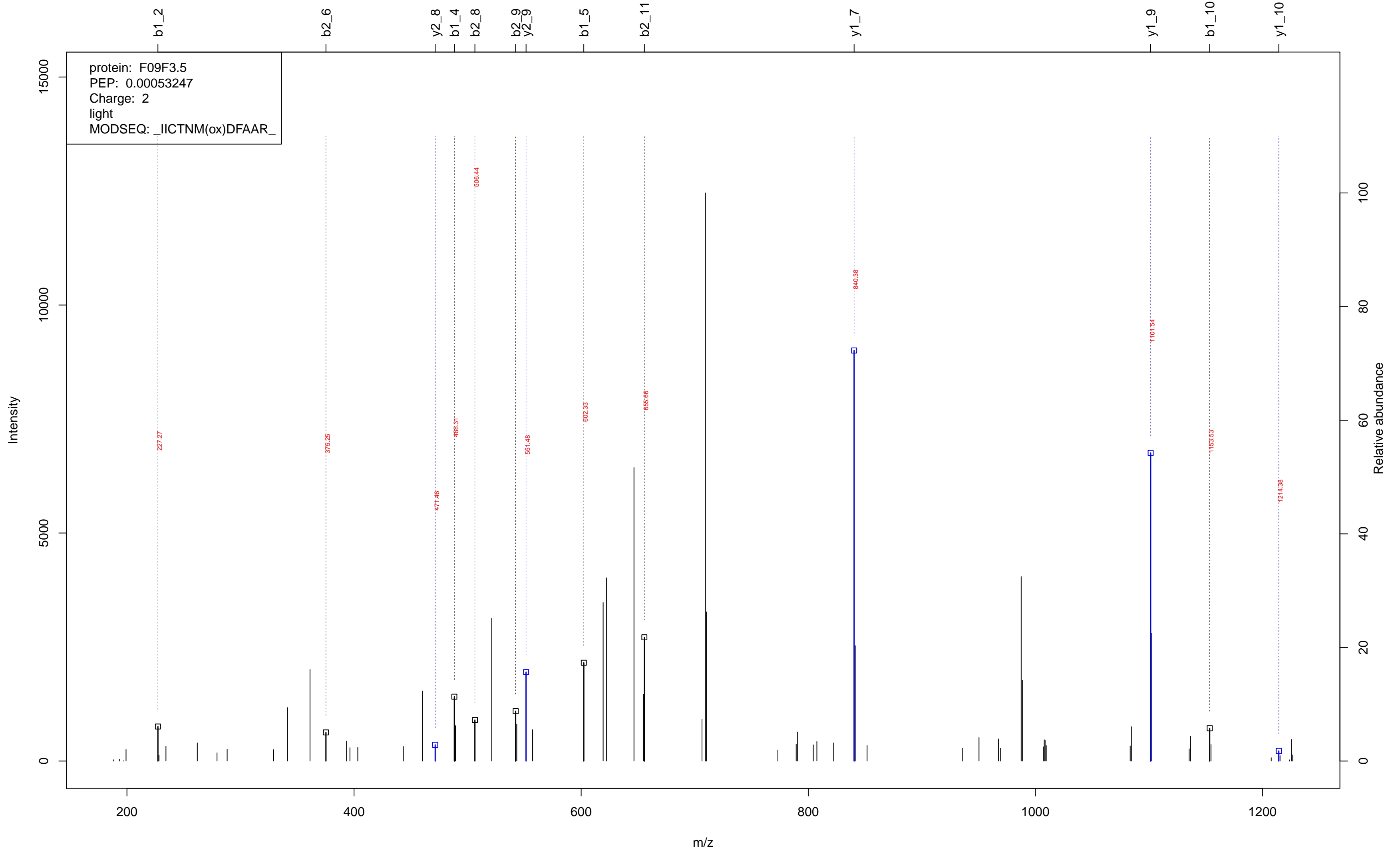


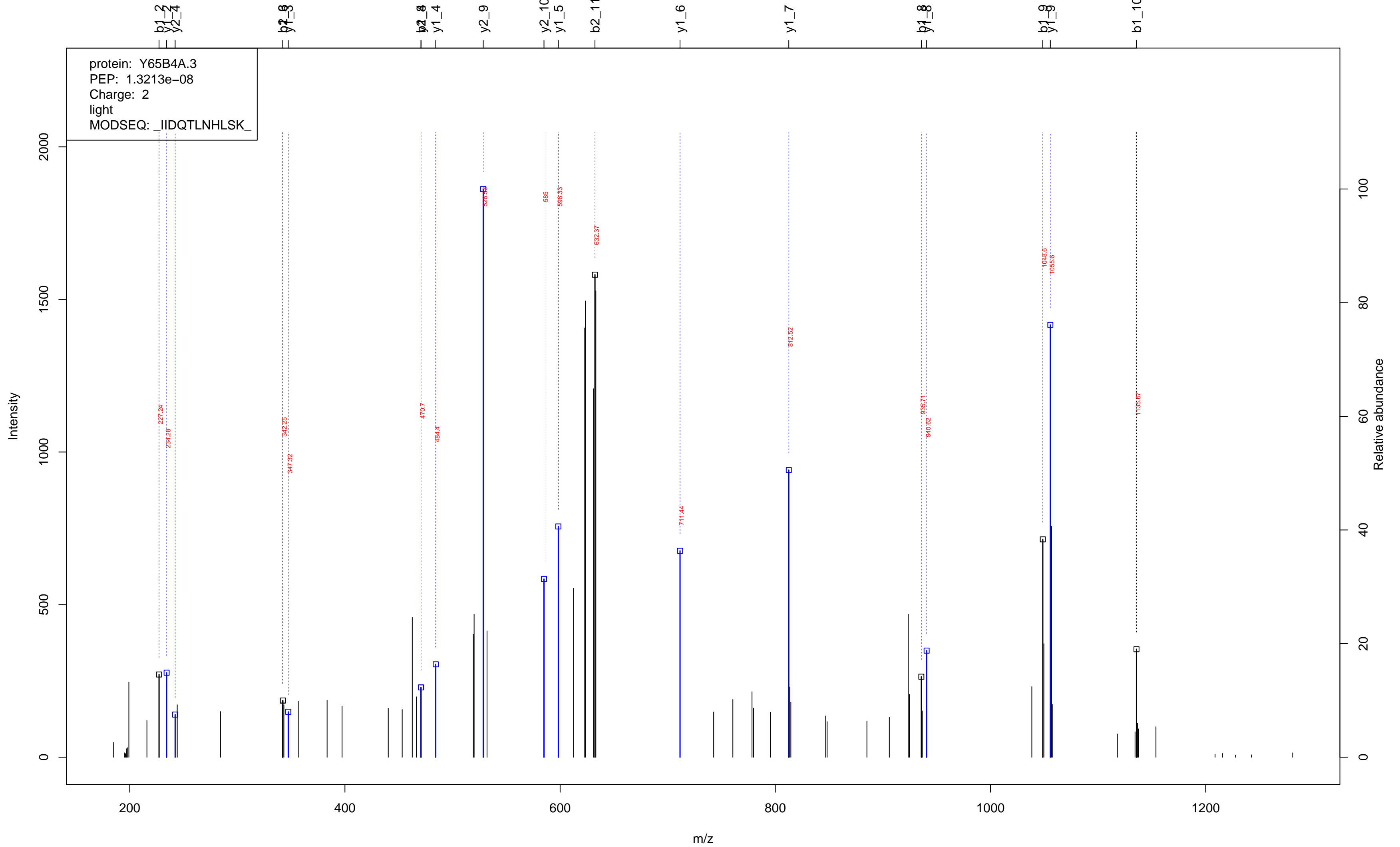


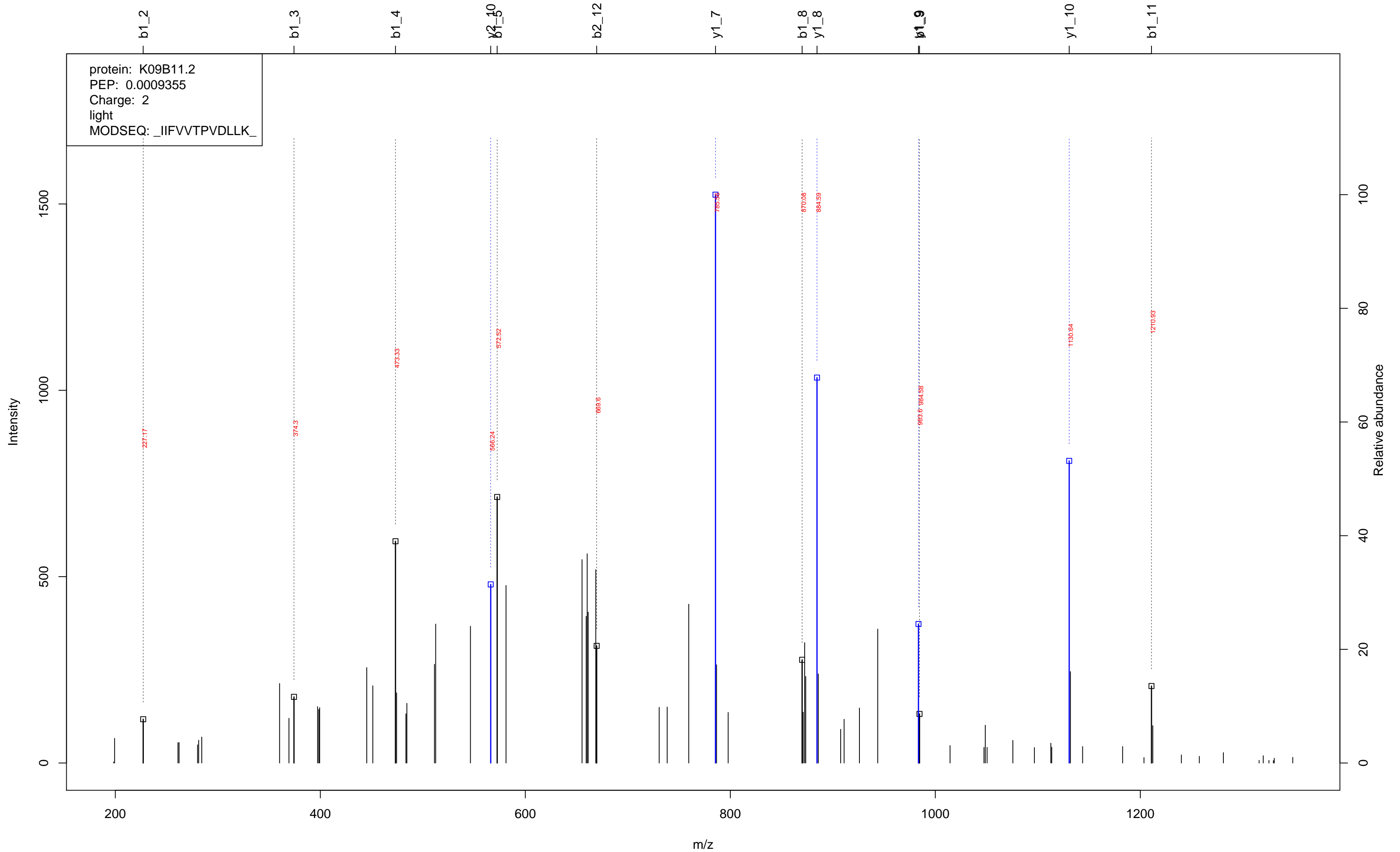


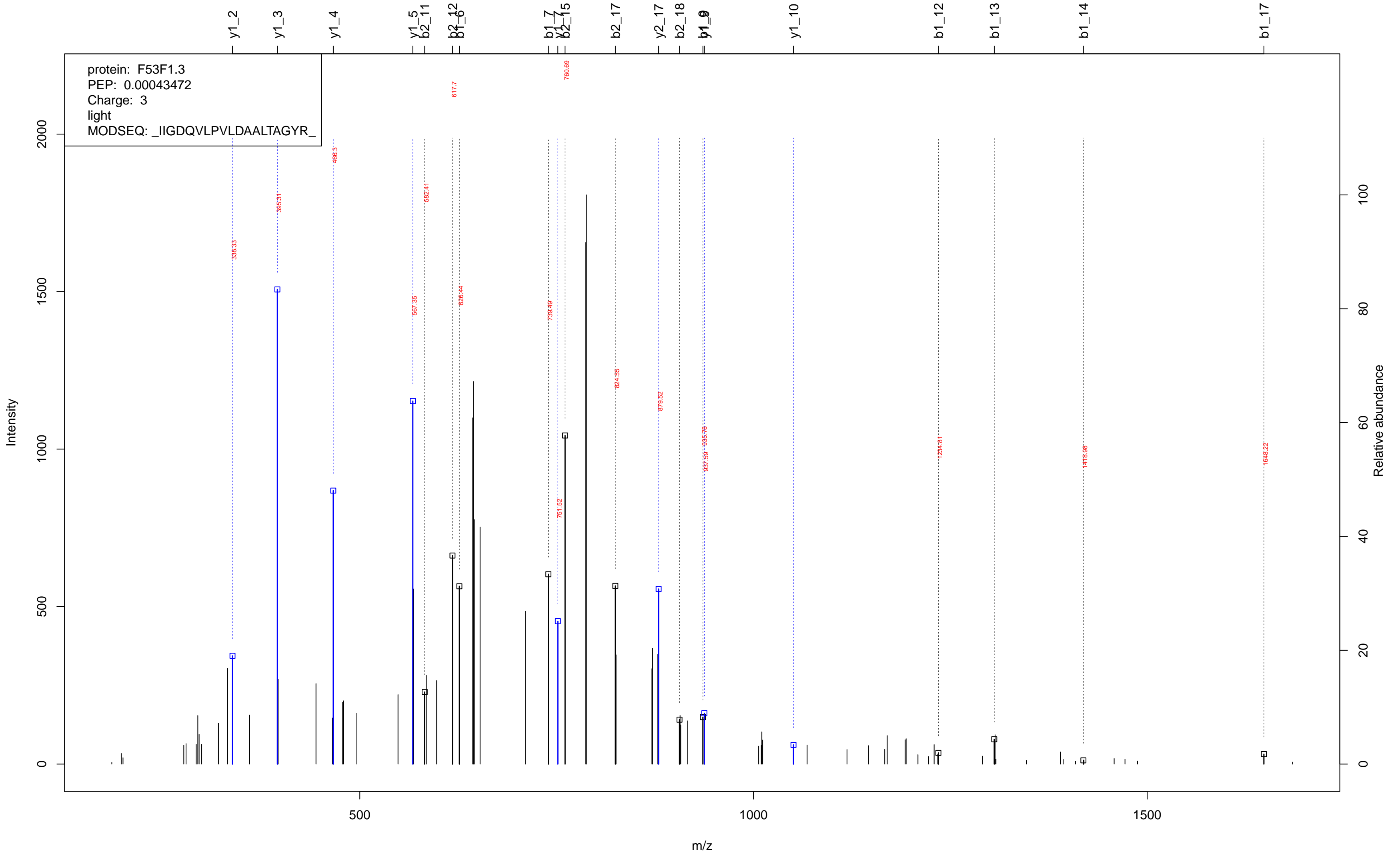


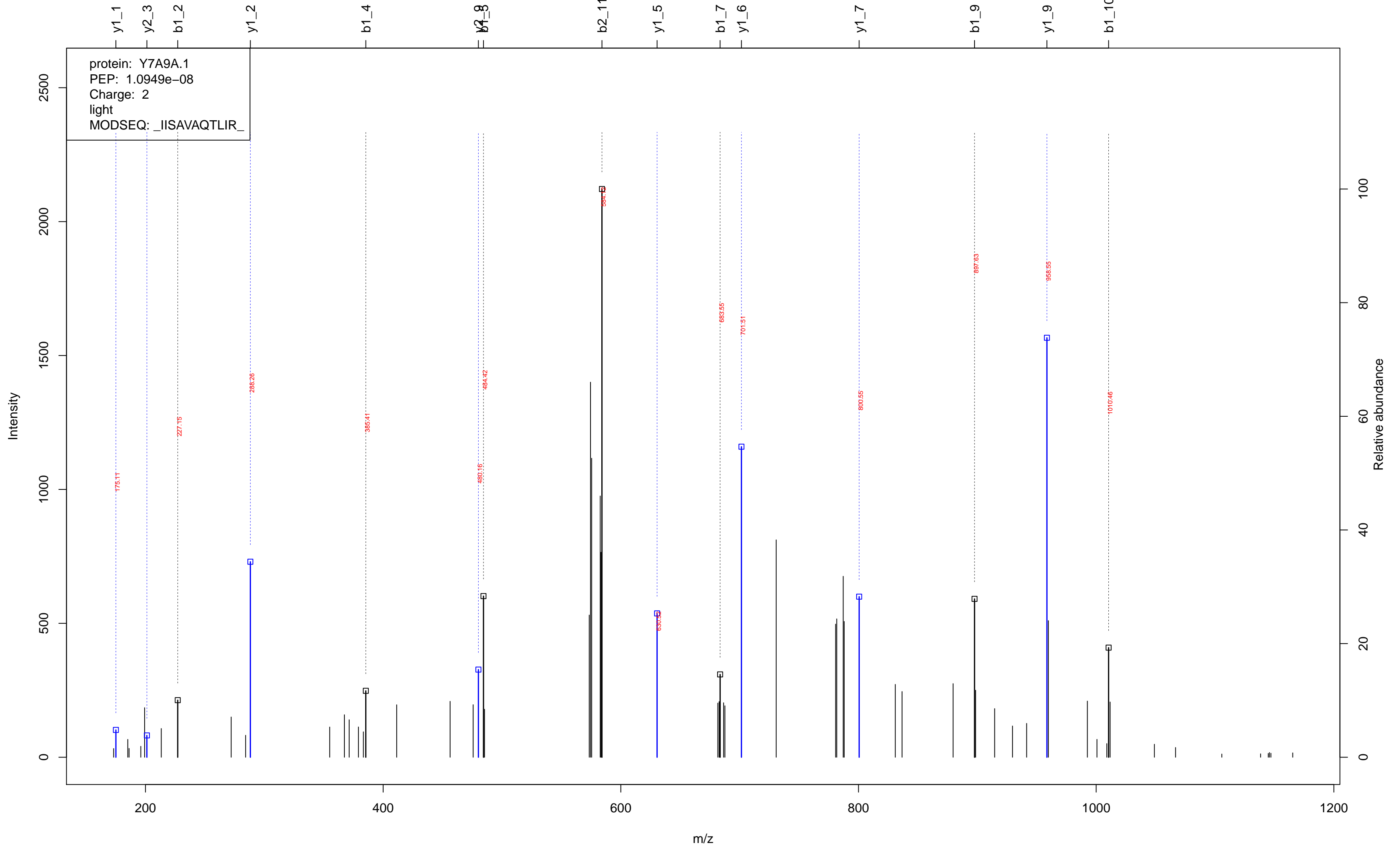
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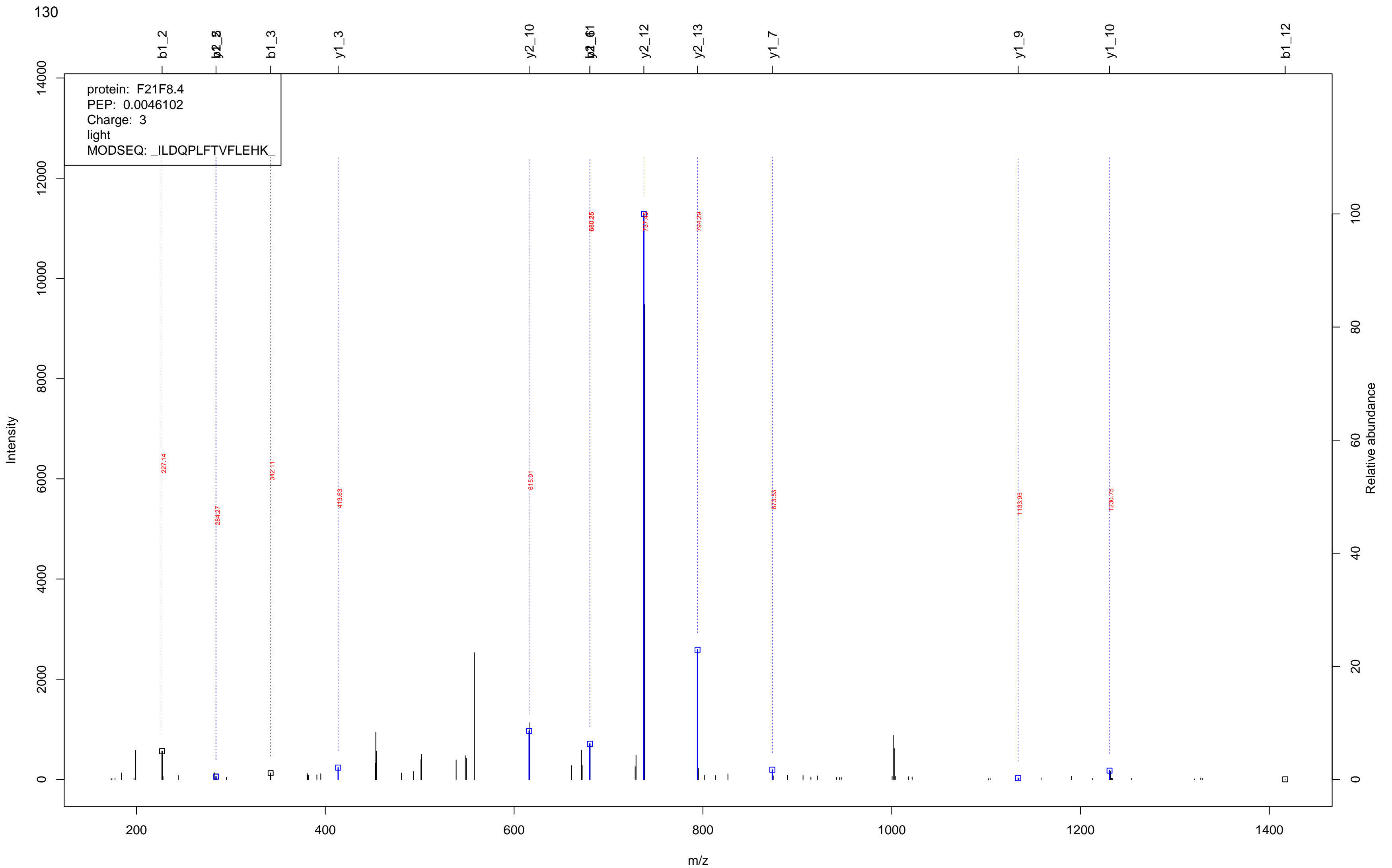


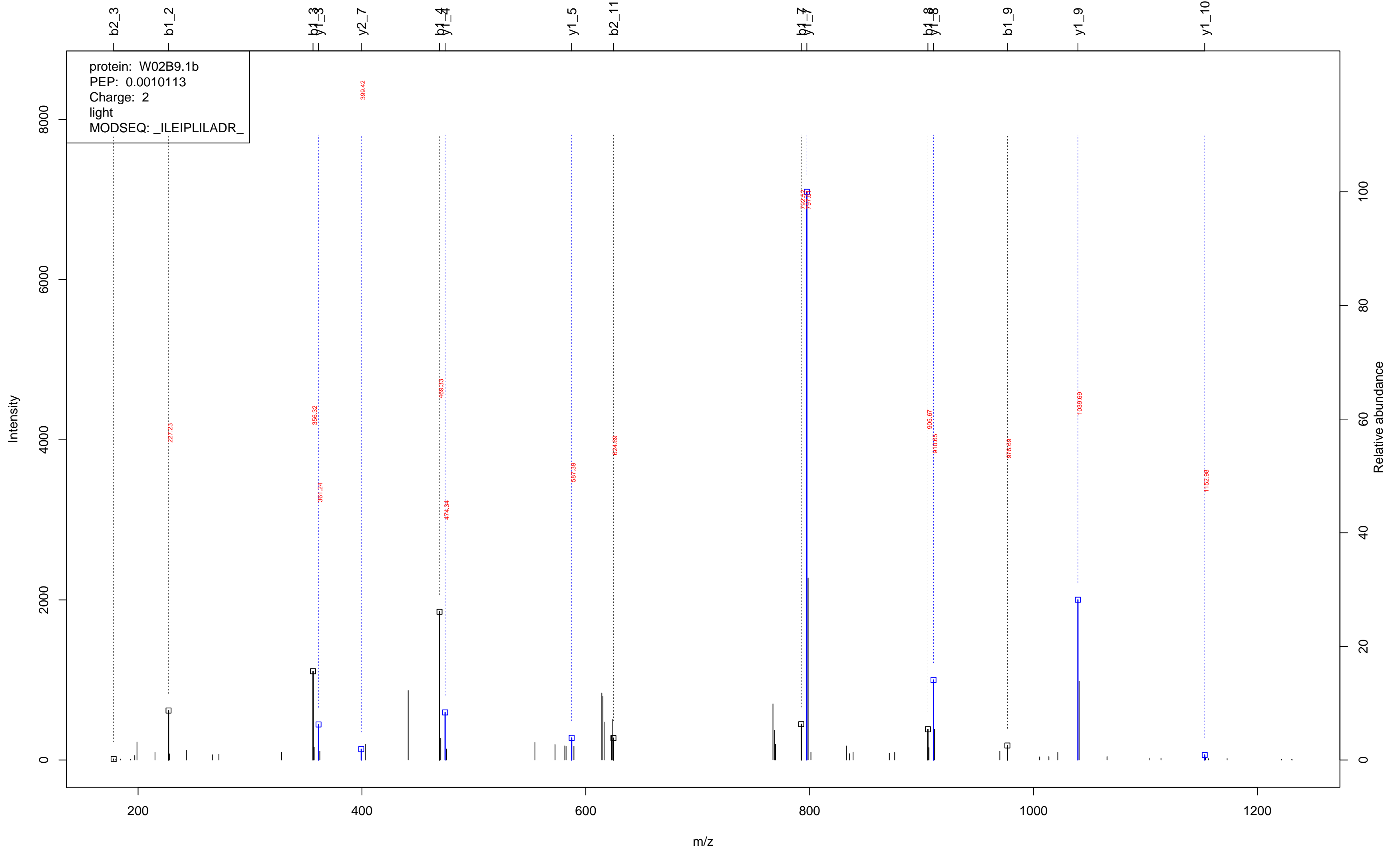


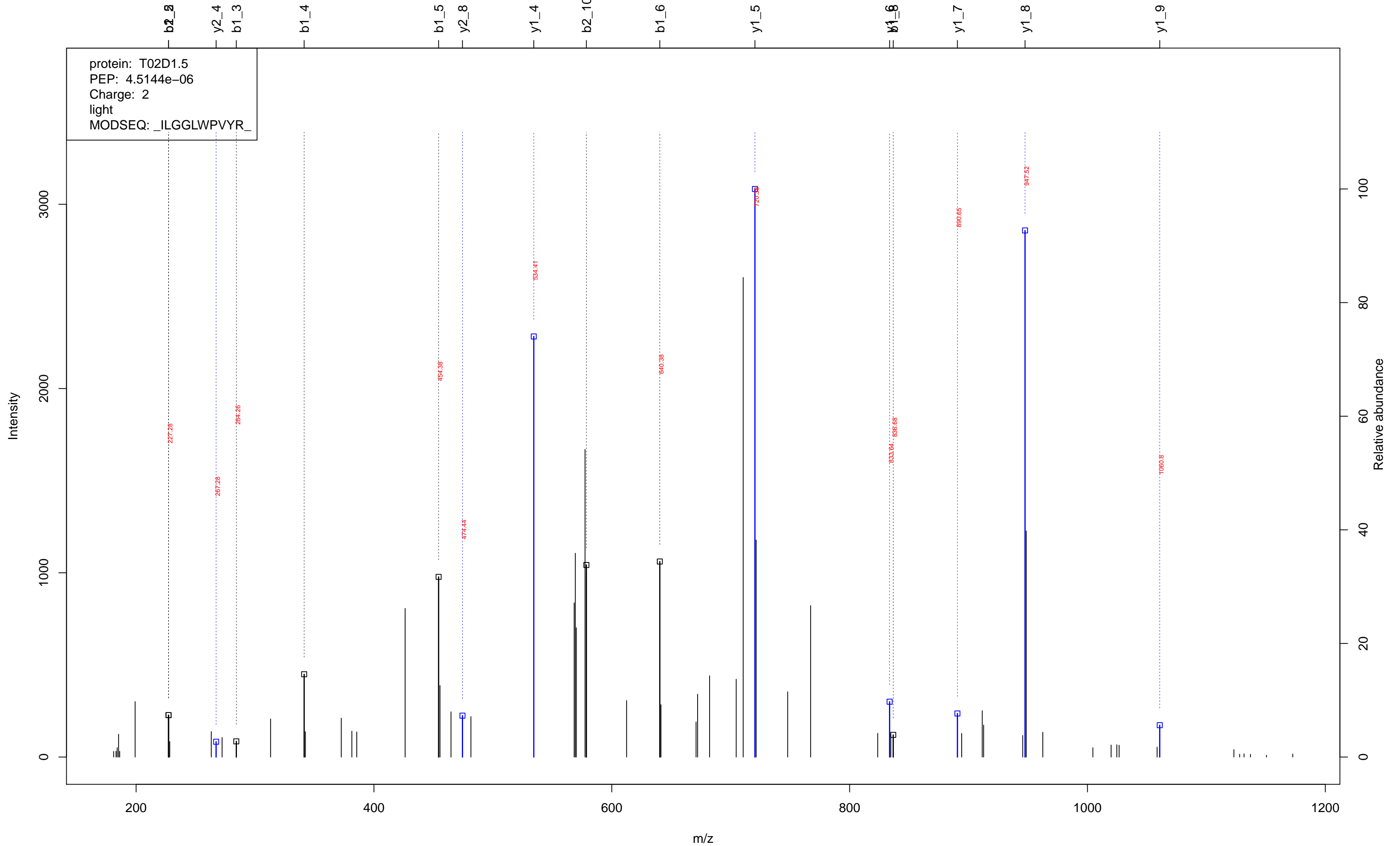


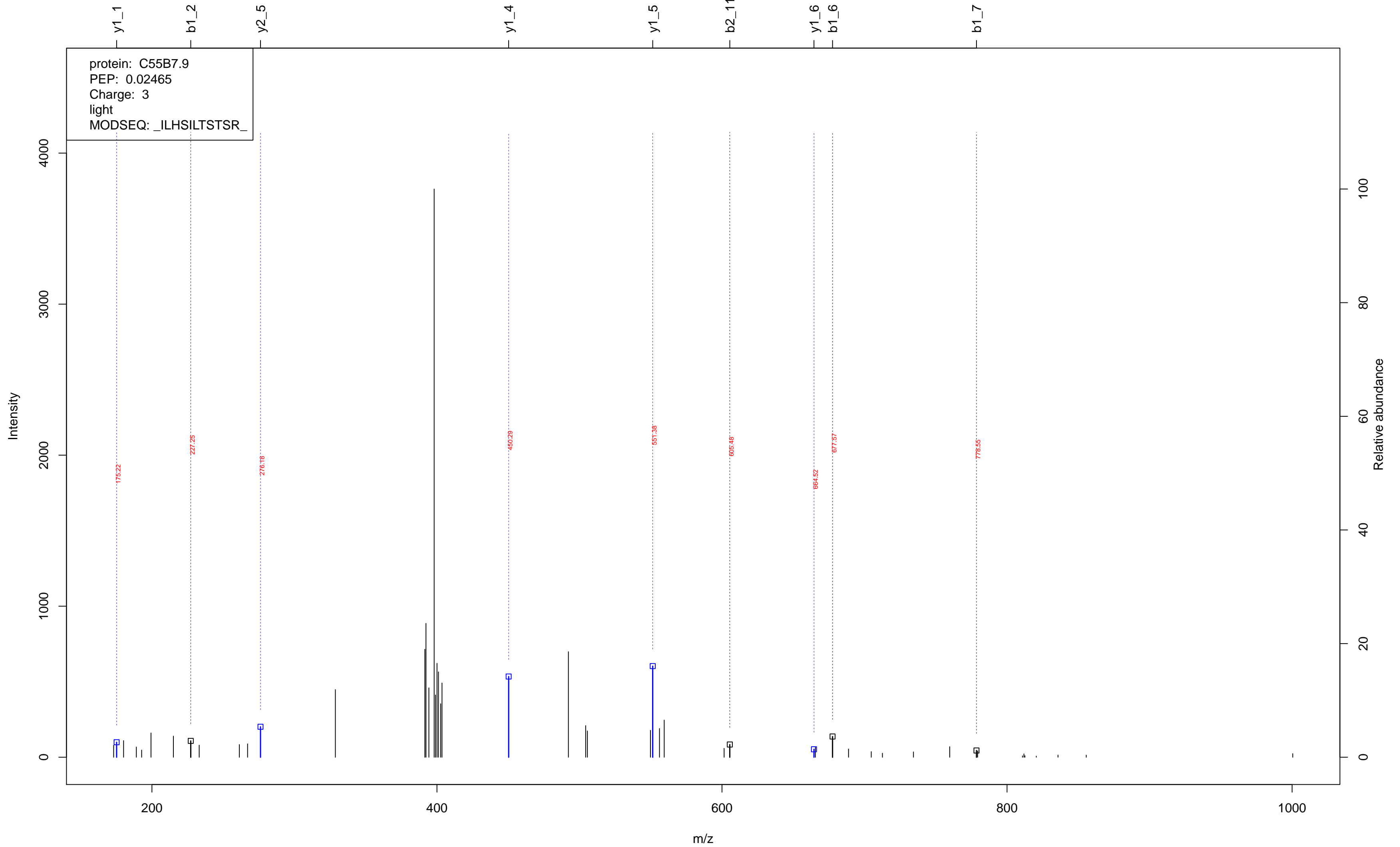


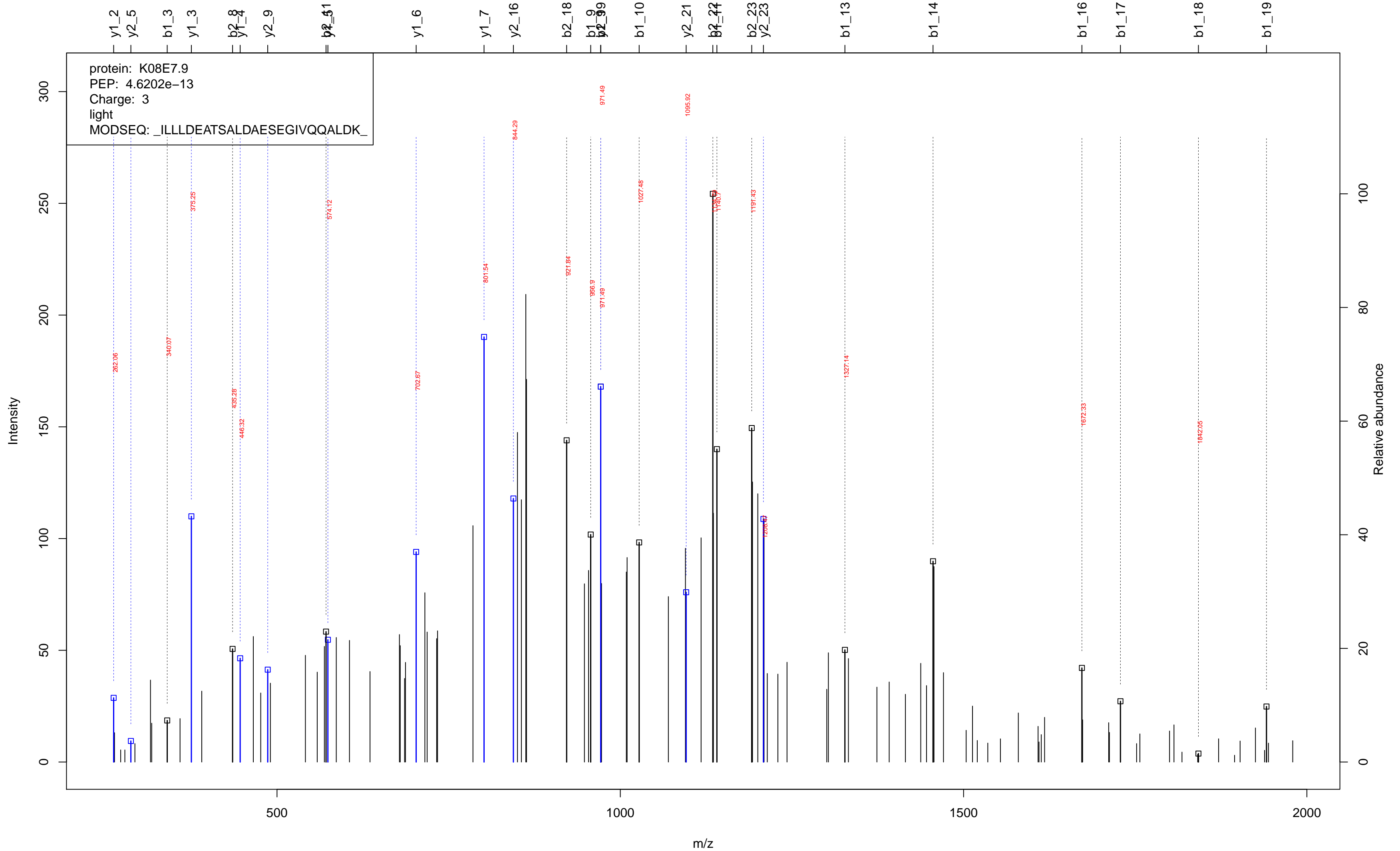


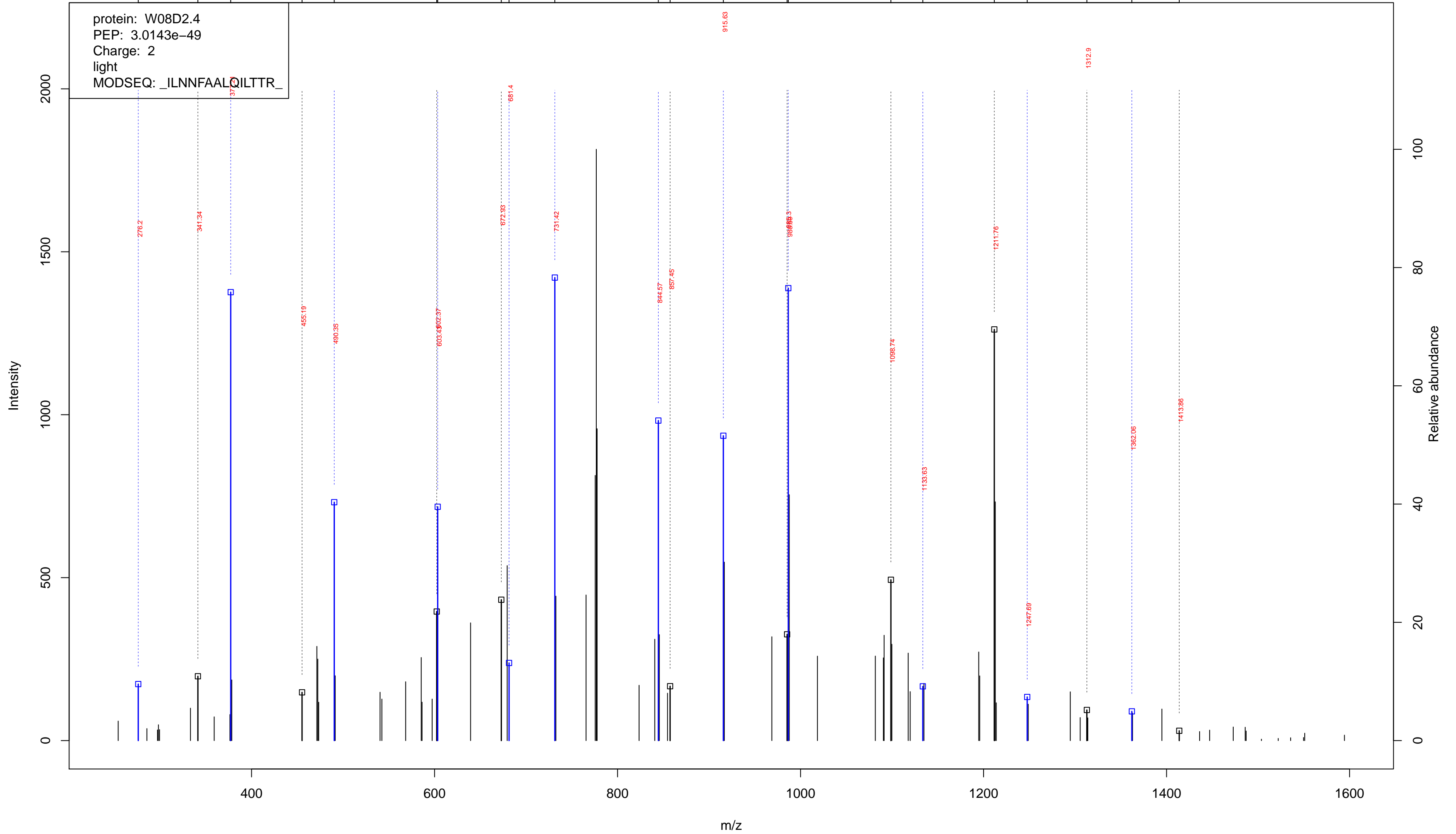


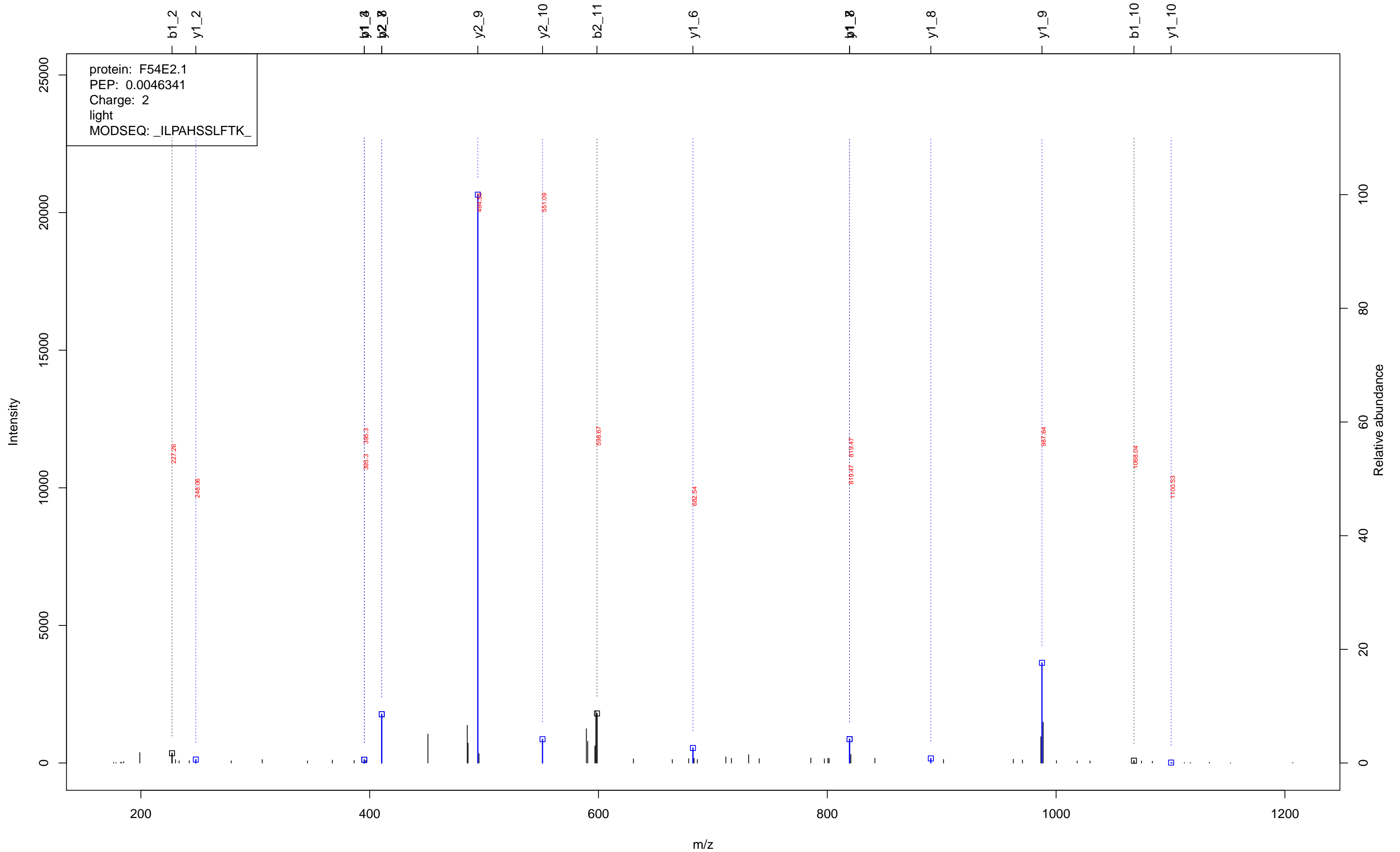


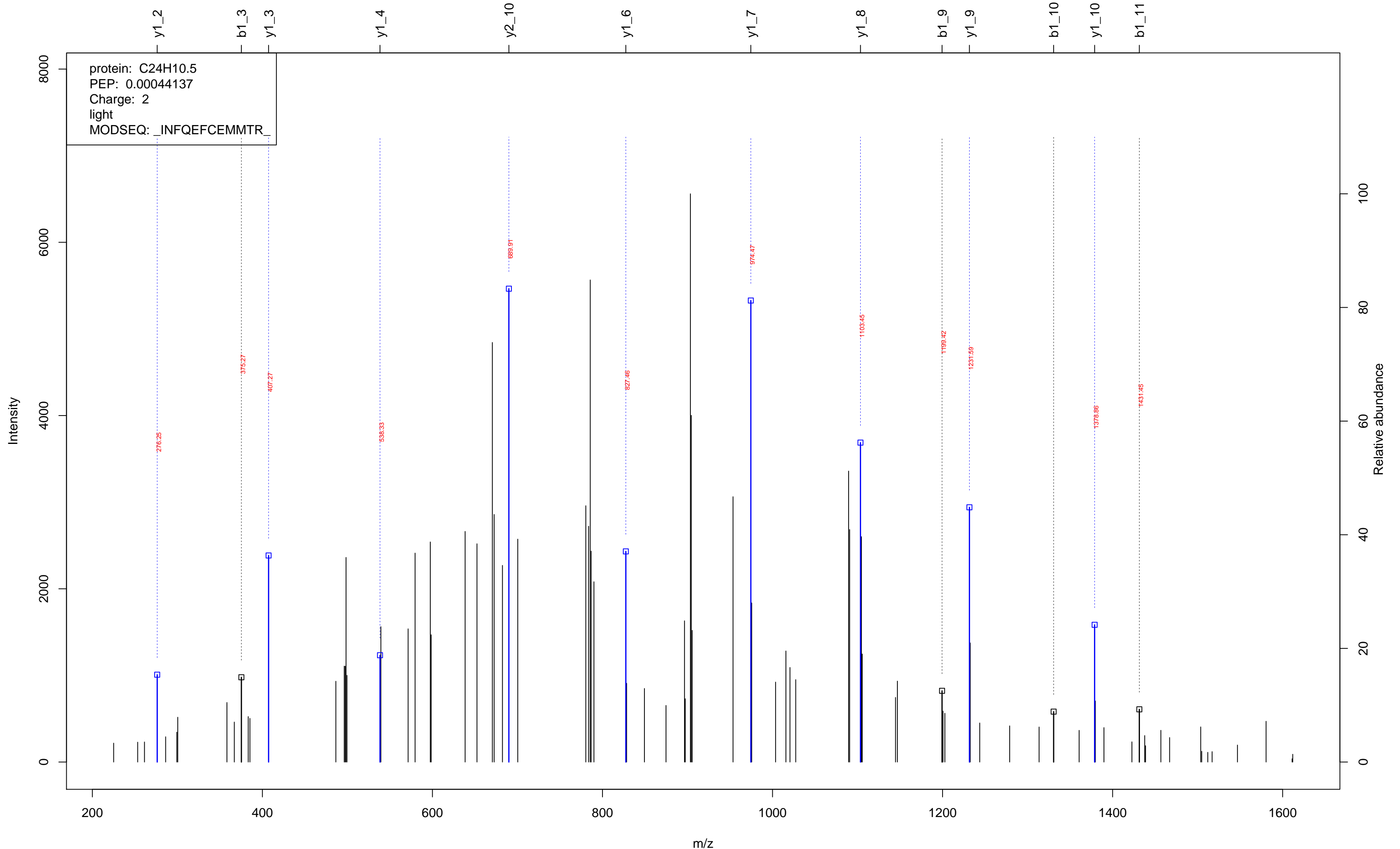


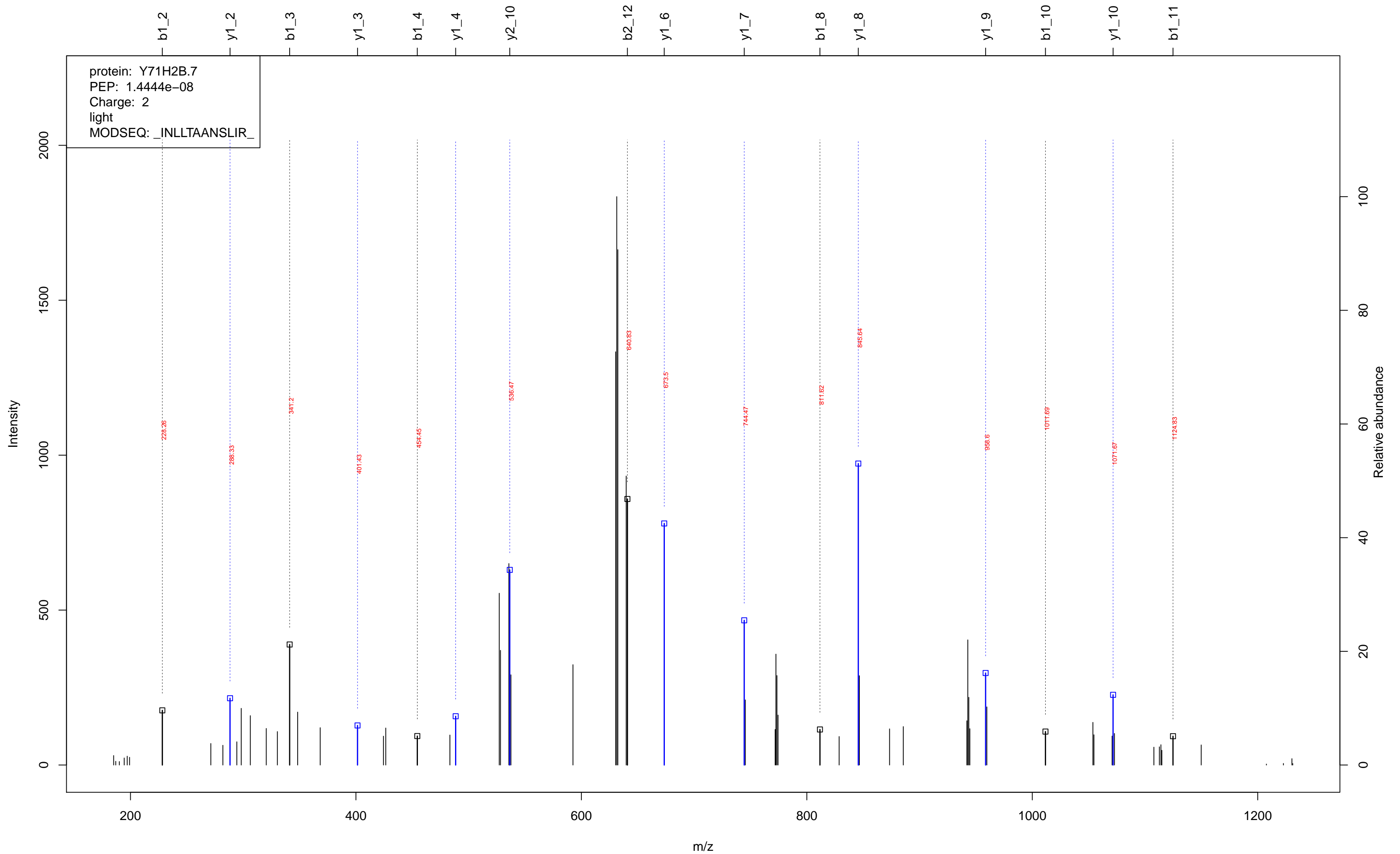


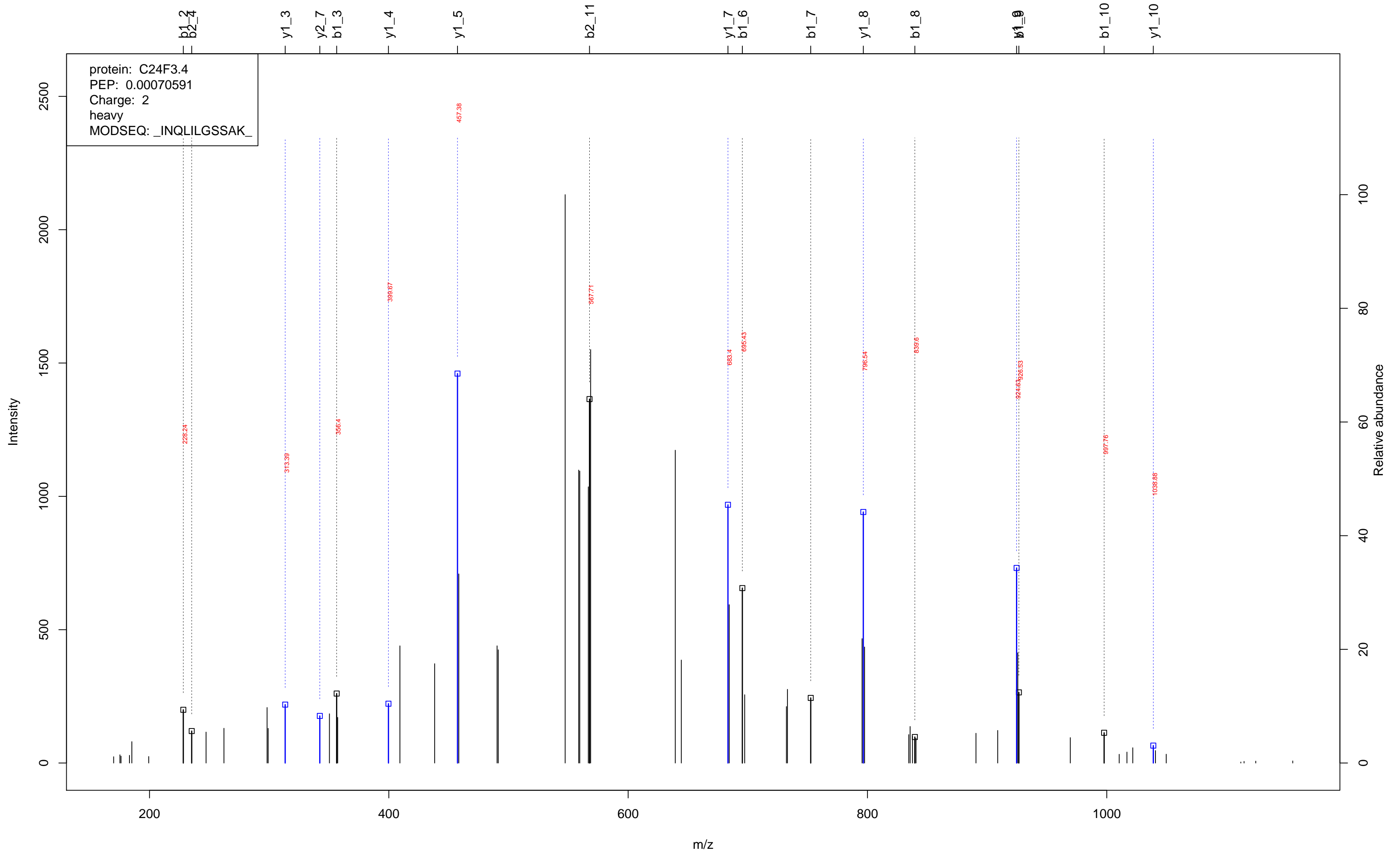


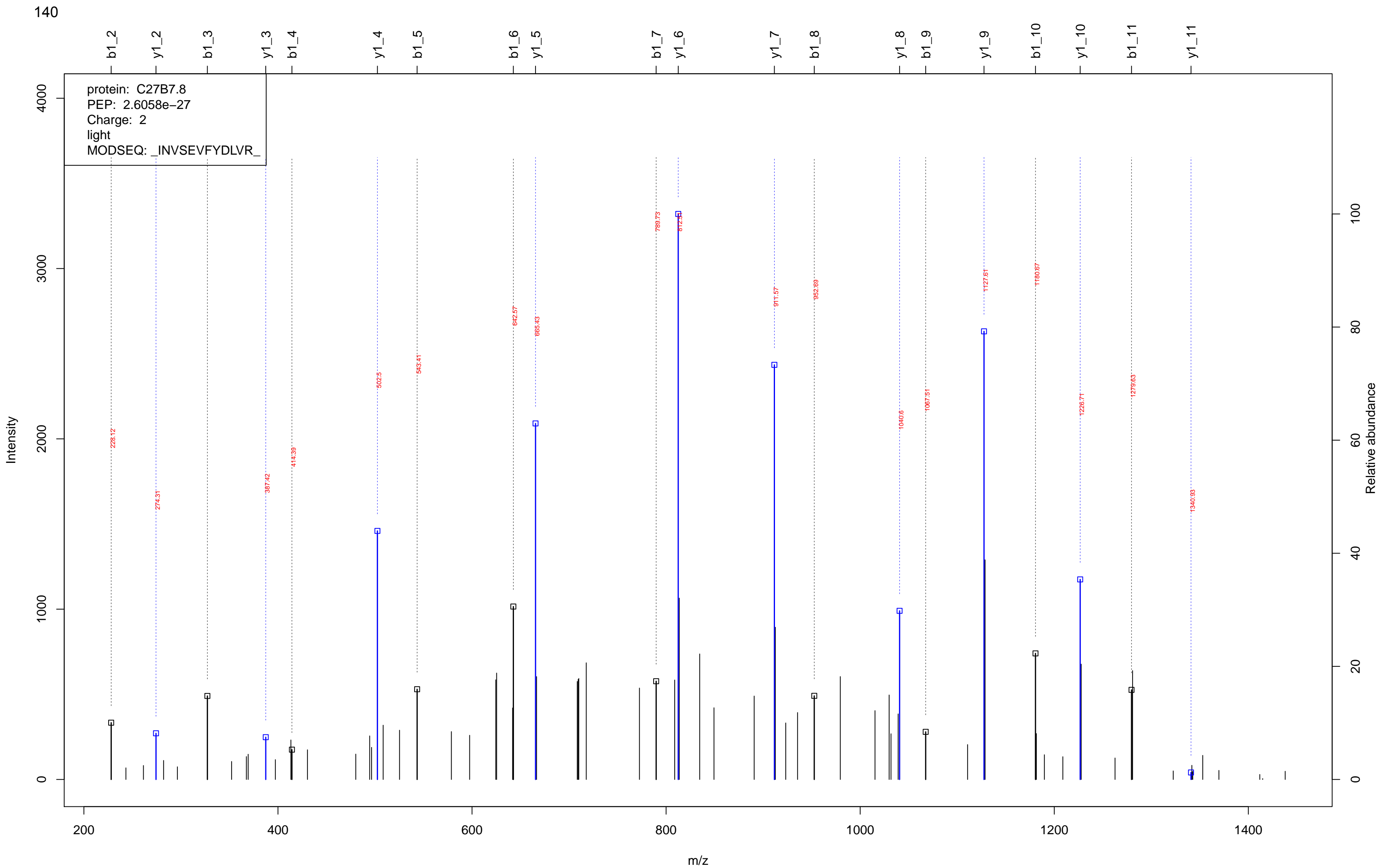


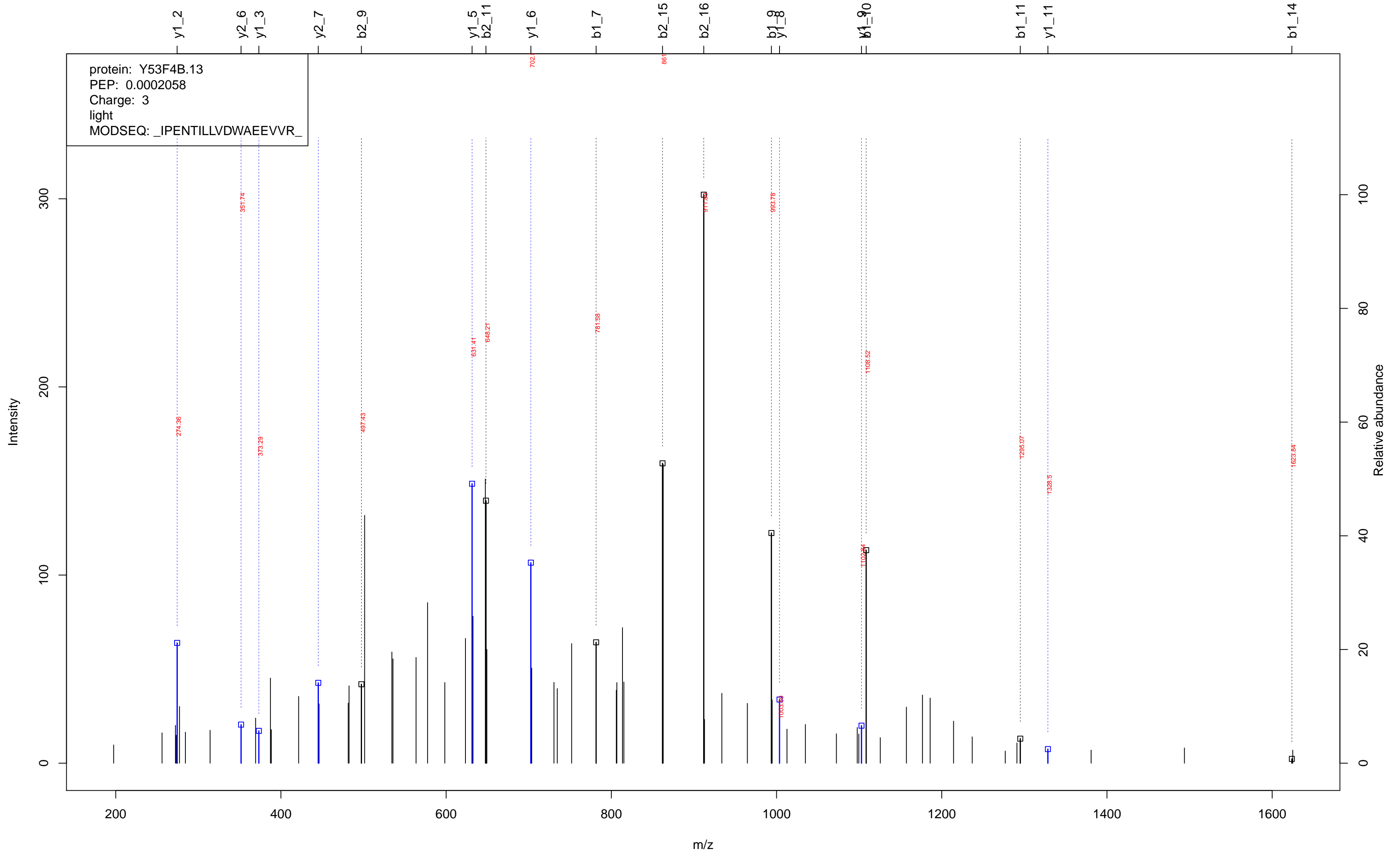


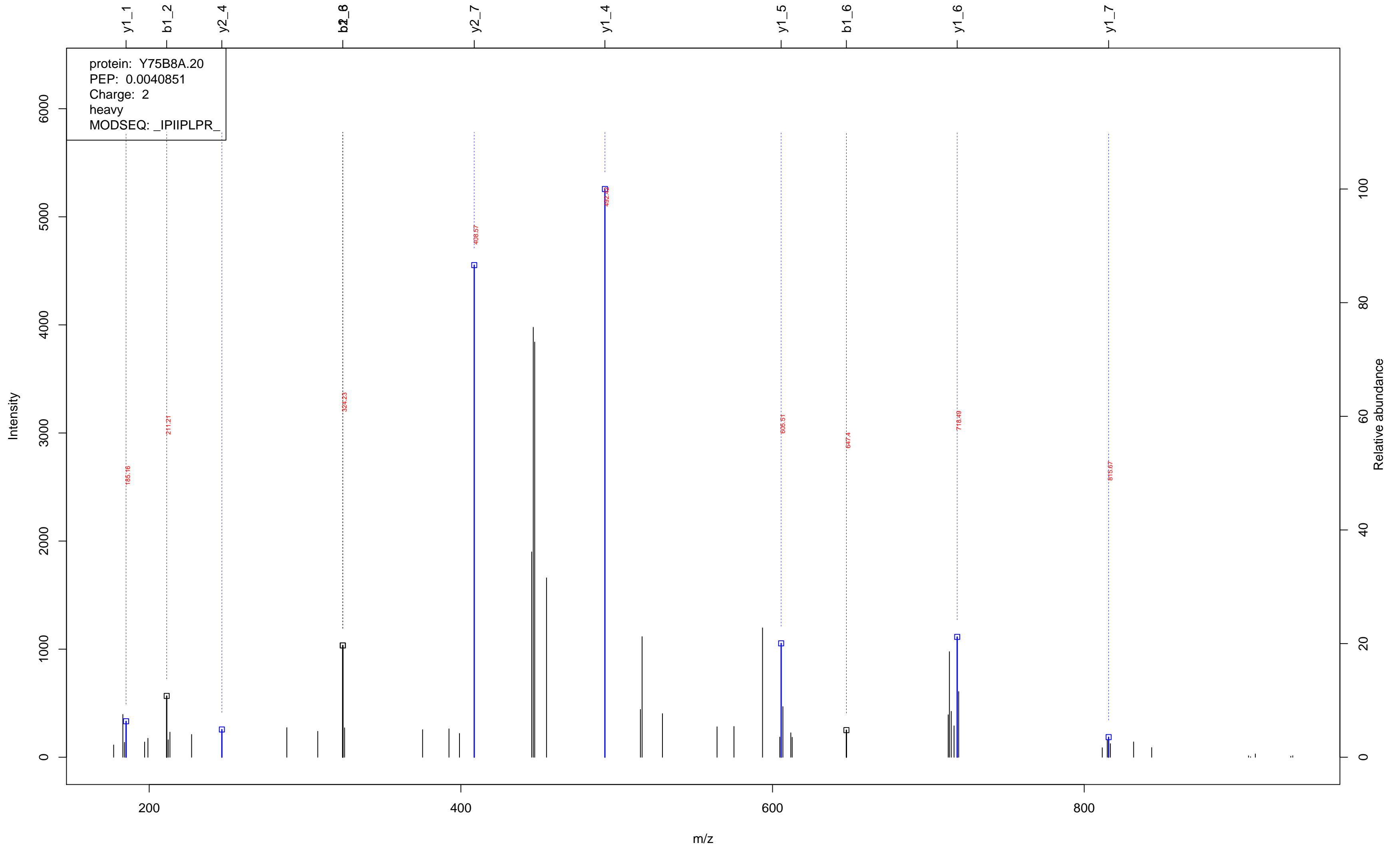


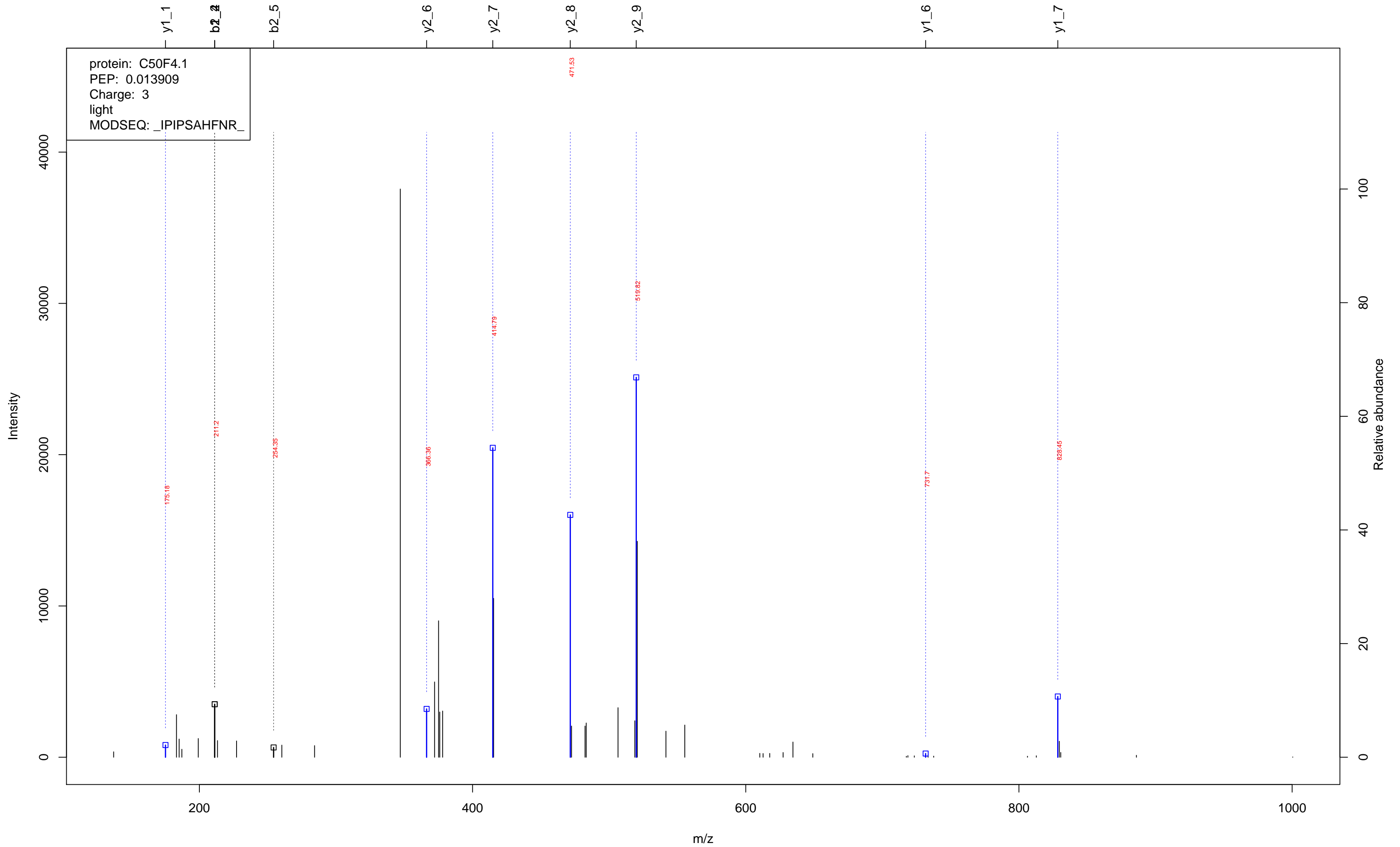


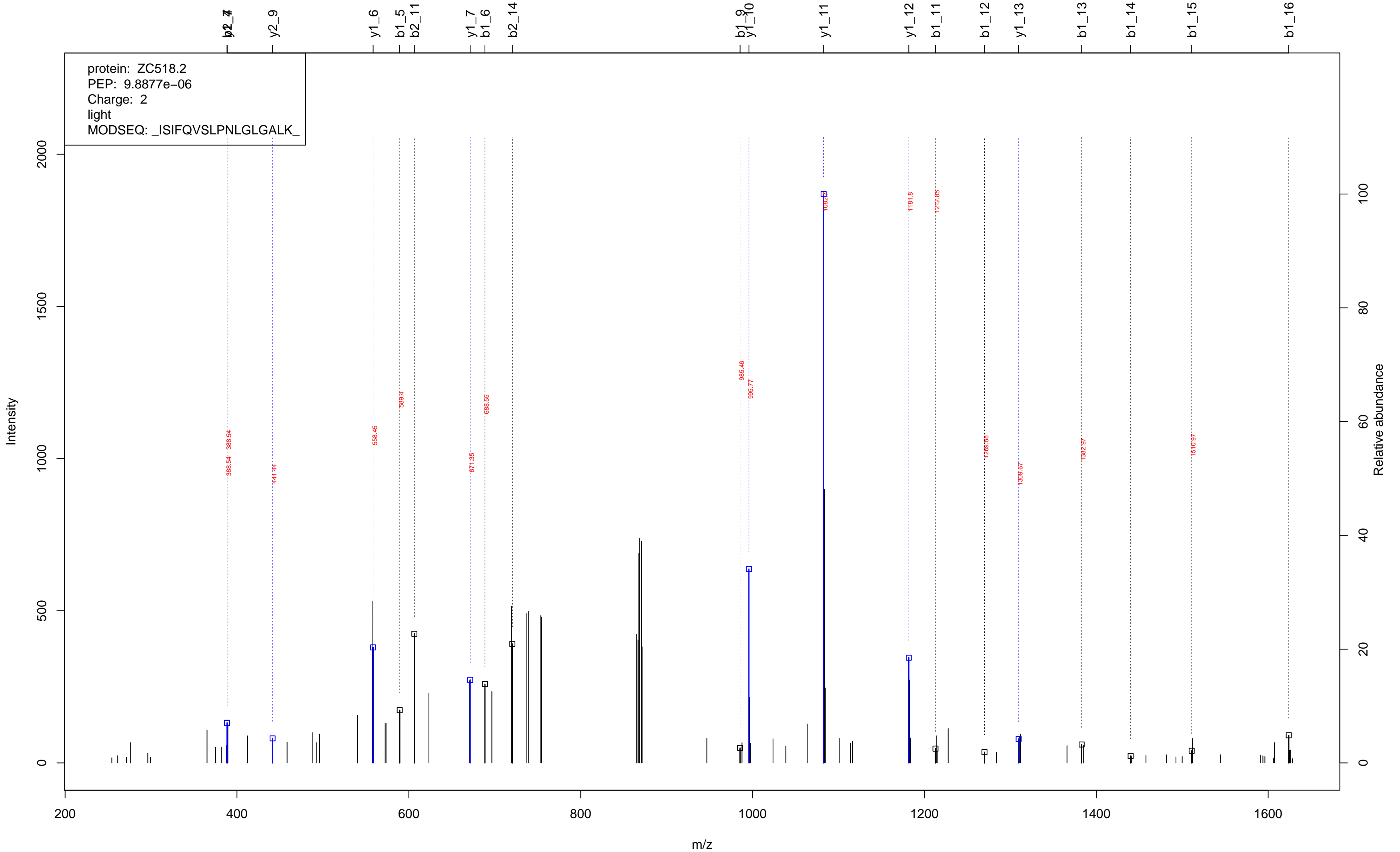


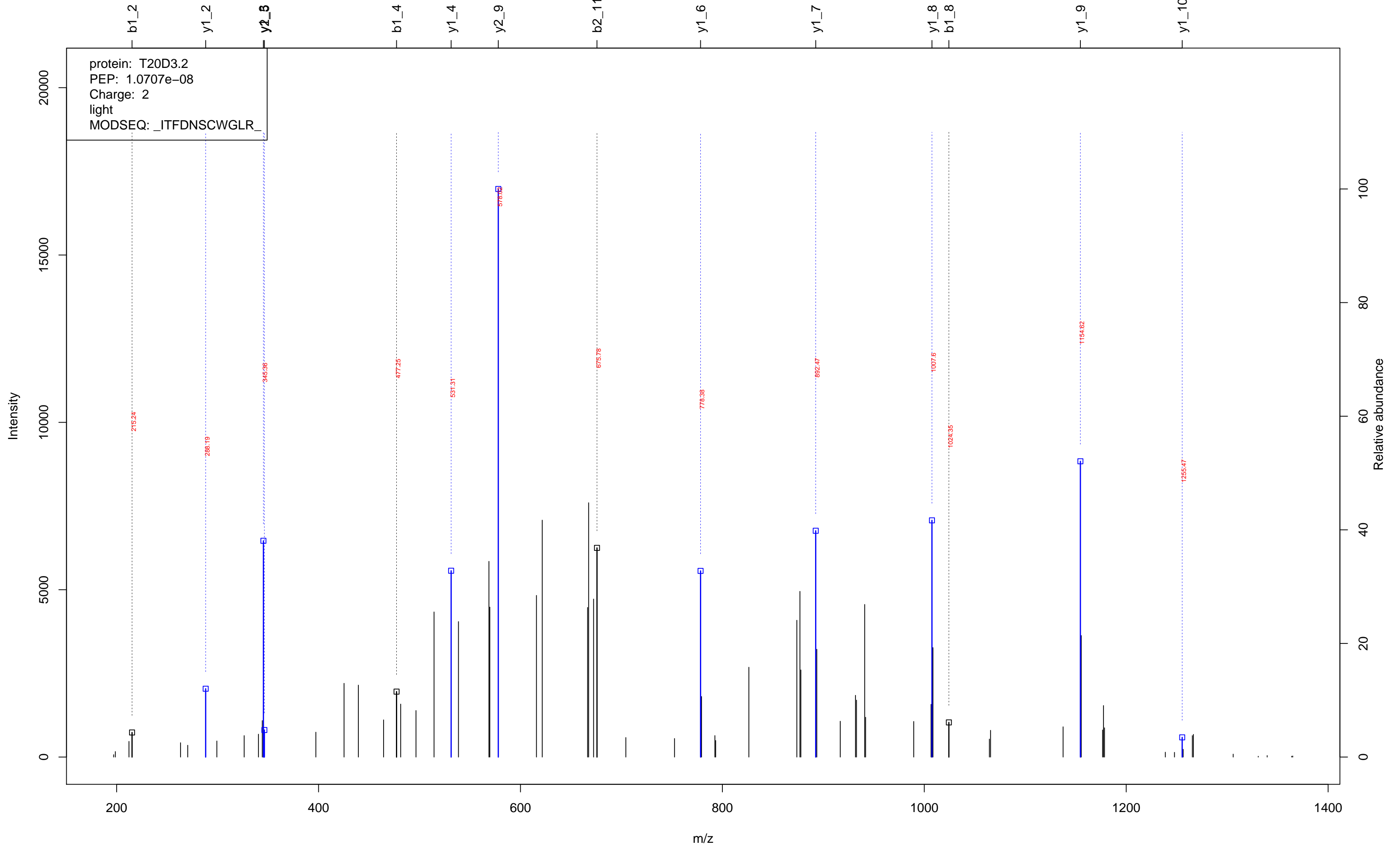


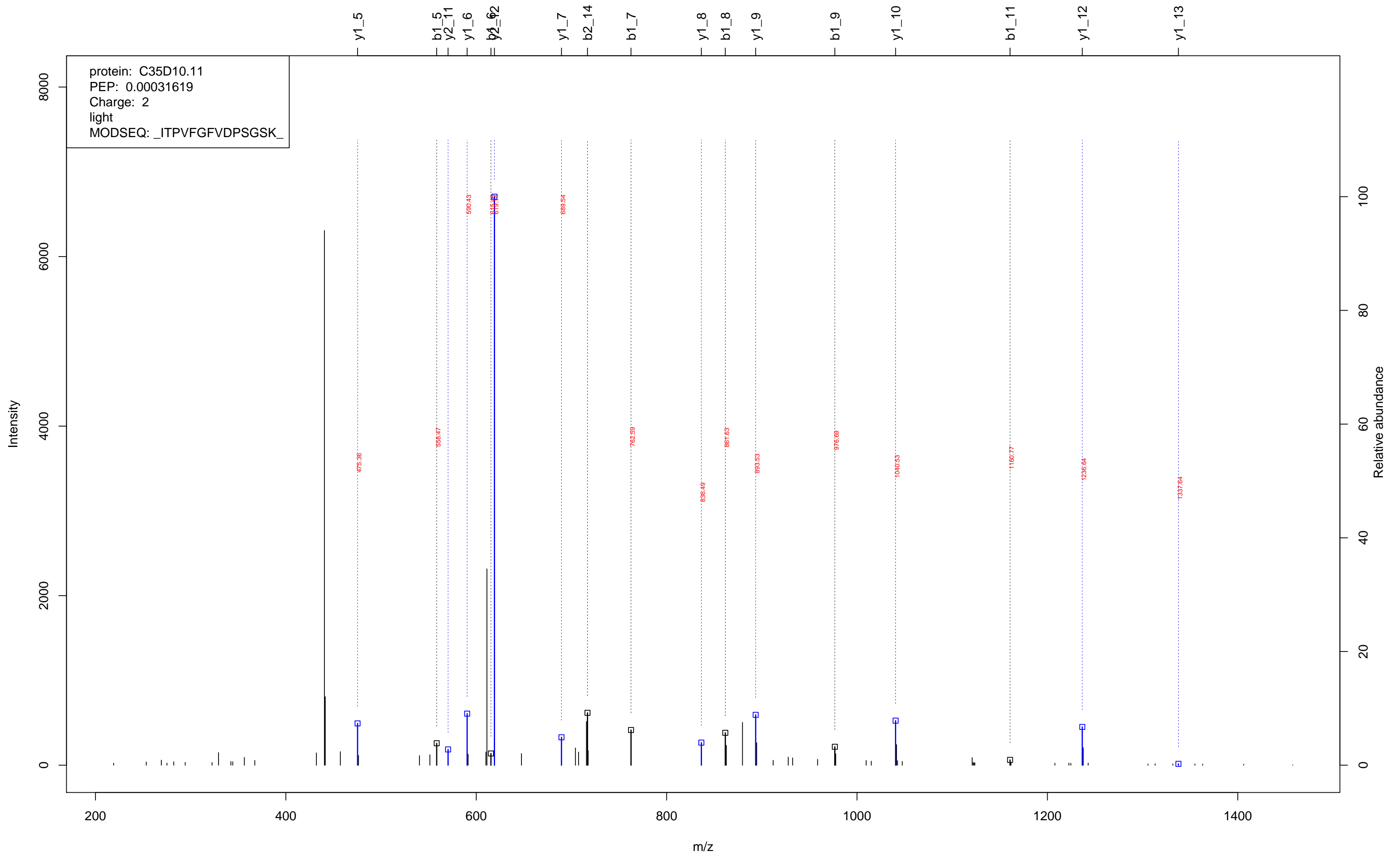


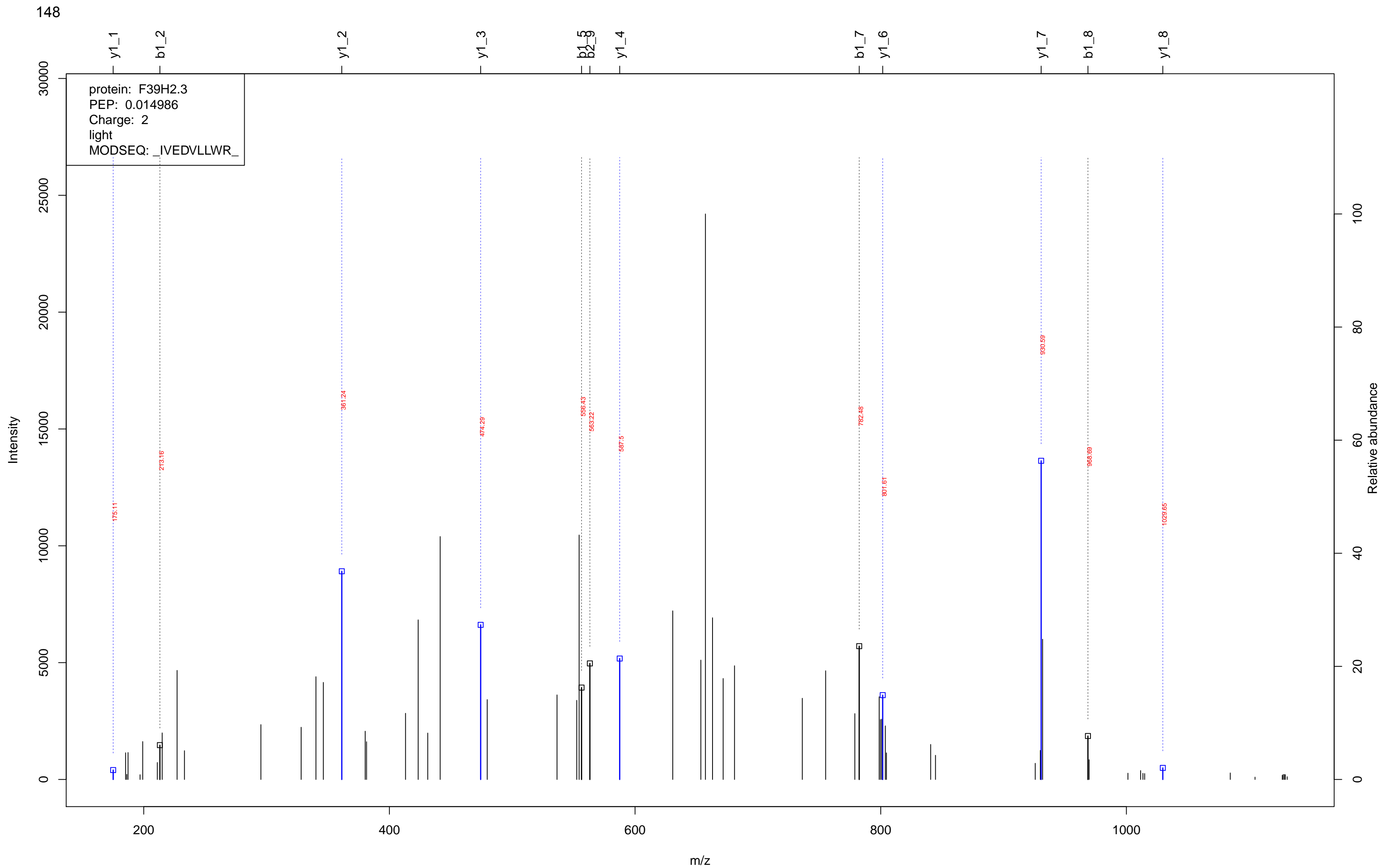


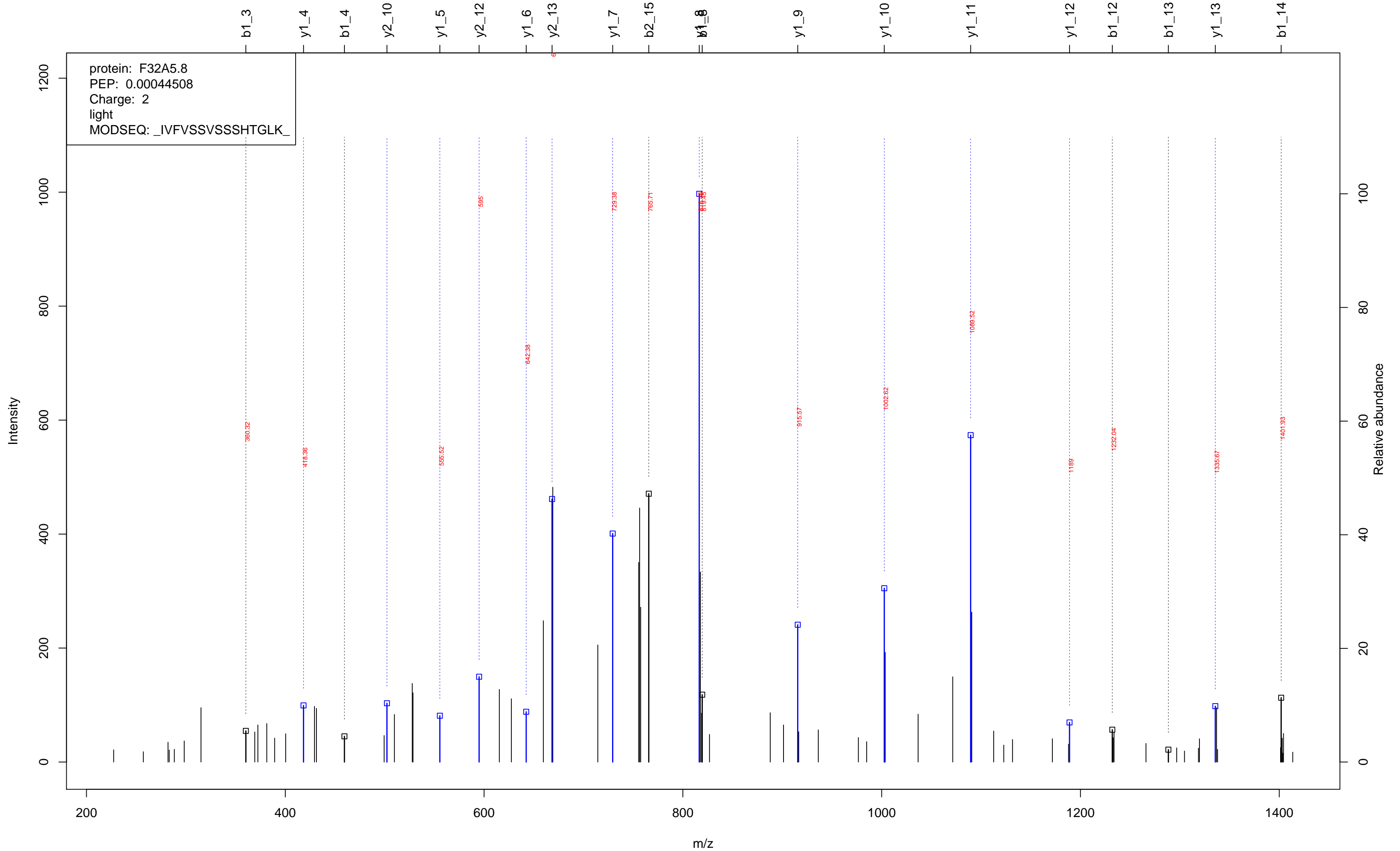




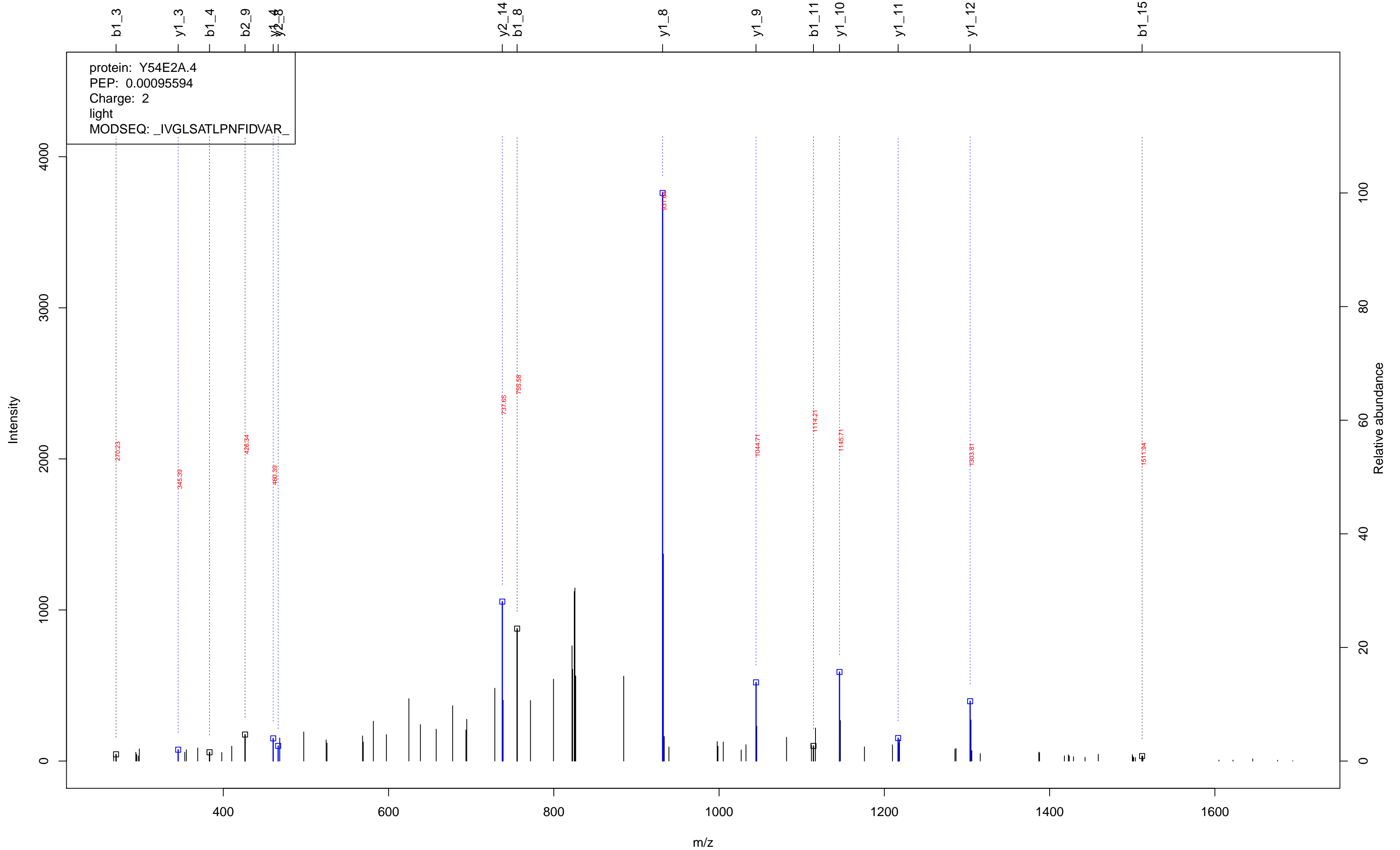


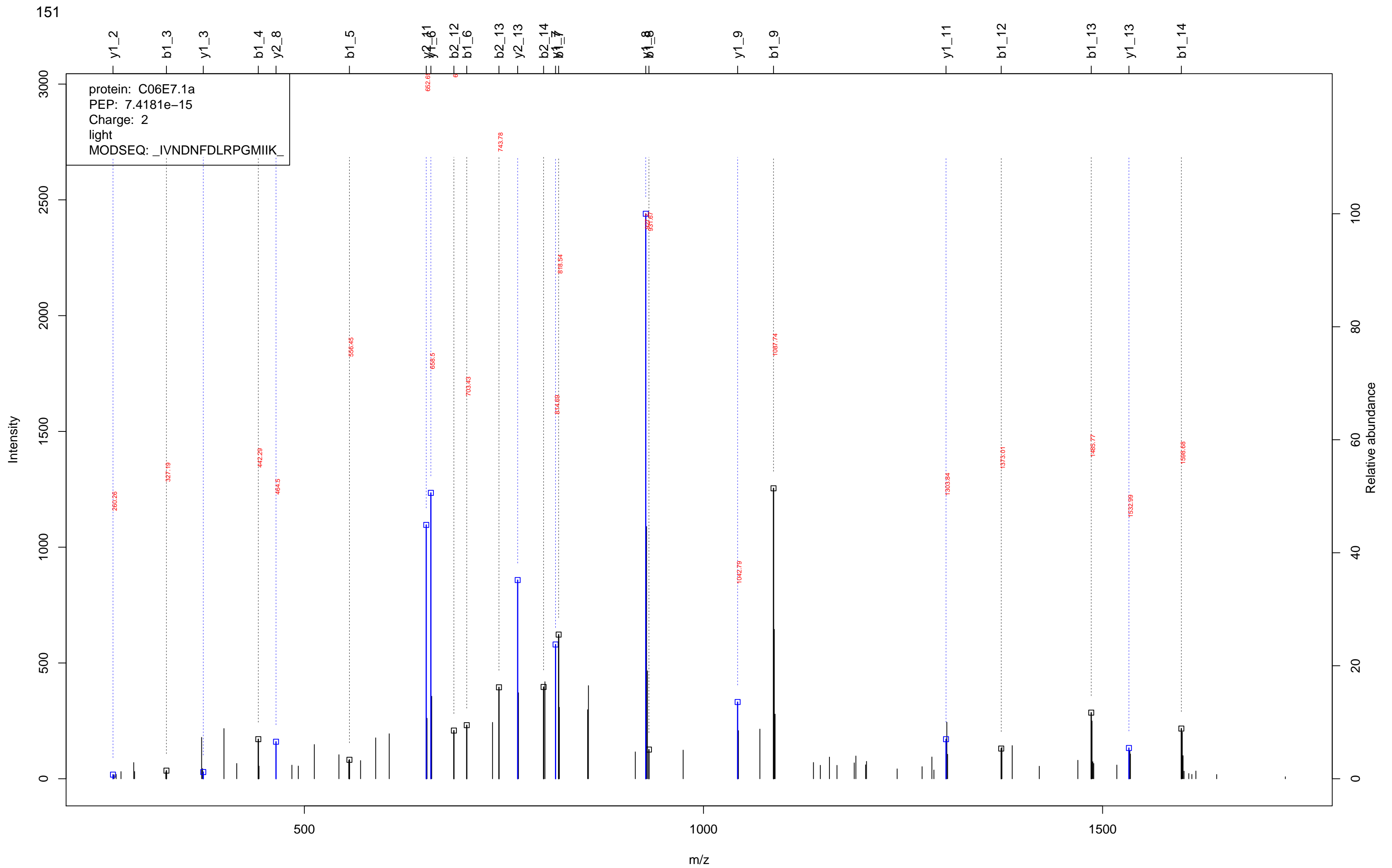


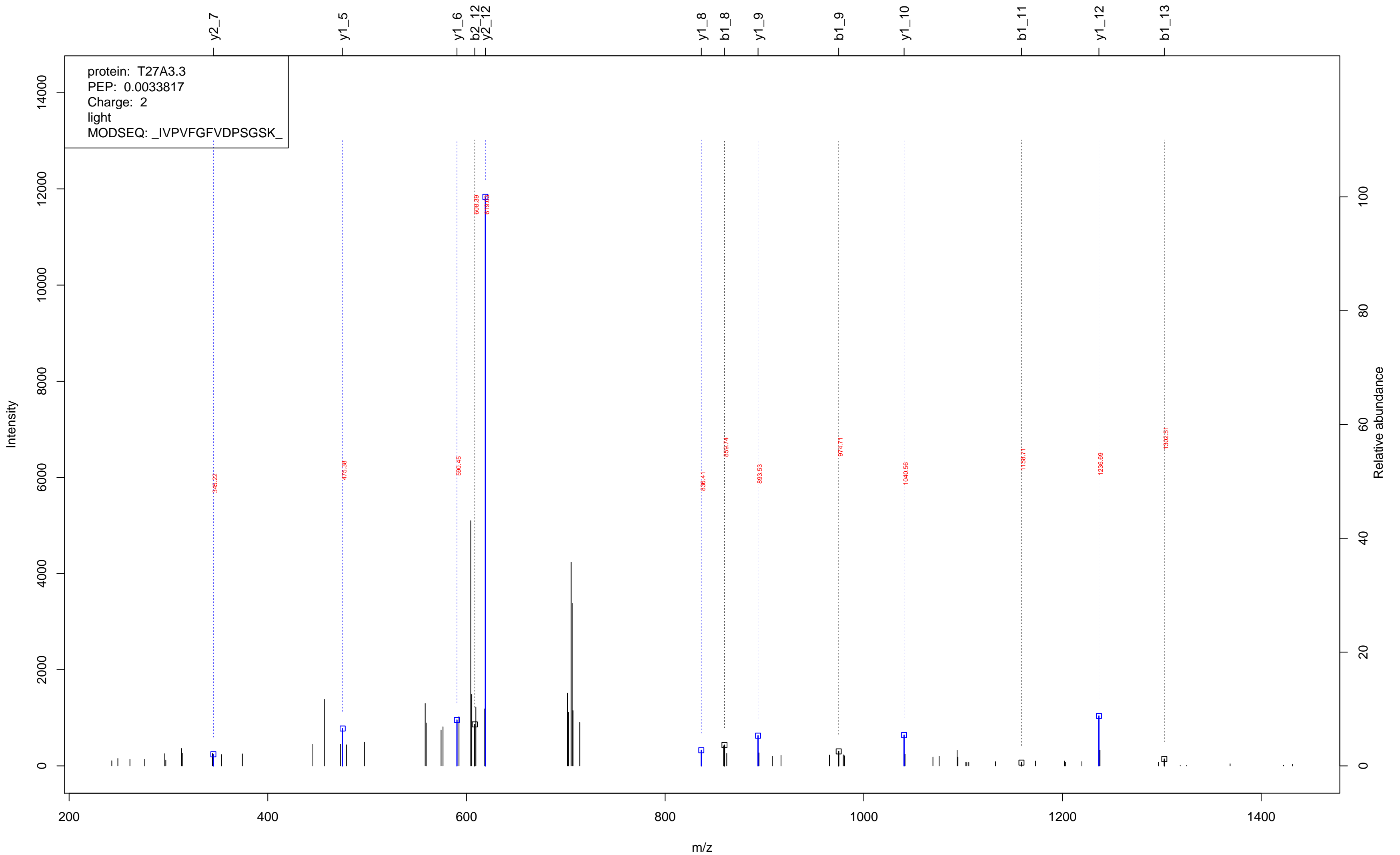


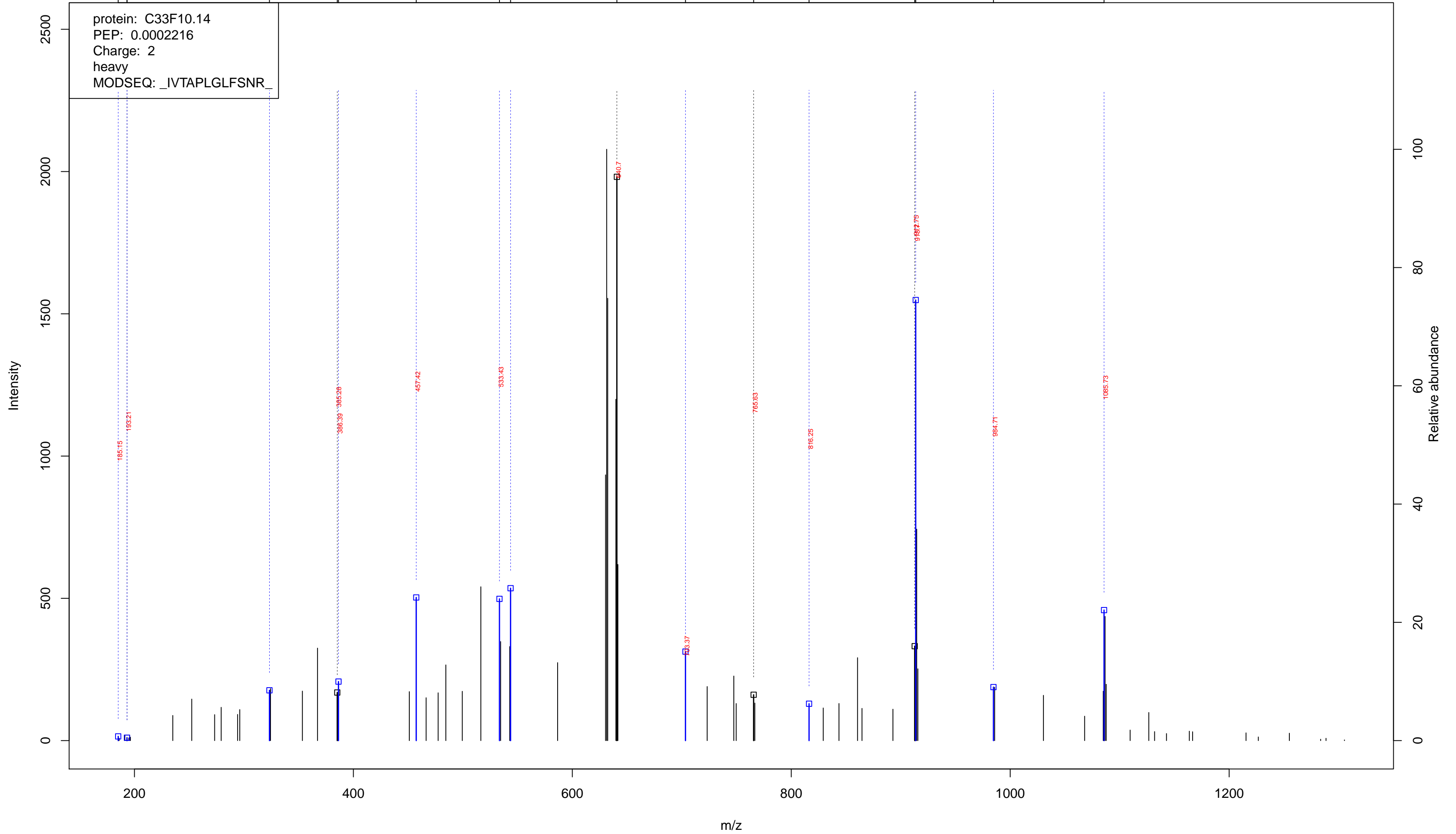


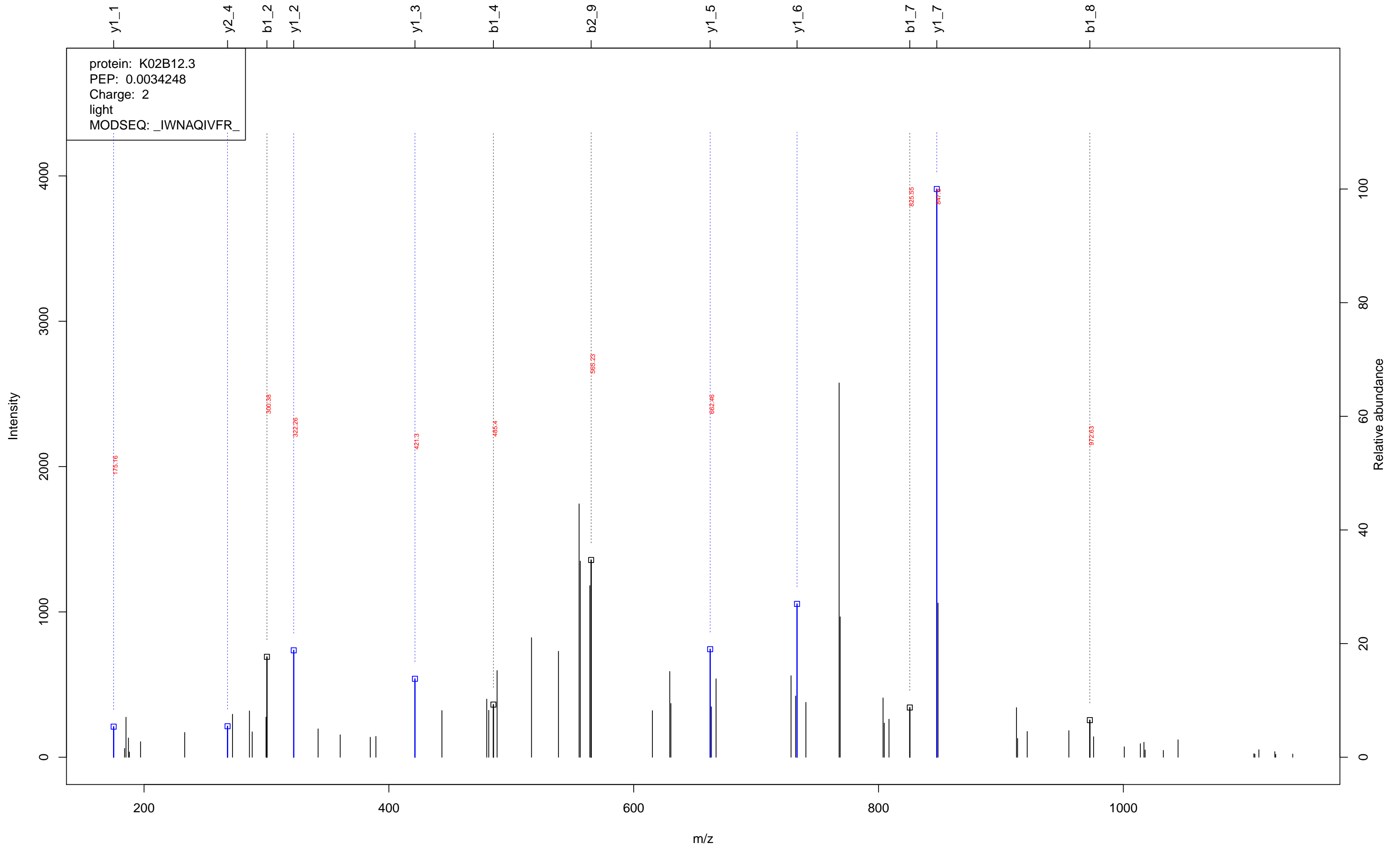
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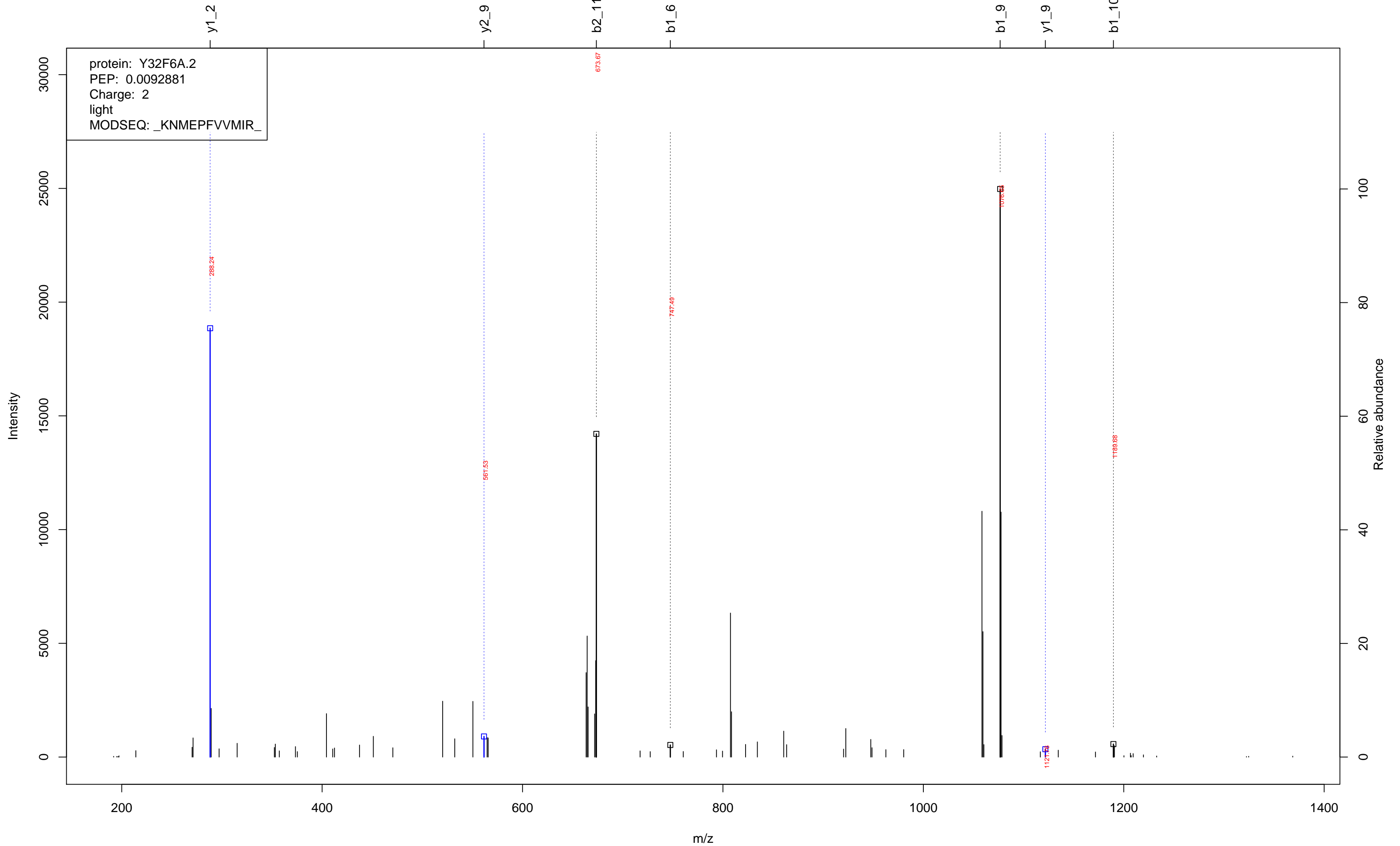


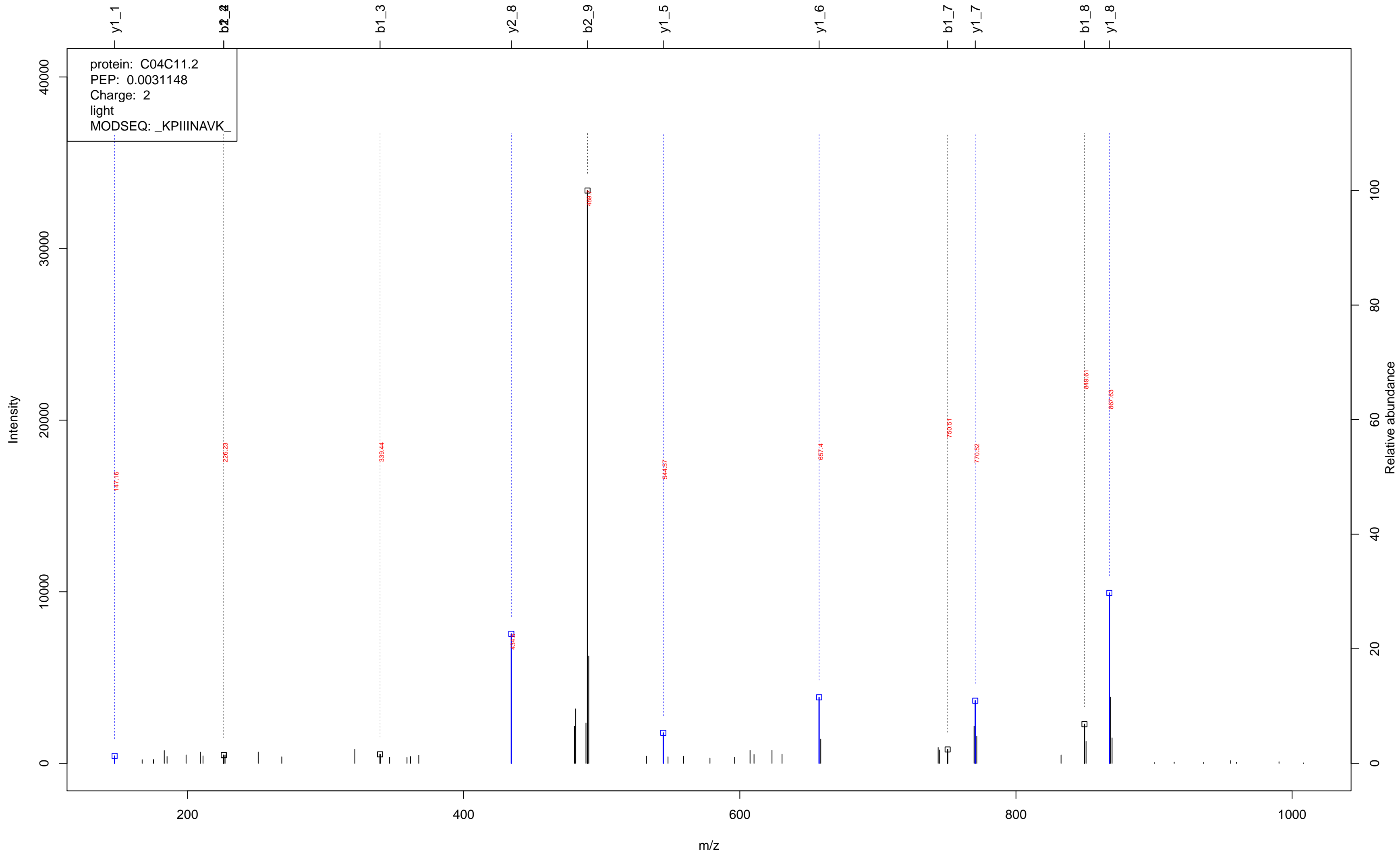


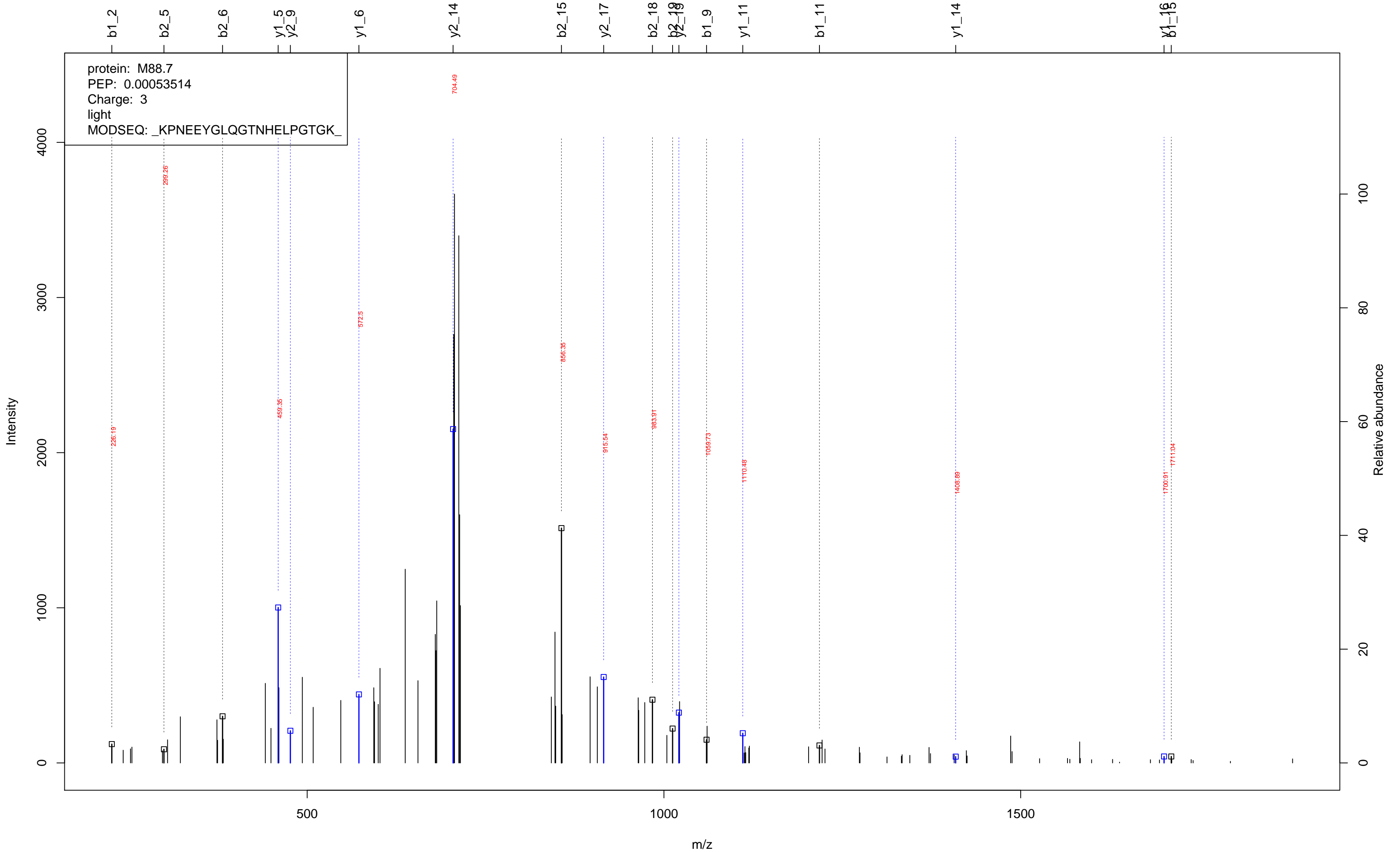


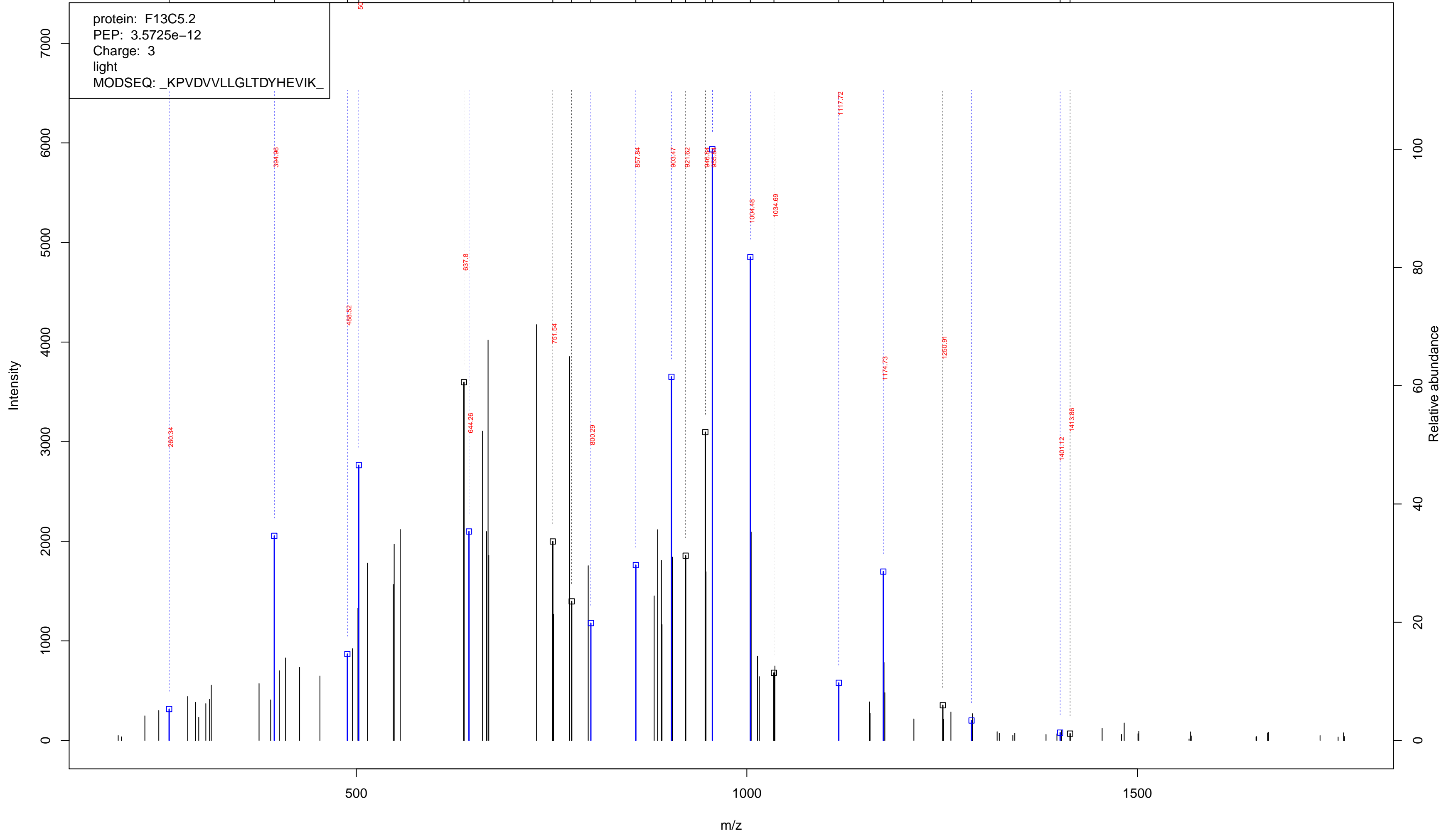


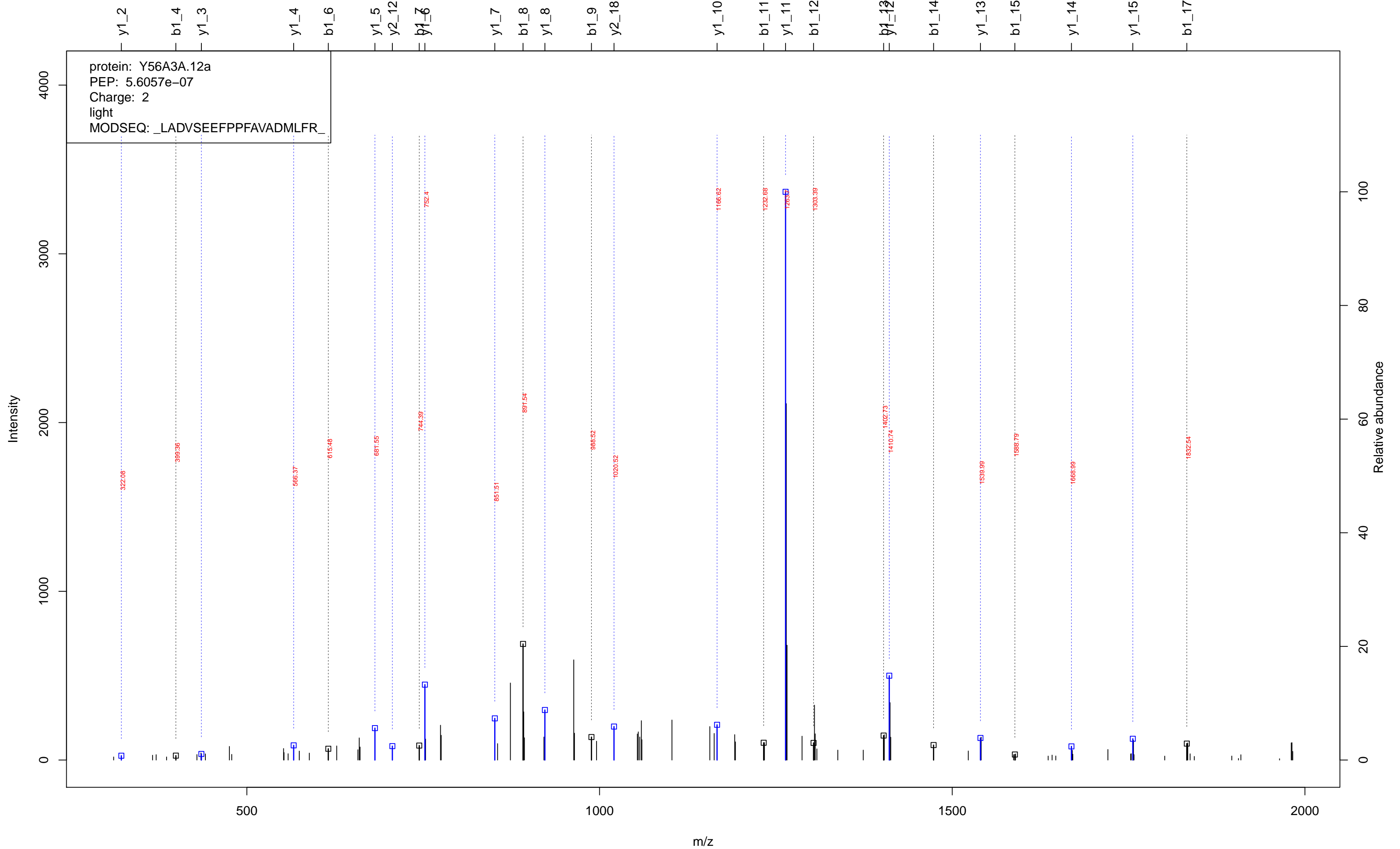


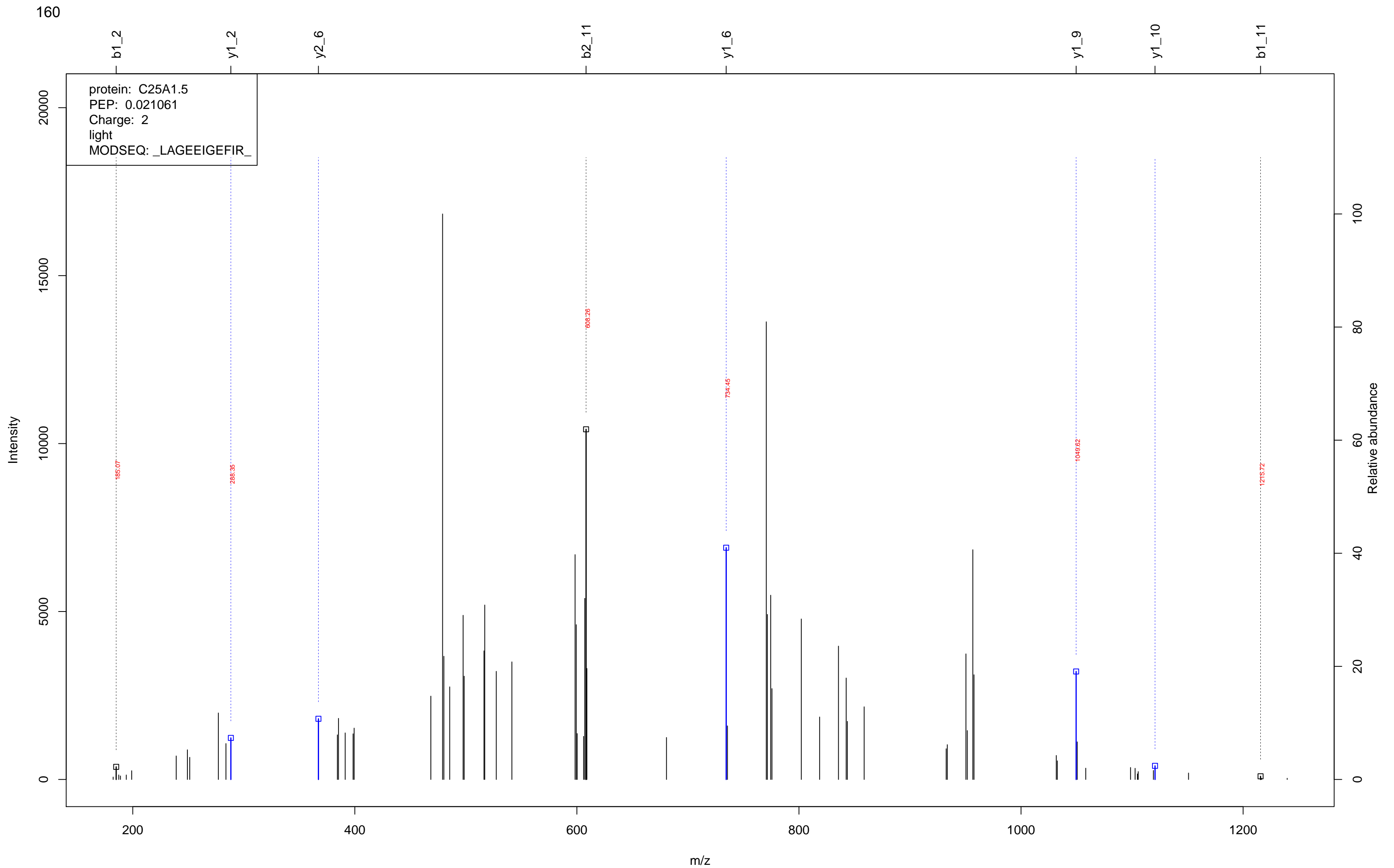


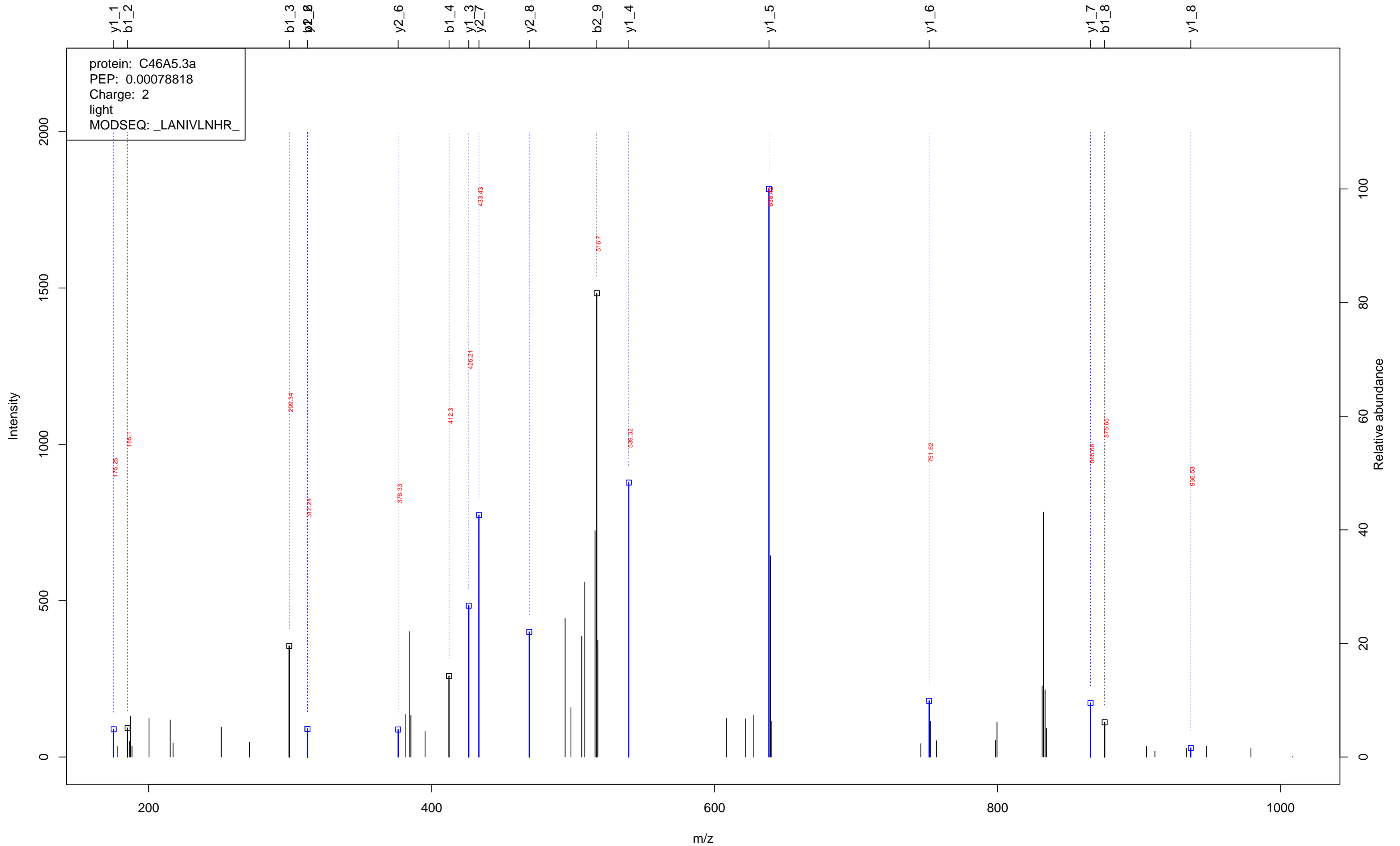


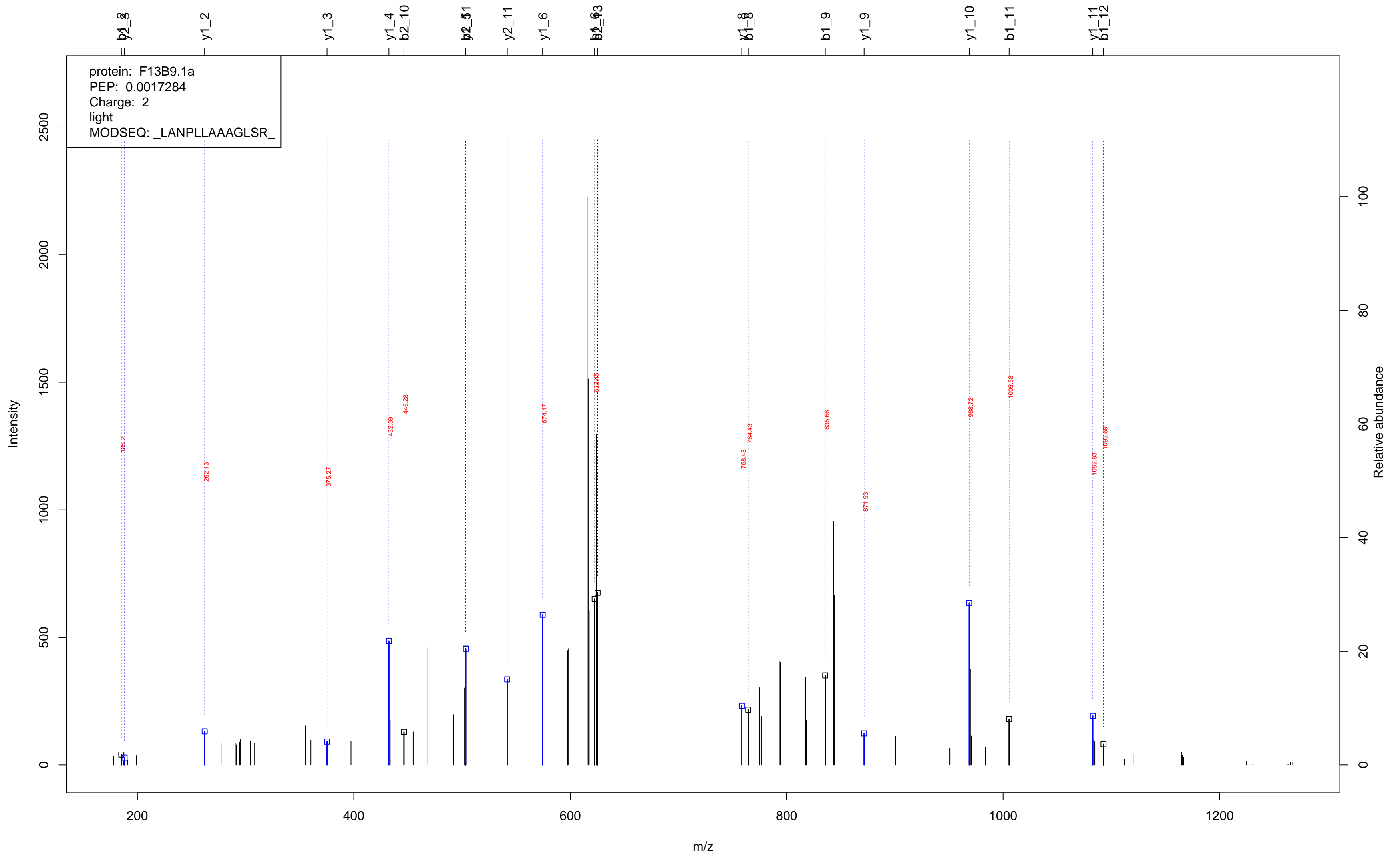


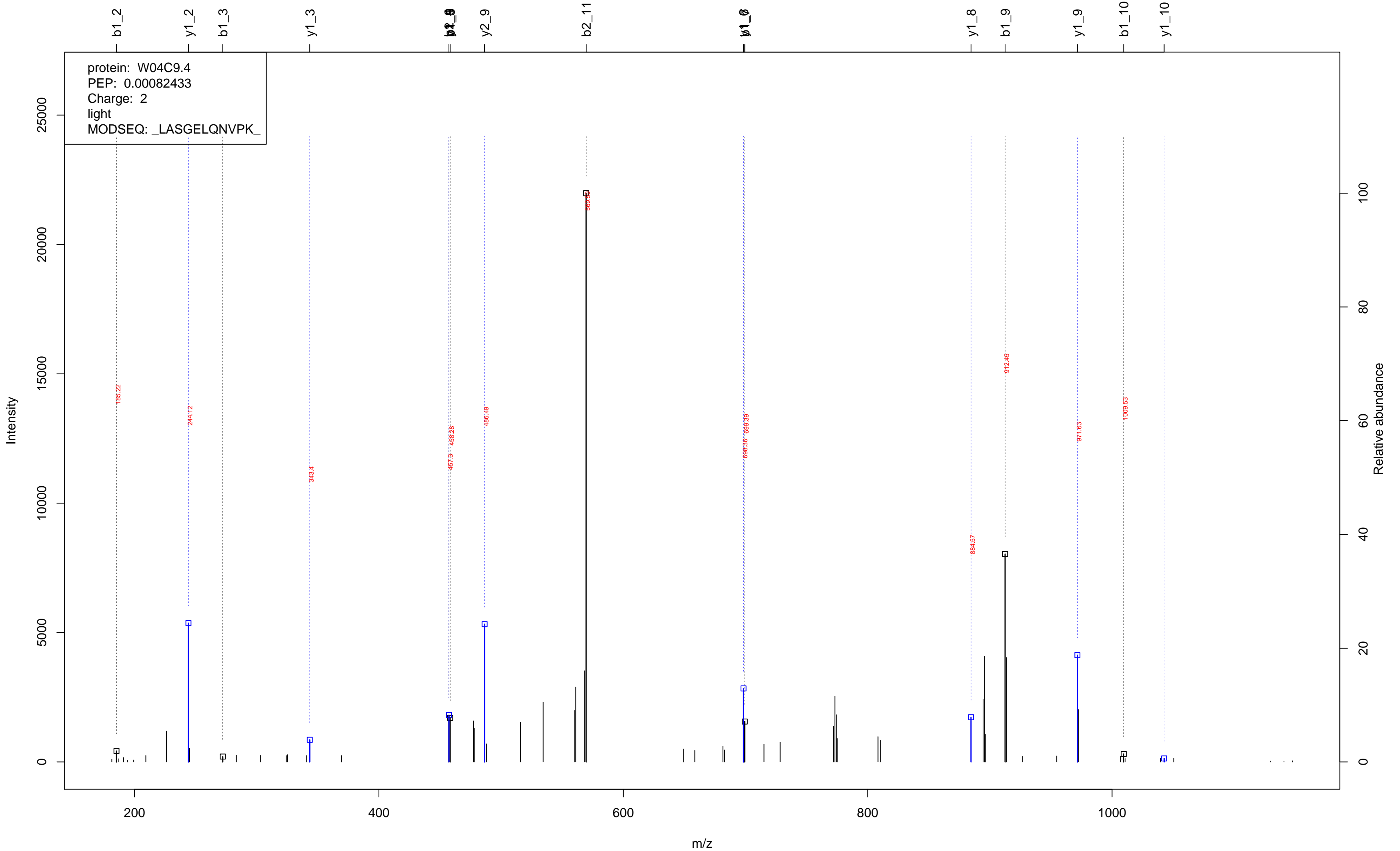


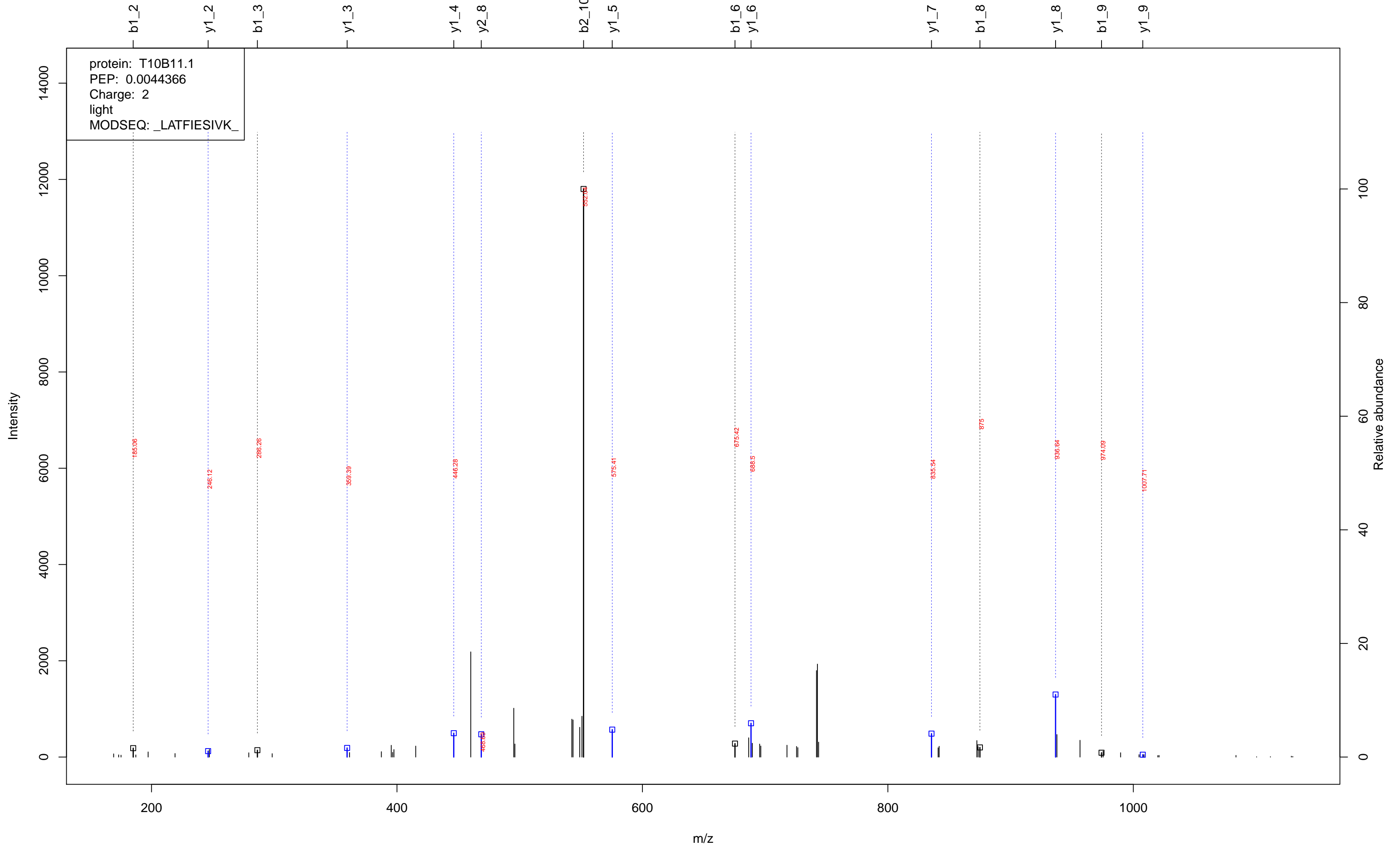


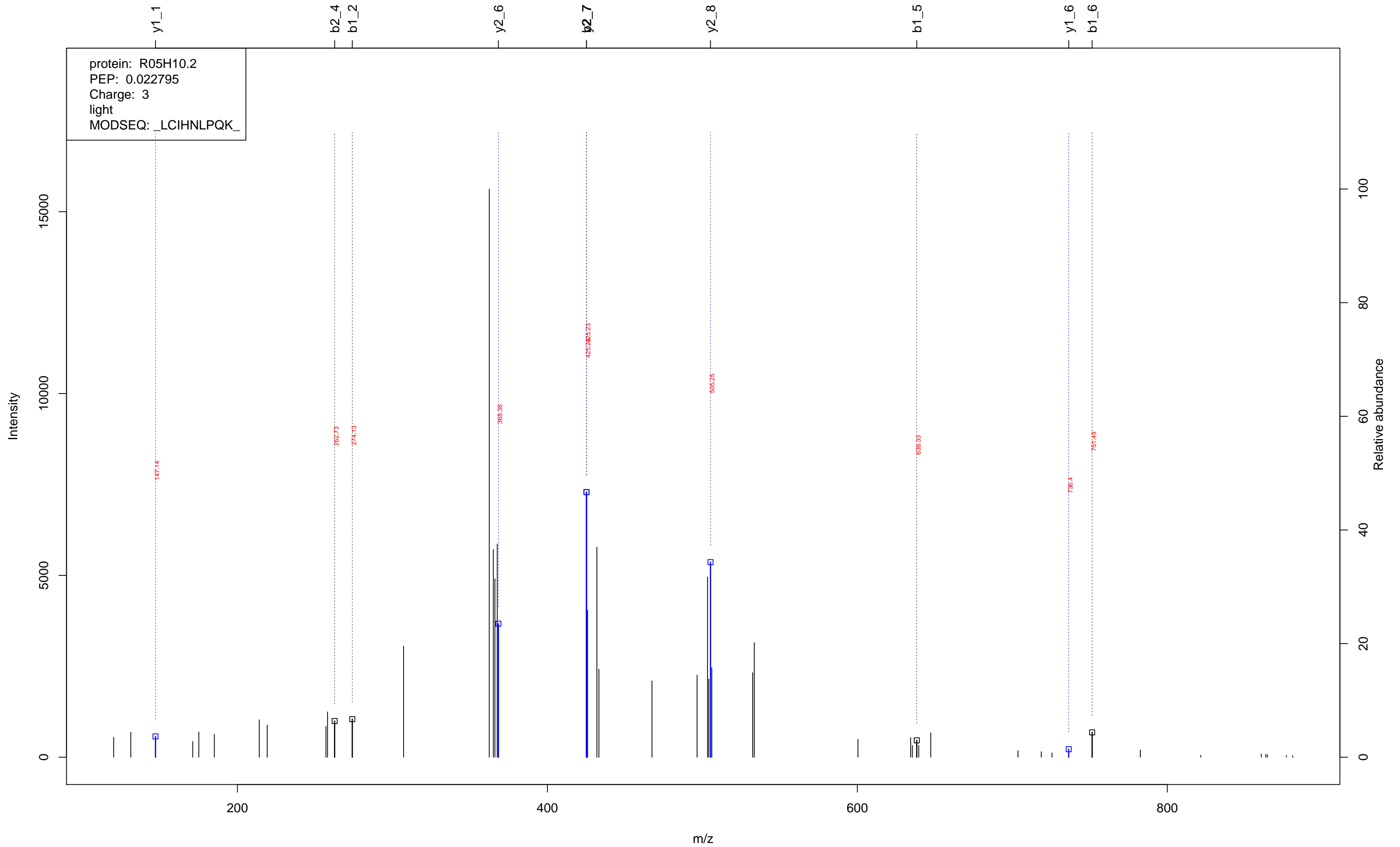


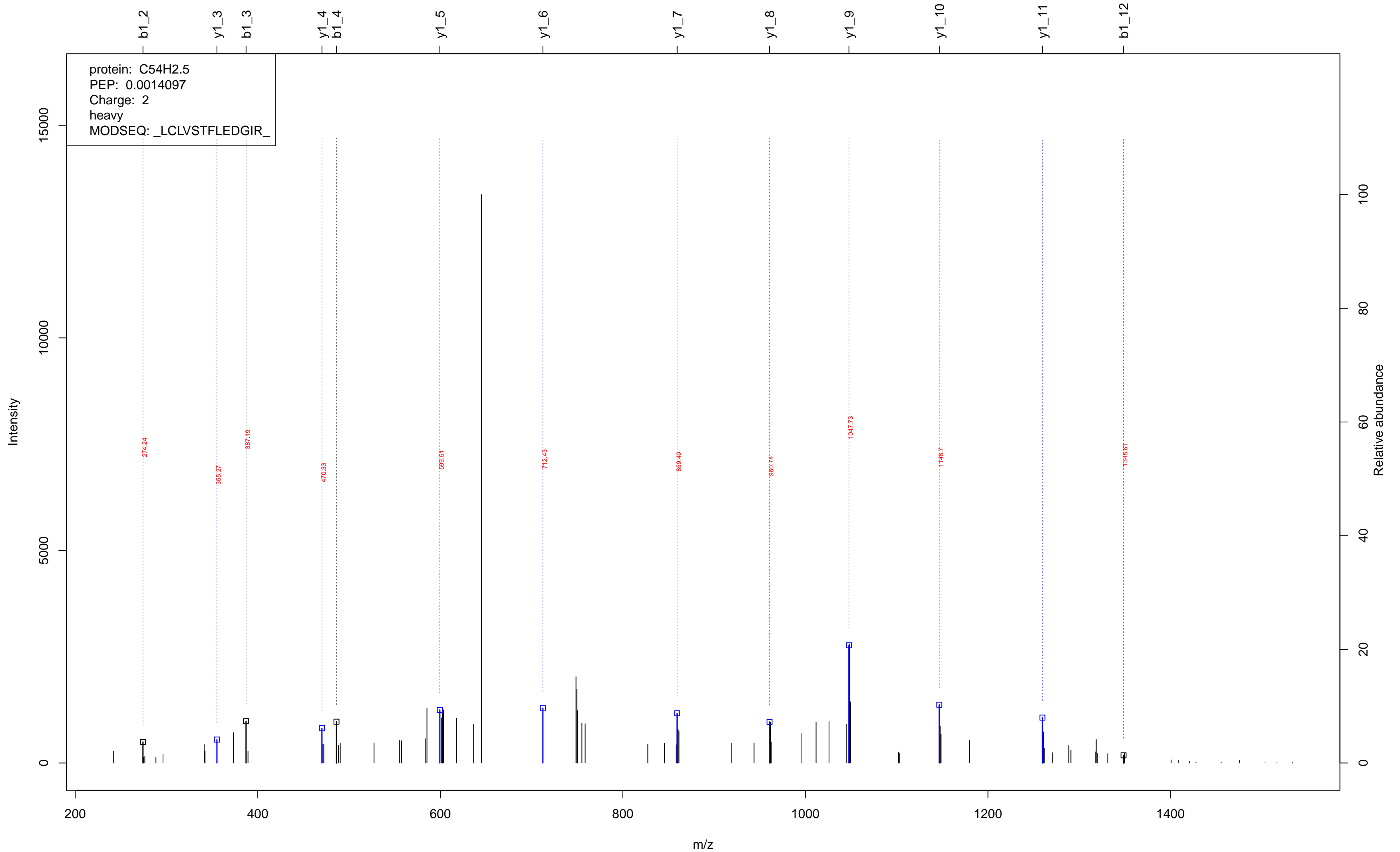


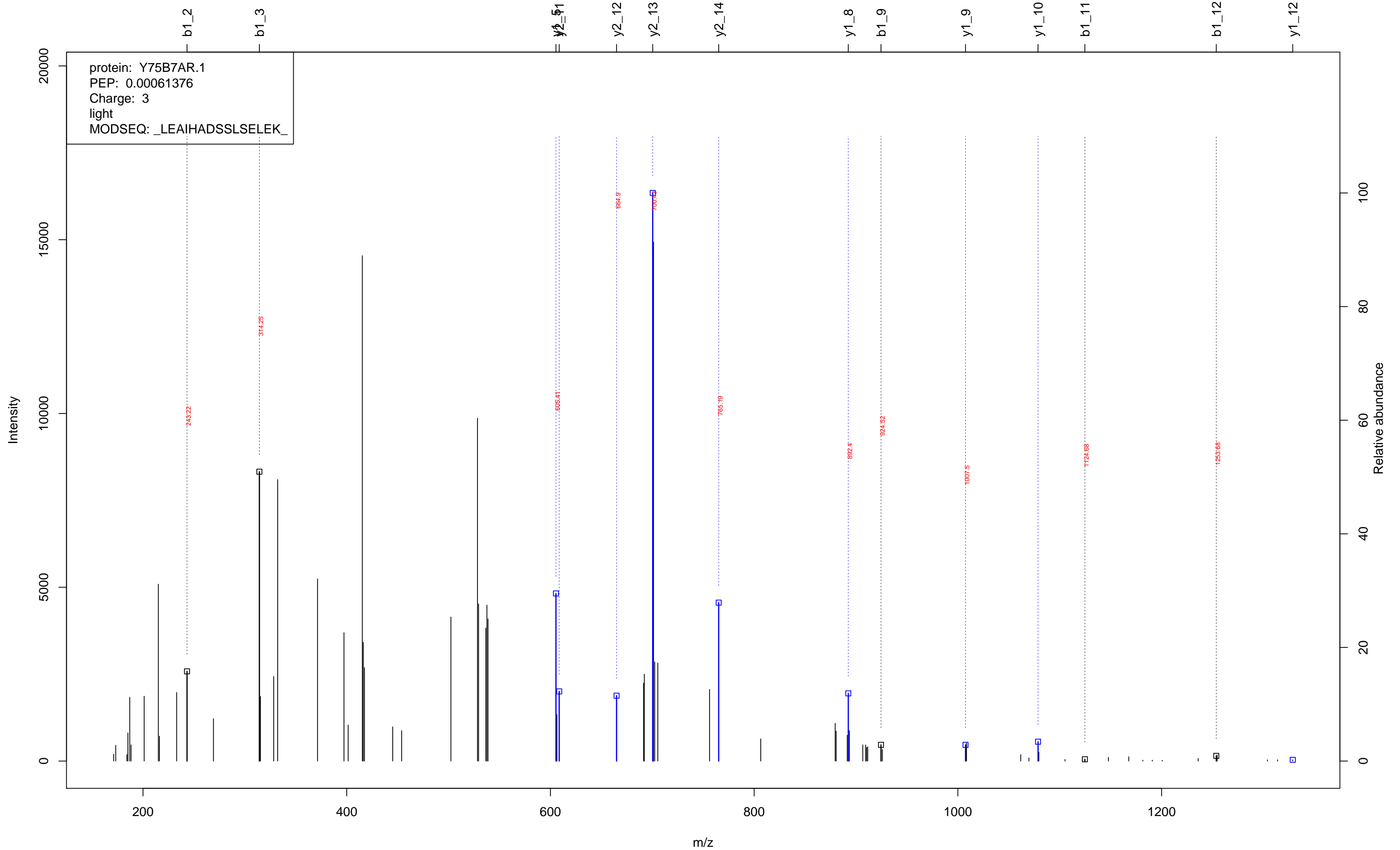


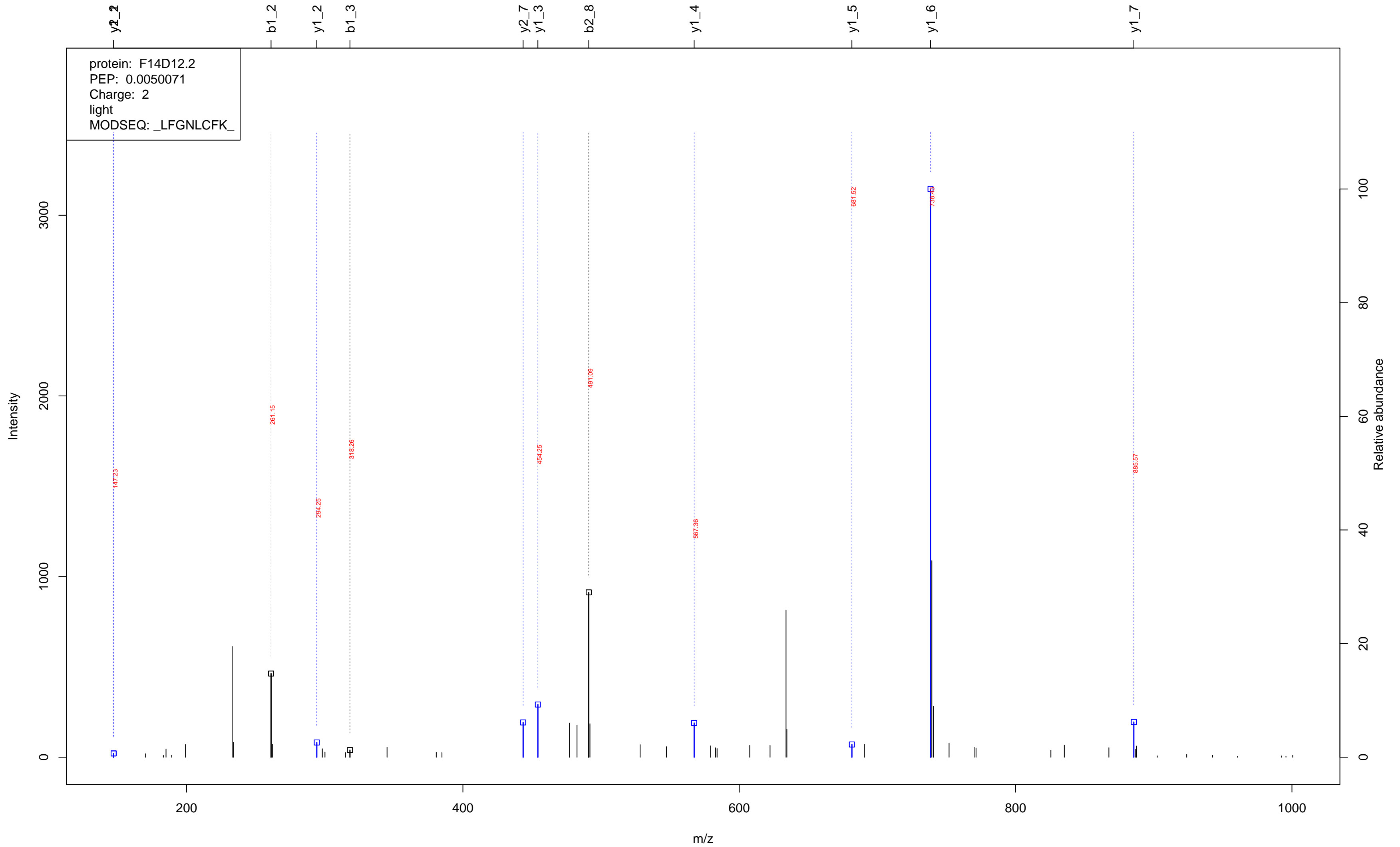


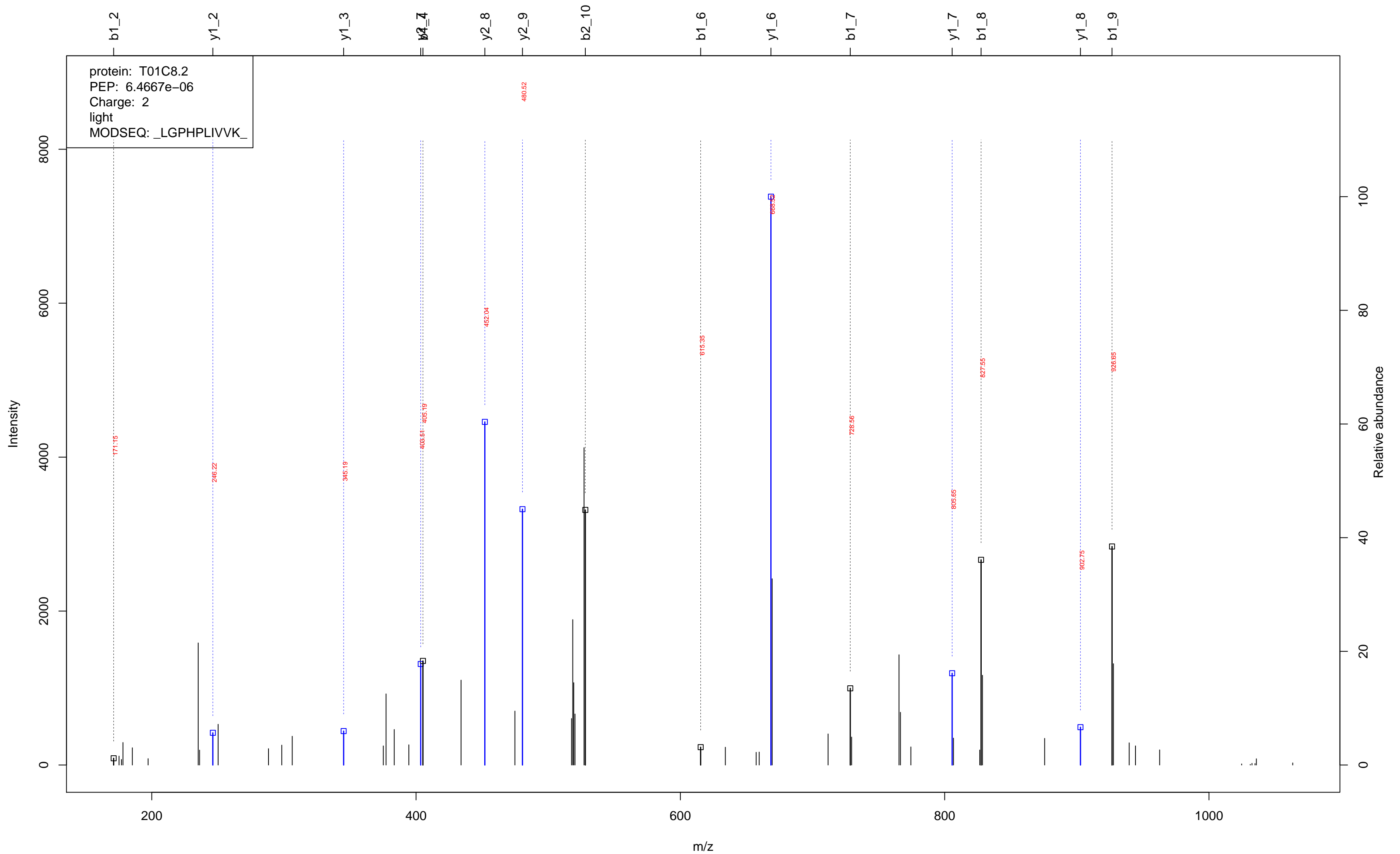


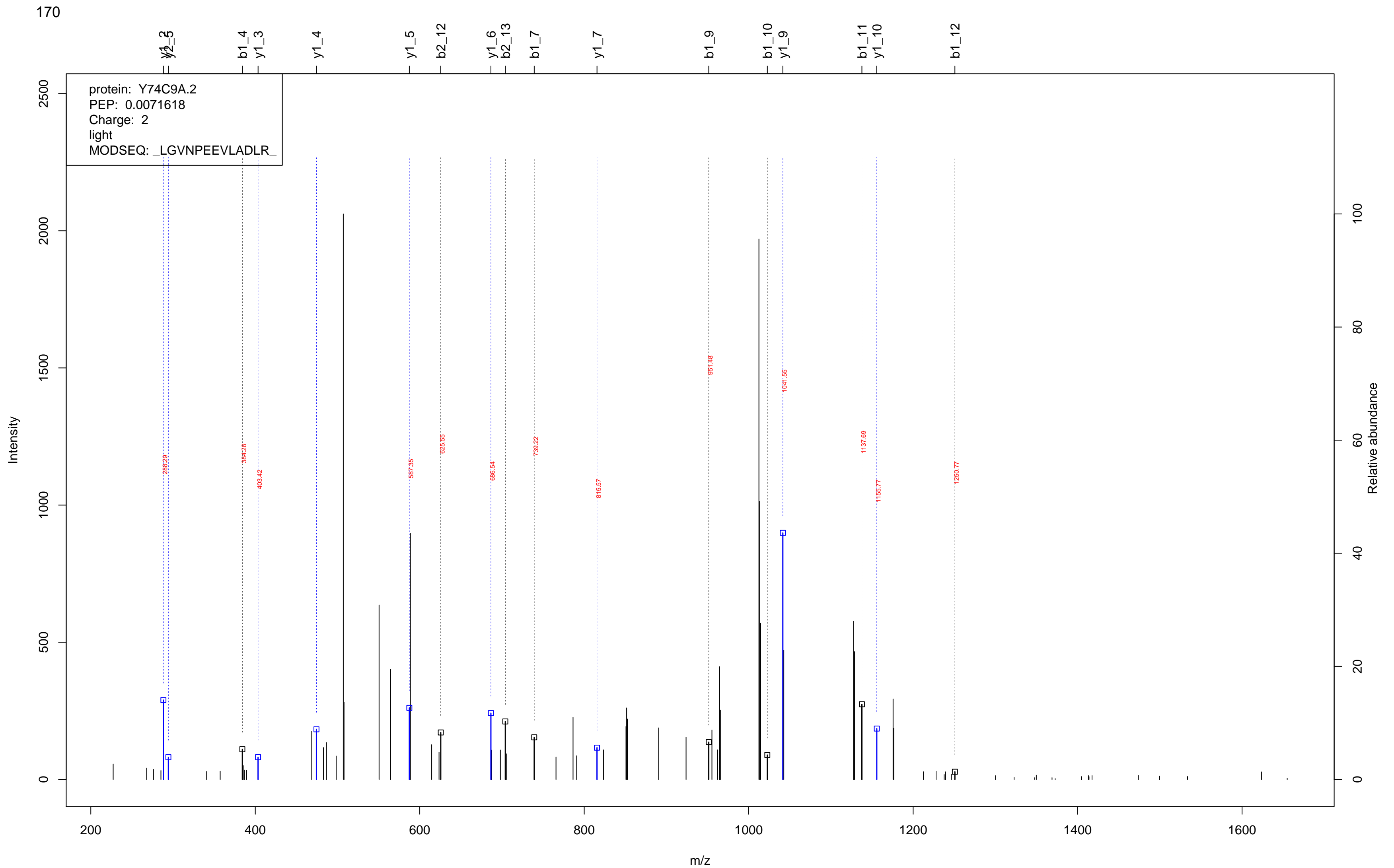


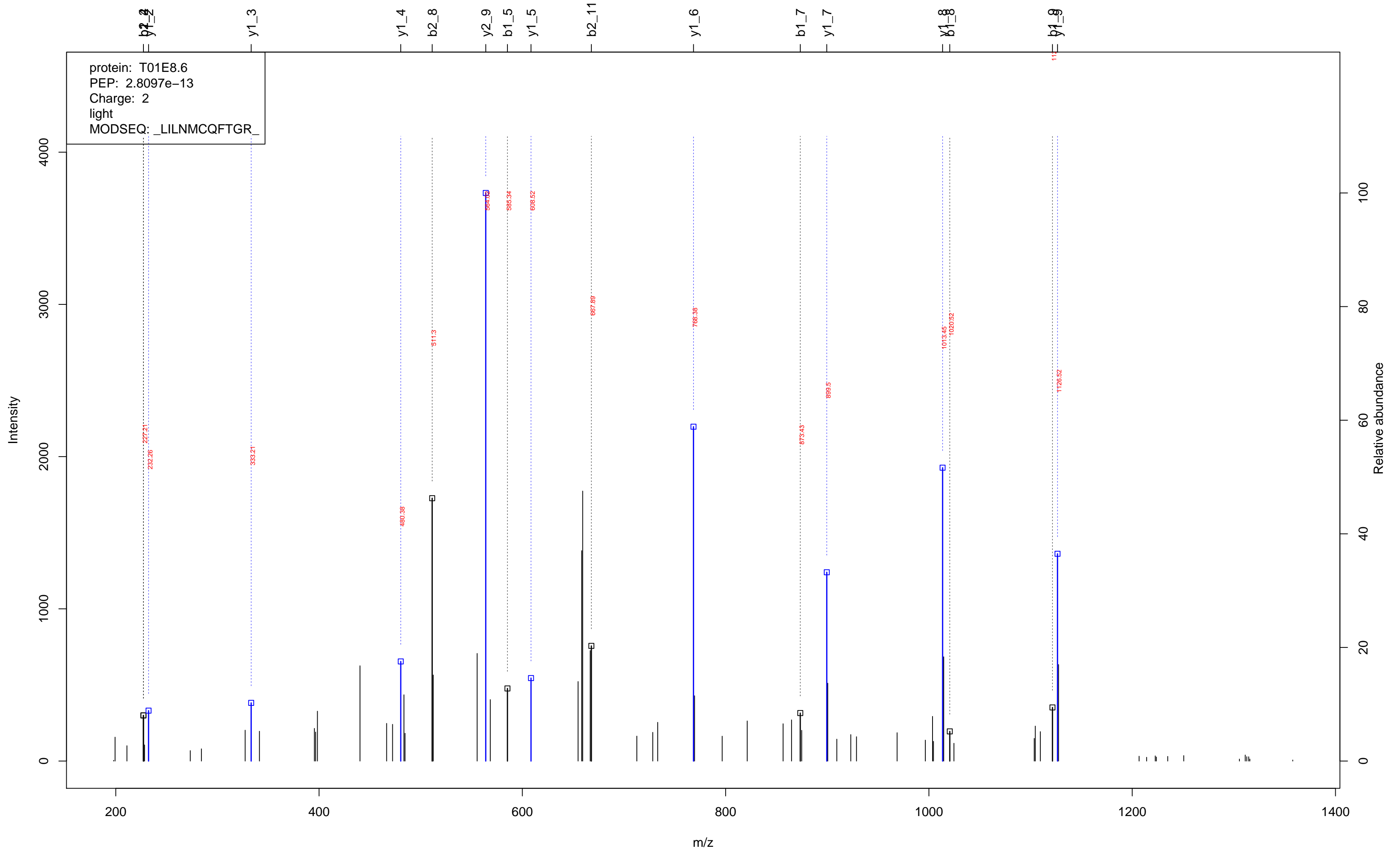


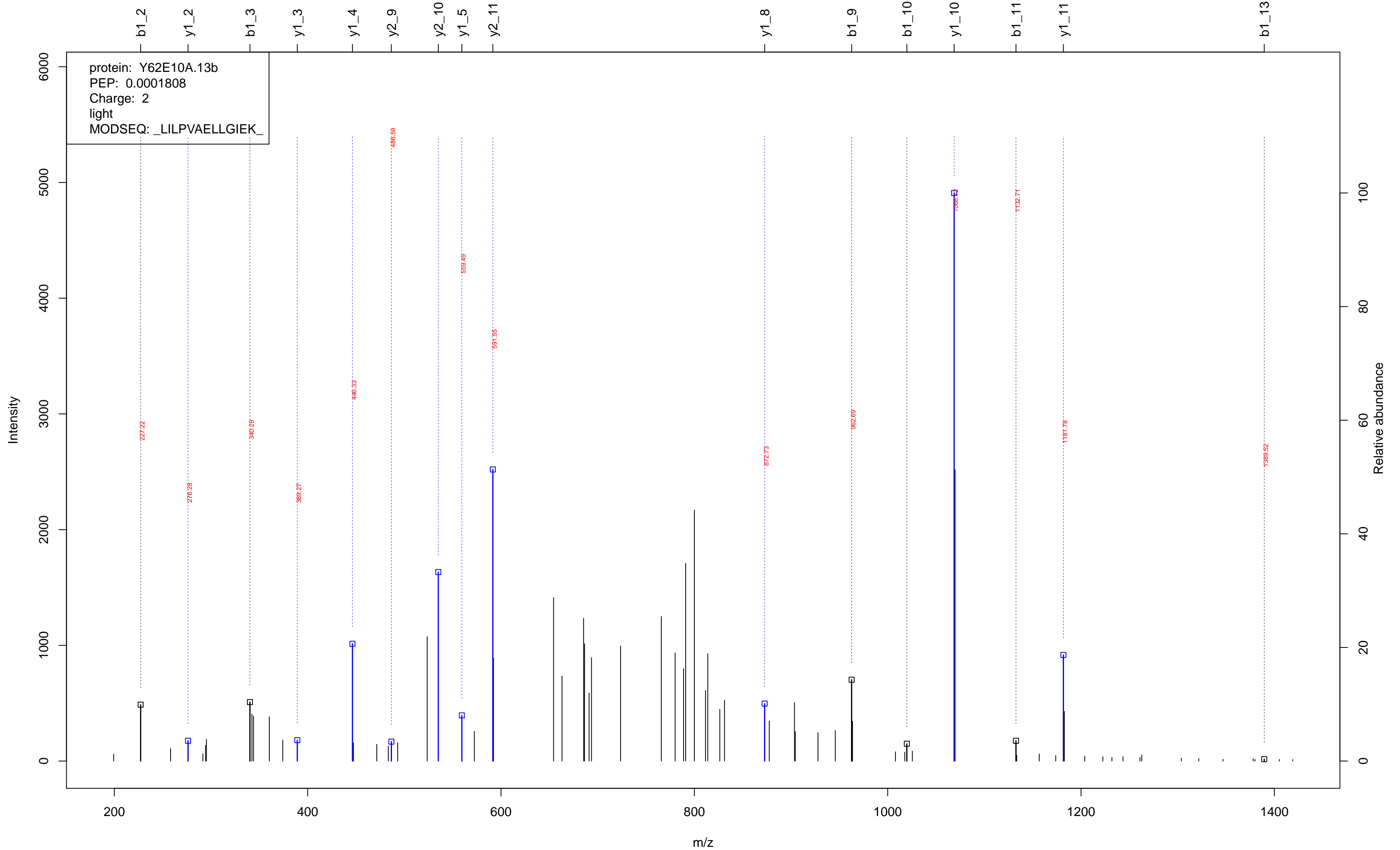


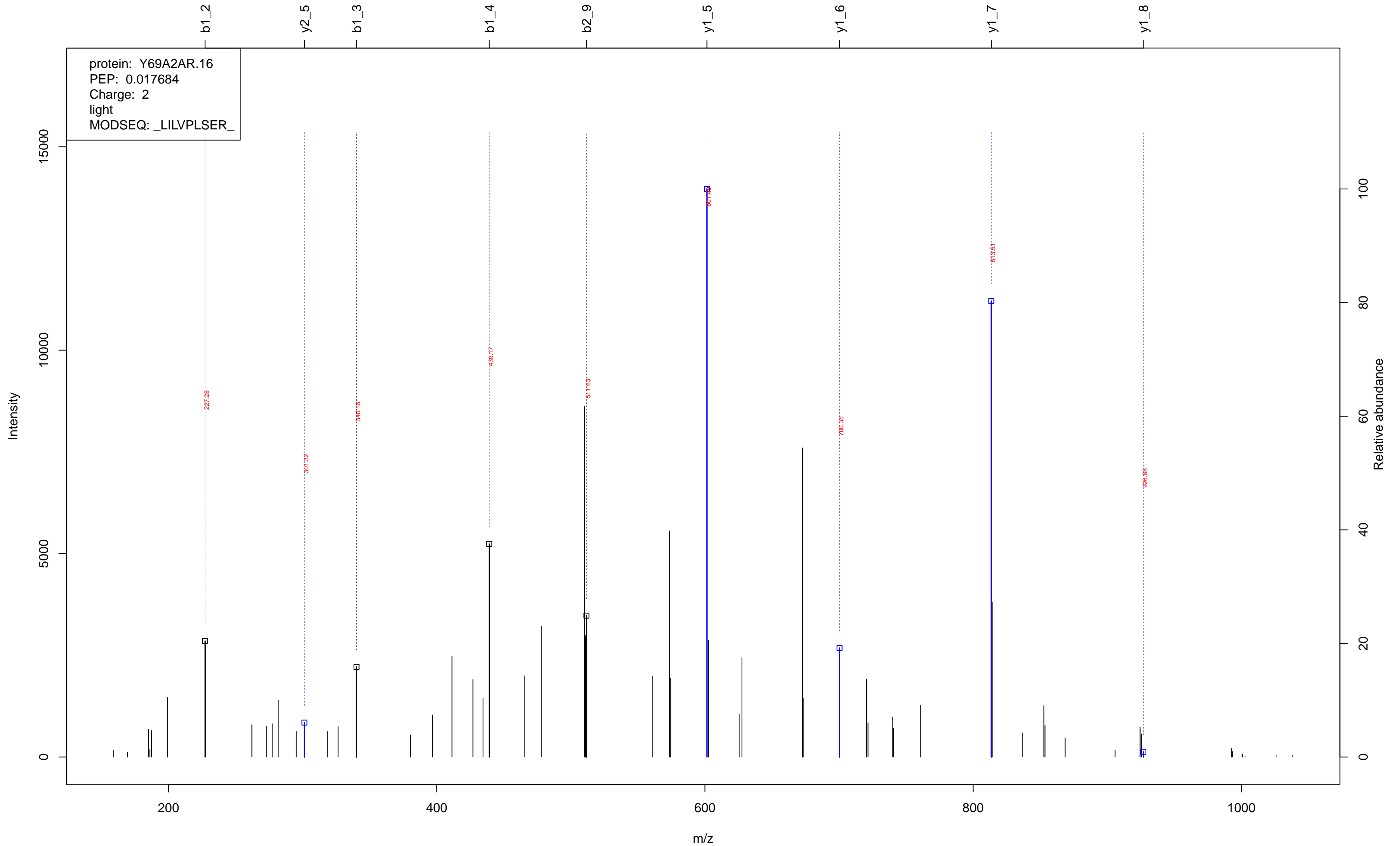


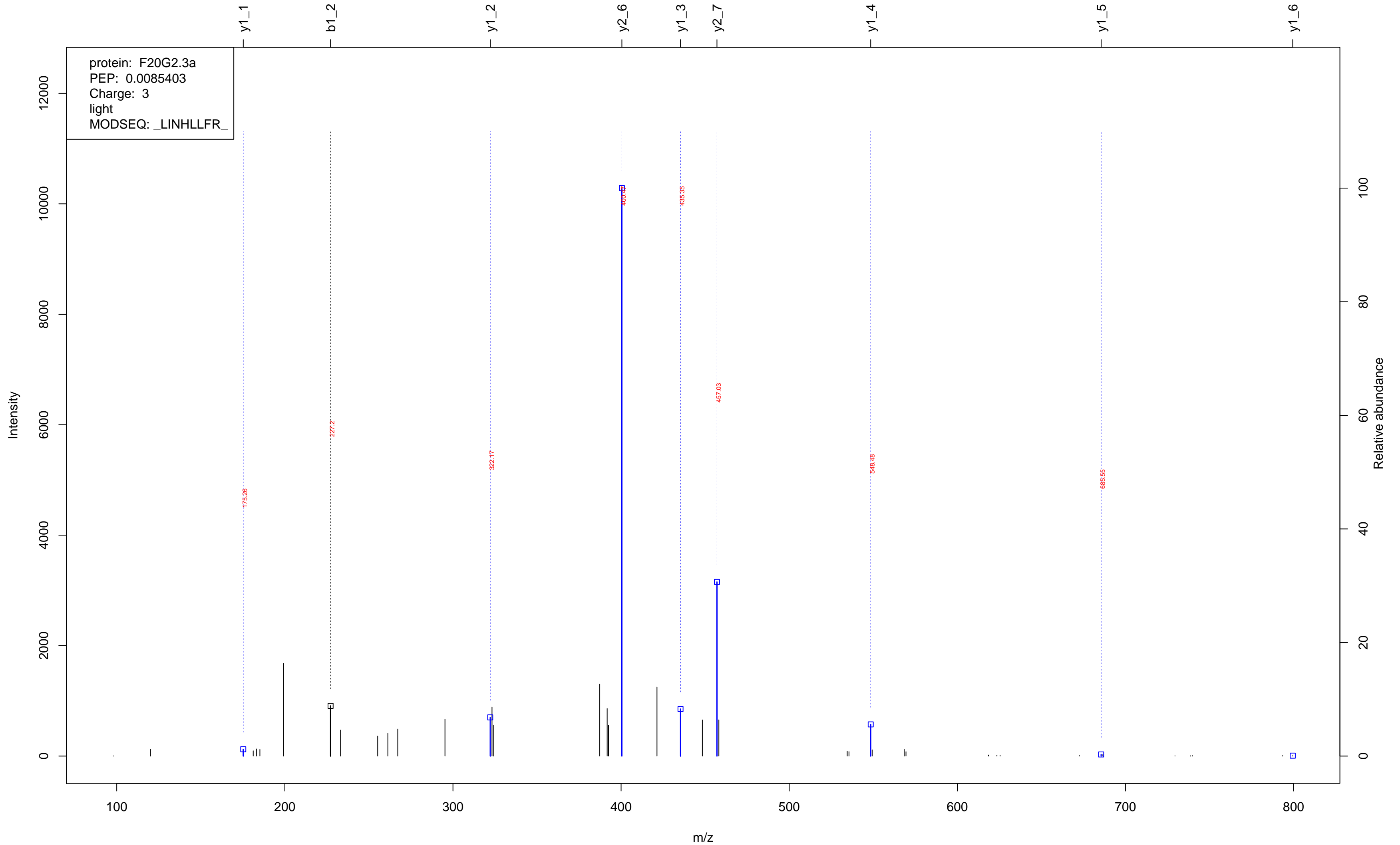


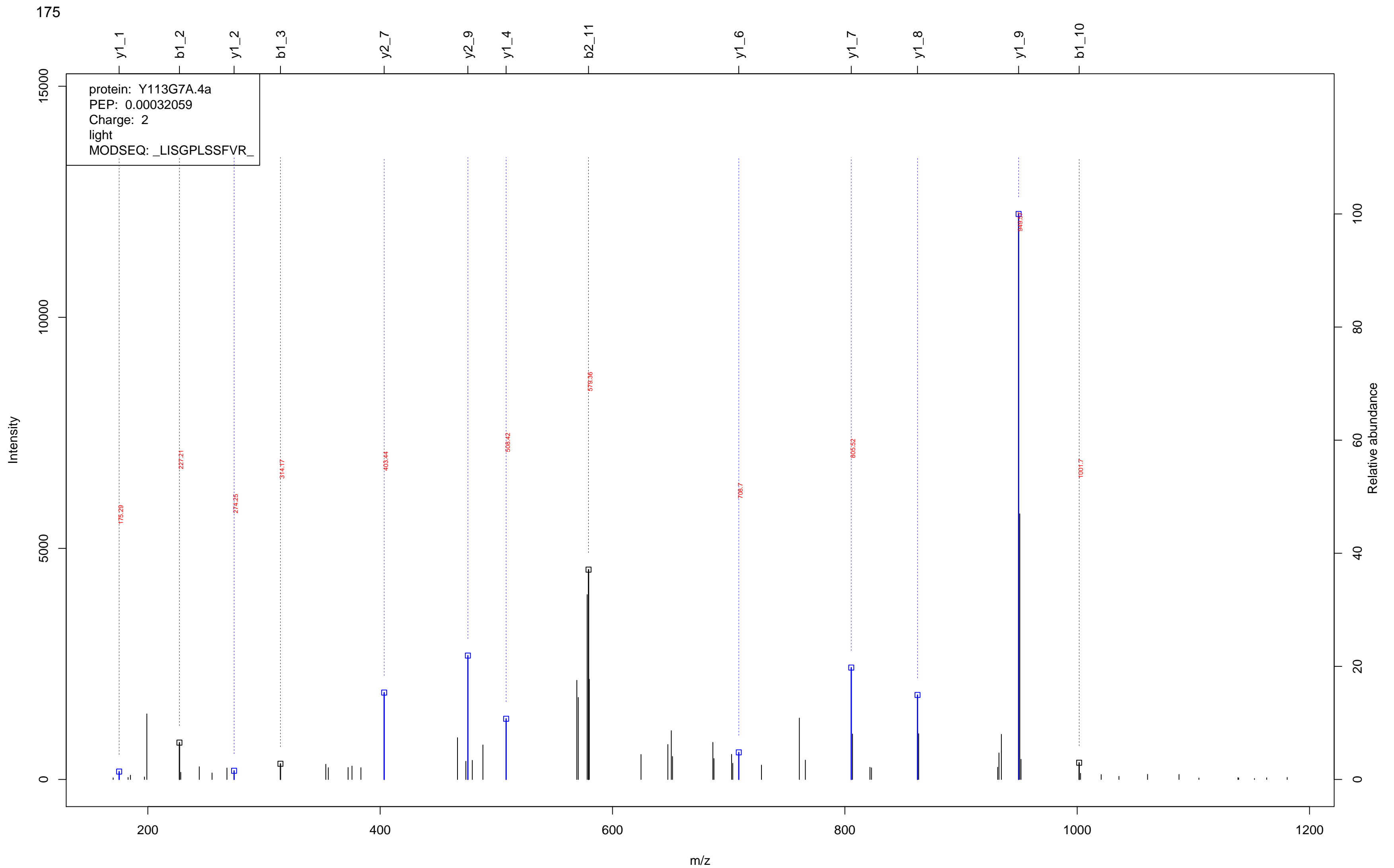


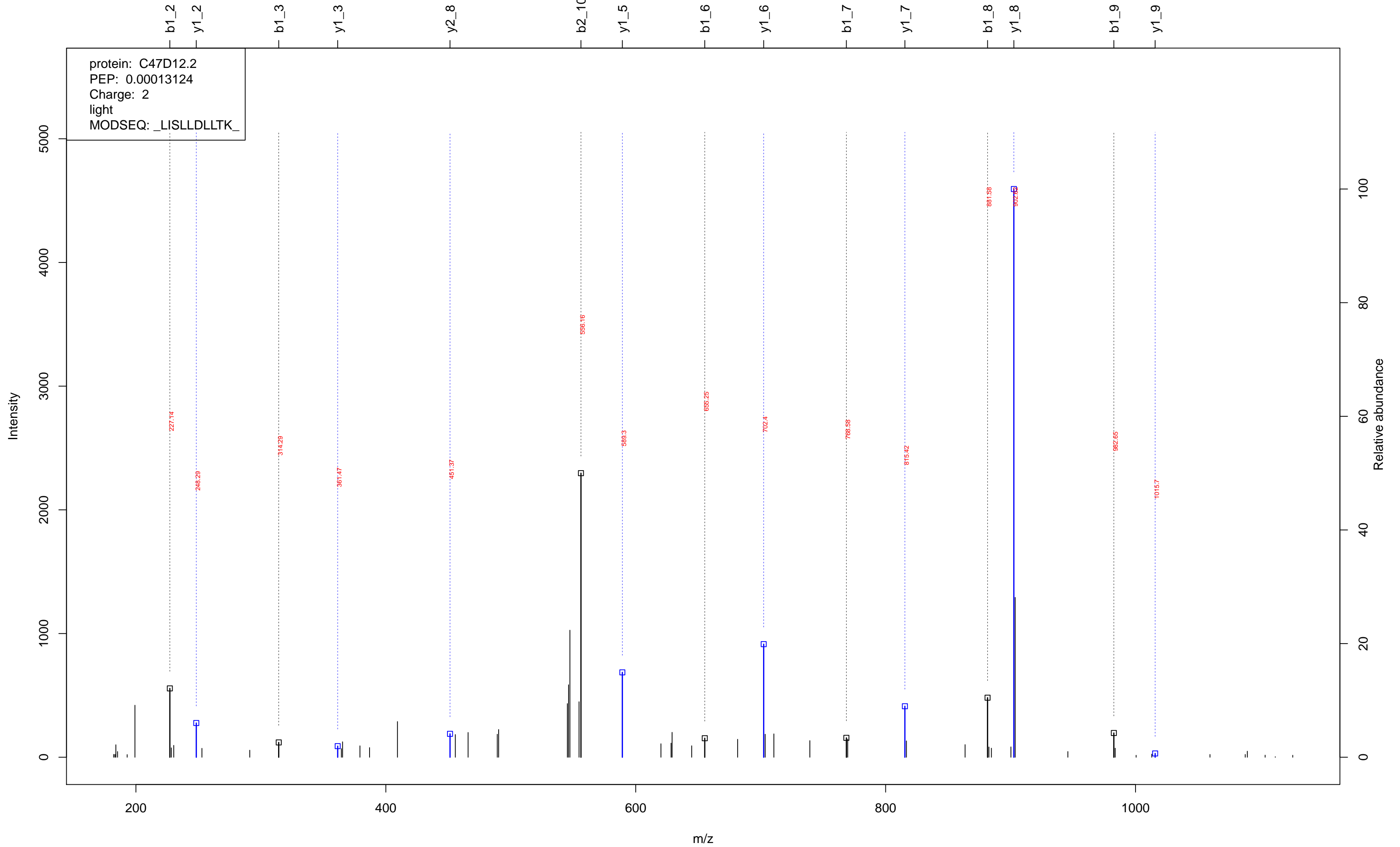


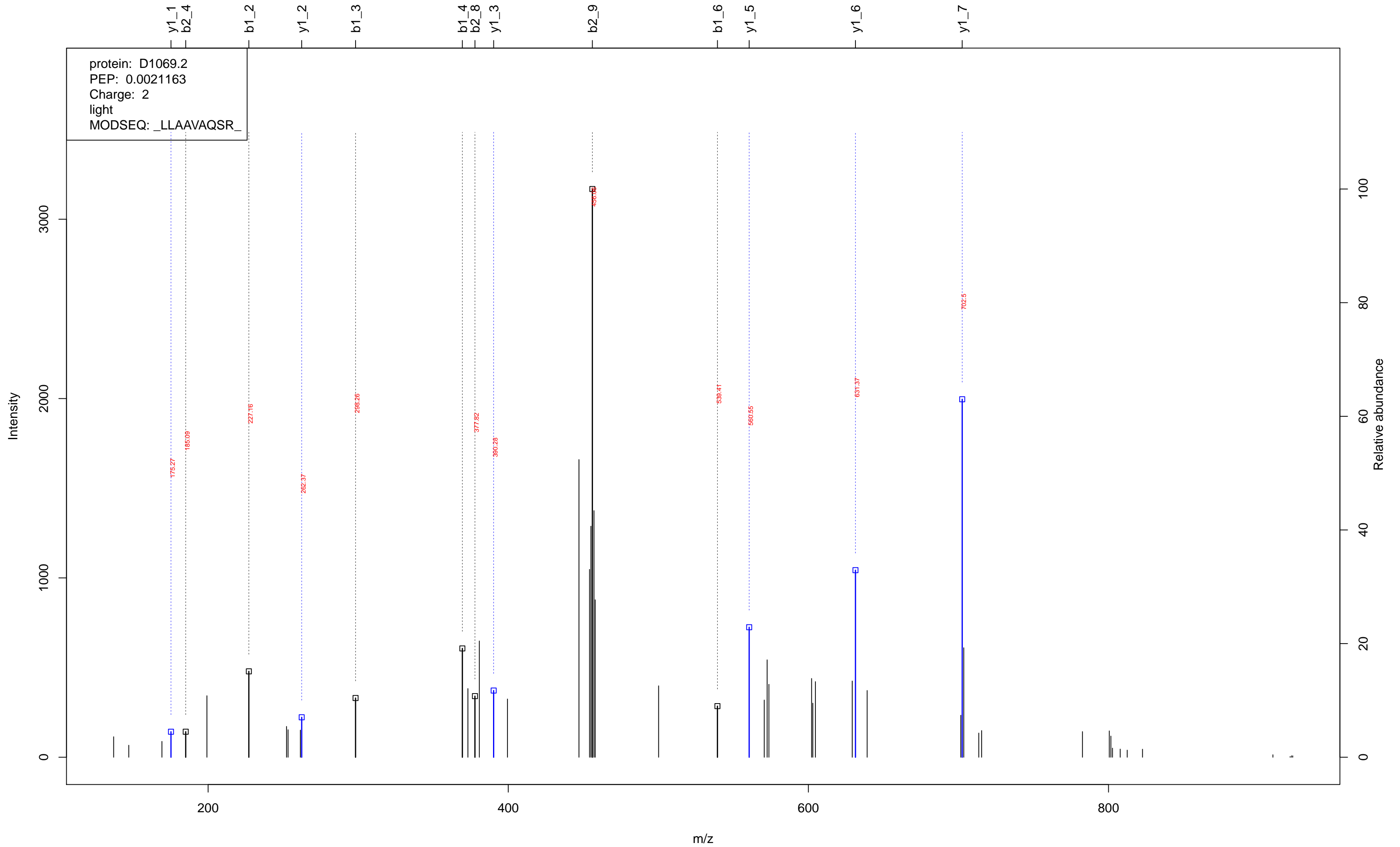


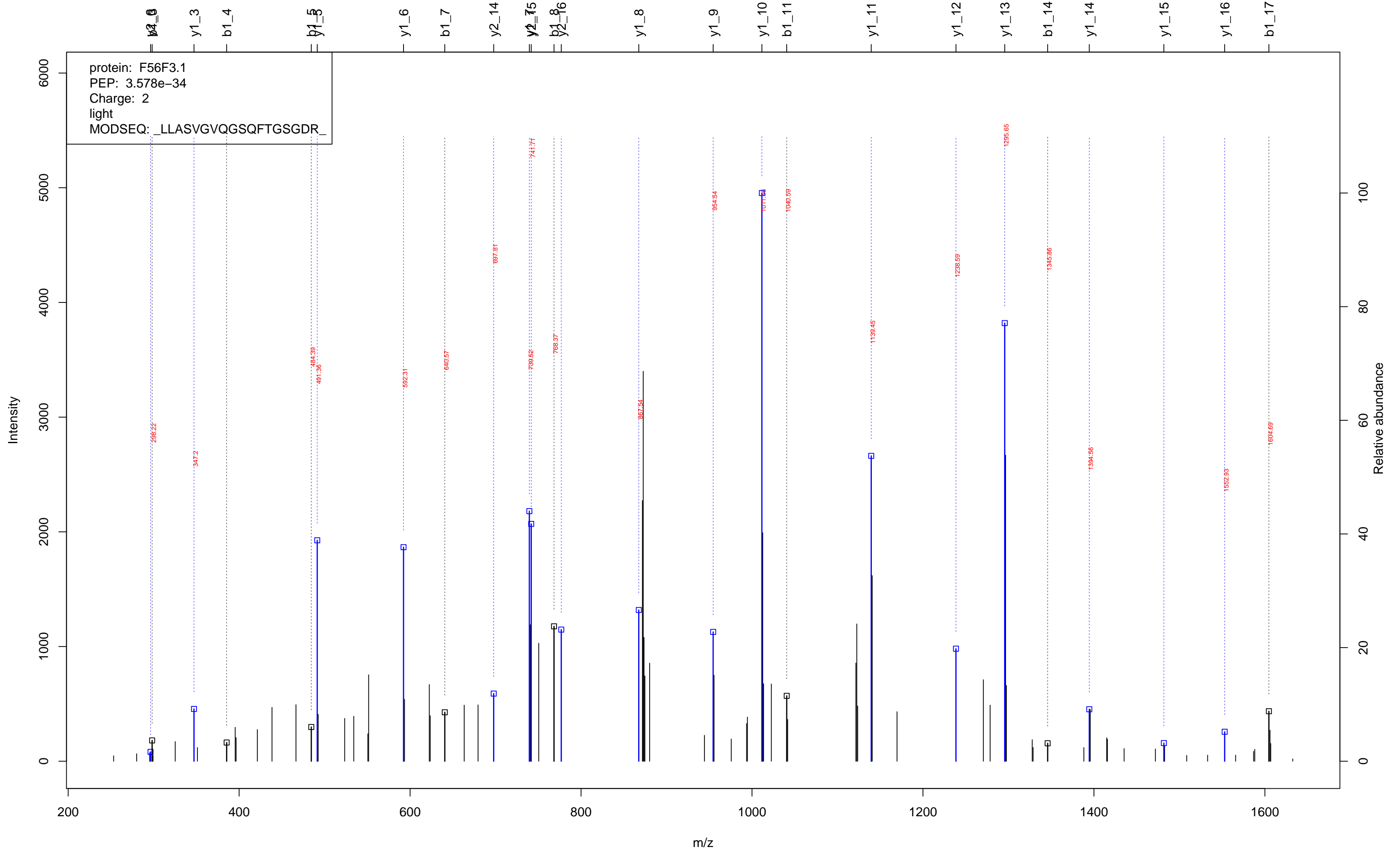






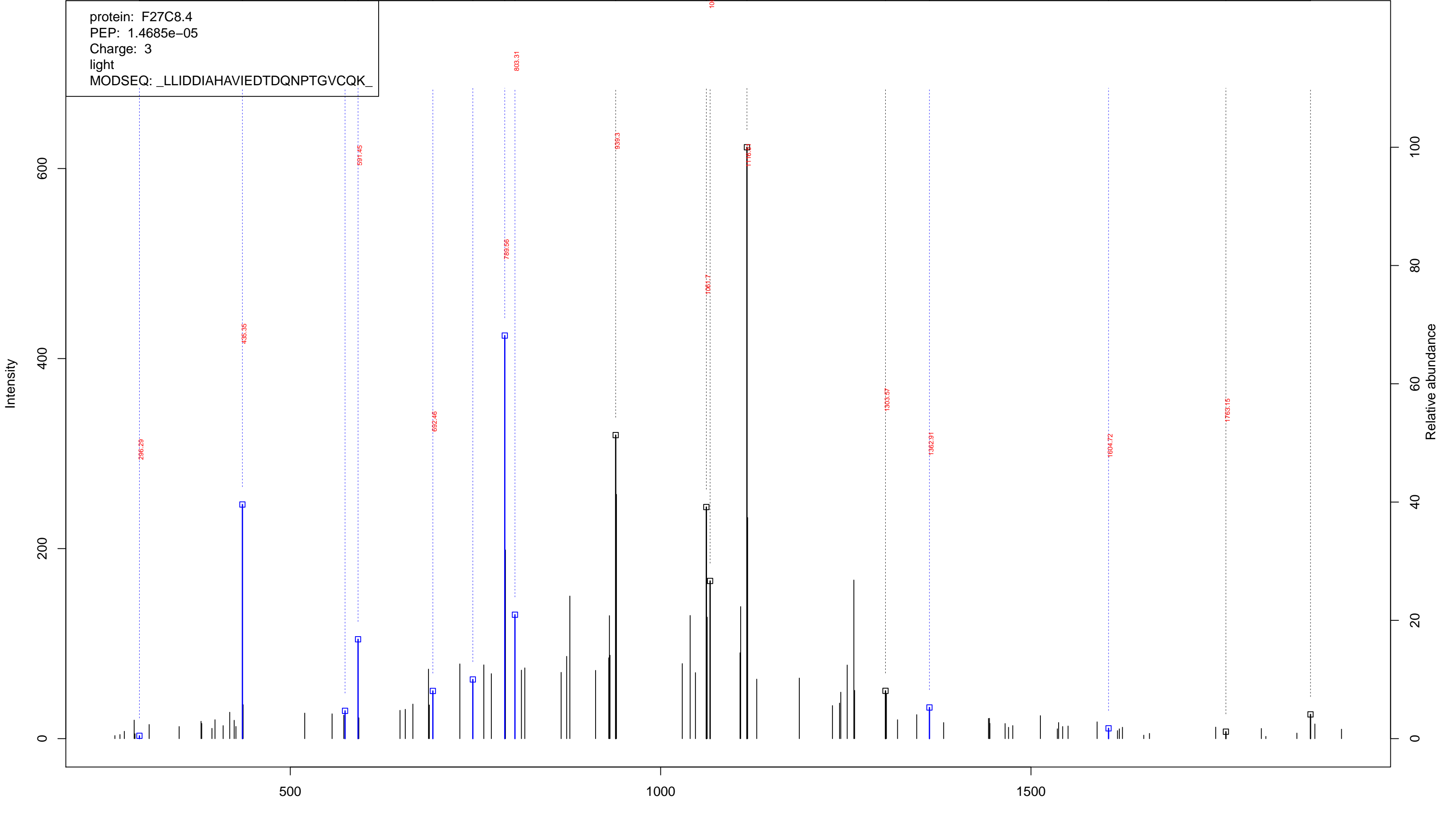


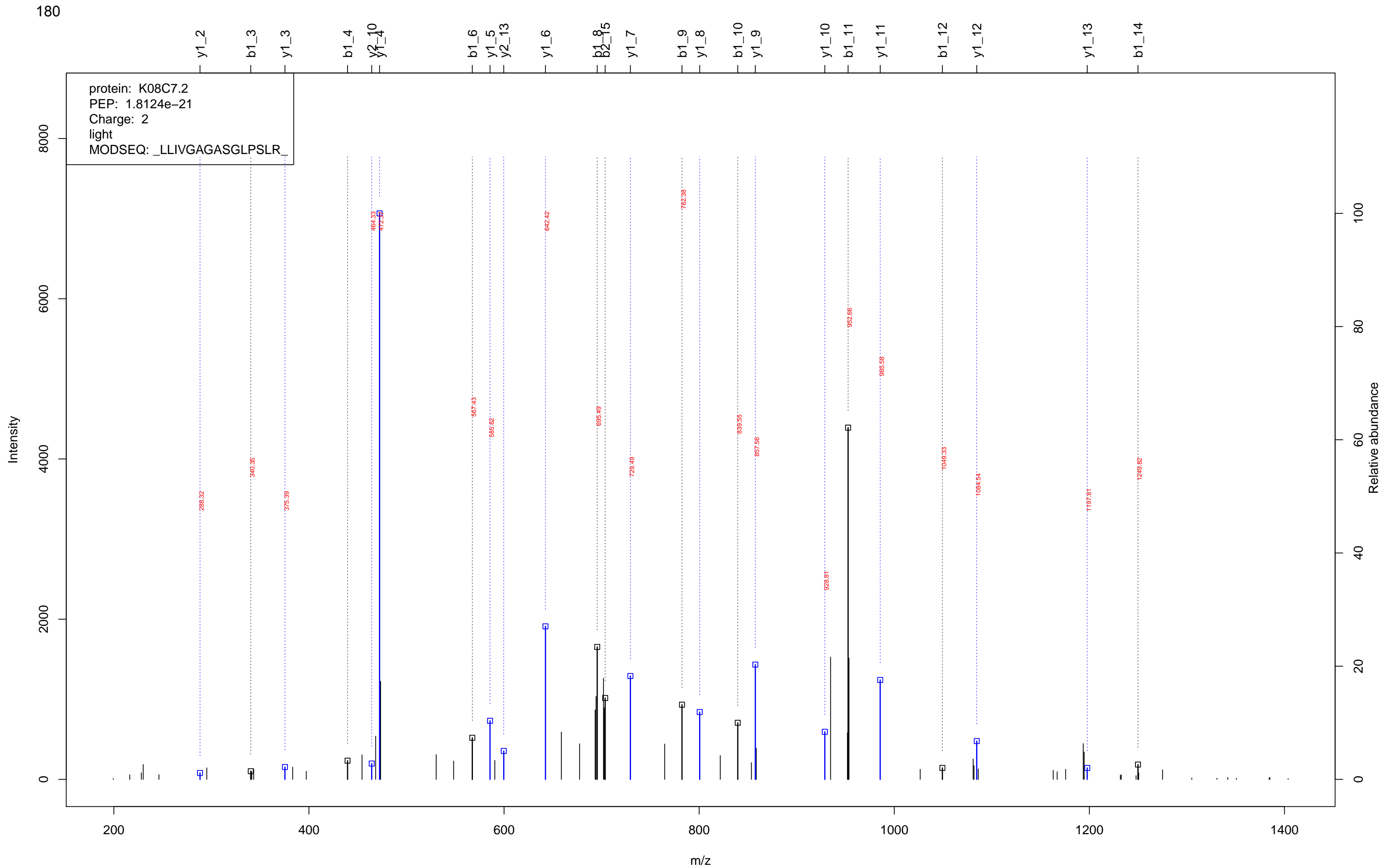


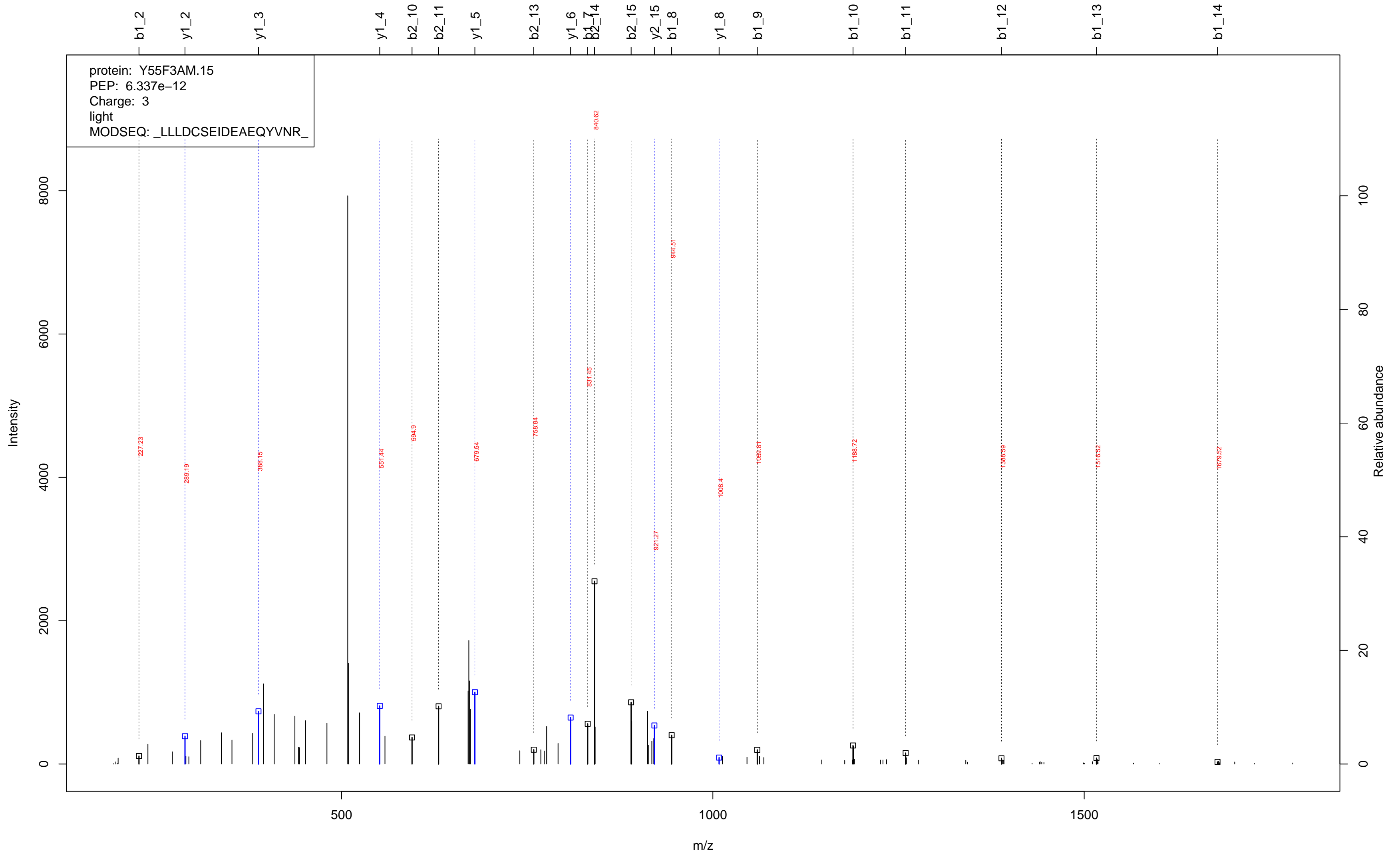


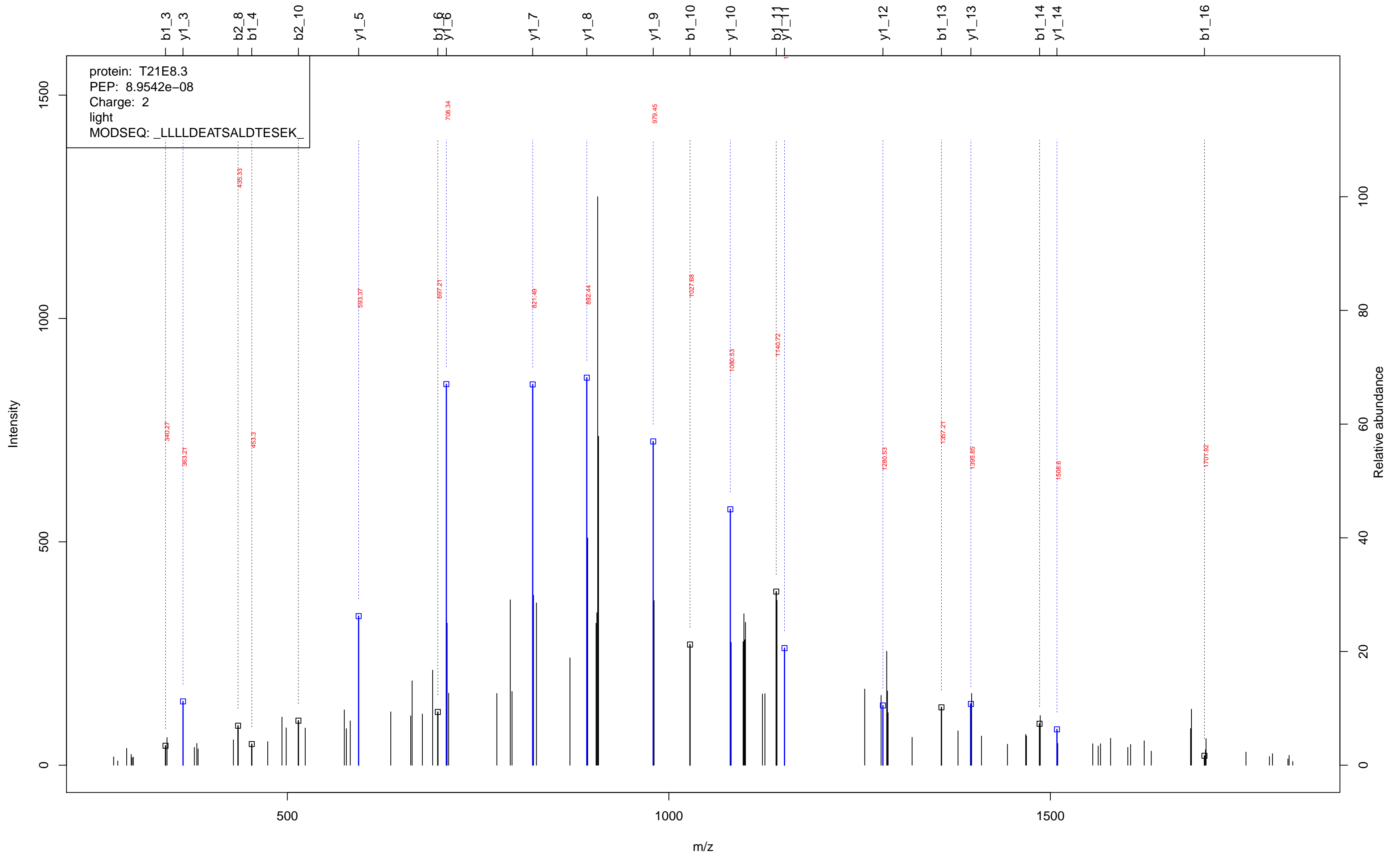
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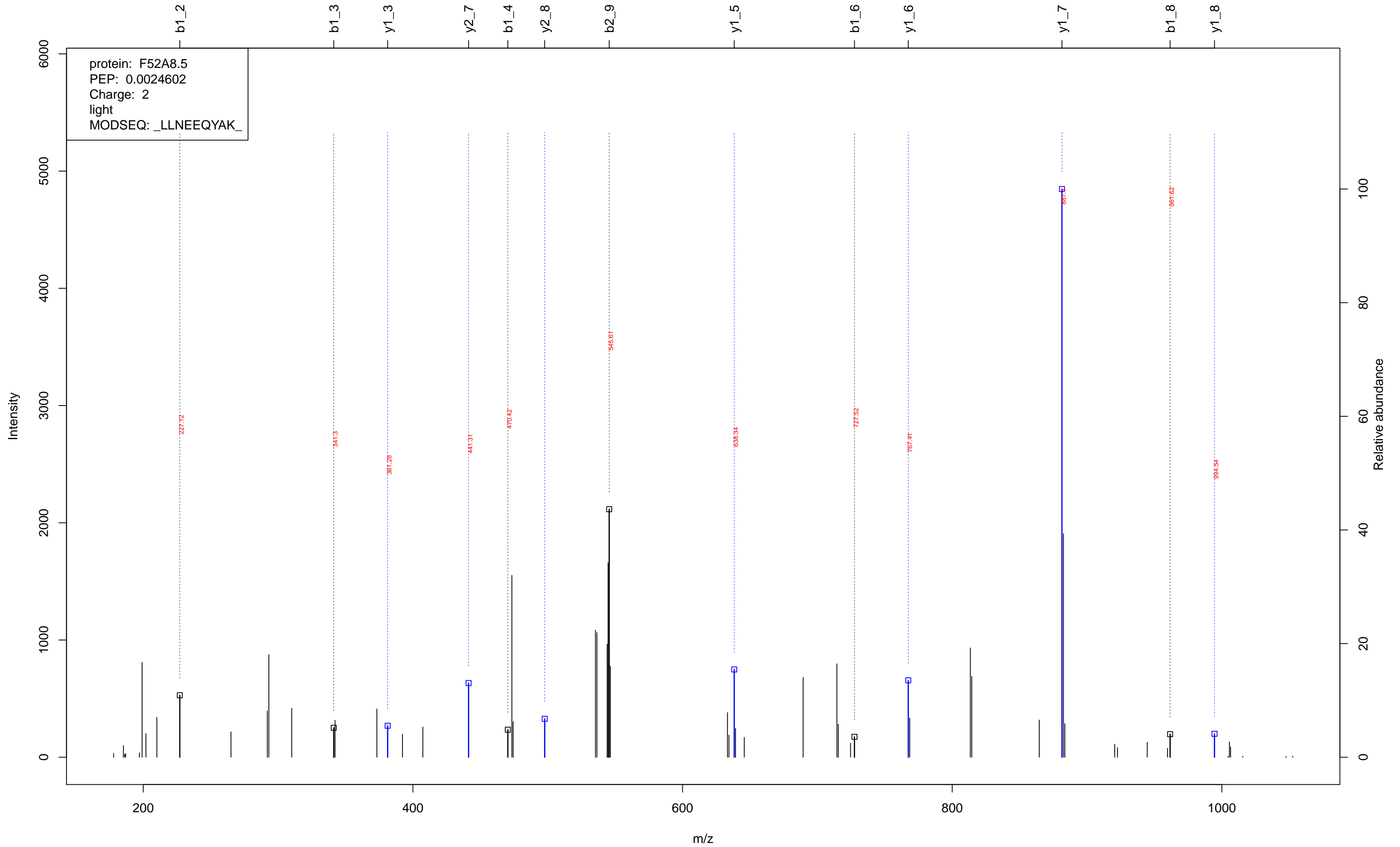
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Charge: 3
light
MODSEQ: _LLIDDIHAVIEDTDQNPTGVCQK_

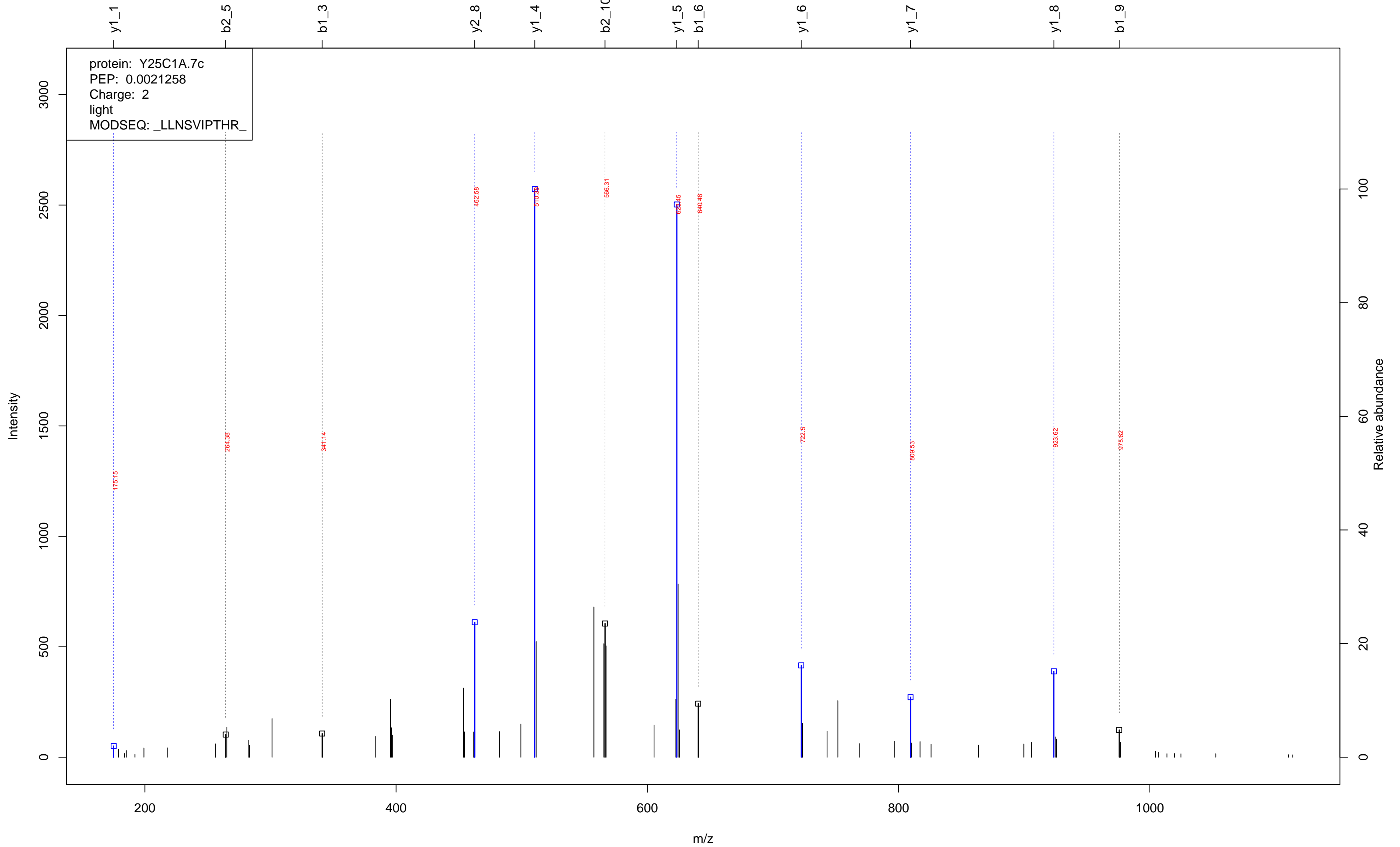


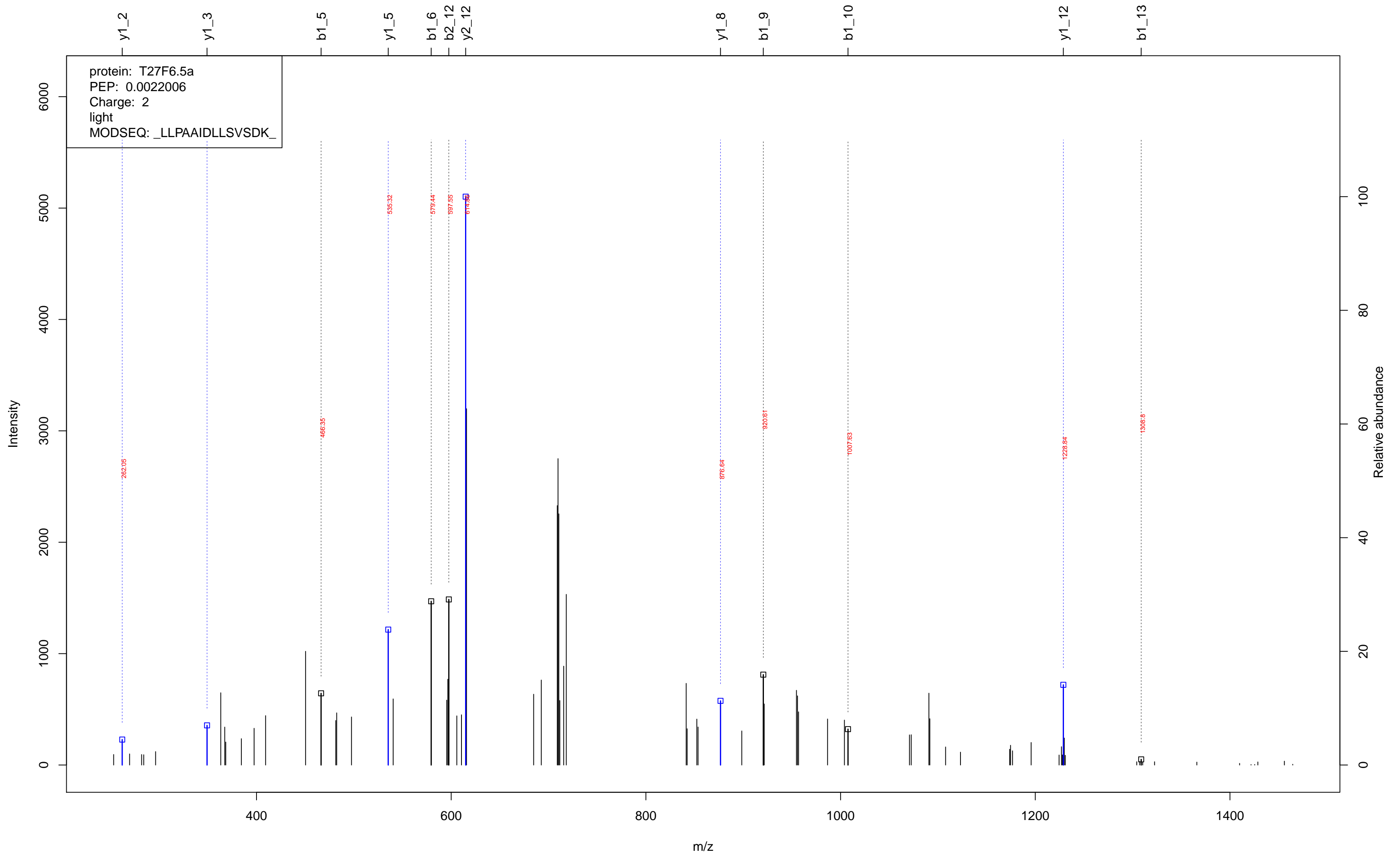


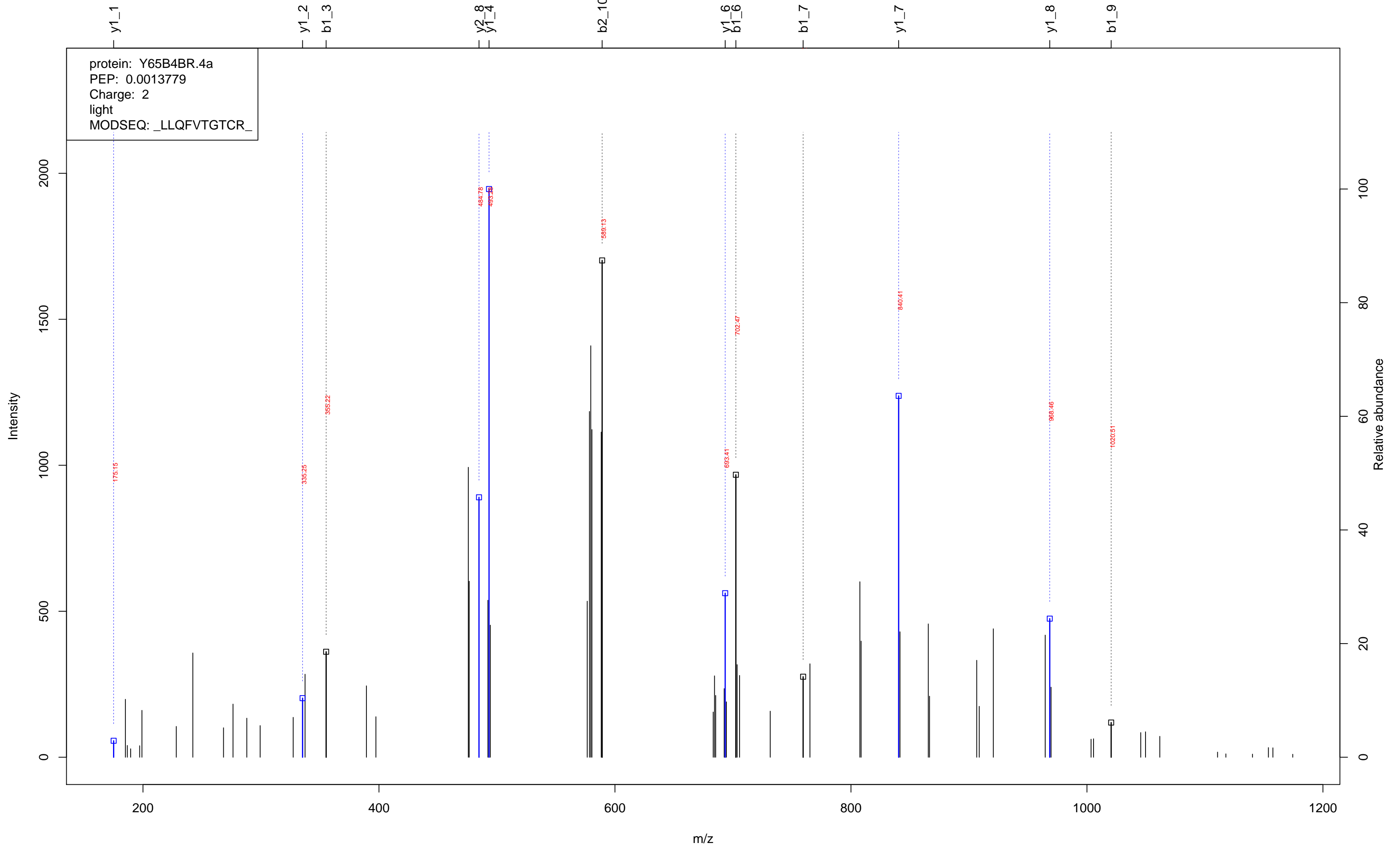


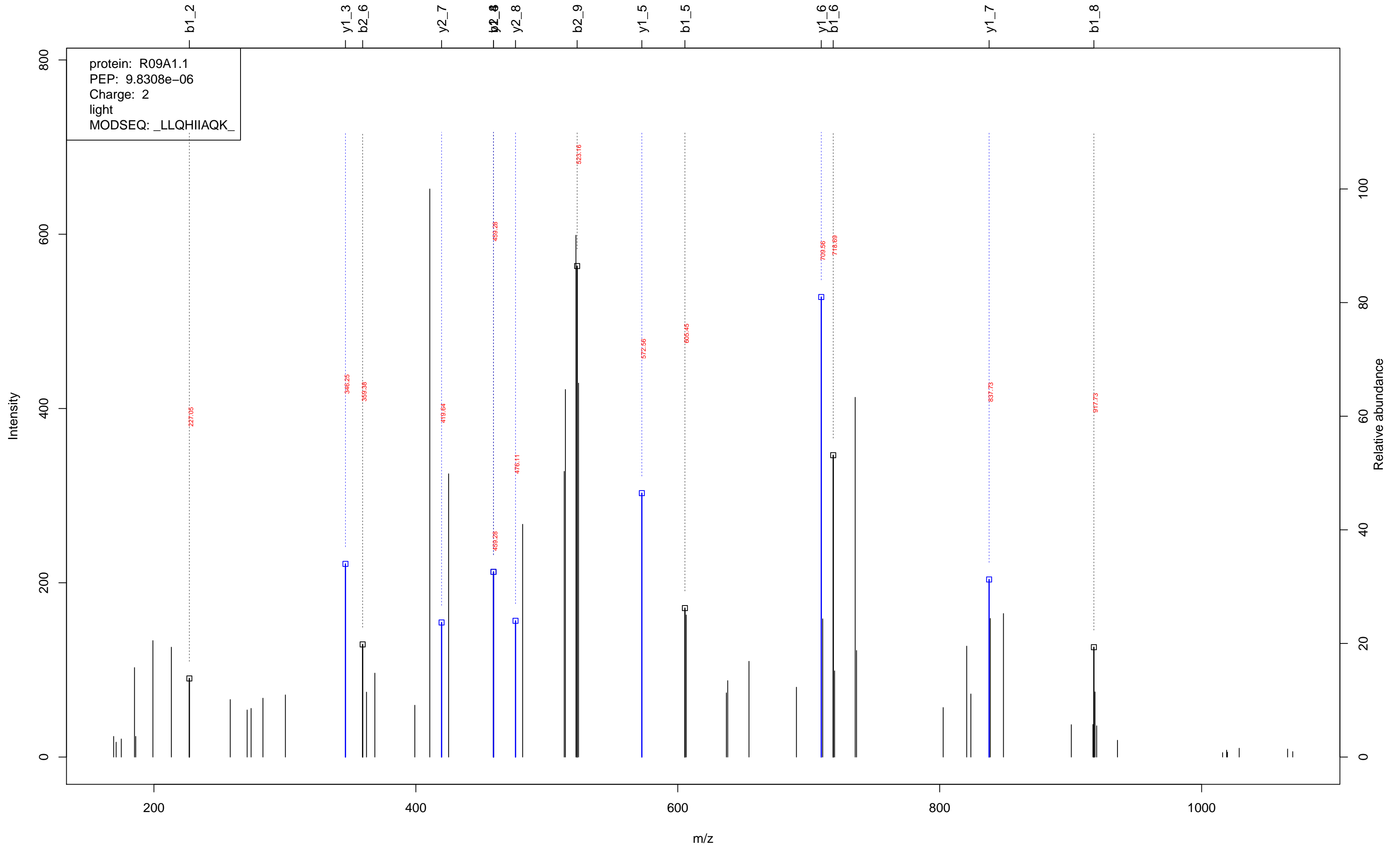


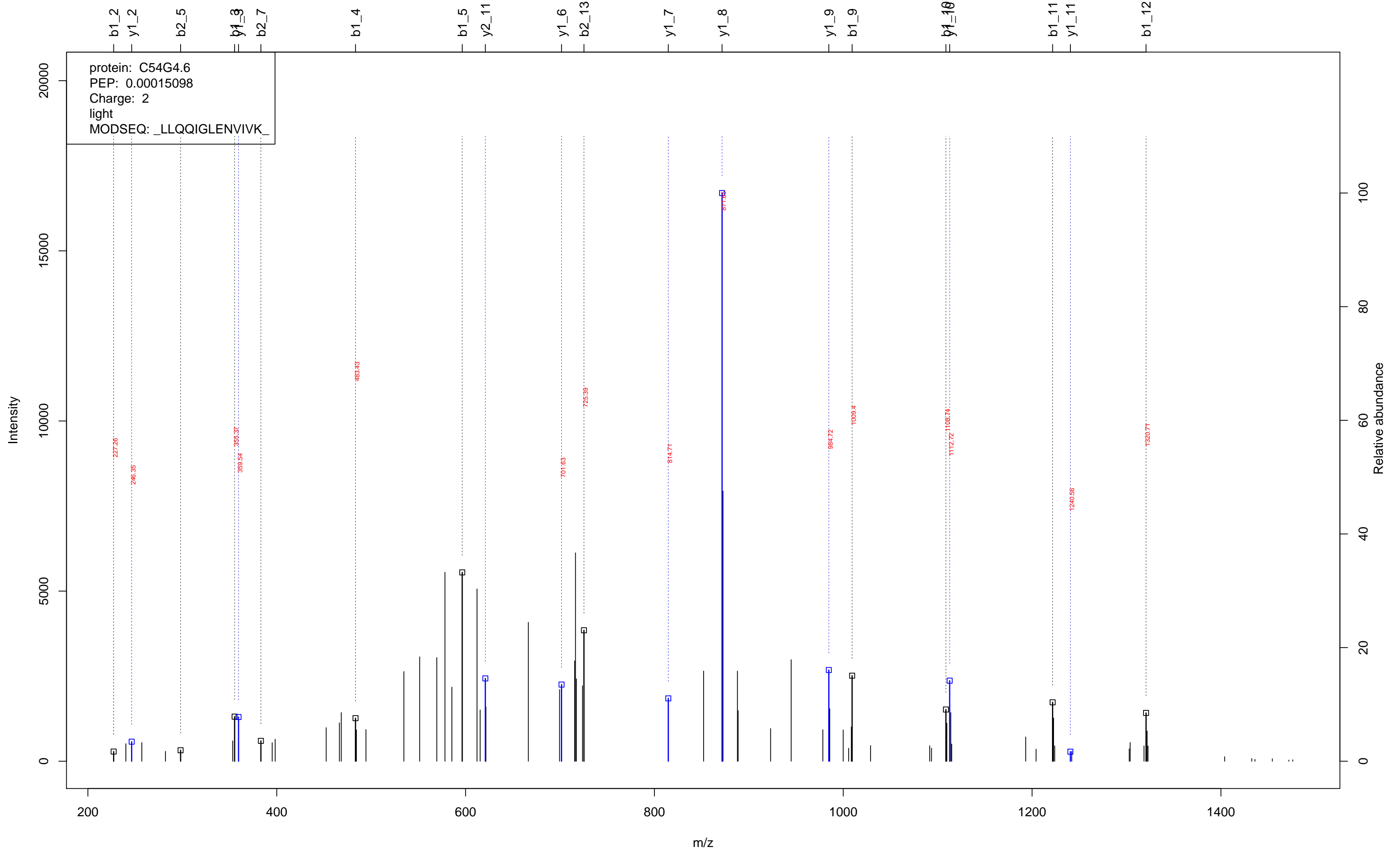


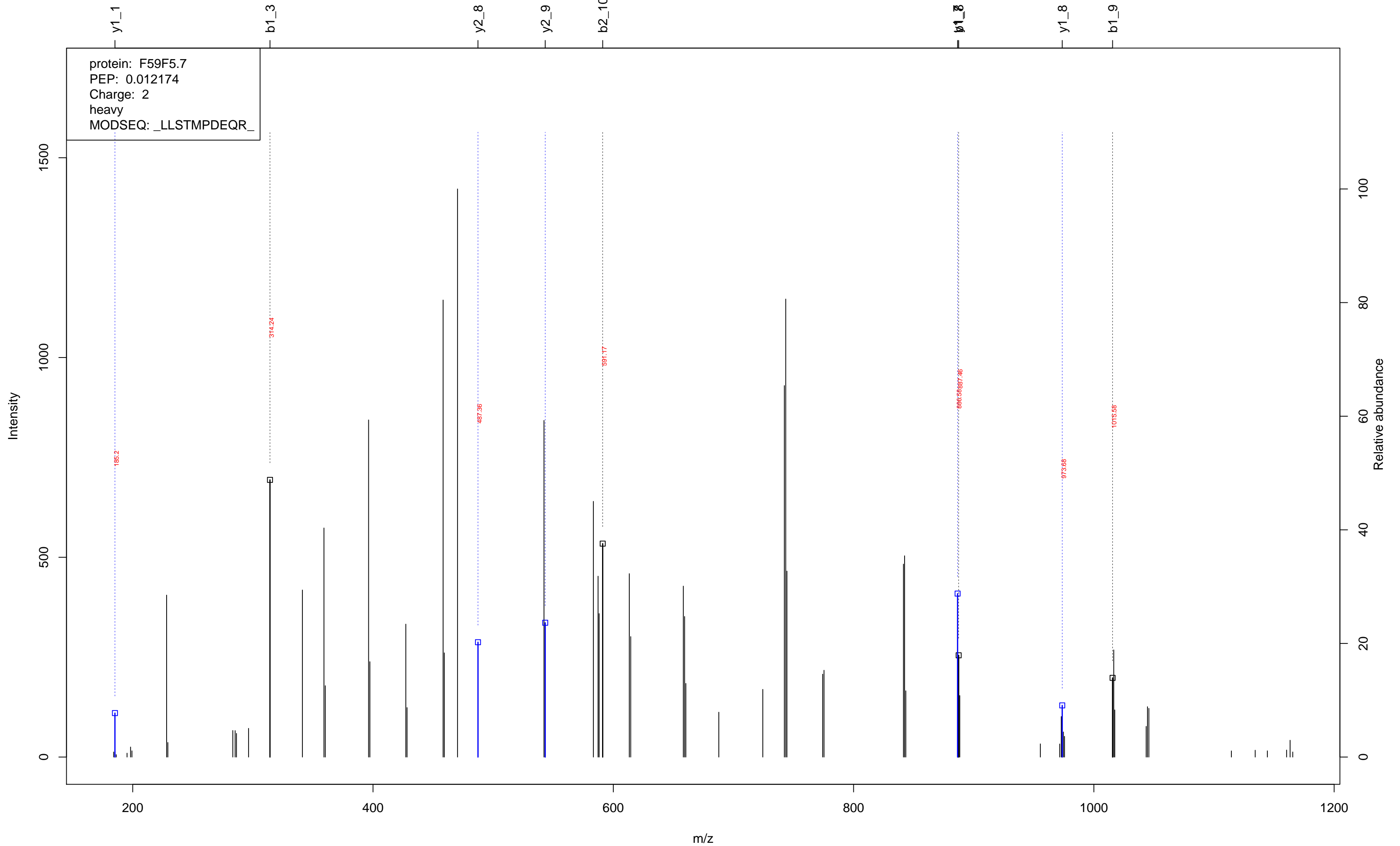


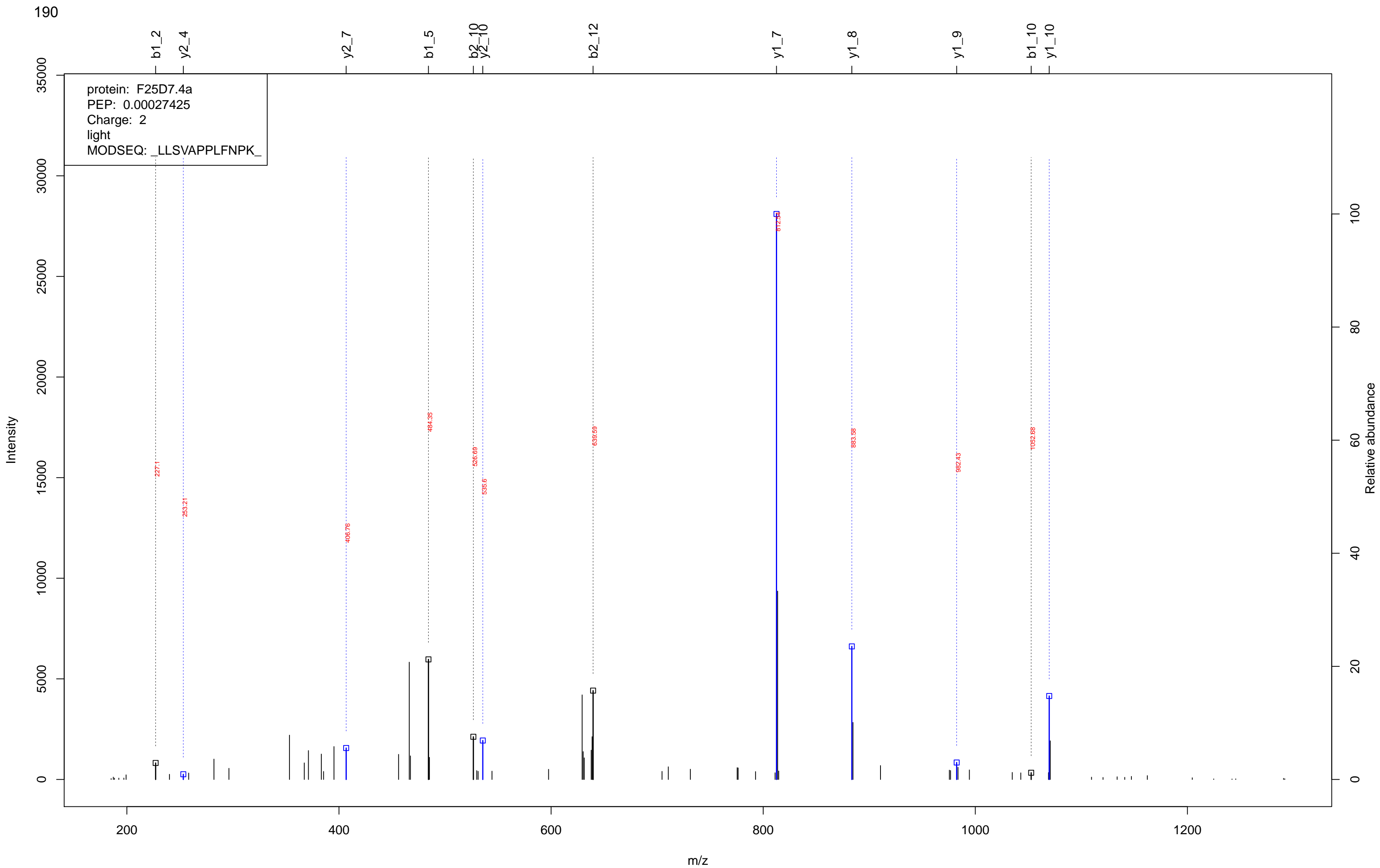


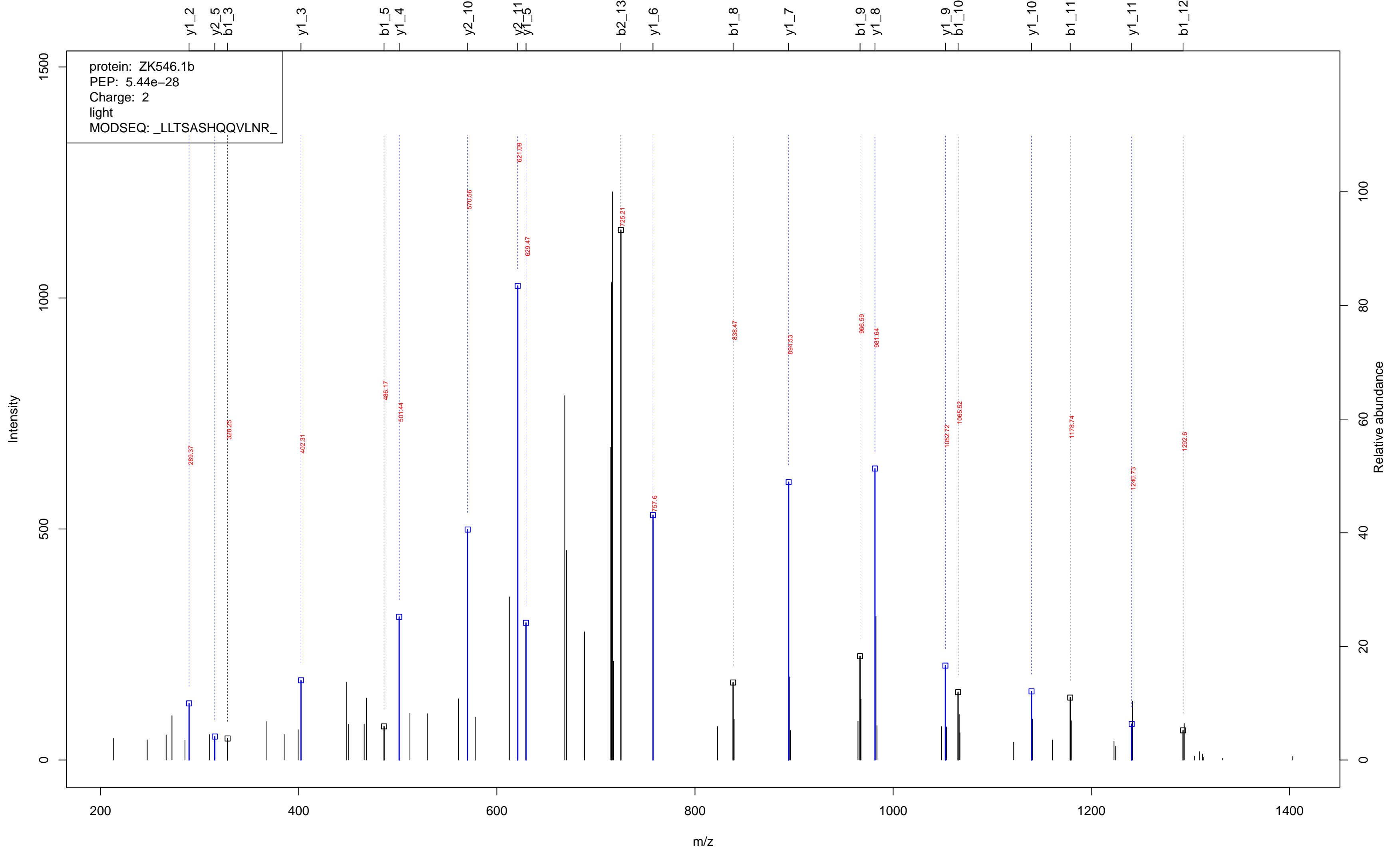


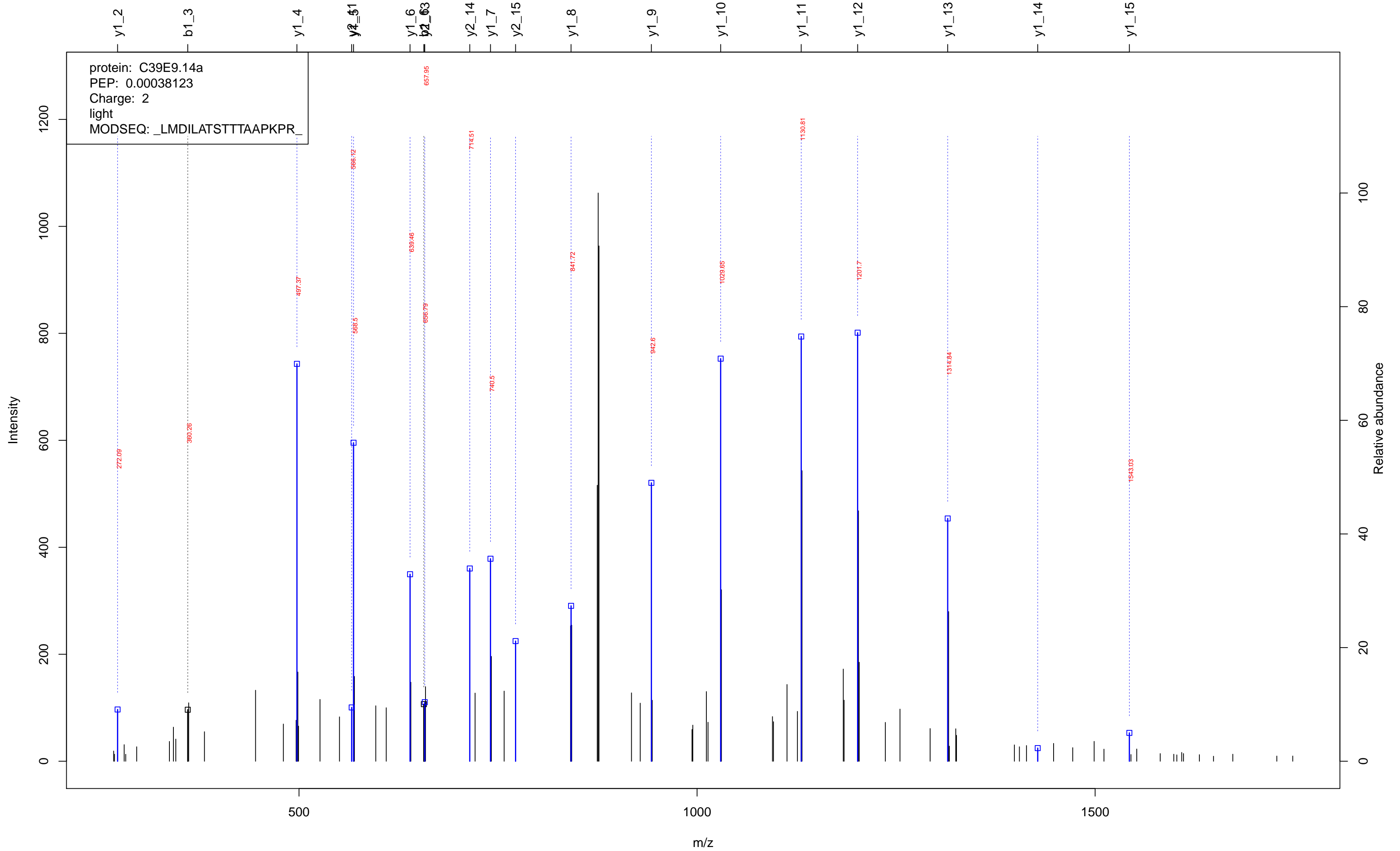


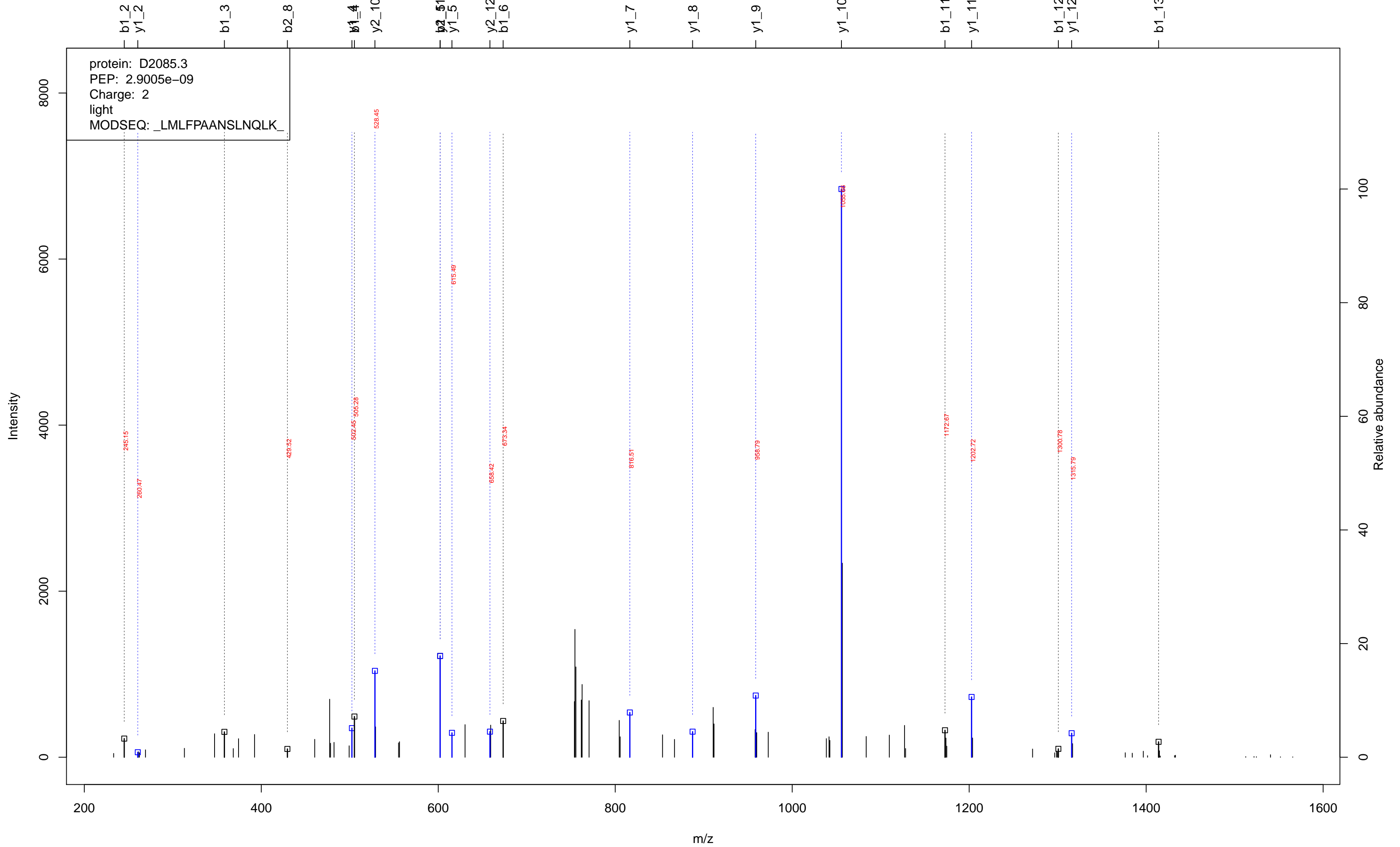


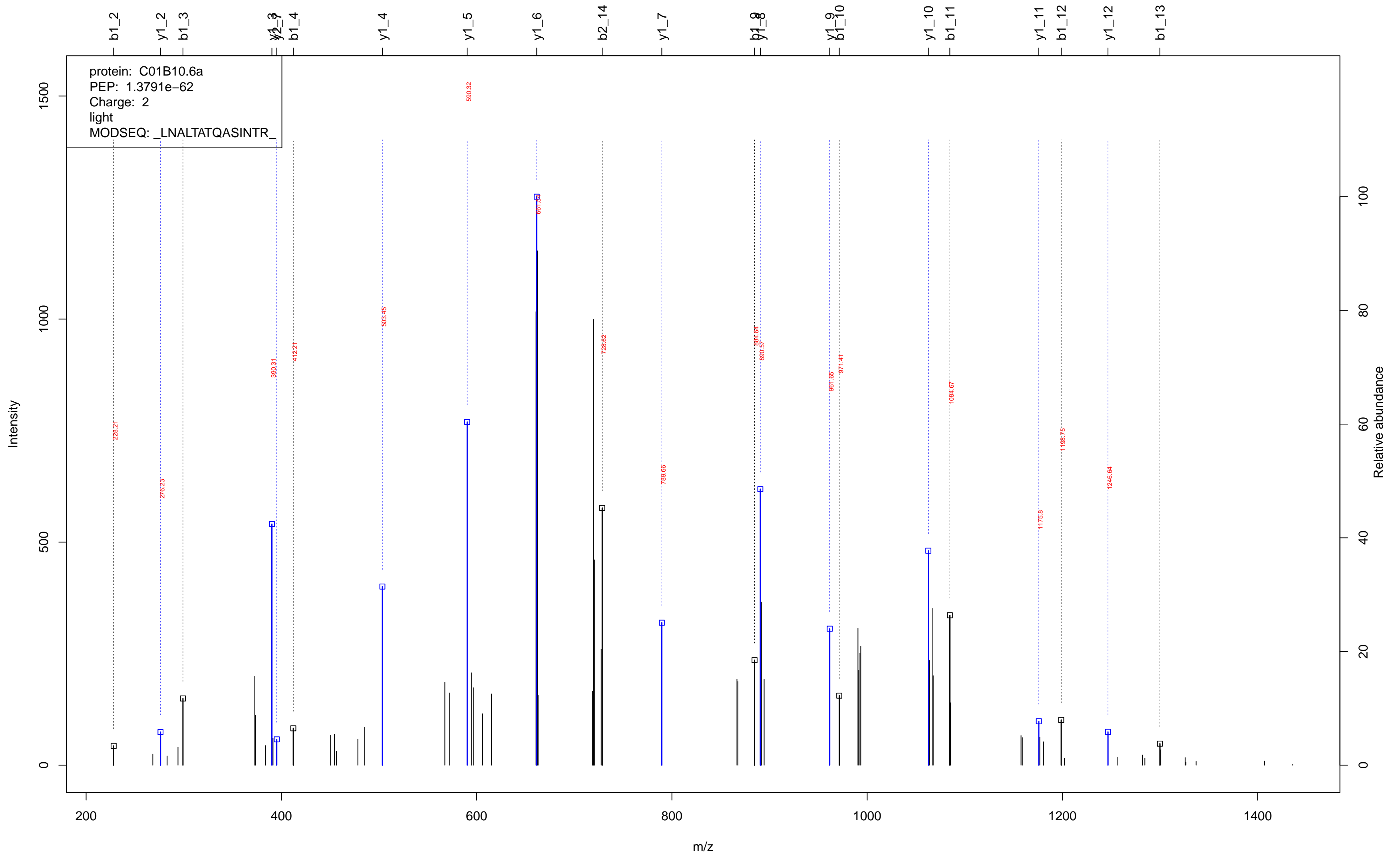




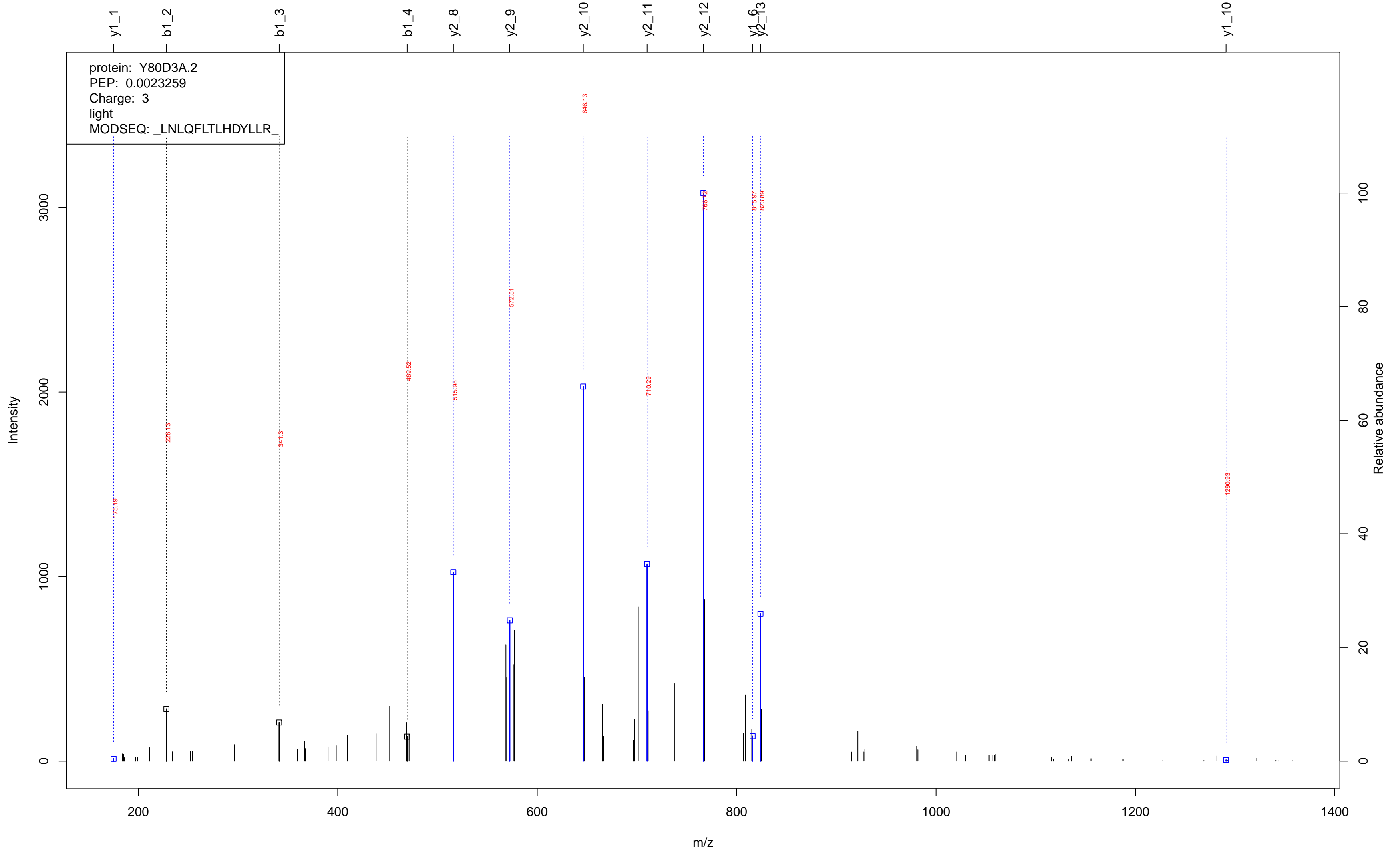


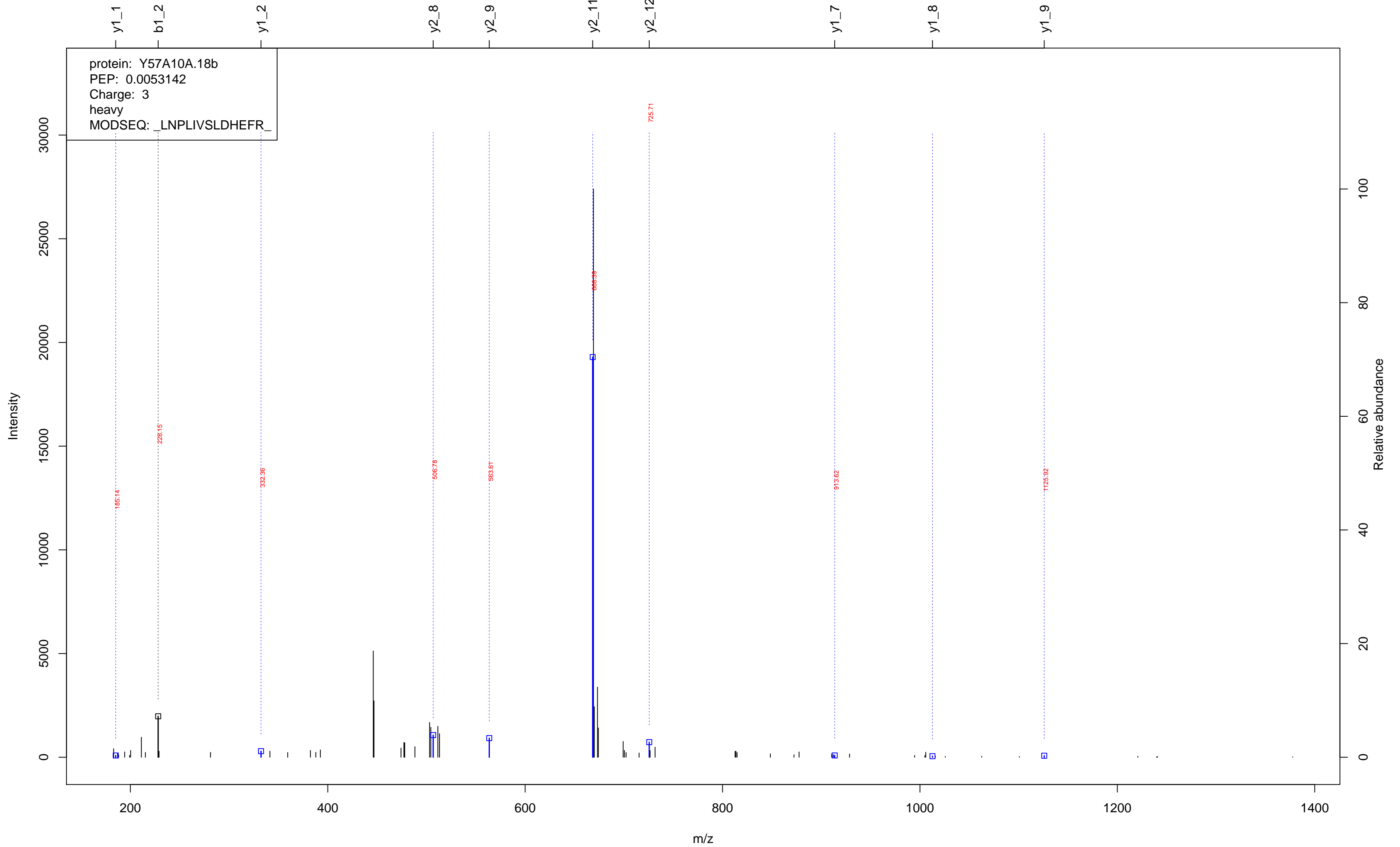


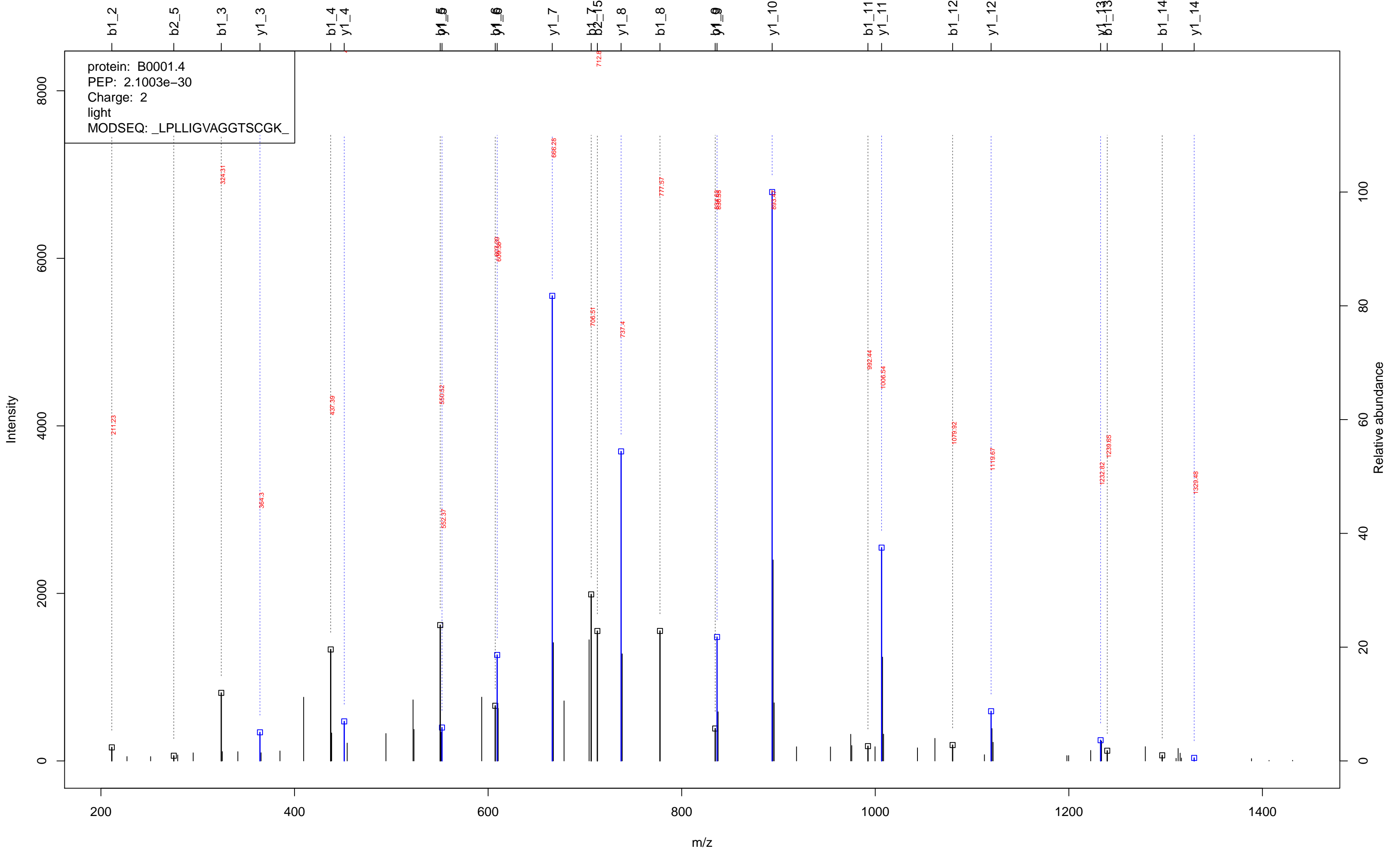


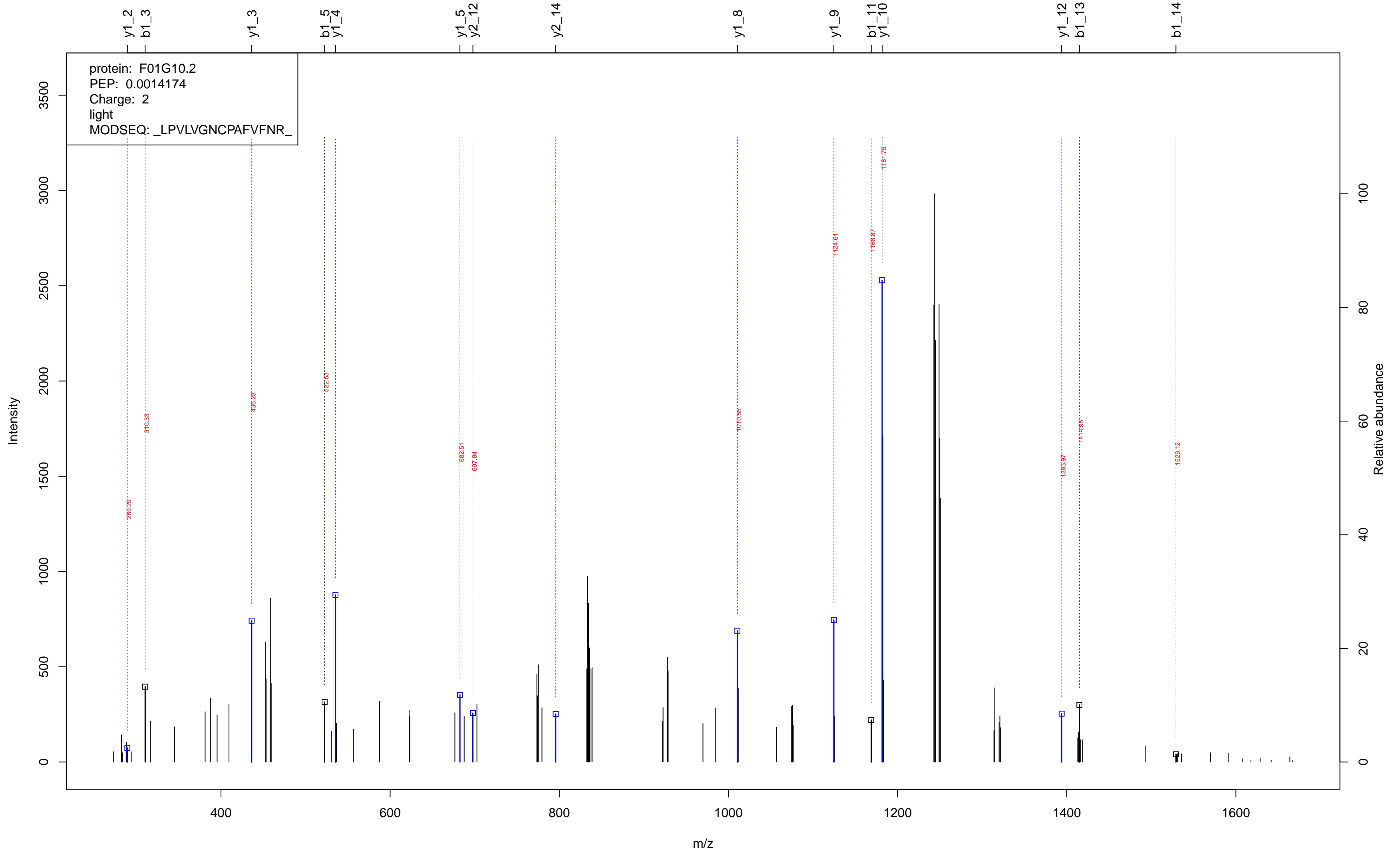


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Charge: 2
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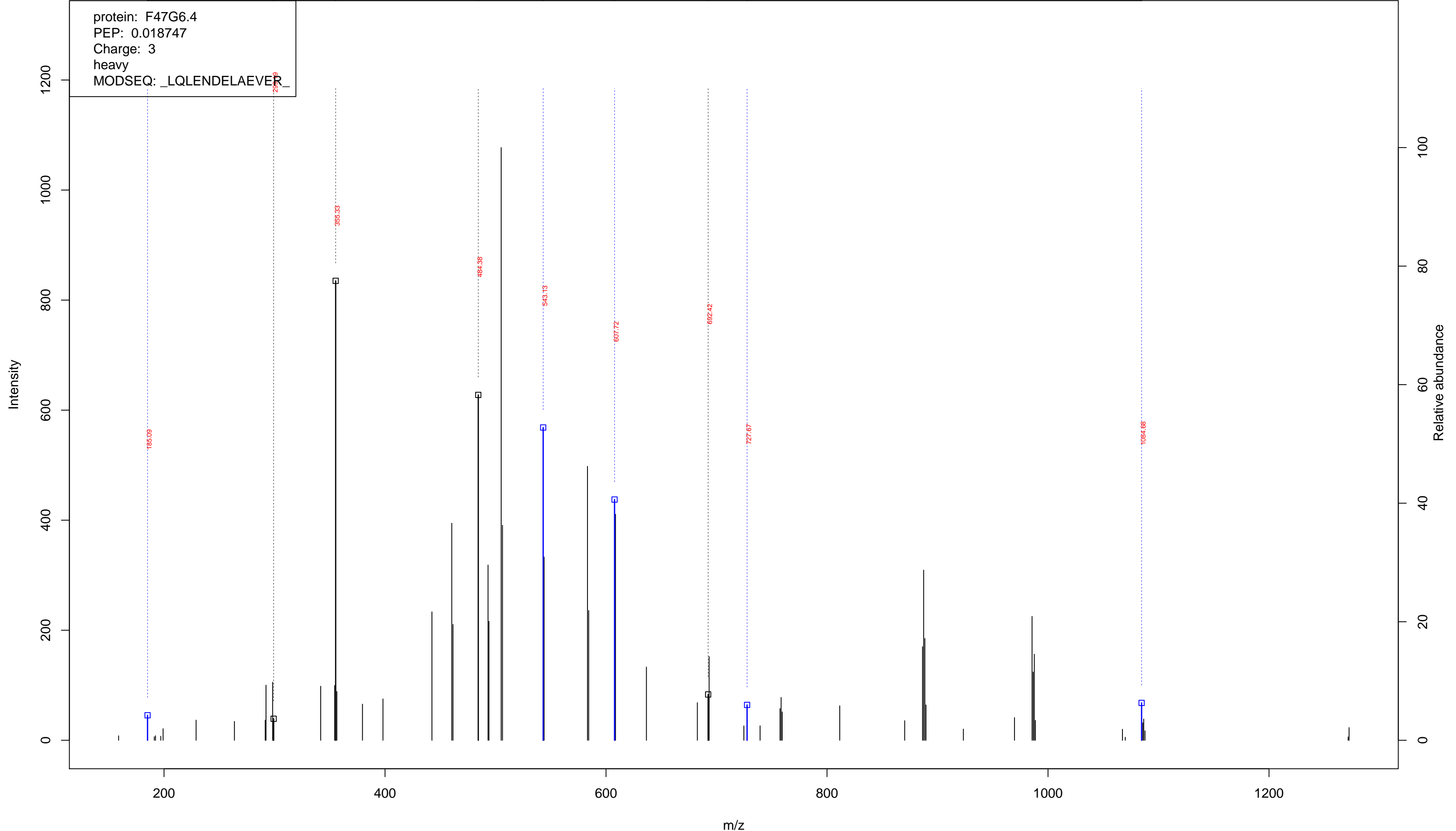


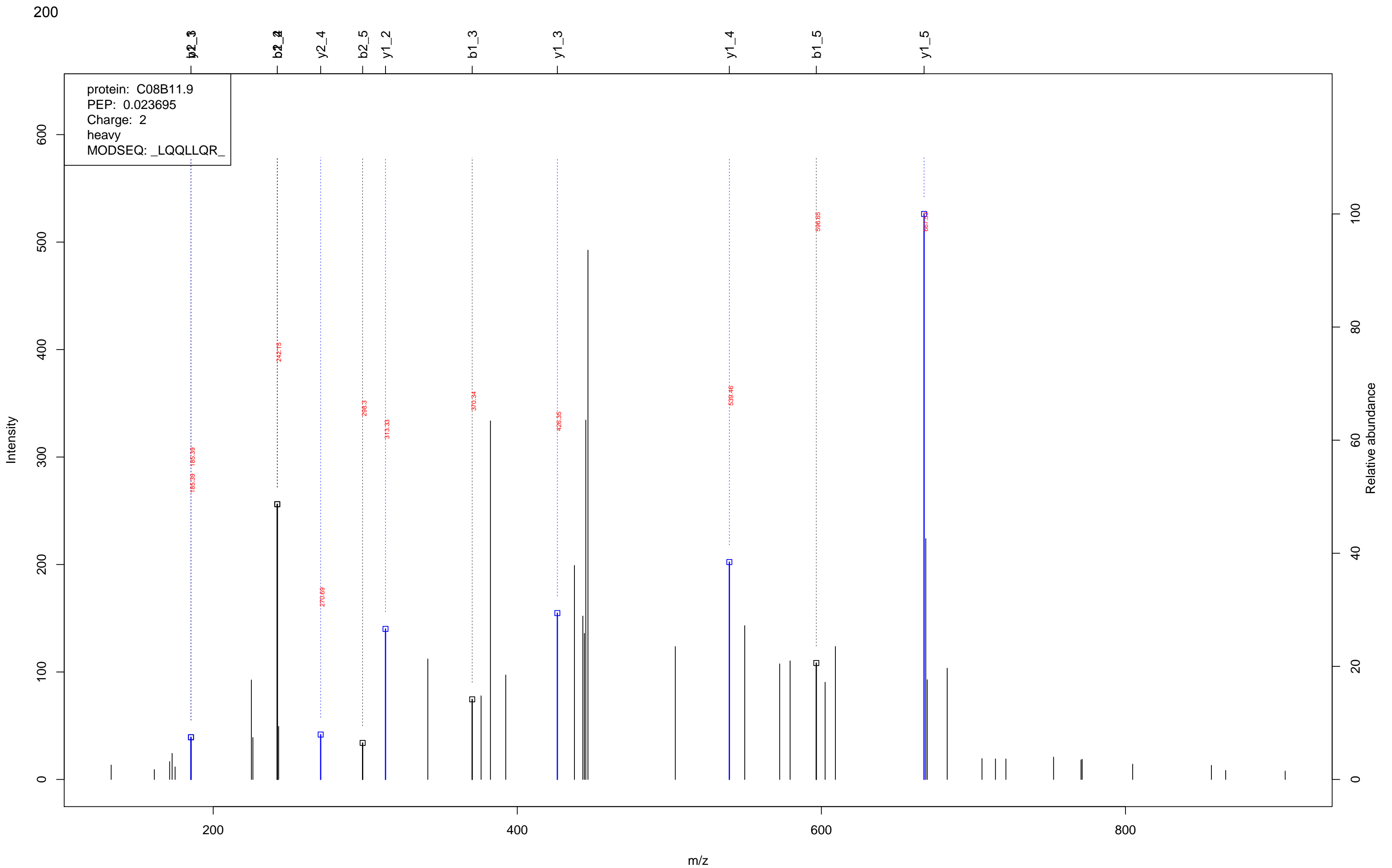




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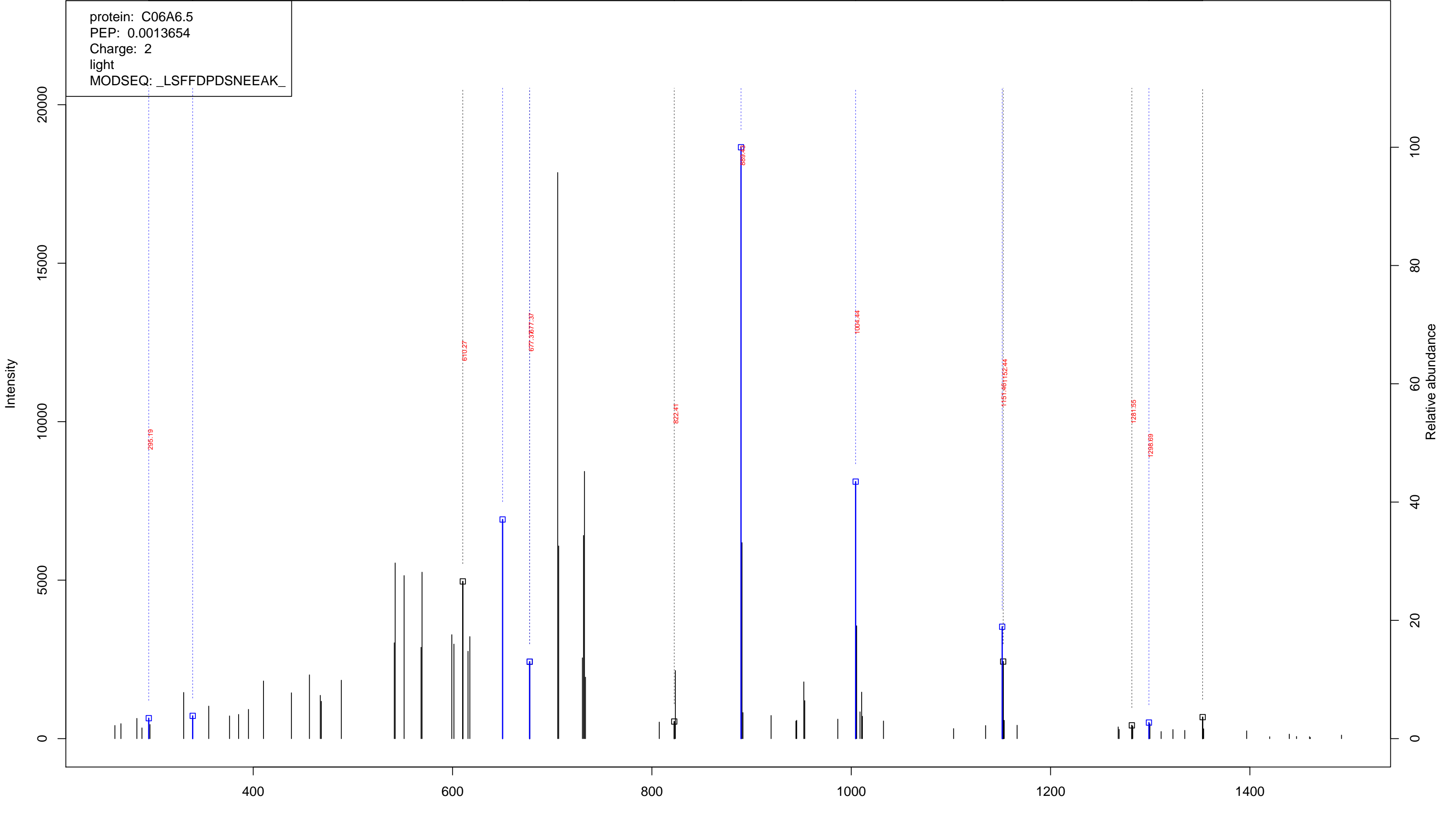
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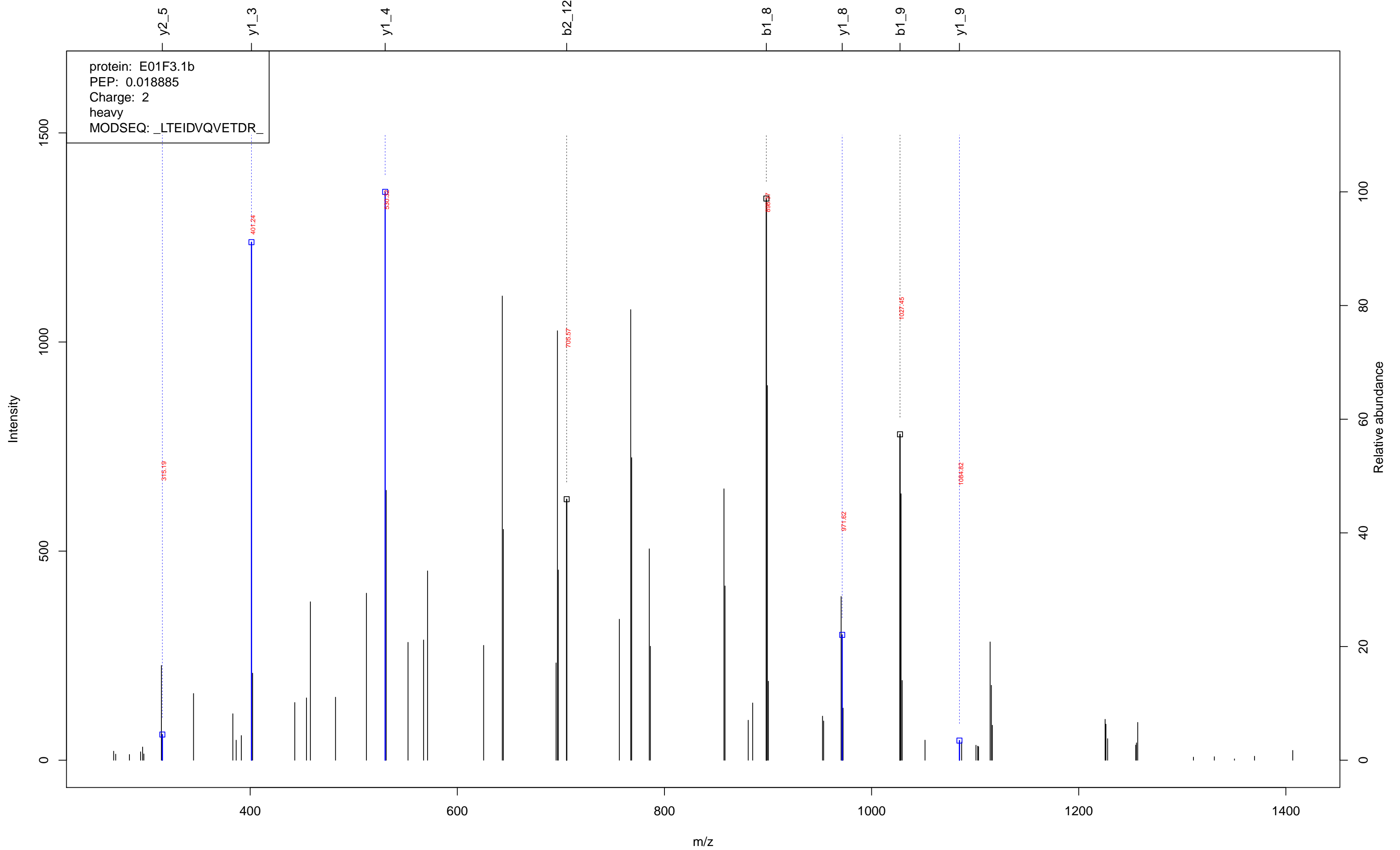


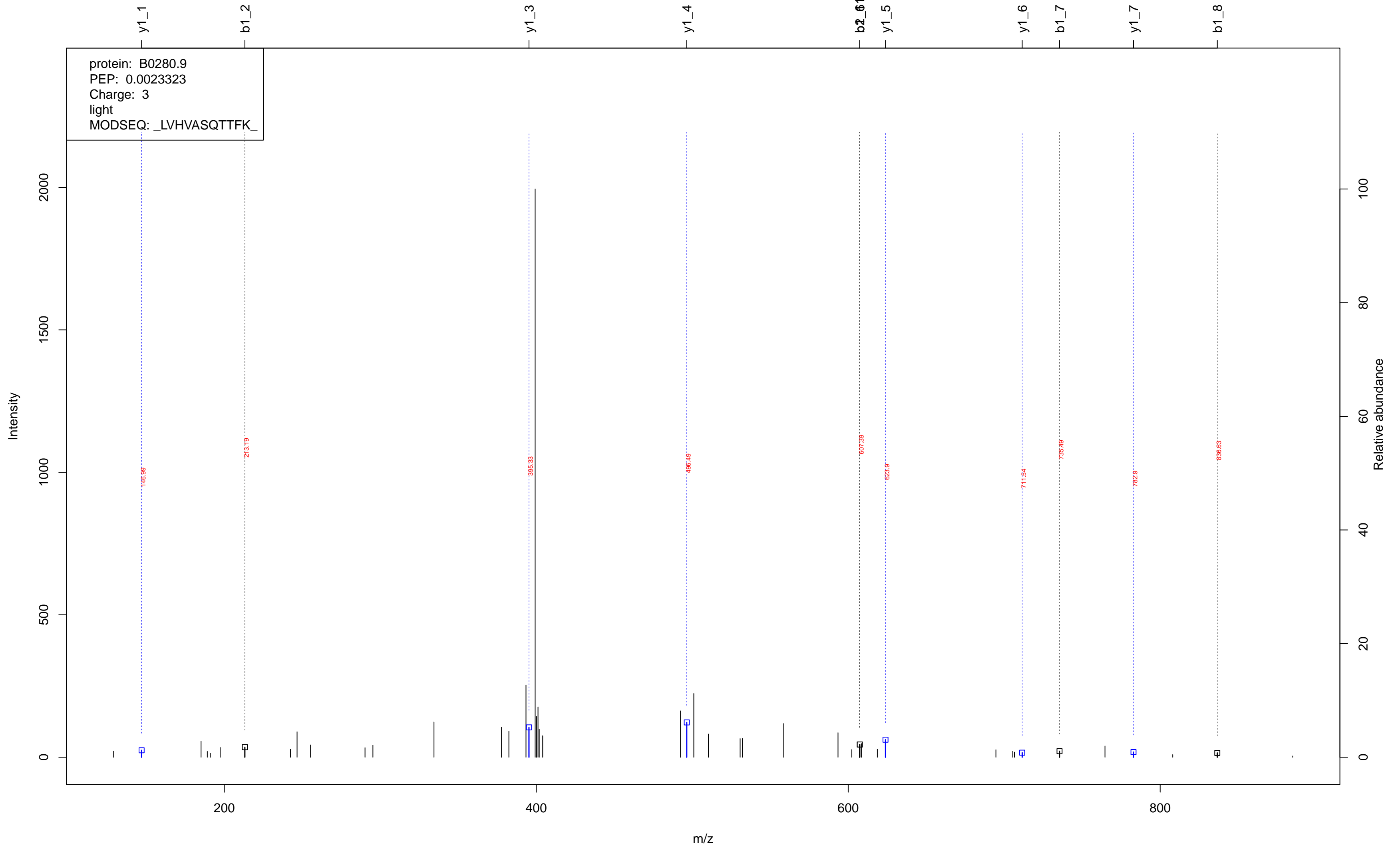


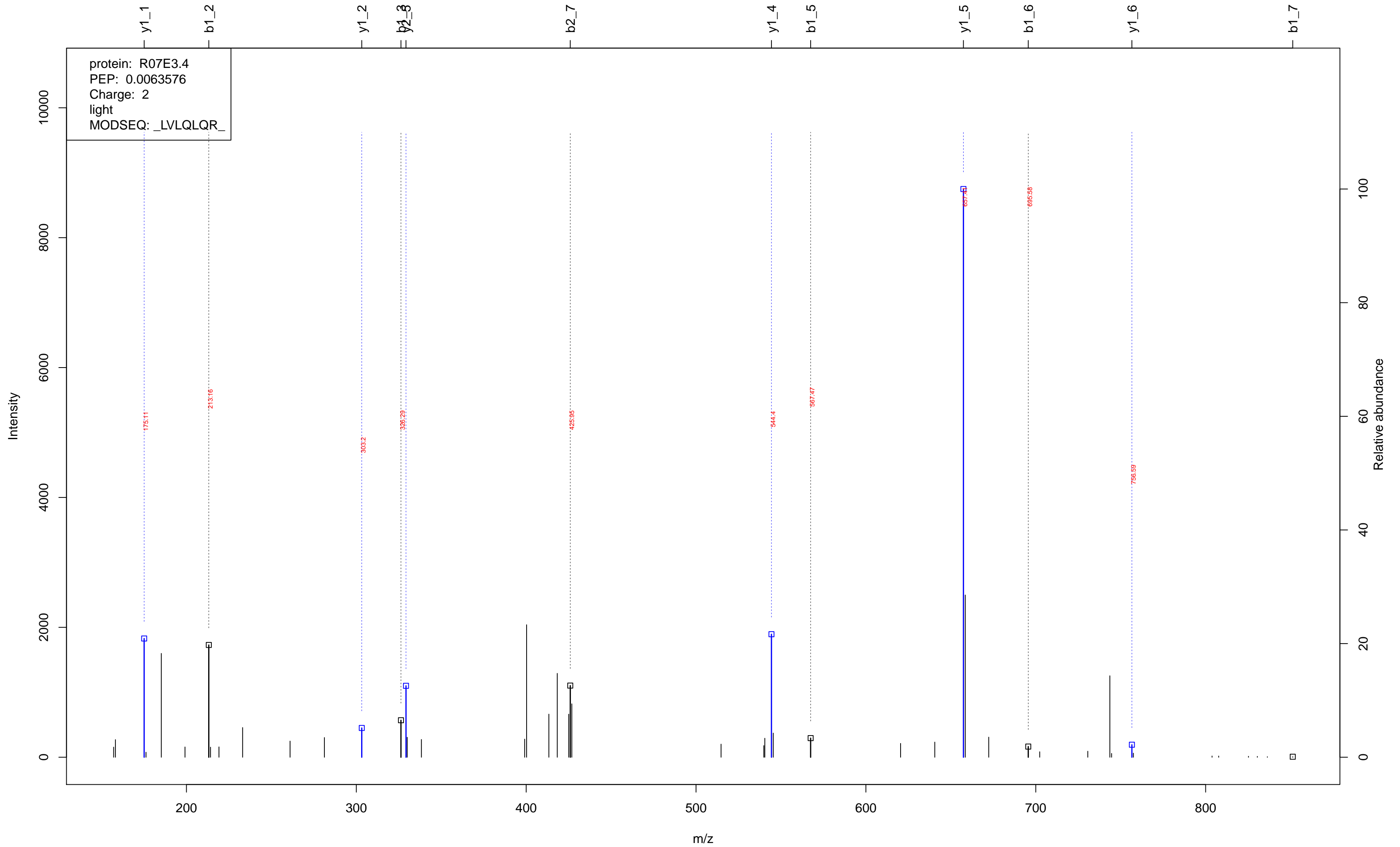
y2_5 y2_6 b1_5 y2_11 ~~y2_62~~ b1_7 y1_8 y1_9 ~~y1_10~~ b1_11 y1_11 b1_12

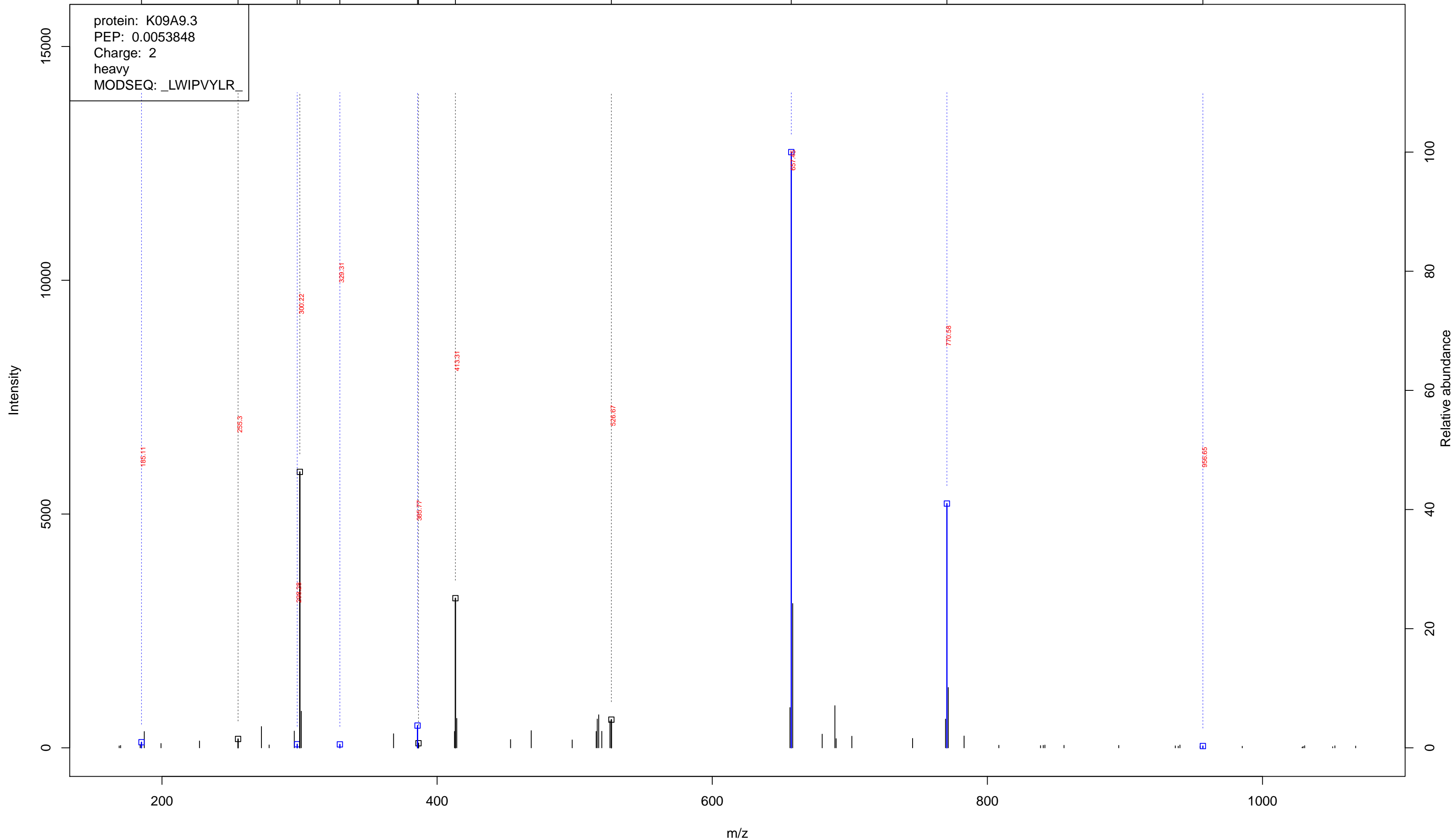
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Charge: 2
light
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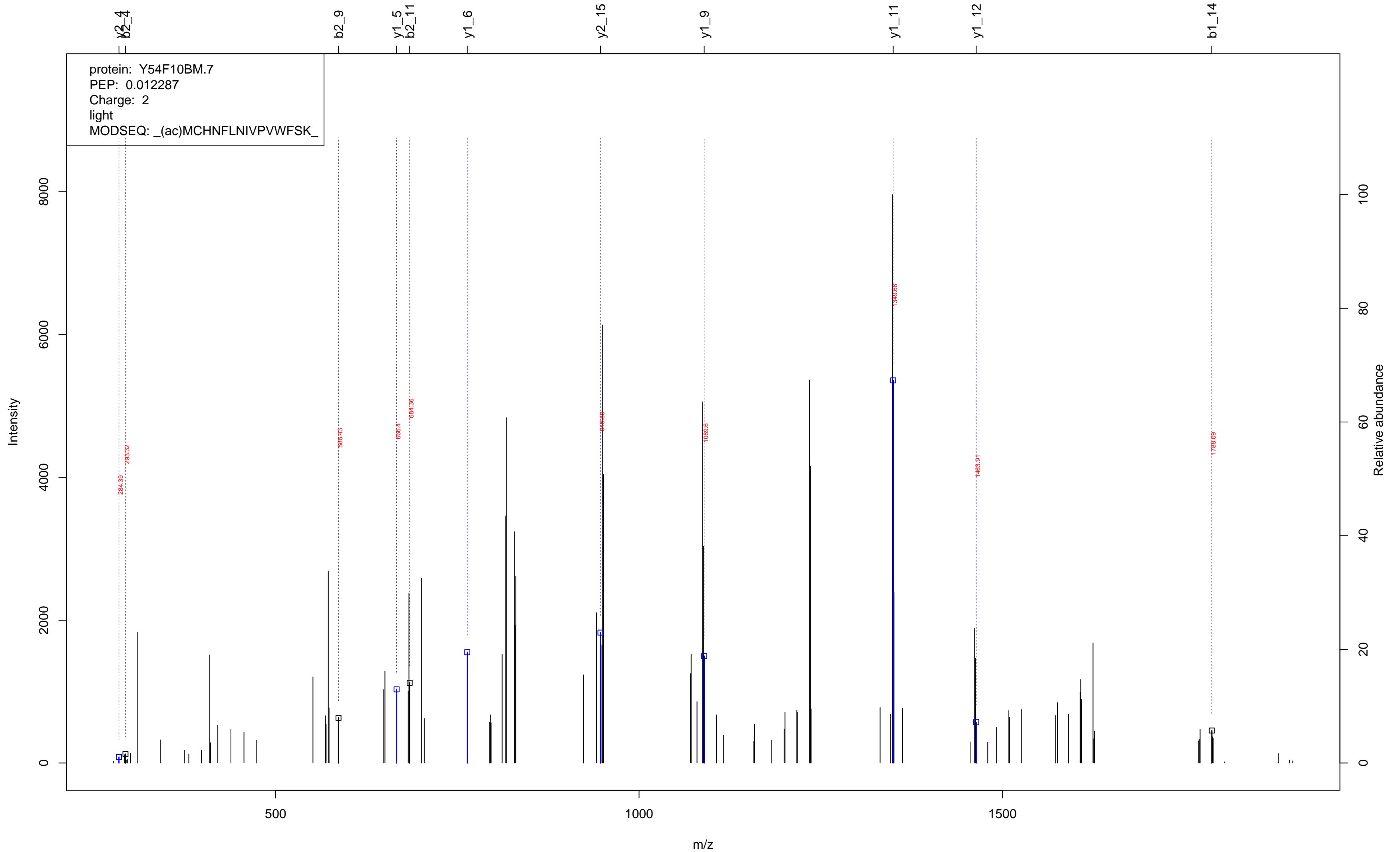


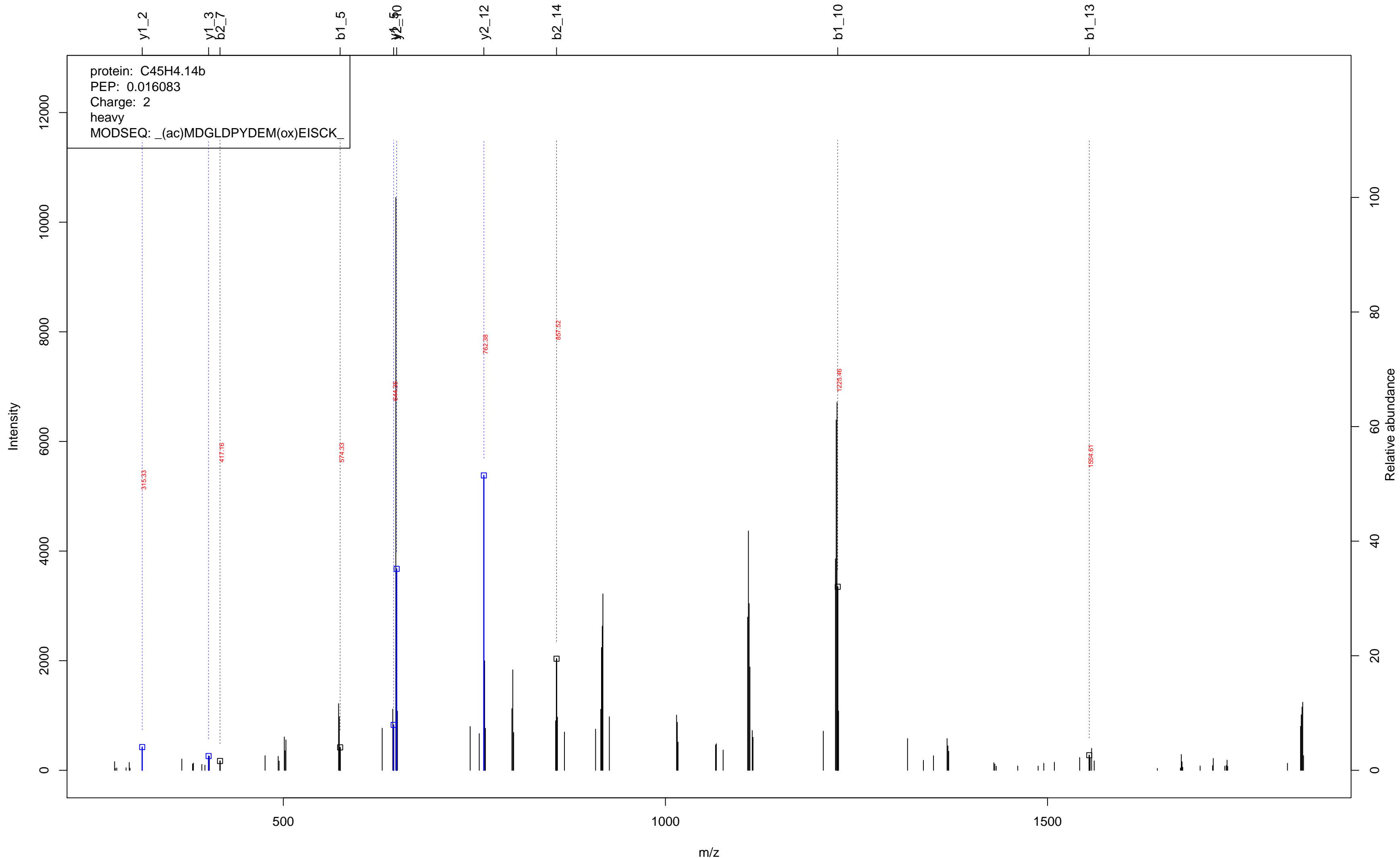






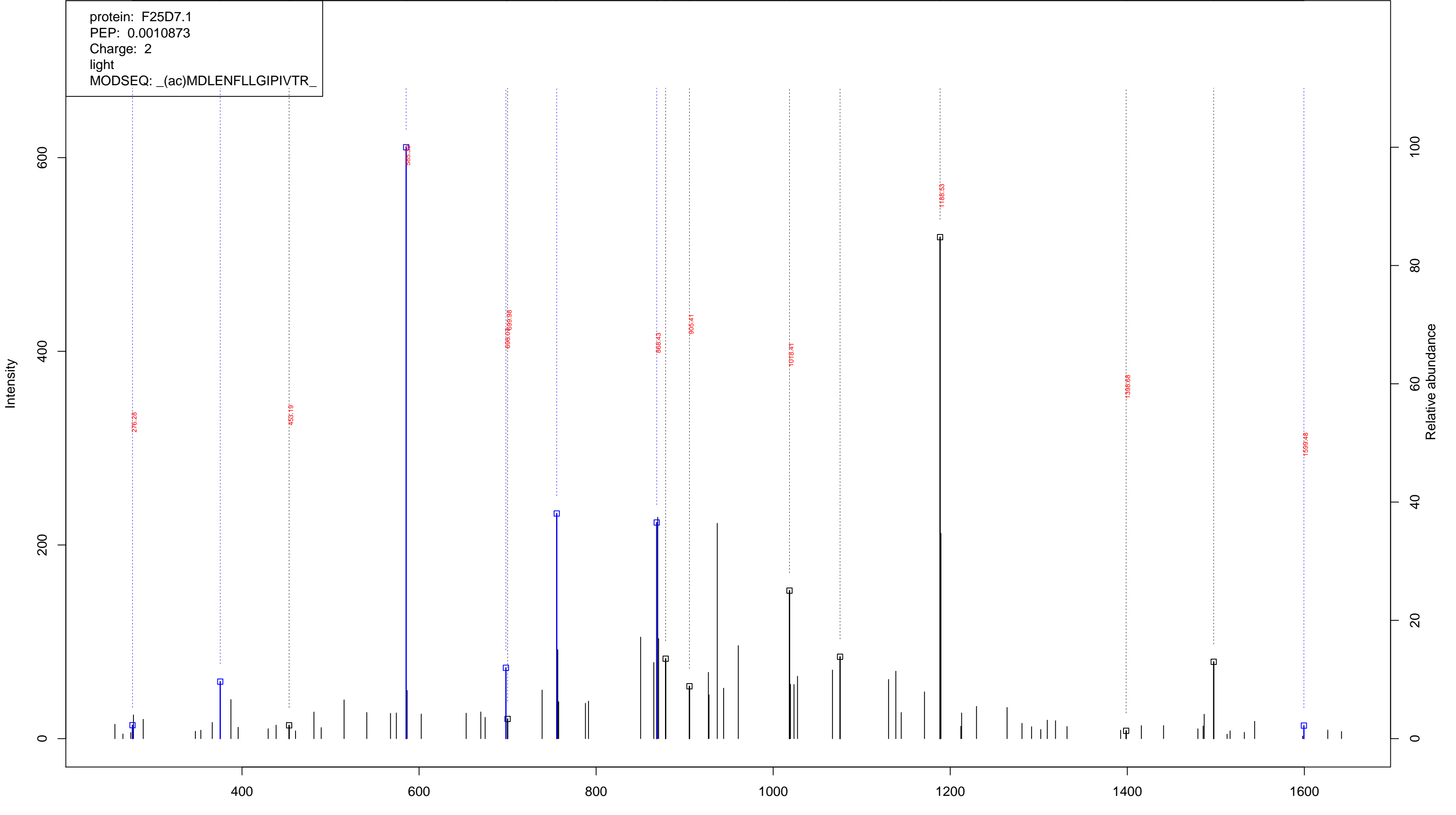


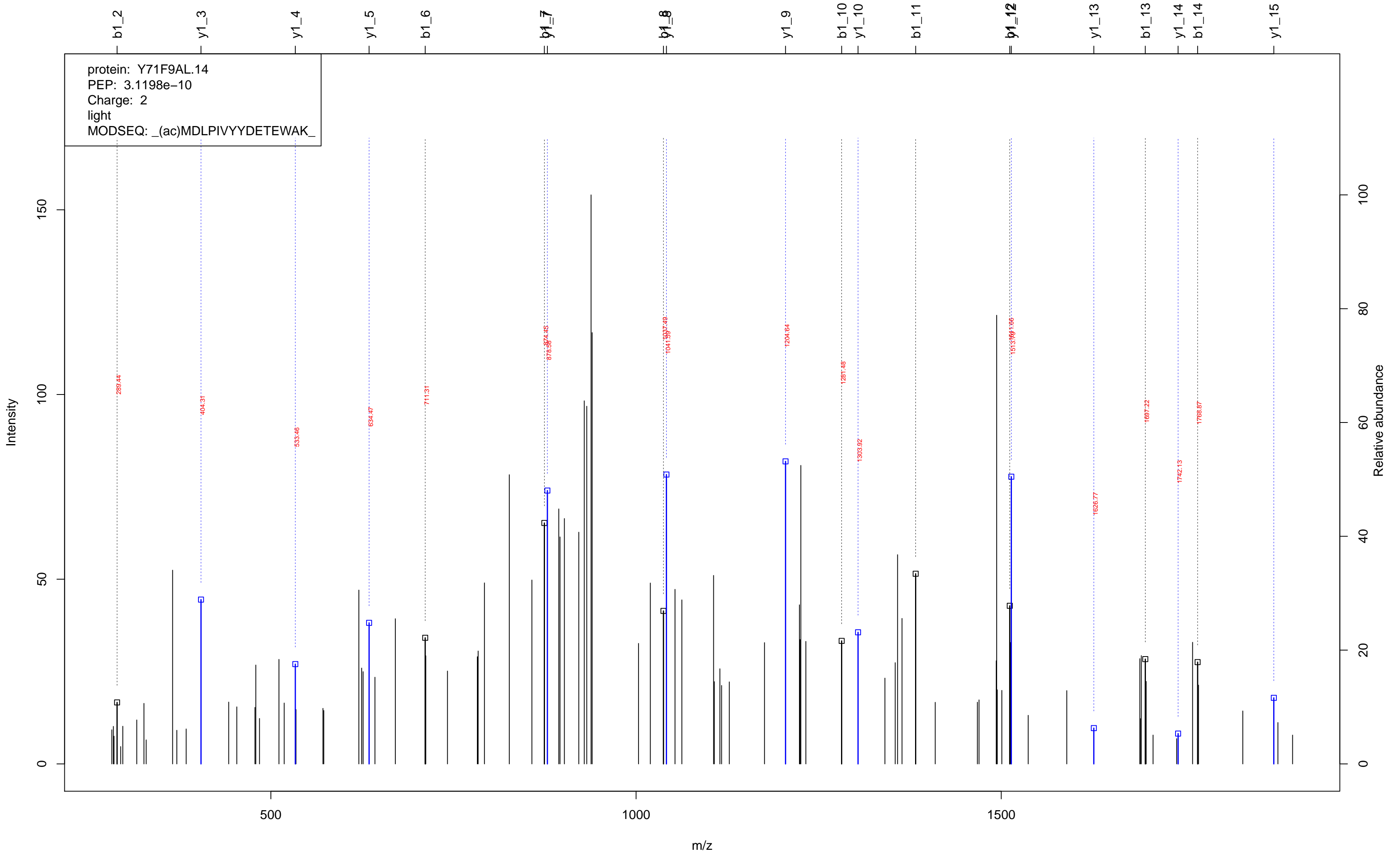


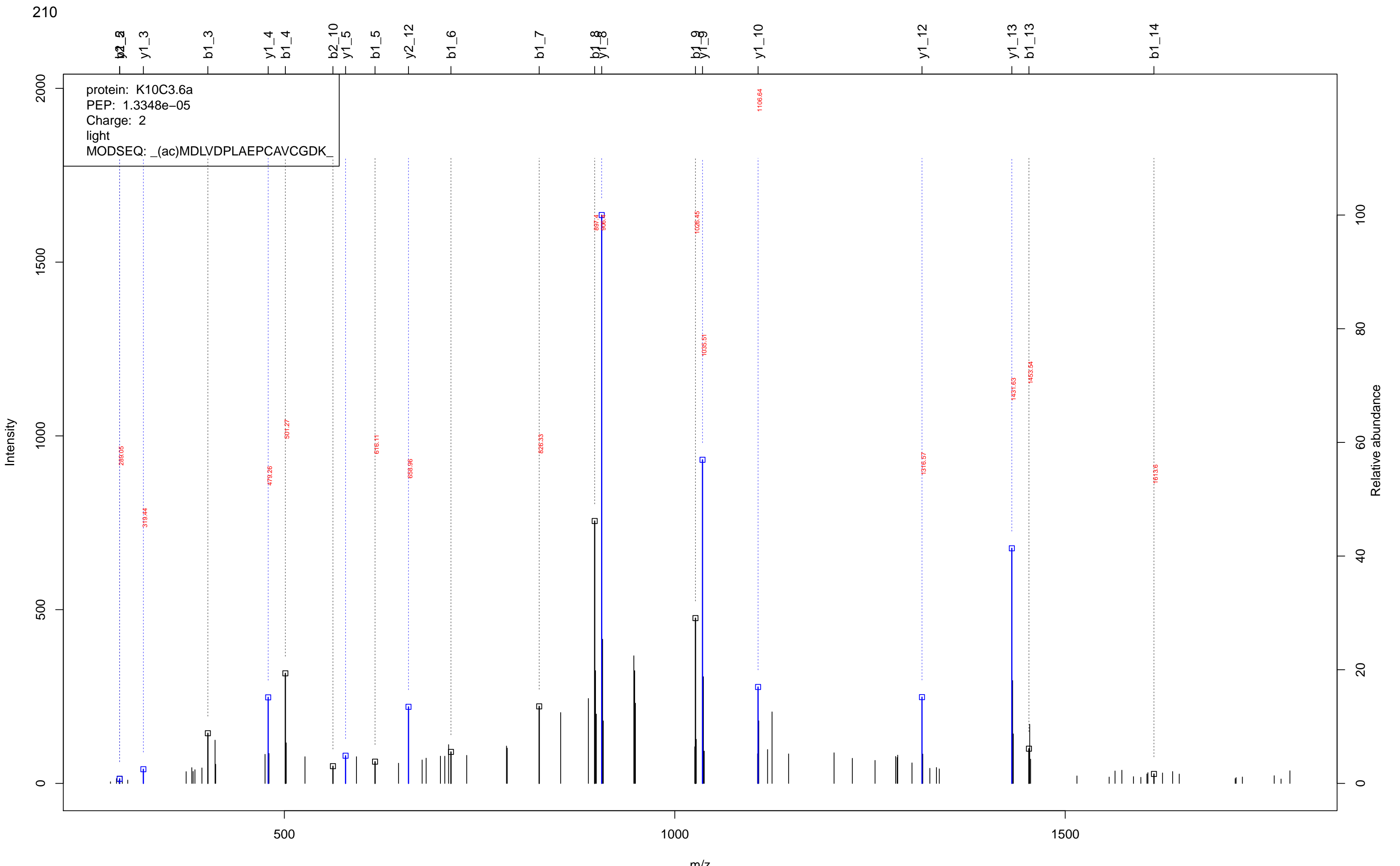


y1_2 y1_3 b2_7 y1_5 y1_6 y1_7 y1_8 b2_15 b1_7 b1_8 b1_9 b1_10 b1_12 b1_13 y1_14

protein: F25D7.1
PEP: 0.0010873
Charge: 2
light
MODSEQ: _(ac)MDLENFLLGIPIVTR_

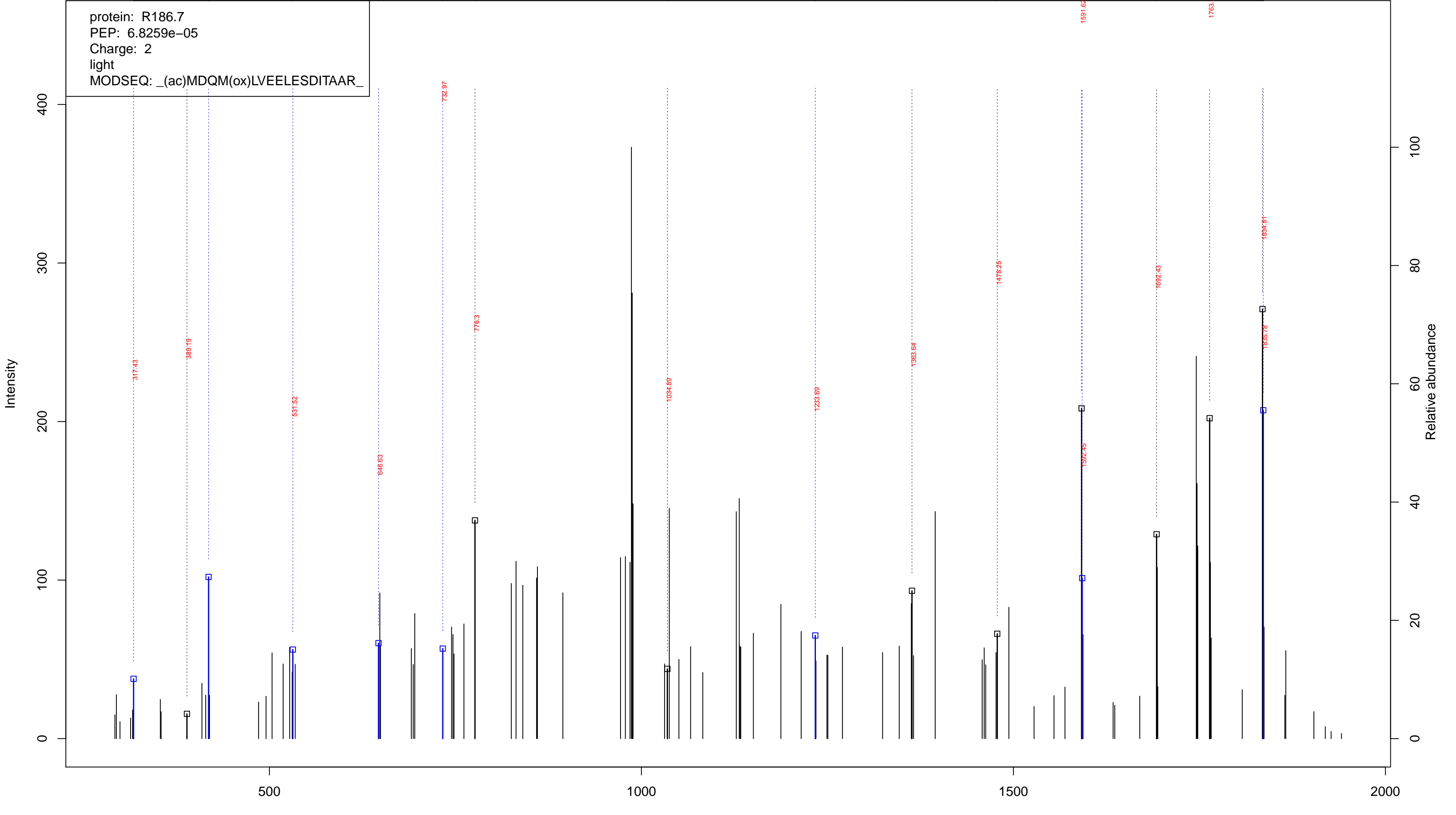


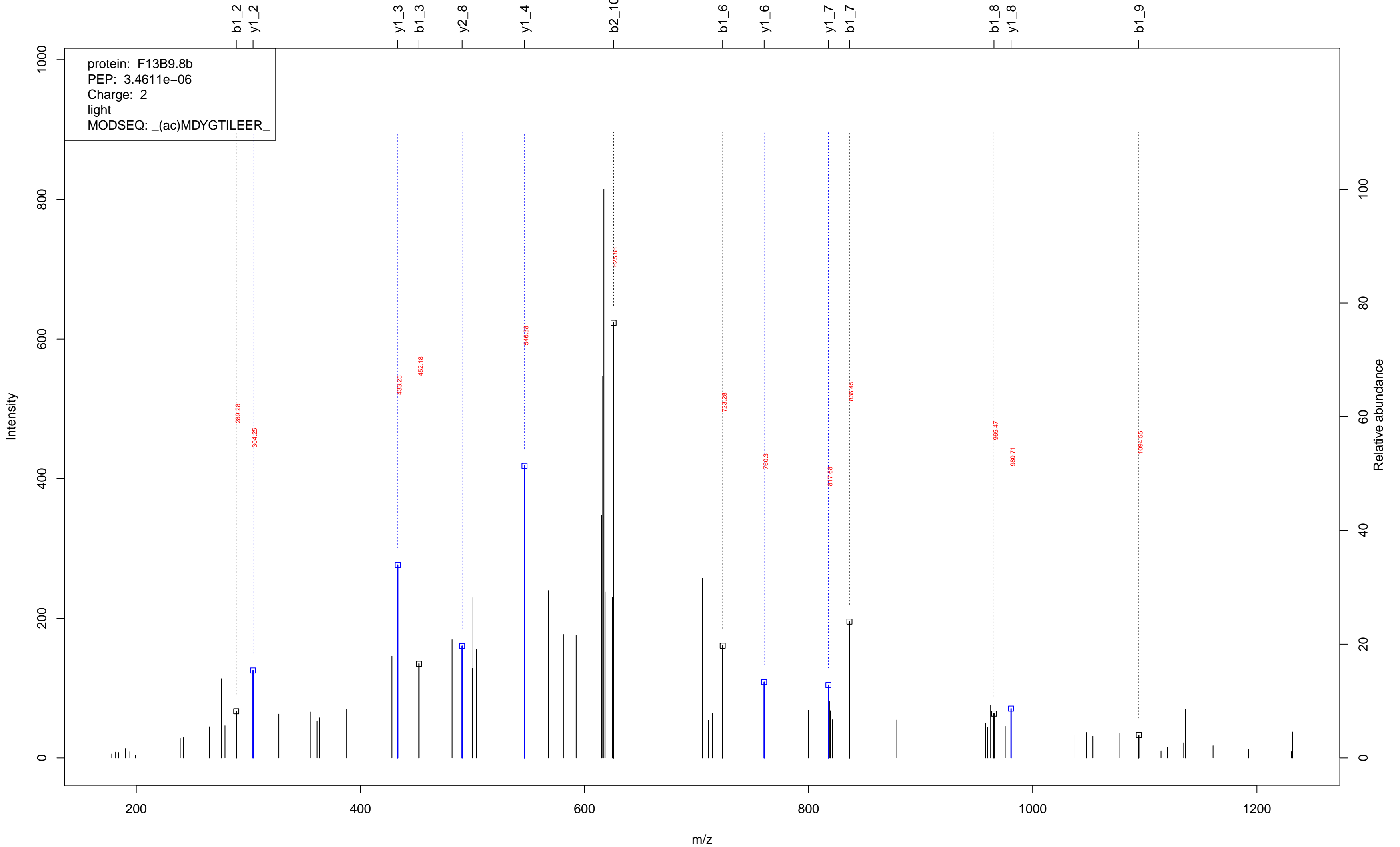


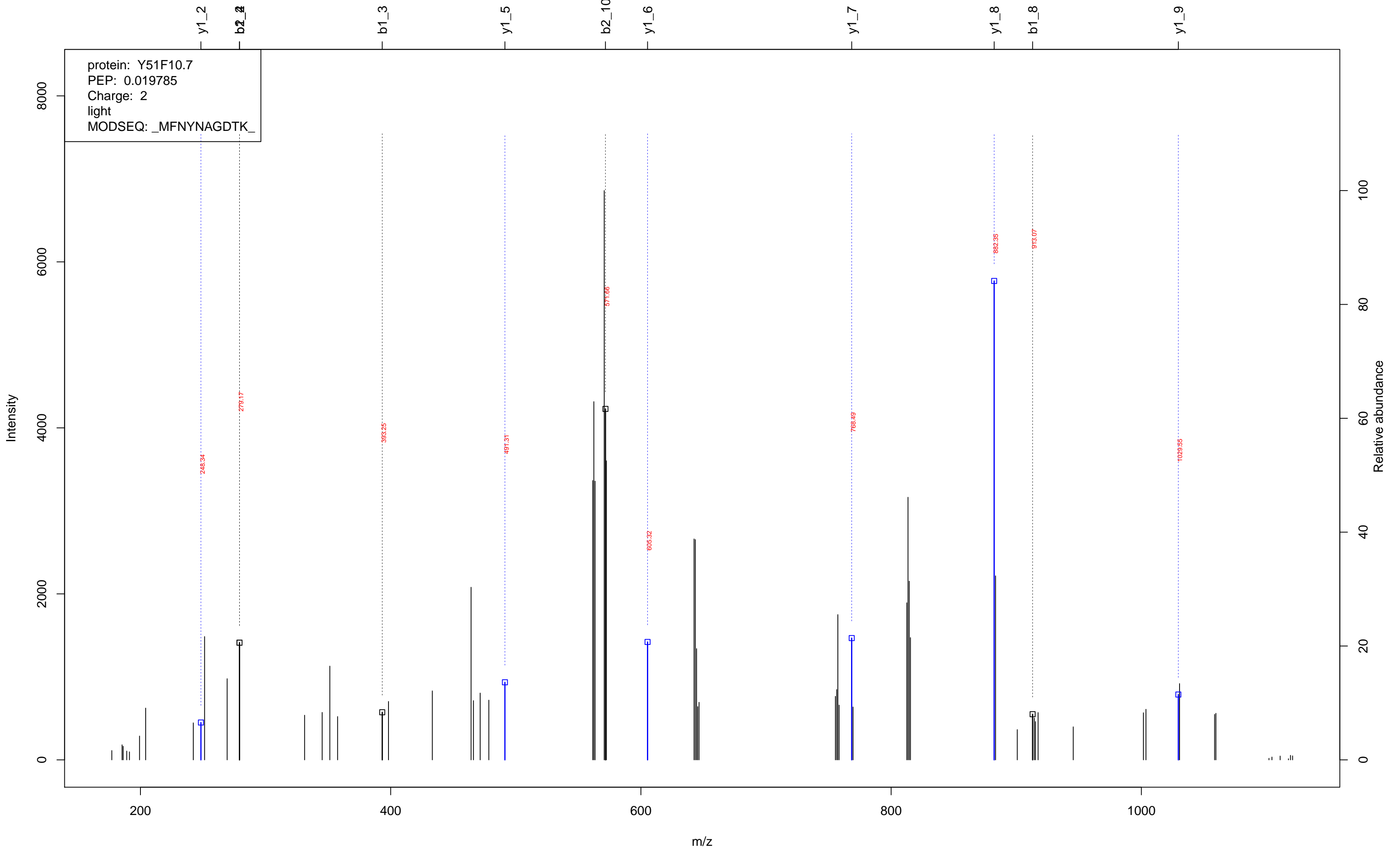


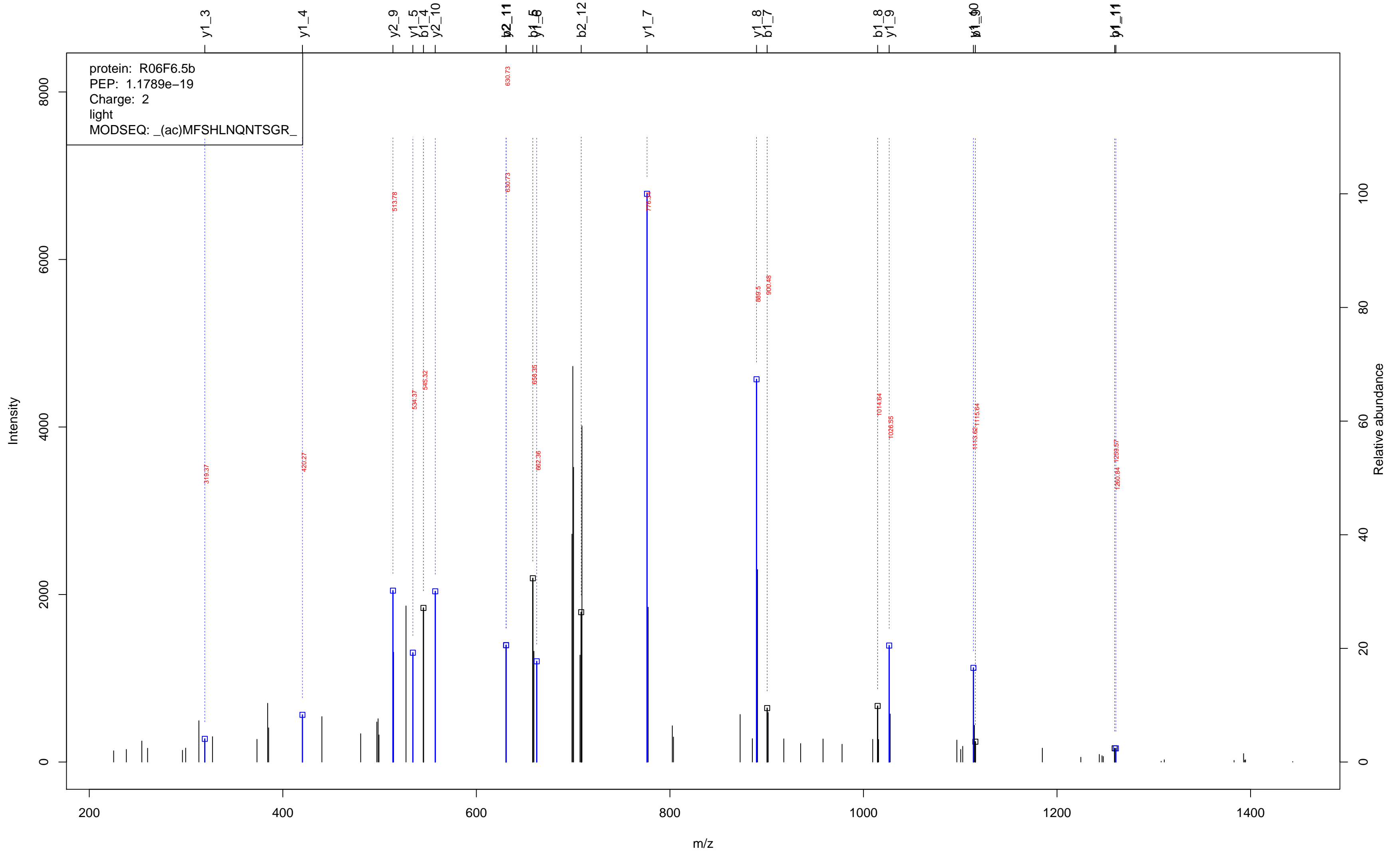
y1_3 b2_6 y1_4 y1_5 y1_6 y1_7 b1_6 b1_8 y1_11 b1_11 b1_12 ~~y1_13~~ b1_14 b1_15 ~~b1_16~~

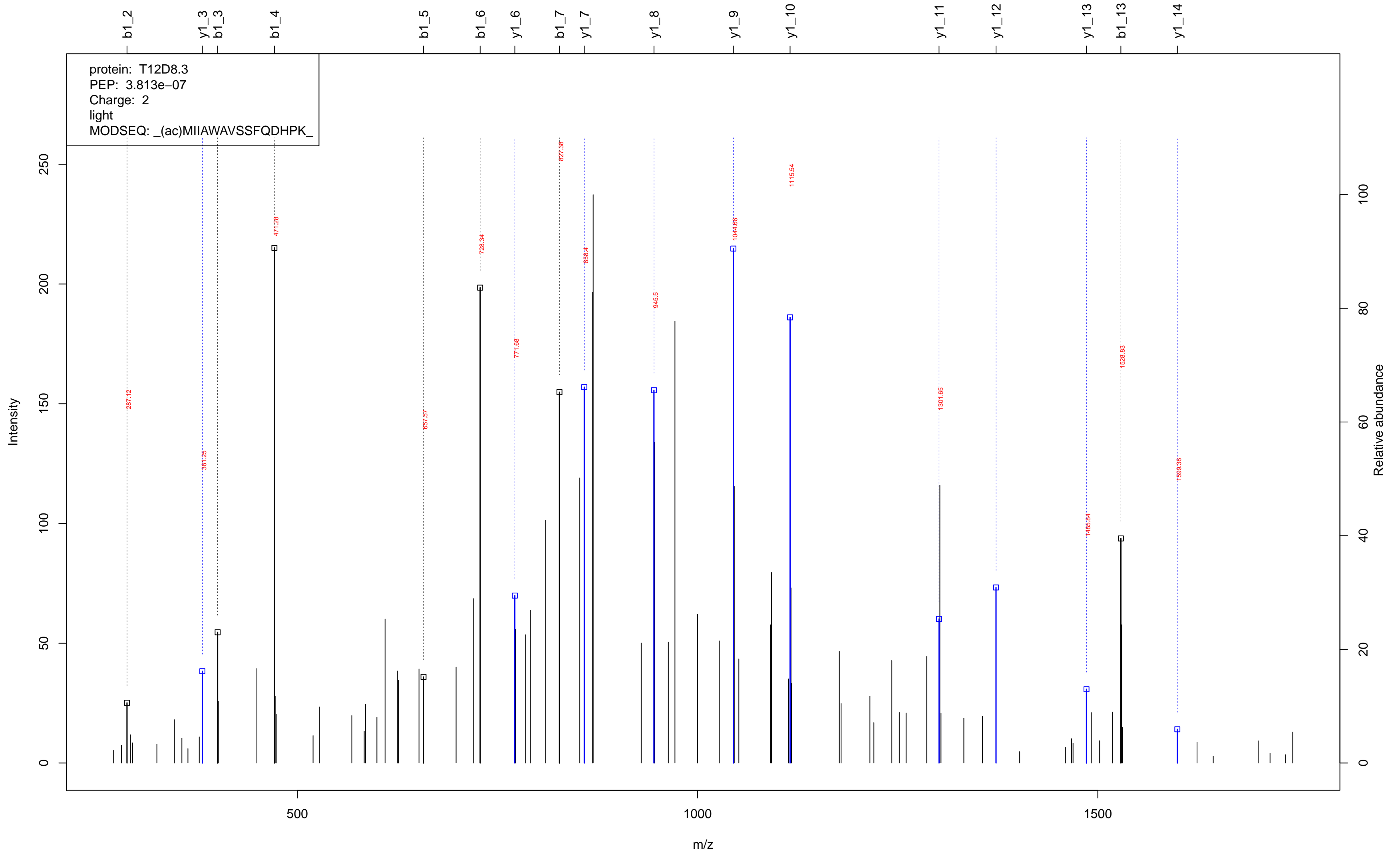
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Charge: 2
light
MODSEQ: (ac)MDQM(ox)LVEELESDITAAR_

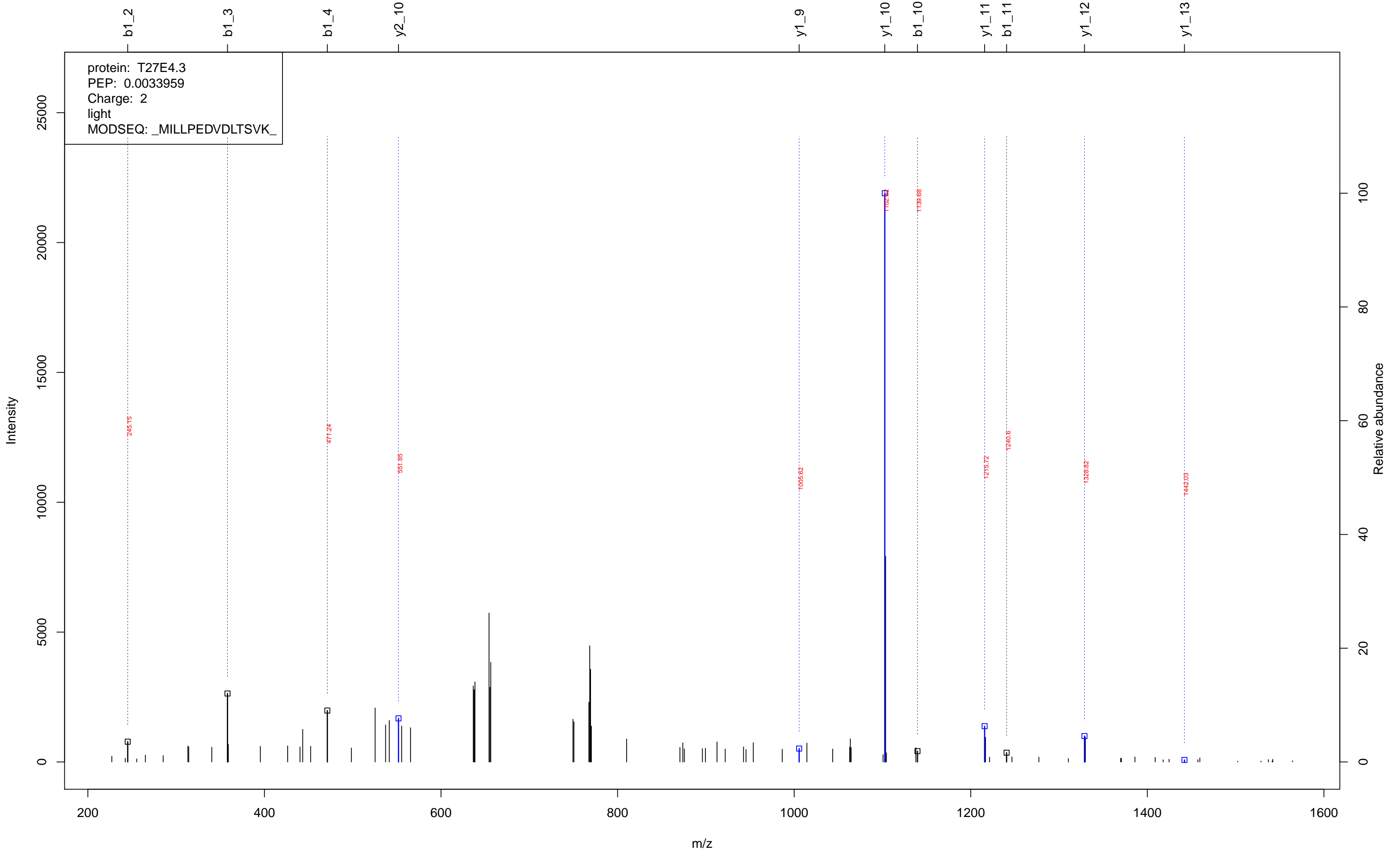


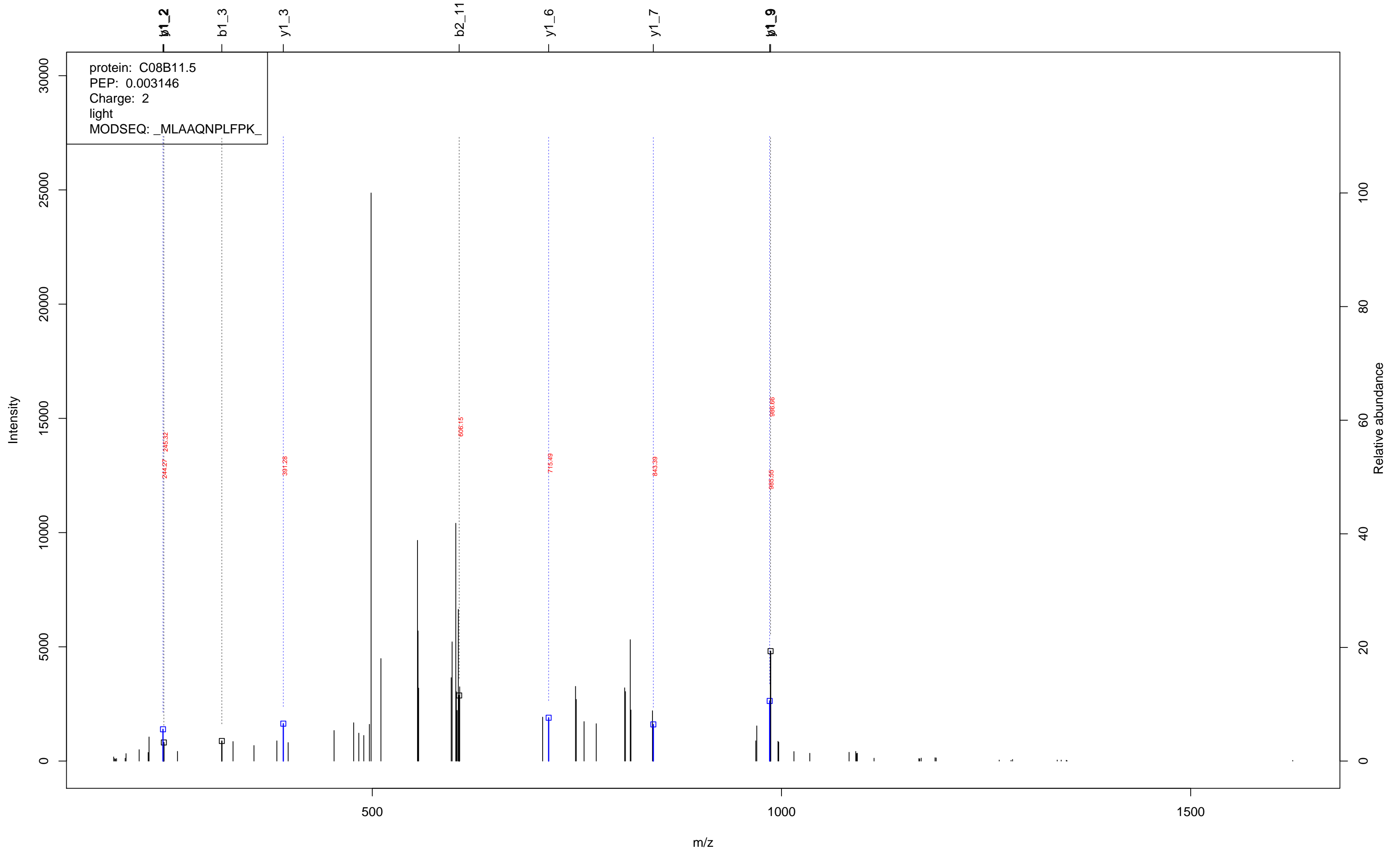


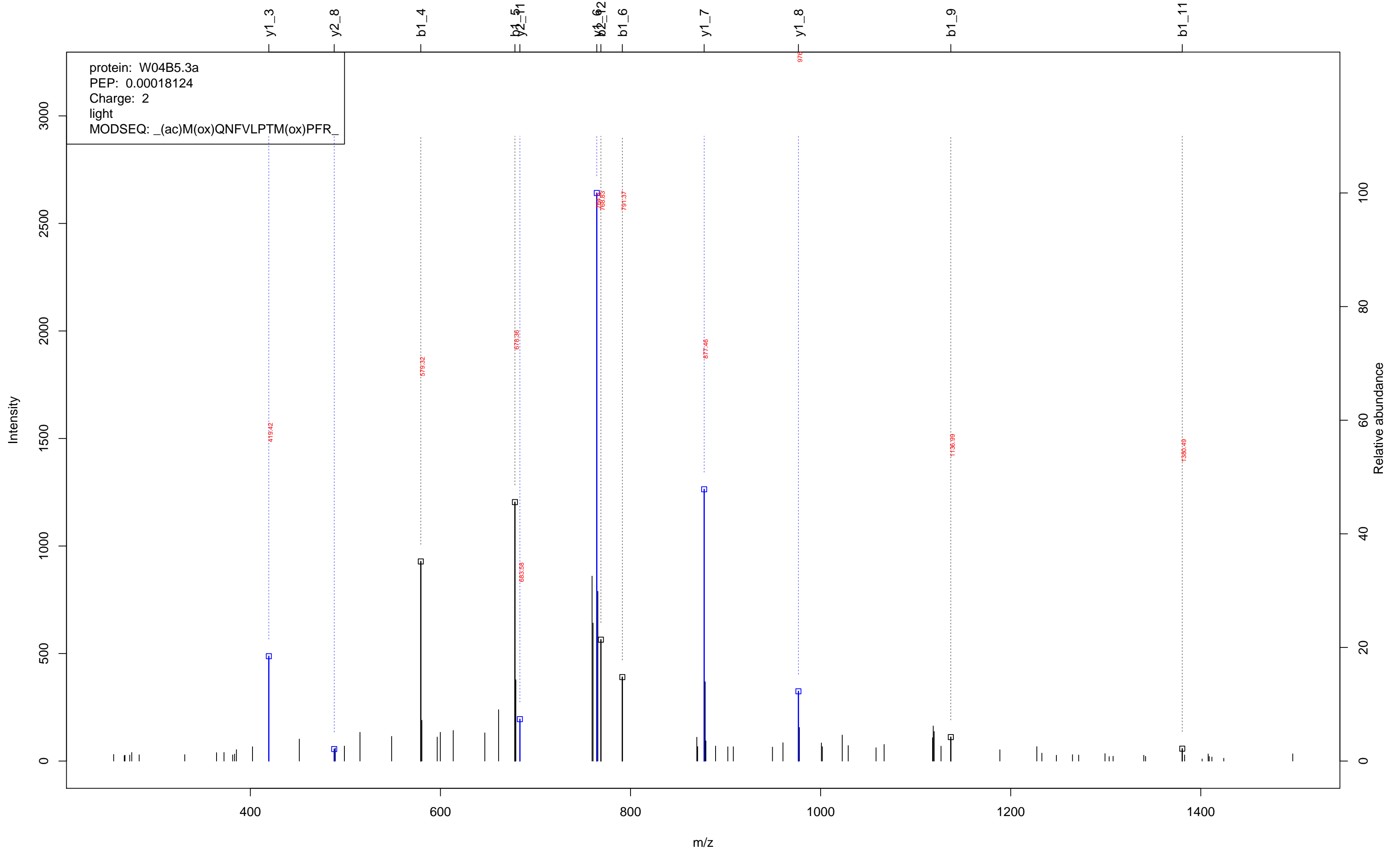


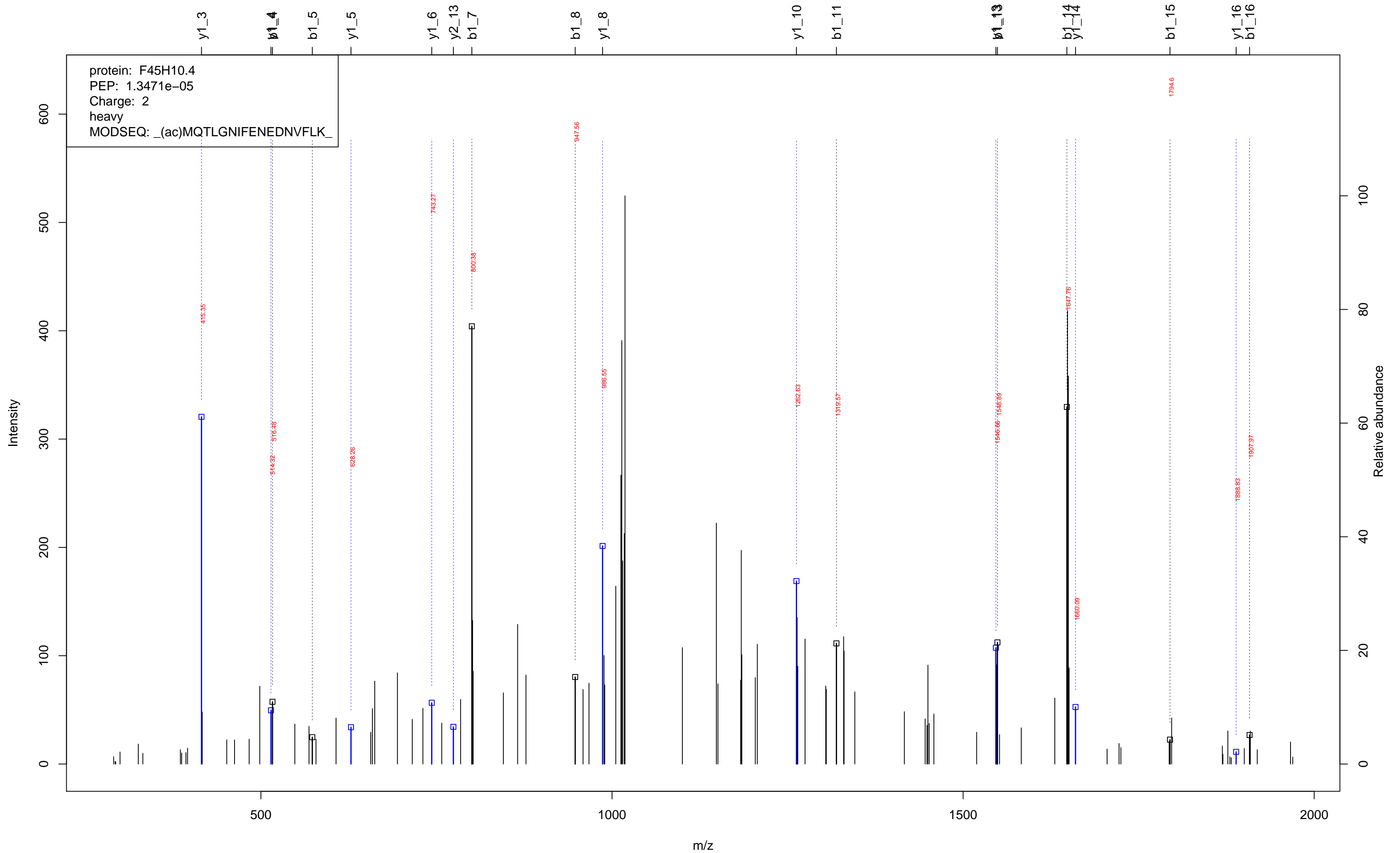


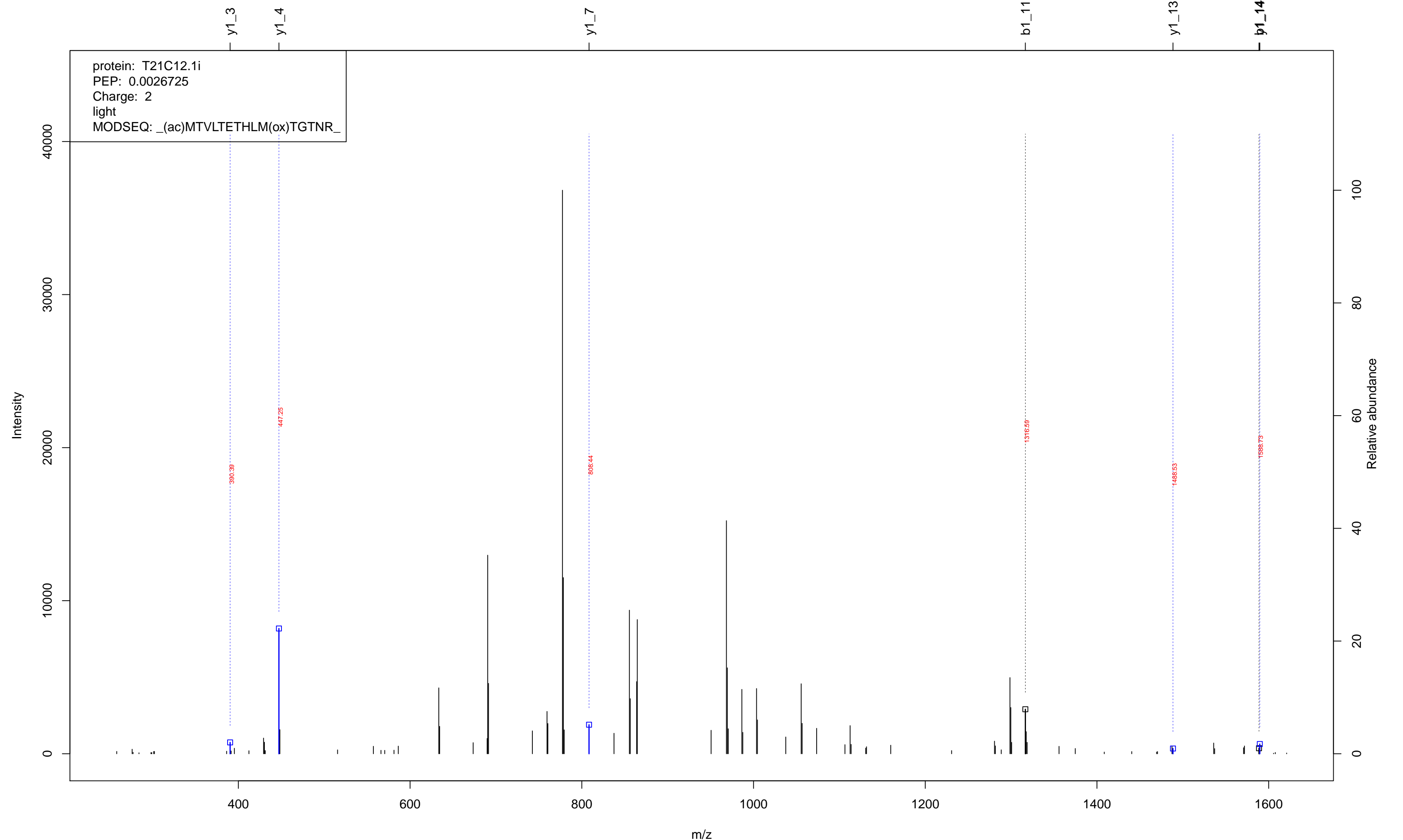


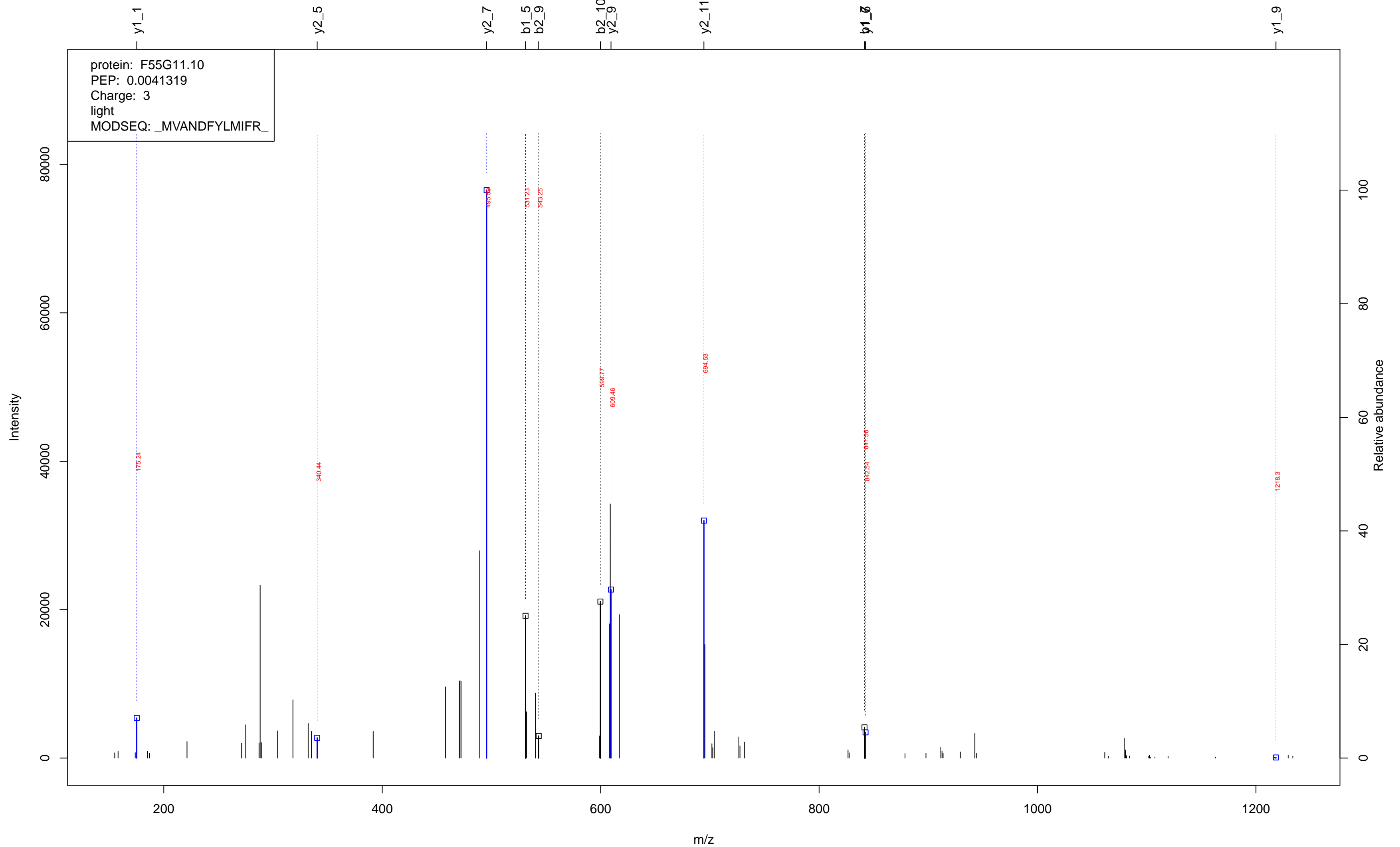


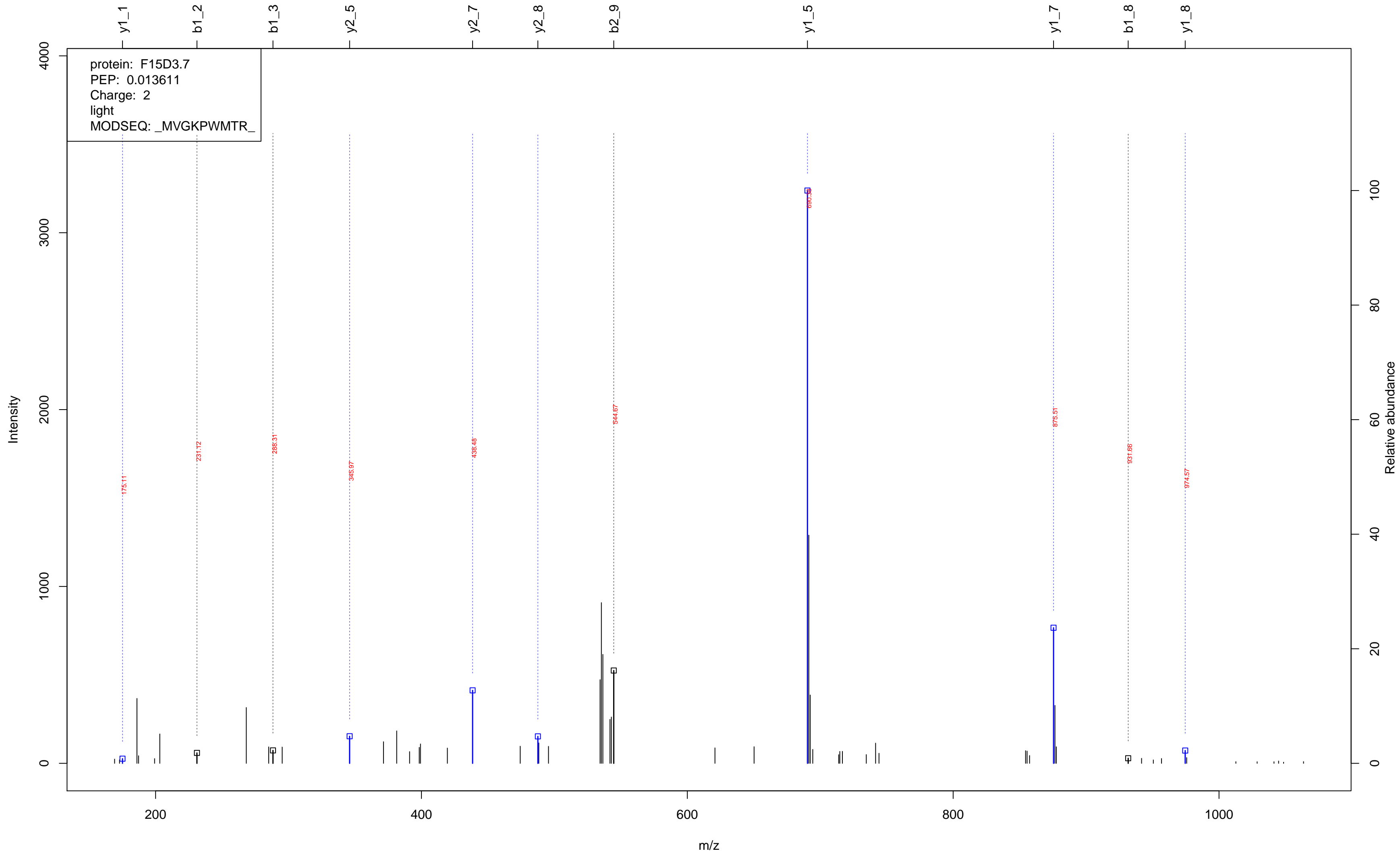


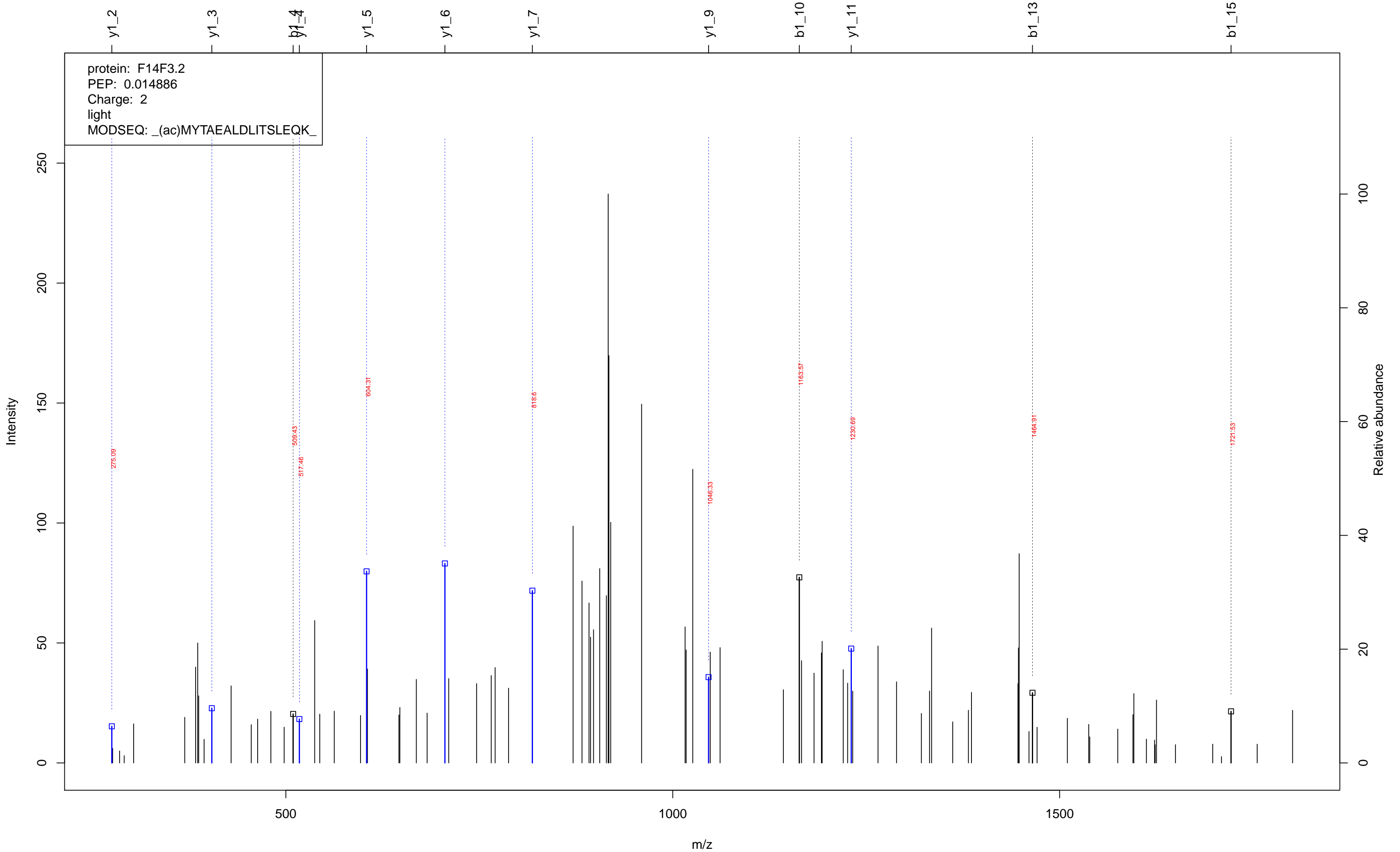


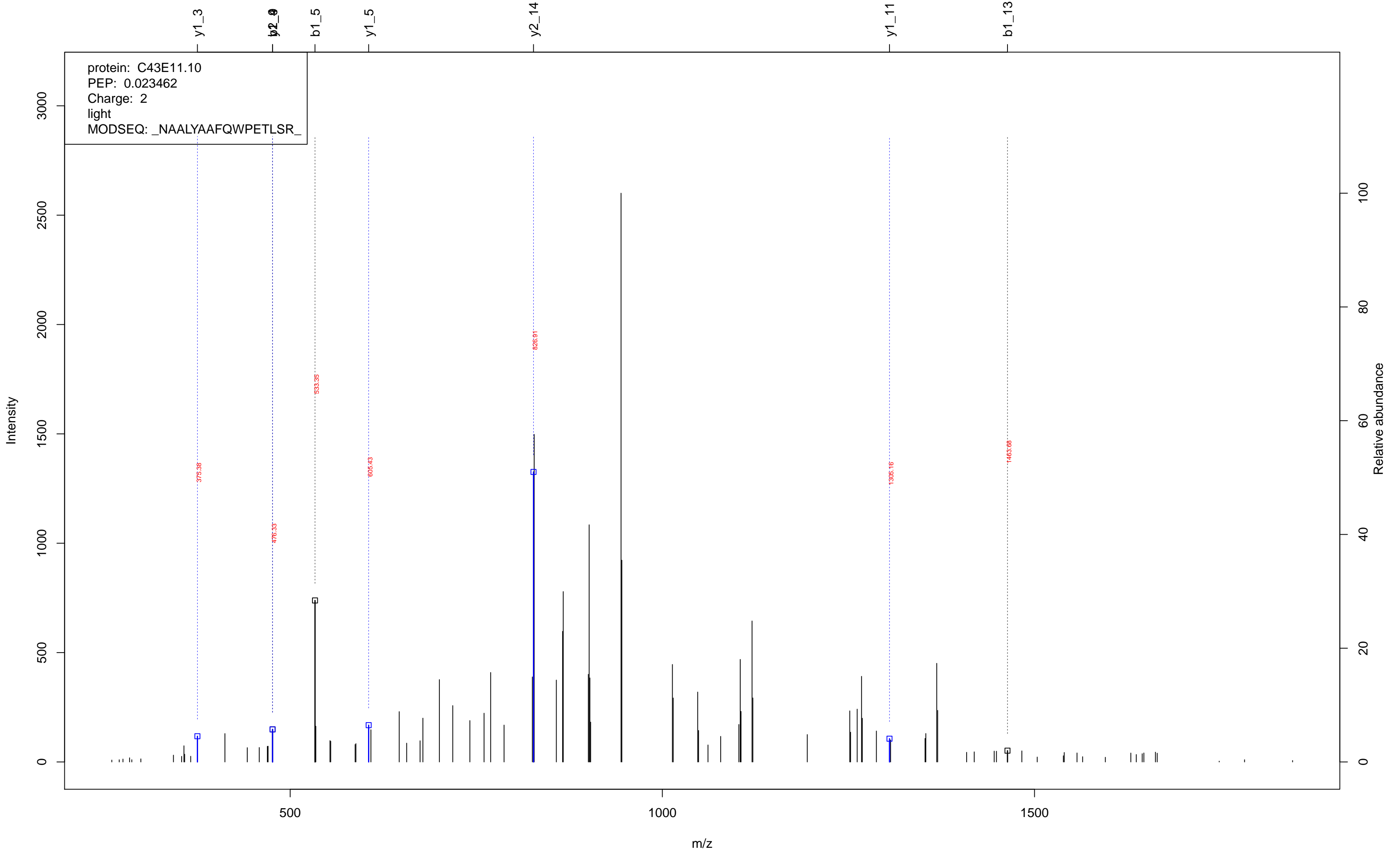


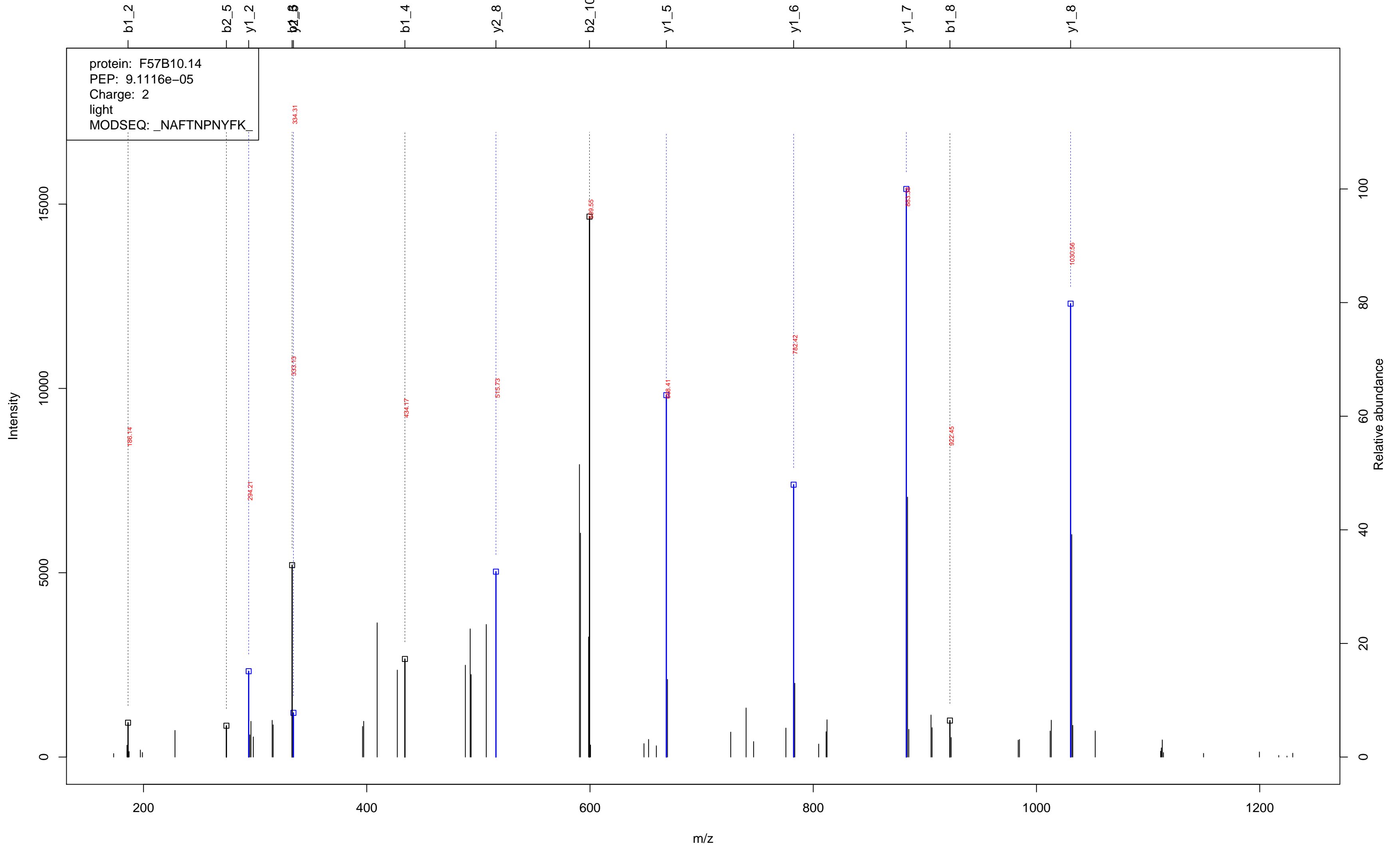


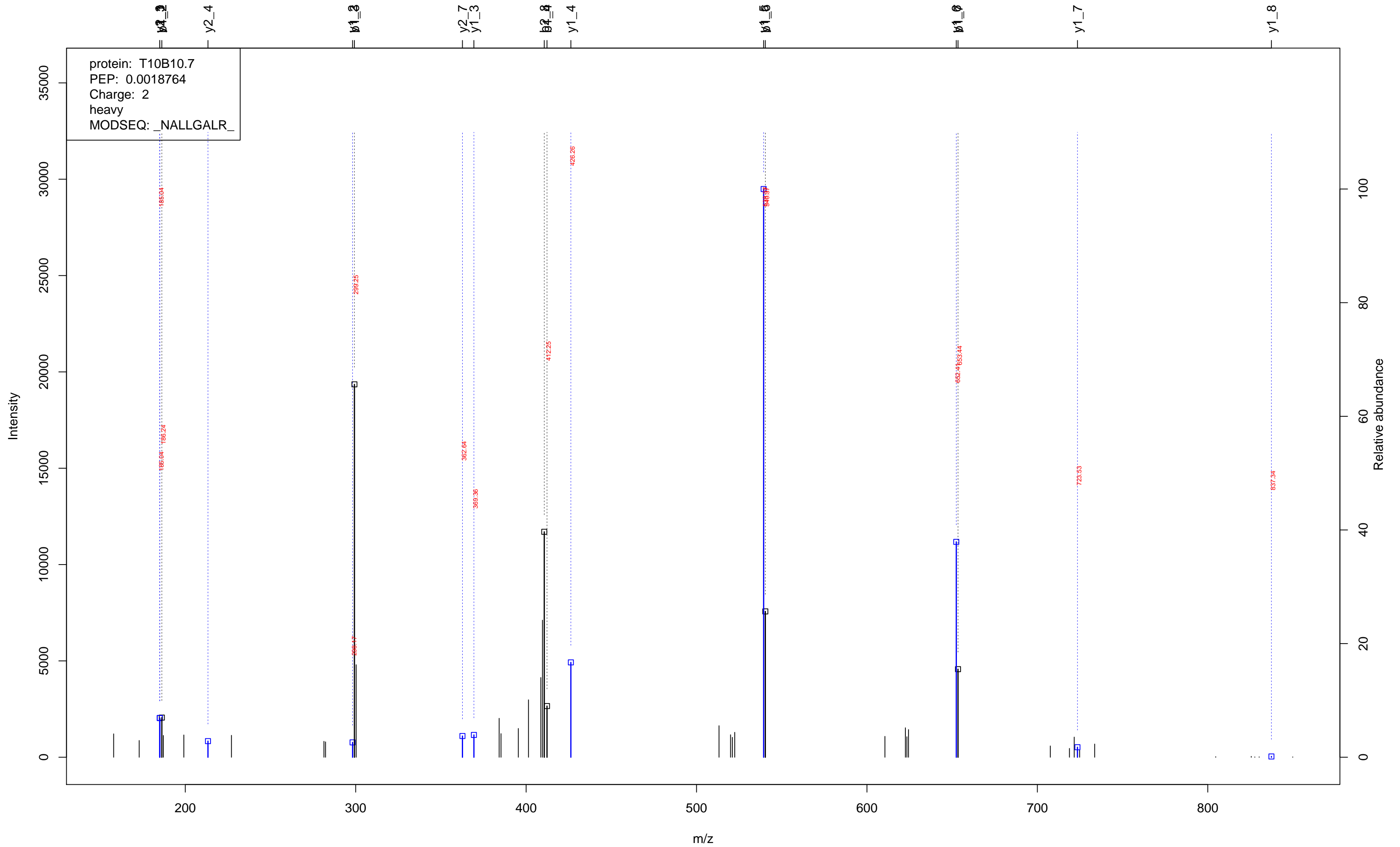


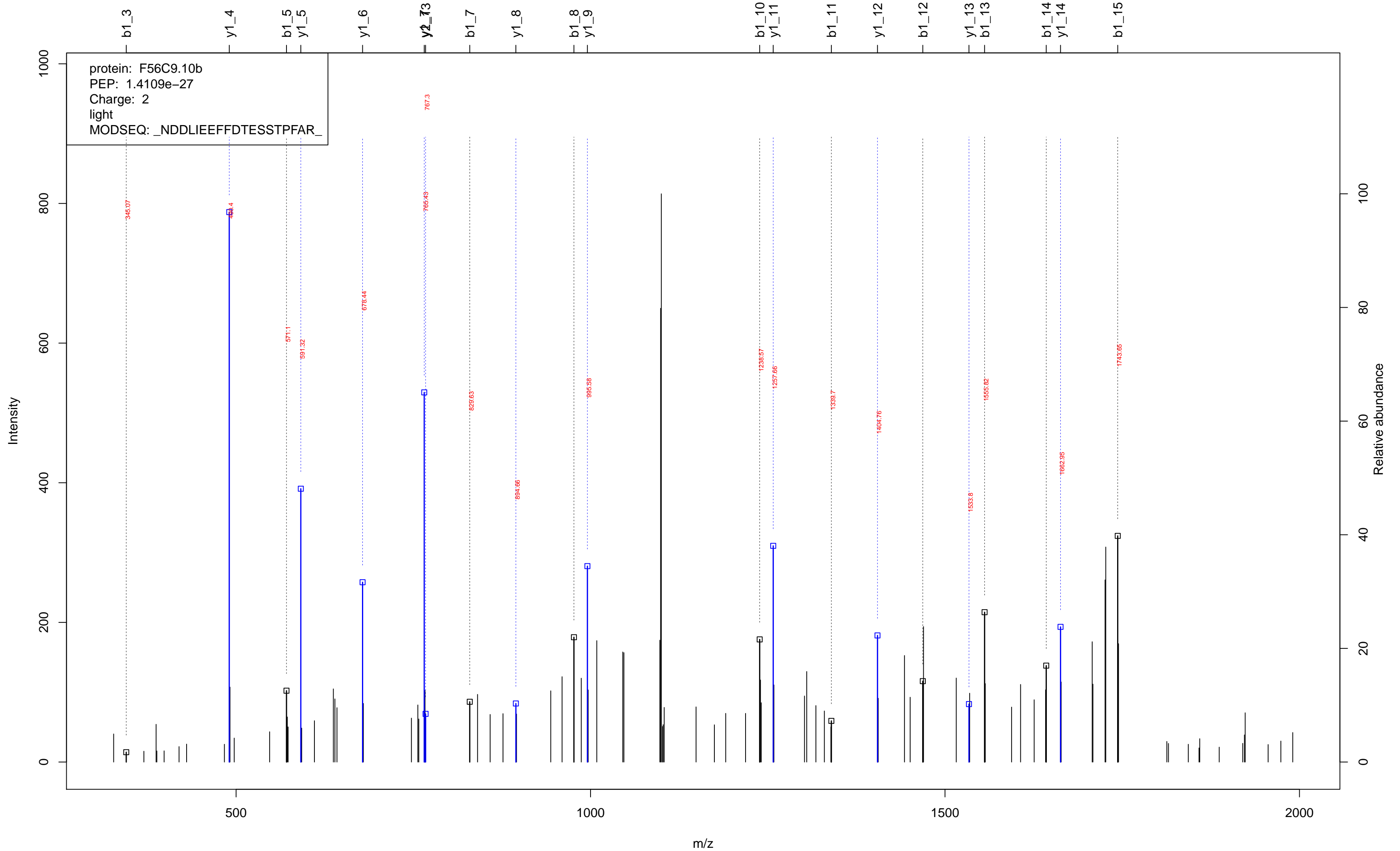


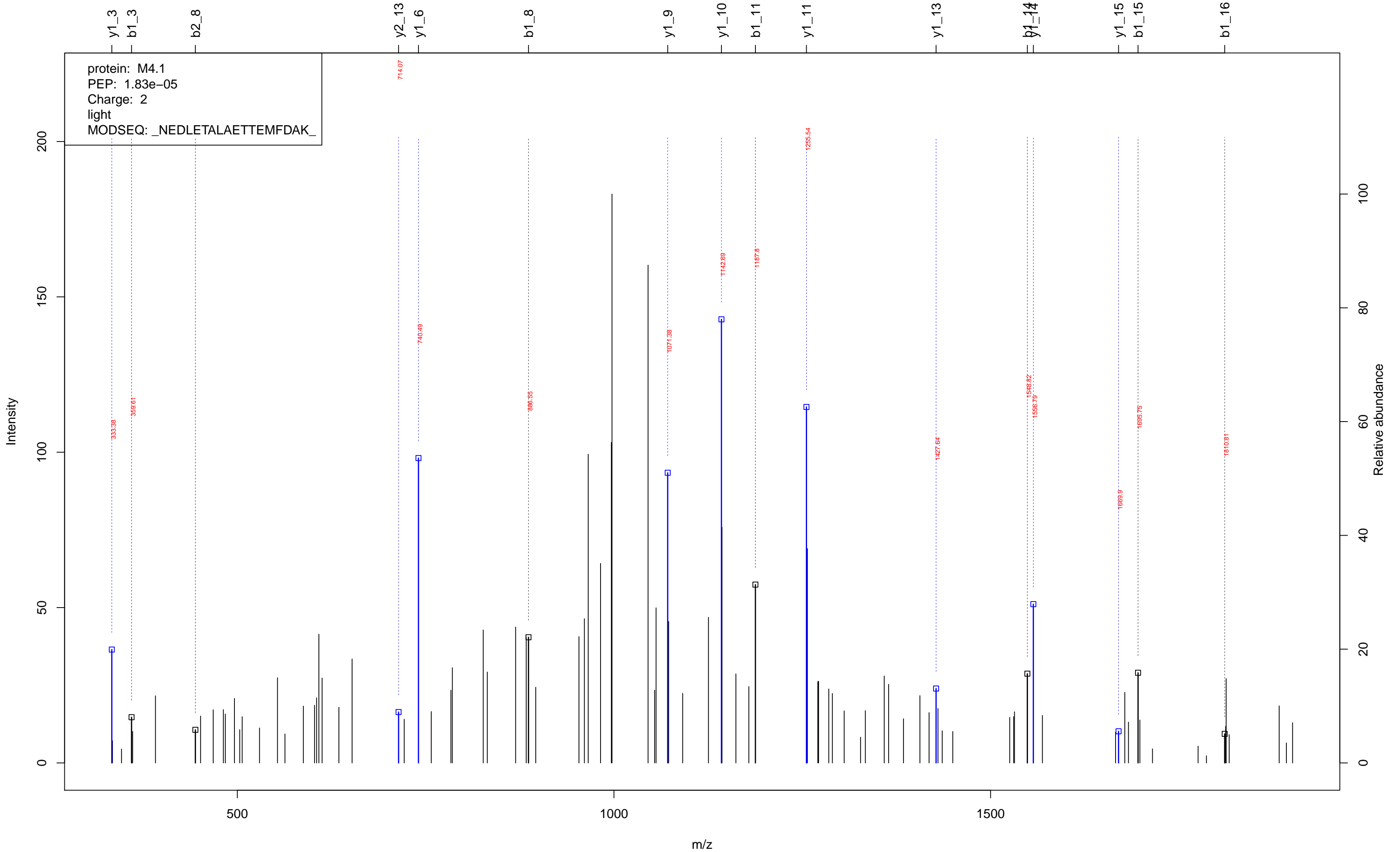


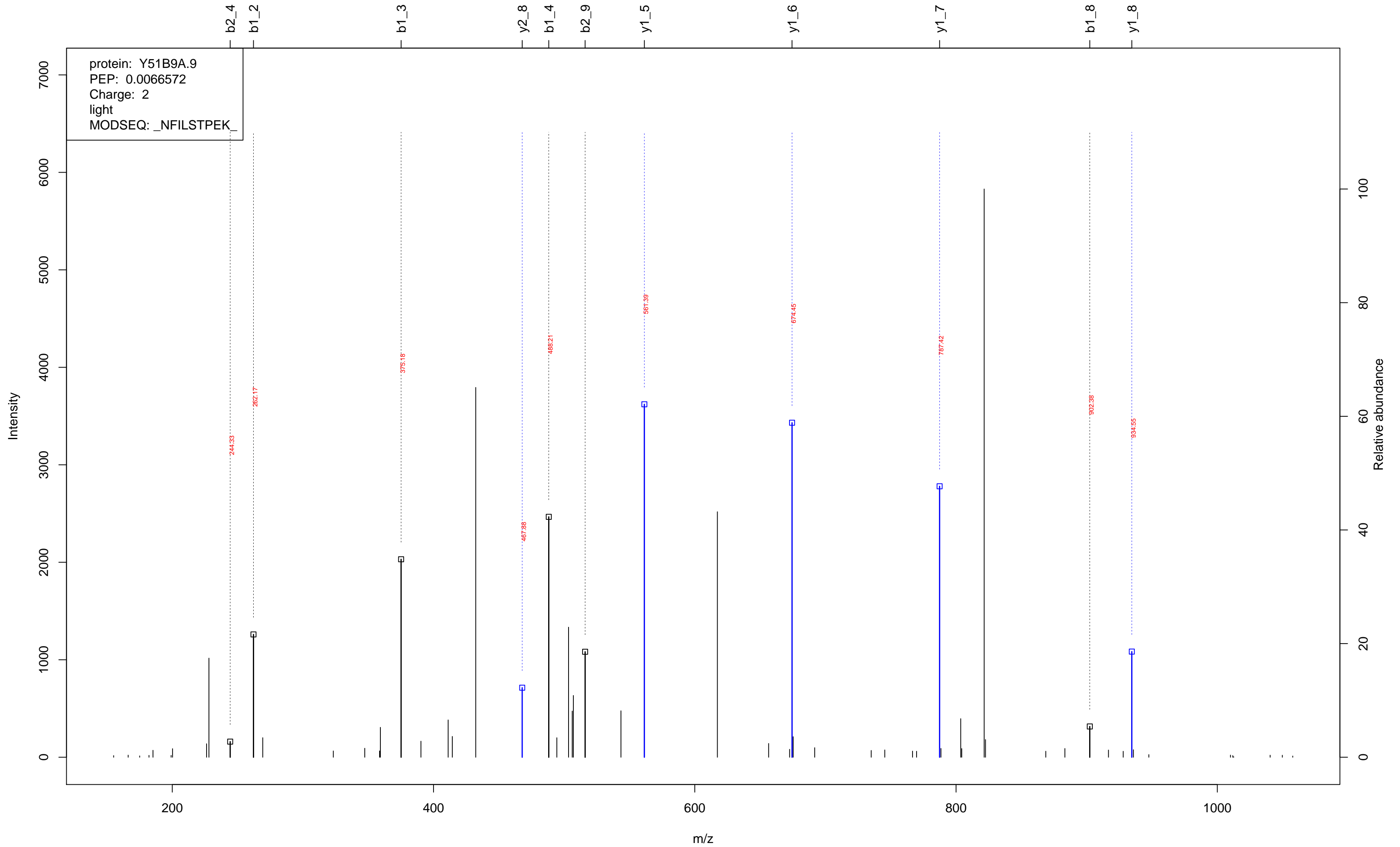












y1_2

y1_4

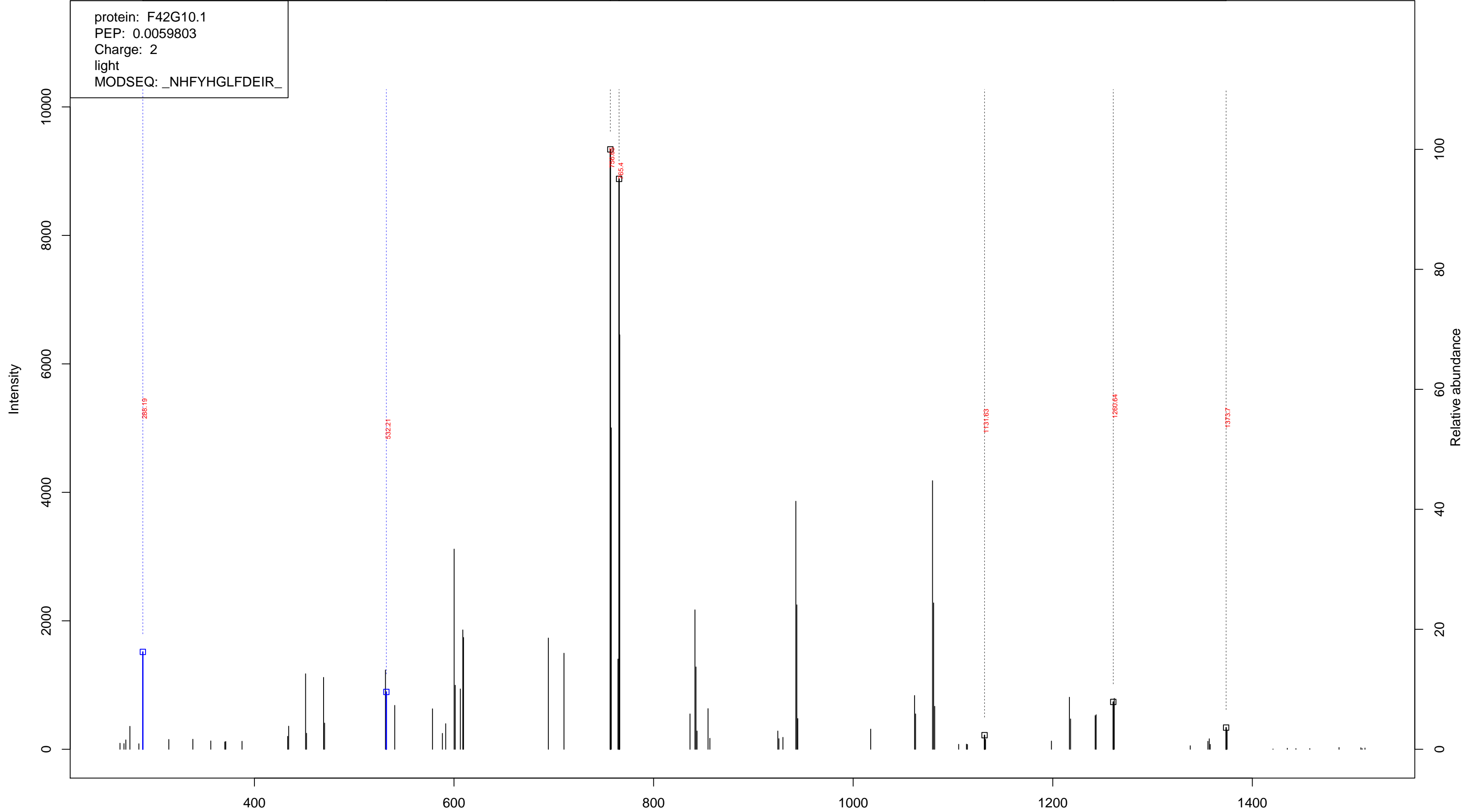
b1_6
b2_12

b1_9

b1_10

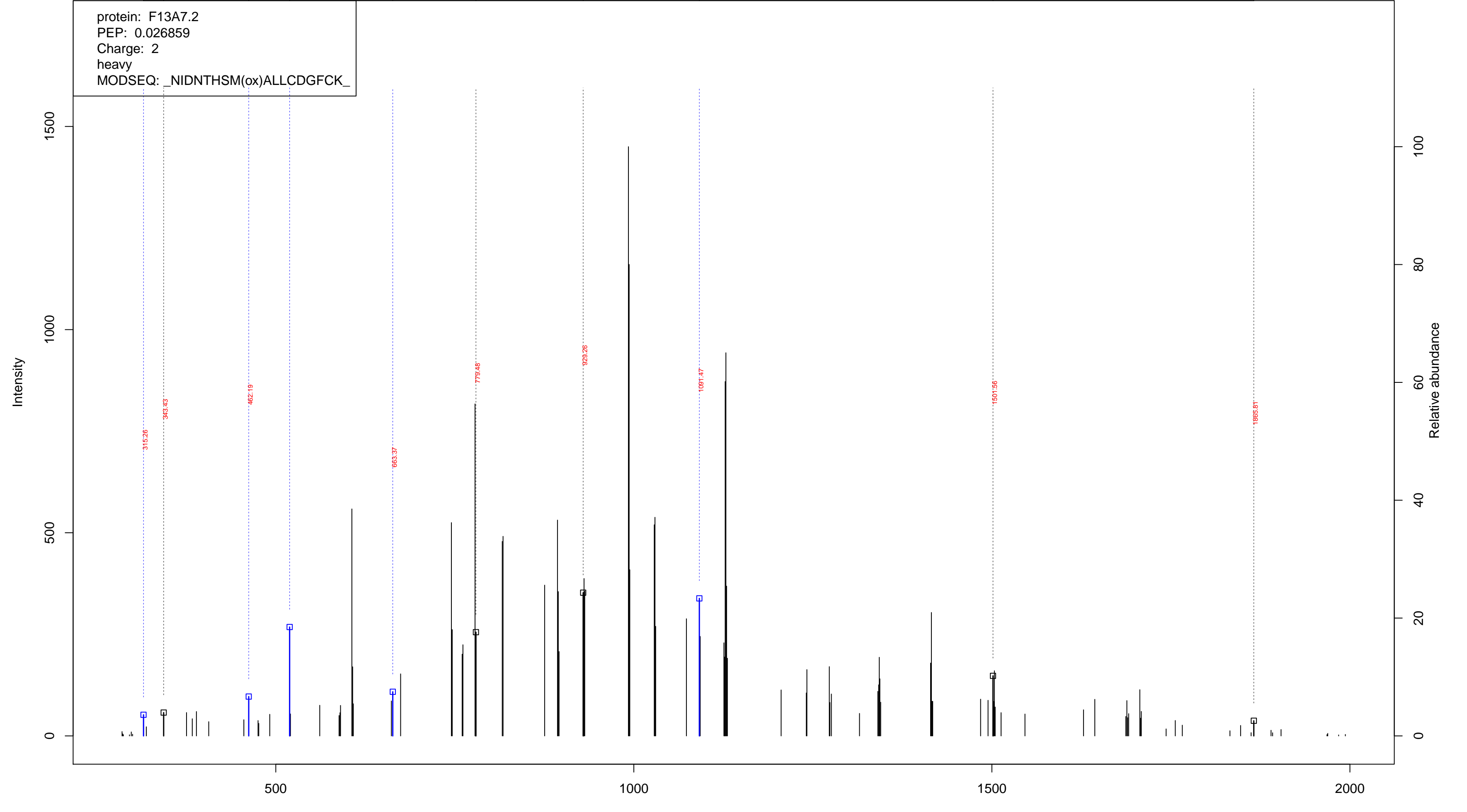
b1_11

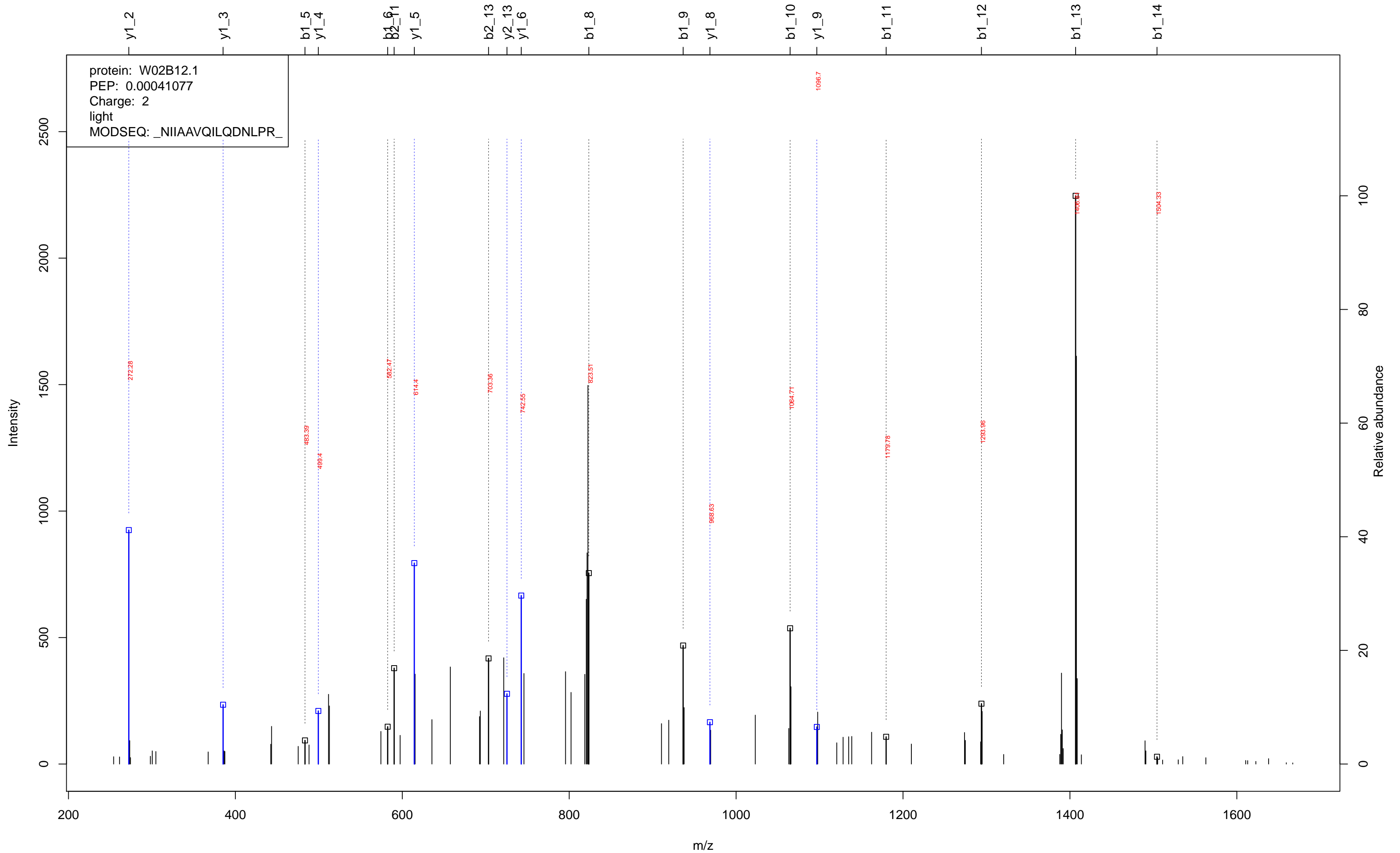
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PEP: 0.0059803
Charge: 2
light
MODSEQ: _NHFYHGLFDEIR_



y1_2 b1_3 y1_3 y1_4 y2_11 b2_14 b1_8 y1_9 b1_13 b1_16

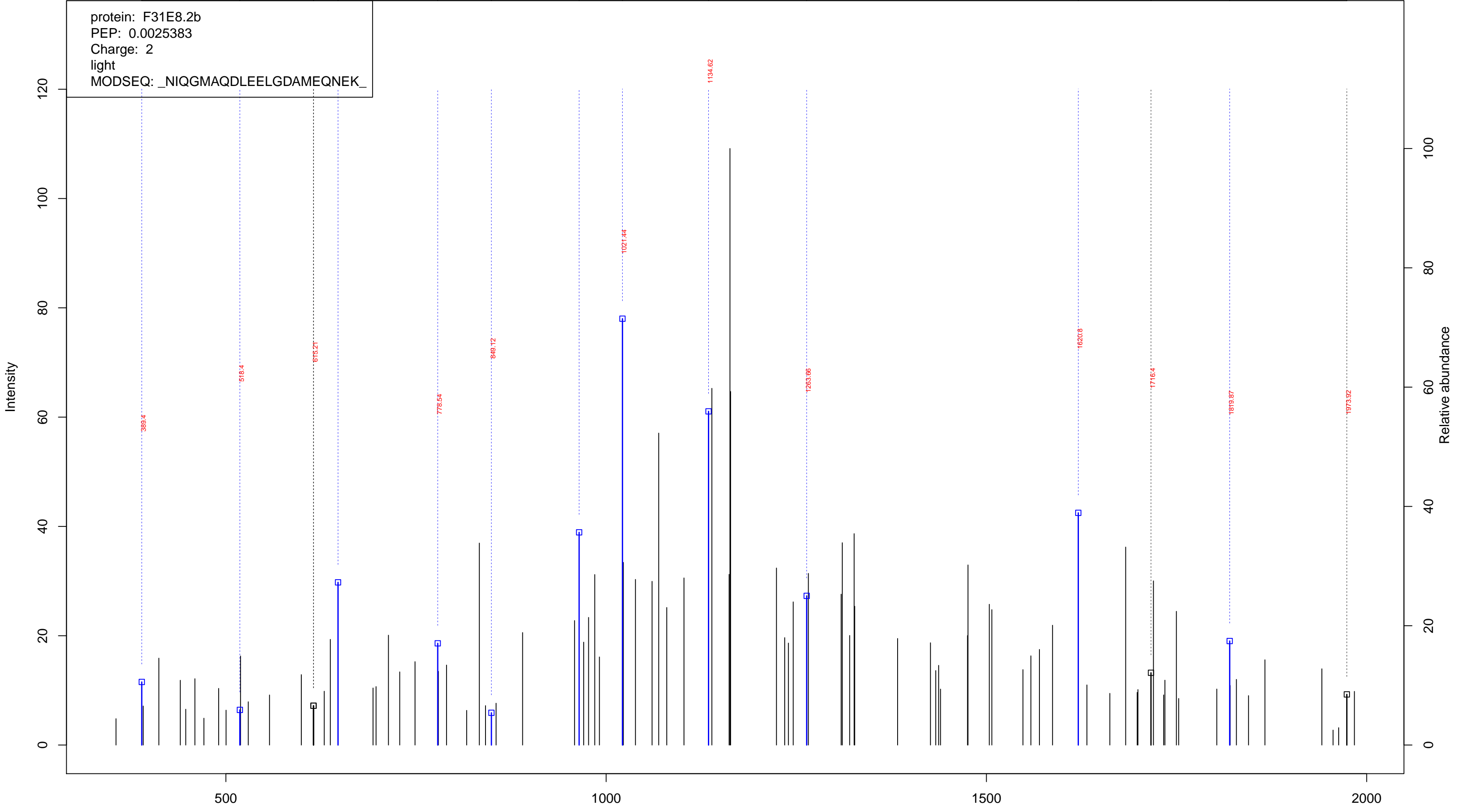
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PEP: 0.026859
Charge: 2
heavy
MODSEQ: _NIDNTHSM(ox)ALLCDGFCK_

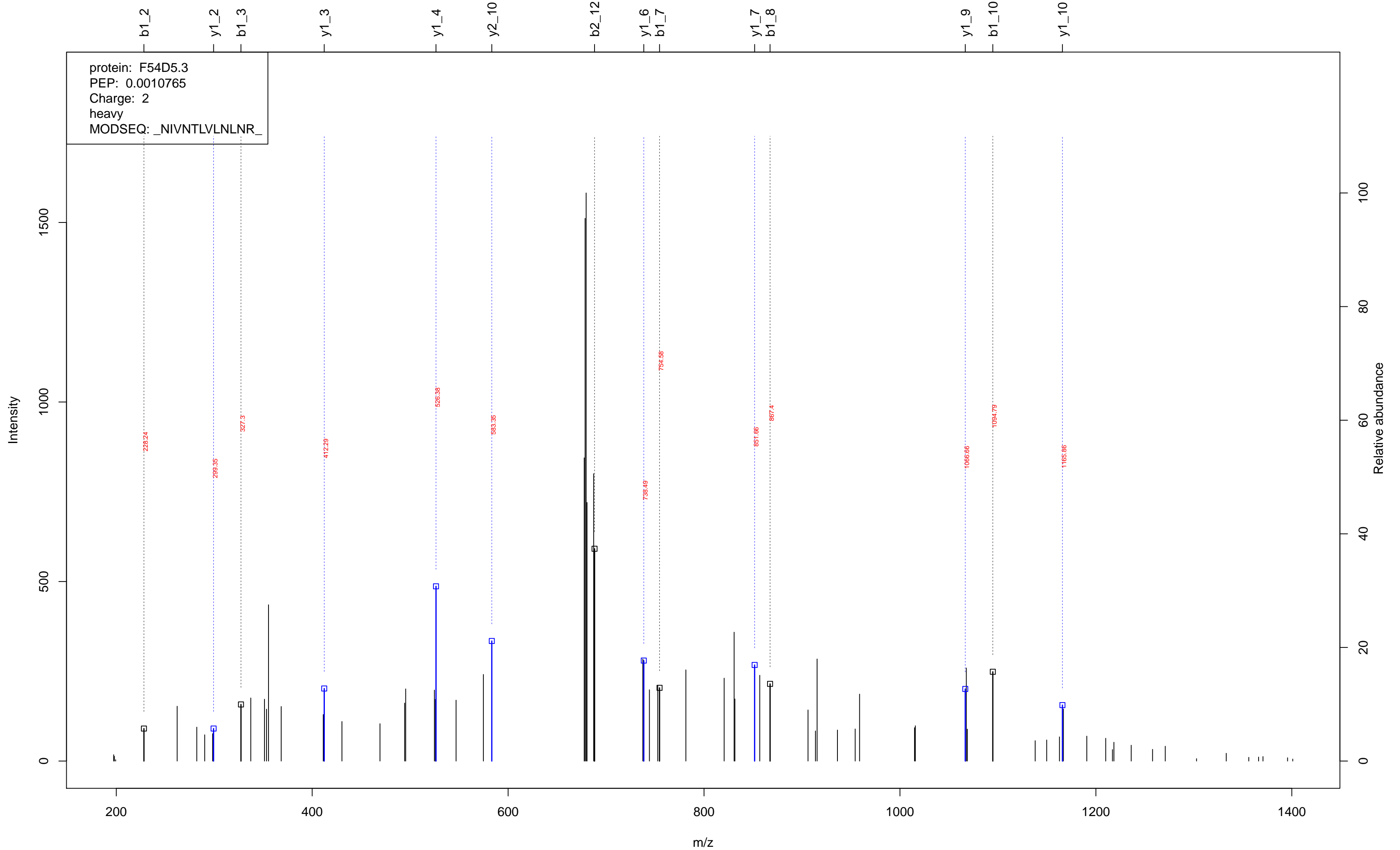


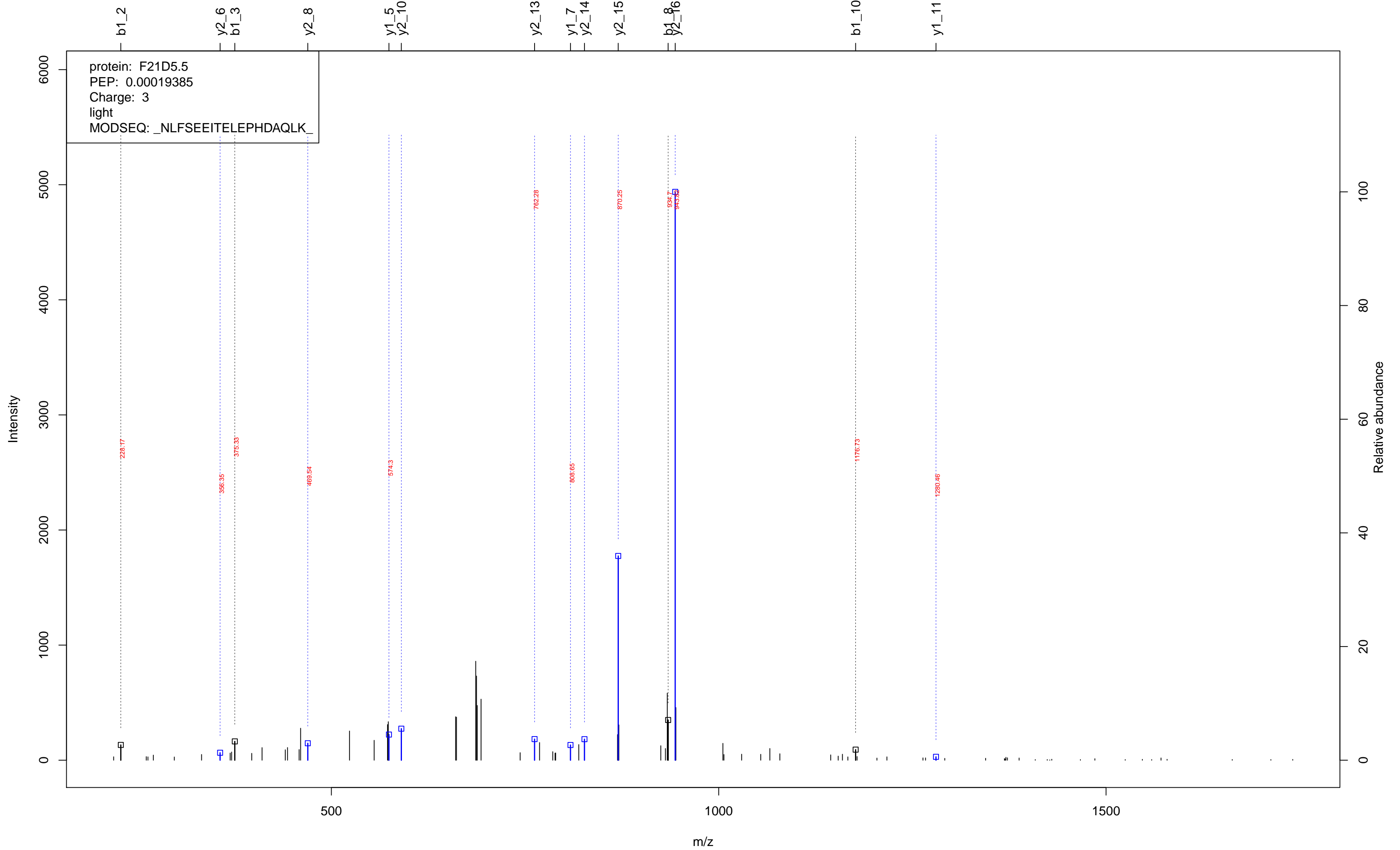


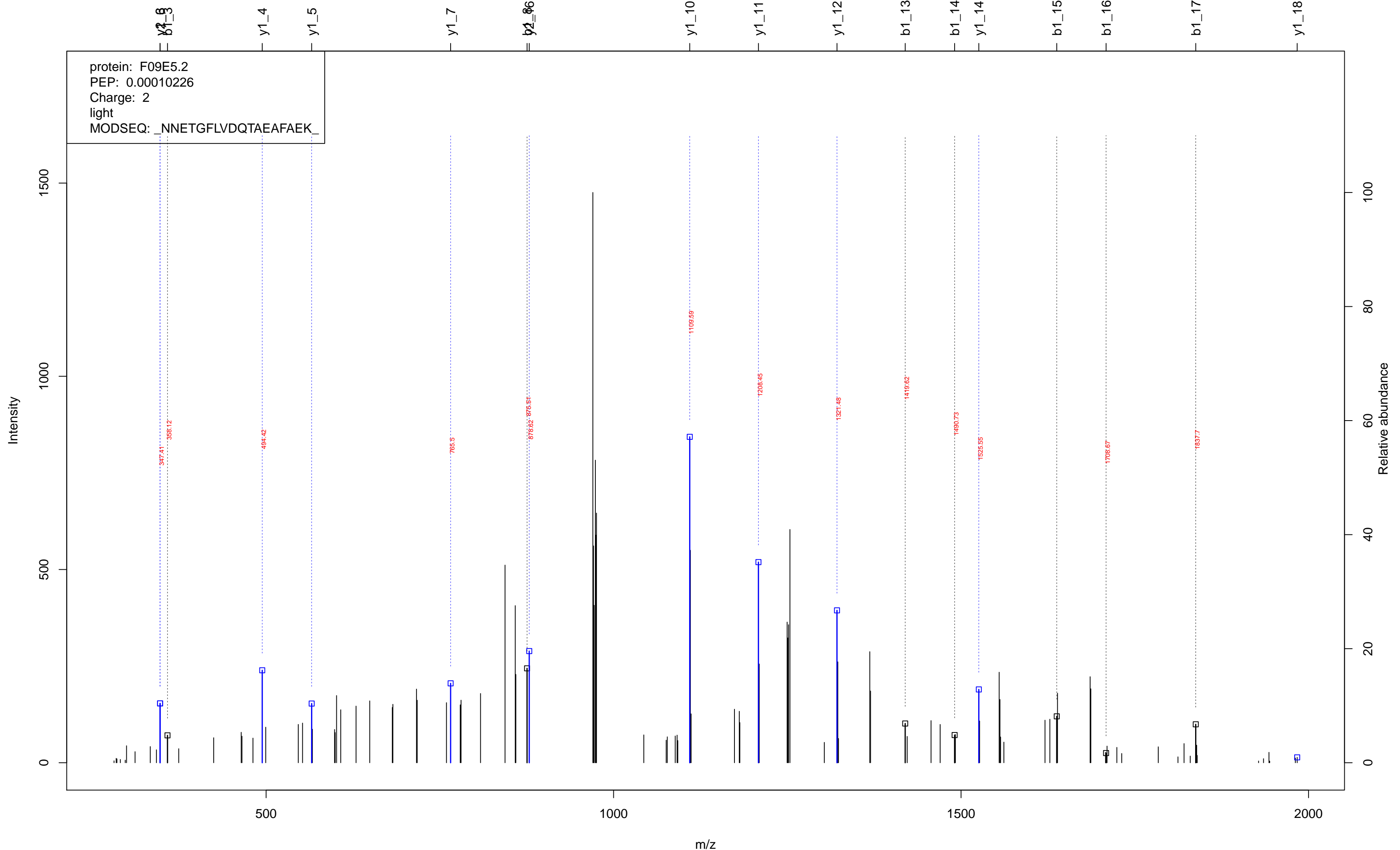
y2_6 y1_4 b2_61 y1_5 y1_6 y1_7 y1_8 y1_9 y1_10 y1_11 y1_14 b1_16 y1_16 b1_18

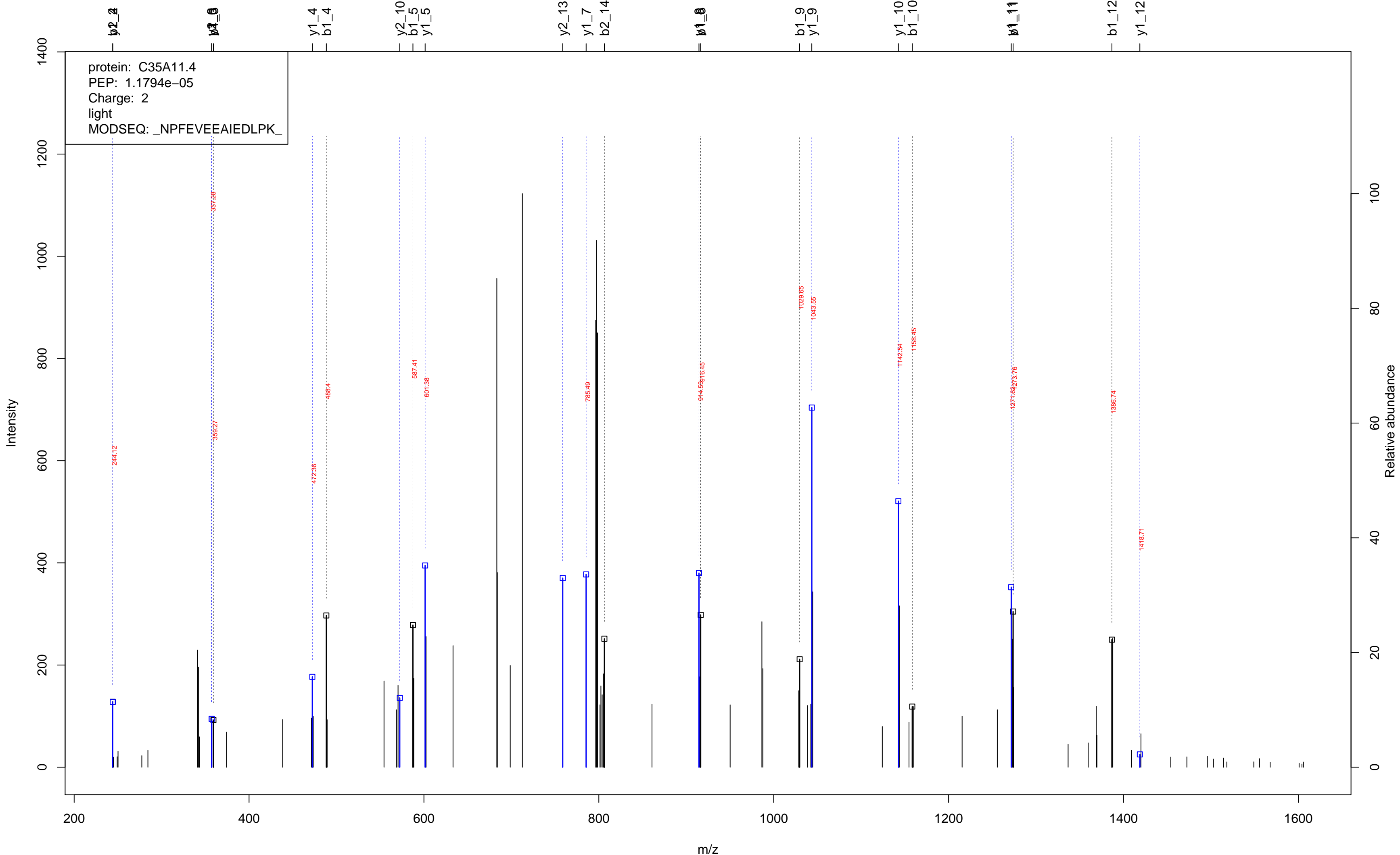
protein: F31E8.2b
PEP: 0.0025383
Charge: 2
light
MODSEQ: _NIQGMAQDLEELGDAMEQNEK_

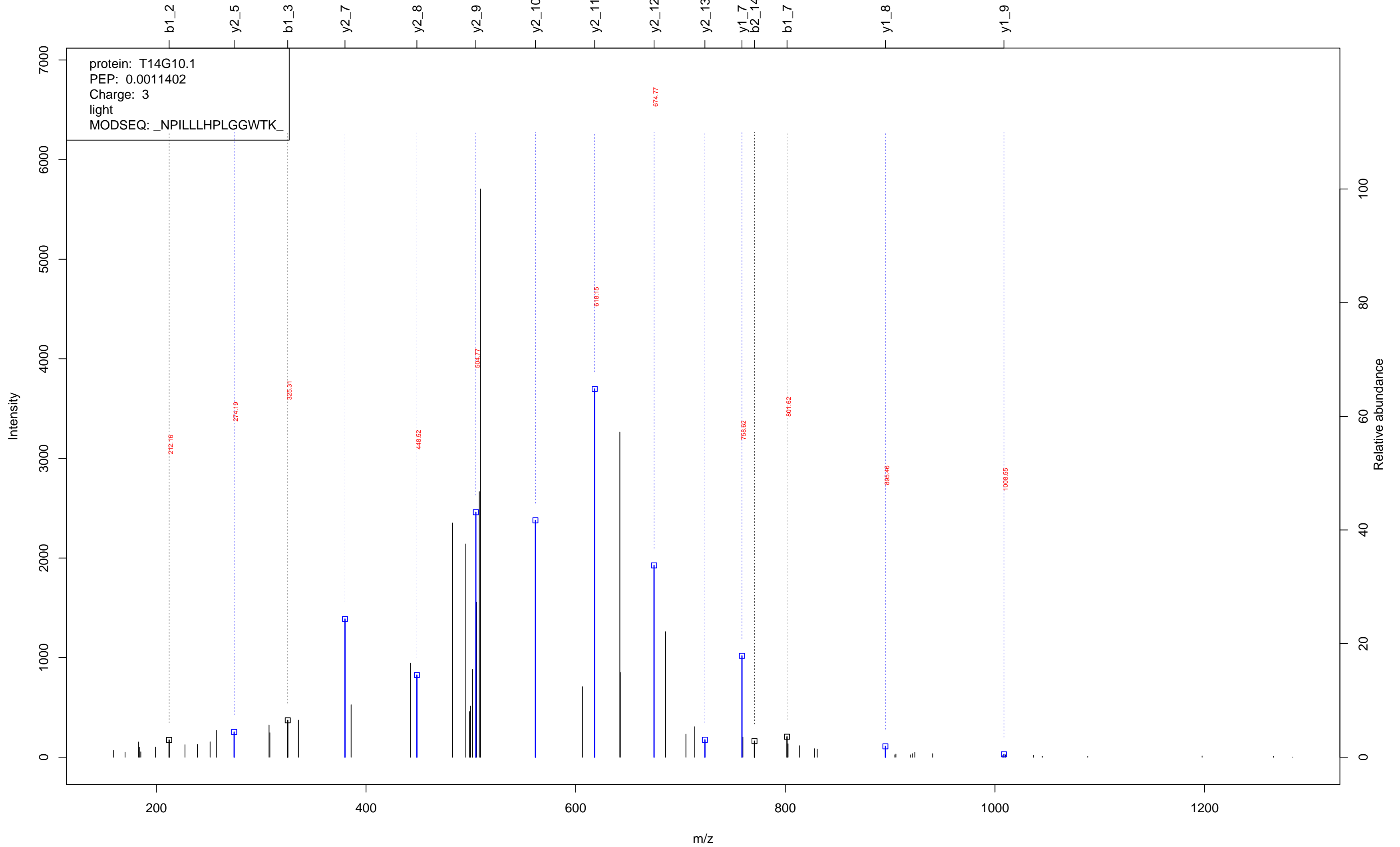


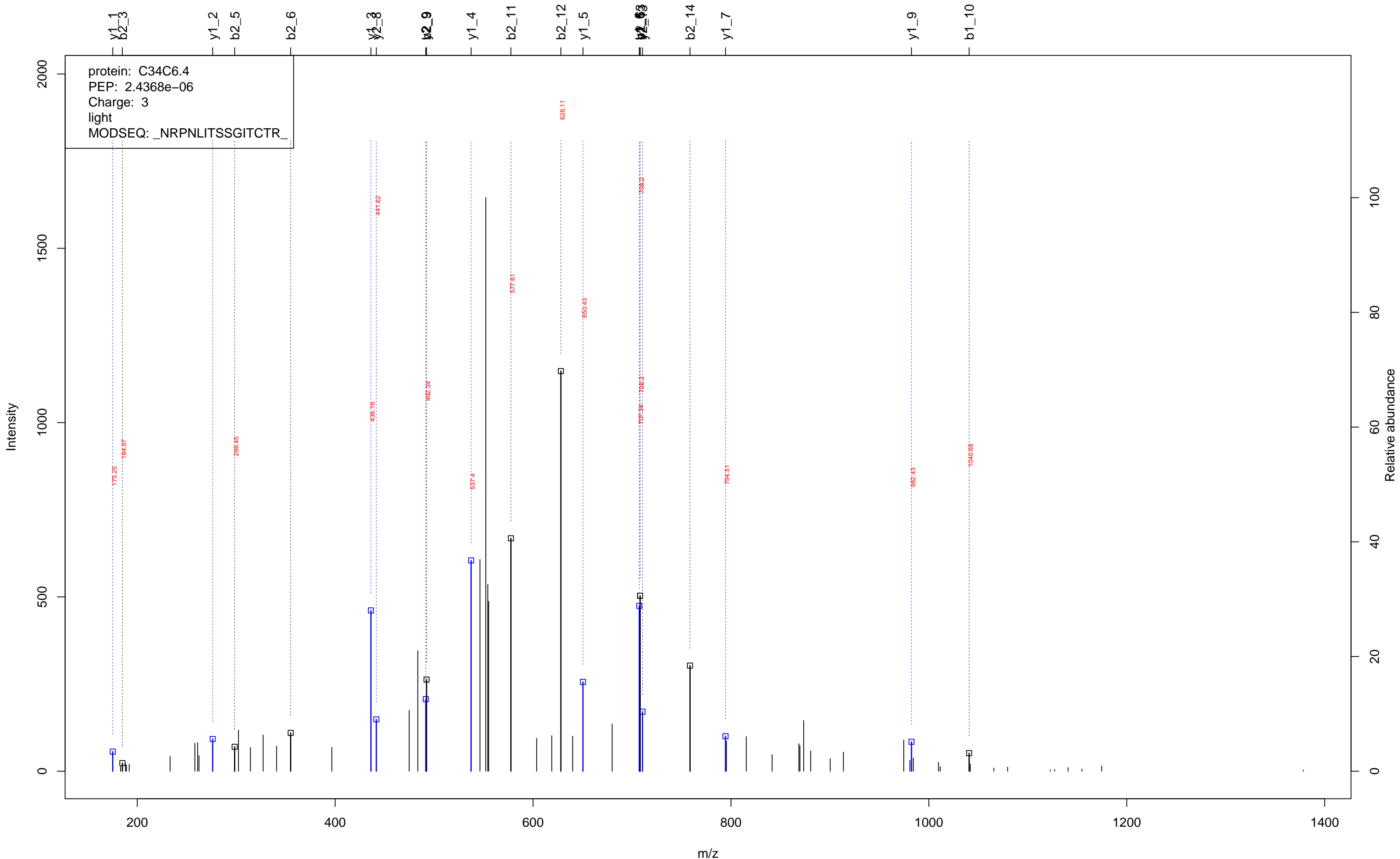








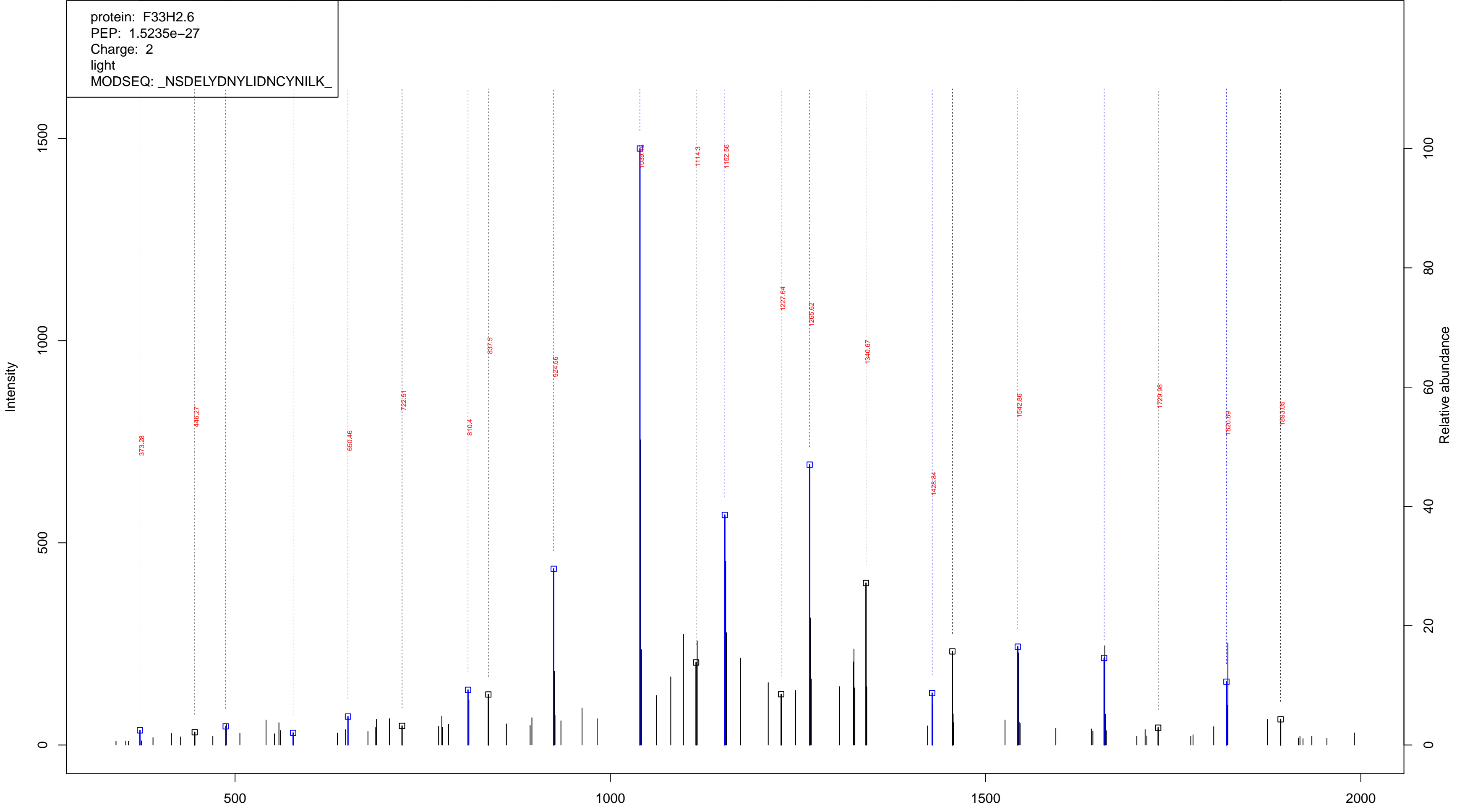


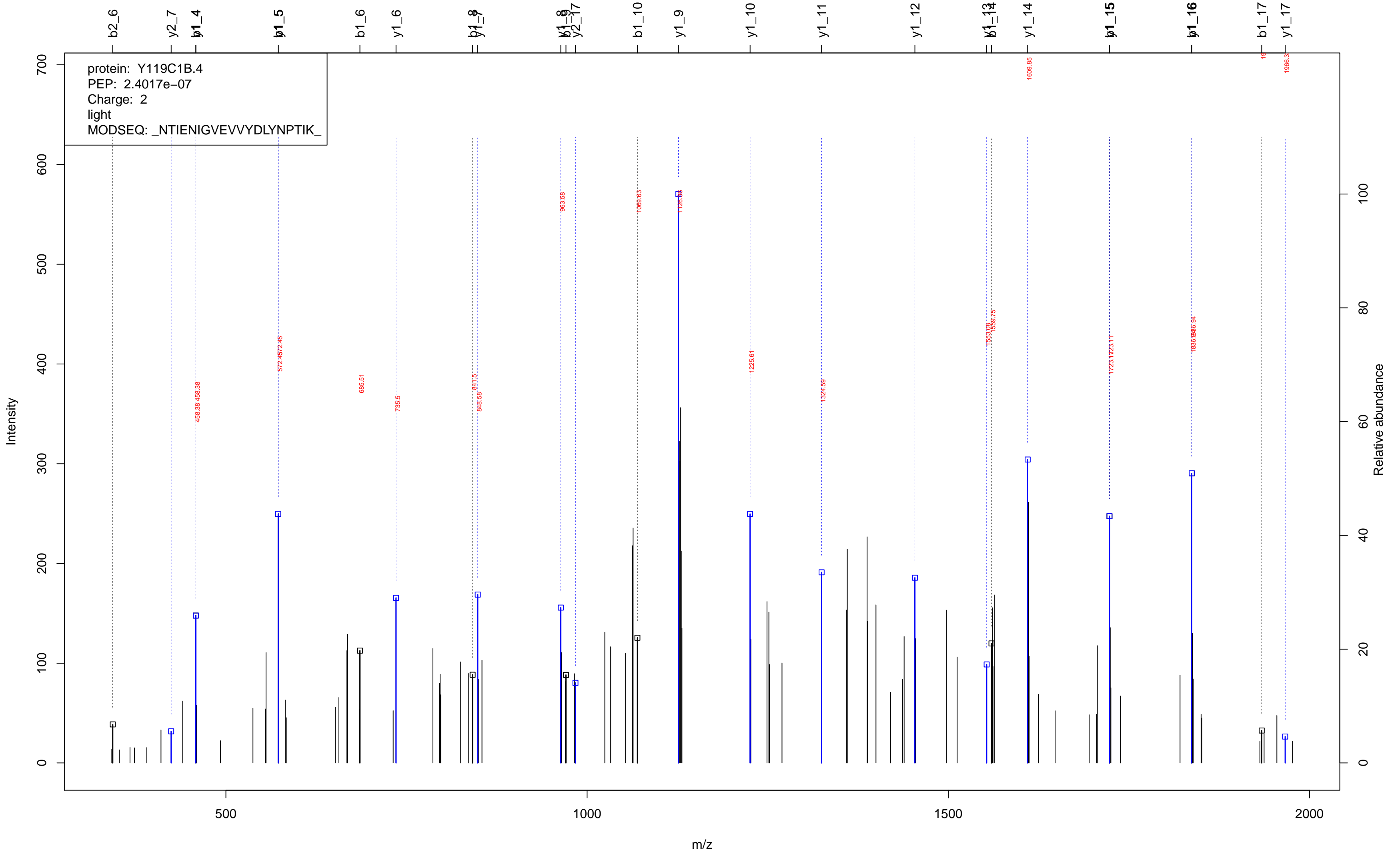


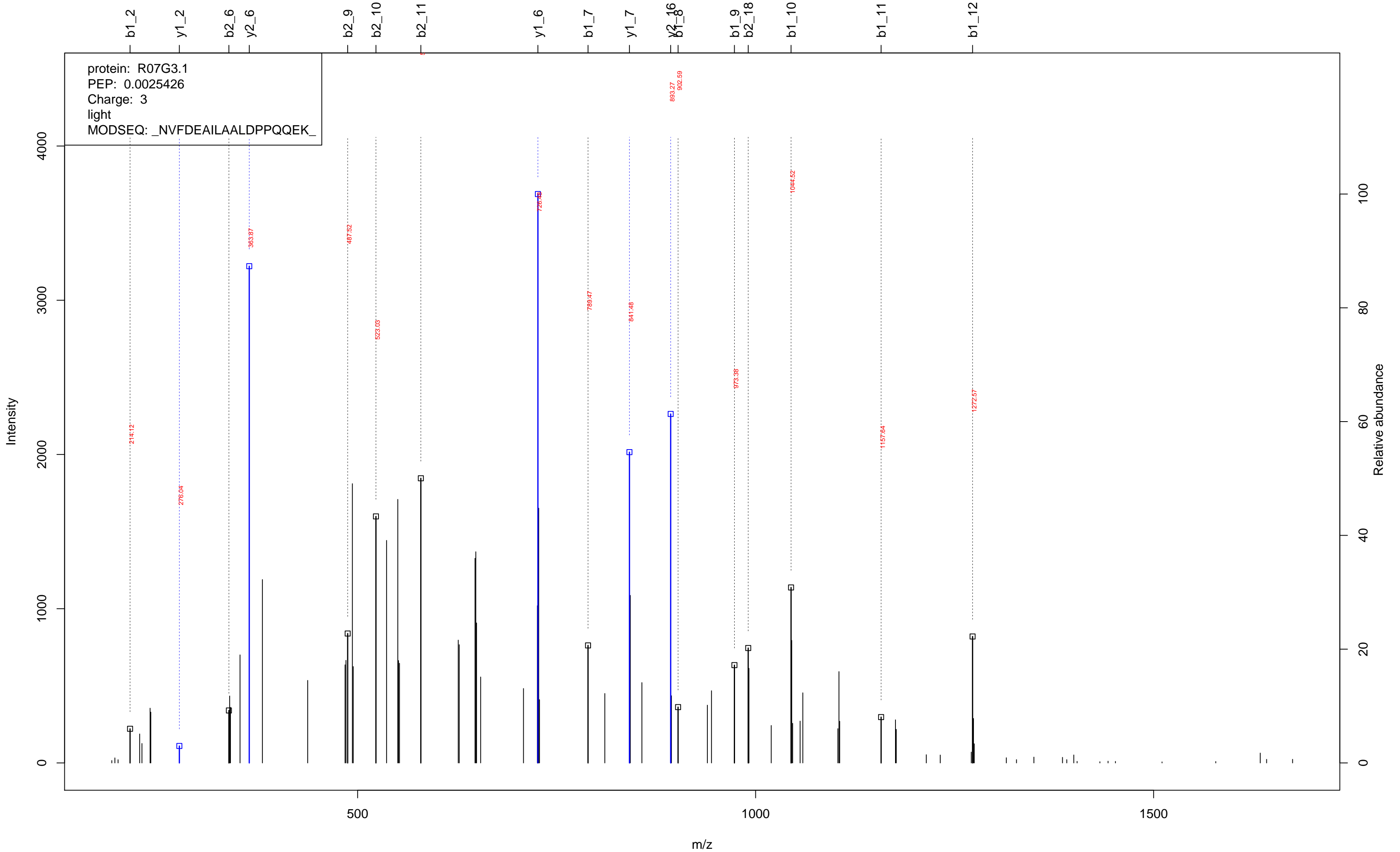
240

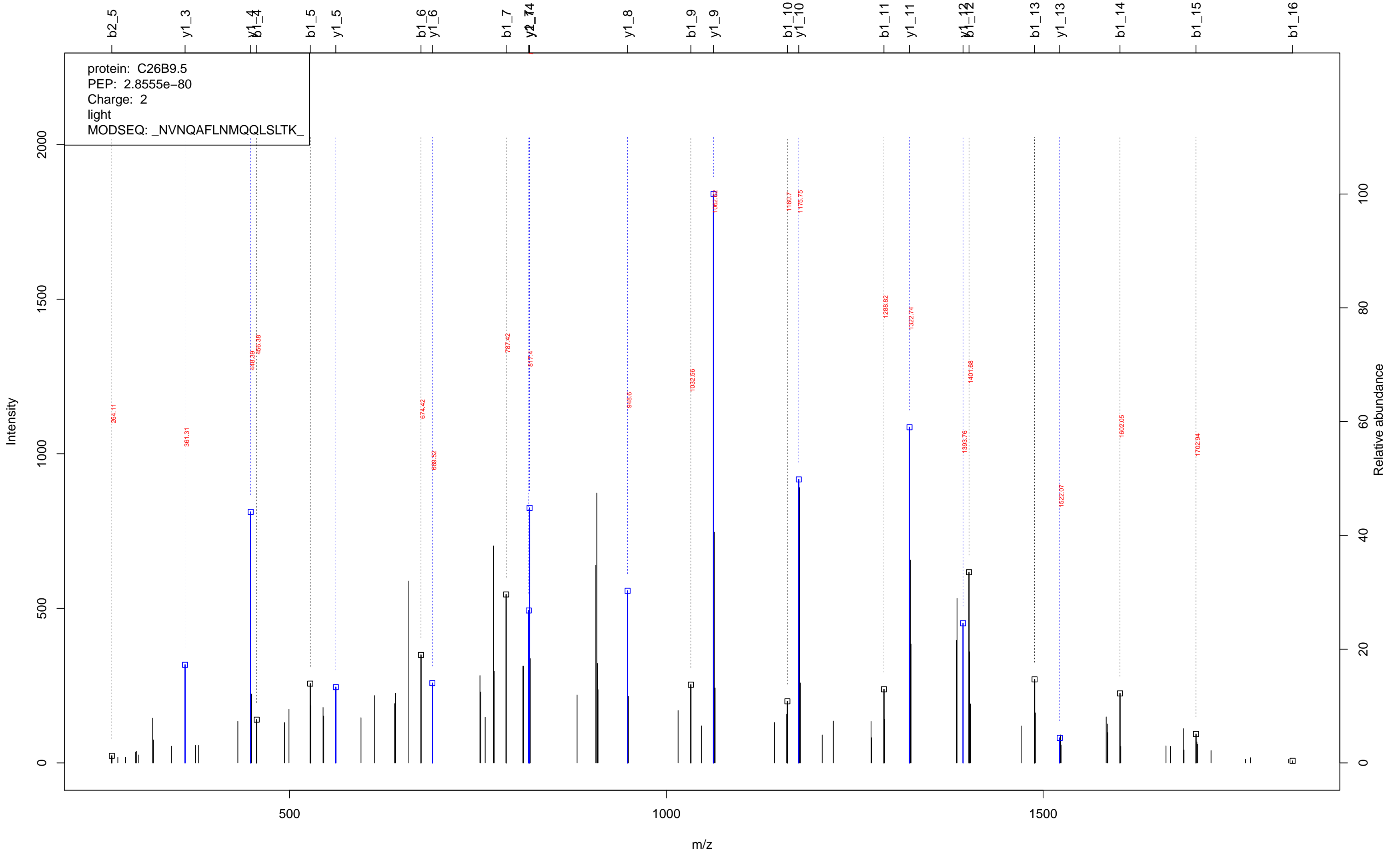
y1_3 b1_4 y1_4 y2_9 y1_5 b1_6 y1_6 b1_7 y1_7 y1_8 b1_9 y1_9 b1_10 y1_10 b1_11 y1_11 b1_12 y1_12 y1_13 b1_14 y1_14 b1_15

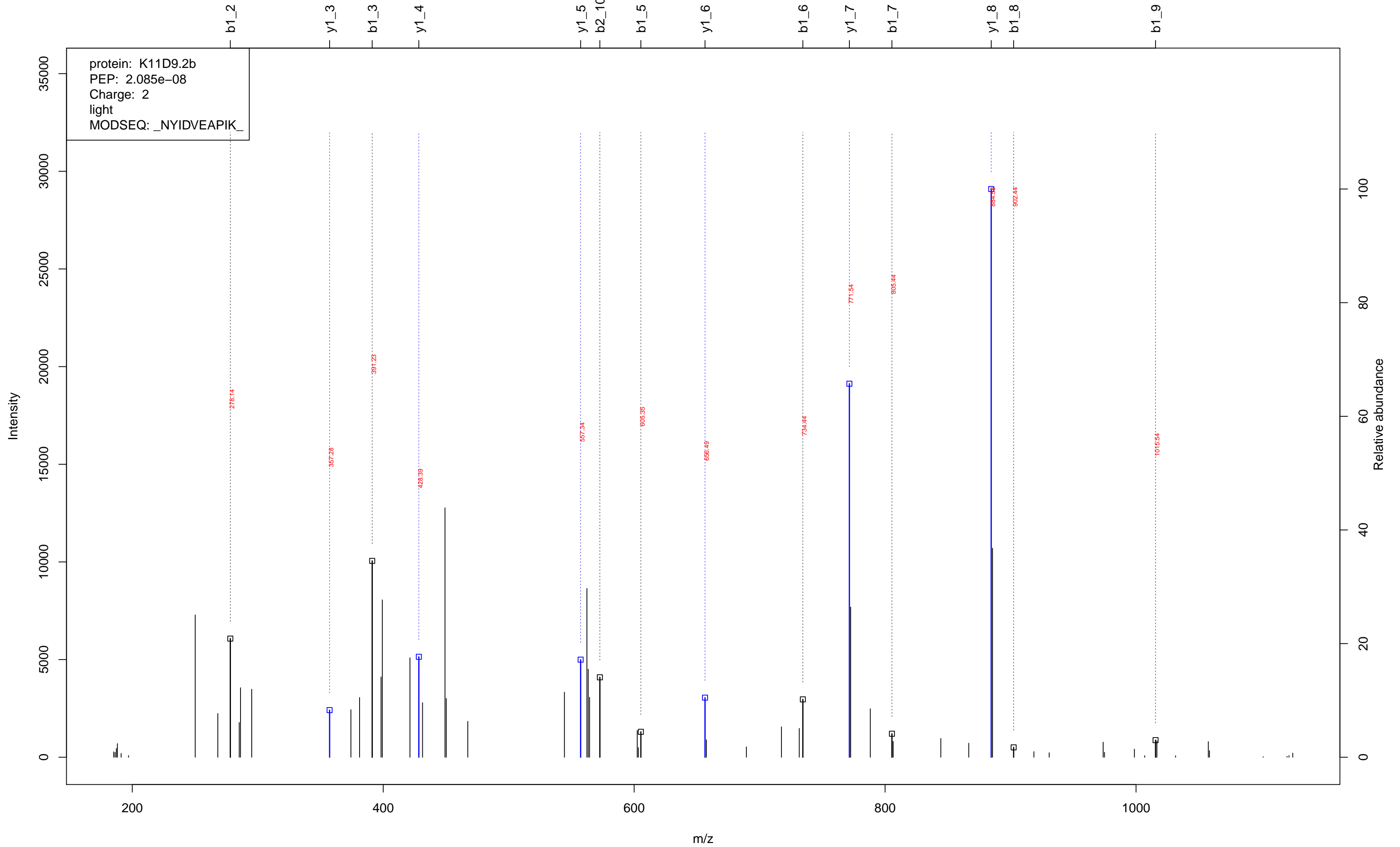
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PEP: 1.5235e-27
Charge: 2
light
MODSEQ: _NSDELYDNYLIDNCYNILK_

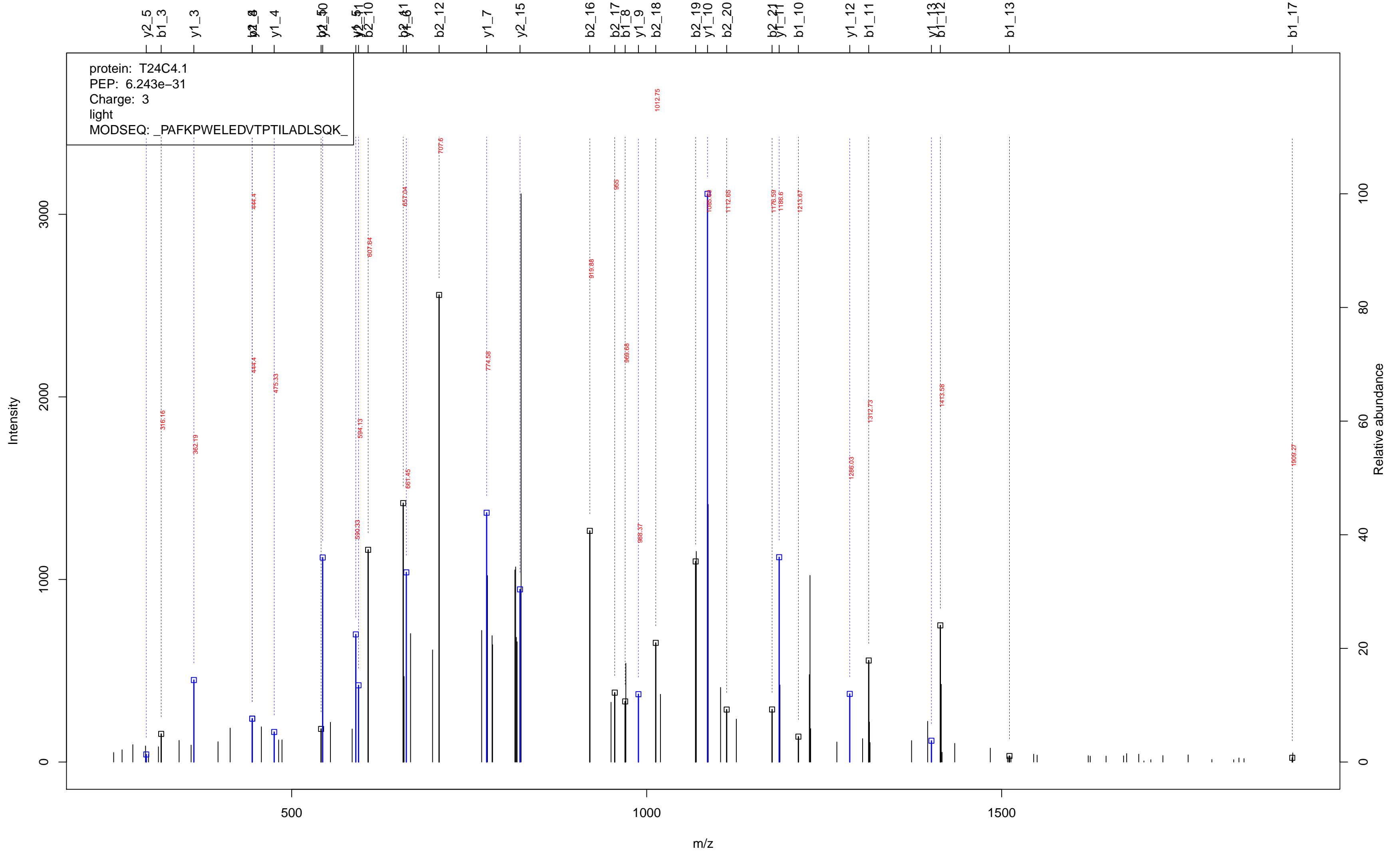


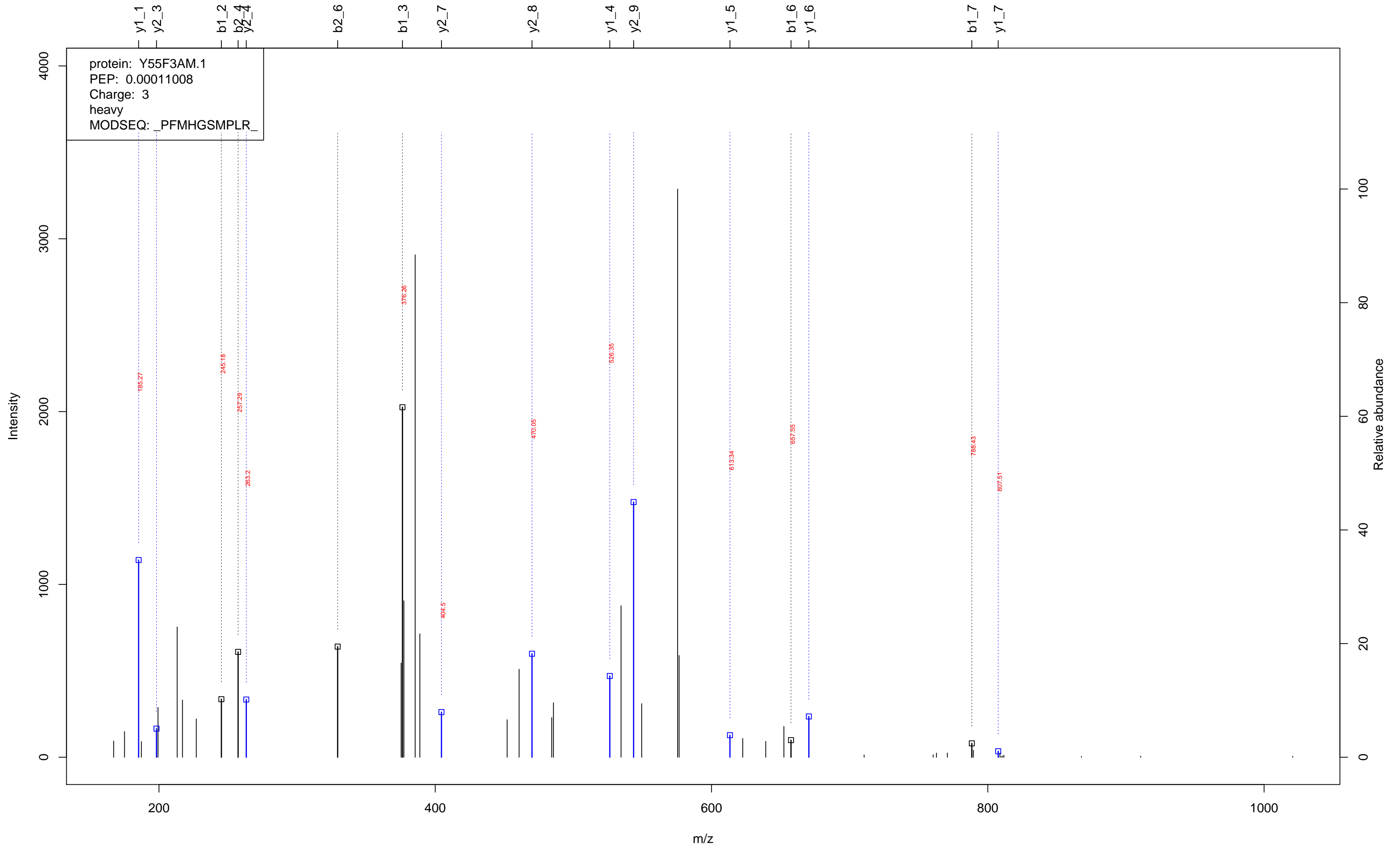


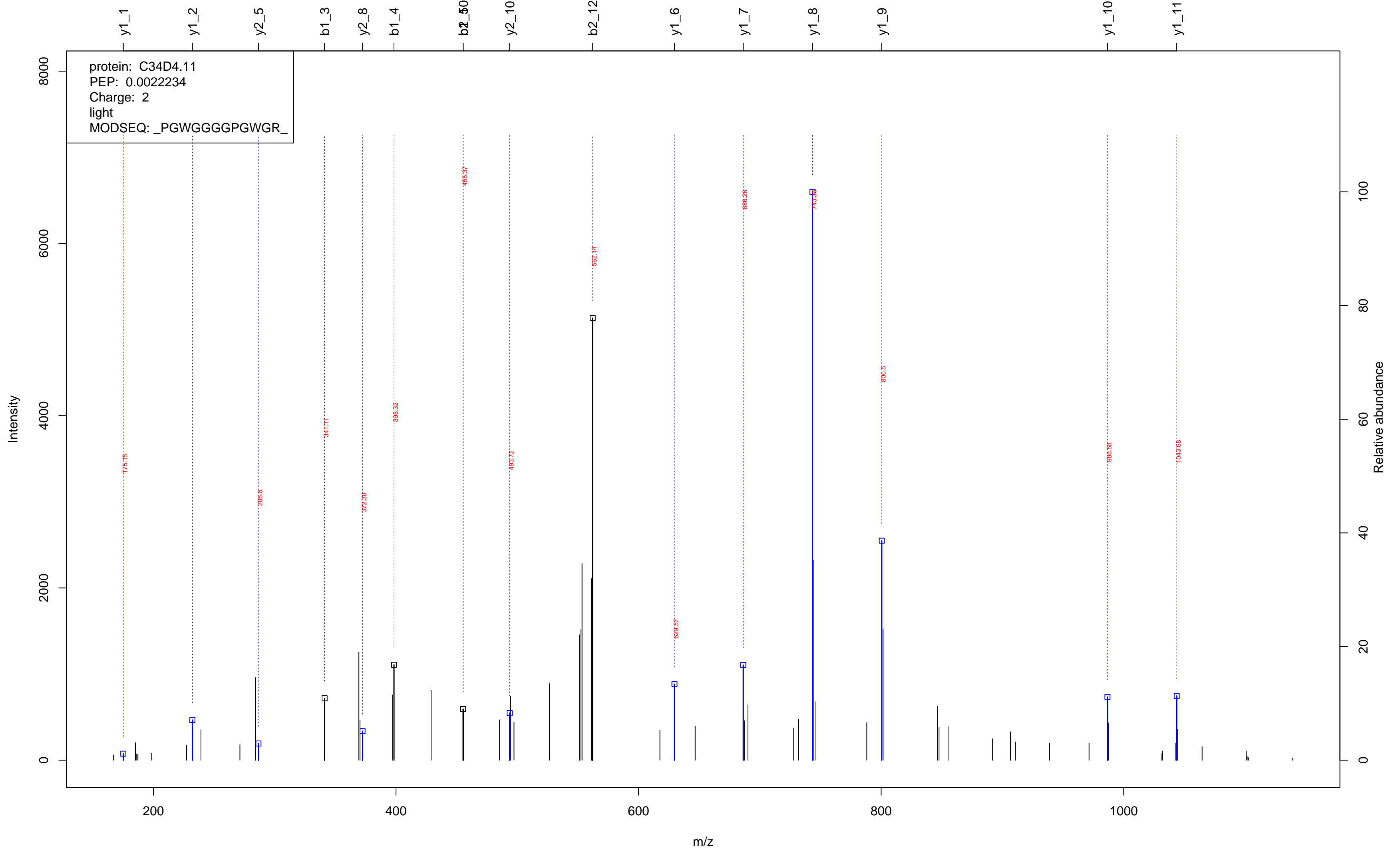


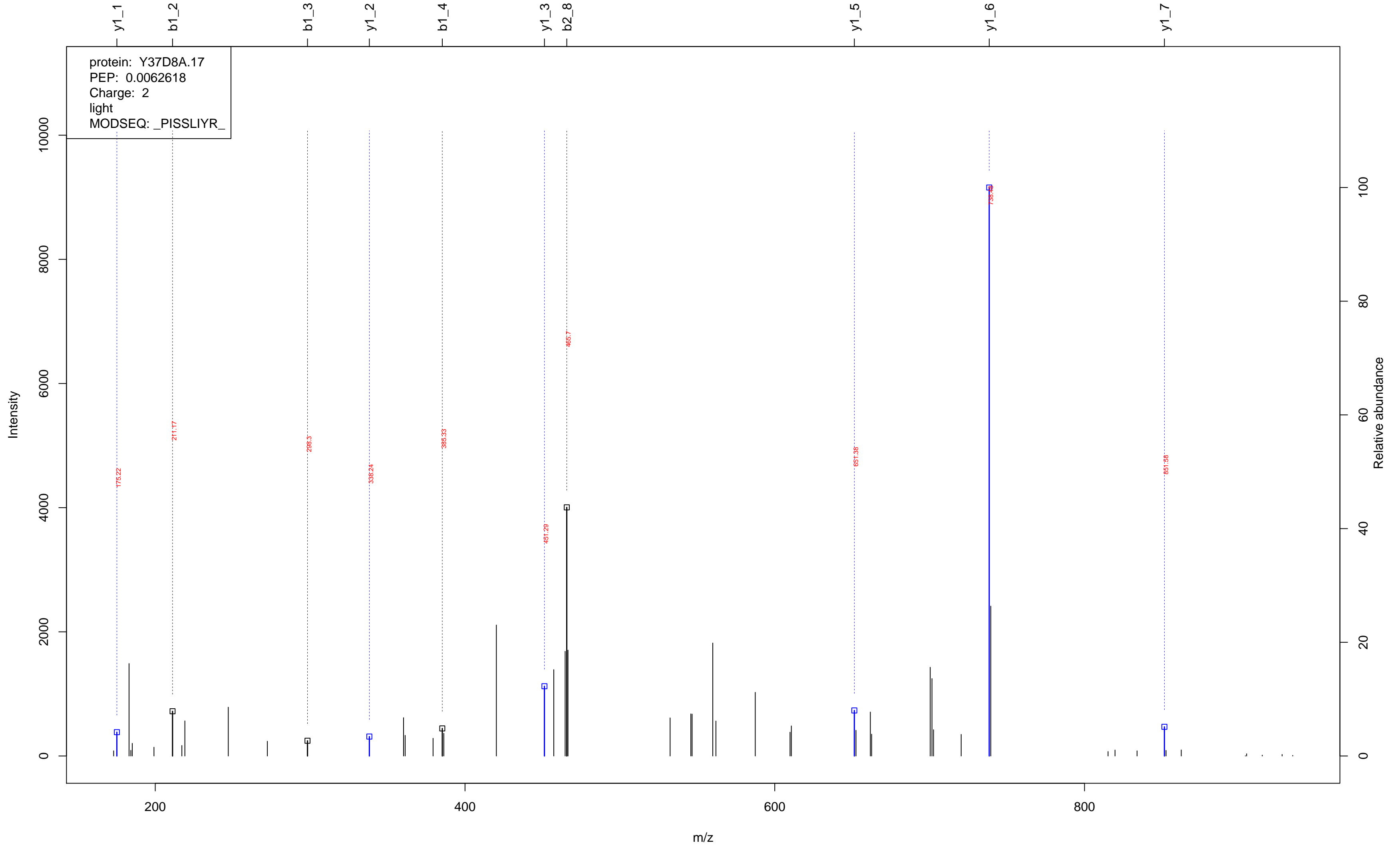


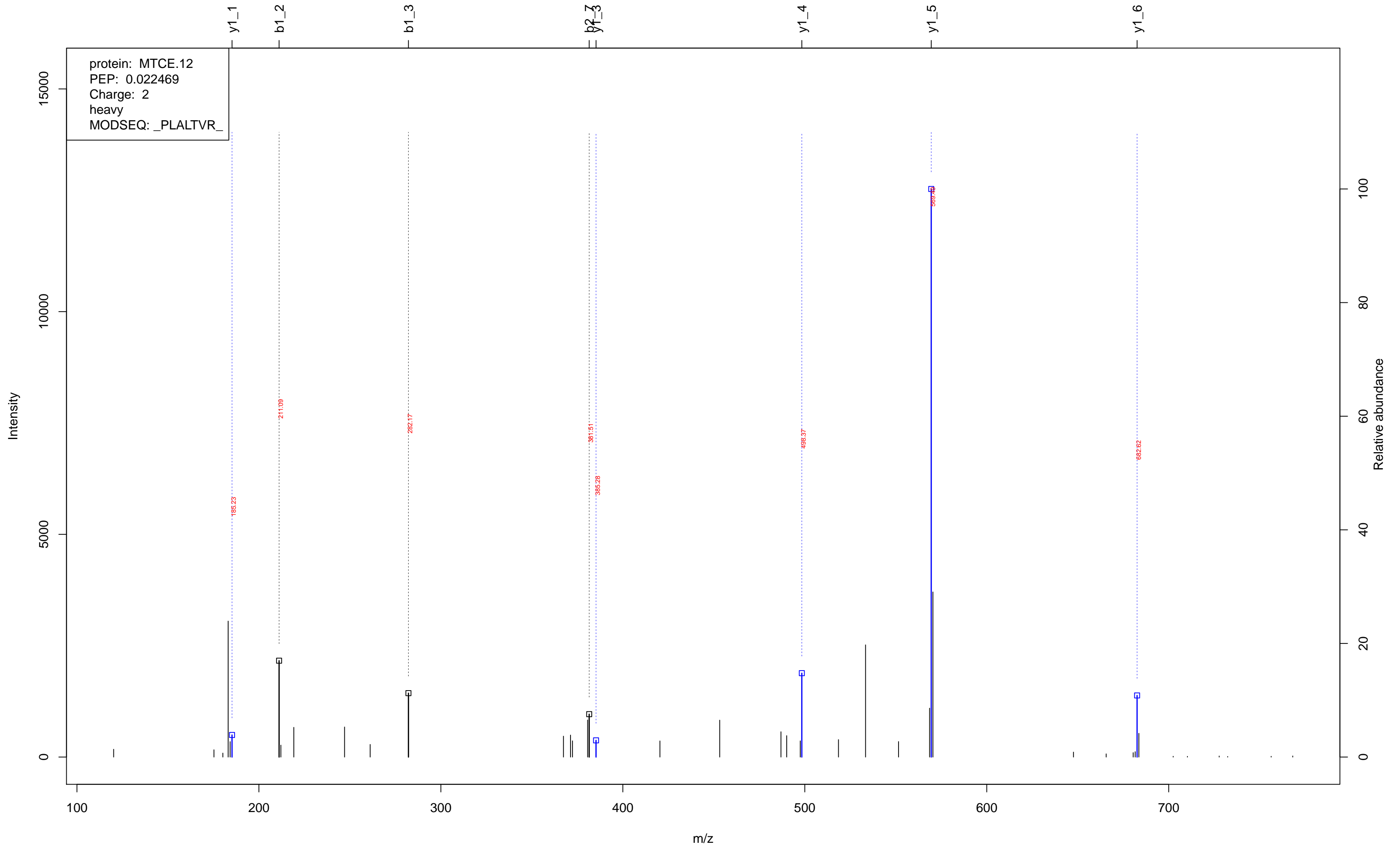


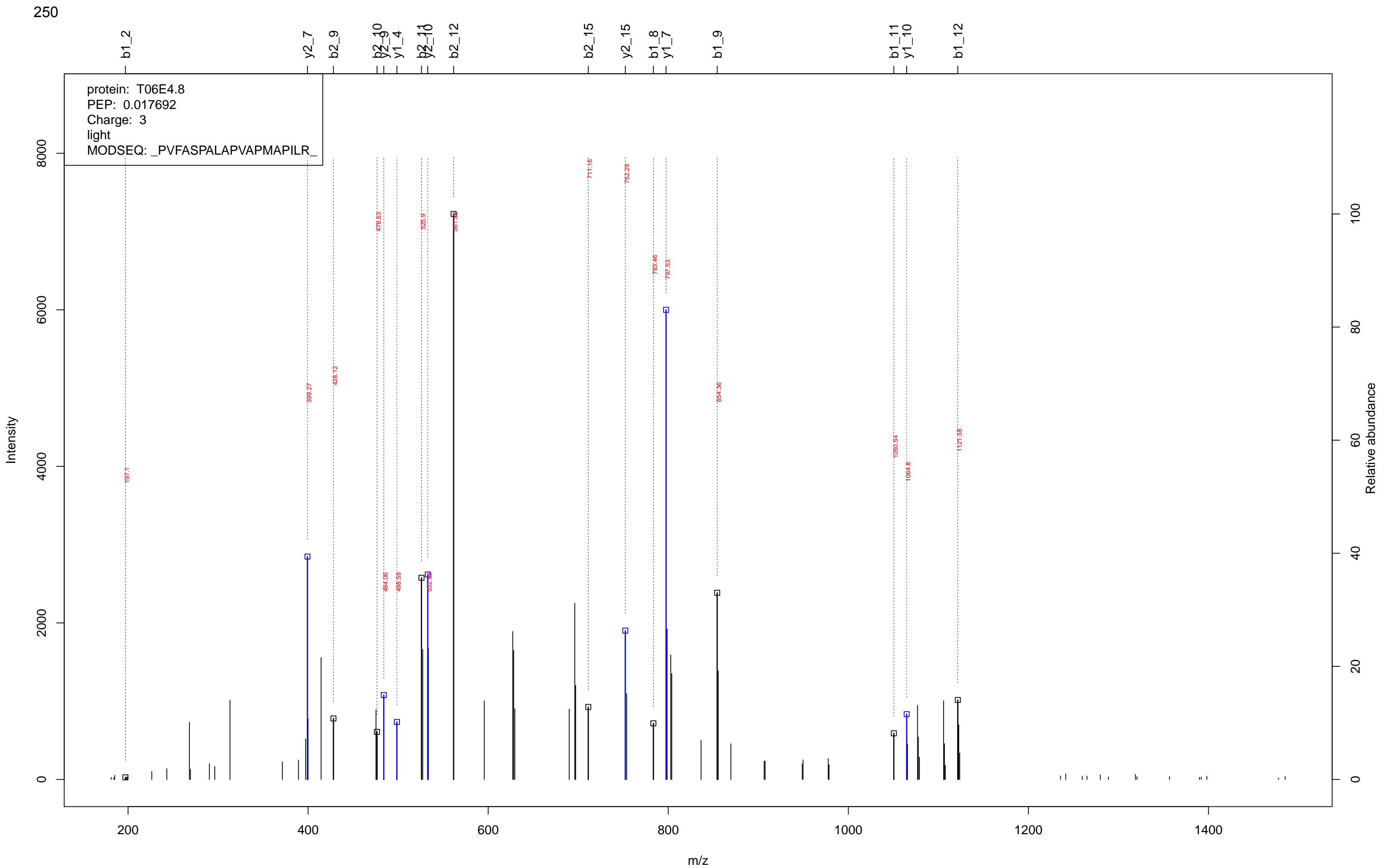


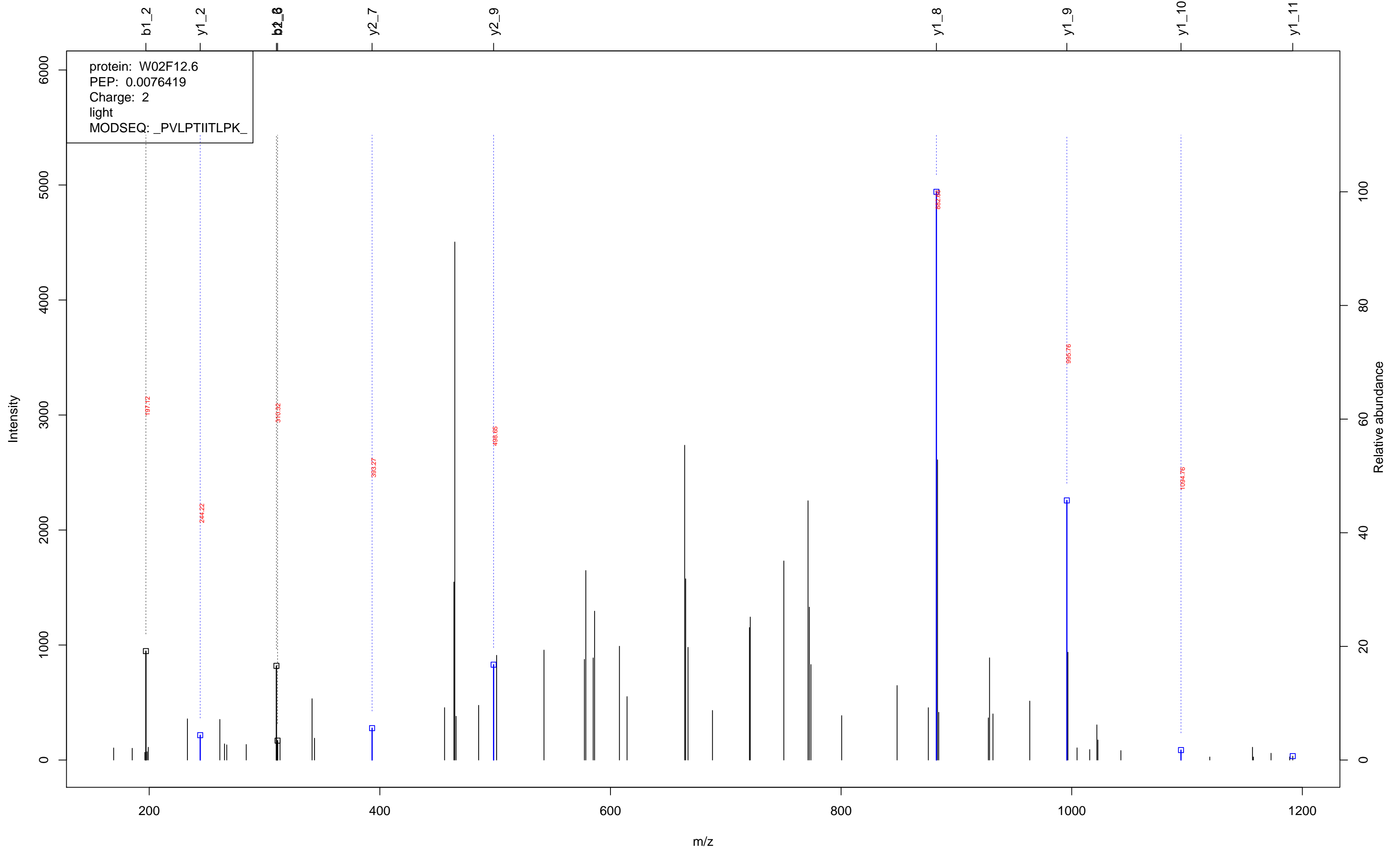


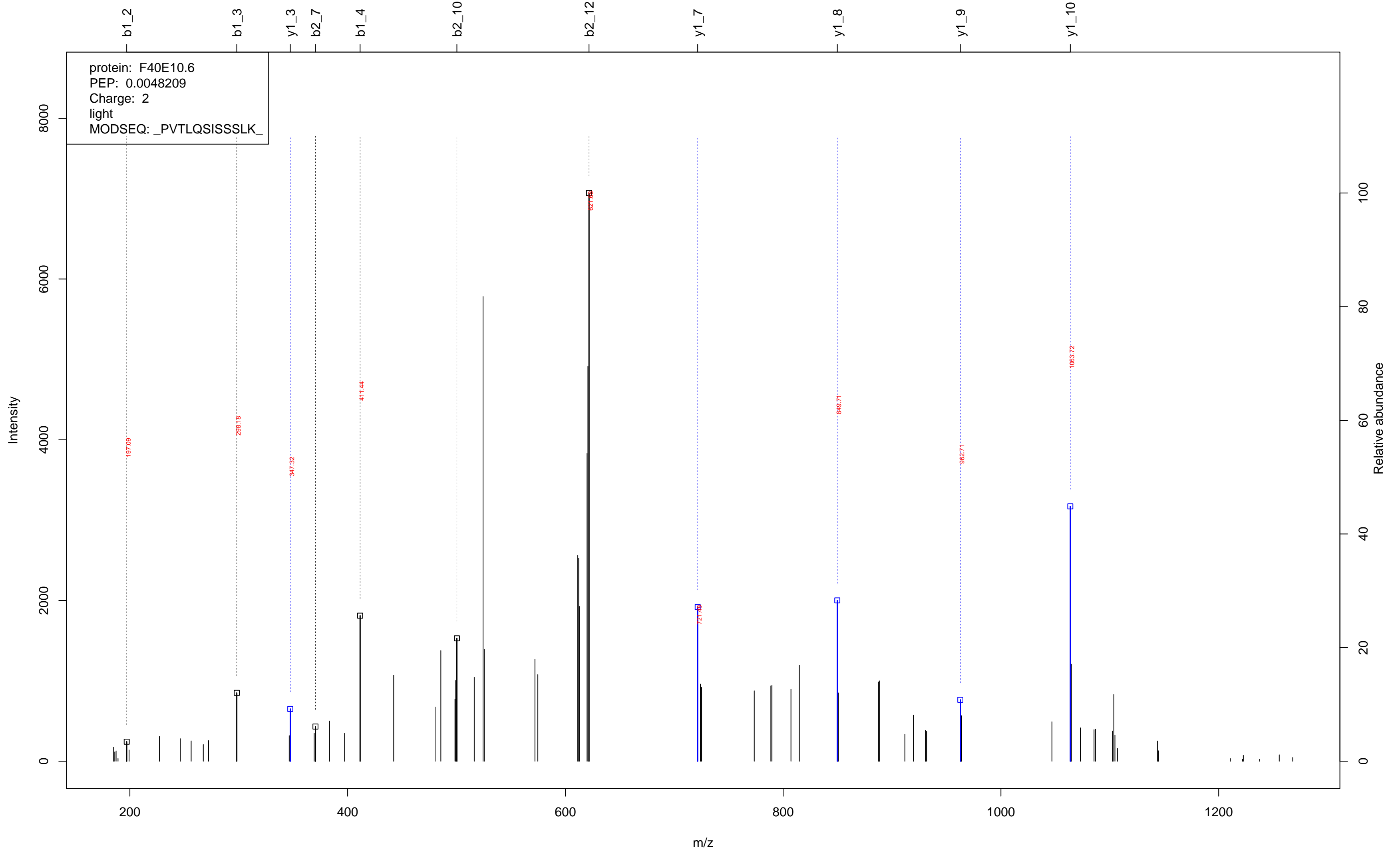


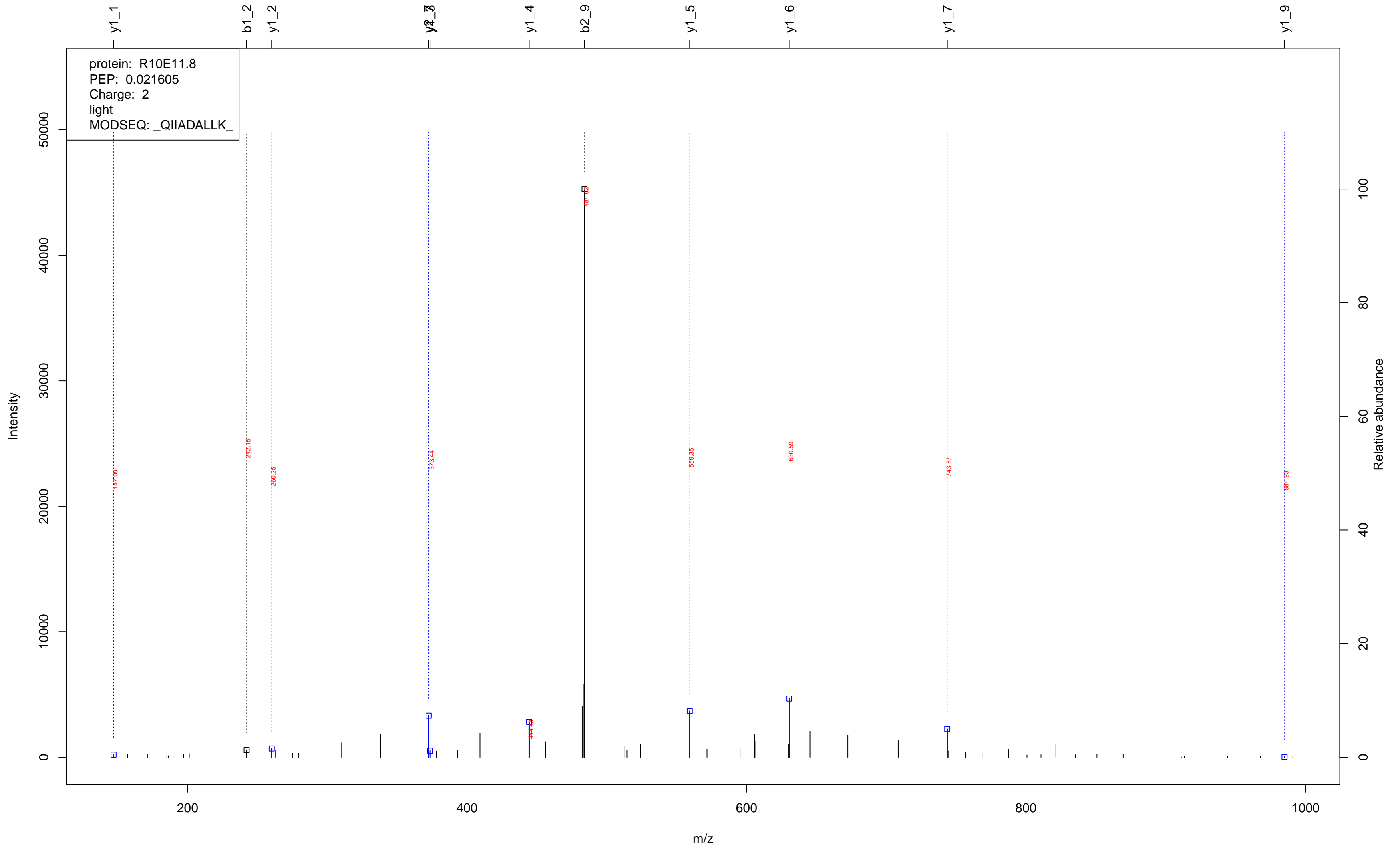


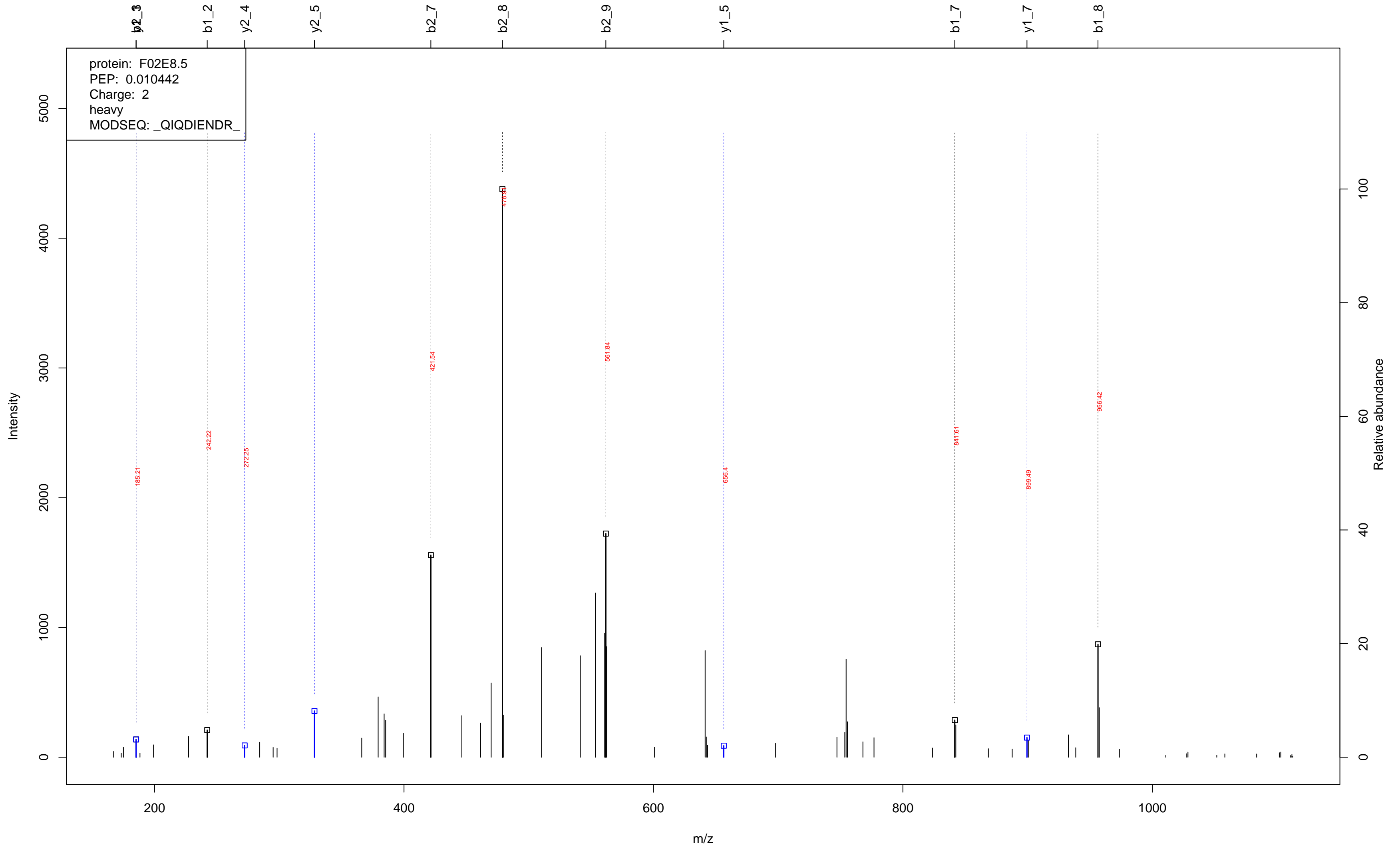


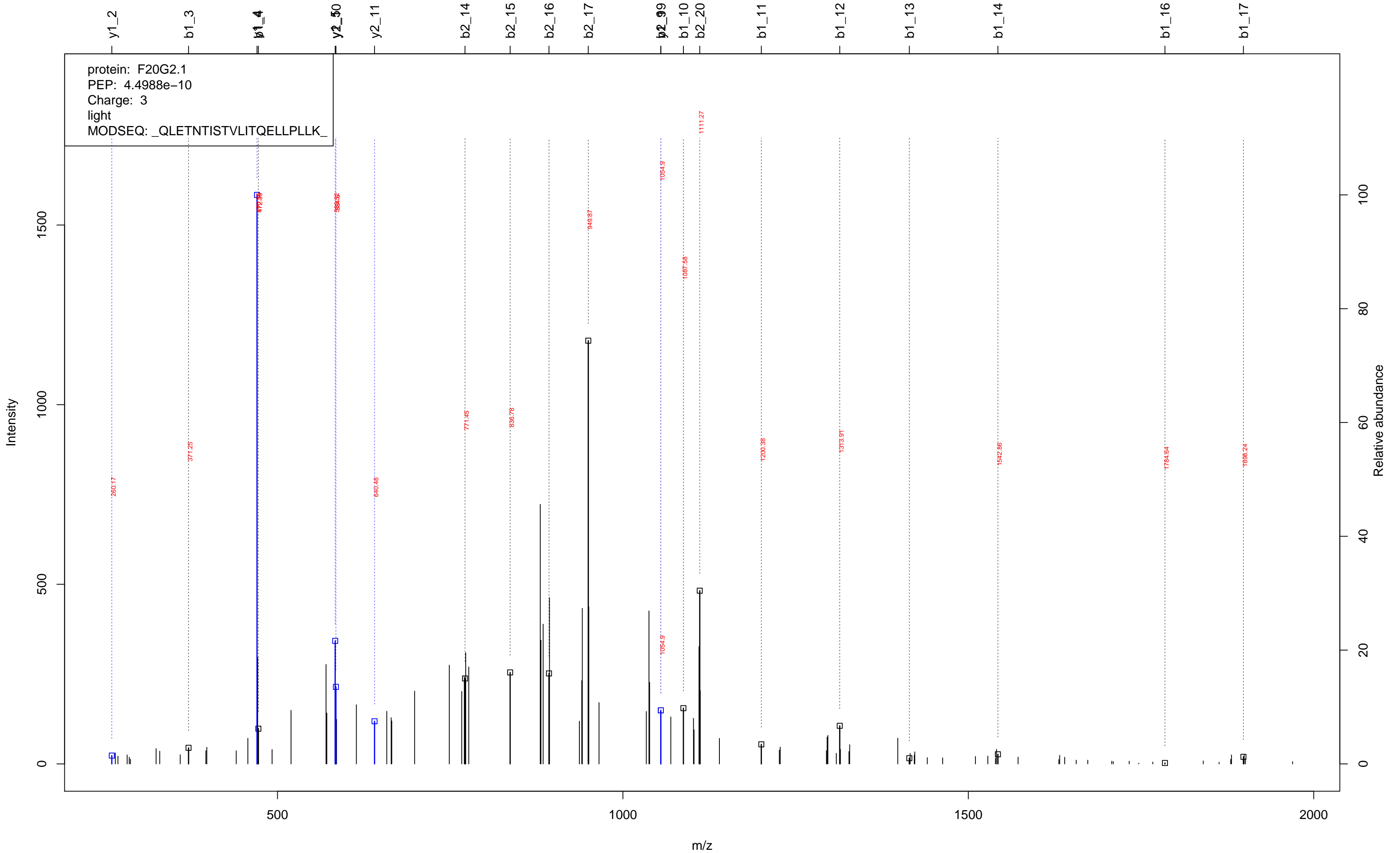


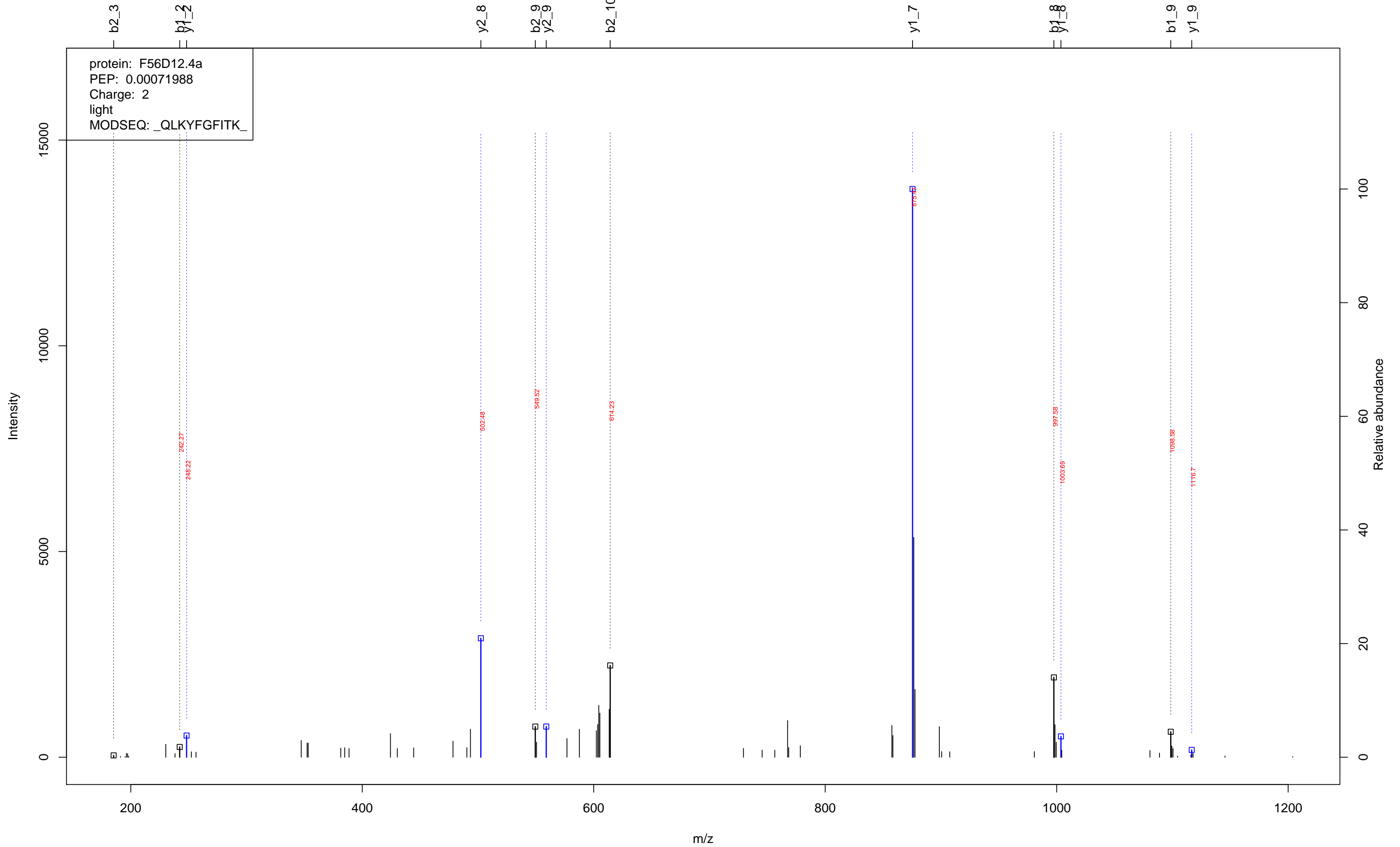


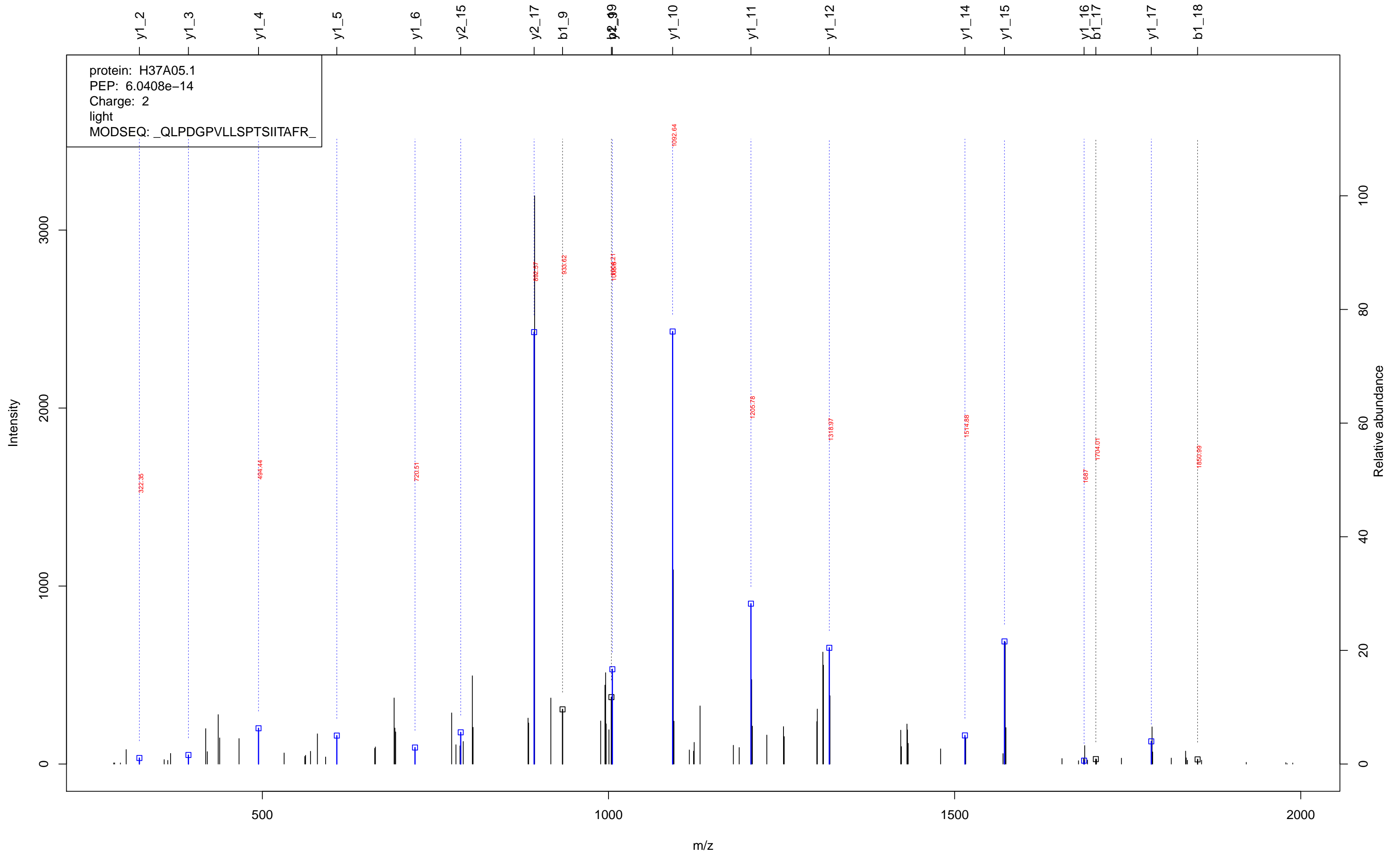


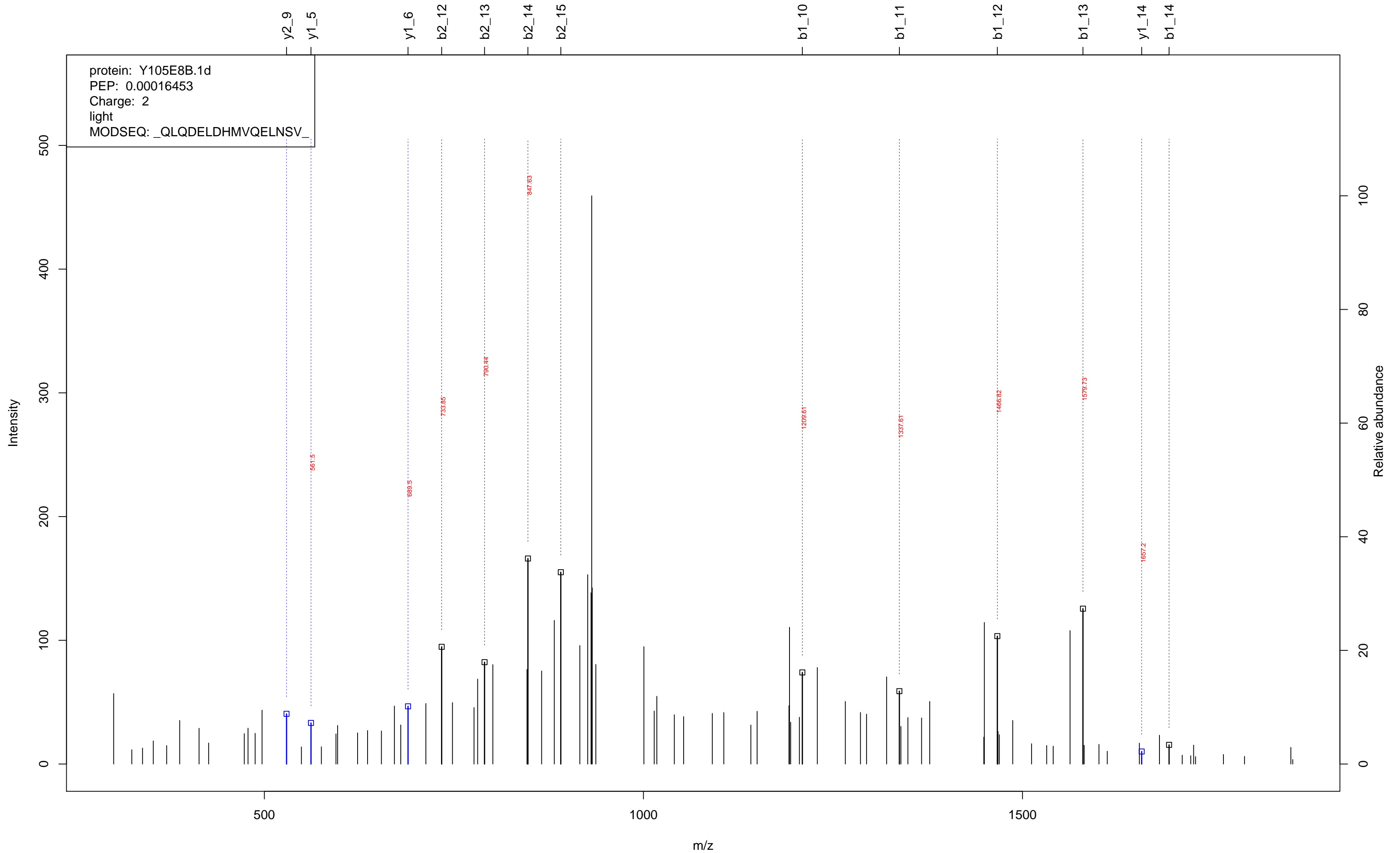


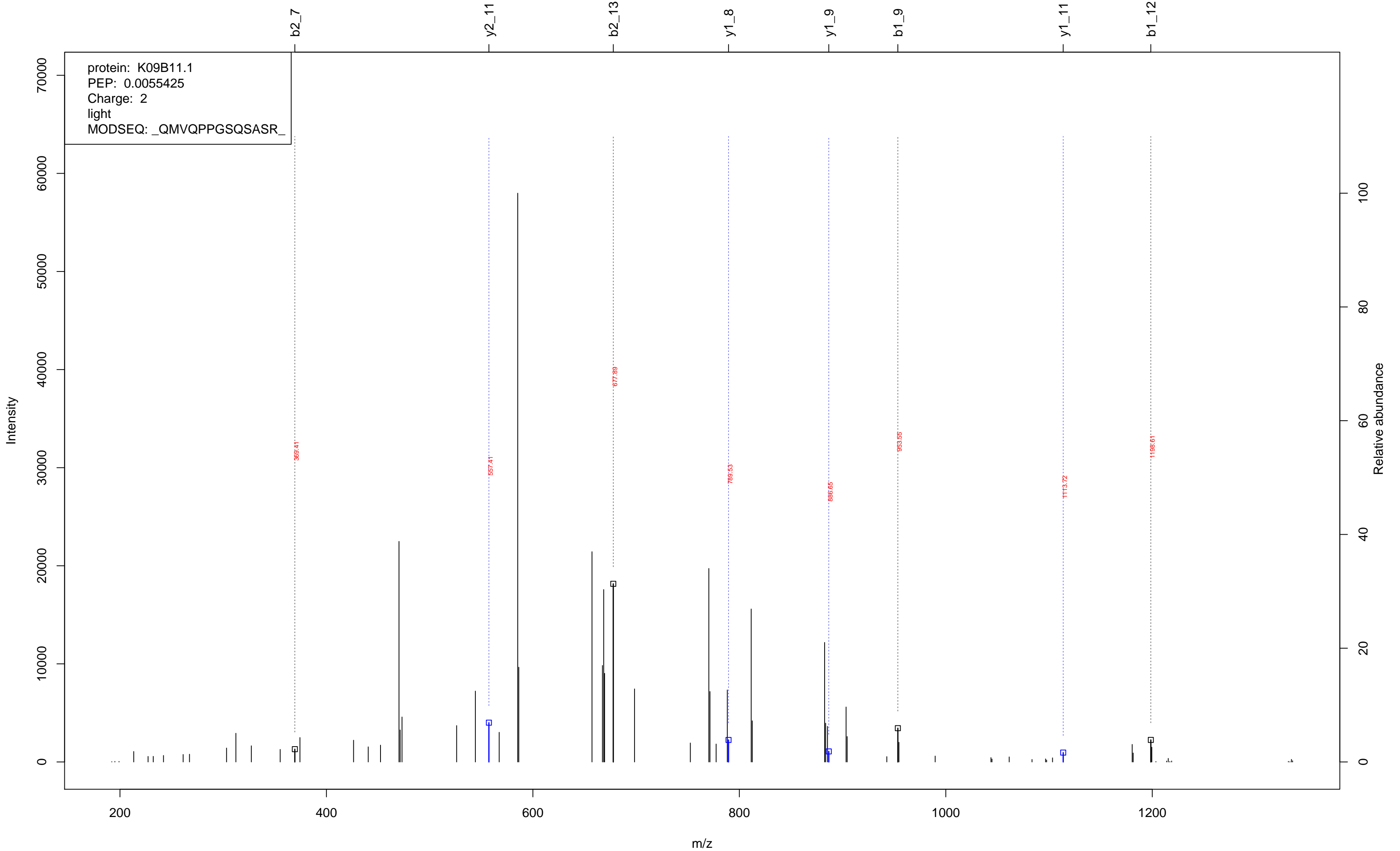


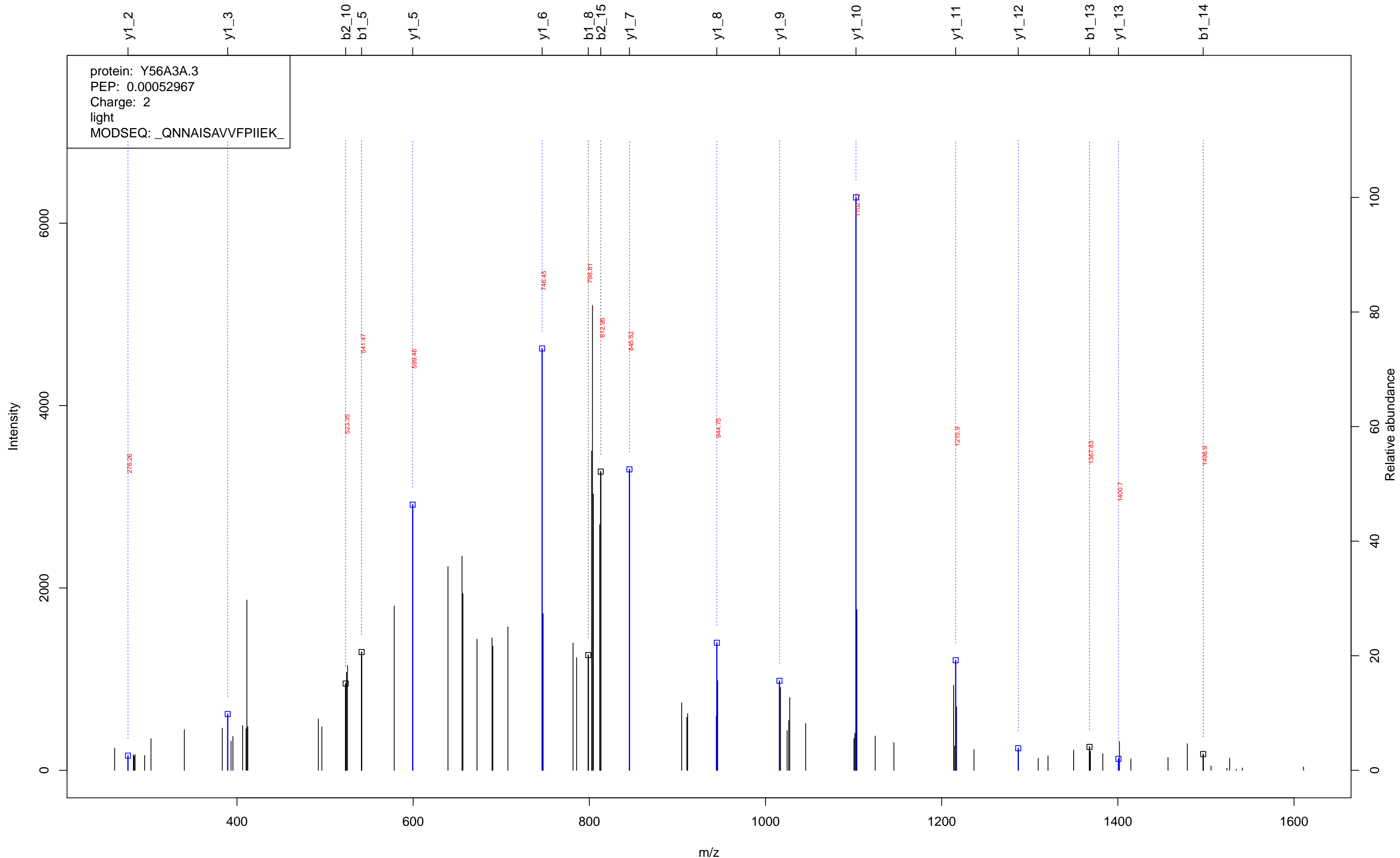


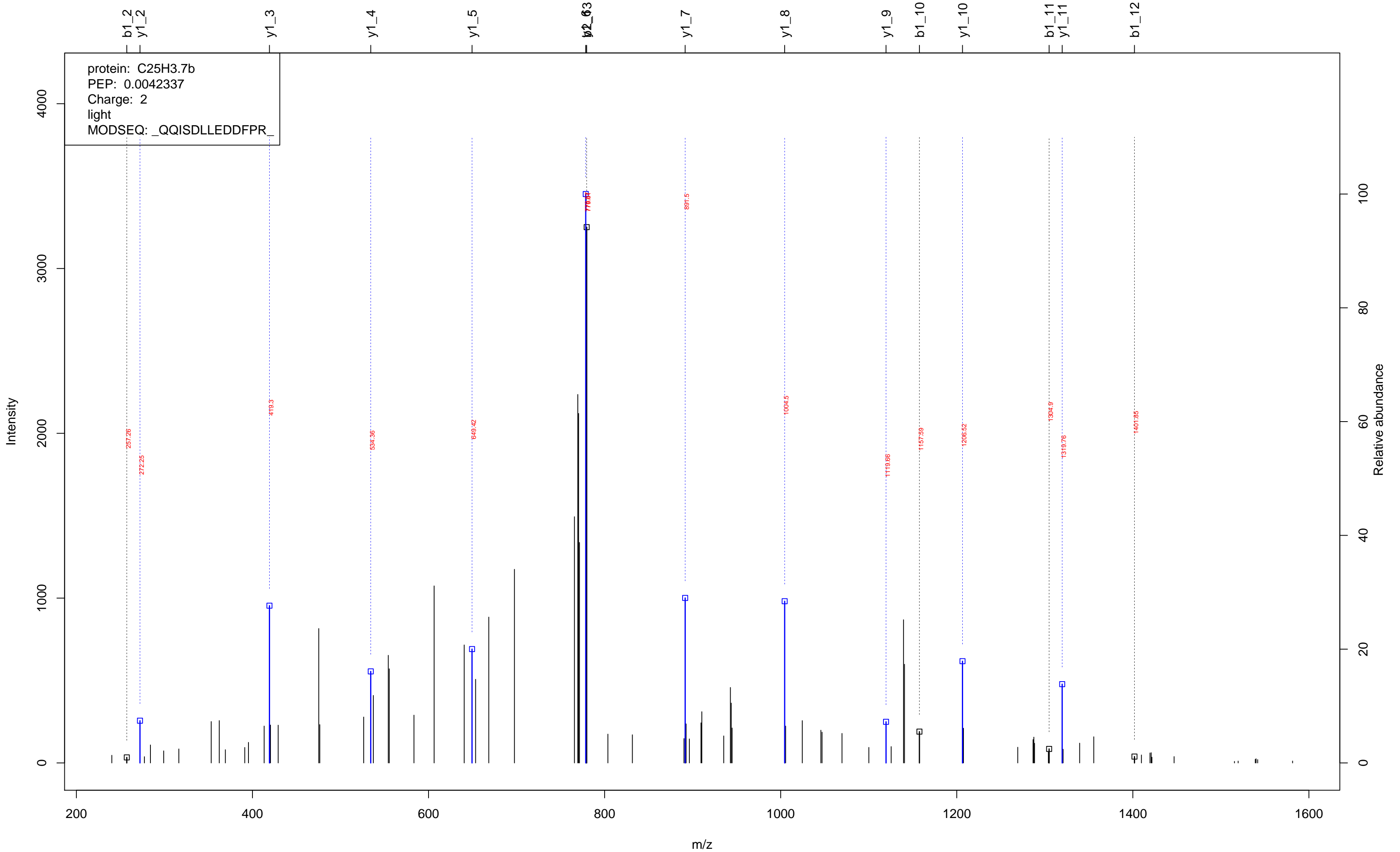


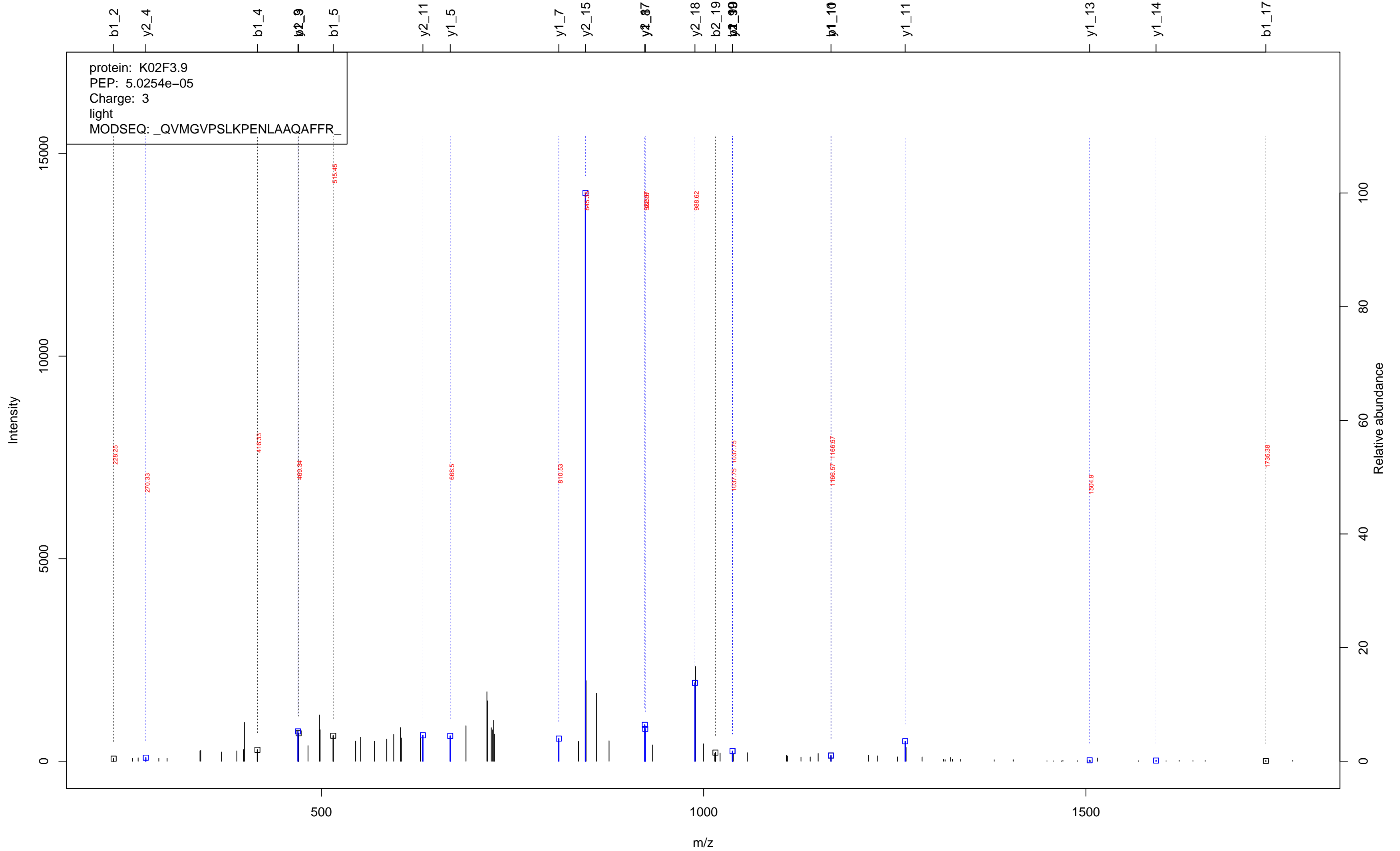


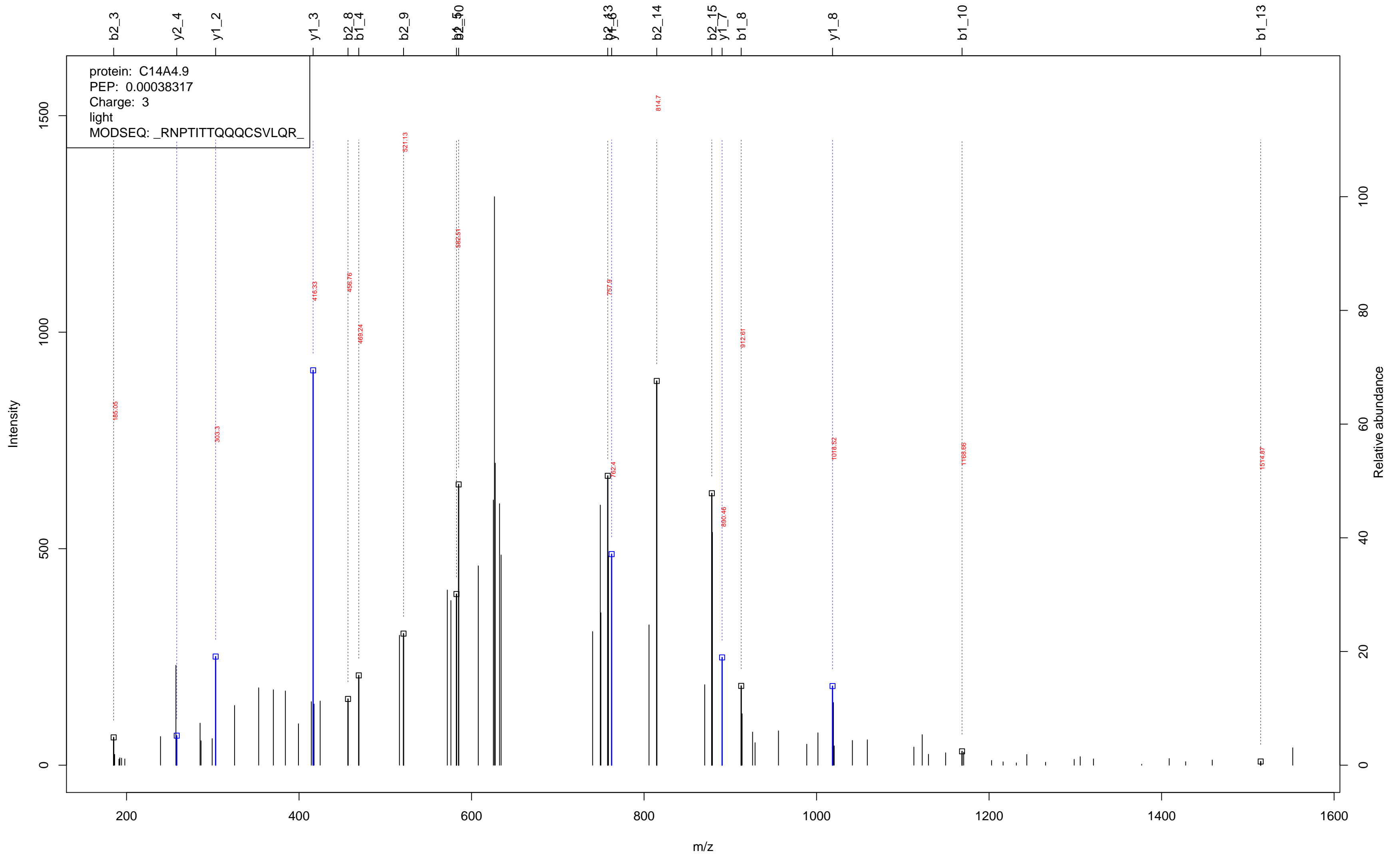


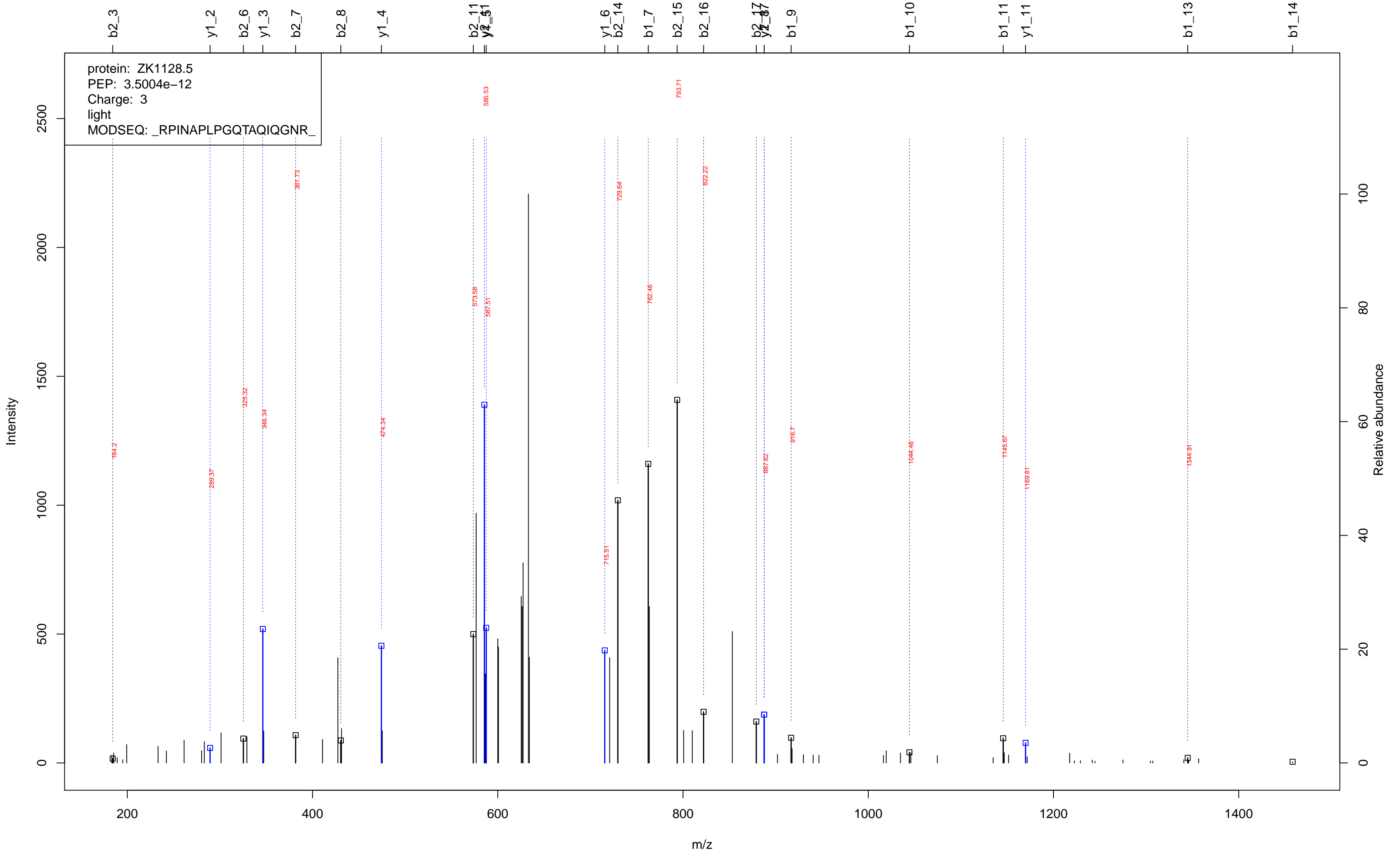


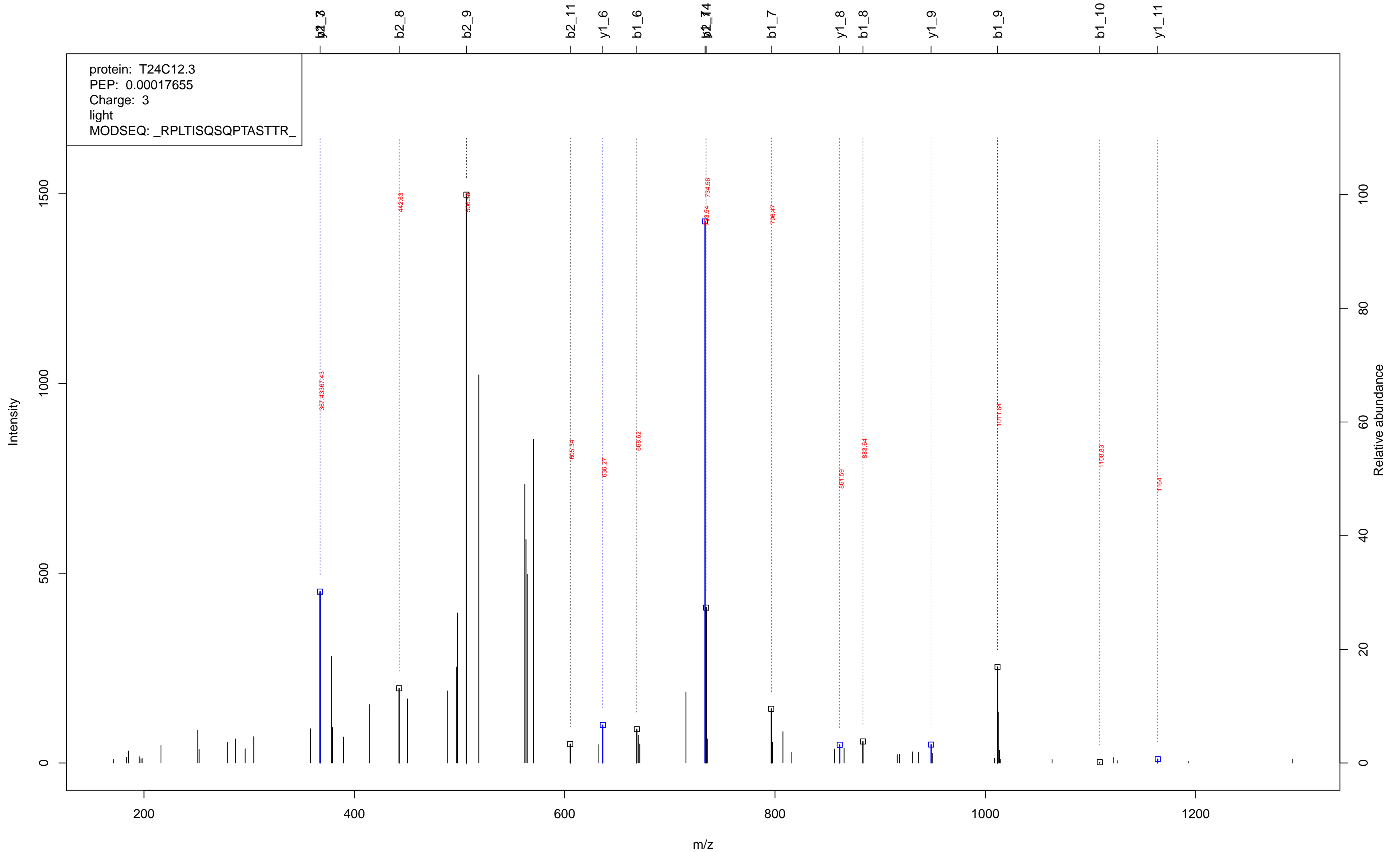


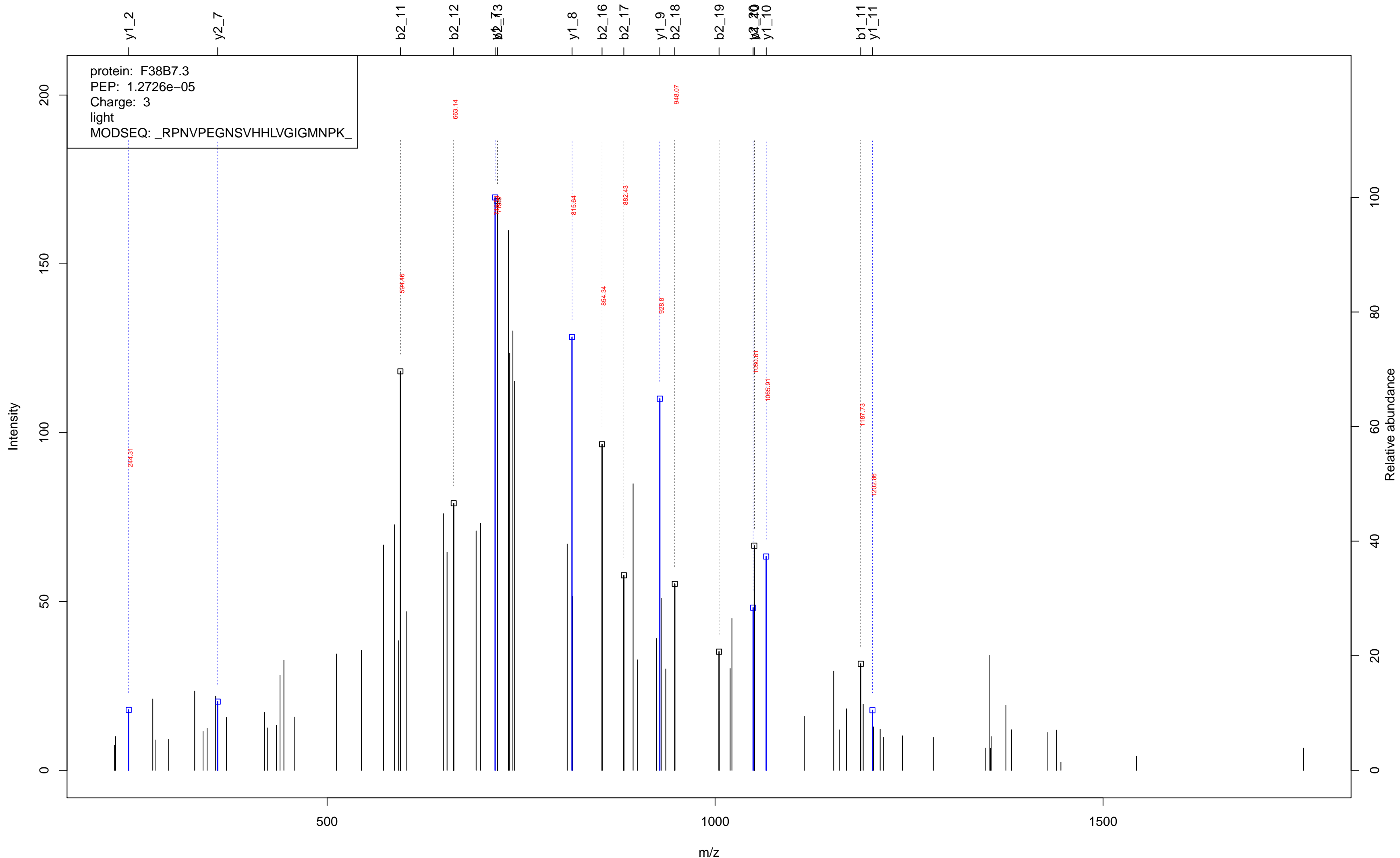


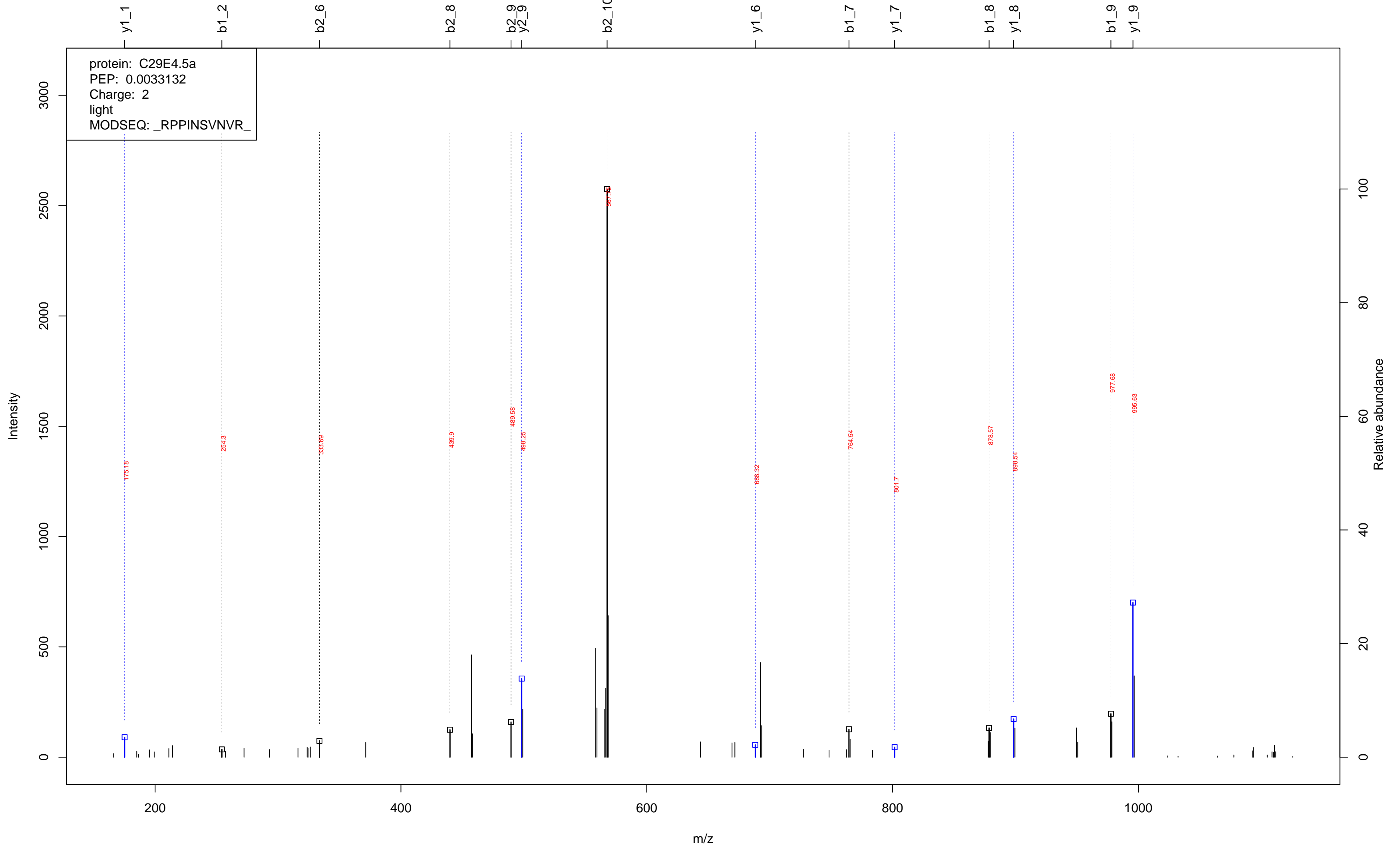


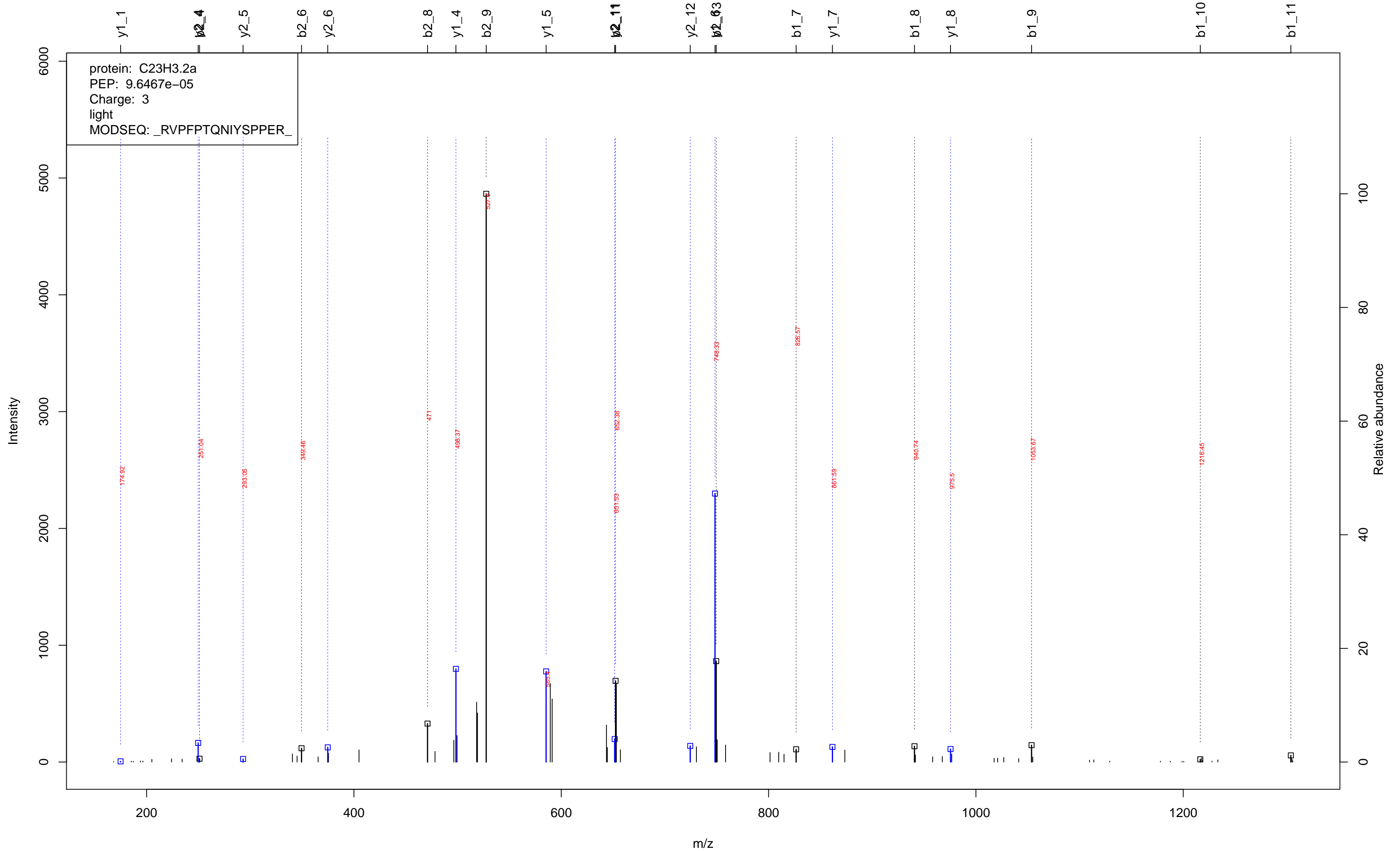


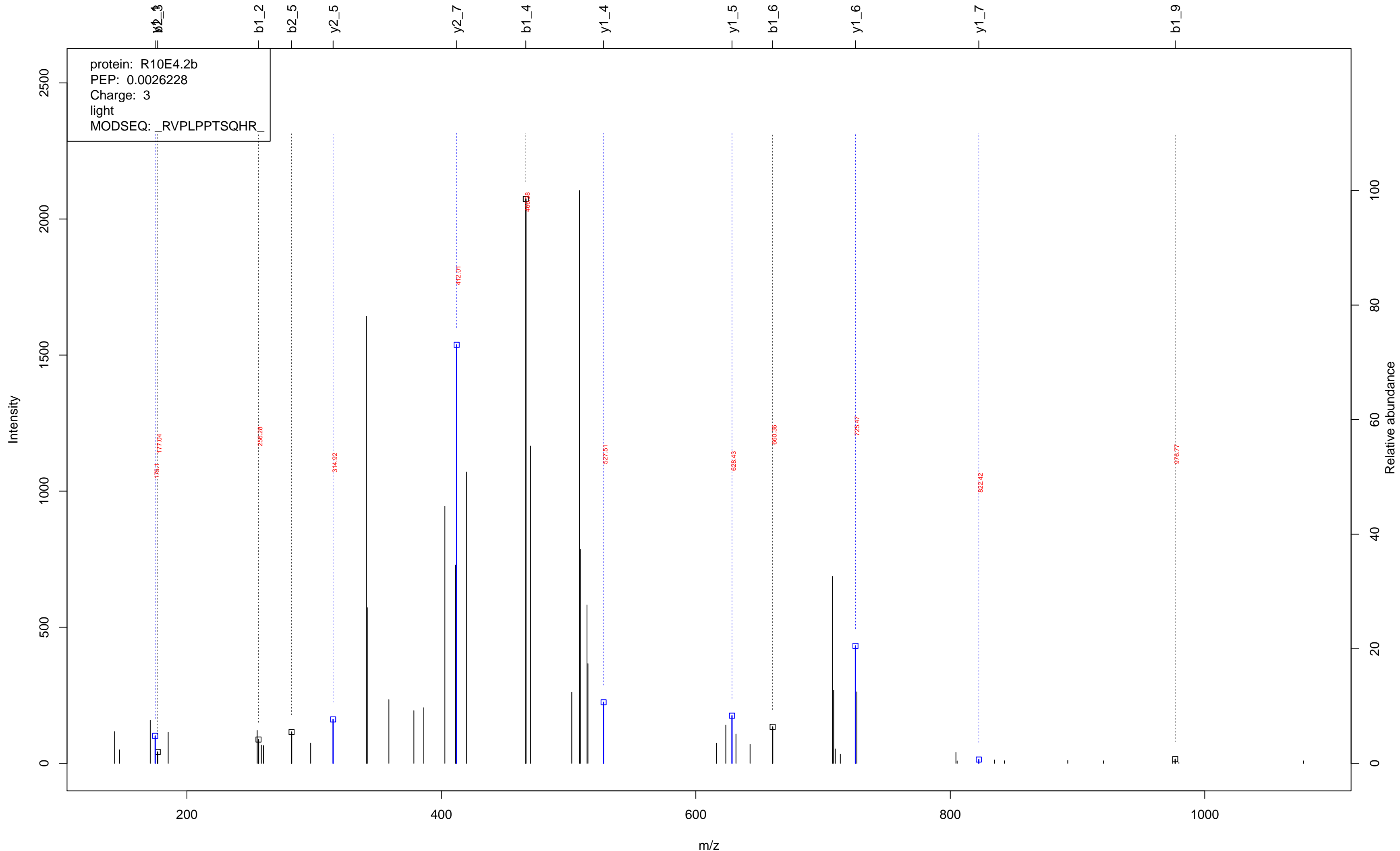


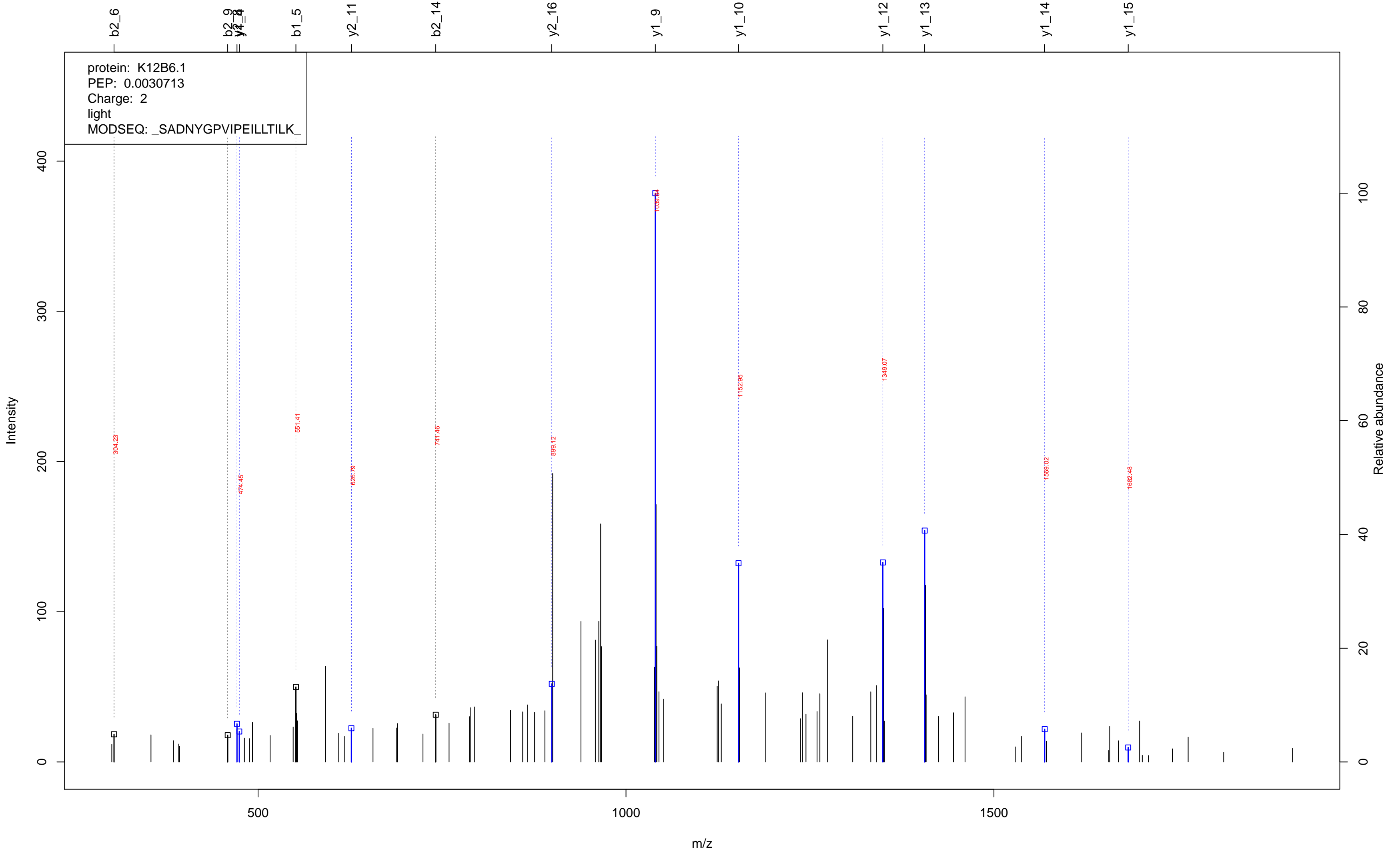


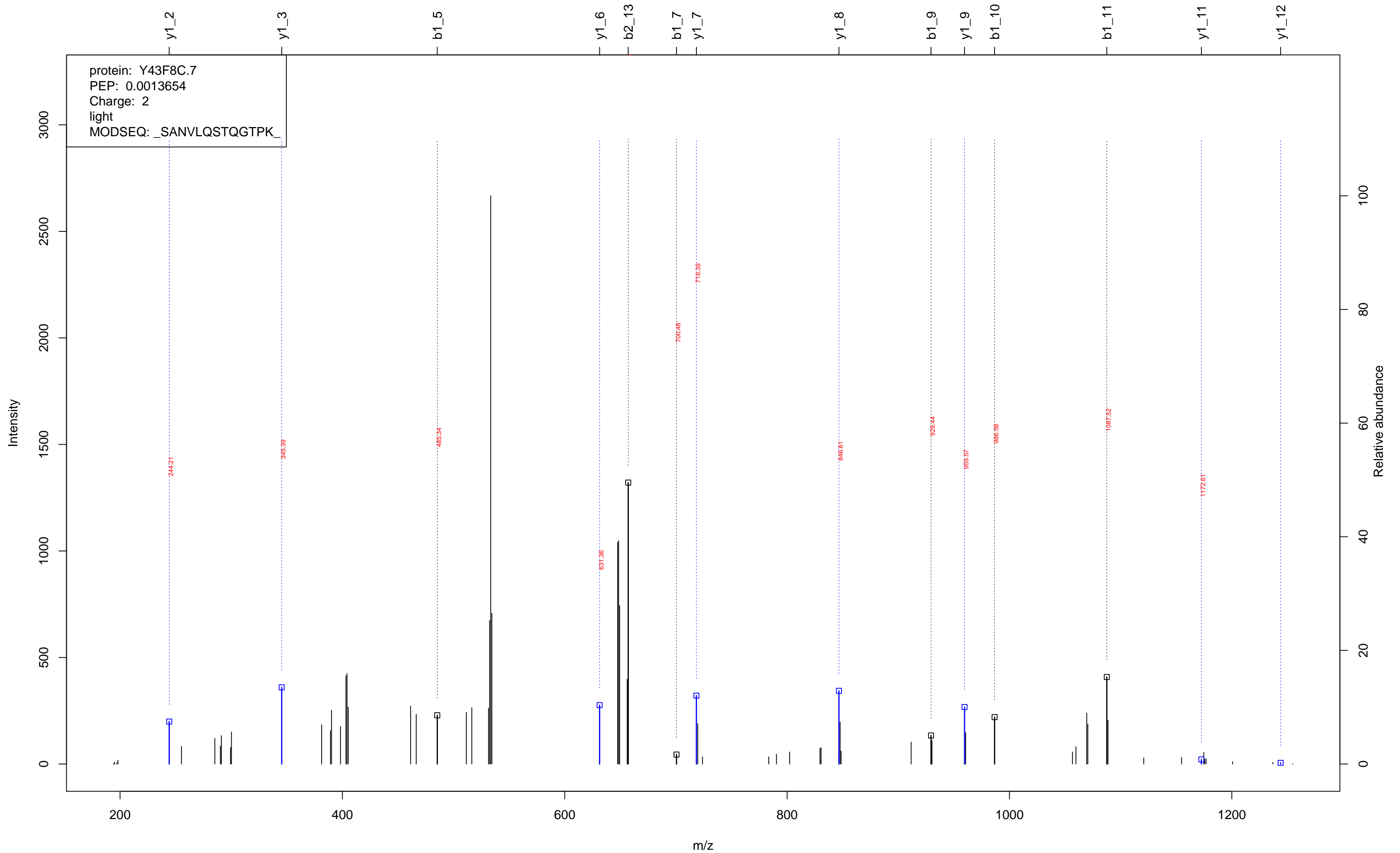


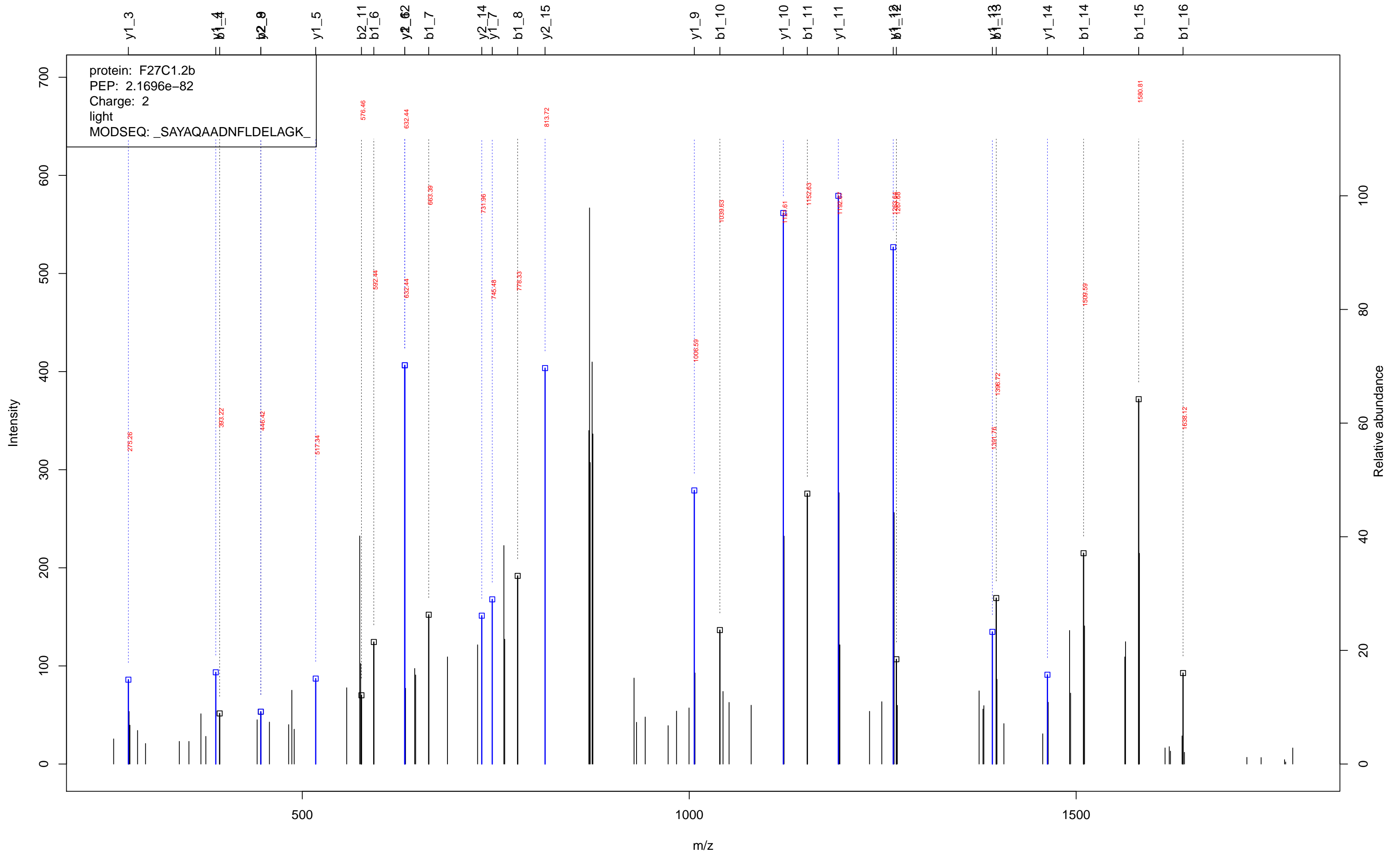


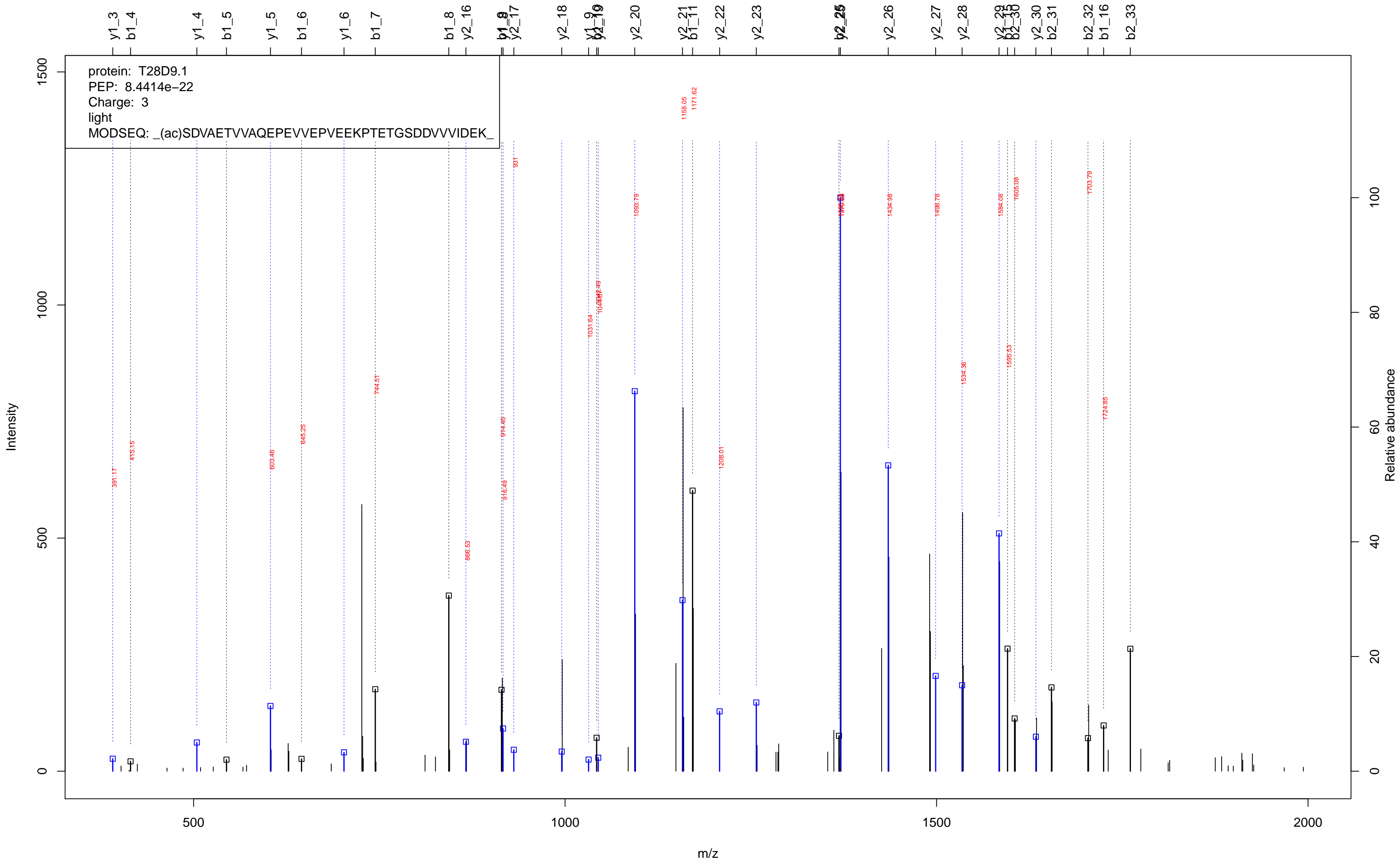


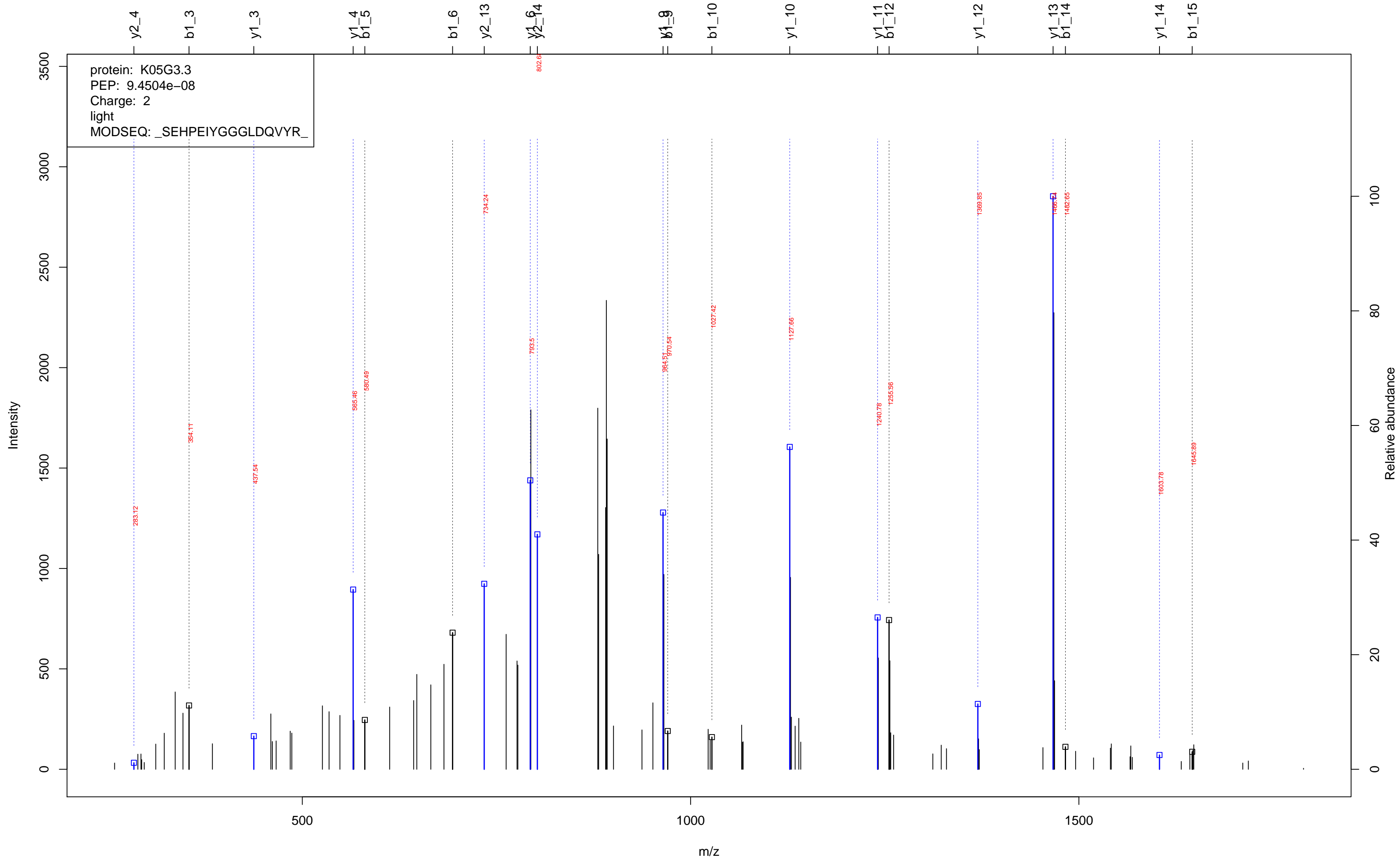


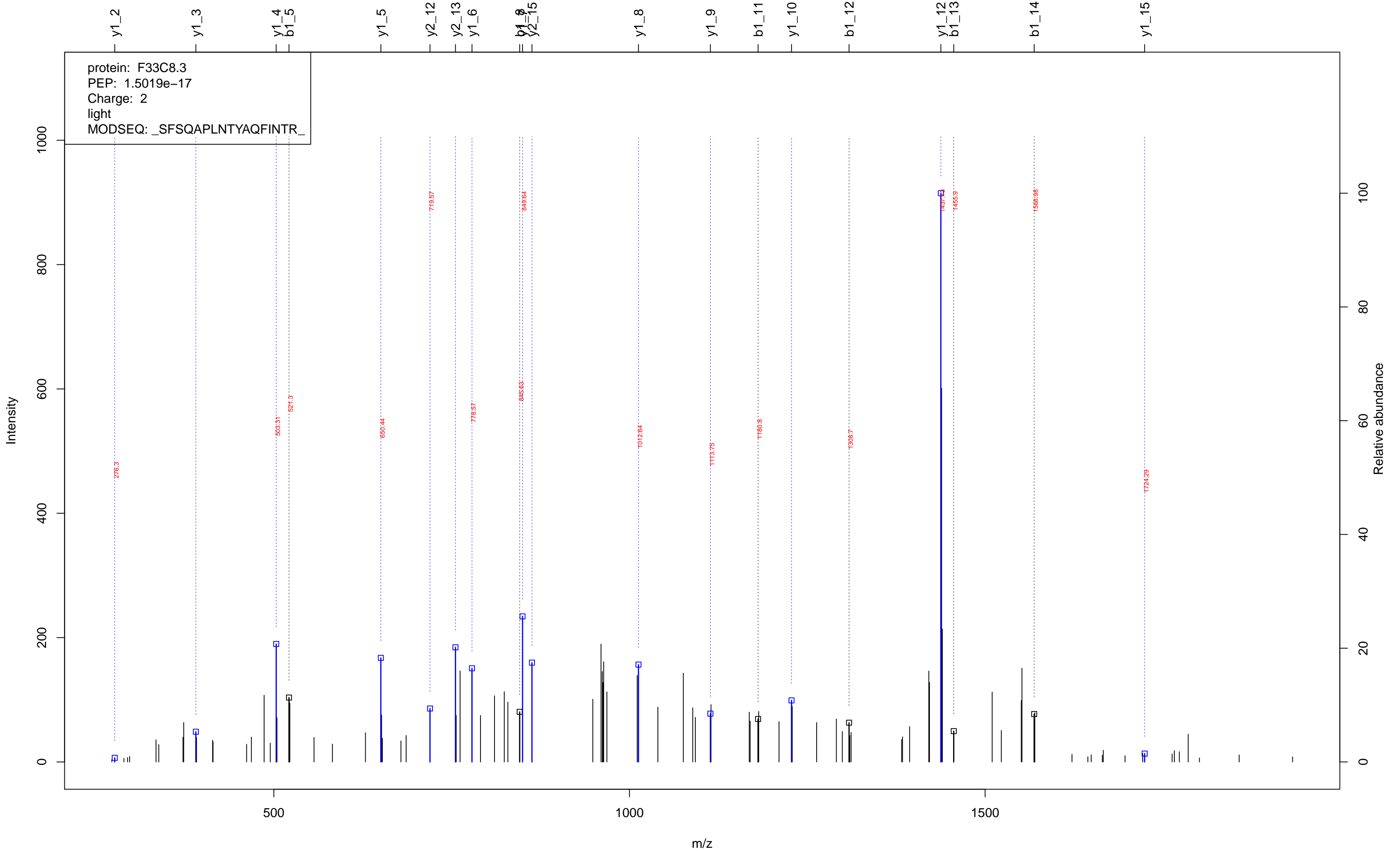


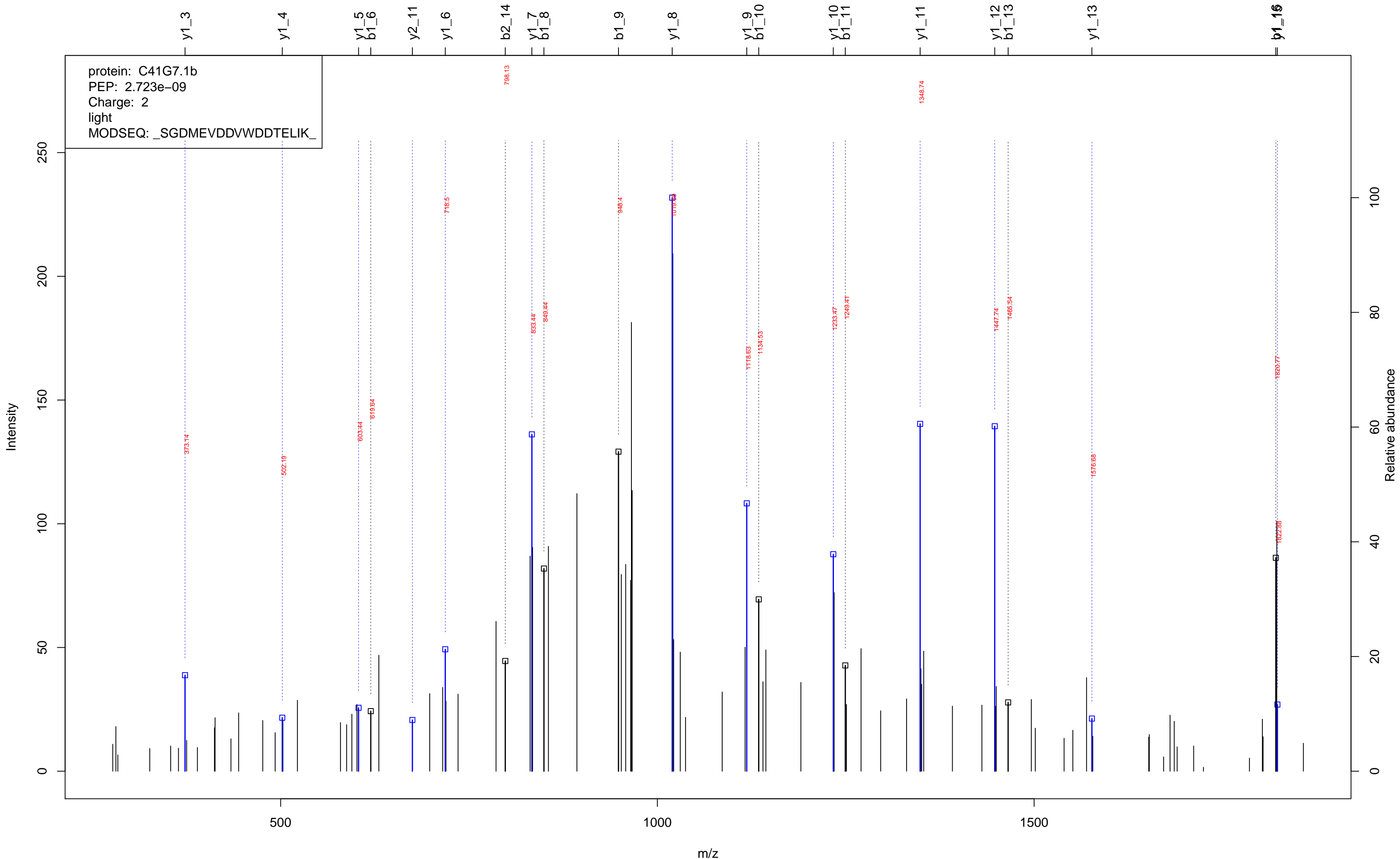


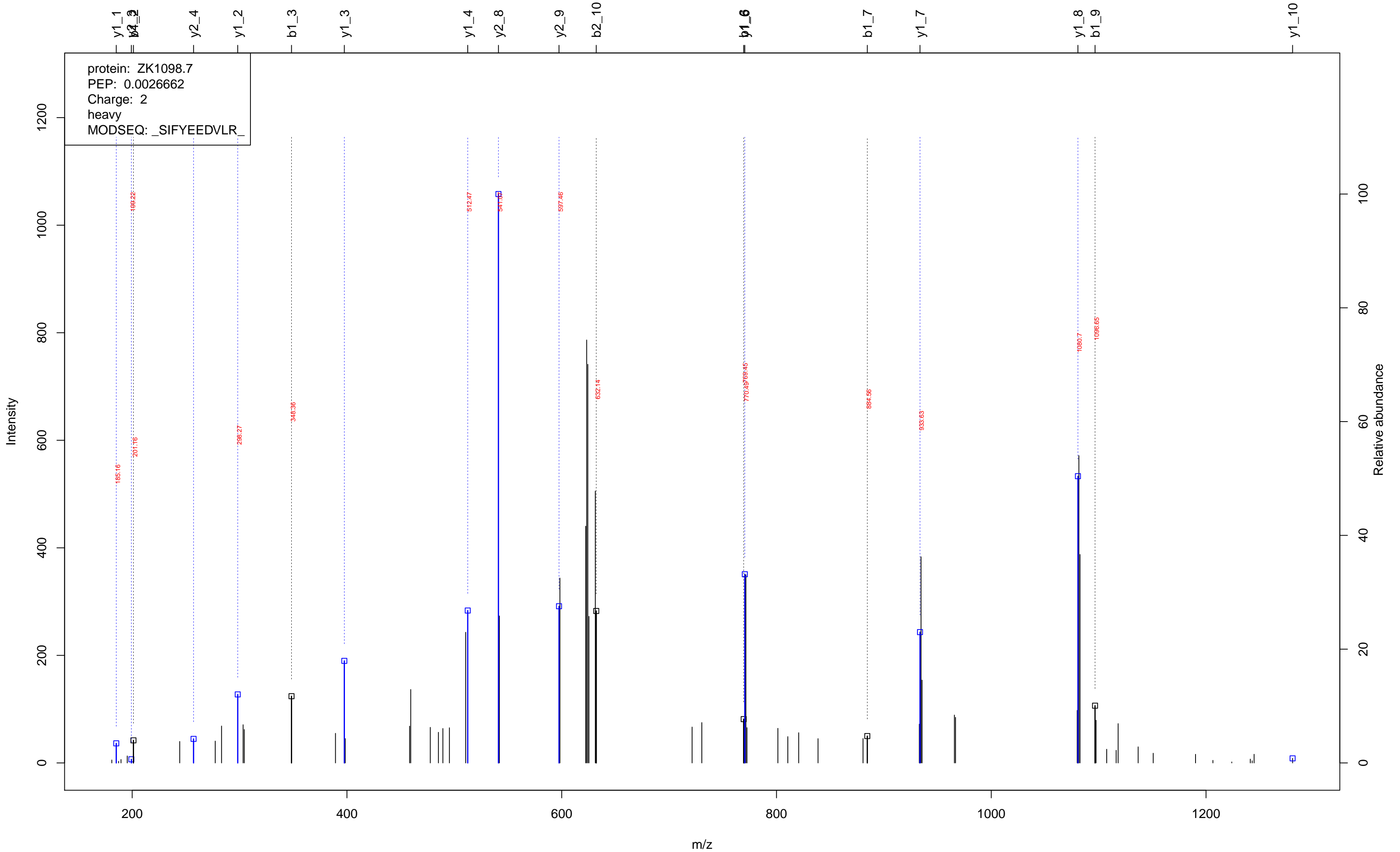


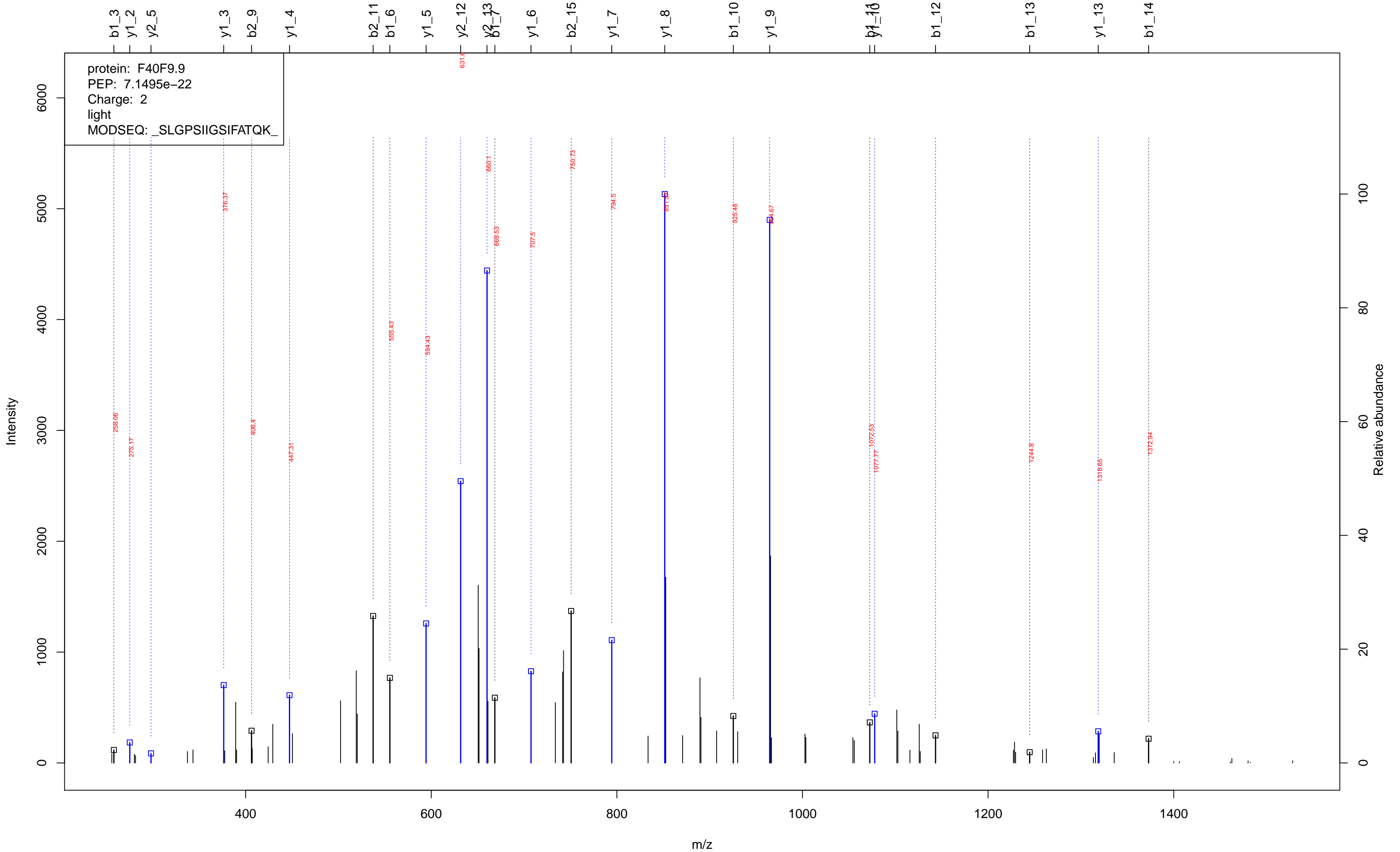








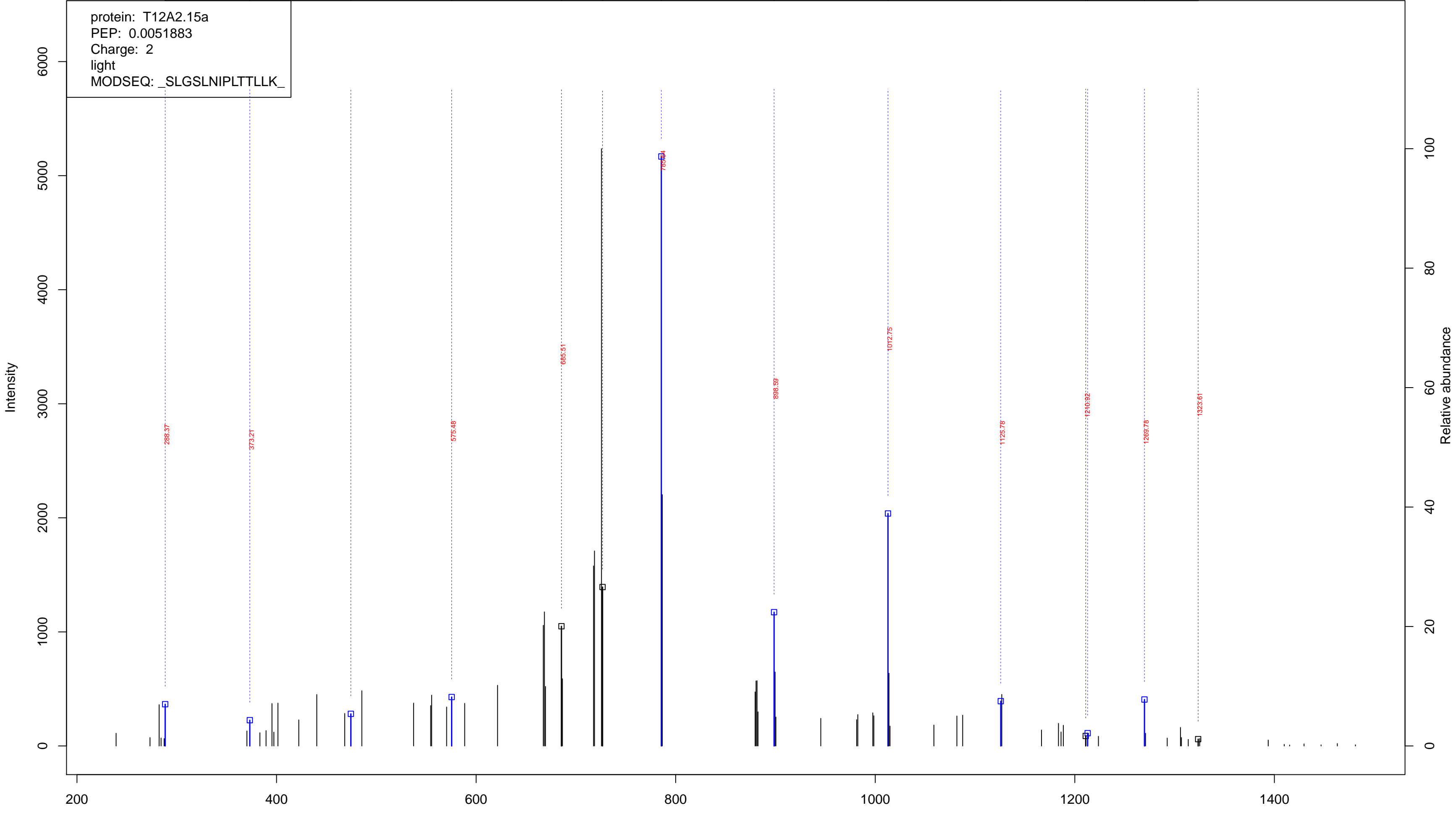


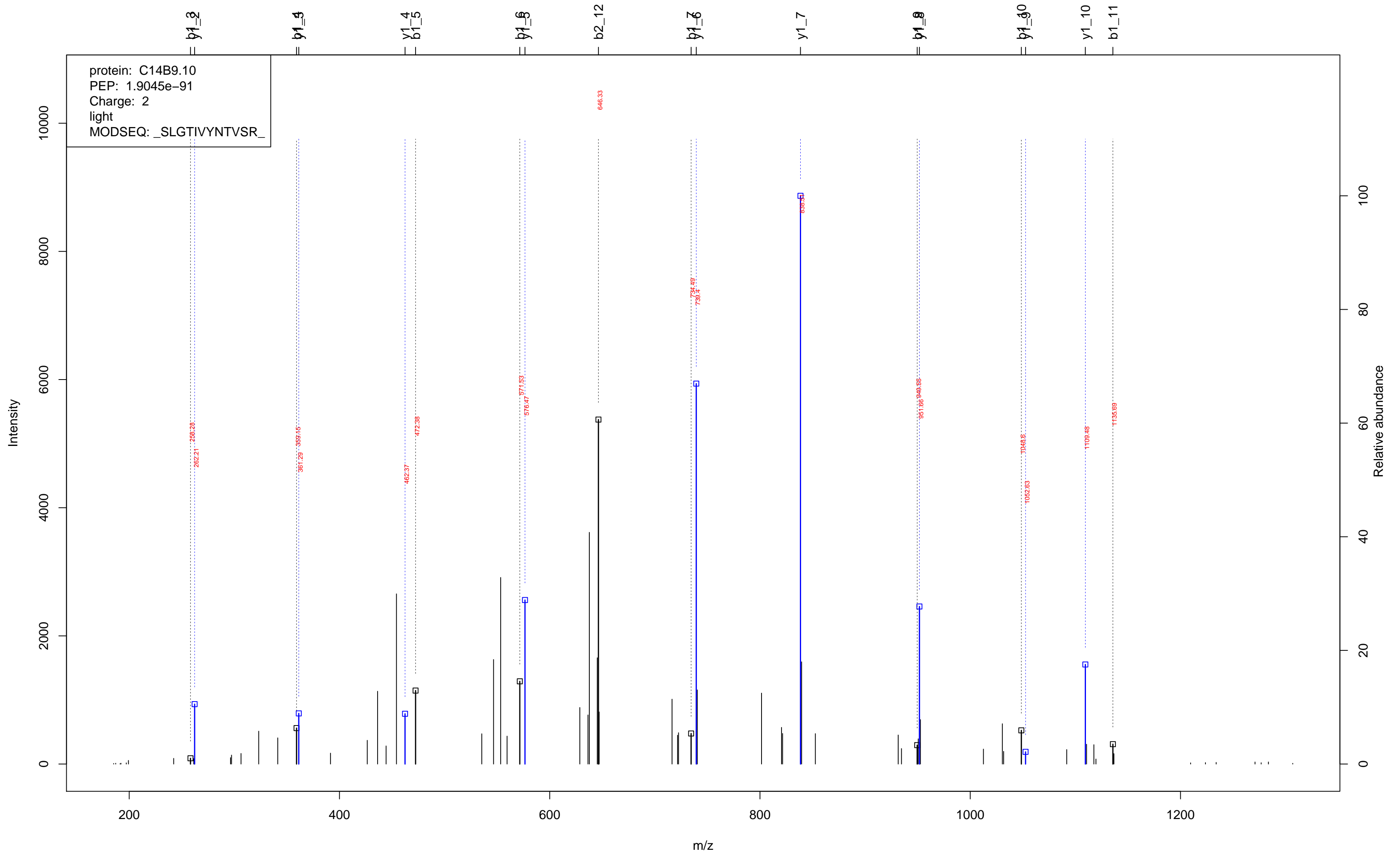


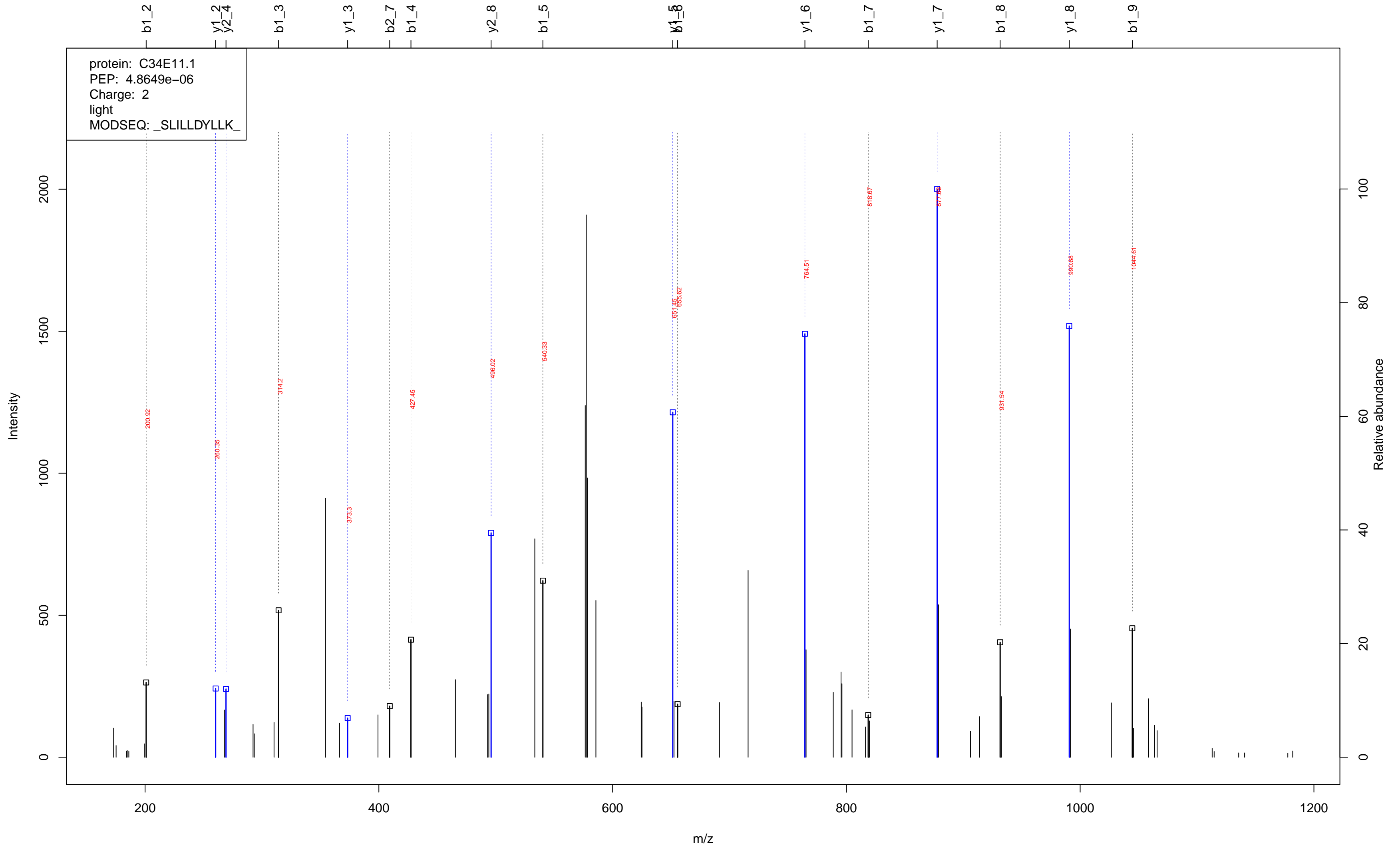
280

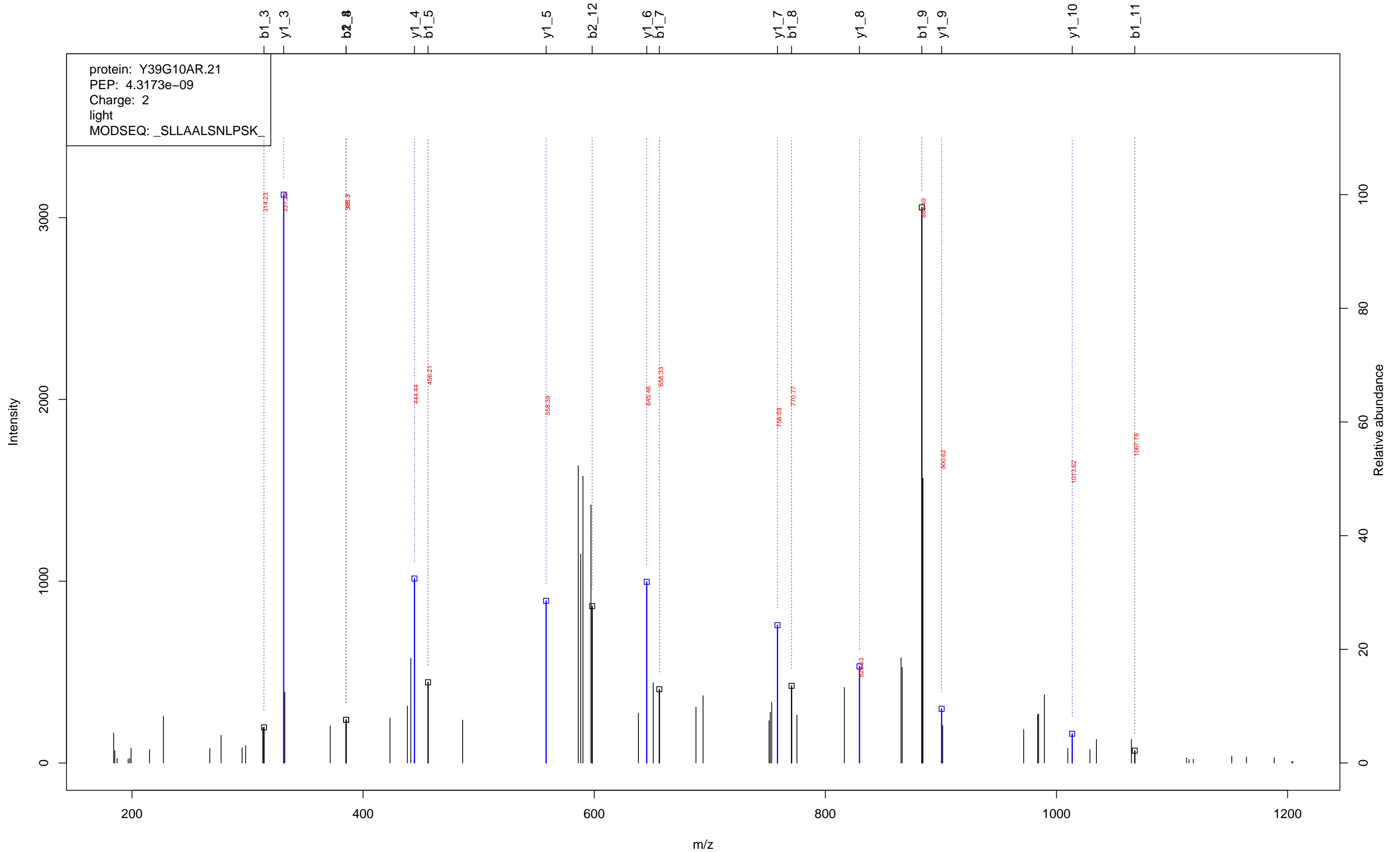
y2_5 y1_3 y1_4 y1_5 b1_7 b2_14 y1_7 y1_8 y1_9 y1_10 y1_12 y1_12 b1_13

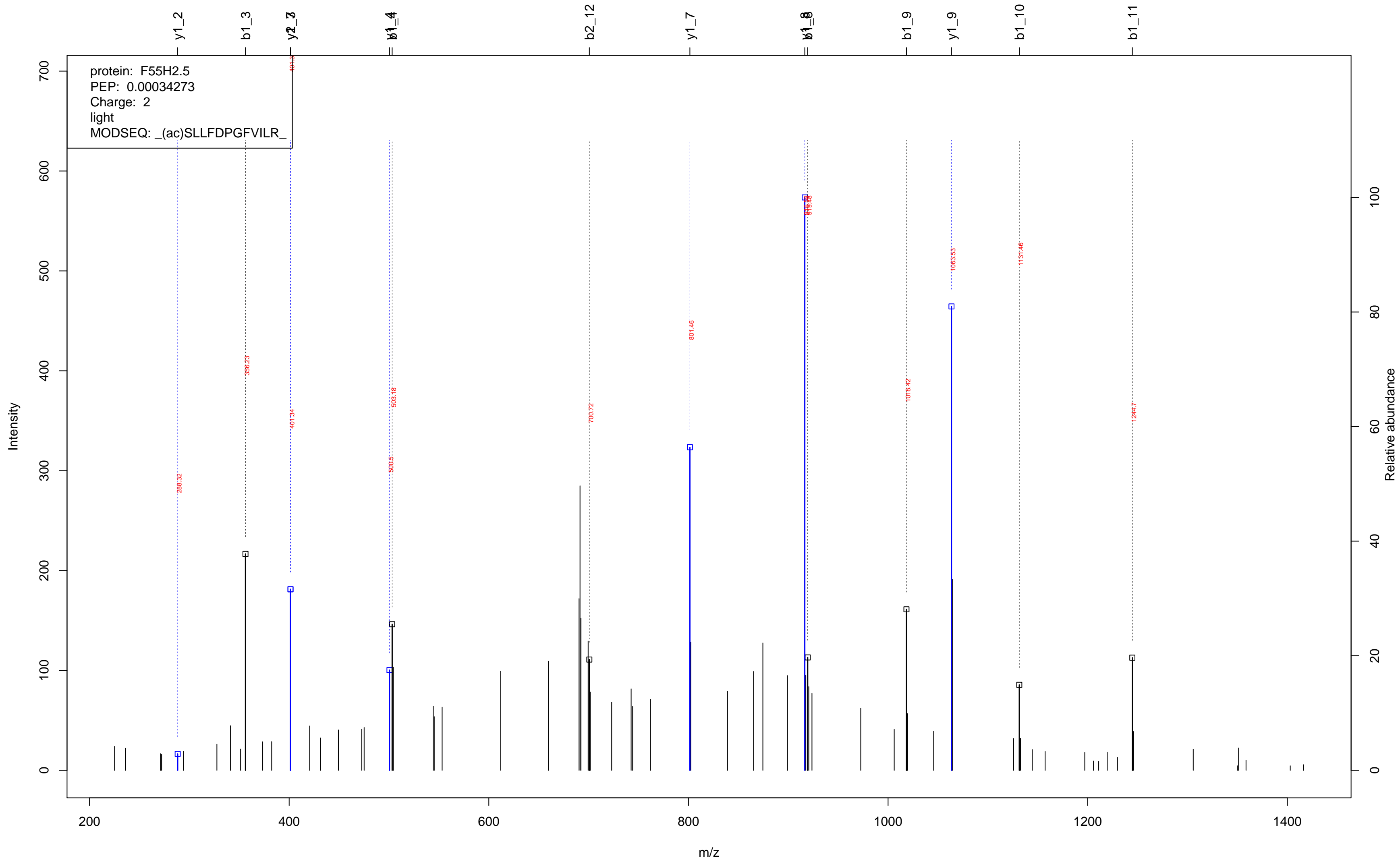
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PEP: 0.0051883
Charge: 2
light
MODSEQ: _SLGSLNIPLTLLK_

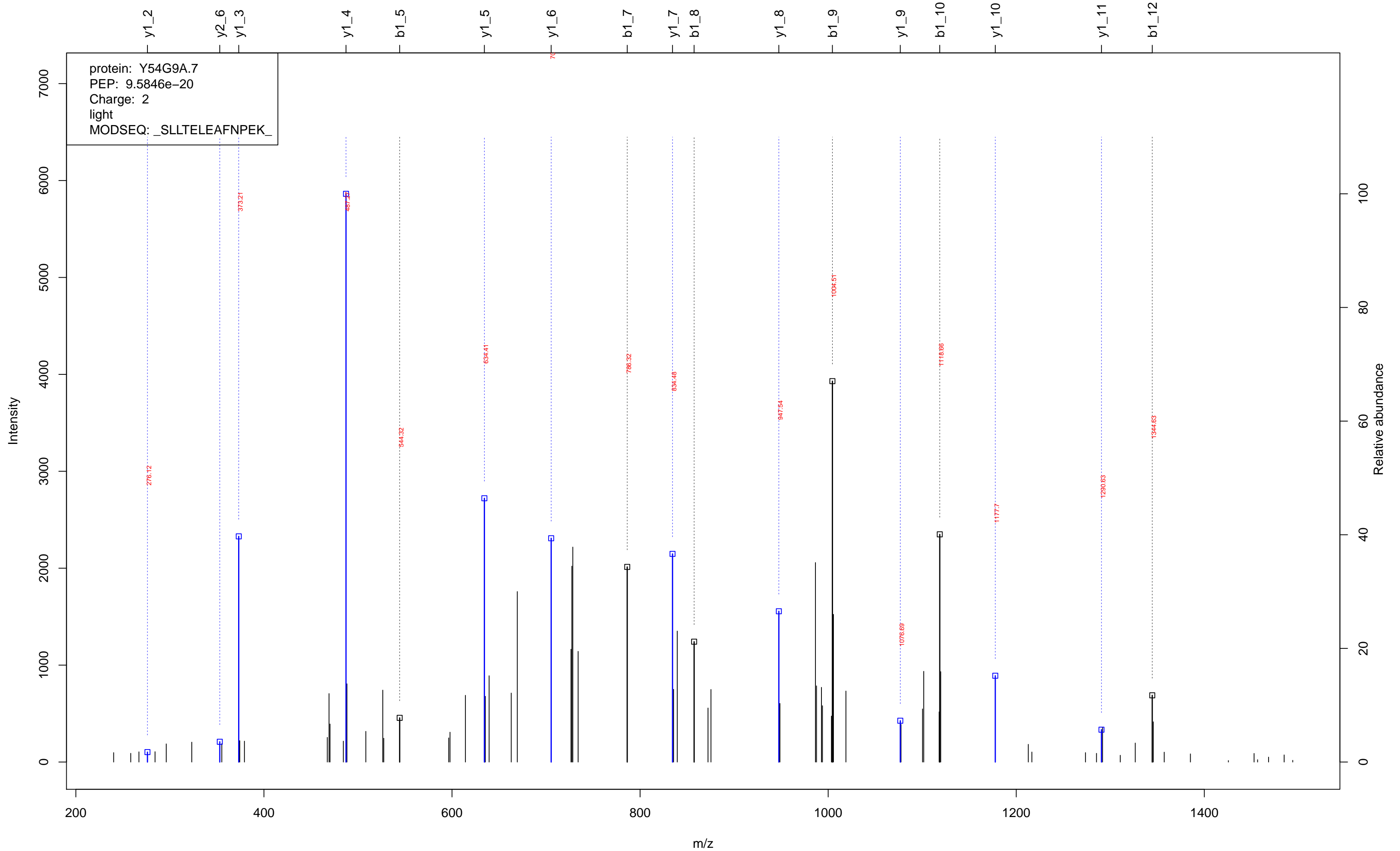


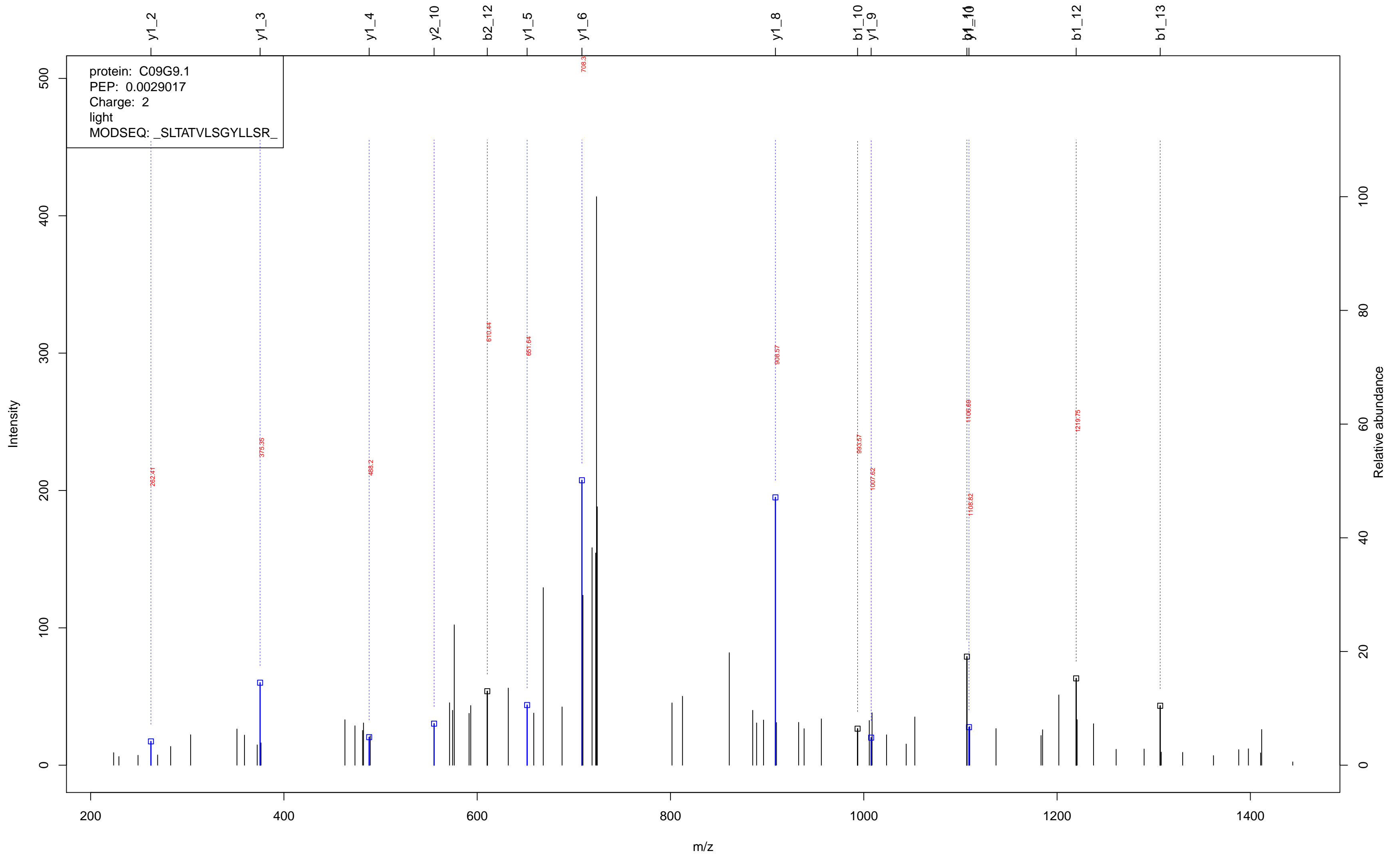


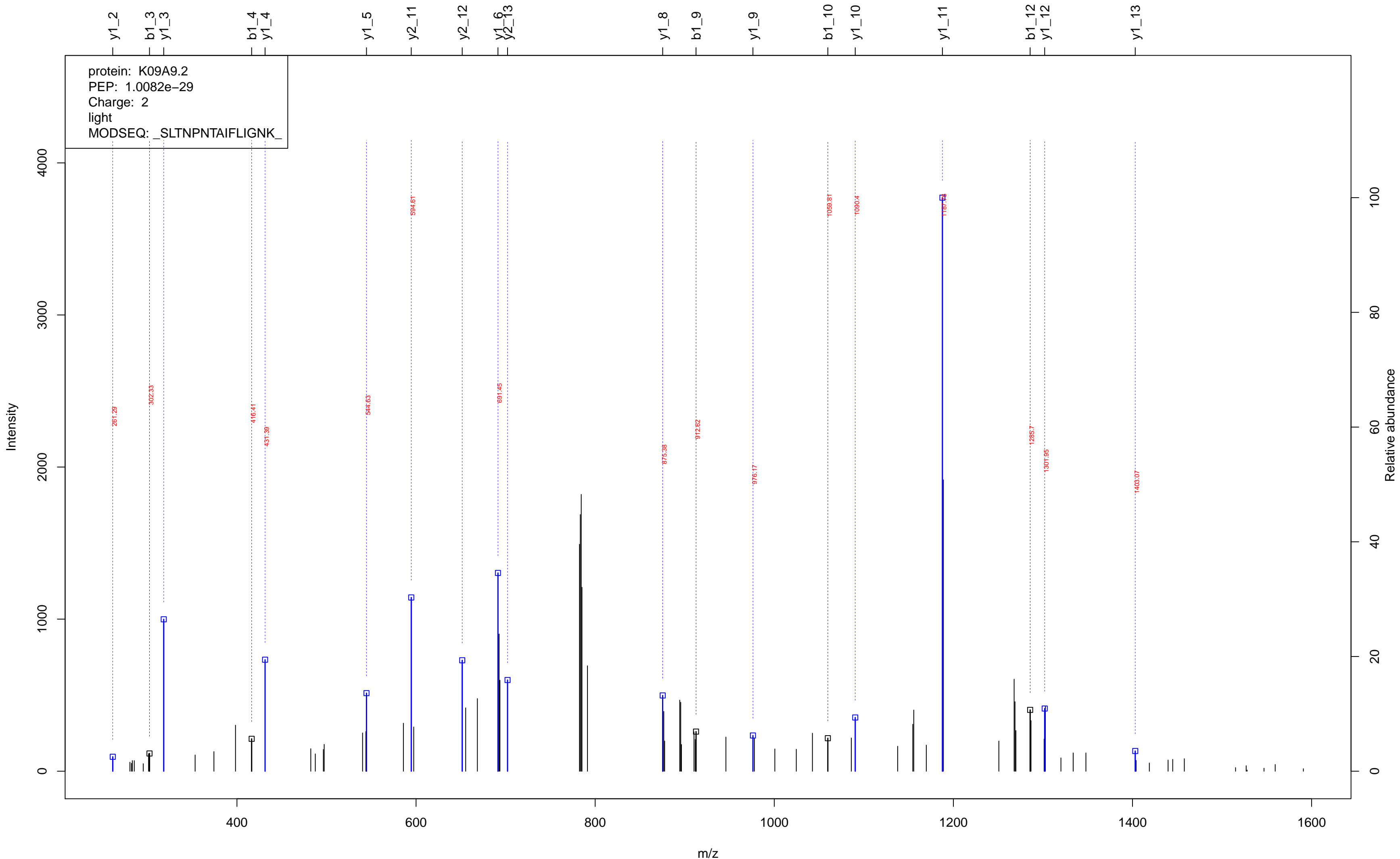


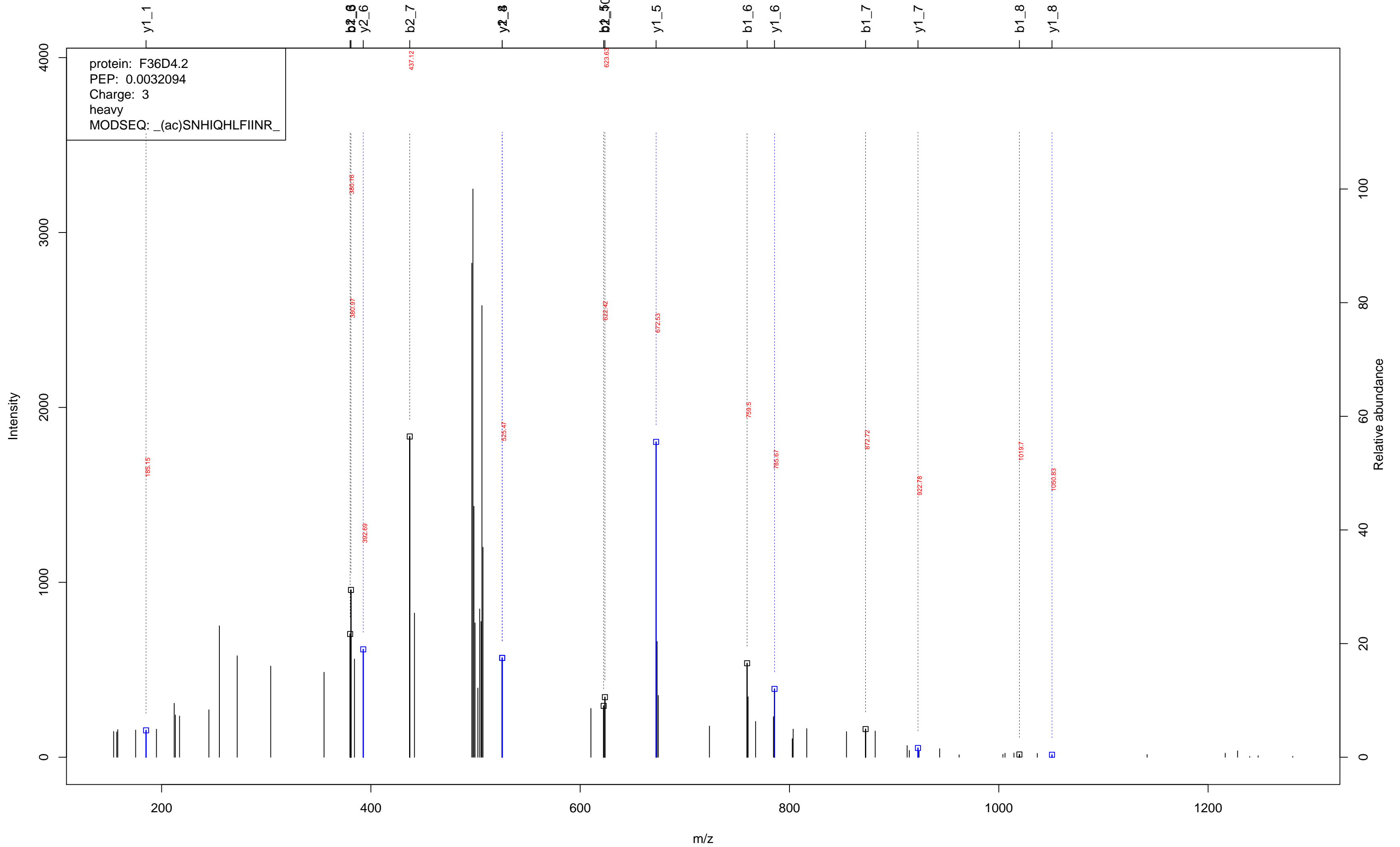


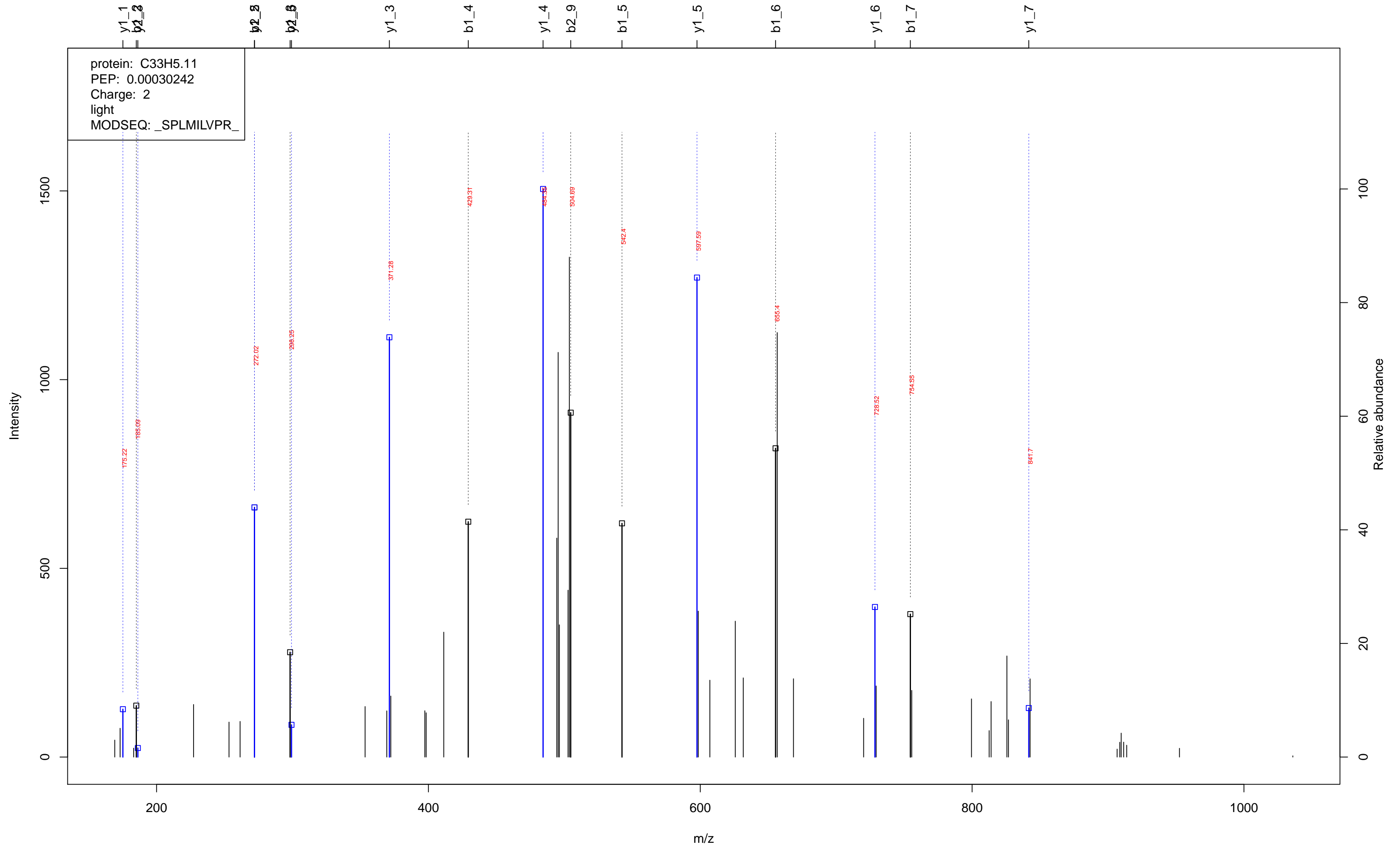


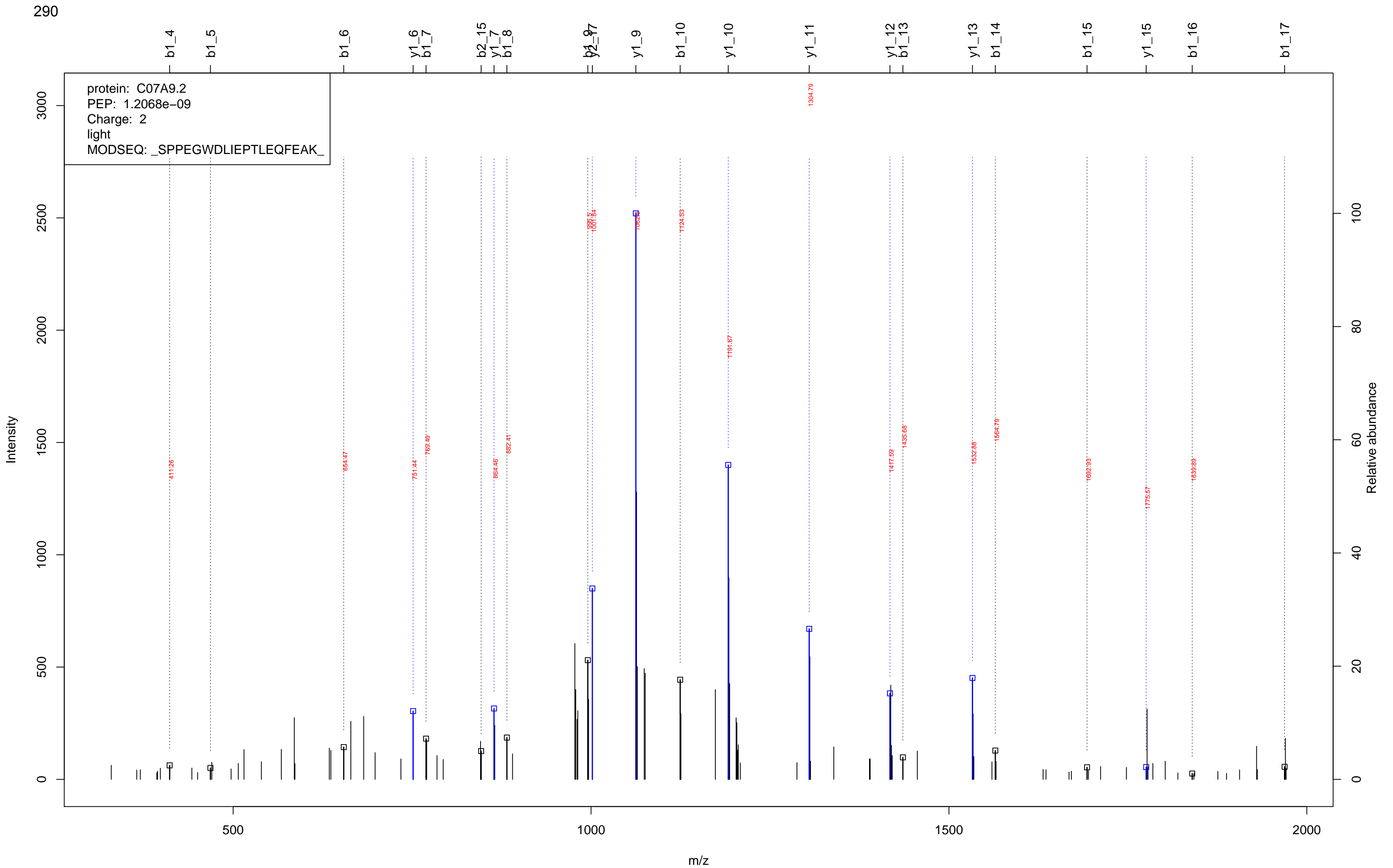


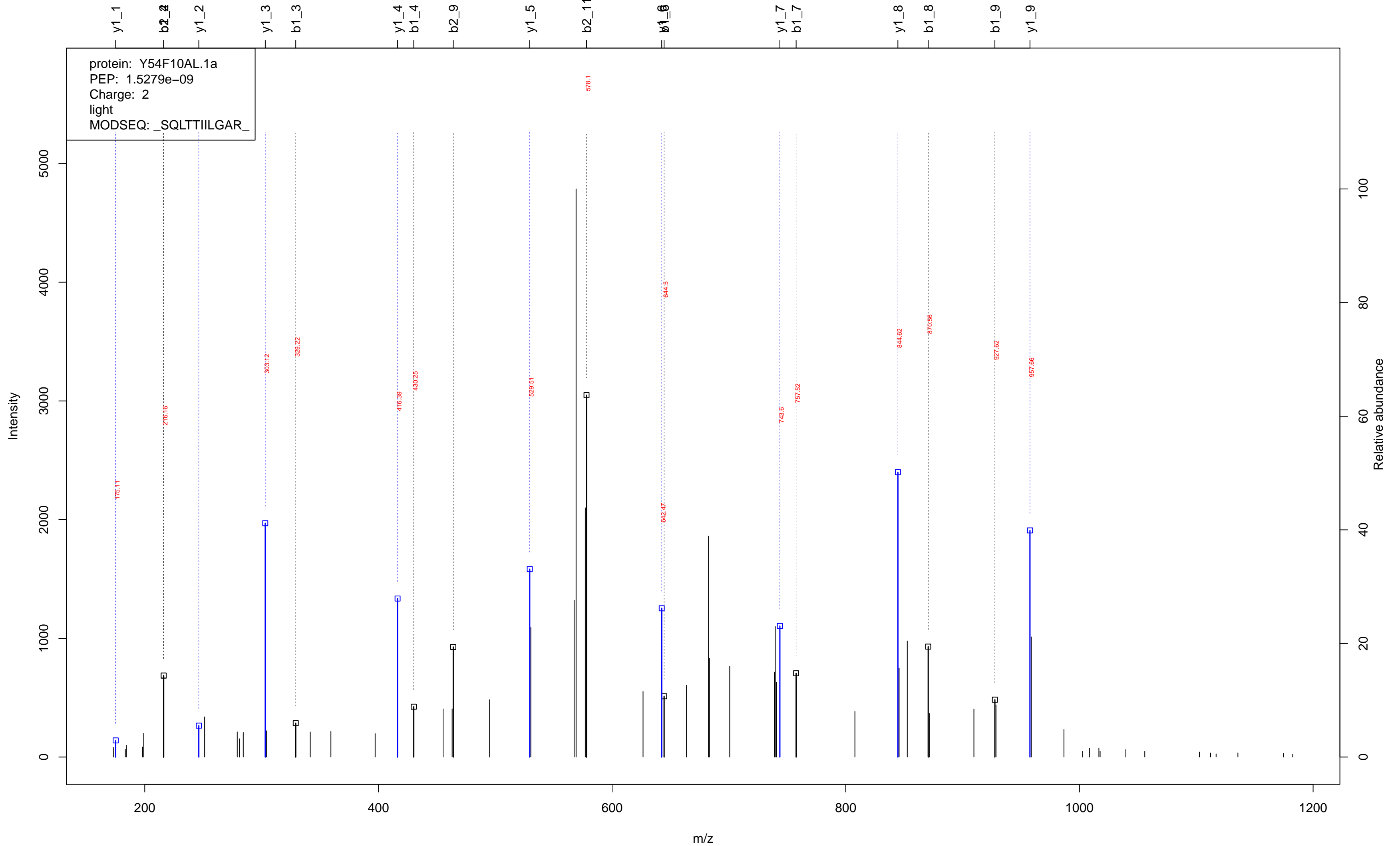


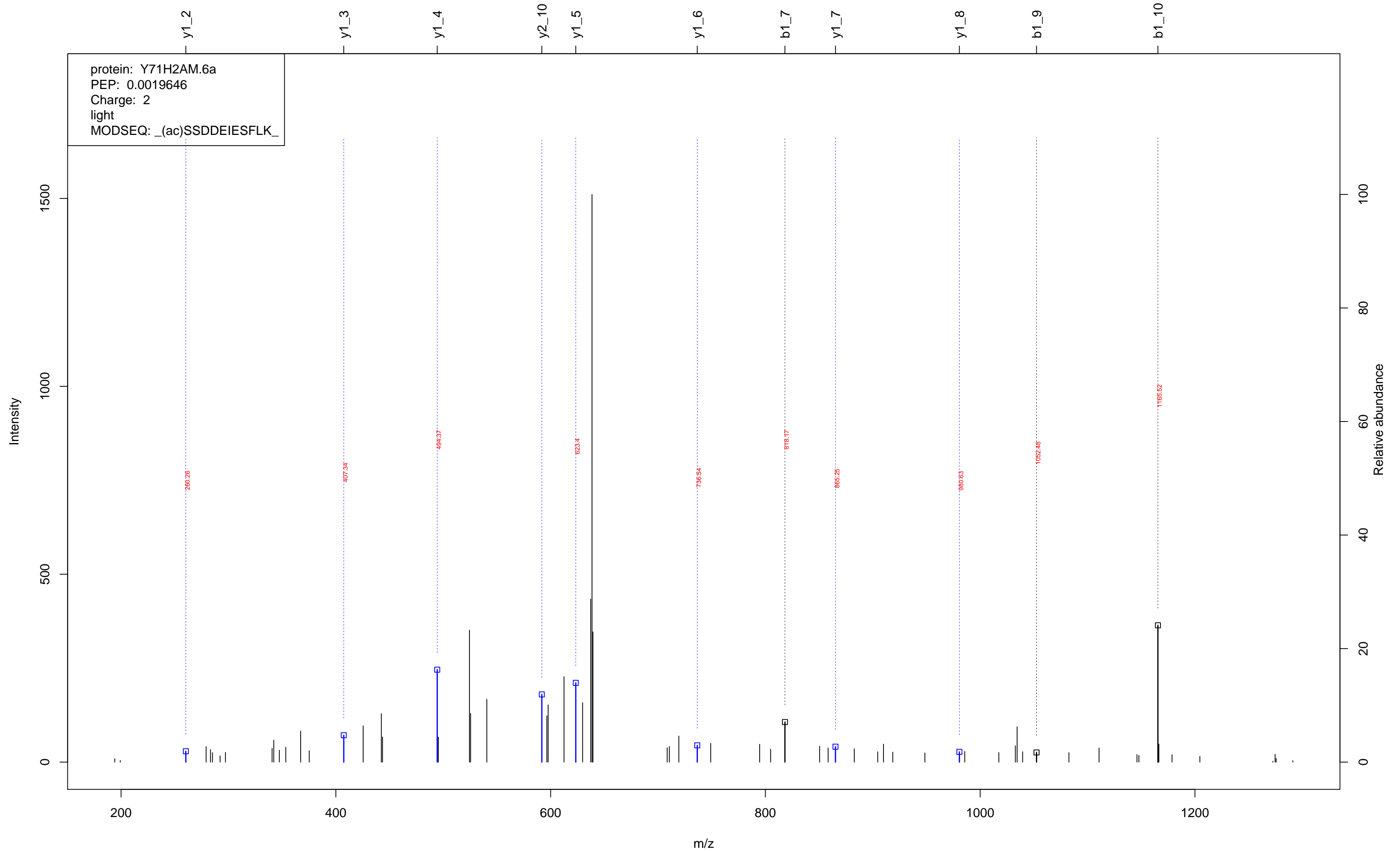


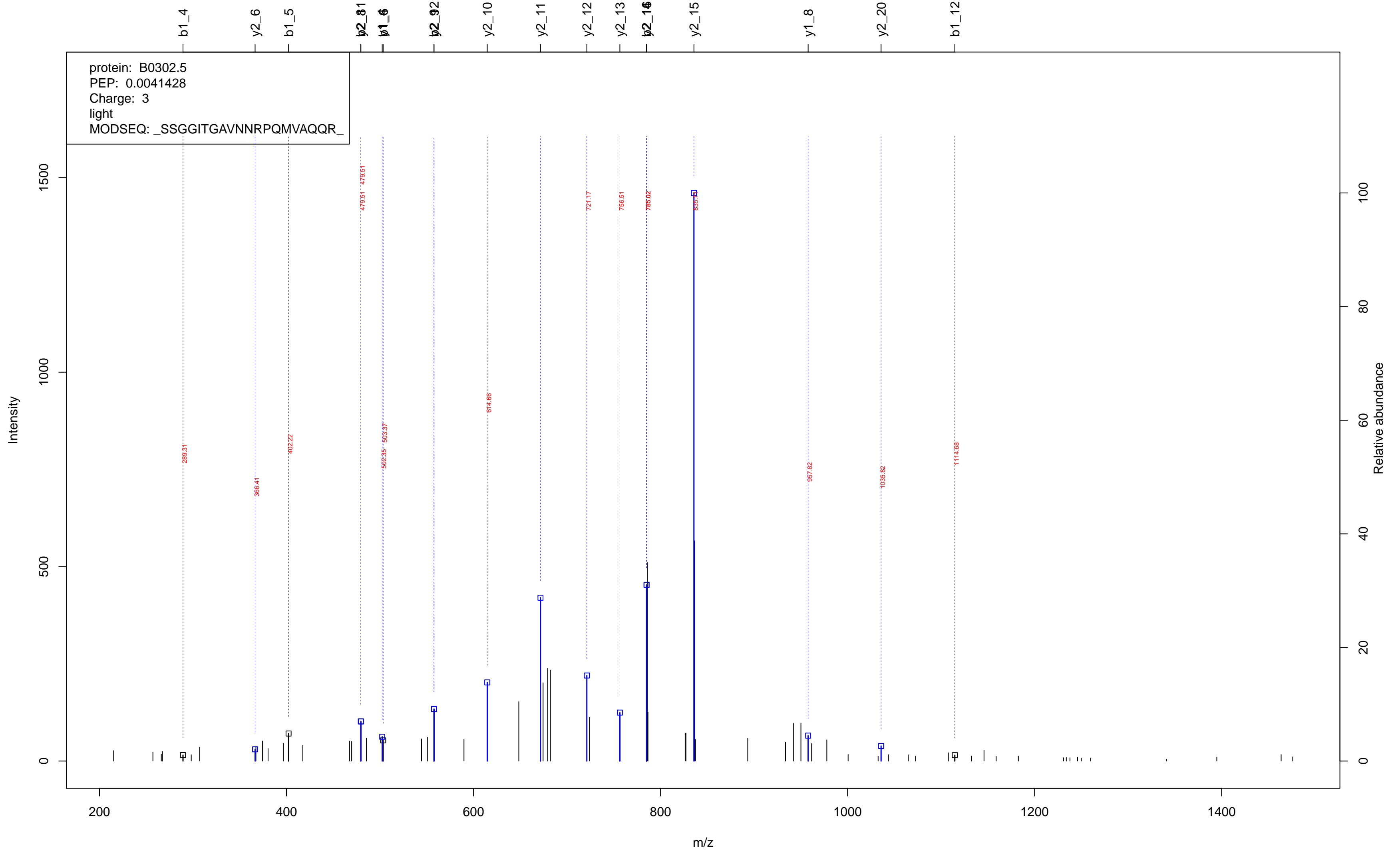






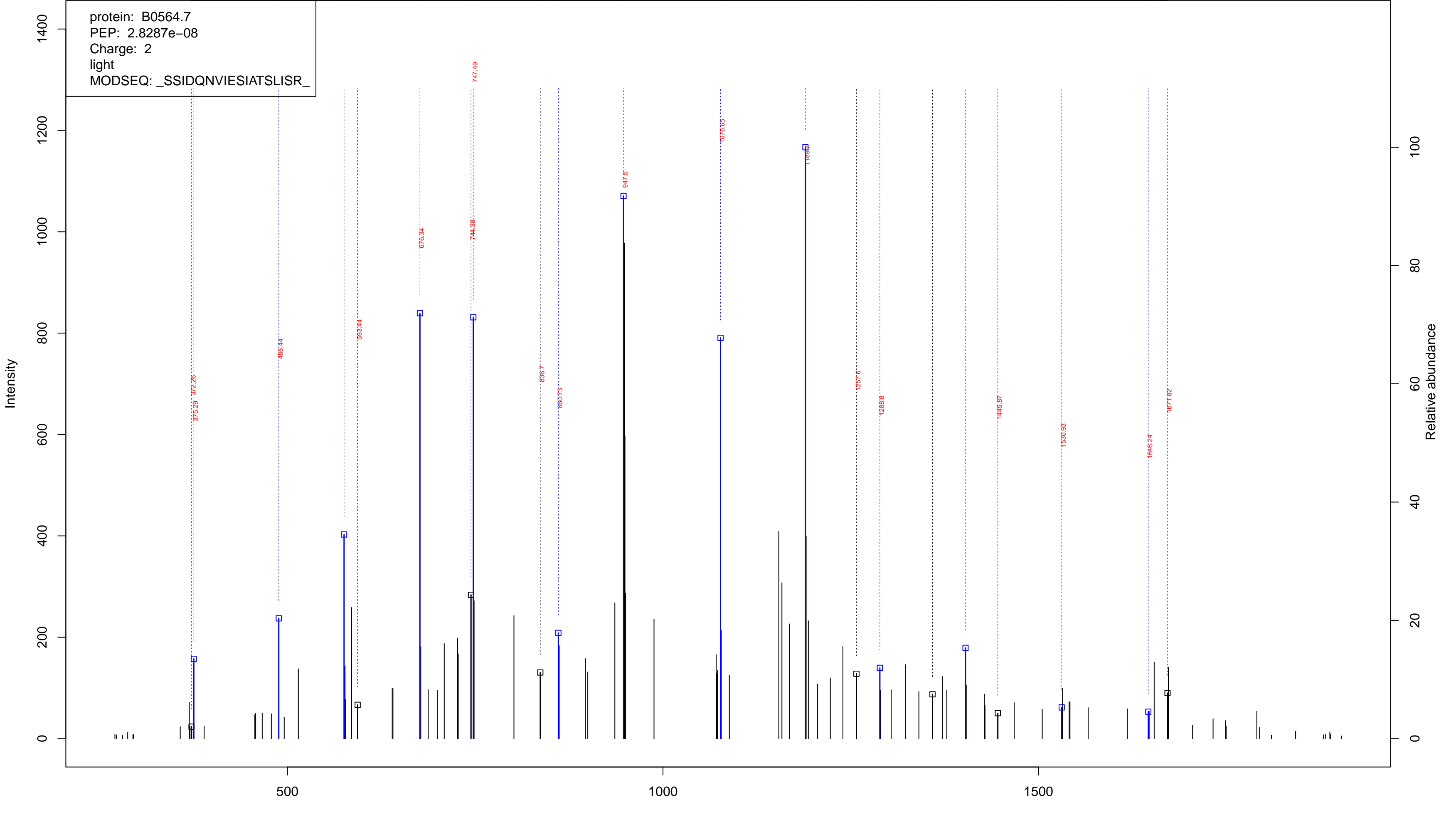


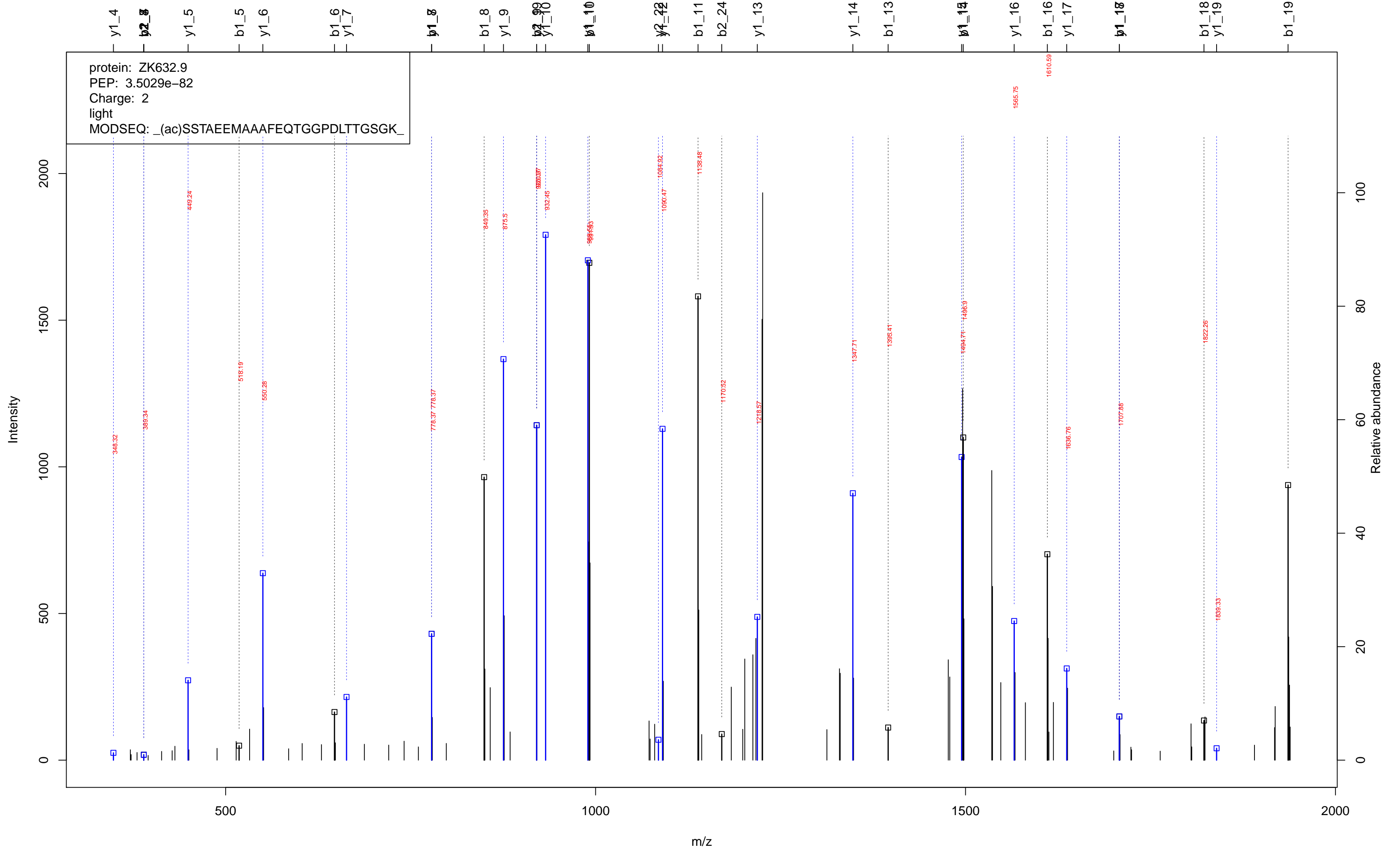


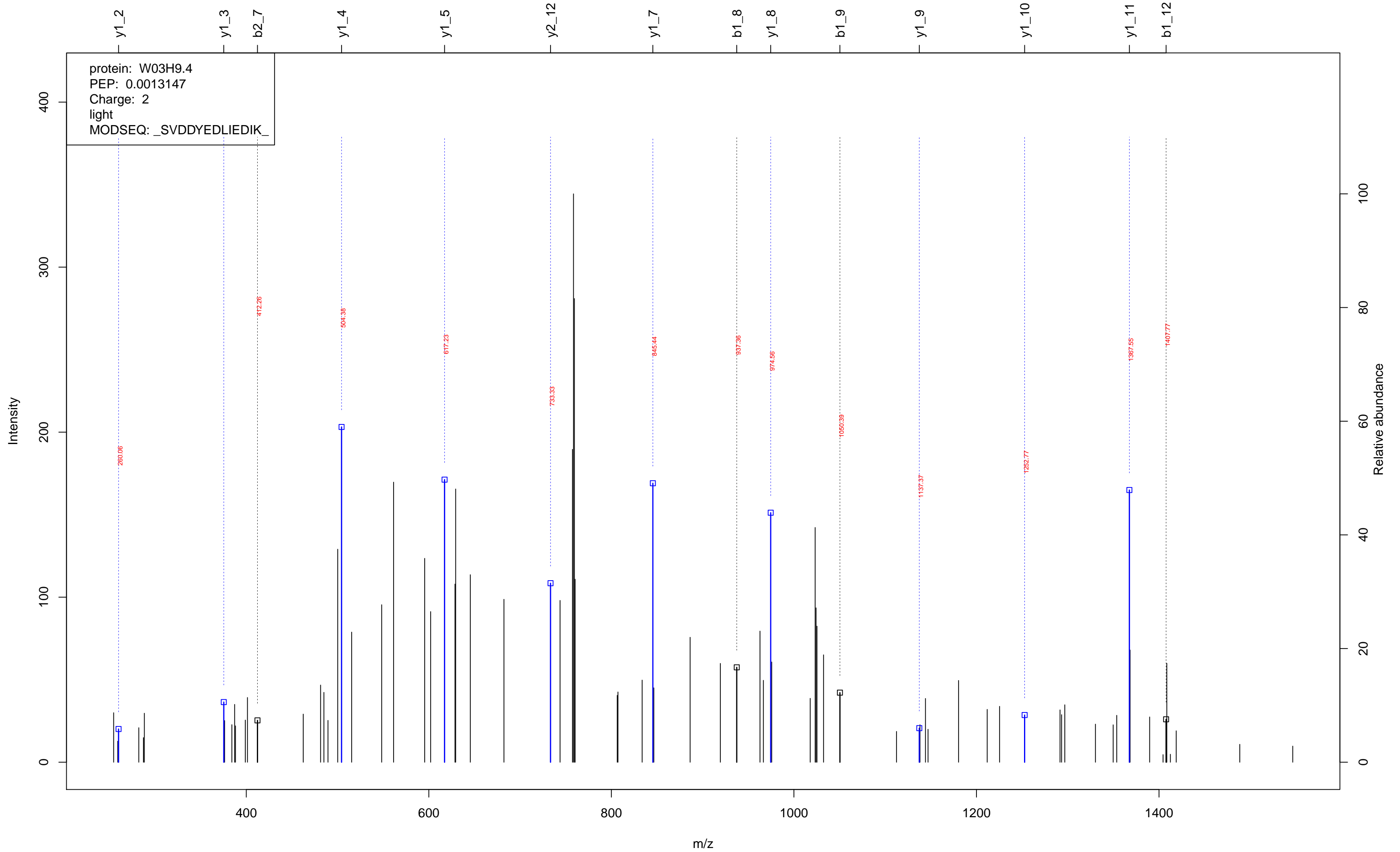


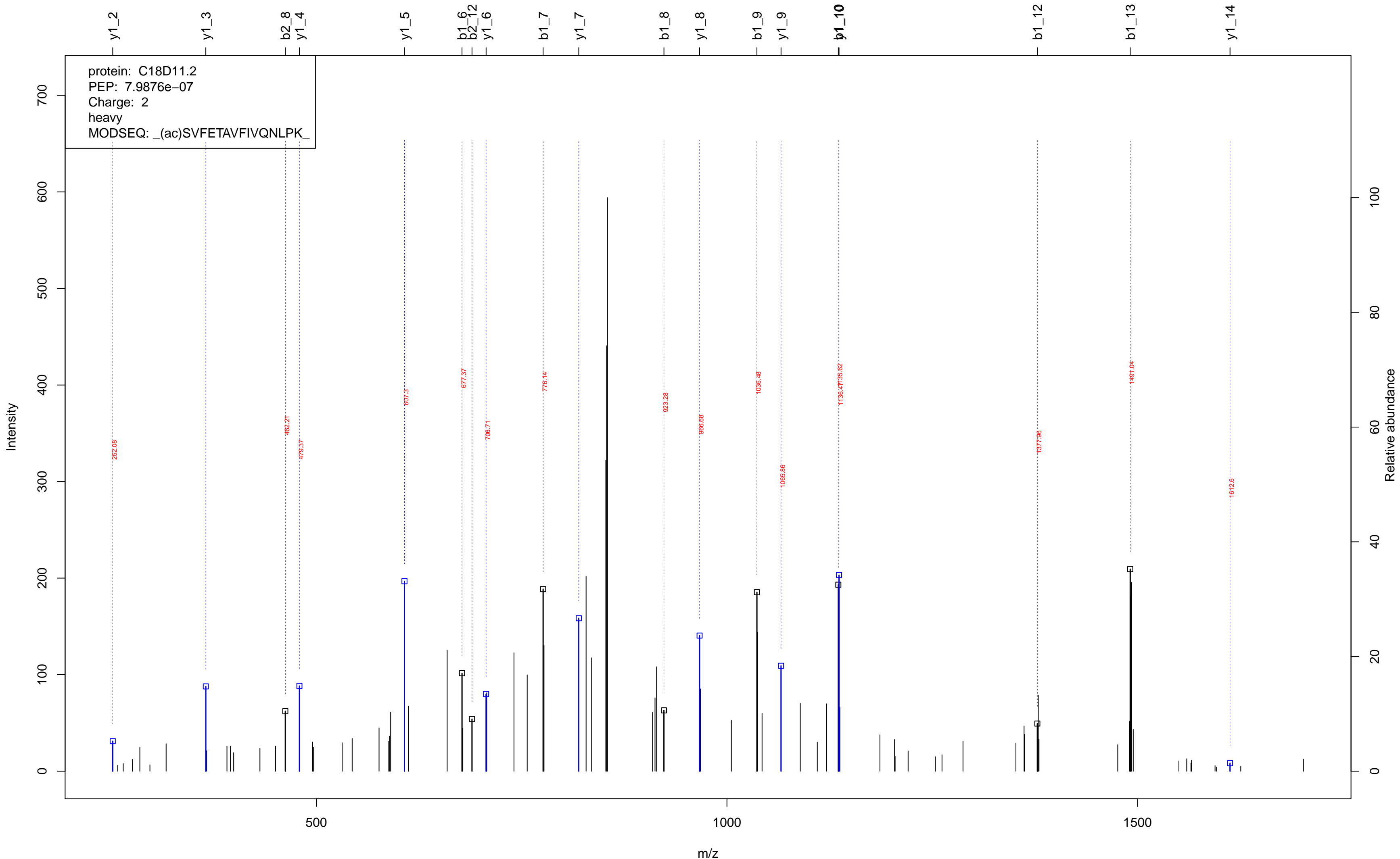
b2_3 y1_4 y1_5 b2_11 y1_6 b1_7 b2_16 y1_8 y1_9 y1_10 y1_11 b1_12 y1_12 b1_13 y1_13 b1_14 y1_14 y1_15 b1_16

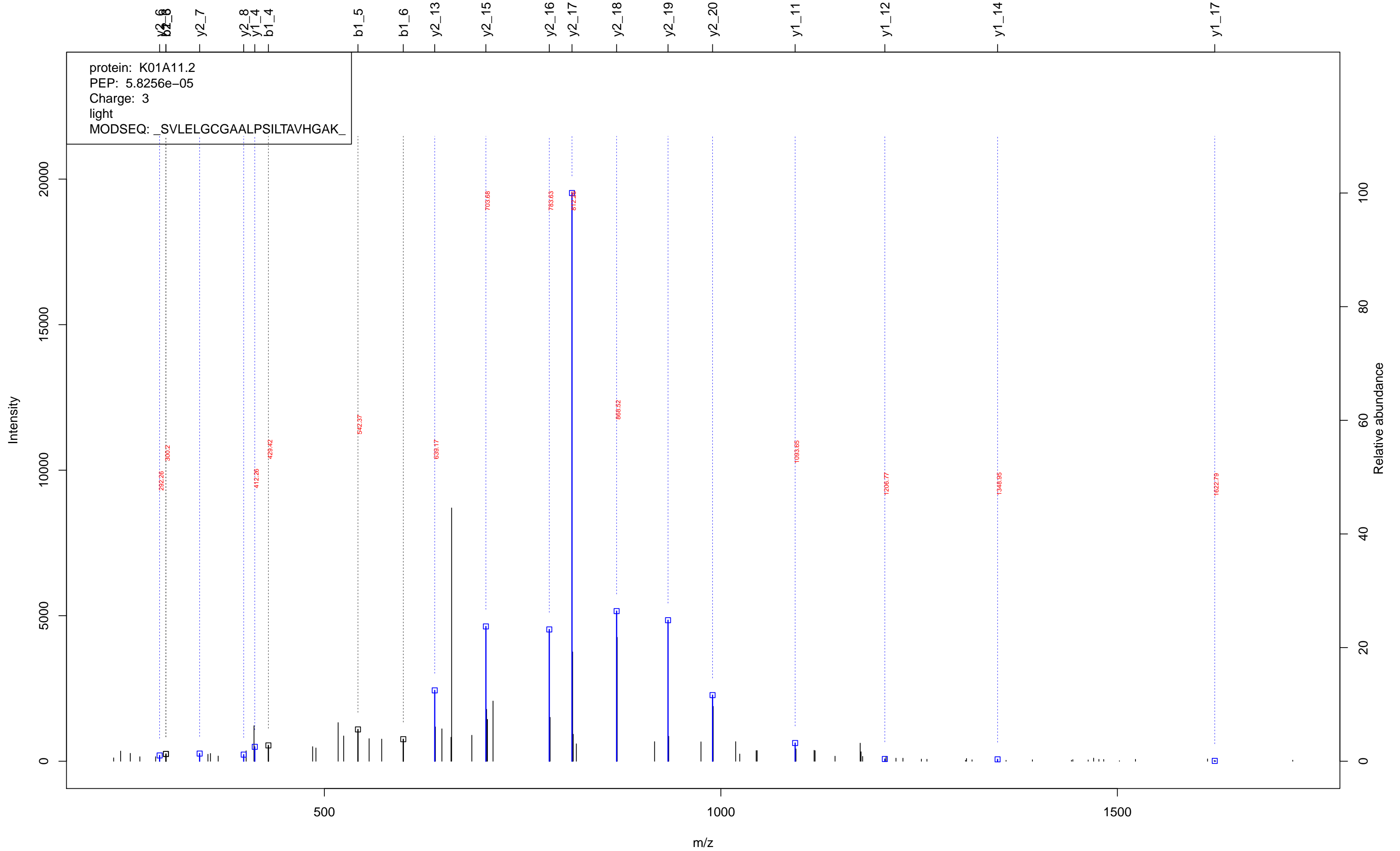
protein: B0564.7
PEP: 2.8287e-08
Charge: 2
light
MODSEQ: _SSIDQNVIESIATSLISR_

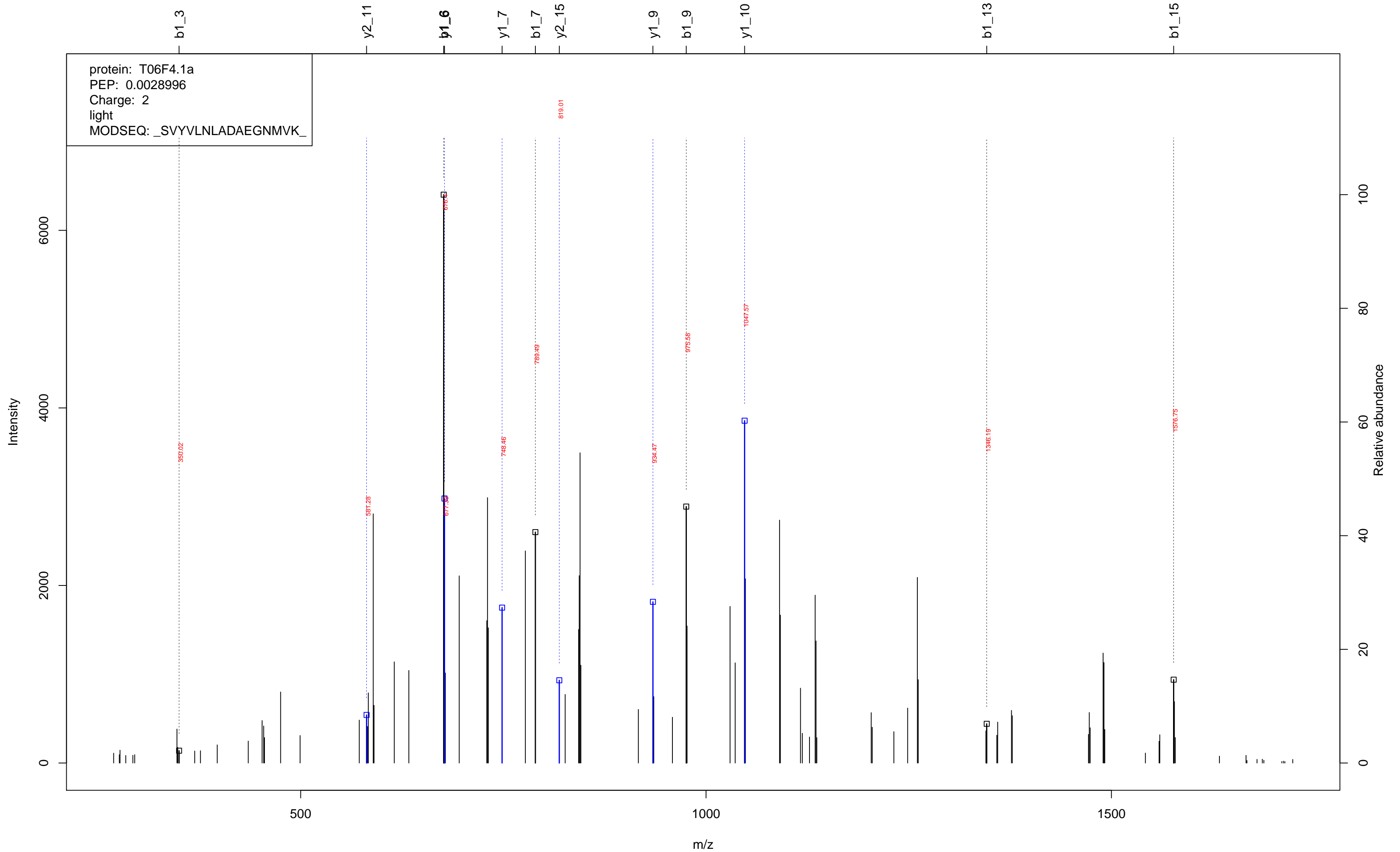


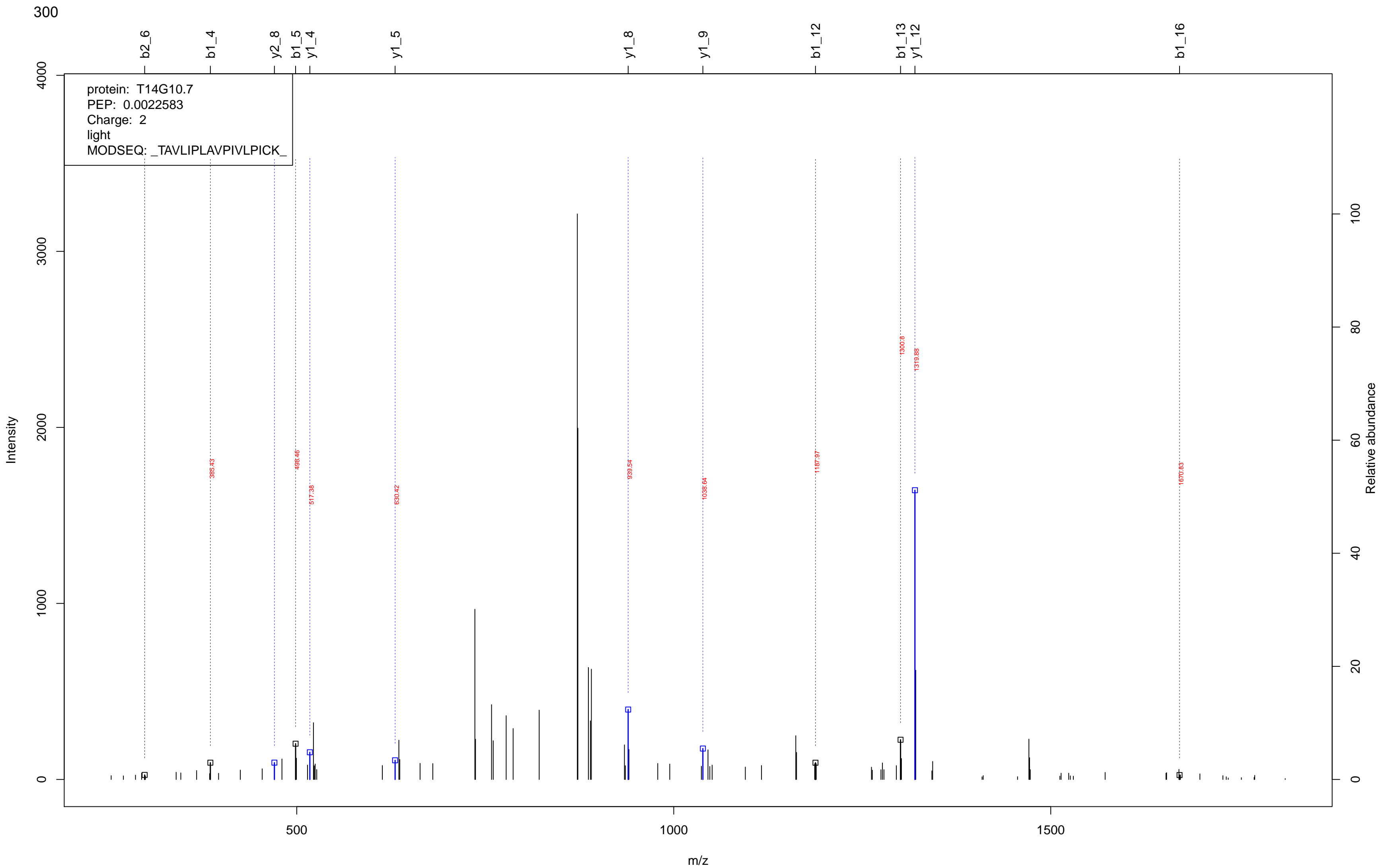


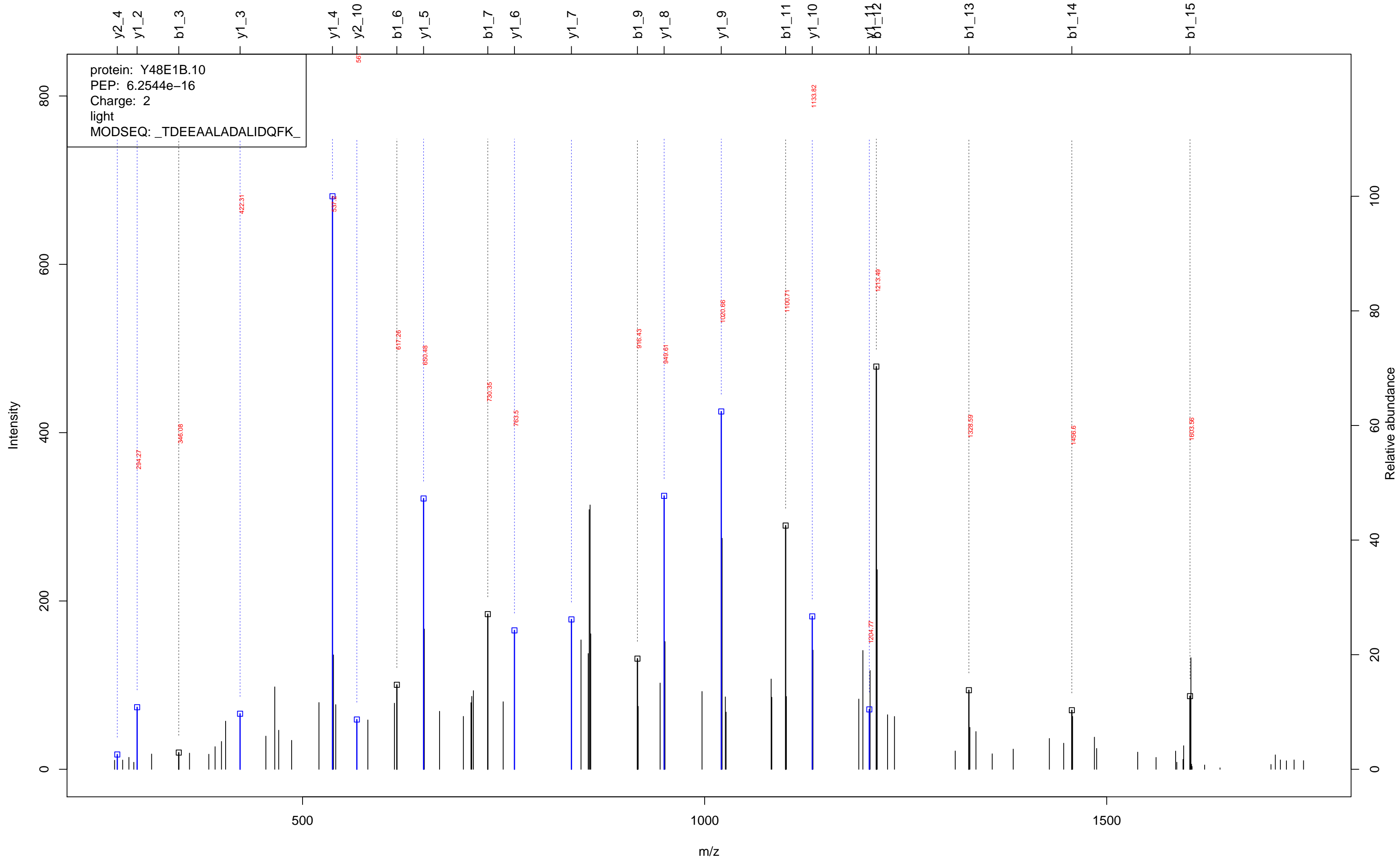


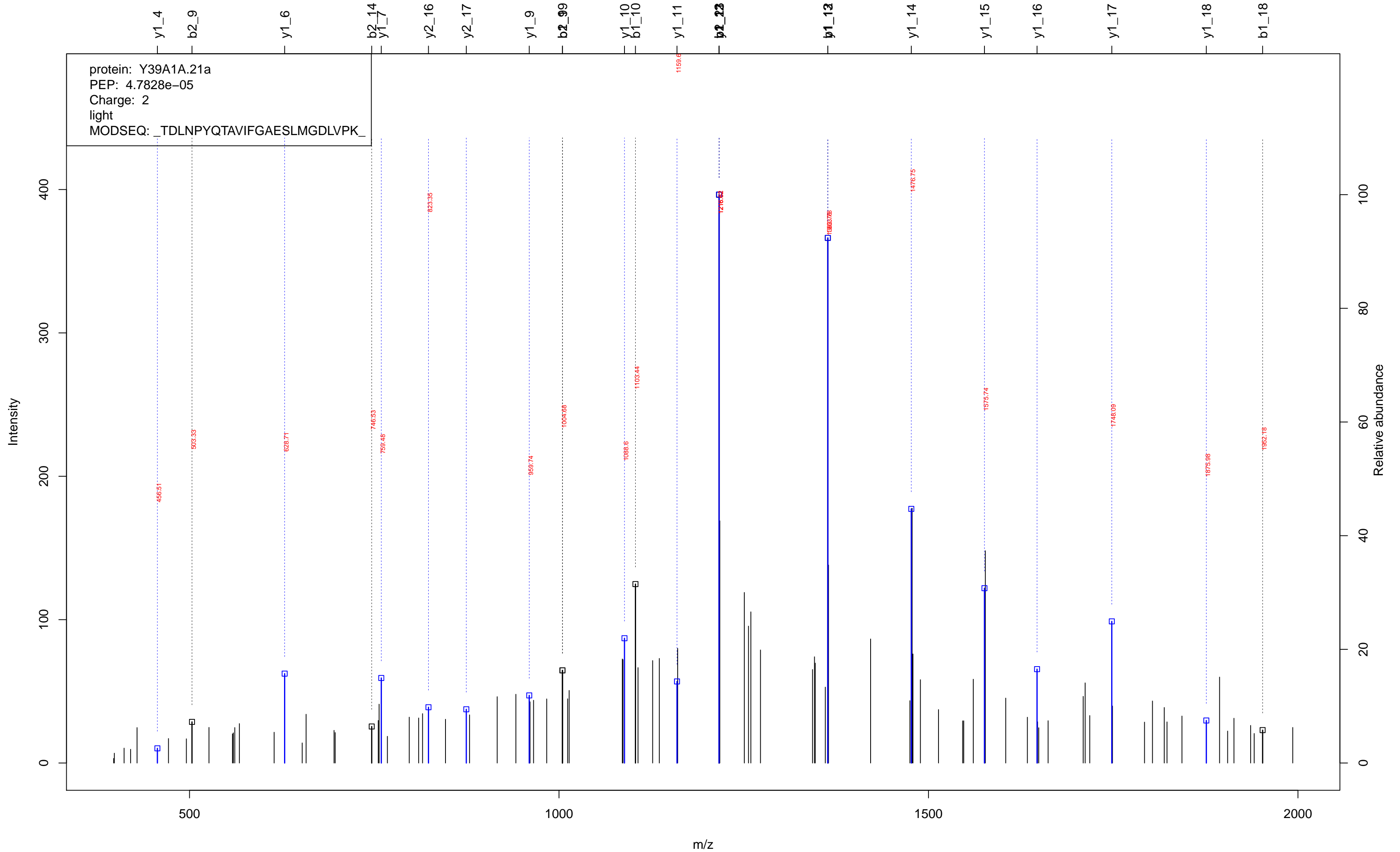


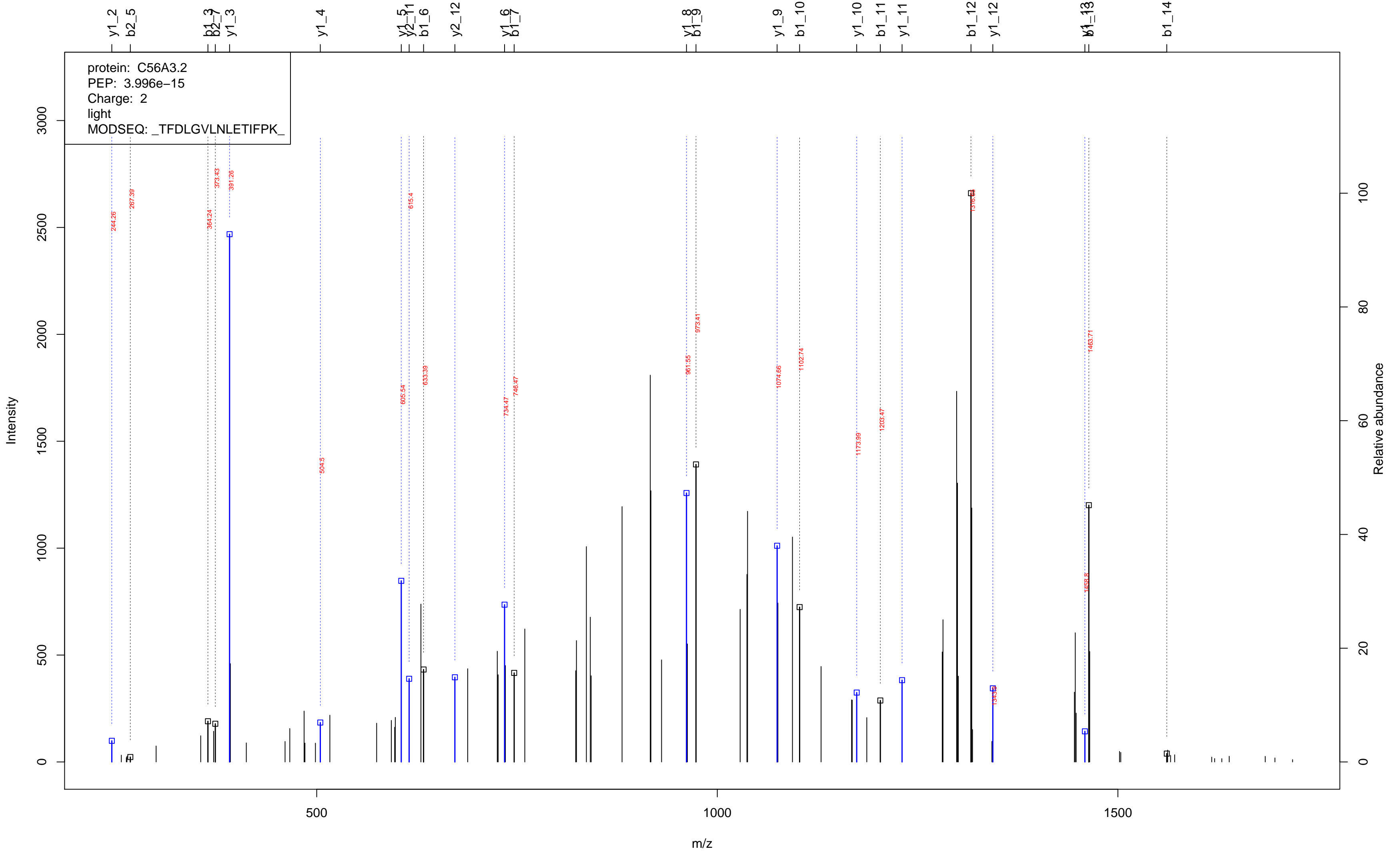


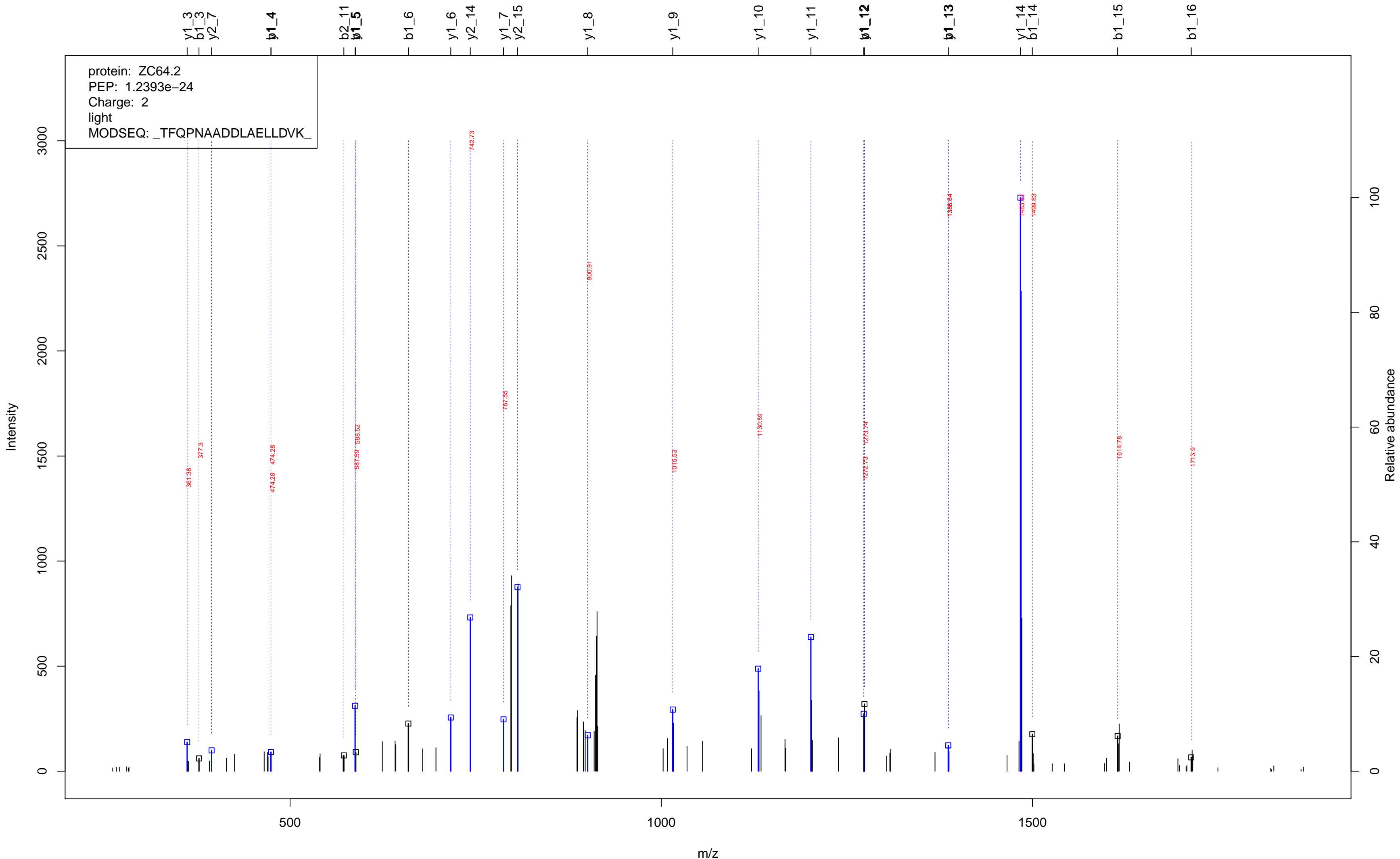






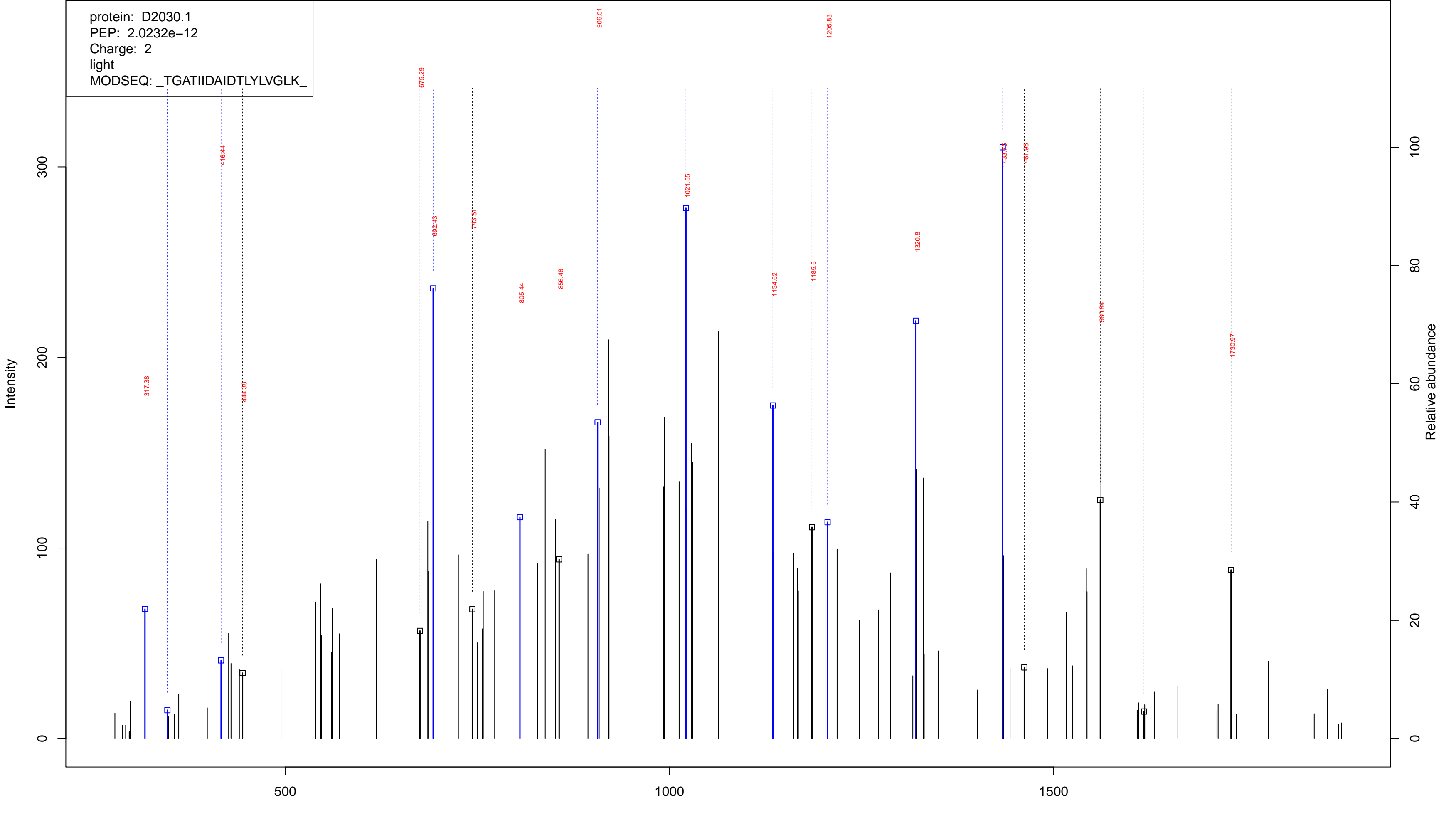


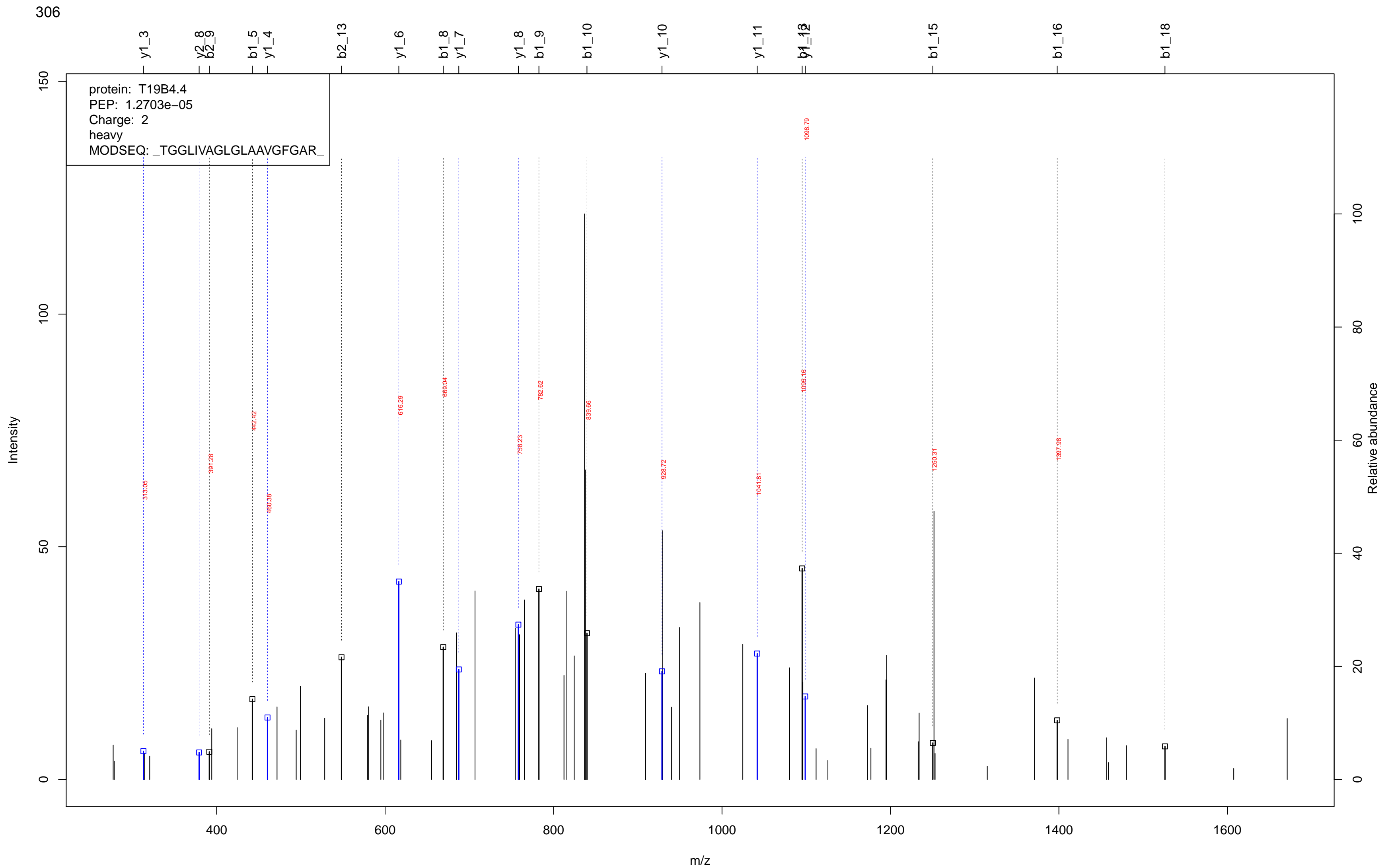


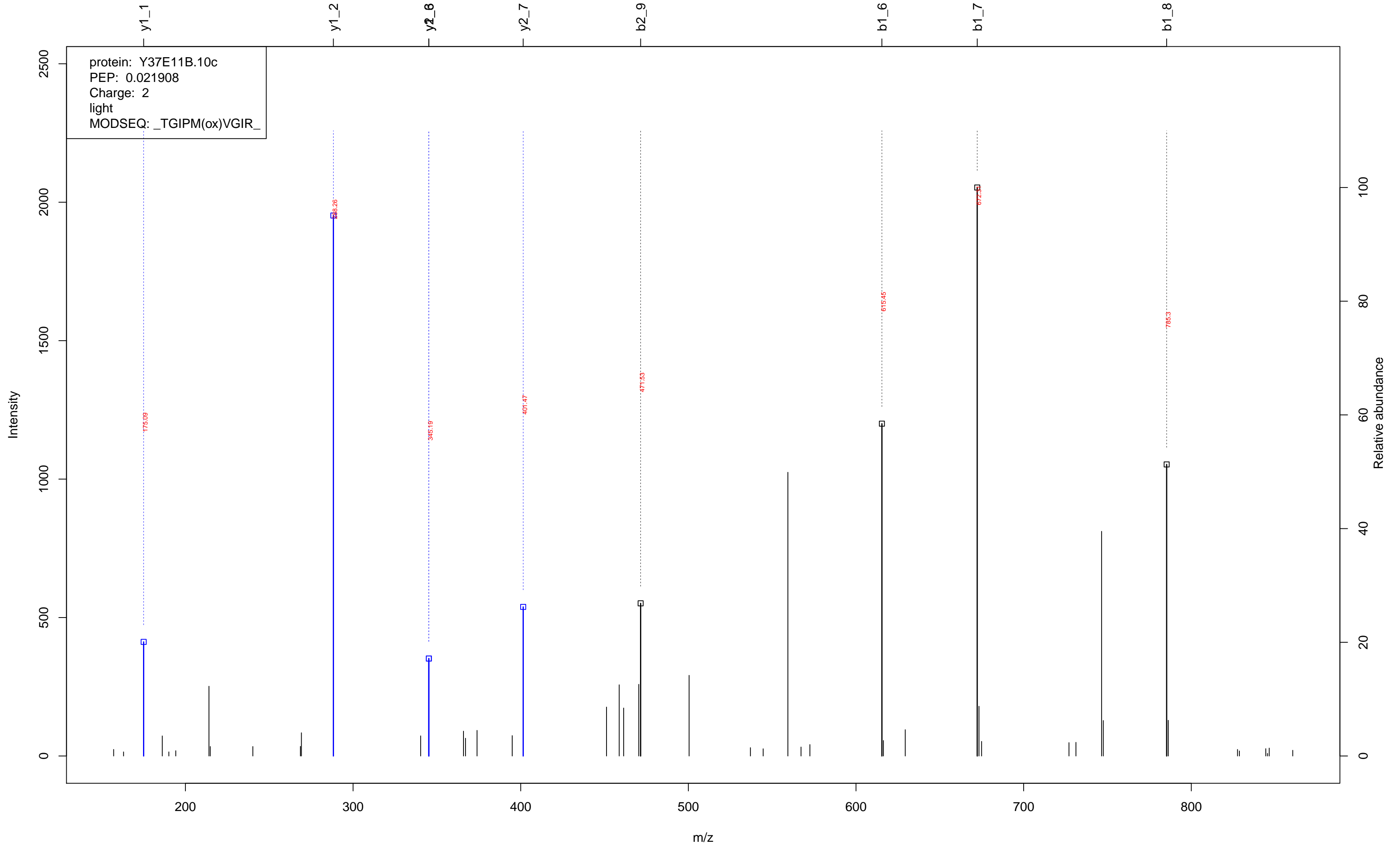


y1_3 y2_6 y1_4 b1_5 b2_13 y1_6 b1_8 y1_7 b1_9 y1_8 y1_9 y1_10 b1_12 y1_11 y1_12 y1_13 b1_14 b1_15 b1_16 b1_17

protein: D2030.1
PEP: 2.0232e-12
Charge: 2
light
MODSEQ: _TGATIIDAIDTLYLVGLK_

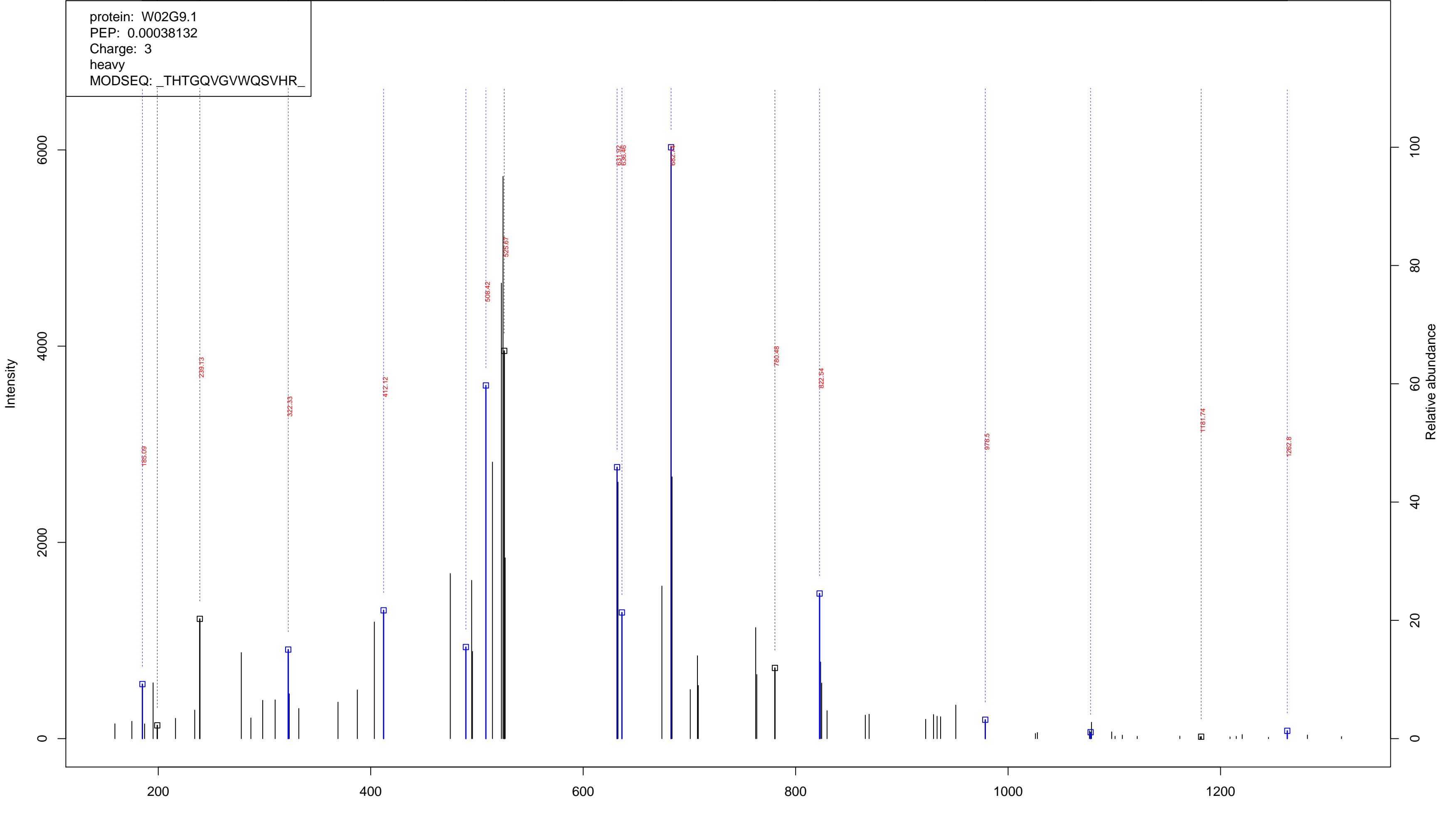


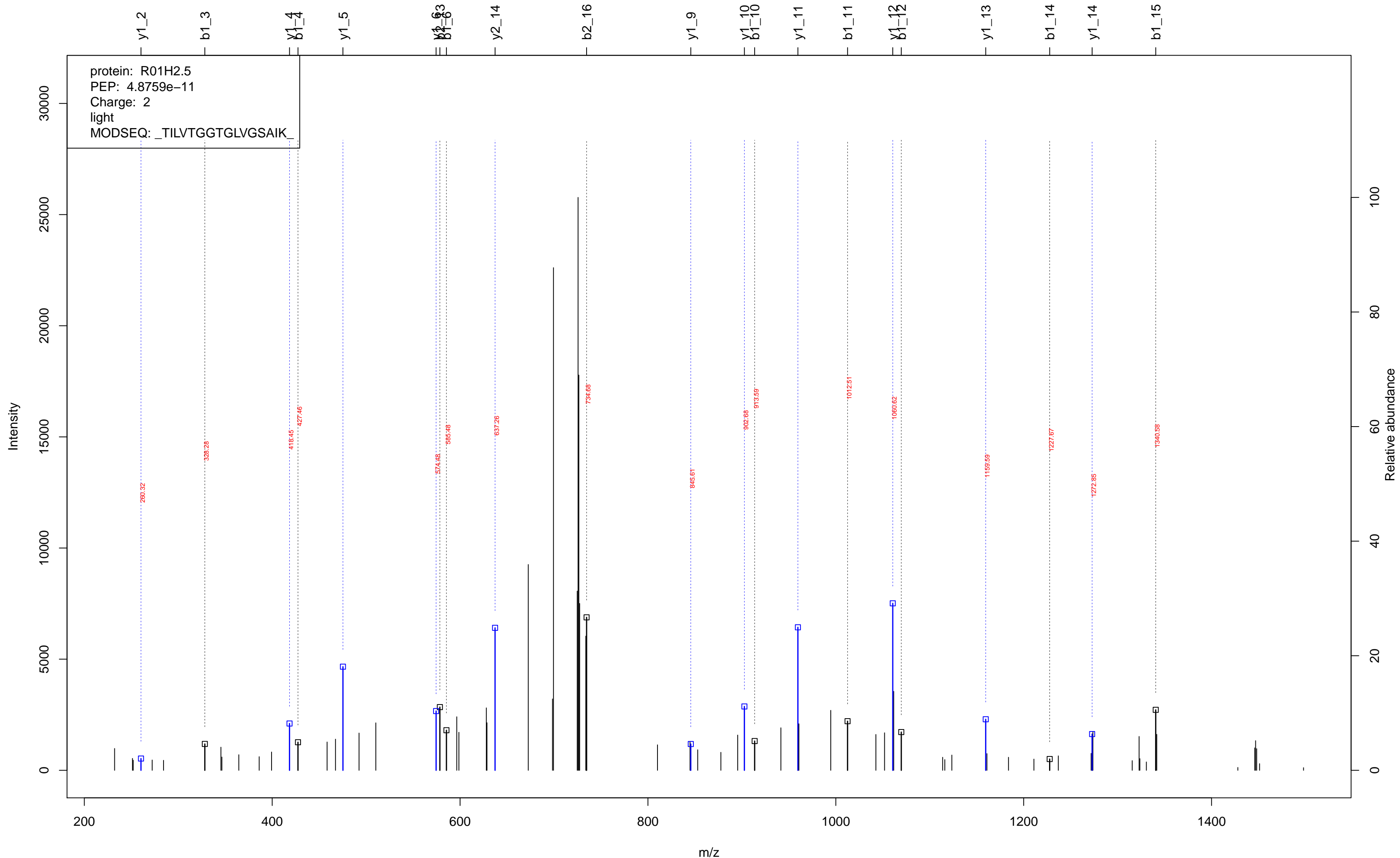


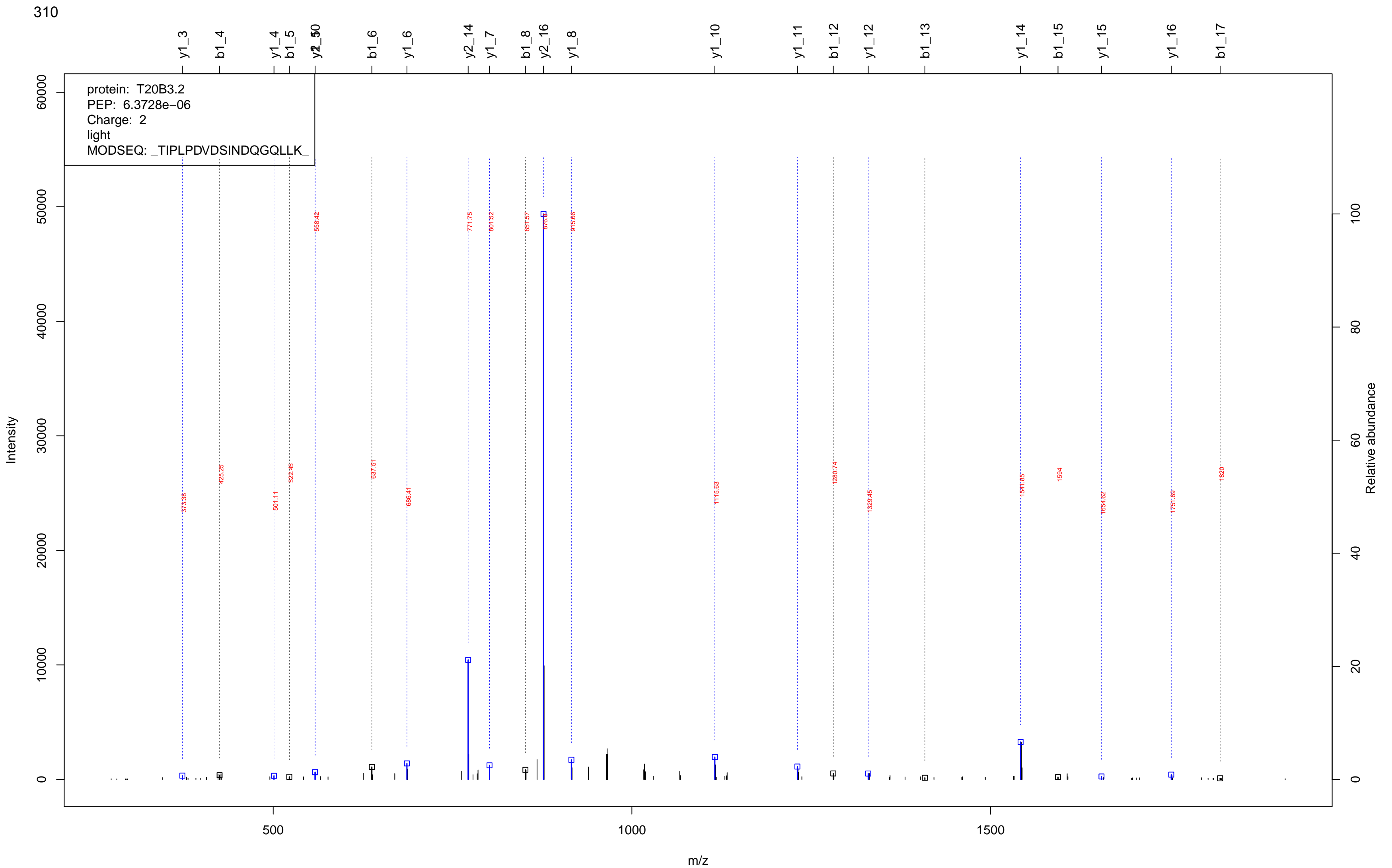


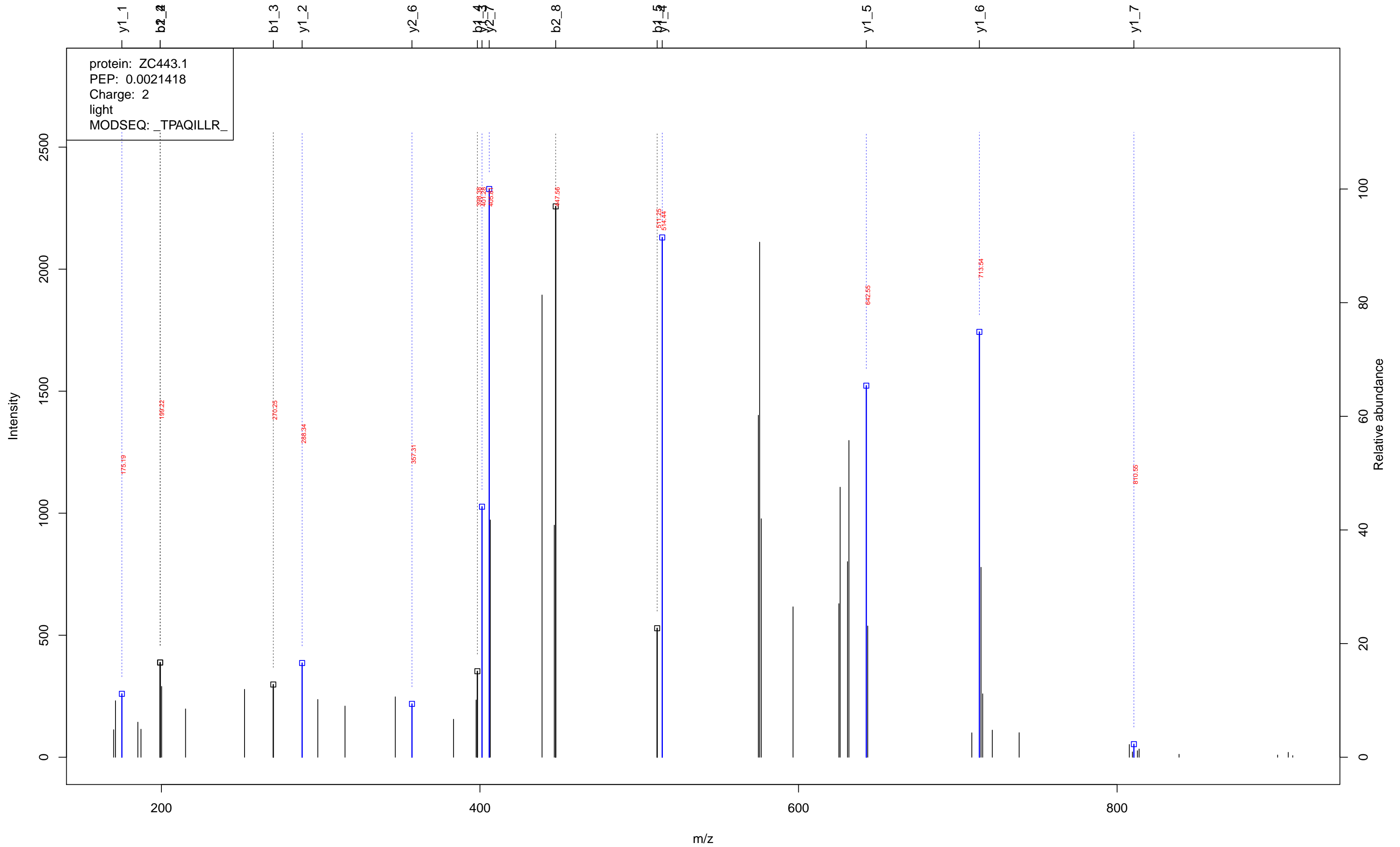
y1_1
b2_4
b1_2
y1_2
y2_6
y2_8
y1_4
b1_5
y2_11
y2_12
b1_8
y1_6
y1_8
y1_9
b1_11
y1_11

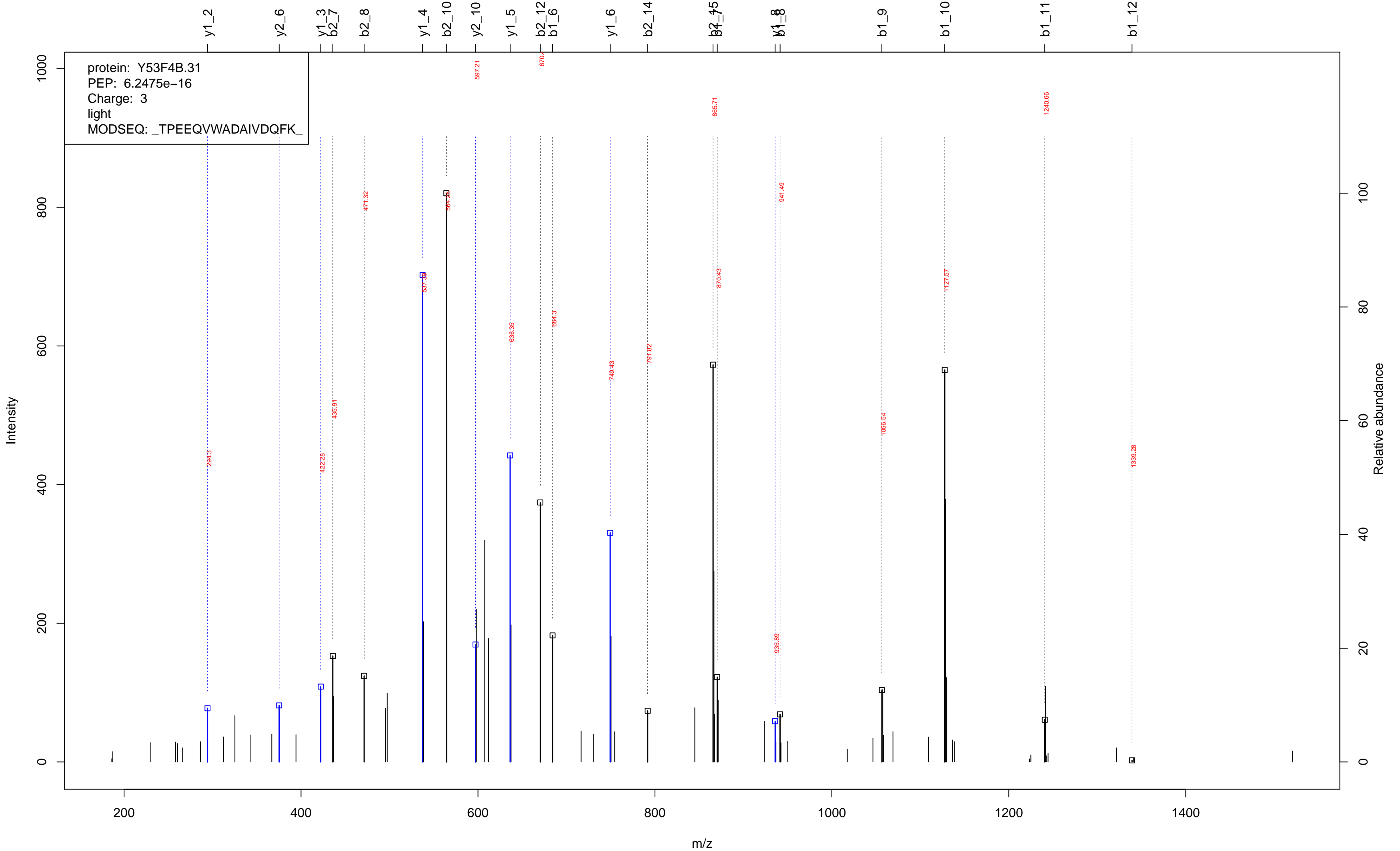
protein: W02G9.1
PEP: 0.00038132
Charge: 3
heavy
MODSEQ: _THTGQVGWQSVHR_

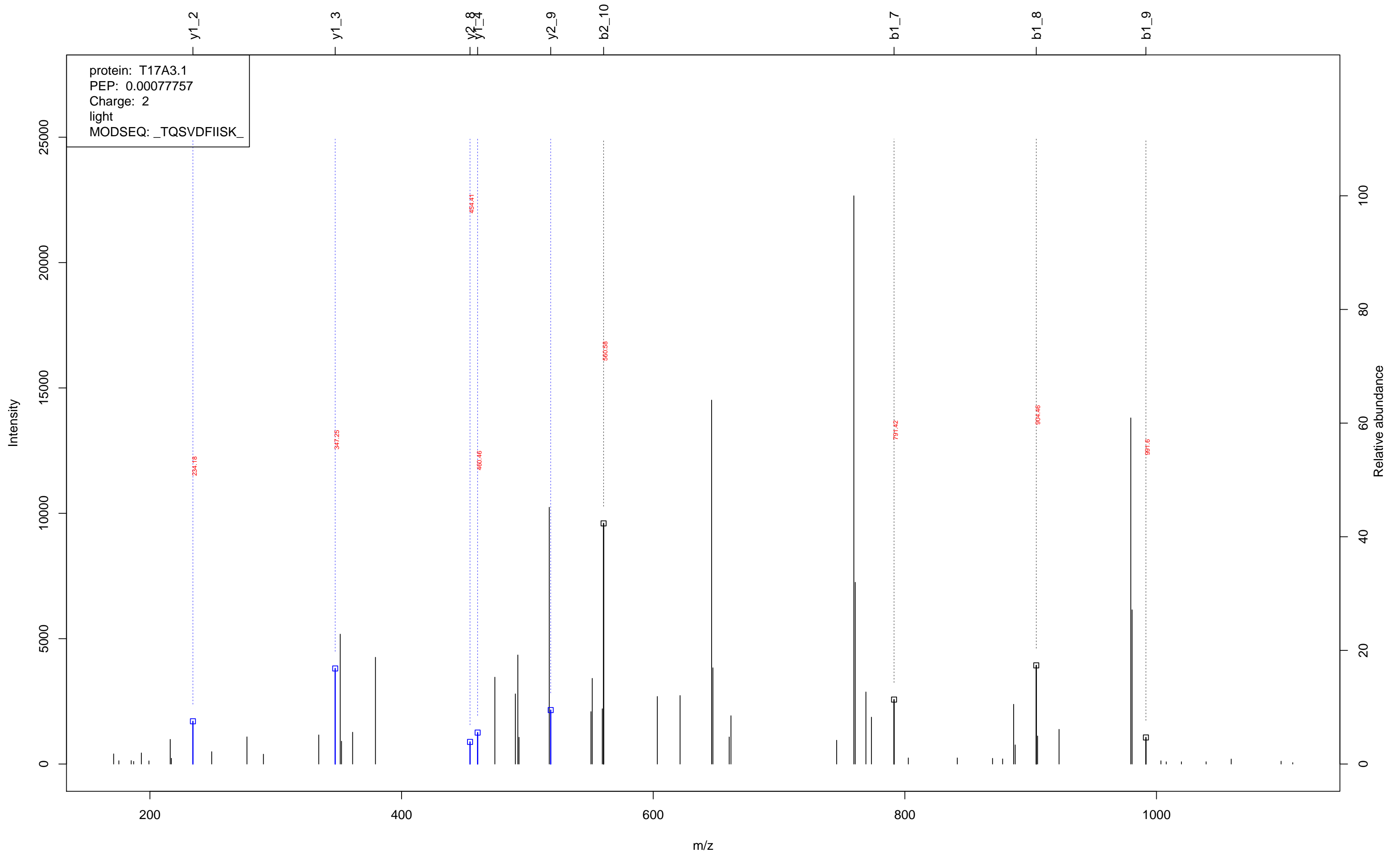


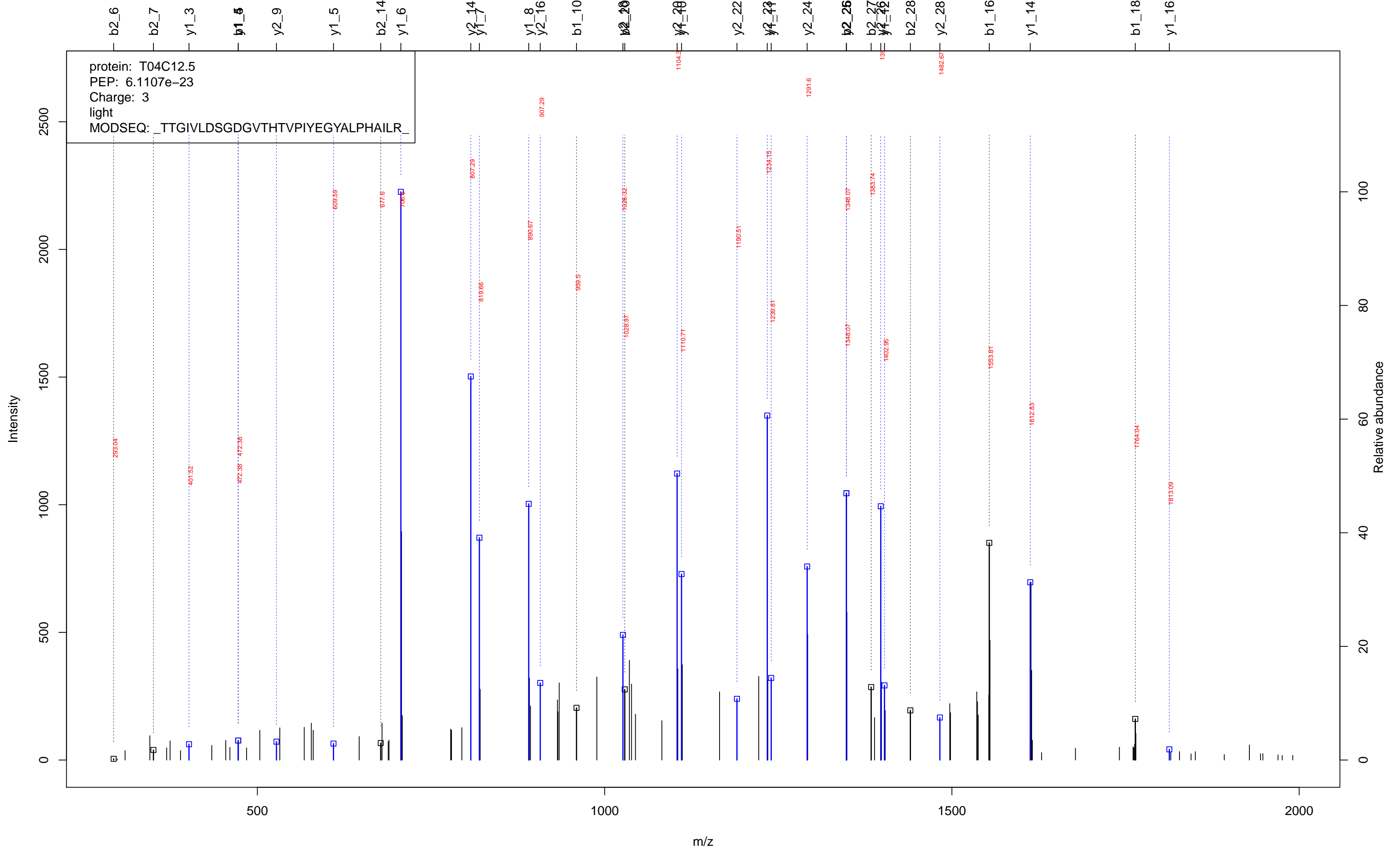








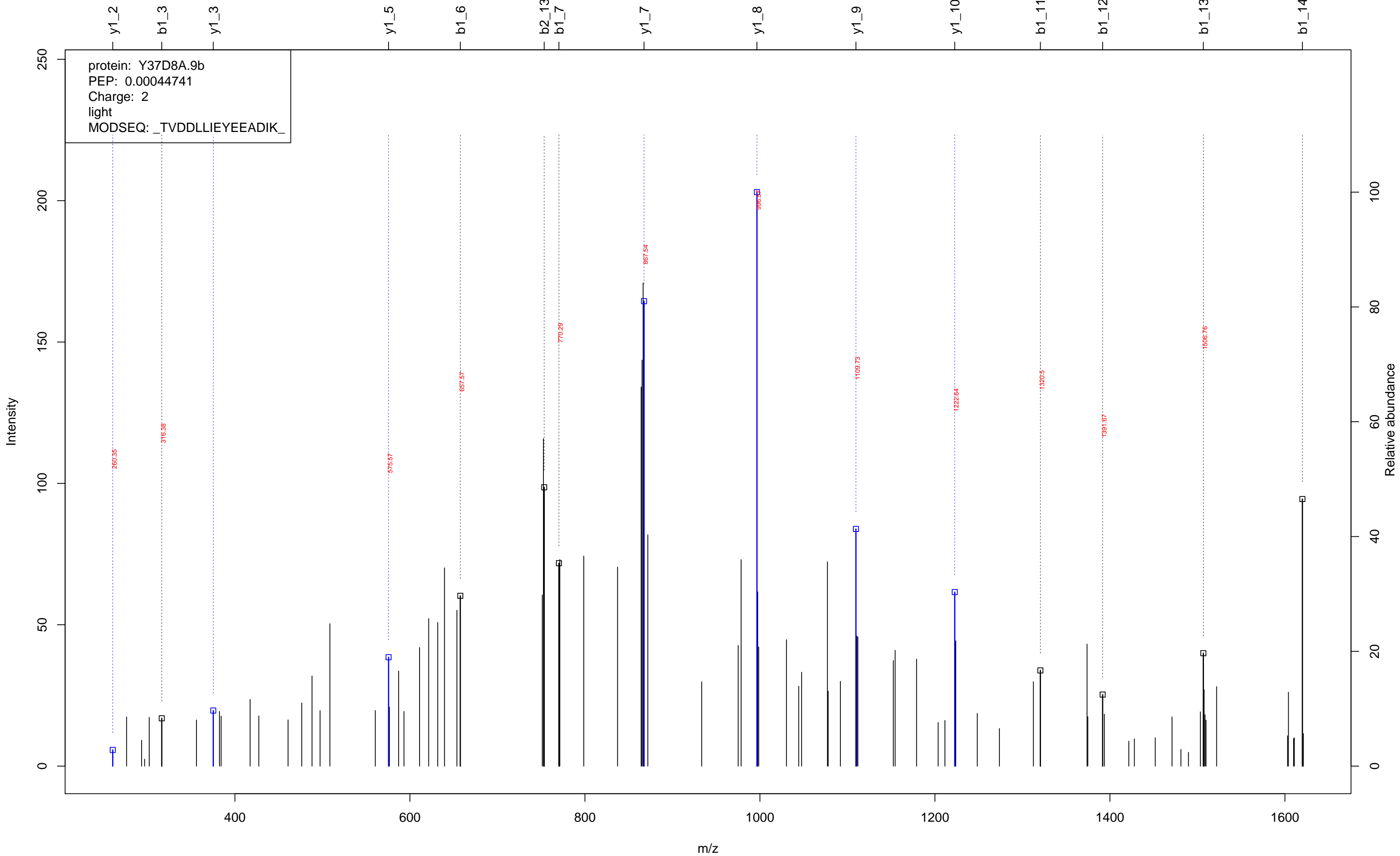


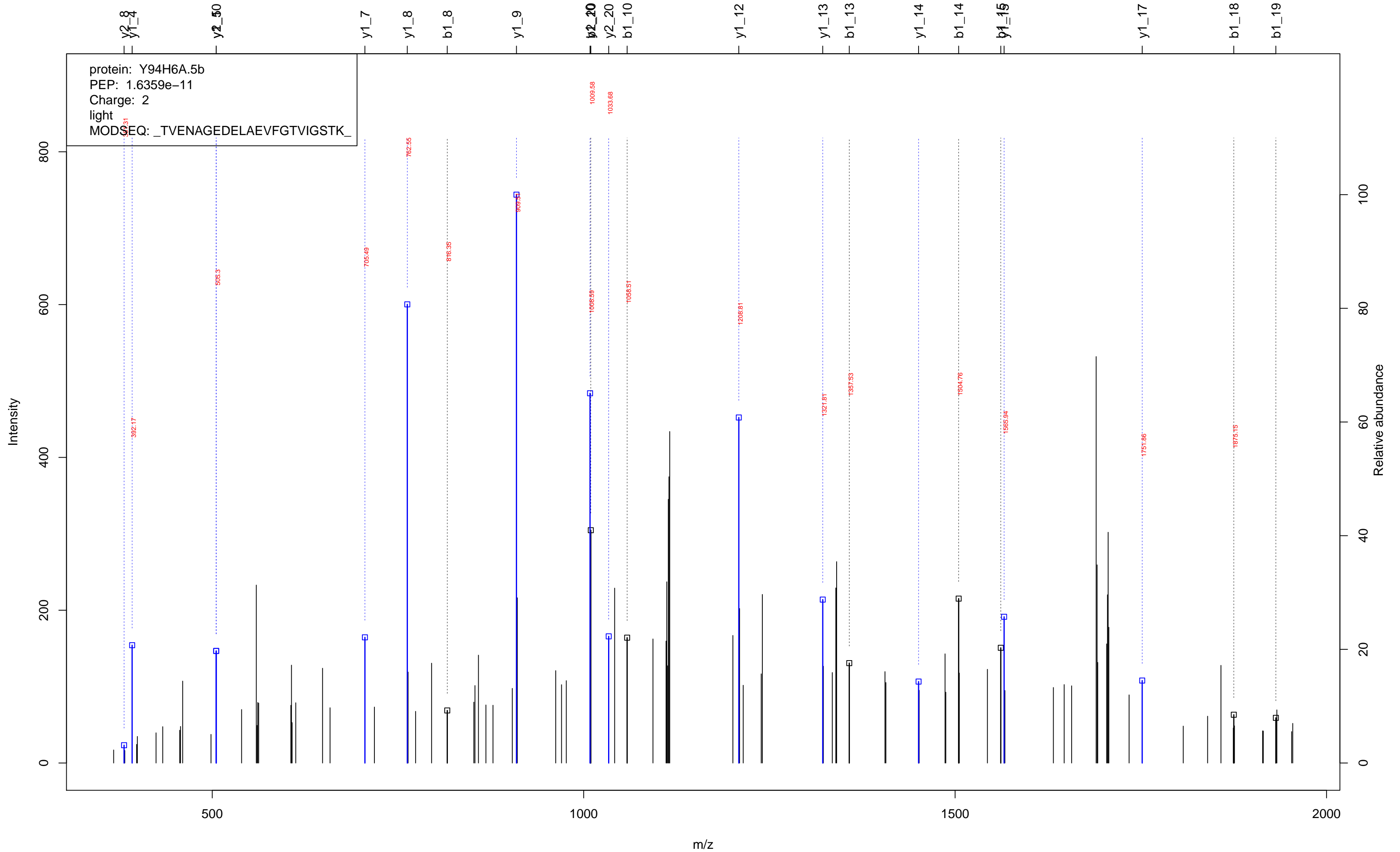


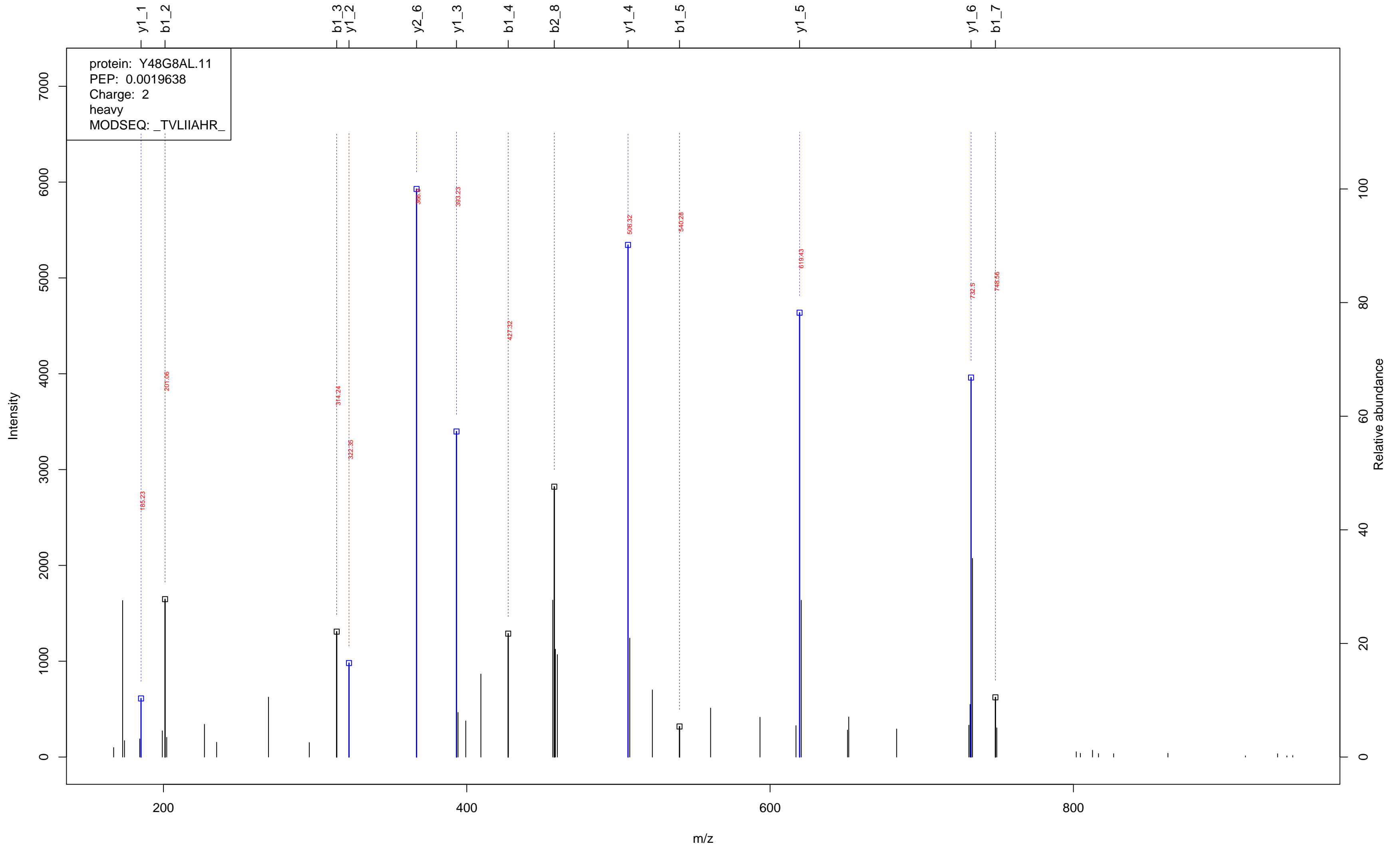
protein: T04C12.5
PEP: 6.1107e-23
Charge: 3
light
MODSEQ: _TTGIVLDSGDGVTHTVPIYEGYALPHAILR_

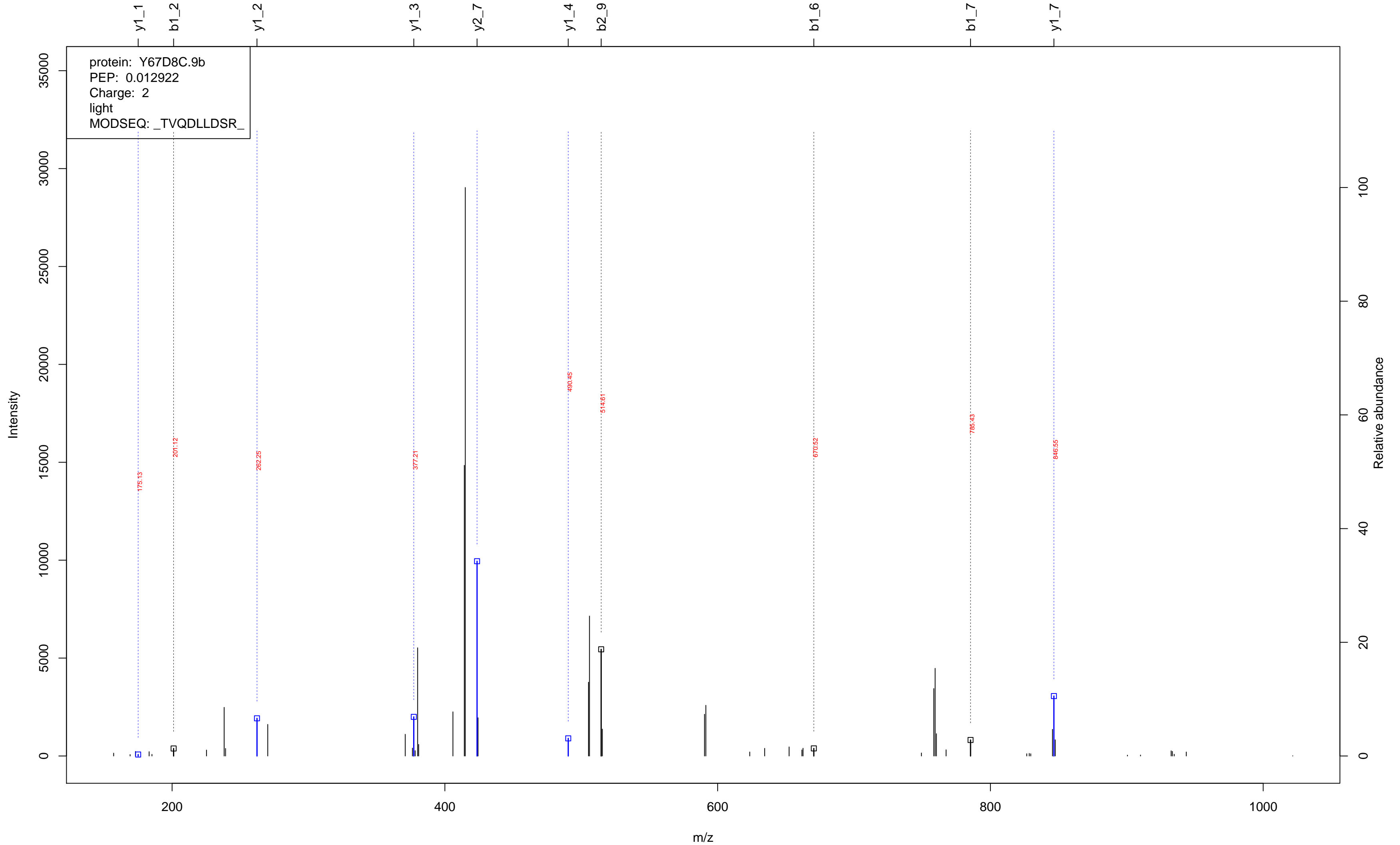
315

protein: Y37D8A.9b
PEP: 0.00044741
Charge: 2
light
MODSEQ: _TVDDLLIEYEEADIK_



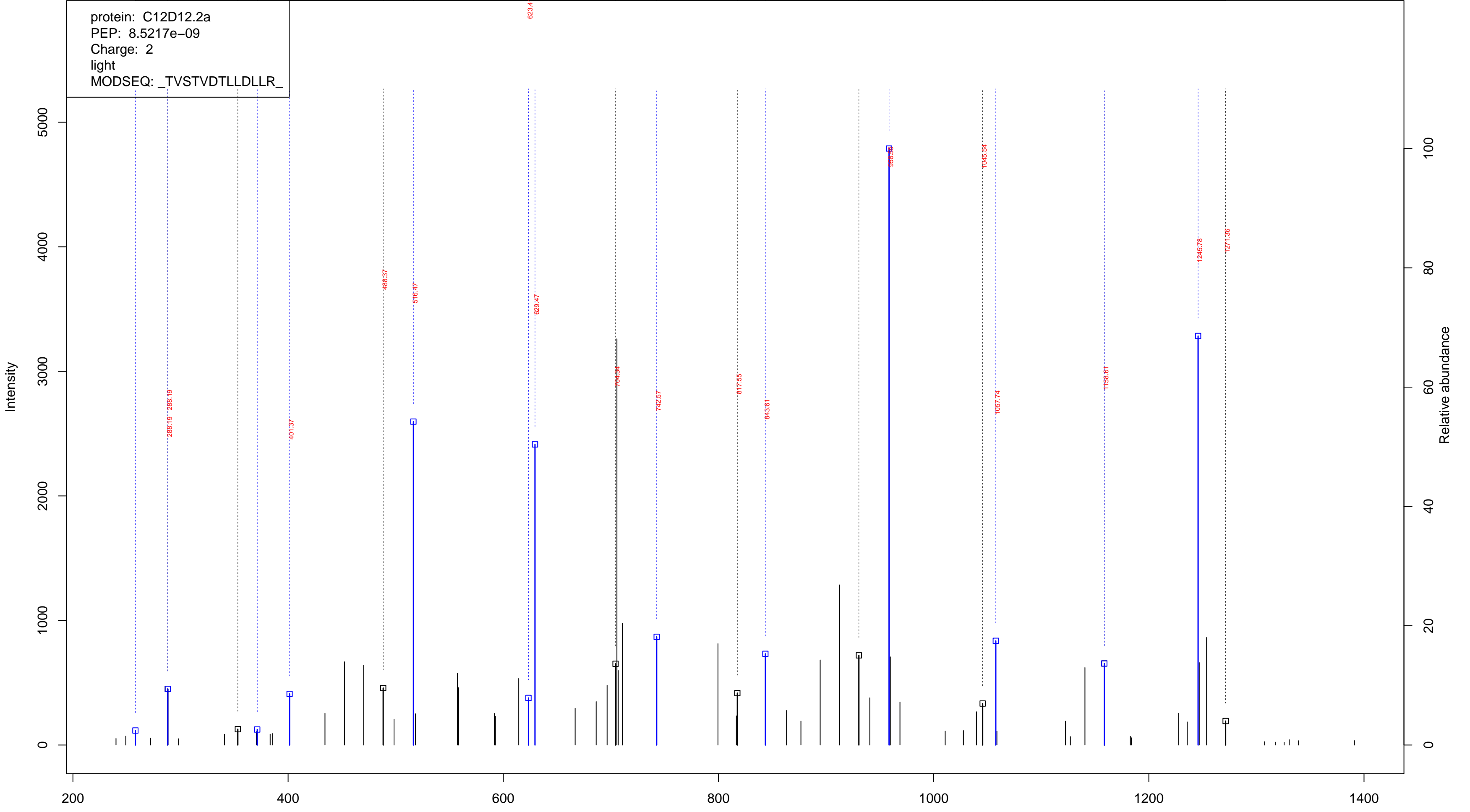


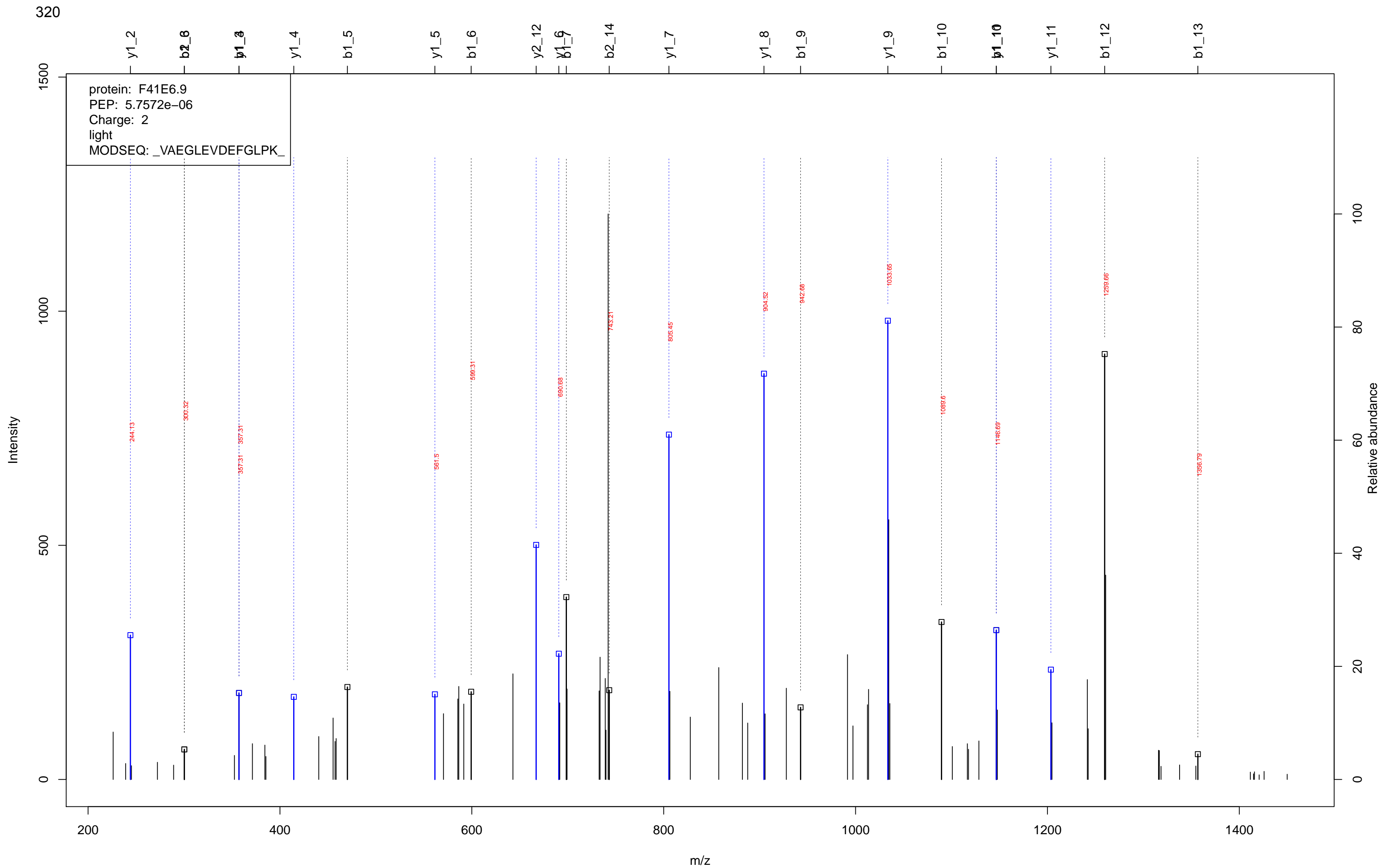


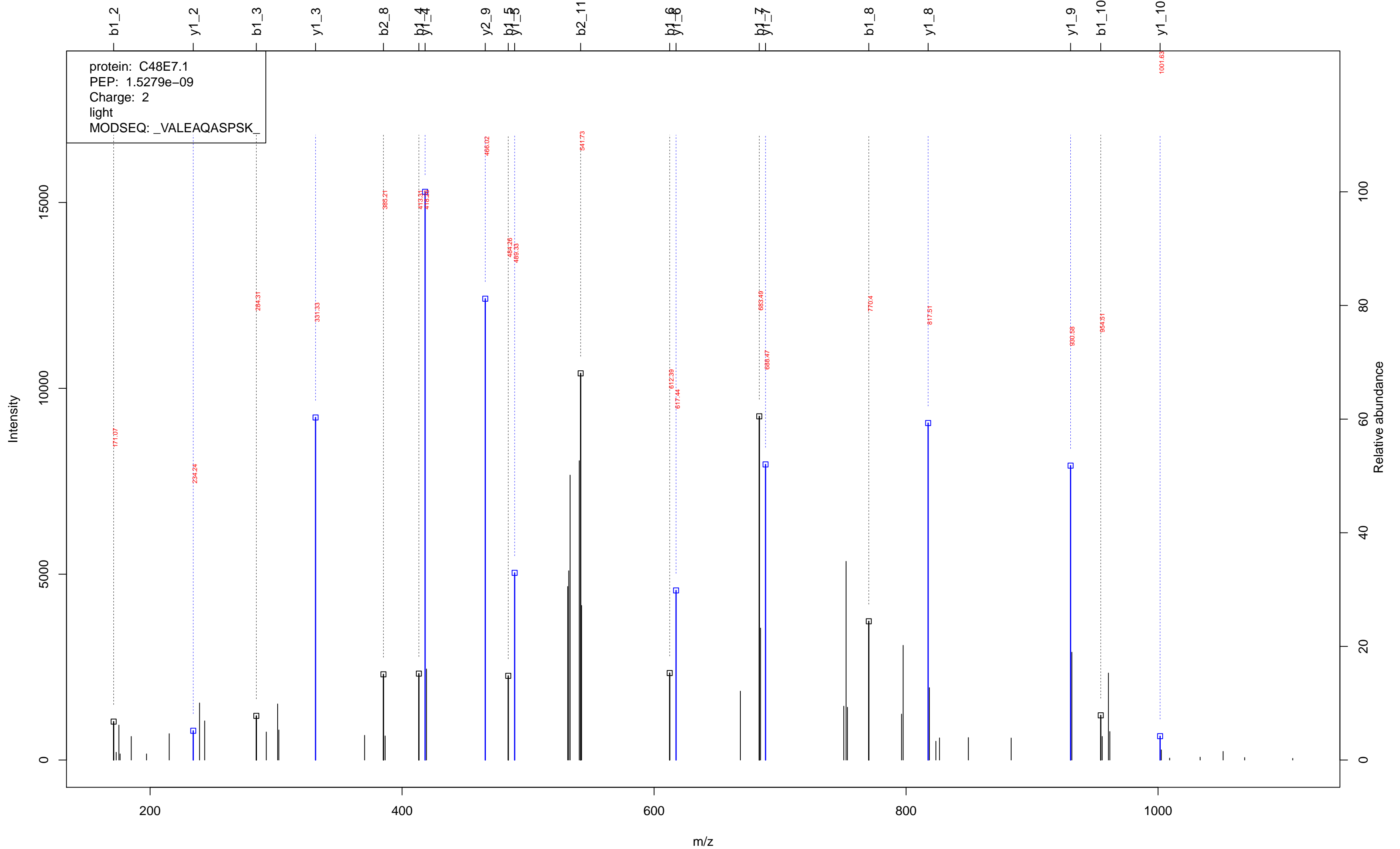


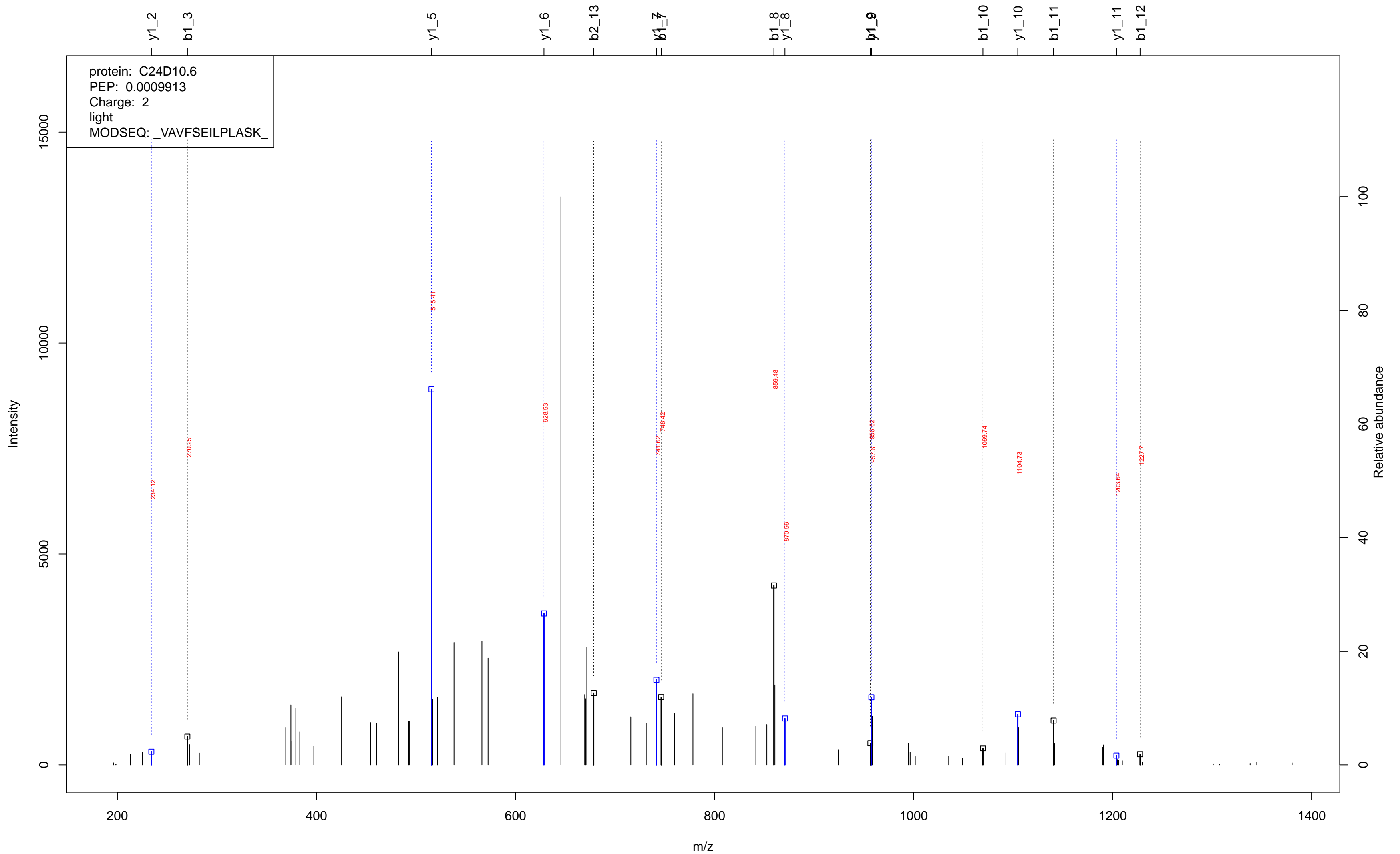
y2_4 p1_3 b2_7 y2_6 y1_3 b1_5 y1_4 y2_5 y1_5 b1_7 y1_6 b1_8 y1_7 b1_9 y1_8 b1_10 y1_9 p1_10 y1_11 b1_12

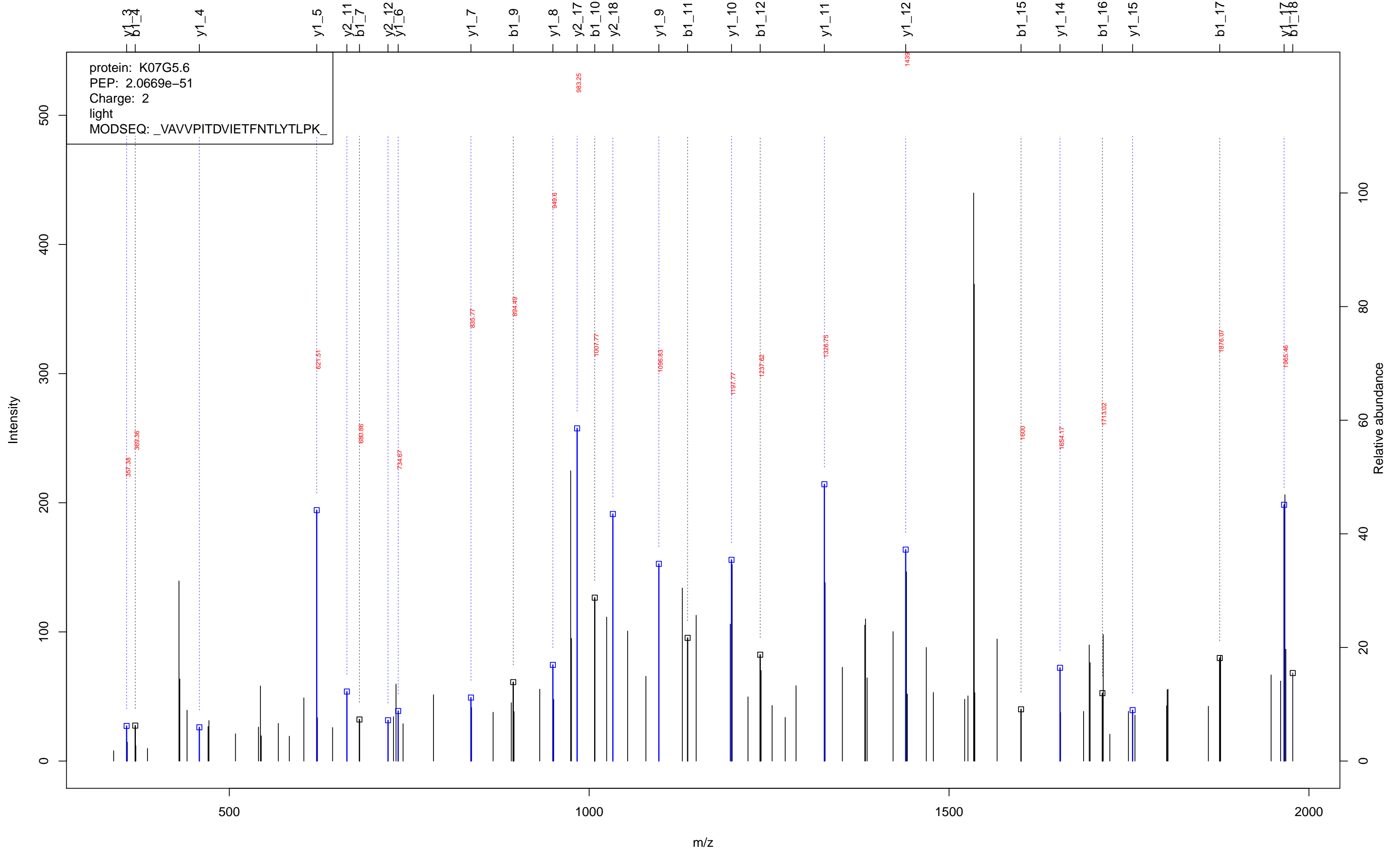
protein: C12D12.2a
PEP: 8.5217e-09
Charge: 2
light
MODSEQ: _TVSTVDTLLDLLR_

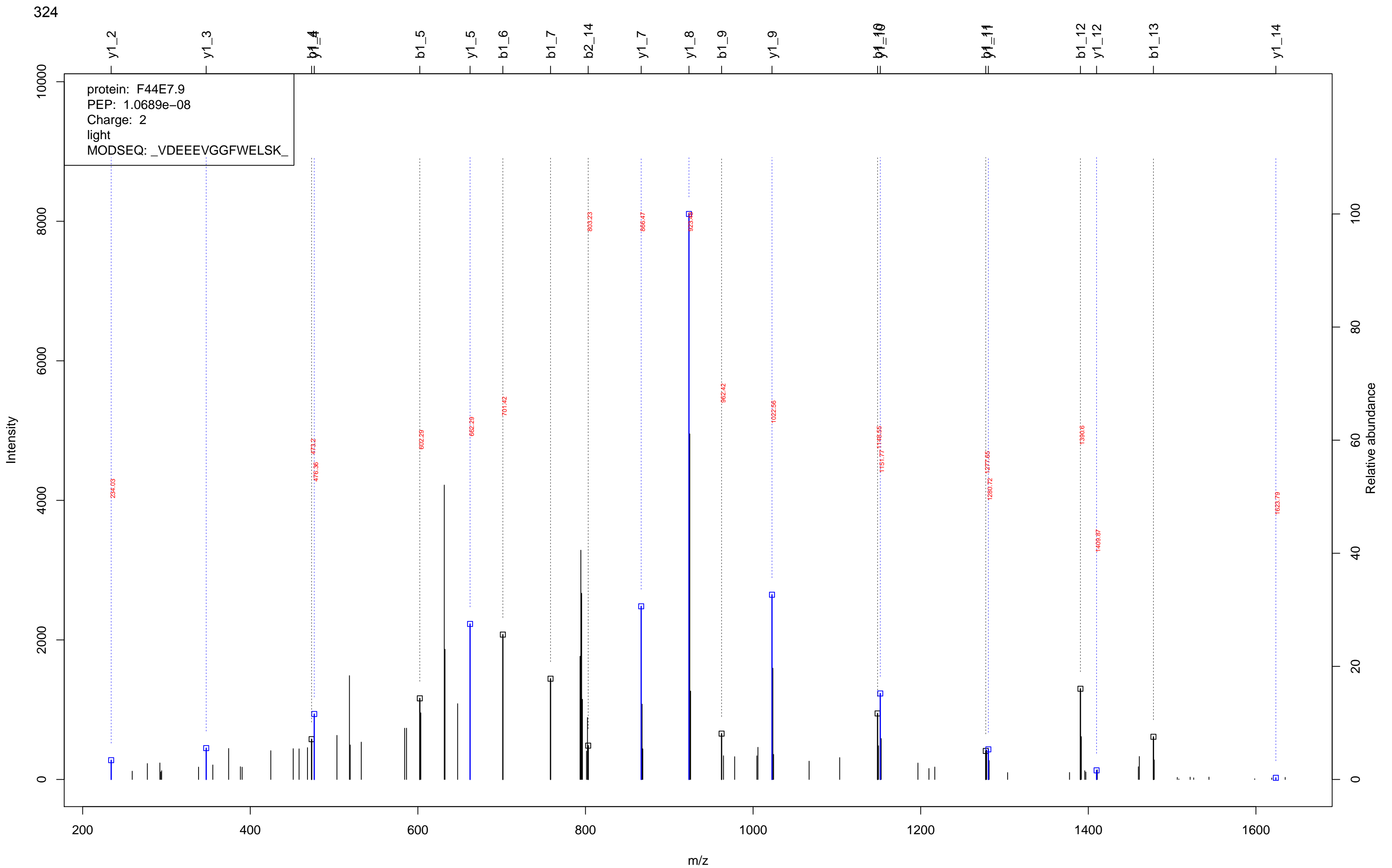


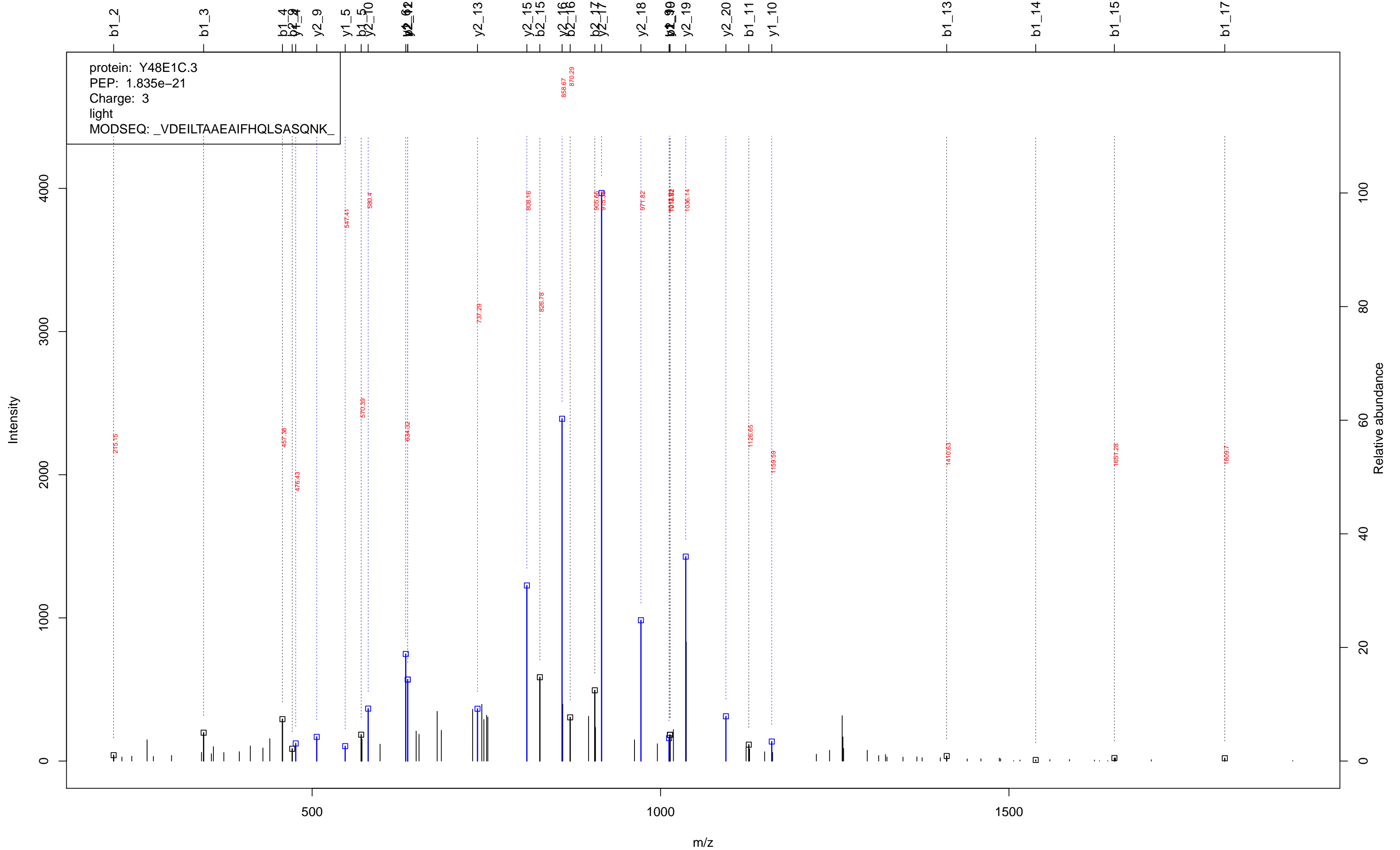


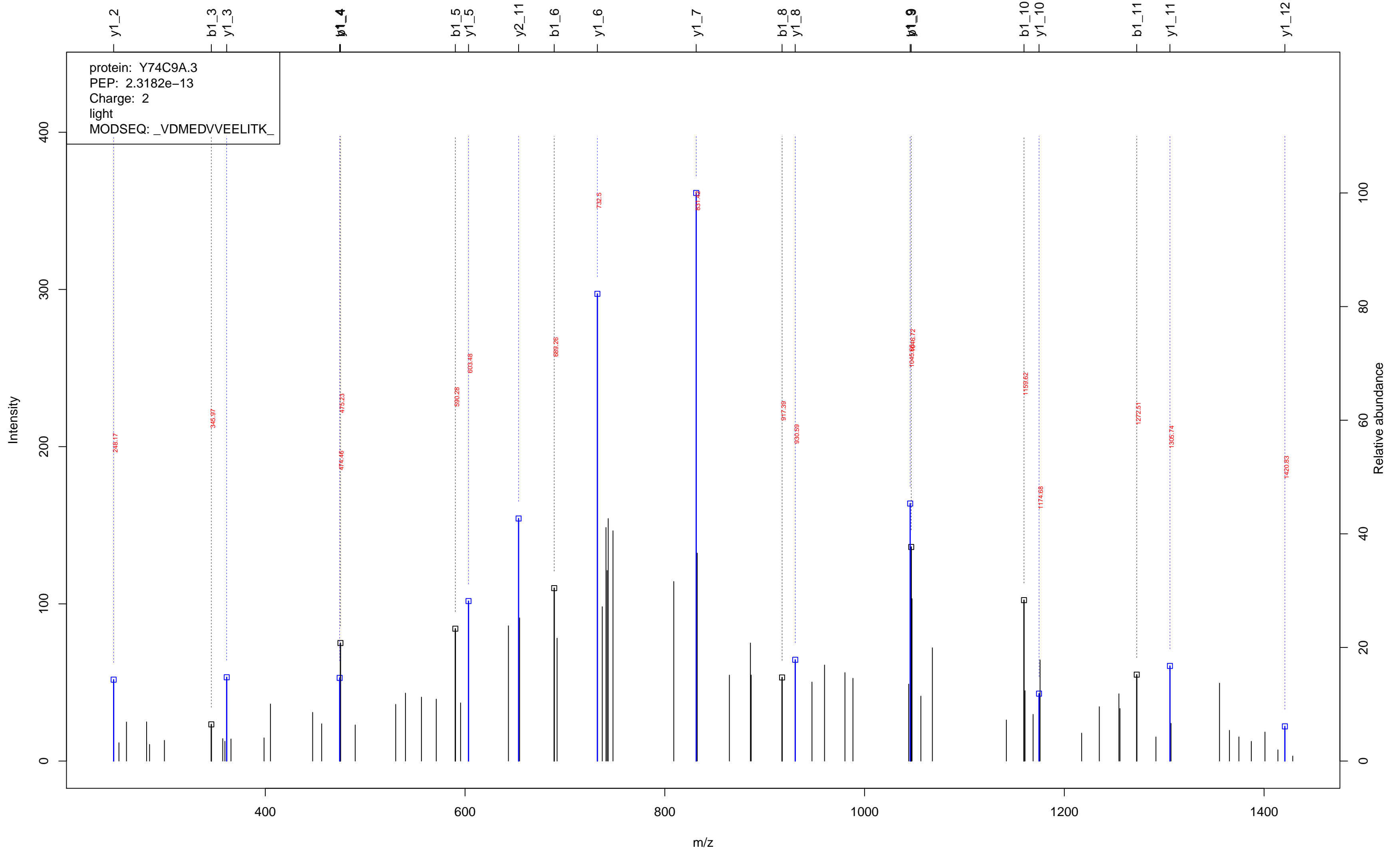


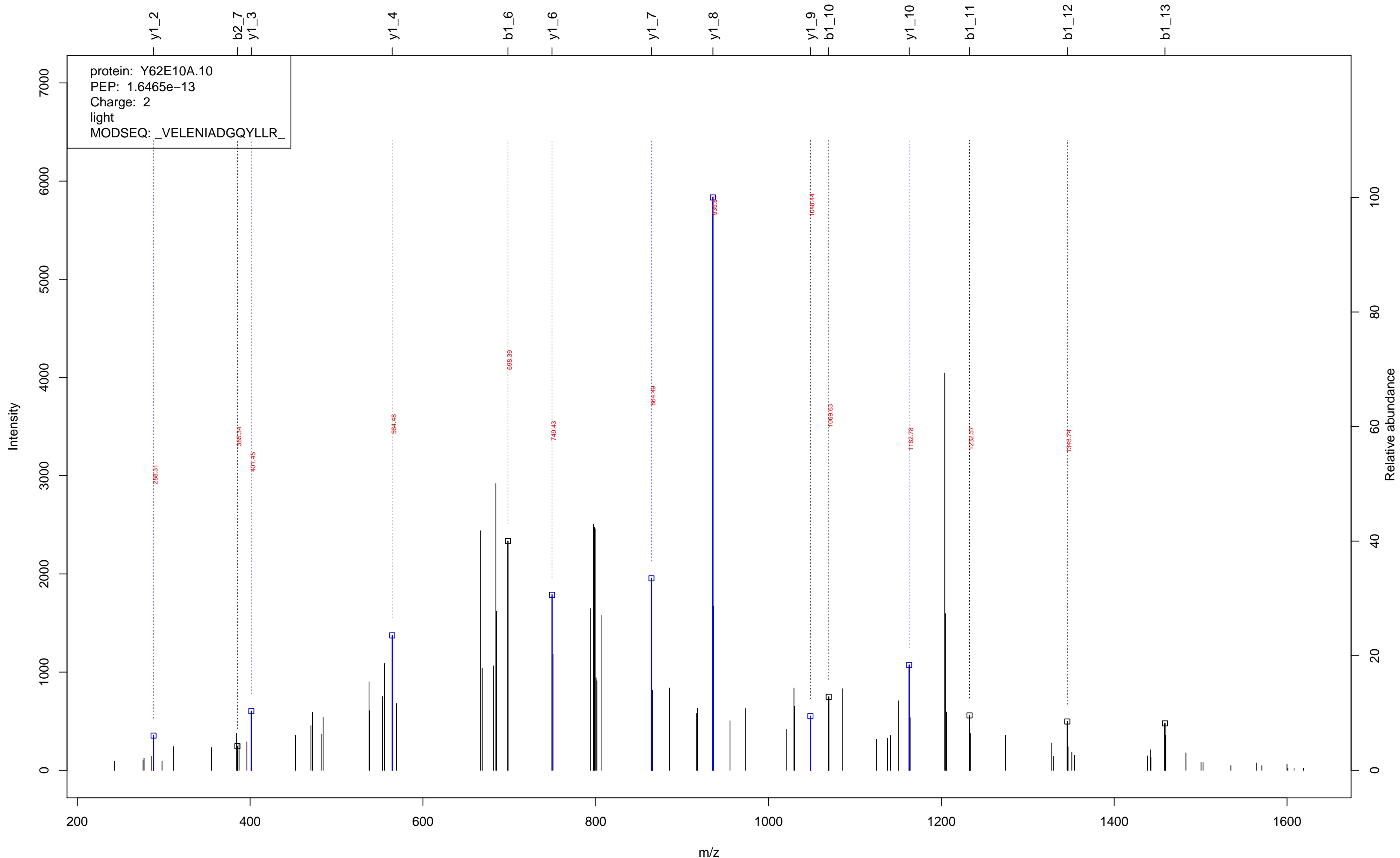


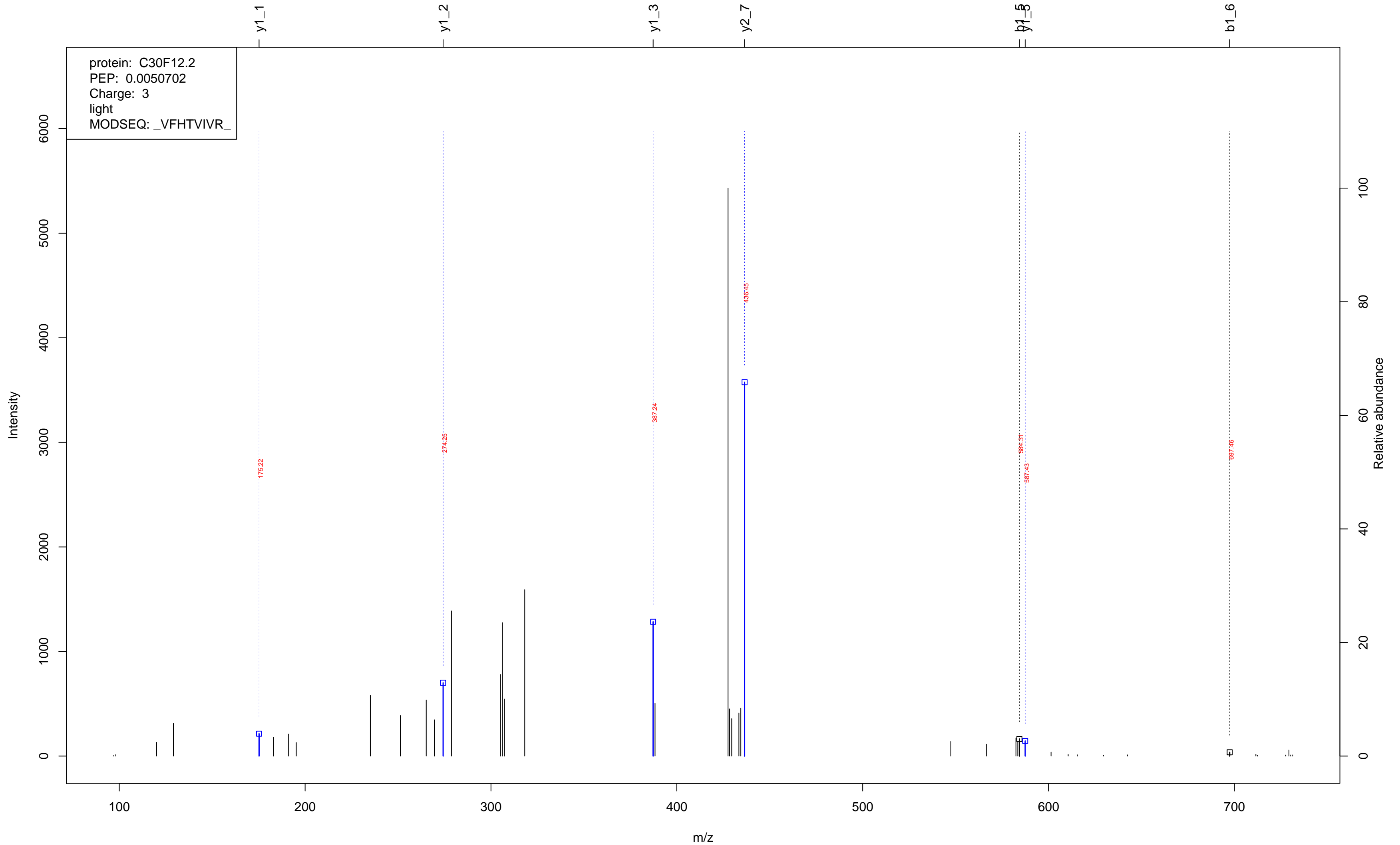


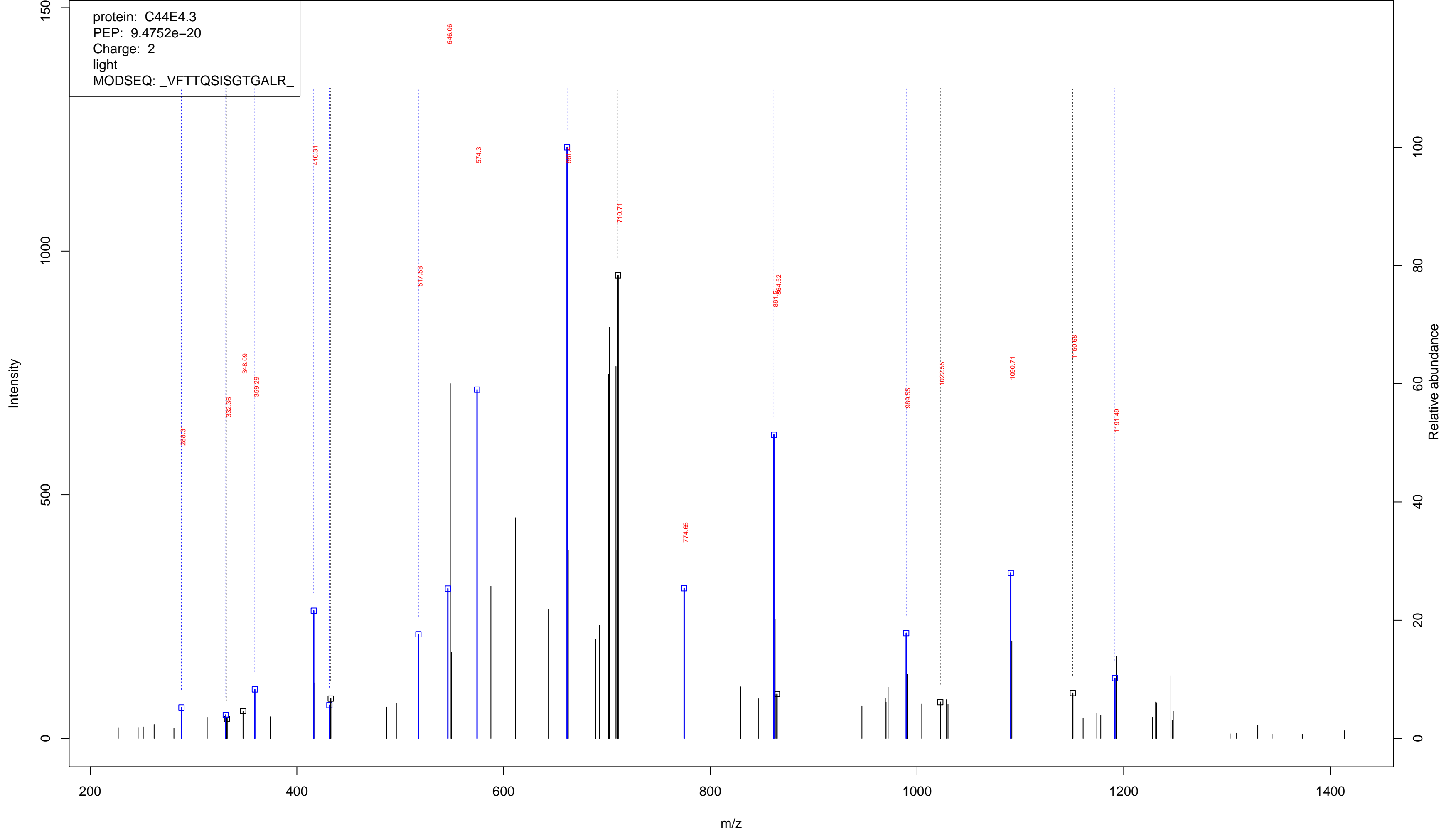




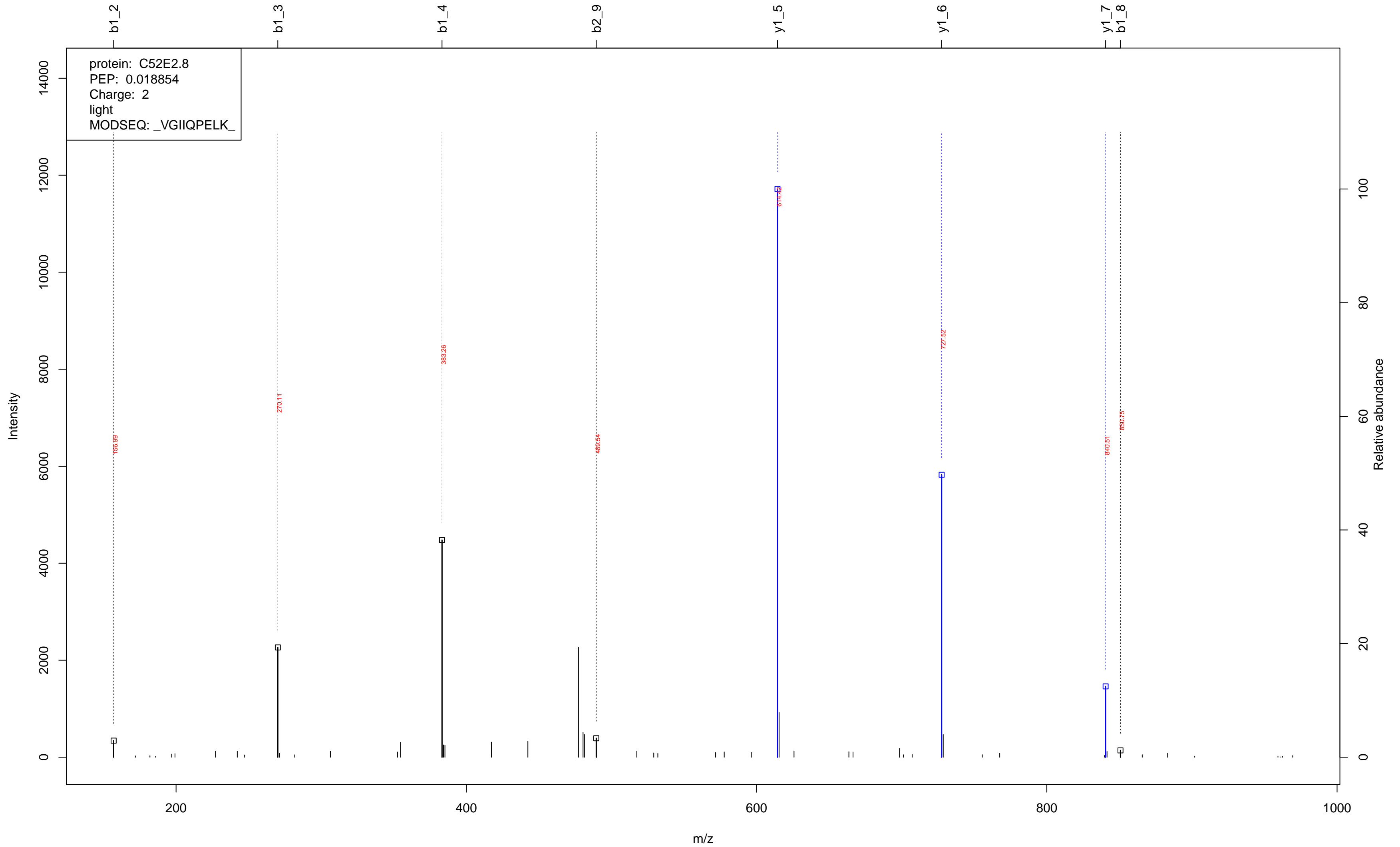


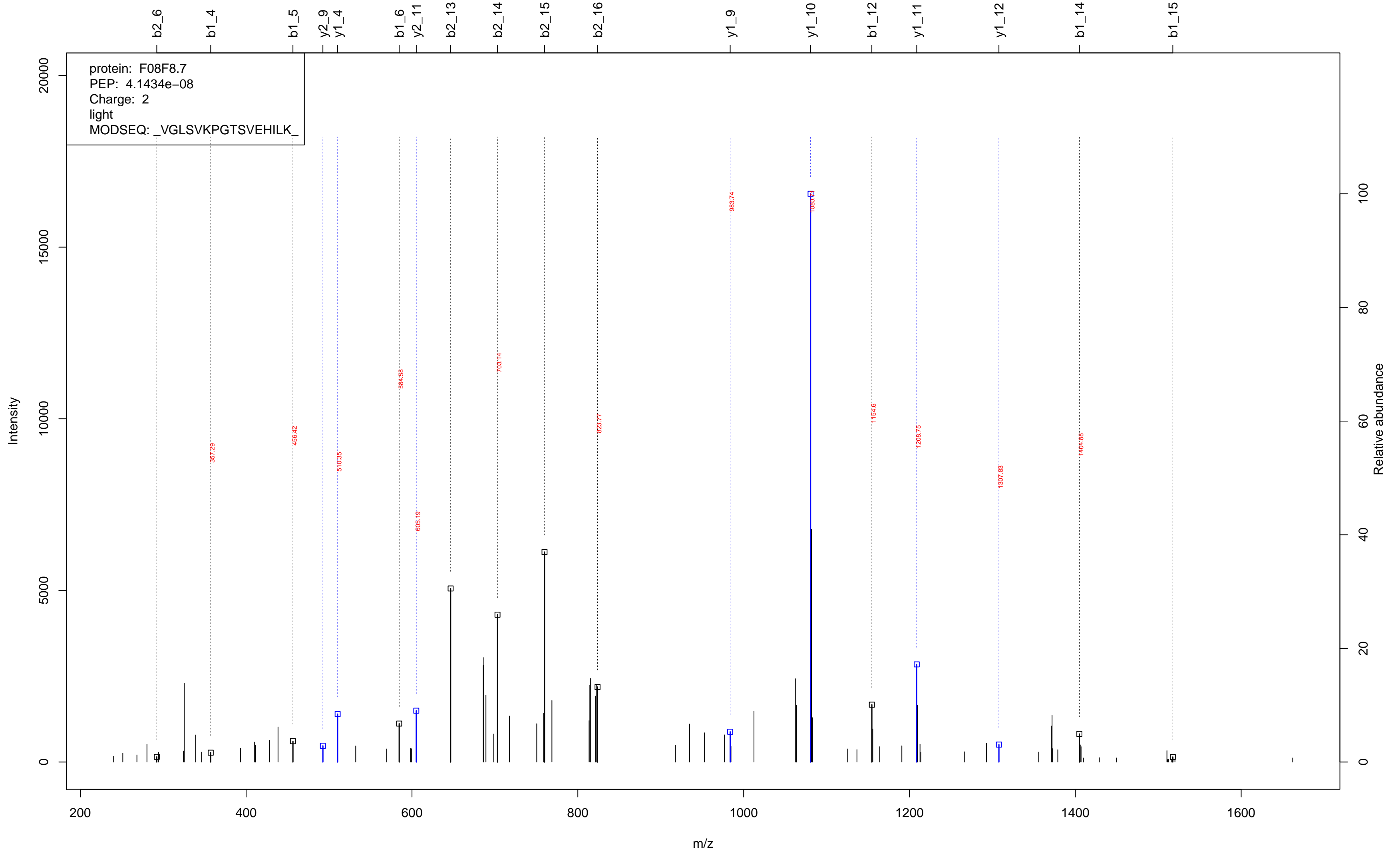


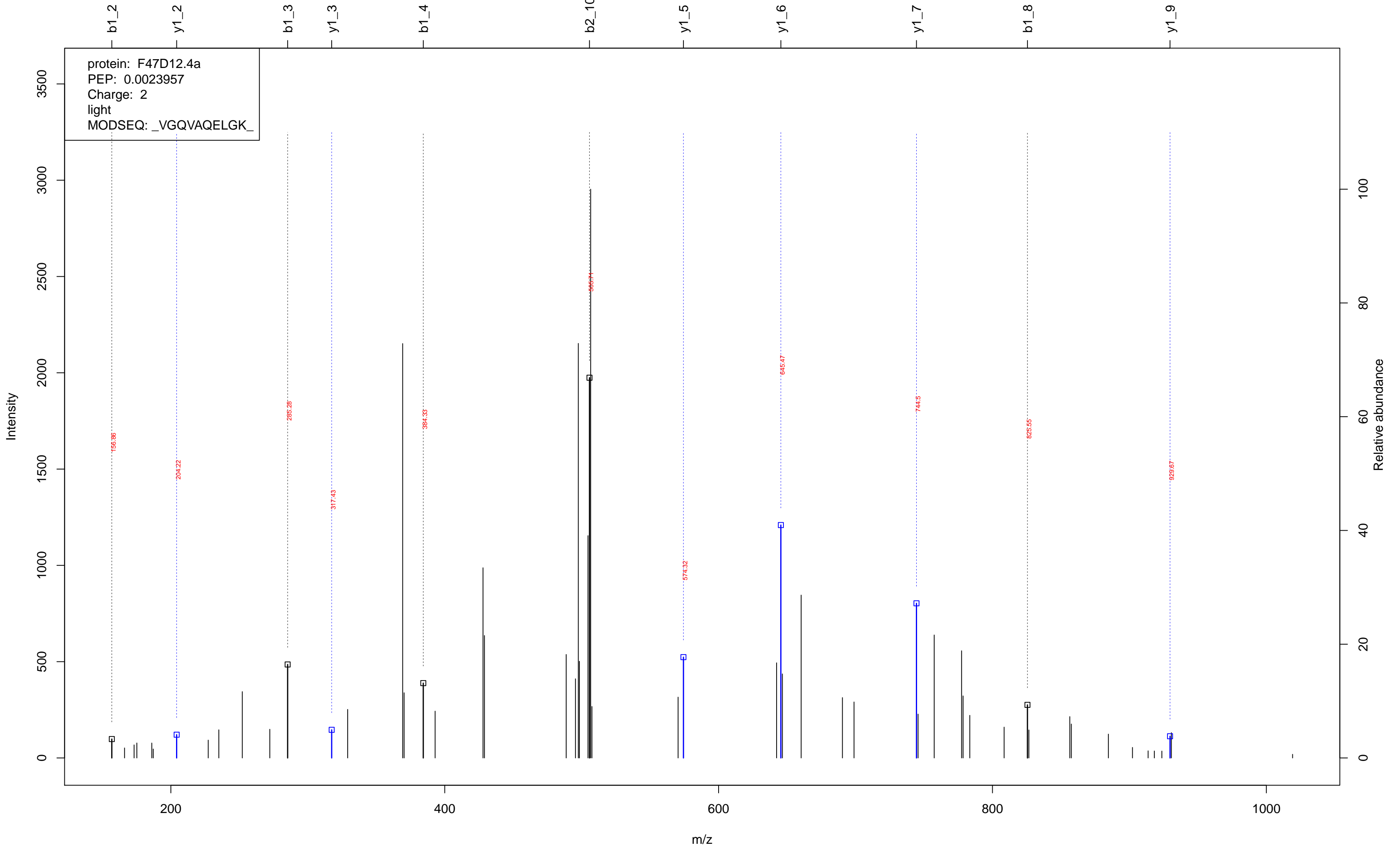


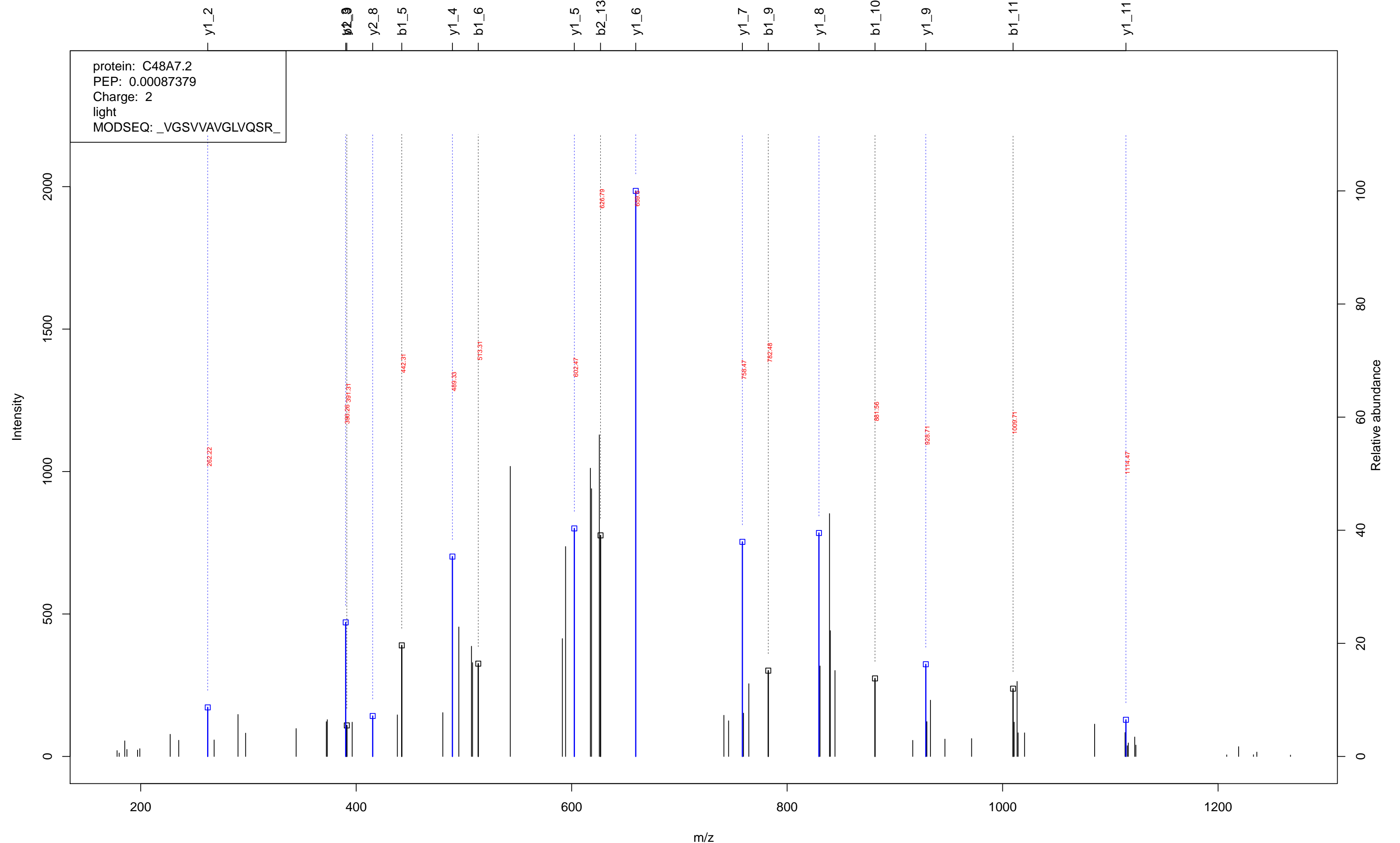


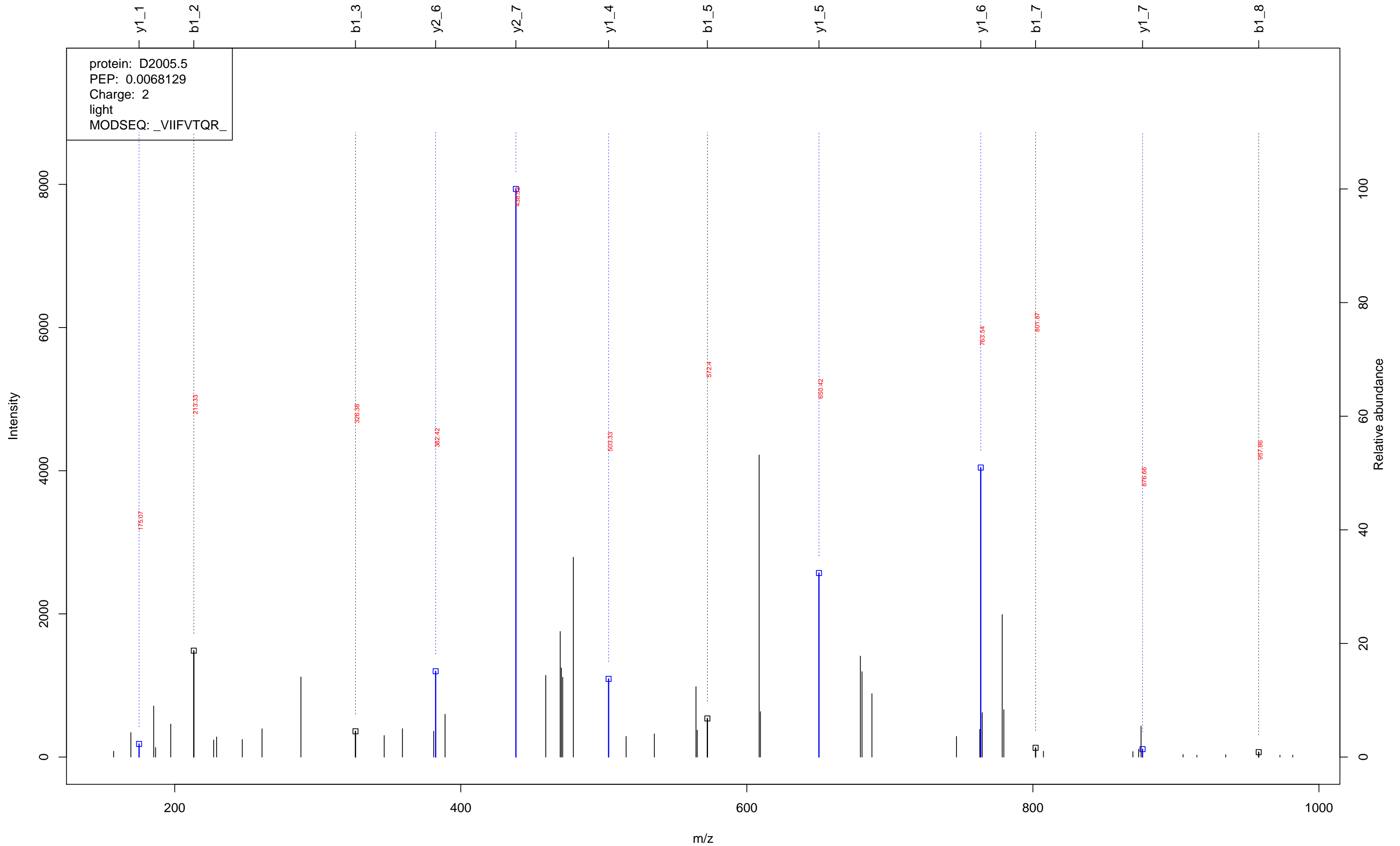
330

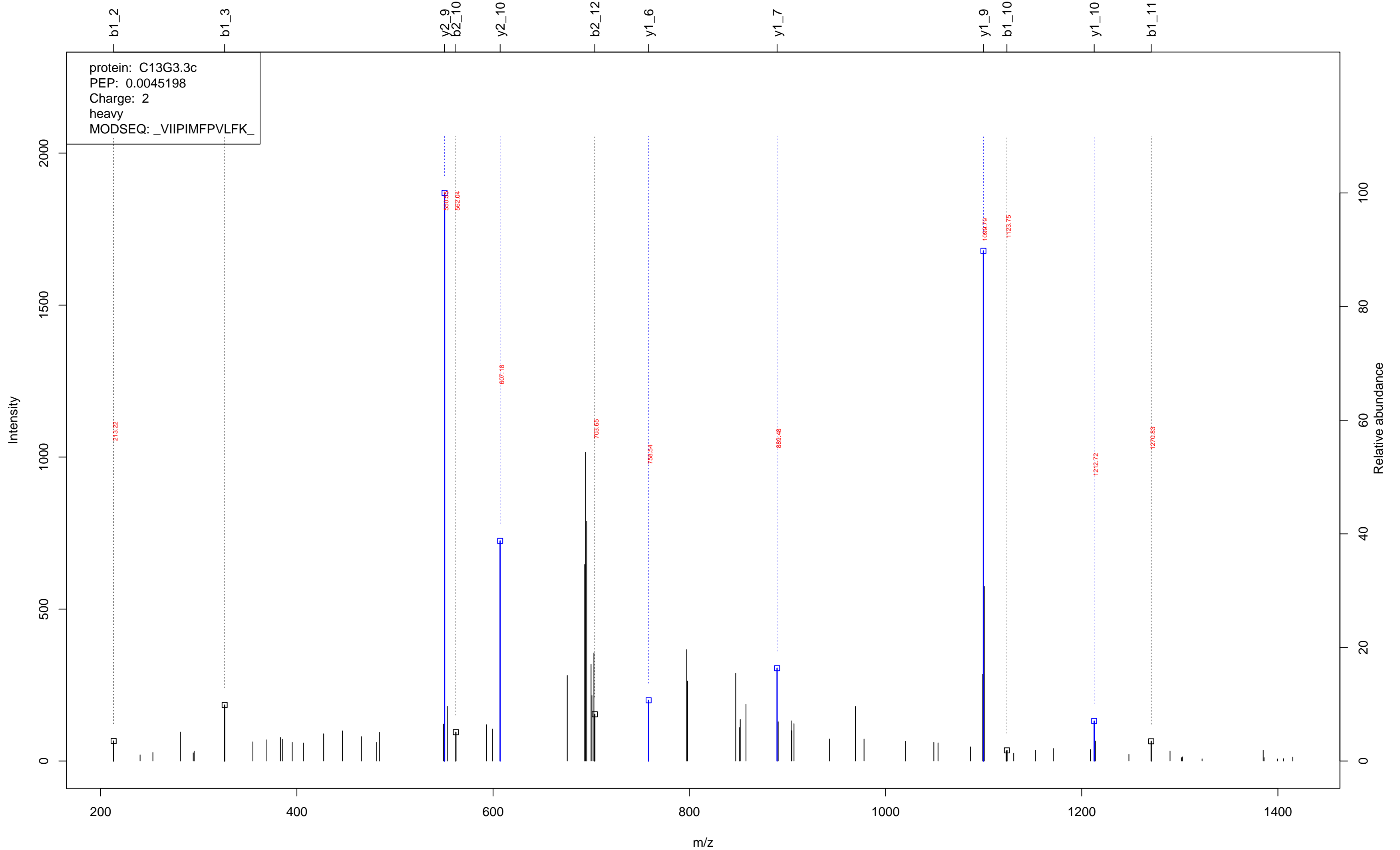


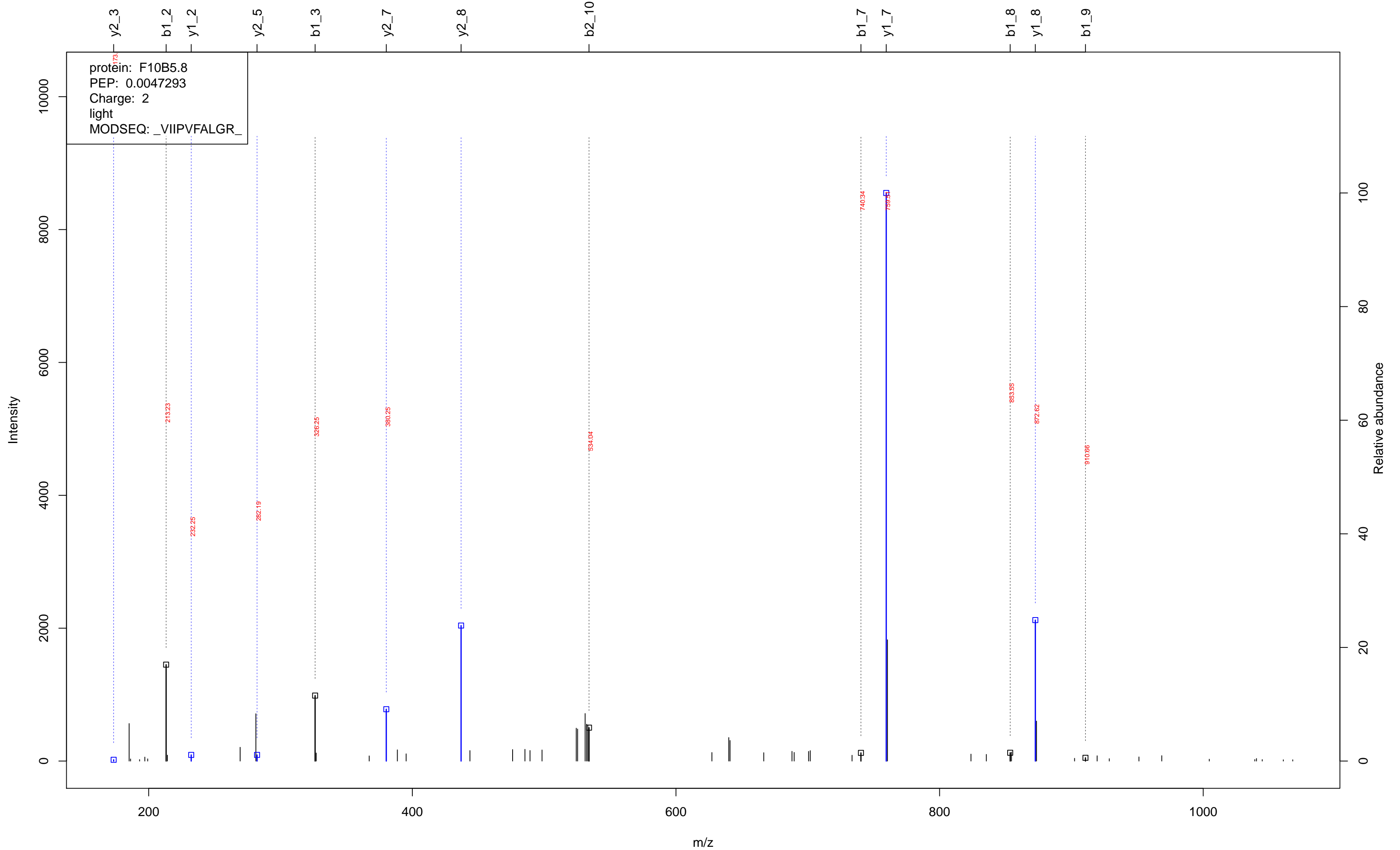


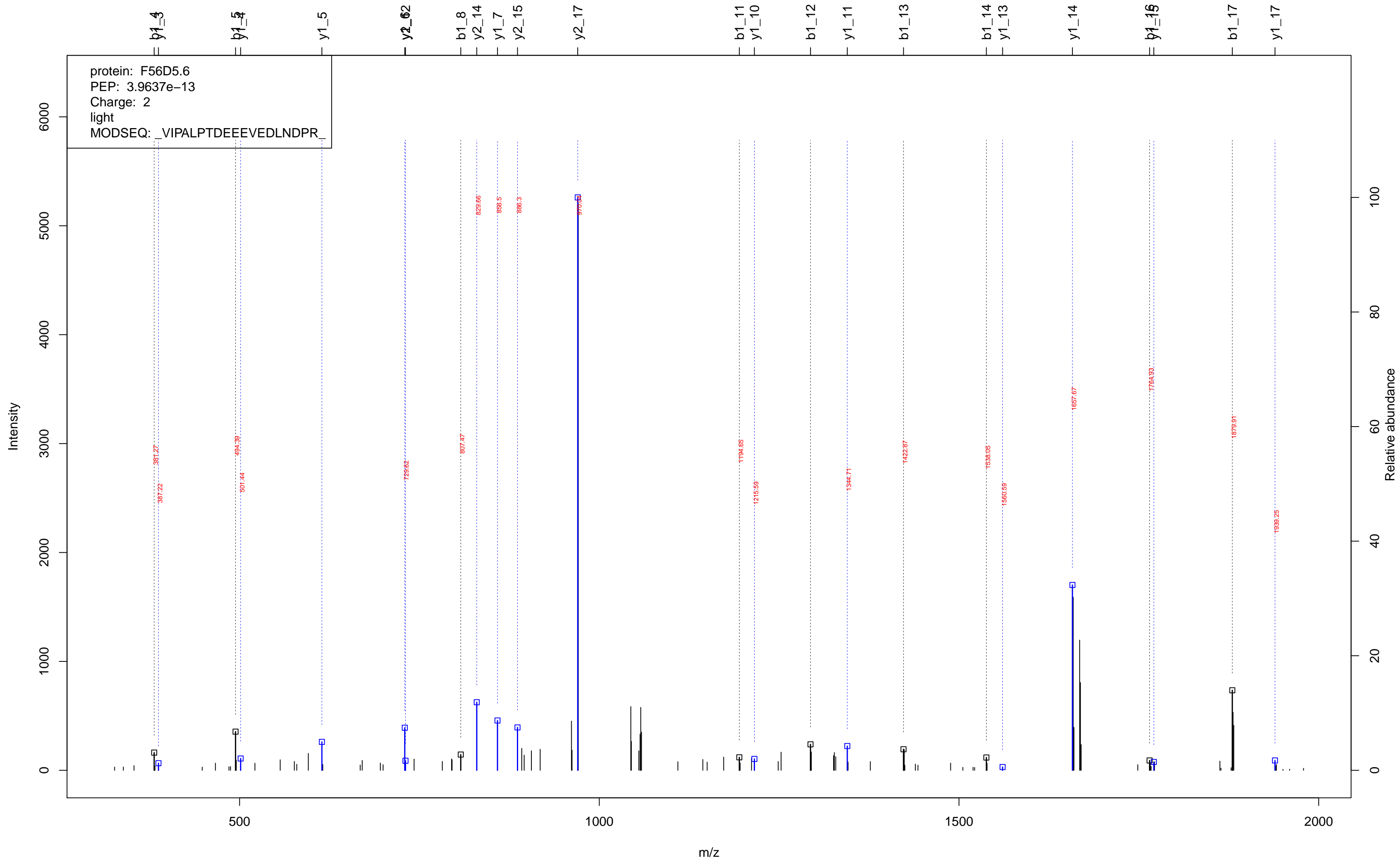


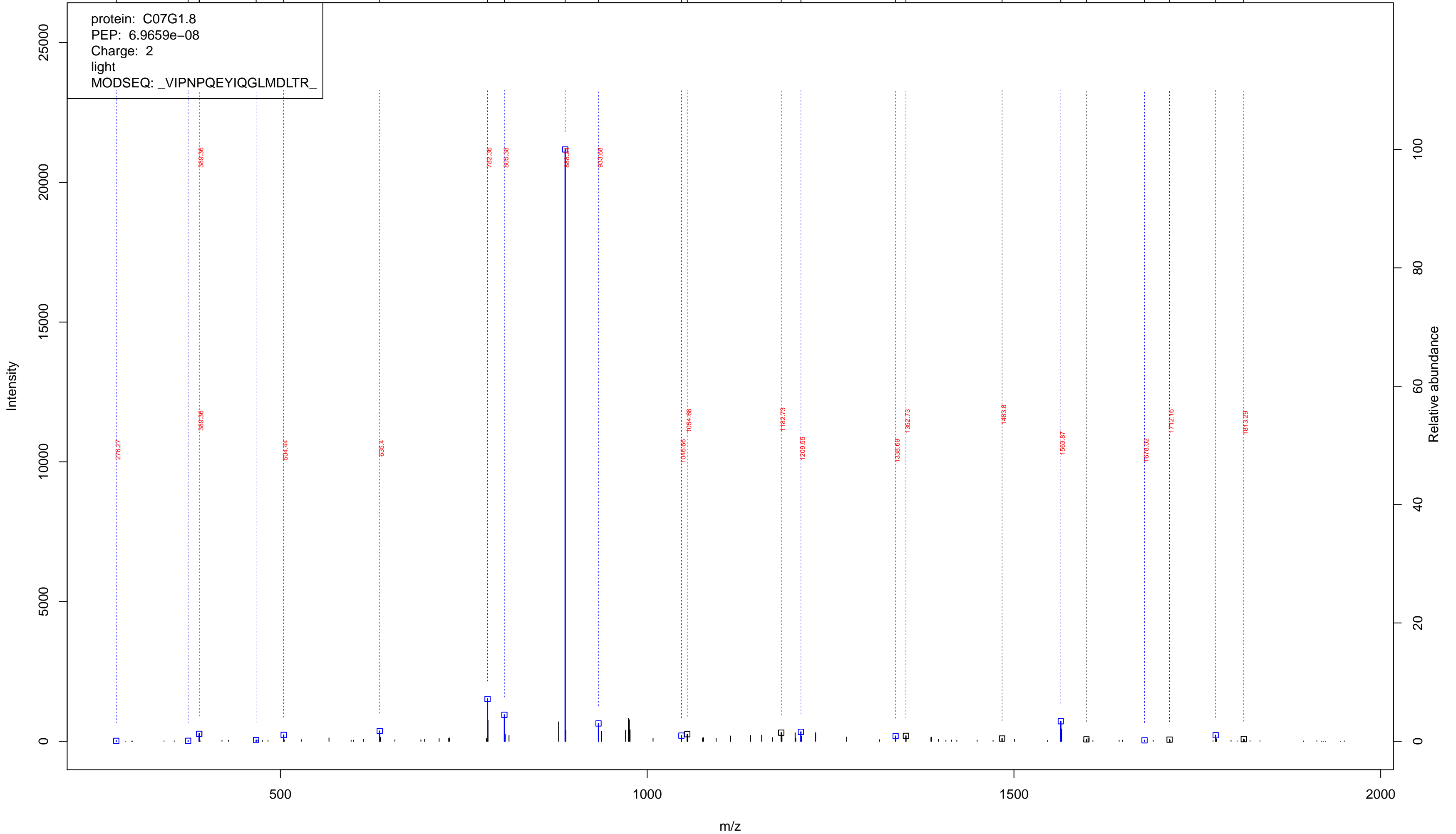


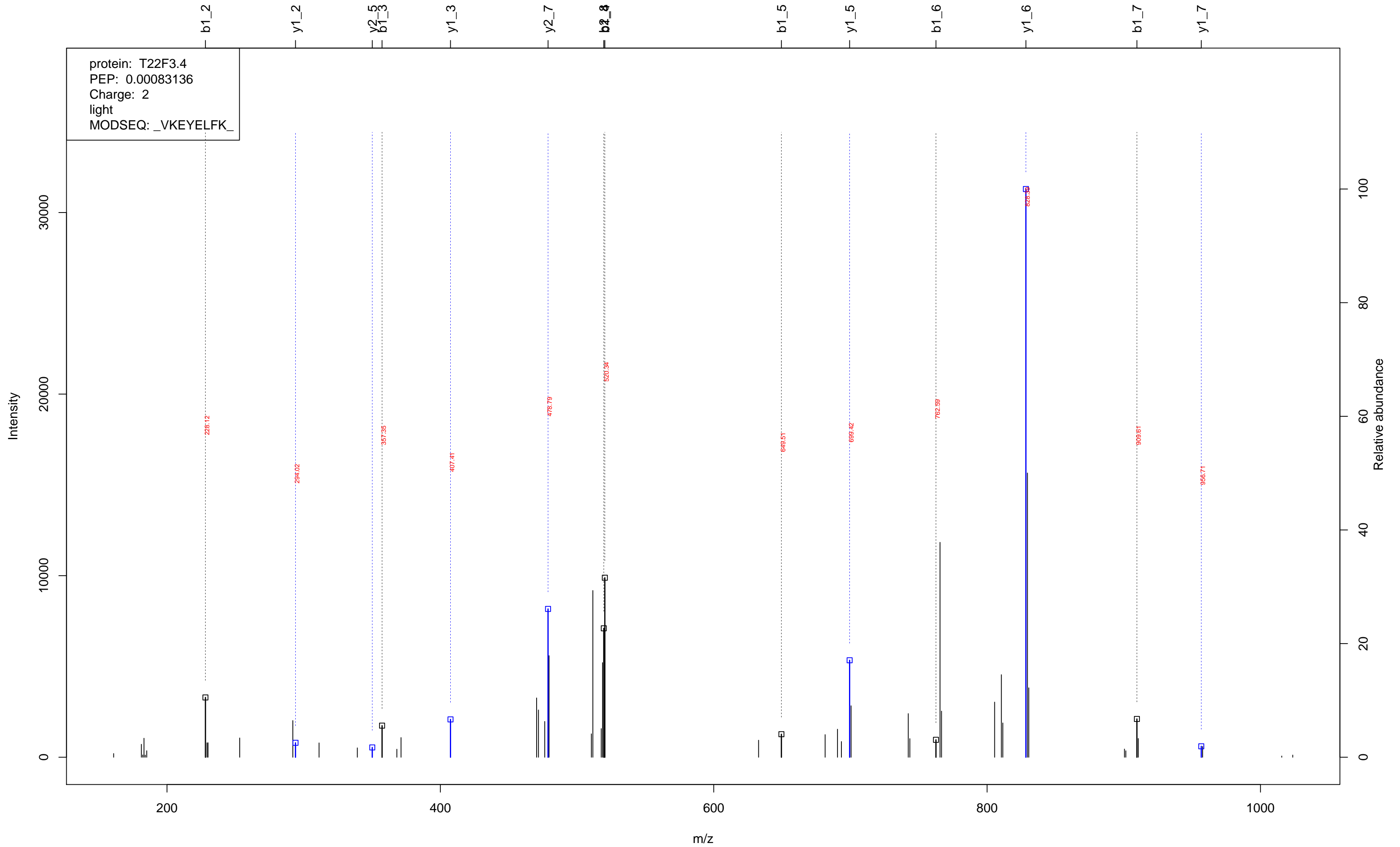


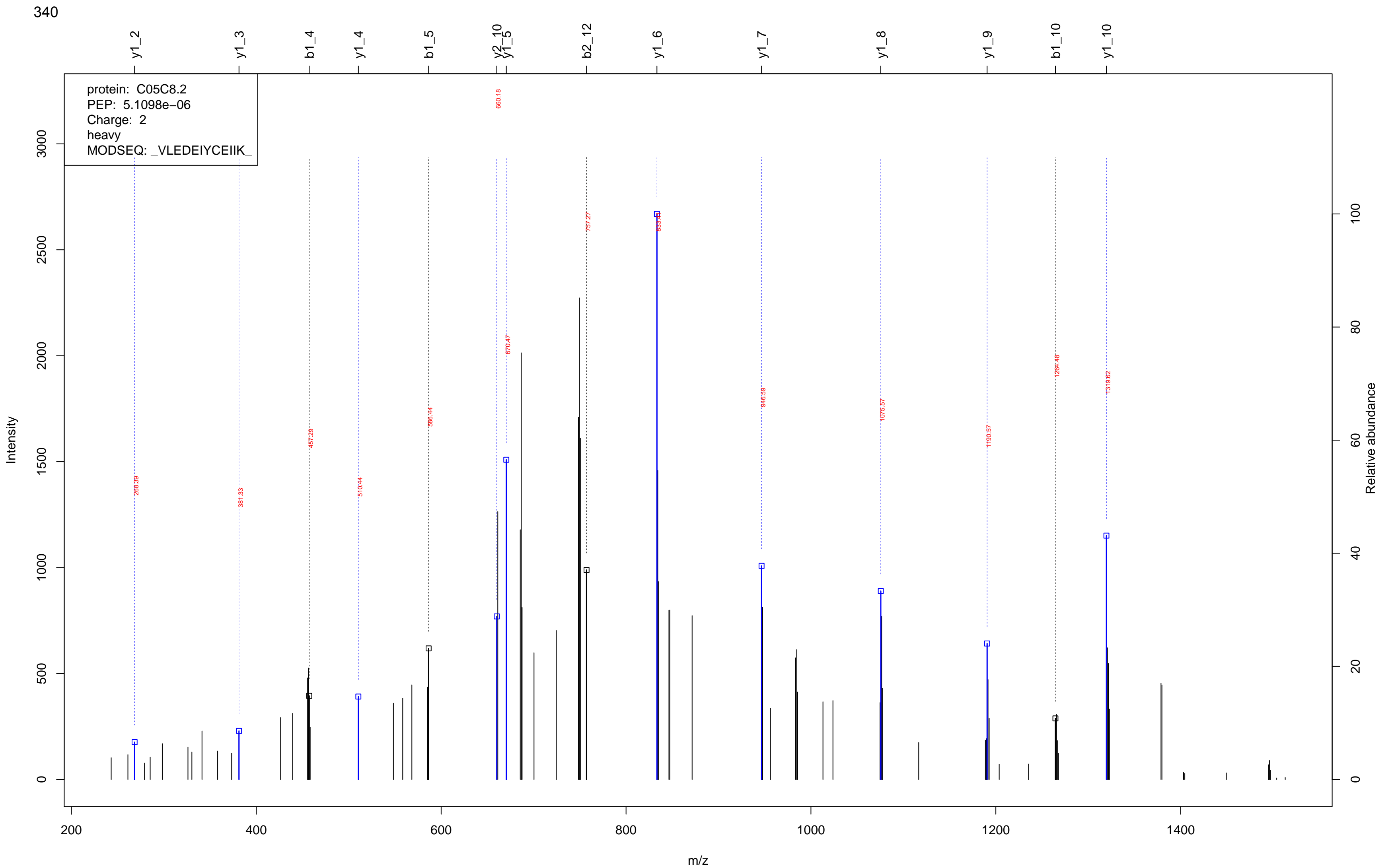






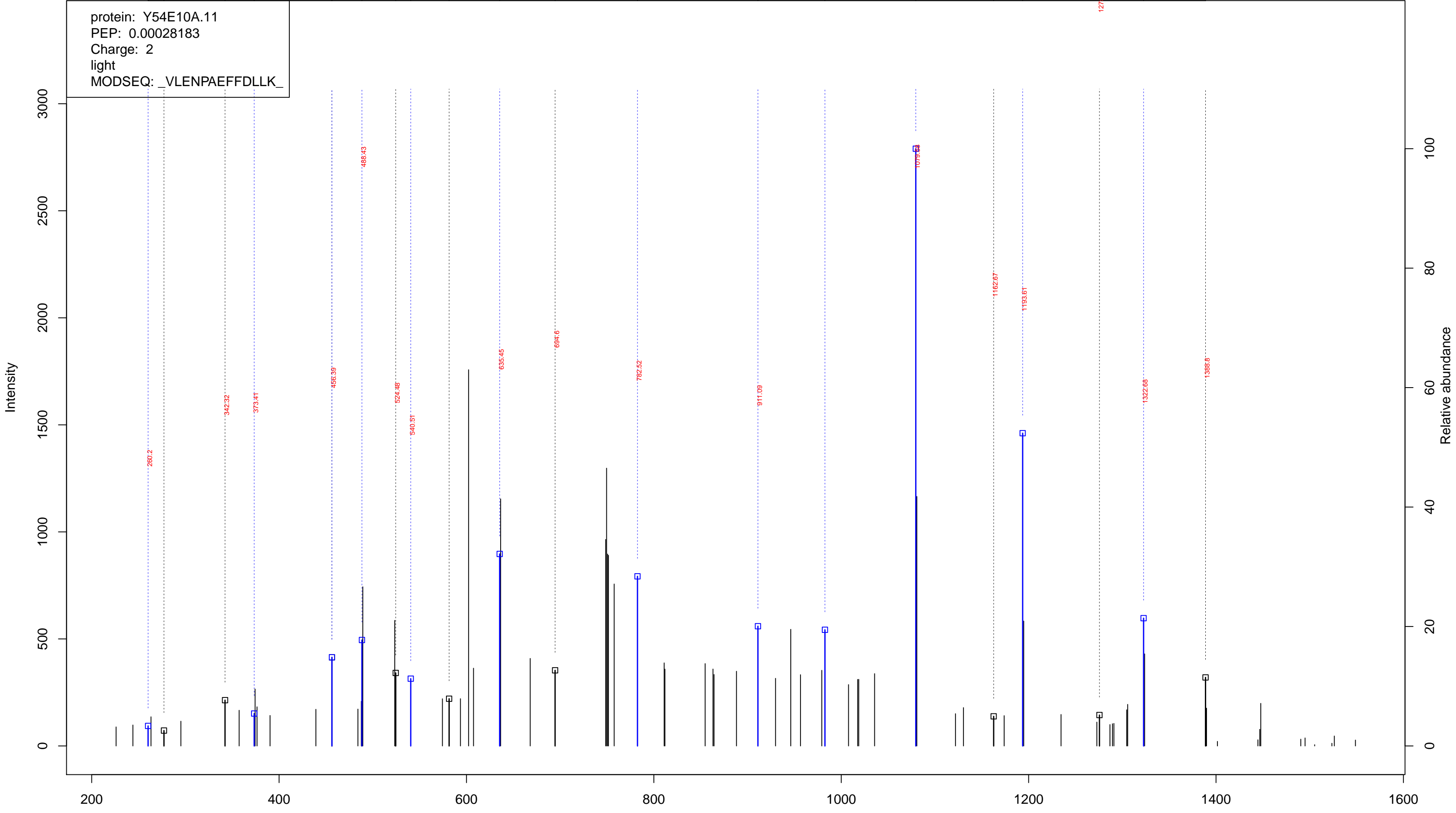


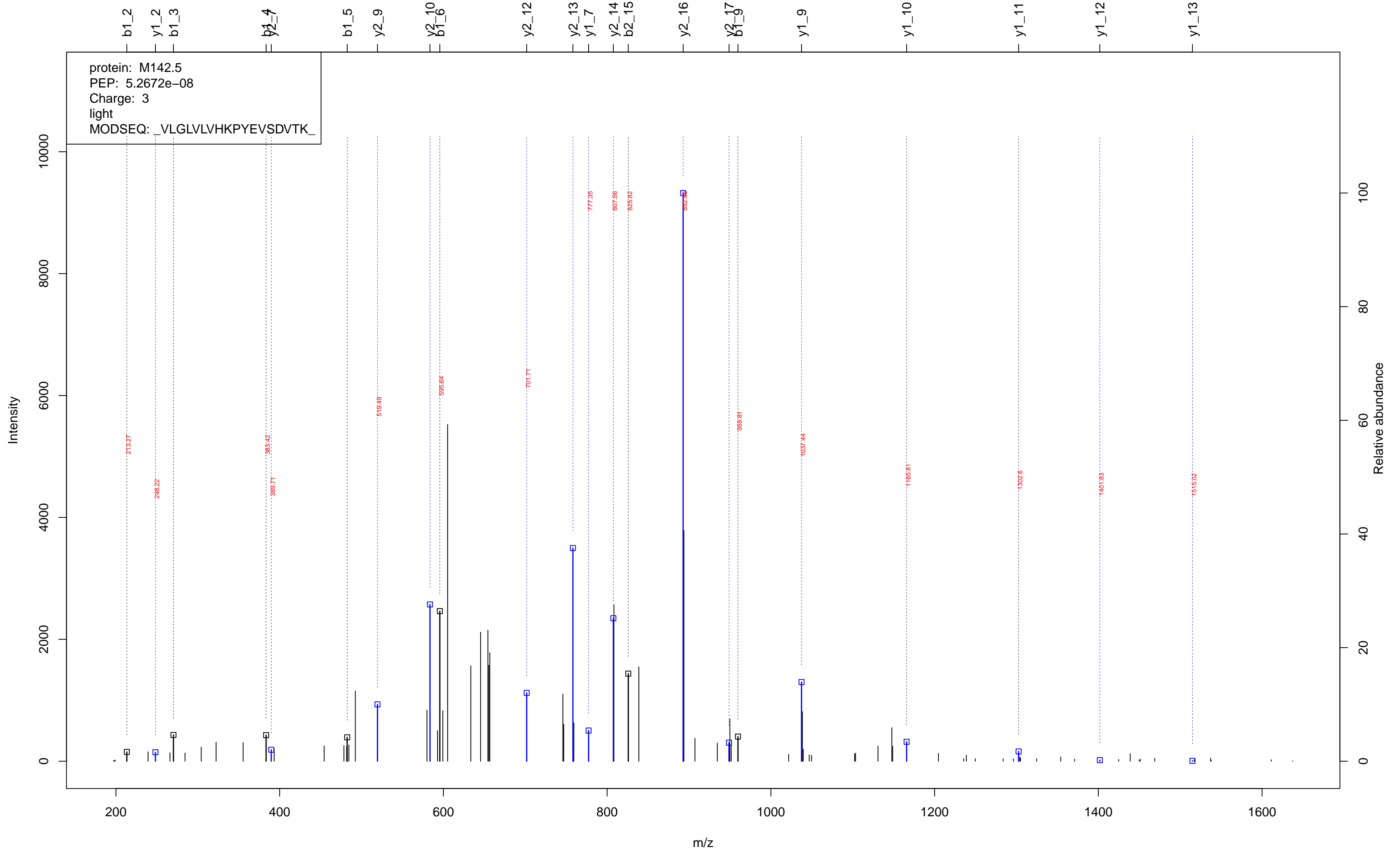


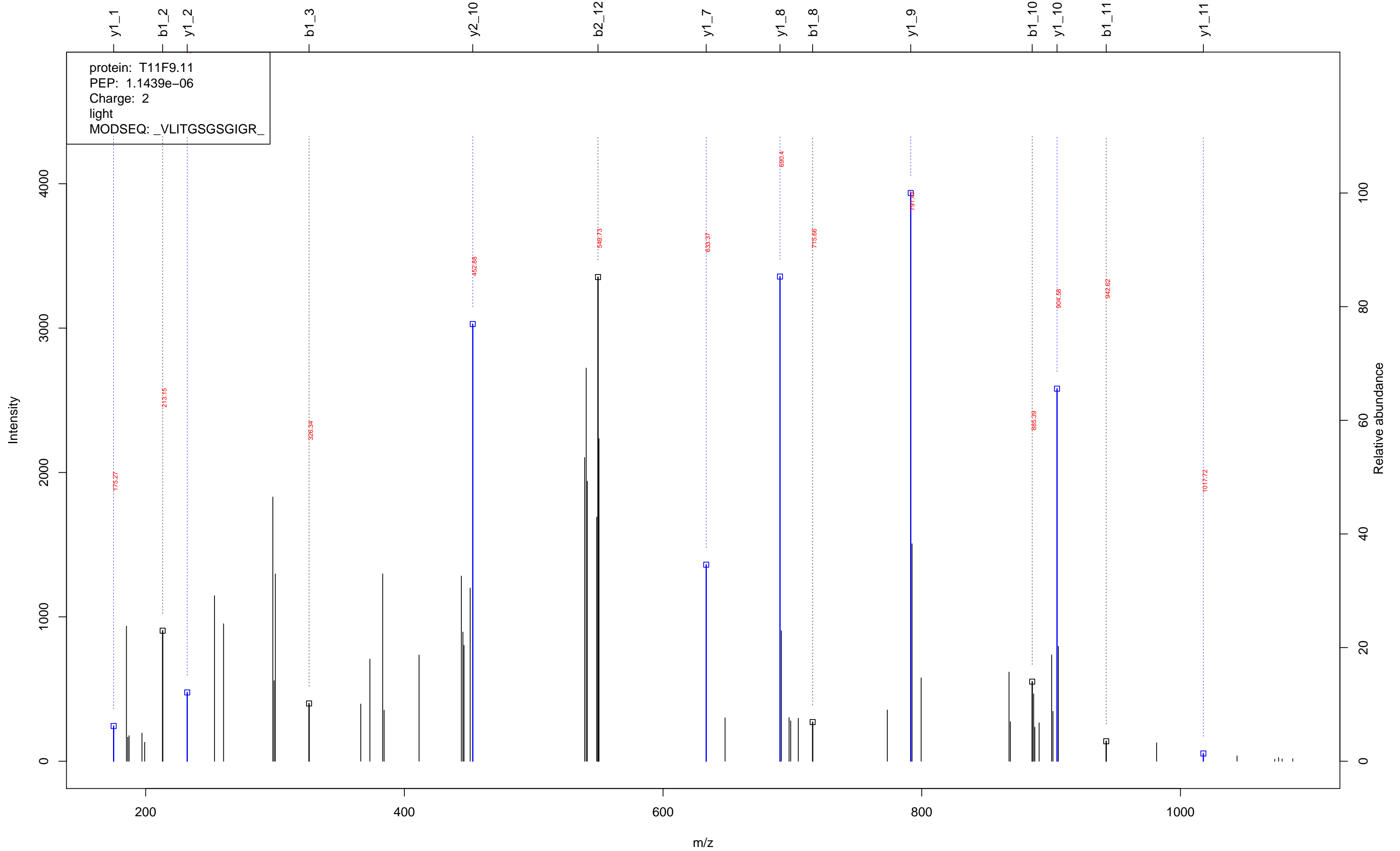


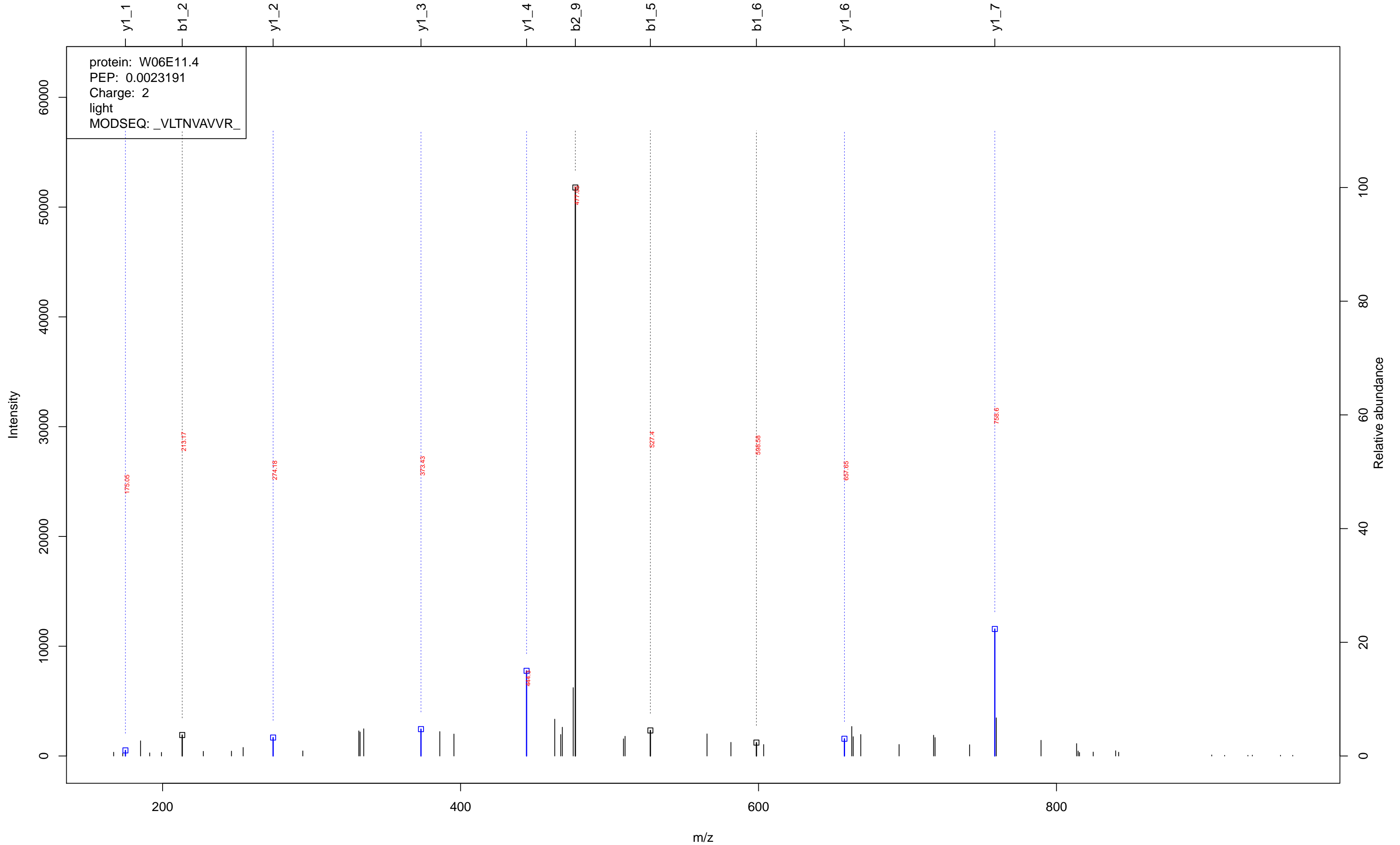
y1_2 b2_5 b1_3 y1_3 b2_7 y1_4 b2_9 y2_9 b2_10 y1_5 b2_12 y1_6 y1_7 y1_8 y1_9 b1_10 y1_10 b1_11 y1_11 b1_12

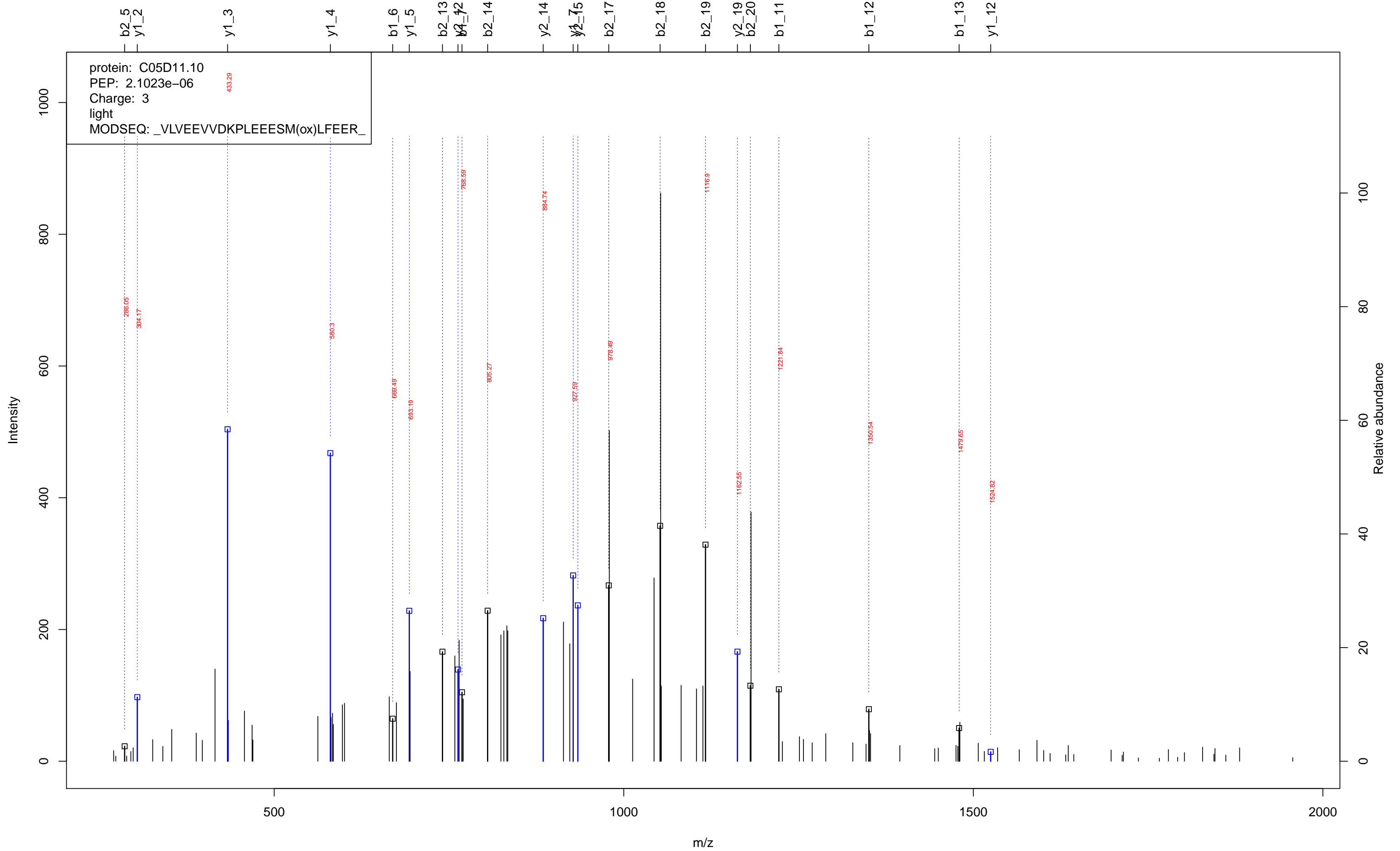
protein: Y54E10A.11
PEP: 0.00028183
Charge: 2
light
MODSEQ: _VLENPAEFFDLLK_

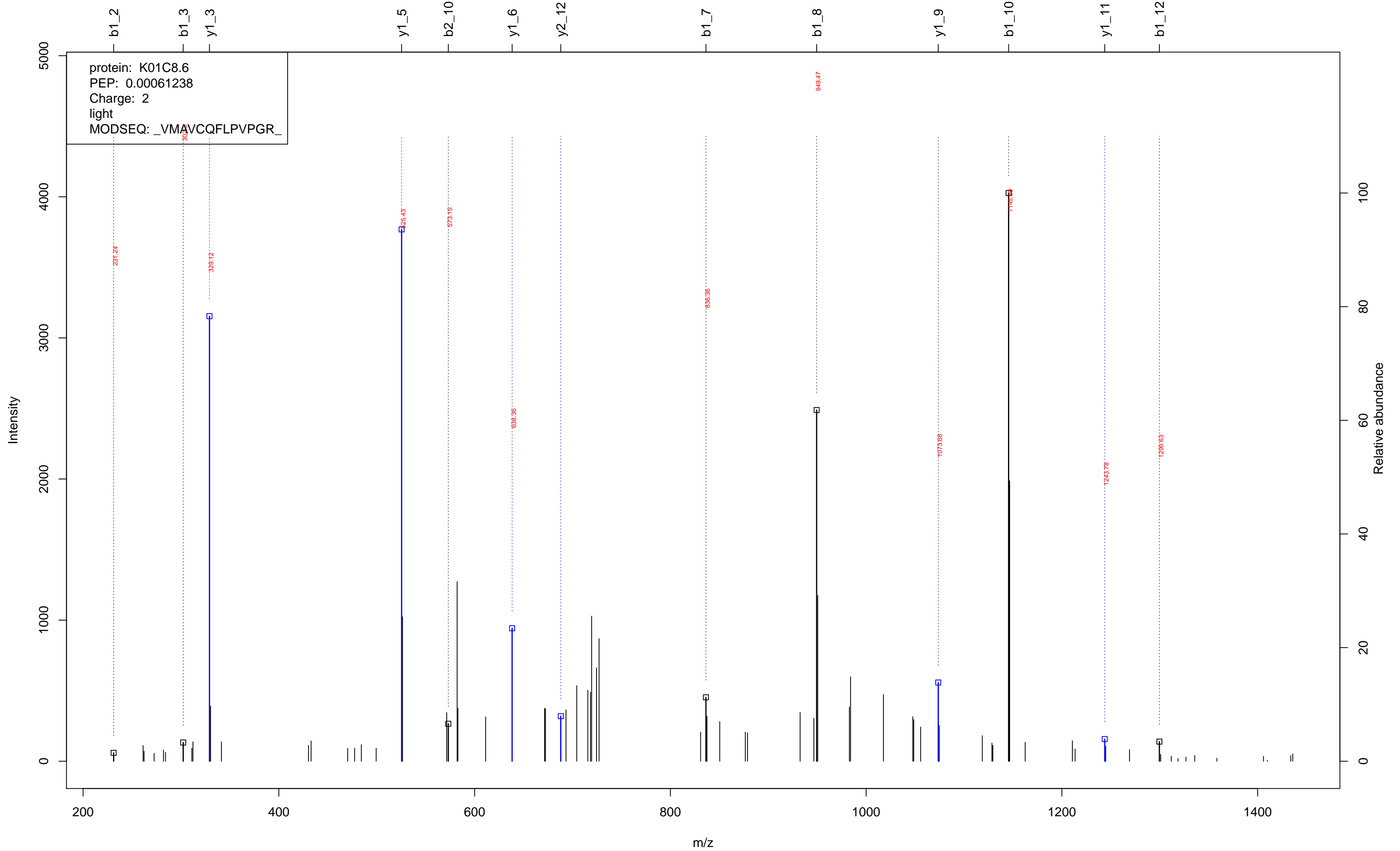


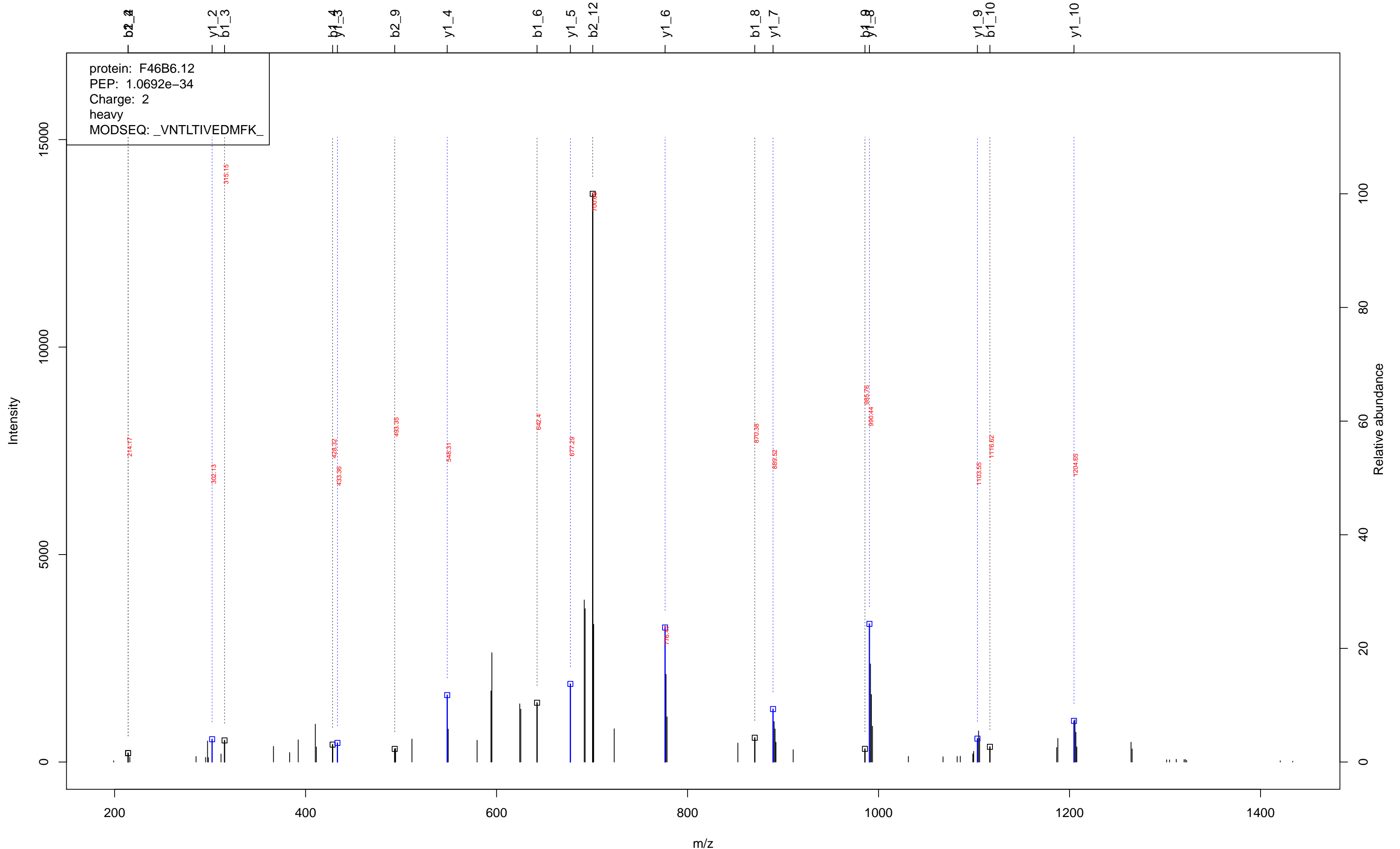


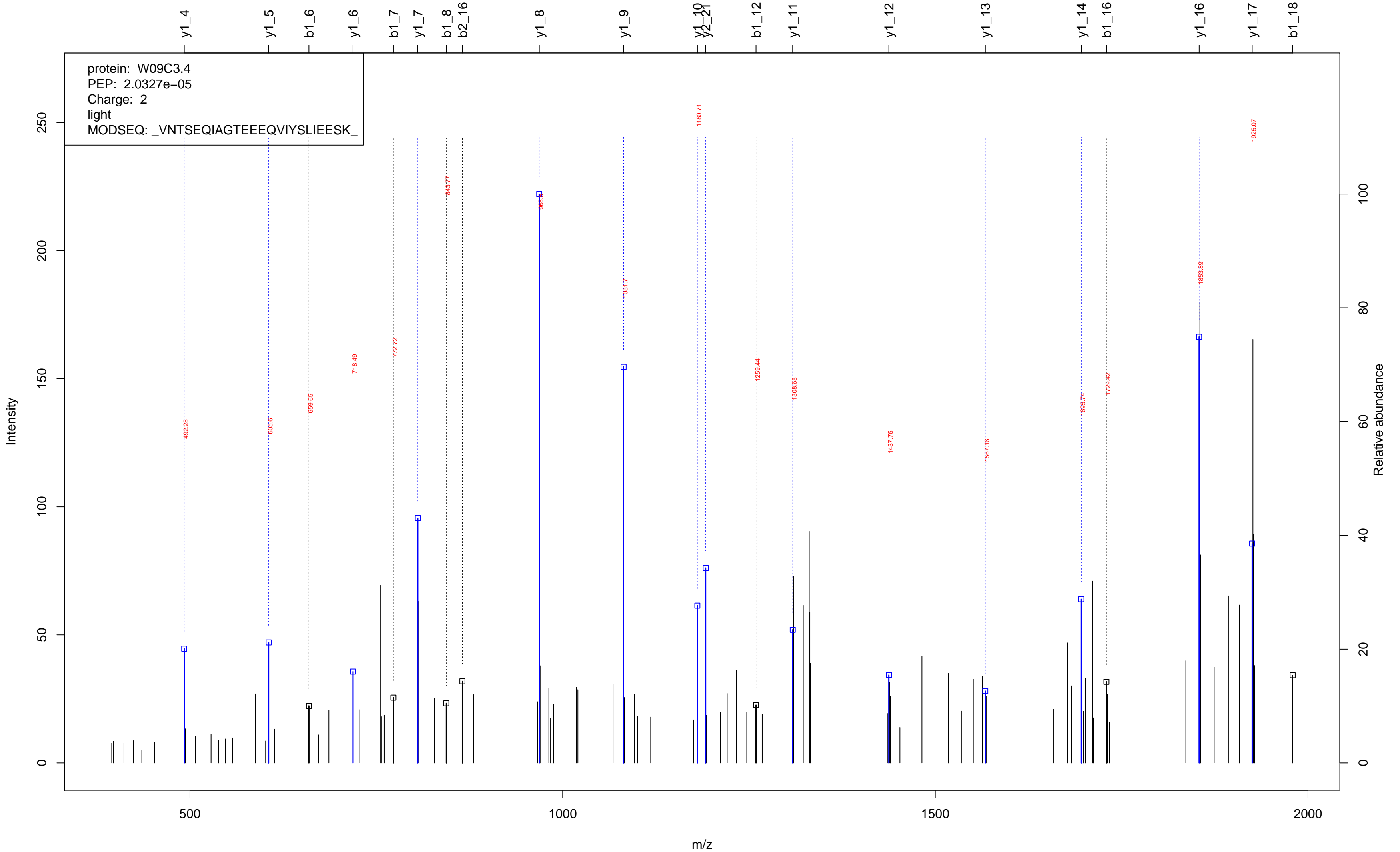


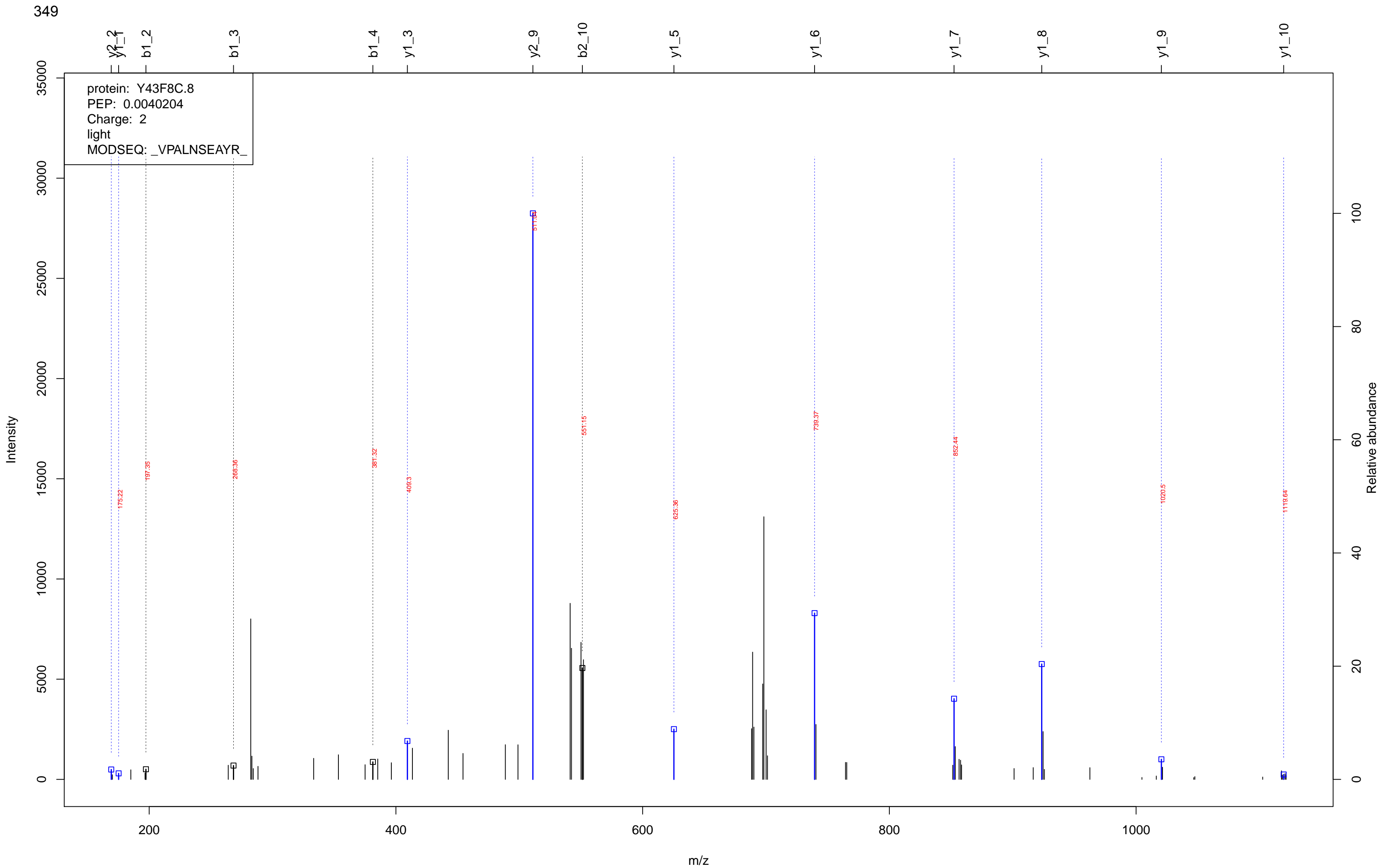


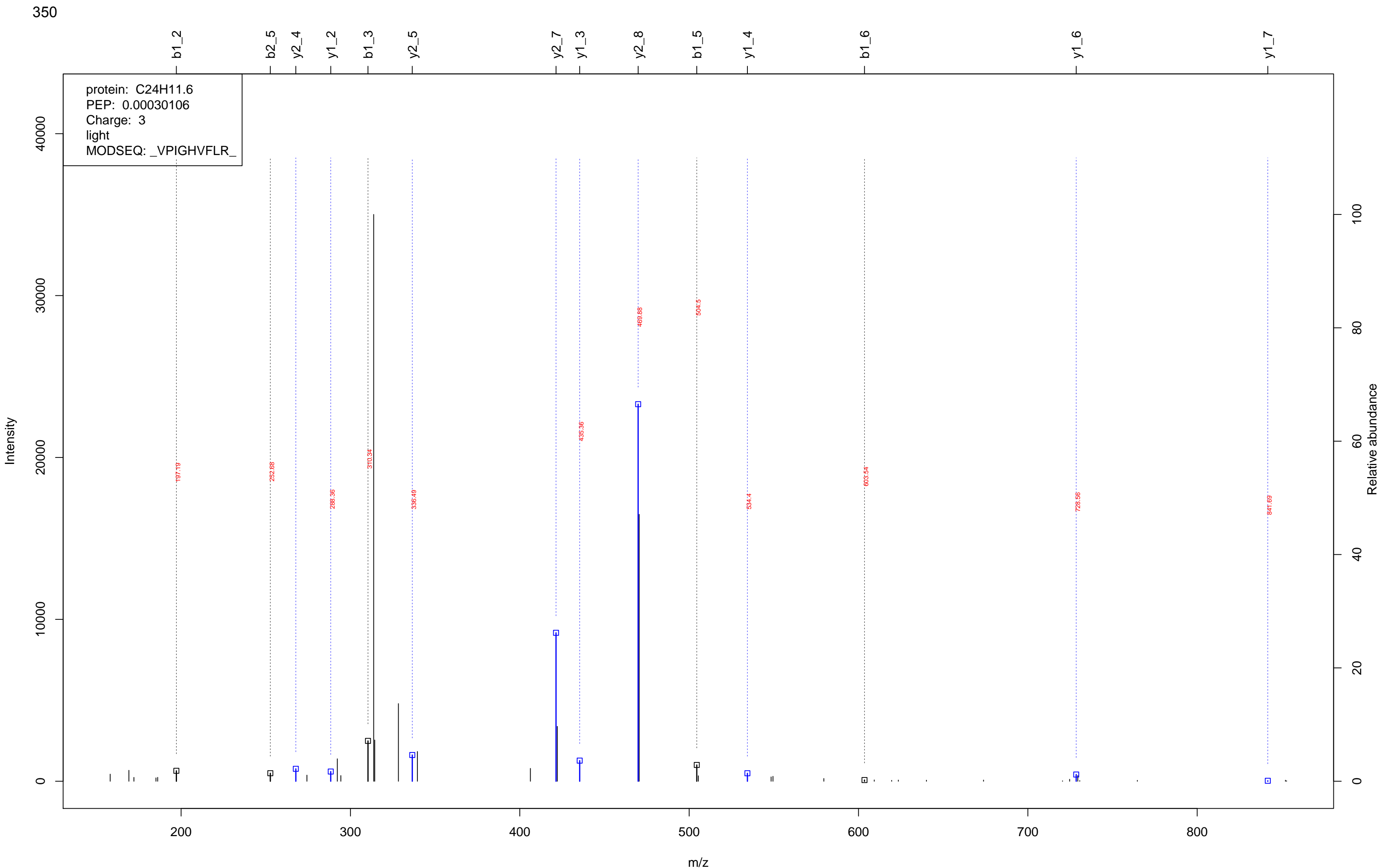


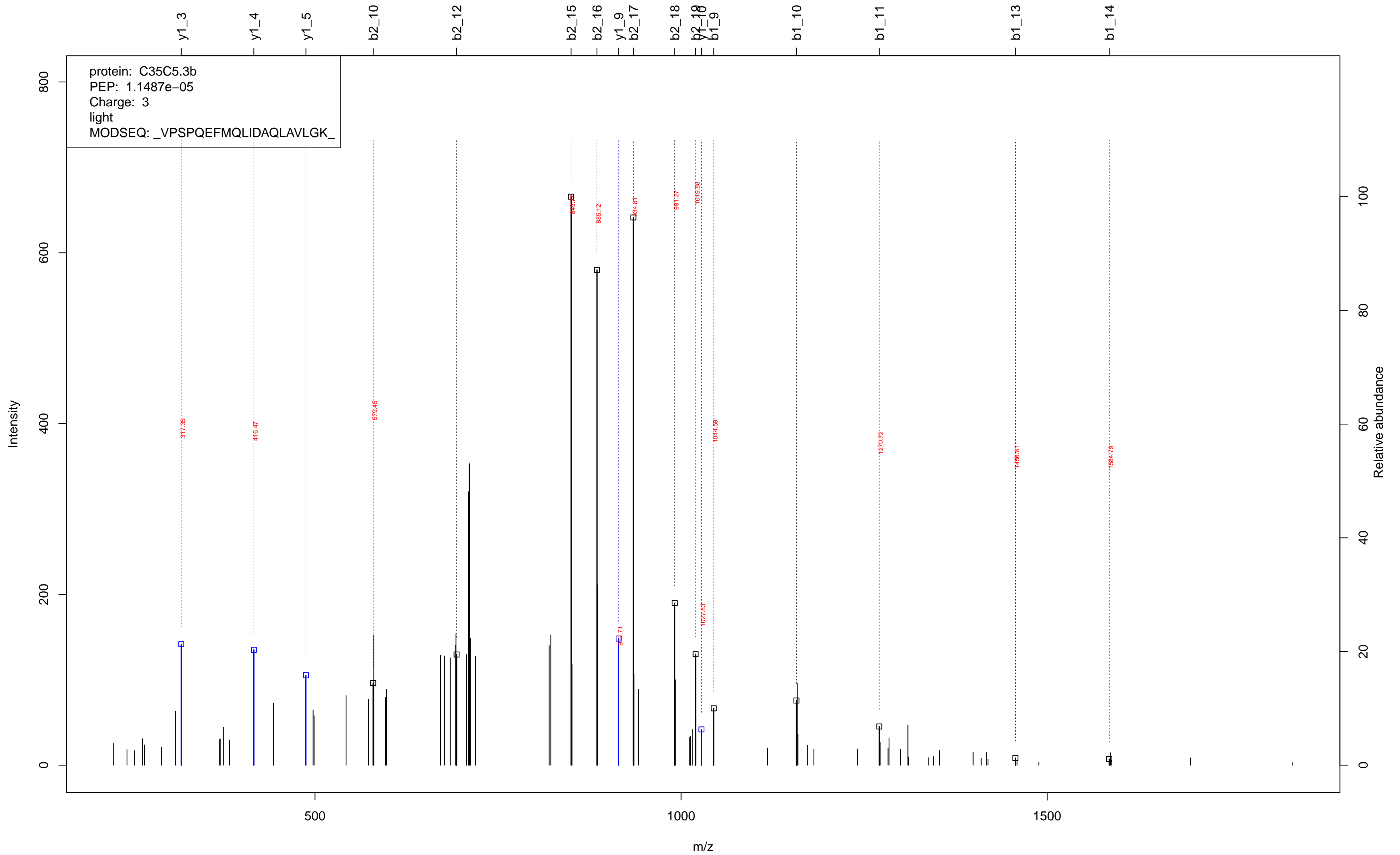


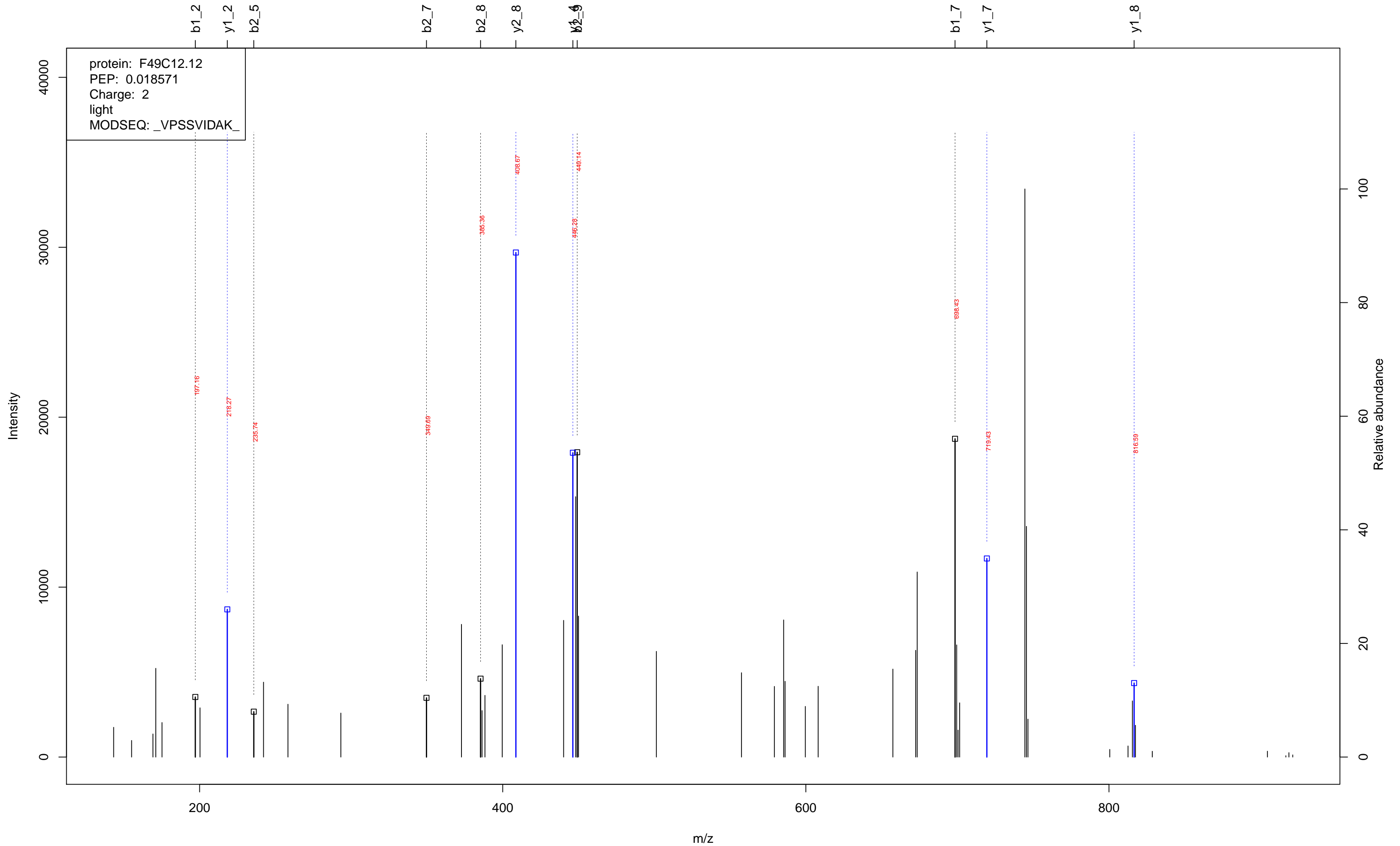


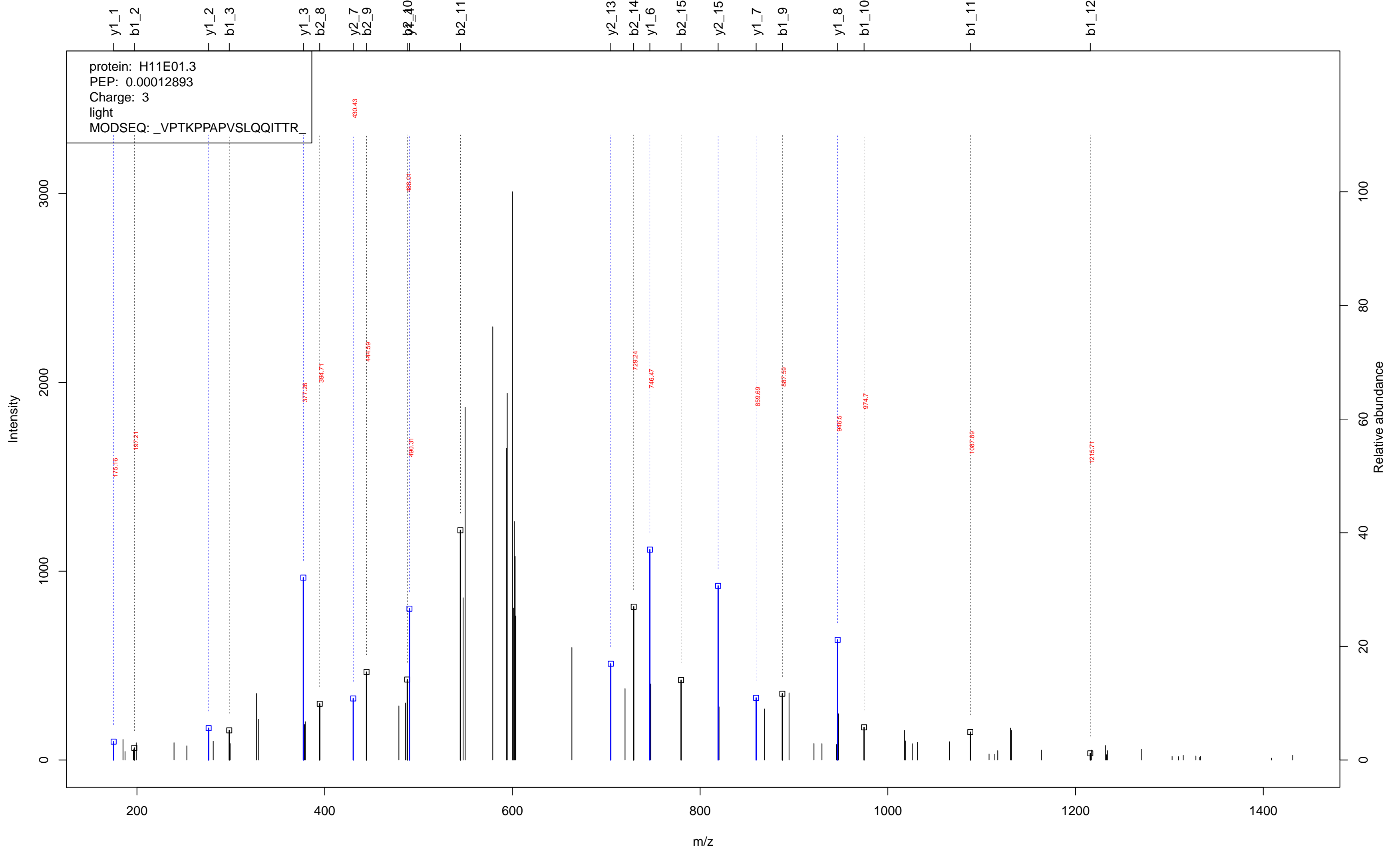


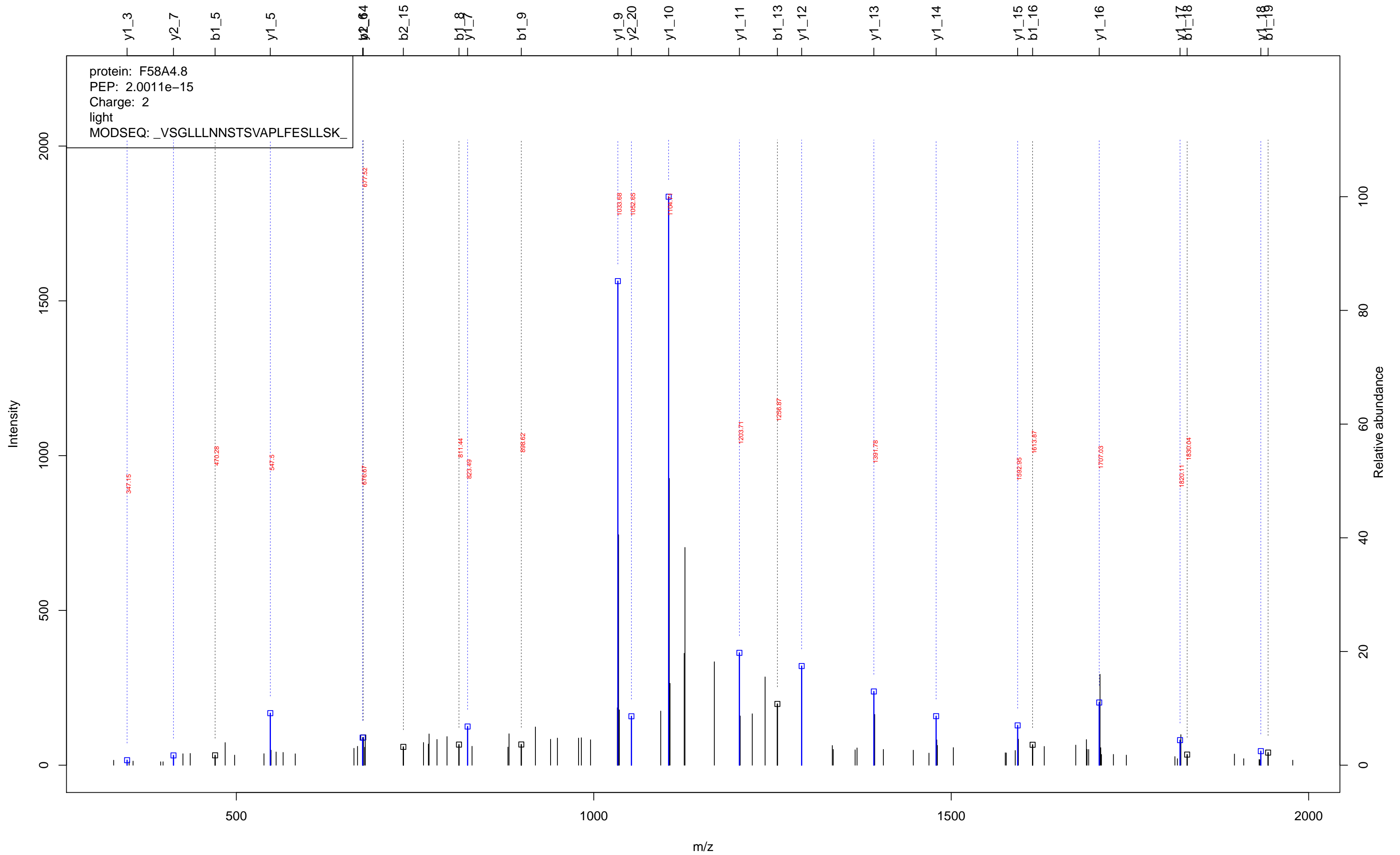


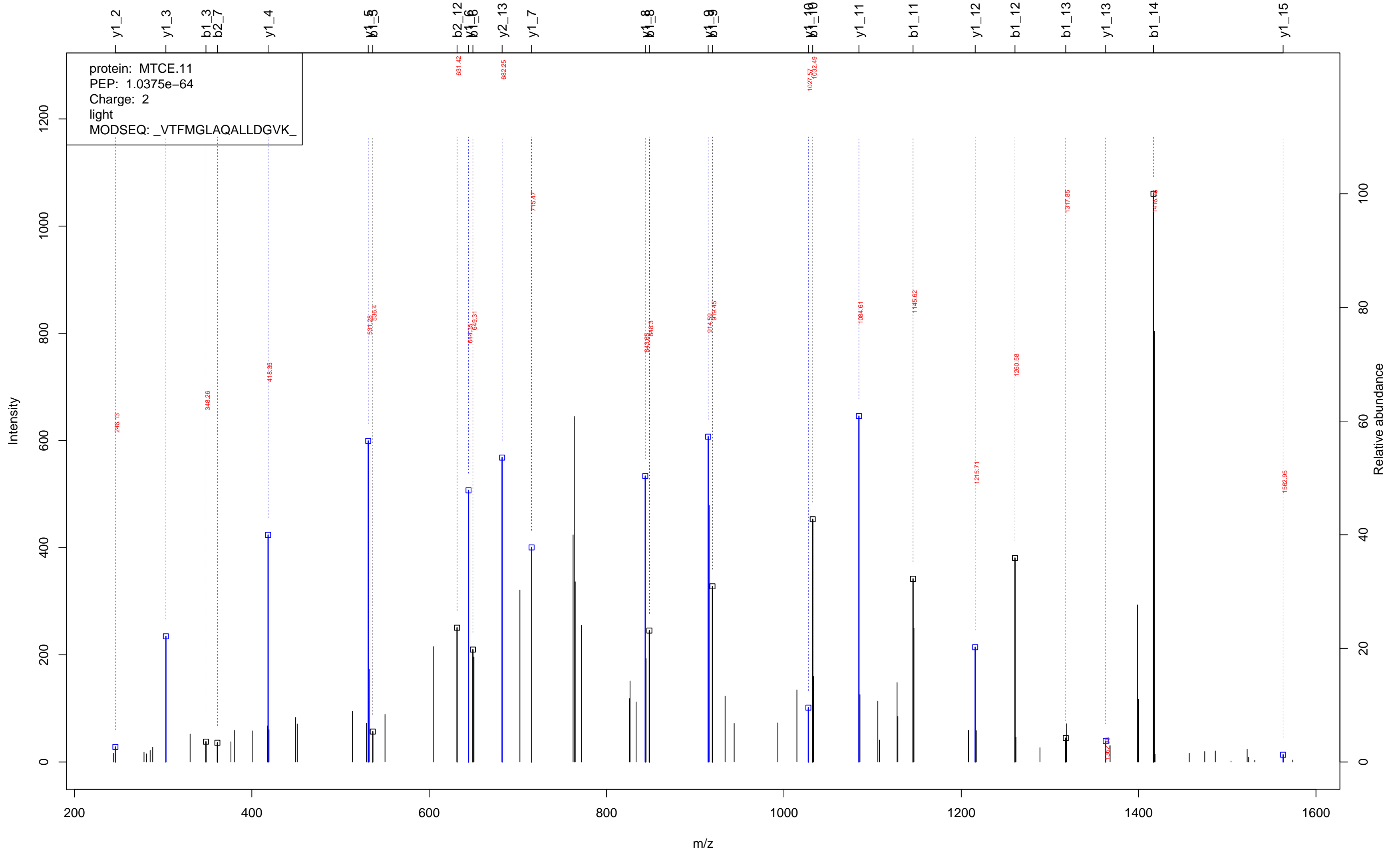


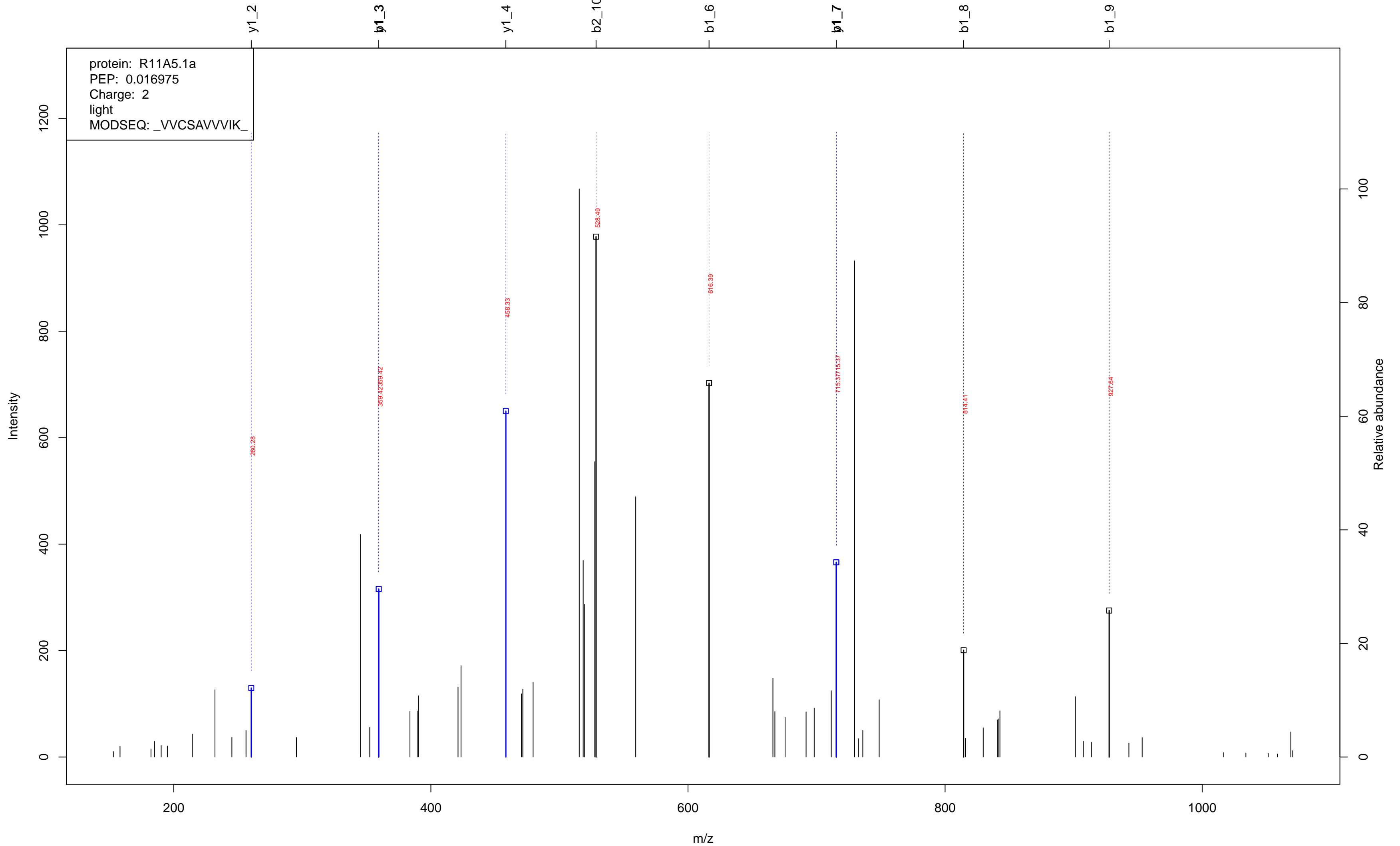


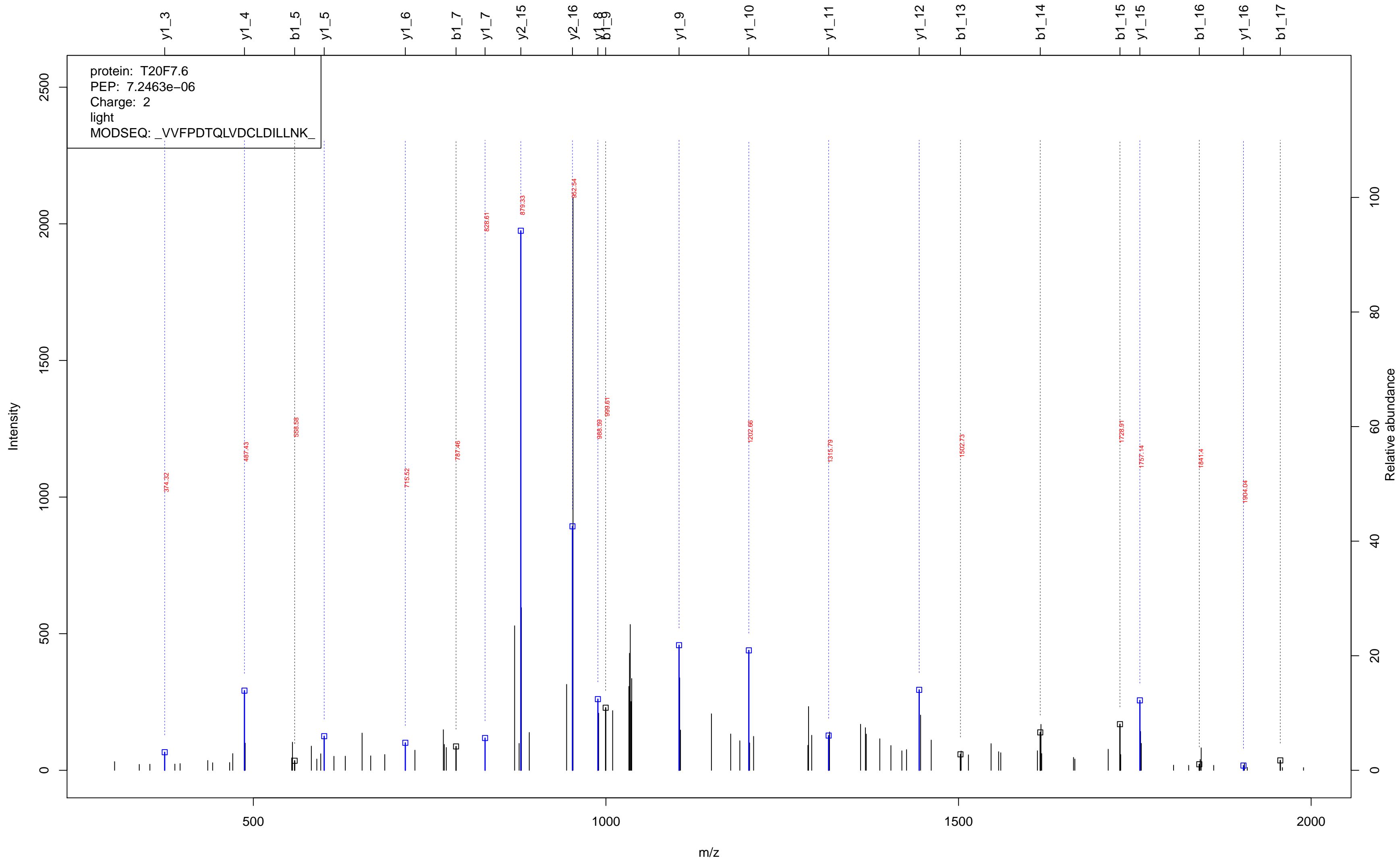


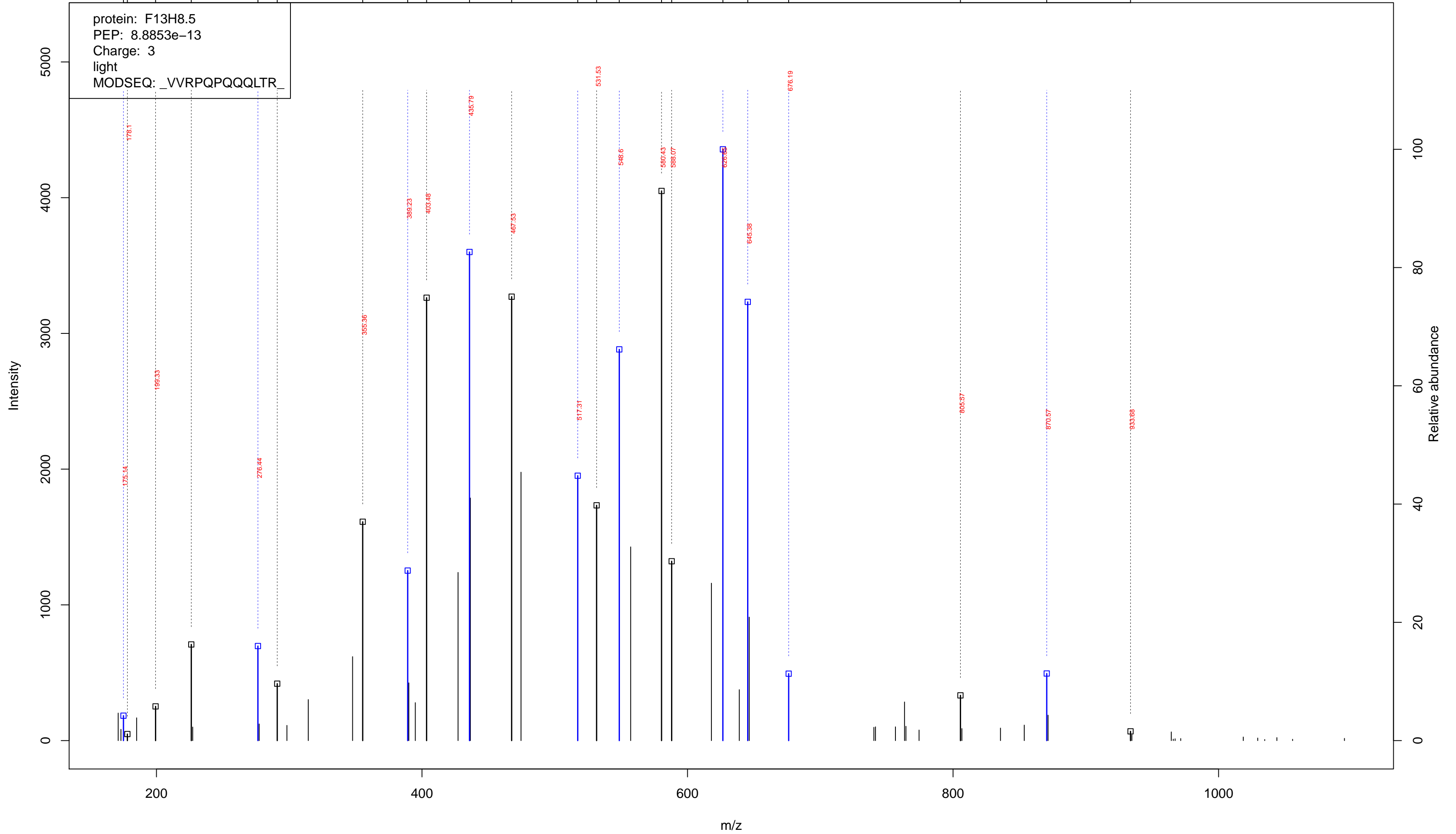


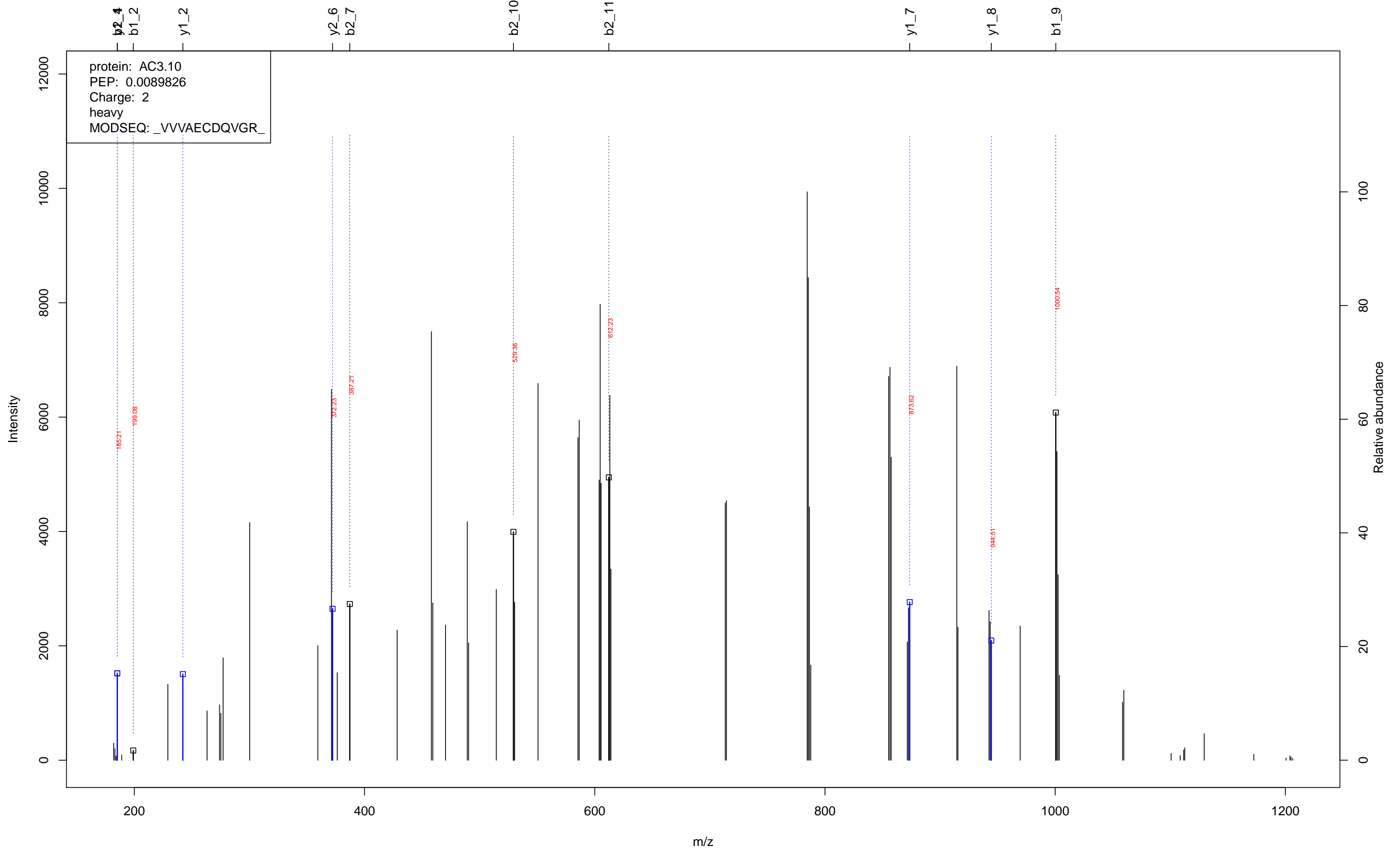


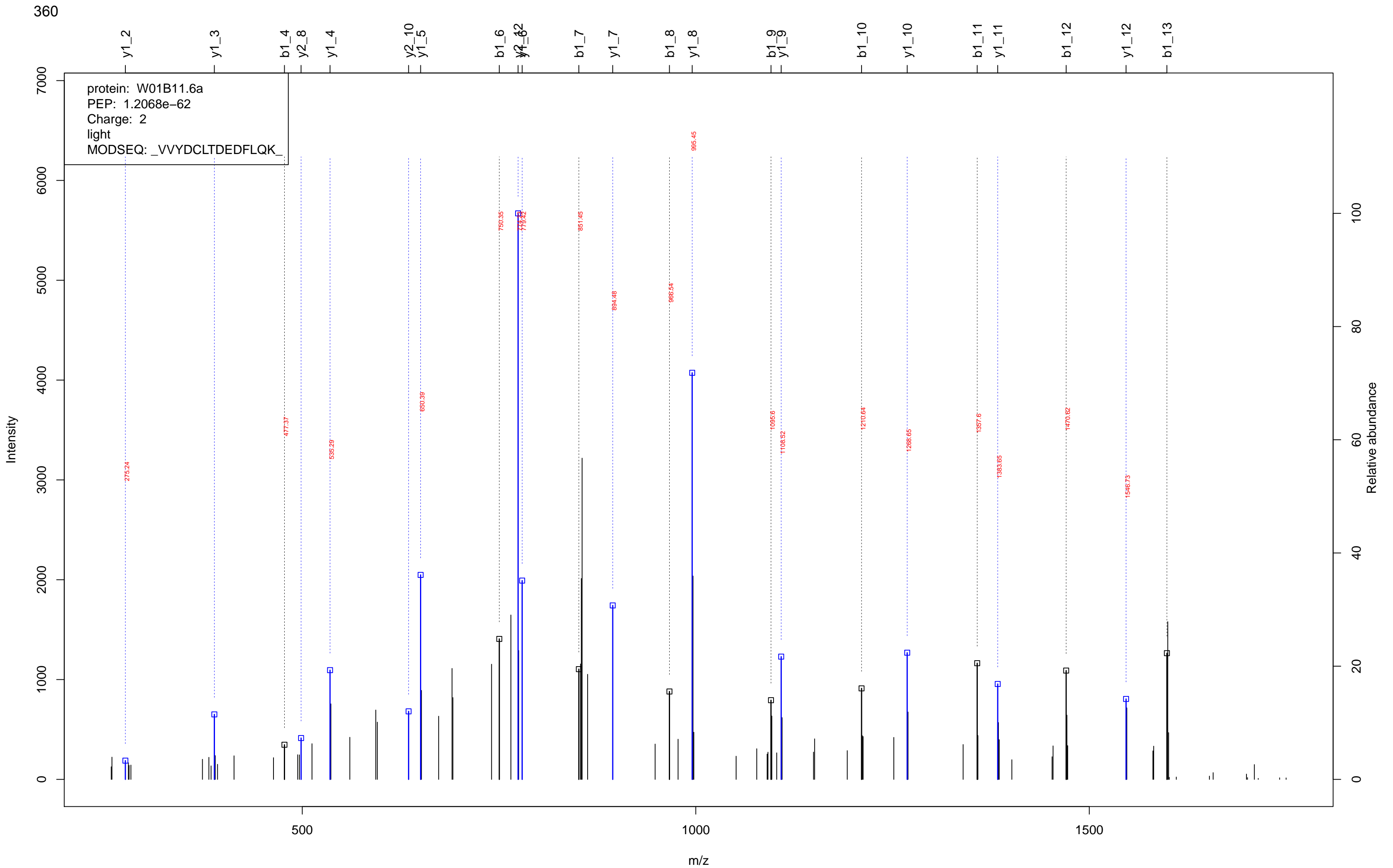


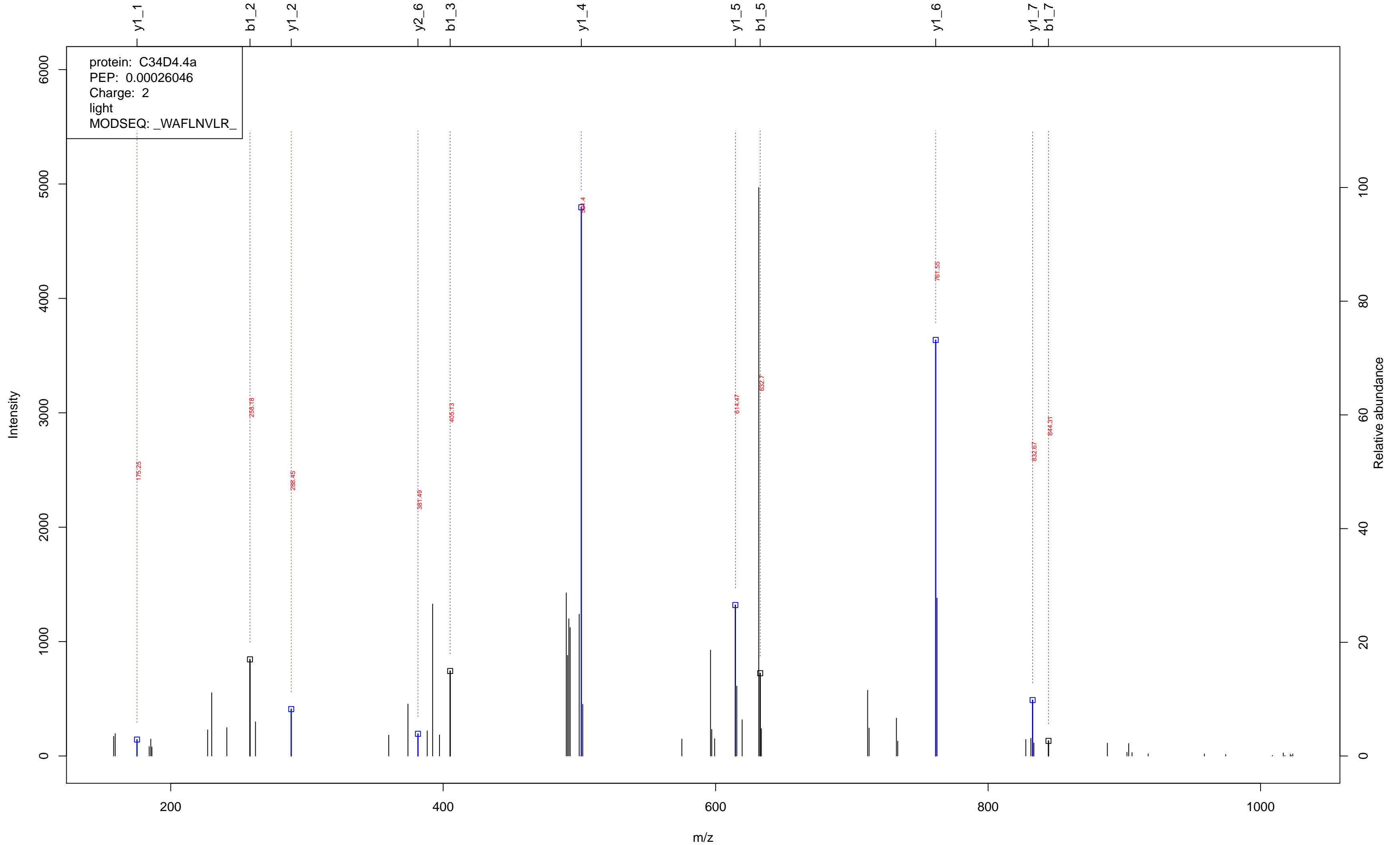


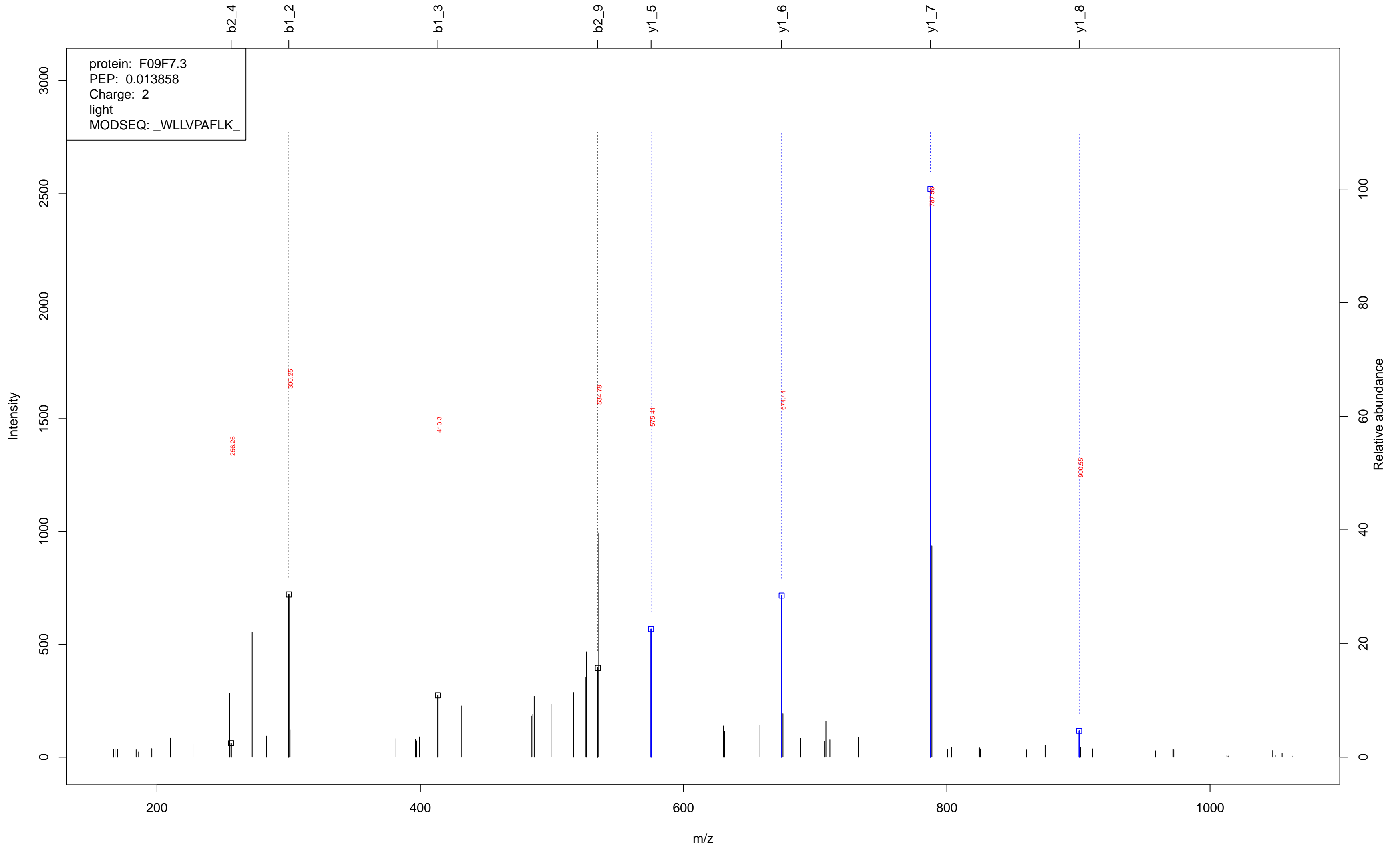


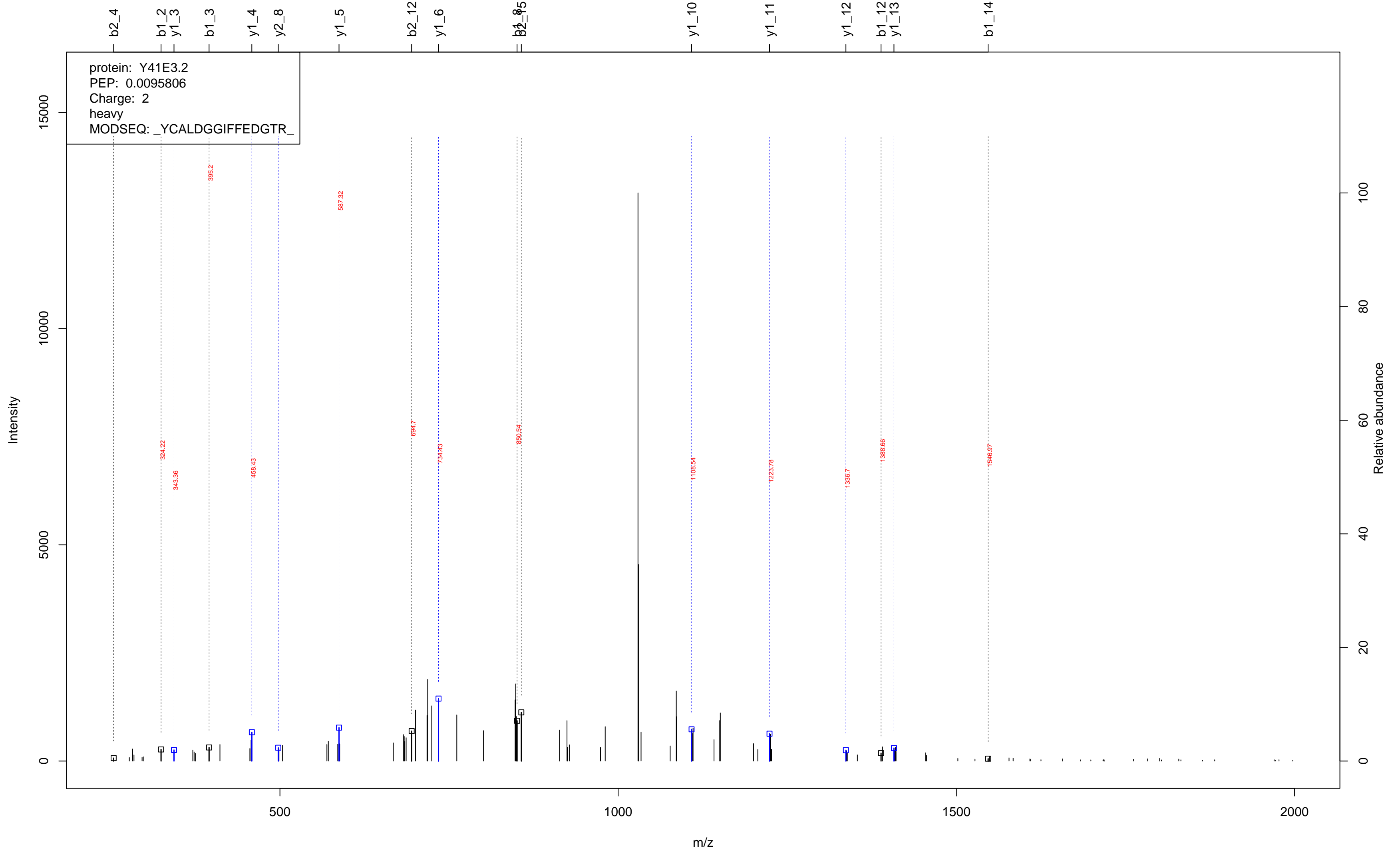






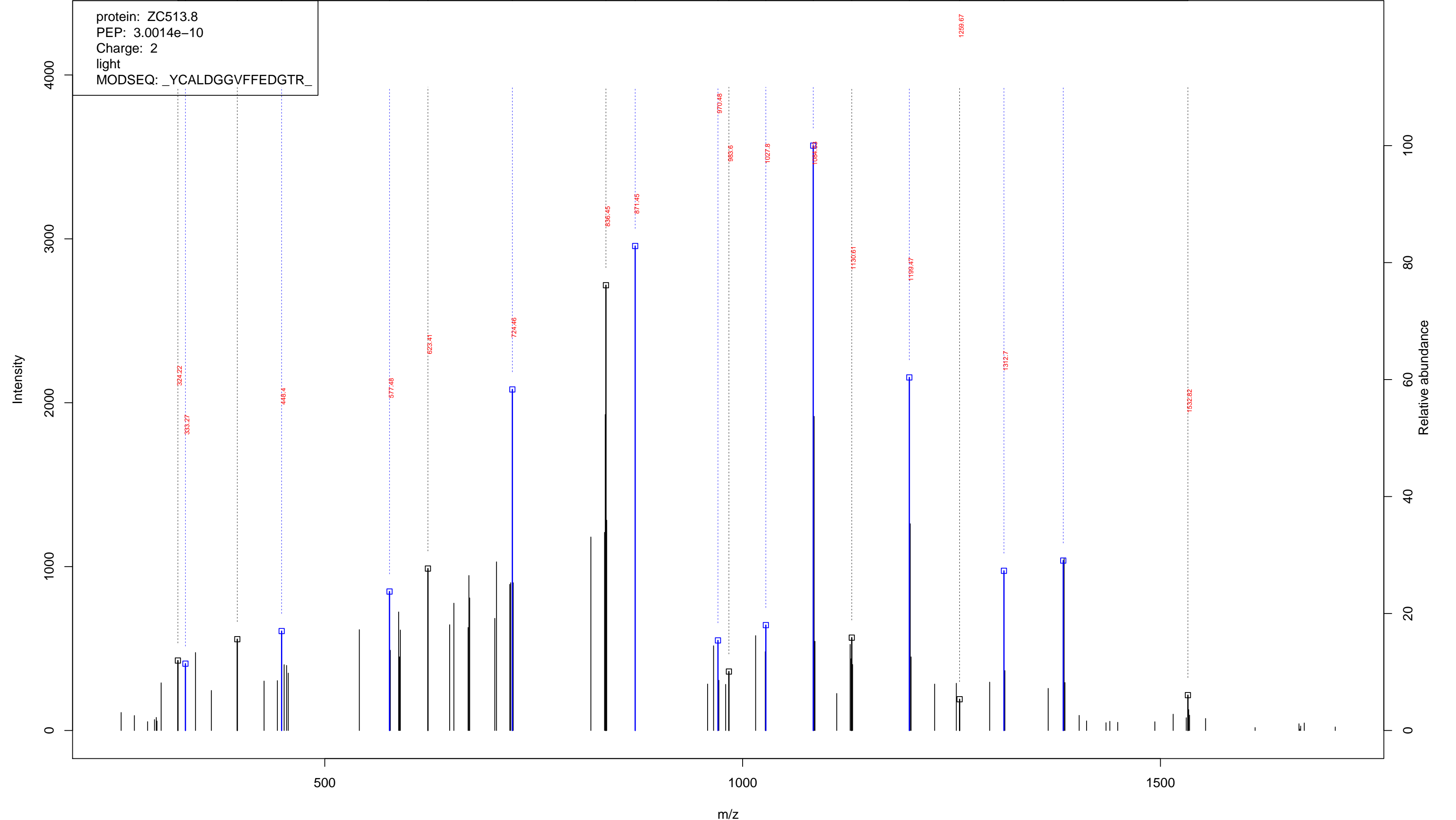


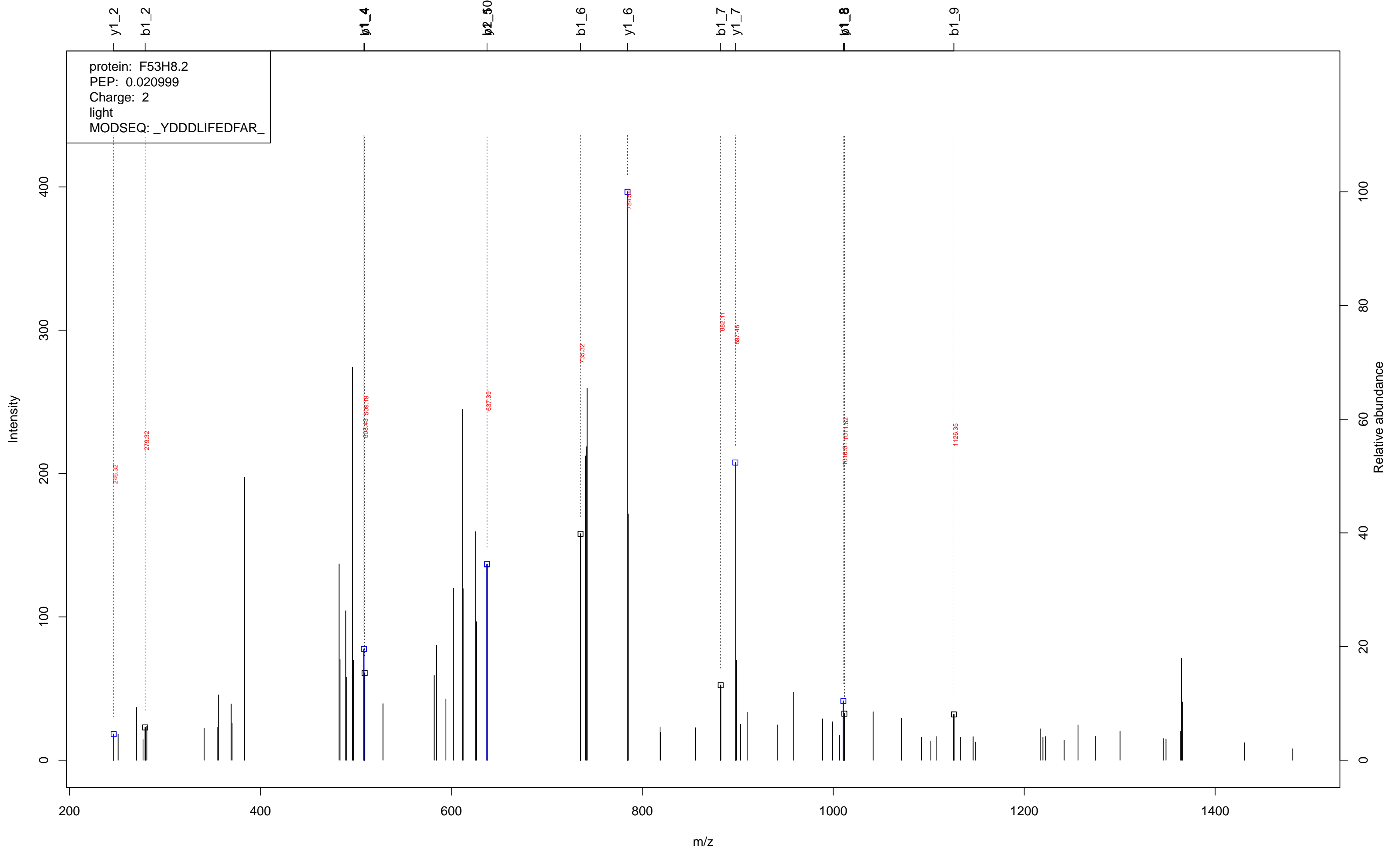


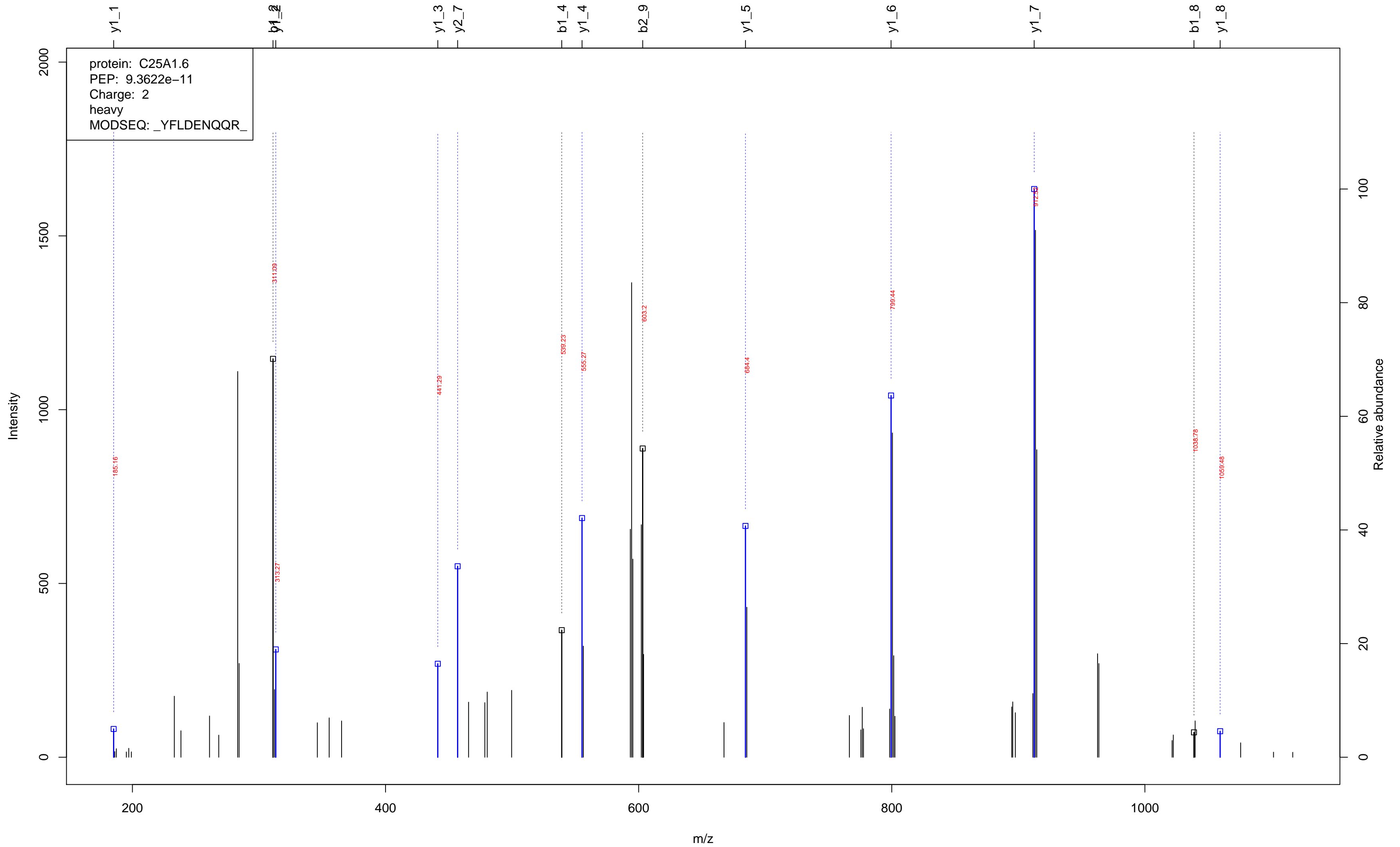


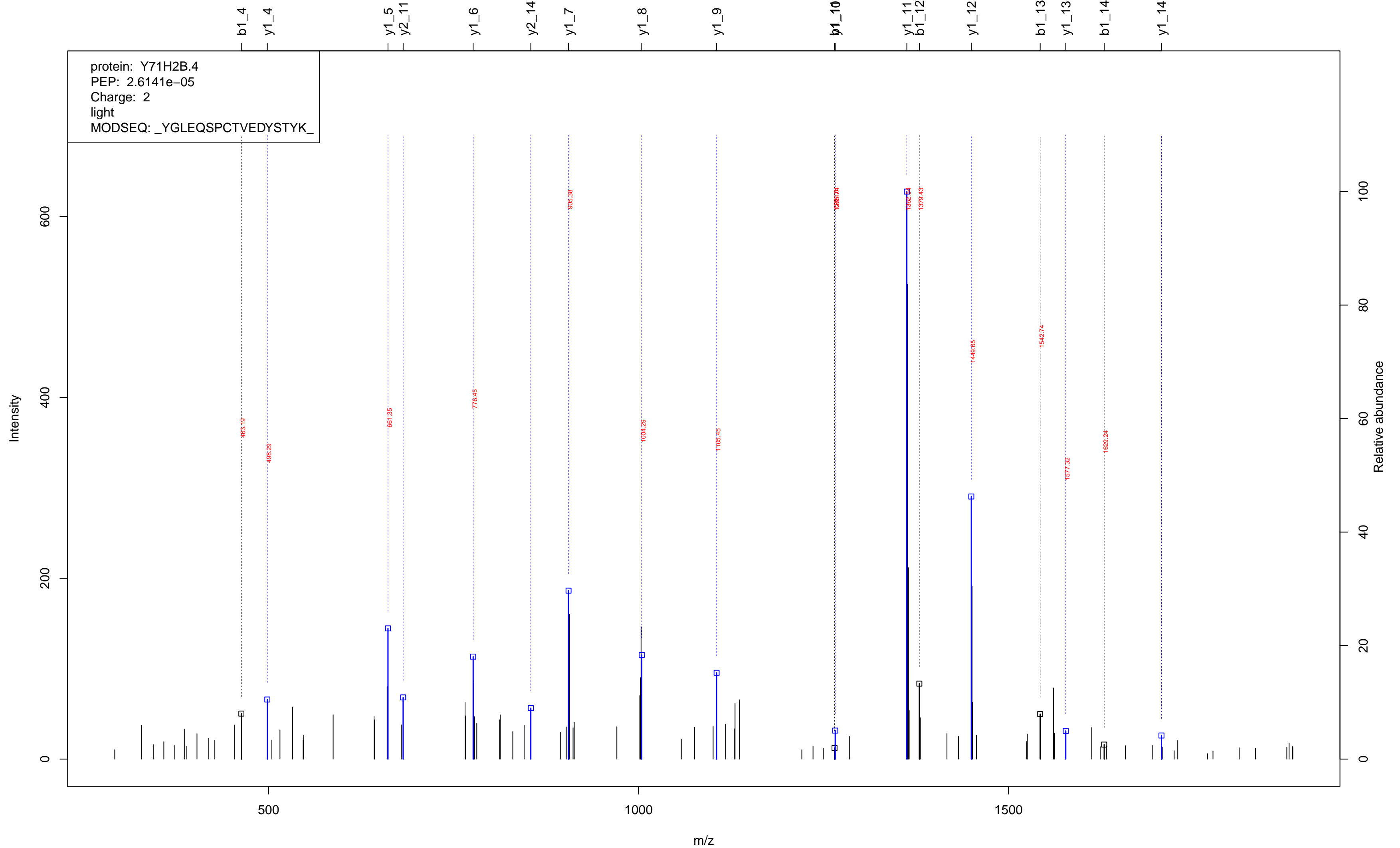
b1_3 y1_3 b1_3 y1_4 y1_5 b1_5 y1_6 b1_8 y1_7 y1_8 b1_9 y1_9 y1_10 b1_10 y1_11 b1_11 y1_12 y1_13 b1_14

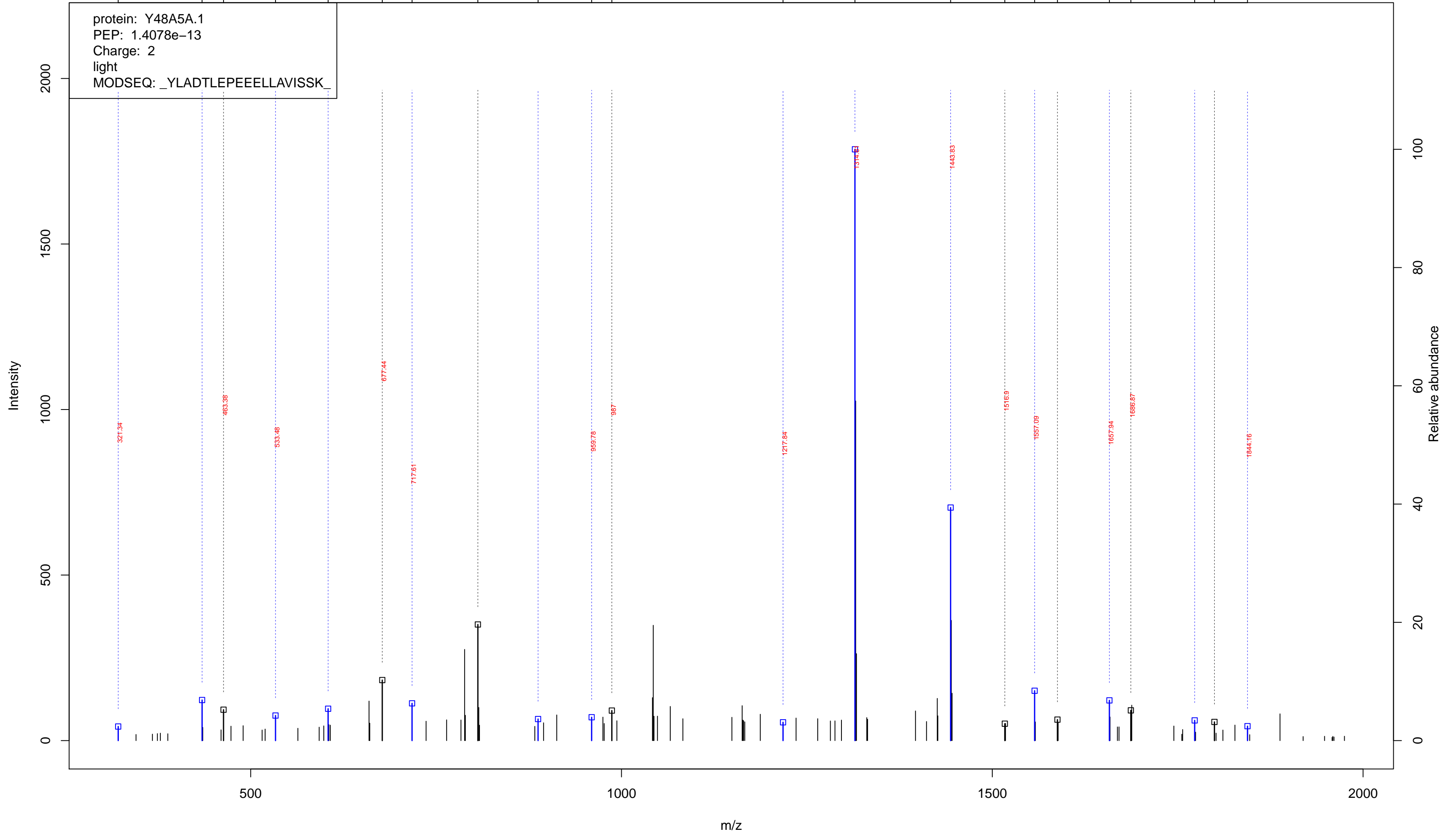
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PEP: 3.0014e-10
Charge: 2
light
MODSEQ: _YCALDGGVFFEDGTR_

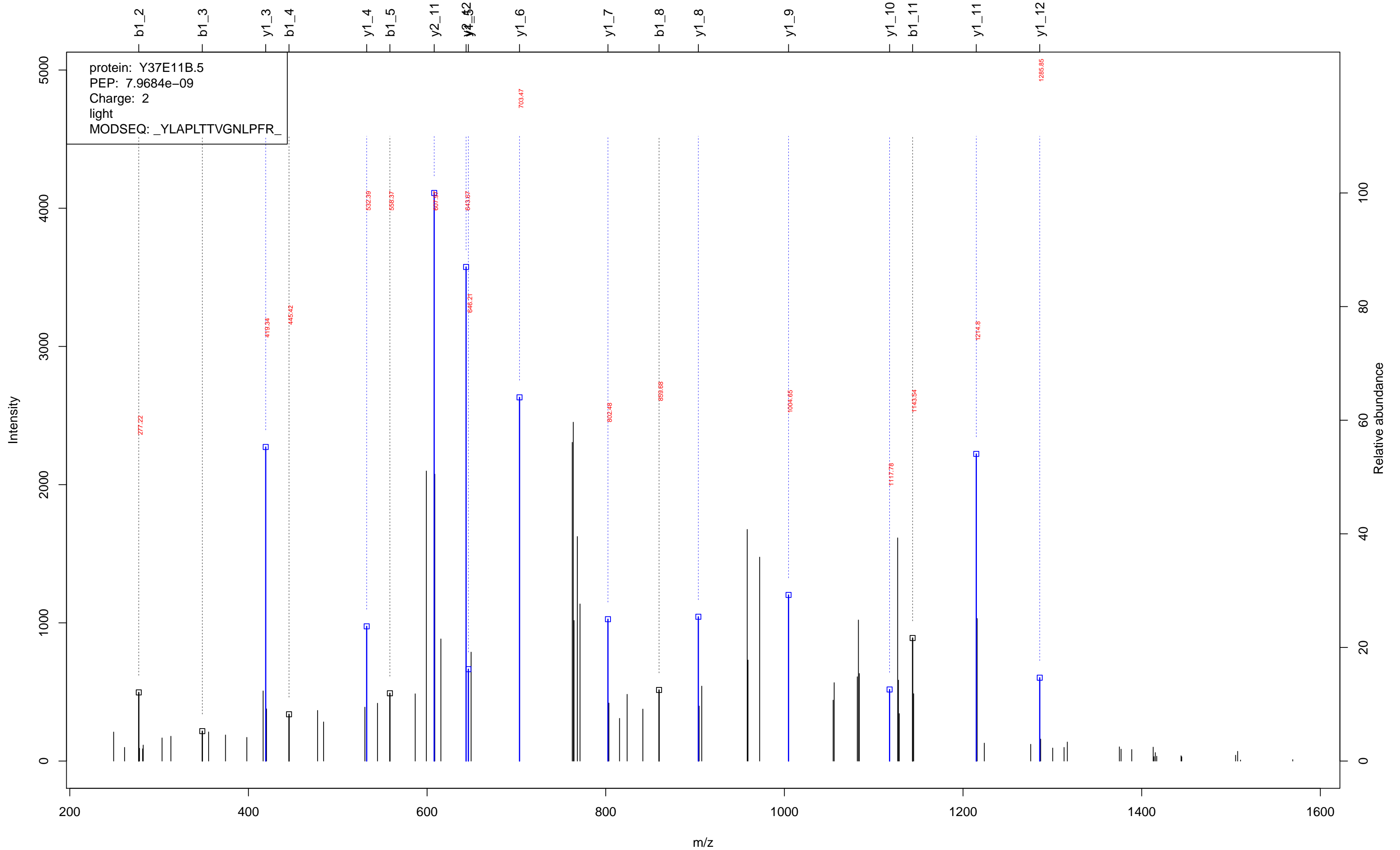






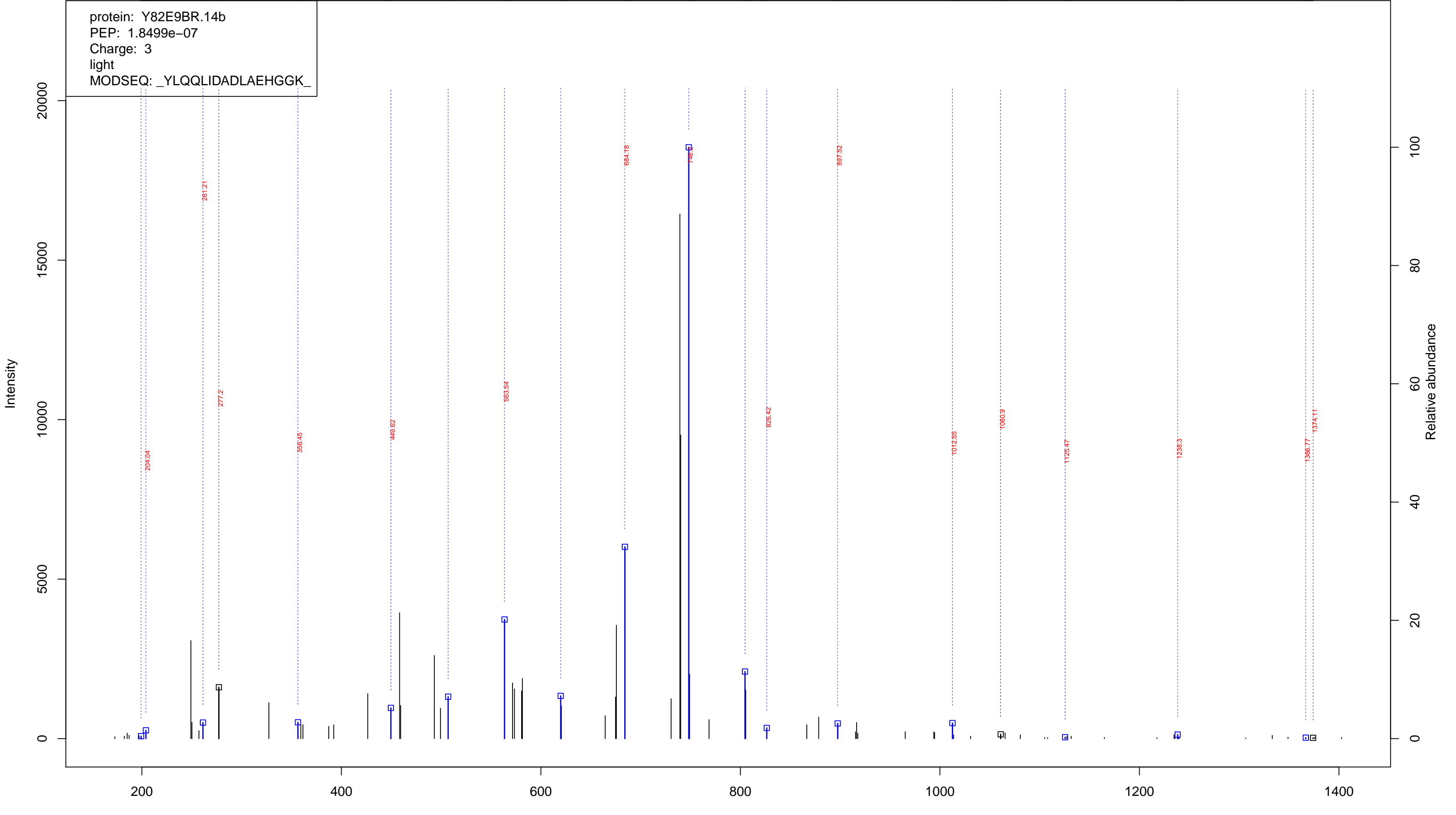


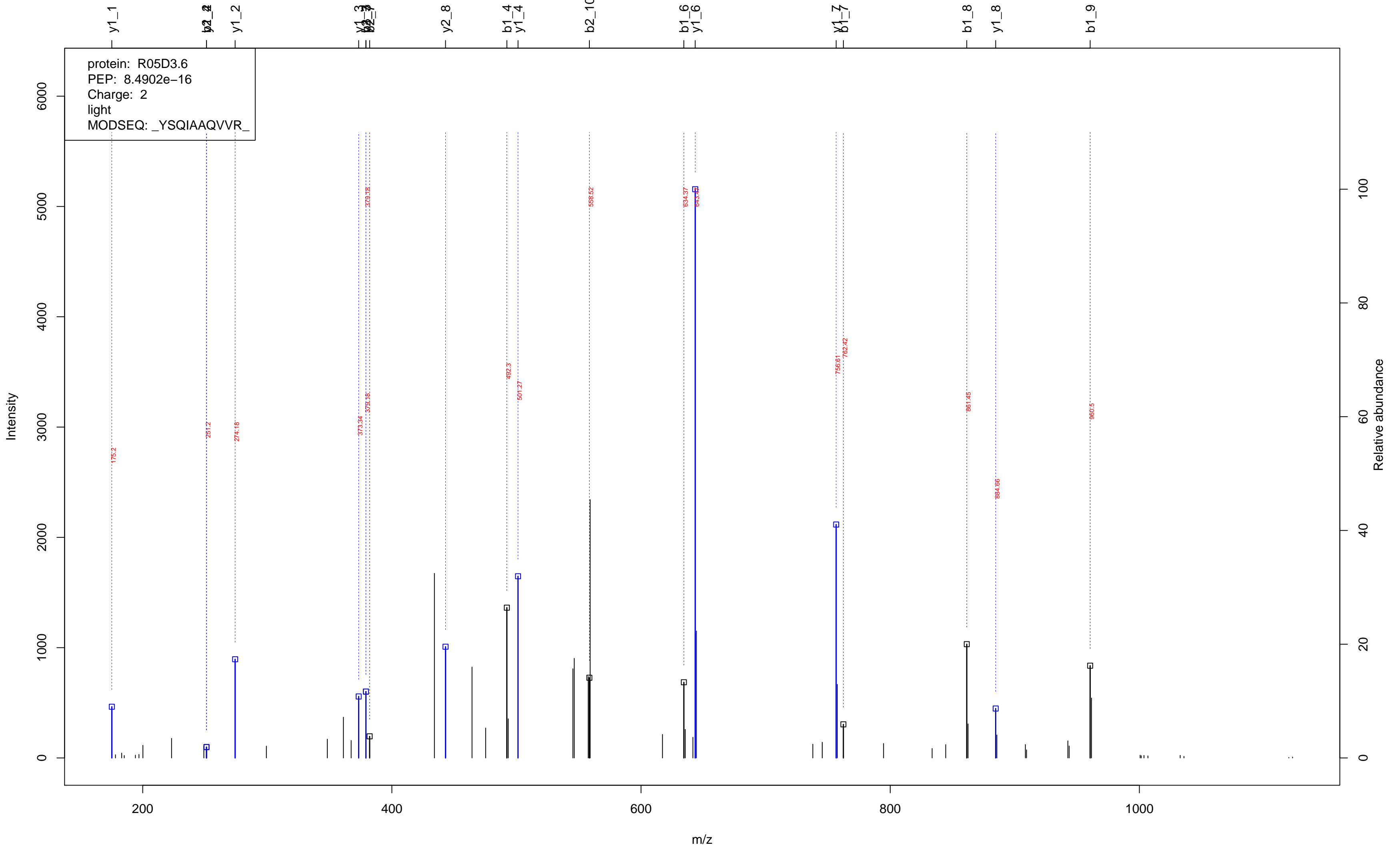


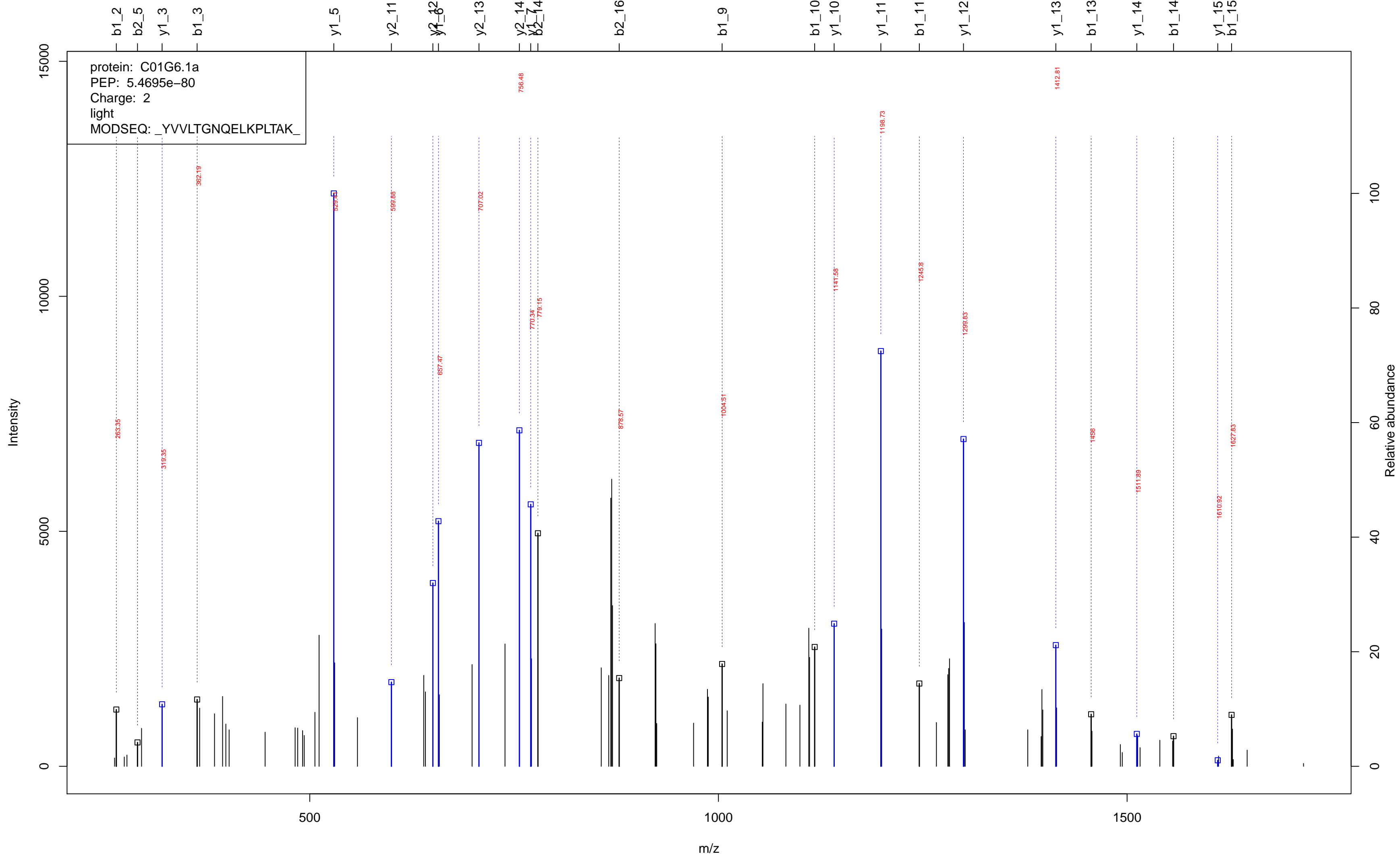


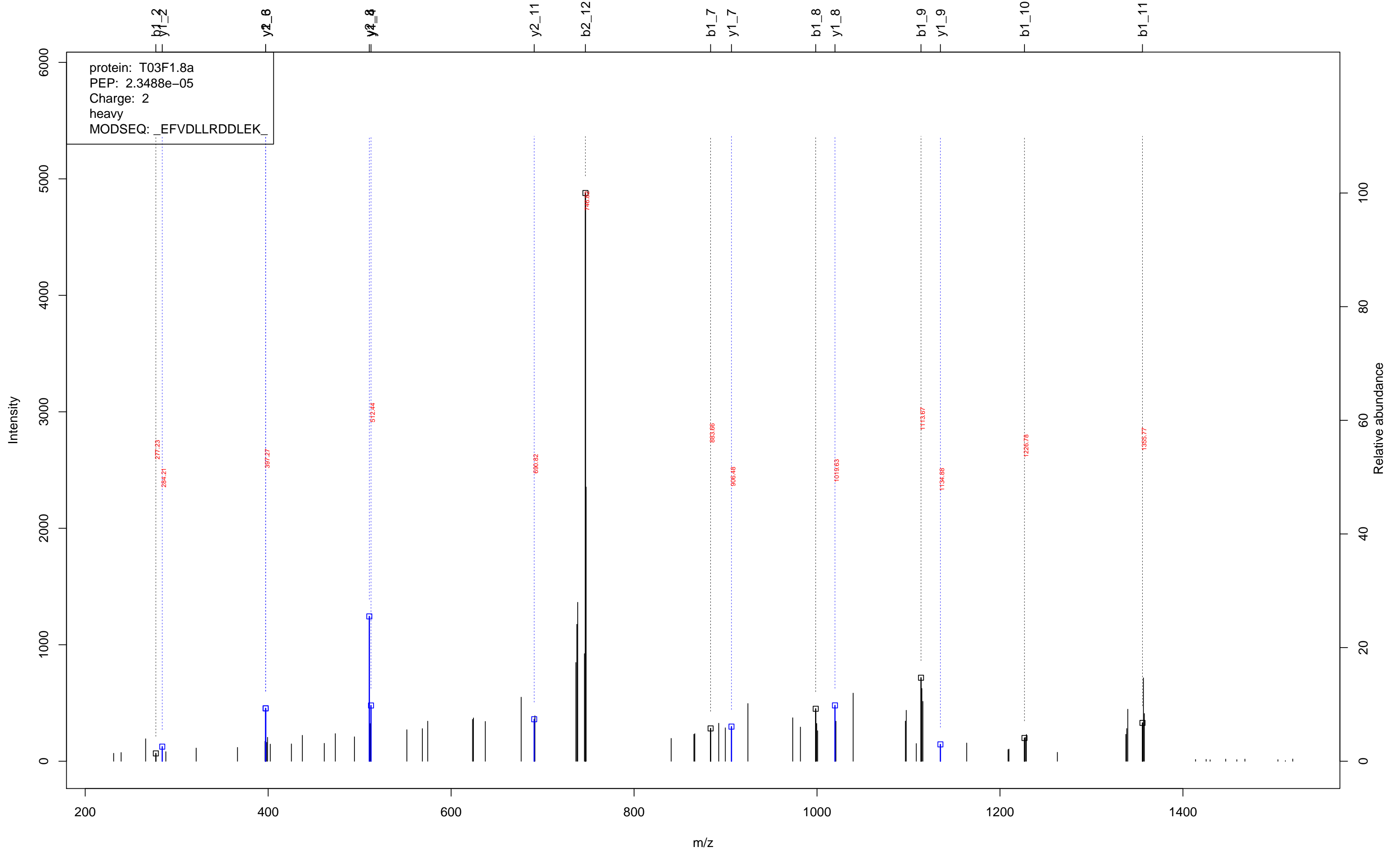
y2_4 y1_3 b1_2 y2_7 y2_9 y2_10 y2_11 y2_12 y2_13 y2_14 y2_15 y1_8 y1_9 y1_10 b1_9 y1_11 y1_12 b1_13

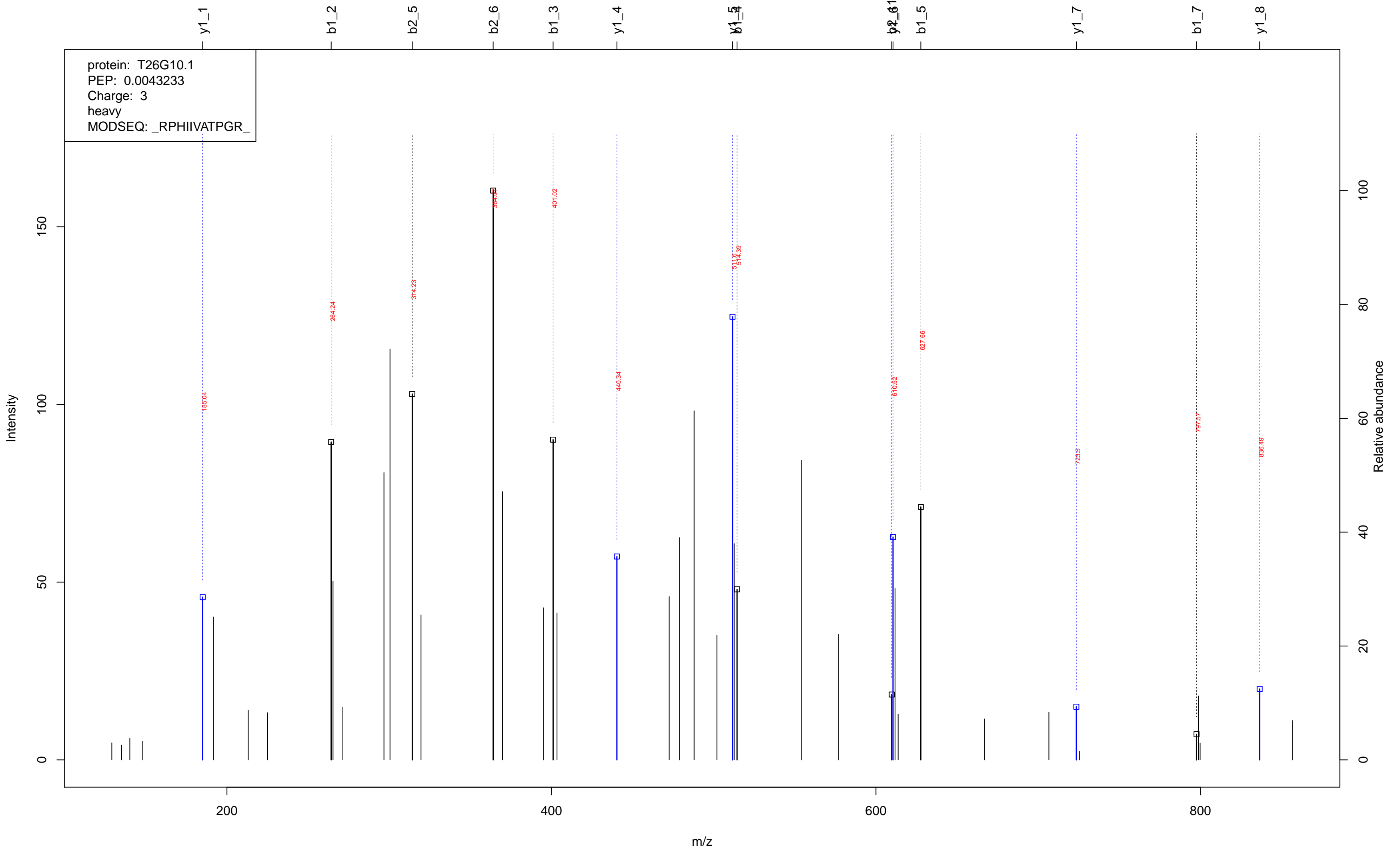
protein: Y82E9BR.14b
PEP: 1.8499e-07
Charge: 3
light
MODSEQ: _YLQQLIDADLAEHGK_

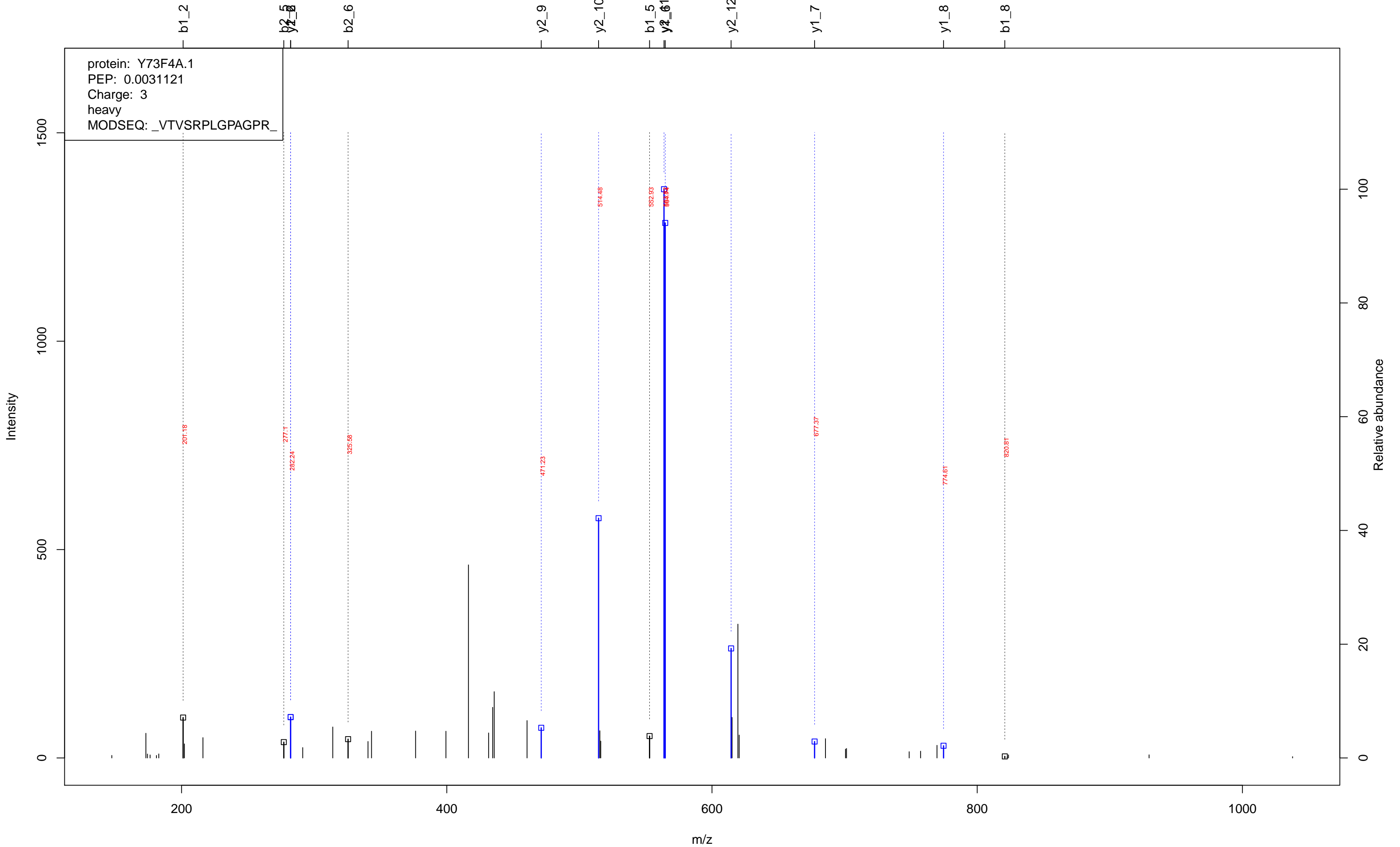












y1_1 b2_4 y1_3 y2_8 b2_9 y2_9 y2_10 b2_11 y2_11 y1_7 b1_8 b1_9

protein: T10C6.5
PEP: 0.00055861
Charge: 3
heavy
MODSEQ: _(ac)TTAHRPTFHPAR_

