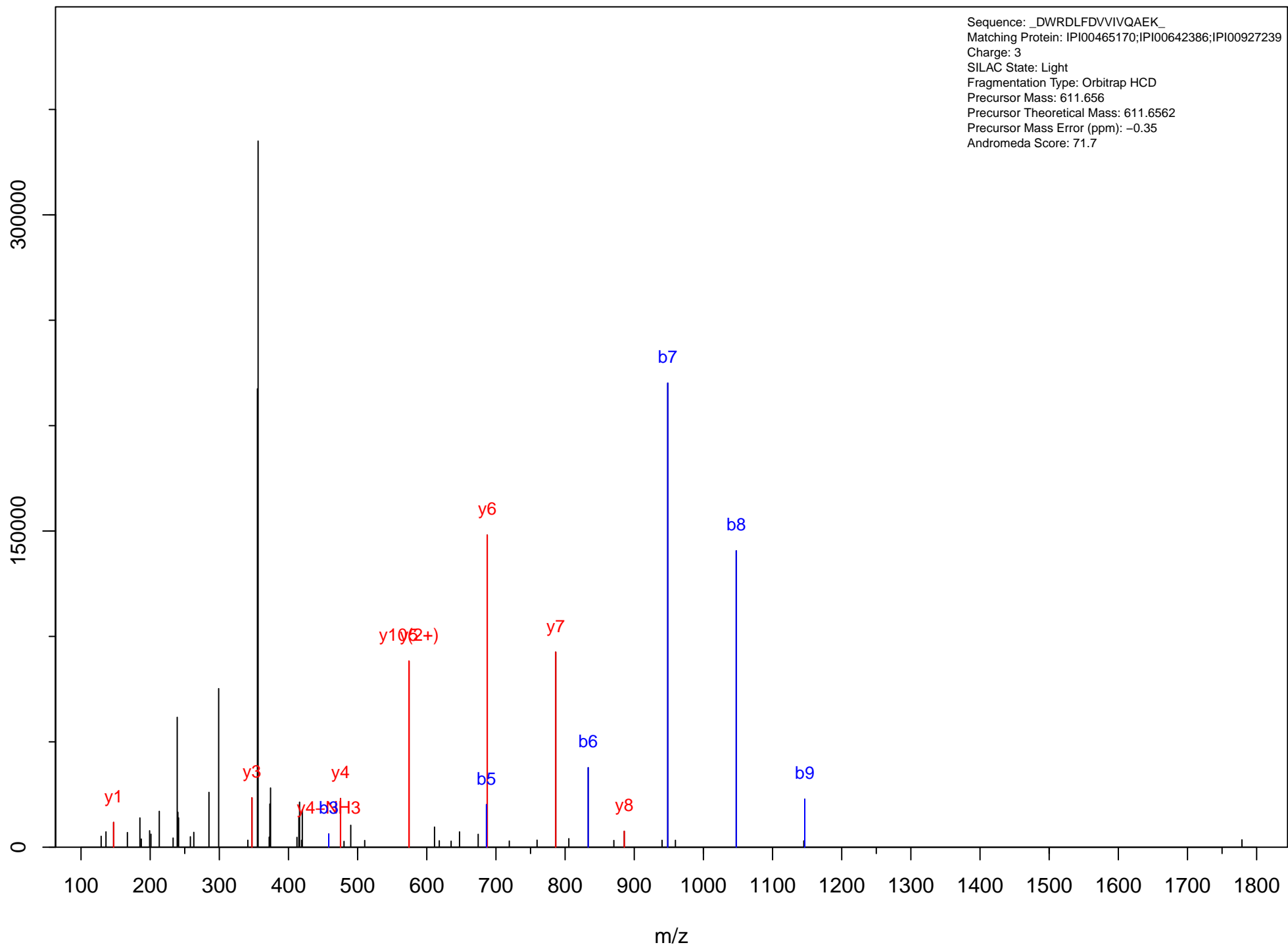
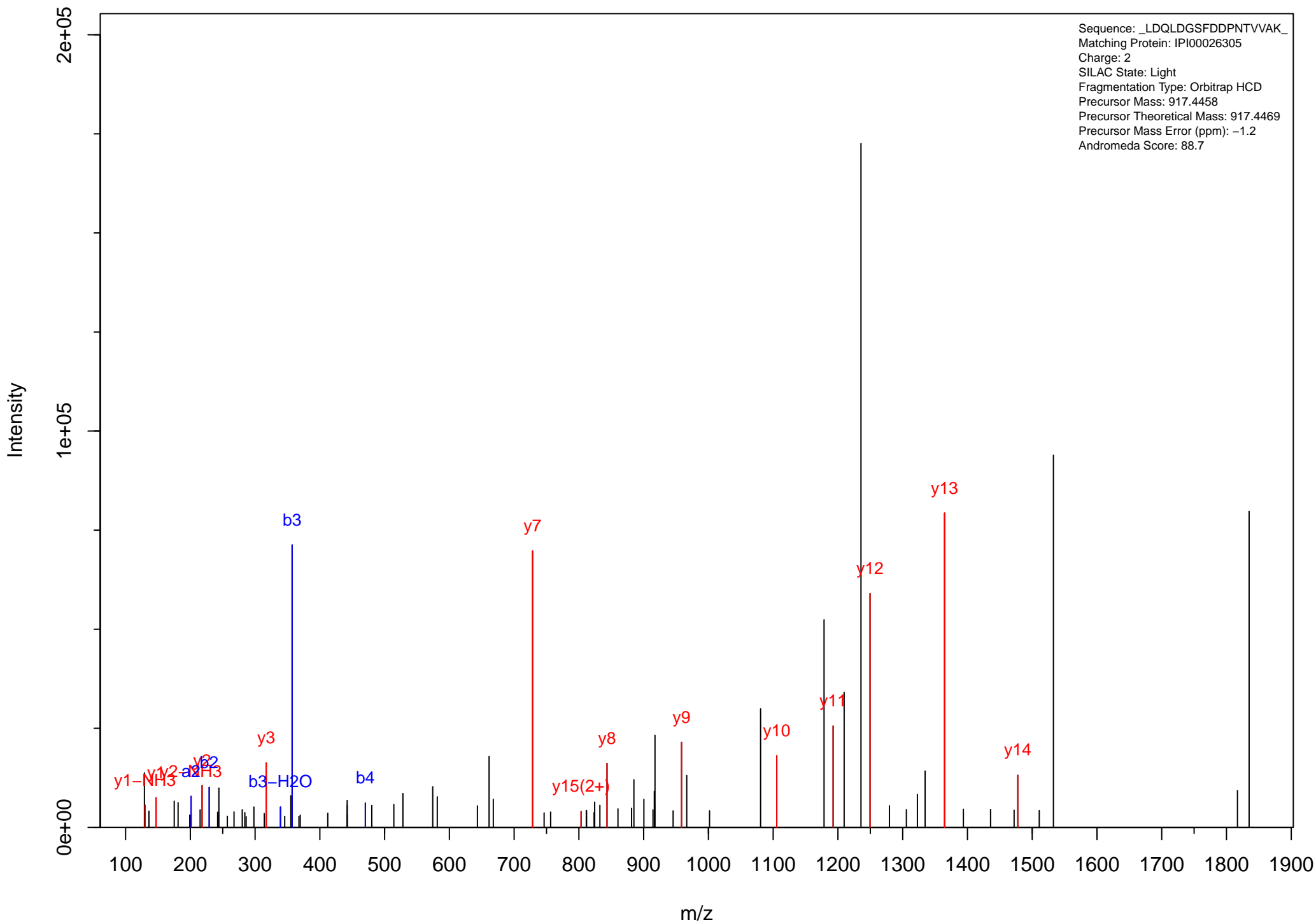
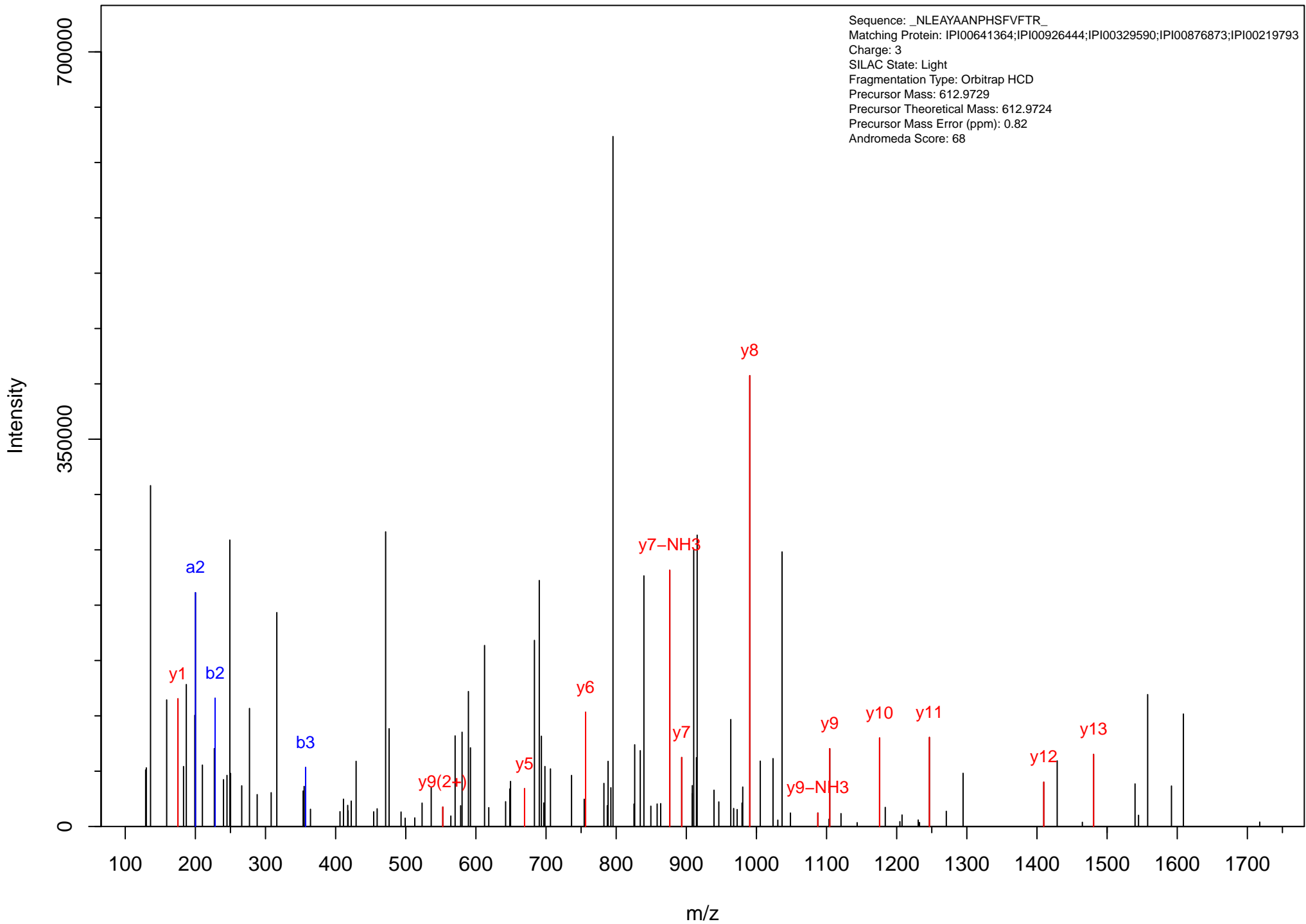
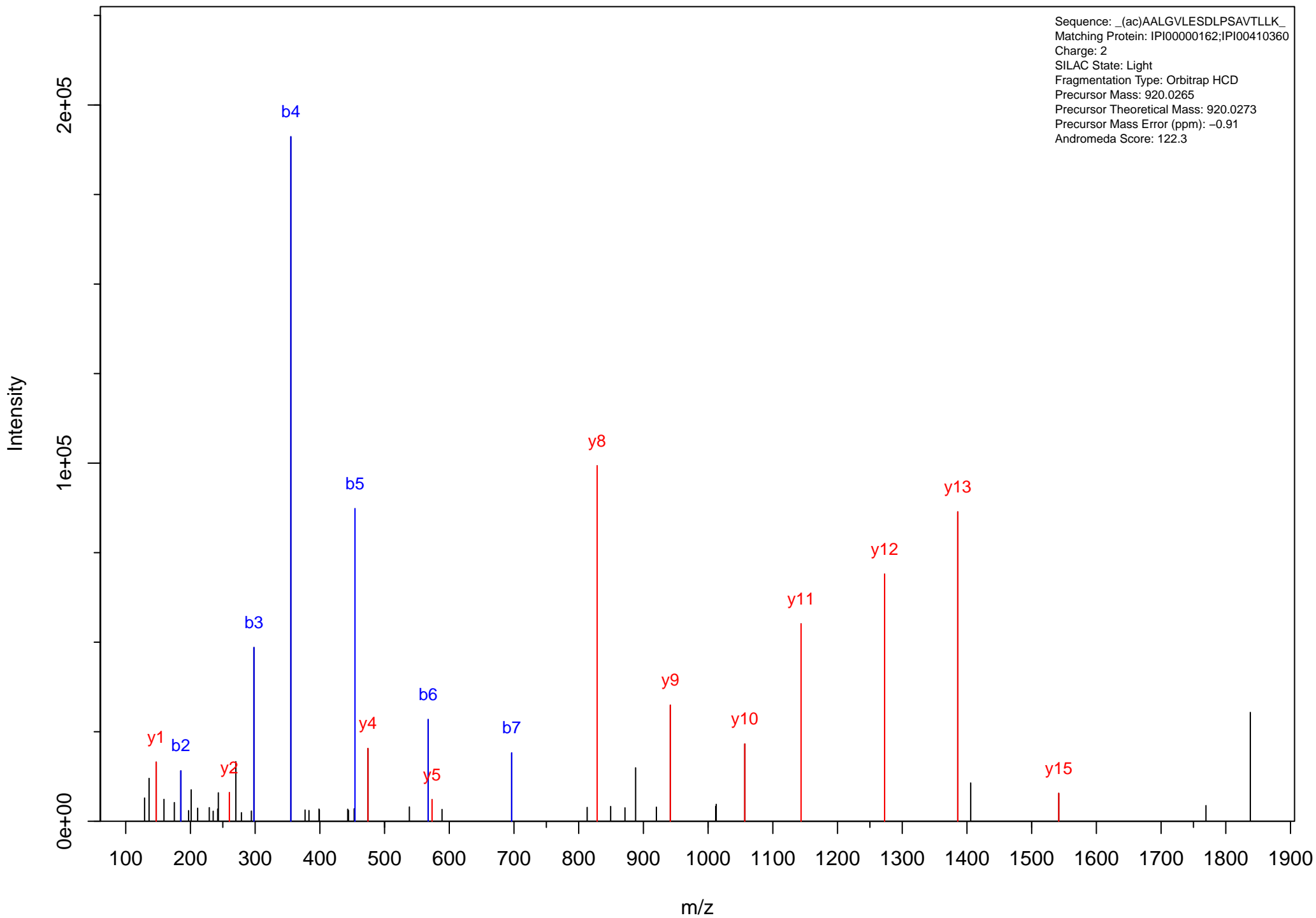


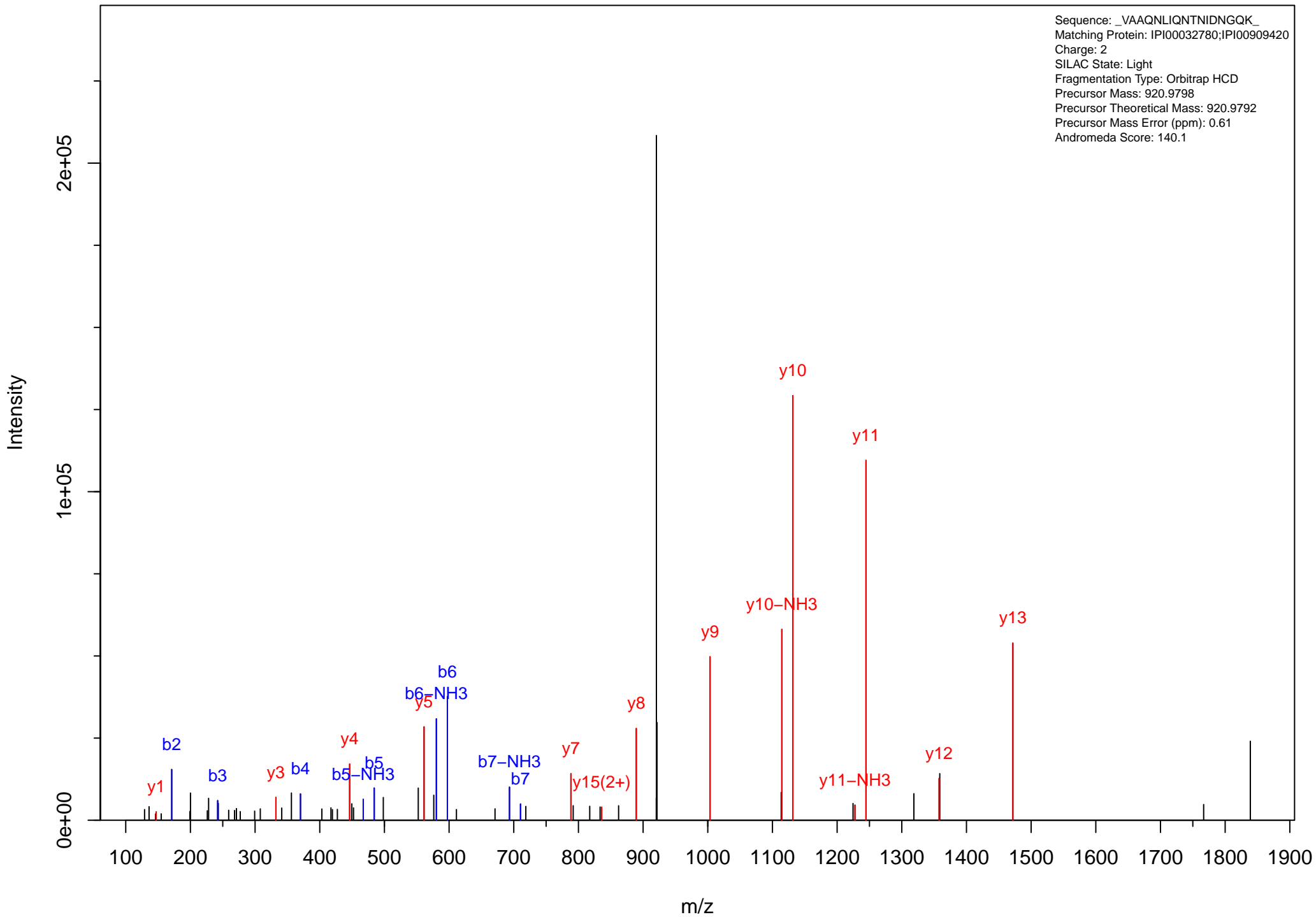
Intensity

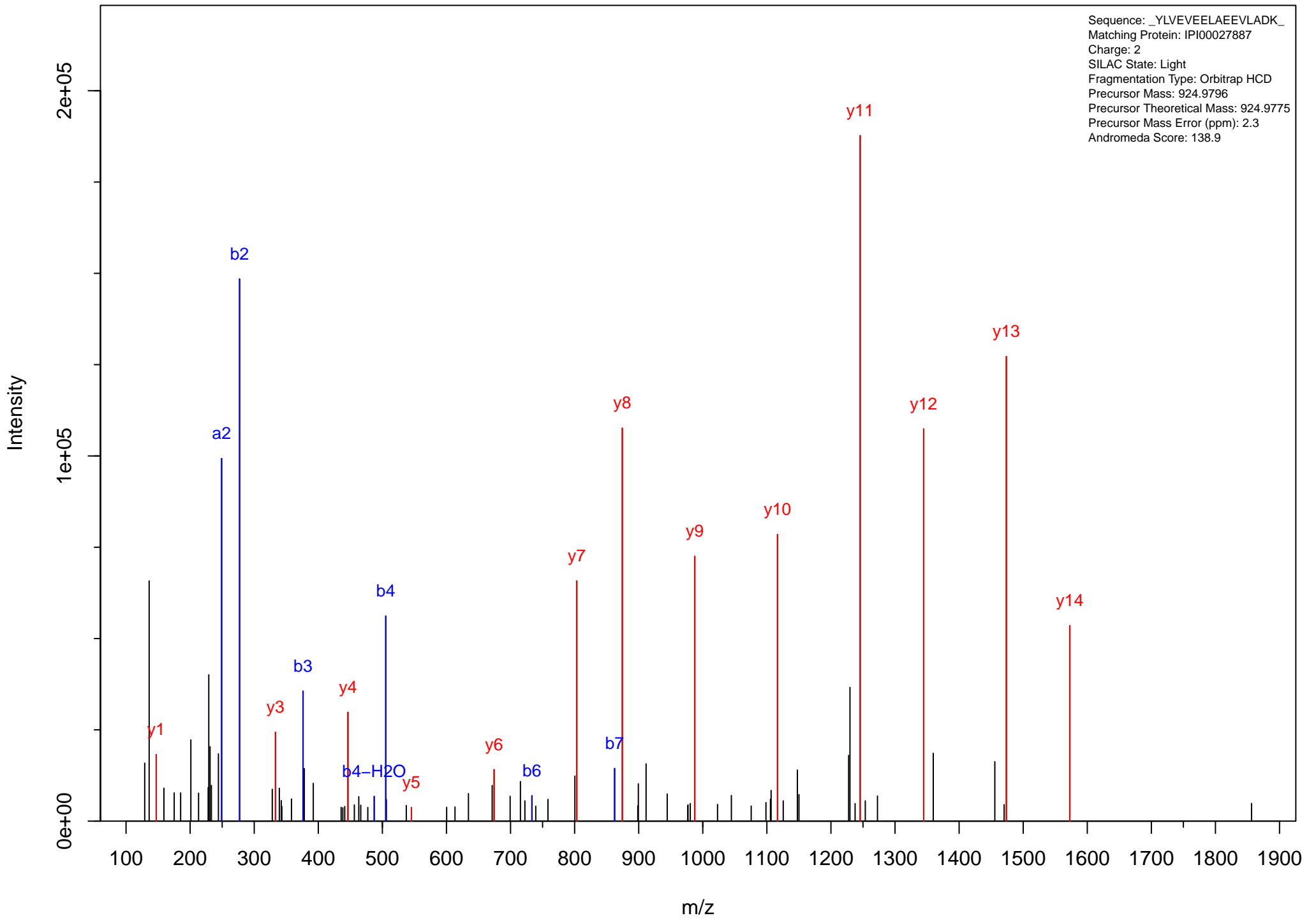


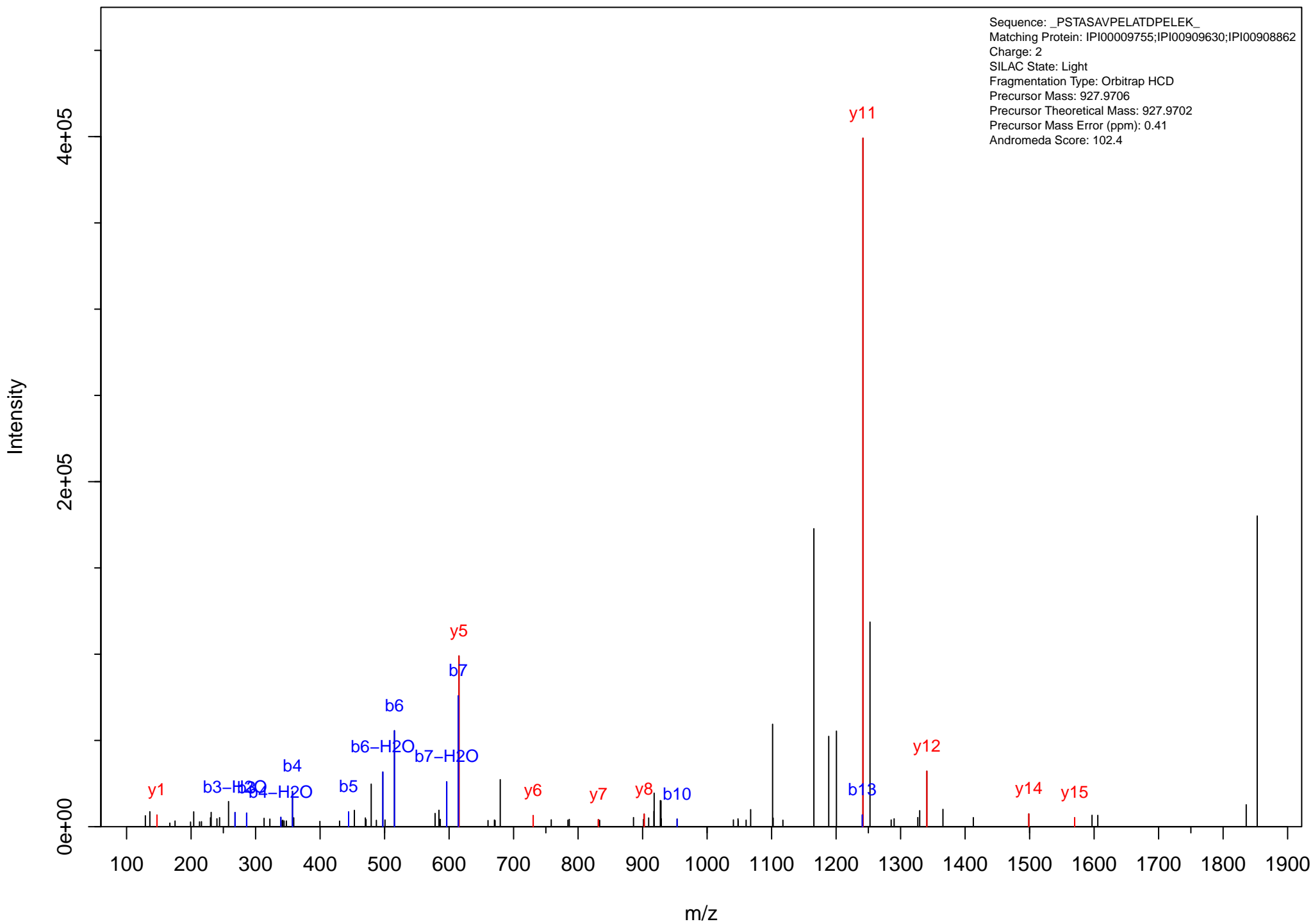


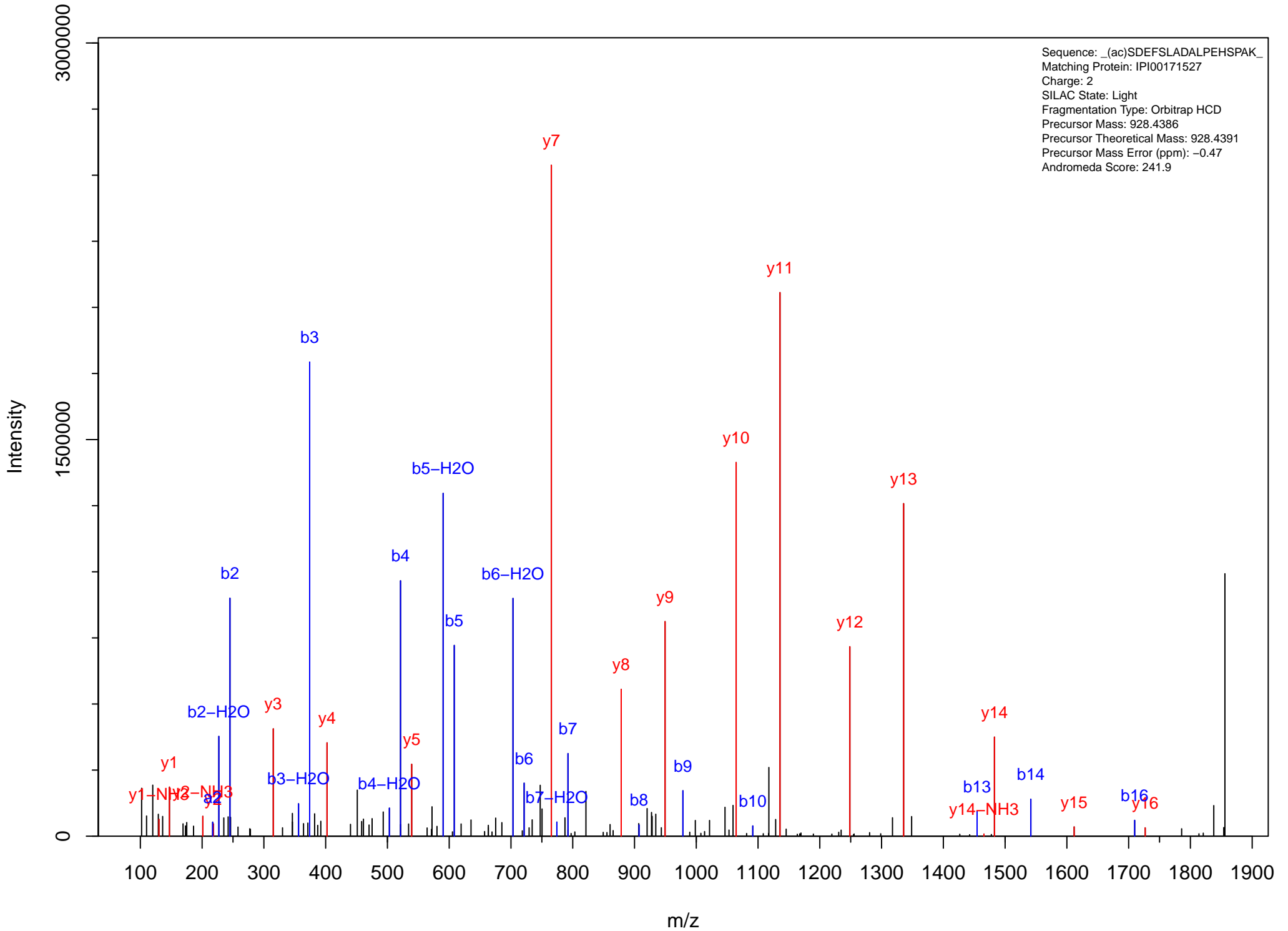


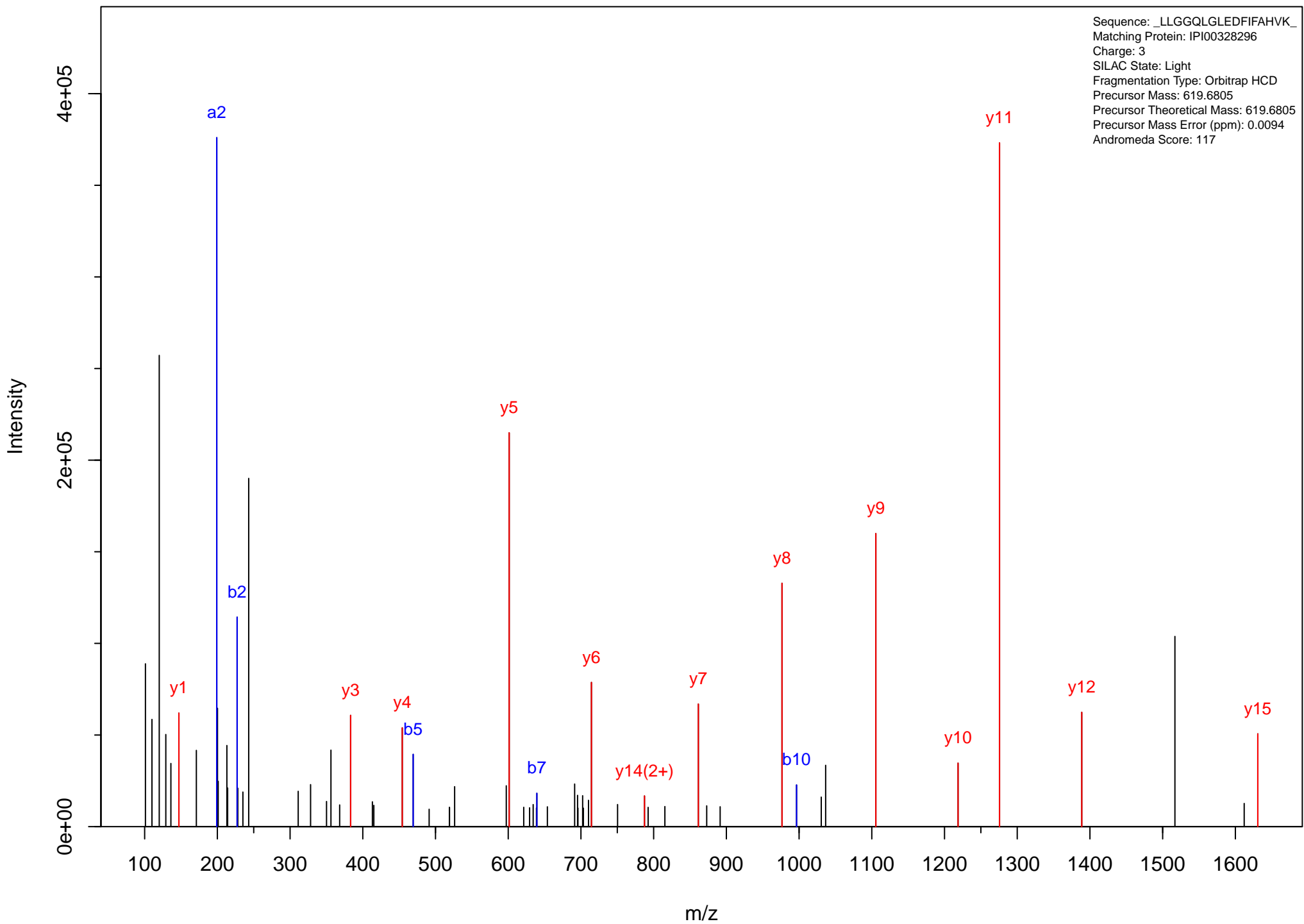


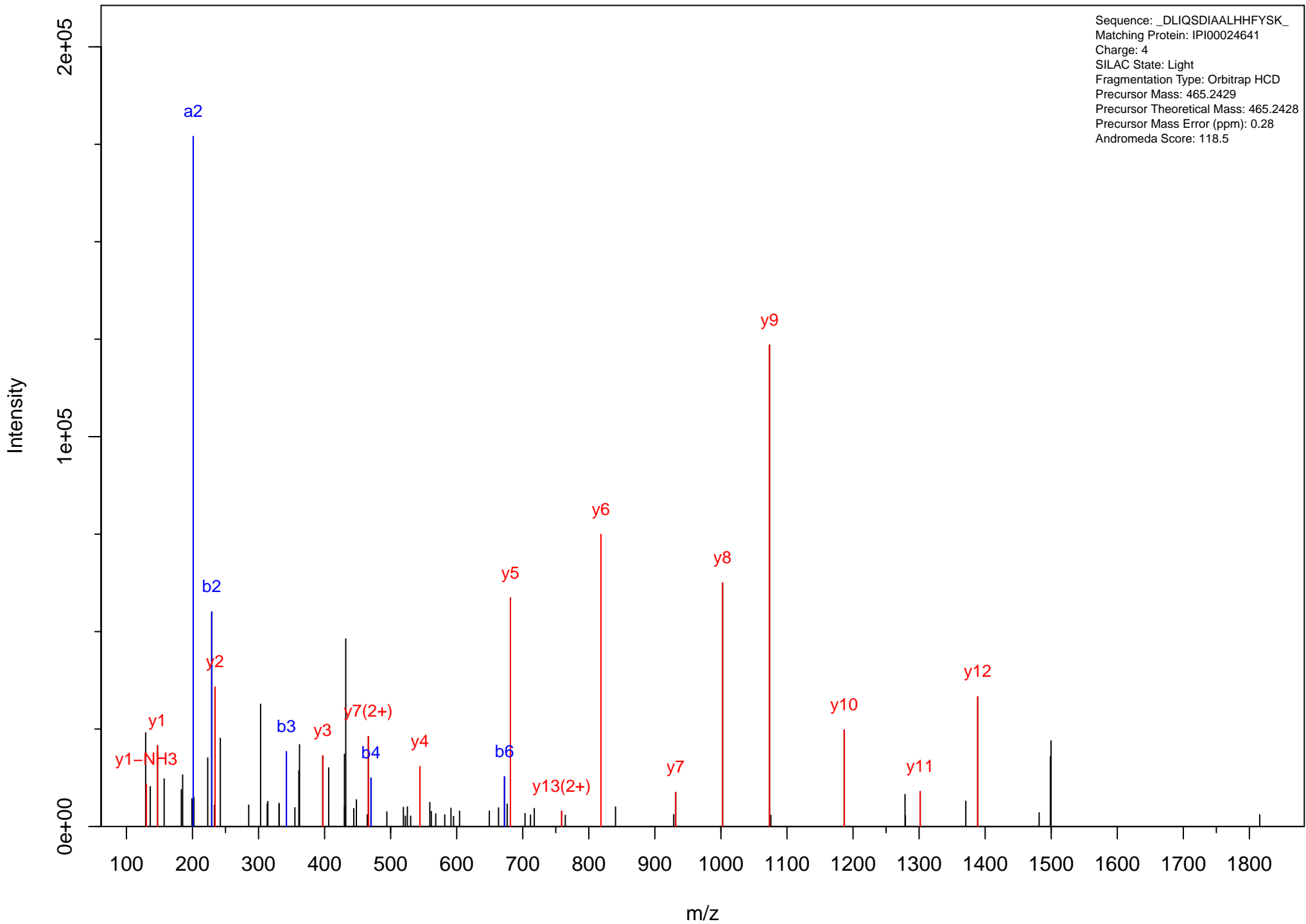


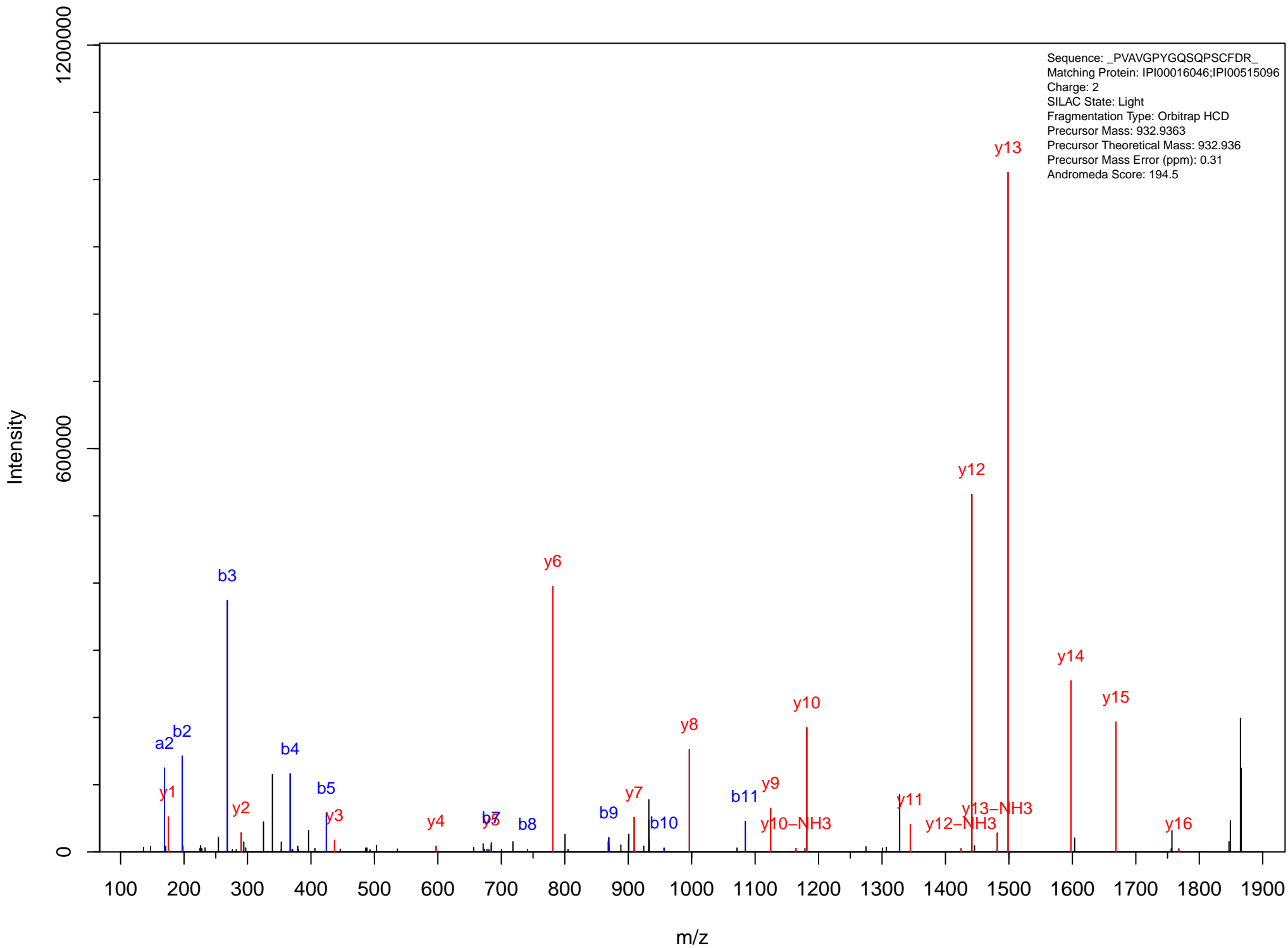


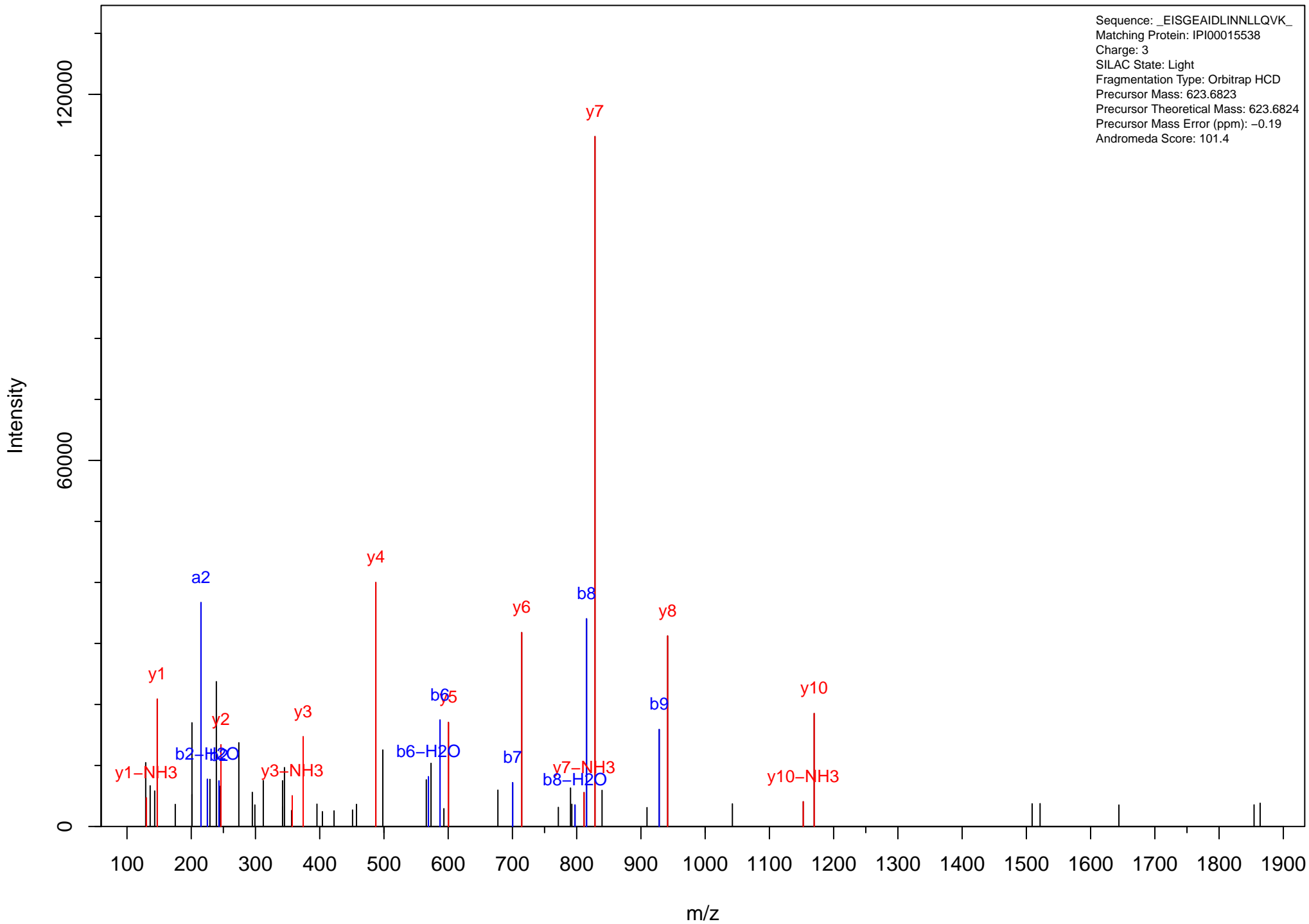


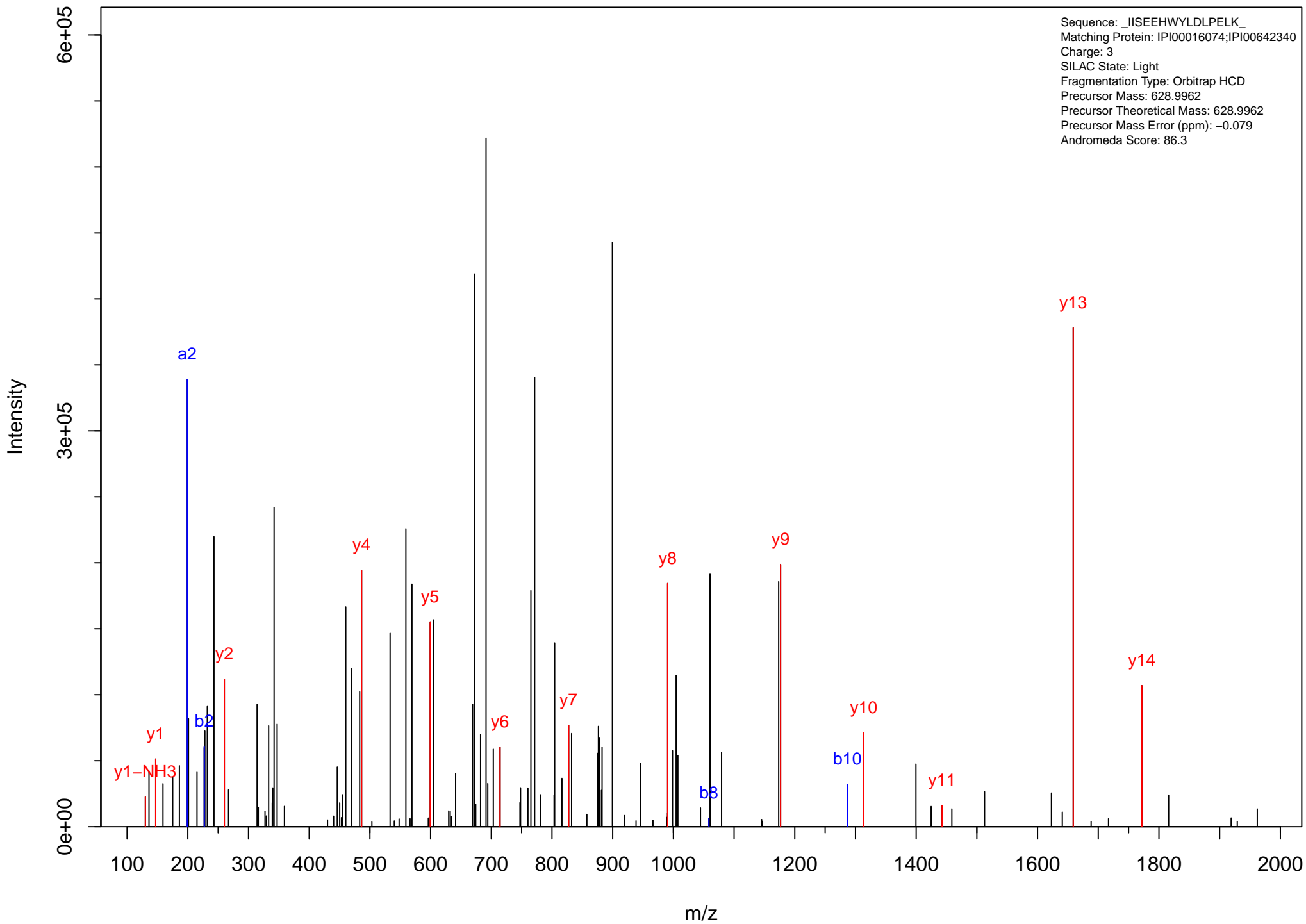


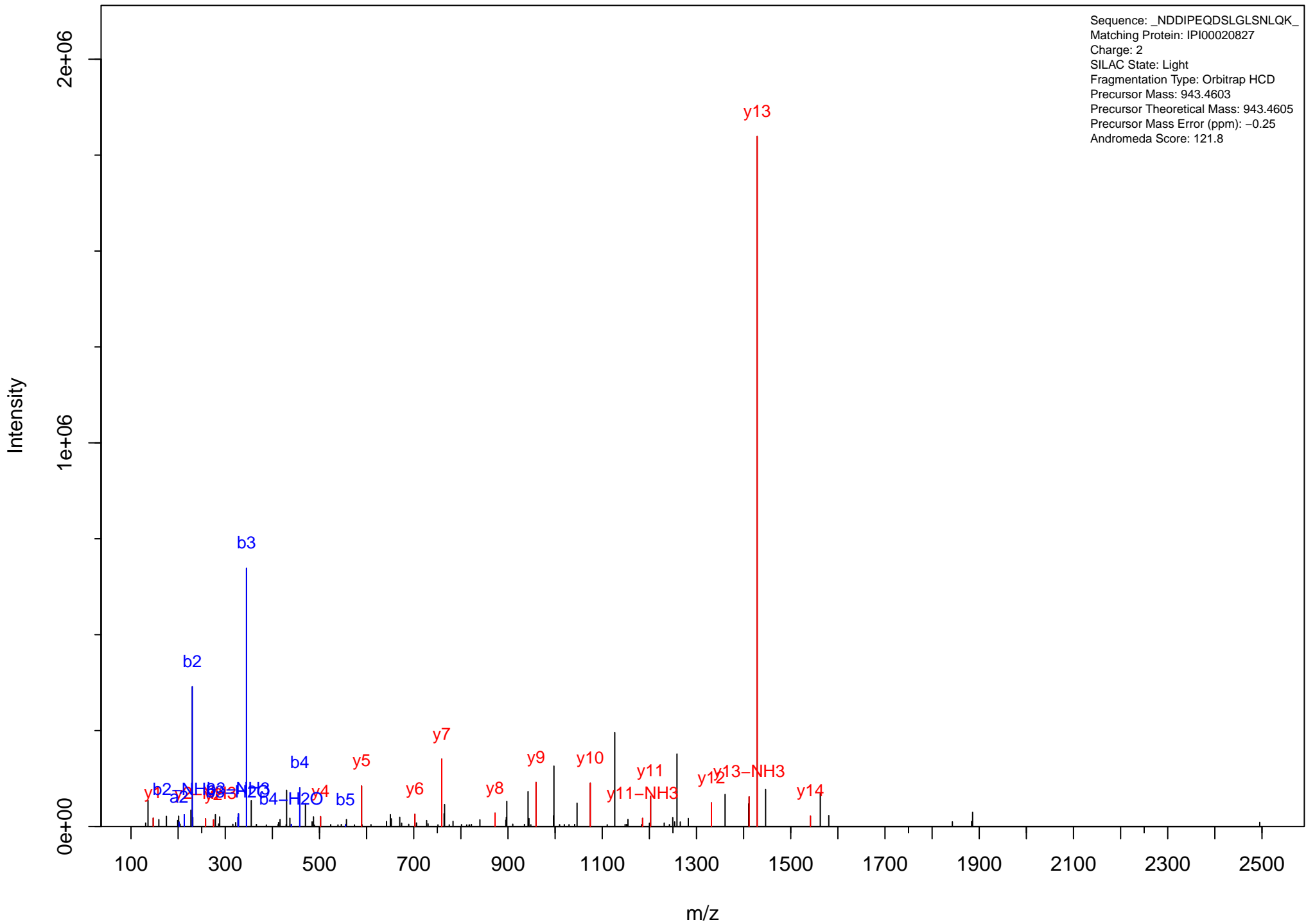


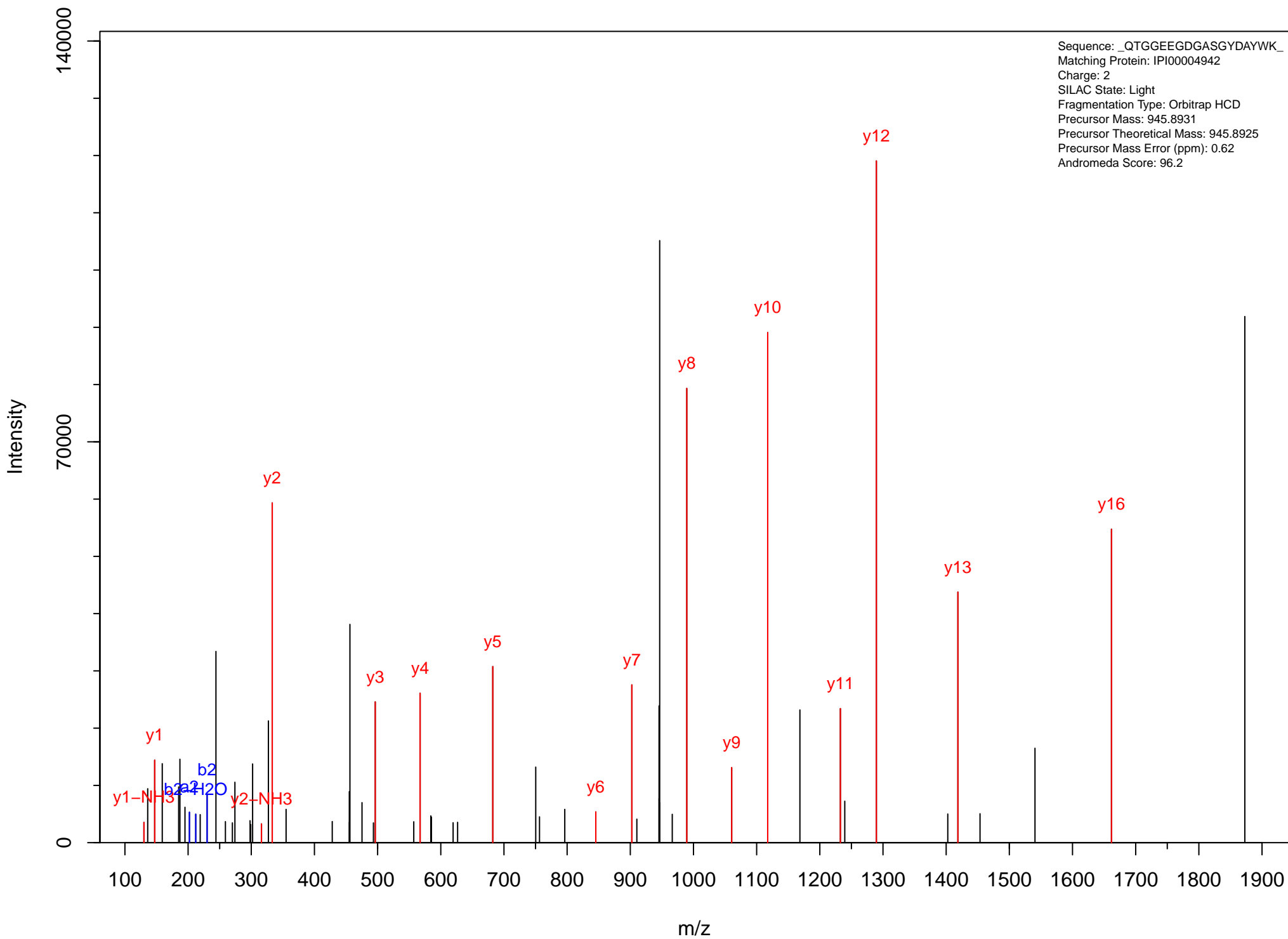


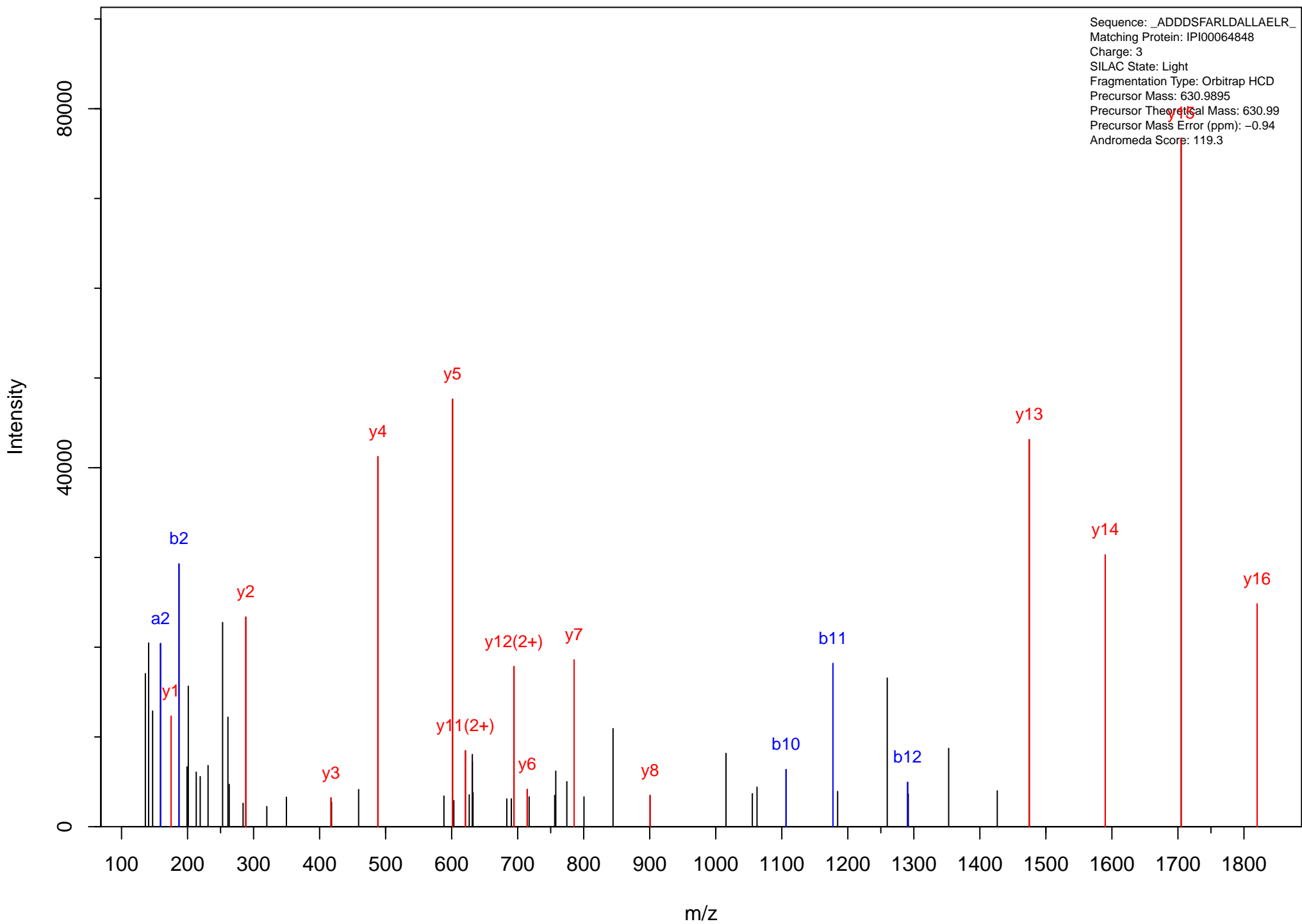


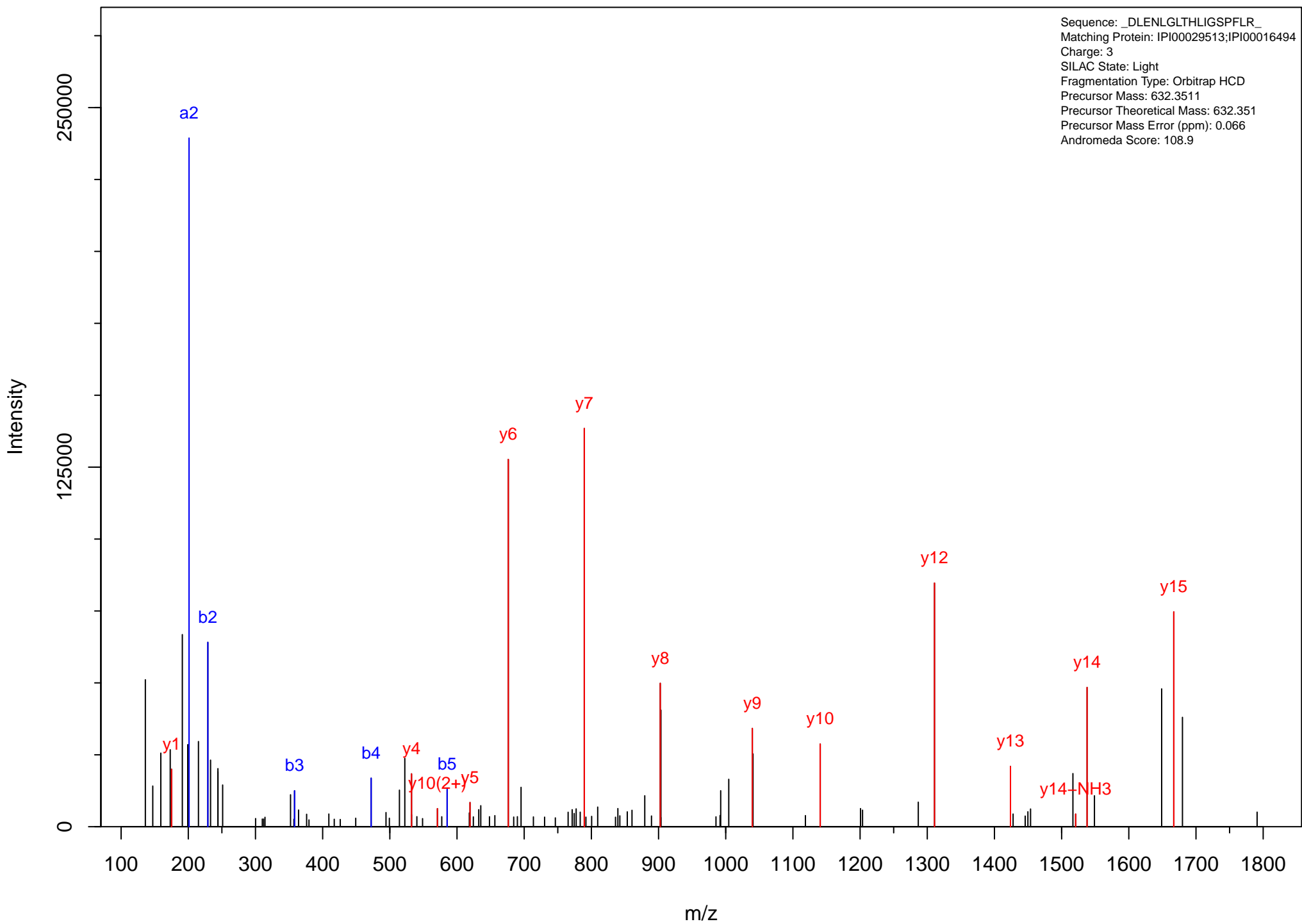


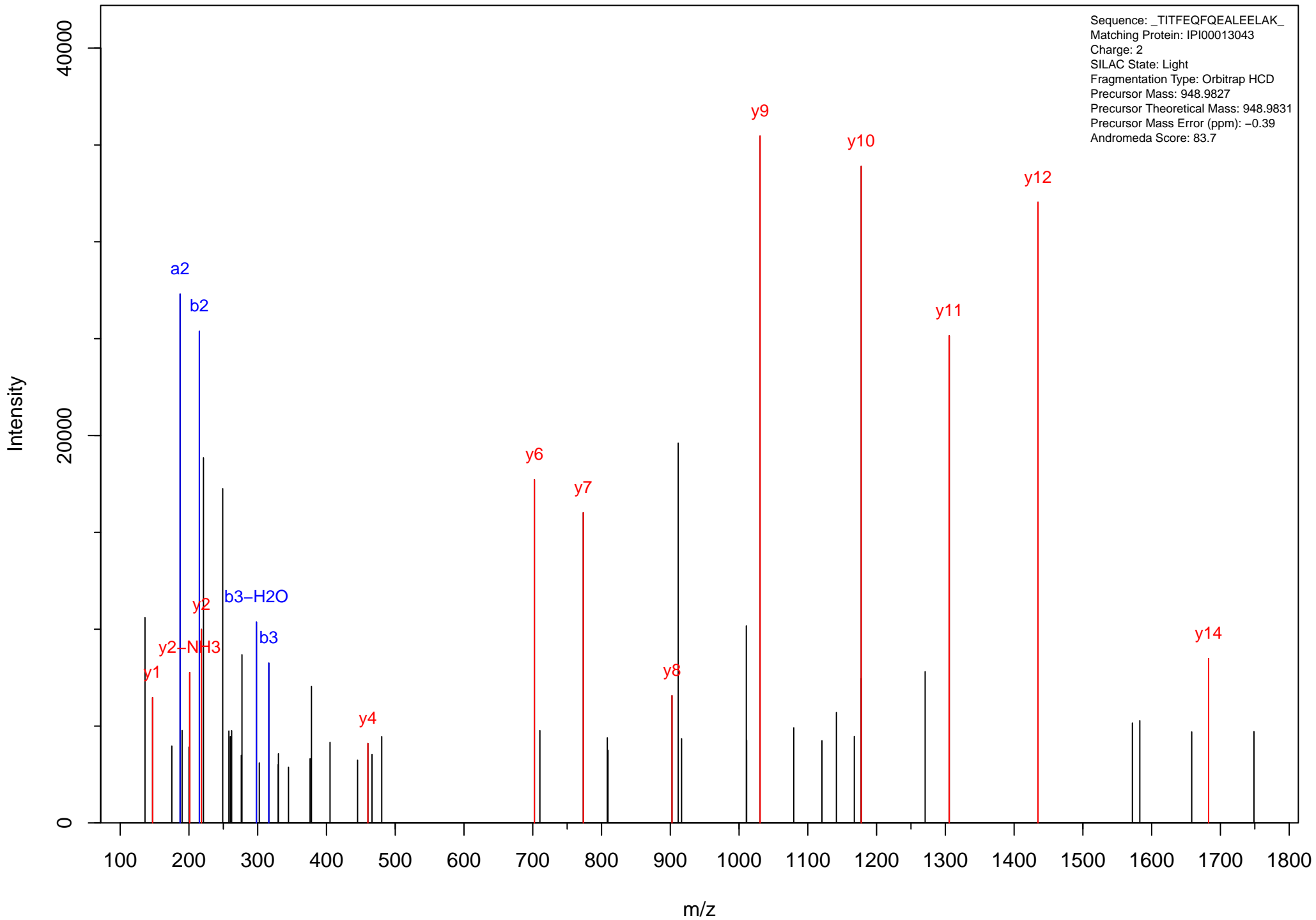


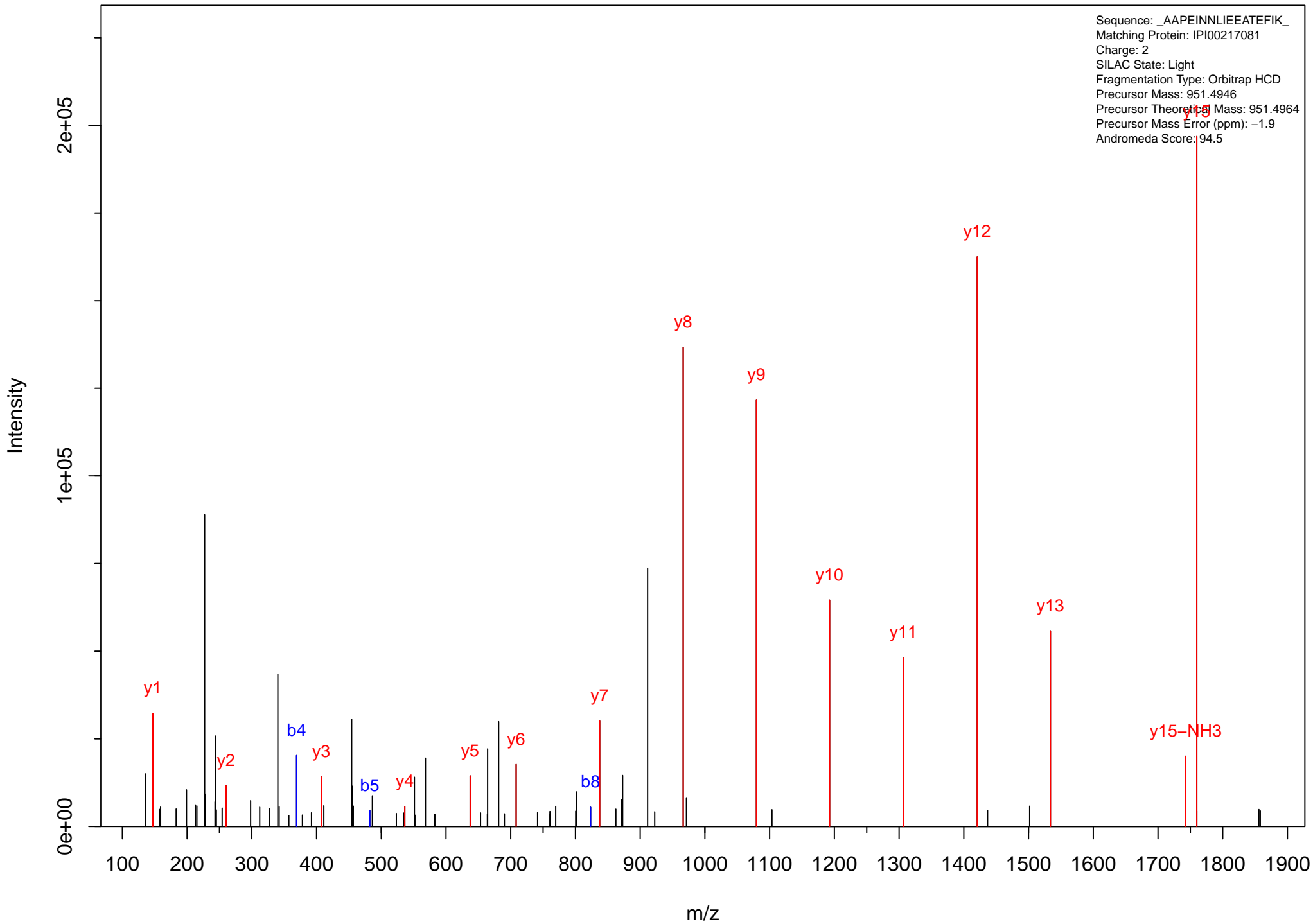


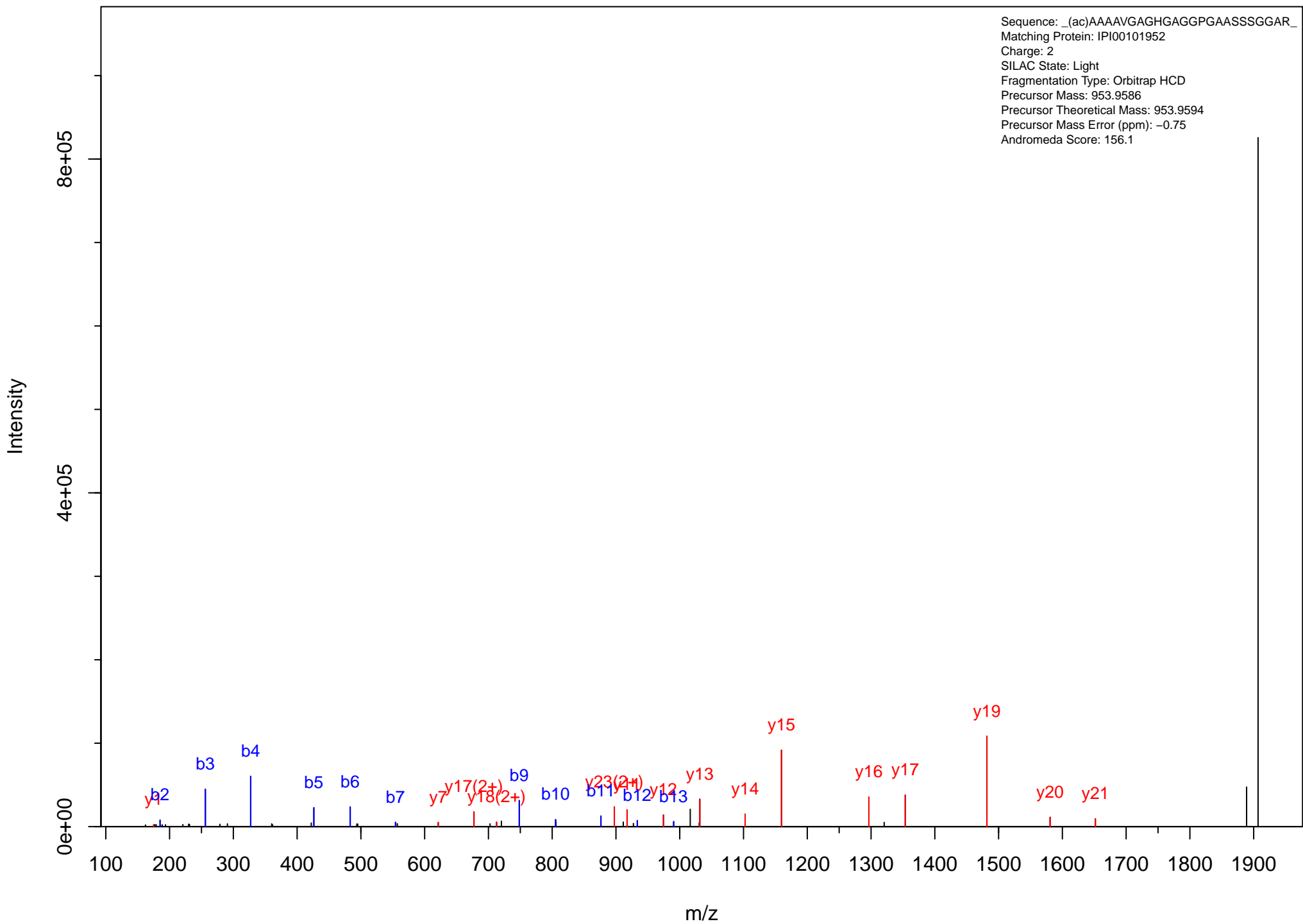


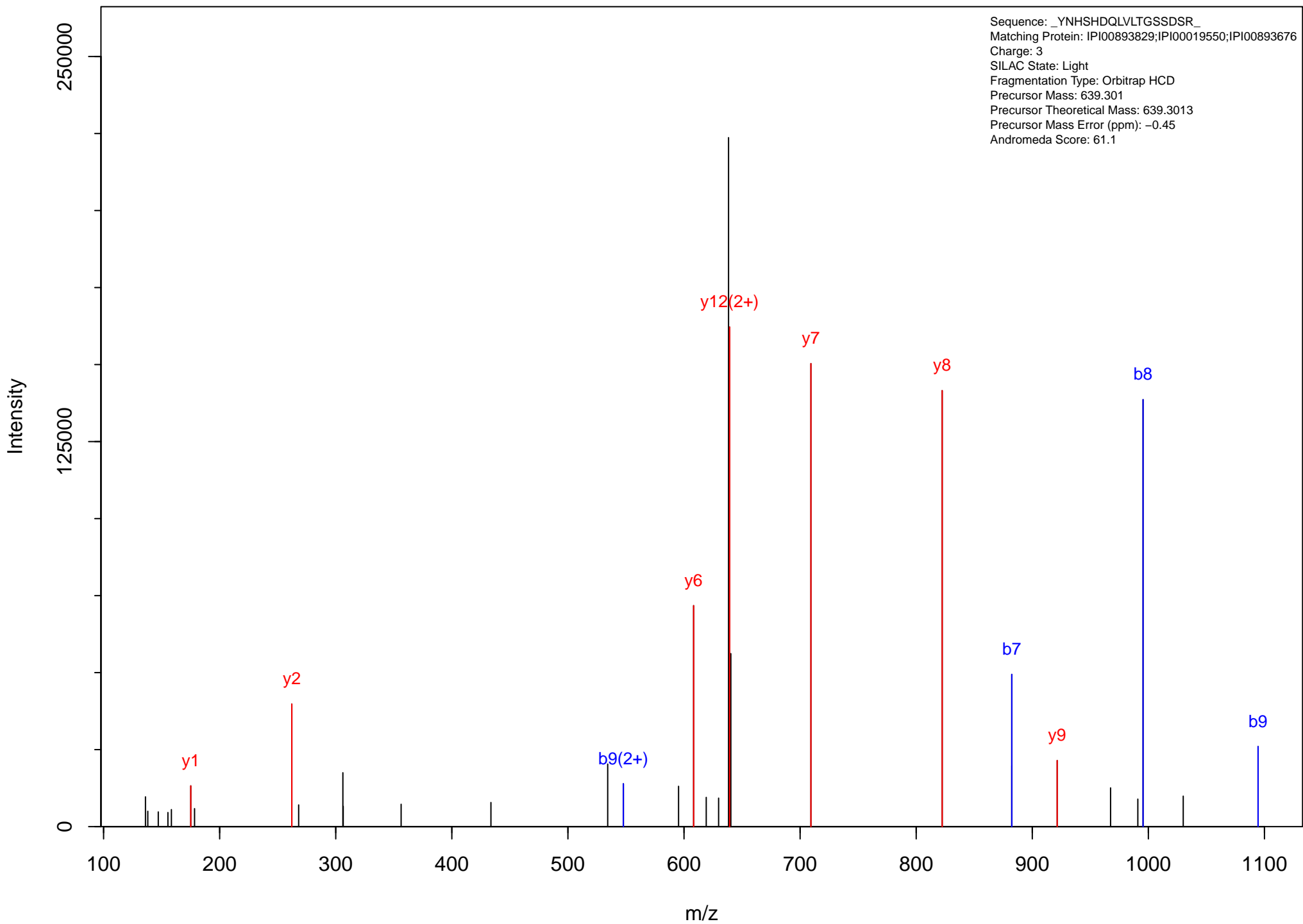


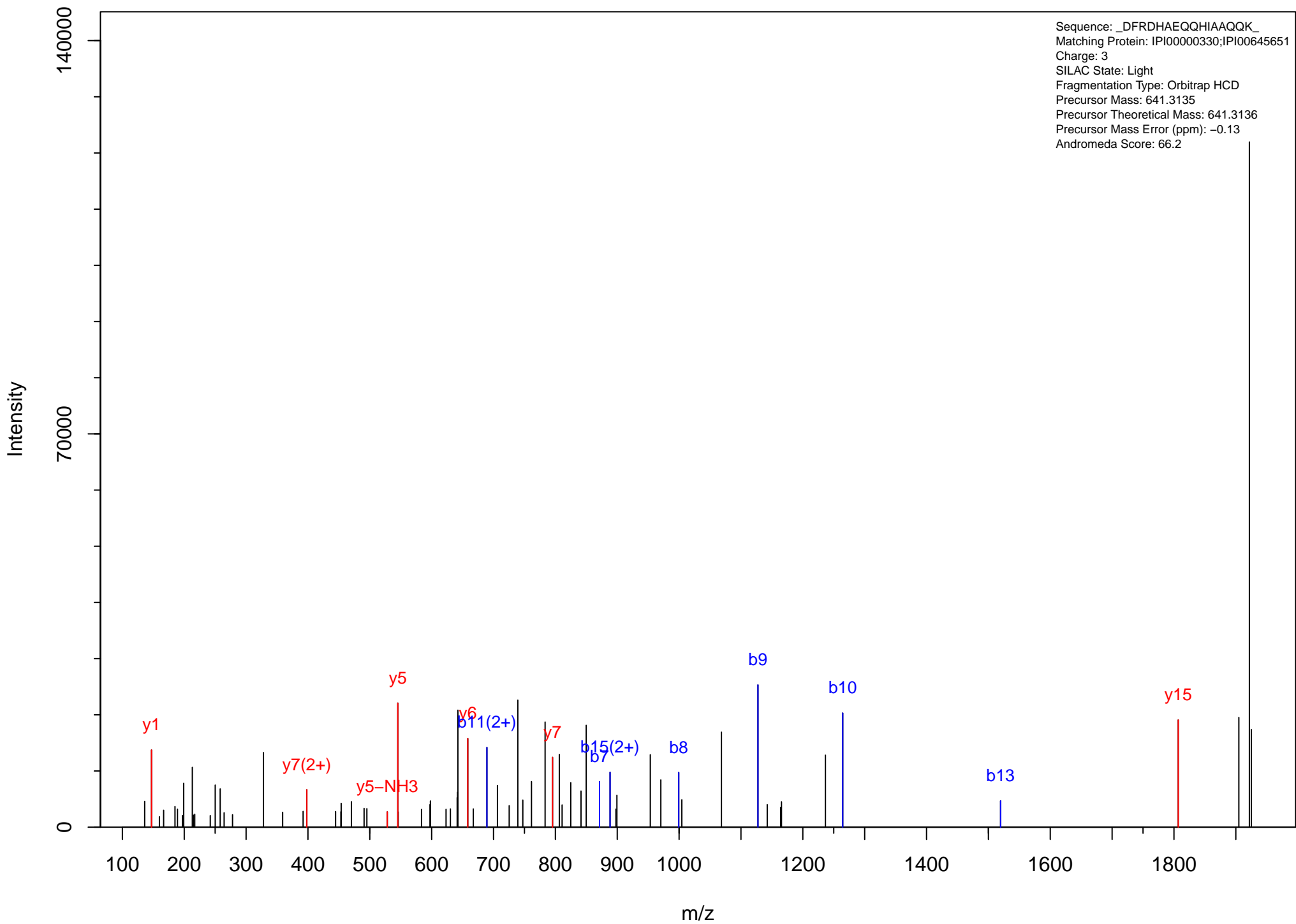


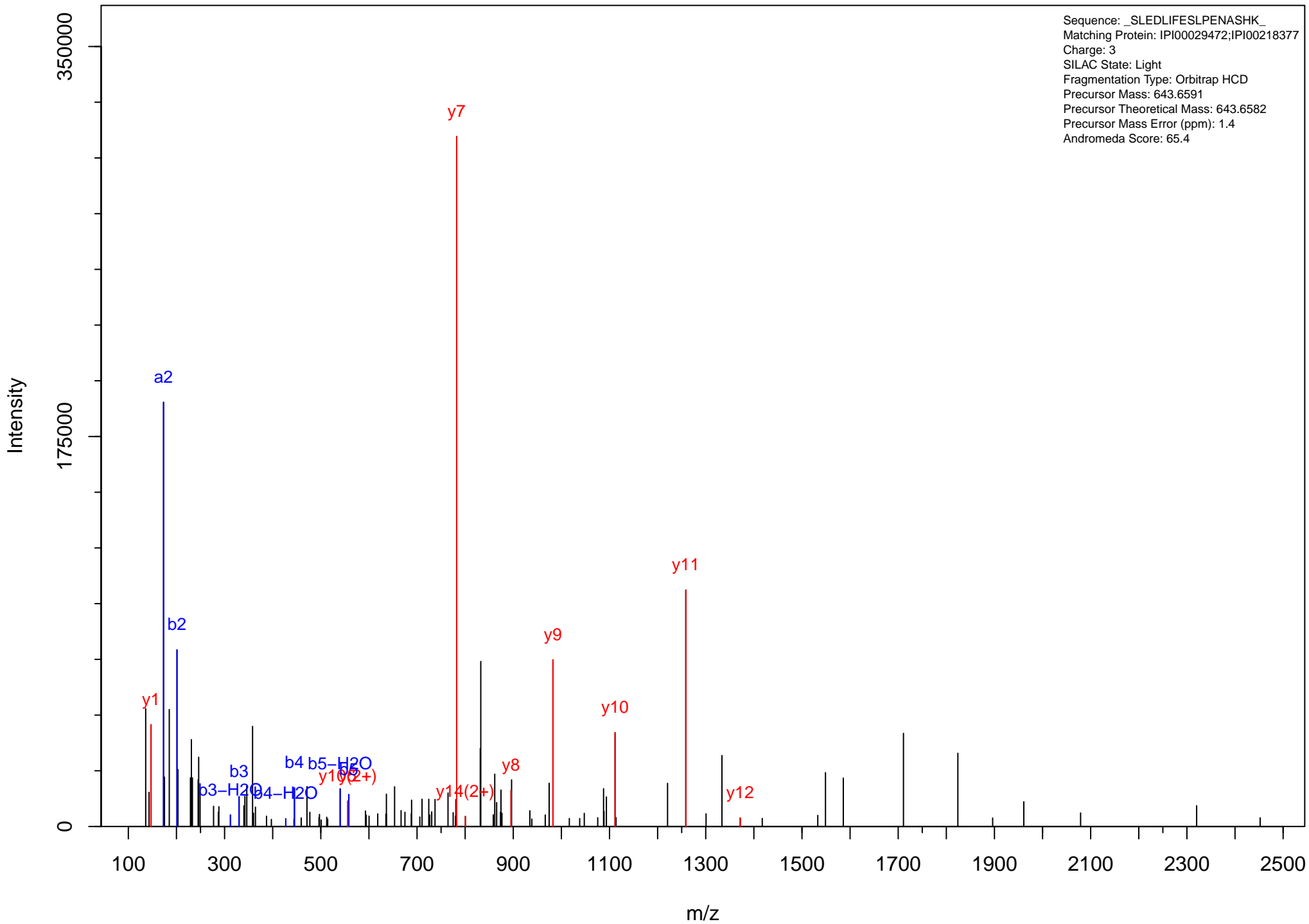


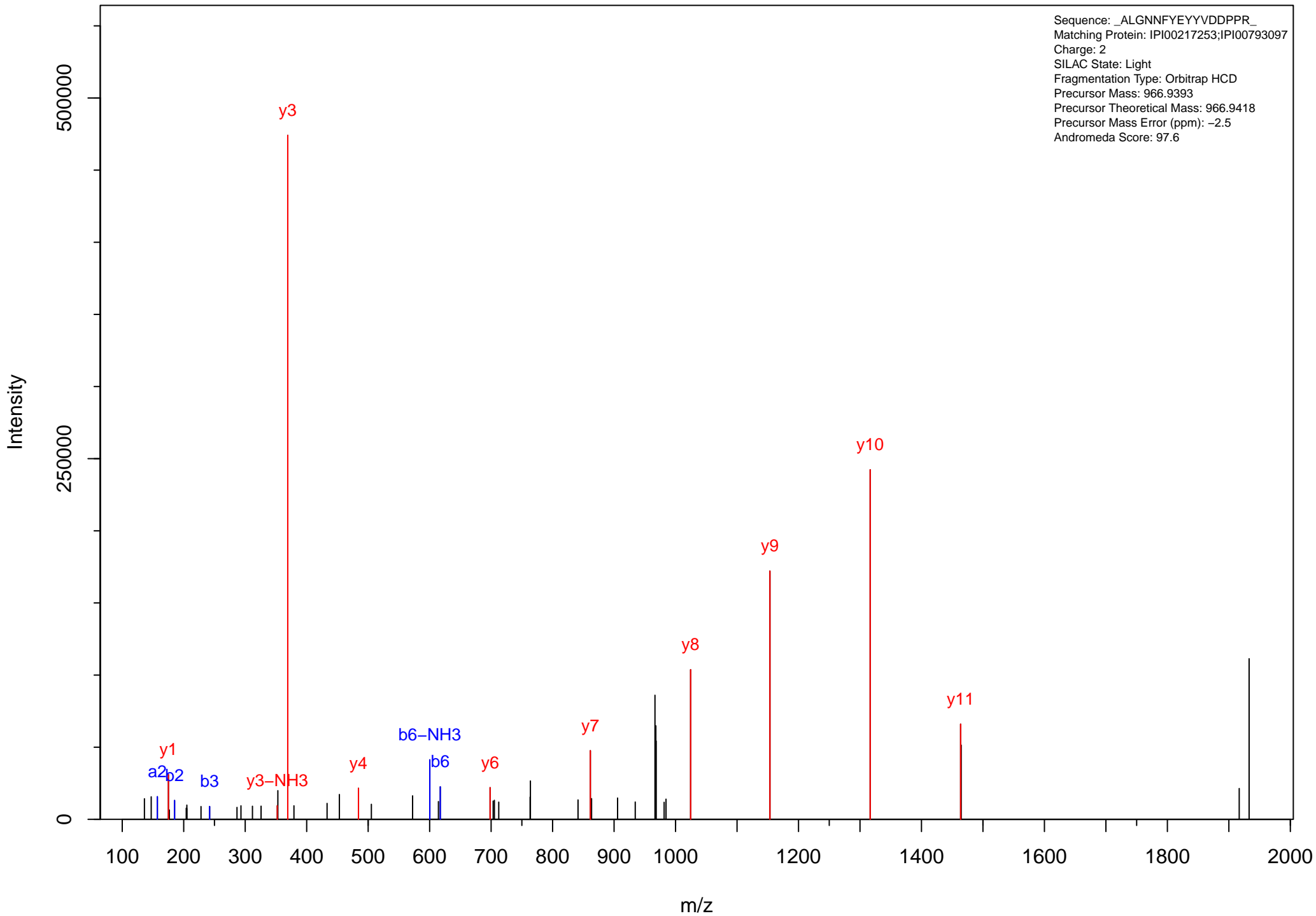


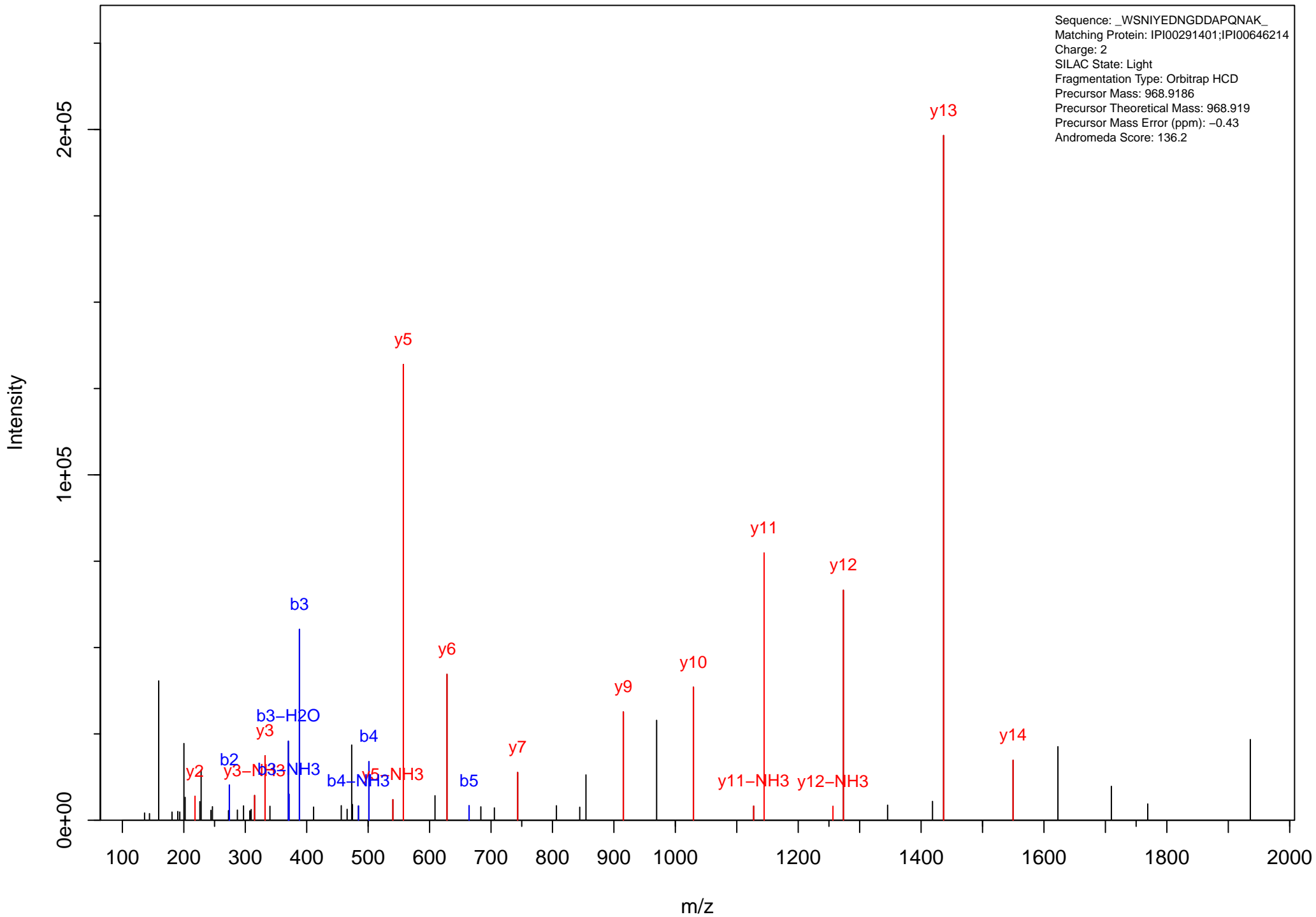


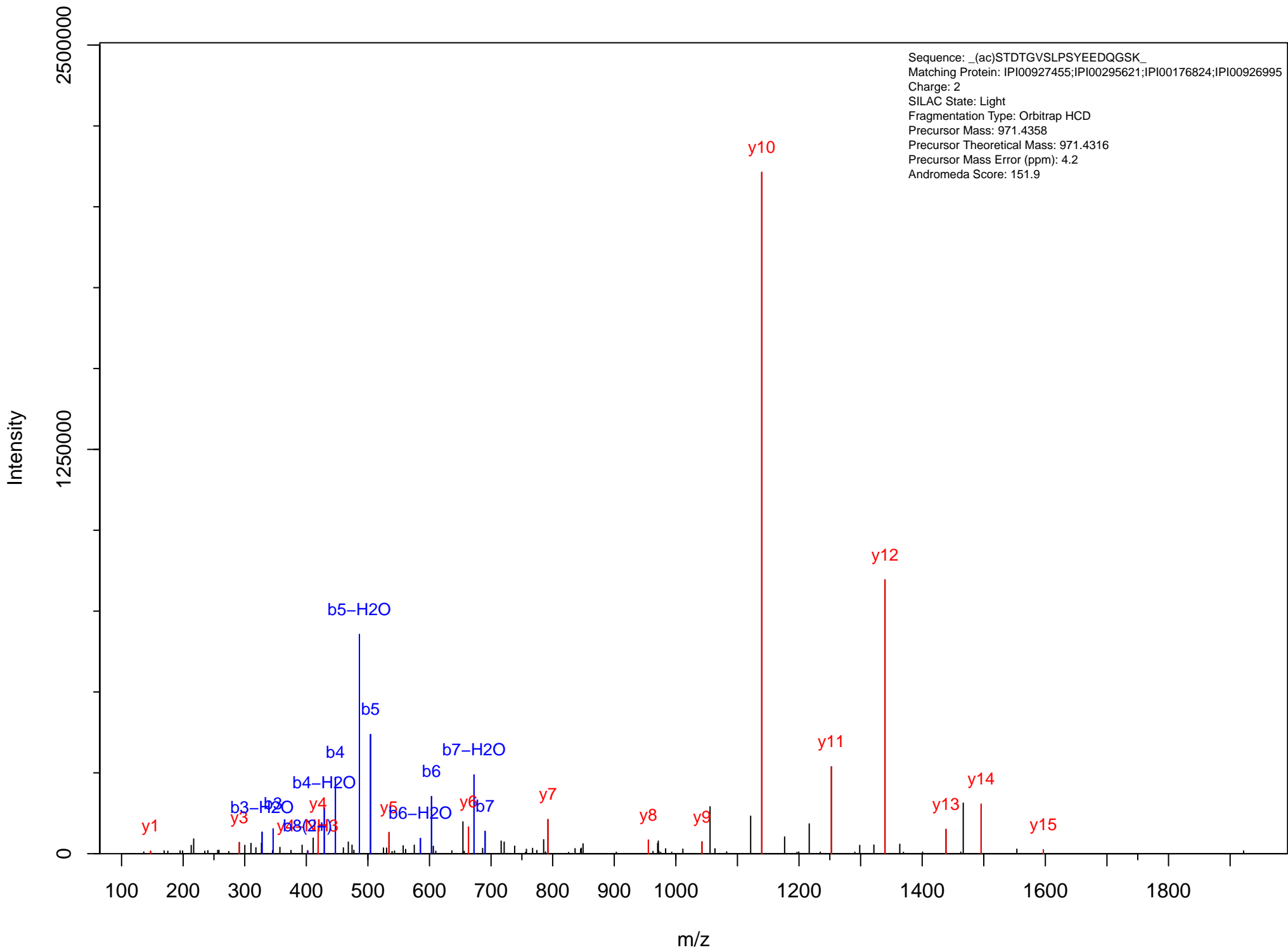


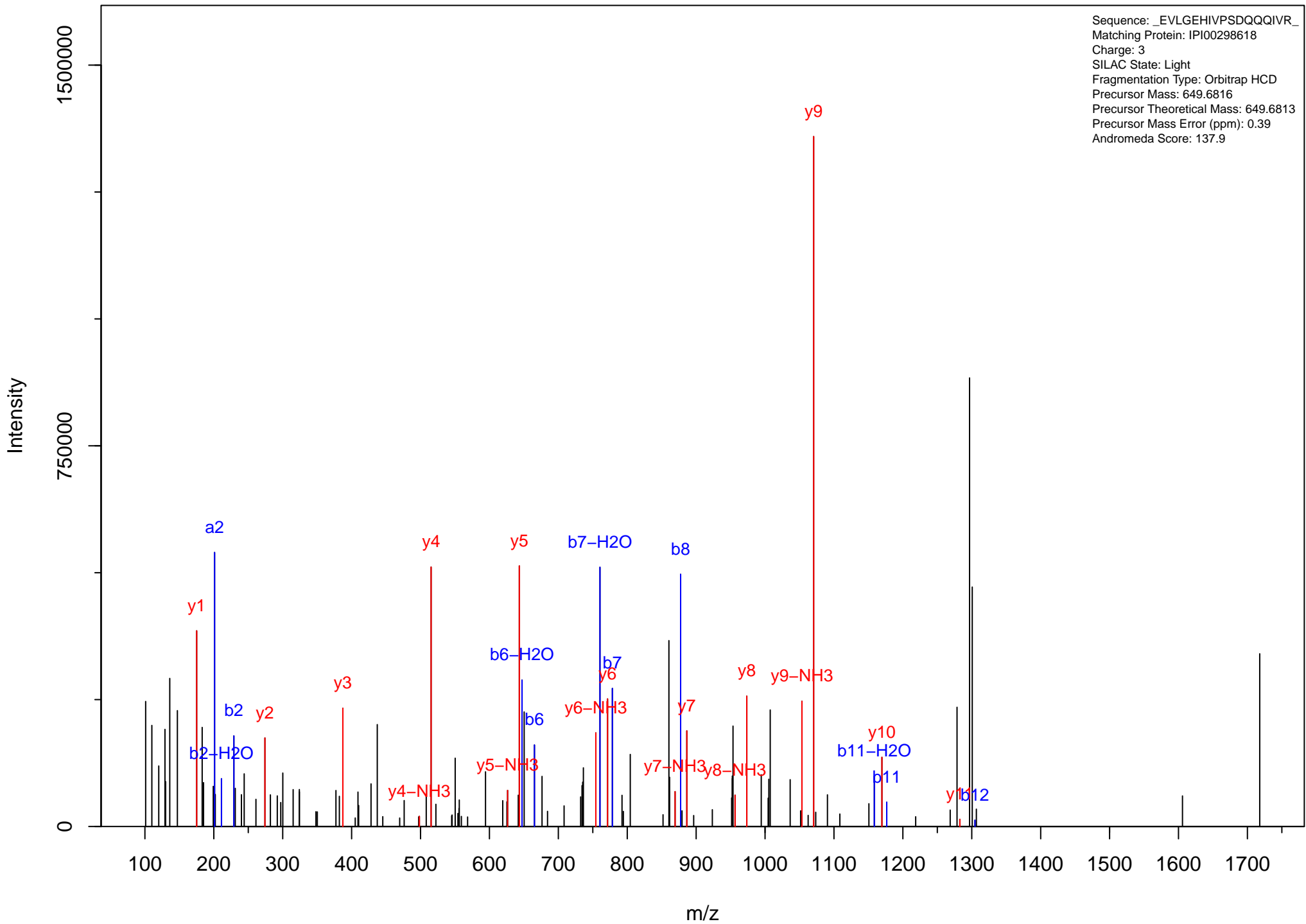


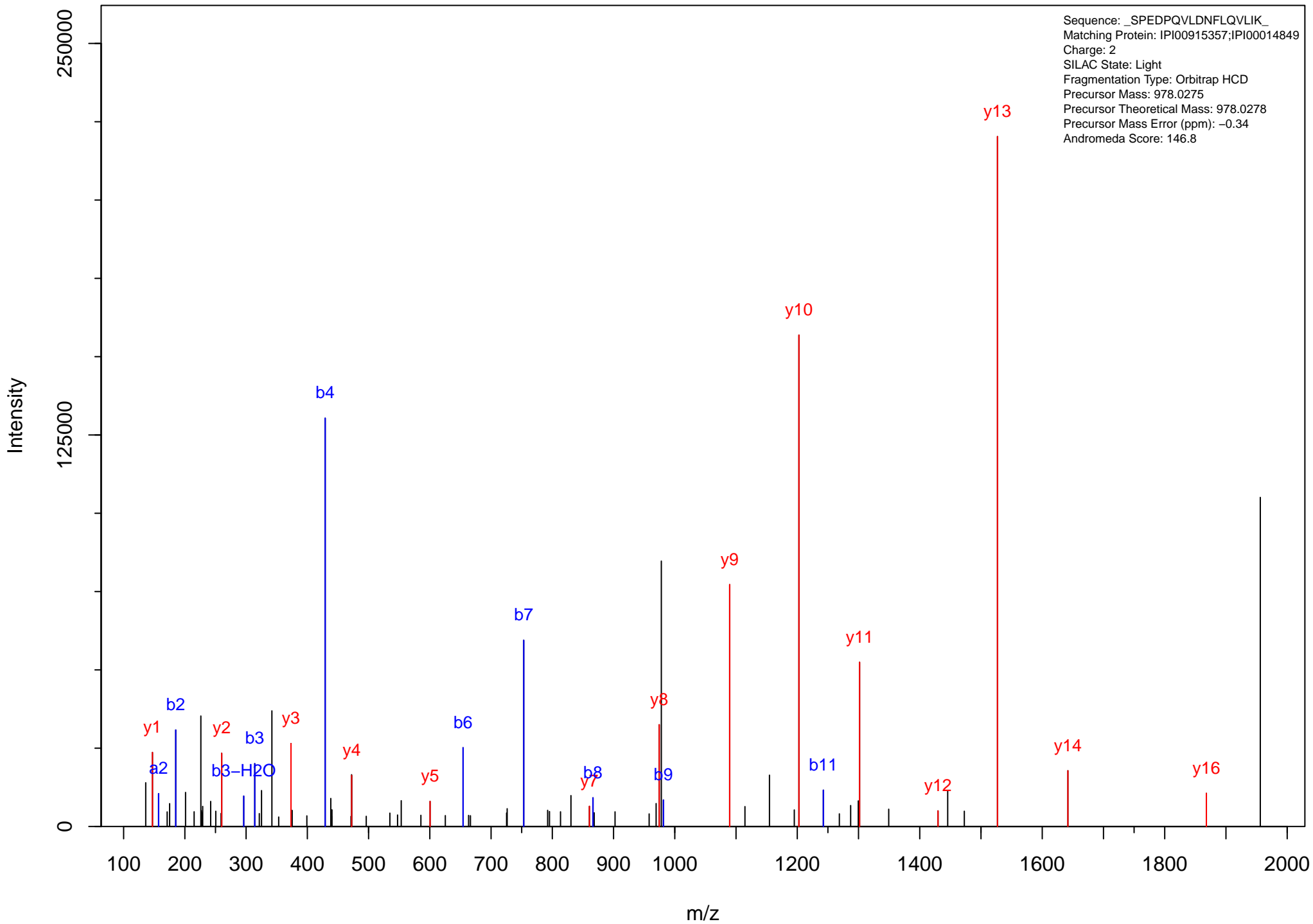


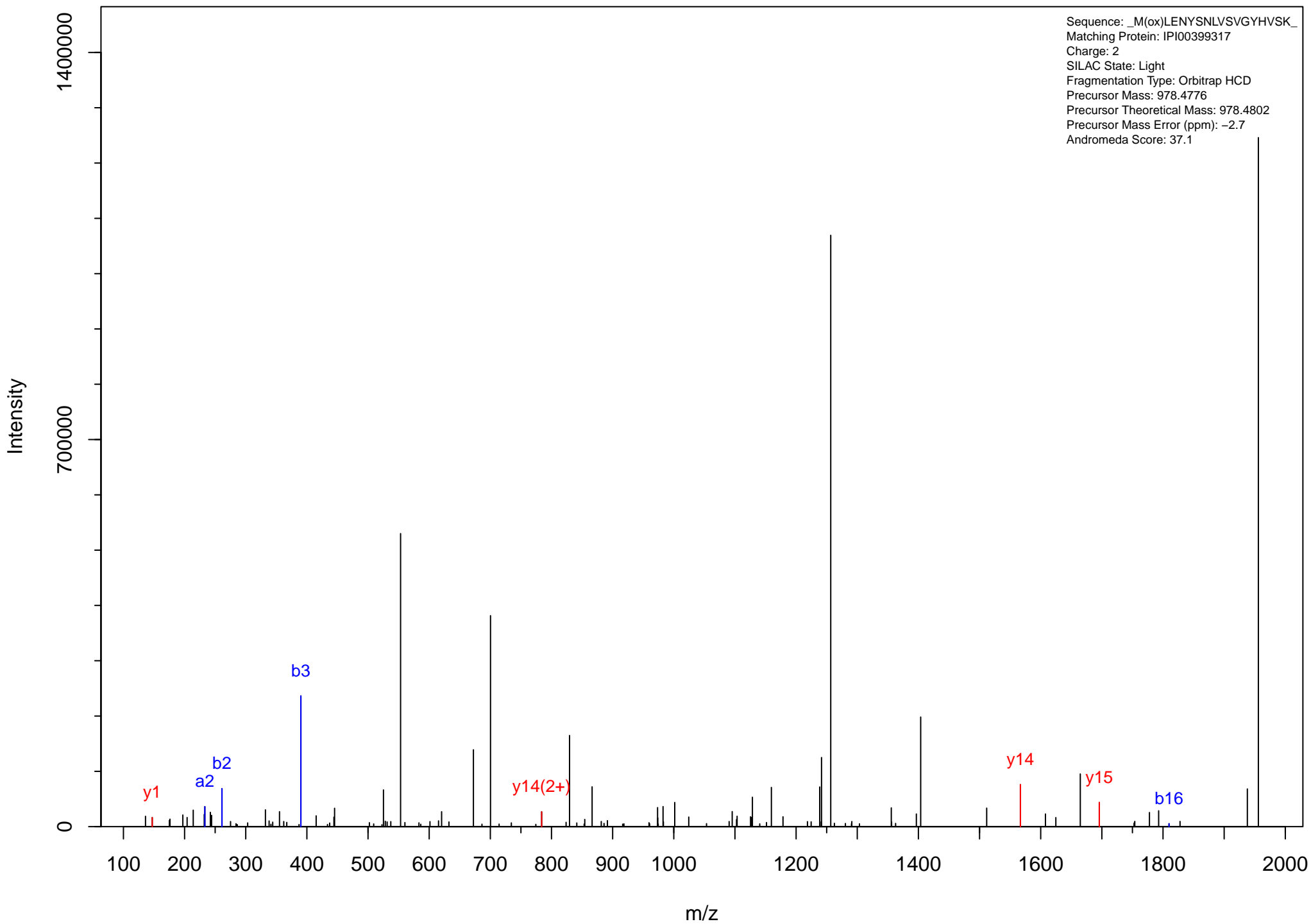


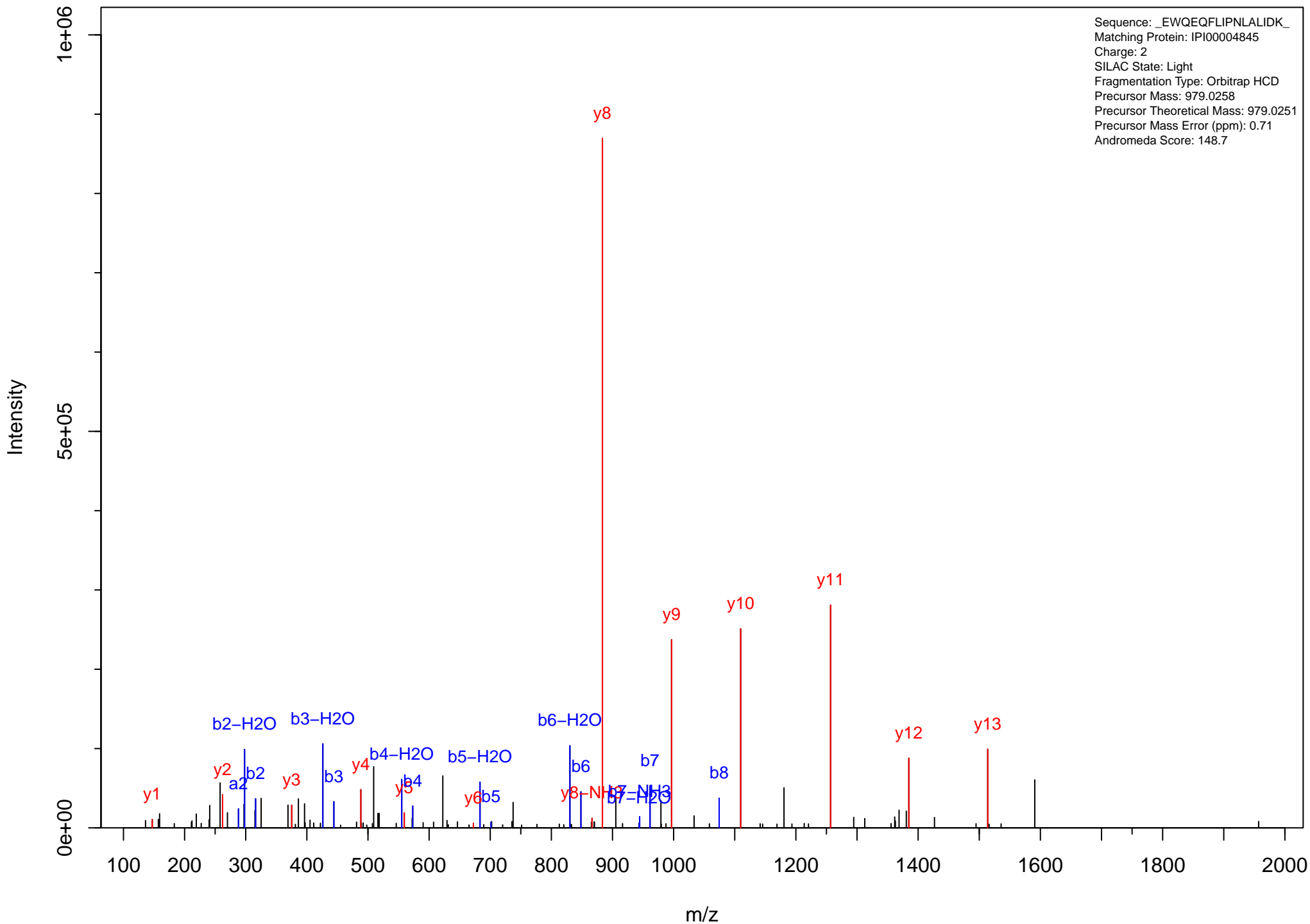


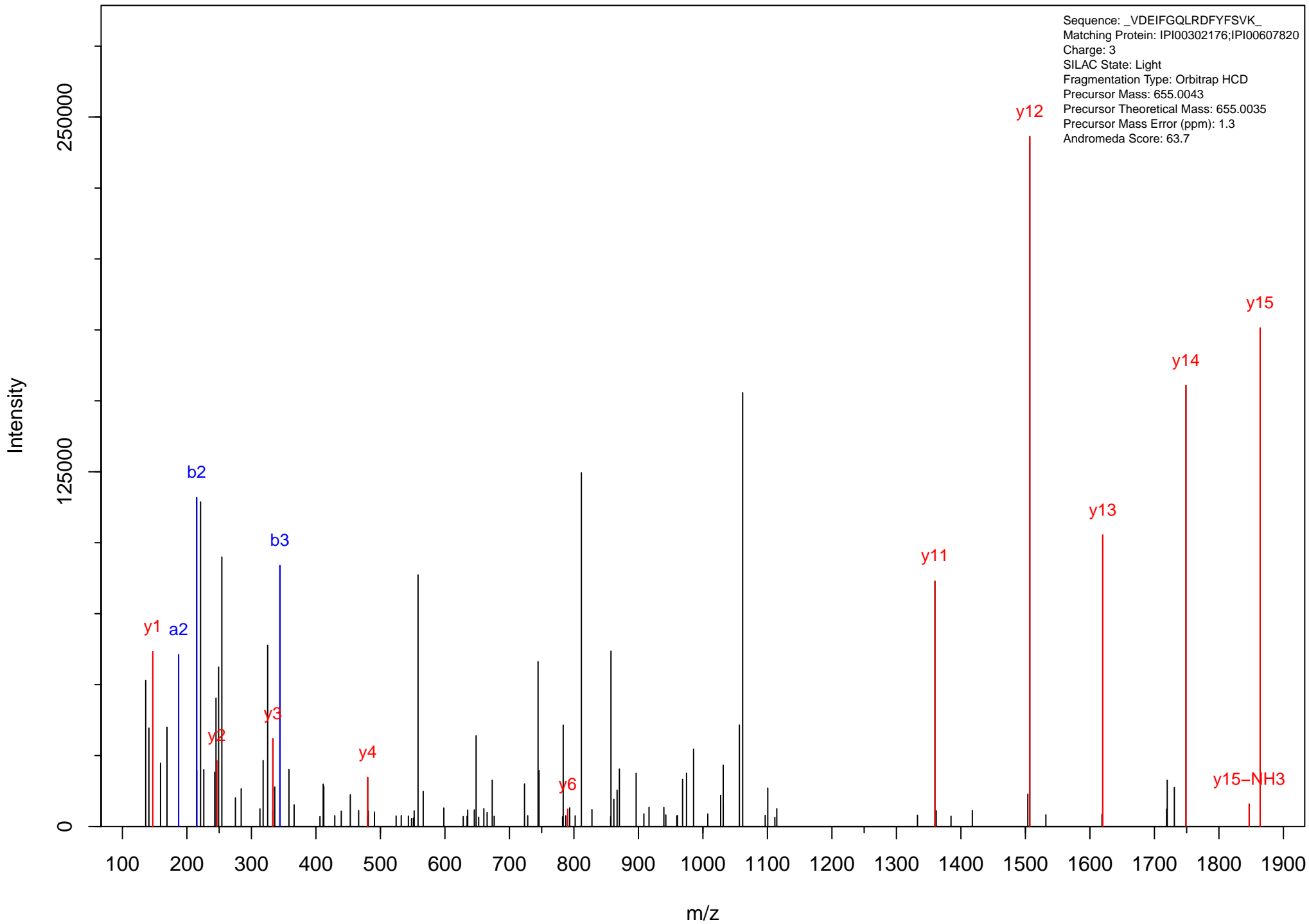


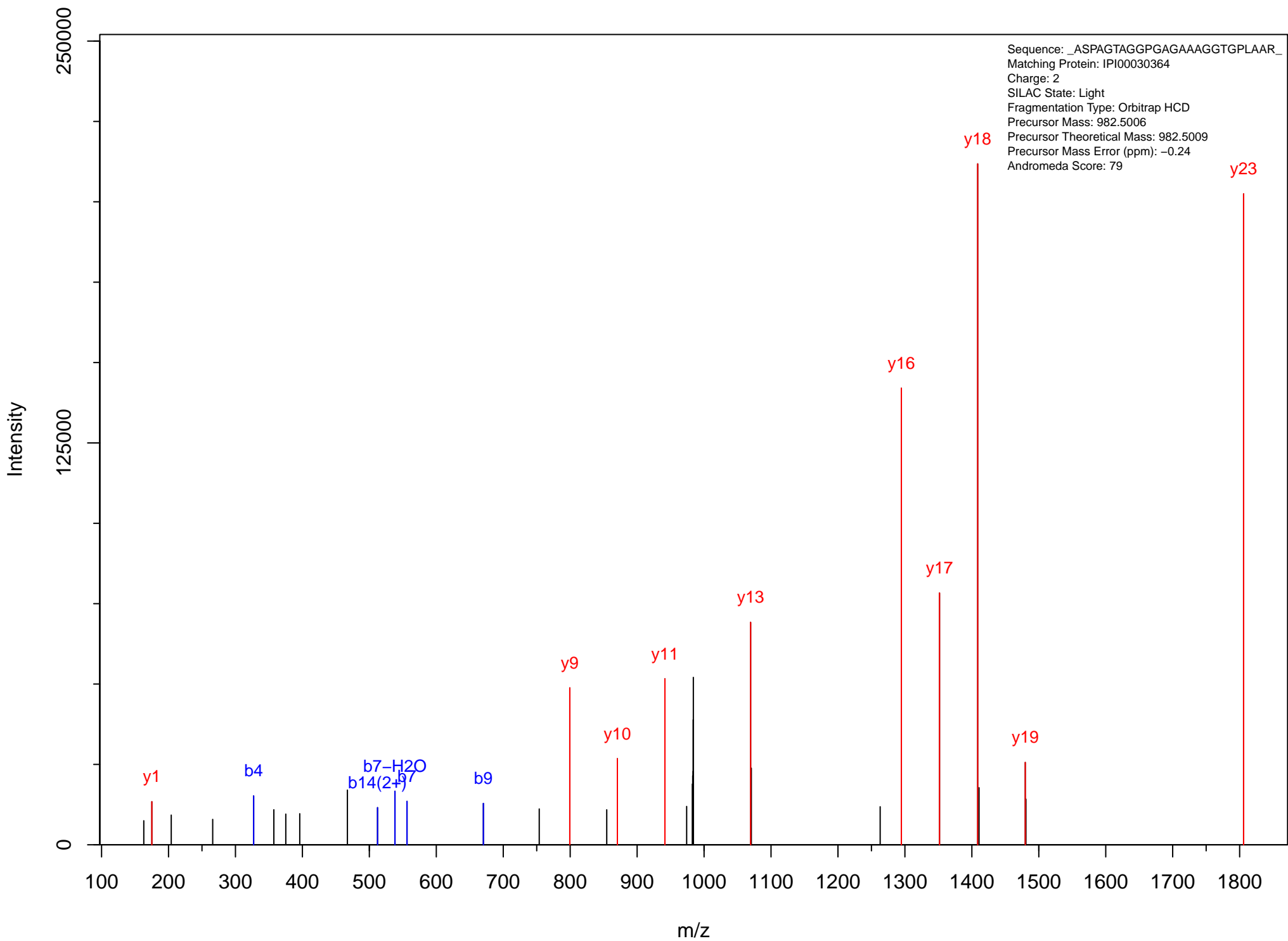


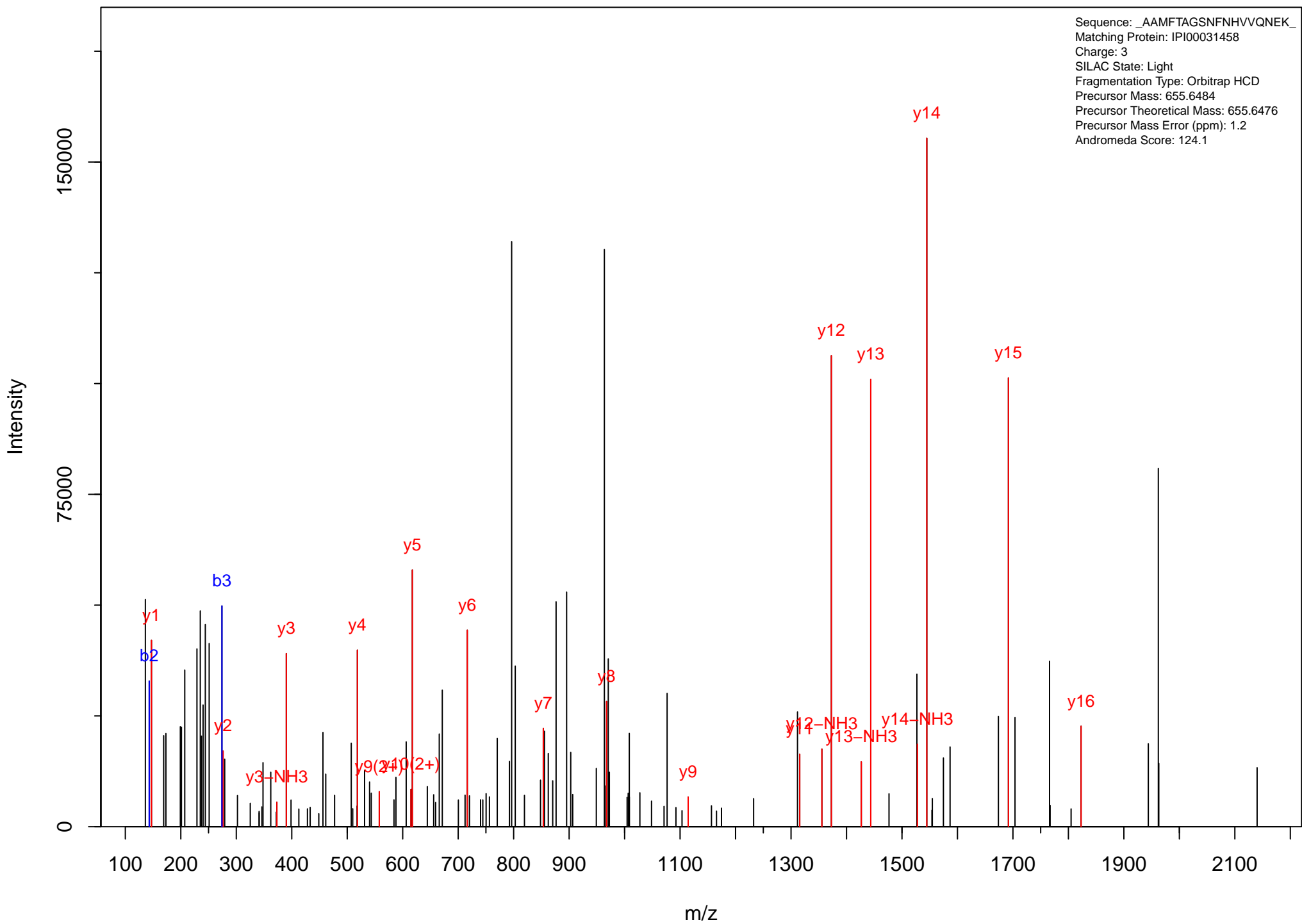




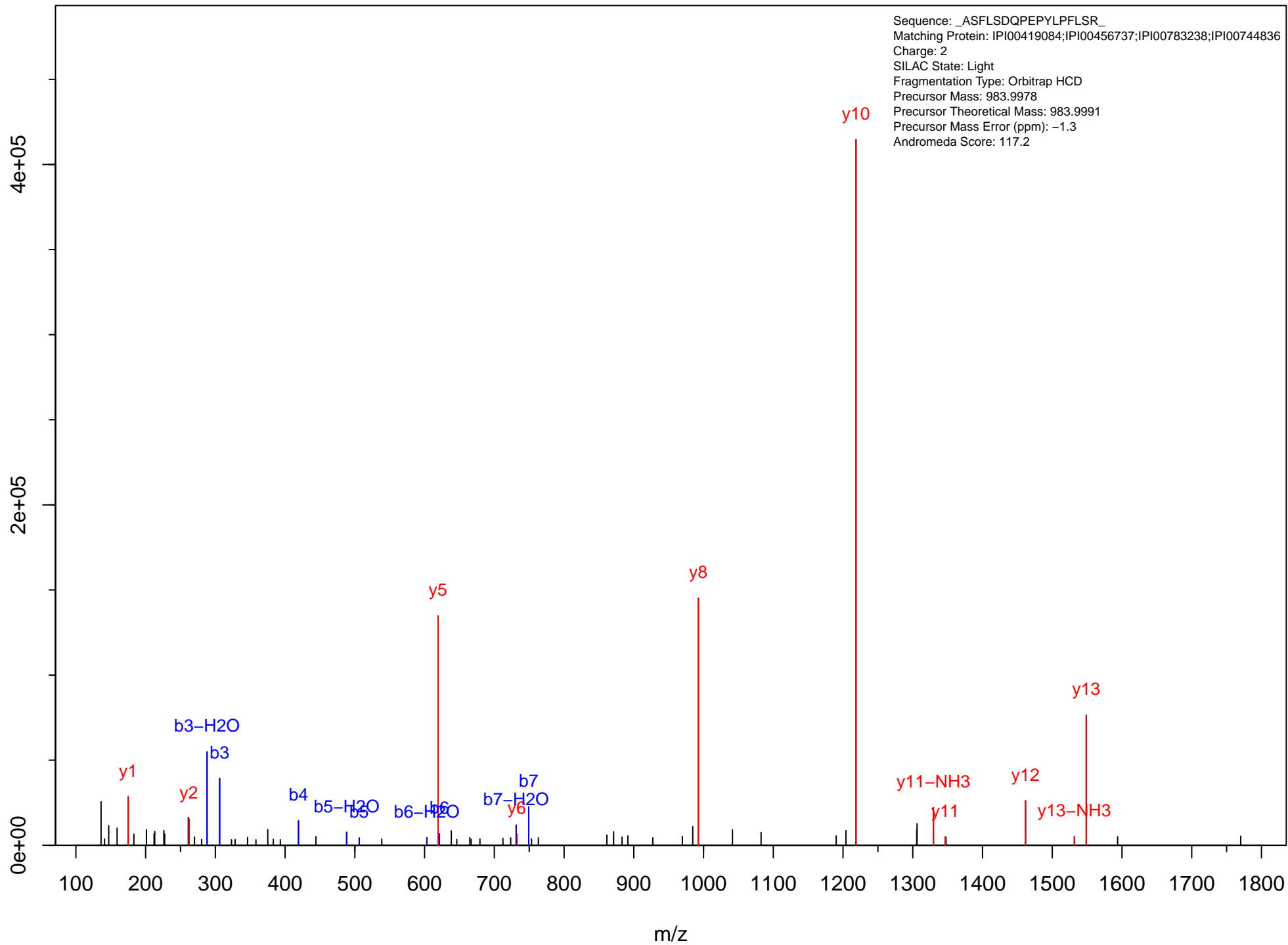


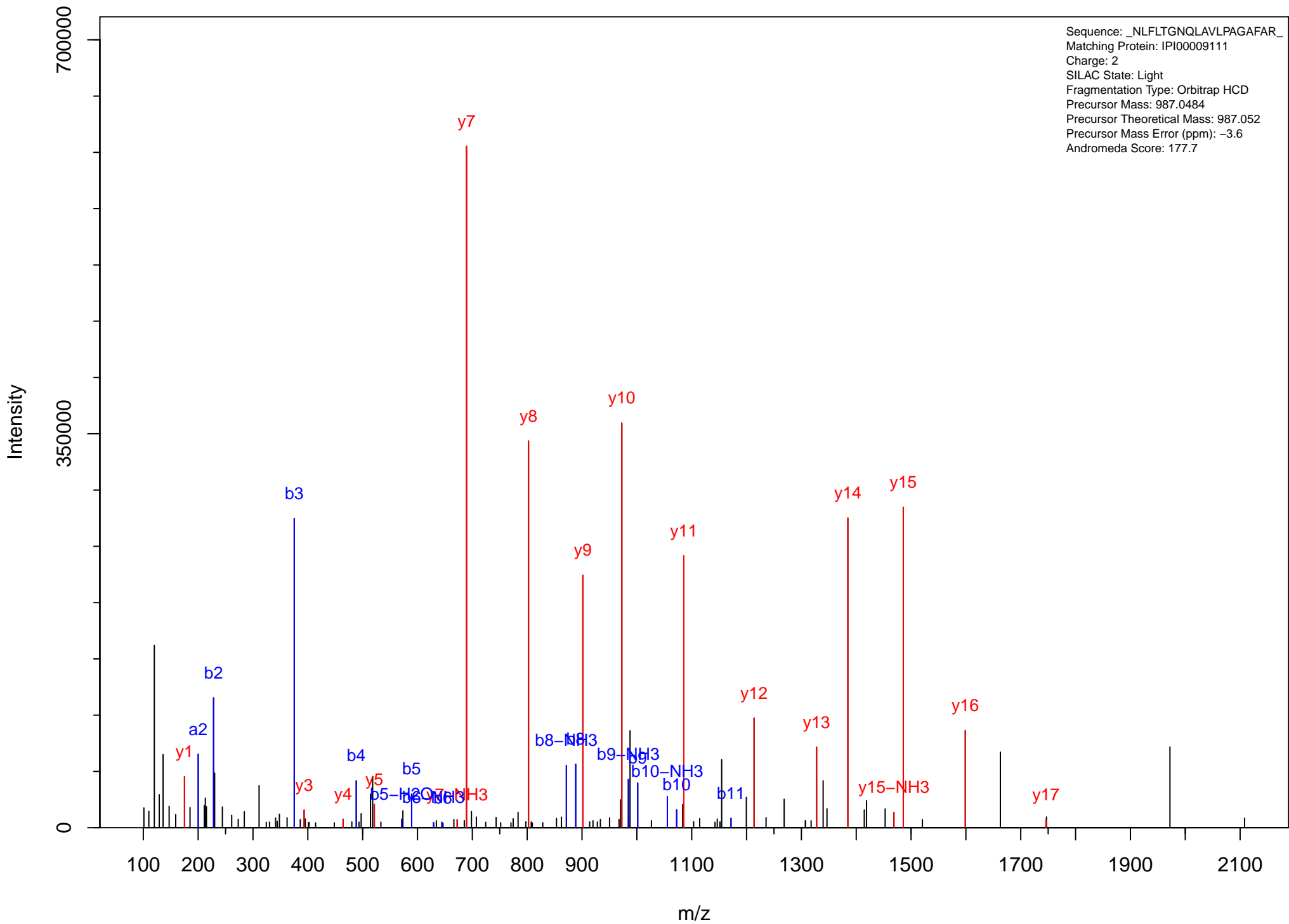


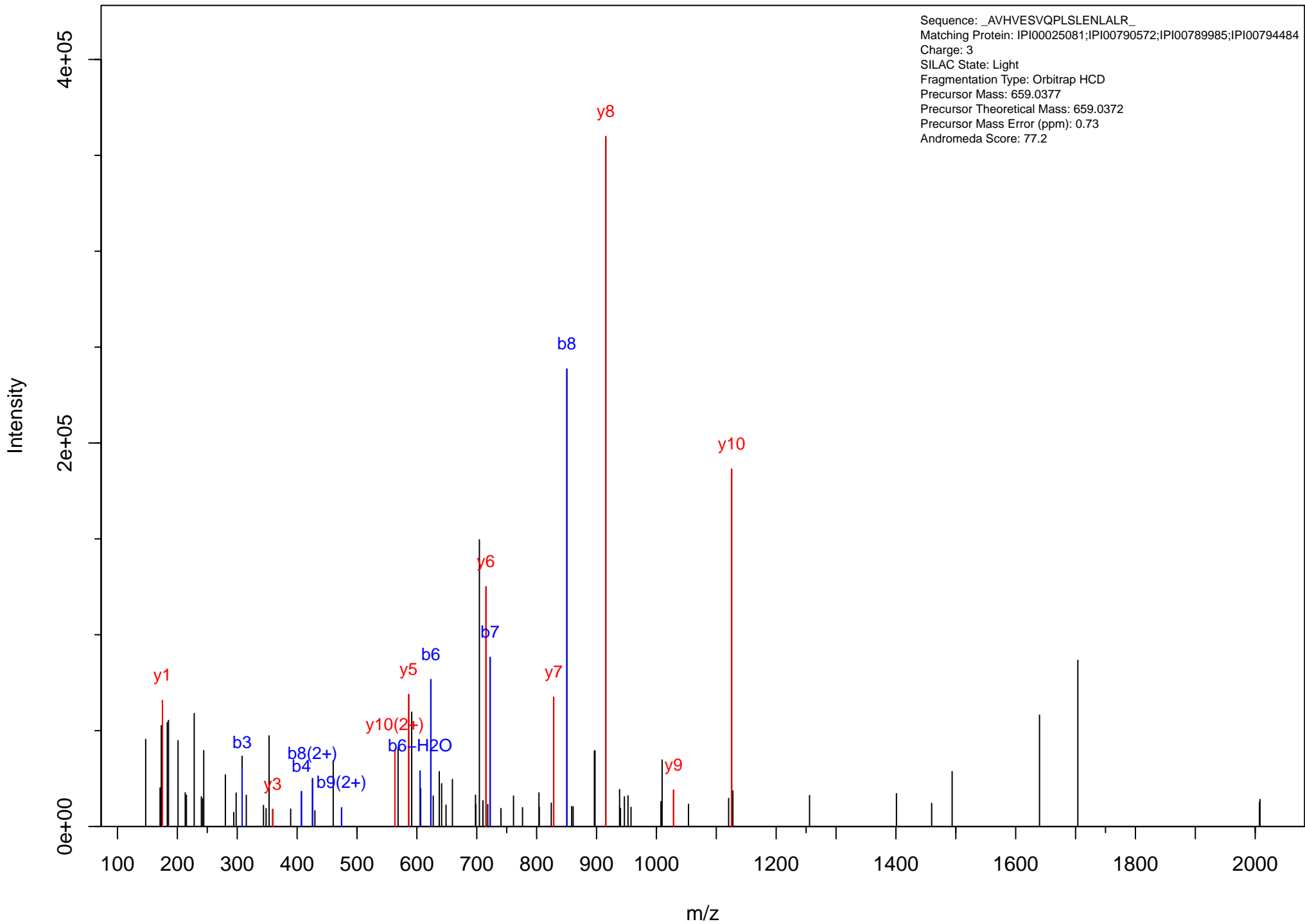


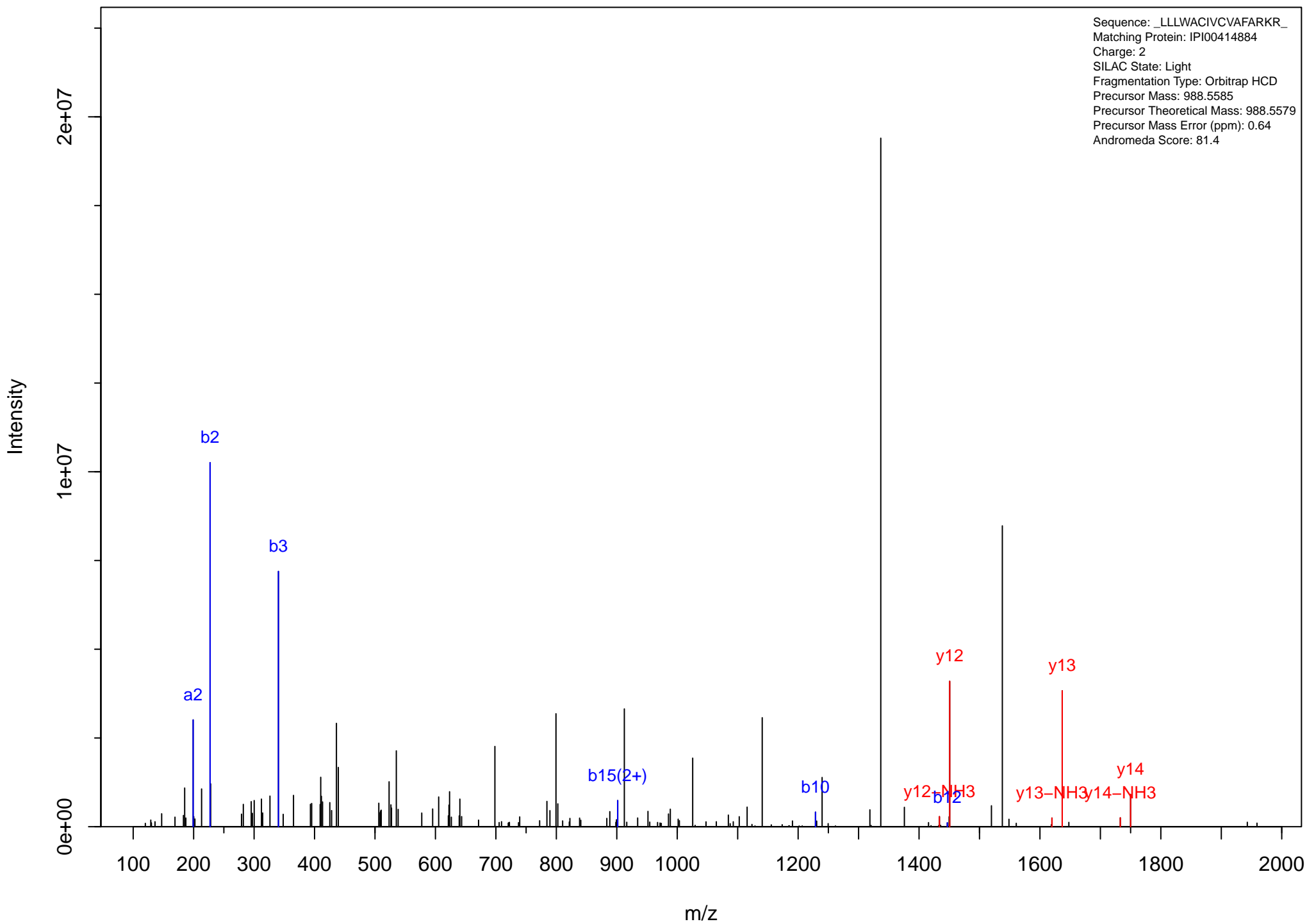


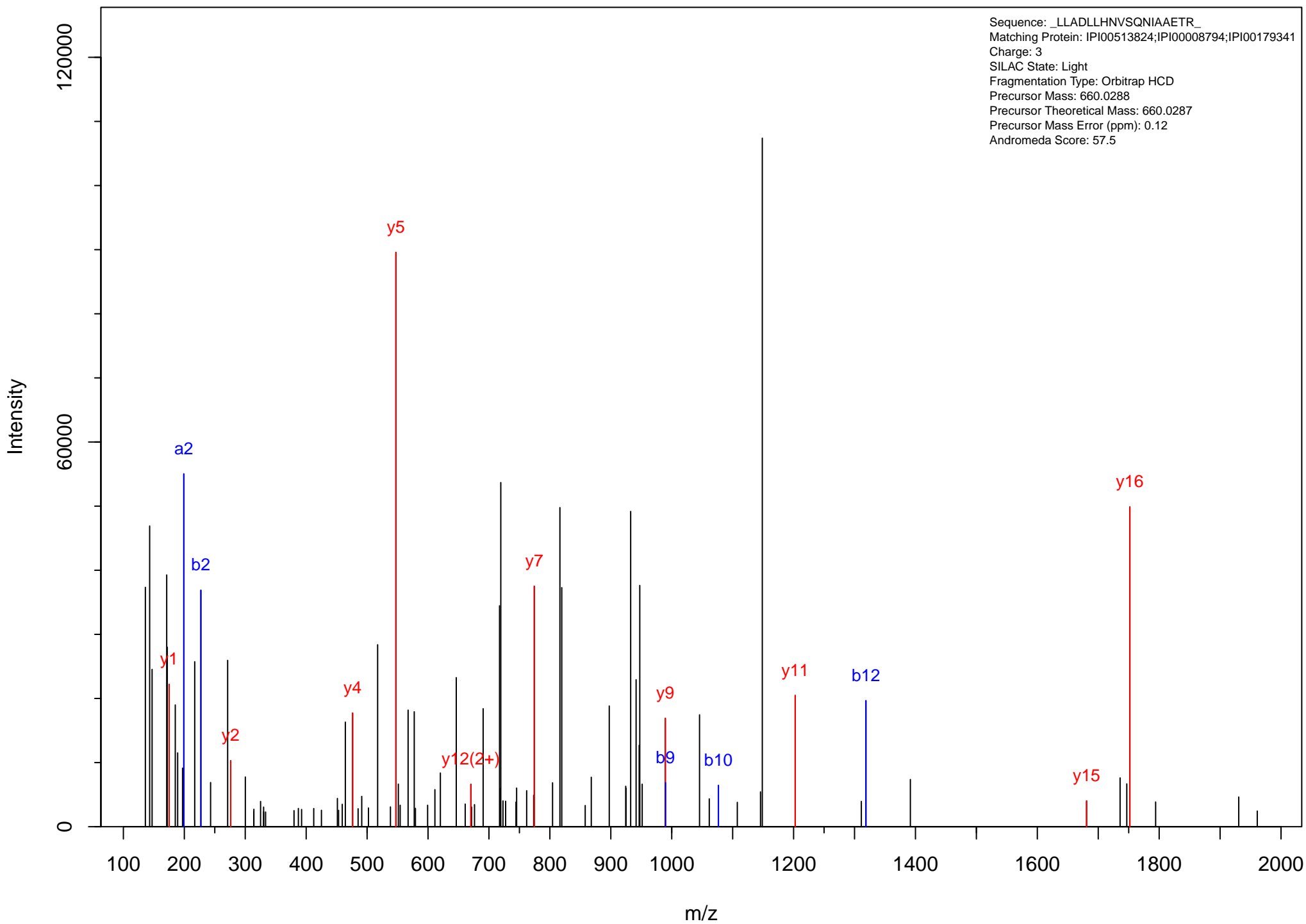
Intensity

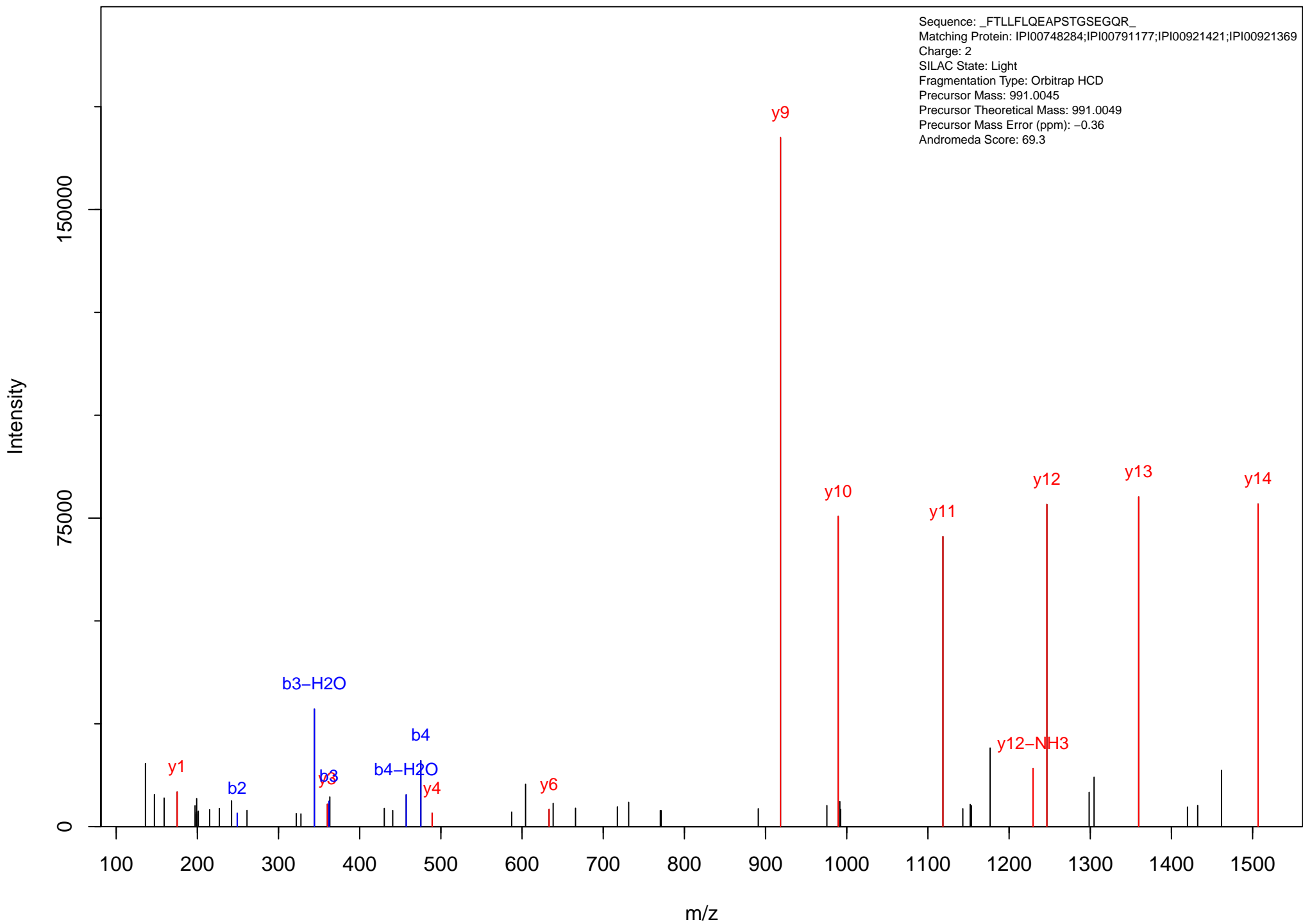


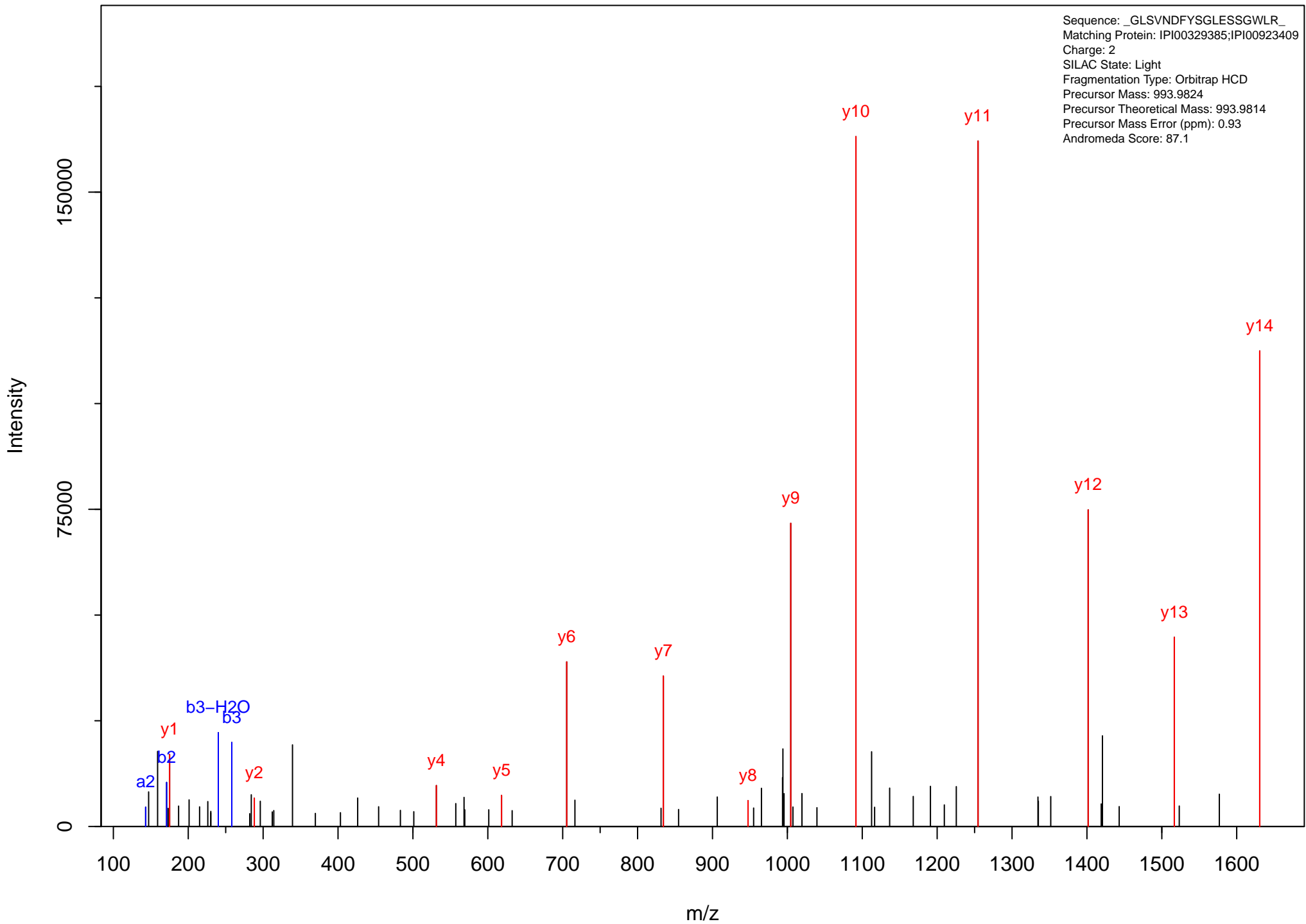


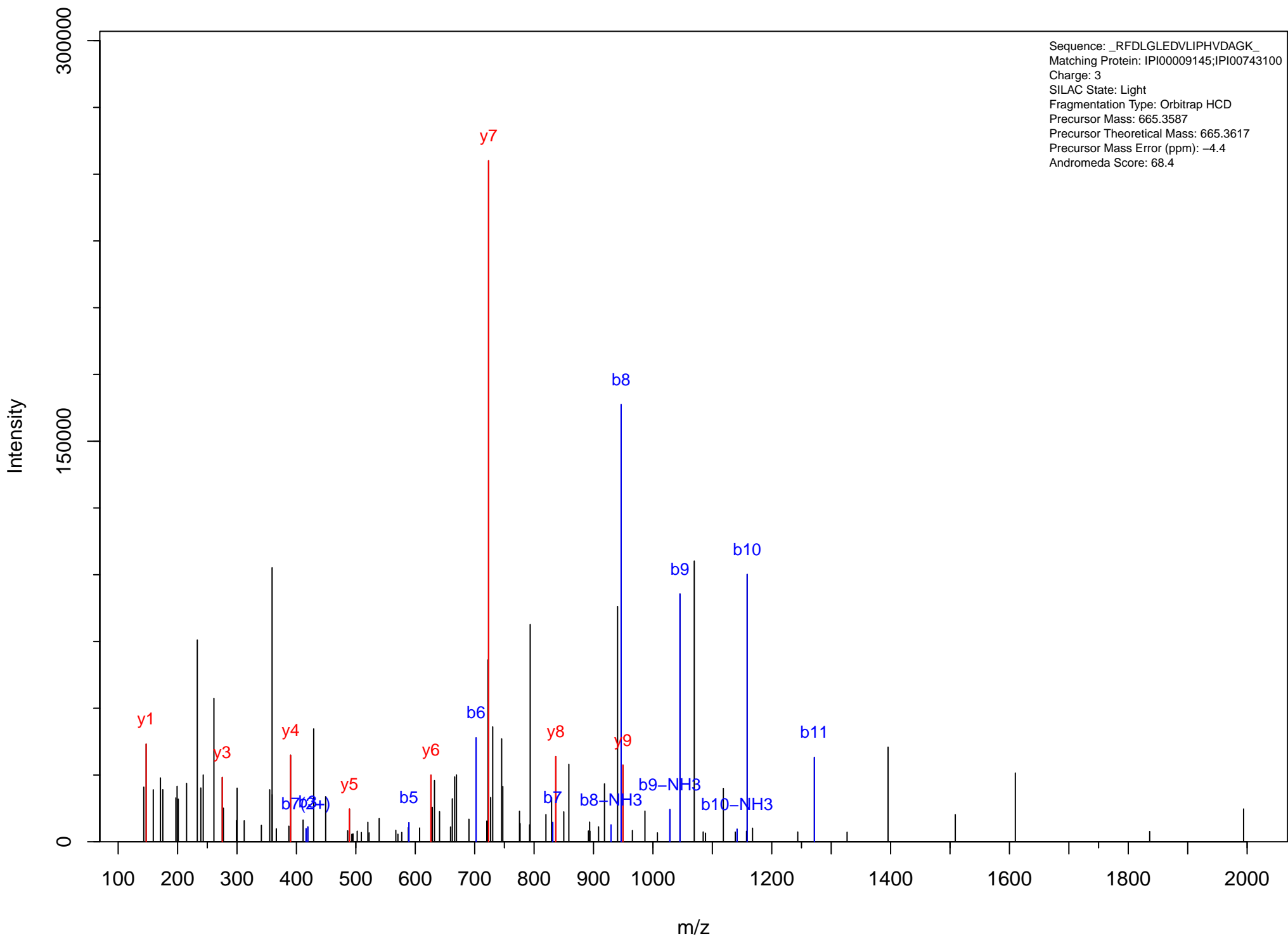


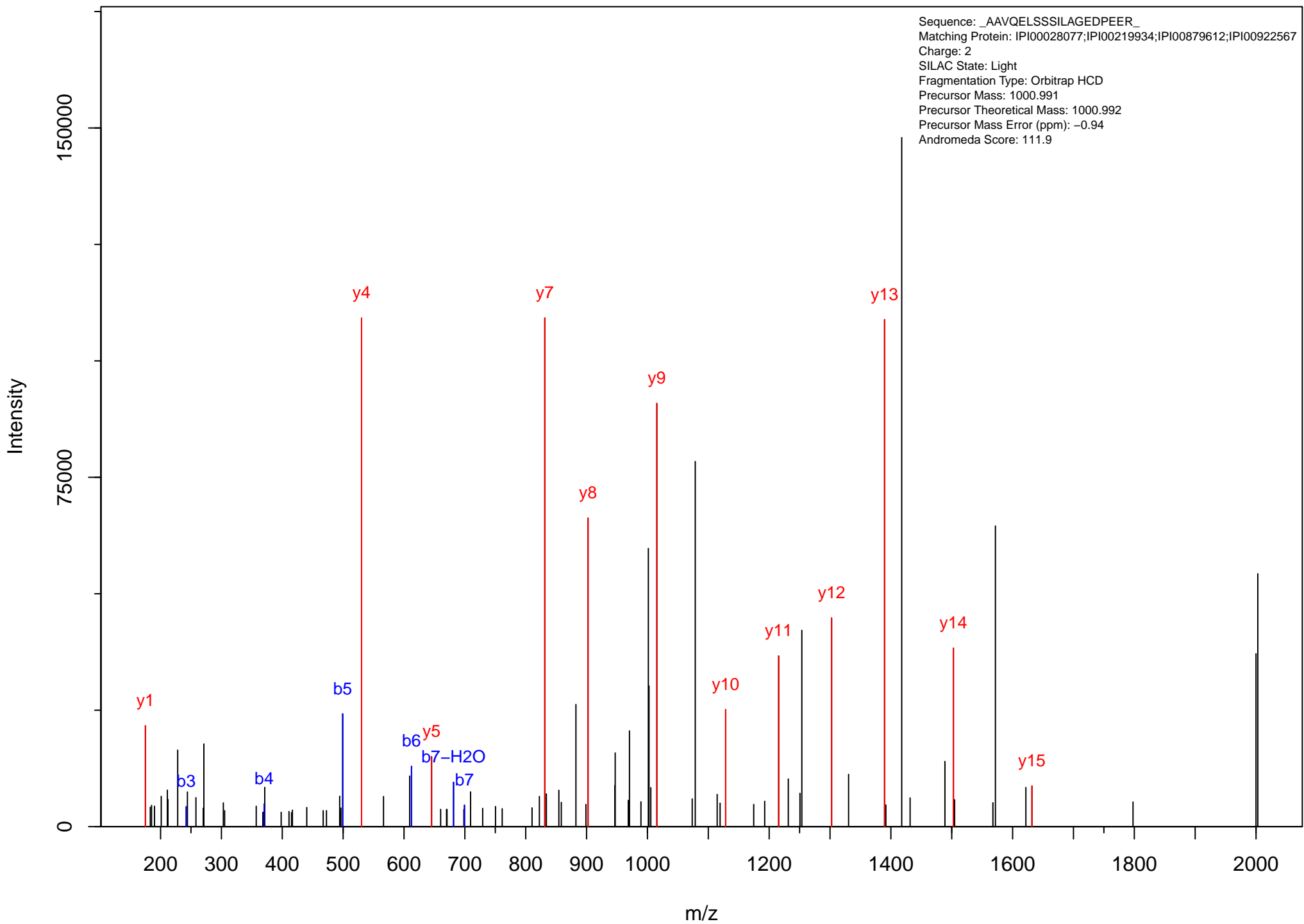


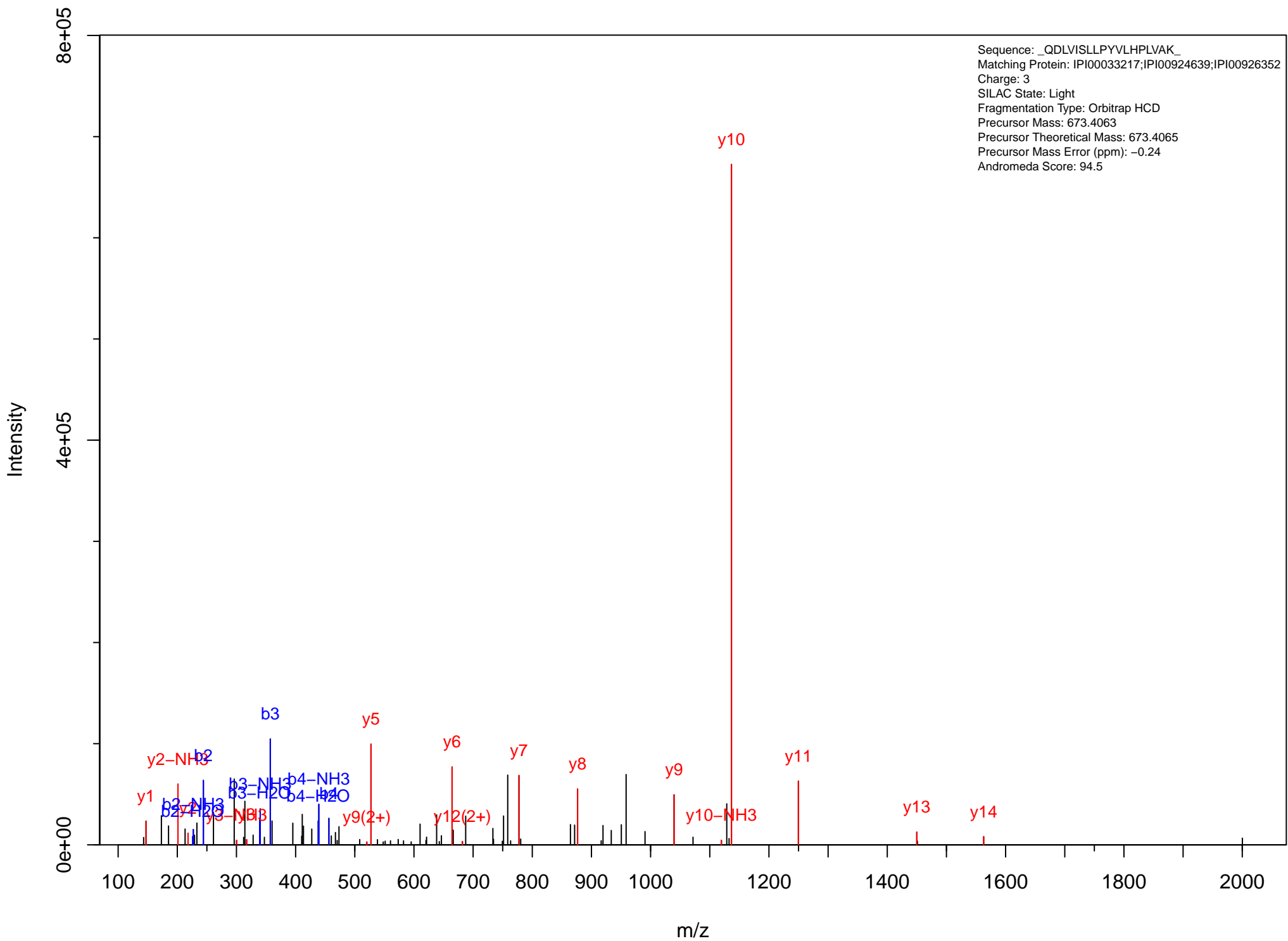


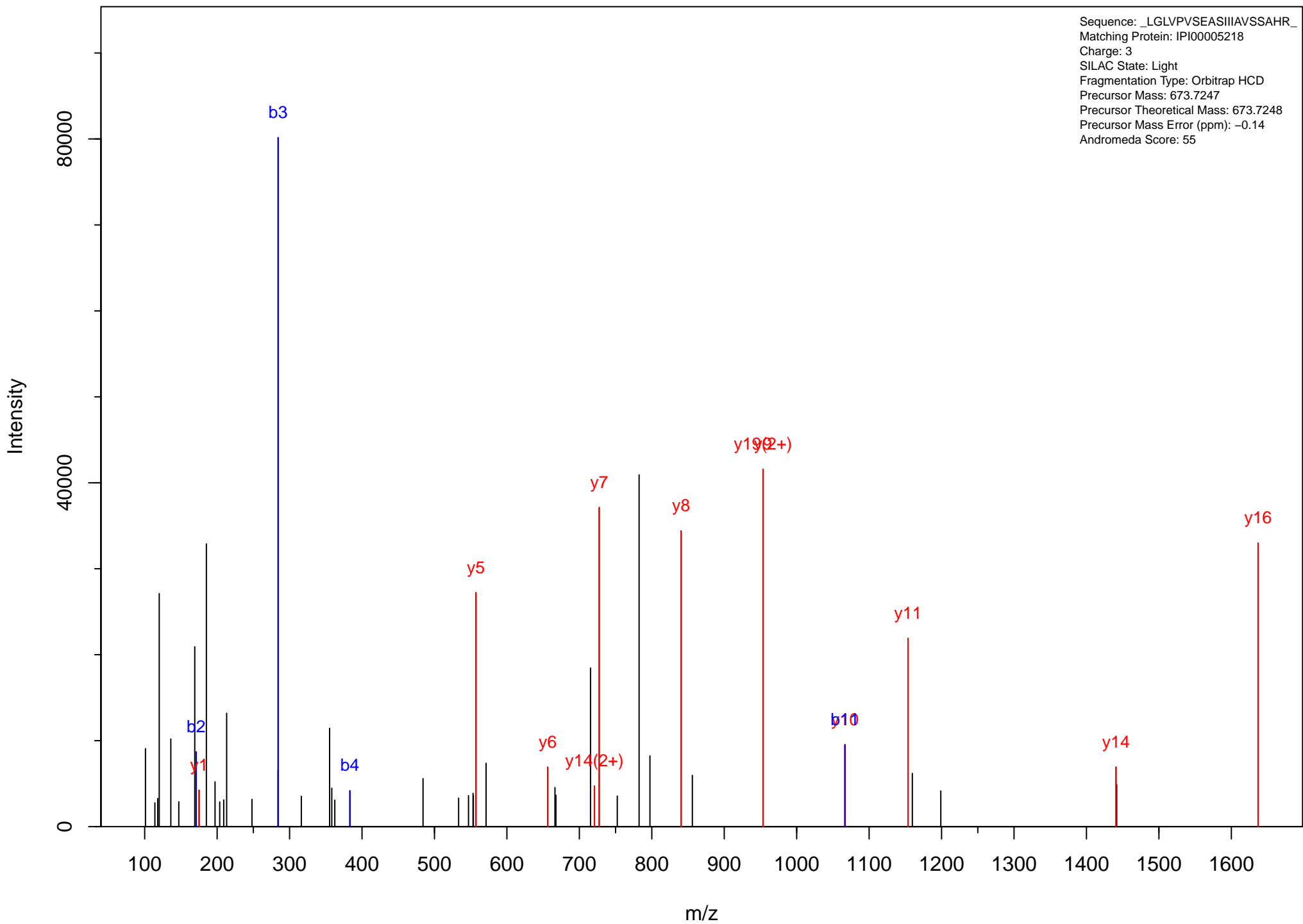


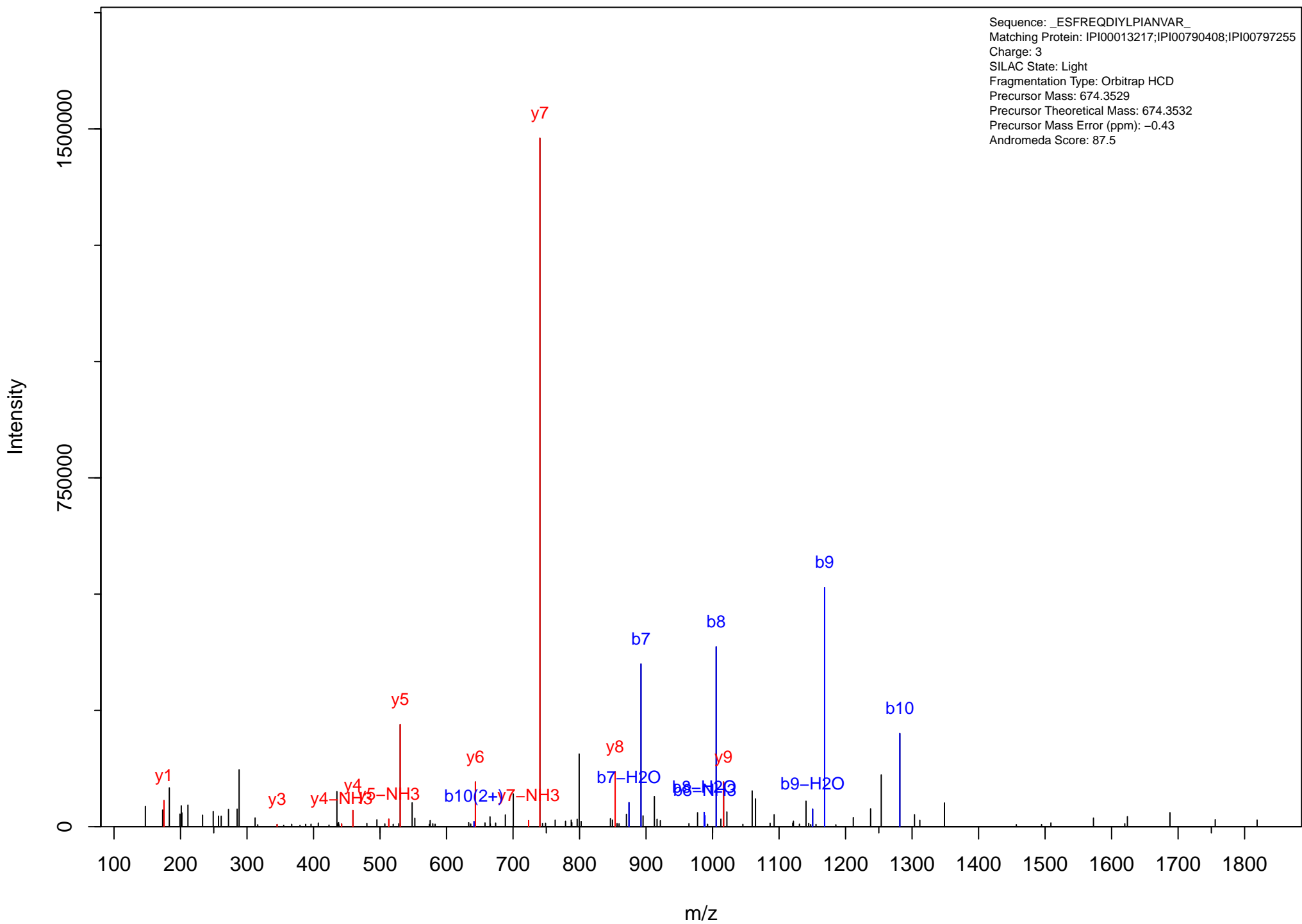


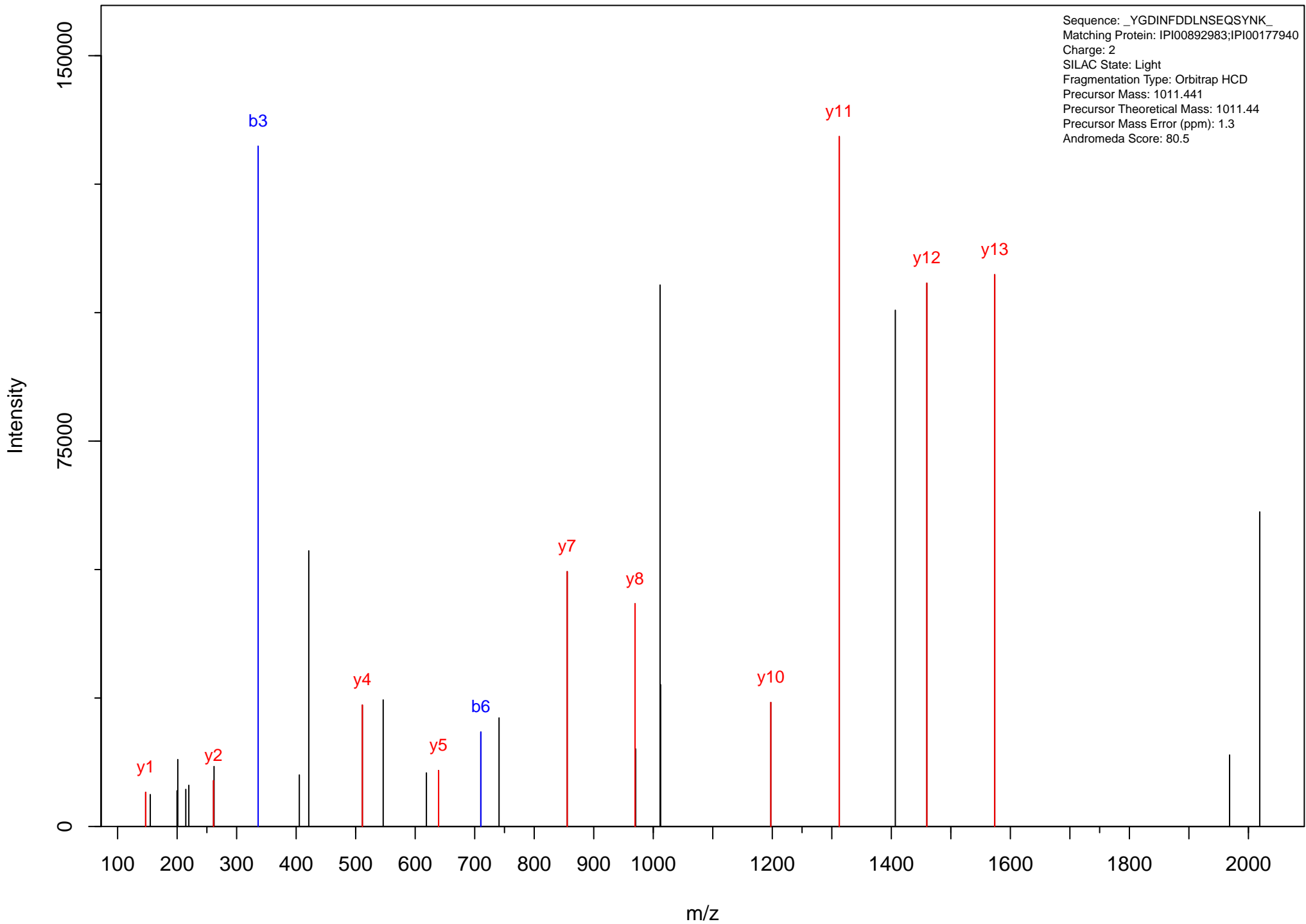


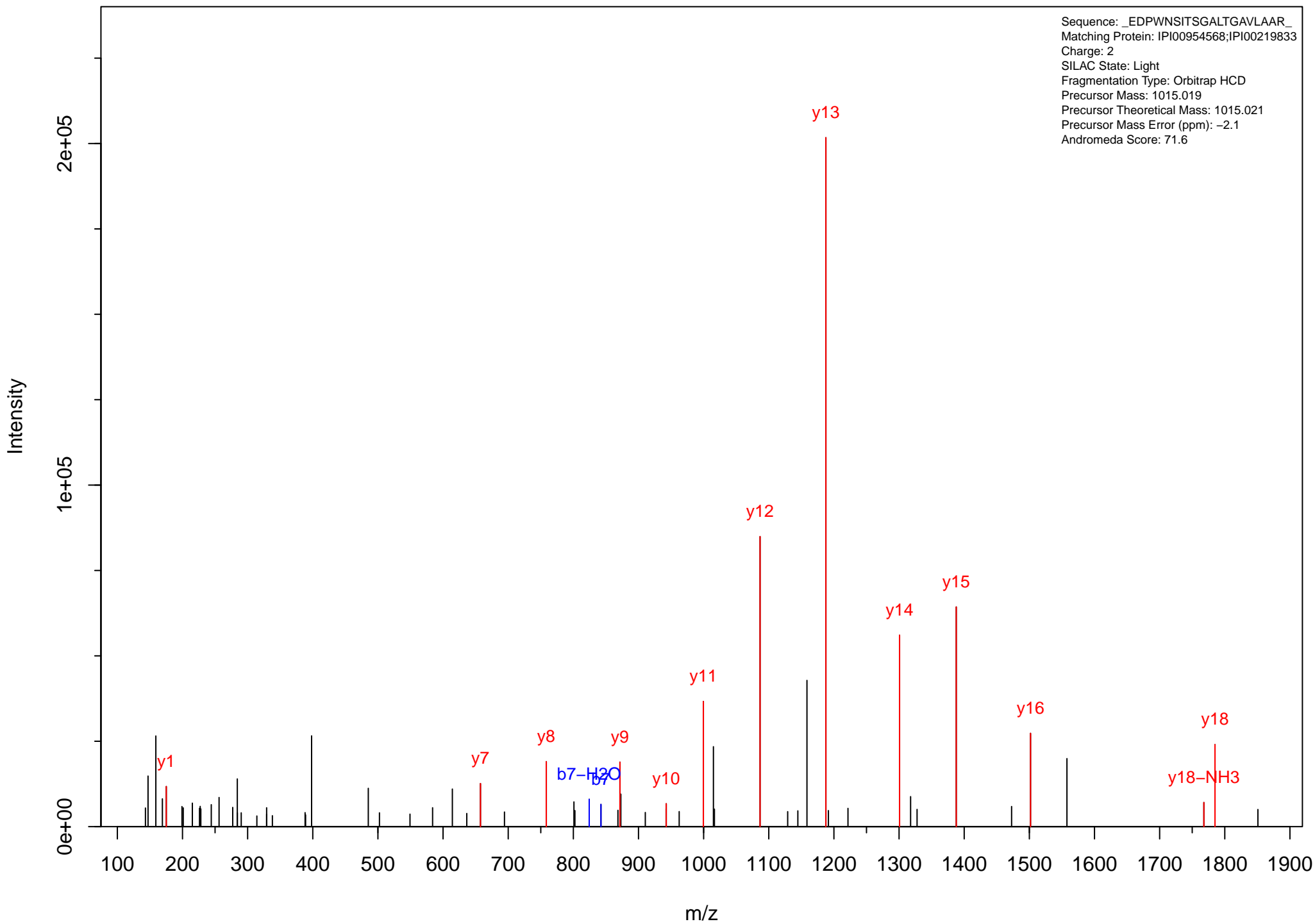


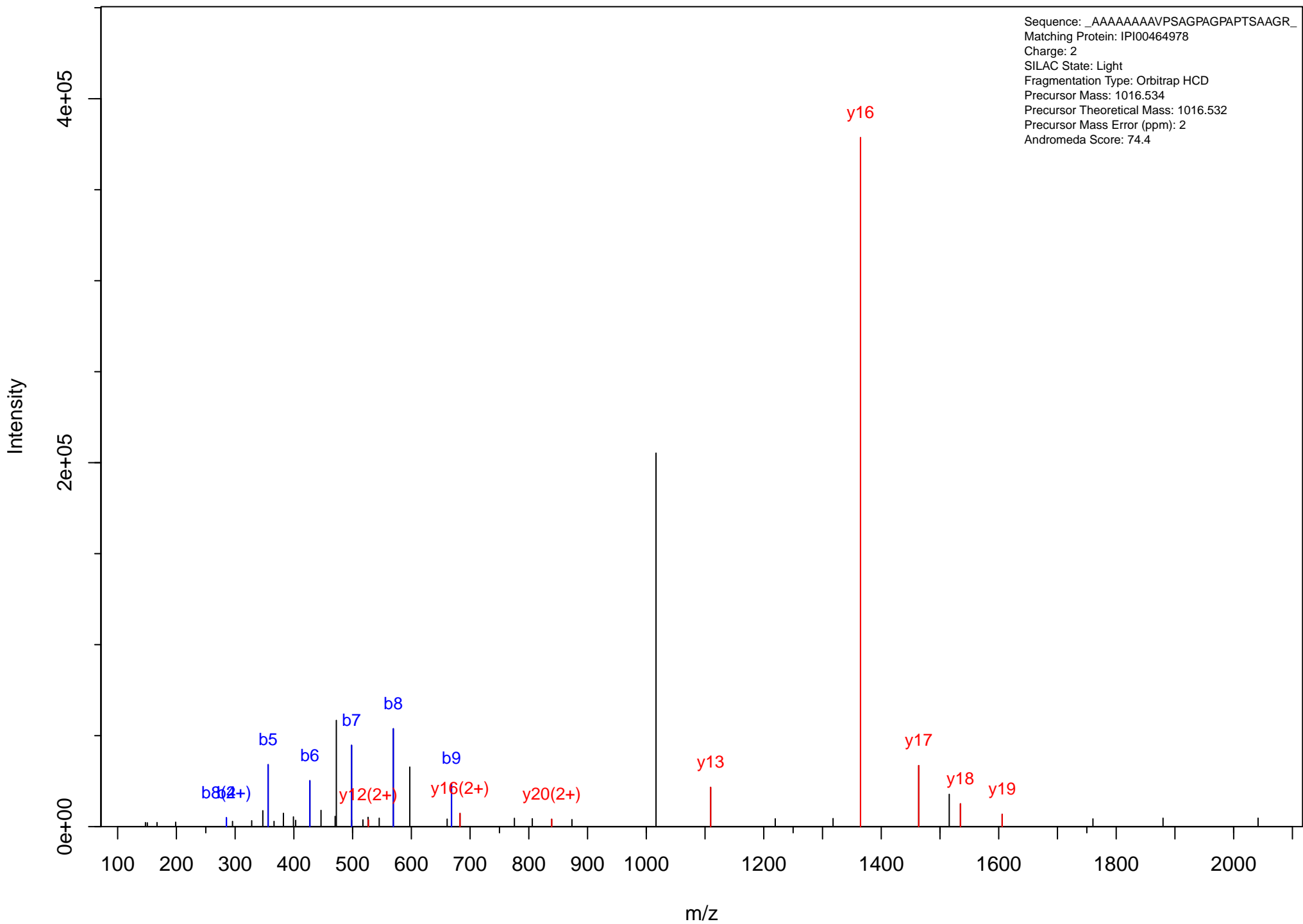


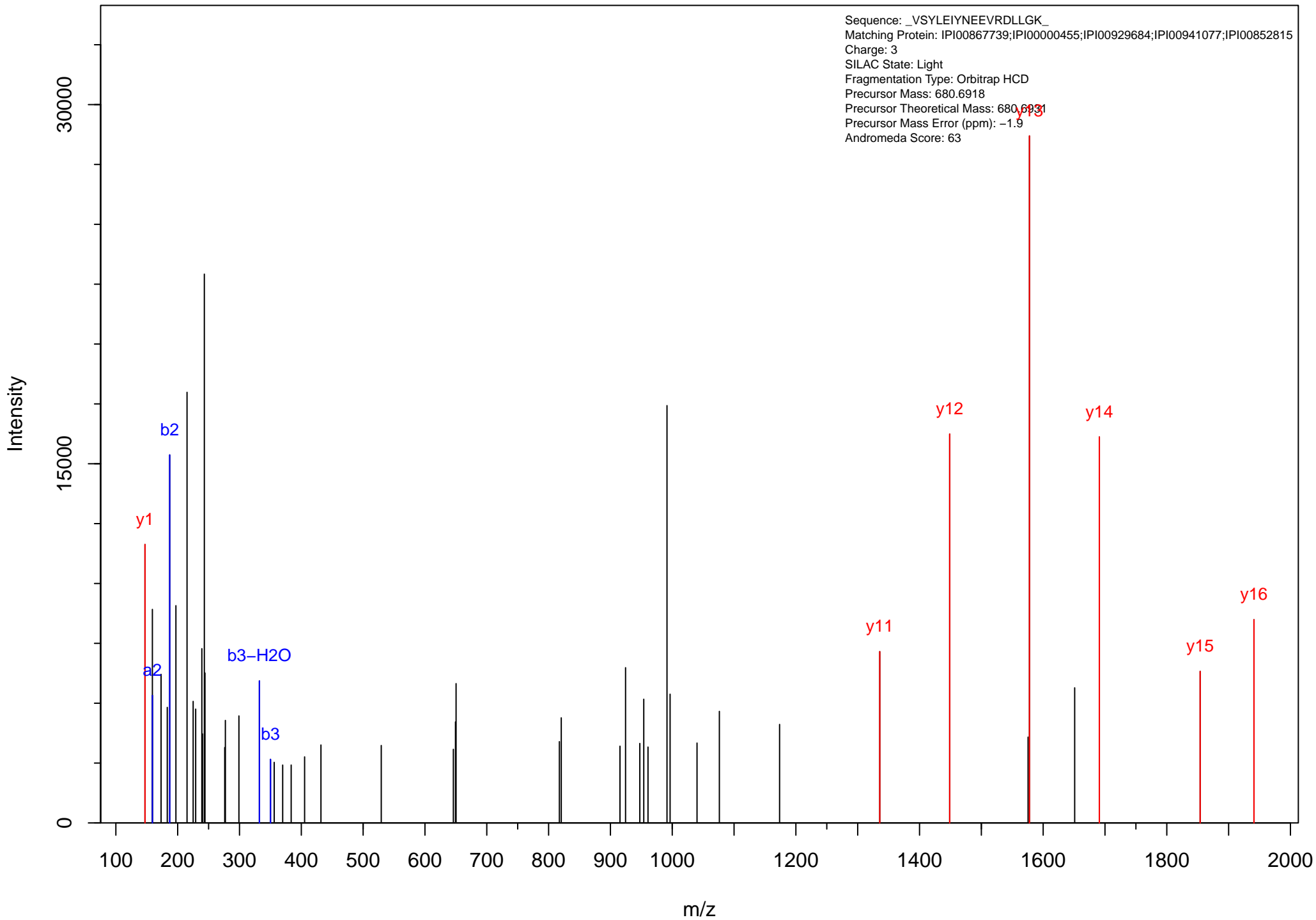


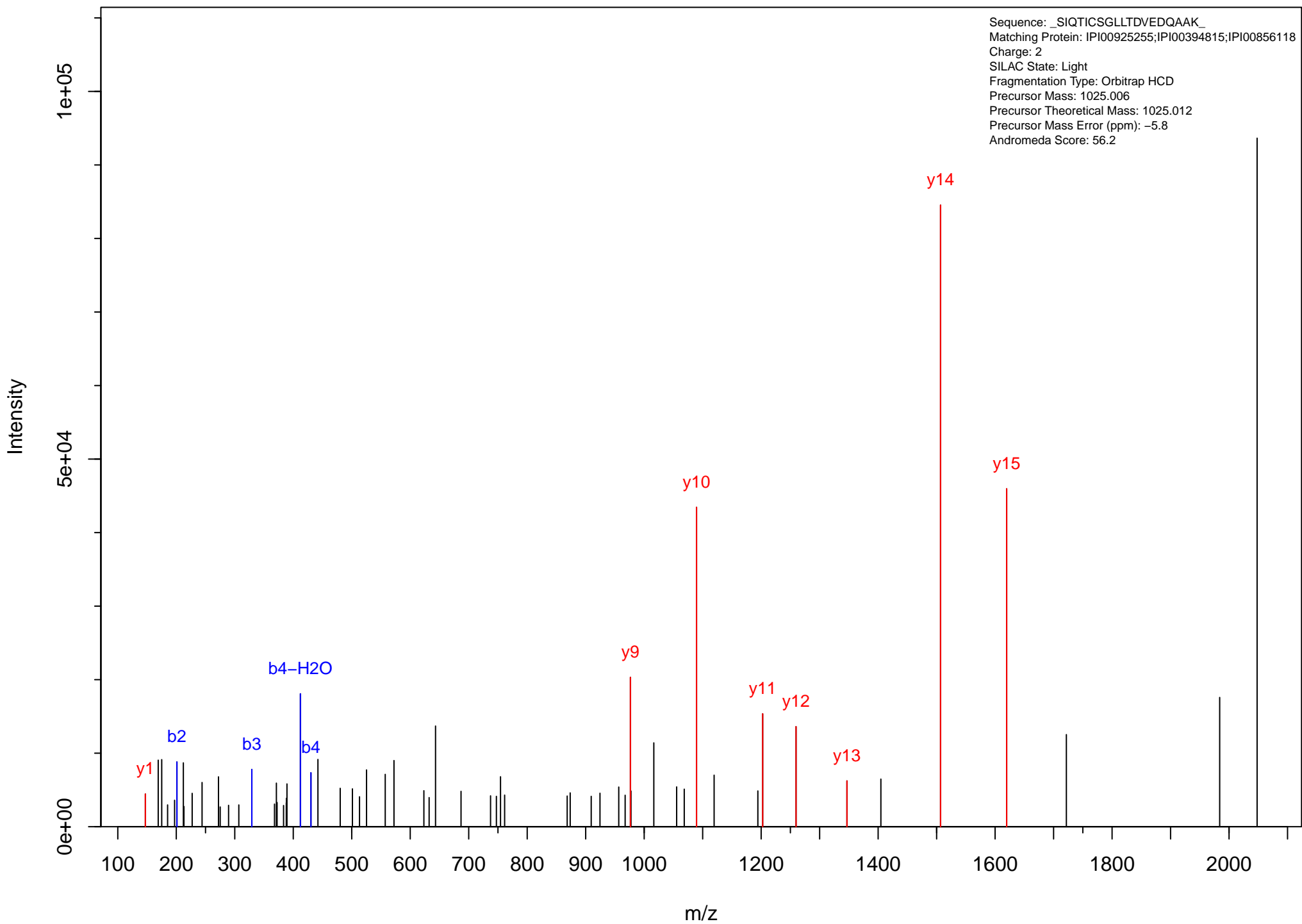


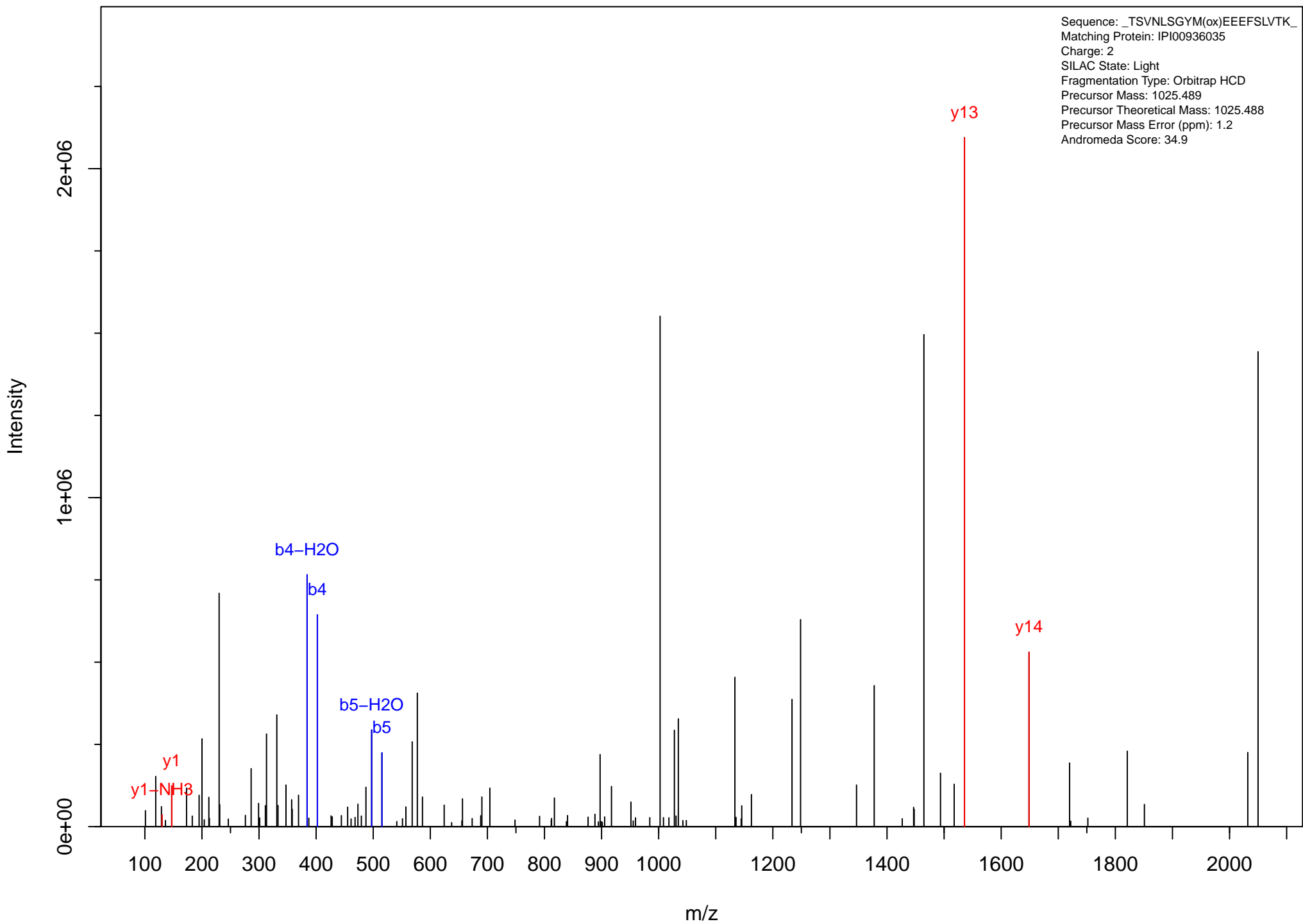


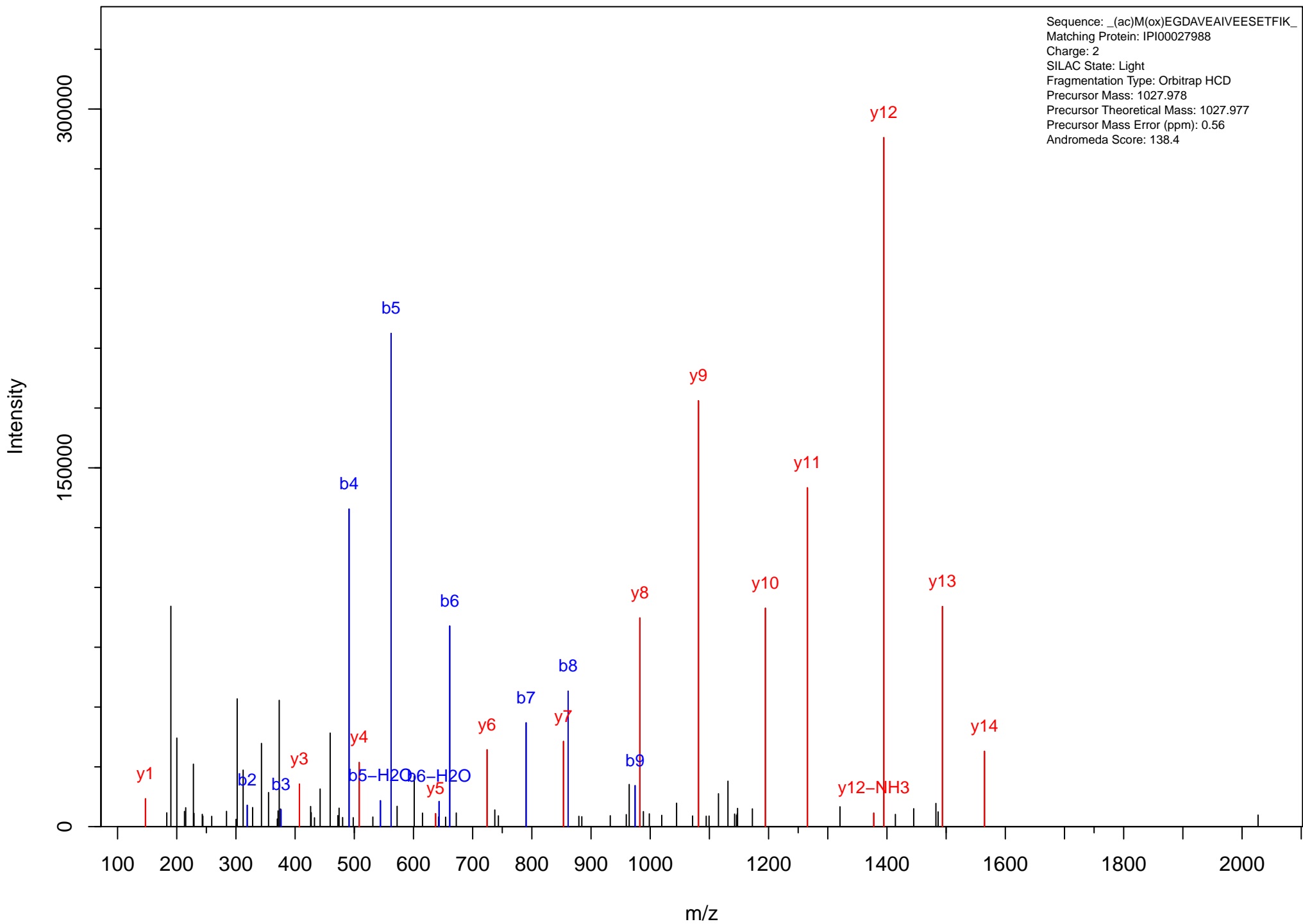


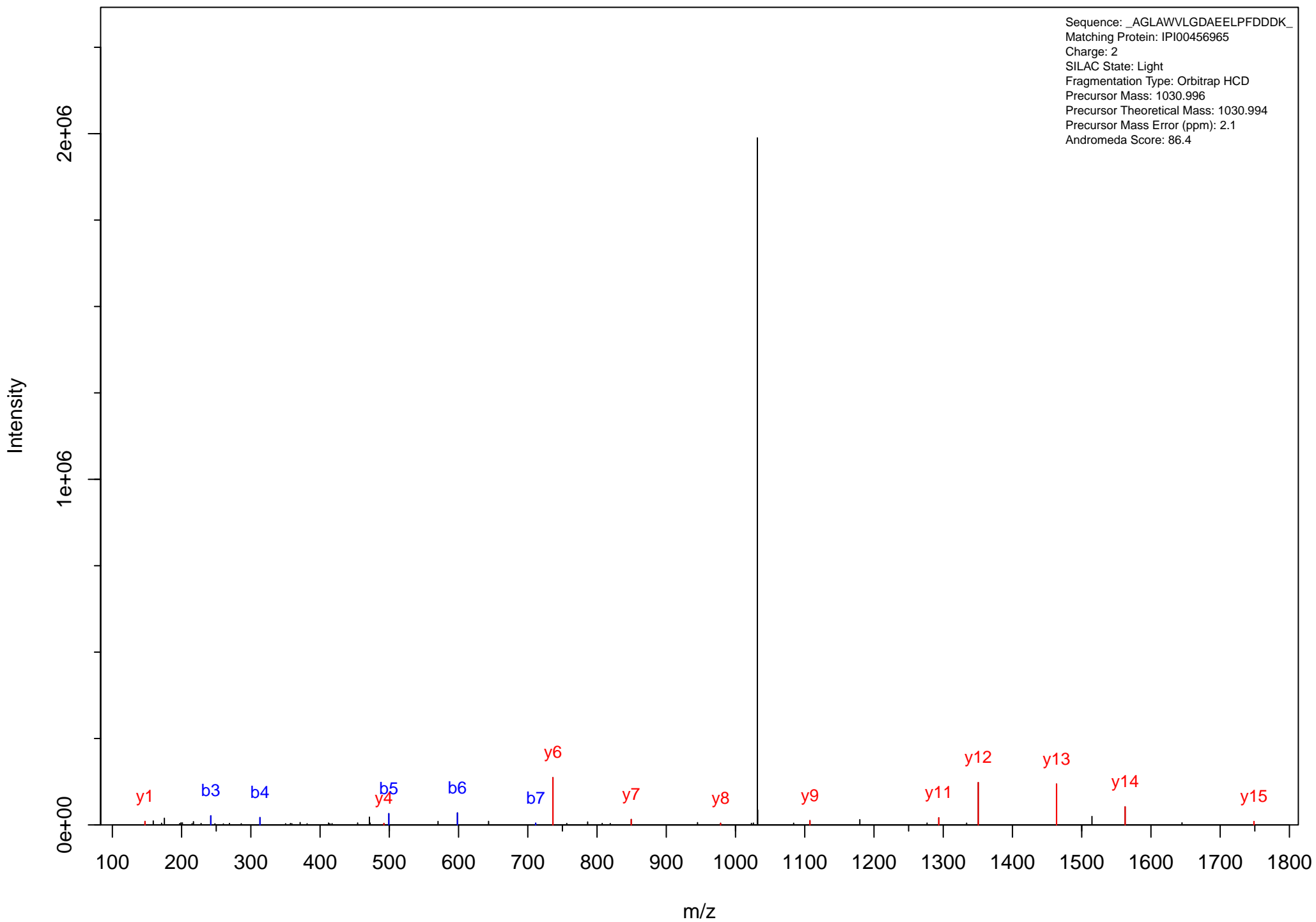


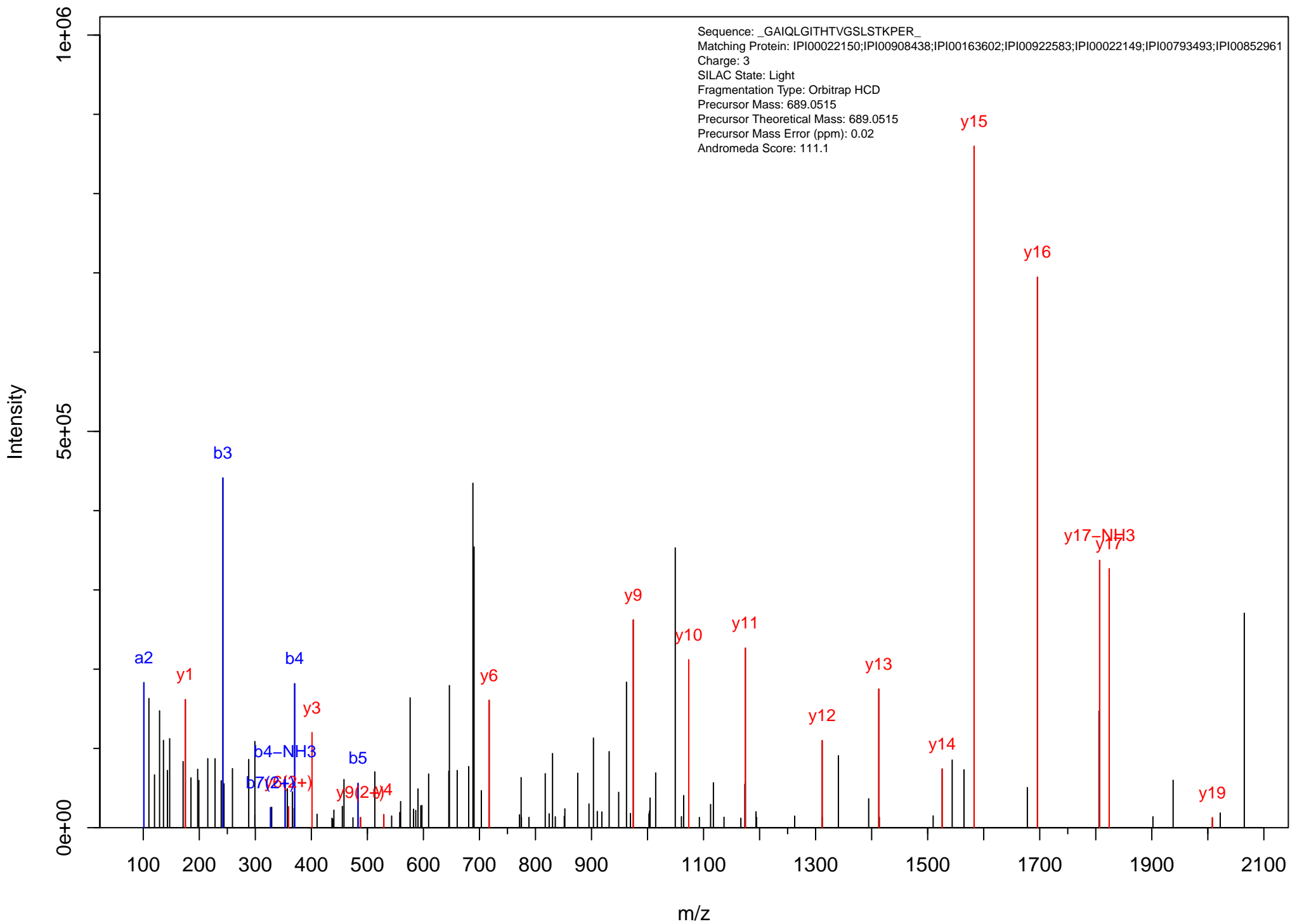


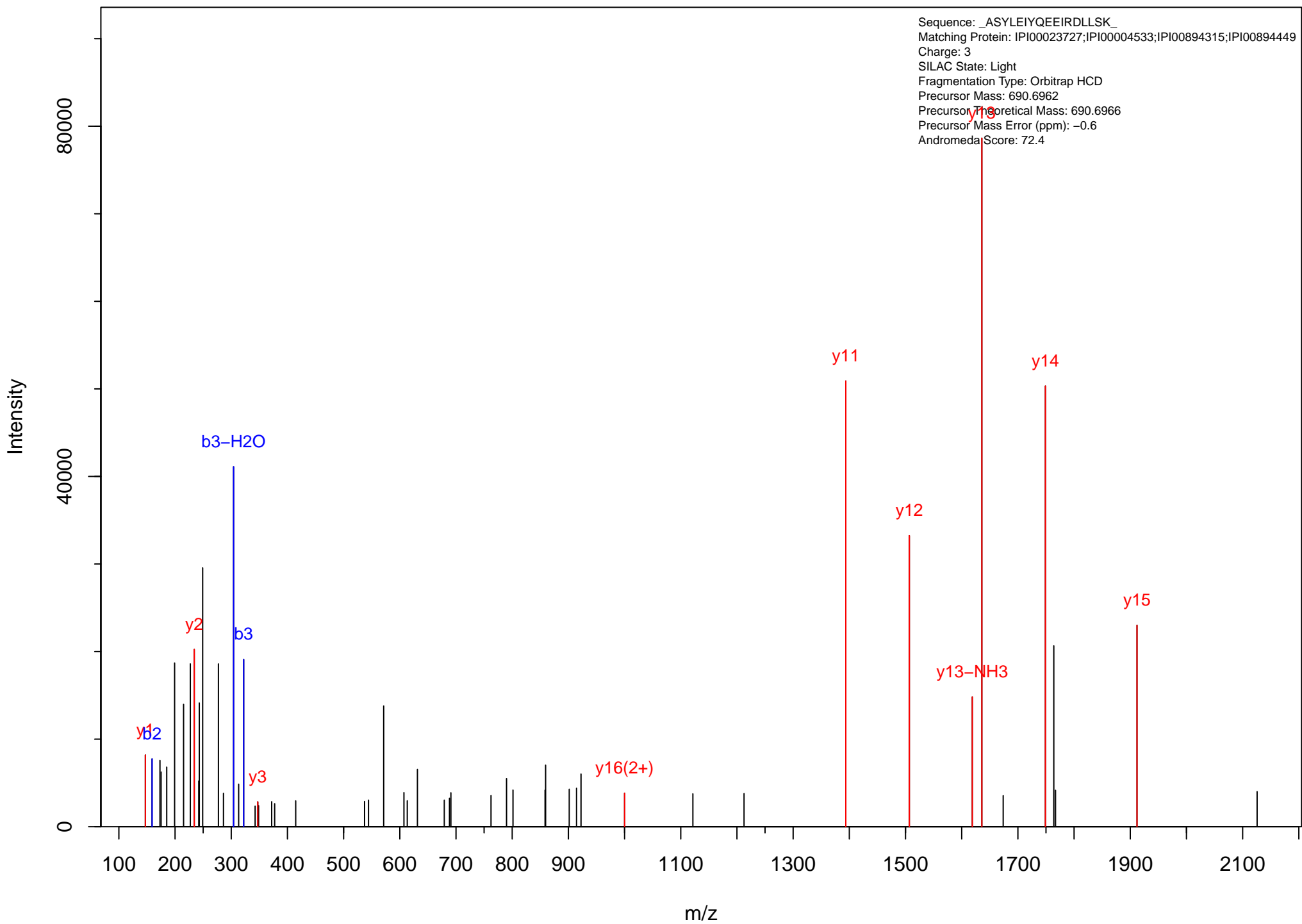


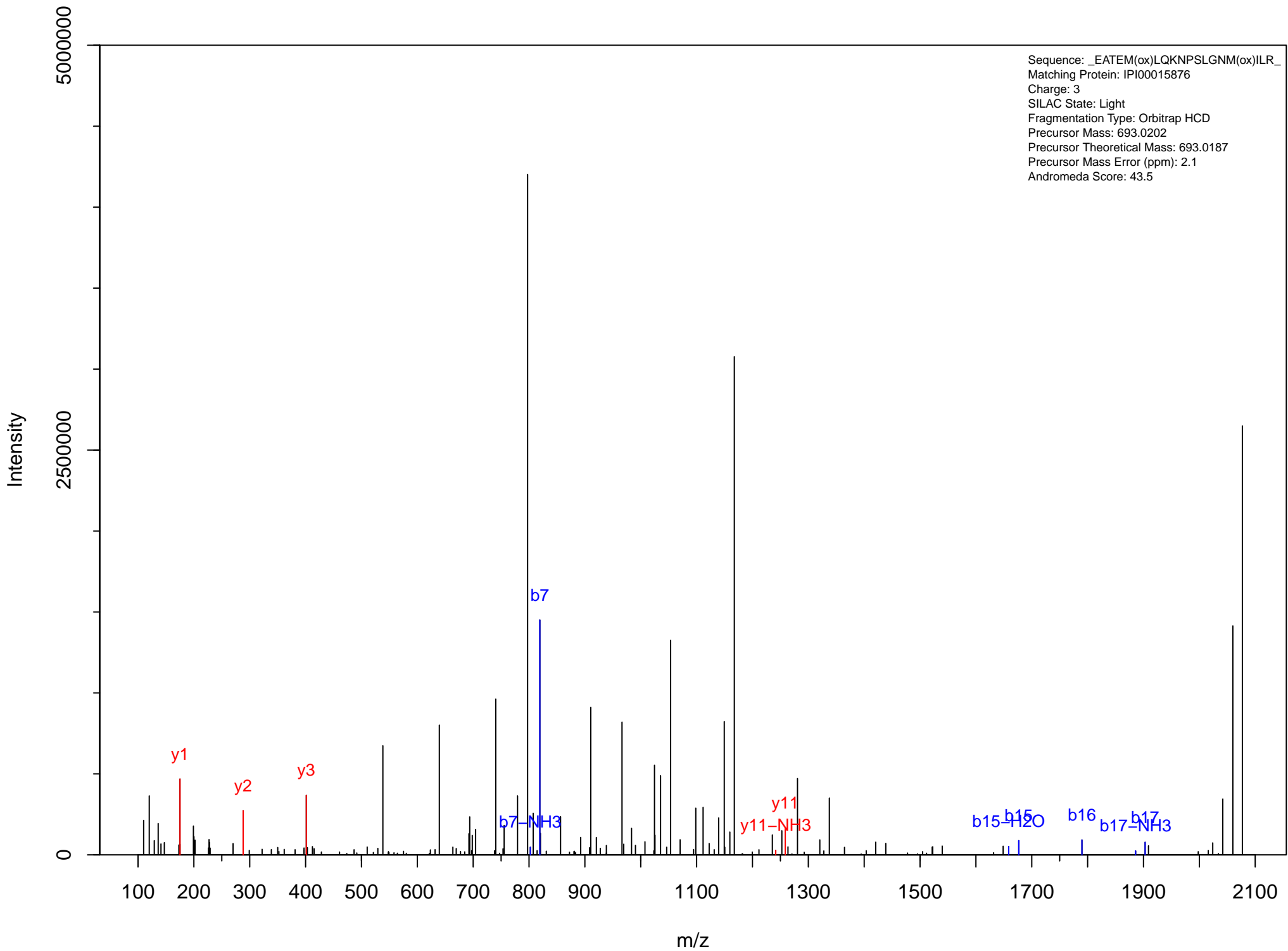


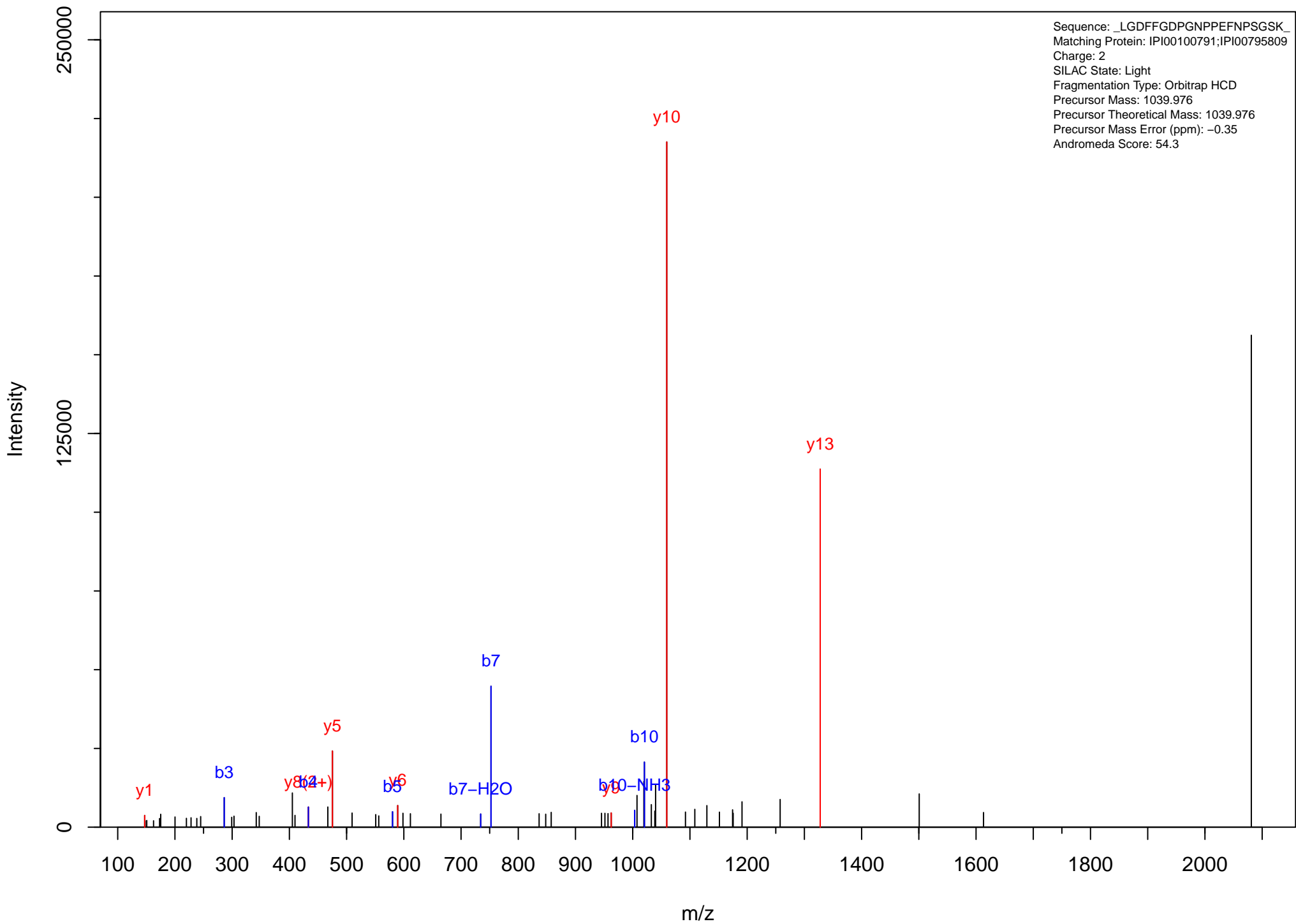


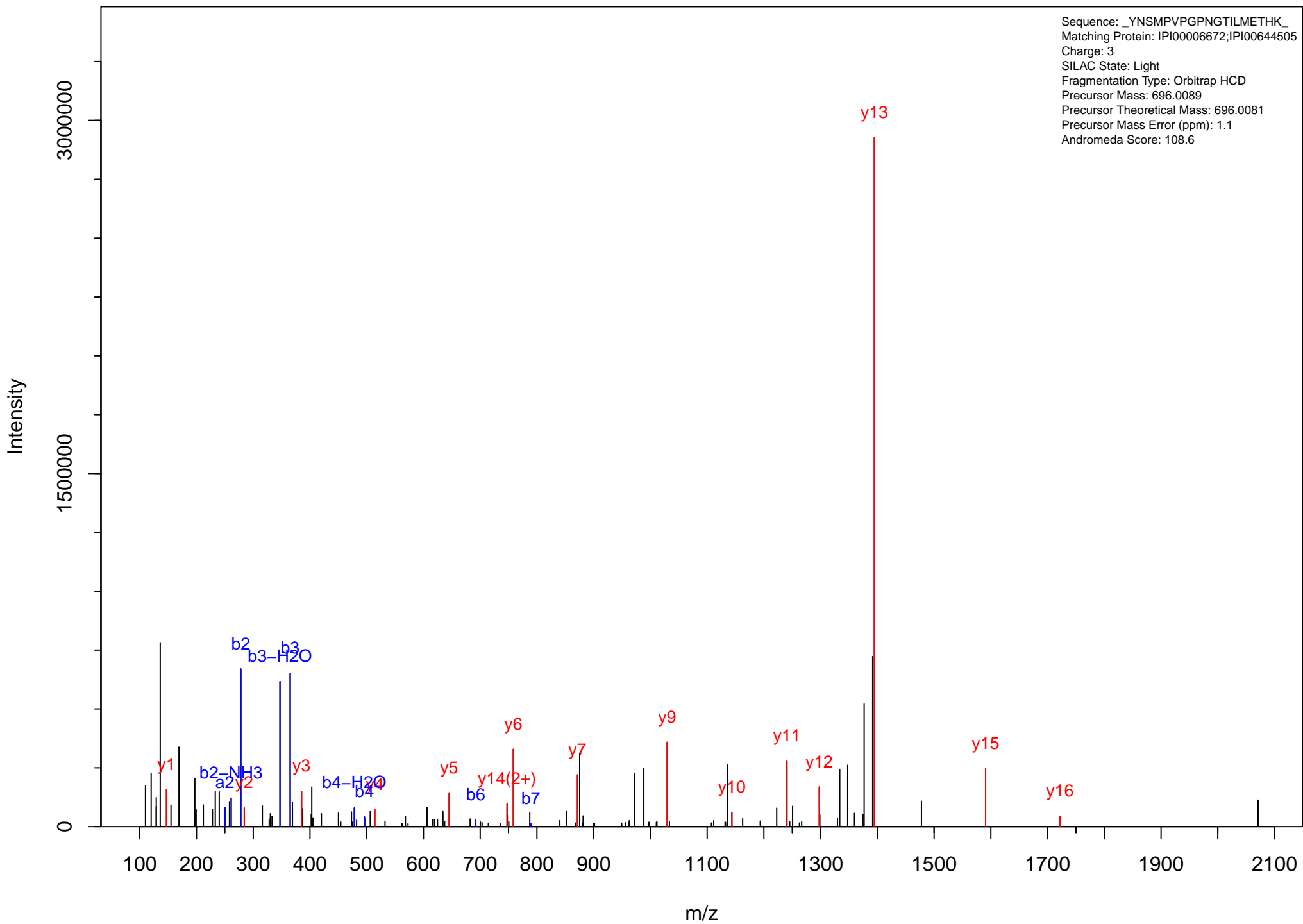


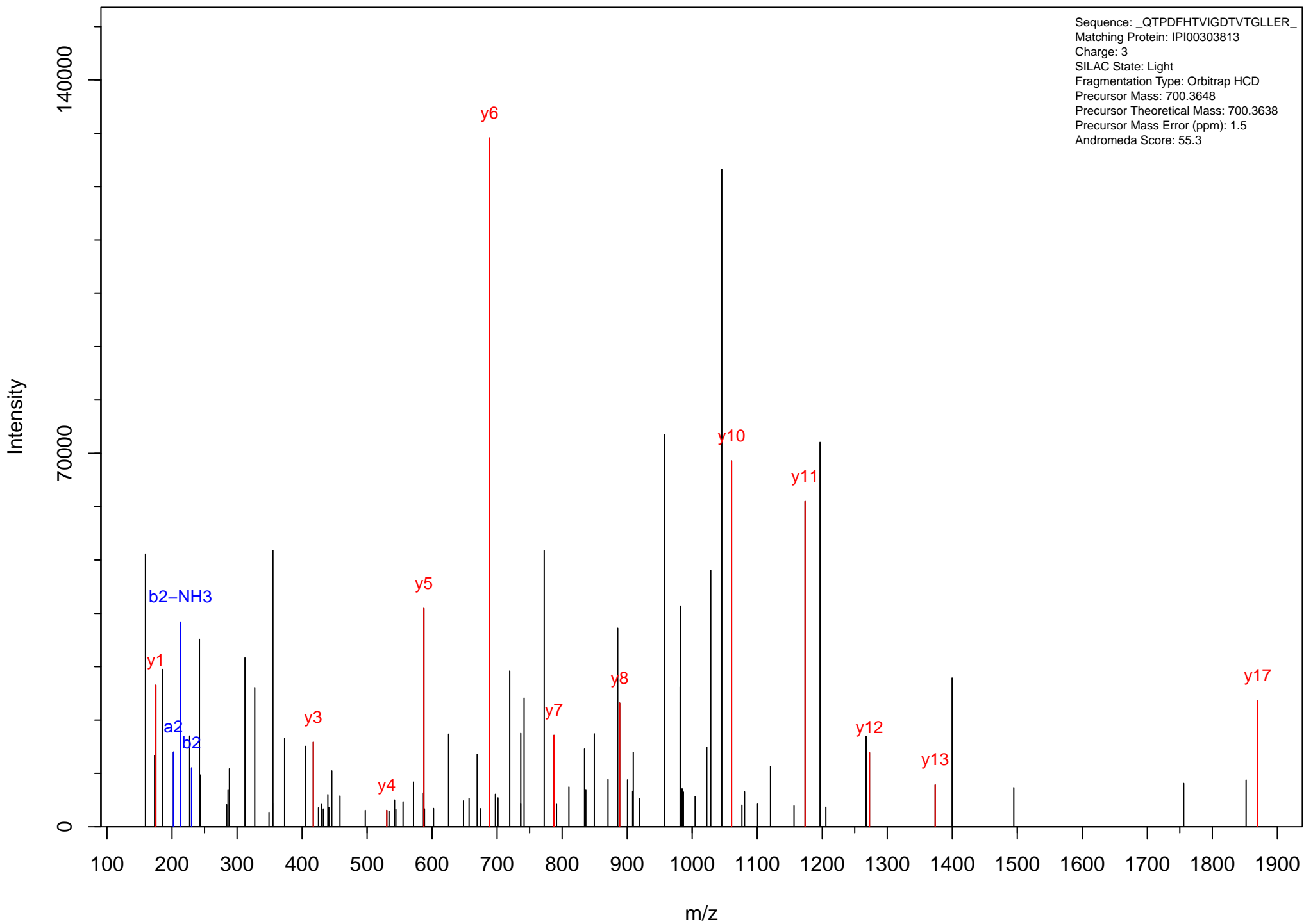


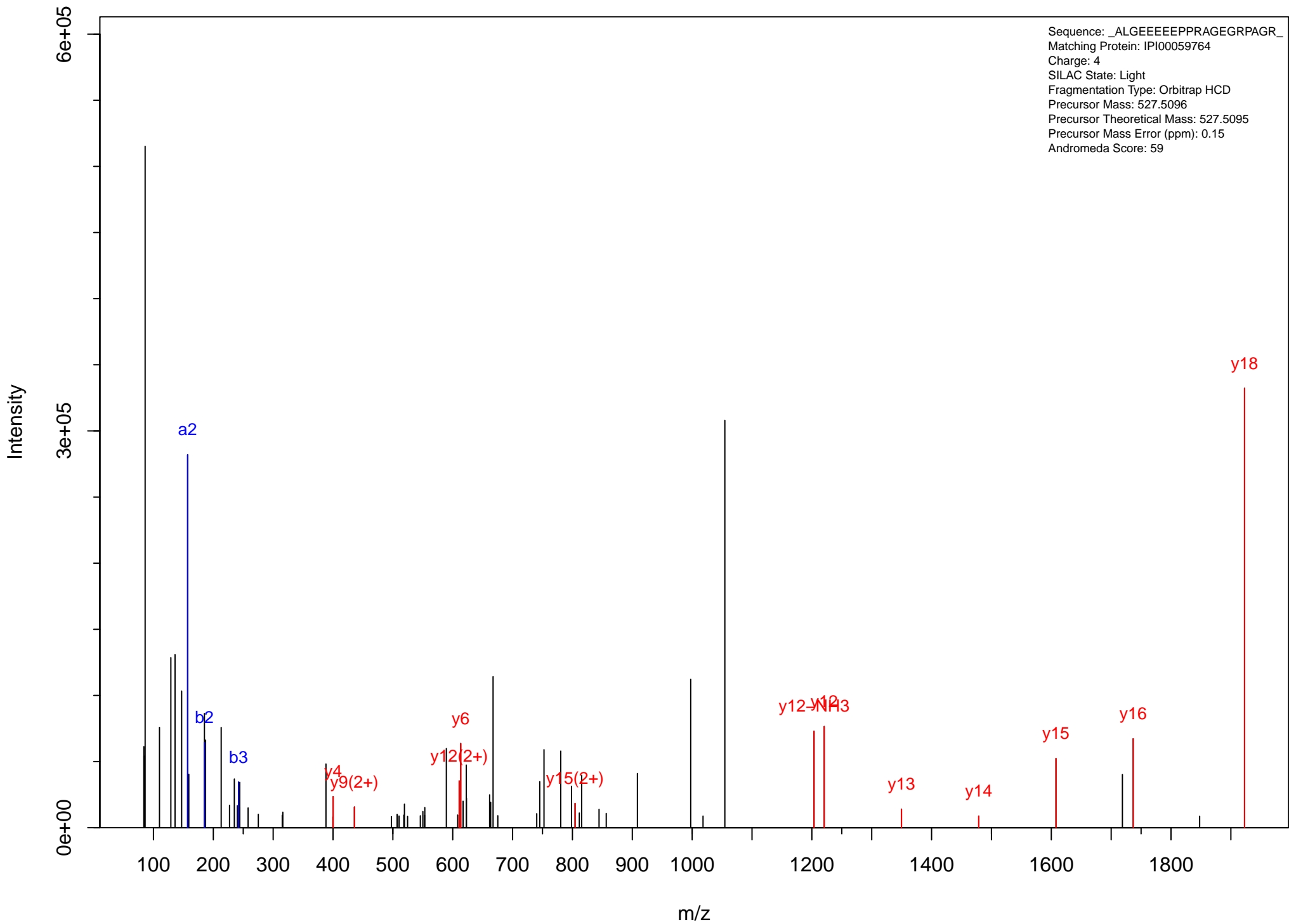


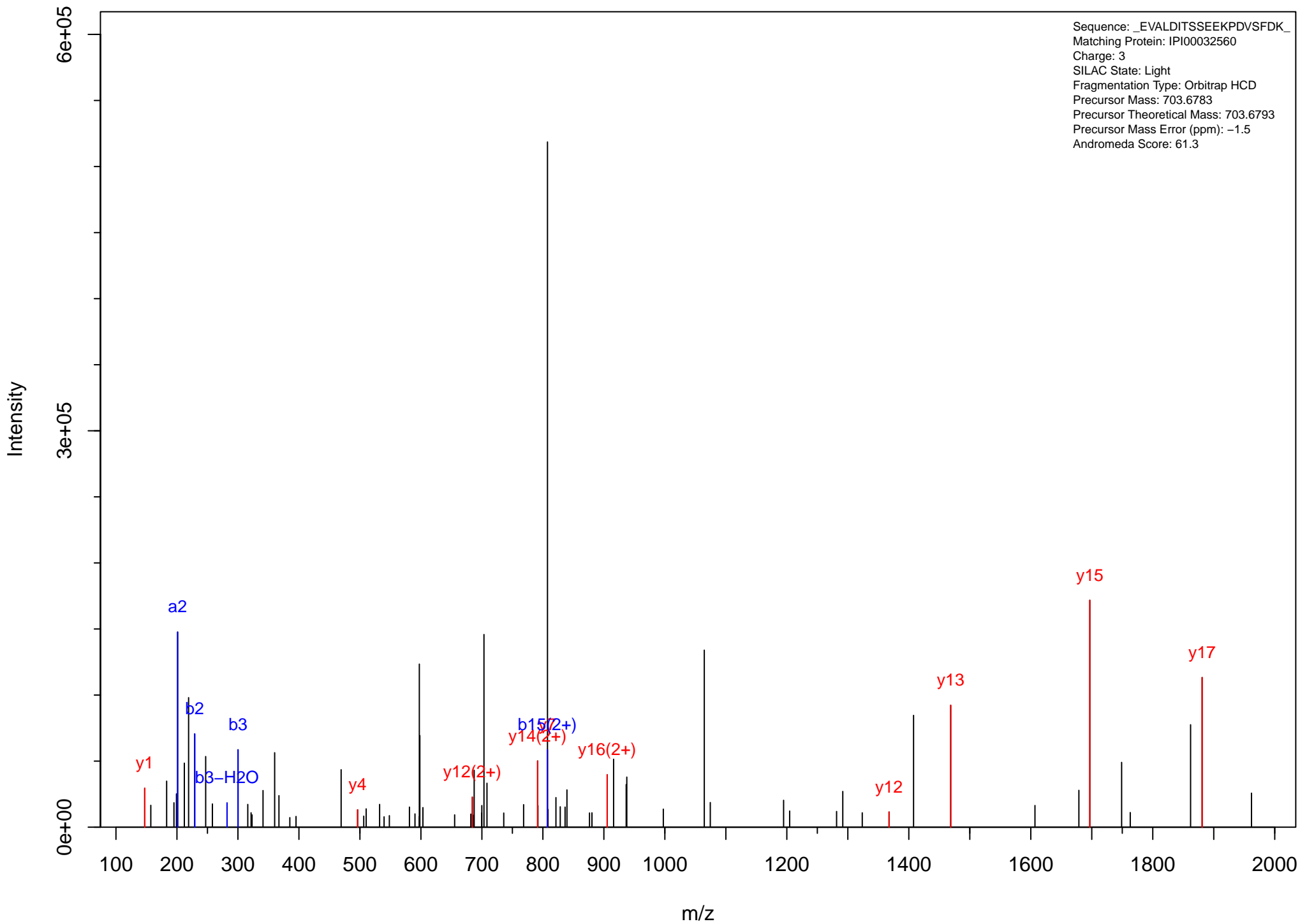


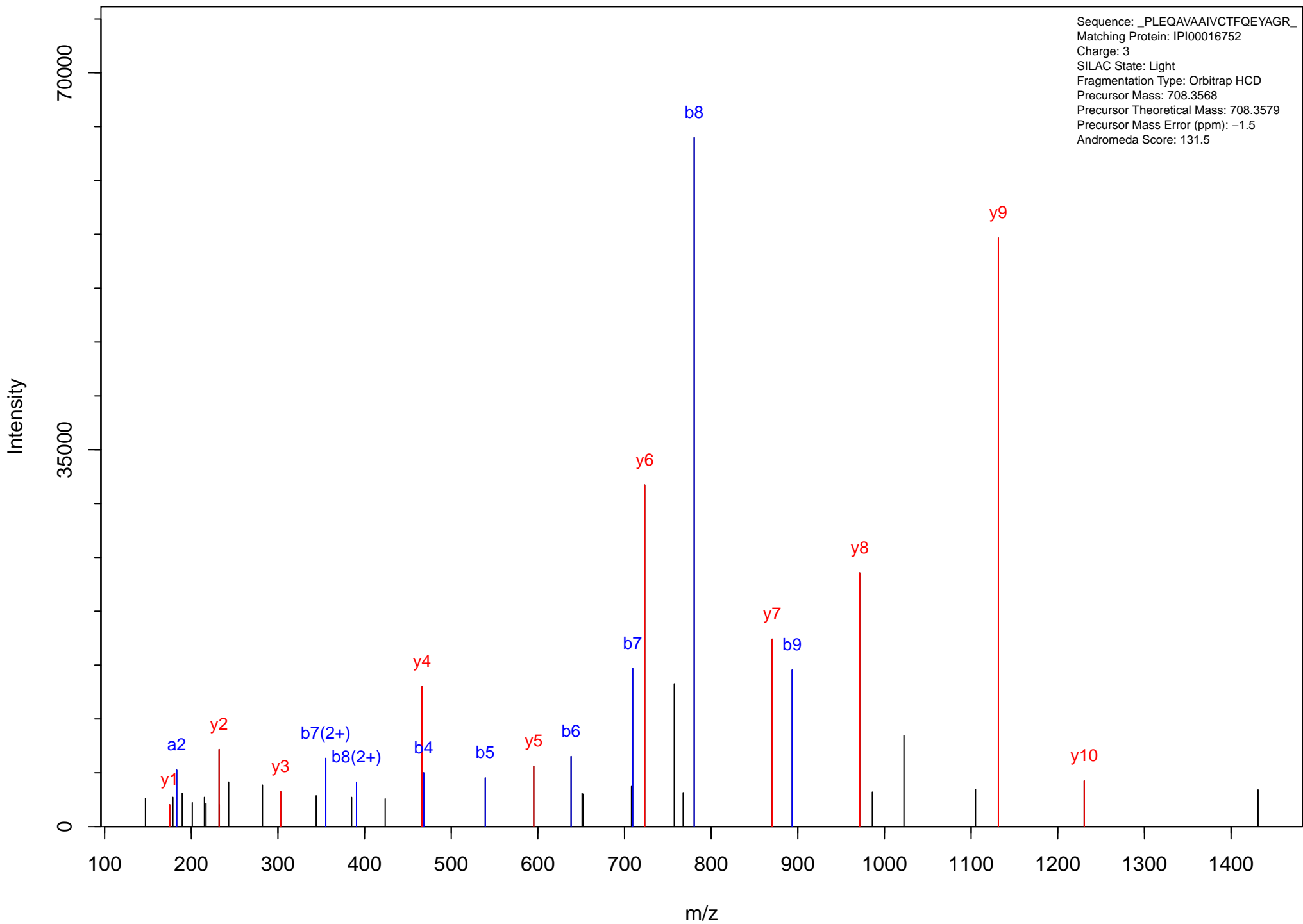


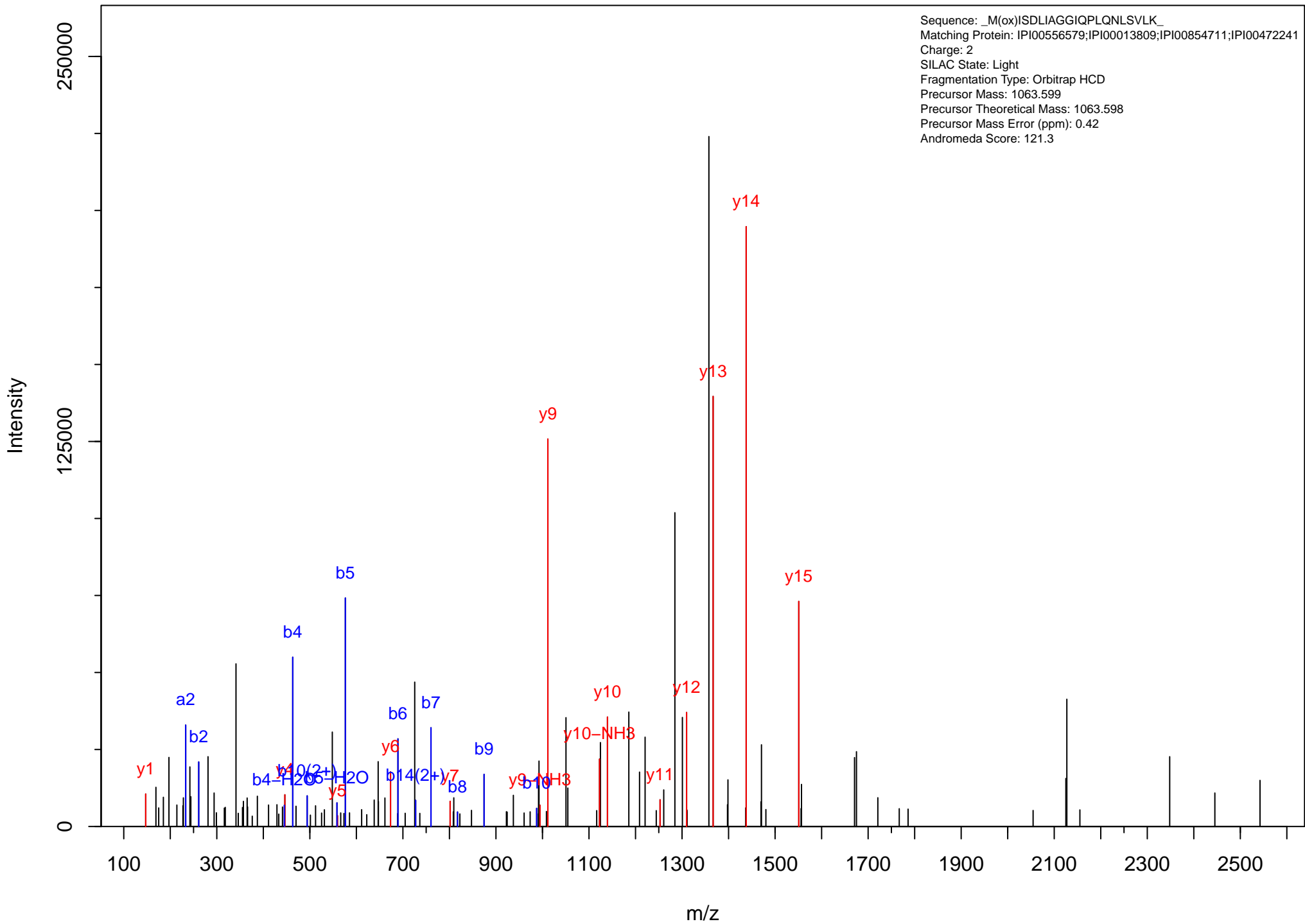


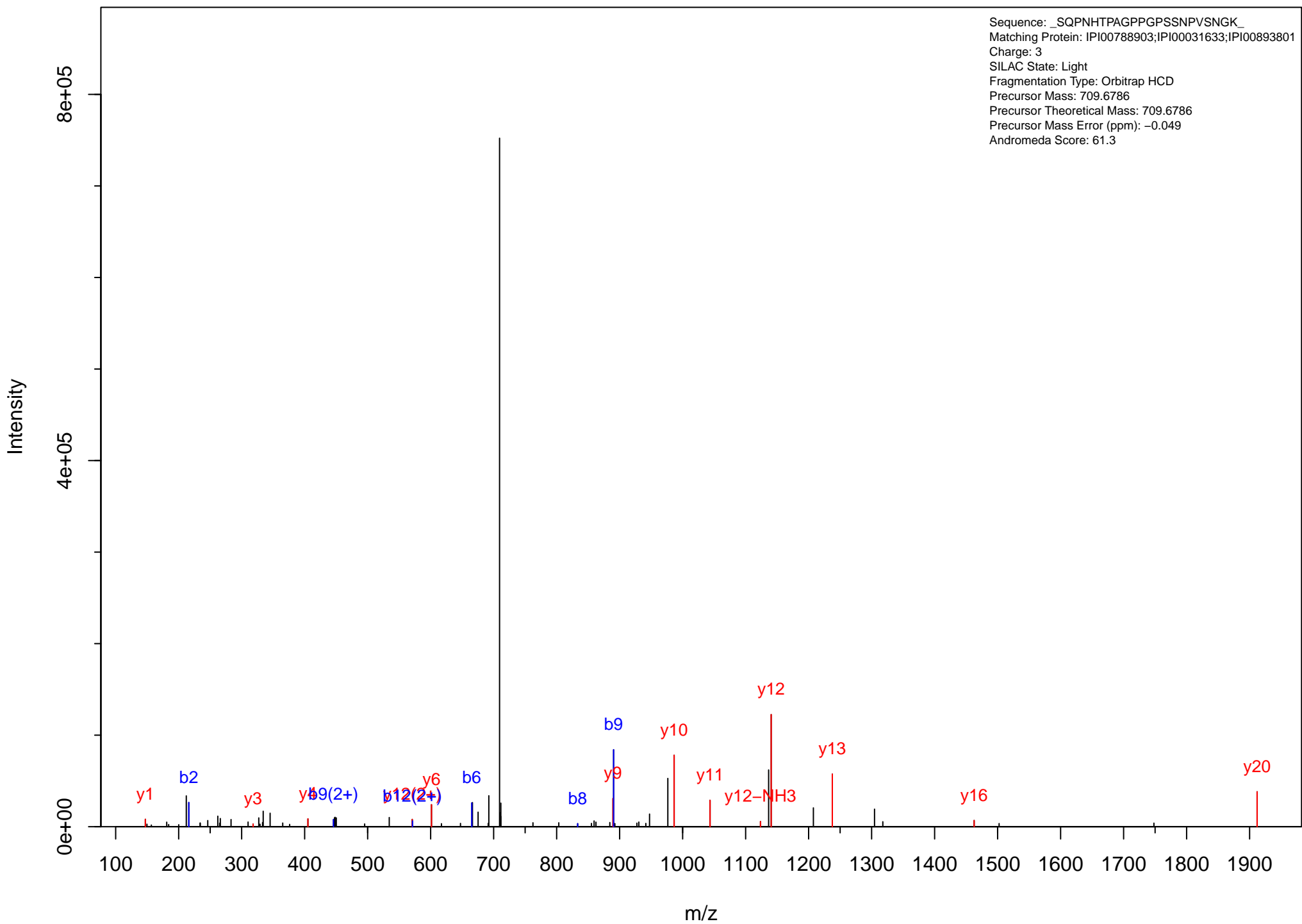


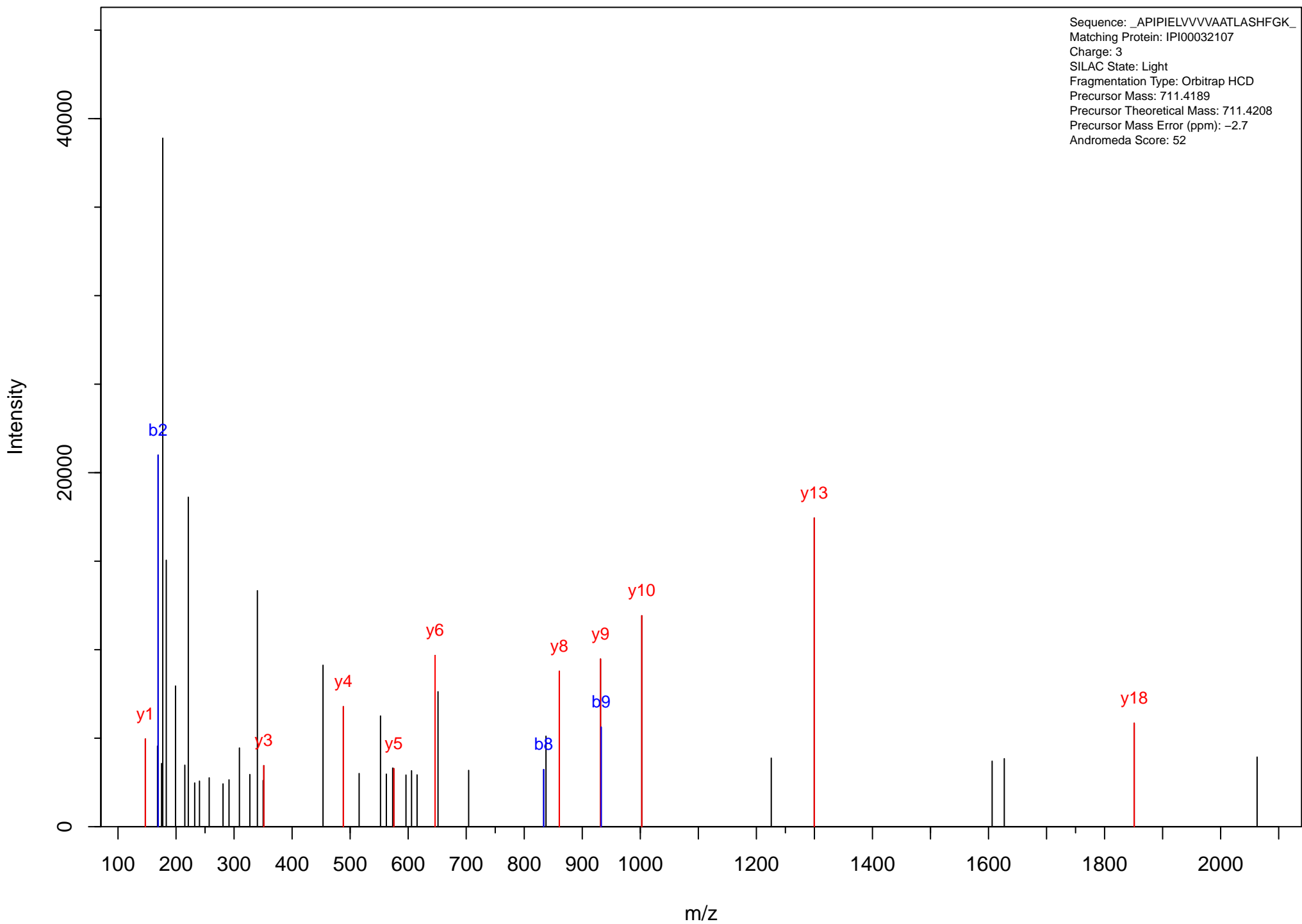


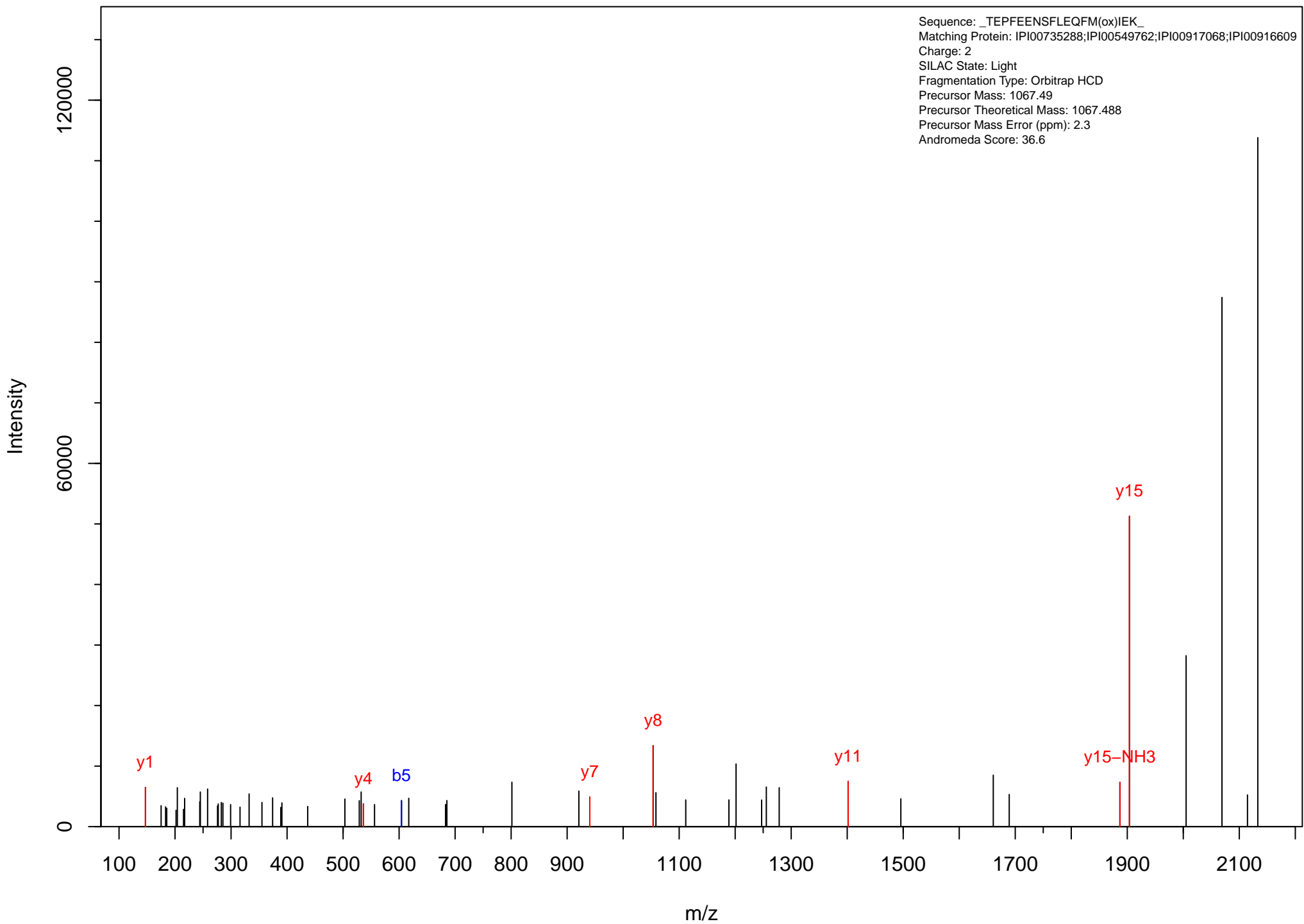


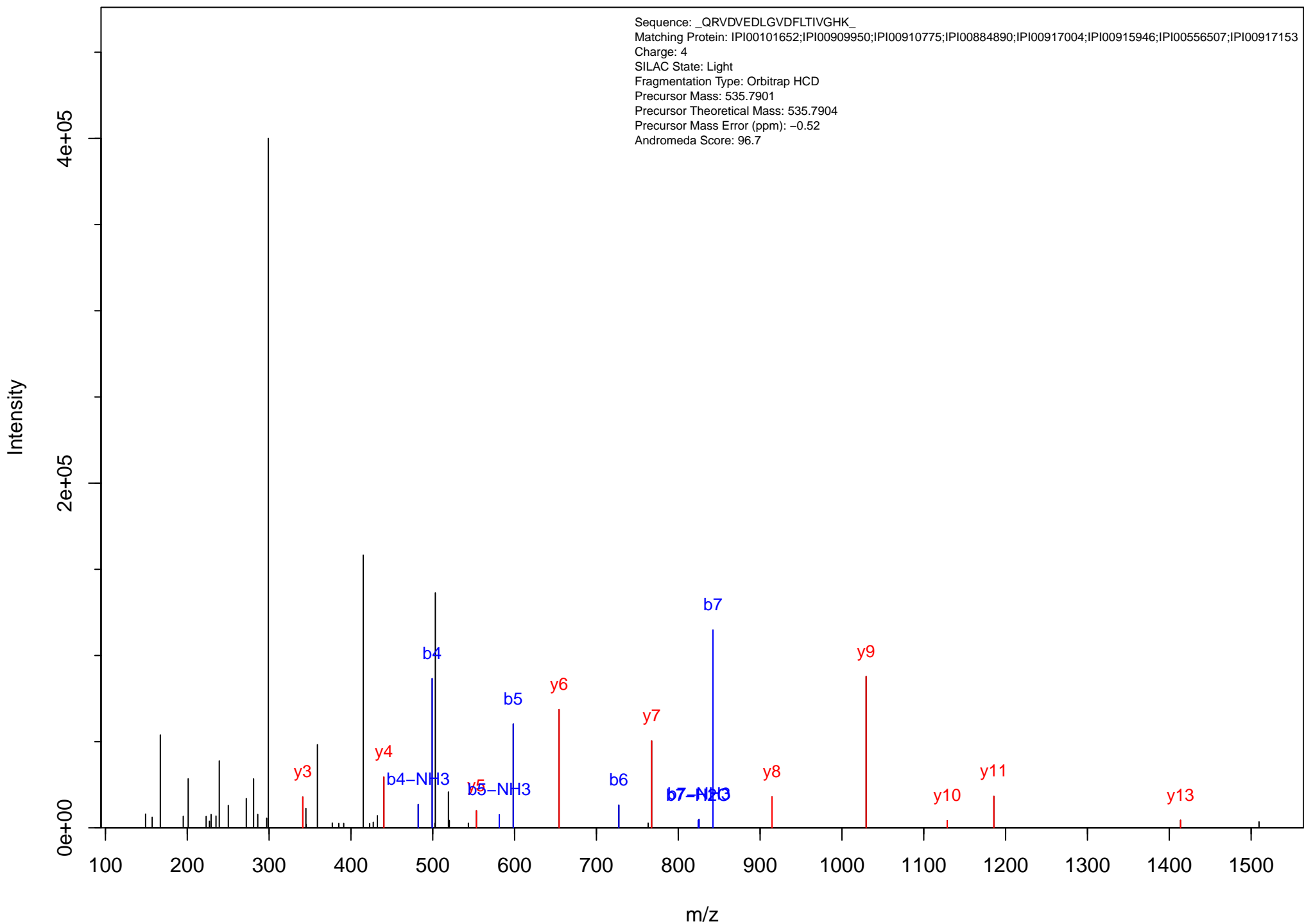


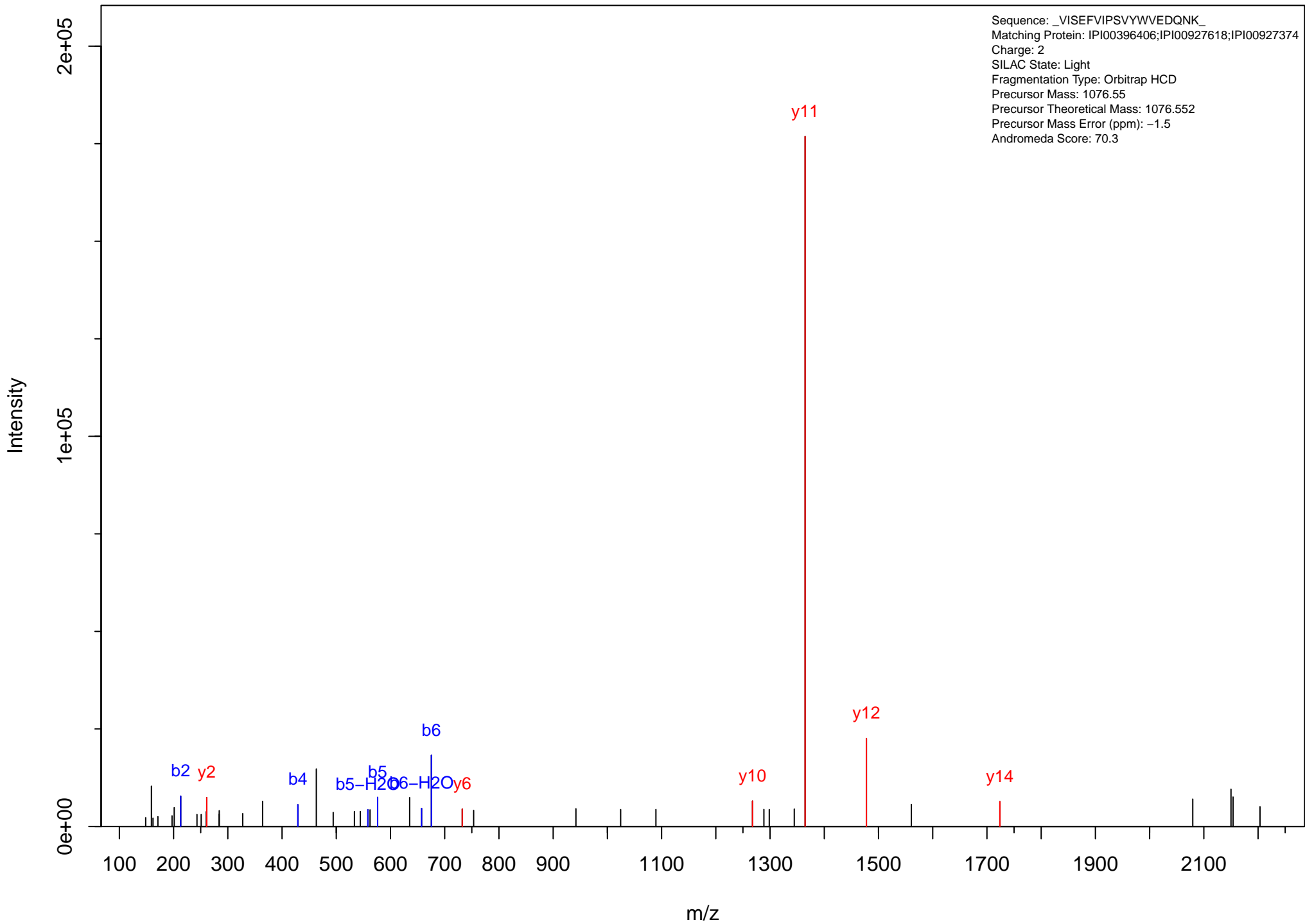


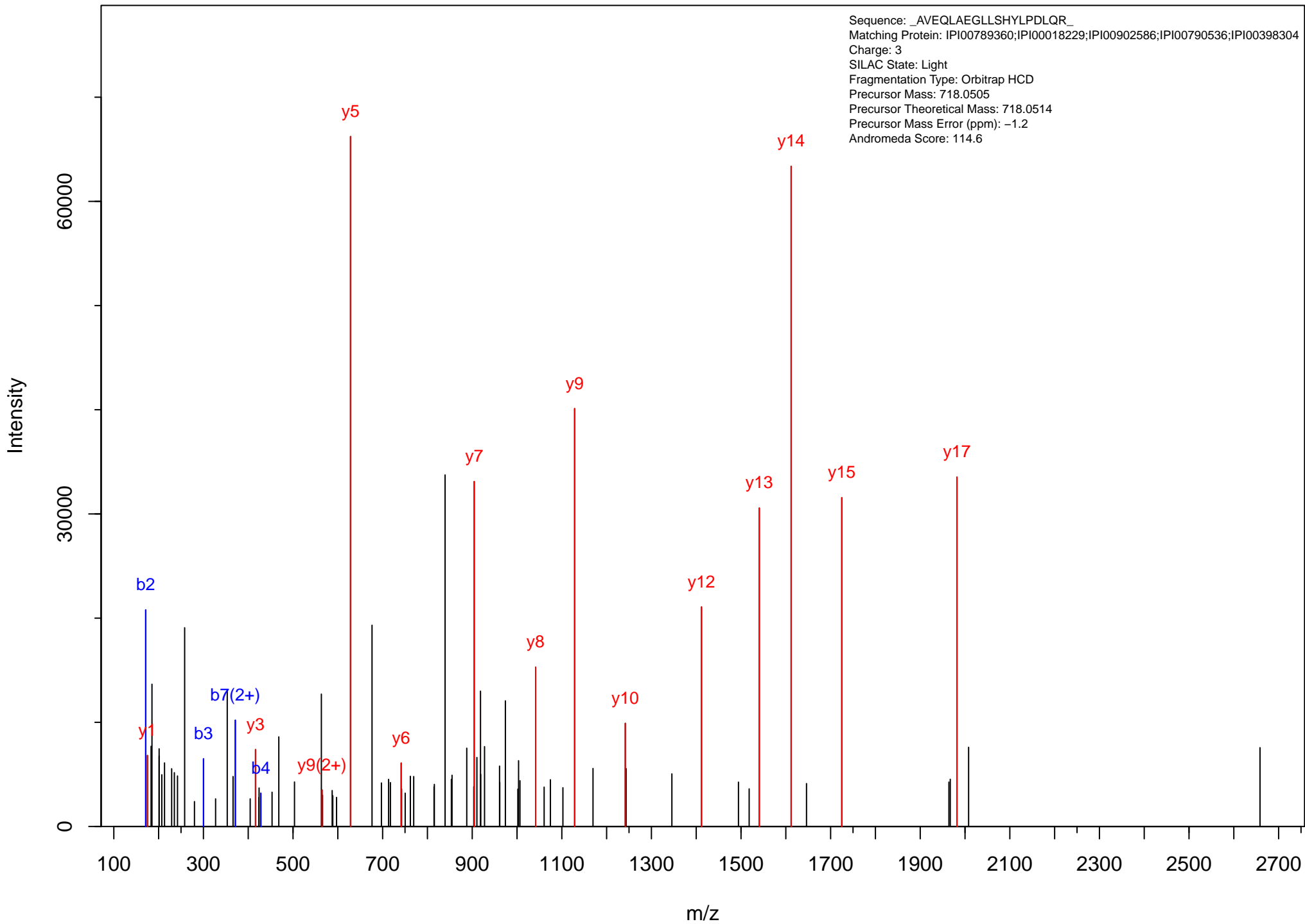


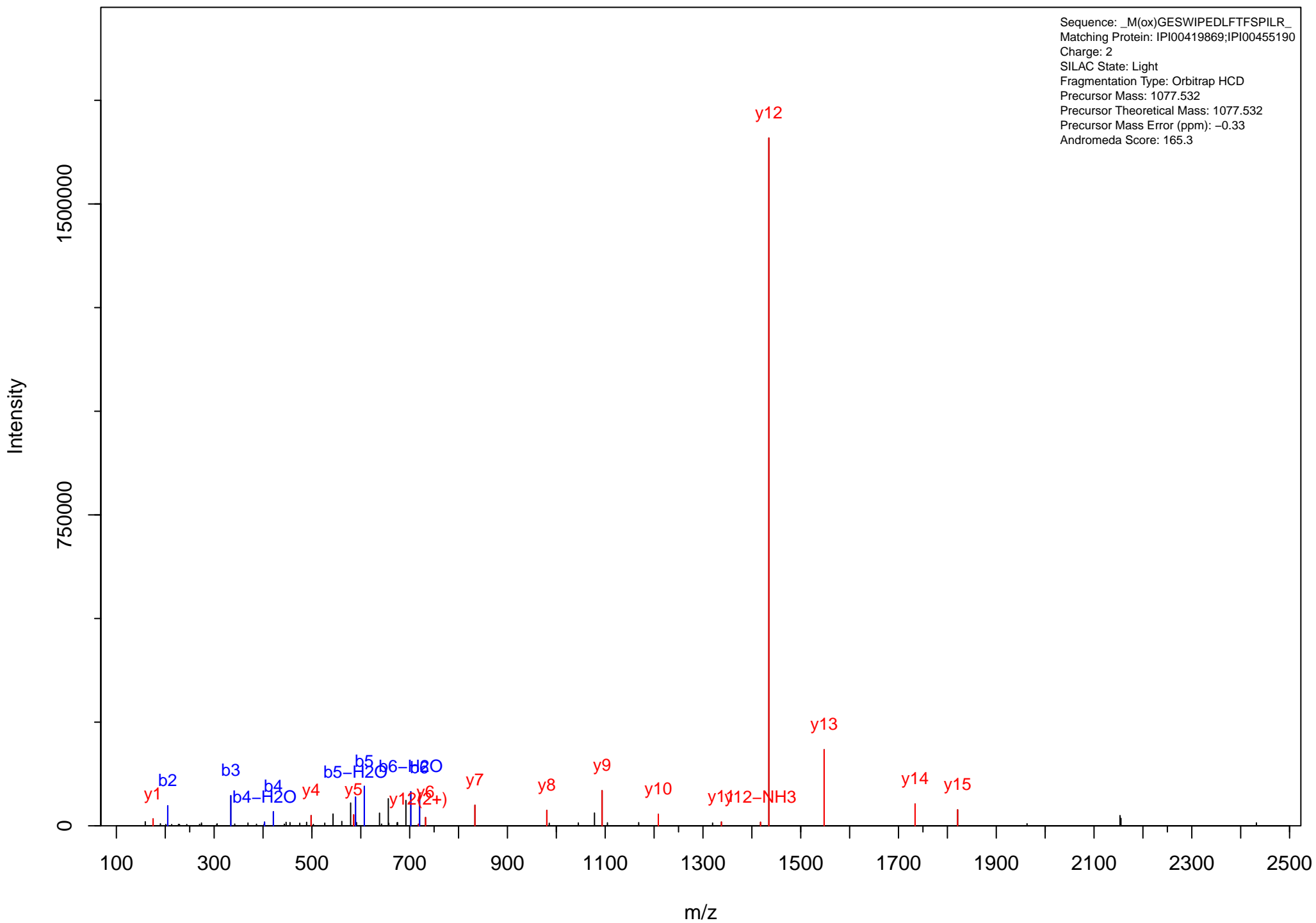


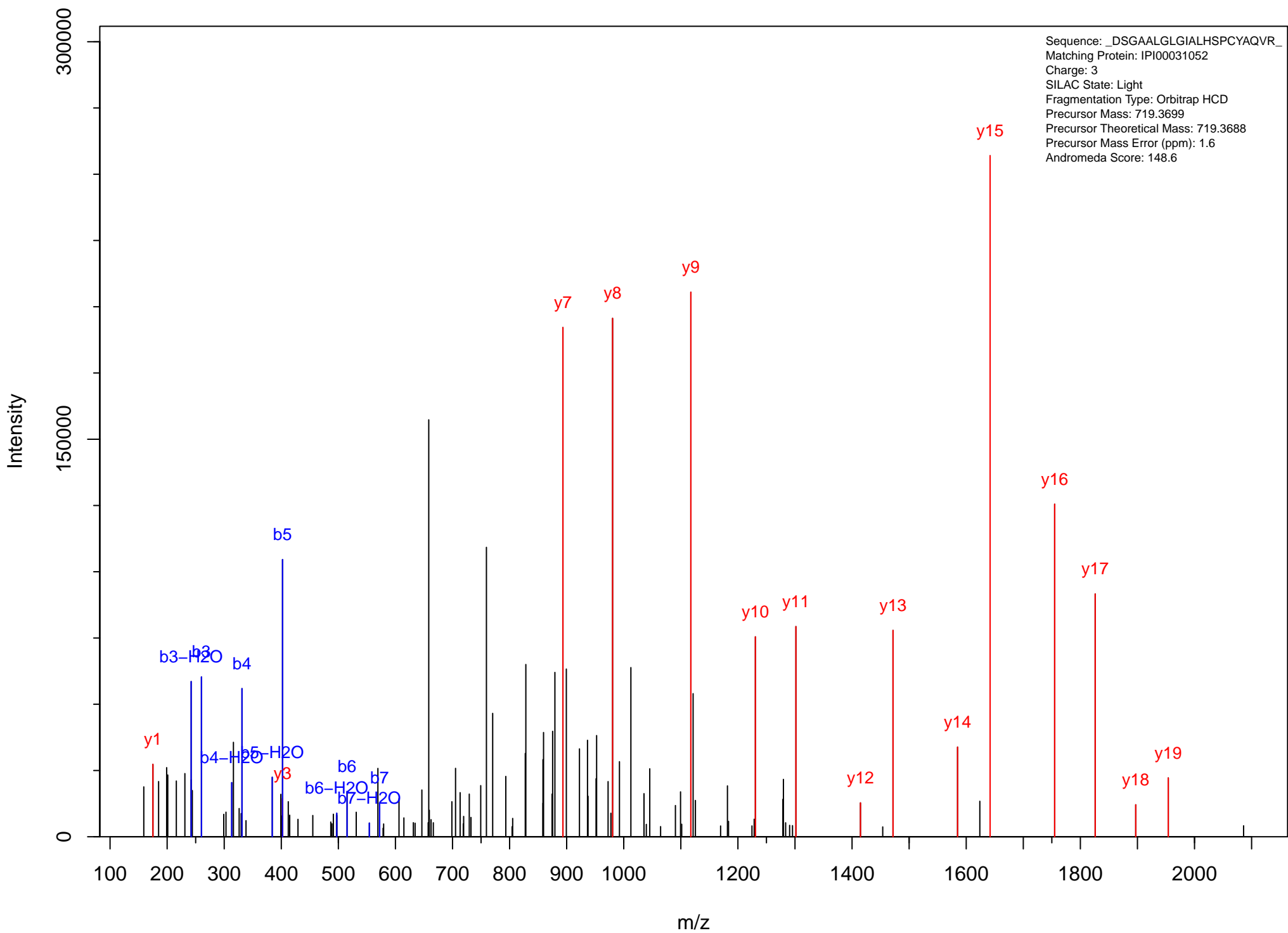












Intensity

6e+05

3e+05

0e+00

100 200 300 400 500 600 700 800 900 1100 1300 1500 1700 1900 2100

m/z

Sequence: _QLLAEESLPTTPFYFILGK_
Matching Protein: IPI00217240
Charge: 2
SILAC State: Light
Fragmentation Type: Orbitrap HCD
Precursor Mass: 1084.088
Precursor Theoretical Mass: 1084.088
Precursor Mass Error (ppm): -0.16
Andromeda Score: 150.3

y11

y8

y13

y12

y14

y15

y16

a2

y2

b2

b3

y3

b3-NH3

b4

y4

b4-NH3

b5

y5

b5-NH3

b6

y6

b6-NH3

b7

y7

b7-H2O

b8

y8

b8-H2O

b9

y9

b9-H2O

b10

y10

b10-H2O

b11

y11

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b13

y13

b13-H2O

b14

y14

b14-H2O

b15

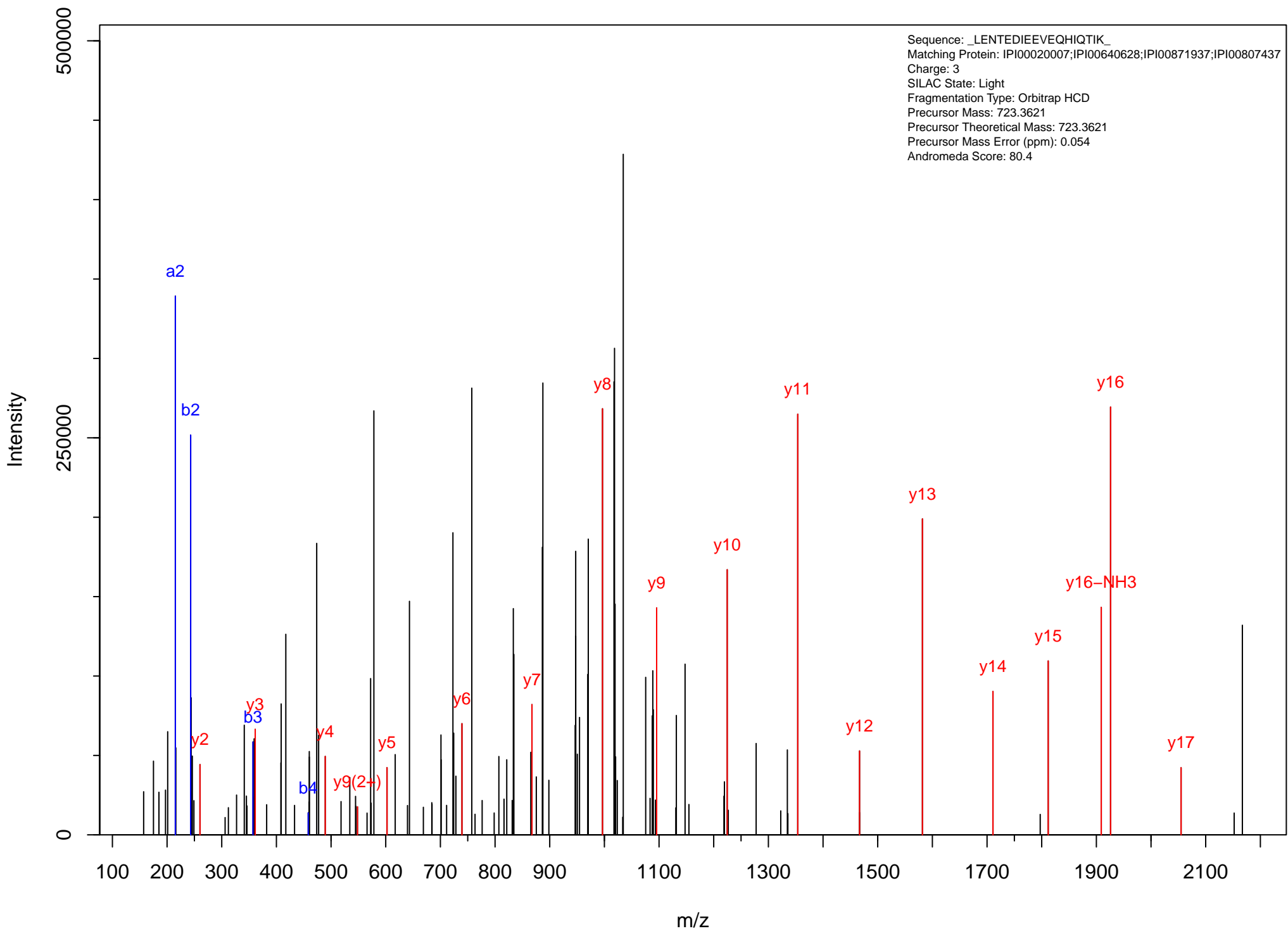
y15

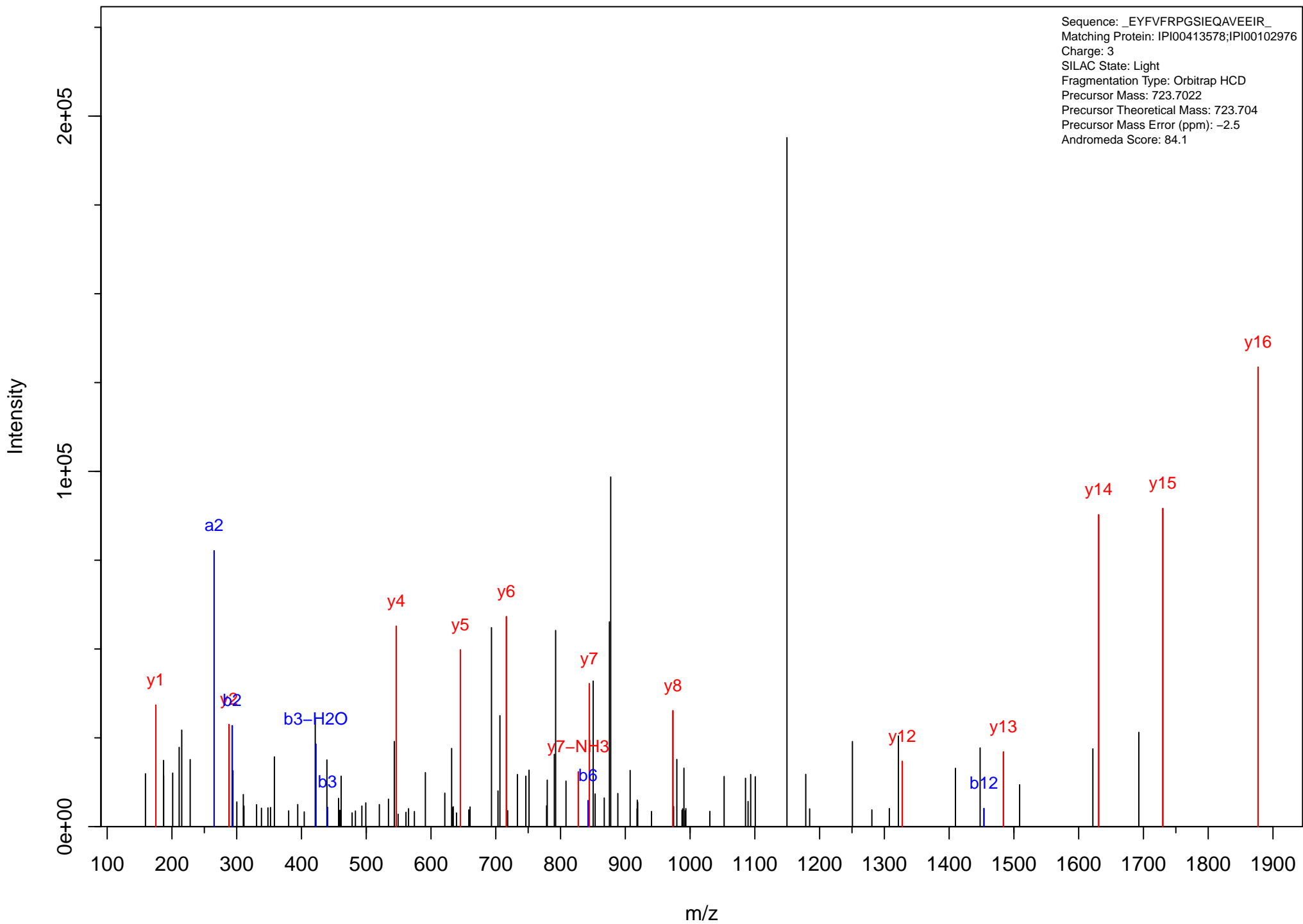
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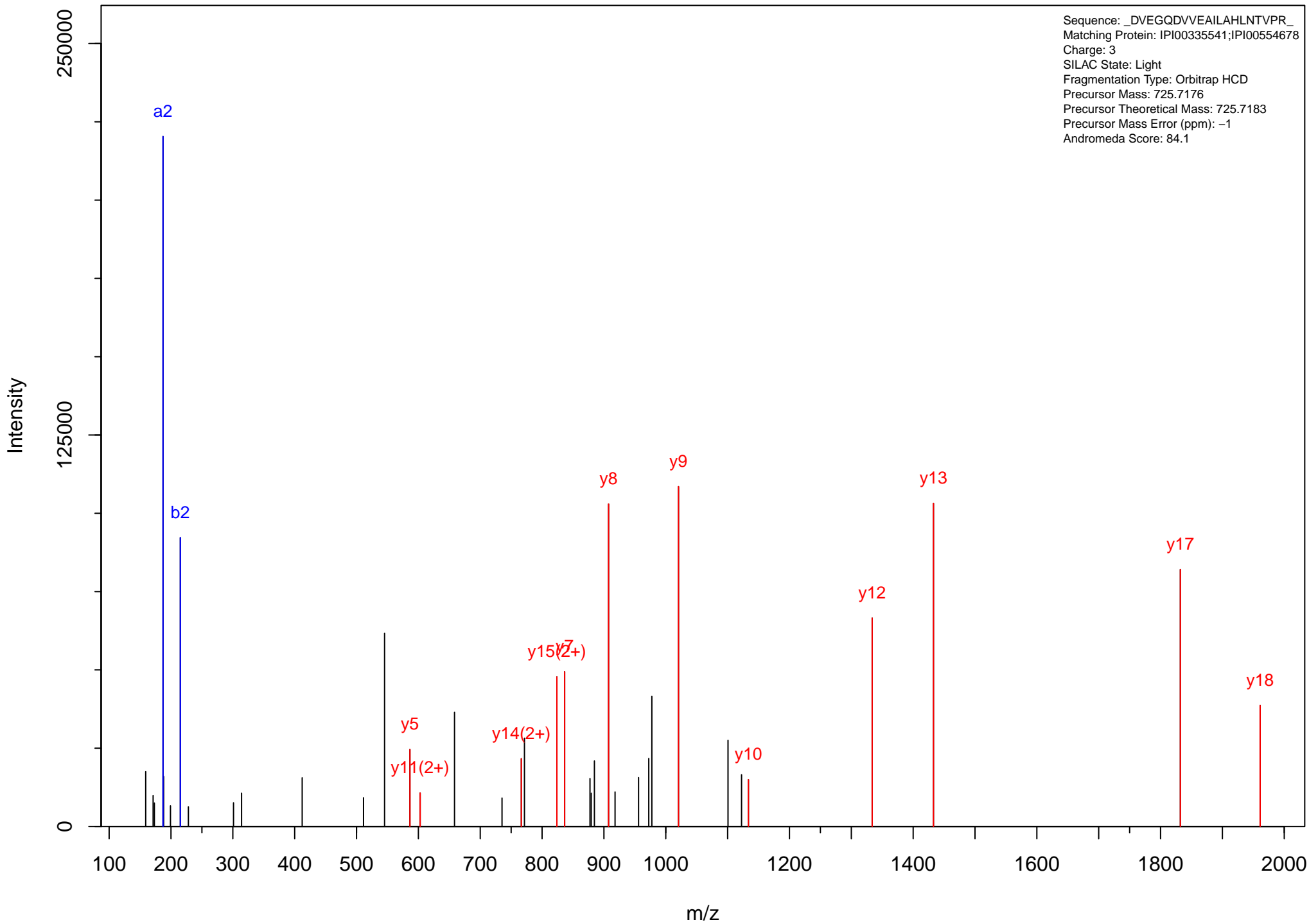
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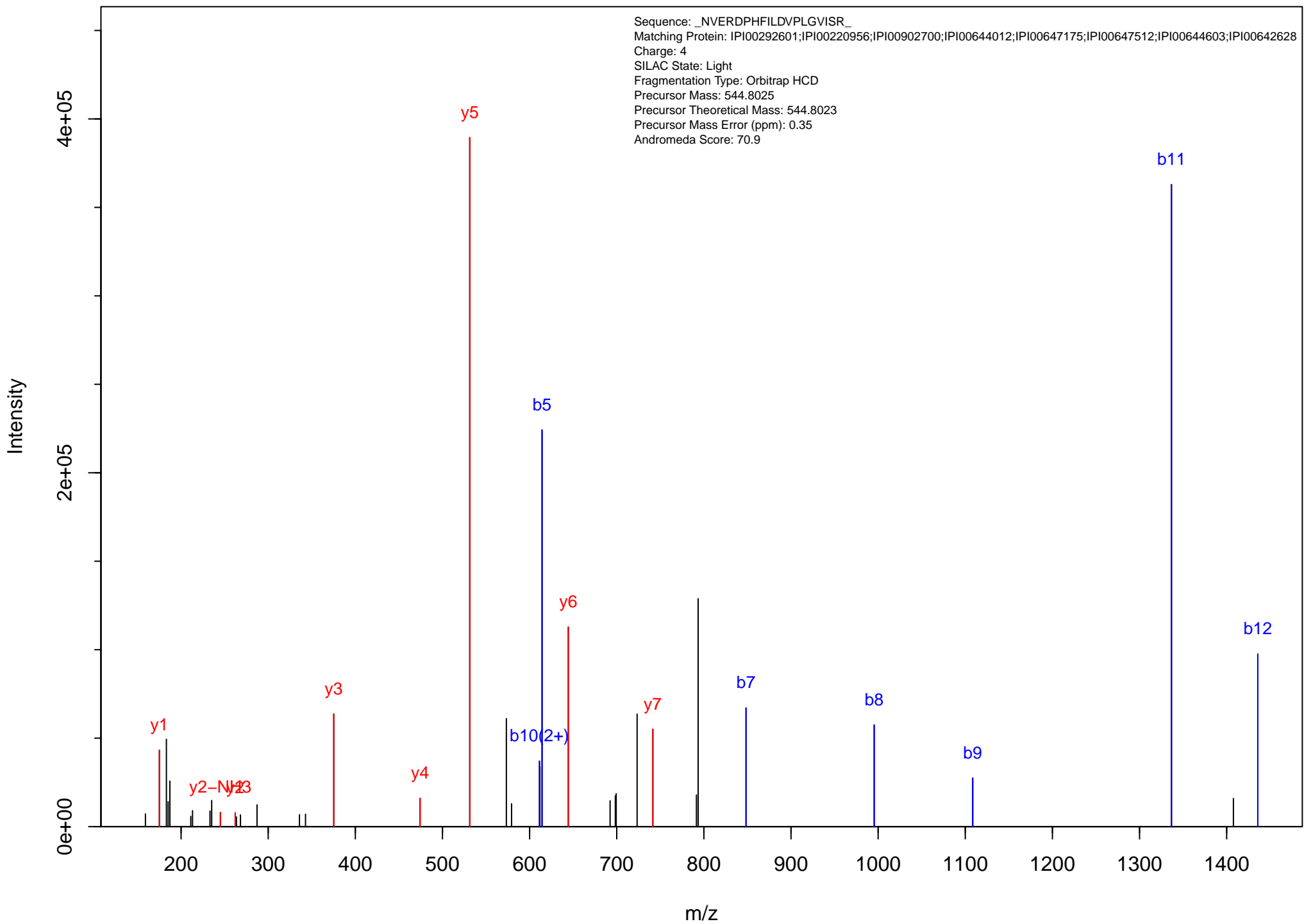
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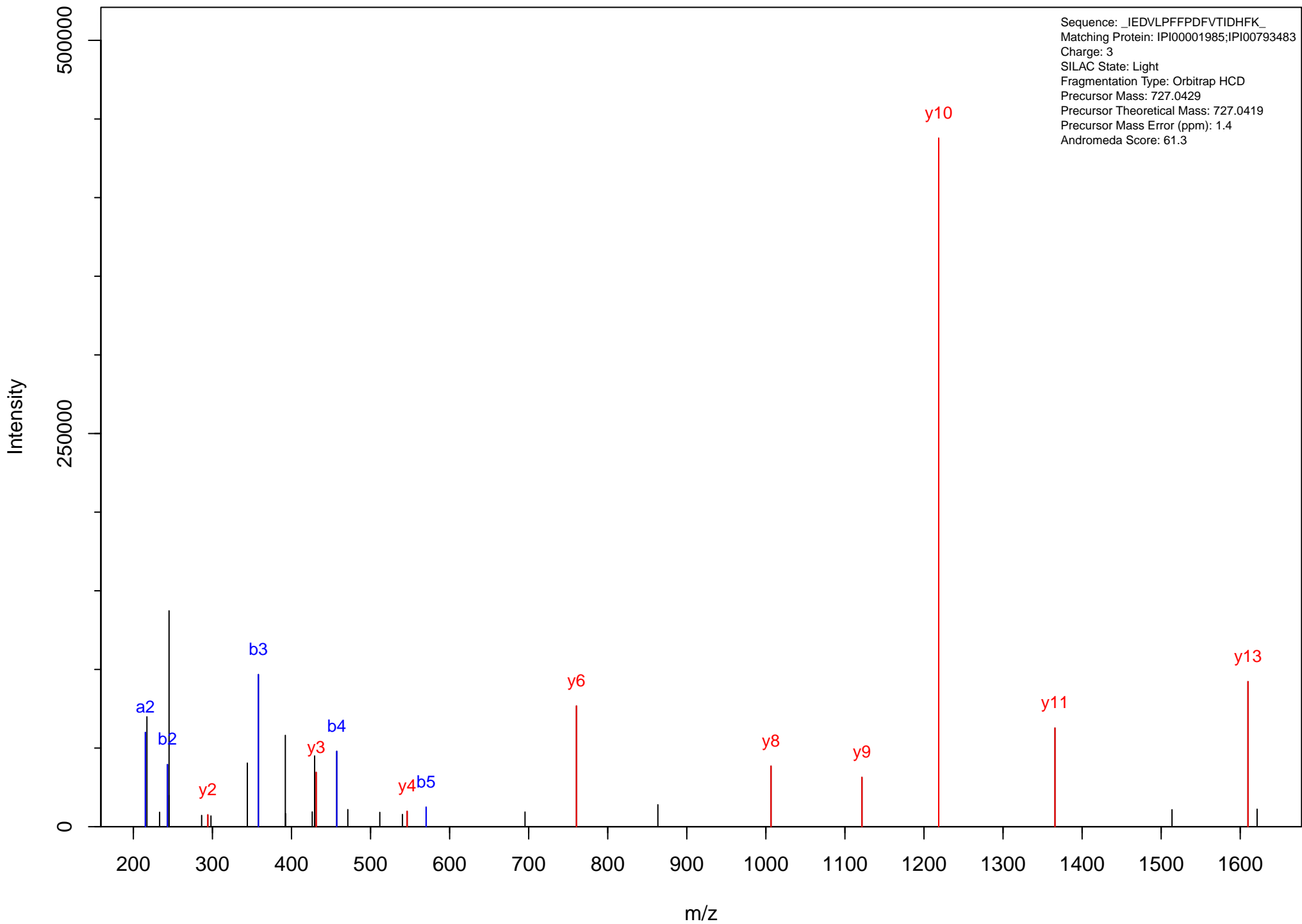
b16-H2O

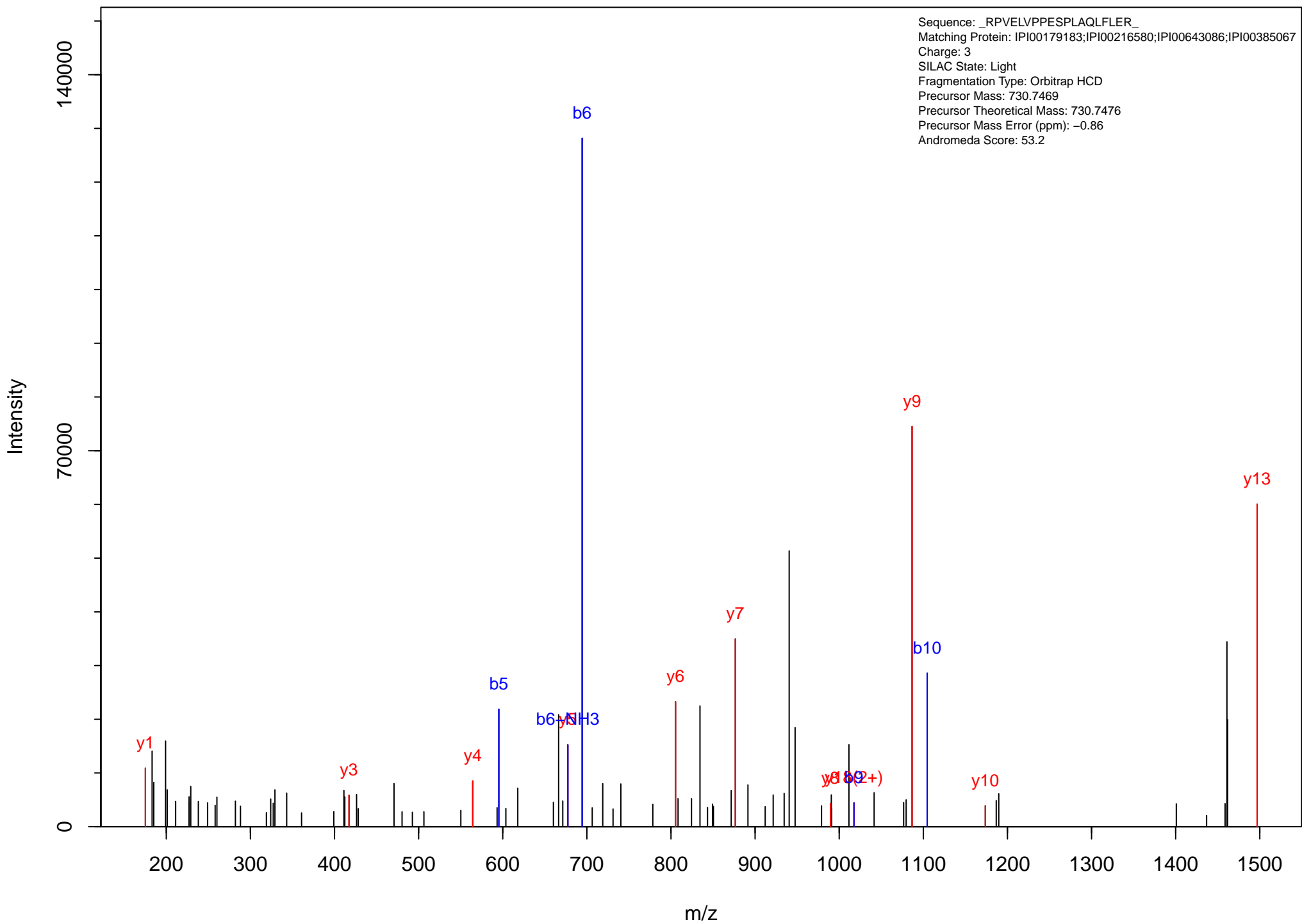


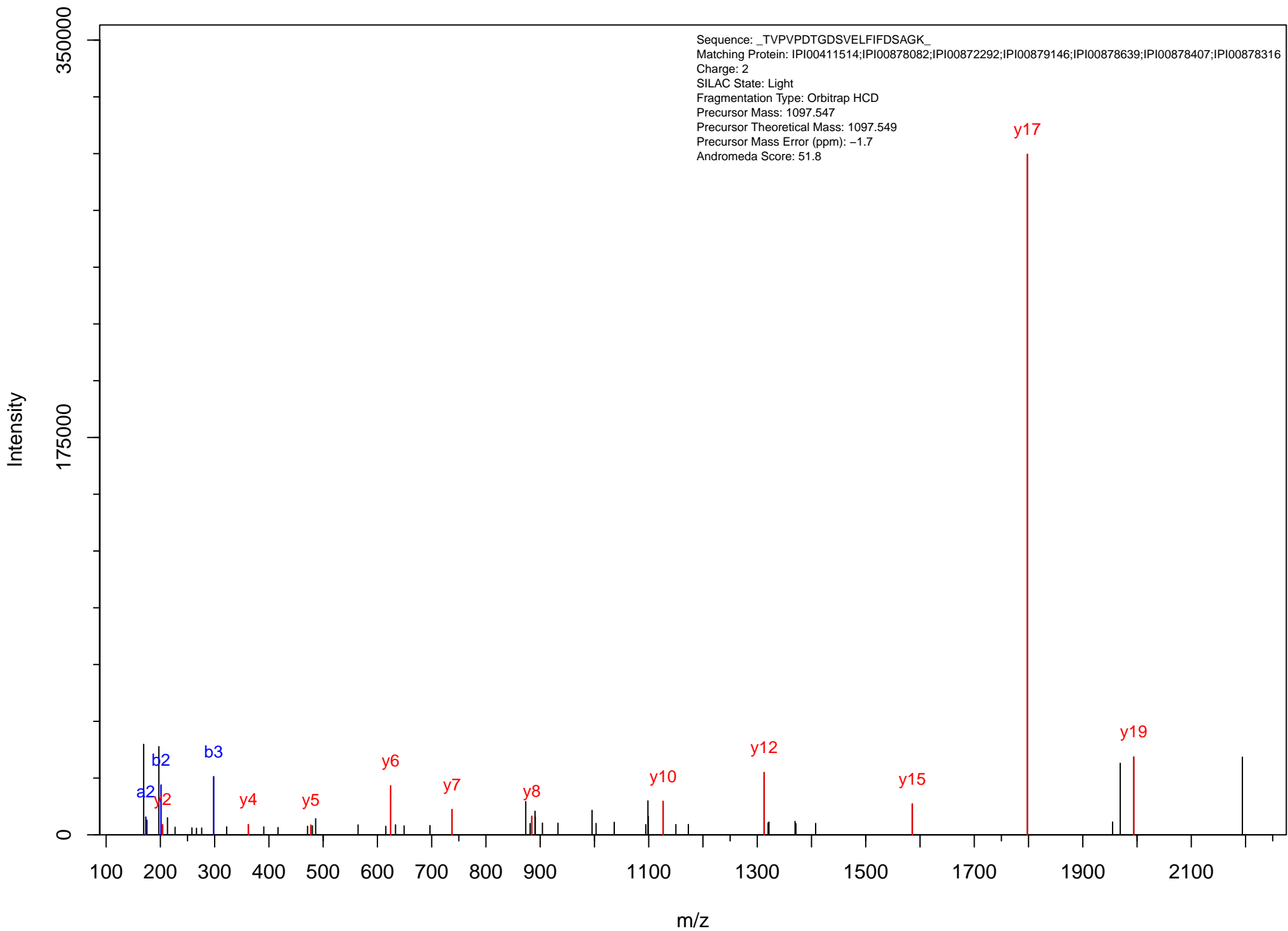


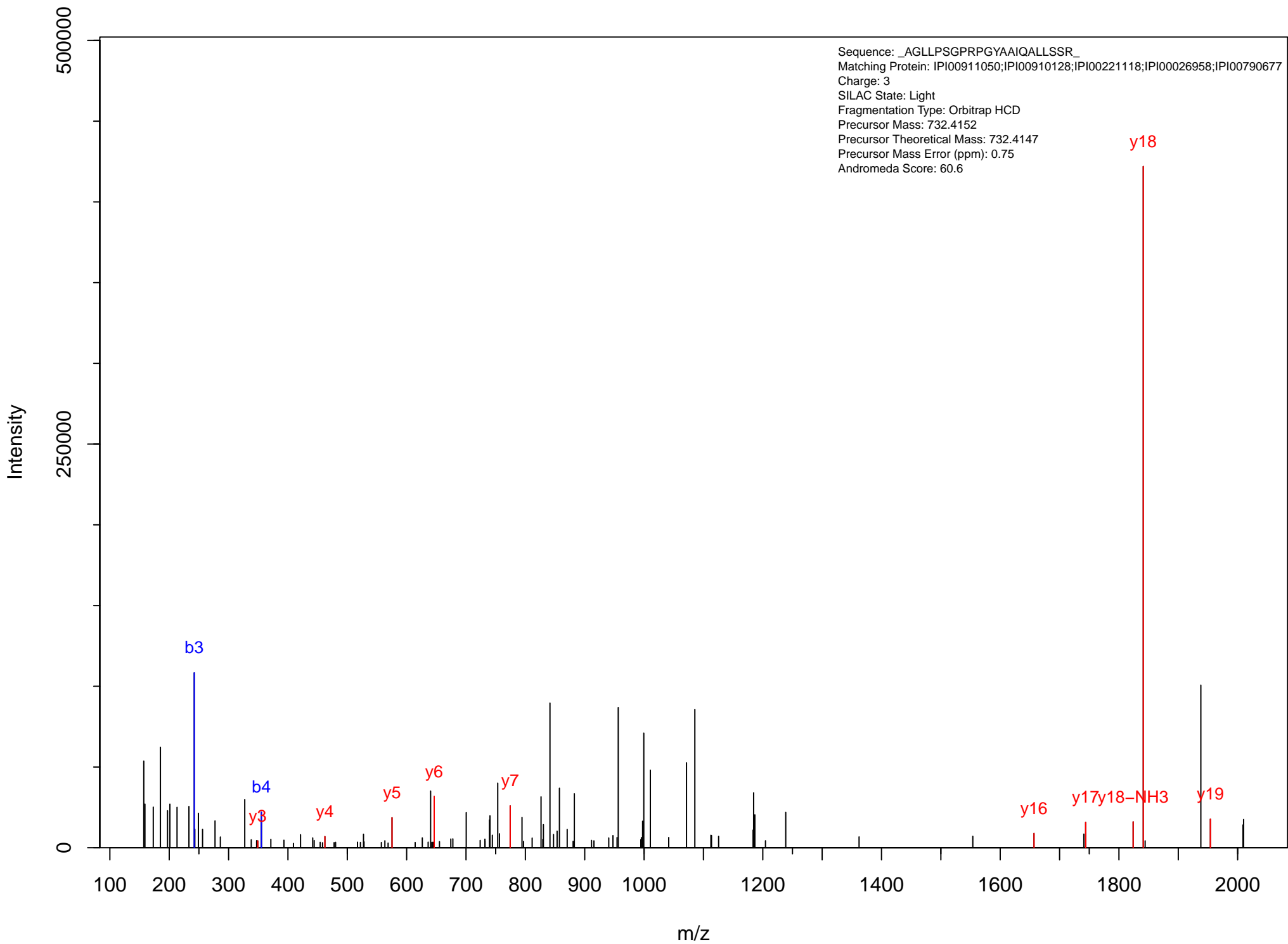


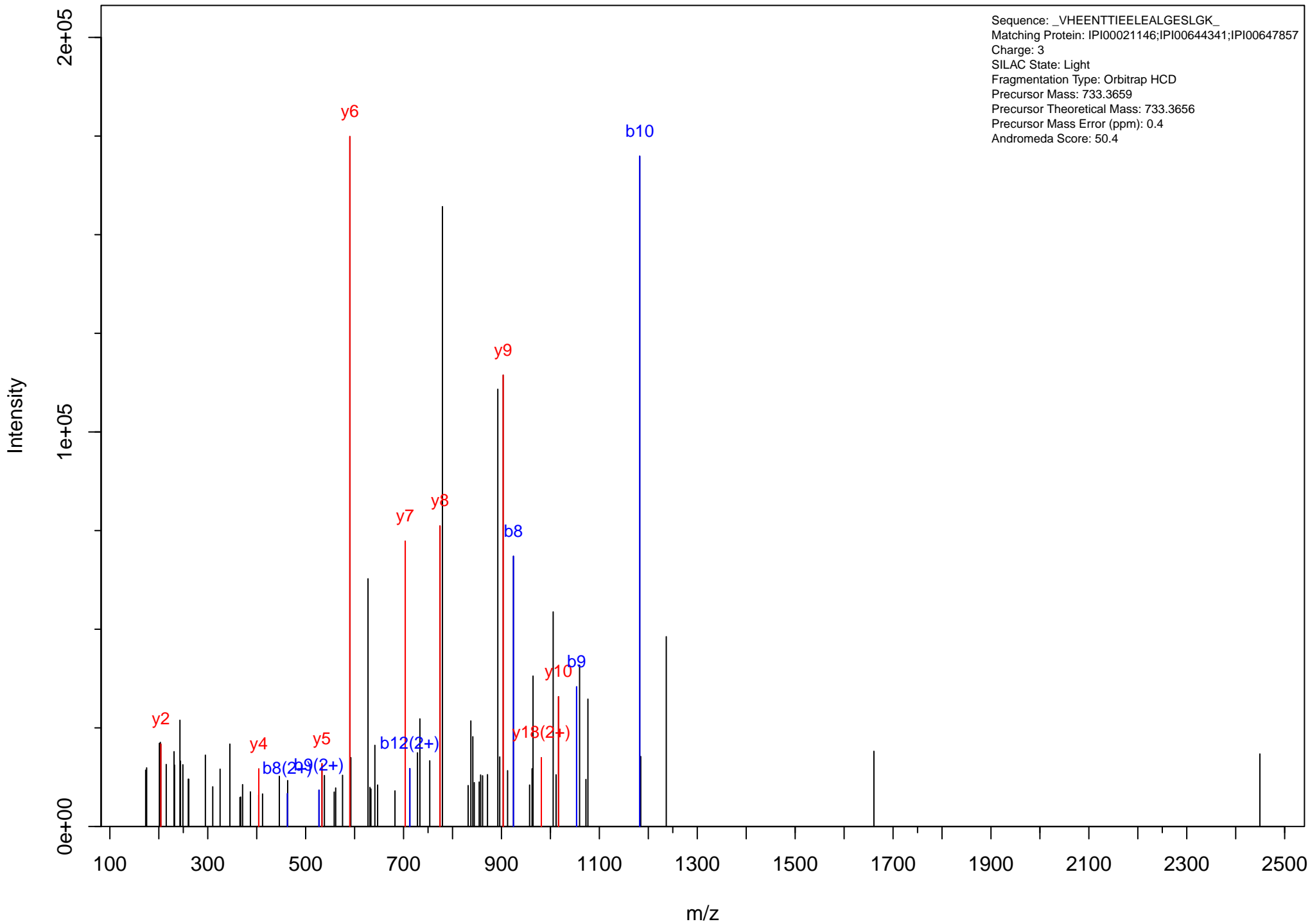


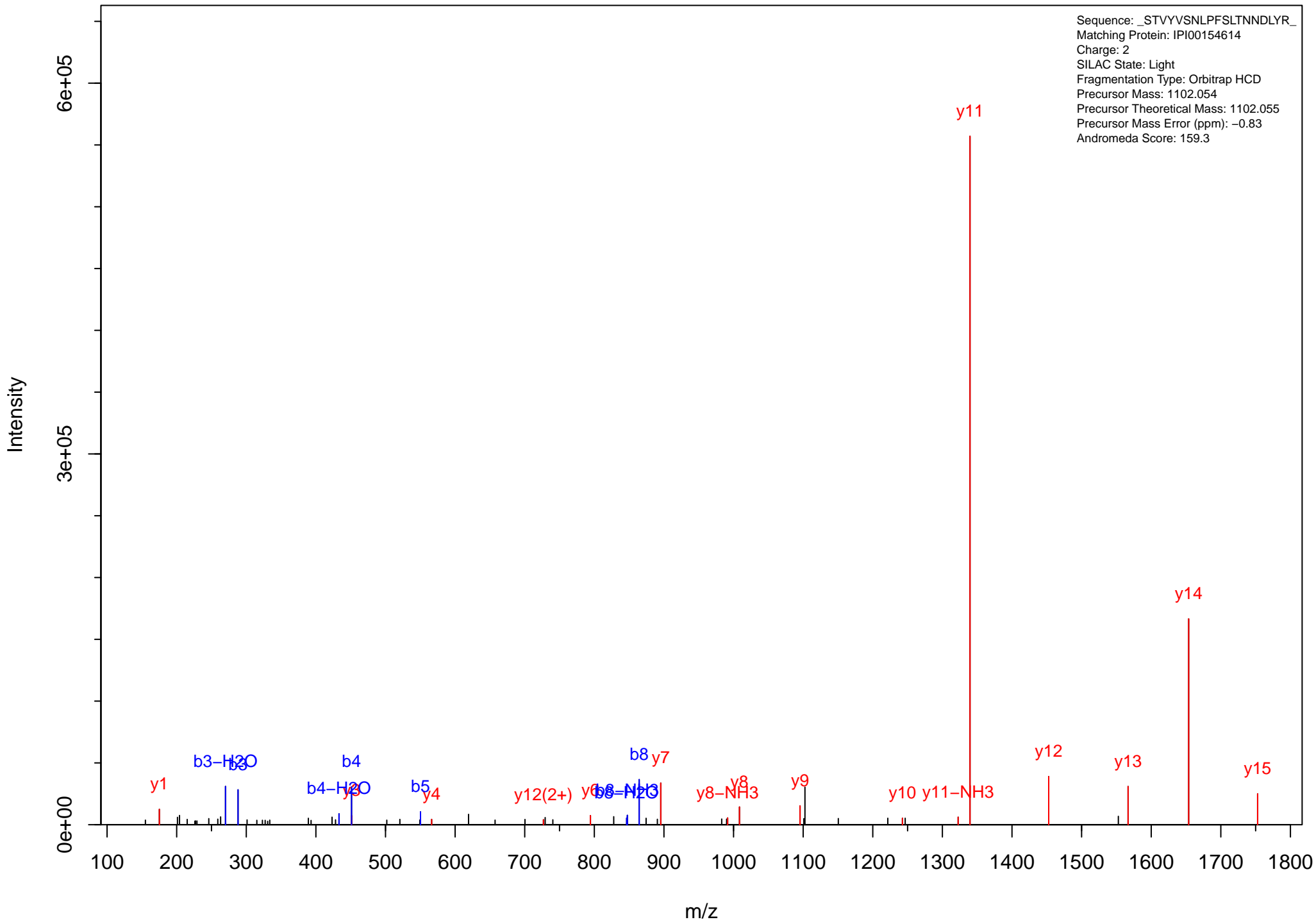


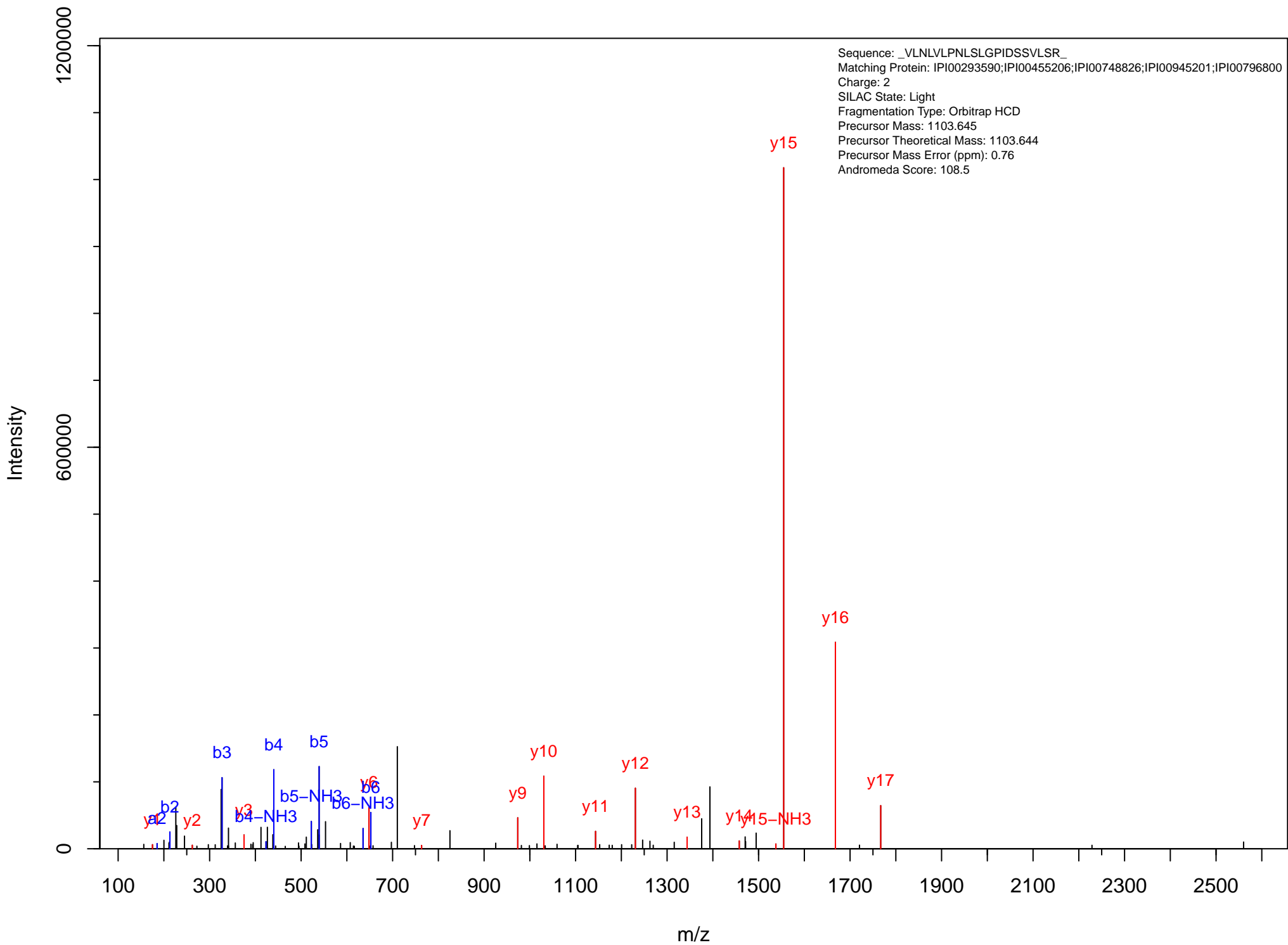


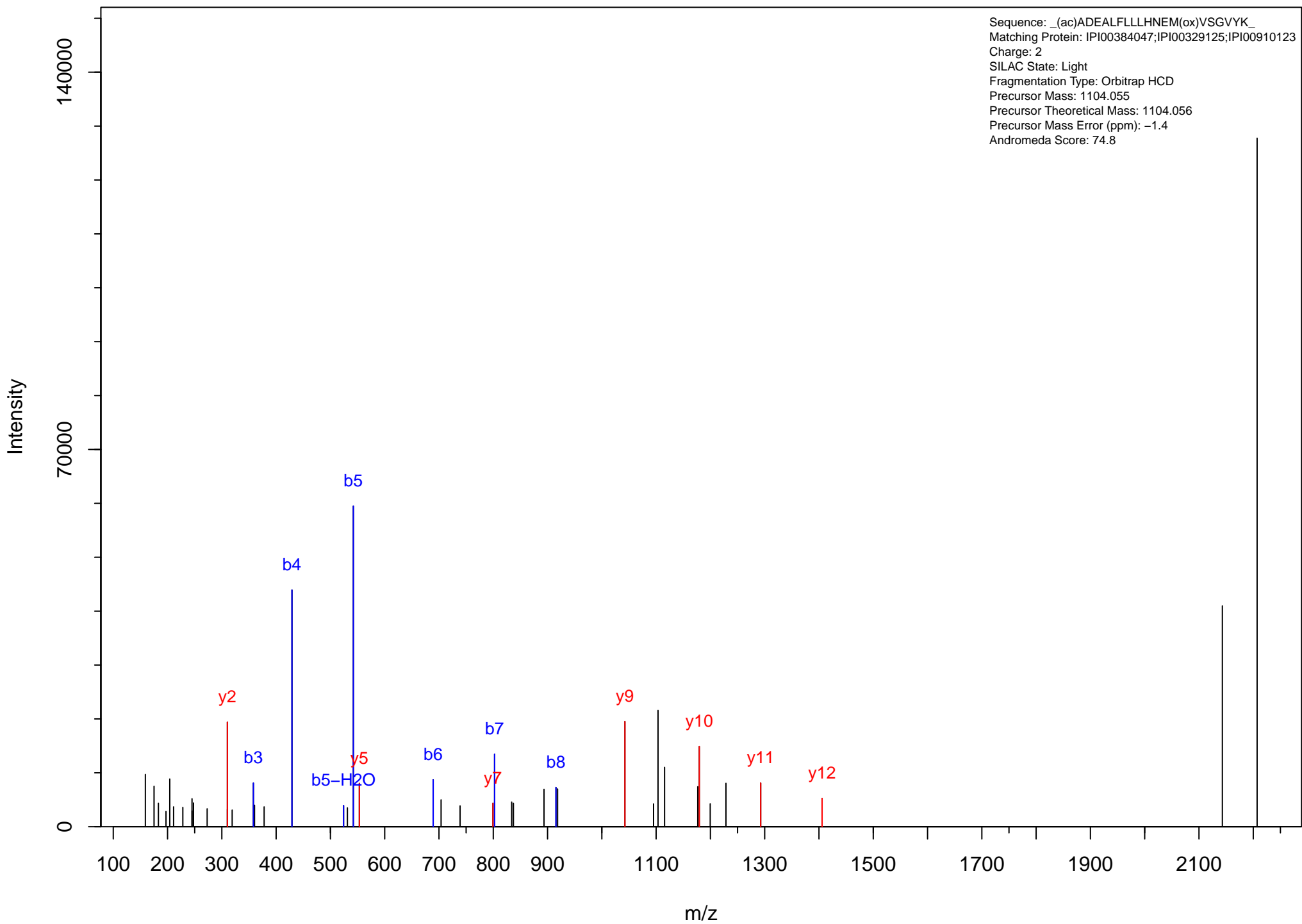


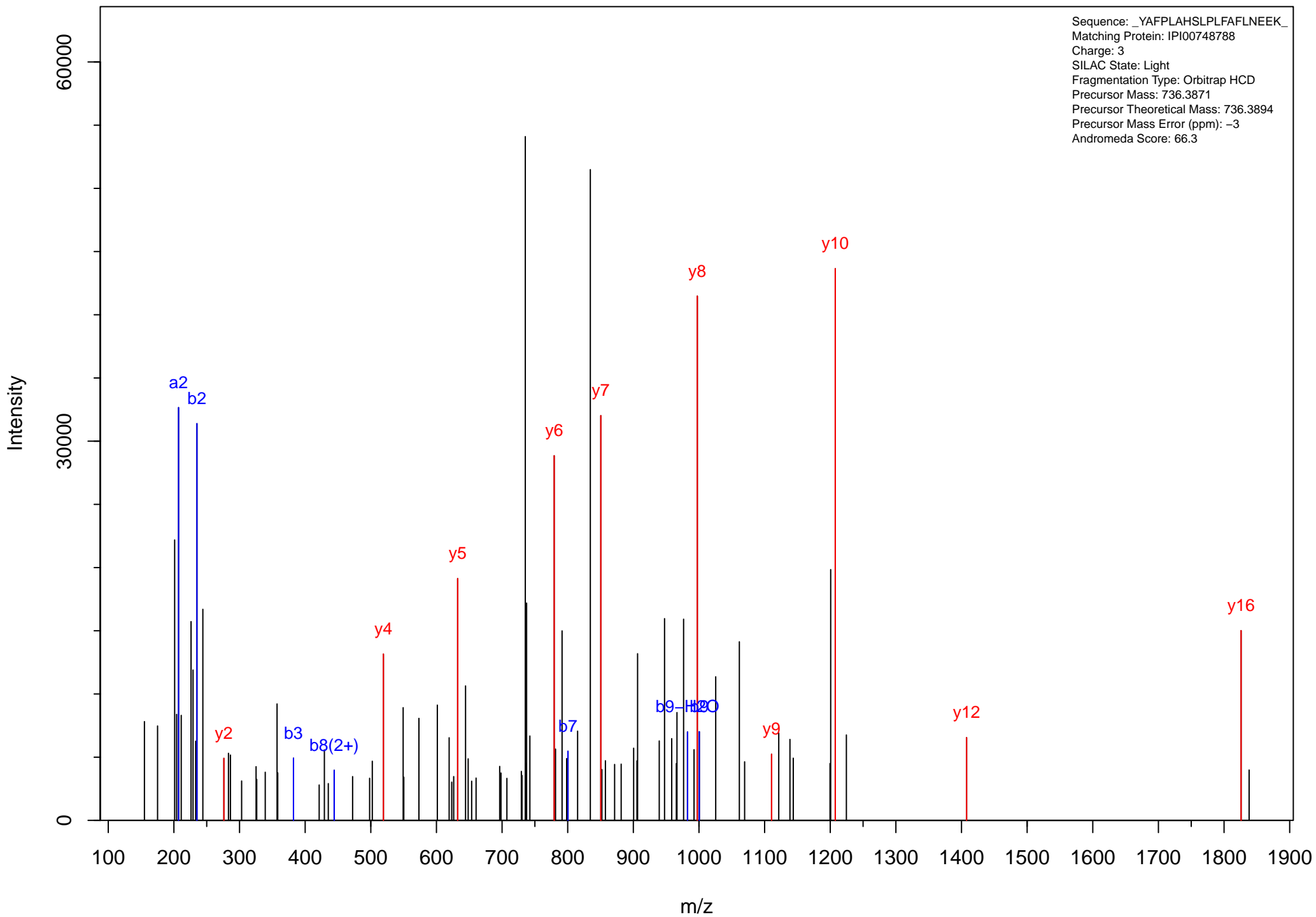


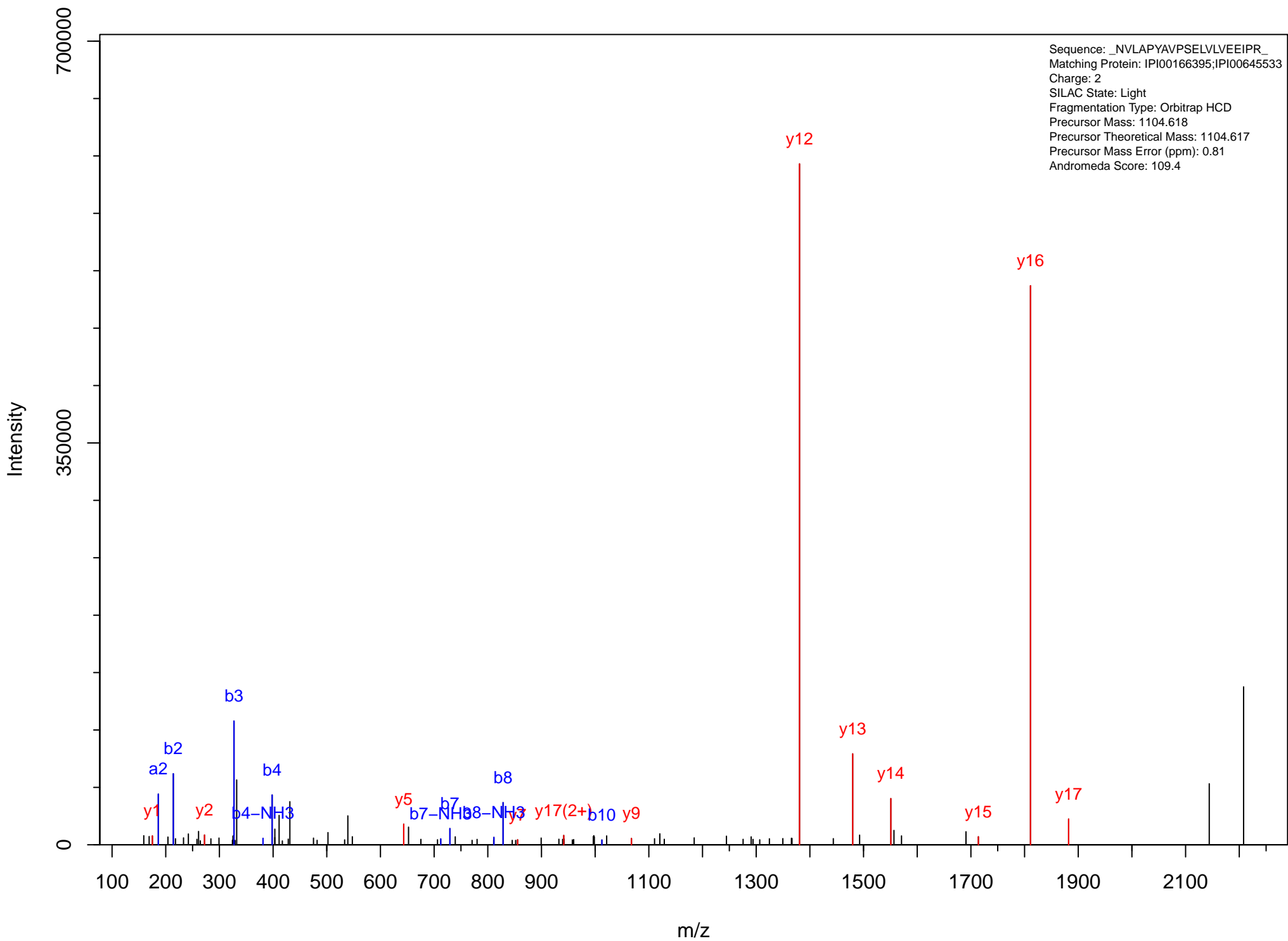


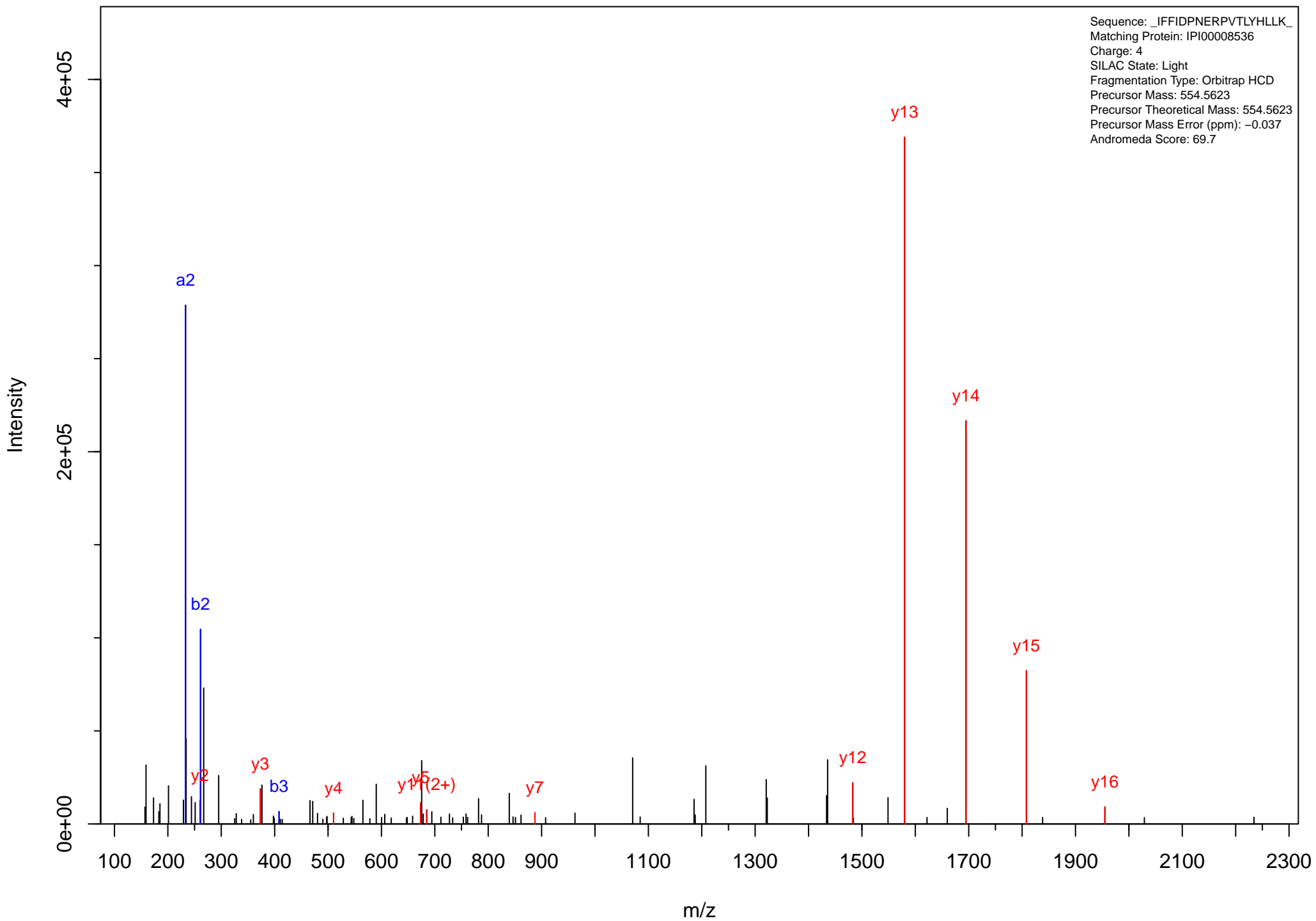


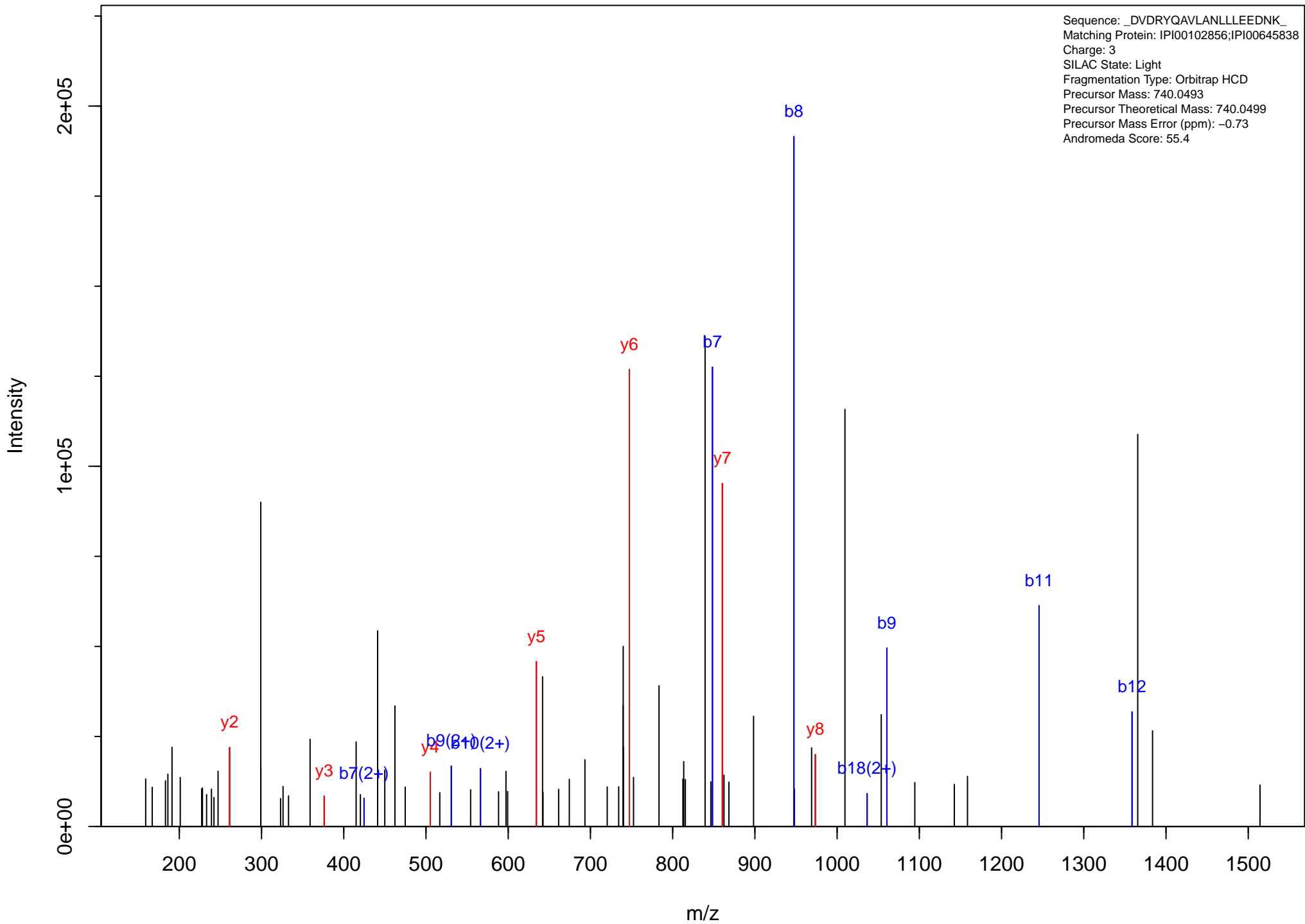


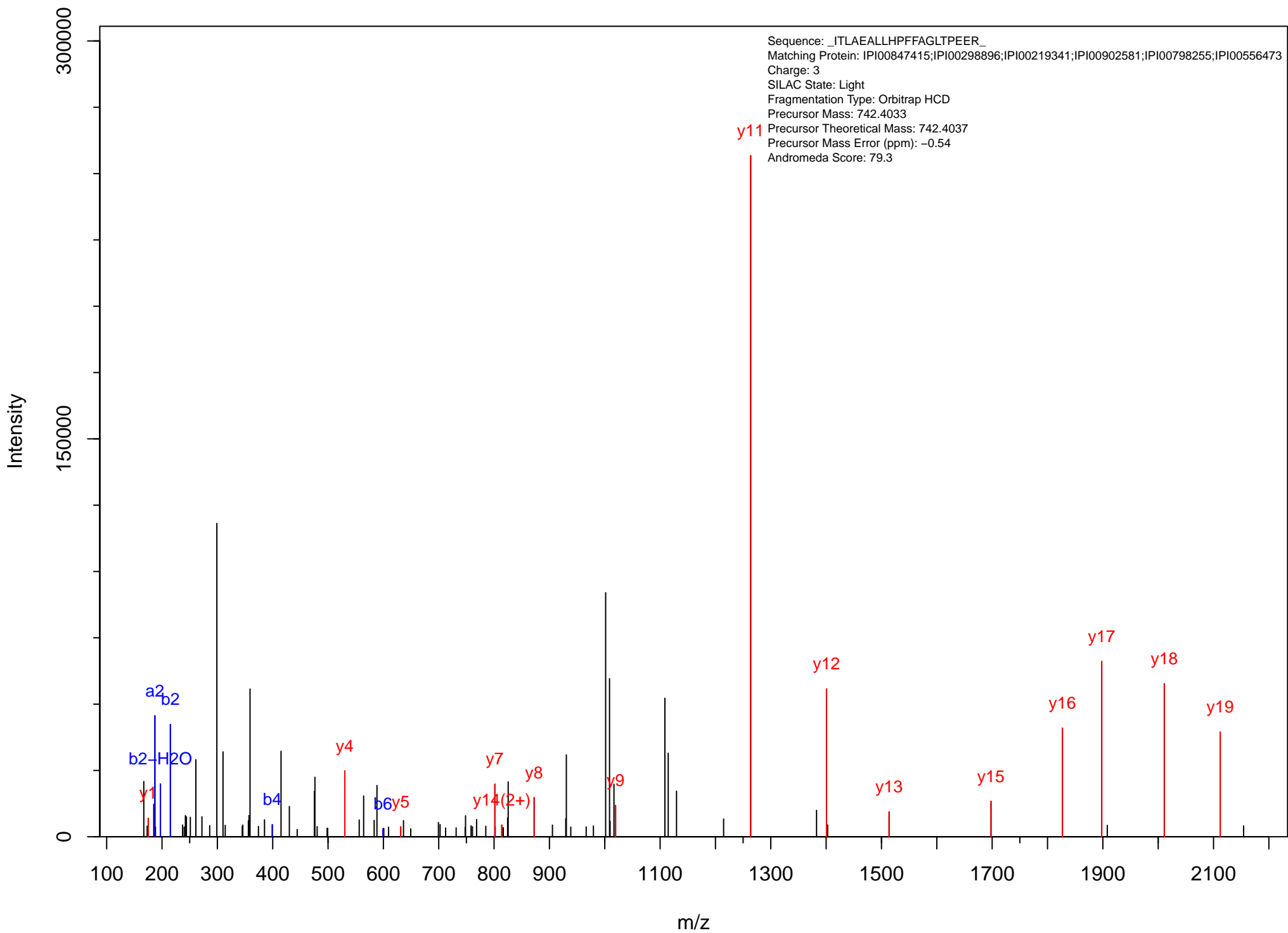


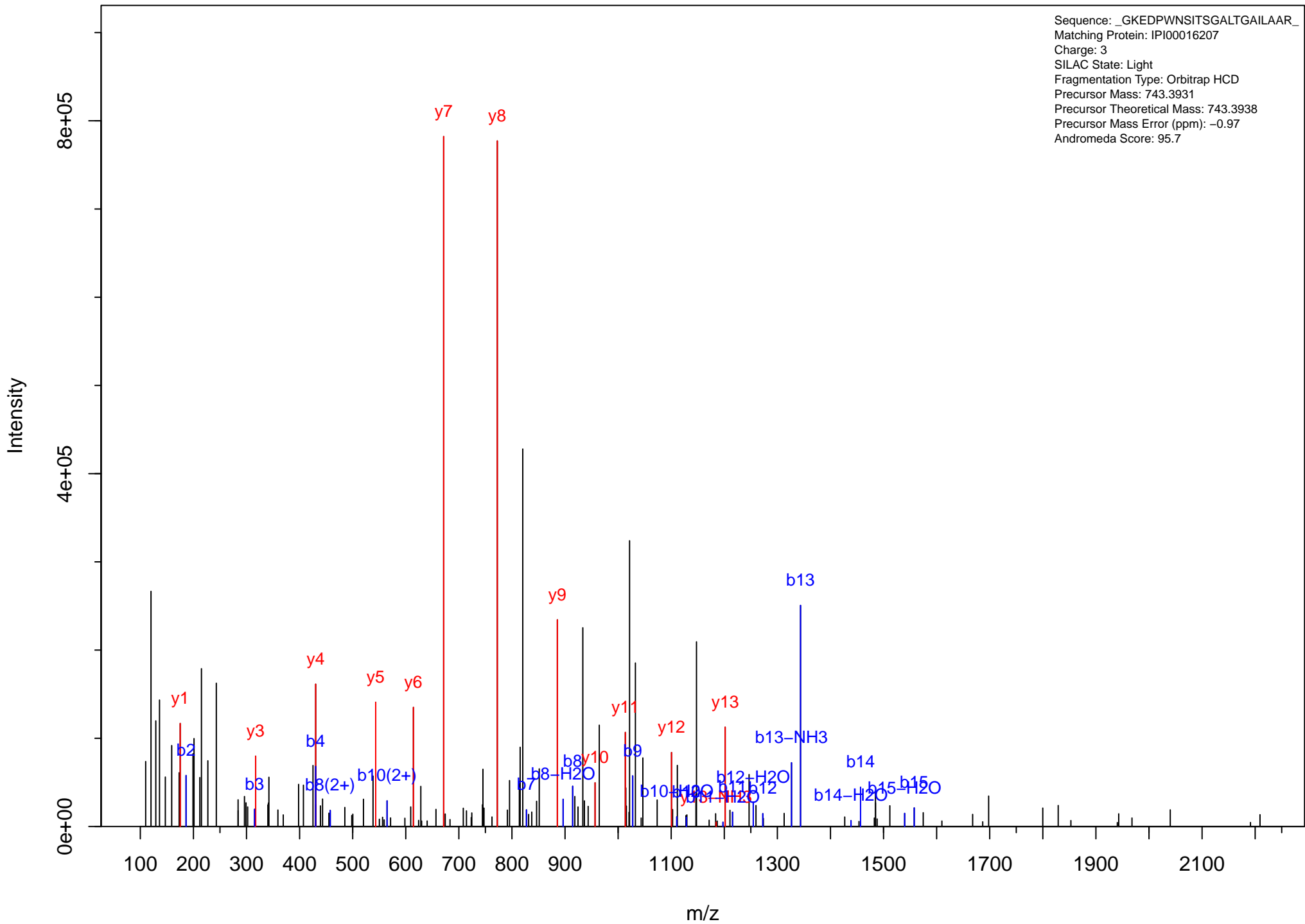


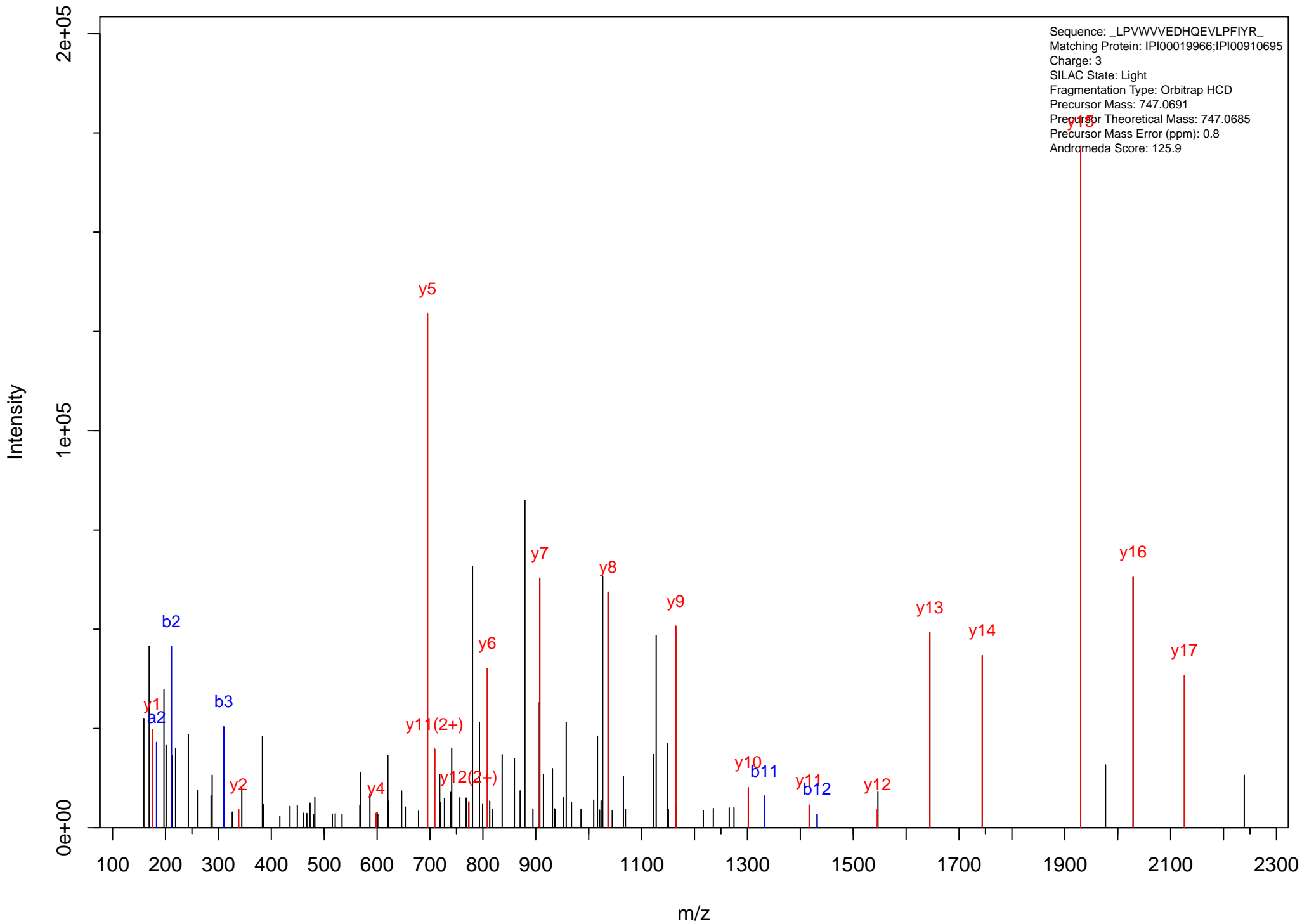


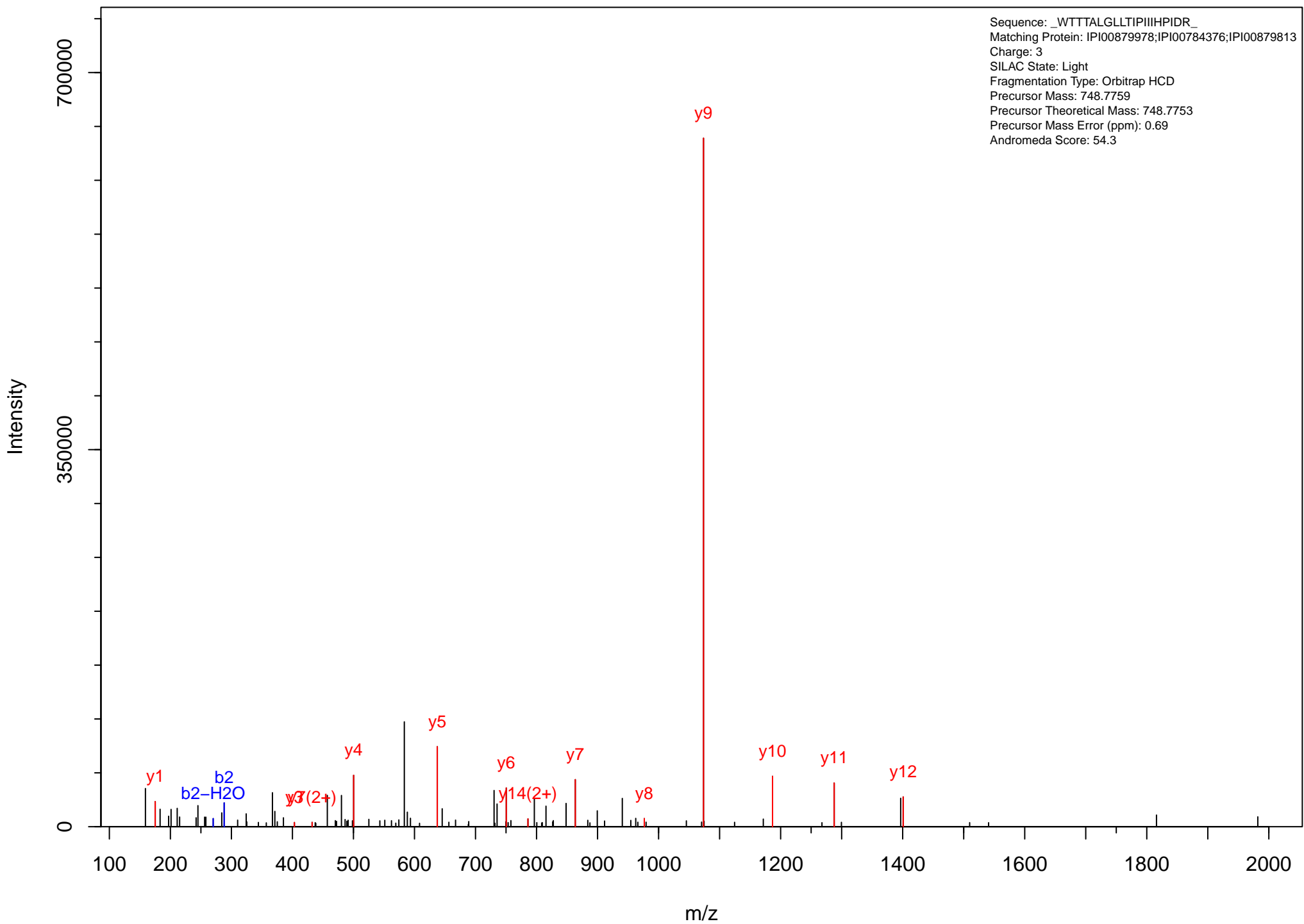


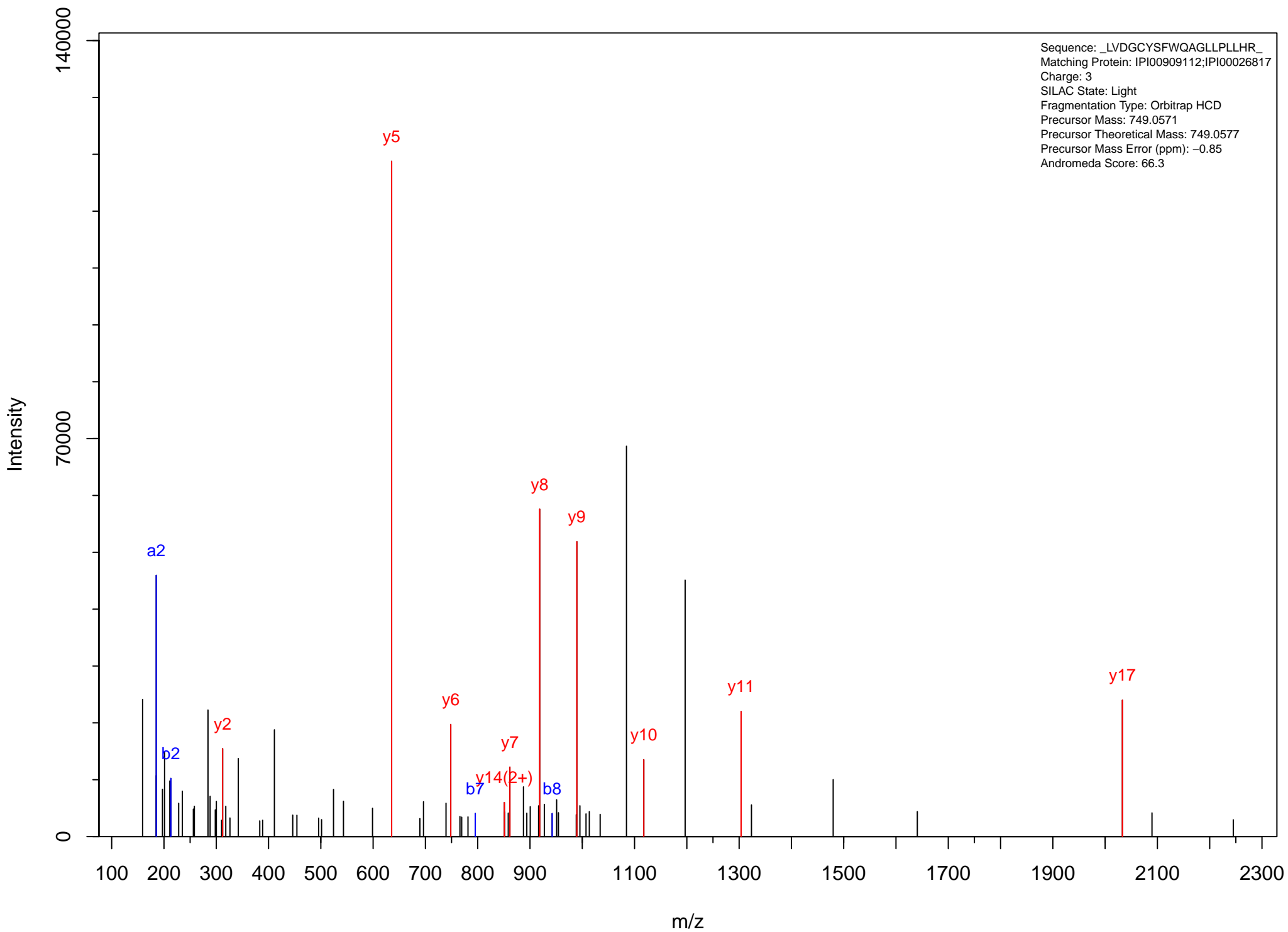


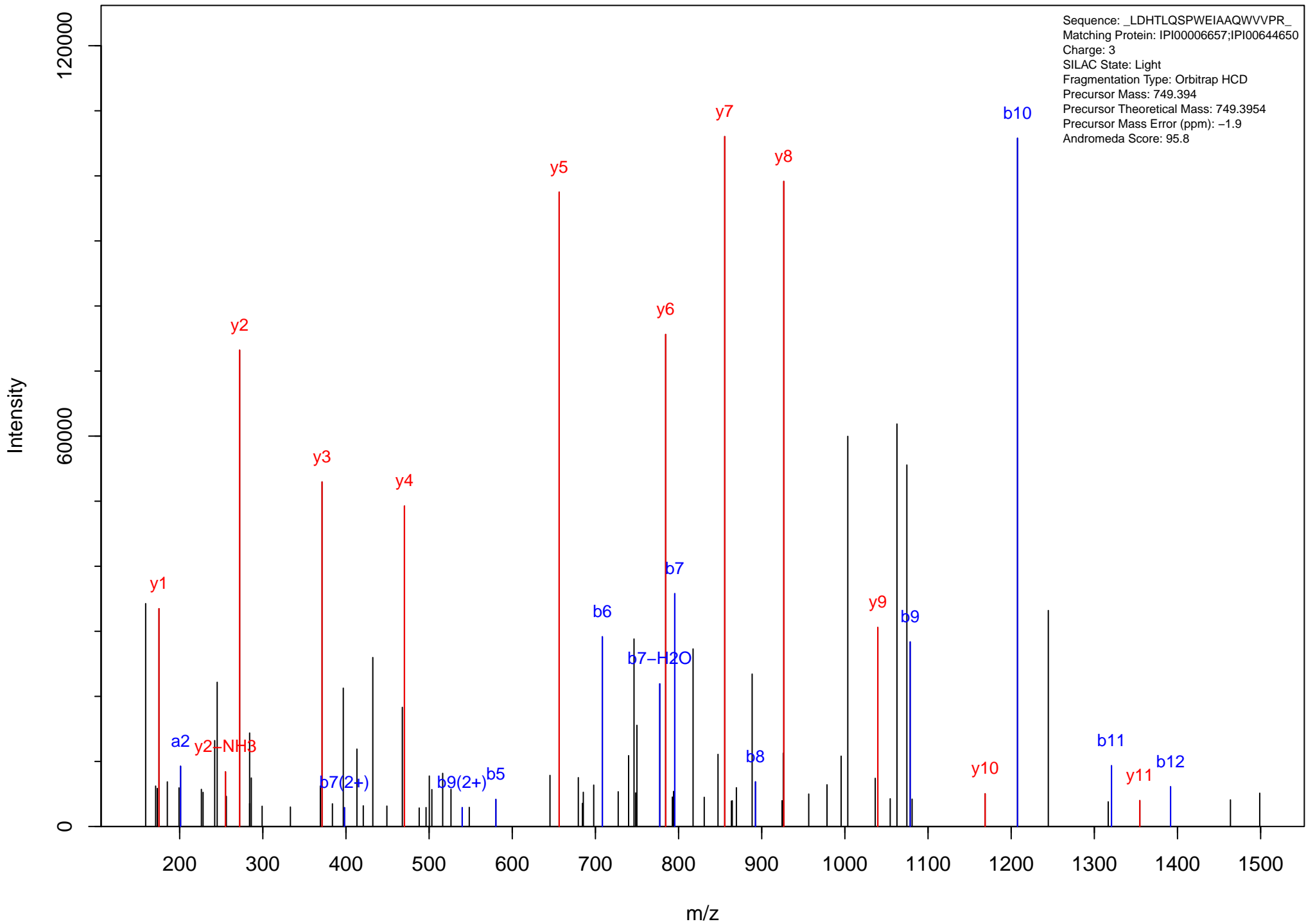


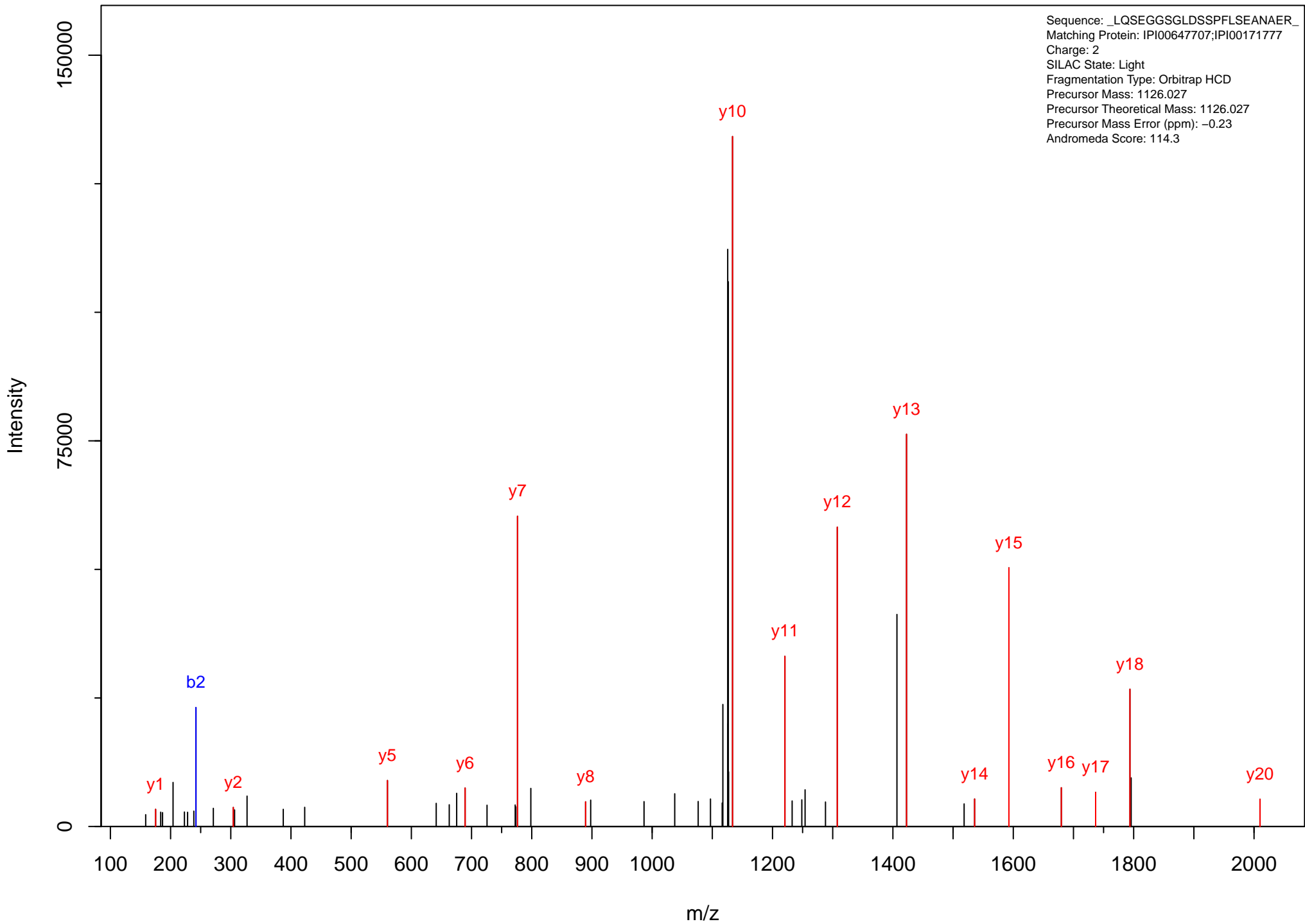


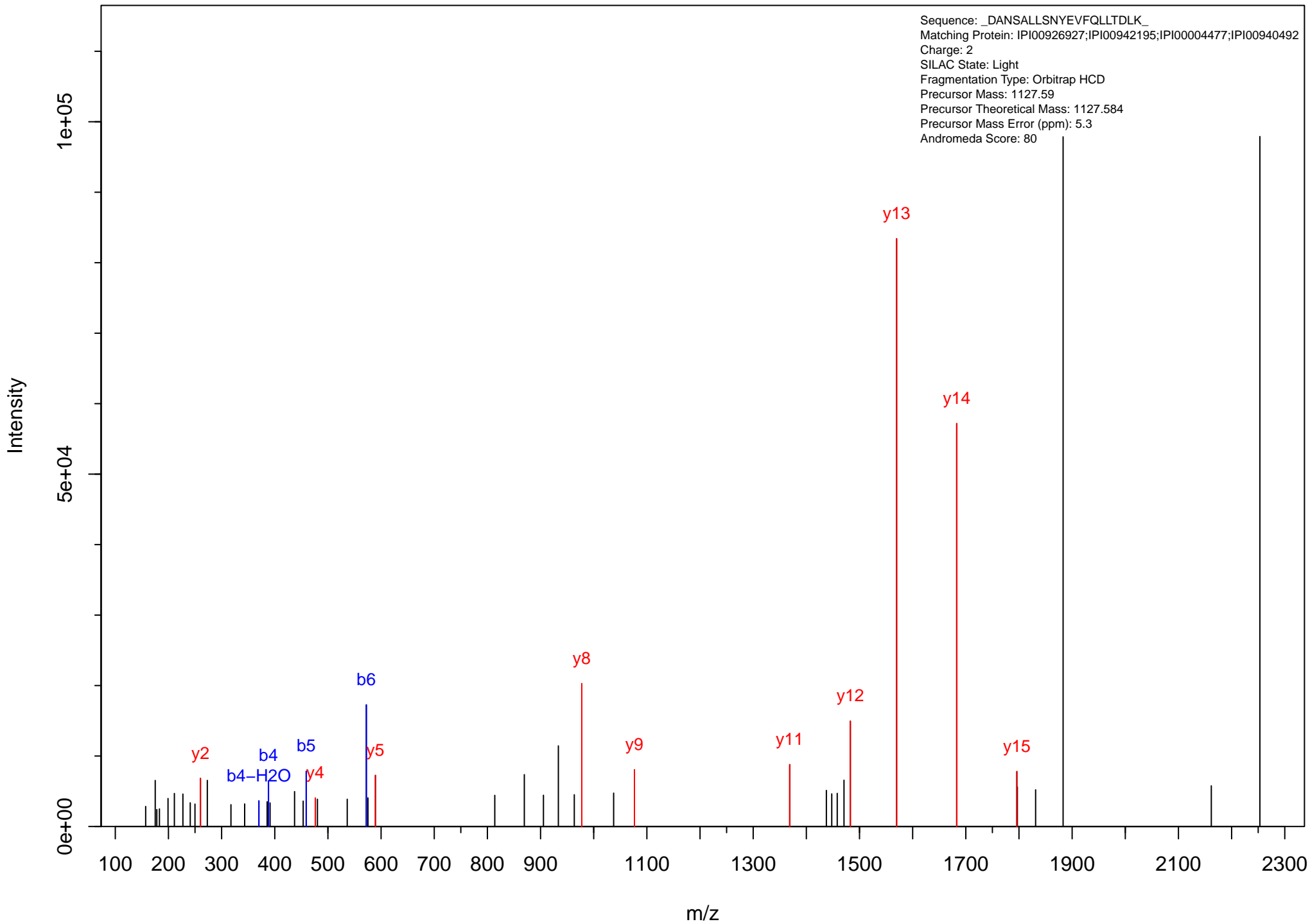


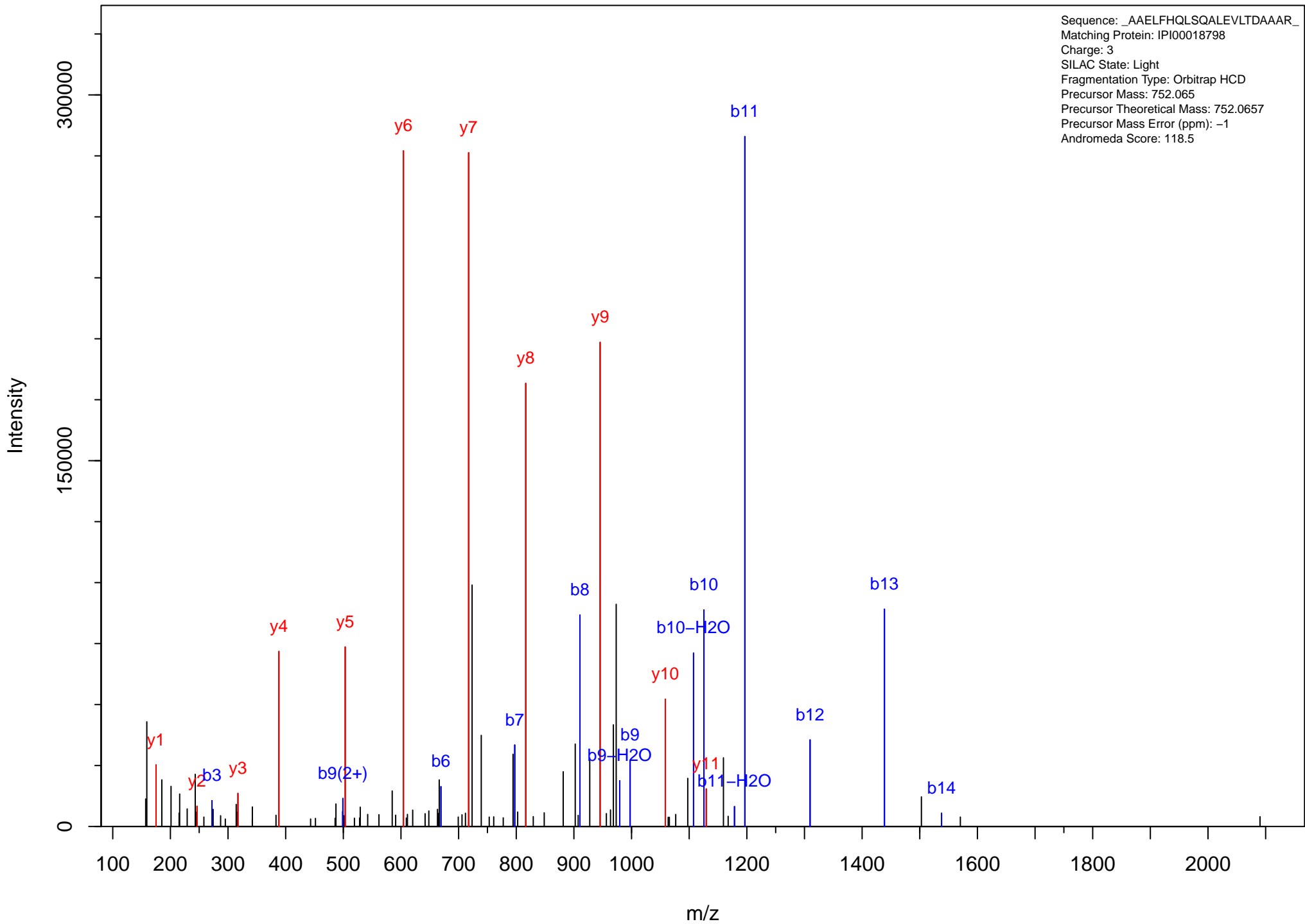


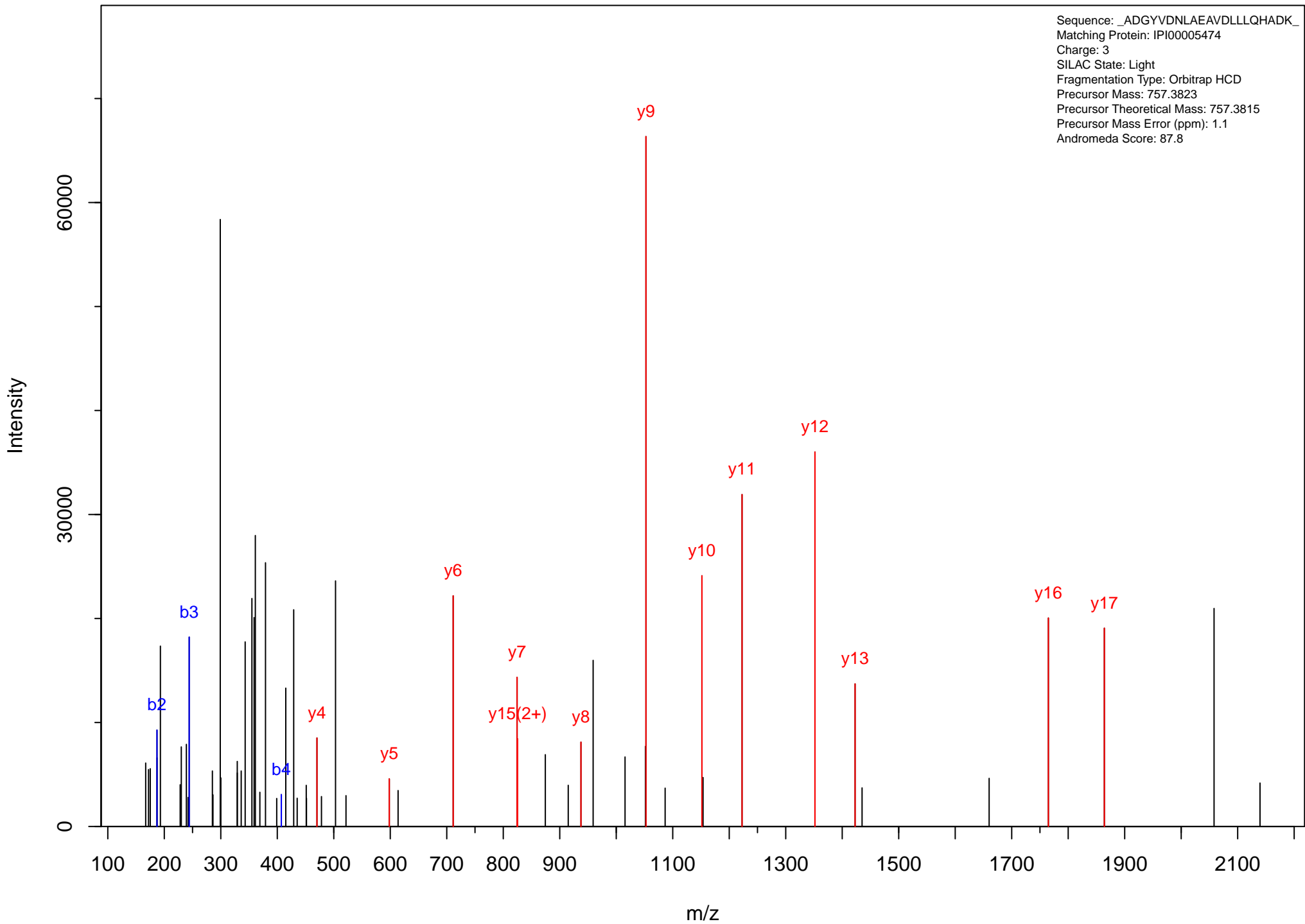


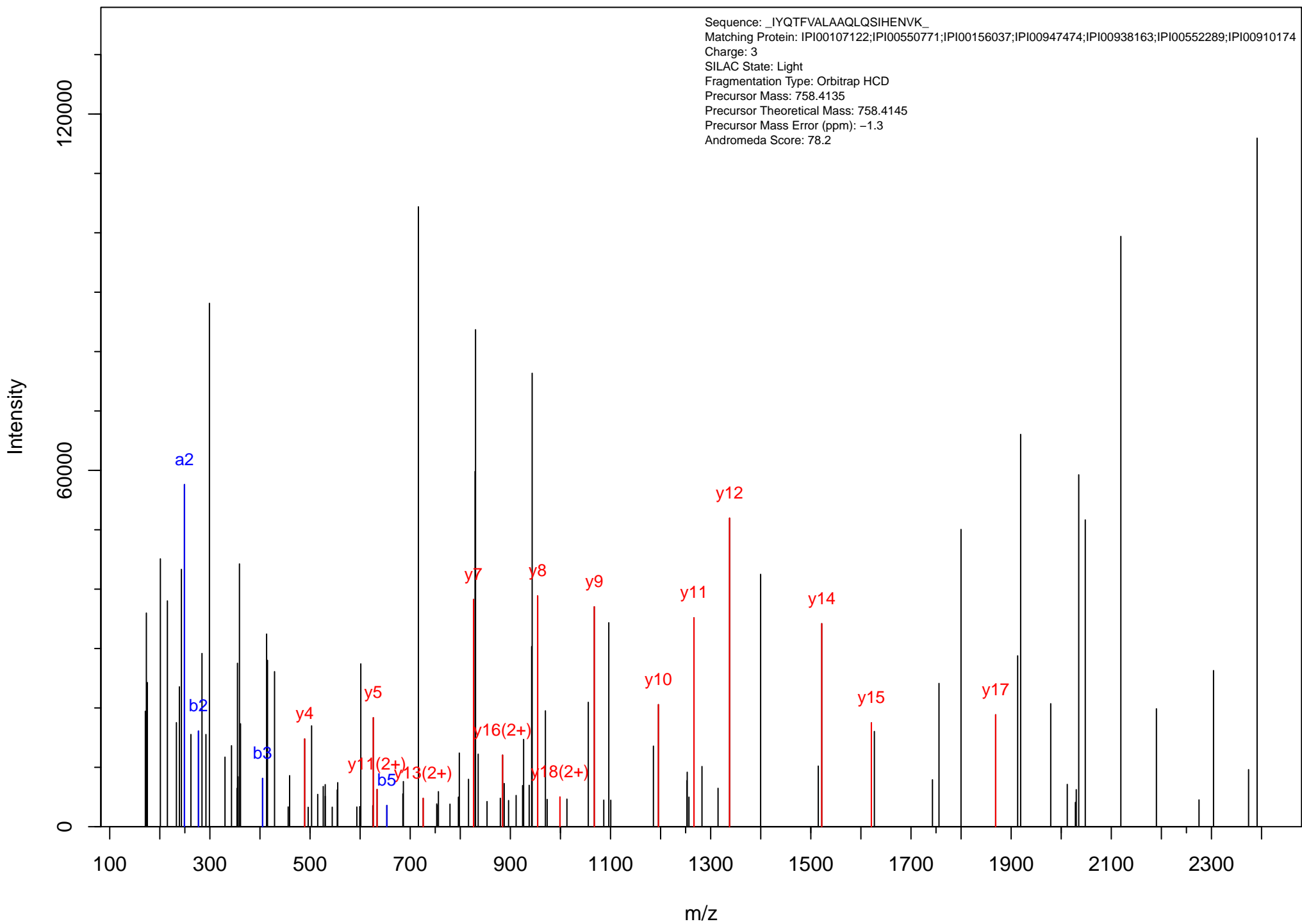


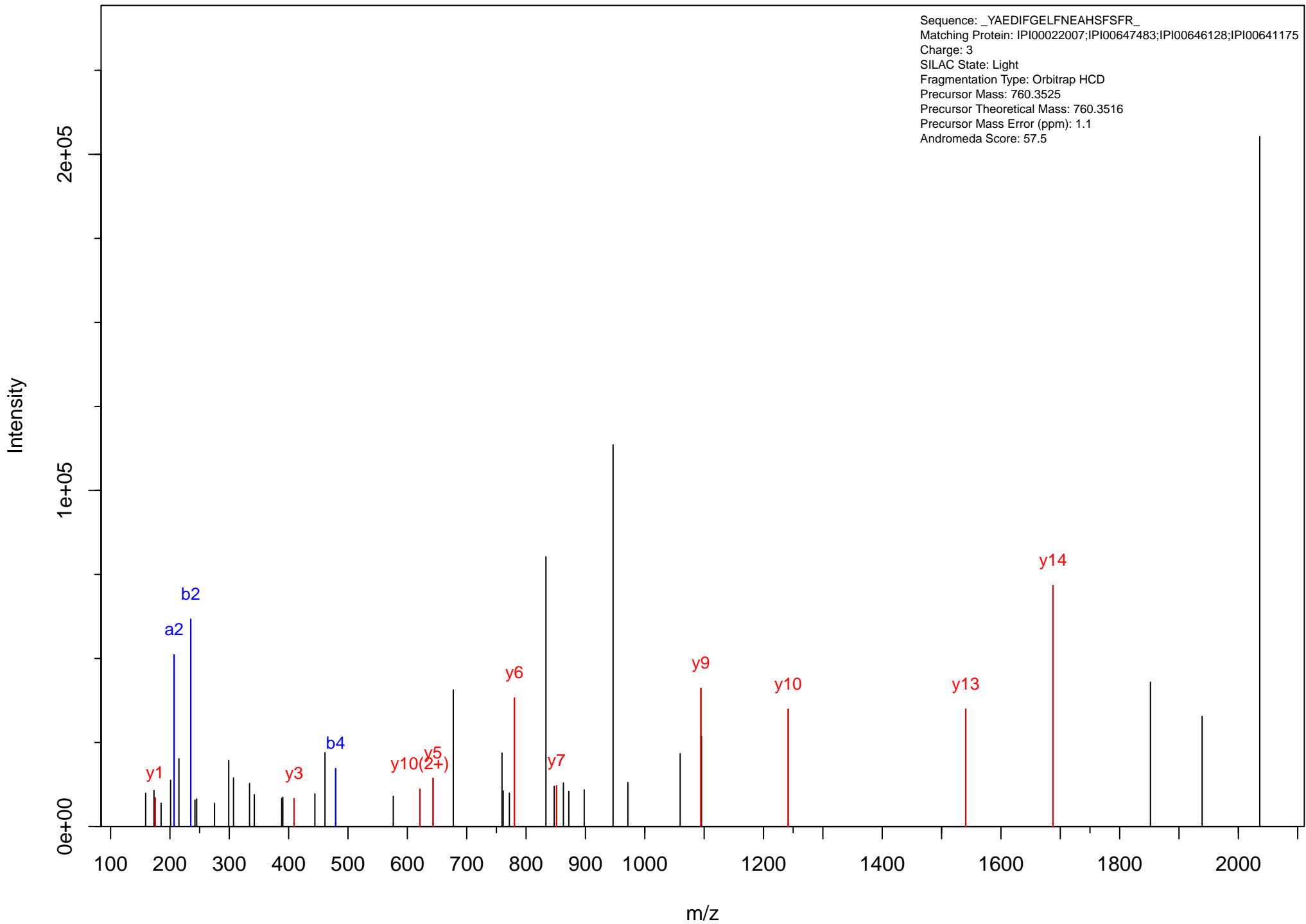


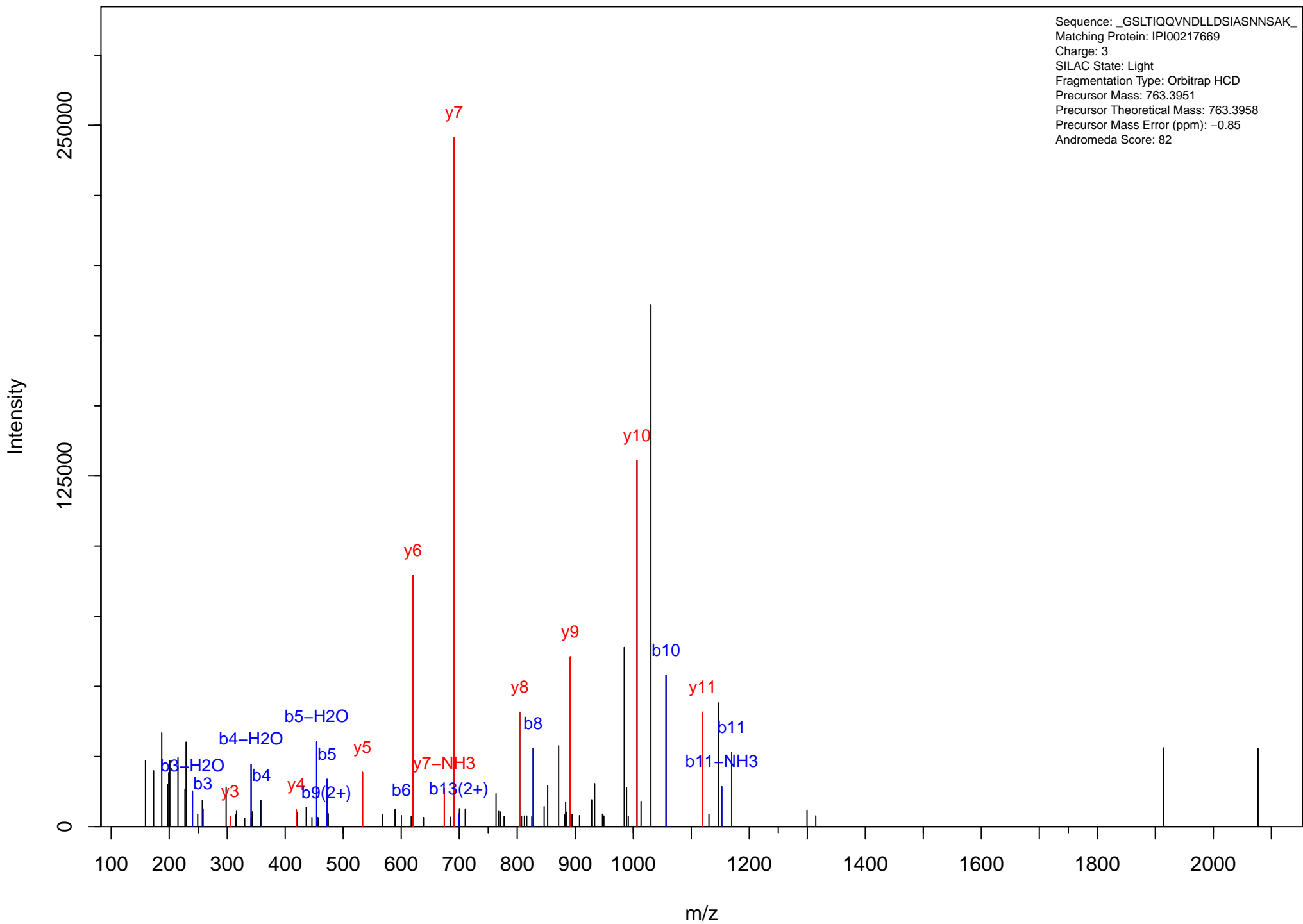


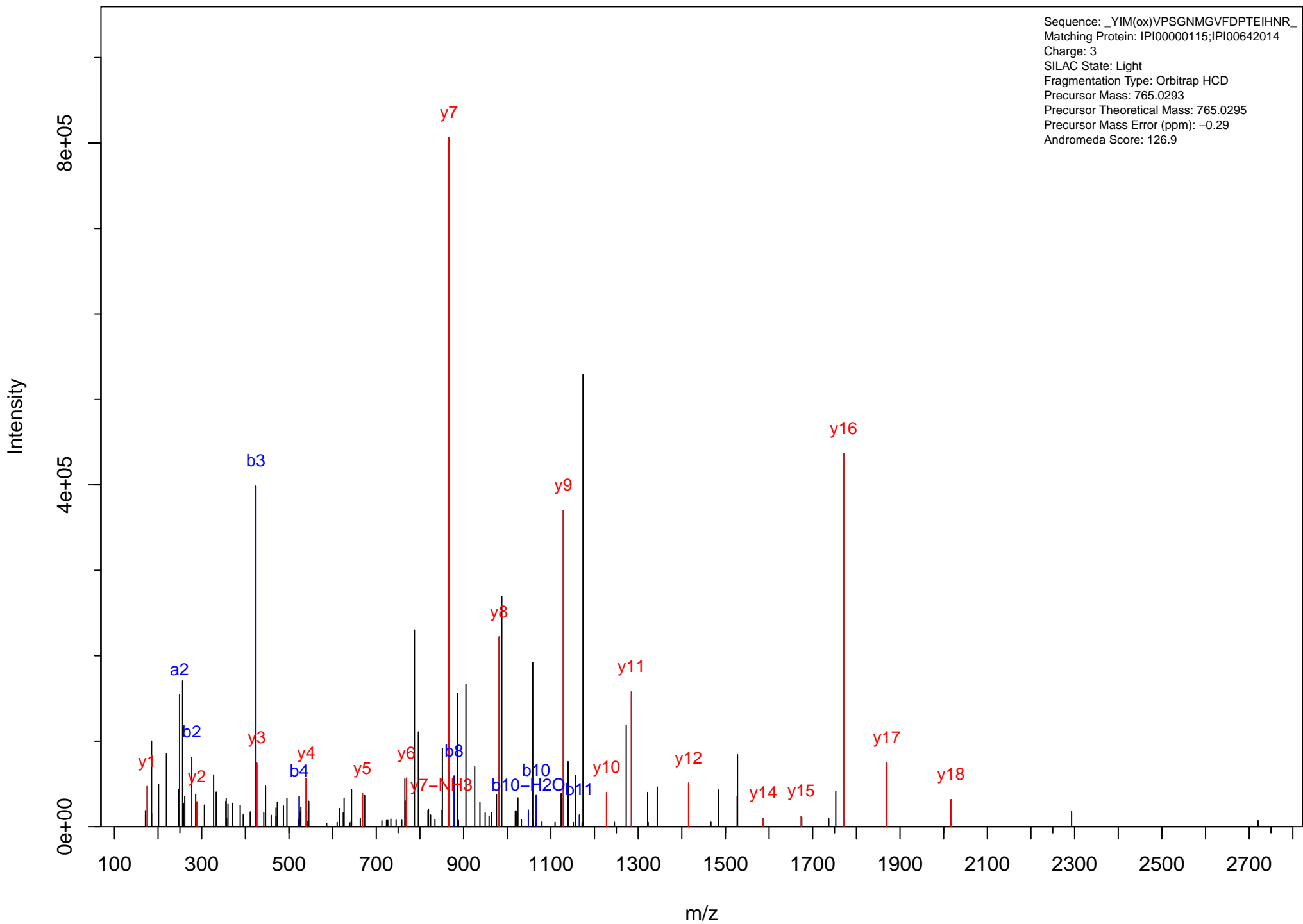


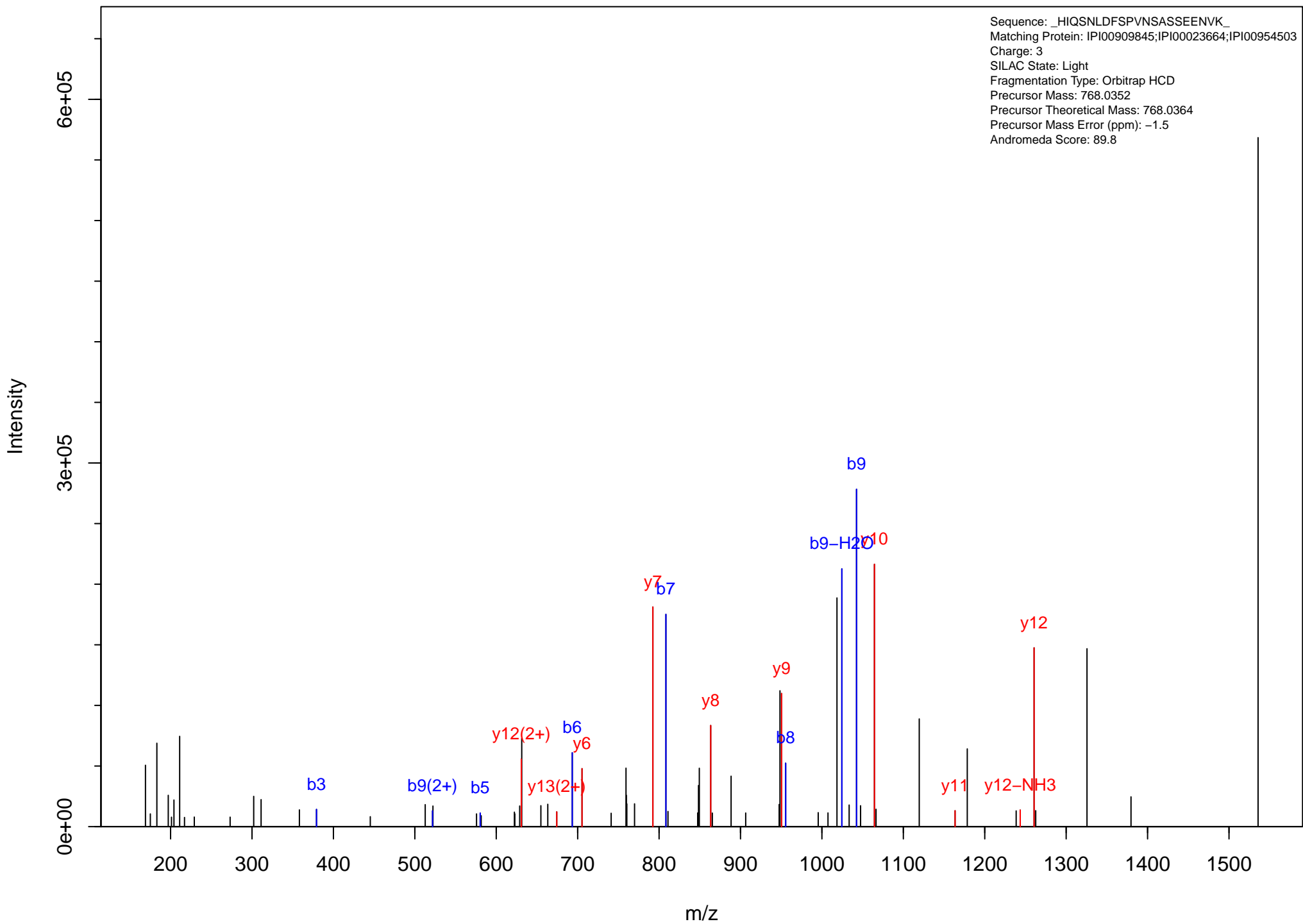


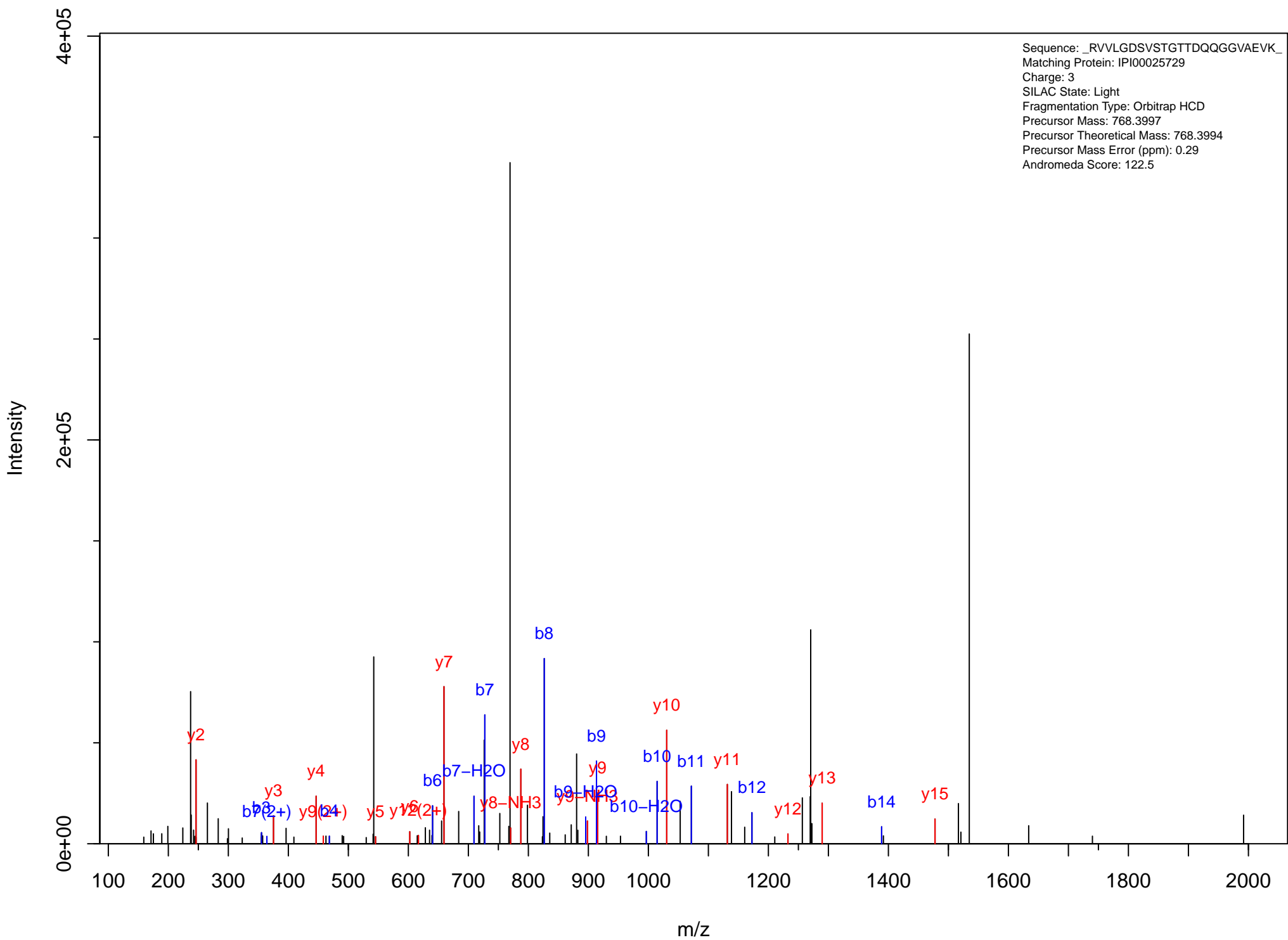












Intensity

4e+05

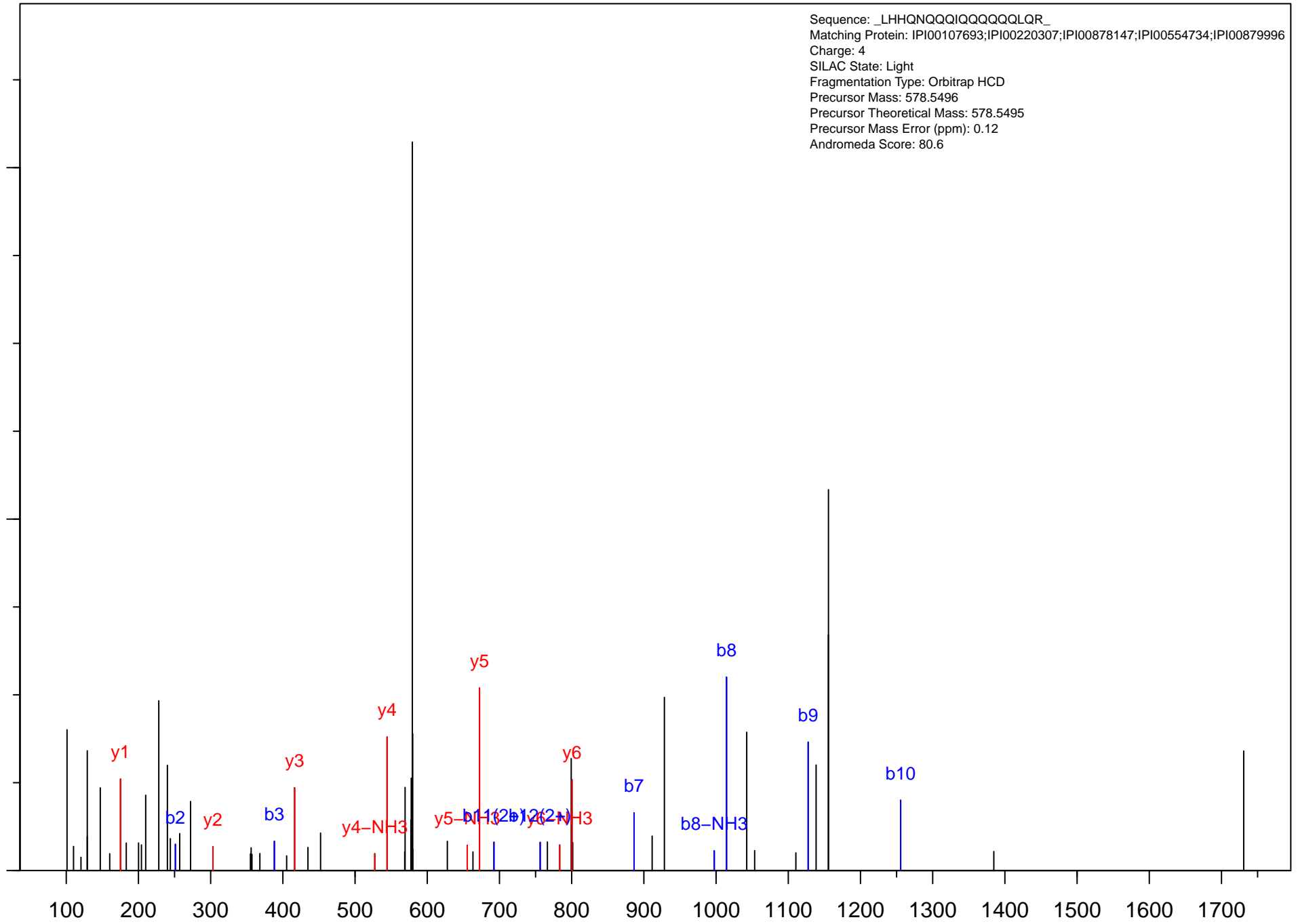
2e+05

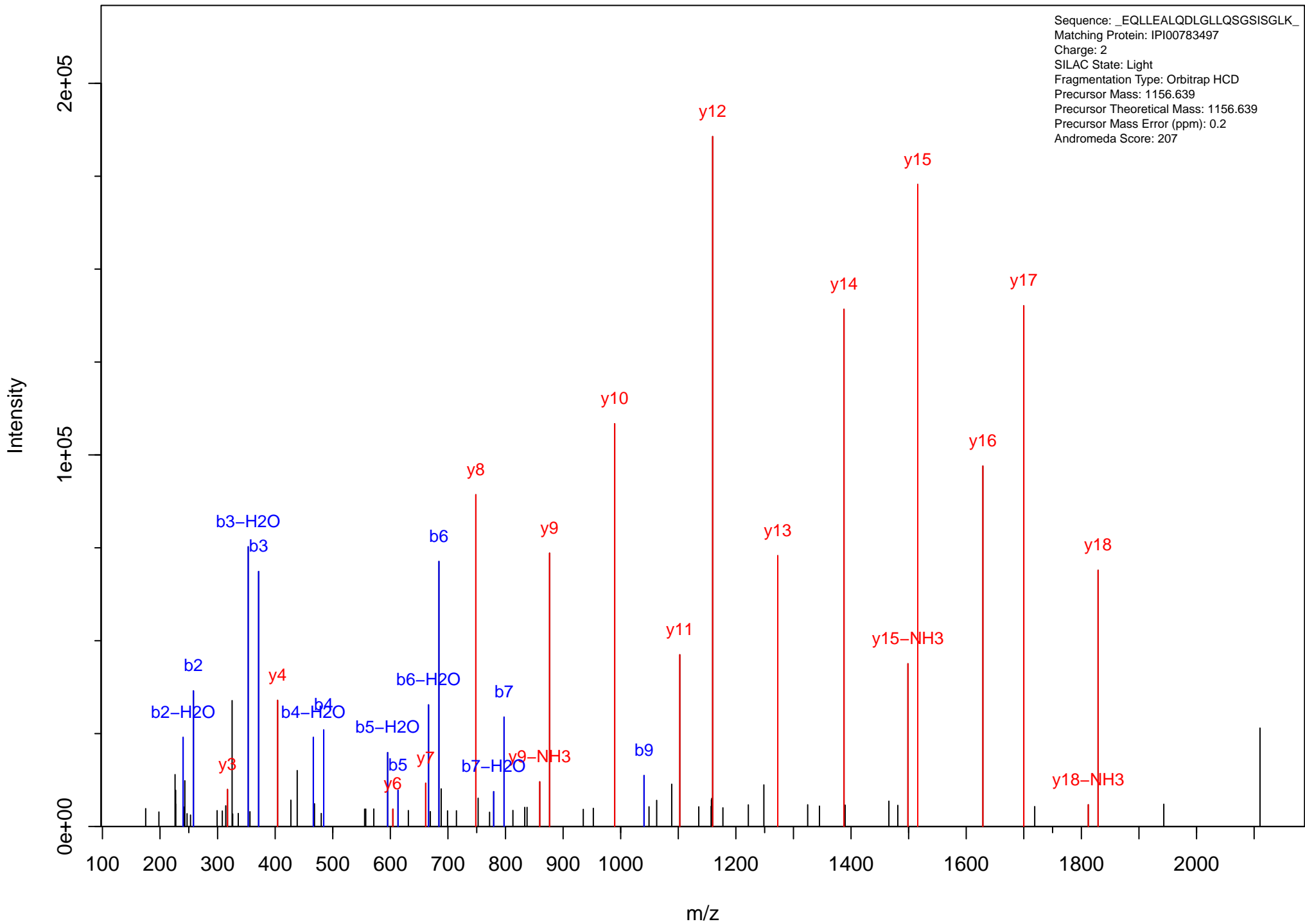
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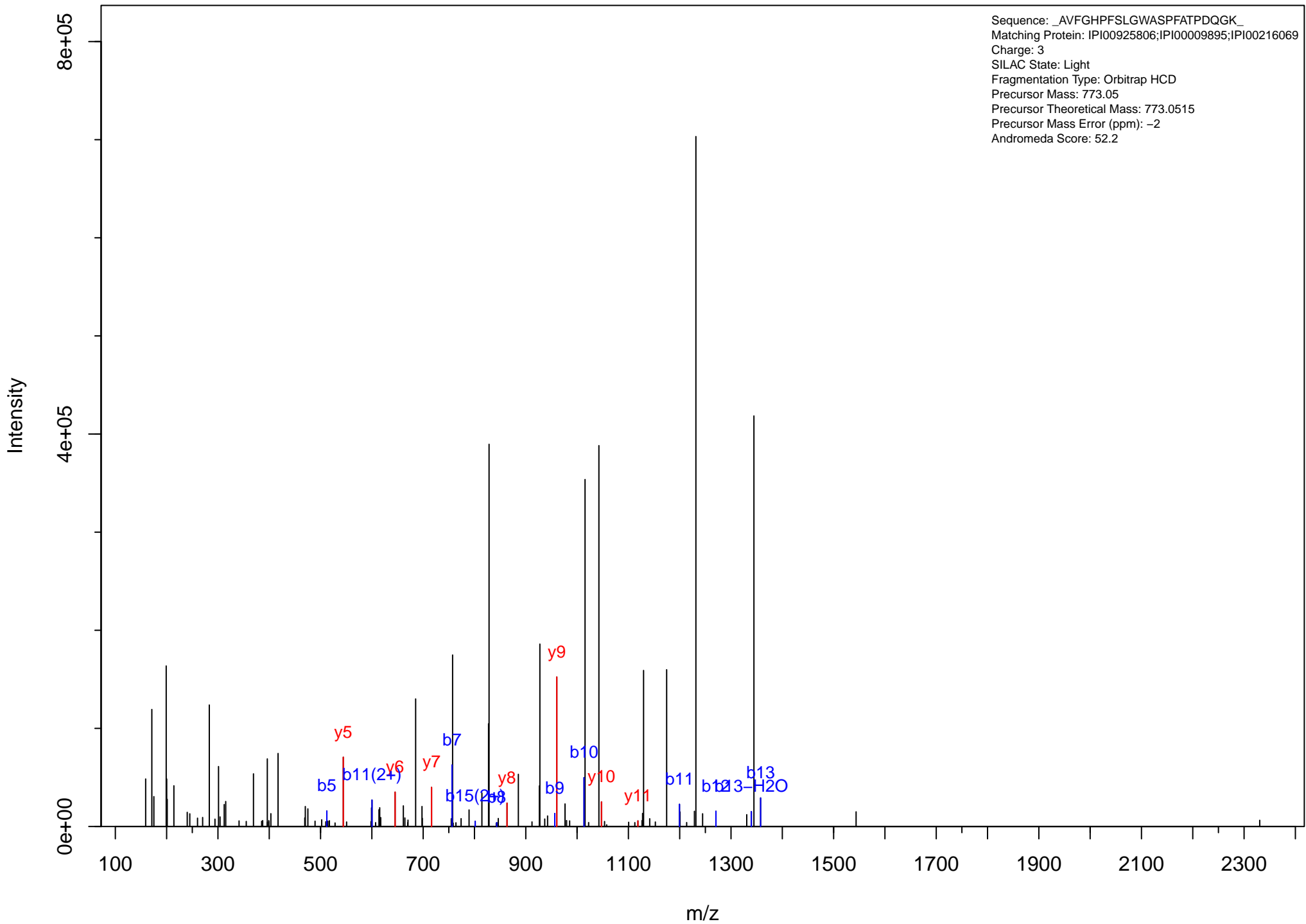
Sequence: _LHHQNQQIQQQQQLQR_
Matching Protein: IPI00107693;IPI00220307;IPI00878147;IPI00554734;IPI00879996
Charge: 4
SILAC State: Light
Fragmentation Type: Orbitrap HCD
Precursor Mass: 578.5496
Precursor Theoretical Mass: 578.5495
Precursor Mass Error (ppm): 0.12
Andromeda Score: 80.6

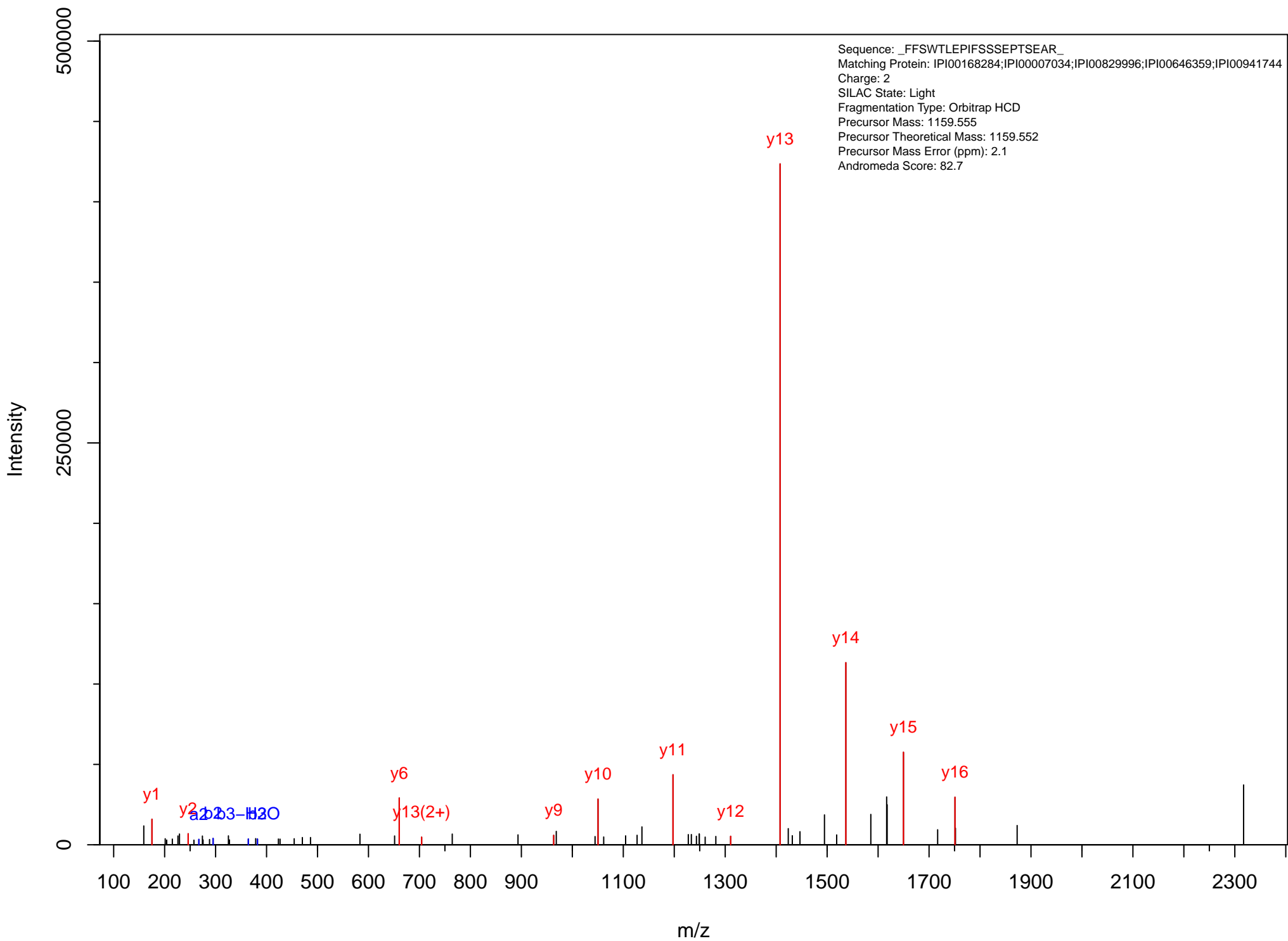
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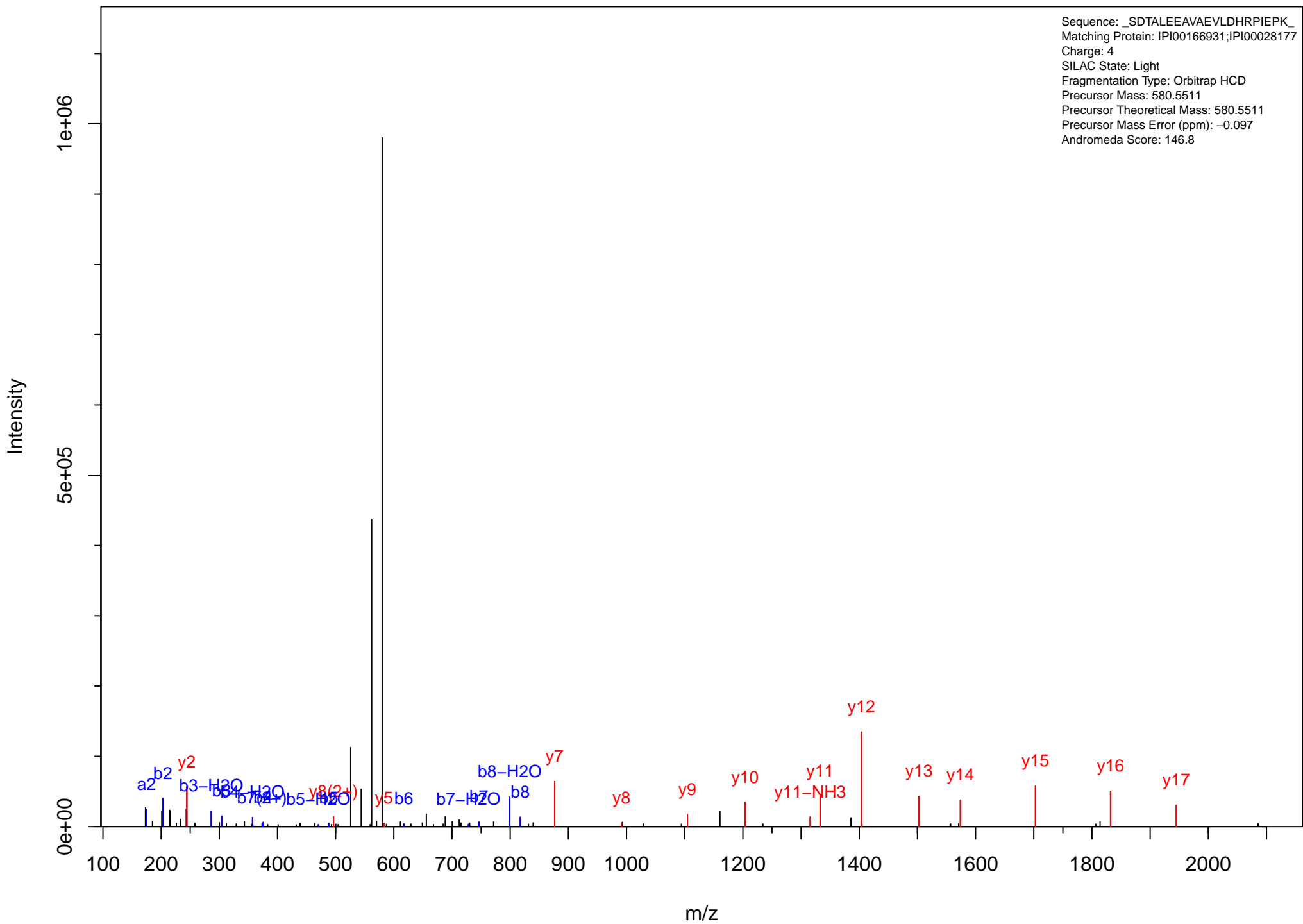
m/z

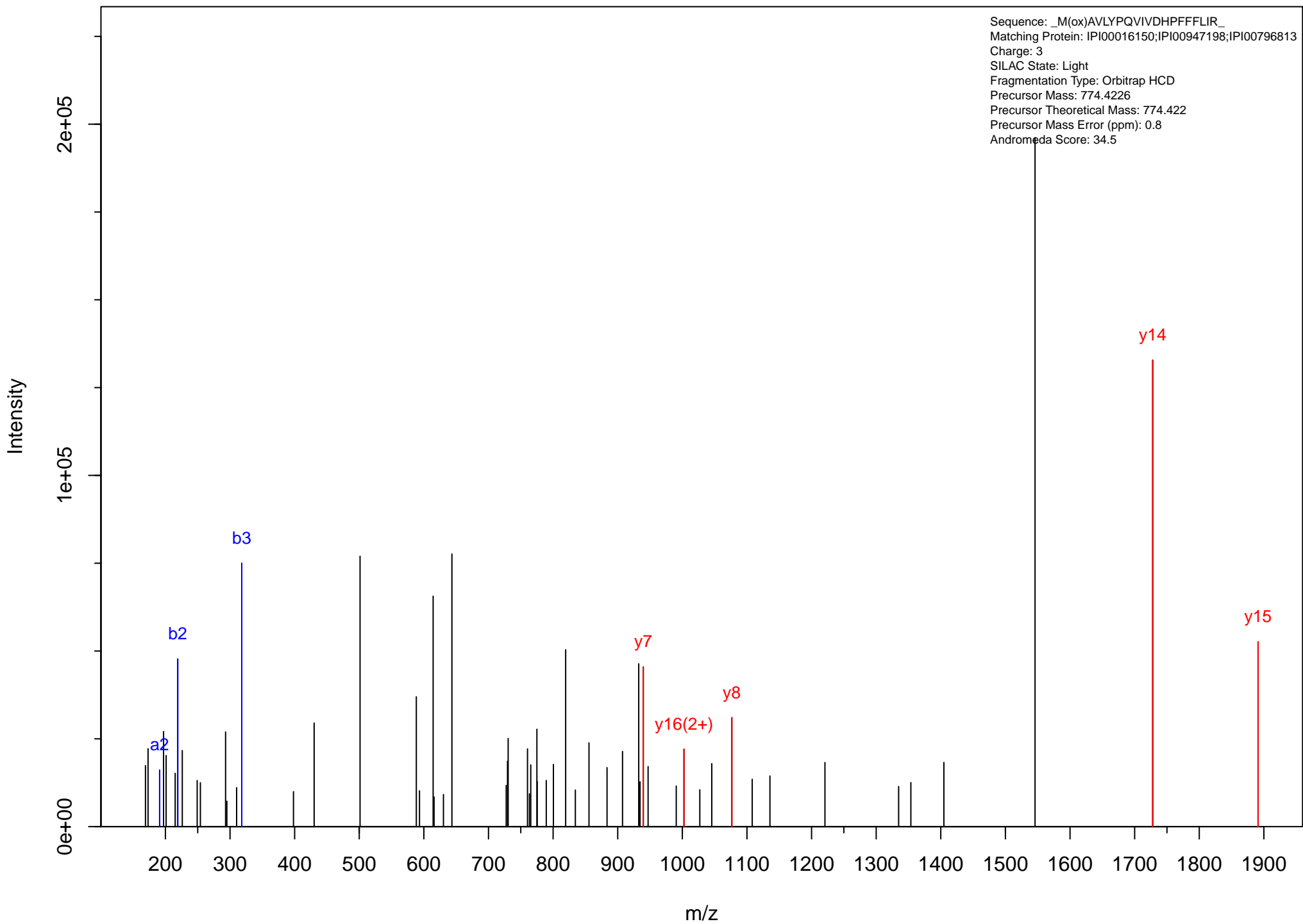


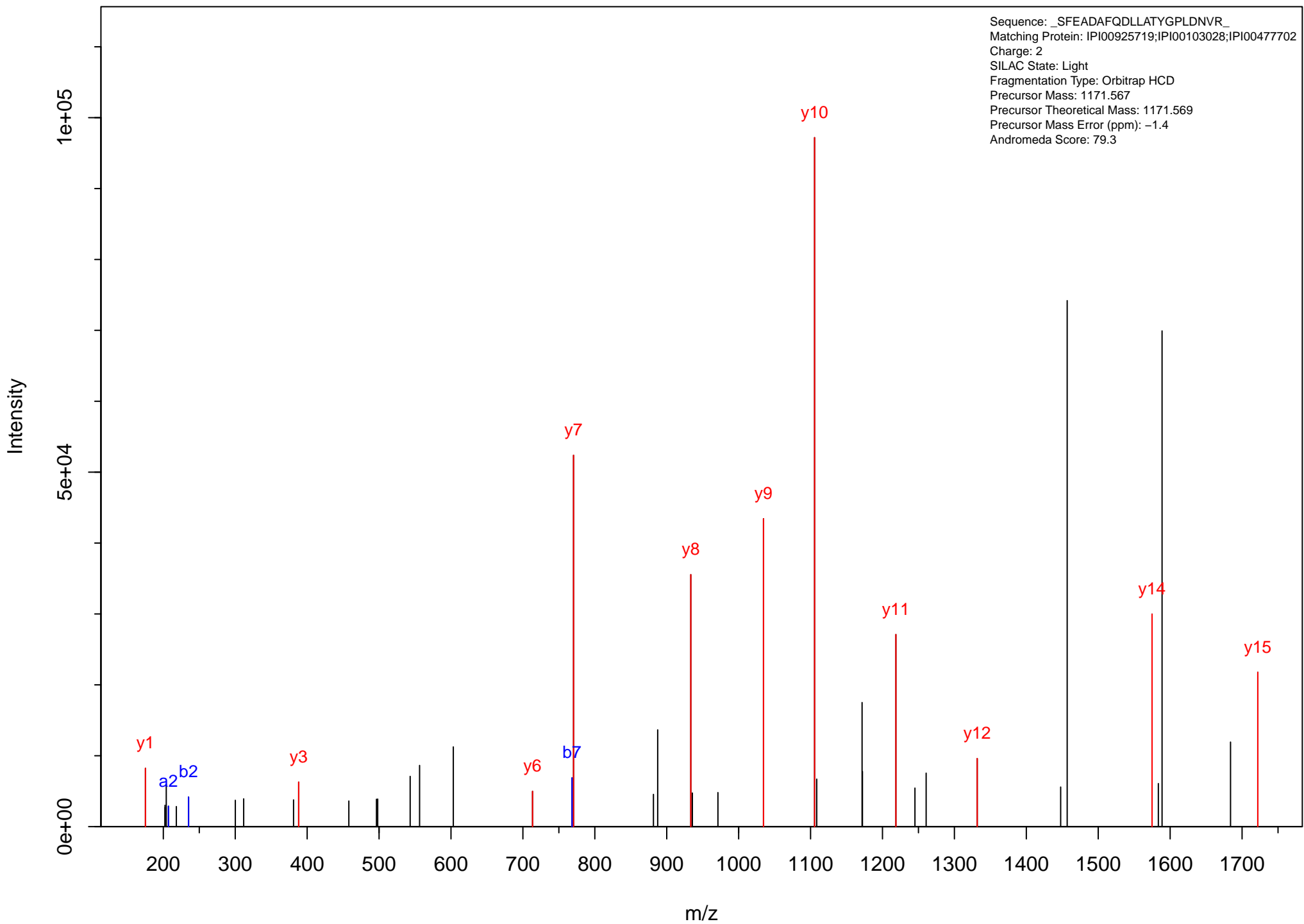


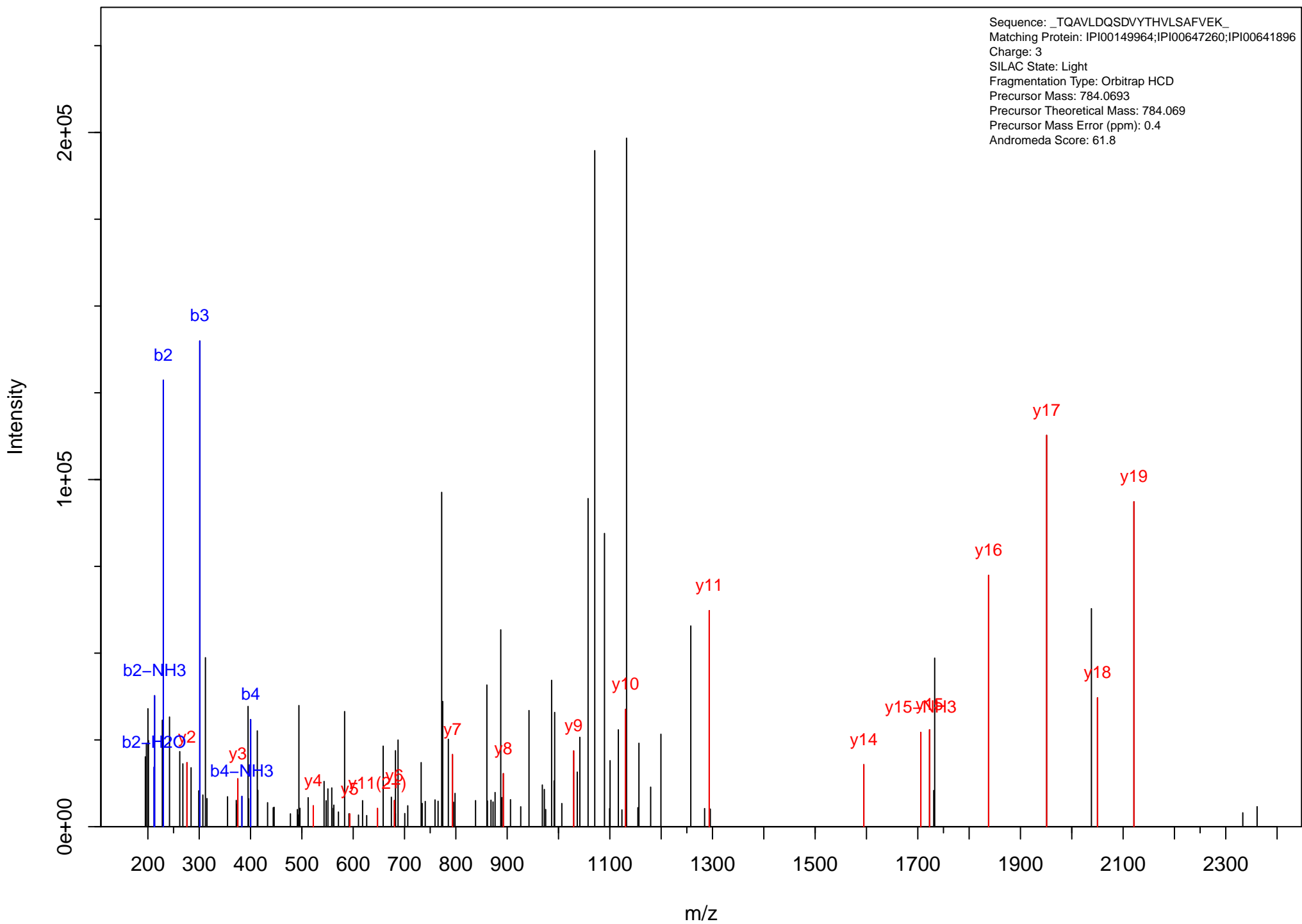


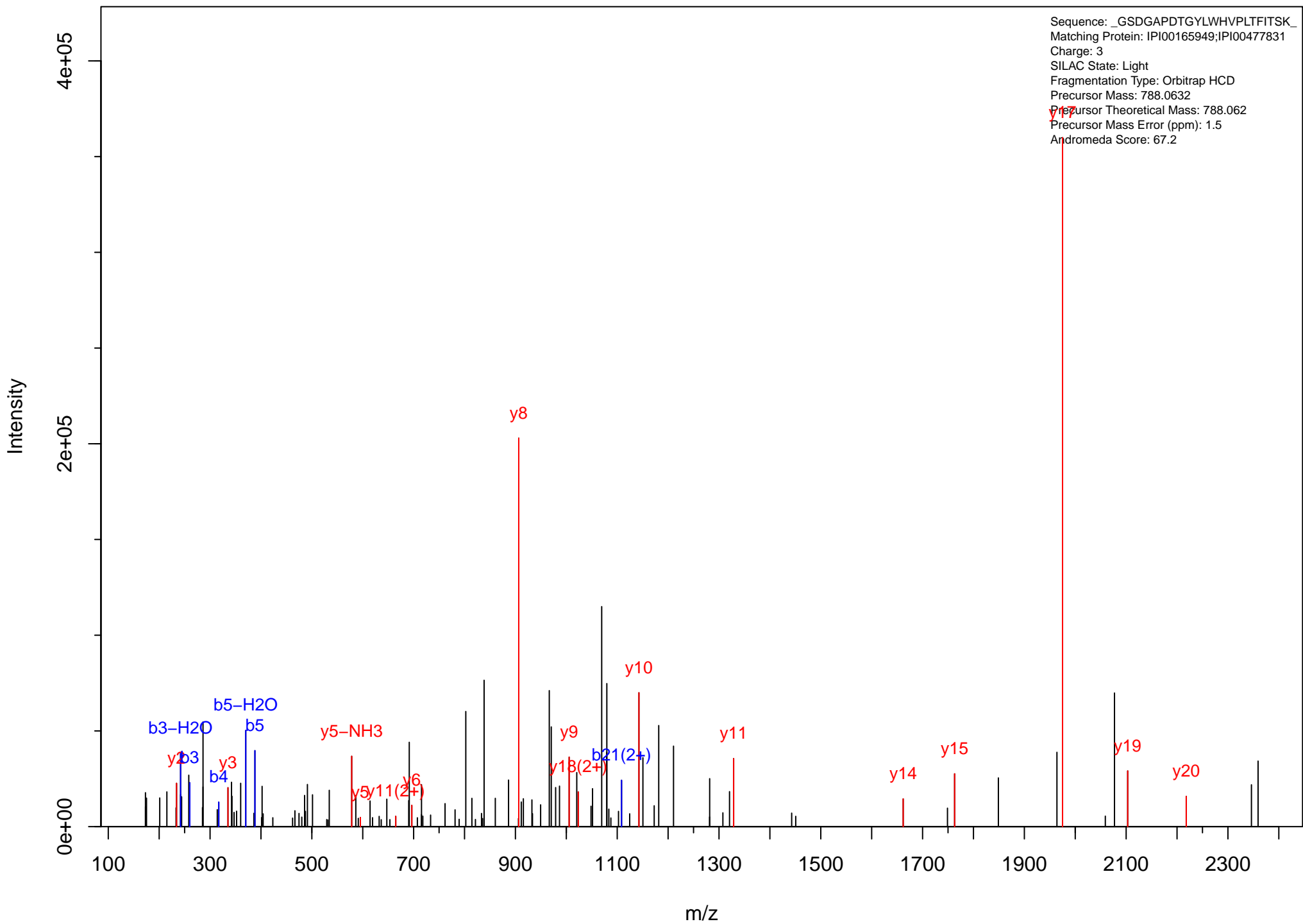


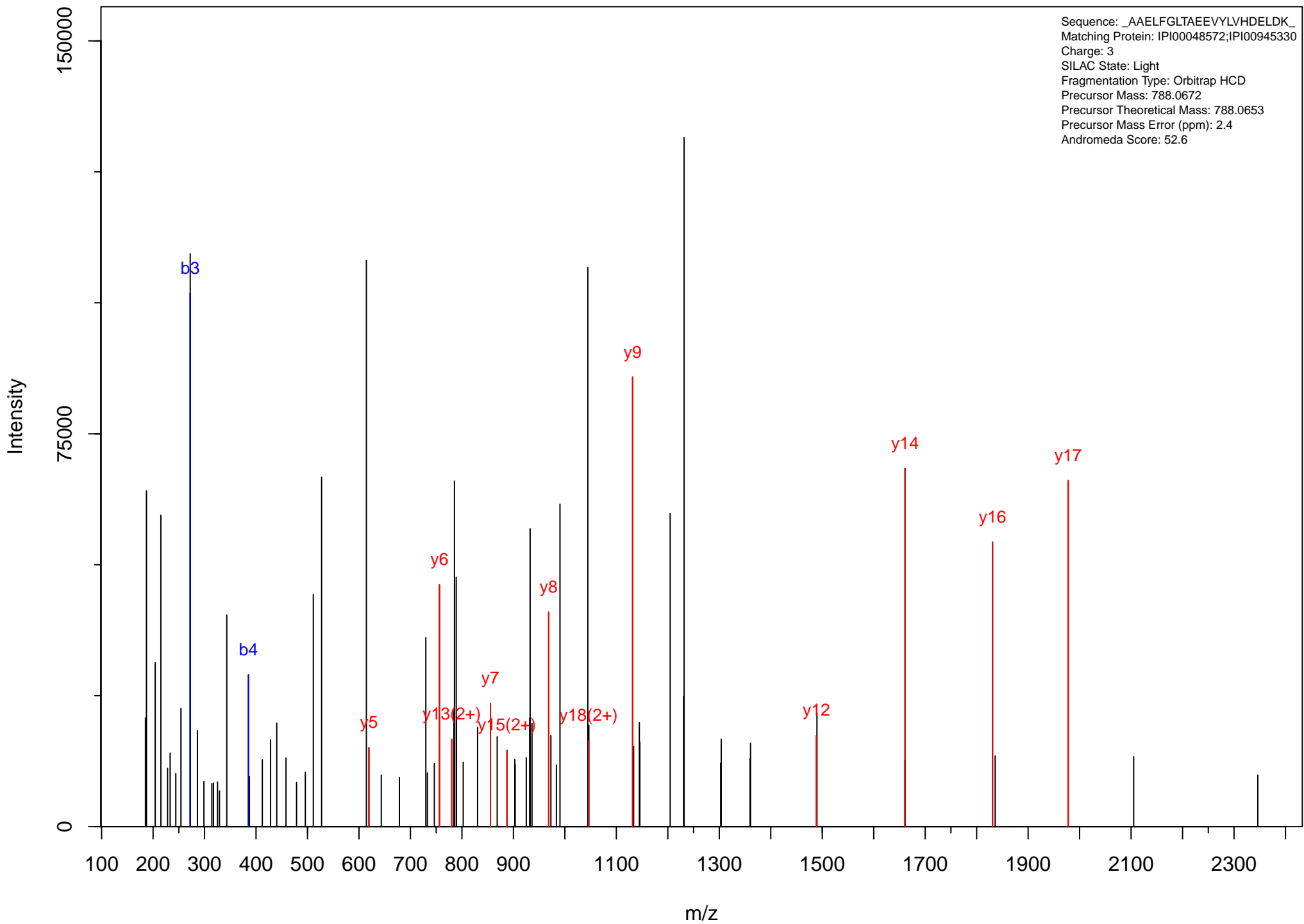


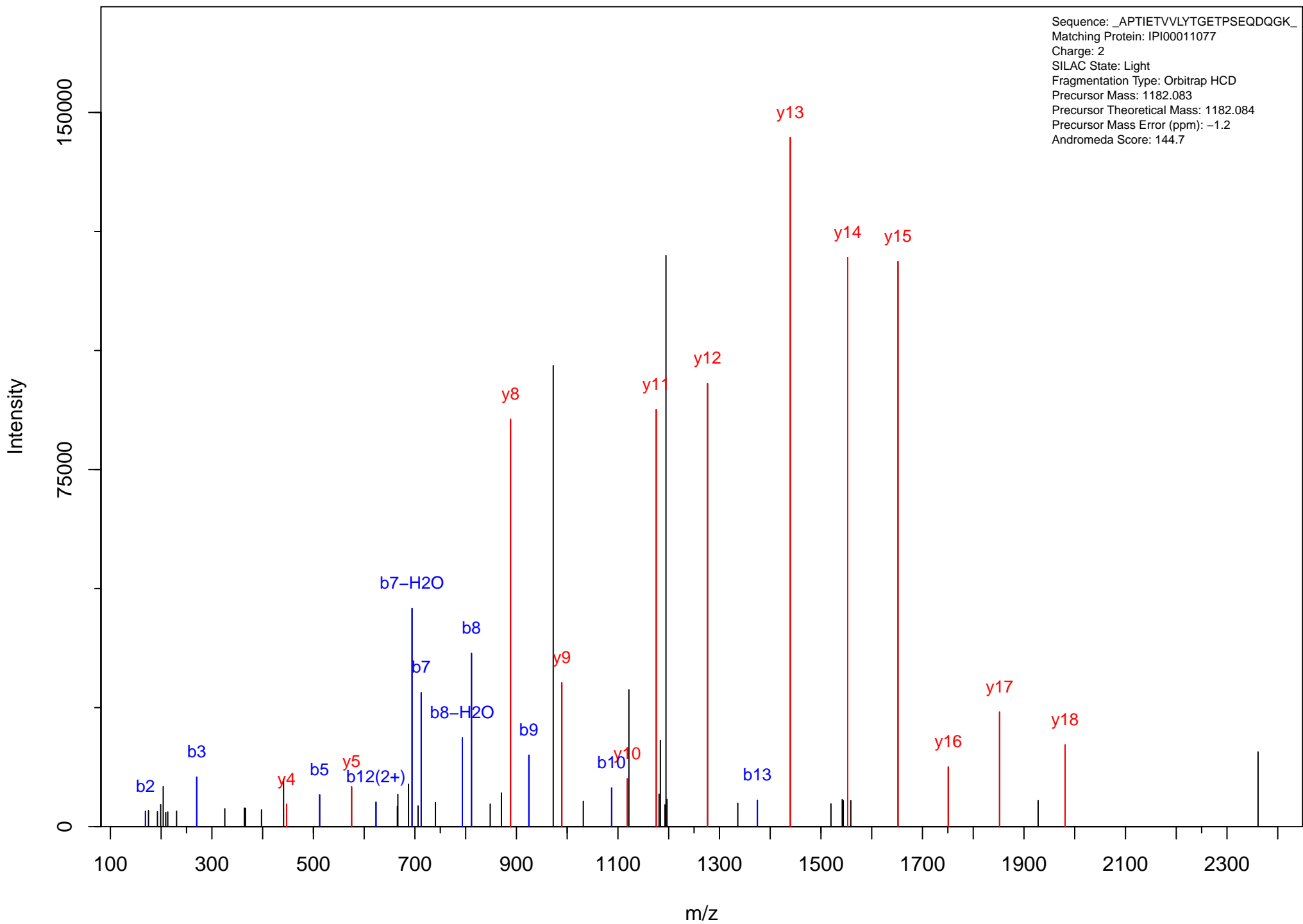


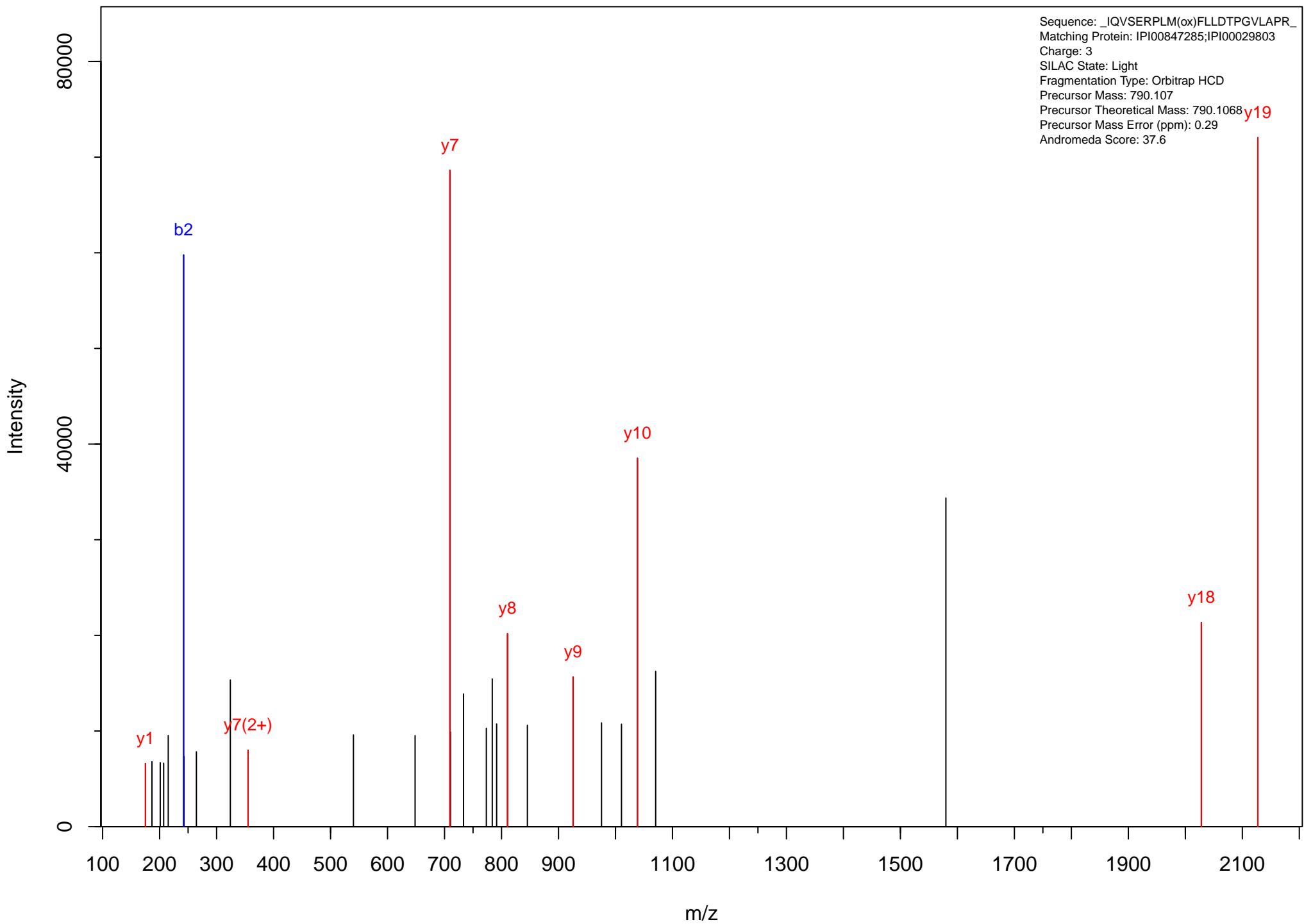


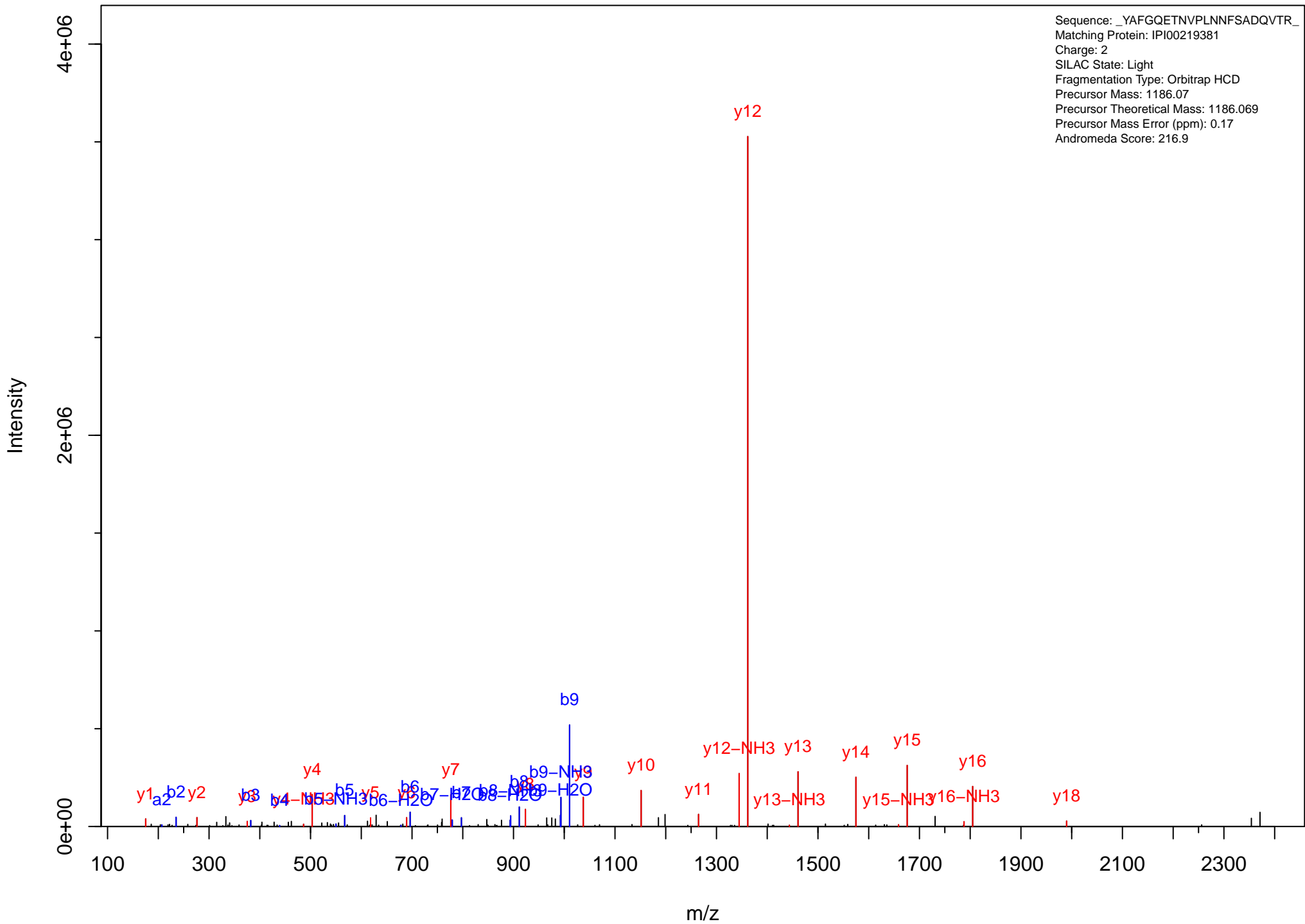


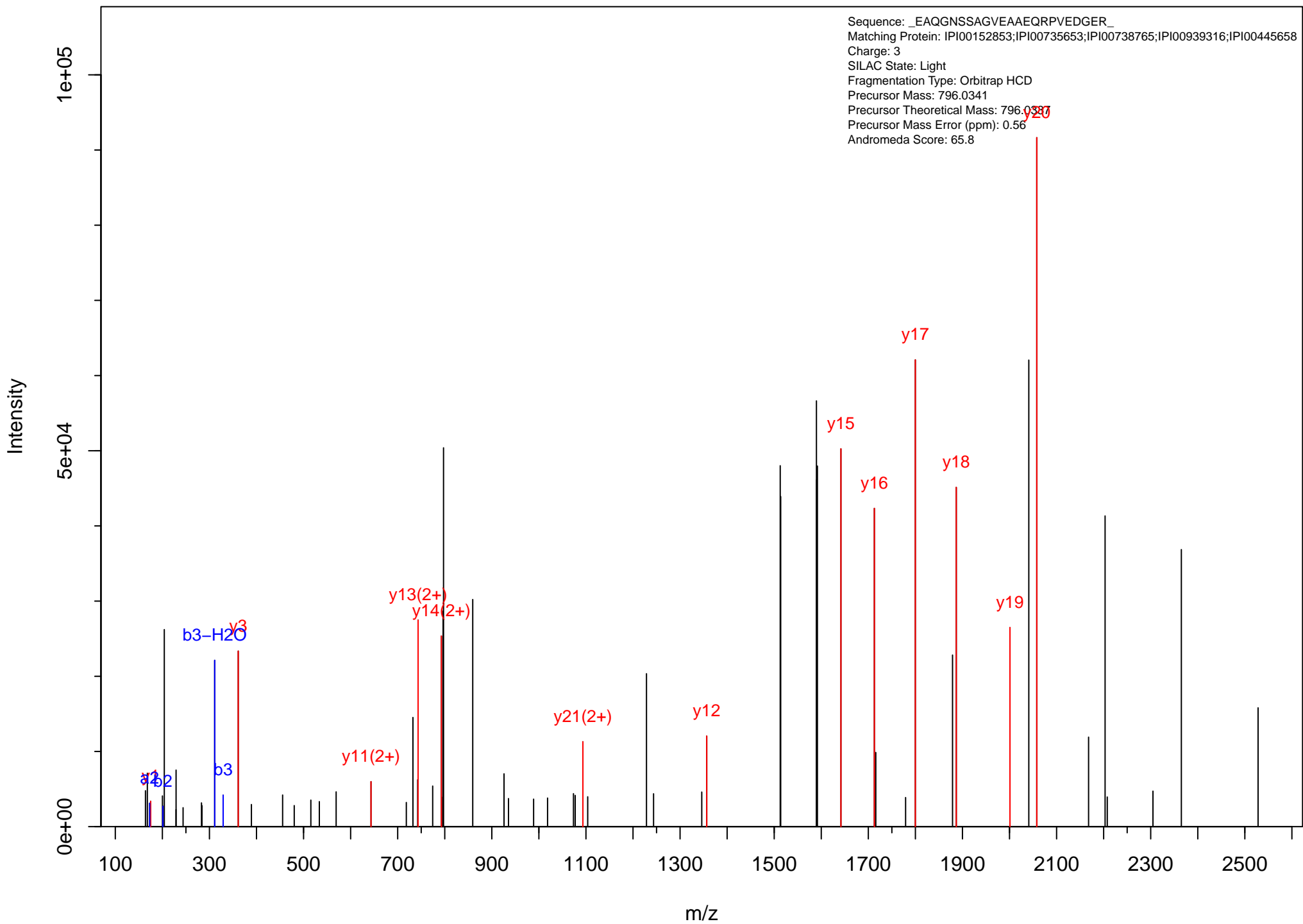


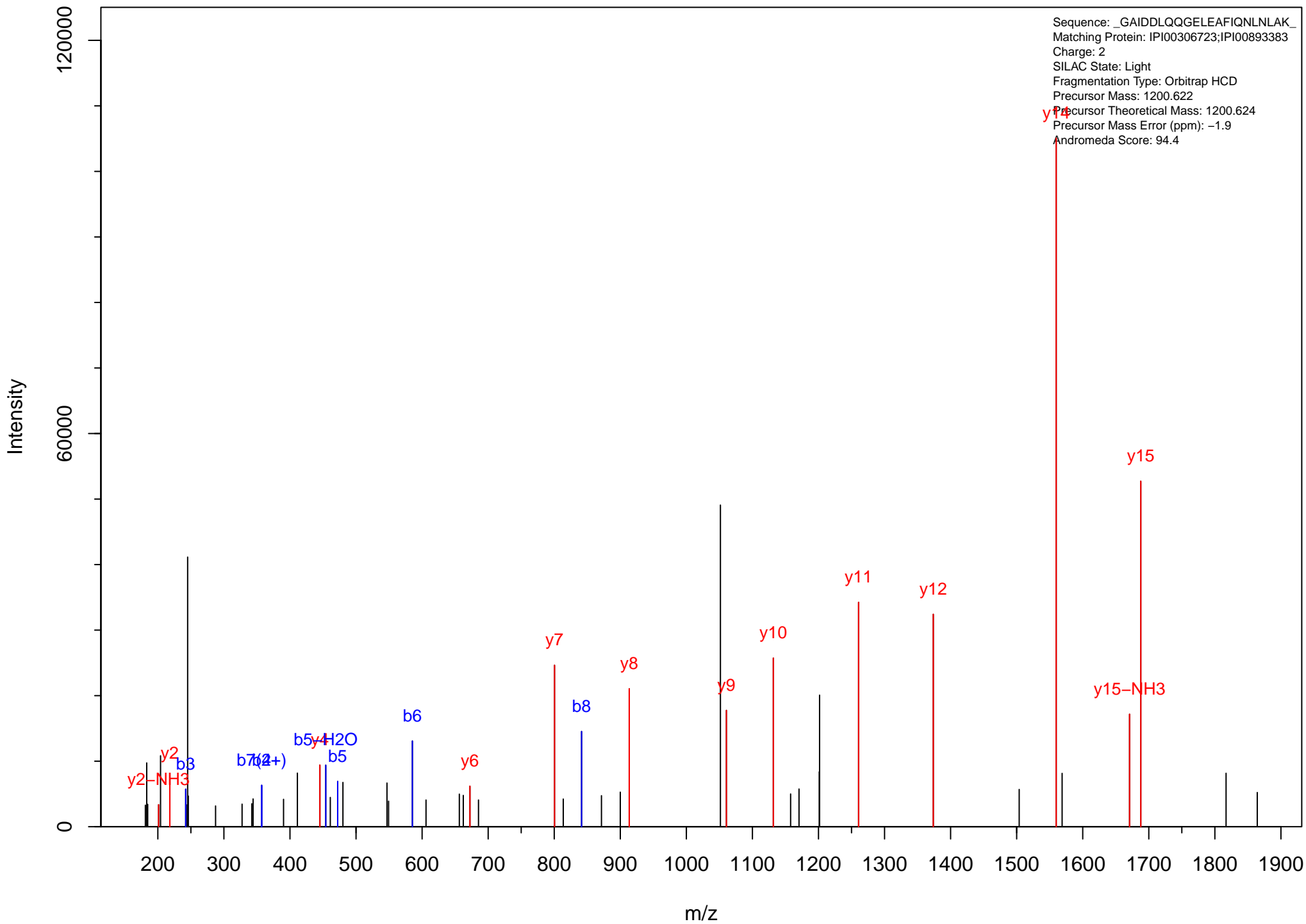


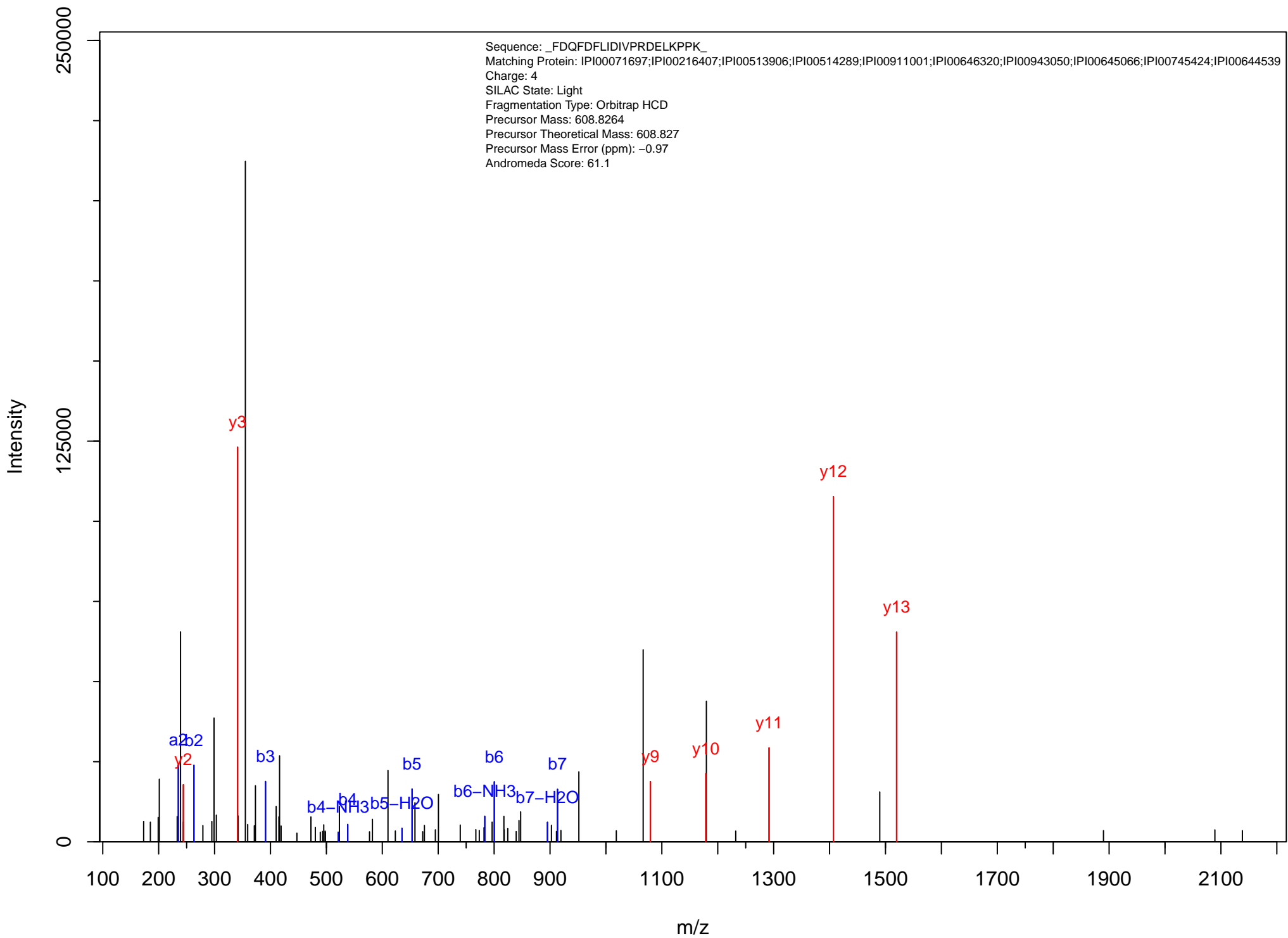


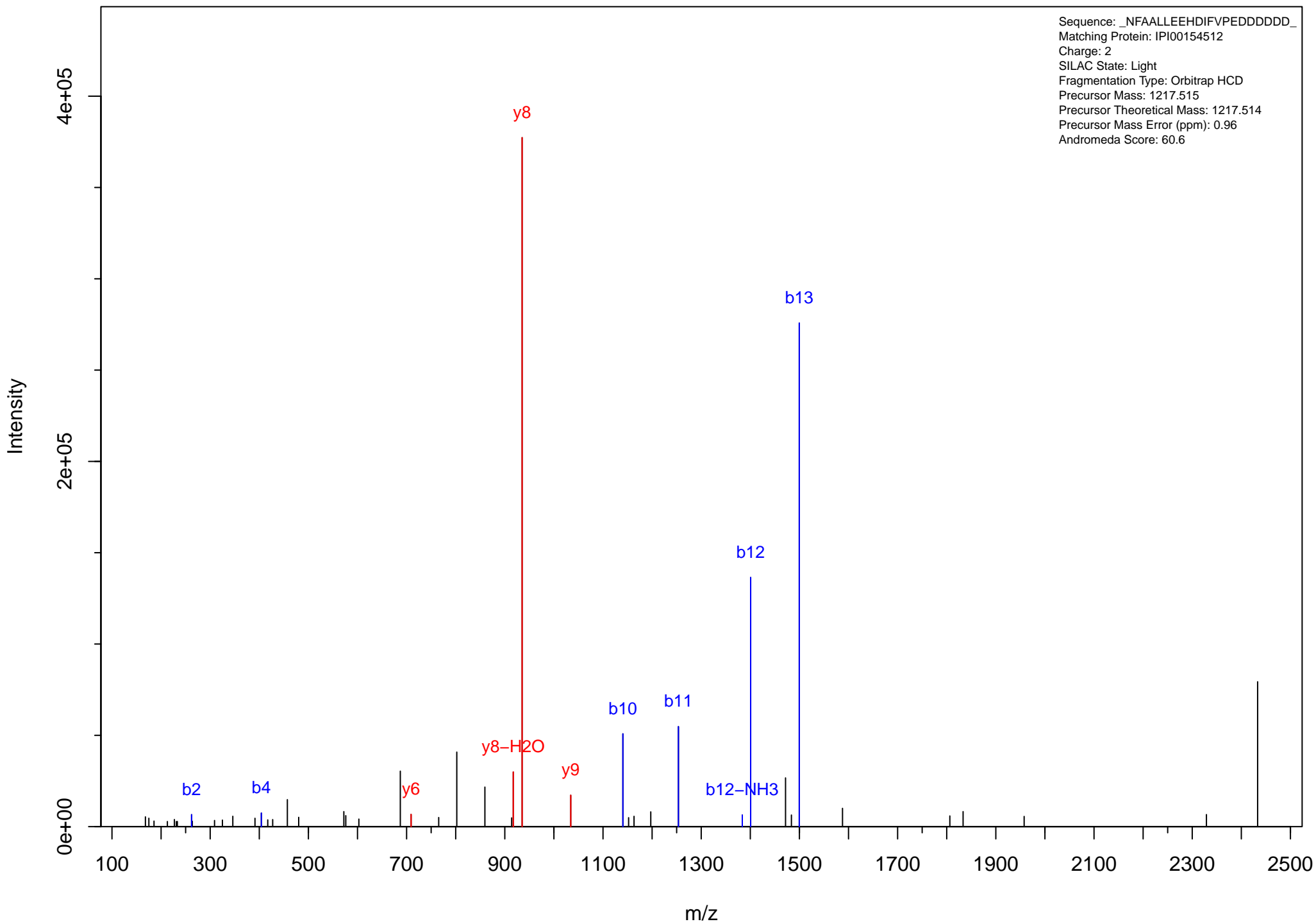


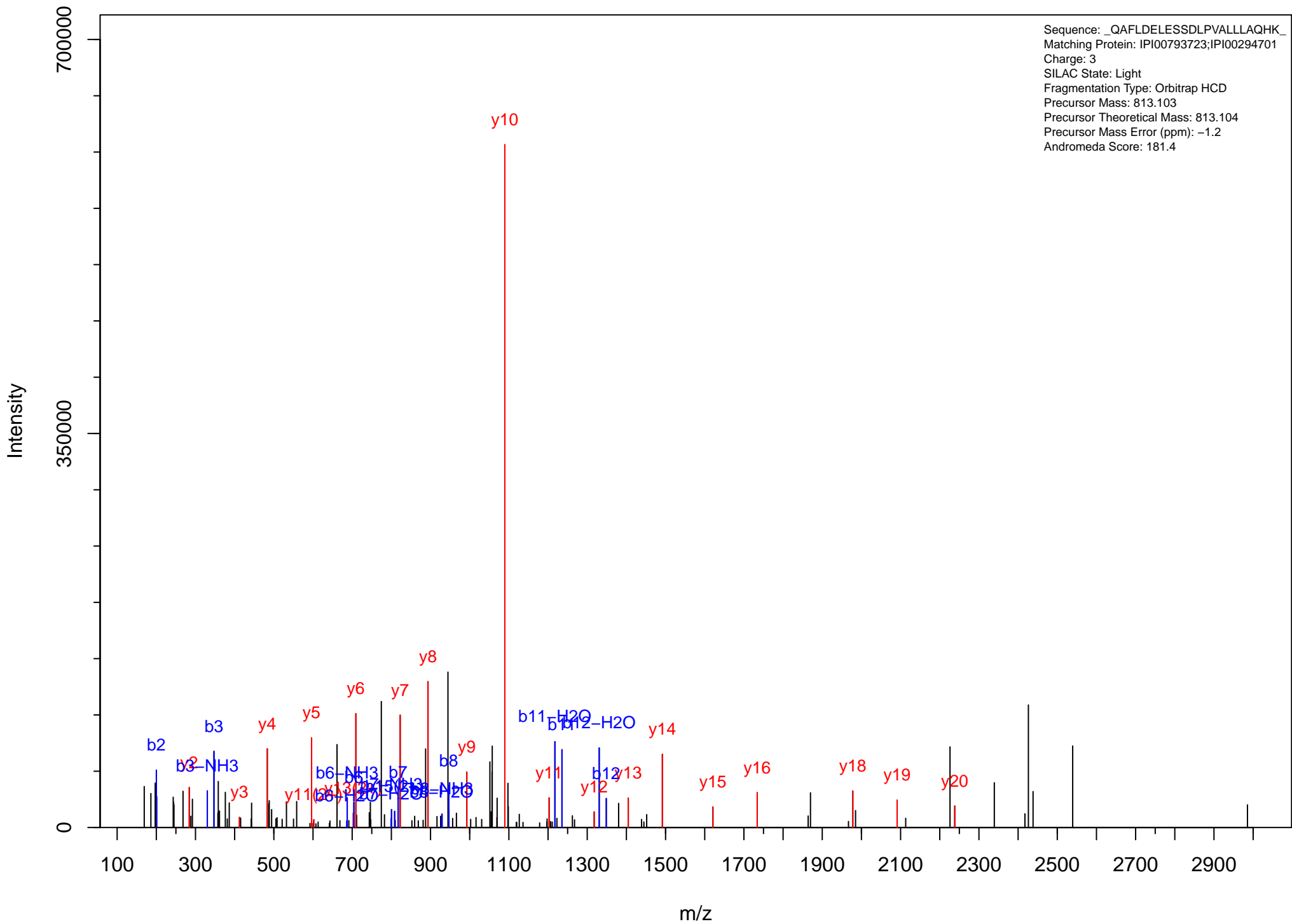


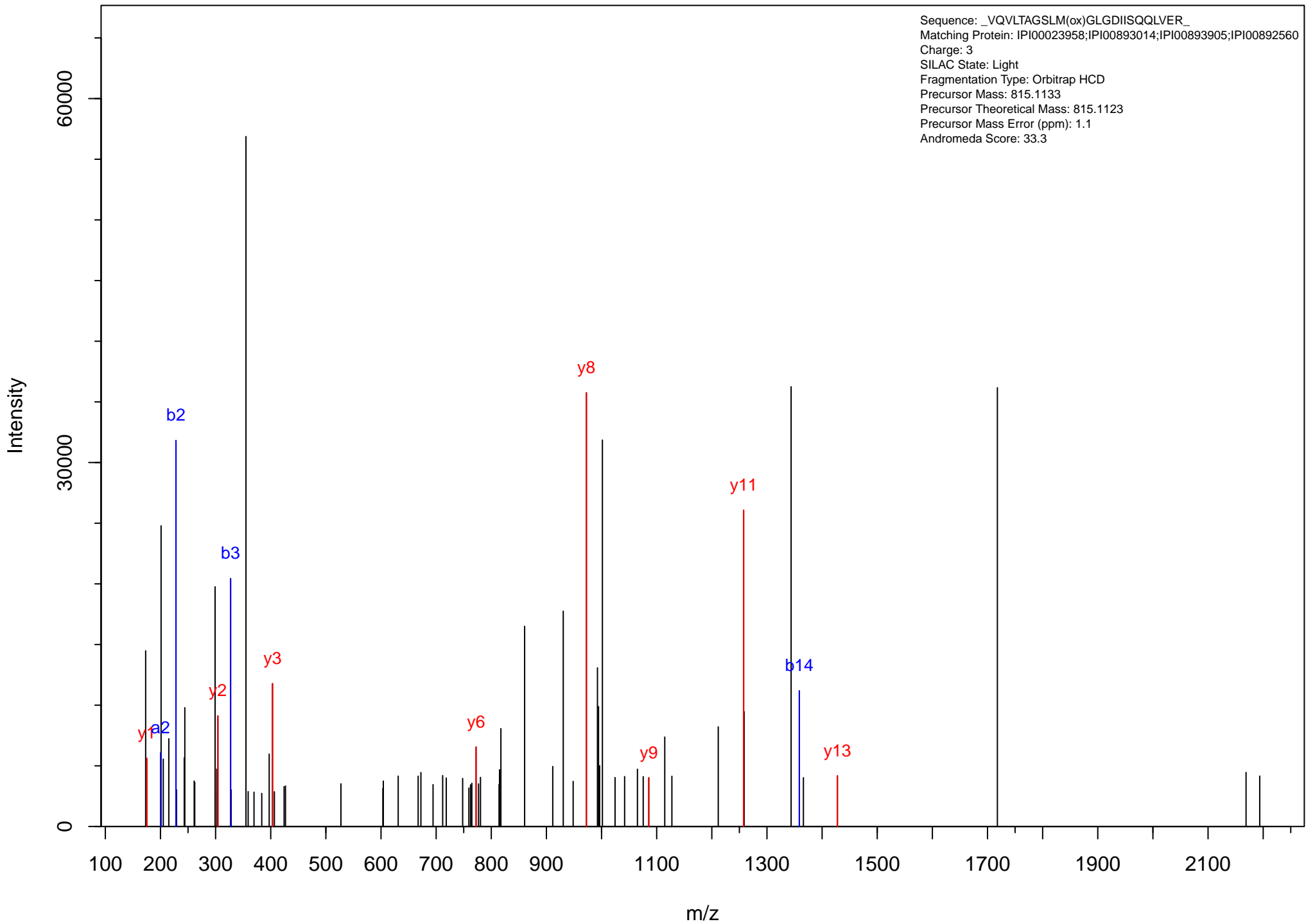


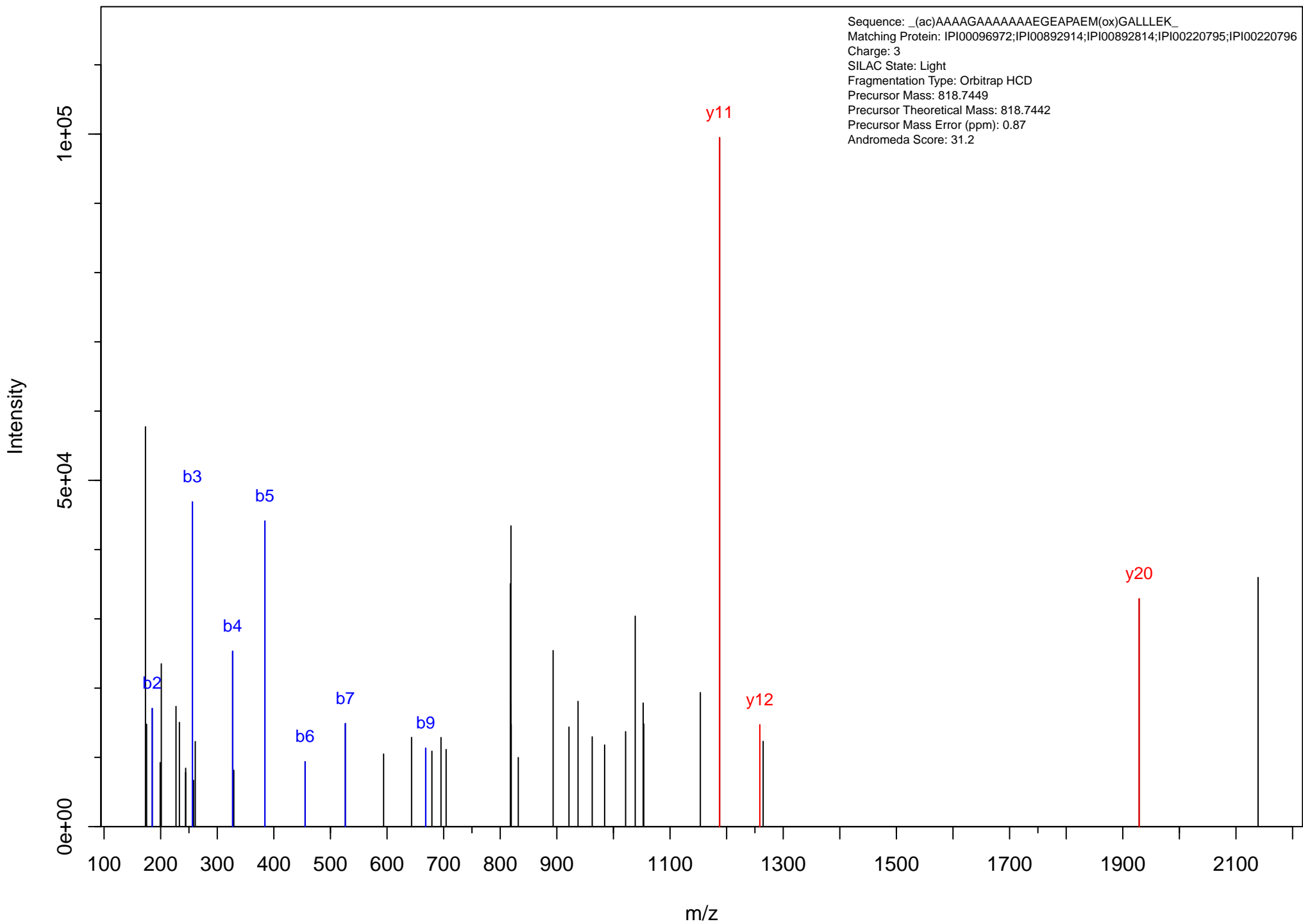


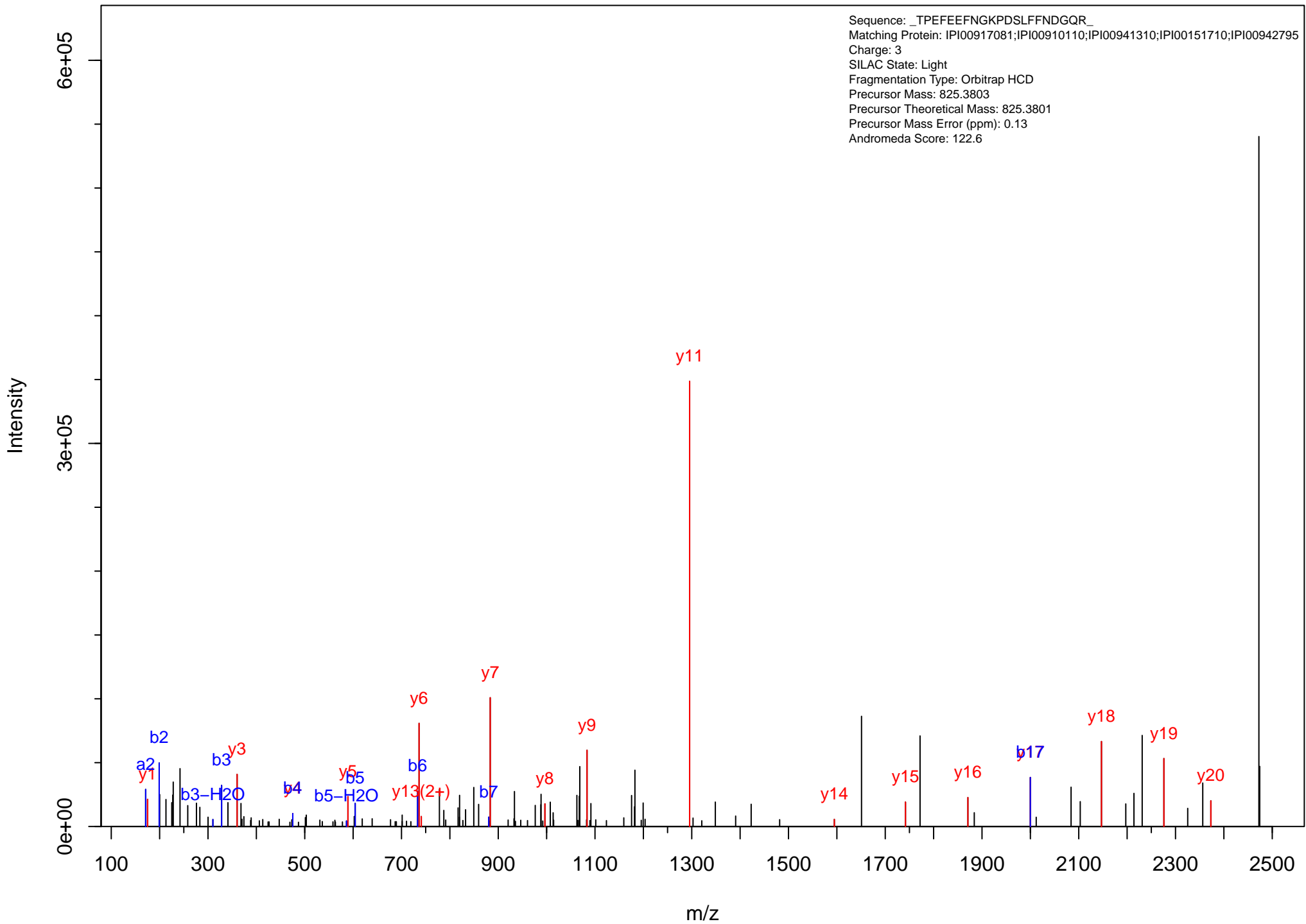


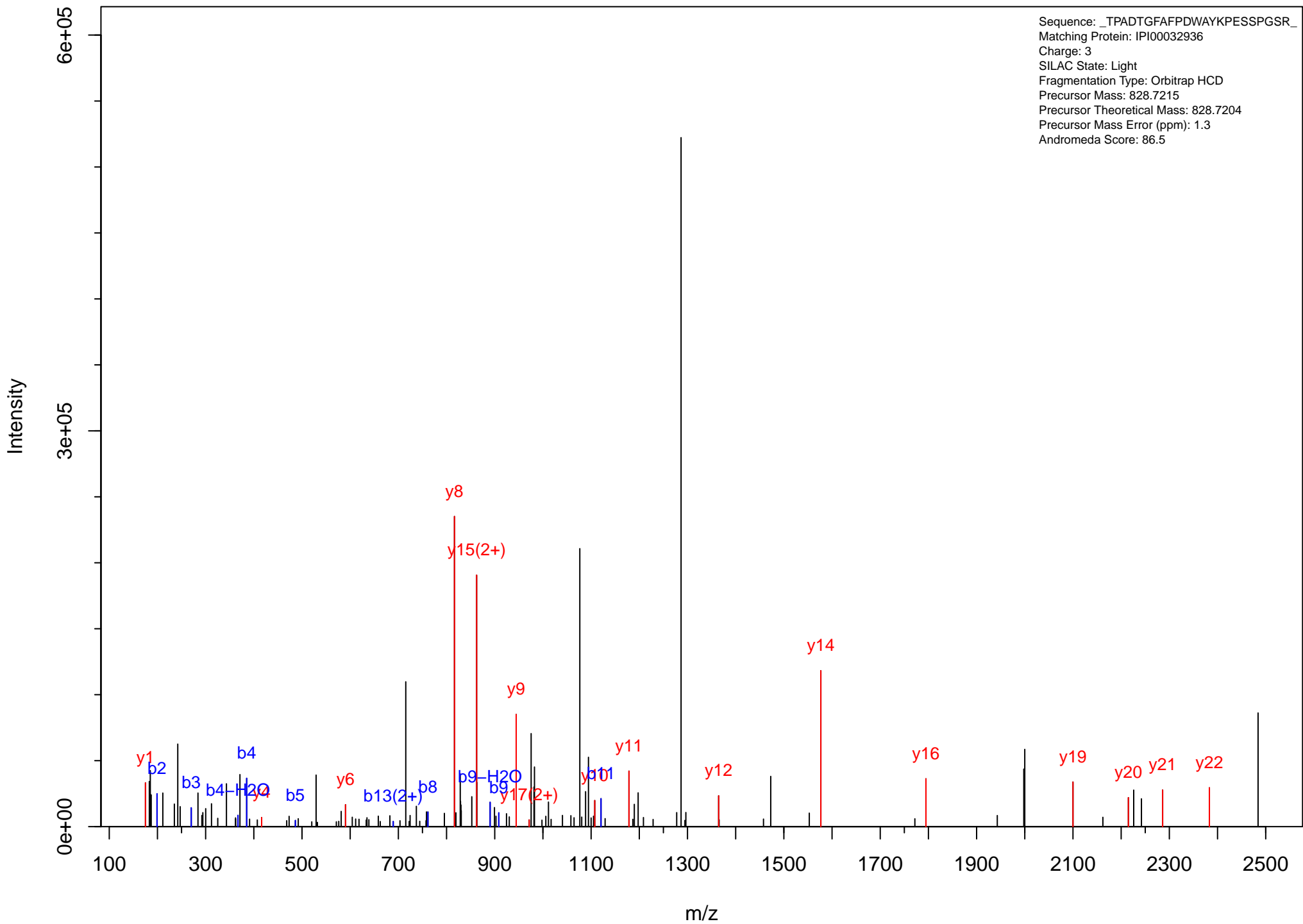


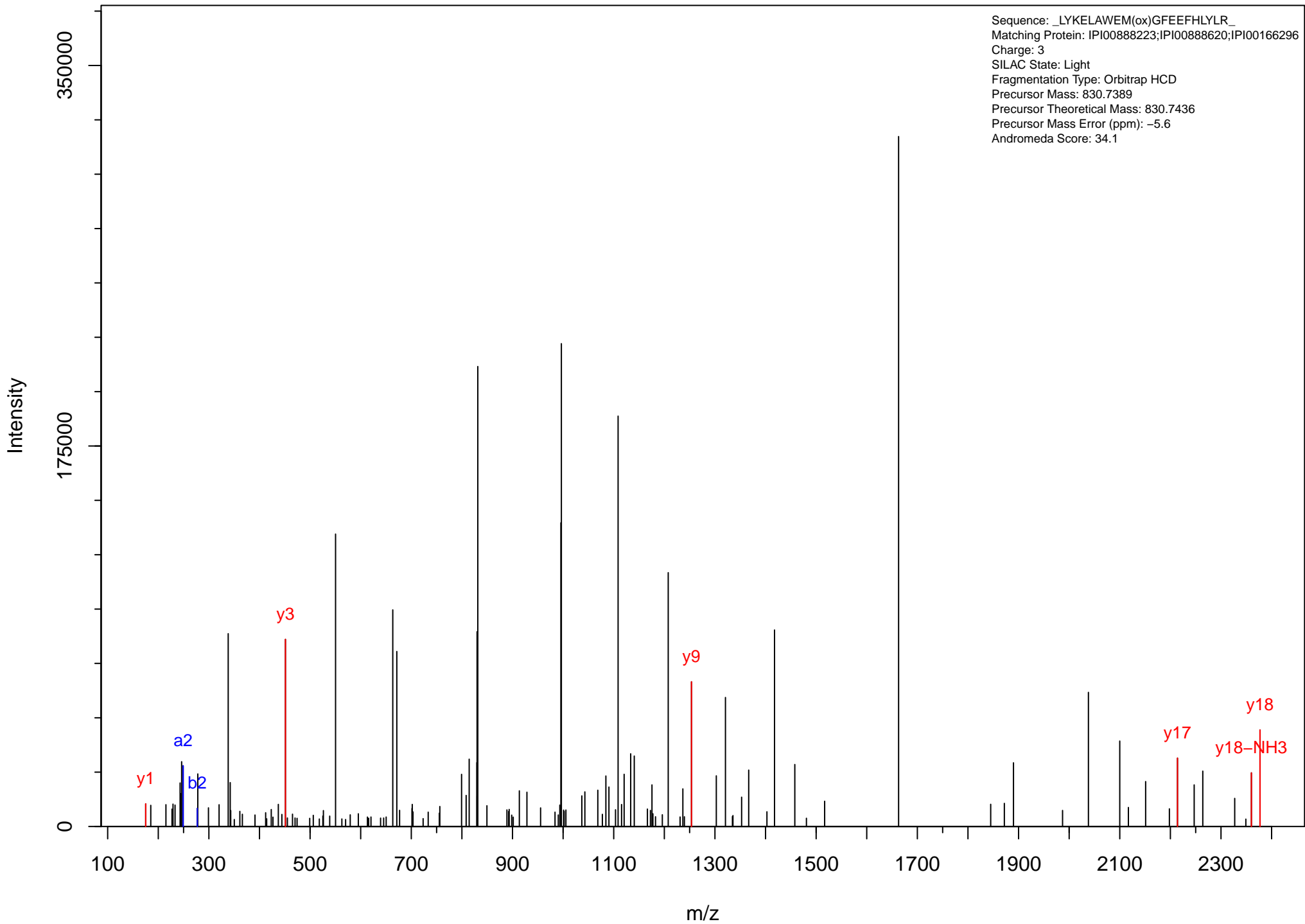


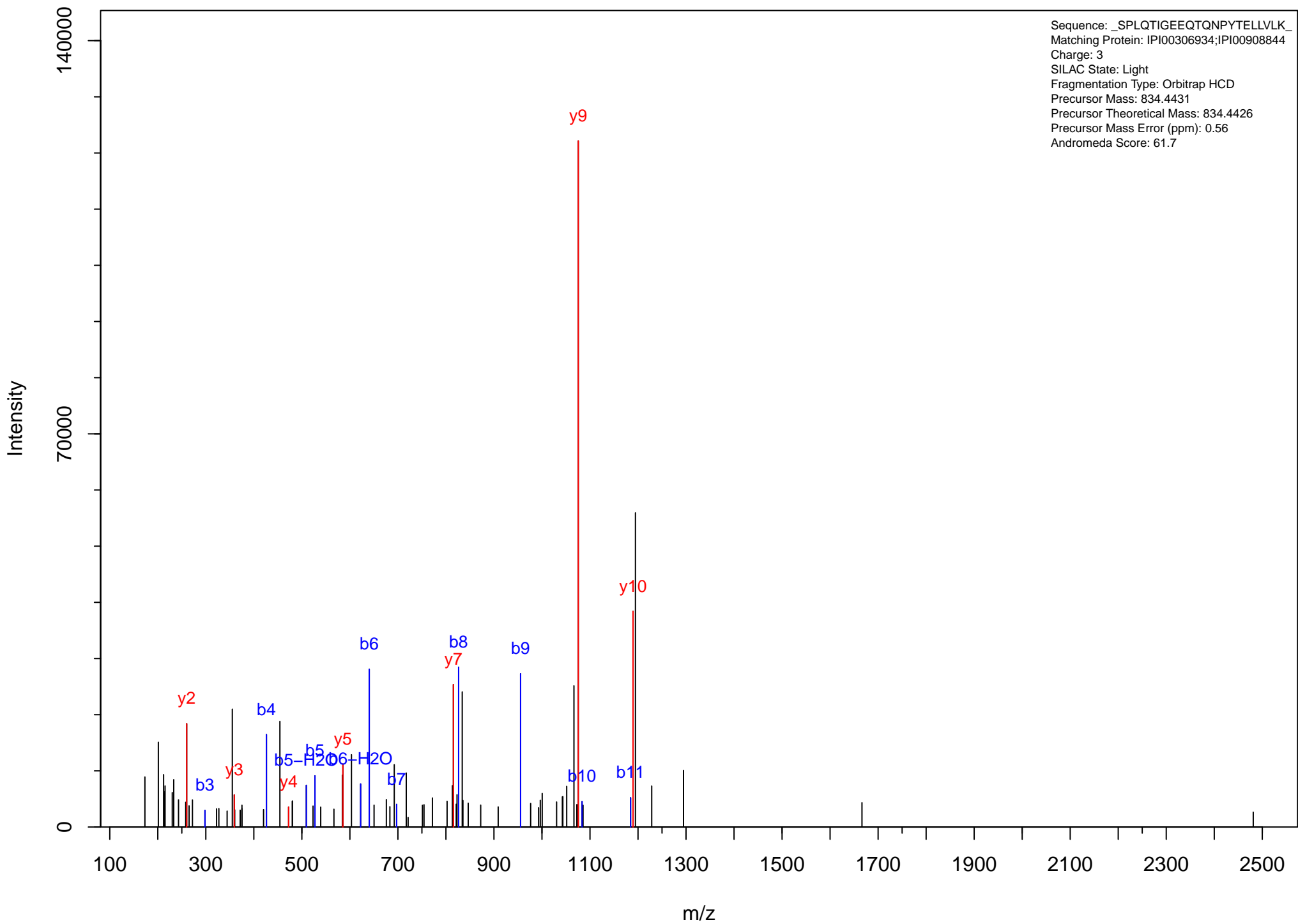


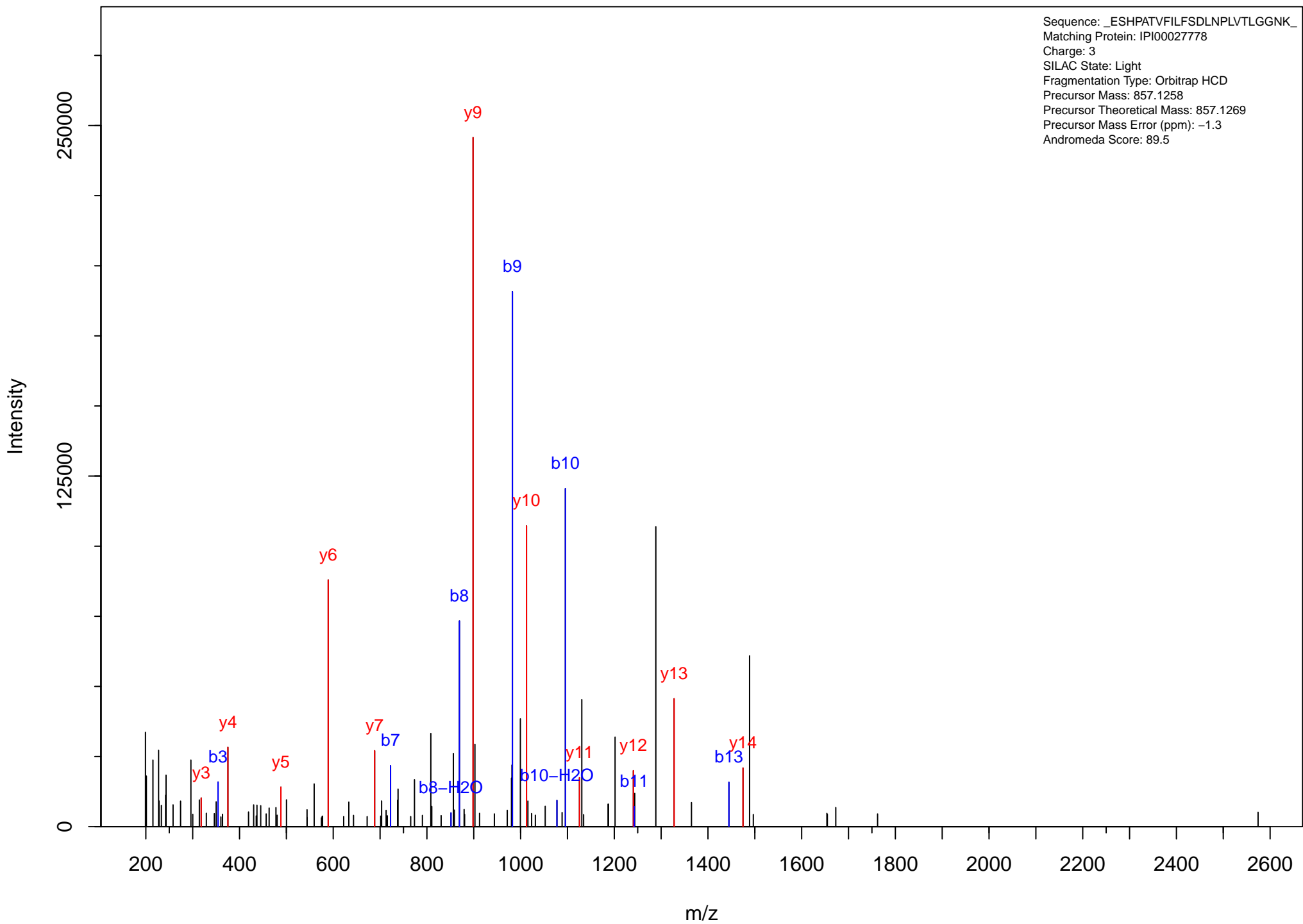


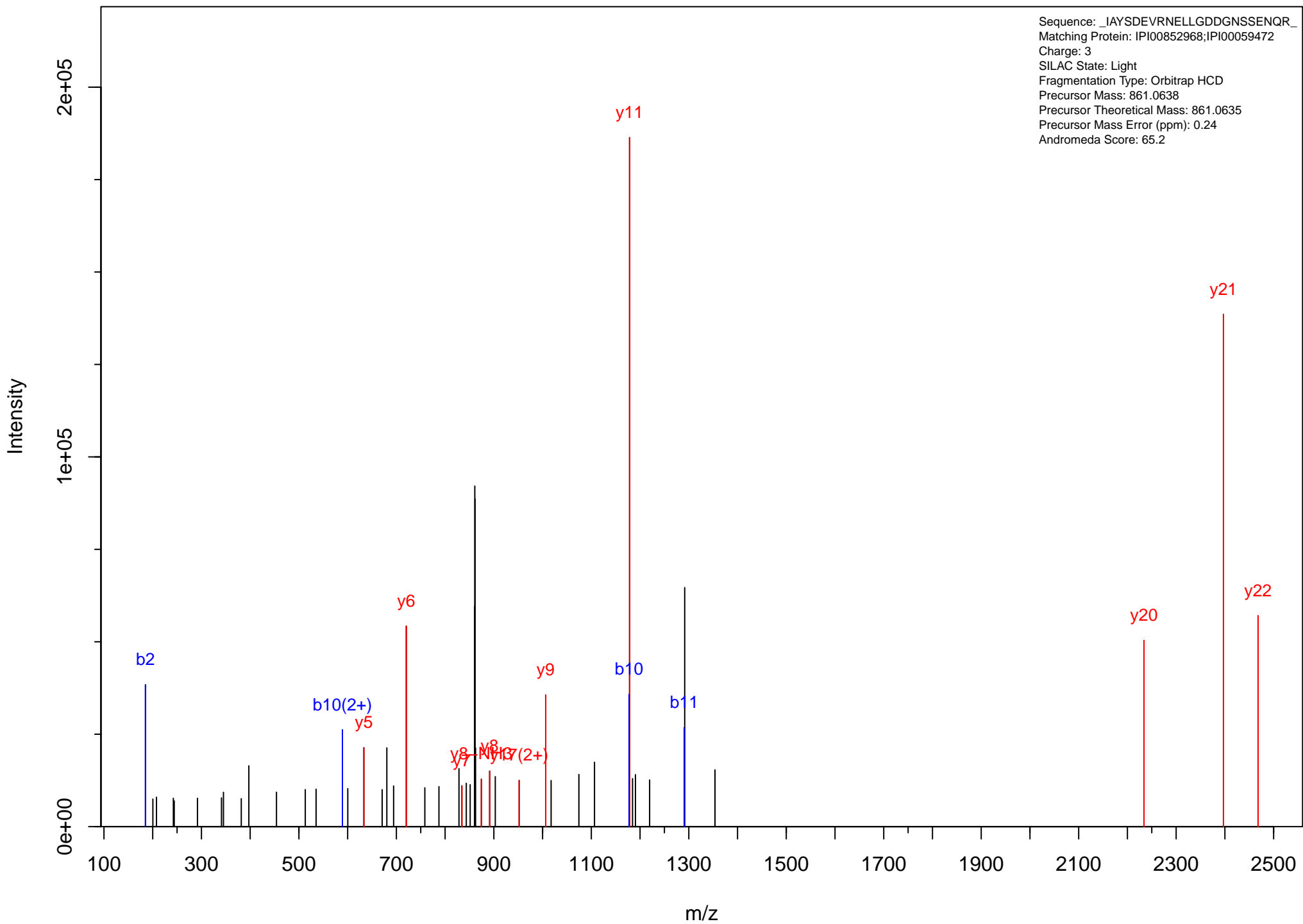


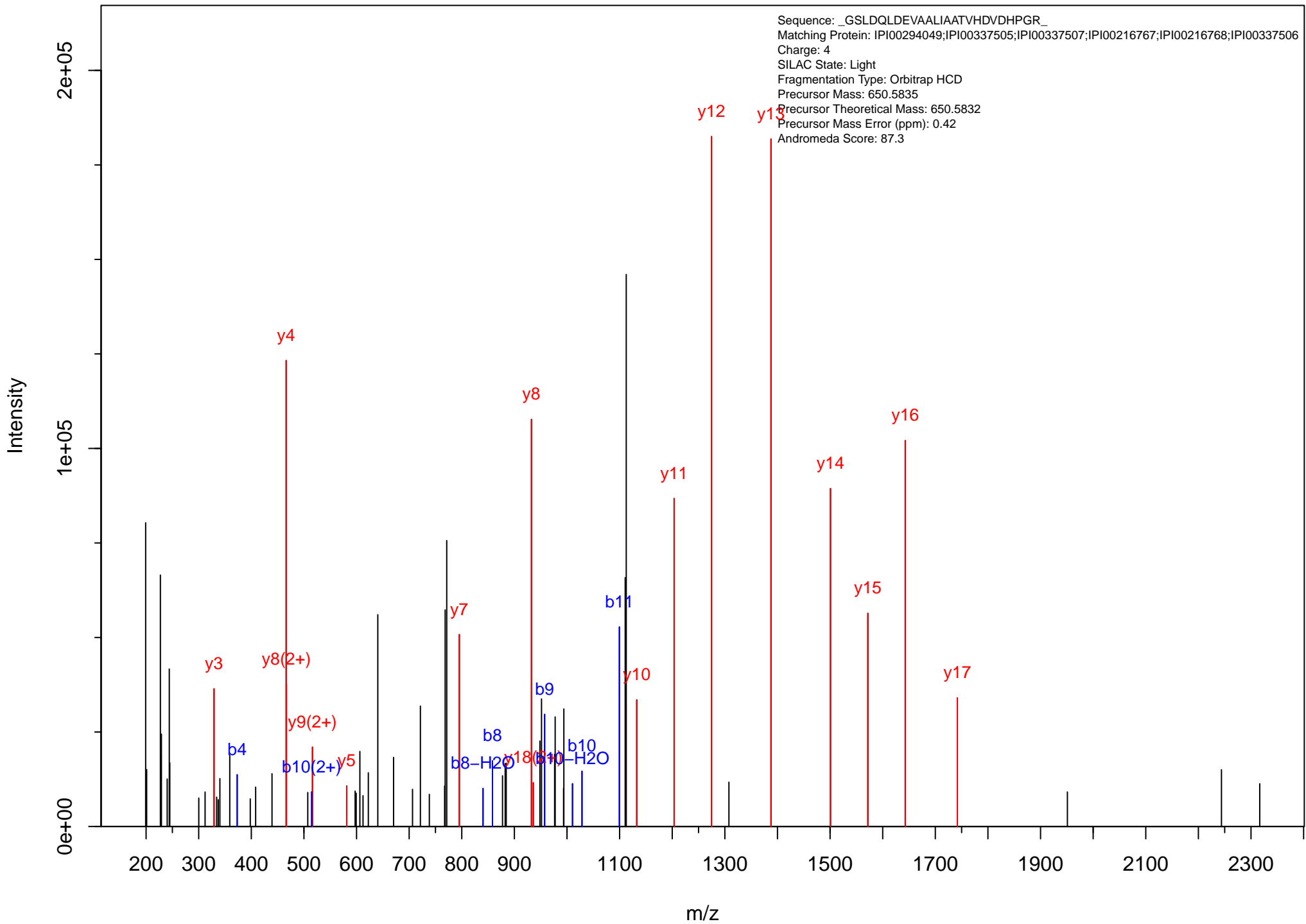


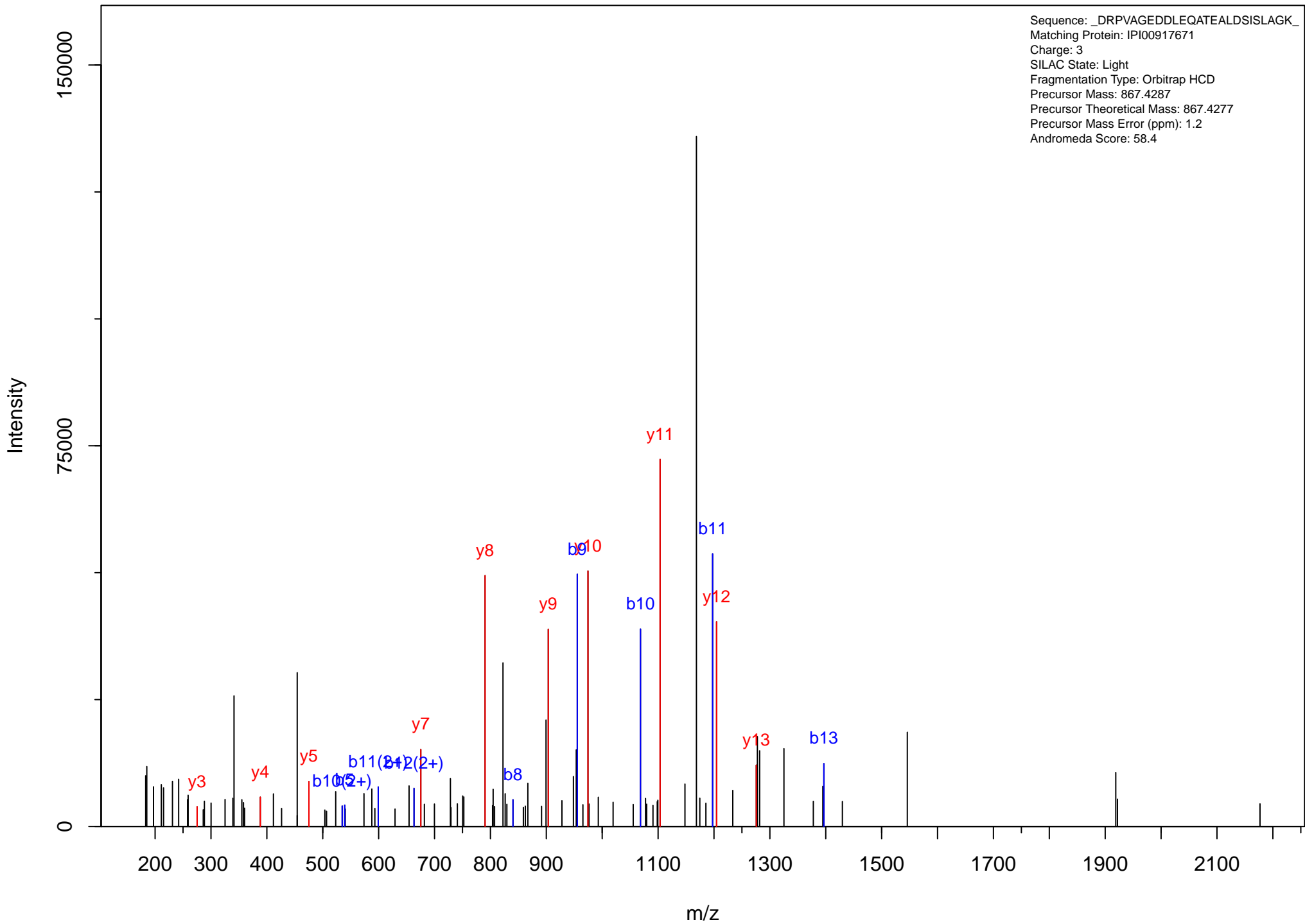


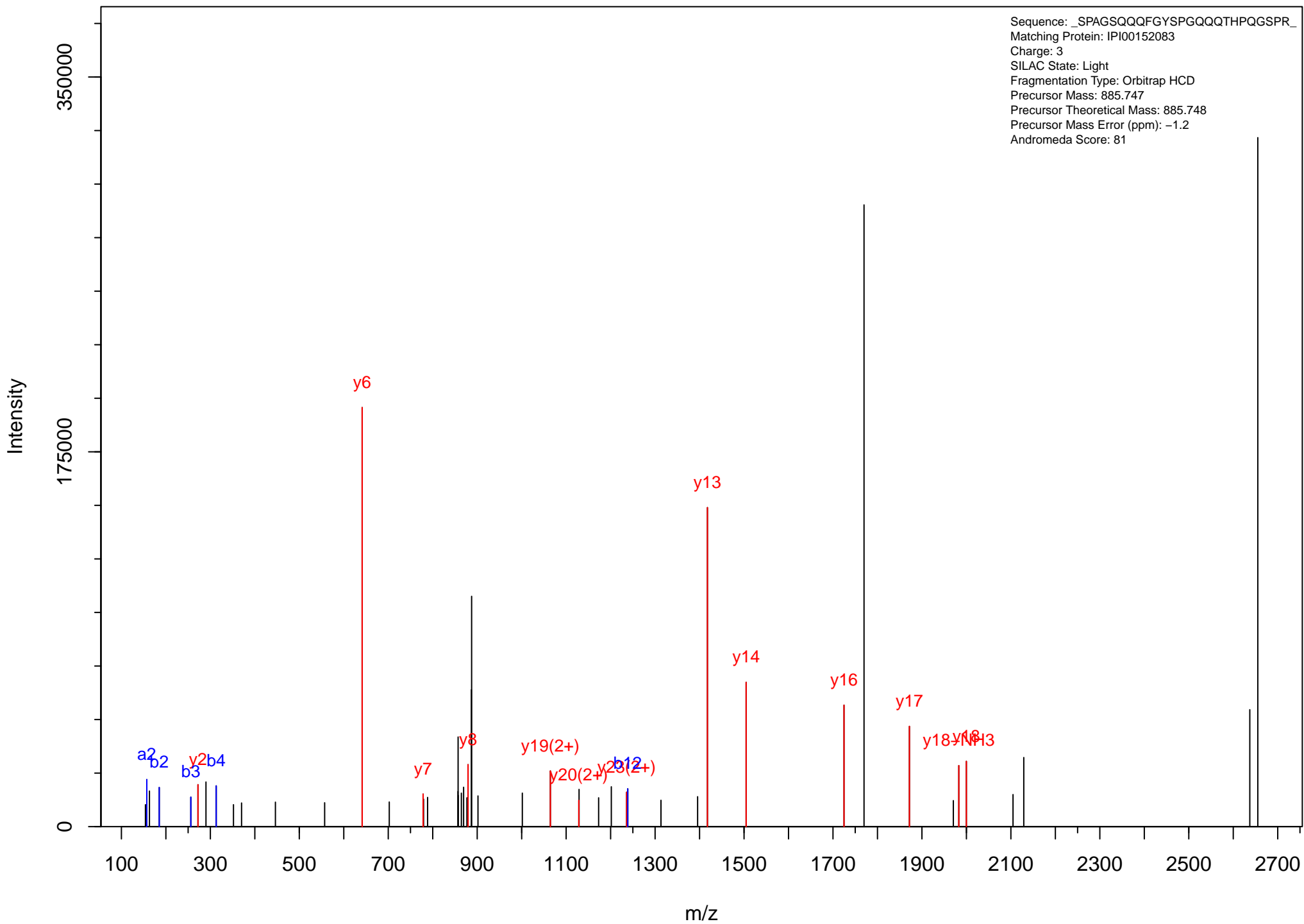


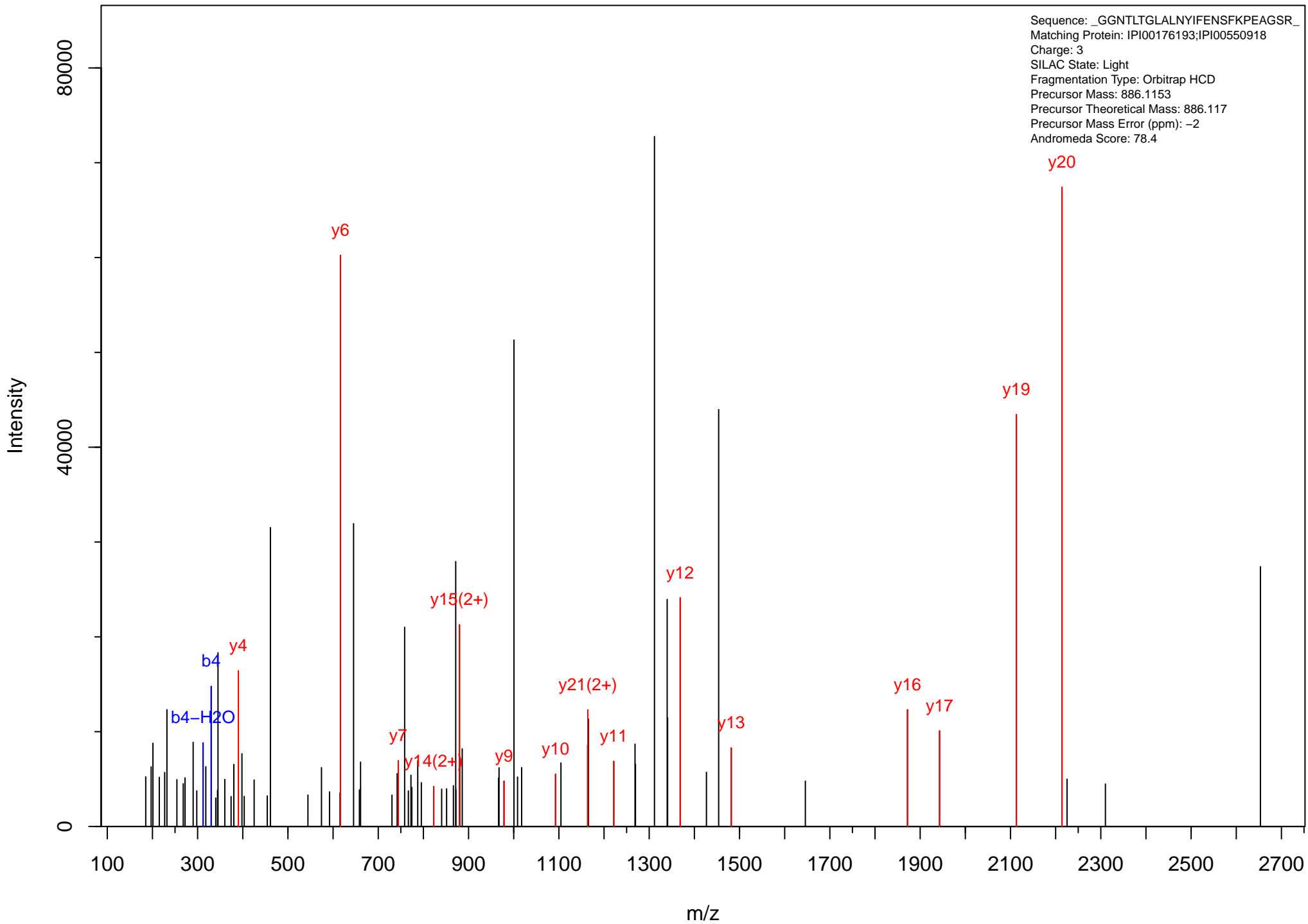


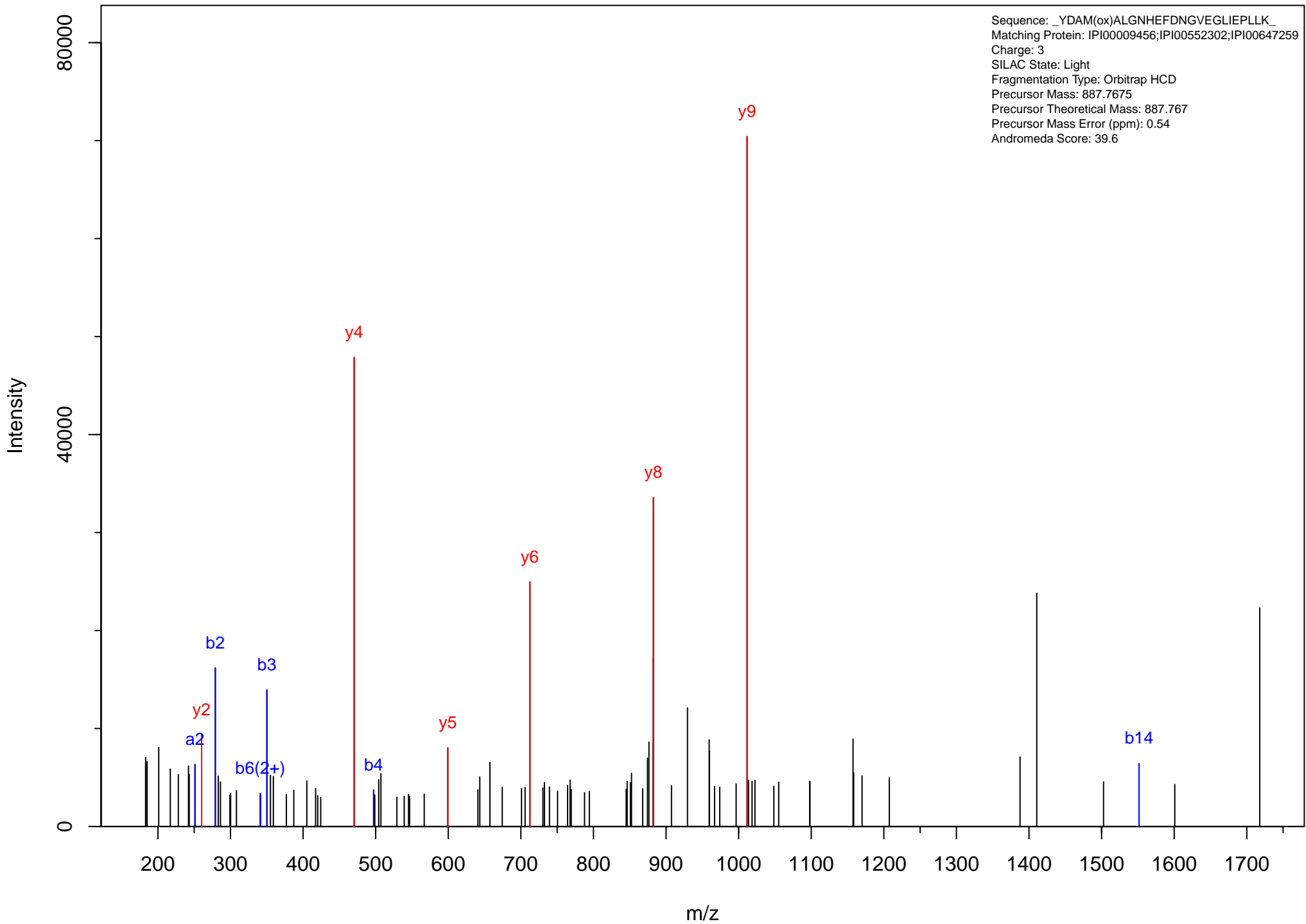


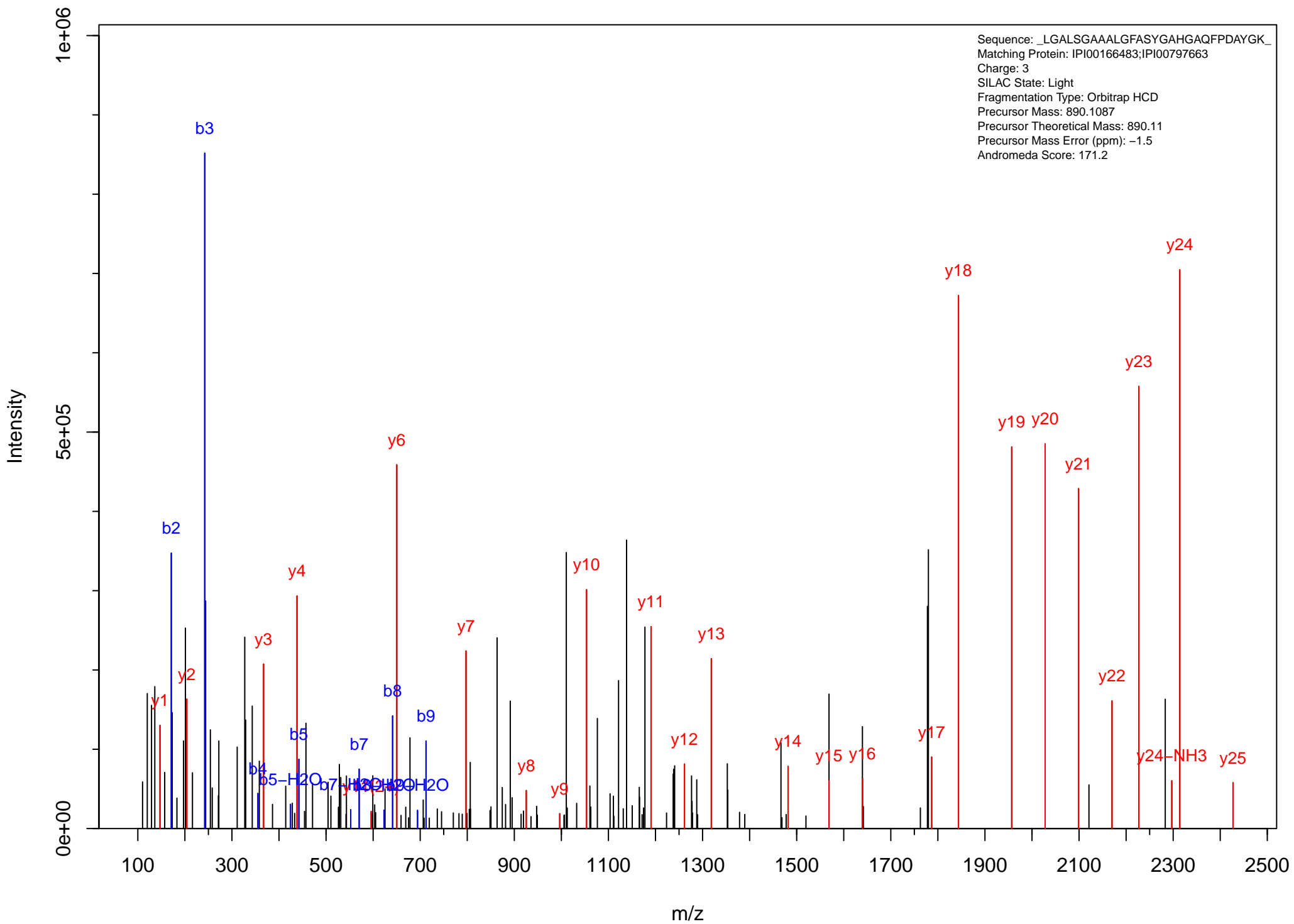


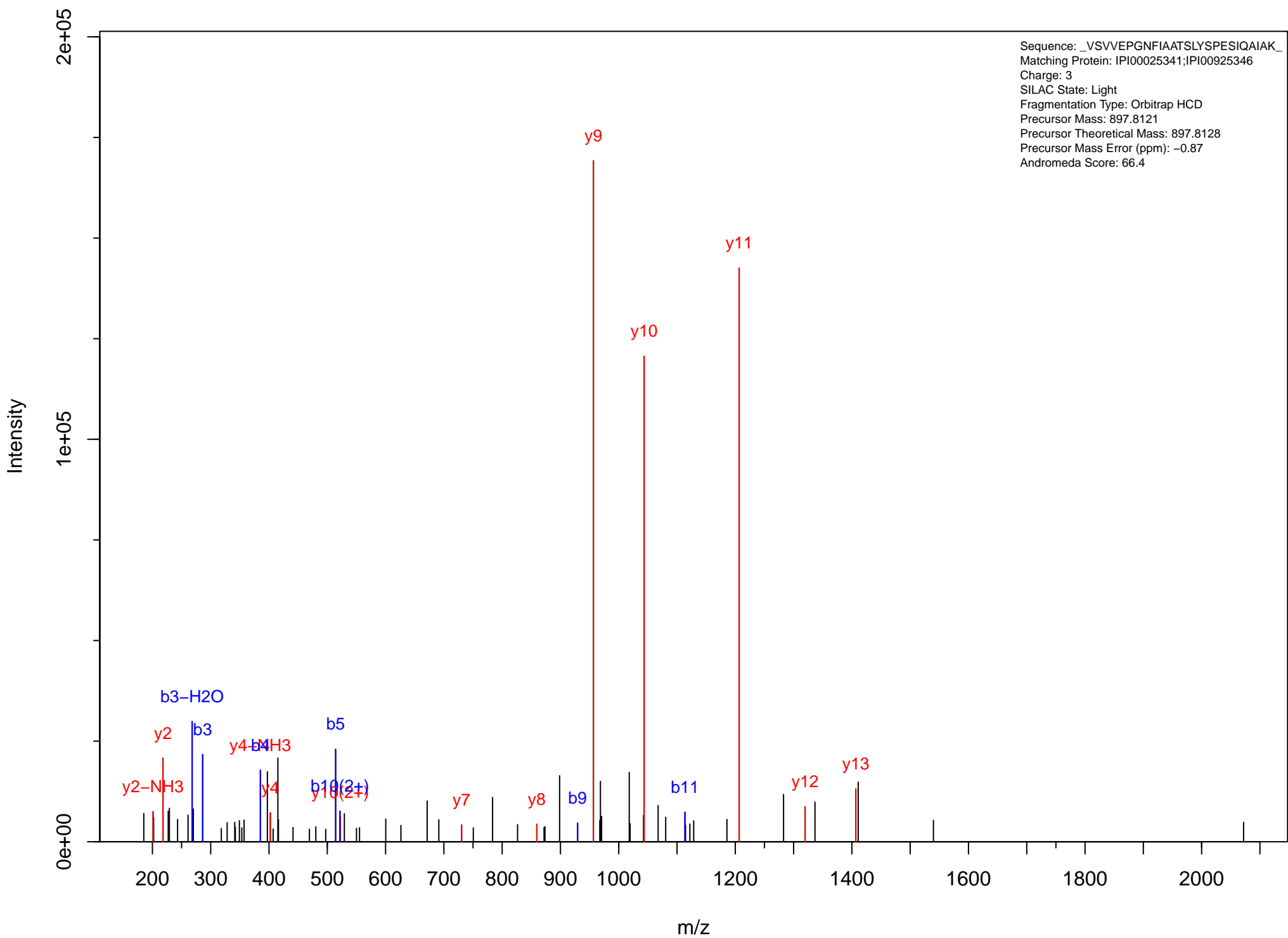


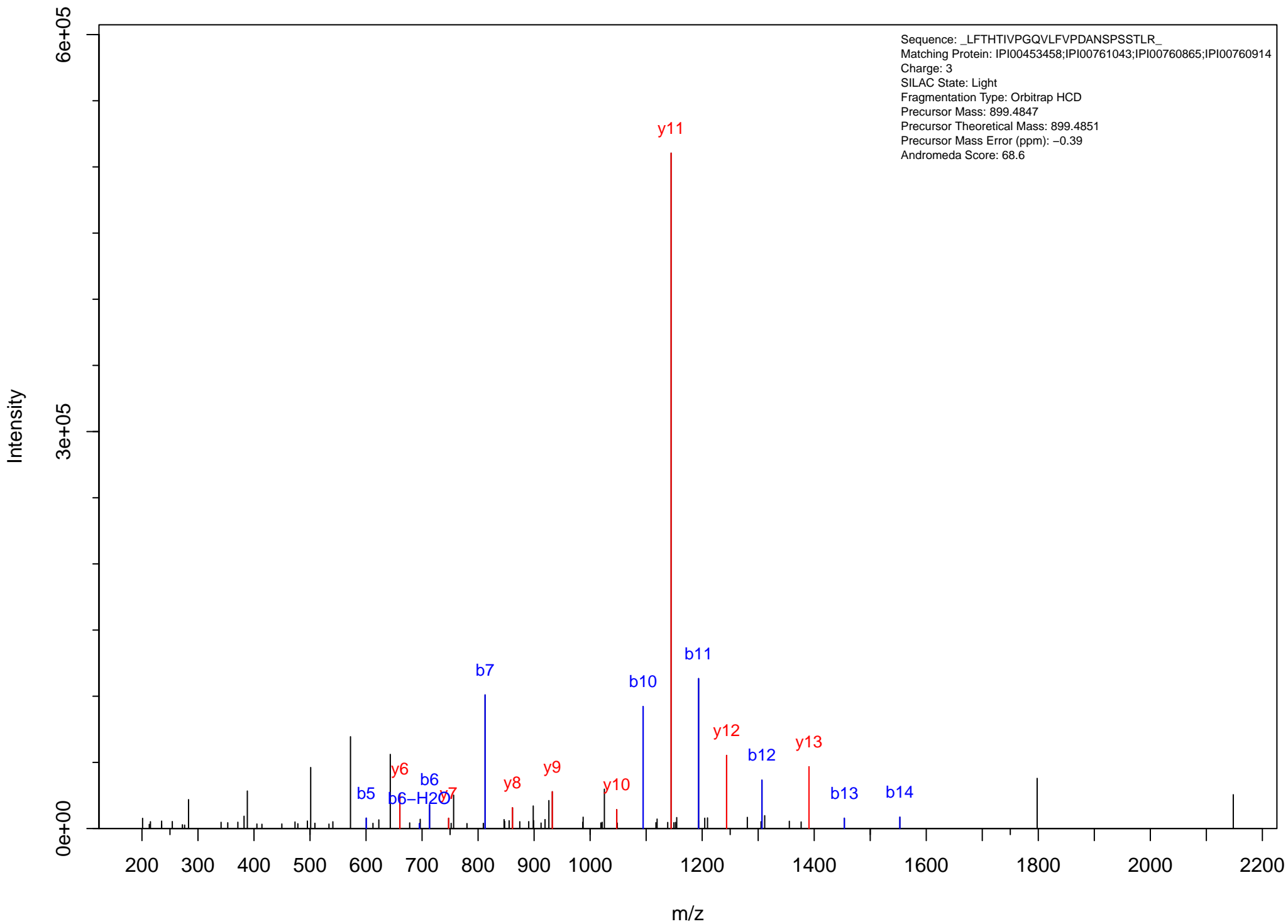


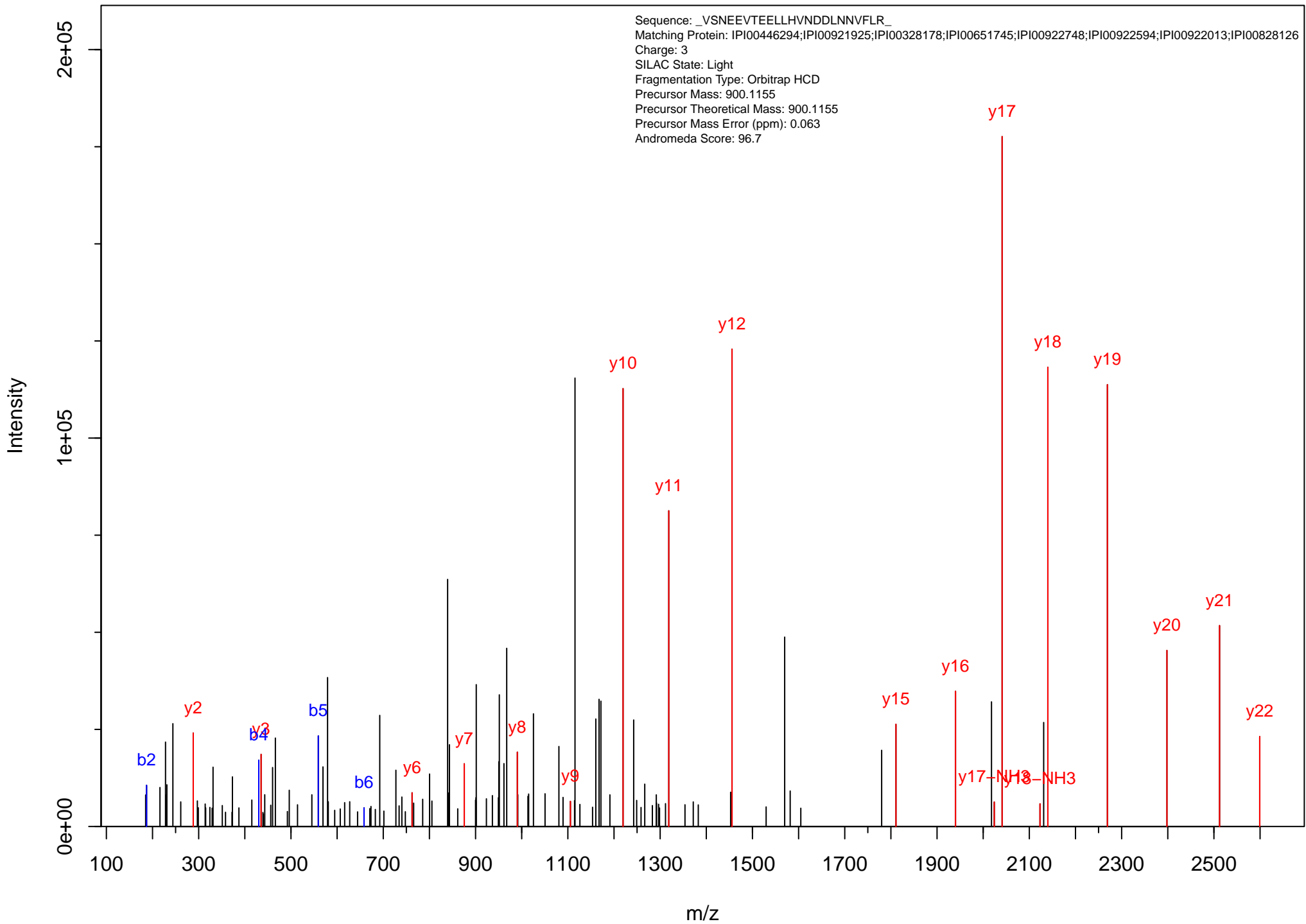


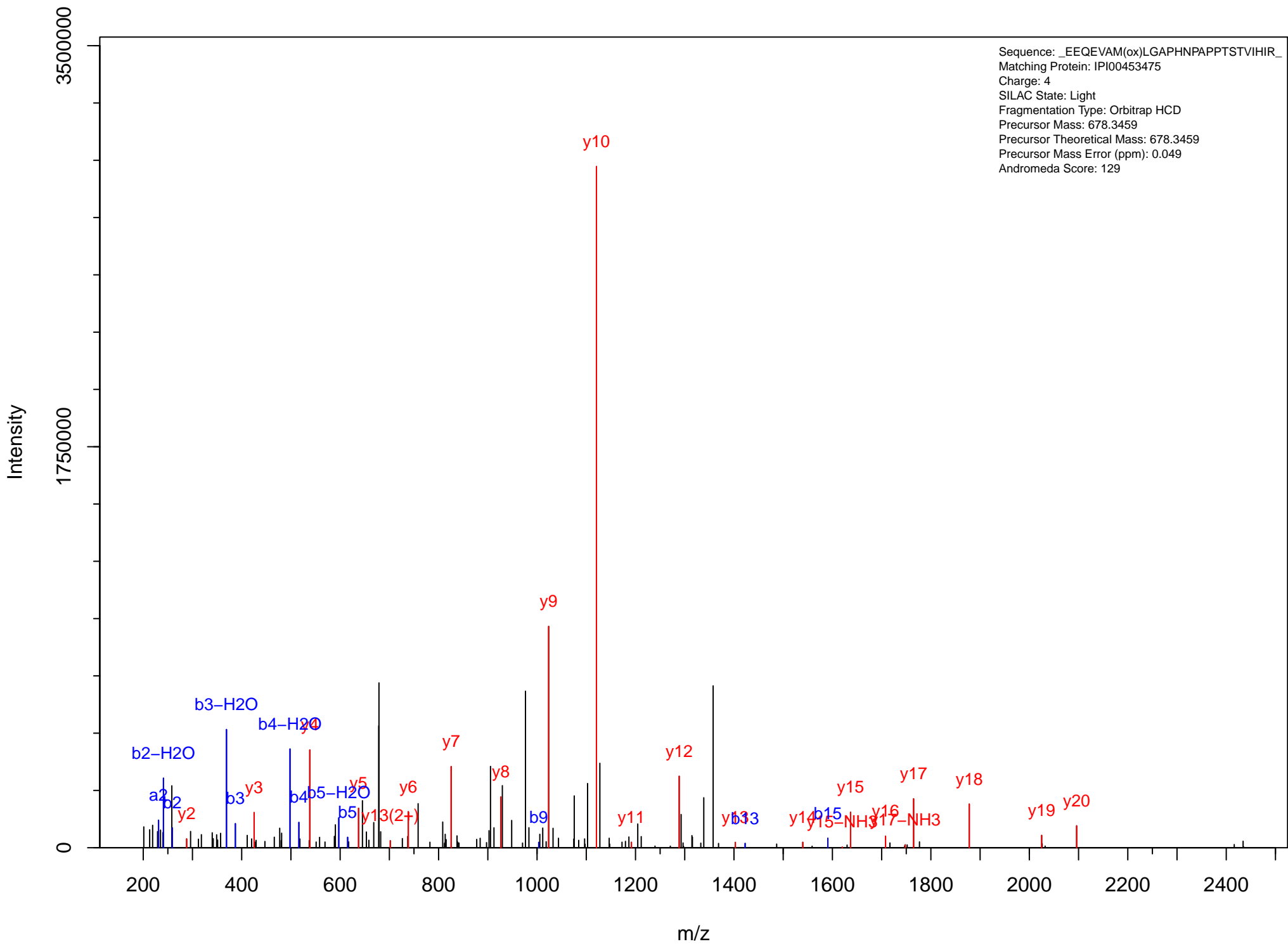


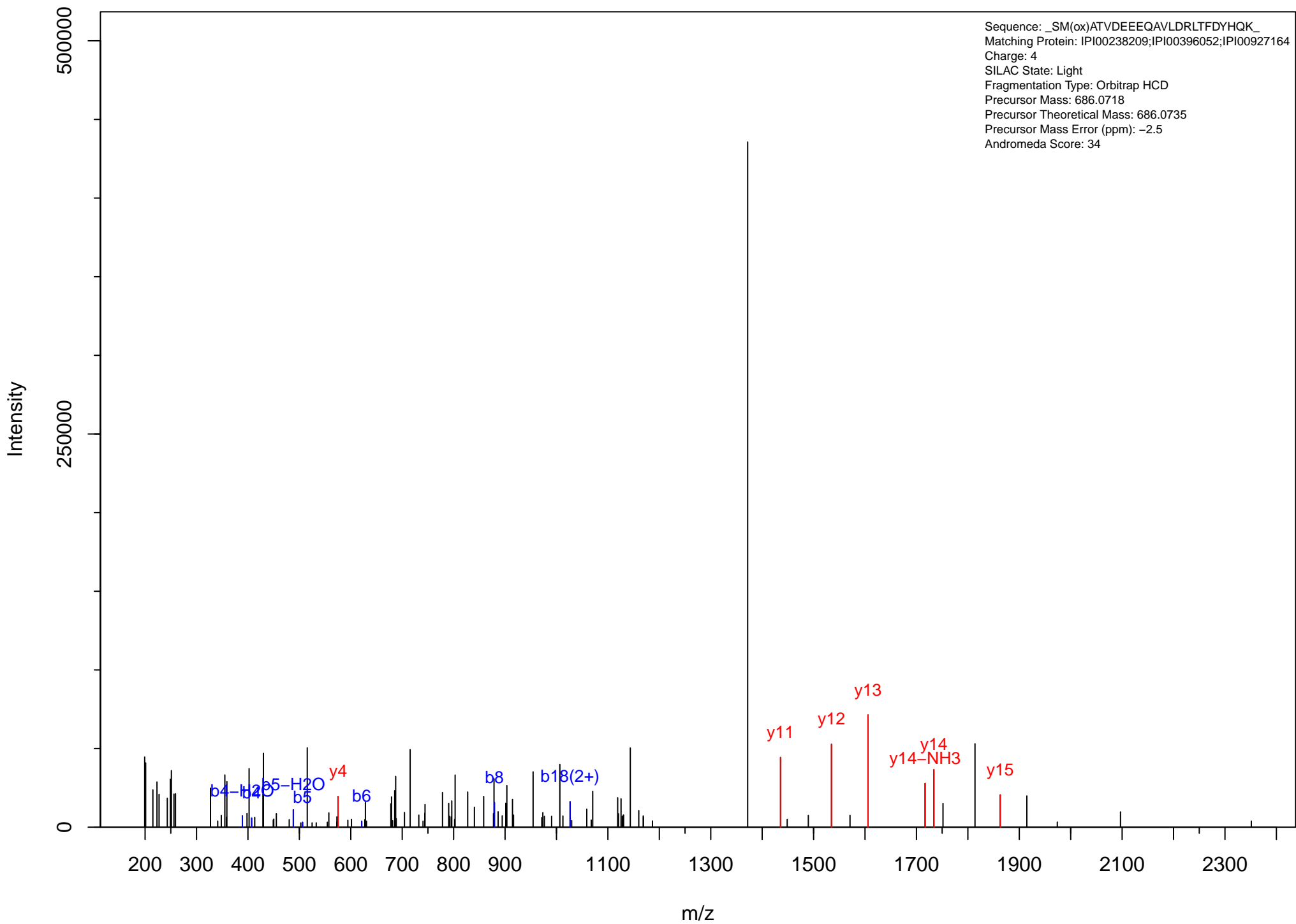


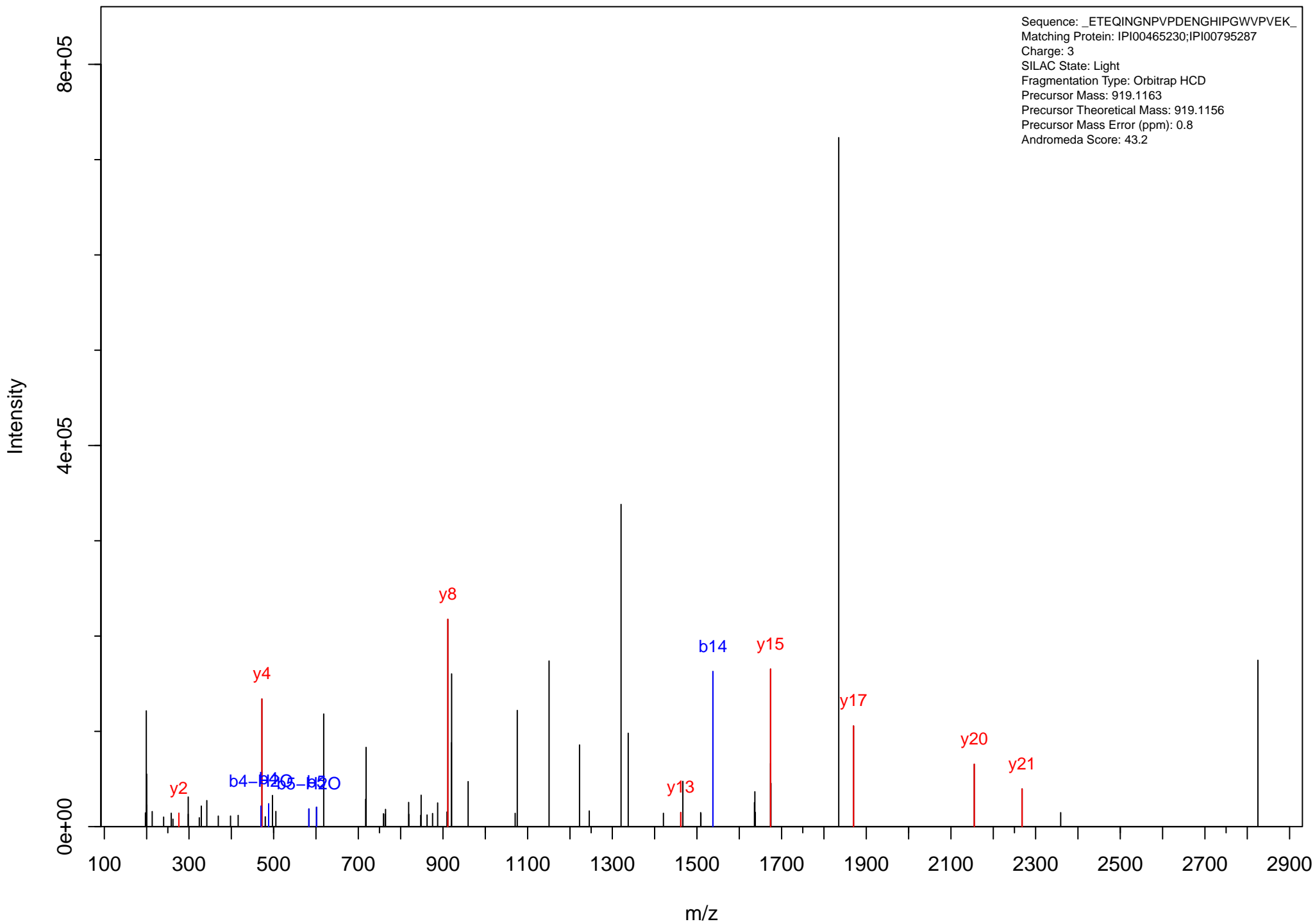


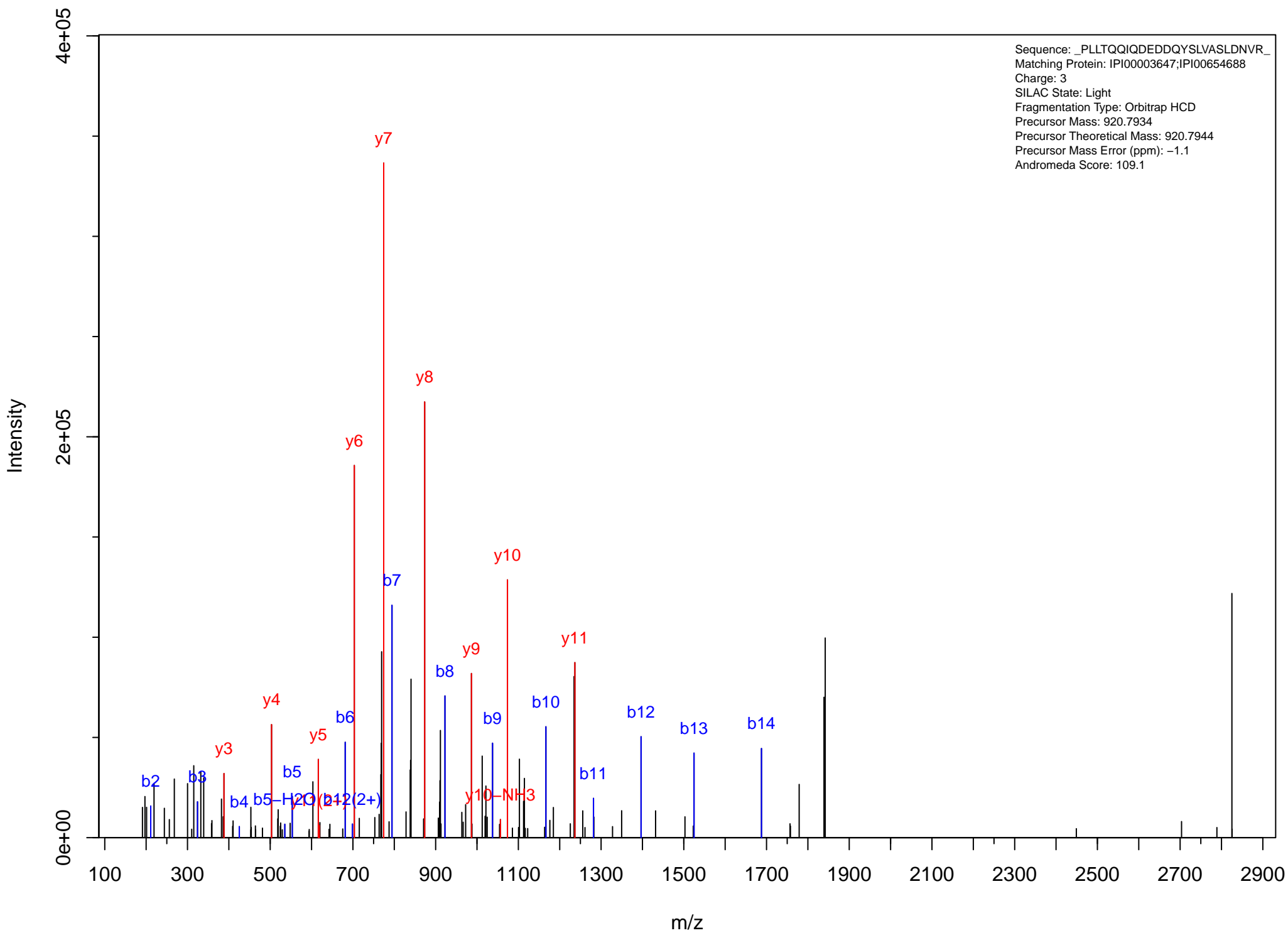


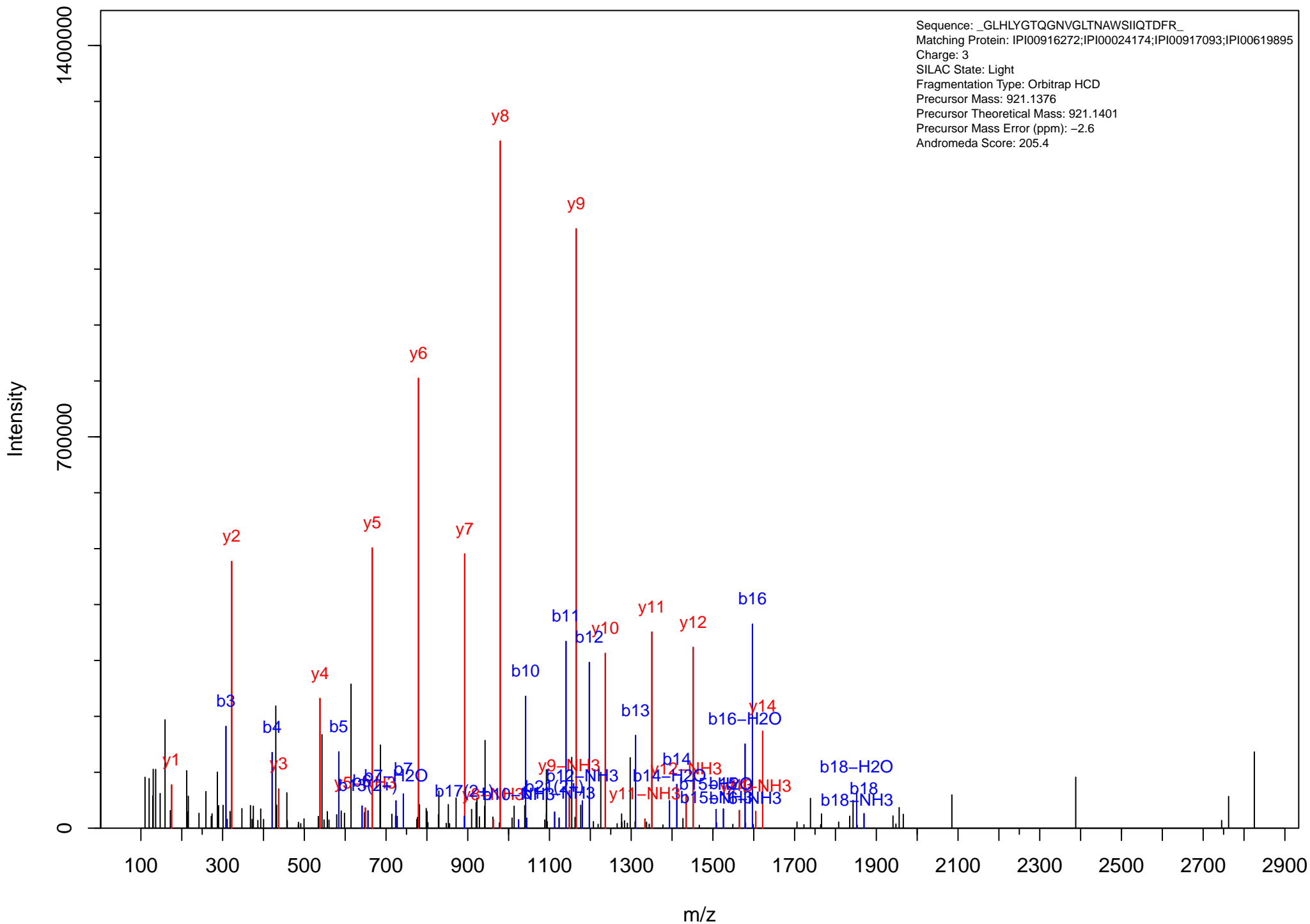


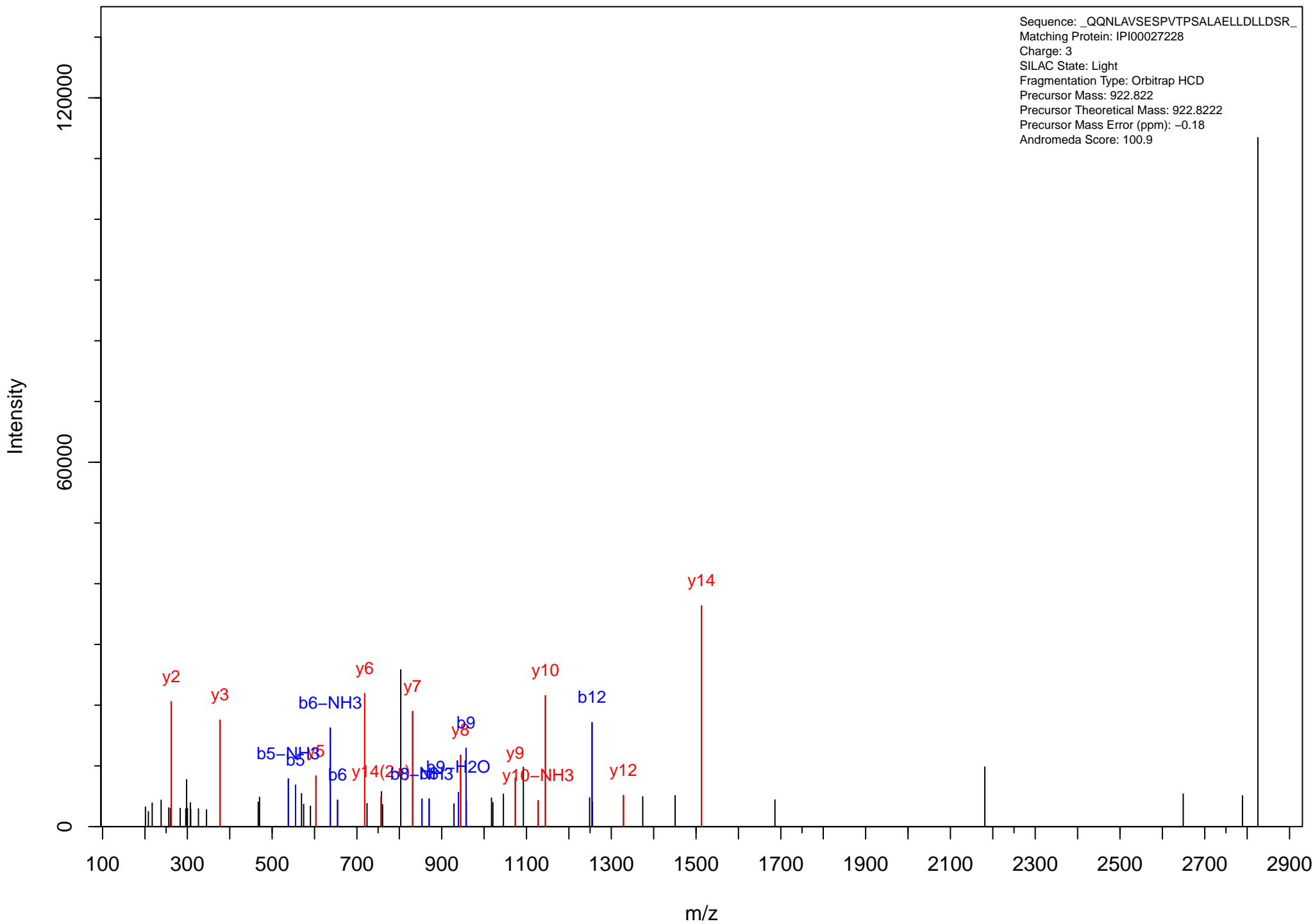


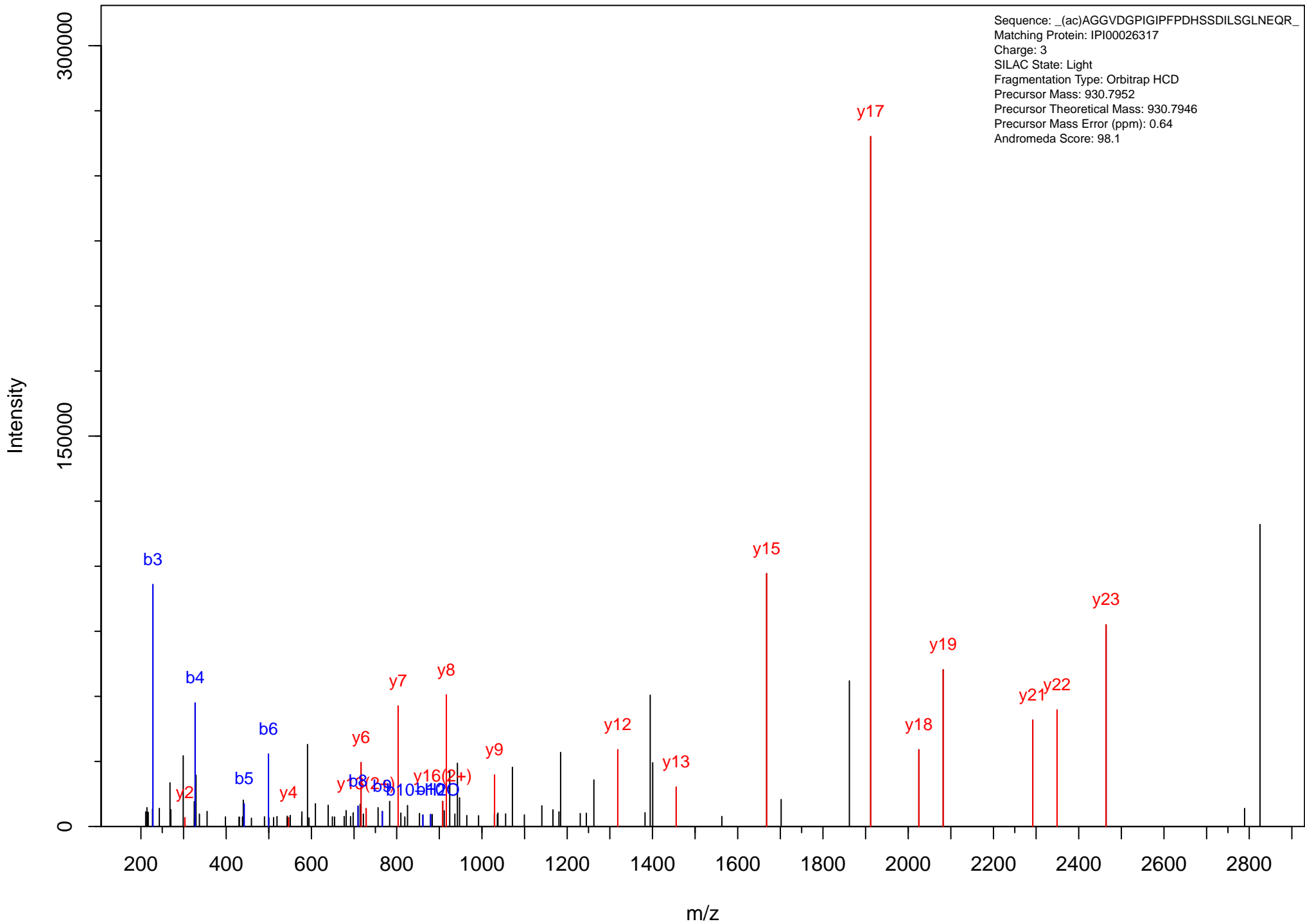


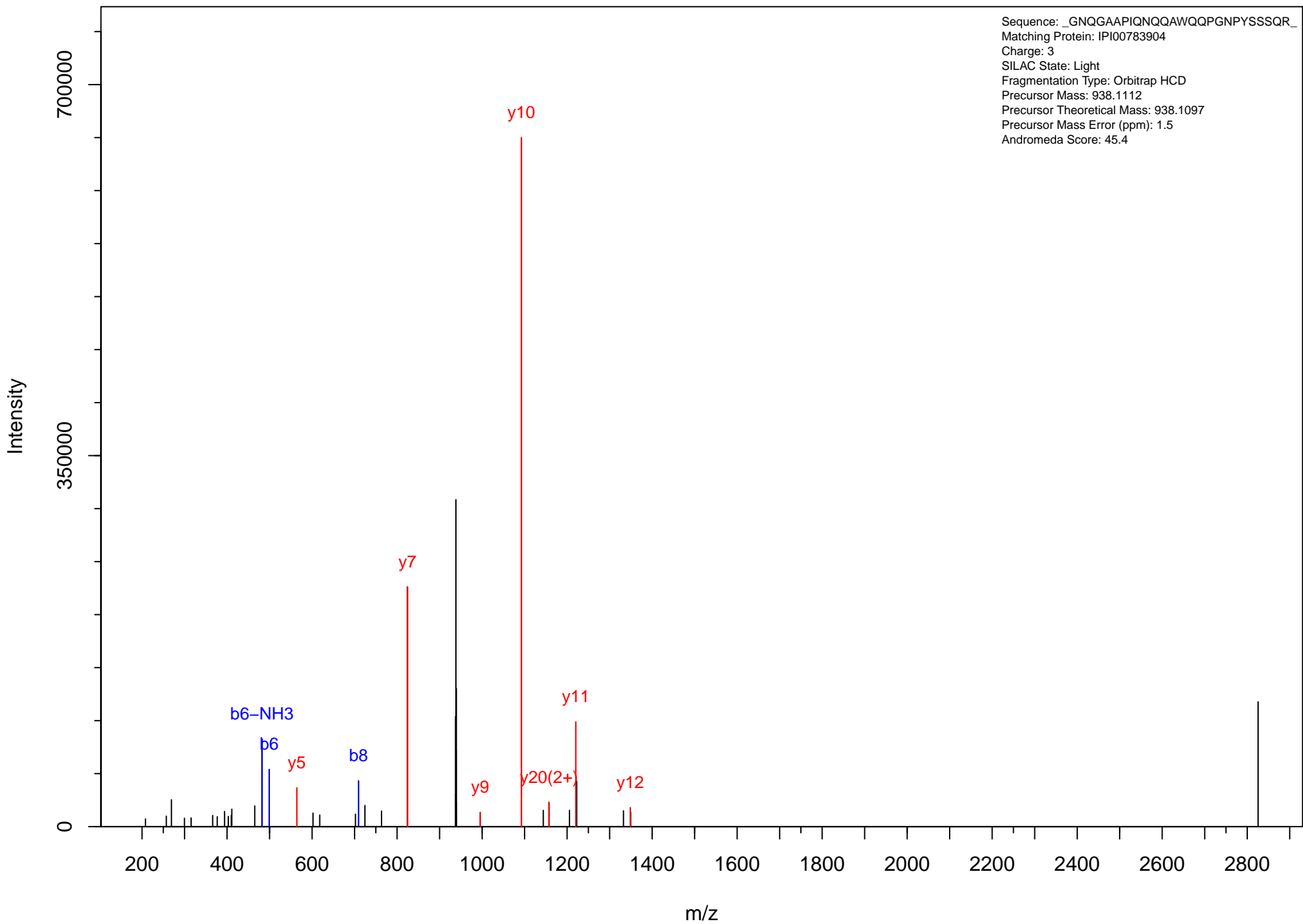


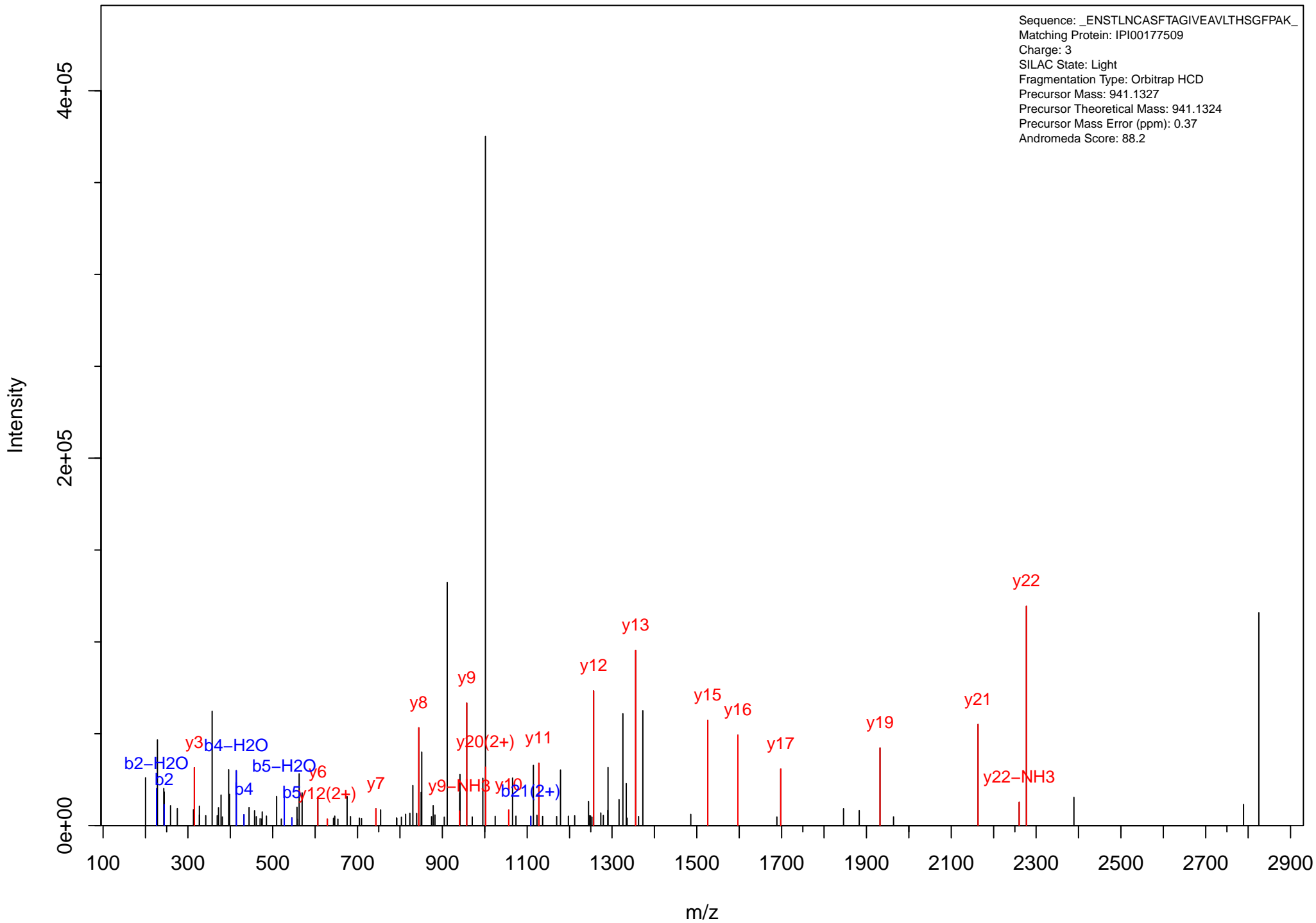


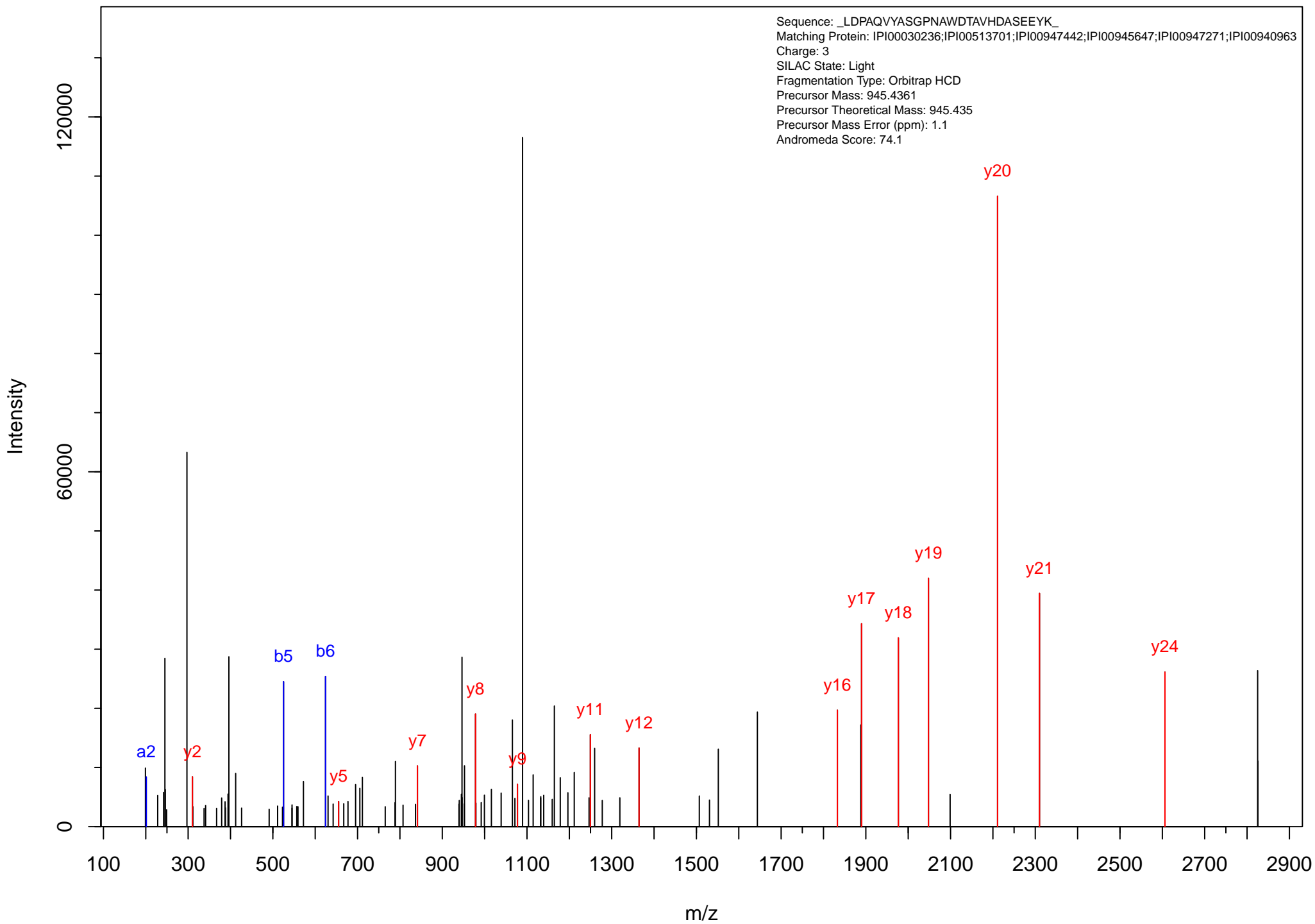


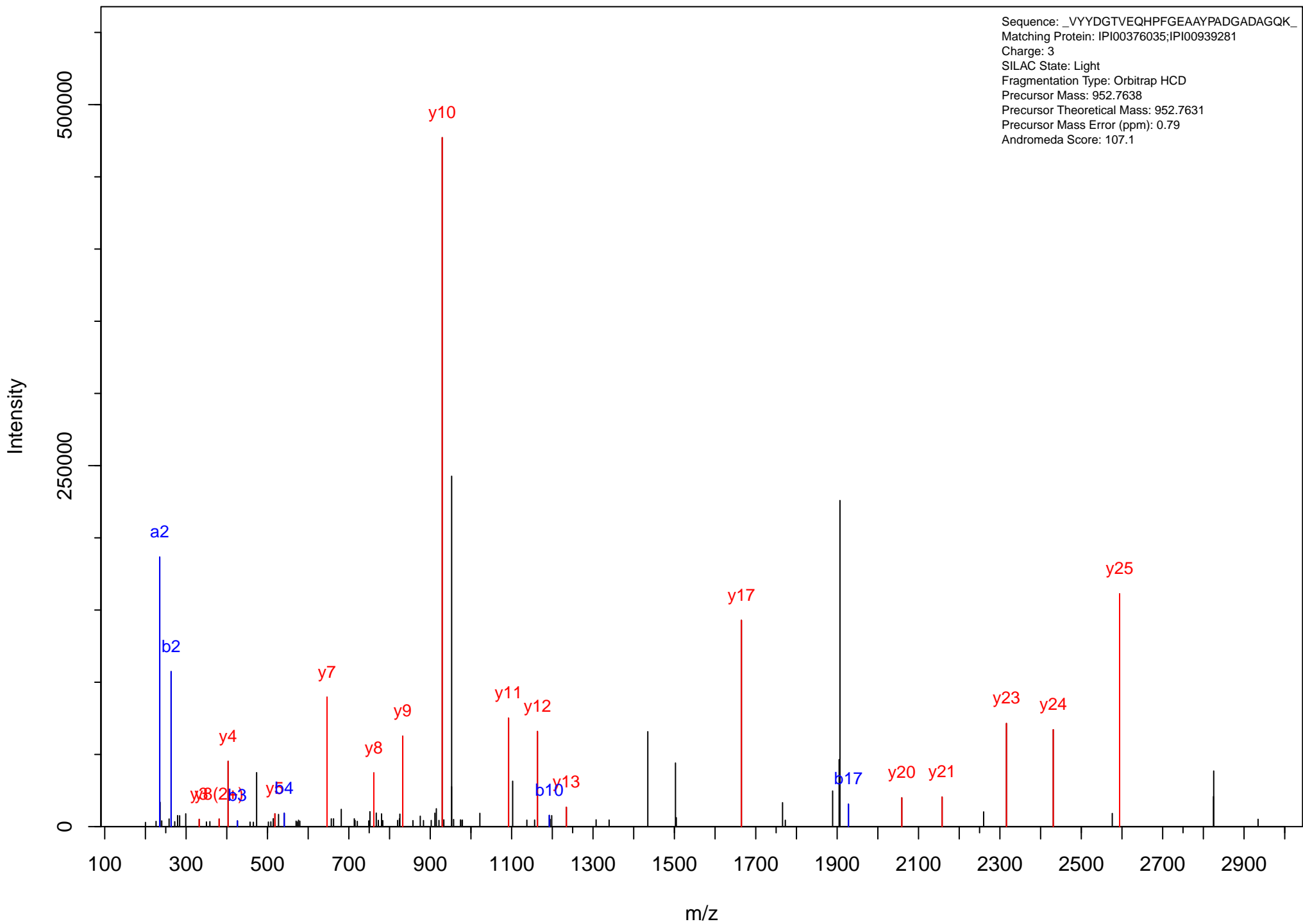


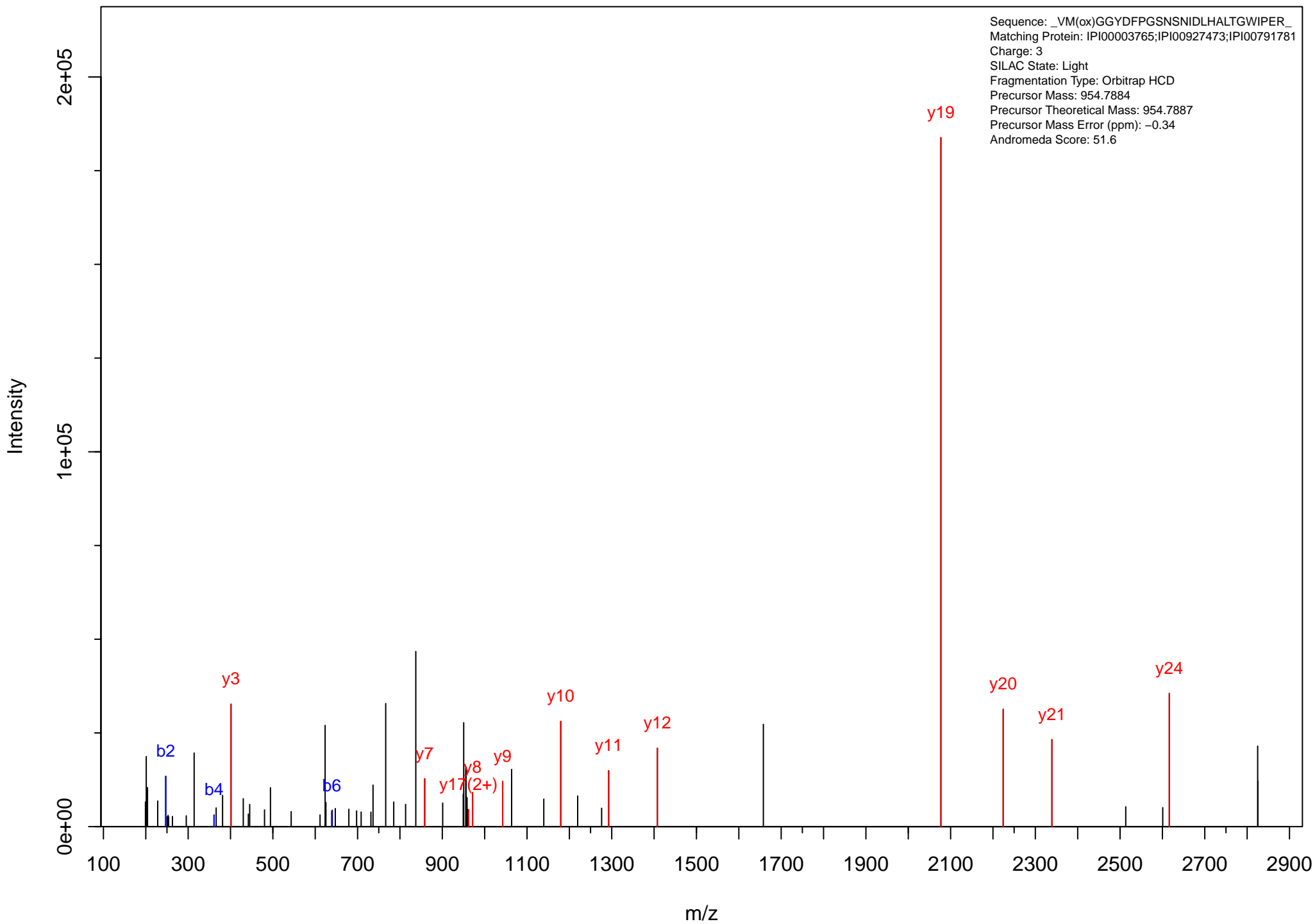




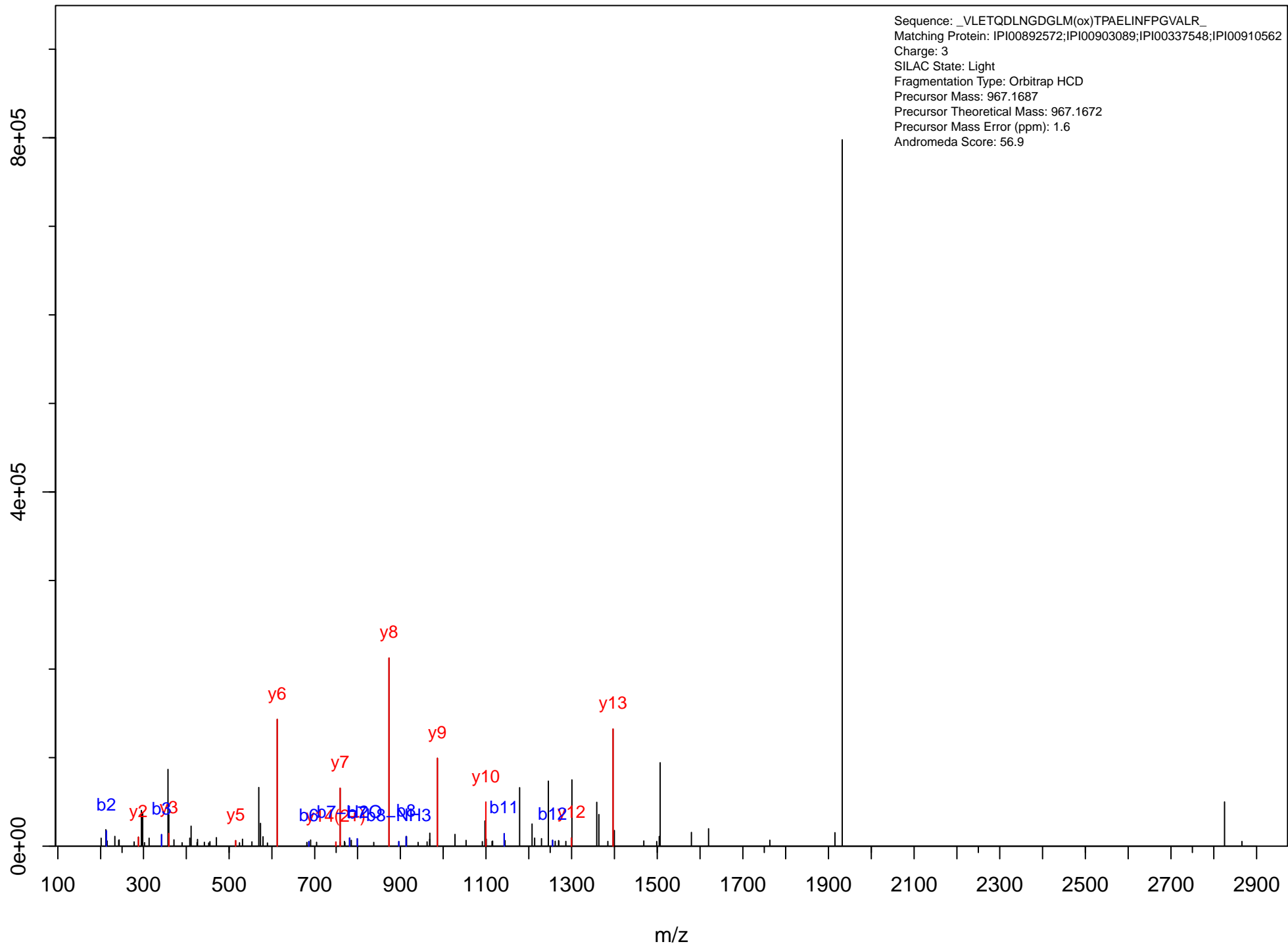


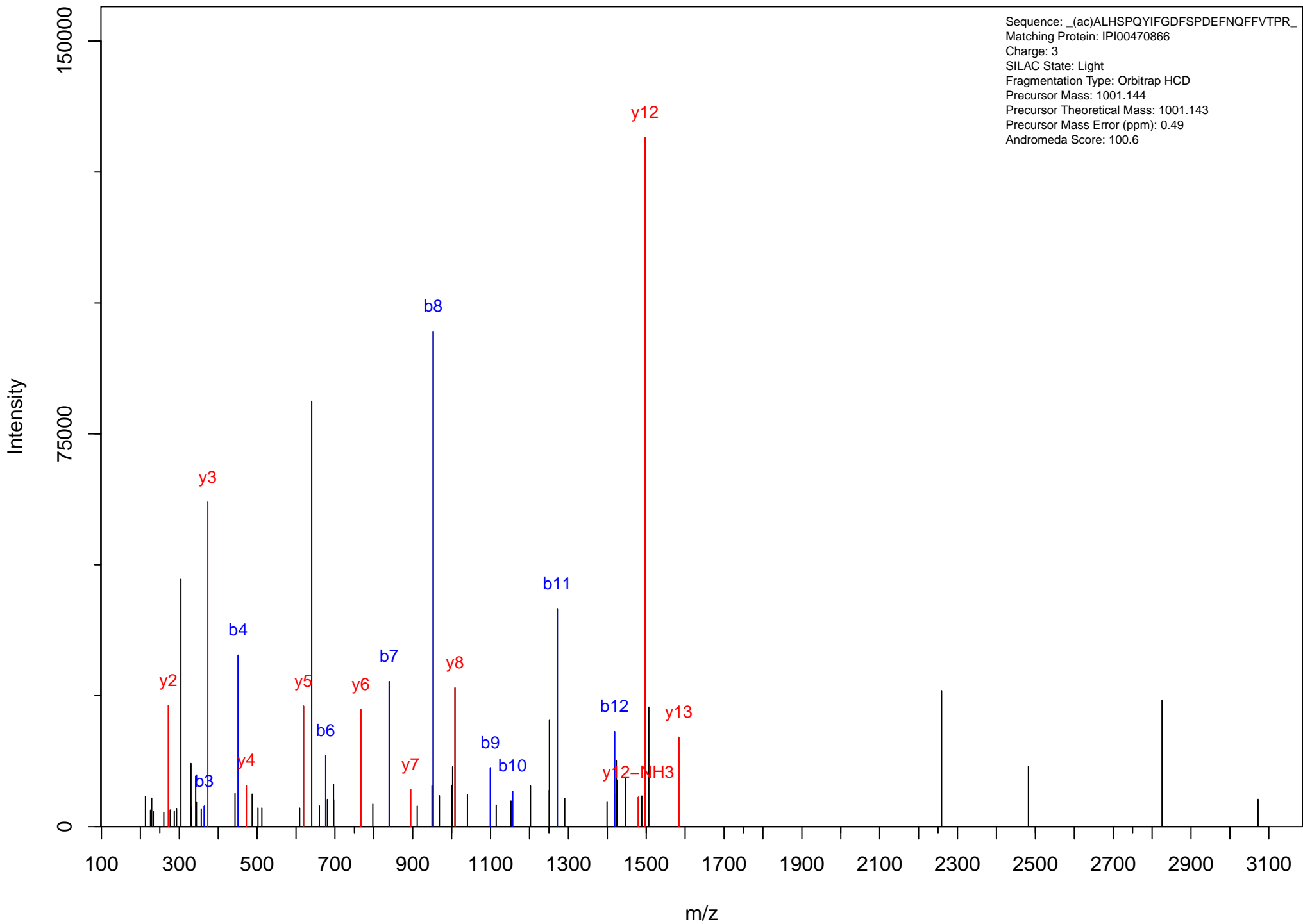


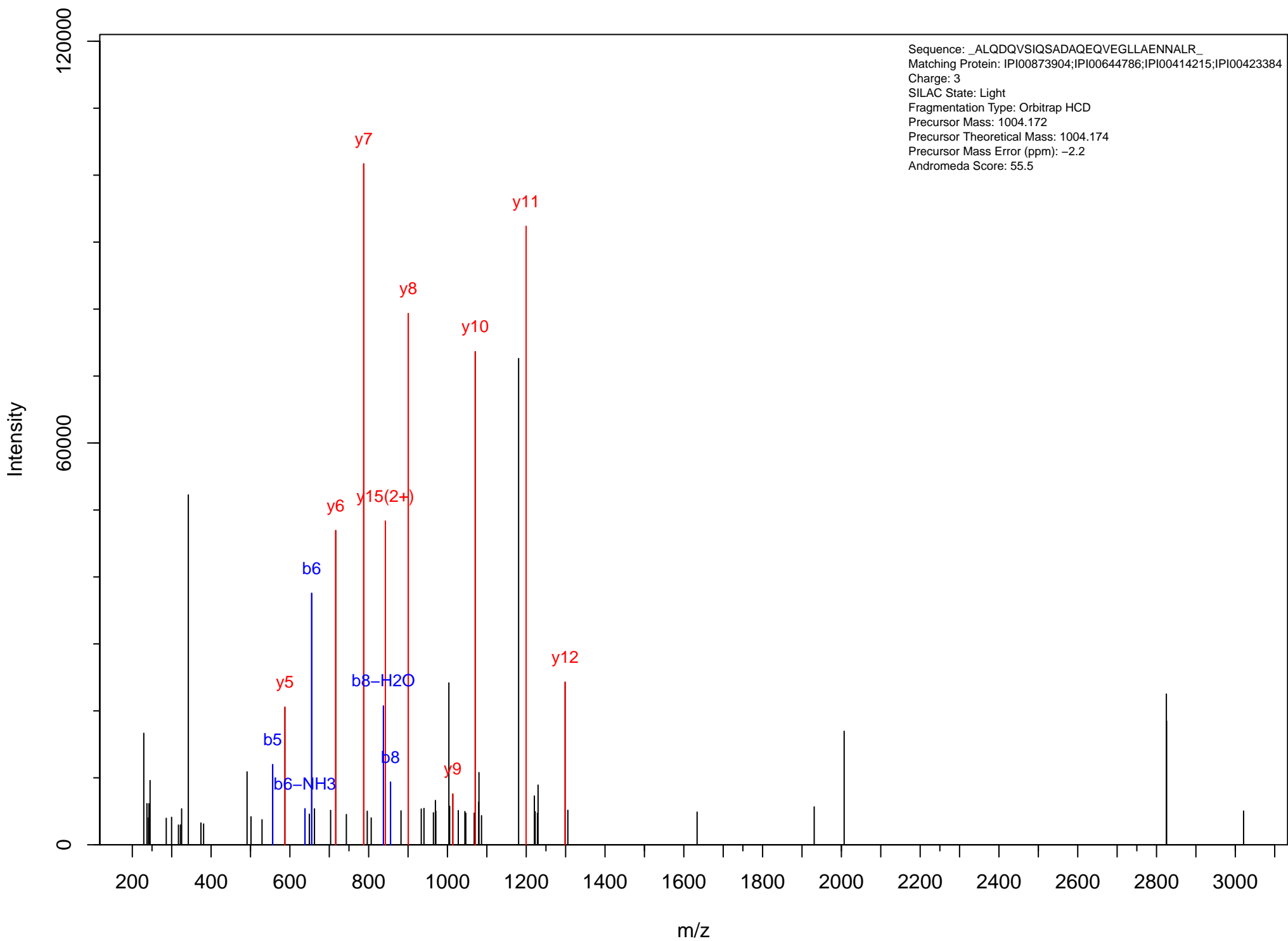


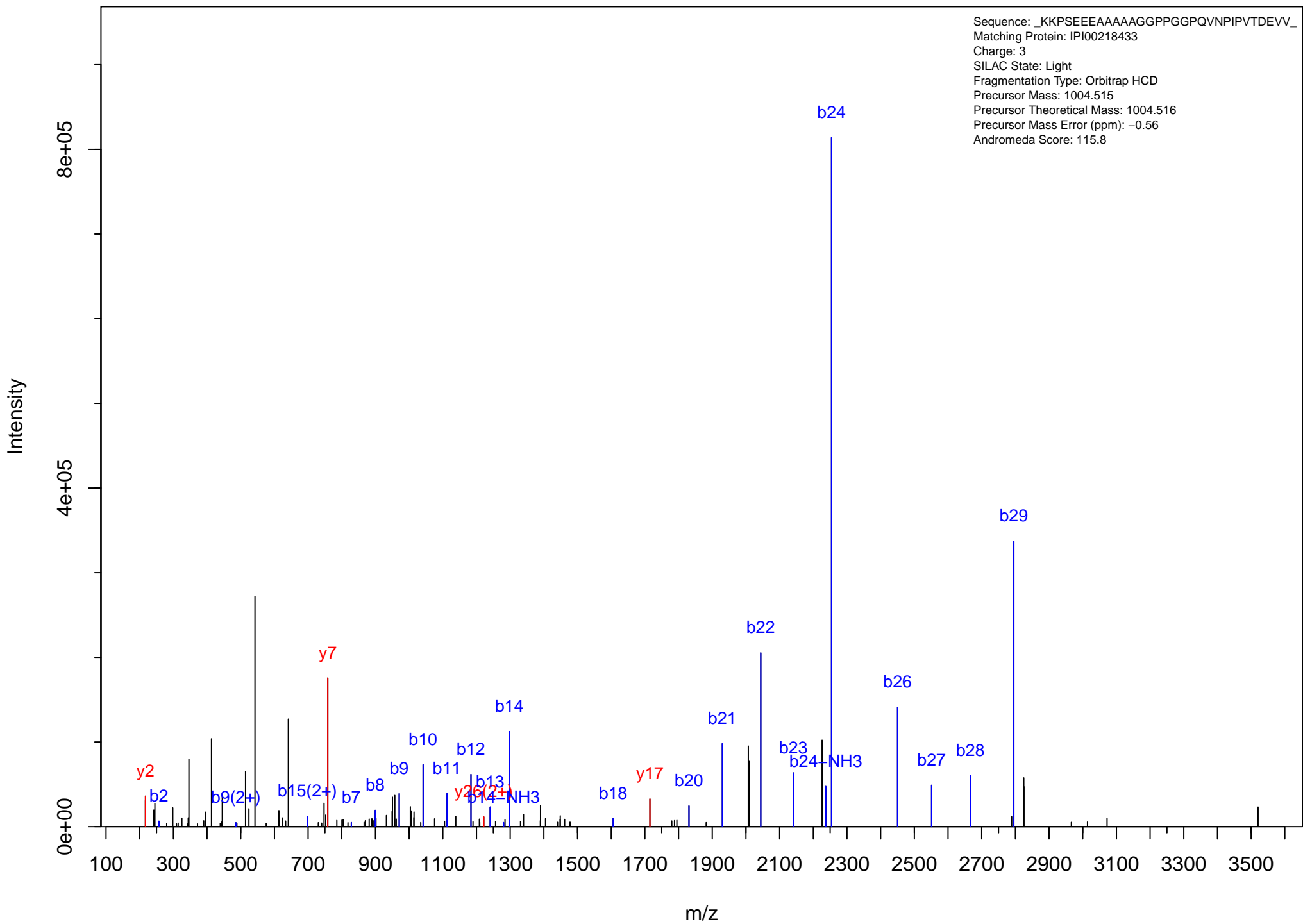


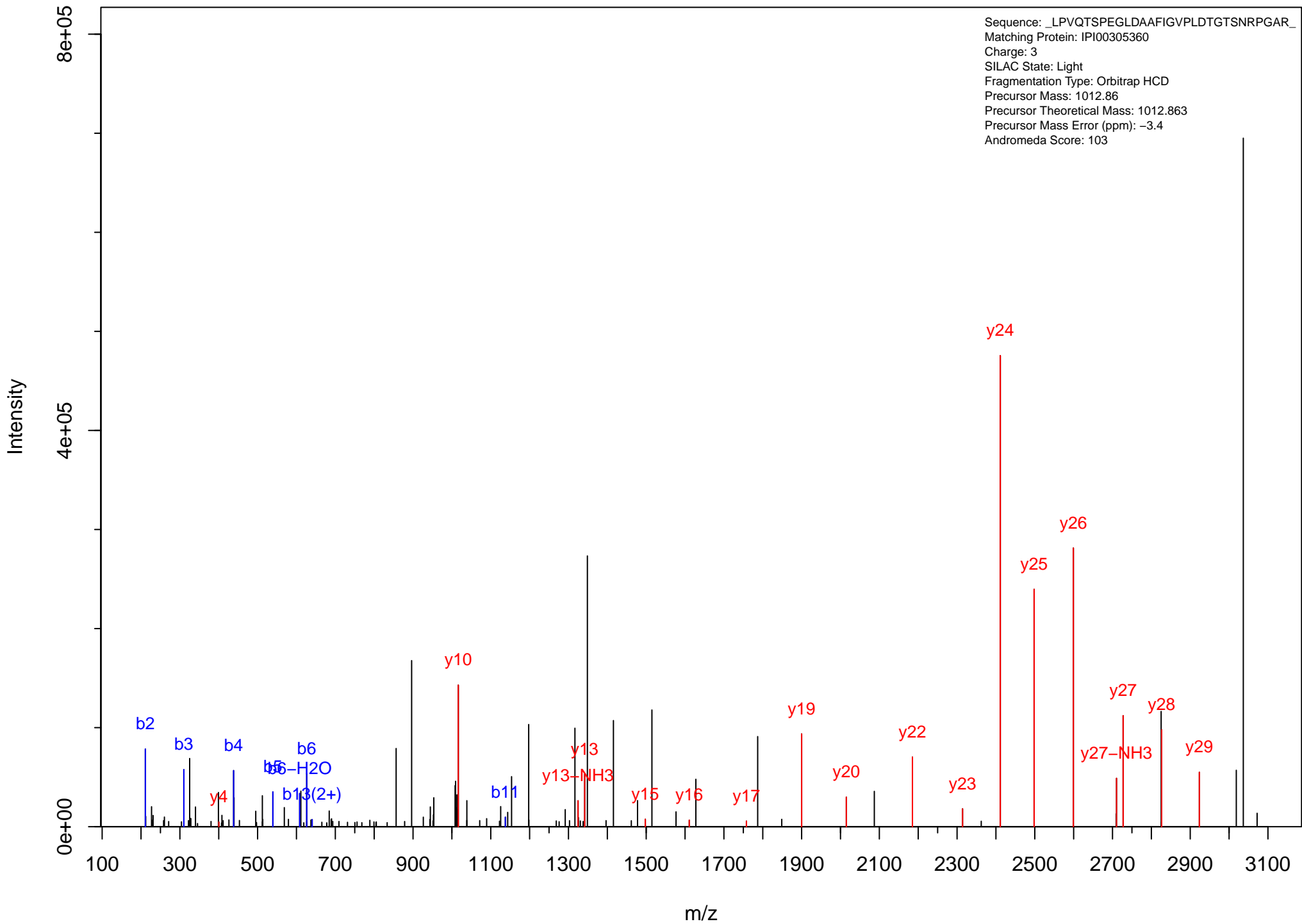
Intensity

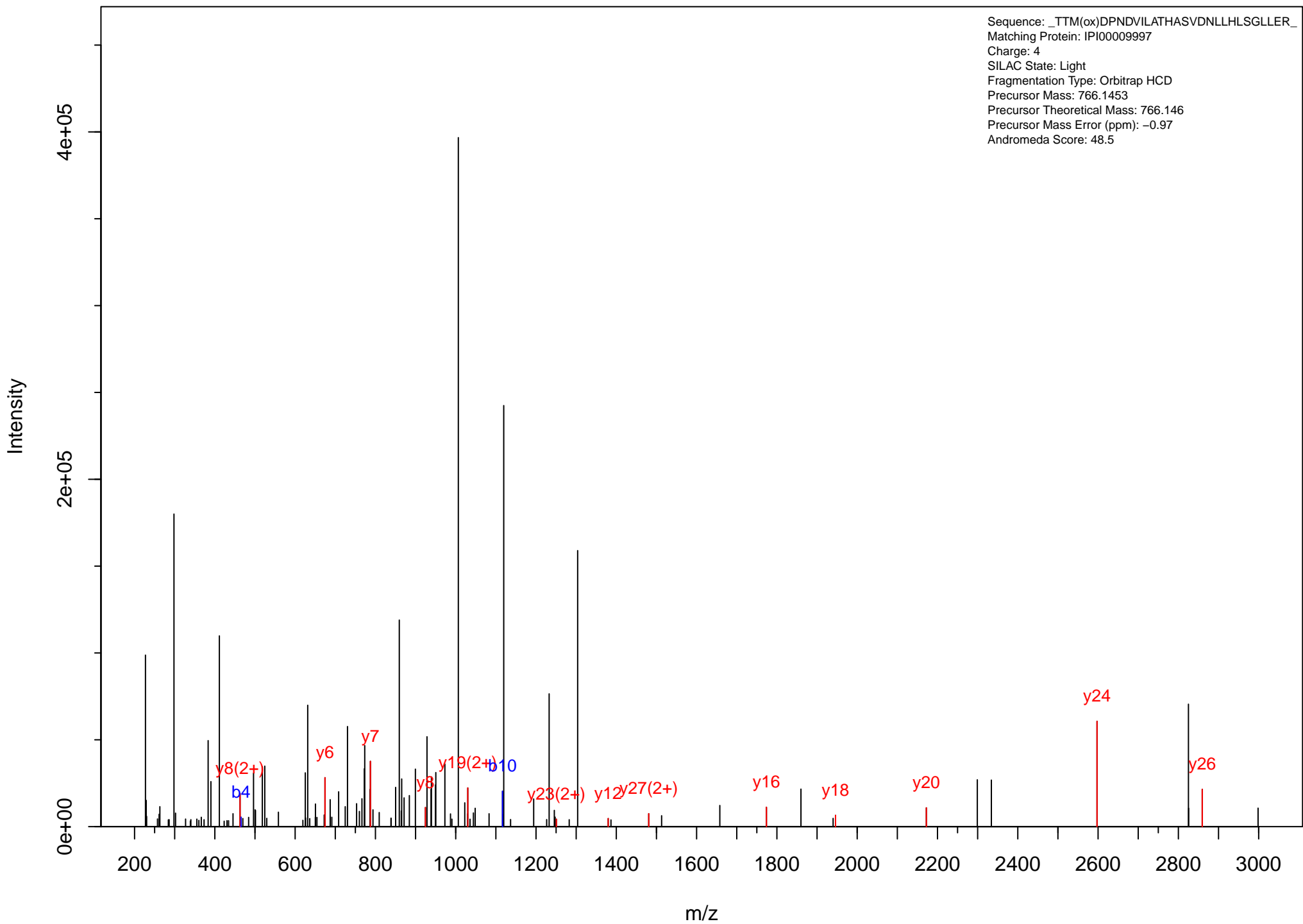


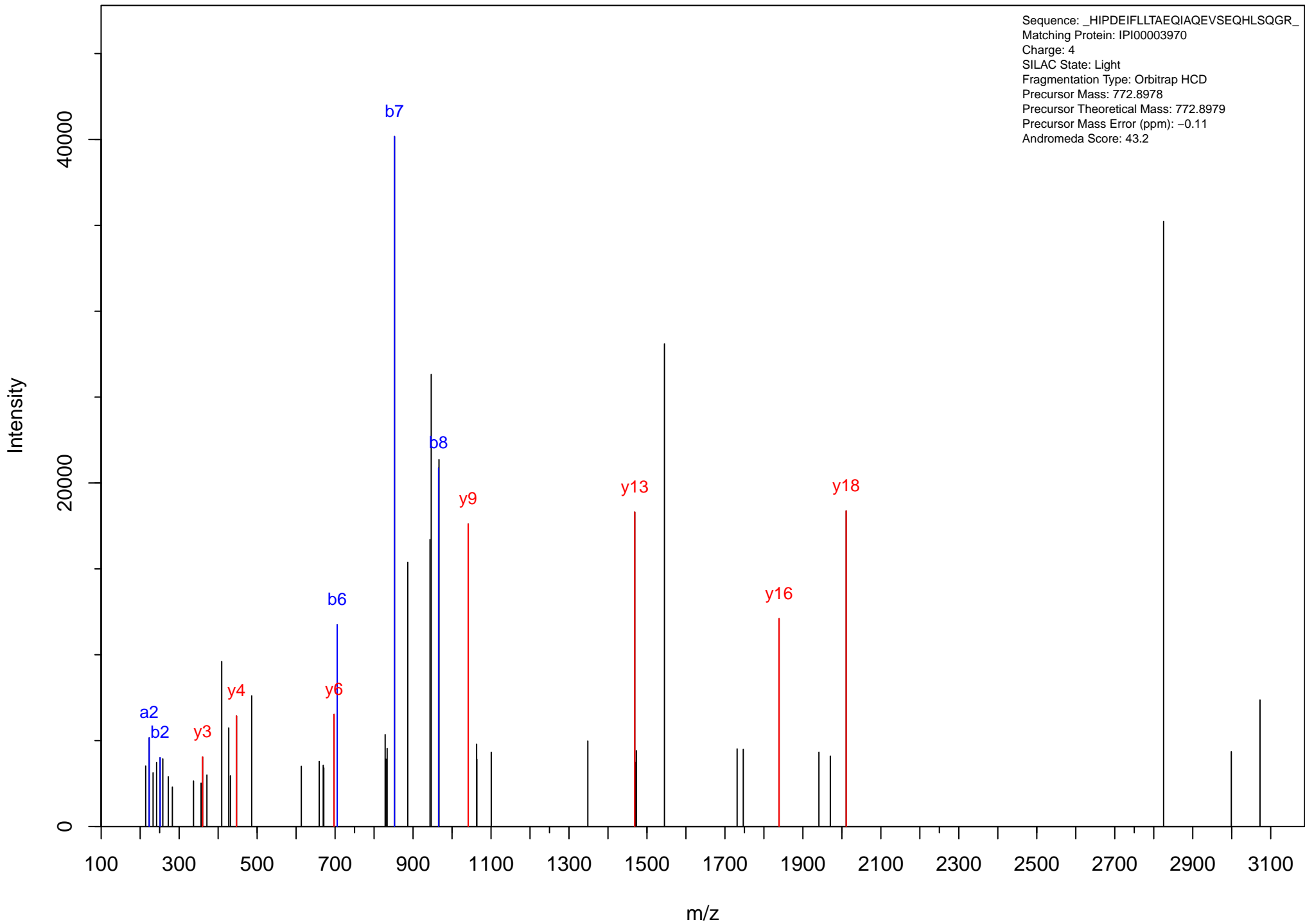


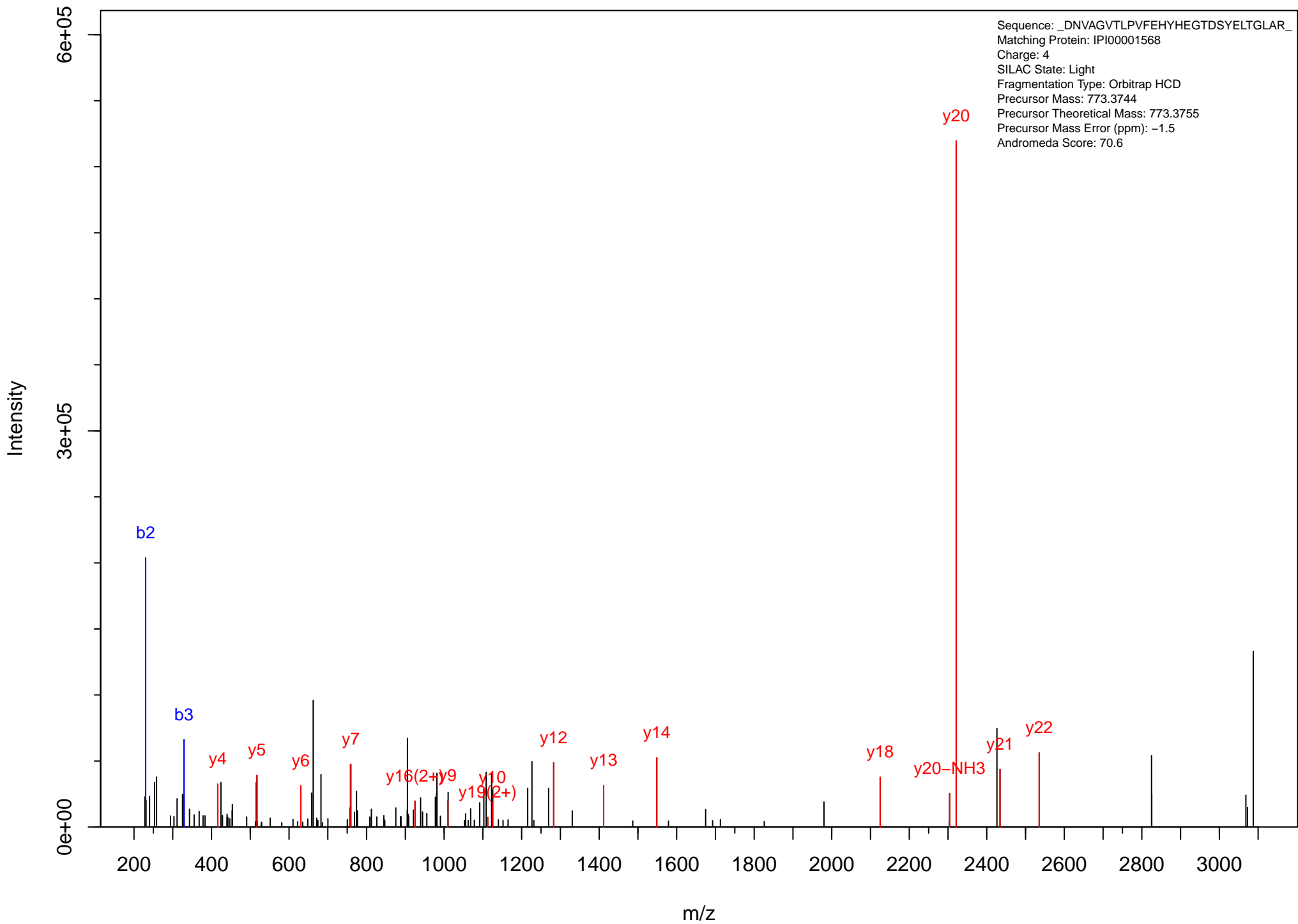


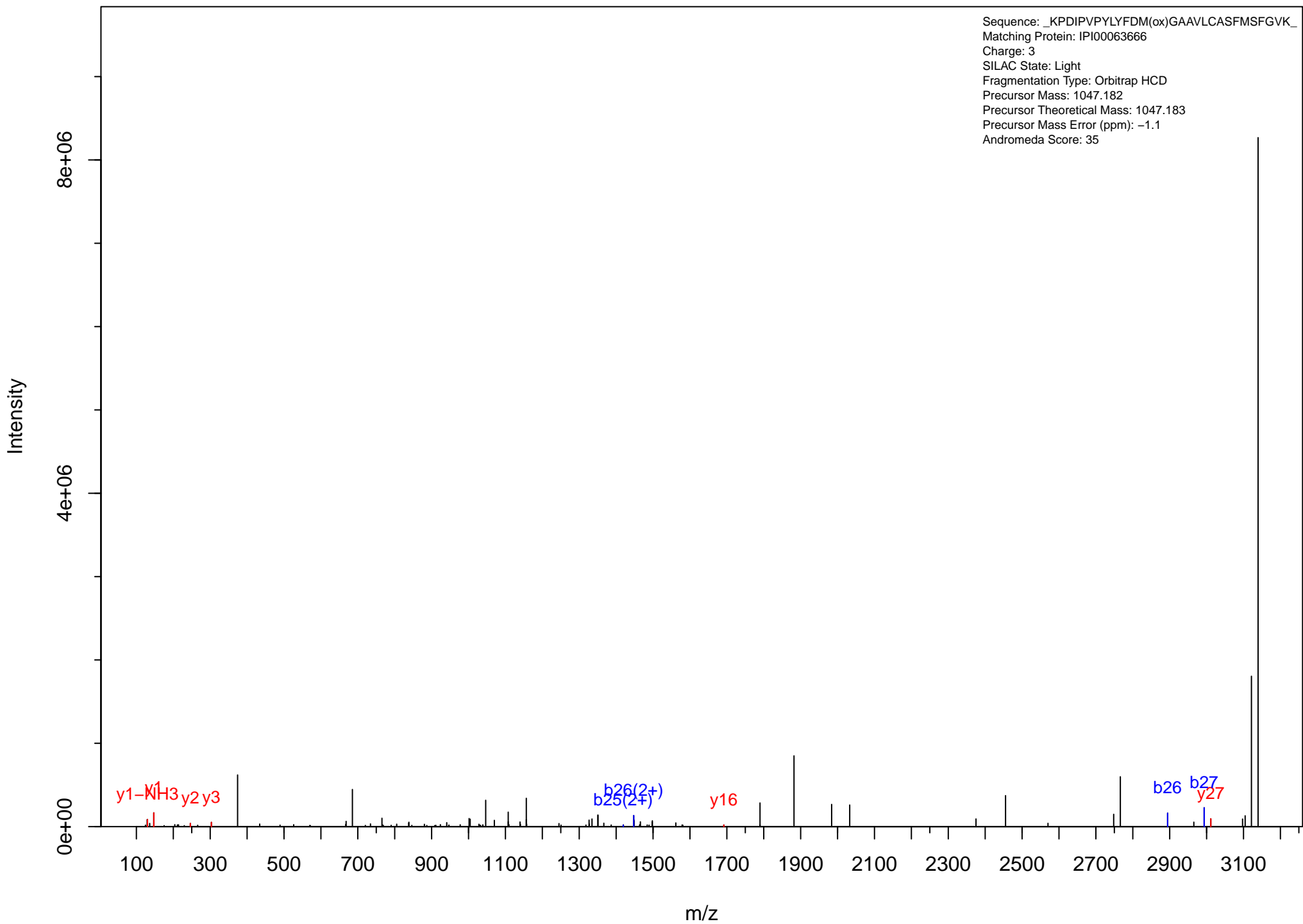


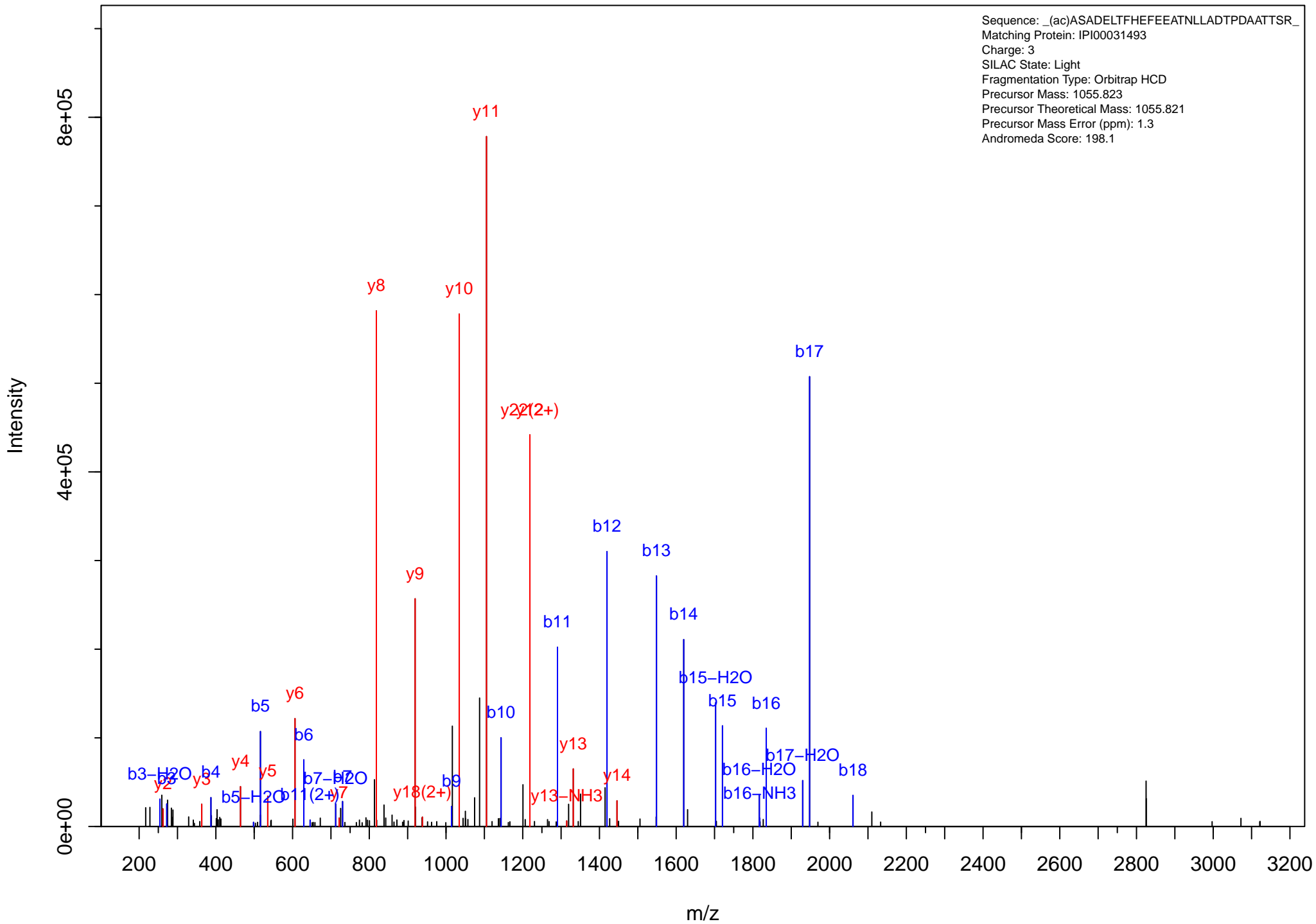












Intensity

4e+05

2e+05

0e+00

200 400 600 800 1000 1200 1400 1600 1800 2000 2200 2400 2600 2800 3000 3200

m/z

Sequence: _SSNDHGSLALTPRPQLELAESSPAASFLSSR_
Matching Protein: IPI00106549;IPI00908597;IPI00737502
Charge: 4
SILAC State: Light
Fragmentation Type: Orbitrap HCD
Precursor Mass: 807.1564
Precursor Theoretical Mass: 807.1588
Precursor Mass Error (ppm): -3
Andromeda Score: 61

y9

y6

y5

y3

y8

y10

y11

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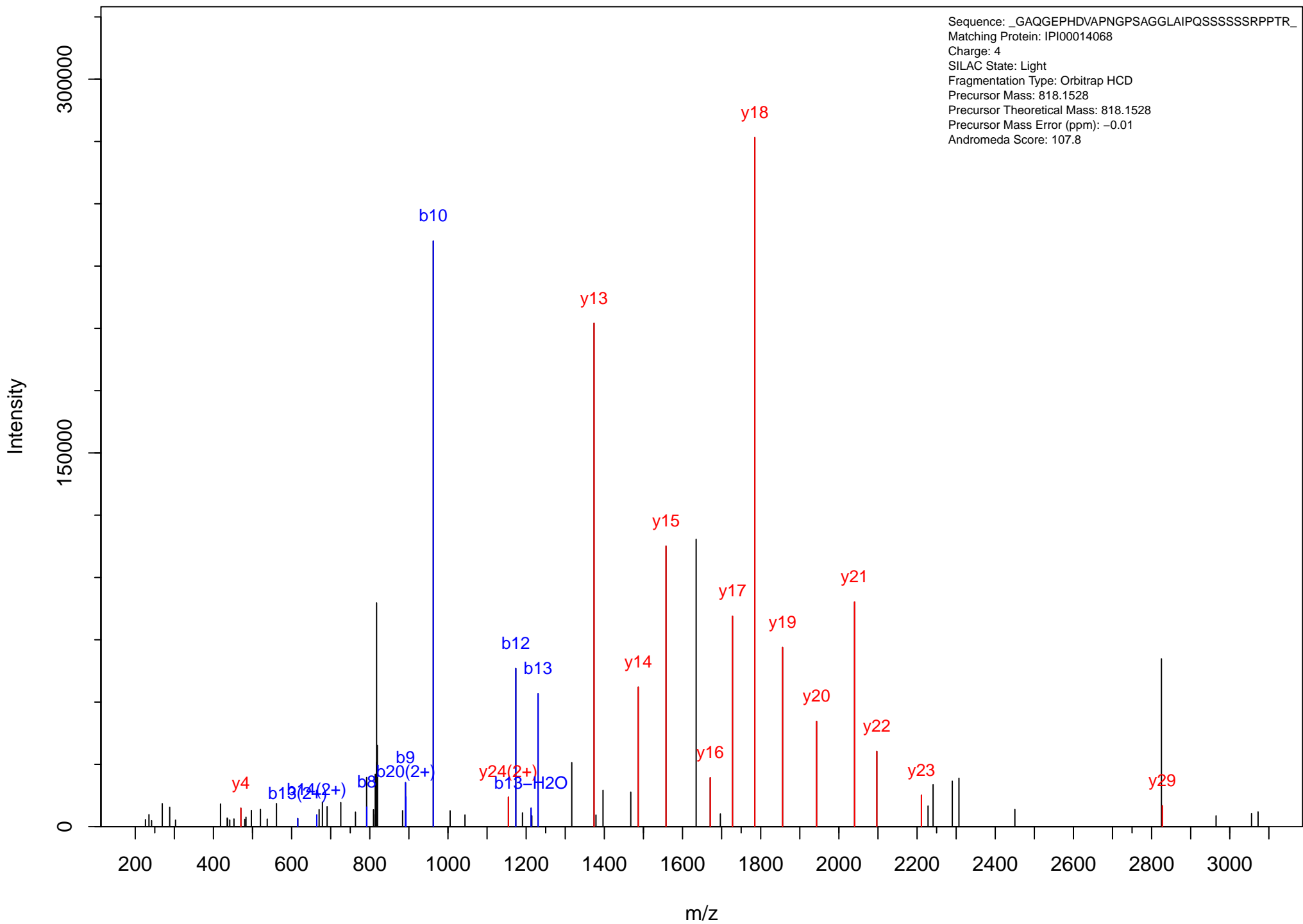
y10

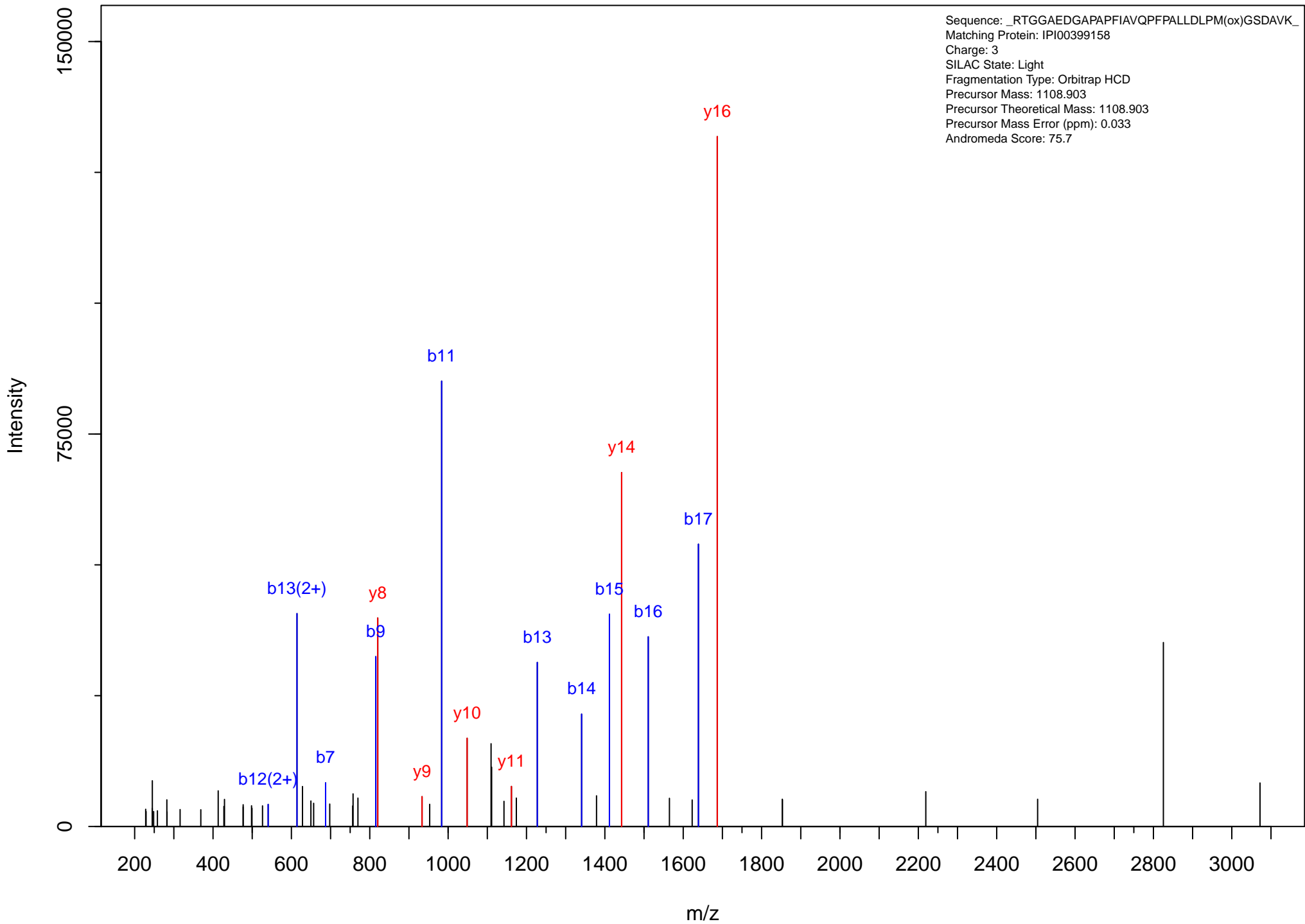
y11

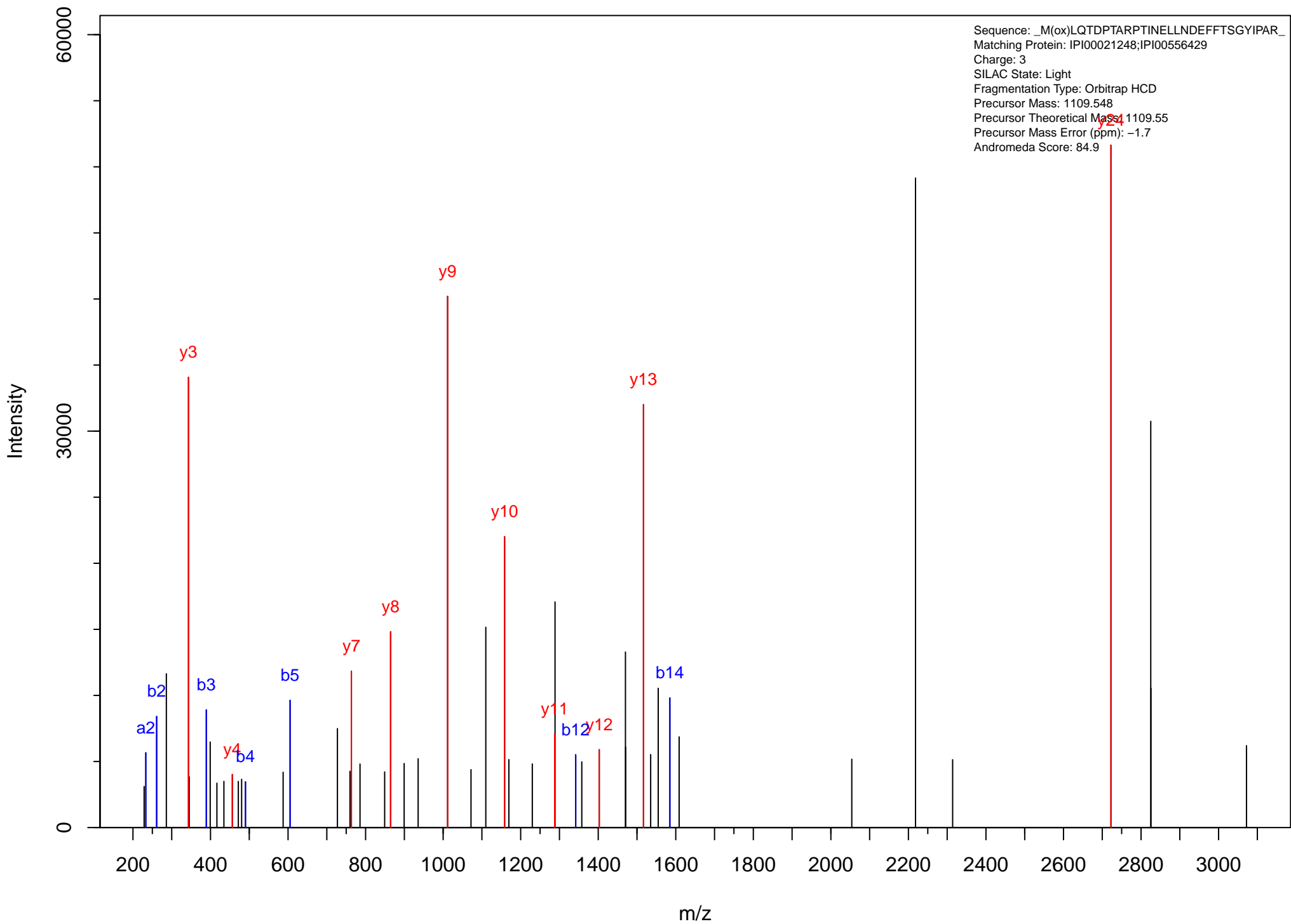
y12

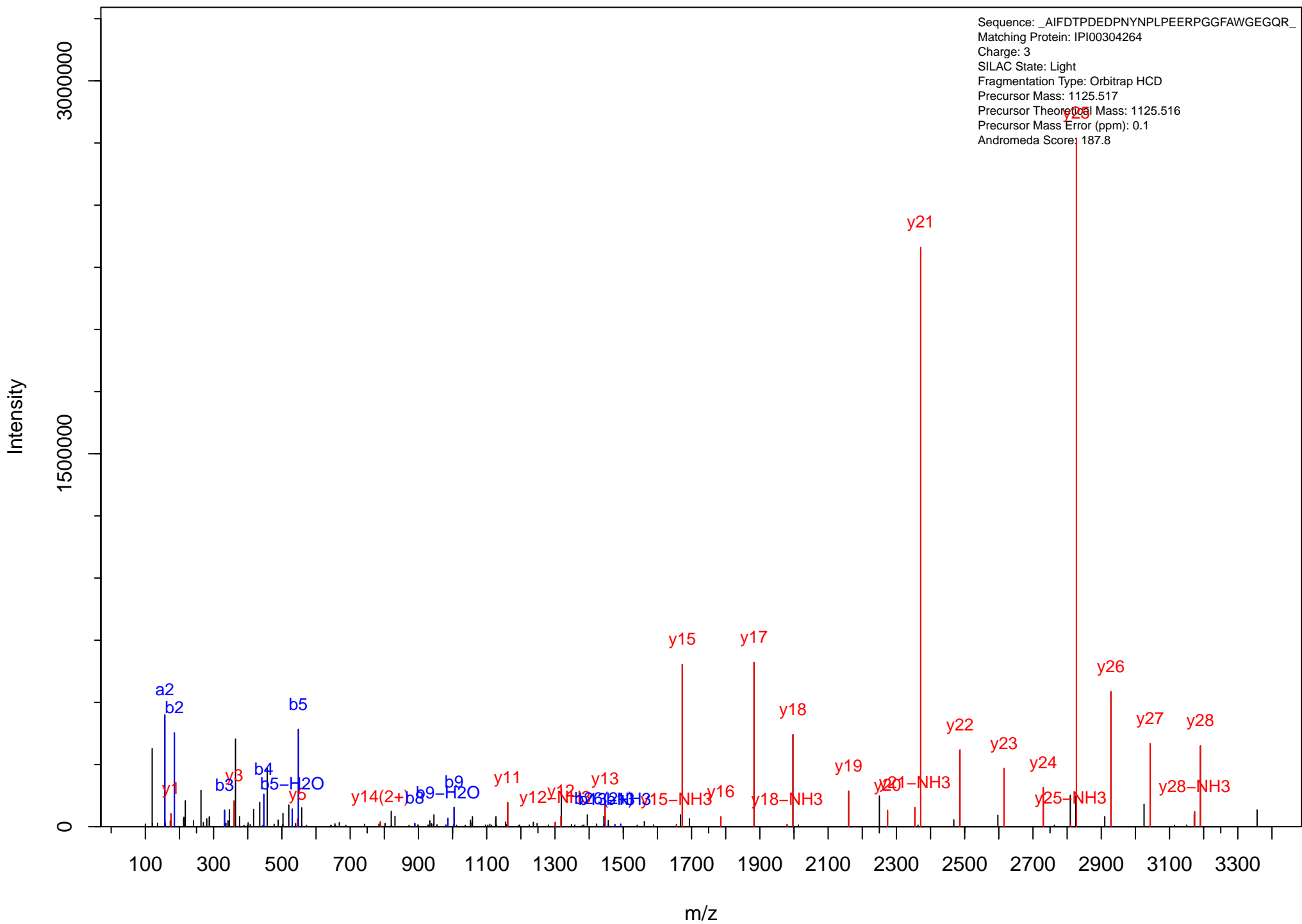
y12

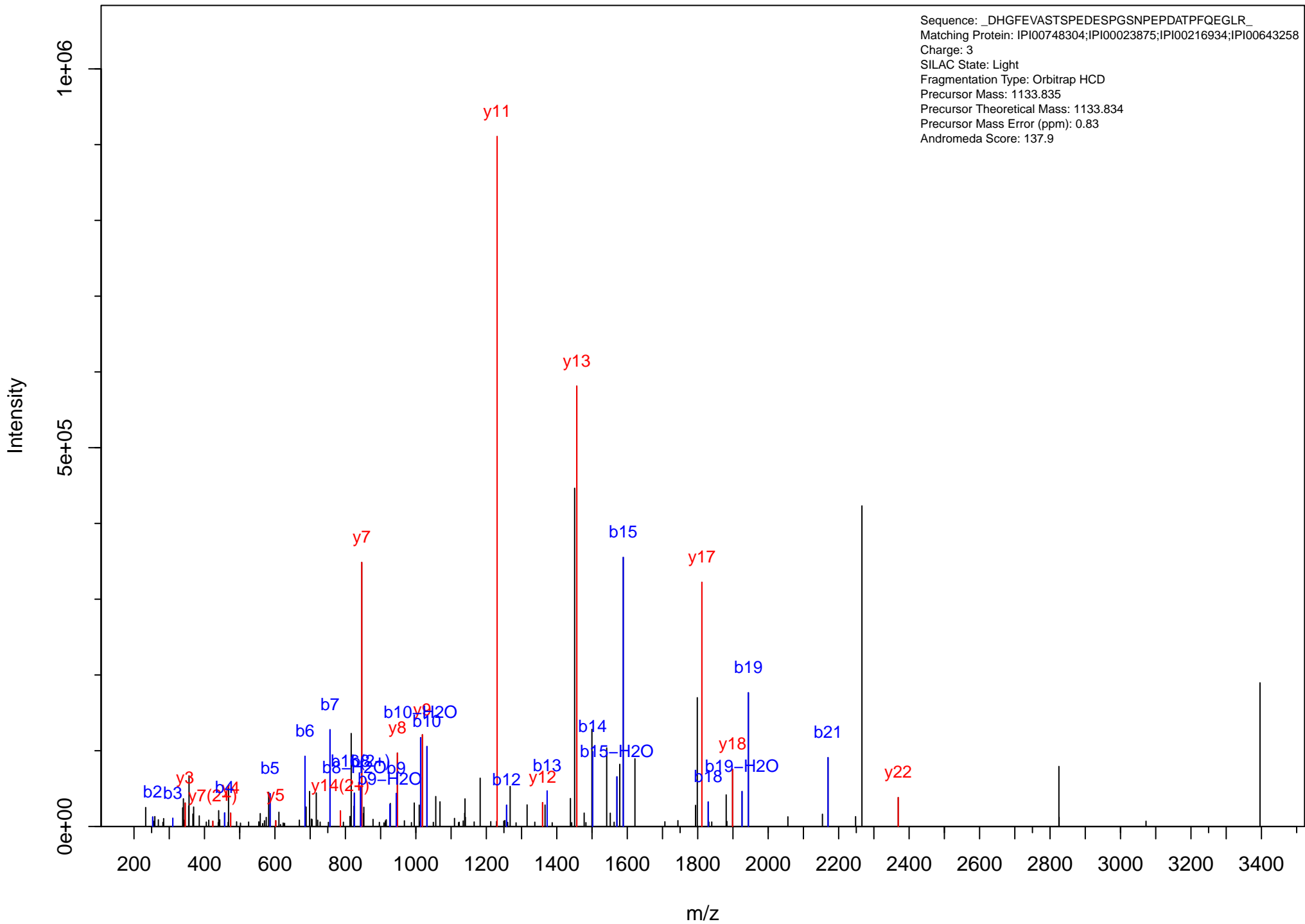
y10

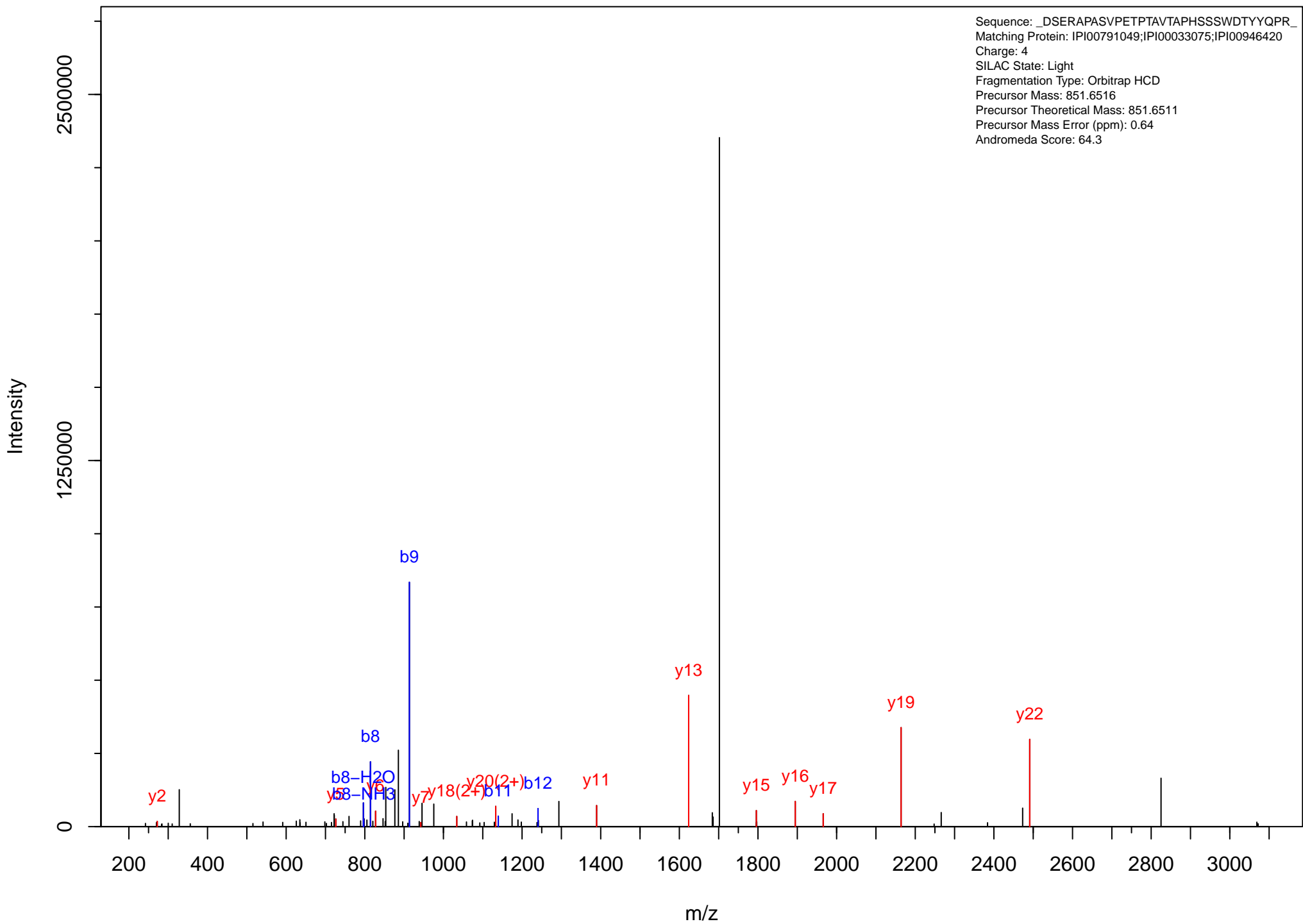


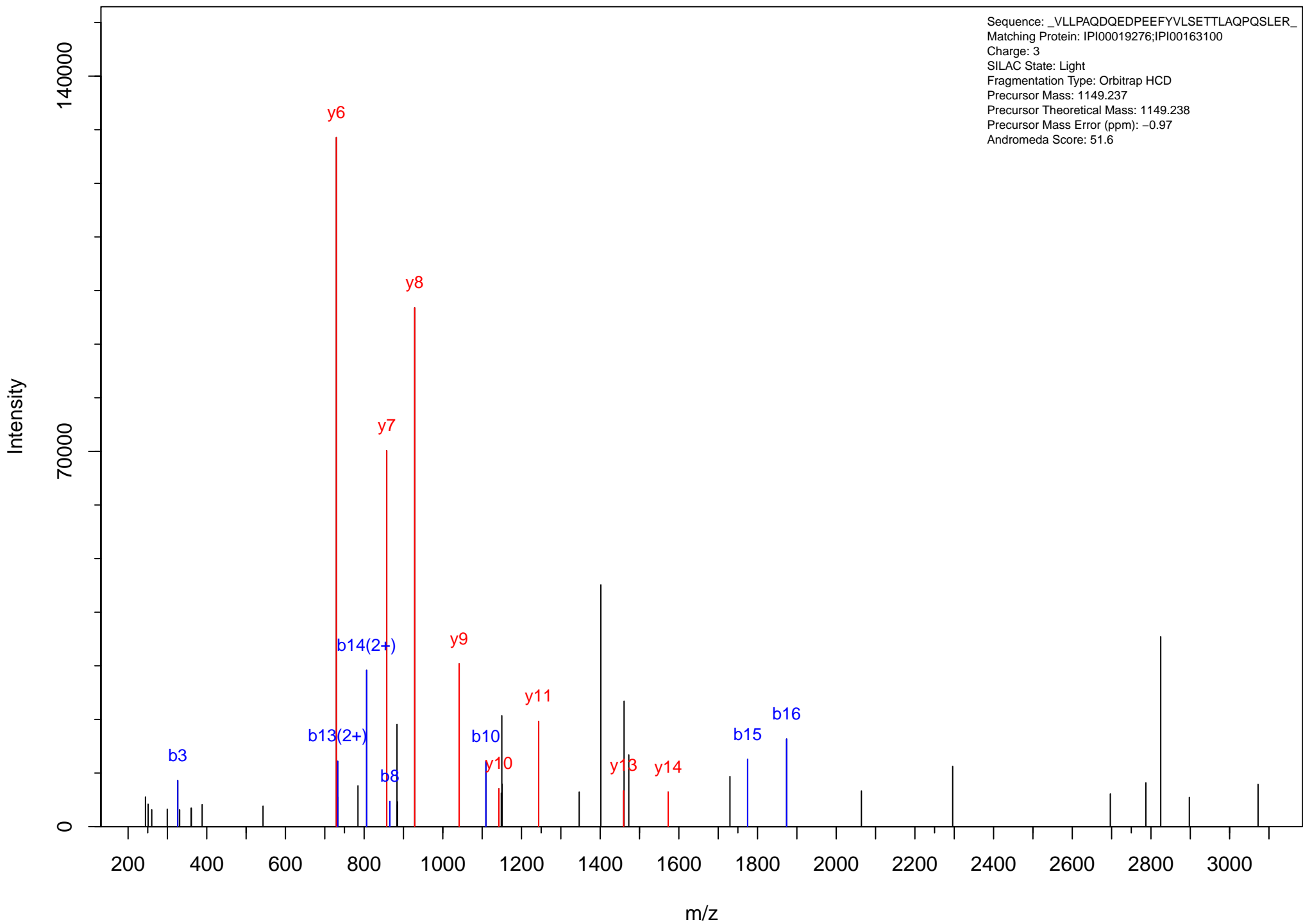


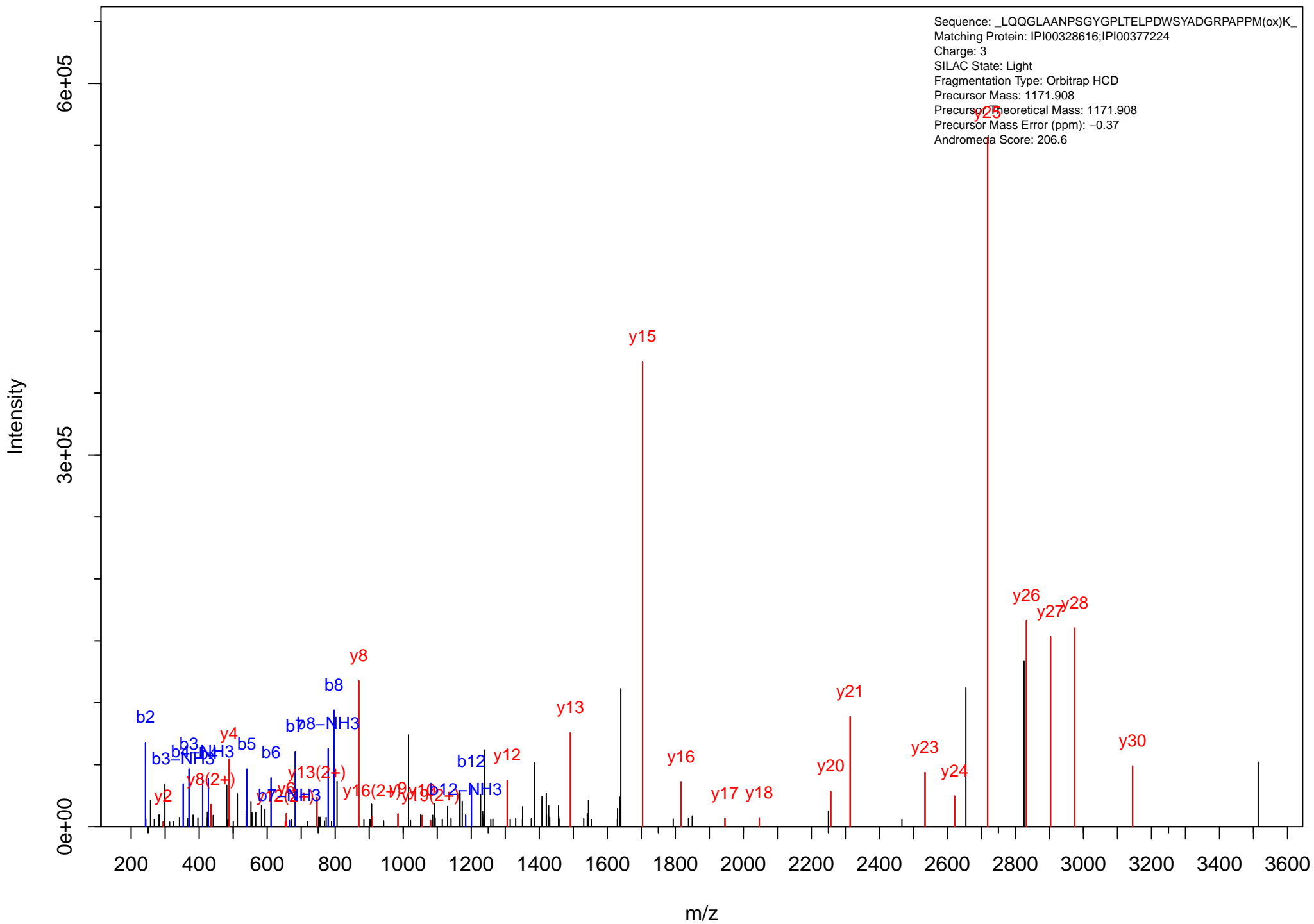


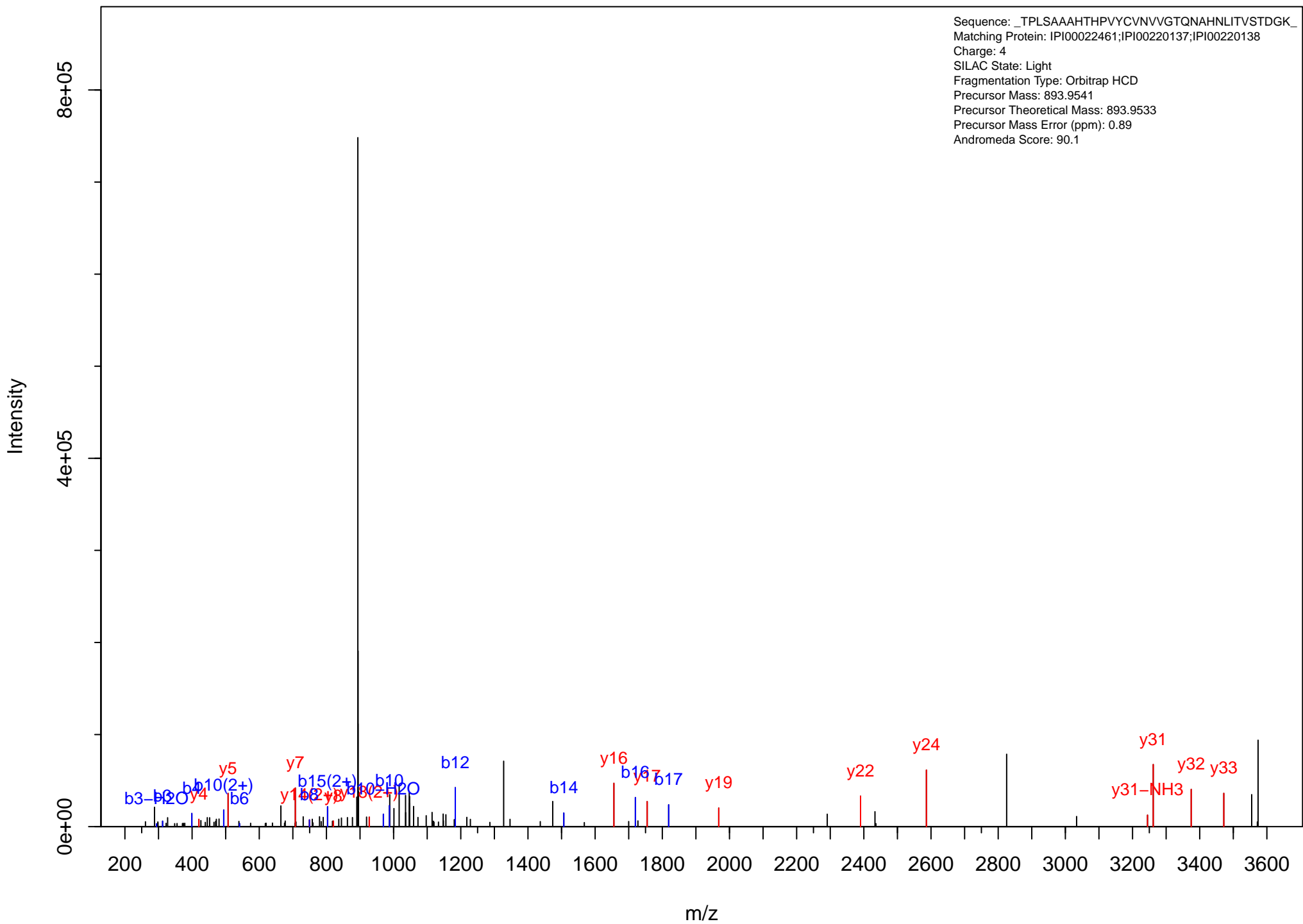


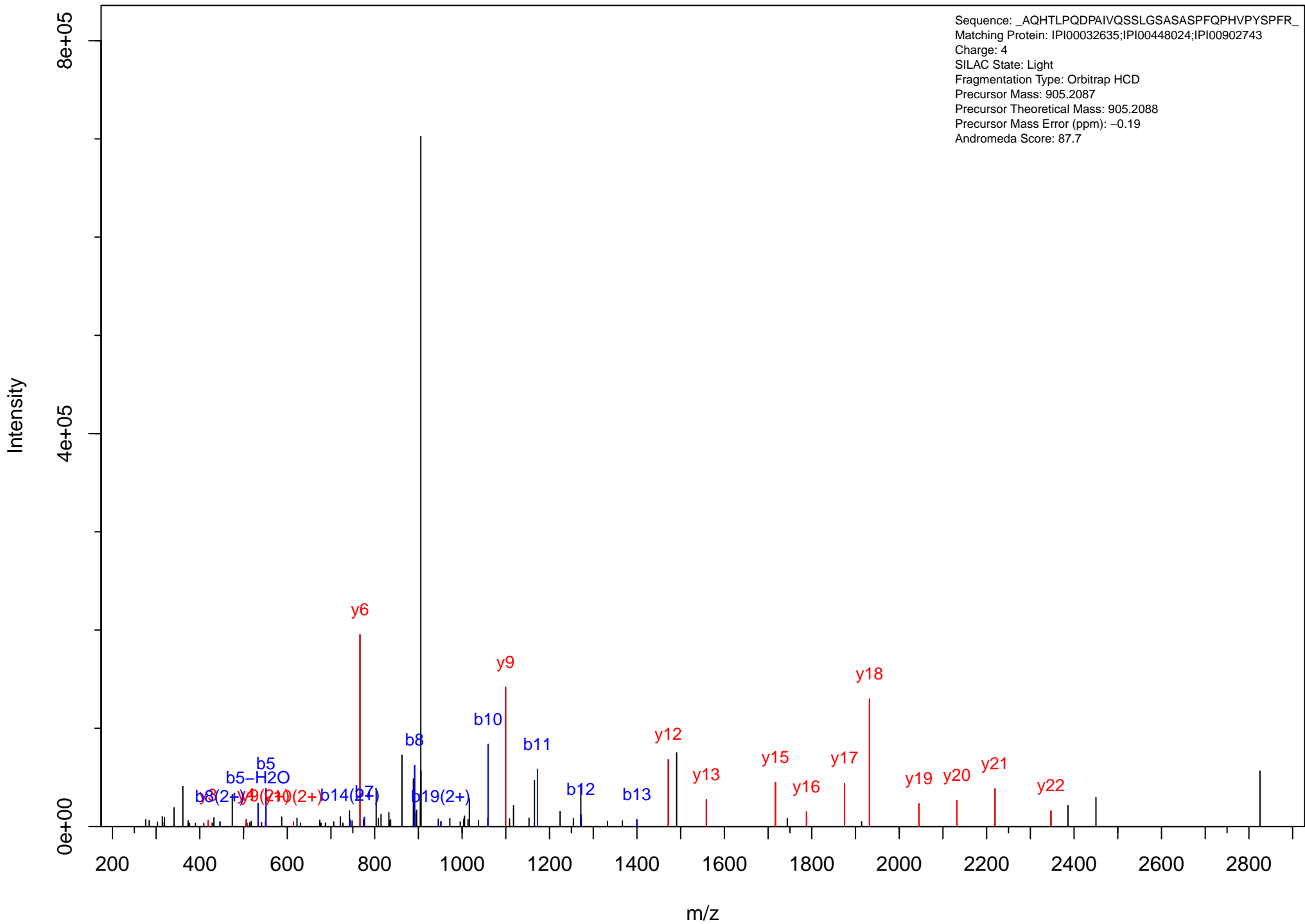


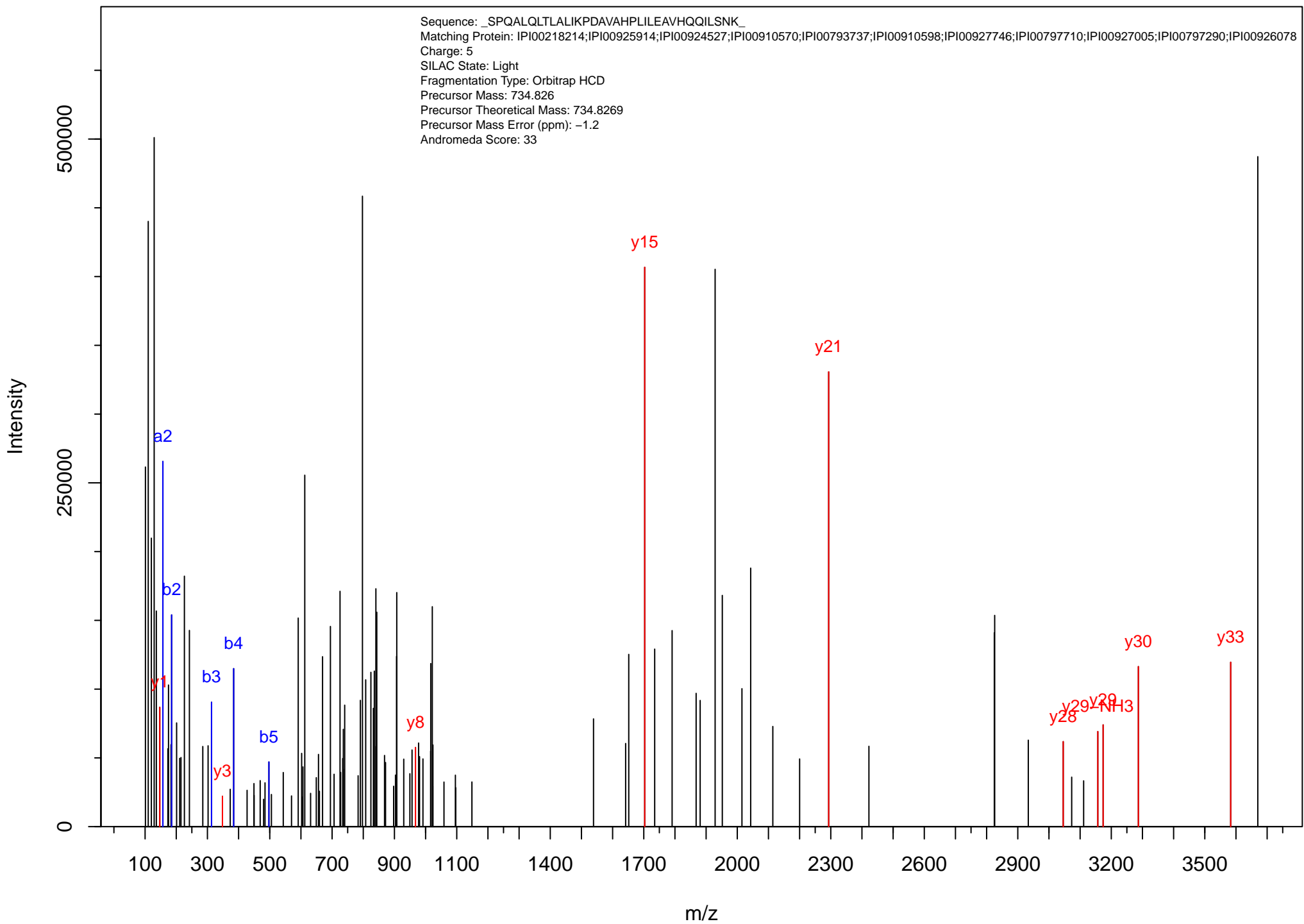


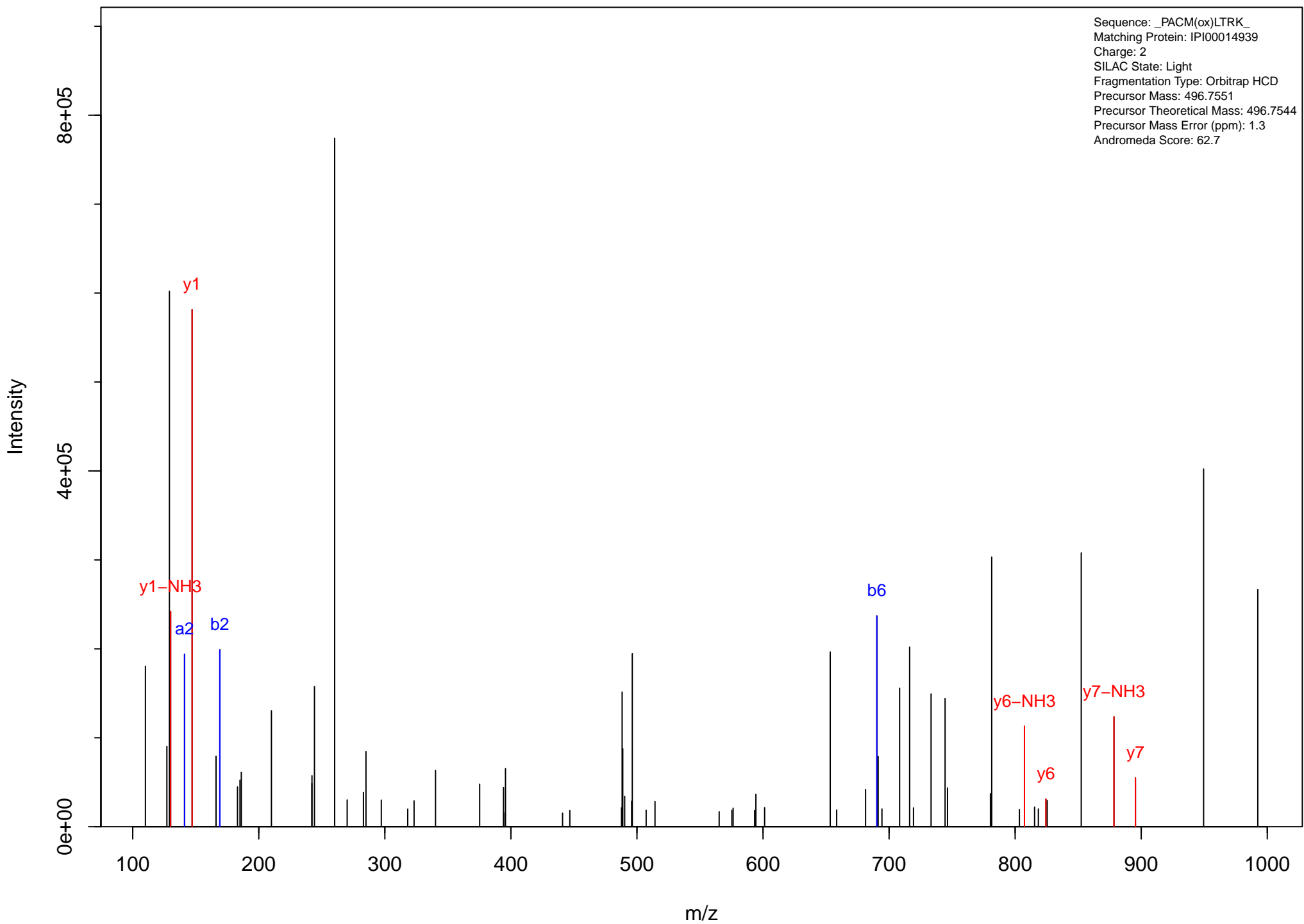


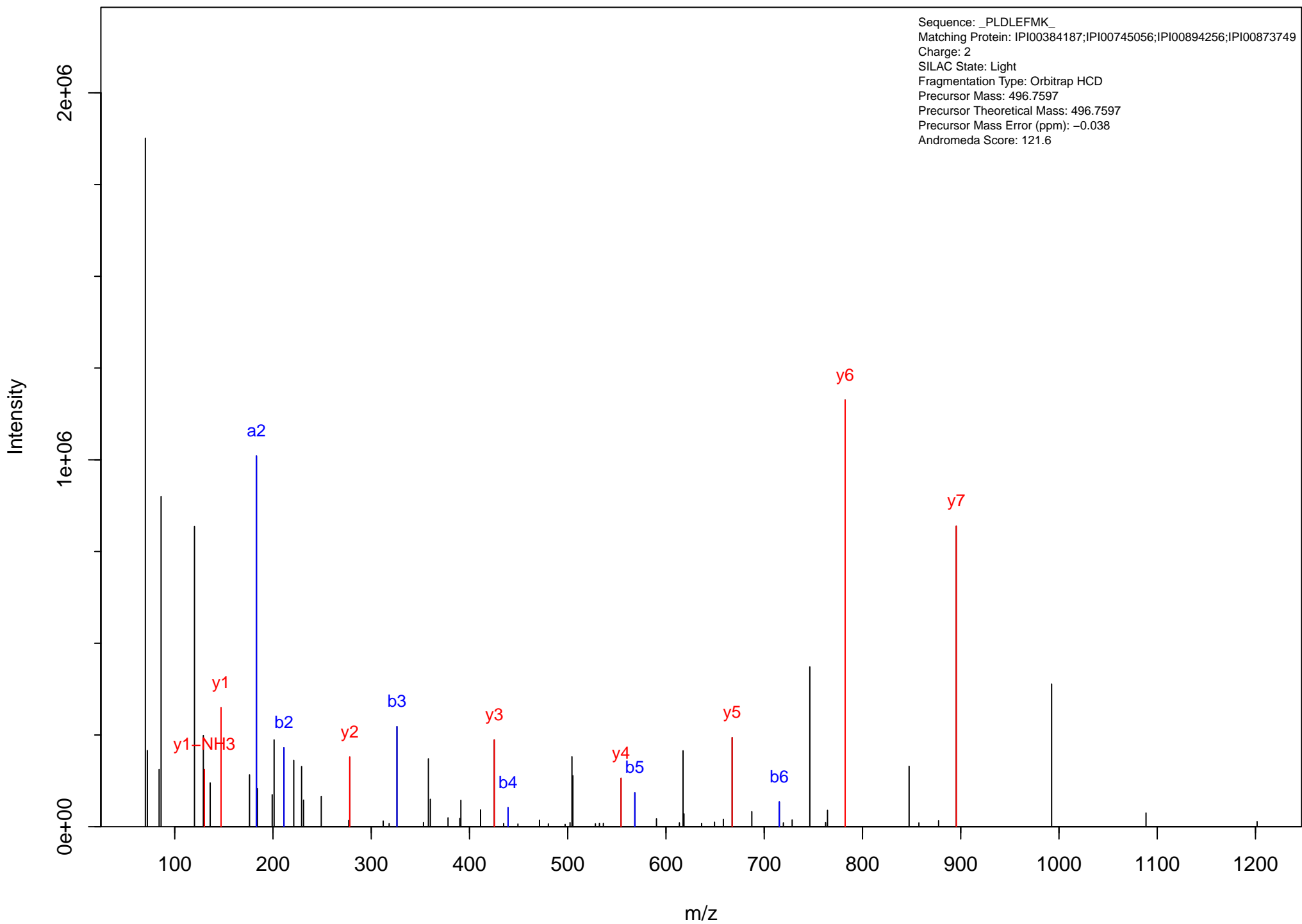


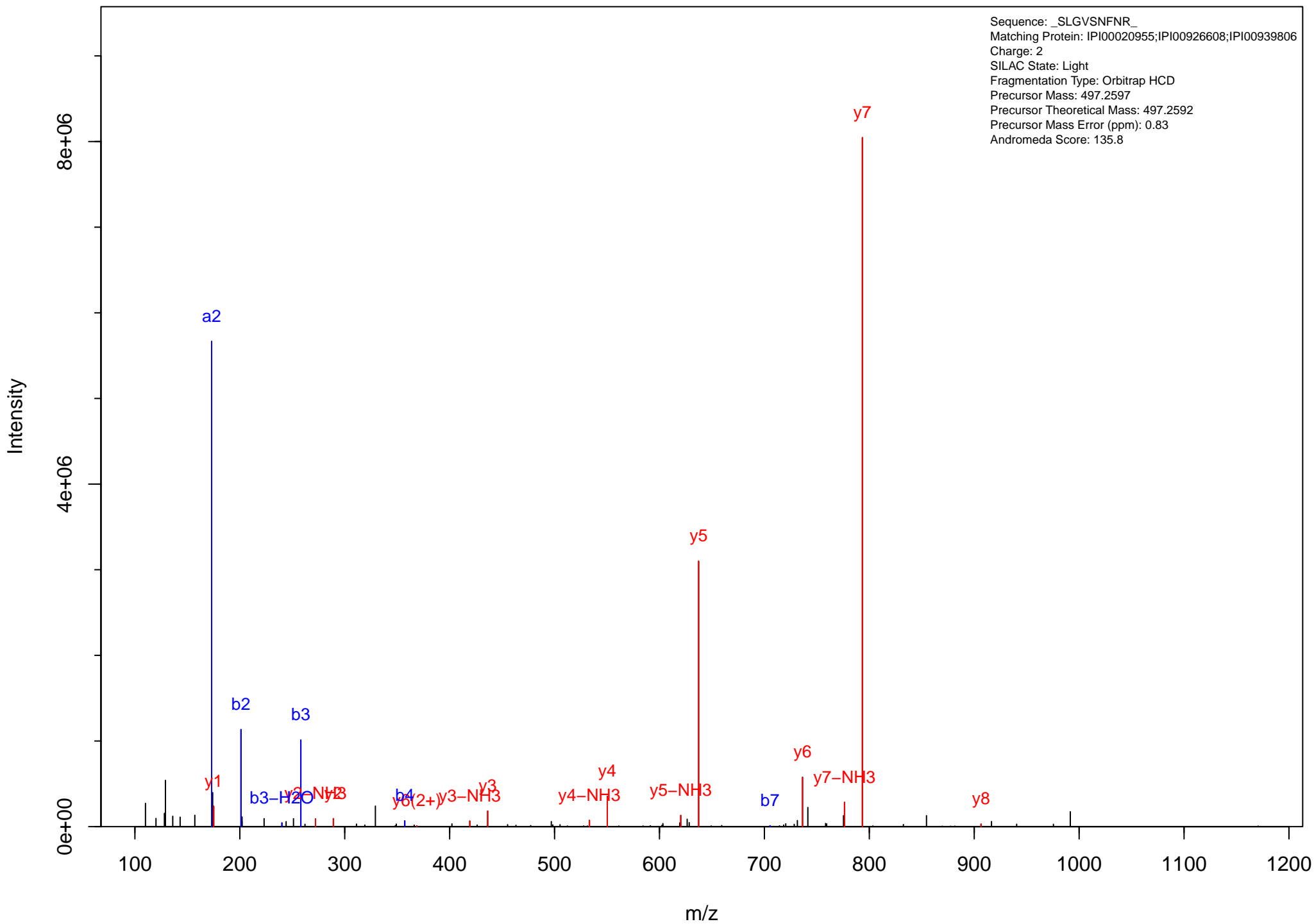


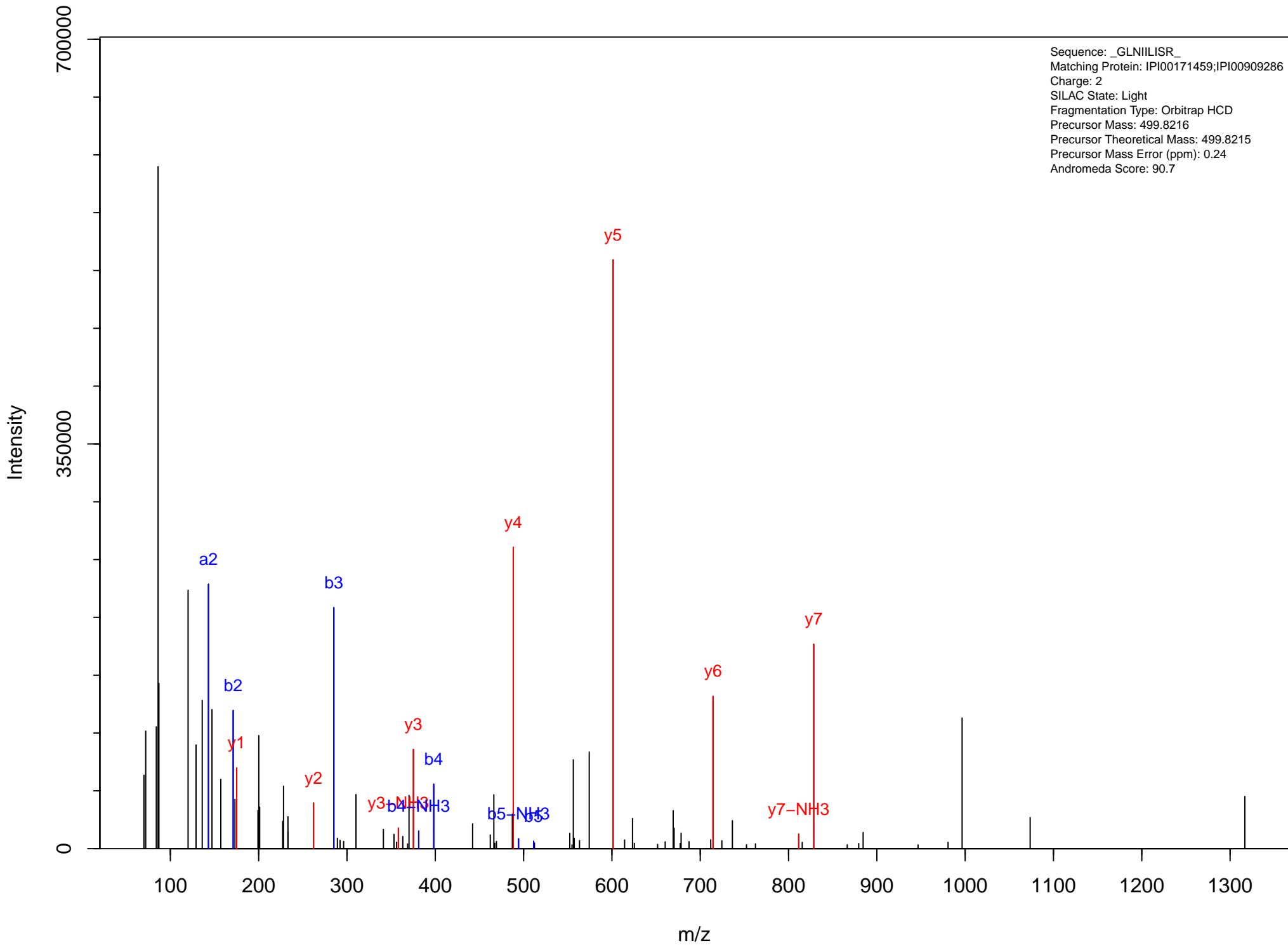


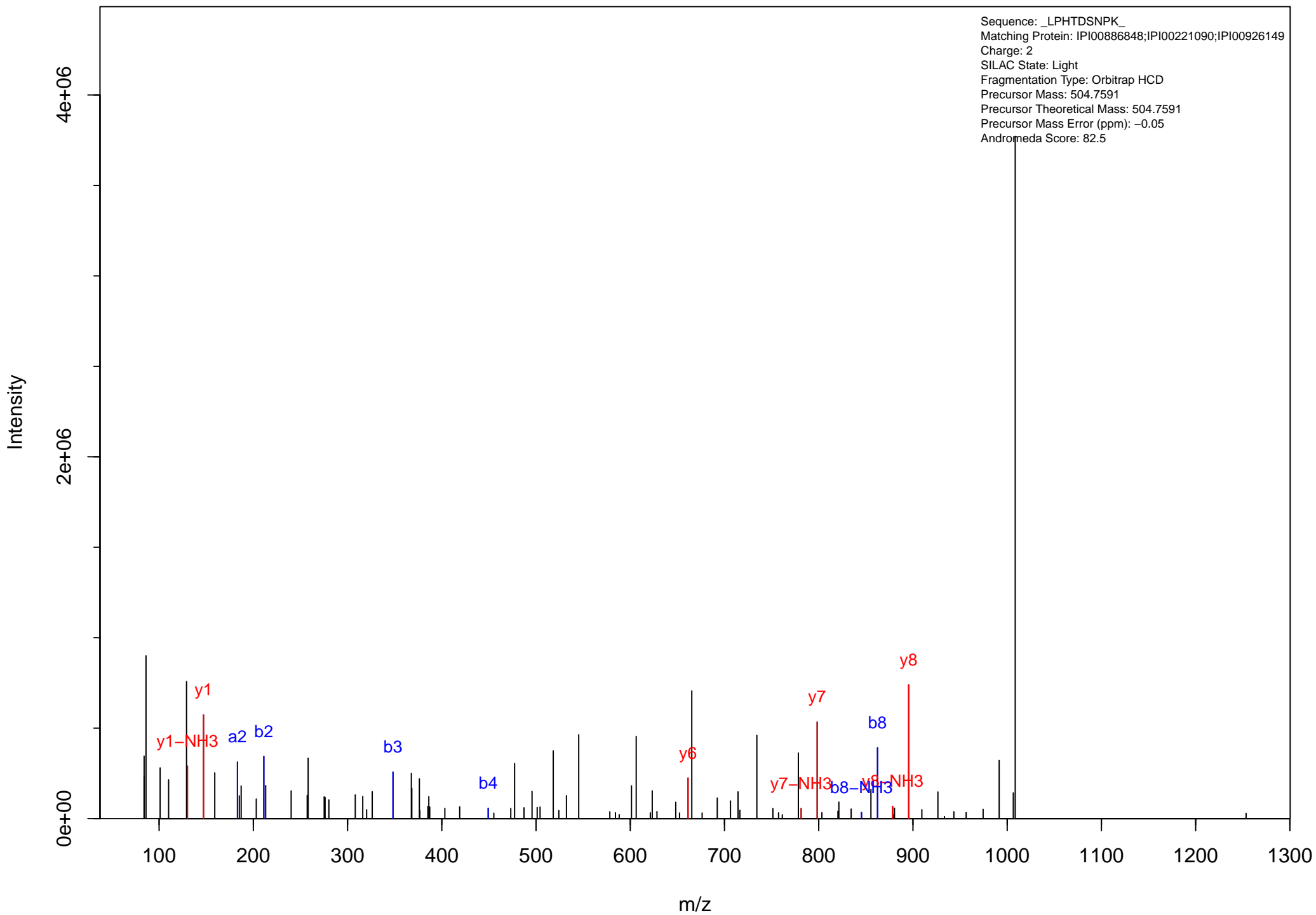


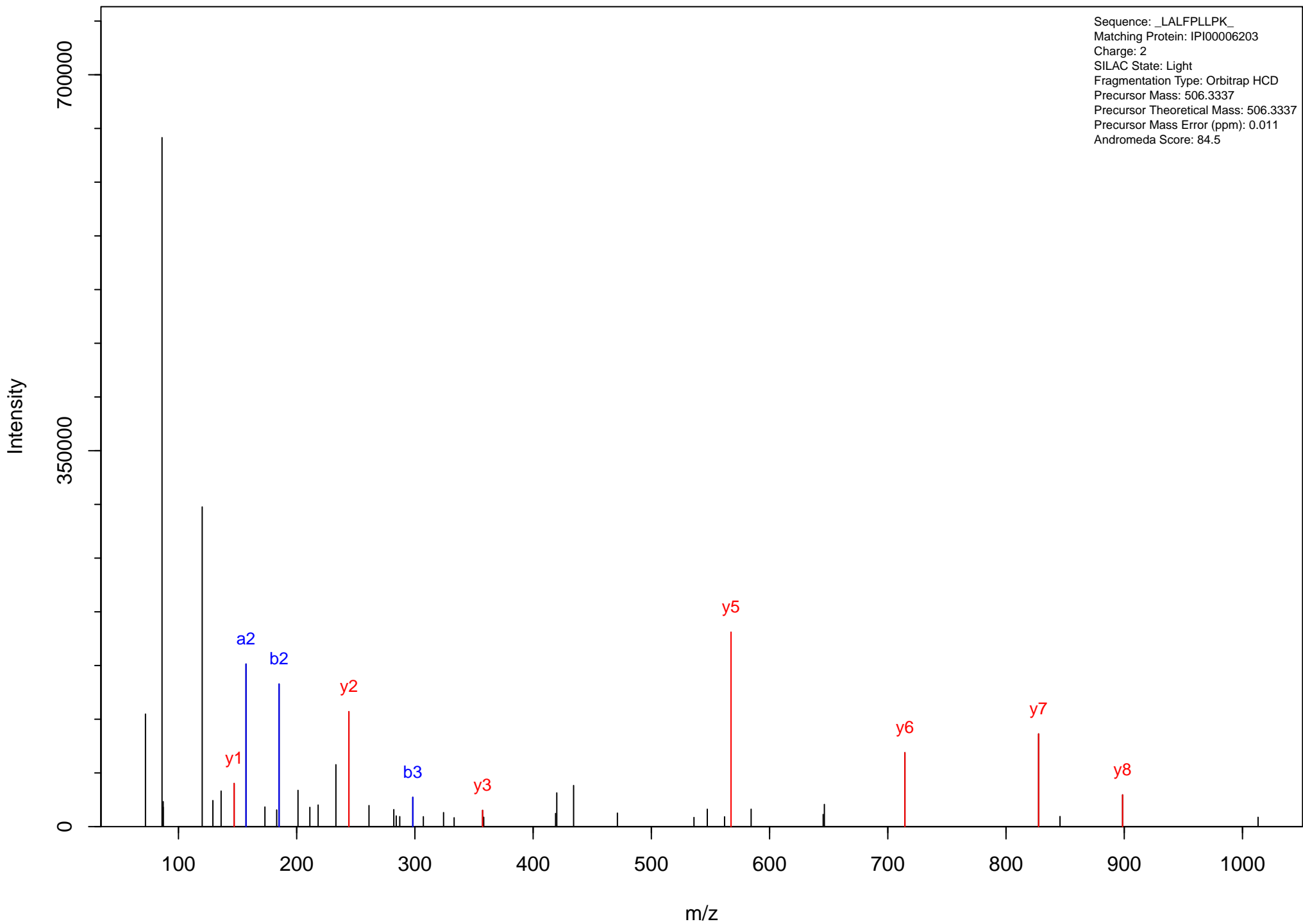


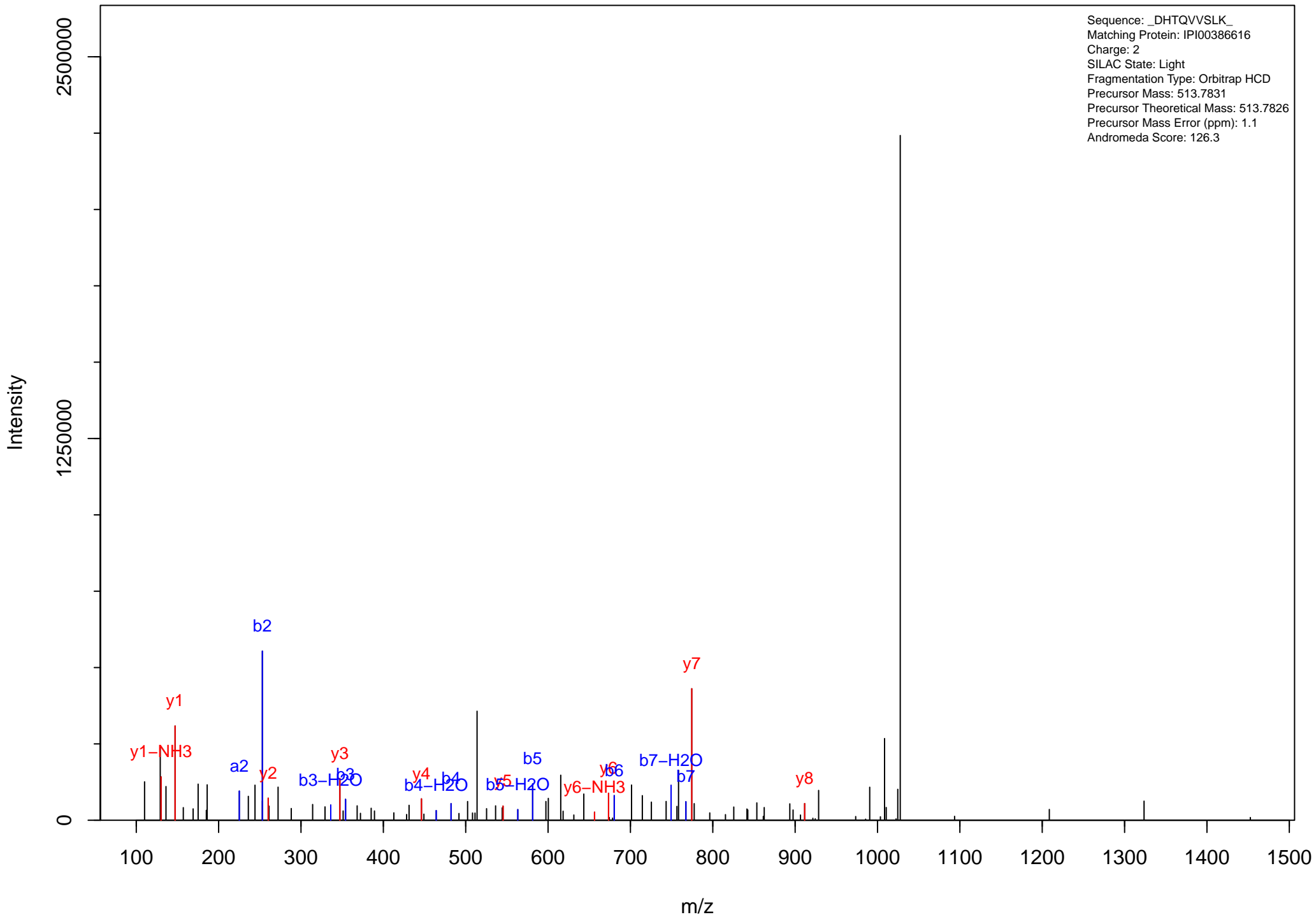


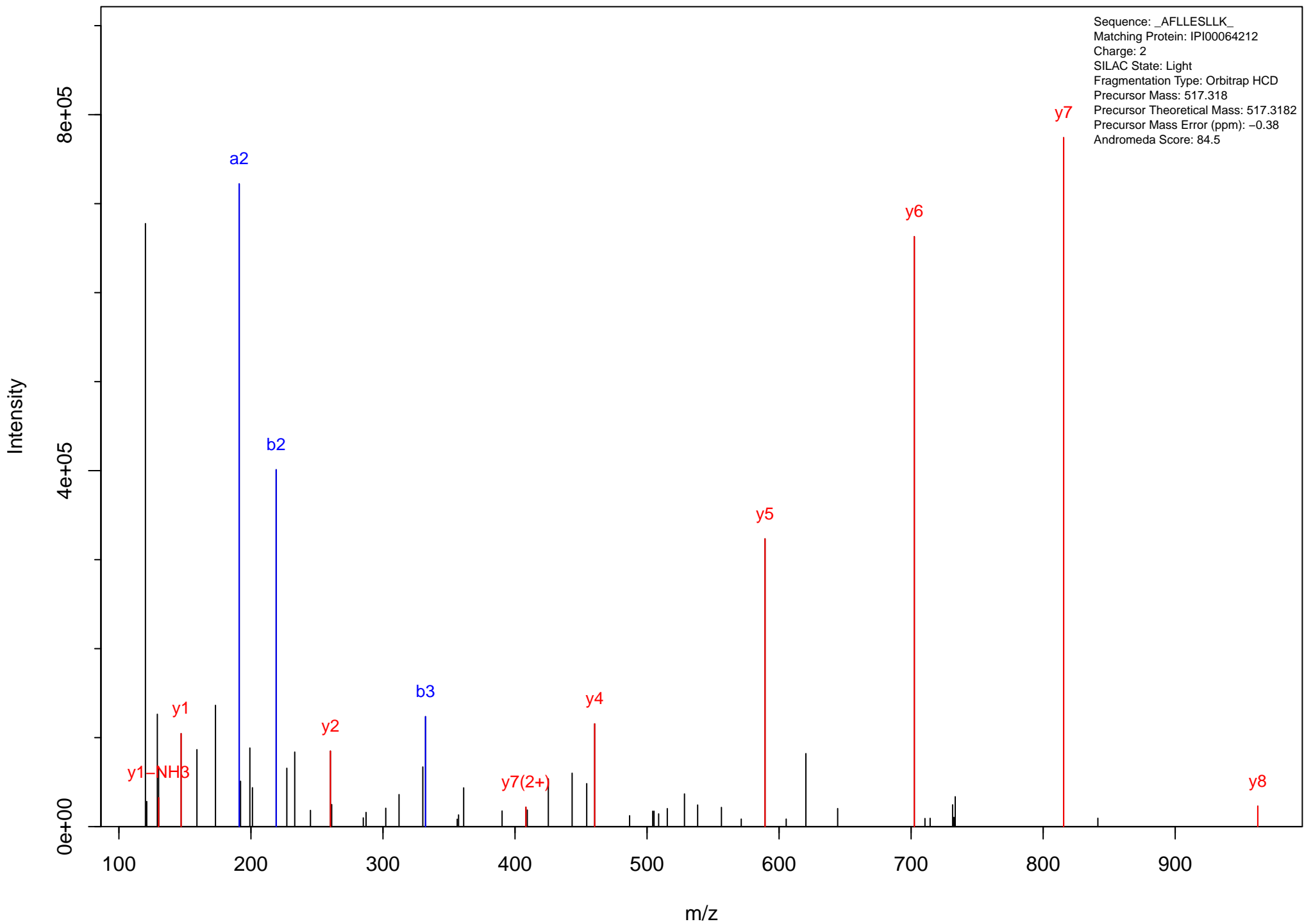


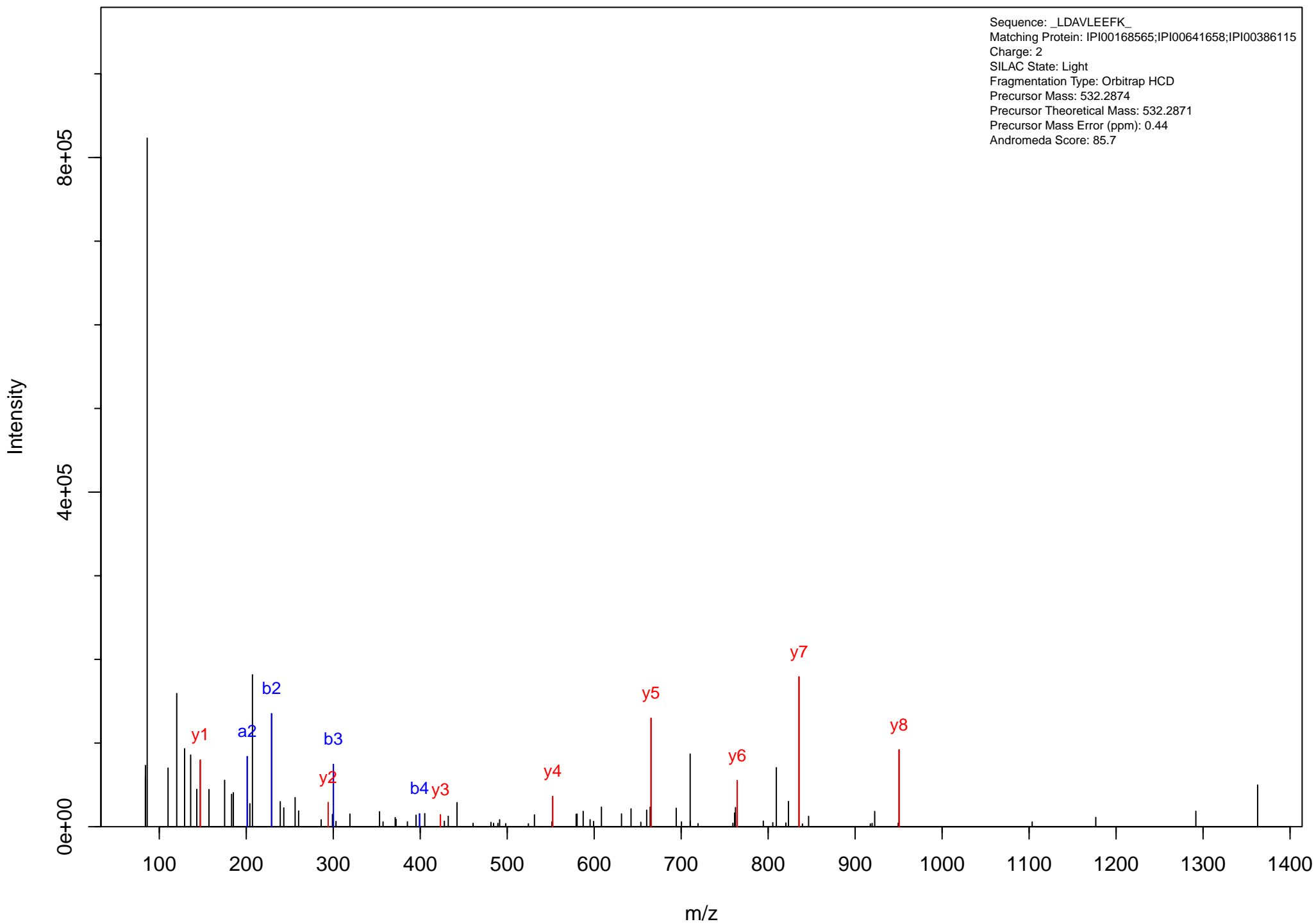


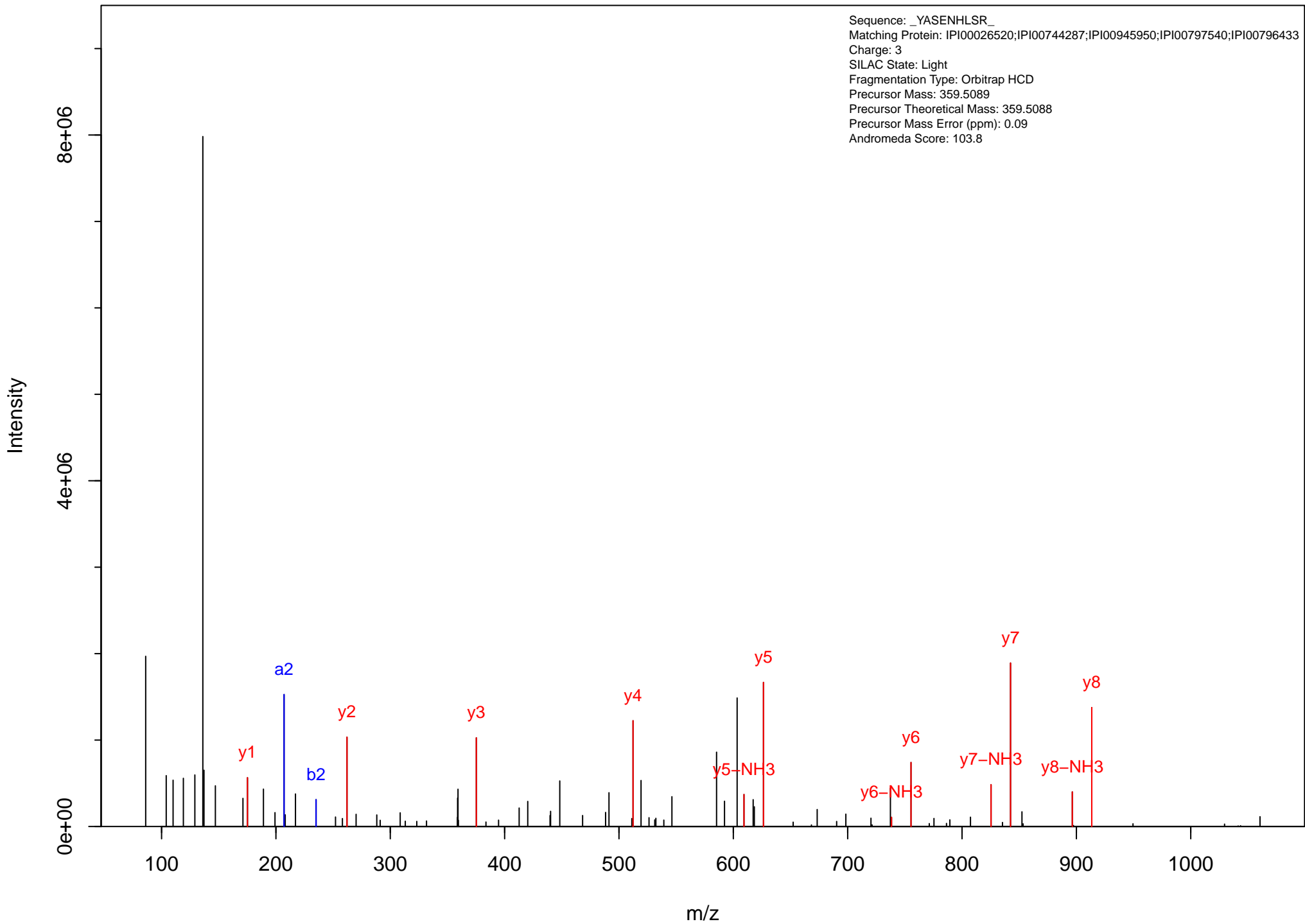


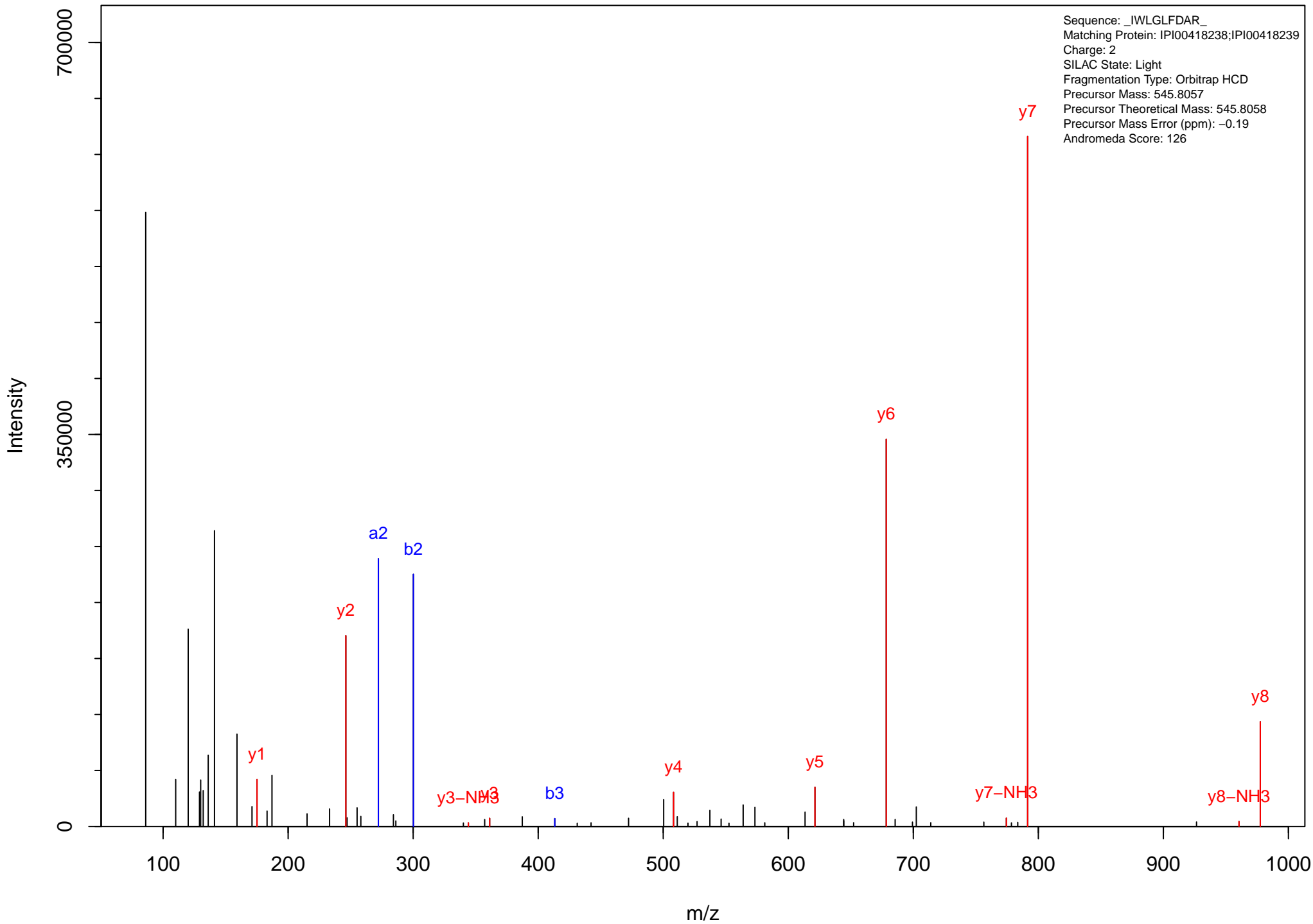


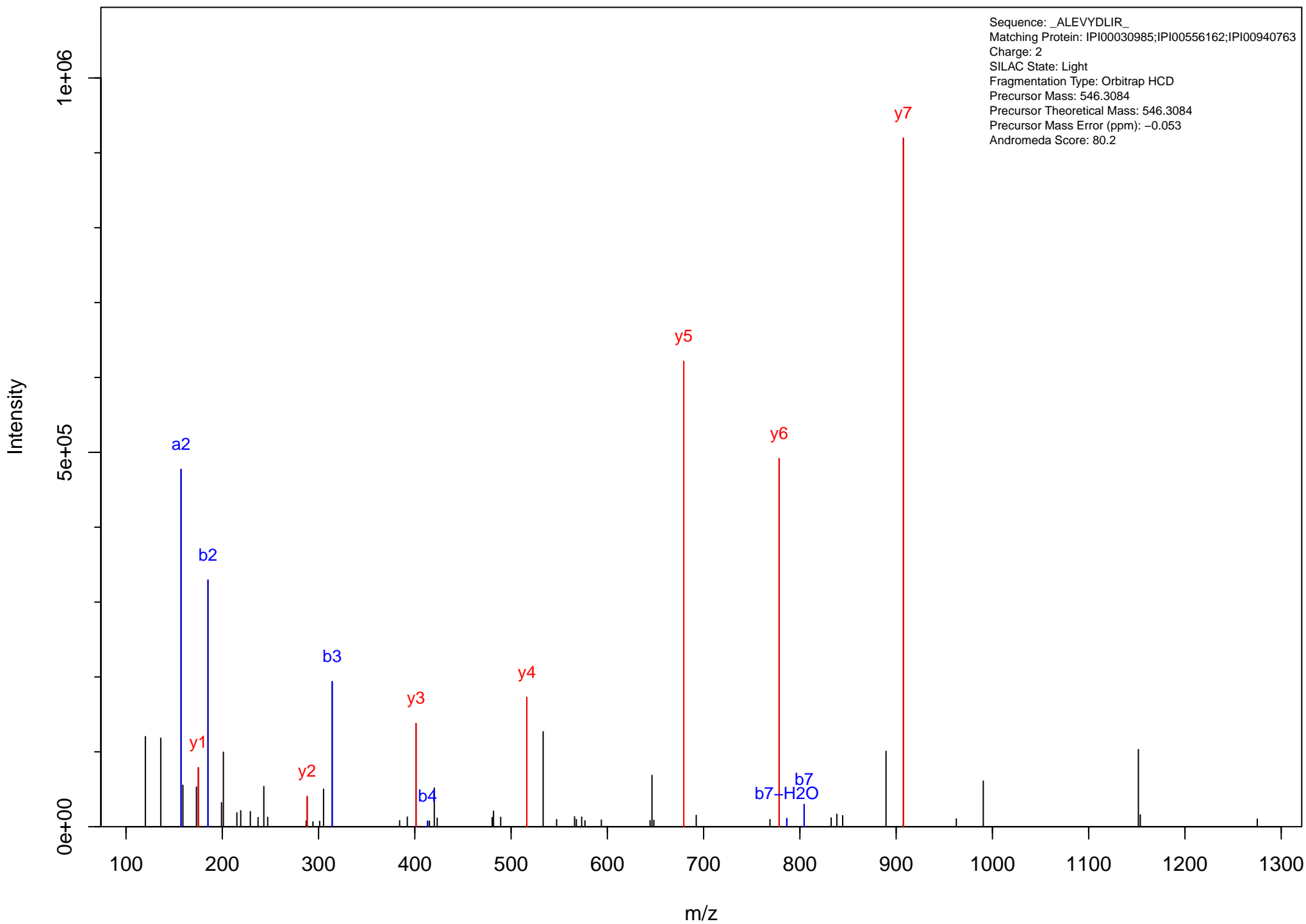


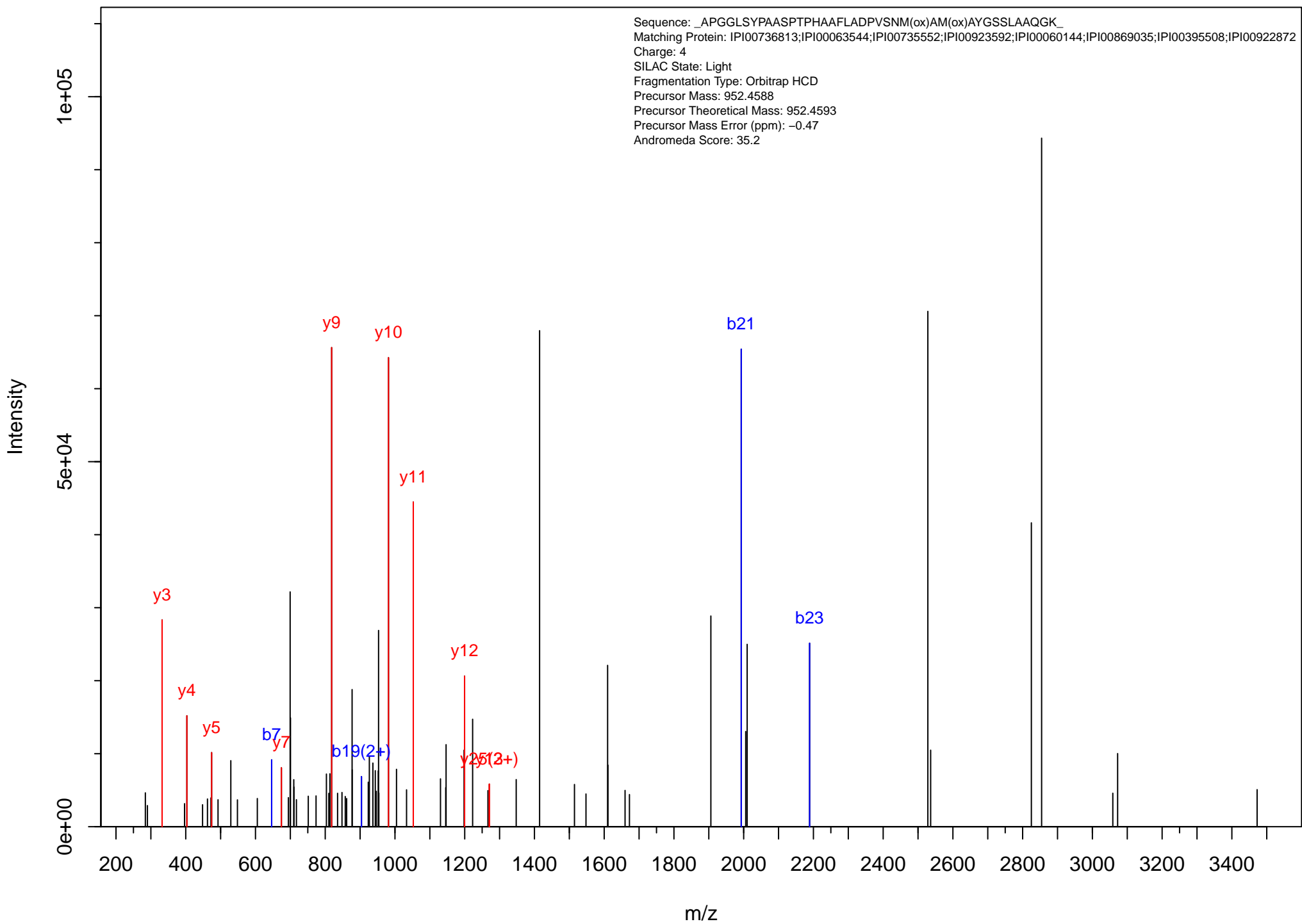


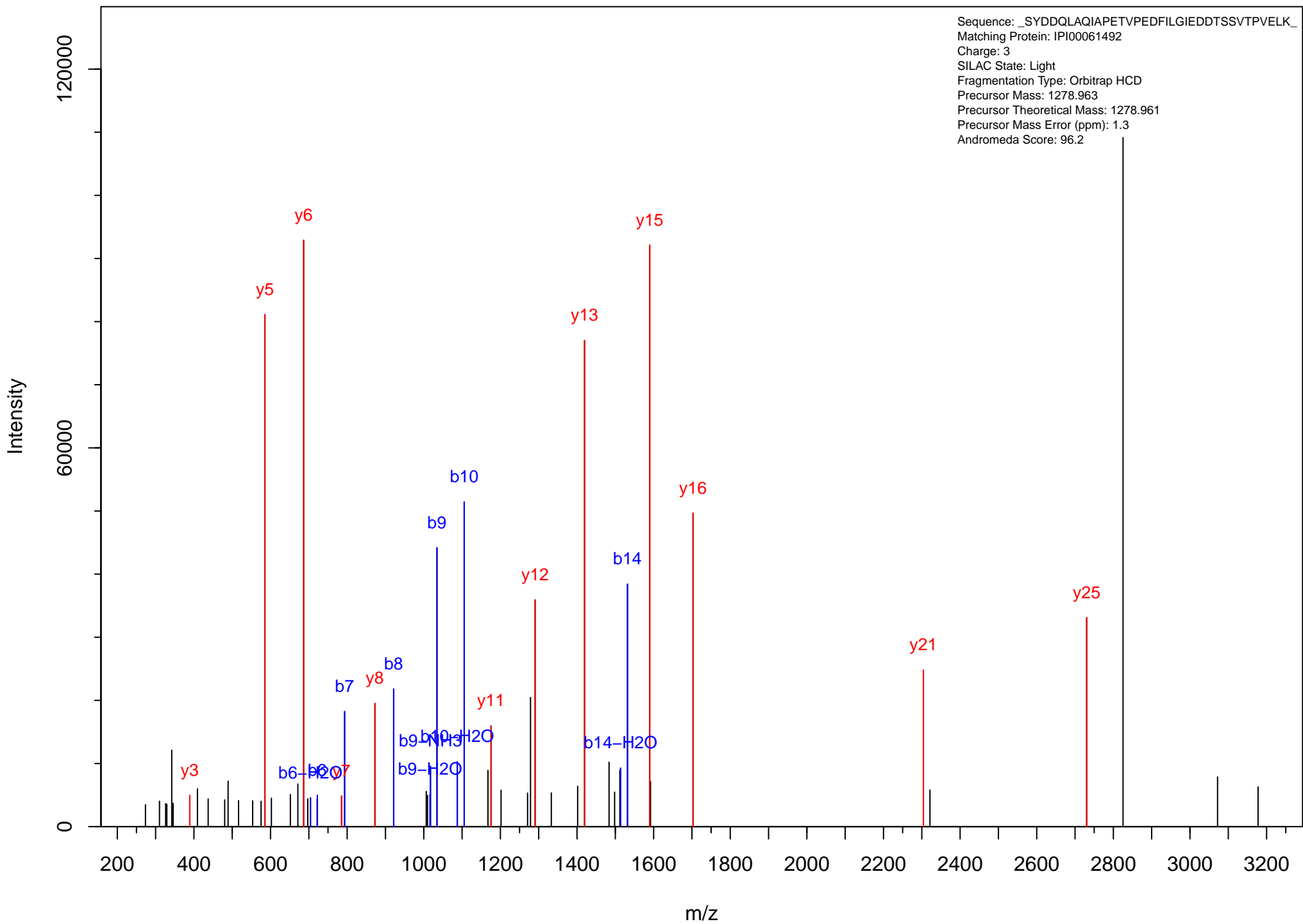


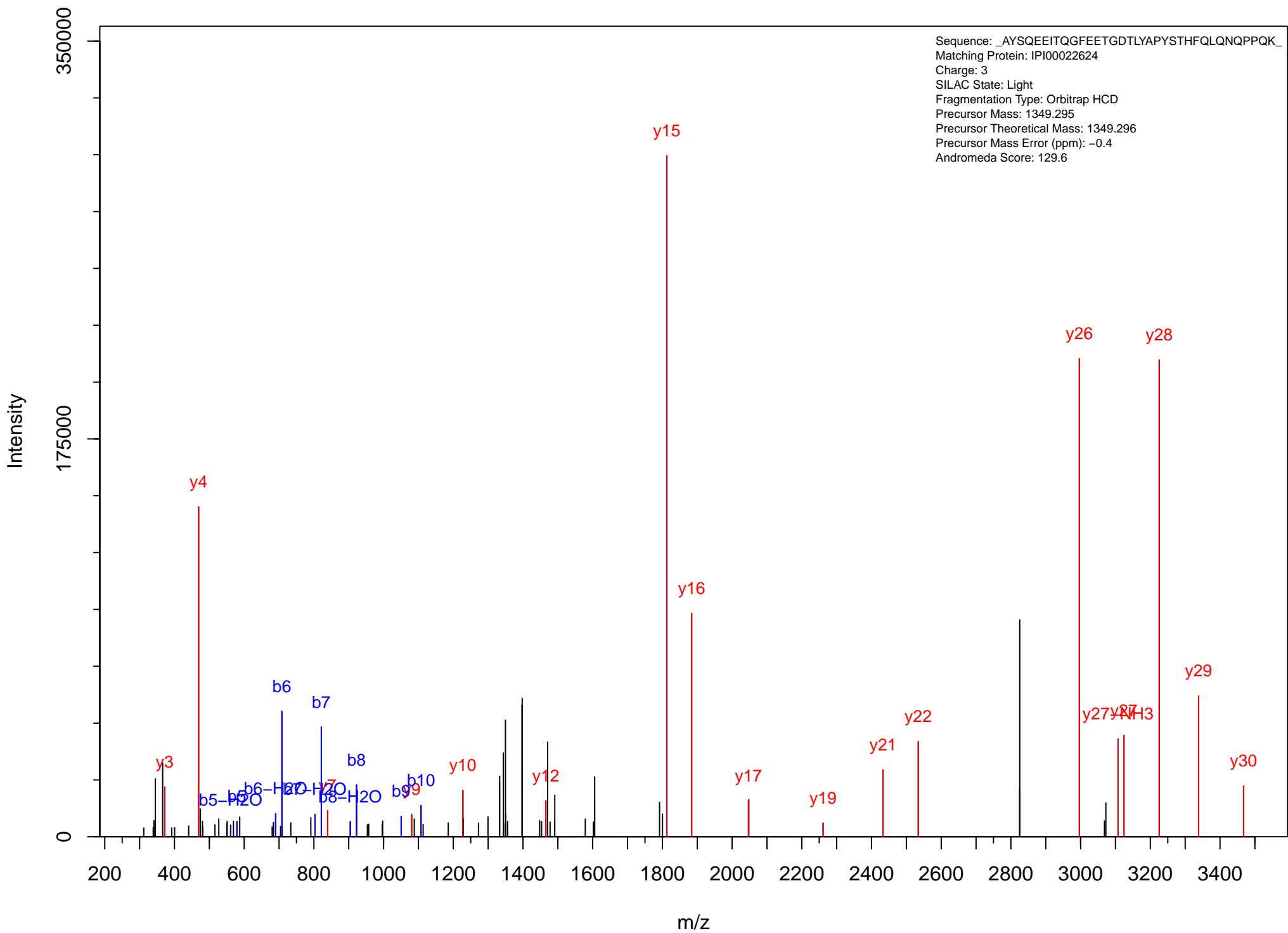


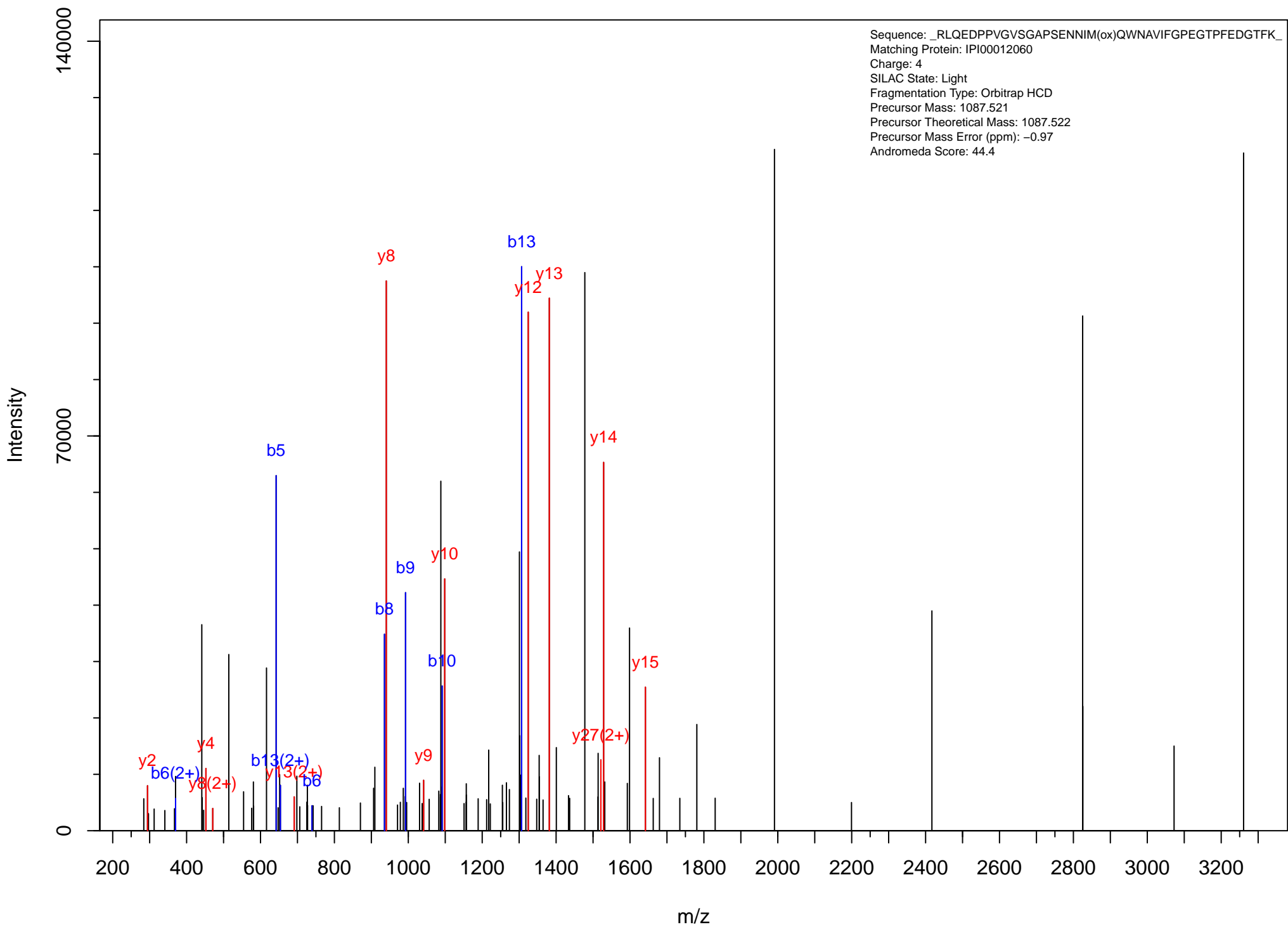


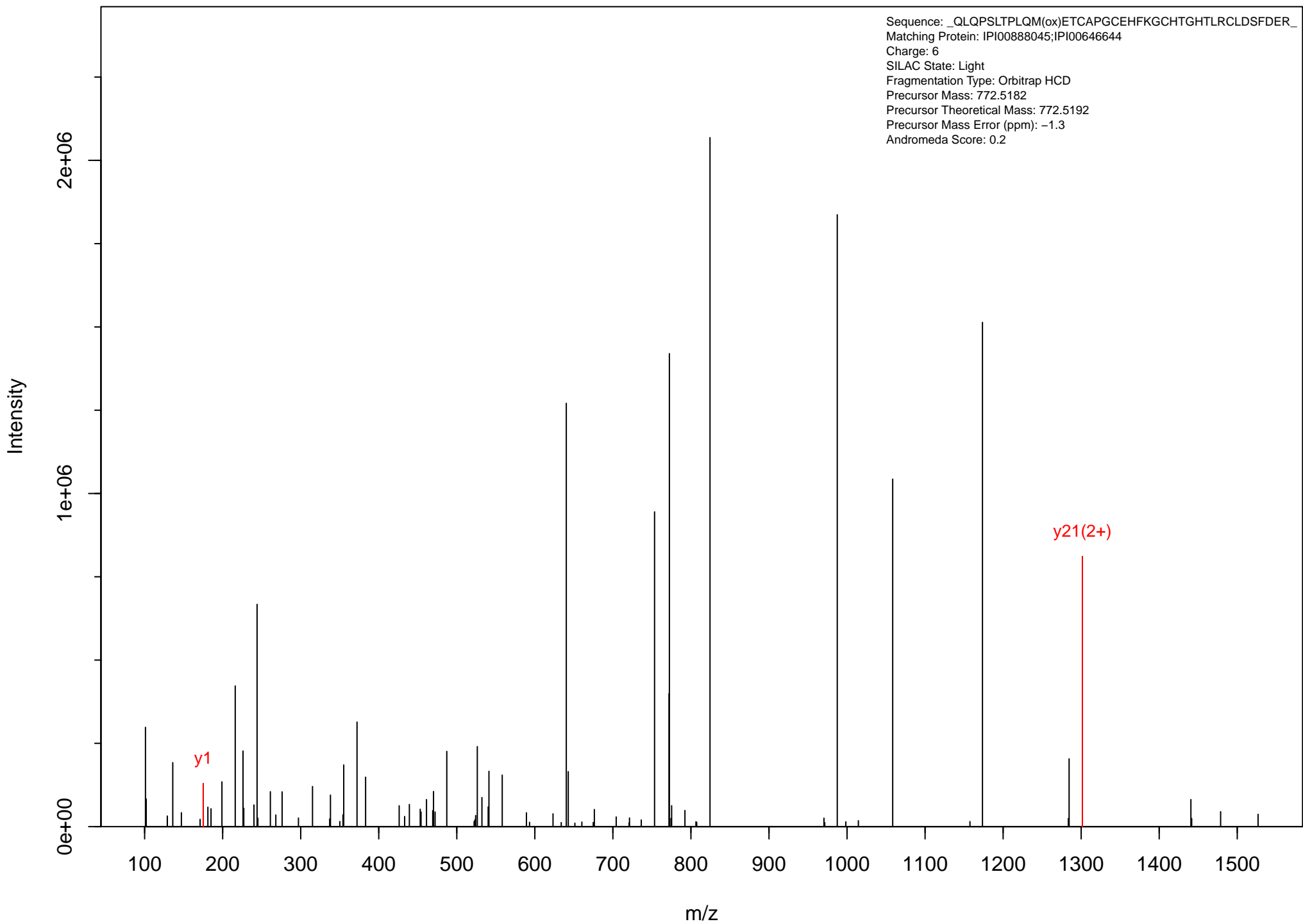


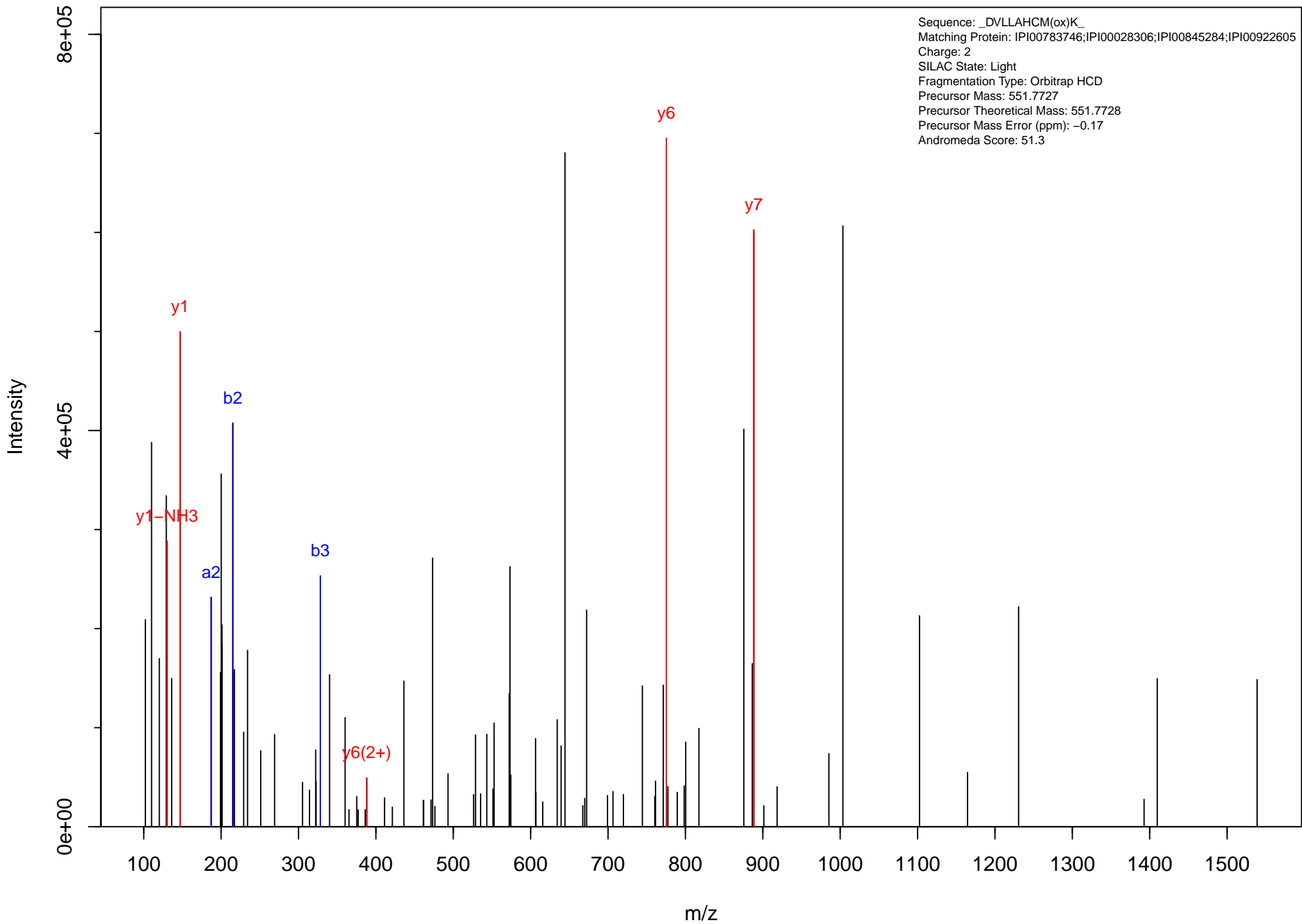


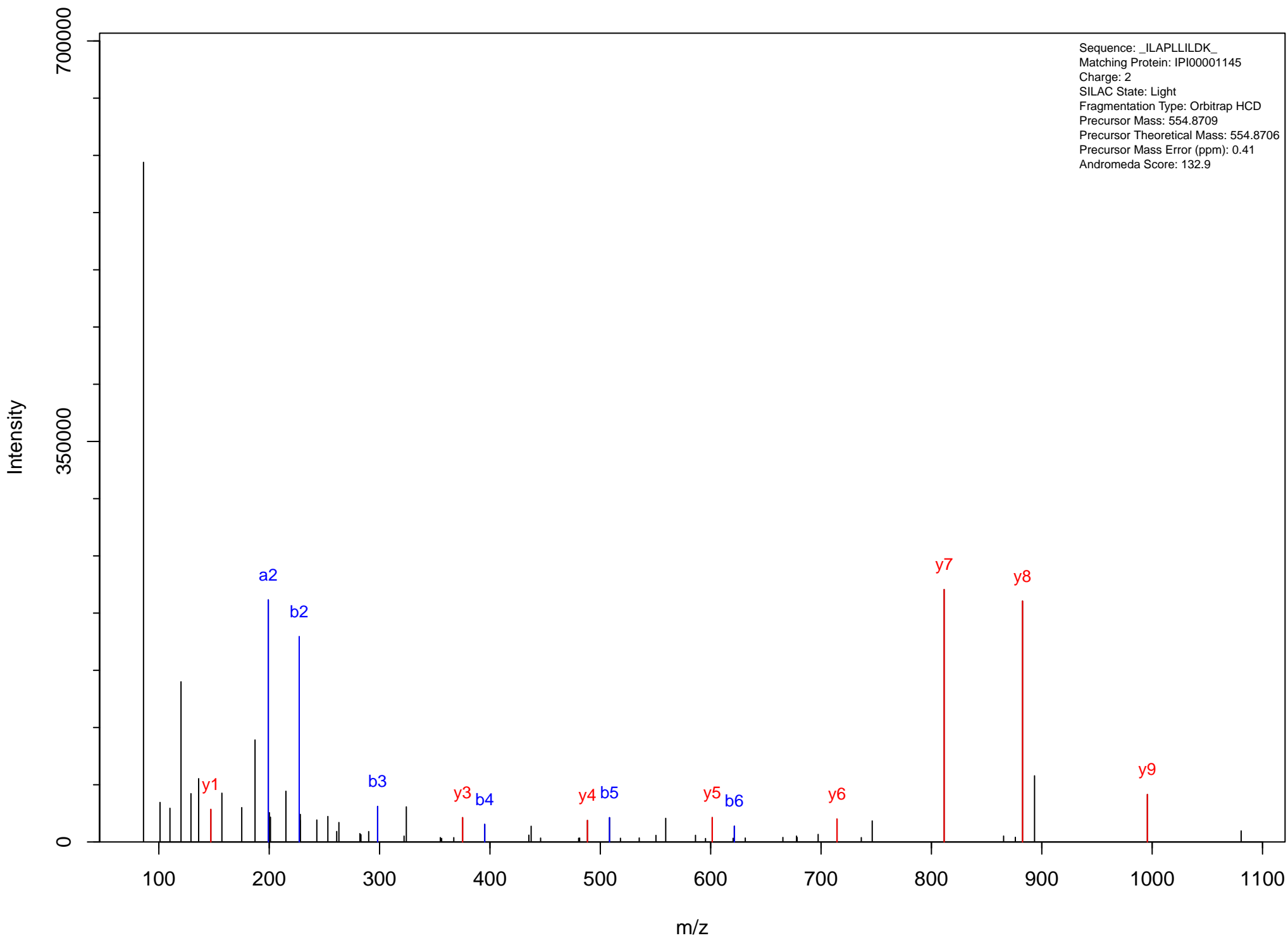


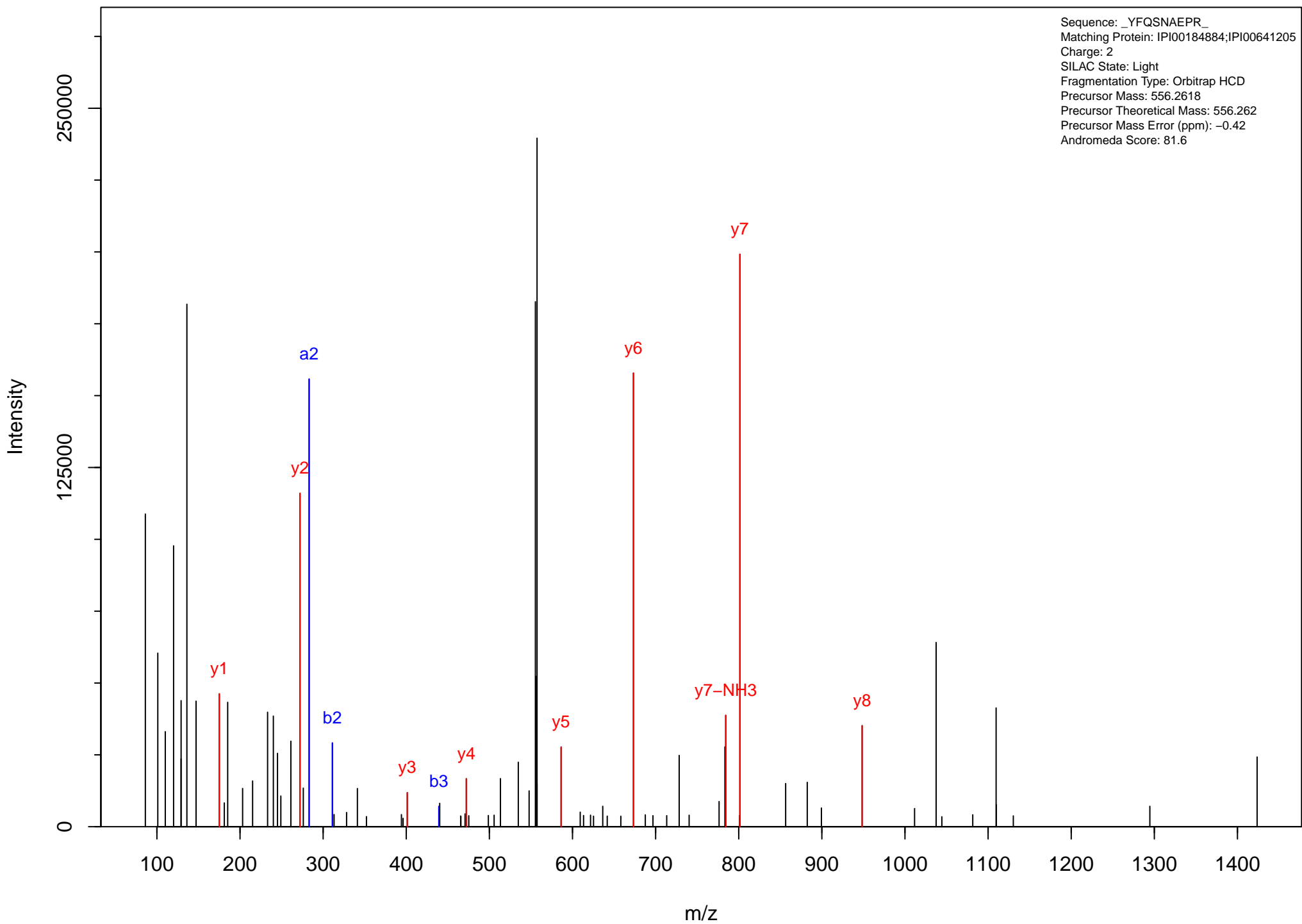


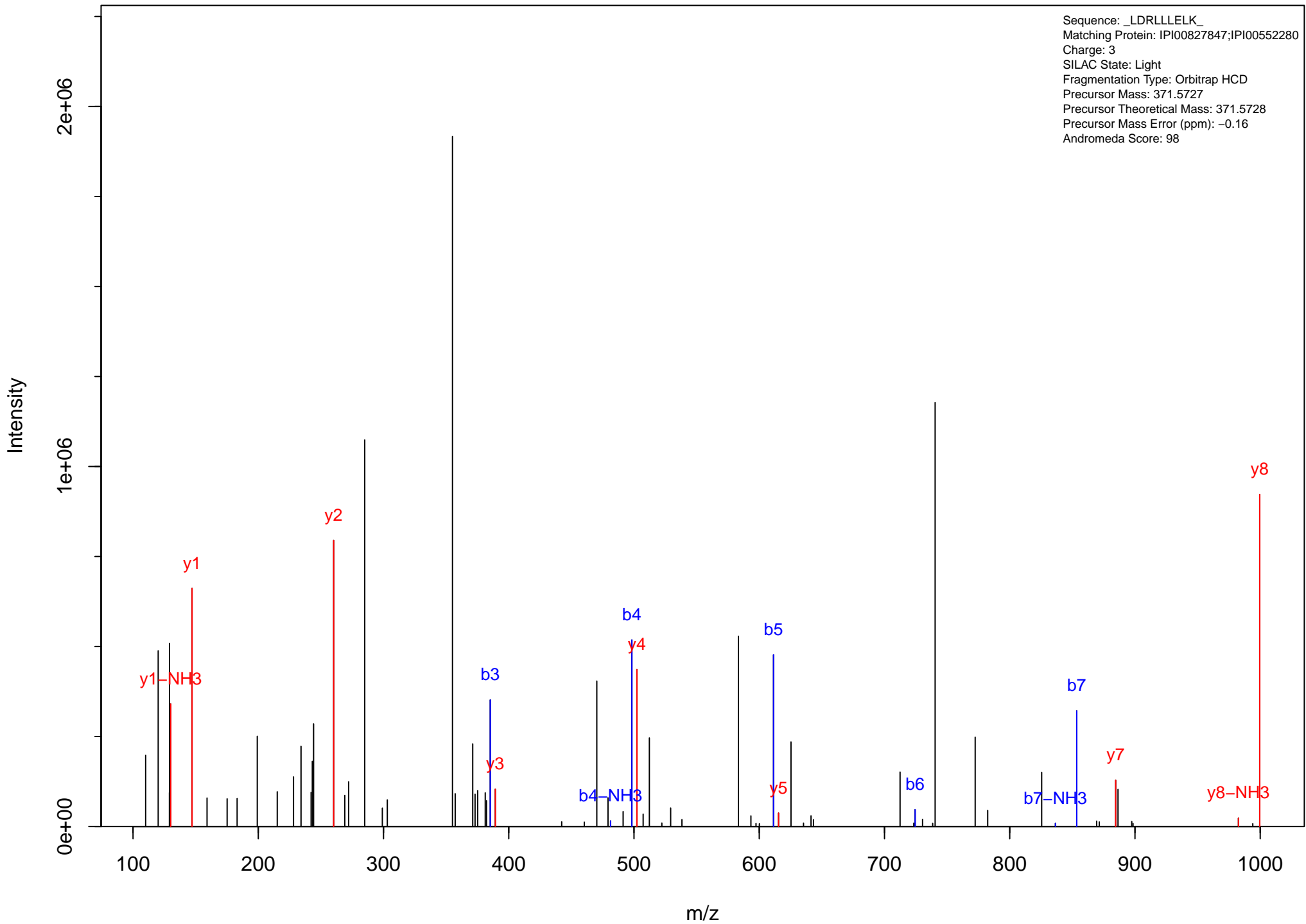


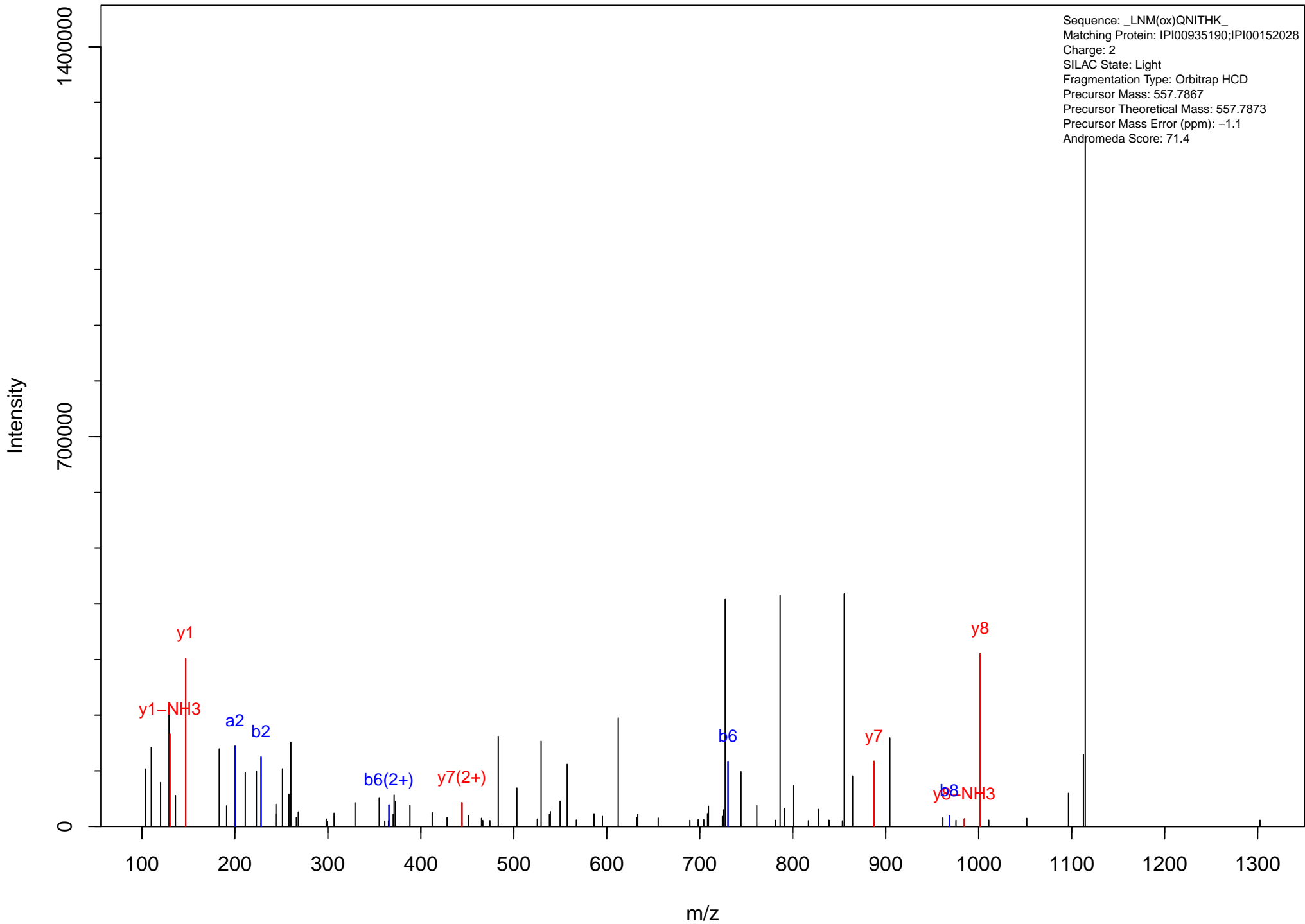


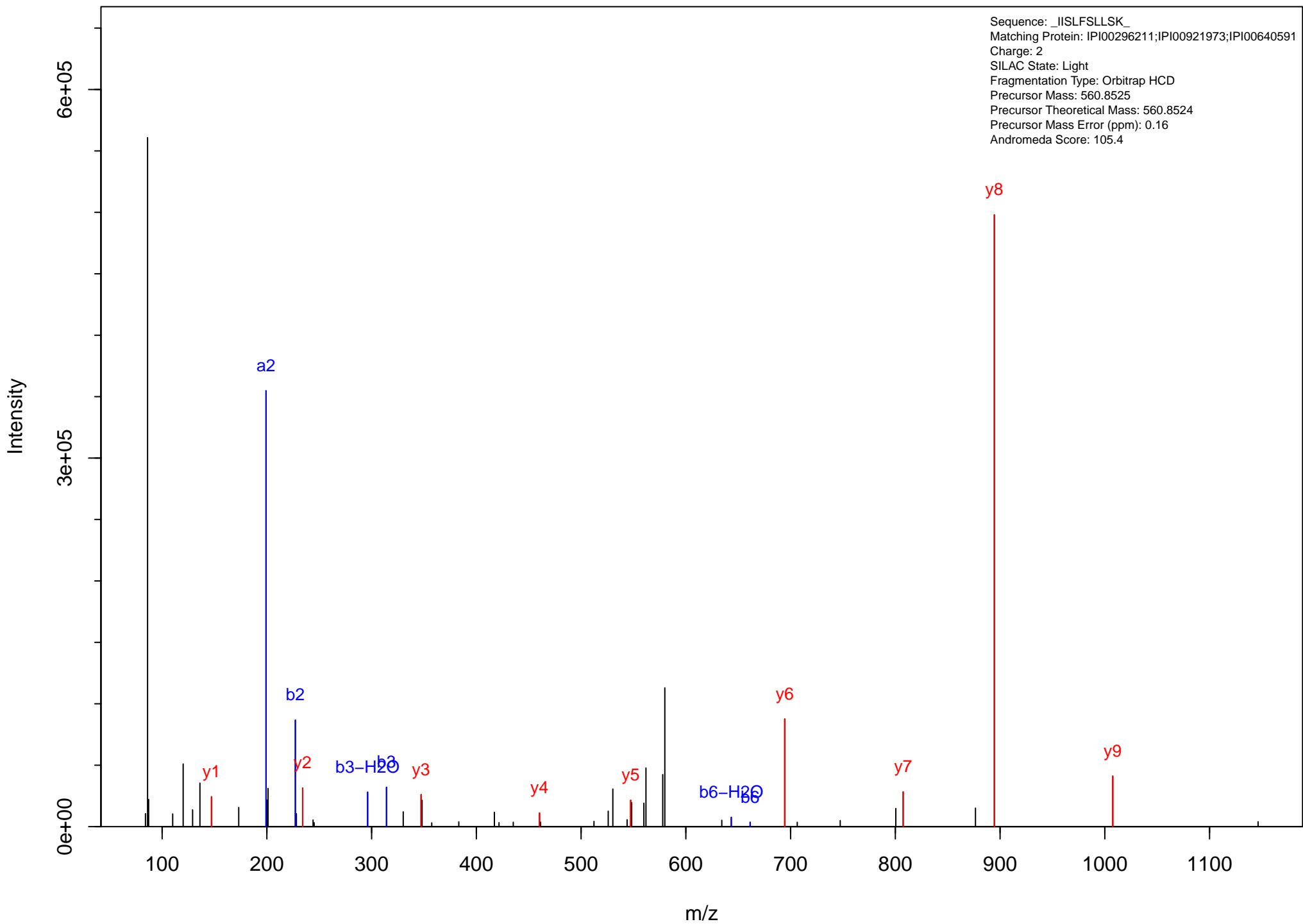


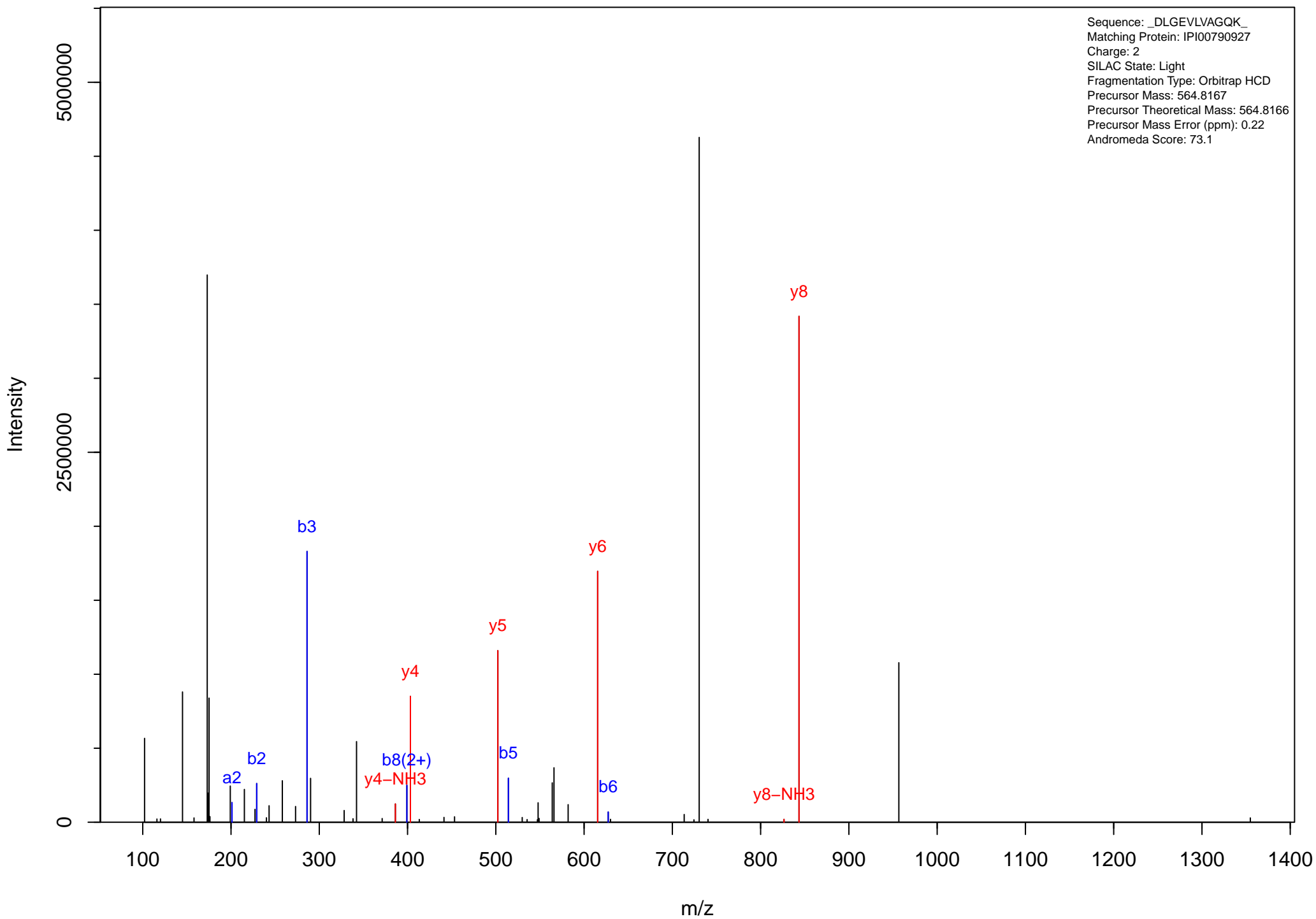


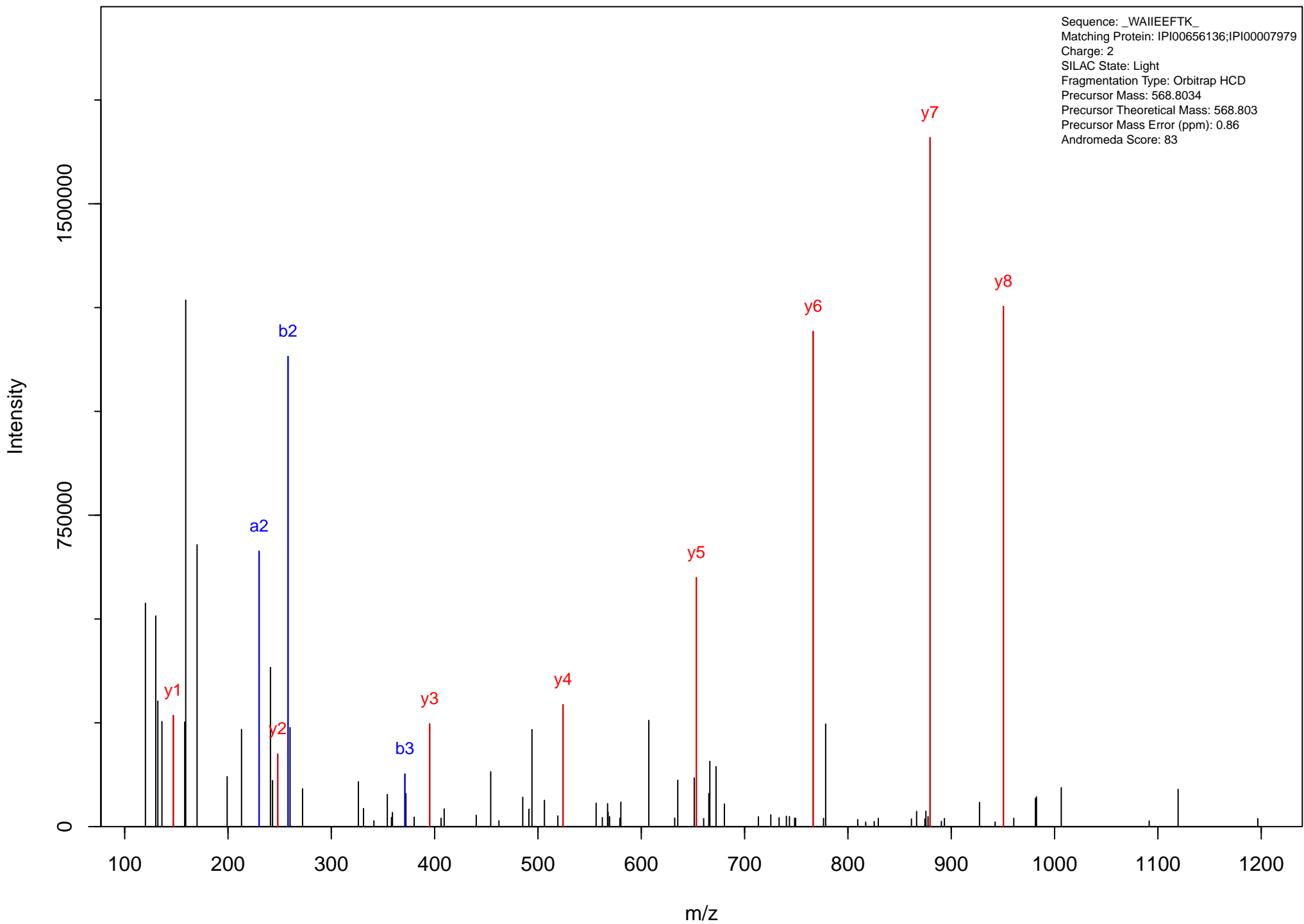


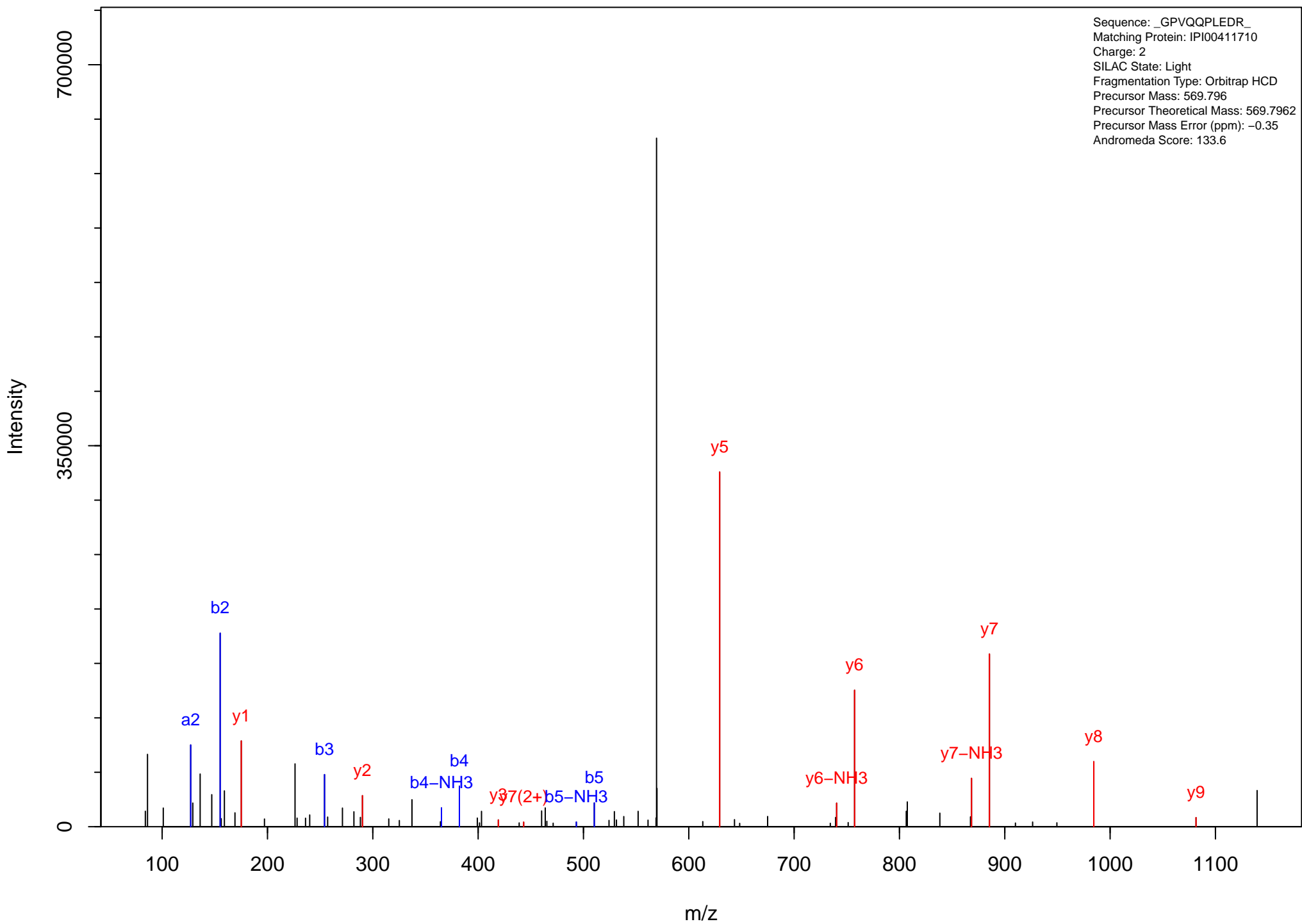


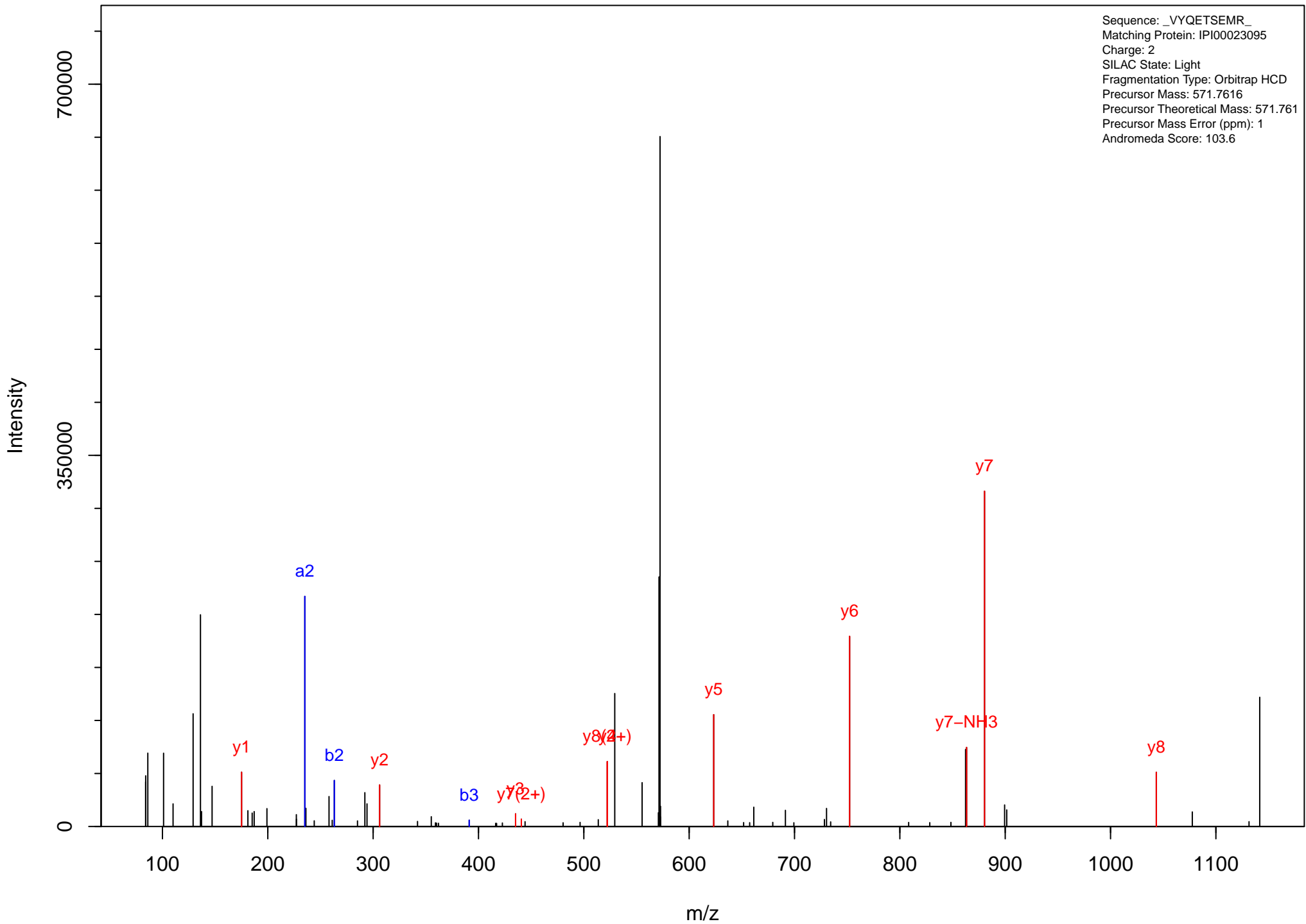


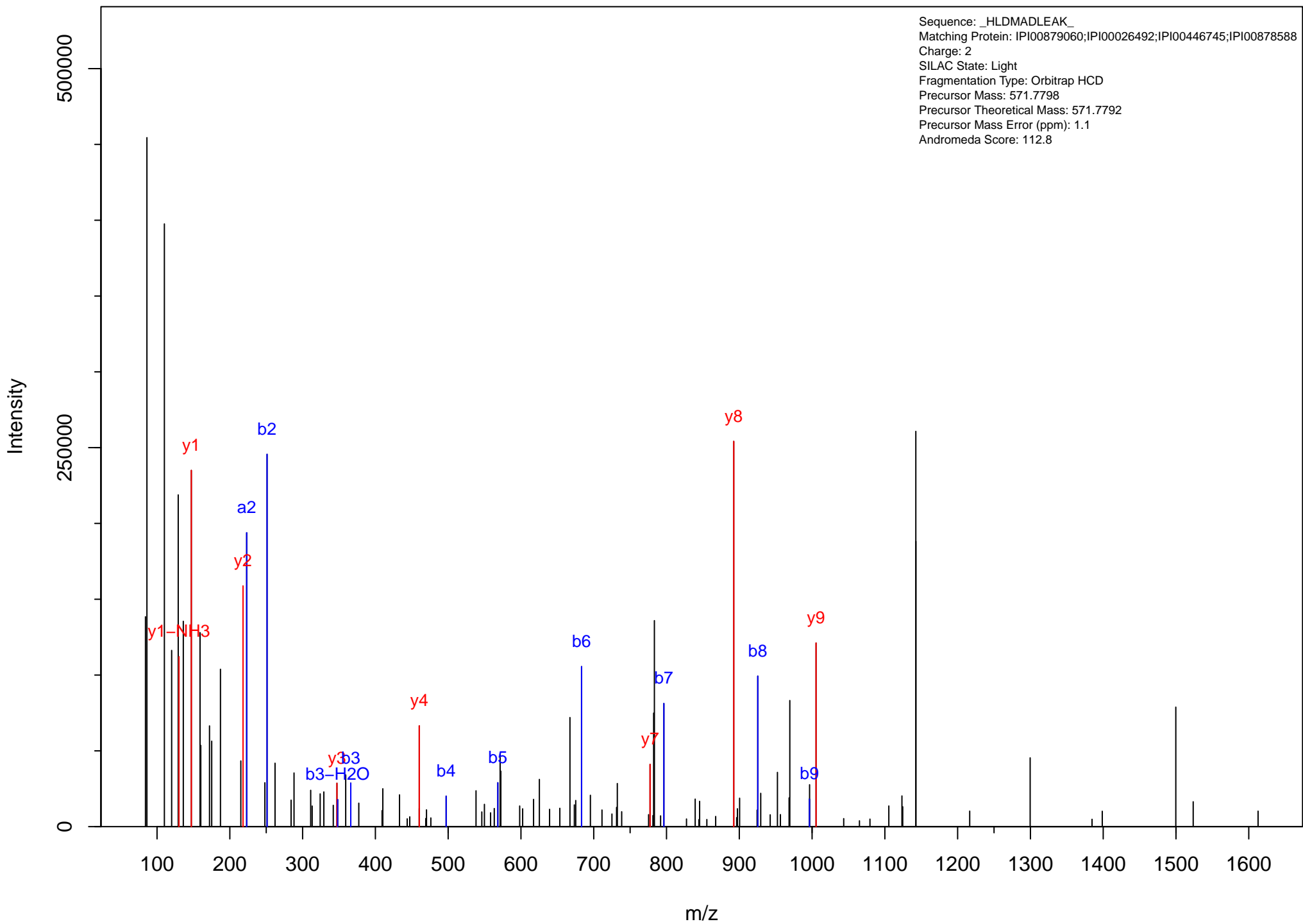


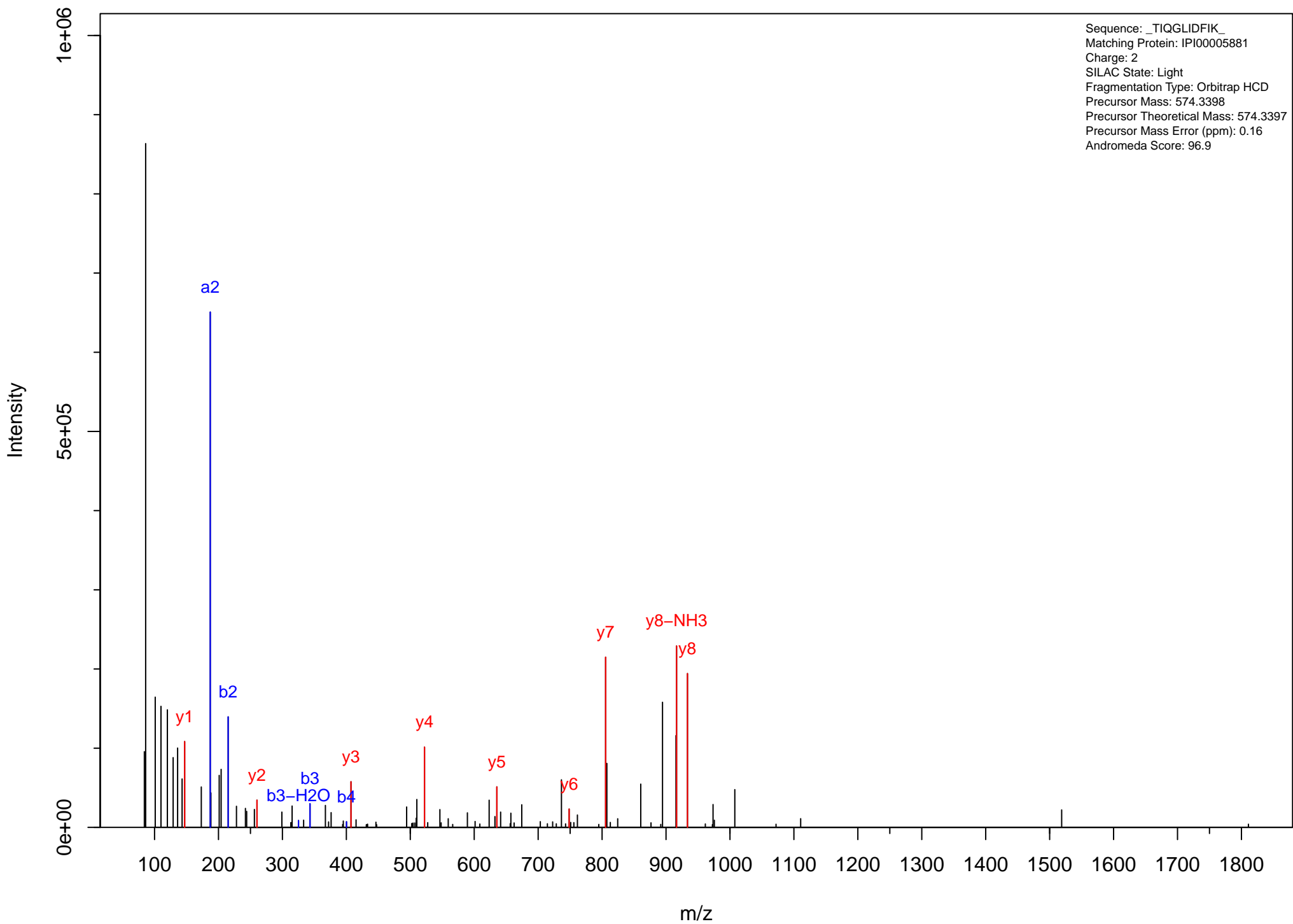


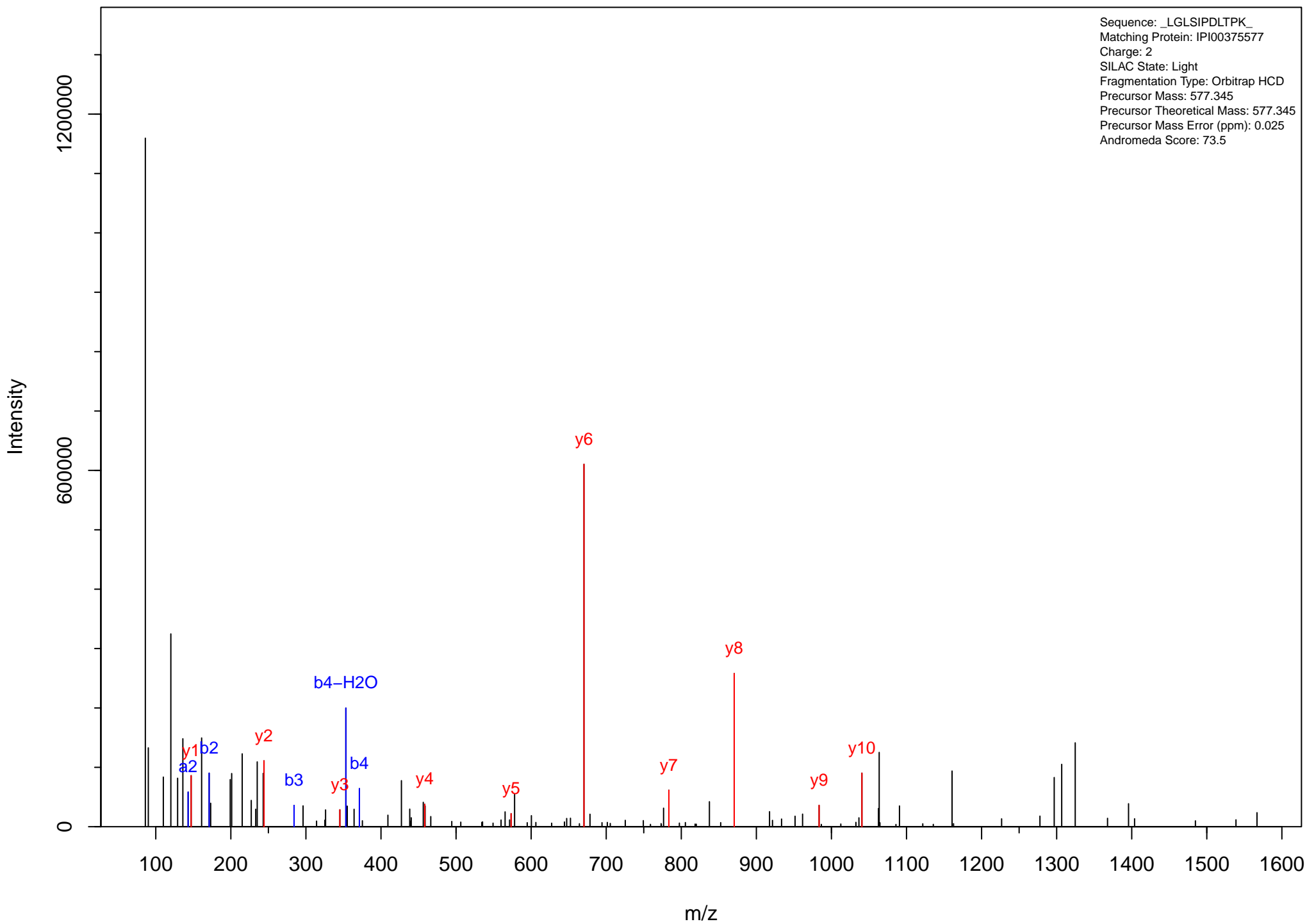


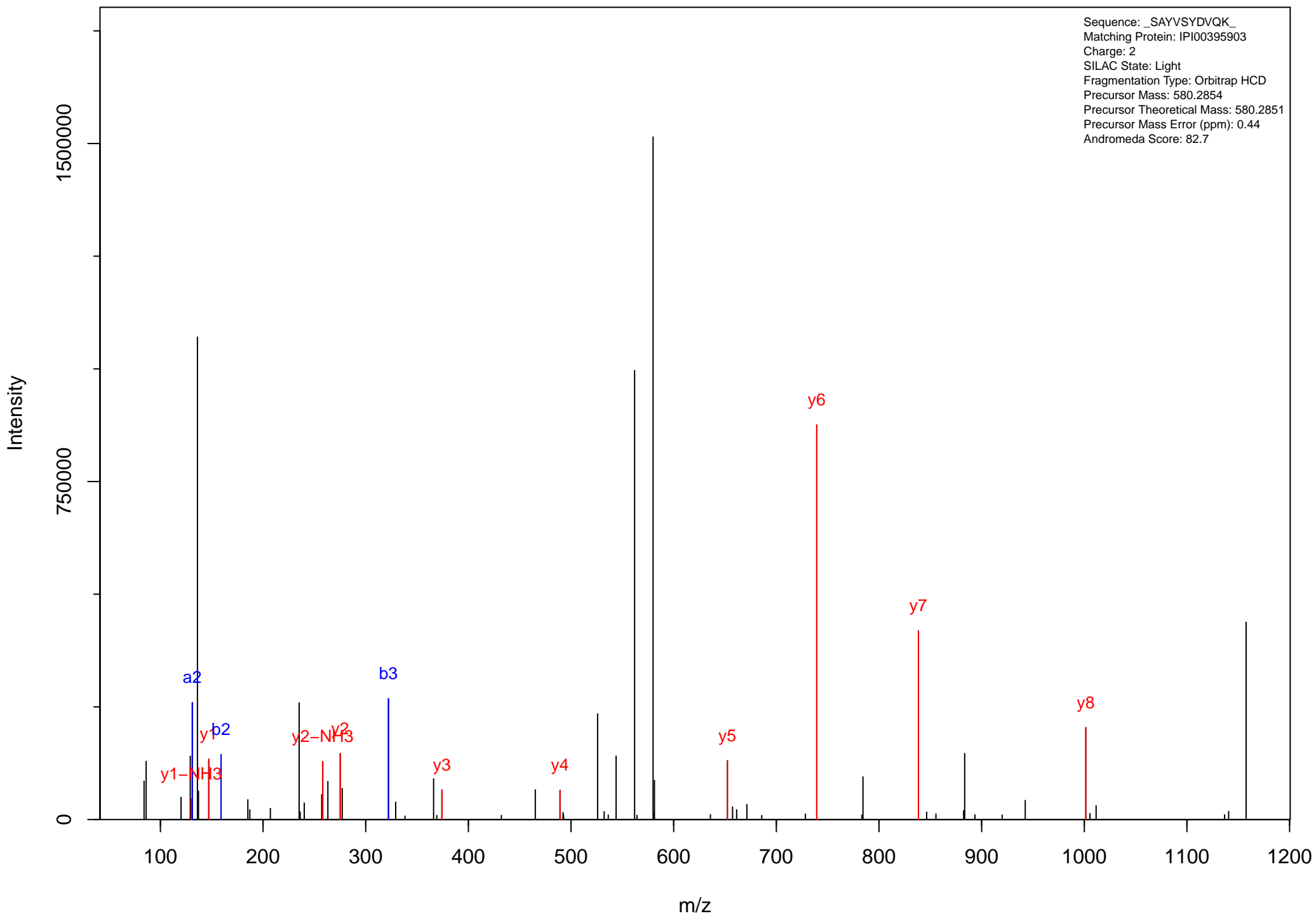


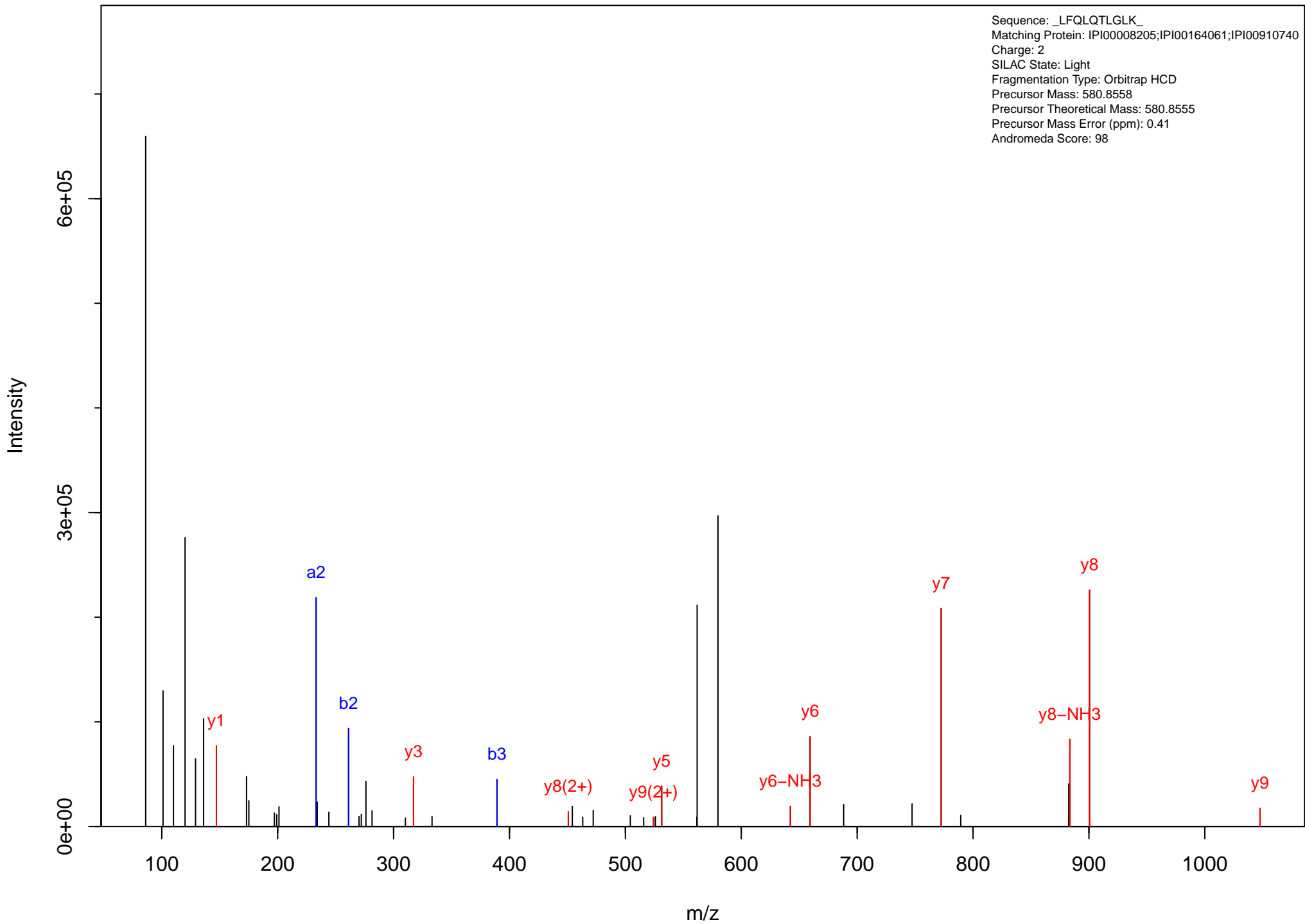


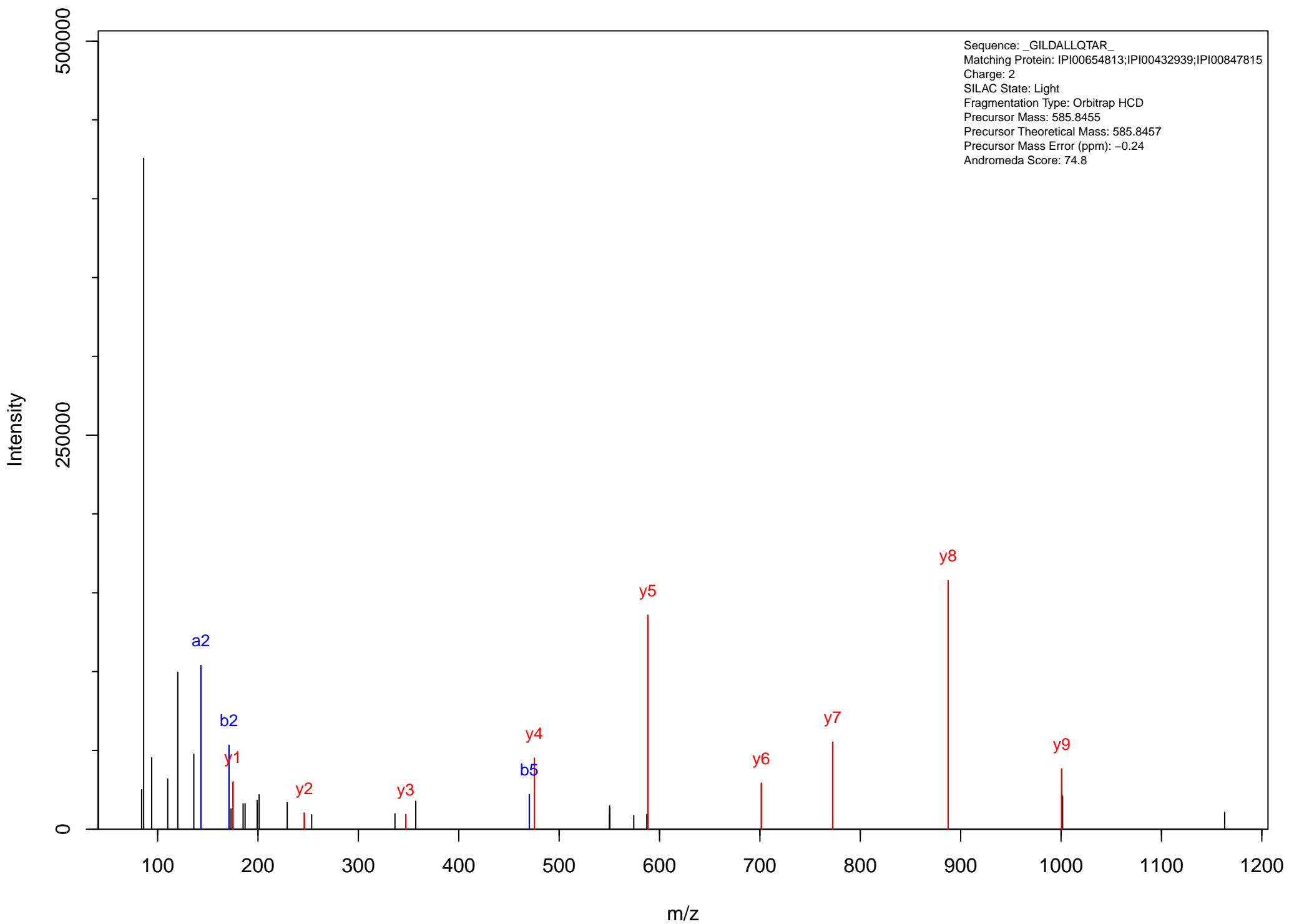


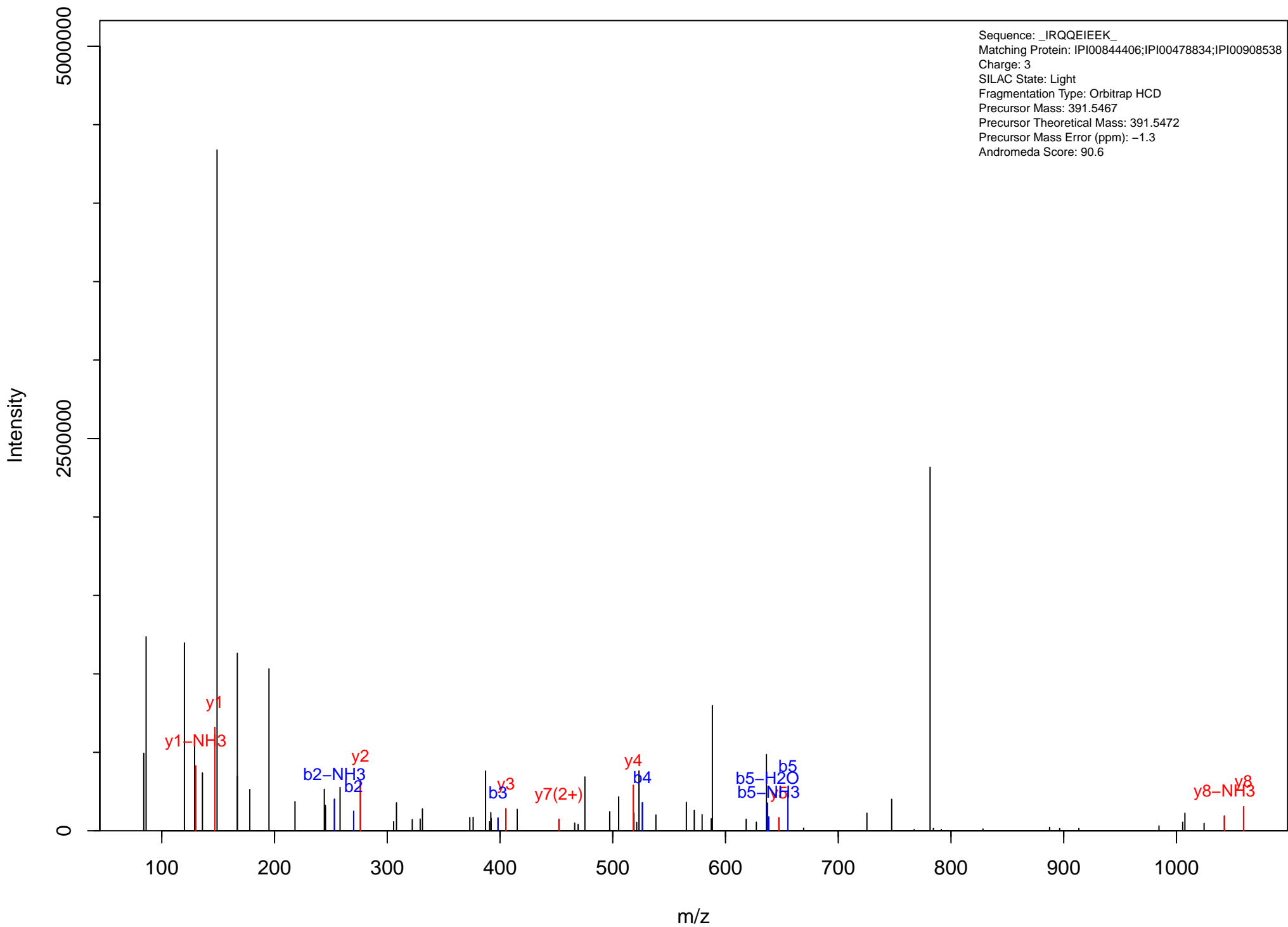


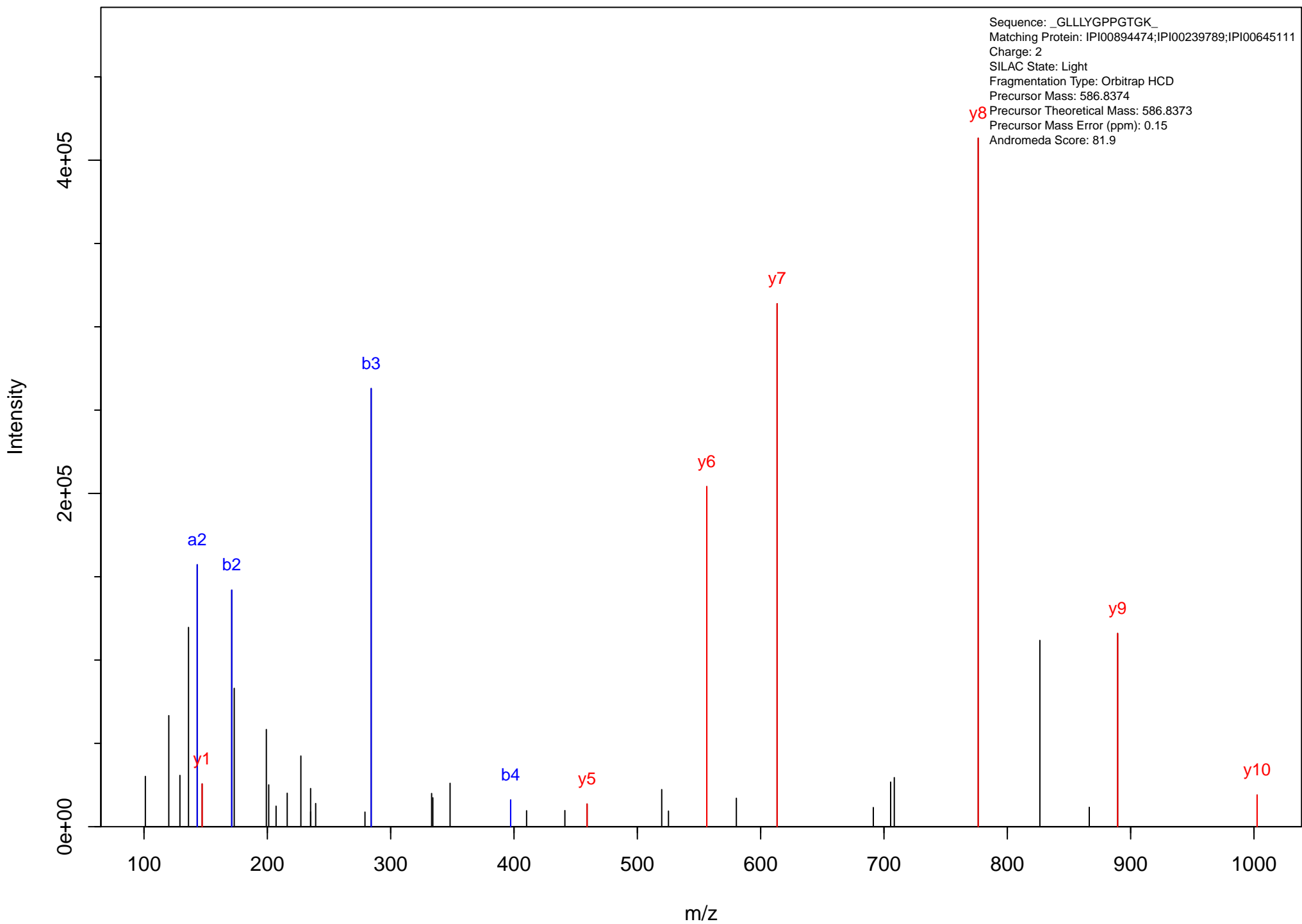


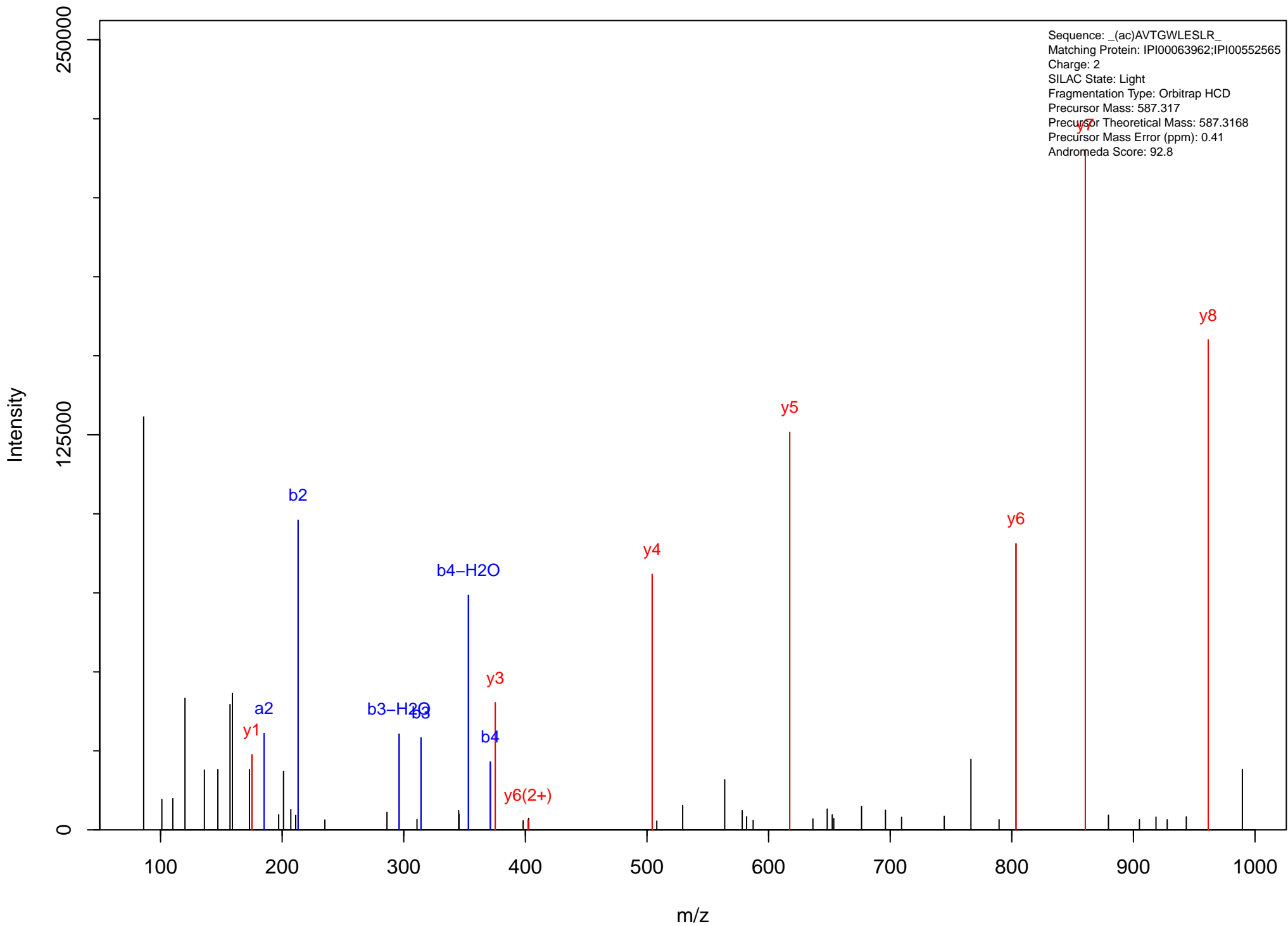


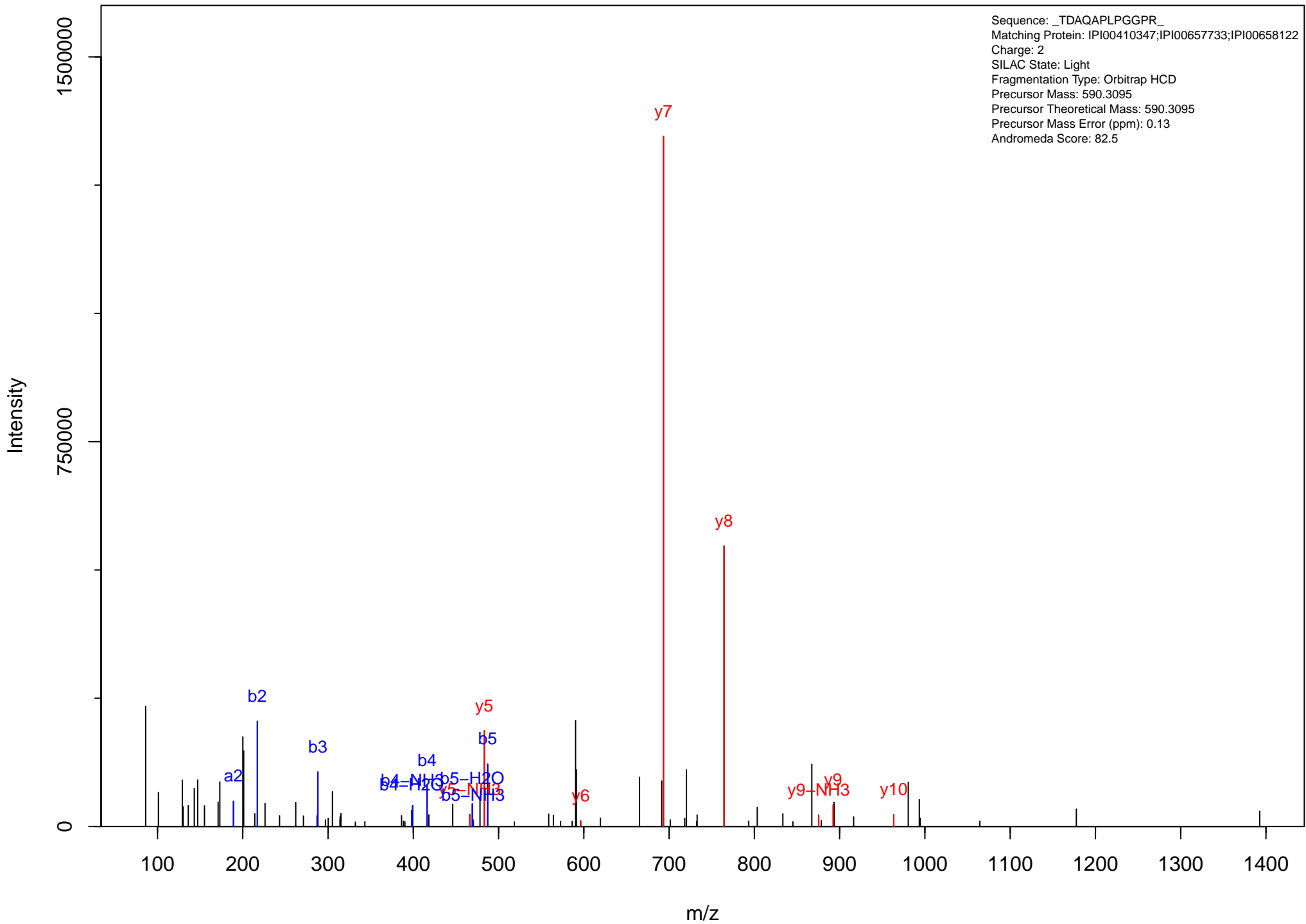


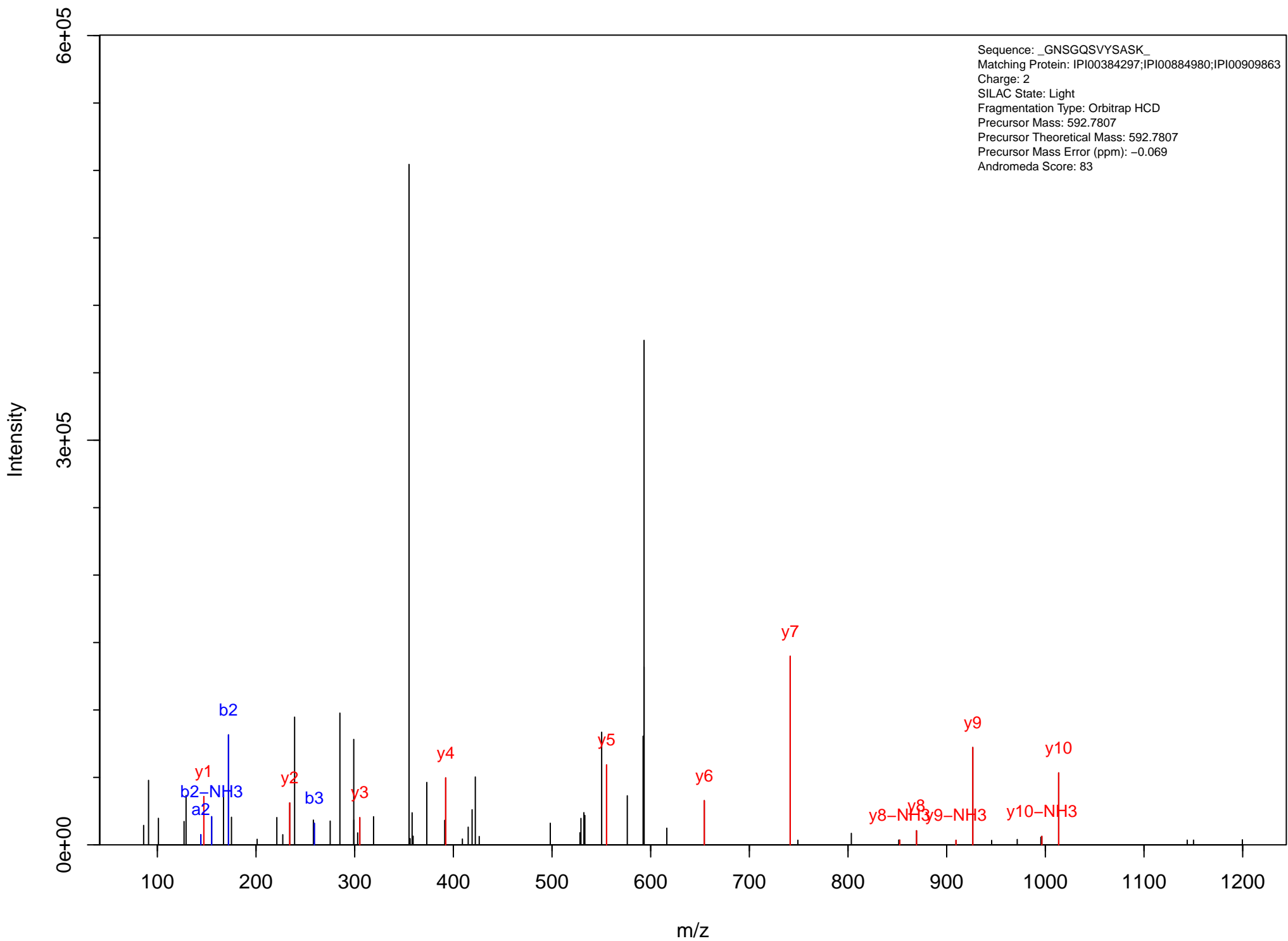


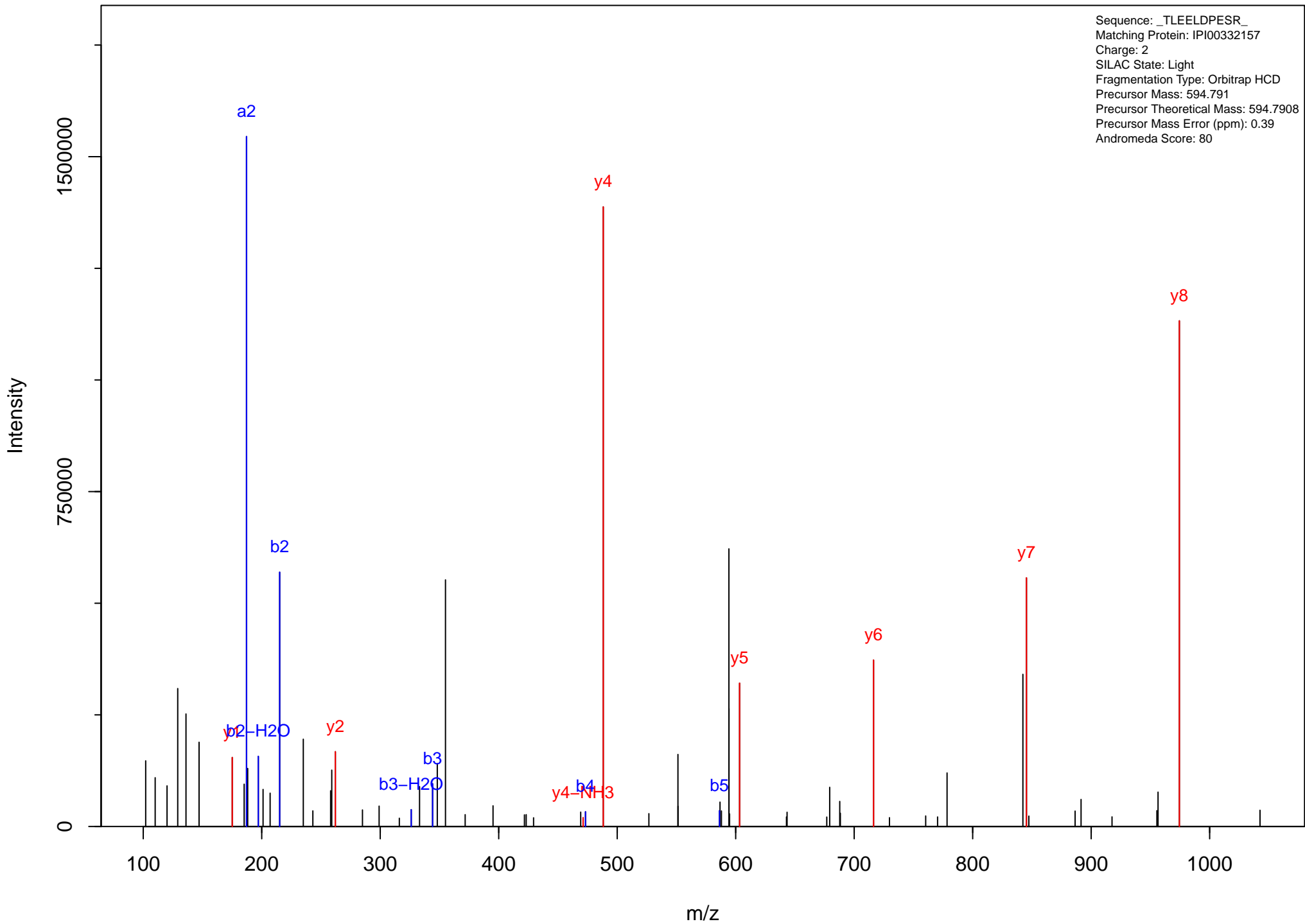


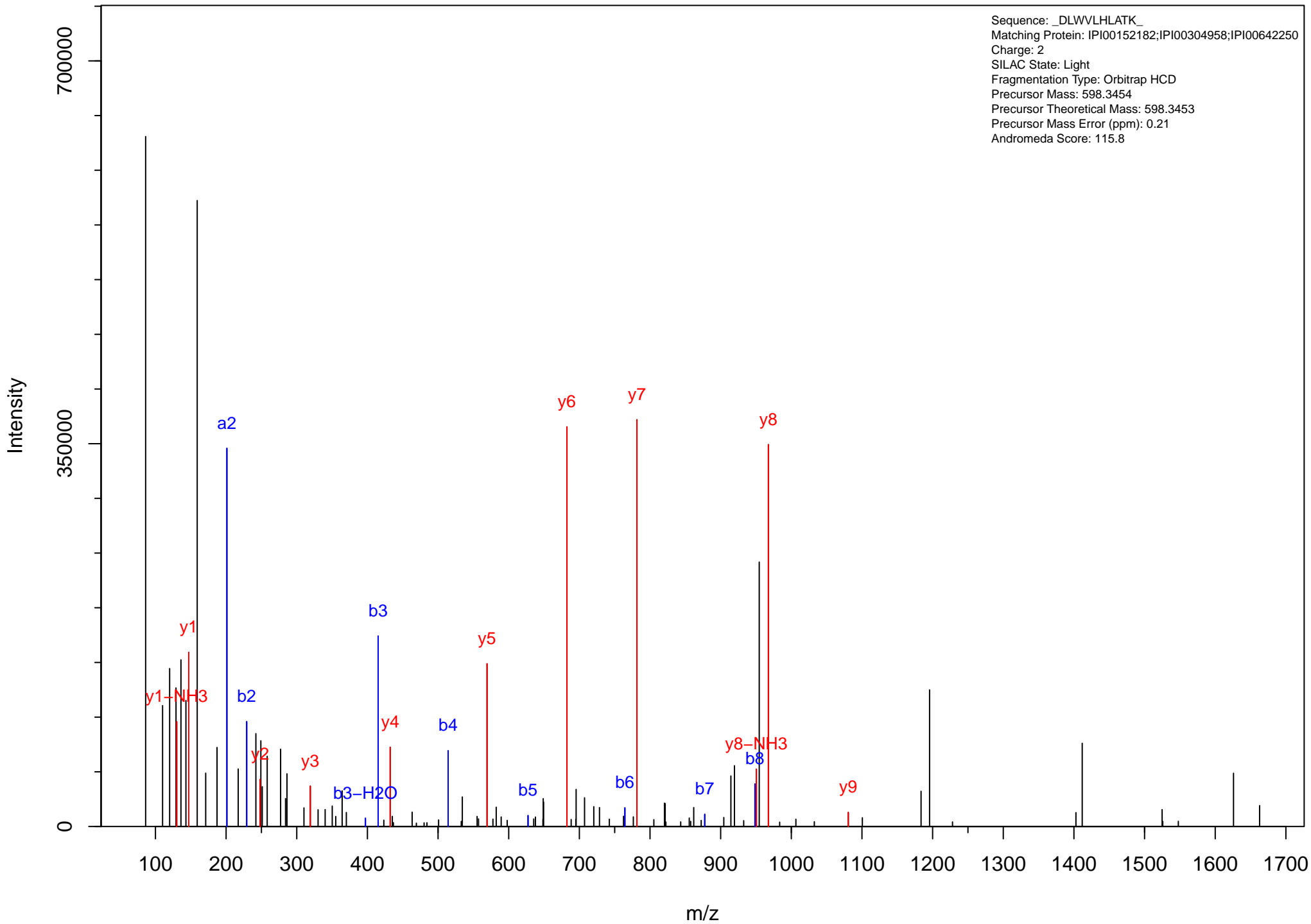


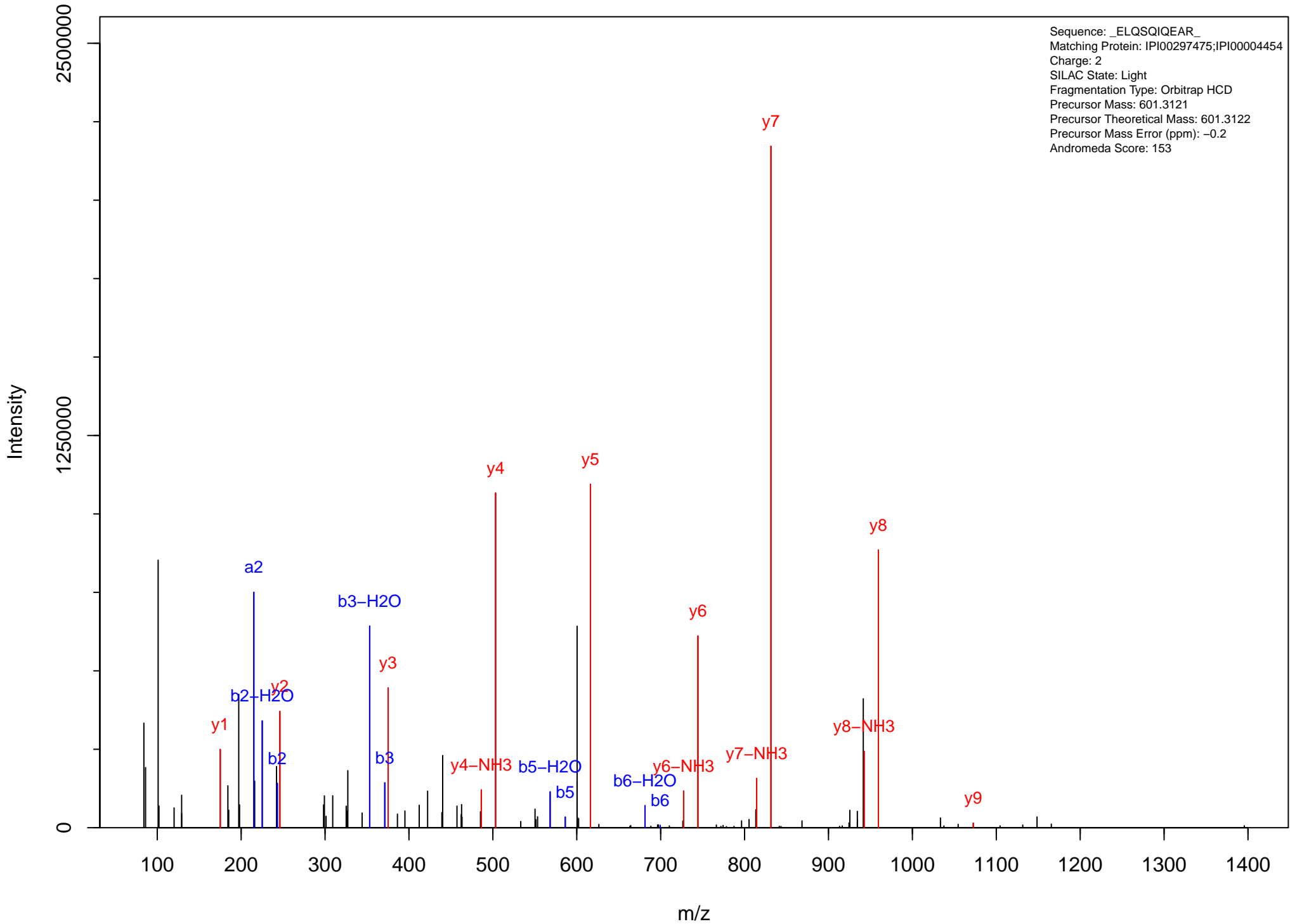


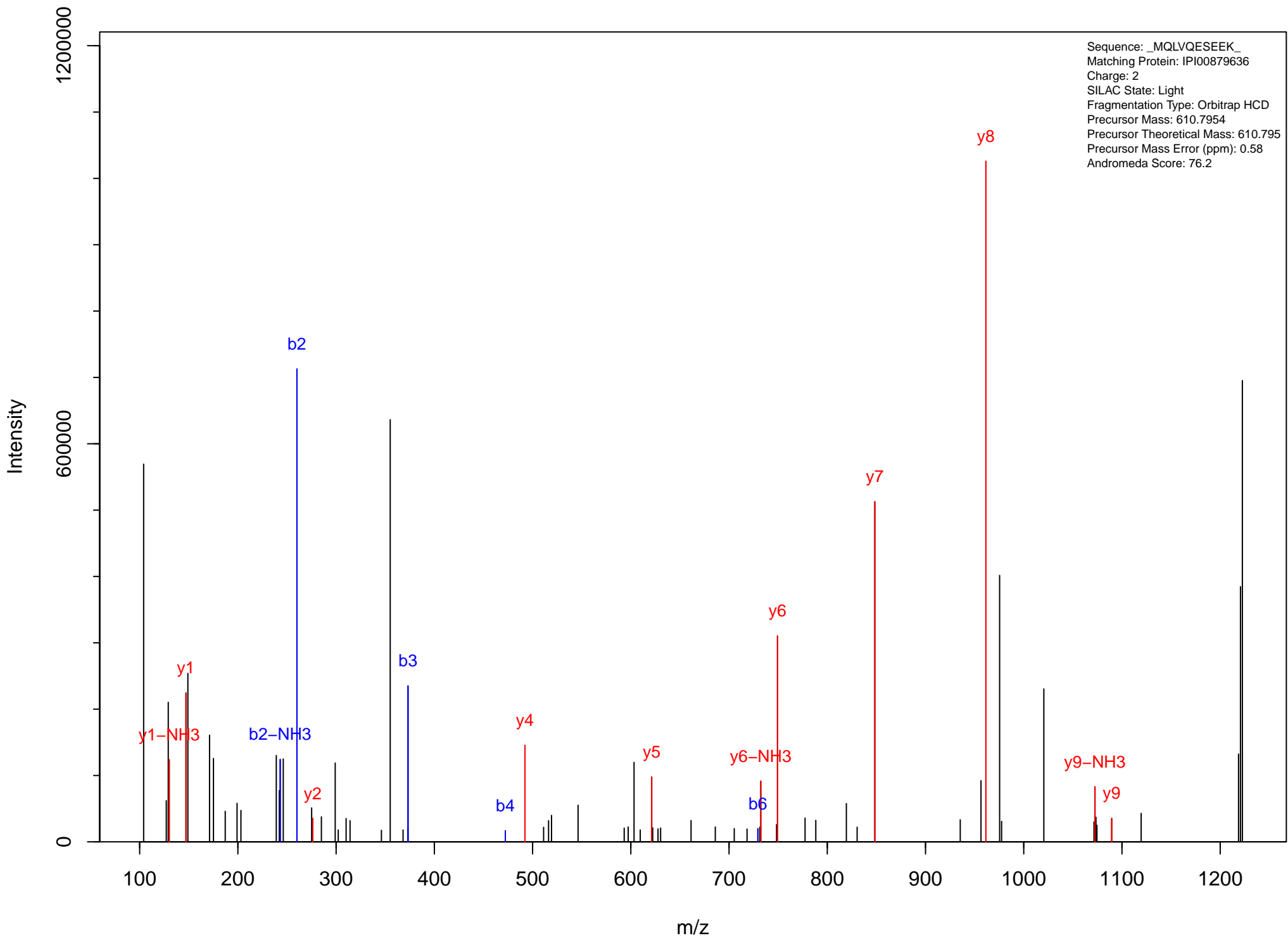


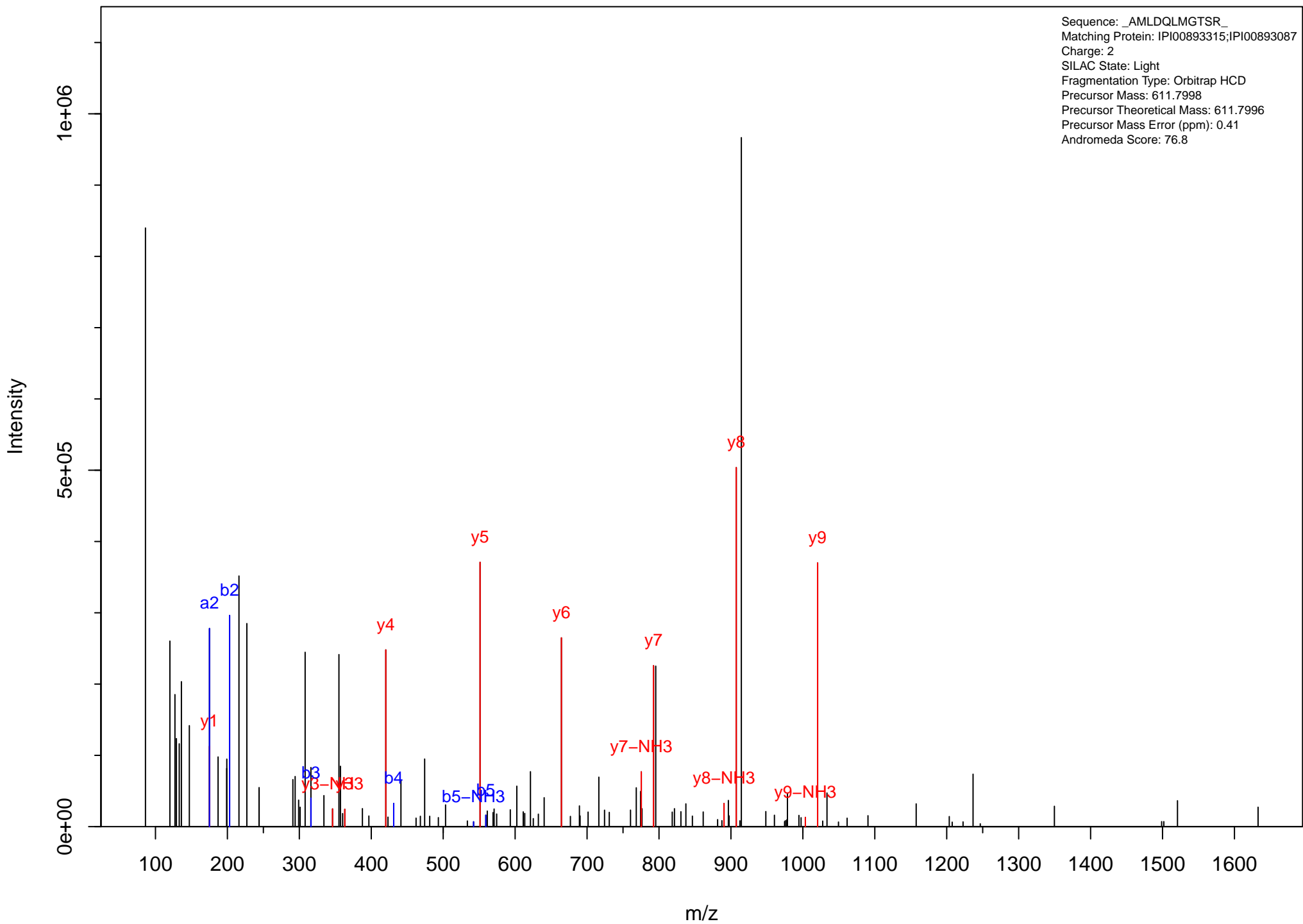


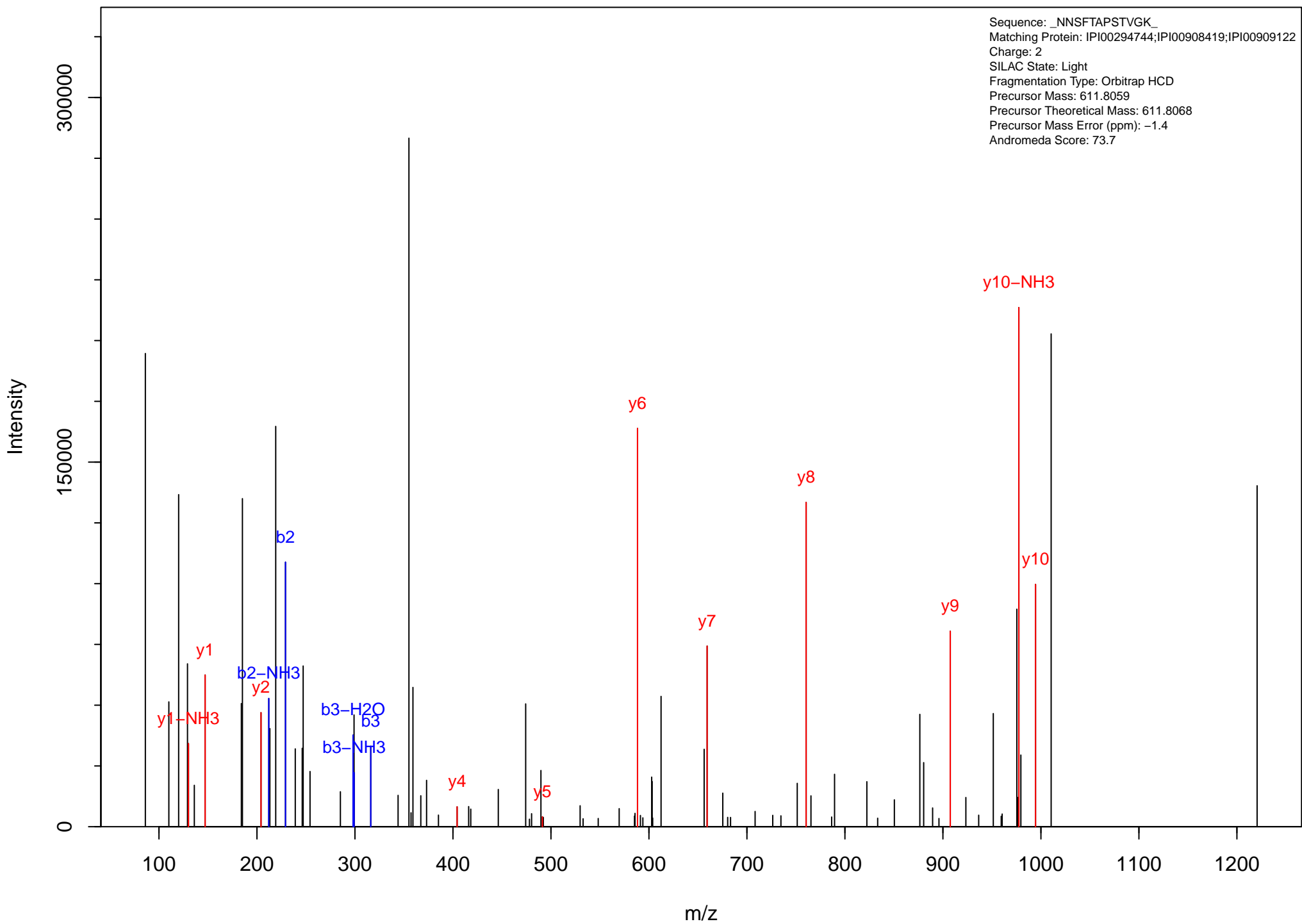


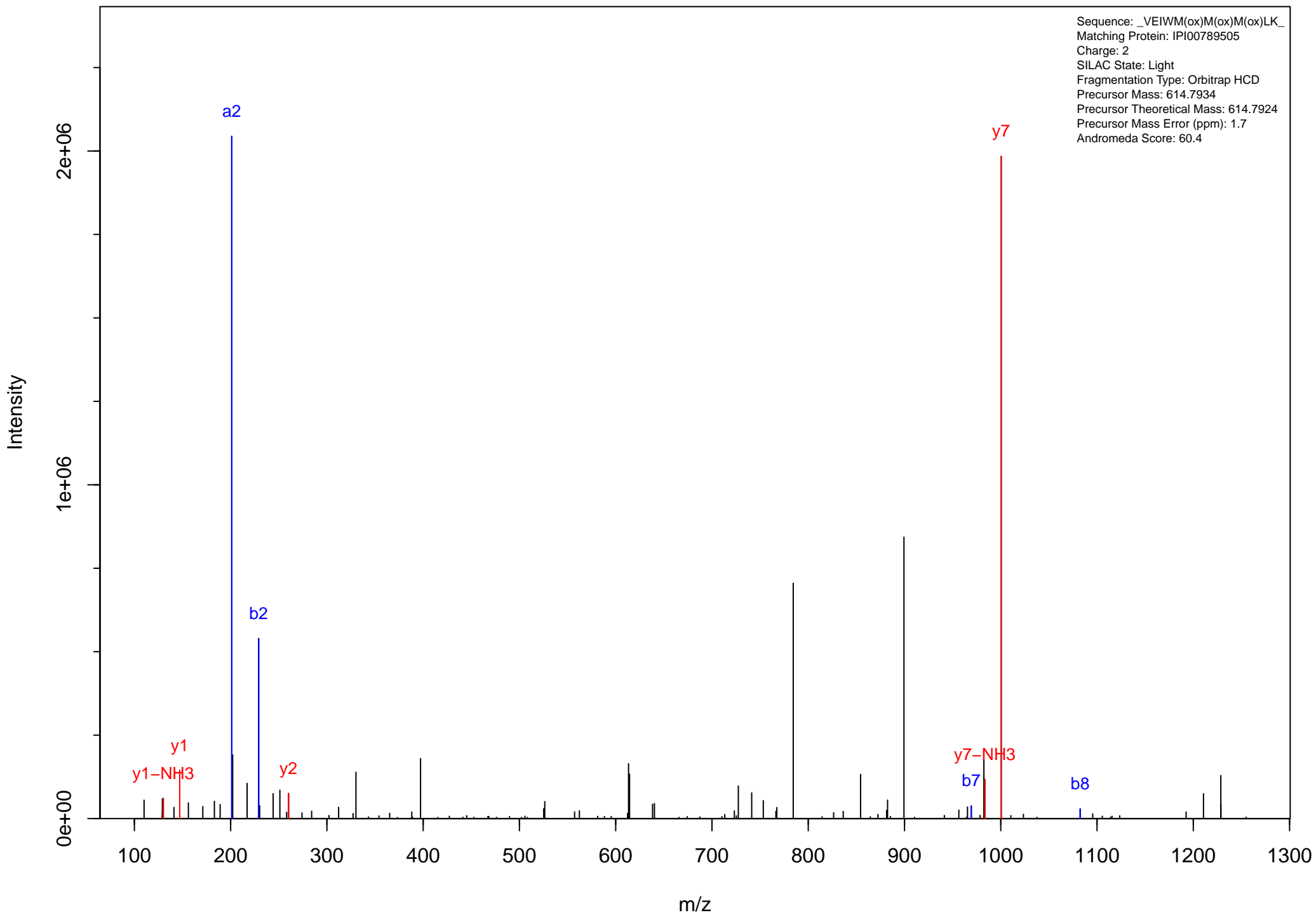


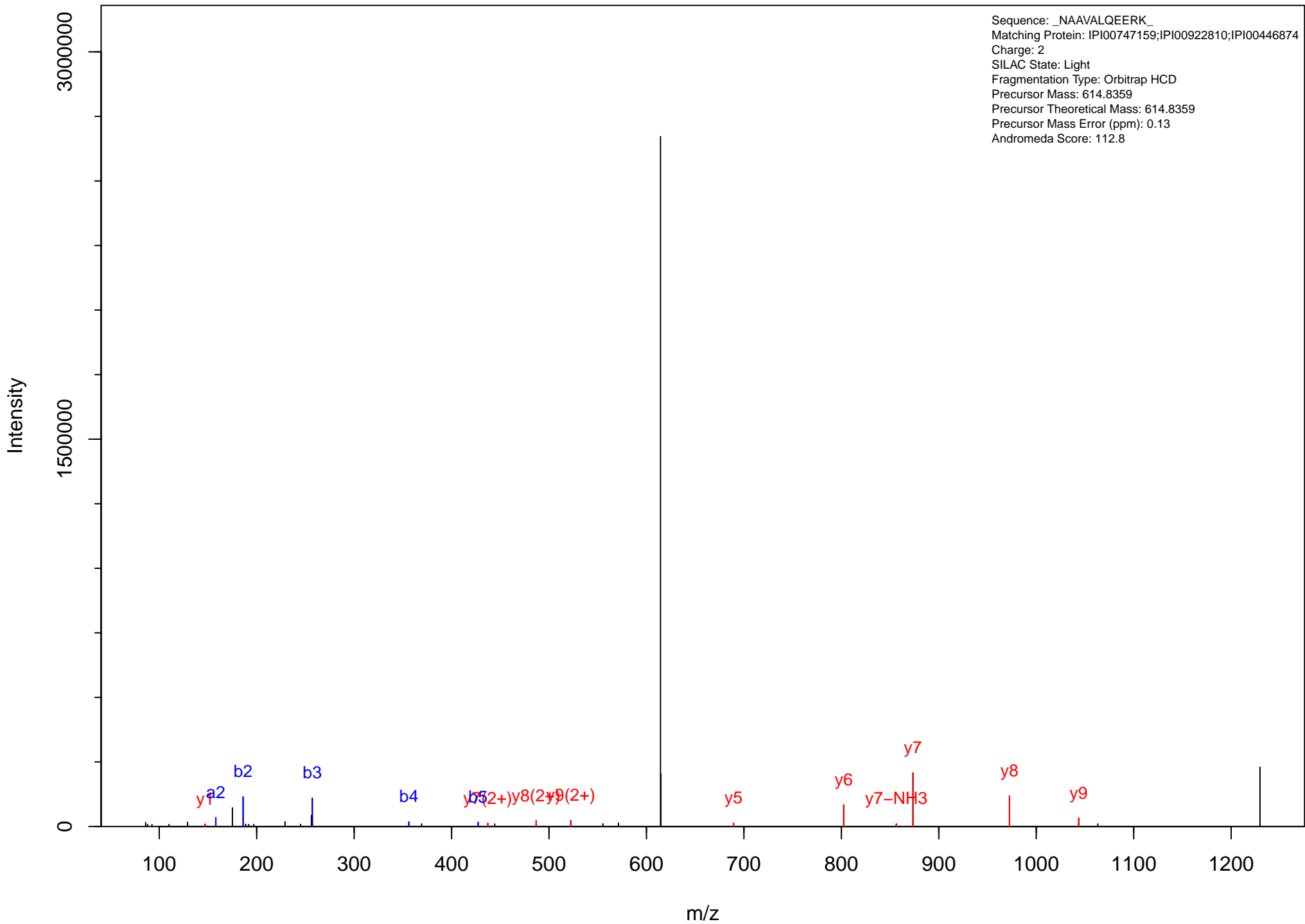


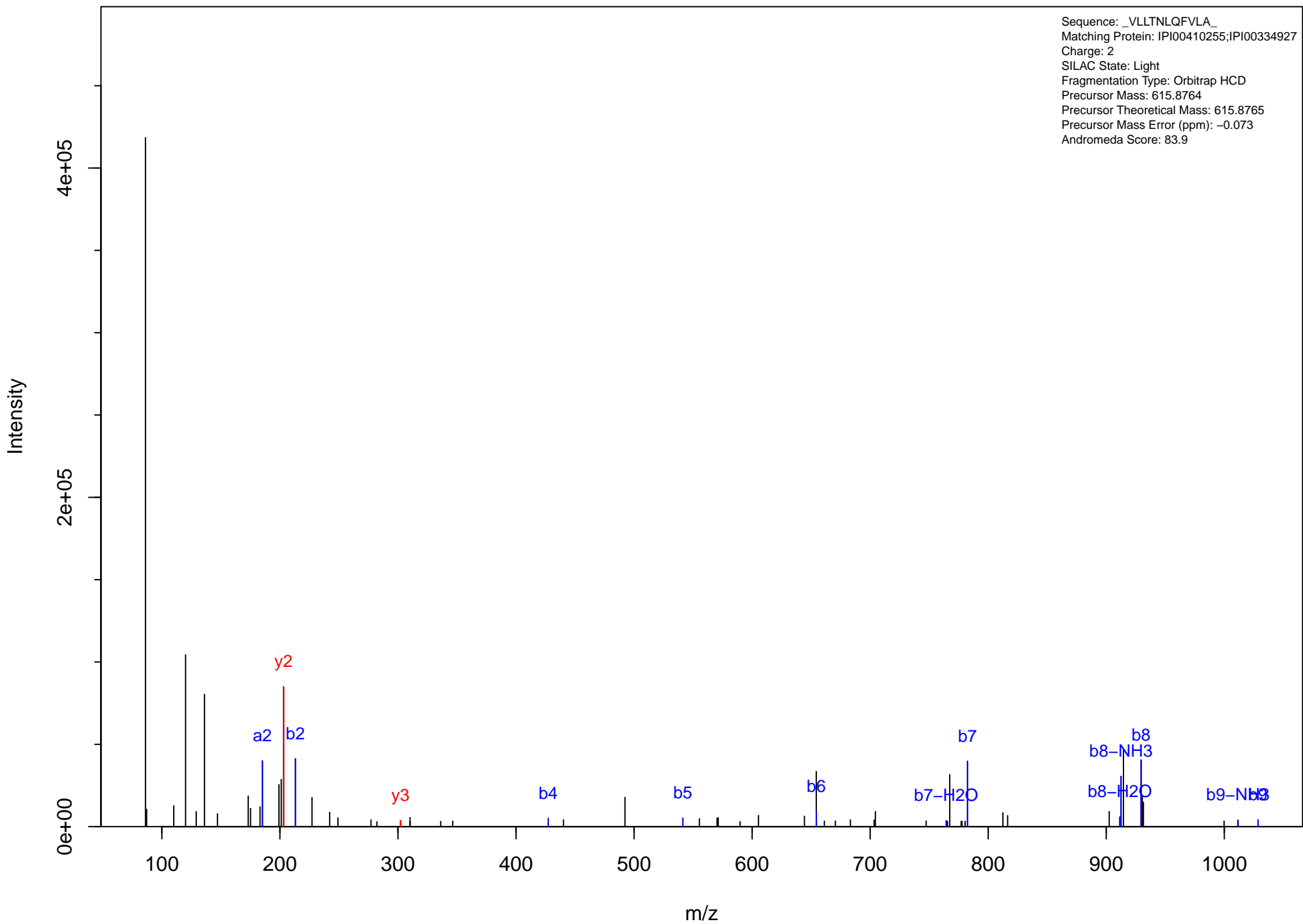


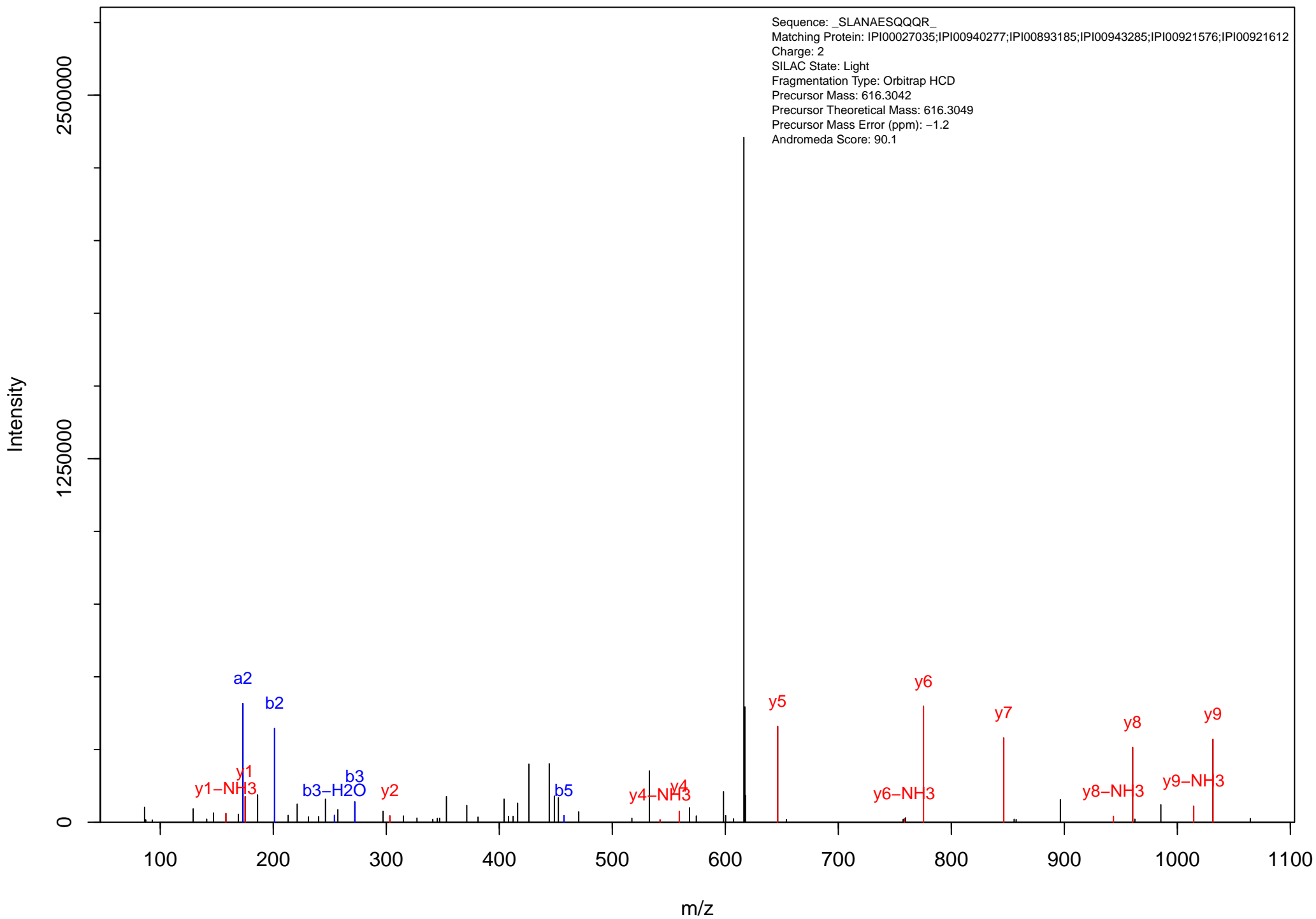


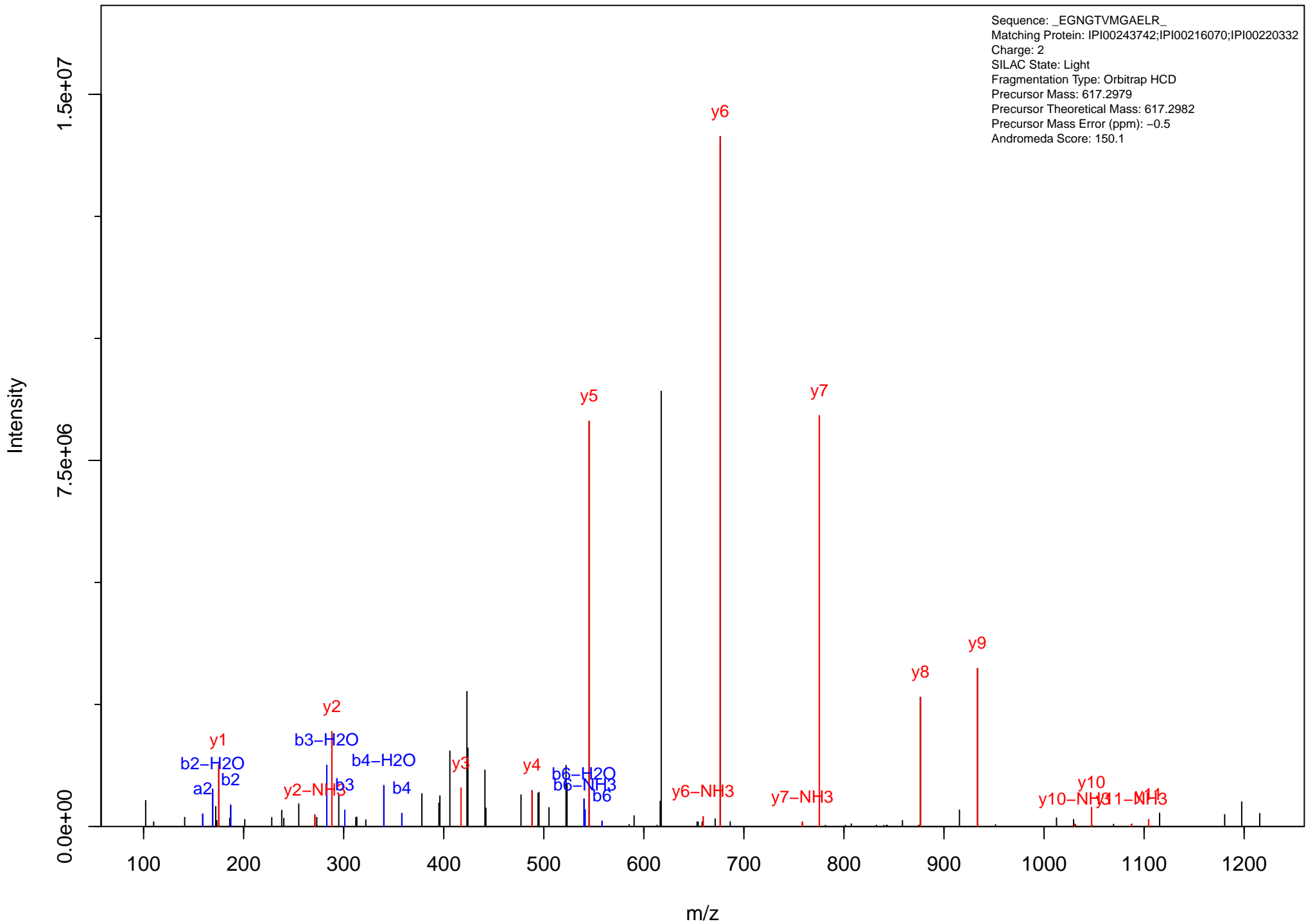


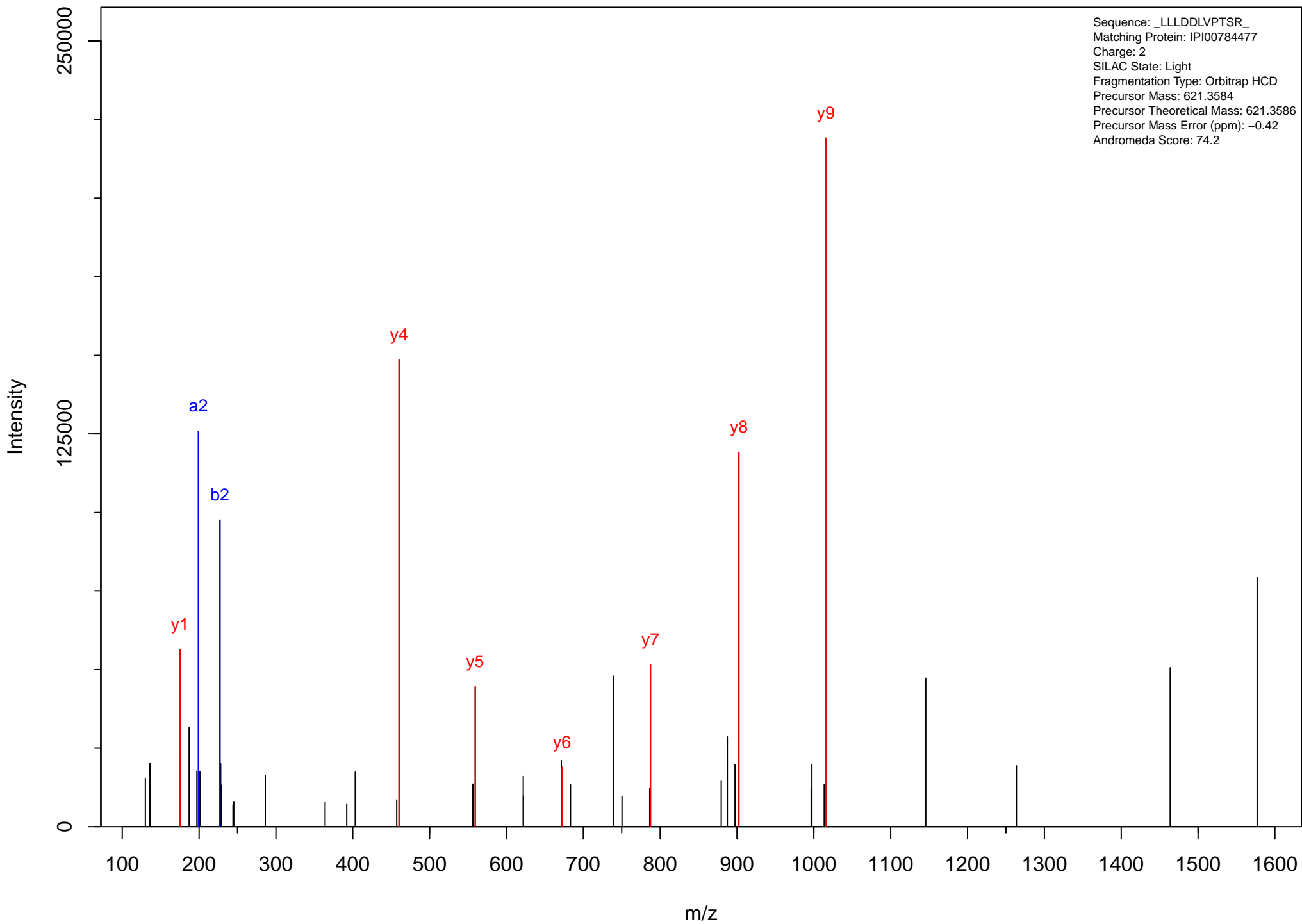


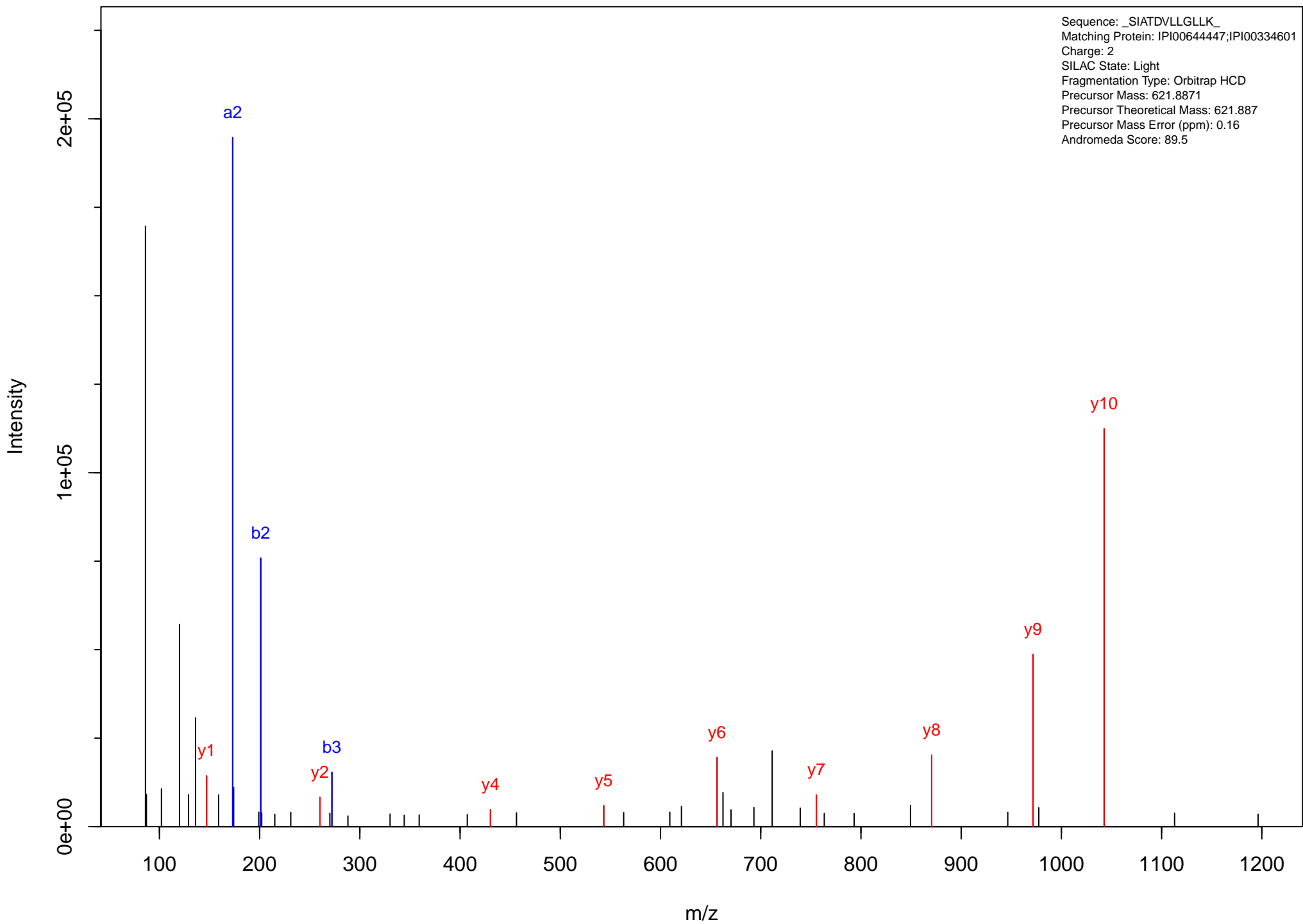


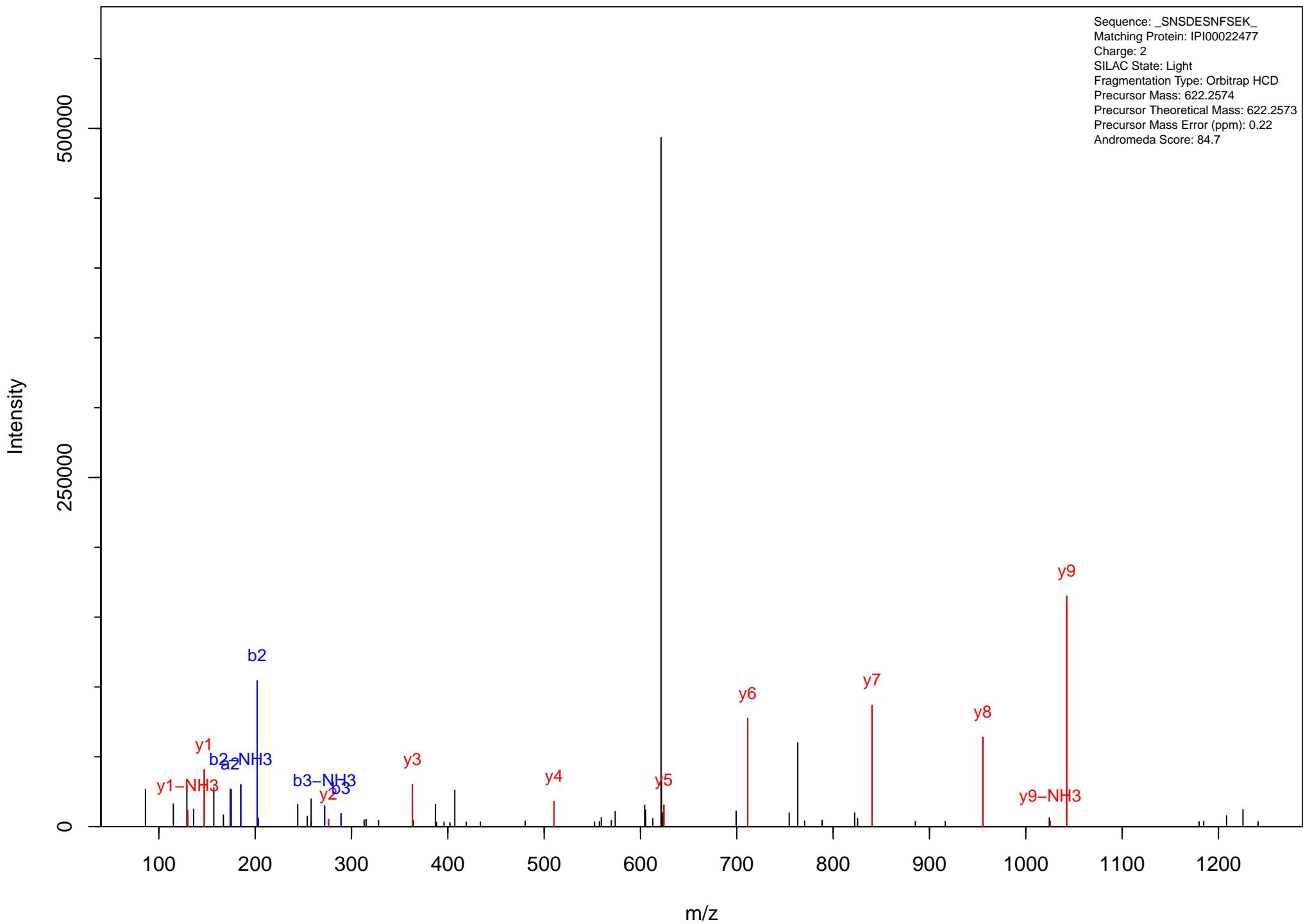


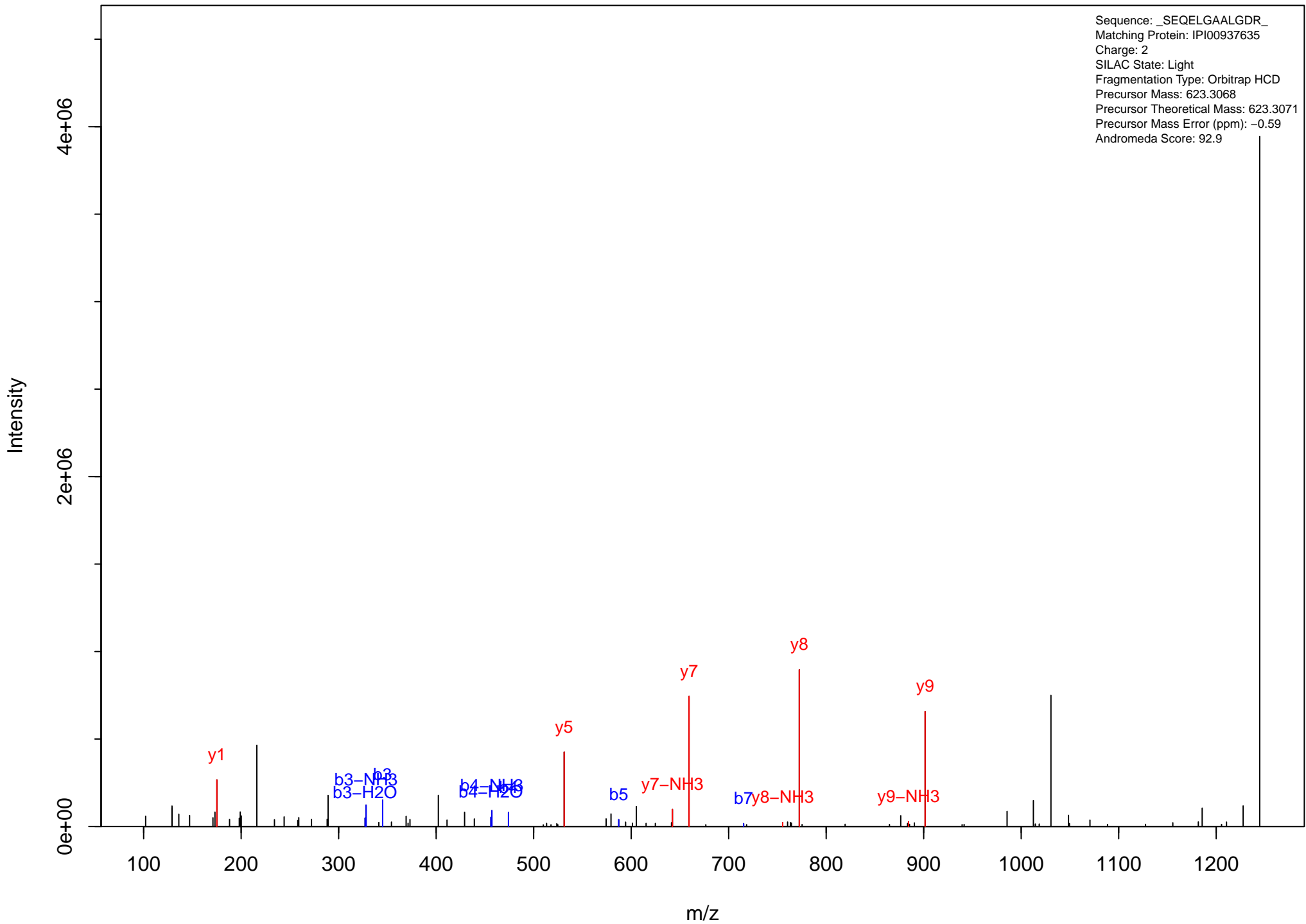


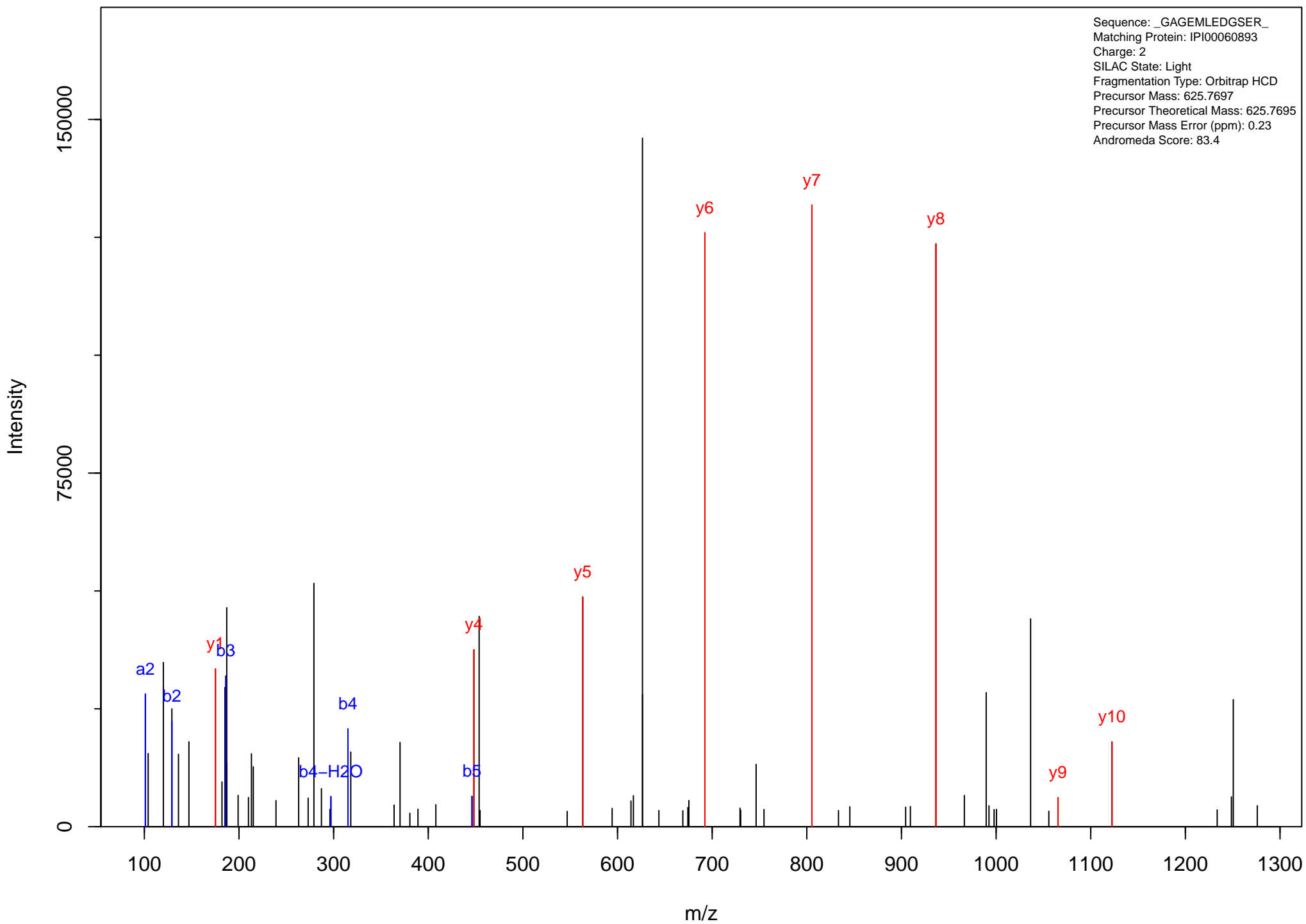


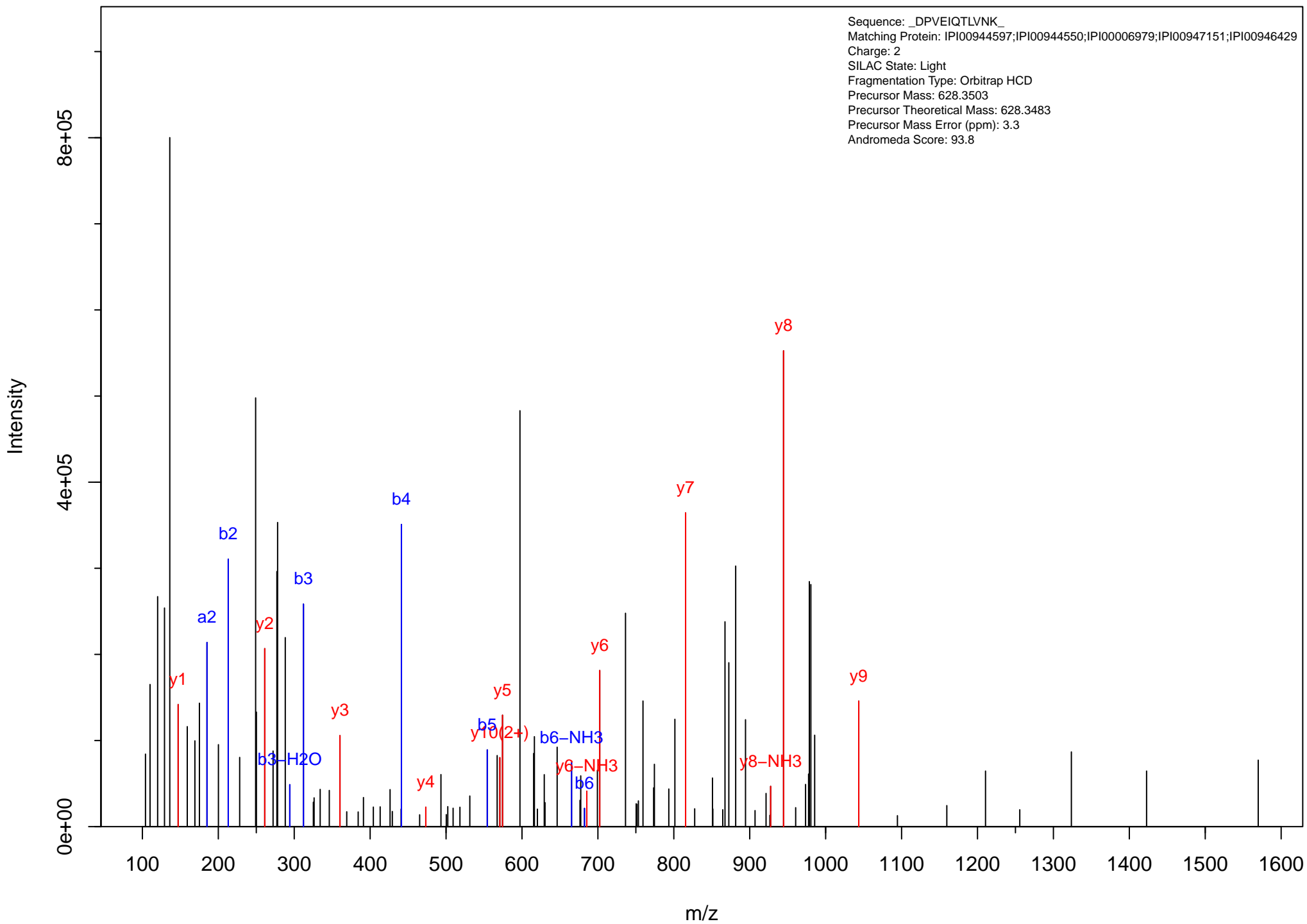


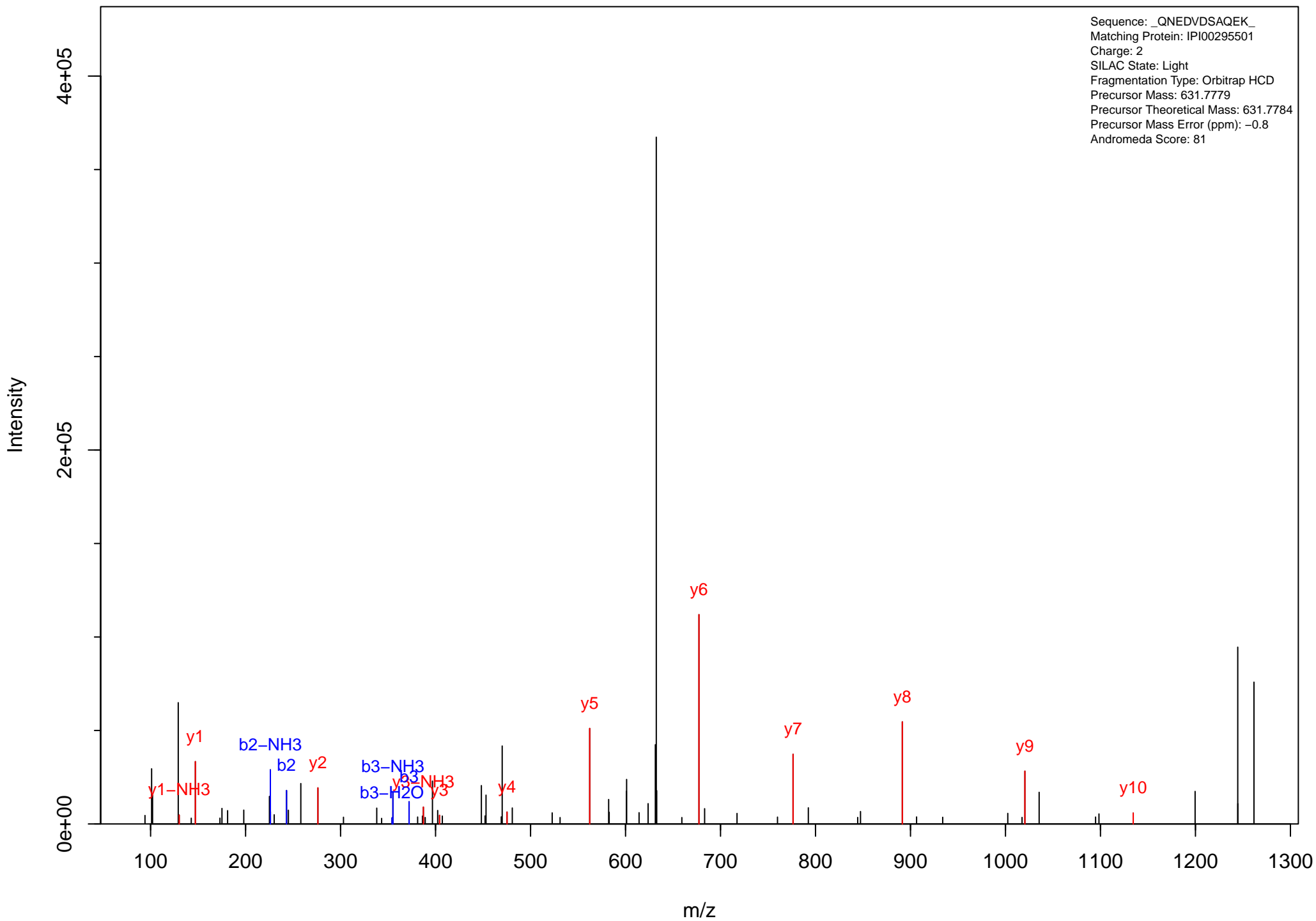


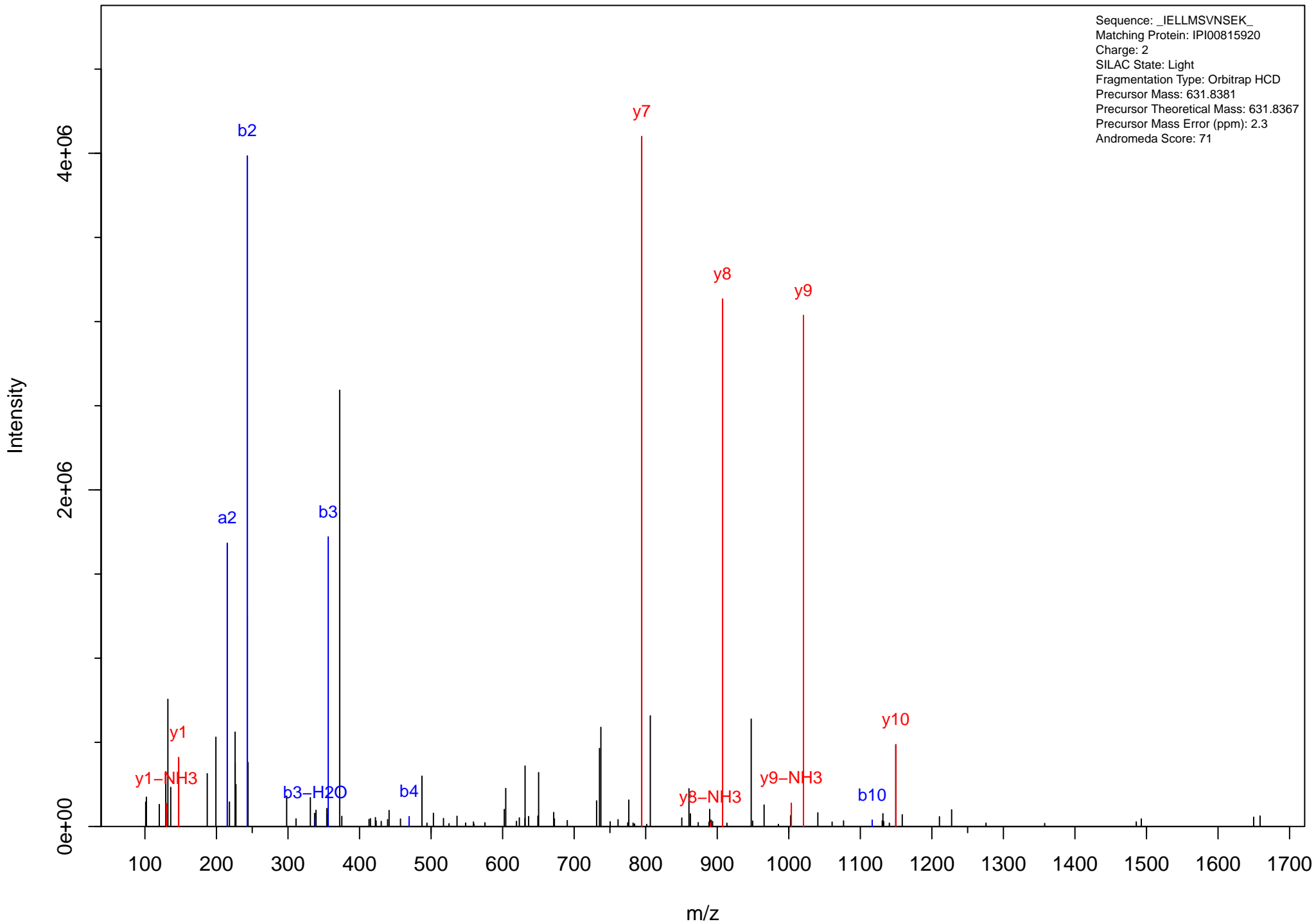


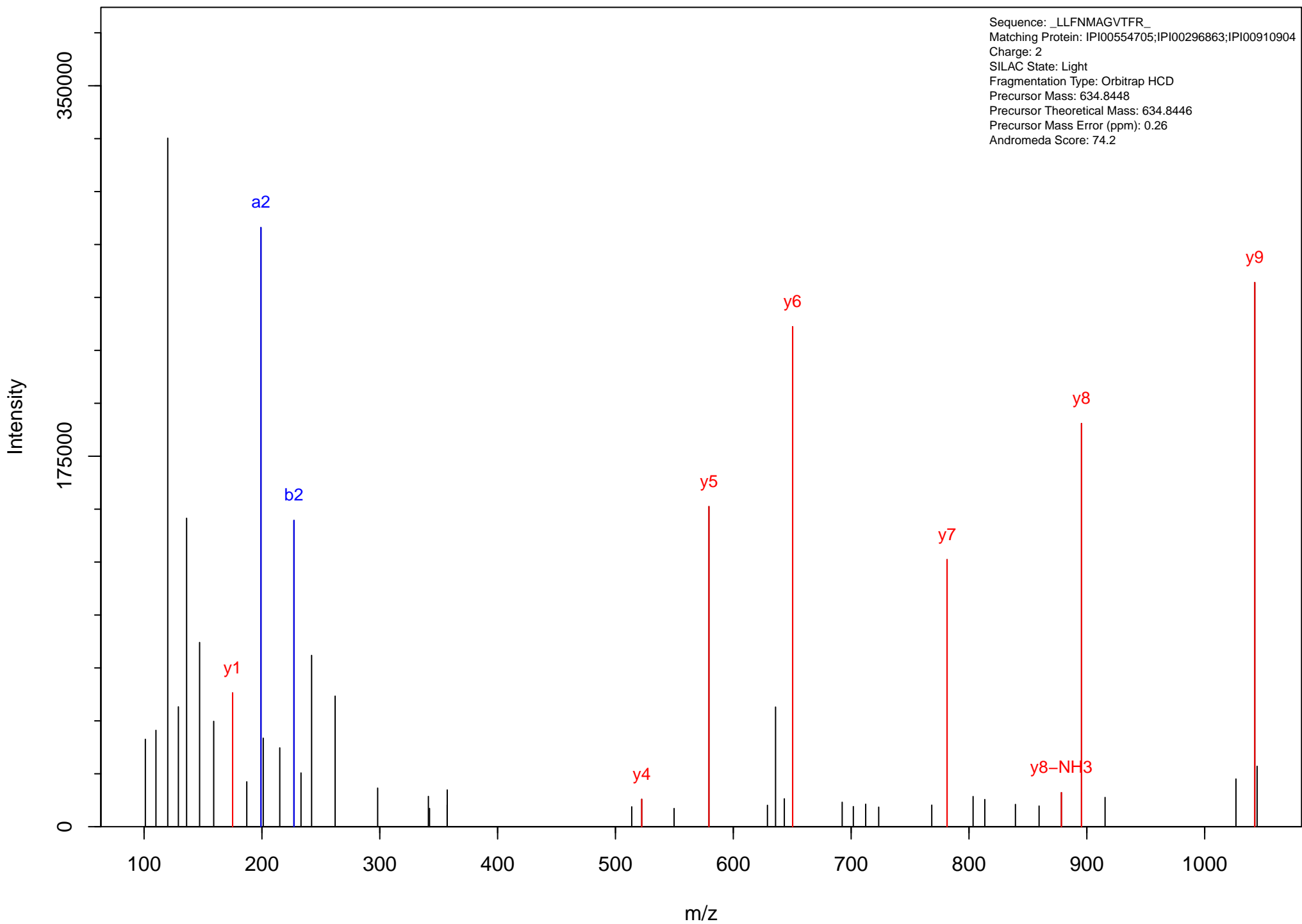


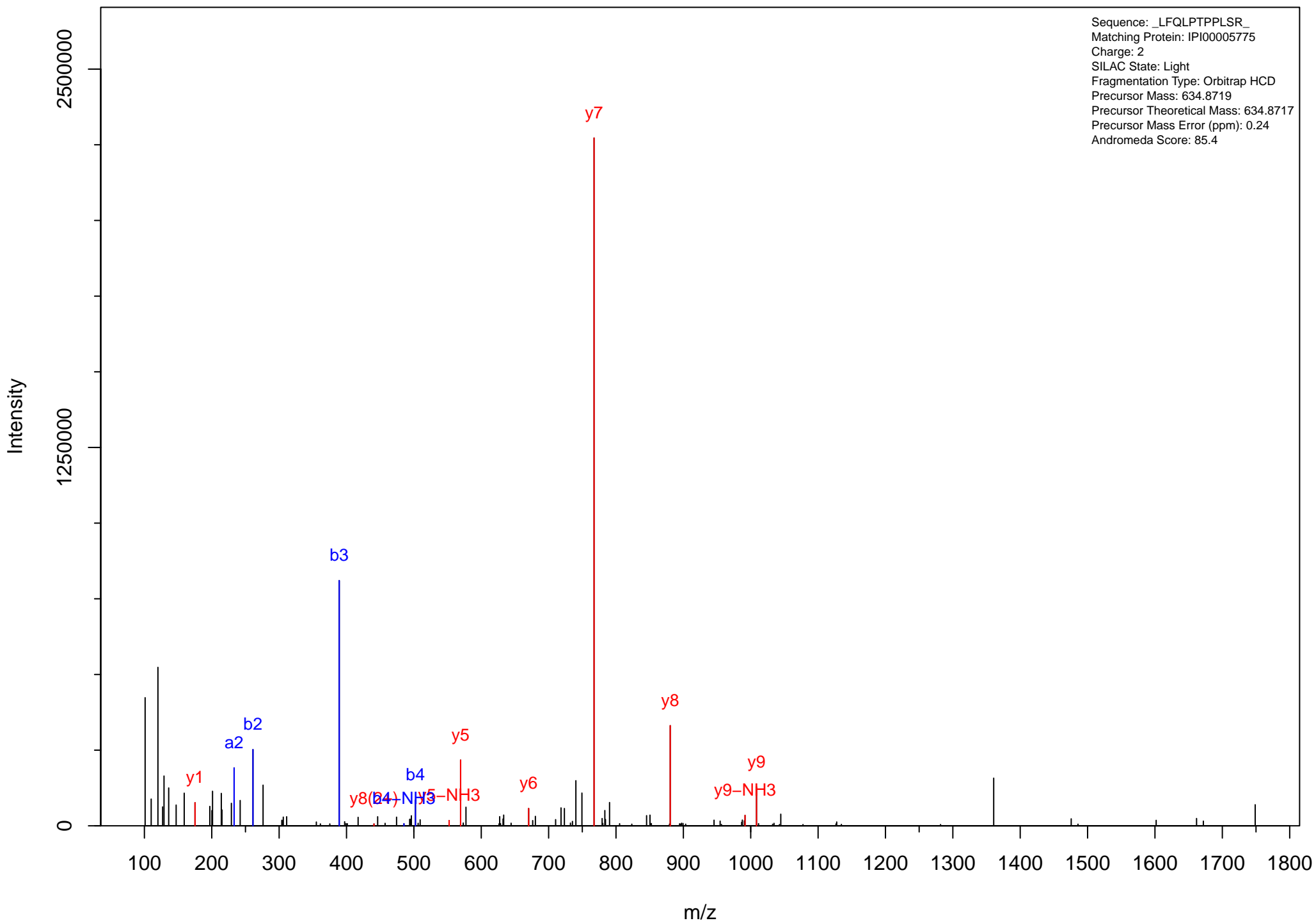


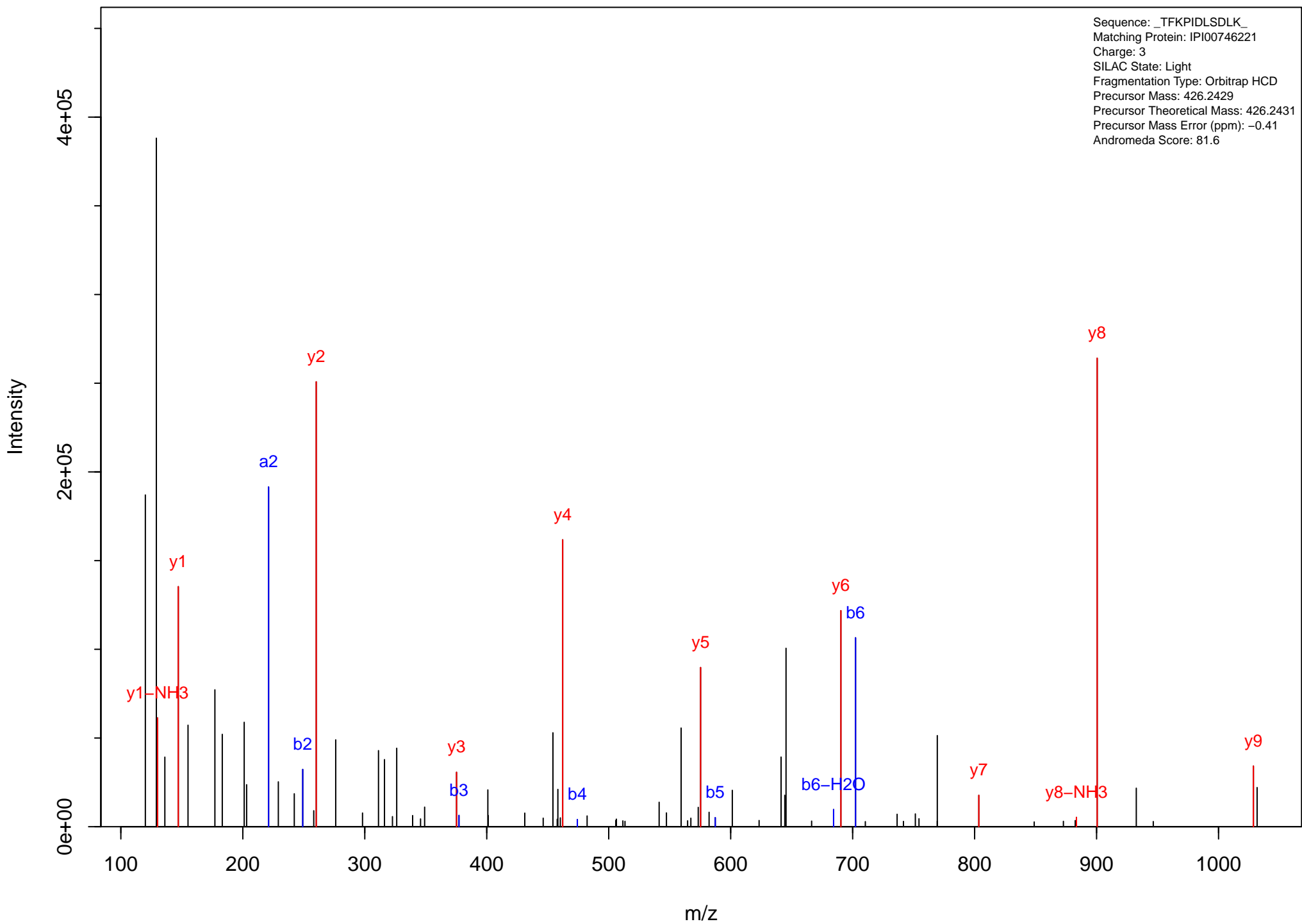


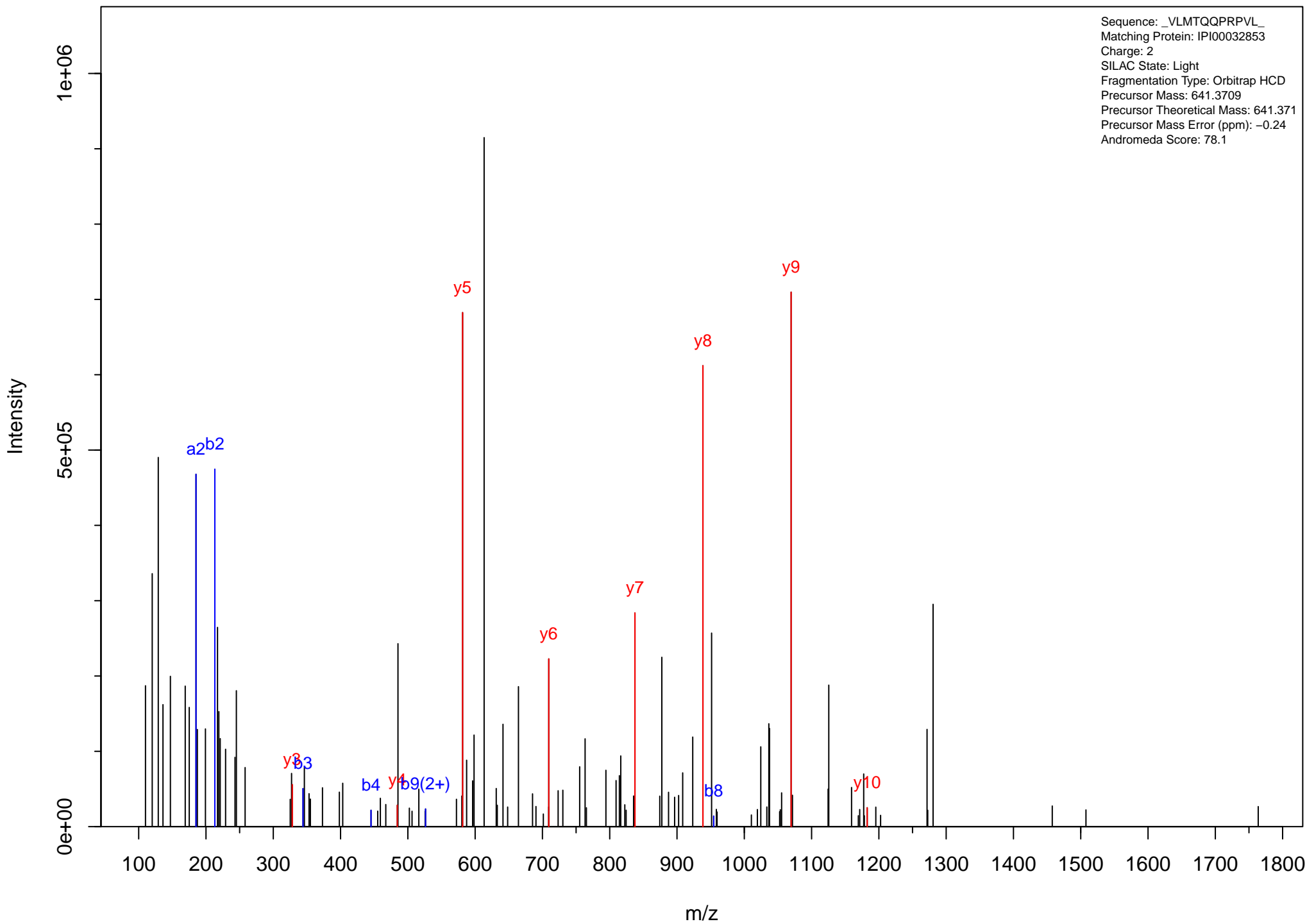


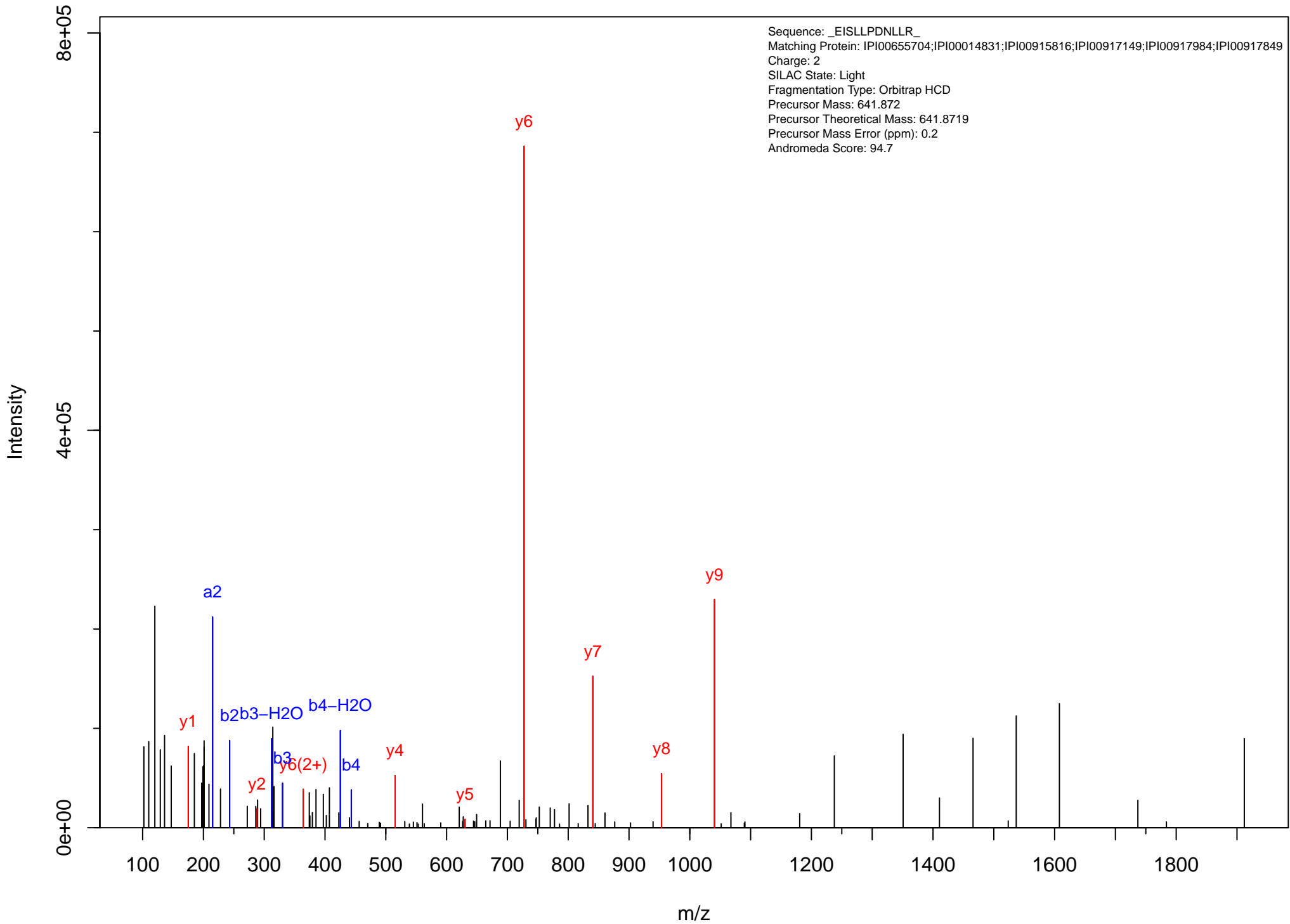


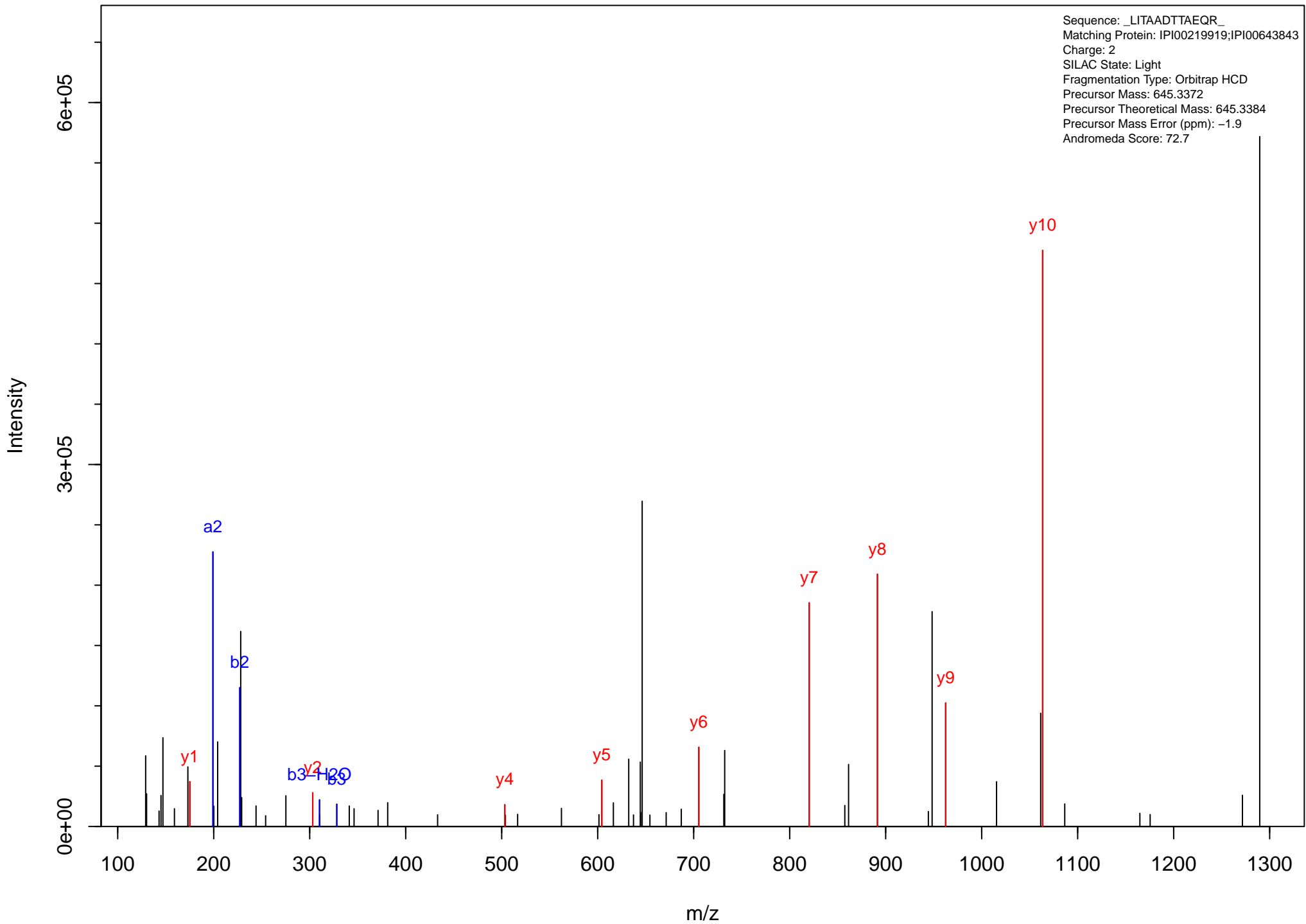


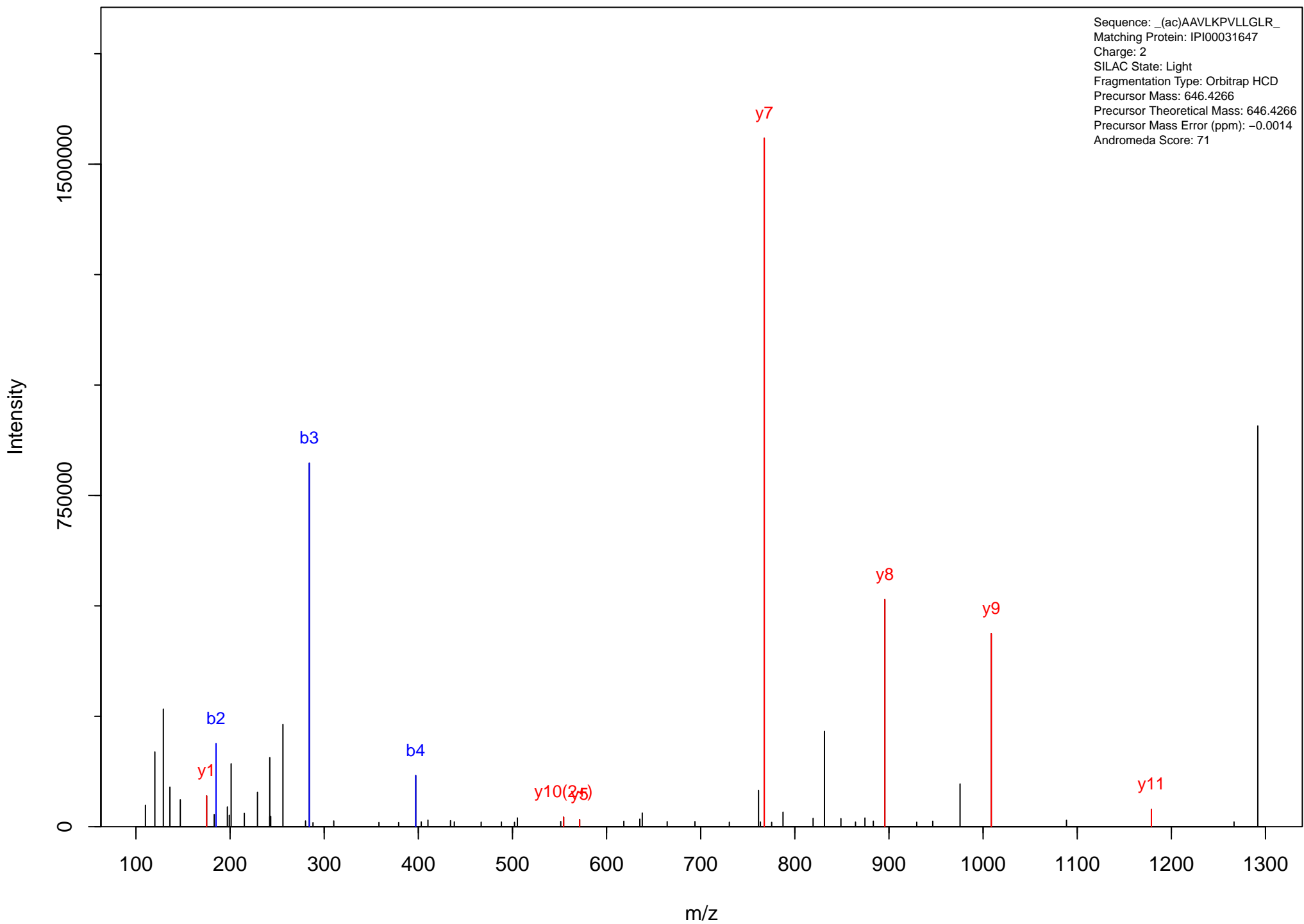


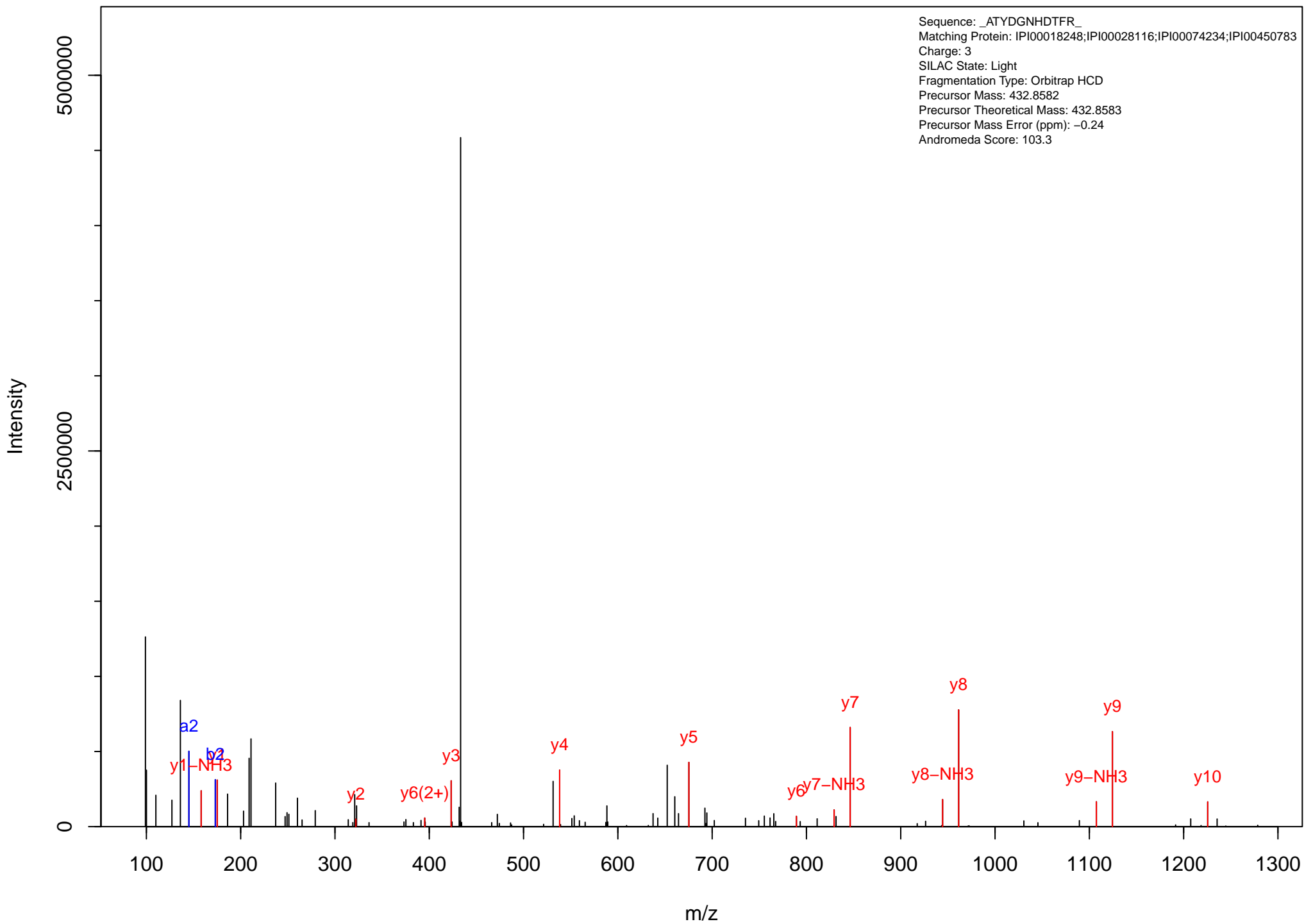


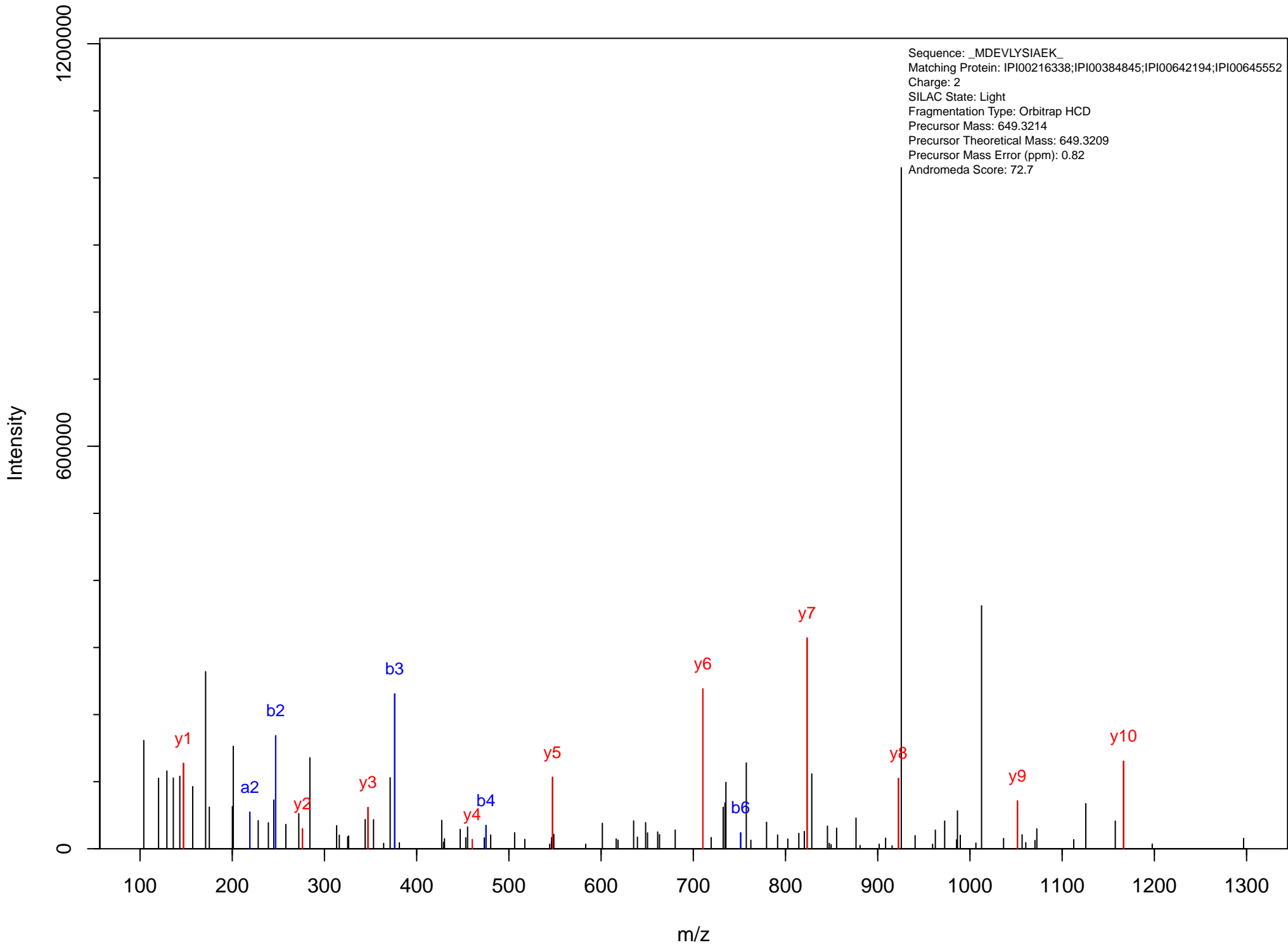


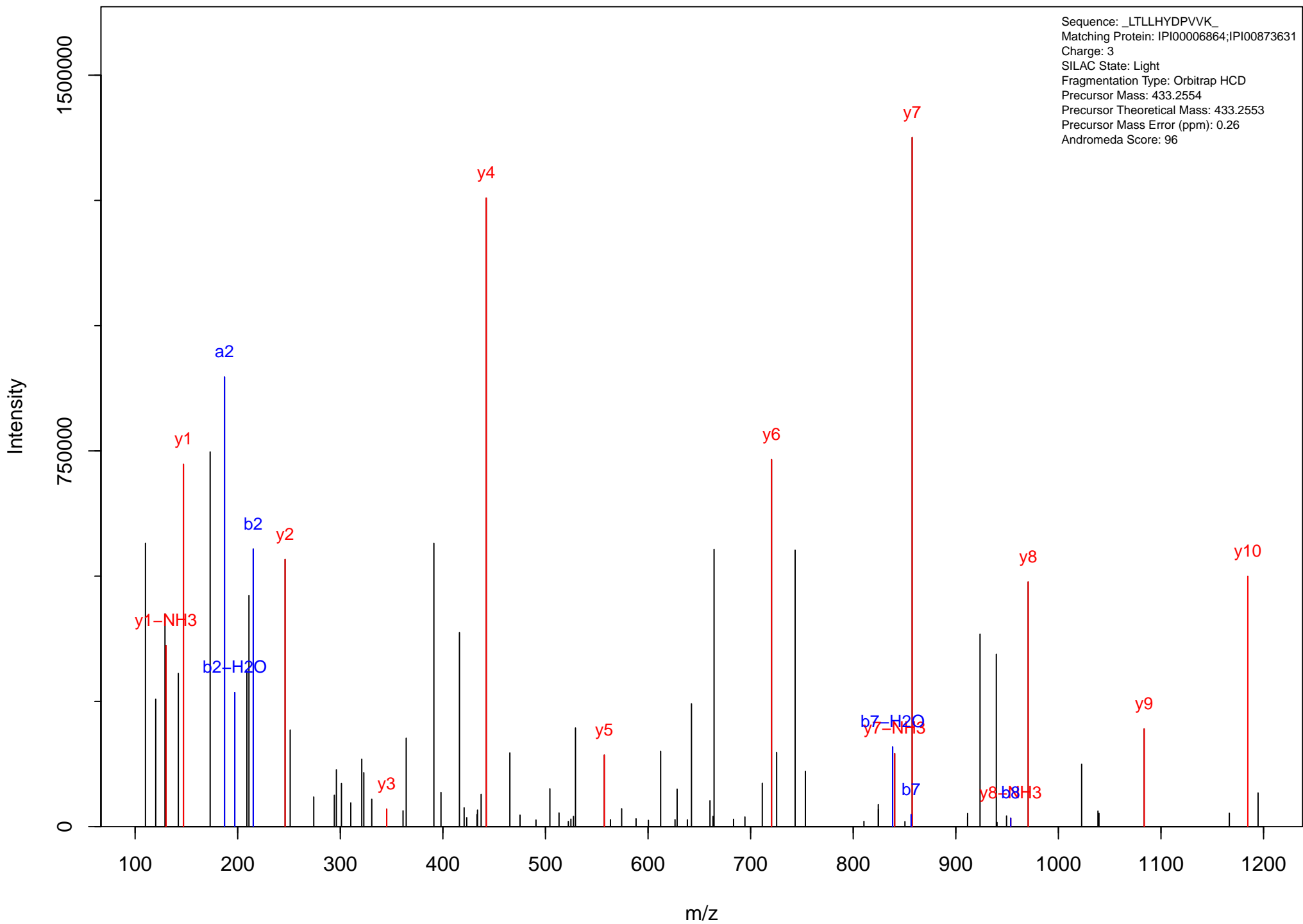


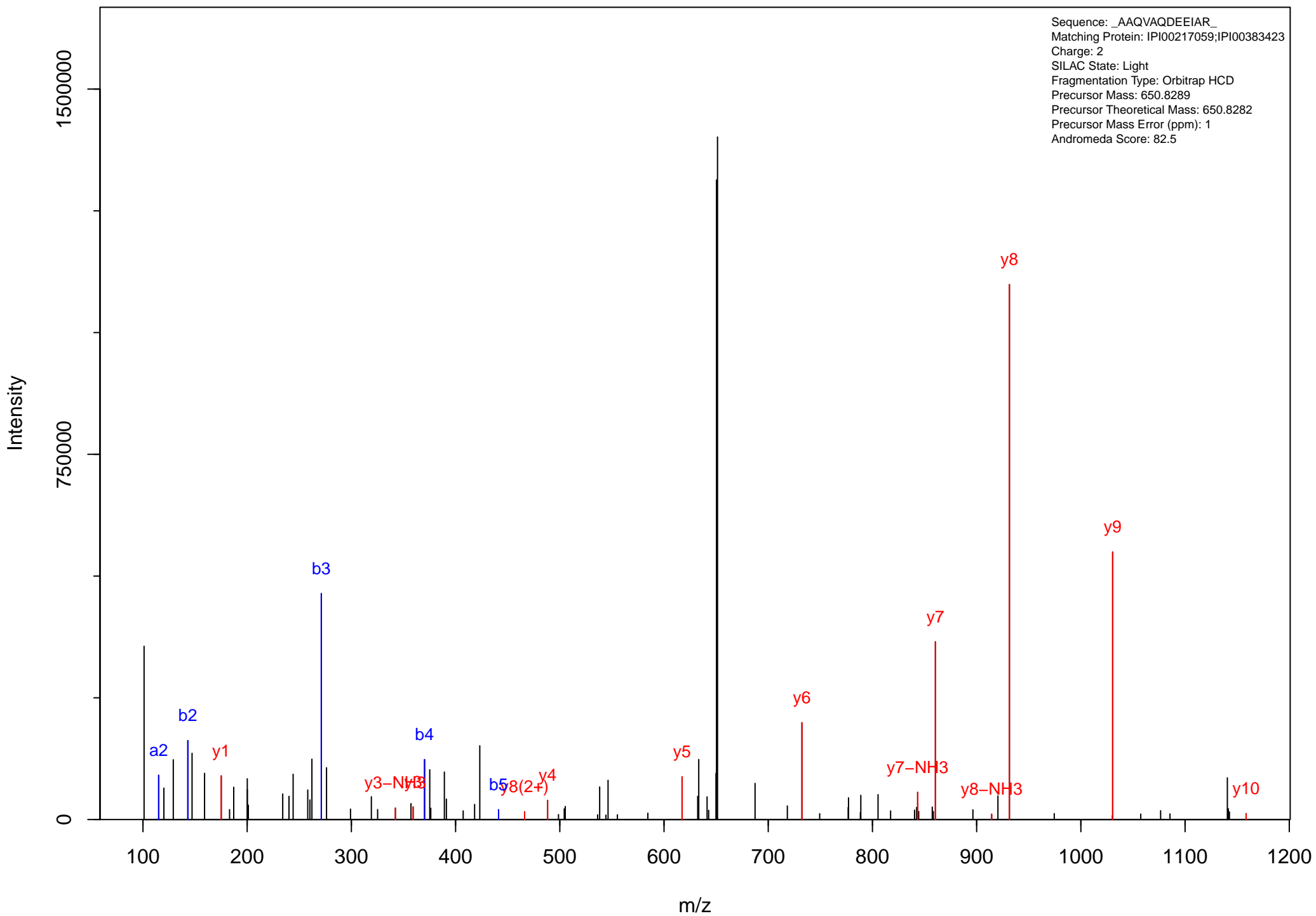


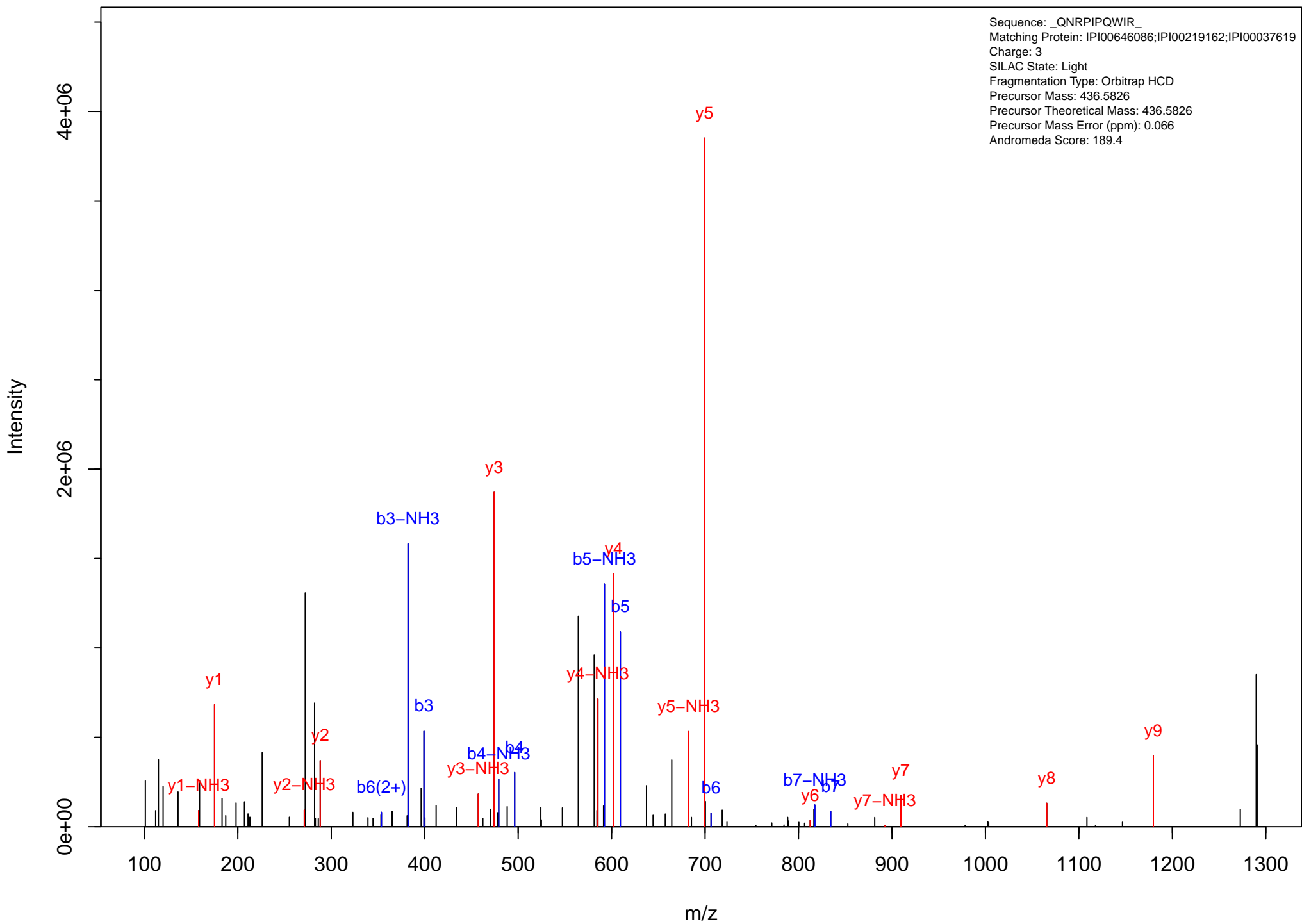


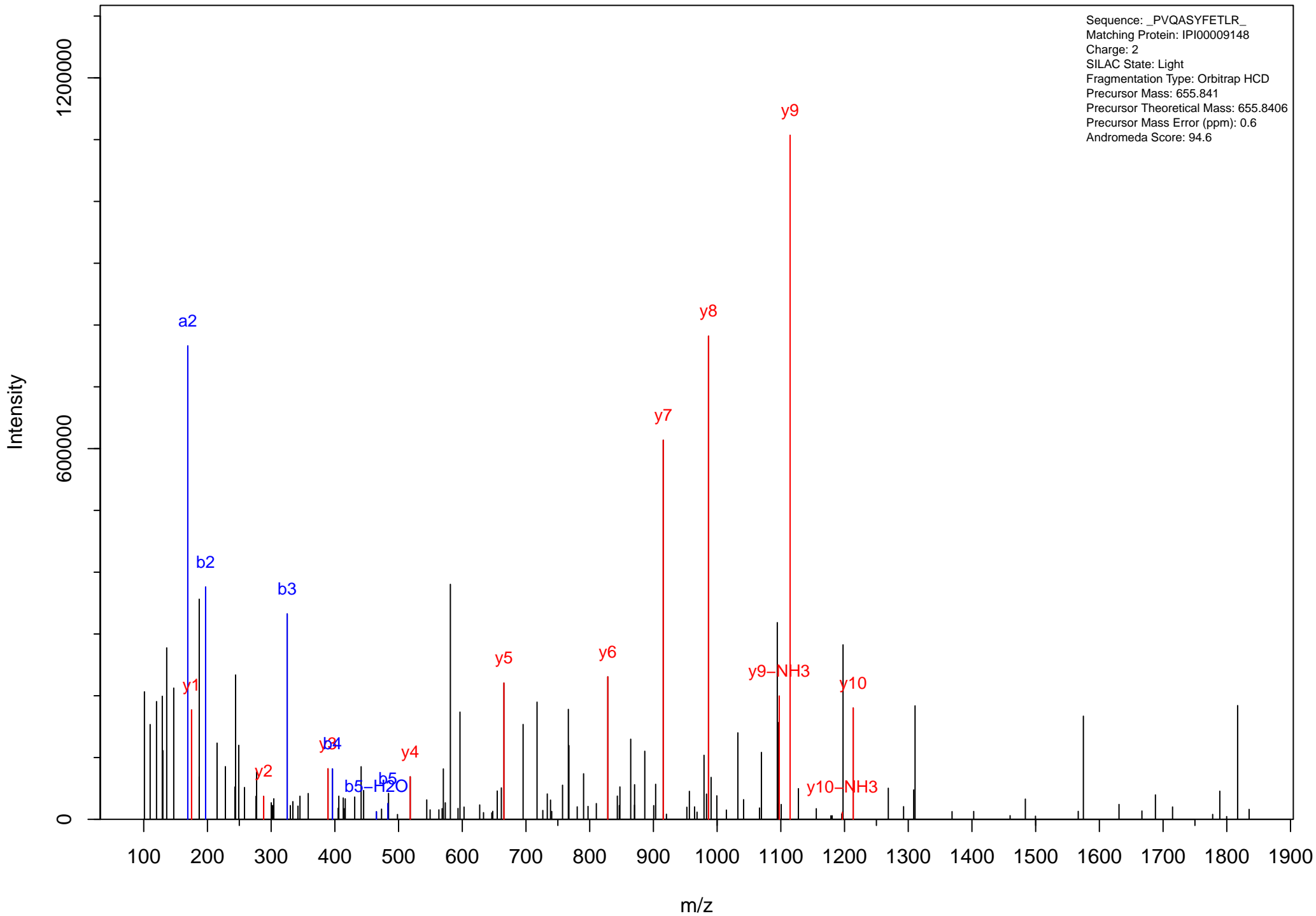


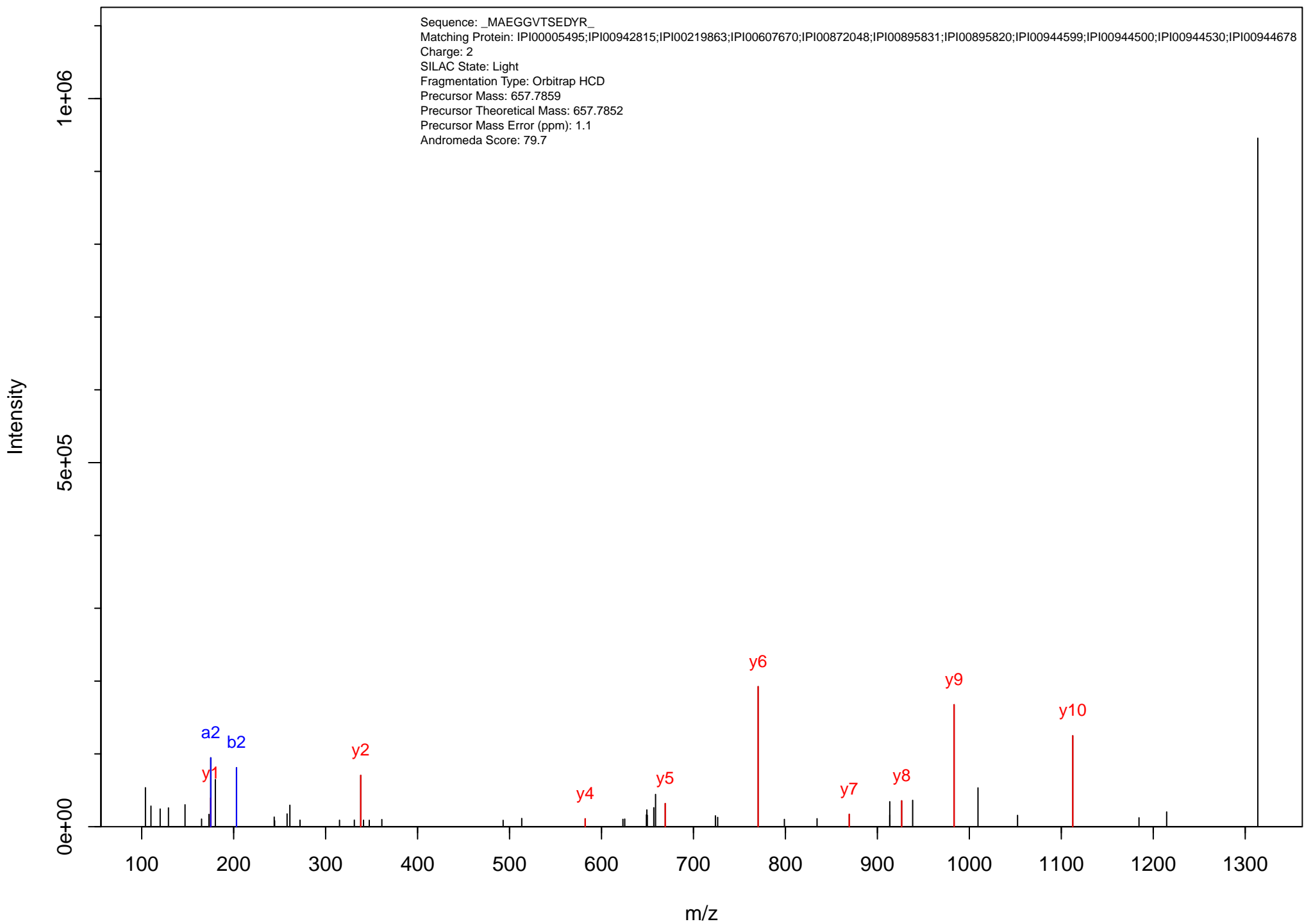


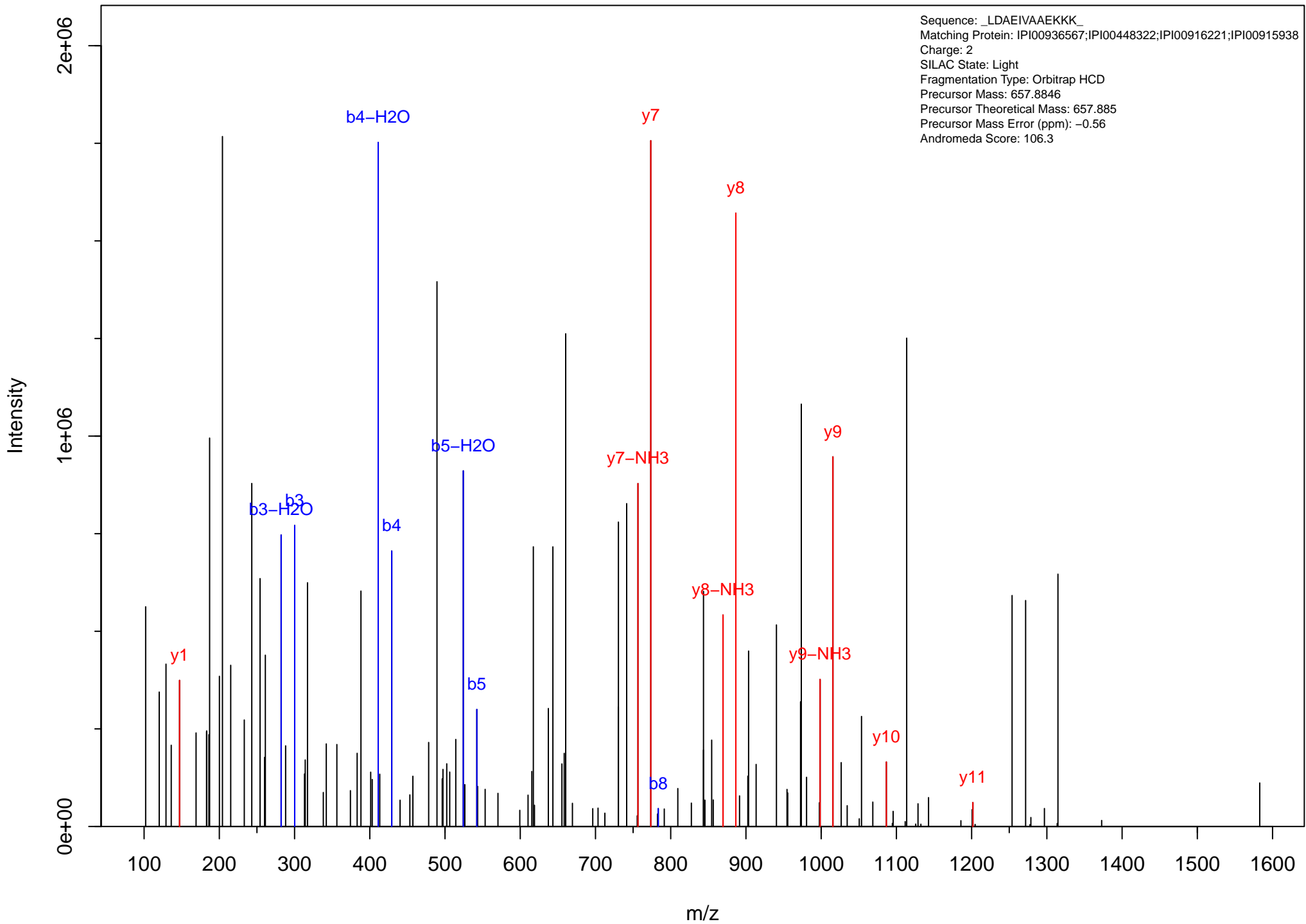


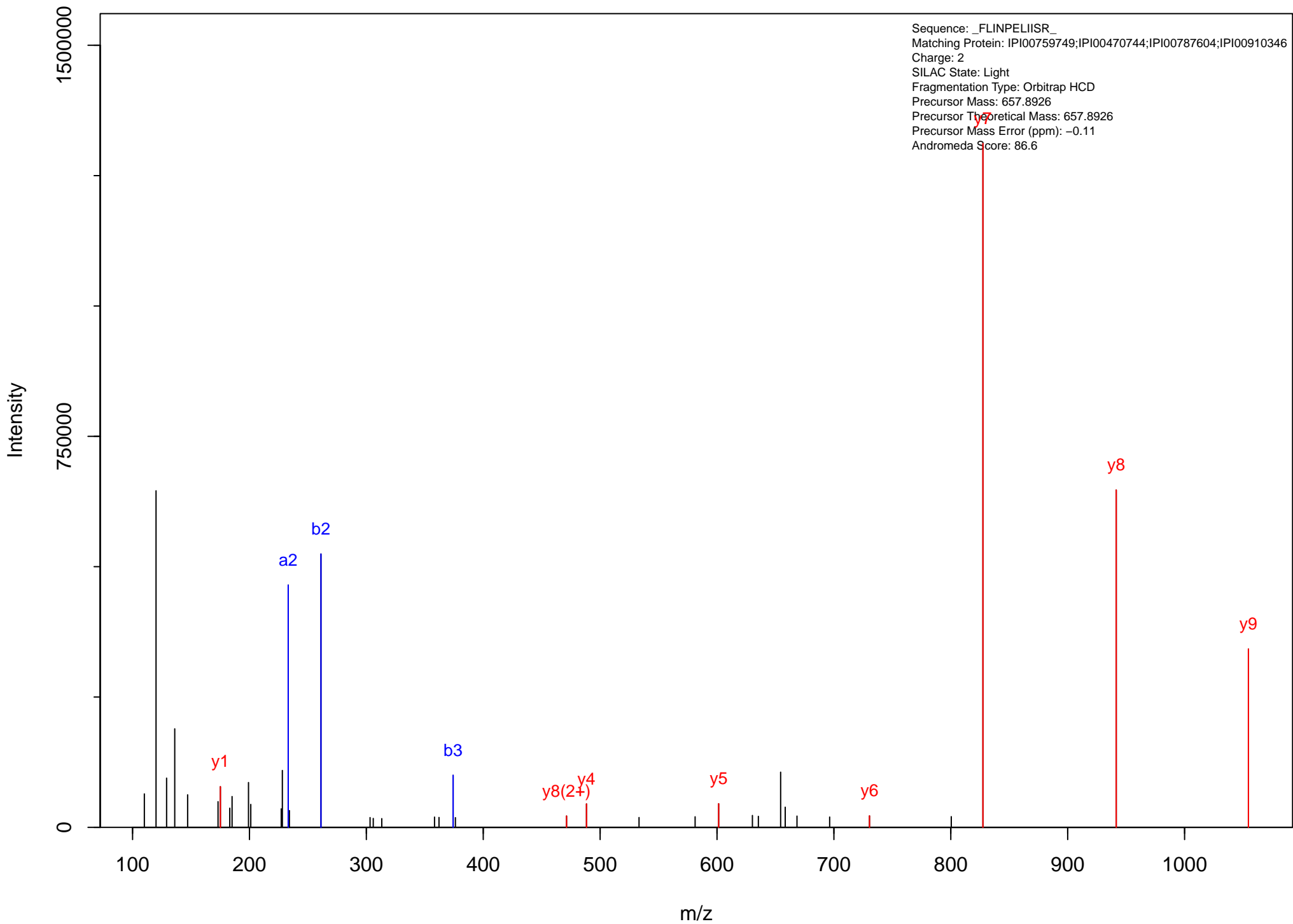


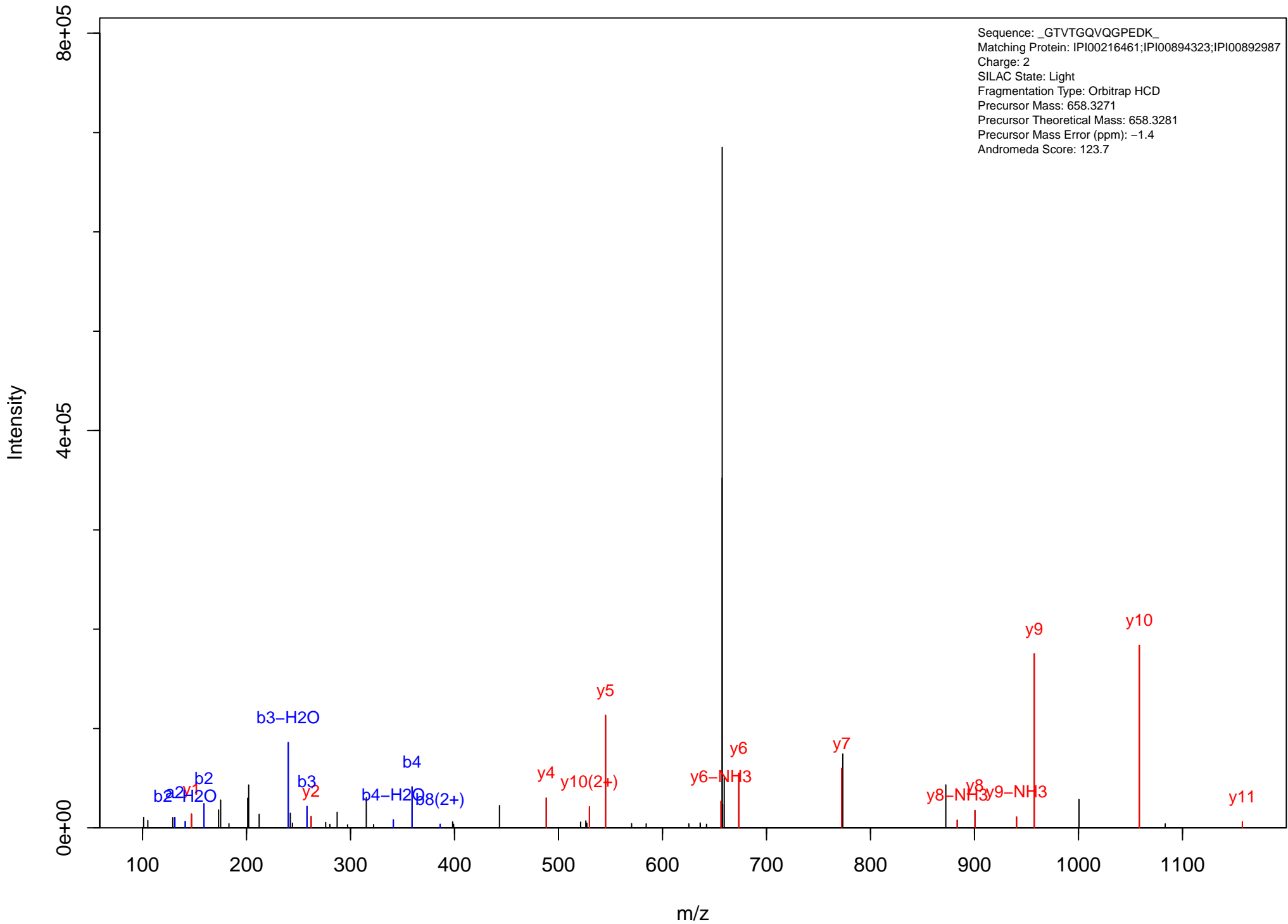


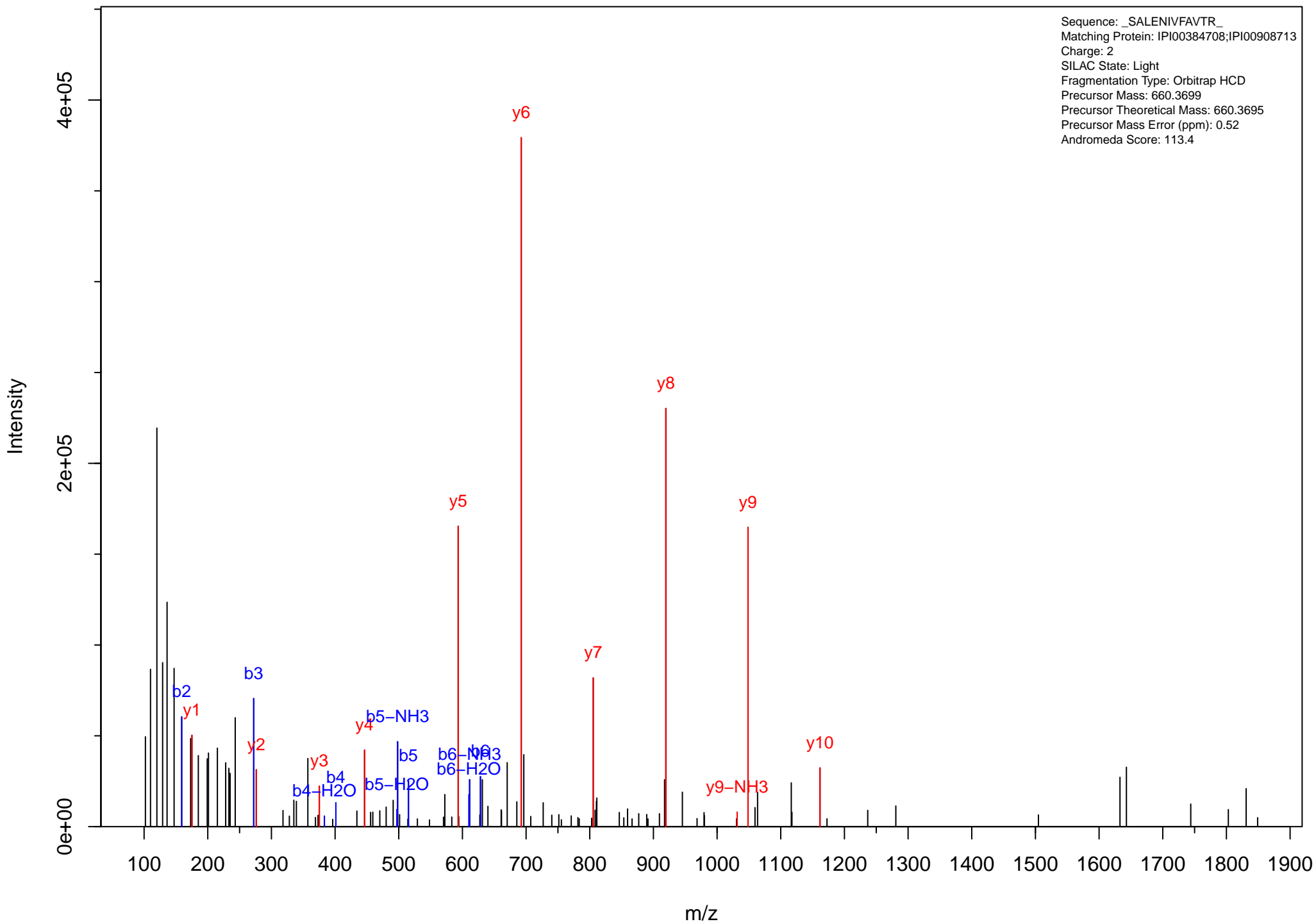


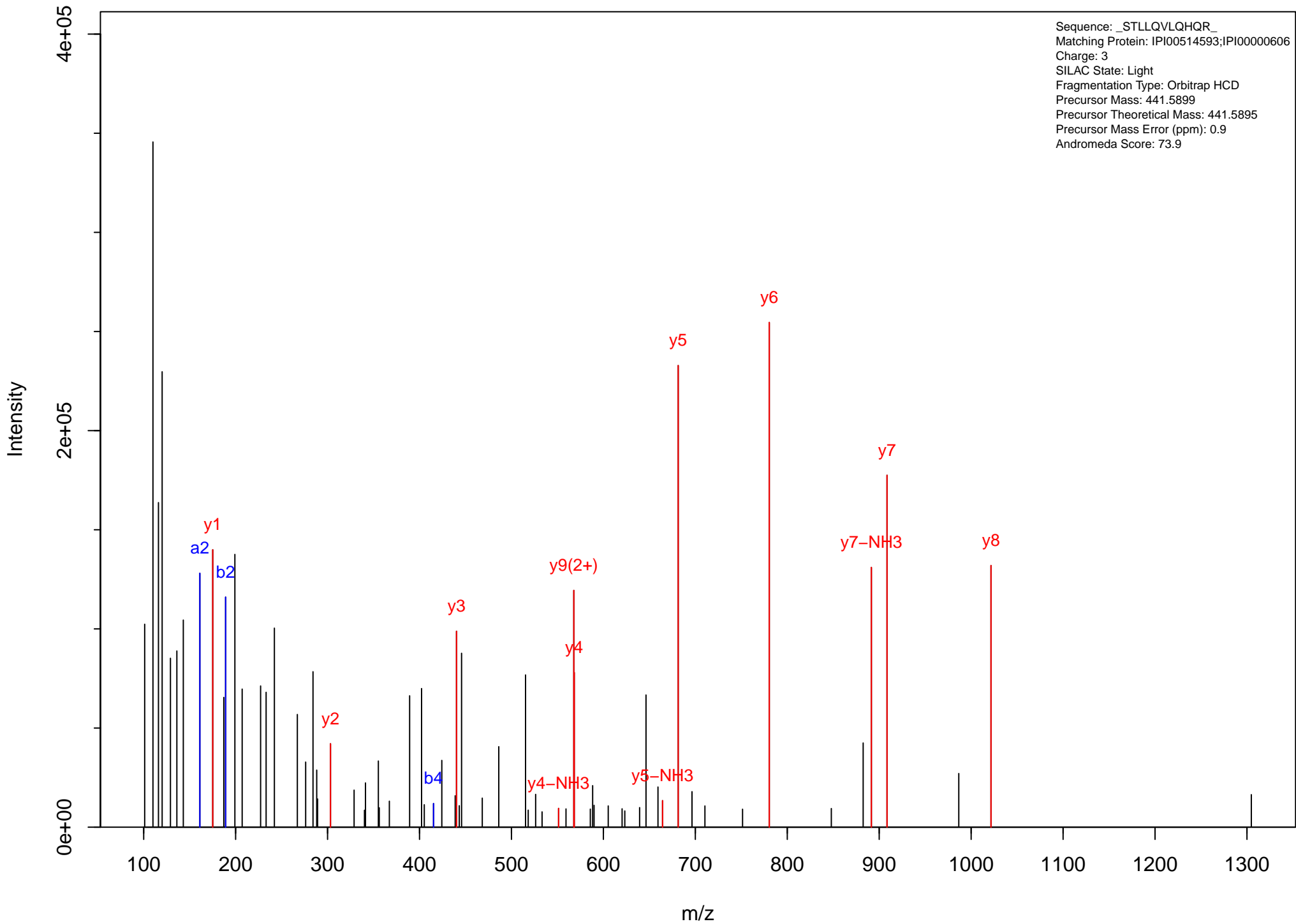


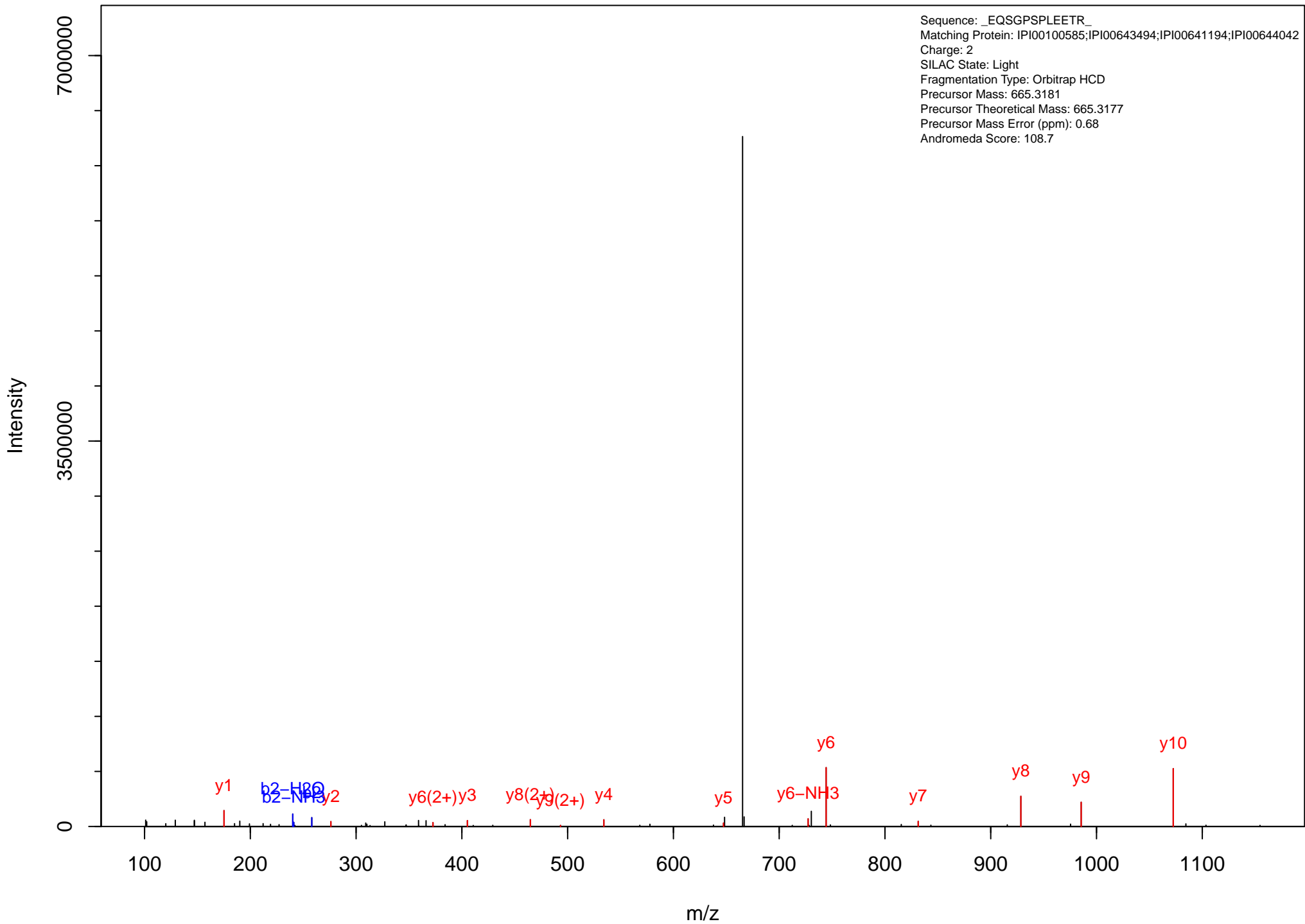


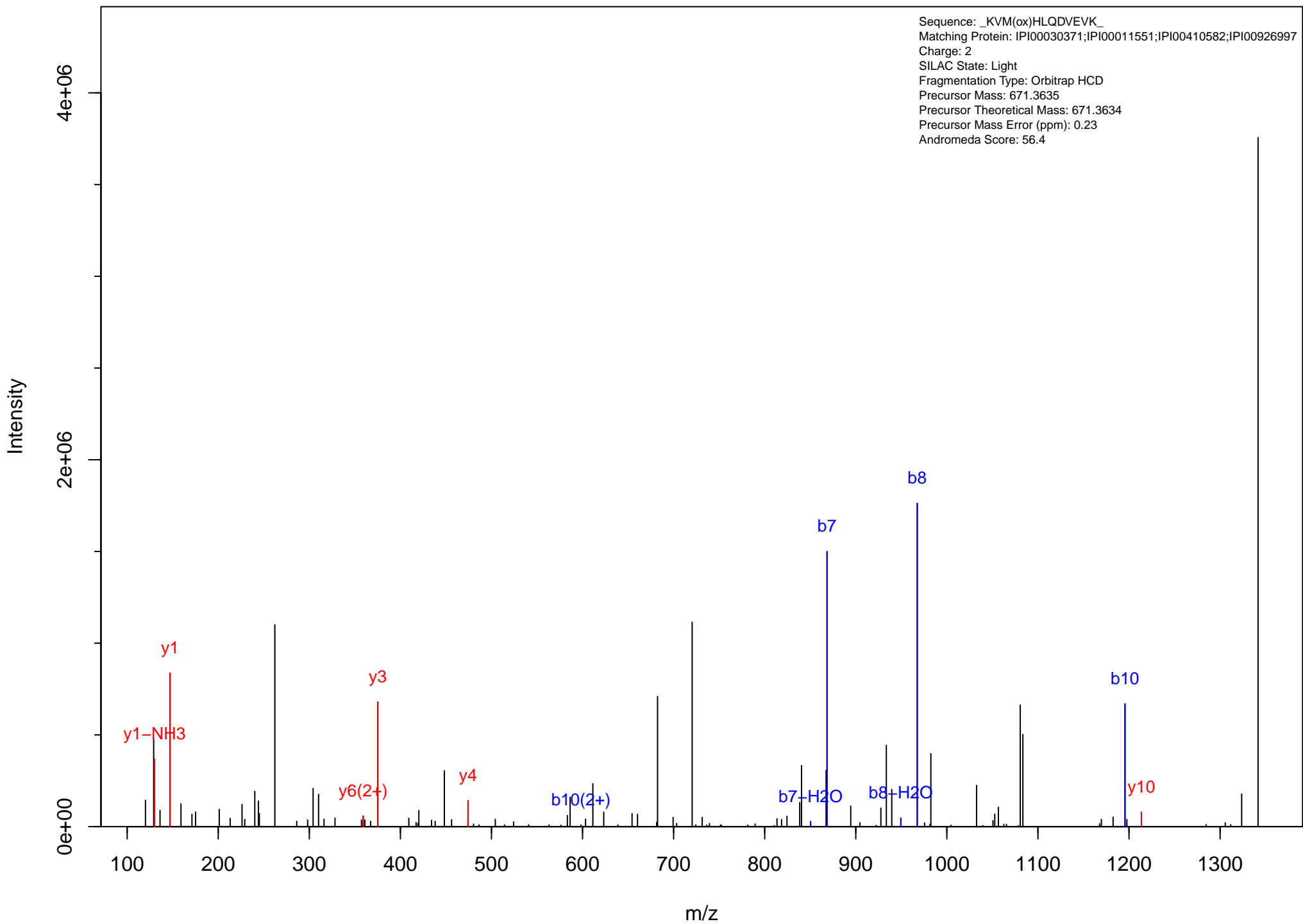




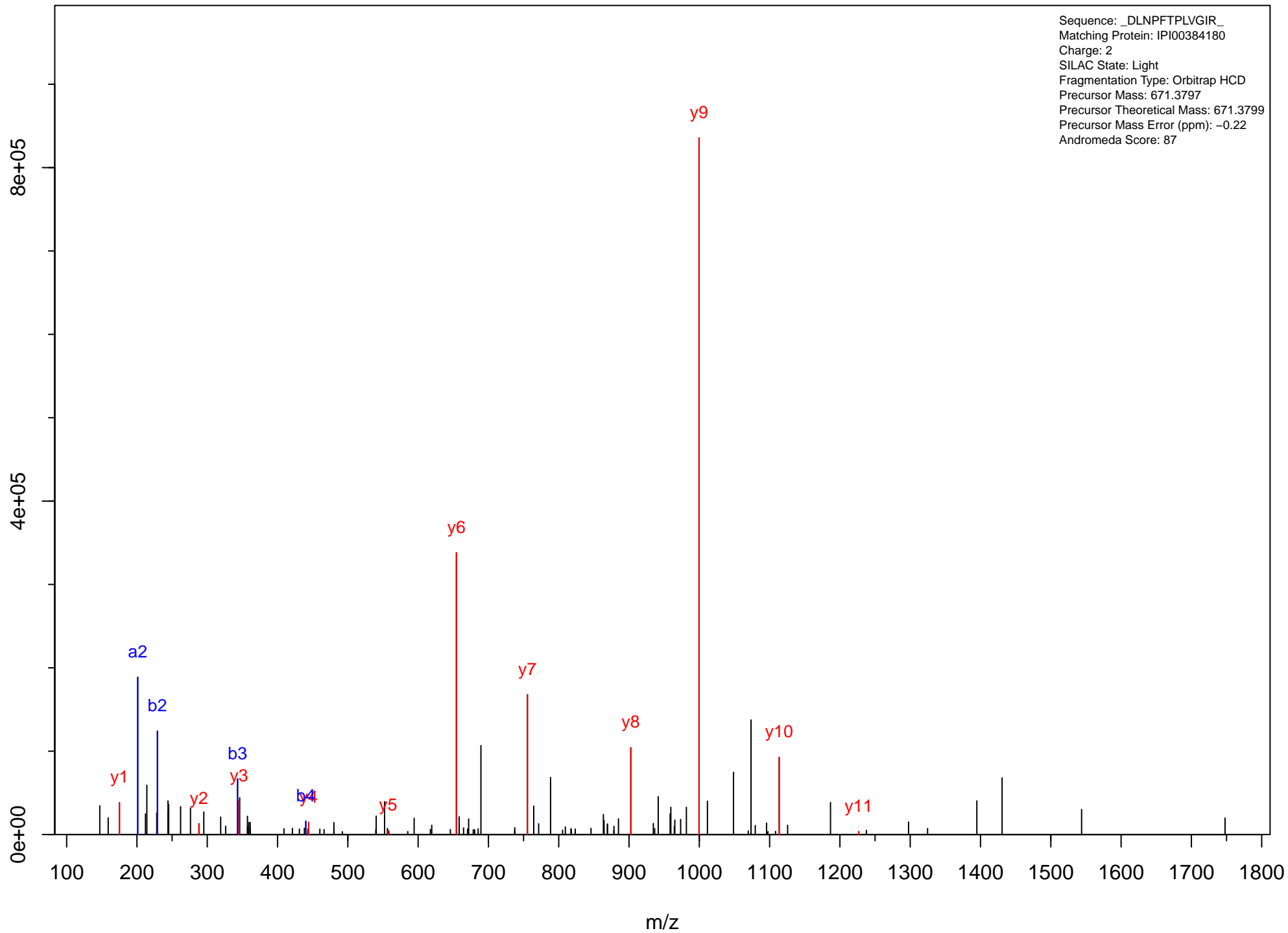


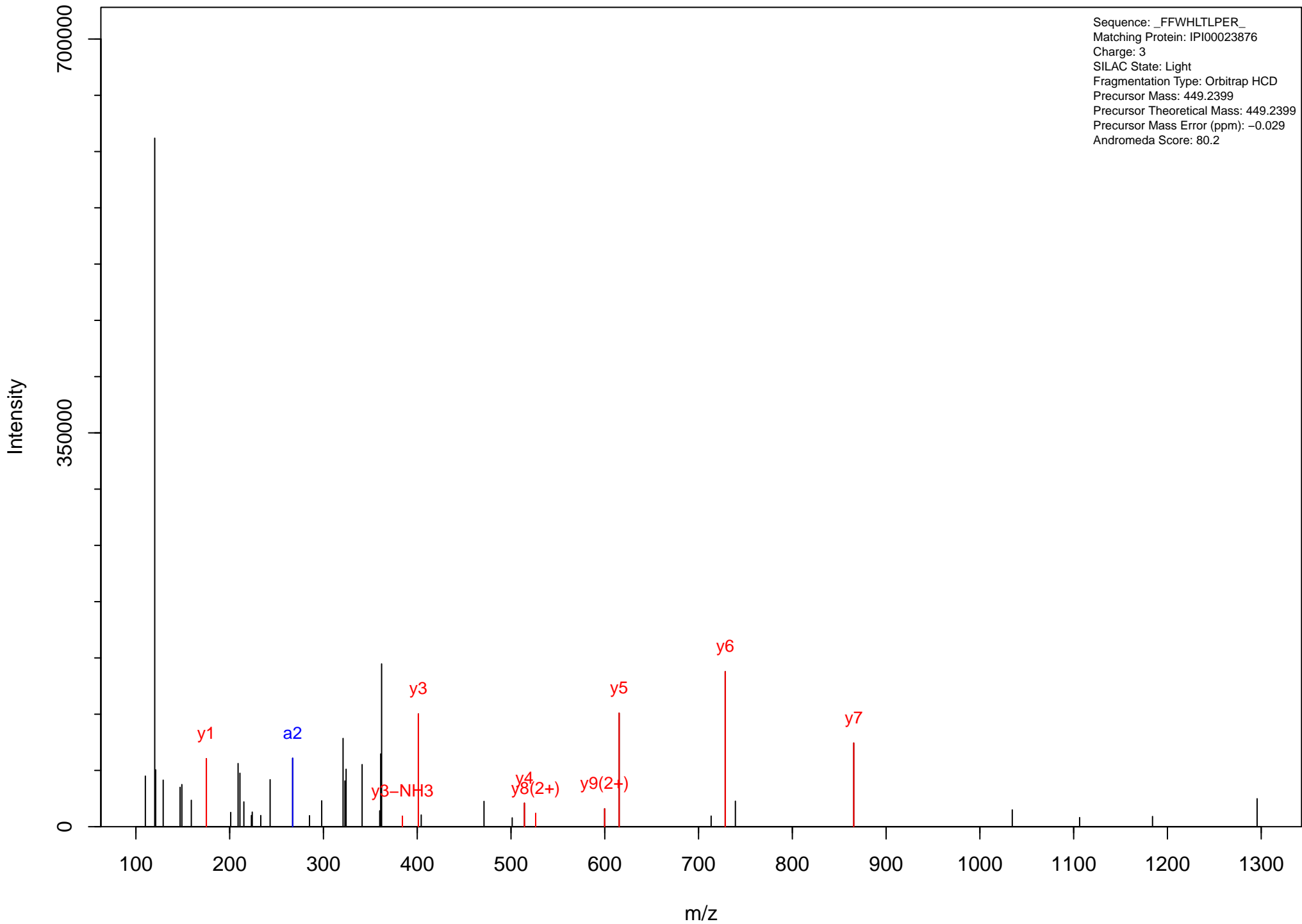




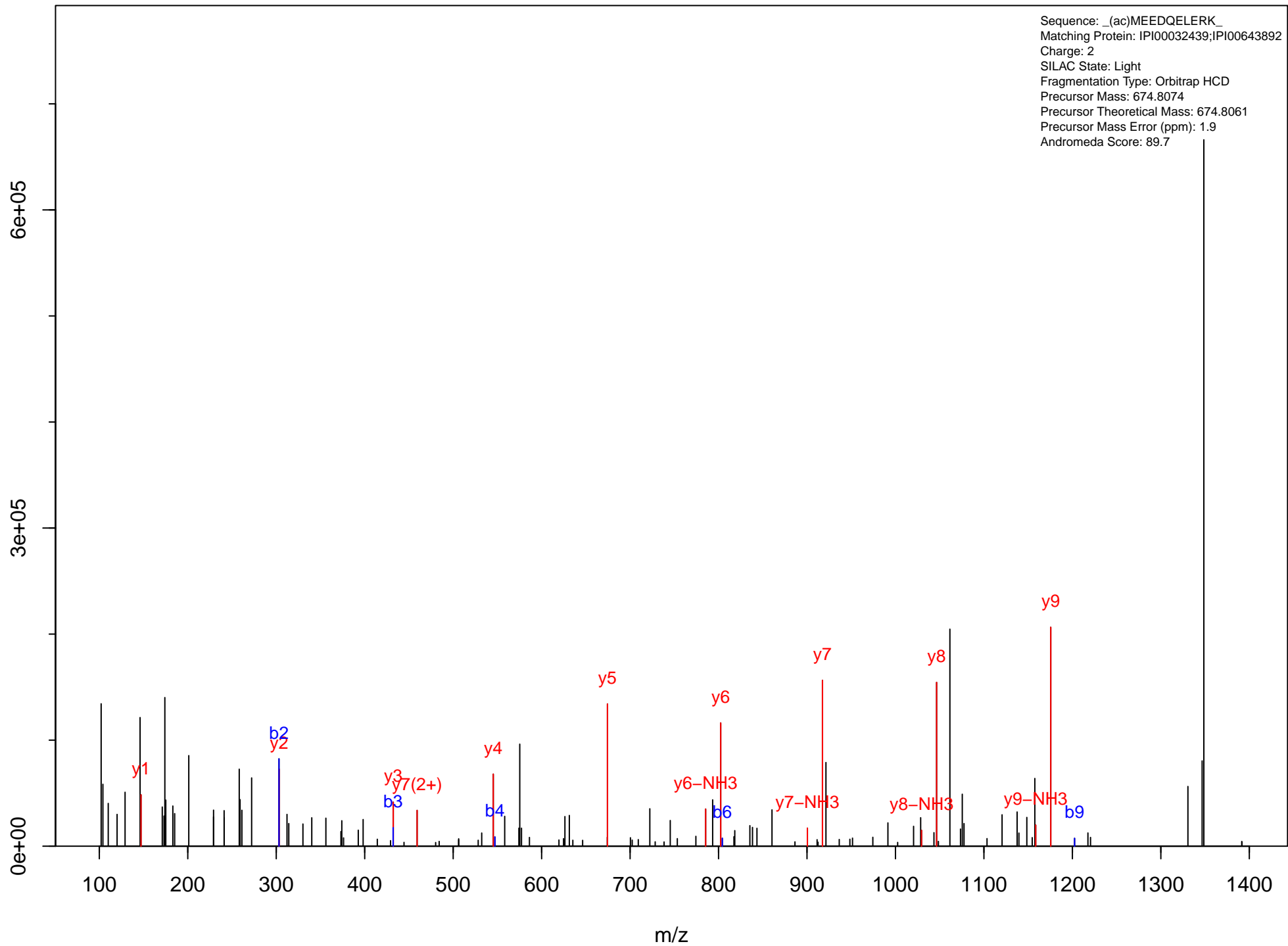


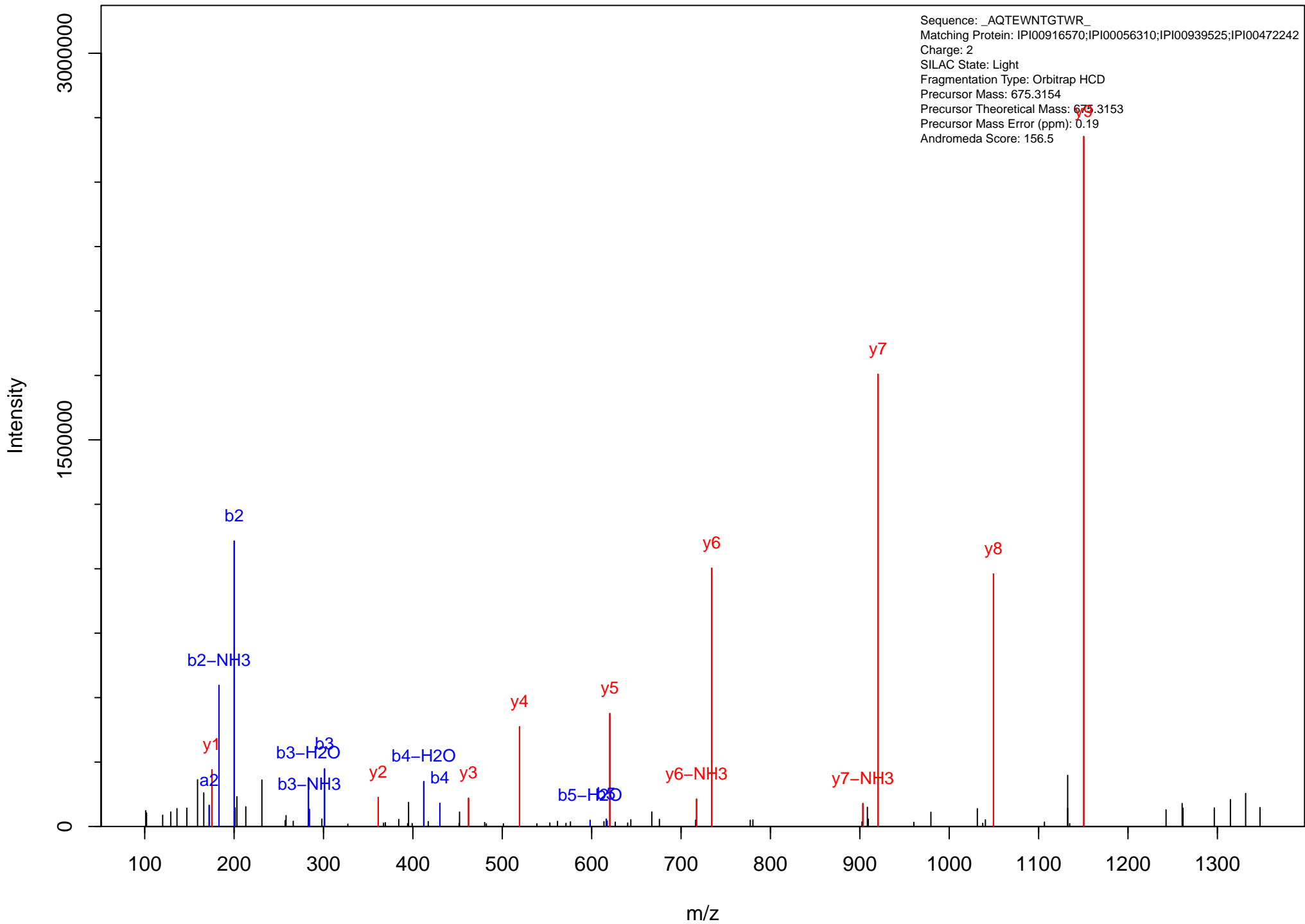
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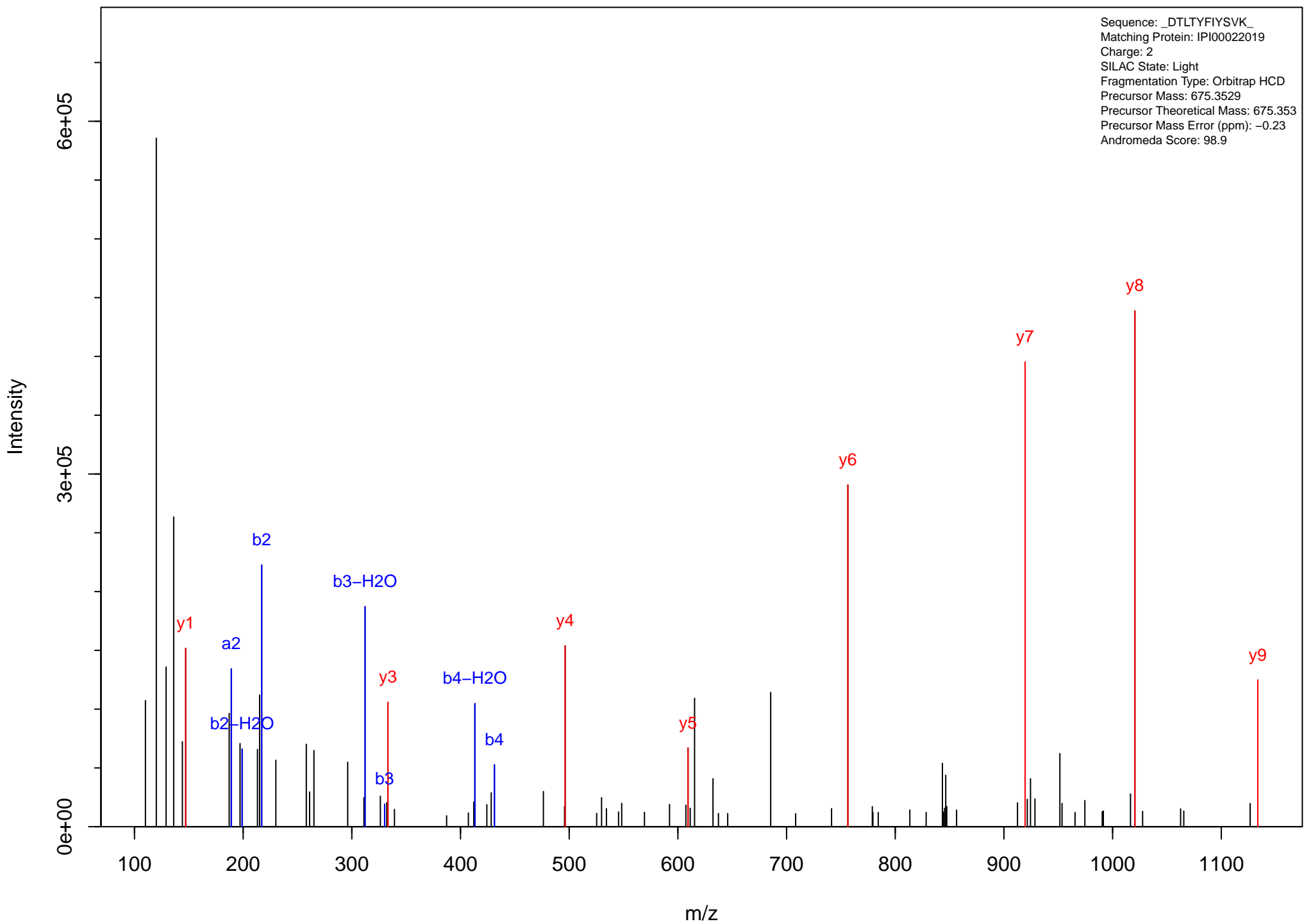


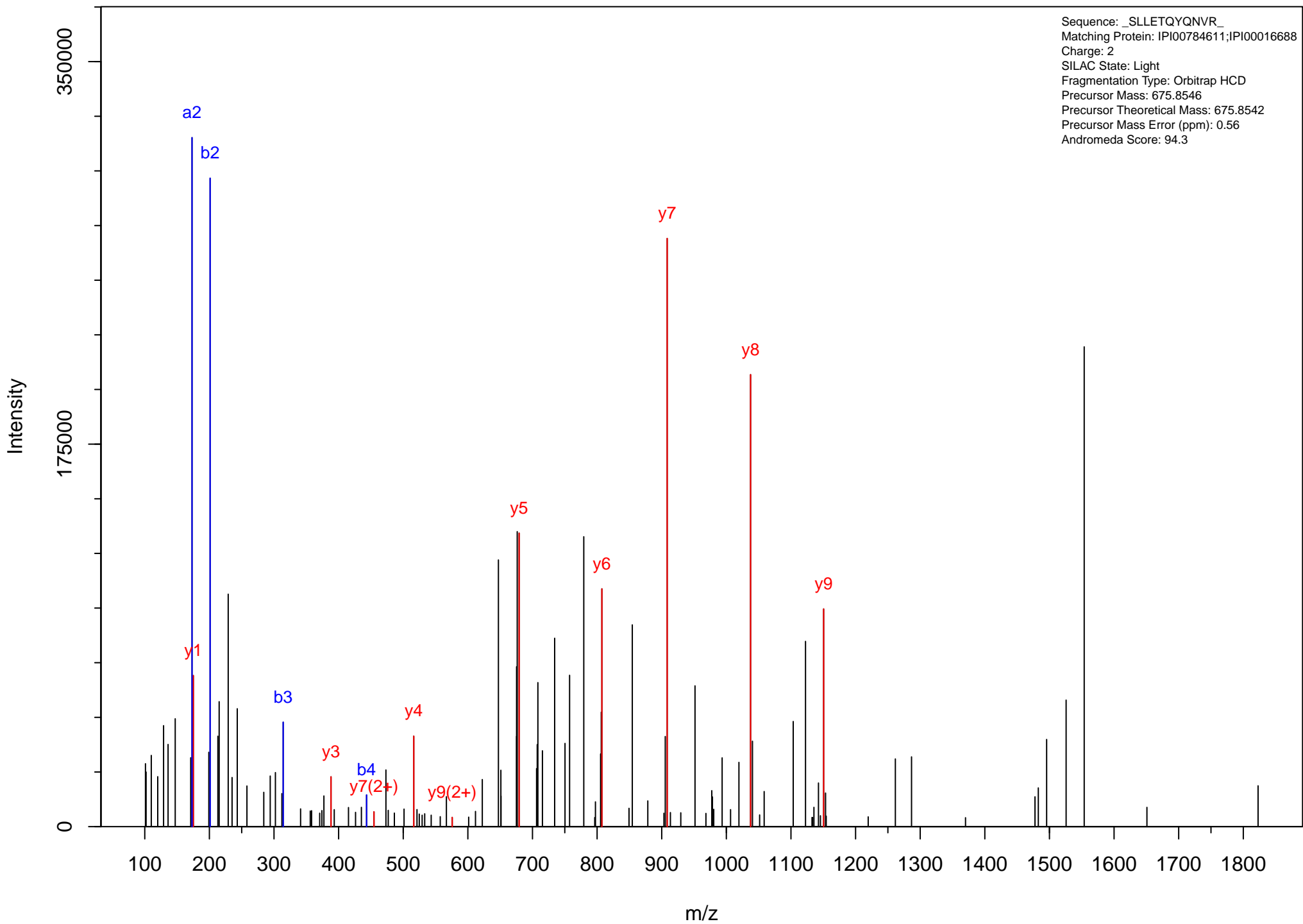


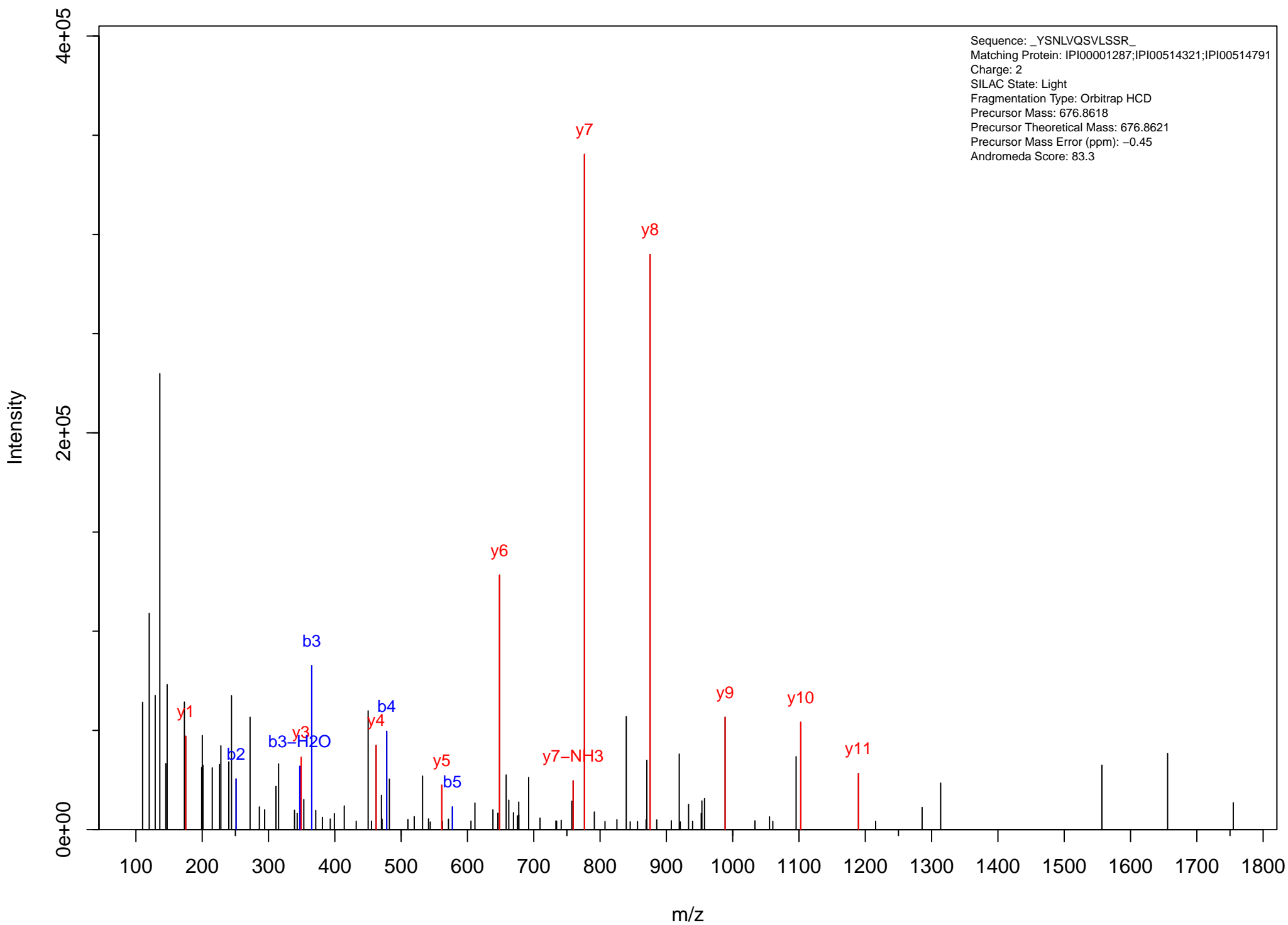
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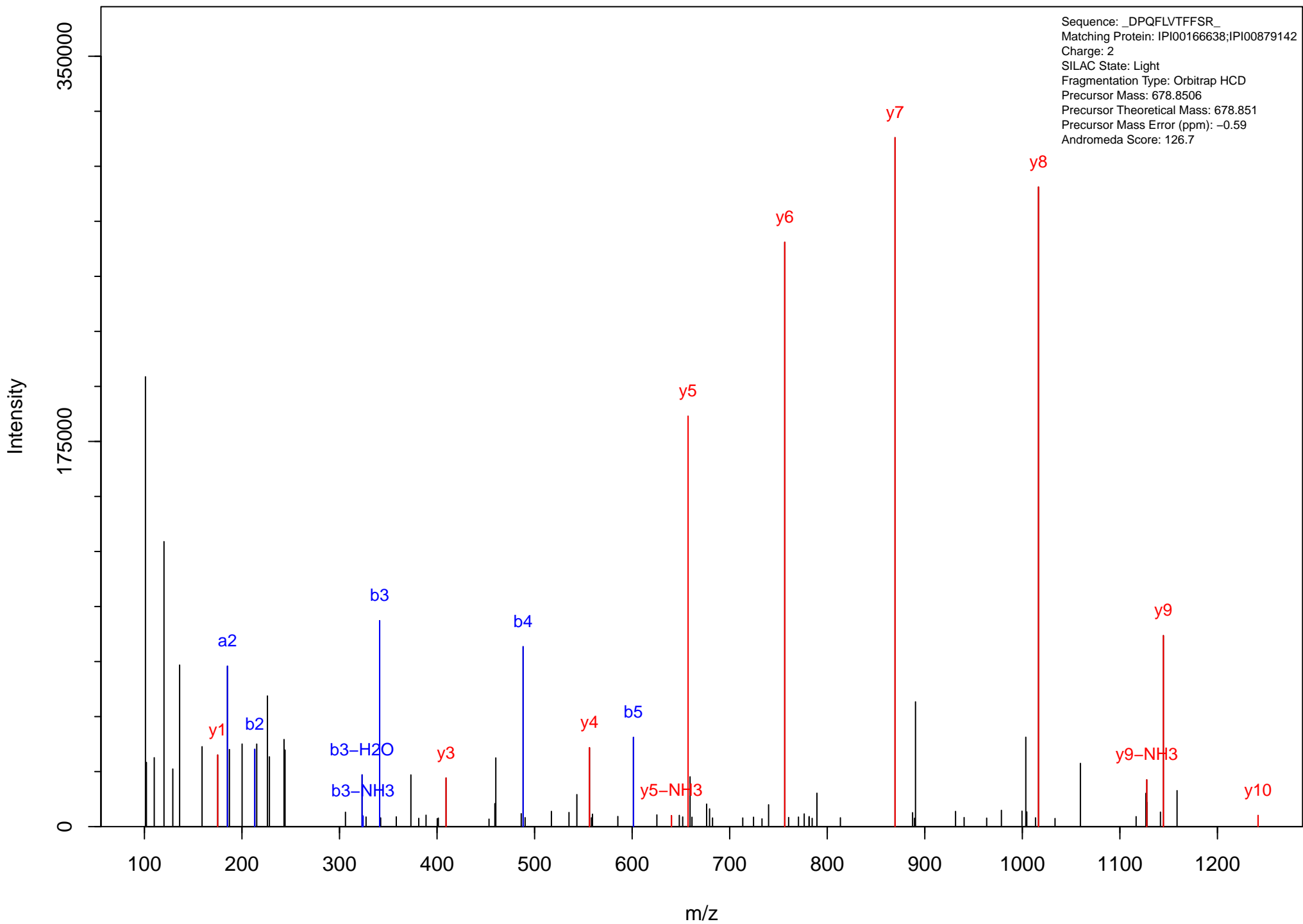


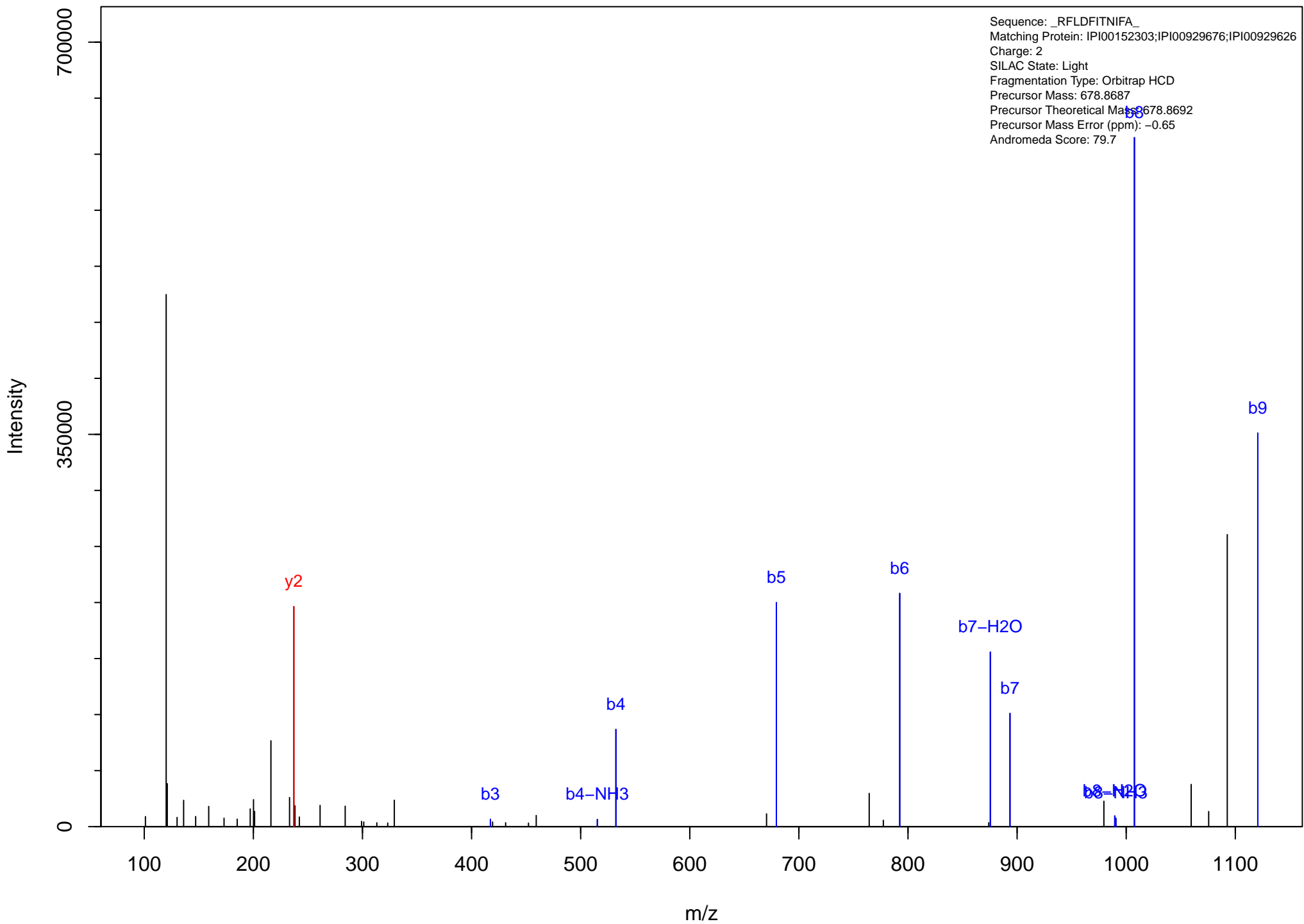


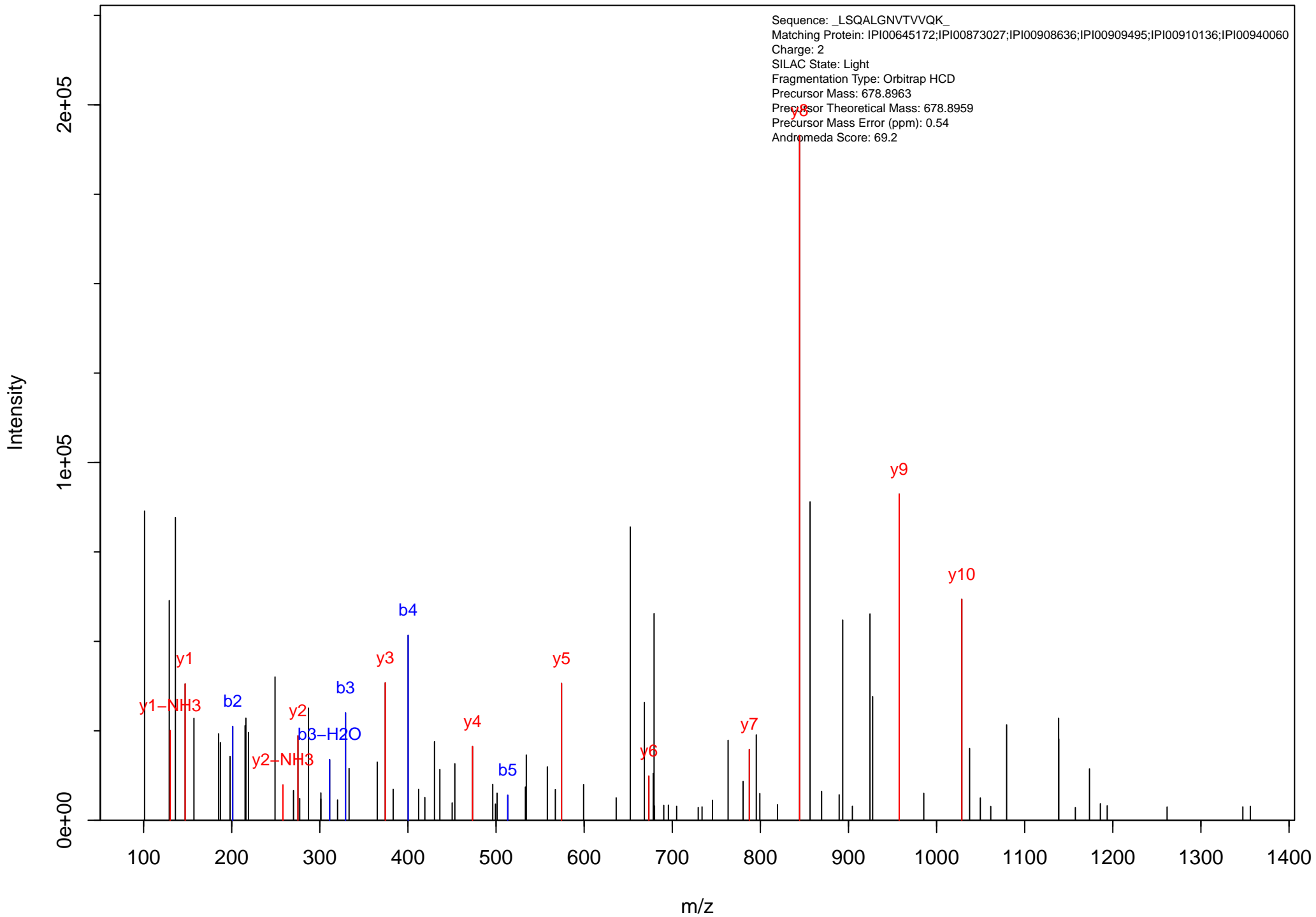


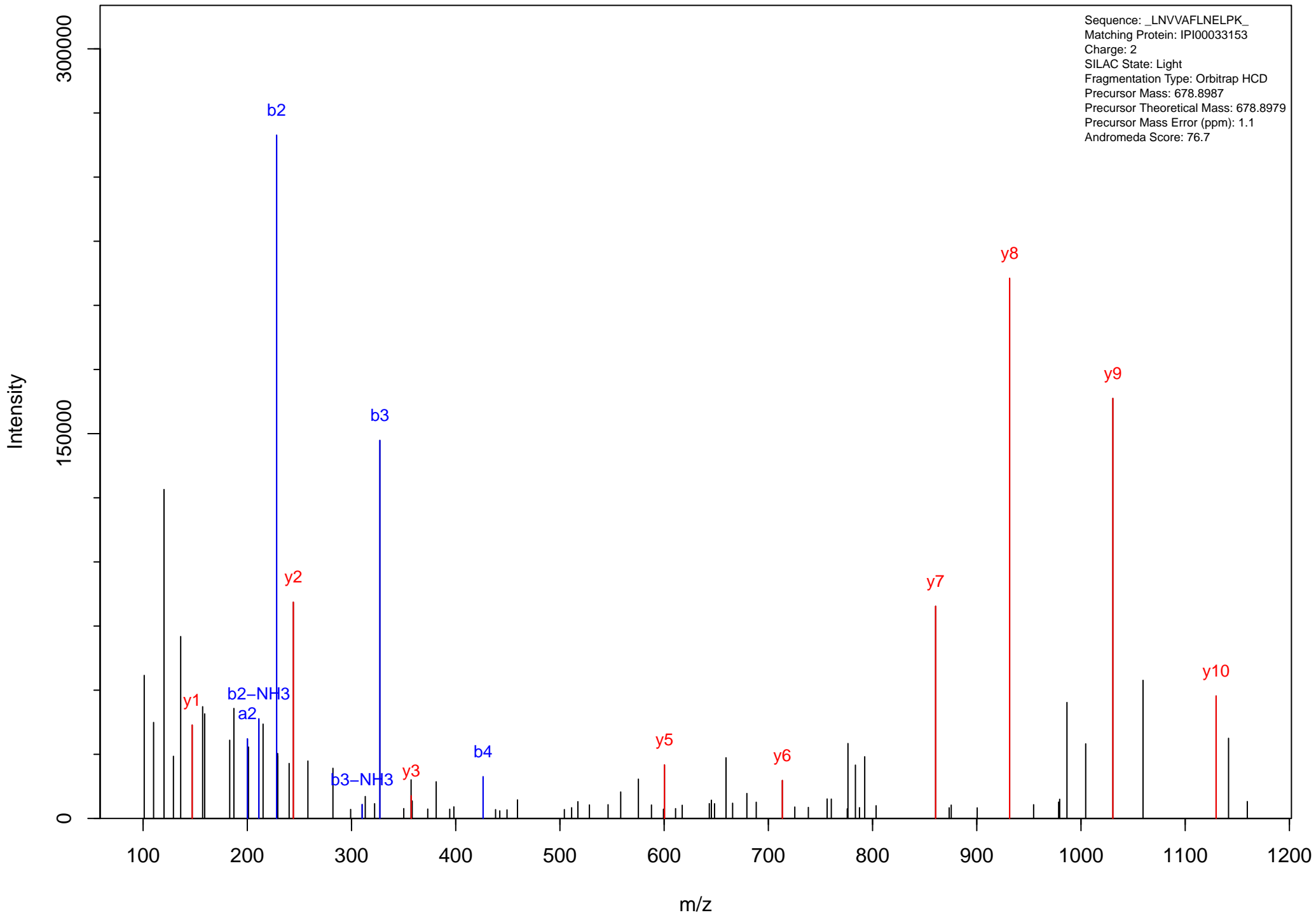


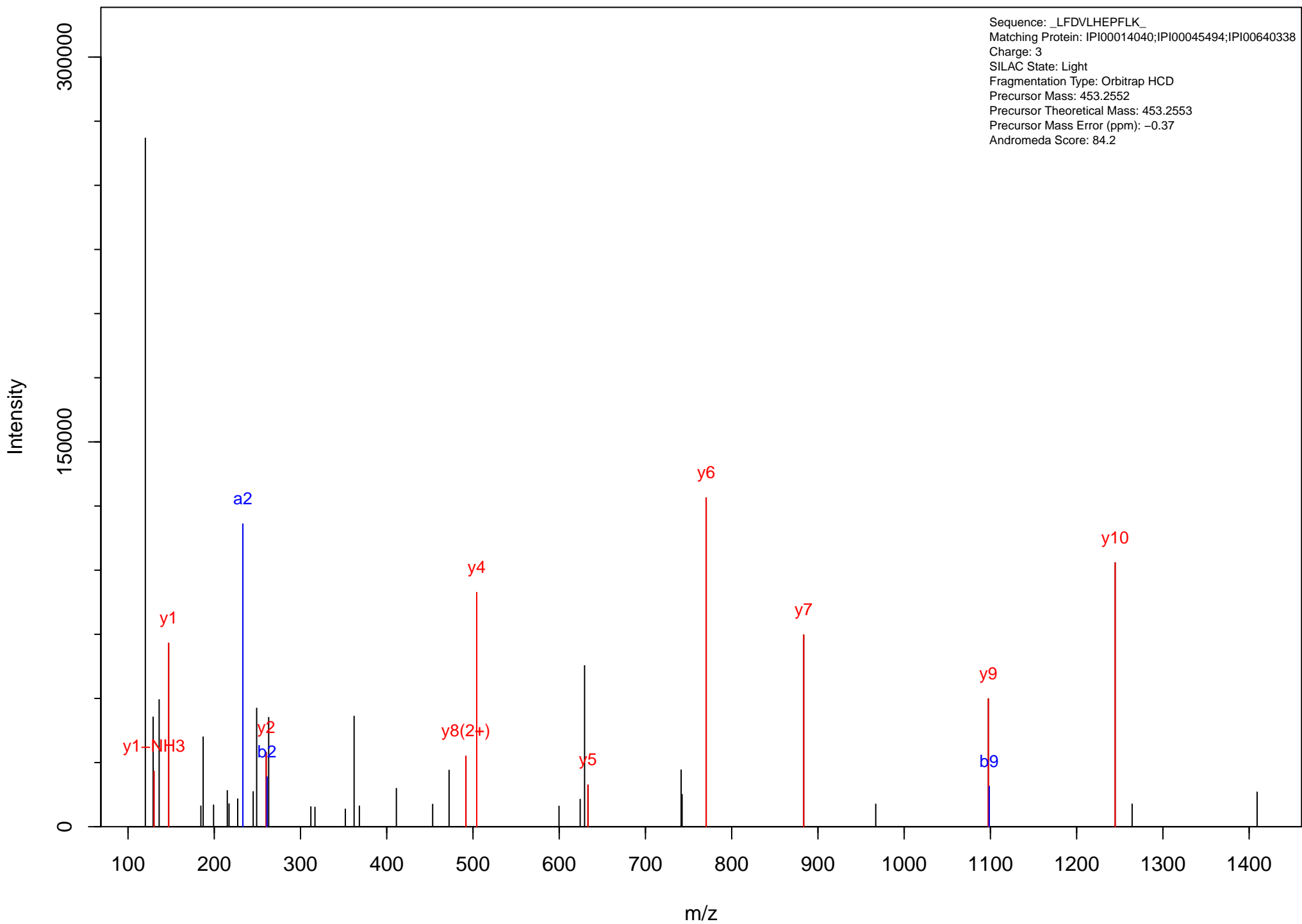


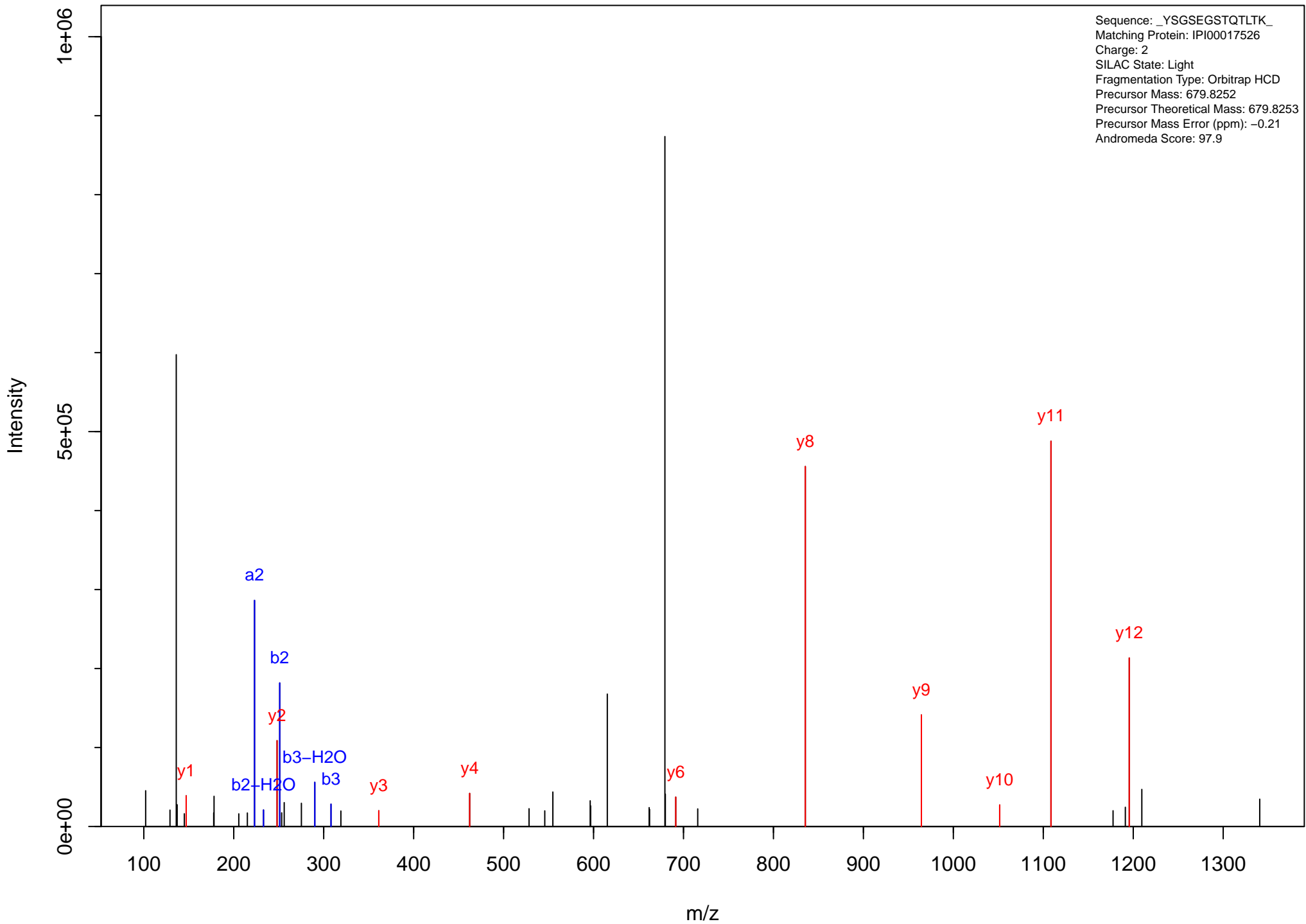


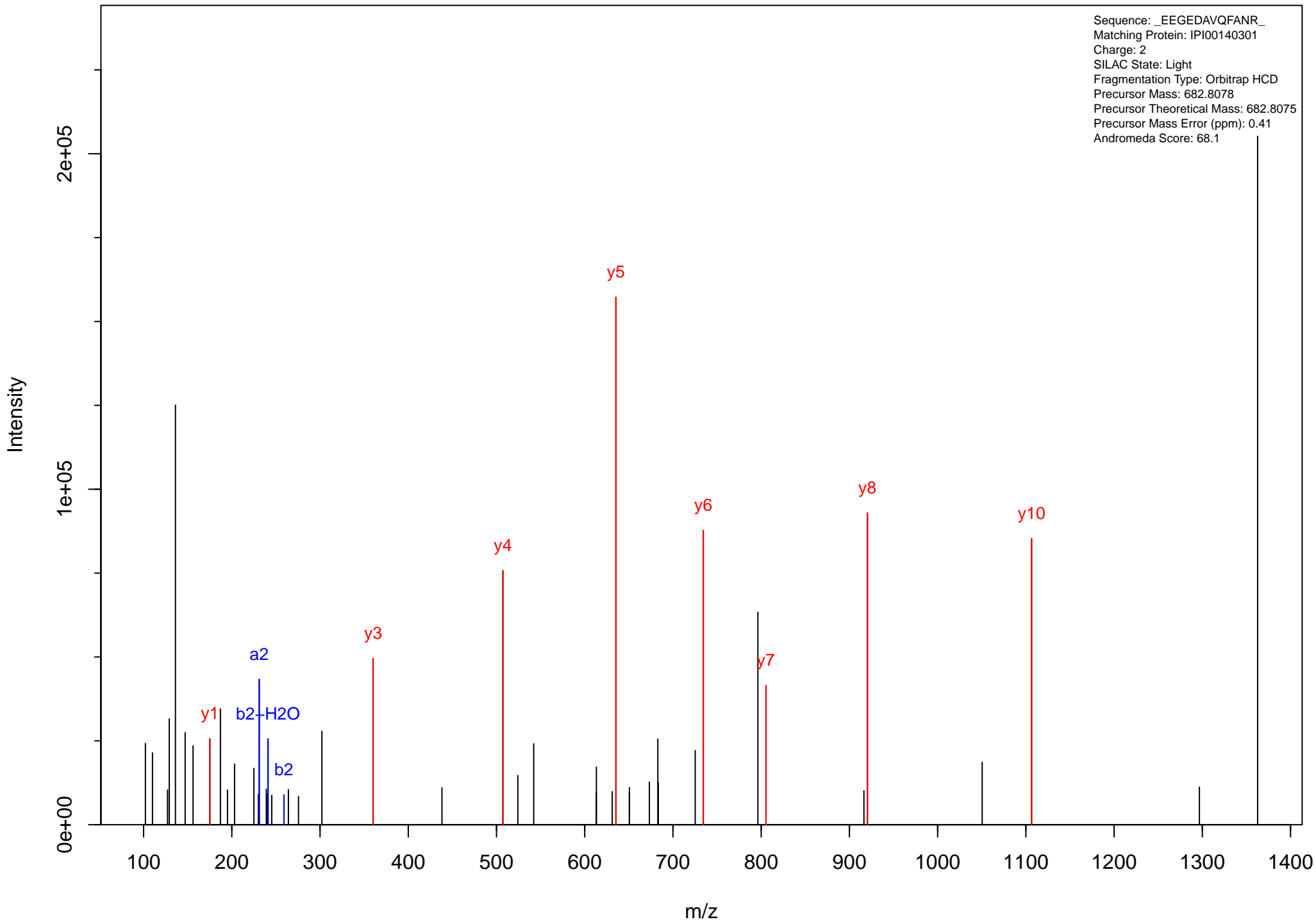


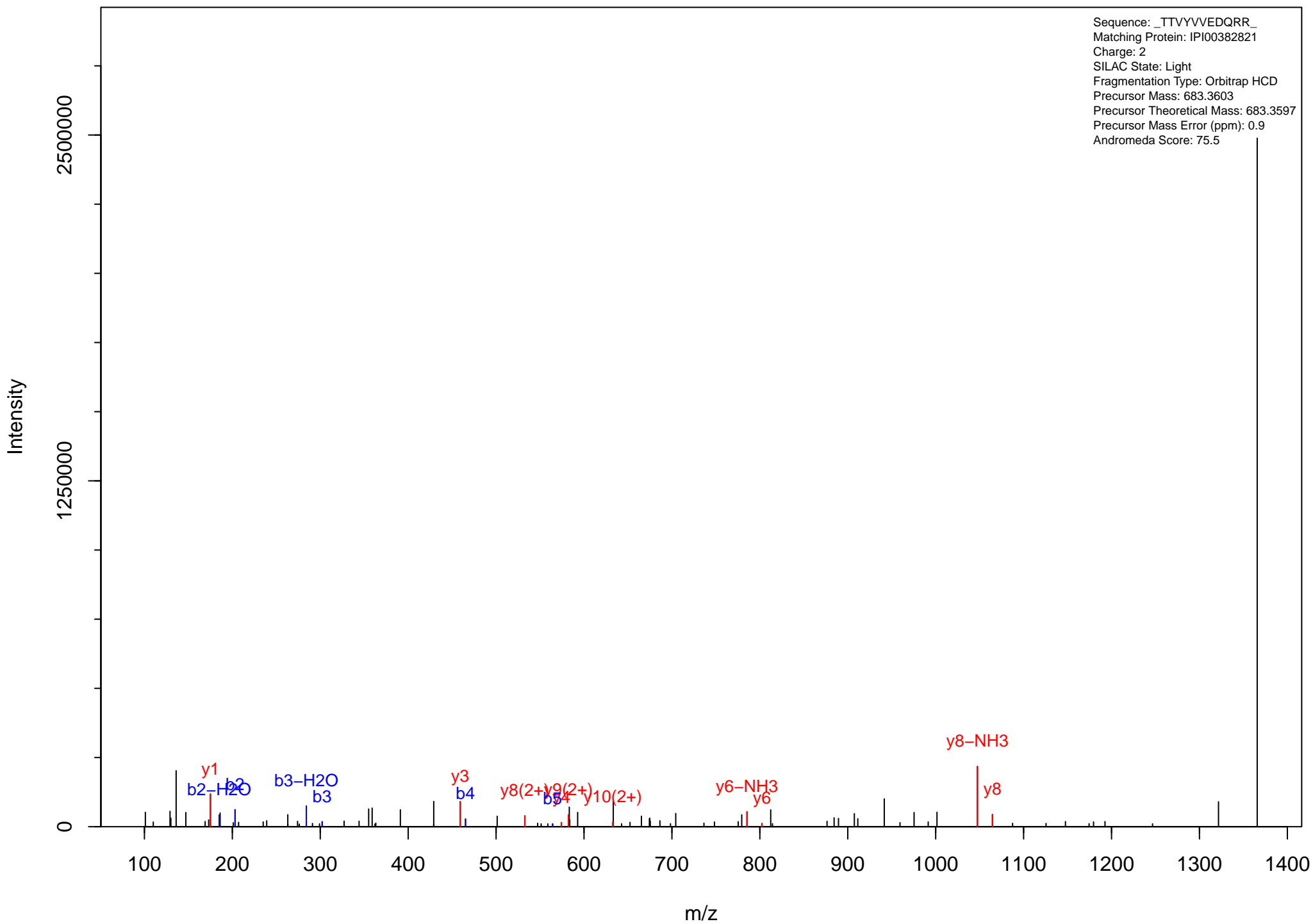


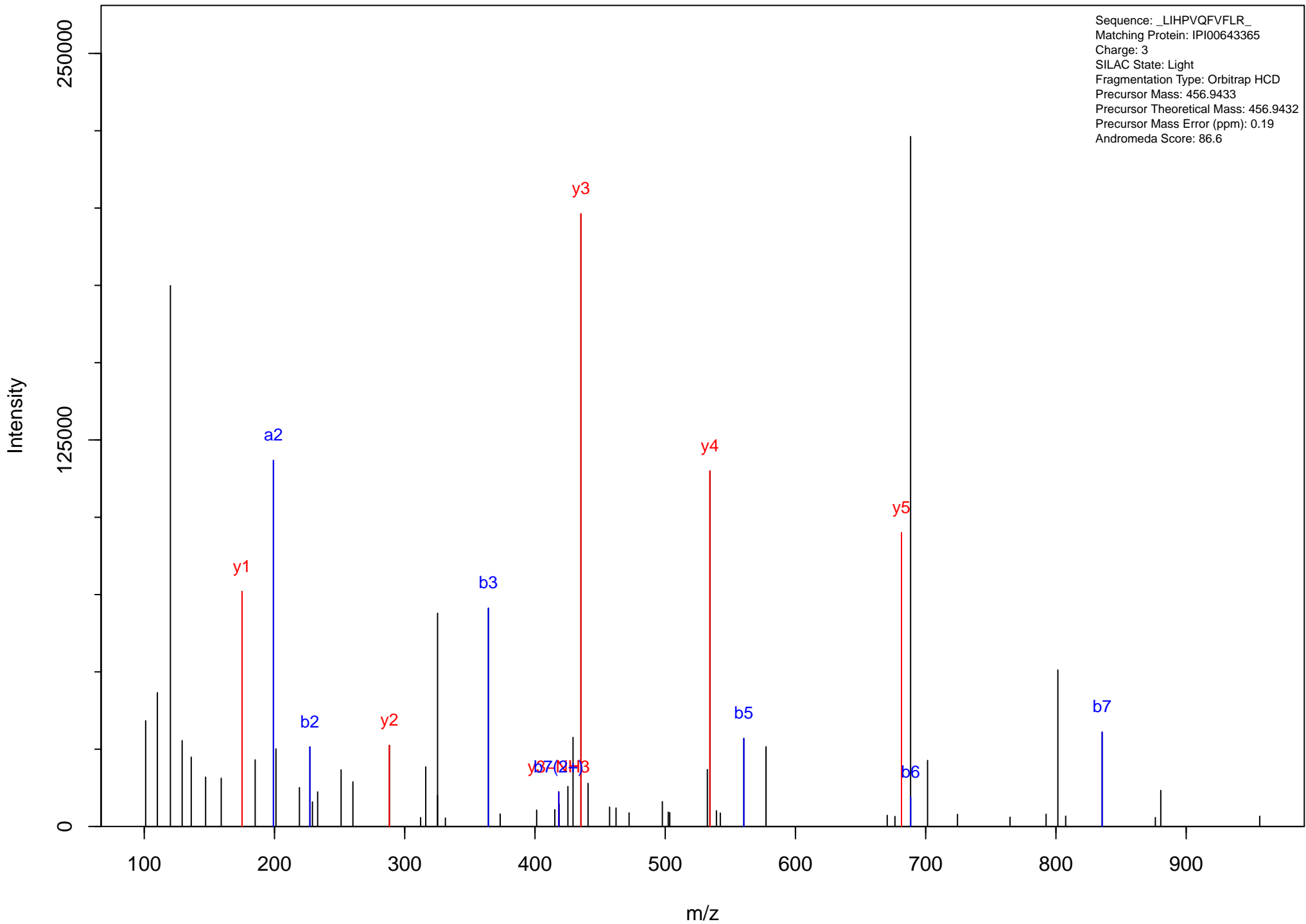


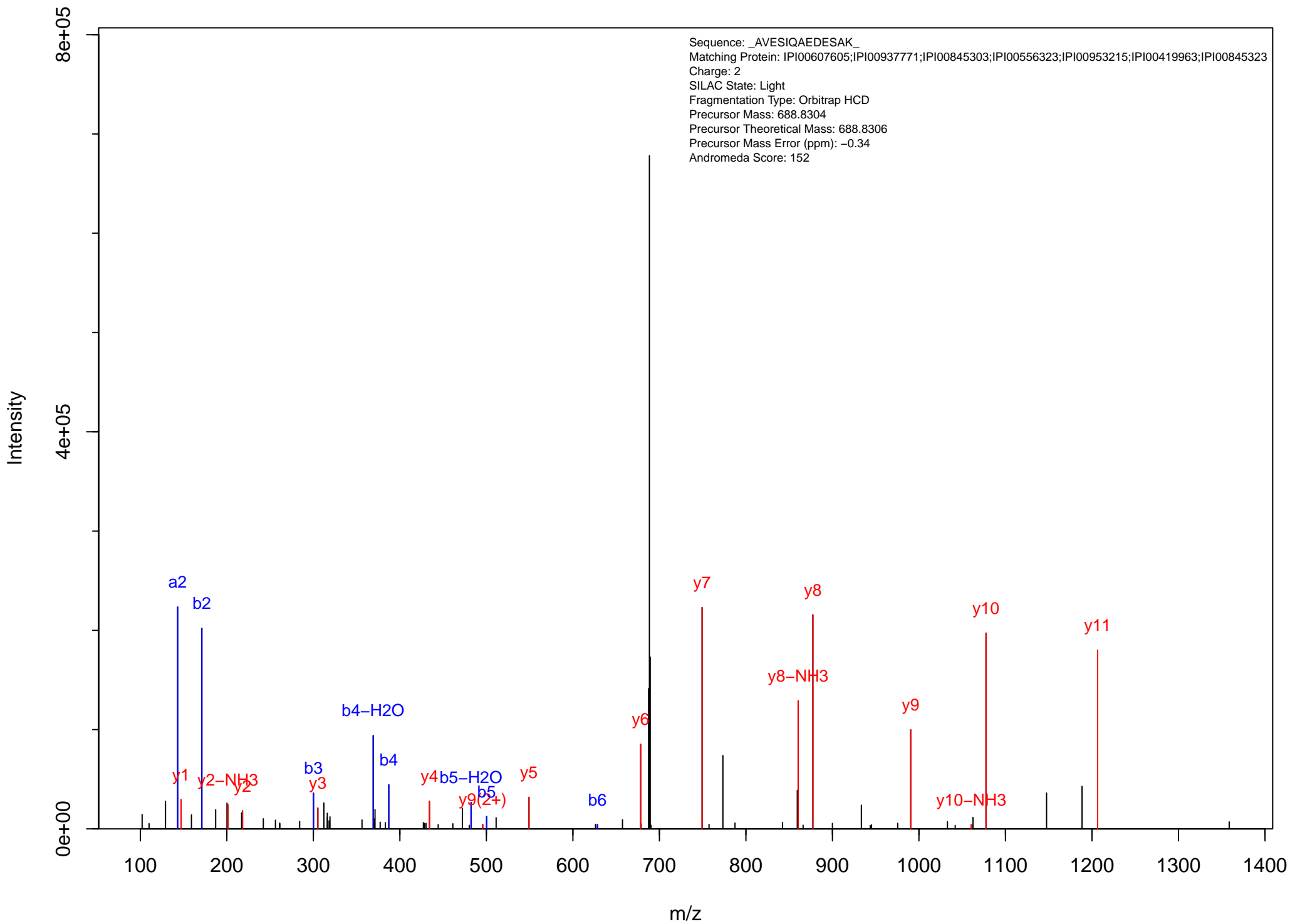


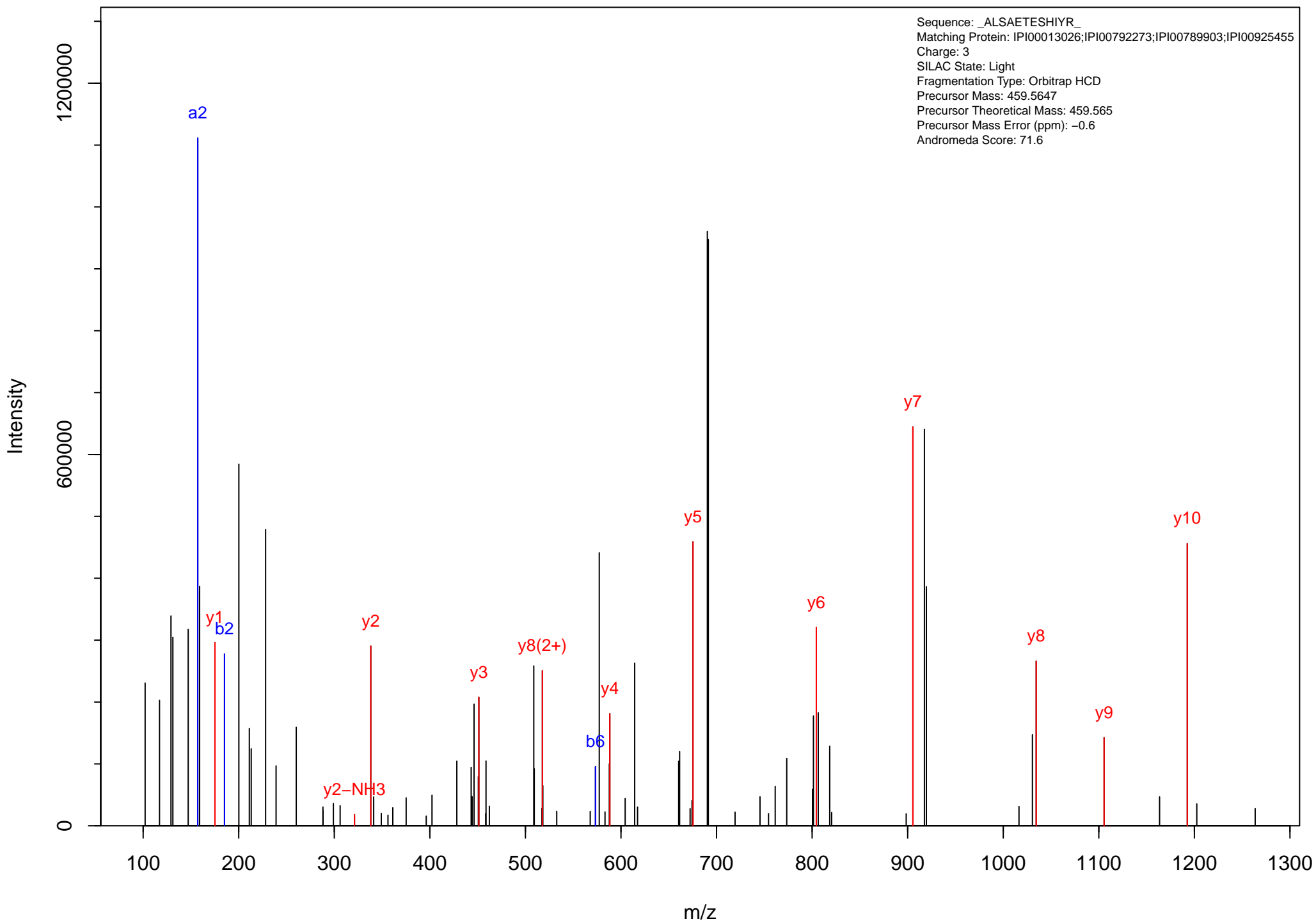


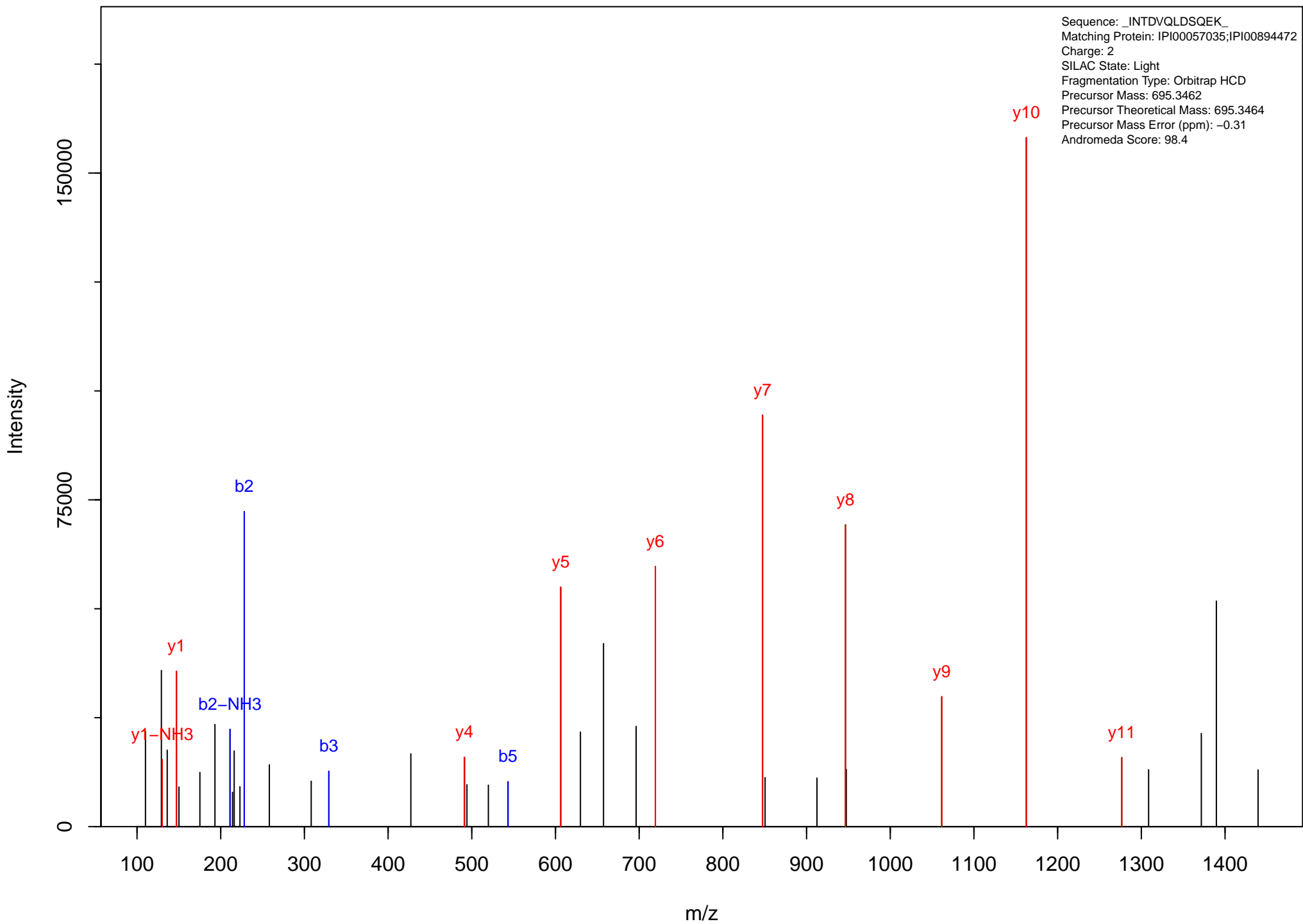


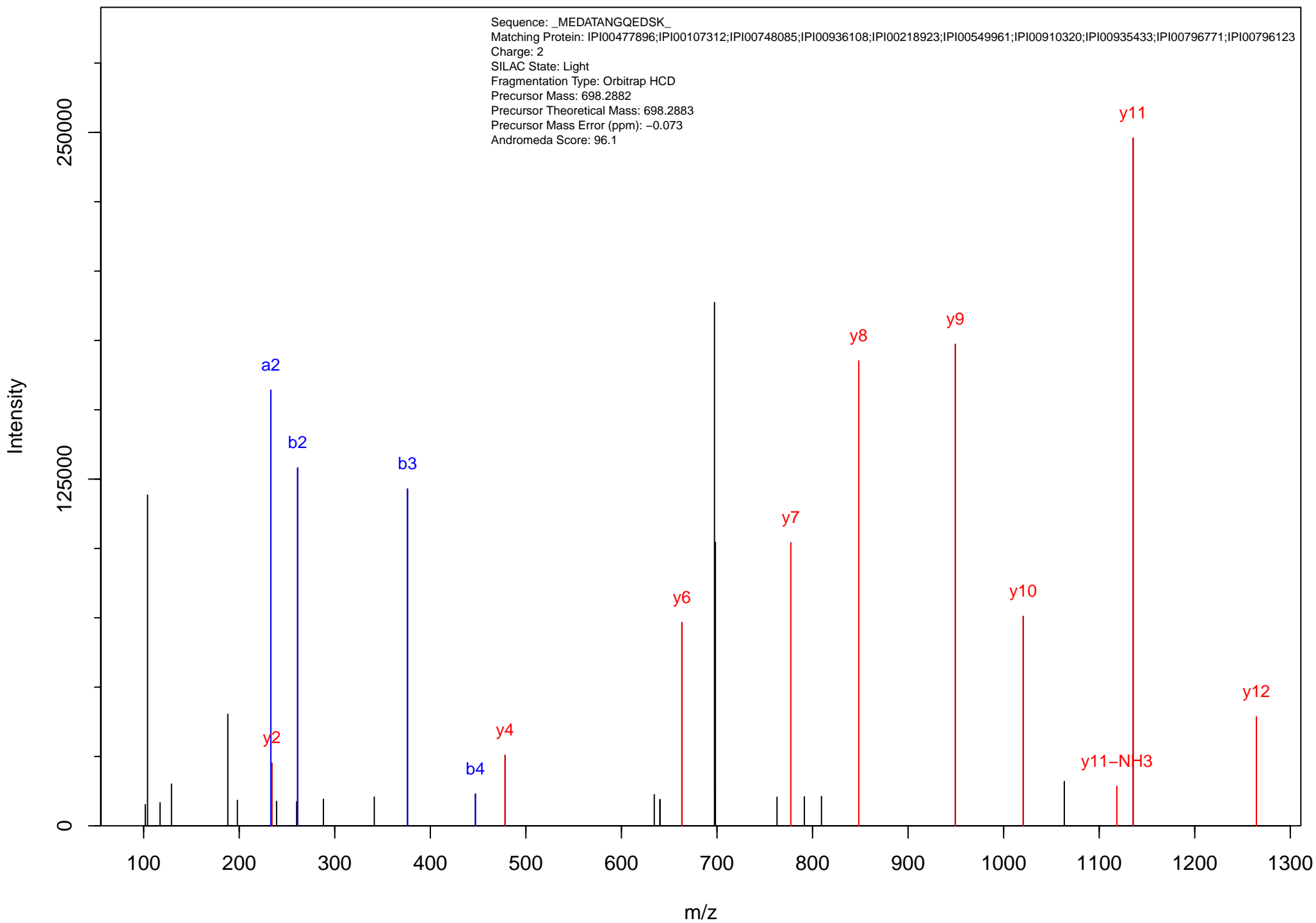


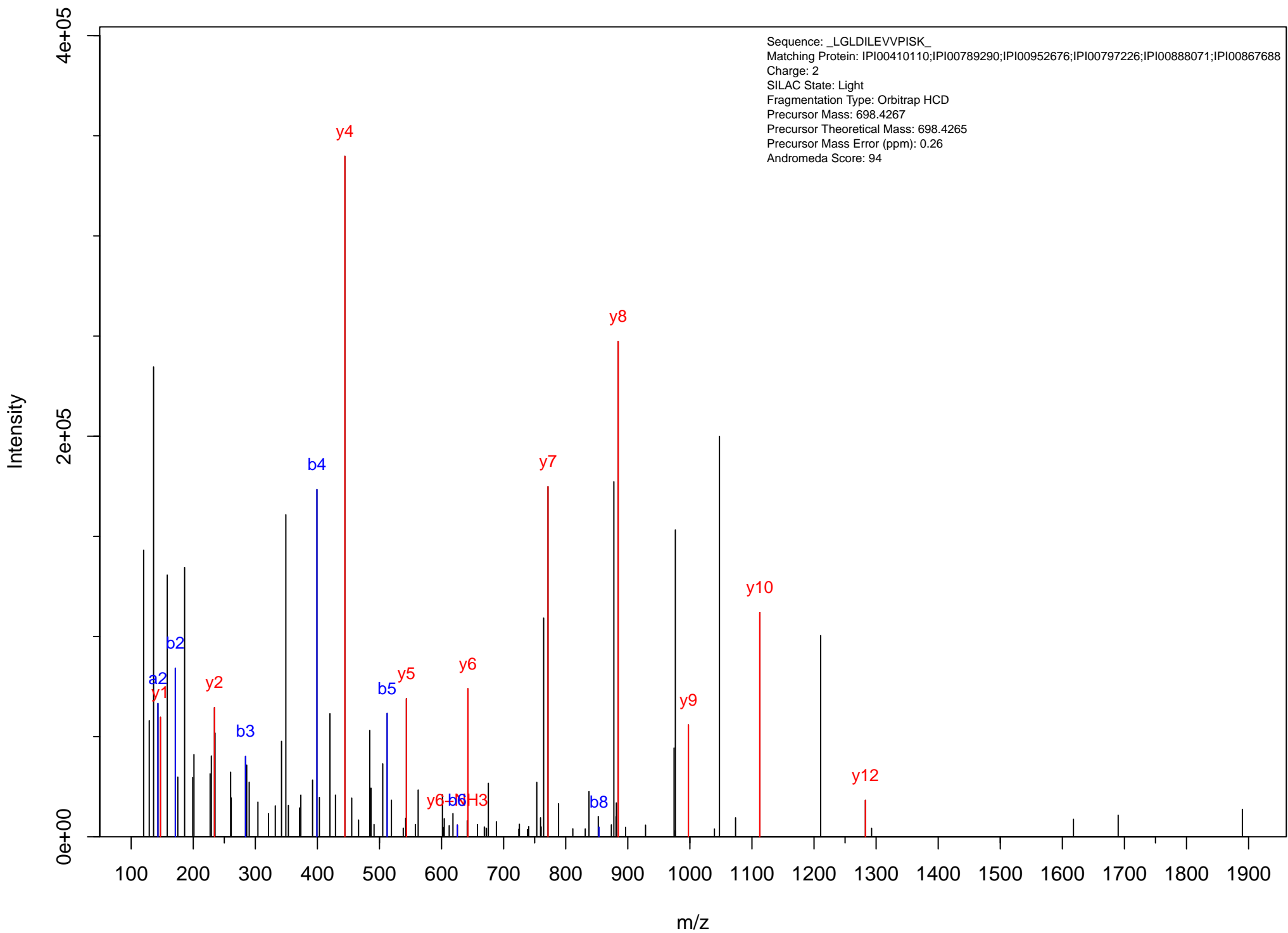


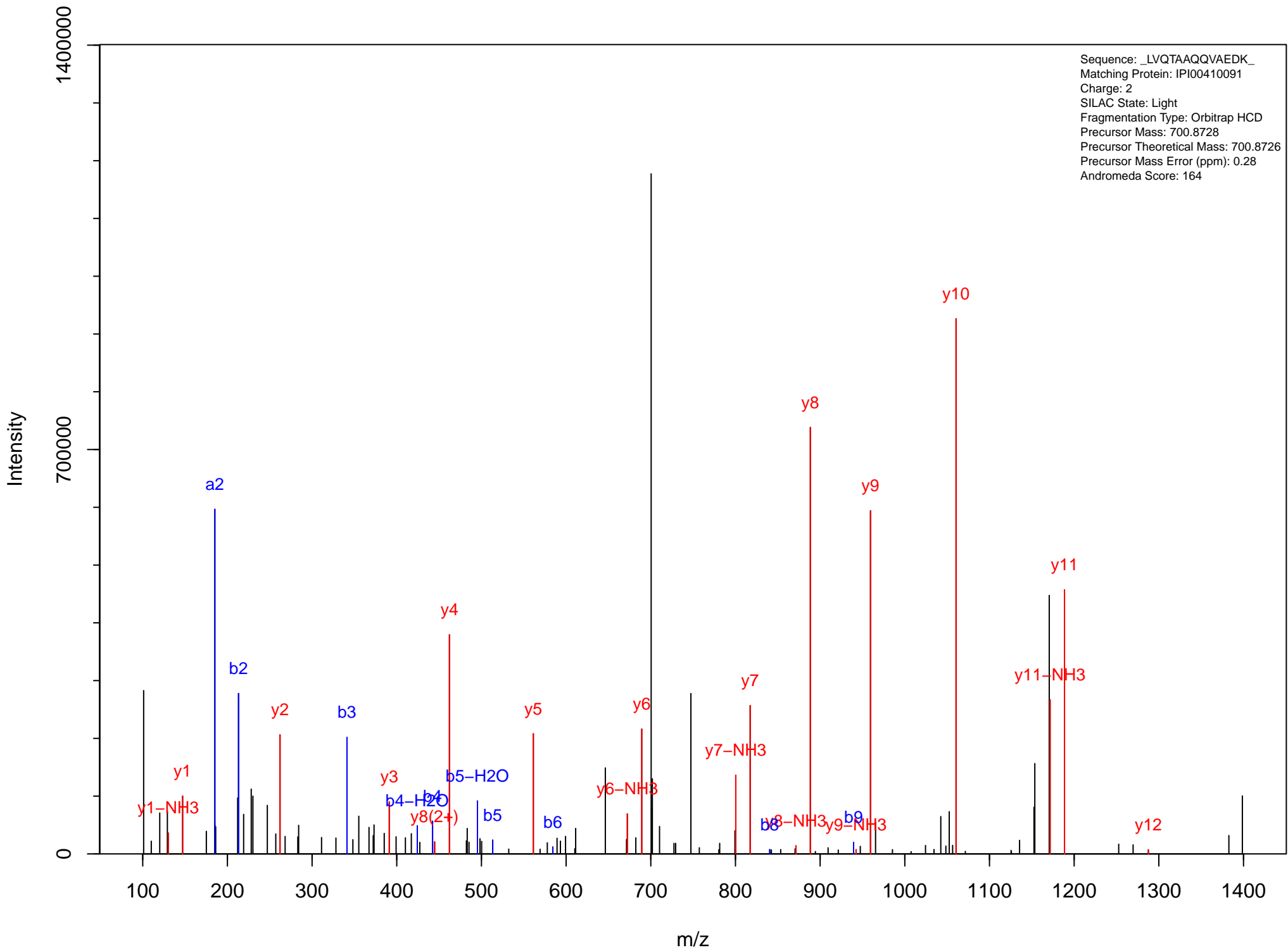


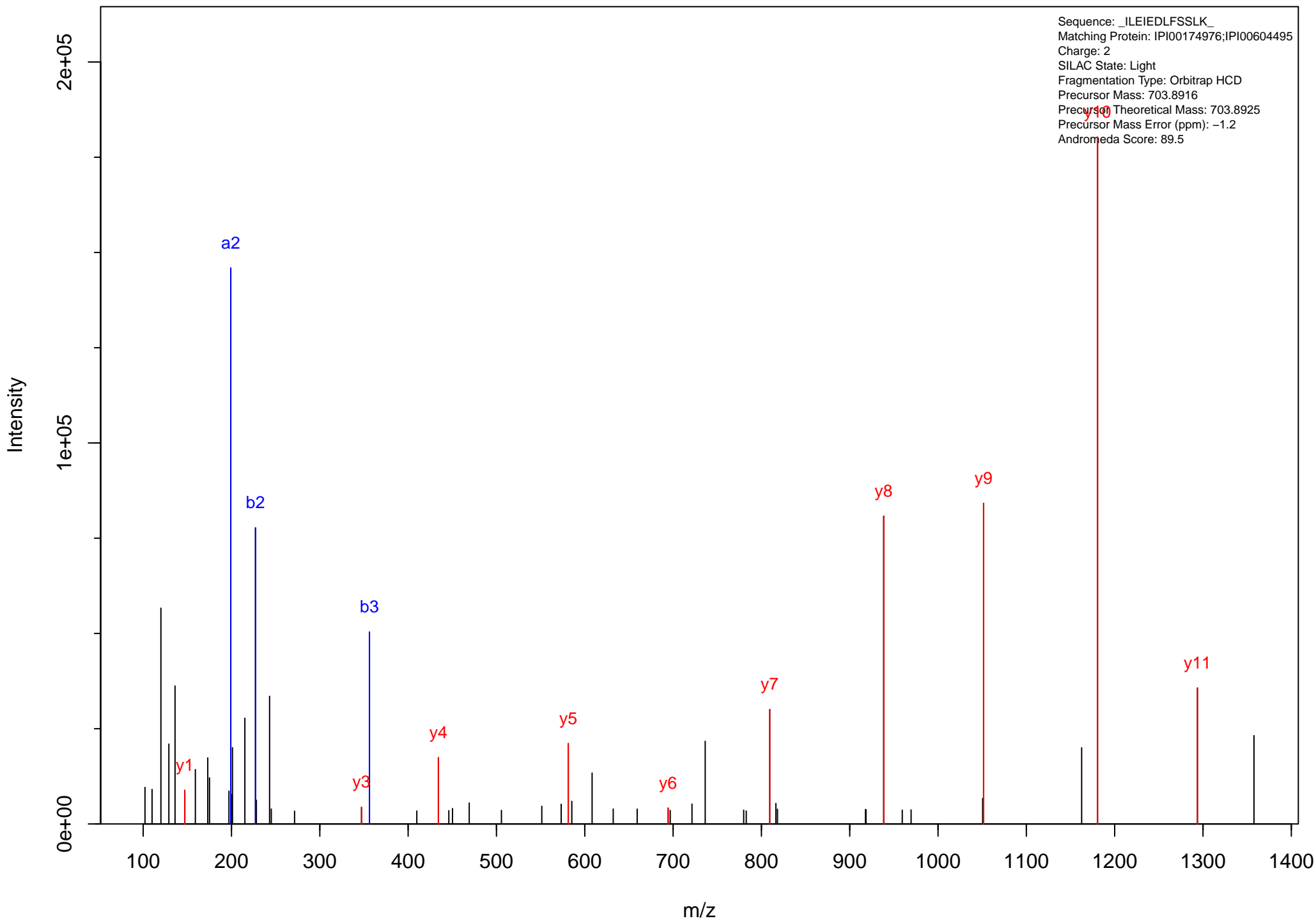


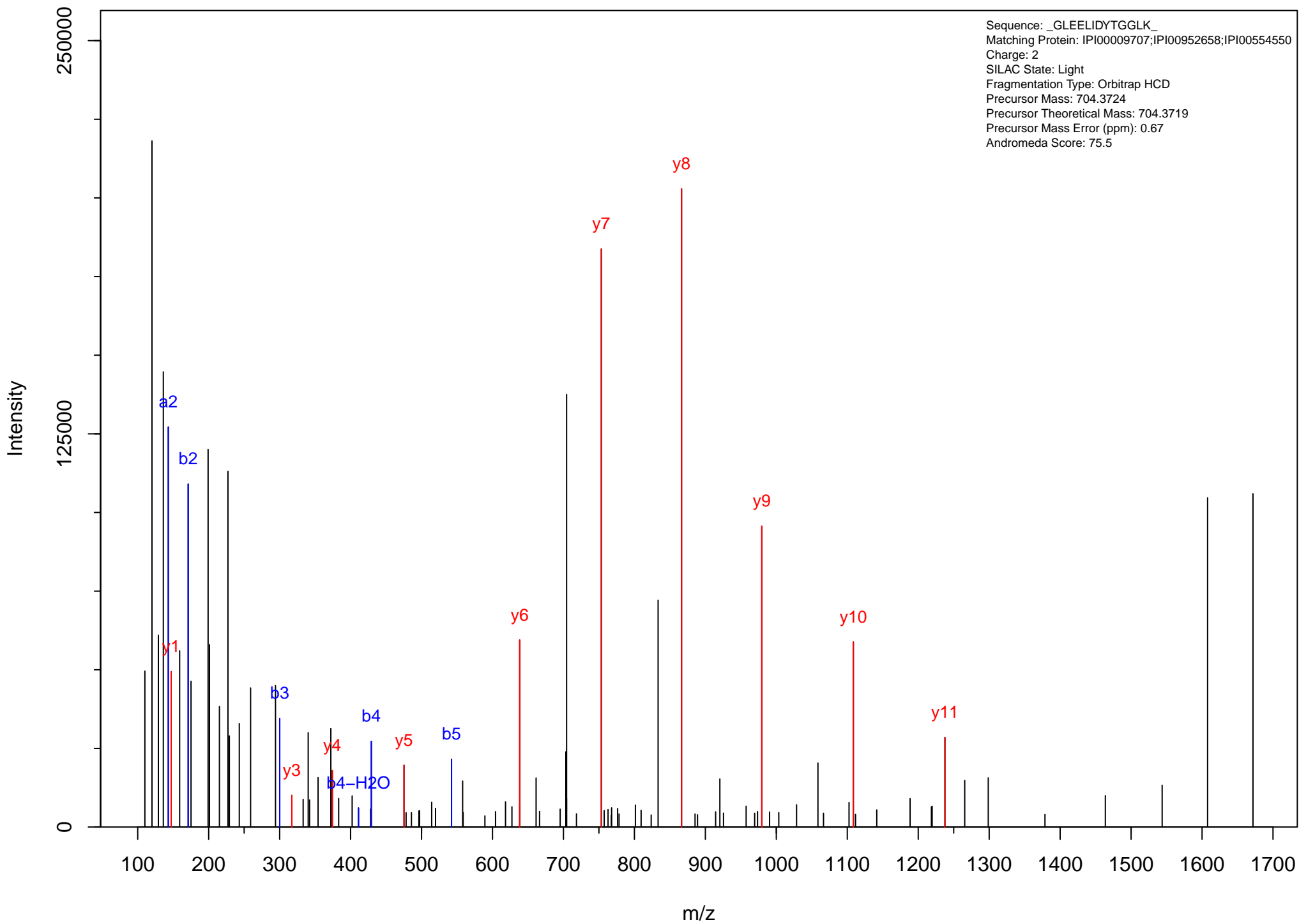


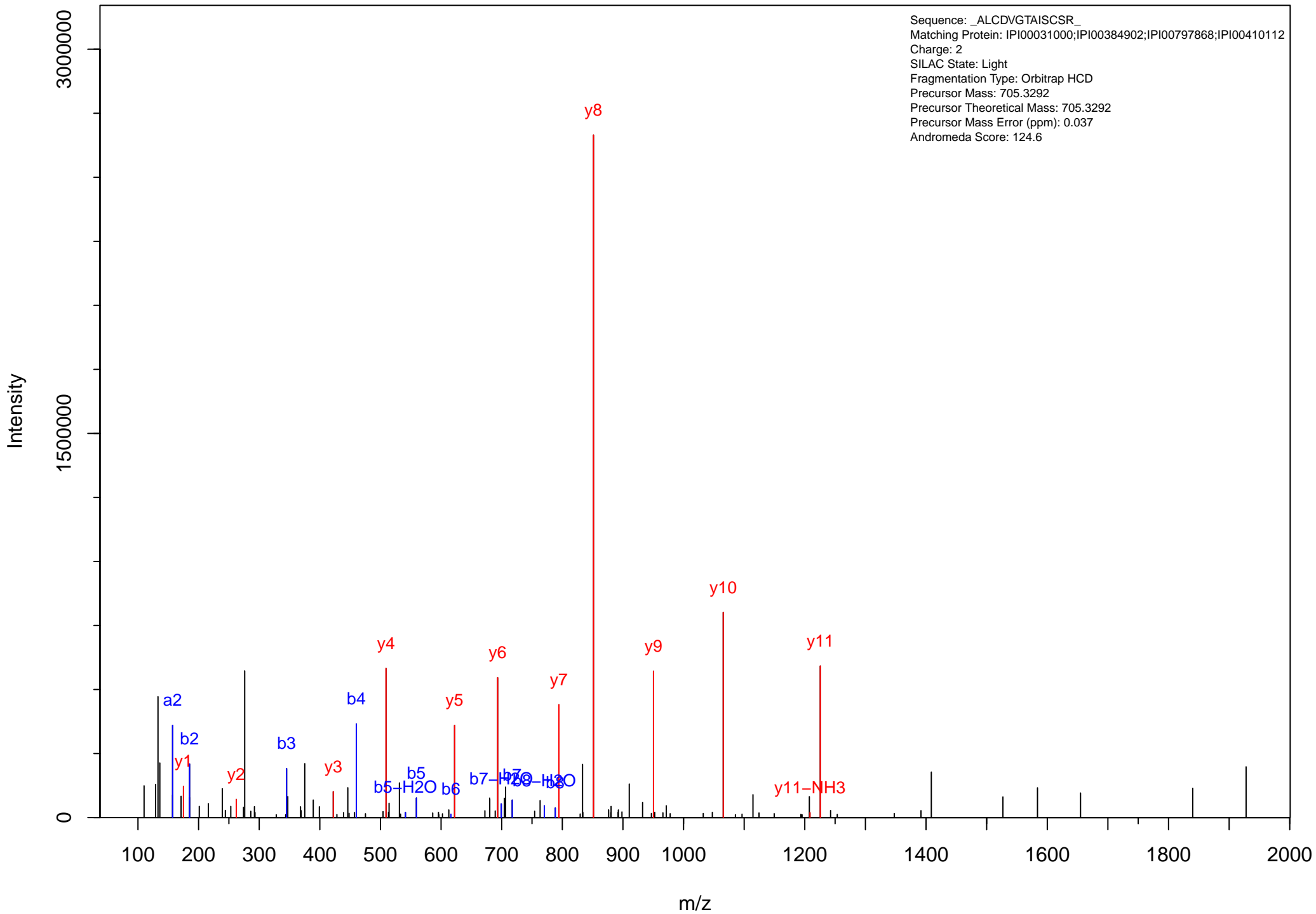




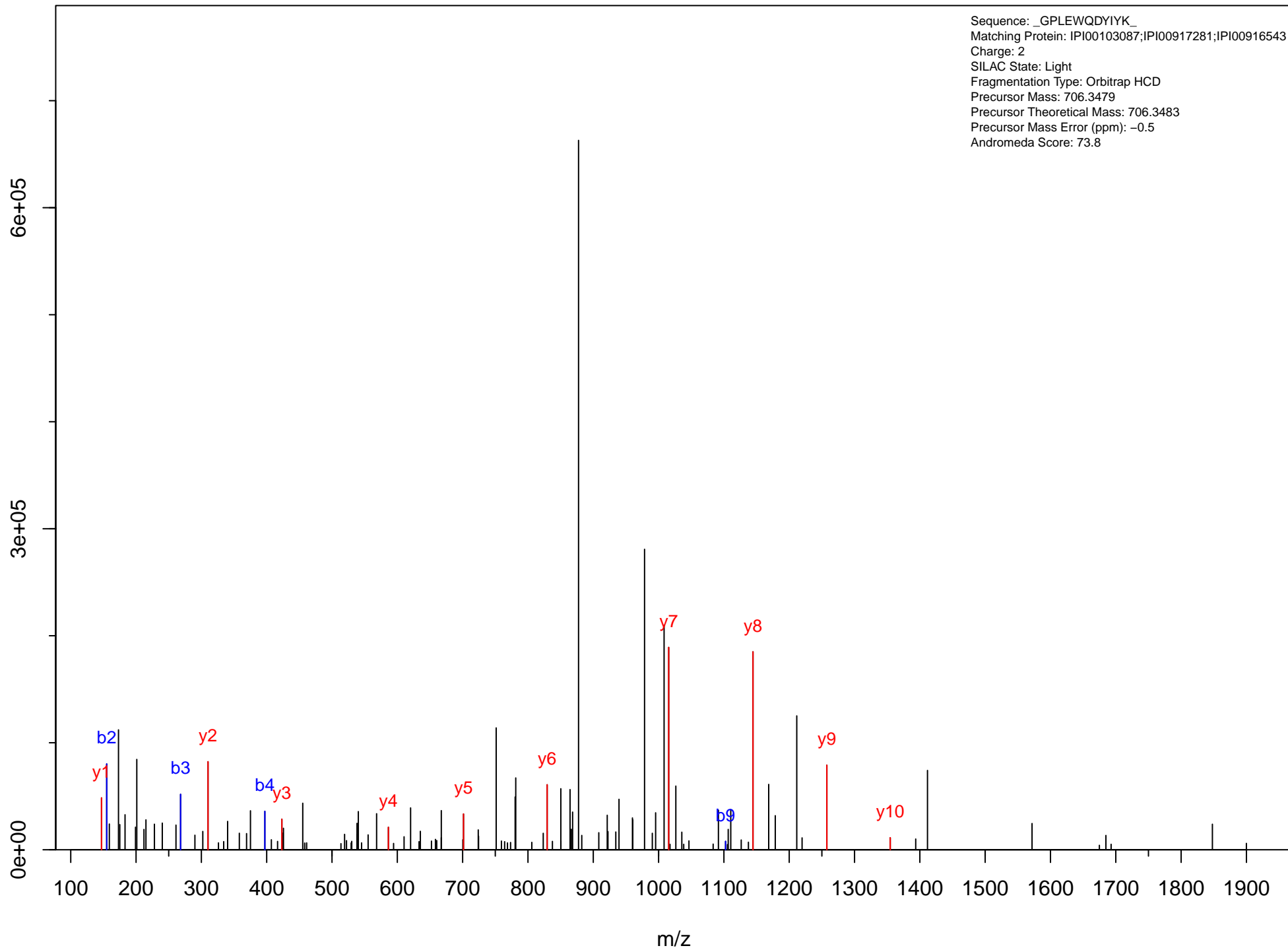


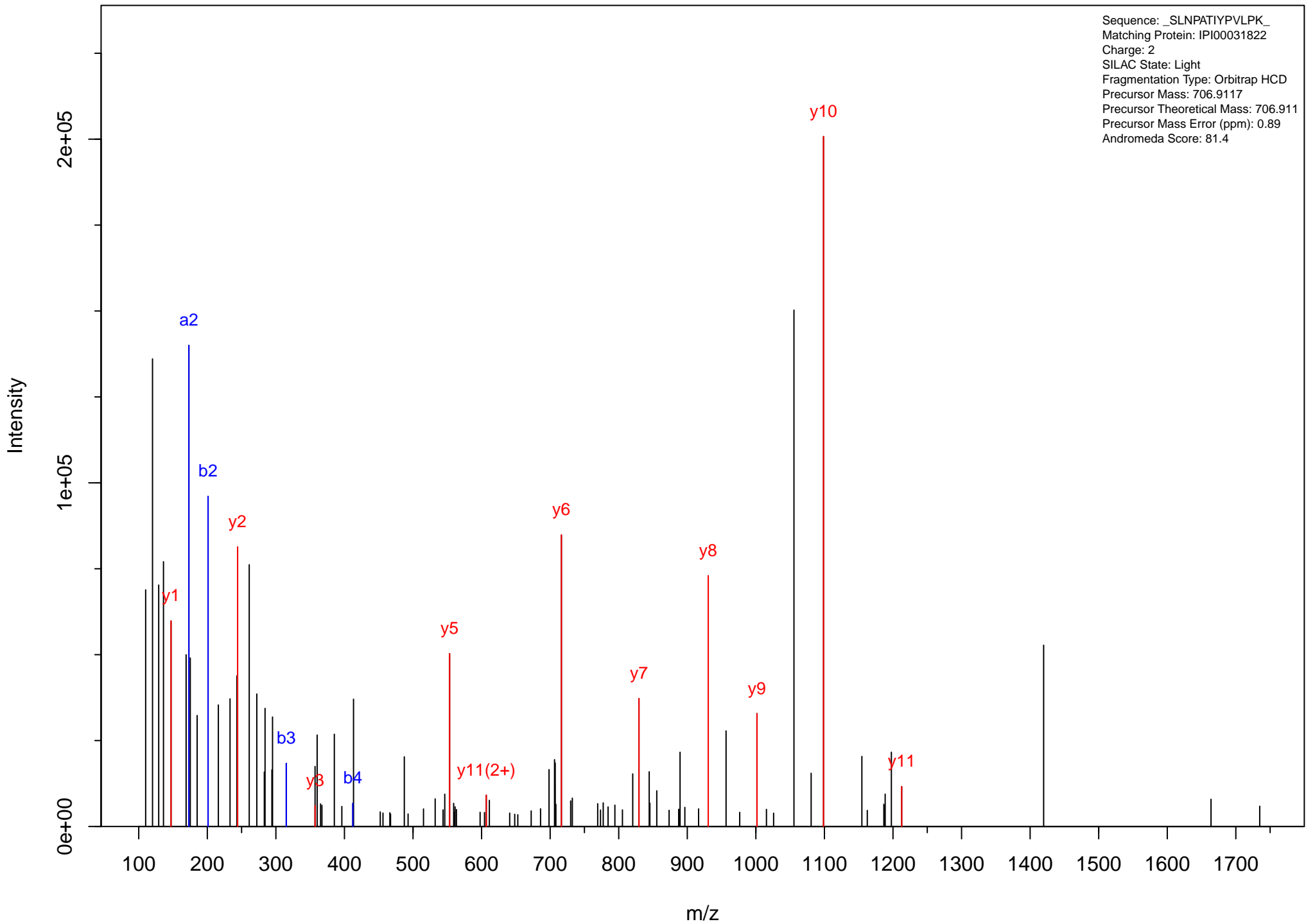


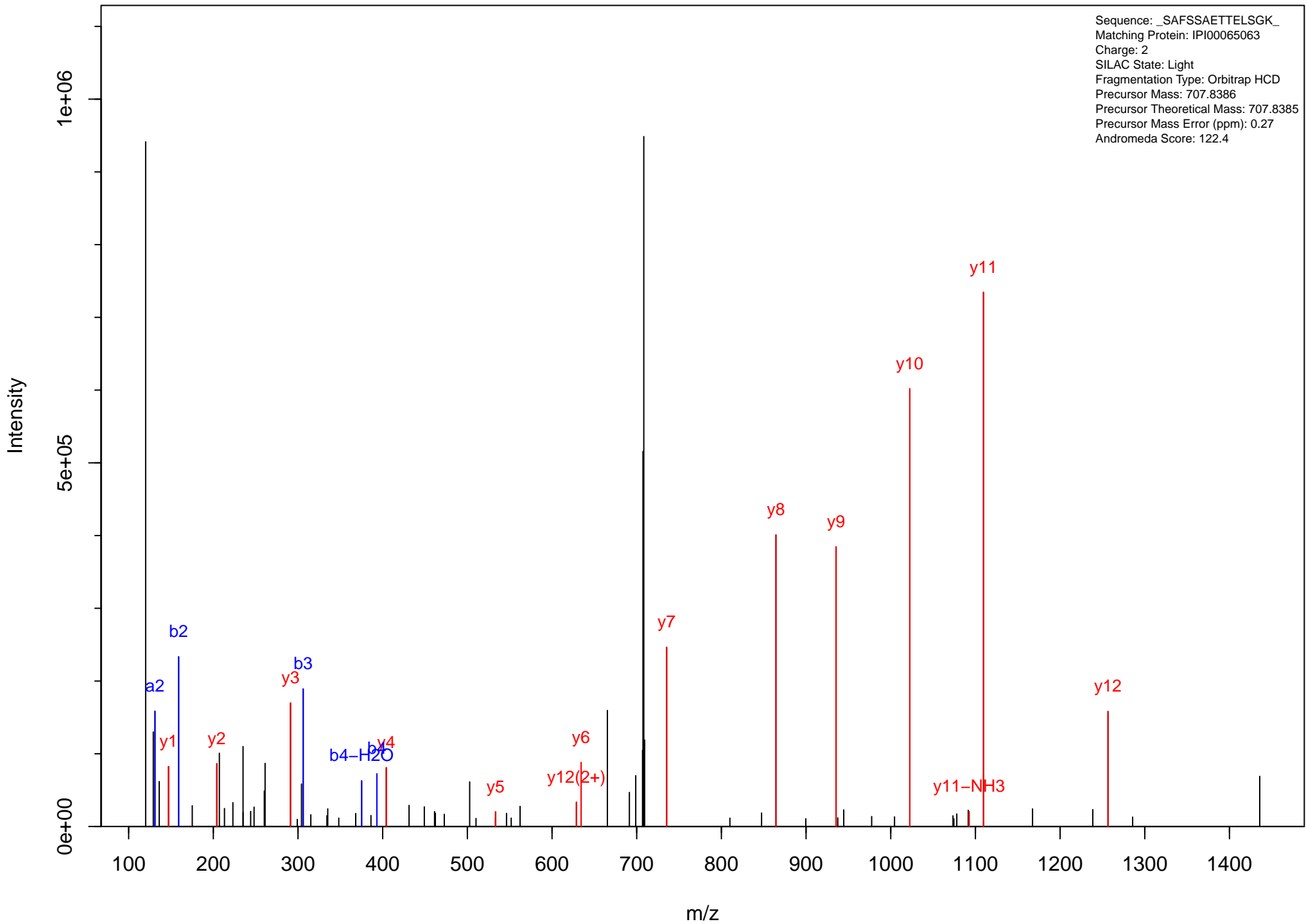


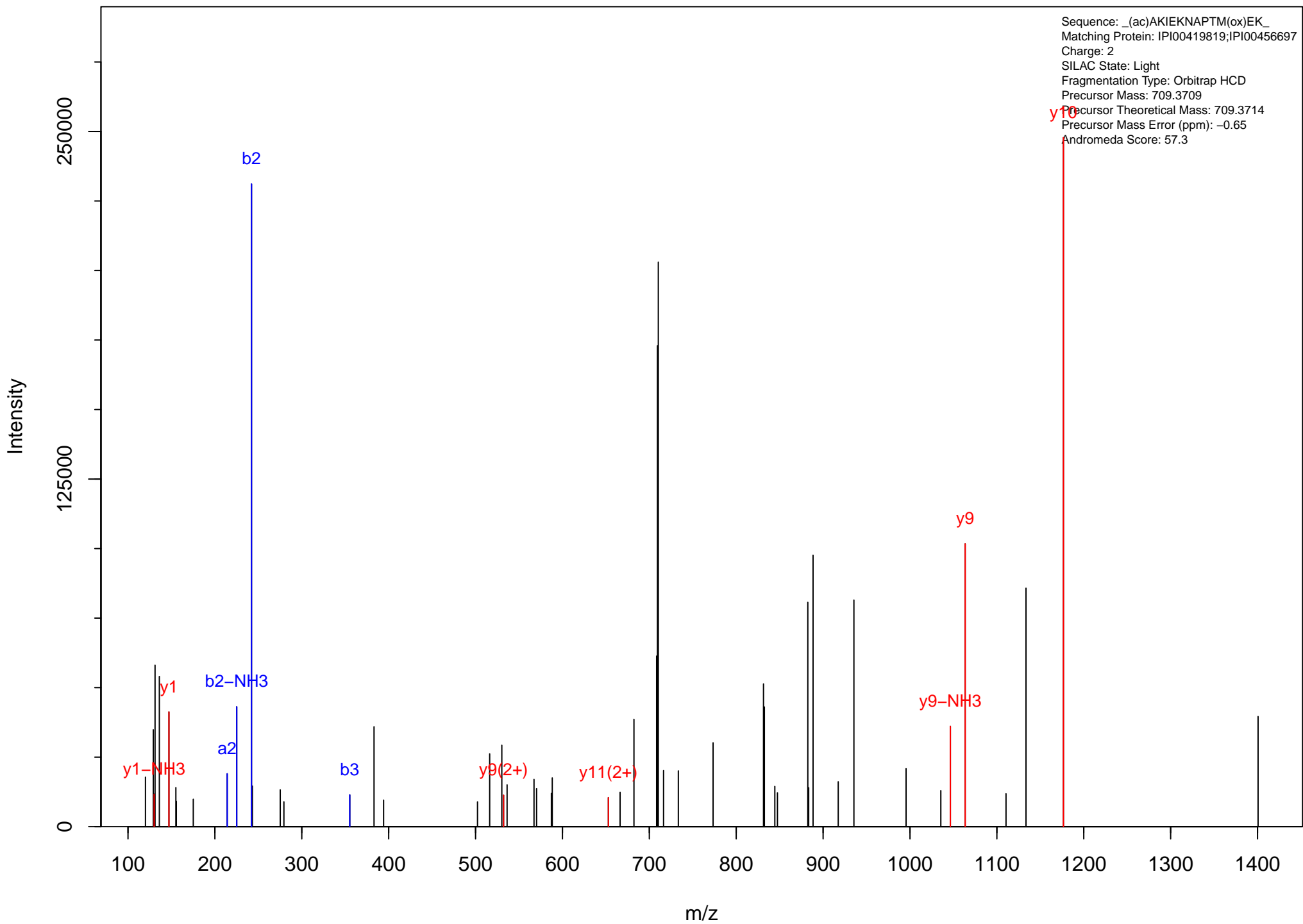


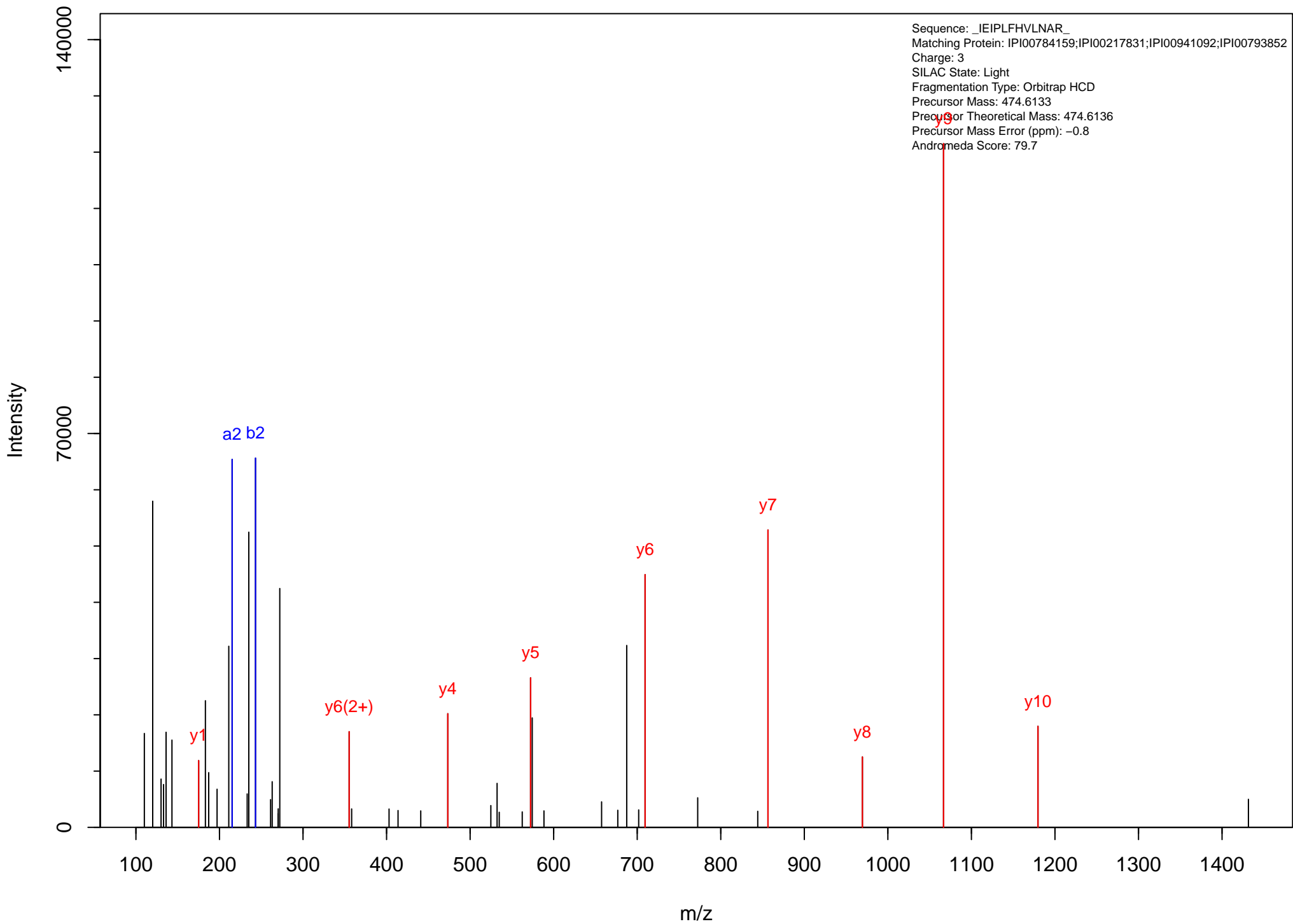
Intensity

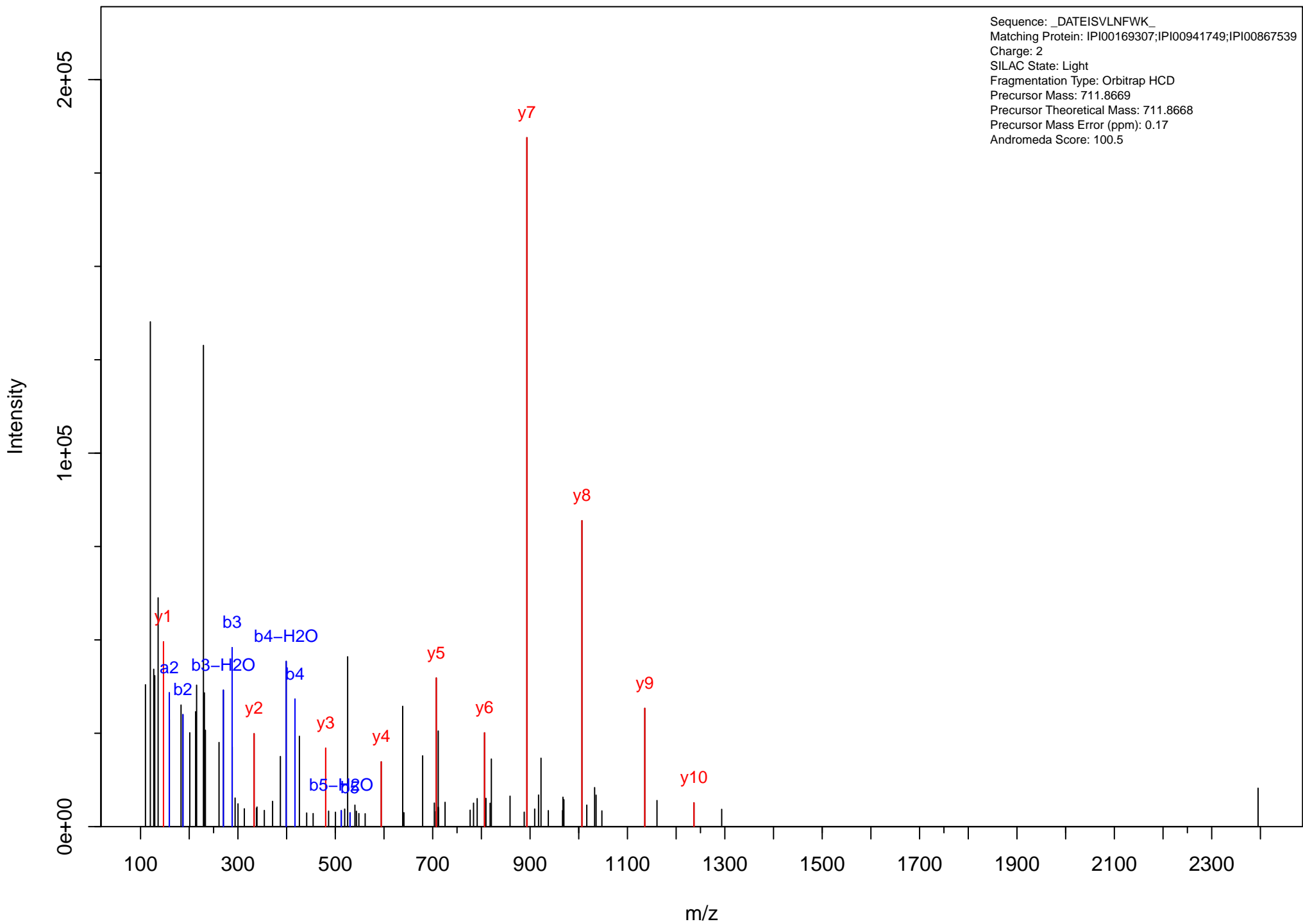


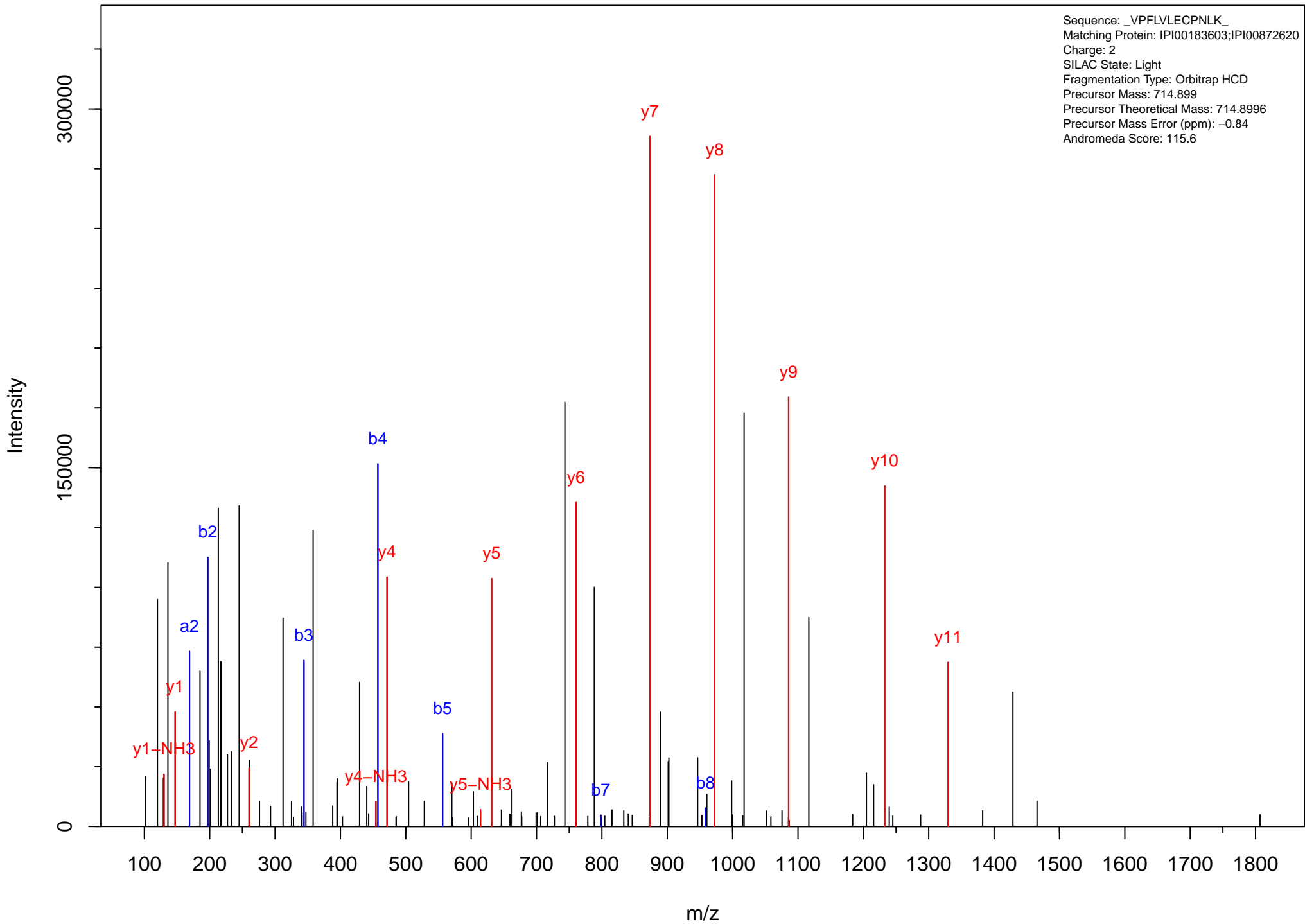


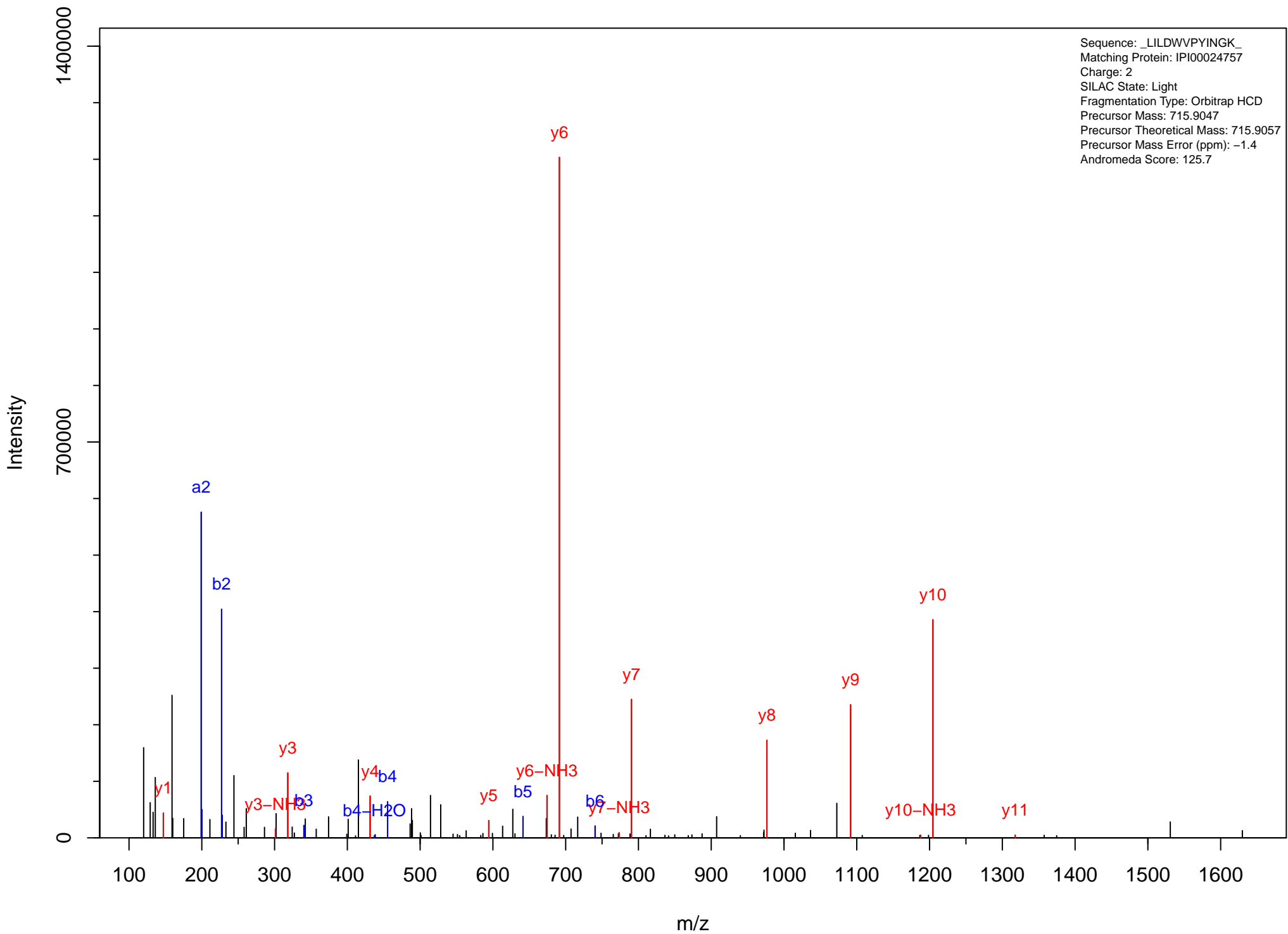


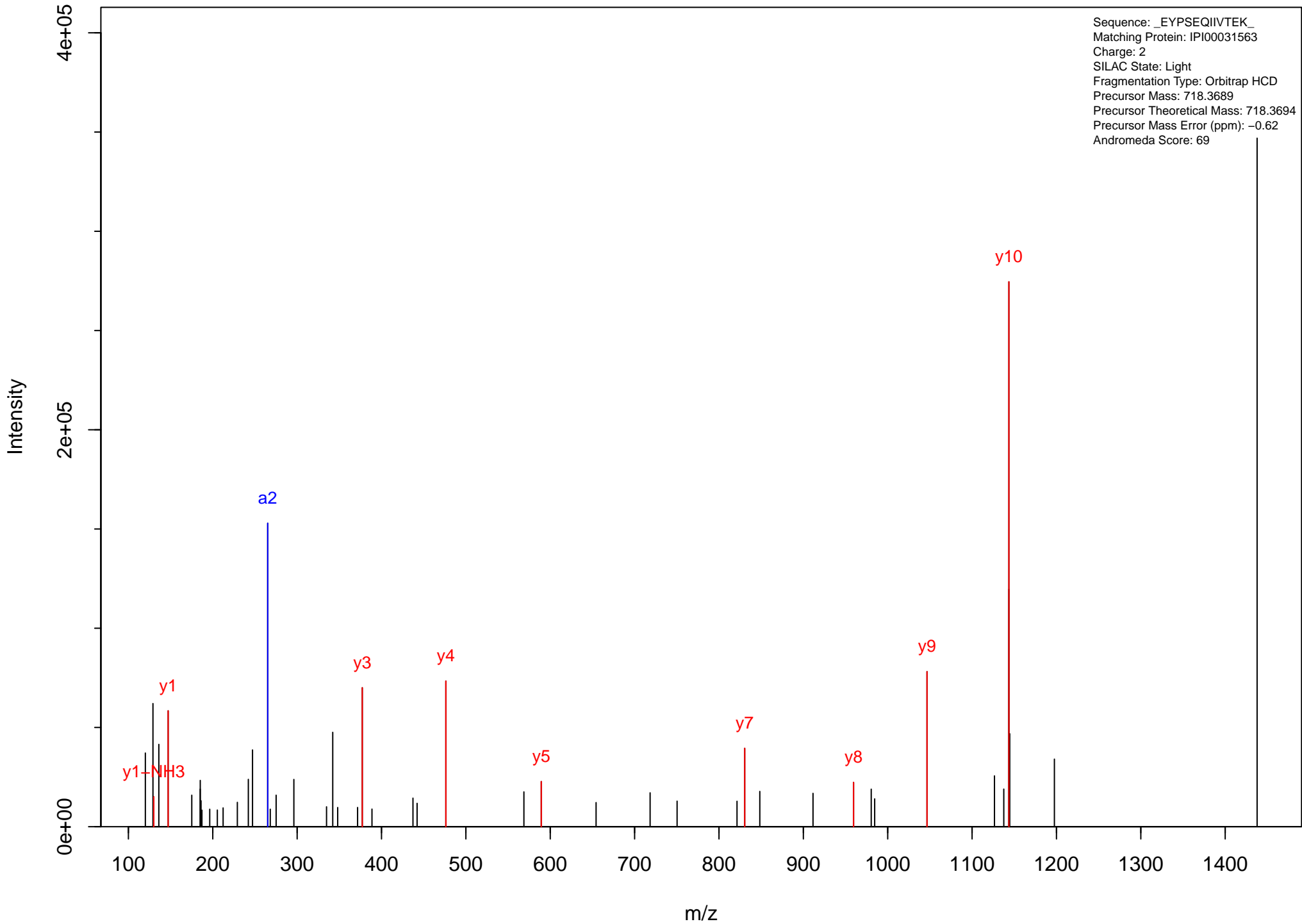


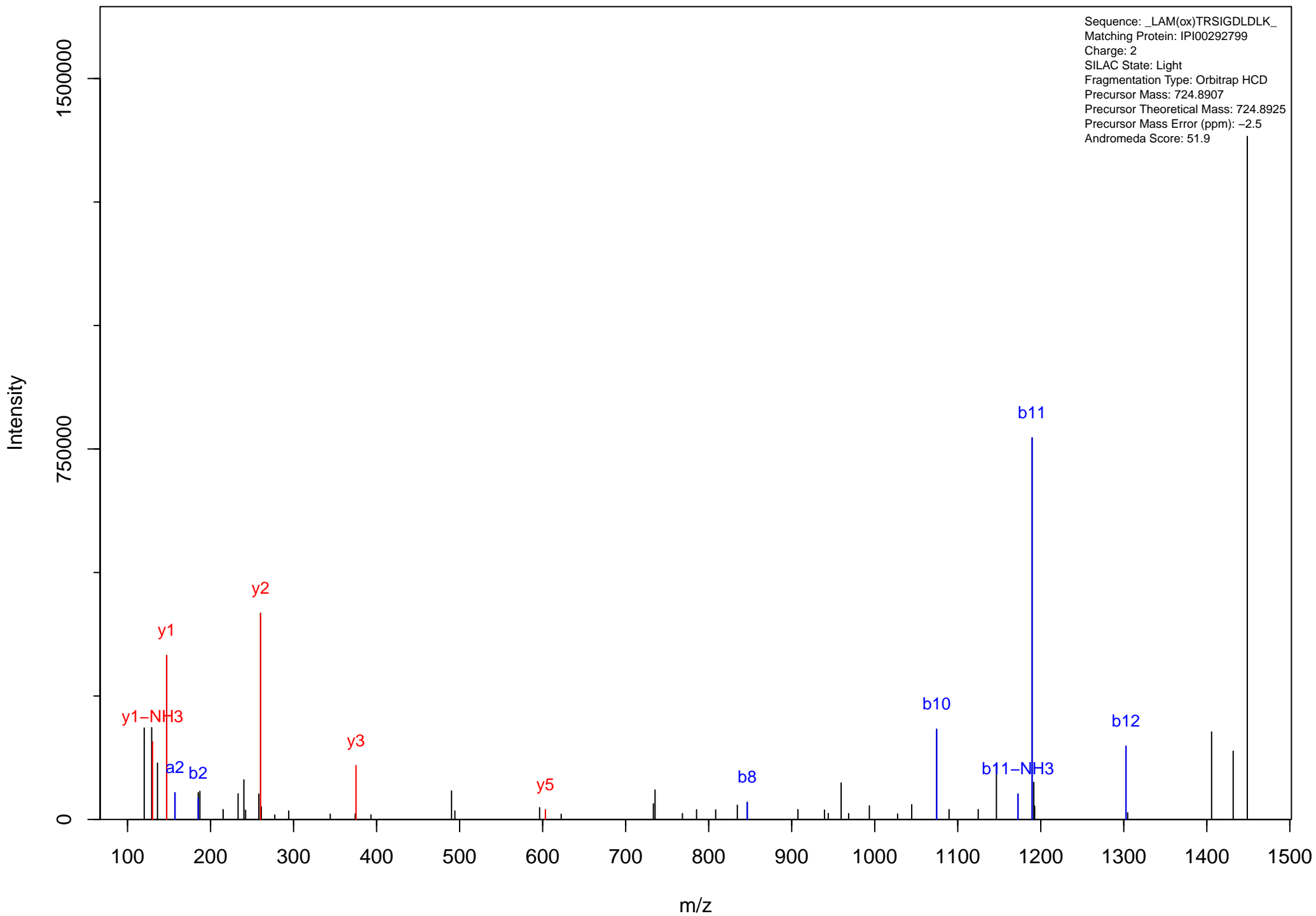


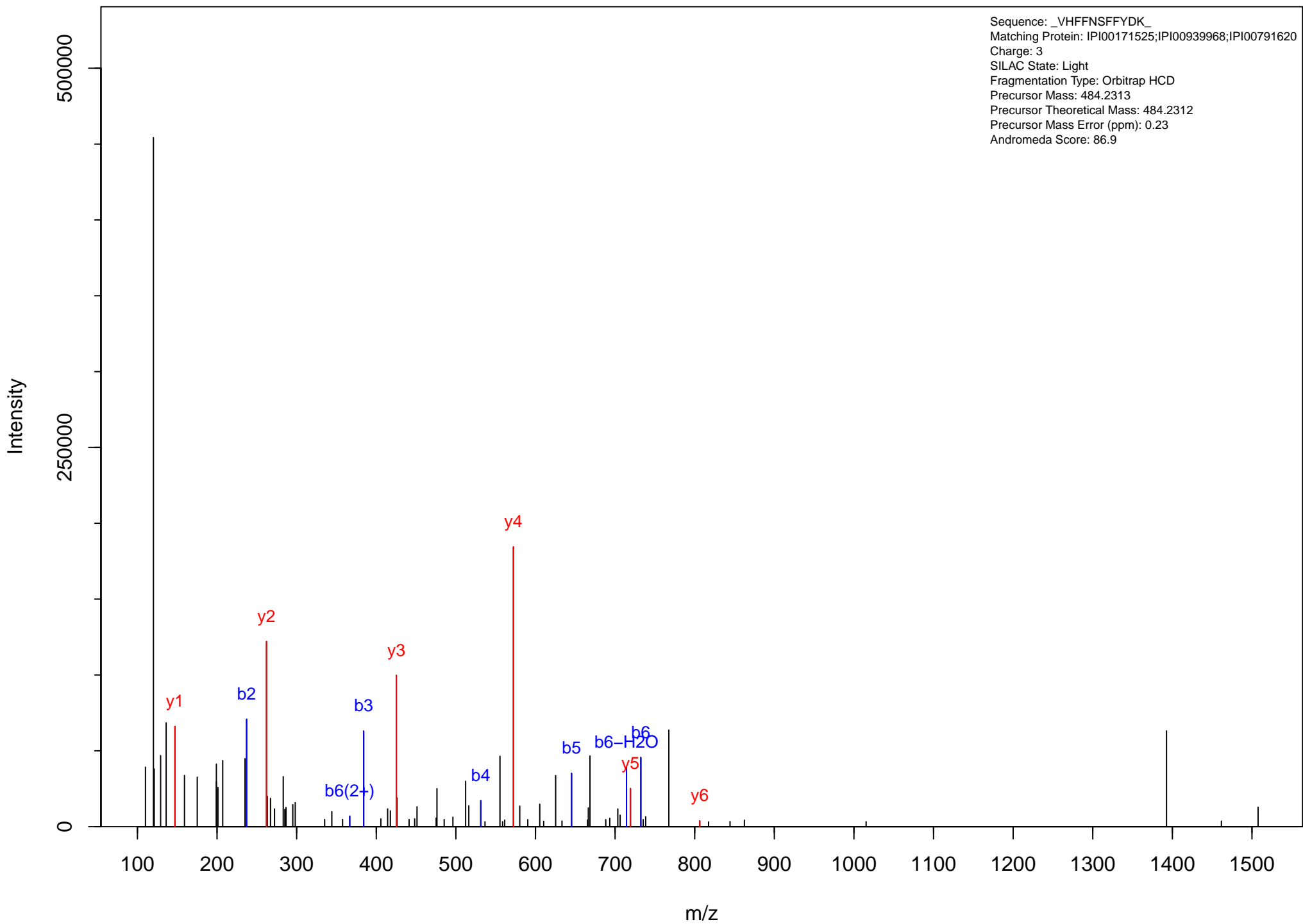


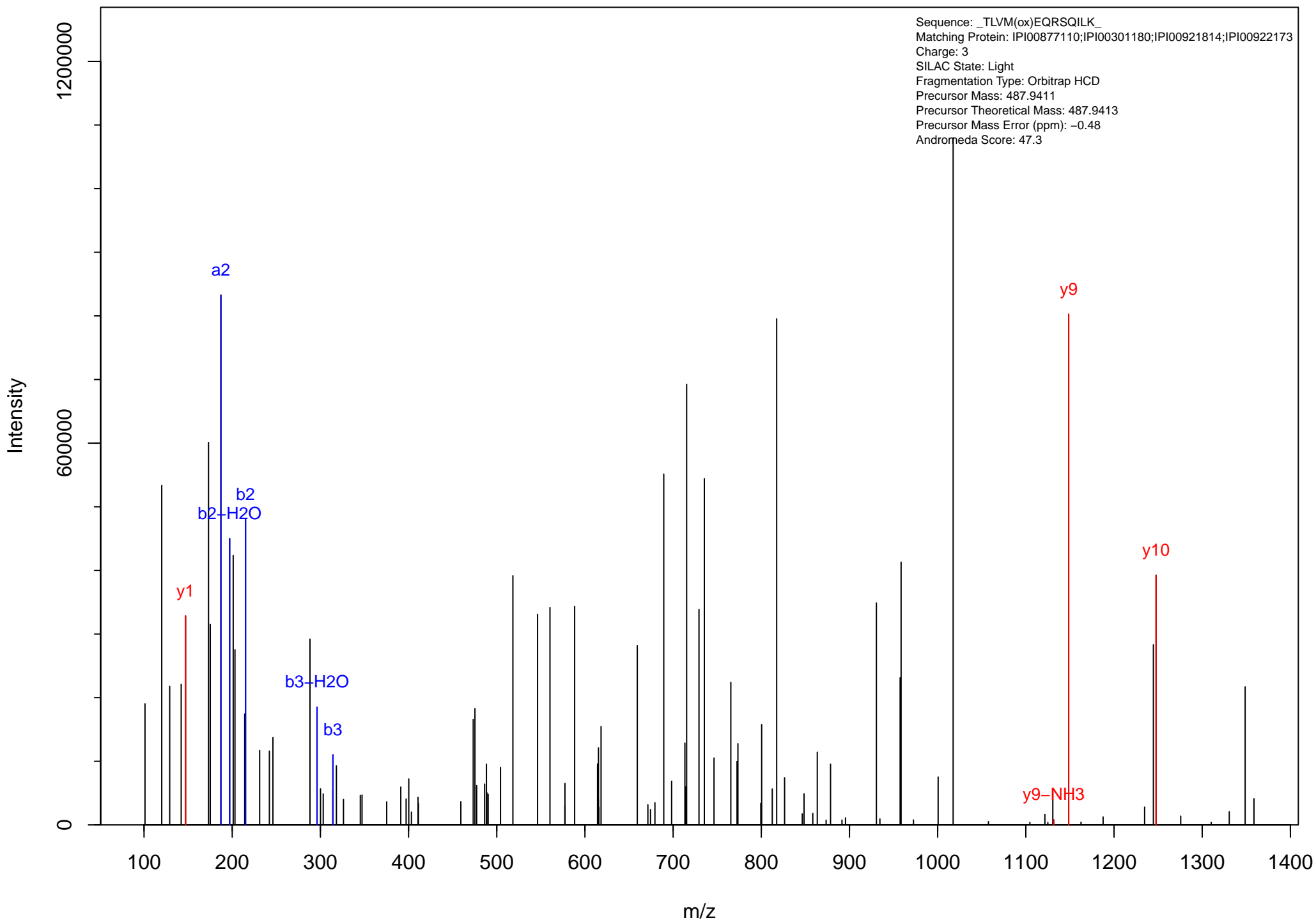


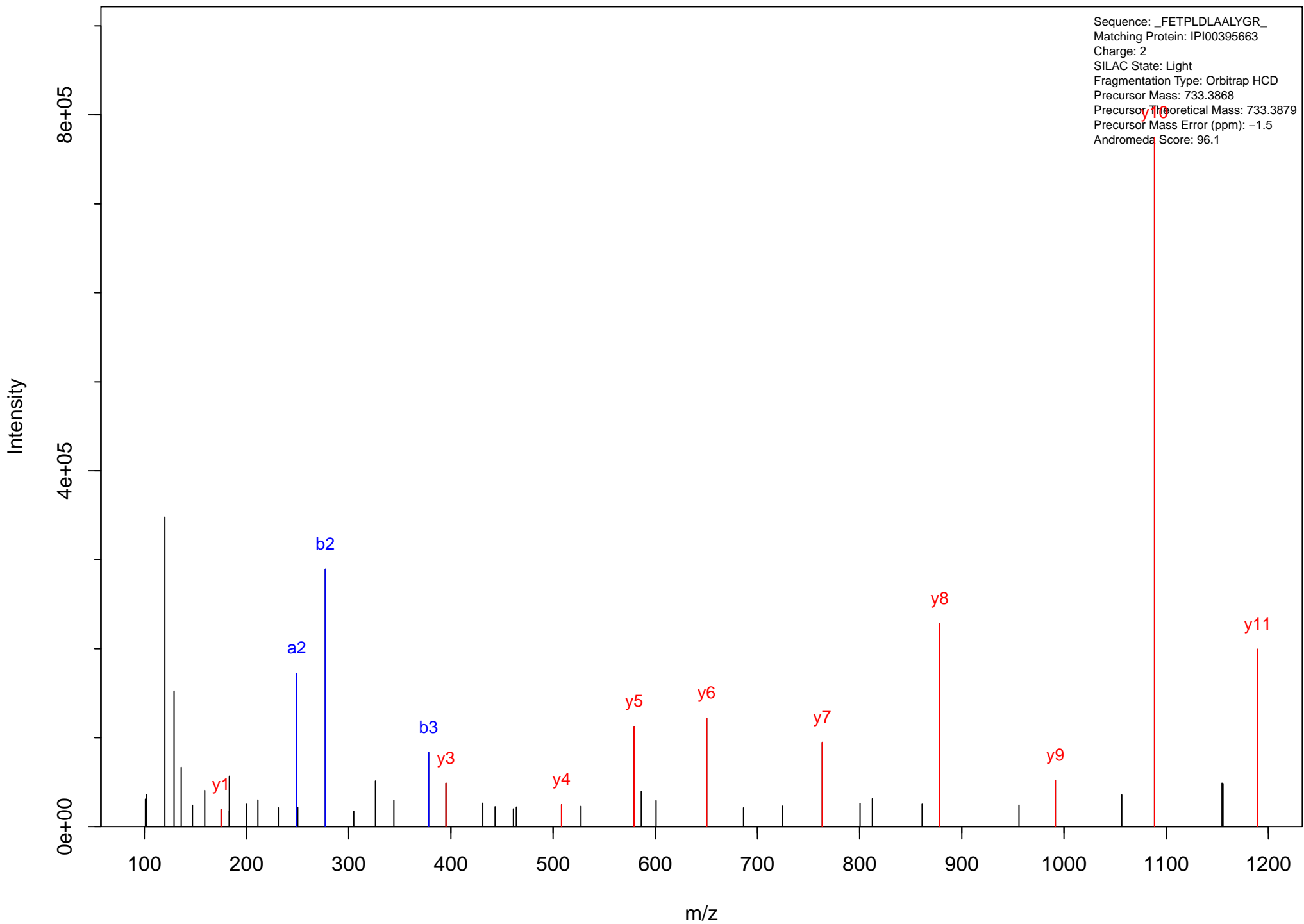


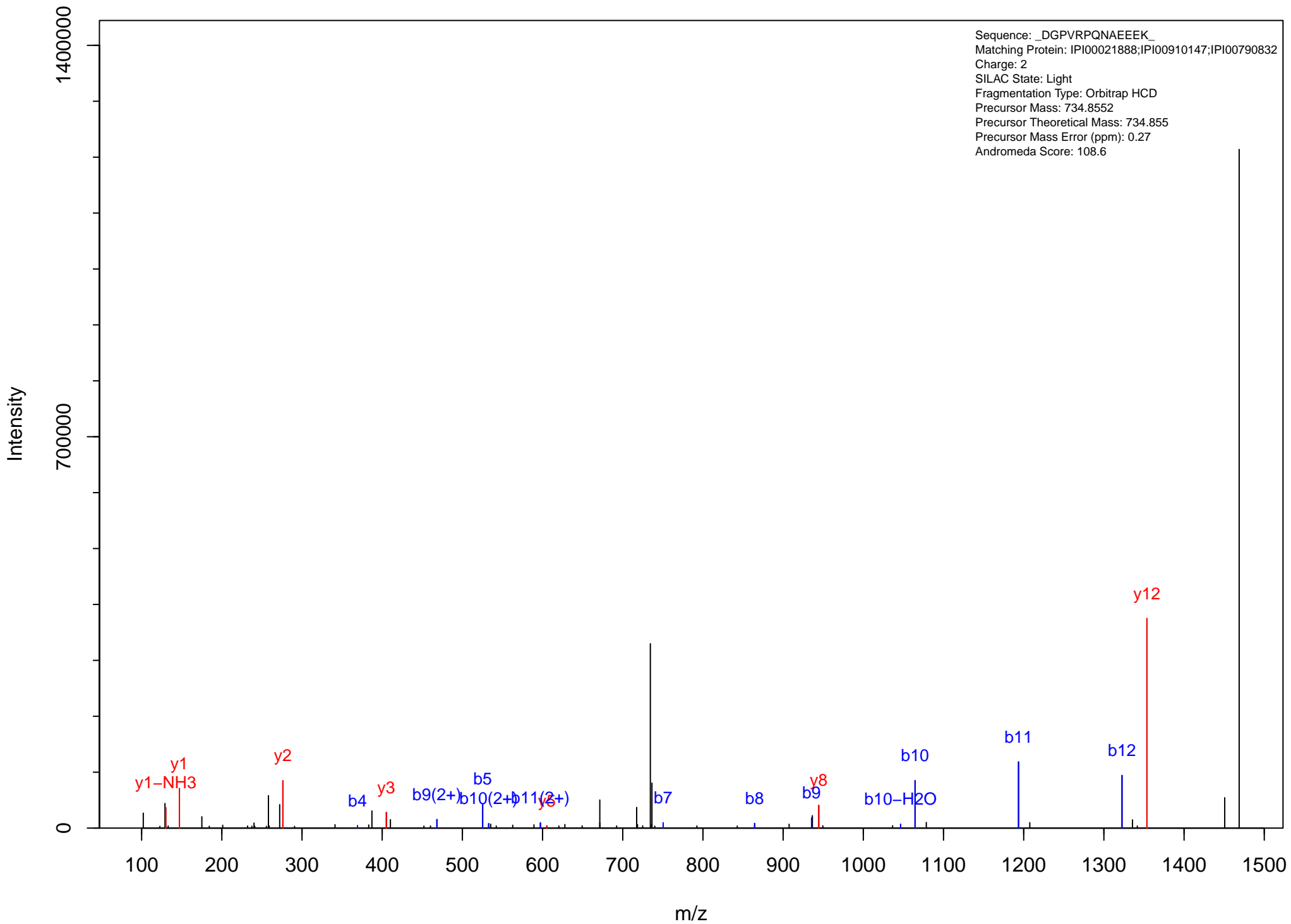


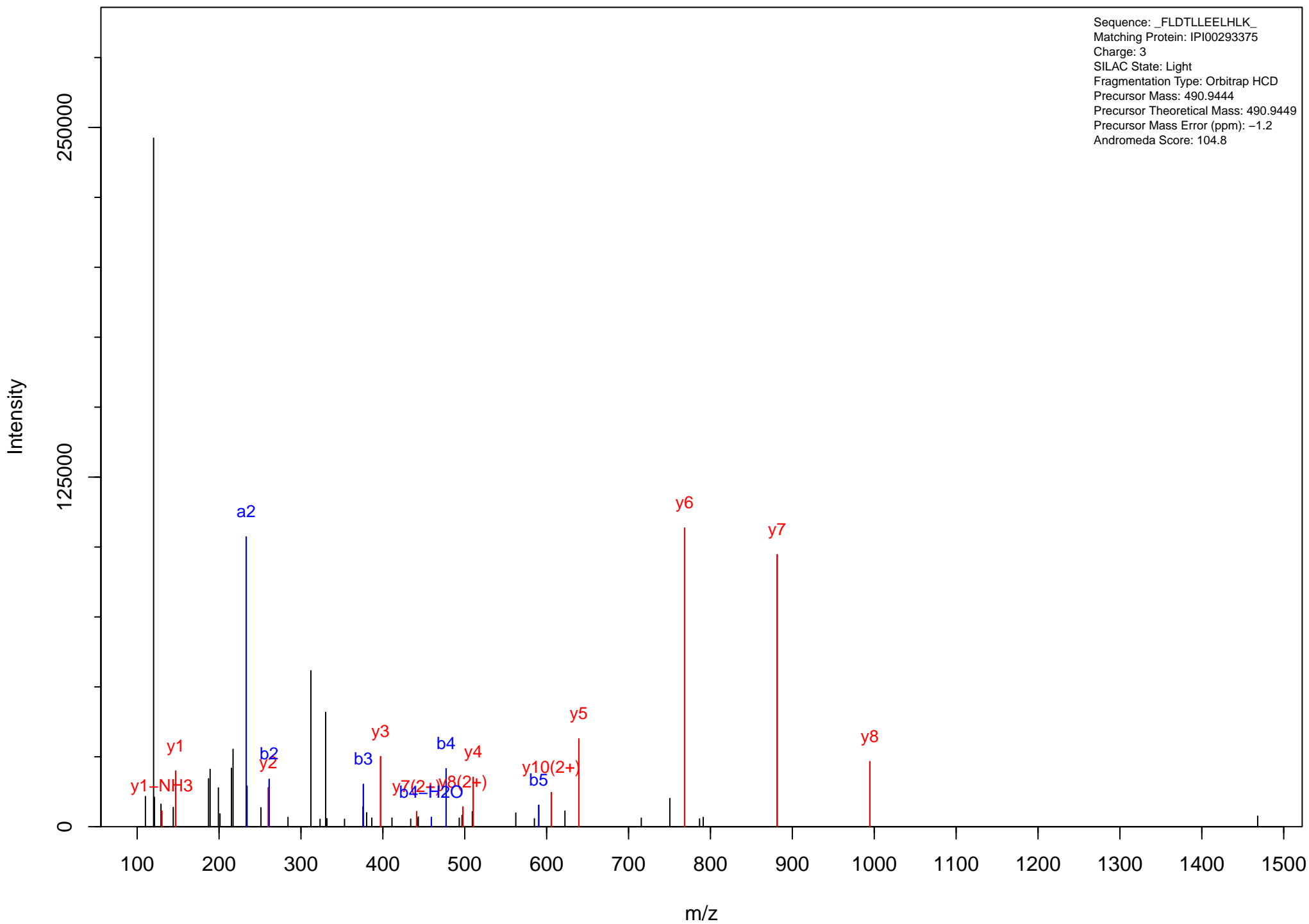


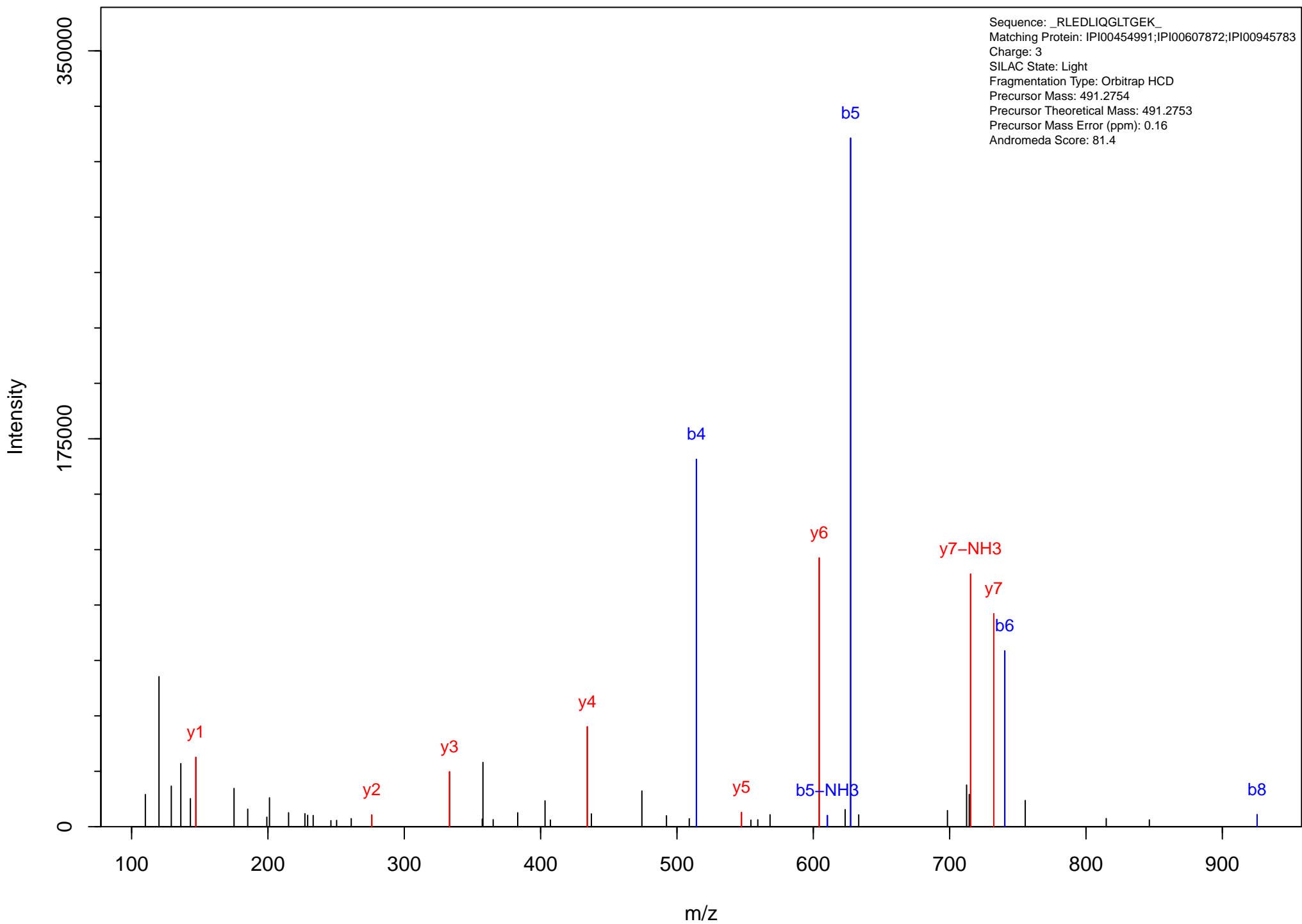


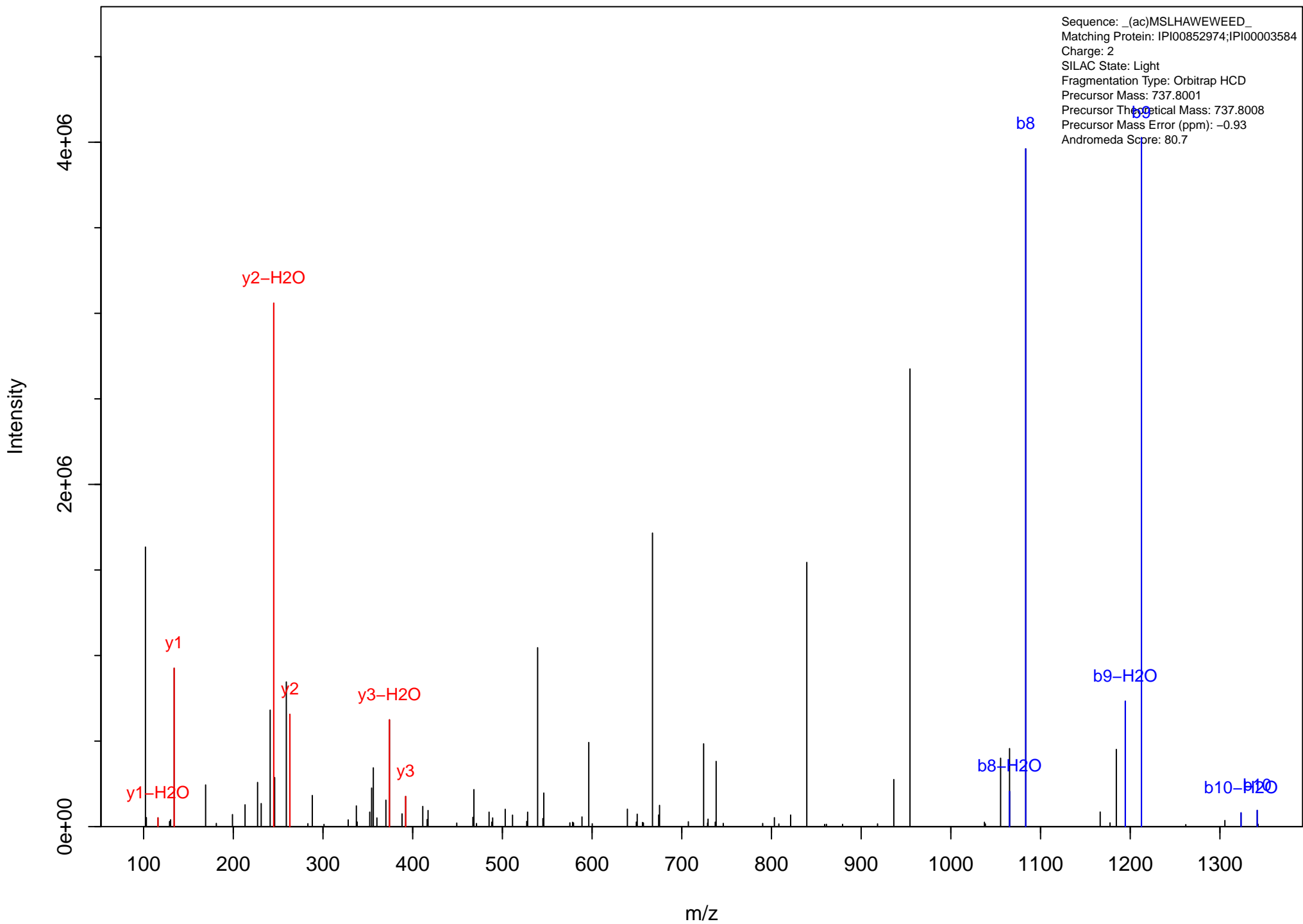


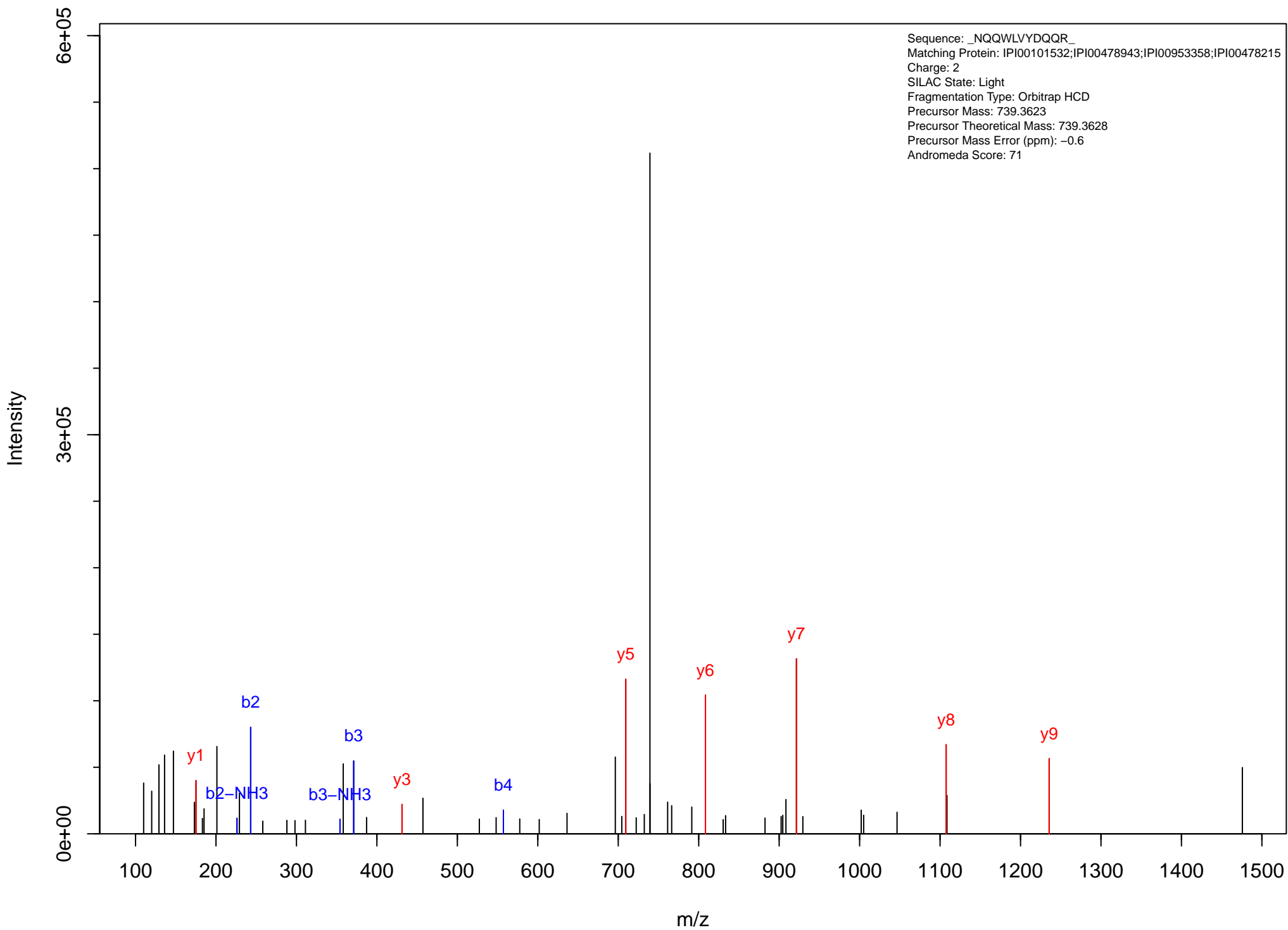


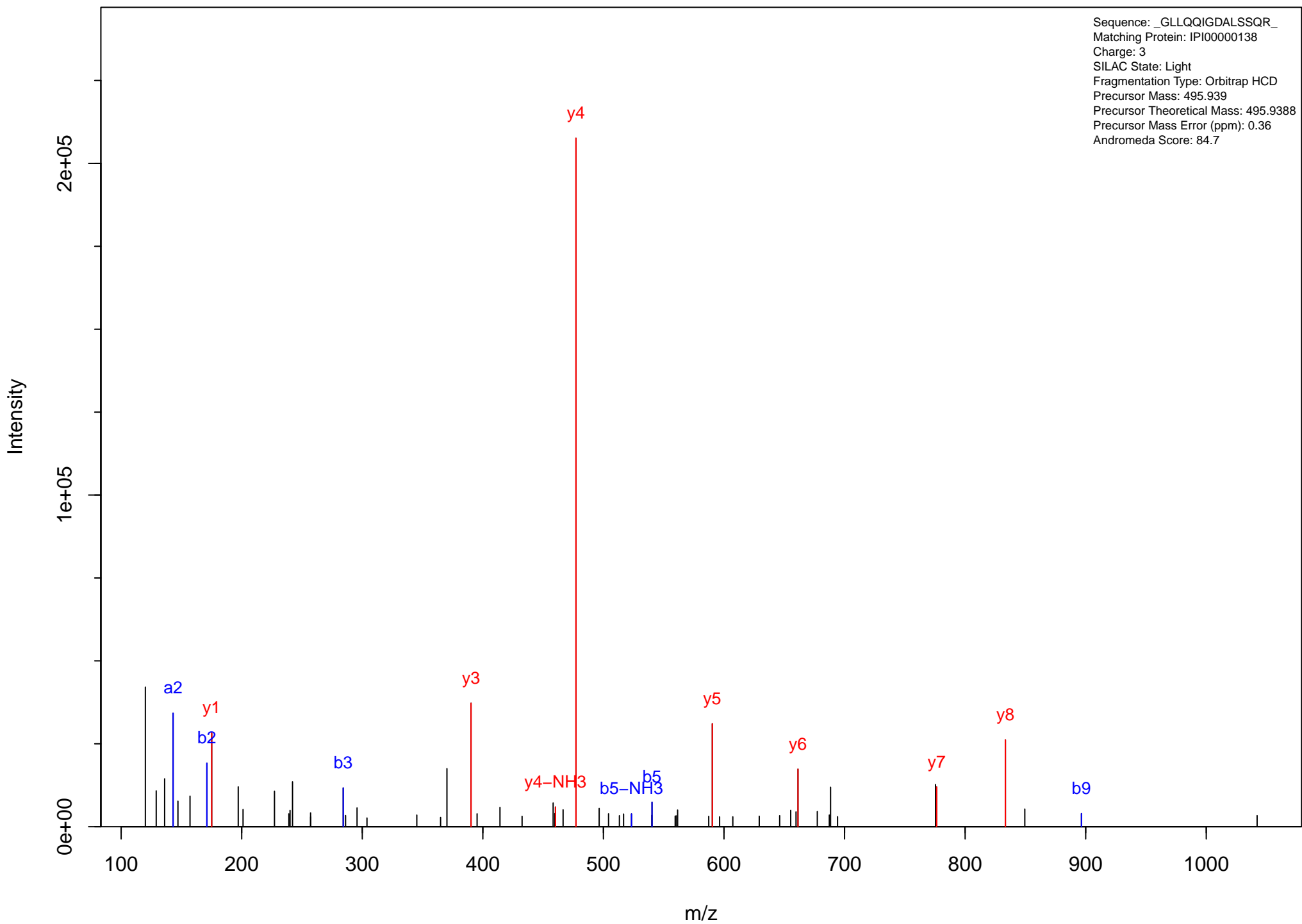




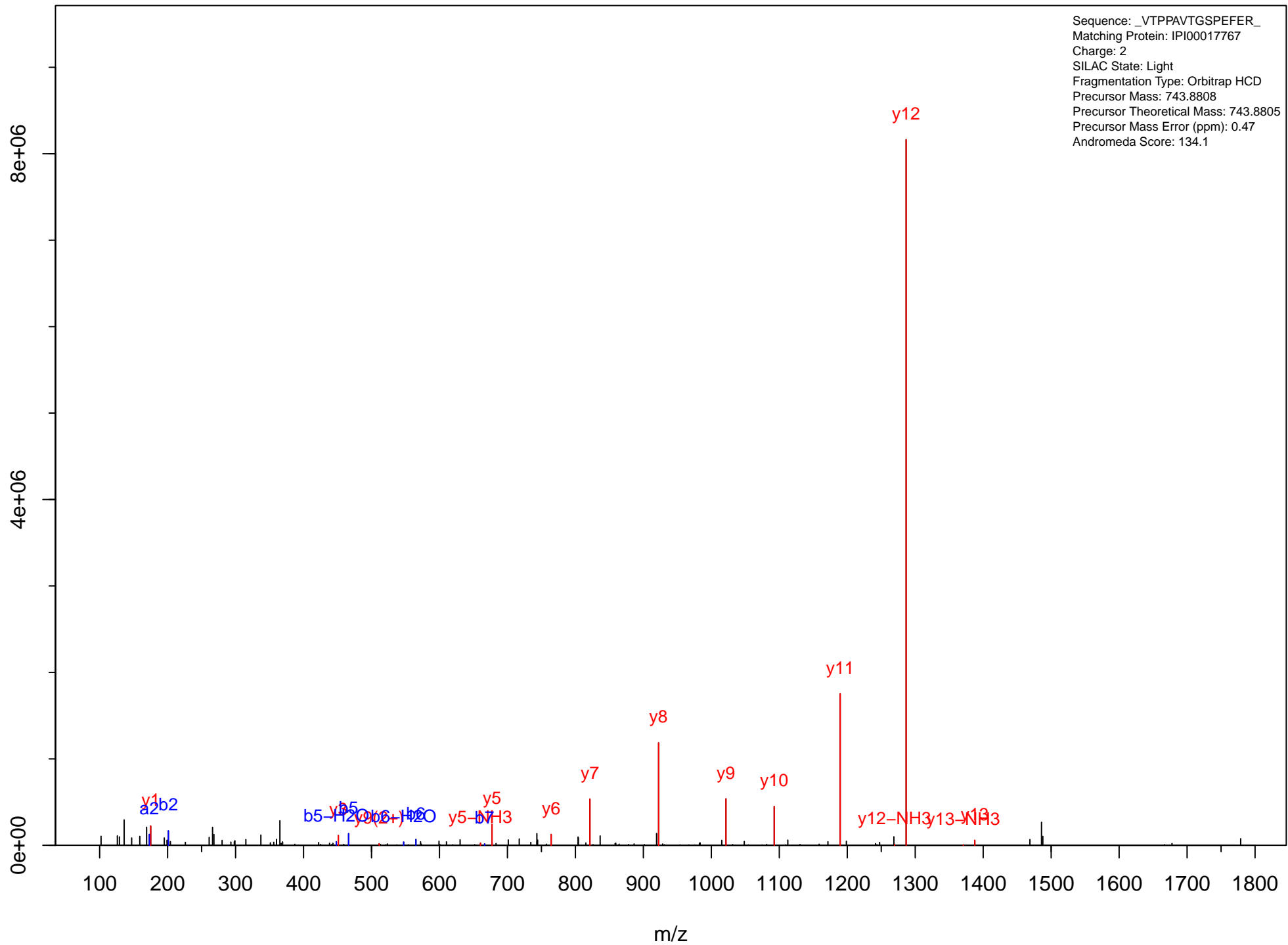


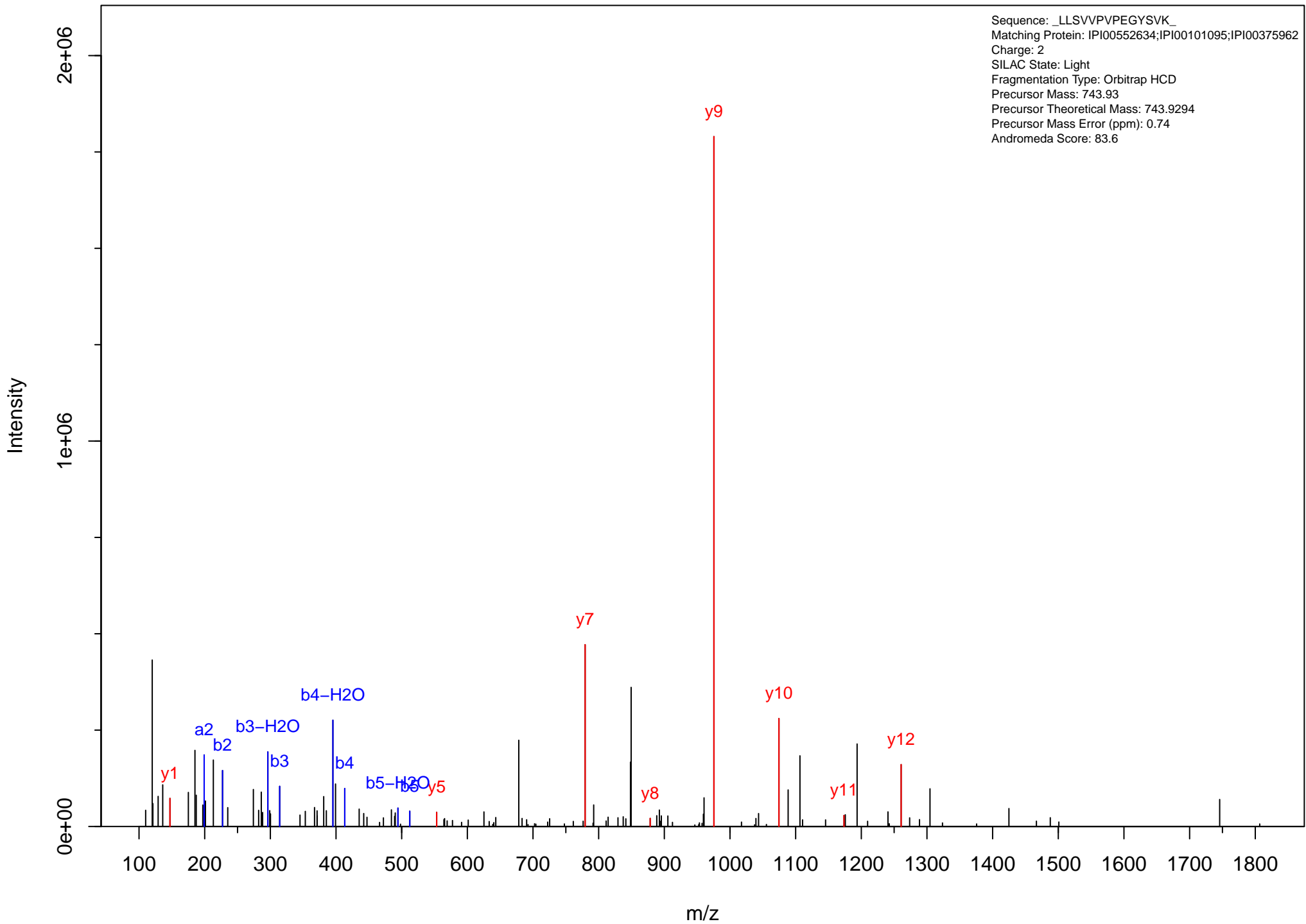


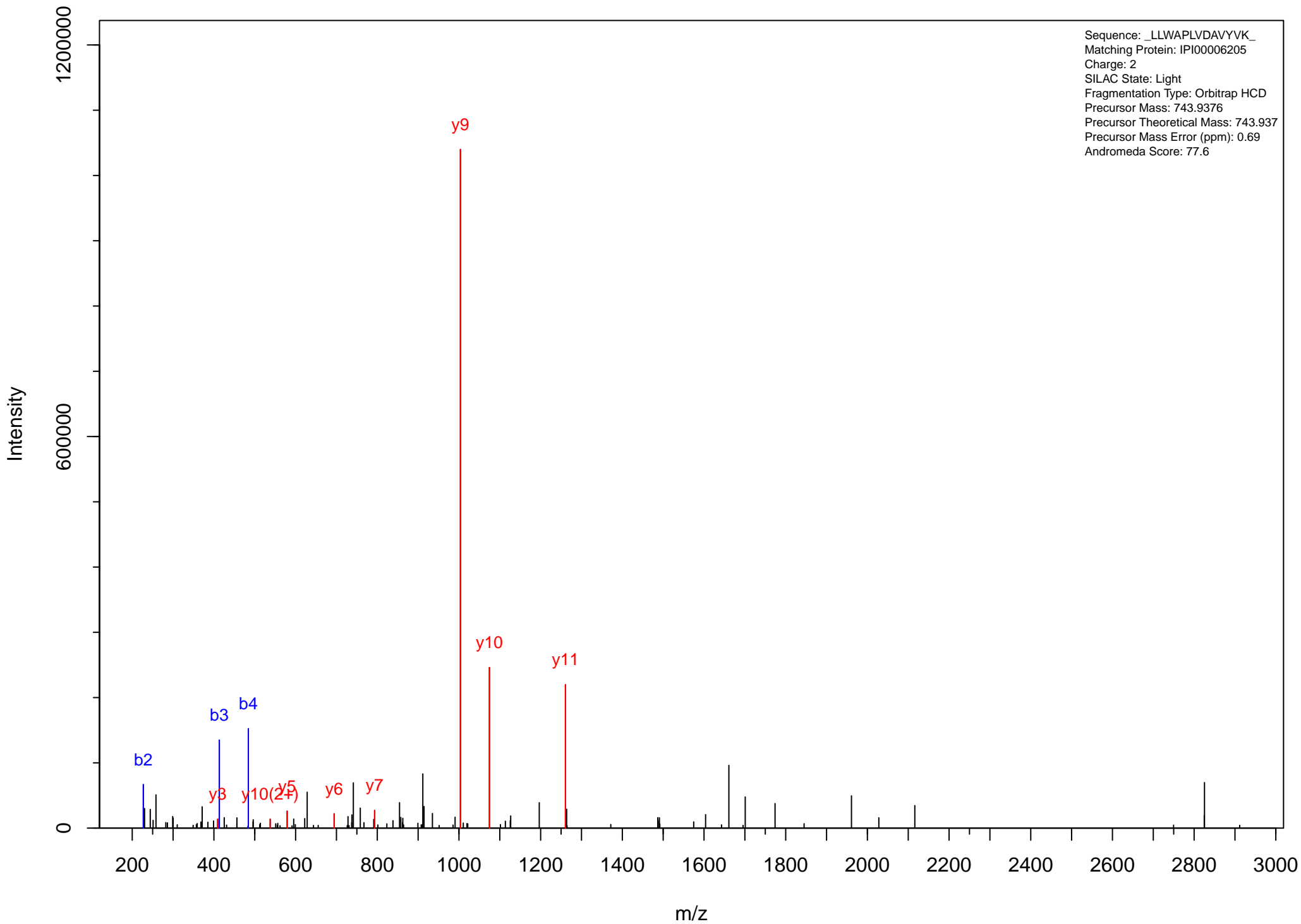


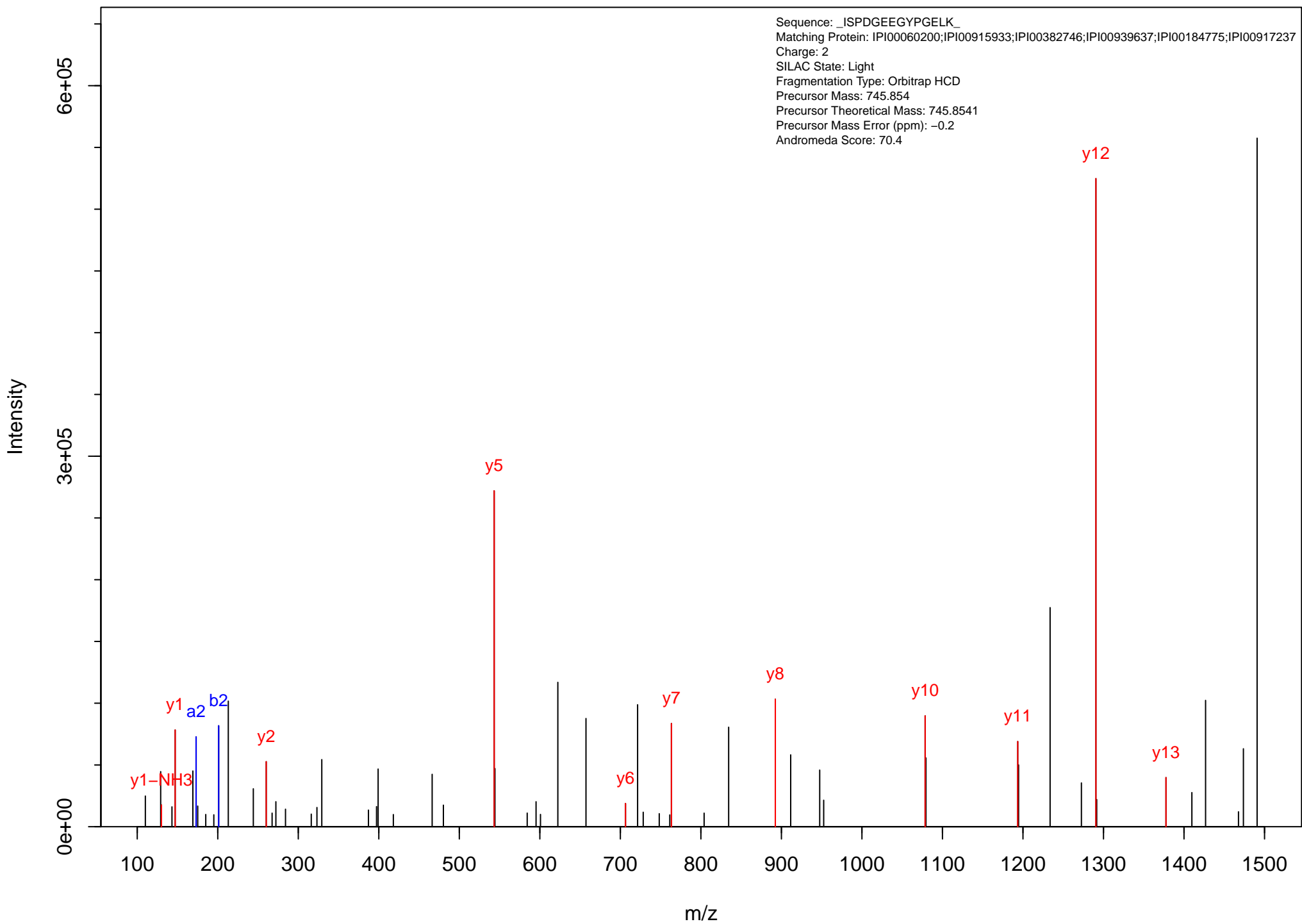


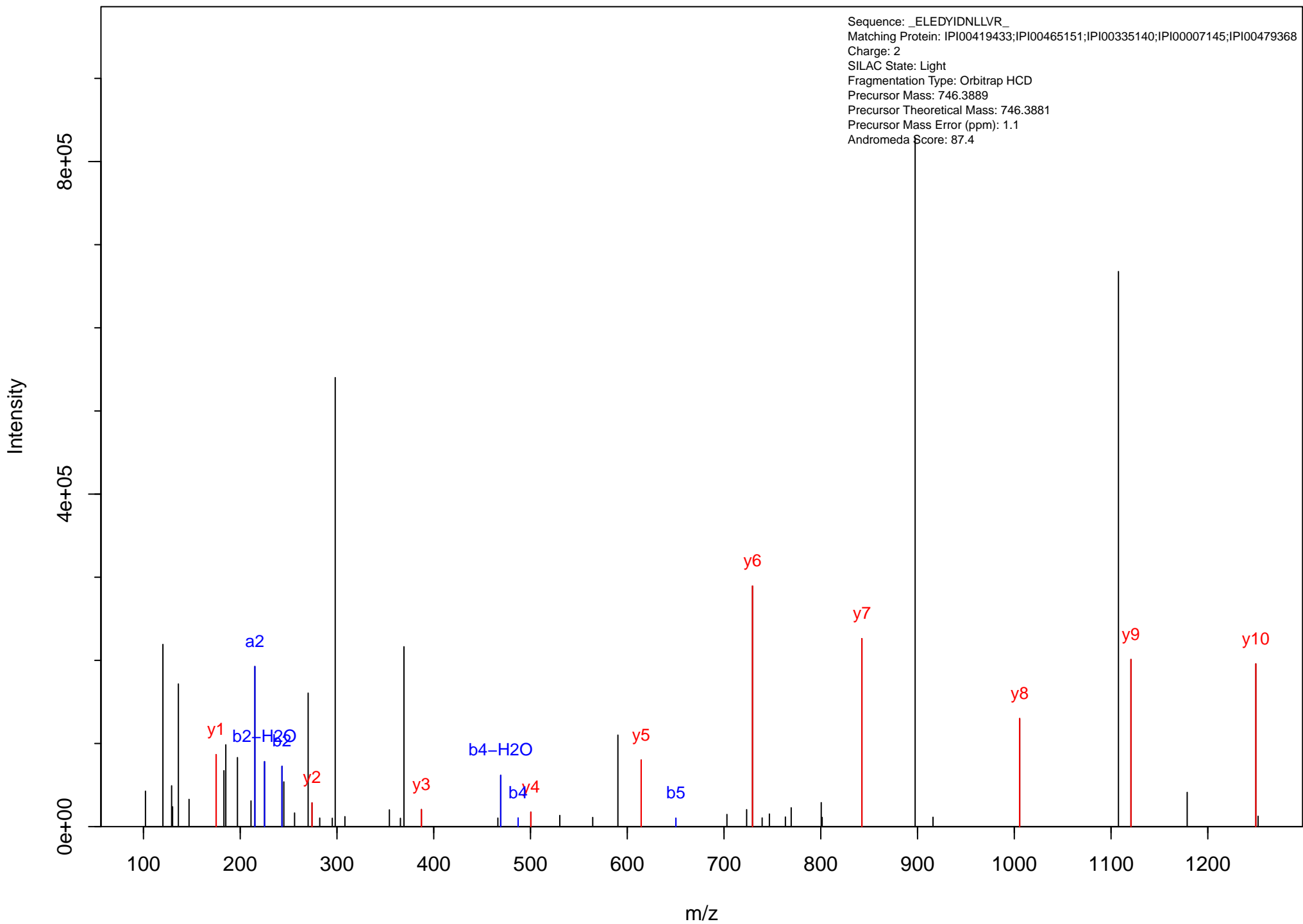
Intensity

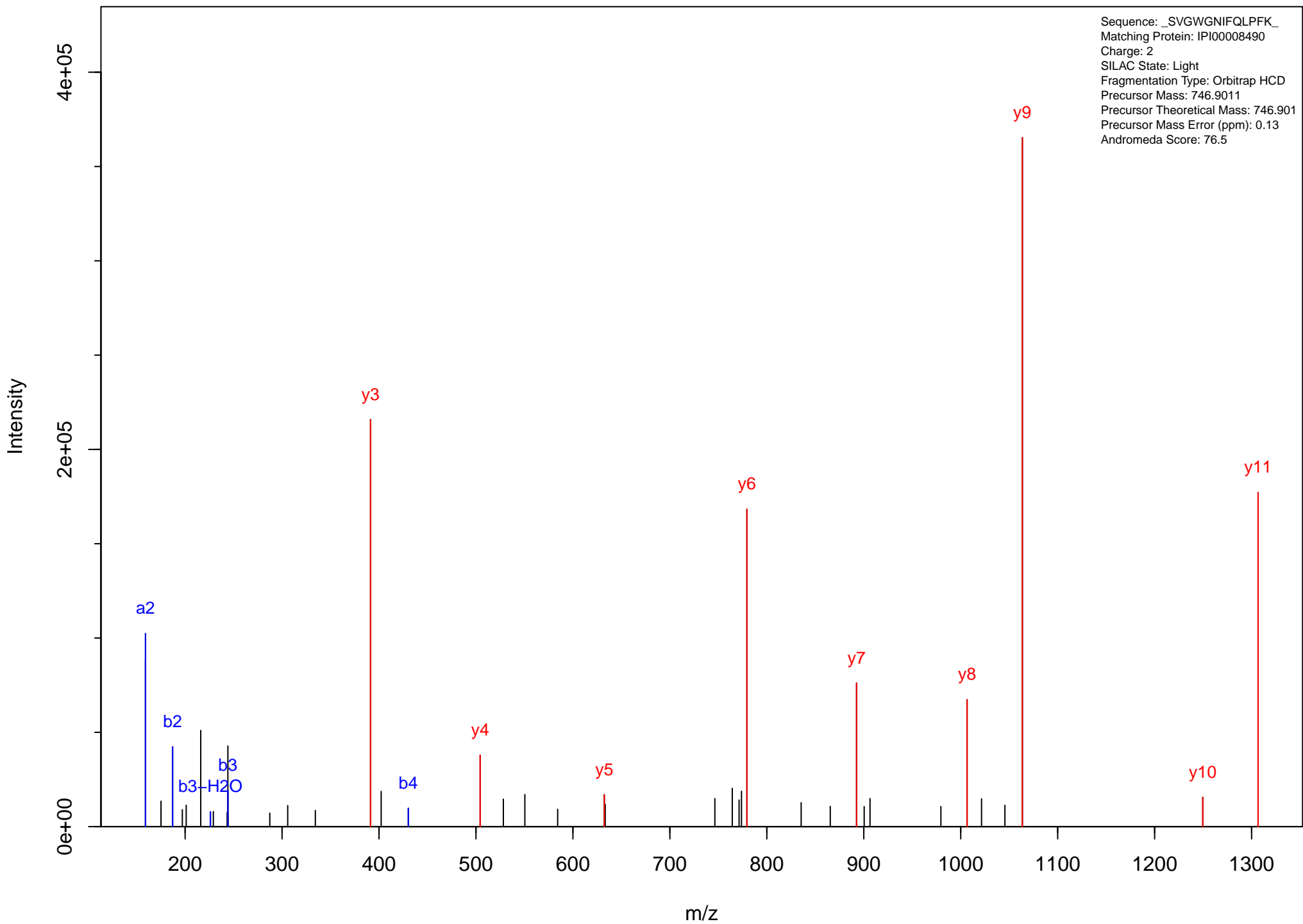


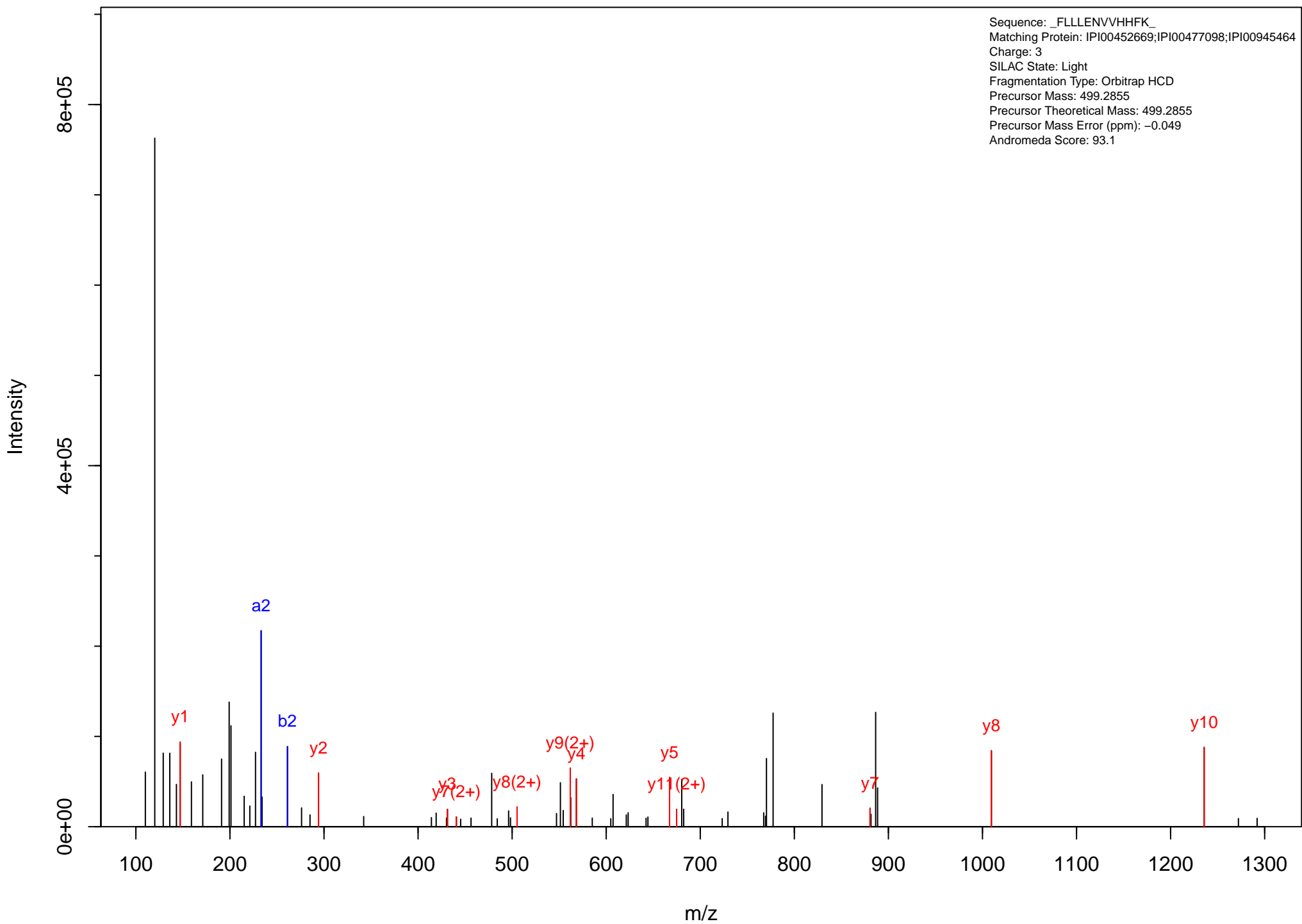


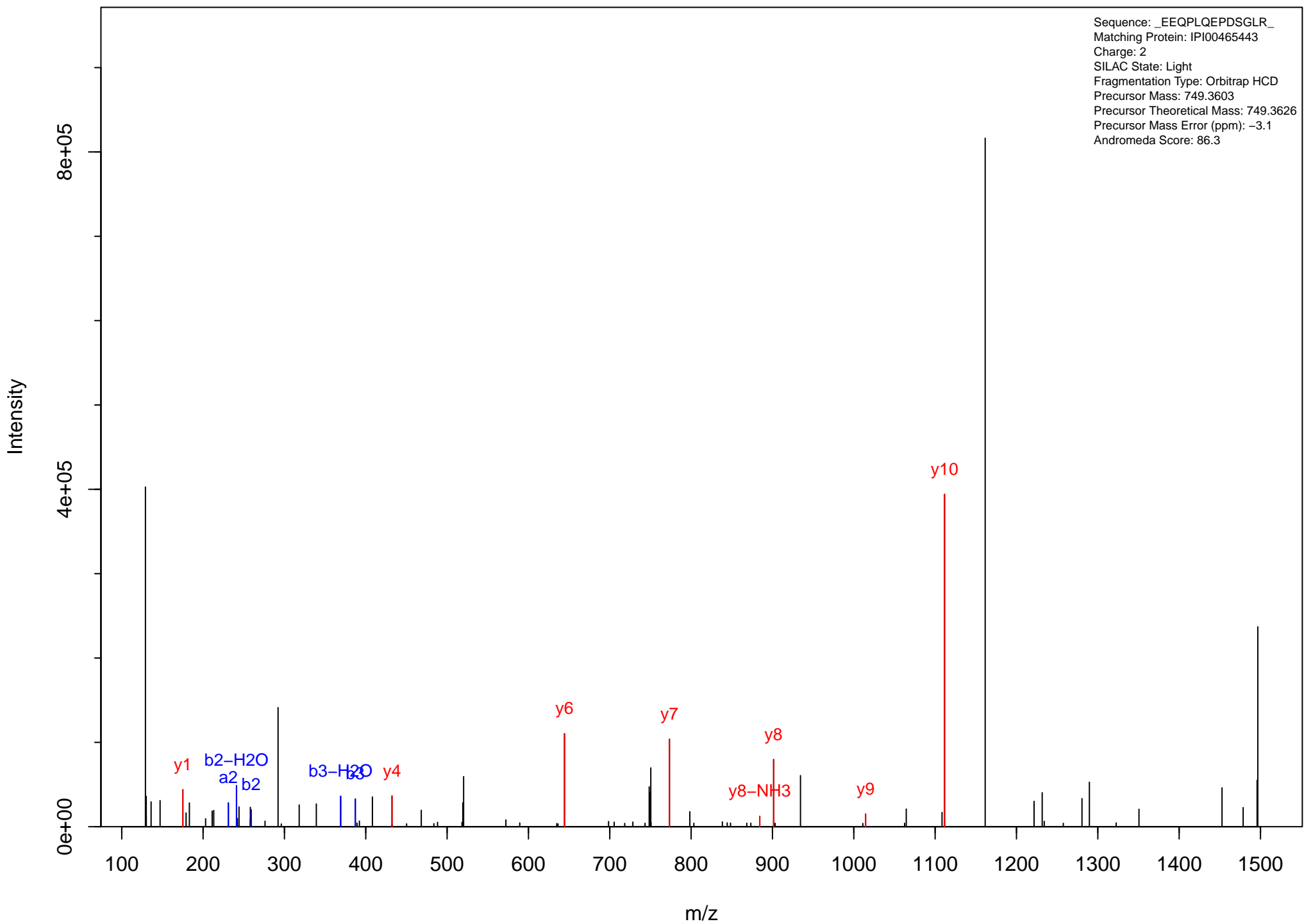


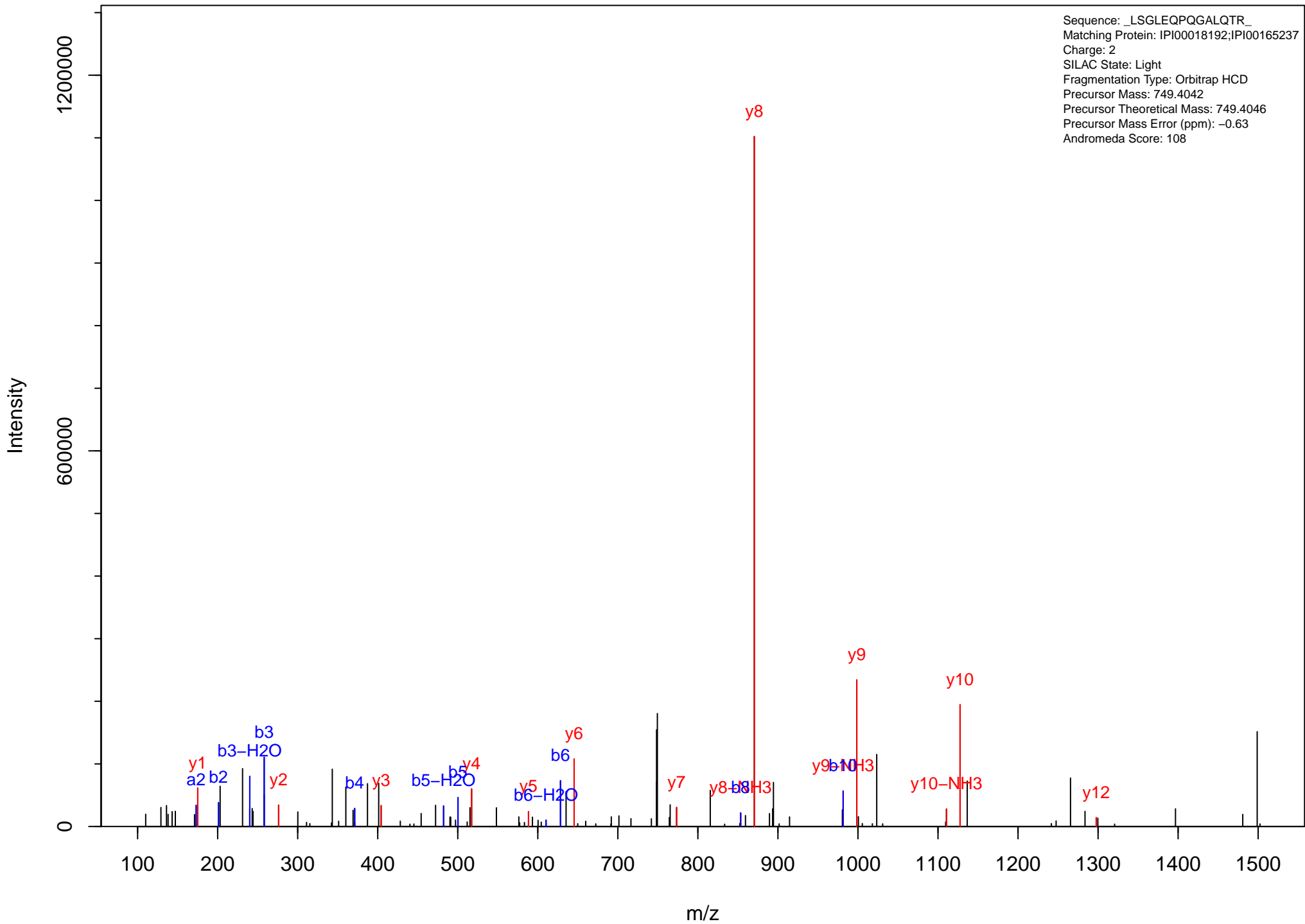


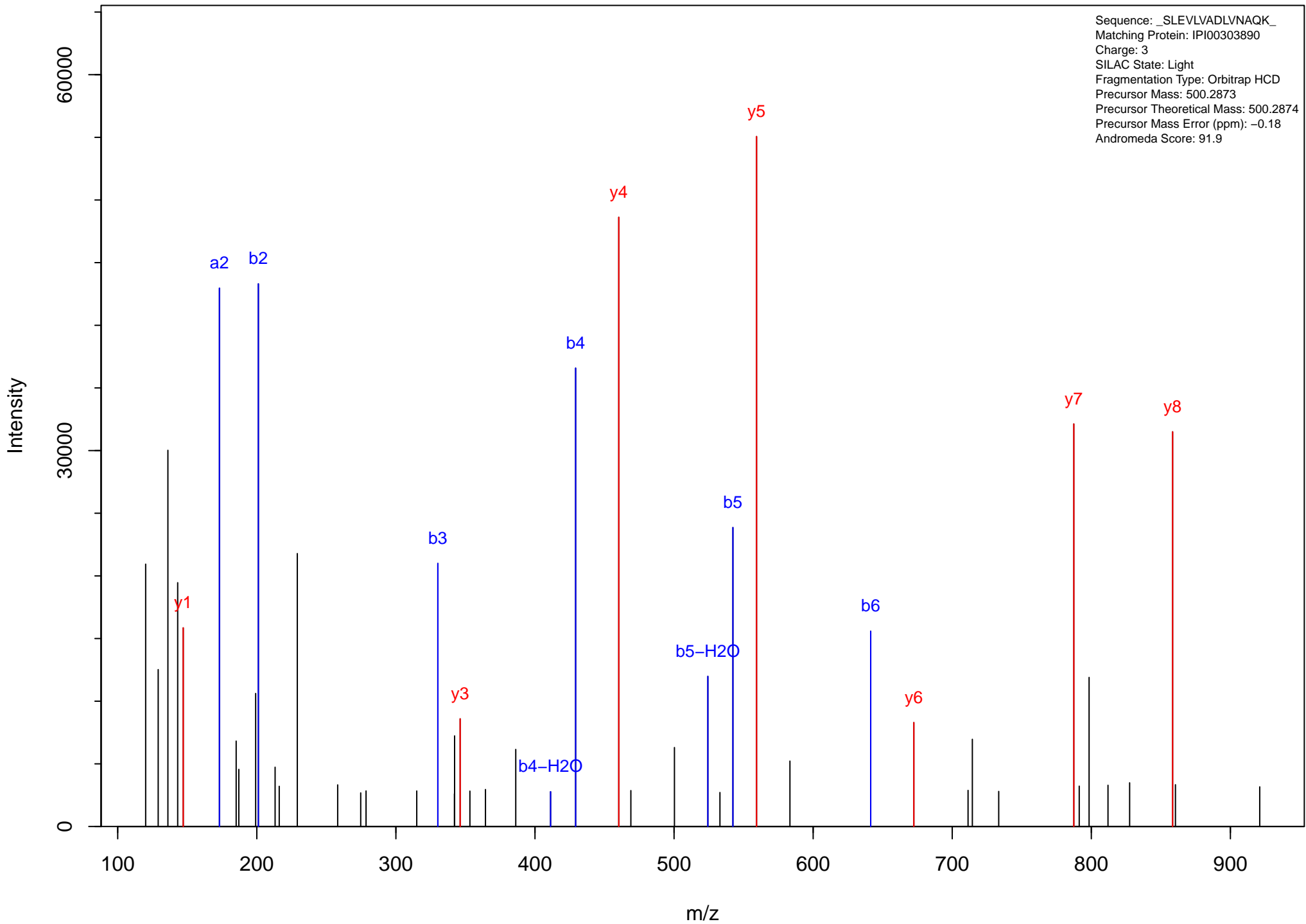


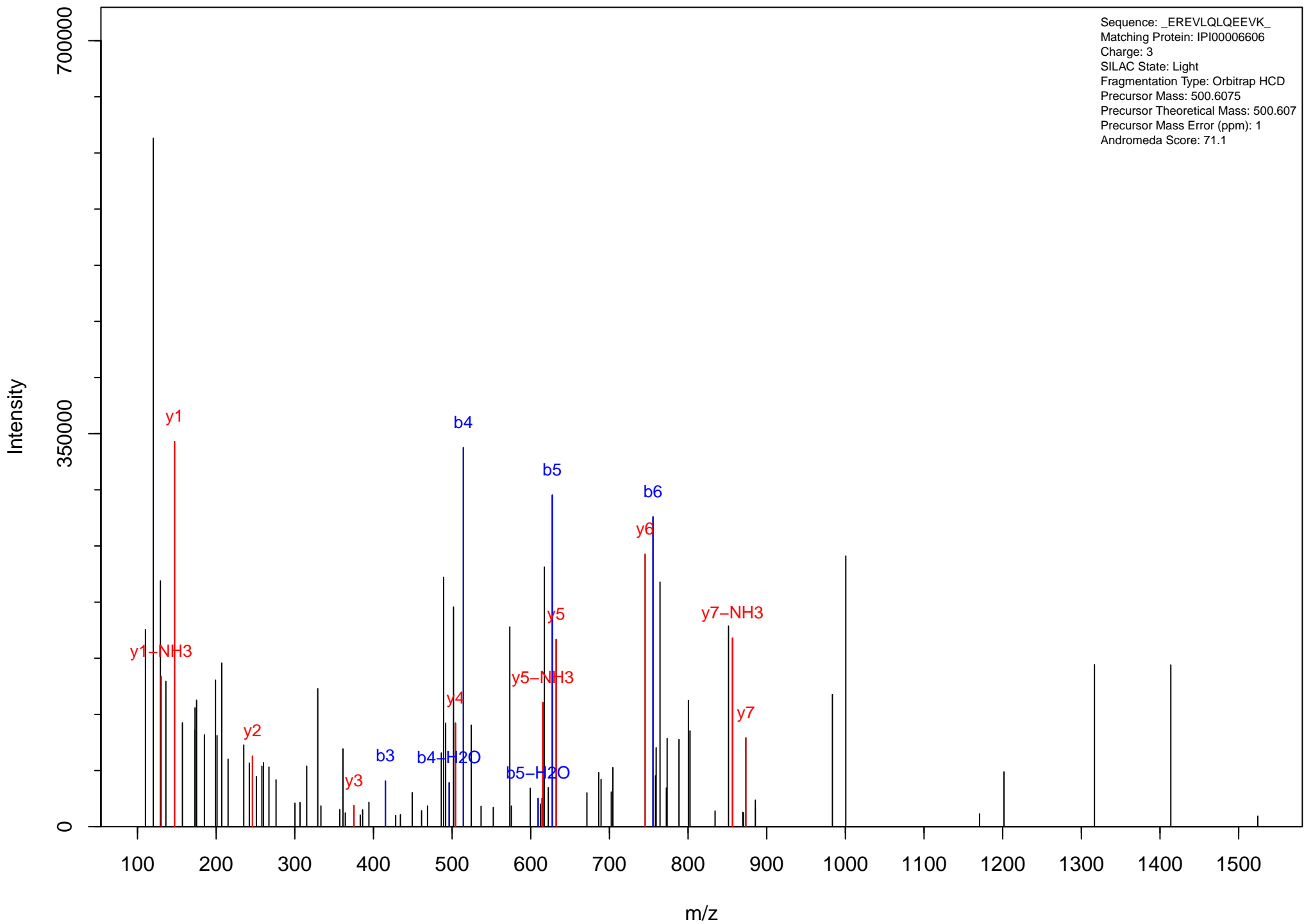


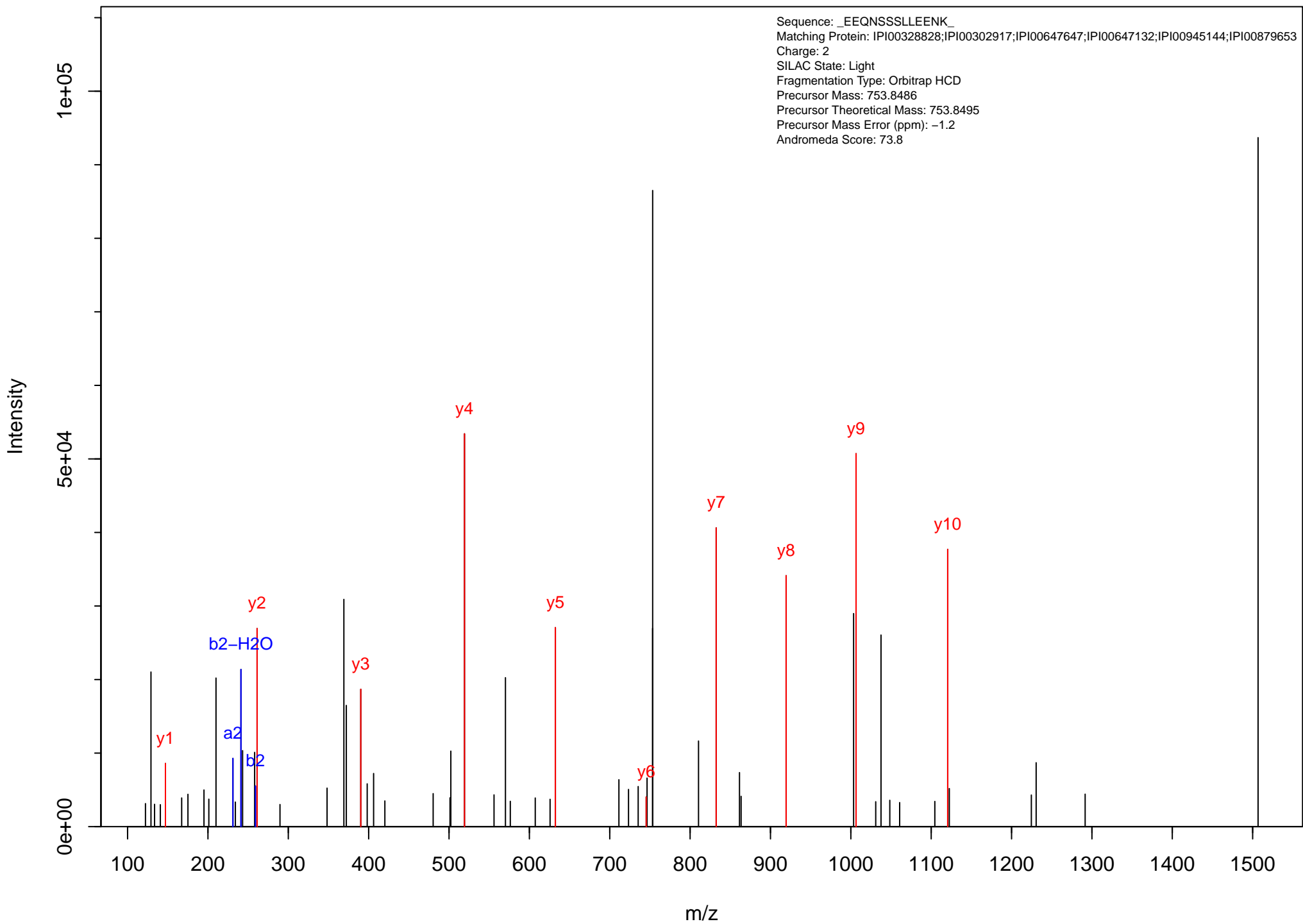


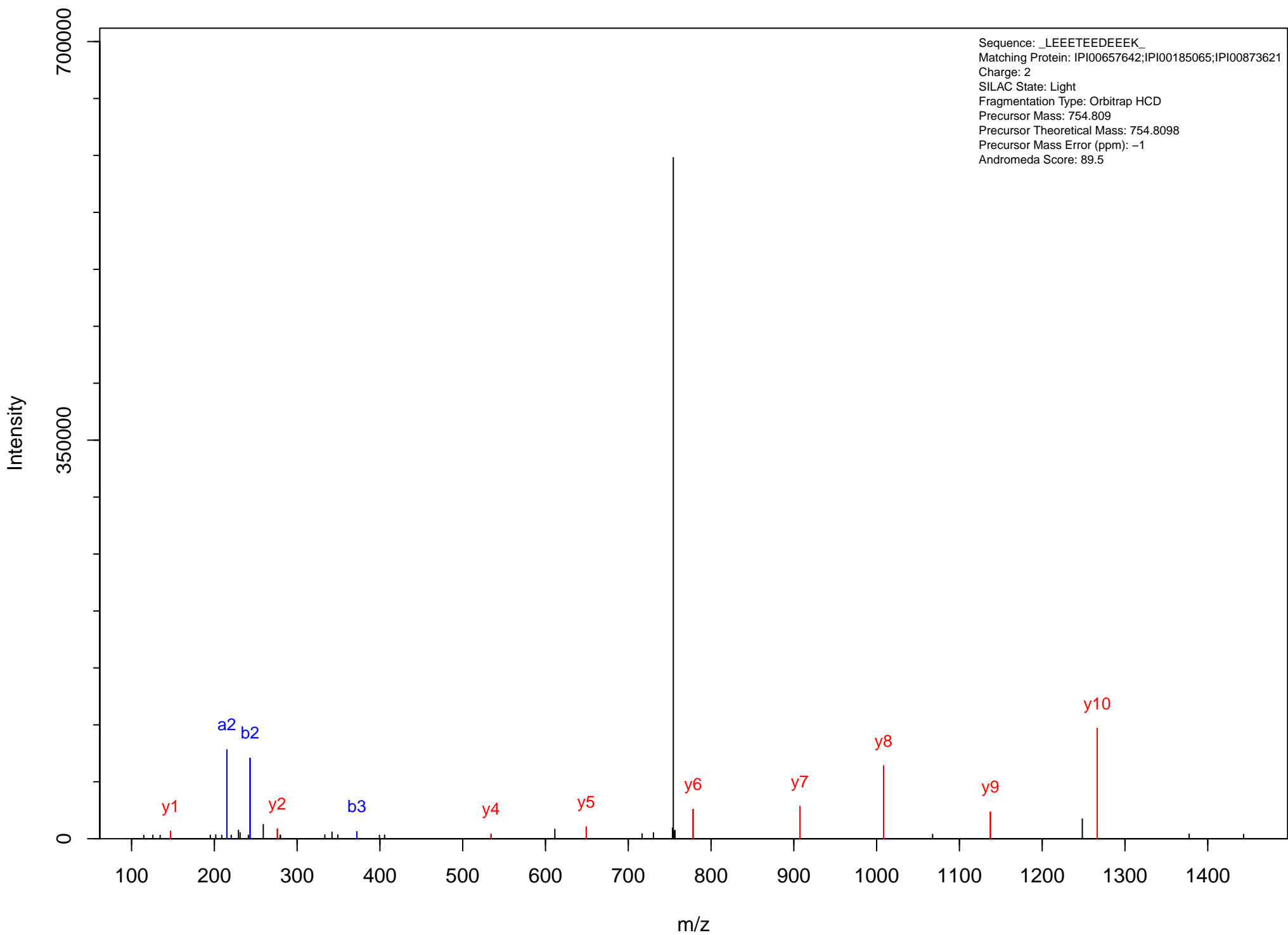


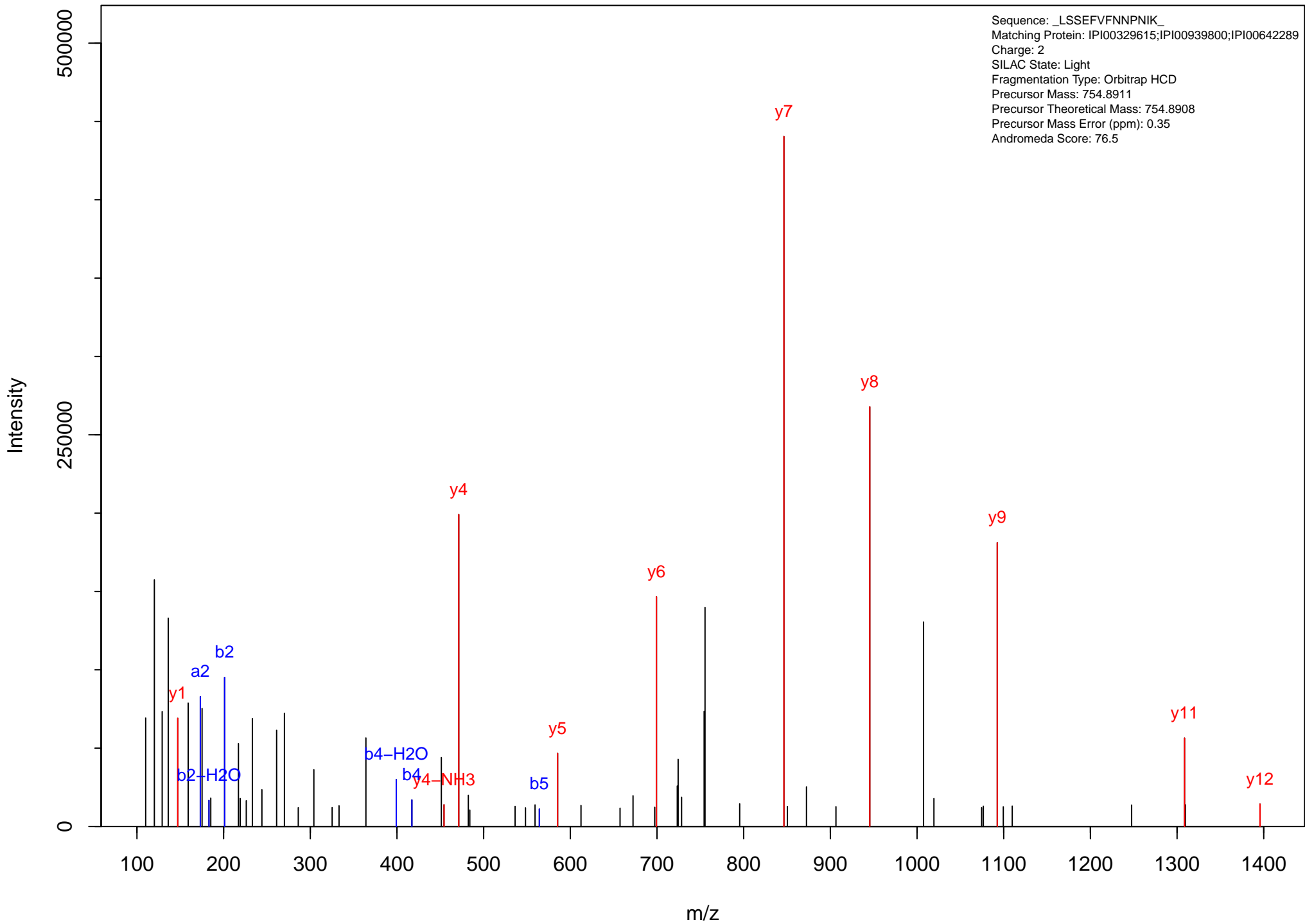


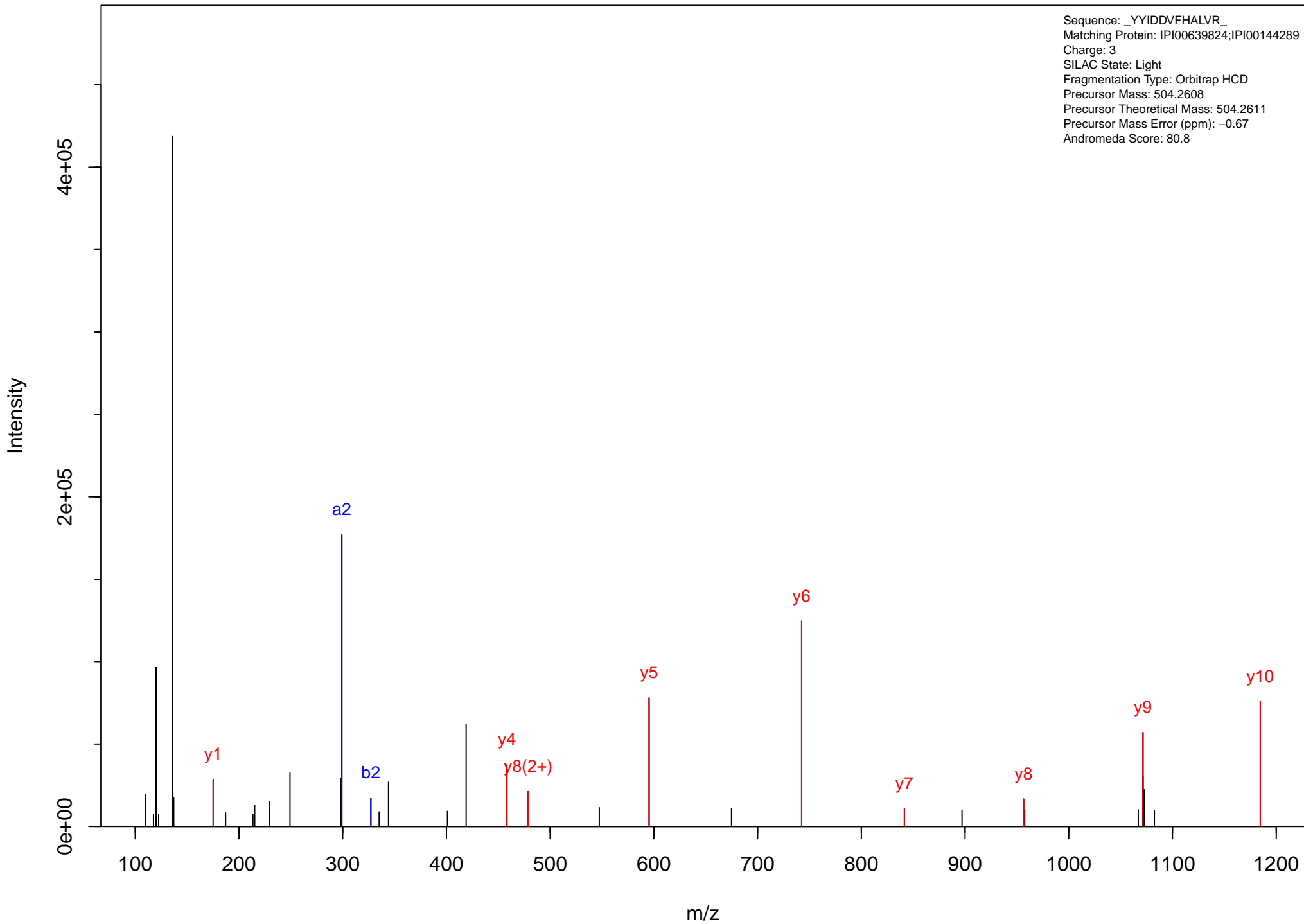


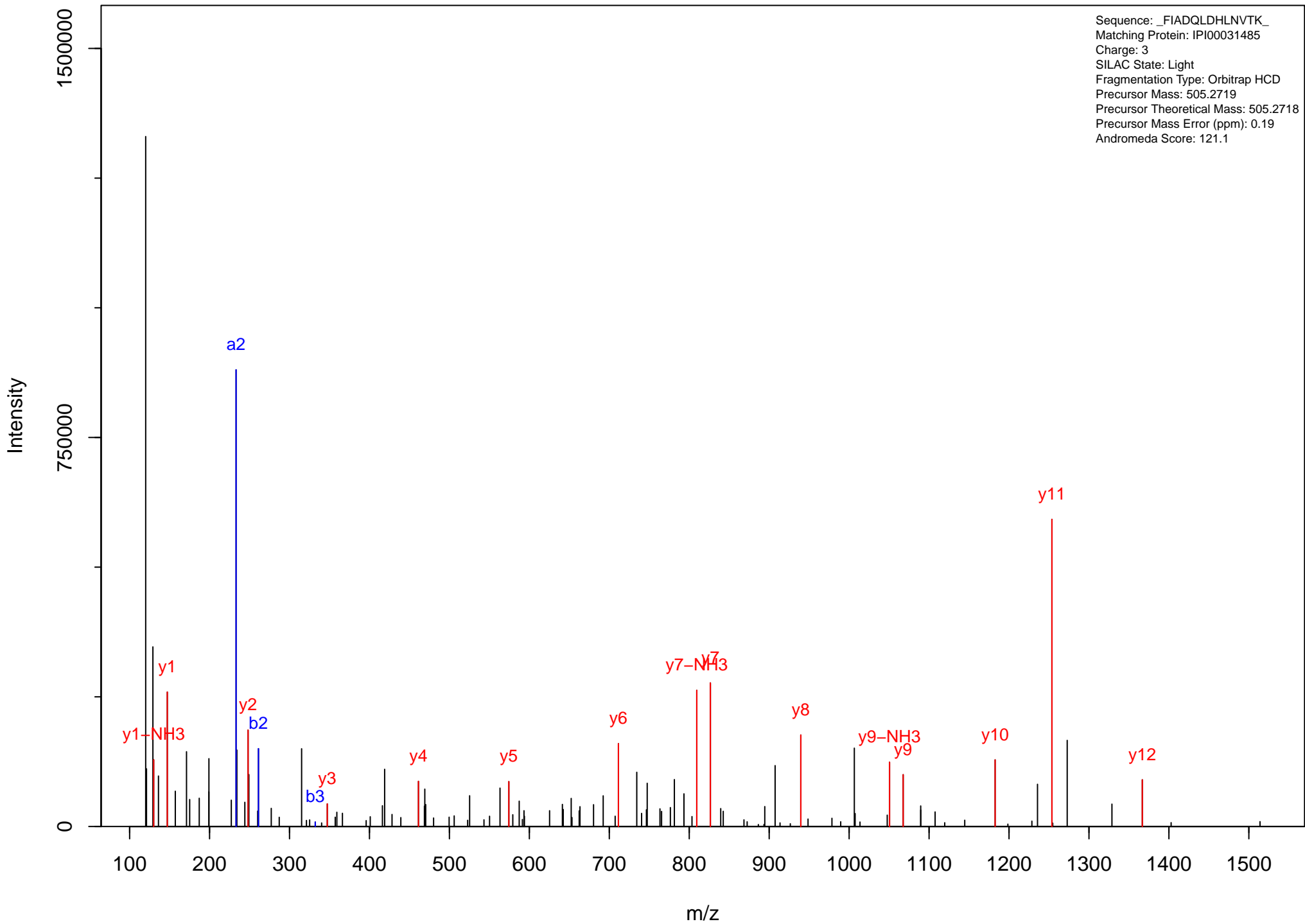


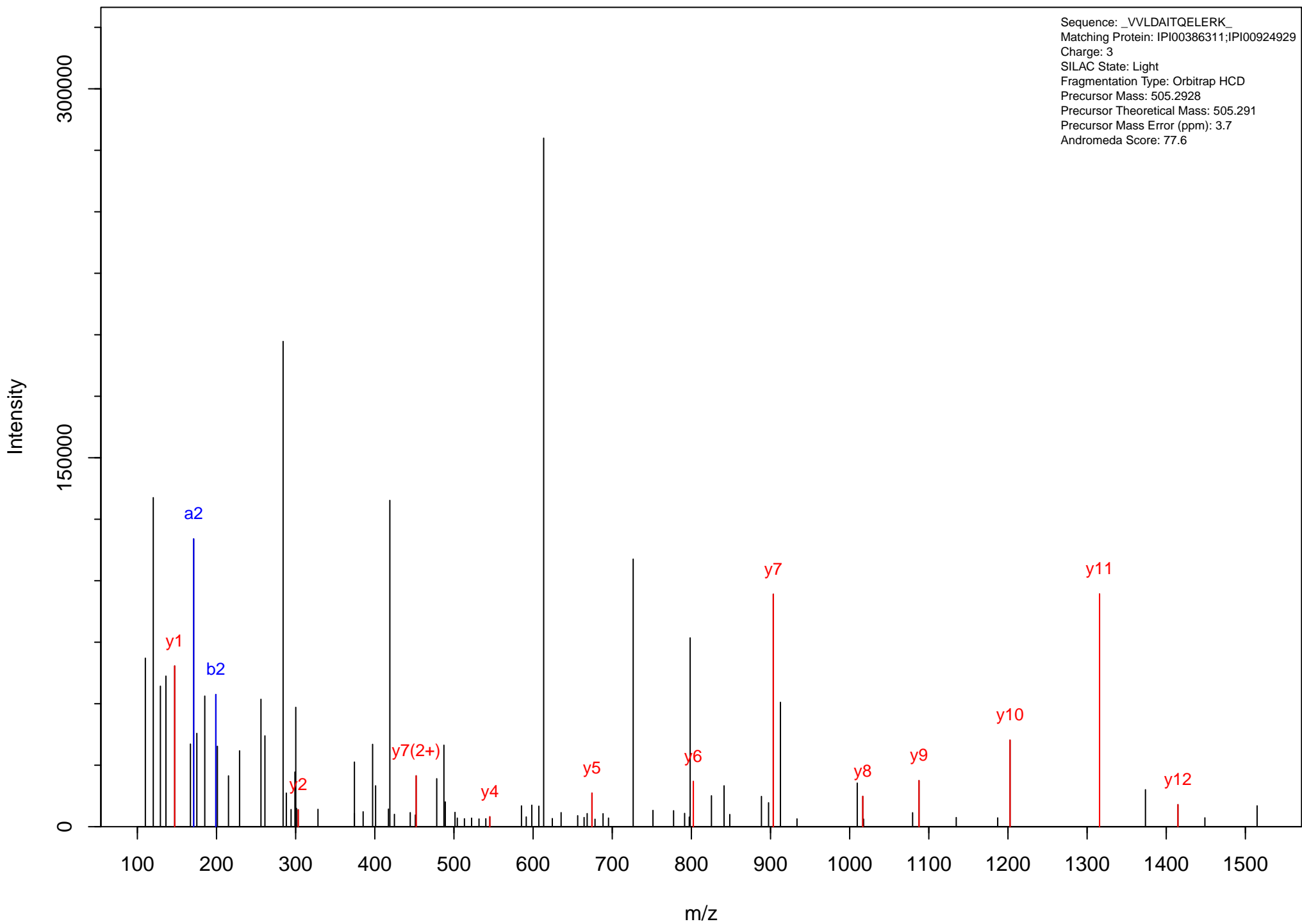


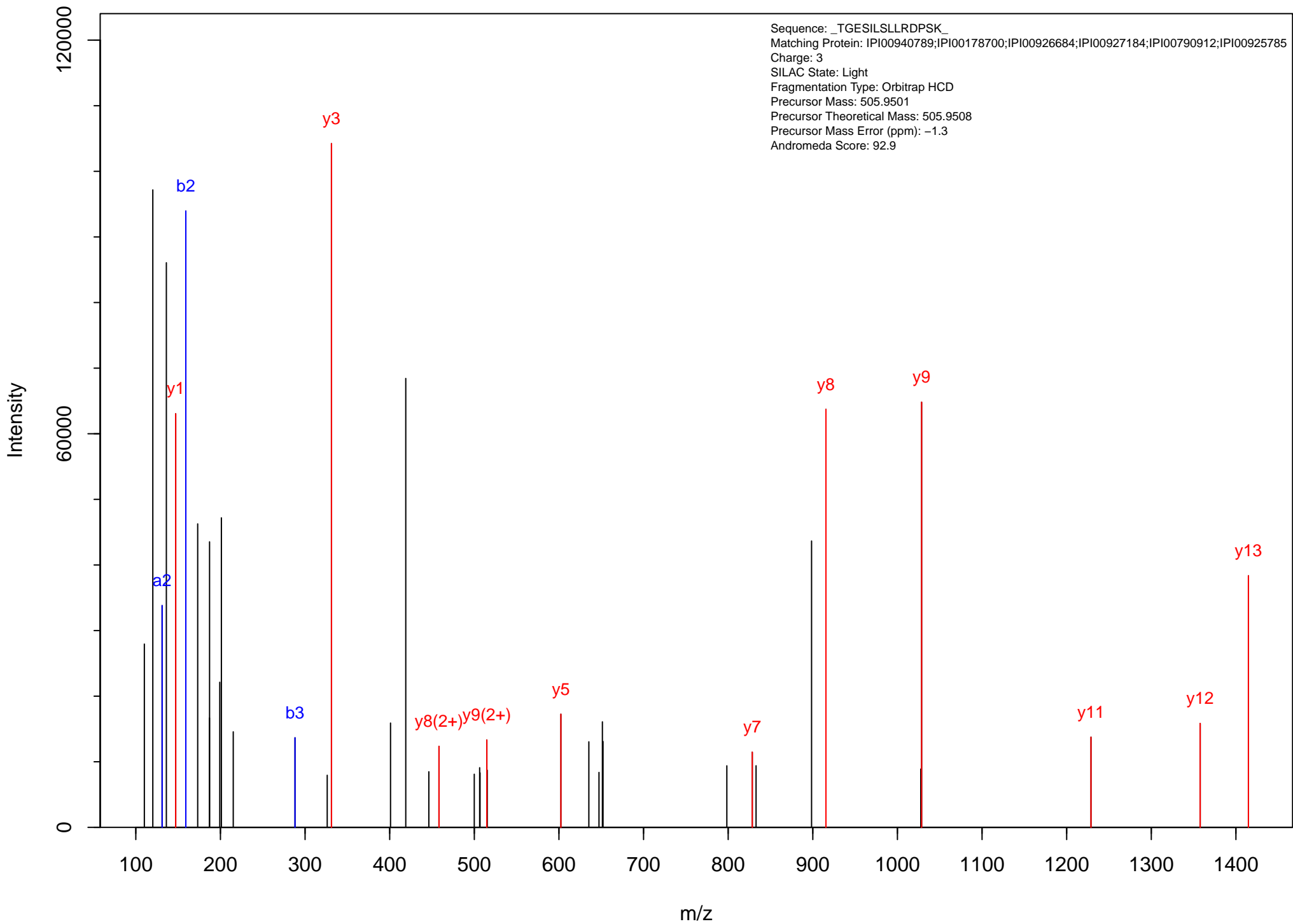


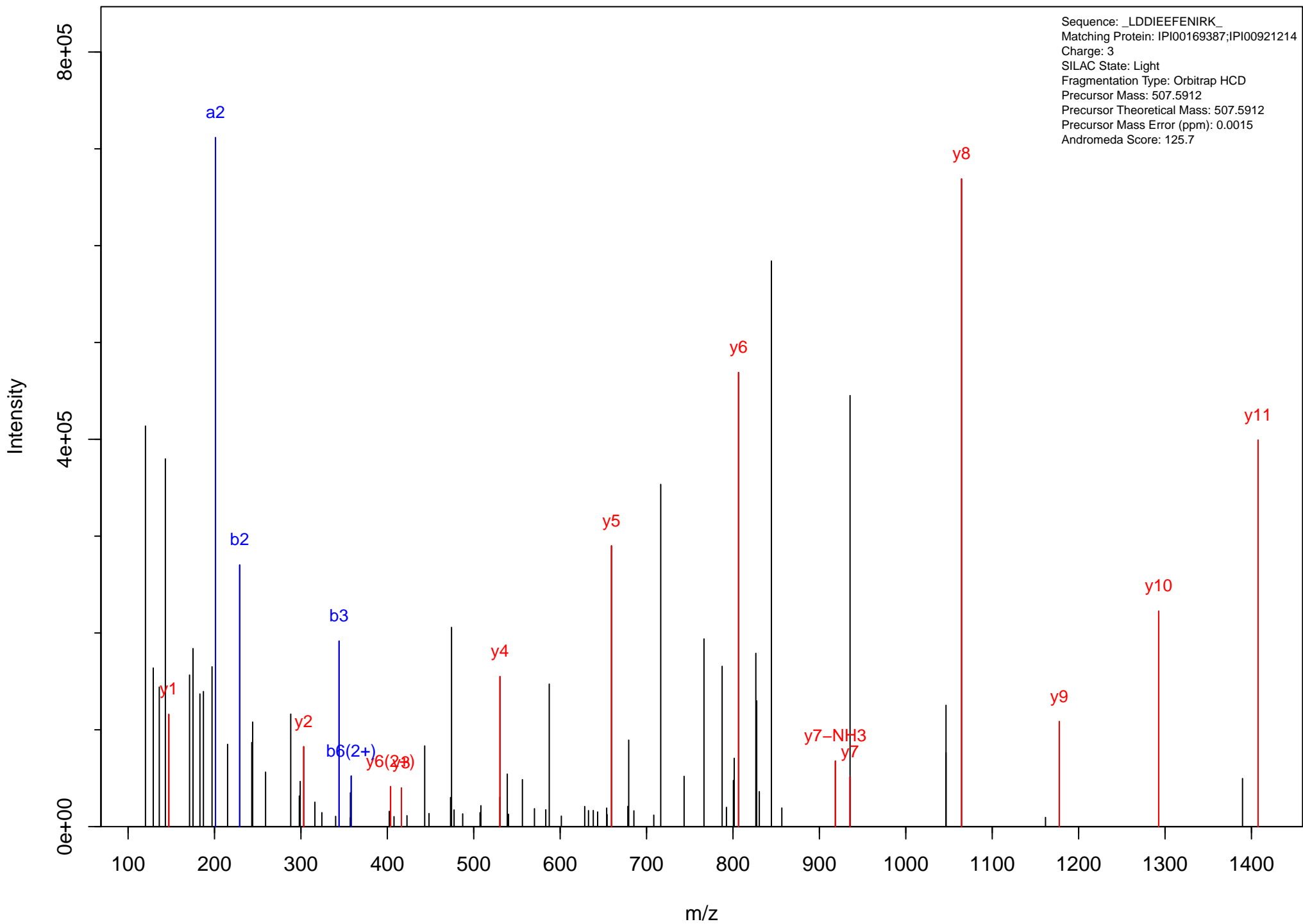


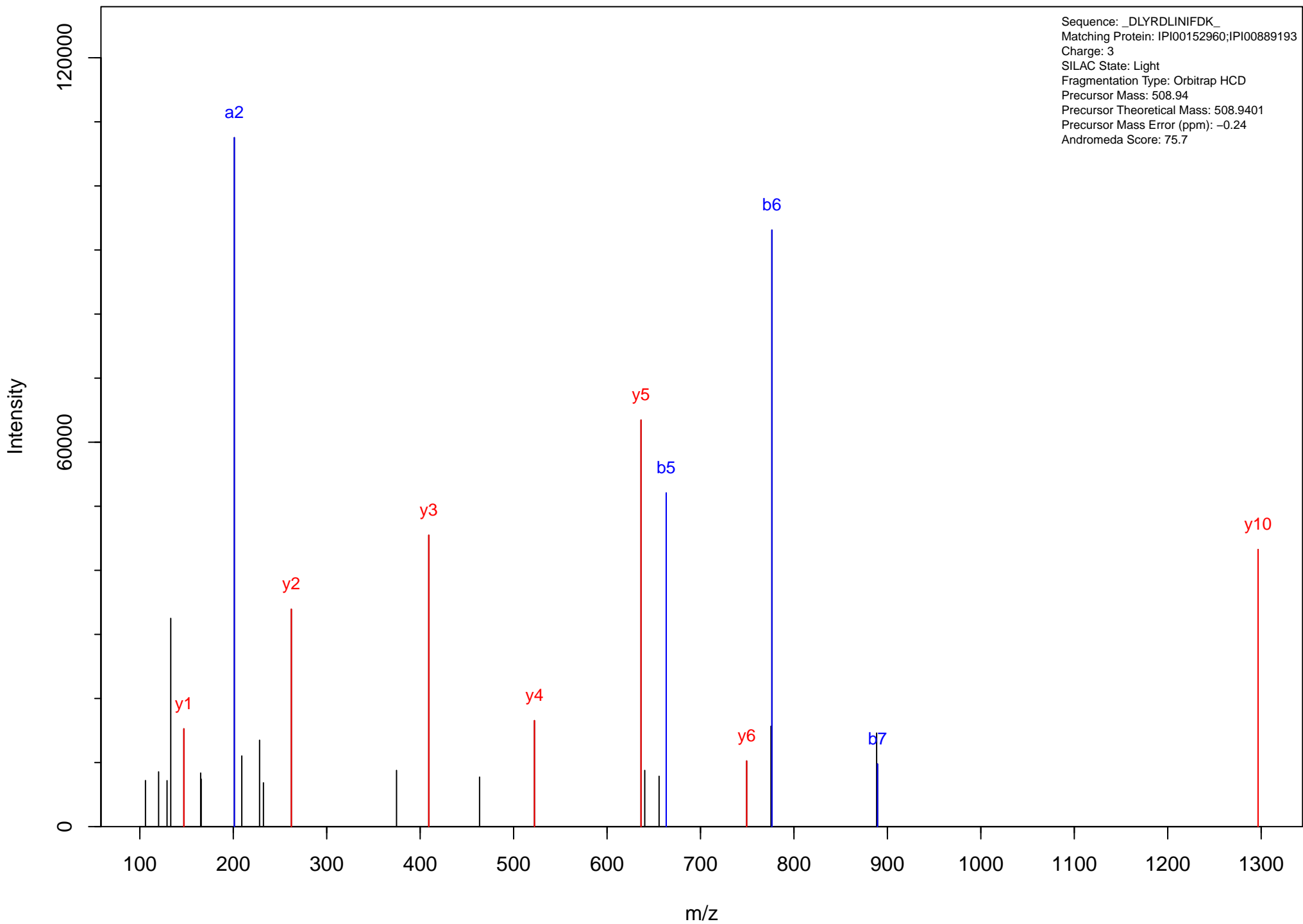


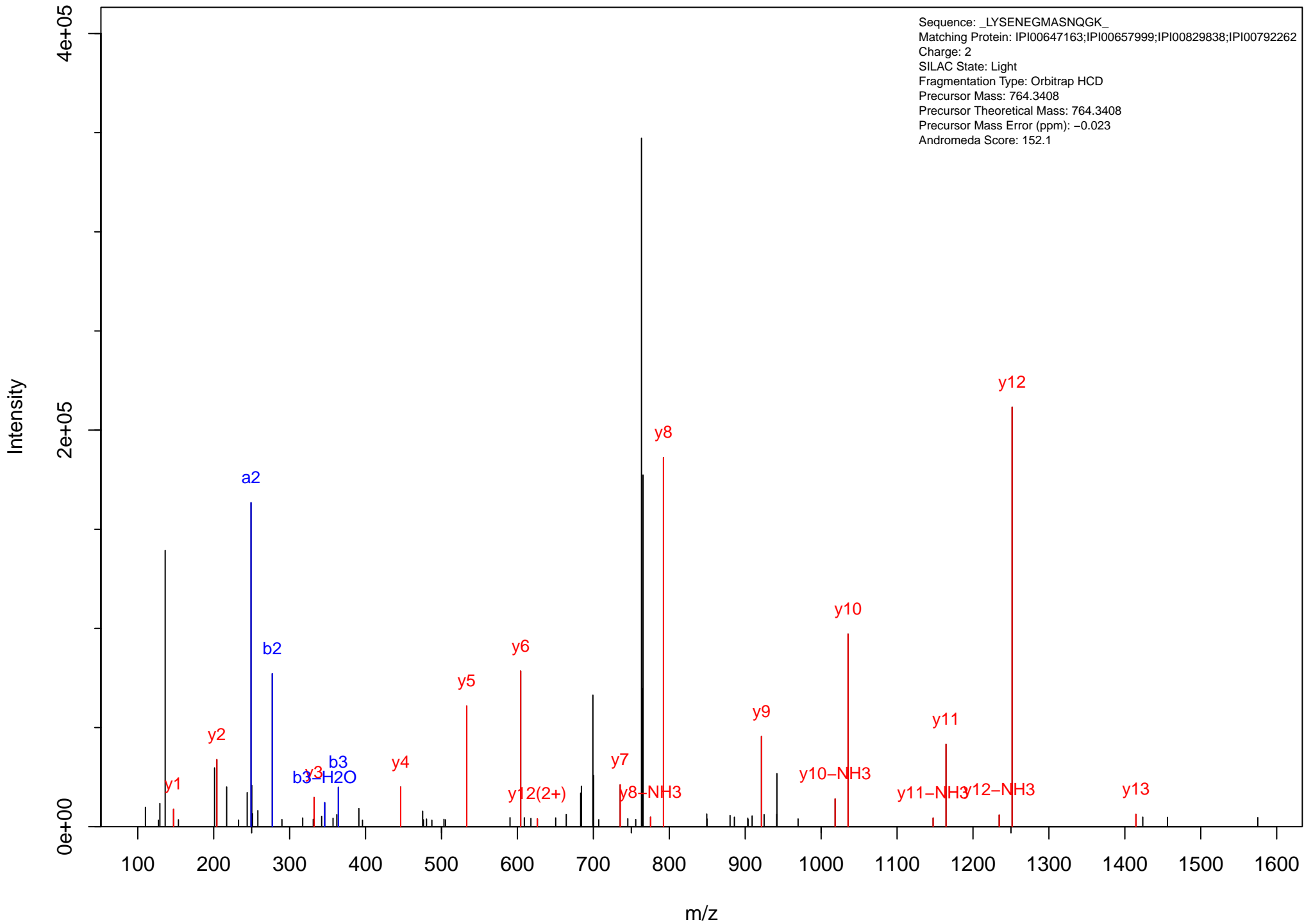


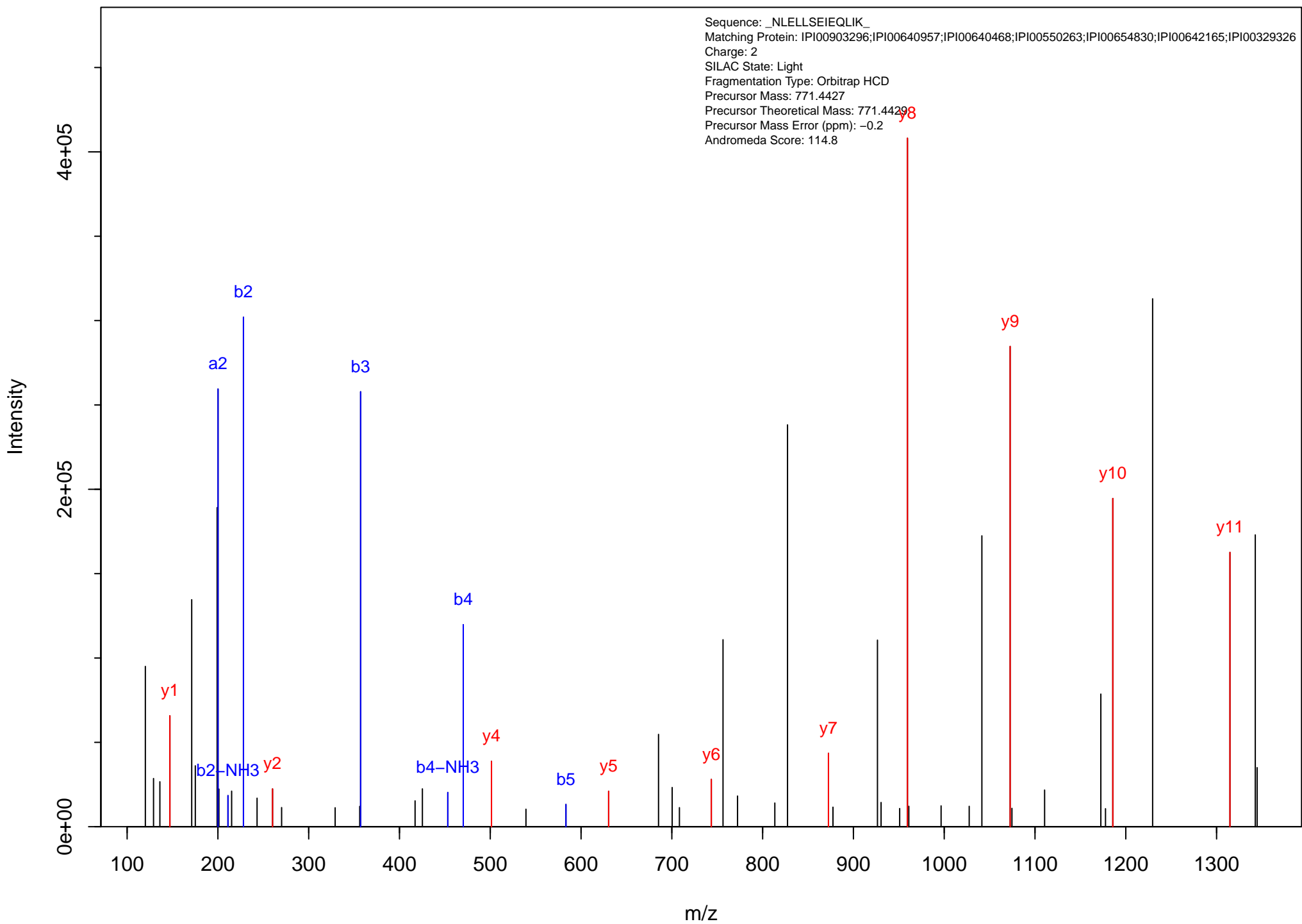


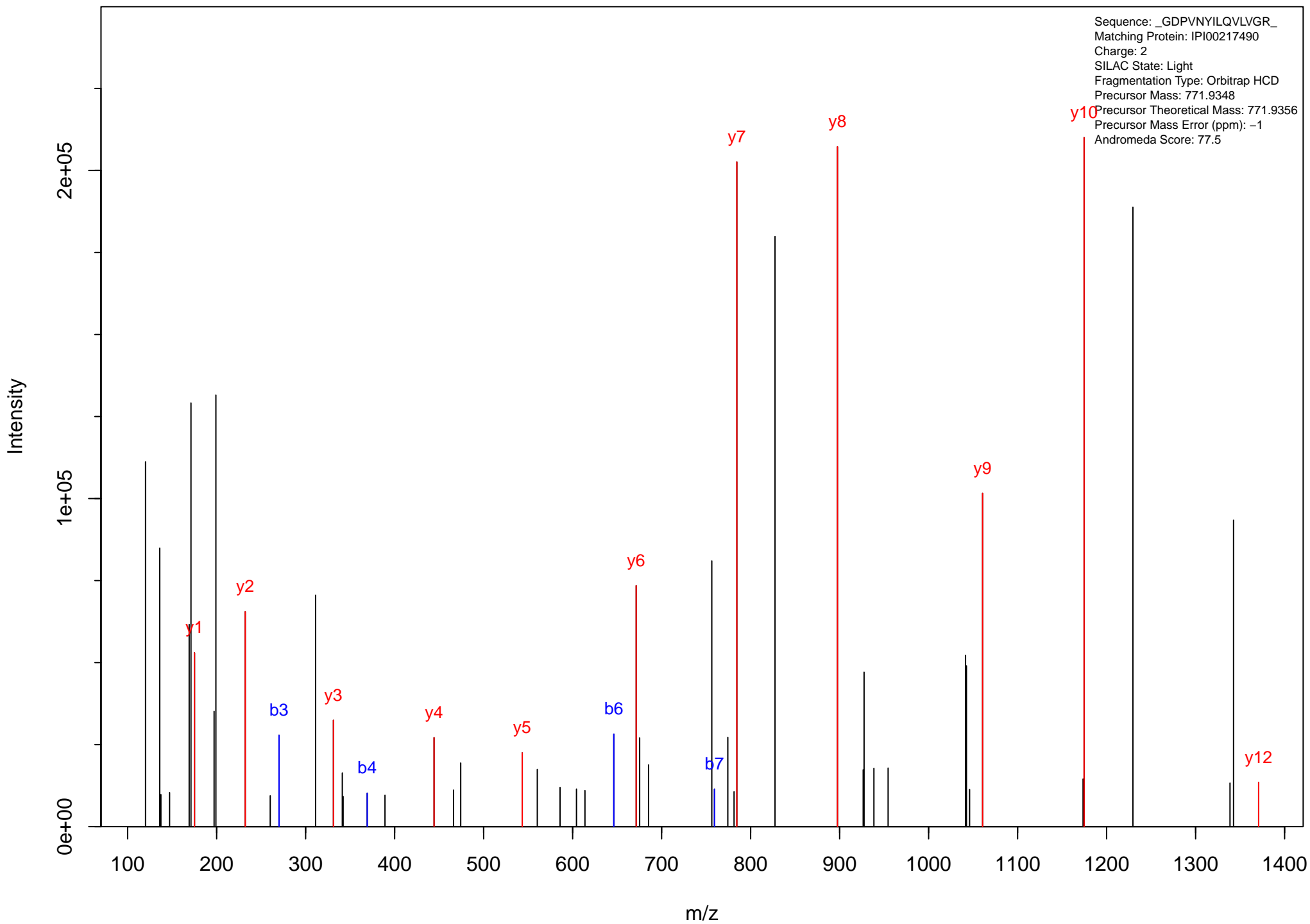


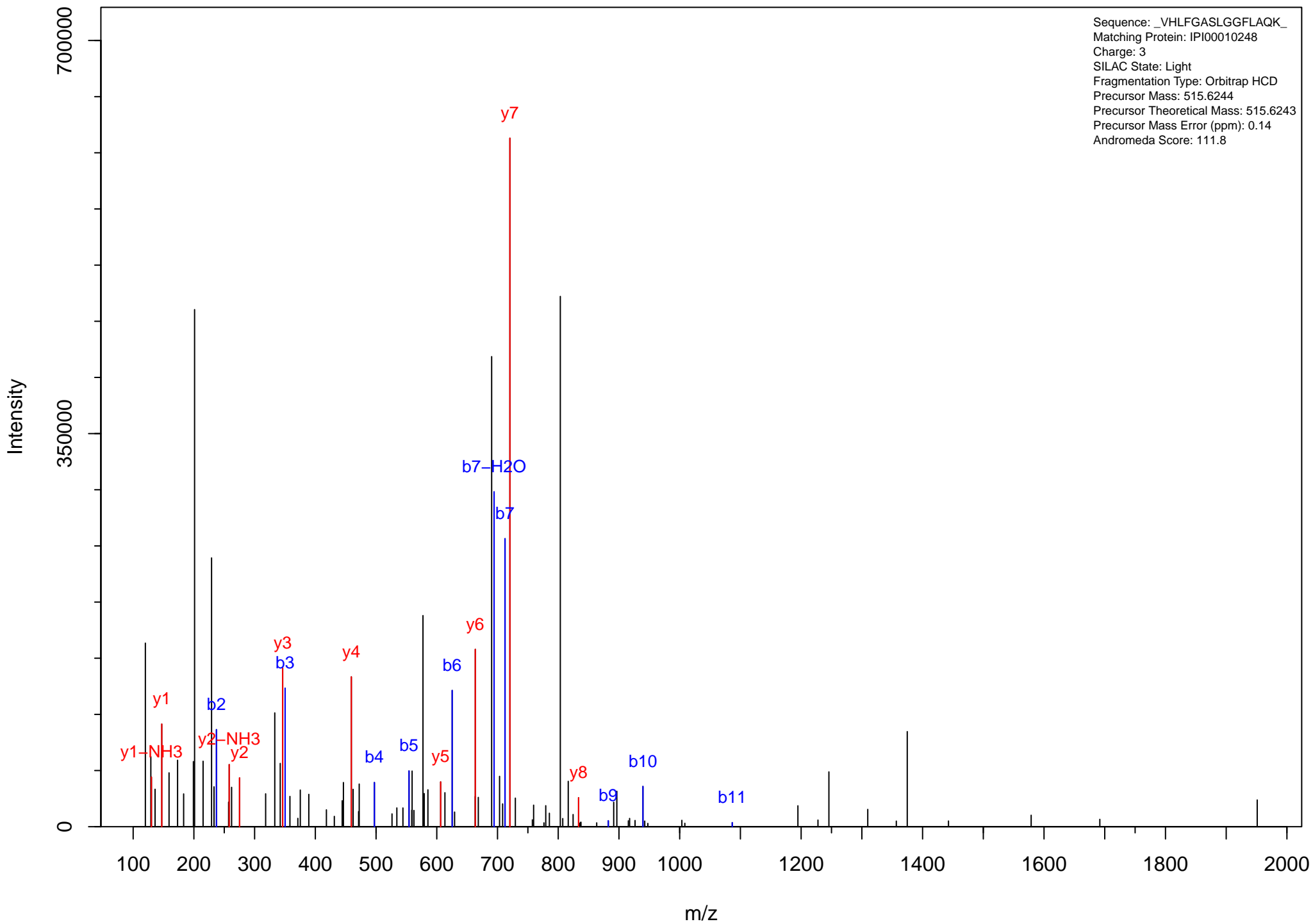


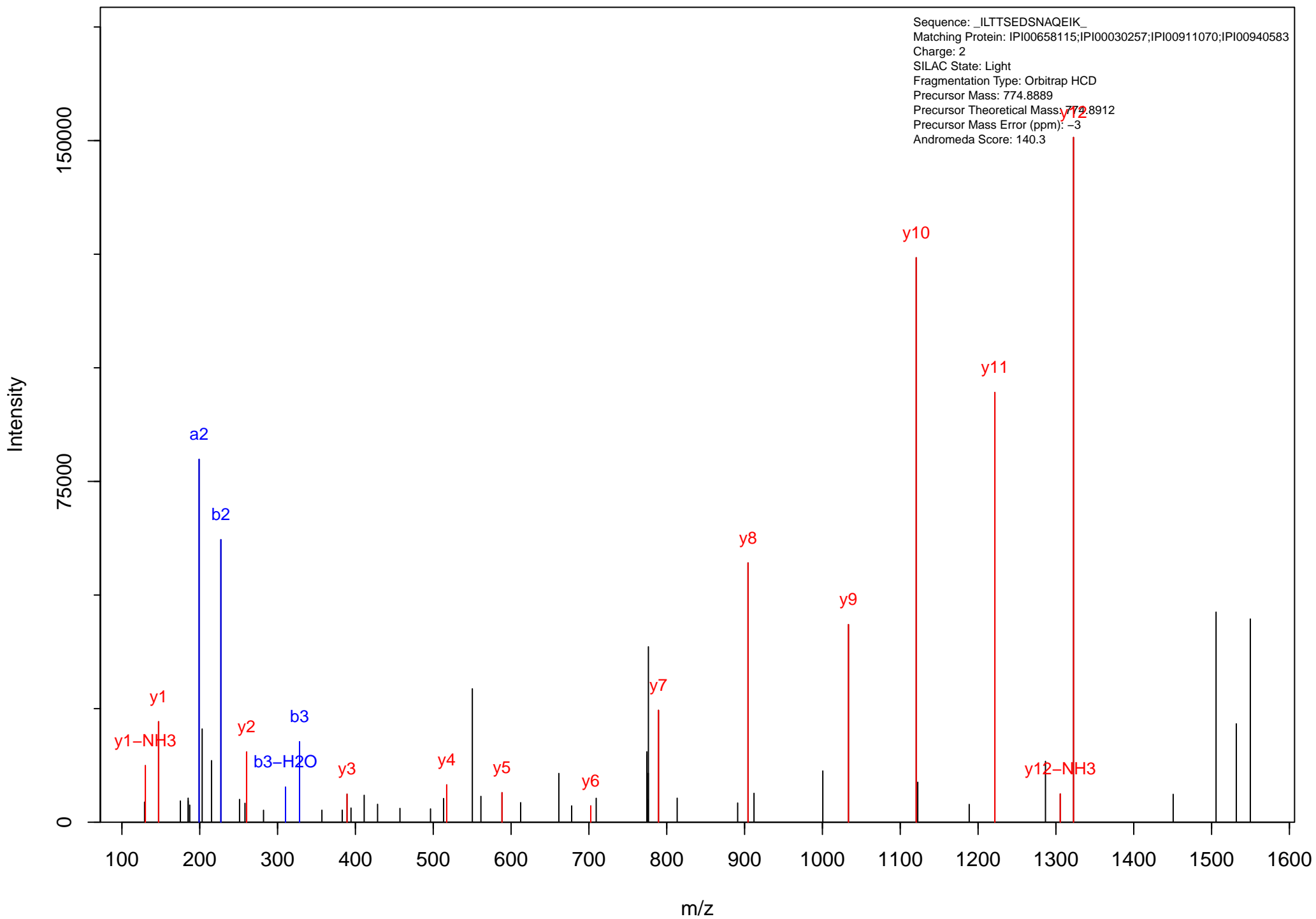


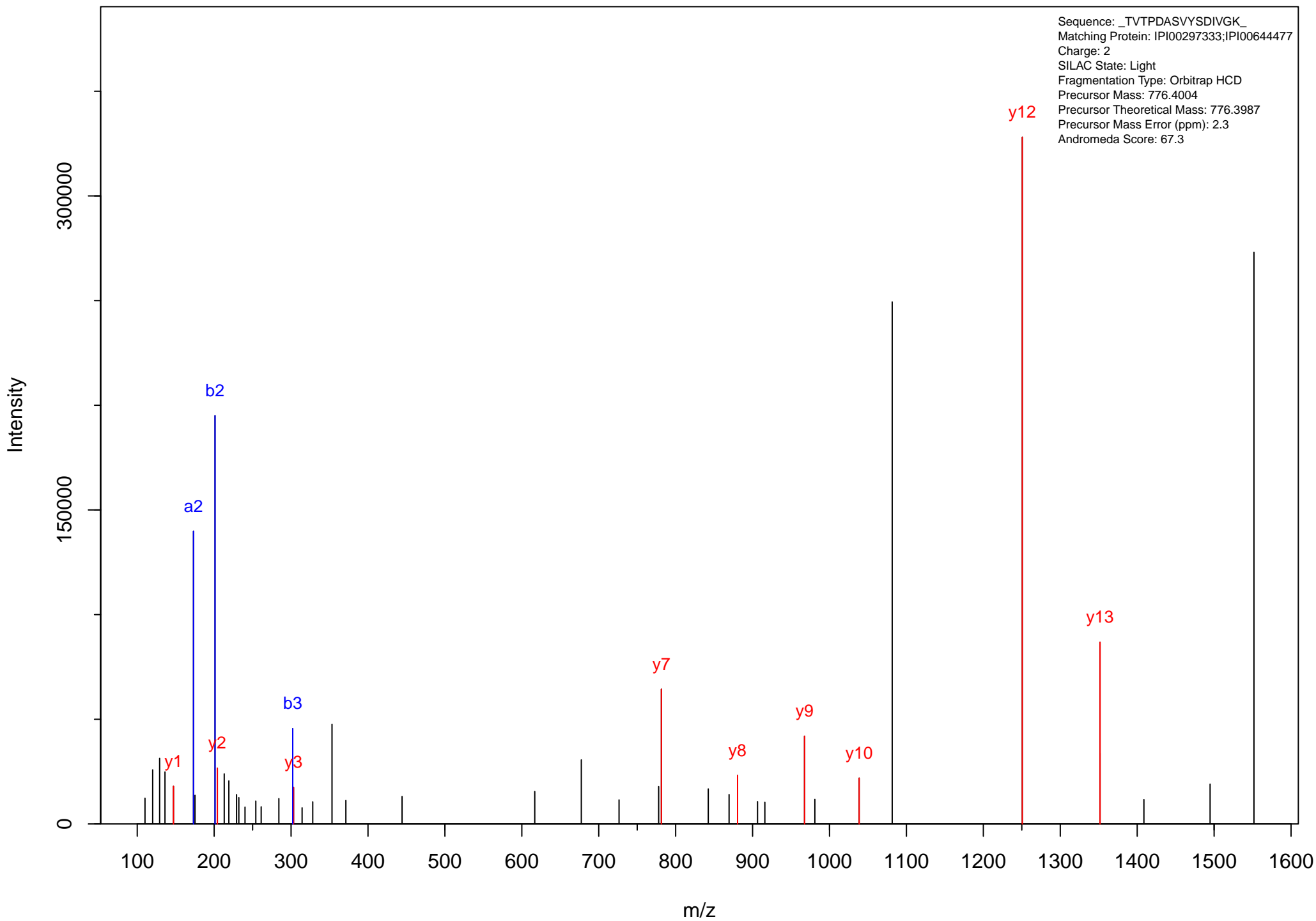


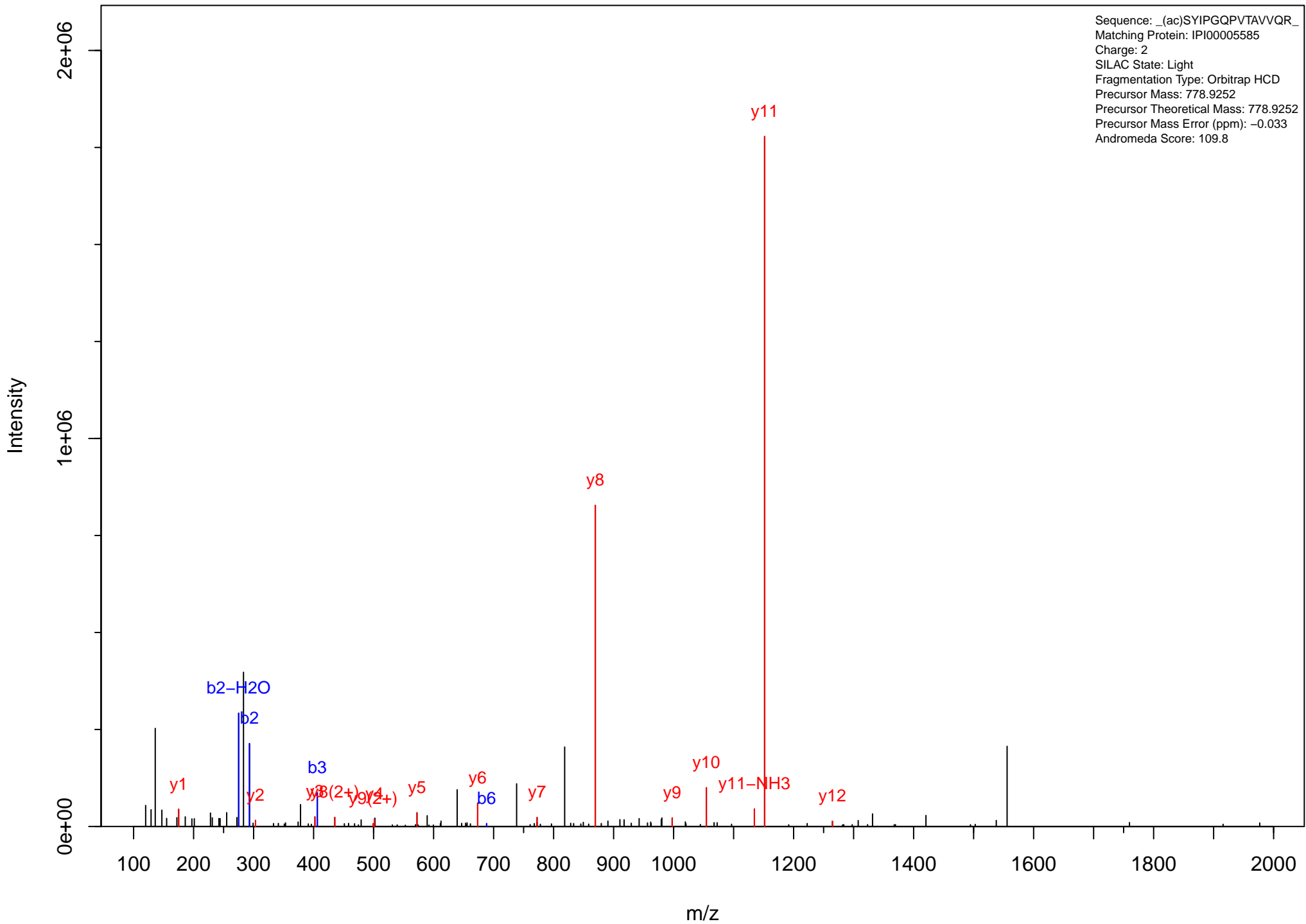


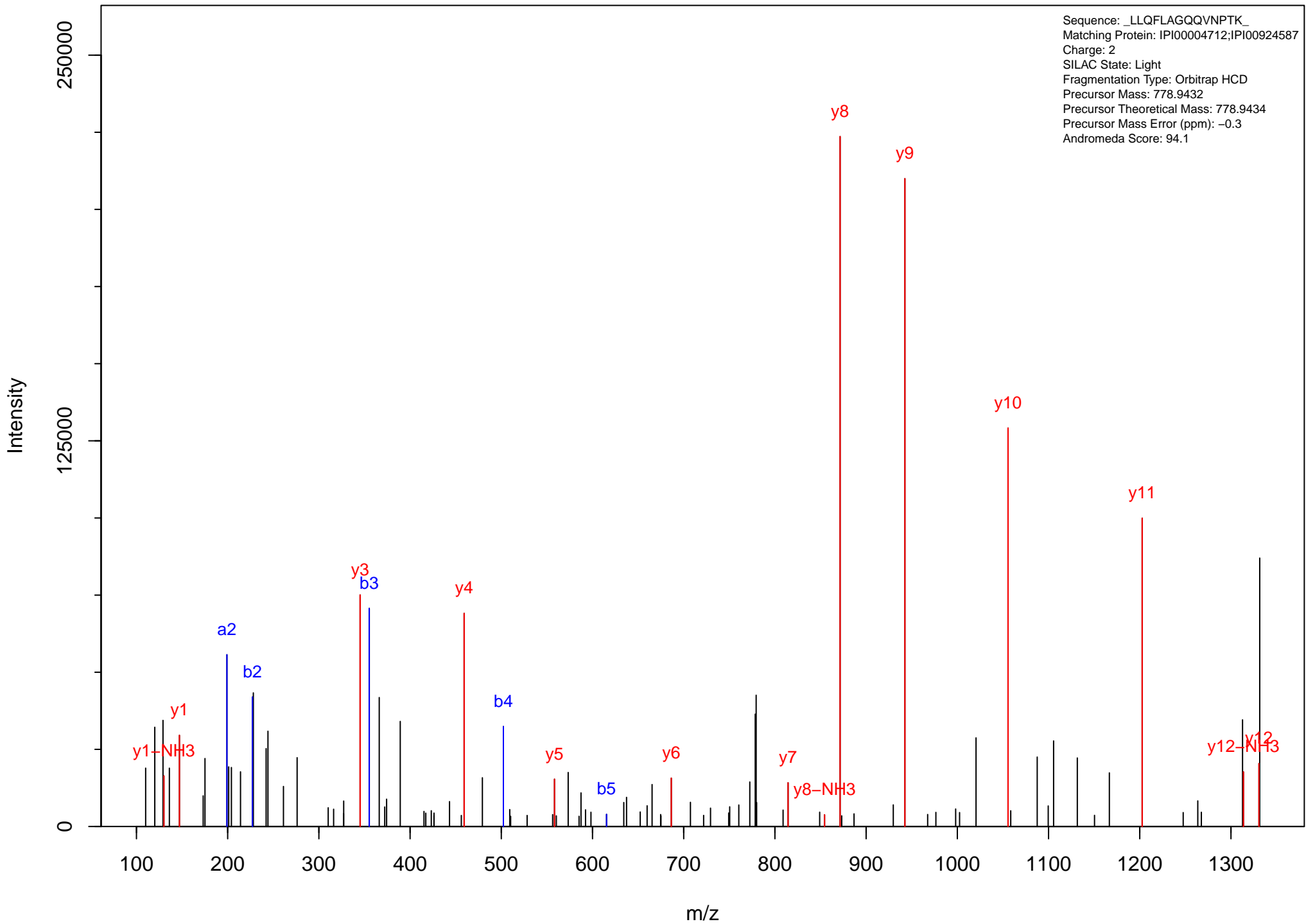


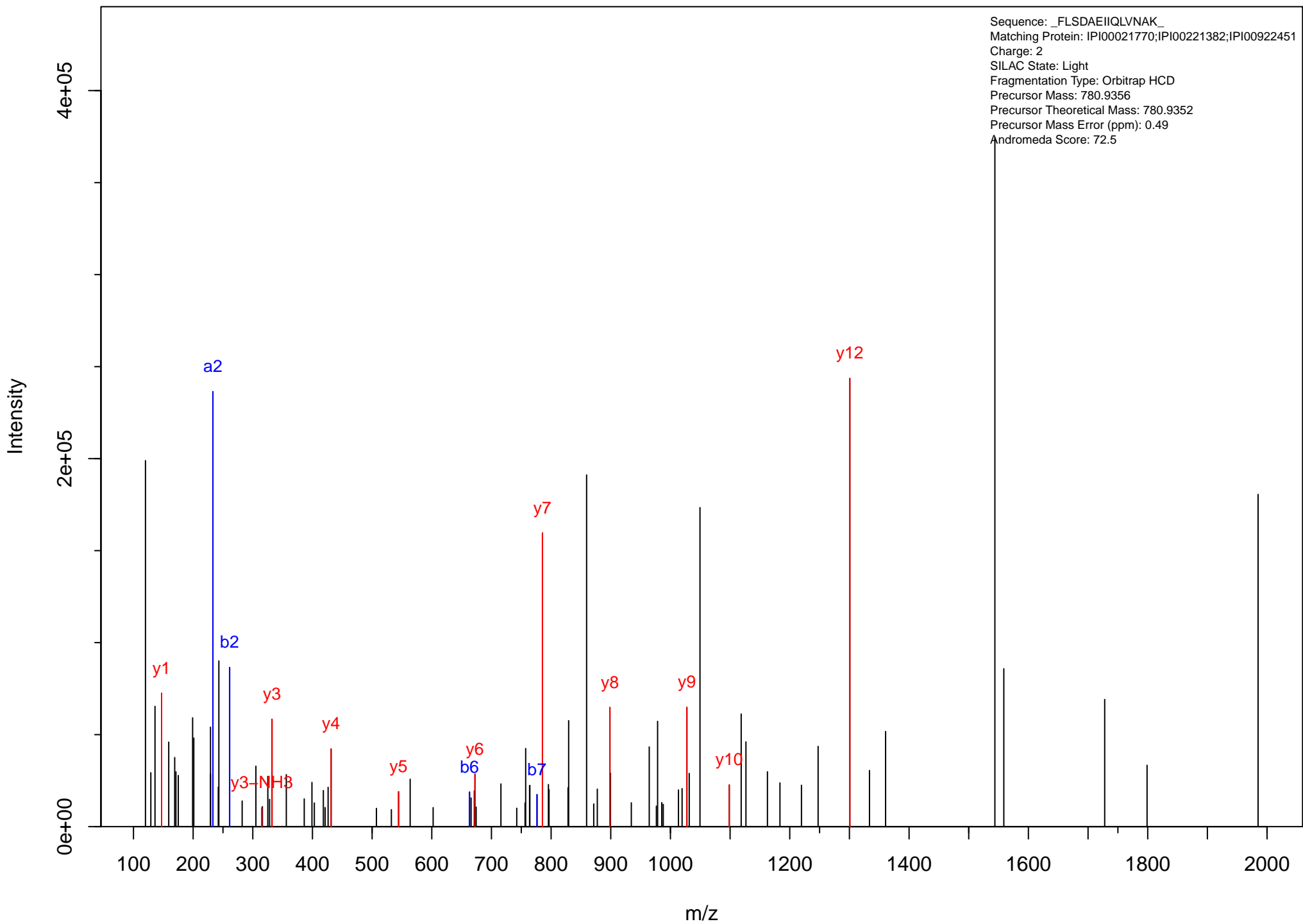


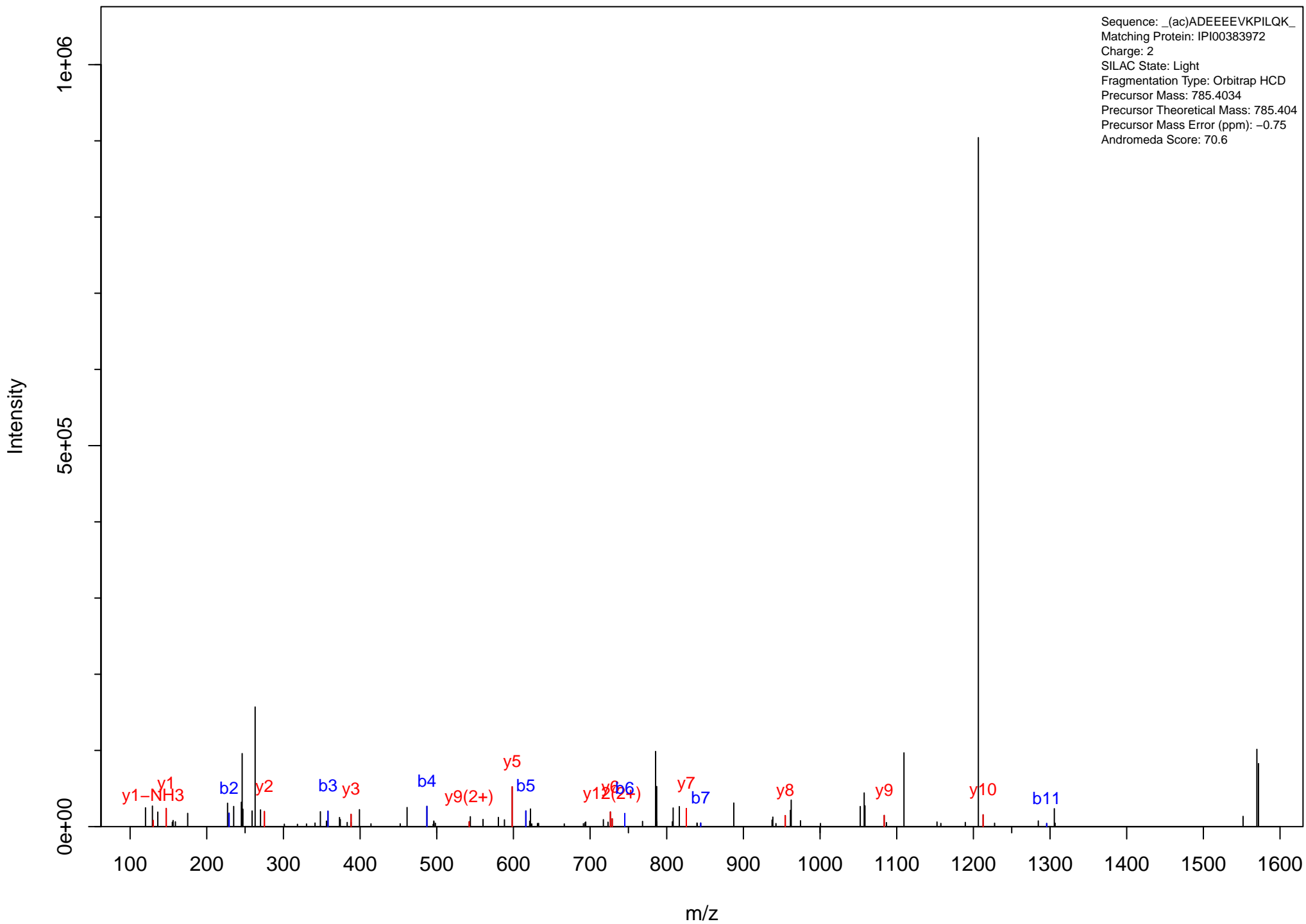


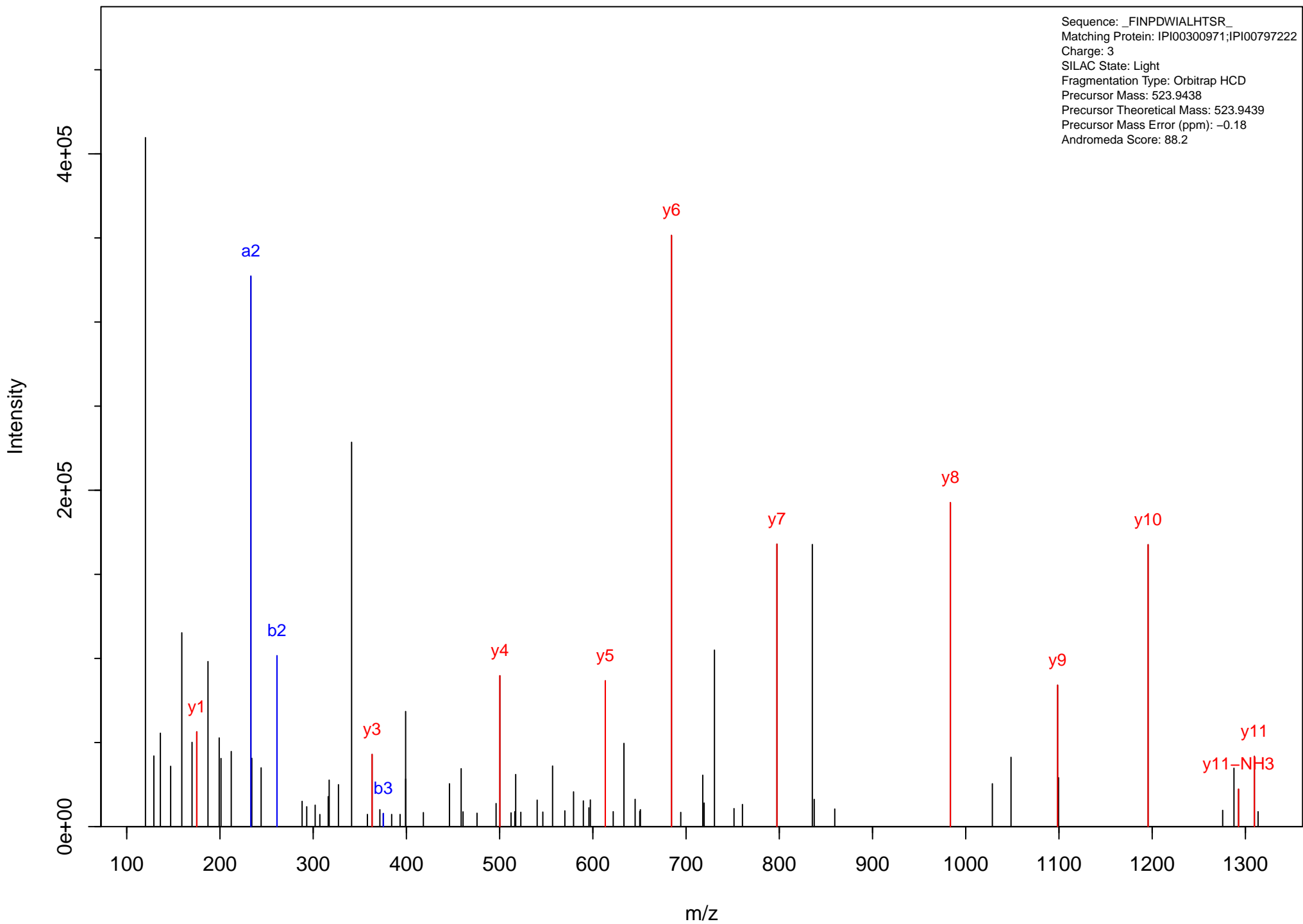


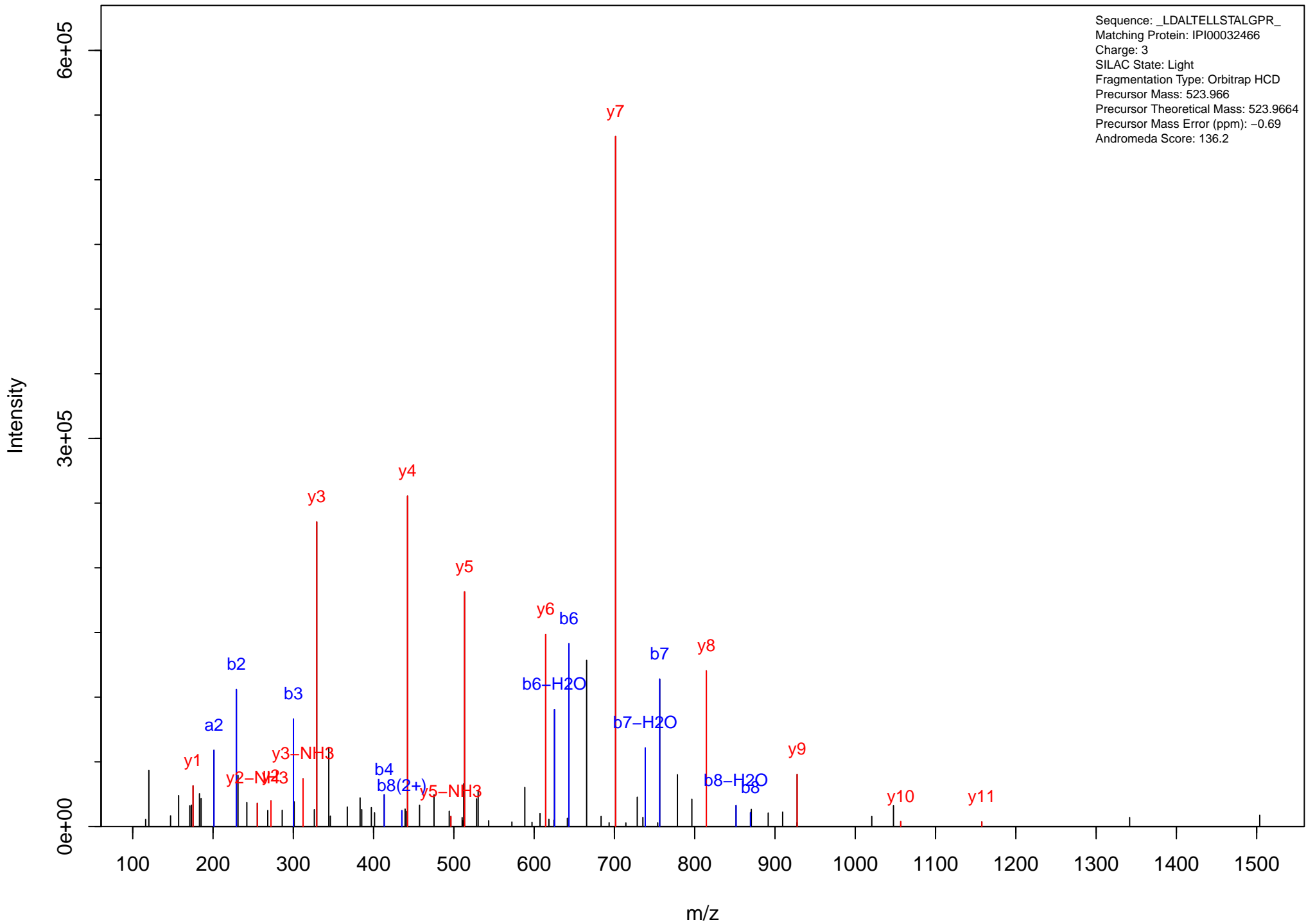


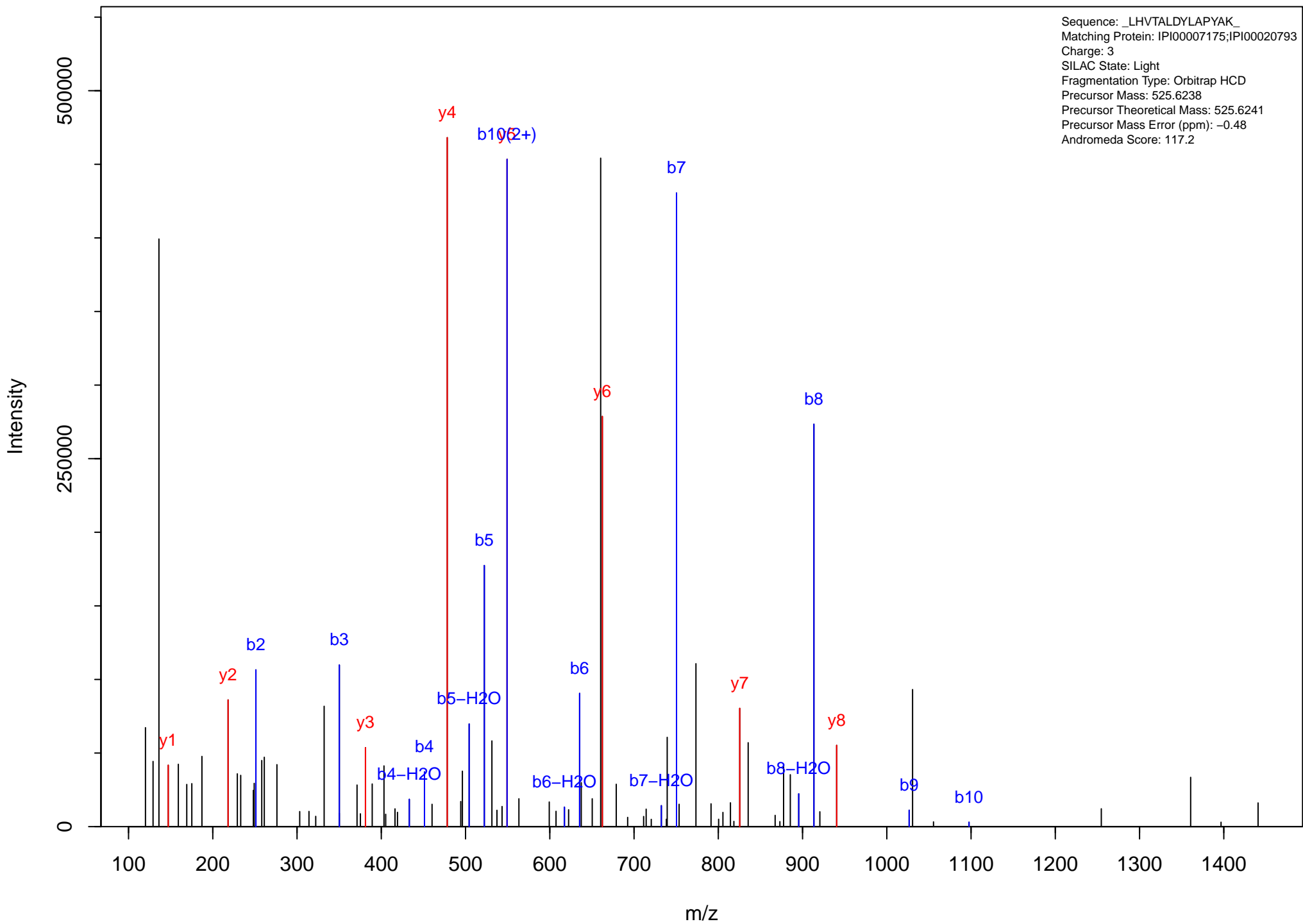


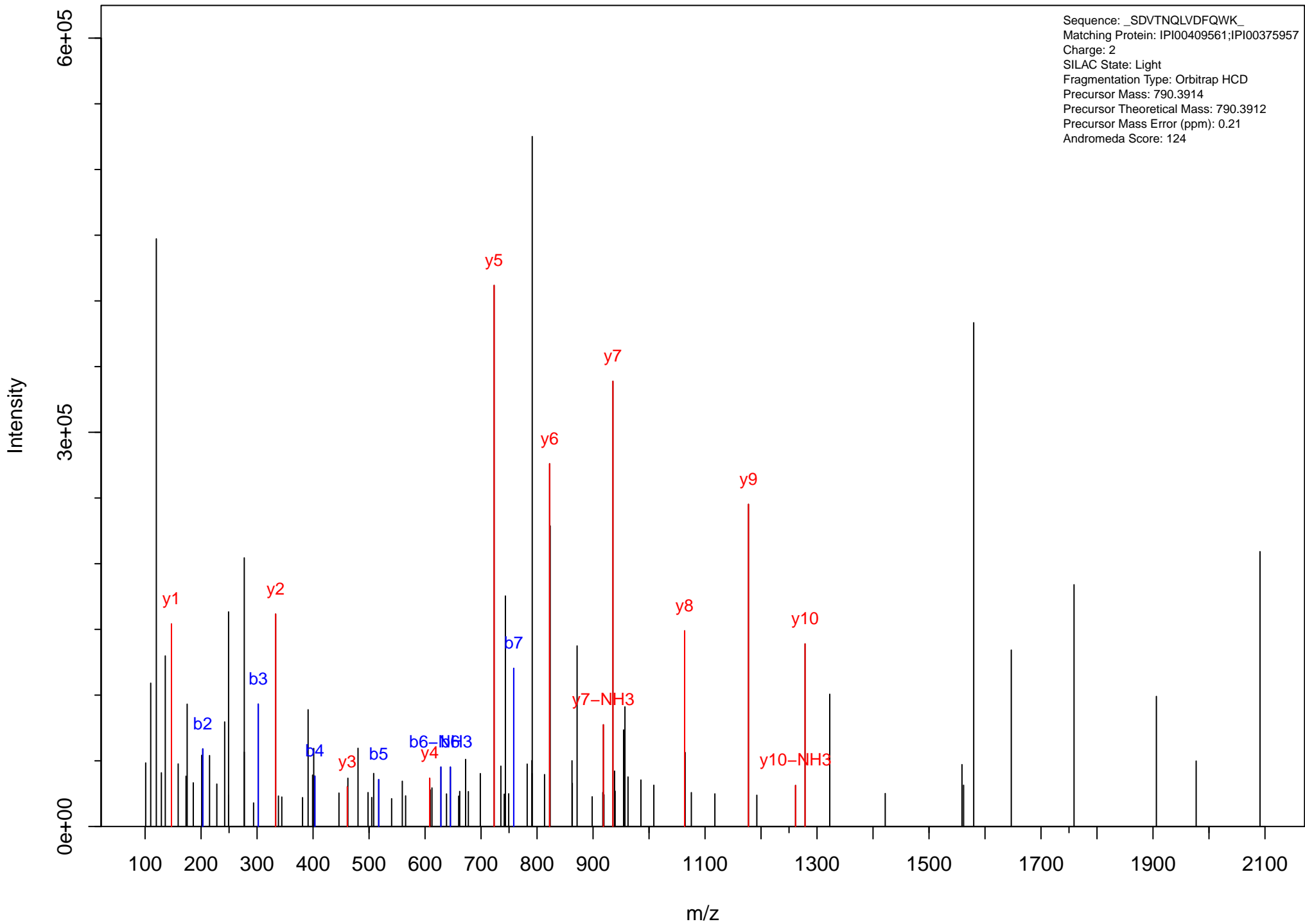


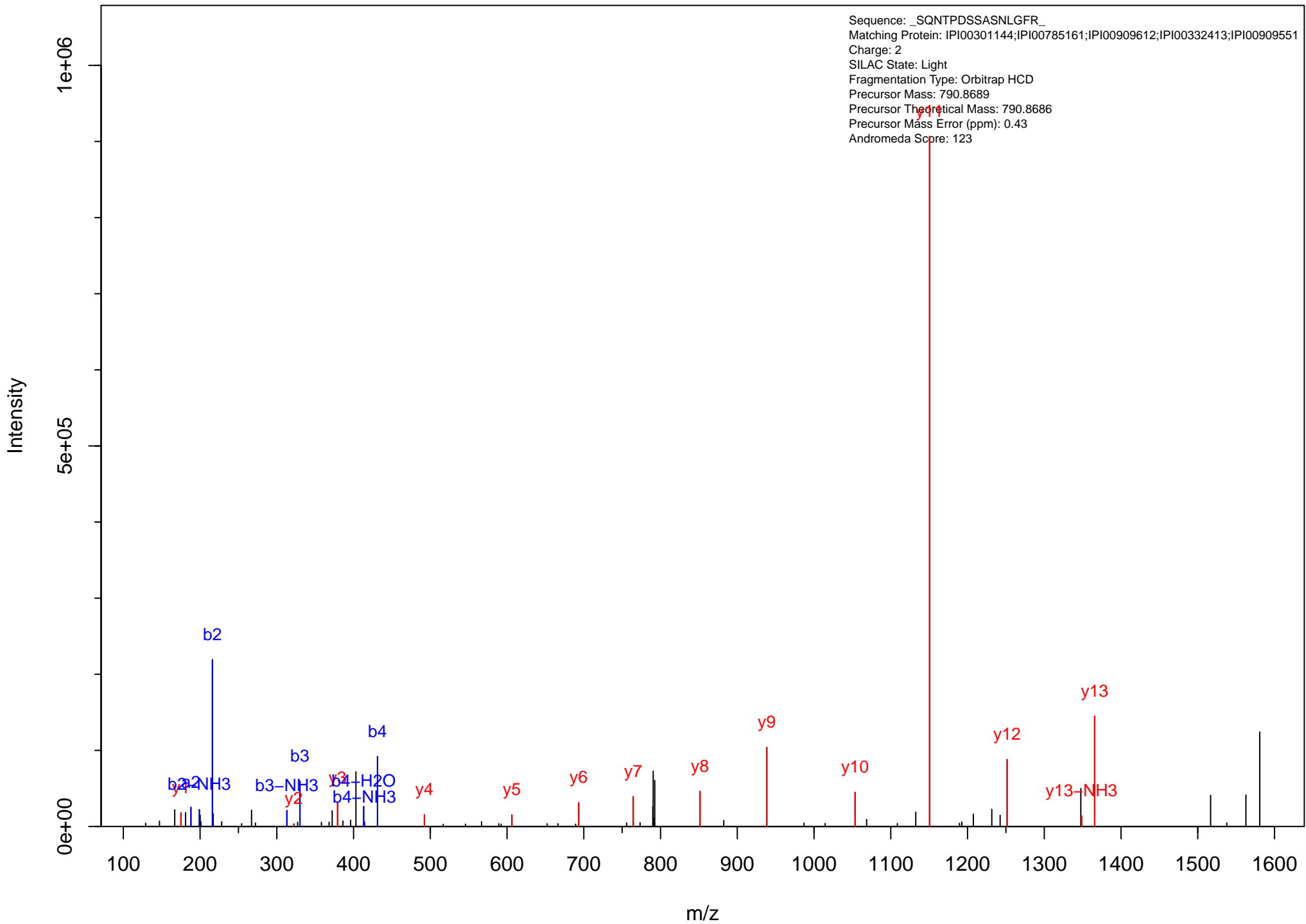


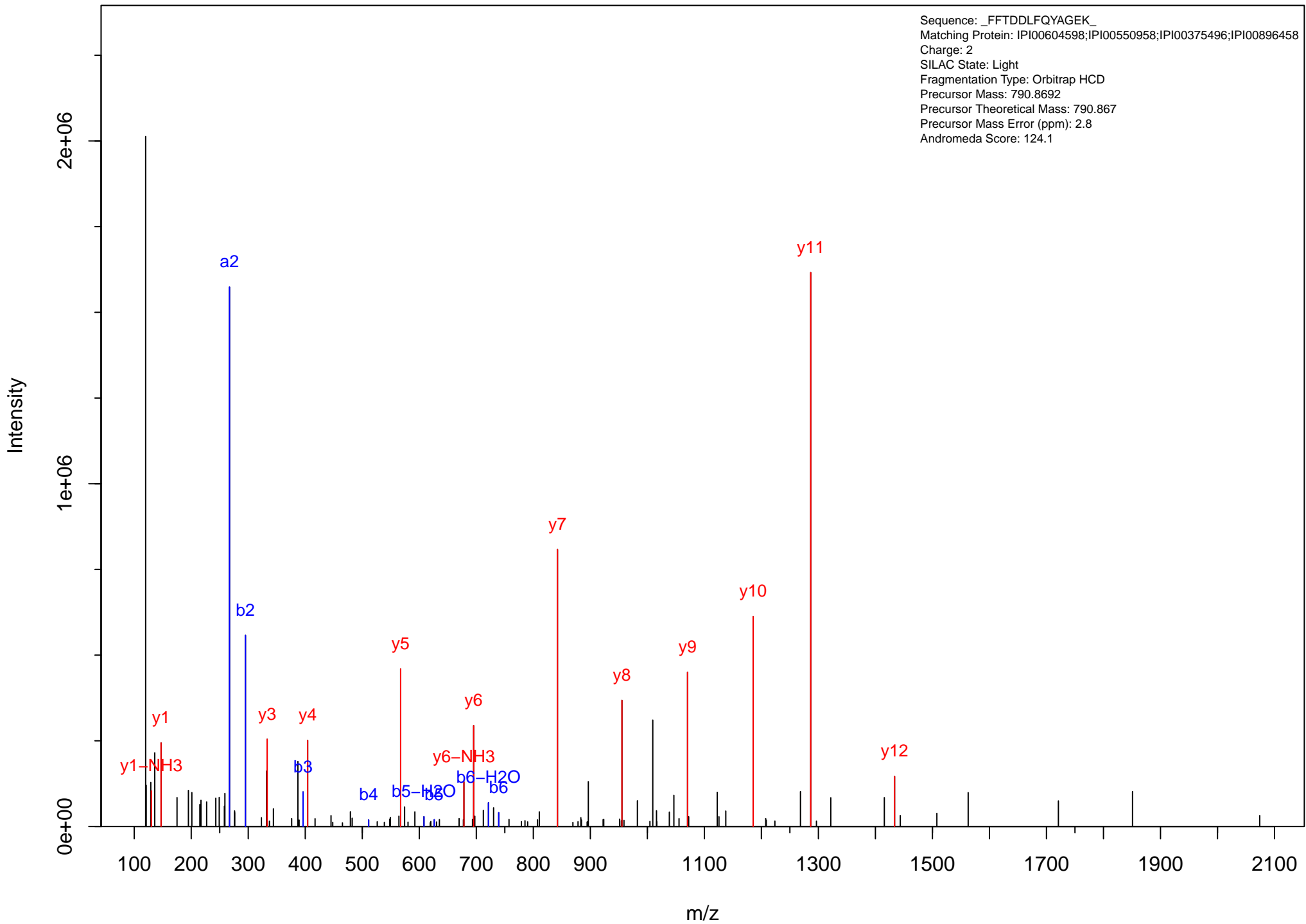


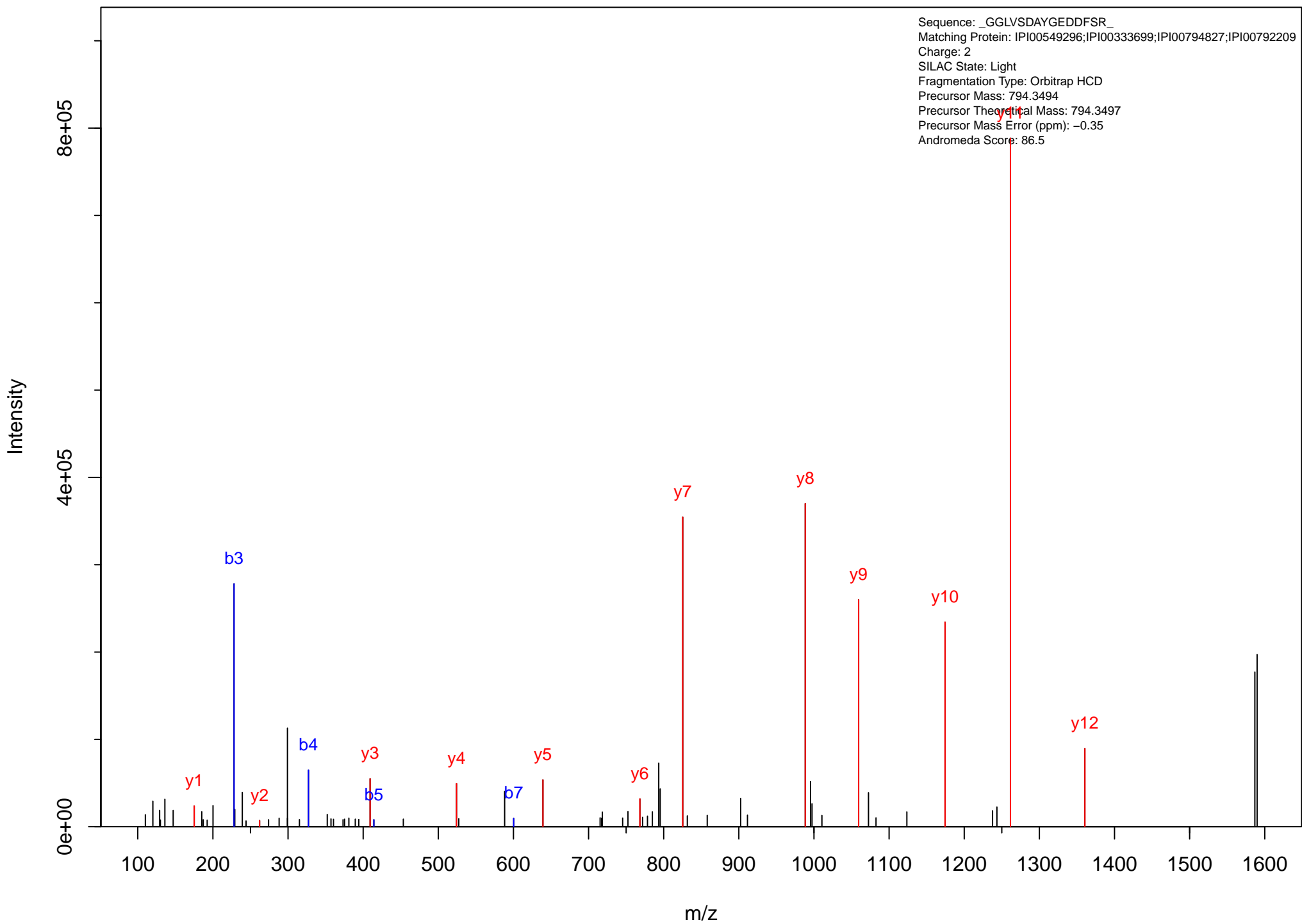


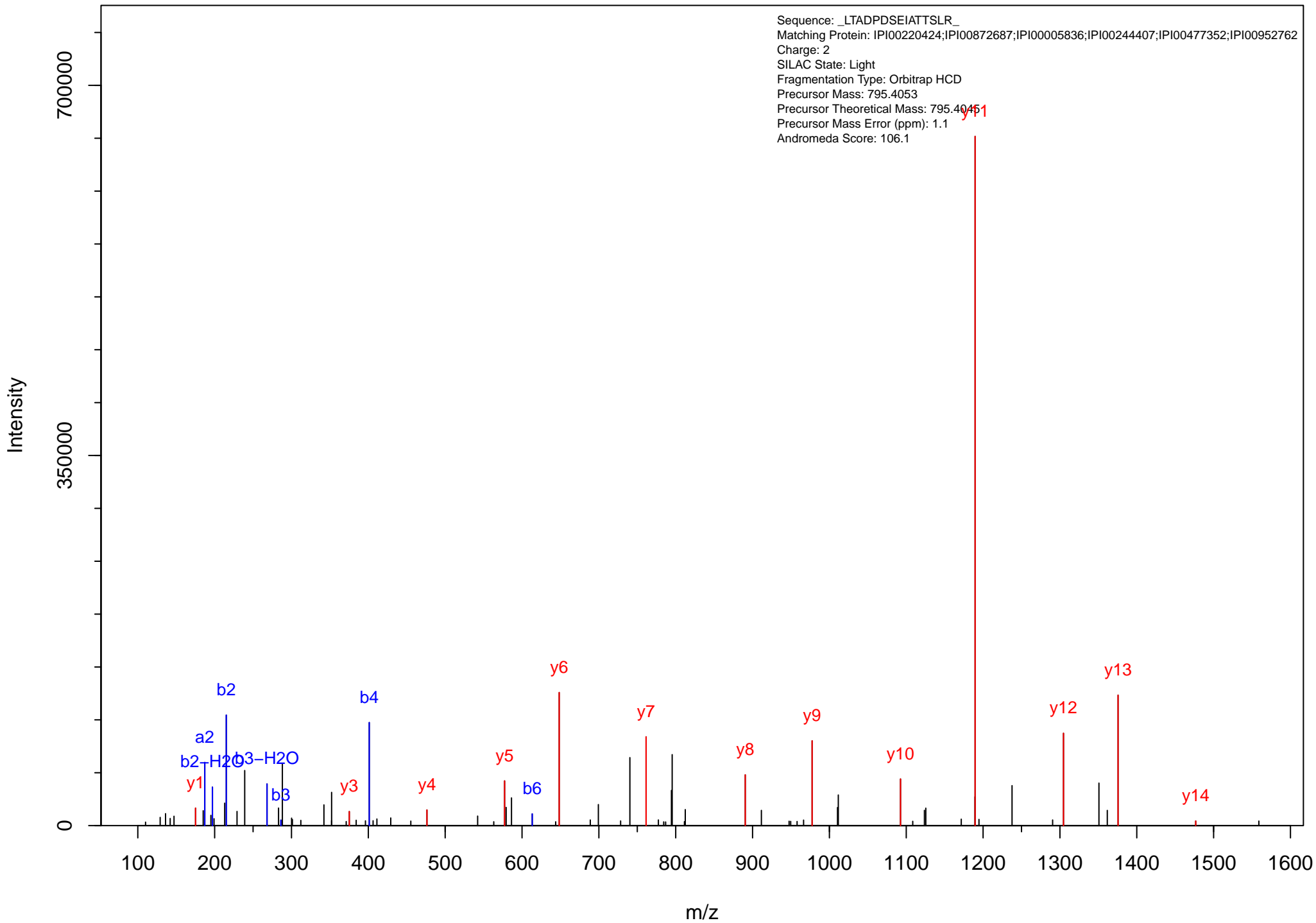


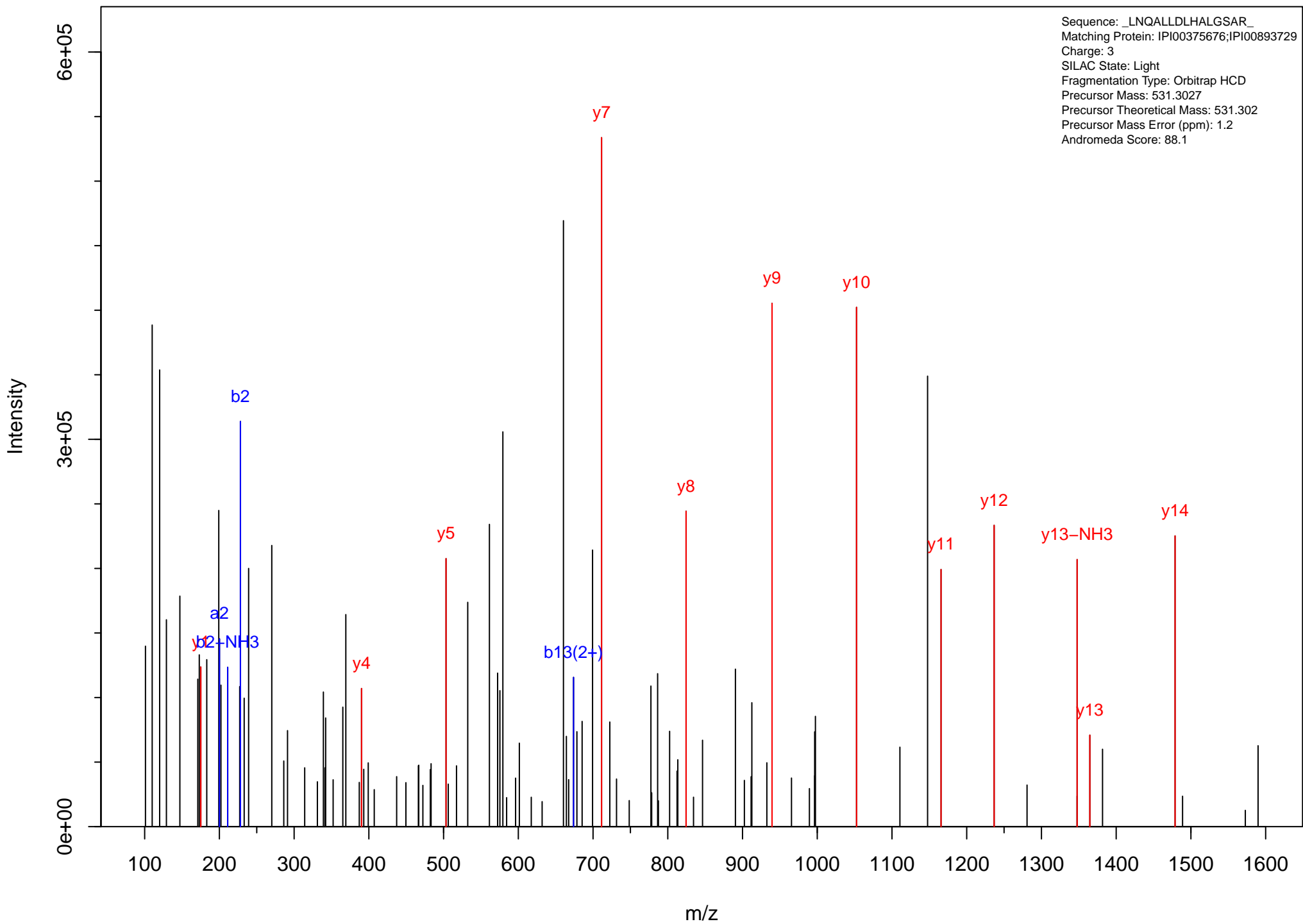


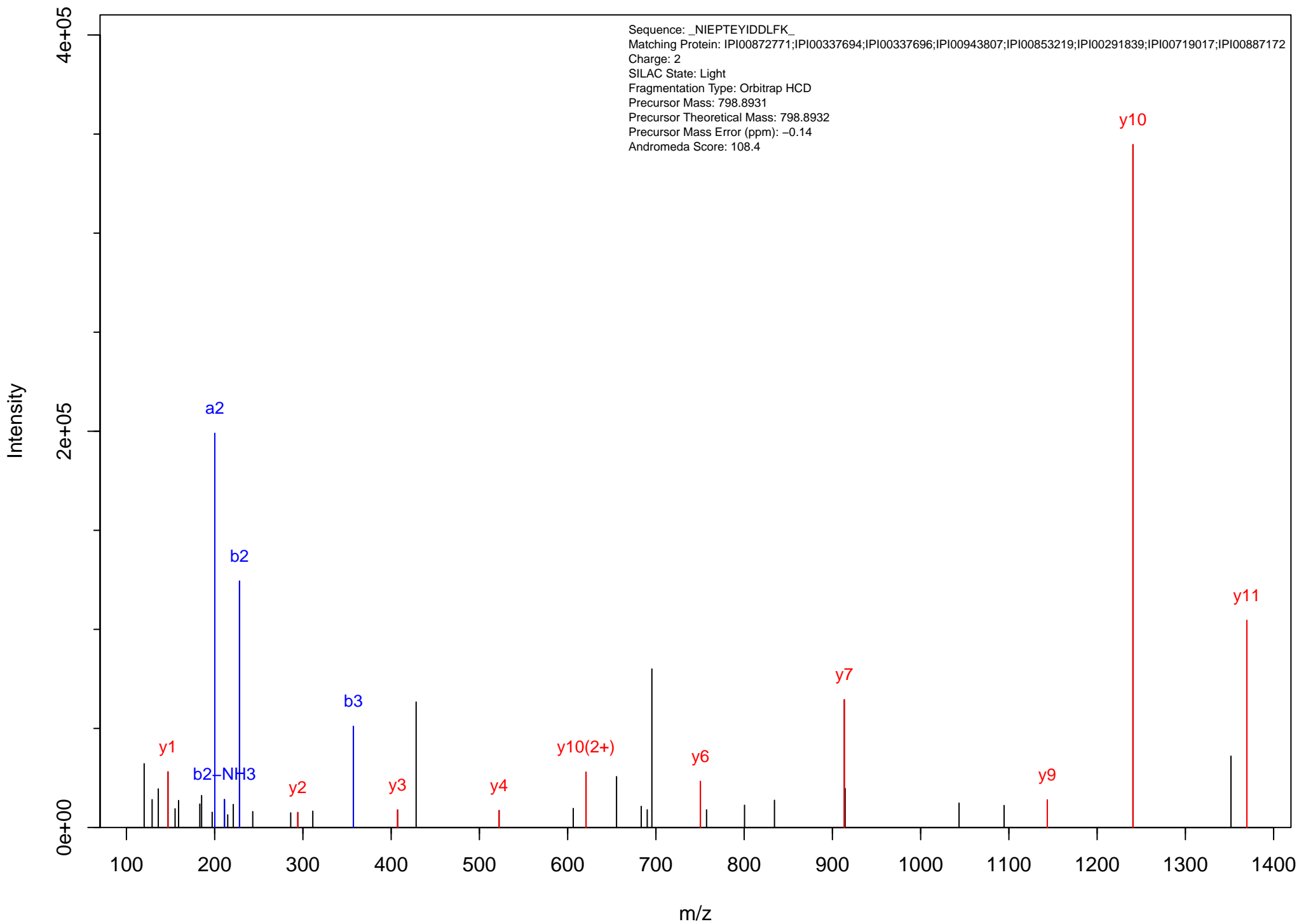


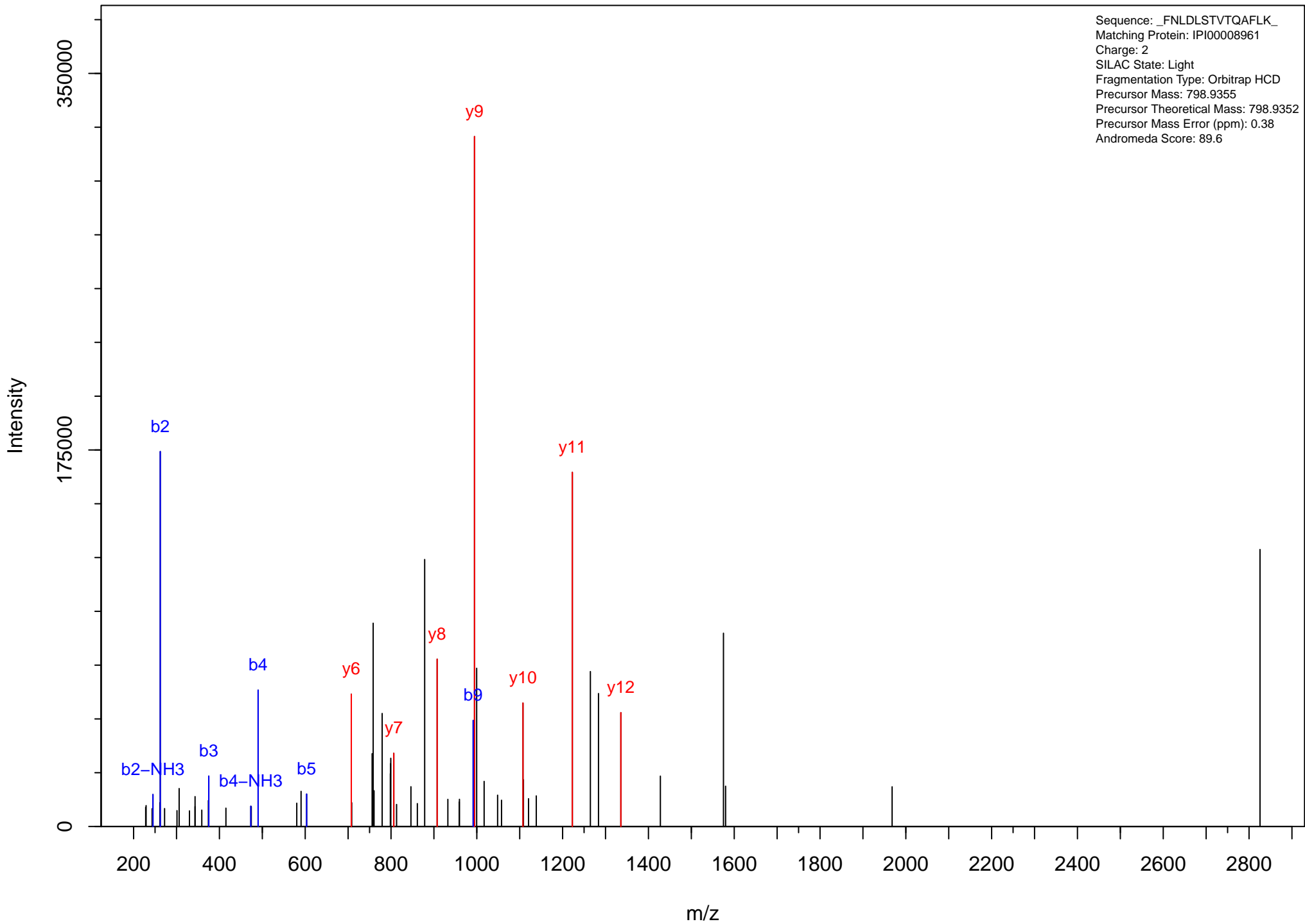












Intensity

