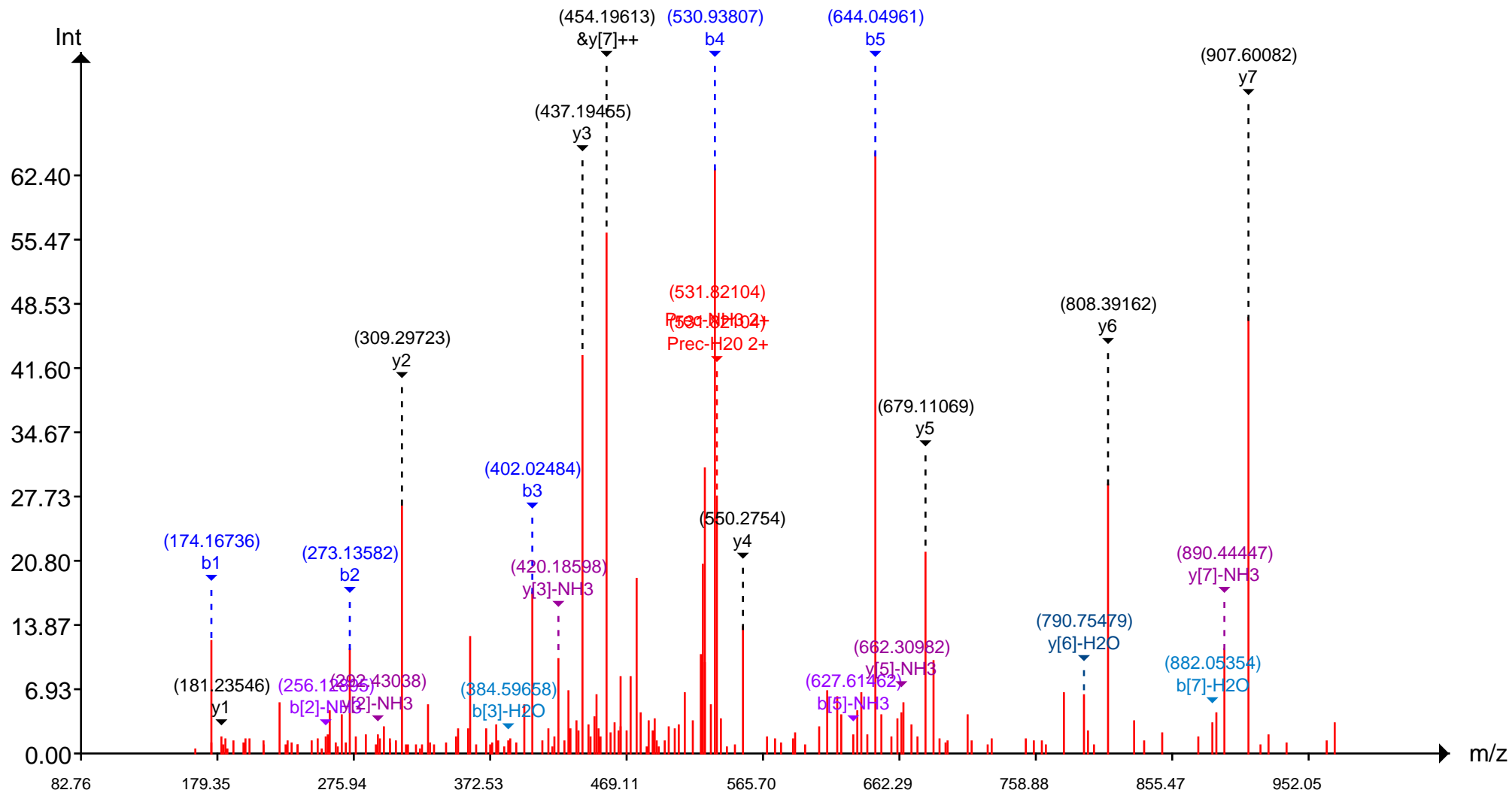


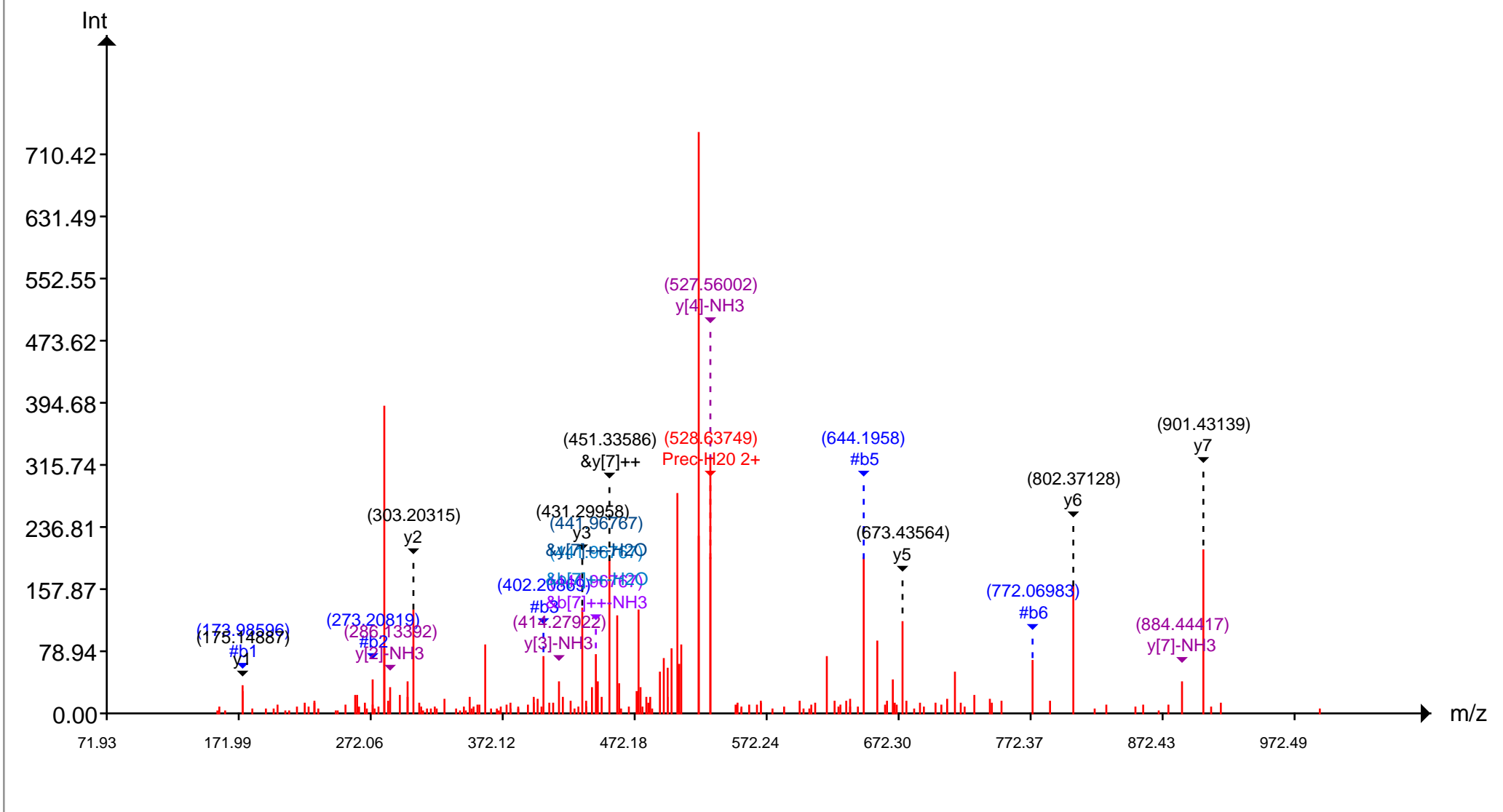
Info

| | | | |
|--------------------------|--------------------------|---------------------------|----------|
| Modified Sequence | AcD3-SQTGVGELTTQNTR-COOH | Ion score | 88.57 |
| Protein | Q96A65 | Confidence level | 99.00% |
| Precursor m/z | 768.88928 | Identity Threshold | 33.72 |
| Charge | 2+ | Mass error | 0.002106 |



Info

| | | | |
|--------------------------|-------------------------|---------------------------|----------|
| Modified Sequence | AcD3-QVEELQQR<C13>-COOH | Ion score | 35.84 |
| Protein | Q9BRP8 | Confidence level | 99.00% |
| Precursor m/z | 540.79439 | Identity Threshold | 34.45 |
| Charge | 2+ | Mass error | -3.57E-4 |



Info

Modified Sequence AcD3-QVEELQQR-COOH

Protein Q9BRP8

Precursor m/z 537.78474

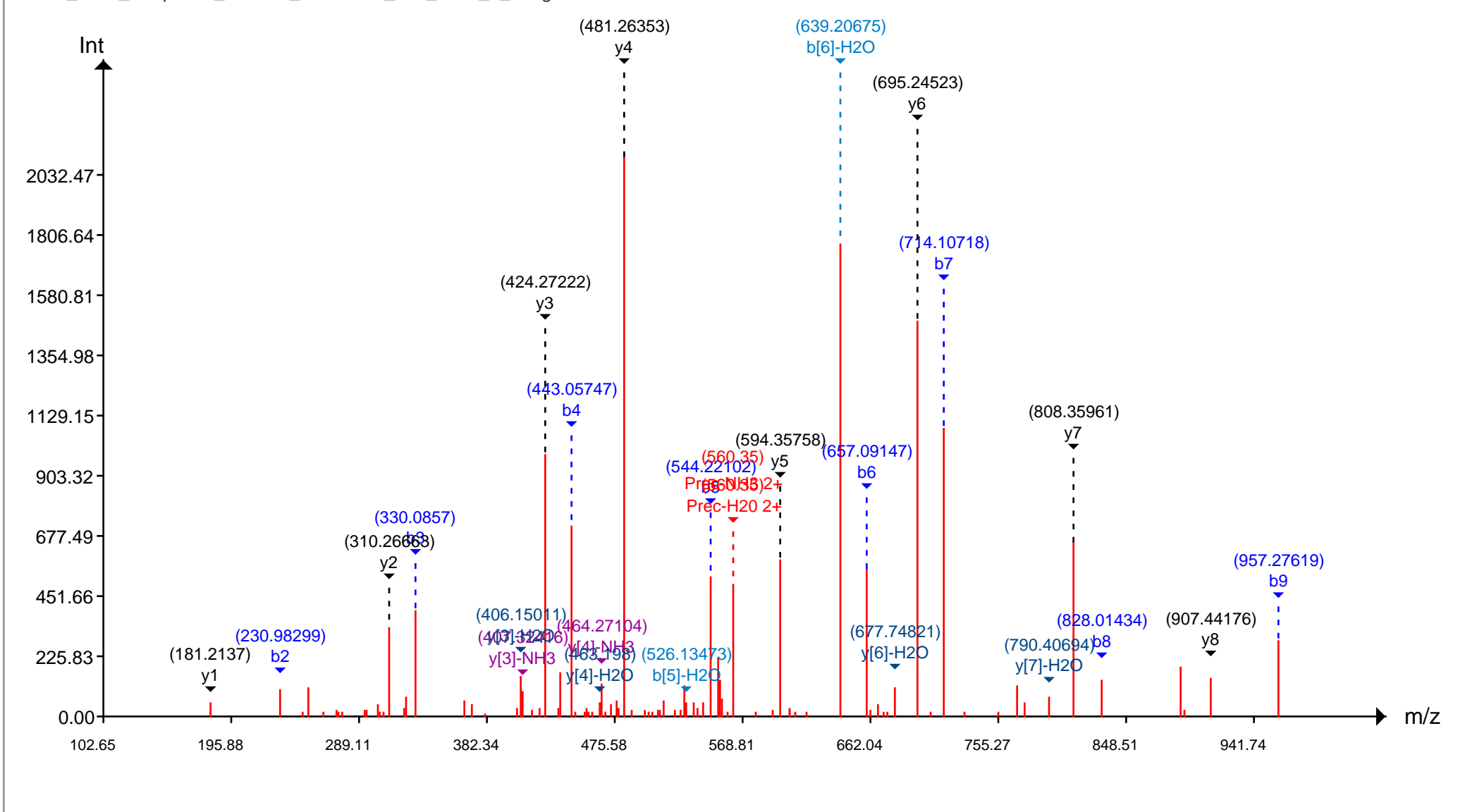
Charge 2+

Ion score 41.86

Confidence level 99.00%

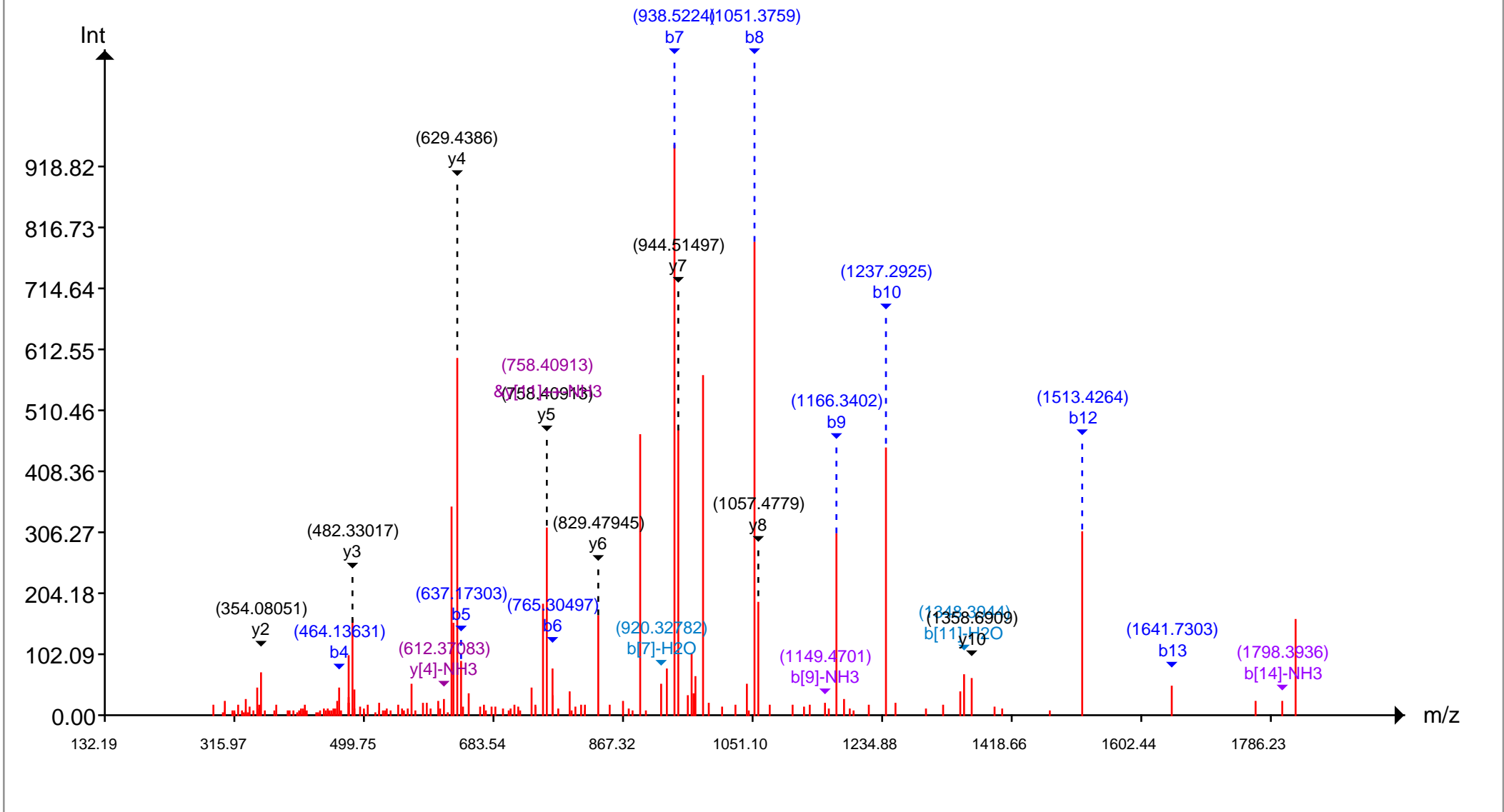
Identity Threshold 34.75

Mass error 4.85E-4



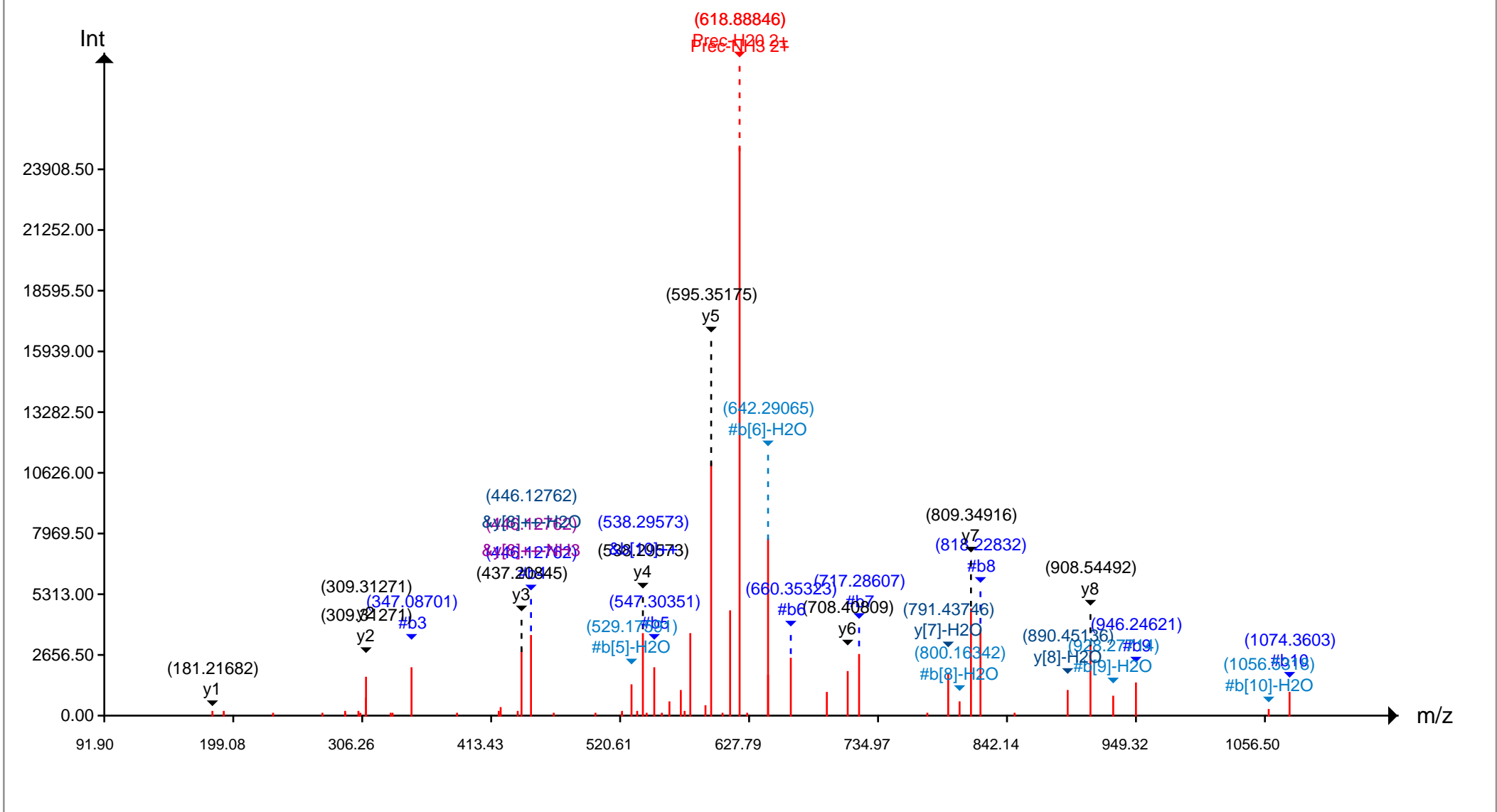
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 70.92 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32312 | Identity Threshold | 48.32 |
| Charge | 2+ | Mass error | -7.58E-4 |



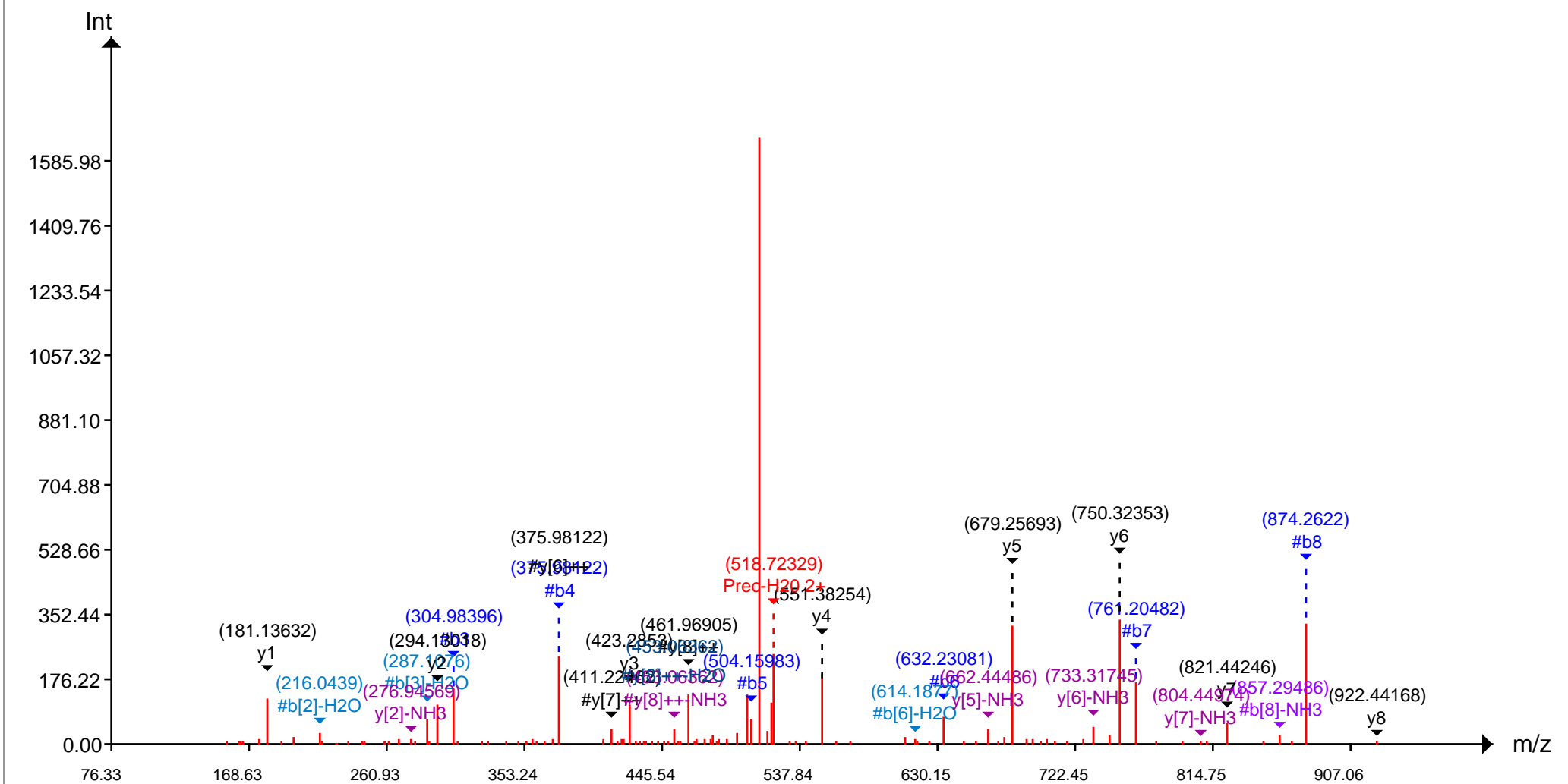
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-AM<Mox*>AEK<AcD3K*>QK<AcD3K*>LDAEFQK<AcD3K*>R<C13>-COOH | Ion score | 49.0 |
| Protein | Q9H875 | Confidence level | 99.00% |
| Precursor m/z | 998.03925 | Identity Threshold | 48.55 |
| Charge | 2+ | Mass error | 0.001189 |



Info

| | | | |
|--------------------------|------------------------------------|---------------------------|-----------|
| Modified Sequence | AcD3-AGK<AcD3K*>VTLGTQQR<C13>-COOH | Ion score | 78.94 |
| Protein | Q08AF3 | Confidence level | 99.00% |
| Precursor m/z | 627.87179 | Identity Threshold | 46.57 |
| Charge | 2+ | Mass error | -0.001578 |



Info

Modified Sequence AcD3-STAAQQELR<C13>-COOH

Ion score 72.4

Protein P35579

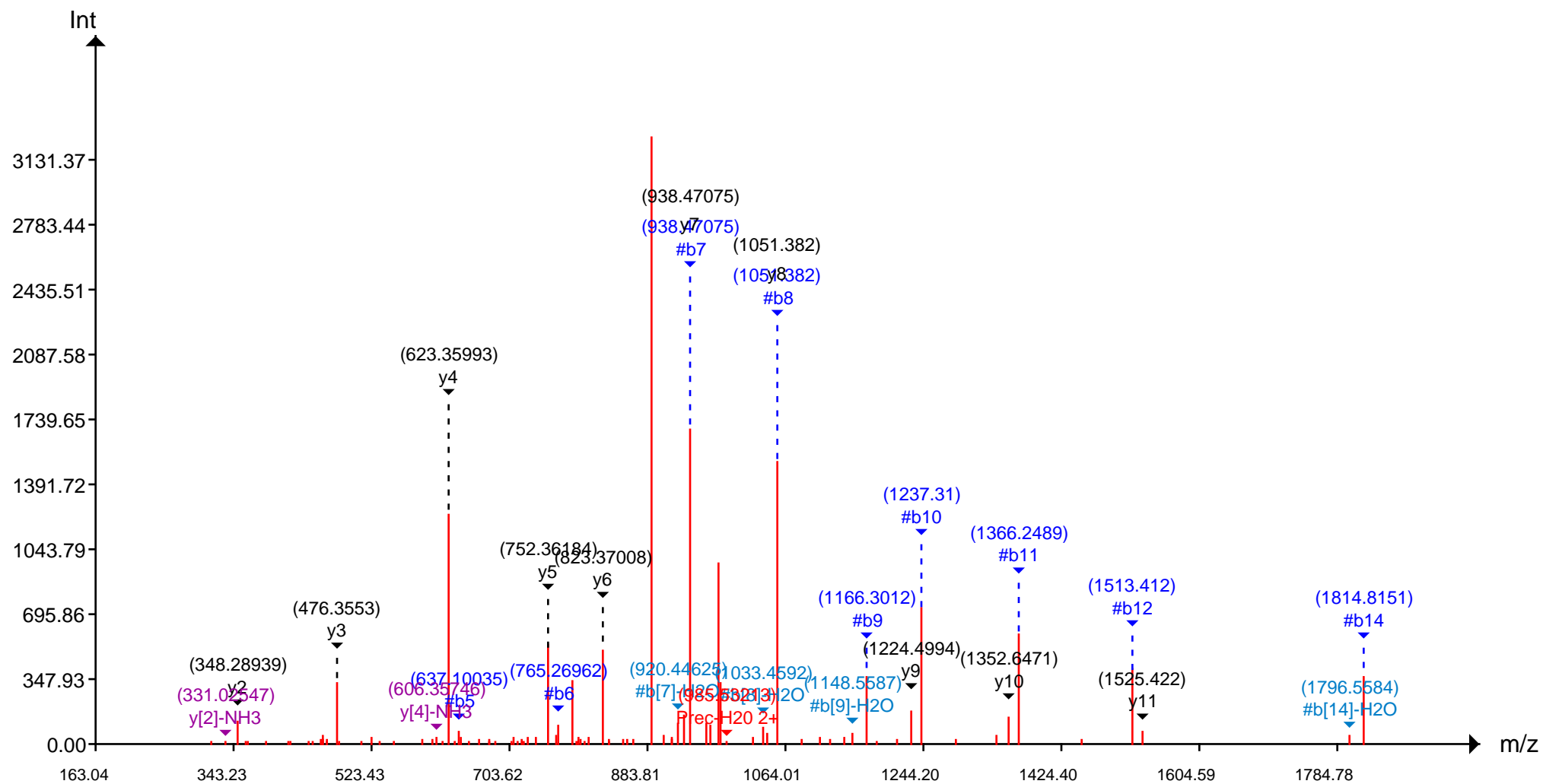
Confidence level 99.00%

Precursor m/z 527.78688

Identity Threshold 48.71

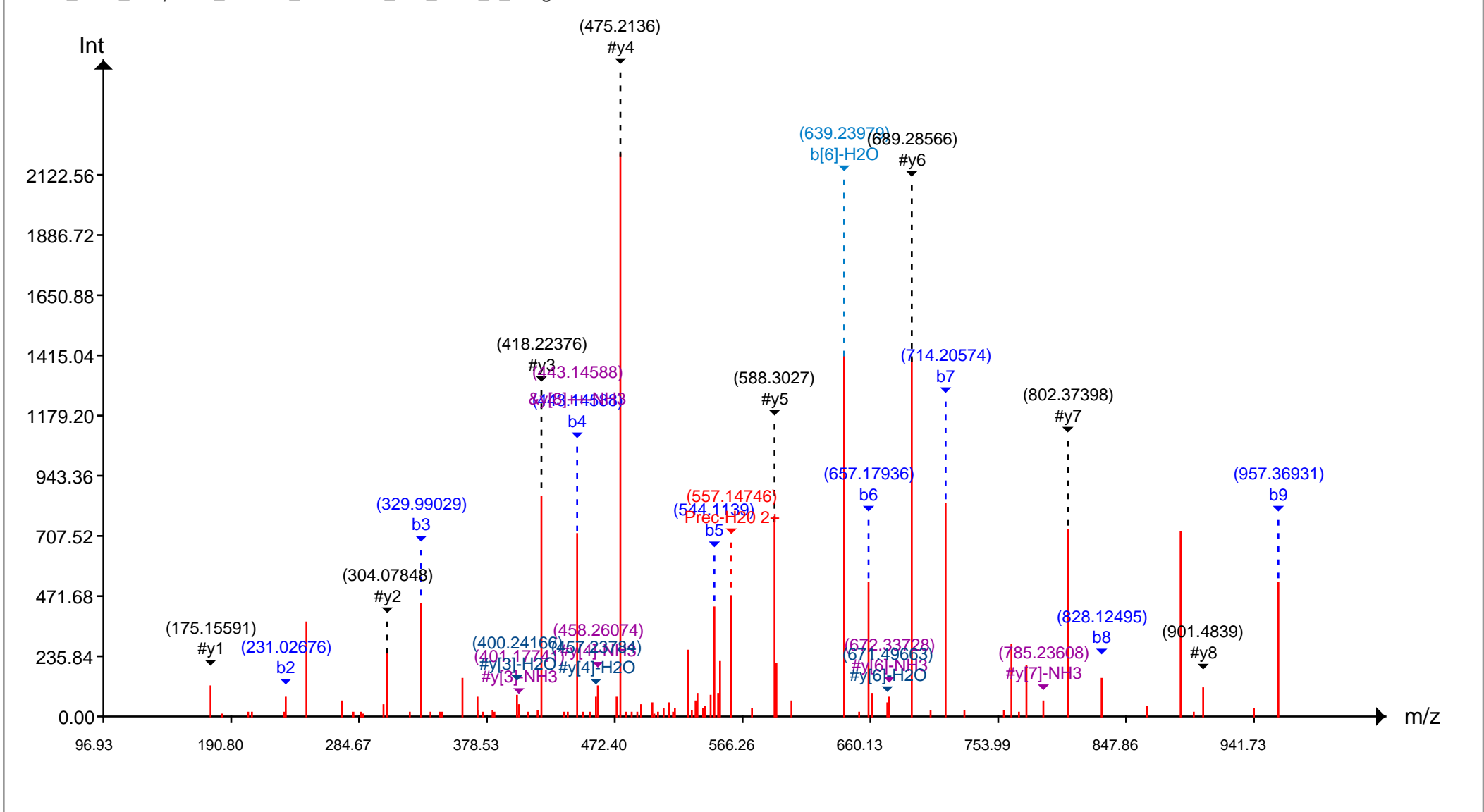
Charge 2+

Mass error 2.79E-4



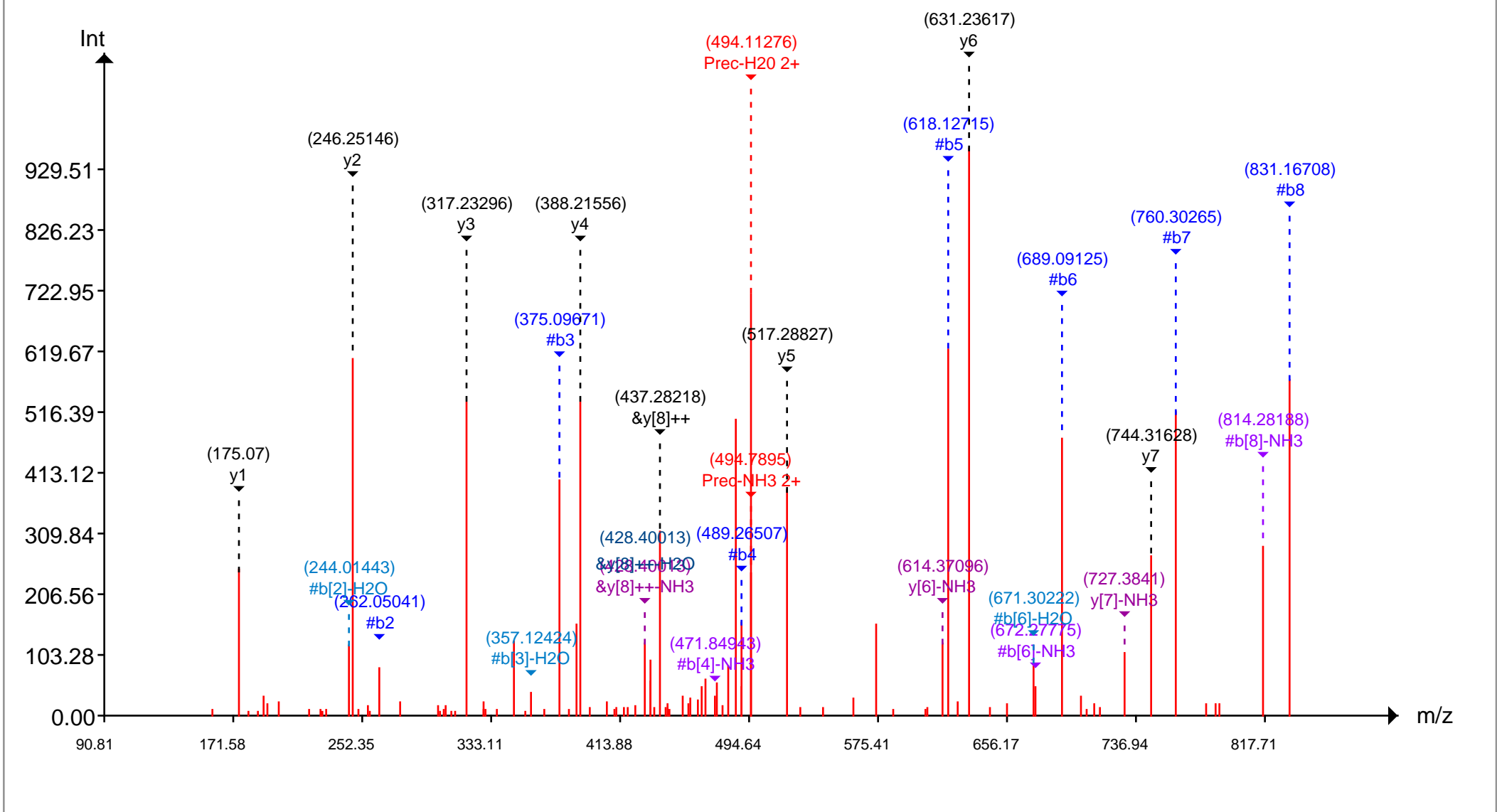
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-AM<Mox*>AEK<AcD3K*>QK<AcD3K*>LDAEFQK<AcD3K*>R-COOH | Ion score | 80.69 |
| Protein | Q9H875 | Confidence level | 99.00% |
| Precursor m/z | 995.02923 | Identity Threshold | 48.88 |
| Charge | 2+ | Mass error | 0.00129 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 72.59 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31739 | Identity Threshold | 48.79 |
| Charge | 2+ | Mass error | 0.007923 |



Info

Modified Sequence AcD3-SELNEAAAR-COOH

Protein Q12968

Precursor m/z 503.25541

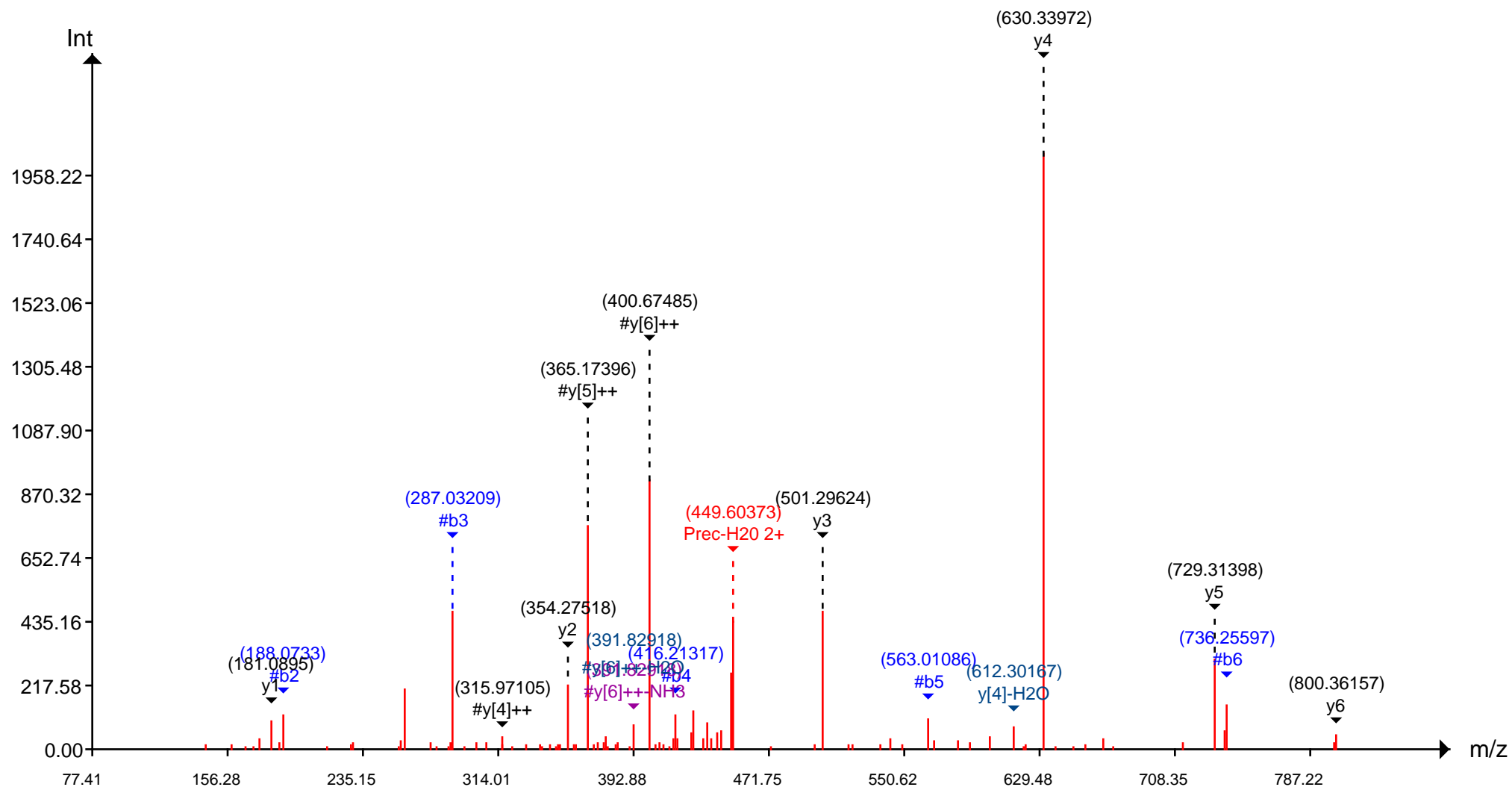
Charge 2+

Ion score 64.23

Confidence level 99.00%

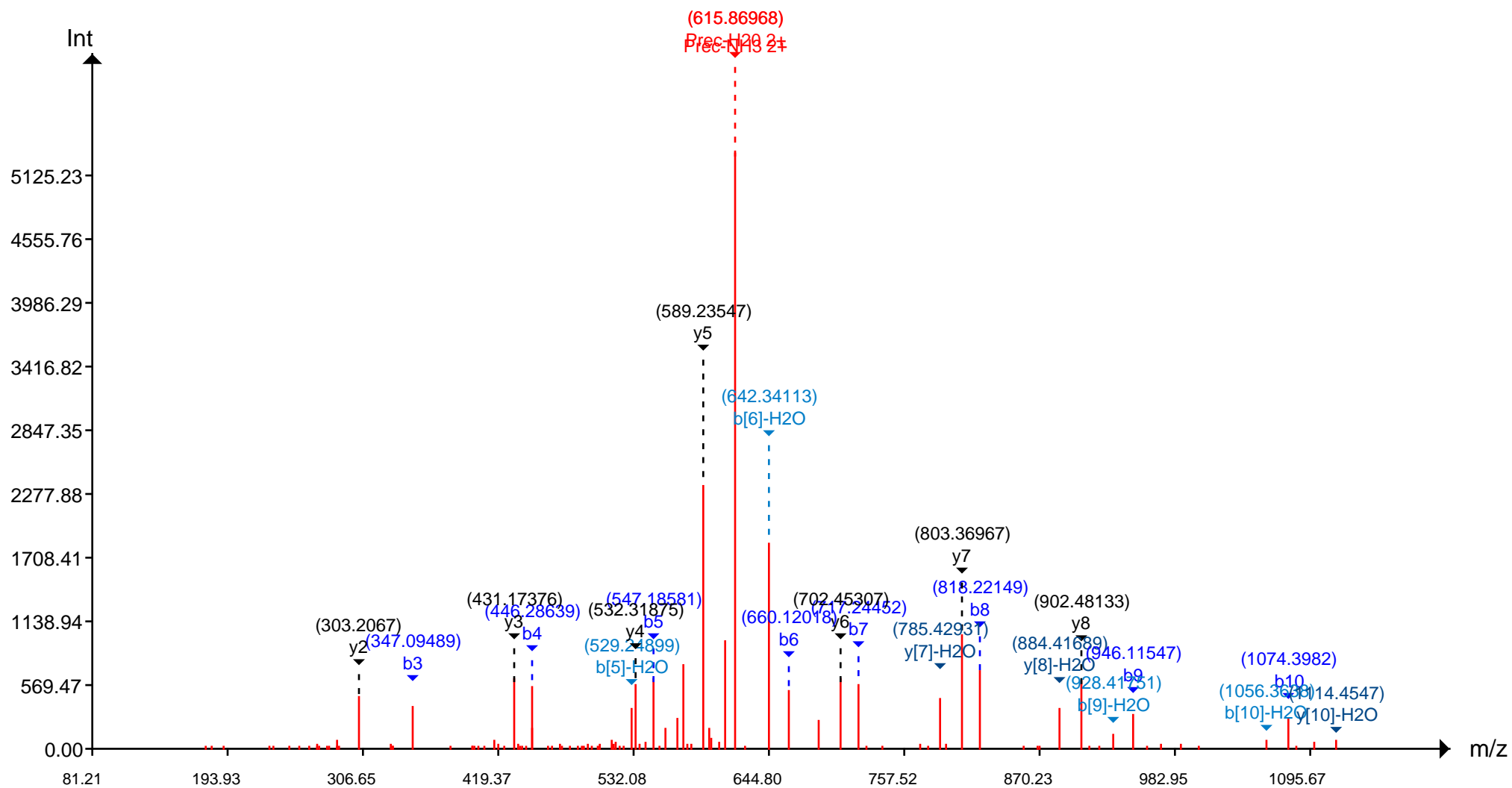
Identity Threshold 48.50

Mass error -2.99E-4



Info

| | | | |
|--------------------------|--------------------------------------|---------------------------|--------|
| Modified Sequence | AcD3-AAVEM<Mox*>K<AcD3K*>R<C13>-COOH | Ion score | 48.67 |
| Protein | Q16643 | Confidence level | 99.00% |
| Precursor m/z | 458.76041 | Identity Threshold | 48.18 |
| Charge | 2+ | Mass error | 8.8E-5 |



Info

Modified Sequence AcD3-AGK<AcD3K*>VTLGTQQR-COOH

Ion score 58.47

Protein Q08AF3

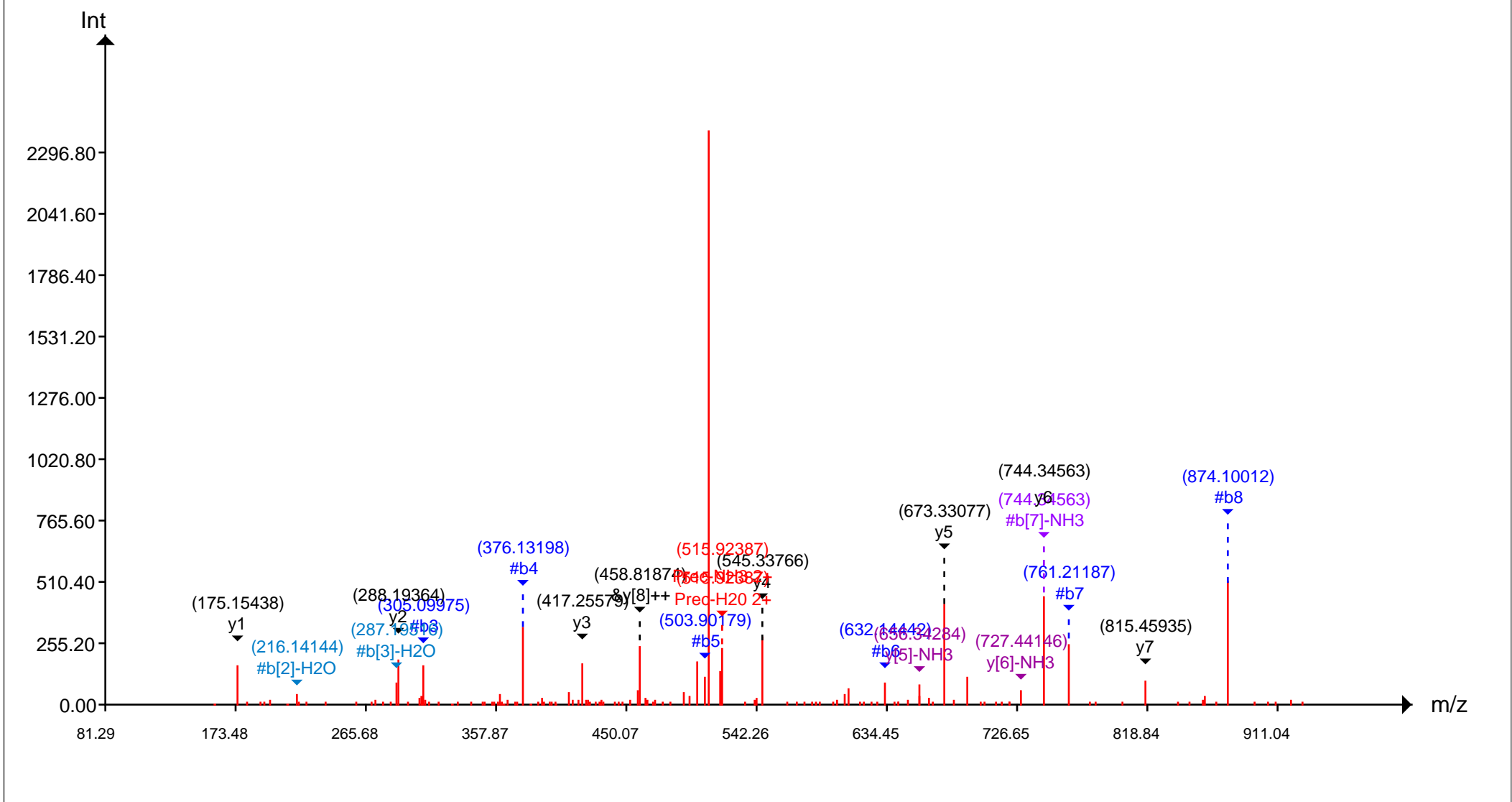
Confidence level 99.00%

Precursor m/z 624.86194

Identity Threshold 47.61

Charge 2+

Mass error -0.001136



Info

Modified Sequence AcD3-STAAQQELR-COOH

Protein P35579

Precursor m/z 524.77666

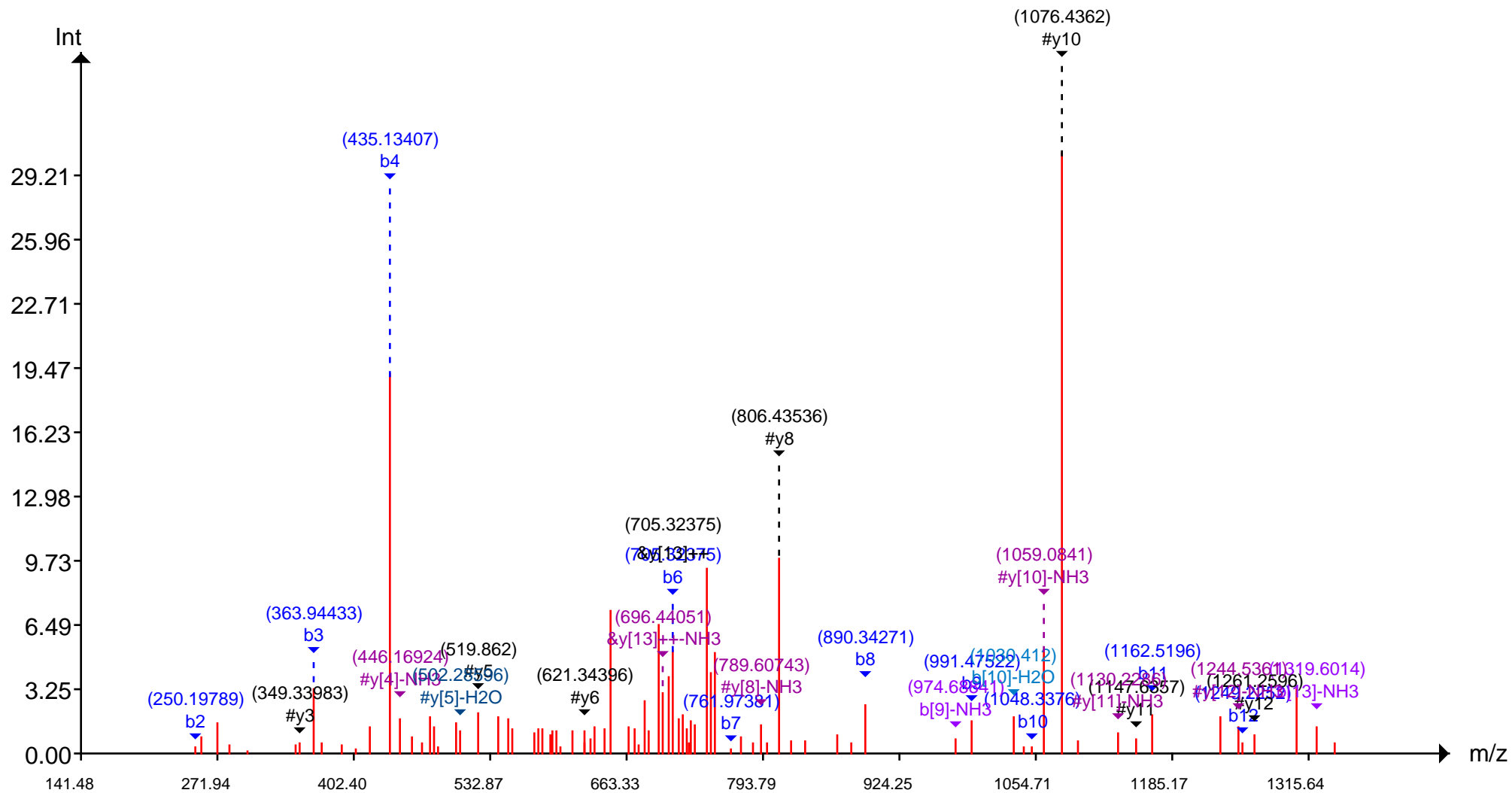
Charge 2+

Ion score 73.8

Confidence level 99.00%

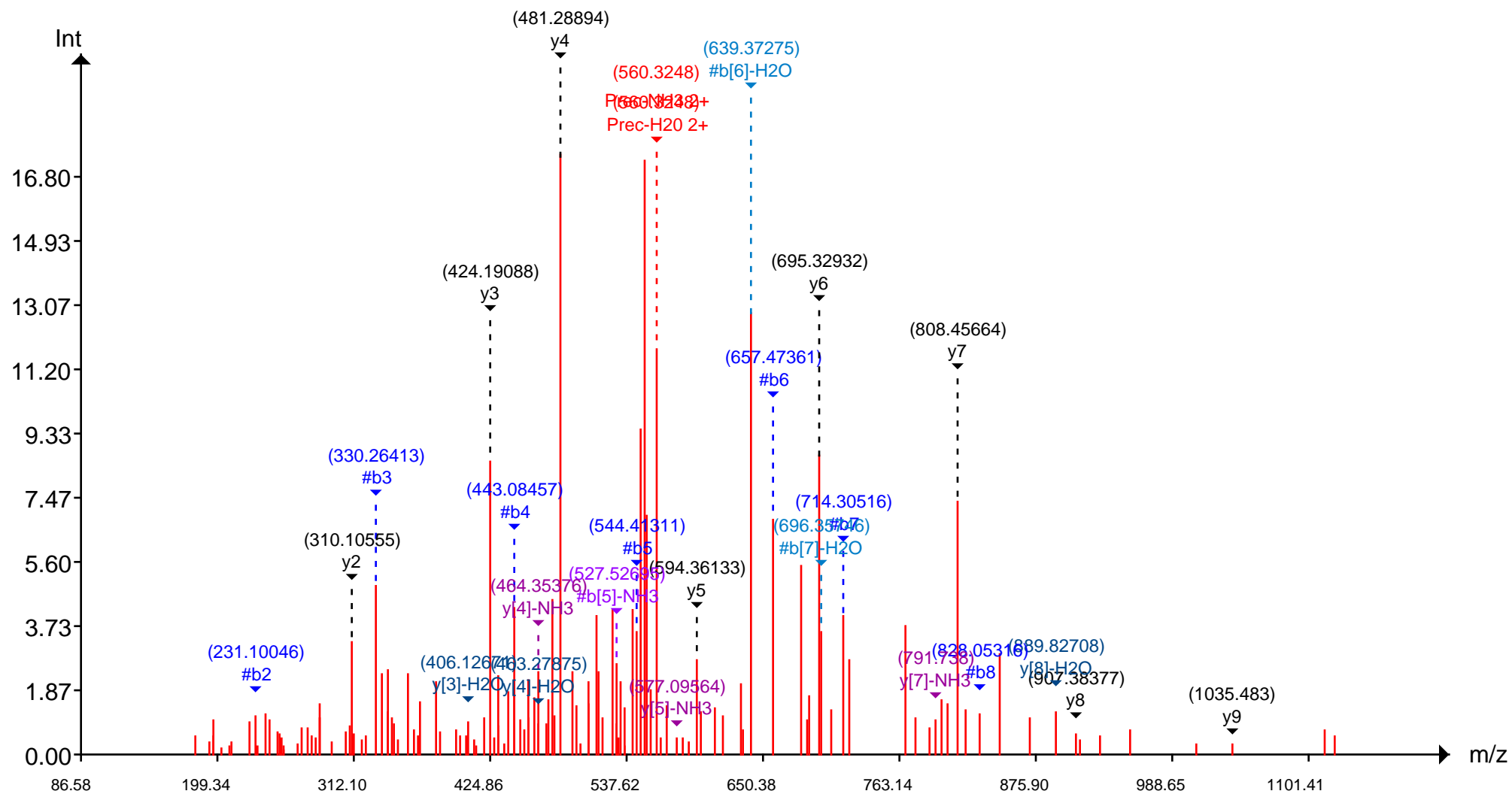
Identity Threshold 49.07

Mass error -2.0E-5



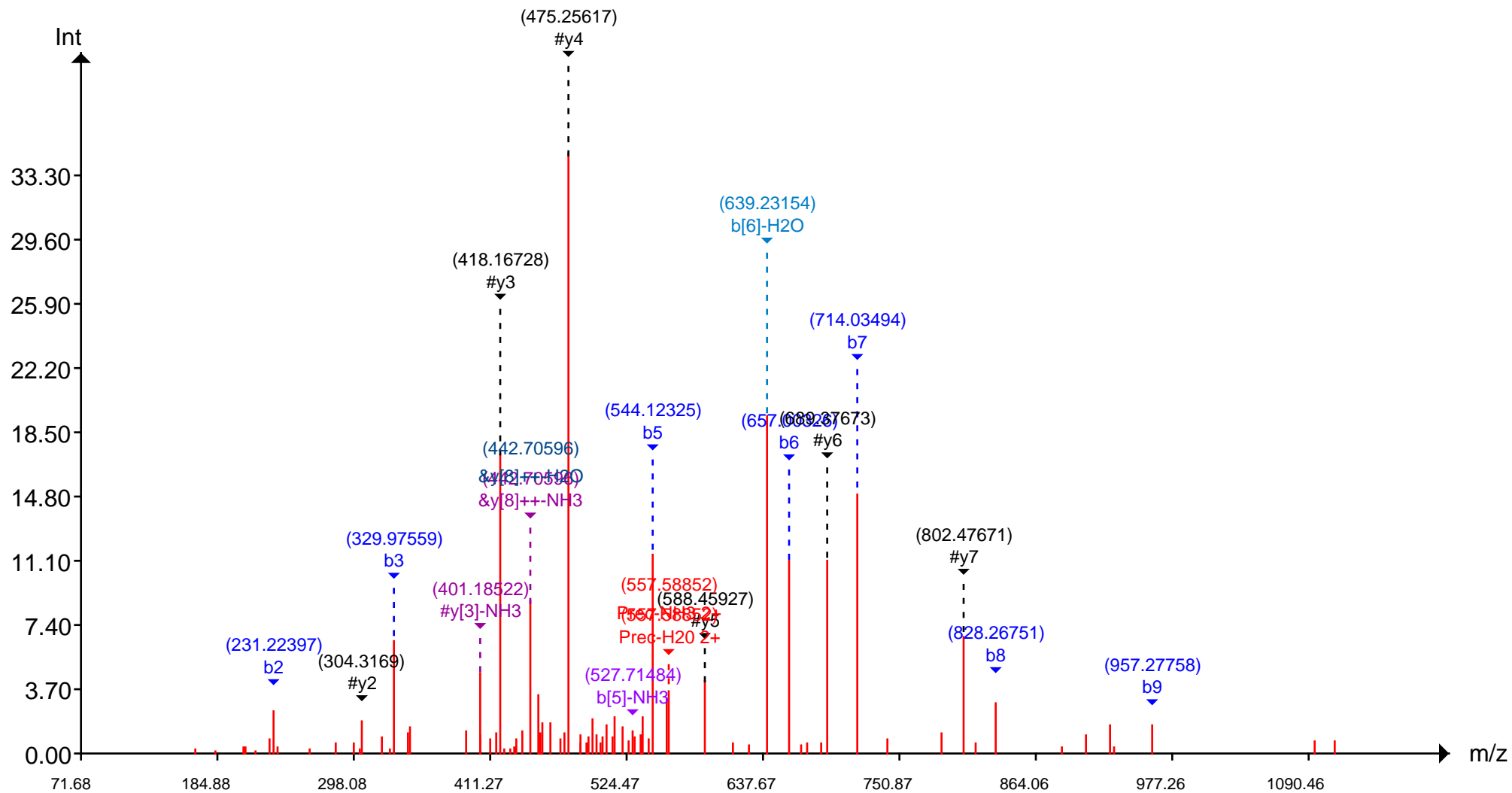
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR-COOH | Ion score | 49.41 |
| Protein | P23588 | Confidence level | 99.00% |
| Precursor m/z | 755.86322 | Identity Threshold | 47.76 |
| Charge | 2+ | Mass error | 6.3E-4 |



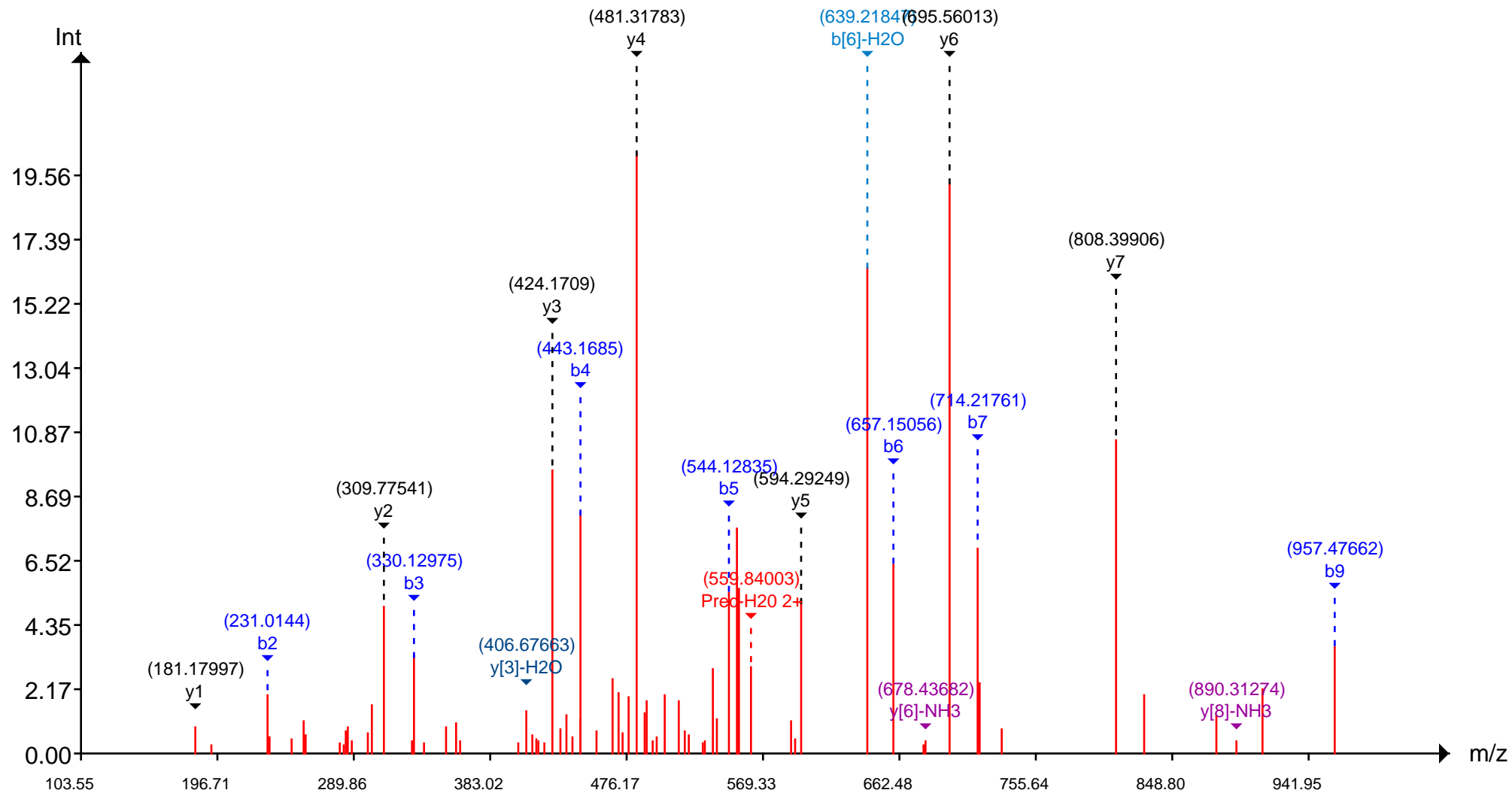
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 58.2 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32391 | Identity Threshold | 47.94 |
| Charge | 2+ | Mass error | 8.22E-4 |



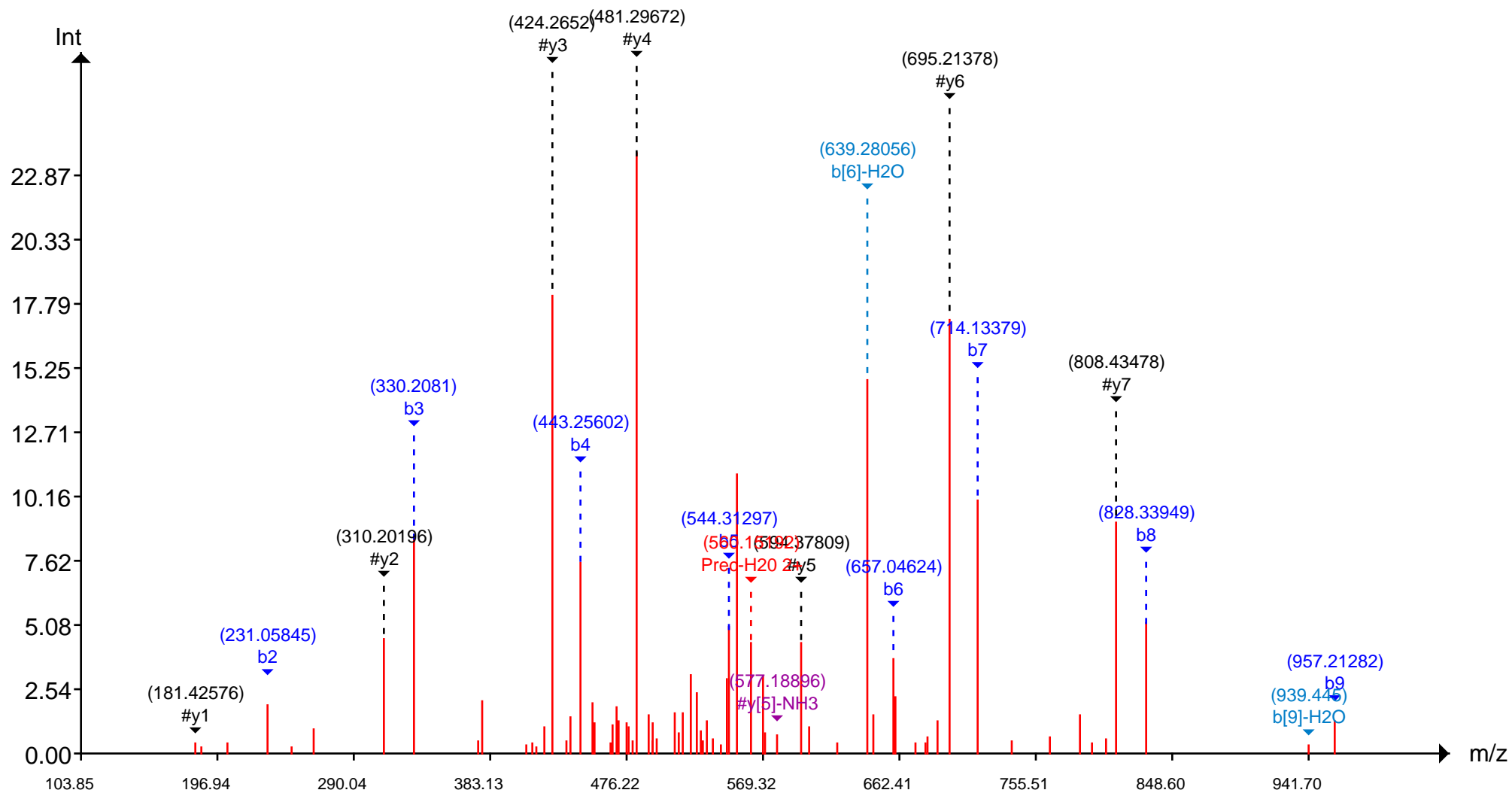
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 61.26 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31338 | Identity Threshold | 48.99 |
| Charge | 2+ | Mass error | -9.7E-5 |



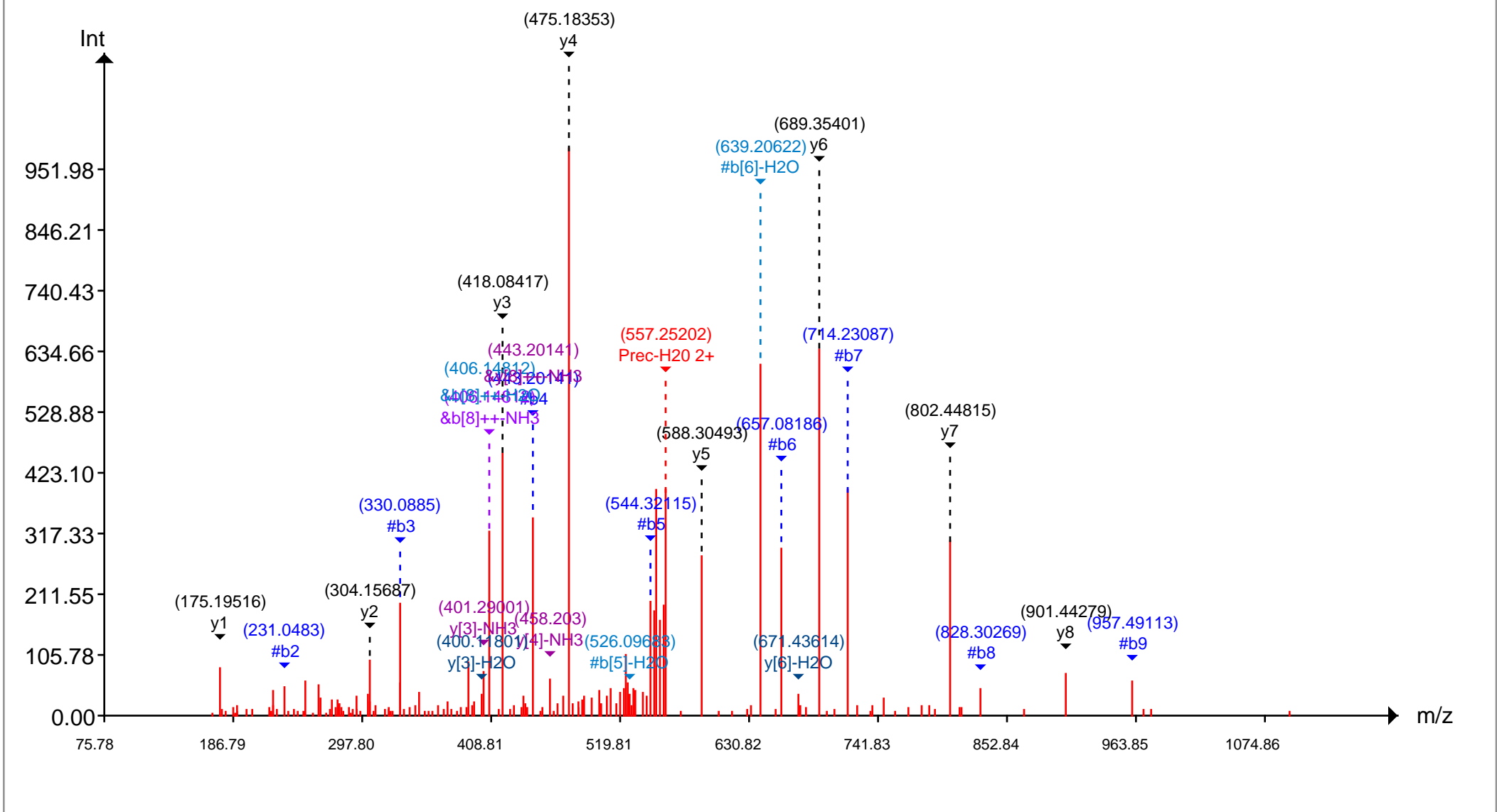
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 54.08 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32361 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 2.22E-4 |



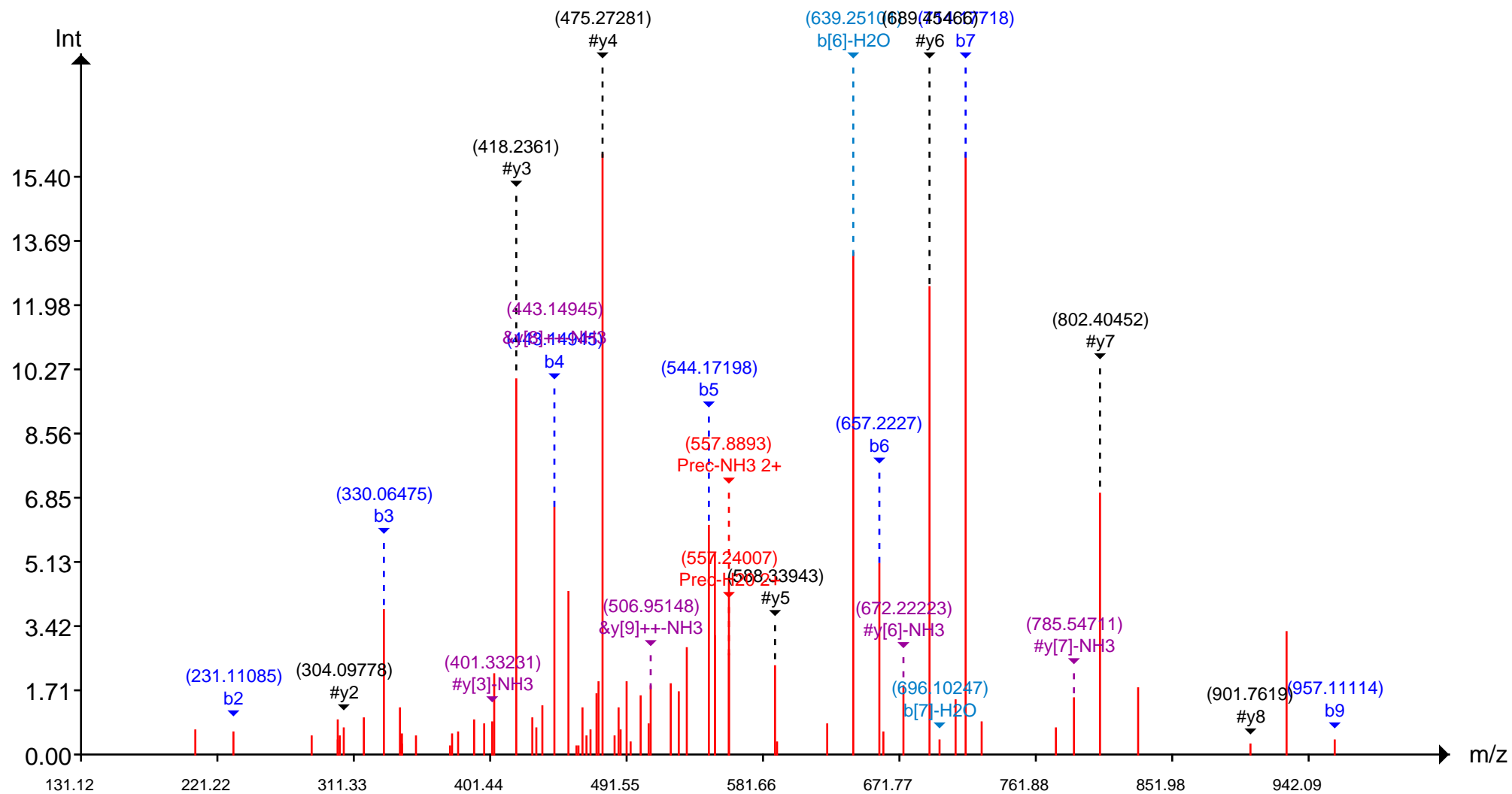
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 69.55 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.3236 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 2.02E-4 |



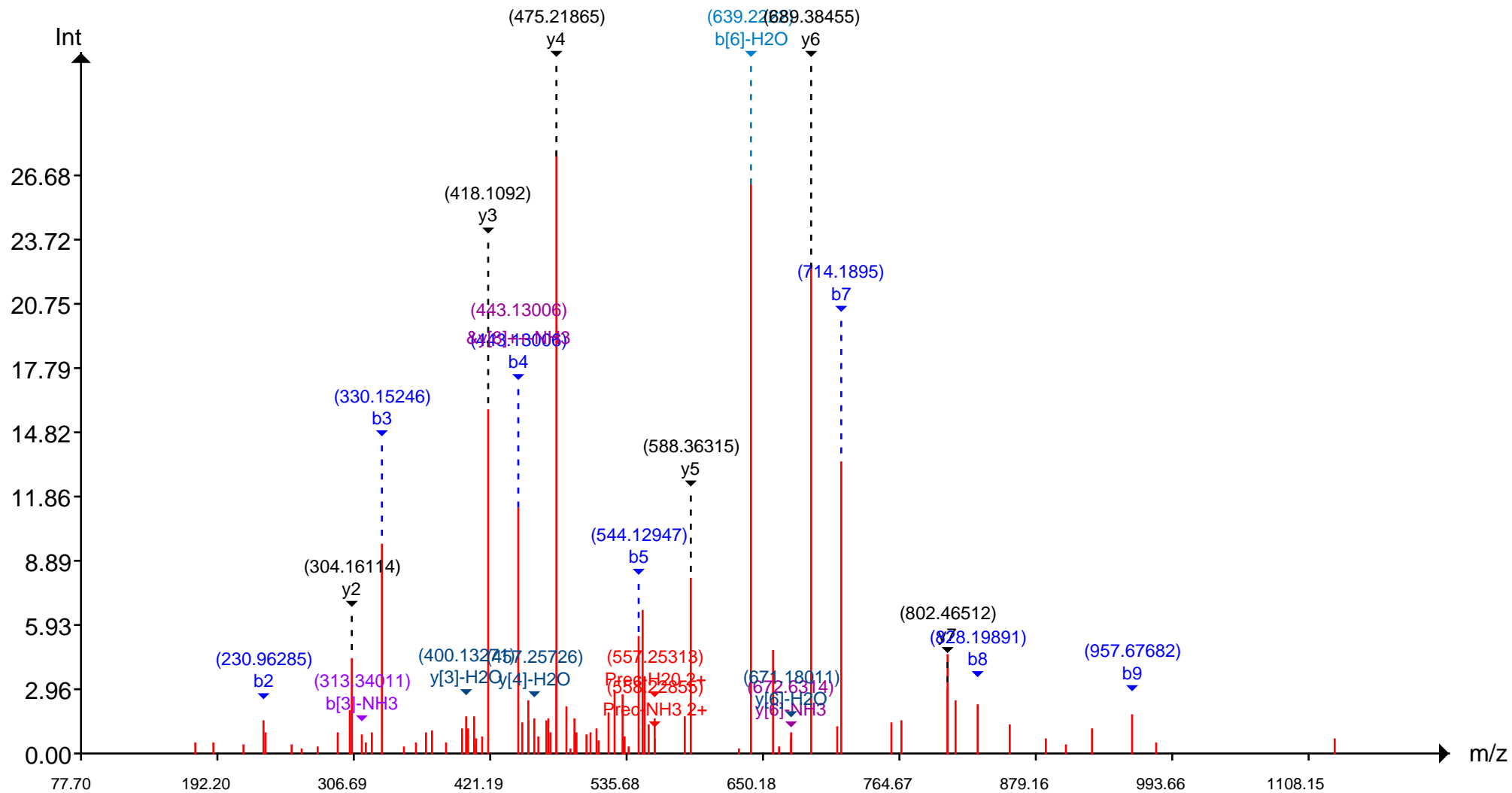
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 63.11 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31316 | Identity Threshold | 48.90 |
| Charge | 2+ | Mass error | -5.37E-4 |



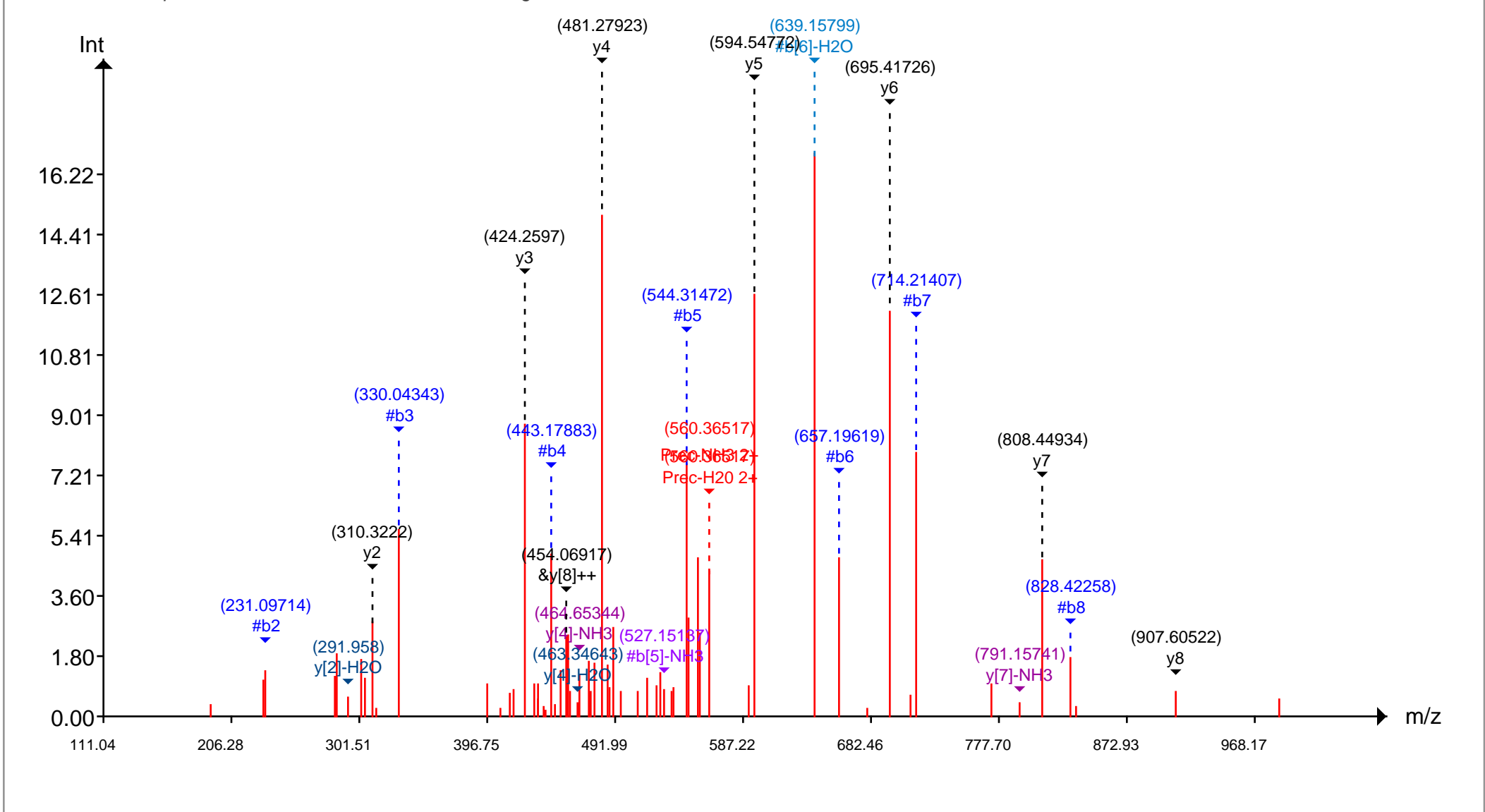
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 49.98 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31324 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | -3.77E-4 |



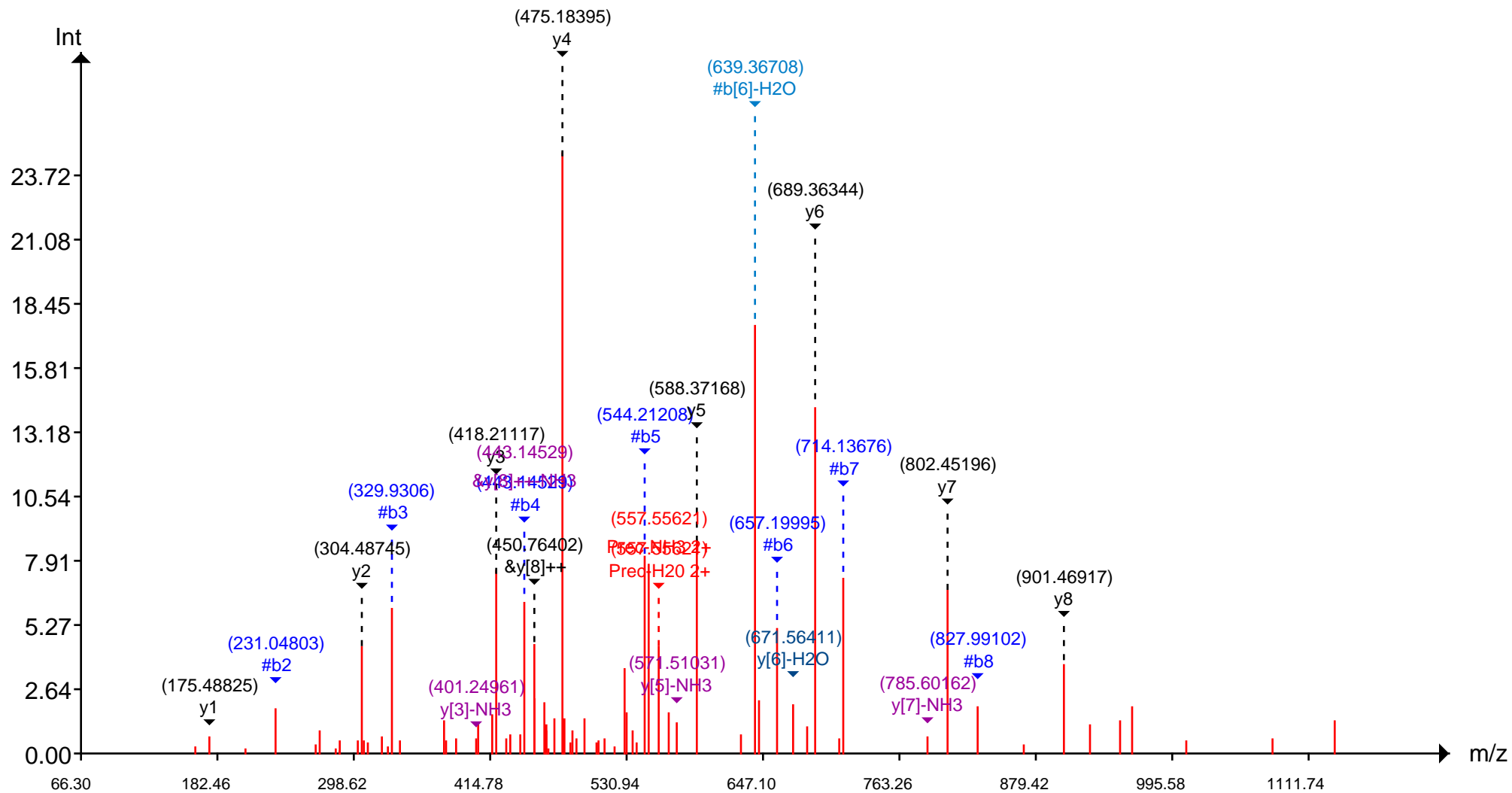
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 50.46 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31346 | Identity Threshold | 48.97 |
| Charge | 2+ | Mass error | 6.3E-5 |



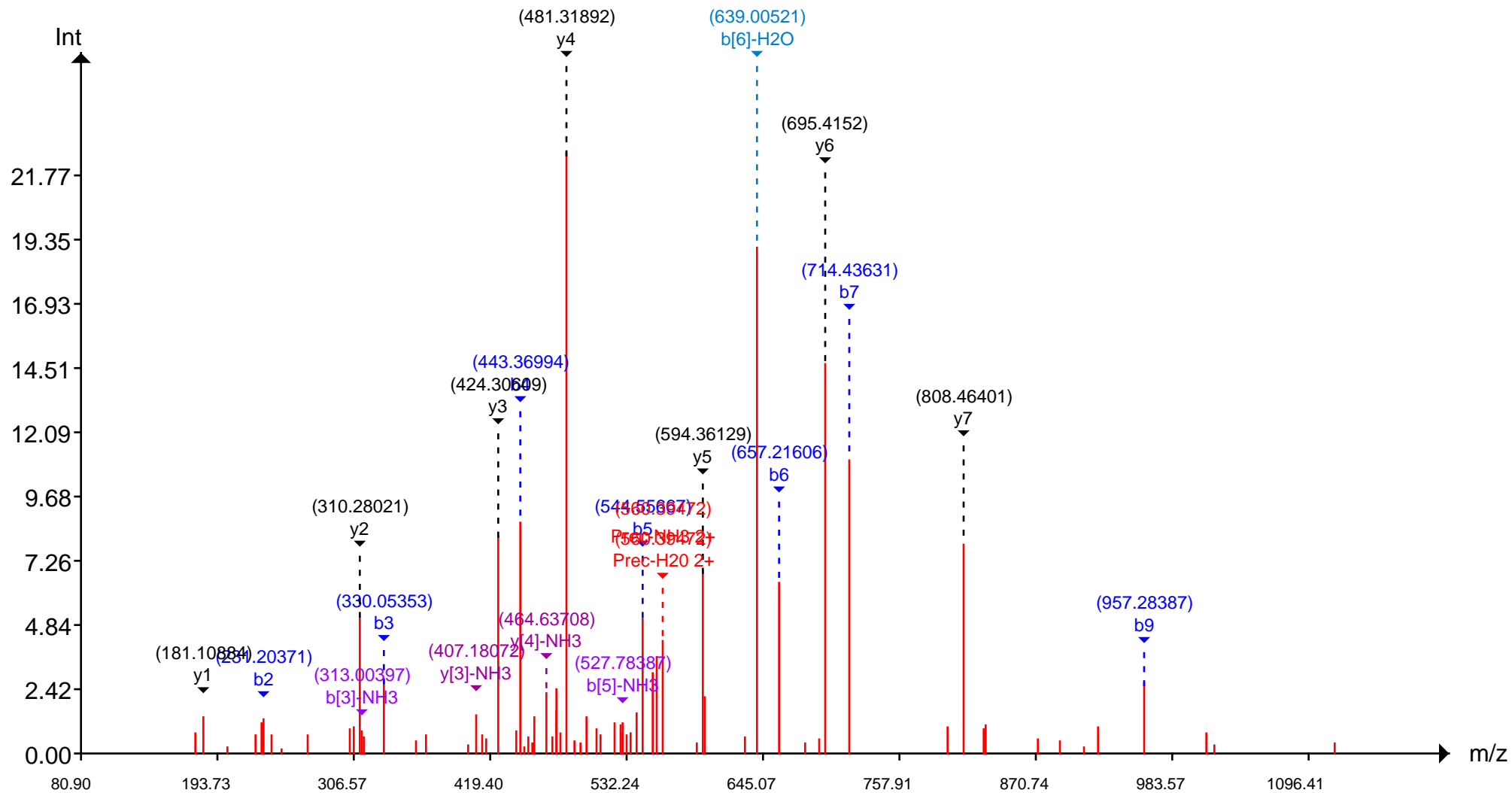
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 67.23 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32348 | Identity Threshold | 48.30 |
| Charge | 2+ | Mass error | -3.8E-5 |



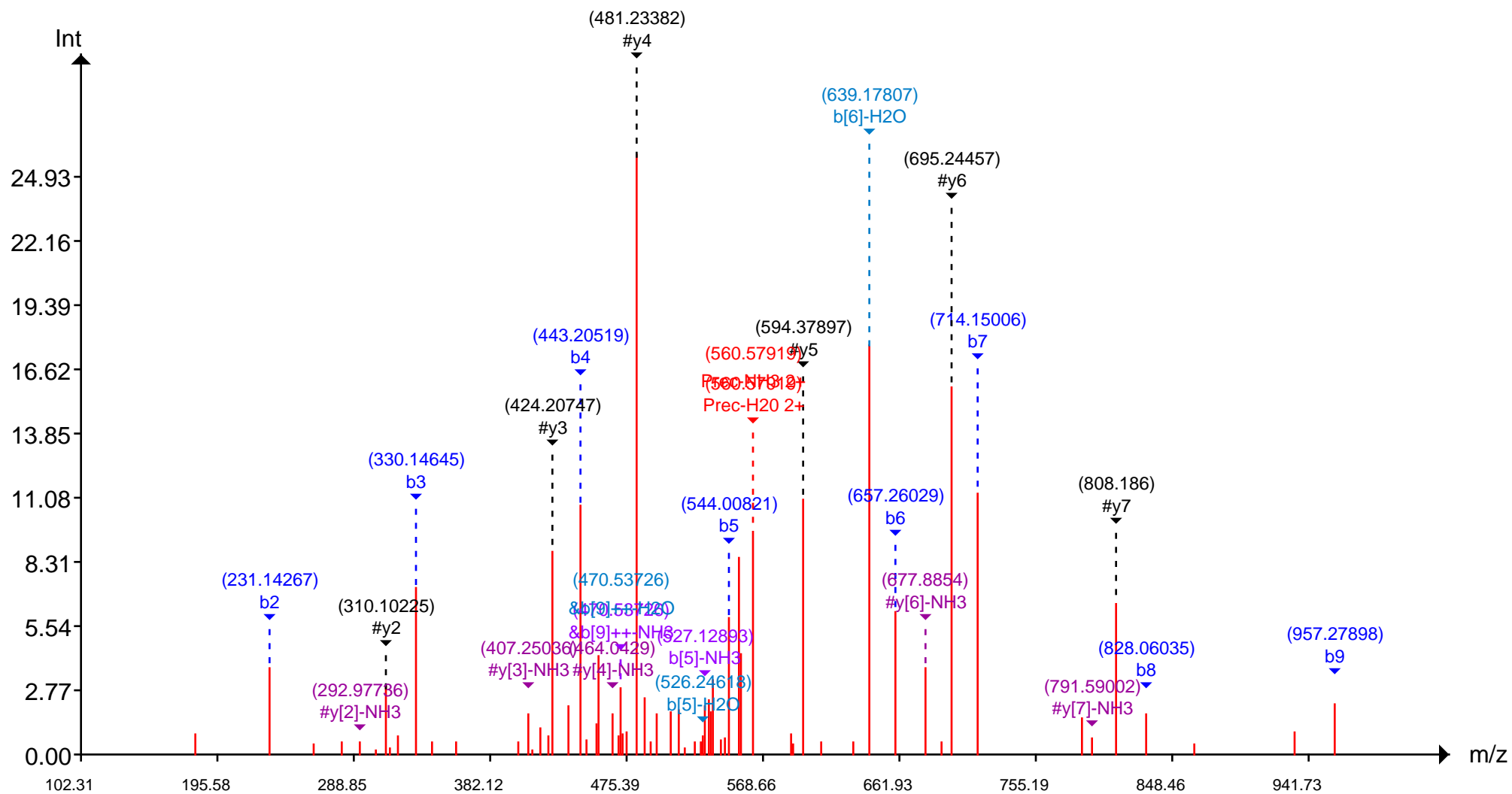
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 77.09 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31345 | Identity Threshold | 49.03 |
| Charge | 2+ | Mass error | 4.3E-5 |



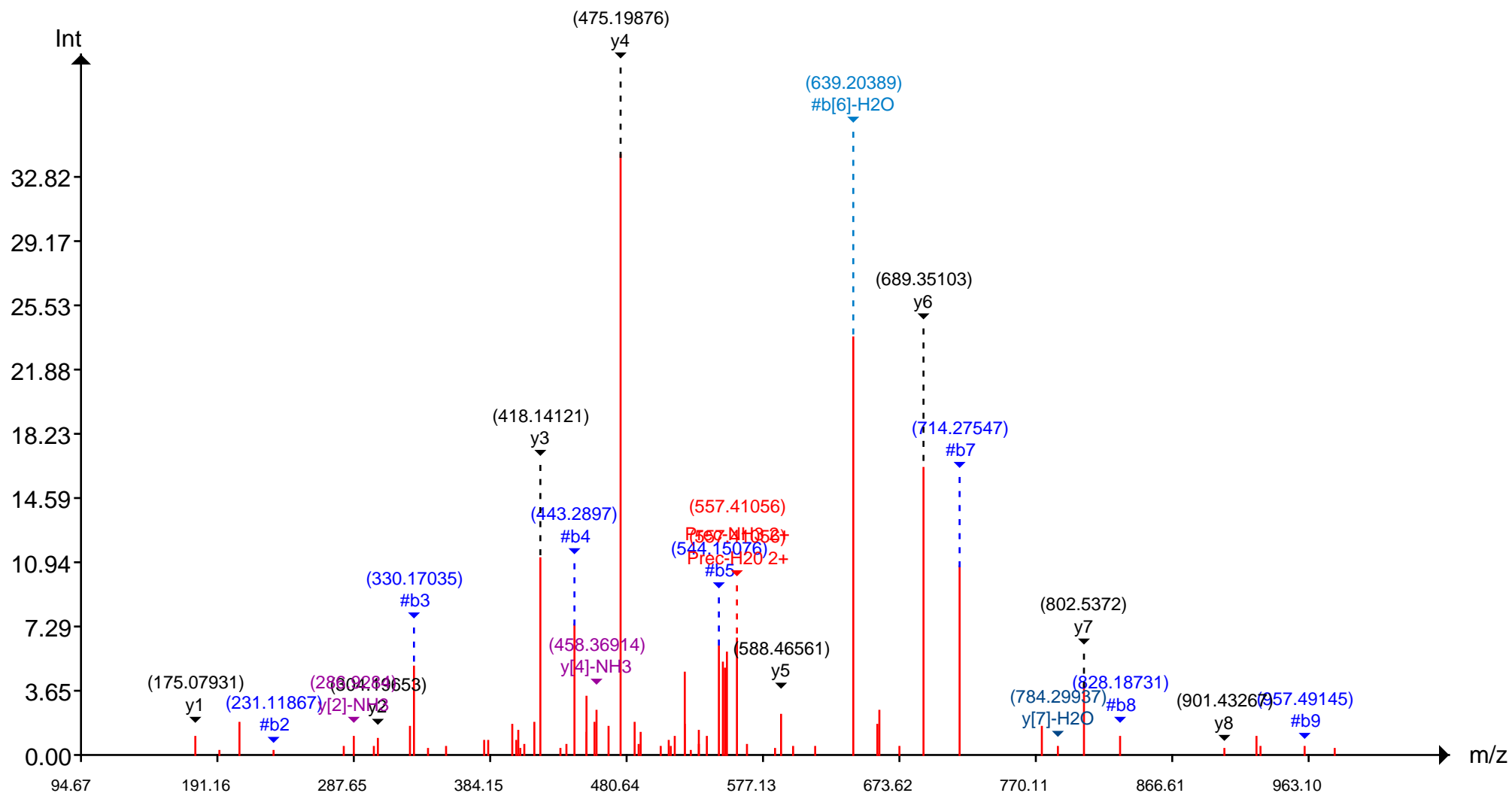
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 60.53 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32322 | Identity Threshold | 48.30 |
| Charge | 2+ | Mass error | -5.58E-4 |



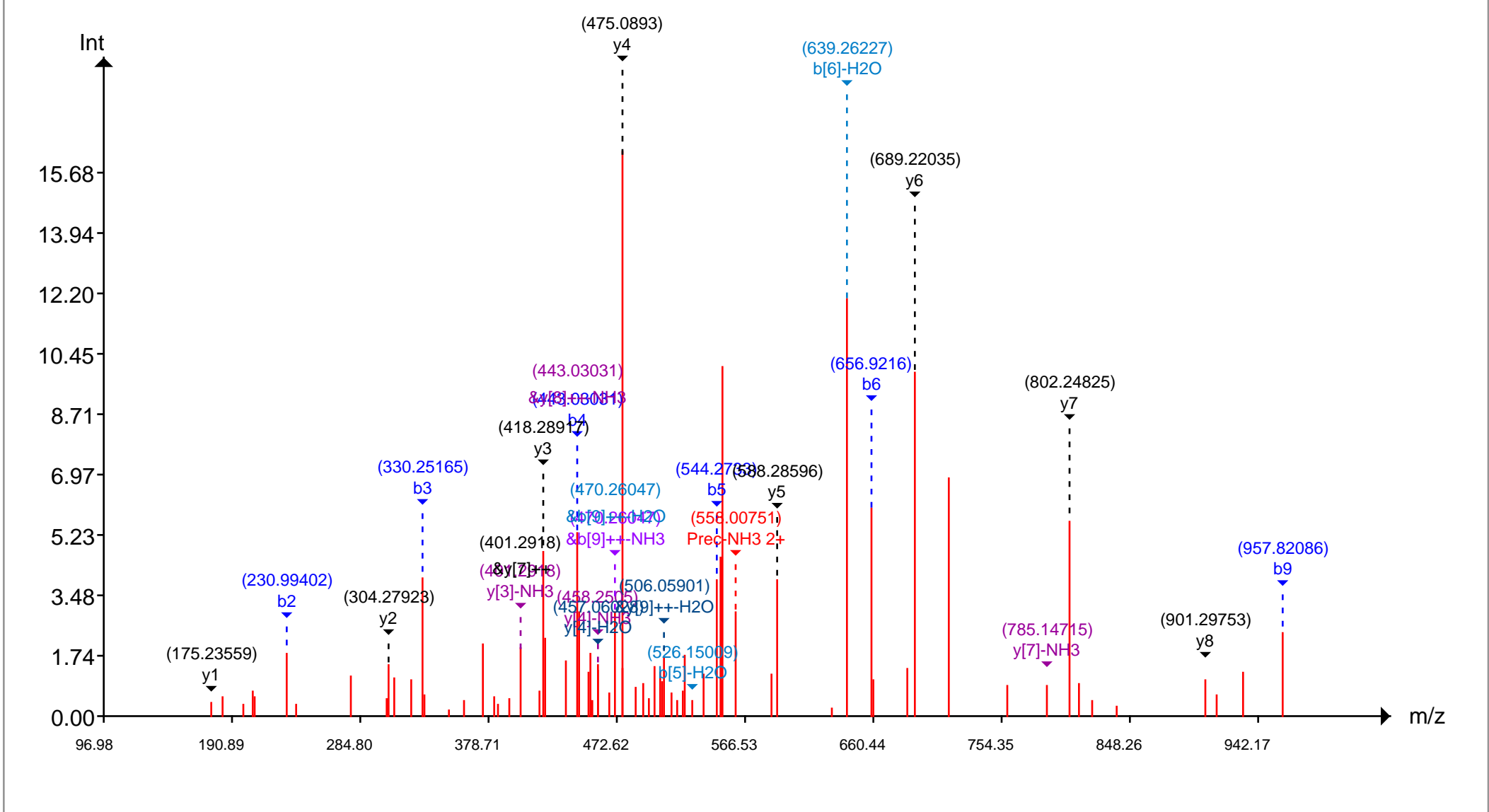
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 64.64 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32315 | Identity Threshold | 48.30 |
| Charge | 2+ | Mass error | -6.98E-4 |



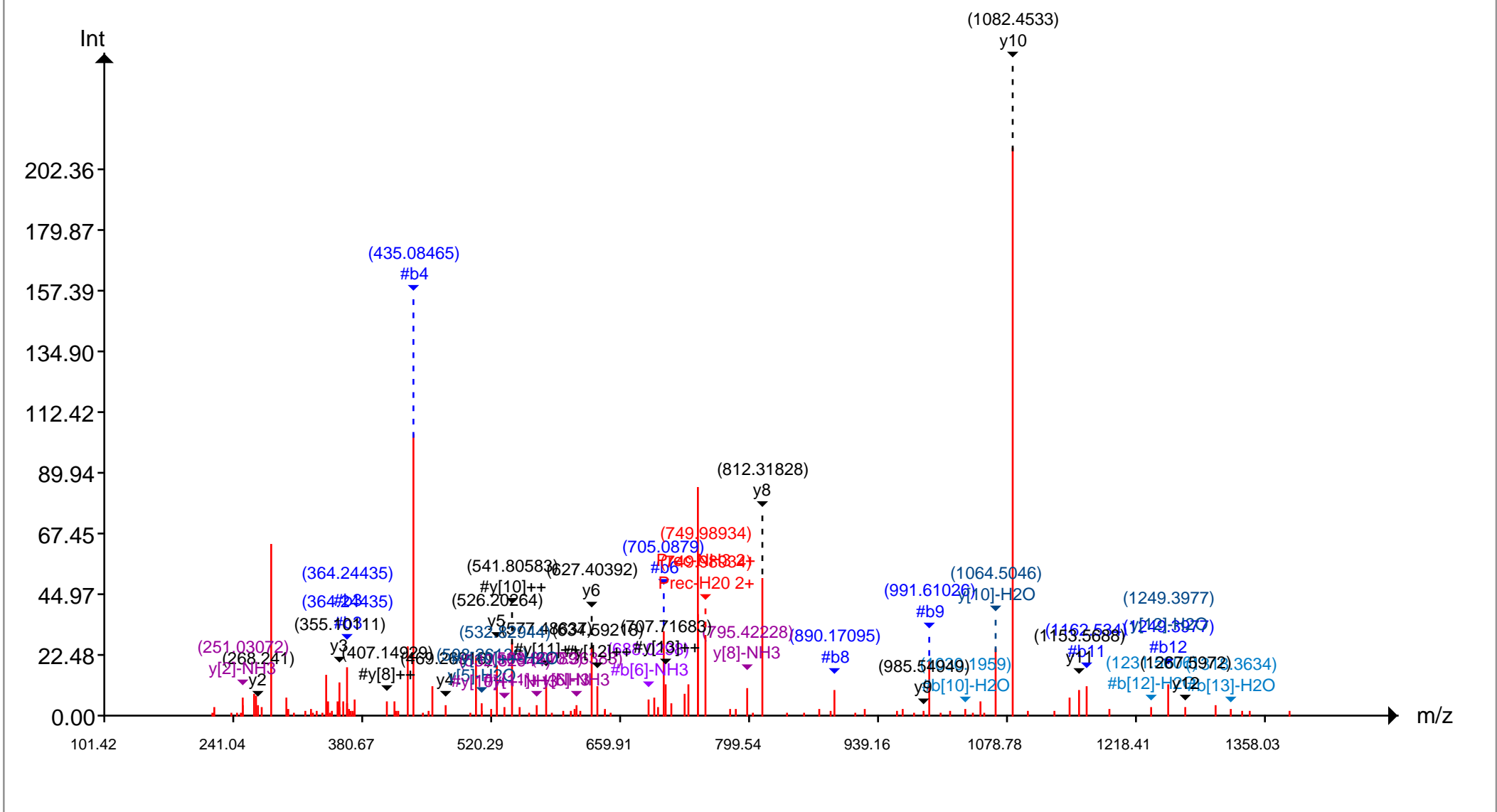
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 58.91 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31356 | Identity Threshold | 48.80 |
| Charge | 2+ | Mass error | 2.63E-4 |



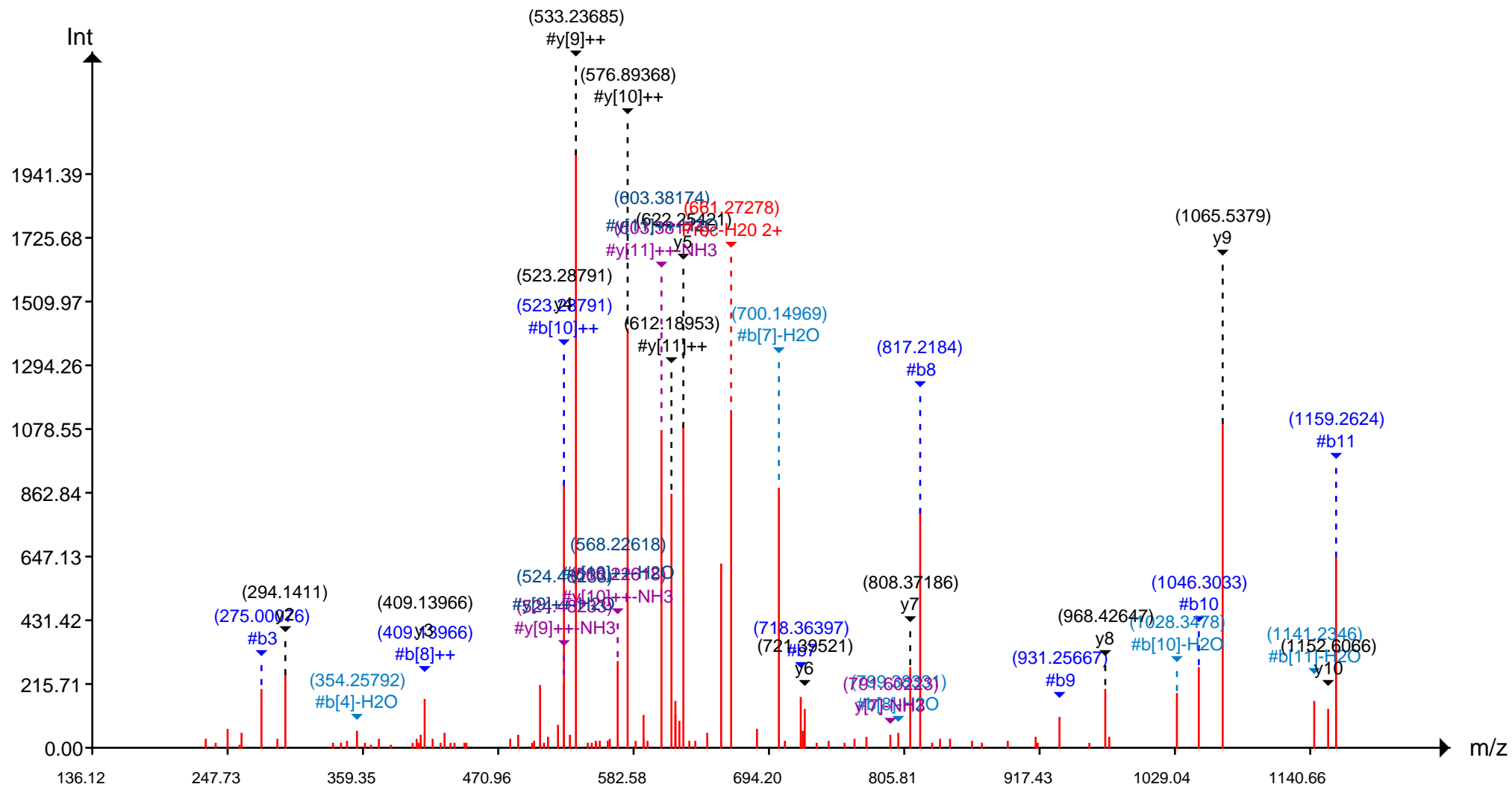
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 53.65 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.3135 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 1.43E-4 |



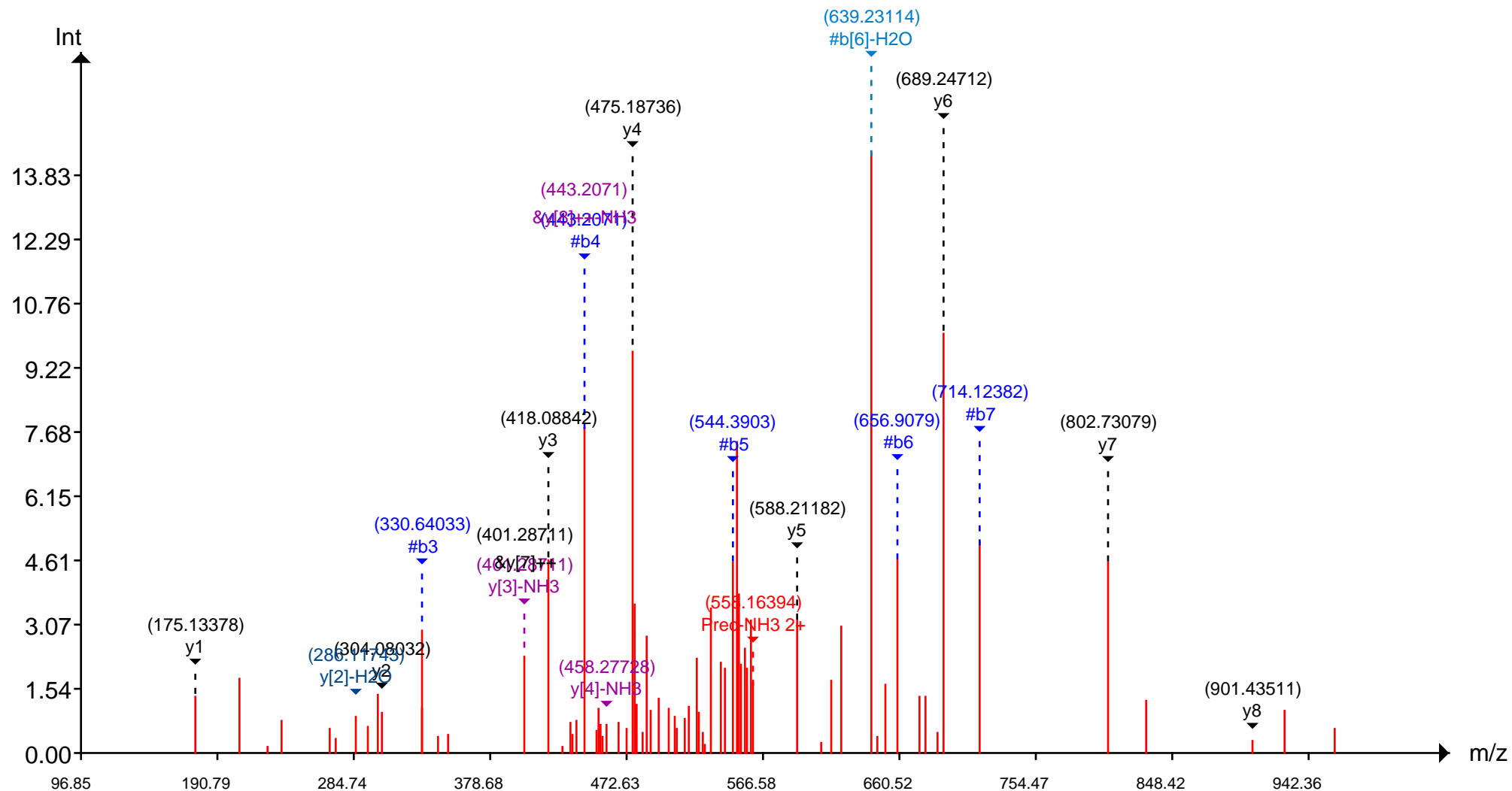
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR<C13>-COOH | Ion score | 56.41 |
| Protein | P23588 | Confidence level | 99.00% |
| Precursor m/z | 758.87325 | Identity Threshold | 48.20 |
| Charge | 2+ | Mass error | 5.48E-4 |



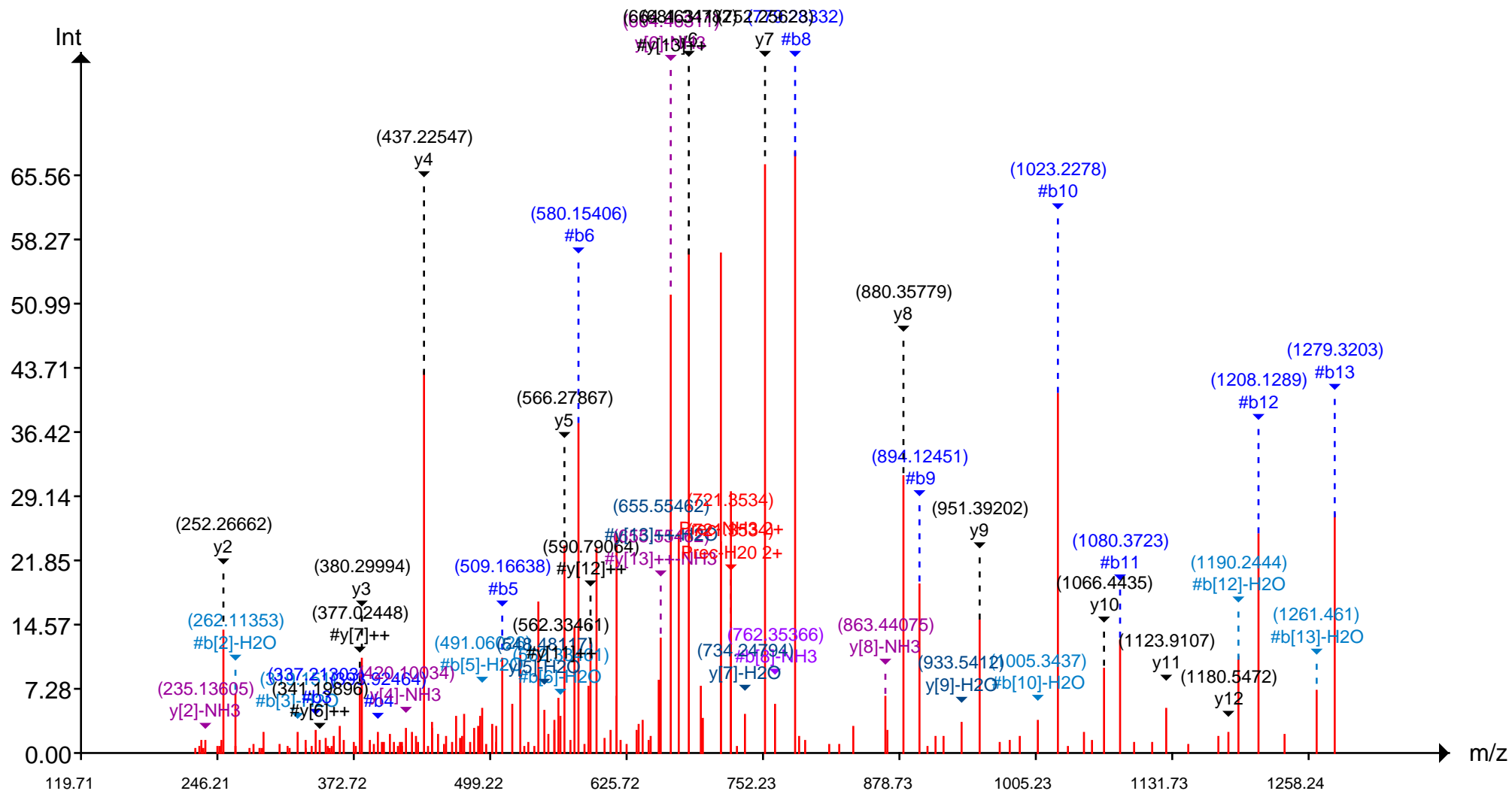
Info

| | | | |
|--------------------------|-----------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-AASPC<Cmm*>SVVNDLR<C13>-COOH | Ion score | 66.68 |
| Protein | P52888 | Confidence level | 99.00% |
| Precursor m/z | 670.34422 | Identity Threshold | 48.88 |
| Charge | 2+ | Mass error | 2.28E-4 |

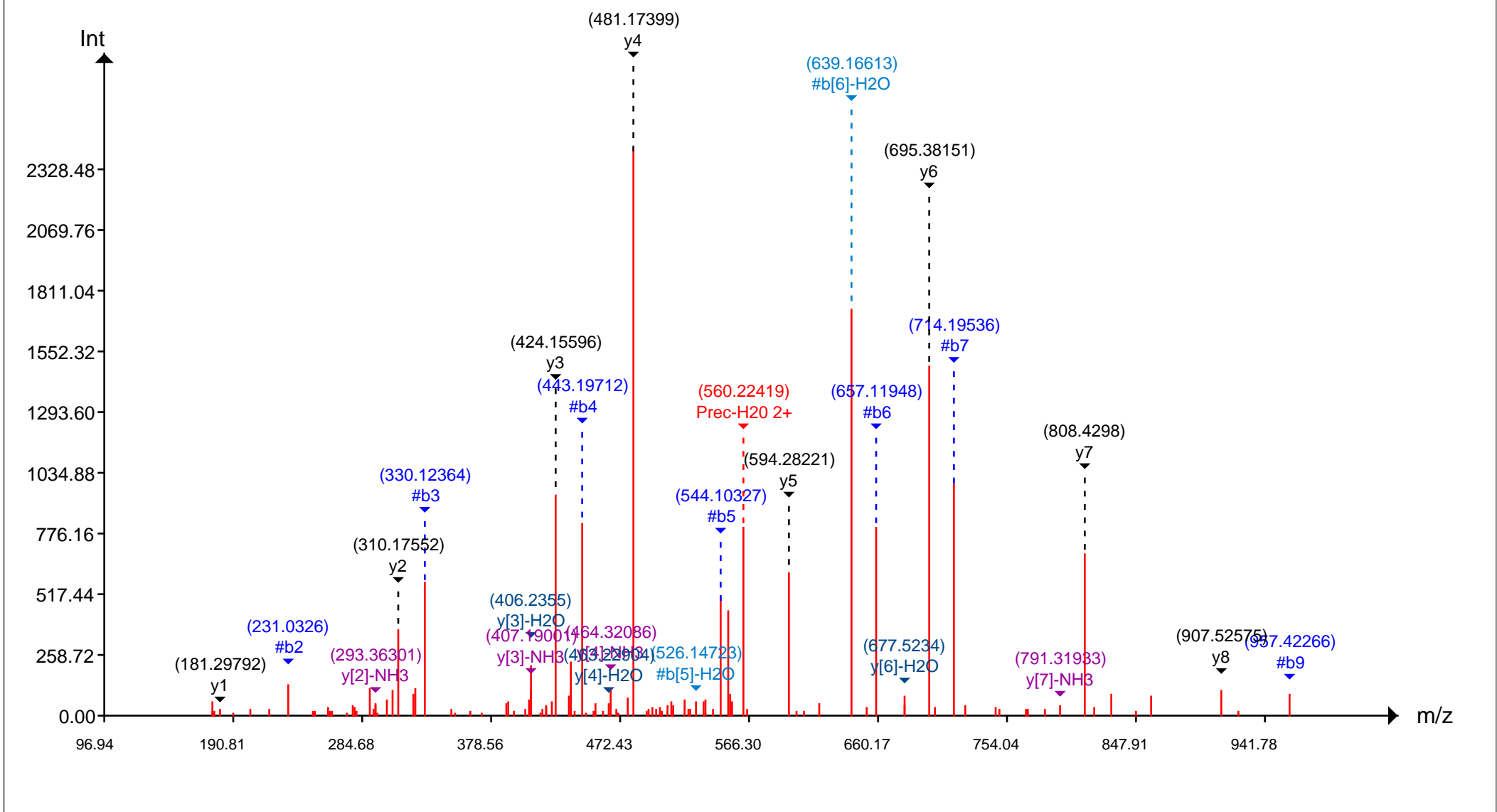


Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 55.5 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31366 | Identity Threshold | 48.78 |
| Charge | 2+ | Mass error | 4.63E-4 |

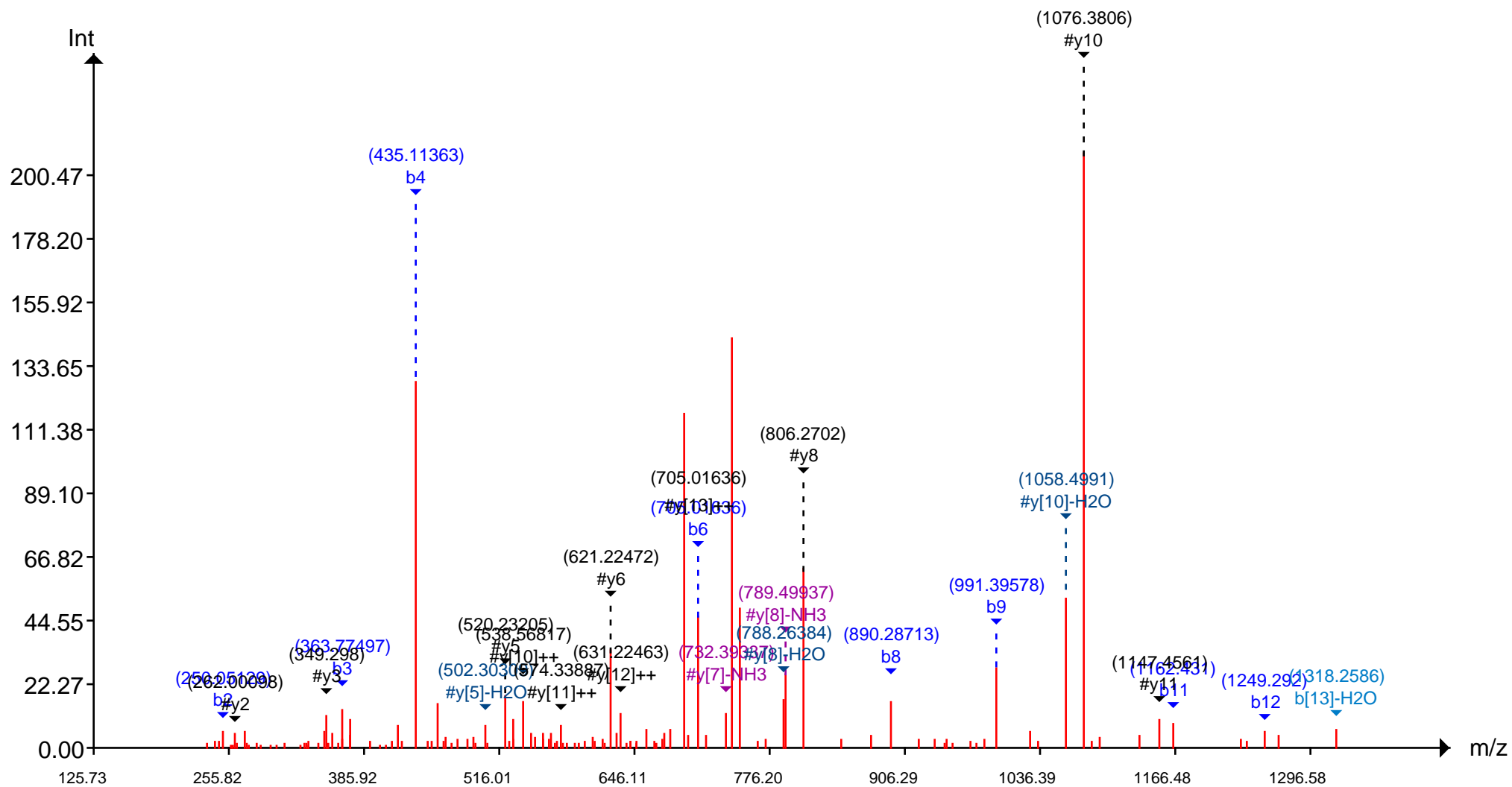


| Info | | | |
|--------------------------|-------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-SFGGDAQADEGQAR<C13>-COOH | Ion score | 109.88 |
| Protein | P33992 | Confidence level | 99.00% |
| Precursor m/z | 730.33284 | Identity Threshold | 45.42 |
| Charge | 2+ | Mass error | 1.12E-4 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 59.49 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32306 | Identity Threshold | 48.31 |
| Charge | 2+ | Mass error | -8.78E-4 |



Info

Modified Sequence AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR-COOH

Ion score 68.51

Protein P23588

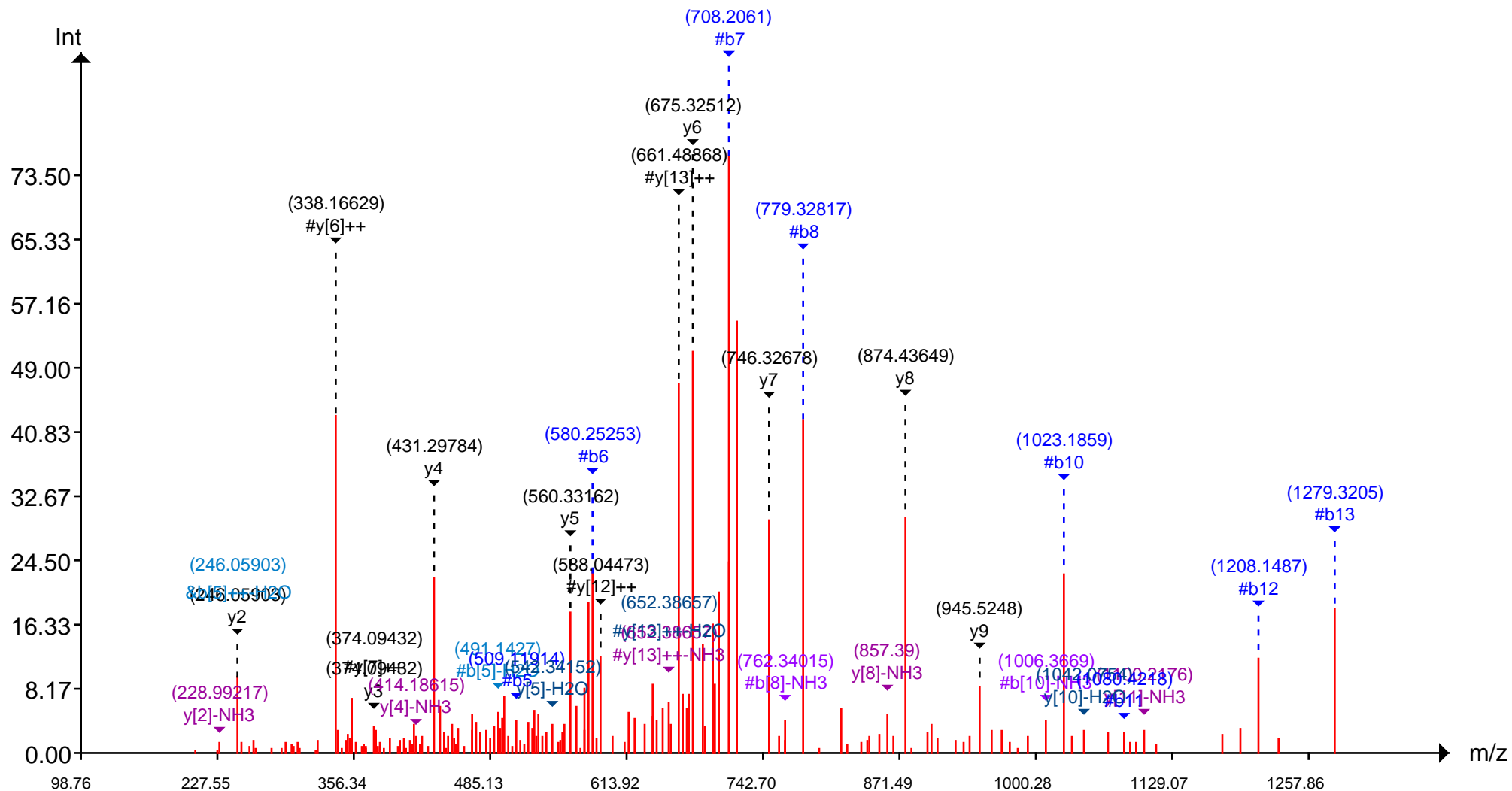
Confidence level 99.00%

Precursor m/z 755.86311

Identity Threshold 47.73

Charge 2+

Mass error 4.1E-4



Info

Modified Sequence AcD3-SFGGDAQADEGQAR-COOH

Protein P33992

Precursor m/z 727.32307

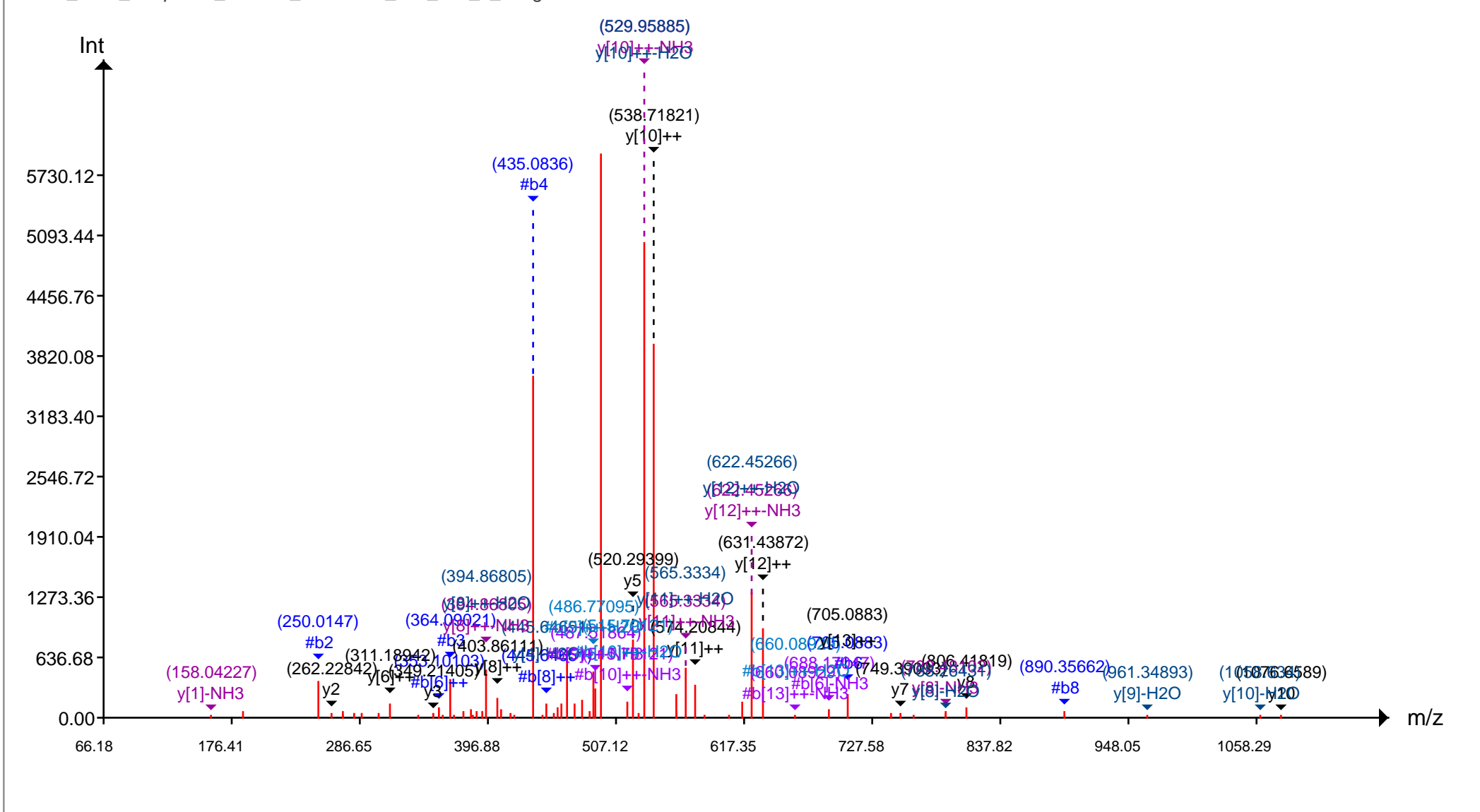
Charge 2+

Ion score 45.09

Confidence level 99.00%

Identity Threshold 44.30

Mass error 7.13E-4



Info

Modified Sequence AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR-COOH

Ion score 59.98

Protein P23588

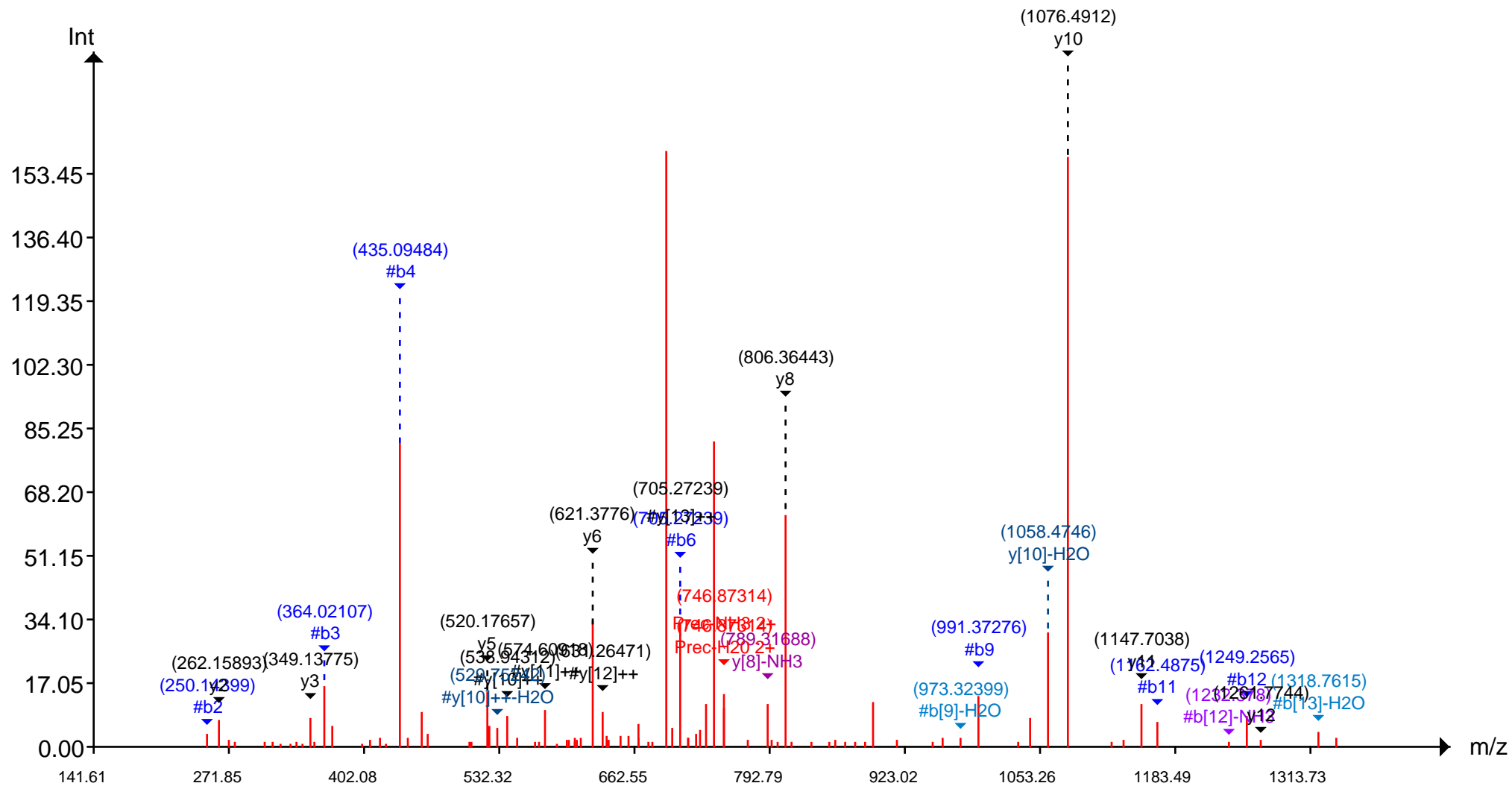
Confidence level 99.00%

Precursor m/z 504.24422

Identity Threshold 47.76

Charge 3+

Mass error -4.26E-4



Info

Modified Sequence AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR-COOH

Ion score 58.83

Protein P23588

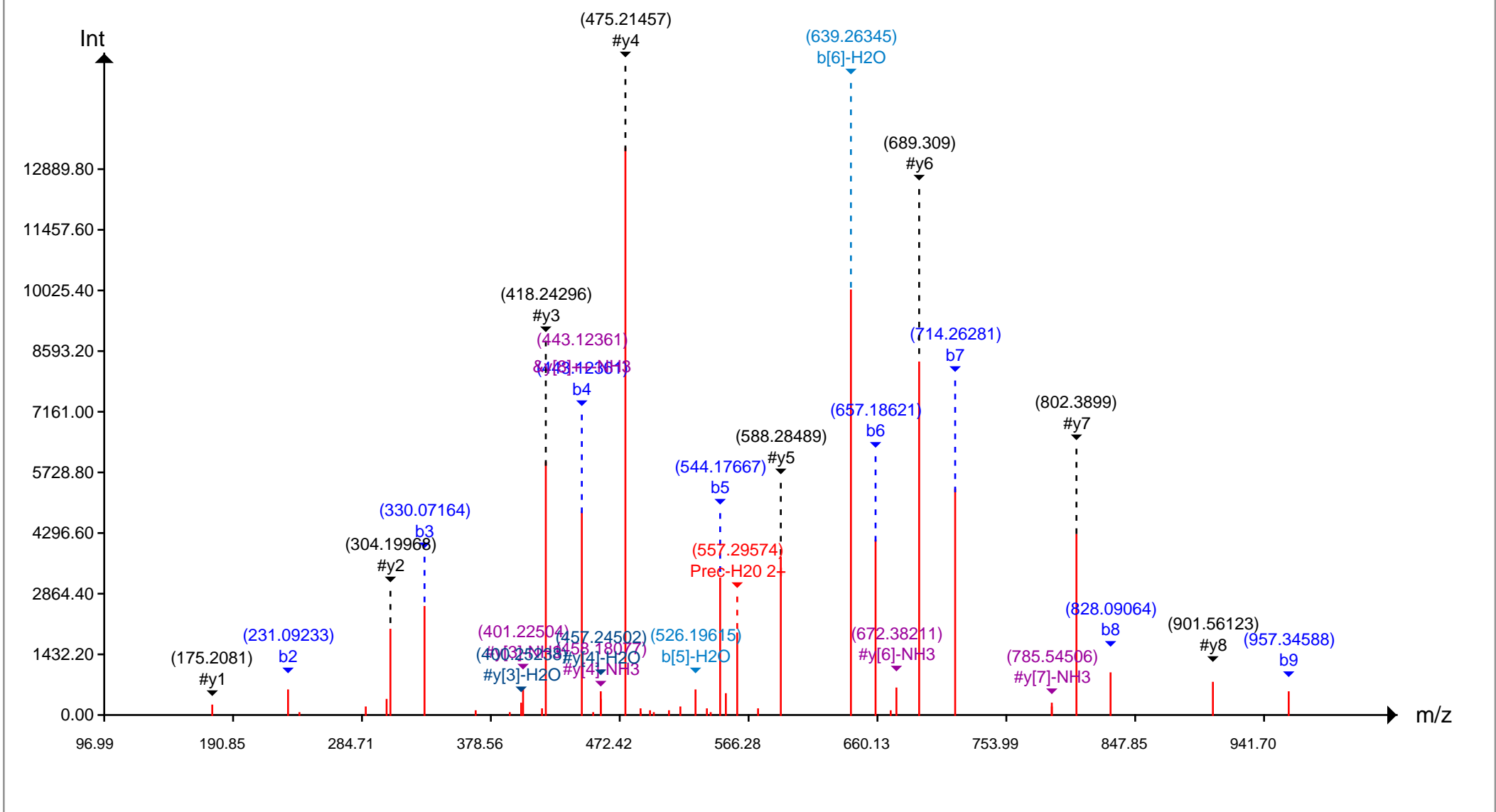
Confidence level 99.00%

Precursor m/z 755.86321

Identity Threshold 47.75

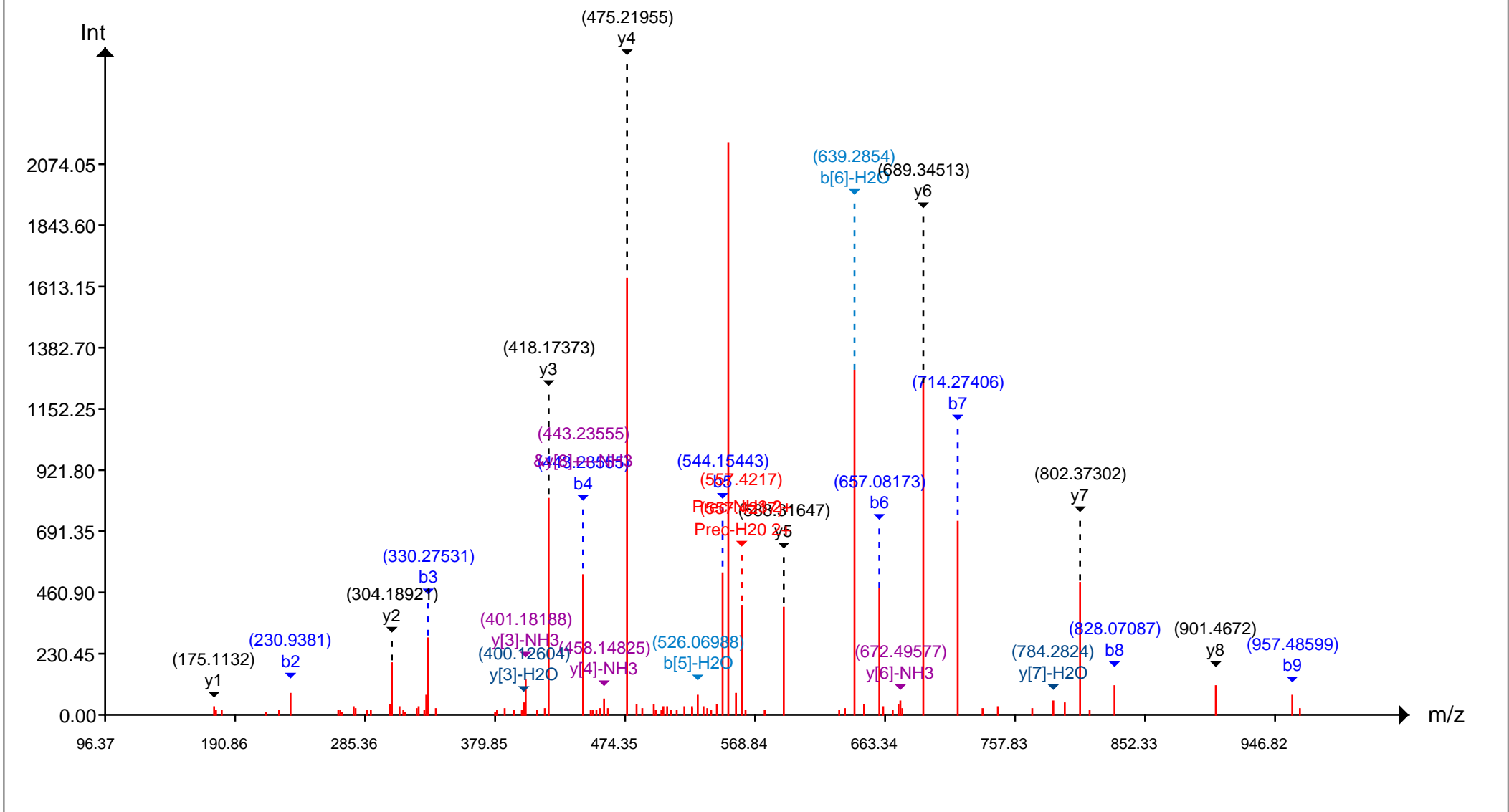
Charge 2+

Mass error 6.1E-4



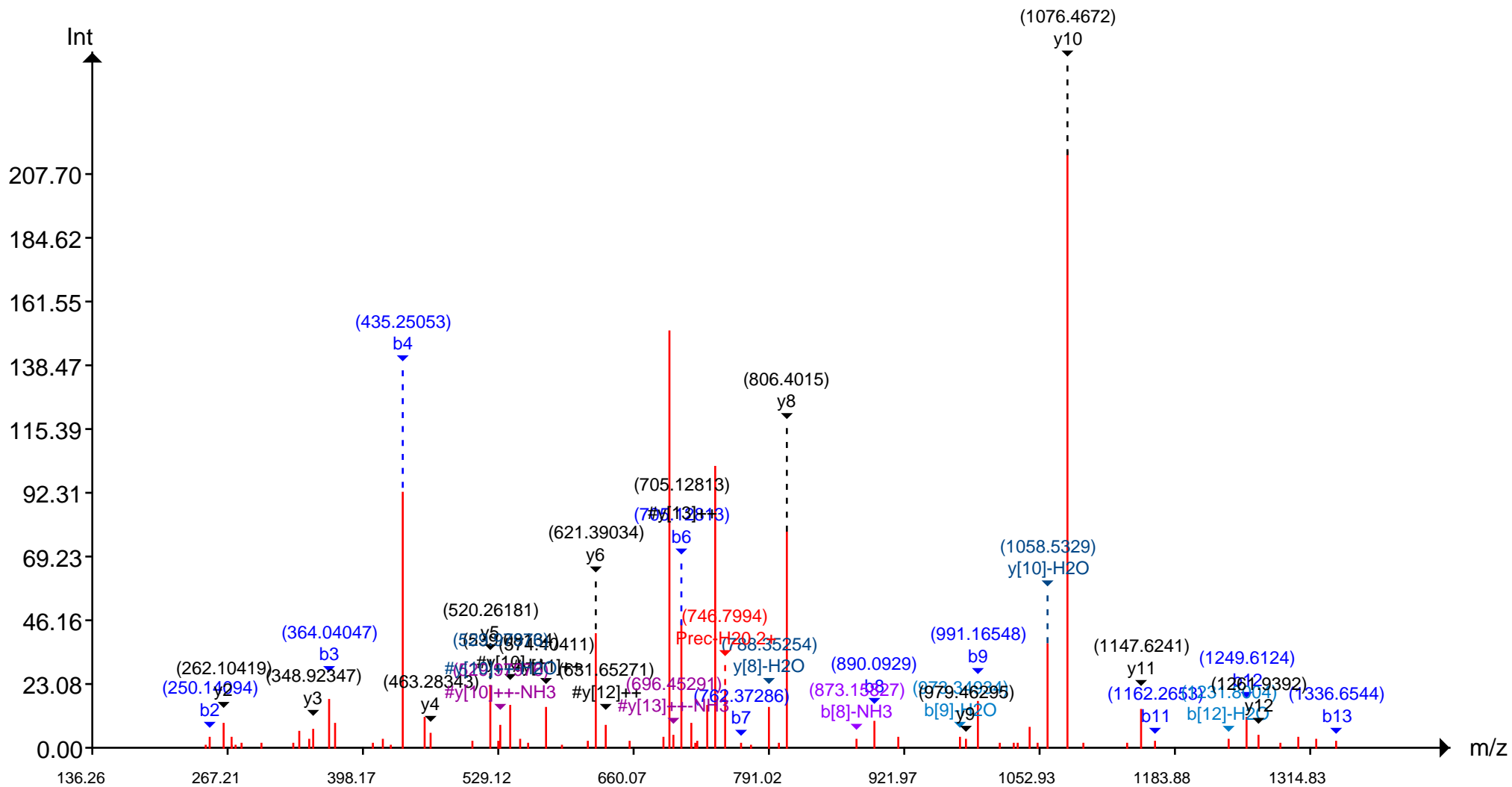
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 92.1 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31347 | Identity Threshold | 48.85 |
| Charge | 2+ | Mass error | 8.3E-5 |



Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 81.46 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31347 | Identity Threshold | 48.85 |
| Charge | 2+ | Mass error | 8.3E-5 |



Info

Modified Sequence AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR-COOH

Ion score 73.51

Protein P23588

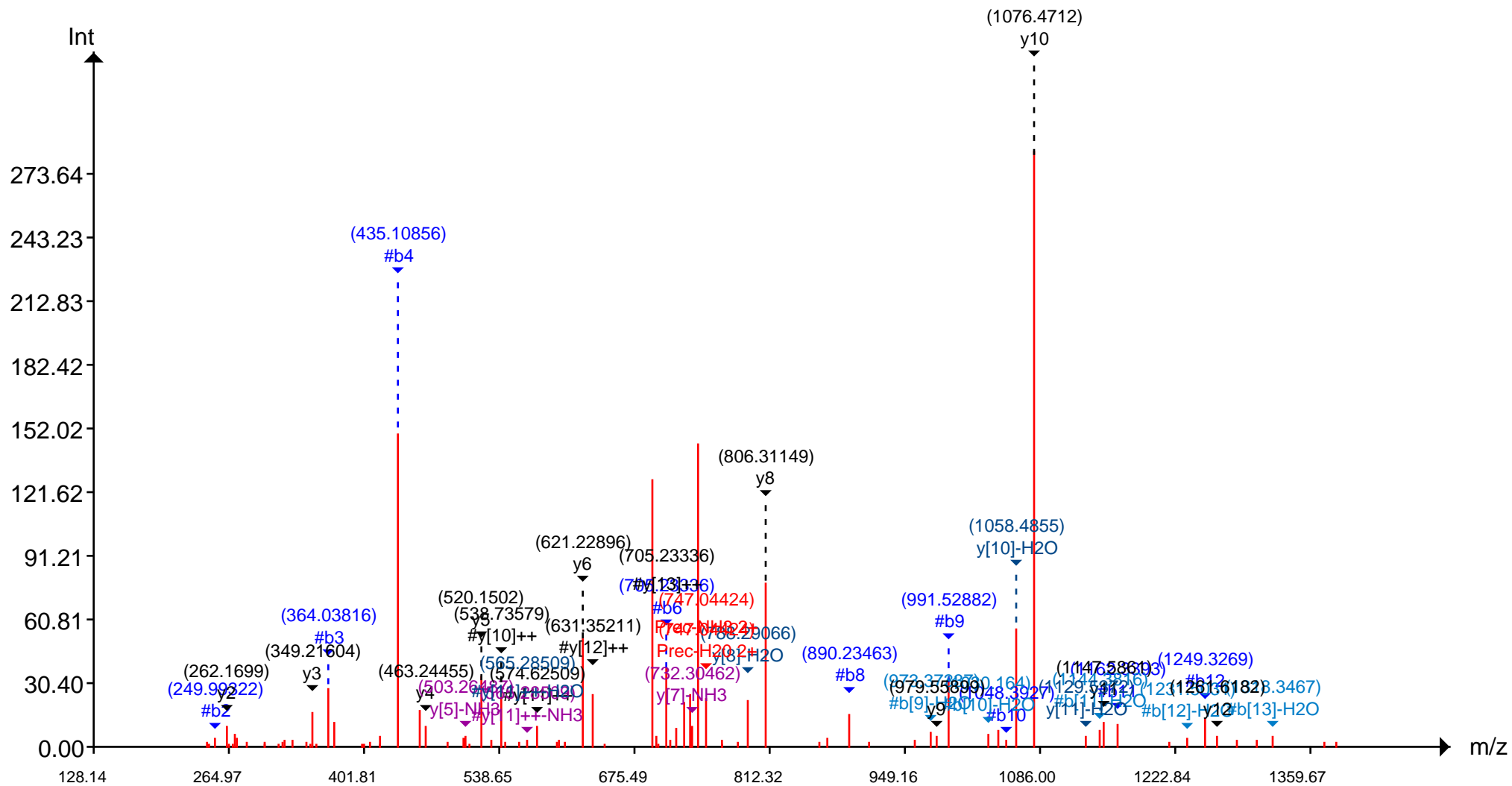
Confidence level 99.00%

Precursor m/z 755.86326

Identity Threshold 47.74

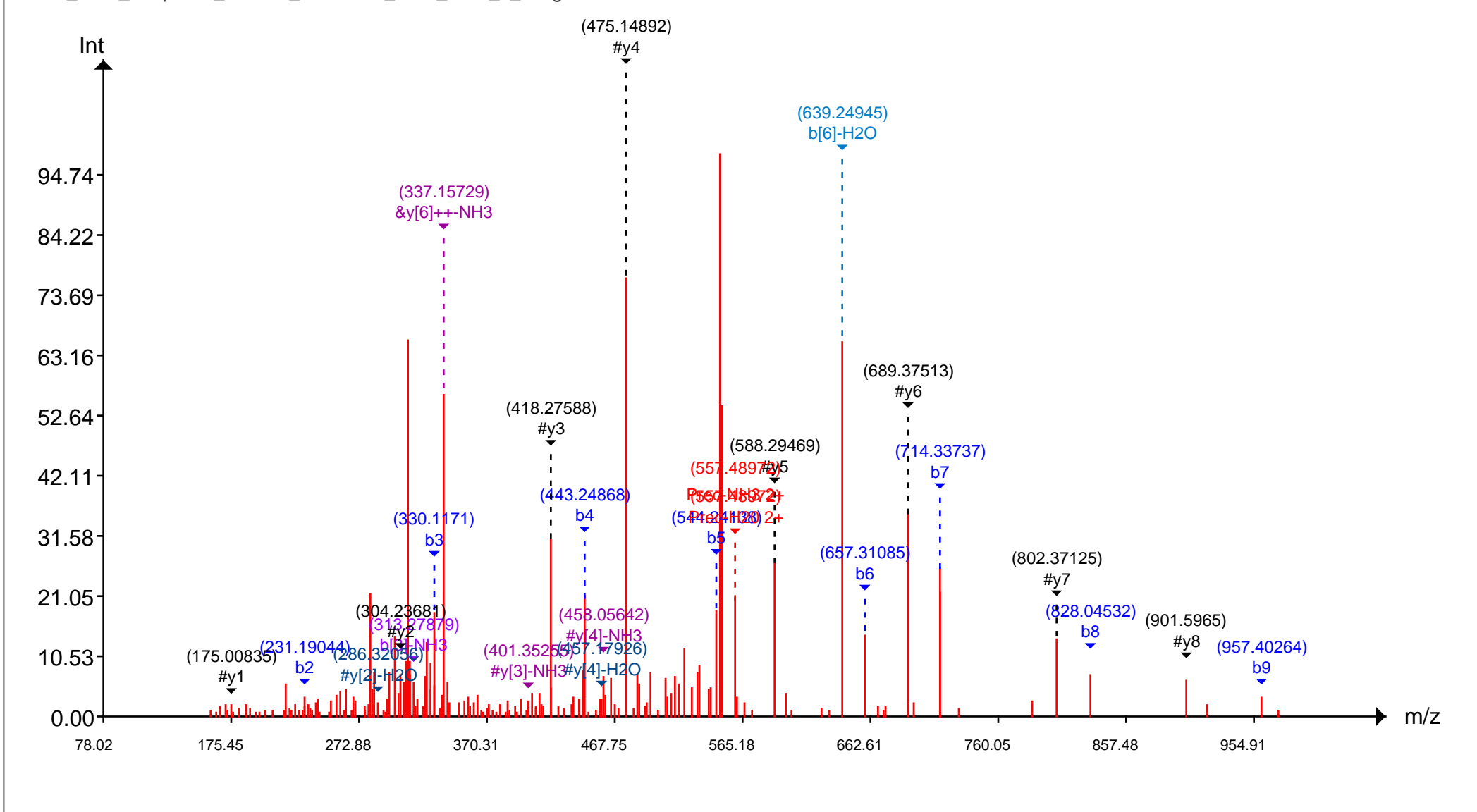
Charge 2+

Mass error 7.1E-4



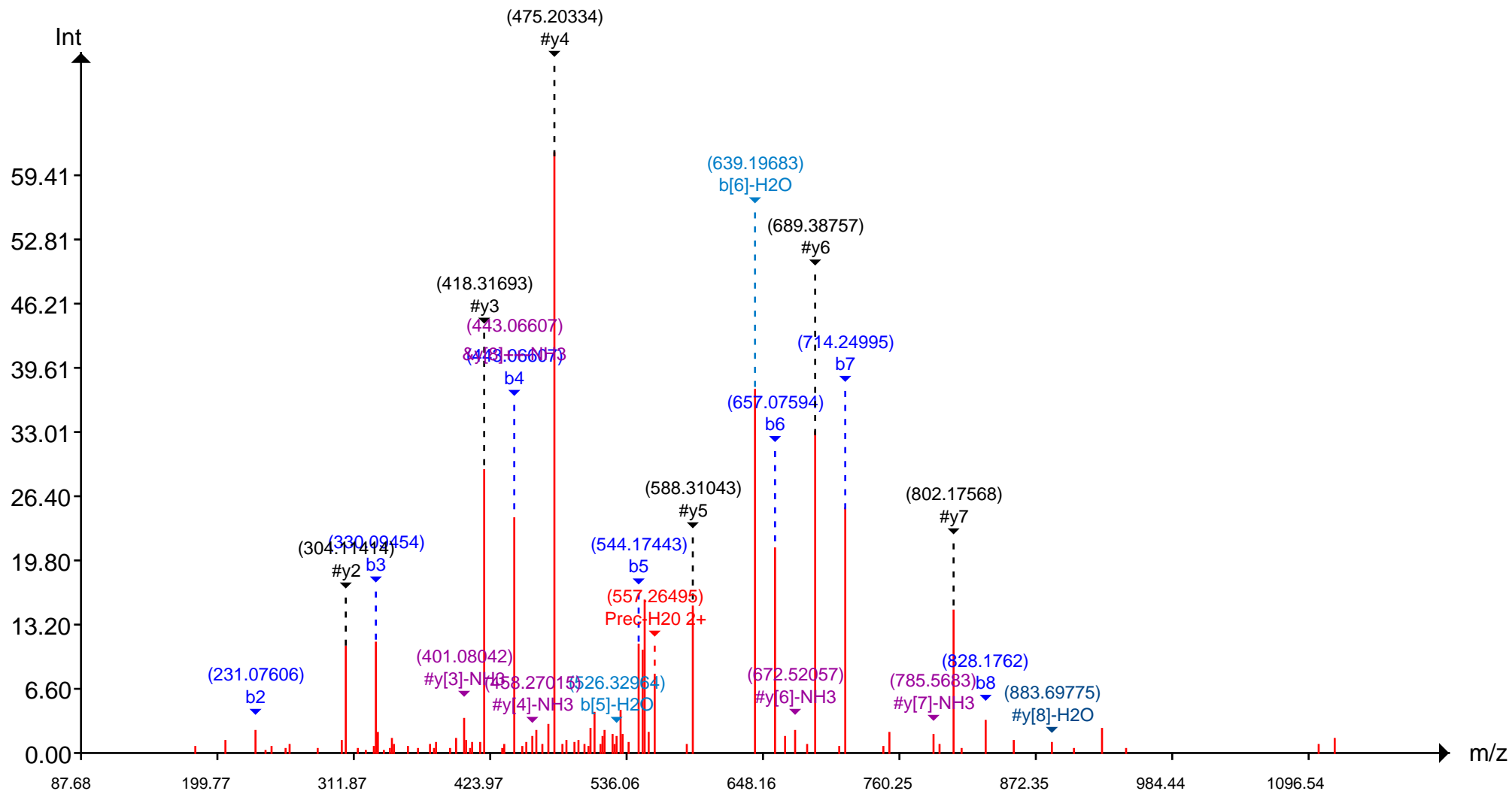
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR-COOH | Ion score | 67.87 |
| Protein | P23588 | Confidence level | 99.00% |
| Precursor m/z | 755.86299 | Identity Threshold | 47.69 |
| Charge | 2+ | Mass error | 1.7E-4 |



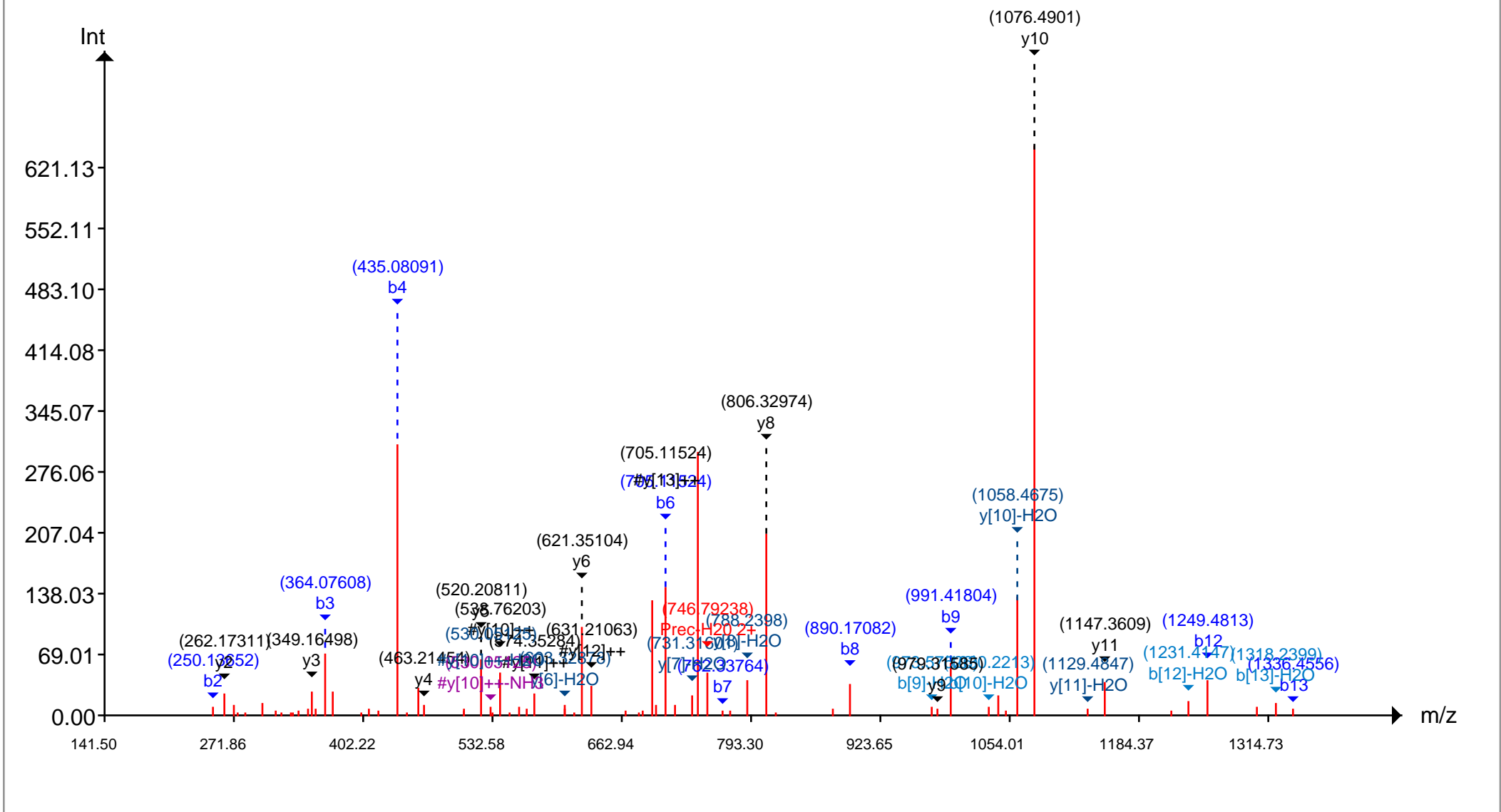
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 50.15 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31347 | Identity Threshold | 48.85 |
| Charge | 2+ | Mass error | 8.3E-5 |



Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 52.01 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31346 | Identity Threshold | 48.97 |
| Charge | 2+ | Mass error | 6.3E-5 |



Info

Modified Sequence AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR-COOH

Ion score 77.32

Protein P23588

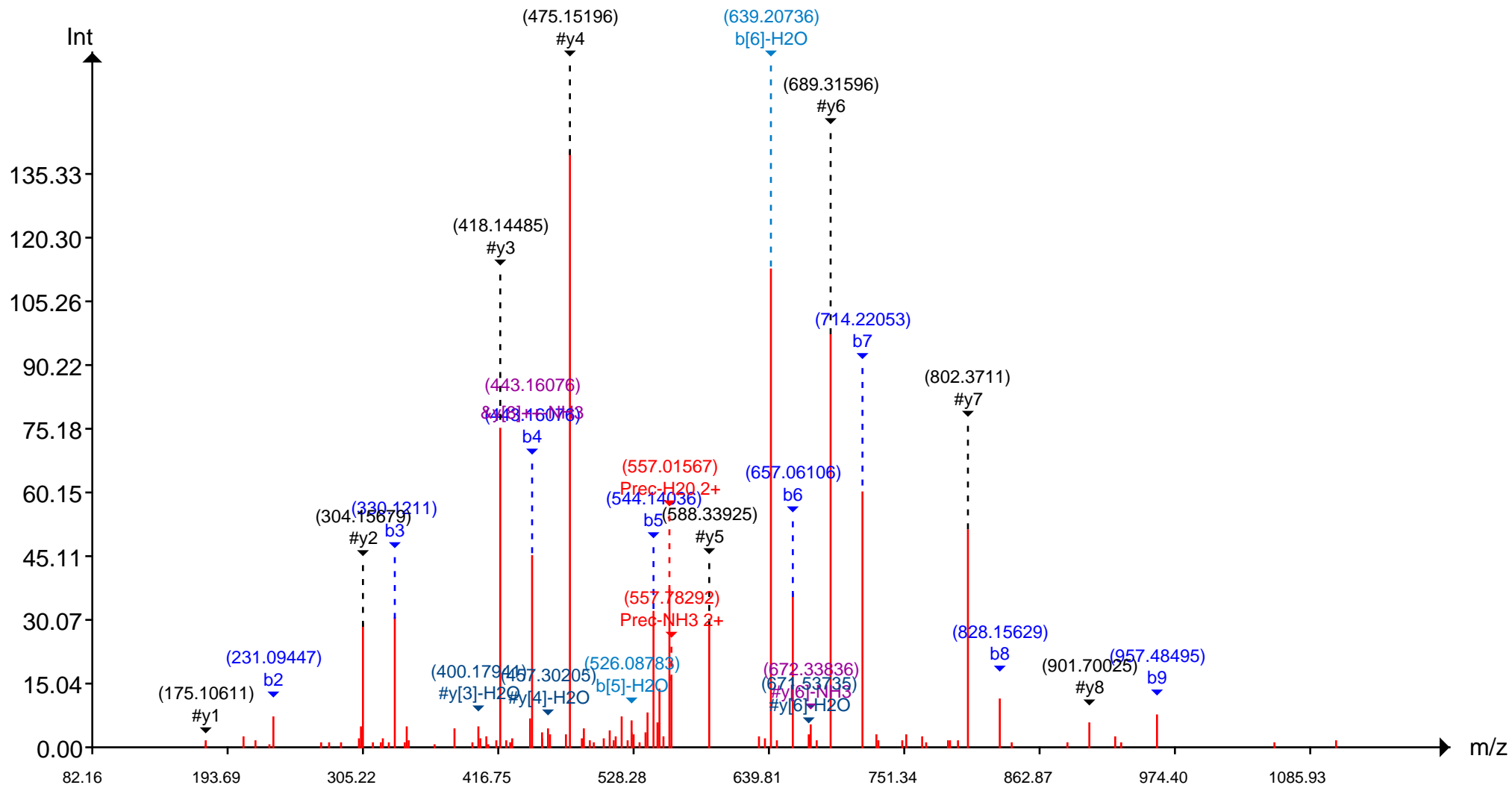
Confidence level 99.00%

Precursor m/z 755.86313

Identity Threshold 47.74

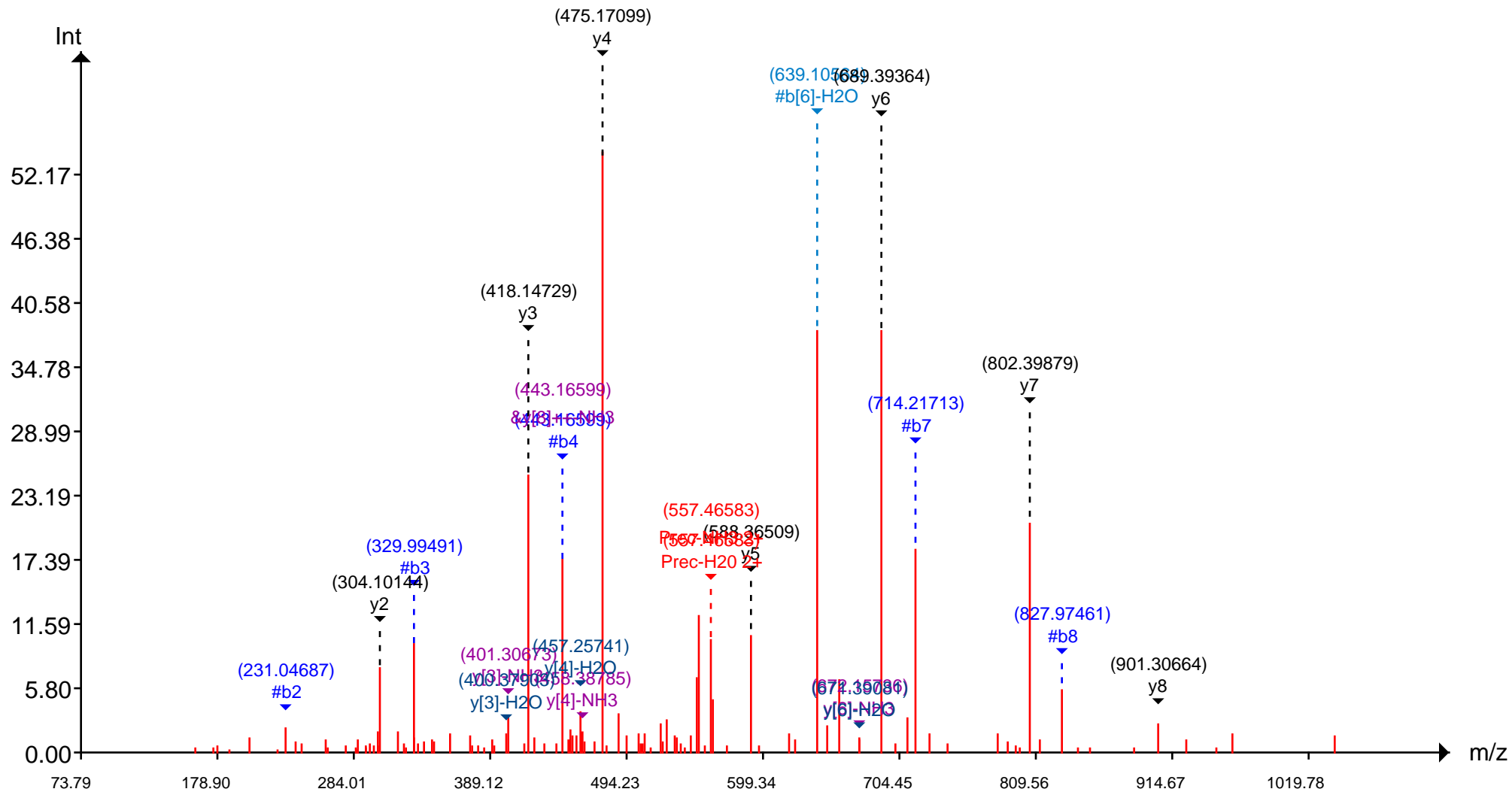
Charge 2+

Mass error 4.5E-4



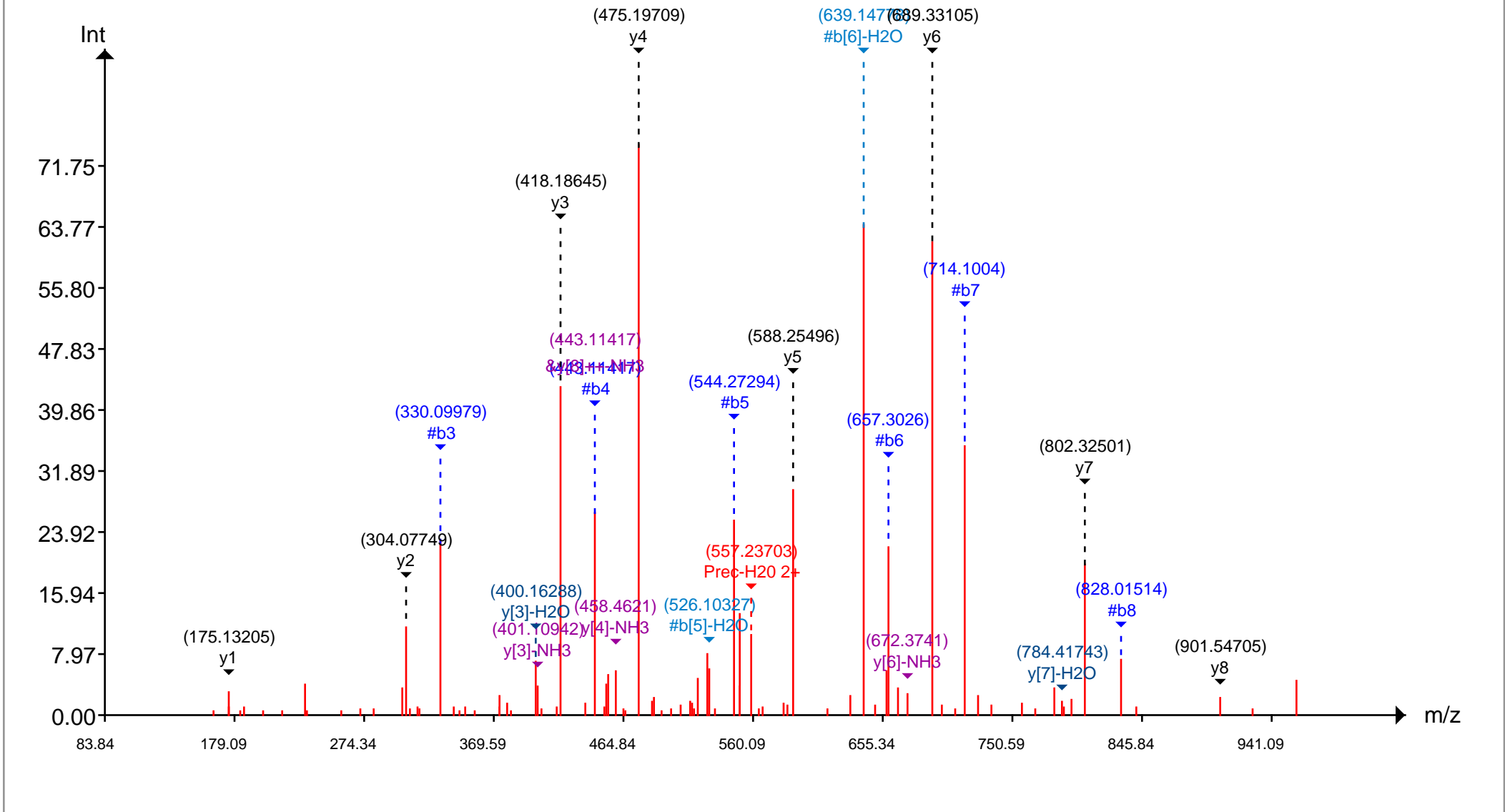
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 66.17 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31346 | Identity Threshold | 48.97 |
| Charge | 2+ | Mass error | 6.3E-5 |



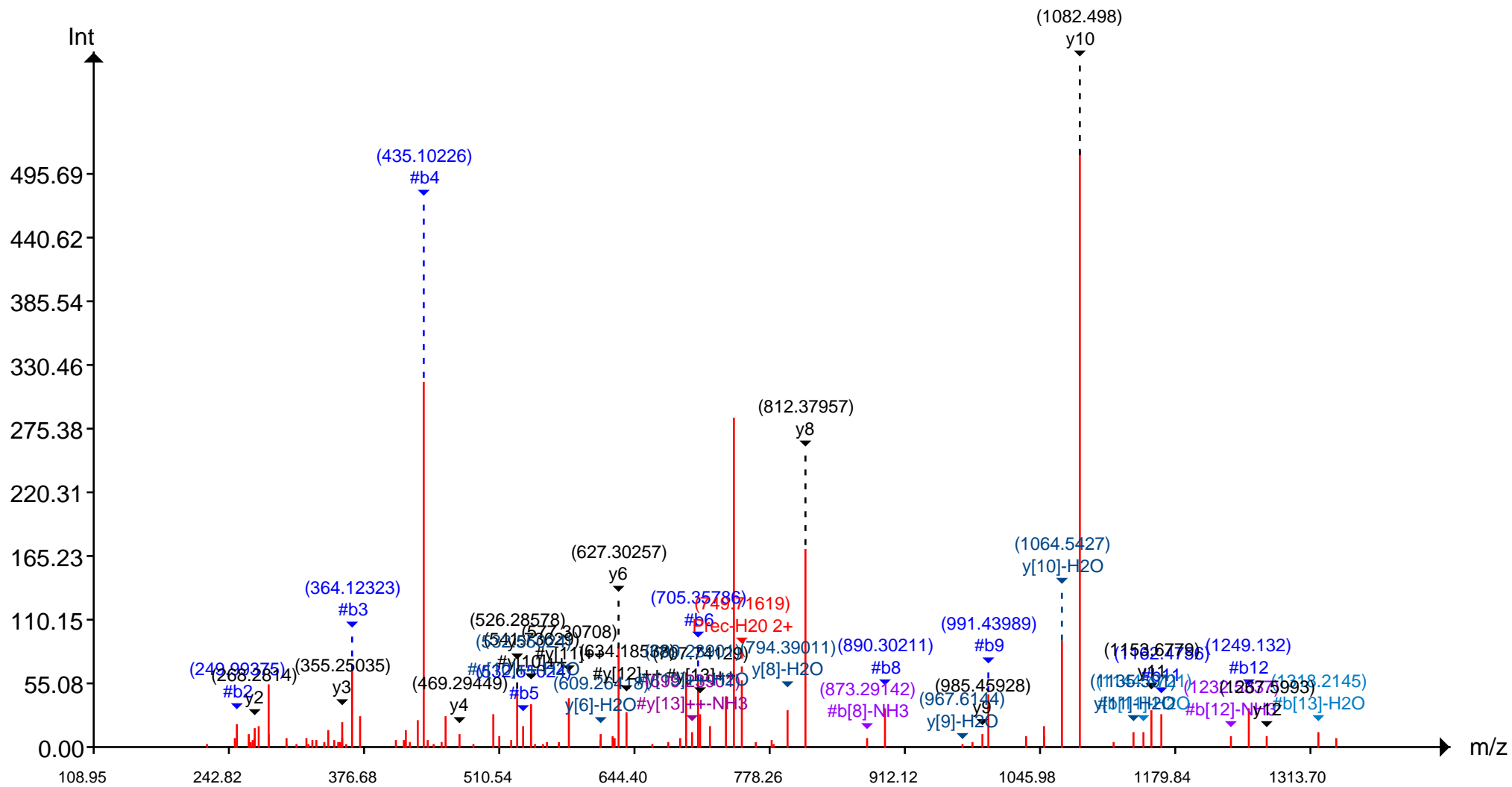
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 64.18 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31346 | Identity Threshold | 48.97 |
| Charge | 2+ | Mass error | 6.3E-5 |

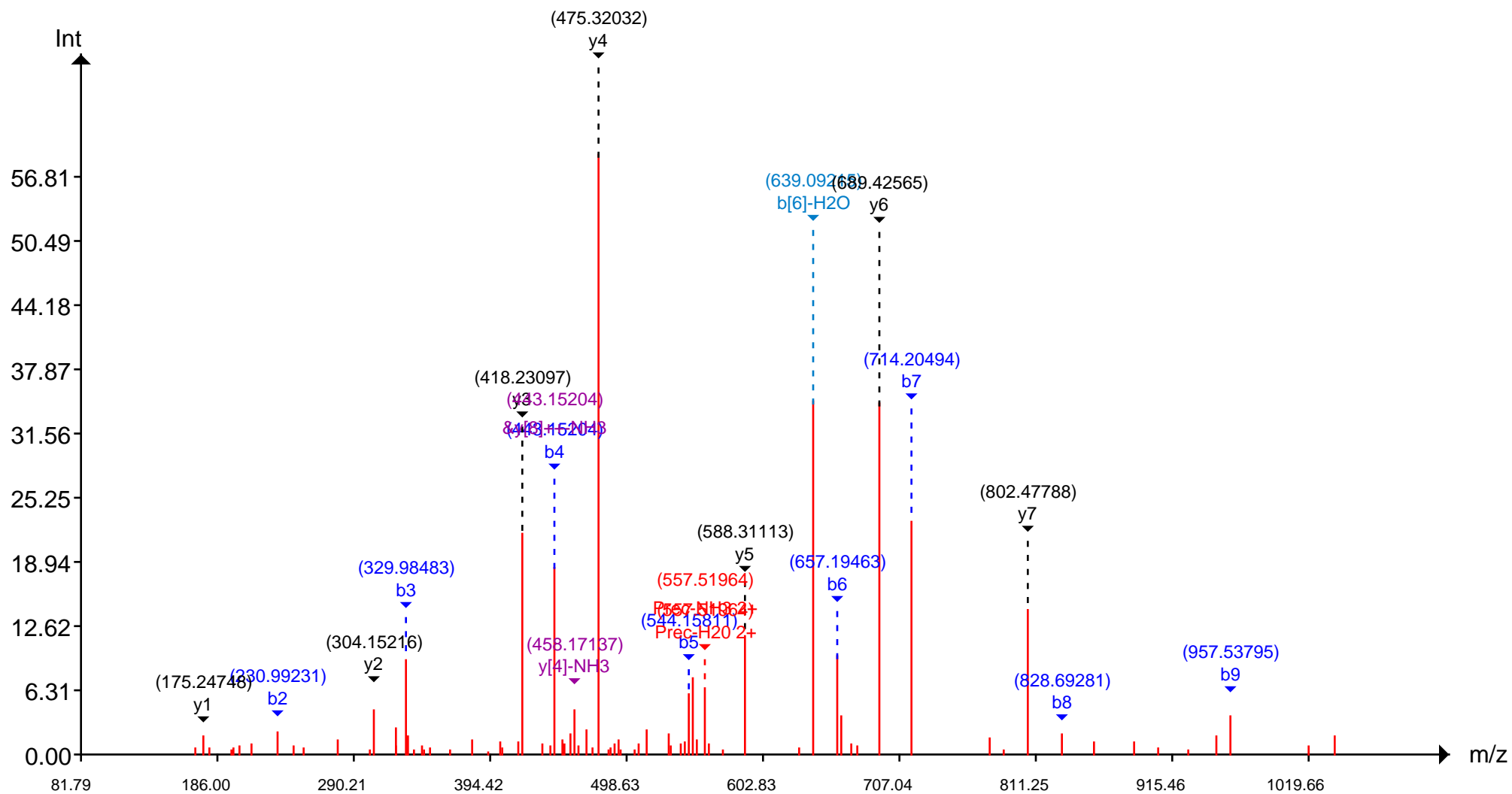


Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 81.05 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31345 | Identity Threshold | 49.03 |
| Charge | 2+ | Mass error | 4.3E-5 |

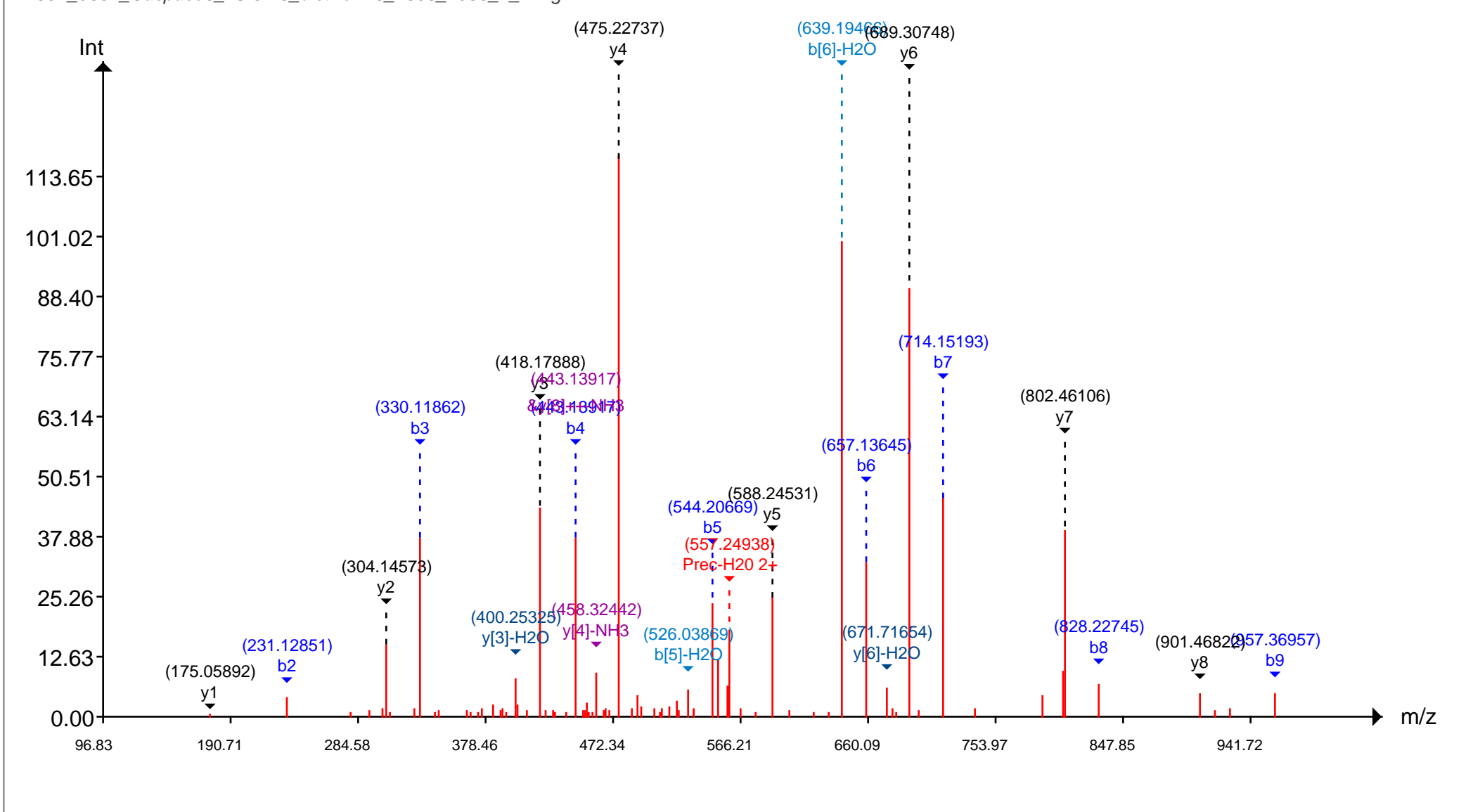


| Info | | Ion score | 72.43 |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR<C13>-COOH | Confidence level | 99.00% |
| Protein | P23588 | Identity Threshold | 48.10 |
| Precursor m/z | 758.87308 | Mass error | 2.08E-4 |
| Charge | 2+ | | |



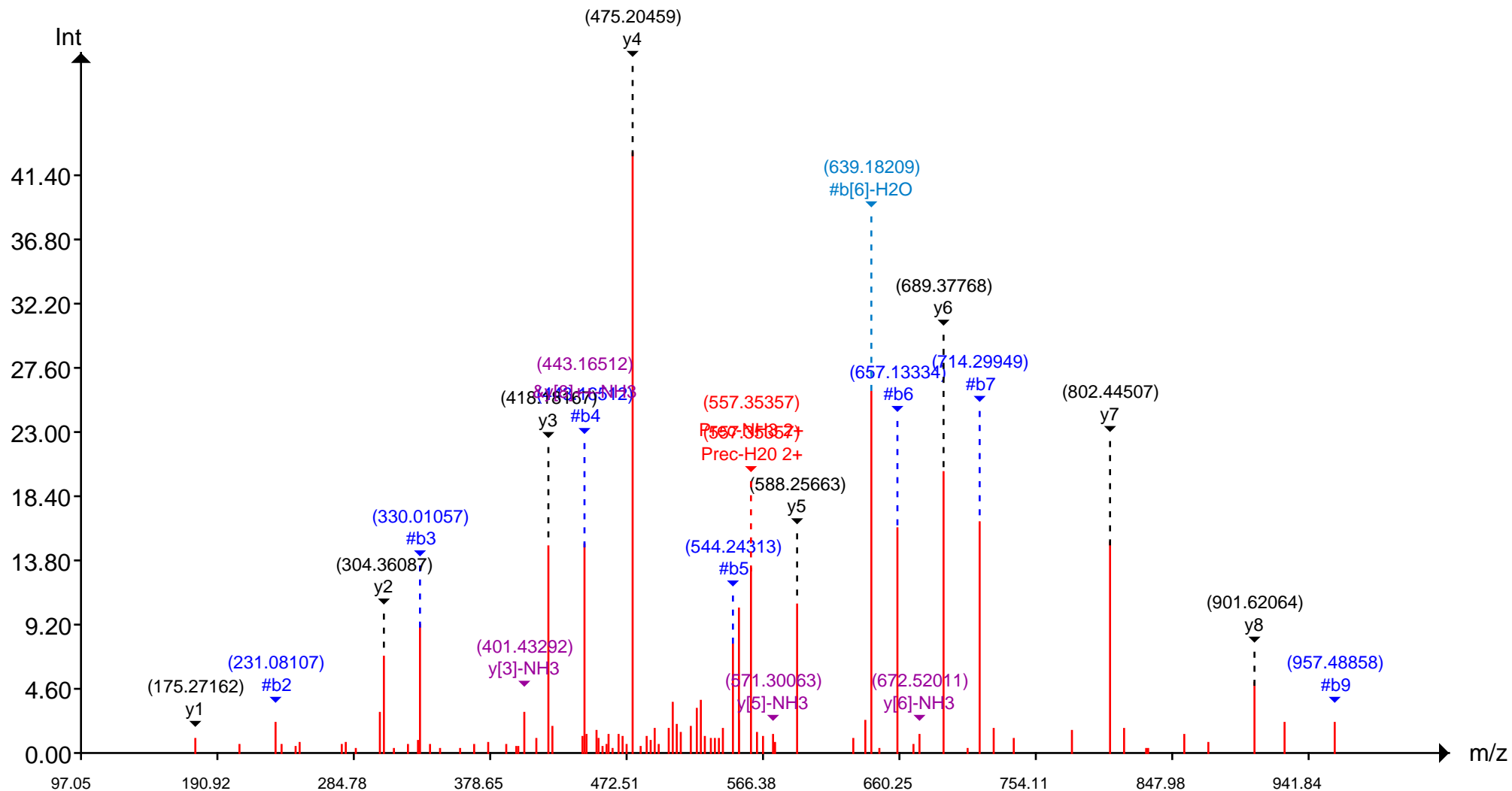
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 67.29 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31342 | Identity Threshold | 49.03 |
| Charge | 2+ | Mass error | -1.7E-5 |



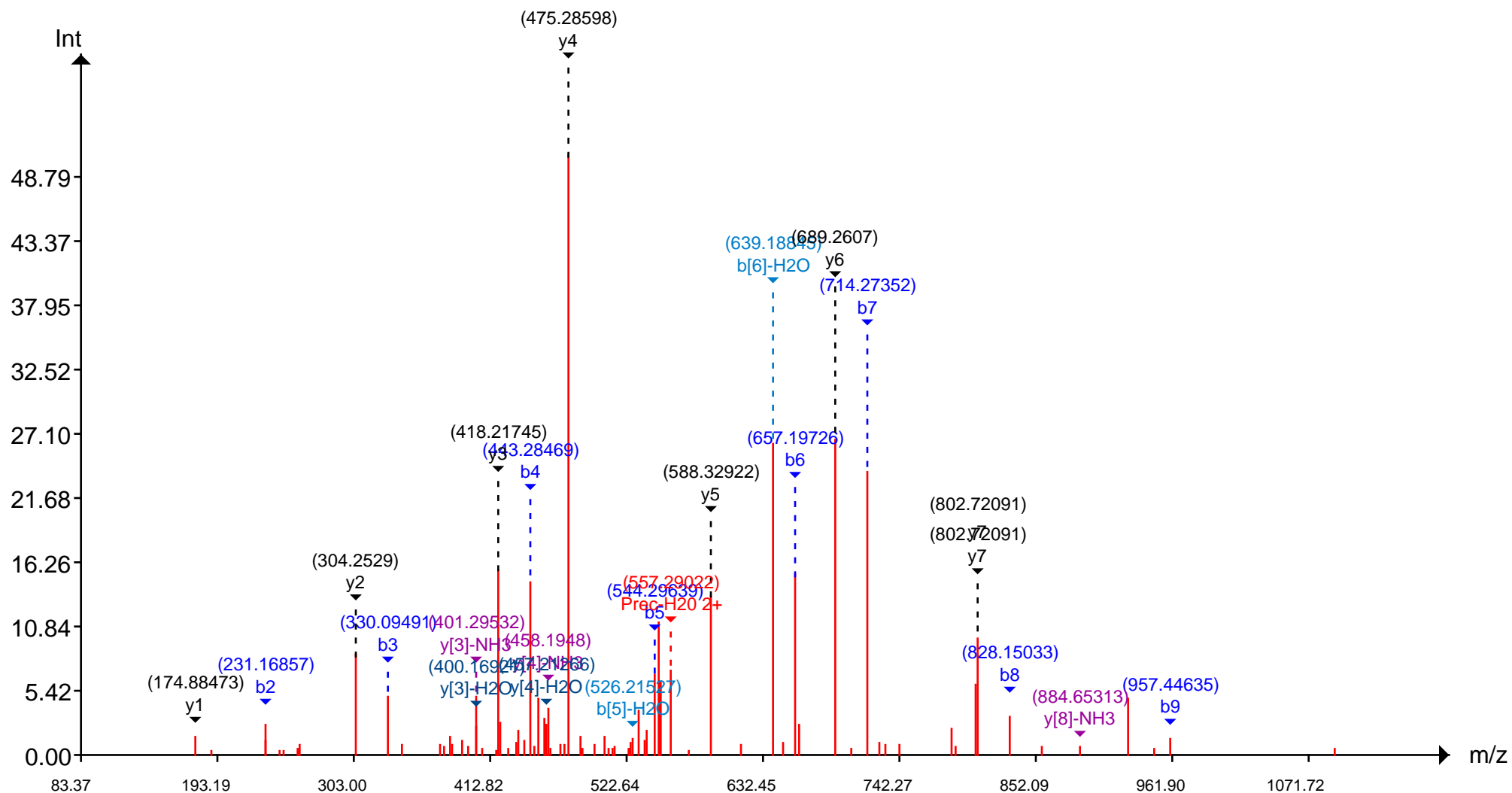
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 74.99 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31336 | Identity Threshold | 48.86 |
| Charge | 2+ | Mass error | -1.37E-4 |



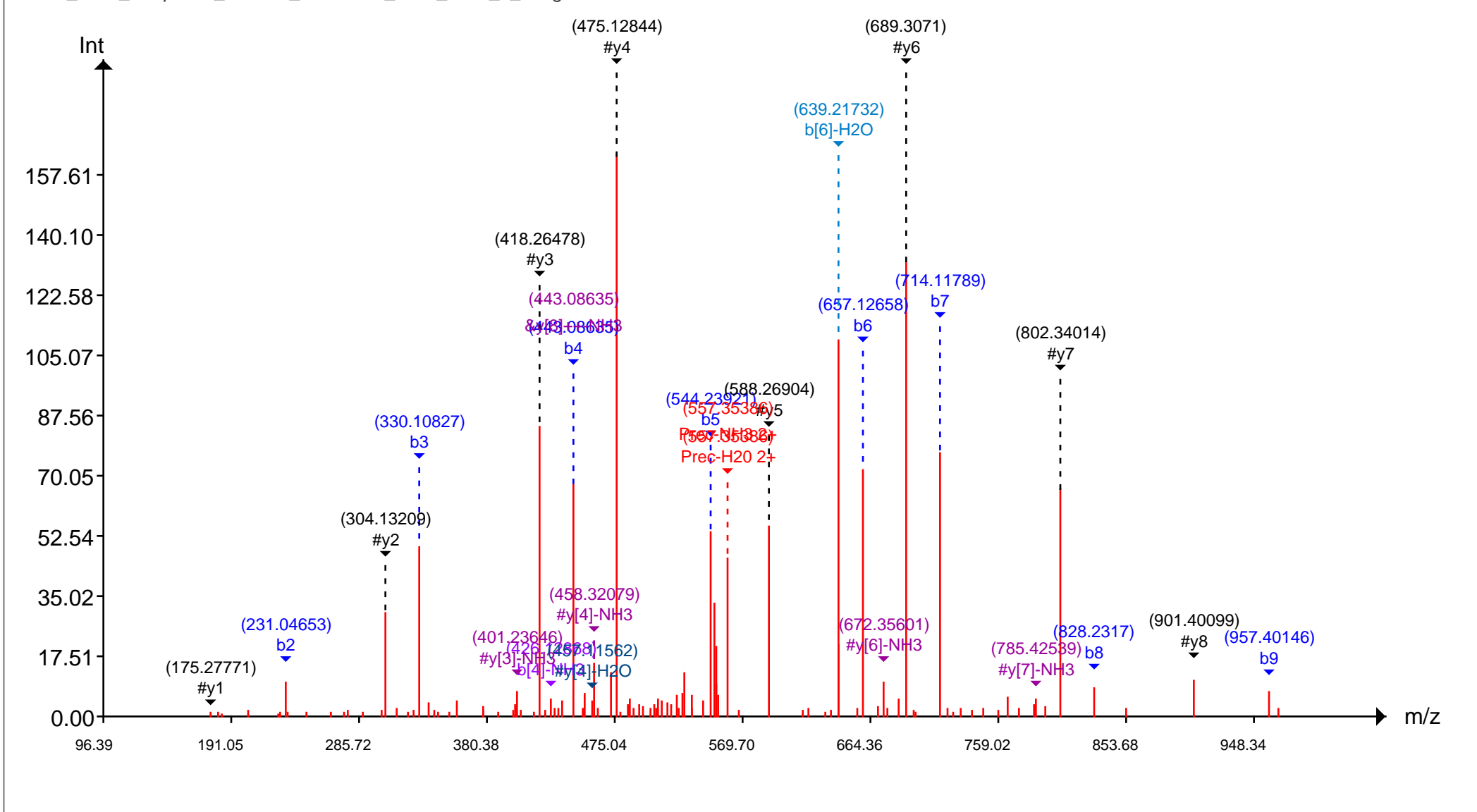
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 67.76 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31335 | Identity Threshold | 48.86 |
| Charge | 2+ | Mass error | -1.57E-4 |



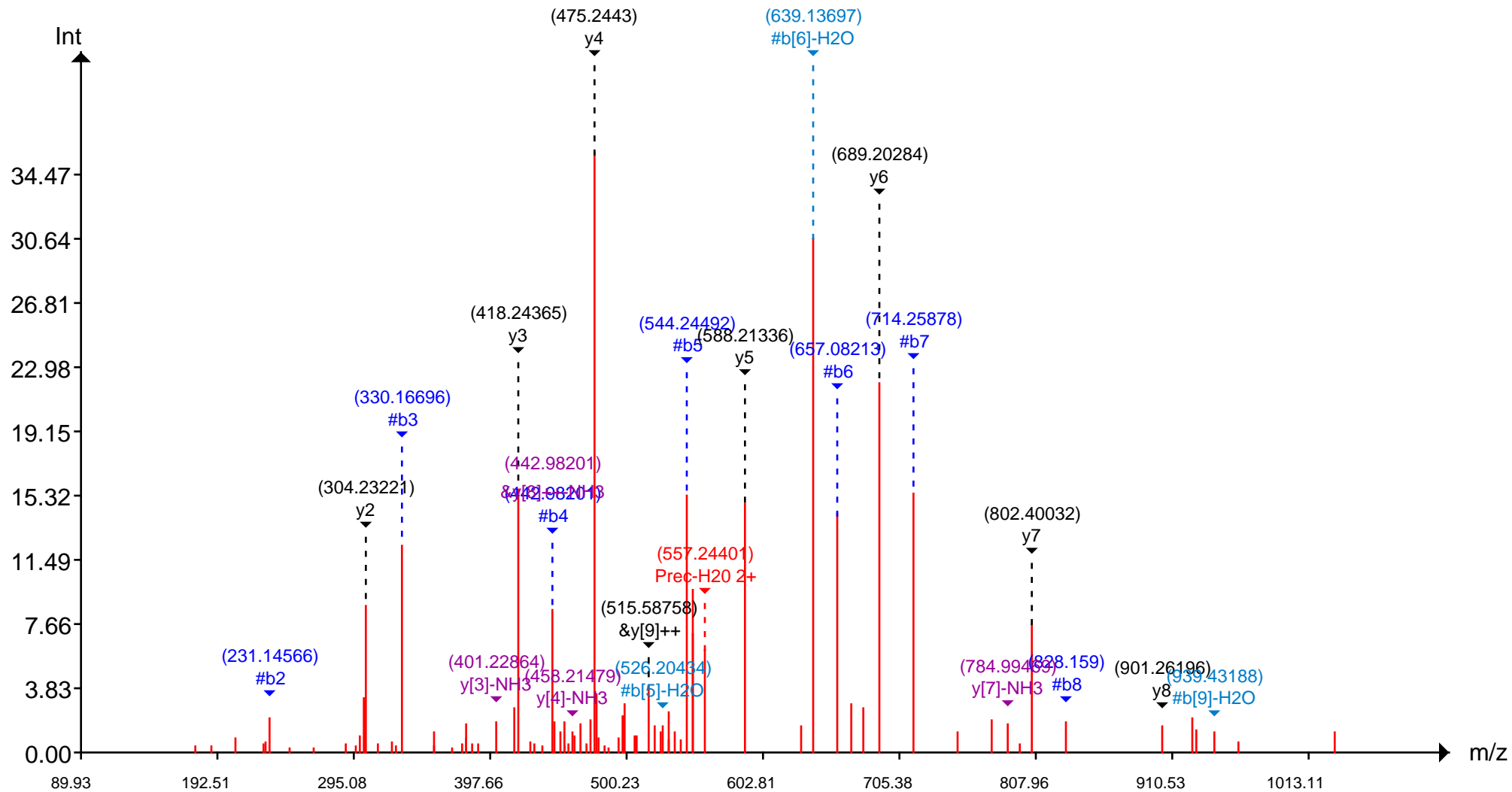
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 62.94 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.3136 | Identity Threshold | 48.77 |
| Charge | 2+ | Mass error | 3.43E-4 |



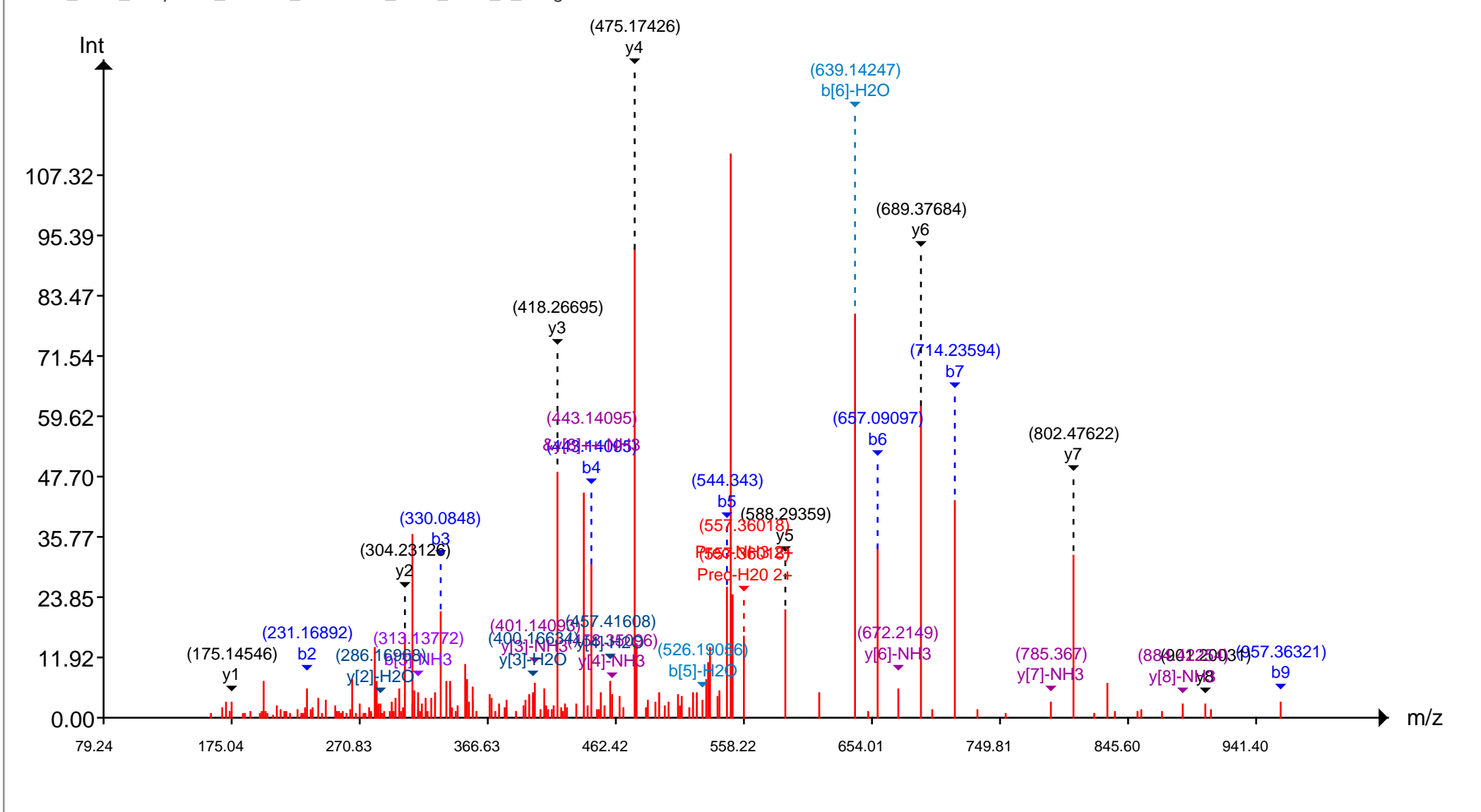
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 70.25 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31353 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 2.03E-4 |



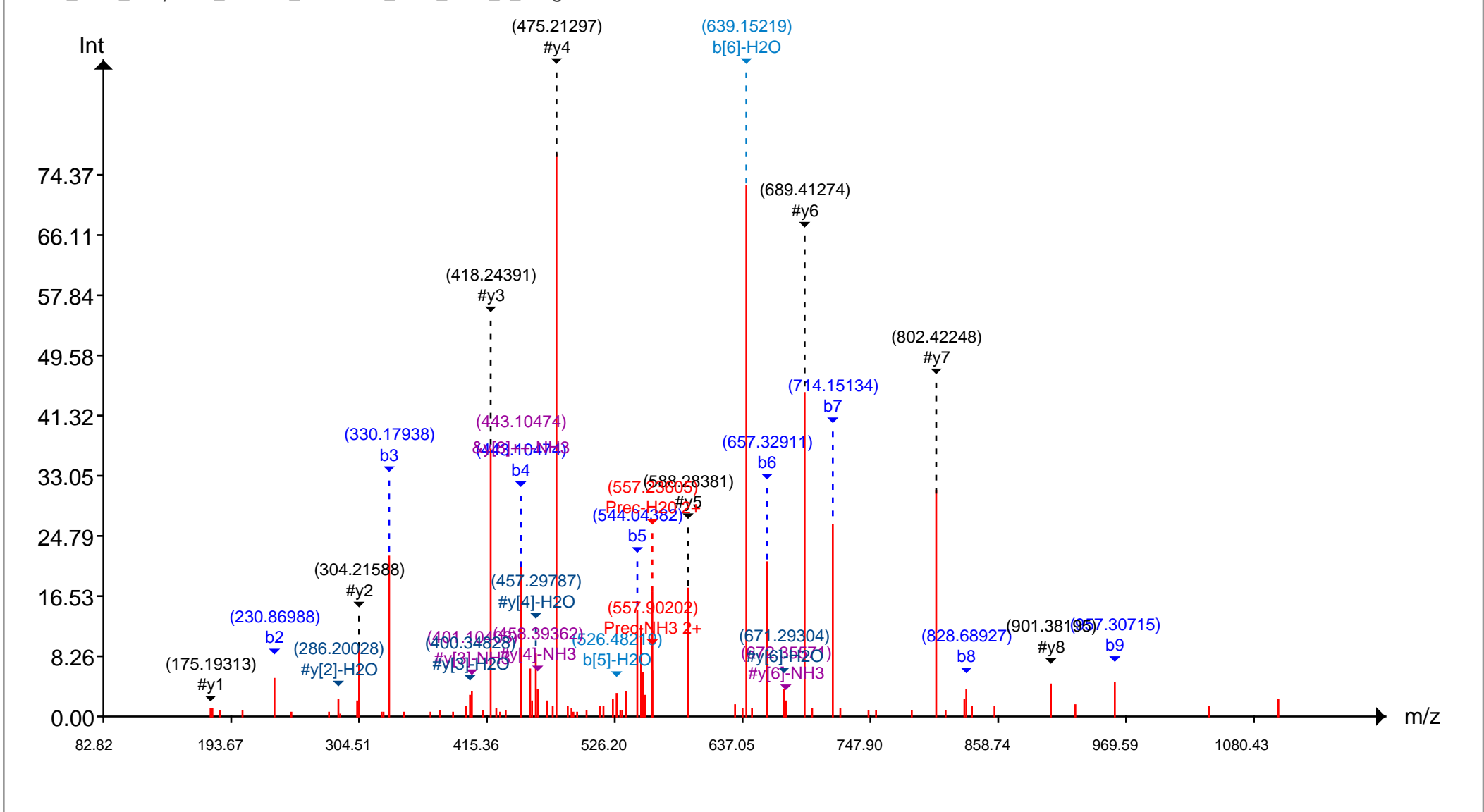
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 63.8 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31352 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 1.83E-4 |



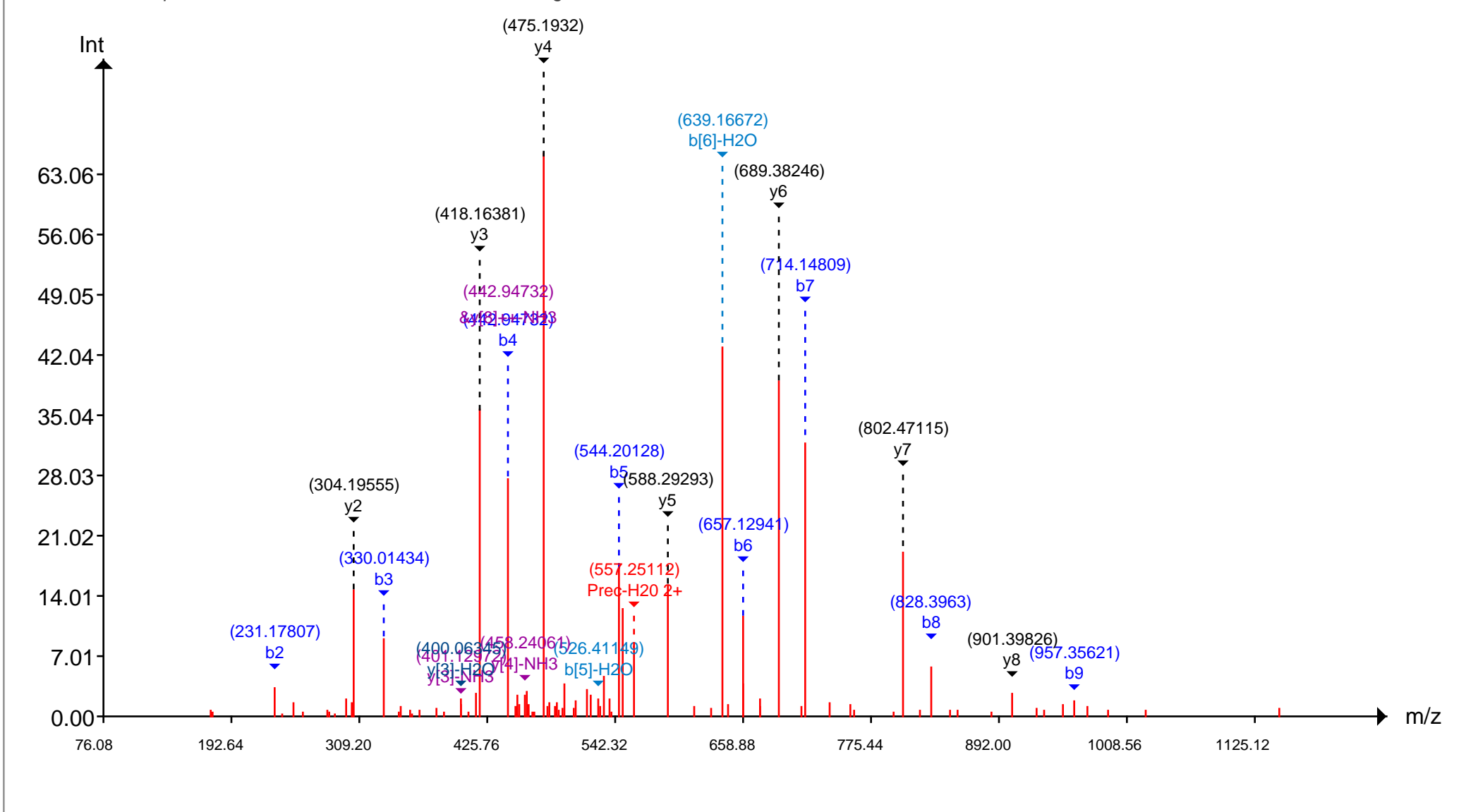
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 61.22 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31353 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 2.03E-4 |



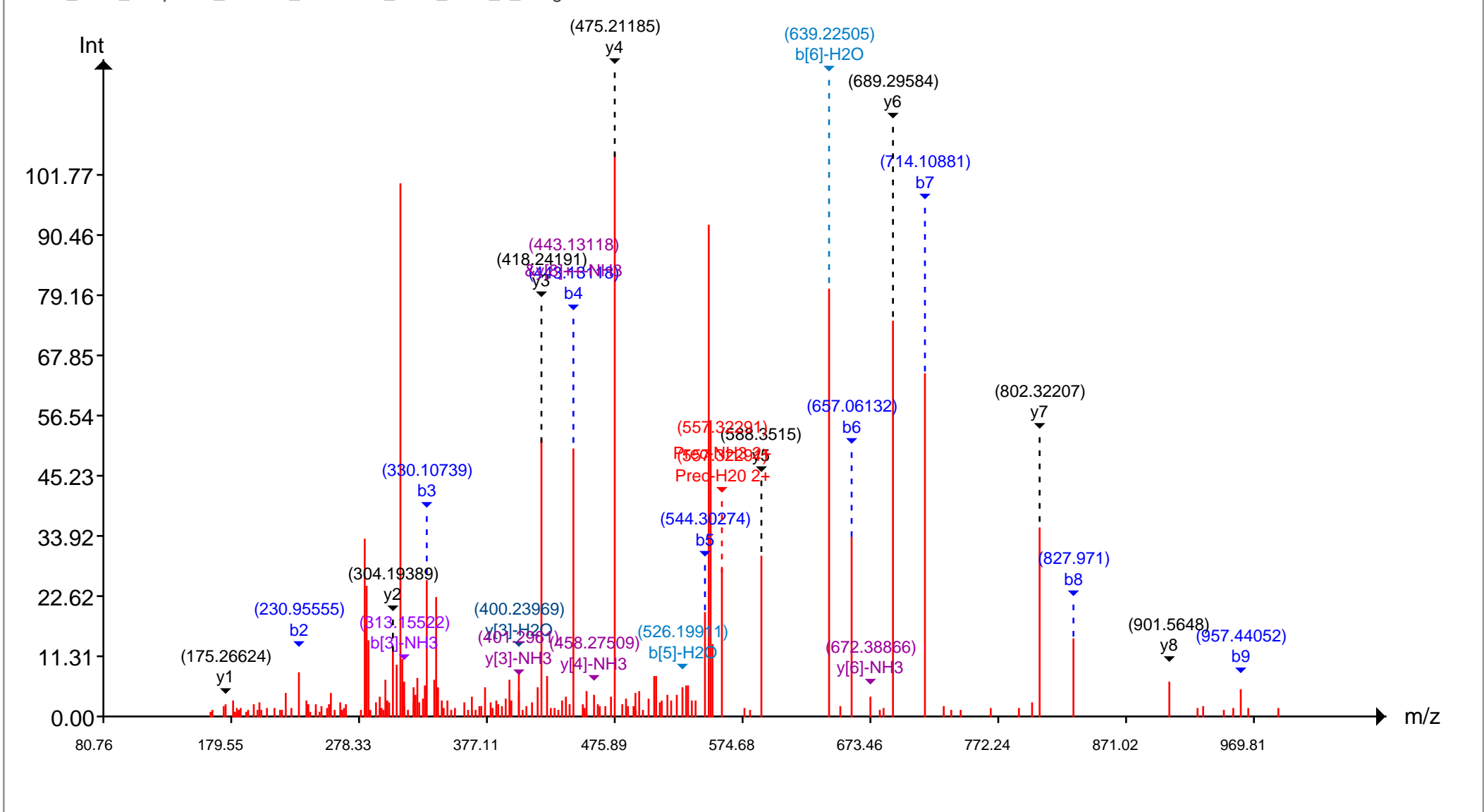
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 68.39 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.3135 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 1.43E-4 |



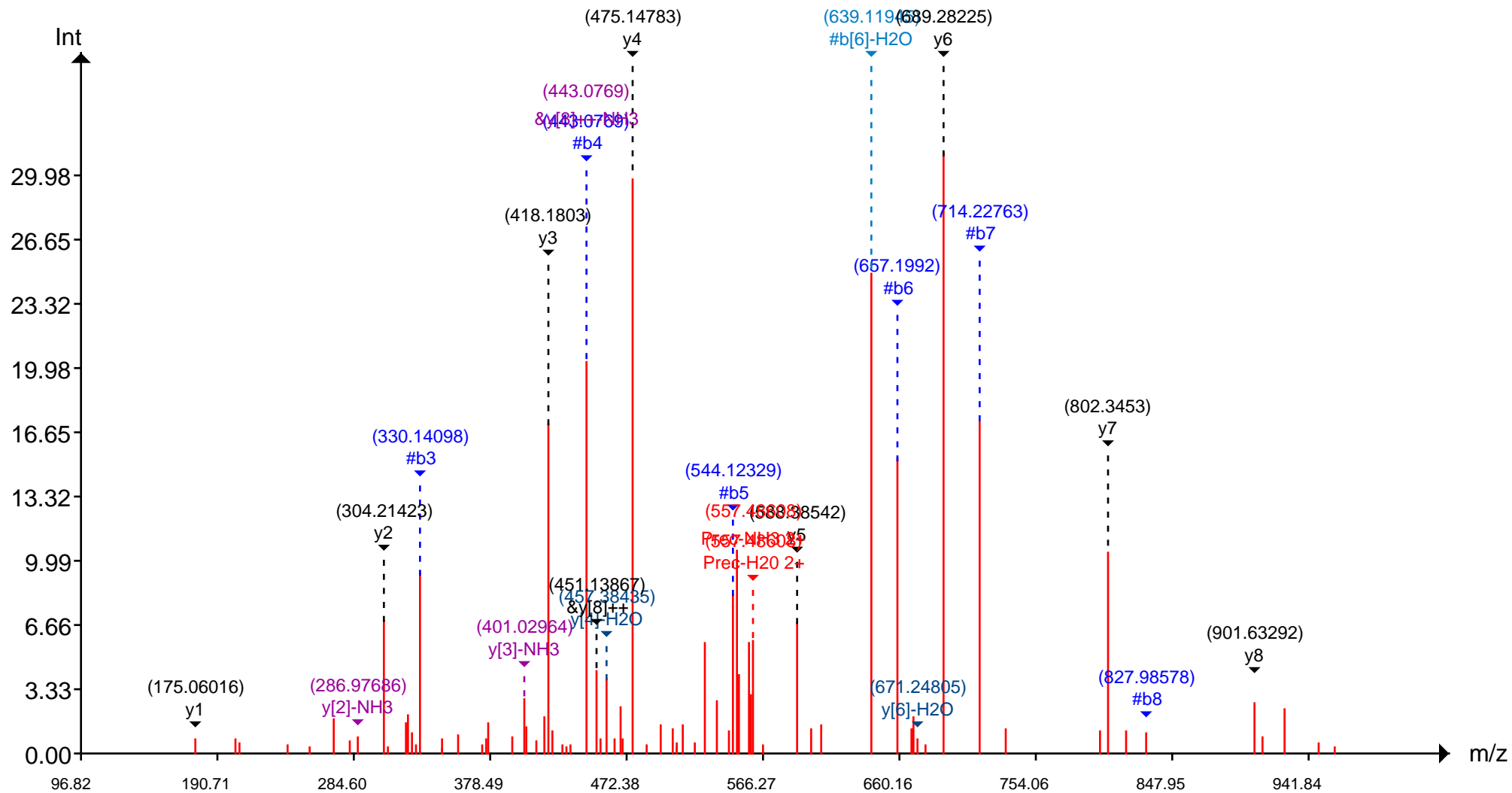
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 71.12 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31352 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 1.83E-4 |



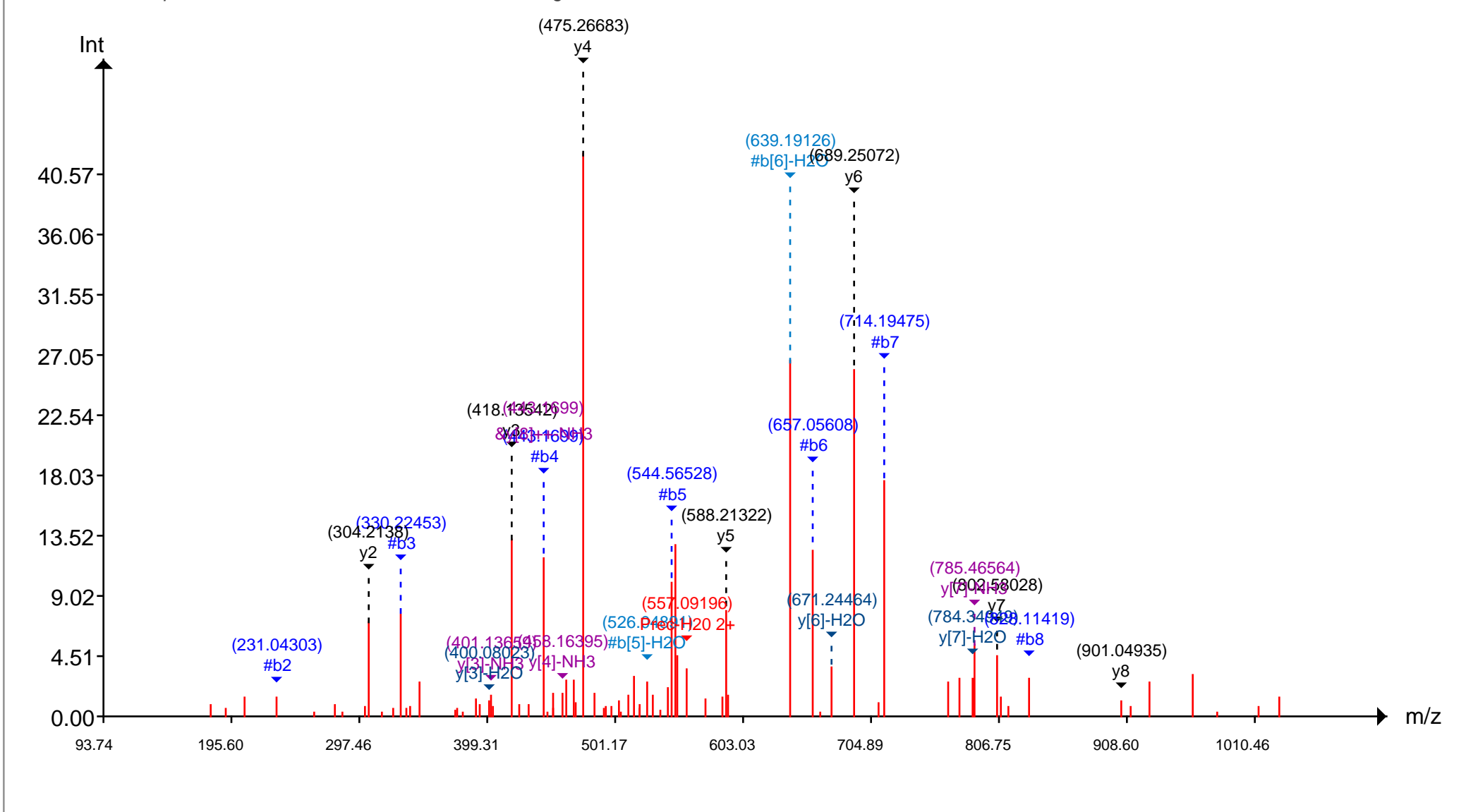
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 51.61 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31348 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 1.03E-4 |



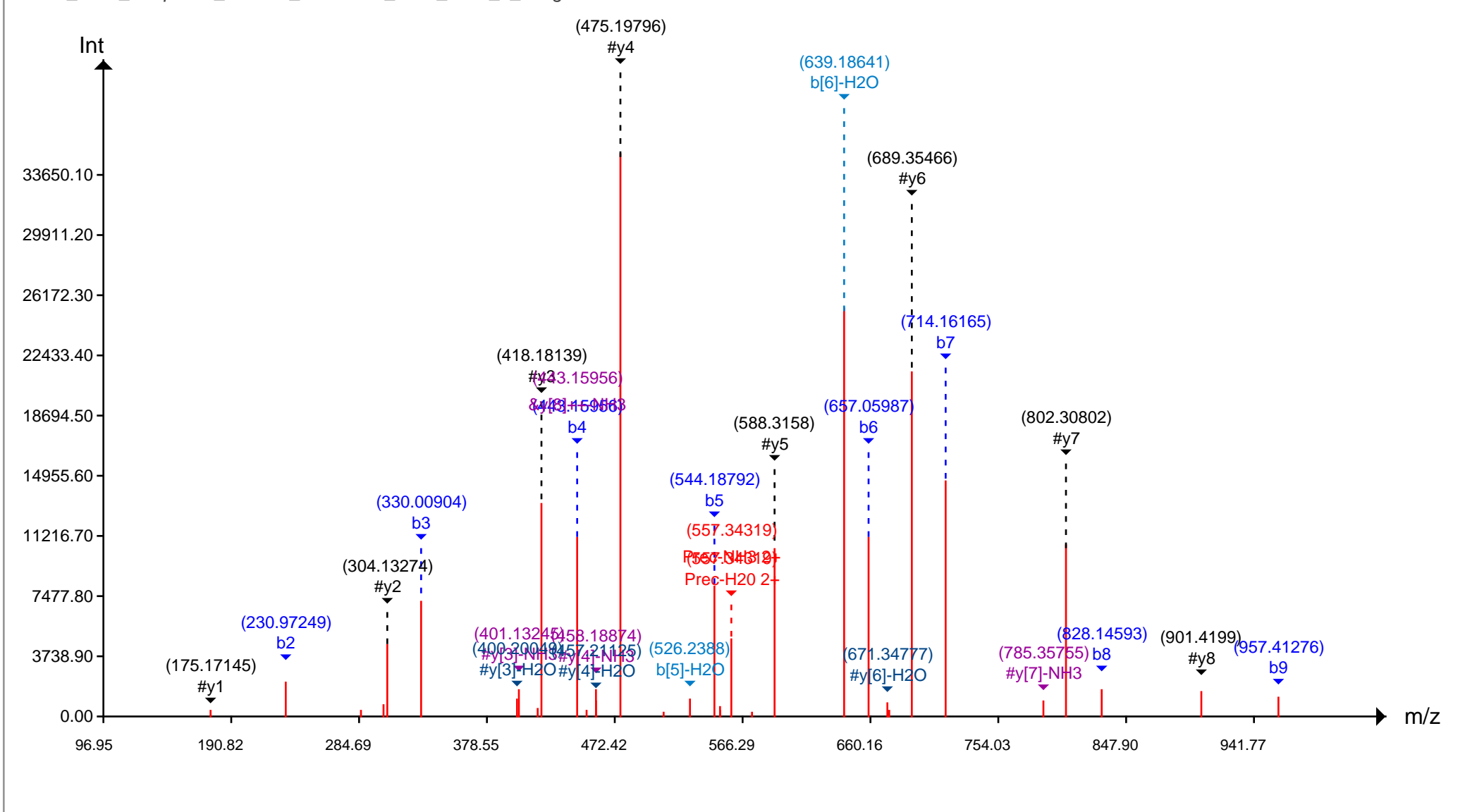
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 60.15 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31348 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 1.03E-4 |



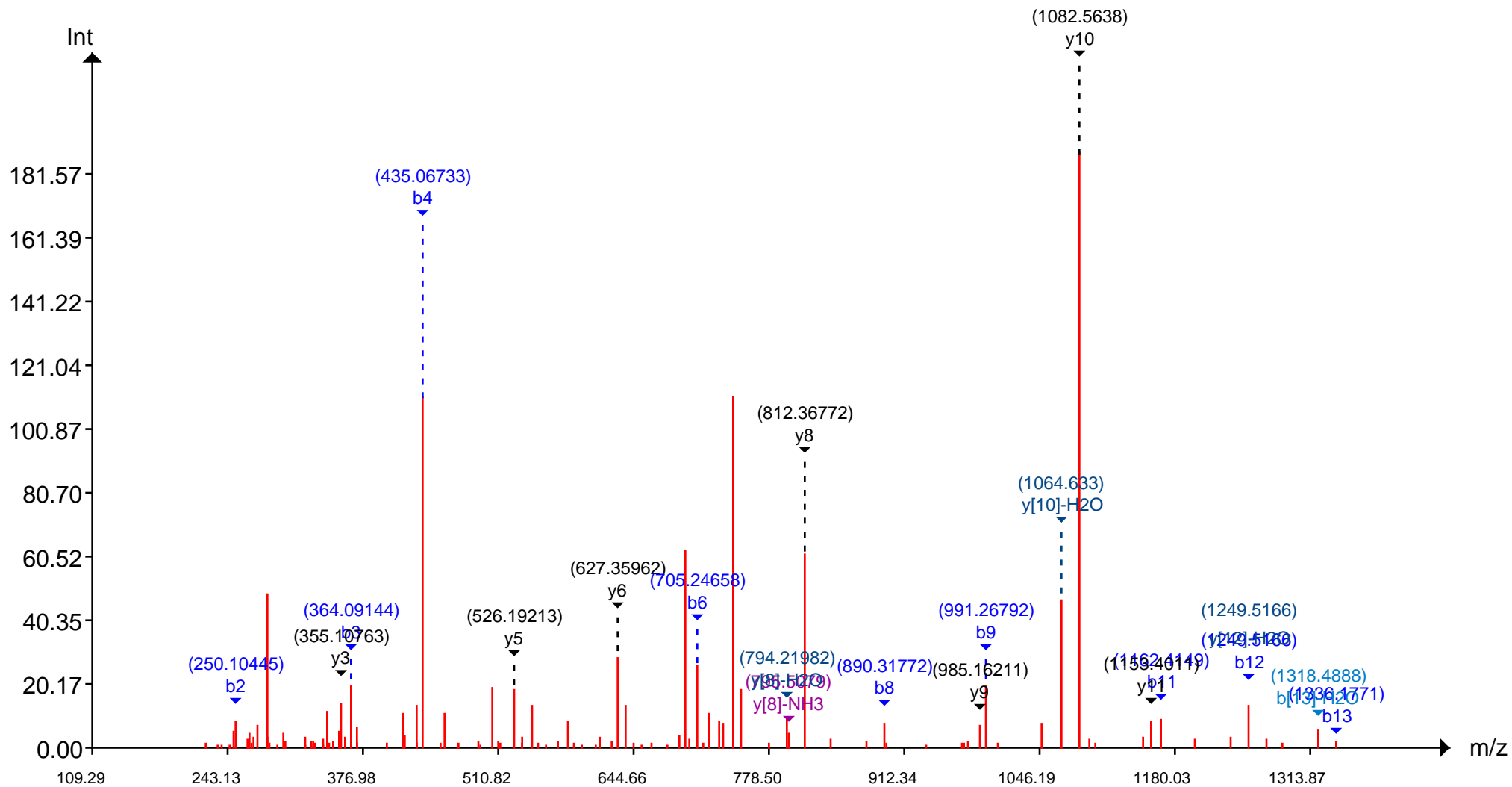
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 53.76 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31325 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | -3.57E-4 |



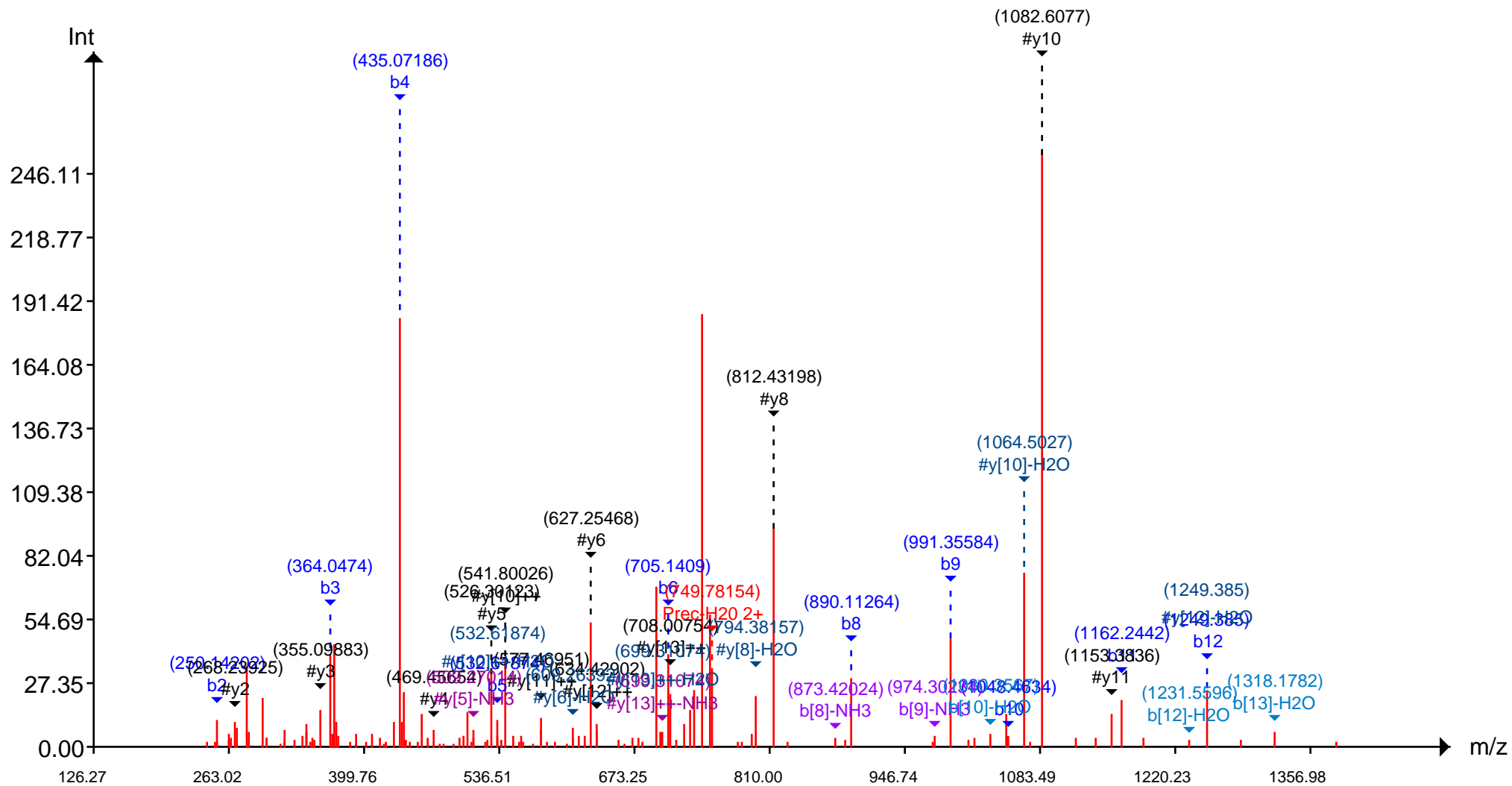
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 92.14 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31331 | Identity Threshold | 48.86 |
| Charge | 2+ | Mass error | -2.37E-4 |



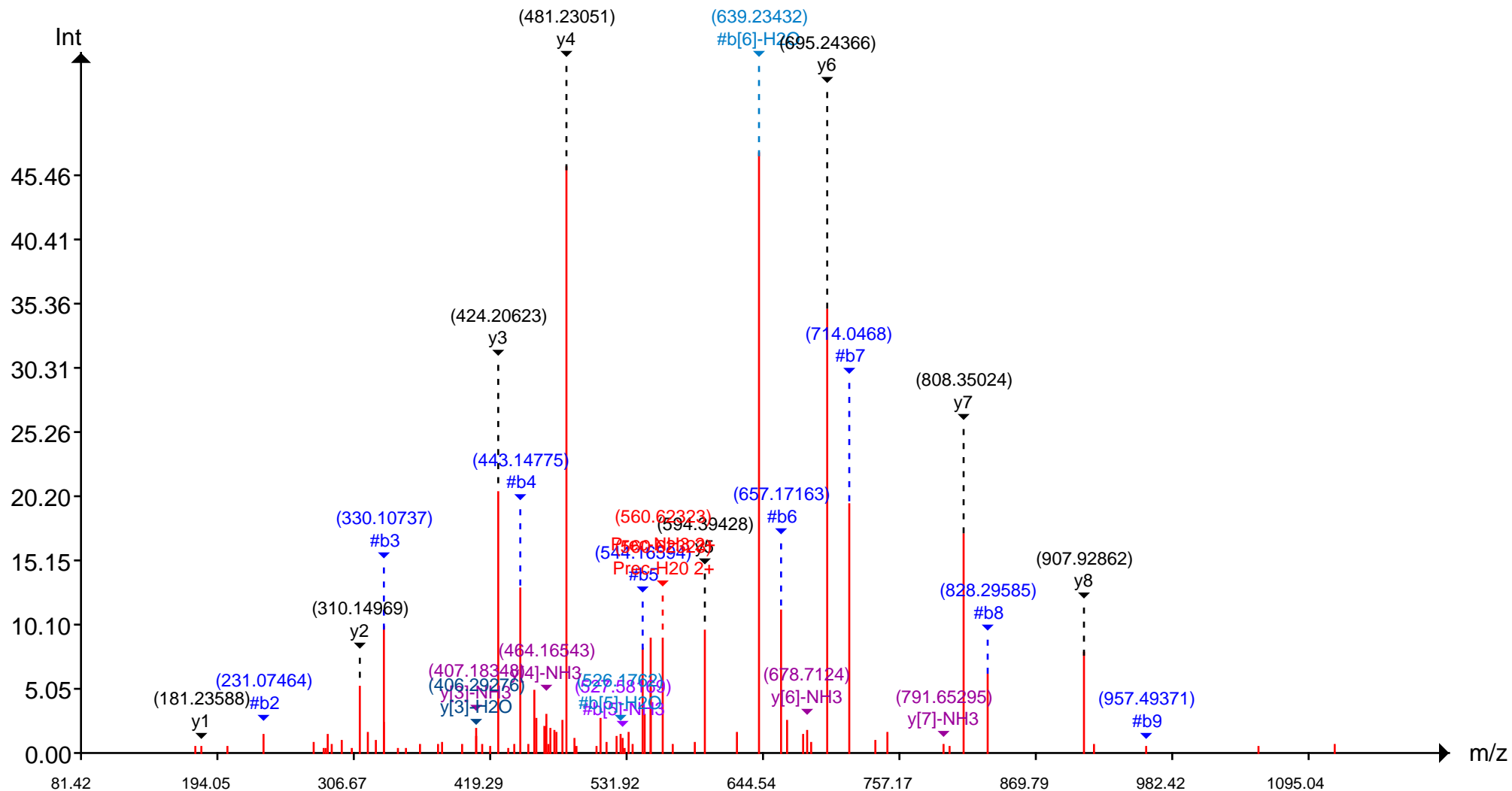
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR<C13>-COOH | Ion score | 73.41 |
| Protein | P23588 | Confidence level | 99.00% |
| Precursor m/z | 758.87319 | Identity Threshold | 48.10 |
| Charge | 2+ | Mass error | 4.28E-4 |



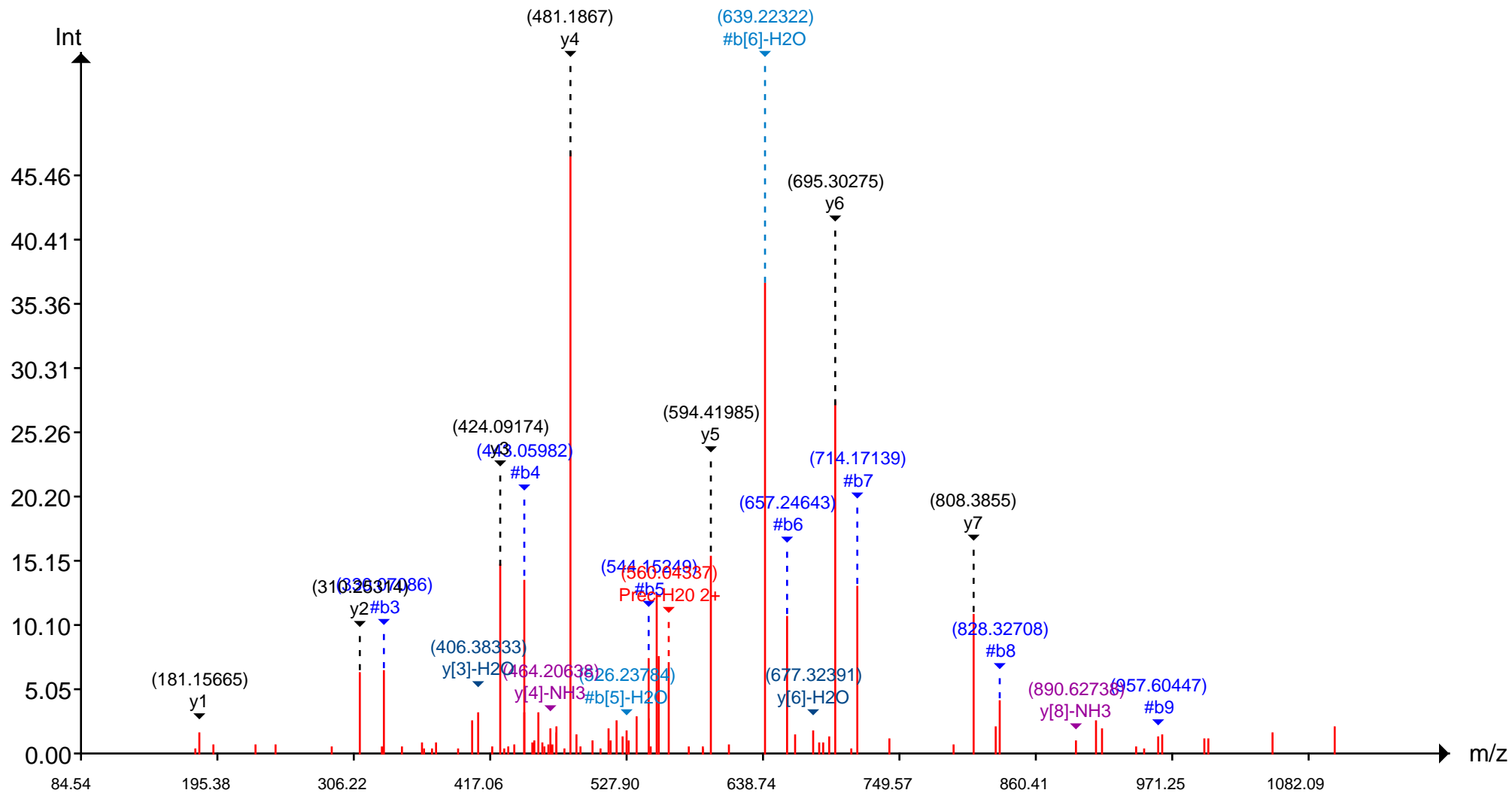
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-GM<Mox*>NAPK<AcD3K*>GQTGNSSR<C13>-COOH | Ion score | 71.59 |
| Protein | P23588 | Confidence level | 99.00% |
| Precursor m/z | 758.87312 | Identity Threshold | 48.07 |
| Charge | 2+ | Mass error | 2.88E-4 |



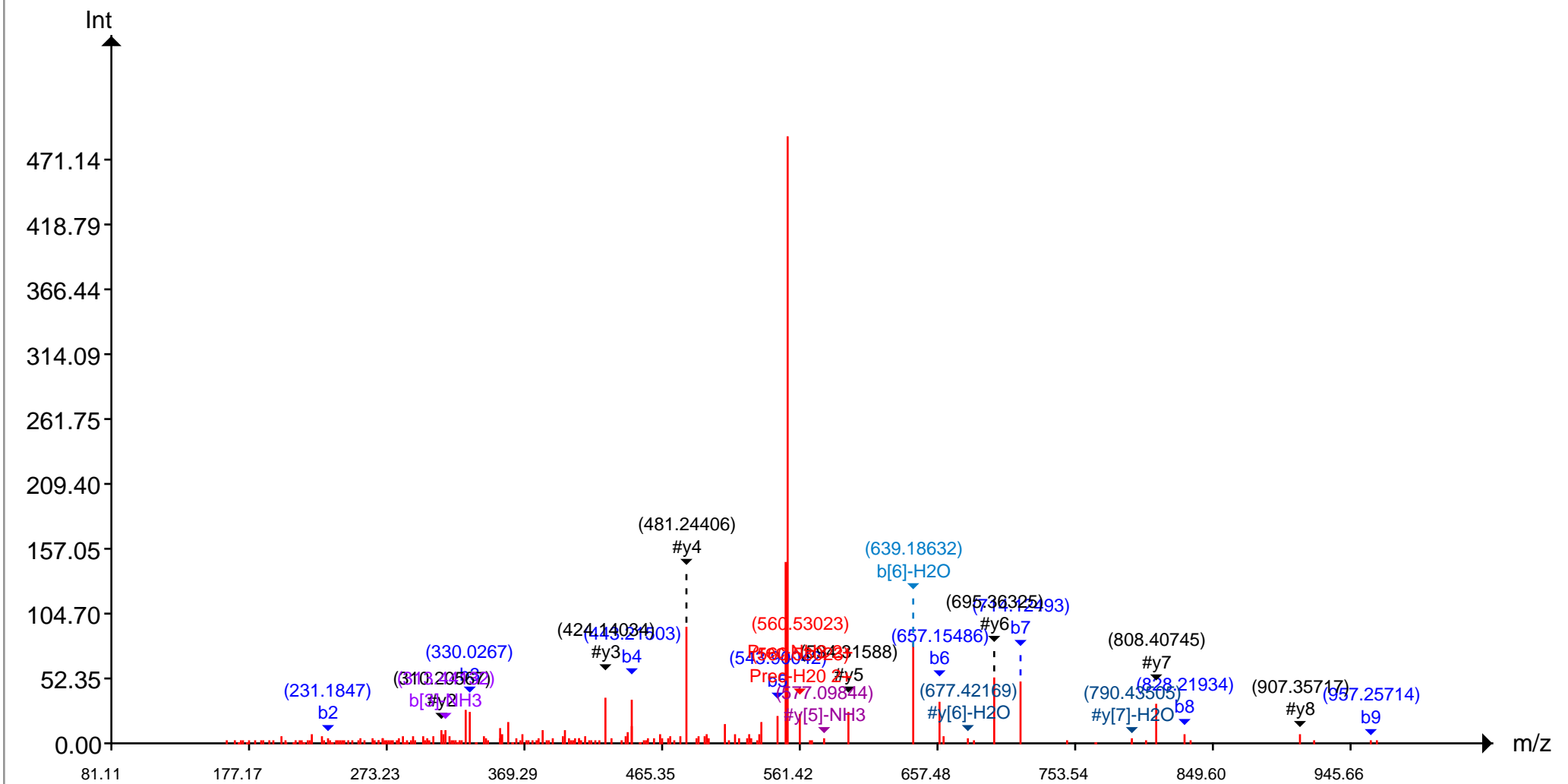
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 68.27 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32374 | Identity Threshold | 47.93 |
| Charge | 2+ | Mass error | 4.82E-4 |



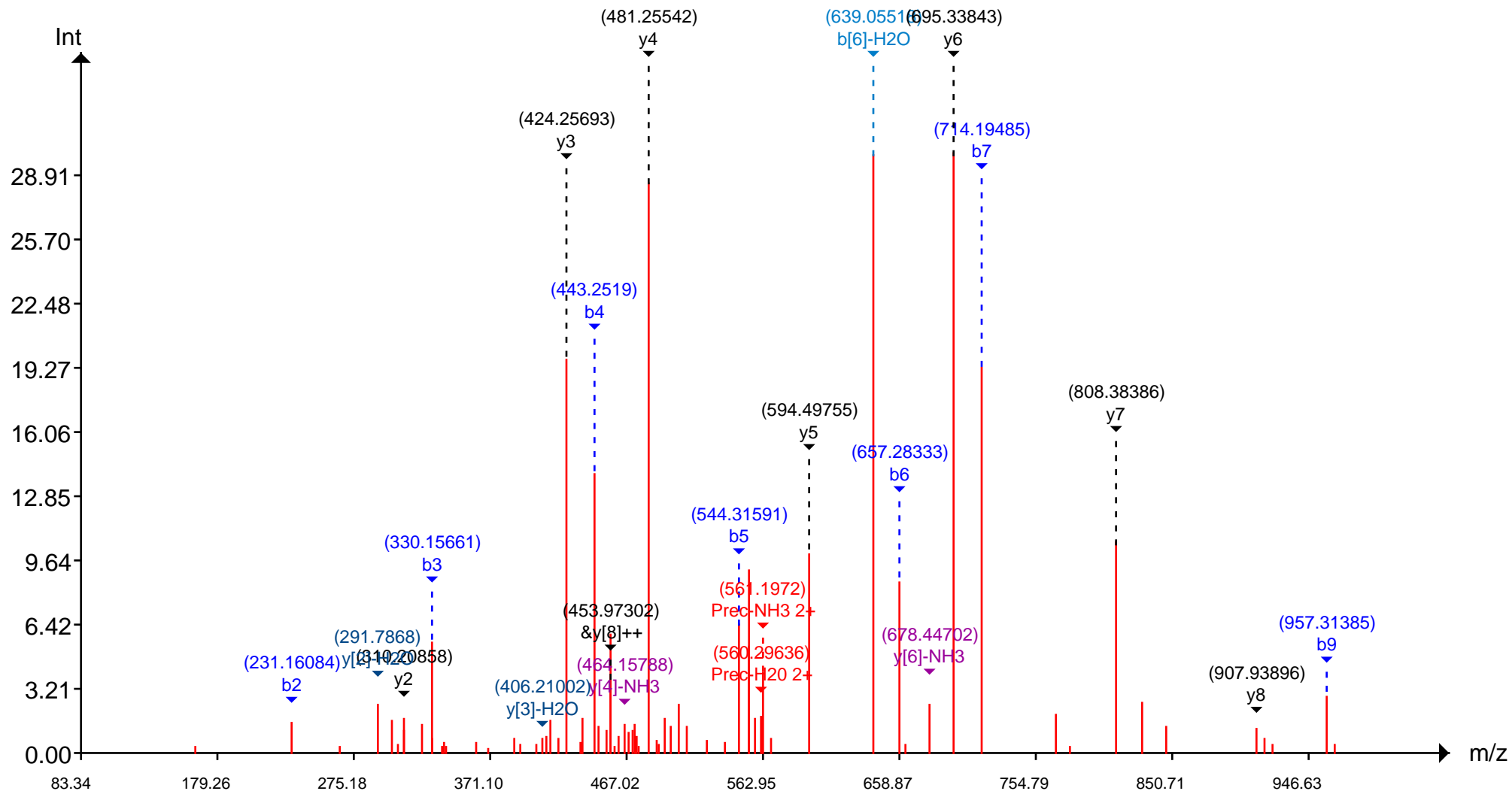
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 52.98 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32372 | Identity Threshold | 47.92 |
| Charge | 2+ | Mass error | 4.42E-4 |



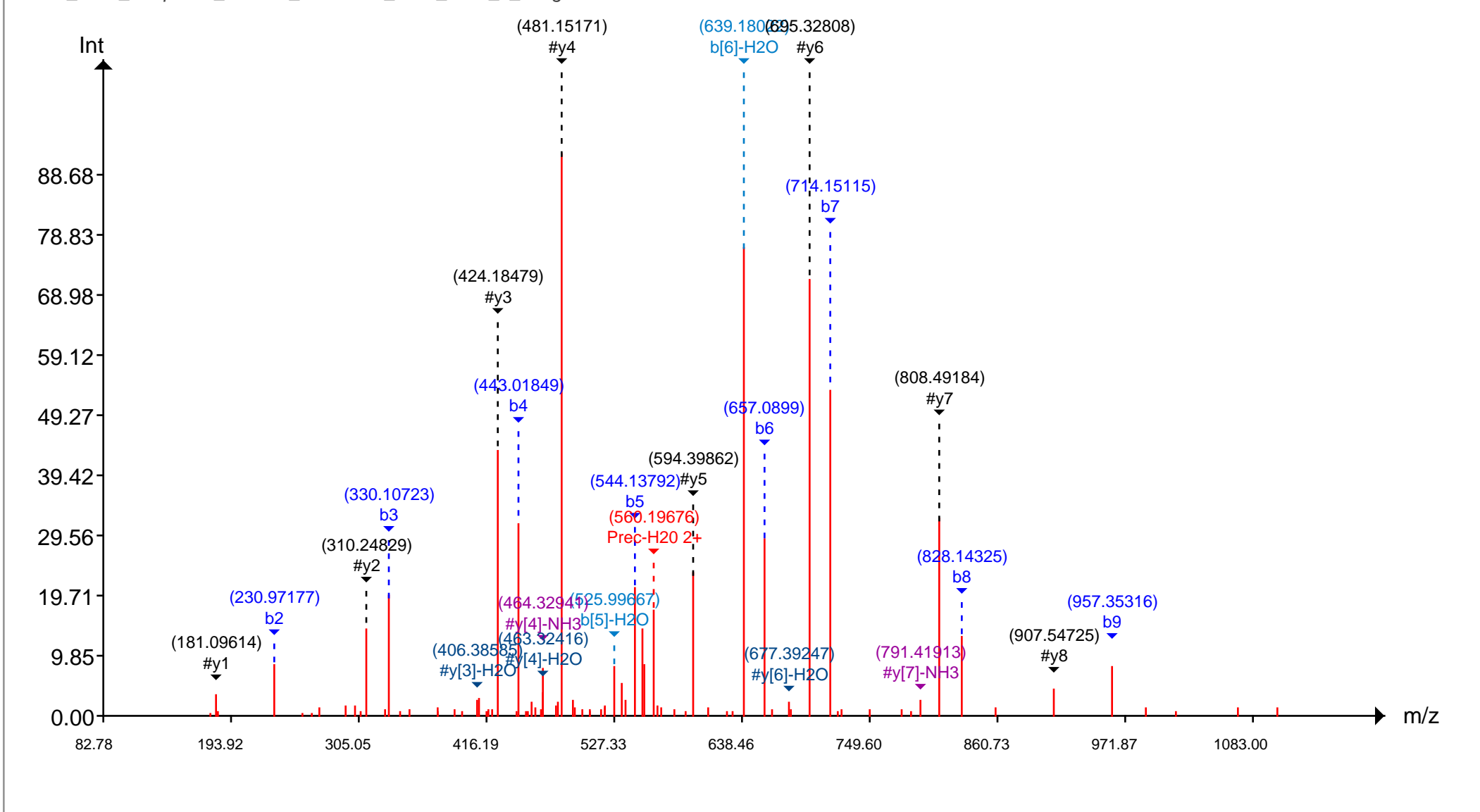
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 50.54 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32404 | Identity Threshold | 47.93 |
| Charge | 2+ | Mass error | 0.001082 |

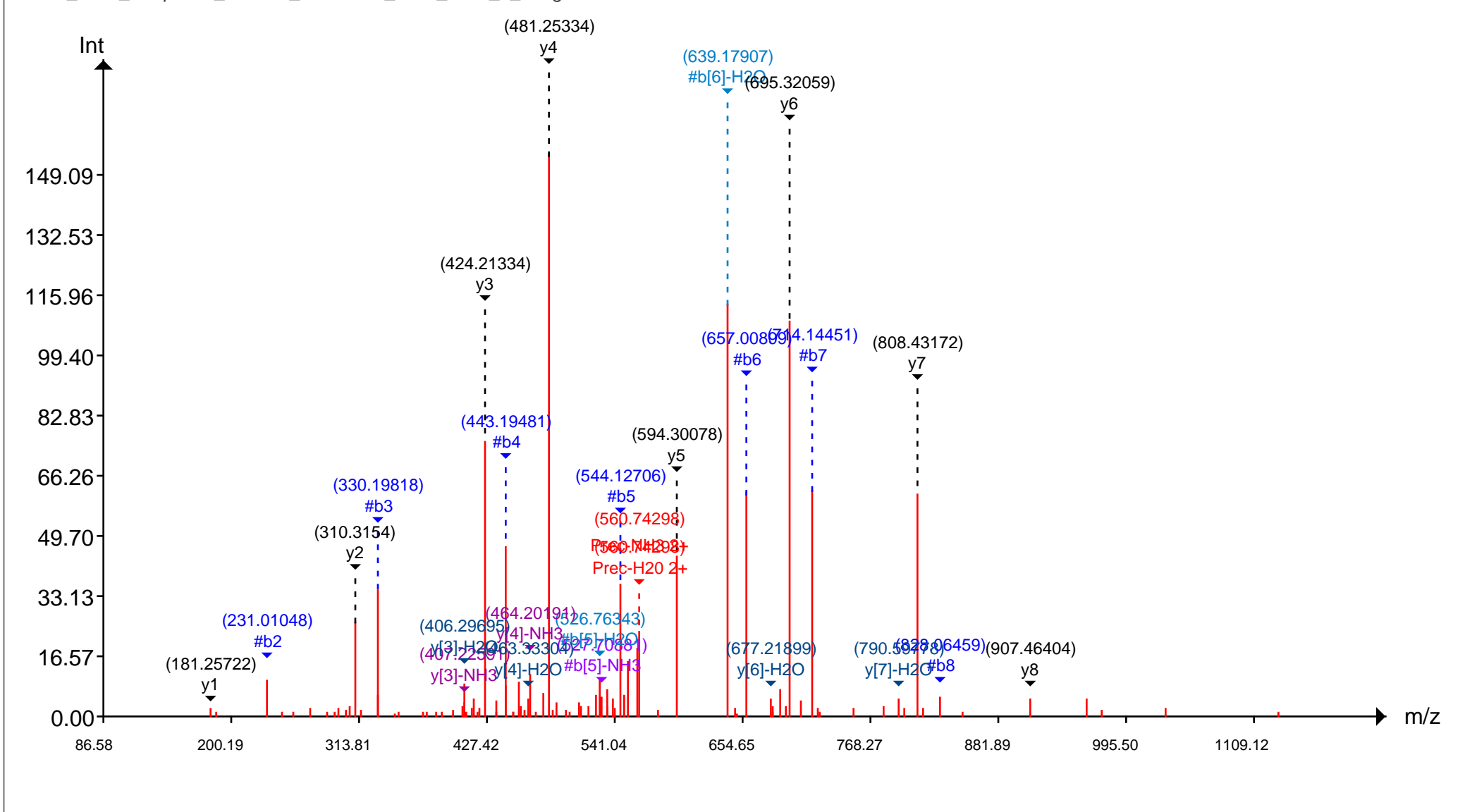


Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 65.44 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32375 | Identity Threshold | 47.93 |
| Charge | 2+ | Mass error | 5.02E-4 |

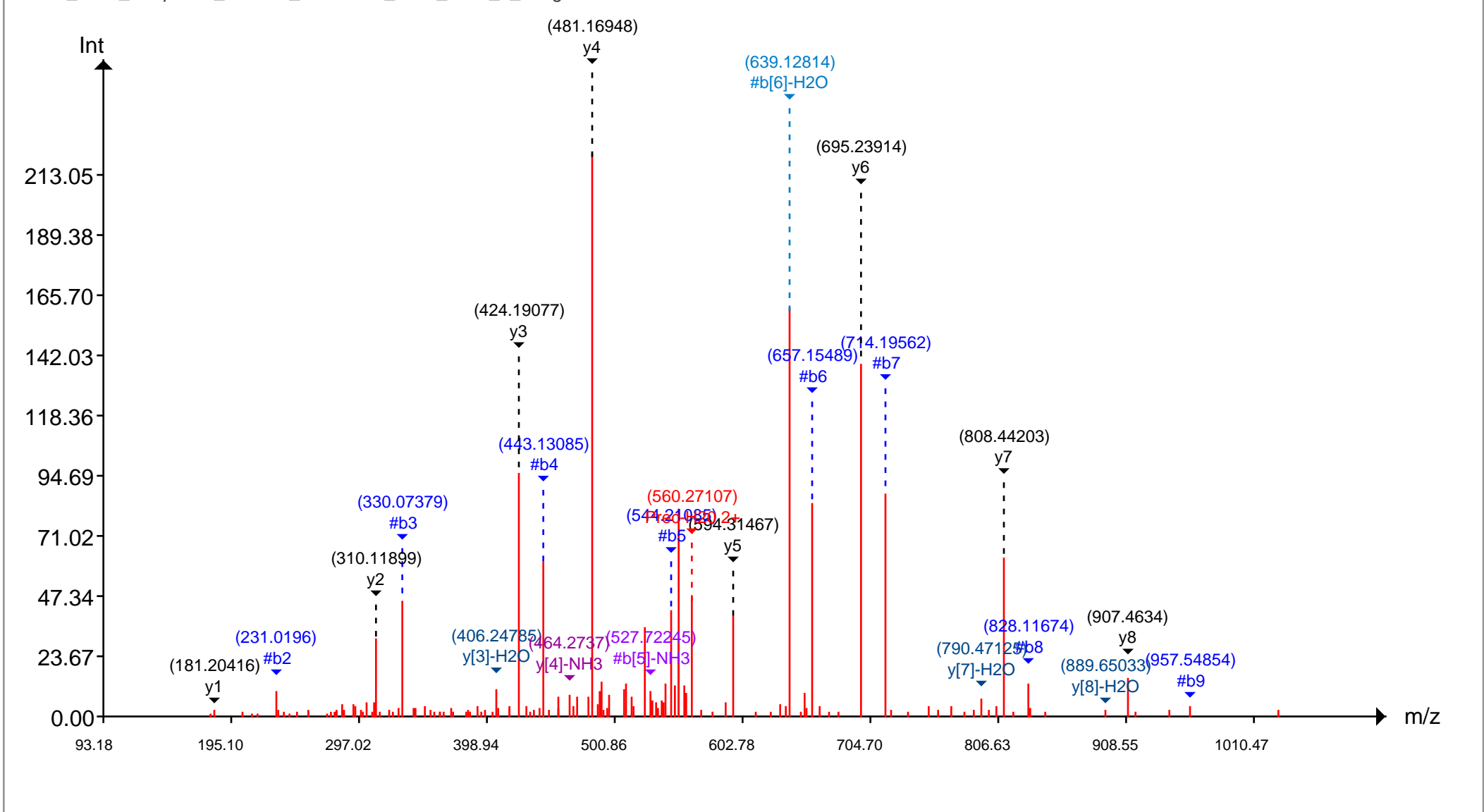


| Info | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 81.84 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32344 | Identity Threshold | 48.31 |
| Charge | 2+ | Mass error | -1.18E-4 |



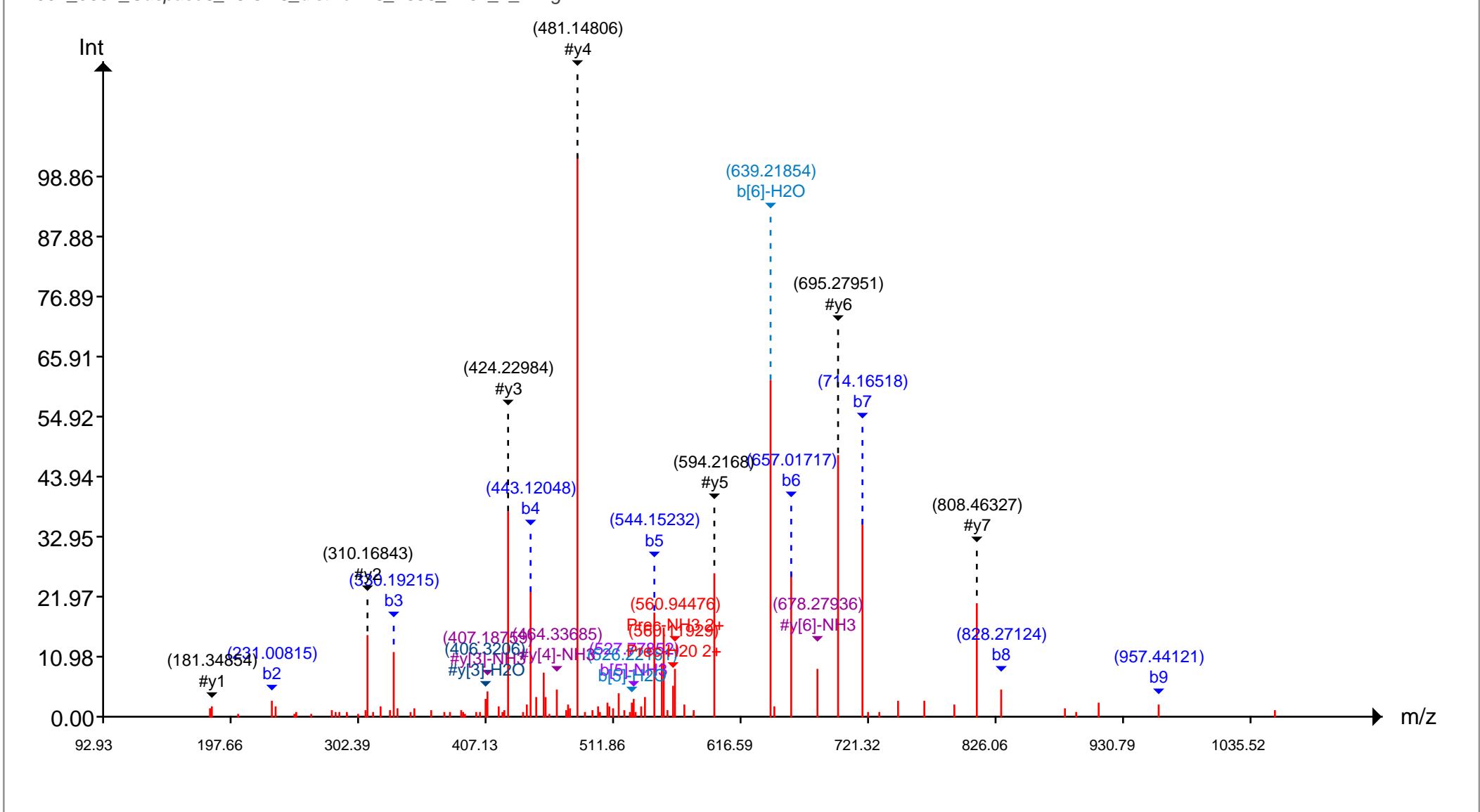
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 63.92 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32343 | Identity Threshold | 48.31 |
| Charge | 2+ | Mass error | -1.38E-4 |



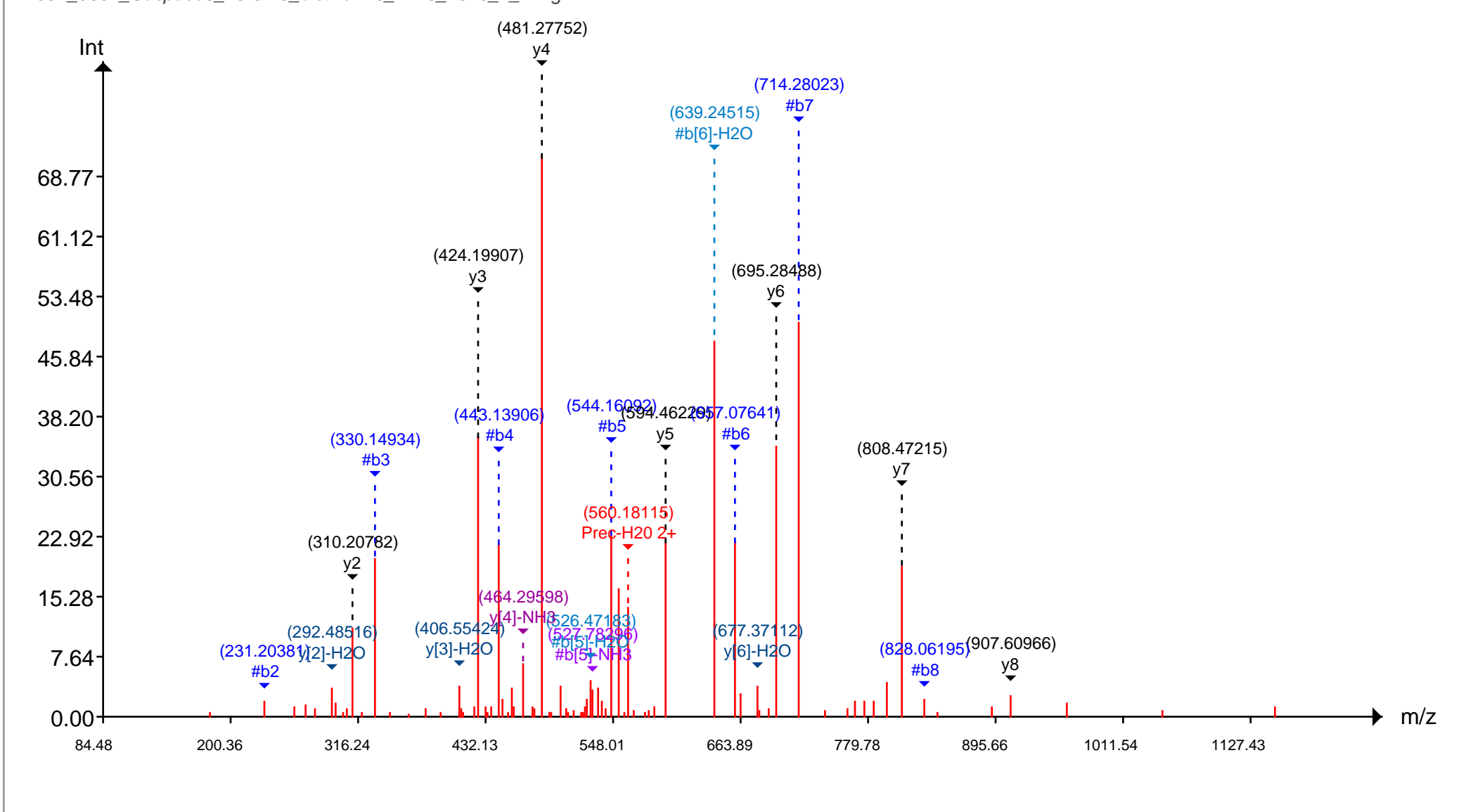
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 75.82 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32347 | Identity Threshold | 48.30 |
| Charge | 2+ | Mass error | -5.8E-5 |



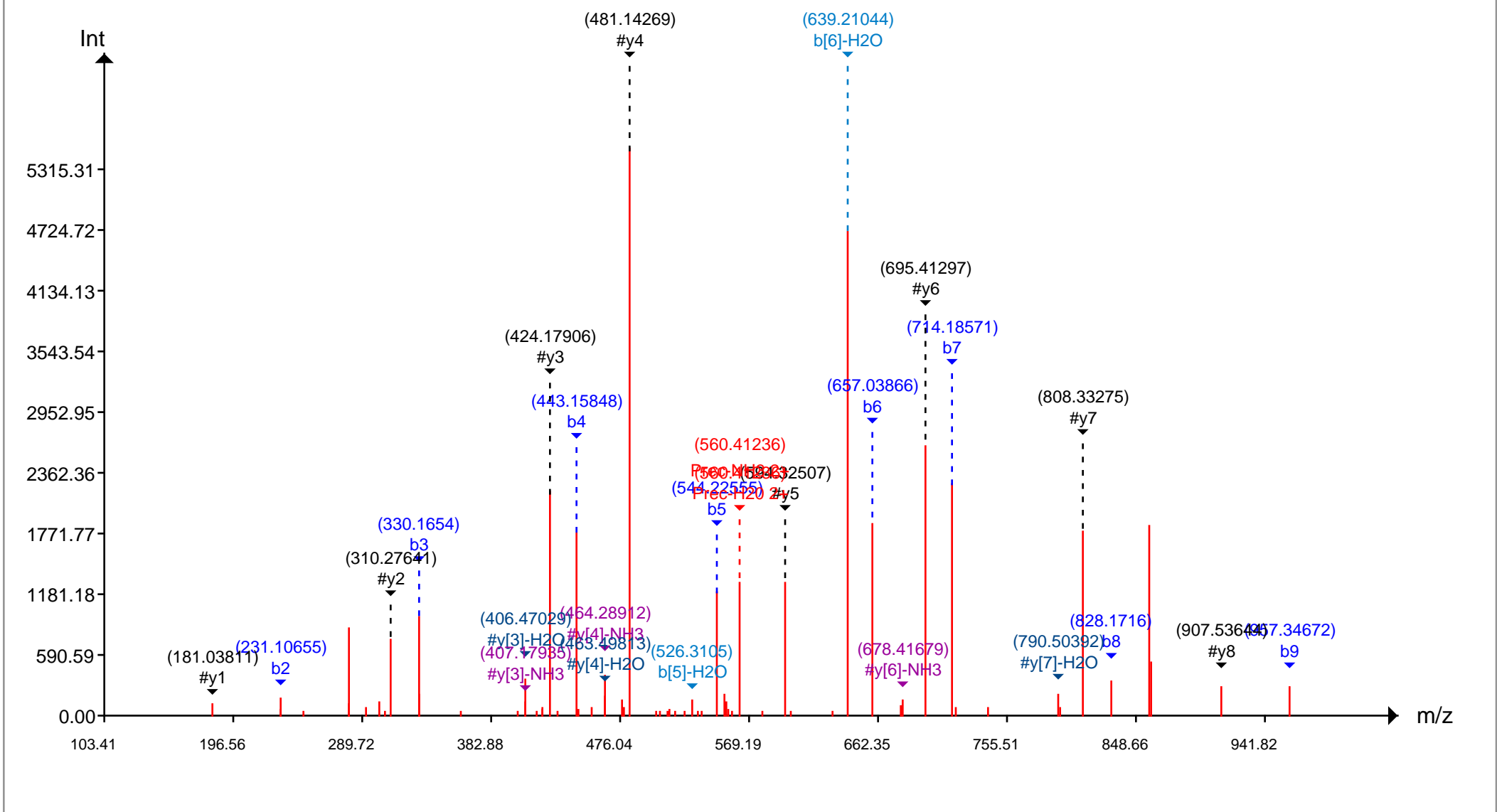
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 81.28 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32353 | Identity Threshold | 48.15 |
| Charge | 2+ | Mass error | 6.2E-5 |



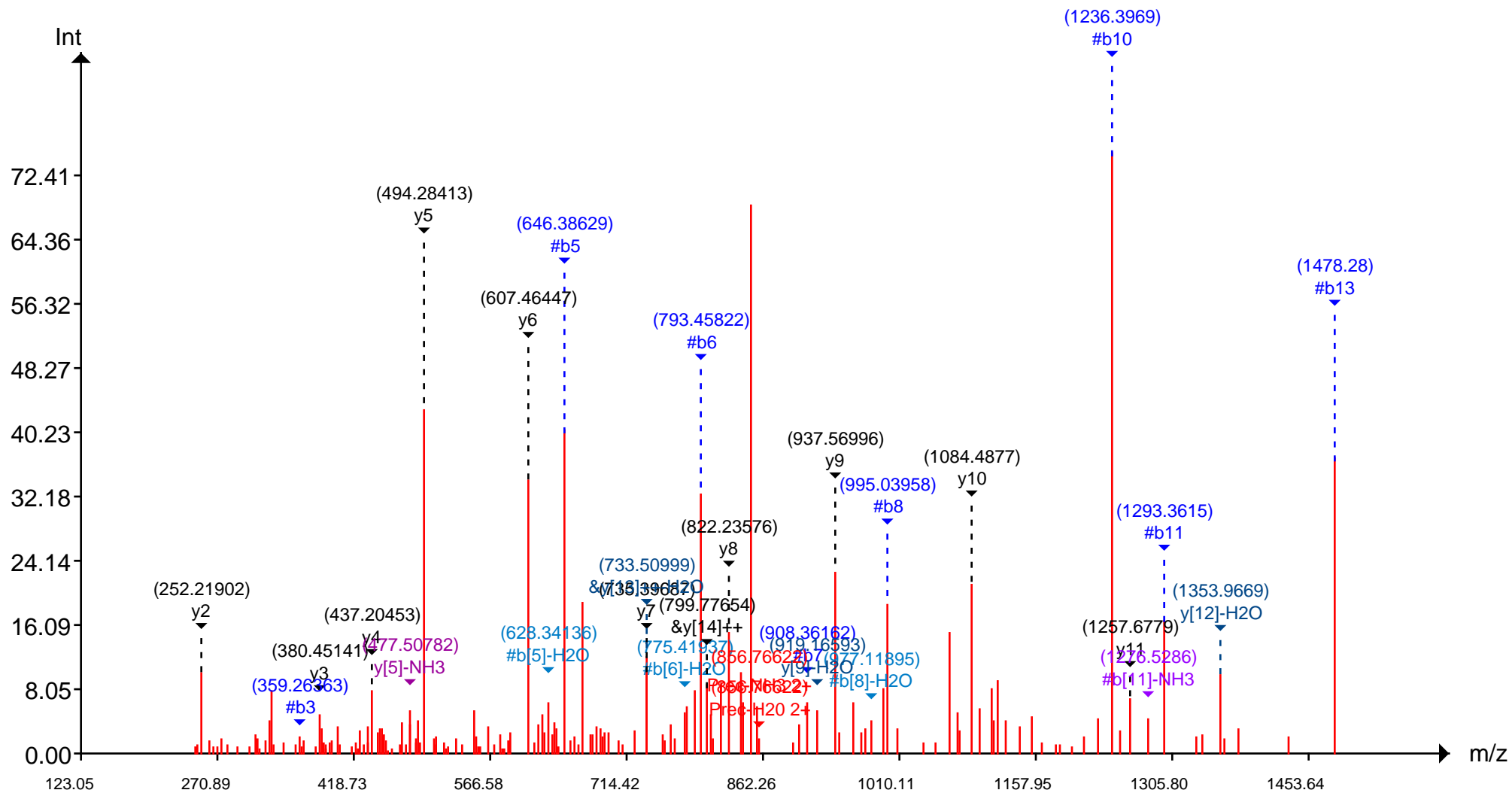
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 61.01 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32358 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 1.62E-4 |



Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 63.1 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32356 | Identity Threshold | 48.03 |
| Charge | 2+ | Mass error | 1.22E-4 |



Info

Modified Sequence AcD3-SLINK<AcD3K*>FDSQLGGQAR<C13>-COOH

Ion score 71.63

Protein Q9P2M7

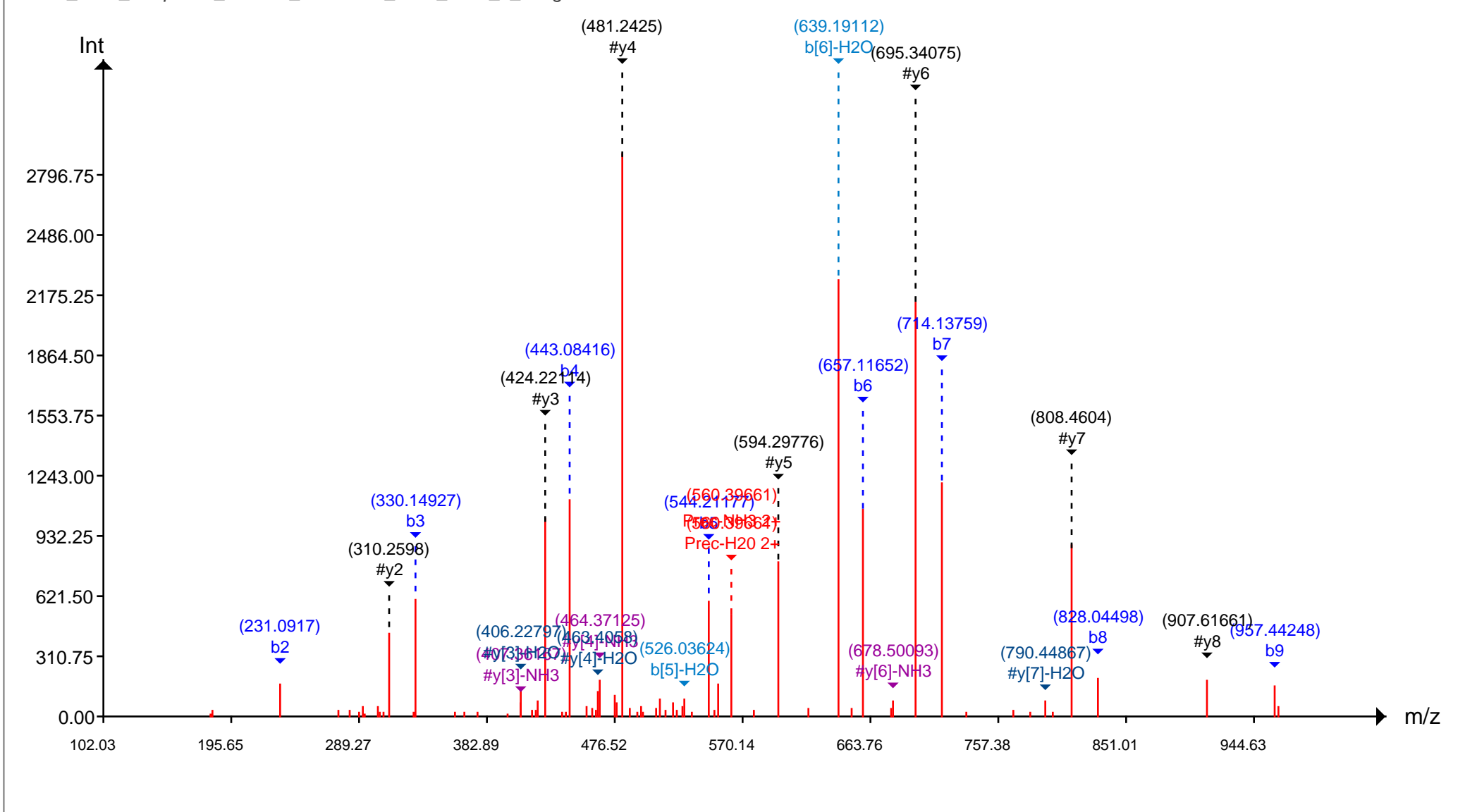
Confidence level 99.00%

Precursor m/z 865.4757

Identity Threshold 48.42

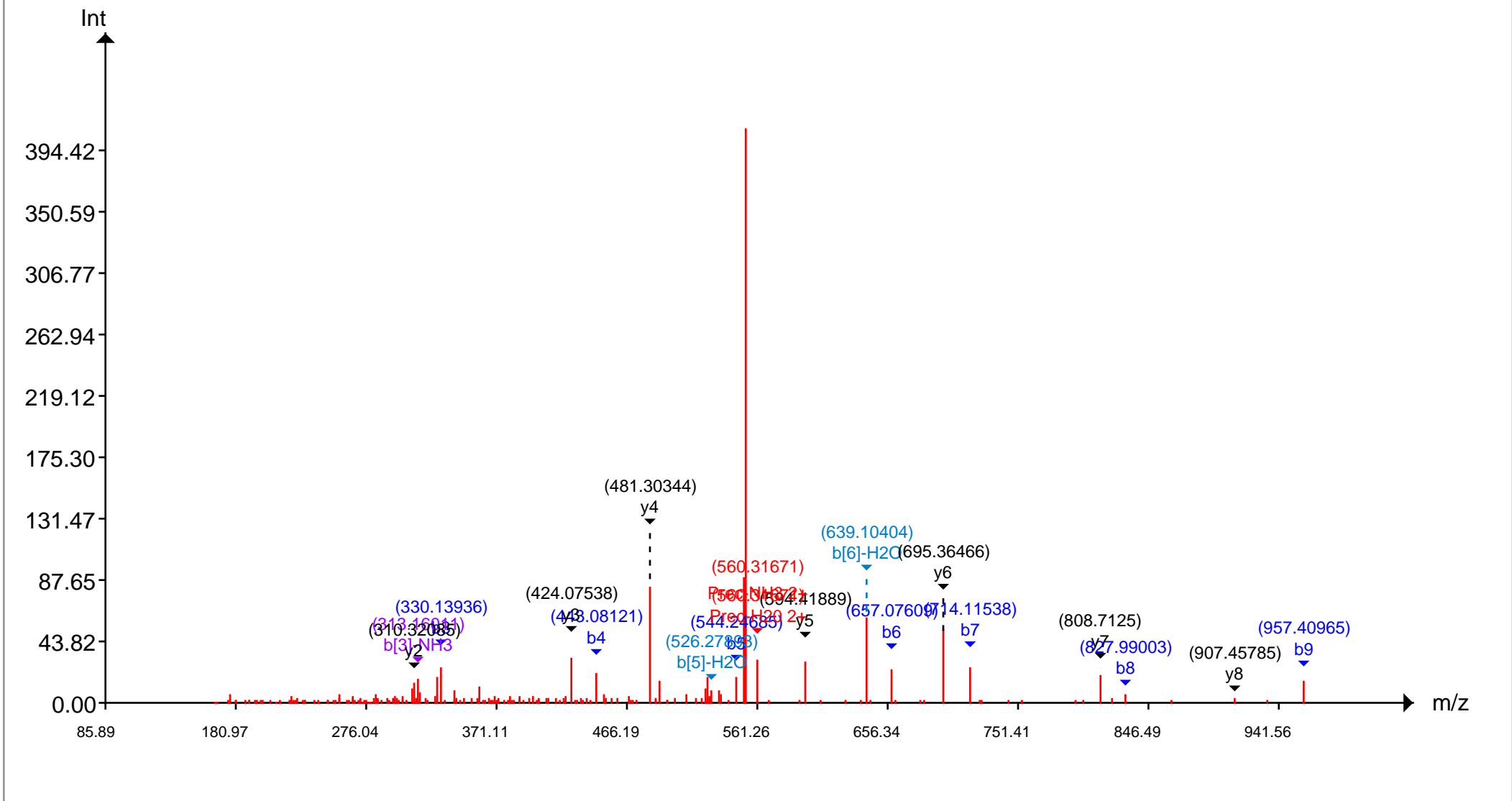
Charge 2+

Mass error -4.38E-4



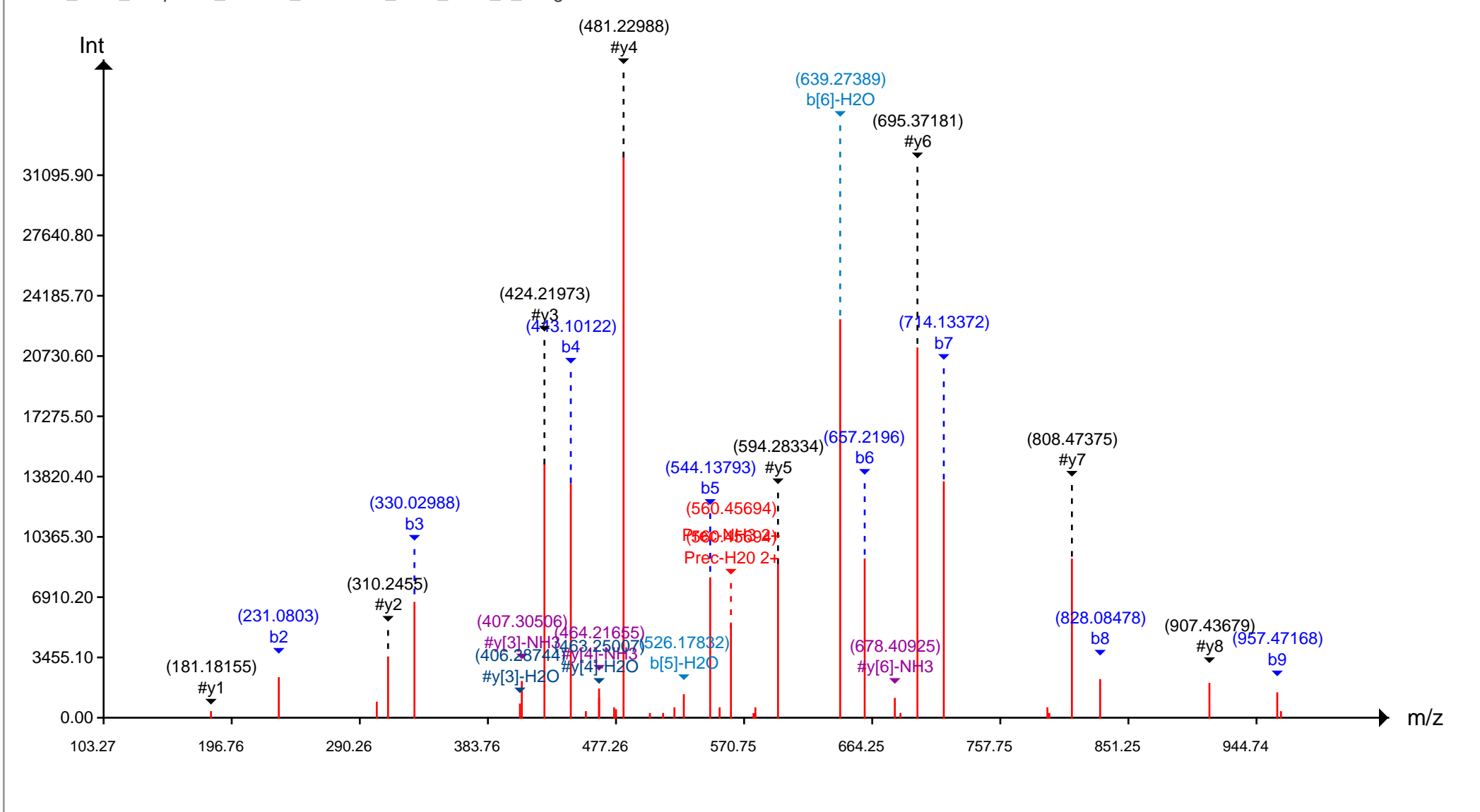
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 86.14 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32322 | Identity Threshold | 48.30 |
| Charge | 2+ | Mass error | -5.58E-4 |



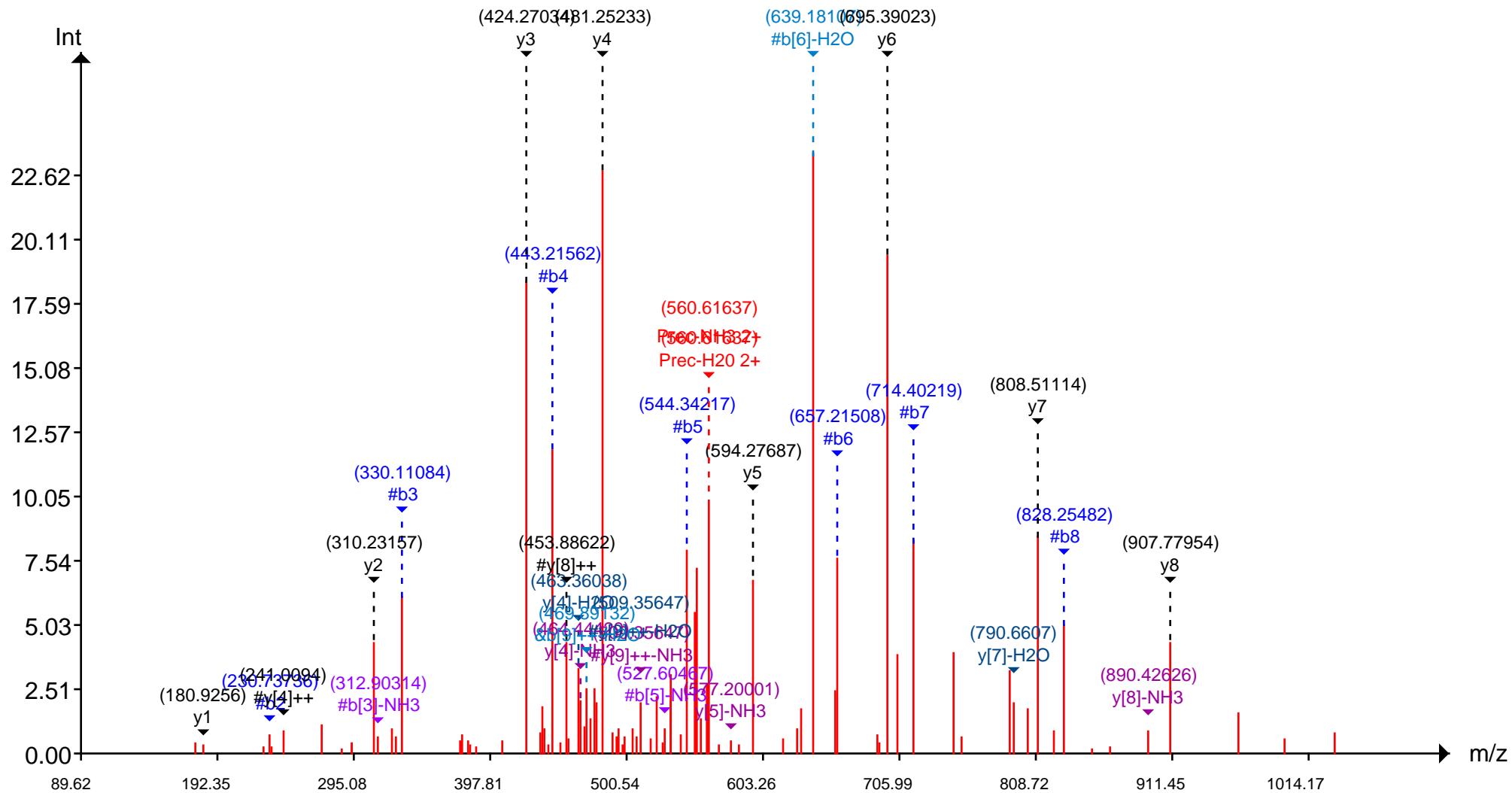
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 48.92 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32335 | Identity Threshold | 48.29 |
| Charge | 2+ | Mass error | -2.98E-4 |



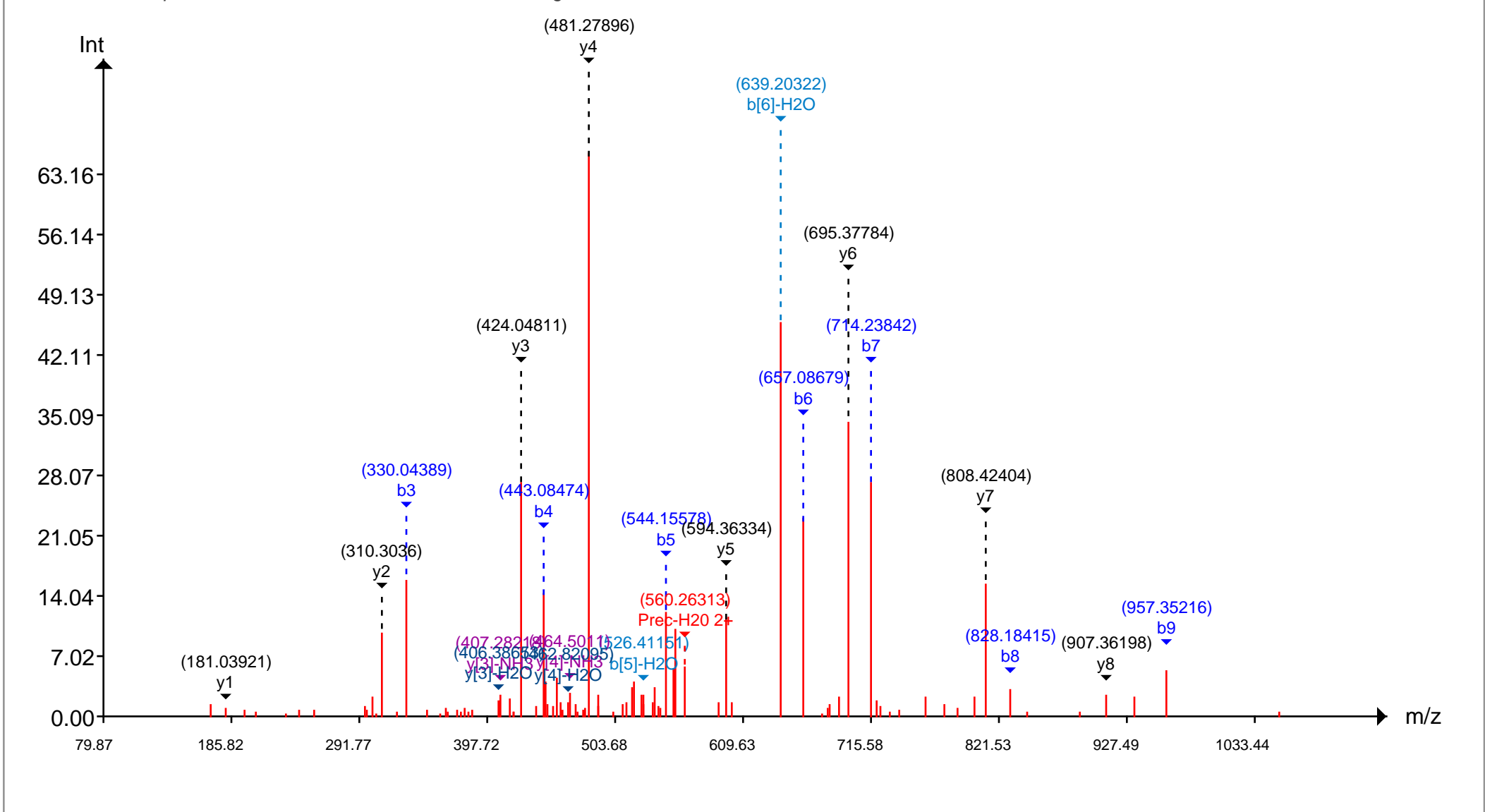
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 92.1 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32323 | Identity Threshold | 48.29 |
| Charge | 2+ | Mass error | -5.38E-4 |



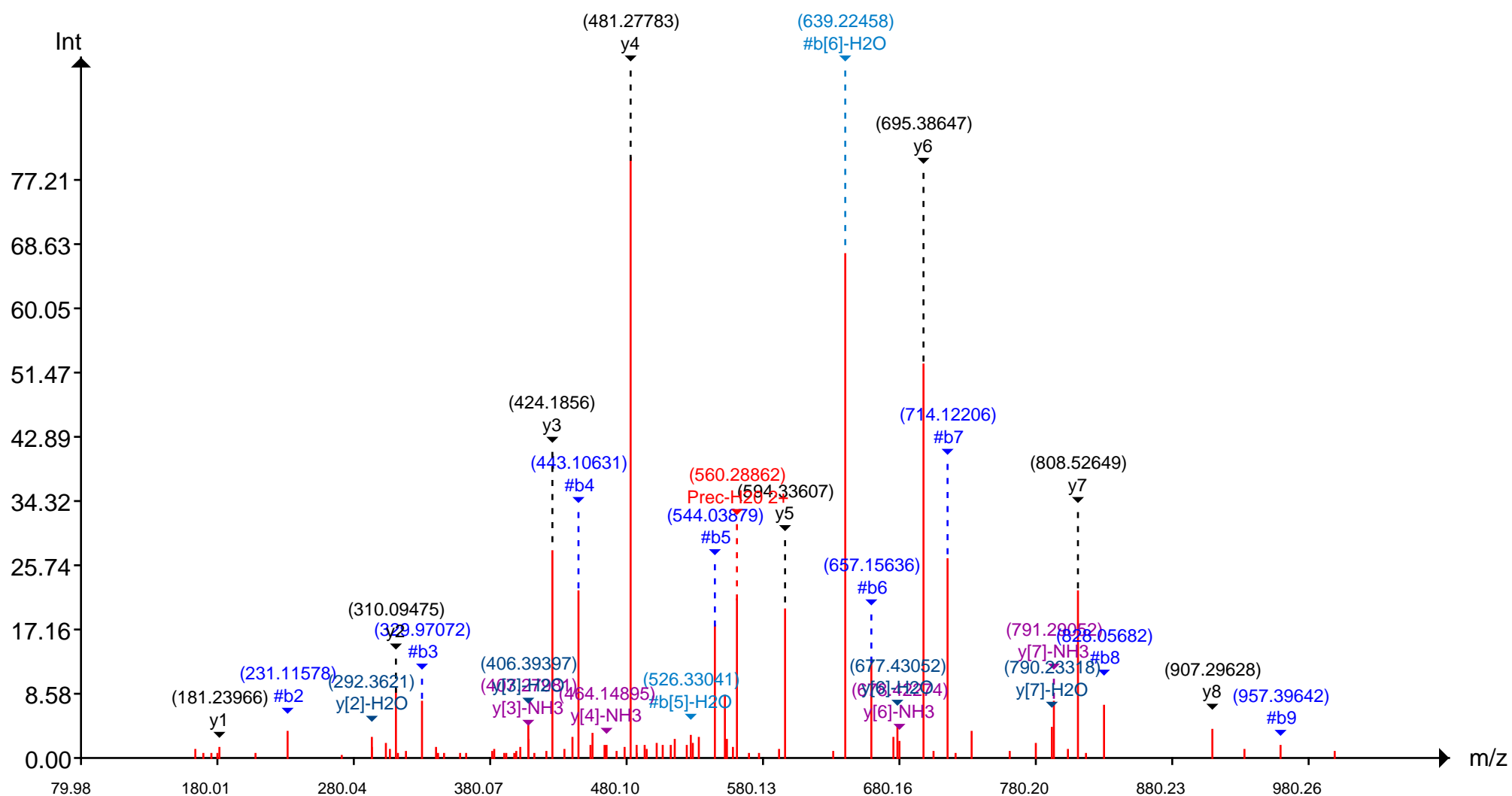
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 55.47 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32365 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 3.02E-4 |



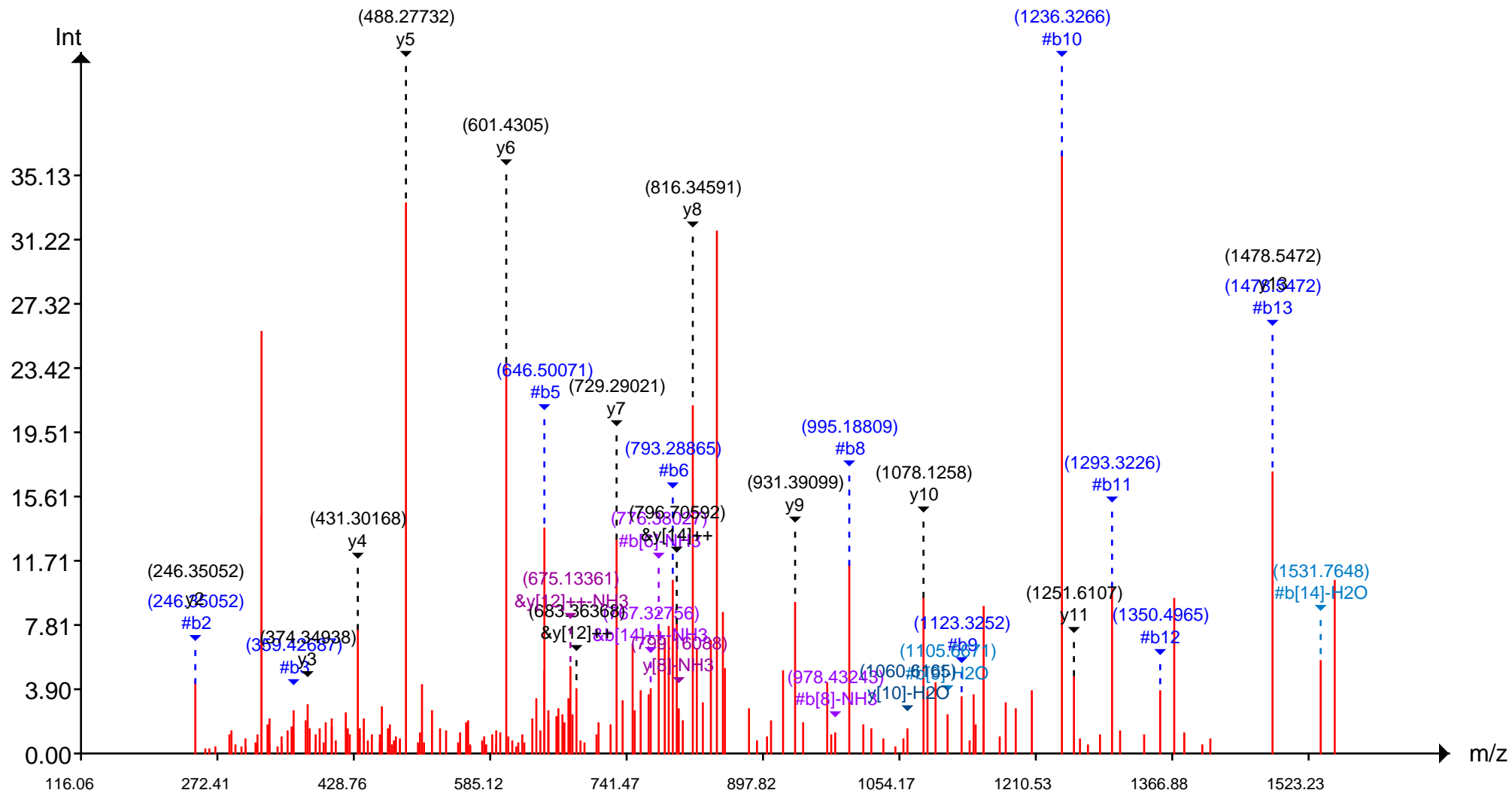
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 75.0 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32365 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 3.02E-4 |



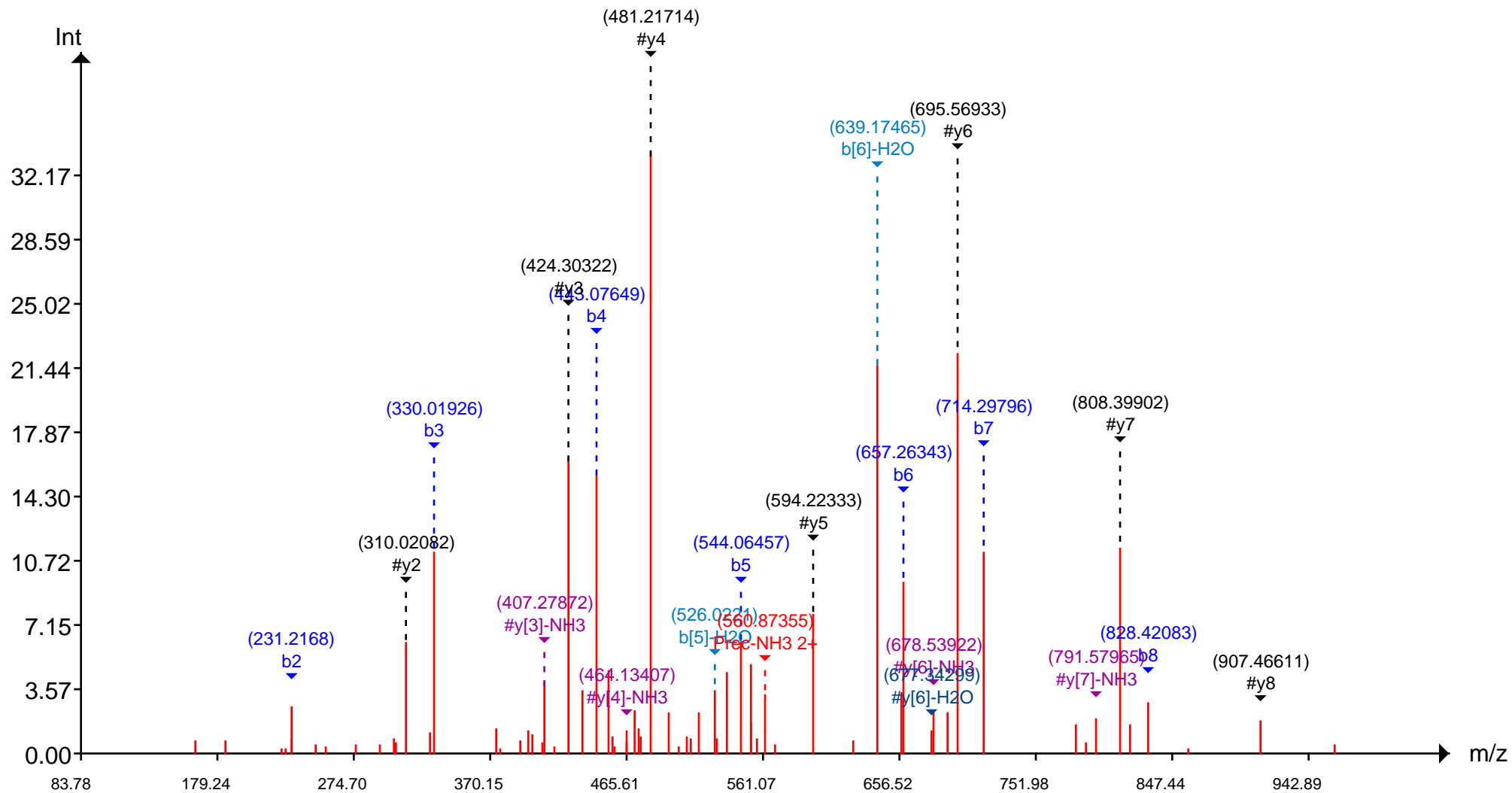
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 81.41 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32367 | Identity Threshold | 47.93 |
| Charge | 2+ | Mass error | 3.42E-4 |



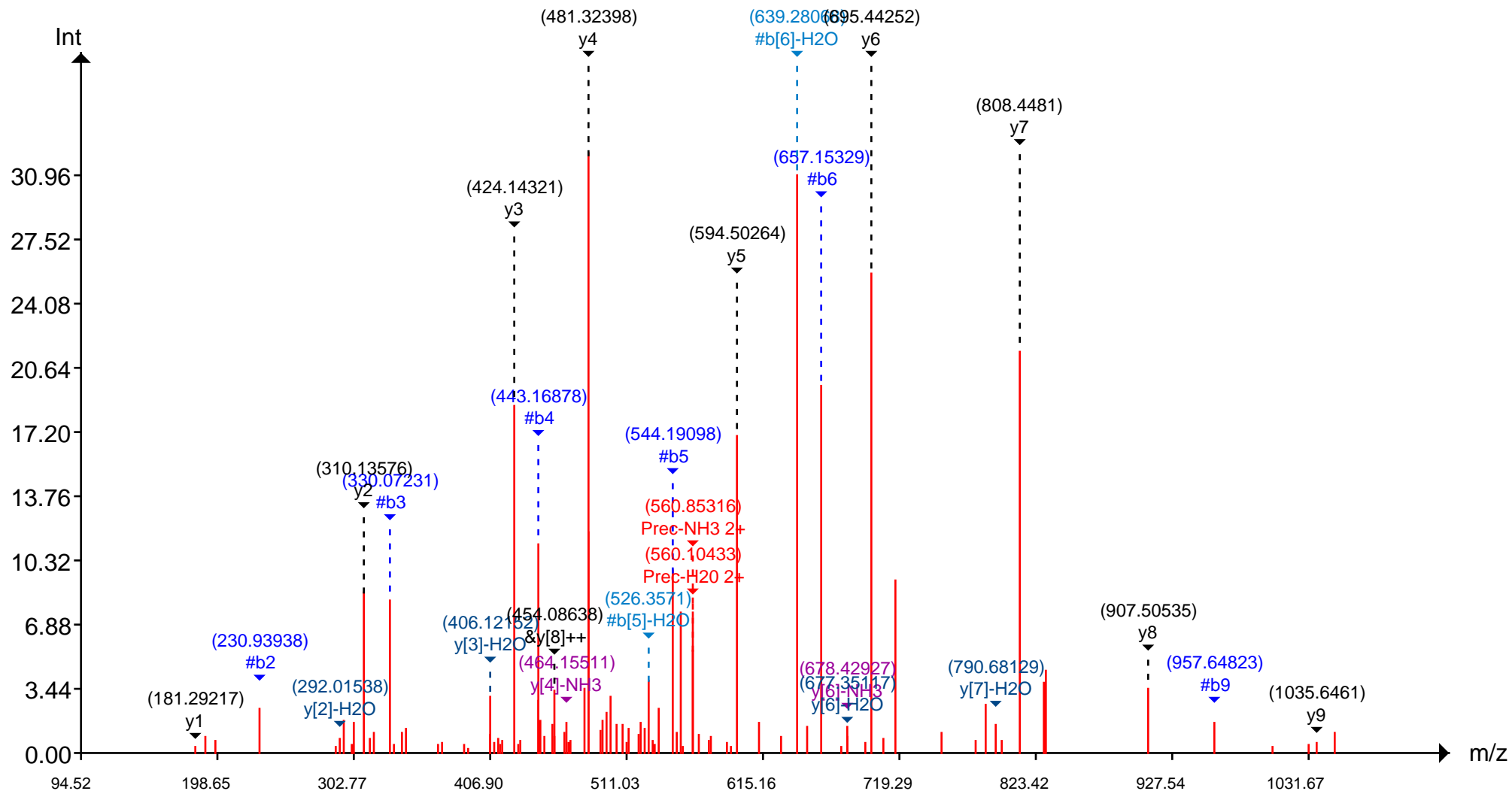
Info

| | | | |
|--------------------------|-----------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-SLINK<AcD3K*>FDSQLGGQAR-COOH | Ion score | 69.23 |
| Protein | Q9P2M7 | Confidence level | 99.00% |
| Precursor m/z | 862.46654 | Identity Threshold | 48.79 |
| Charge | 2+ | Mass error | 0.001383 |



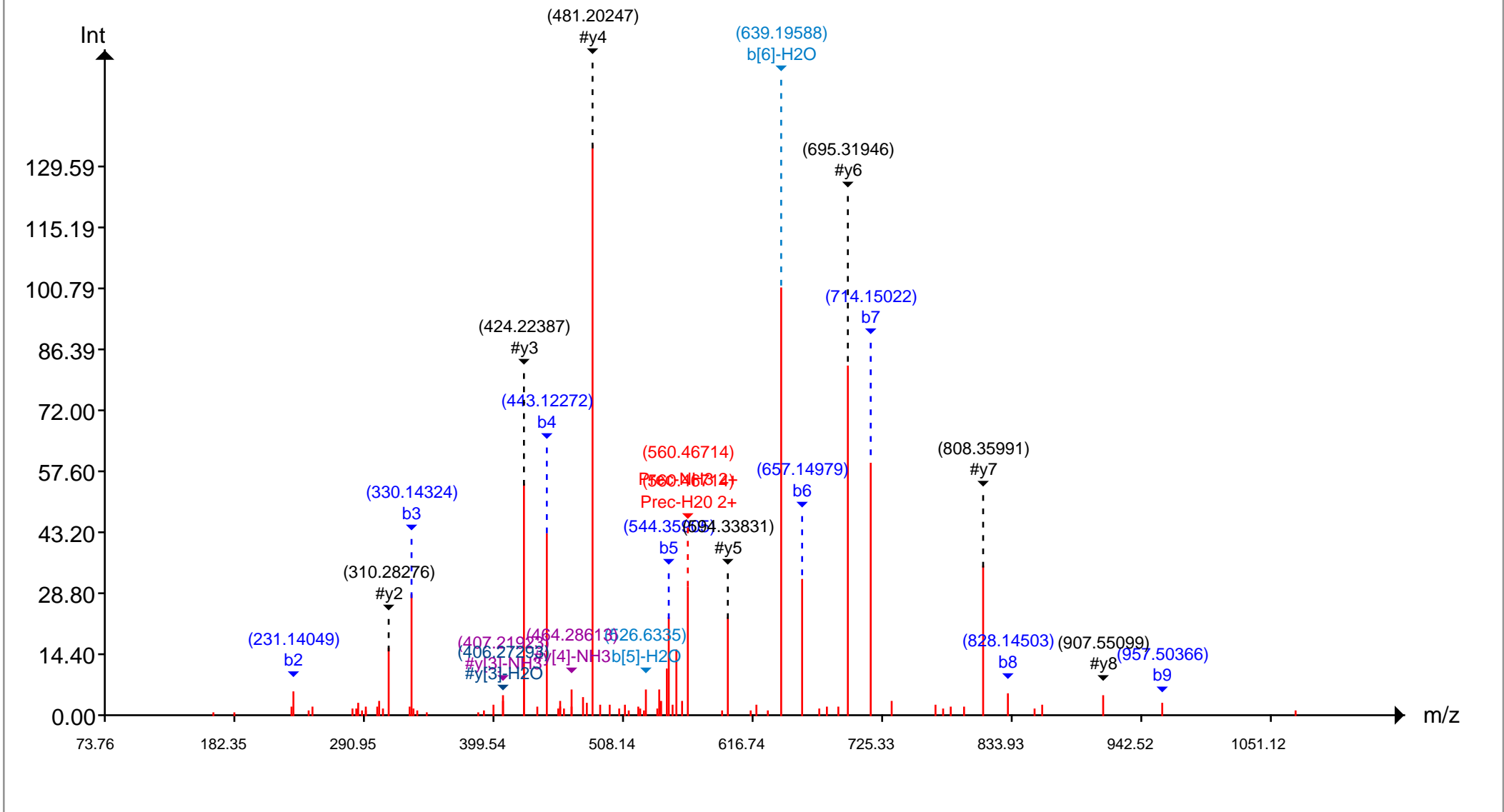
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 67.4 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32368 | Identity Threshold | 47.93 |
| Charge | 2+ | Mass error | 3.62E-4 |



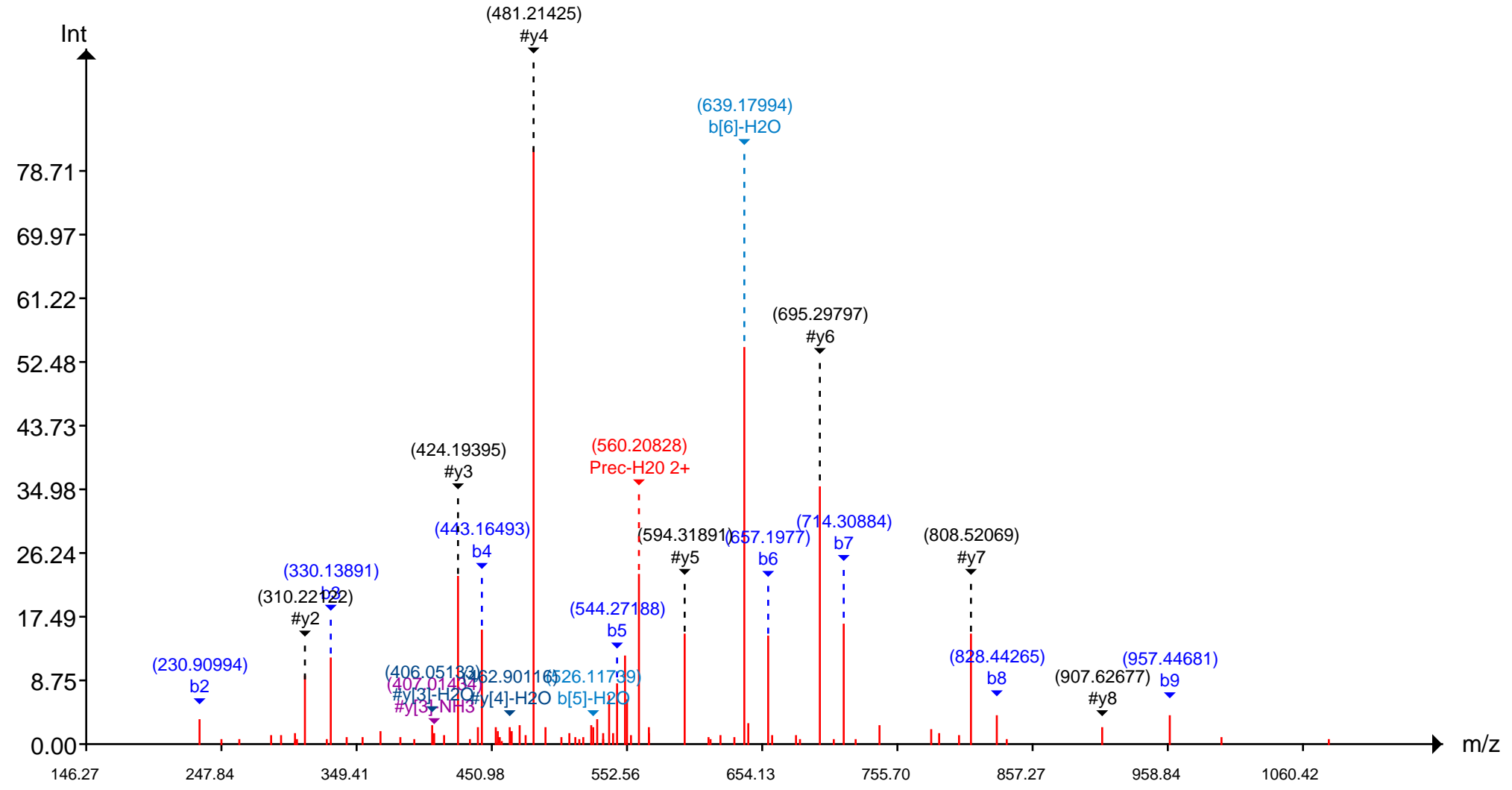
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 62.51 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32359 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 1.82E-4 |



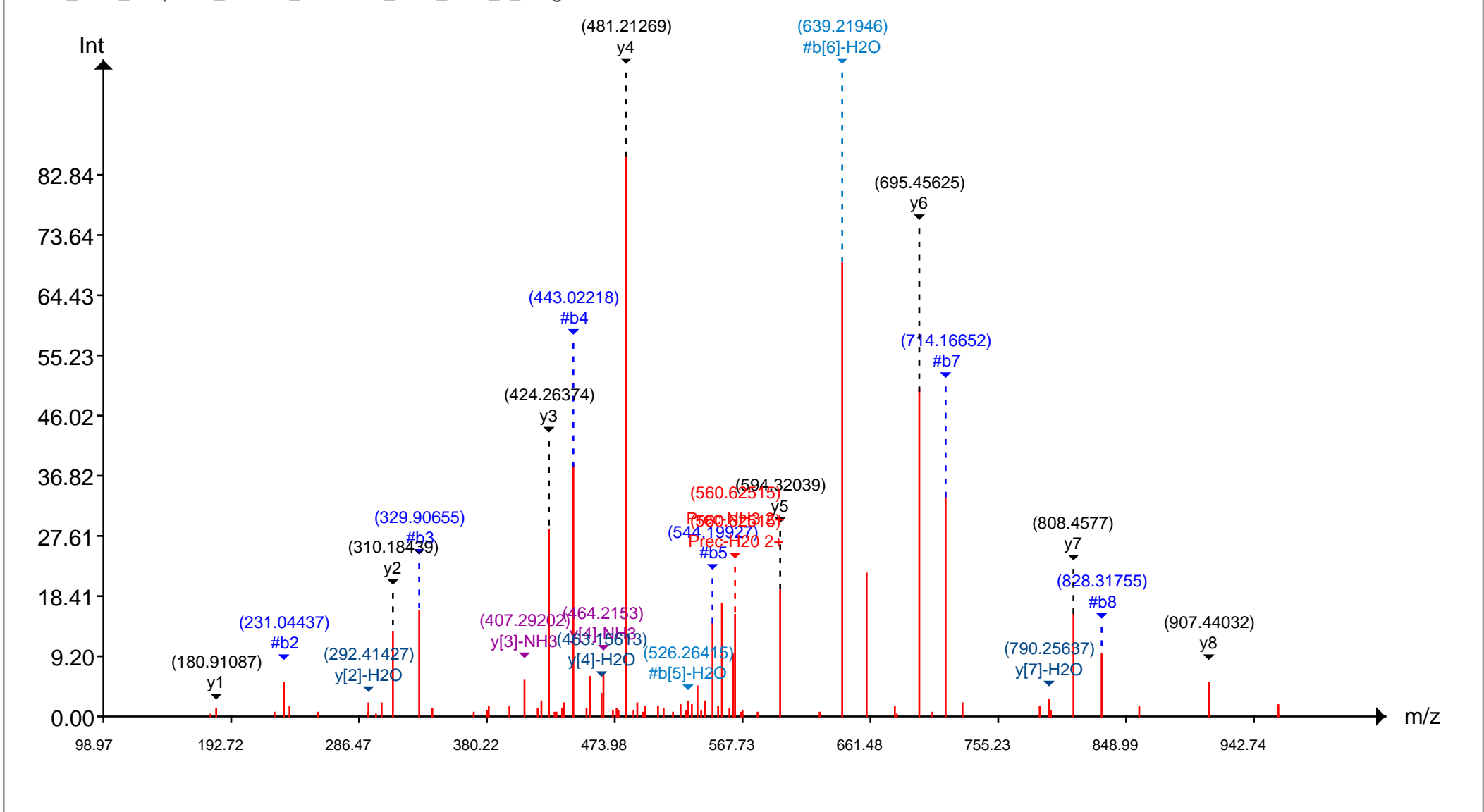
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 65.53 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32359 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 1.82E-4 |



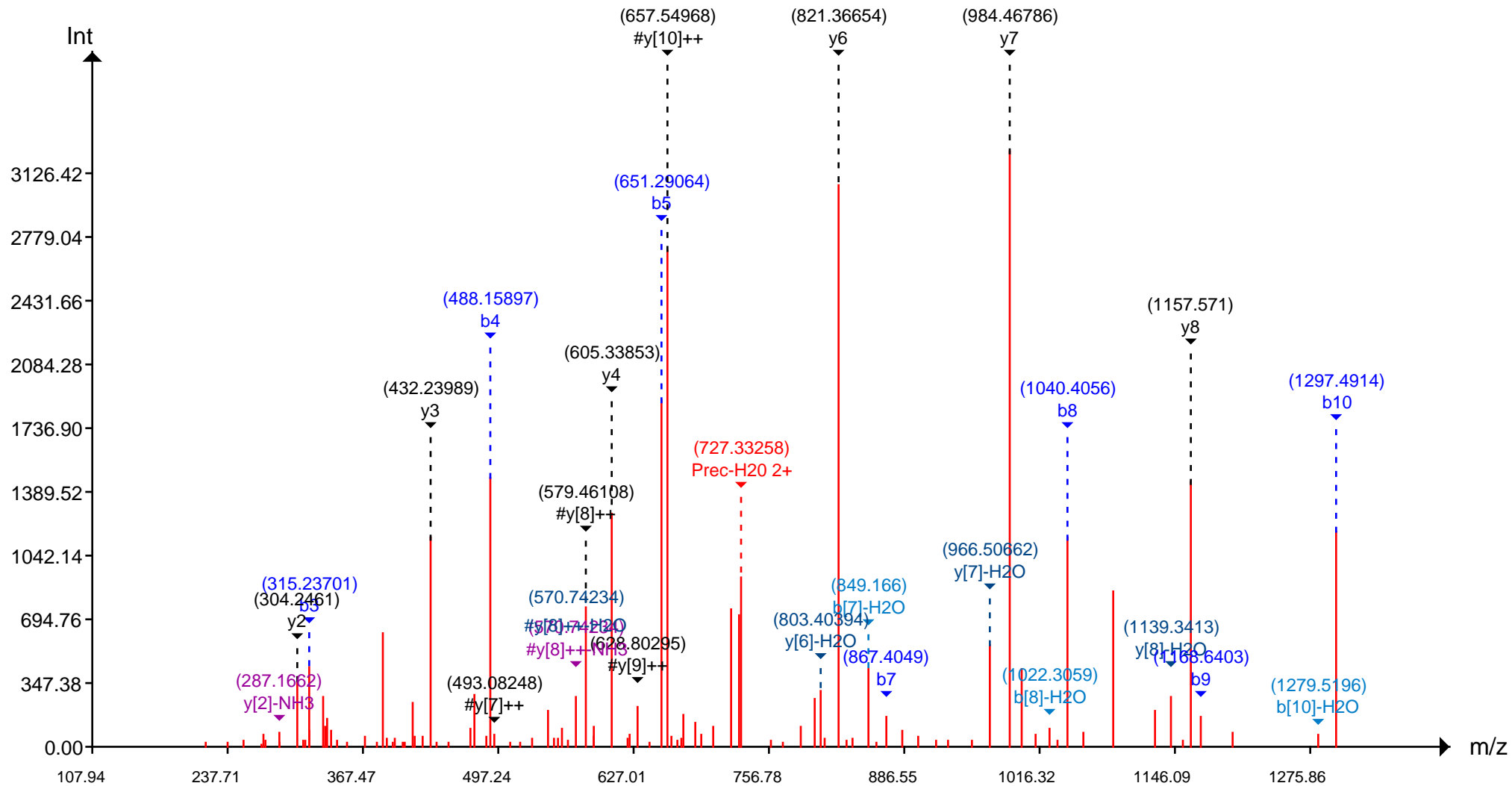
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 62.7 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32362 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 2.42E-4 |



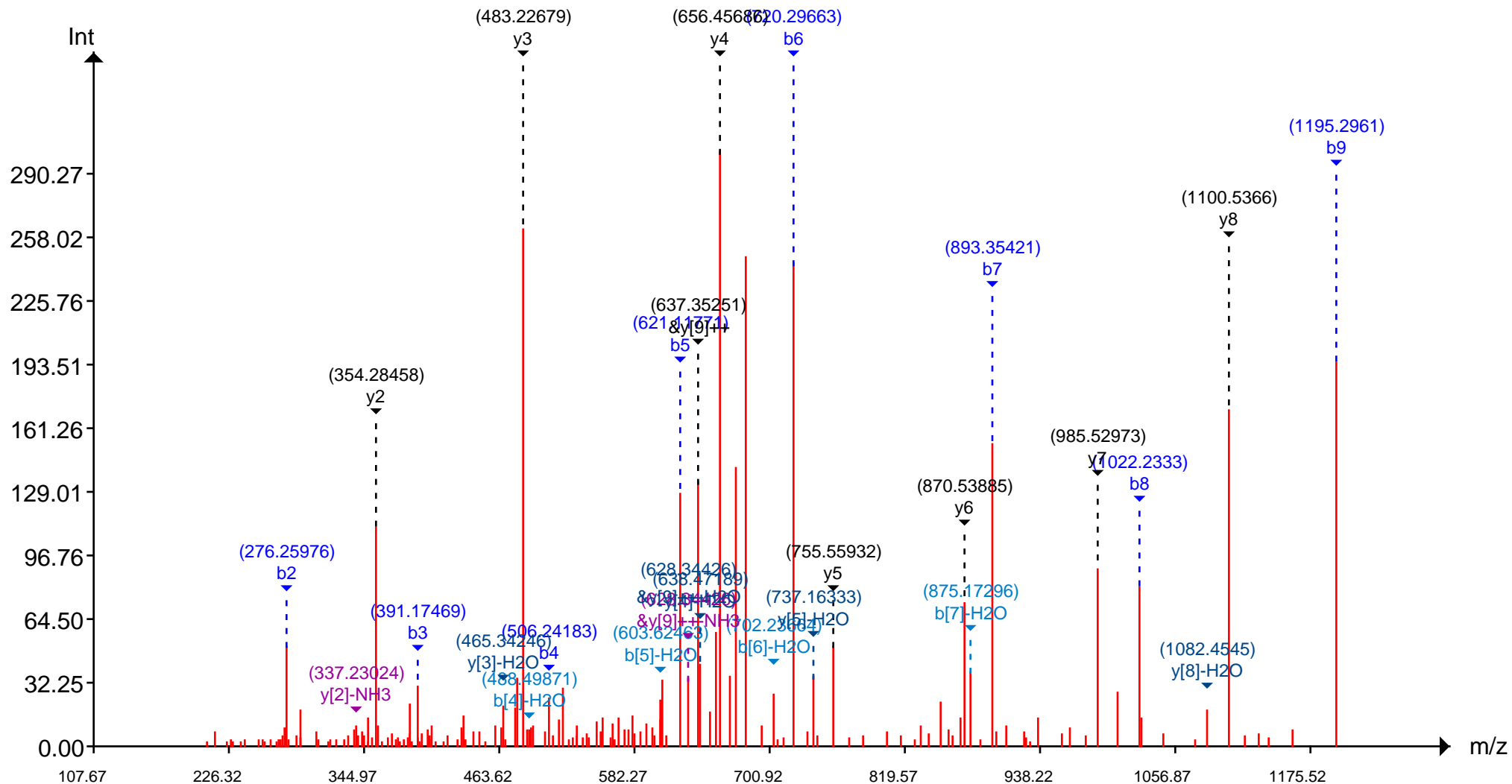
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 72.74 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.32362 | Identity Threshold | 47.96 |
| Charge | 2+ | Mass error | 2.42E-4 |



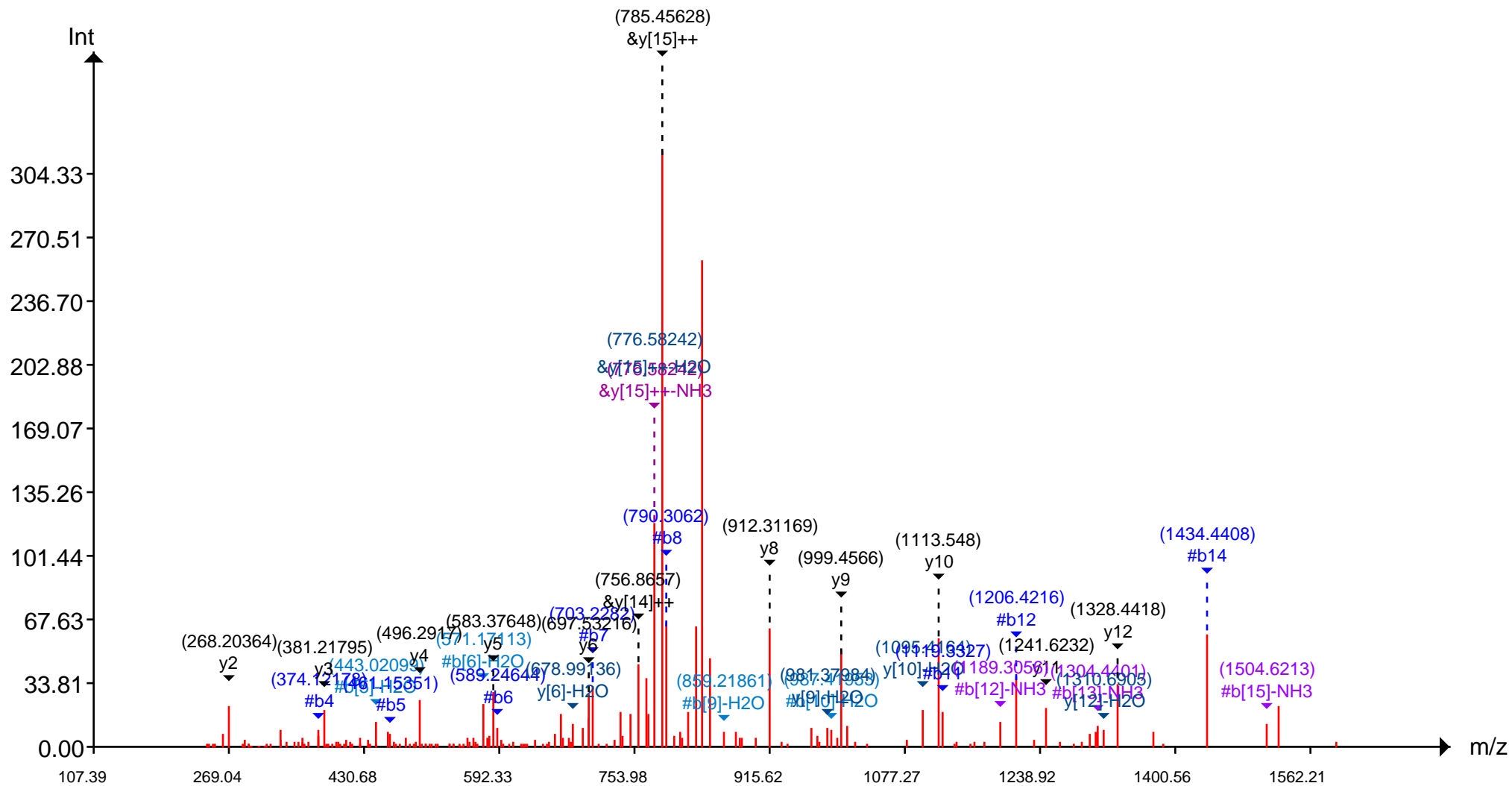
Info

| | | | |
|--------------------------|---------------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-IGVK<AcD3K*>YSEK<AcD3K*>QER-COOH | Ion score | 48.99 |
| Protein | Q99439 | Confidence level | 99.00% |
| Precursor m/z | 736.40908 | Identity Threshold | 48.38 |
| Charge | 2+ | Mass error | 7.82E-4 |



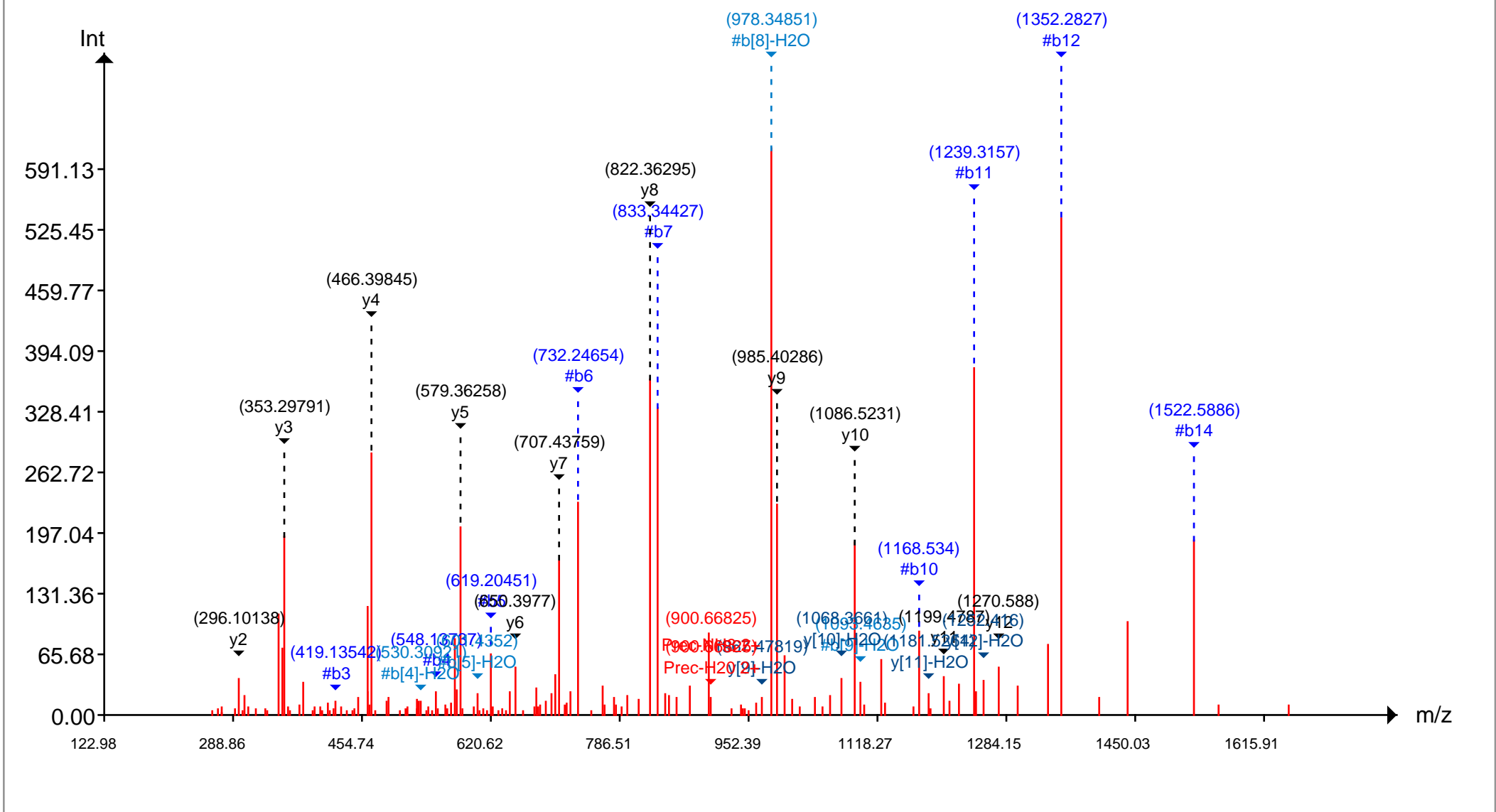
Info

| | | | |
|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-GK<AcD3K*>DDDVK<AcD3K*>EK<AcD3K*>R<C13>-COOH | Ion score | 51.5 |
| Protein | O94913 | Confidence level | 99.00% |
| Precursor m/z | 688.38076 | Identity Threshold | 48.78 |
| Charge | 2+ | Mass error | -6.21E-4 |



Info

| | | | |
|--------------------------|--------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-SGPSSQNSSQNSDLR<C13>-COOH | Ion score | 84.19 |
| Protein | Q96N21 | Confidence level | 99.00% |
| Precursor m/z | 851.39451 | Identity Threshold | 46.26 |
| Charge | 2+ | Mass error | 8.32E-4 |



Info

Modified Sequence AcD3-SLK<AcD3K*>EALTYDGALLGDR<C13>-COOH

Ion score 65.32

Protein Q15056

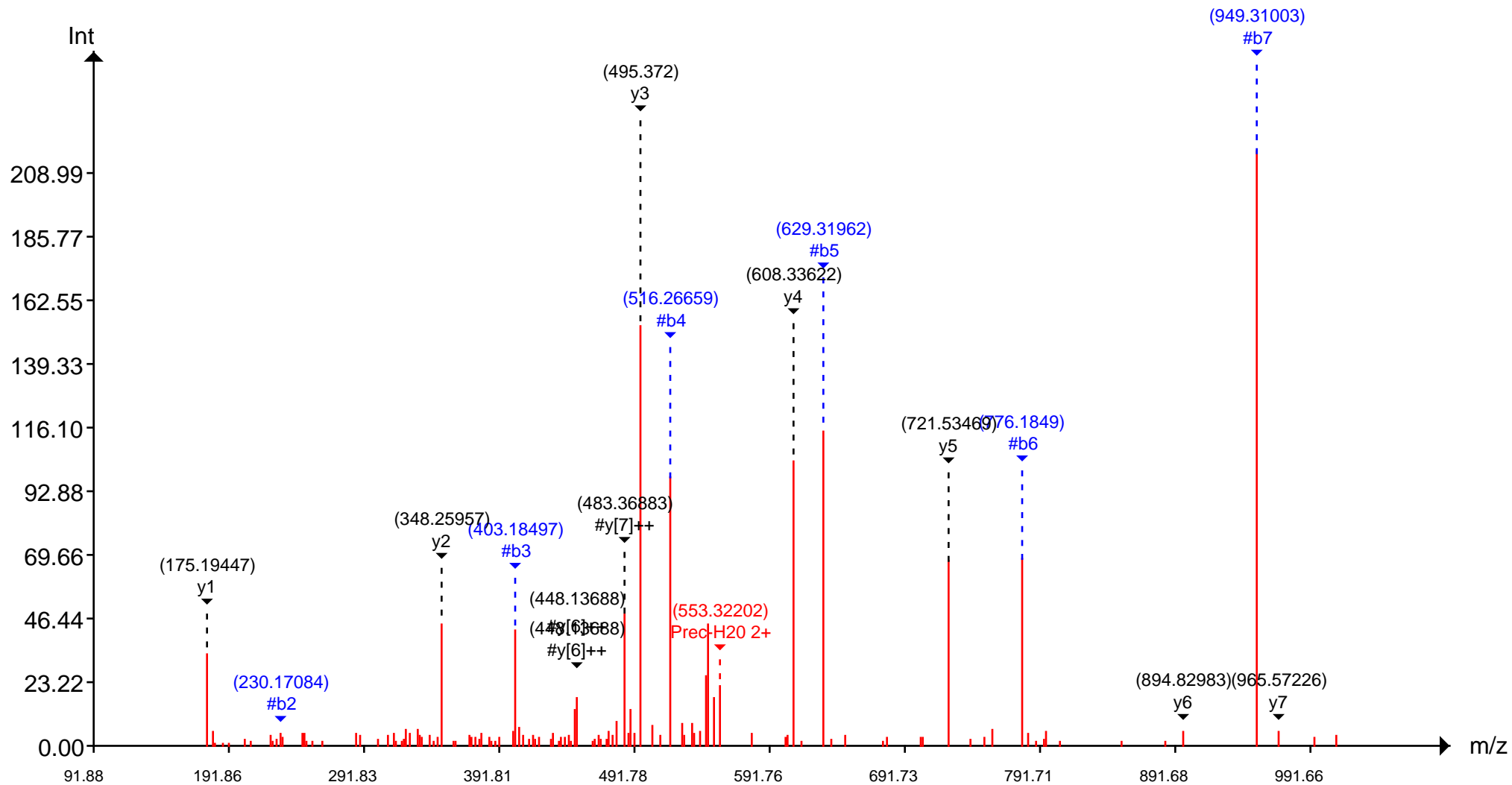
Confidence level 99.00%

Precursor m/z 909.49715

Identity Threshold 48.50

Charge 2+

Mass error 0.001278



Info

Modified Sequence AcD3-LAK<AcD3K*>IIFK<AcD3K*>R-COOH

Ion score 48.04

Protein P49321

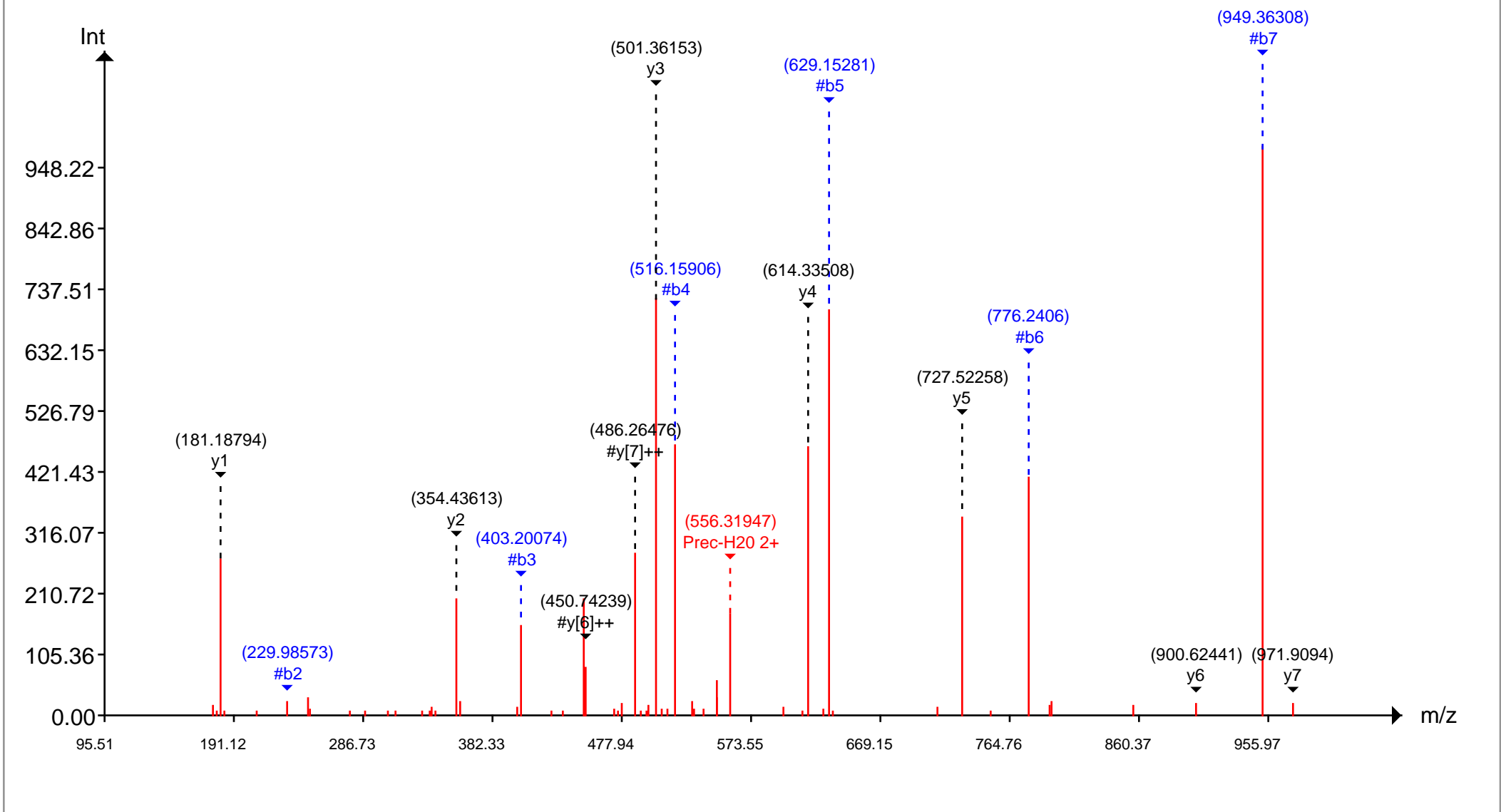
Confidence level 99.00%

Precursor m/z 562.38095

Identity Threshold 35.61

Charge 2+

Mass error -1.34E-4



Info

Modified Sequence AcD3-LAK<AcD3K*>IIFK<AcD3K*>R<C13>-COOH

Ion score 51.65

Protein P49321

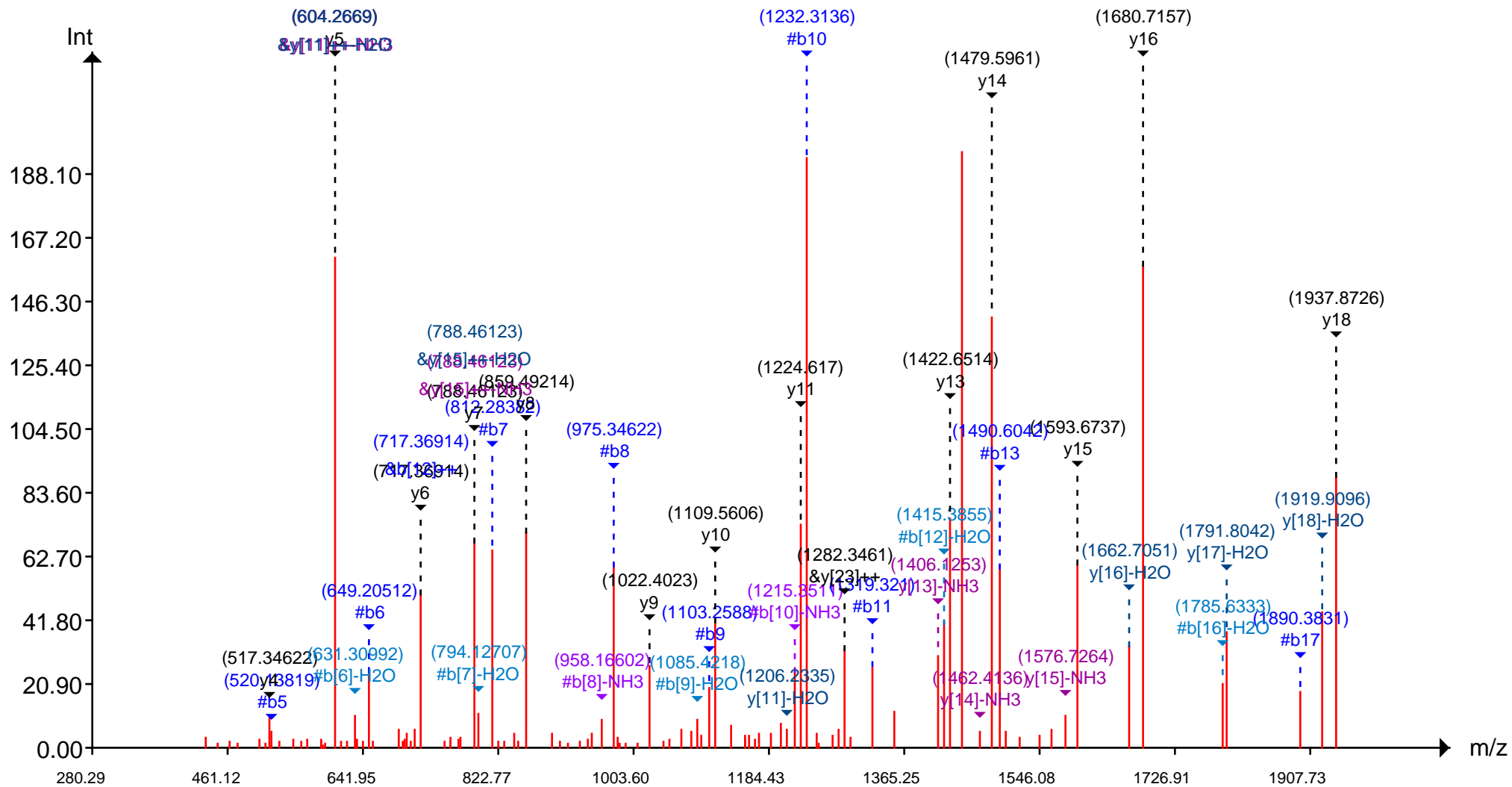
Confidence level 99.00%

Precursor m/z 565.39103

Identity Threshold 32.94

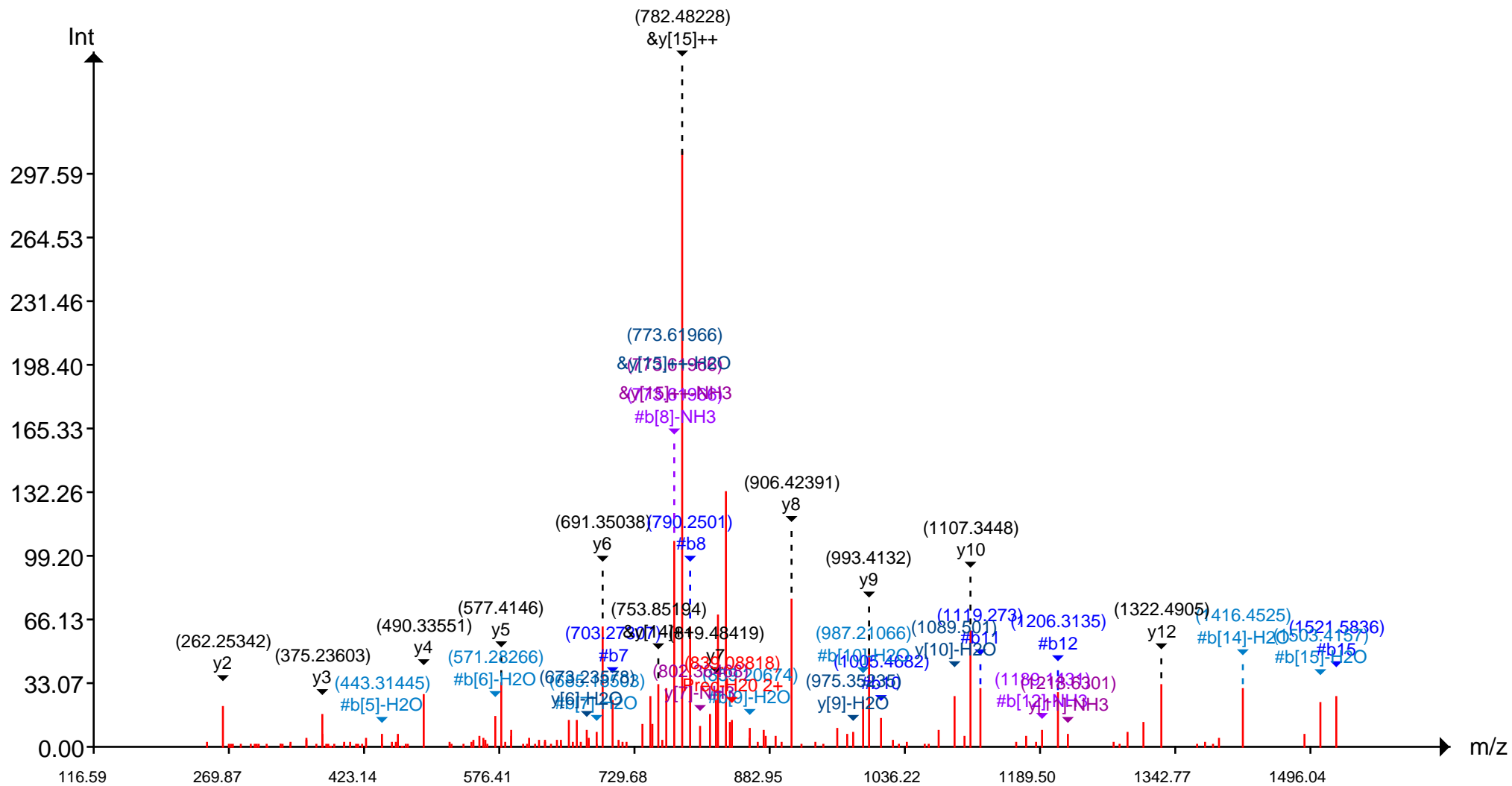
Charge 2+

Mass error -1.16E-4



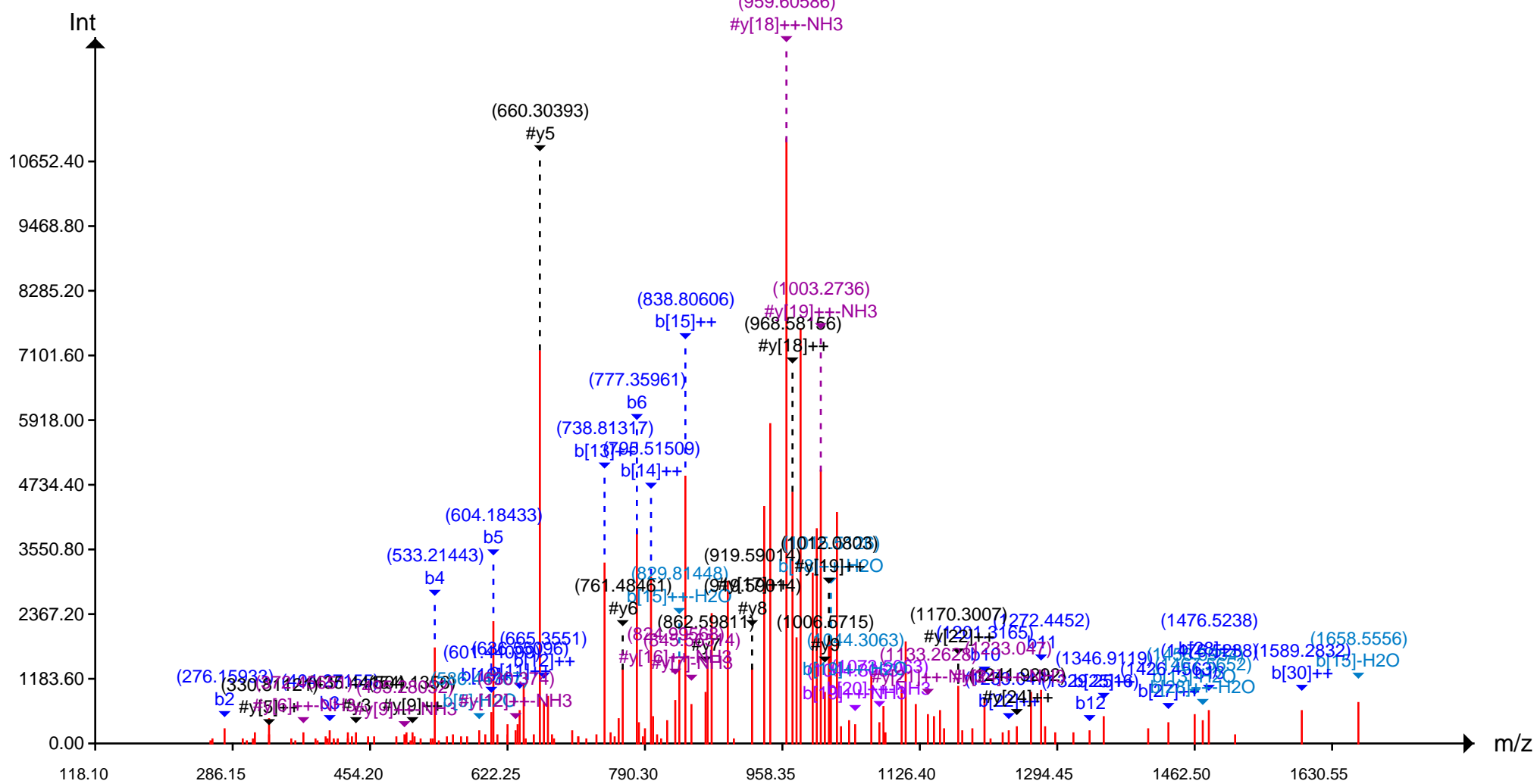
Info

| | | | |
|--------------------------|--|---------------------------|-----------|
| Modified Sequence | AcD3-GVM<Mox*>NGEYYQESNGPTDSYAAISQVDR-COOH | Ion score | 126.03 |
| Protein | P09496 | Confidence level | 99.00% |
| Precursor m/z | 1456.638 | Identity Threshold | 41.48 |
| Charge | 2+ | Mass error | -0.003109 |

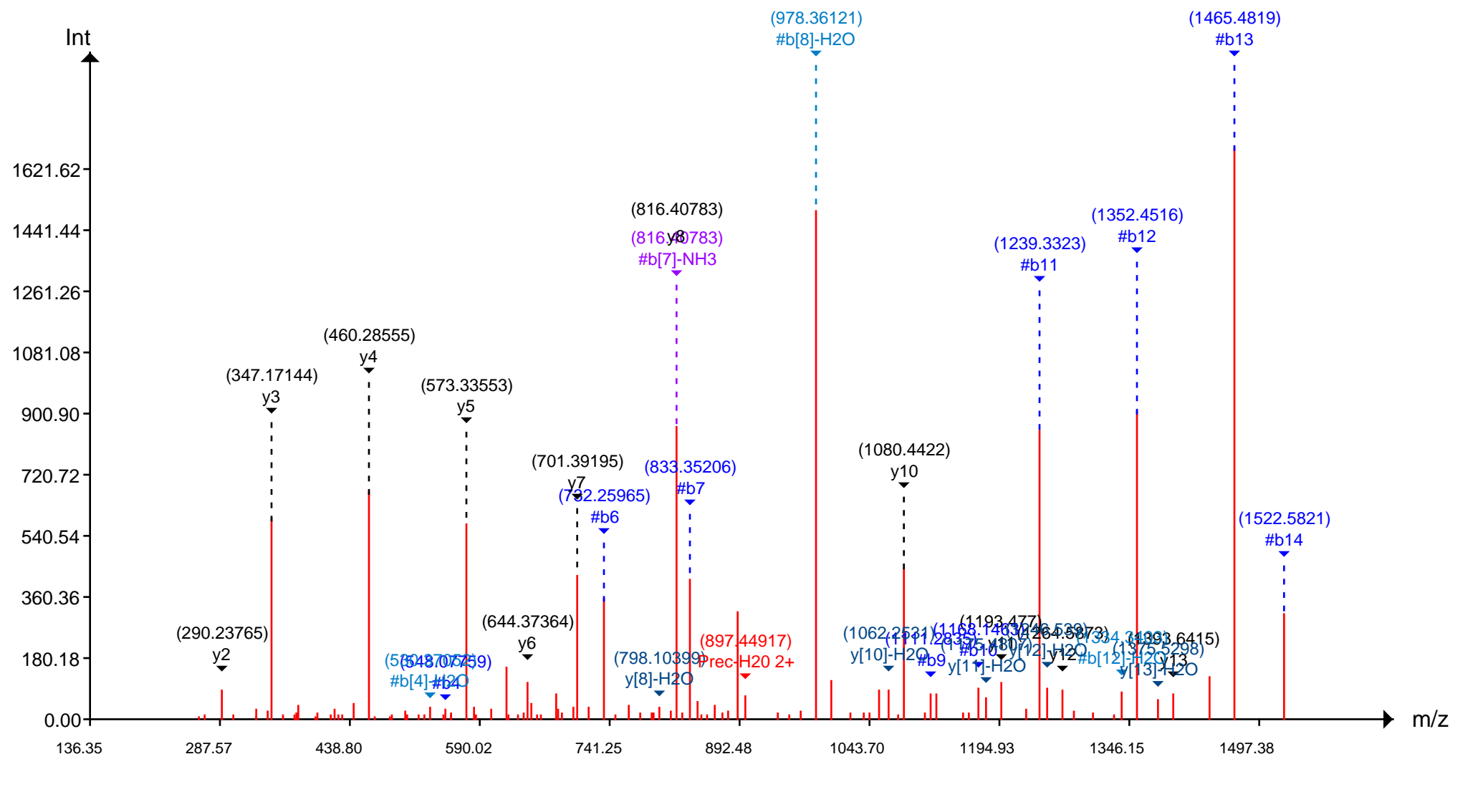


Info

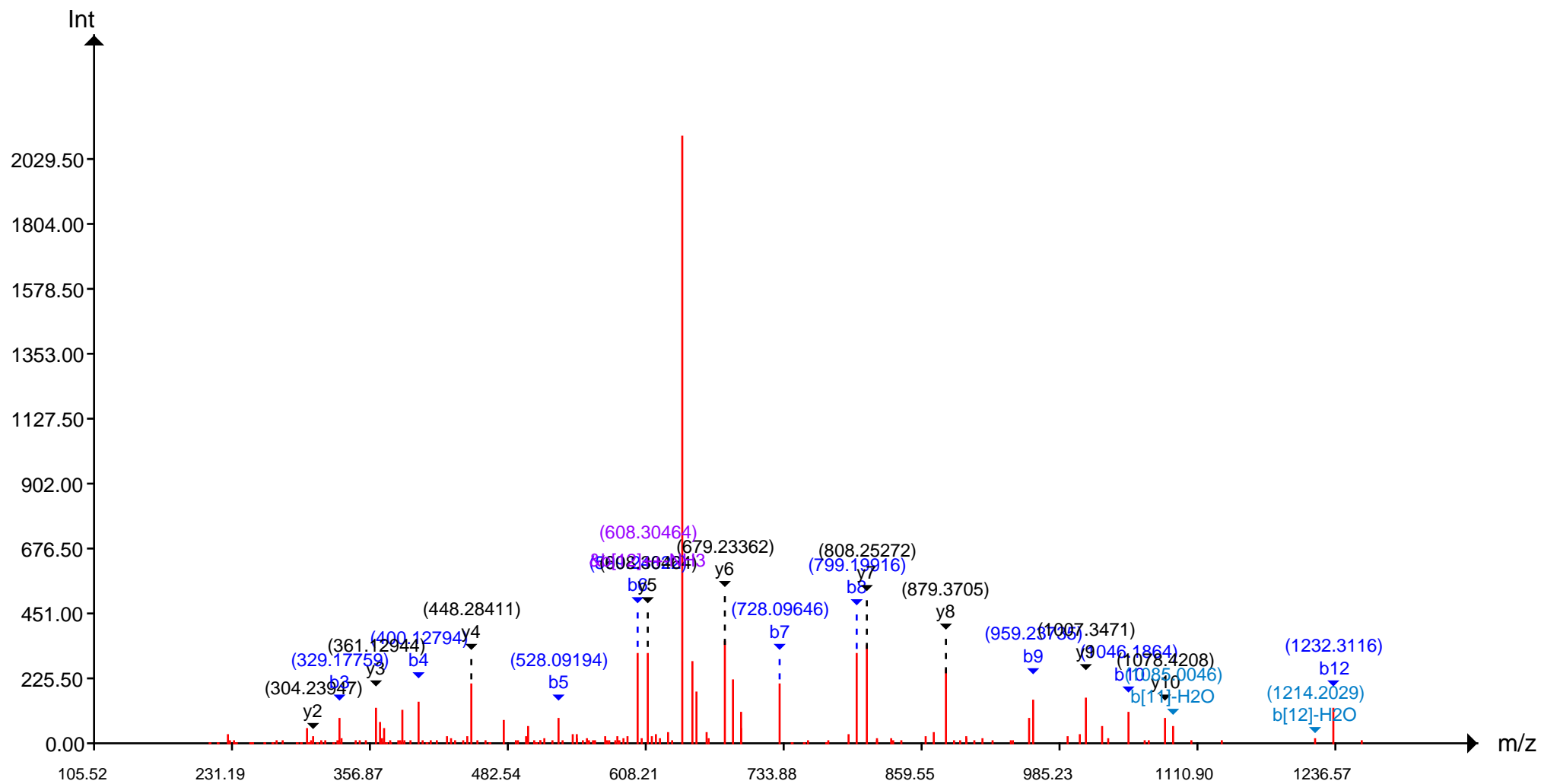
| | | | |
|--------------------------|---------------------------|---------------------------|---------|
| Modified Sequence | AcD3-SGPSSQNSSQNSDLR-COOH | Ion score | 87.08 |
| Protein | Q96N21 | Confidence level | 99.00% |
| Precursor m/z | 848.384 | Identity Threshold | 45.18 |
| Charge | 2+ | Mass error | -4.6E-5 |



| Info | |
|-----------|--|
| Mo... | AcD3-GK<AcD3K*>QEAK<AcD3K*>PQQAAGM<Mox*>LSPK<AcD3K*>TGGK<AcD3K*>EAASGTPQK<AcD3K*>SR-COOH |
| | Ion score |
| Q13428 | Confidence level |
| 903.73039 | Identity Threshold |
| 4+ | Mass error |



| Info | | | |
|--------------------------|------------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-SLK<AcD3K*>EALTYDGALLGDR-COOH | Ion score | 86.58 |
| Protein | Q15056 | Confidence level | 99.00% |
| Precursor m/z | 906.48705 | Identity Threshold | 48.88 |
| Charge | 2+ | Mass error | 0.00122 |



Info

Modified Sequence AcD3-GPEAQAEAC<Cmm*>SGER-COOH

Ion score 64.47

Protein Q9NX24

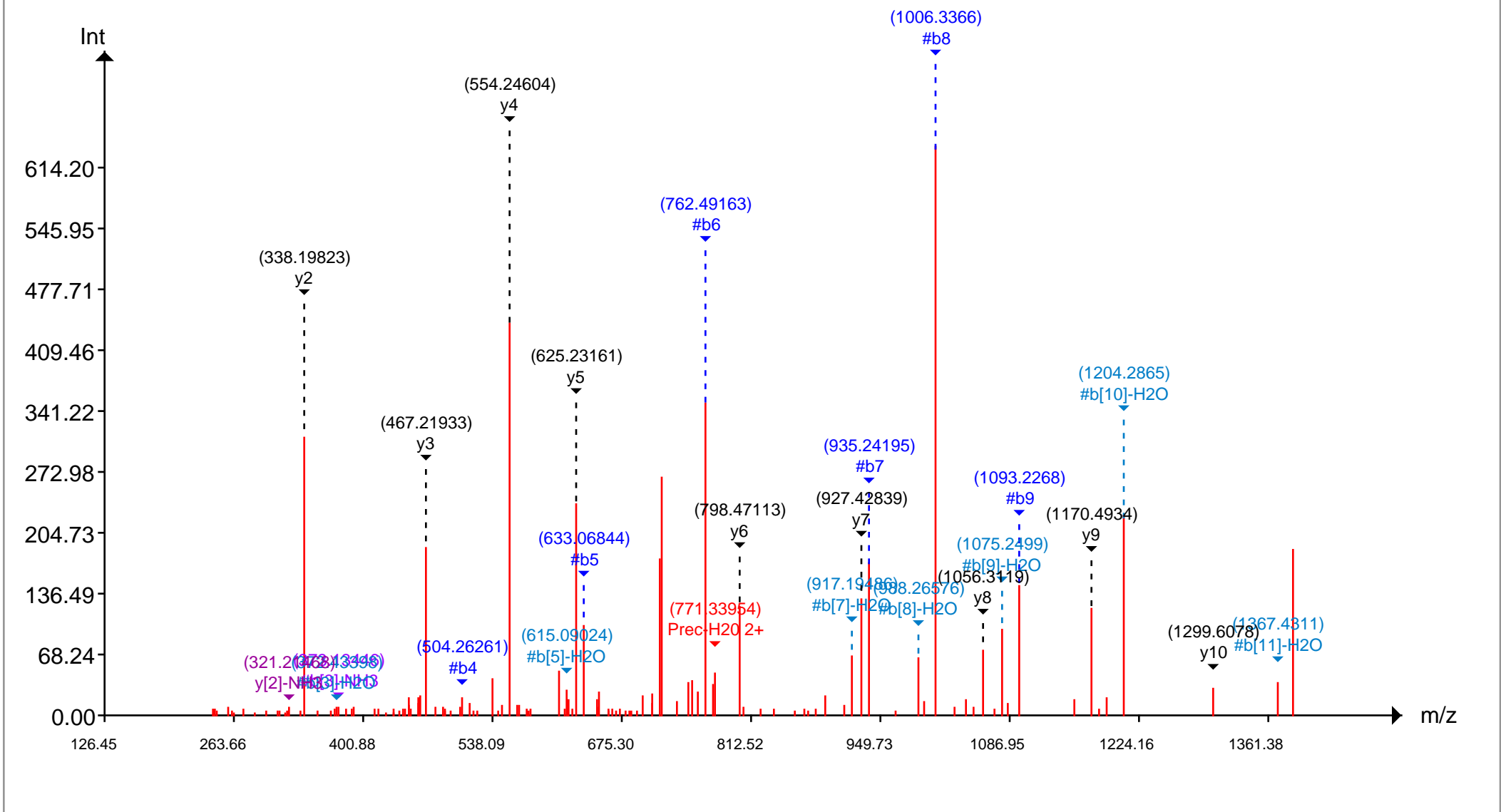
Confidence level 99.00%

Precursor m/z 703.80601

Identity Threshold 43.21

Charge 2+

Mass error 3.61E-4



Info

Modified Sequence AcD3-SQENEEK<AcD3K*>ASEYR-COOH

Ion score 68.89

Protein O94782

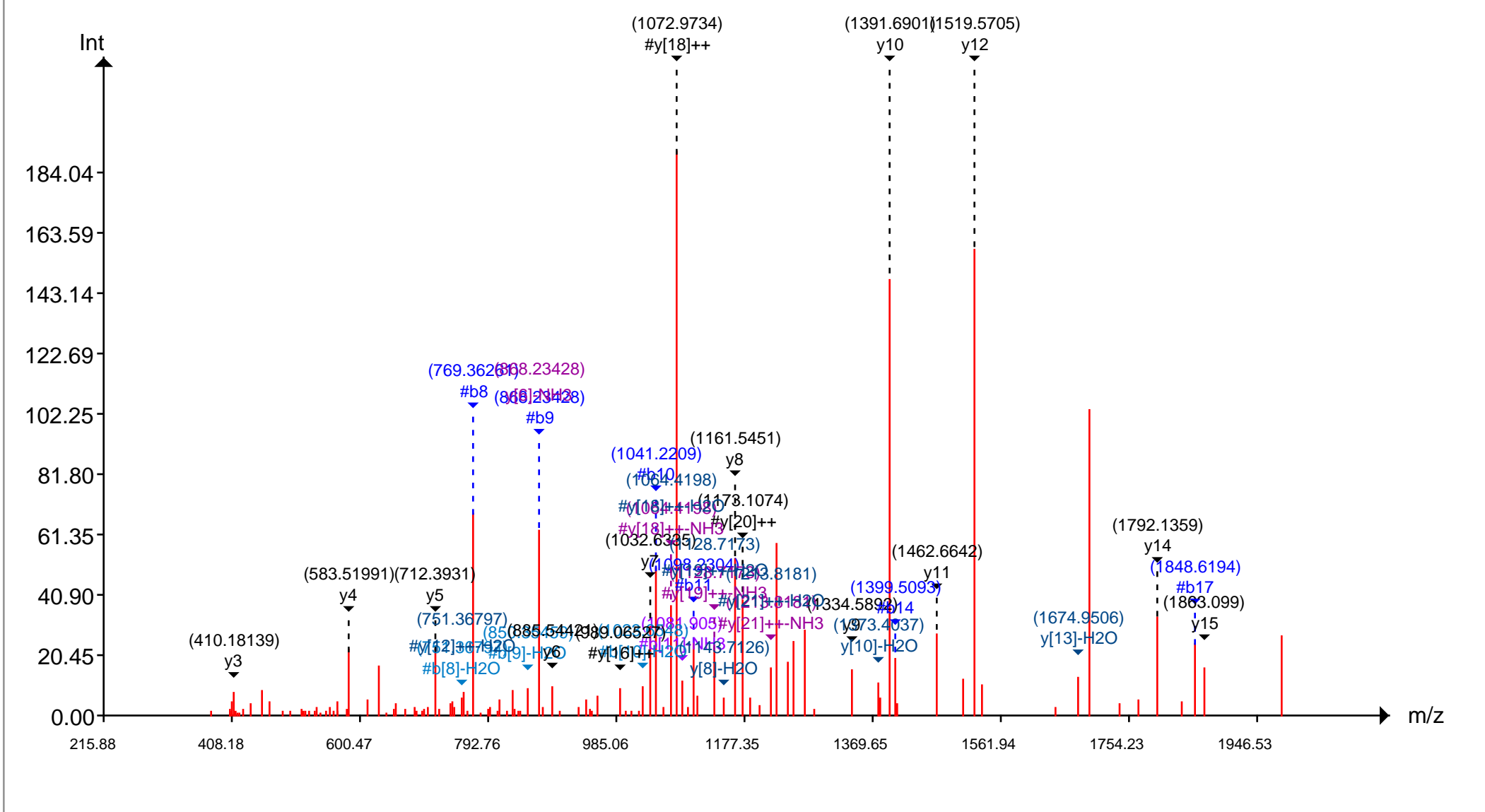
Confidence level 99.00%

Precursor m/z 780.35824

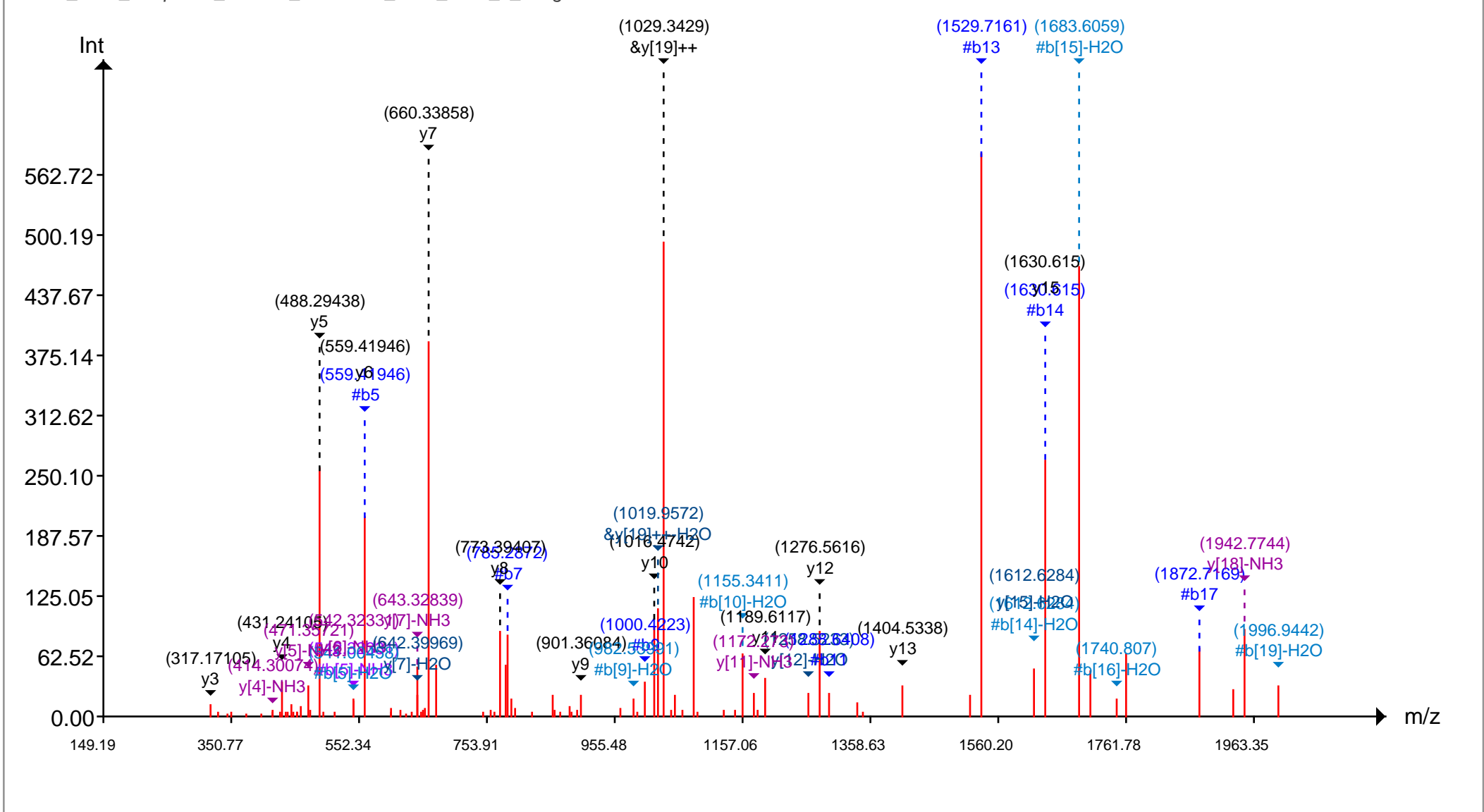
Identity Threshold 45.66

Charge 2+

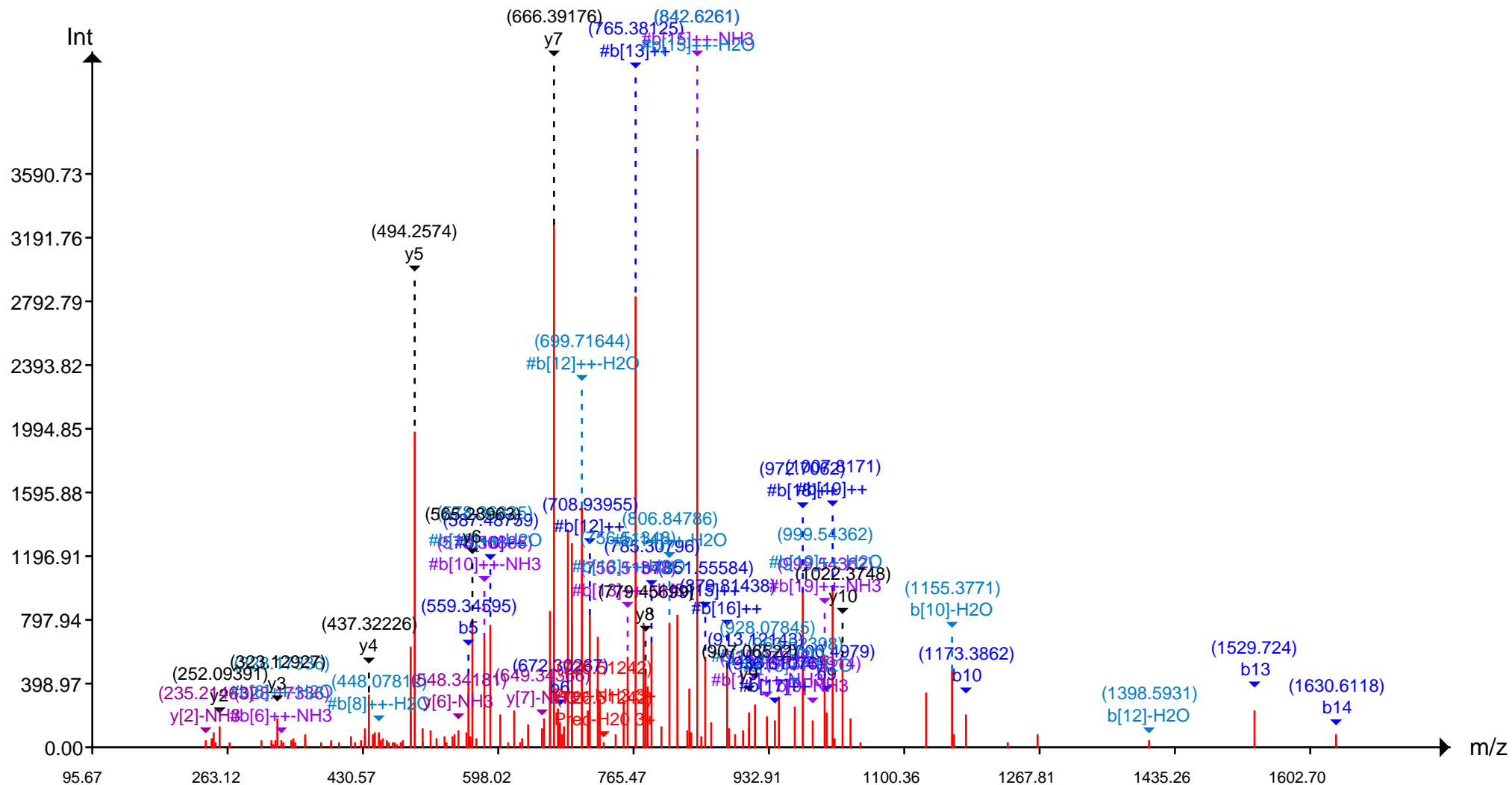
Mass error 1.76E-4



| Info | | | |
|----------------|---|--------------------|------|
| Modified Se... | AcD3-AVAEPANAVK<AcD3K*>GAGK<AcD3K*>EM<Mox*>K<AcD3K*>EK<AcD3K*>TQR<C13>-COOH | Ion score | 62.2 |
| ... | Q9HCF4 | Confidence level | 99.0 |
| Precurs... | 1280.7044 | Identity Threshold | 47.2 |
| ... | 2+ | Mass error | 0.00 |

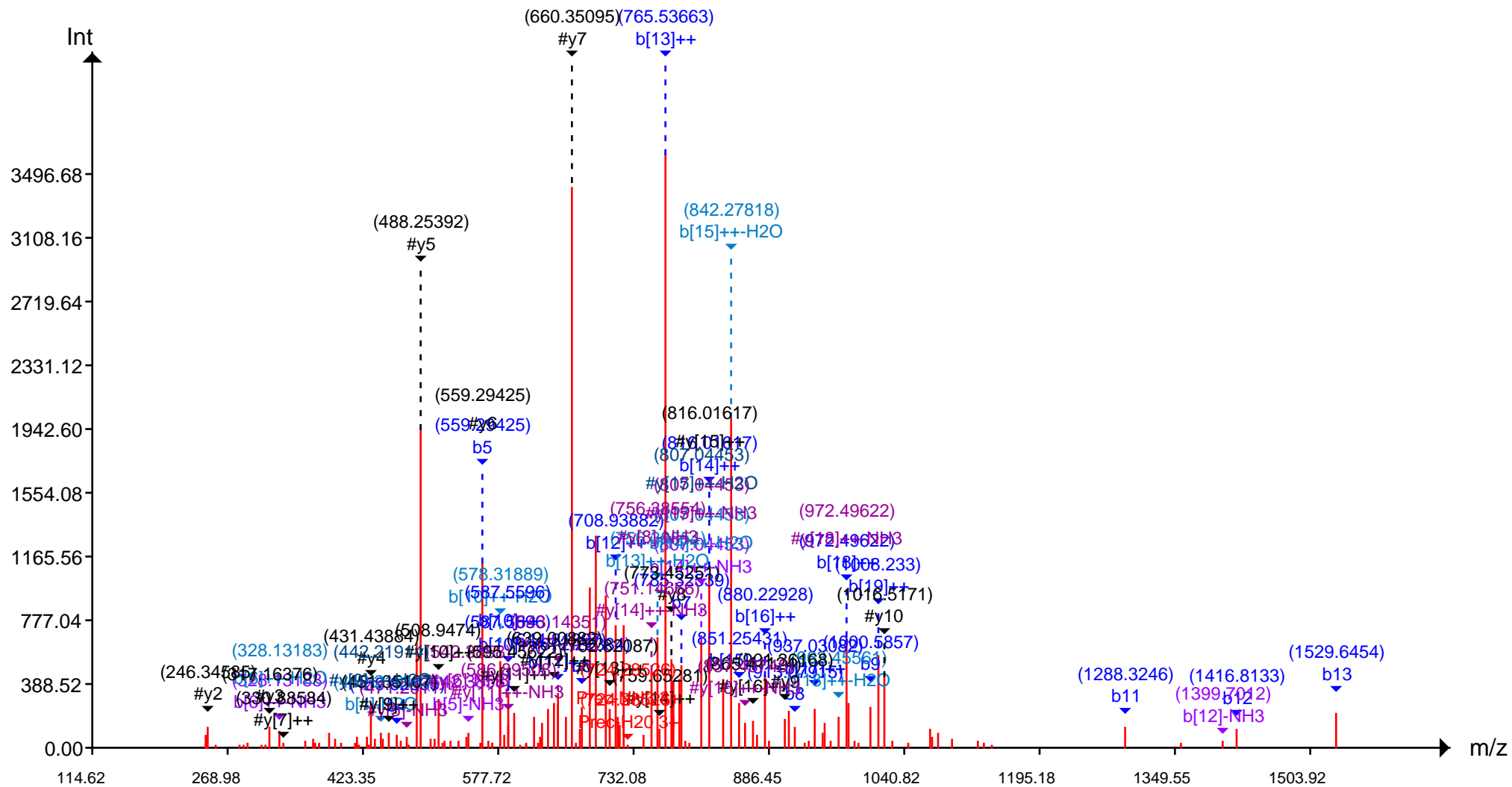


| Info | | | |
|--------------------------|--|---------------------------|-----------|
| Modified Sequence | AcD3-SPGK<AcD3K*>VLLQSK<AcD3K*>DQITAGNAAR-COOH | Ion score | 103.94 |
| Protein | P22234 | Confidence level | 99.00% |
| Precursor m/z | 1095.1148 | Identity Threshold | 46.92 |
| Charge | 2+ | Mass error | -0.001123 |



Info

| | | | |
|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-SPGK<AcD3K*>VLLQSK<AcD3K*>DQITAGNAAR<C13>-COOH | Ion score | 53.23 |
| Protein | P22234 | Confidence level | 99.00% |
| Precursor m/z | 732.41921 | Identity Threshold | 45.90 |
| Charge | 3+ | Mass error | -5.11E-4 |



Info

Modified Sequence AcD3-SPGK<AcD3K*>VLLQSK<AcD3K*>DQITAGNAAR-COOH

Ion score 63.35

Protein P22234

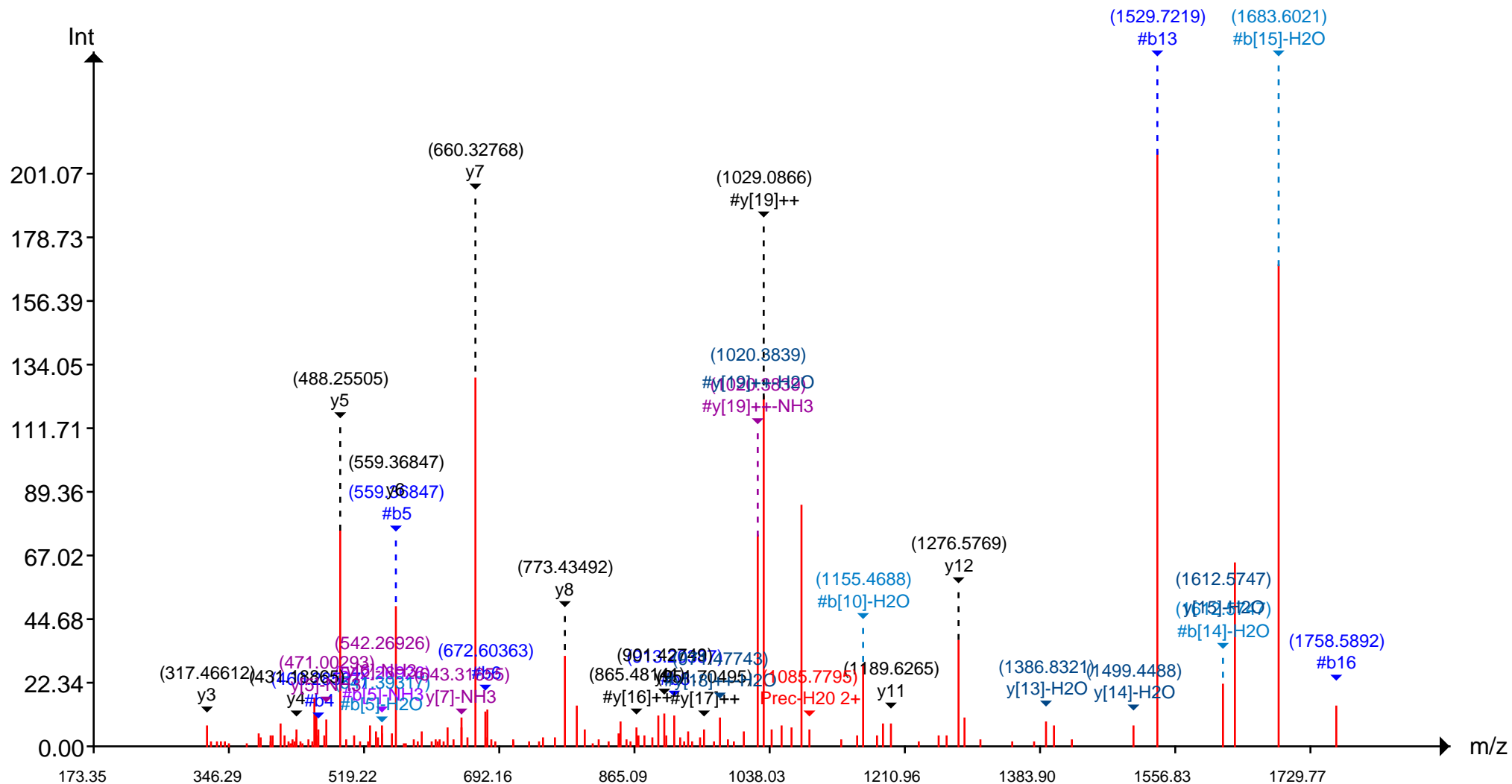
Confidence level 99.00%

Precursor m/z 730.4126

Identity Threshold 46.89

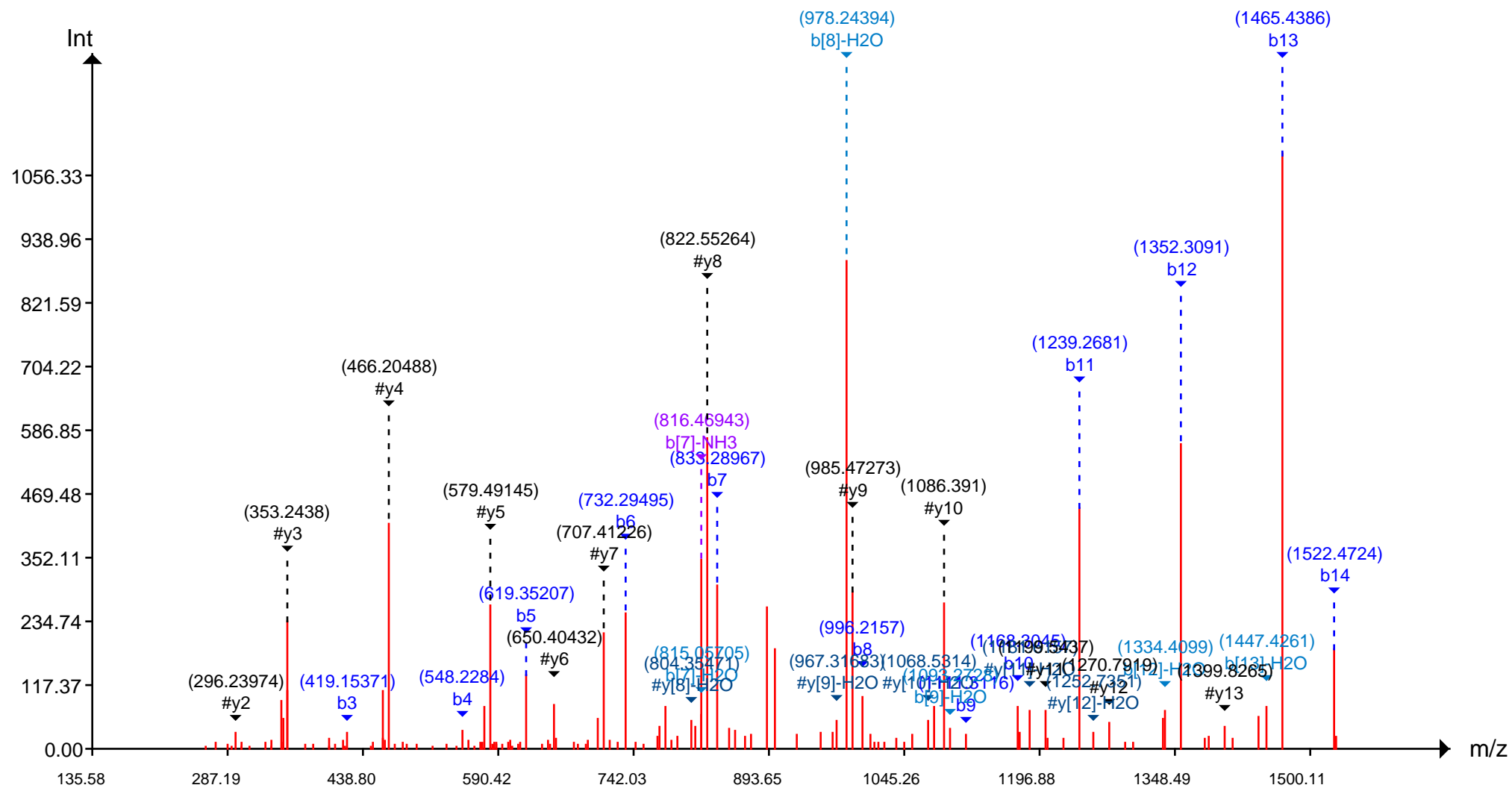
Charge 3+

Mass error -1.99E-4



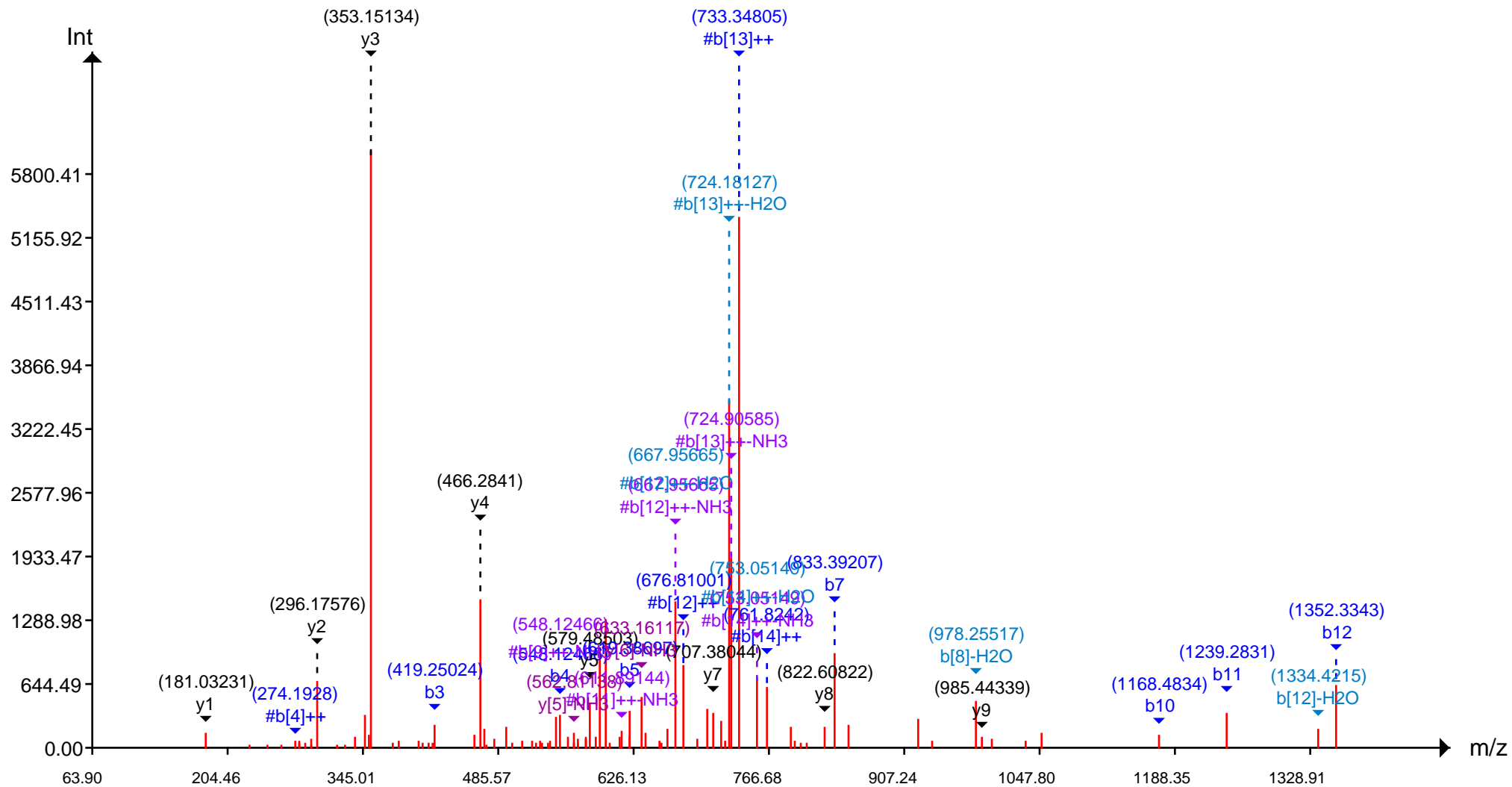
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-SPGK<AcD3K*>VLLQSK<AcD3K*>DQITAGNAAR-COOH | Ion score | 50.97 |
| Protein | P22234 | Confidence level | 99.00% |
| Precursor m/z | 1095.1158 | Identity Threshold | 46.81 |
| Charge | 2+ | Mass error | 8.77E-4 |



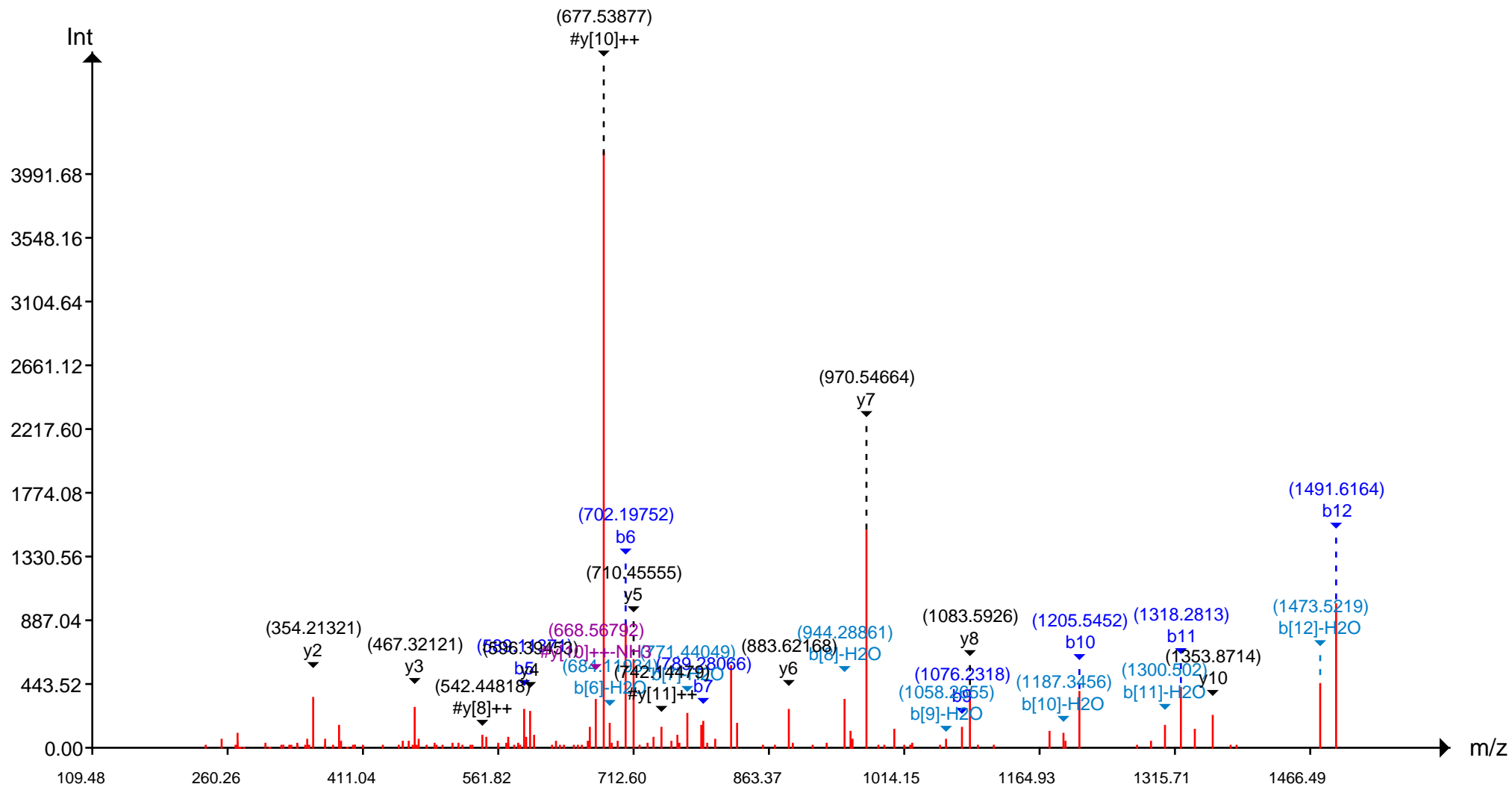
Info

| | | | |
|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-SLK<AcD3K*>EALTYDGALLGDR<C13>-COOH | Ion score | 81.41 |
| Protein | Q15056 | Confidence level | 99.00% |
| Precursor m/z | 909.49637 | Identity Threshold | 48.54 |
| Charge | 2+ | Mass error | -2.82E-4 |



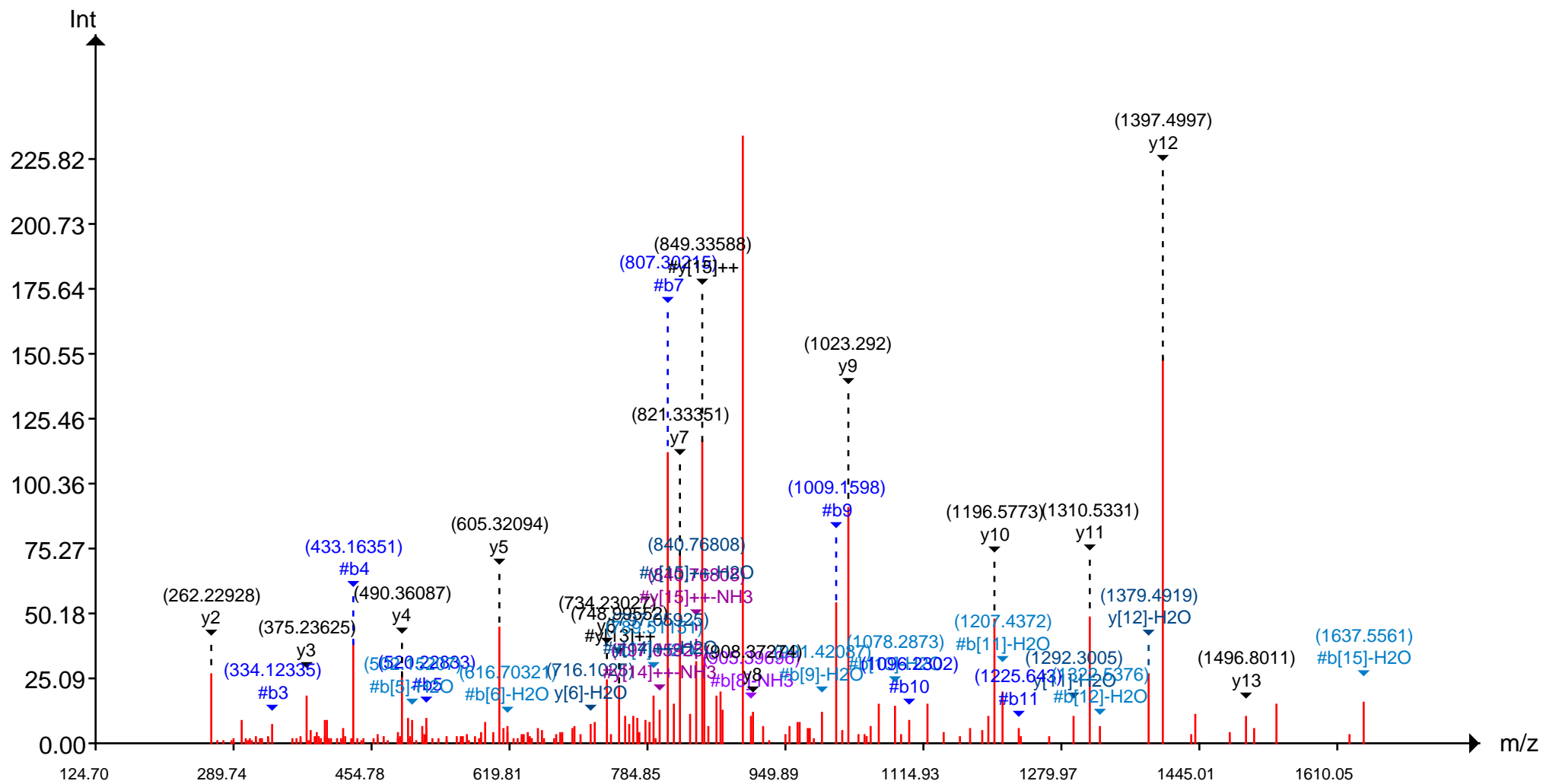
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-SLK<AcD3K*>EALTYDGALLGDR<C13>-COOH | Ion score | 57.25 |
| Protein | Q15056 | Confidence level | 99.00% |
| Precursor m/z | 606.66694 | Identity Threshold | 48.51 |
| Charge | 3+ | Mass error | 5.22E-4 |

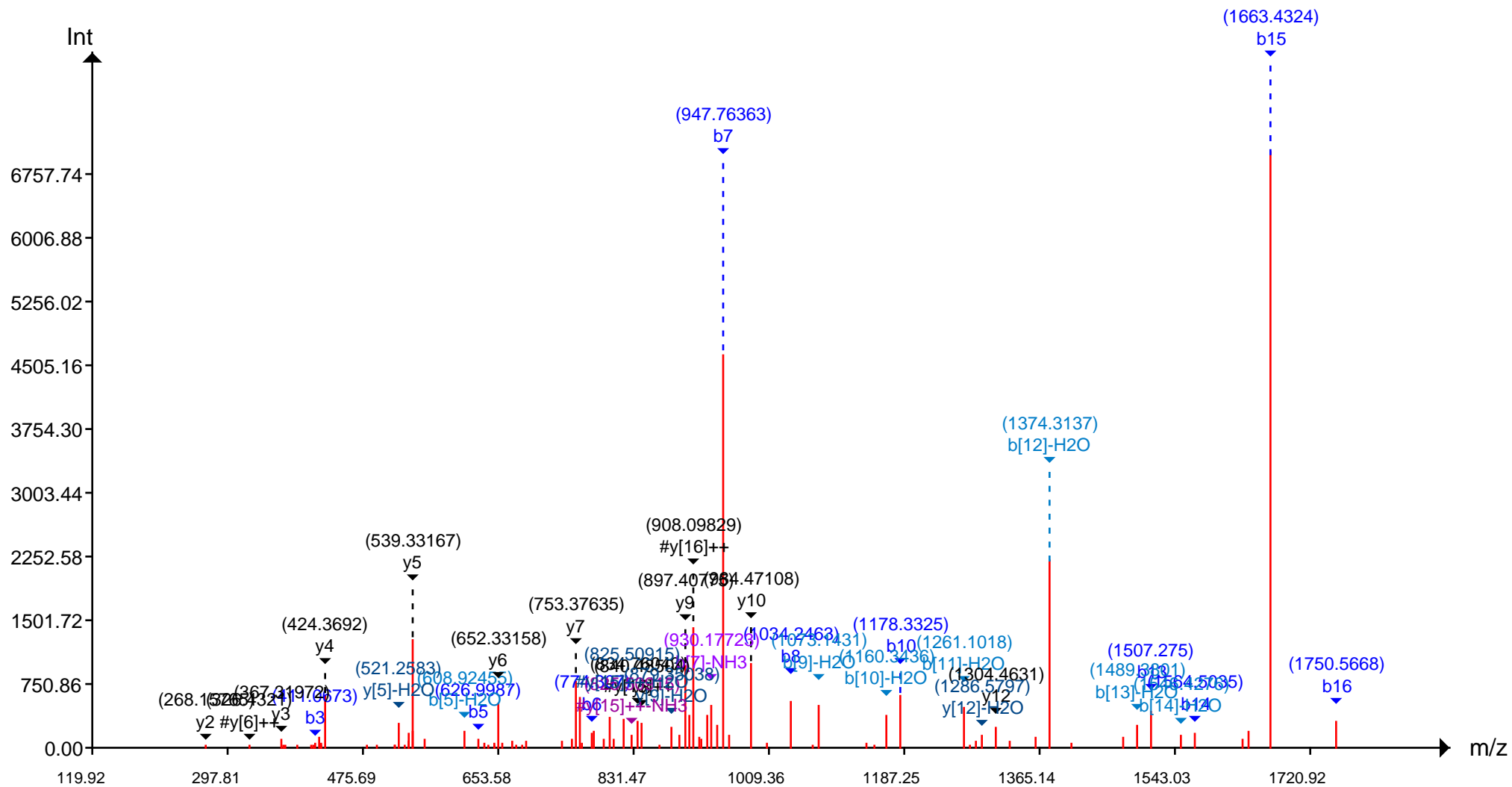


Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GSEPK<AcD3K*>LSK<AcD3K*>NELK<AcD3K*>R<C13>-COOH | Ion score | 50.67 |
| Protein | Q15046 | Confidence level | 99.00% |
| Precursor m/z | 836.49218 | Identity Threshold | 45.41 |
| Charge | 2+ | Mass error | 0.001027 |

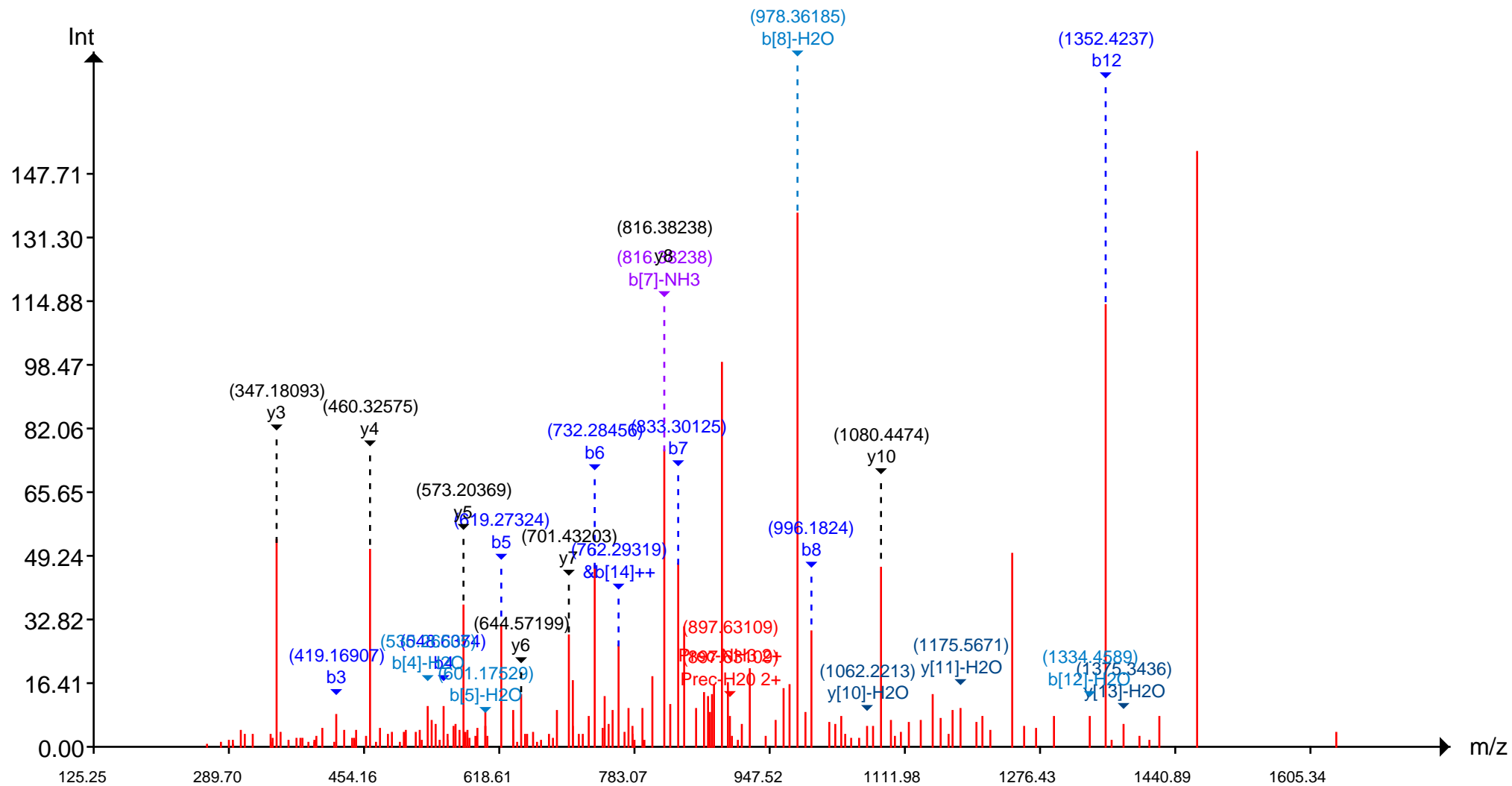


| Info | | | |
|--------------------------|------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-SSNVSNK<AcD3K*>DSSEDDISR-COOH | Ion score | 82.92 |
| Protein | Q5W0B1 | Confidence level | 99.00% |
| Precursor m/z | 915.41752 | Identity Threshold | 45.57 |
| Charge | 2+ | Mass error | 0.001152 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-AFM<Mox*>SEM<Mox*>K<AcD3K*>SGSTLDGVSR<C13>-COOH | Ion score | 69.66 |
| Protein | Q9BWU0 | Confidence level | 99.00% |
| Precursor m/z | 965.95911 | Identity Threshold | 47.25 |
| Charge | 2+ | Mass error | 0.001034 |



Info

Modified Sequence AcD3-SLK<AcD3K*>EALTYDGALLGDR-COOH

Ion score 51.53

Protein Q15056

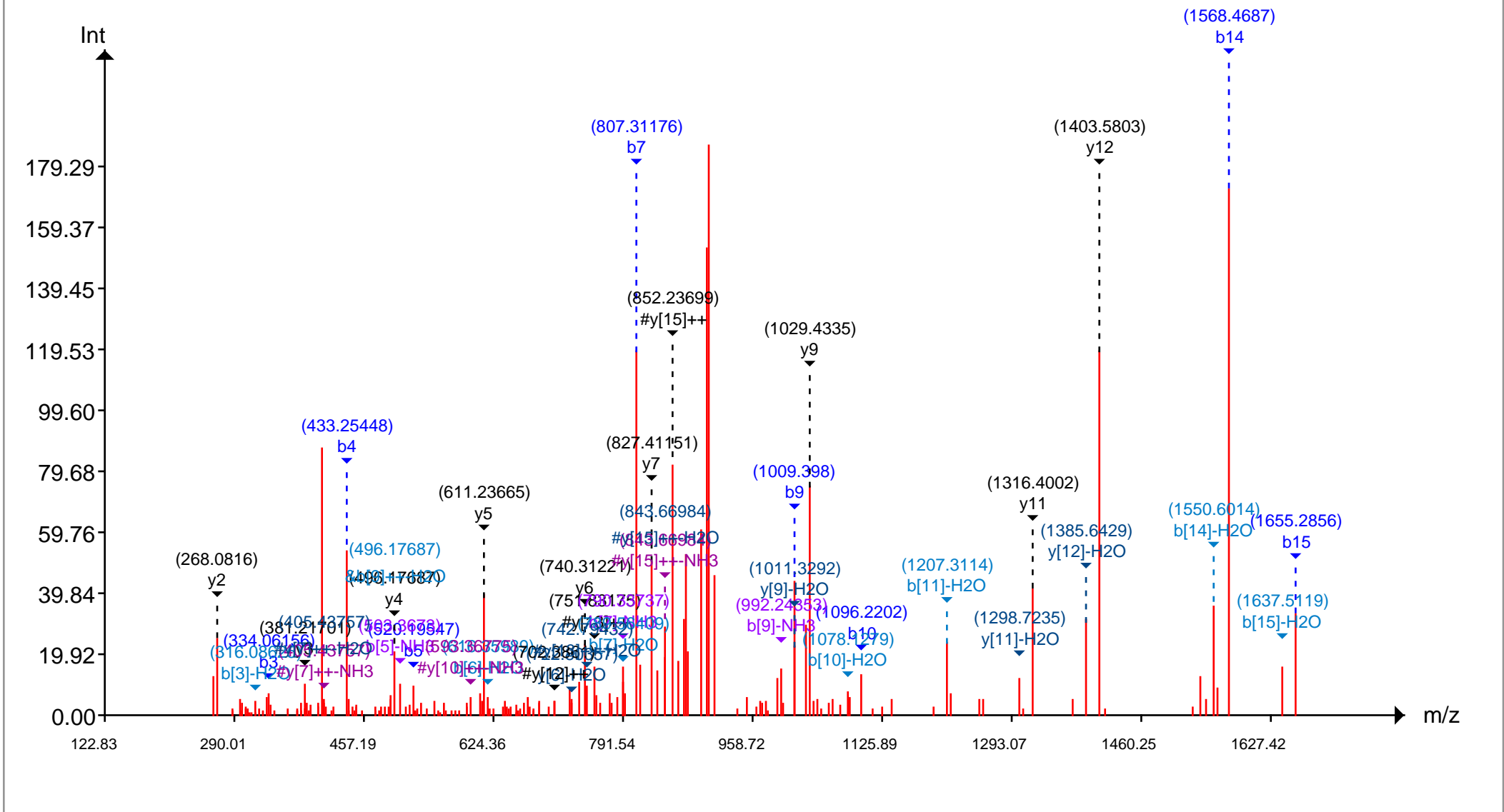
Confidence level 99.00%

Precursor m/z 906.48719

Identity Threshold 48.91

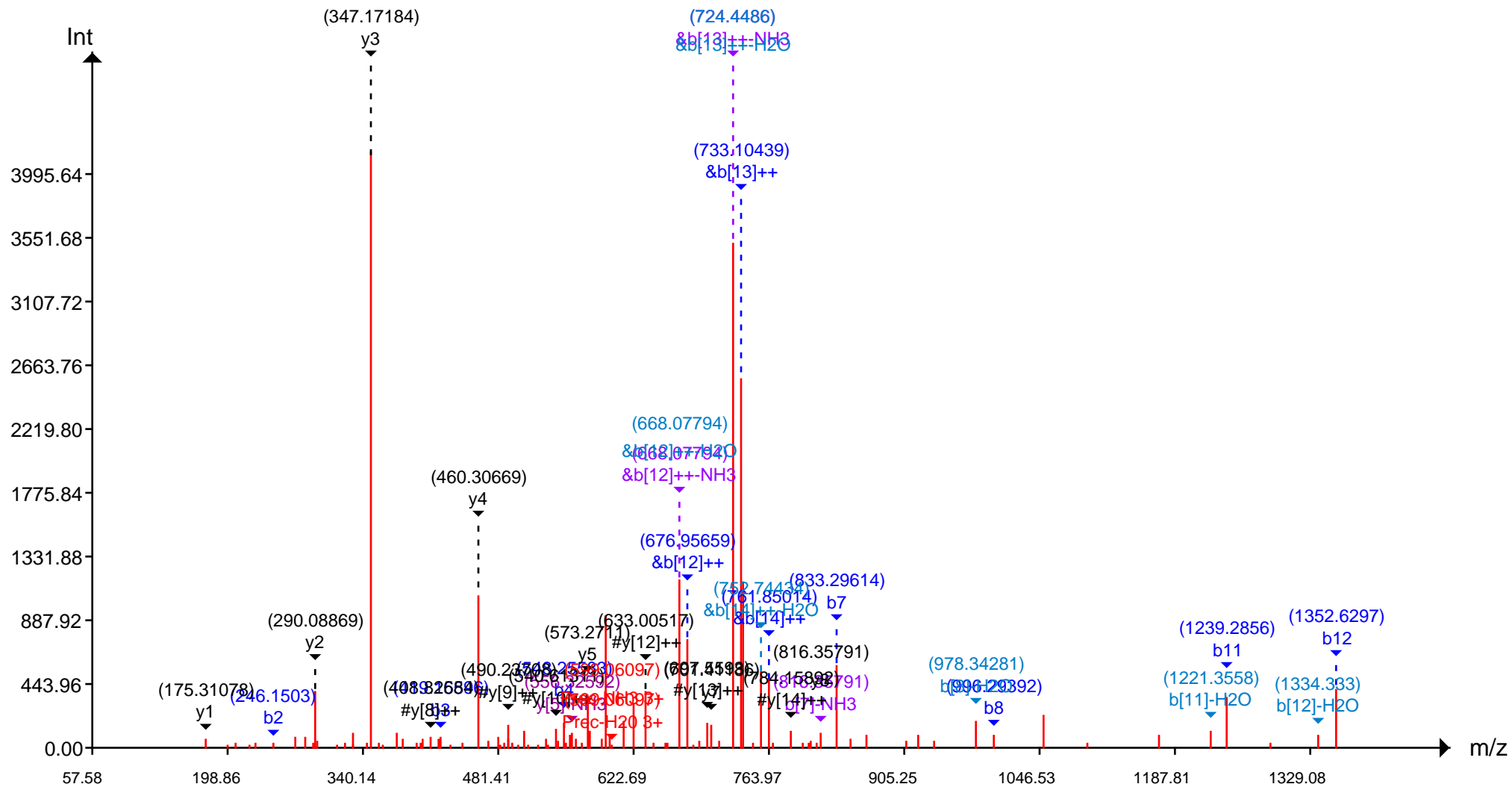
Charge 2+

Mass error 0.0015



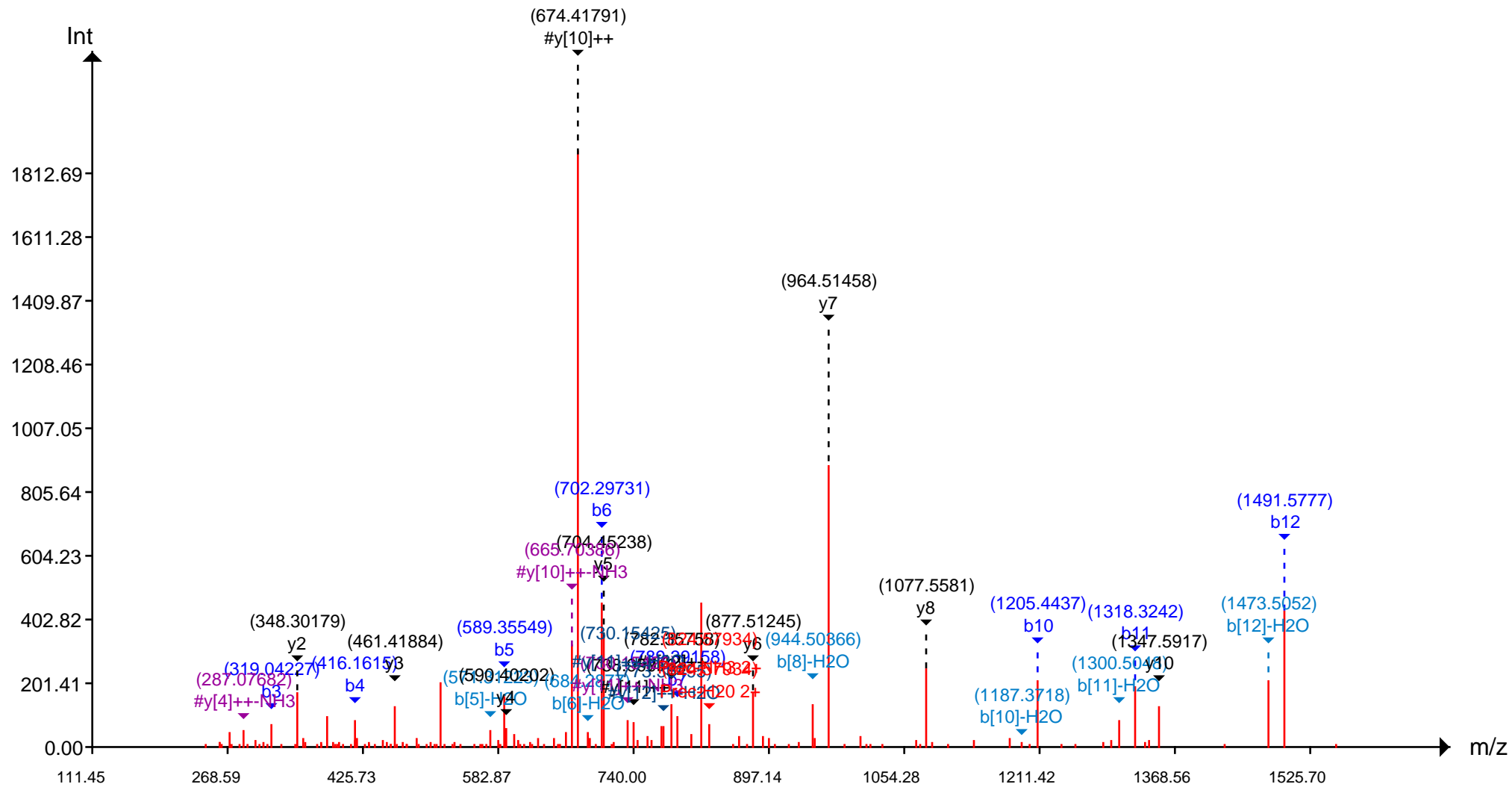
Info

| | | | |
|--------------------------|---|---------------------------|--------|
| Modified Sequence | AcD3-SSNVSNK<AcD3K*>DSSEDDISR<C13>-COOH | Ion score | 63.05 |
| Protein | Q5W0B1 | Confidence level | 99.00% |
| Precursor m/z | 918.42724 | Identity Threshold | 46.41 |
| Charge | 2+ | Mass error | 4.5E-4 |



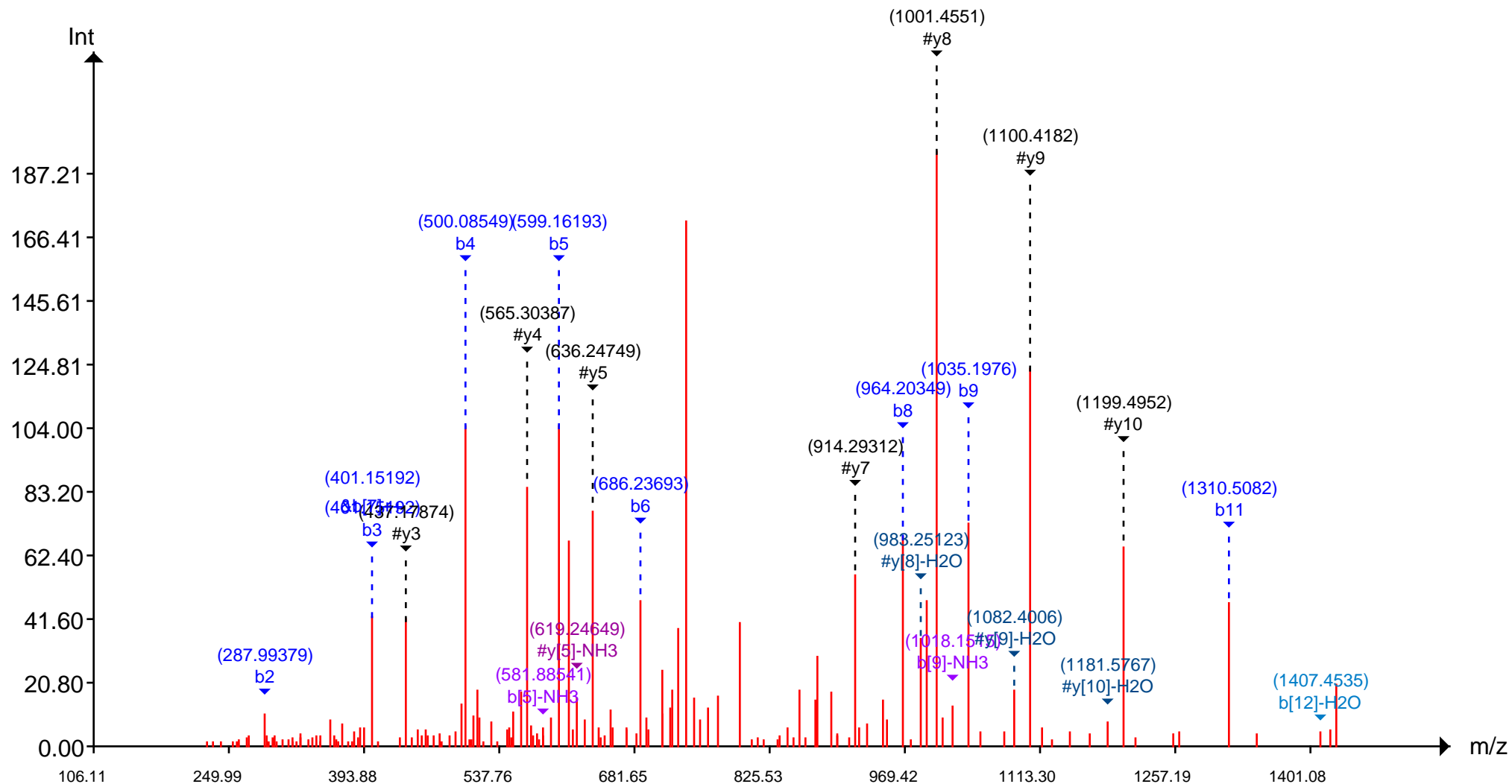
Info

| | | | |
|--------------------------|------------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-SLK<AcD3K*>EALTYDGALLGDR-COOH | Ion score | 51.47 |
| Protein | Q15056 | Confidence level | 99.00% |
| Precursor m/z | 604.66031 | Identity Threshold | 48.90 |
| Charge | 3+ | Mass error | 7.74E-4 |



Info

| | | | |
|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-GSEPK<AcD3K*>LSK<AcD3K*>NELK<AcD3K*>R-COOH | Ion score | 62.29 |
| Protein | Q15046 | Confidence level | 99.00% |
| Precursor m/z | 833.4824 | Identity Threshold | 46.33 |
| Charge | 2+ | Mass error | 0.001609 |



Info

Modified Sequence AcD3-NQIVVSDYAQM<Mox*>DR-COOH

Ion score 56.24

Protein P50991

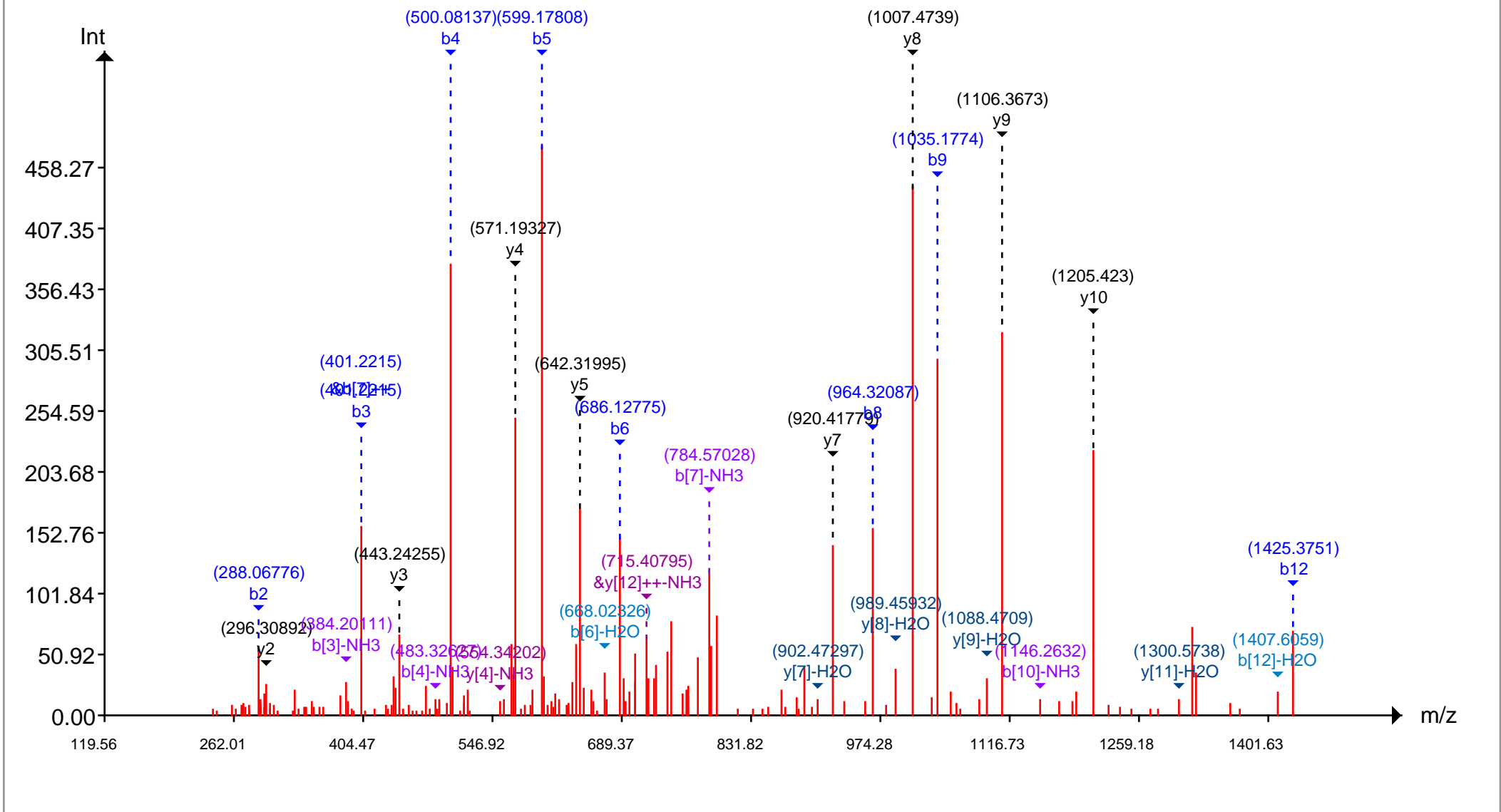
Confidence level 99.00%

Precursor m/z 800.37868

Identity Threshold 46.97

Charge 2+

Mass error -9.82E-4



Info

Modified Sequence AcD3-NQIVVSDYAQM<Mox*>DR<C13>-COOH

Ion score 69.17

Protein P50991

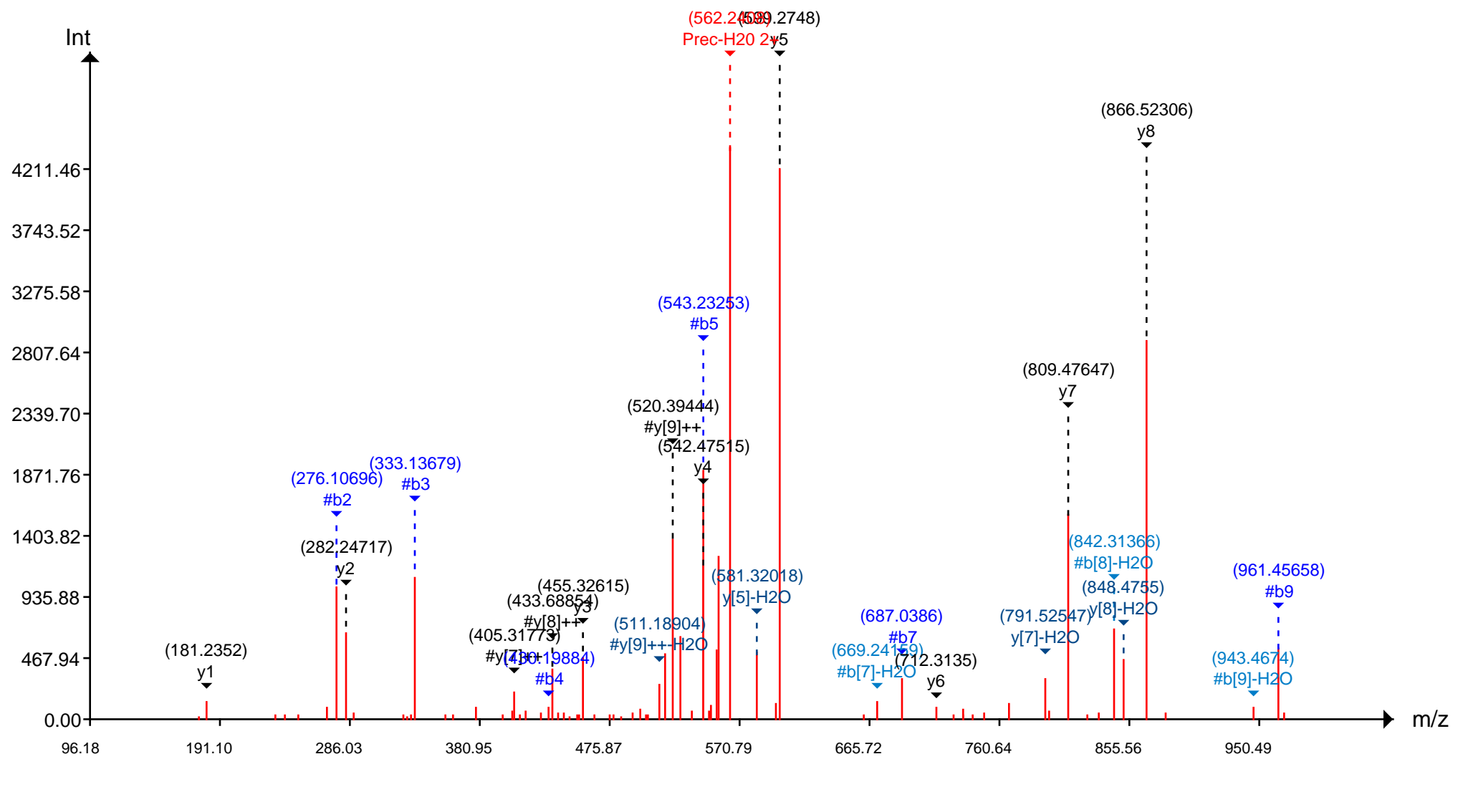
Confidence level 99.00%

Precursor m/z 803.38952

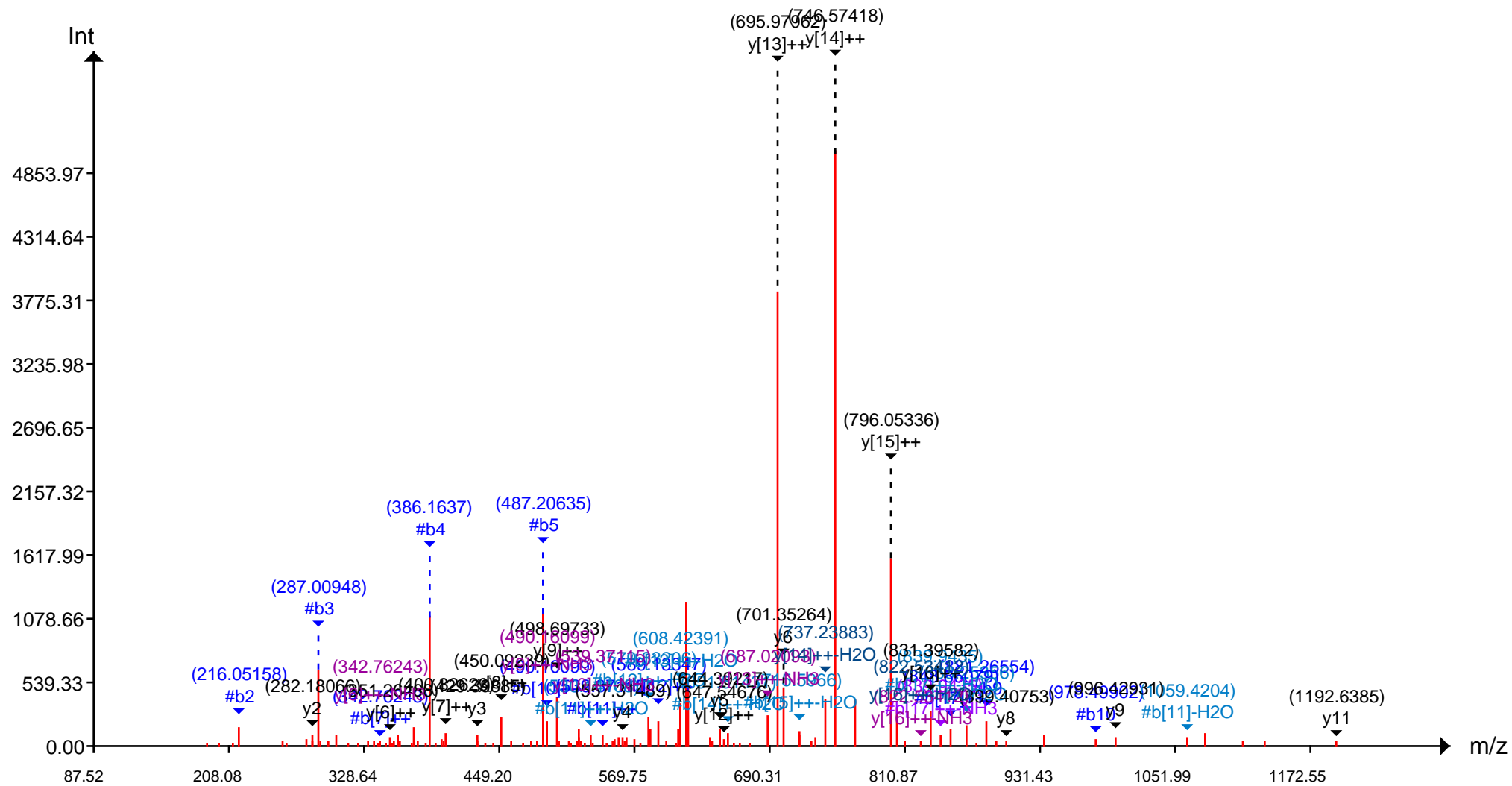
Identity Threshold 47.91

Charge 2+

Mass error 5.57E-4



| Info | | | |
|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-GK<AcD3K*>GPIGSK<AcD3K*>TR<C13>-COOH | Ion score | 59.97 |
| Protein | P25705 | Confidence level | 99.00% |
| Precursor m/z | 571.35264 | Identity Threshold | 44.59 |
| Charge | 2+ | Mass error | -1.29E-4 |



Info

Modified Sequence AcD3-GLAVTPTPVVGSQM<Mox*>TR<C13>-COOH

Ion score 54.67

Protein P26368

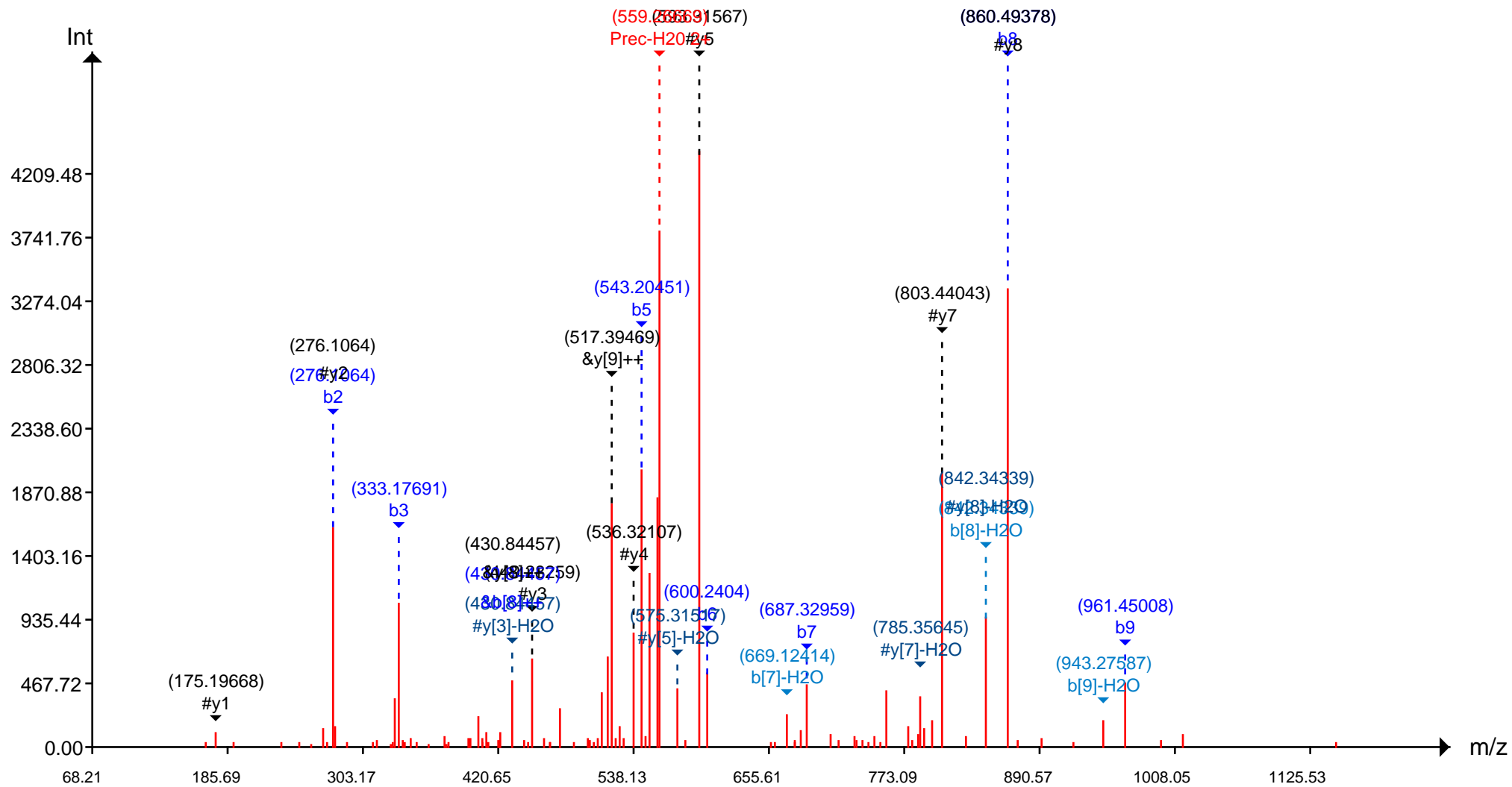
Confidence level 99.00%

Precursor m/z 626.34933

Identity Threshold 48.07

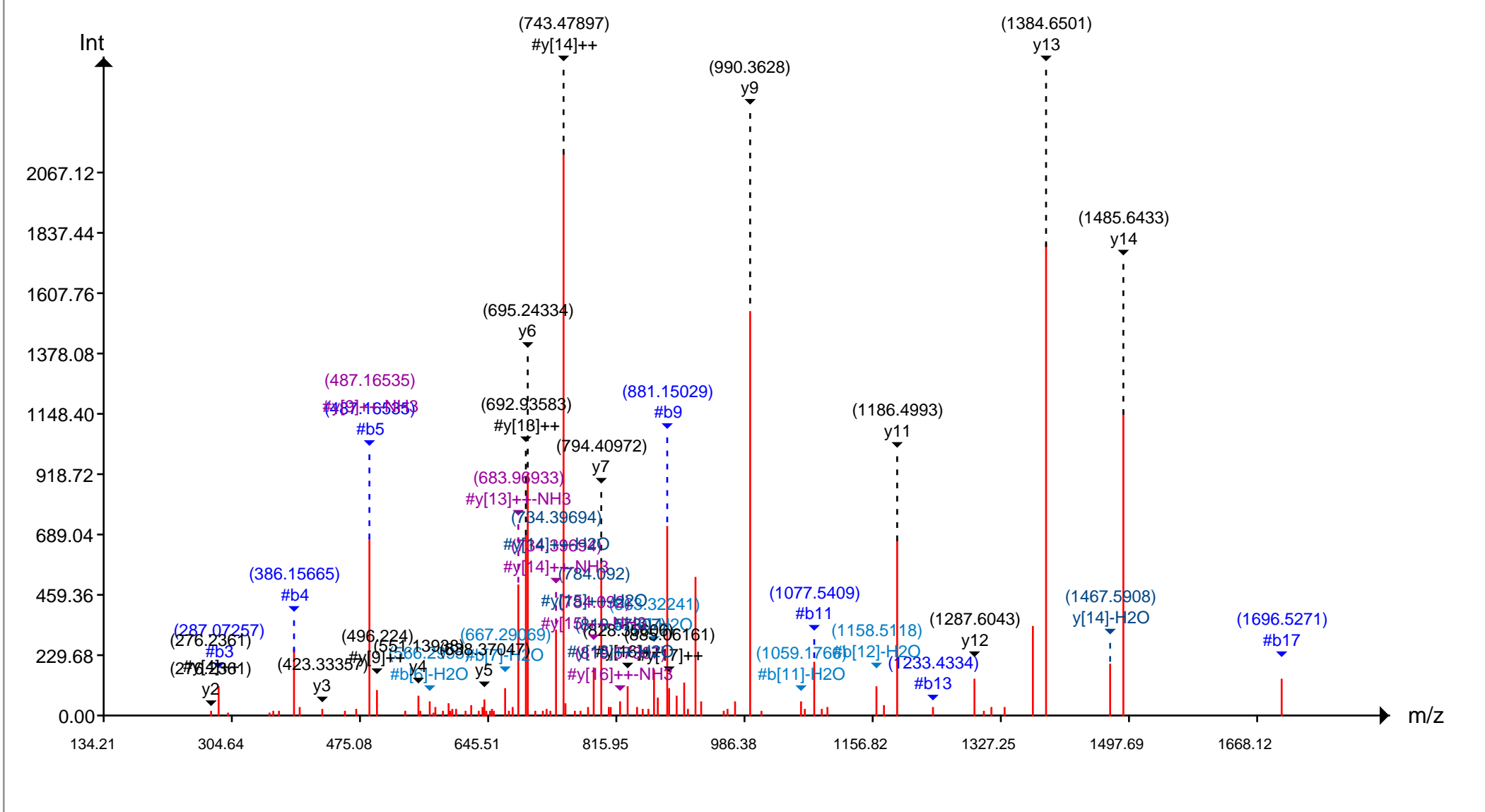
Charge 3+

Mass error -1.75E-4

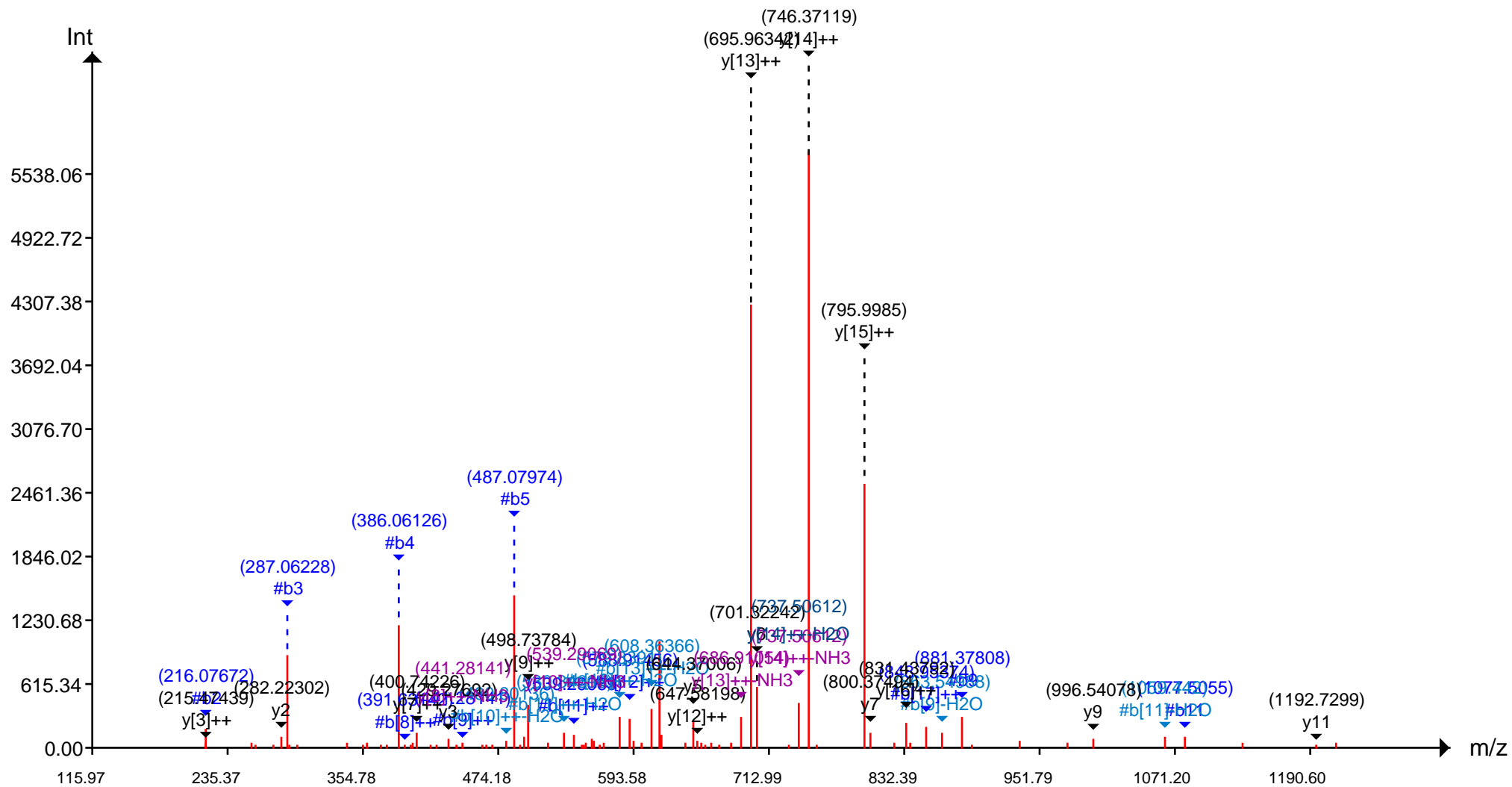


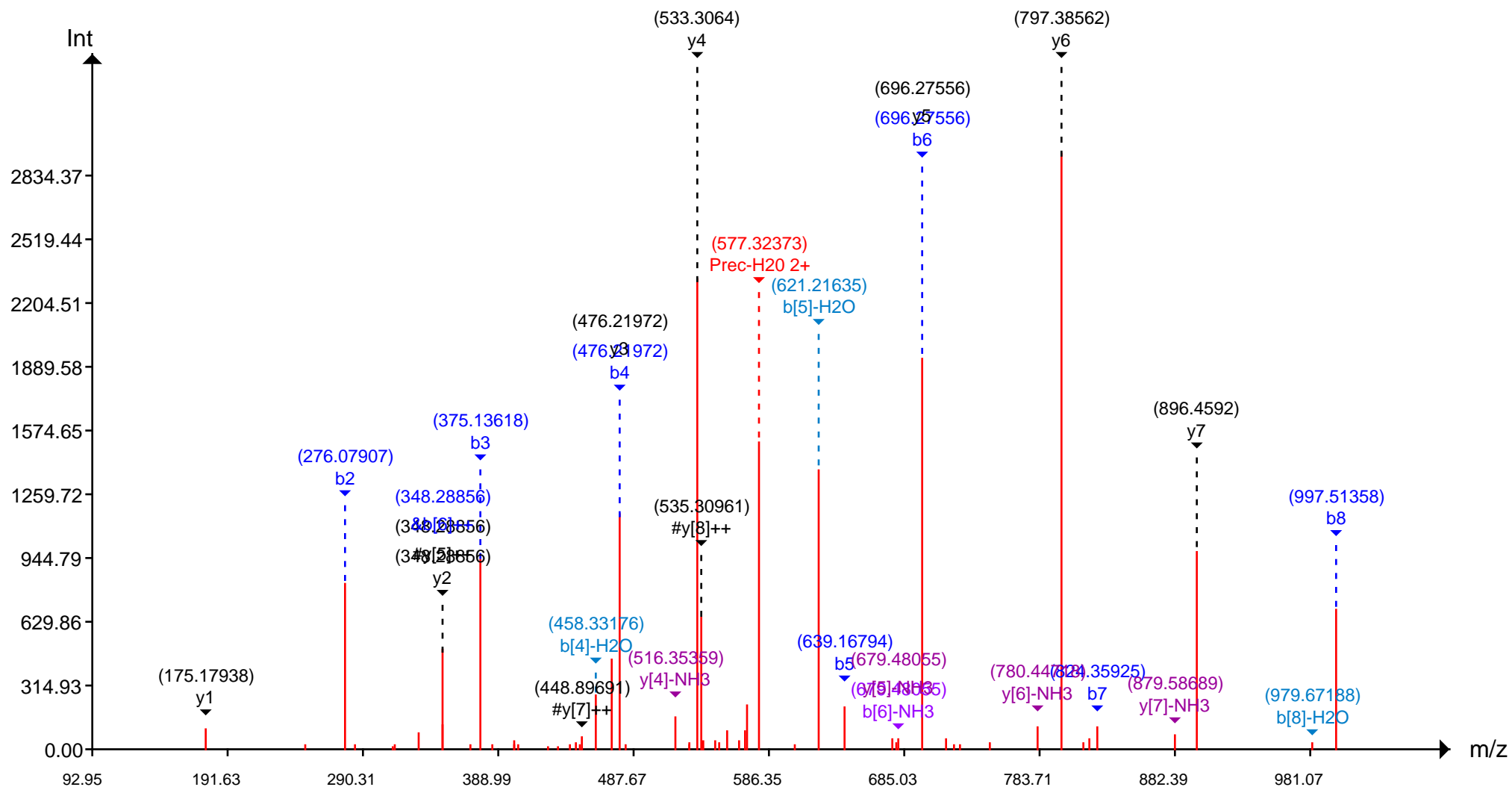
Info

| | | | |
|--------------------------|--------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-GK<AcD3K*>GPIGSK<AcD3K*>TR-COOH | Ion score | 57.69 |
| Protein | P25705 | Confidence level | 99.00% |
| Precursor m/z | 568.34253 | Identity Threshold | 45.68 |
| Charge | 2+ | Mass error | -2.07E-4 |



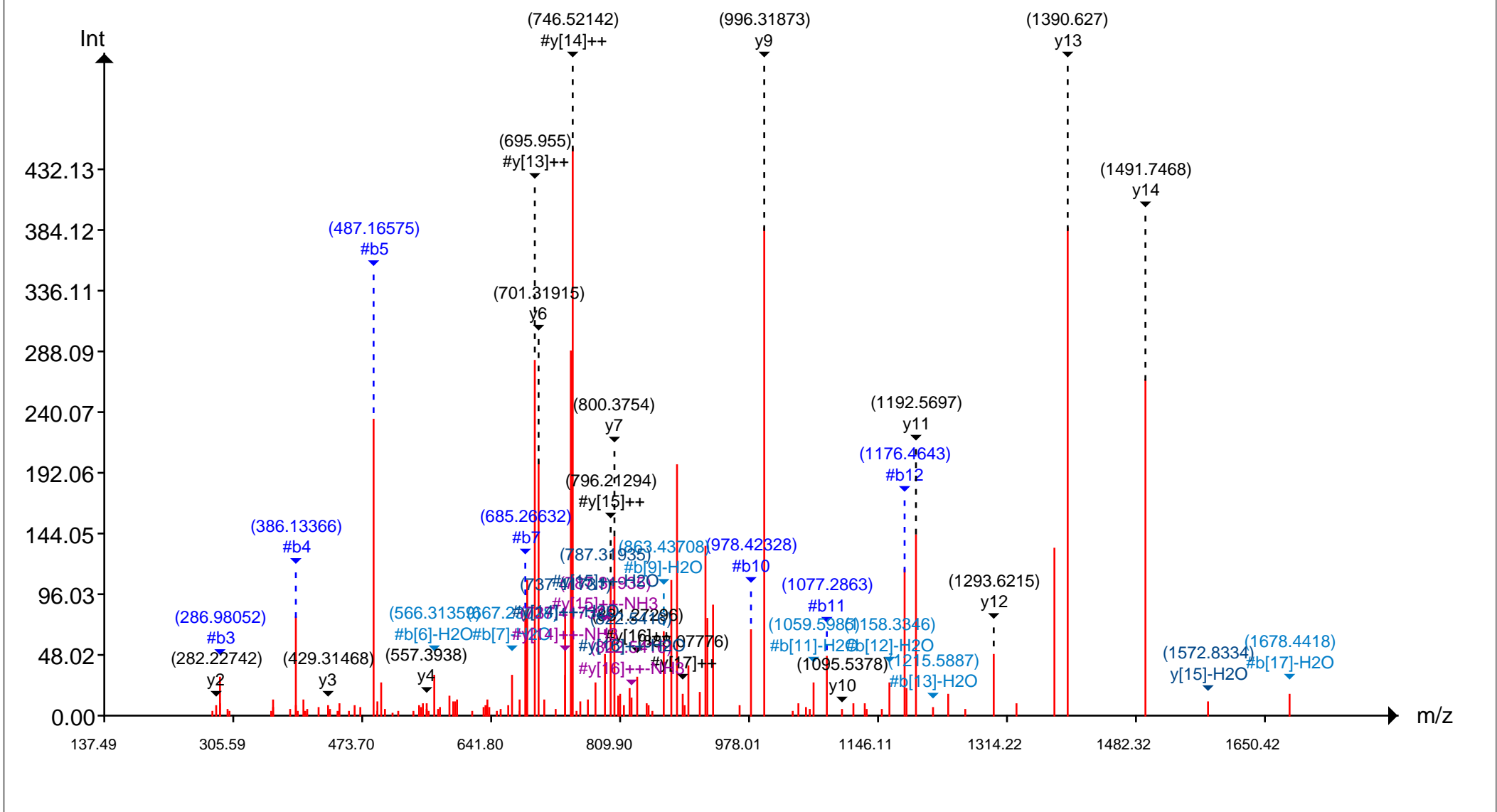
| Info | | | |
|--------------------------|------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-GLAVTPTPVPVVGSMQ<Mox*>TR-COOH | Ion score | 85.95 |
| Protein | P26368 | Confidence level | 99.00% |
| Precursor m/z | 936.01089 | Identity Threshold | 48.49 |
| Charge | 2+ | Mass error | 0.001033 |





Info

| | | | |
|--------------------------|-------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-GK<AcD3K*>VTYGQK<AcD3K*>R-COOH | Ion score | 64.07 |
| Protein | Q92499 | Confidence level | 99.00% |
| Precursor m/z | 586.34258 | Identity Threshold | 47.48 |
| Charge | 2+ | Mass error | -1.07E-4 |



Info

Modified Sequence AcD3-GLAVTPTPVVGSQM<Mox*>TR<C13>-COOH

Ion score 60.33

Protein P26368

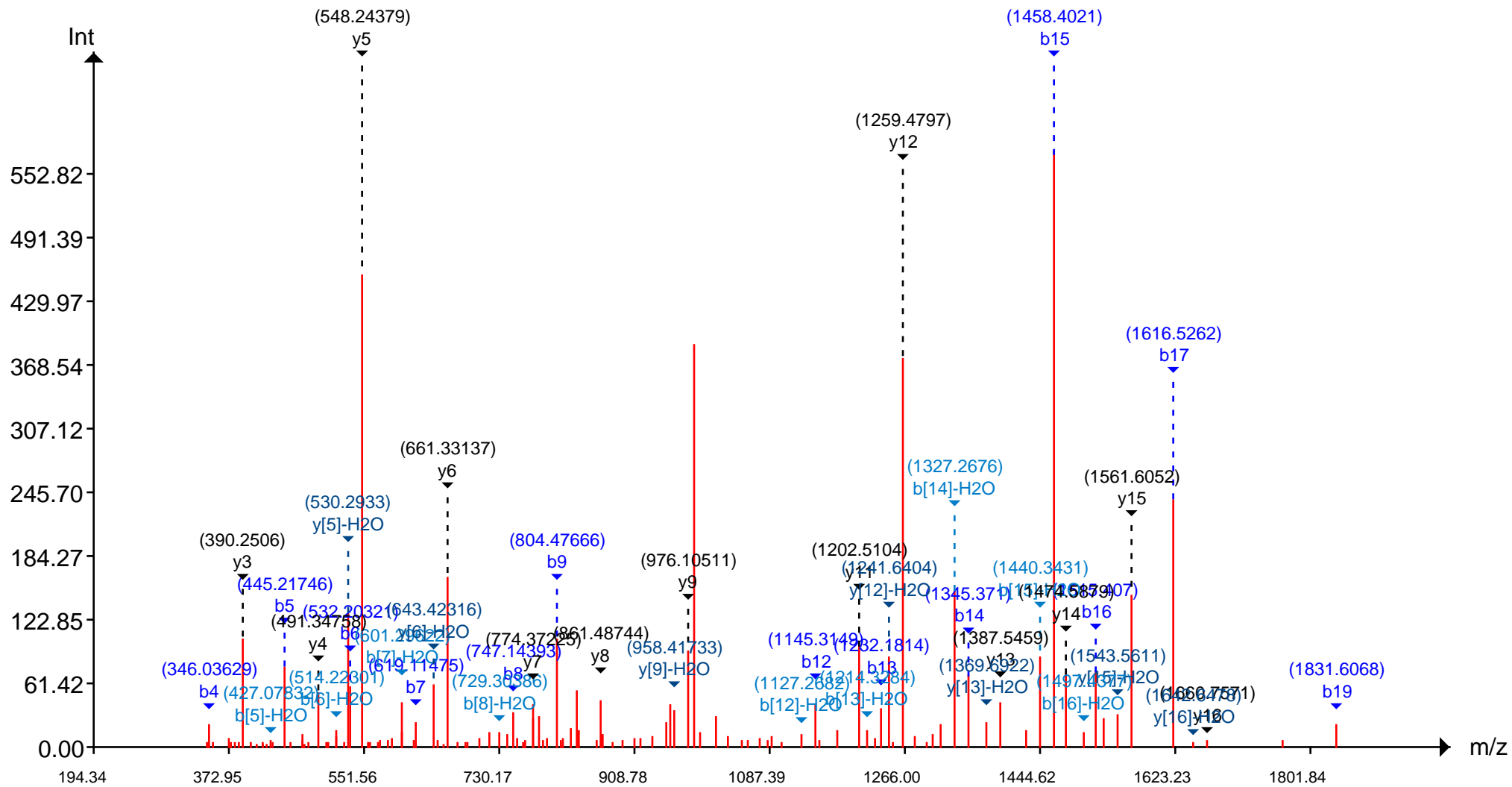
Confidence level 99.00%

Precursor m/z 939.02085

Identity Threshold 47.92

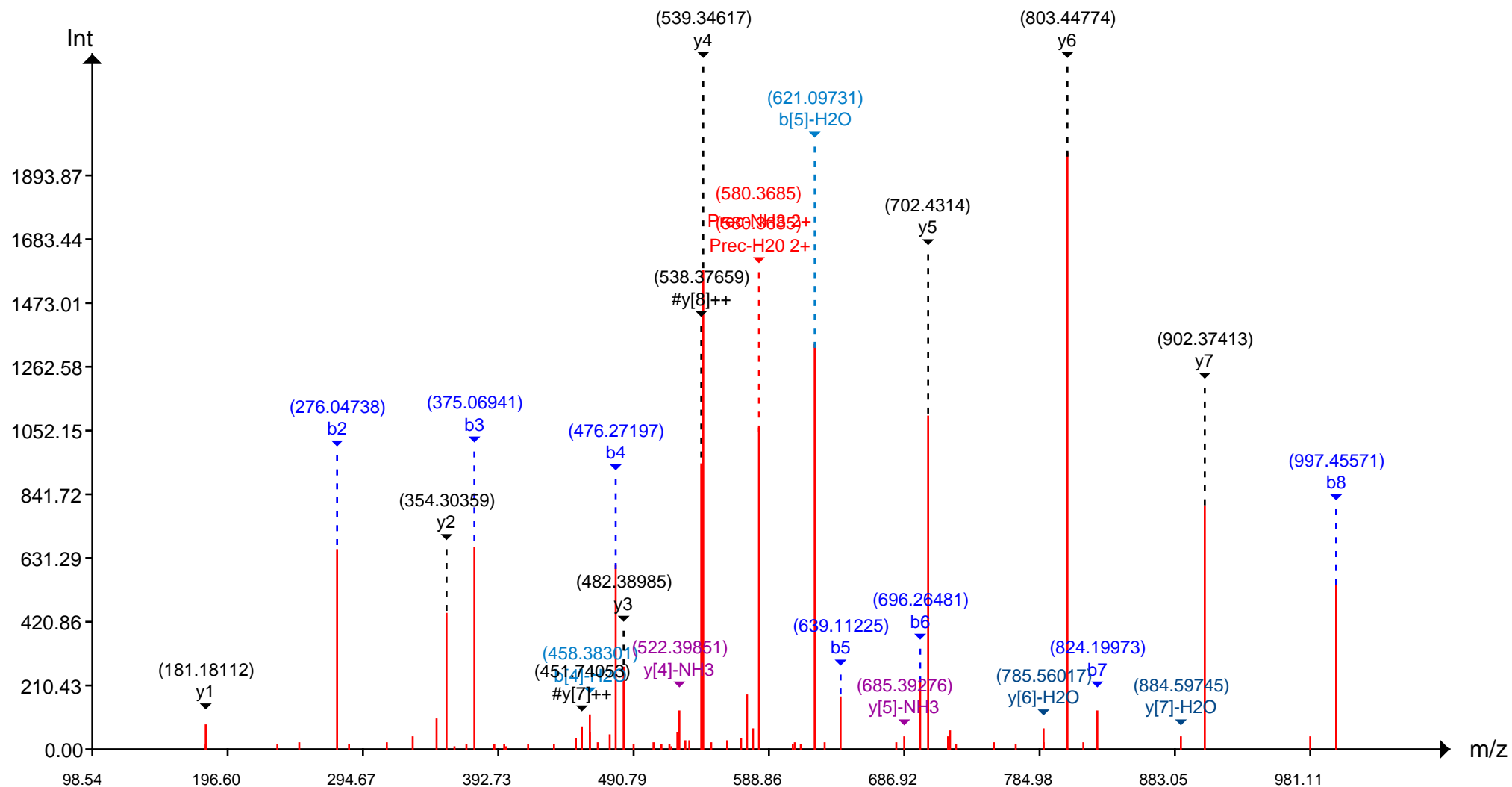
Charge 2+

Mass error 8.11E-4



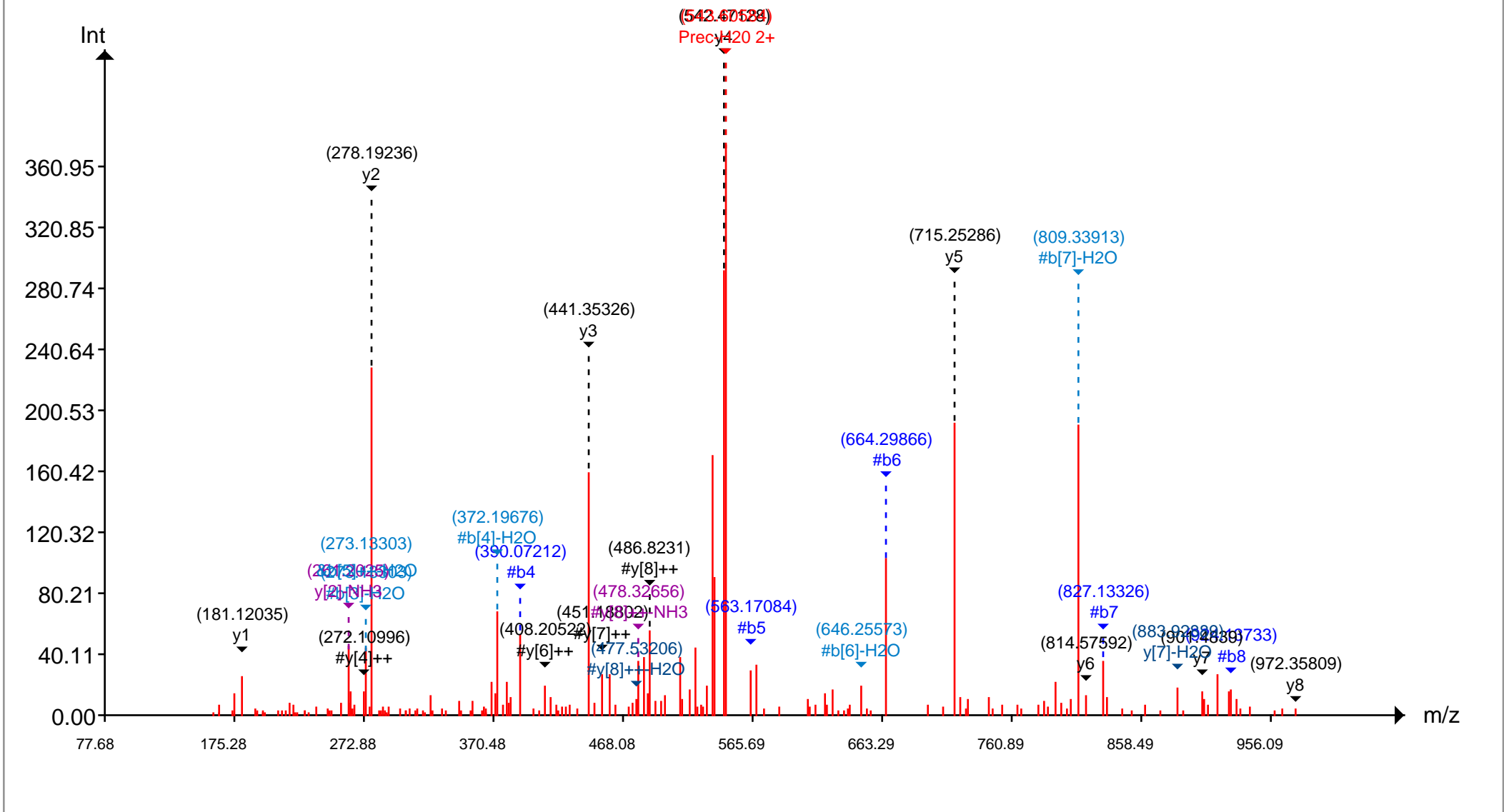
Info

| | | | |
|--------------------------|--------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-GGGEVSSQGPEDSLLGTQSR-COOH | Ion score | 106.96 |
| Protein | Q9P107 | Confidence level | 99.00% |
| Precursor m/z | 1003.4792 | Identity Threshold | 47.43 |
| Charge | 2+ | Mass error | 0.001068 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GK<AcD3K*>VTYGQK<AcD3K*>R<C13>-COOH | Ion score | 61.34 |
| Protein | Q92499 | Confidence level | 99.00% |
| Precursor m/z | 589.35262 | Identity Threshold | 46.55 |
| Charge | 2+ | Mass error | -1.69E-4 |



Info

Modified Sequence AcD3-SASVK<AcD3K*>TYPR<C13>-COOH

Ion score 54.13

Protein Q8N9T8

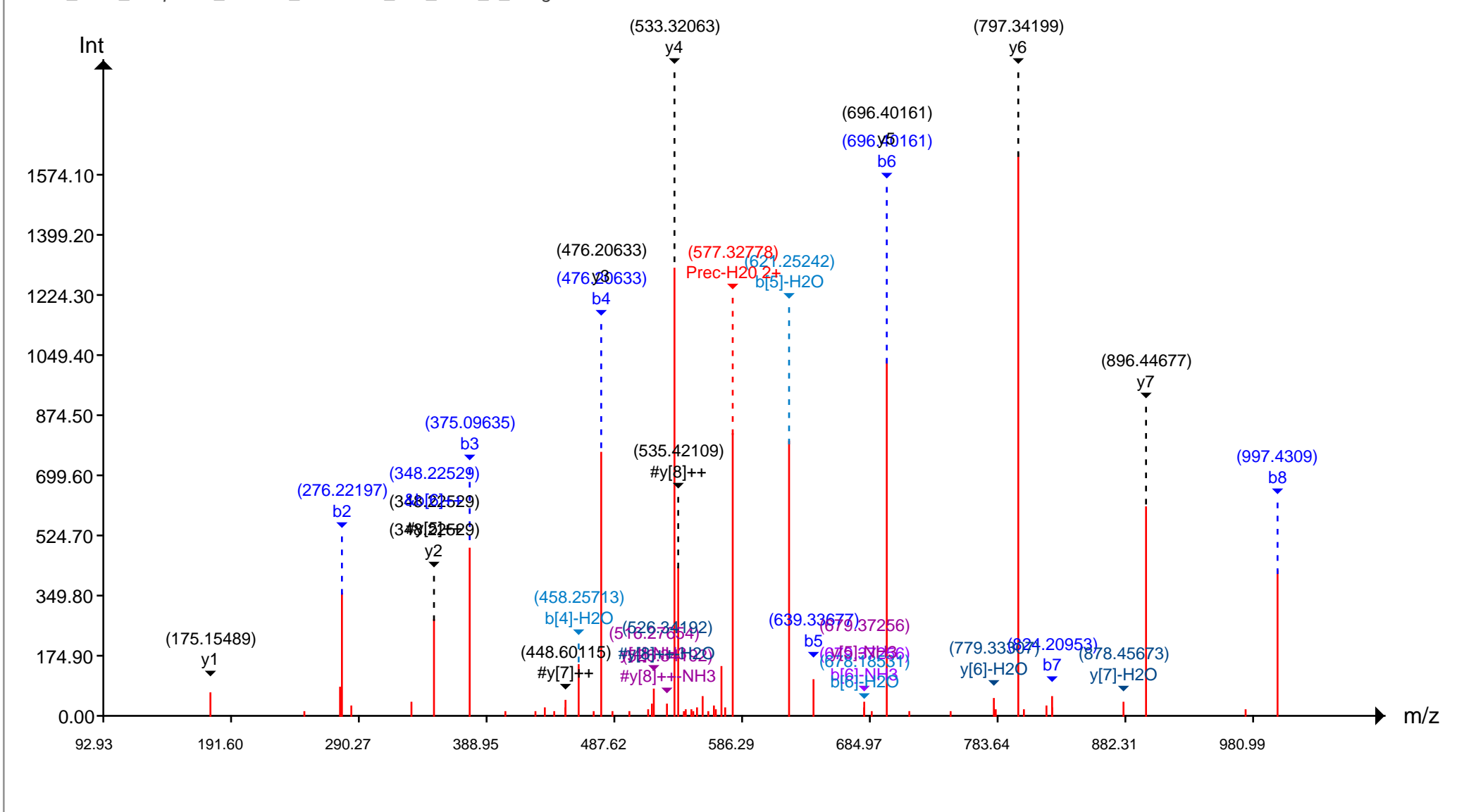
Confidence level 99.00%

Precursor m/z 552.81663

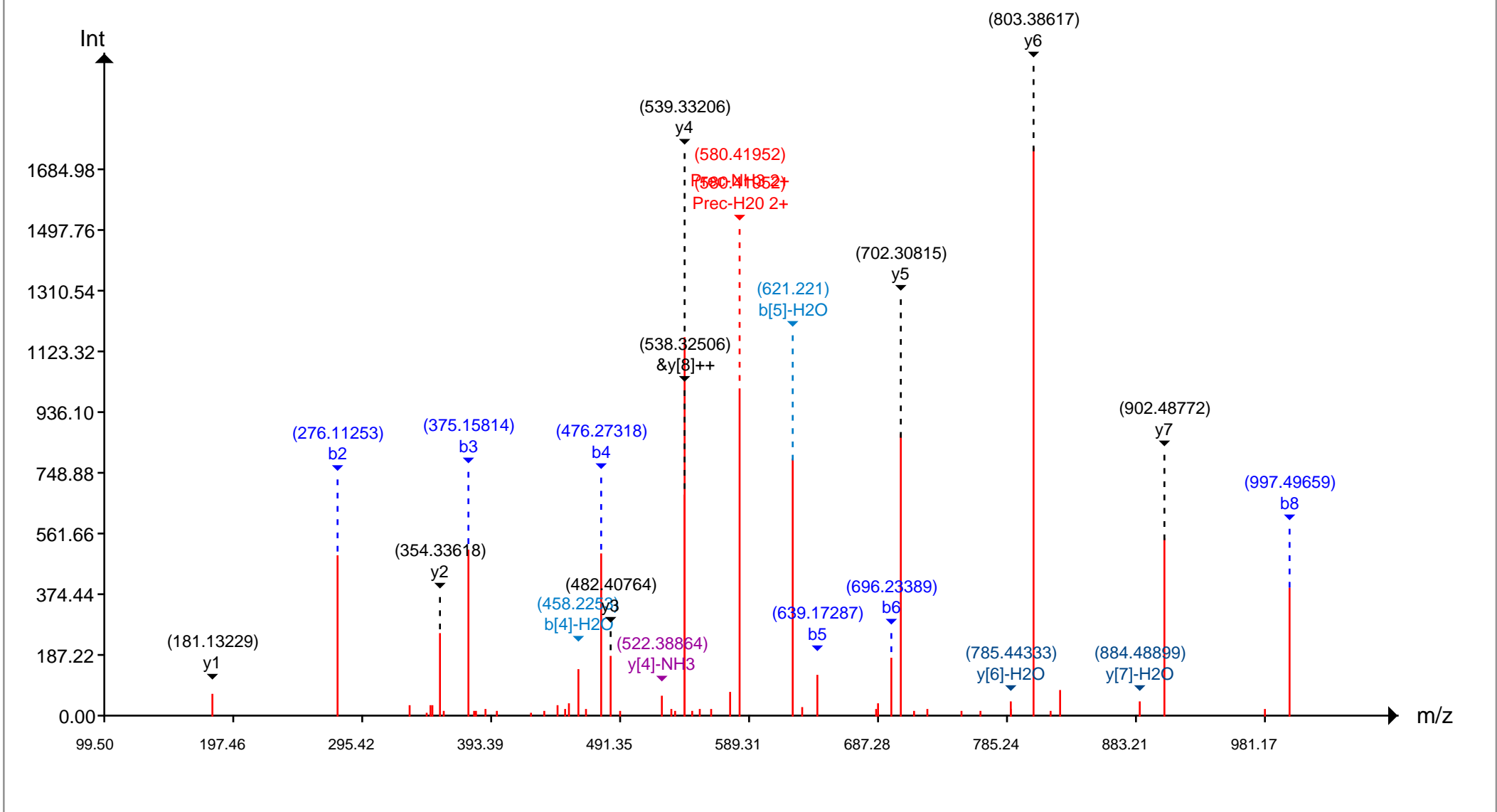
Identity Threshold 48.15

Charge 2+

Mass error -2.04E-4



| Info | | | |
|--------------------------|-------------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-GK<AcD3K*>VTYGQK<AcD3K*>R-COOH | Ion score | 64.18 |
| Protein | Q92499 | Confidence level | 99.00% |
| Precursor m/z | 586.34281 | Identity Threshold | 47.32 |
| Charge | 2+ | Mass error | 3.53E-4 |



Info

Modified Sequence AcD3-GK<AcD3K*>VTYGQK<AcD3K*>R<C13>-COOH

Ion score 62.56

Protein Q92499

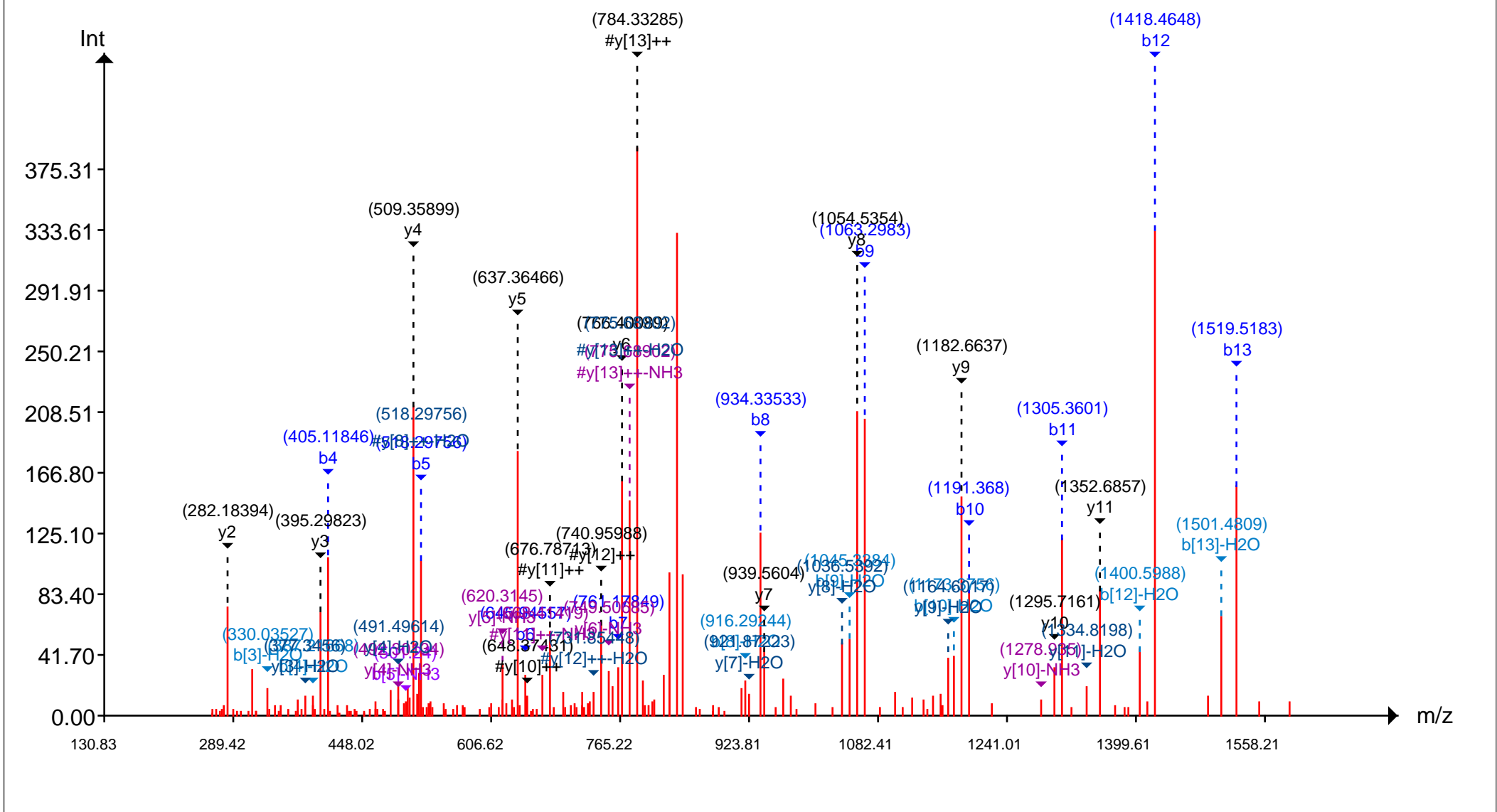
Confidence level 99.00%

Precursor m/z 589.35279

Identity Threshold 46.50

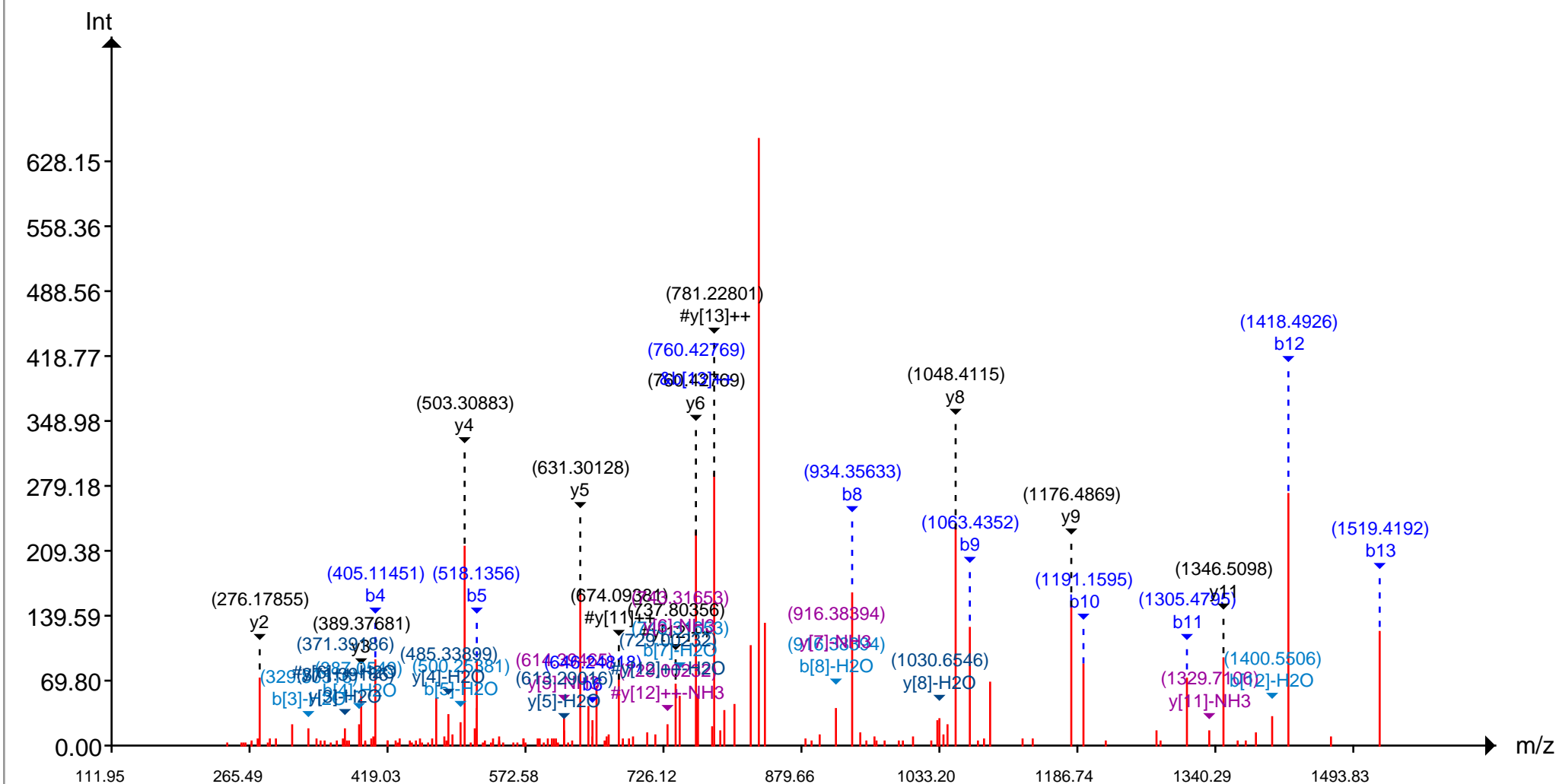
Charge 2+

Mass error 1.71E-4



Info

| | | | |
|--------------------------|---------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-SSQGLQDK<AcD3K*>EQNLTR<C13>-COOH | Ion score | 90.2 |
| Protein | O60293 | Confidence level | 99.00% |
| Precursor m/z | 850.44468 | Identity Threshold | 48.96 |
| Charge | 2+ | Mass error | -2.68E-4 |



Info

Modified Sequence AcD3-SSQGLQDK<AcD3K*>EQNLTR-COOH

Ion score 77.96

Protein O60293

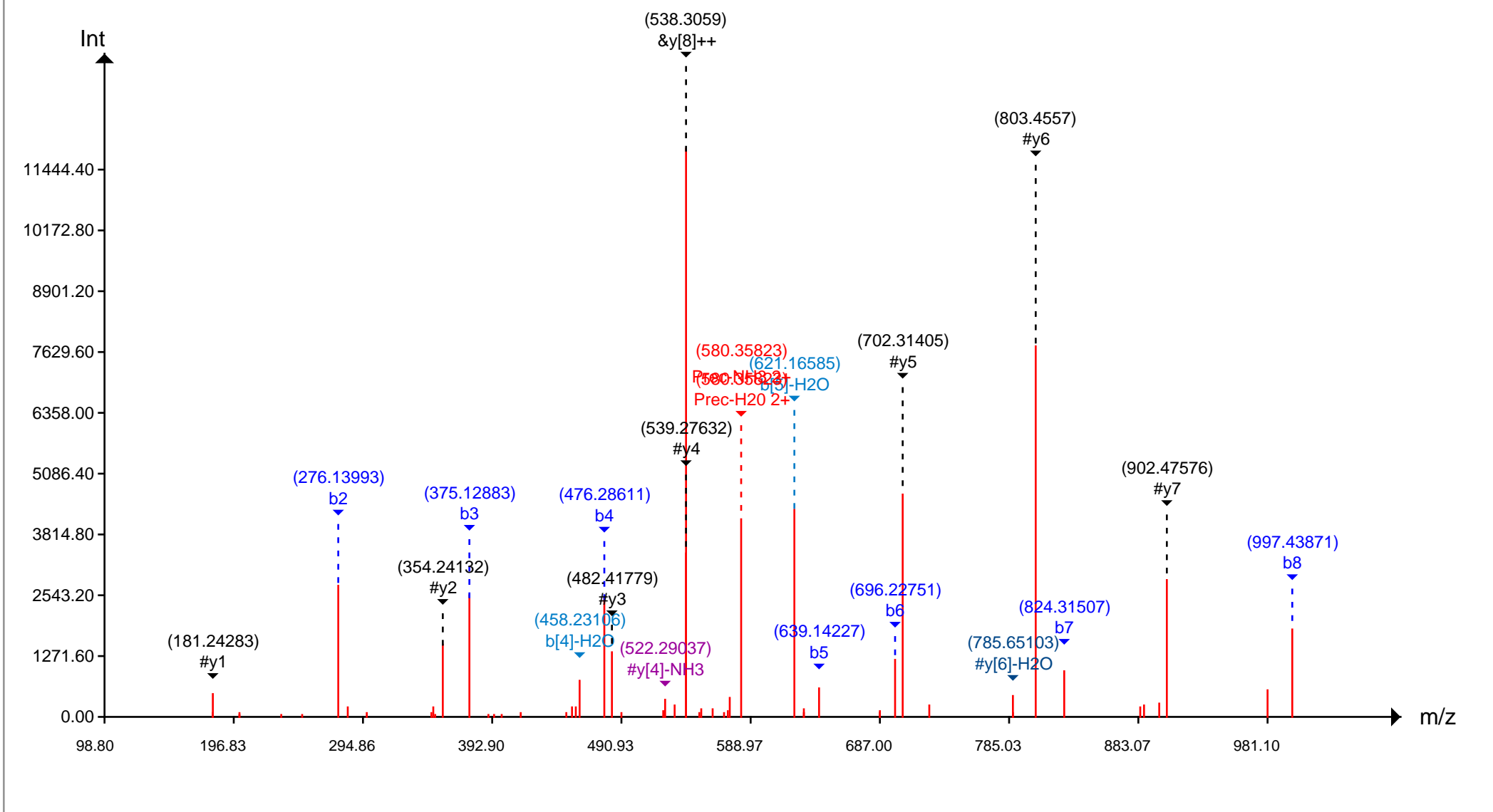
Confidence level 99.00%

Precursor m/z 847.43536

Identity Threshold 49.10

Charge 2+

Mass error 0.001233



Info

Modified Sequence AcD3-GK<AcD3K*>VTYGQK<AcD3K*>R<C13>-COOH

Ion score 64.74

Protein Q92499

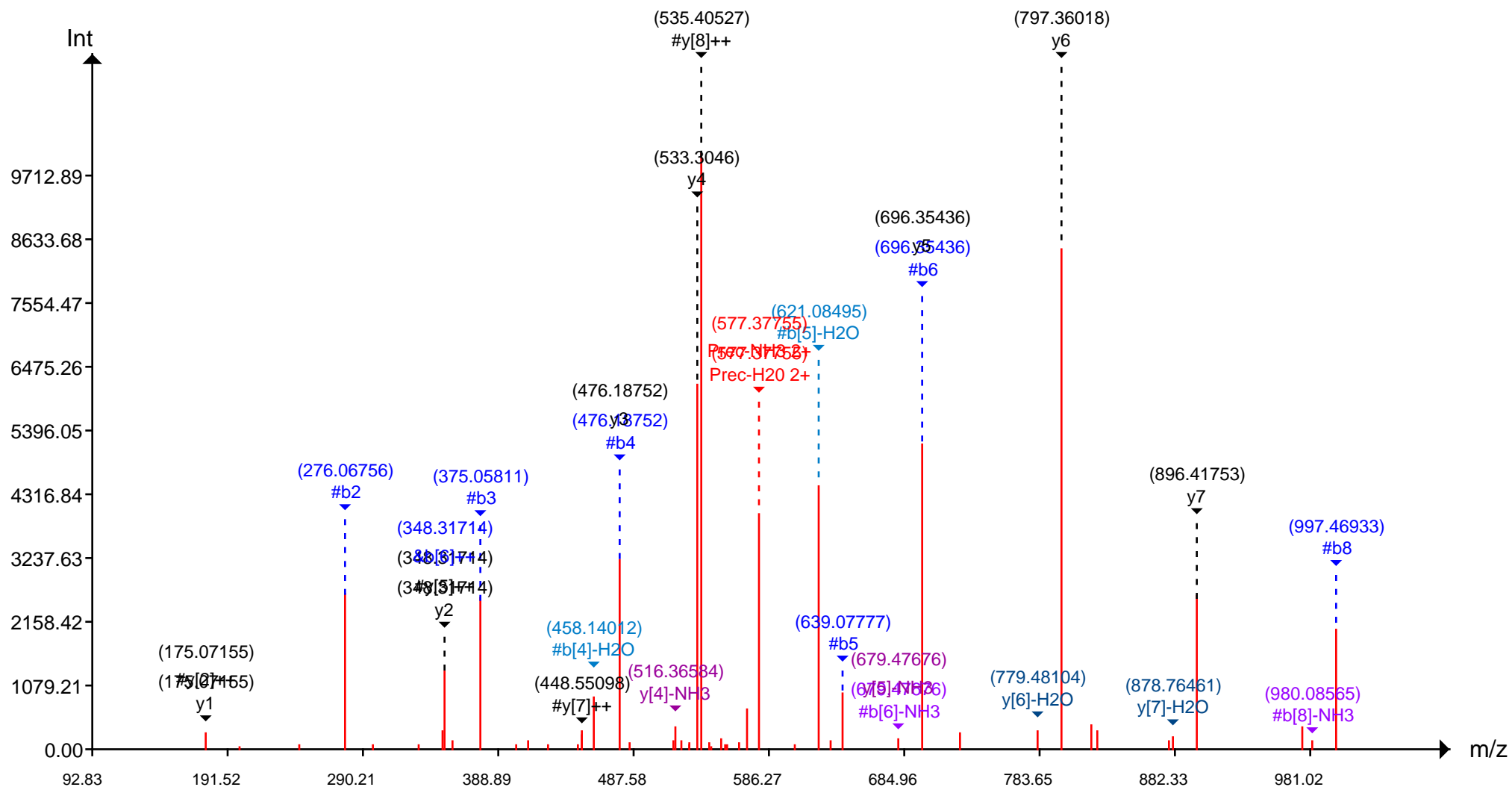
Confidence level 99.00%

Precursor m/z 589.35298

Identity Threshold 46.46

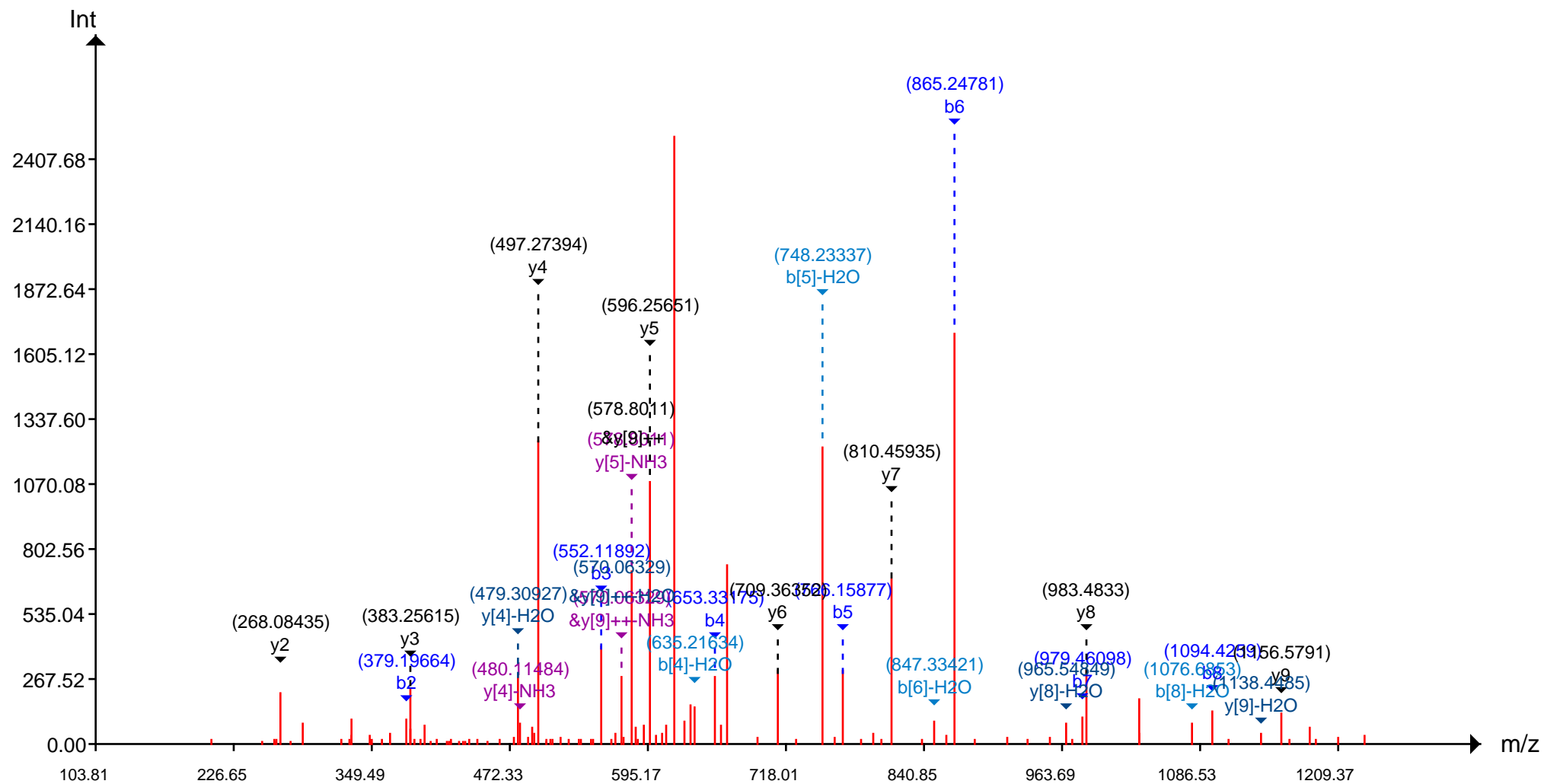
Charge 2+

Mass error 5.51E-4



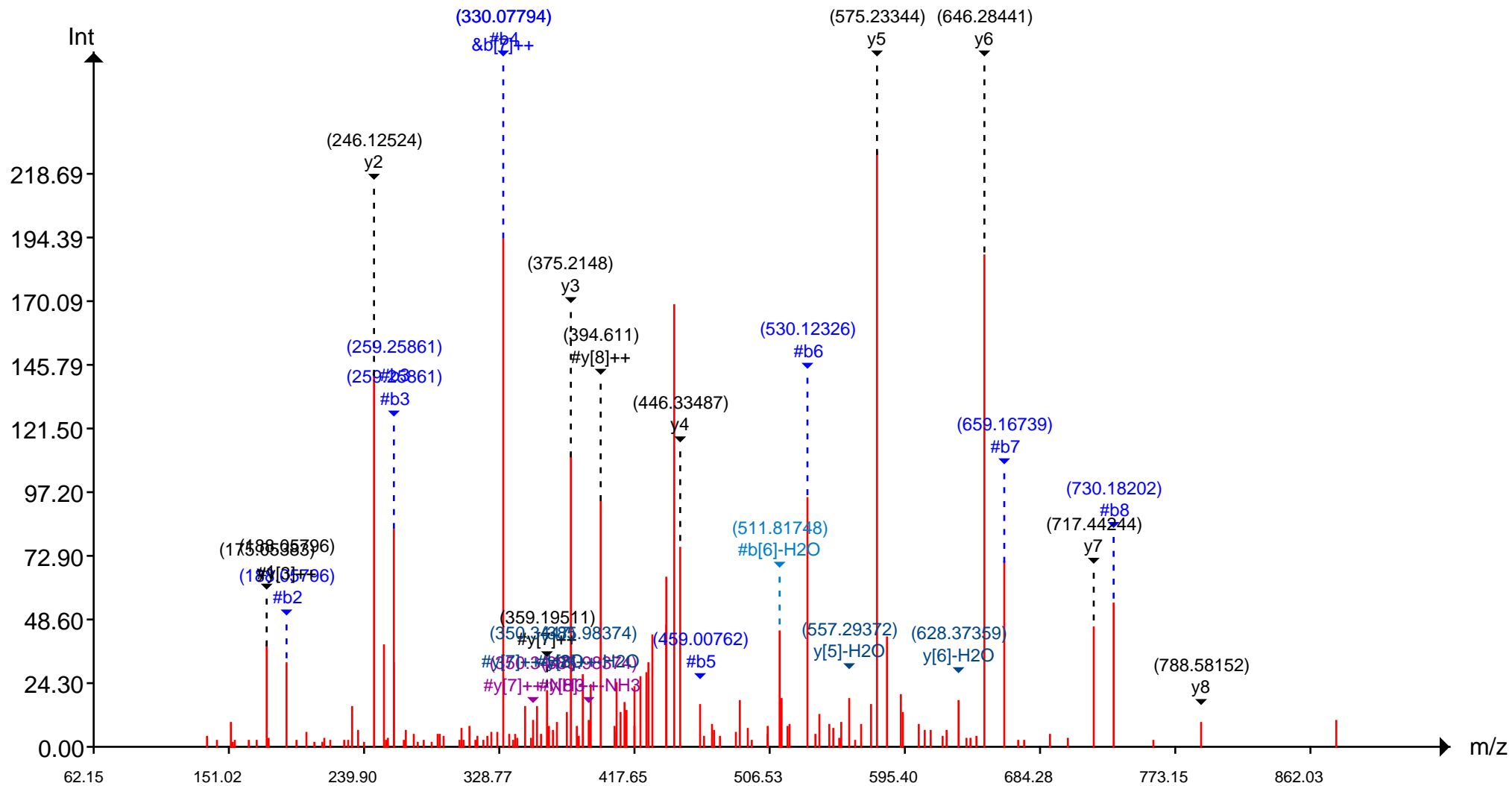
Info

| | | | |
|--------------------------|-------------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-GK<AcD3K*>VTYGQK<AcD3K*>R-COOH | Ion score | 56.62 |
| Protein | Q92499 | Confidence level | 99.00% |
| Precursor m/z | 586.34295 | Identity Threshold | 47.28 |
| Charge | 2+ | Mass error | 6.33E-4 |



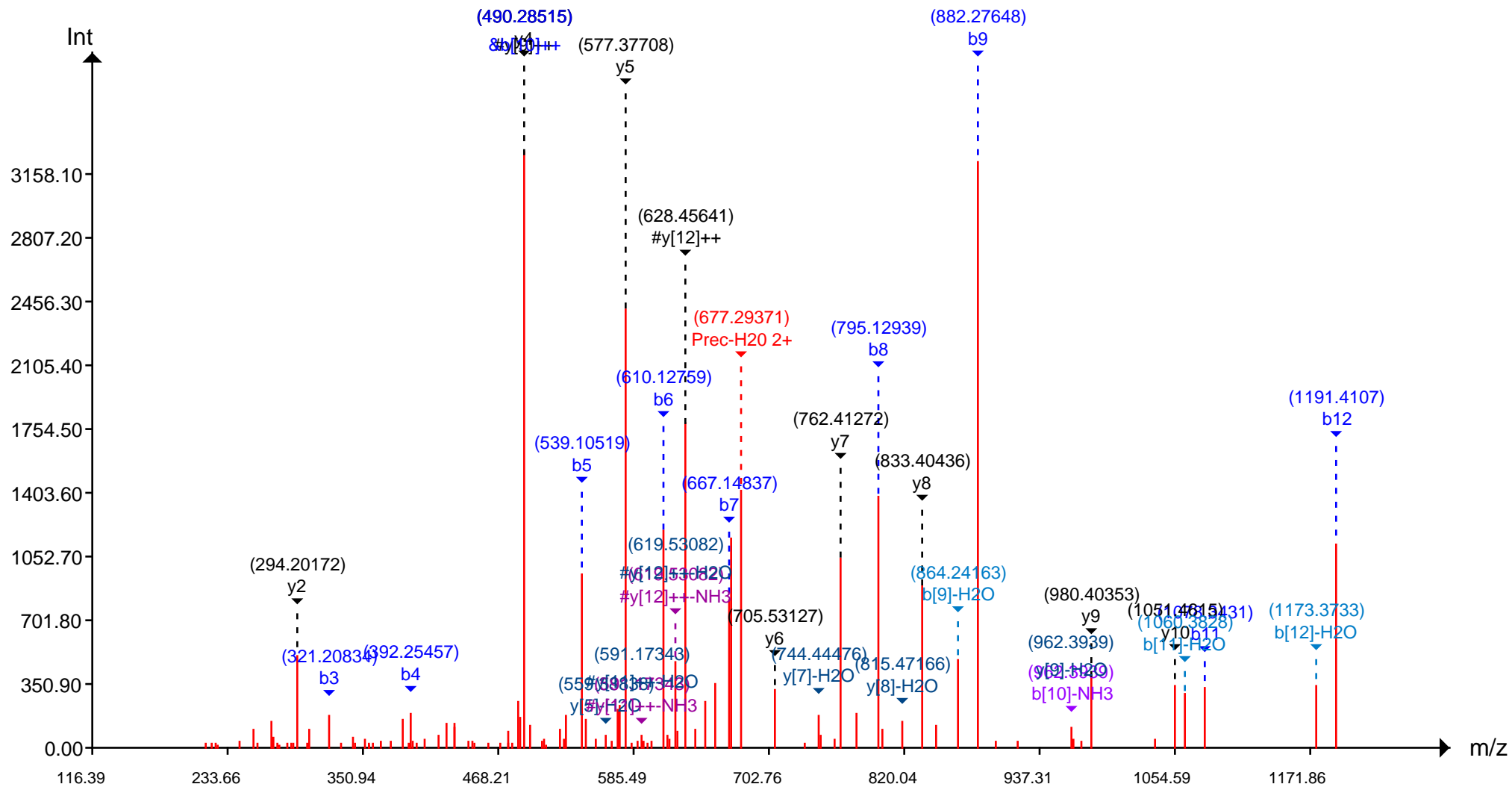
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-C<Cmm*>K<AcD3K*>K<AcD3K*>TIVNDSR<C13>-COOH | Ion score | 69.49 |
| Protein | Q00987 | Confidence level | 99.00% |
| Precursor m/z | 681.37896 | Identity Threshold | 48.55 |
| Charge | 2+ | Mass error | 7.38E-4 |



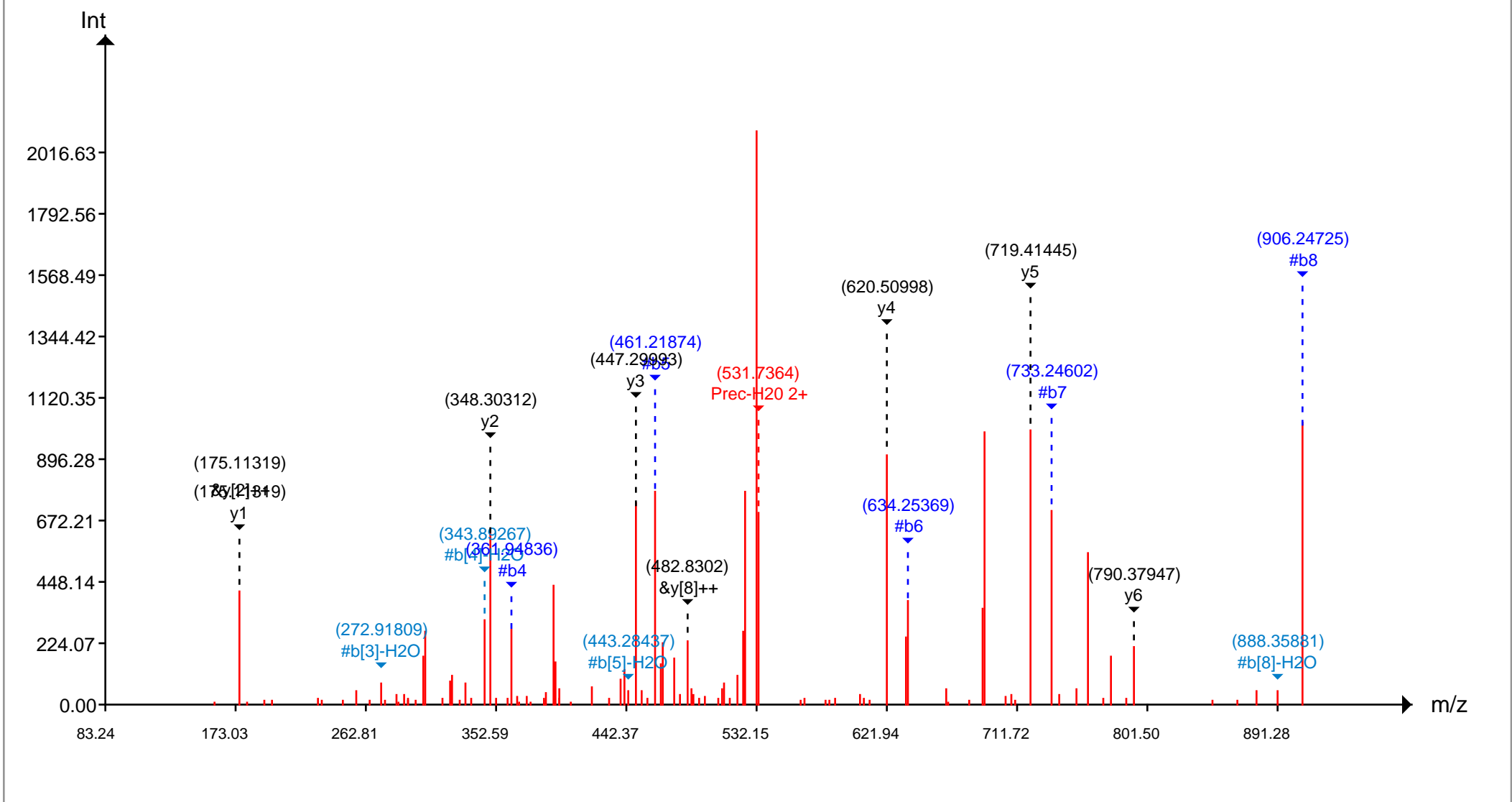
Info

| | | | |
|--------------------------|---------------------|---------------------------|---------|
| Modified Sequence | AcD3-AAAAEAEAR-COOH | Ion score | 66.83 |
| Protein | Q9Y5V3 | Confidence level | 99.00% |
| Precursor m/z | 452.73185 | Identity Threshold | 48.43 |
| Charge | 2+ | Mass error | 2.65E-4 |



Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-AGM<Mox*>AM<Mox*>AGQSPVLR<C13>-COOH | Ion score | 68.04 |
| Protein | P26599 | Confidence level | 99.00% |
| Precursor m/z | 686.34845 | Identity Threshold | 48.74 |
| Charge | 2+ | Mass error | 2.34E-4 |



Info

Modified Sequence AcD3-ASSAVK<AcD3K*>VK<AcD3K*>R-COOH

Ion score 44.87

Protein Q13148

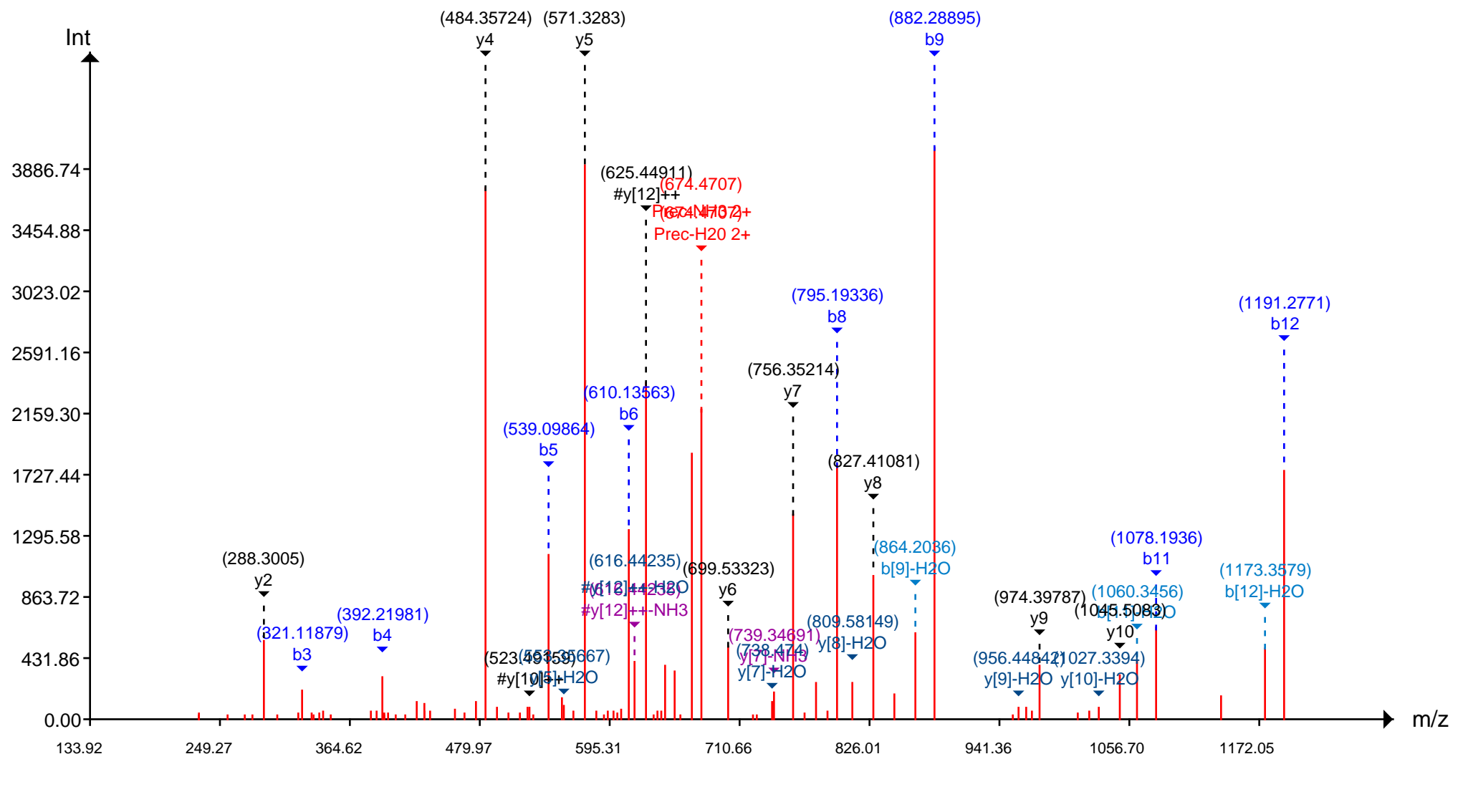
Confidence level 99.00%

Precursor m/z 540.83992

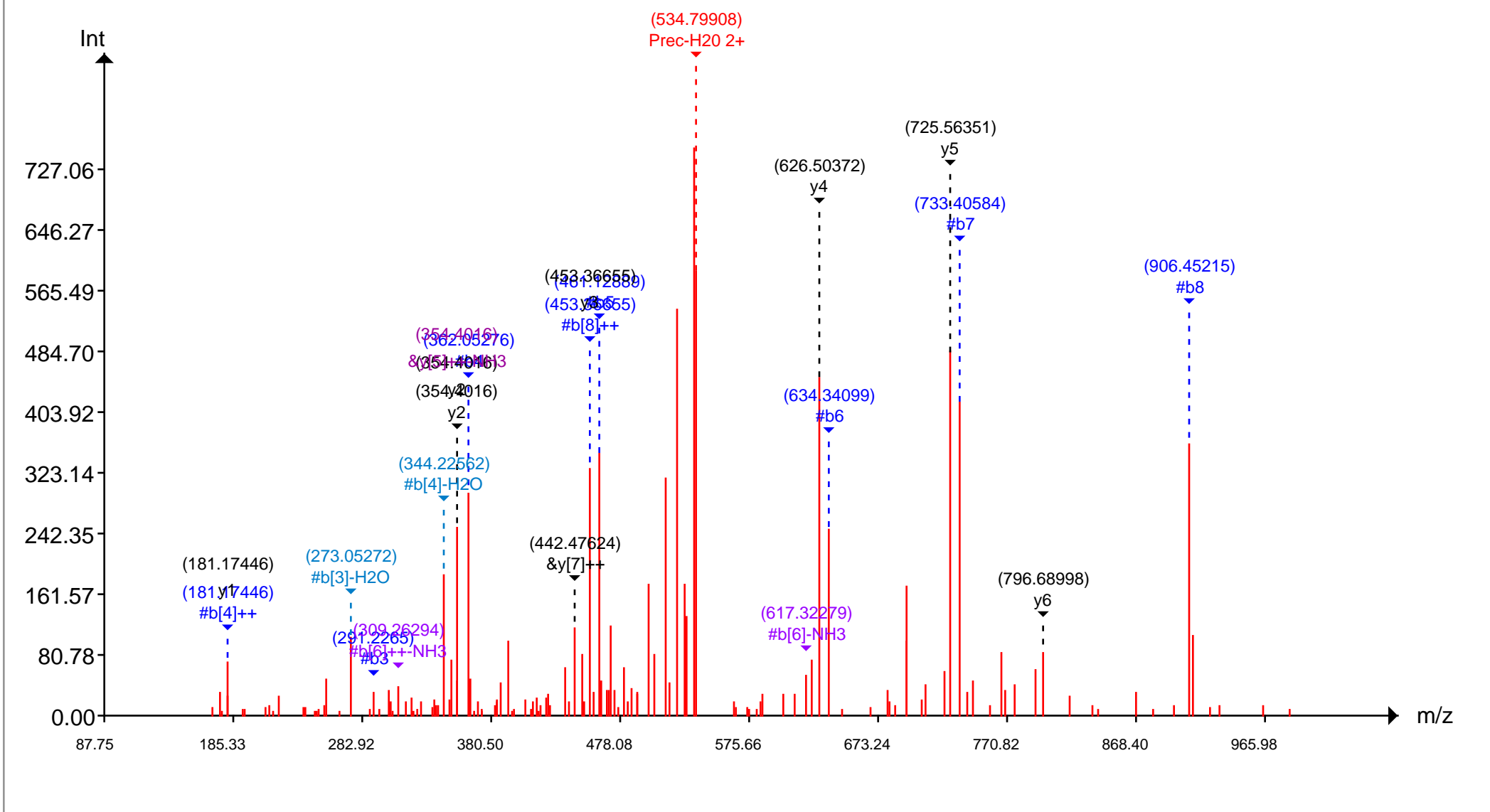
Identity Threshold 43.82

Charge 2+

Mass error 4.02E-4



| Info | | | |
|--------------------------|-------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-AGM<Mox*>AM<Mox*>AGQSPVLR-COOH | Ion score | 66.77 |
| Protein | P26599 | Confidence level | 99.00% |
| Precursor m/z | 683.33821 | Identity Threshold | 48.43 |
| Charge | 2+ | Mass error | -1.04E-4 |



Info

Modified Sequence AcD3-ASSAVK<AcD3K*>VK<AcD3K*>R<C13>-COOH

Ion score 45.94

Protein Q13148

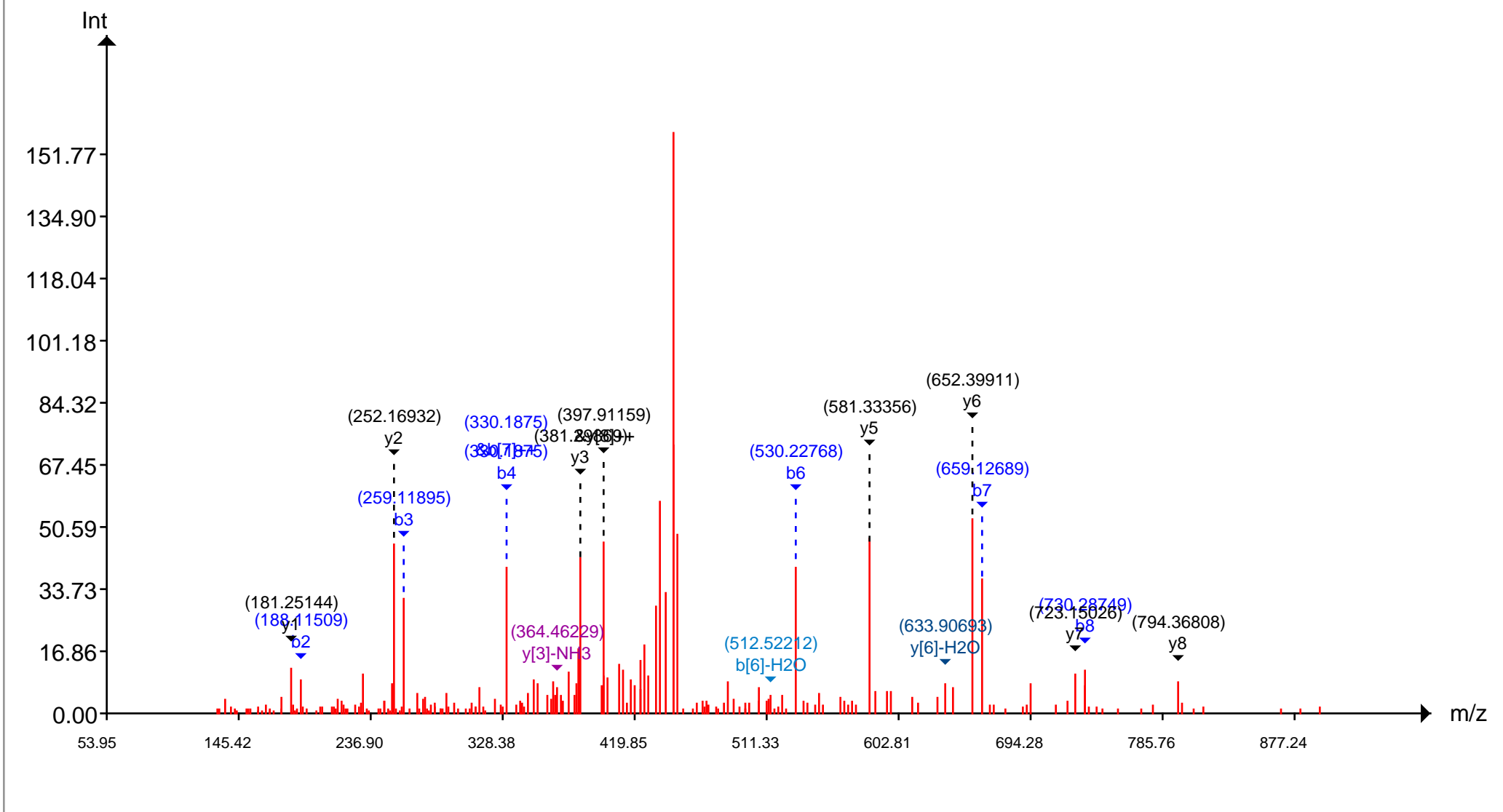
Confidence level 99.00%

Precursor m/z 543.84999

Identity Threshold 42.35

Charge 2+

Mass error 4.0E-4



Info

Modified Sequence AcD3-AAAAEAEAR<C13>-COOH

Ion score 54.14

Protein Q9Y5V3

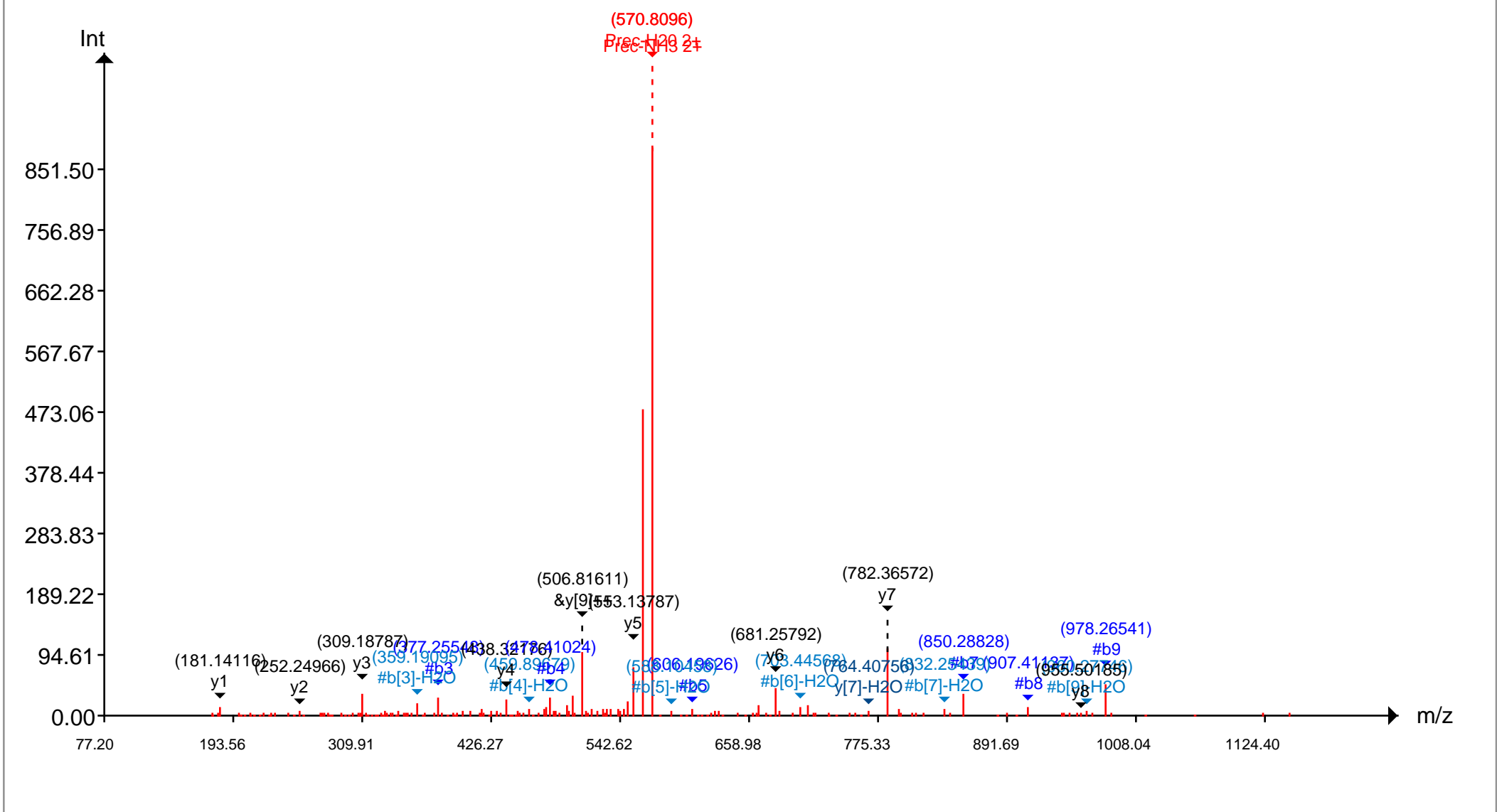
Confidence level 99.00%

Precursor m/z 455.74177

Identity Threshold 48.95

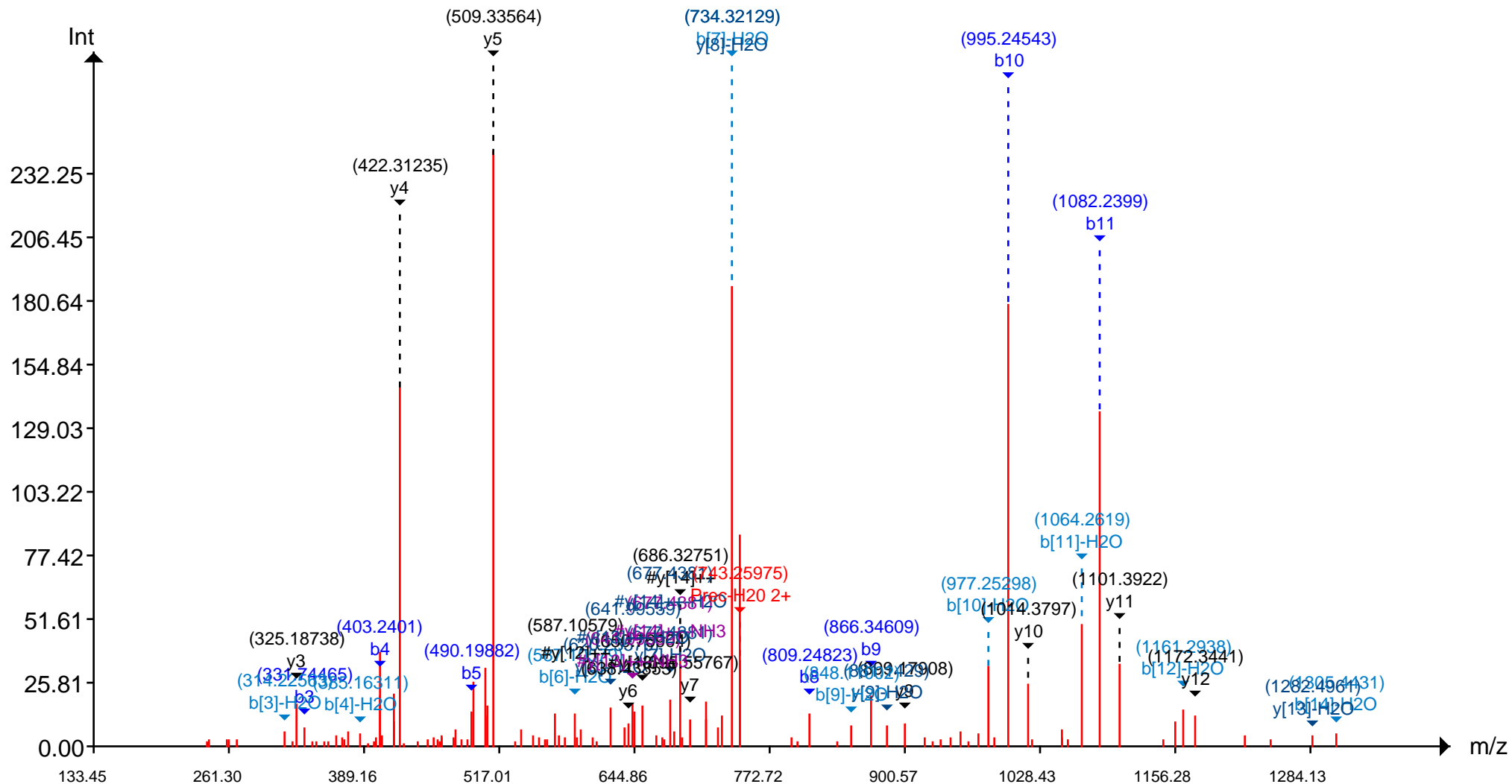
Charge 2+

Mass error -3.7E-5



Info

| | | | |
|--------------------------|-----------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-TGK<AcD3K*>TQDEGAR<C13>-COOH | Ion score | 49.85 |
| Protein | Q9BQC3 | Confidence level | 99.00% |
| Precursor m/z | 579.80161 | Identity Threshold | 48.72 |
| Charge | 2+ | Mass error | -4.13E-4 |



Info

Modified Sequence AcD3-SAQASDM<Mox*>GGESPGSR<C13>-COOH

Ion score 53.99

Protein P50851

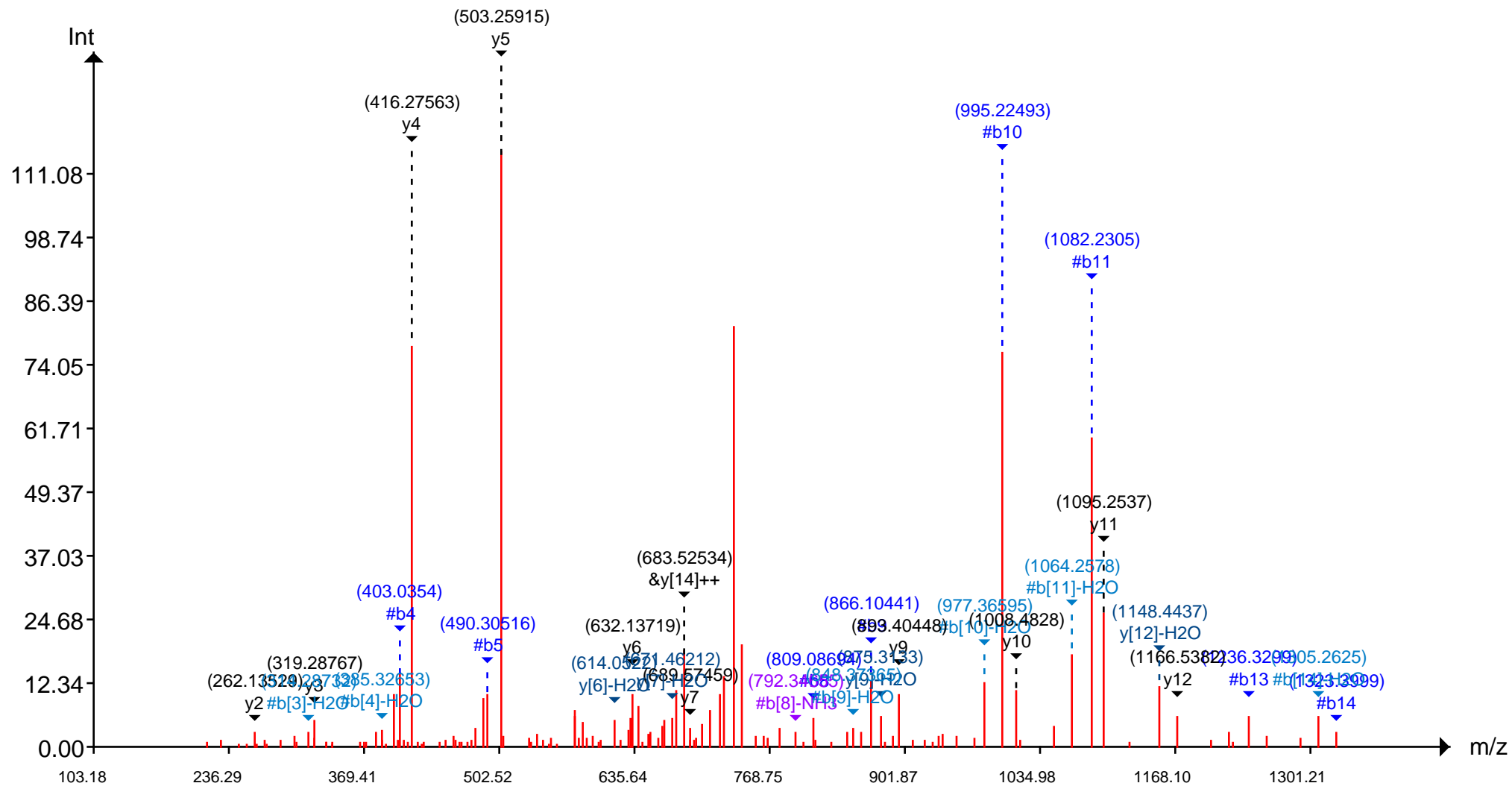
Confidence level 99.00%

Precursor m/z 752.32957

Identity Threshold 43.26

Charge 2+

Mass error 3.77E-4



Info

Modified Sequence AcD3-SAQASDM<Mox*>GGESPGSR-COOH

Ion score 76.06

Protein P50851

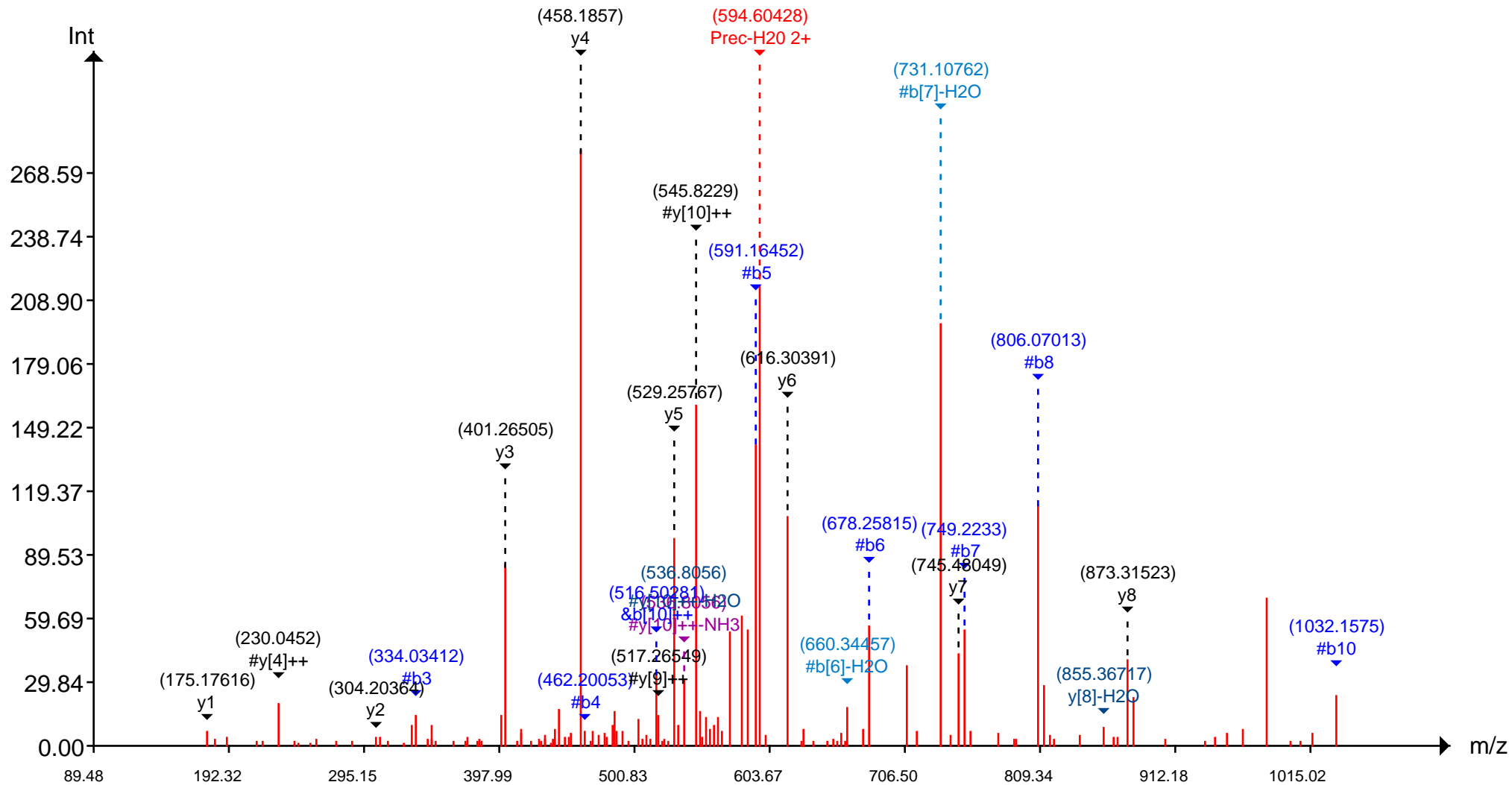
Confidence level 99.00%

Precursor m/z 749.3196

Identity Threshold 42.13

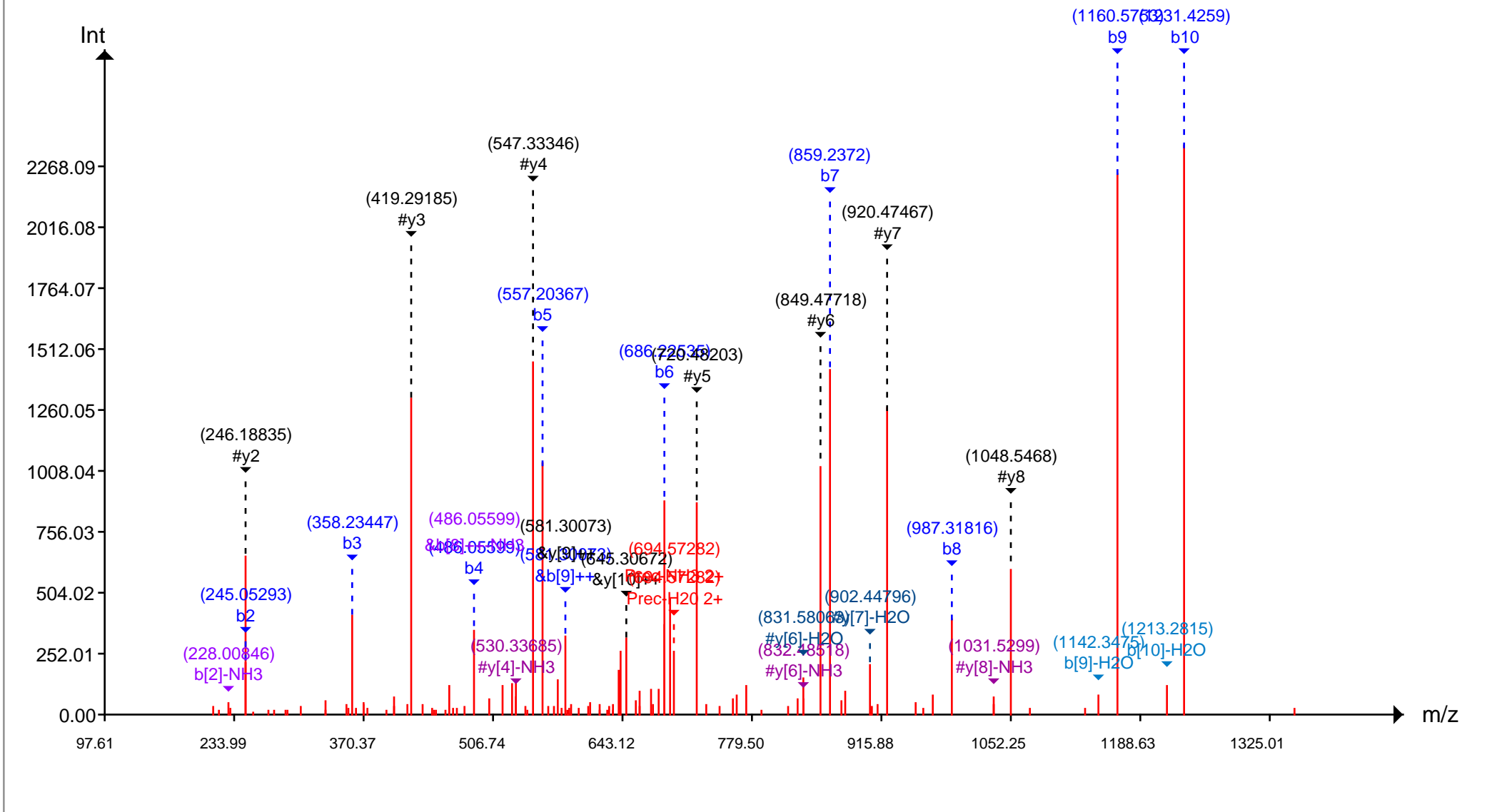
Charge 2+

Mass error 5.79E-4

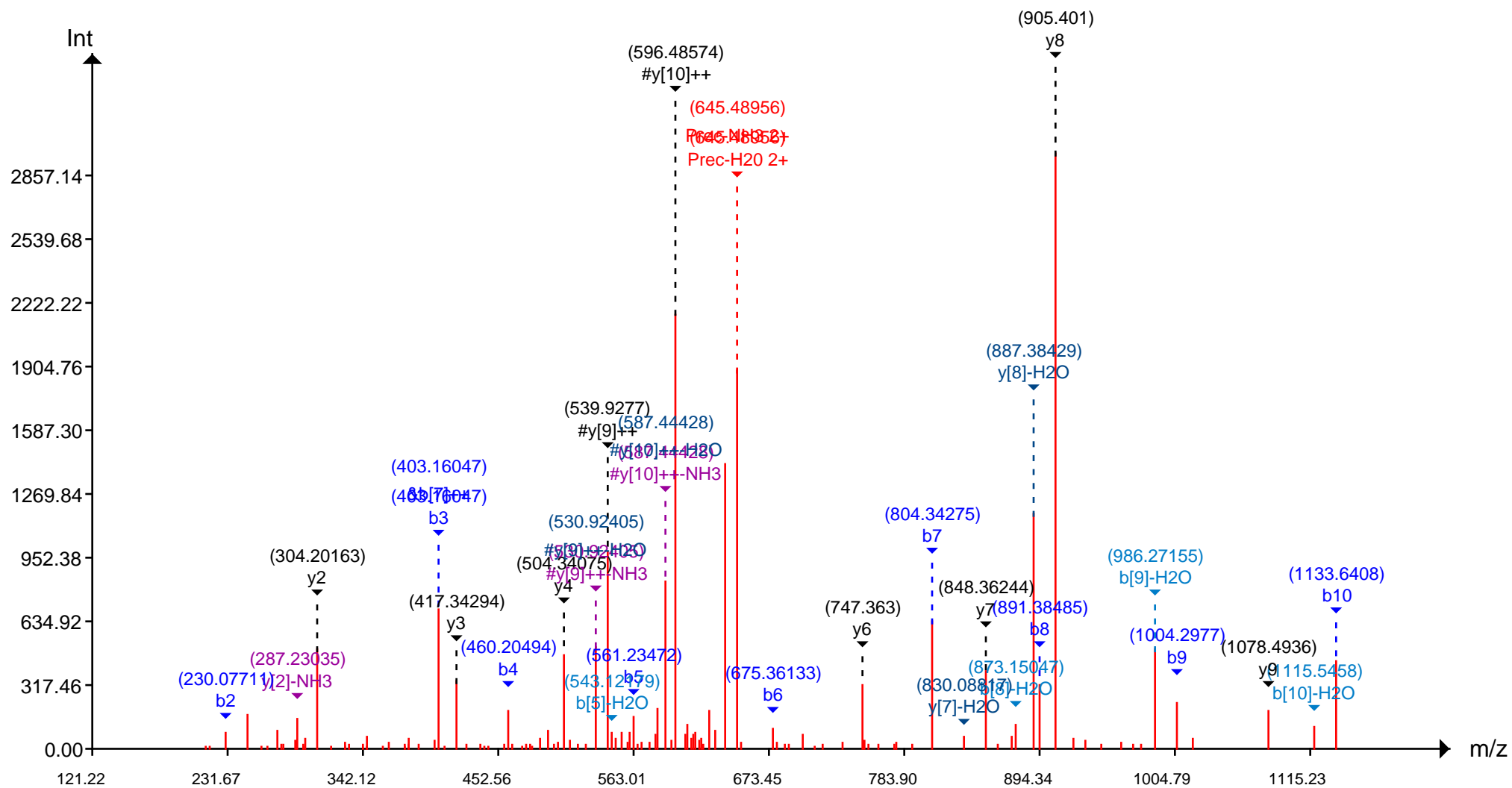


Info

| | | | |
|--------------------------|-----------------------------|---------------------------|----------|
| Modified Sequence | AcD3-AGC<Cmm*>QESAGPER-COOH | Ion score | 54.17 |
| Protein | Q99575 | Confidence level | 99.00% |
| Precursor m/z | 603.76582 | Identity Threshold | 43.79 |
| Charge | 2+ | Mass error | -3.22E-4 |

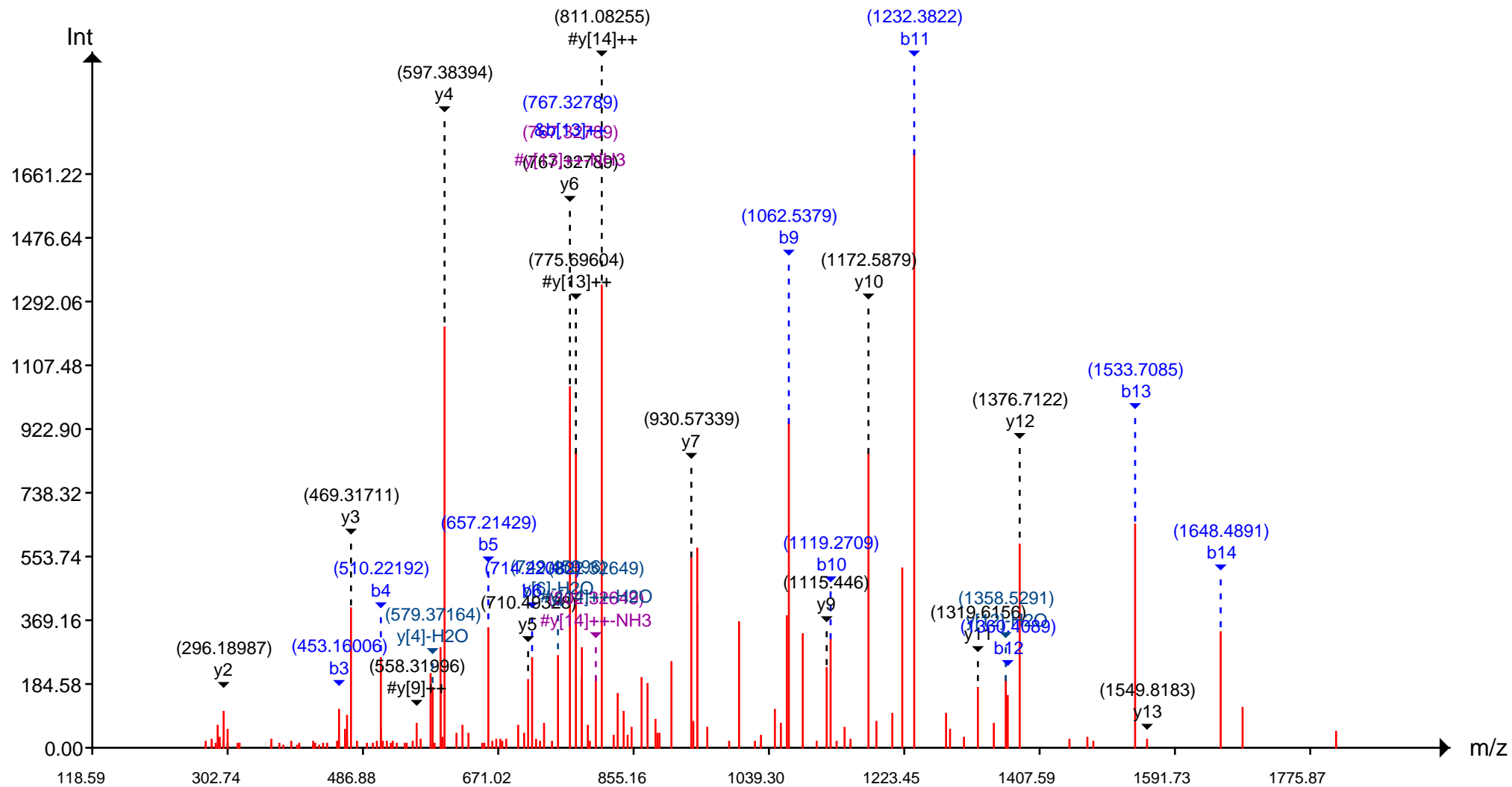


| Info | | | |
|--------------------------|---------------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-AQLQAEK<AcD3K*>QK<AcD3K*>AR-COOH | Ion score | 83.99 |
| Protein | Q13442 | Confidence level | 99.00% |
| Precursor m/z | 703.4092 | Identity Threshold | 47.10 |
| Charge | 2+ | Mass error | 3.66E-4 |



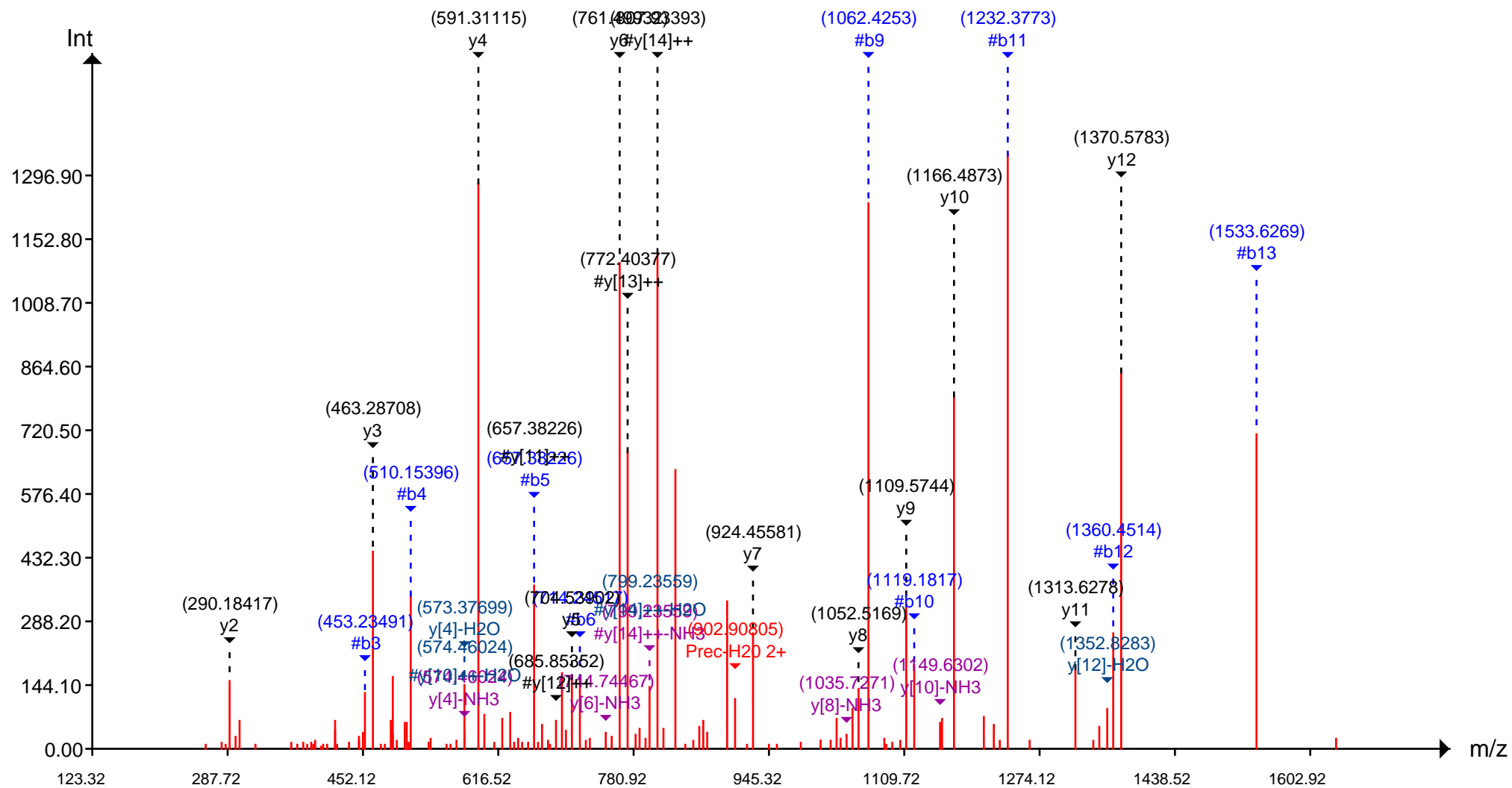
Info

| | | | |
|--------------------------|-------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-ALK<AcD3K*>GTNESLER-COOH | Ion score | 59.8 |
| Protein | P08670 | Confidence level | 99.00% |
| Precursor m/z | 654.35732 | Identity Threshold | 48.69 |
| Charge | 2+ | Mass error | 1.67E-4 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-YAK<AcD3K*>GFGGQYGIQK<AcD3K*>DR<C13>-COOH | Ion score | 75.92 |
| Protein | P14317 | Confidence level | 99.00% |
| Precursor m/z | 914.98527 | Identity Threshold | 49.12 |
| Charge | 2+ | Mass error | -1.43E-4 |



Info

Modified Sequence AcD3-YAK<AcD3K*>GFGGQYGIQK<AcD3K*>DR-COOH

Ion score 85.92

Protein P14317

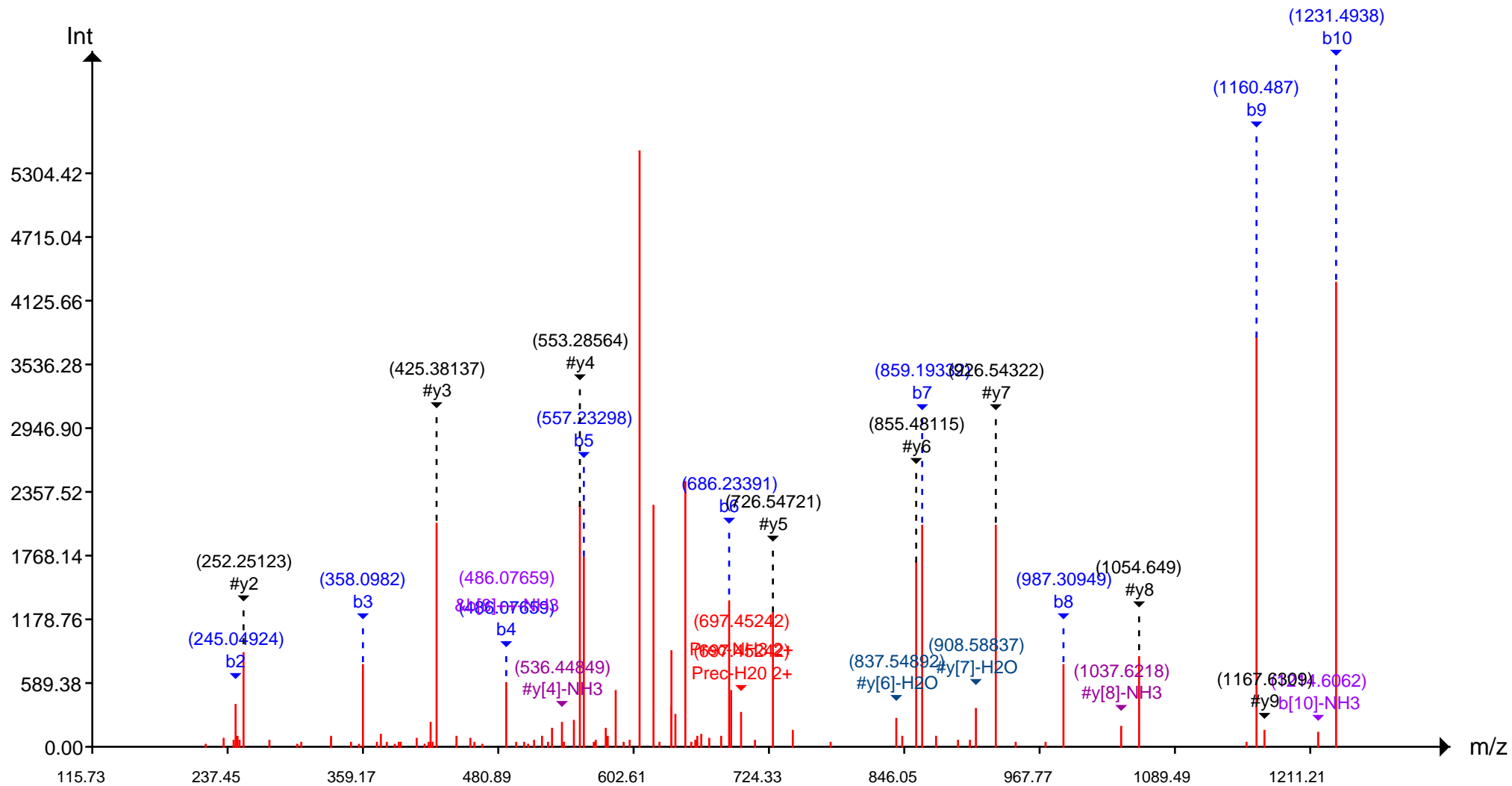
Confidence level 99.00%

Precursor m/z 911.97551

Identity Threshold 49.23

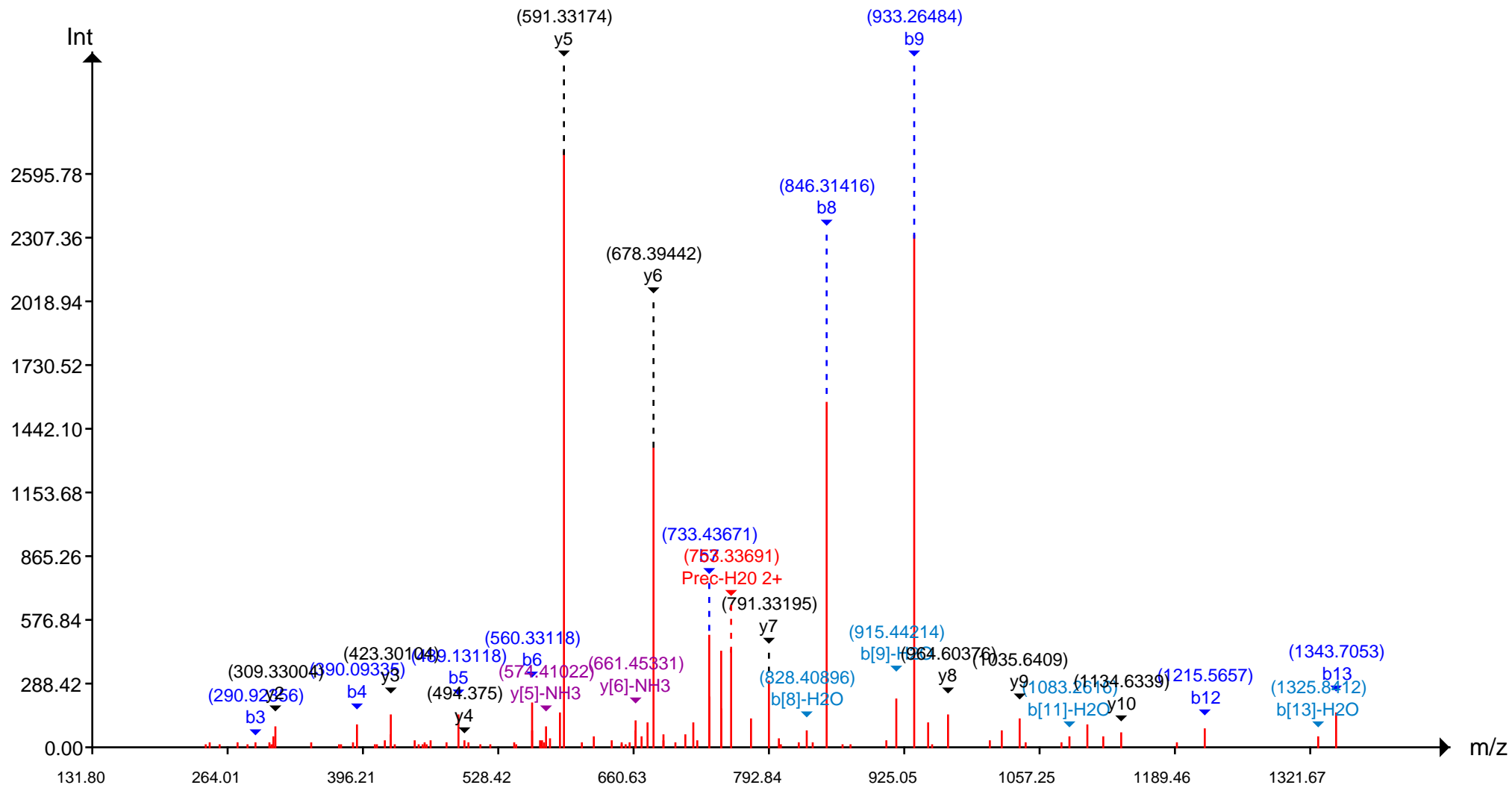
Charge 2+

Mass error 4.79E-4



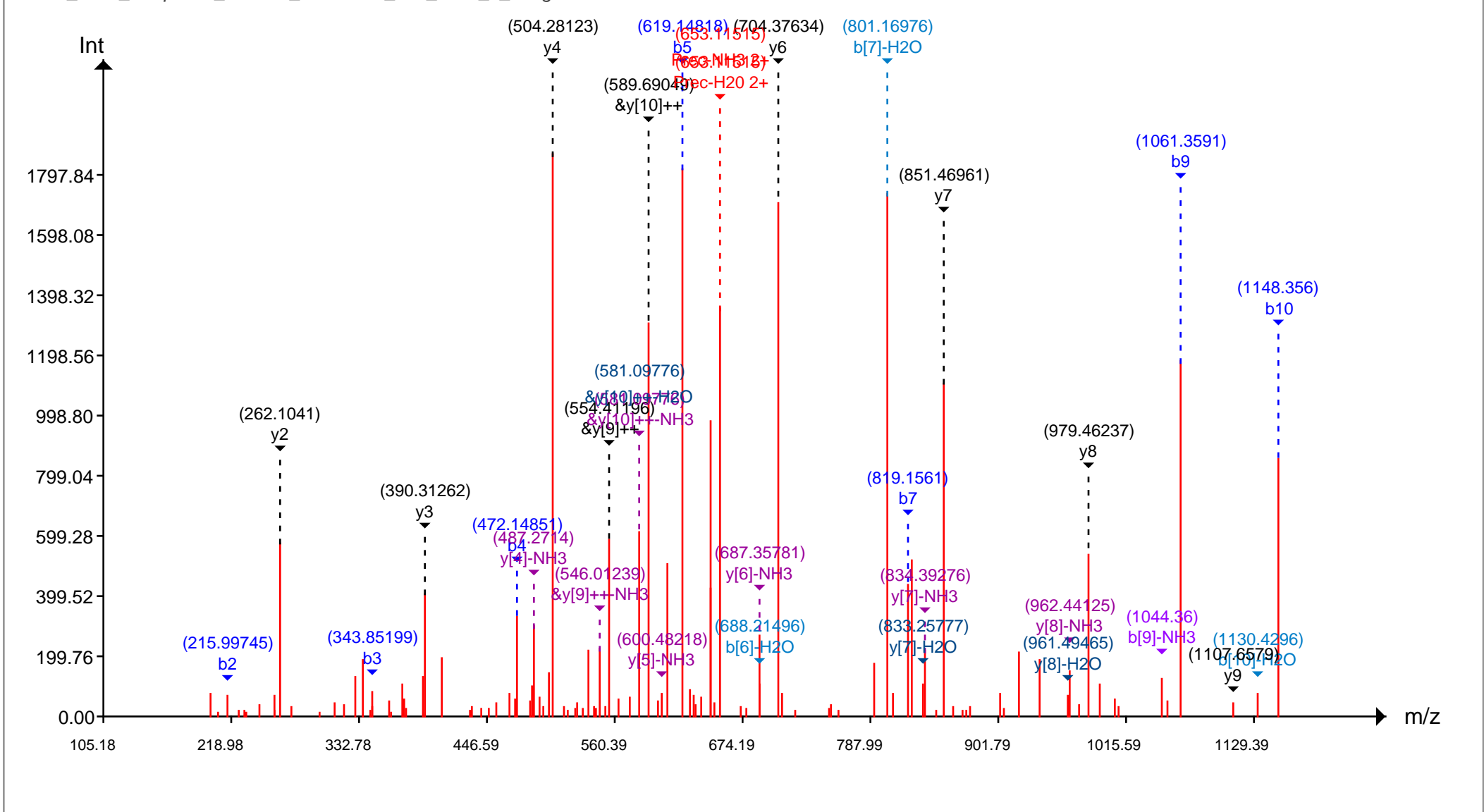
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-AQLQAEK<AcD3K*>QK<AcD3K*>AR<C13>-COOH | Ion score | 71.46 |
| Protein | Q13442 | Confidence level | 99.00% |
| Precursor m/z | 706.4191 | Identity Threshold | 45.62 |
| Charge | 2+ | Mass error | 2.4E-5 |



Info

| | | | |
|--------------------------|---------------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-STGVVAK<AcD3K*>LSPANQR<C13>-COOH | Ion score | 75.89 |
| Protein | Q13263 | Confidence level | 99.00% |
| Precursor m/z | 762.44152 | Identity Threshold | 46.68 |
| Charge | 2+ | Mass error | 3.39E-4 |



Info

Modified Sequence AcD3-VAQQFSLNQSR-COOH

Ion score 56.28

Protein O60216

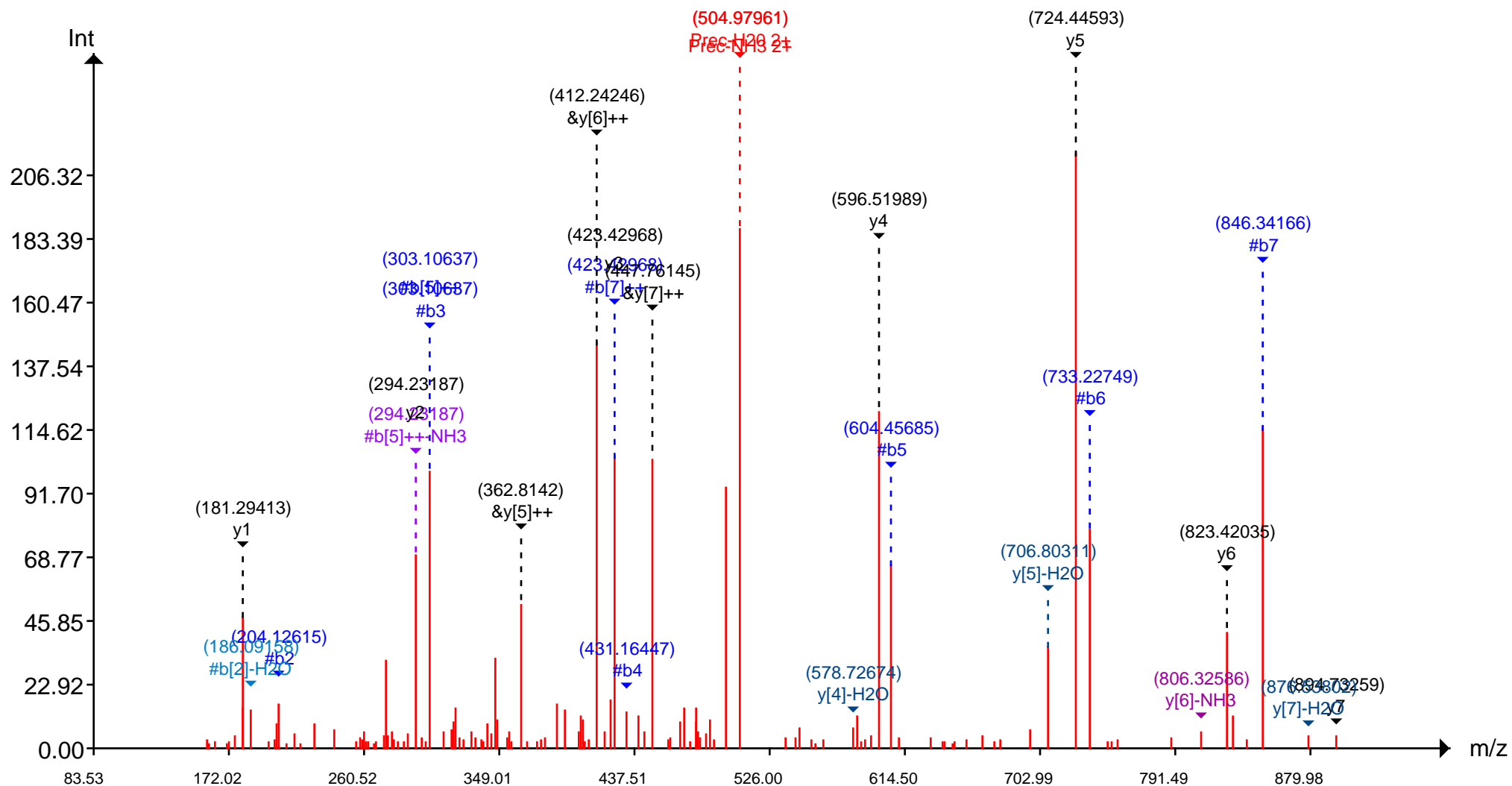
Confidence level 99.00%

Precursor m/z 661.84667

Identity Threshold 49.07

Charge 2+

Mass error -0.00299



Info

Modified Sequence AcD3-SAVQK<AcD3K*>ELR<C13>-COOH

Ion score 73.59

Protein Q9P265

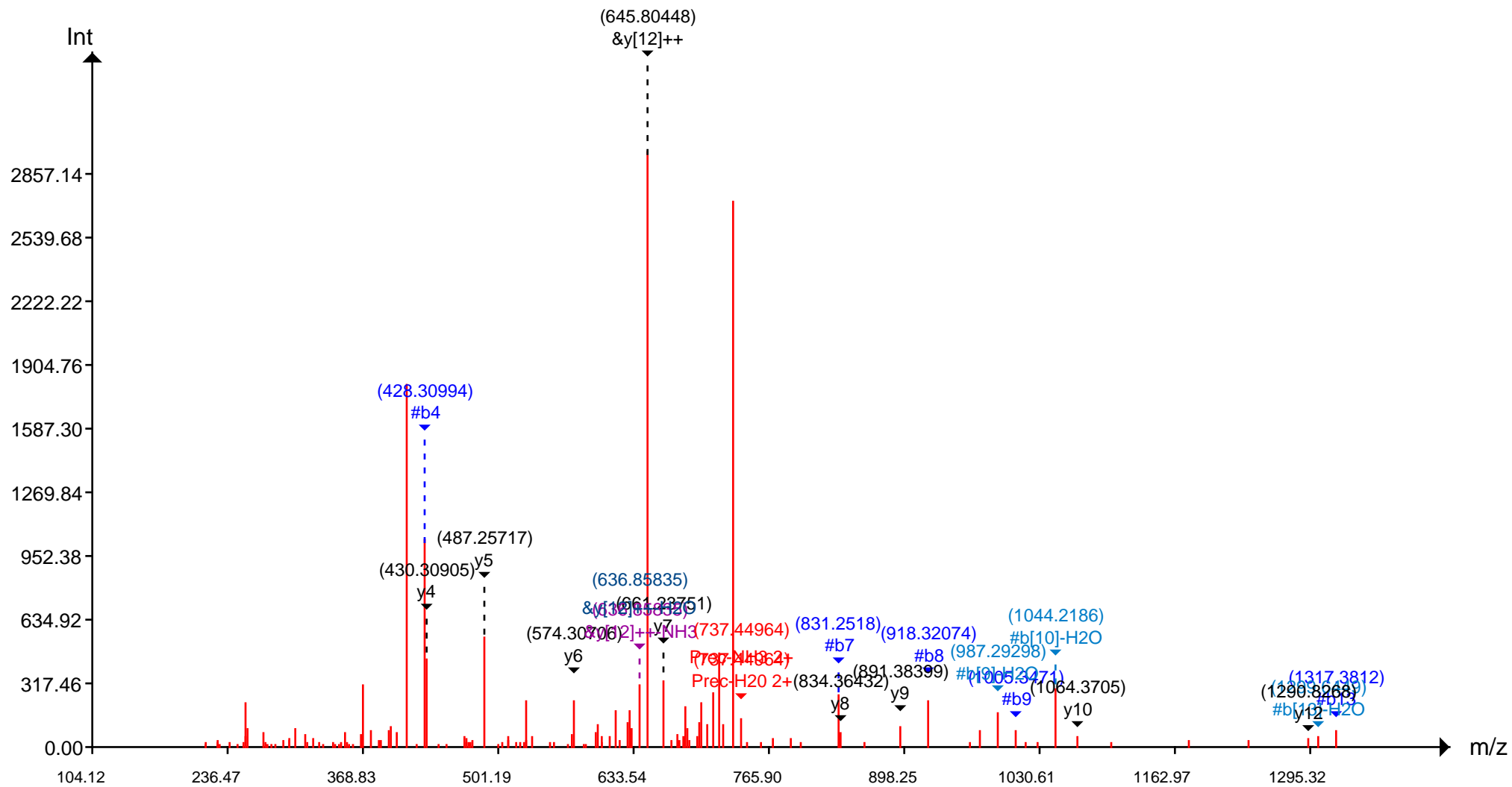
Confidence level 99.00%

Precursor m/z 513.81135

Identity Threshold 46.30

Charge 2+

Mass error -1.89E-4



Info

Modified Sequence AcD3-GVPEK<AcD3K*>GK<AcD3K*>SSGPSAR-COOH

Ion score 51.13

Protein Q96IZ0

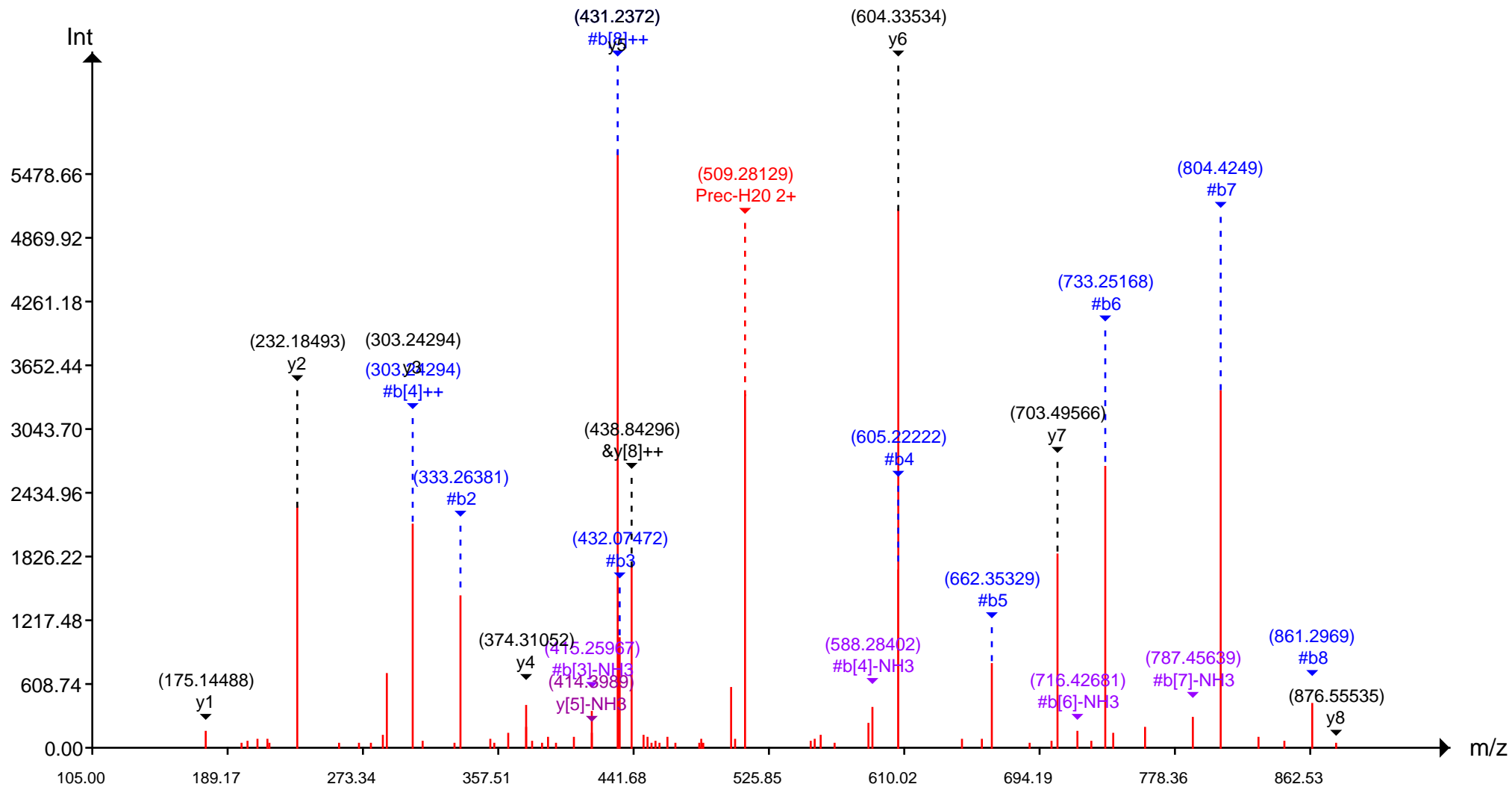
Confidence level 99.00%

Precursor m/z 746.41346

Identity Threshold 48.56

Charge 2+

Mass error 0.008459



Info

Modified Sequence AcD3-NK<AcD3K*>VK<AcD3K*>GAAGR-COOH

Ion score 74.23

Protein P78527

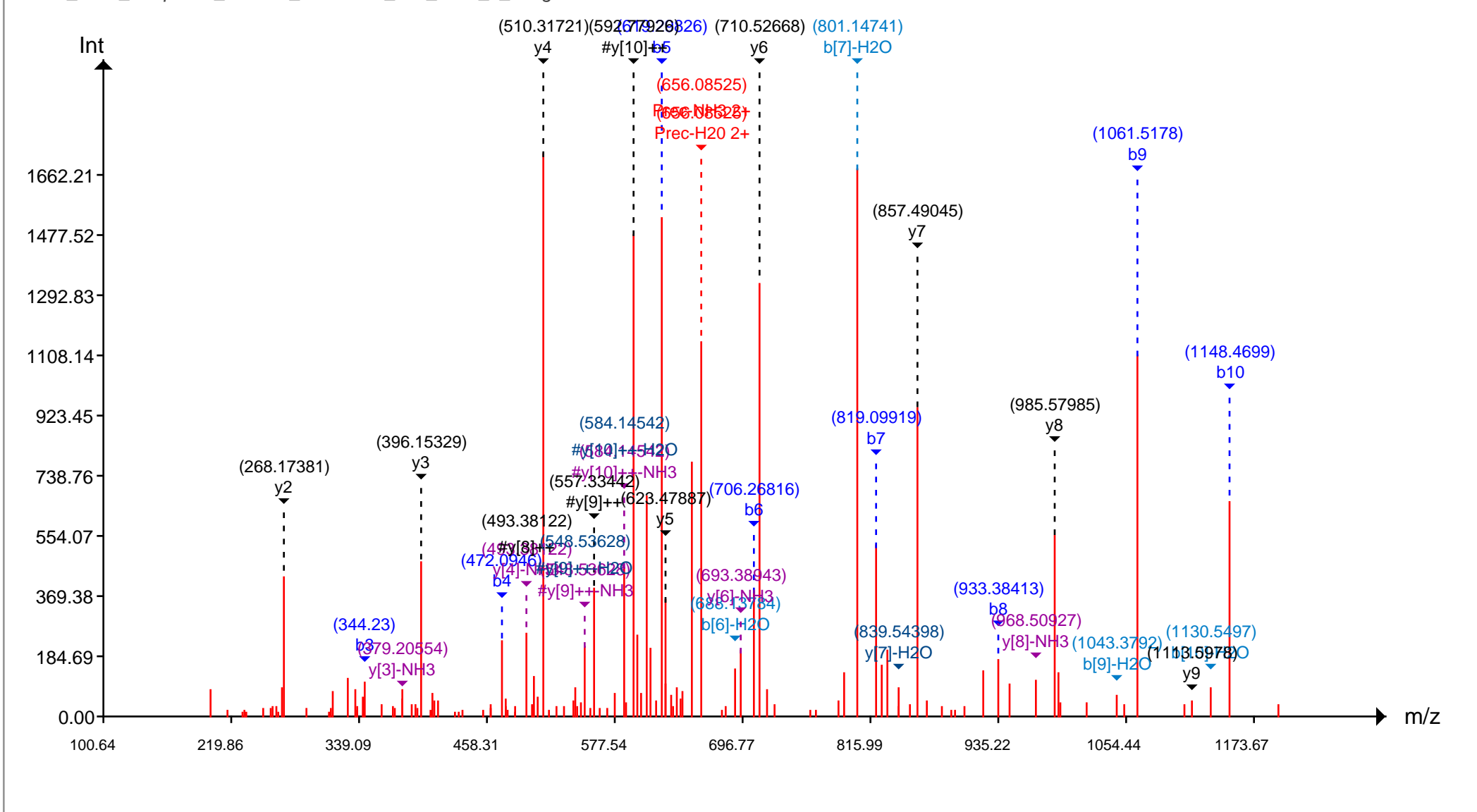
Confidence level 99.00%

Precursor m/z 518.31674

Identity Threshold 45.74

Charge 2+

Mass error 6.57E-4



Info

Modified Sequence AcD3-VAQQFSLNQSR<C13>-COOH

Ion score 65.48

Protein O60216

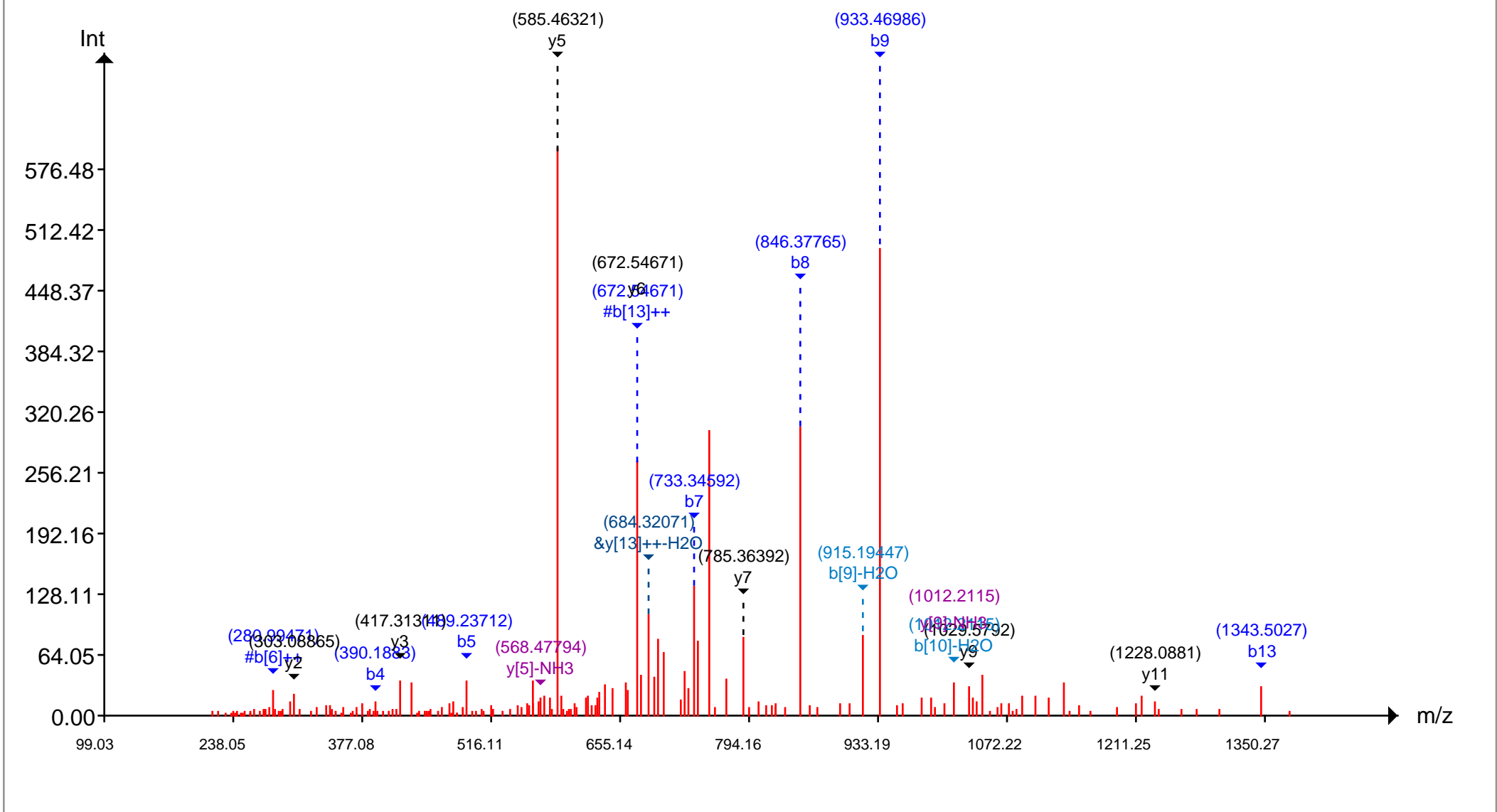
Confidence level 99.00%

Precursor m/z 664.8574

Identity Threshold 49.00

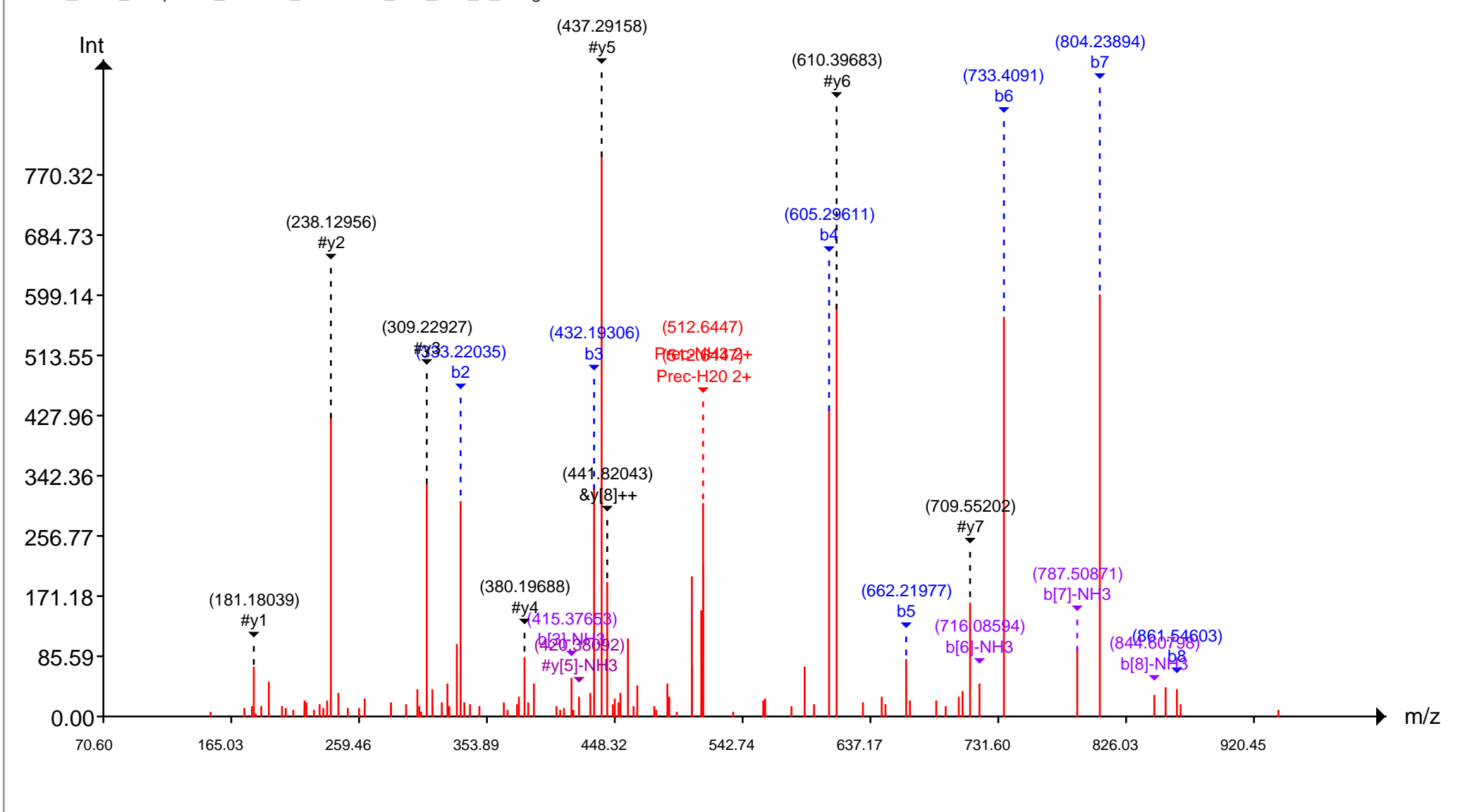
Charge 2+

Mass error -0.001672



Info

| | | | |
|--------------------------|----------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-STGVVAK<AcD3K*>LSPANQR-COOH | Ion score | 48.13 |
| Protein | Q13263 | Confidence level | 99.00% |
| Precursor m/z | 759.43162 | Identity Threshold | 47.76 |
| Charge | 2+ | Mass error | 6.81E-4 |



Info

Modified Sequence AcD3-NK<AcD3K*>VK<AcD3K*>GAAGR<C13>-COOH

Ion score 55.29

Protein P78527

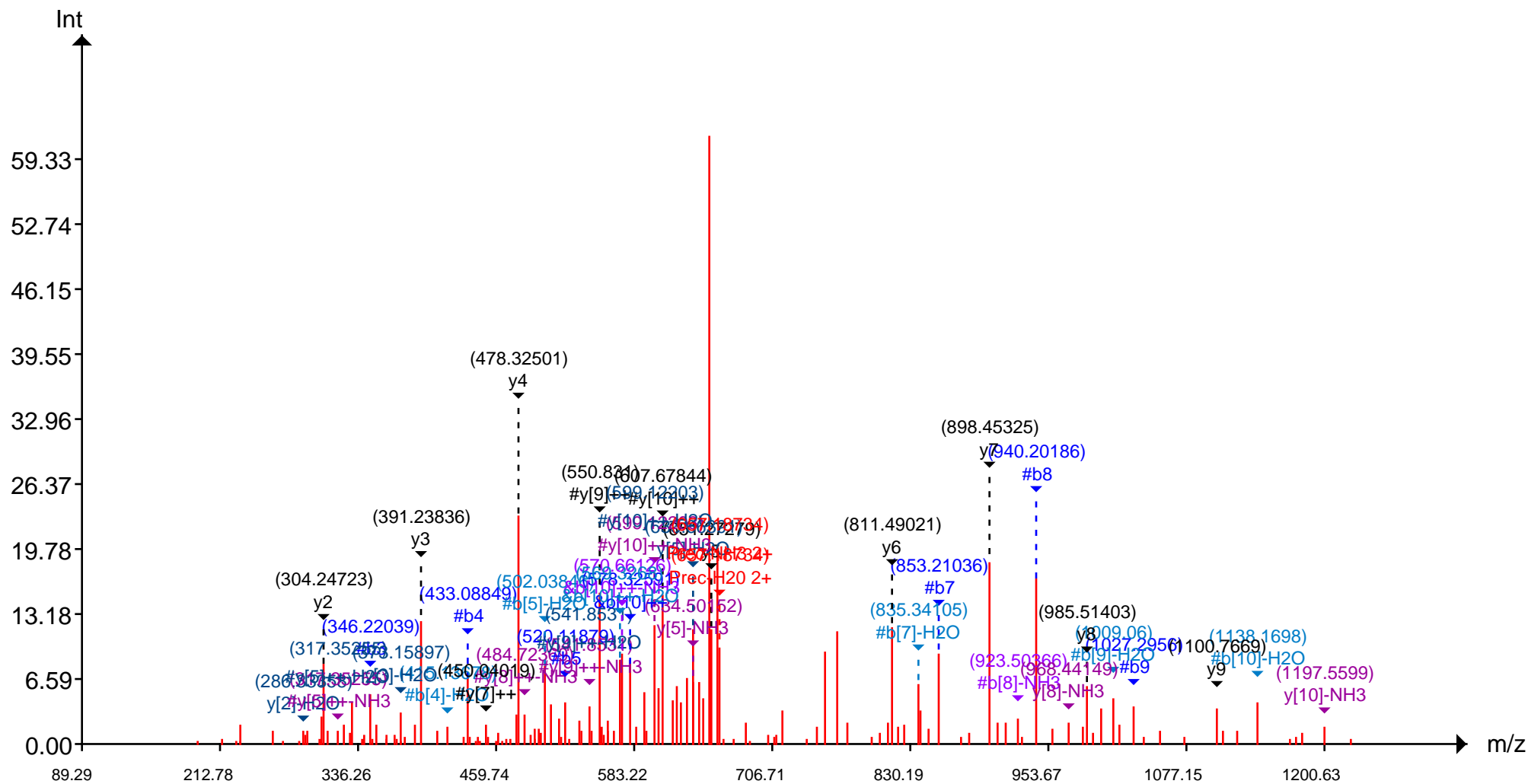
Confidence level 99.00%

Precursor m/z 521.3265

Identity Threshold 43.74

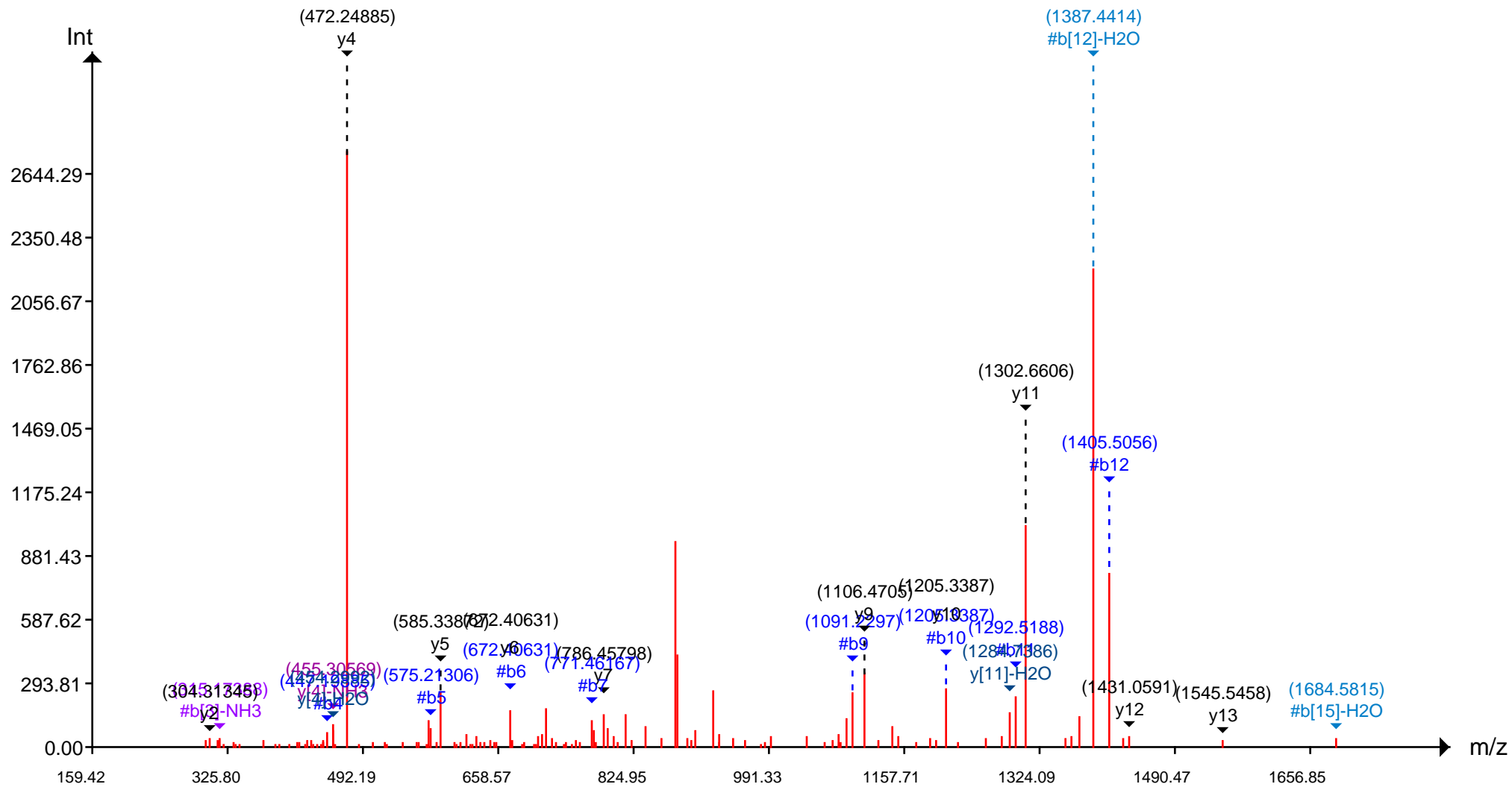
Charge 2+

Mass error 3.6E-5



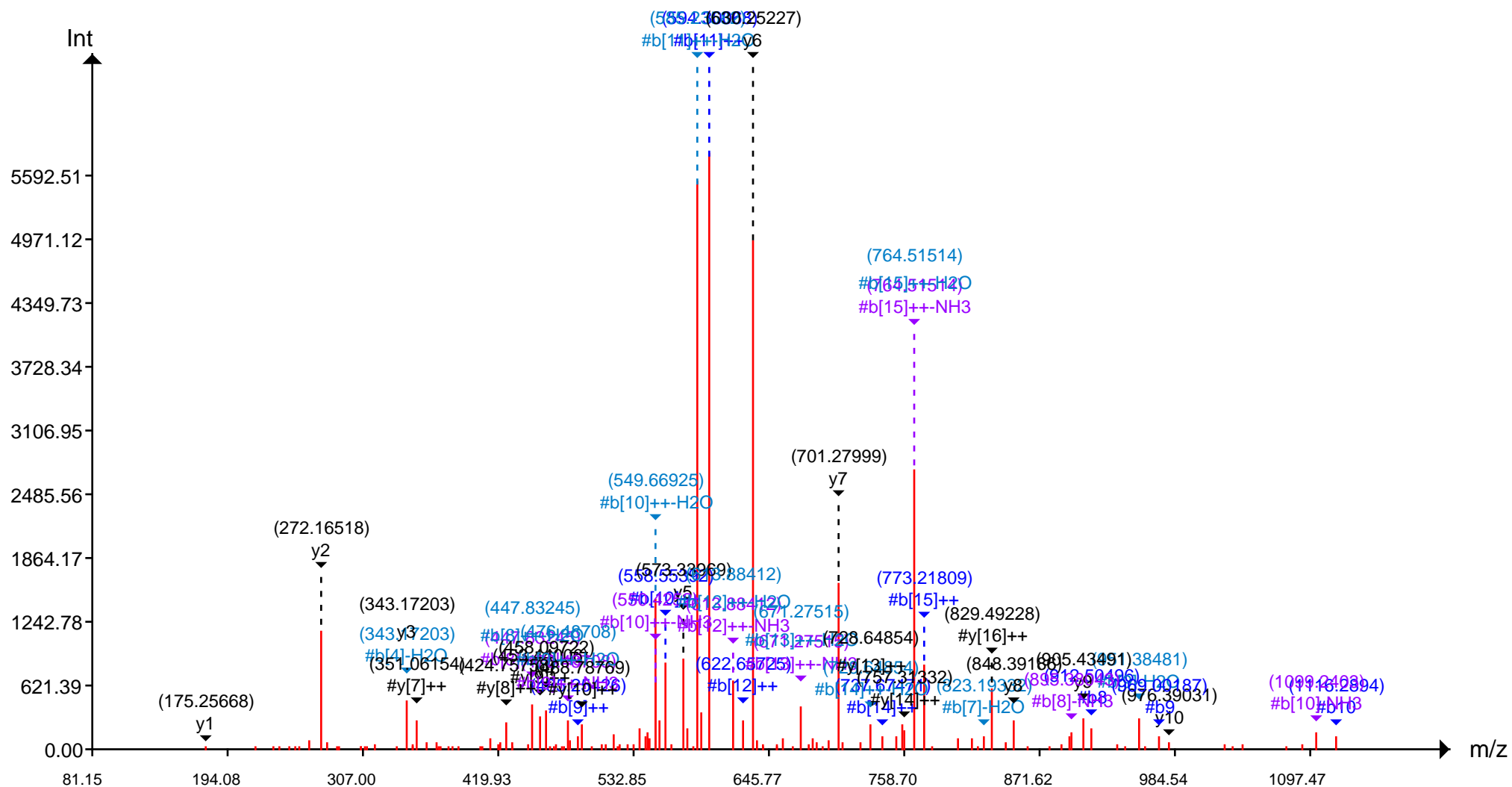
Info

| | | | |
|--------------------------|-------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-ANDSSC<Cmm*>K<AcD3K*>SSER-COOH | Ion score | 49.88 |
| Protein | Q9Y2H0 | Confidence level | 99.00% |
| Precursor m/z | 665.7937 | Identity Threshold | 43.71 |
| Charge | 2+ | Mass error | -8.98E-4 |



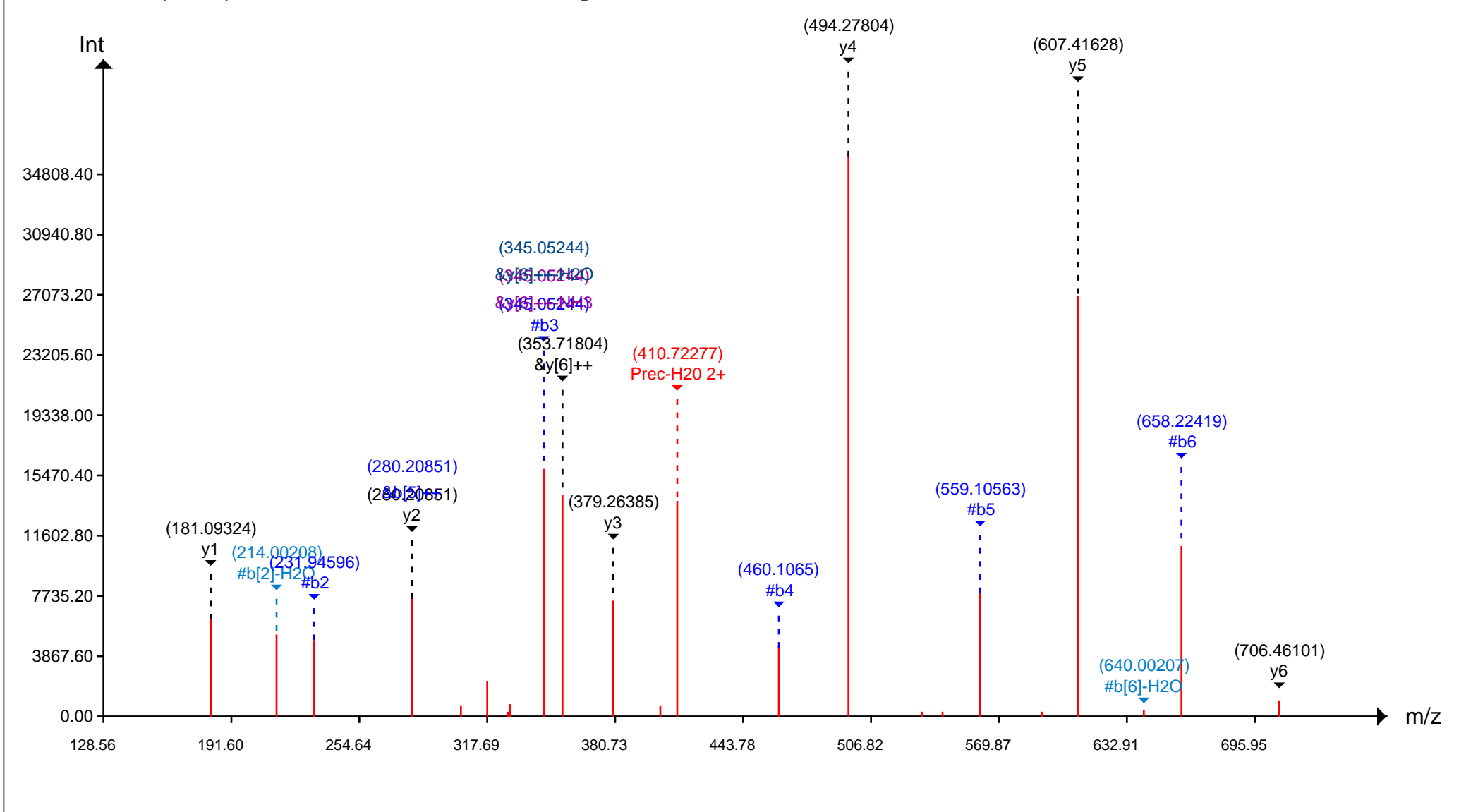
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-SAQDQPVK<AcD3K*>M<Mox*>NSLPAER-COOH | Ion score | 77.07 |
| Protein | P30419 | Confidence level | 99.00% |
| Precursor m/z | 938.97079 | Identity Threshold | 48.81 |
| Charge | 2+ | Mass error | 3.16E-4 |



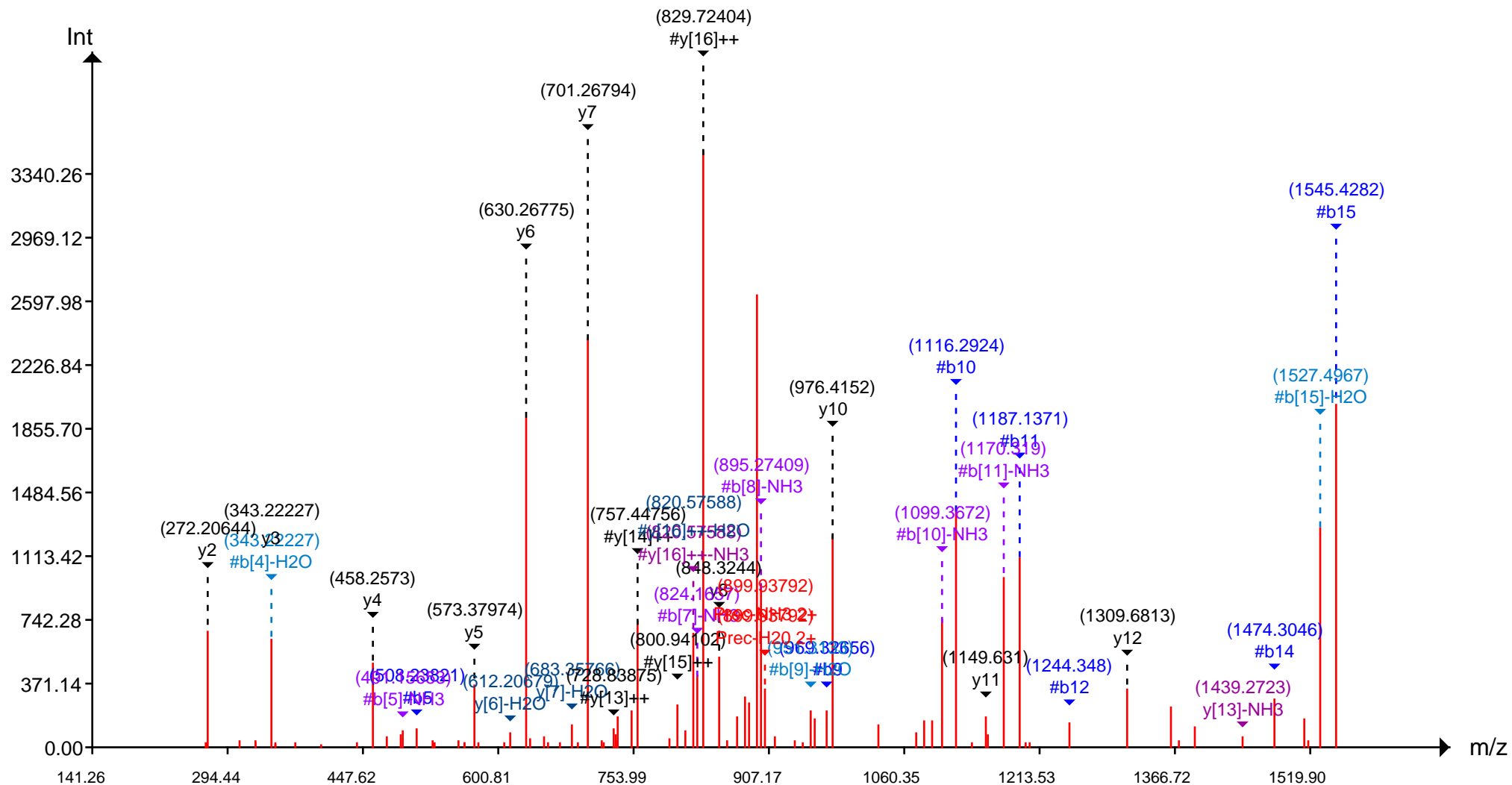
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH | Ion score | 87.21 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 606.26681 | Identity Threshold | 42.96 |
| Charge | 3+ | Mass error | -1.0E-4 |



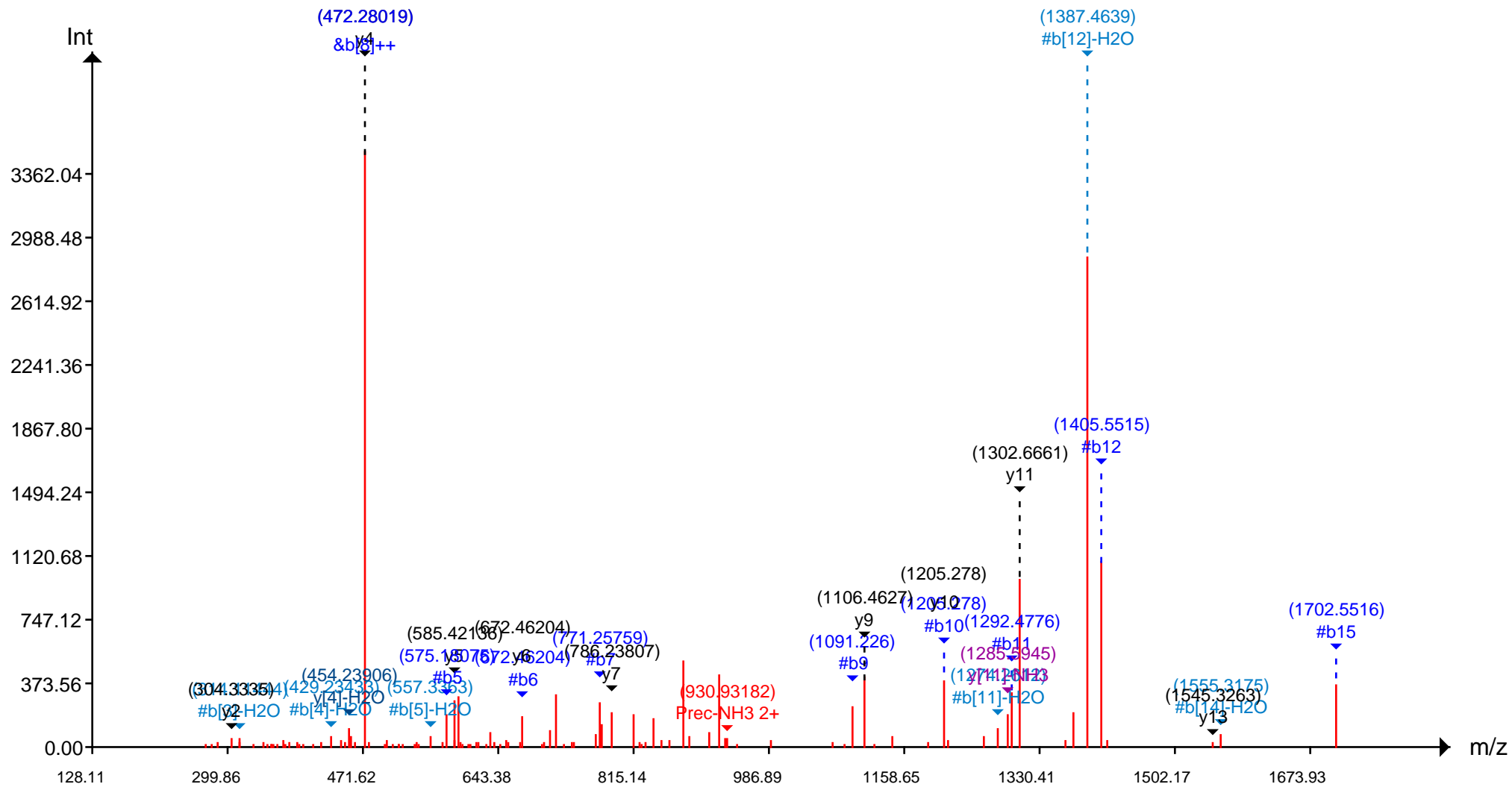
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-SVLDVVR<C13>-COOH | Ion score | 58.45 |
| Protein | Q13885, Q9BVA1, P68371, Q13509, P07437 | Confidence level | 99.00% |
| Precursor m/z | 419.762 | Identity Threshold | 45.46 |
| Charge | 2+ | Mass error | -4.3E-5 |



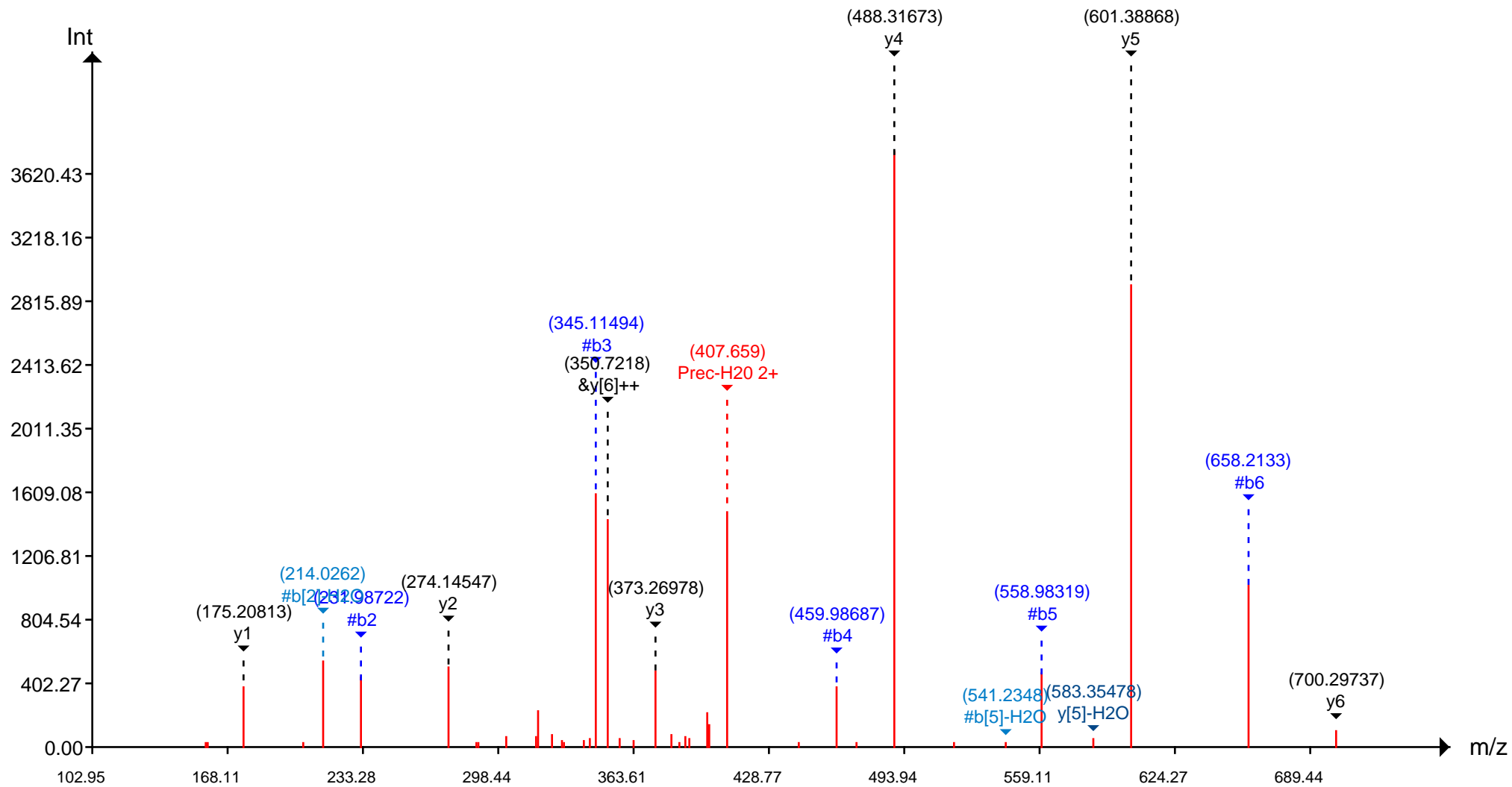
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH | Ion score | 80.83 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 908.89686 | Identity Threshold | 42.99 |
| Charge | 2+ | Mass error | 4.66E-4 |



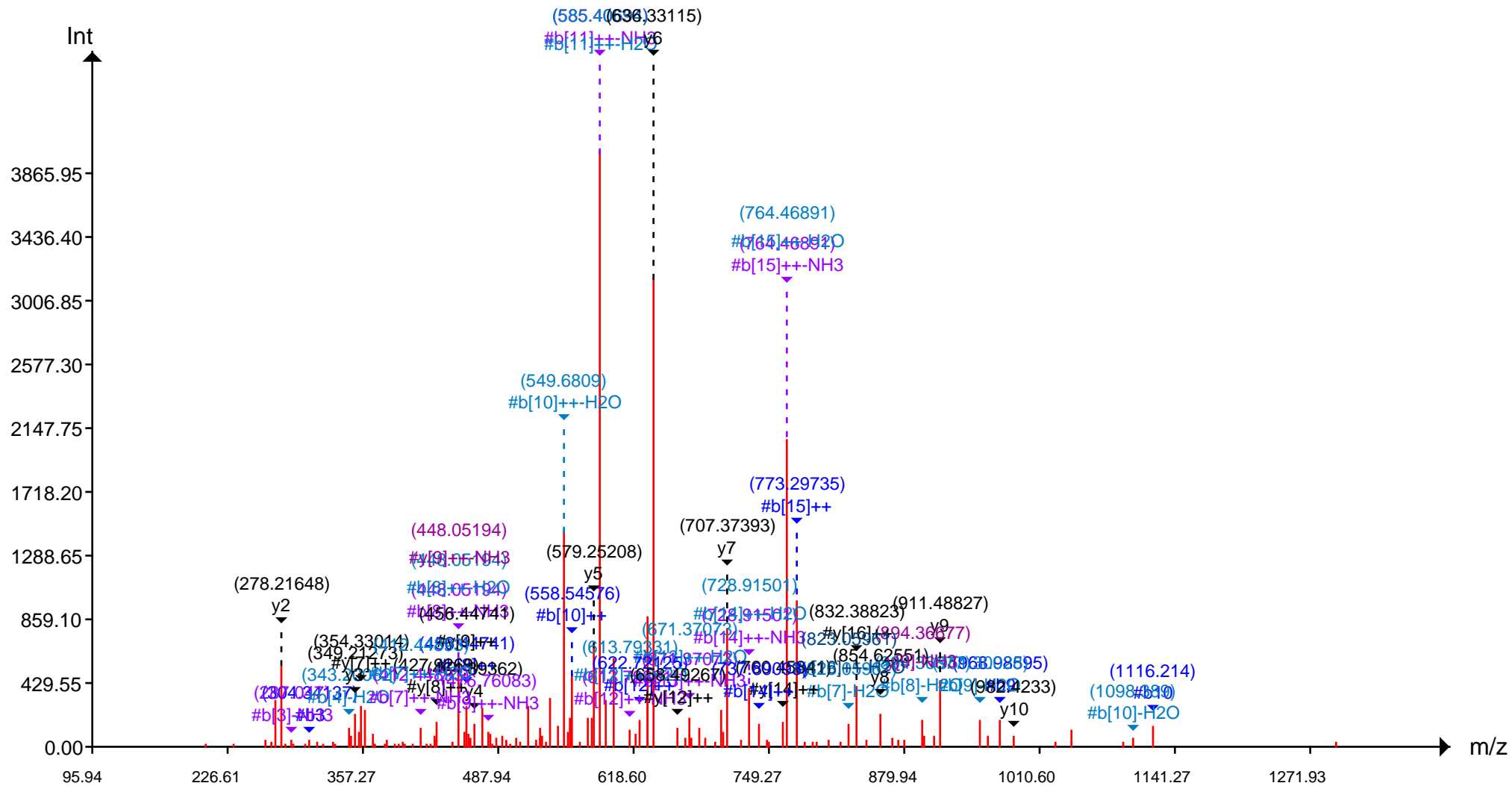
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-SAQDQPVK<AcD3K*>M<Mox*>NSLPAER-COOH | Ion score | 71.83 |
| Protein | P30419 | Confidence level | 99.00% |
| Precursor m/z | 938.971 | Identity Threshold | 48.84 |
| Charge | 2+ | Mass error | 7.36E-4 |



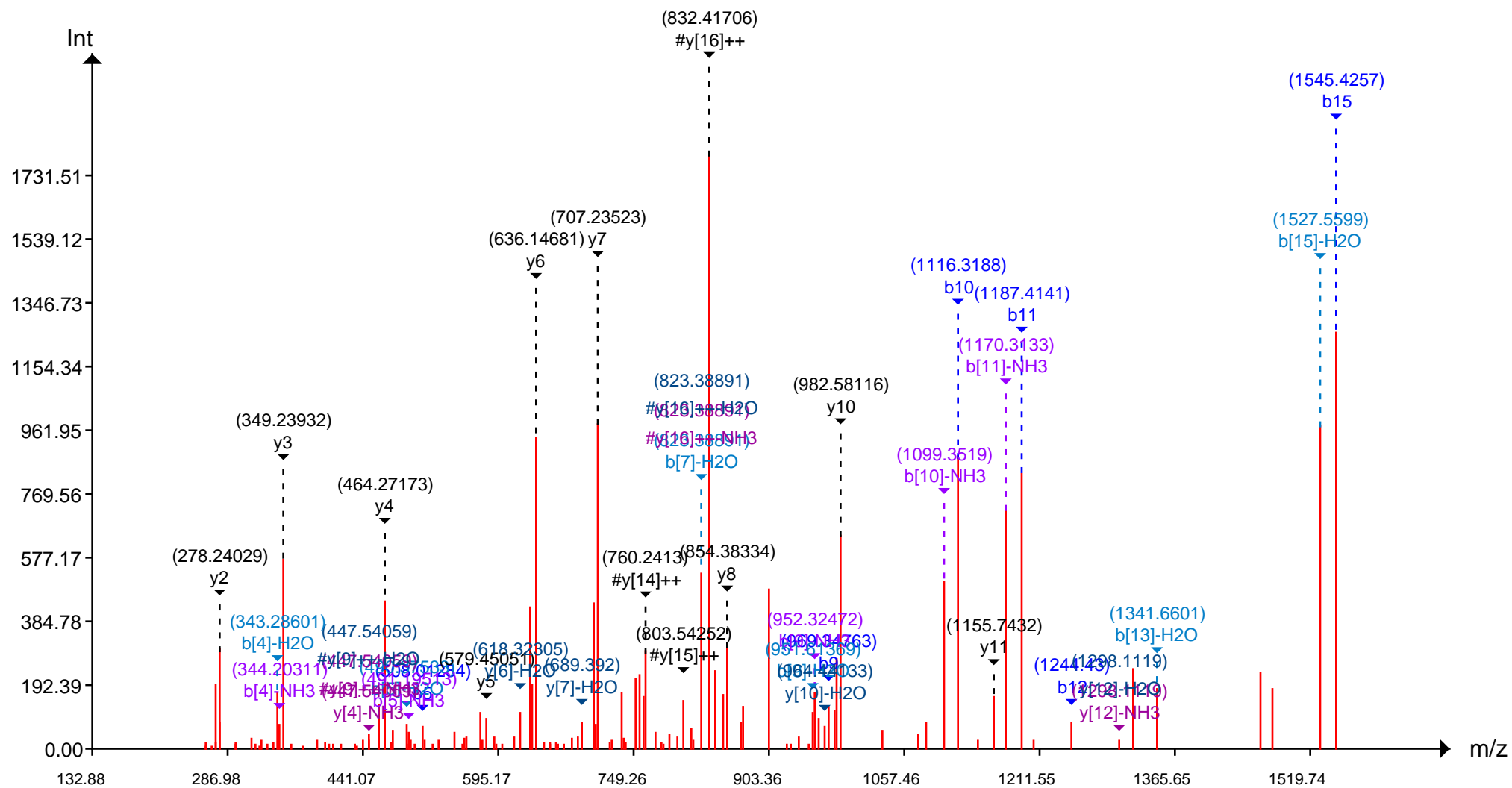
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-SVLDVVR-COOH | Ion score | 61.74 |
| Protein | Q13885, Q9BVA1, P68371, Q13509, P07437 | Confidence level | 99.00% |
| Precursor m/z | 416.75204 | Identity Threshold | 47.58 |
| Charge | 2+ | Mass error | 1.79E-4 |



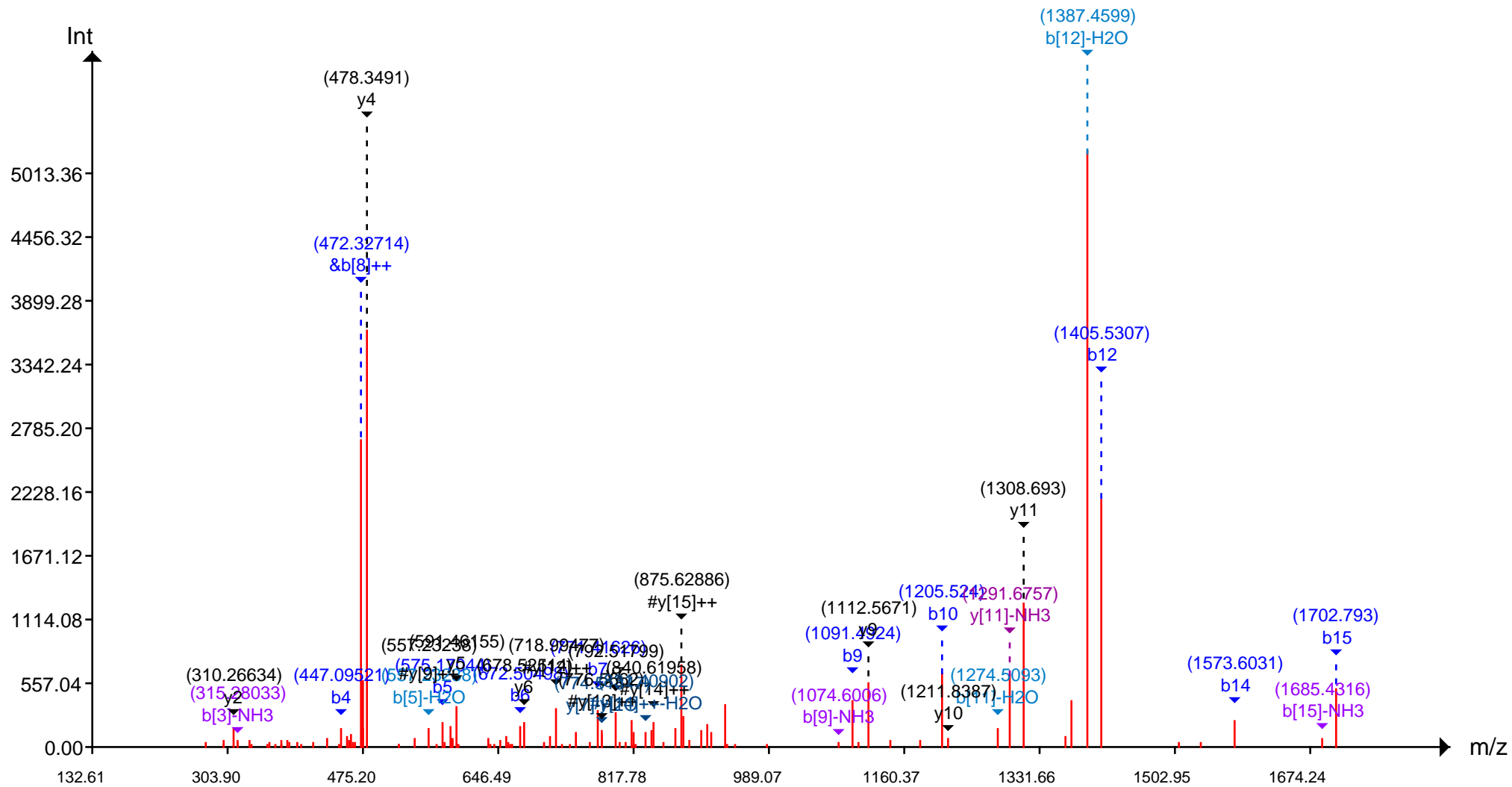
Info

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|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 57.99 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 608.27339 | Identity Threshold | 44.19 |
| Charge | 3+ | Mass error | -5.01E-4 |



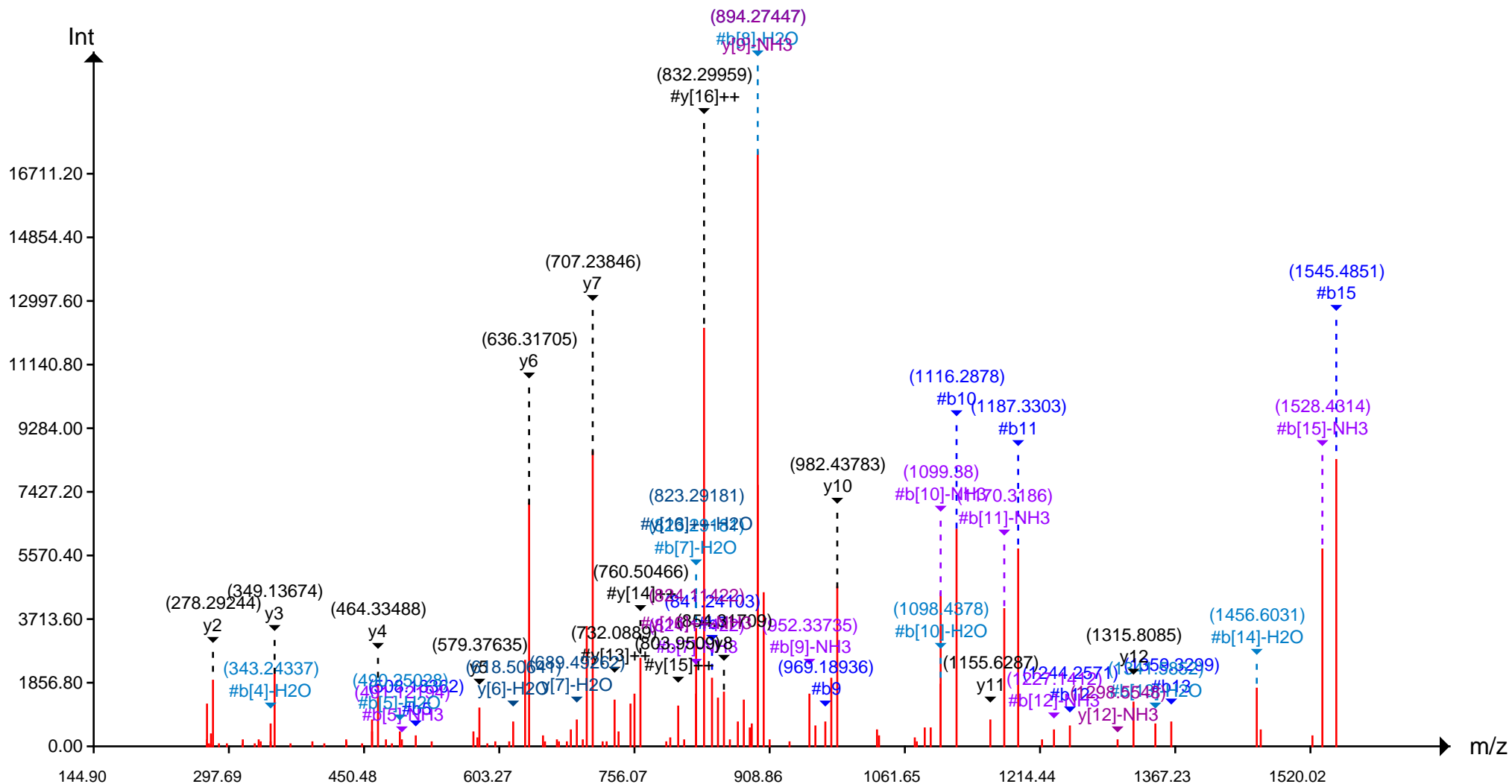
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 54.86 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 911.90679 | Identity Threshold | 44.25 |
| Charge | 2+ | Mass error | 1.85E-4 |



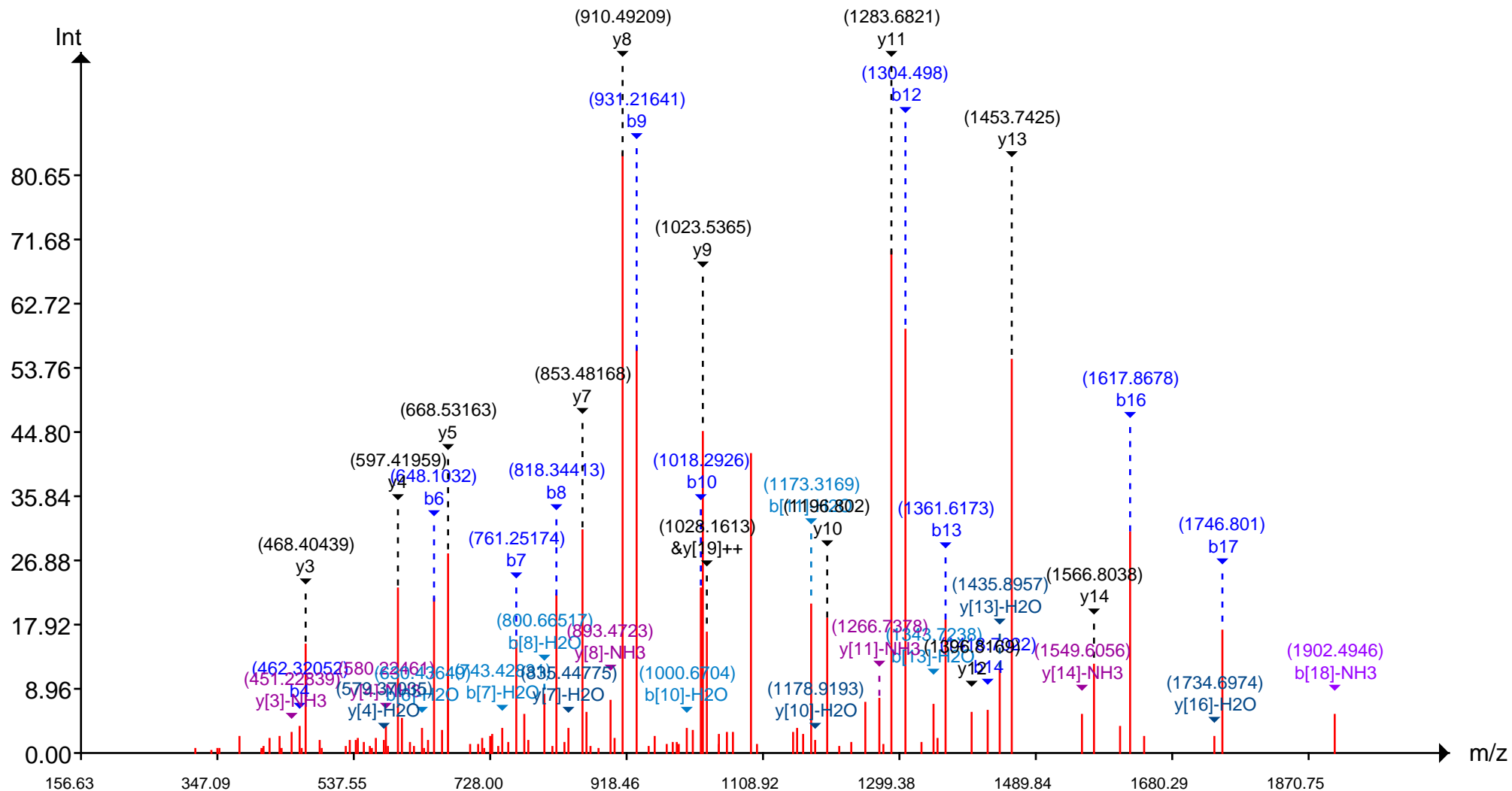
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-SAQDQPVK<AcD3K*>M<Mox*>NSLPAER<C13>-COOH | Ion score | 63.36 |
| Protein | P30419 | Confidence level | 99.00% |
| Precursor m/z | 941.98093 | Identity Threshold | 48.86 |
| Charge | 2+ | Mass error | 4.54E-4 |



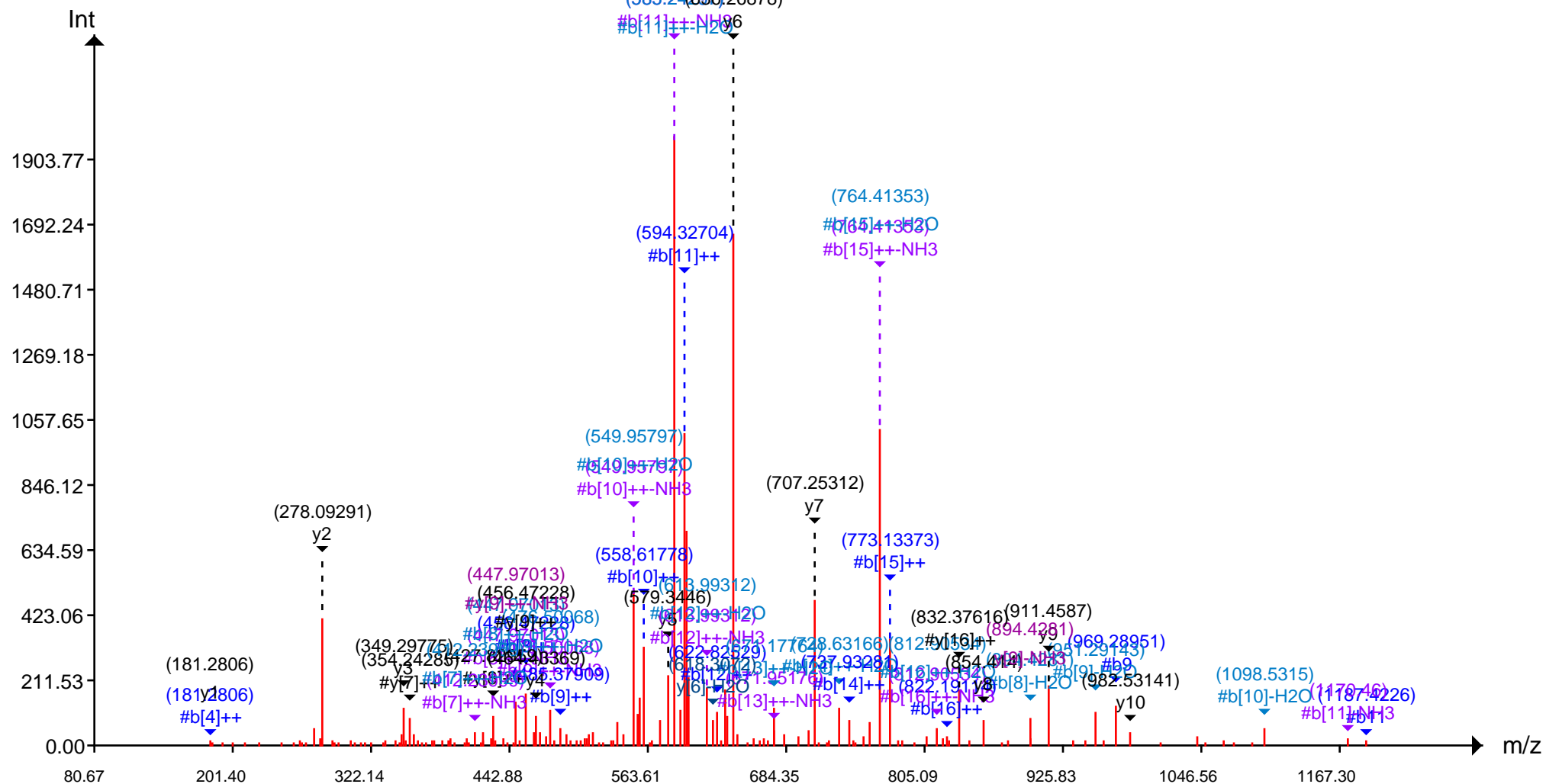
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 61.03 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 911.9069 | Identity Threshold | 44.24 |
| Charge | 2+ | Mass error | 4.05E-4 |



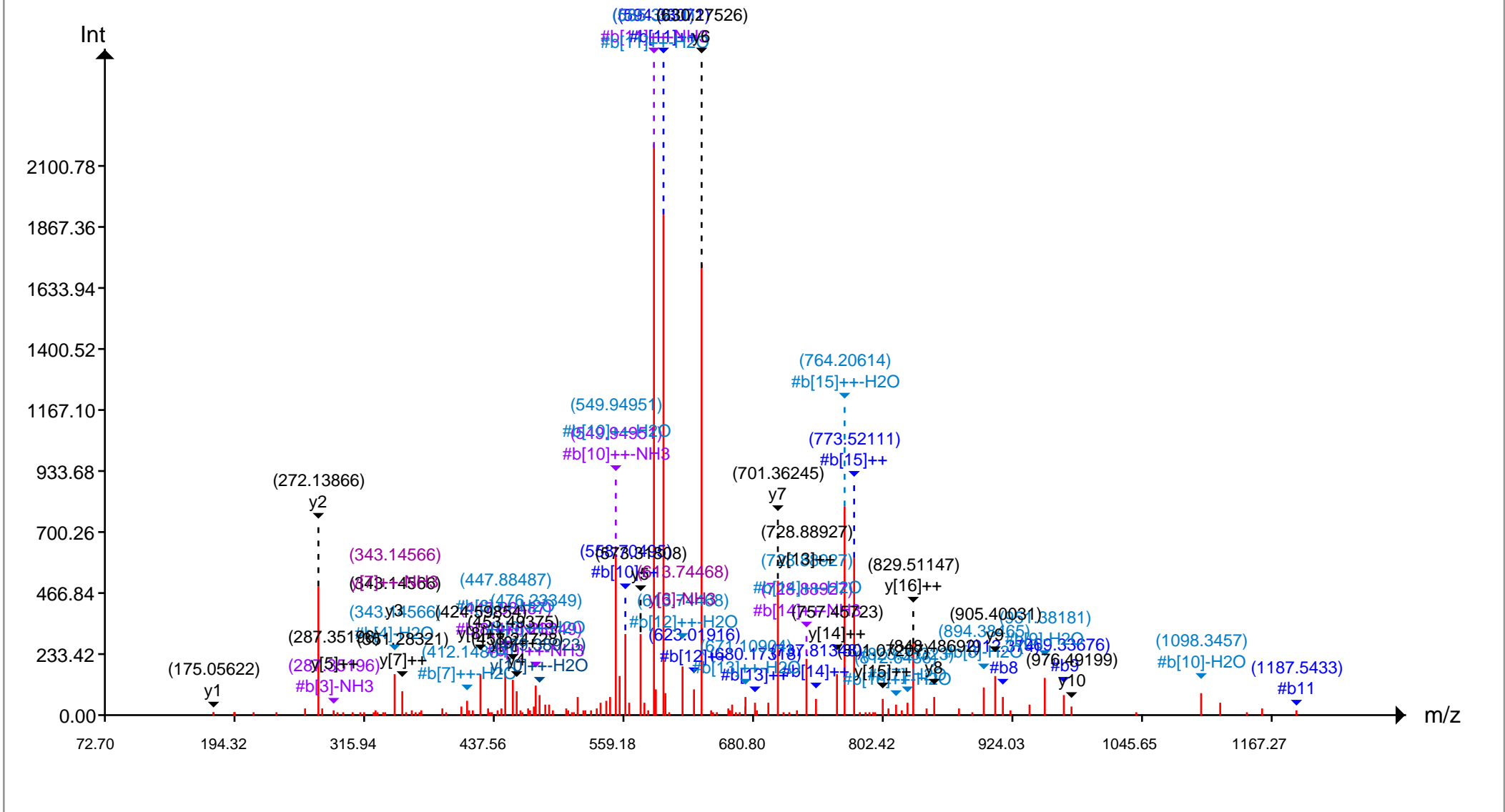
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-NSSQDALGLSK<AcD3K*>LGGQAEK<AcD3K*>NR<C13>-COOH | Ion score | 94.22 |
| Protein | Q13136 | Confidence level | 99.00% |
| Precursor m/z | 1107.5922 | Identity Threshold | 48.64 |
| Charge | 2+ | Mass error | 5.06E-4 |



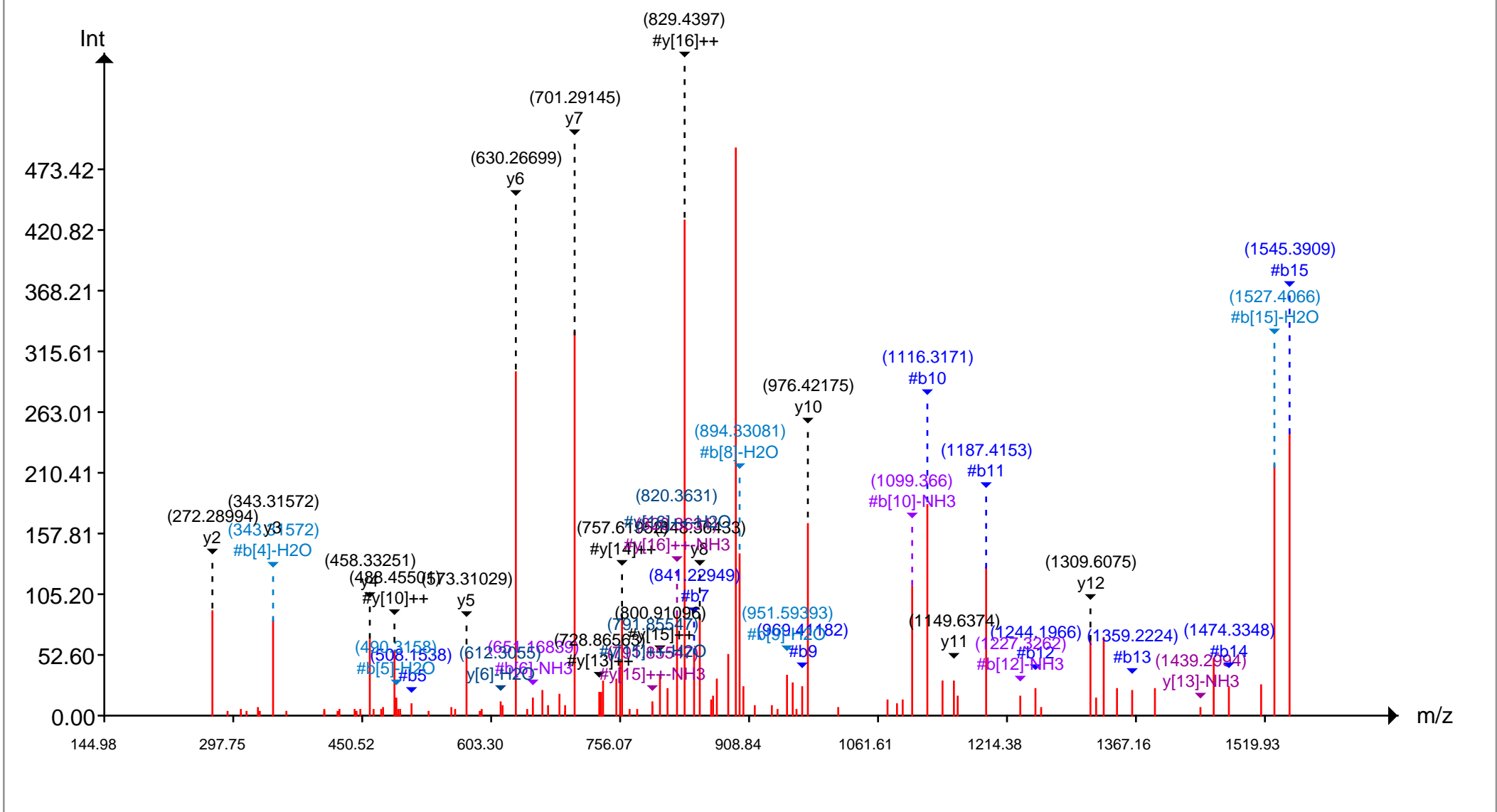
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 65.21 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 608.27356 | Identity Threshold | 44.24 |
| Charge | 3+ | Mass error | 9.0E-6 |



Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH | Ion score | 92.04 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 606.26684 | Identity Threshold | 42.97 |
| Charge | 3+ | Mass error | -1.0E-5 |



Info

Modified Sequence AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH

Ion score 76.28

Protein P60709, P63261, Q6S8J3, A5A3E0

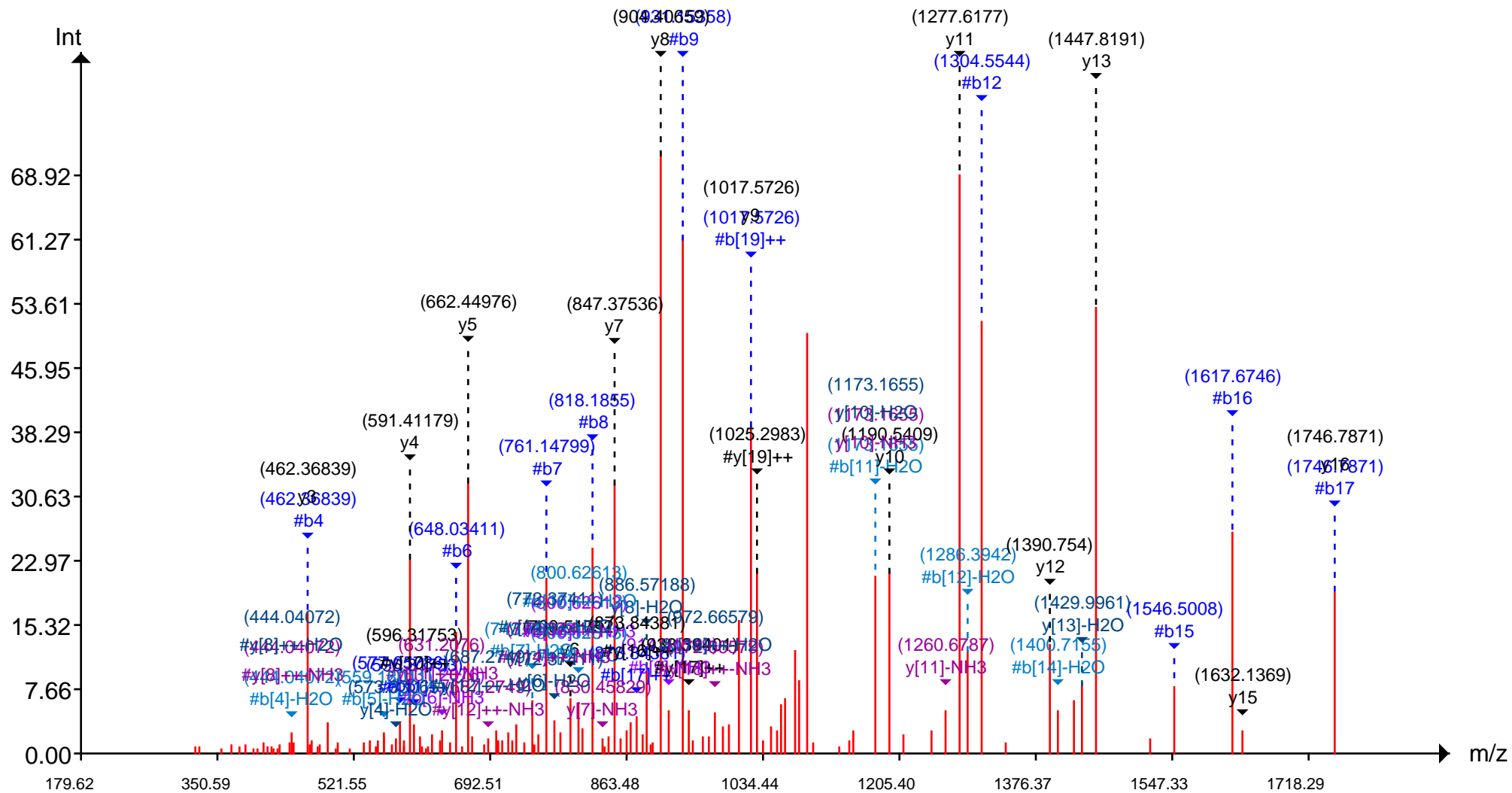
Confidence level 99.00%

Precursor m/z 908.89694

Identity Threshold 43.00

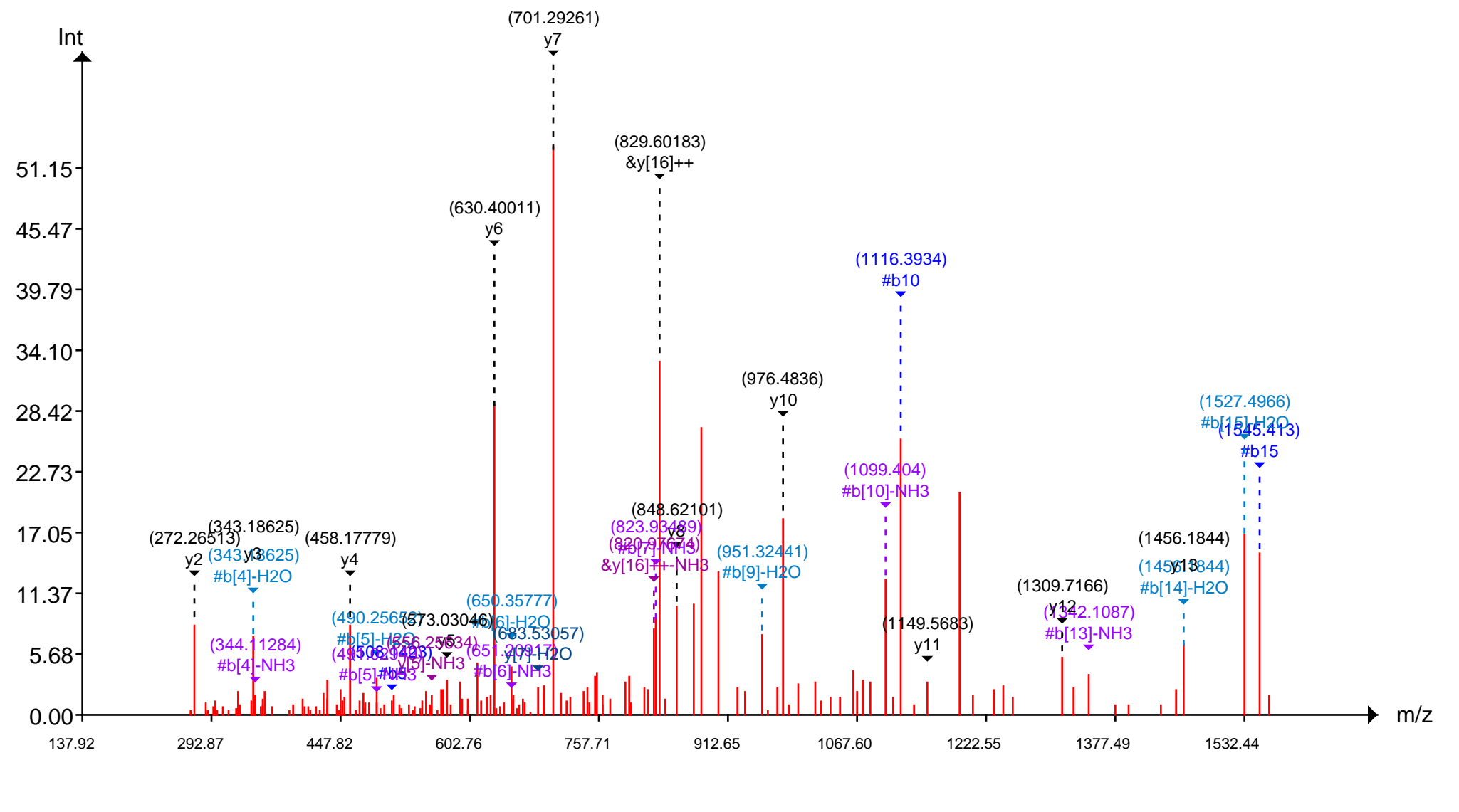
Charge 2+

Mass error 6.26E-4

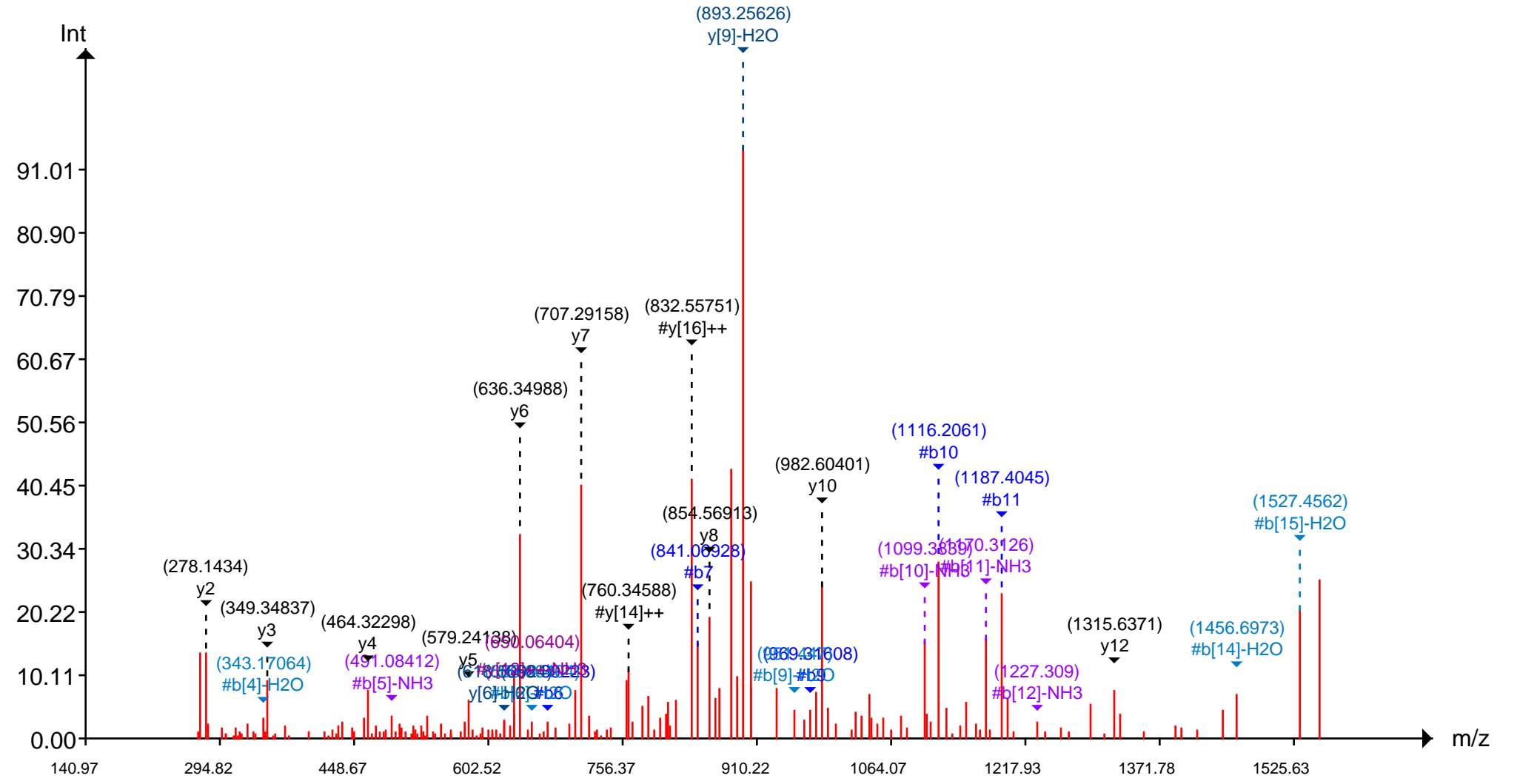


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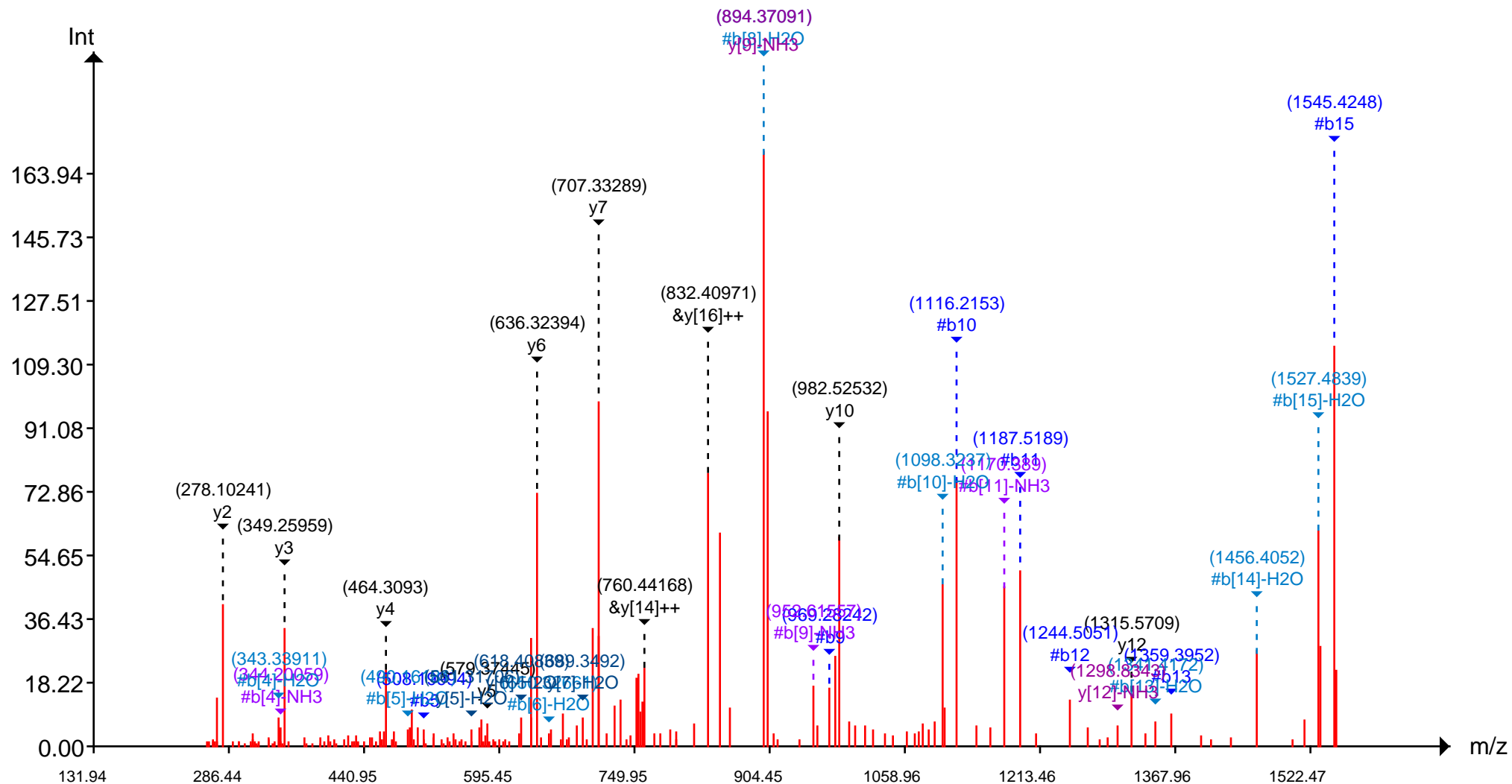
| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-NSSQDALGLSK<AcD3K*>LGGQAEK<AcD3K*>NR-COOH | Ion score | 109.15 |
| Protein | Q13136 | Confidence level | 99.00% |
| Precursor m/z | 1104.5822 | Identity Threshold | 48.86 |
| Charge | 2+ | Mass error | 6.48E-4 |



| Info | | | |
|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH | Ion score | 62.71 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 908.89746 | Identity Threshold | 43.04 |
| Charge | 2+ | Mass error | 0.001666 |

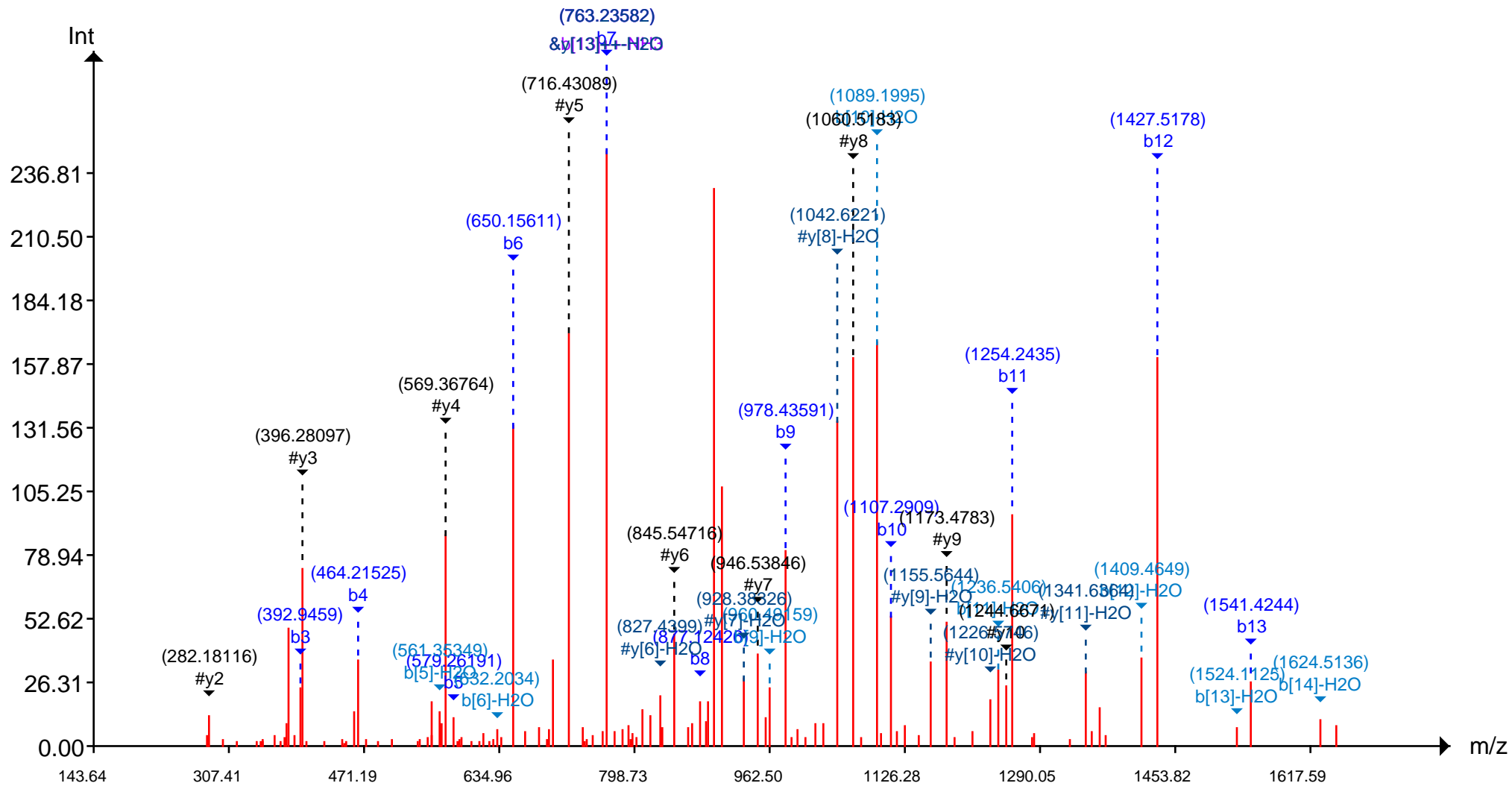


| Info | | | |
|--------------------------|--|---------------------------|-----------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 49.7 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 911.90576 | Identity Threshold | 44.13 |
| Charge | 2+ | Mass error | -0.001875 |



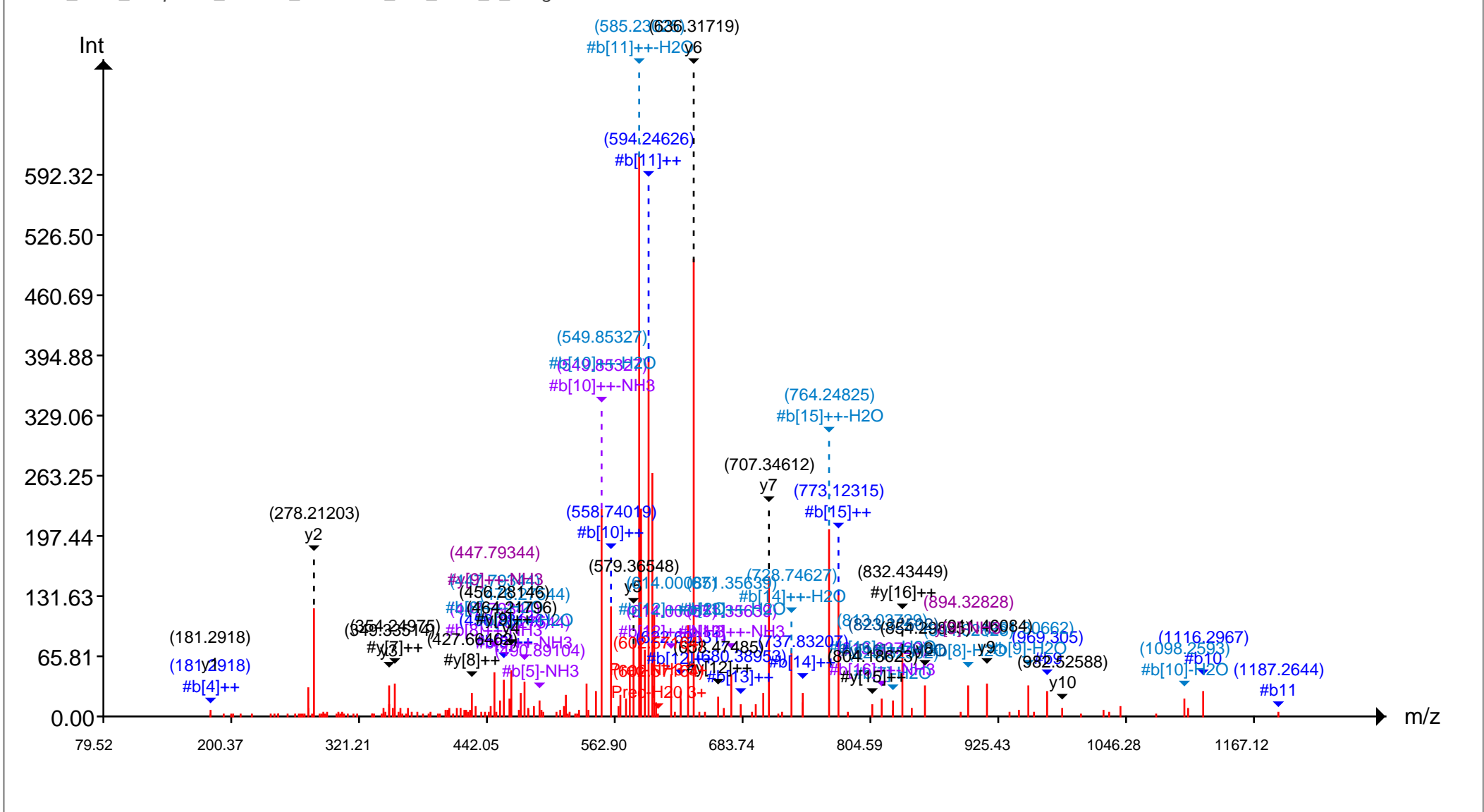
Info

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|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 46.01 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 911.90679 | Identity Threshold | 44.25 |
| Charge | 2+ | Mass error | 1.85E-4 |



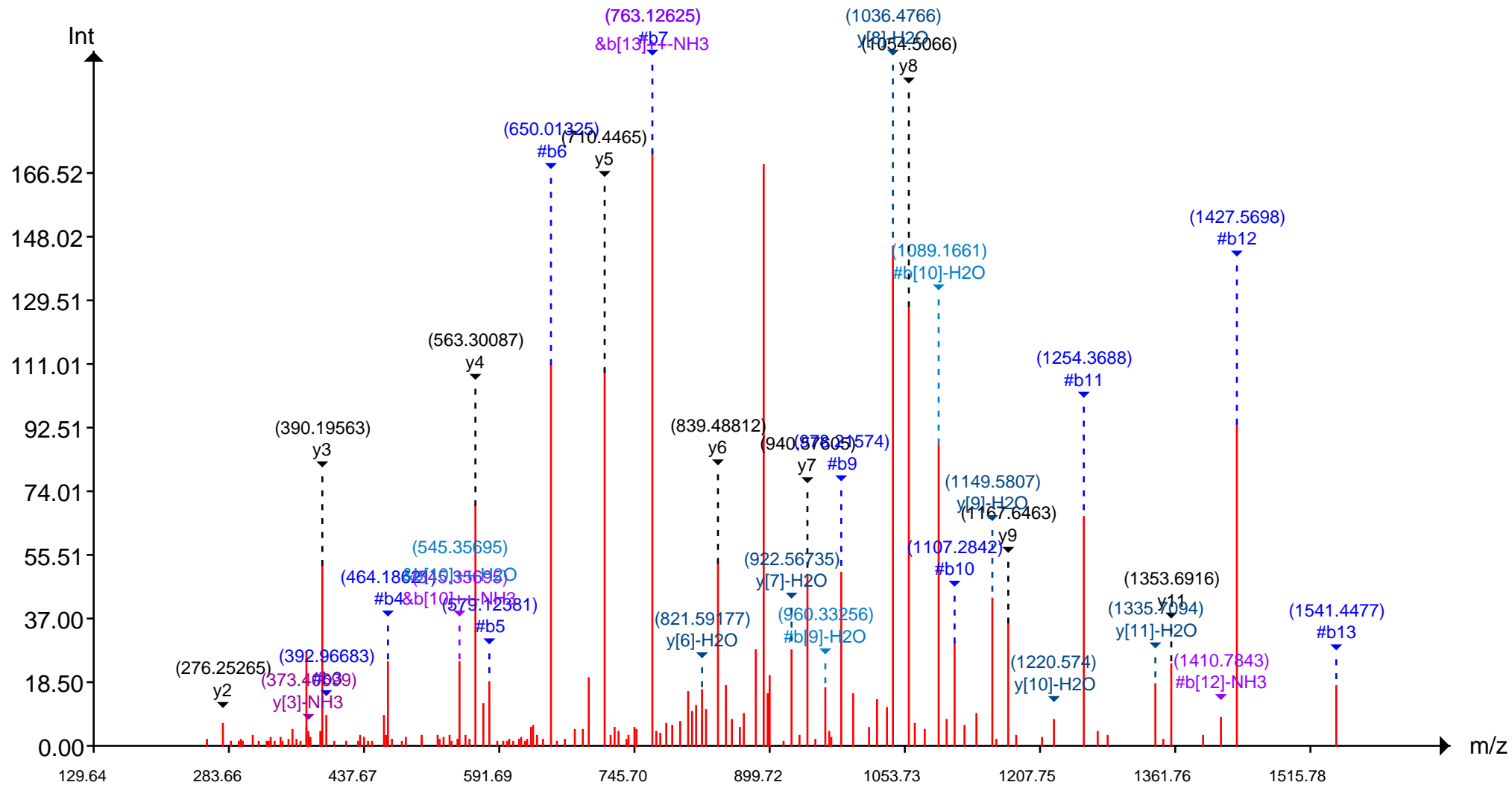
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-FSLADAINTEFK<AcD3K*>NTR<C13>-COOH | Ion score | 86.12 |
| Protein | P08670 | Confidence level | 99.00% |
| Precursor m/z | 911.98138 | Identity Threshold | 49.13 |
| Charge | 2+ | Mass error | 6.98E-4 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 75.95 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 608.27342 | Identity Threshold | 44.20 |
| Charge | 3+ | Mass error | -4.11E-4 |



Info

Modified Sequence AcD3-FSLADAINTEFK<AcD3K*>NTR-COOH

Ion score 88.45

Protein P08670

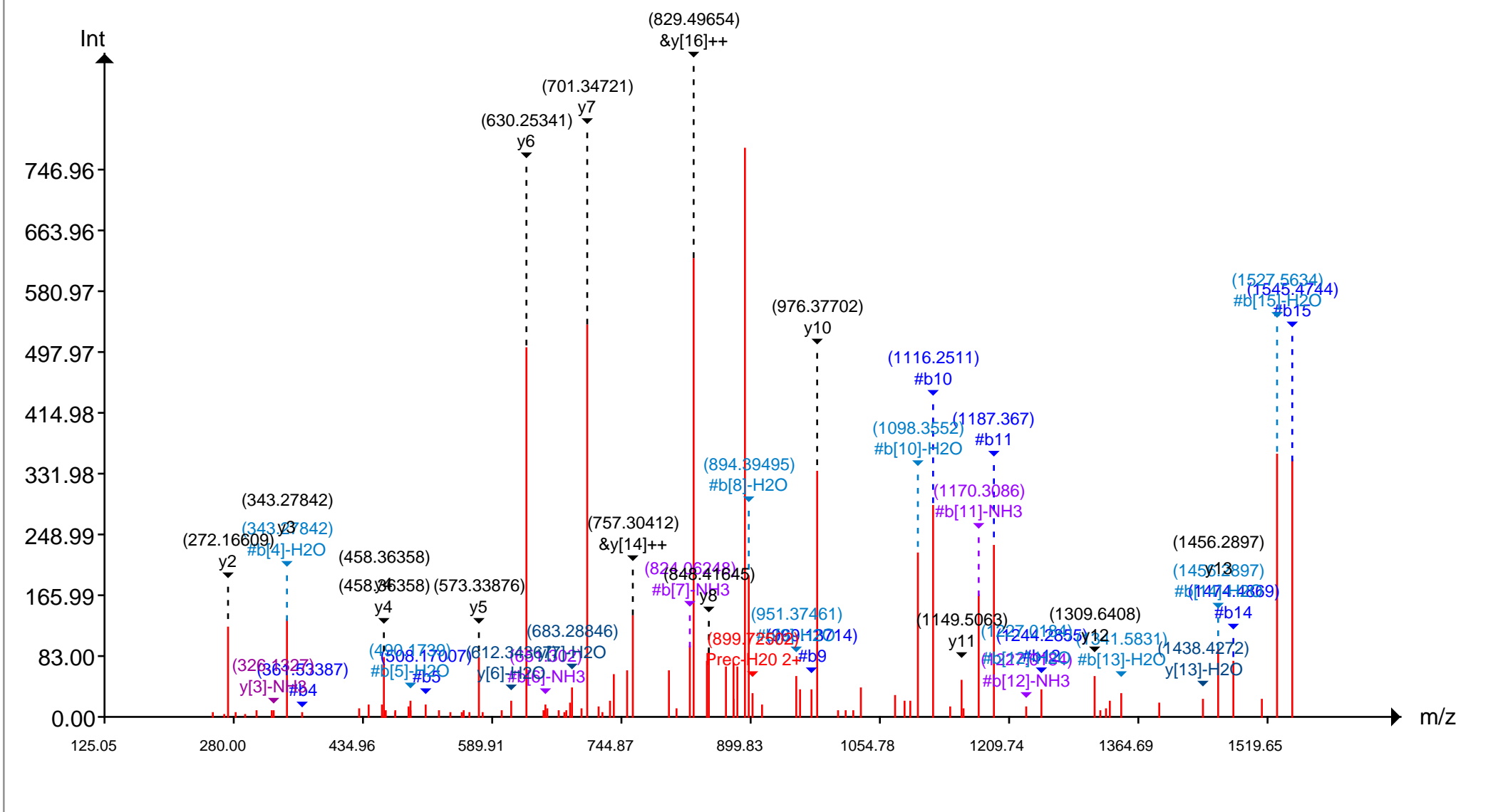
Confidence level 99.00%

Precursor m/z 908.97148

Identity Threshold 49.15

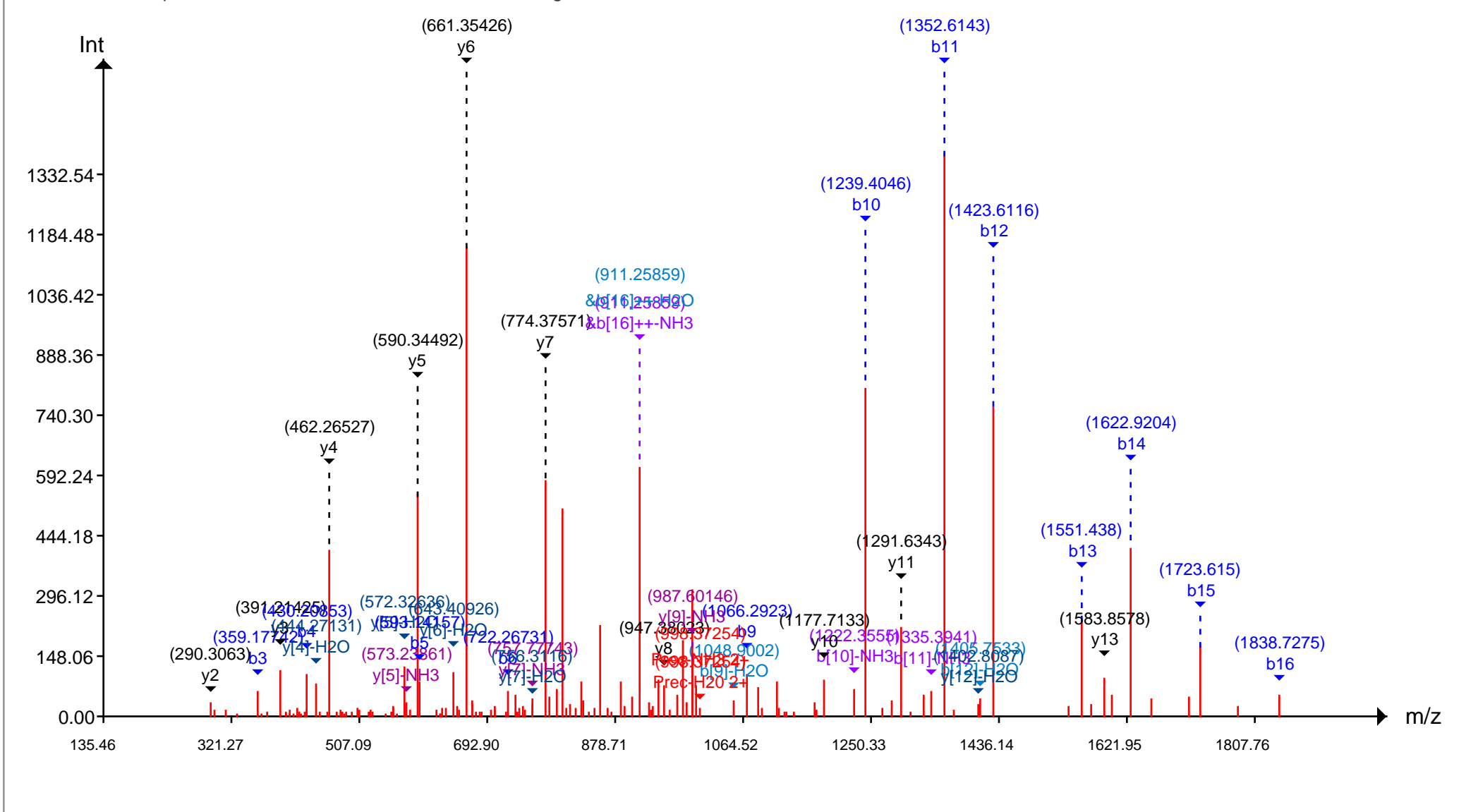
Charge 2+

Mass error 0.00104



Info

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|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH | Ion score | 86.12 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 908.89656 | Identity Threshold | 42.94 |
| Charge | 2+ | Mass error | -1.34E-4 |



Info

Modified Sequence AcD3-TVLAYENK<AcD3K*>GK<AcD3K*>IAQATDR-COOH

Ion score 89.01

Protein Q8N806

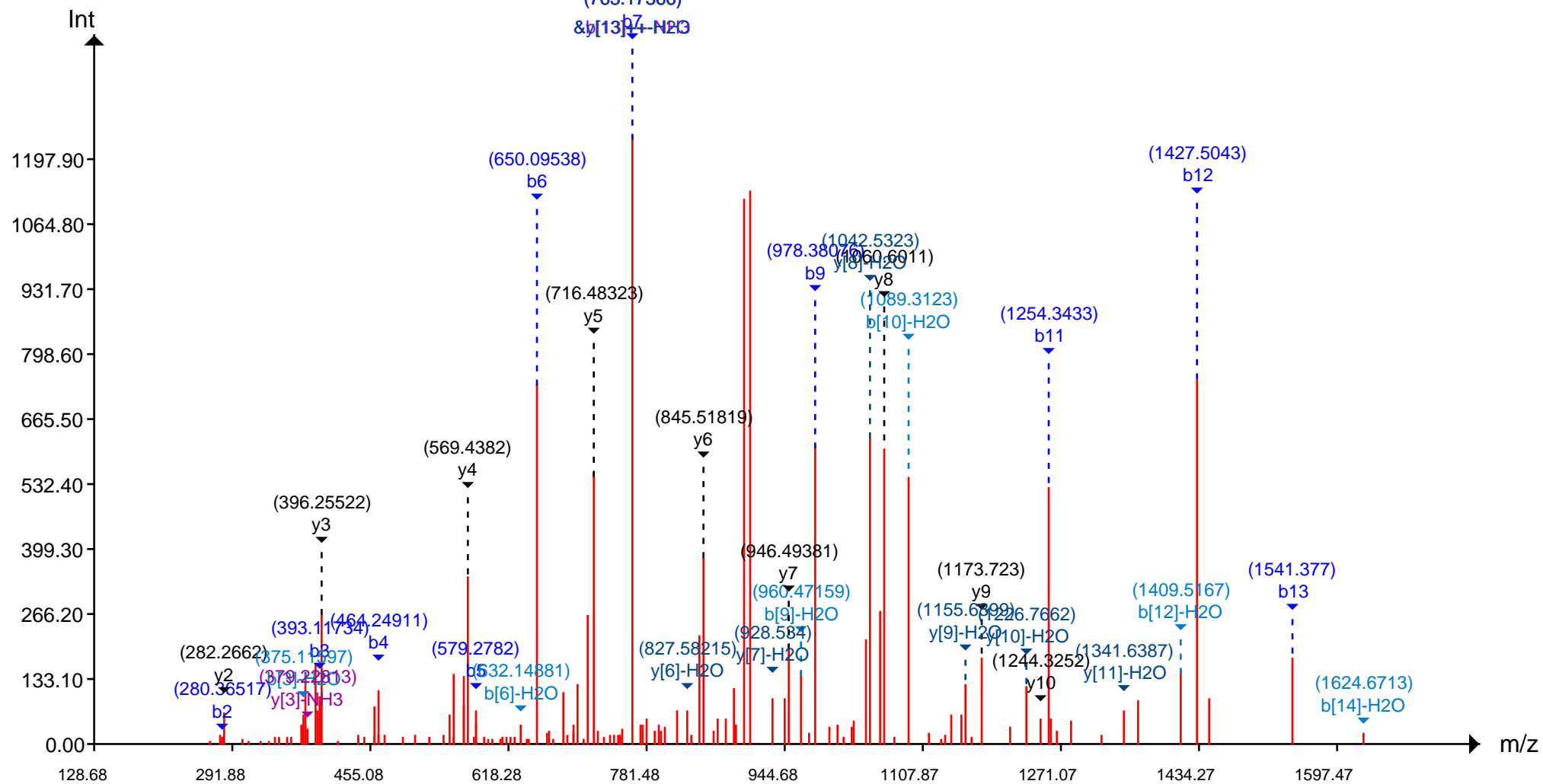
Confidence level 99.00%

Precursor m/z 1007.0522

Identity Threshold 48.19

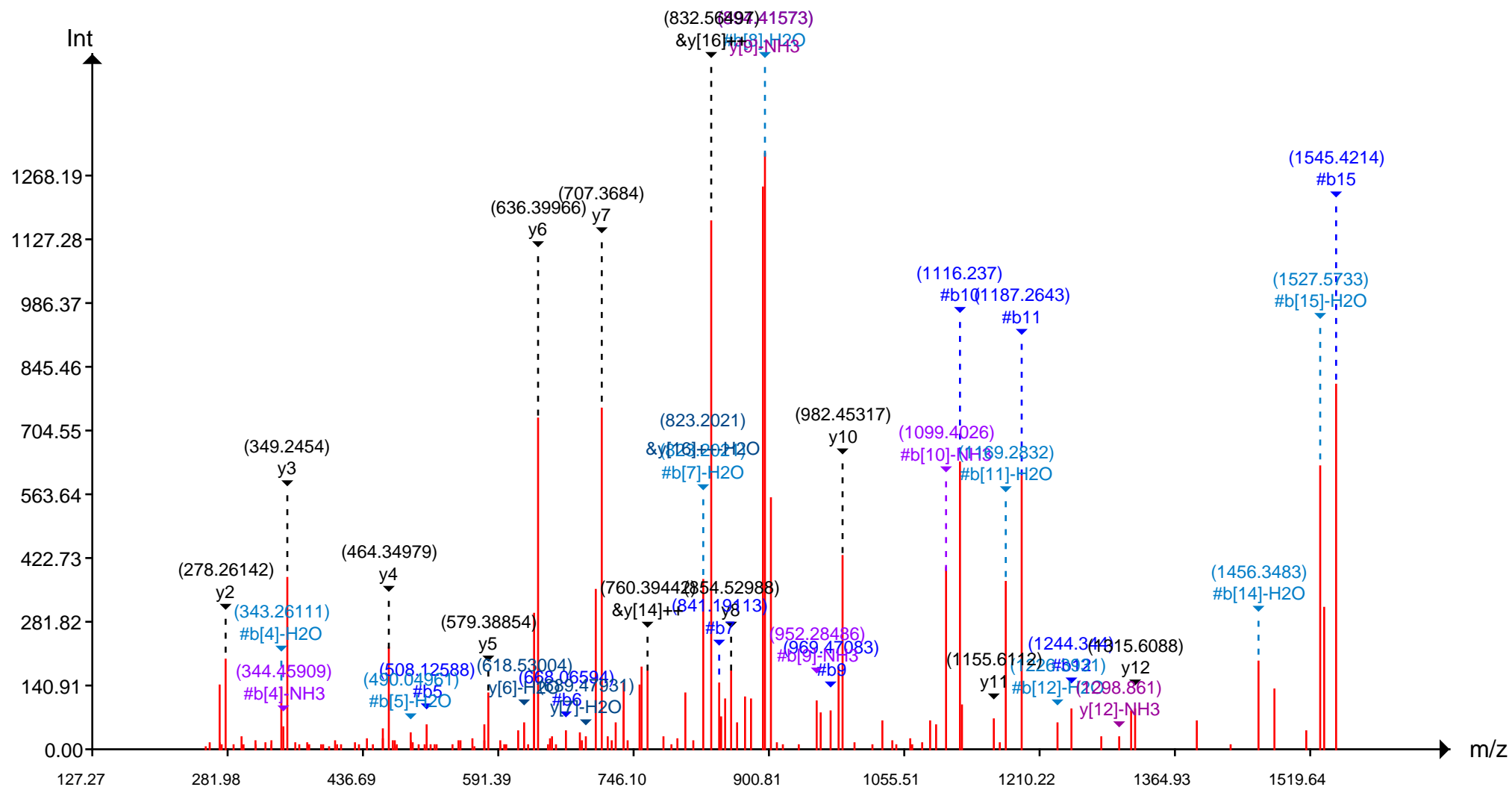
Charge 2+

Mass error 0.001011



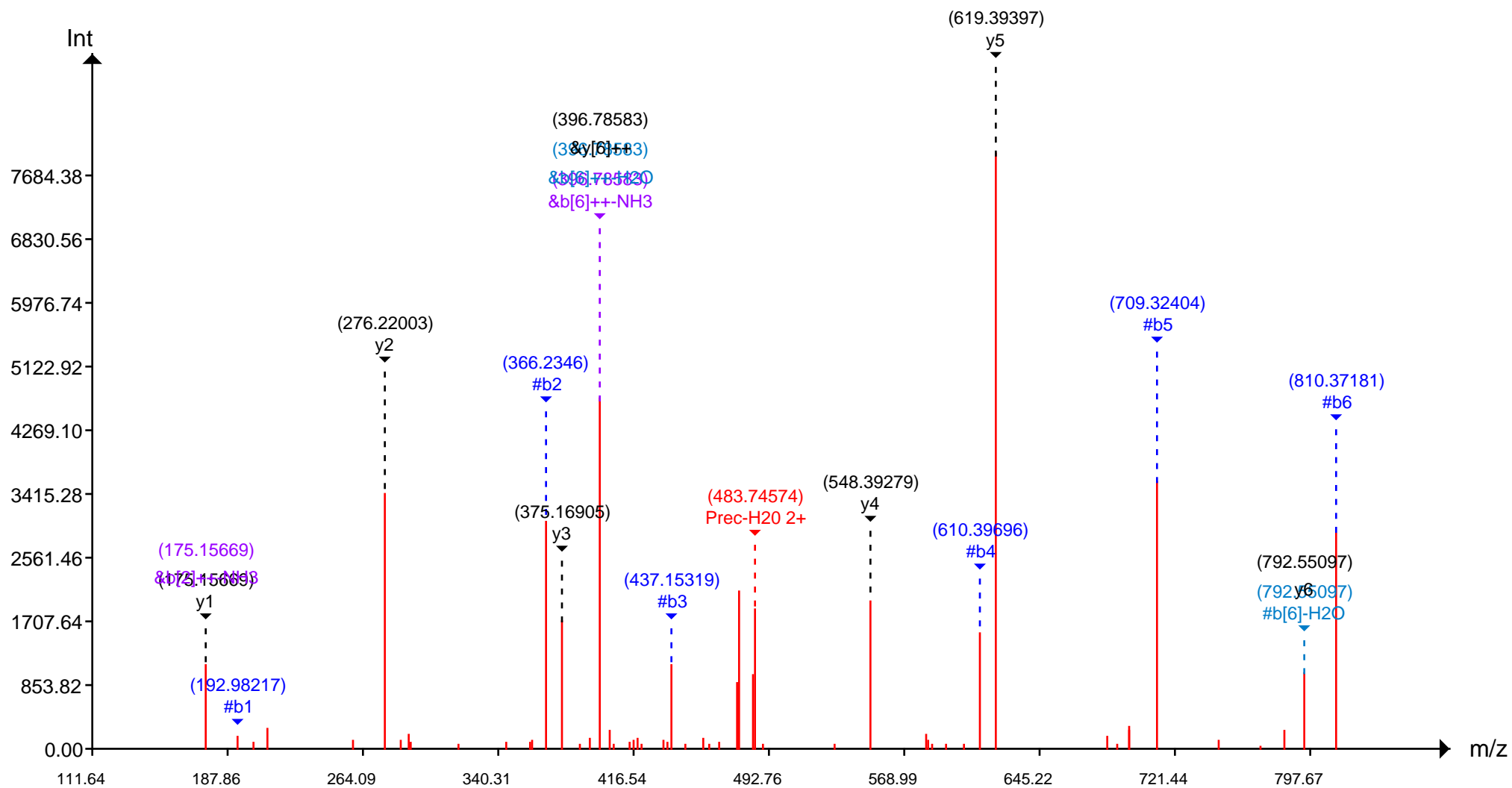
Info

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|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-FSLADAINTEFK<AcD3K*>NTR<C13>-COOH | Ion score | 64.52 |
| Protein | P08670 | Confidence level | 99.00% |
| Precursor m/z | 911.98148 | Identity Threshold | 49.11 |
| Charge | 2+ | Mass error | 8.98E-4 |



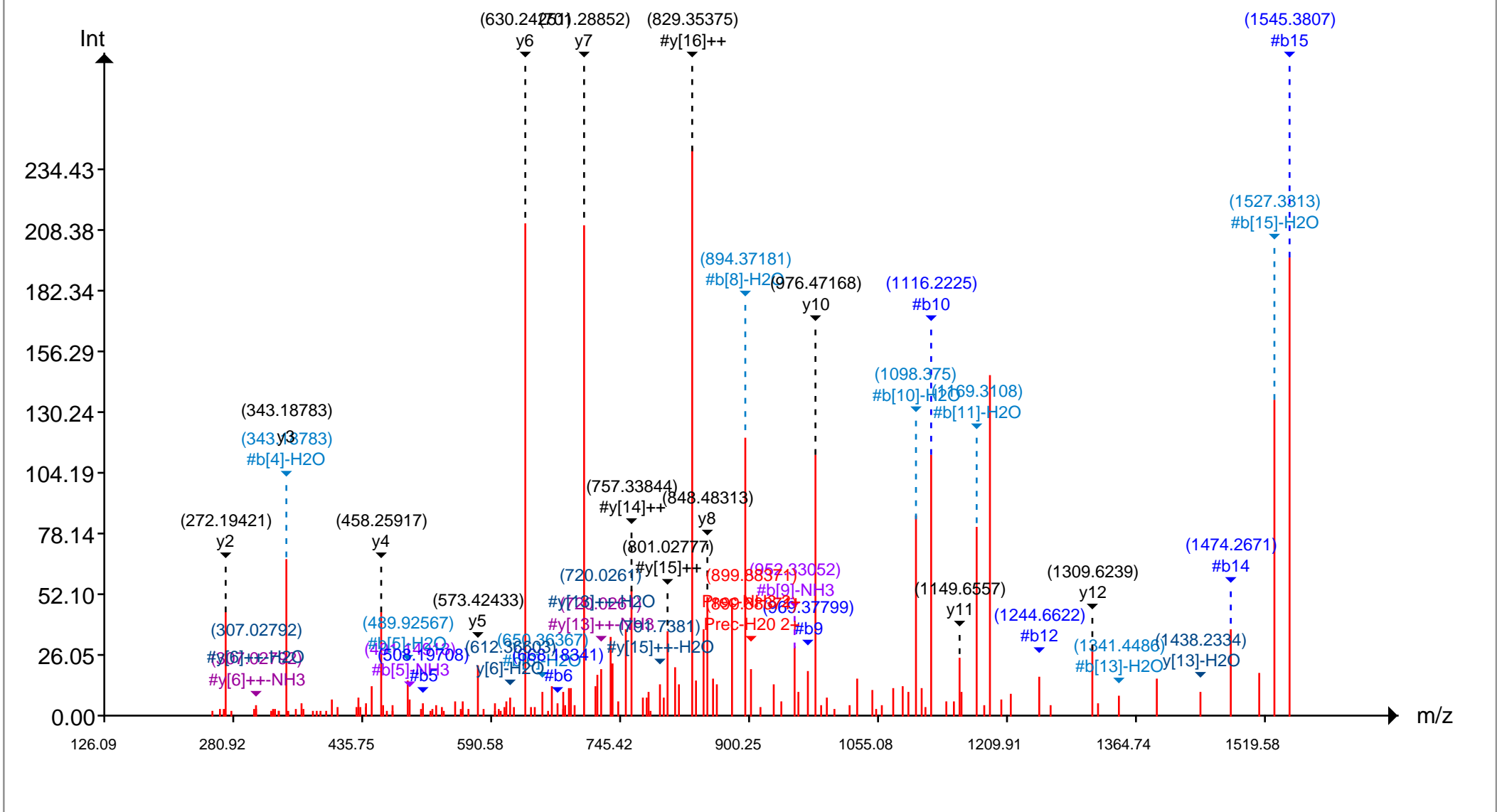
Info

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|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR<C13>-COOH | Ion score | 55.39 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 911.90643 | Identity Threshold | 44.20 |
| Charge | 2+ | Mass error | -5.35E-4 |



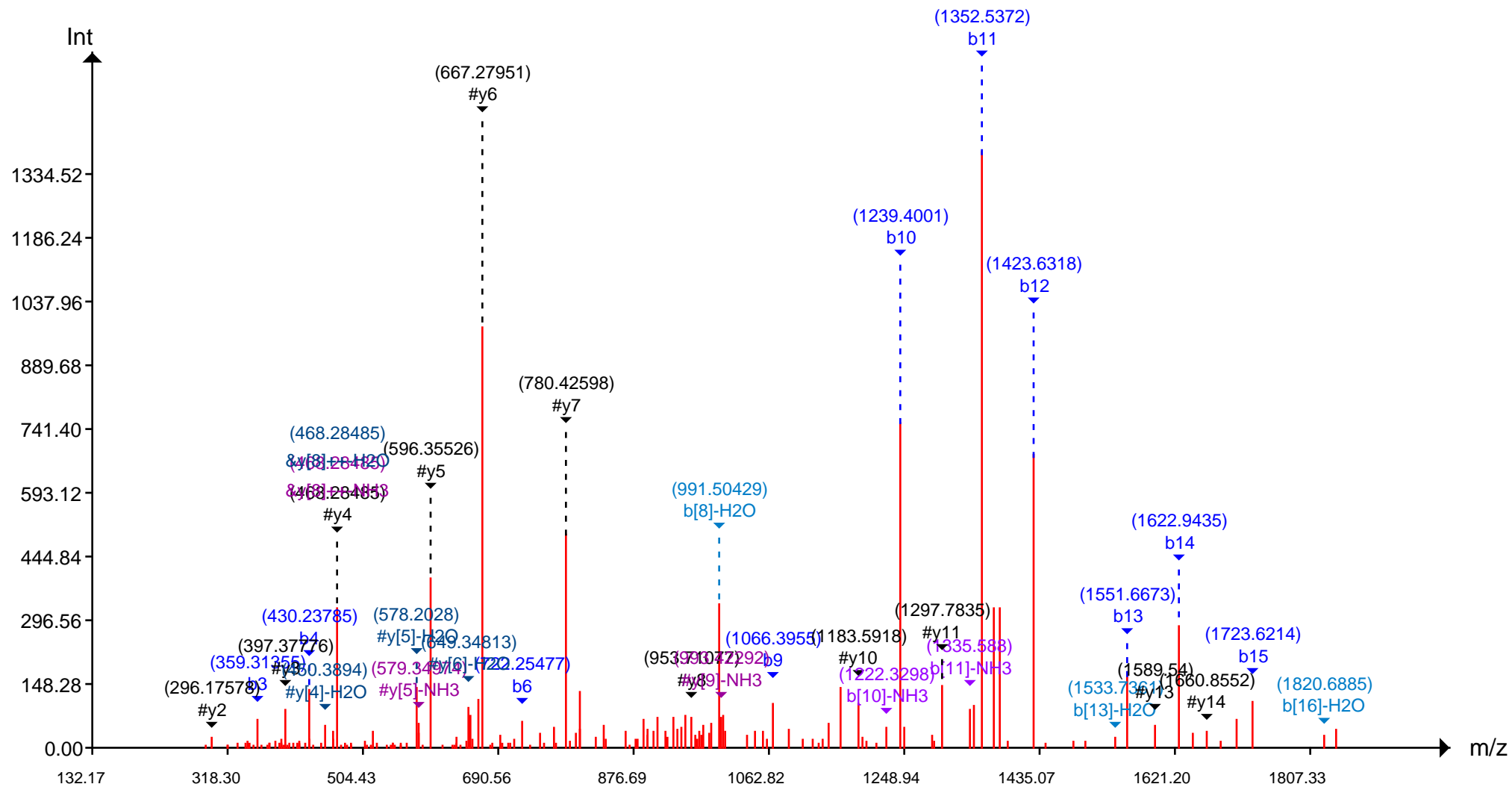
Info

| | | | |
|--------------------------|-----------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-FK<AcD3K*>AK<AcD3K*>VTR-COOH | Ion score | 45.15 |
| Protein | Q9Y4L1 | Confidence level | 99.00% |
| Precursor m/z | 492.81274 | Identity Threshold | 43.72 |
| Charge | 2+ | Mass error | -4.76E-4 |



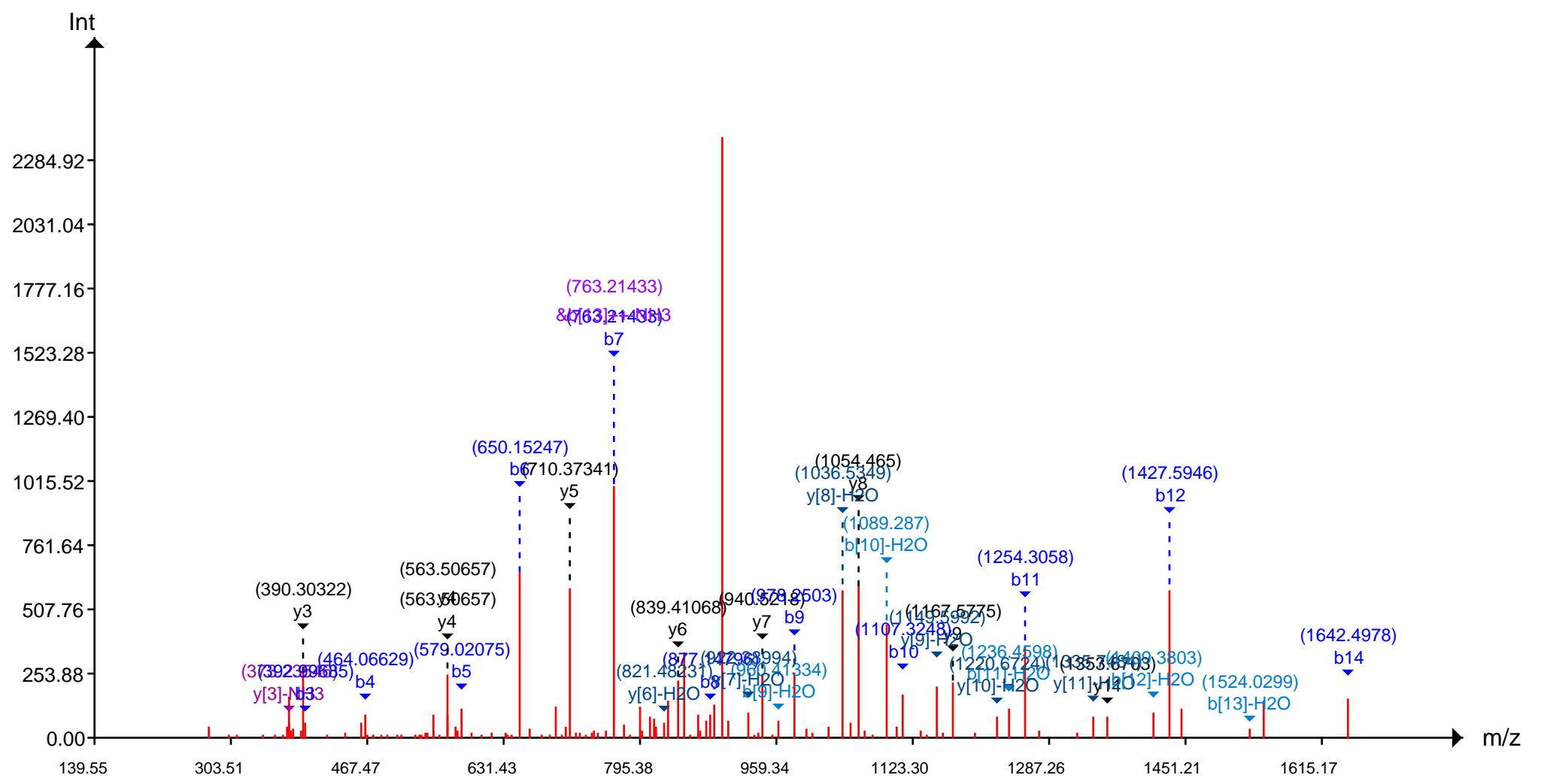
Info

| | | | |
|--------------------------|---|---------------------------|----------|
| Modified Sequence | AcD3-NGSGM<Mox*>C<Cmm*>K<AcD3K*>AGFAGDDAPR-COOH | Ion score | 75.62 |
| Protein | P60709, P63261, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 908.89647 | Identity Threshold | 42.90 |
| Charge | 2+ | Mass error | -3.14E-4 |

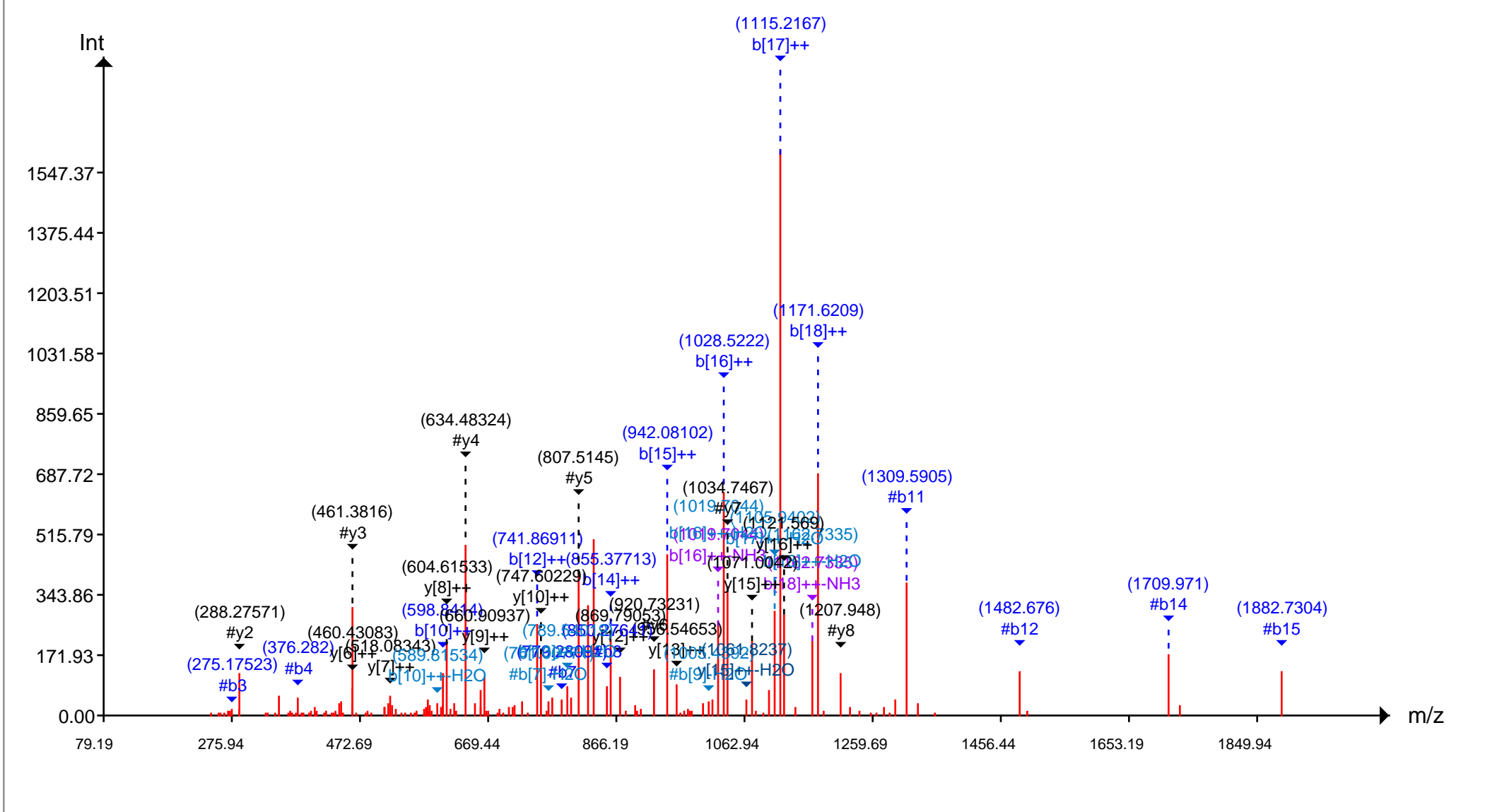


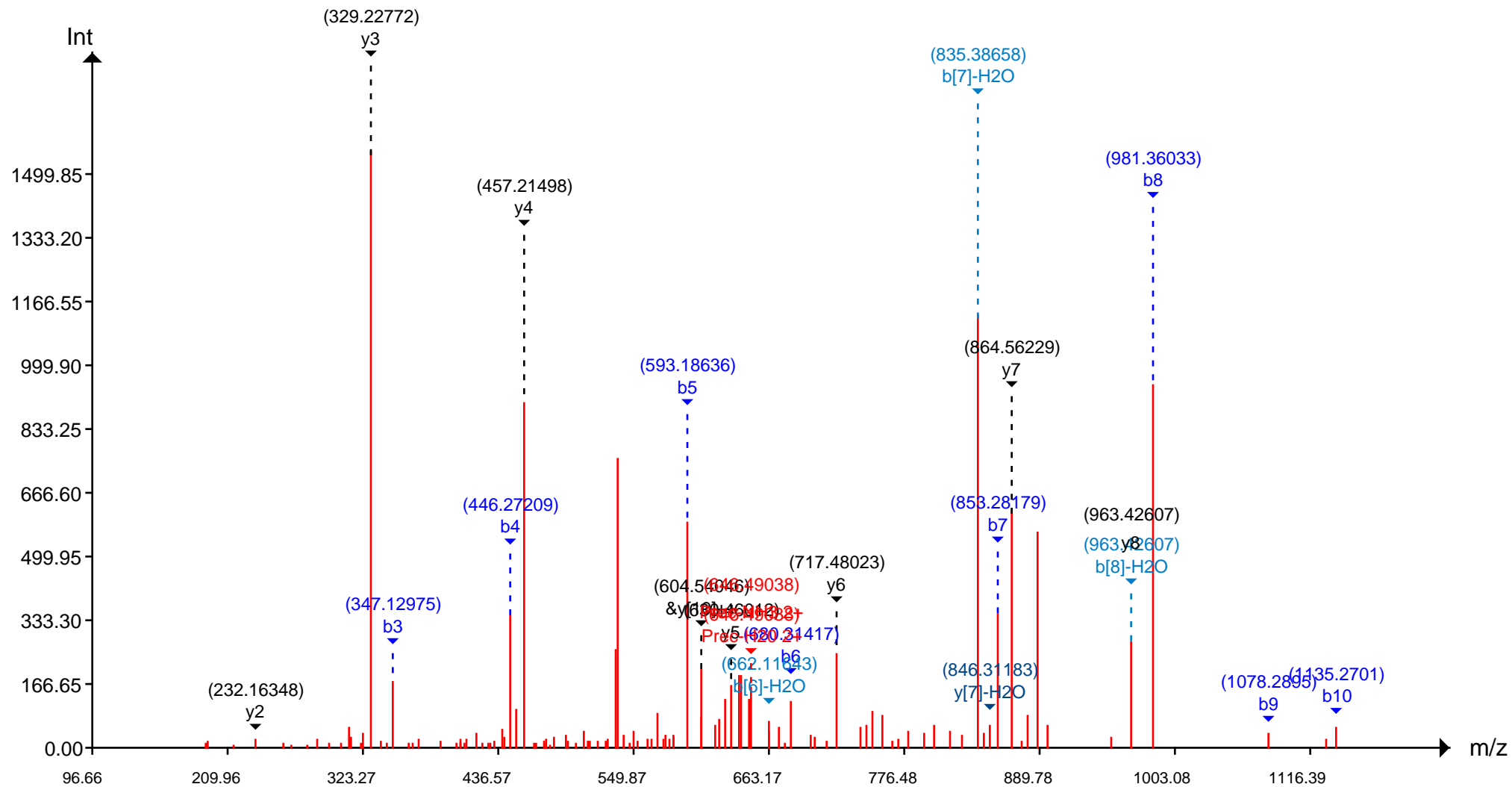
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-TVLAYENK<AcD3K*>GK<AcD3K*>IAQATDR<C13>-COOH | Ion score | 73.06 |
| Protein | Q8N806 | Confidence level | 99.00% |
| Precursor m/z | 1010.0623 | Identity Threshold | 47.58 |
| Charge | 2+ | Mass error | 0.00107 |



| Info | | | |
|--------------------------|-----------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-FSLADAINTEFK<AcD3K*>NTR-COOH | Ion score | 77.08 |
| Protein | P08670 | Confidence level | 99.00% |
| Precursor m/z | 909.48047 | Identity Threshold | 49.07 |
| Charge | 2+ | Mass error | 1.01902 |





Info

Modified Sequence AcD3-GAK<AcD3K*>VFSK<AcD3K*>QPGR-COOH

Protein P61011

Precursor m/z 655.38224

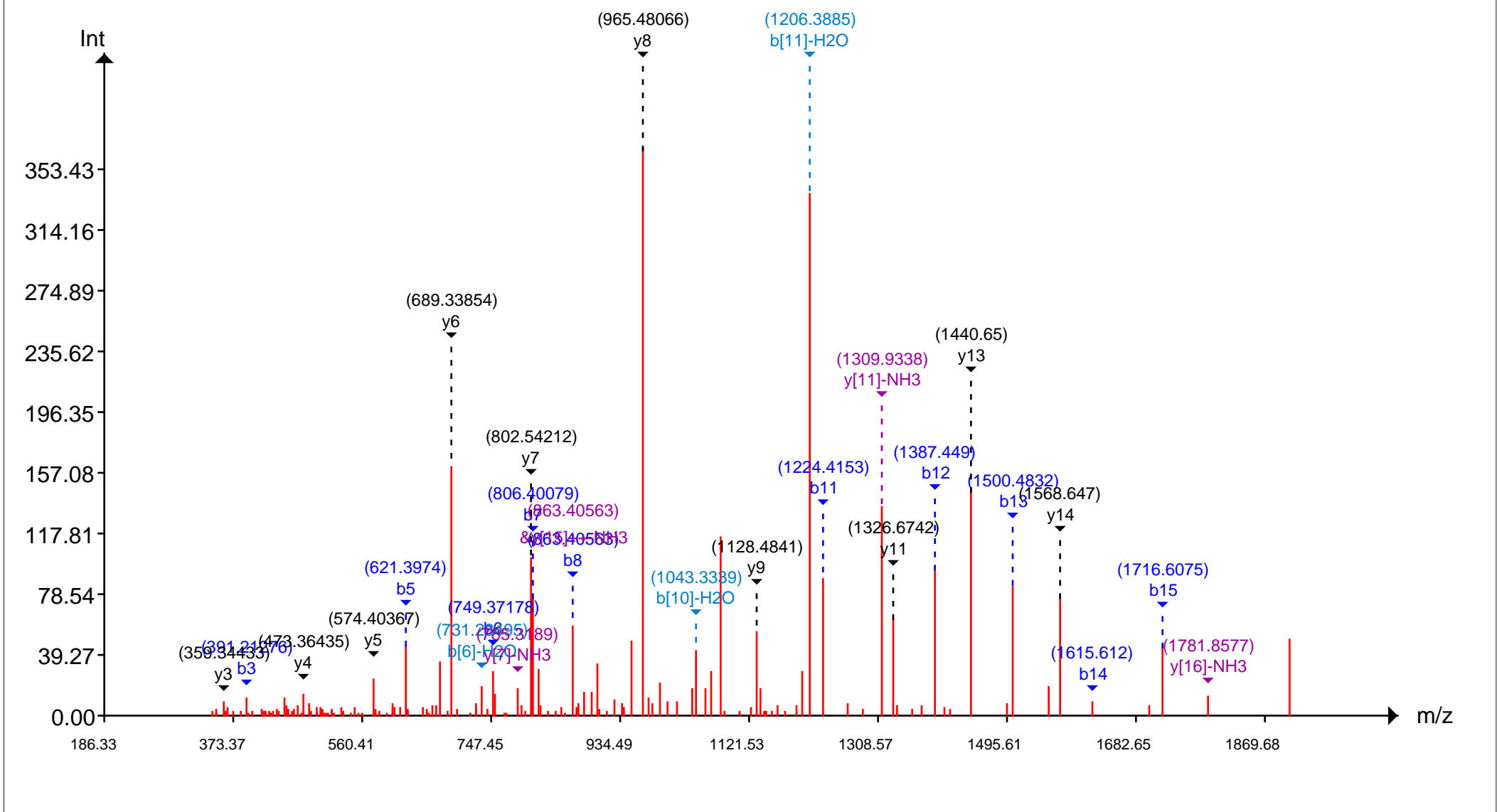
Charge 2+

Ion score 47.91

Confidence level 99.00%

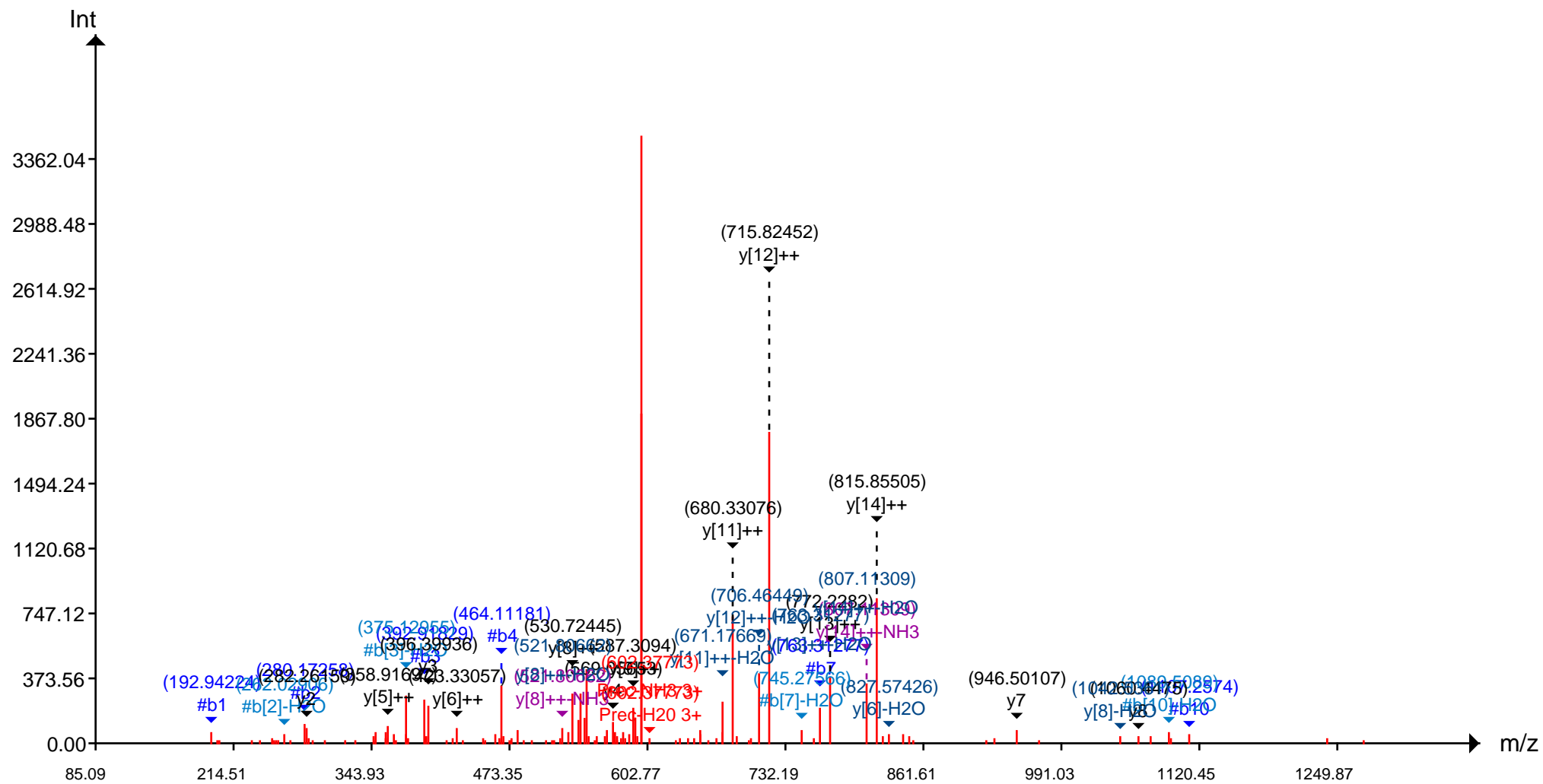
Identity Threshold 46.98

Mass error -1.03E-4

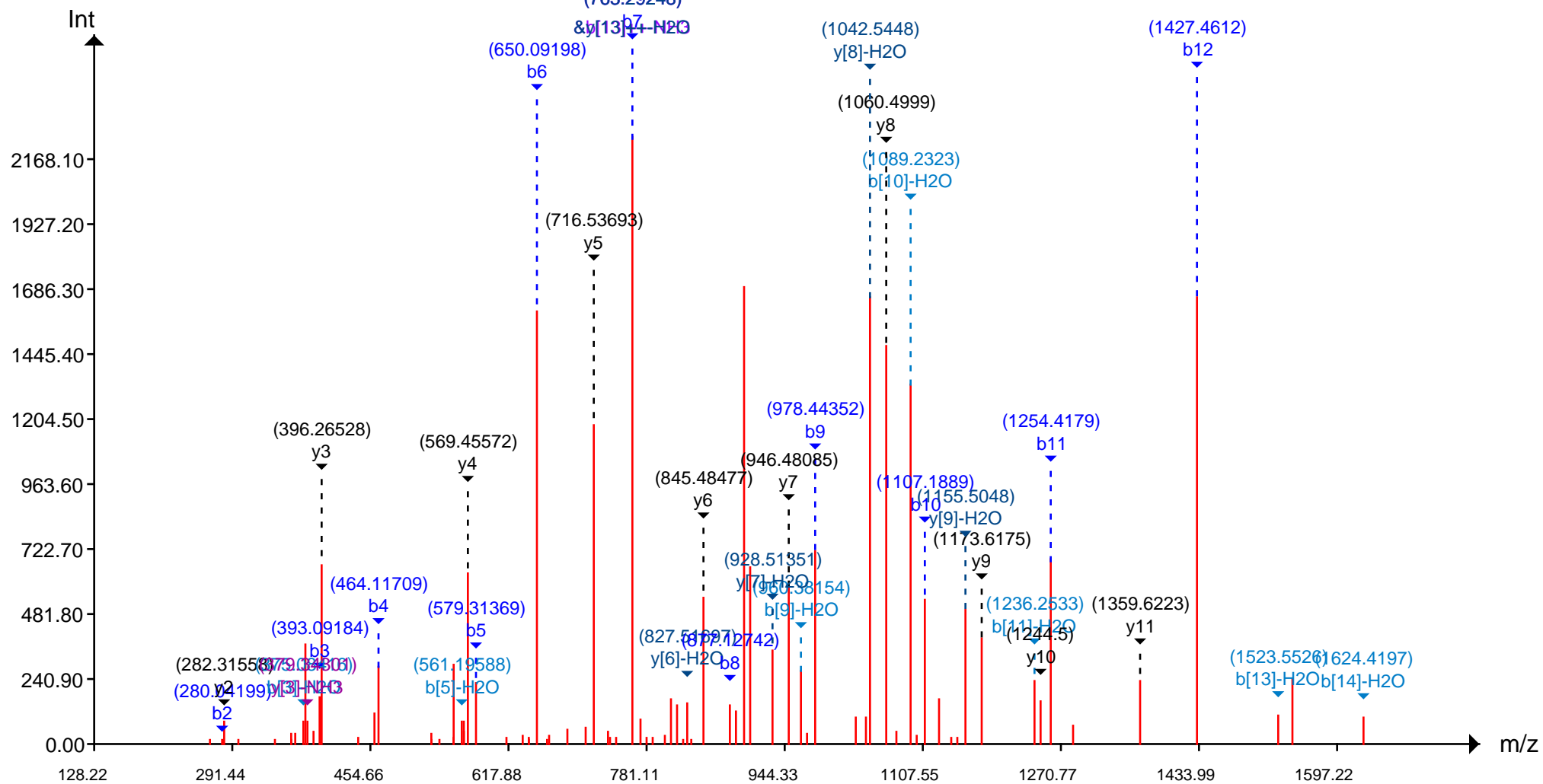


Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GDK<AcD3K*>GK<AcD3K*>QGGPTYIYIDTNALR-COOH | Ion score | 87.61 |
| Protein | O96019 | Confidence level | 99.00% |
| Precursor m/z | 1095.0638 | Identity Threshold | 49.06 |
| Charge | 2+ | Mass error | 0.001995 |

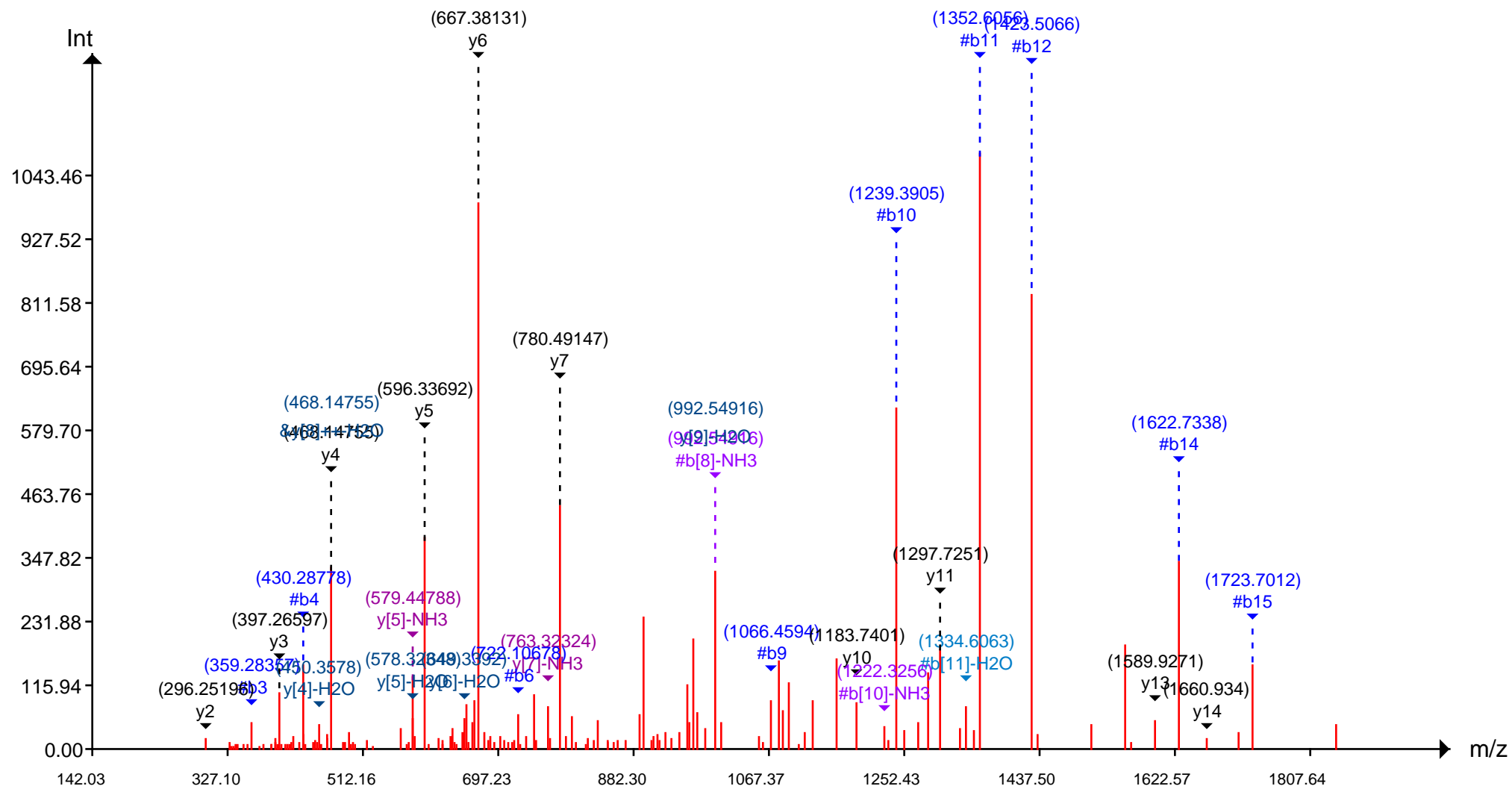


| Info | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-FSLADAINTEFK<AcD3K*>NTR<C13>-COOH | Ion score | 51.1 |
| Protein | P08670 | Confidence level | 99.00% |
| Precursor m/z | 608.32307 | Identity Threshold | 49.11 |
| Charge | 3+ | Mass error | -1.28E-4 |



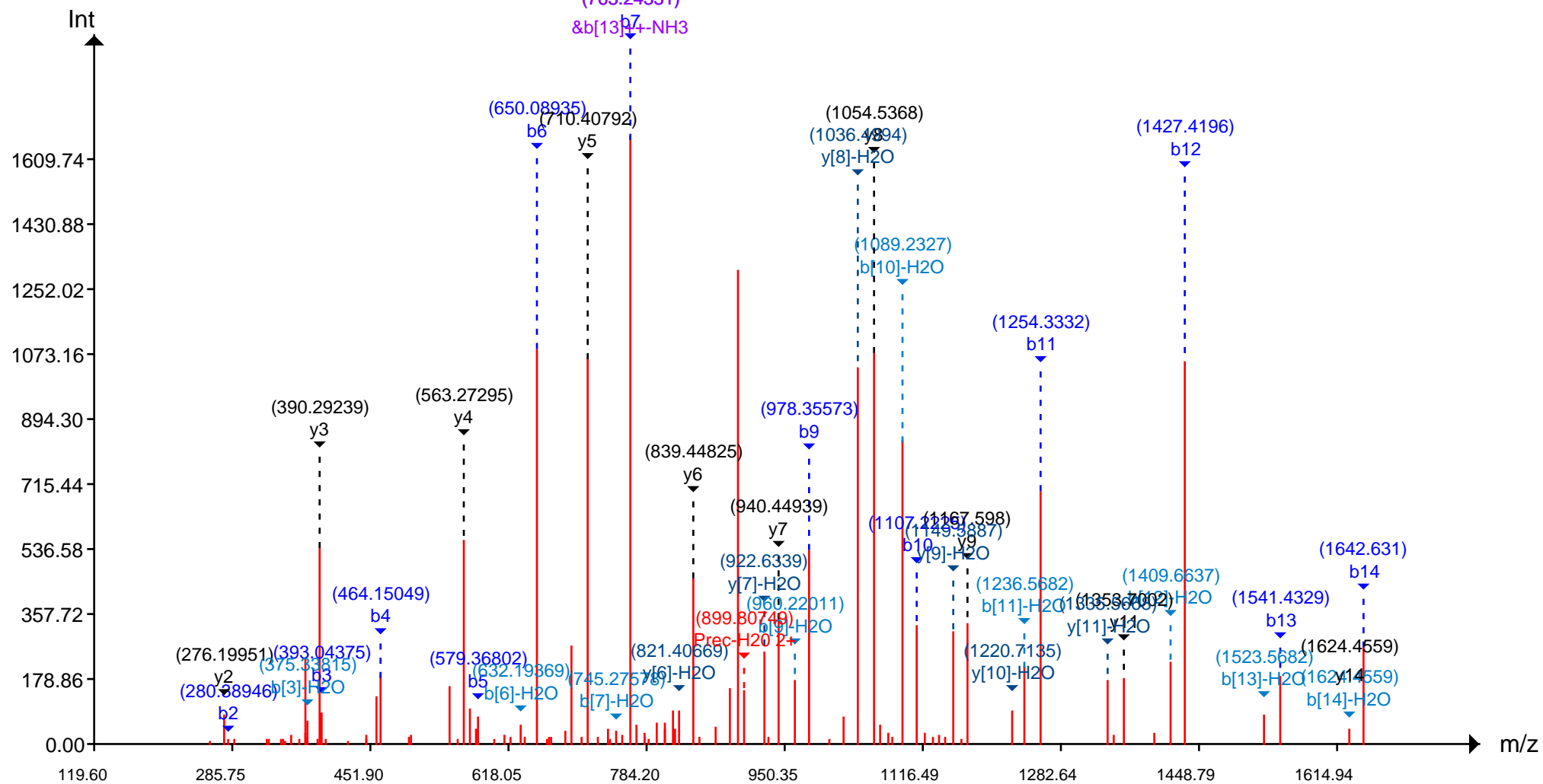
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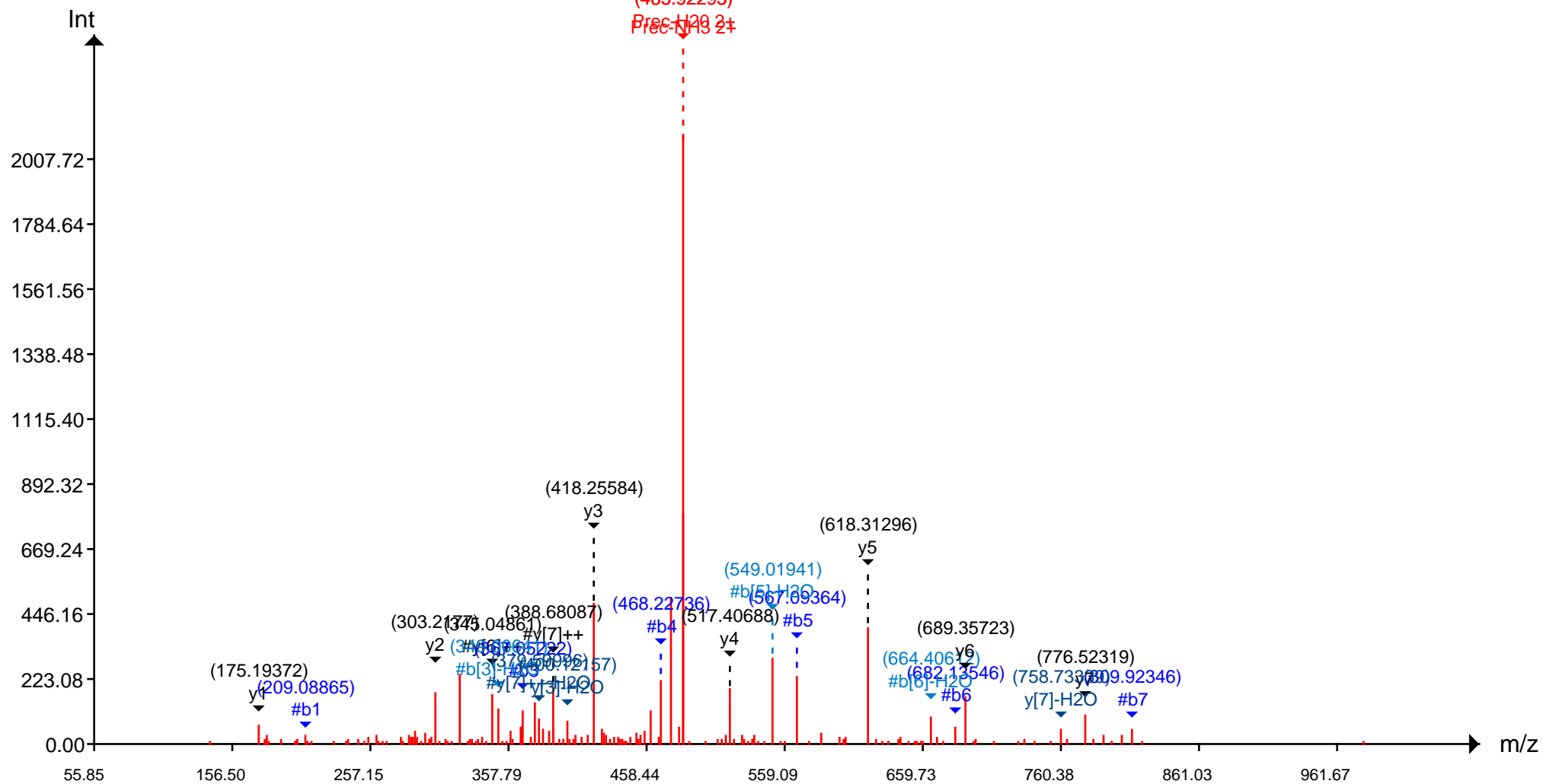
| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-FSLADAINTEFK<AcD3K*>NTR<C13>-COOH | Ion score | 87.58 |
| Protein | P08670 | Confidence level | 99.00% |
| Precursor m/z | 911.98139 | Identity Threshold | 49.13 |
| Charge | 2+ | Mass error | 7.18E-4 |



Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-TVLAYENK<AcD3K*>GK<AcD3K*>IAQATDR<C13>-COOH | Ion score | 56.05 |
| Protein | Q8N806 | Confidence level | 99.00% |
| Precursor m/z | 1010.0613 | Identity Threshold | 47.67 |
| Charge | 2+ | Mass error | -9.3E-4 |





Info

Modified Sequence AcD3-YSATVDQR-COOH

Protein O00232

Precursor m/z 492.74477

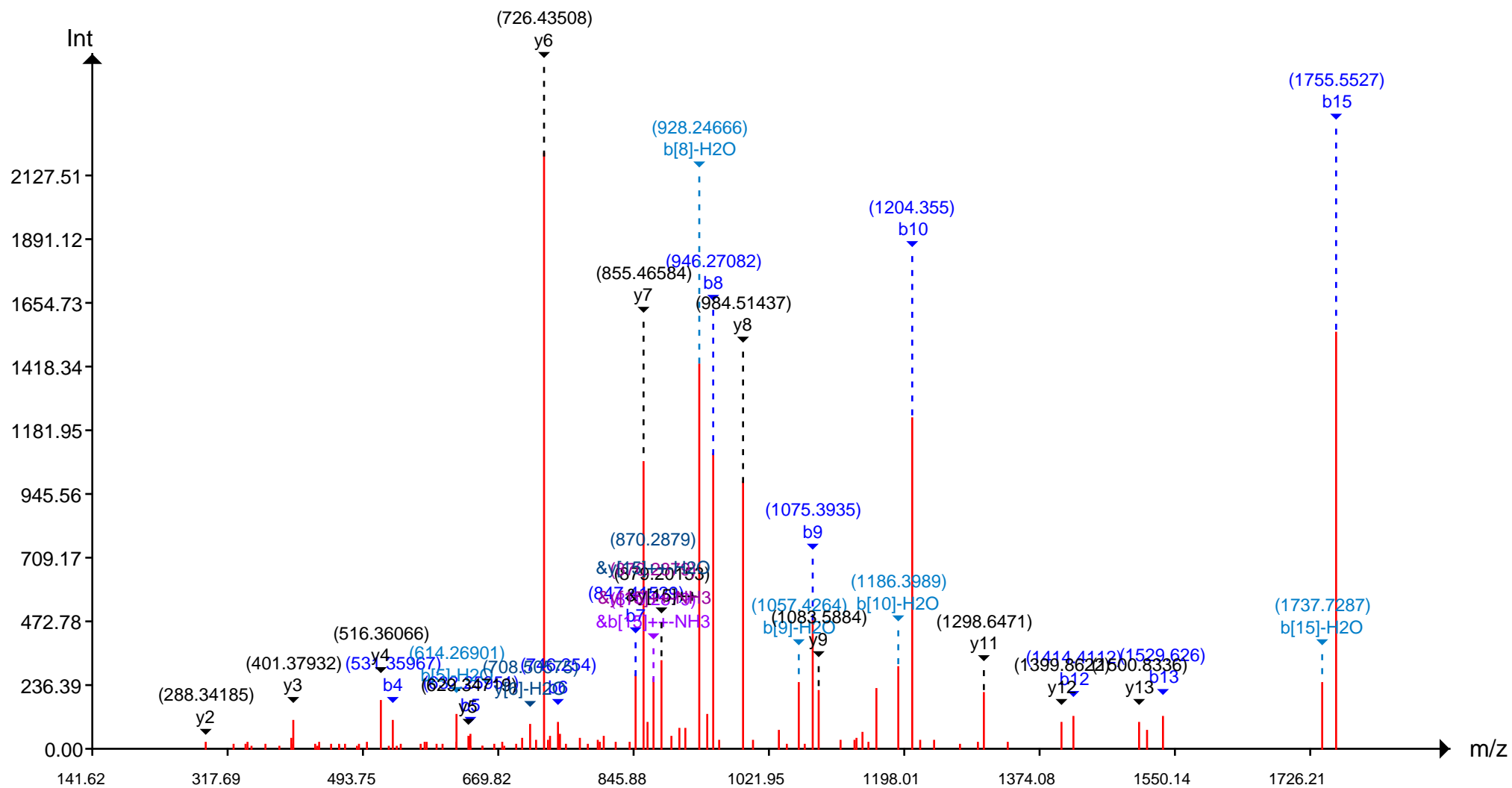
Charge 2+

Ion score 48.2

Confidence level 99.00%

Identity Threshold 47.91

Mass error -1.55E-4



Info

Modified Sequence AcD3-QQTTNTVEEPLDLIR-COOH

Ion score 86.48

Protein P62310

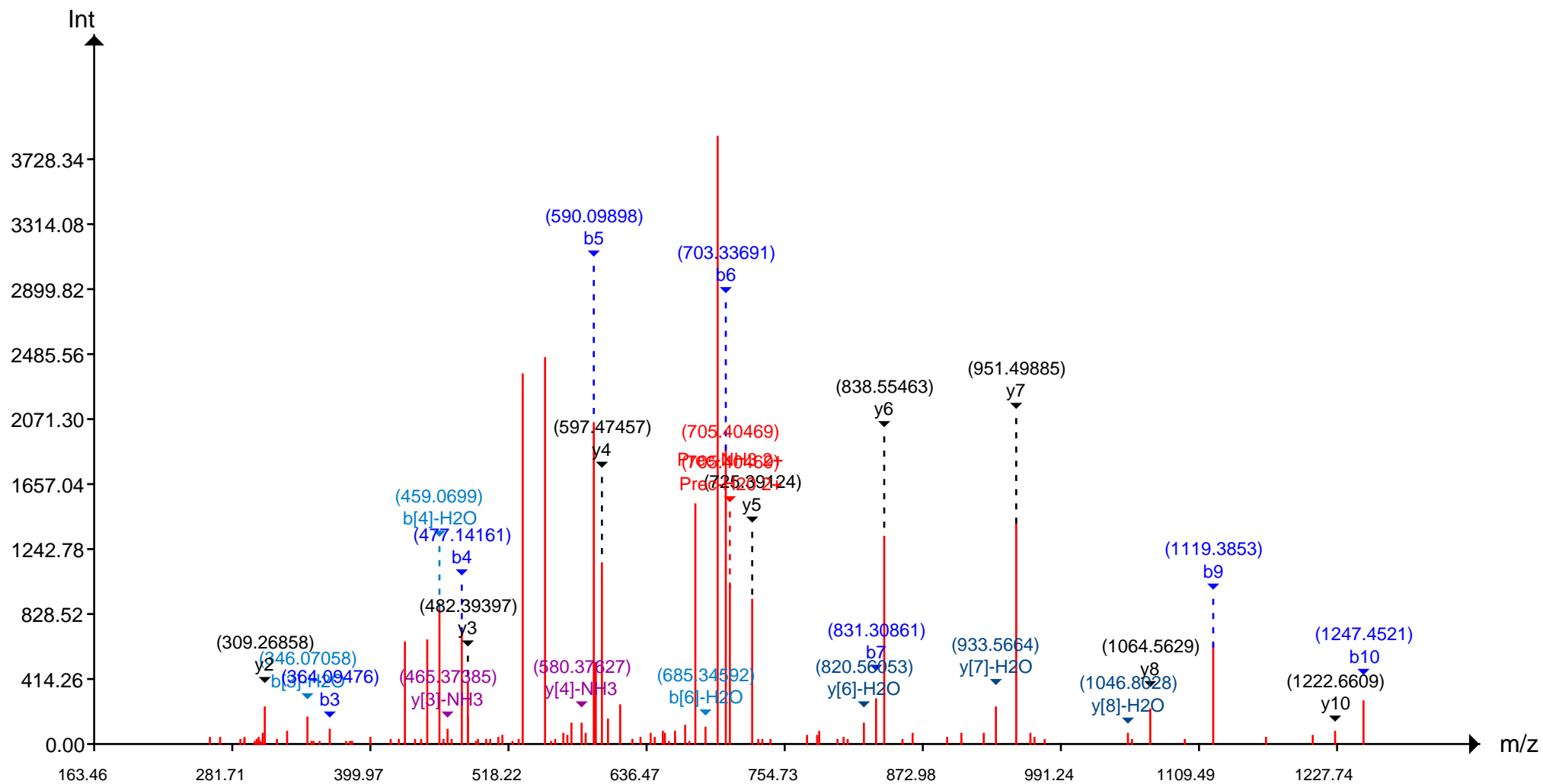
Confidence level 99.00%

Precursor m/z 965.50197

Identity Threshold 49.02

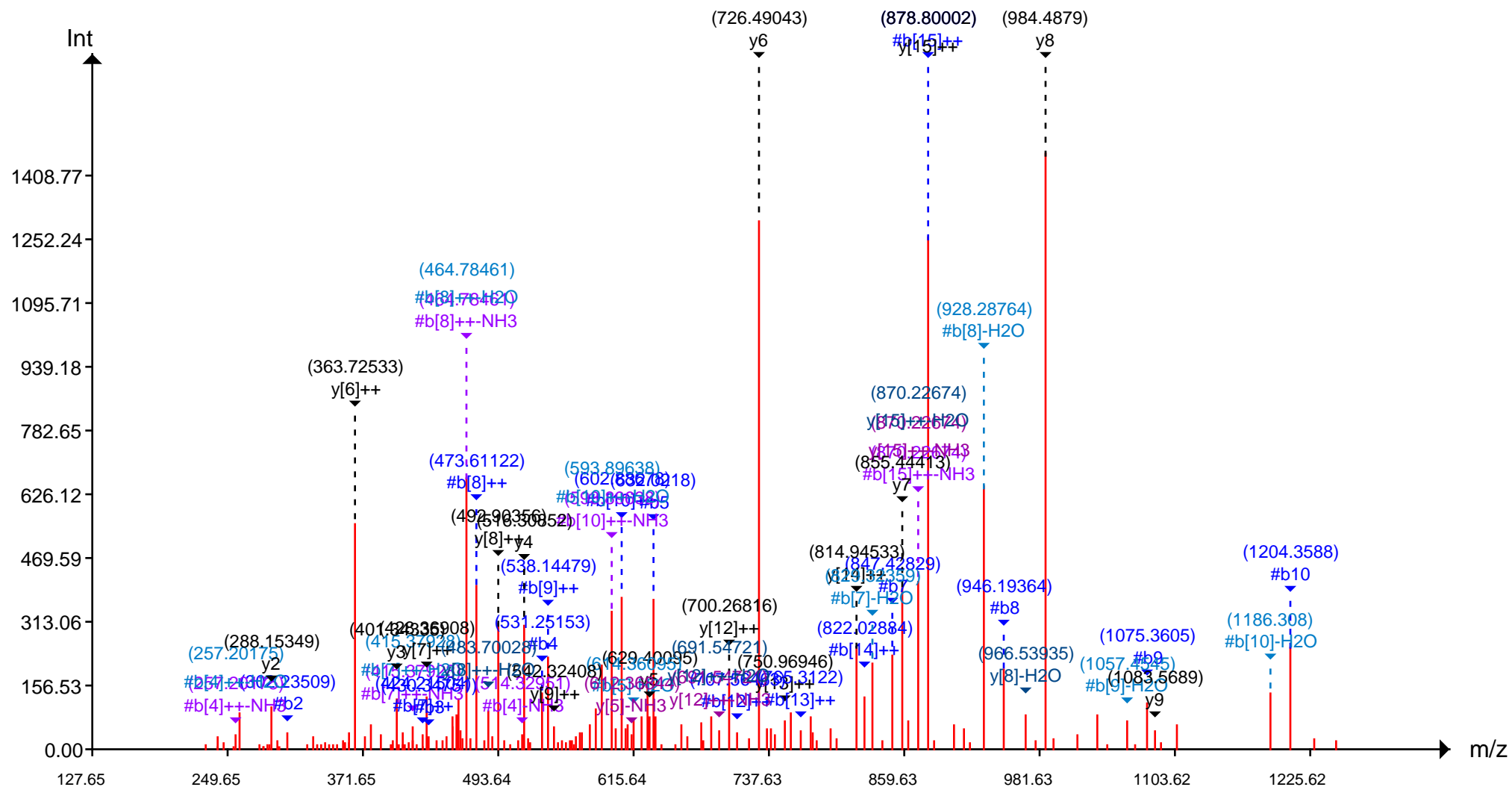
Charge 2+

Mass error 0.001137



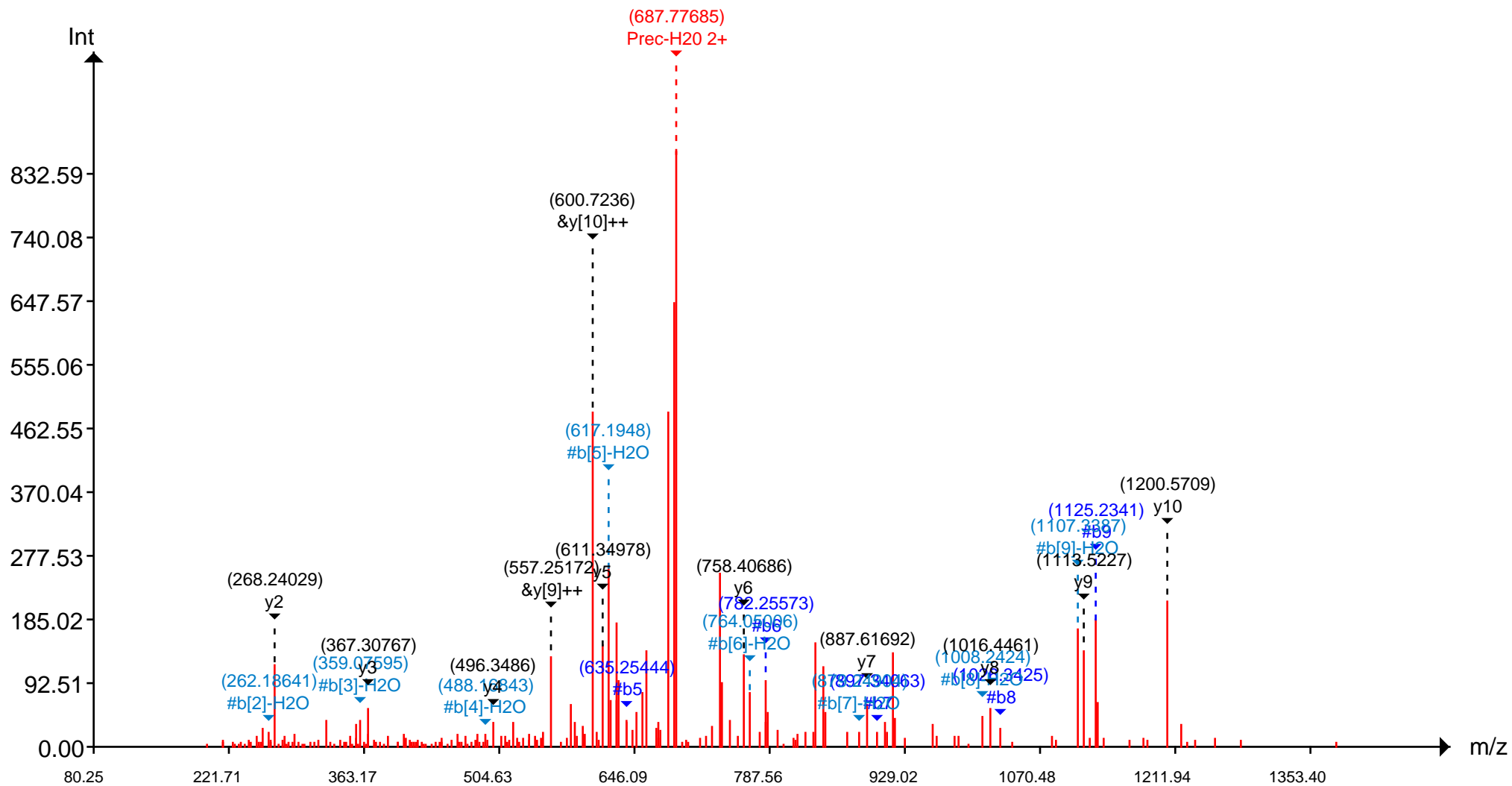
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-C<Cmm*>GTILLQDK<AcD3K*>QR<C13>-COOH | Ion score | 48.37 |
| Protein | O60232 | Confidence level | 99.00% |
| Precursor m/z | 714.39816 | Identity Threshold | 48.25 |
| Charge | 2+ | Mass error | 1.21E-4 |



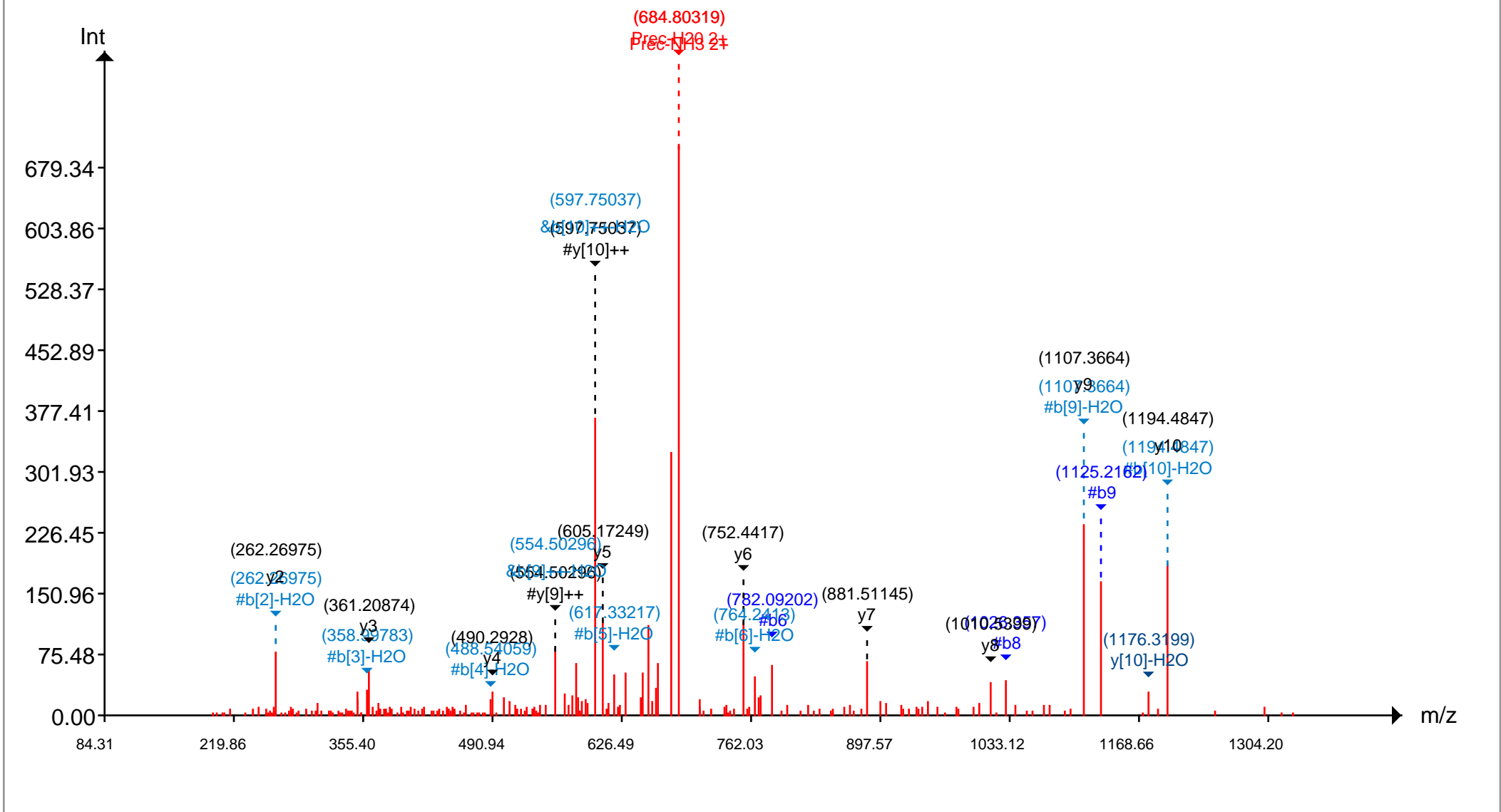
Info

| | | | |
|--------------------------|---------------------------|---------------------------|--------|
| Modified Sequence | AcD3-QQTTNTVEEPLDLIR-COOH | Ion score | 56.87 |
| Protein | P62310 | Confidence level | 99.00% |
| Precursor m/z | 644.00337 | Identity Threshold | 49.00 |
| Charge | 3+ | Mass error | 3.1E-5 |



Info

| | | | |
|--------------------------|----------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-M<Mox*>SPEEFDEVSR<C13>-COOH | Ion score | 44.96 |
| Protein | P42224 | Confidence level | 99.00% |
| Precursor m/z | 696.81012 | Identity Threshold | 44.45 |
| Charge | 2+ | Mass error | 7.23E-4 |



Info

Modified Sequence AcD3-M<Mox*>SPEEFDEVSR-COOH

Ion score 54.75

Protein P42224

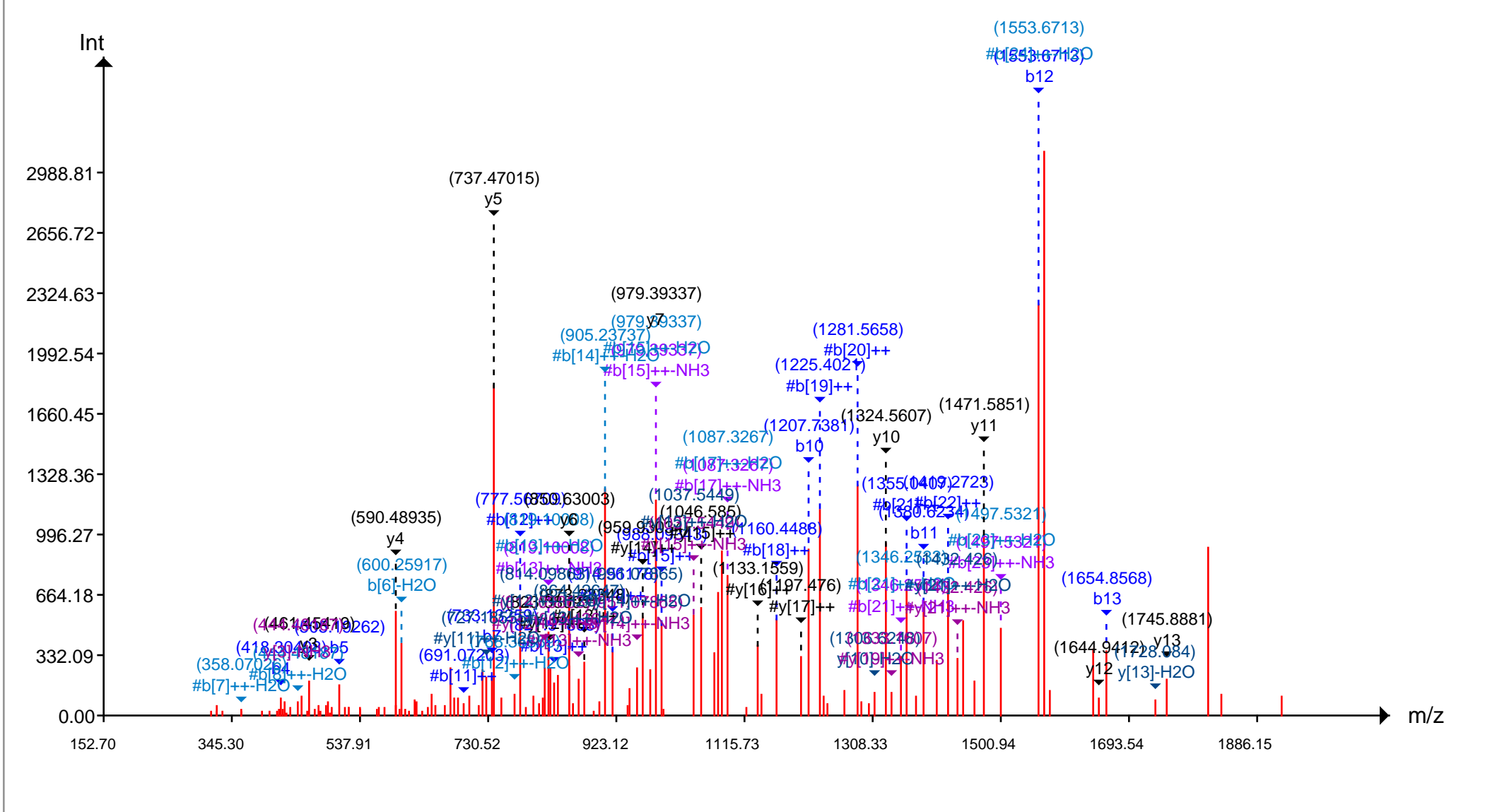
Confidence level 99.00%

Precursor m/z 693.79997

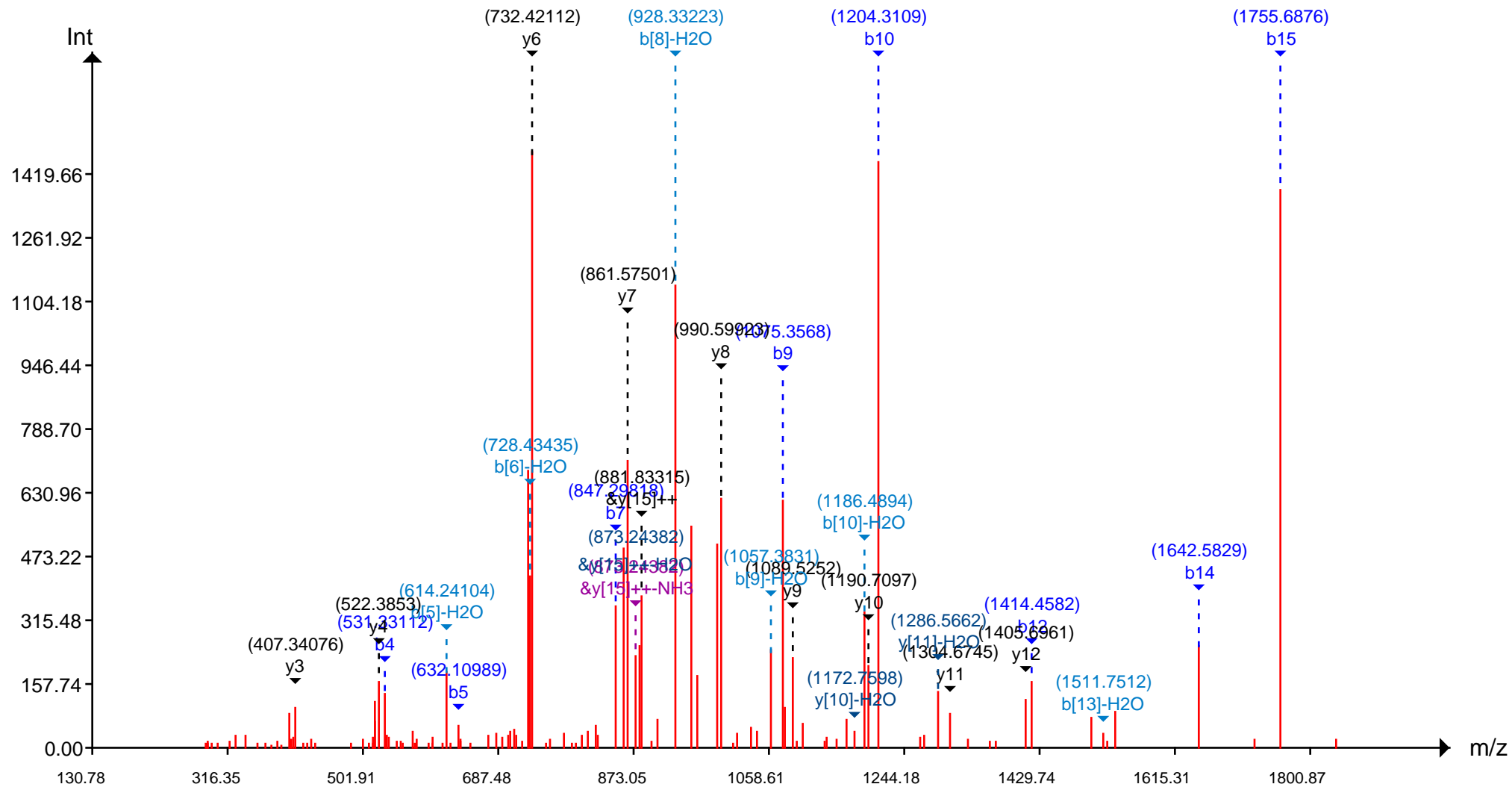
Identity Threshold 42.73

Charge 2+

Mass error 5.64E-4



| Info | Ion score | Confidence |
|---|-----------|------------|
| AcD3-GAAK<AcD3K*>SLDK<AcD3K*>QK<AcD3K*>K<AcD3K*>K<AcD3K*>TK<AcD3K*>M<Mox*>TDEEIM<Mox*>EK<AcD3K*>LR-COOH | 1100.2738 | High |
| Q13177 | | Identified |
| 3+ | | Mass error |



Info

Modified Sequence AcD3-QQTTNTVEEPLDLIR<C13>-COOH

Ion score 57.09

Protein P62310

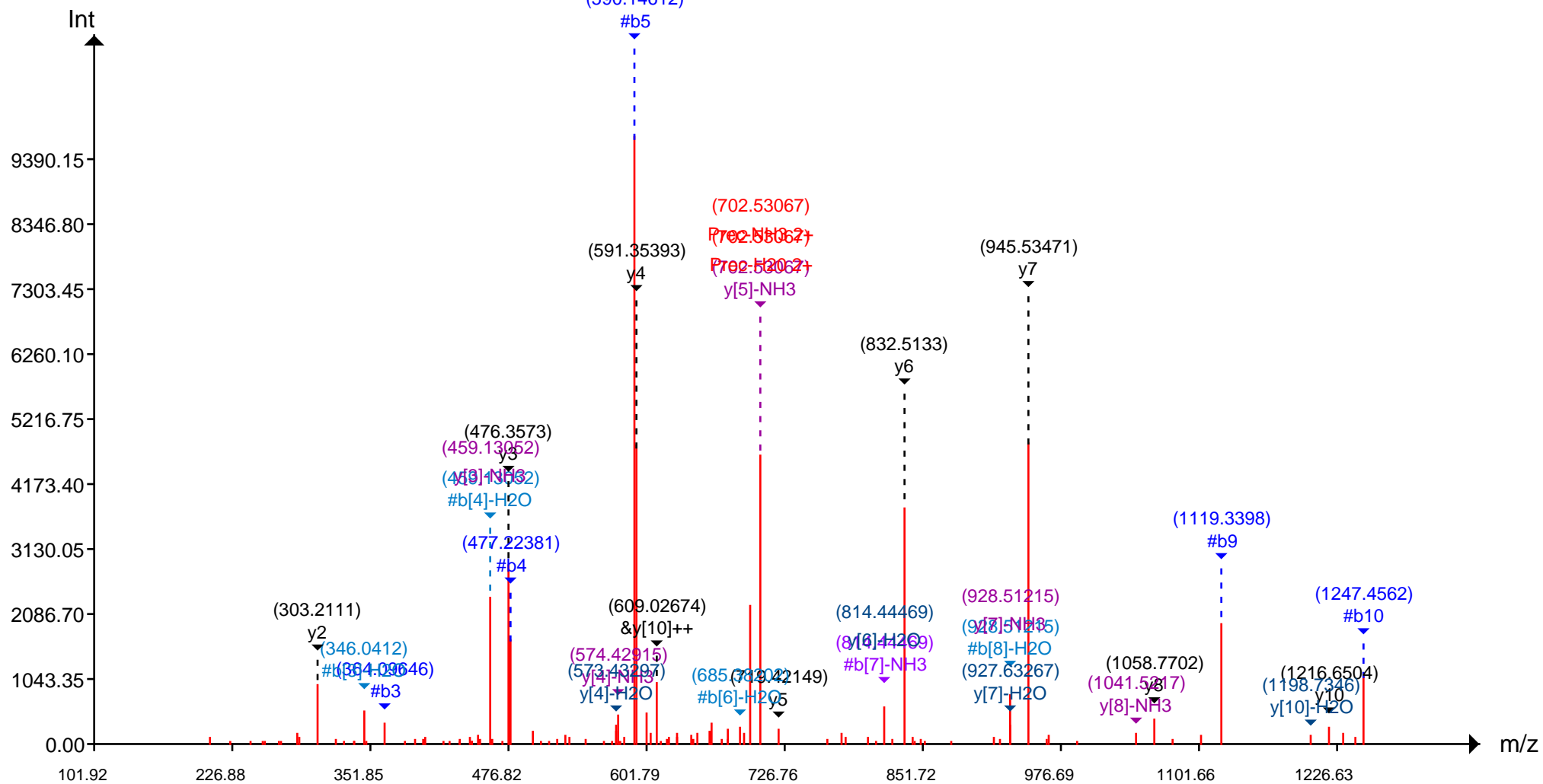
Confidence level 99.00%

Precursor m/z 968.51201

Identity Threshold 48.98

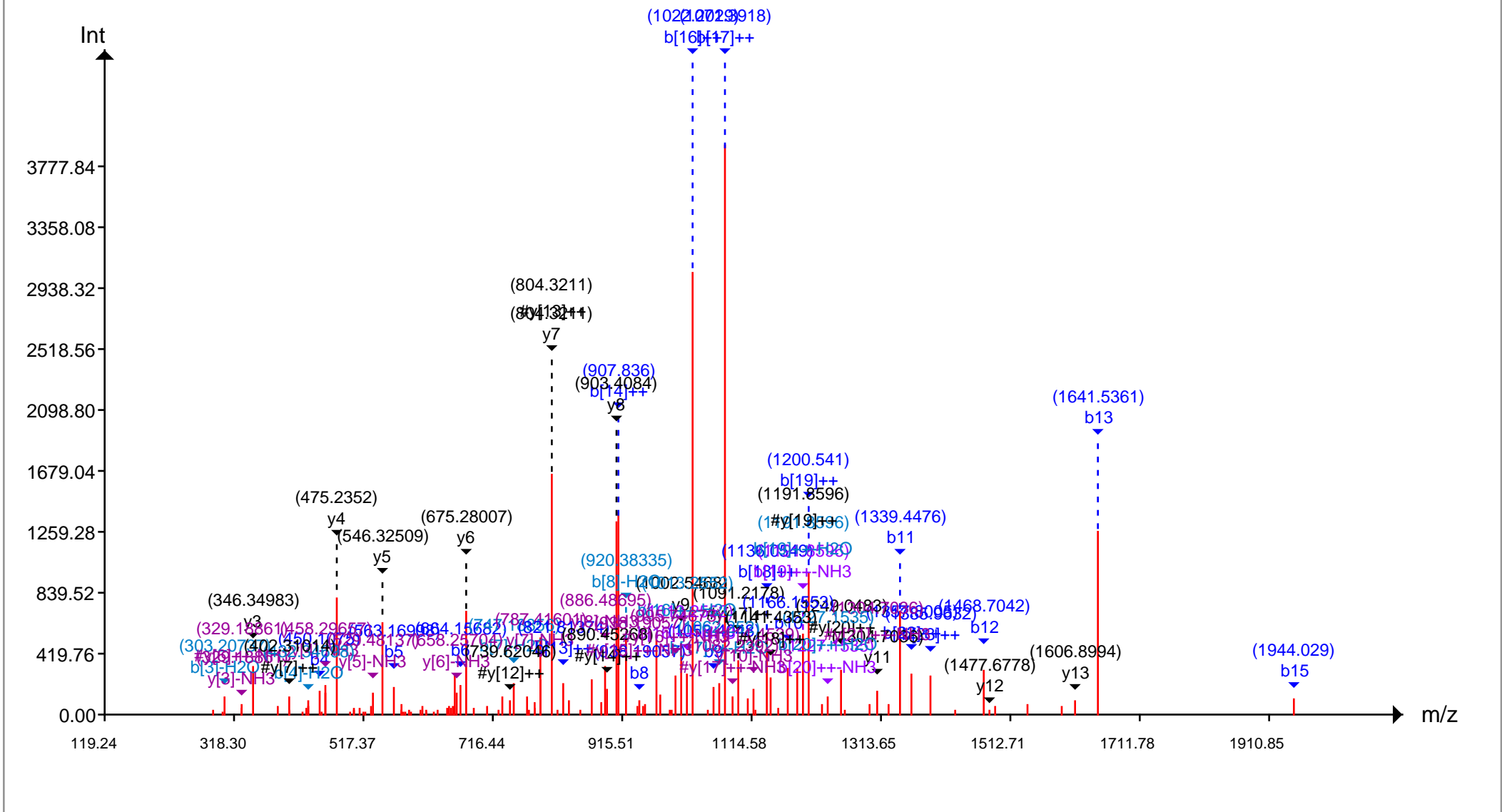
Charge 2+

Mass error 0.001076



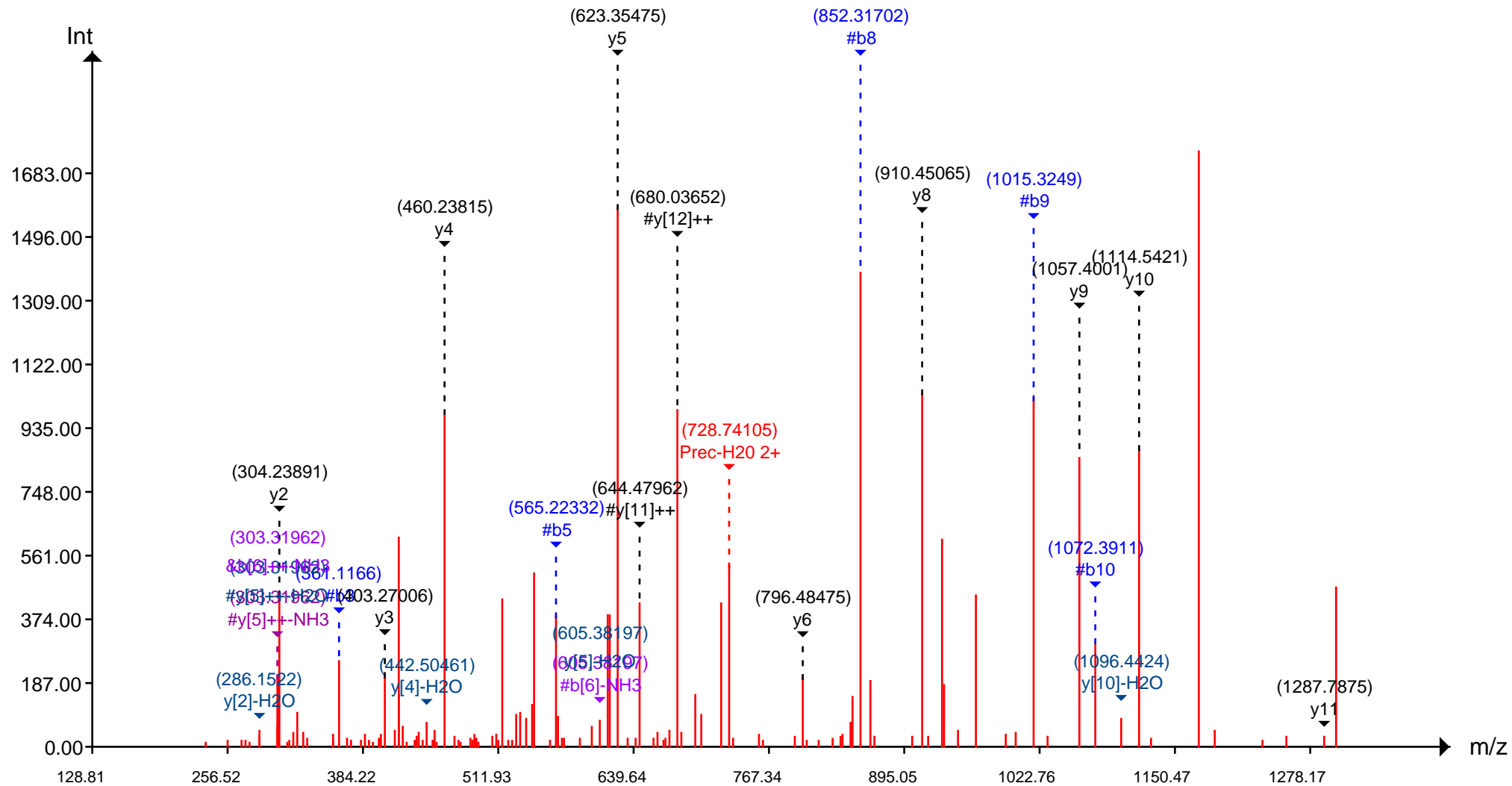
Info

| | | | |
|--------------------------|-------------------------------------|---------------------------|--------|
| Modified Sequence | AcD3-C<Cmm*>GTILLQDK<AcD3K*>QR-COOH | Ion score | 78.02 |
| Protein | O60232 | Confidence level | 99.00% |
| Precursor m/z | 711.38805 | Identity Threshold | 48.85 |
| Charge | 2+ | Mass error | 4.3E-5 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-TSSEITTK<AcD3K*>DLK<AcD3K*>EK<AcD3K*>K<AcD3K*>EVVEEAENGR-COOH | Ion score | 86.64 |
| Protein | P06454 | Confidence level | 99.00% |
| Precursor m/z | 982.85492 | Identity Threshold | 48.24 |
| Charge | 3+ | Mass error | 1.008401 |



Info

Modified Sequence AcD3-AAK<AcD3K*>GFGGK<AcD3K*>YGVER-COOH

Ion score 53.79

Protein P14317

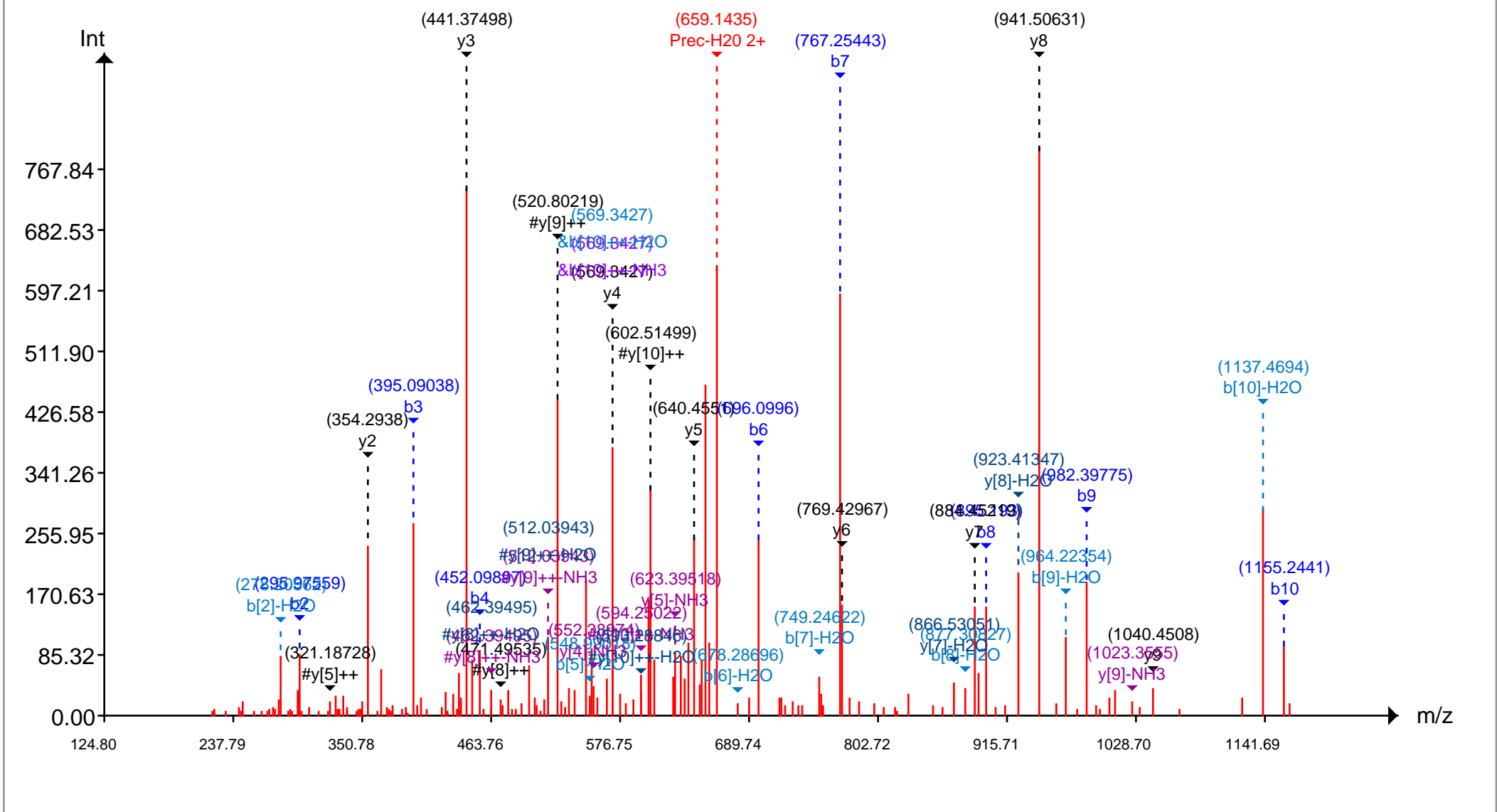
Confidence level 99.00%

Precursor m/z 737.90398

Identity Threshold 48.85

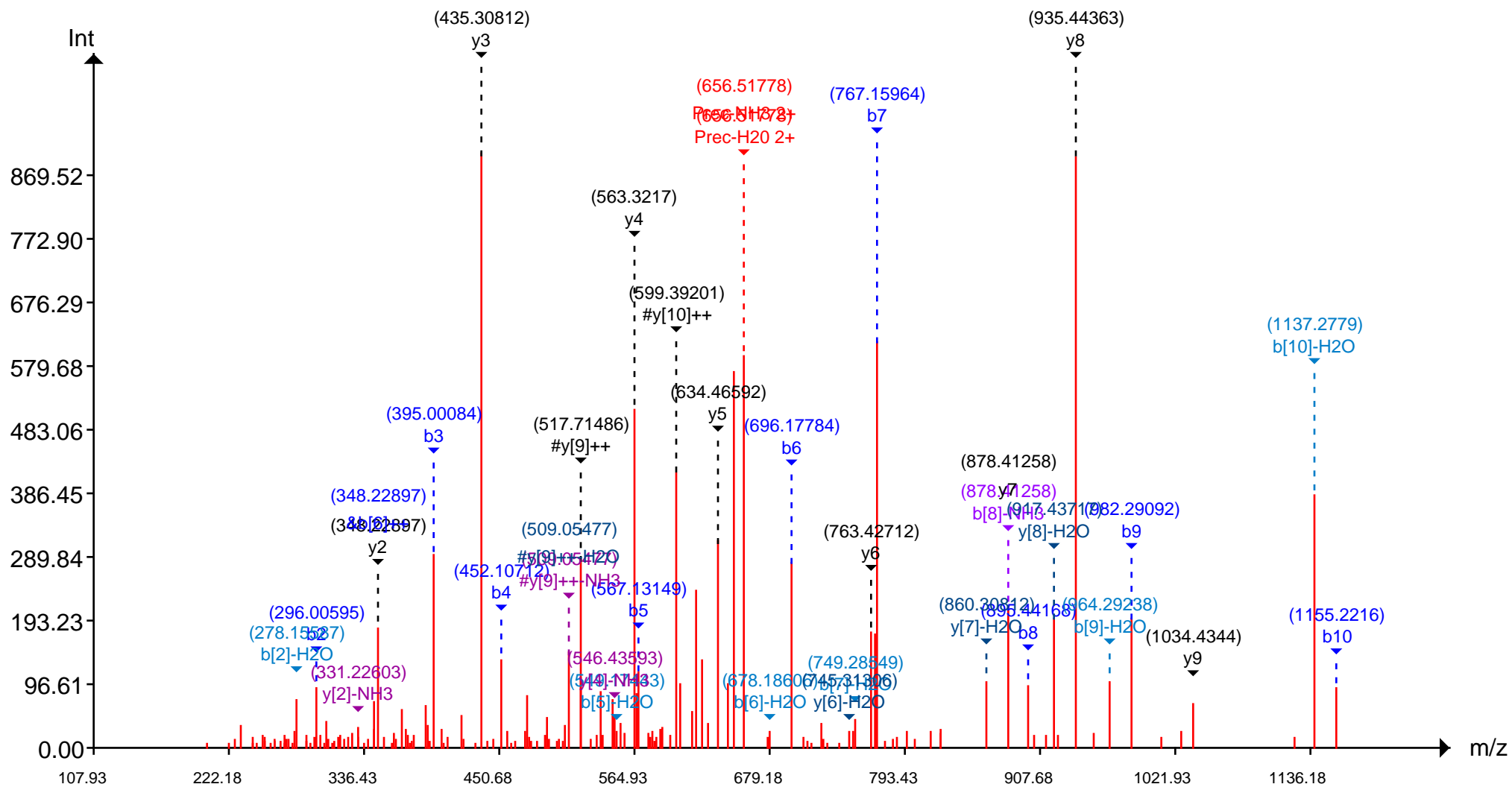
Charge 2+

Mass error 7.9E-4



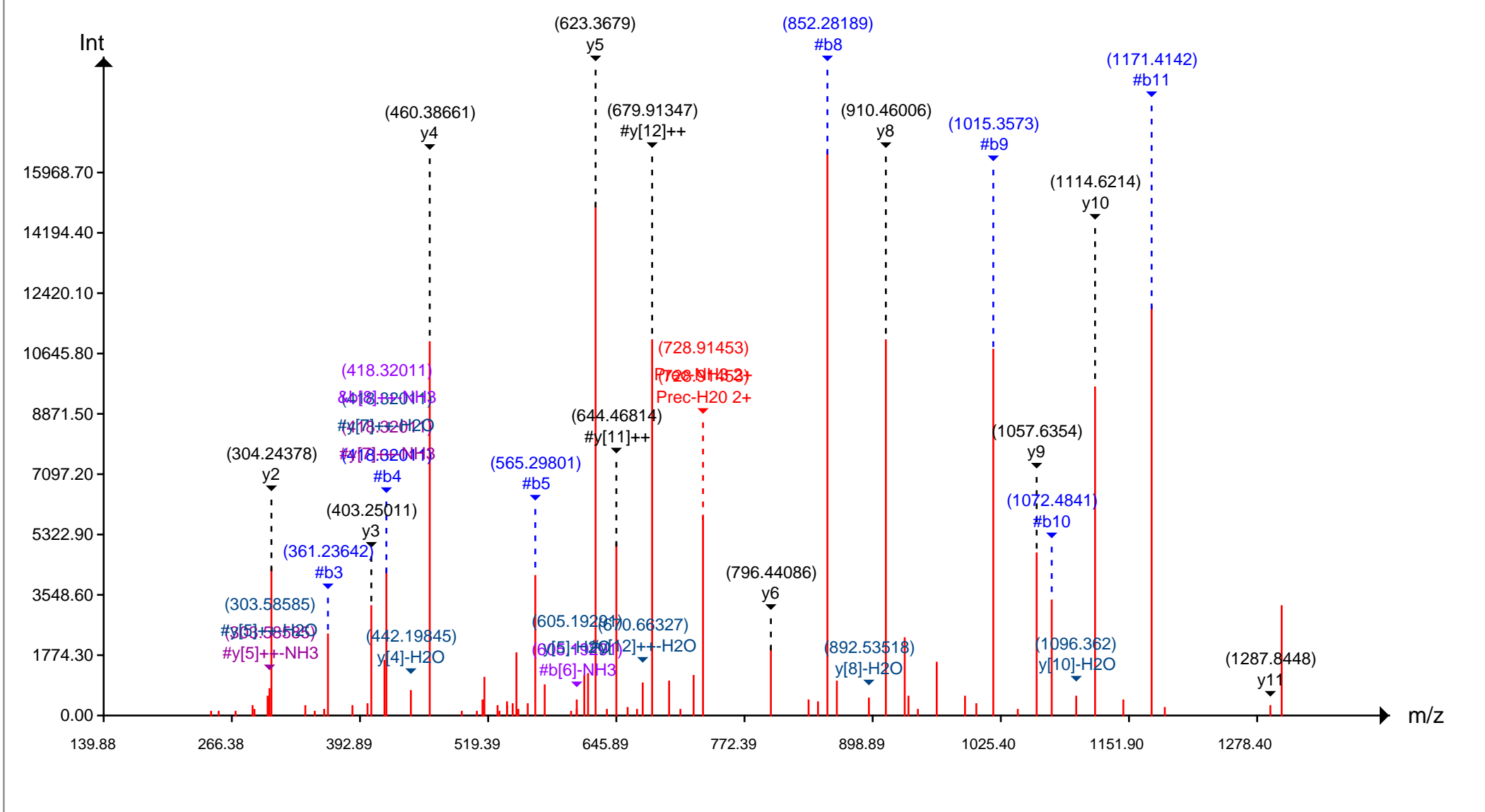
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-SYVGDEAQS<AcD3K*>R<C13>-COOH | Ion score | 68.55 |
| Protein | P62736, P60709, P68032, P63261, P63267, P68133 | Confidence level | 99.00% |
| Precursor m/z | 668.34153 | Identity Threshold | 48.89 |
| Charge | 2+ | Mass error | 4.63E-4 |

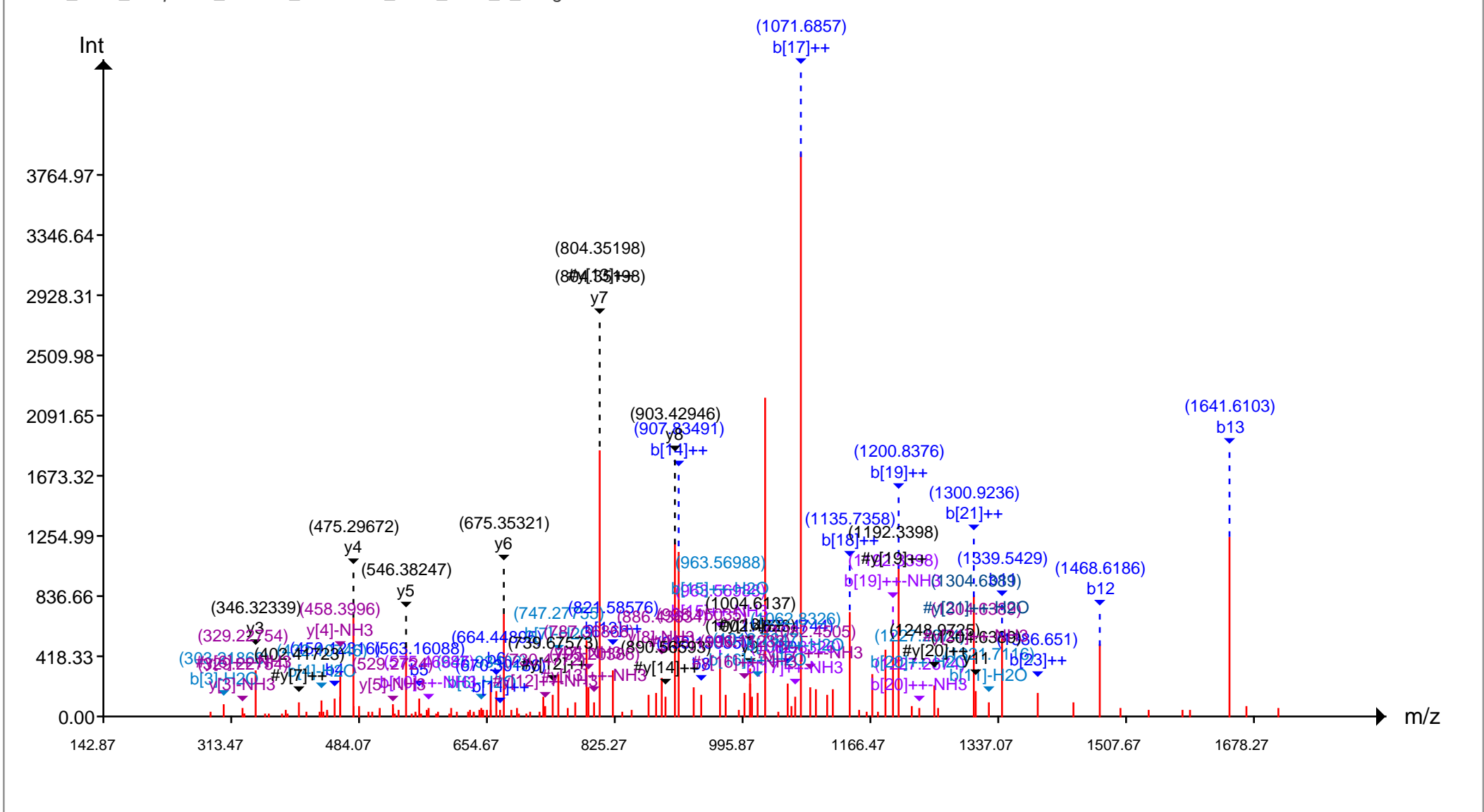


Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-SYVGDEAQSK<AcD3K*>R-COOH | Ion score | 65.32 |
| Protein | P62736, P60709, P68032, P63261, P63267, P68133 | Confidence level | 99.00% |
| Precursor m/z | 665.3316 | Identity Threshold | 48.48 |
| Charge | 2+ | Mass error | 7.45E-4 |

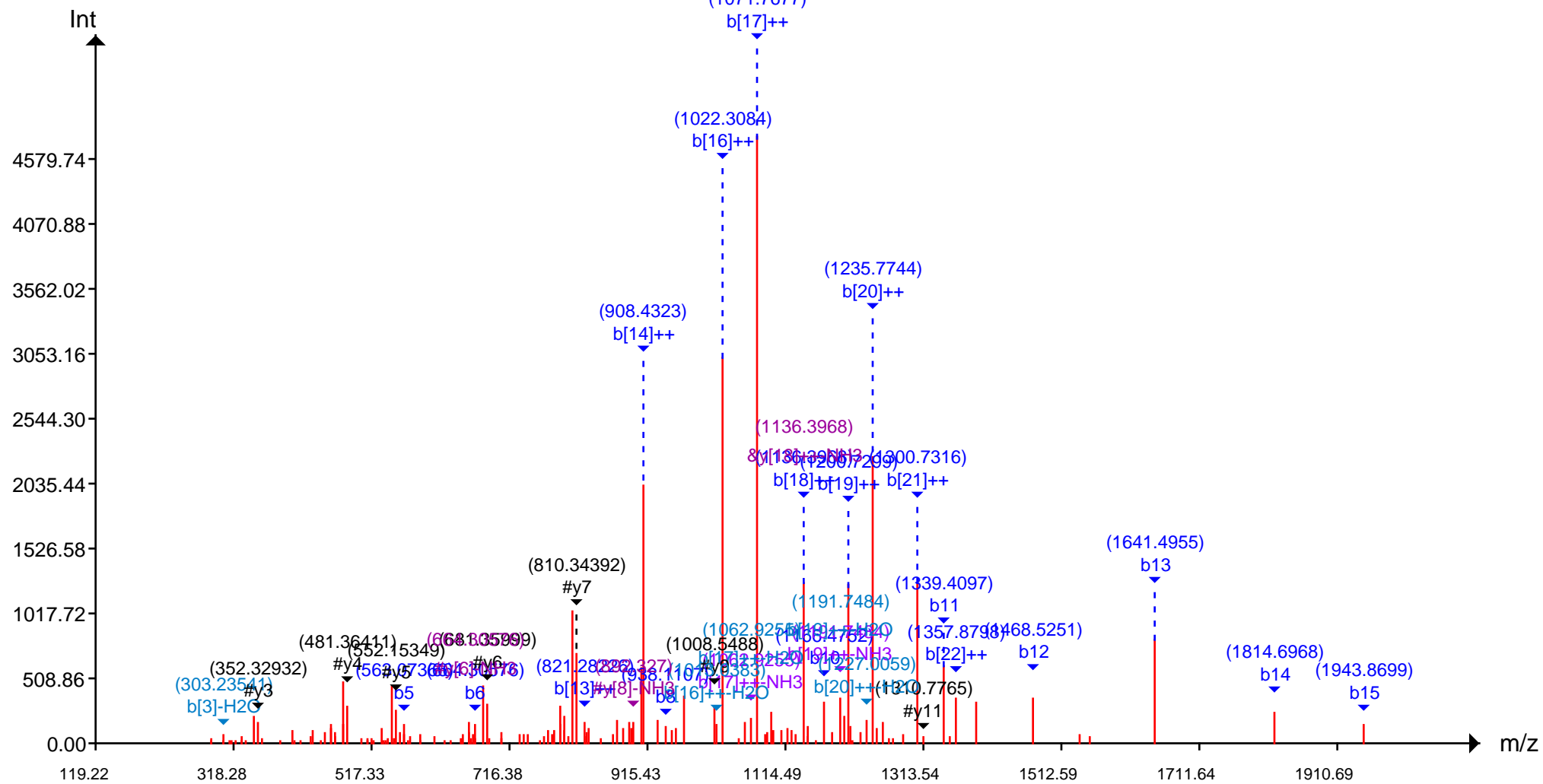


| Info | | | |
|--------------------------|---|---------------------------|--------|
| Modified Sequence | AcD3-AAK<AcD3K*>GFGGK<AcD3K*>YGVER-COOH | Ion score | 75.52 |
| Protein | P14317 | Confidence level | 99.00% |
| Precursor m/z | 737.90381 | Identity Threshold | 48.87 |
| Charge | 2+ | Mass error | 4.5E-4 |

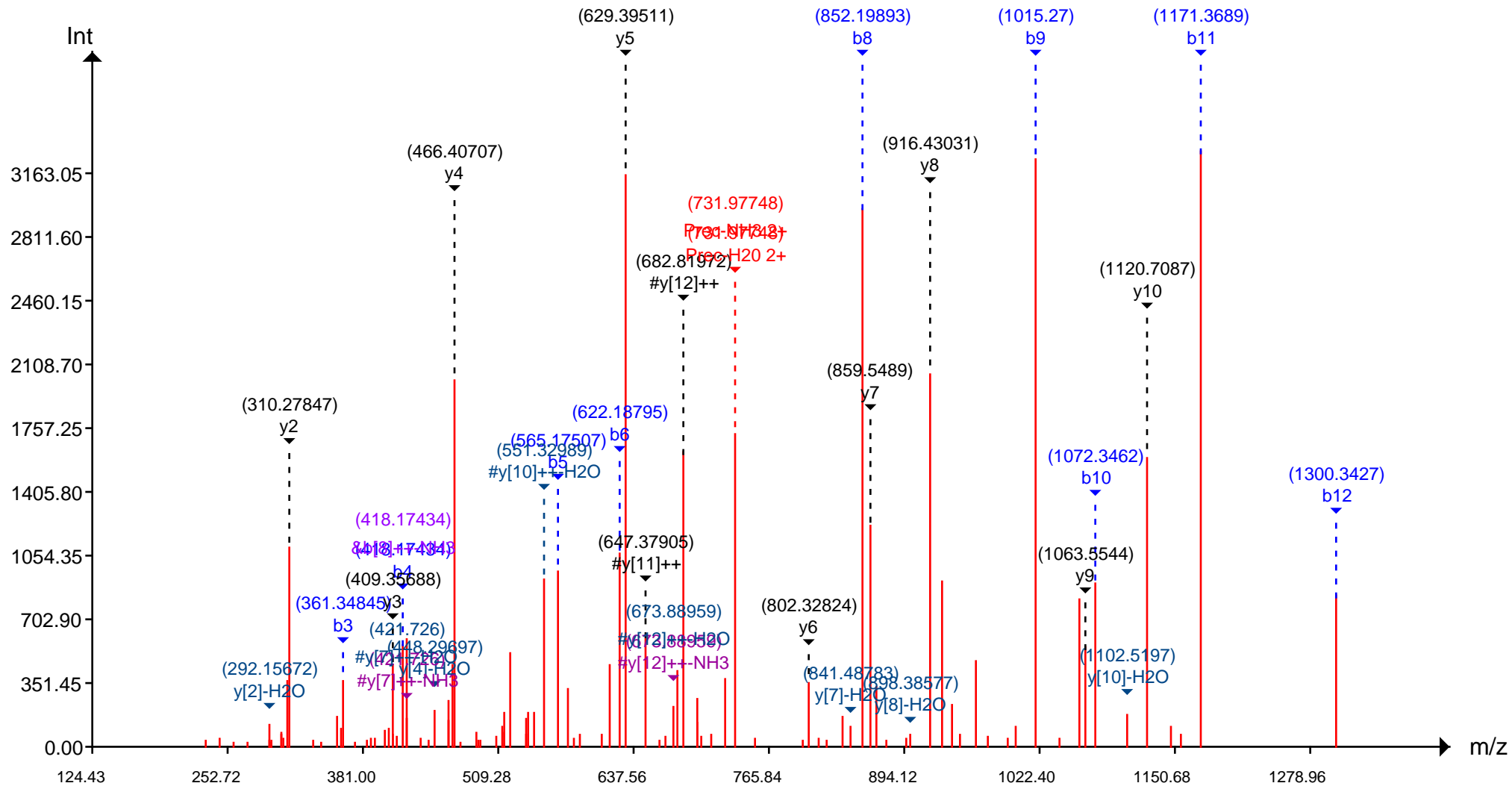


Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-TSSEITTK<AcD3K*>DLK<AcD3K*>EK<AcD3K*>K<AcD3K*>EVVEEAENGR-COOH | Ion score | 59.22 |
| Protein | P06454 | Confidence level | 99.00% |
| Precursor m/z | 982.51923 | Identity Threshold | 48.35 |
| Charge | 3+ | Mass error | 0.001331 |

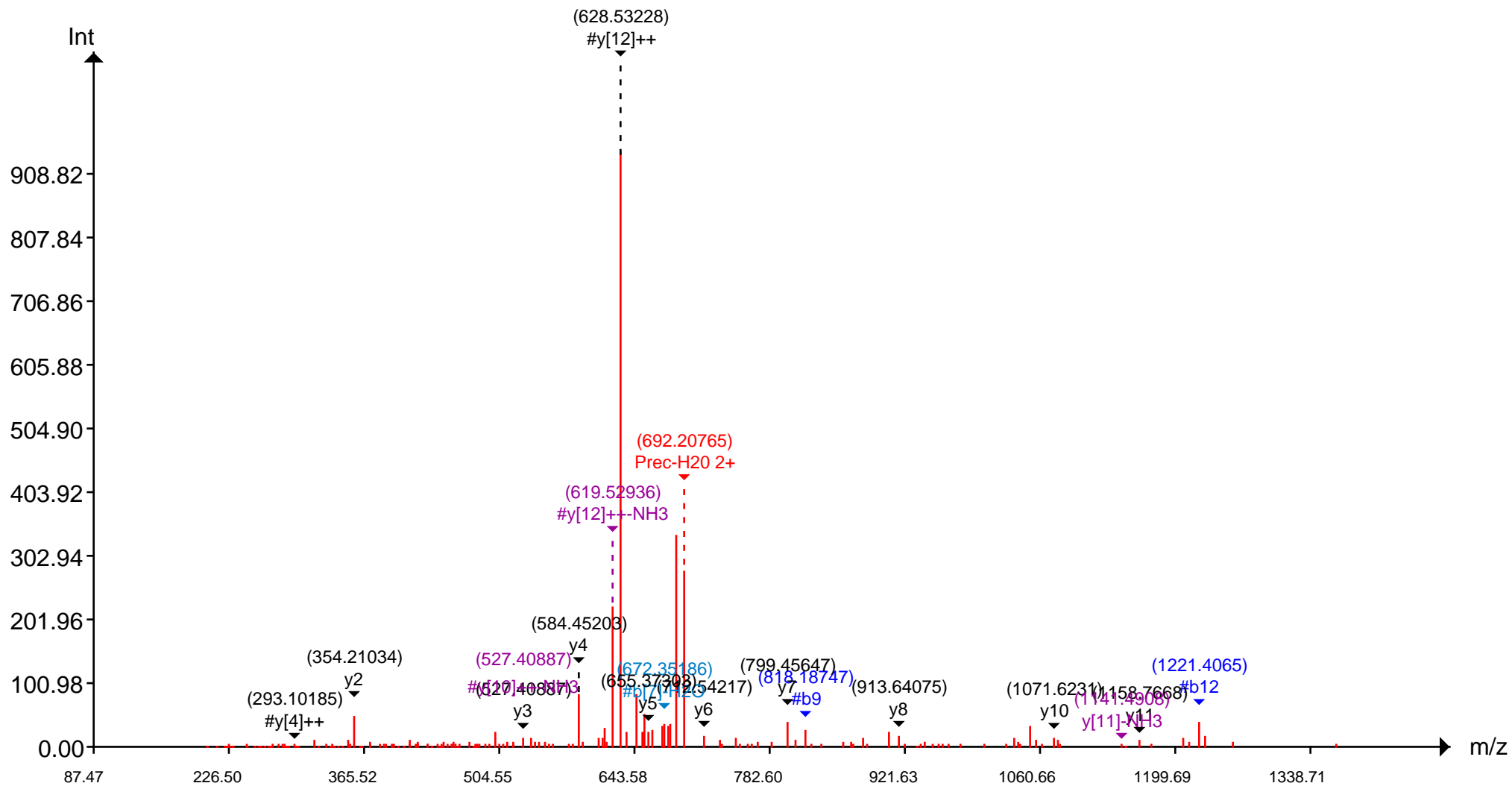


| Info | | | |
|--------------------------|---|---------------------------|--------|
| Modified Sequence | AcD3-TSSEITTK<AcD3K*>DLK<AcD3K*>EK<AcD3K*>K<AcD3K*>EVVEEAENGR<C13>-COOH | Ion score | 54.05 |
| Protein | P06454 | Confidence level | 99.00% |
| Precursor Ion | 984.52594 | Identity Threshold | 48.03 |
| Charge | 3+ | Mass error | 0.0013 |



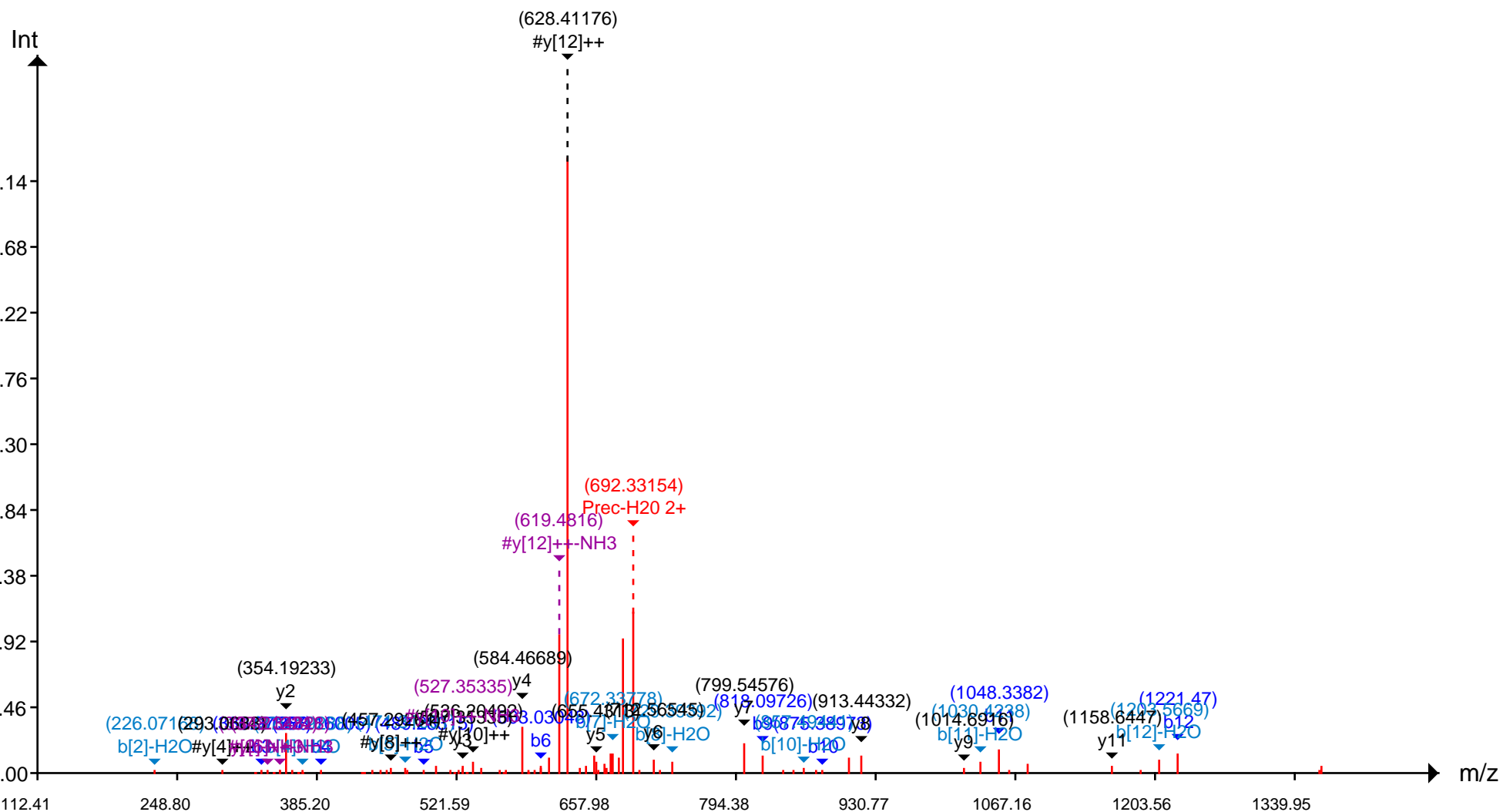
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-AAK<AcD3K*>GFGGK<AcD3K*>YGVER<C13>-COOH | Ion score | 63.04 |
| Protein | P14317 | Confidence level | 99.00% |
| Precursor m/z | 740.91381 | Identity Threshold | 48.30 |
| Charge | 2+ | Mass error | 3.09E-4 |



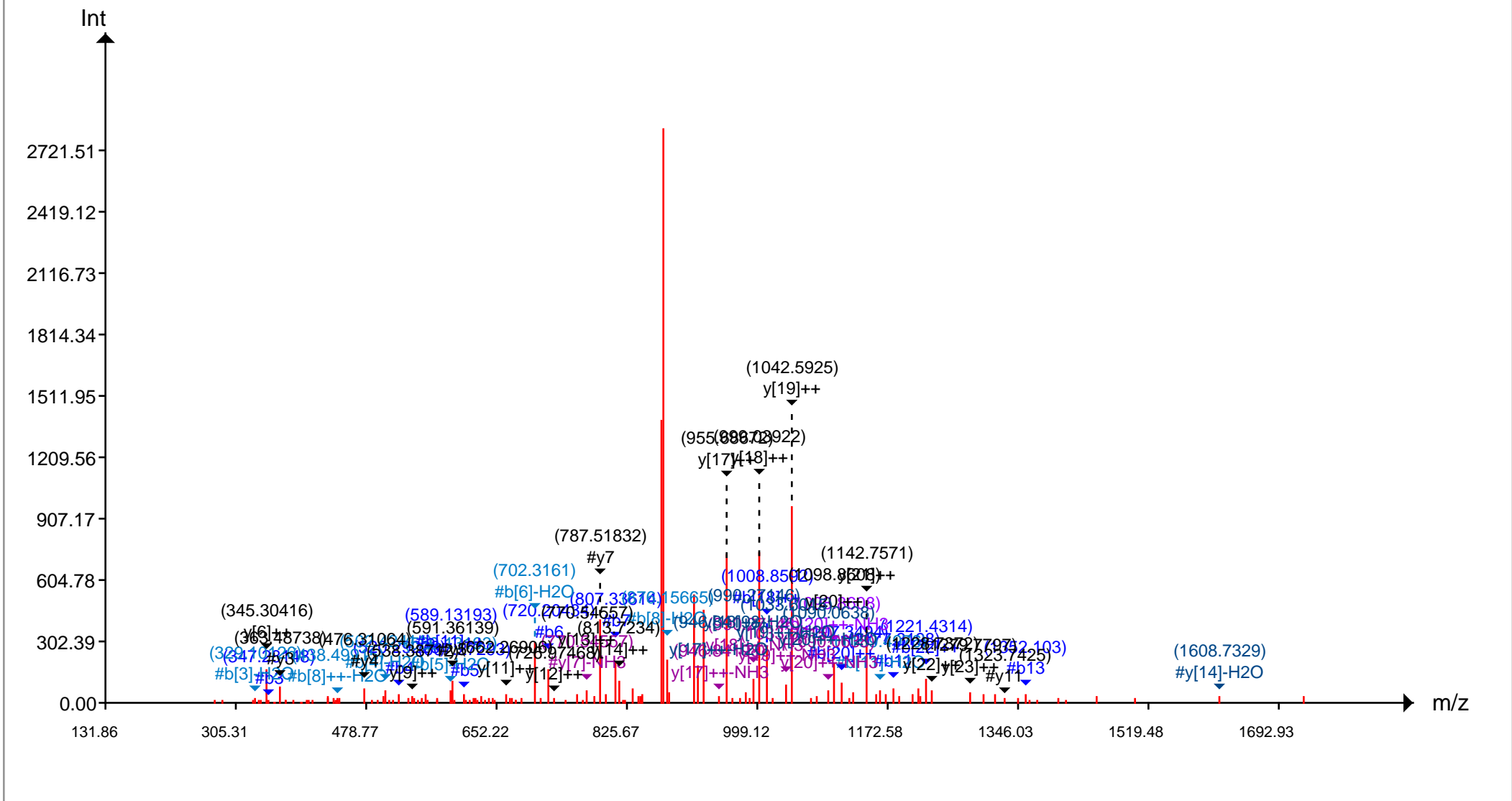
Info

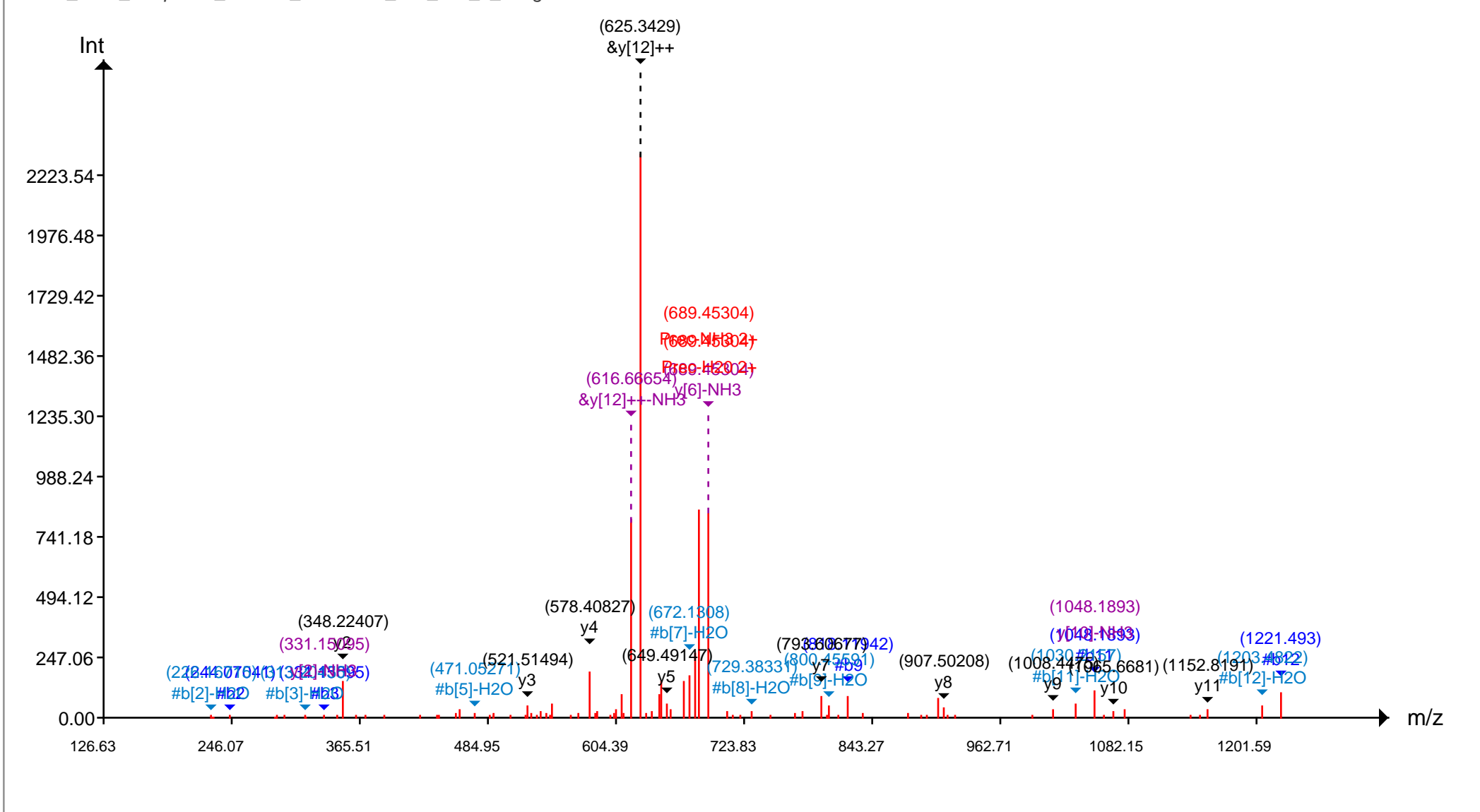
| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-TPSGTNSGAGK<AcD3K*>K<AcD3K*>R<C13>-COOH | Ion score | 56.93 |
| Protein | P62877 | Confidence level | 99.00% |
| Precursor m/z | 701.39054 | Identity Threshold | 48.53 |
| Charge | 2+ | Mass error | -1.3E-5 |



Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-TPSGTNSGAGK<AcD3K*>K<AcD3K*>R<C13>-COOH | Ion score | 60.04 |
| Protein | P62877 | Confidence level | 99.00% |
| Precursor m/z | 701.39031 | Identity Threshold | 48.58 |
| Charge | 2+ | Mass error | -4.73E-4 |





Info

Modified Sequence AcD3-TPSGTNSGAGK<AcD3K*>K<AcD3K*>R-COOH

Ion score 55.78

Protein P62877

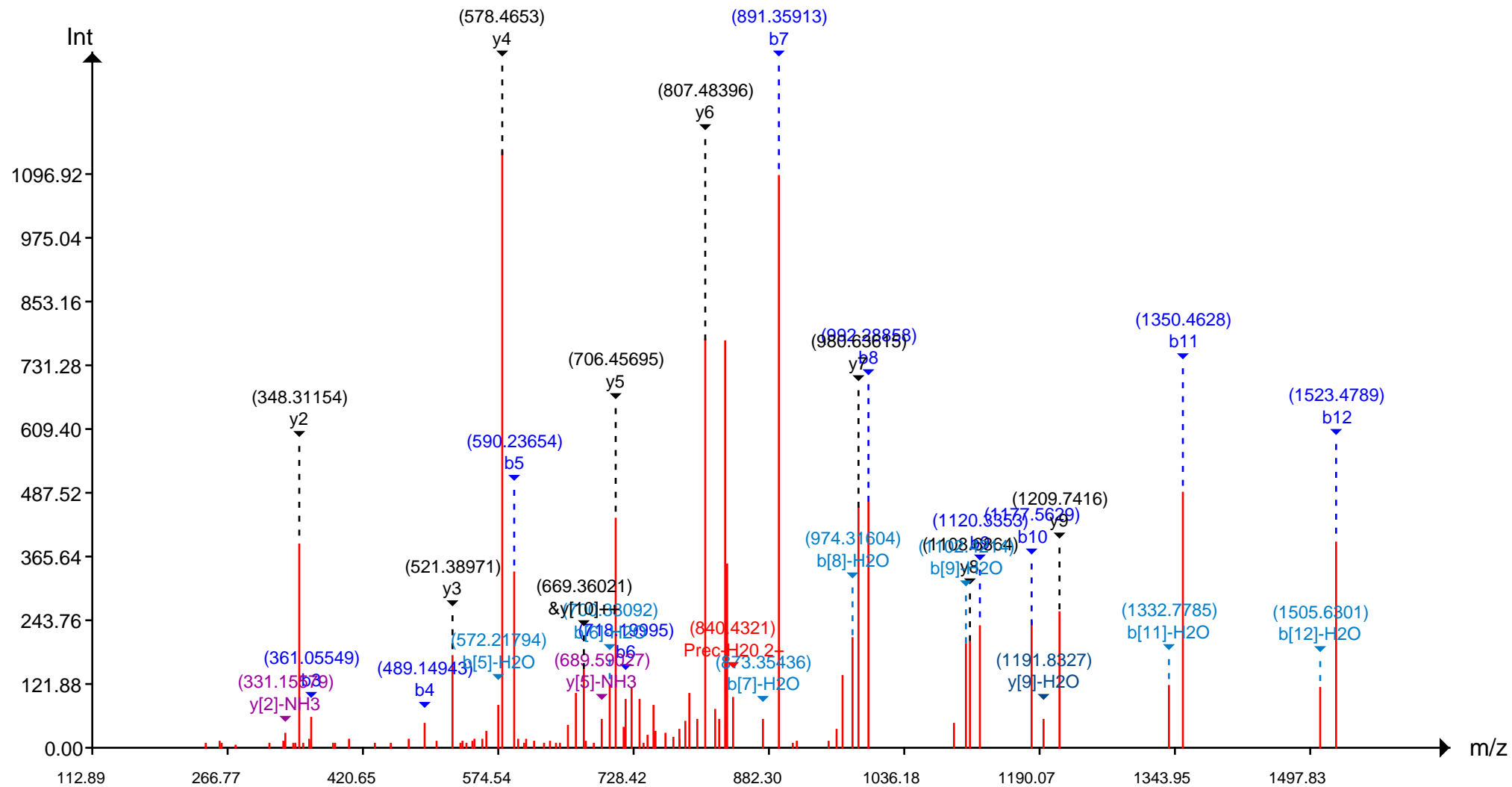
Confidence level 99.00%

Precursor m/z 698.38003

Identity Threshold 48.85

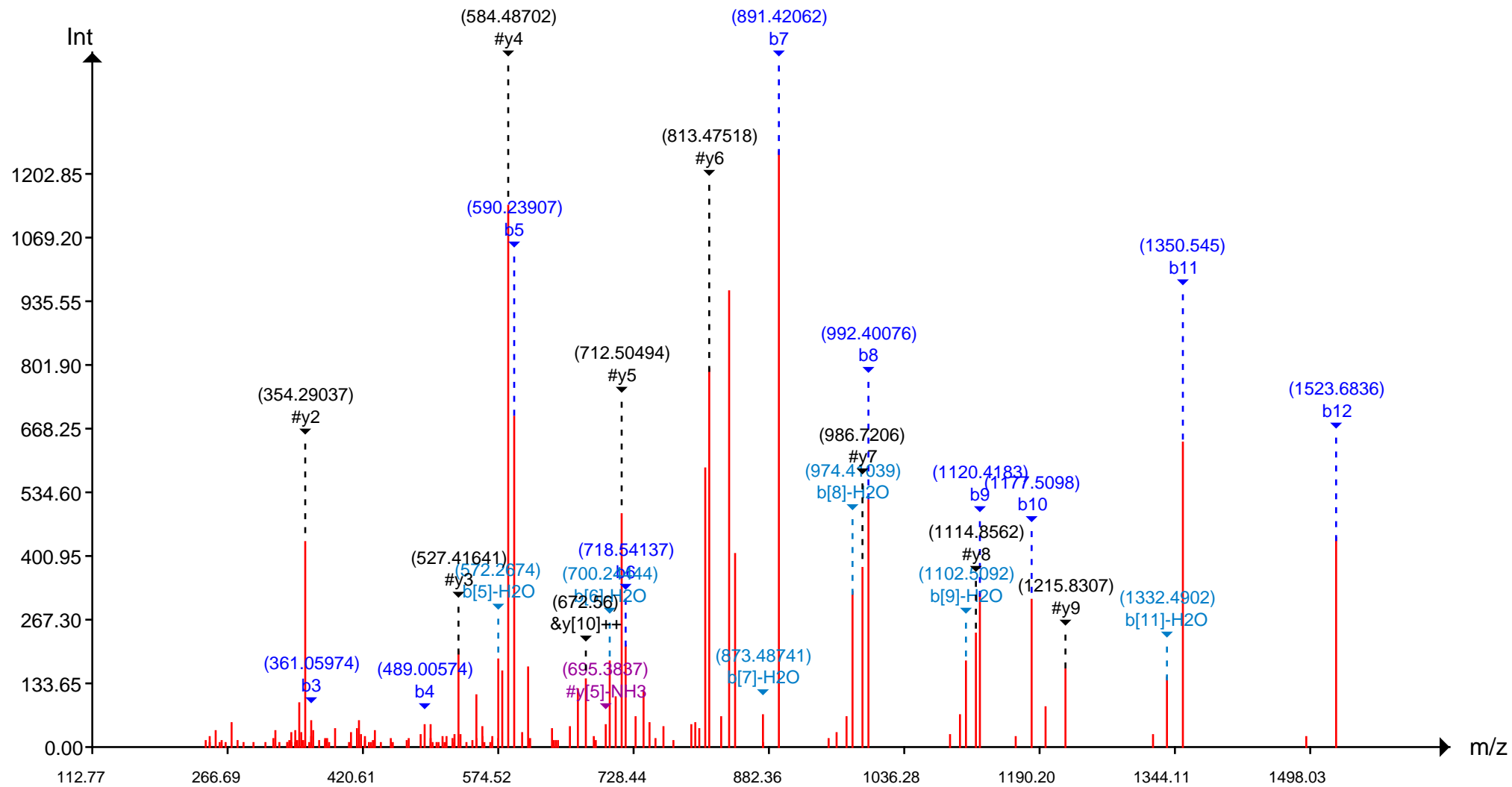
Charge 2+

Mass error -8.91E-4



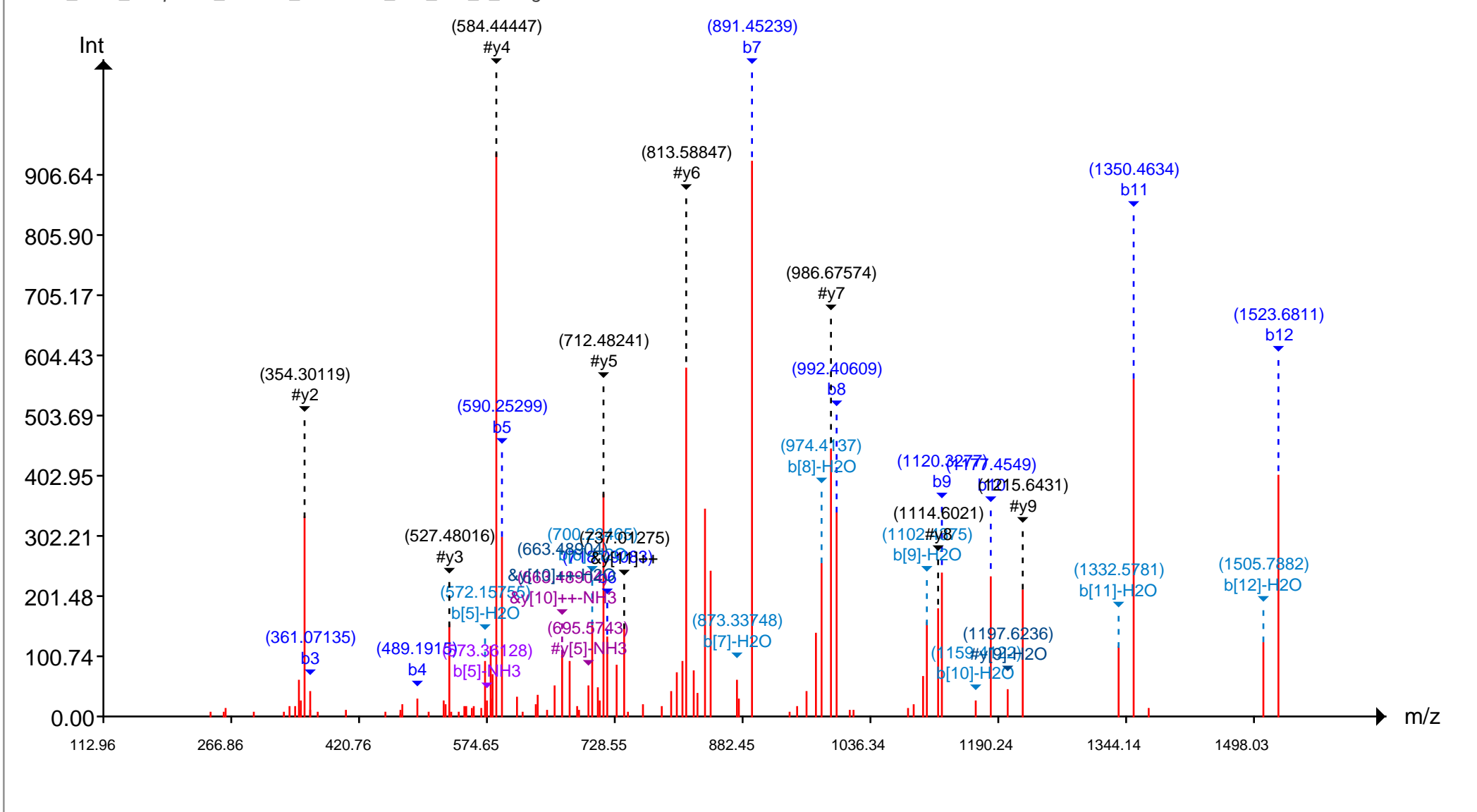
Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-GEEQTQK<AcD3K*>TQGK<AcD3K*>K<AcD3K*>R-COOH | Ion score | 90.52 |
| Protein | Q9UEE9 | Confidence level | 99.00% |
| Precursor m/z | 849.46415 | Identity Threshold | 48.60 |
| Charge | 2+ | Mass error | 4.03E-4 |



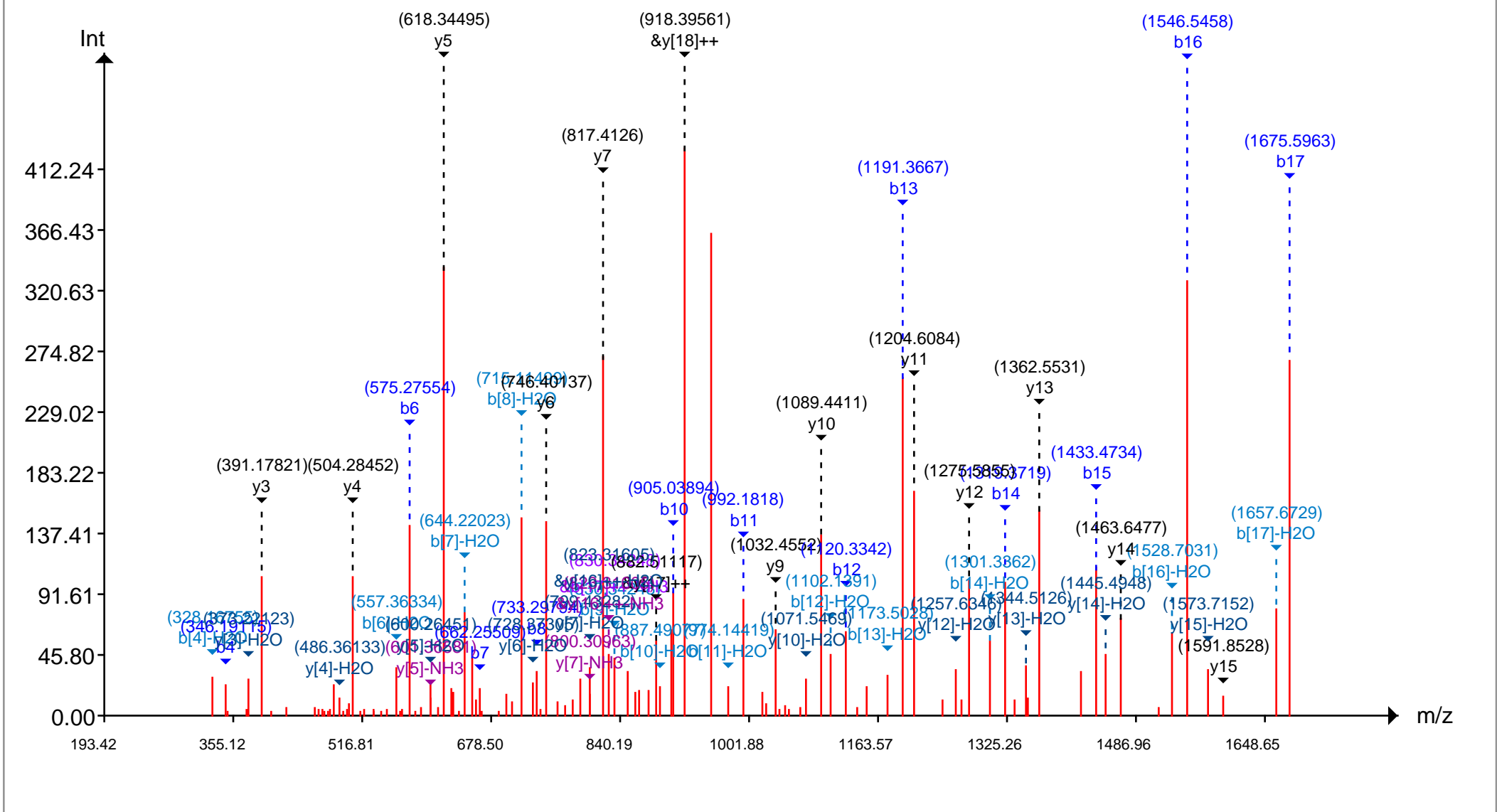
Info

| | | | |
|--------------------------|--|---------------------------|--------|
| Modified Sequence | AcD3-GEEQTQK<AcD3K*>TQGK<AcD3K*>K<AcD3K*>R<C13>-COOH | Ion score | 79.28 |
| Protein | Q9UEE9 | Confidence level | 99.00% |
| Precursor m/z | 852.47404 | Identity Threshold | 48.07 |
| Charge | 2+ | Mass error | 4.1E-5 |



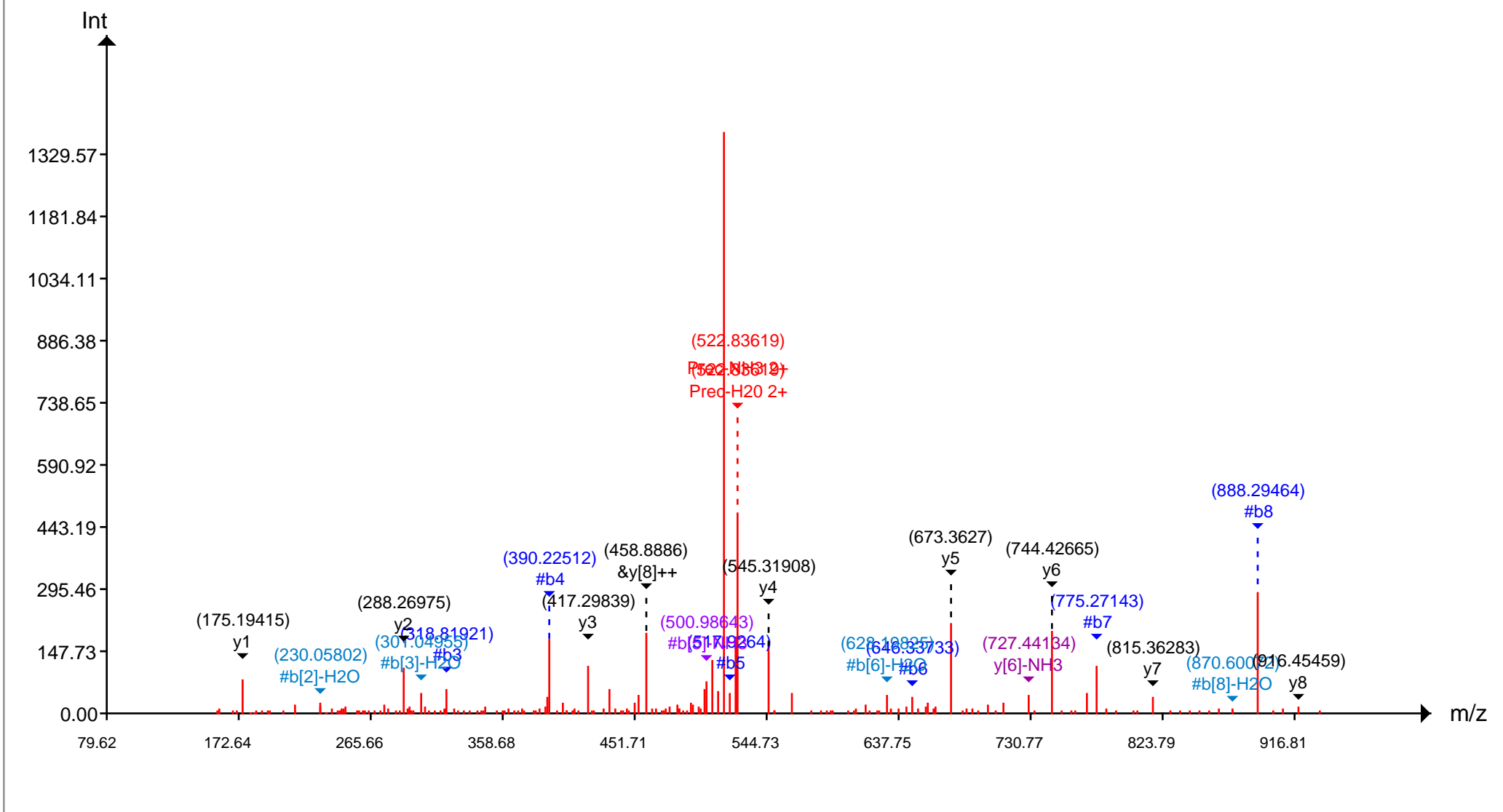
Info

| | | | |
|--------------------------|--|---------------------------|---------|
| Modified Sequence | AcD3-GEEQTQK<AcD3K*>TQGK<AcD3K*>K<AcD3K*>R<C13>-COOH | Ion score | 73.84 |
| Protein | Q9UEE9 | Confidence level | 99.00% |
| Precursor m/z | 852.47415 | Identity Threshold | 48.08 |
| Charge | 2+ | Mass error | 2.61E-4 |



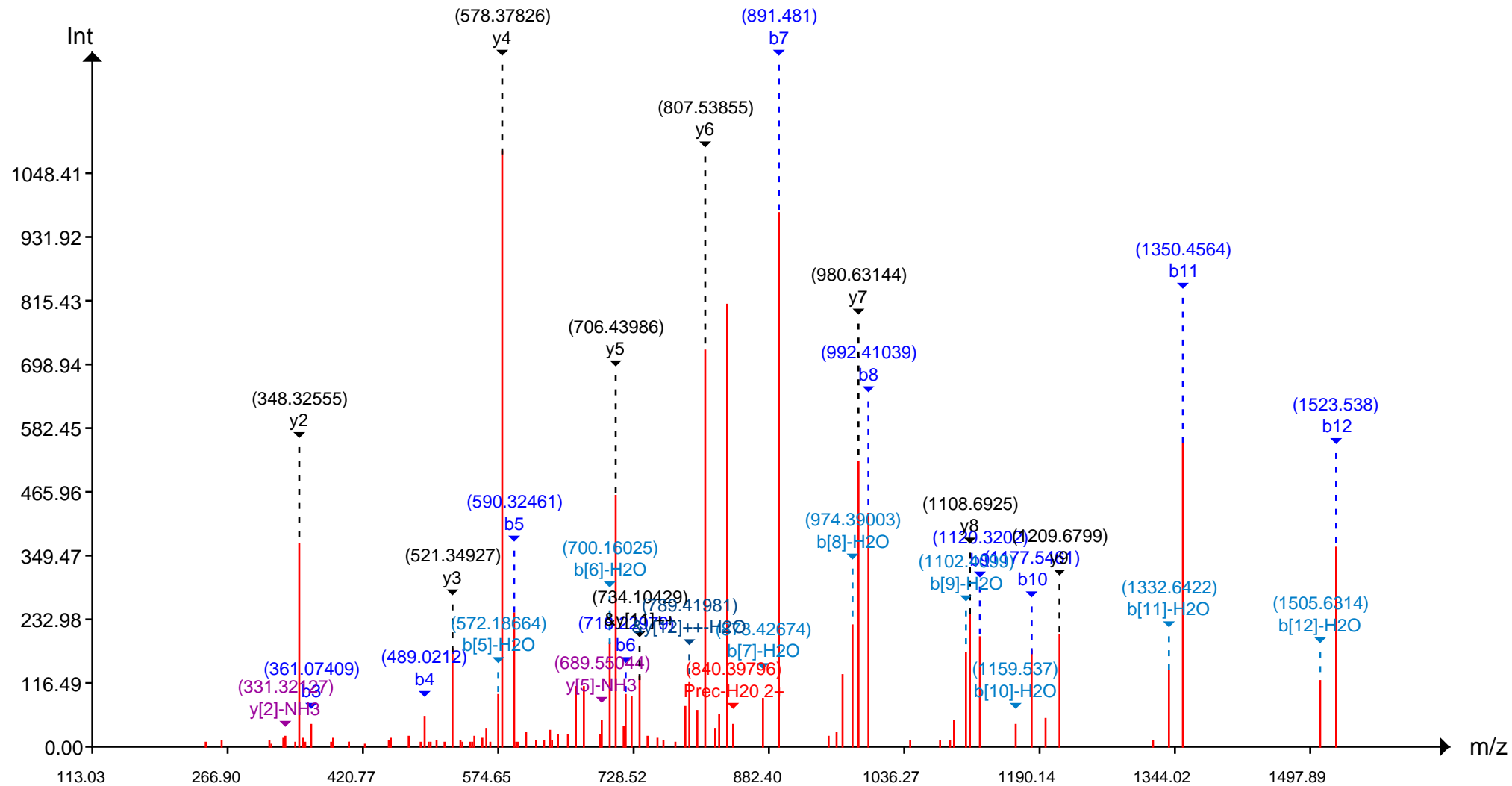
Info

| | | | |
|--------------------------|--------------------------------|---------------------------|---------|
| Modified Sequence | AcD3-GATAQTSADGGSQAQNLESR-COOH | Ion score | 109.62 |
| Protein | Q9Y5V3 | Confidence level | 99.00% |
| Precursor m/z | 968.95541 | Identity Threshold | 46.77 |
| Charge | 2+ | Mass error | 1.35E-4 |



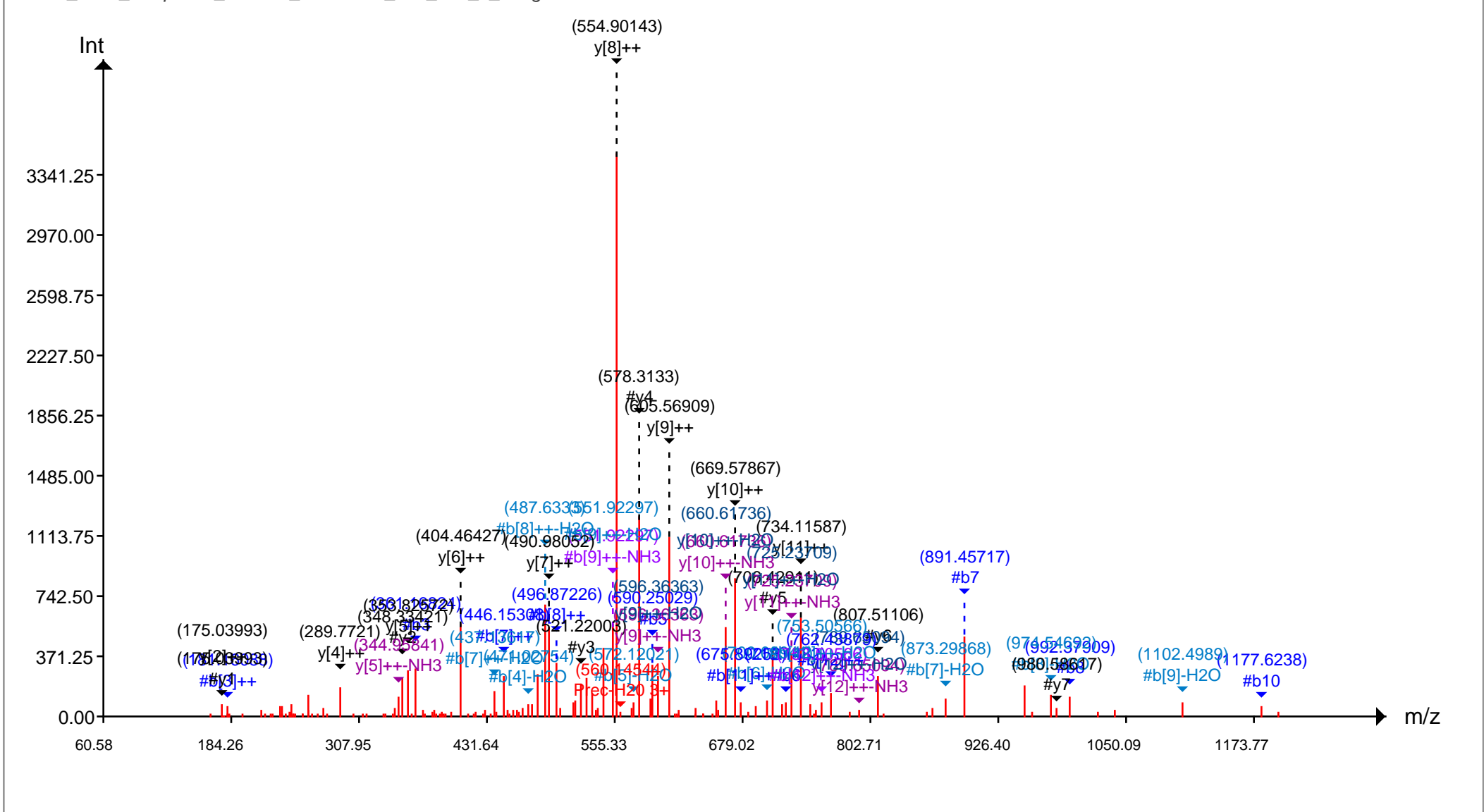
Info

| | | | |
|--------------------------|---------------------|---------------------------|---------|
| Modified Sequence | AcD3-TTAAQQELR-COOH | Ion score | 59.34 |
| Protein | P35580 | Confidence level | 99.00% |
| Precursor m/z | 531.78459 | Identity Threshold | 49.03 |
| Charge | 2+ | Mass error | 1.85E-4 |

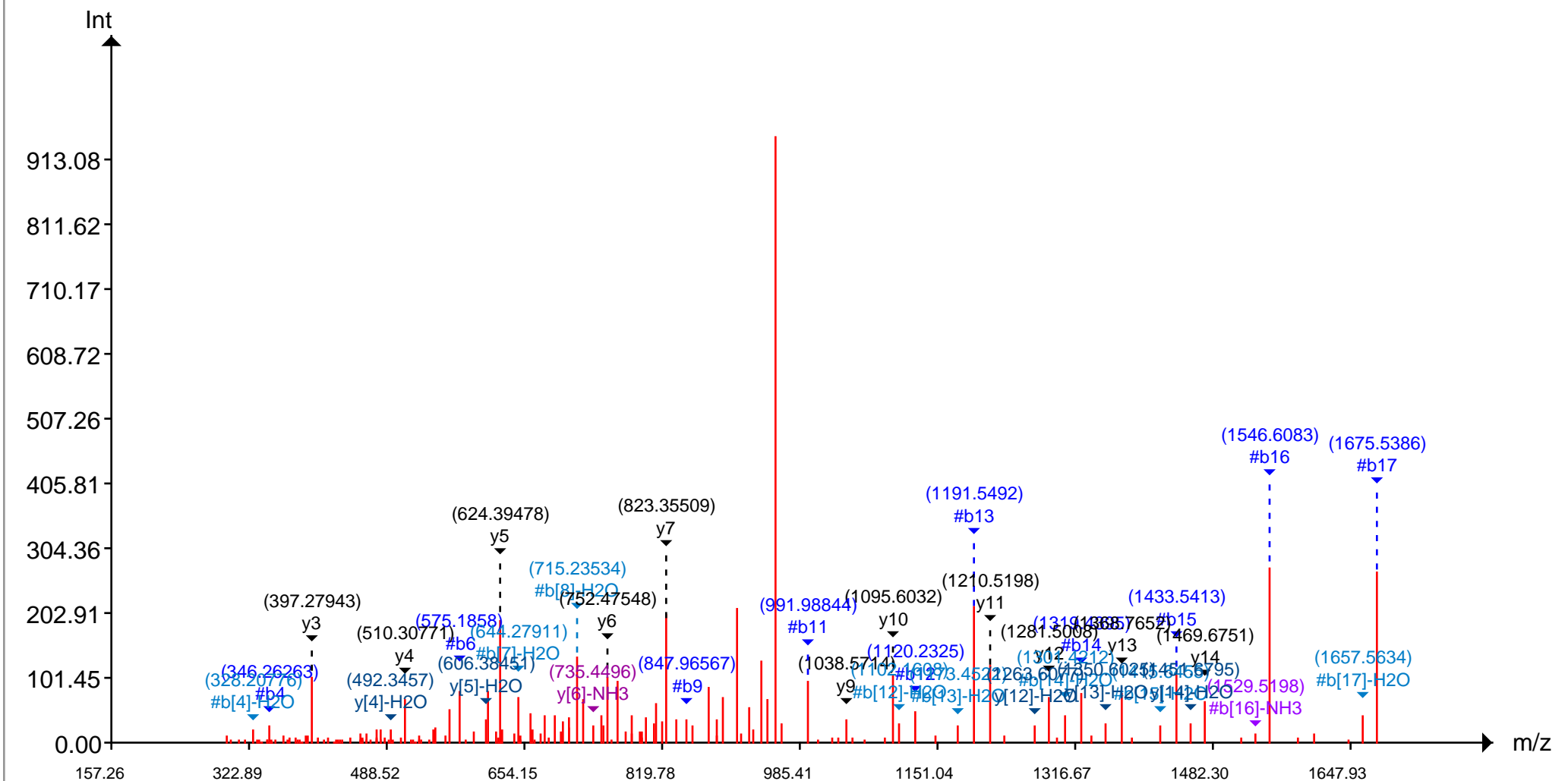


Info

| | | | |
|--------------------------|---|---------------------------|---------|
| Modified Sequence | AcD3-GEEQTQK<AcD3K*>TQGK<AcD3K*>K<AcD3K*>R-COOH | Ion score | 90.68 |
| Protein | Q9UEE9 | Confidence level | 99.00% |
| Precursor m/z | 849.46419 | Identity Threshold | 48.60 |
| Charge | 2+ | Mass error | 4.83E-4 |



| Info | | | |
|--------------------------|---|---------------------------|-----------|
| Modified Sequence | AcD3-GEEQTQK<AcD3K*>TQGK<AcD3K*>K<AcD3K*>R-COOH | Ion score | 68.57 |
| Protein | Q9UEE9 | Confidence level | 99.00% |
| Precursor m/z | 566.64469 | Identity Threshold | 48.60 |
| Charge | 3+ | Mass error | -0.001103 |



Info

Modified Sequence AcD3-GATAQTSADGSQAQNLESR<C13>-COOH

Ion score 97.77

Protein Q9Y5V3

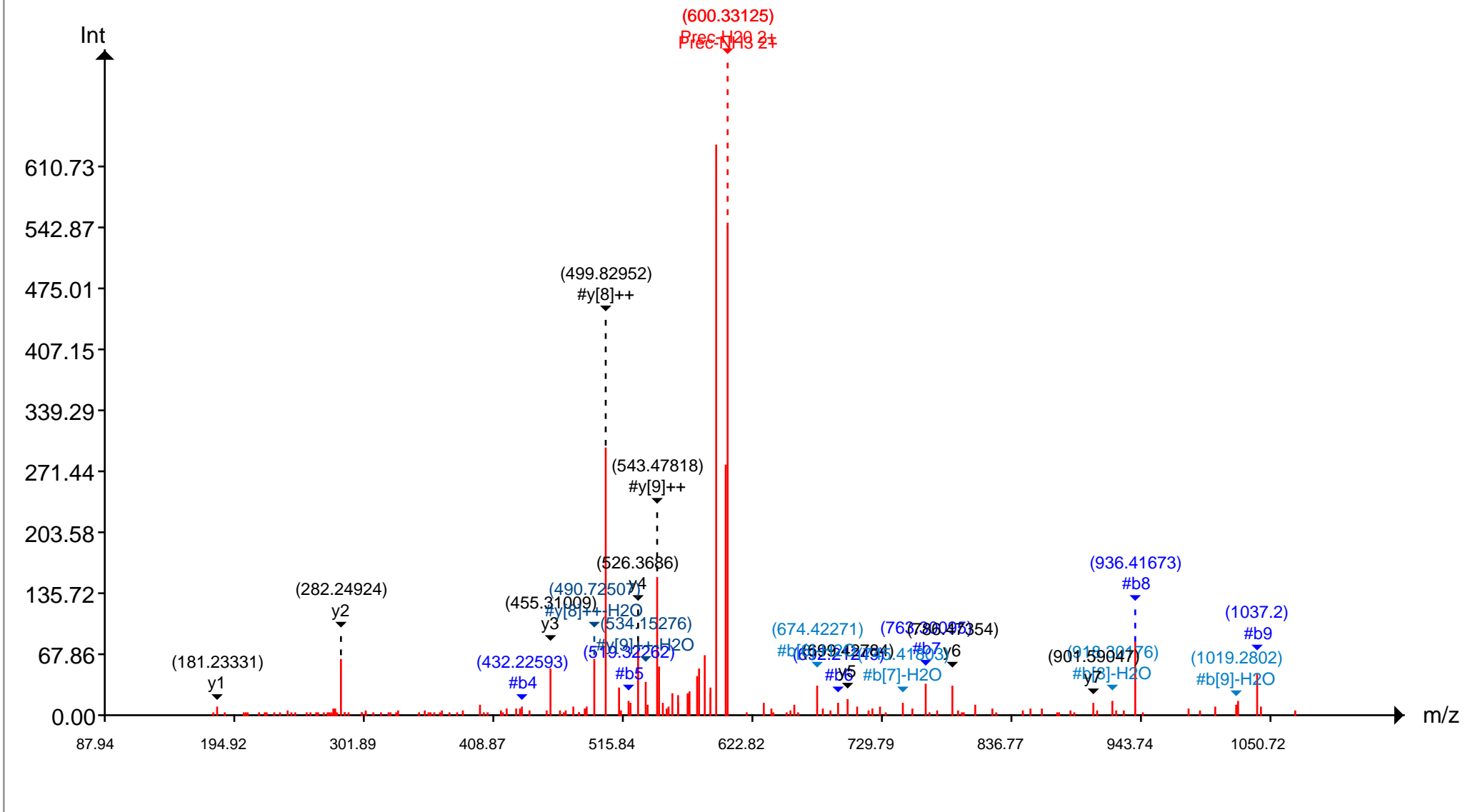
Confidence level 99.00%

Precursor m/z 971.96531

Identity Threshold 47.55

Charge 2+

Mass error -2.07E-4



Info

Modified Sequence AcD3-SSPDSK<AcD3K*>AK<AcD3K*>TR<C13>-COOH

Ion score 49.62

Protein Q9UQ35

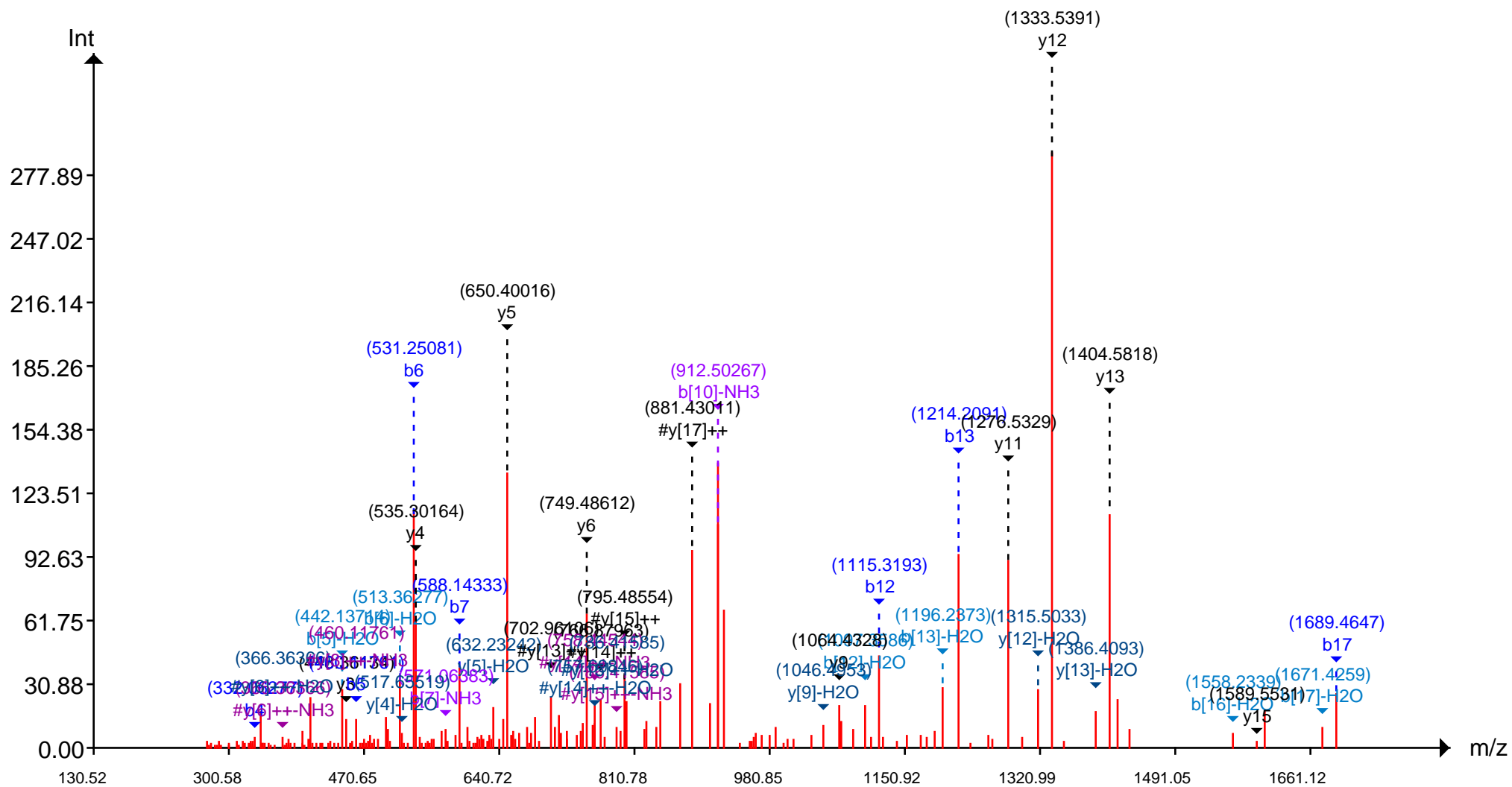
Confidence level 99.00%

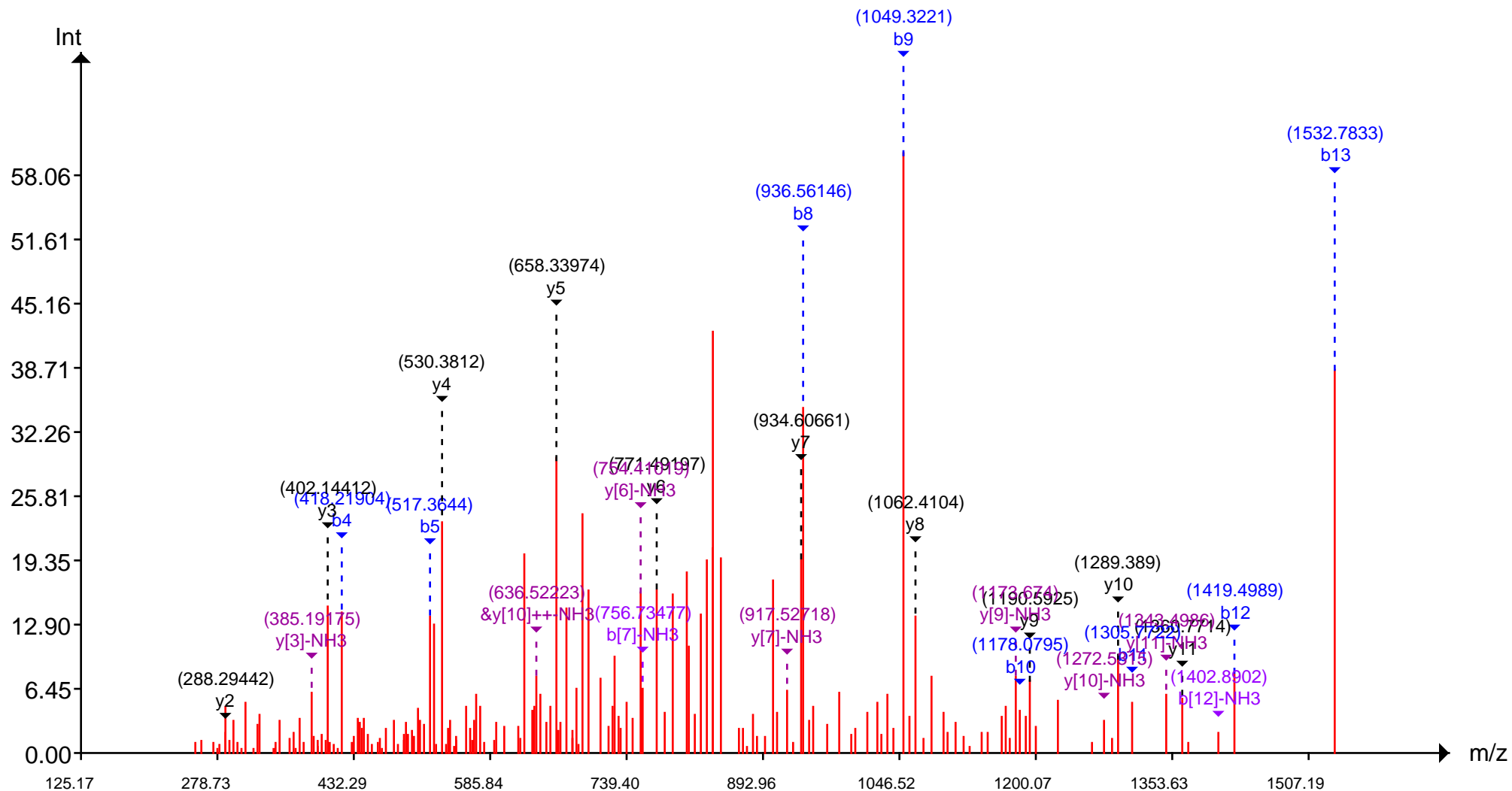
Precursor m/z 609.34227

Identity Threshold 48.51

Charge 2+

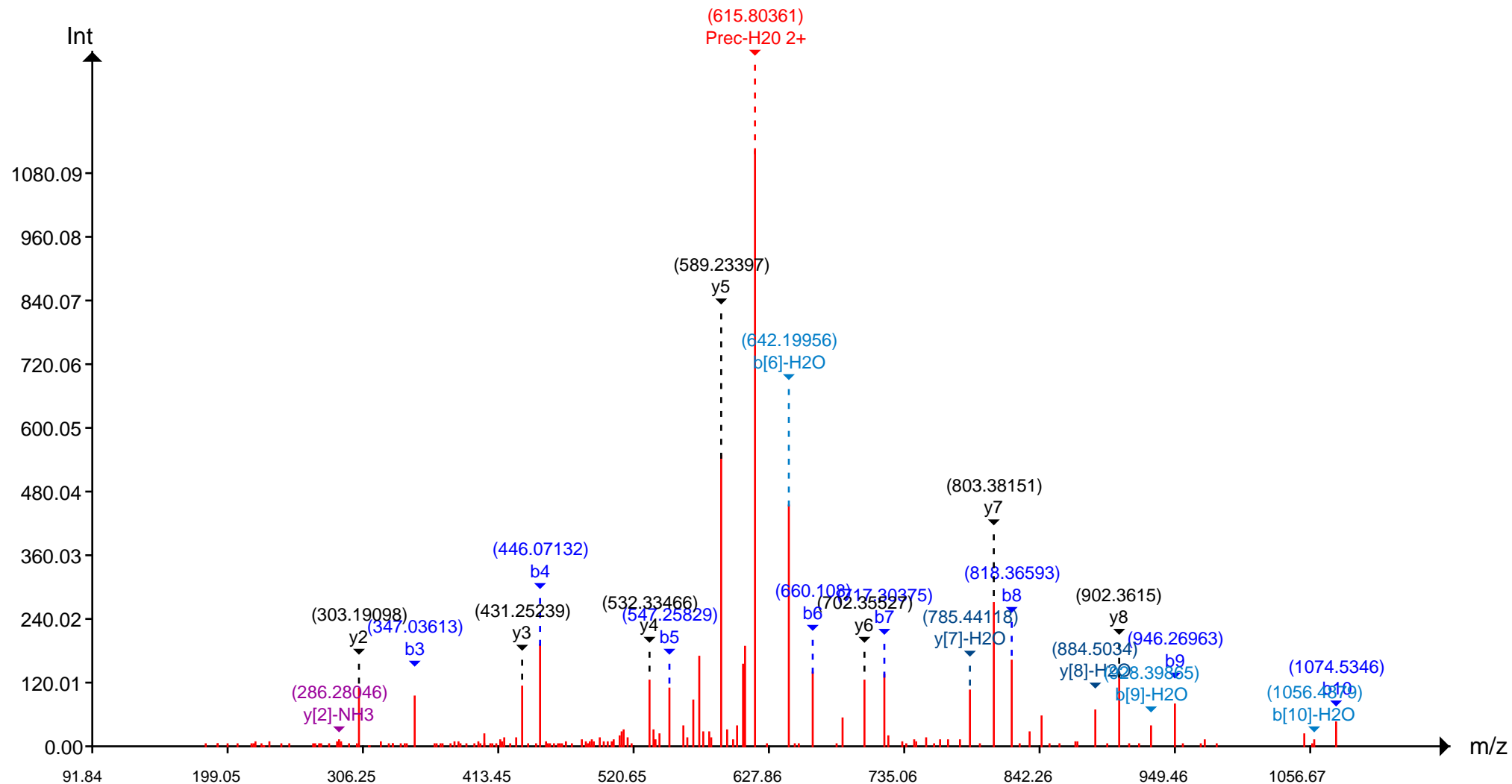
Mass error -5.14E-4





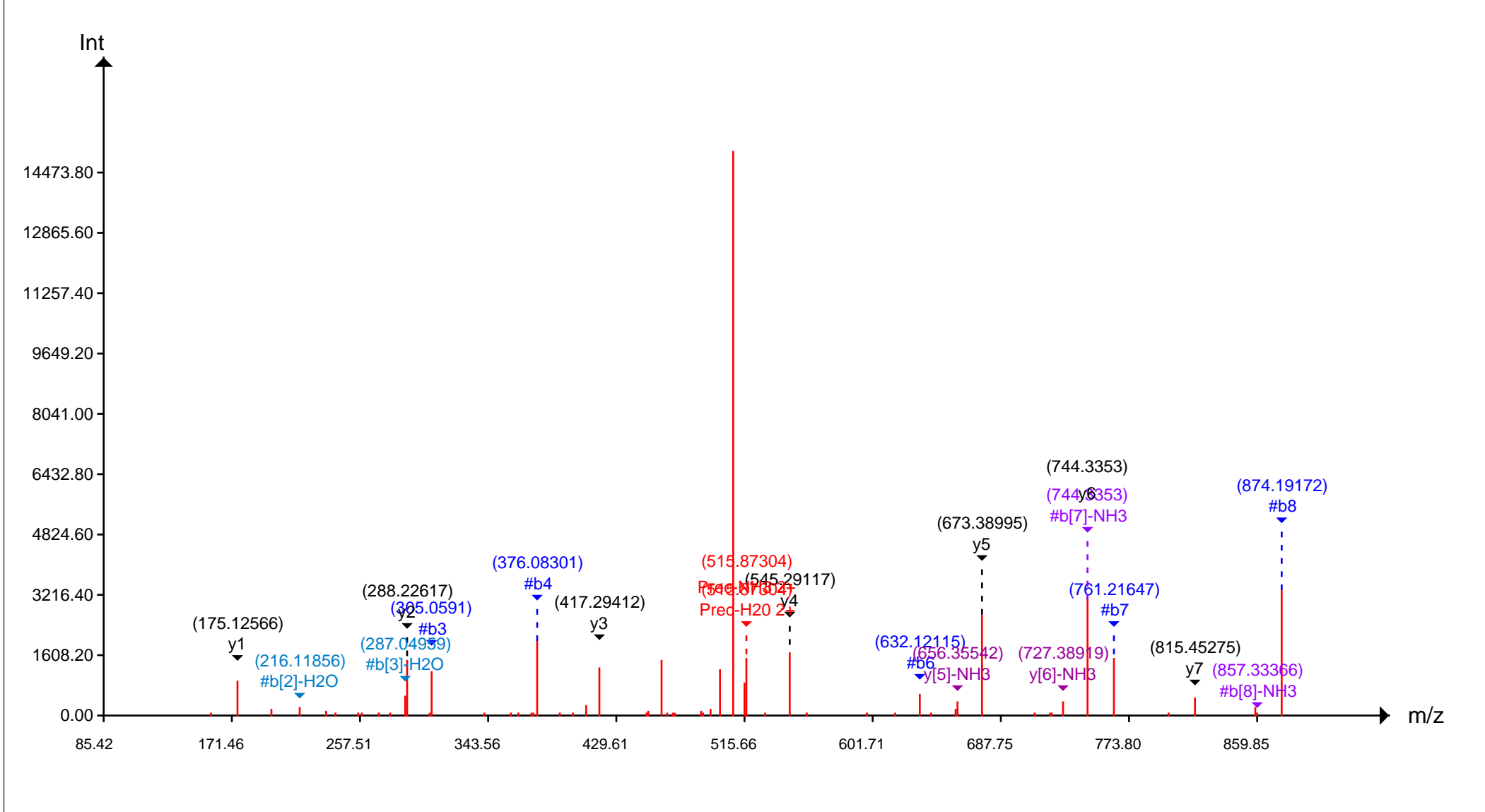
Info

| | | | |
|--------------------------|--------------------------|---------------------------|---------|
| Modified Sequence | AcD3-STLAVQQYIQQNIR-COOH | Ion score | 61.52 |
| Protein | Q9Y3A3 | Confidence level | 99.00% |
| Precursor m/z | 853.96685 | Identity Threshold | 48.57 |
| Charge | 2+ | Mass error | 1.11E-4 |

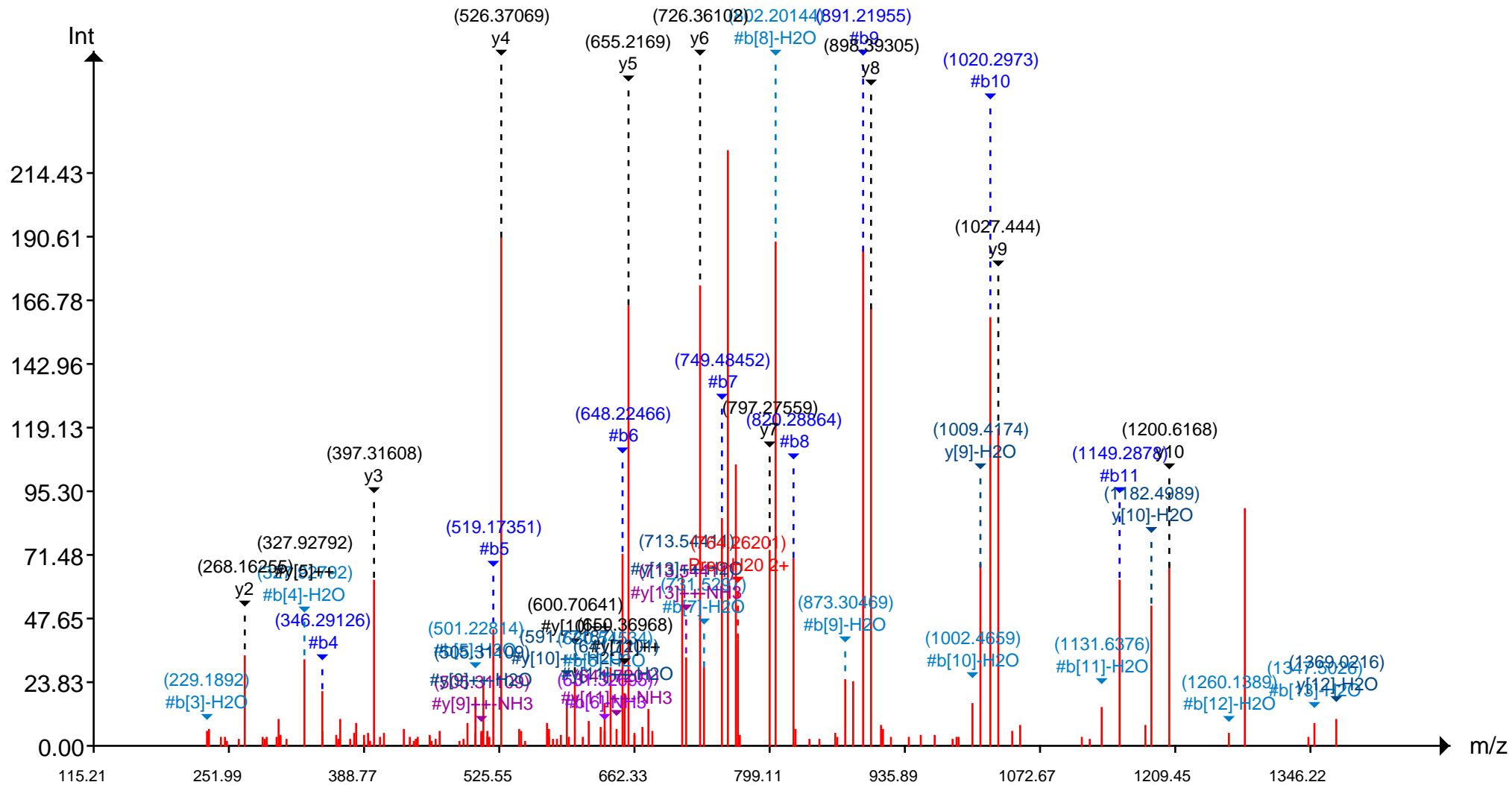


Info

| | | | |
|--------------------------|-------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-AGK<AcD3K*>VTLGTQQR-COOH | Ion score | 64.2 |
| Protein | Q08AF3 | Confidence level | 99.00% |
| Precursor m/z | 624.86245 | Identity Threshold | 47.59 |
| Charge | 2+ | Mass error | -1.16E-4 |



| Info | | | |
|--------------------------|---------------------|---------------------------|---------|
| Modified Sequence | AcD3-STAAQQELR-COOH | Ion score | 79.64 |
| Protein | P35579 | Confidence level | 99.00% |
| Precursor m/z | 524.77666 | Identity Threshold | 49.07 |
| Charge | 2+ | Mass error | -2.0E-5 |



Info

Modified Sequence AcD3-GGSVK<AcD3K*>ETAAEEESR<C13>-COOH

Protein Q7Z5W3

Precursor m/z 773.38403

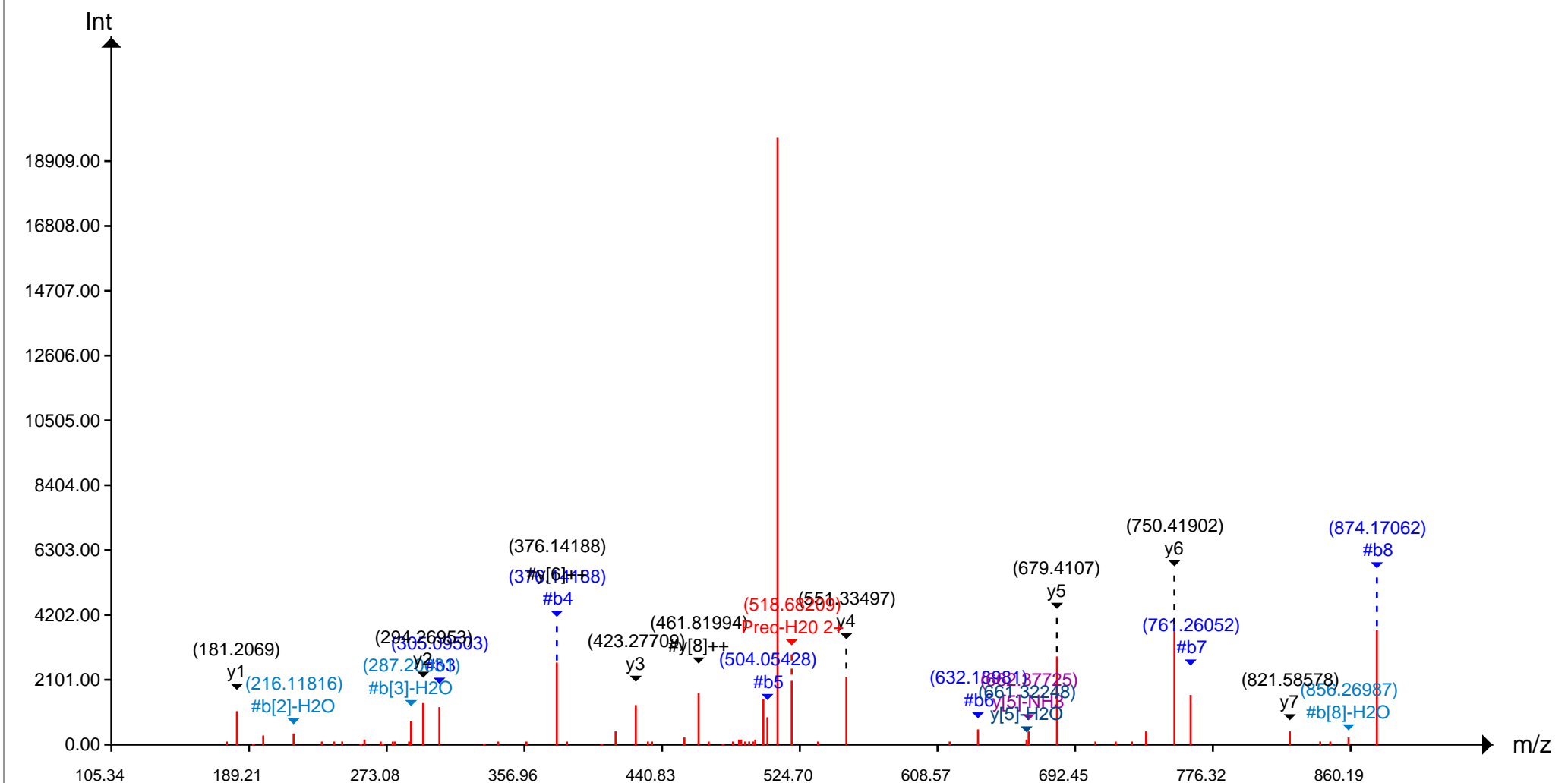
Charge 2+

Ion score 71.73

Confidence level 99.00%

Identity Threshold 48.45

Mass error 2.88E-4



Info

Modified Sequence AcD3-STAAQQELR<C13>-COOH

Protein P35579

Precursor m/z 527.78667

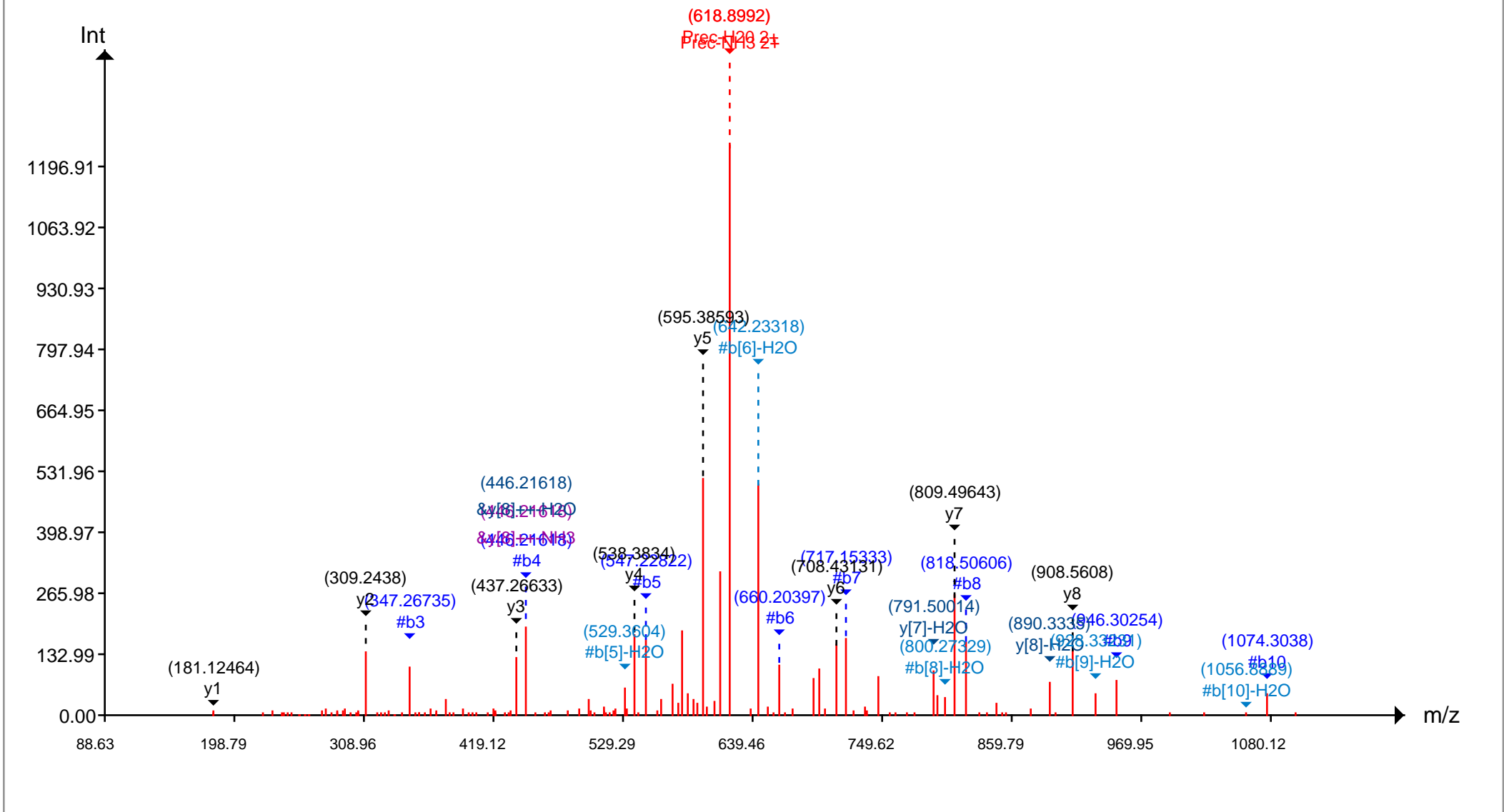
Charge 2+

Ion score 71.03

Confidence level 99.00%

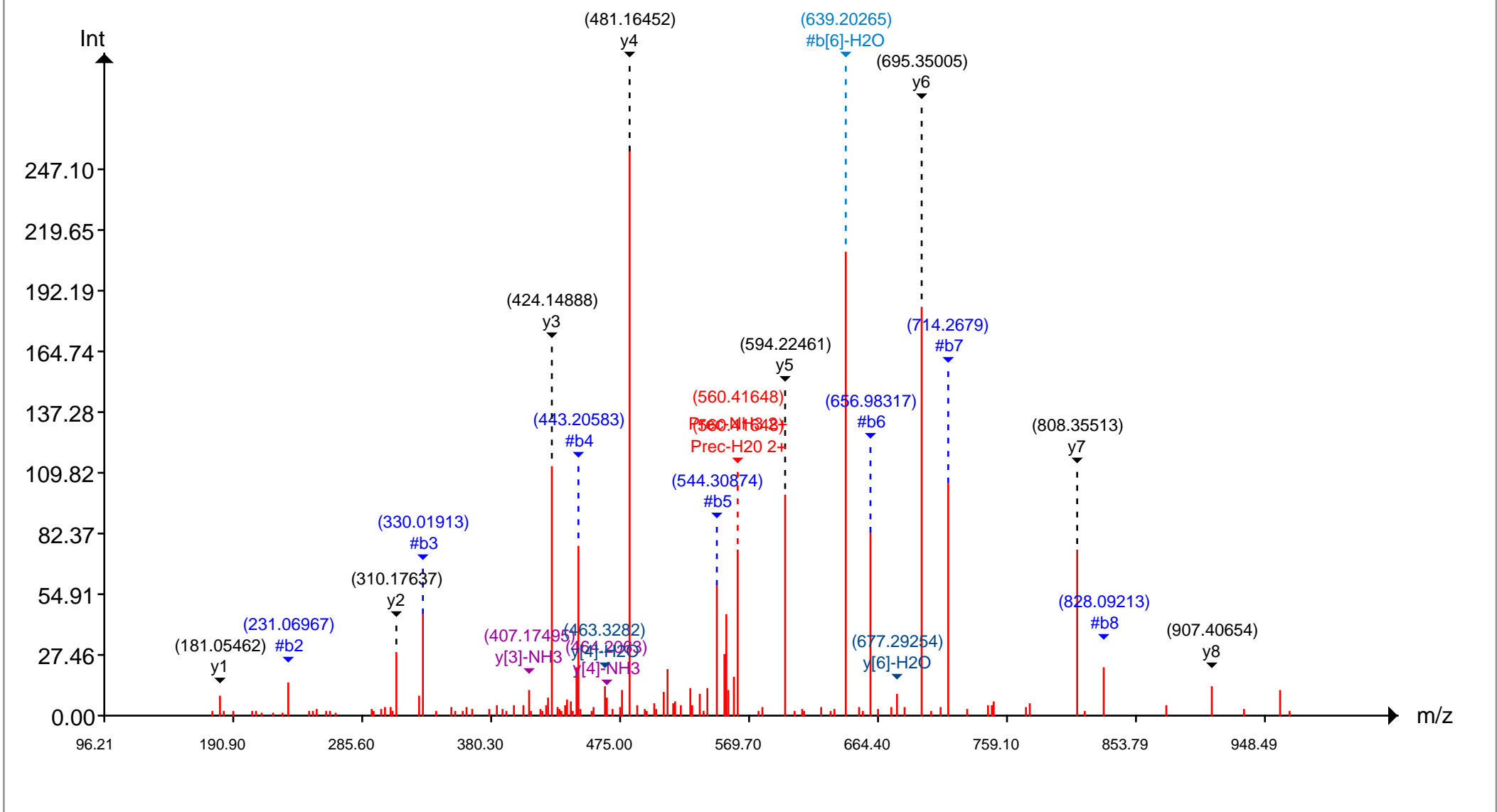
Identity Threshold 48.62

Mass error -1.41E-4



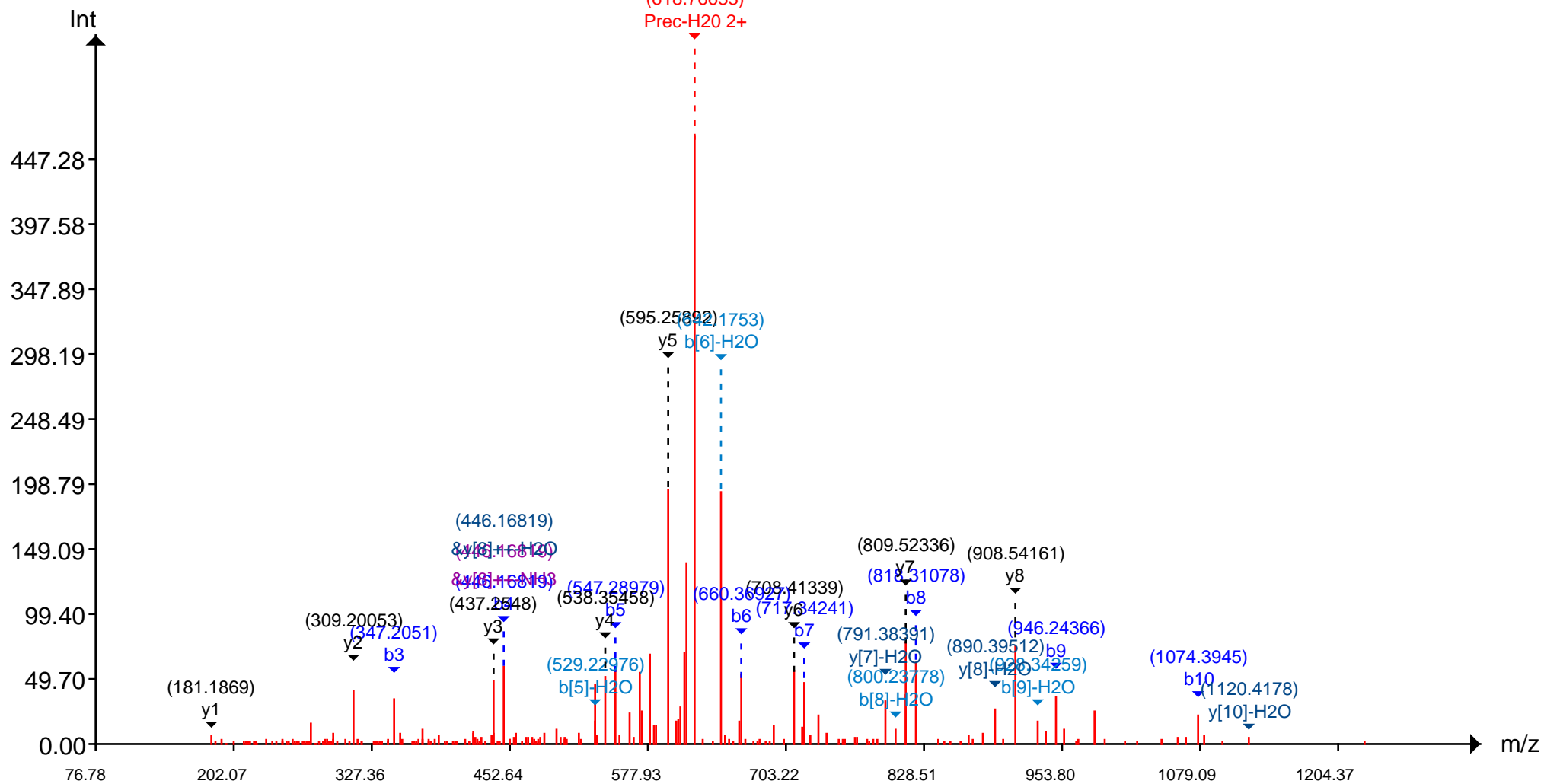
Info

| | | | |
|--------------------------|------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-AGK<AcD3K*>VTLGTQQR<C13>-COOH | Ion score | 63.96 |
| Protein | Q08AF3 | Confidence level | 99.00% |
| Precursor m/z | 627.87236 | Identity Threshold | 46.36 |
| Charge | 2+ | Mass error | -4.38E-4 |



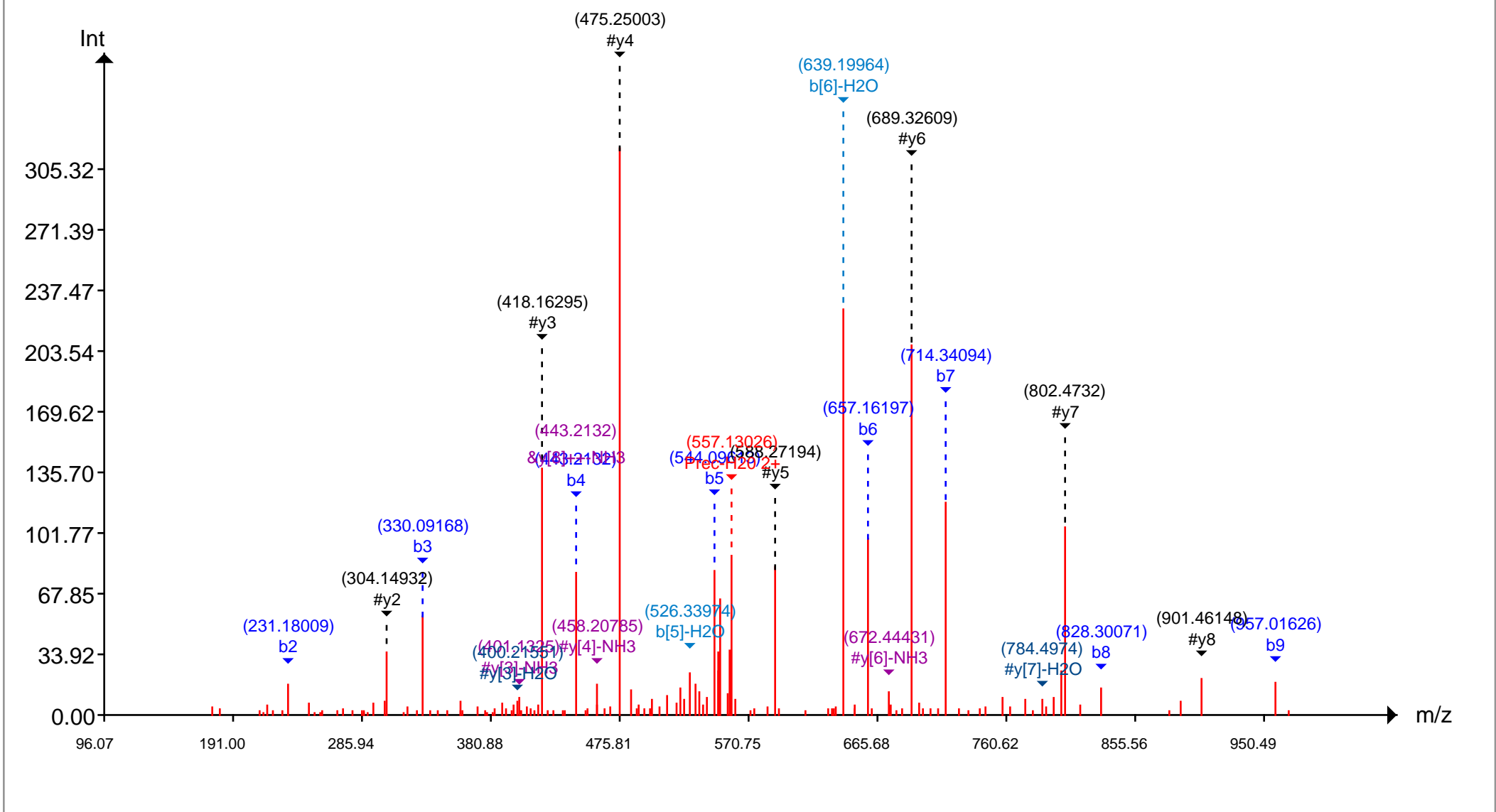
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER<C13>-COOH | Ion score | 81.1 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 569.3234 | Identity Threshold | 48.34 |
| Charge | 2+ | Mass error | -1.98E-4 |



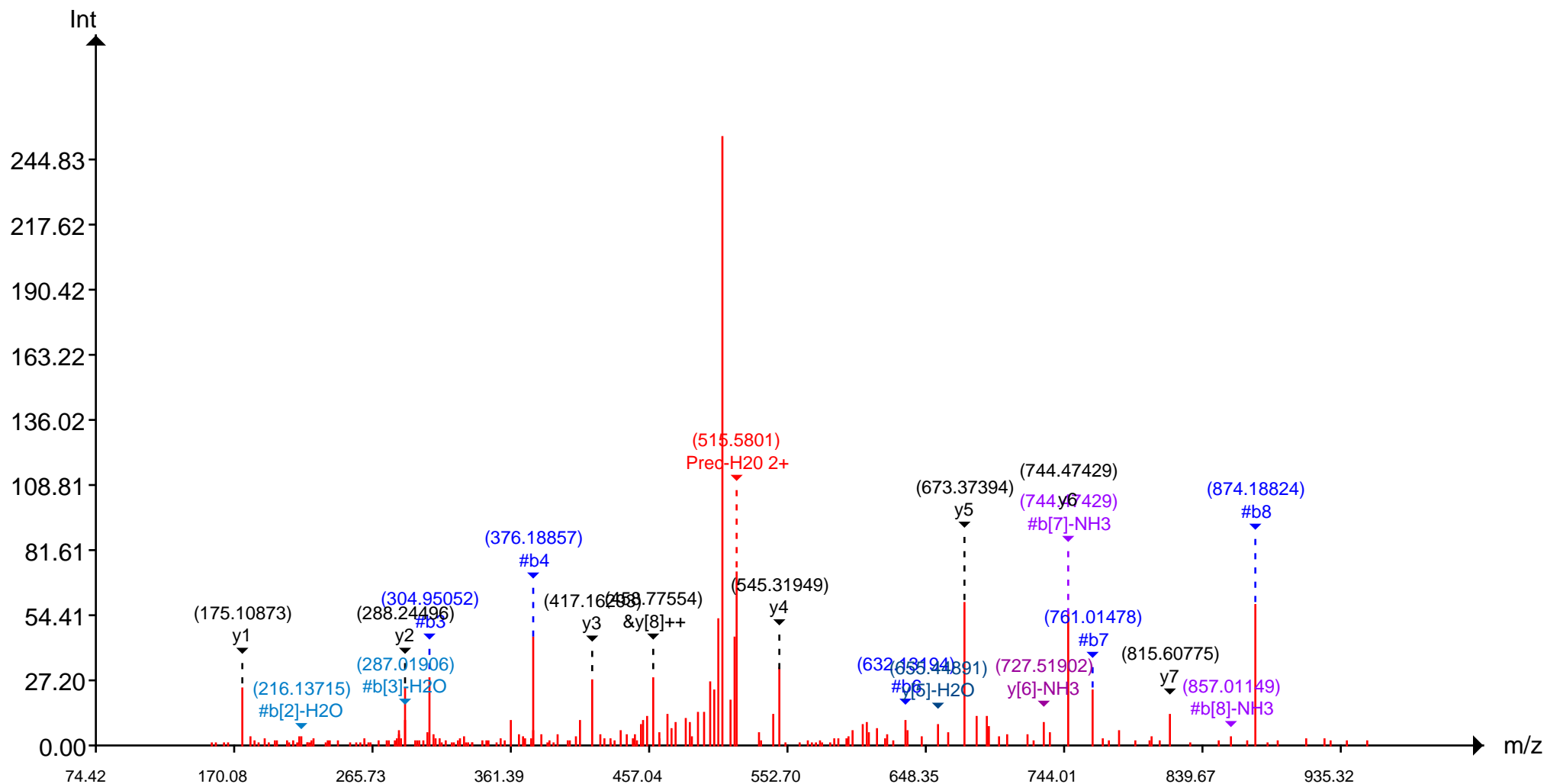
Info

| | | | |
|--------------------------|------------------------------------|---------------------------|----------|
| Modified Sequence | AcD3-AGK<AcD3K*>VTLGTQQR<C13>-COOH | Ion score | 61.82 |
| Protein | Q08AF3 | Confidence level | 99.00% |
| Precursor m/z | 627.87242 | Identity Threshold | 46.35 |
| Charge | 2+ | Mass error | -3.18E-4 |



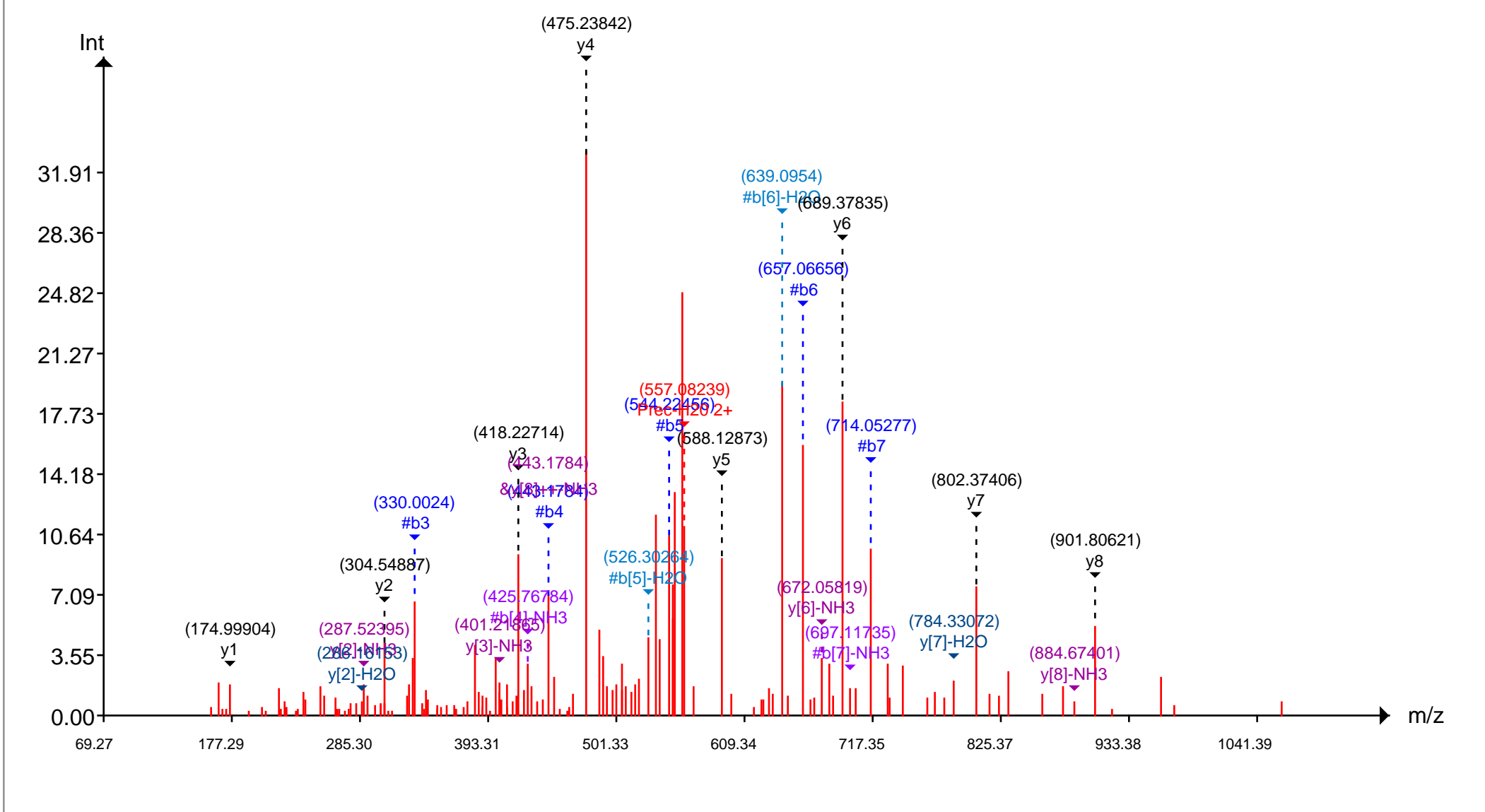
Info

| | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 67.63 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31335 | Identity Threshold | 48.86 |
| Charge | 2+ | Mass error | -1.57E-4 |



Info

| | | | |
|--------------------------|---------------------|---------------------------|--------|
| Modified Sequence | AcD3-STAAQQELR-COOH | Ion score | 55.45 |
| Protein | P35579 | Confidence level | 99.00% |
| Precursor m/z | 524.77674 | Identity Threshold | 49.05 |
| Charge | 2+ | Mass error | 1.4E-4 |



| Info | | | |
|--------------------------|--|---------------------------|----------|
| Modified Sequence | AcD3-GQVITIGNER-COOH | Ion score | 59.63 |
| Protein | P62736, Q562R1, Q9BYX7, P60709, P68032, P63261, P63267, P68133, Q6S8J3, A5A3E0 | Confidence level | 99.00% |
| Precursor m/z | 566.31336 | Identity Threshold | 48.86 |
| Charge | 2+ | Mass error | -1.37E-4 |