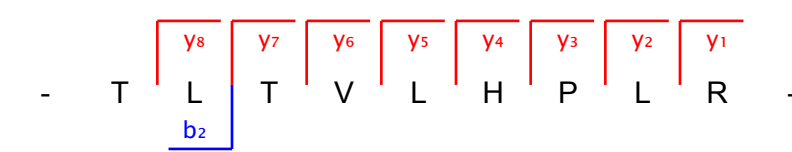
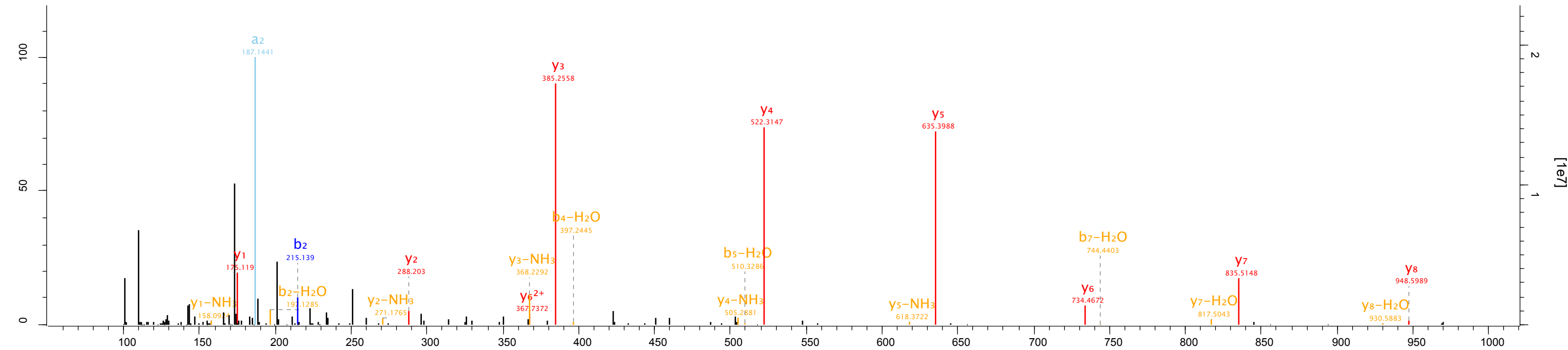
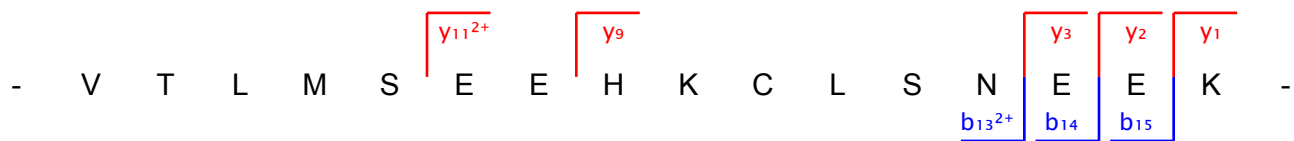
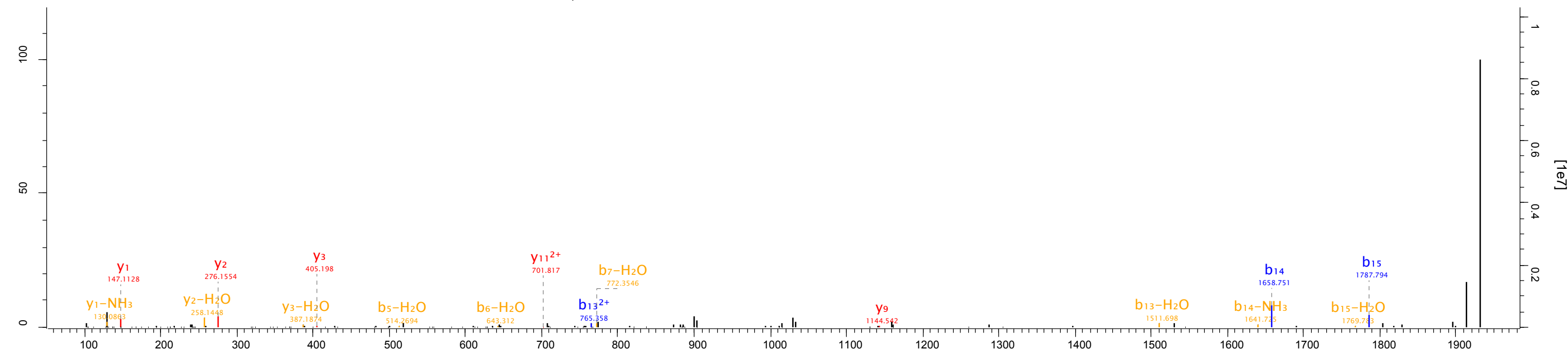


Raw file: 20130528_EXQ4_MaZe_SA_P1_11
 Scan: 36636 Method: FTMS; HCD Score: 122.52 m/z: 350.55 Gene names: Yipf5

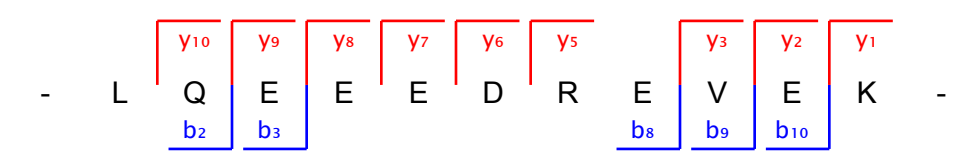
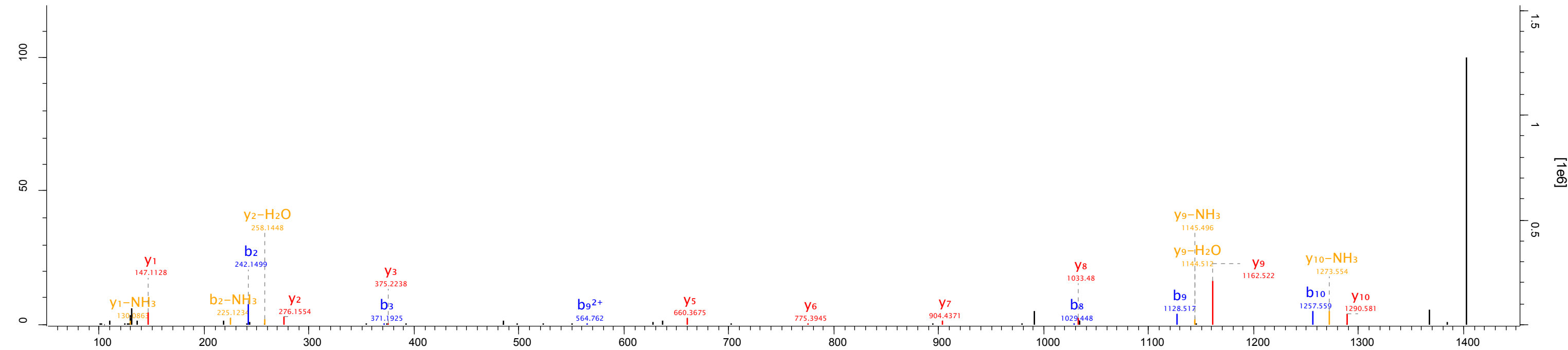


Raw file
20130529_EXQ4_MaZe_SA_P2_3

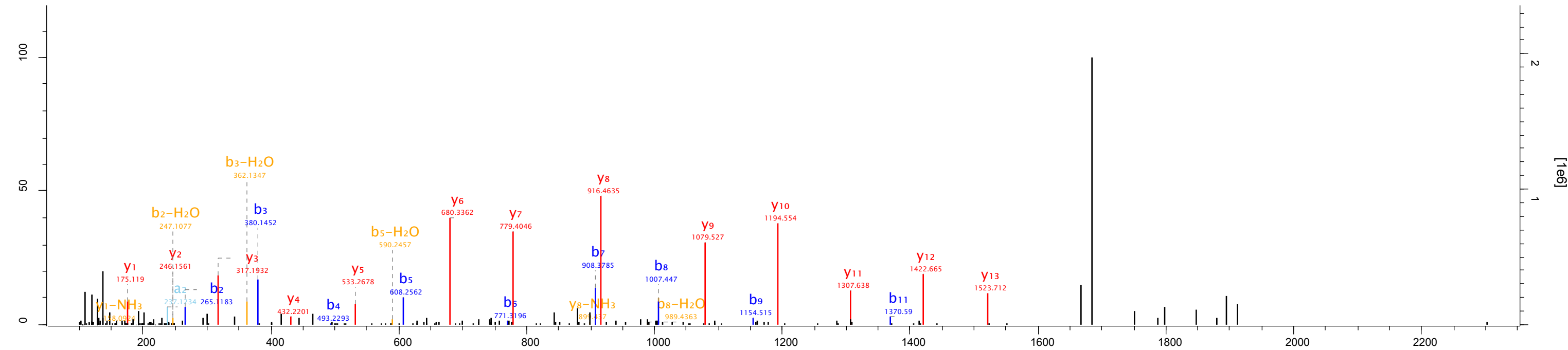
Scan 8484 Method FTMS; HCD Score 53.3 m/z 645.64 Gene names Ptprz1



Raw file: 20130529_EXQ4_MaZe_SA_P2_4
 Scan: 8698 Method: FTMS; HCD Score: 148.18 m/z: 702.34 Gene names: Rassf2



Raw file: 20130529_EXQ4_MaZe_SA_P2_4
 Scan: 47218 Method: FTMS; HCD Score: 122.63 m/z: 844.4 Gene names: Pigm



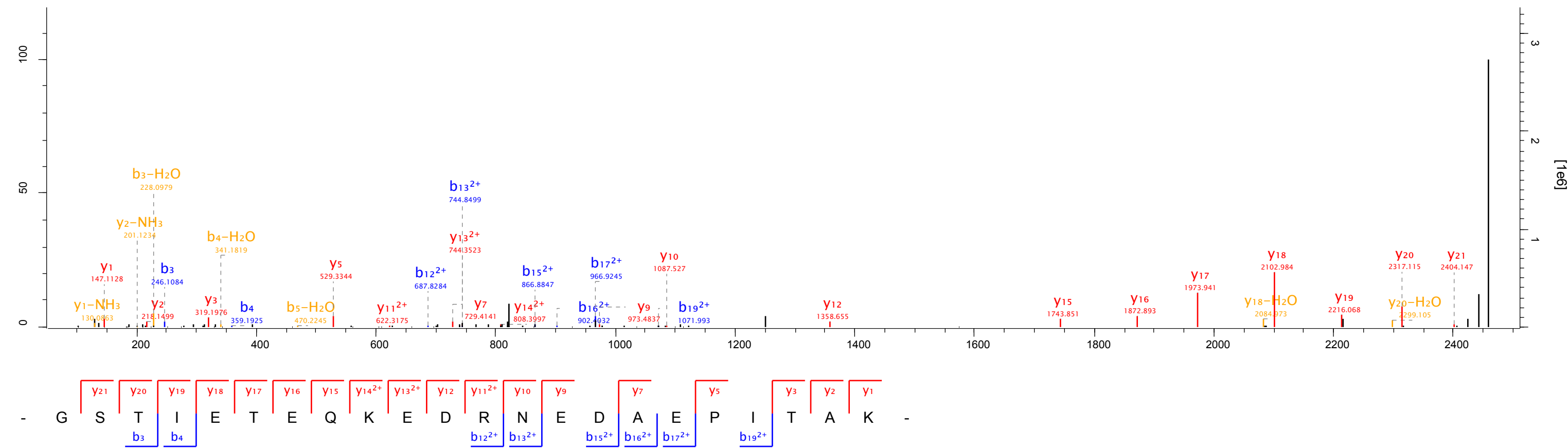
Y - T D I D Y H V F T D A A R -

Fragmentation mapping (b and y ions):

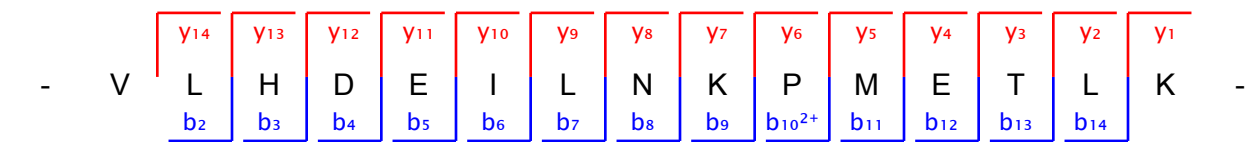
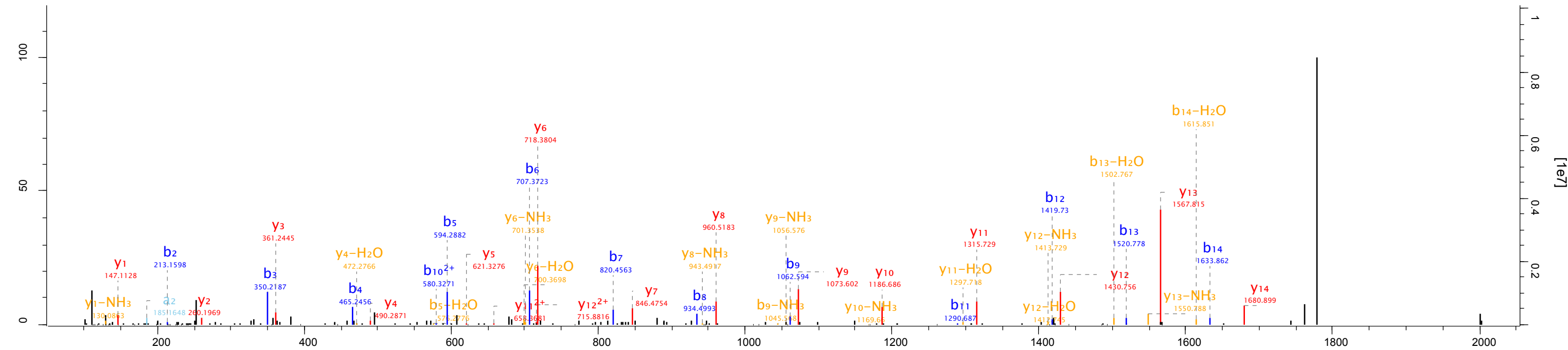
- b₂ (blue) covers T
- b₃ (blue) covers D
- b₄ (blue) covers I
- b₅ (blue) covers D
- b₆ (blue) covers Y
- b₇ (blue) covers H
- b₈ (blue) covers V
- b₉ (blue) covers F
- b₁₁ (blue) covers T
- y₁ (red) covers T
- y₂ (red) covers A
- y₃ (red) covers A
- y₄ (red) covers R
- y₅ (red) covers T
- y₆ (red) covers F
- y₇ (red) covers V
- y₈ (red) covers H
- y₉ (red) covers Y
- y₁₀ (red) covers D
- y₁₁ (red) covers I
- y₁₂ (red) covers D
- y₁₃ (red) covers T

Raw file
20130529_EXQ4_MaZe_SA_P2_8

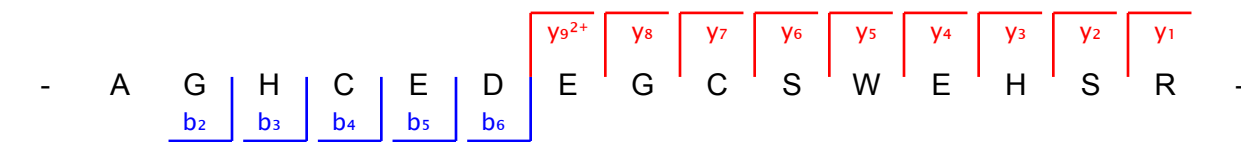
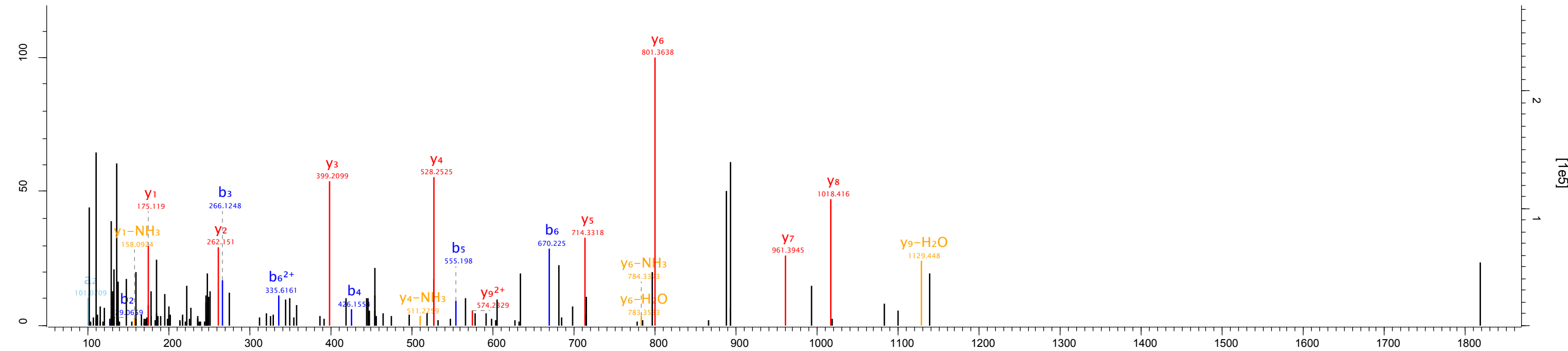
Scan 19281 Method FTMS; HCD Score 117.03 m/z 821.4 Gene names Enah



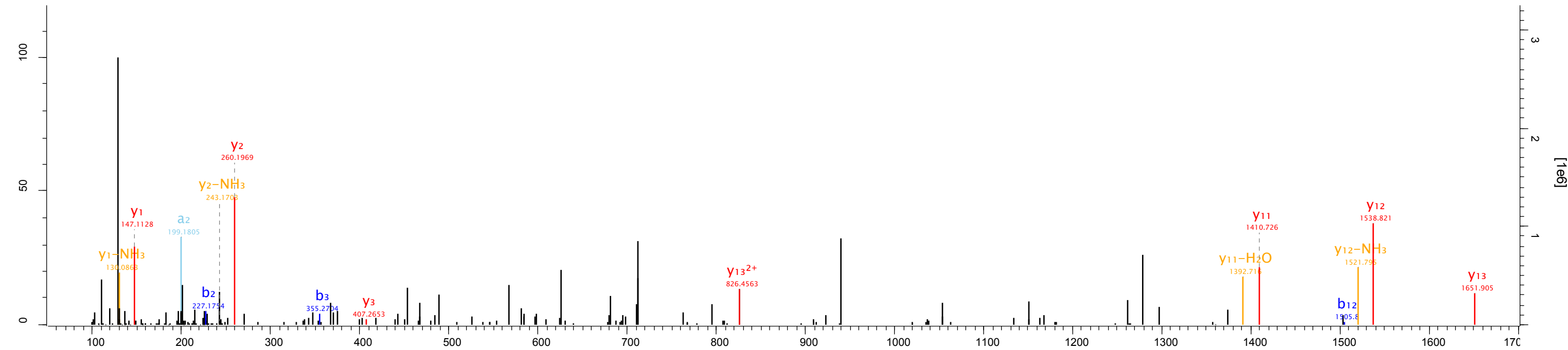
Raw file 20130529_EXQ4_MaZe_SA_P2_11 Scan 45630 Method FTMS; HCD Score 213.55 m/z 890.49 Gene names Slc35a3



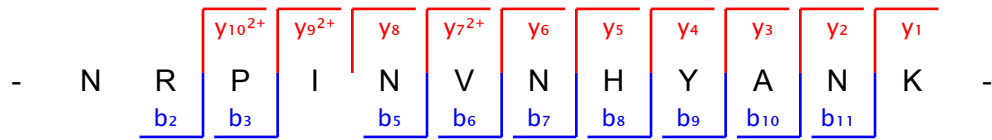
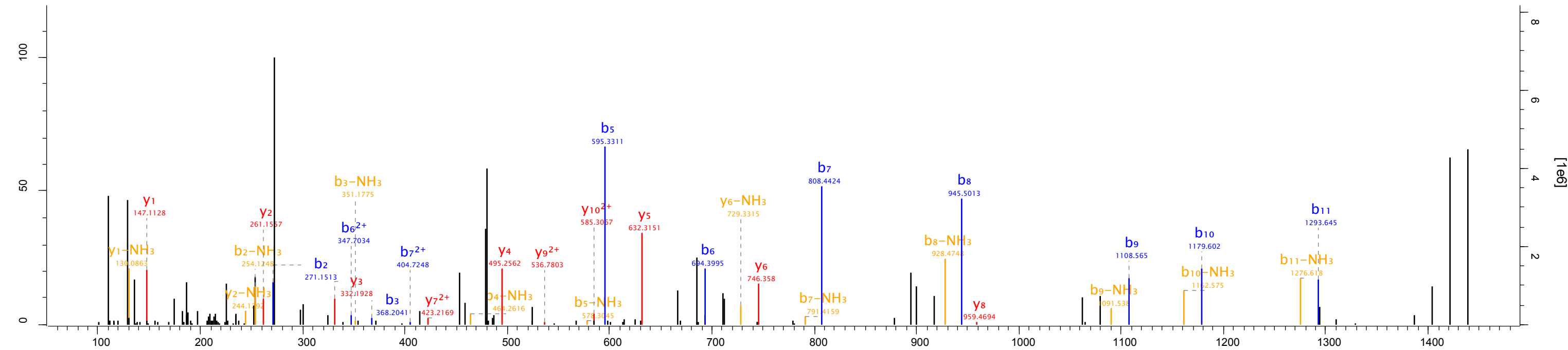
Raw file 20130531_EXQ4_MaZe_SA_P4_6 Scan 10234 Method FTMS; HCD Score 54.4 m/z 454.92 Gene names Gpr137



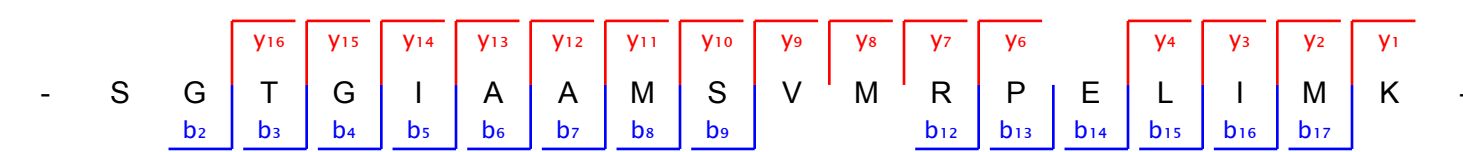
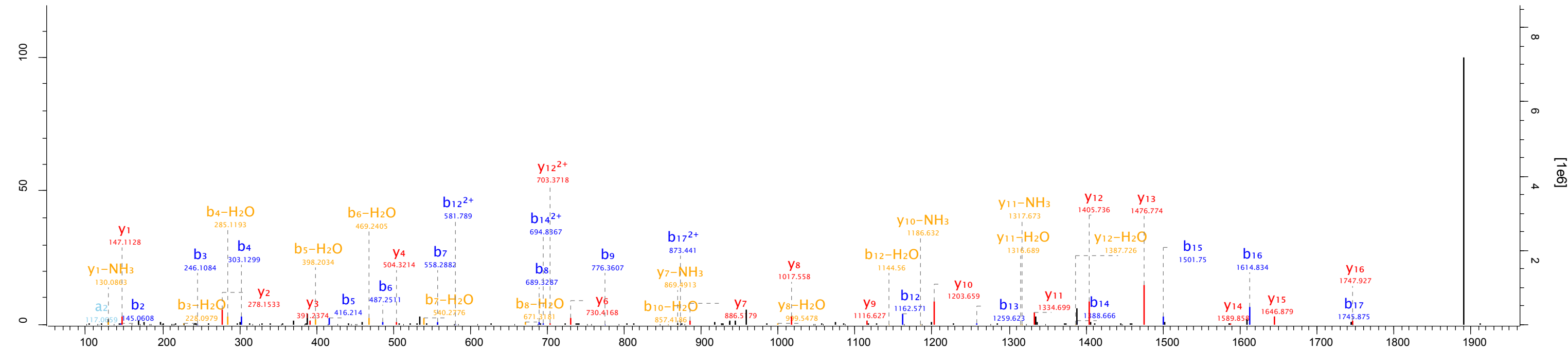
Raw file: 20130531_EXQ4_MaZe_SA_P4_6
 Scan: 49857 Method: FTMS; HCD Score: 50.9 m/z: 442 Gene names: Ppp1r9a



Raw file 20130531_EXQ4_MaZe_SA_P4_11 Scan 10202 Method FTMS; HCD Score 183.03 m/z 480.59 Gene names Ninj1

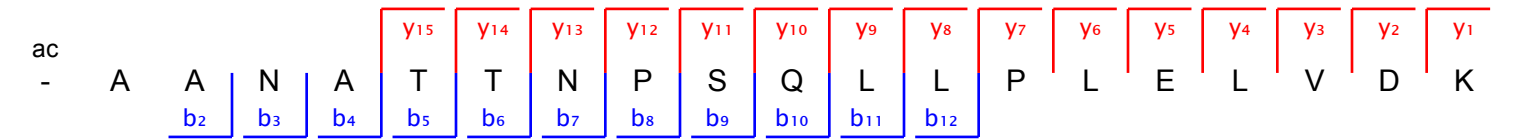
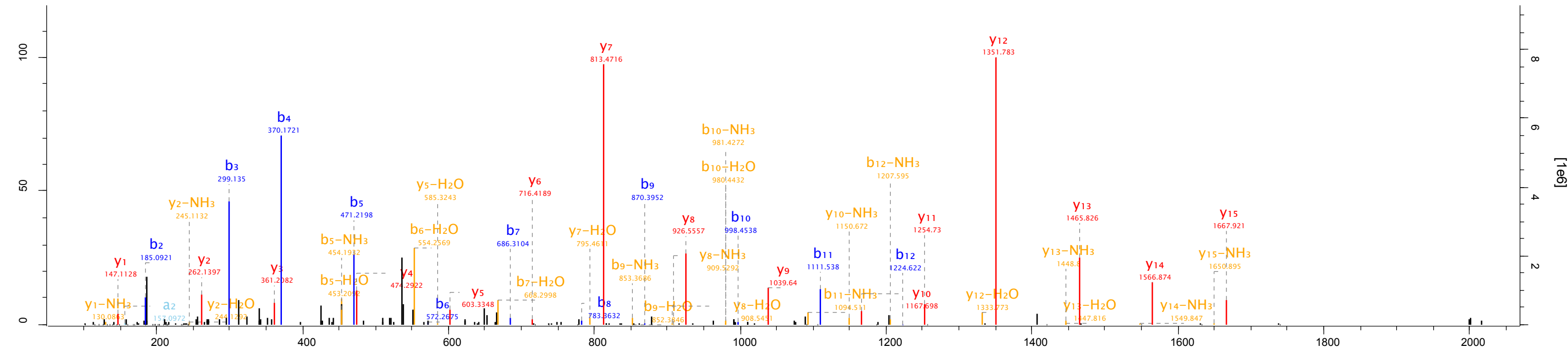


Raw file 20130531_EXQ4_MaZe_SA_P4_11 Scan 75396 Method FTMS; HCD Score 182.82 m/z 946.5 Gene names Atp6v0c

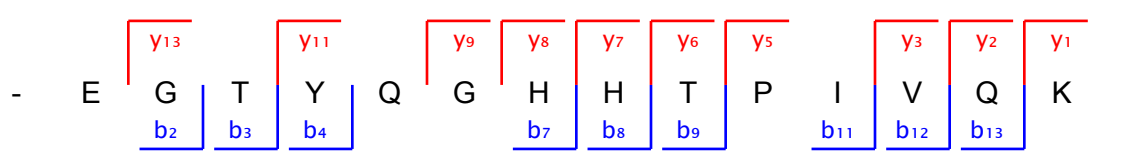
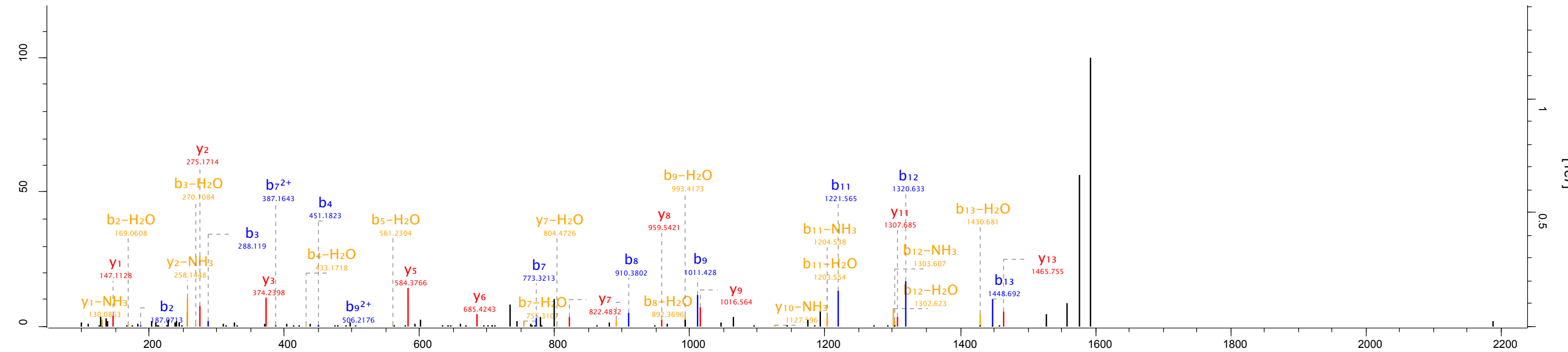


Raw file
20130602_EXQ4_MaZe_SA_M1_8

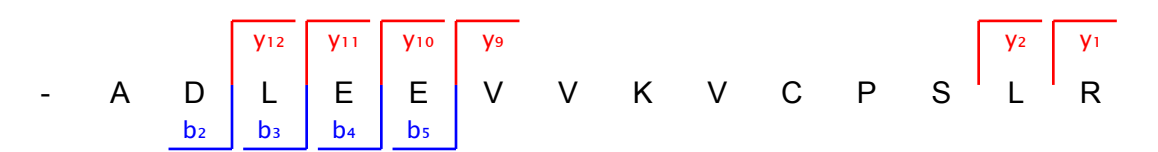
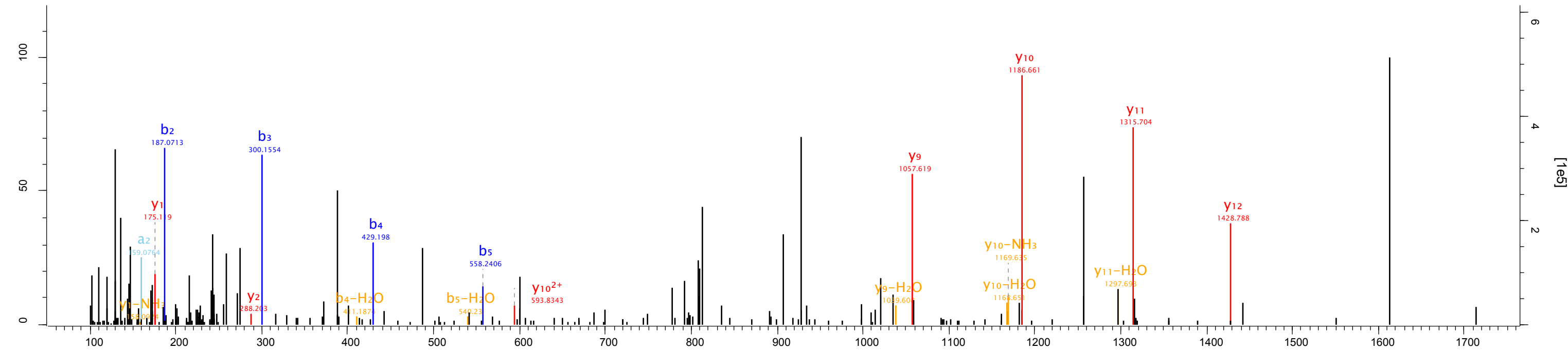
Scan 78671 Method FTMS; HCD Score 194.92 m/z 1019.55 Gene names Lsm5



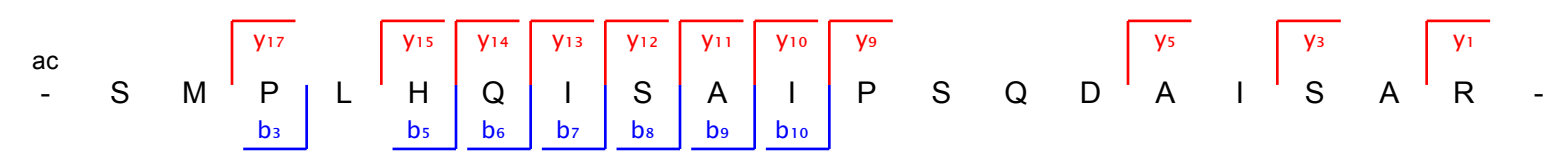
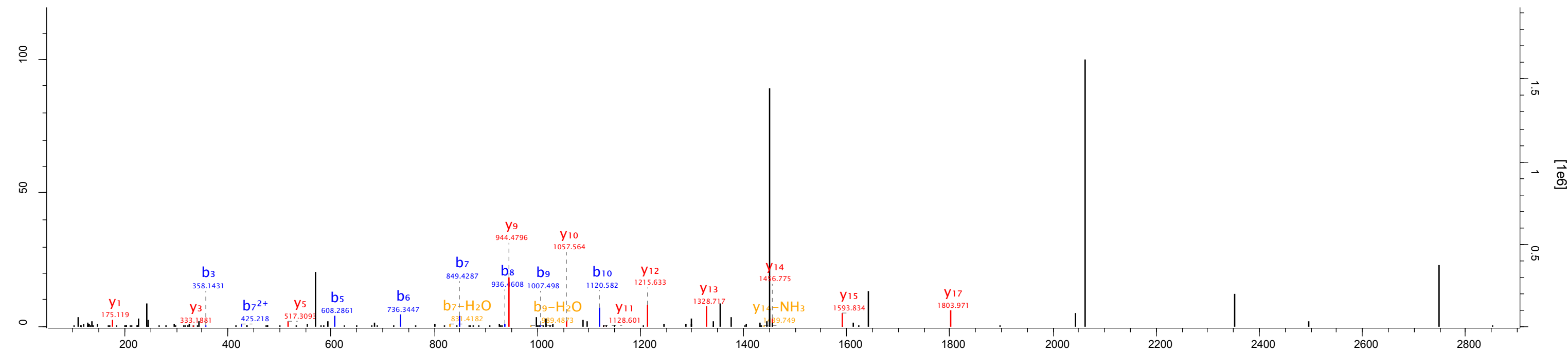
Raw file: 20130602_EXQ4_MaZe_SA_M1_11
 Scan: 12206 Method: FTMS; HCD Score: 164.93 m/z: 797.9 Gene names: mt-Co3;Mtco3



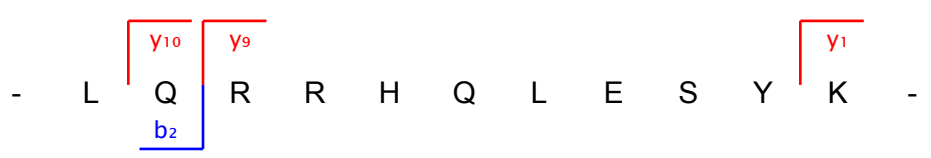
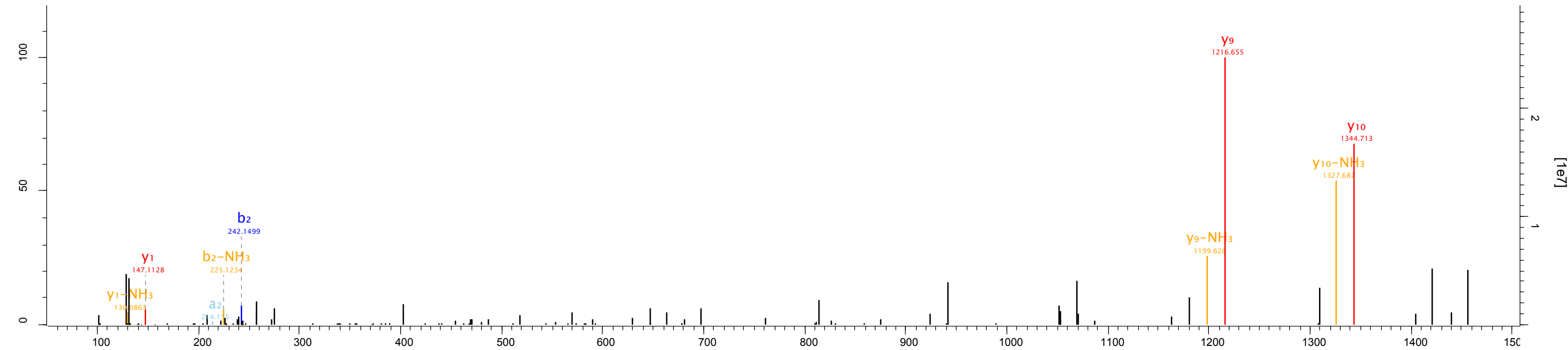
Raw file: 20130605_EXQ4_MaZe_SA_M3_3
 Scan: 40777 Method: FTMS; HCD Score: 66 m/z: 808.43 Gene names: Ttc22



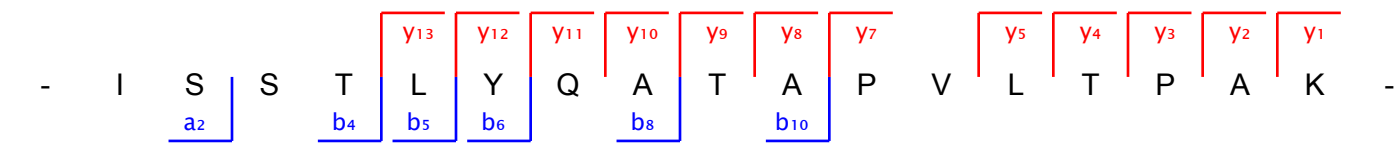
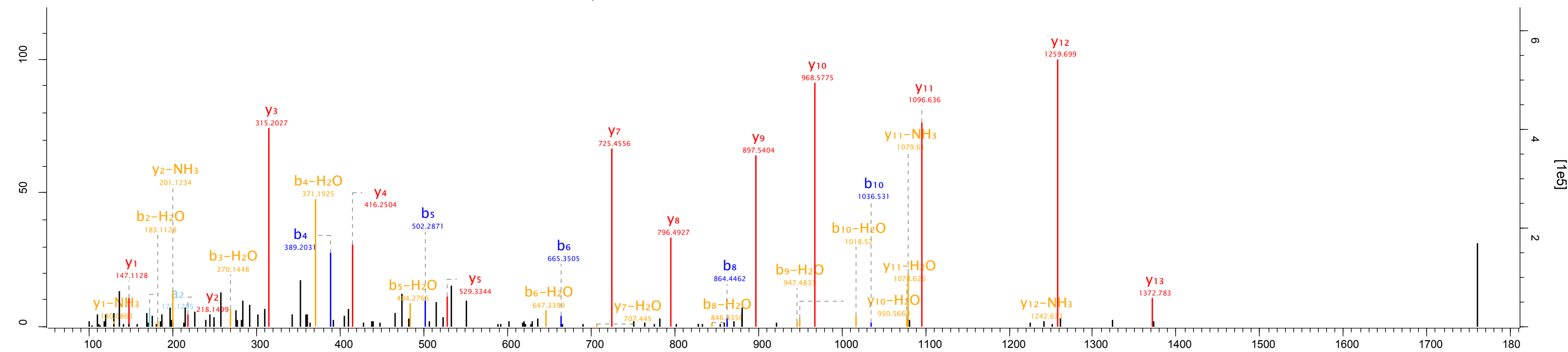
Raw file 20130605_EXQ4_MaZe_SA_M3_8 Scan 63341 Method FTMS; HCD Score 59.34 m/z 1032.15 Gene names March8



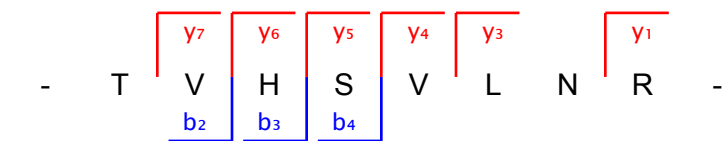
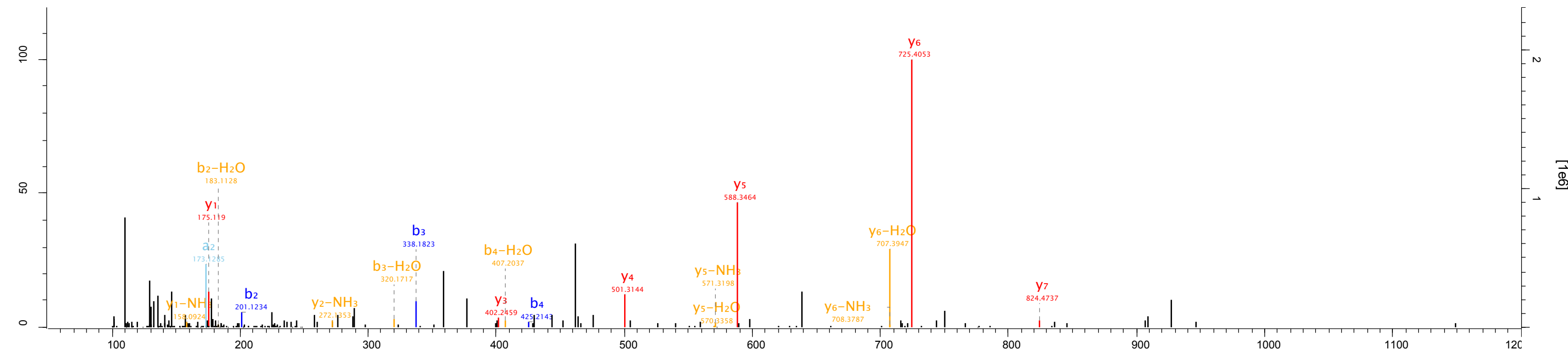
Raw file: 20130605_EXQ4_MaZe_SA_M3_11
 Scan: 4220 Method: FTMS; HCD Score: 41.4 m/z: 486.6 Gene names: Zswim4



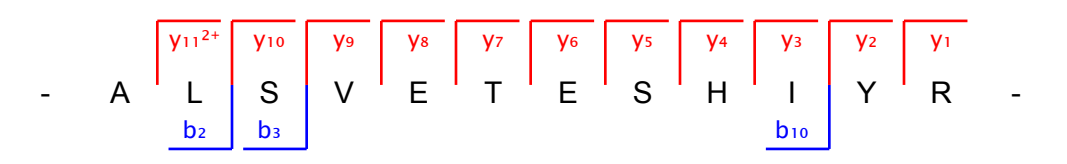
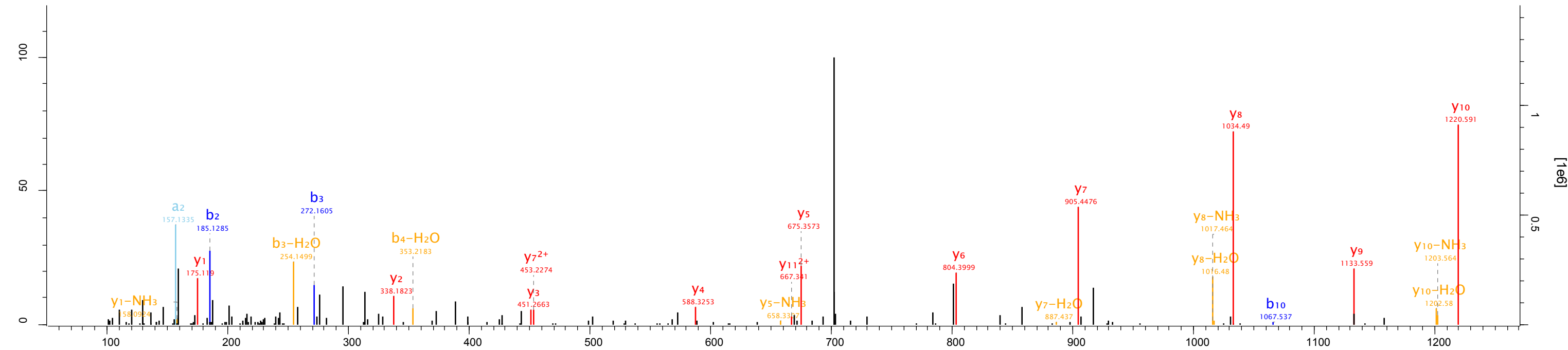
Raw file 20130605_EXQ4_MaZe_SA_M4_5 Scan 50807 Method FTMS; HCD Score 127.3 m/z 881.5 Gene names Krtcap2



Raw file 20130605_EXQ4_MaZe_SA_M4_11 Scan 8738 Method FTMS; HCD Score 84.74 m/z 463.26 Gene names Galnt10



Raw file 20130628_EXQ4_MaZe_SA_P4_11 Scan 34876 Method FTMS; HCD Score 132.32 m/z 702.86 Gene names Cyb561d2



Raw file 20130628_EXQ4_MaZe_SA_P4_11 Scan 37623 Method FTMS; HCD Score 123.75 m/z 403.88 Gene names Ndufb2

