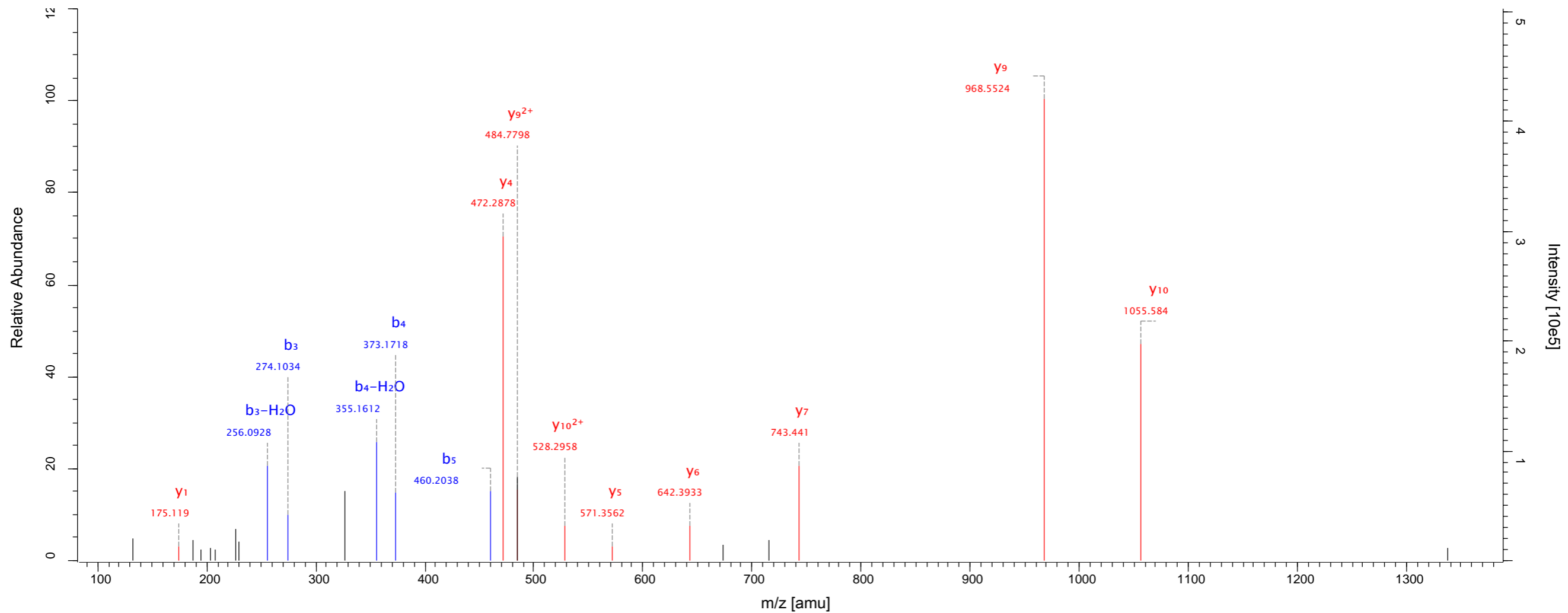
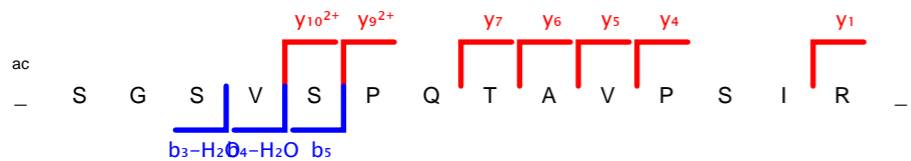
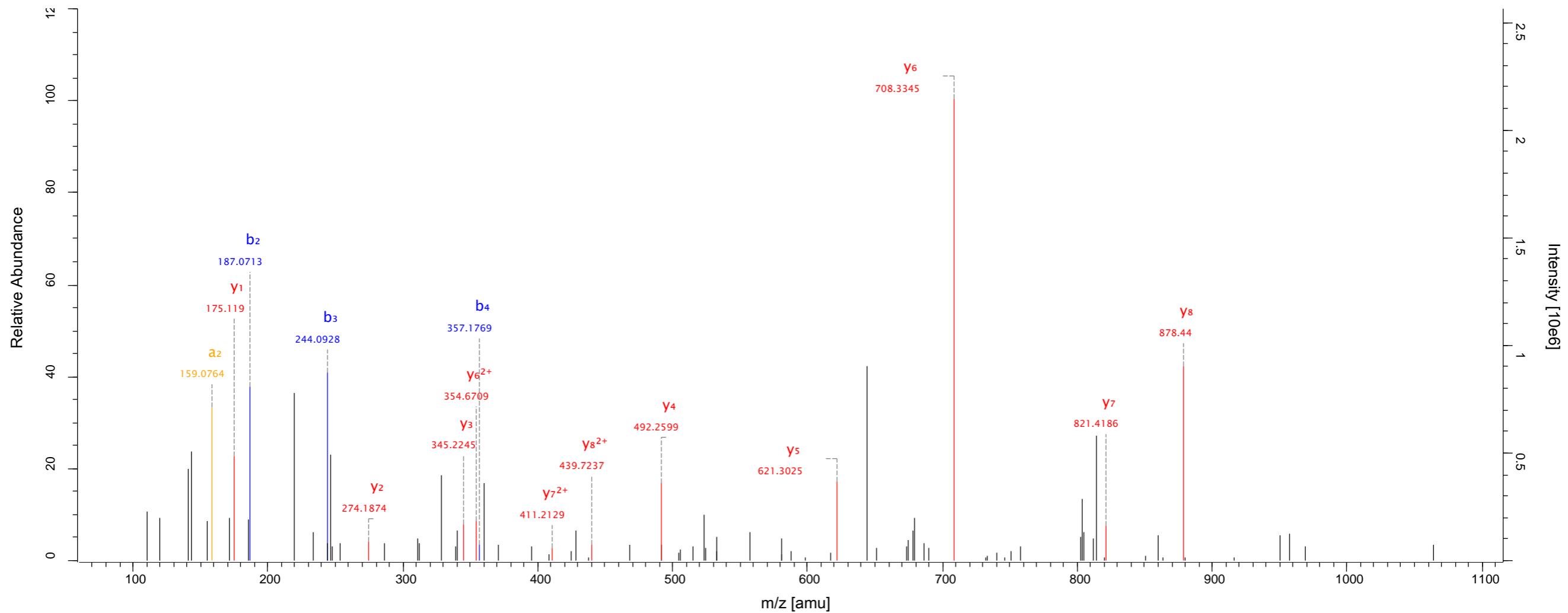
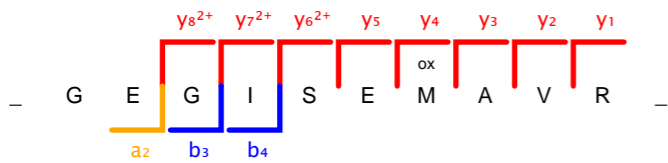


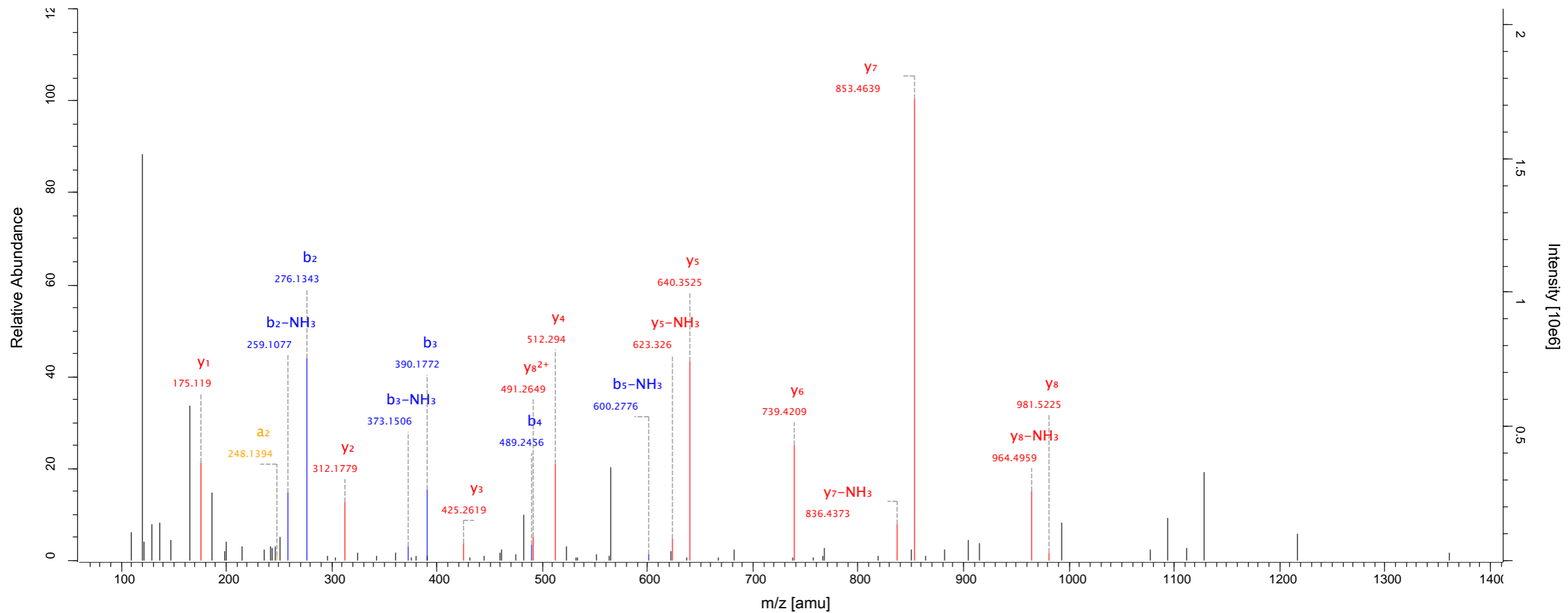
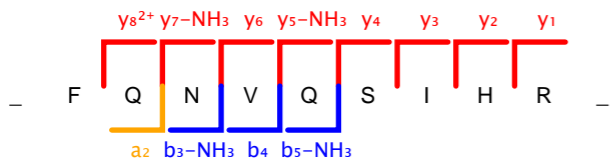
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_1
Scan number: 27670
Protein: ORFS334W + NC_006273.2[204331,204558]+ <unknown description>
Peptide score: 84.72
Method: FTMS; HCD



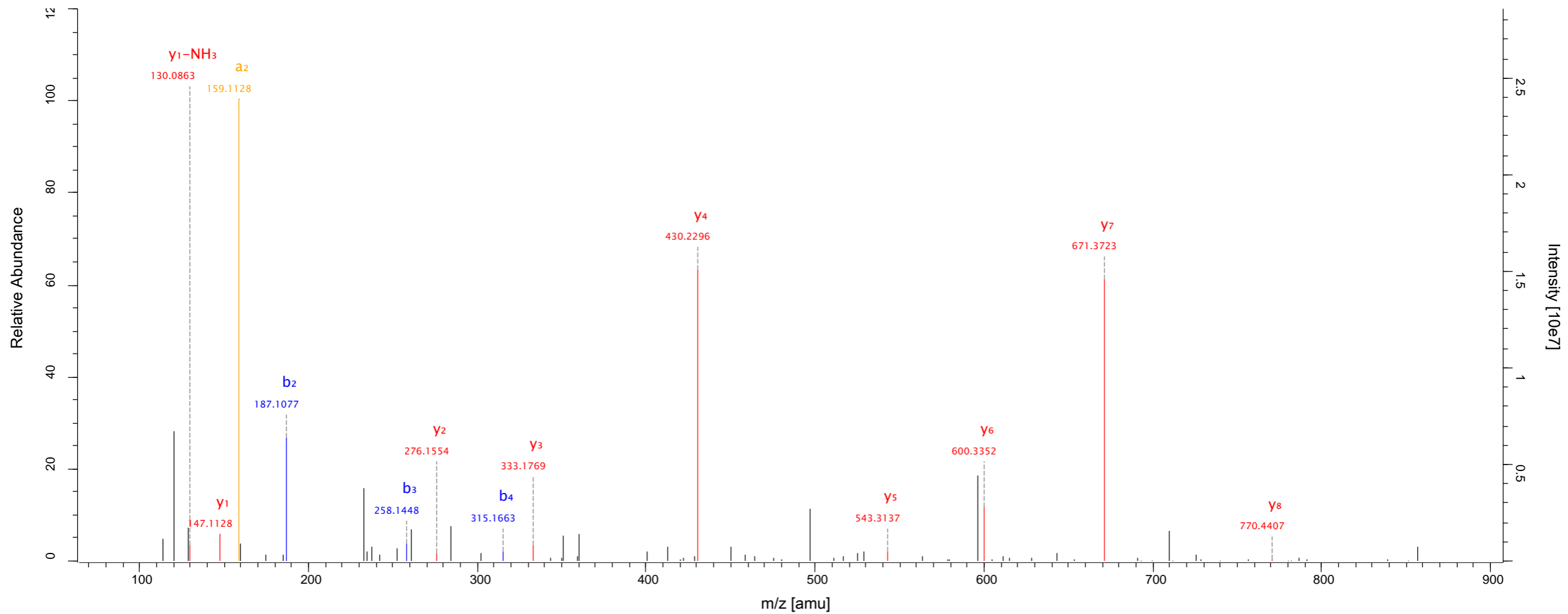
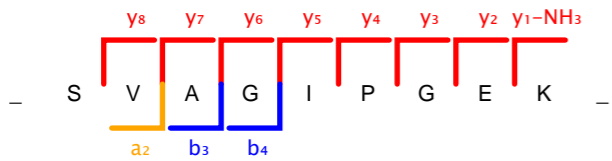
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 10249
Protein: ORFL18W + NC_006273.2[8020,8253]+ <unknown description>
Peptide score: 89.05
Method: FTMS; HCD



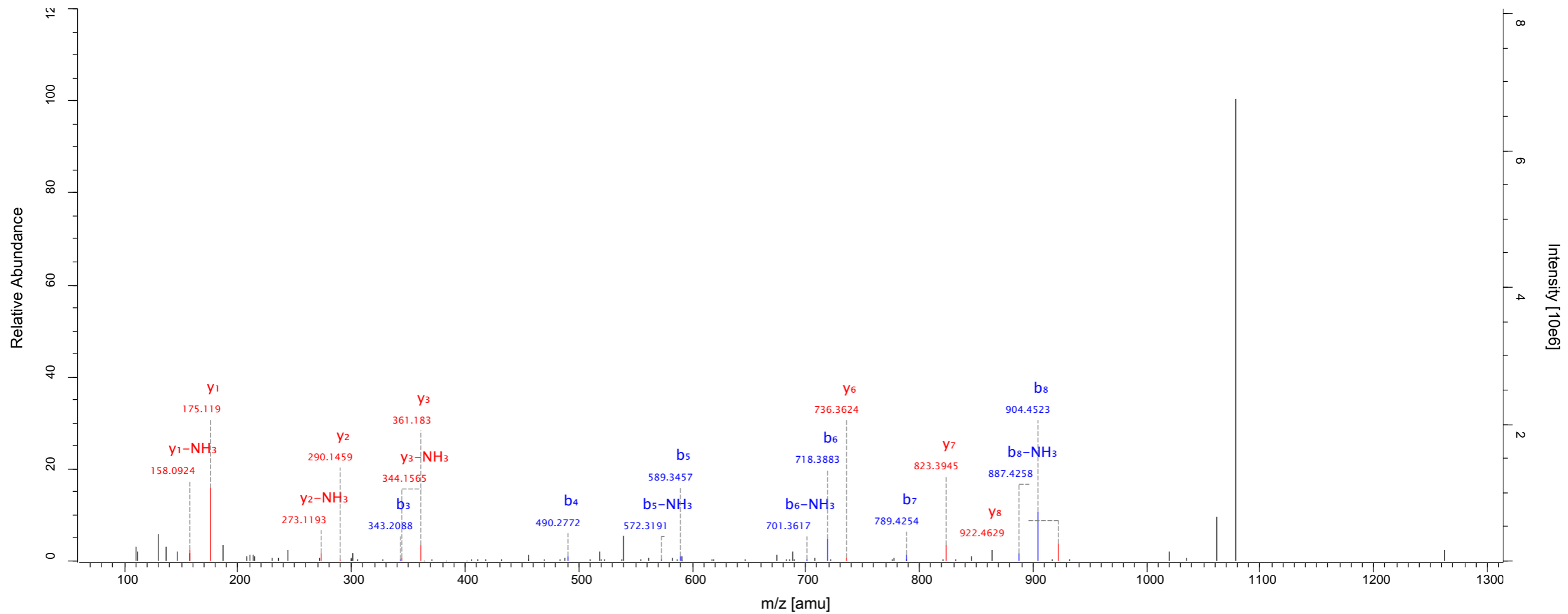
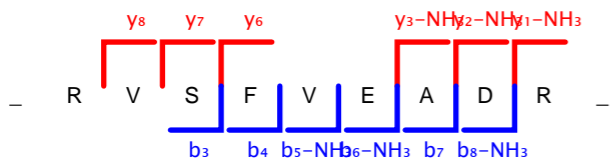
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 10712
Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
Peptide score: 155.1
Method: FTMS; HCD



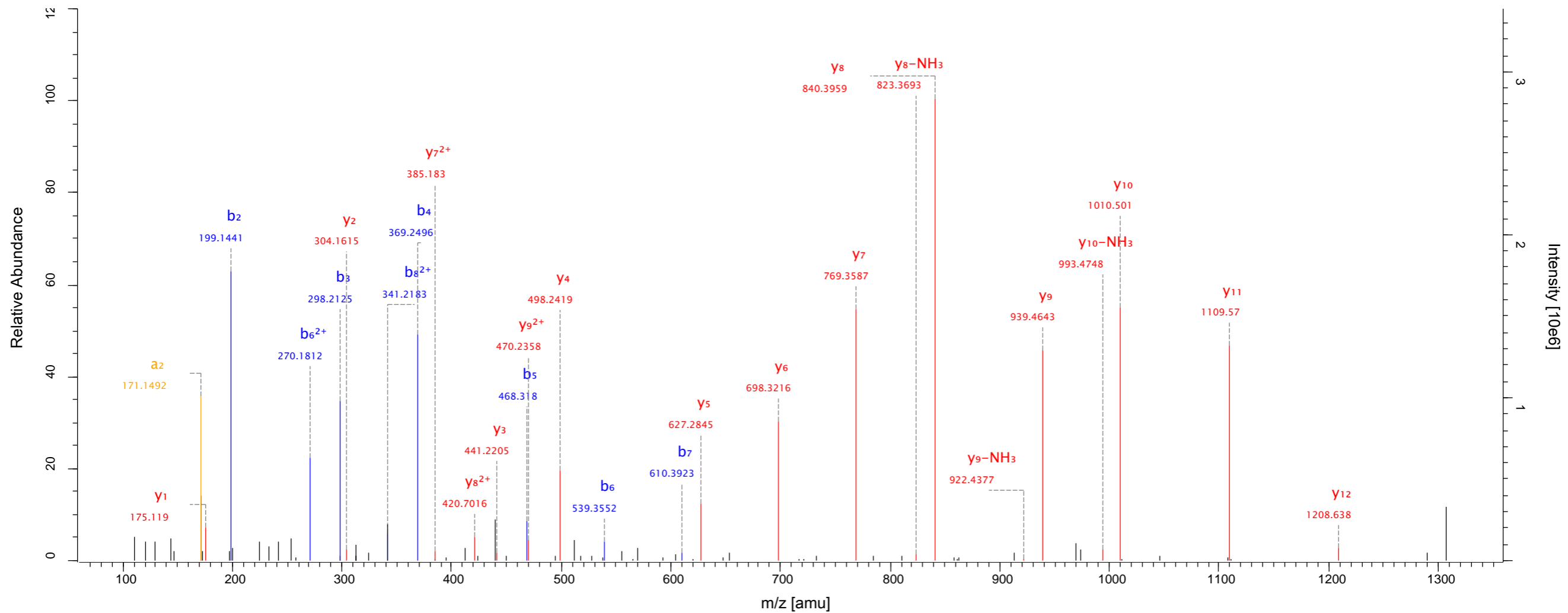
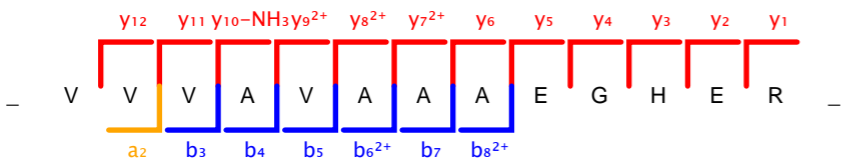
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_10
Scan number: 11413
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 64.71
Method: FTMS; HCD



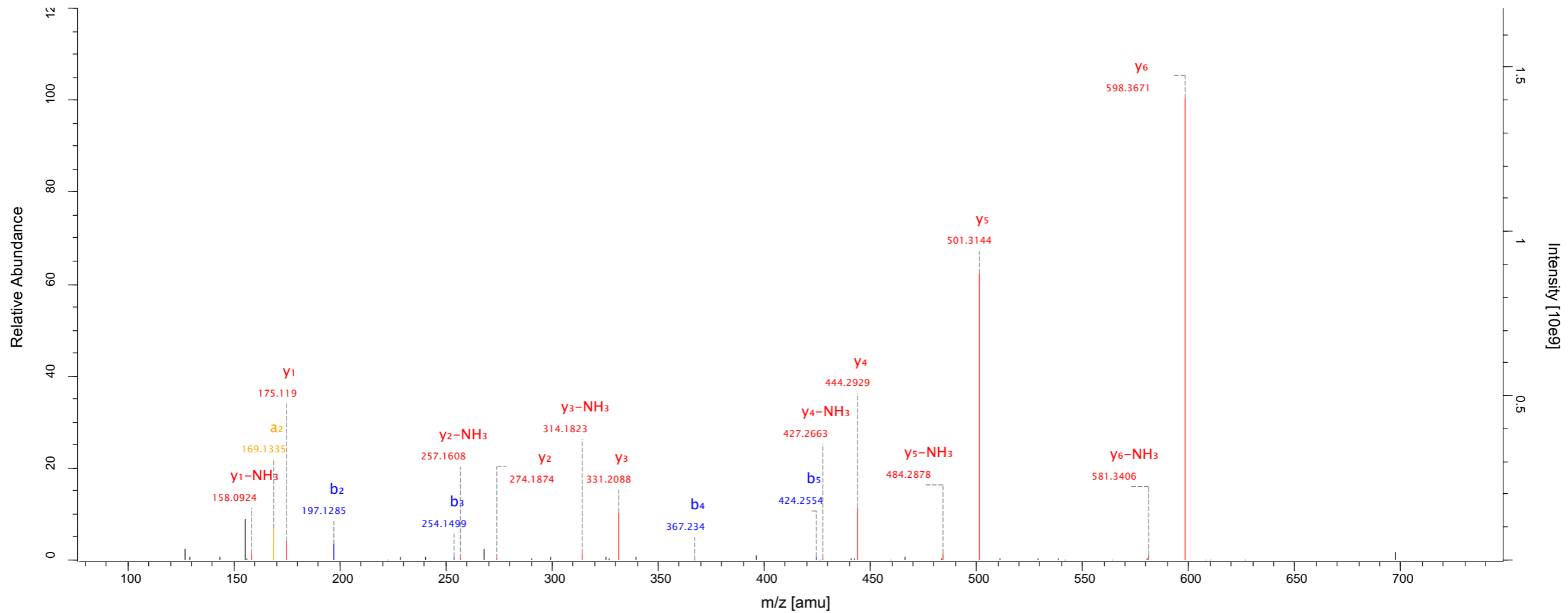
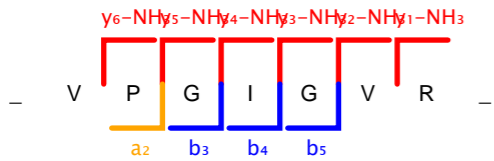
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 11896
Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>; ORFL27C.iORF1 - NC_006273.2[7912,8184]- <unknown description>
Peptide score: 93.56
Method: FTMS; HCD



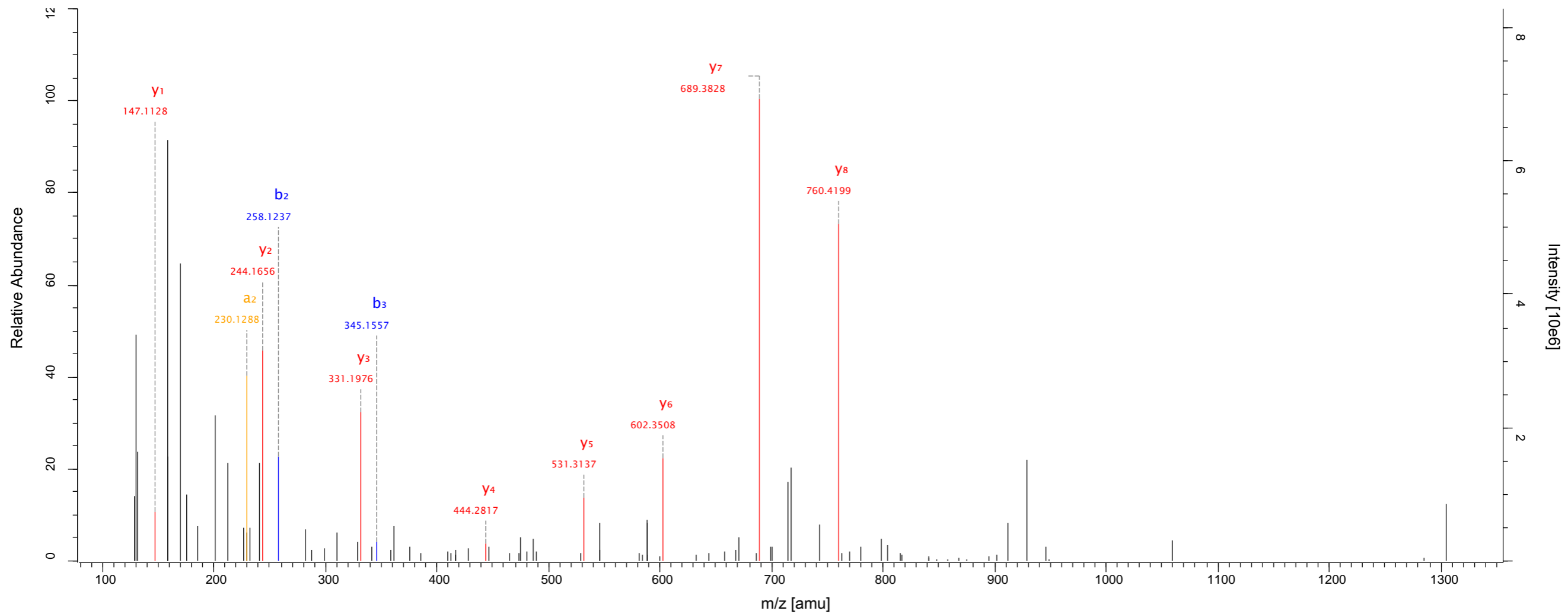
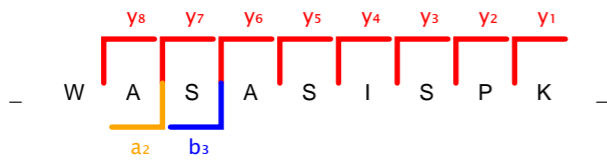
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 13953
Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
Peptide score: 195.51
Method: FTMS; HCD



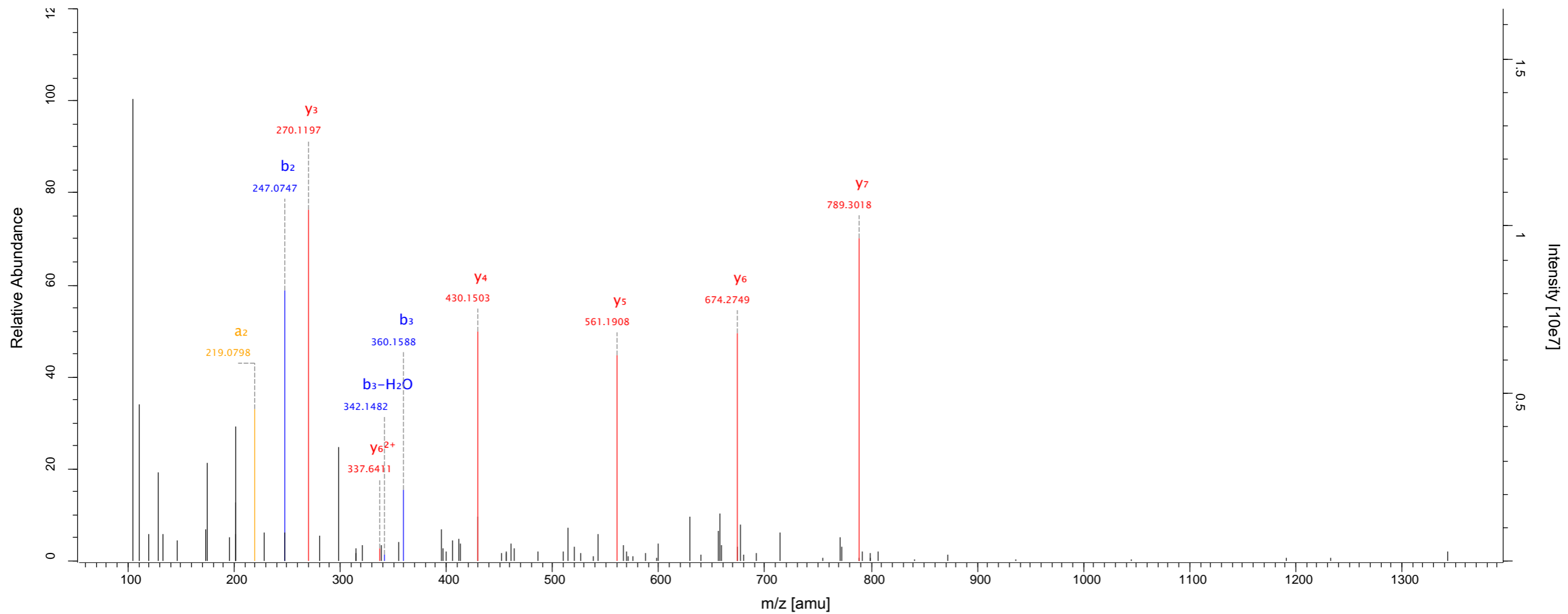
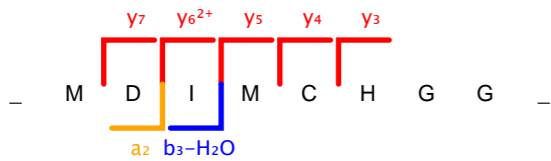
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 14087
Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
Peptide score: 156.93
Method: FTMS; HCD



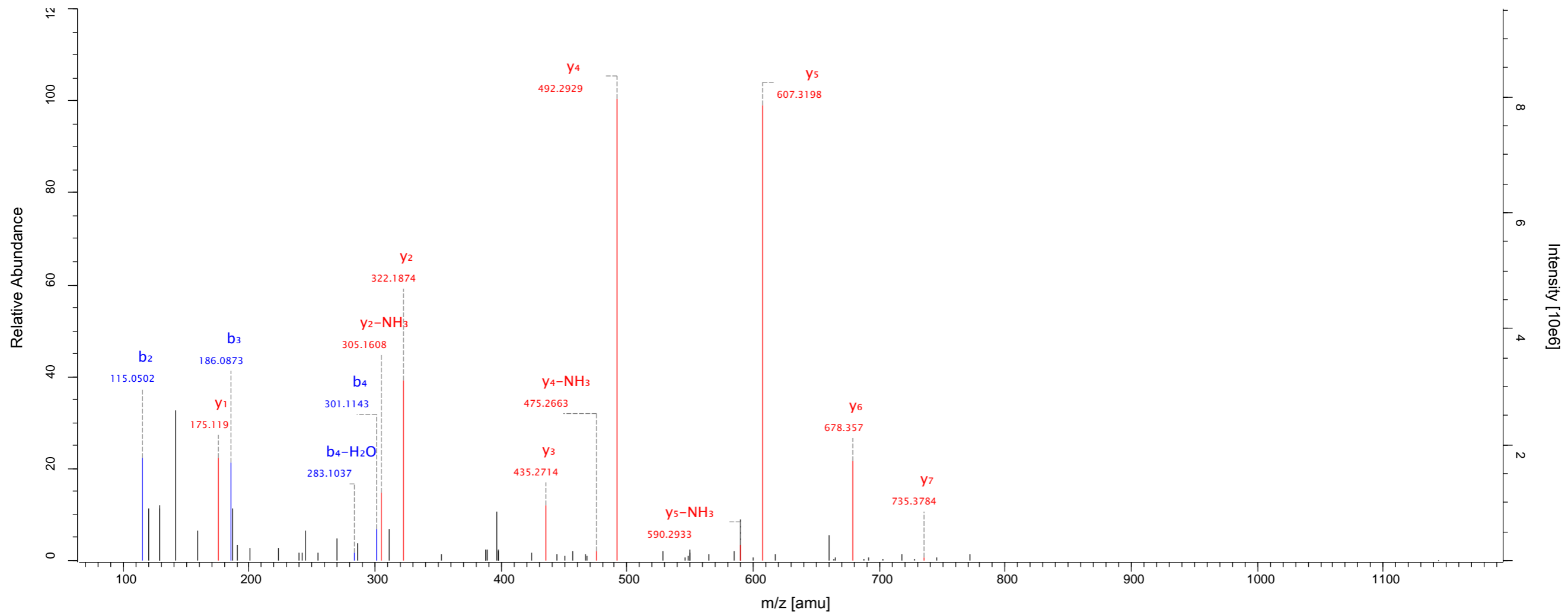
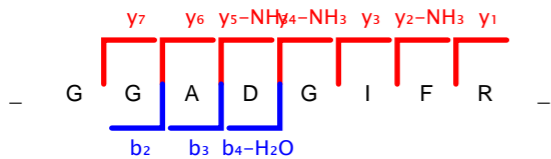
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 14568
Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
Peptide score: 66.62
Method: FTMS; HCD



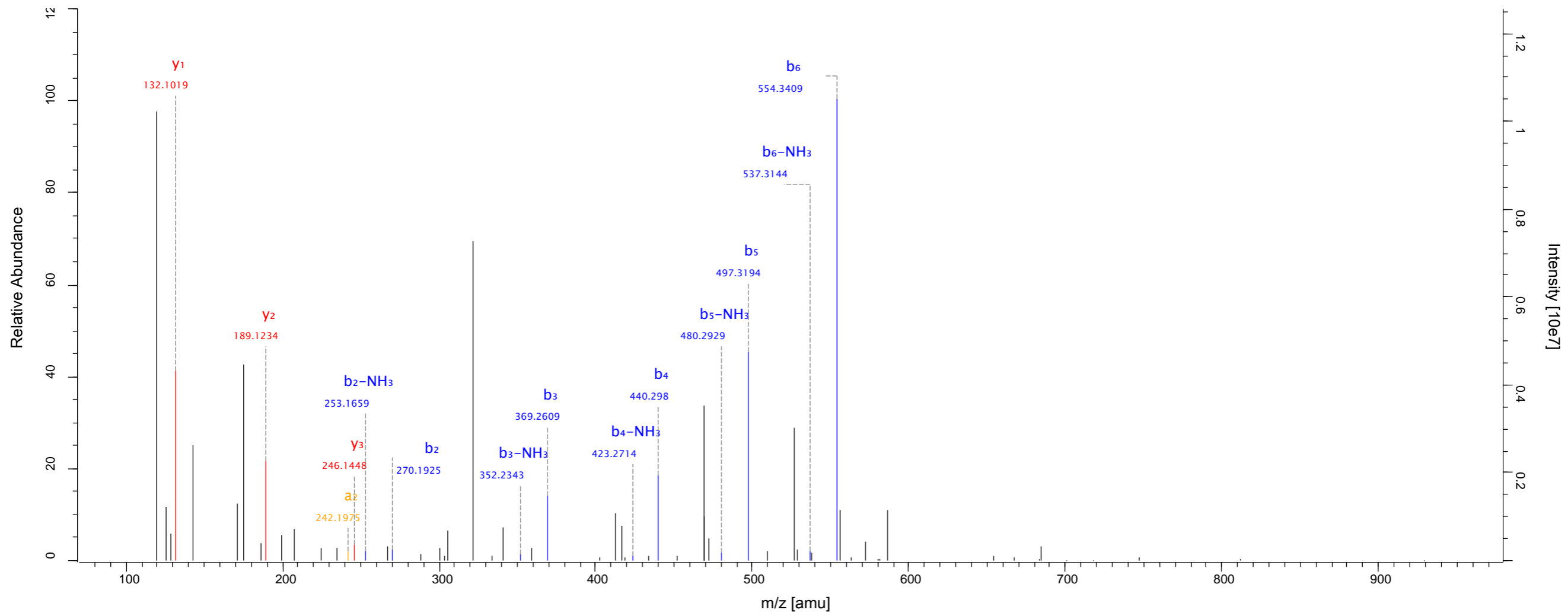
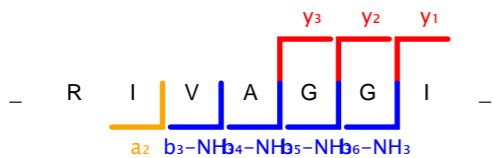
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 14703
Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>; rORF58 - NC_006273.2[37894,37920]- <unknown description>
Peptide score: 85.85
Method: FTMS; HCD



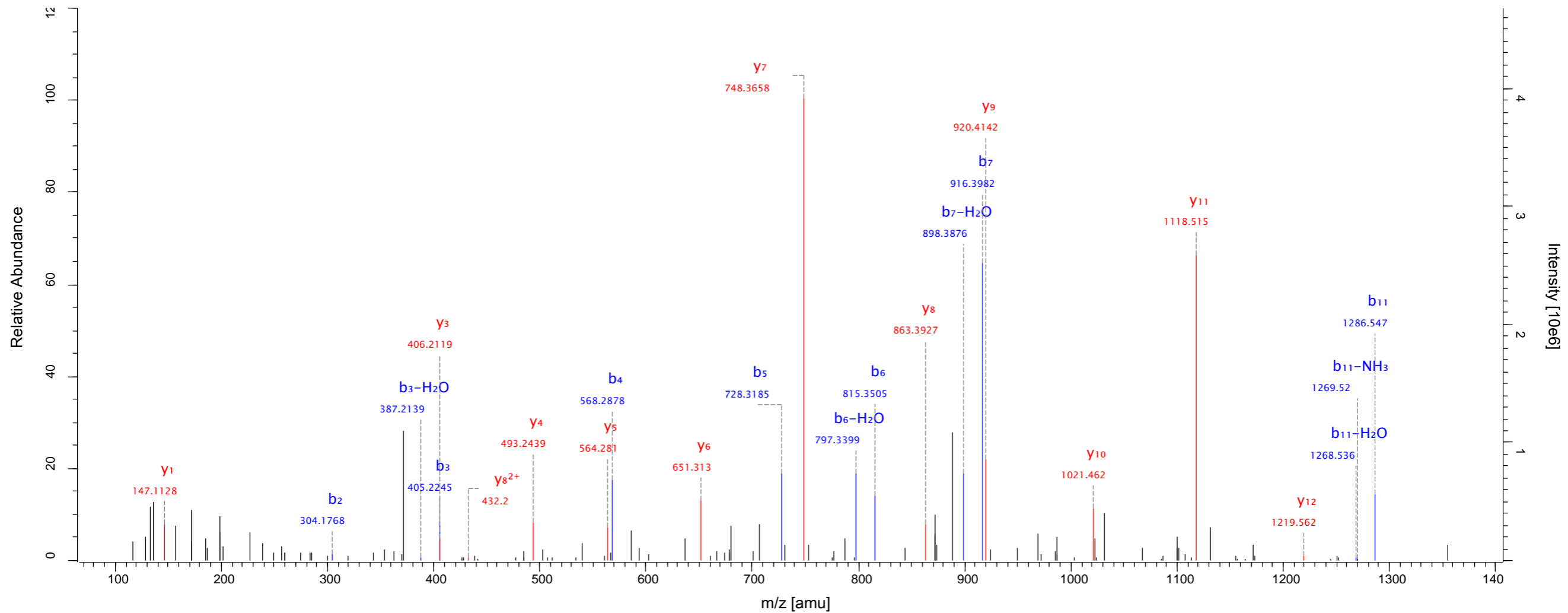
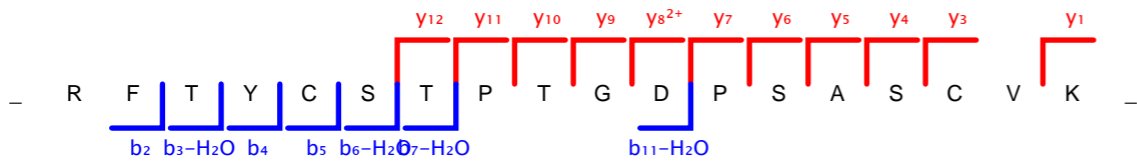
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 15240
Protein: ORFL159W + NC_006273.2[101475,109052]+ <unknown description>; ORFL159W.iORF1 + NC_006273.2[101526,109052]+ <unknown description>
Peptide score: 101.64
Method: FTMS; HCD



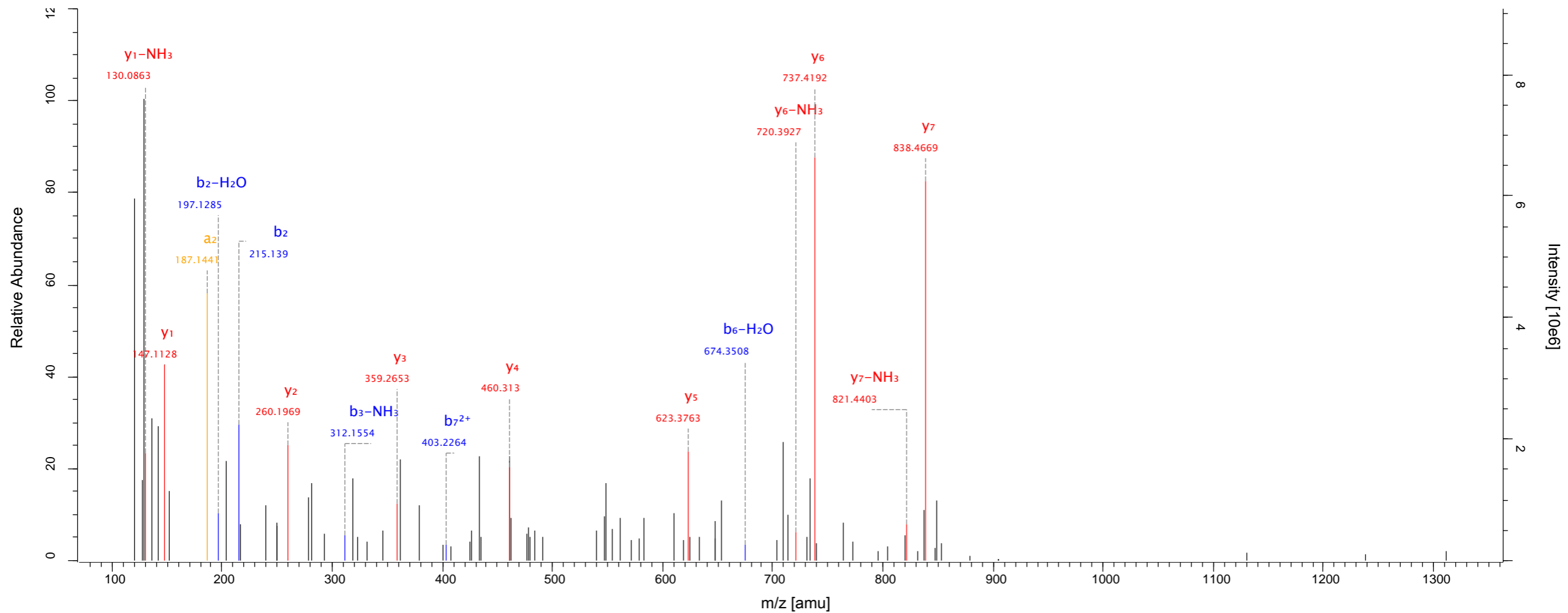
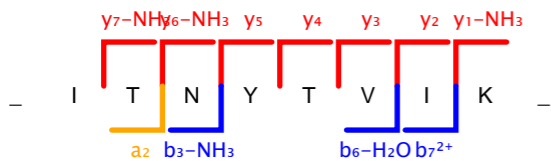
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 15840
Protein: ORFL19W + NC_006273.2[8136,8387]+ <unknown description>
Peptide score: 107.65
Method: FTMS; HCD



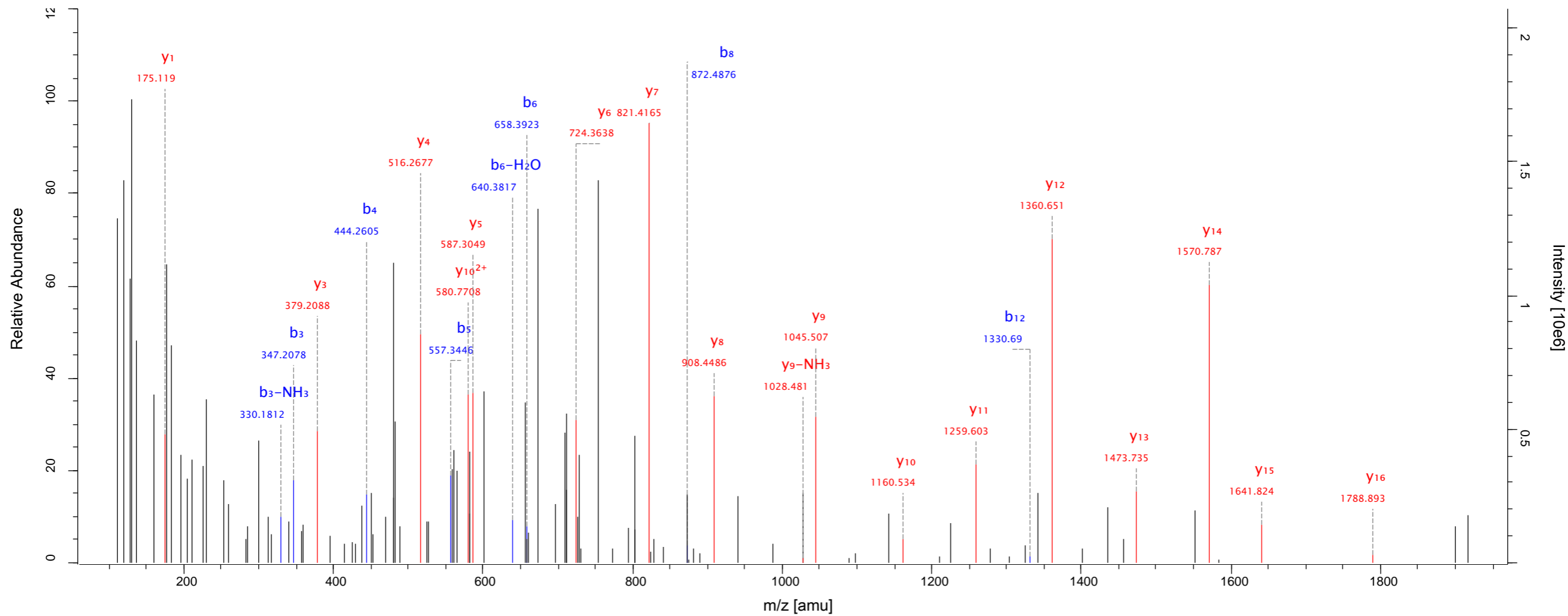
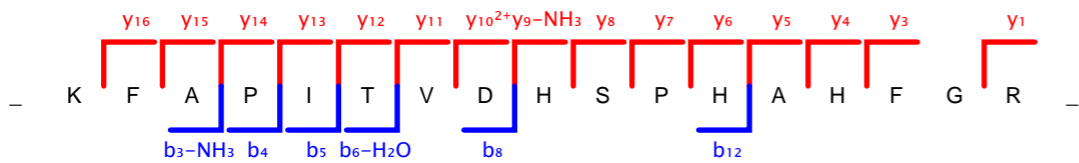
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 16769
 Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
 Peptide score: 93.23
 Method: FTMS; HCD



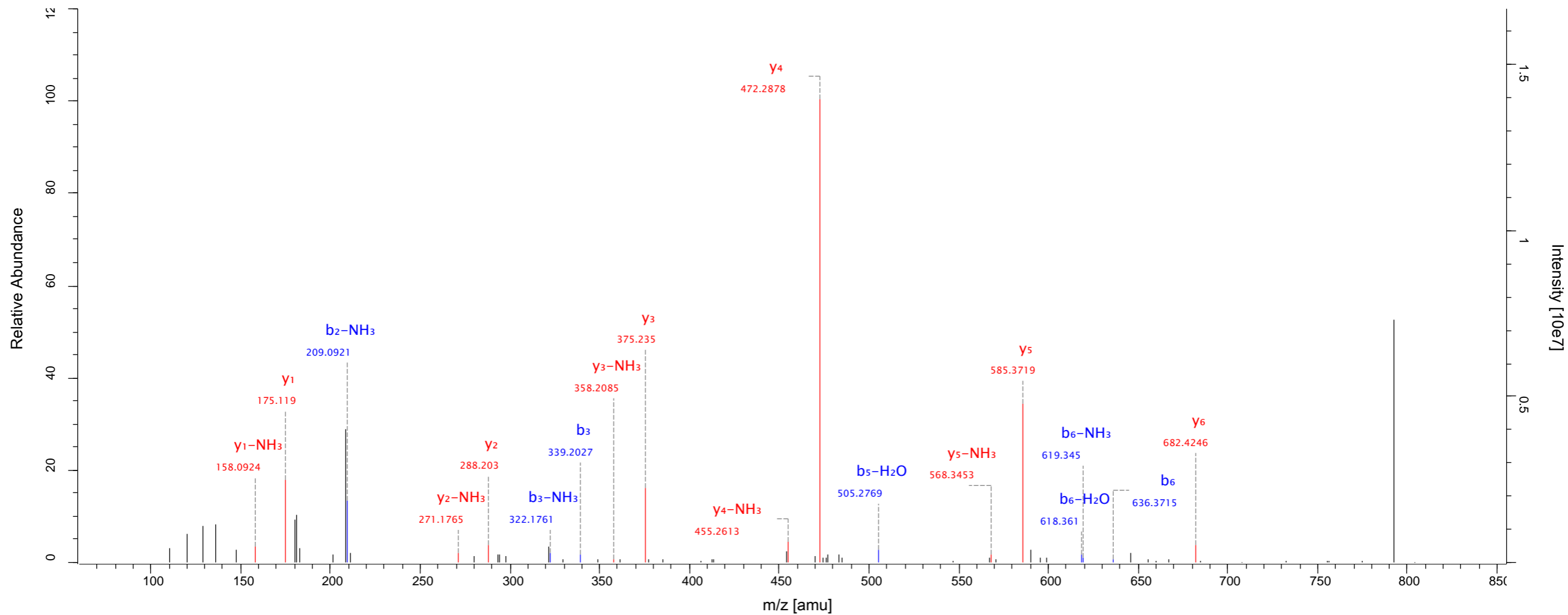
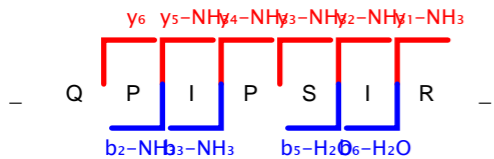
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 16907
 Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
 Peptide score: 91.9
 Method: FTMS; HCD



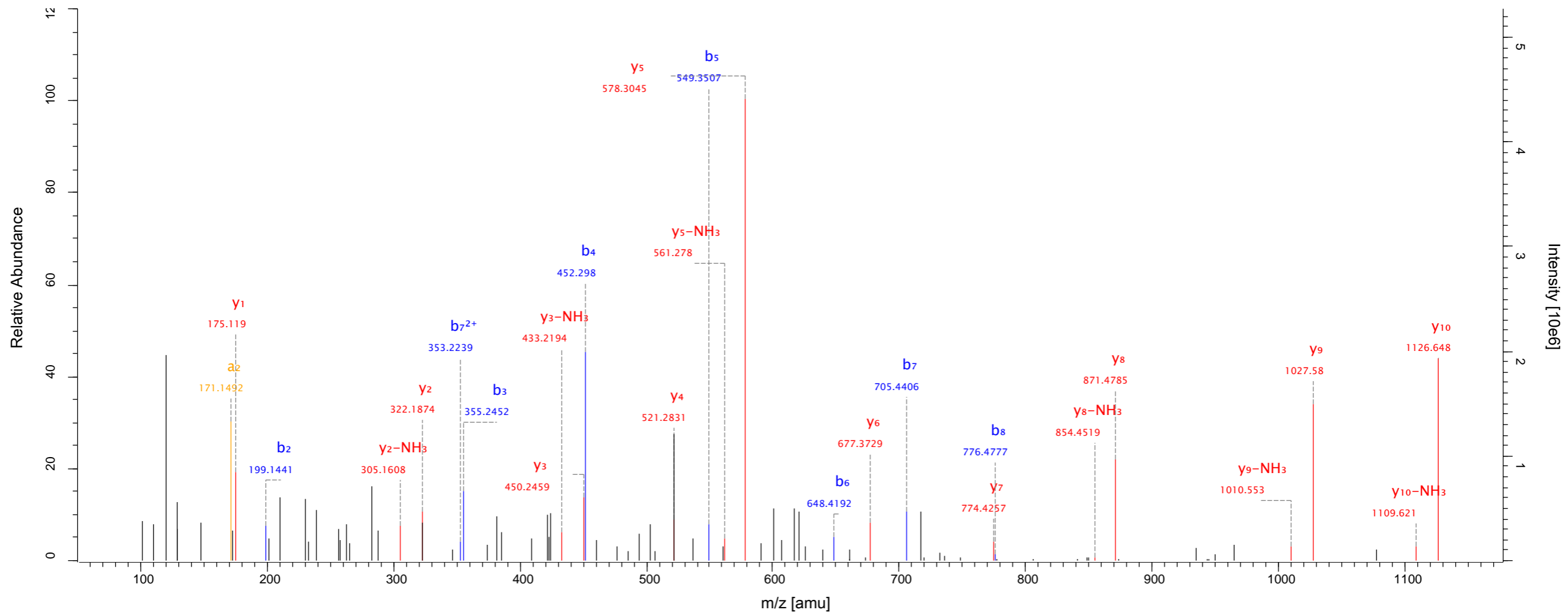
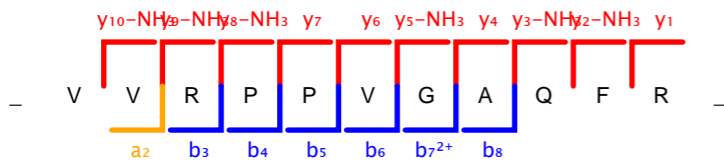
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 16939
Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
Peptide score: 115.8
Method: FTMS; HCD



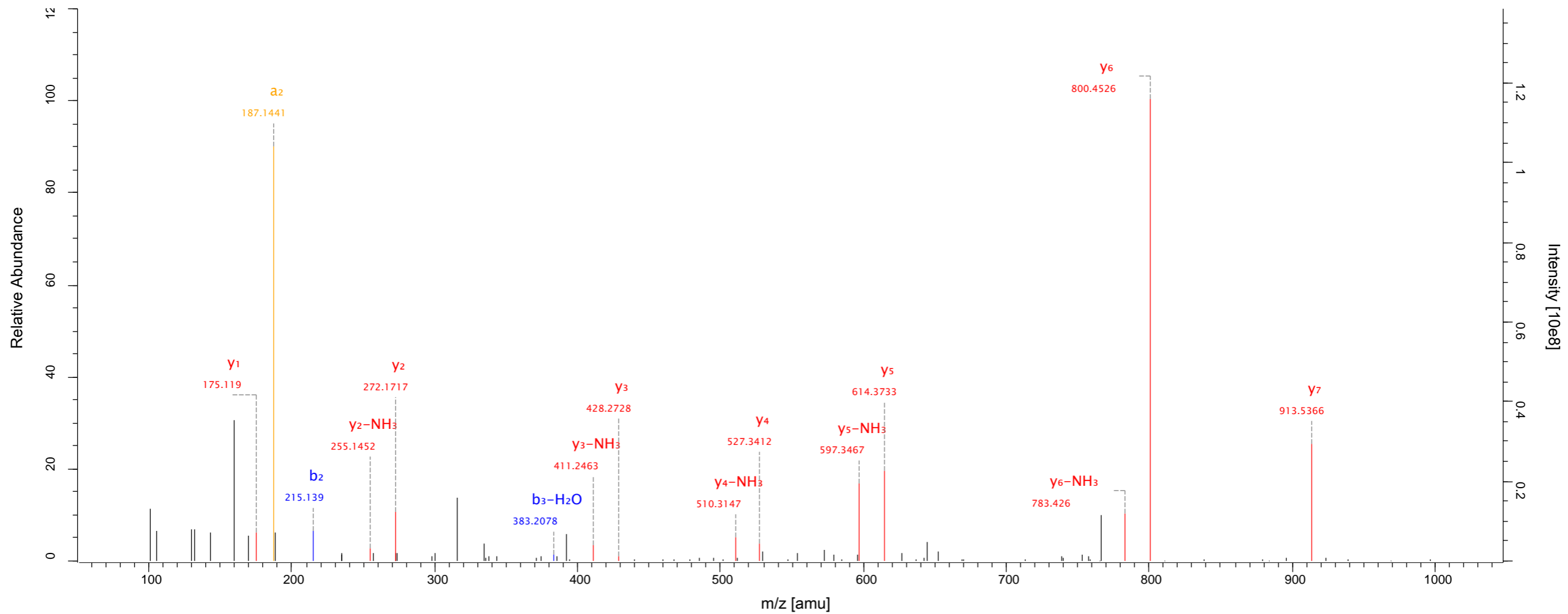
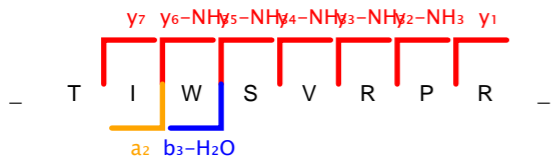
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 17005
Protein: ORFL137W.IORF1 + NC_006273.2[75299,75529]+ <unknown description>
Peptide score: 123.99
Method: FTMS; HCD



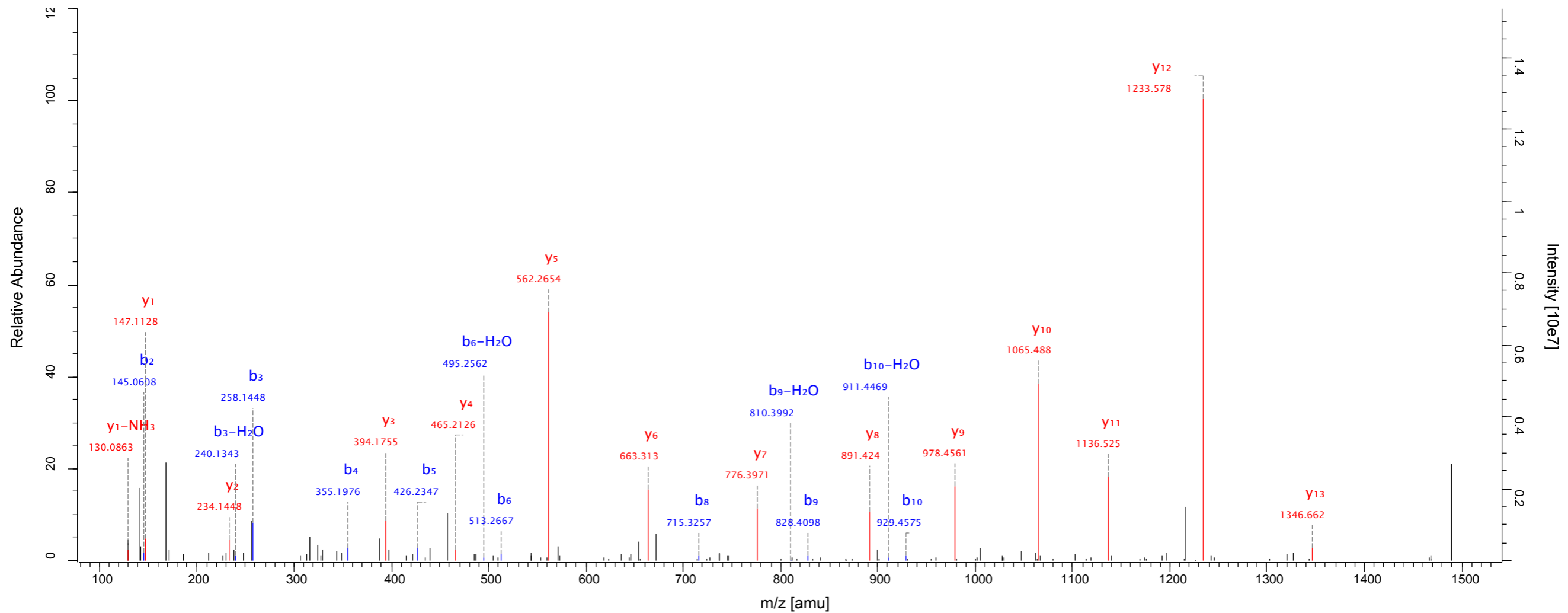
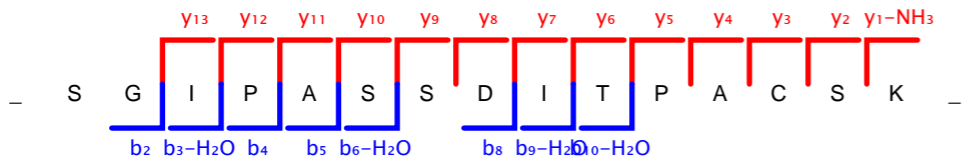
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 17481
 Protein: ORFS334W + NC_006273.2[204331,204558]+ <unknown description>
 Peptide score: 166.66
 Method: FTMS; HCD



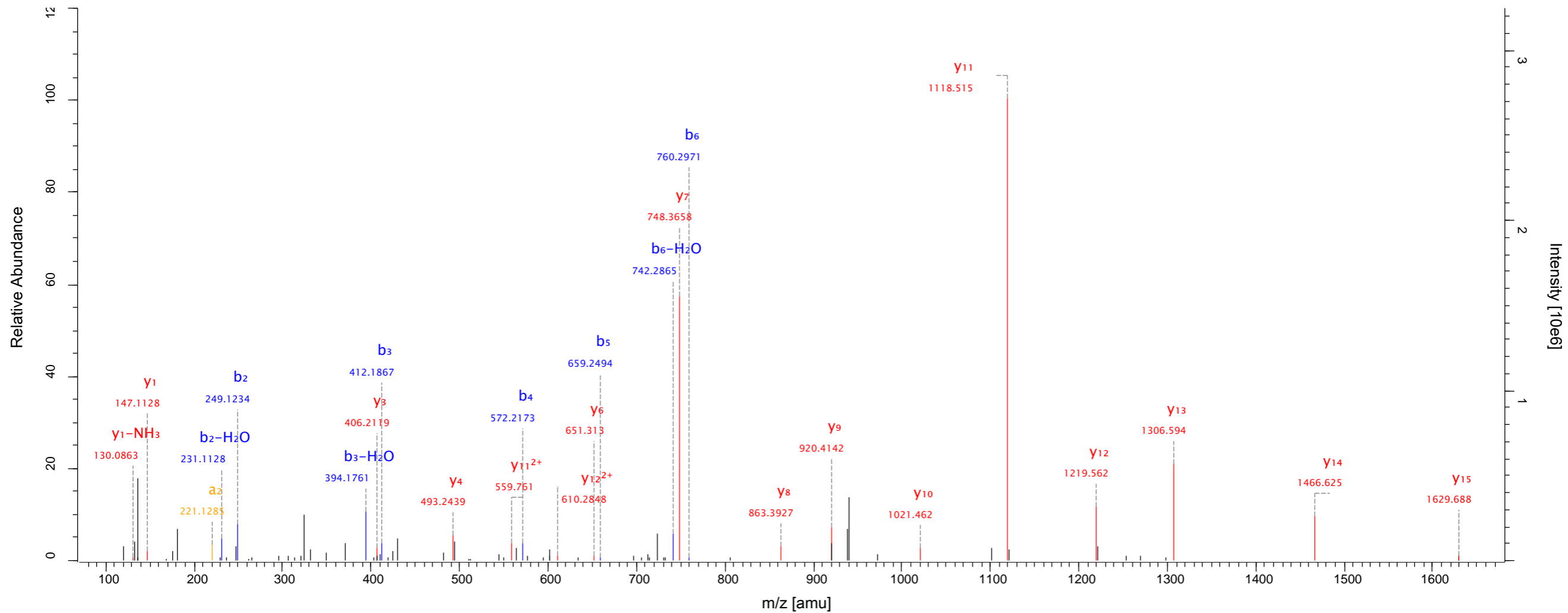
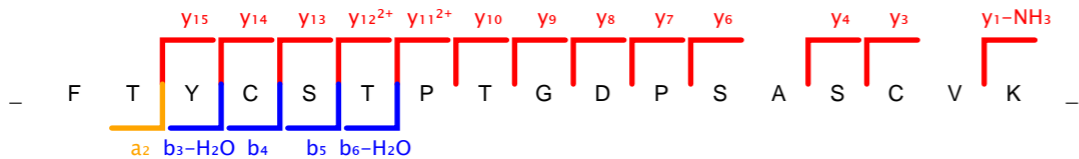
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 18718
Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>
Peptide score: 119.21
Method: FTMS; HCD



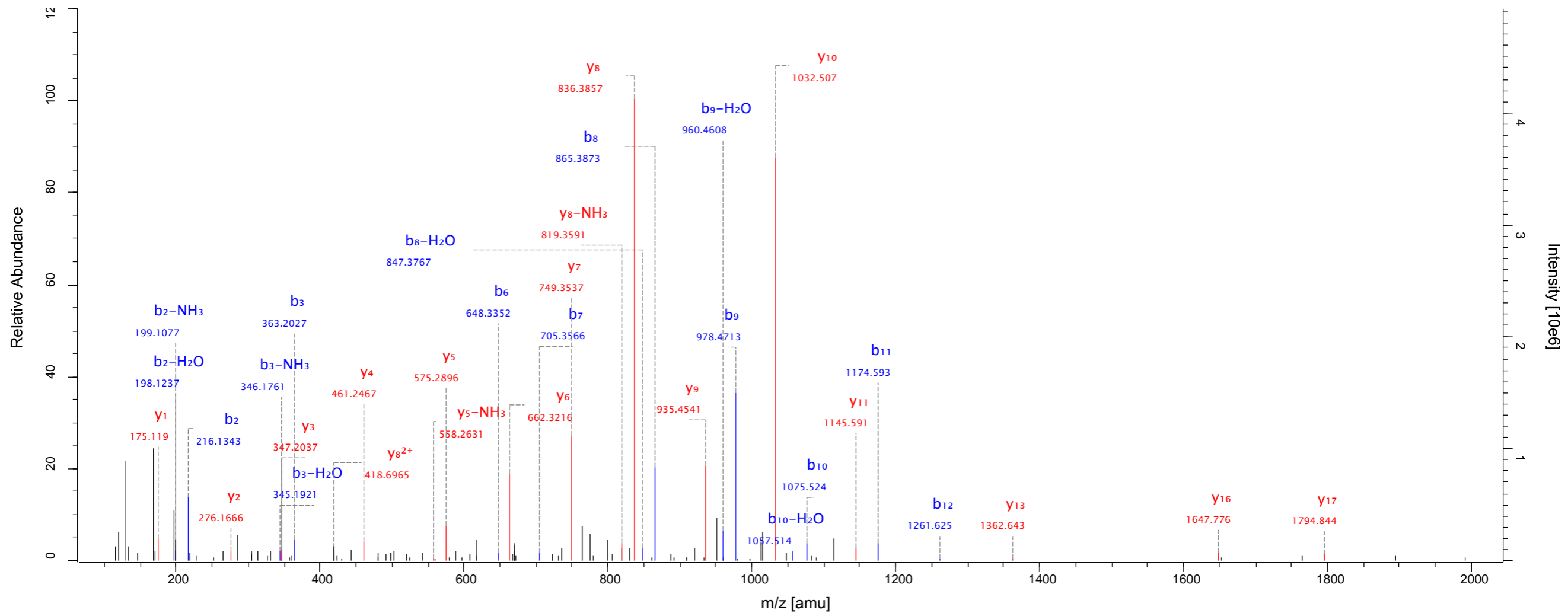
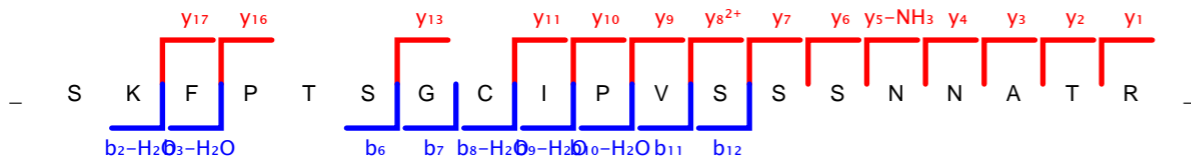
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 18744
Protein: ORFL298W + NC_006273.2[184737,184916]+ <unknown description>
Peptide score: 103.67
Method: FTMS; HCD



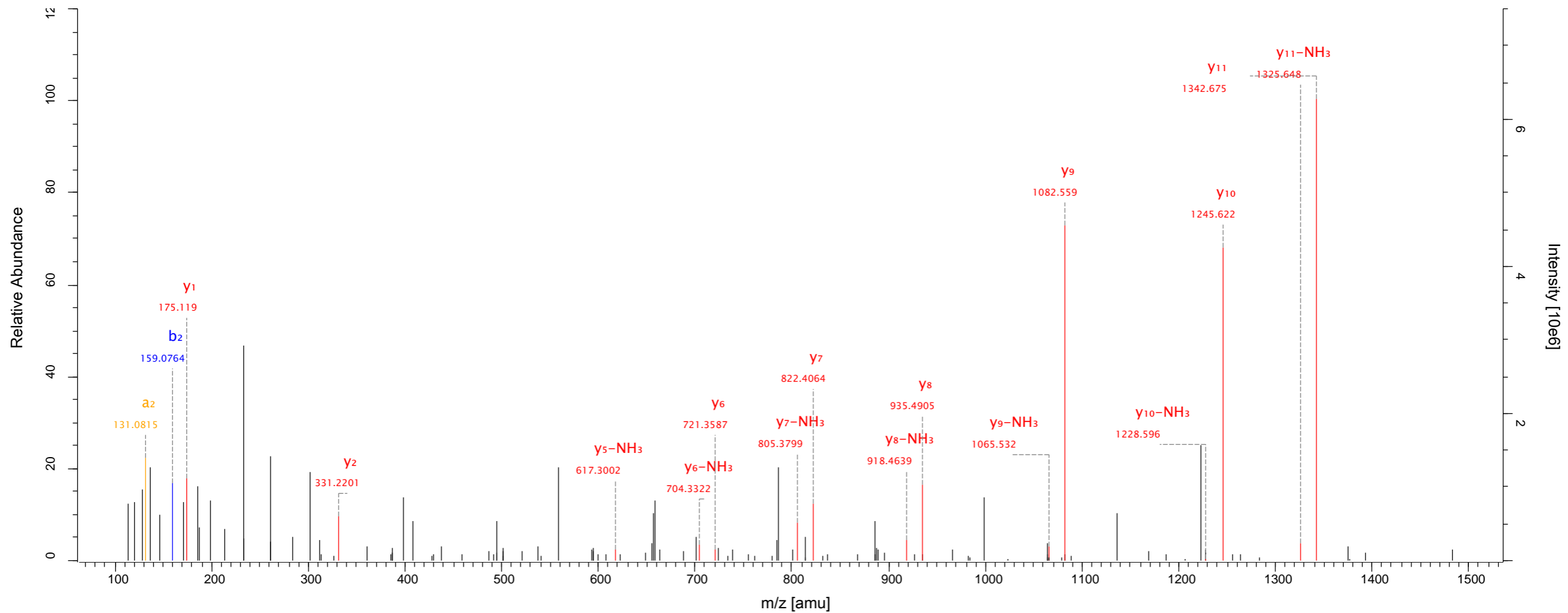
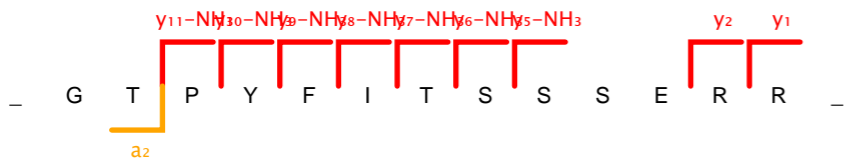
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 19168
 Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
 Peptide score: 120.3
 Method: FTMS; HCD



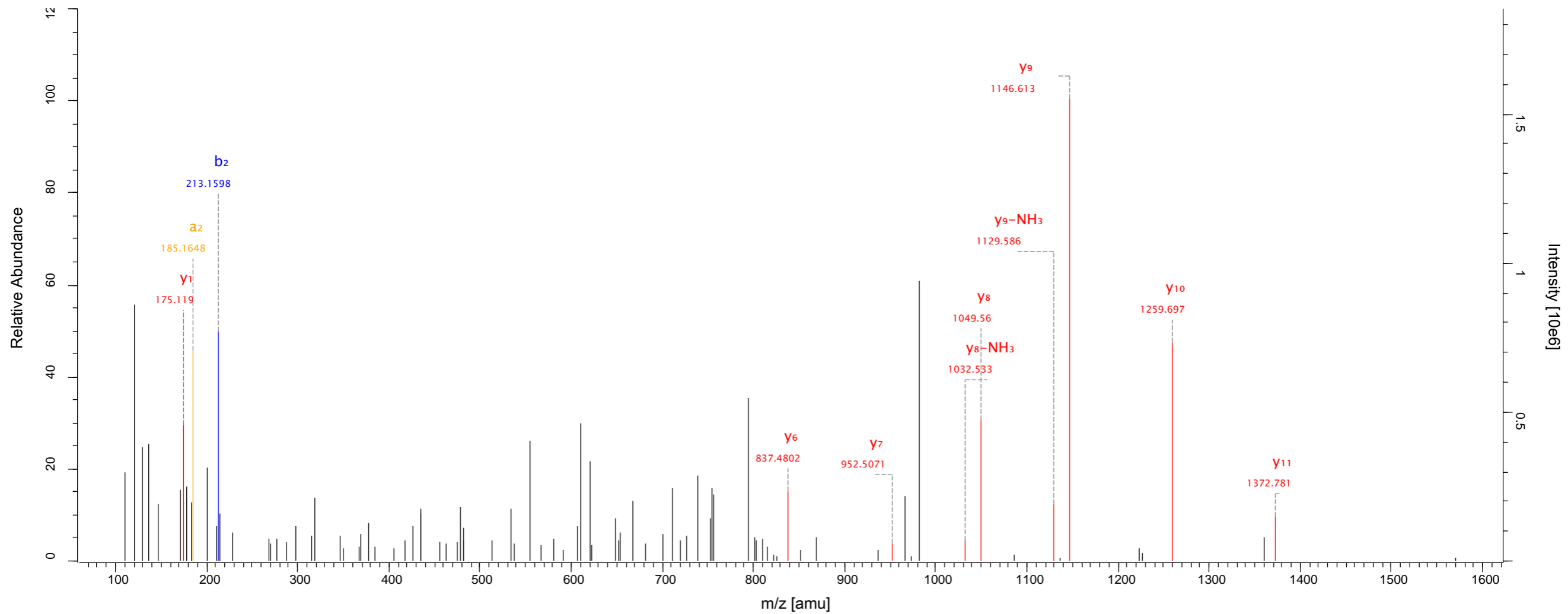
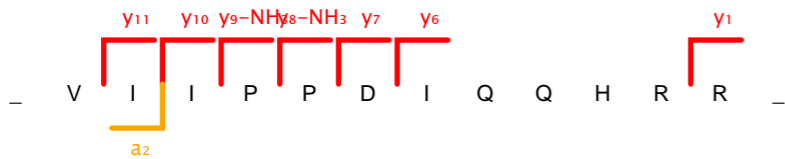
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 19236
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 129.43
 Method: FTMS; HCD



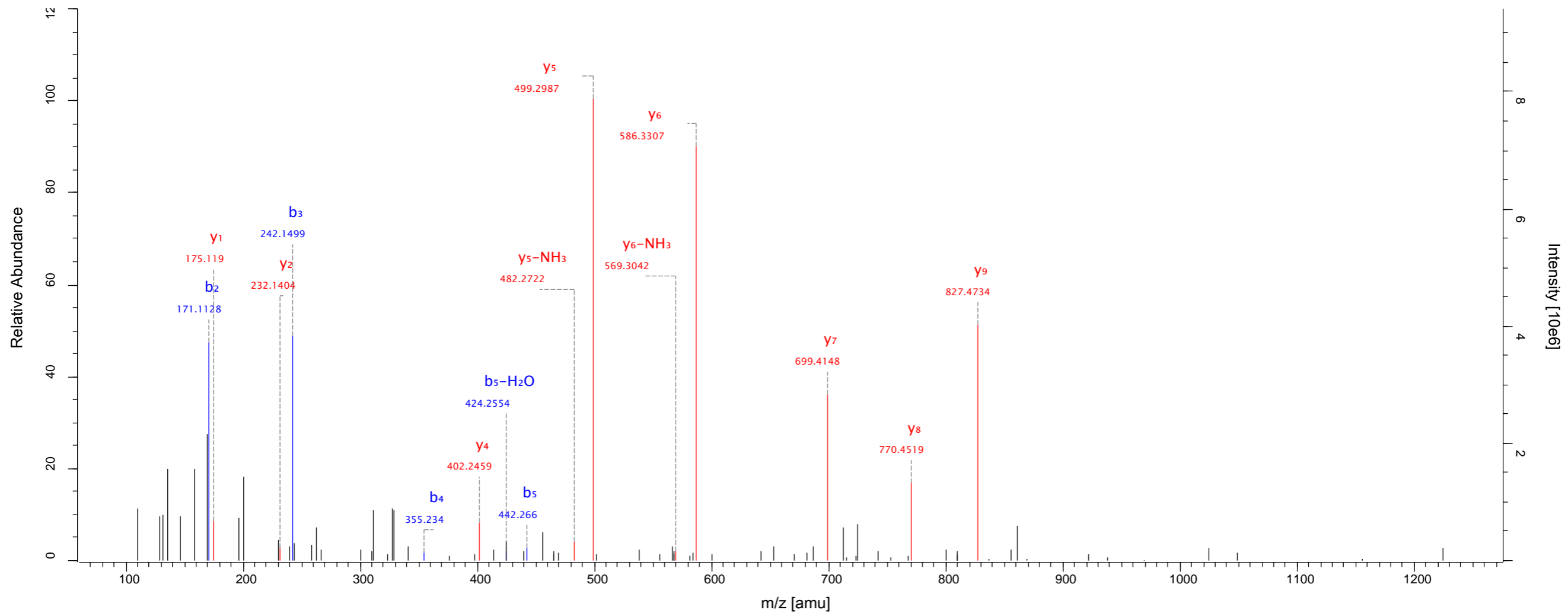
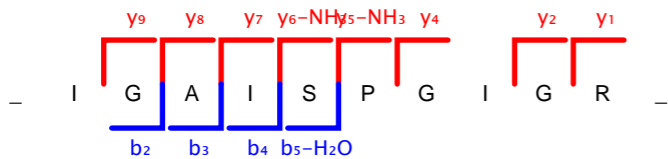
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 19316
Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
Peptide score: 83.05
Method: FTMS; HCD



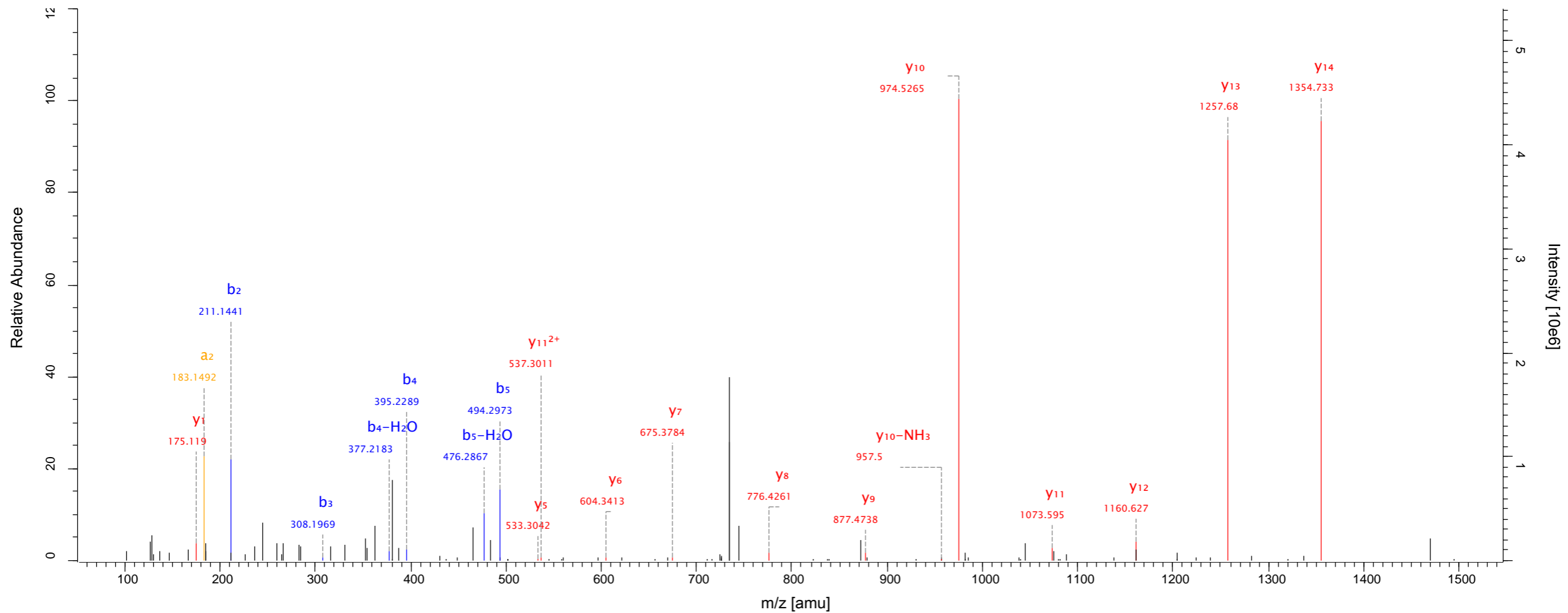
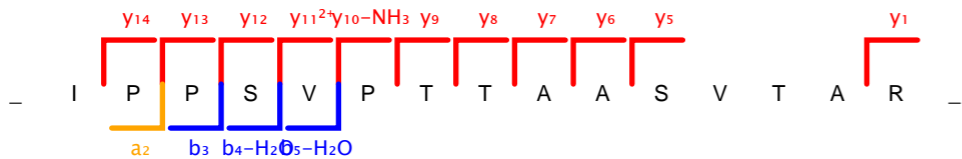
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 19495
Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
Peptide score: 76.14
Method: FTMS; HCD



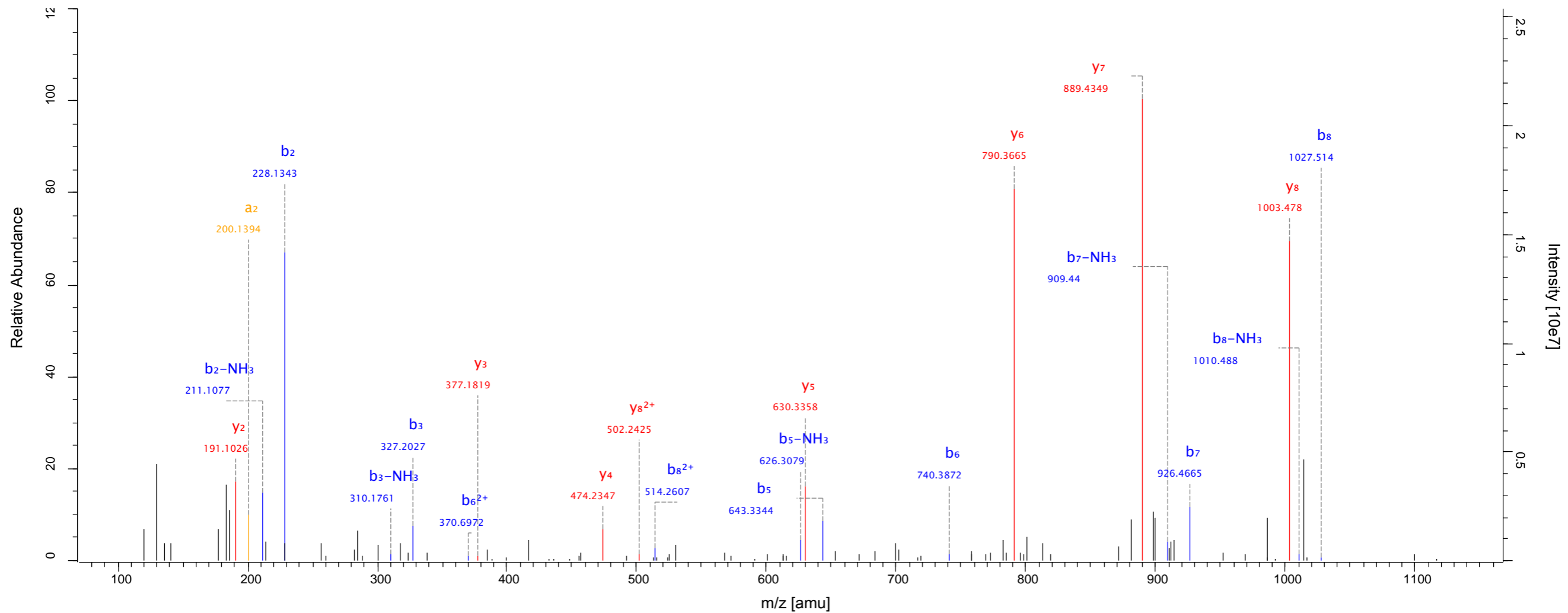
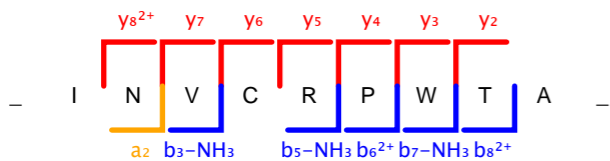
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 20603
Protein: ORFL154W + NC_006273.2[96843,97115]+ <unknown description>; ORFL154W.iORF1 + NC_006273.2[96972,97115]+ <unknown description>
Peptide score: 64.3
Method: FTMS; HCD



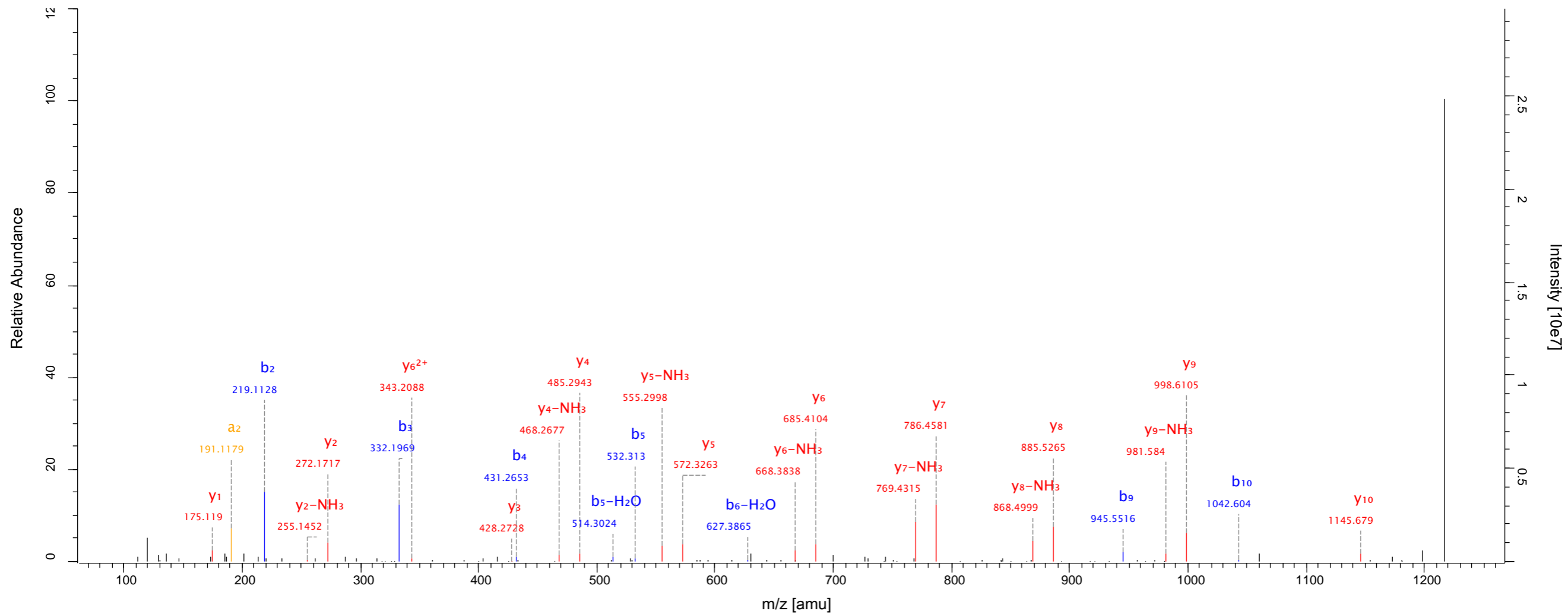
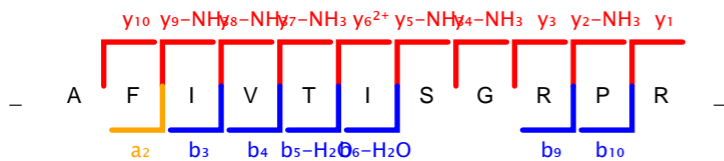
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 22274
Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
Peptide score: 76.73
Method: FTMS; HCD



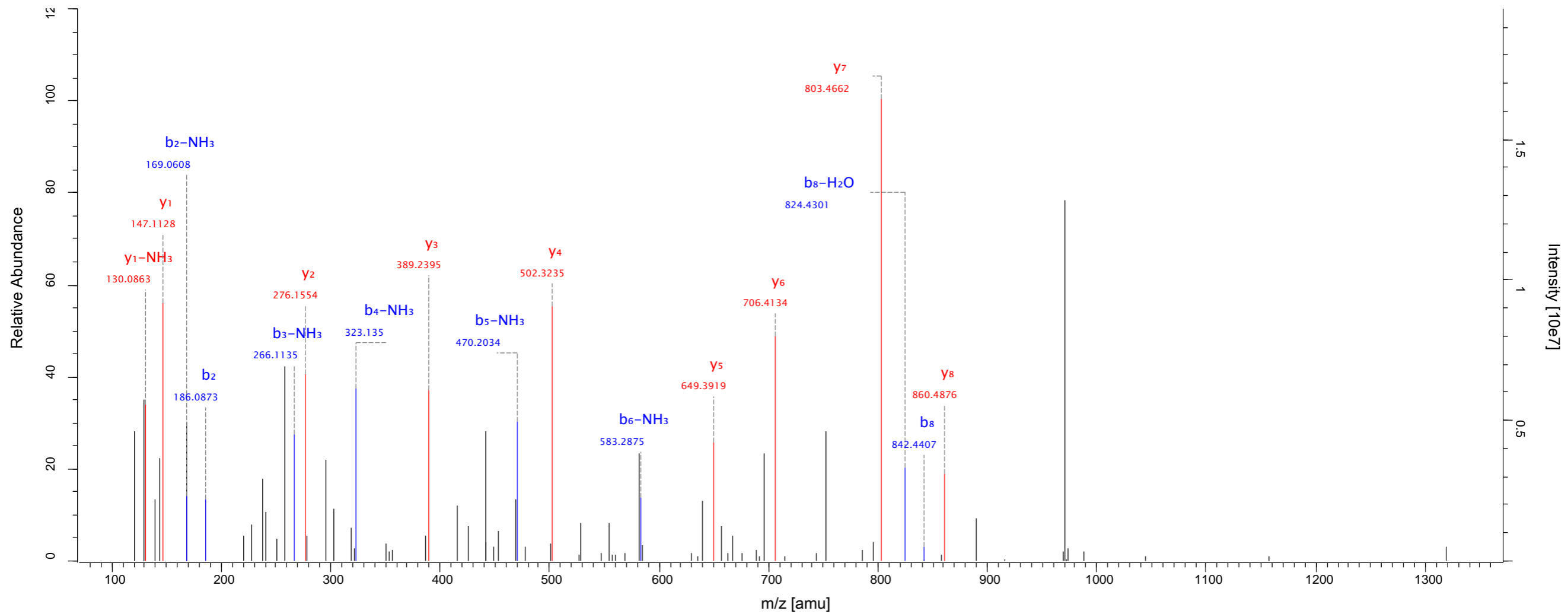
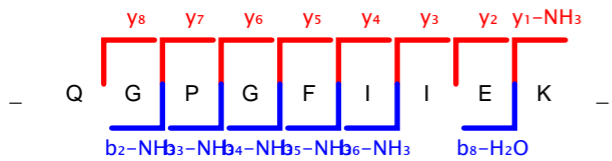
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 25634
Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
Peptide score: 166.78
Method: FTMS; HCD



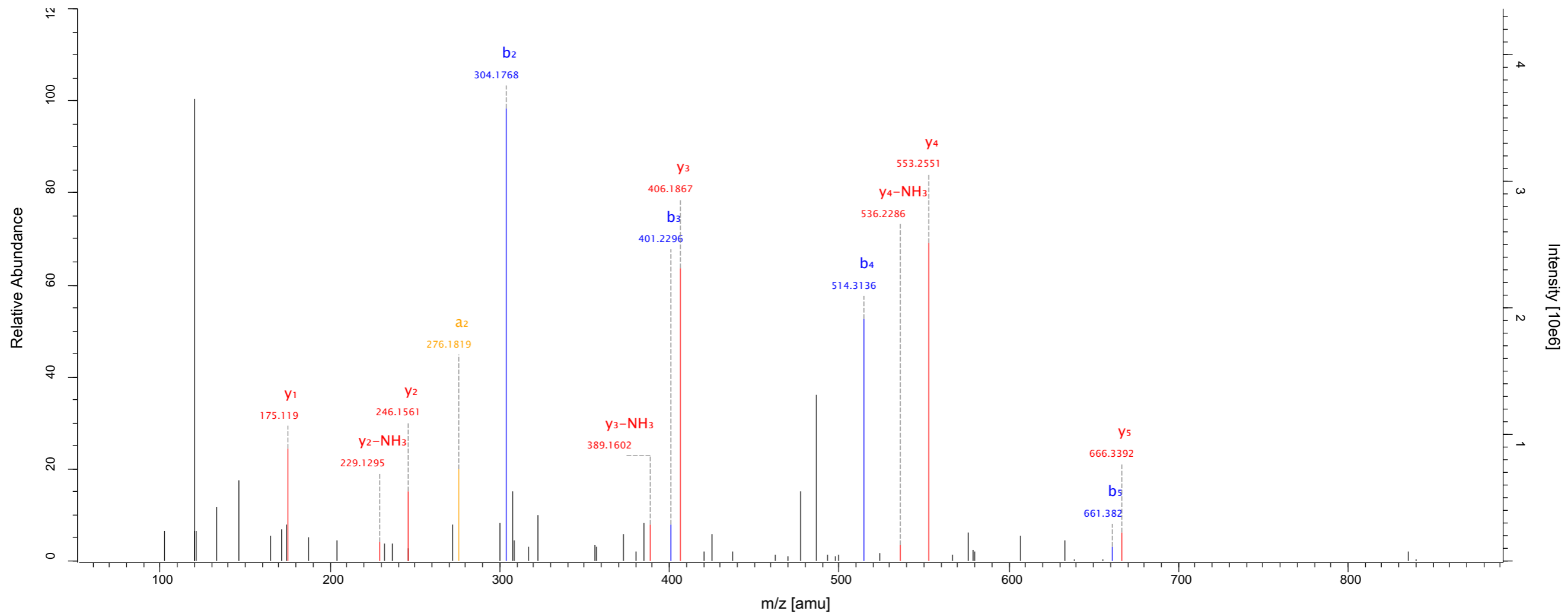
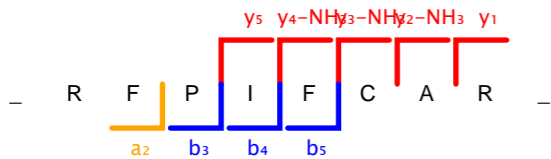
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 26093
 Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
 Peptide score: 199.11
 Method: FTMS; HCD



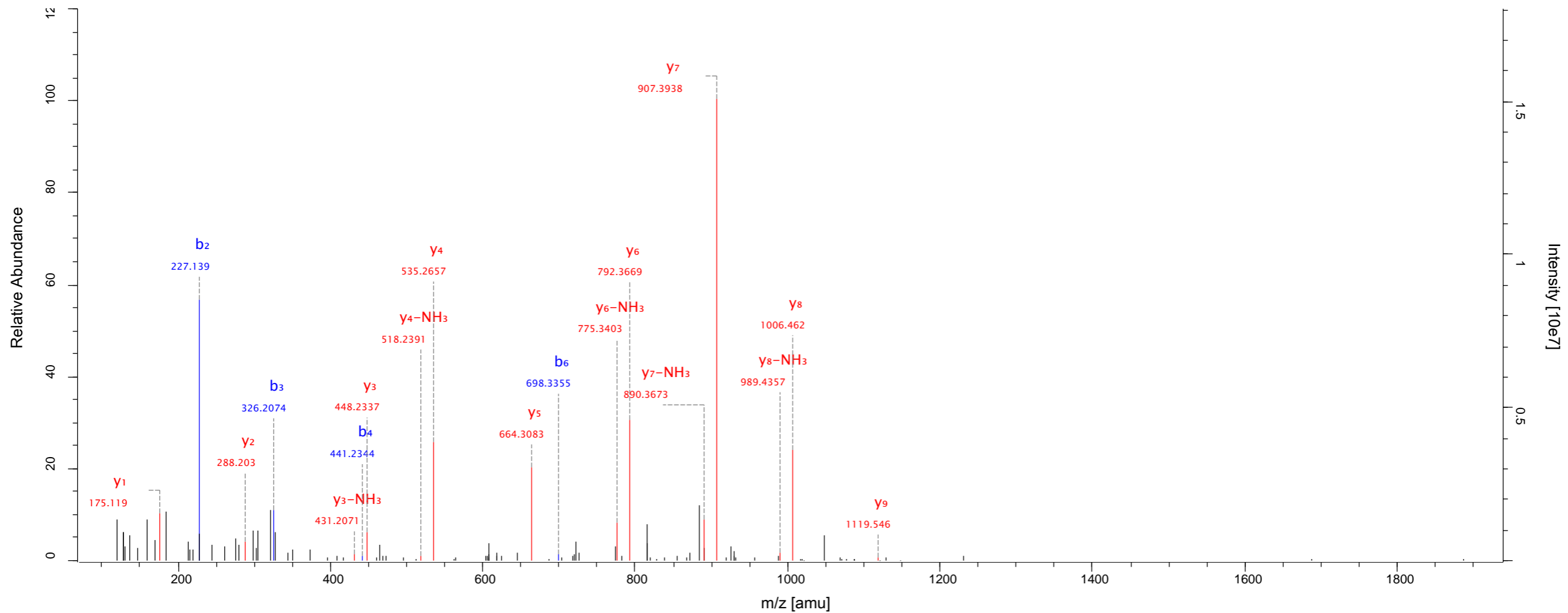
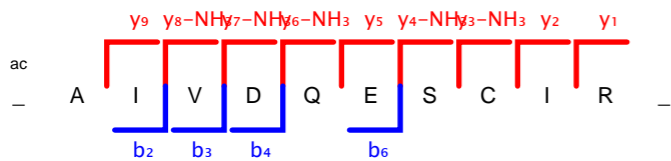
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 26187
Protein: ORFL19W + NC_006273.2[8136,8387]+ <unknown description>
Peptide score: 78.93
Method: FTMS; HCD



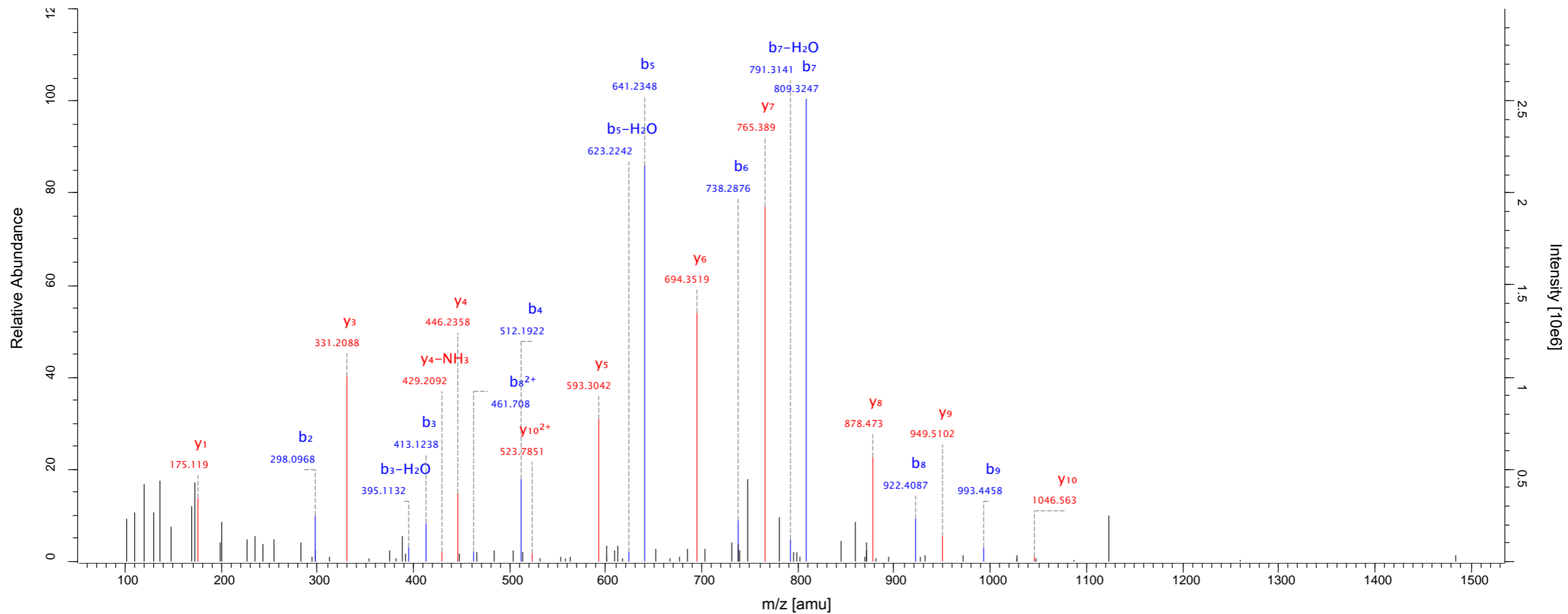
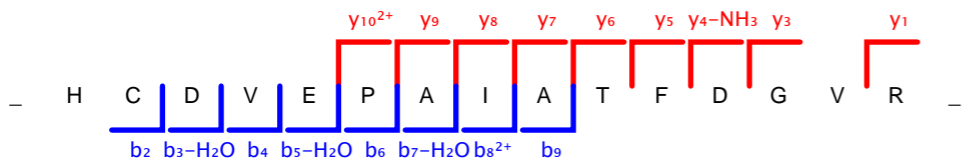
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 26311
Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
Peptide score: 89.3
Method: FTMS; HCD



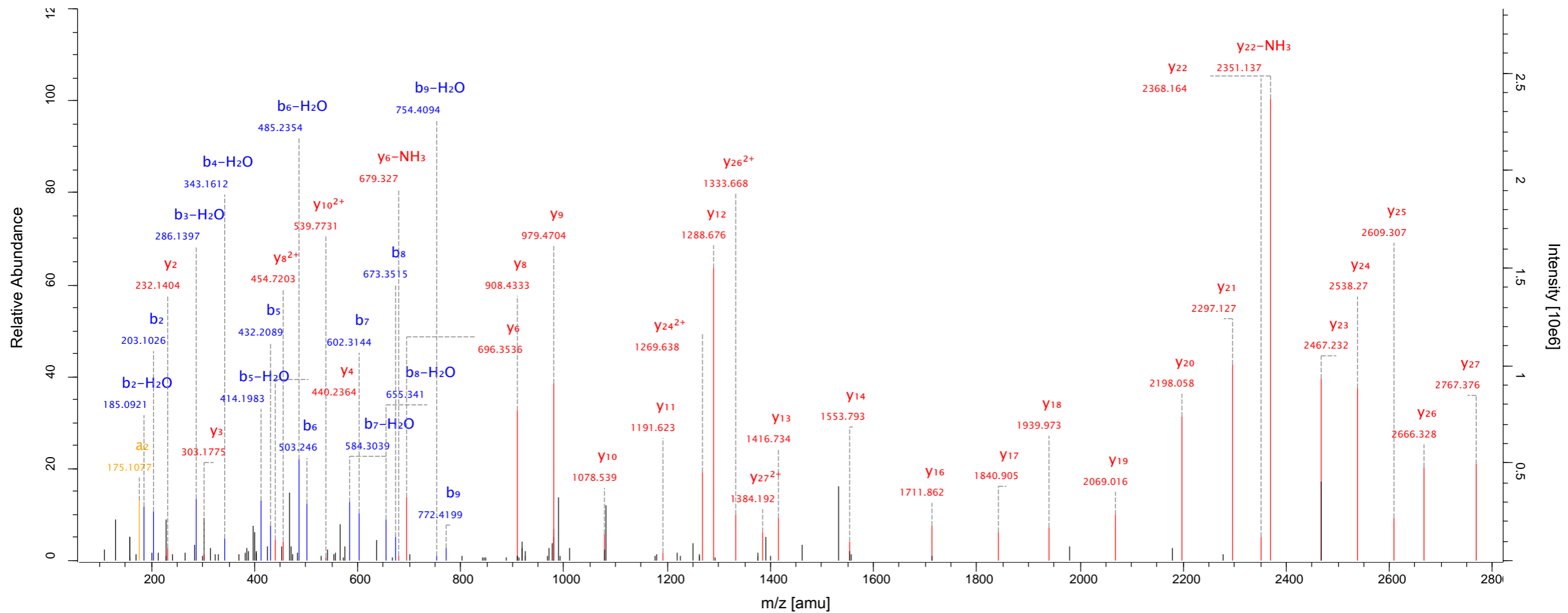
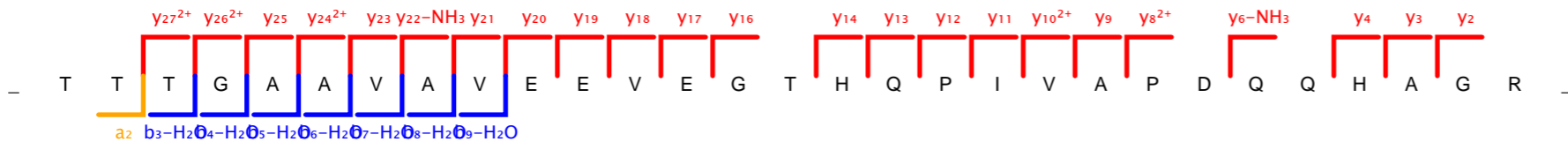
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 27548
Protein: ORFL151C - NC_006273.2[92499,92870]- <unknown description>
Peptide score: 136.57
Method: FTMS; HCD



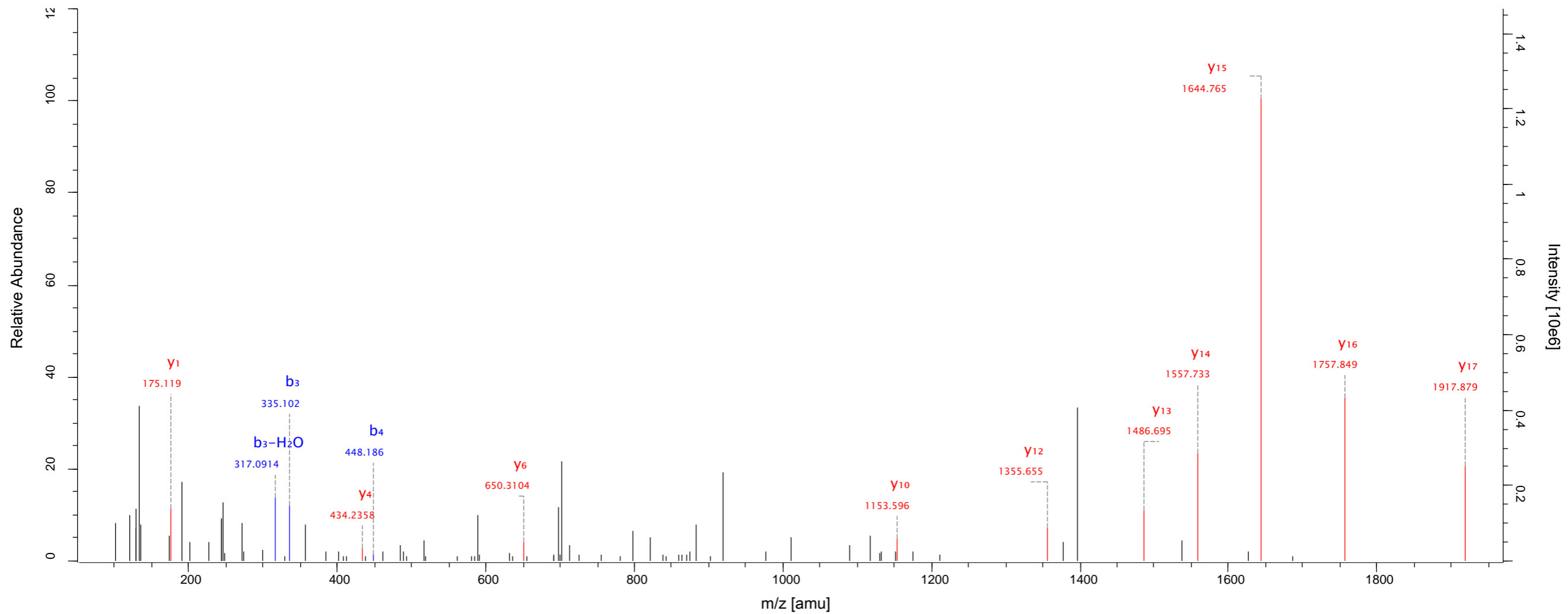
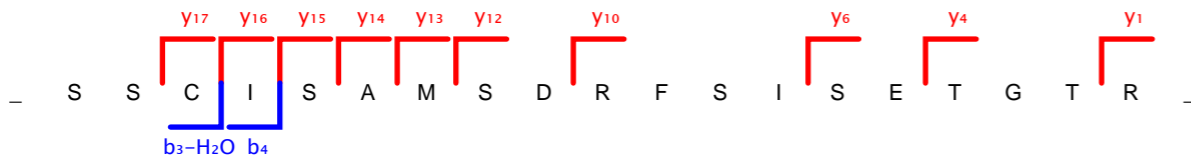
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 29560
Protein: ORFL151C - NC_006273.2[92499,92870]- <unknown description>
Peptide score: 119.25
Method: FTMS; HCD



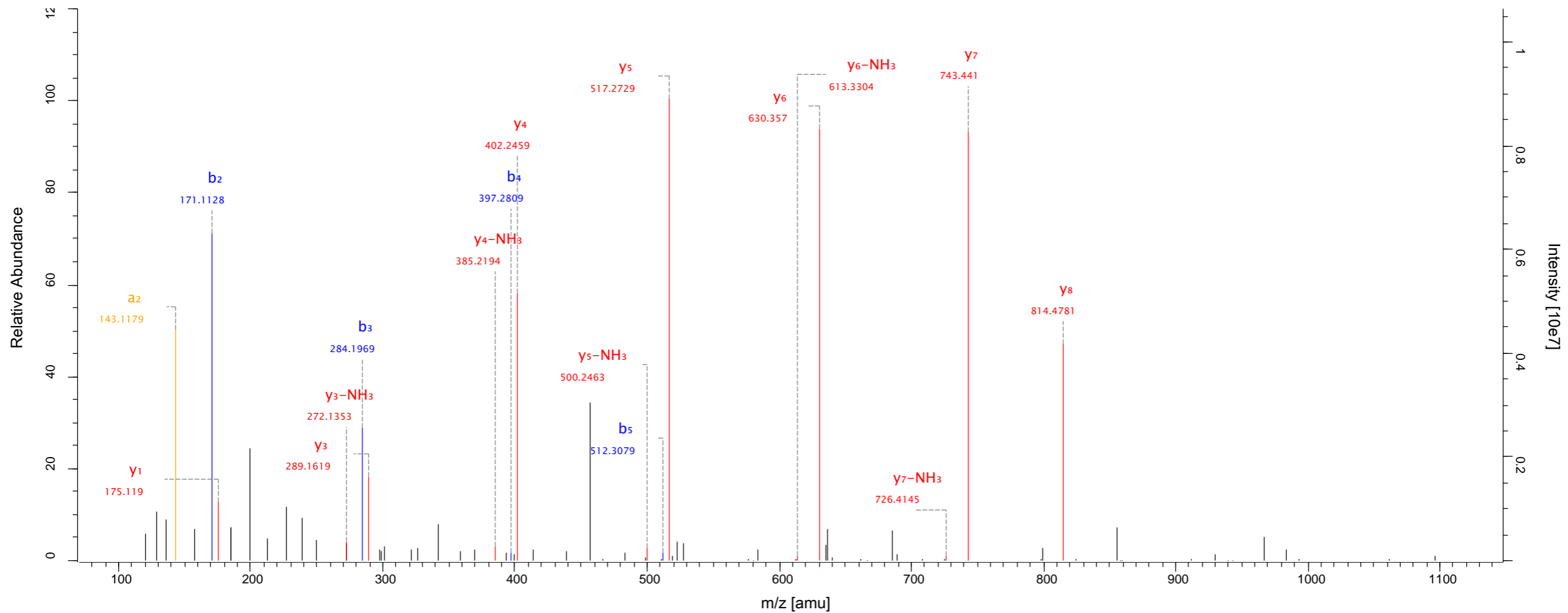
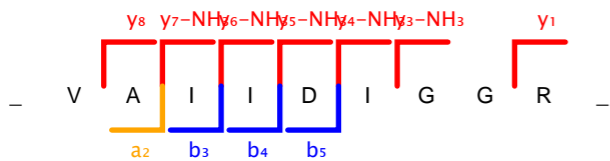
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 29616
 Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
 Peptide score: 258.85
 Method: FTMS; HCD



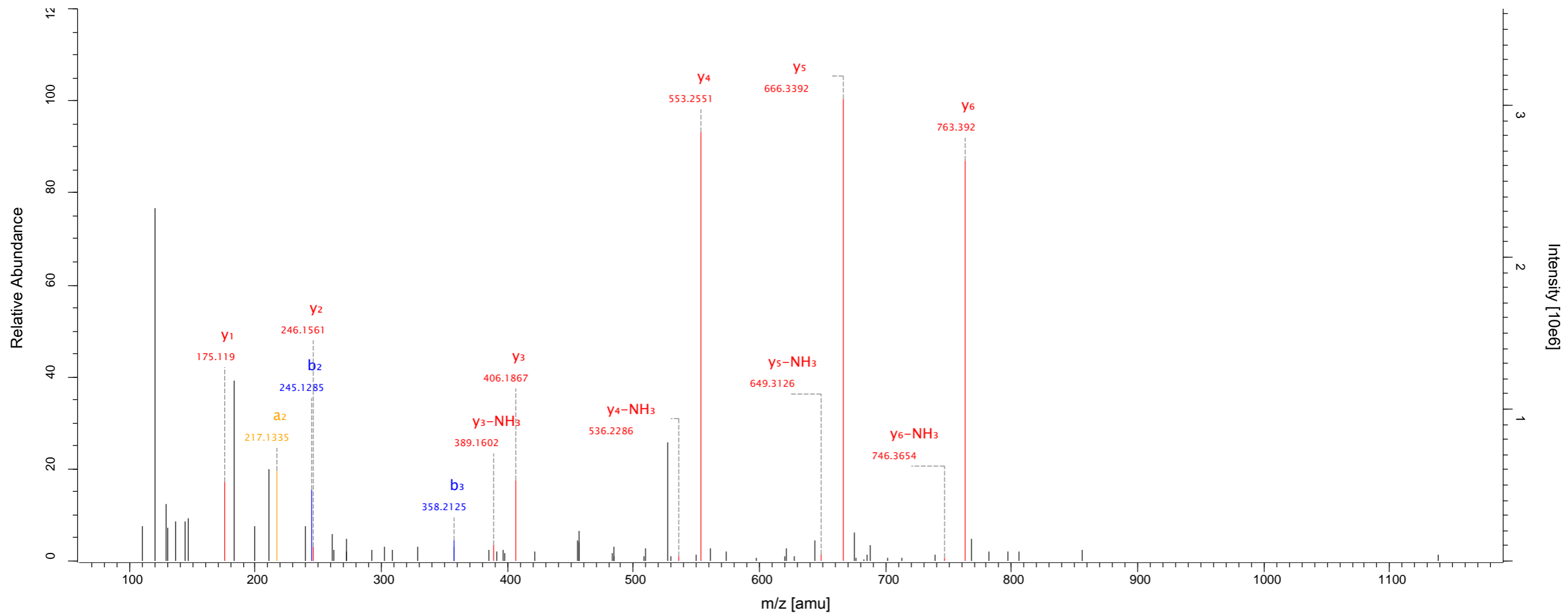
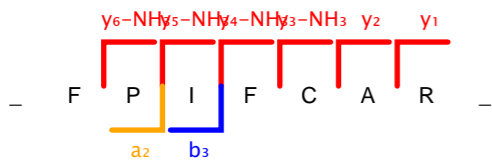
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
Scan number: 30033
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 65.72
Method: FTMS; HCD



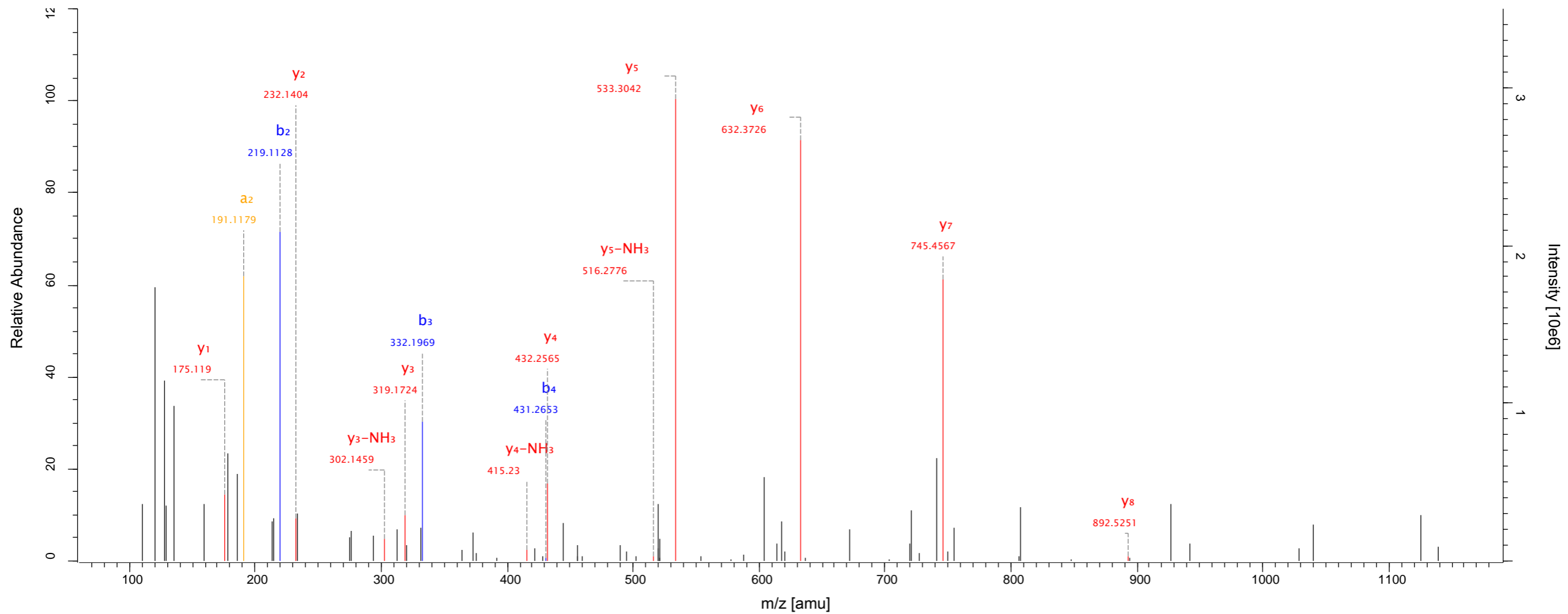
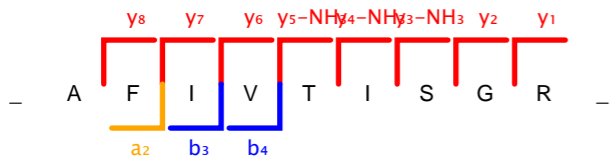
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpI1_10
Scan number: 32268
Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
Peptide score: 119.41
Method: FTMS; HCD



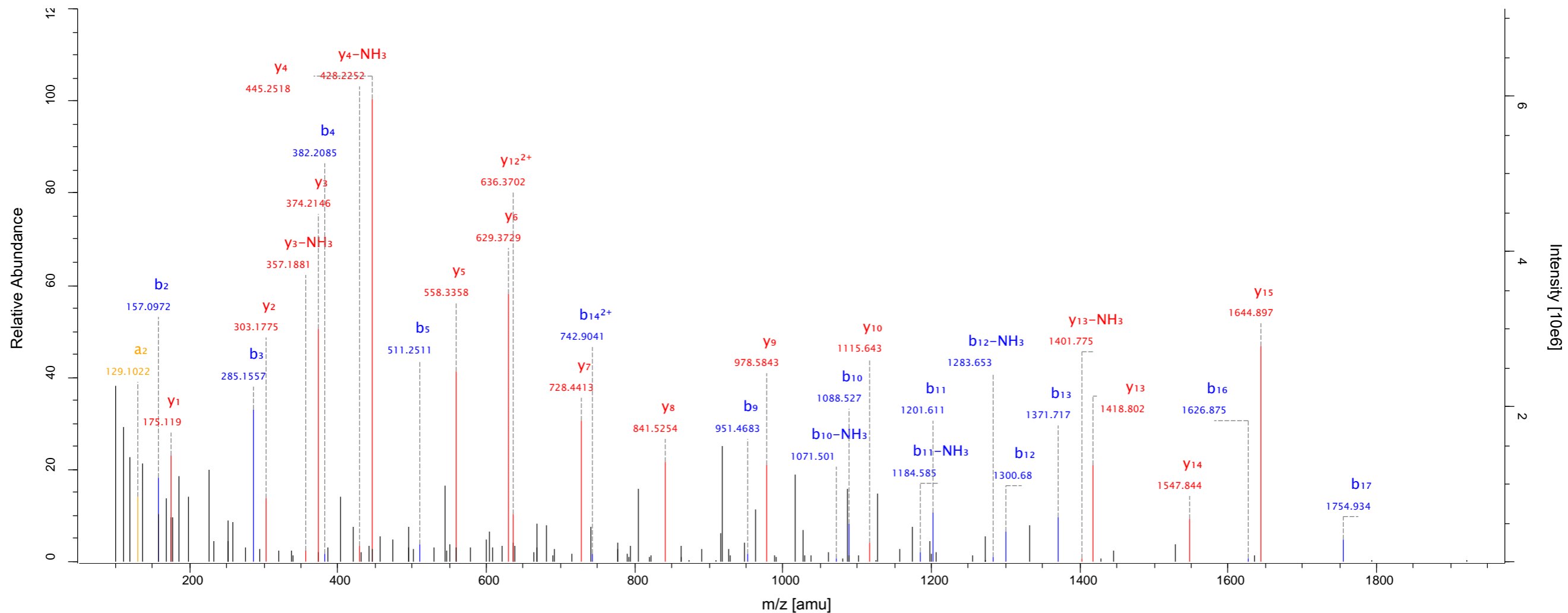
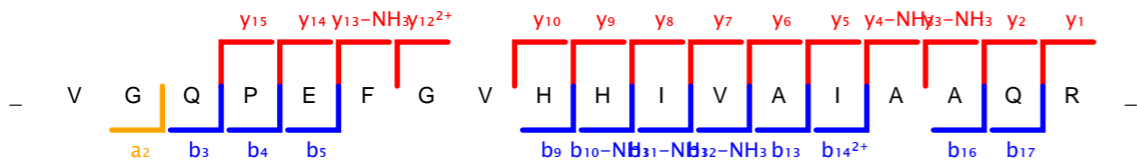
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Scan number: 32465
Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
Peptide score: 89.7
Method: FTMS; HCD



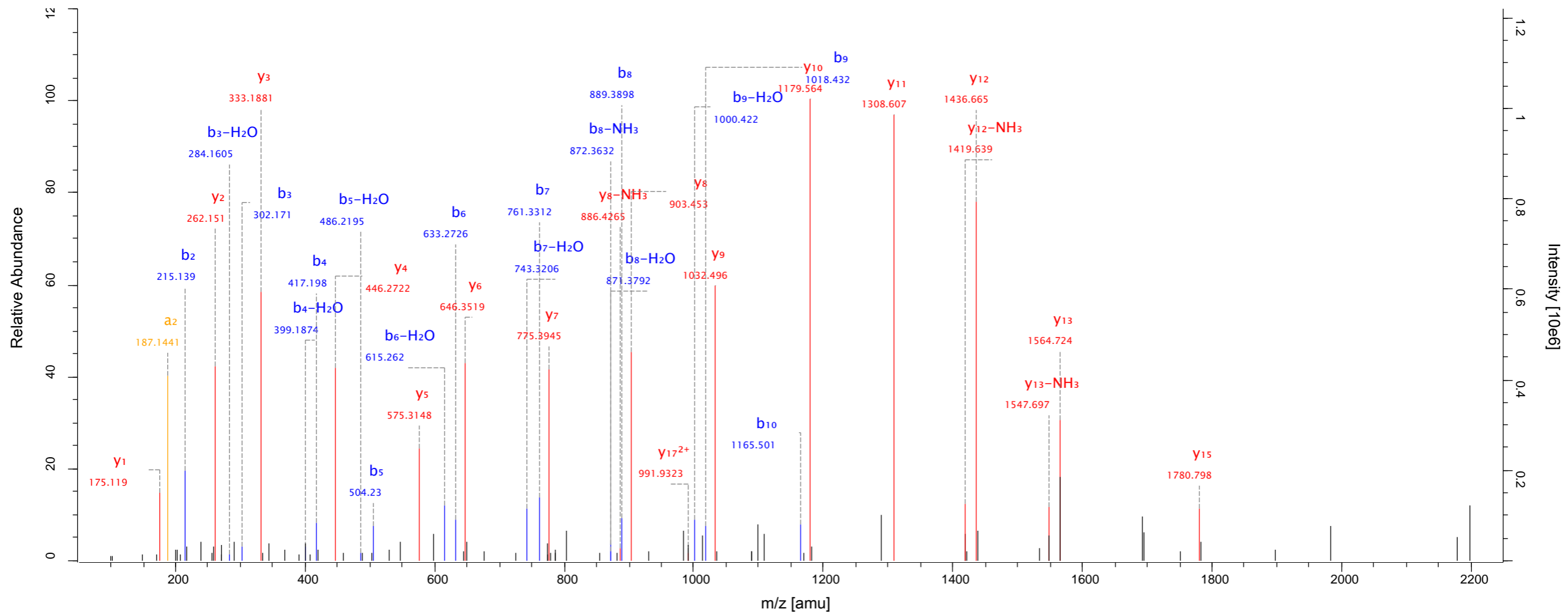
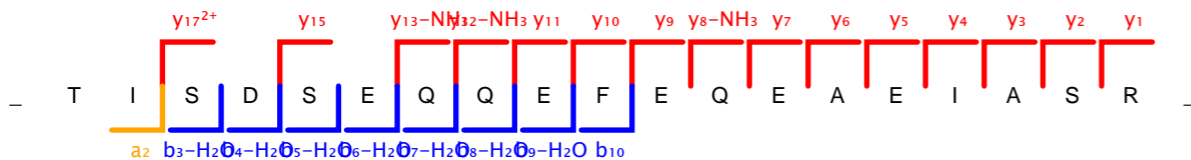
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 32518
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 80.68
Method: FTMS; HCD



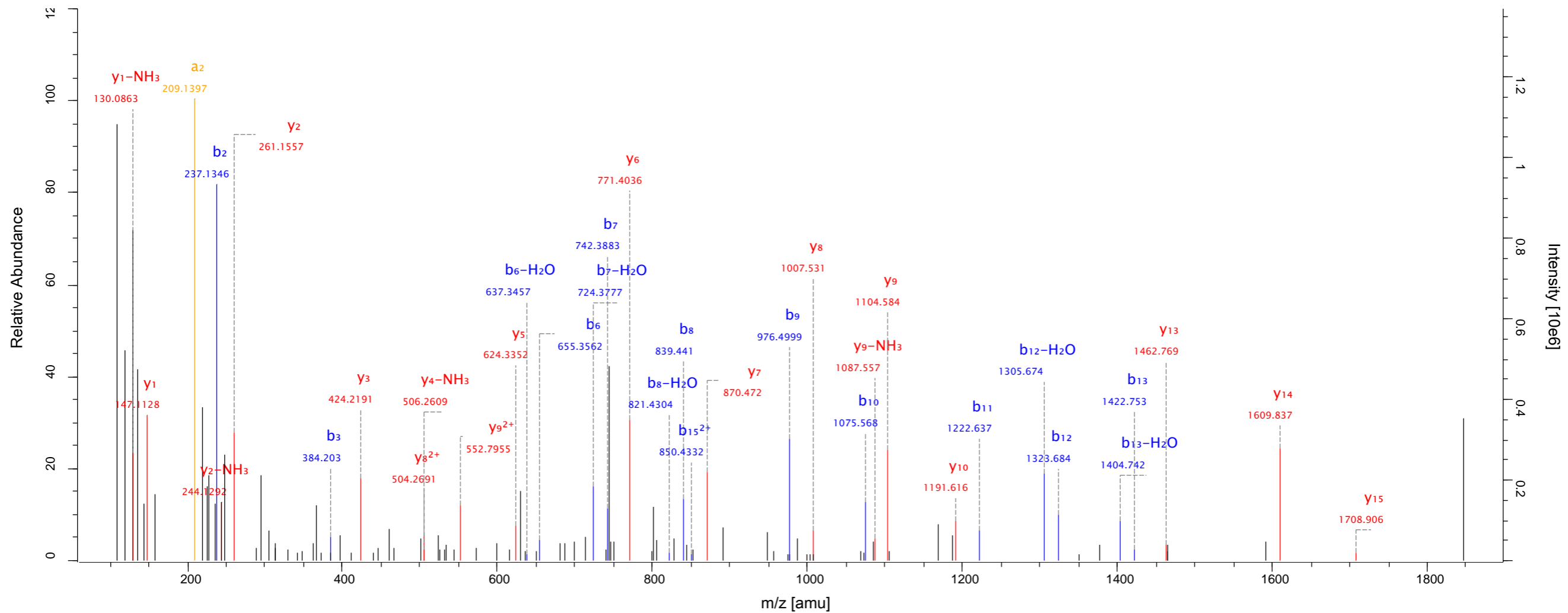
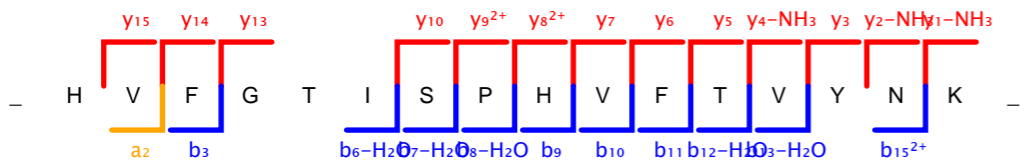
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
 Scan number: 32895
 Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
 Peptide score: 136.35
 Method: FTMS; HCD



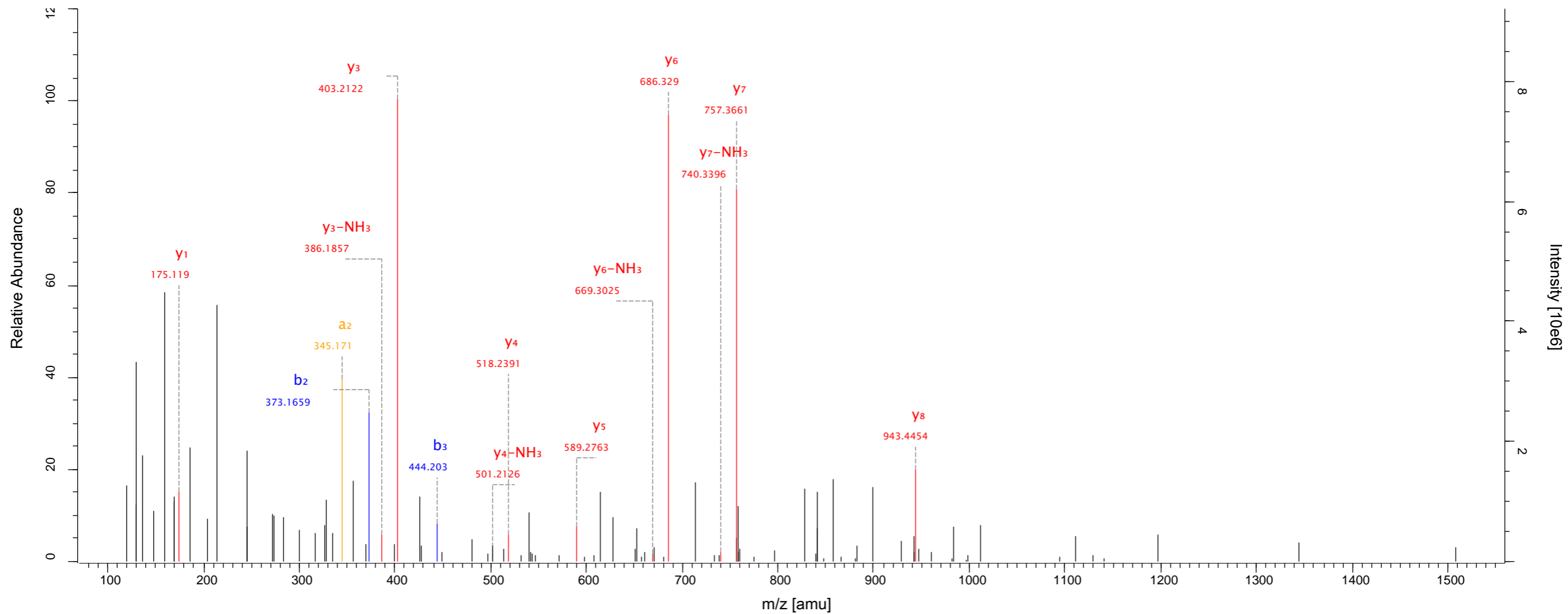
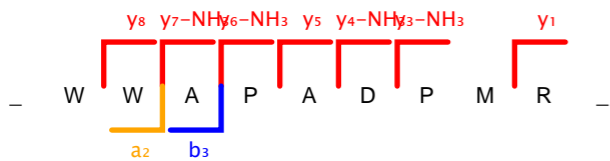
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 33109
 Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>
 Peptide score: 215.79
 Method: FTMS; HCD



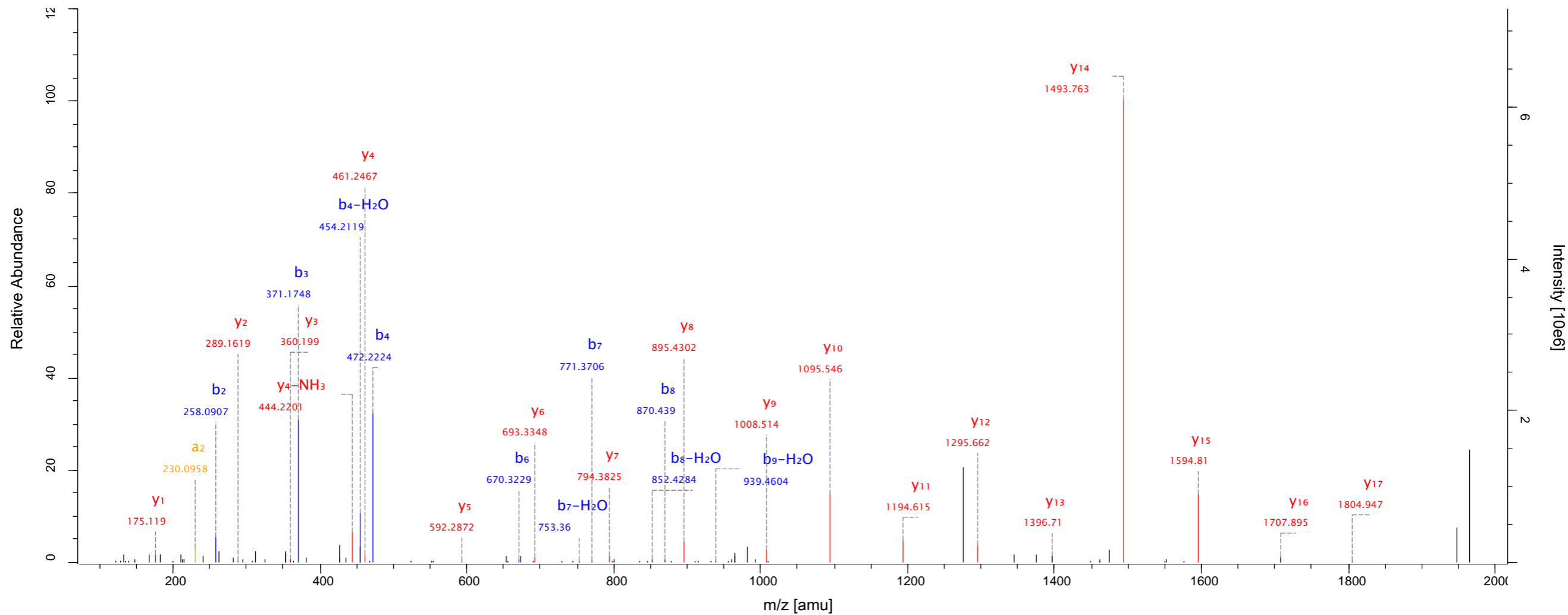
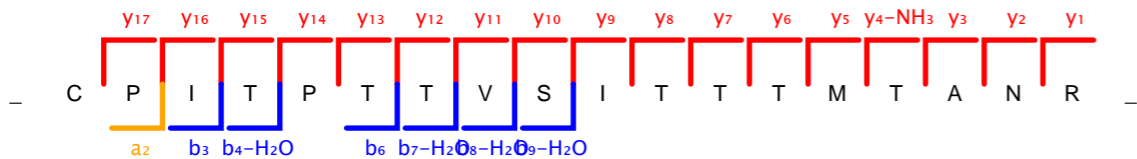
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 34442
 Protein: ORFL154W + NC_006273.2[96843,97115]+ <unknown description>
 Peptide score: 192.18
 Method: FTMS; HCD



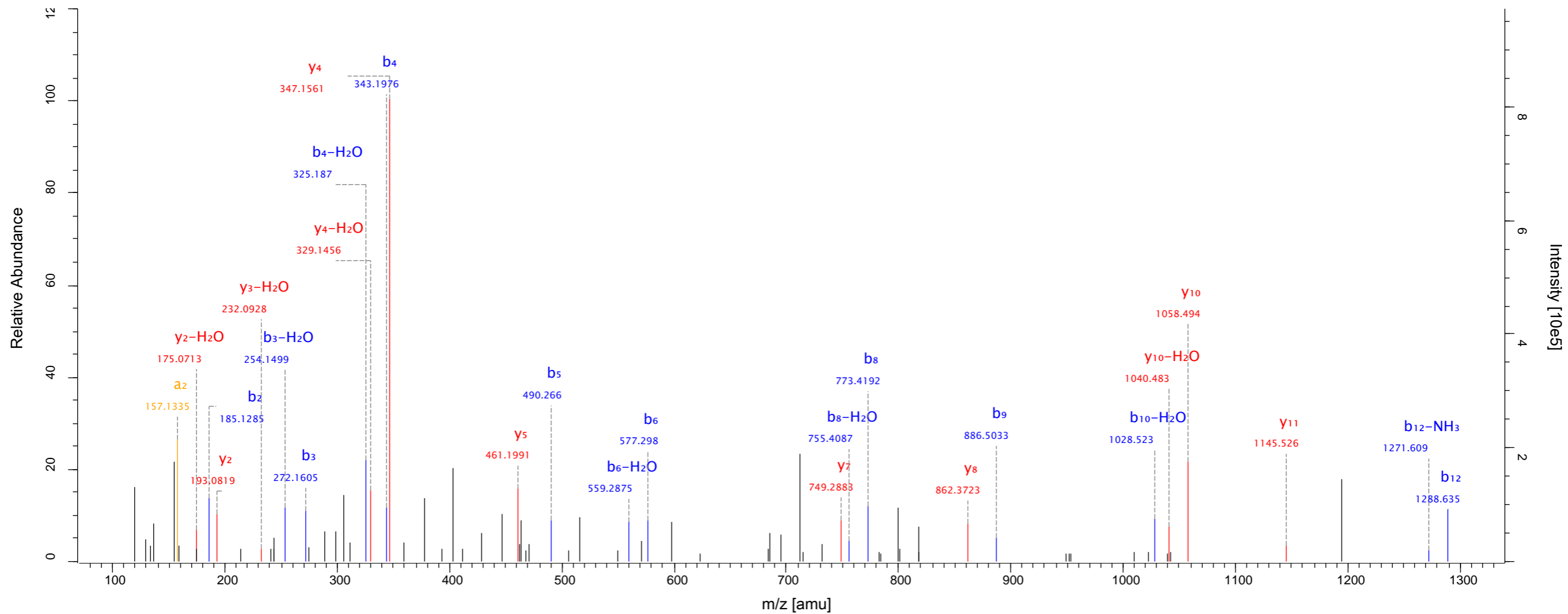
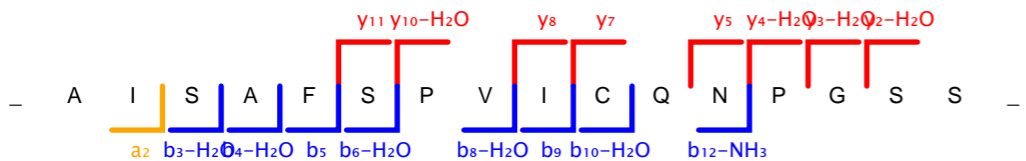
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 34656
Protein: ORFL253W.iORF3 + NC_006273.2[162059,162286]+ <unknown description>
Peptide score: 73.66
Method: FTMS; HCD



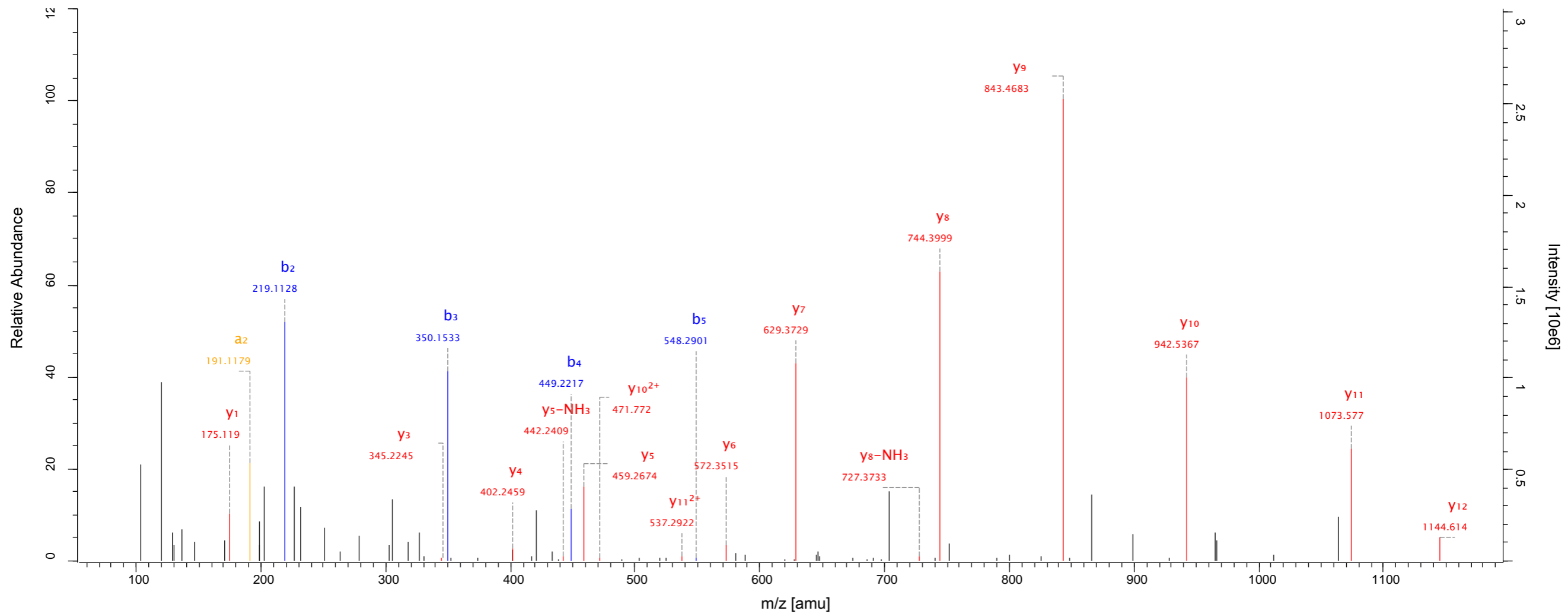
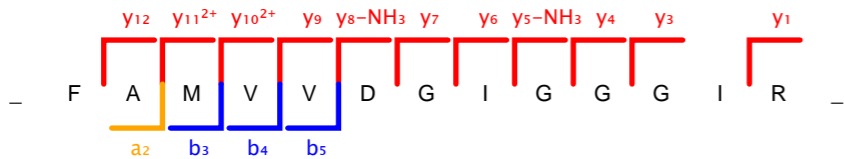
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 35559
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 160.47
Method: FTMS; HCD



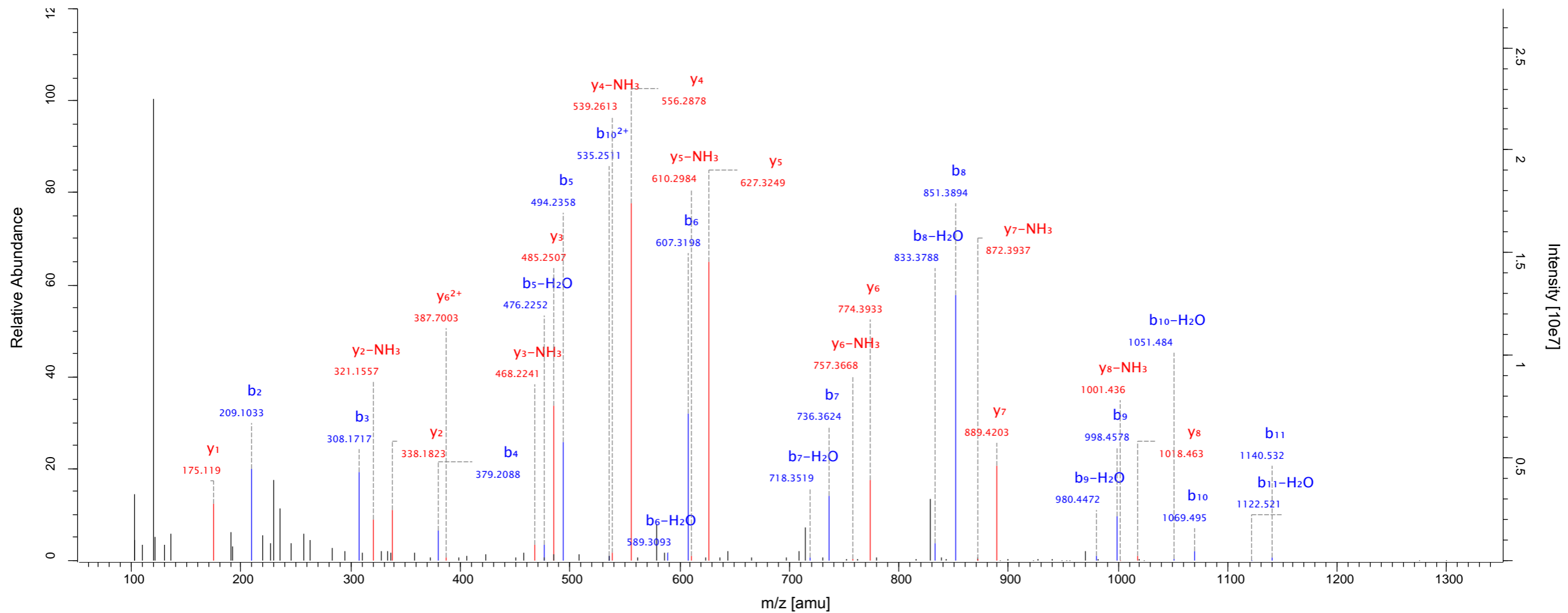
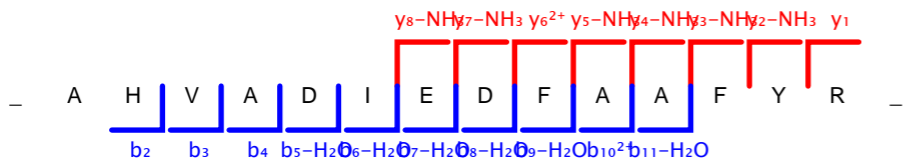
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 38279
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 127.79
 Method: FTMS; HCD



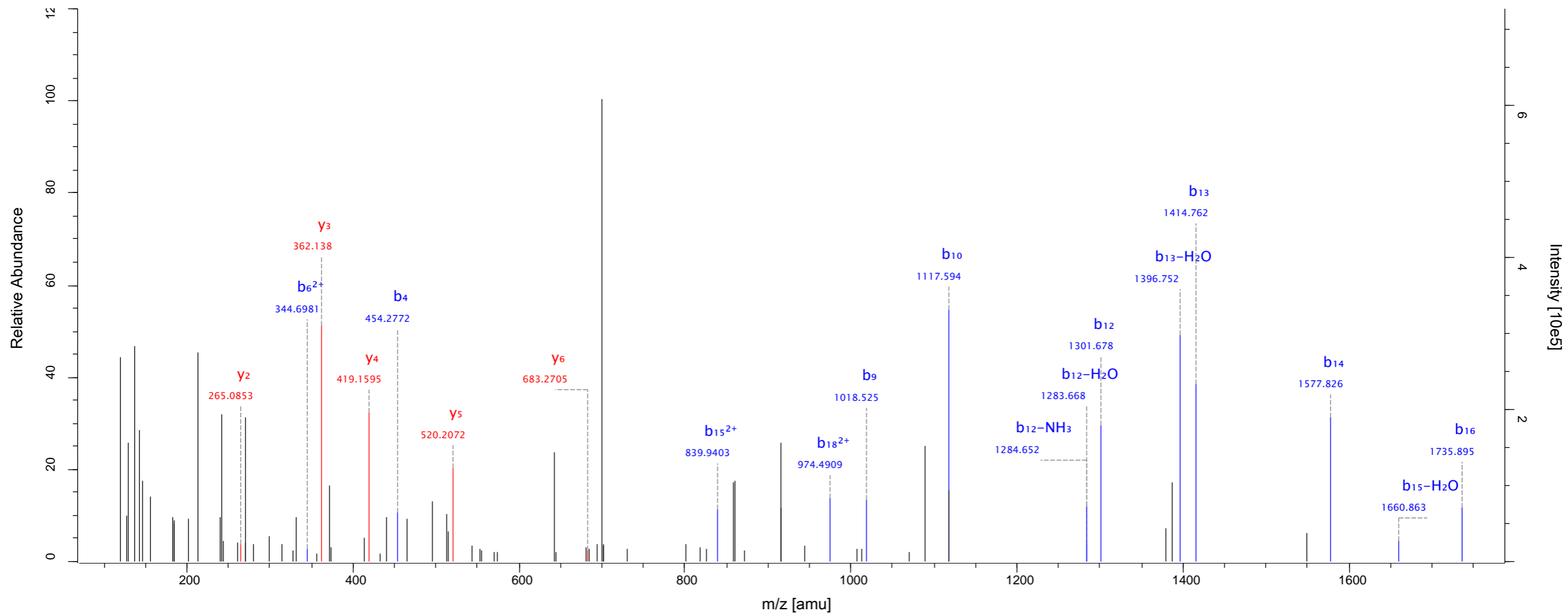
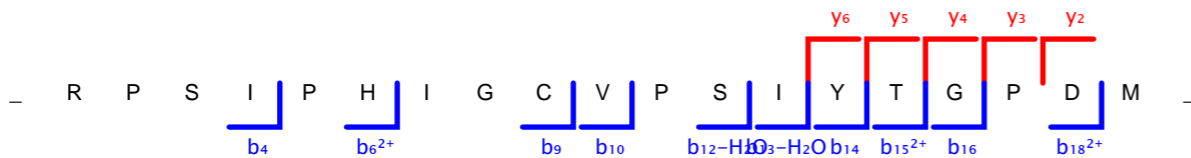
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 41372
Protein: ORFL211W + NC_006273.2[125859,126083]+ <unknown description>
Peptide score: 110.08
Method: FTMS; HCD



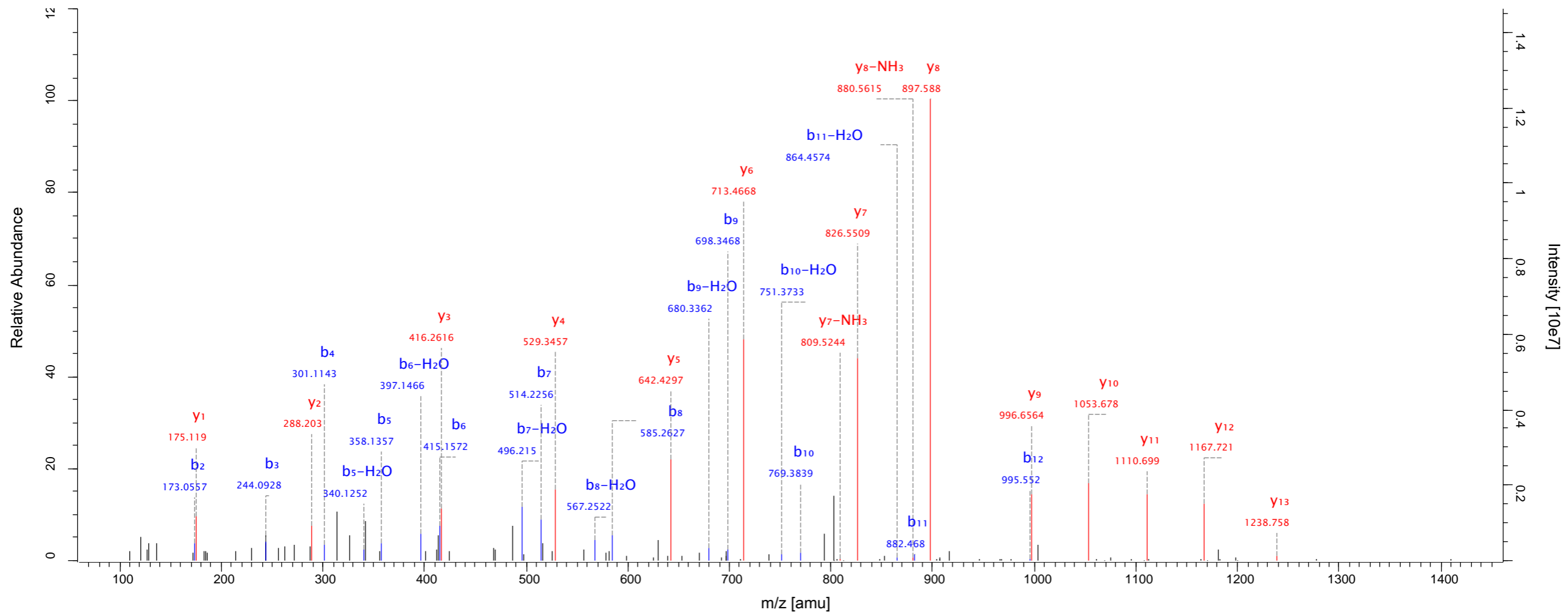
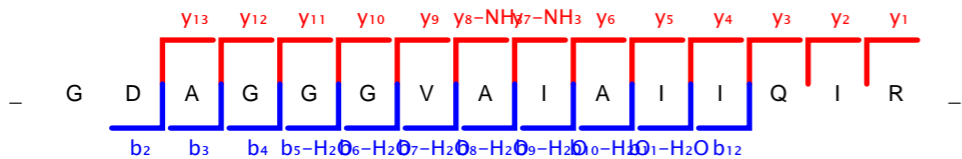
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
 Scan number: 44022
 Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>
 Peptide score: 207.88
 Method: FTMS; HCD



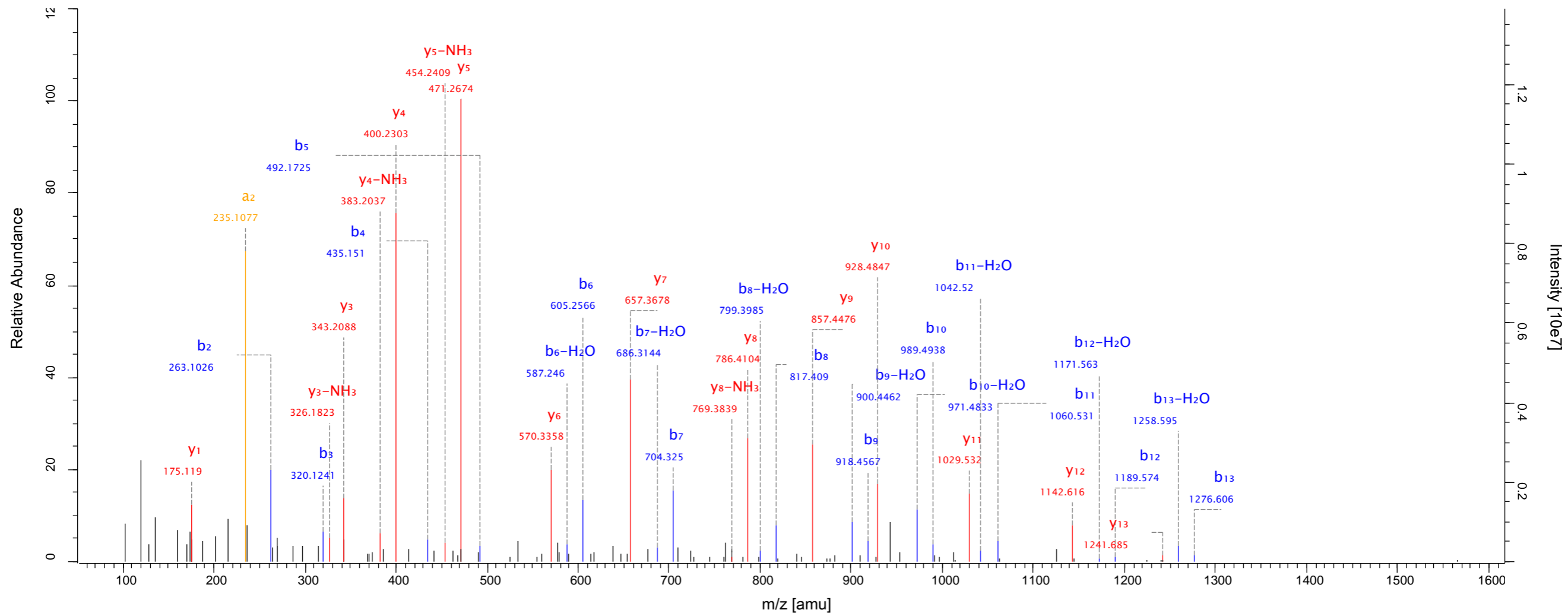
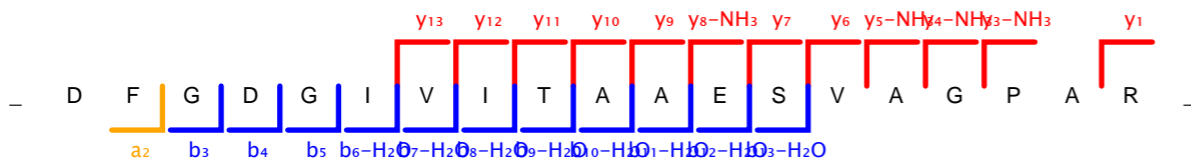
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 44445
 Protein: ORFL172W + NC_006273.2[105961,109027]+ <unknown description>; ORFL159W.iORF2 + NC_006273.2[108854,109027]+ <unknown description>
 Peptide score: 81.43
 Method: FTMS; HCD



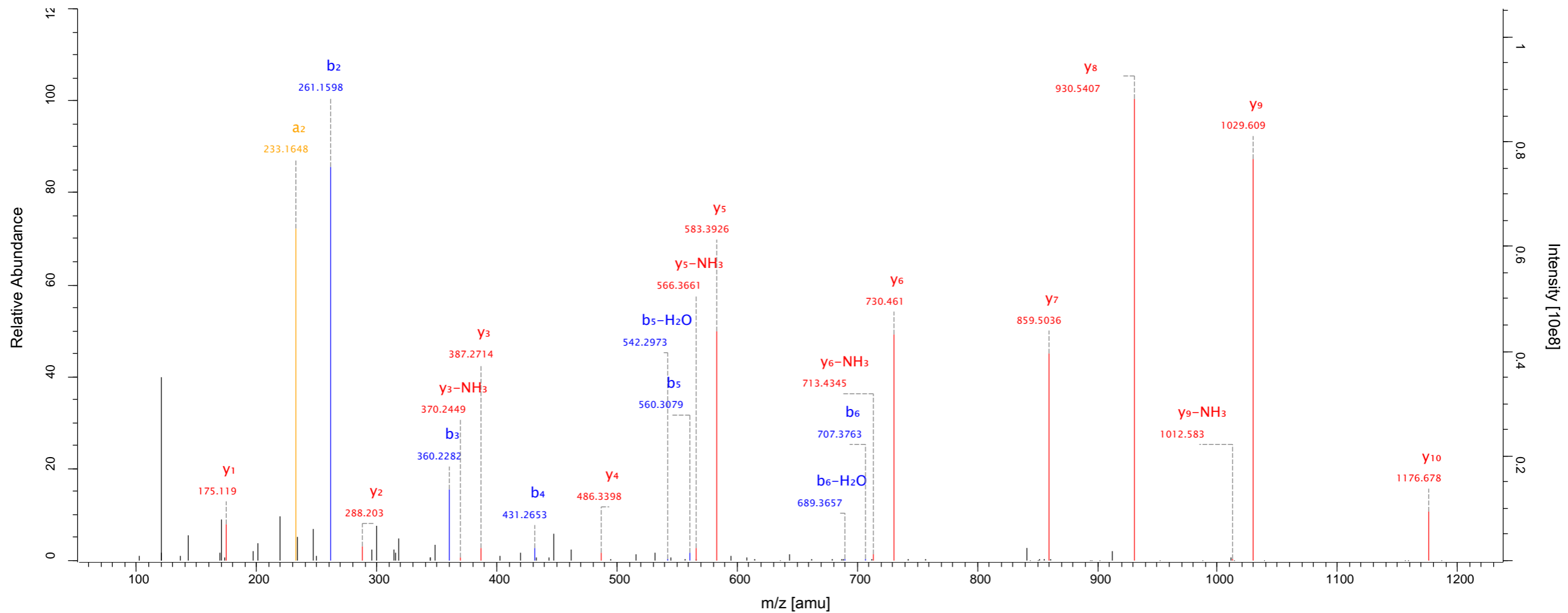
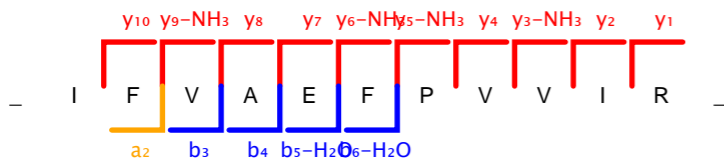
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_10
 Scan number: 50163
 Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
 Peptide score: 212.83
 Method: FTMS; HCD



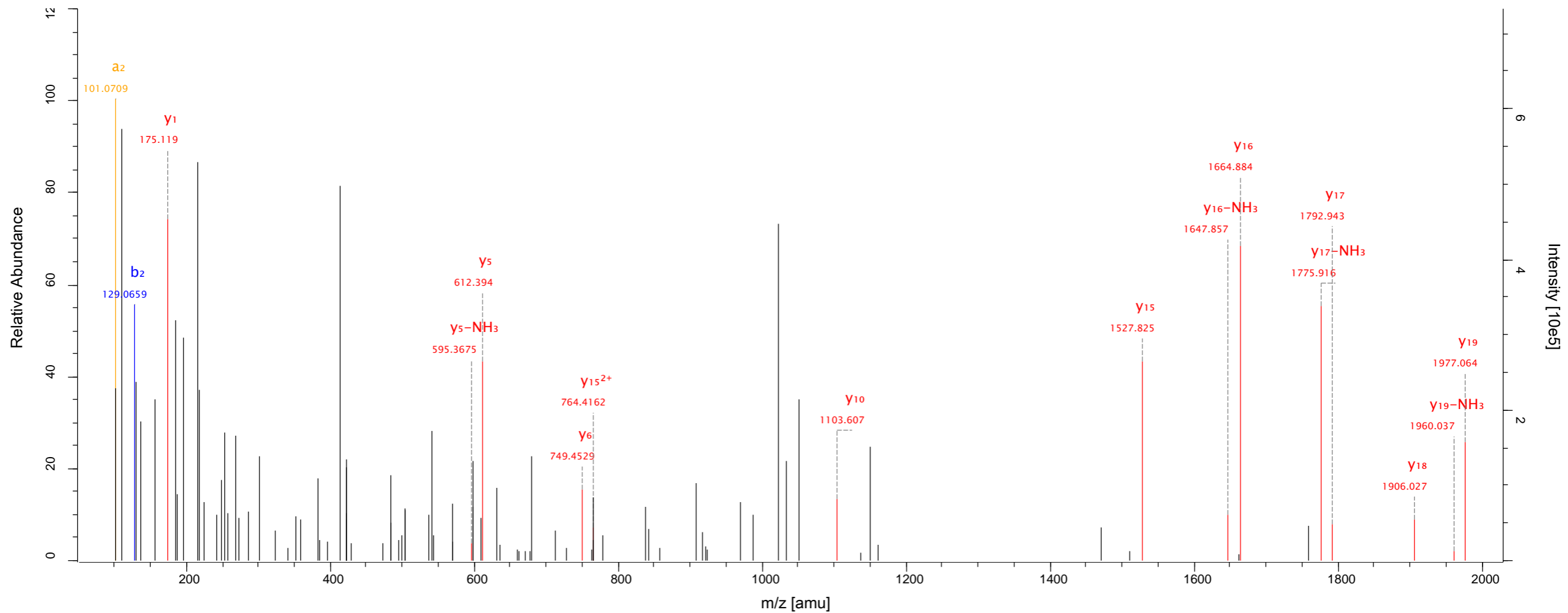
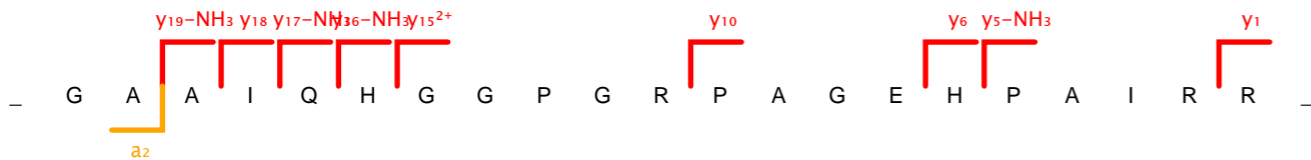
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 50473
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 256.26
 Method: FTMS; HCD



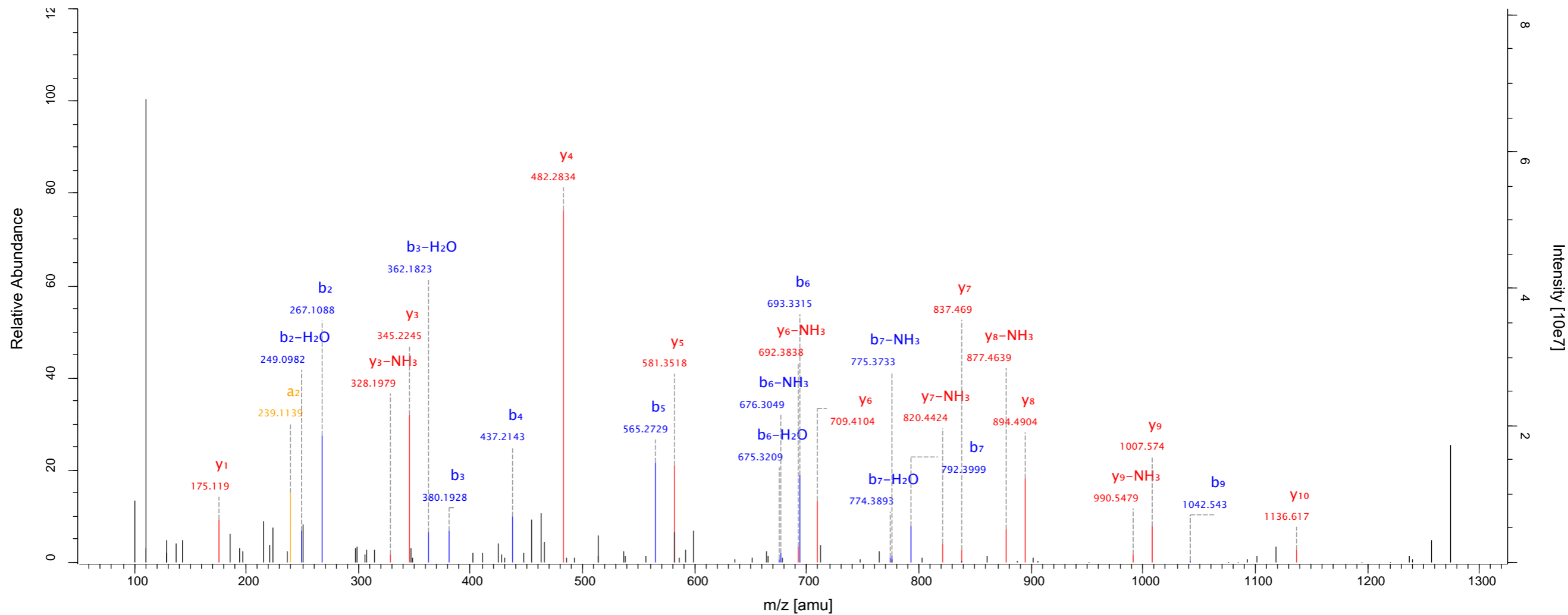
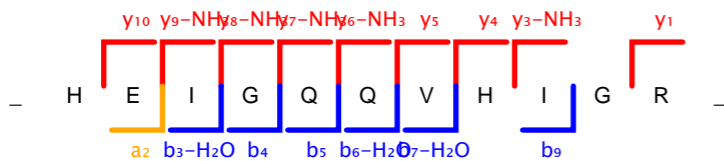
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
Scan number: 50512
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 140.16
Method: FTMS; HCD



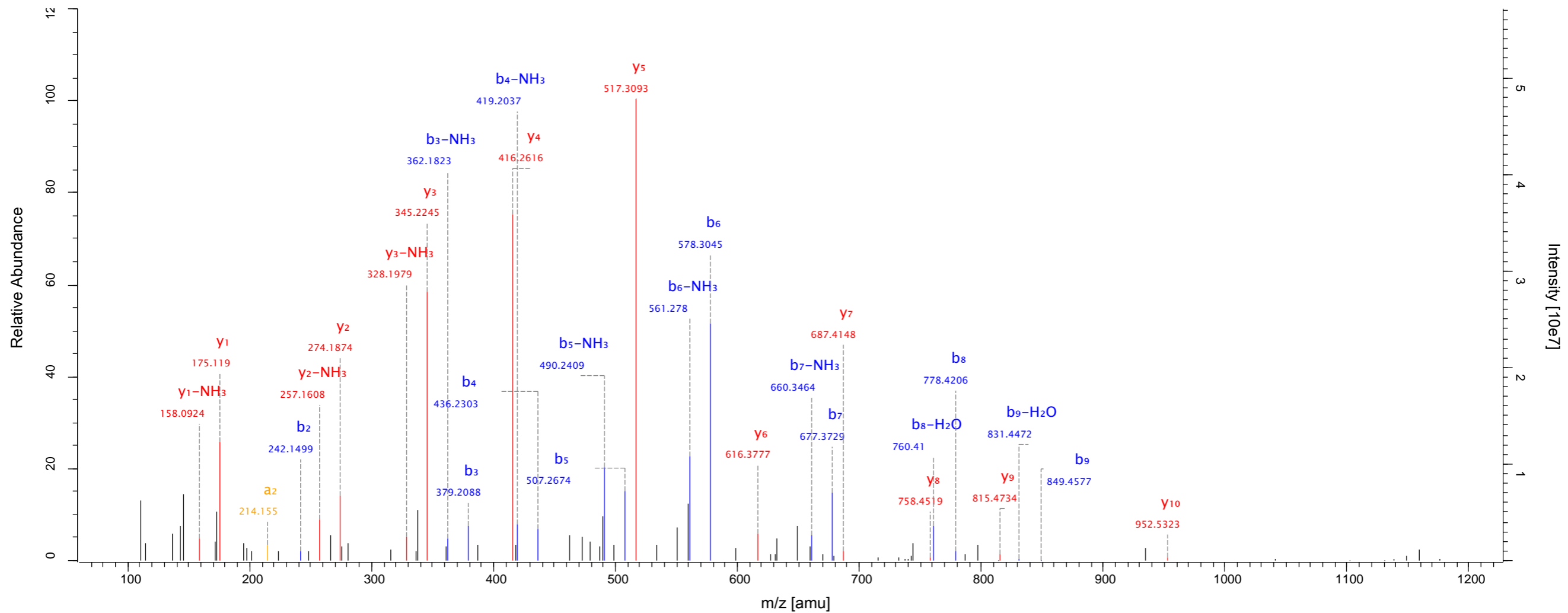
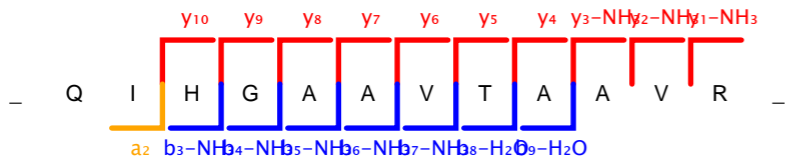
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_10
 Scan number: 8645
 Protein: ORFS359W + NC_006273.2[221986,222303]+ <unknown description>
 Peptide score: 82.01
 Method: FTMS; HCD



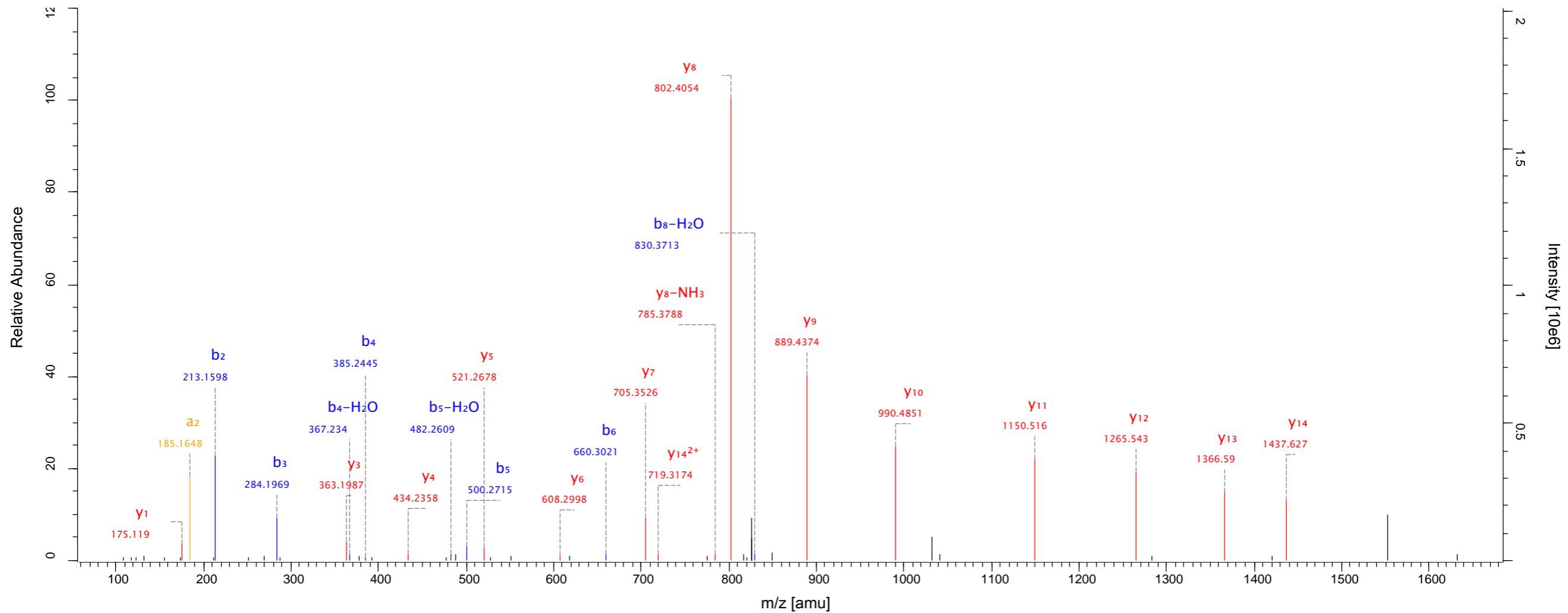
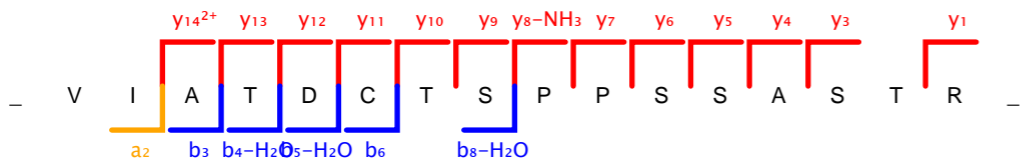
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 11282
Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
Peptide score: 185.77
Method: FTMS; HCD



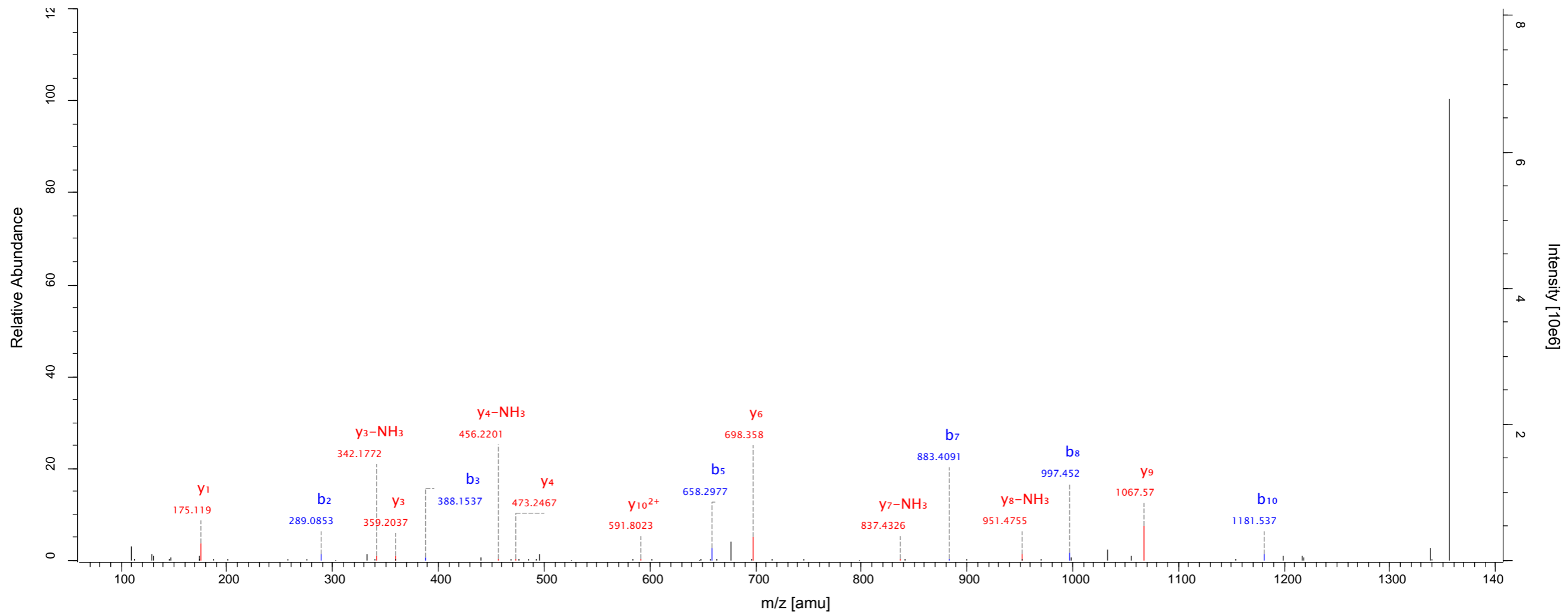
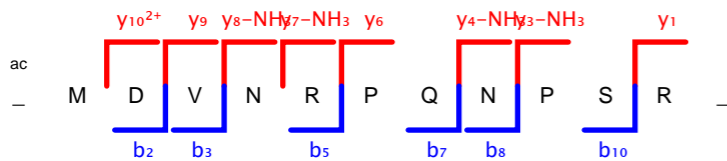
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_11
 Scan number: 11667
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 179.19
 Method: FTMS; HCD



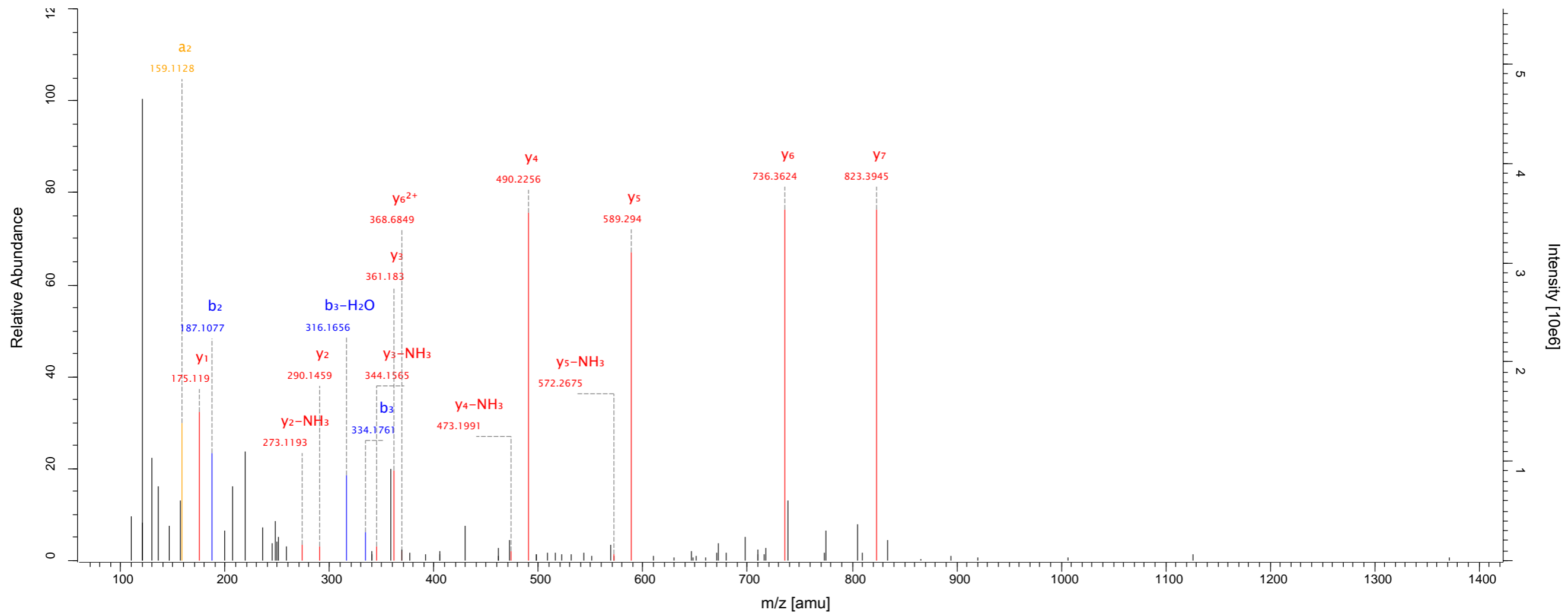
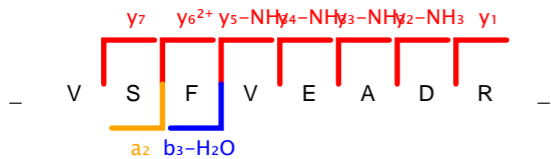
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 Scan number: 12708
 Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
 Peptide score: 164.33
 Method: FTMS; HCD



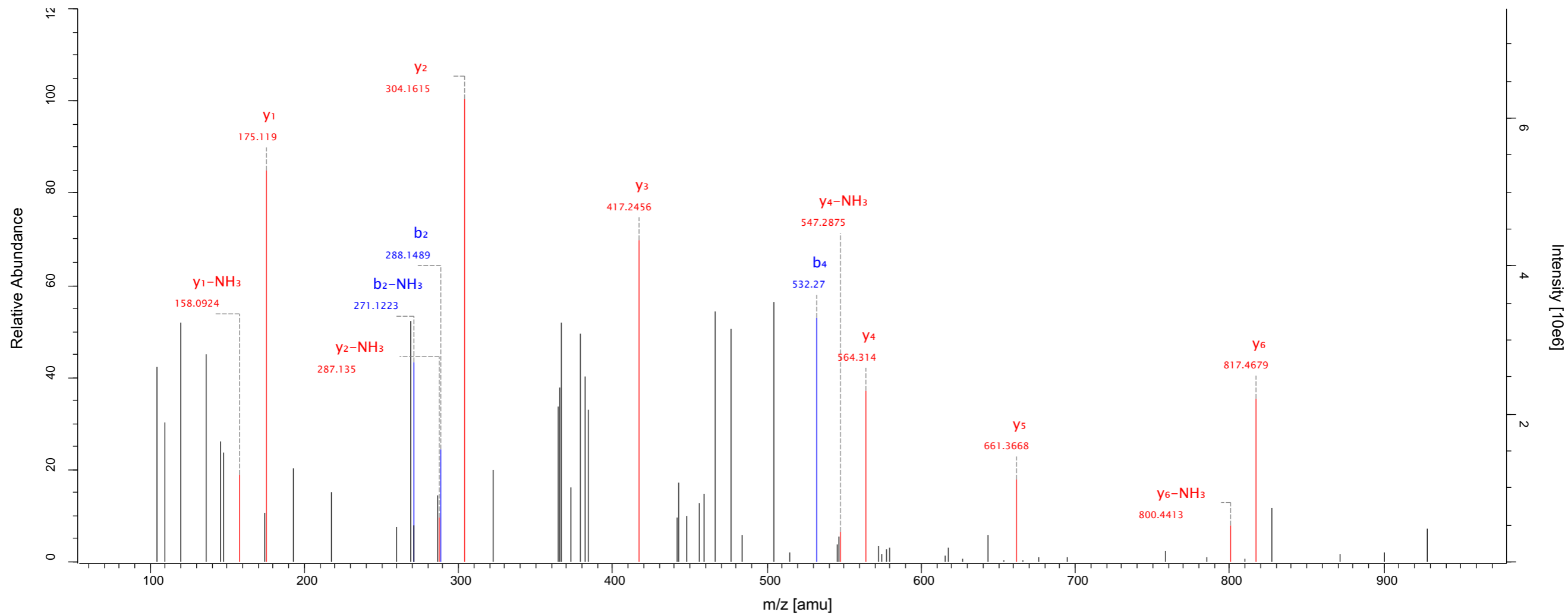
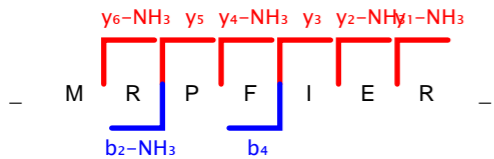
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 12833
Protein: ORFL52C - NC_006273.2[25013,25198]- <unknown description>
Peptide score: 93.26
Method: FTMS; HCD



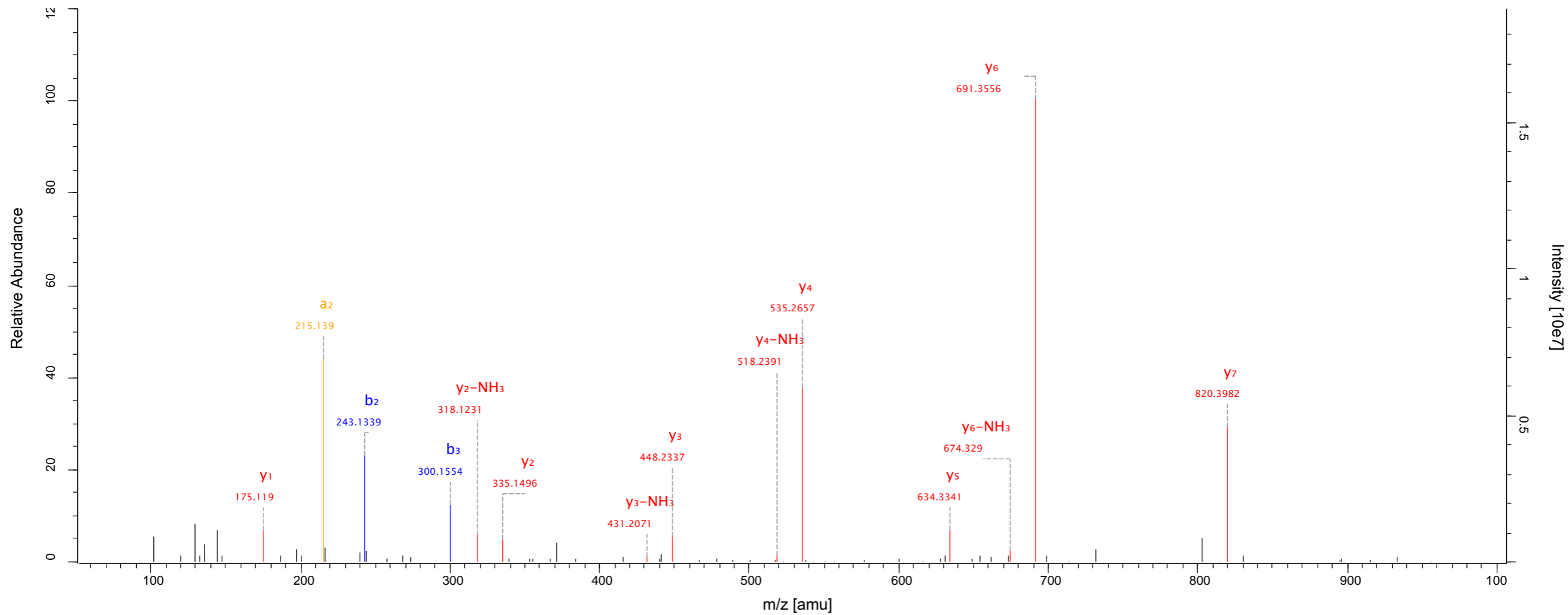
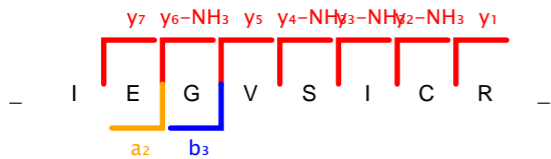
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 15475
Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>; ORFL27C.iORF1 - NC_006273.2[7912,8184]- <unknown description>
Peptide score: 108.09
Method: FTMS; HCD



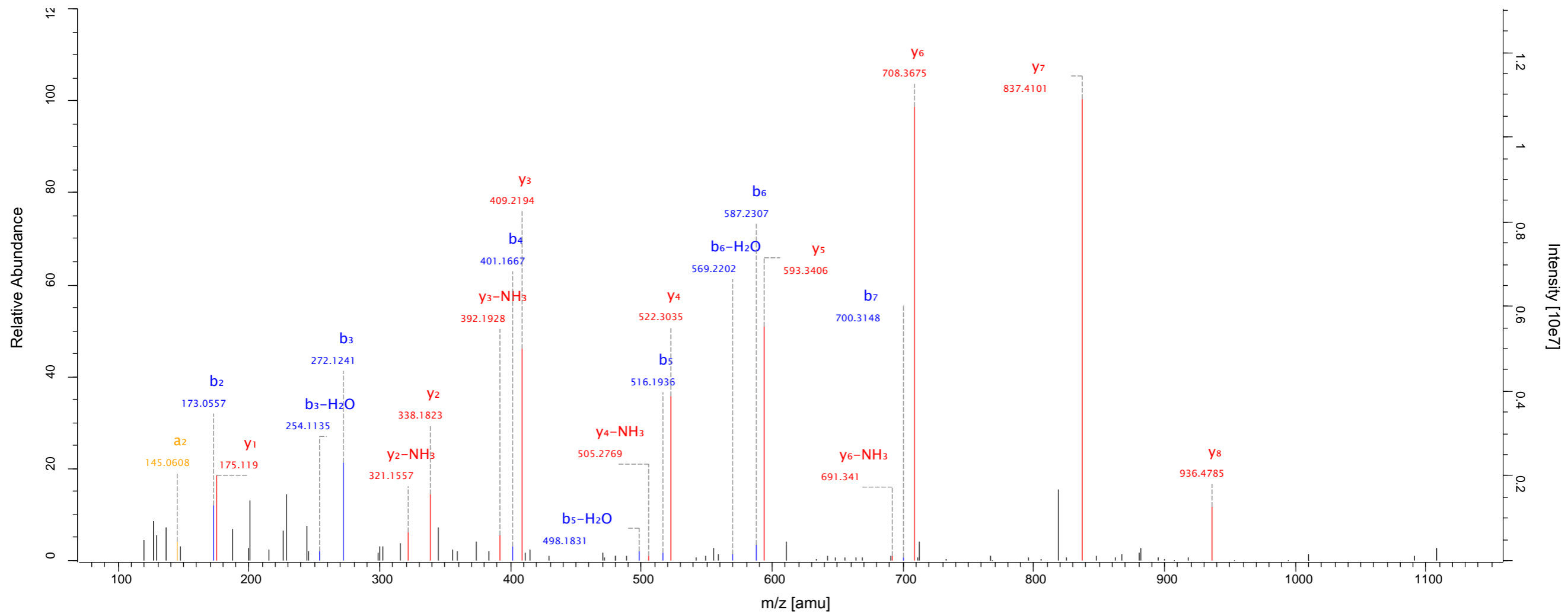
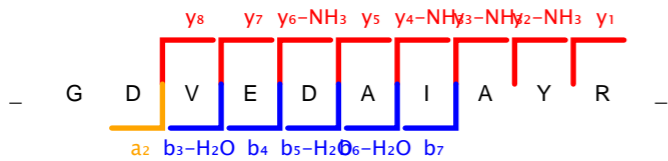
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
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Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 97.07
Method: FTMS; HCD



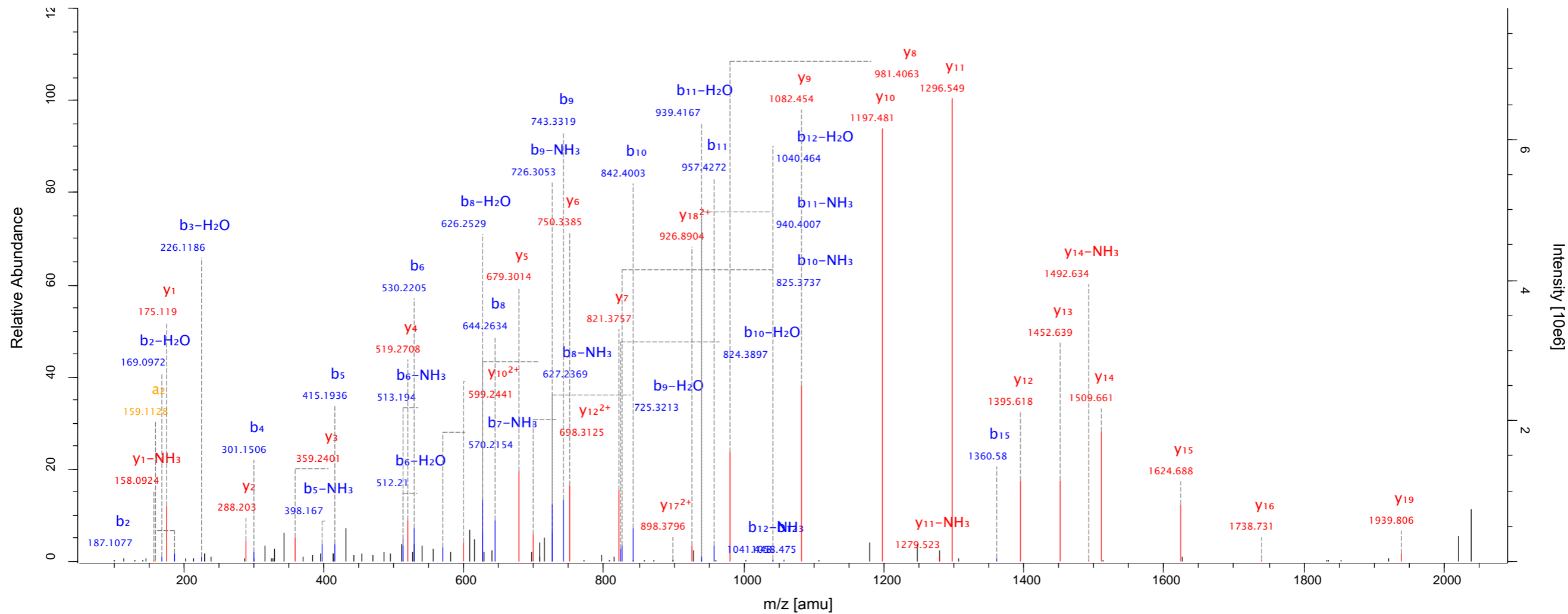
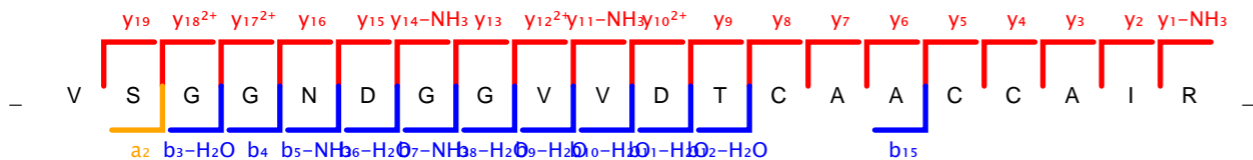
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Peptide score: 126.67
Method: FTMS; HCD



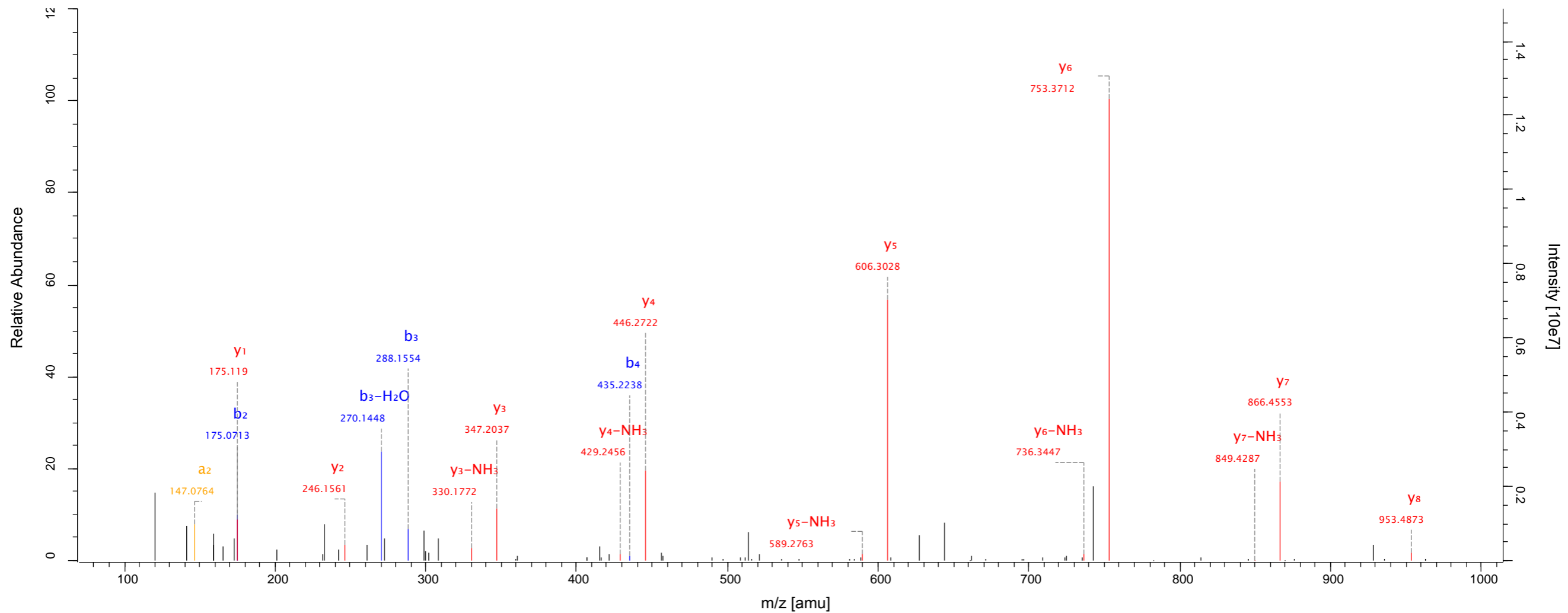
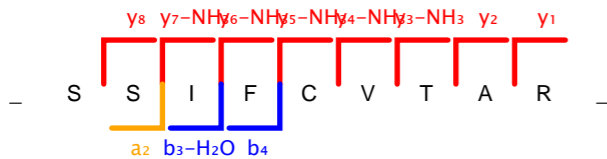
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 20789
Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
Peptide score: 136.99
Method: FTMS; HCD



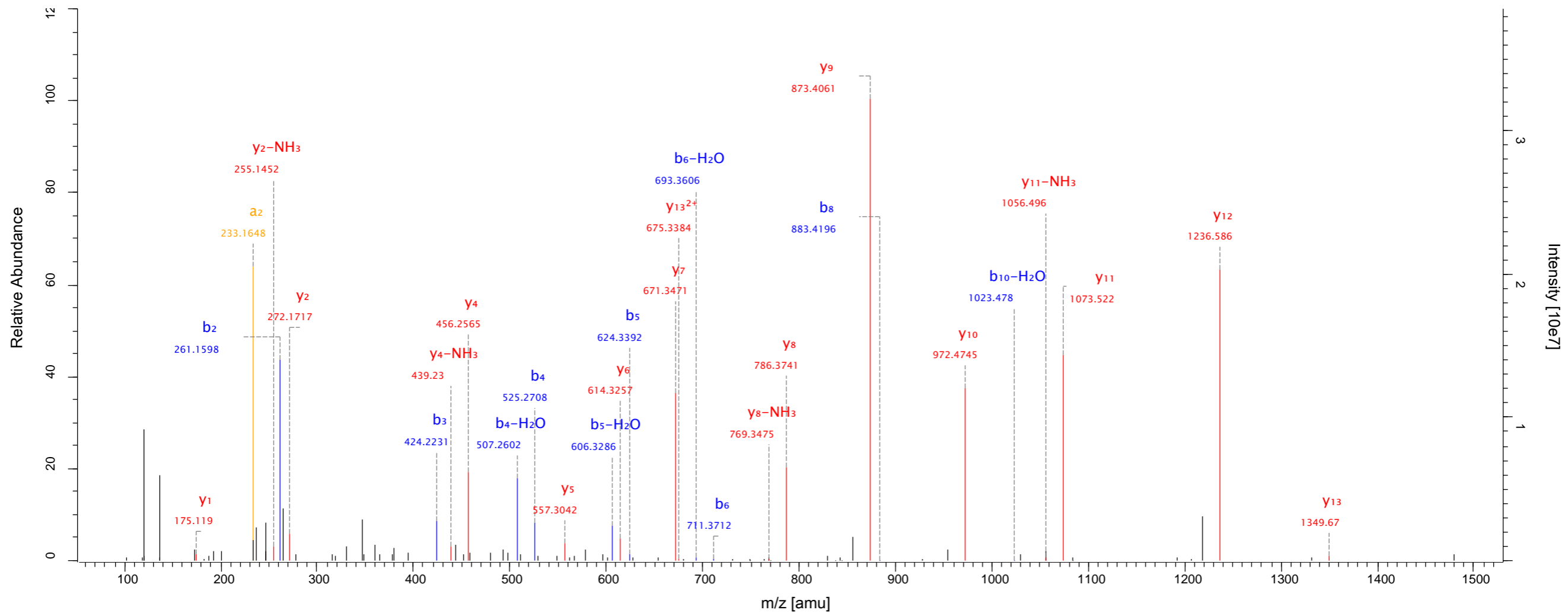
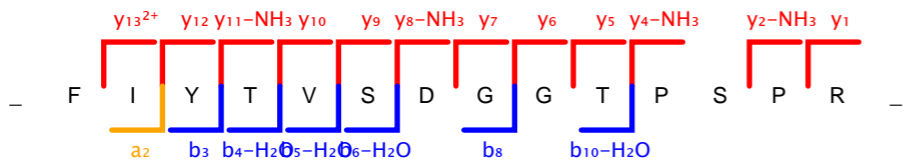
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 Scan number: 22830
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 295.43
 Method: FTMS; HCD



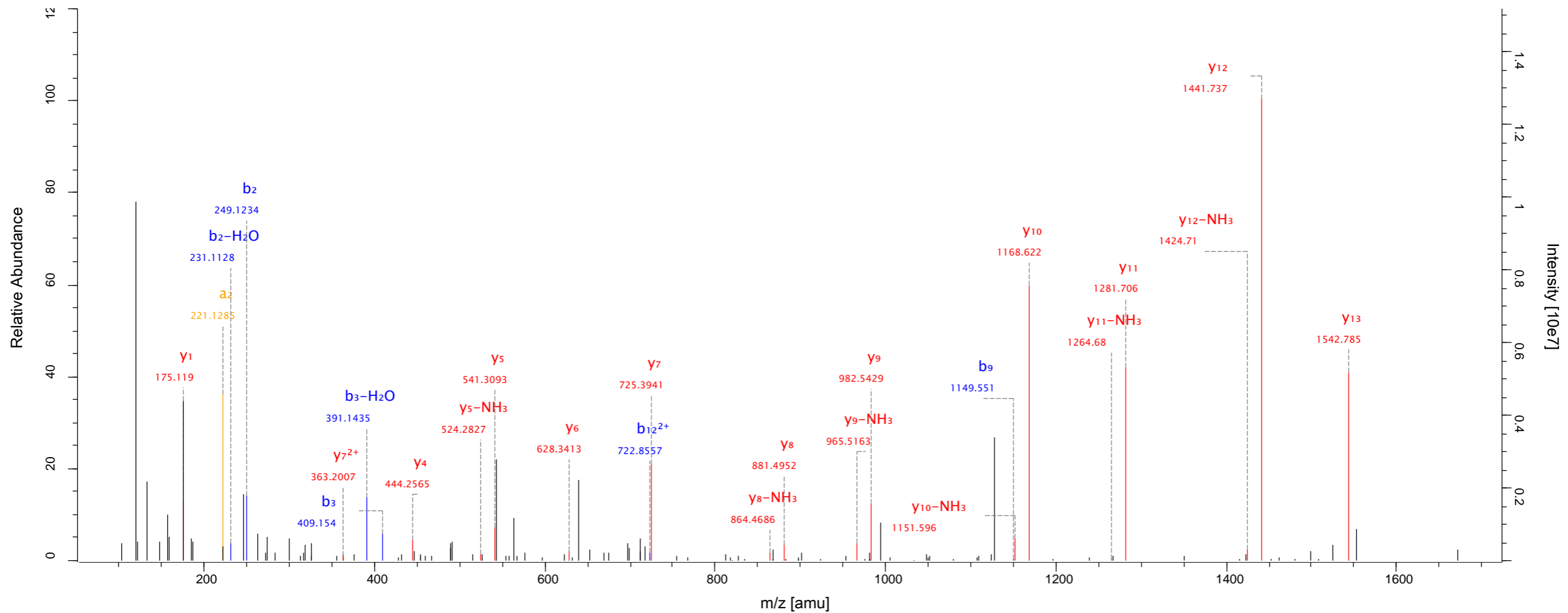
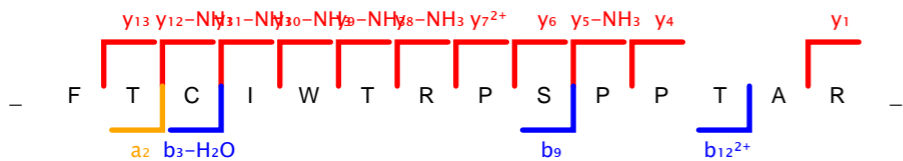
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 25209
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 122.19
Method: FTMS; HCD



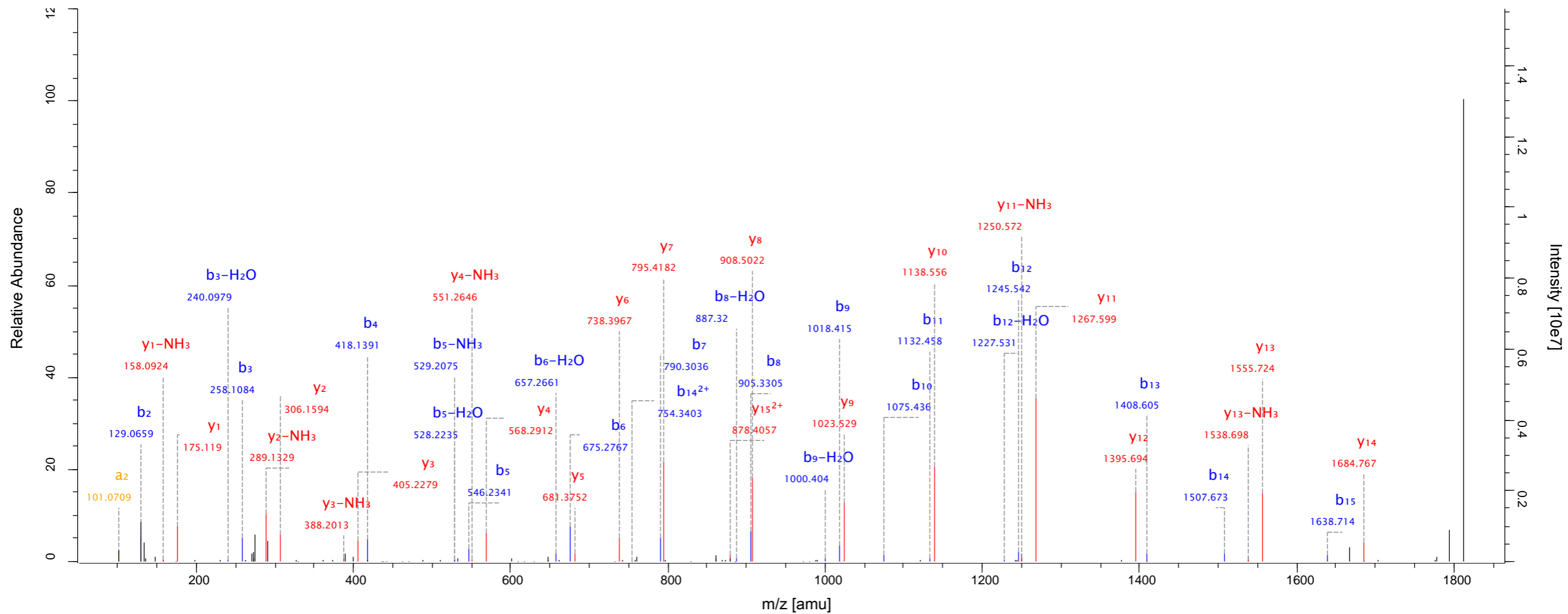
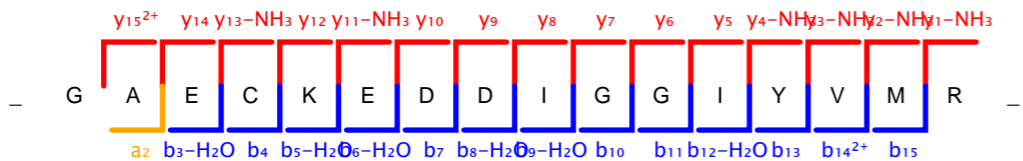
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 25365
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 162.49
 Method: FTMS; HCD



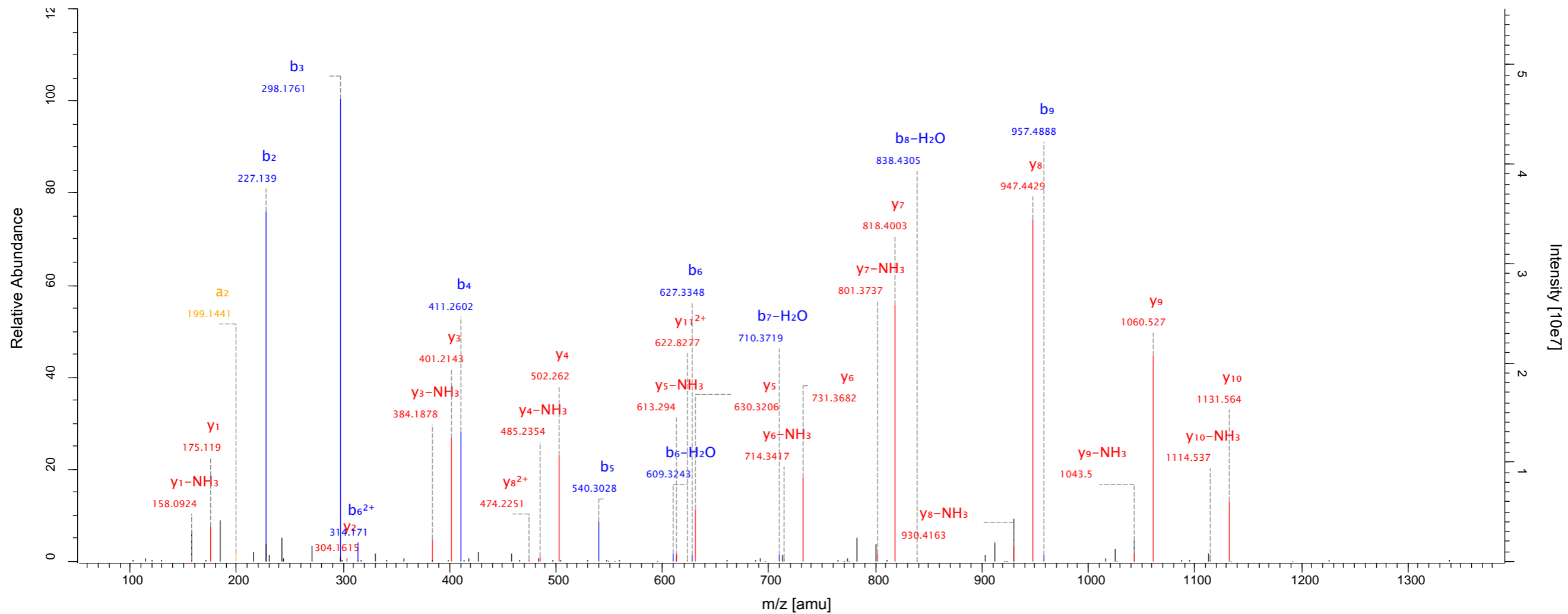
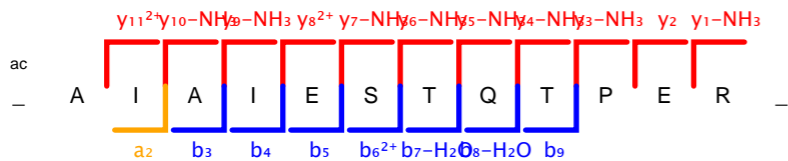
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 27975
 Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
 Peptide score: 134.99
 Method: FTMS; HCD



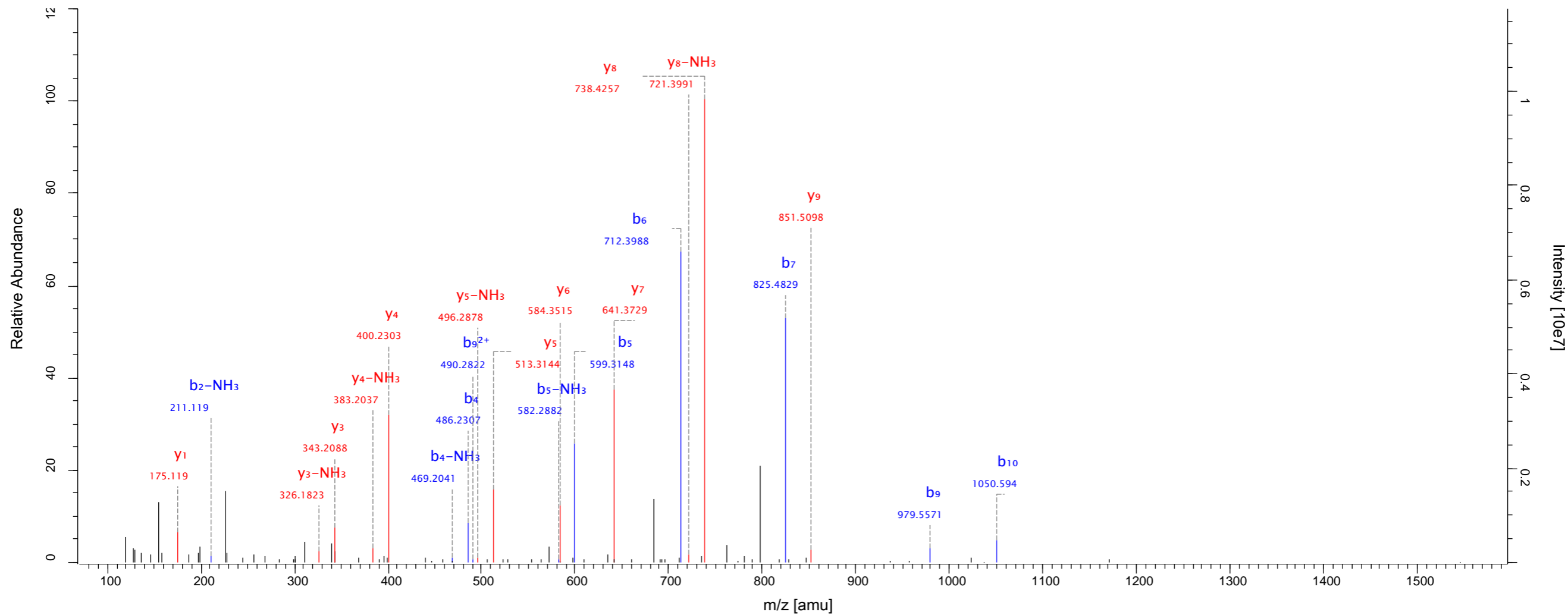
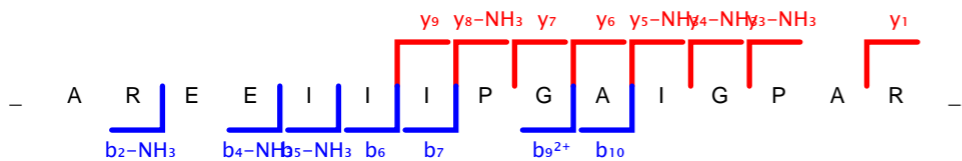
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 29833
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 266.28
 Method: FTMS; HCD



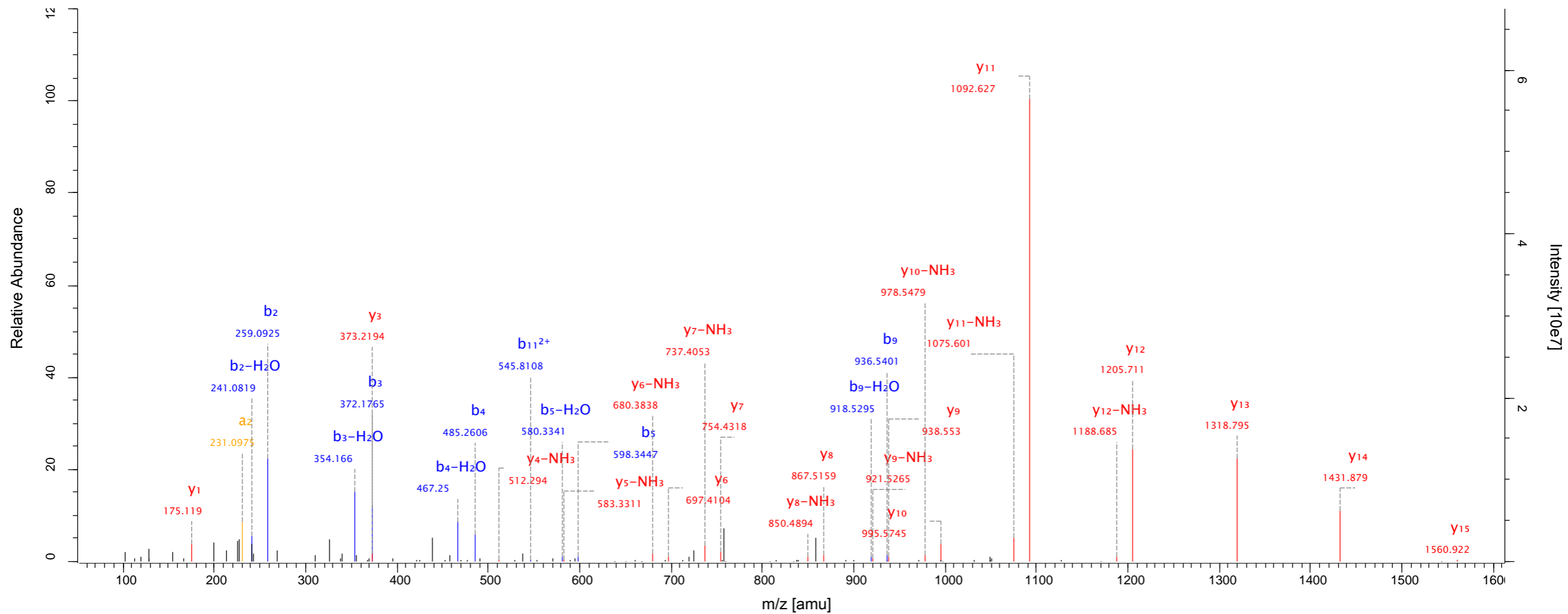
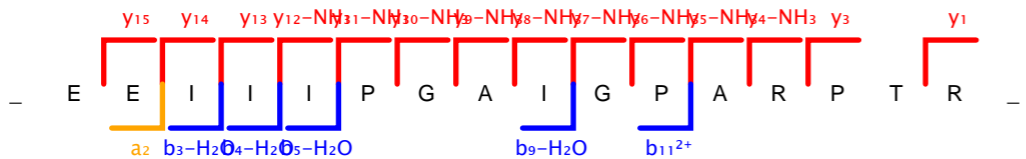
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 30525
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 206.73
 Method: FTMS; HCD



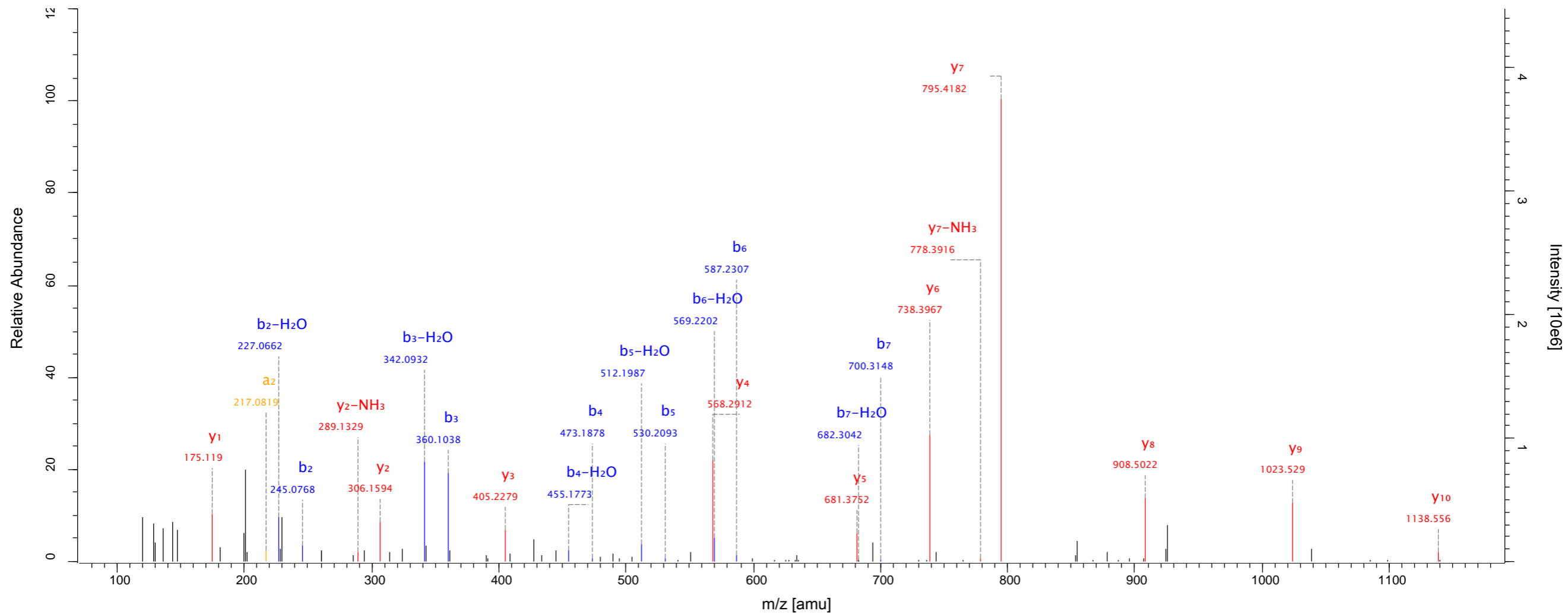
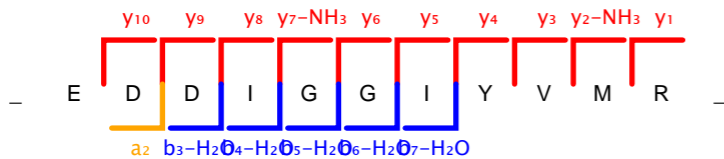
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 33549
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 103.21
Method: FTMS; HCD



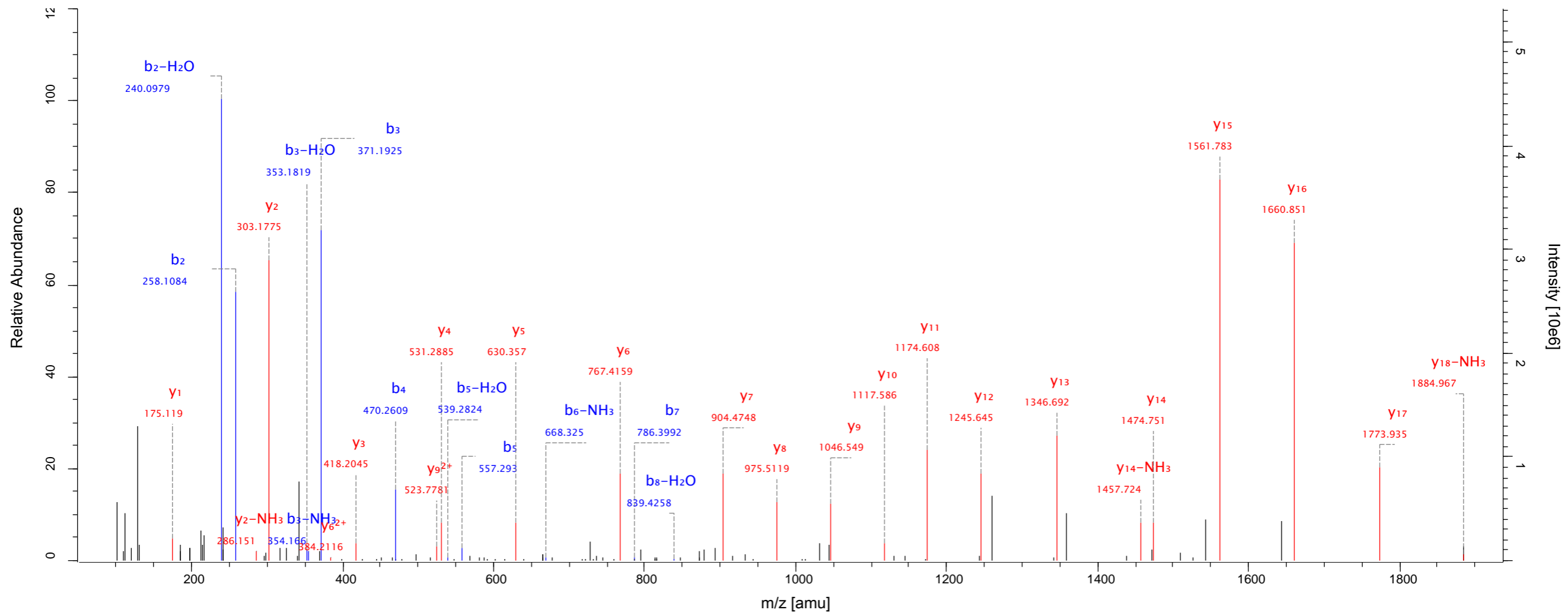
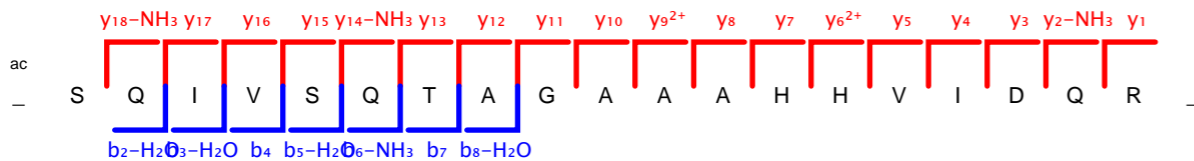
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_11
 Scan number: 34783
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 235.73
 Method: FTMS; HCD



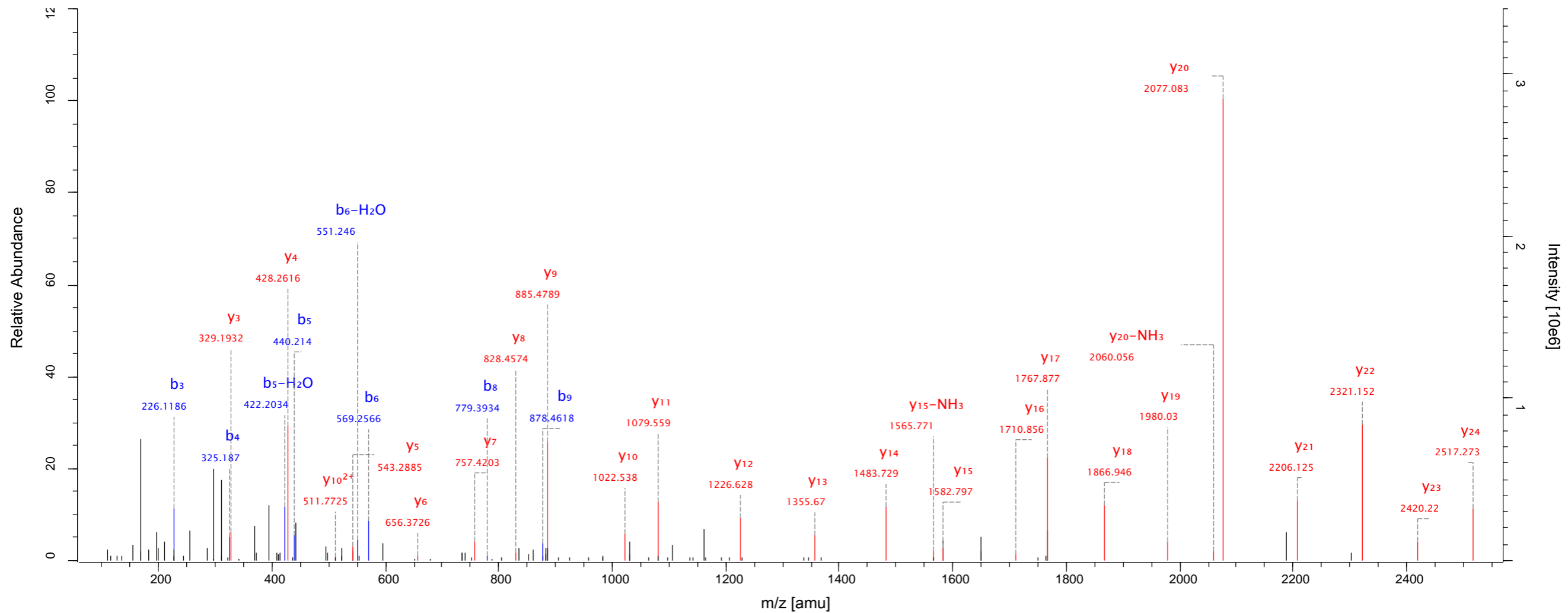
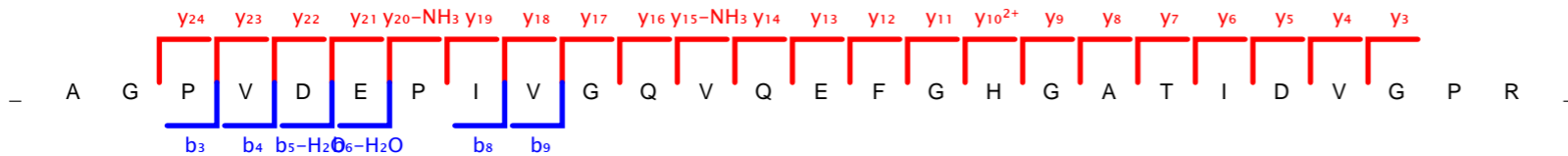
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_11
Scan number: 35425
Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
Peptide score: 143.96
Method: FTMS; HCD



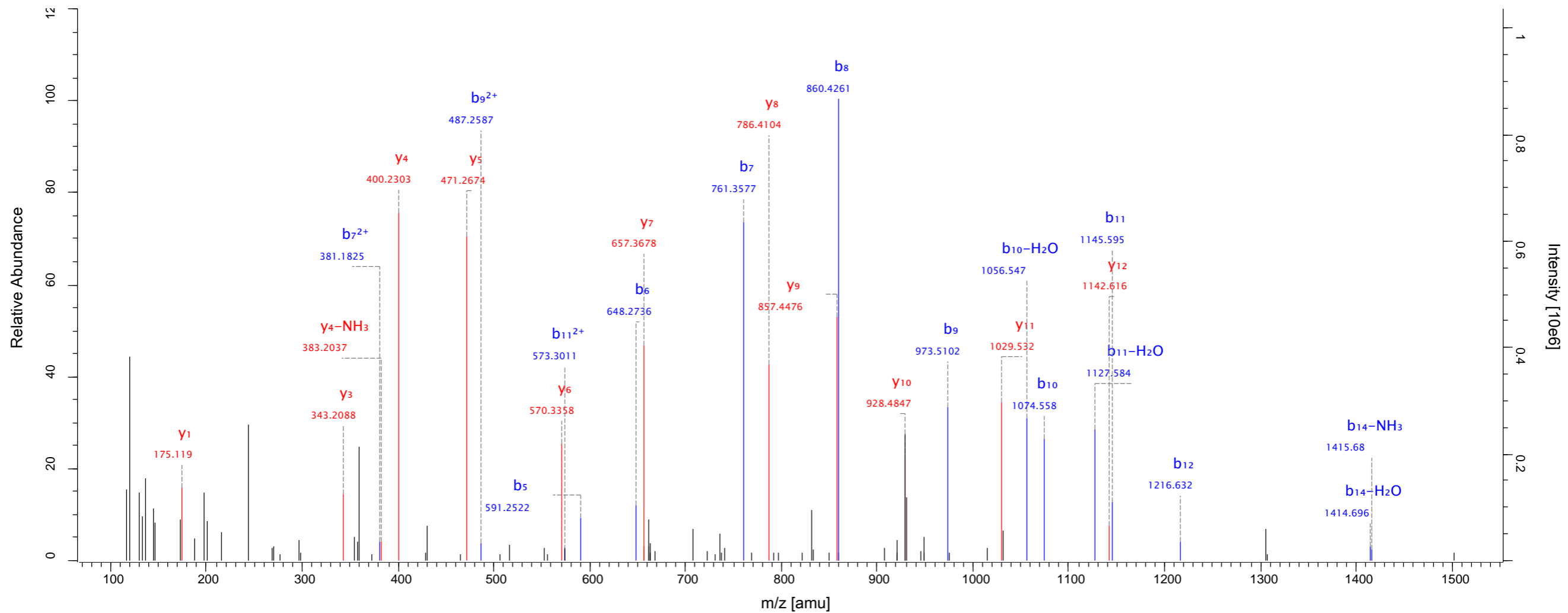
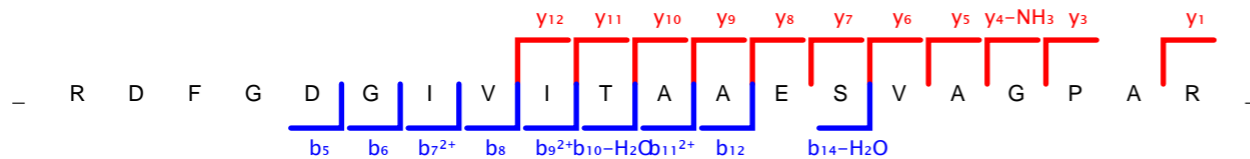
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_11
 Scan number: 39148
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 161.32
 Method: FTMS; HCD



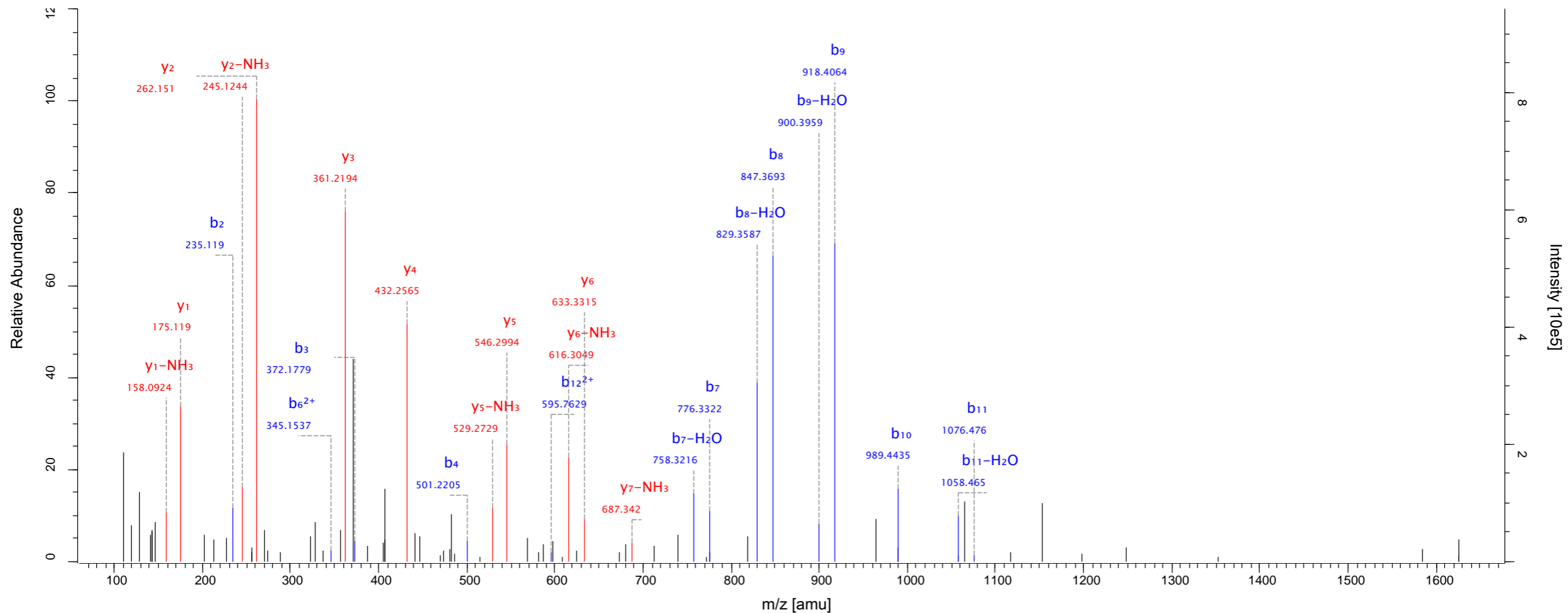
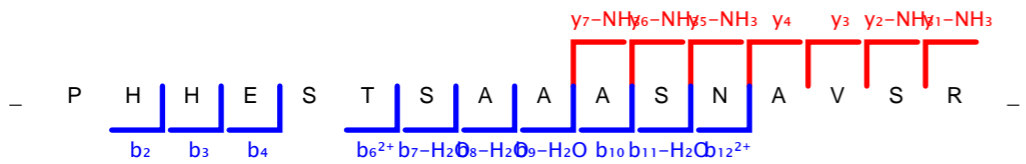
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 44065
 Protein: ORFL118C - NC_006273.2[61896,62396]- <unknown description>
 Peptide score: 169.06
 Method: FTMS; HCD



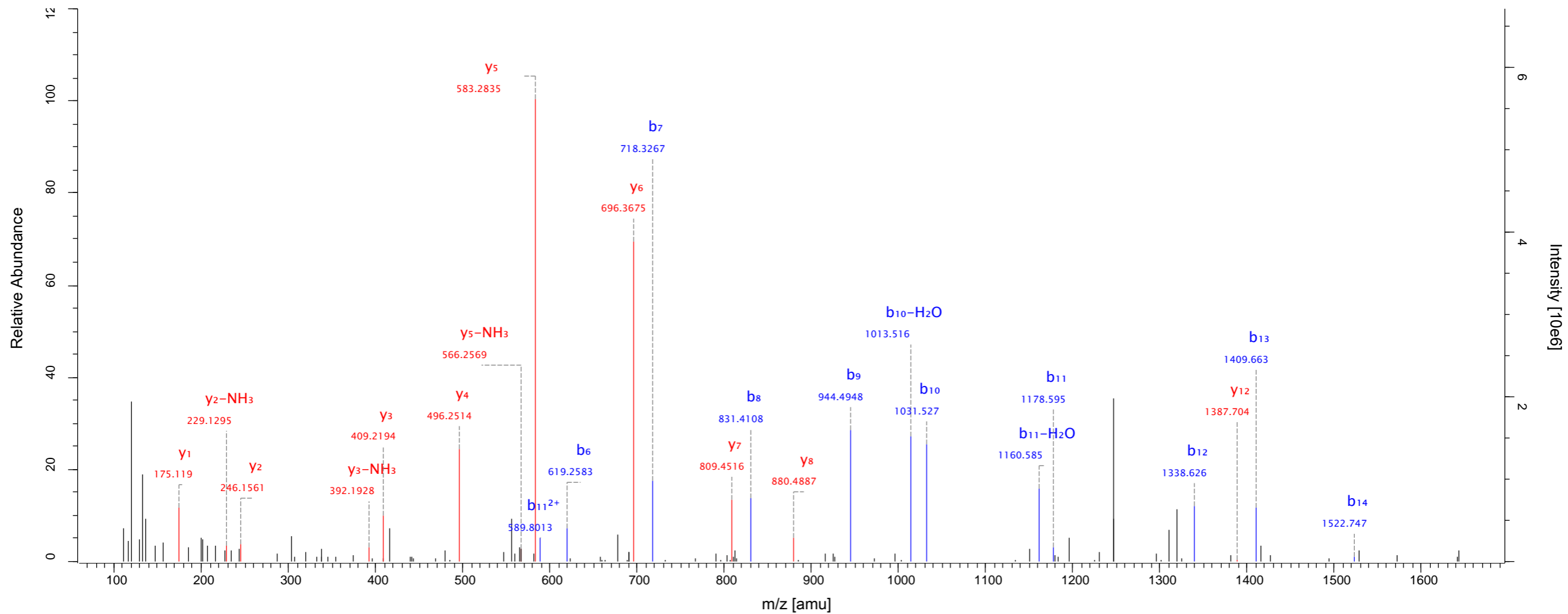
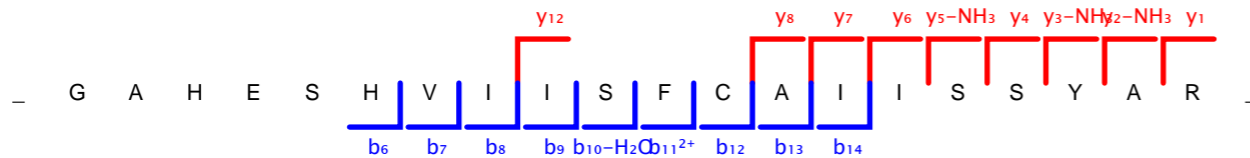
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 44452
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 134.37
 Method: FTMS; HCD



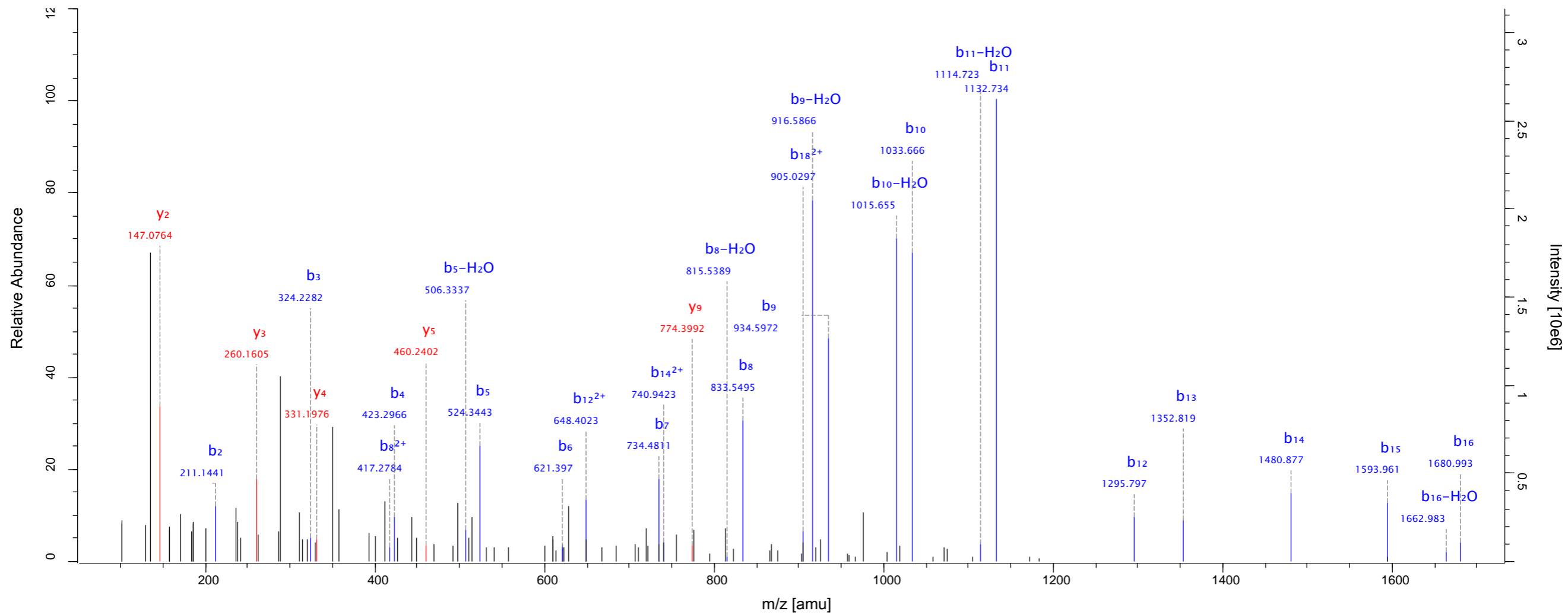
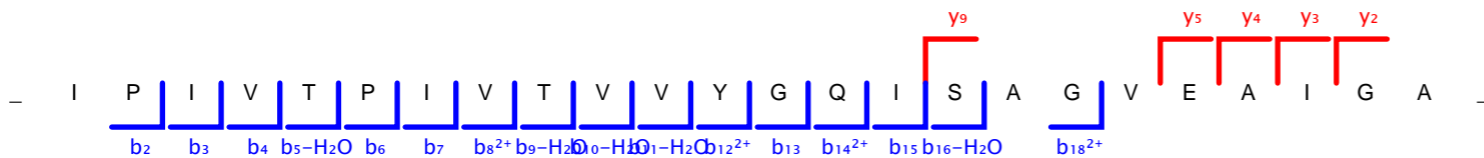
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 5266
 Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
 Peptide score: 143.86
 Method: FTMS; HCD



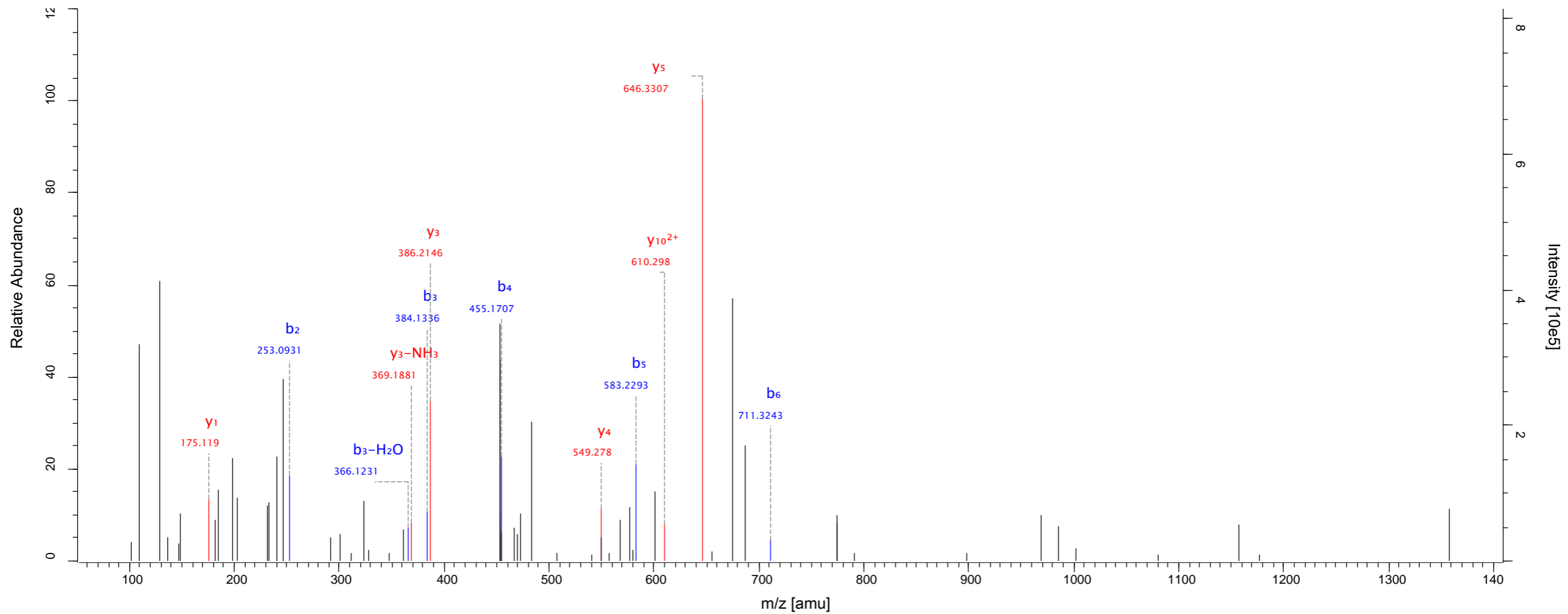
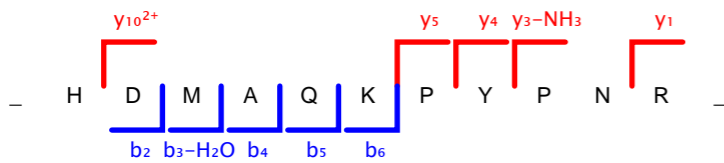
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 52923
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 122.31
Method: FTMS; HCD



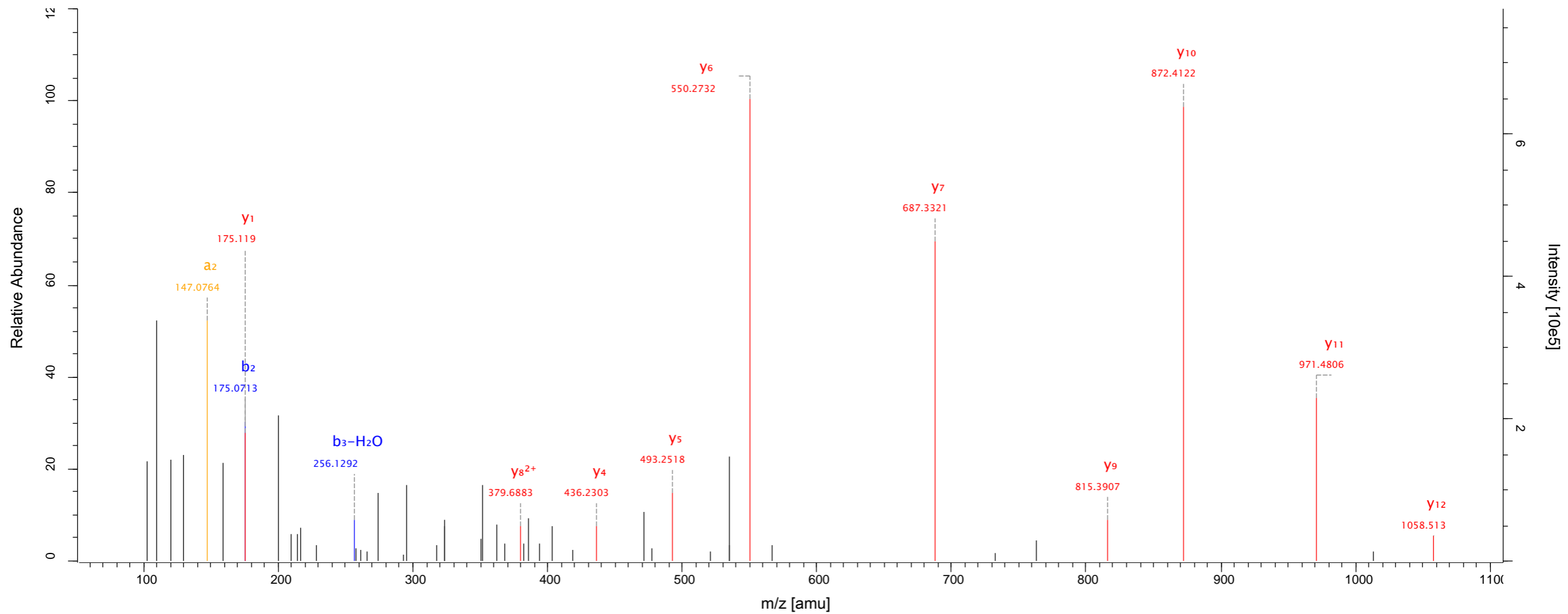
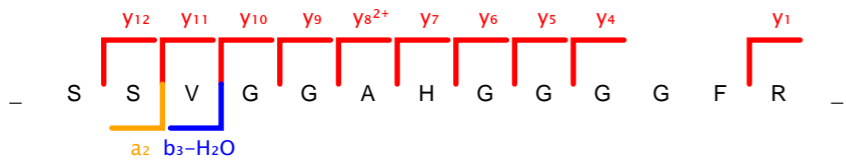
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 53553
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 141.86
 Method: FTMS; HCD



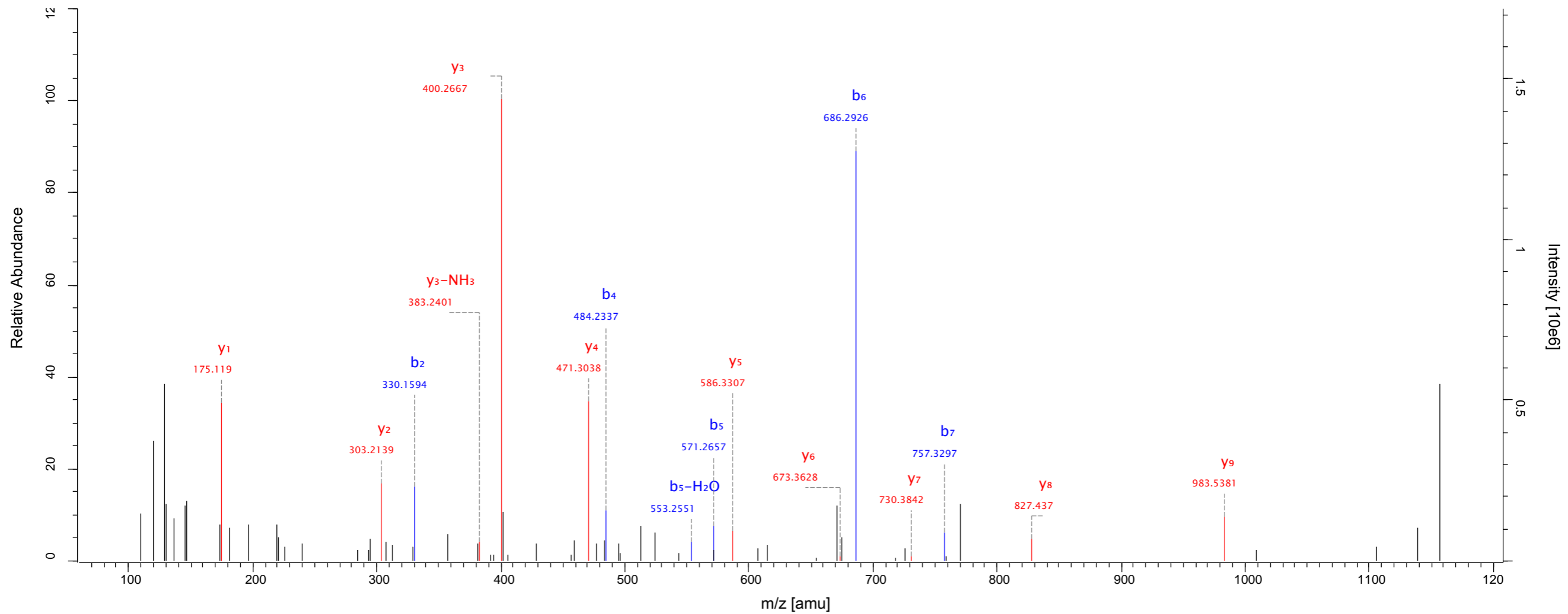
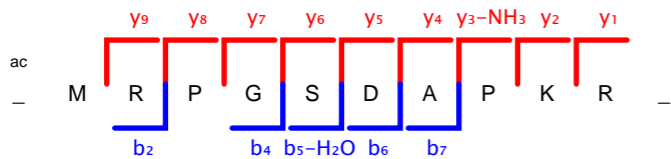
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 6368
Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>
Peptide score: 54.78
Method: FTMS; HCD



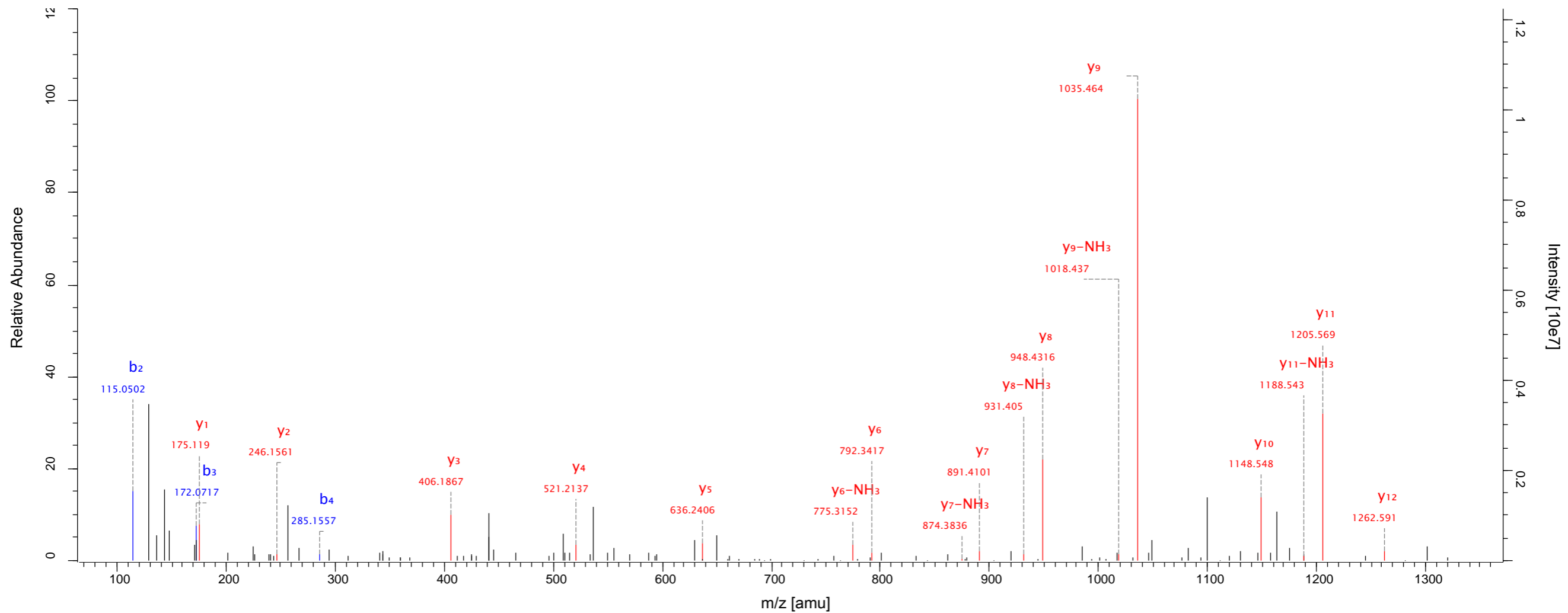
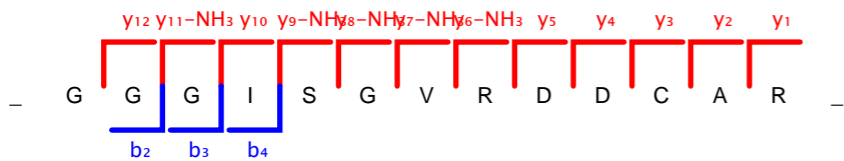
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 6663
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 65.18
Method: FTMS; HCD



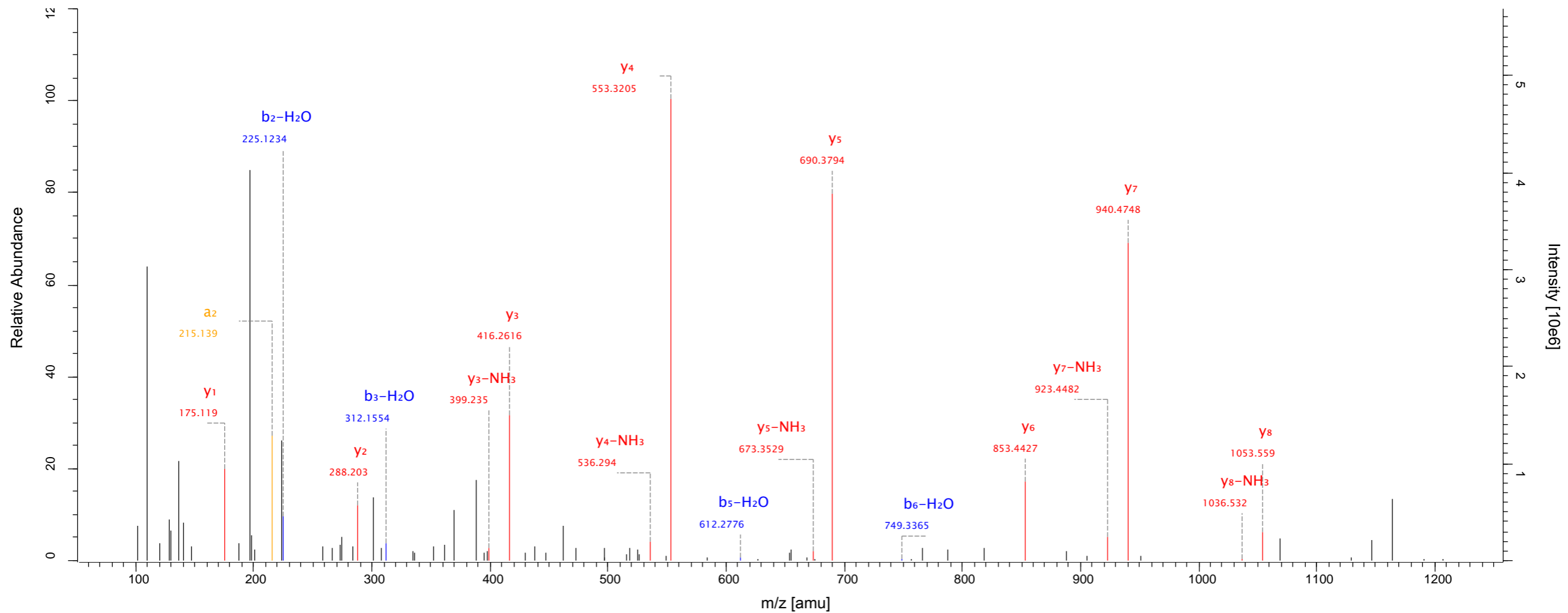
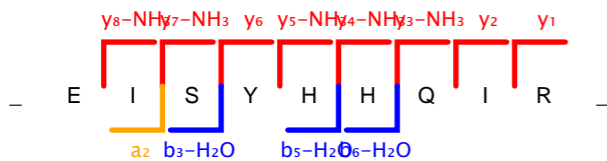
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 7284
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 104.52
Method: FTMS; HCD



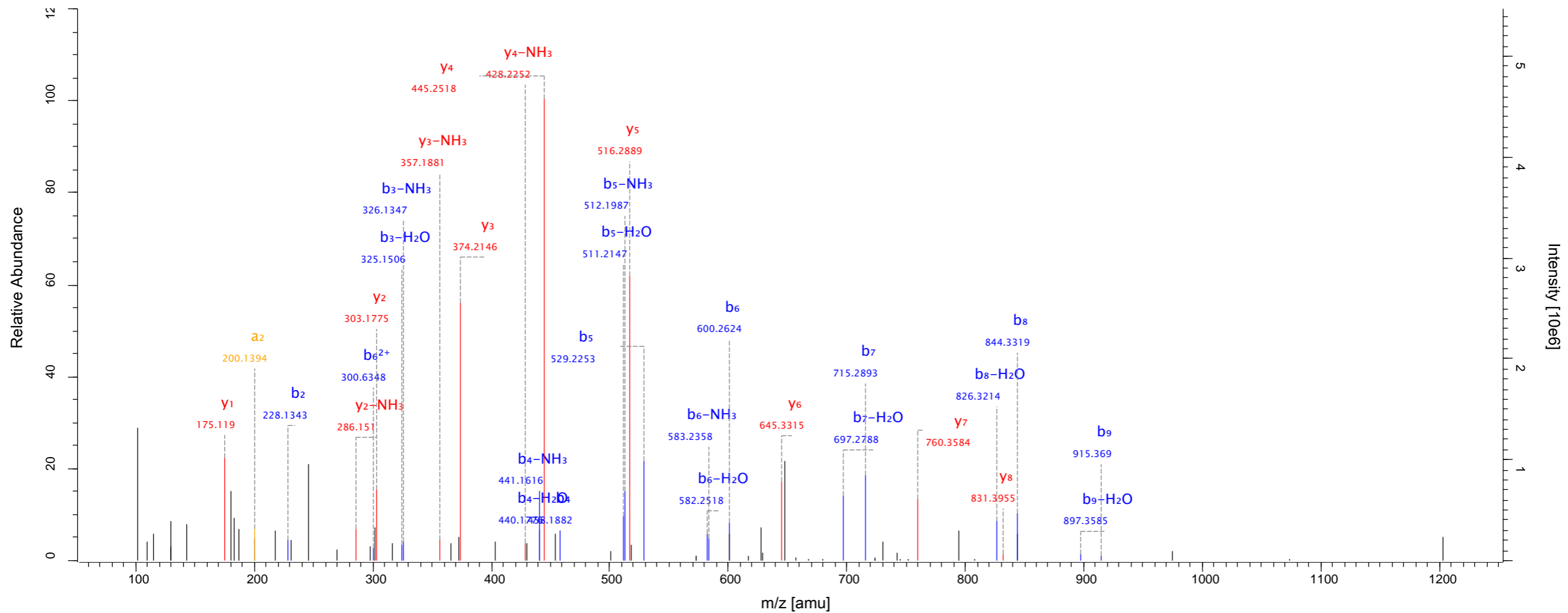
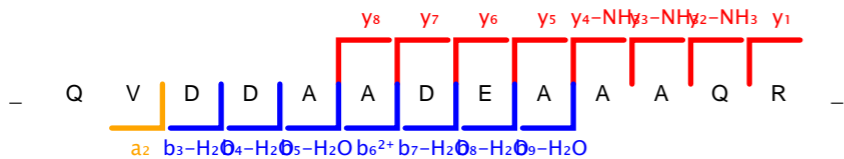
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 8900
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 102.4
Method: FTMS; HCD



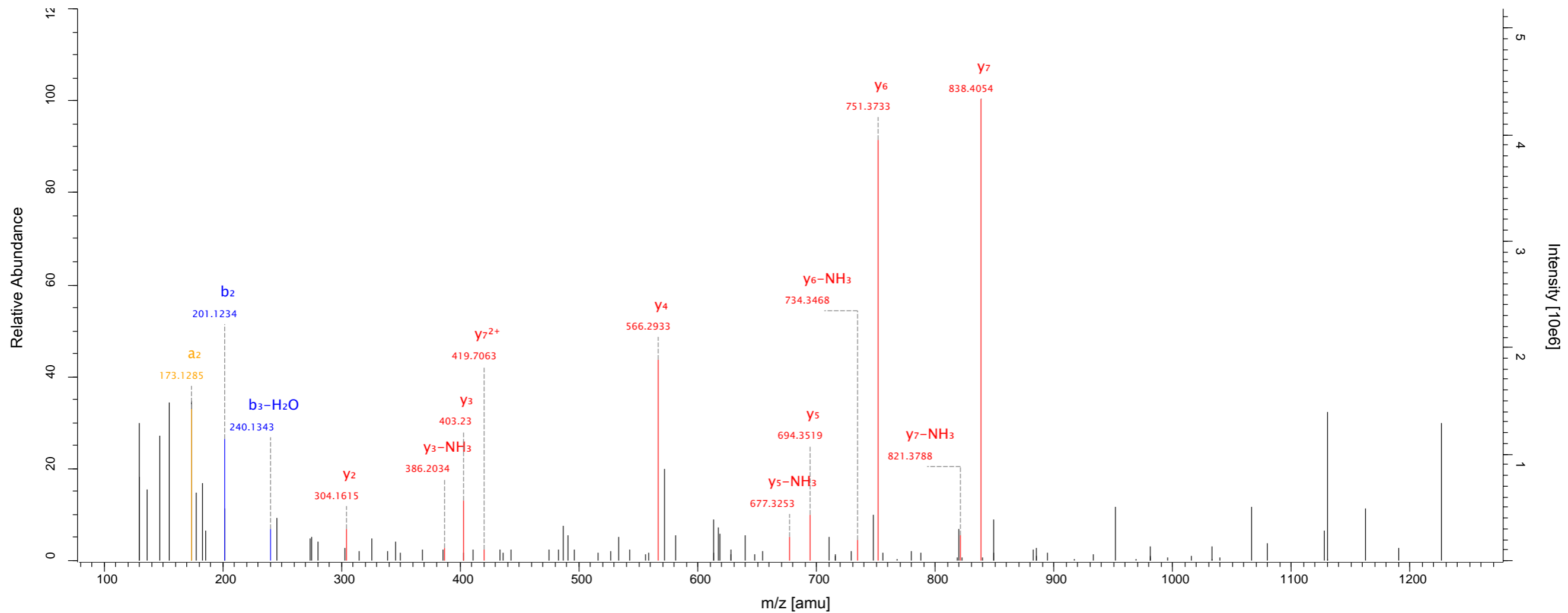
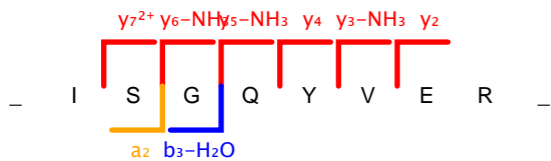
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 9058
Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
Peptide score: 105.57
Method: FTMS; HCD



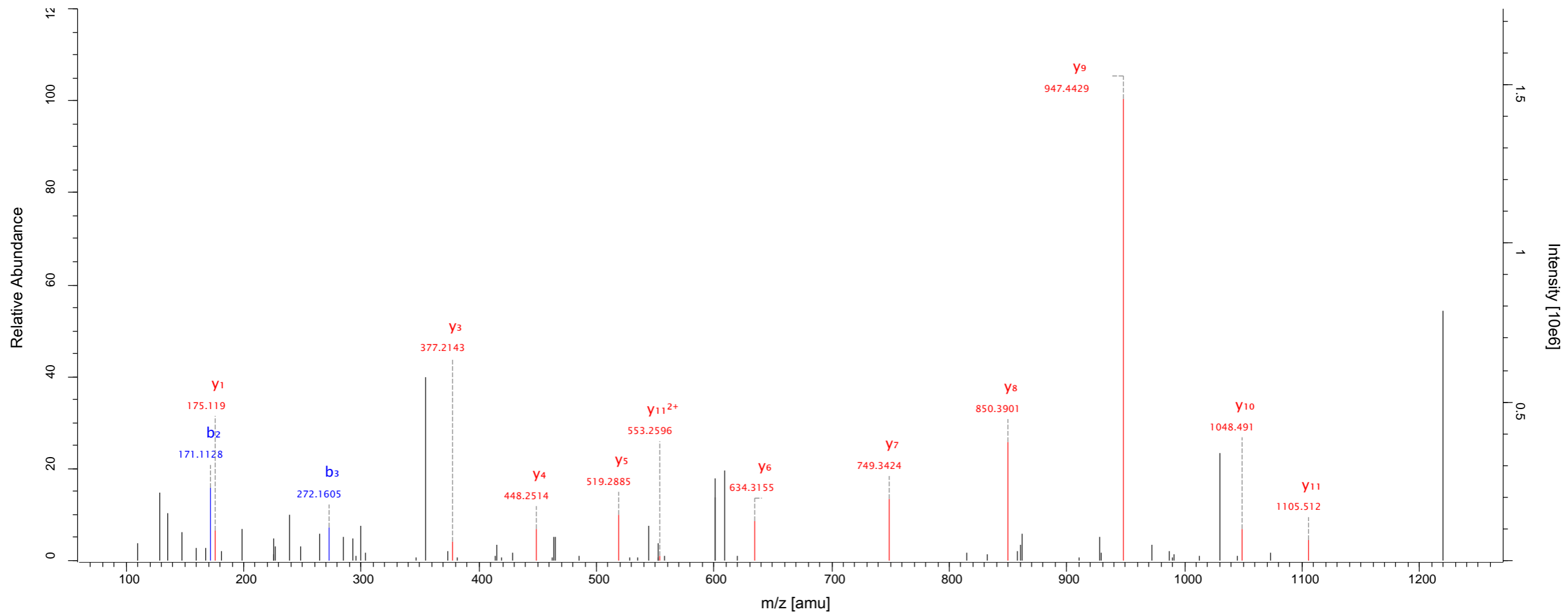
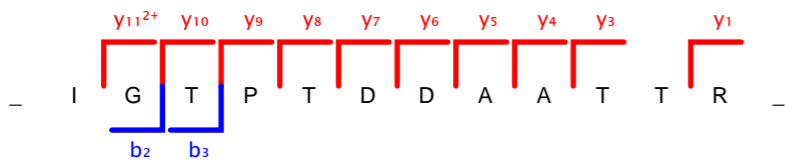
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
 Scan number: 9344
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 179.35
 Method: FTMS; HCD



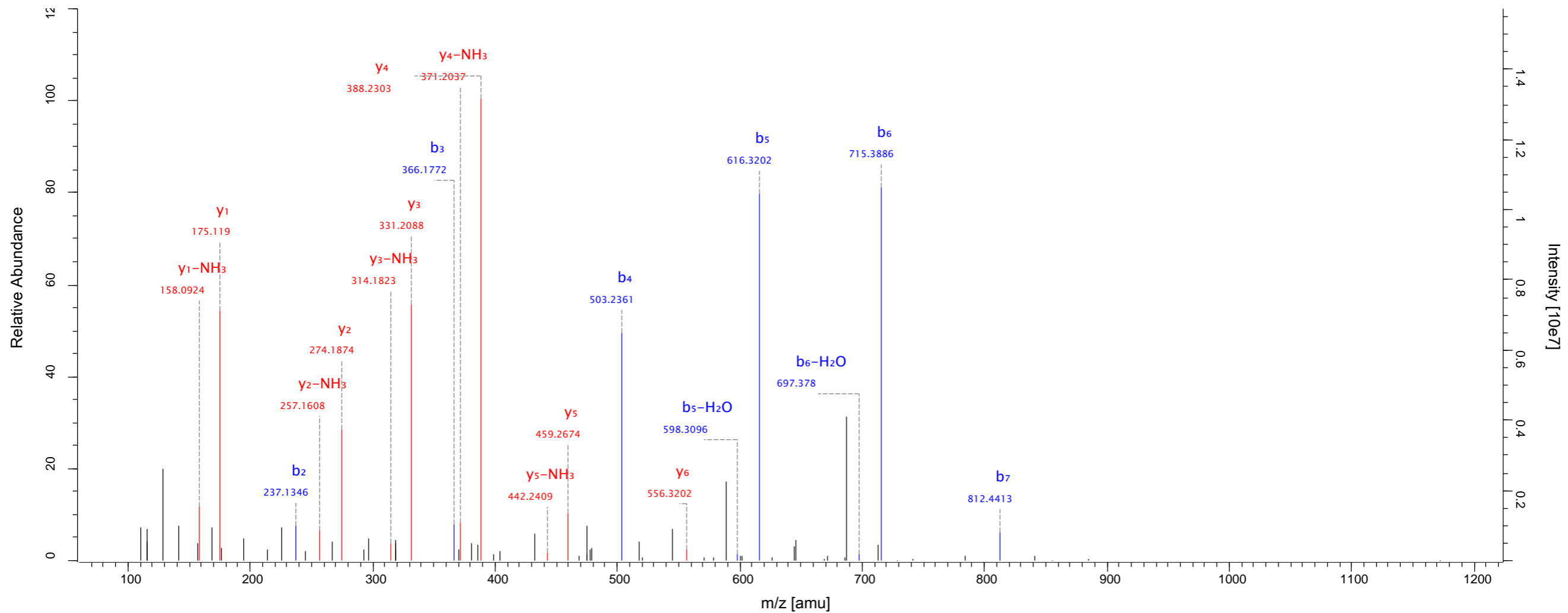
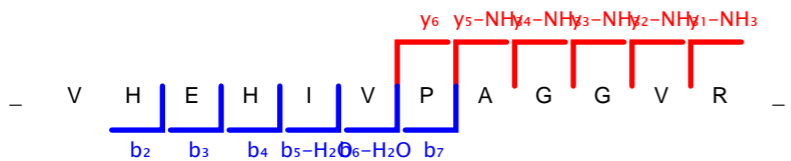
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_11
Scan number: 9835
Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
Peptide score: 86.79
Method: FTMS; HCD



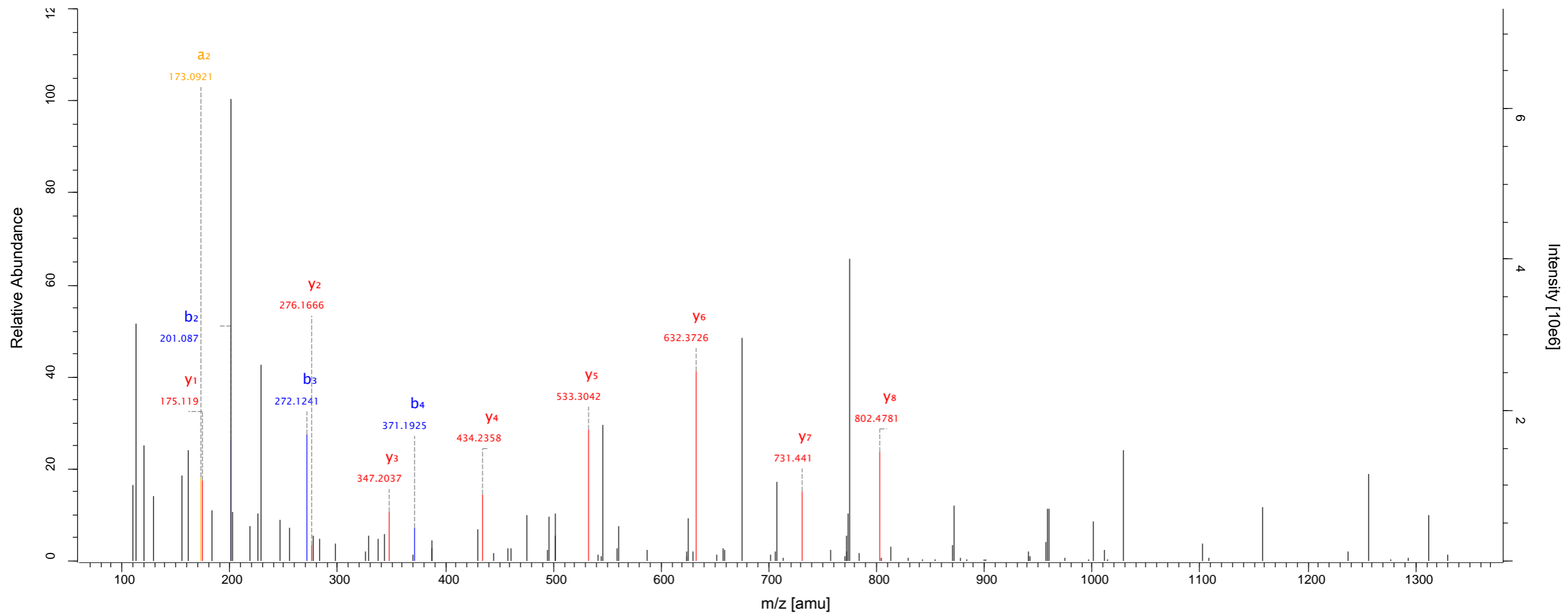
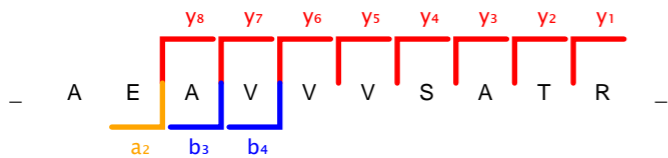
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_12
Scan number: 10528
Protein: ORFL114W + NC_006273.2[60151,60708]+ <unknown description>
Peptide score: 66.26
Method: FTMS; HCD



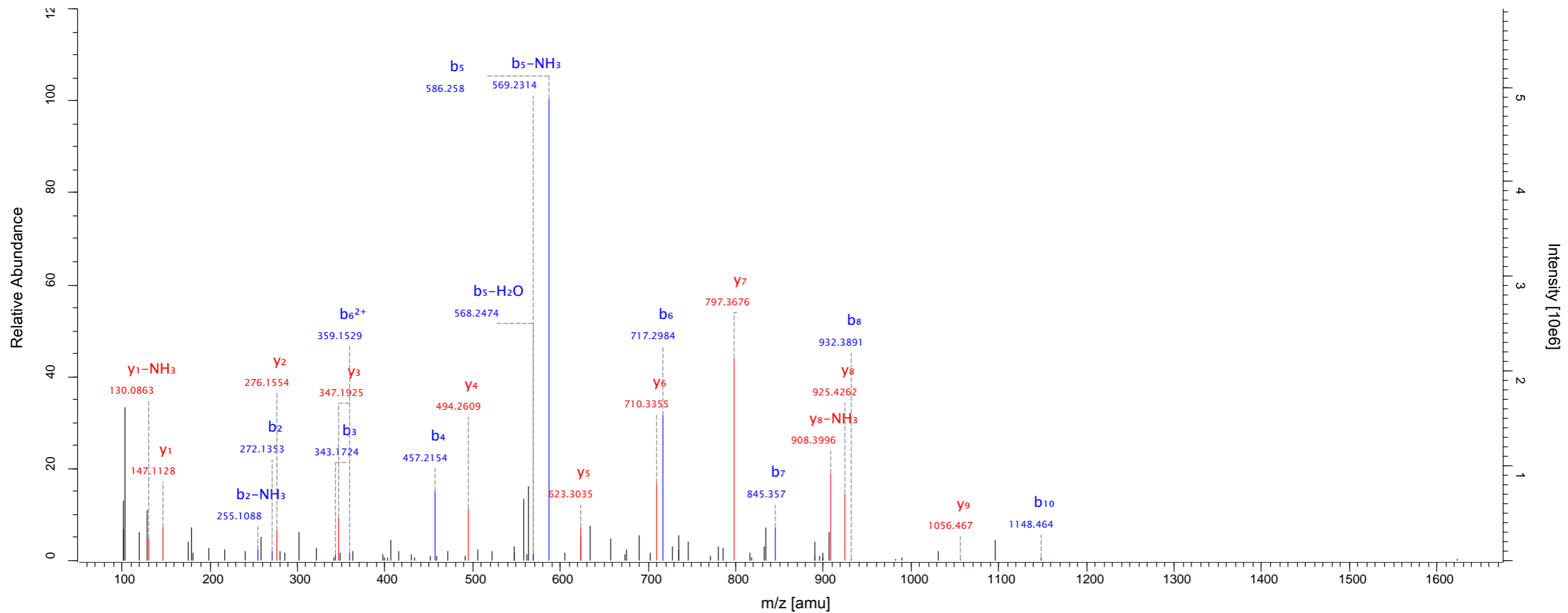
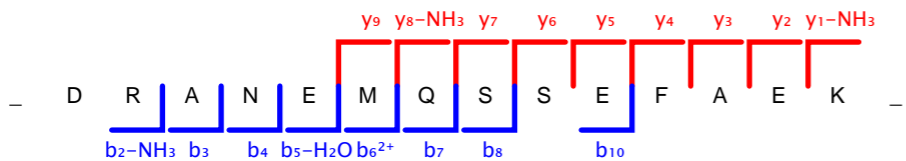
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
Scan number: 11217
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 110.39
Method: FTMS; HCD



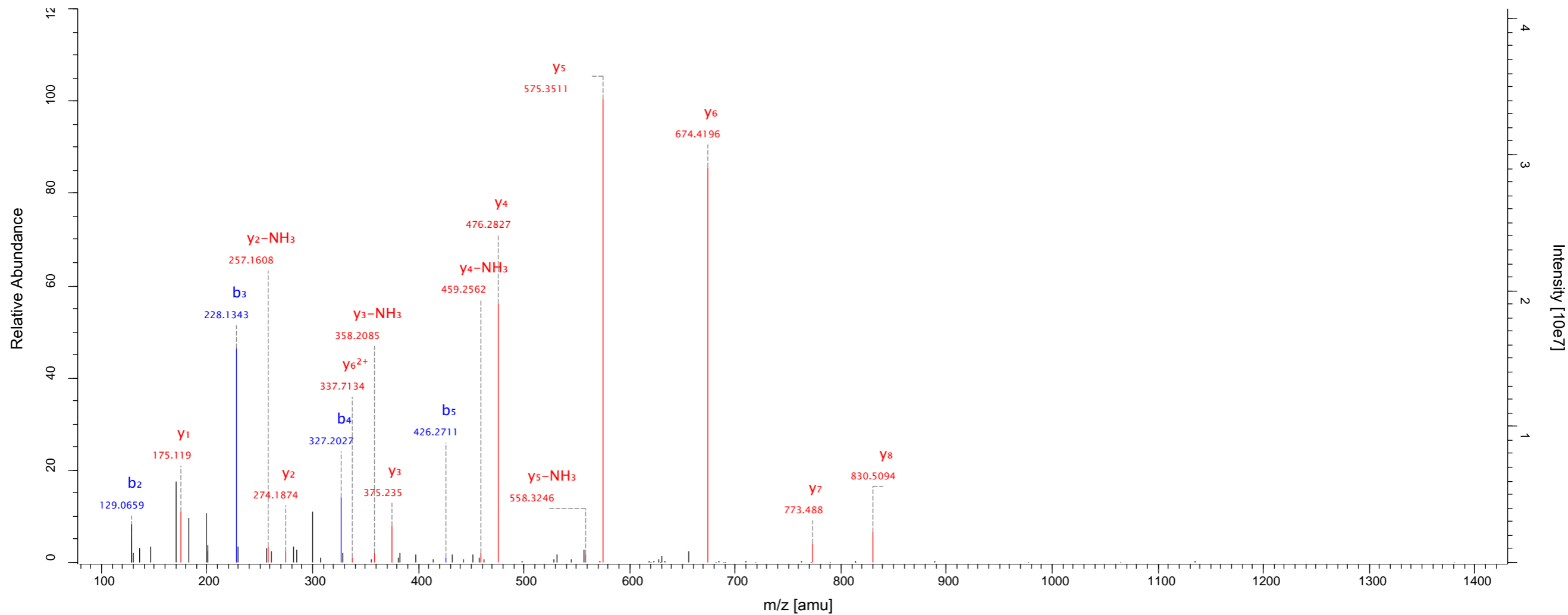
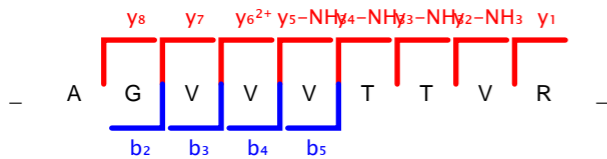
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
Scan number: 12247
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 59.43
Method: FTMS; HCD



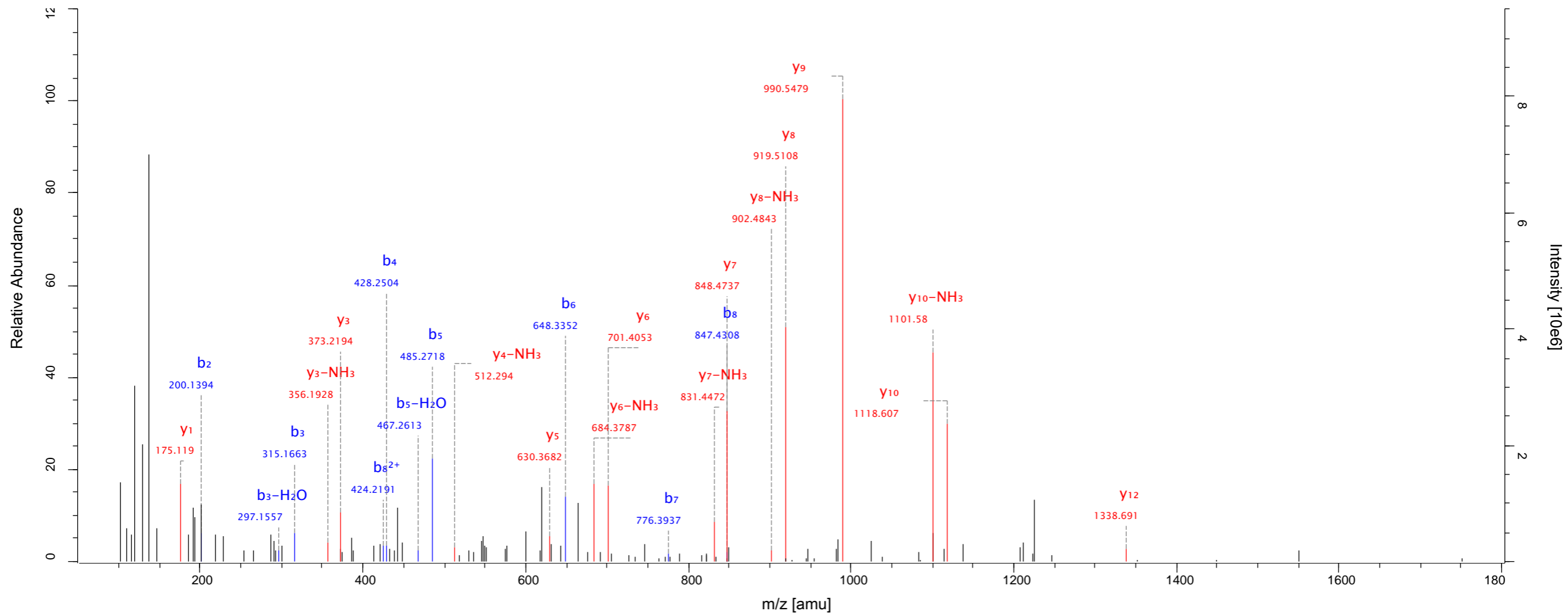
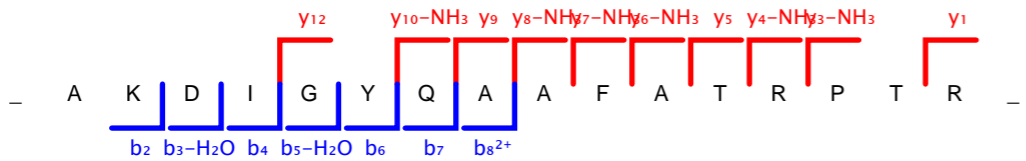
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
Scan number: 12830
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 80.53
Method: FTMS; HCD



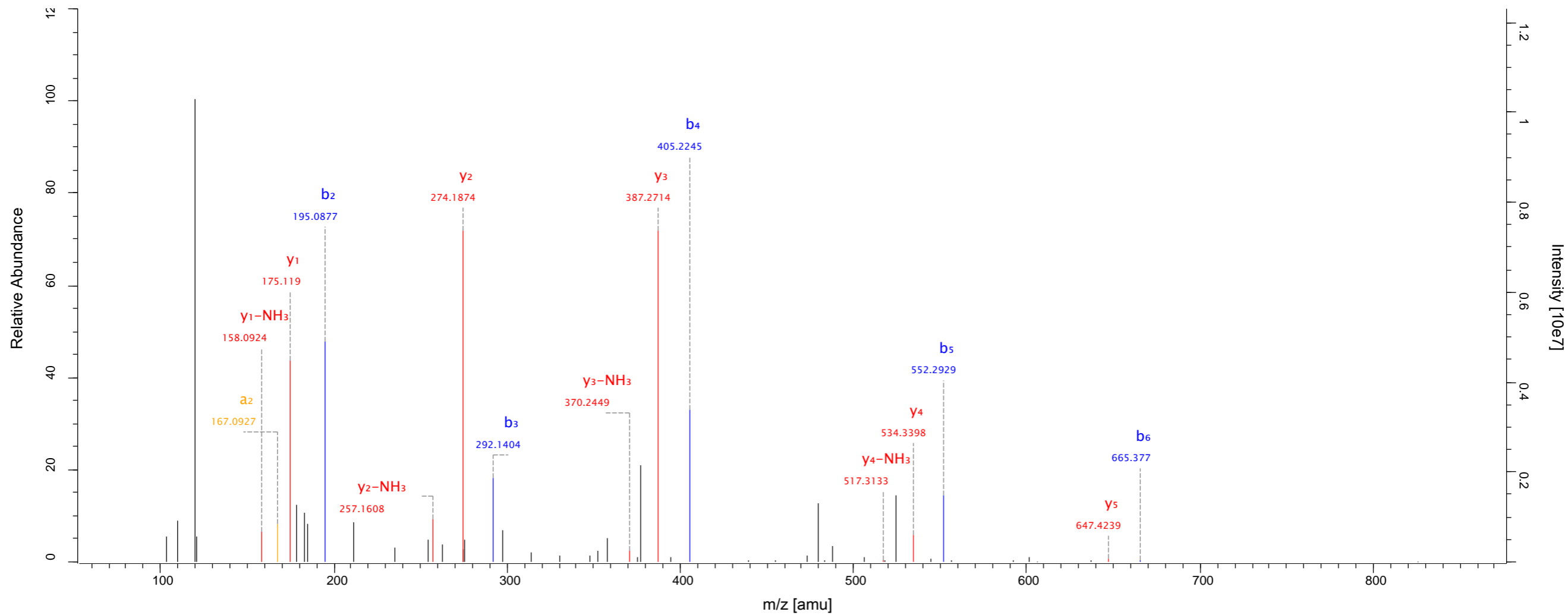
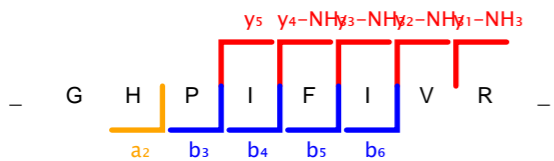
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
Scan number: 14417
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 116.73
Method: FTMS; HCD



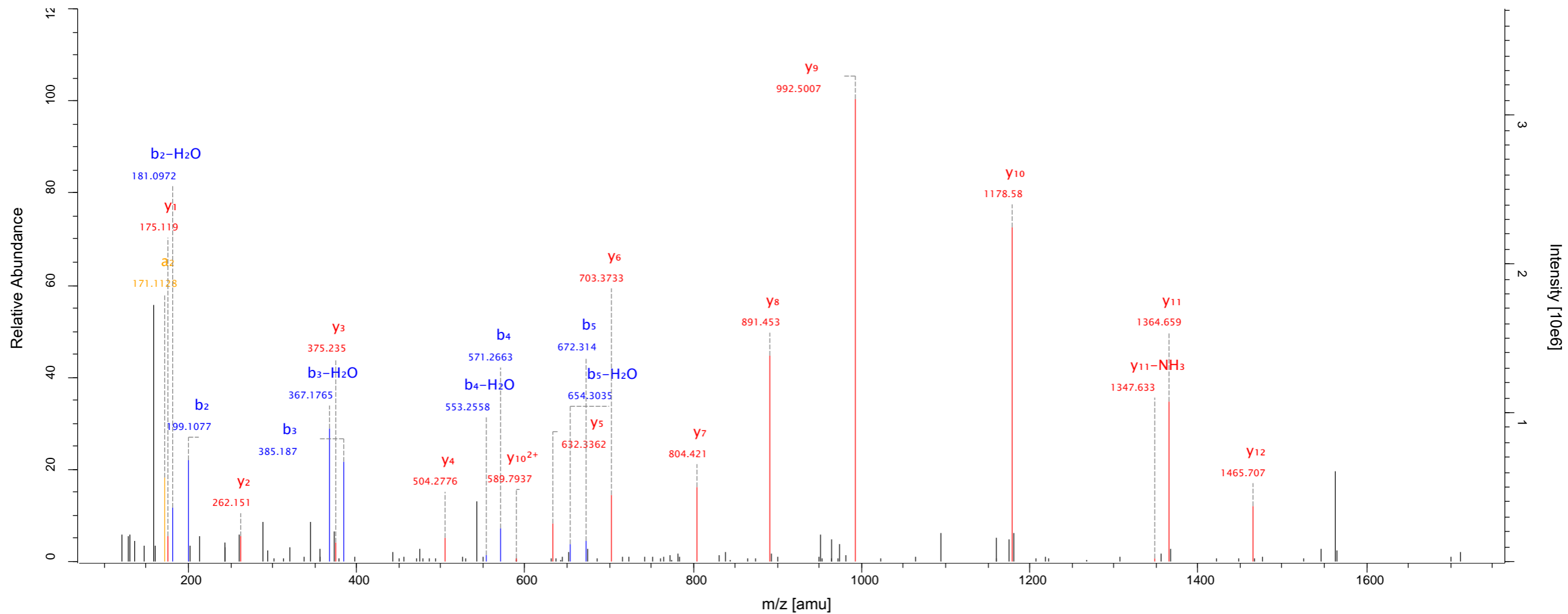
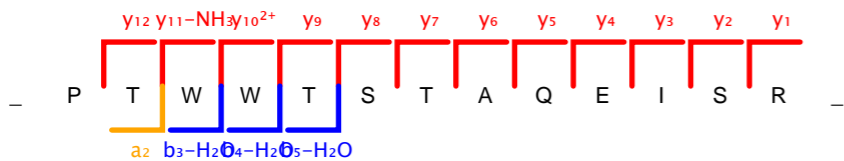
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
 Scan number: 19337
 Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
 Peptide score: 105.14
 Method: FTMS; HCD



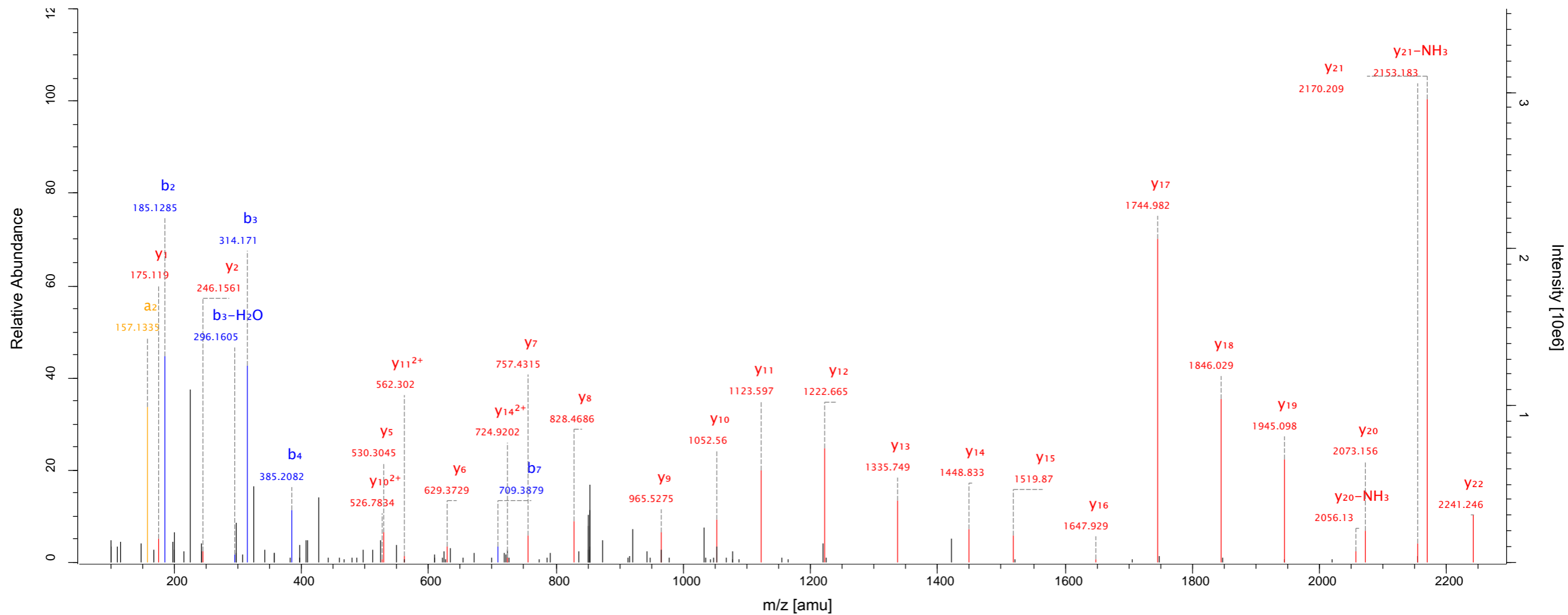
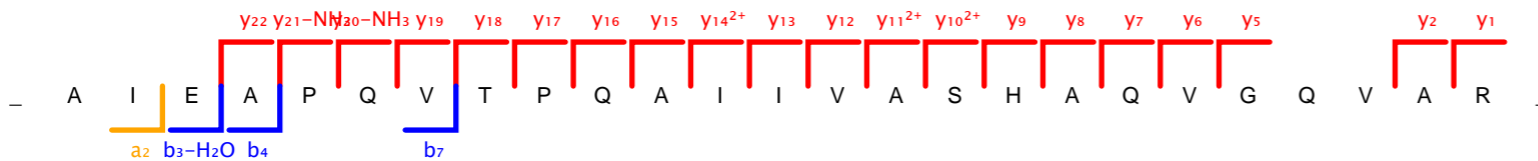
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_12
Scan number: 24065
Protein: ORFL114W + NC_006273.2[60151,60708]+ <unknown description>
Peptide score: 107.66
Method: FTMS; HCD



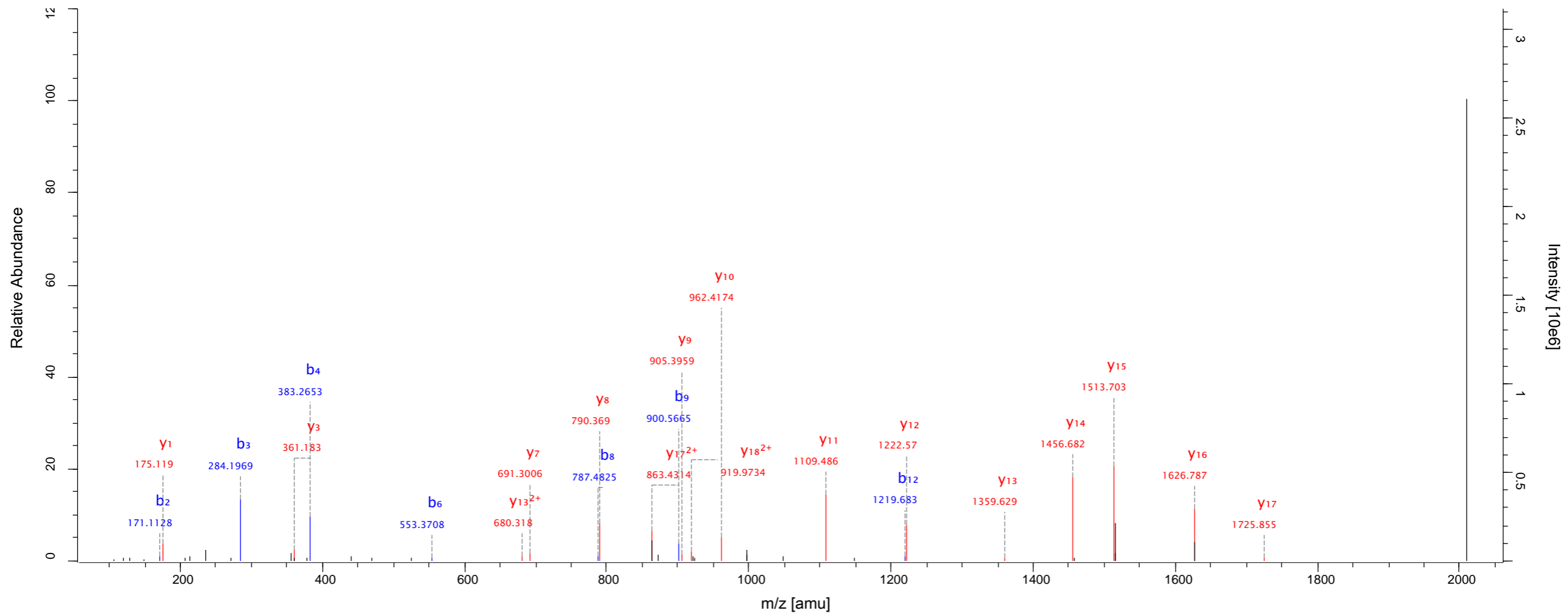
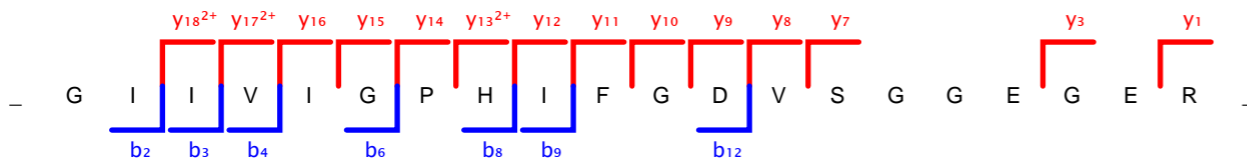
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_12
 Scan number: 42505
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 133.05
 Method: FTMS; HCD



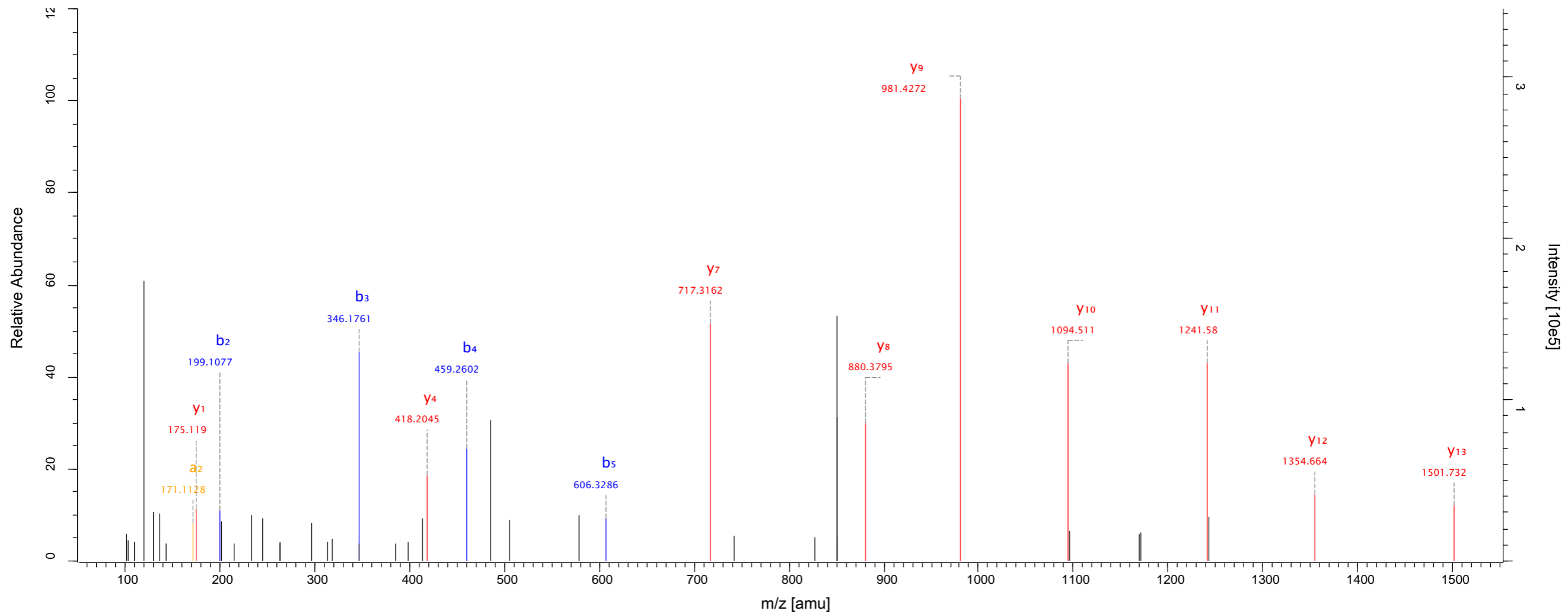
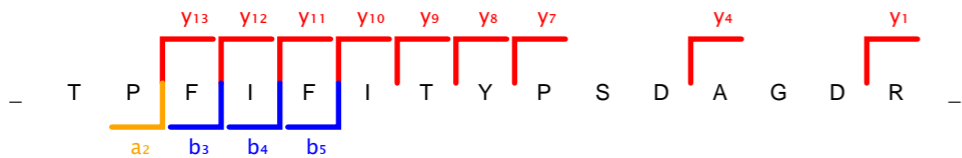
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
 Scan number: 46474
 Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
 Peptide score: 174.74
 Method: FTMS; HCD



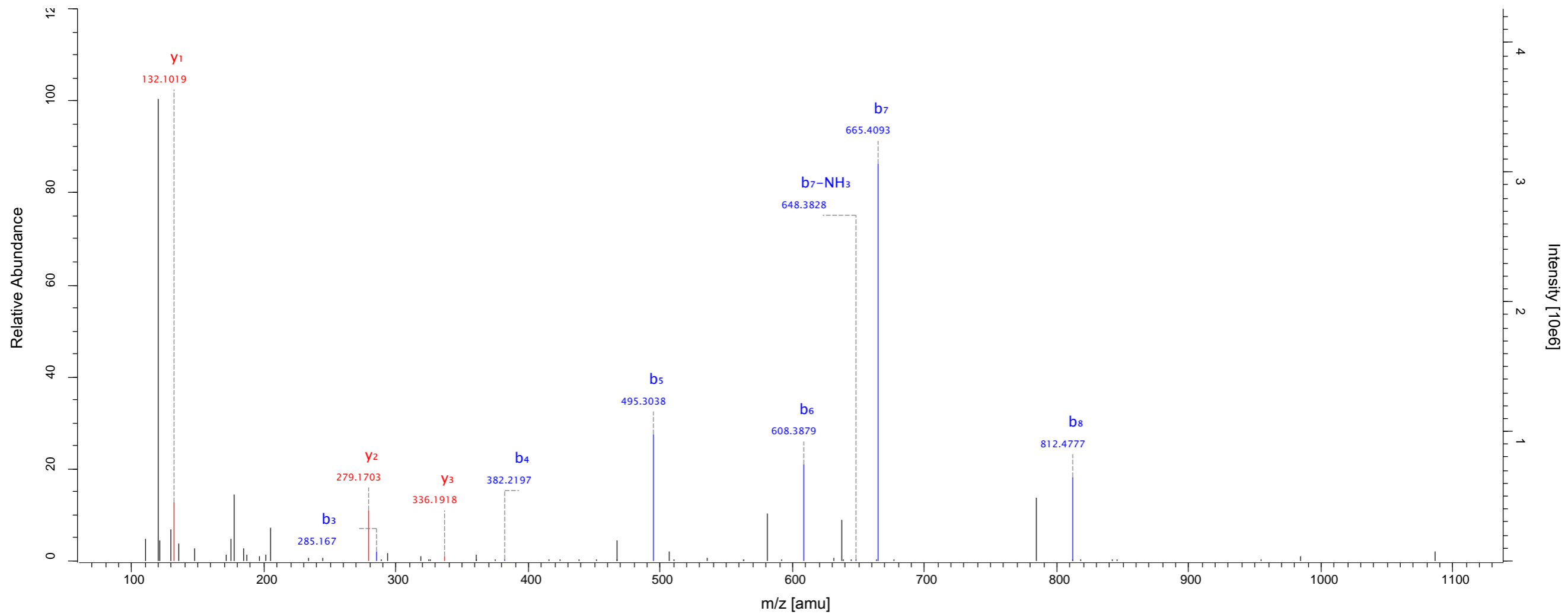
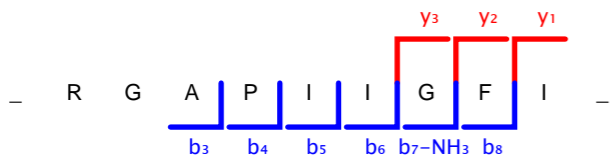
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
Scan number: 50028
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 159.31
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_12
Scan number: 52281
Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
Peptide score: 87.18
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_12
Scan number: 53234
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 52.48
Method: FTMS; HCD



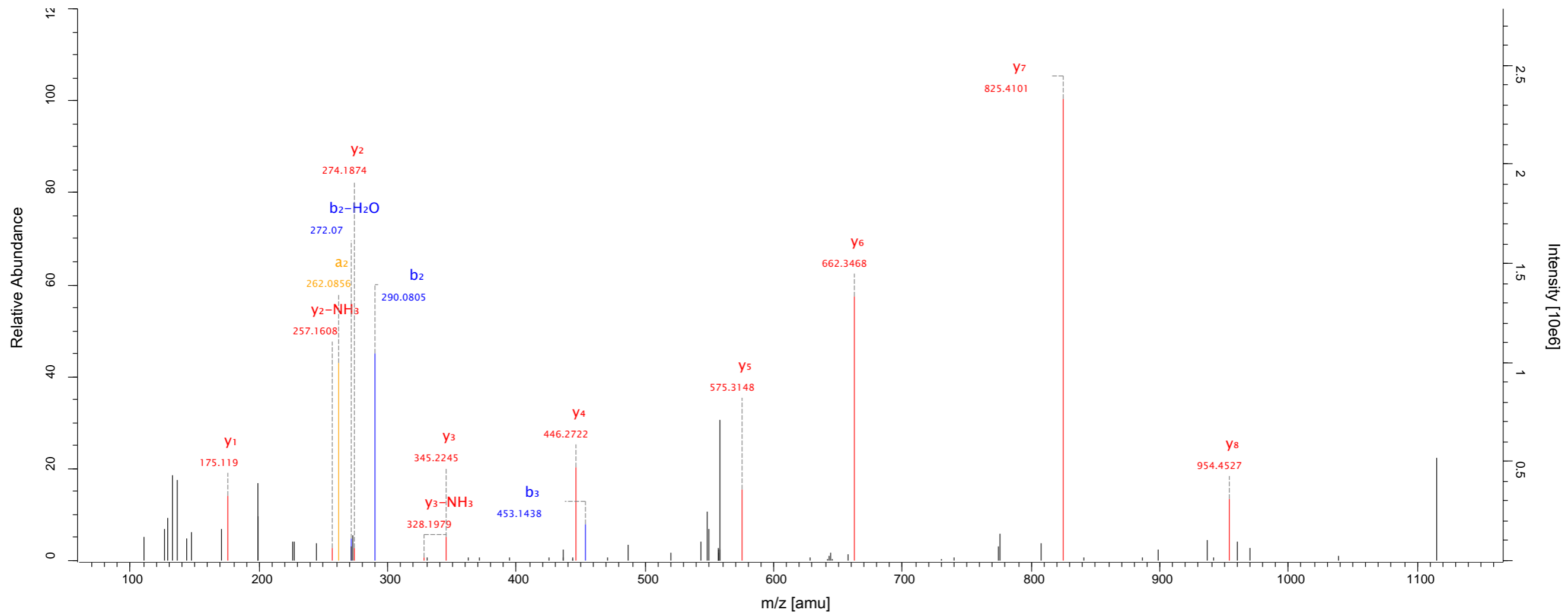
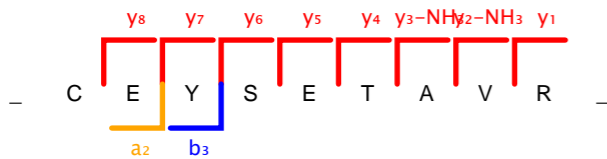
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_13

Scan number: 10425

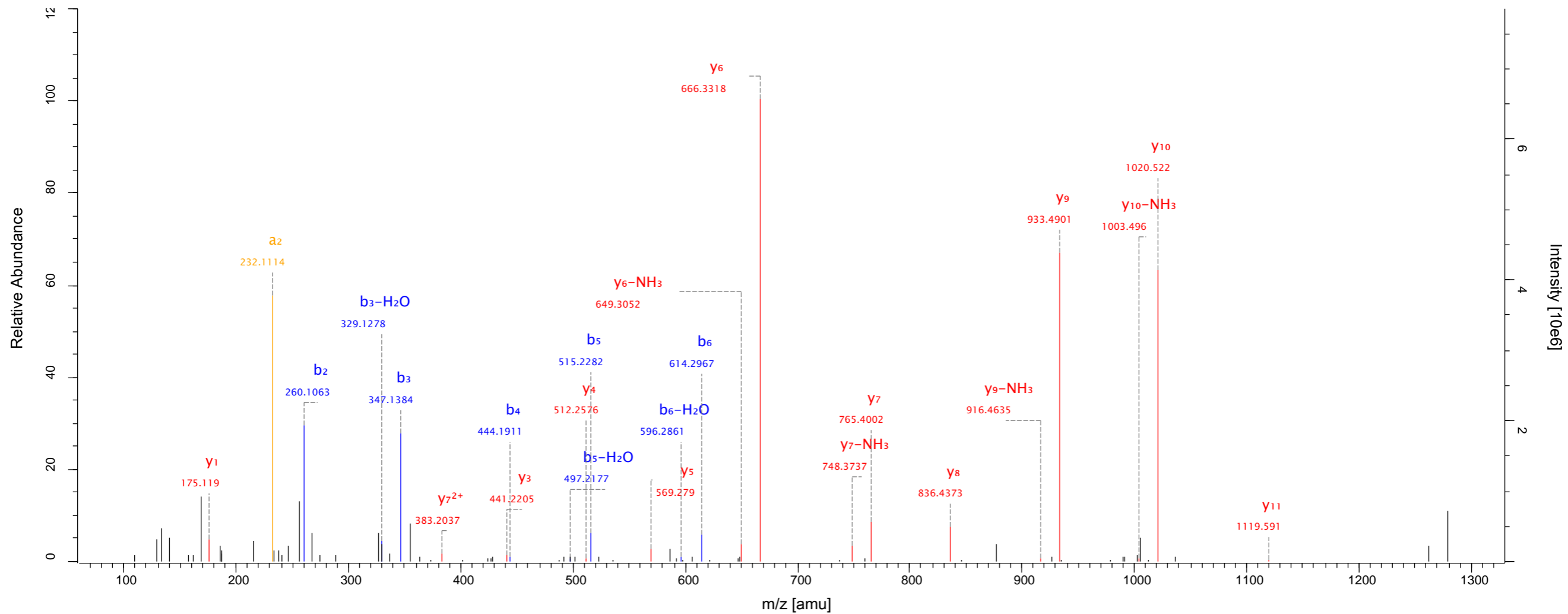
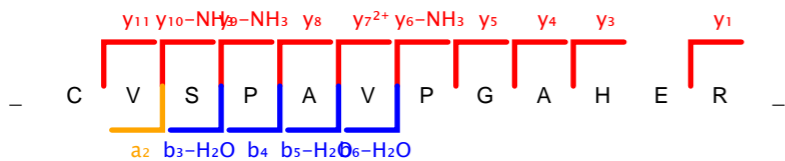
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 80.24

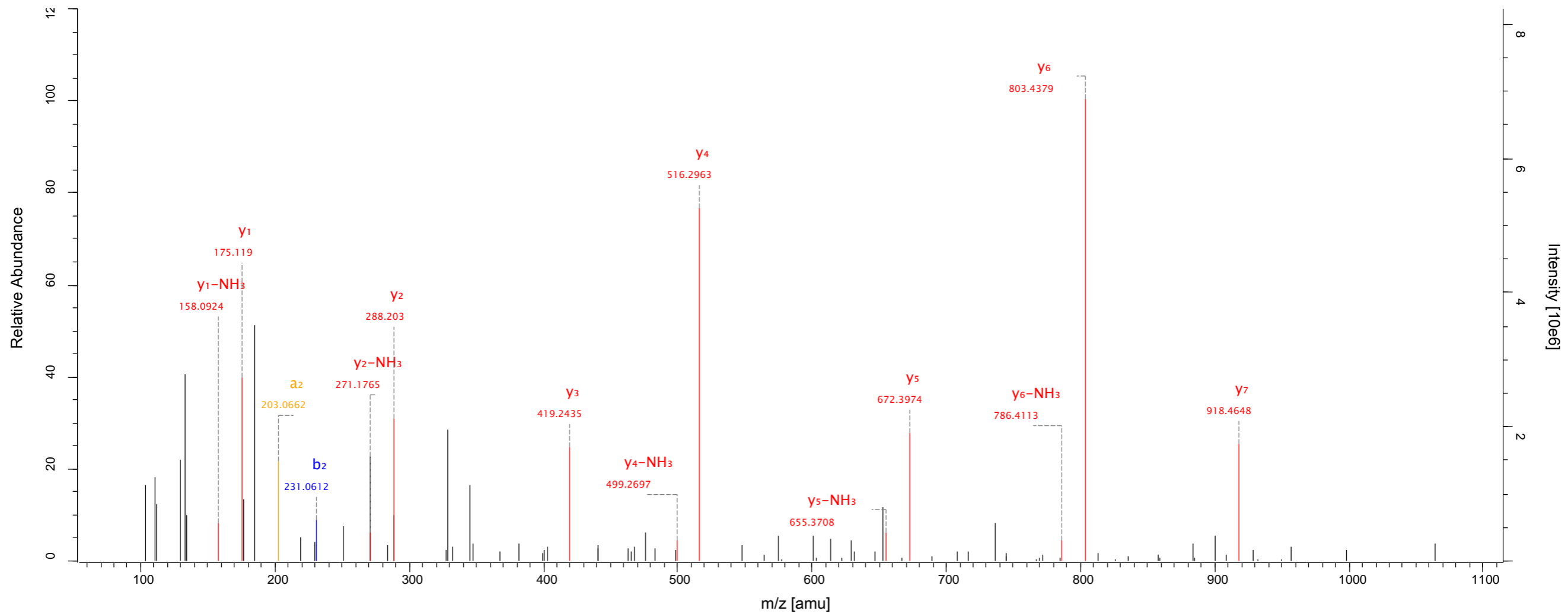
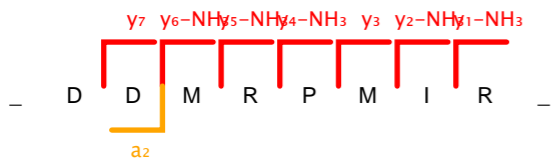
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_13
 Scan number: 11022
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 164.64
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13
Scan number: 15298
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 100.45
Method: FTMS; HCD



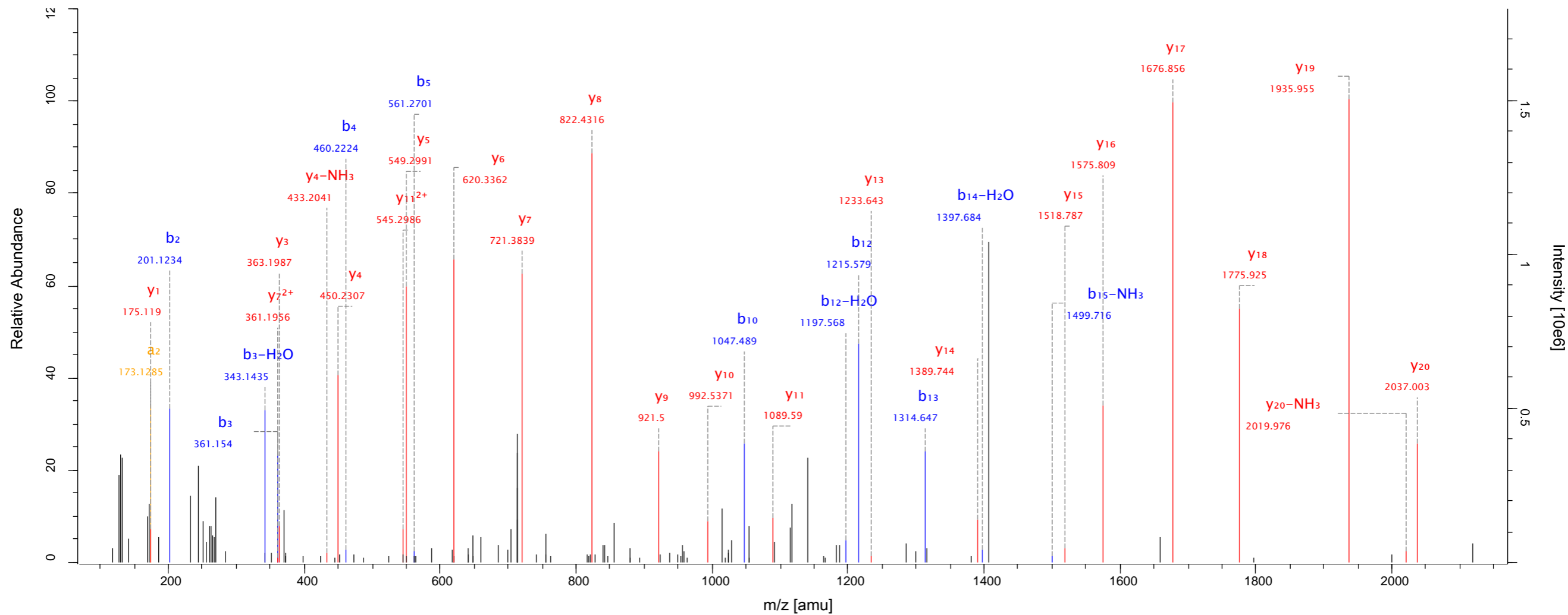
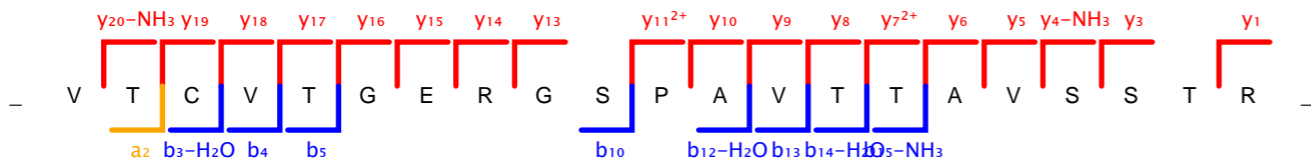
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13

Scan number: 18294

Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 199.56

Method: FTMS; HCD



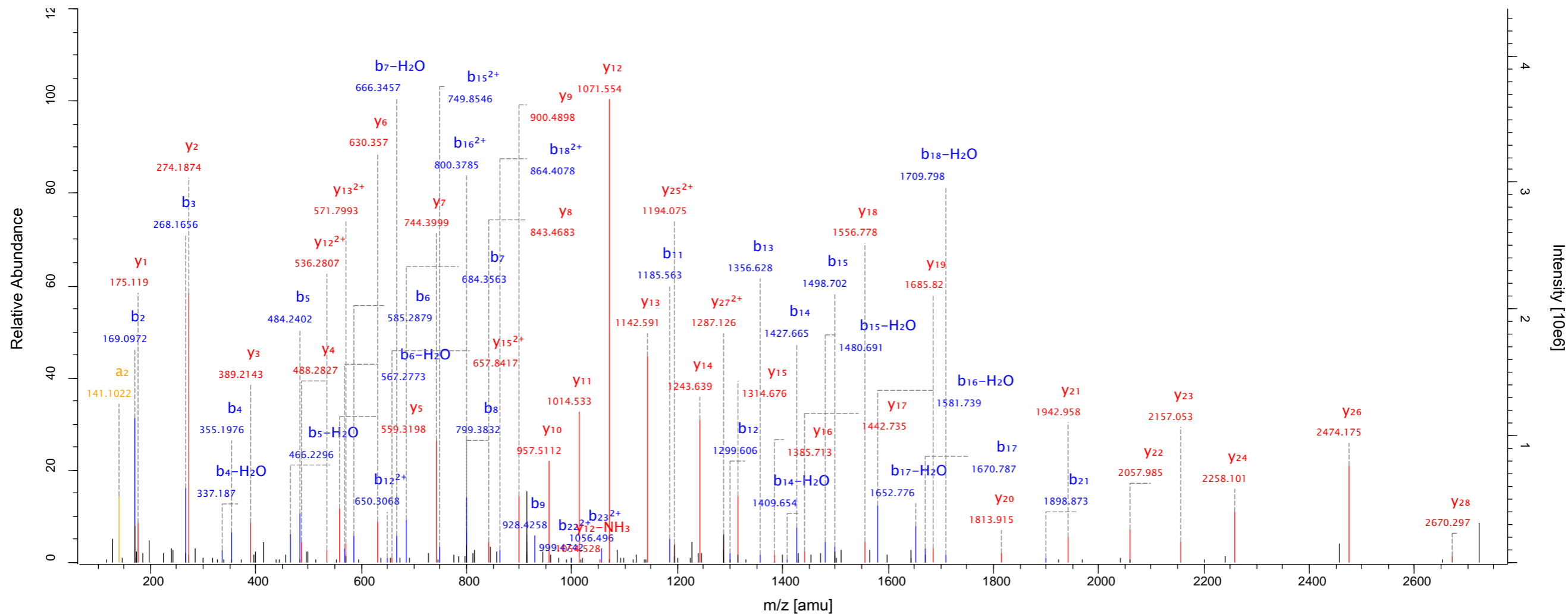
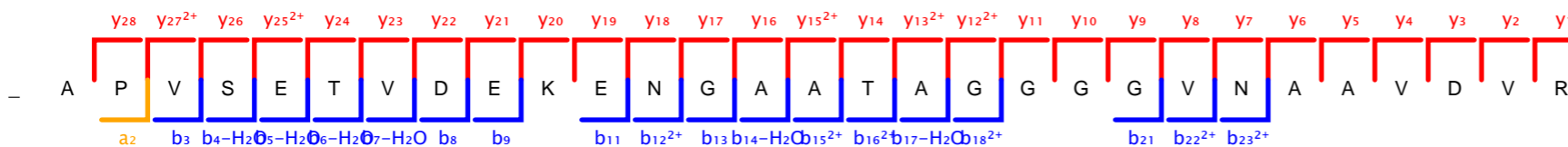
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13

Scan number: 23271

Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 346.34

Method: FTMS; HCD



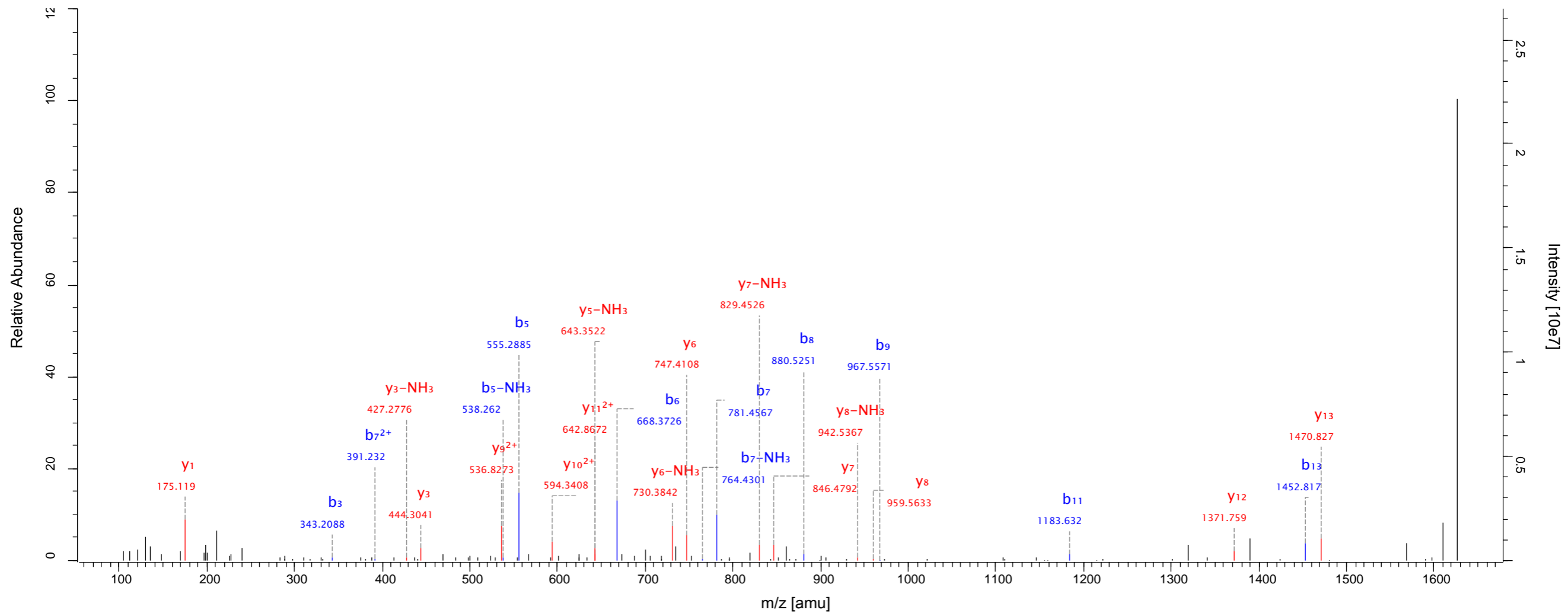
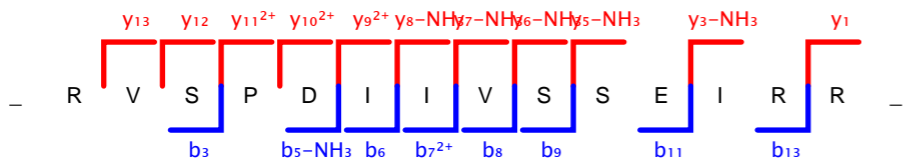
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13

Scan number: 28961

Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 134.21

Method: FTMS; HCD



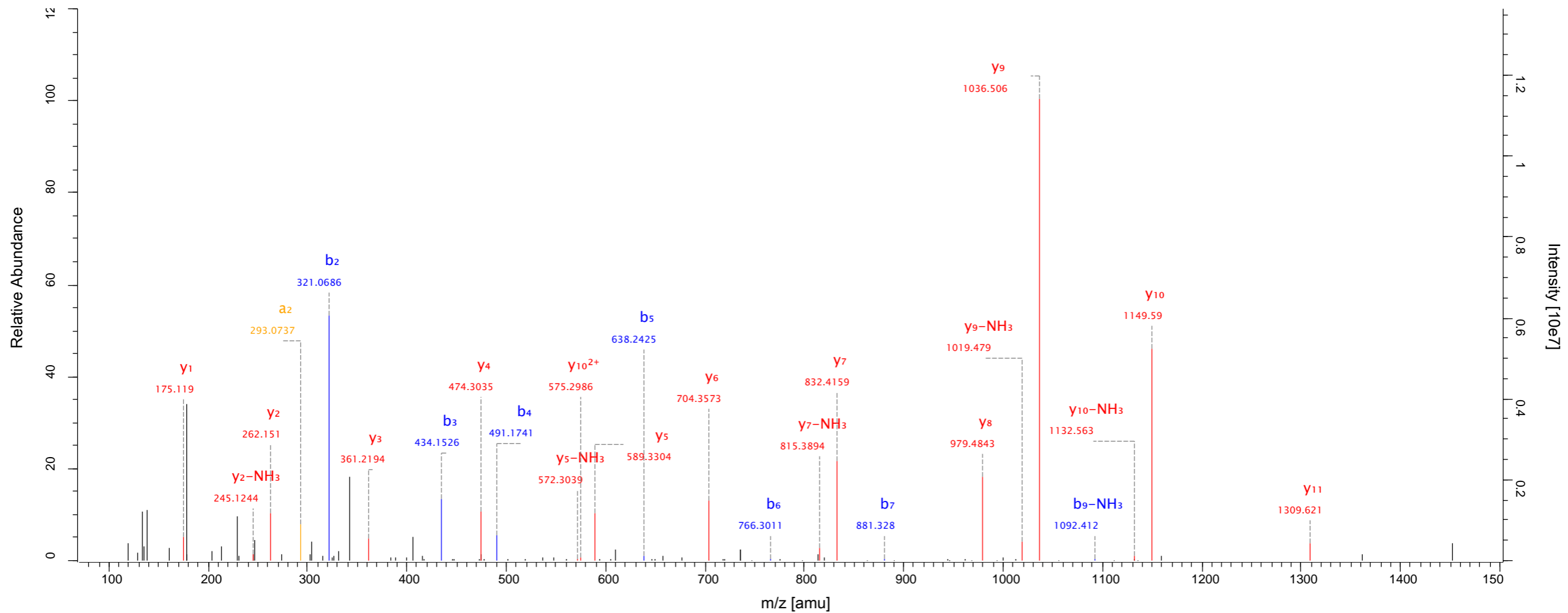
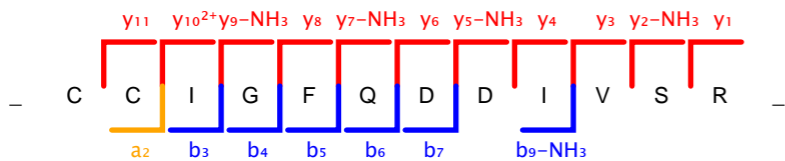
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13

Scan number: 33225

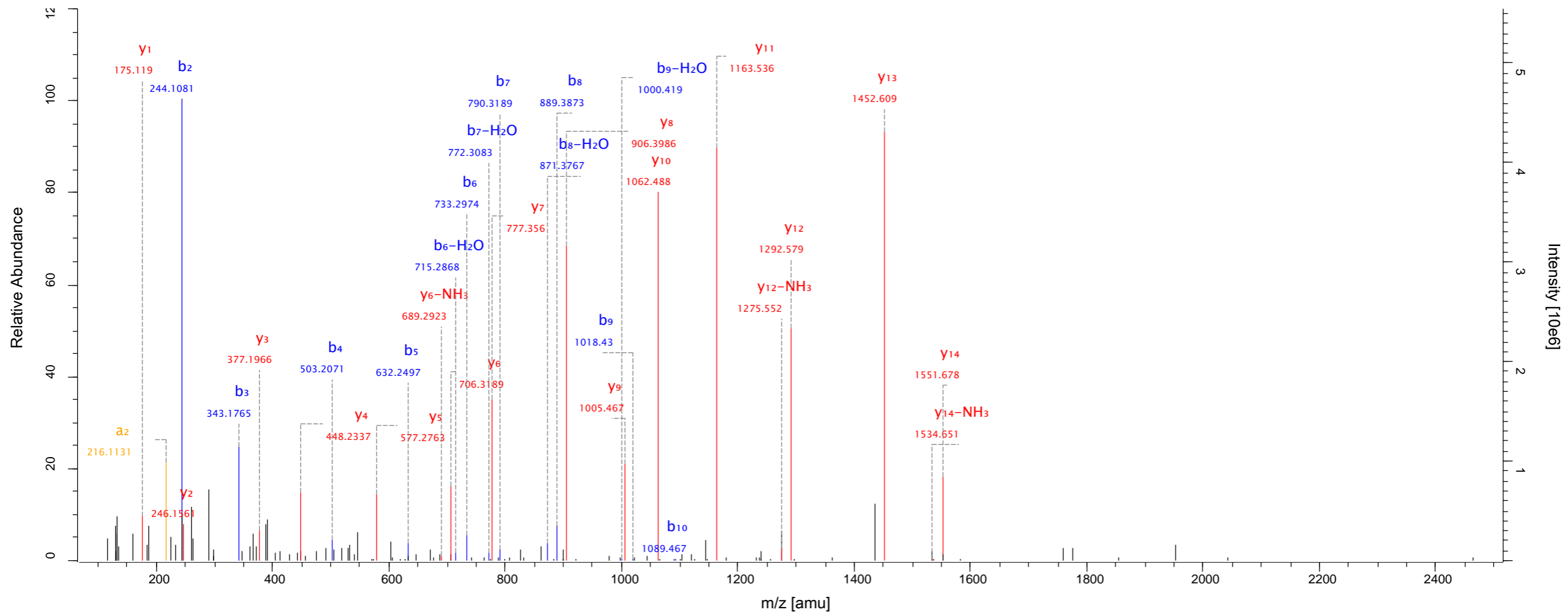
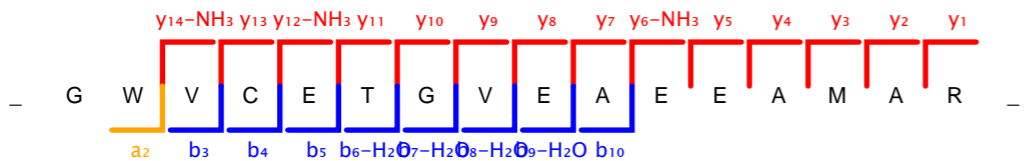
Protein: ORFL159W + NC_006273.2[101475,109052]+ <unknown description>; ORFL159W.iORF1 + NC_006273.2[101526,109052]+ <unknown description>; ORFL178W + NC_006273.2[108142,109052]+ <unknown description>; ORFL163W + NC_006273.2[101475,109052]+ <unknown description>

Peptide score: 164.64

Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_13
 Scan number: 38875
 Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>
 Peptide score: 169.04
 Method: FTMS; HCD



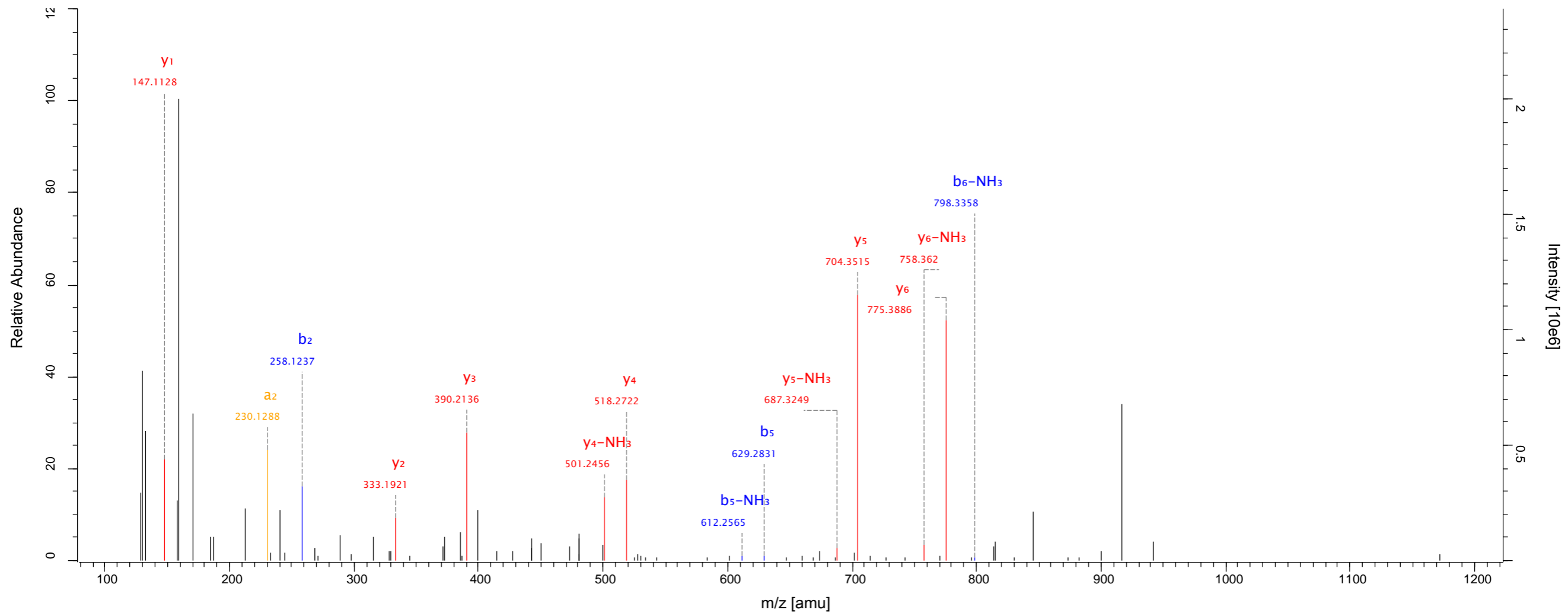
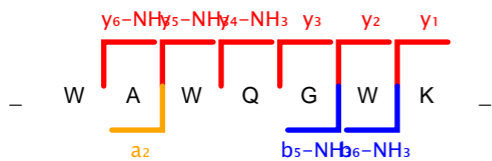
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_13

Scan number: 50936

Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 132.22

Method: FTMS; HCD



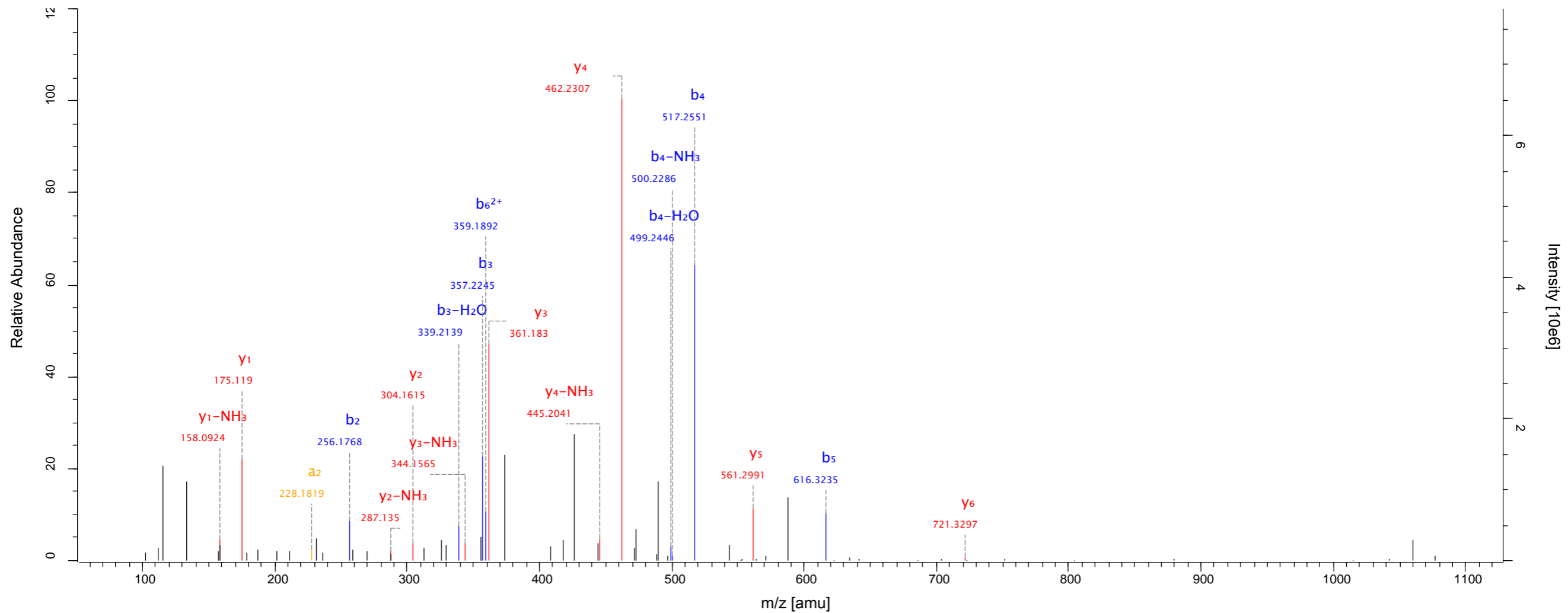
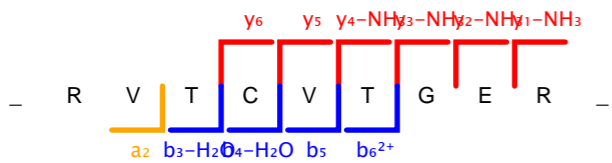
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_13

Scan number: 5923

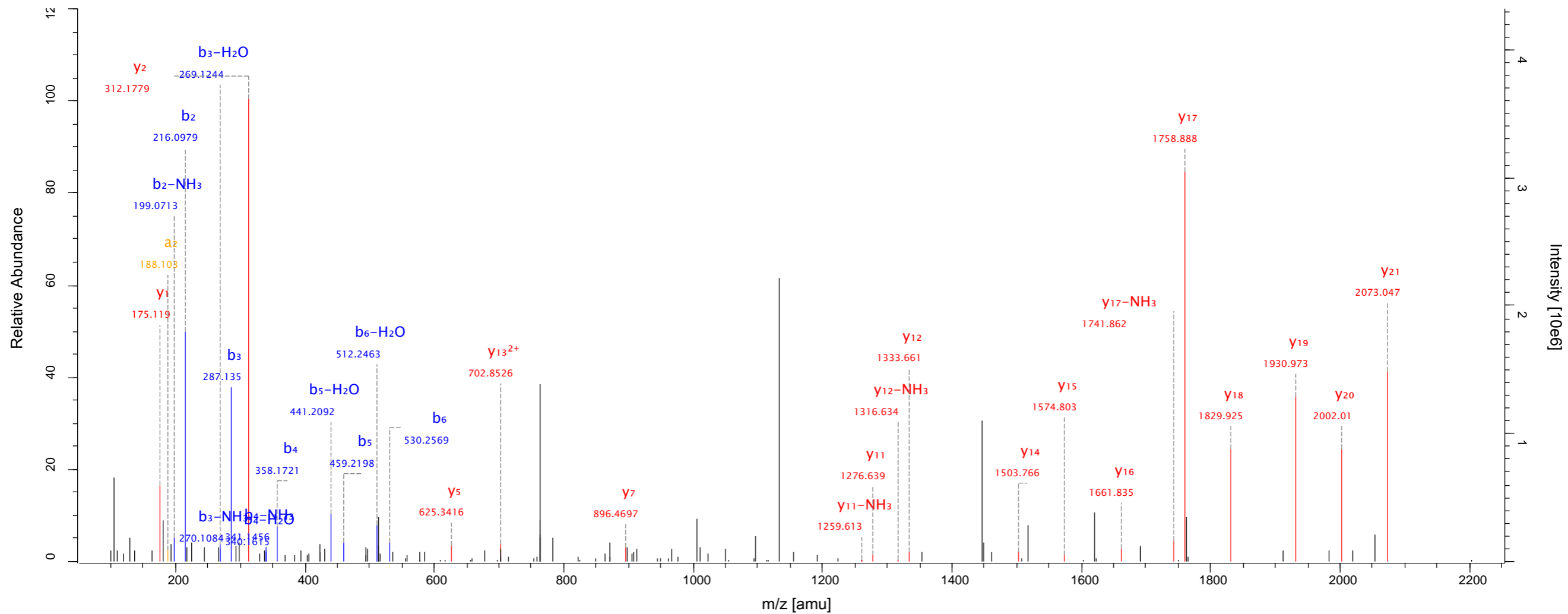
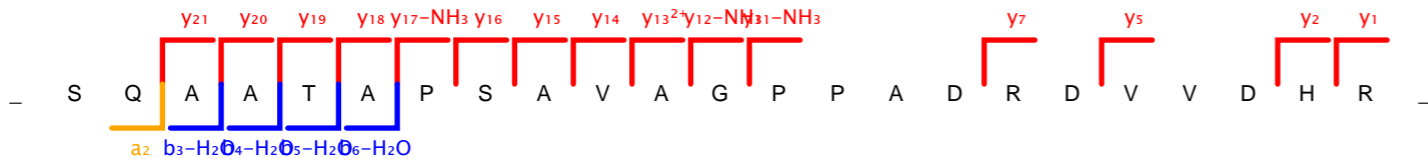
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 108.81

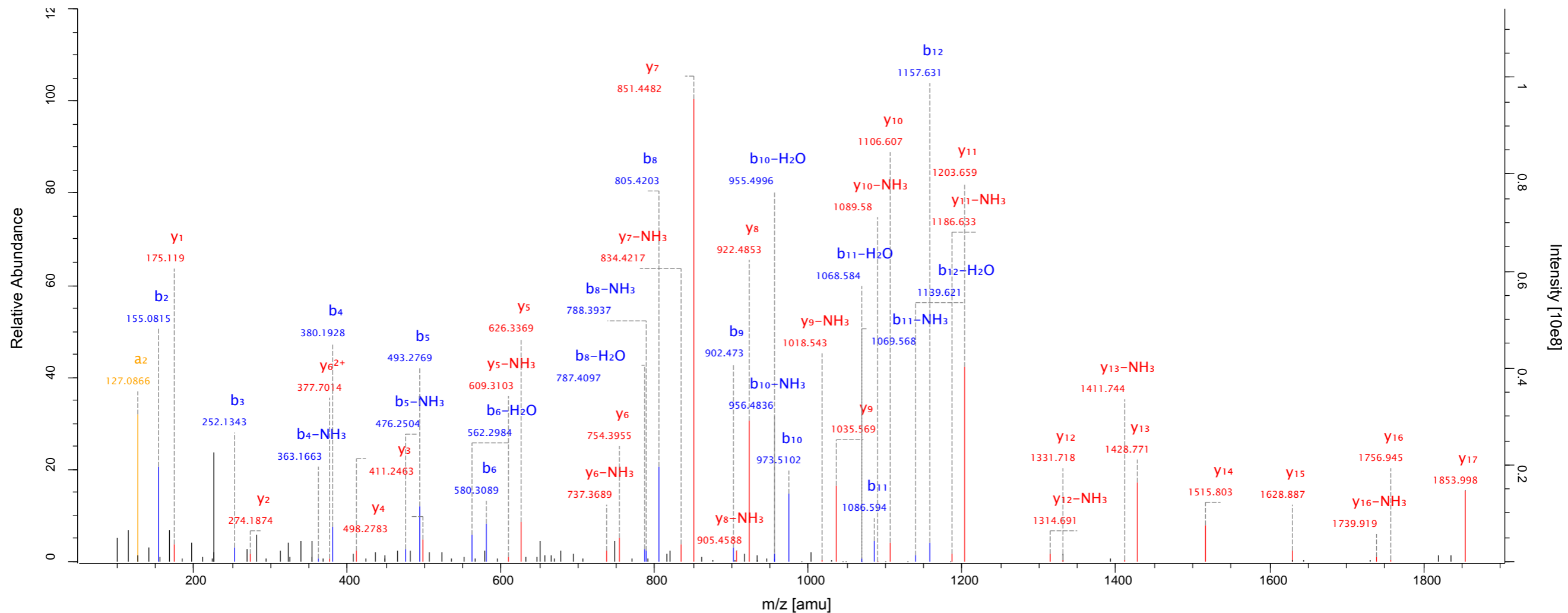
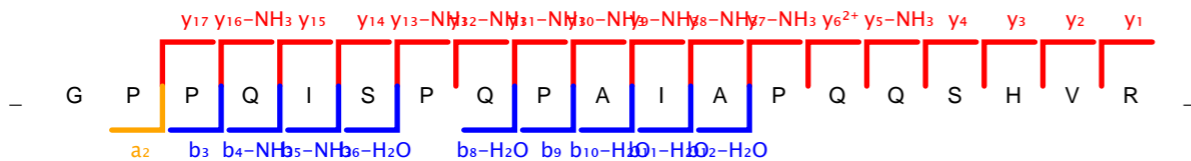
Method: FTMS; HCD



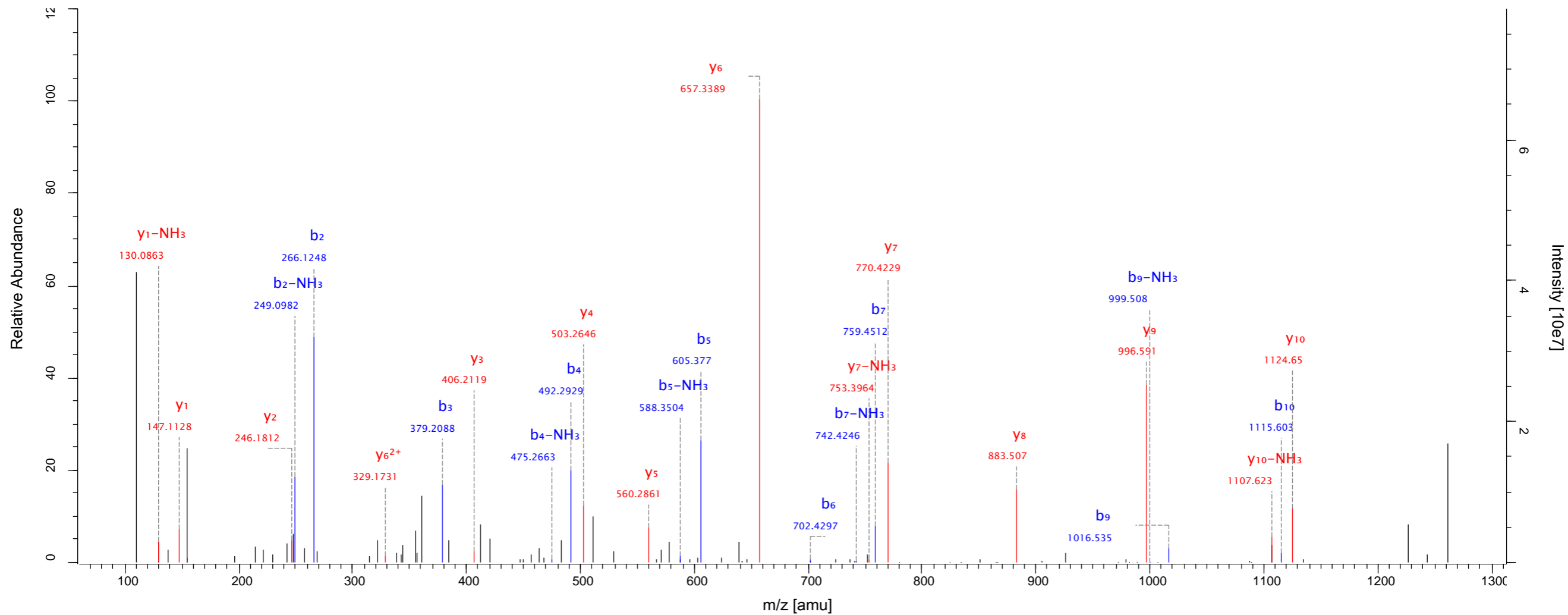
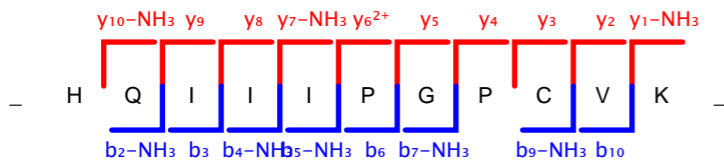
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_14
 Scan number: 16371
 Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>
 Peptide score: 137.43
 Method: FTMS; HCD



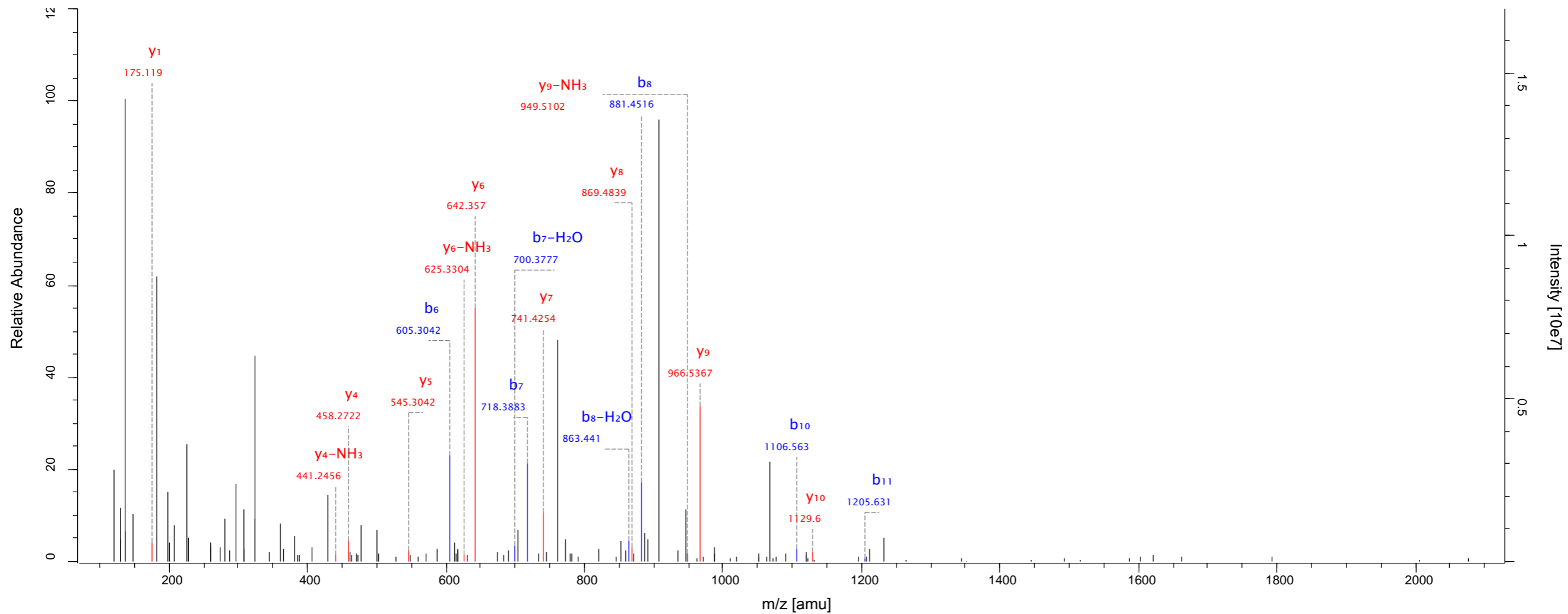
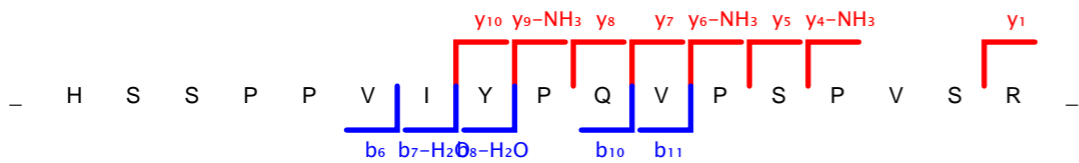
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_14
 Scan number: 22387
 Protein: ORFL251W + NC_006273.2[161872,163748]+ <unknown description>; ORFL253W.iORF1 + NC_006273.2[161917,163748]+ <unknown description>
 Peptide score: 290.62
 Method: FTMS; HCD



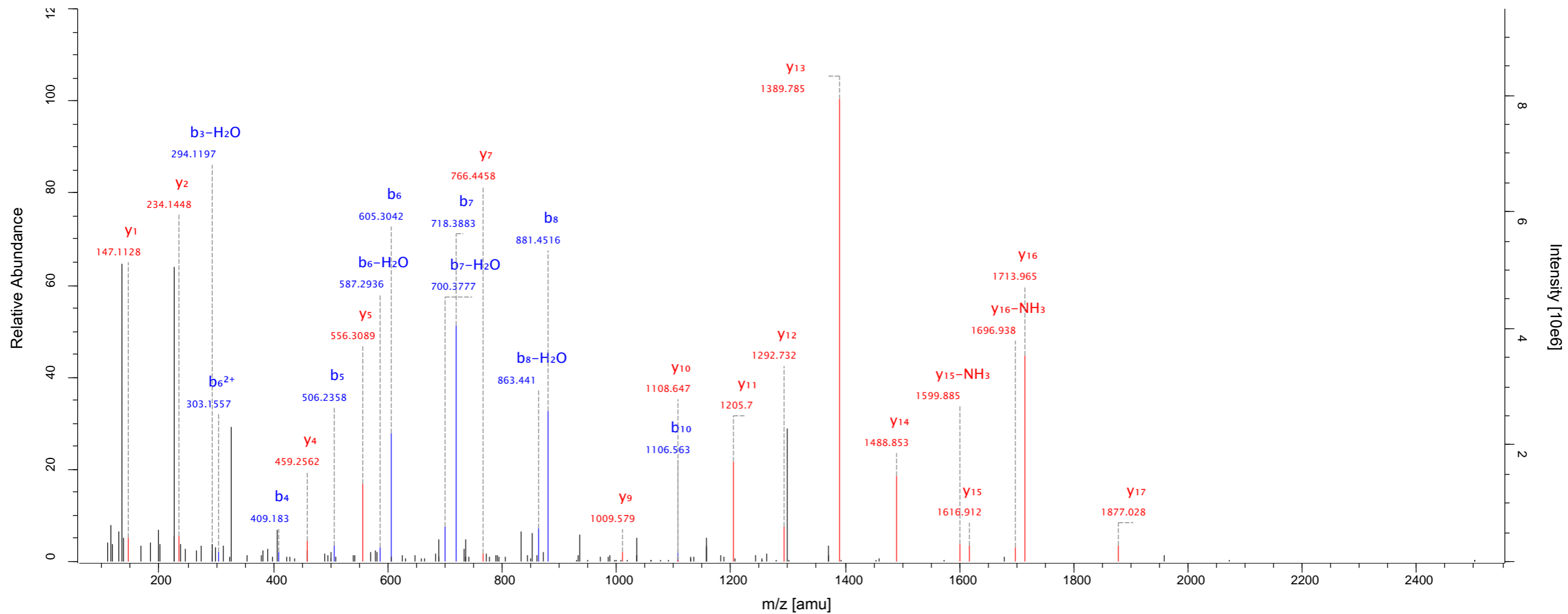
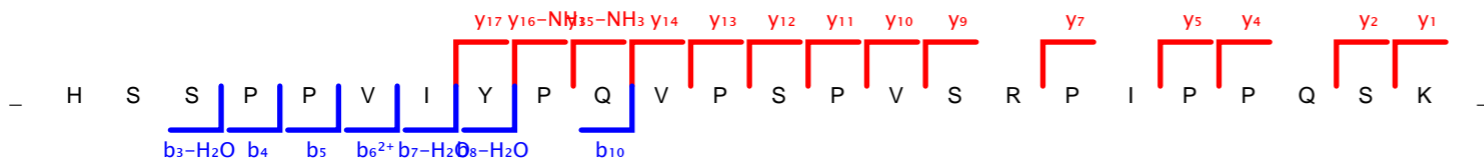
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_14
 Scan number: 23268
 Protein: ORFL251W + NC_006273.2[161872,163748]+ <unknown description>; ORFL253W.iORF1 + NC_006273.2[161917,163748]+ <unknown description>
 Peptide score: 169.65
 Method: FTMS; HCD



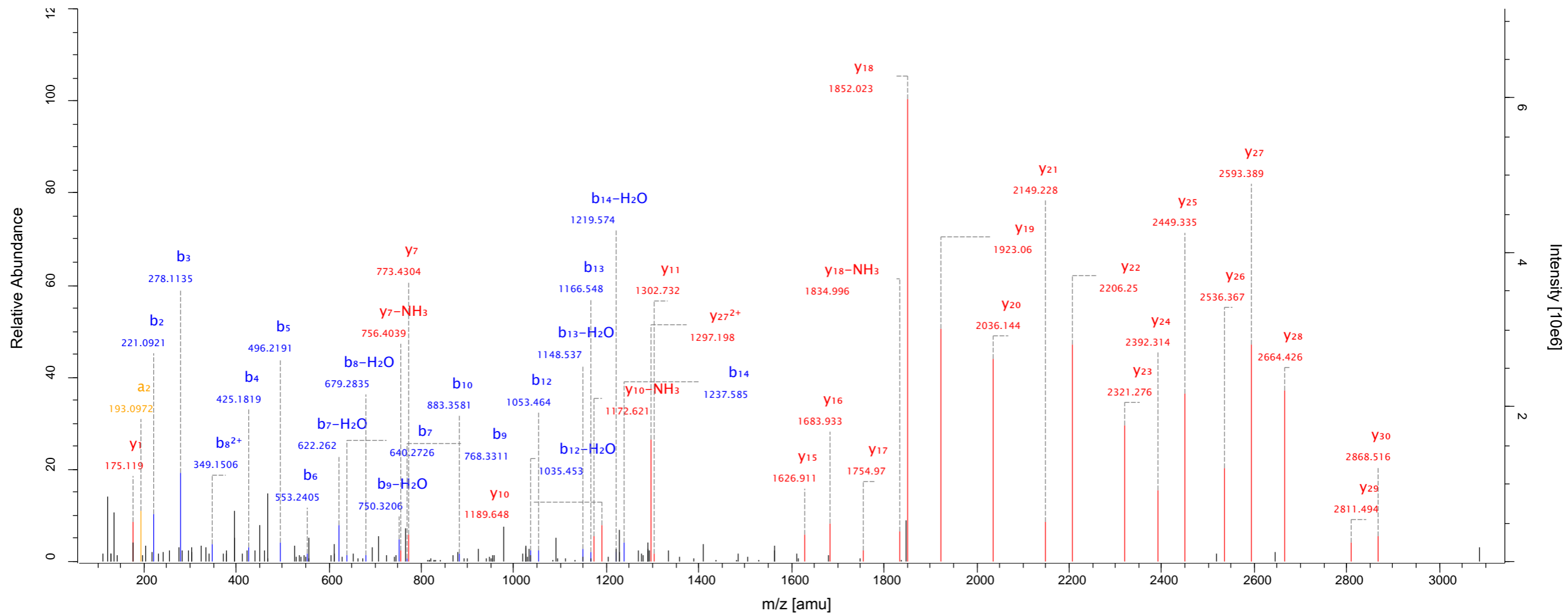
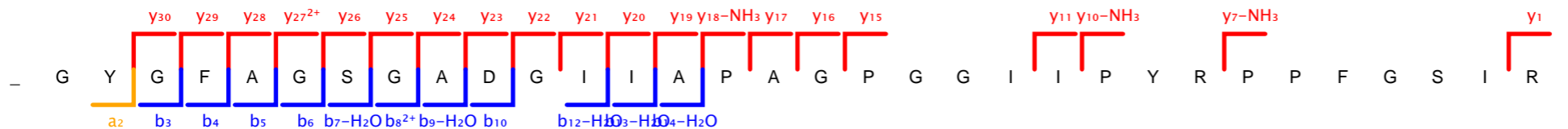
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_14
 Scan number: 31890
 Protein: ORFL251W + NC_006273.2[161872,163748]+ <unknown description>; ORFL253W.iORF1 + NC_006273.2[161917,163748]+ <unknown description>
 Peptide score: 45.46
 Method: FTMS; HCD



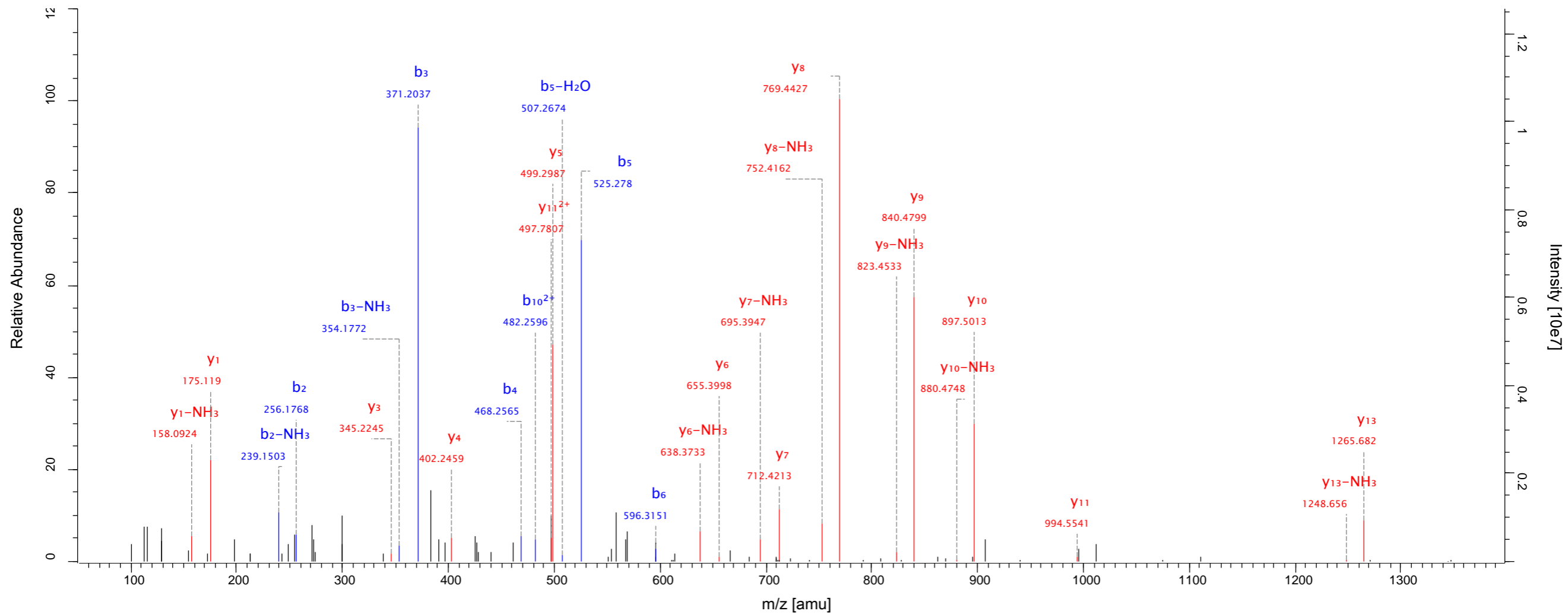
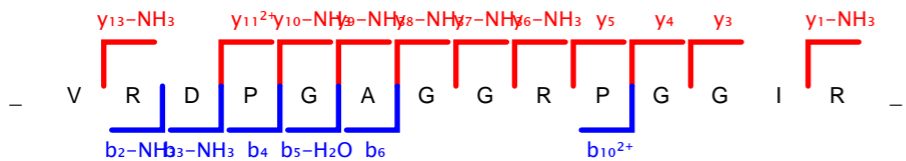
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_14
 Scan number: 33839
 Protein: ORFL251W + NC_006273.2[161872,163748]+ <unknown description>; ORFL253W.iORF1 + NC_006273.2[161917,163748]+ <unknown description>
 Peptide score: 125.88
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_14
 Scan number: 58621
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 176.53
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_14
 Scan number: 8179
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 143.31
 Method: FTMS; HCD



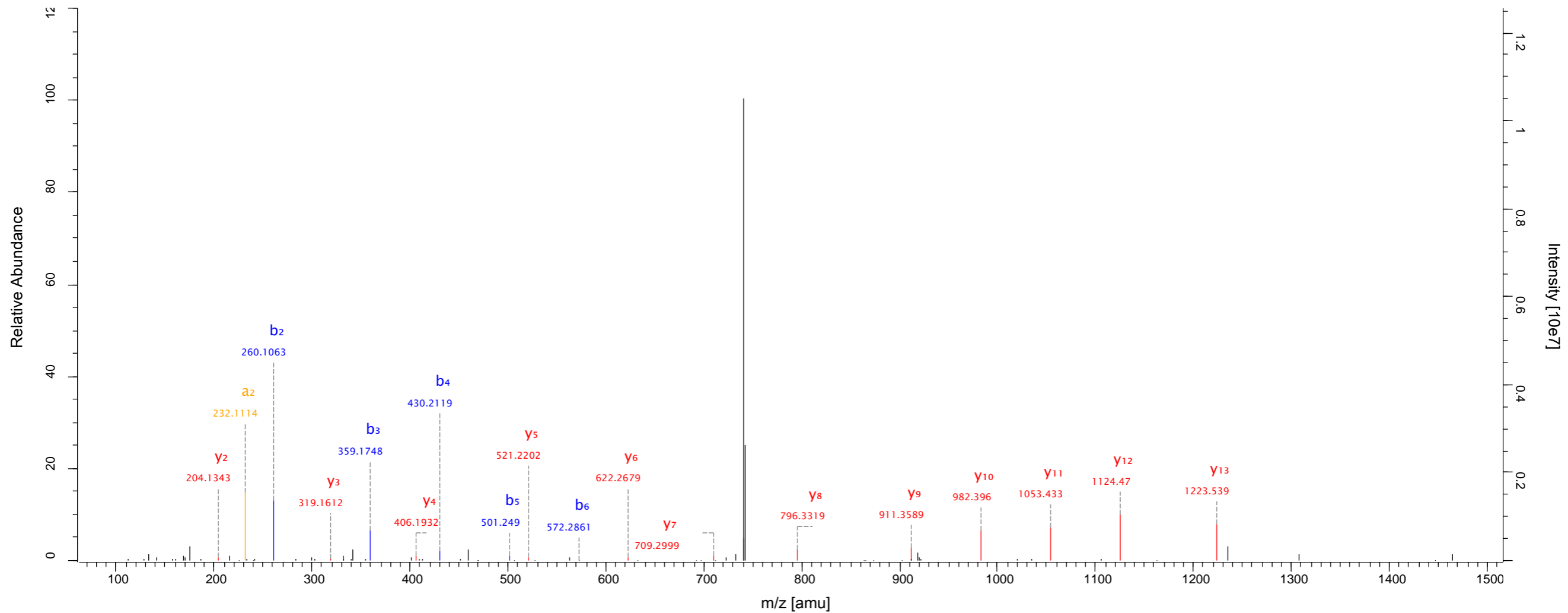
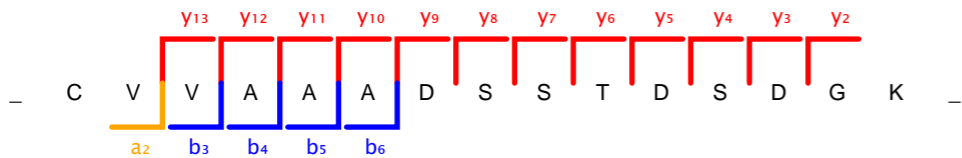
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_14

Scan number: 9685

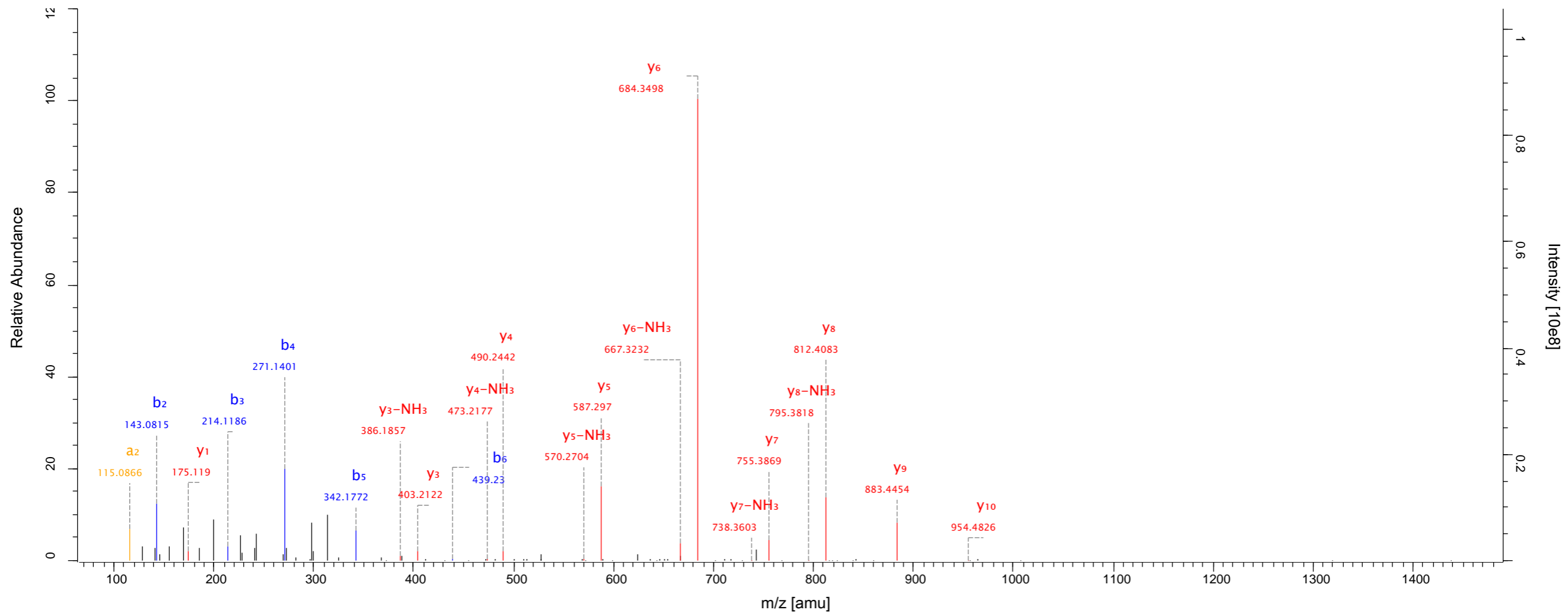
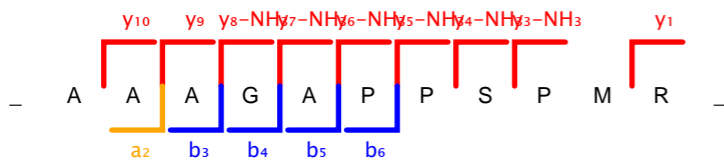
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 99.53

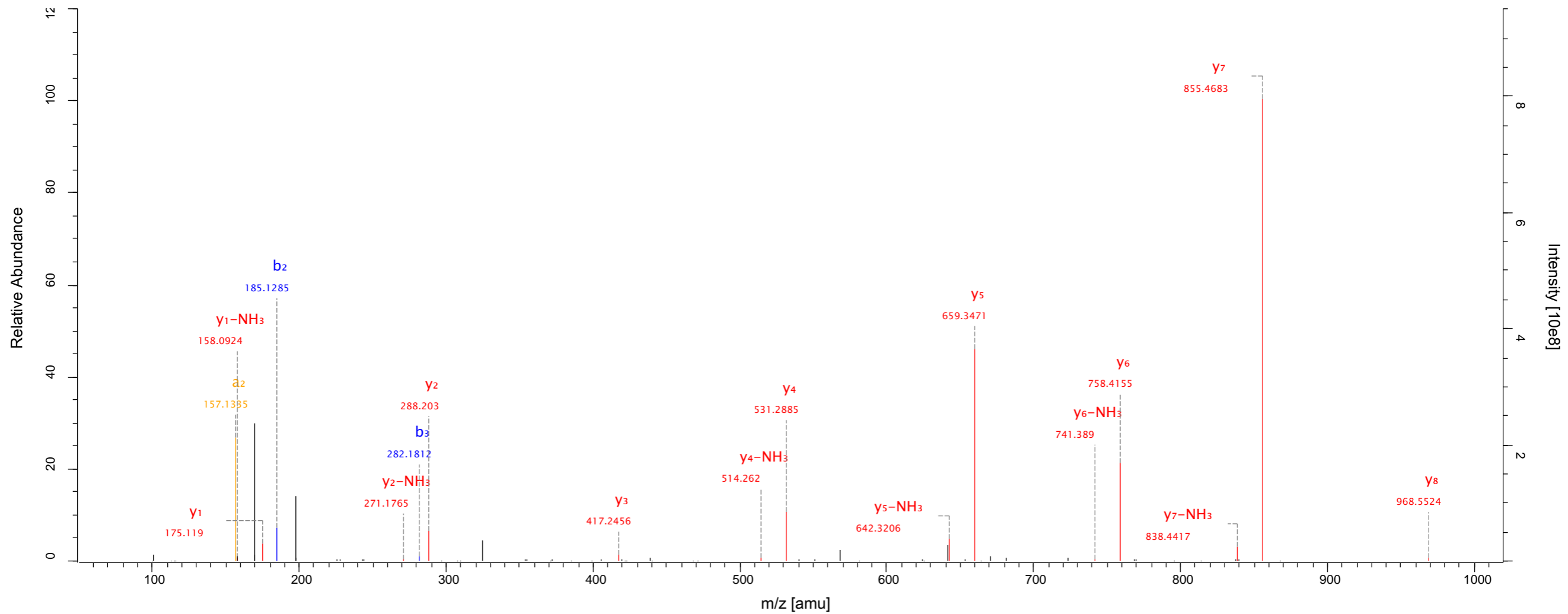
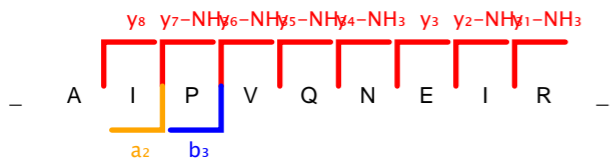
Method: FTMS; HCD



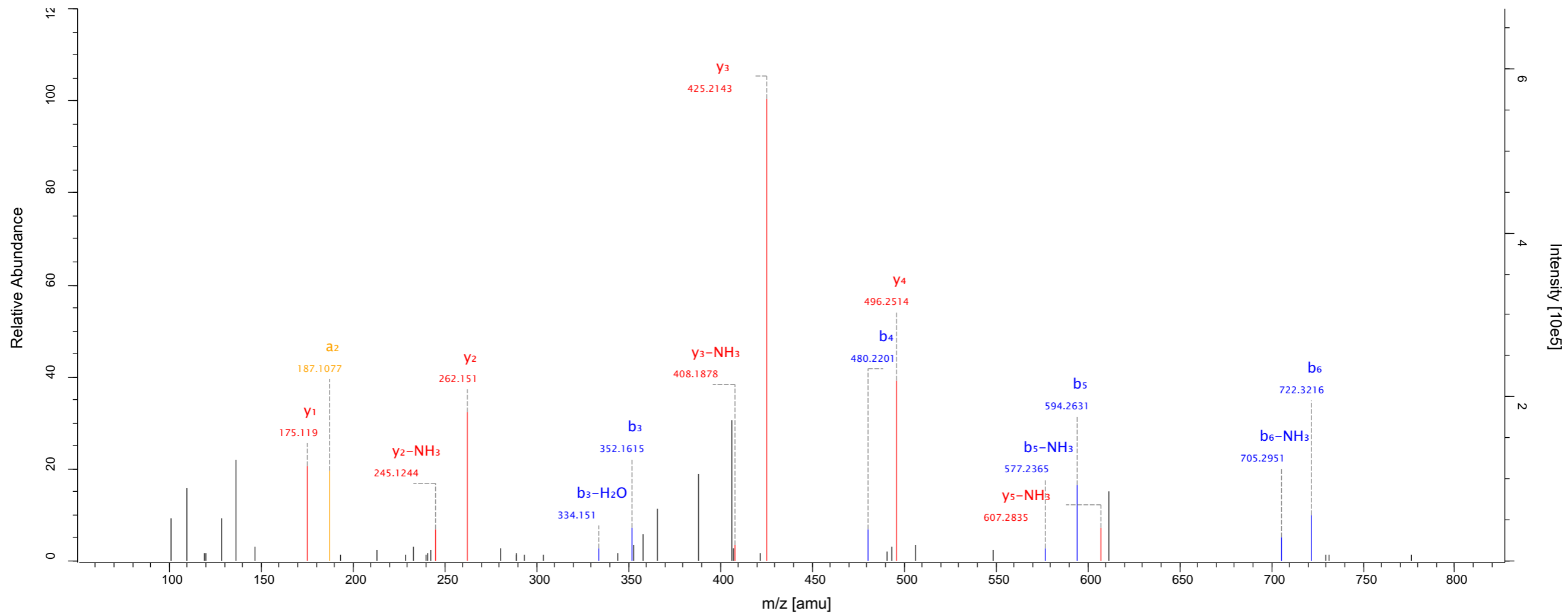
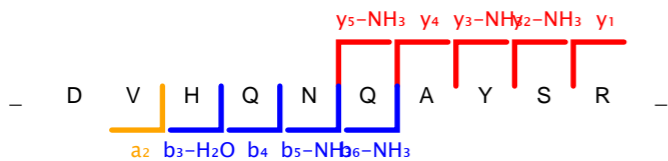
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_15
Scan number: 11201
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 130.66
Method: FTMS; HCD



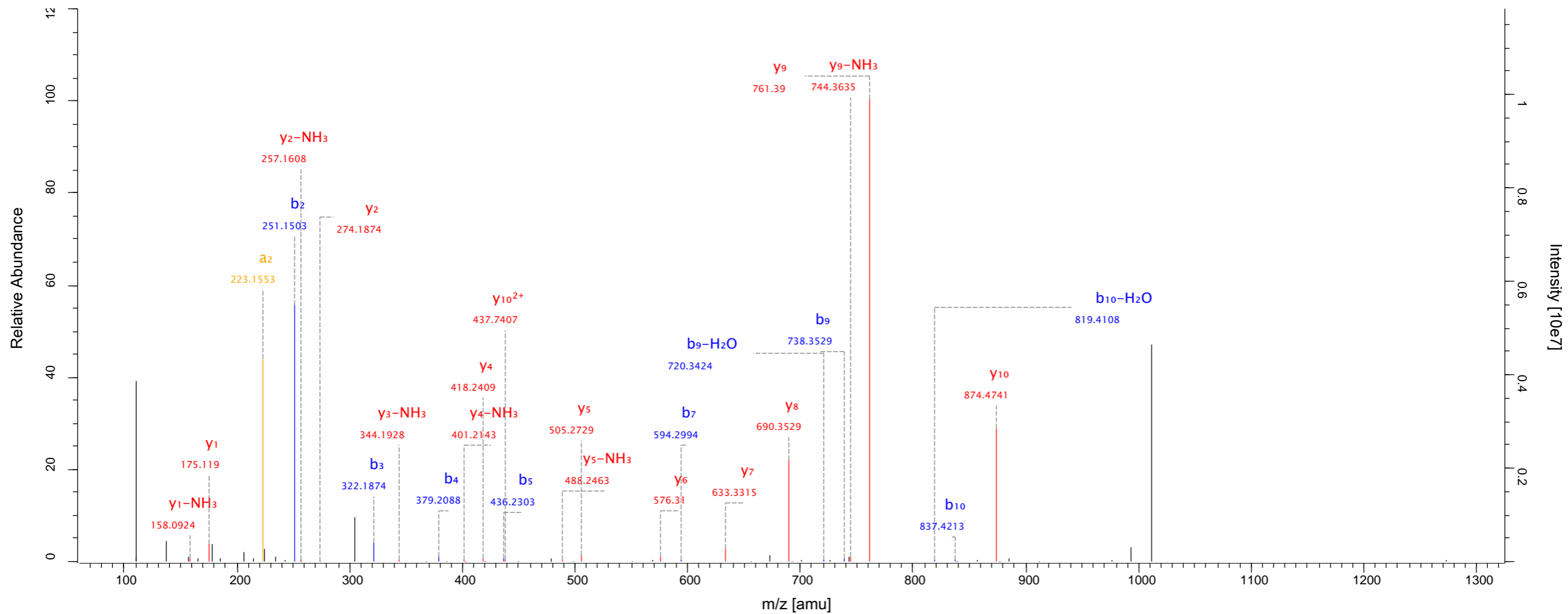
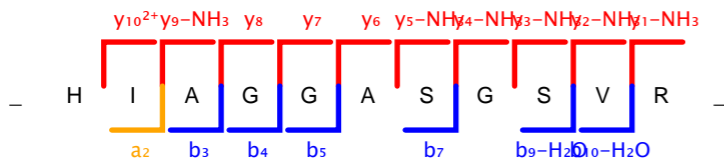
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_15
Scan number: 22097
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 130.27
Method: FTMS; HCD



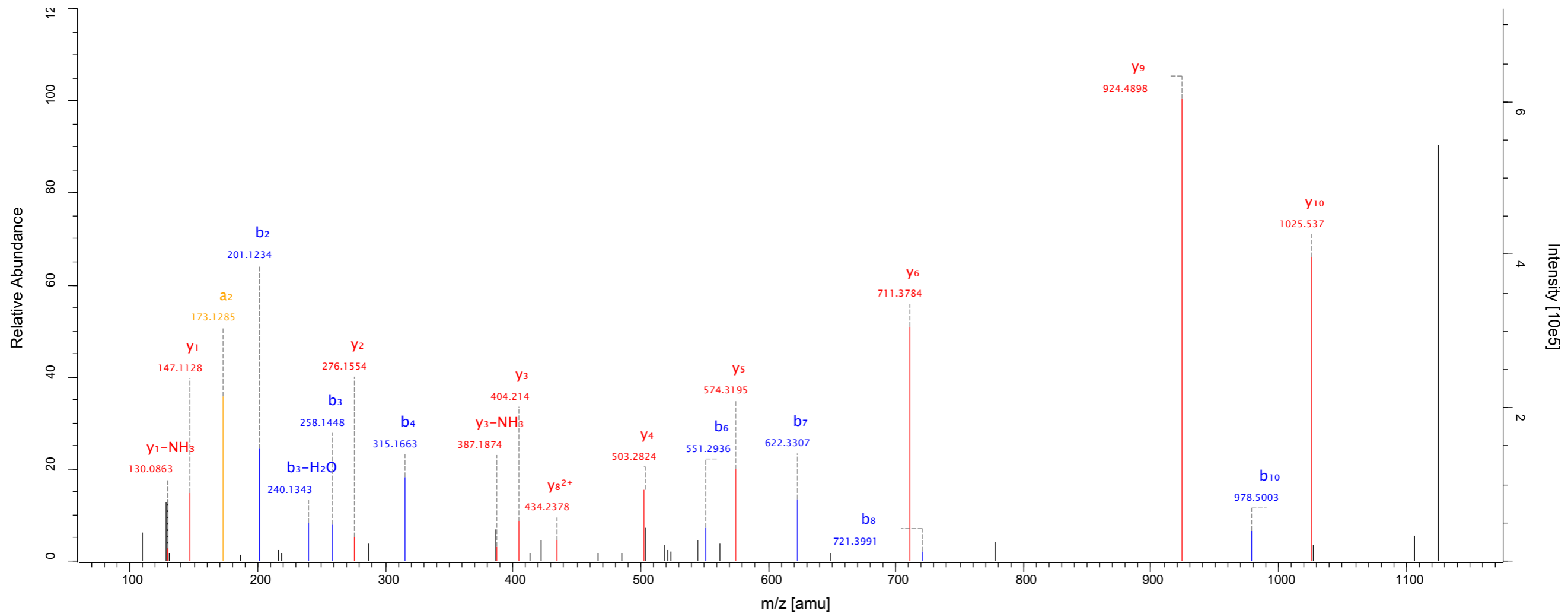
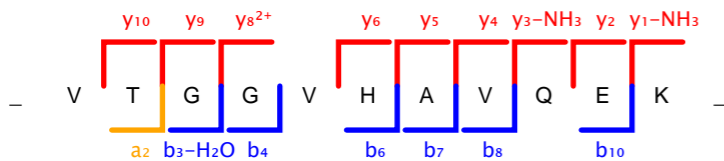
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_15
Scan number: 5033
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 89.05
Method: FTMS; HCD



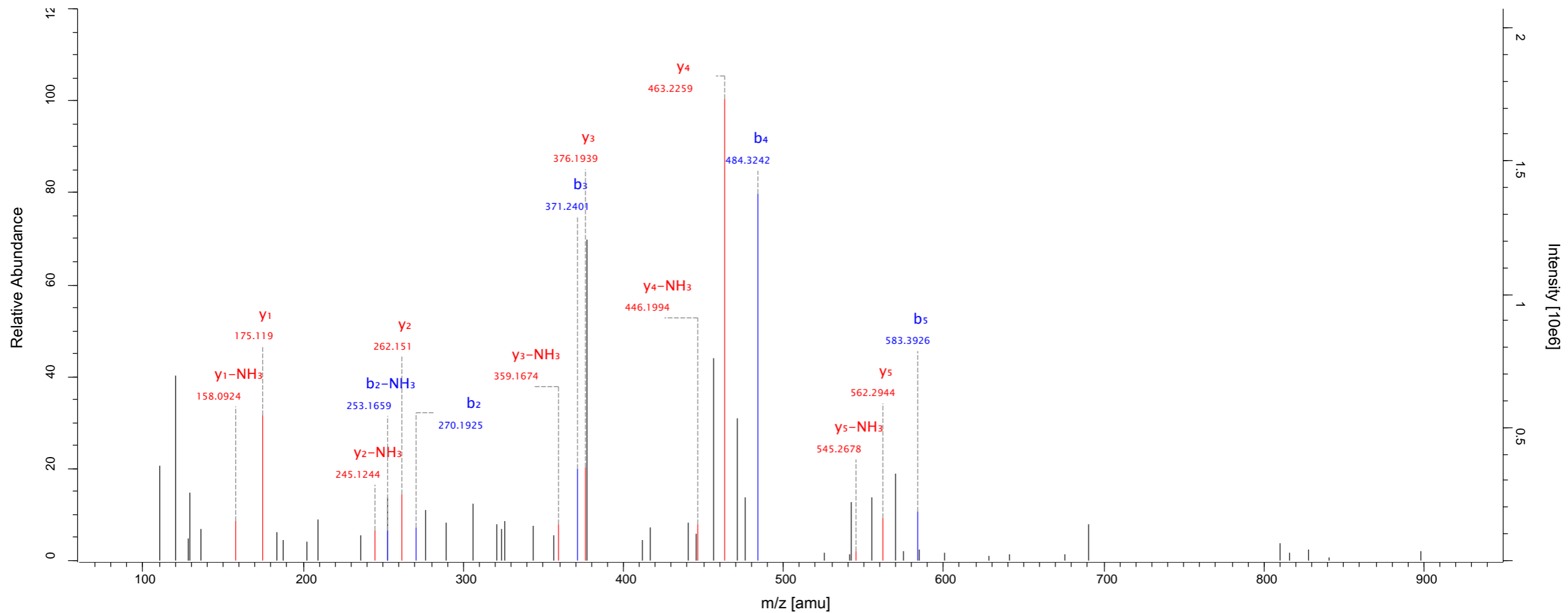
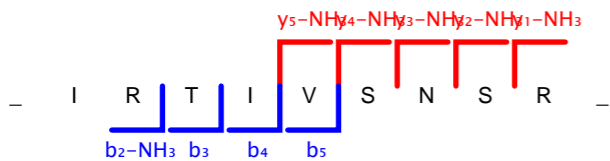
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_15
 Scan number: 5387
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 167.12
 Method: FTMS; HCD



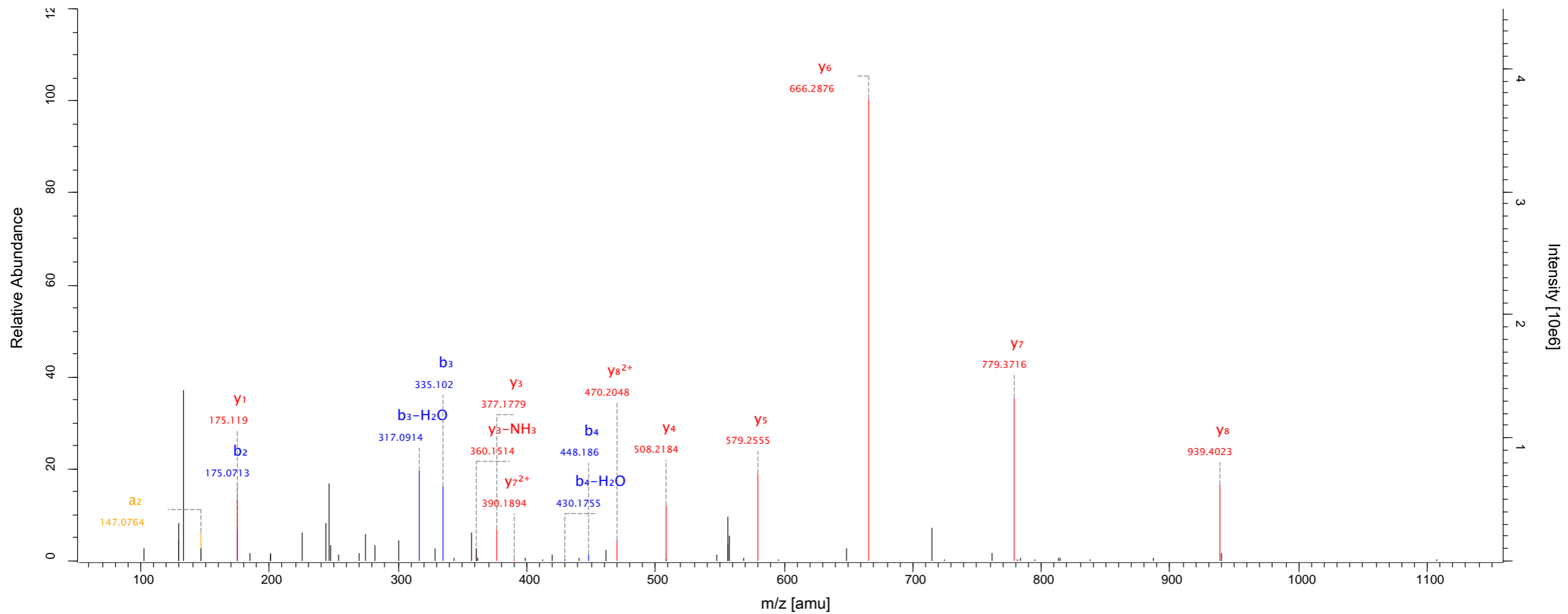
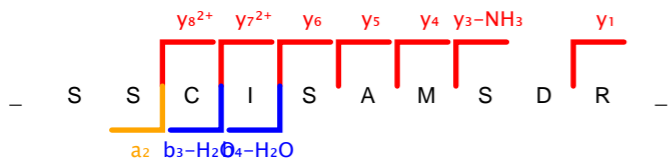
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_15
Scan number: 6241
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 140.3
Method: FTMS; HCD



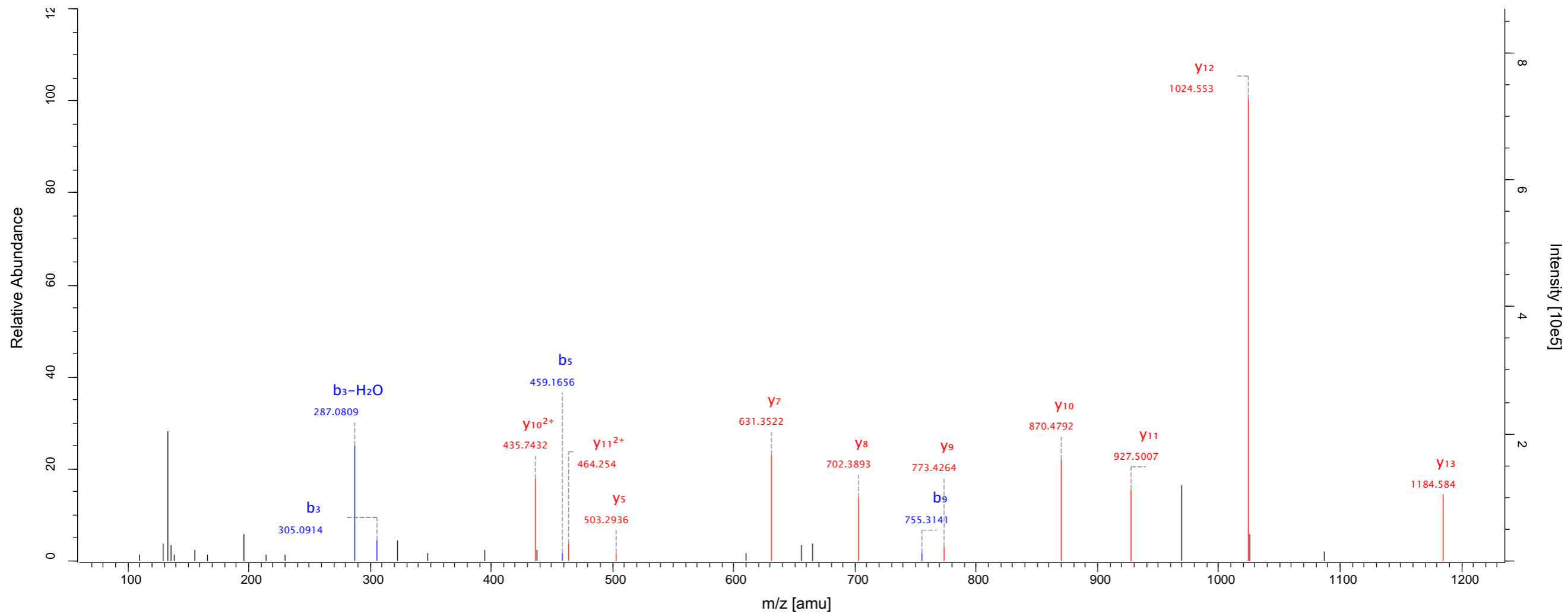
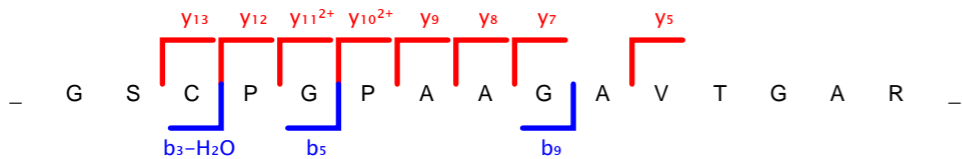
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_18
Scan number: 10129
Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
Peptide score: 78.26
Method: FTMS; HCD



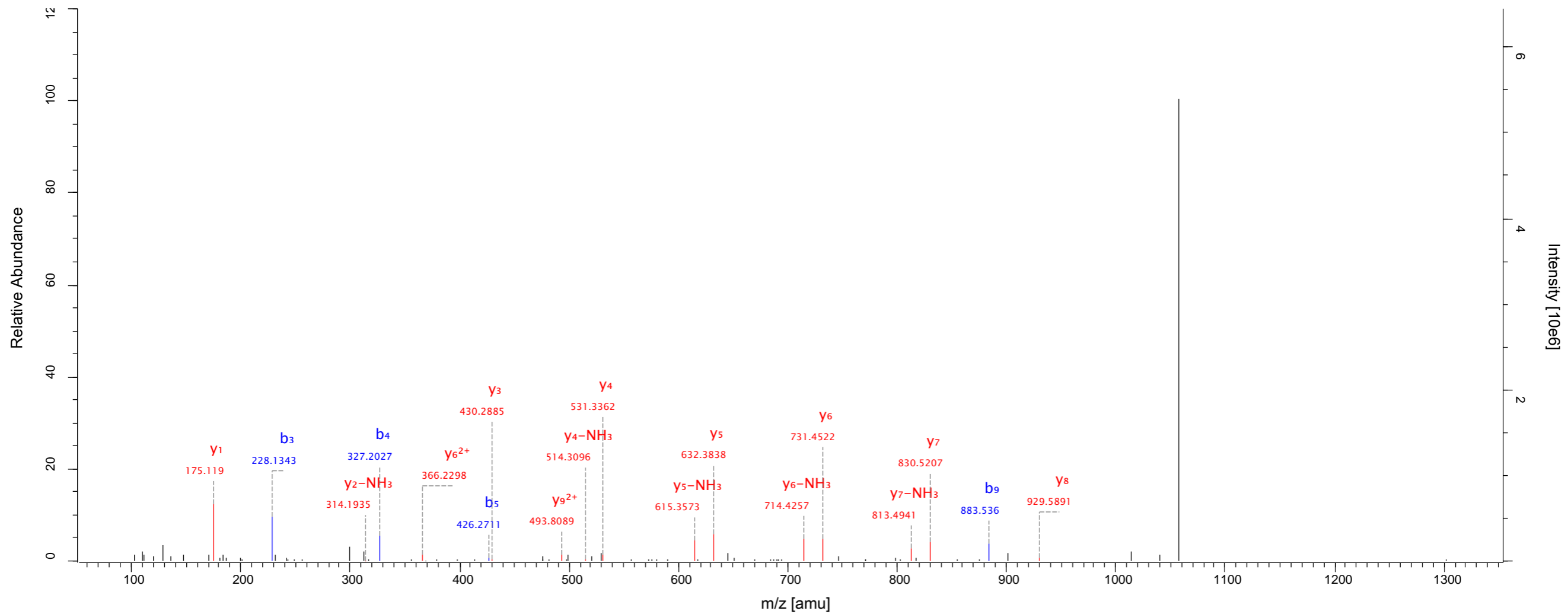
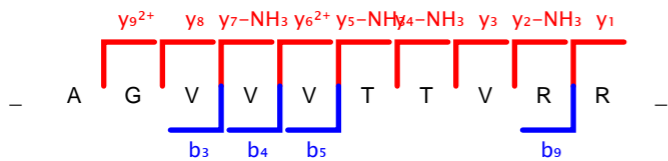
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 10823
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 87.91
Method: FTMS; HCD



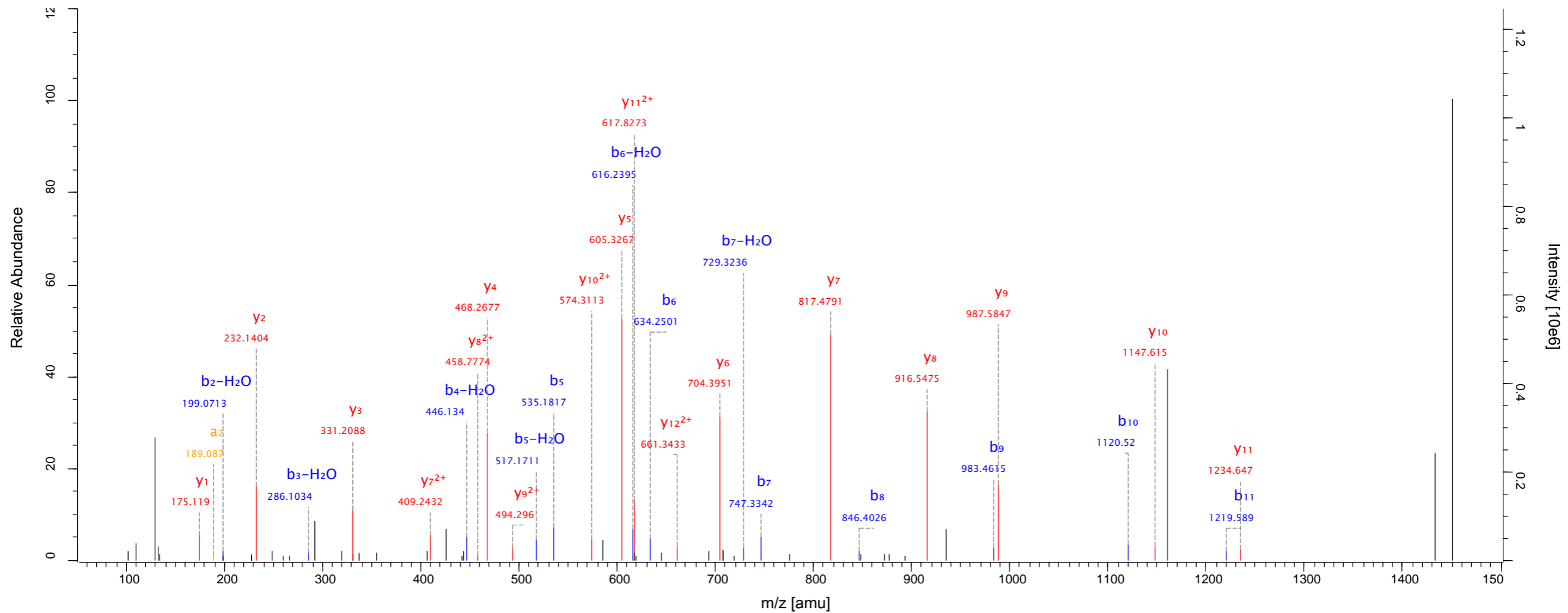
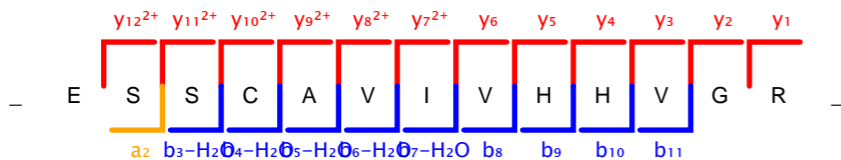
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 11007
Protein: ORFL235W + NC_006273.2[149239,150022]+ <unknown description>
Peptide score: 100.55
Method: FTMS; HCD



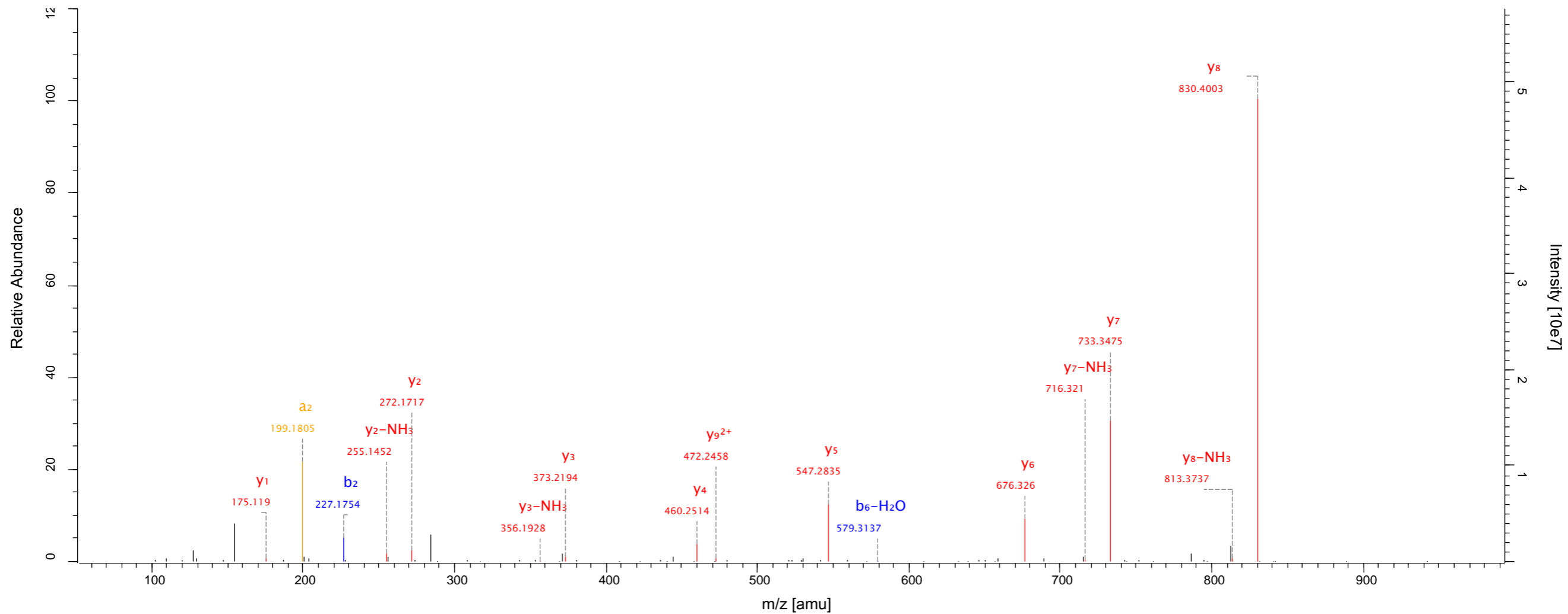
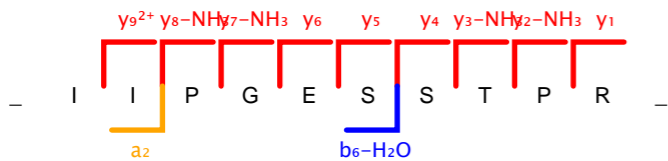
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 11044
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 122.13
Method: FTMS; HCD



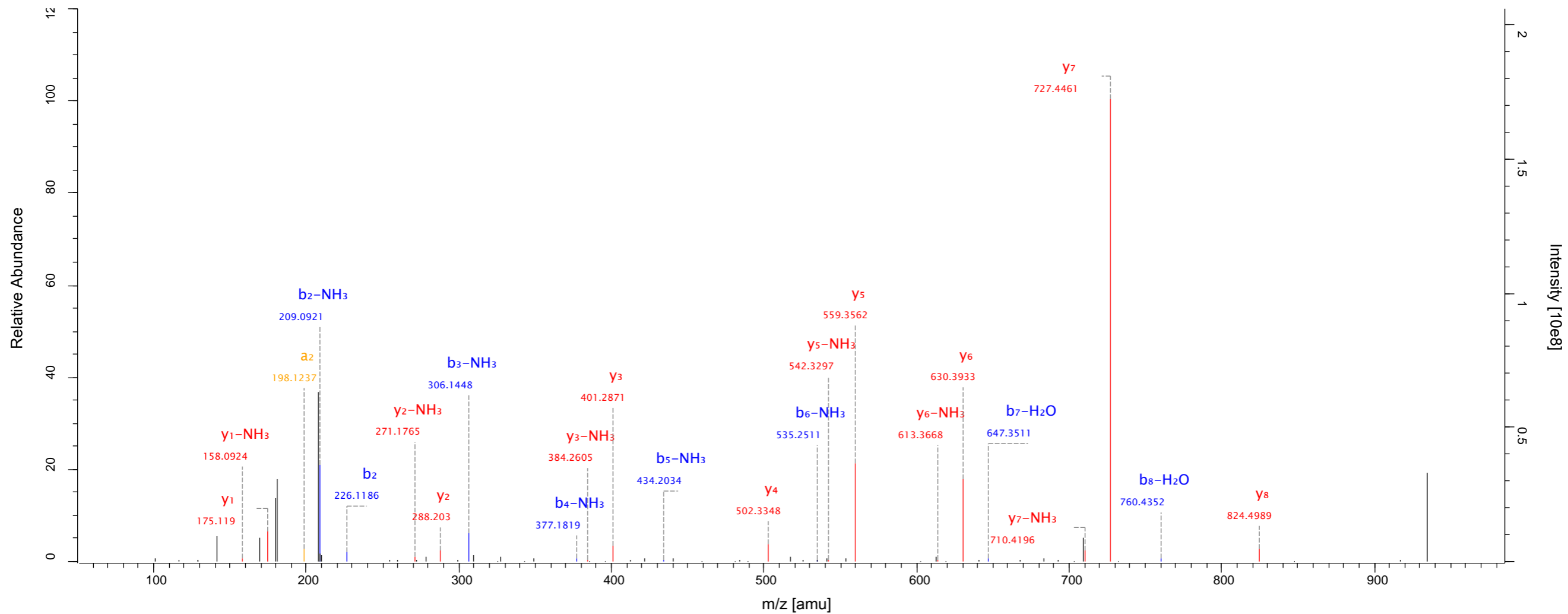
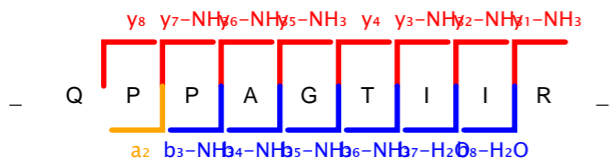
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_18
 Scan number: 11986
 Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
 Peptide score: 198.81
 Method: FTMS; HCD



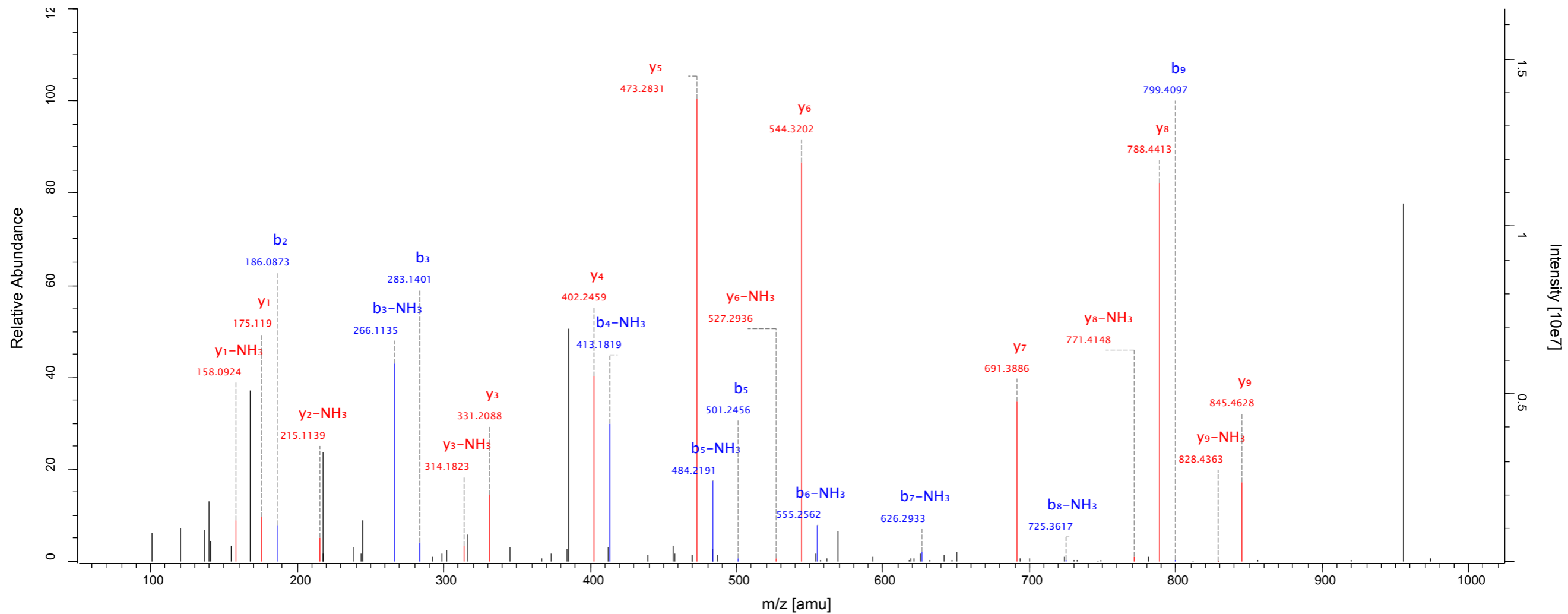
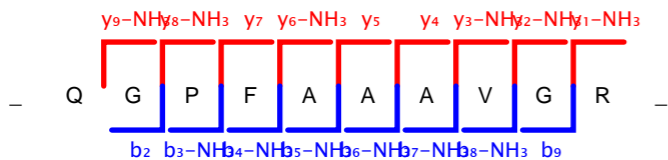
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 12489
Protein: ORFL137W.iORF1 + NC_006273.2[75299,75529]+ <unknown description>
Peptide score: 84.61
Method: FTMS; HCD



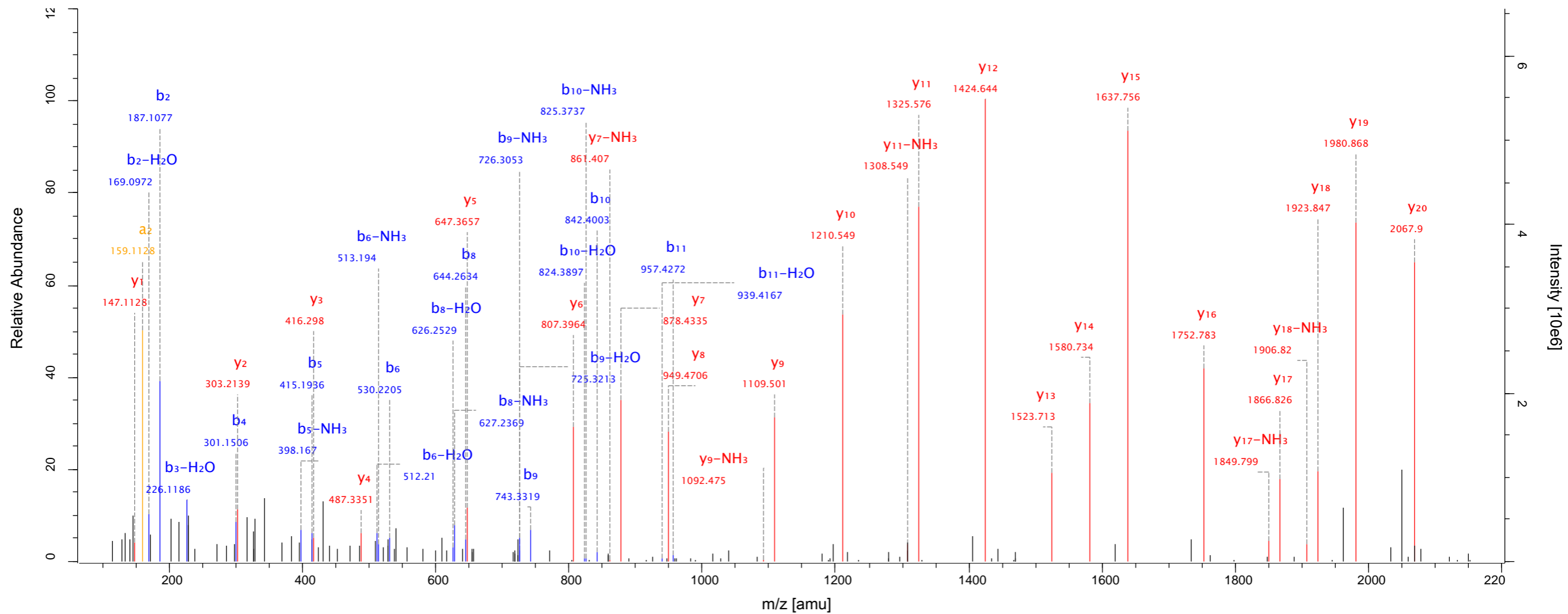
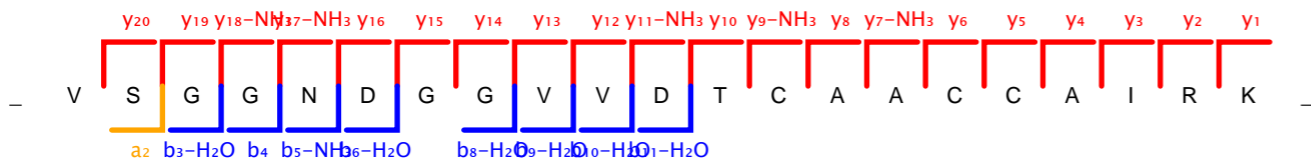
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 16808
 Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
 Peptide score: 116.25
 Method: FTMS; HCD



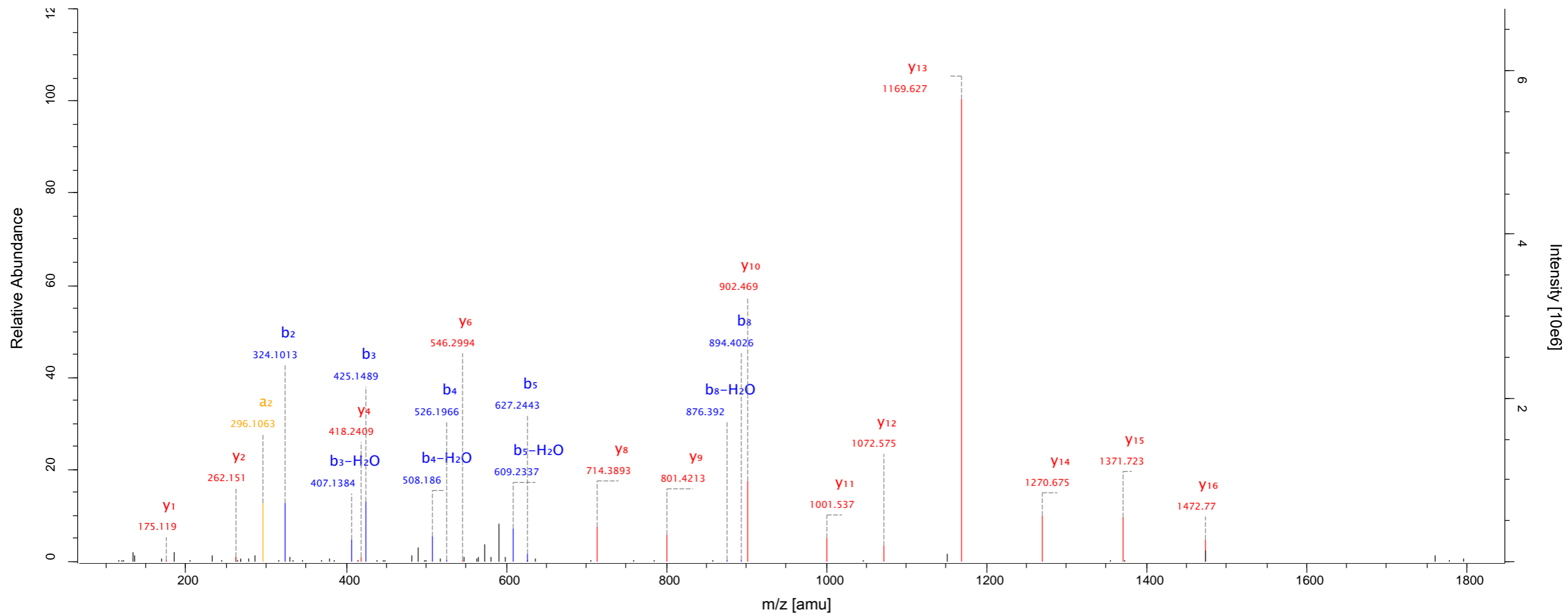
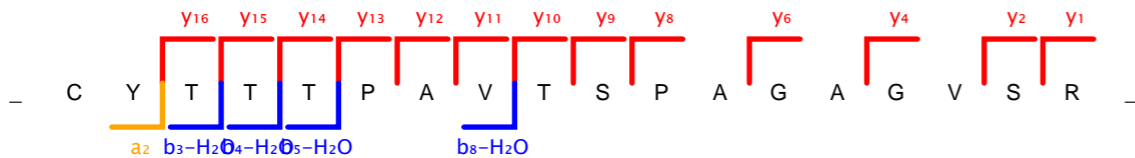
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 18638
 Protein: ORFL121C - NC_006273.2[65579,65851]- <unknown description>
 Peptide score: 108.43
 Method: FTMS; HCD



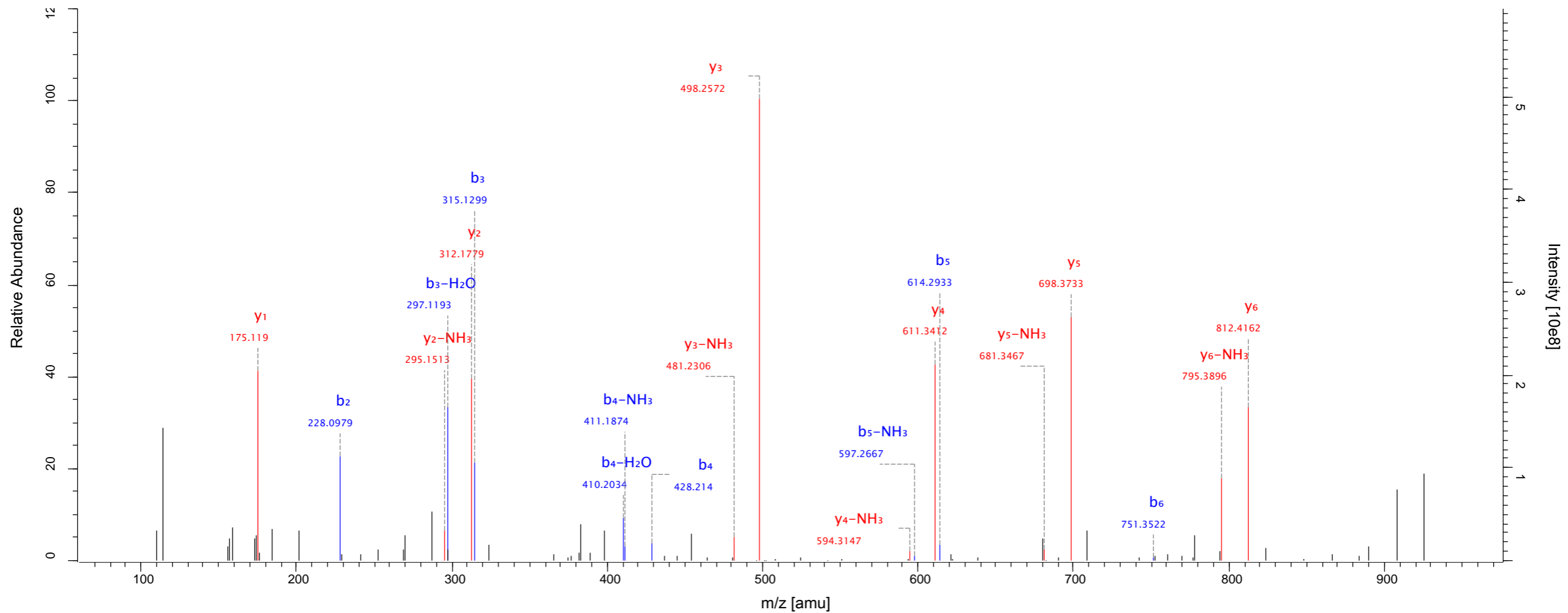
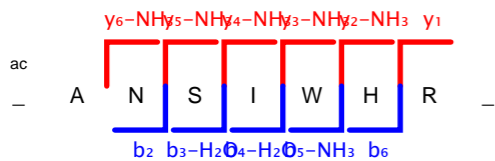
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 19088
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 255.19
 Method: FTMS; HCD



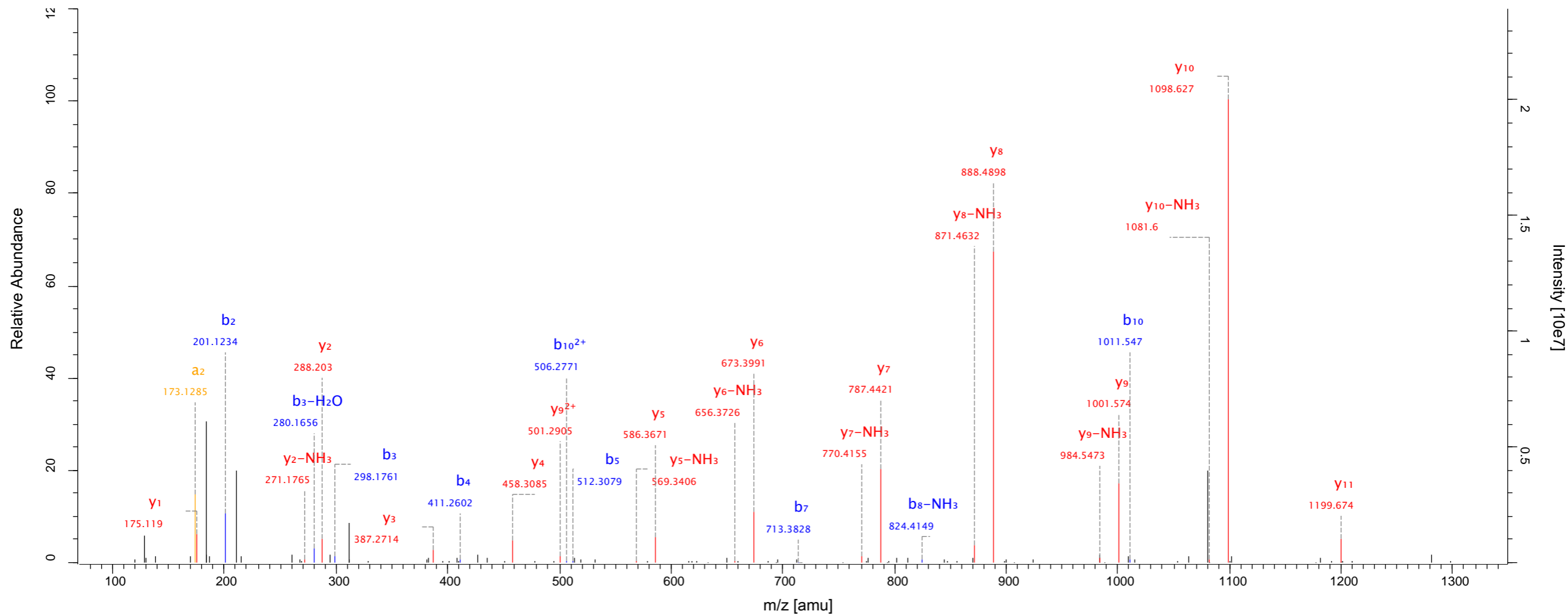
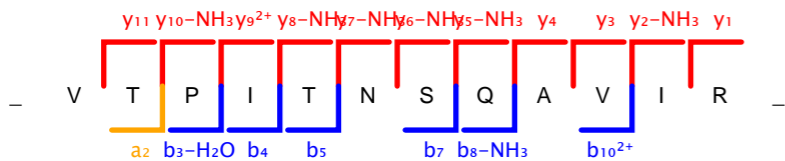
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 19728
 Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
 Peptide score: 111.43
 Method: FTMS; HCD



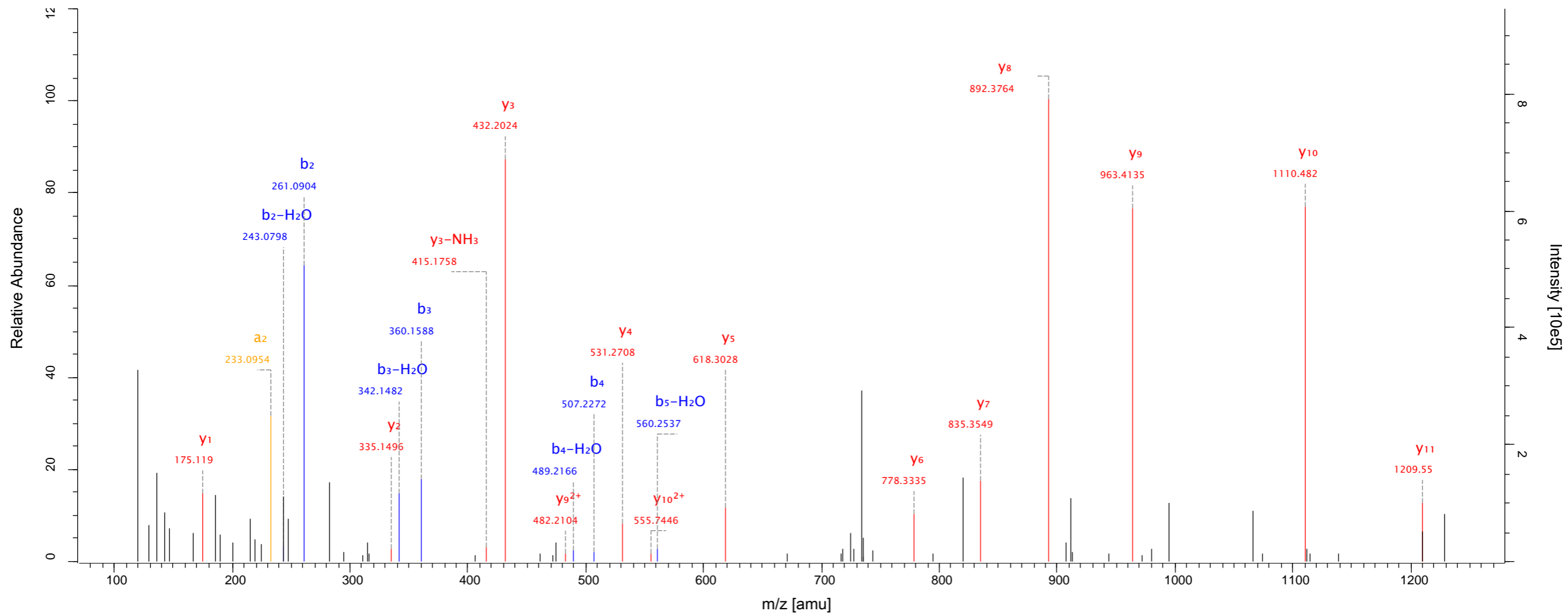
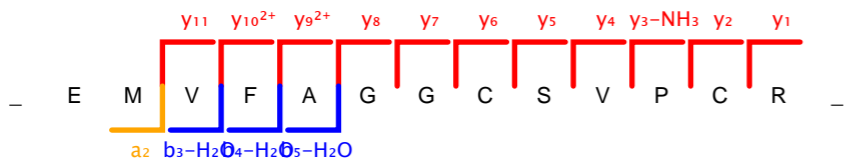
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 21557
Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
Peptide score: 161.68
Method: FTMS; HCD



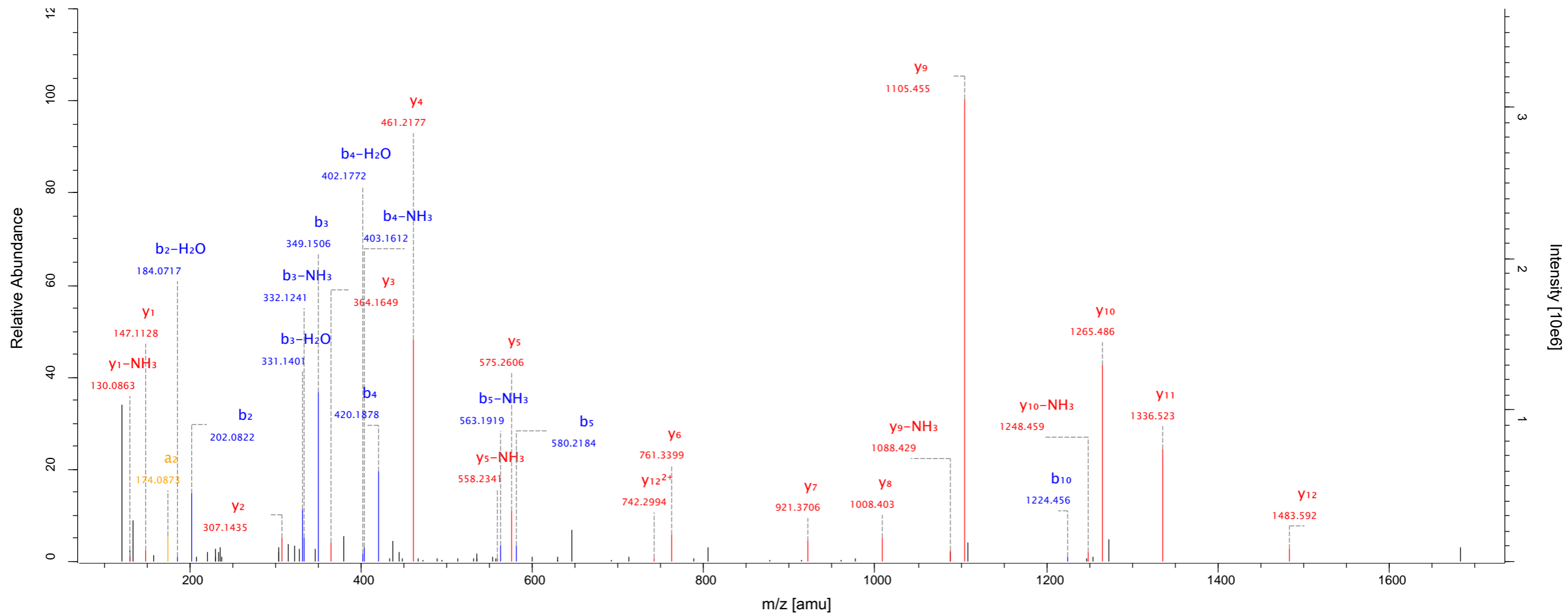
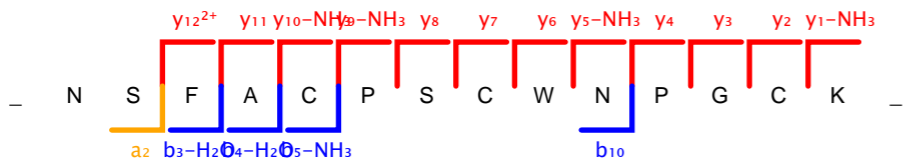
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 23426
 Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
 Peptide score: 155.33
 Method: FTMS; HCD



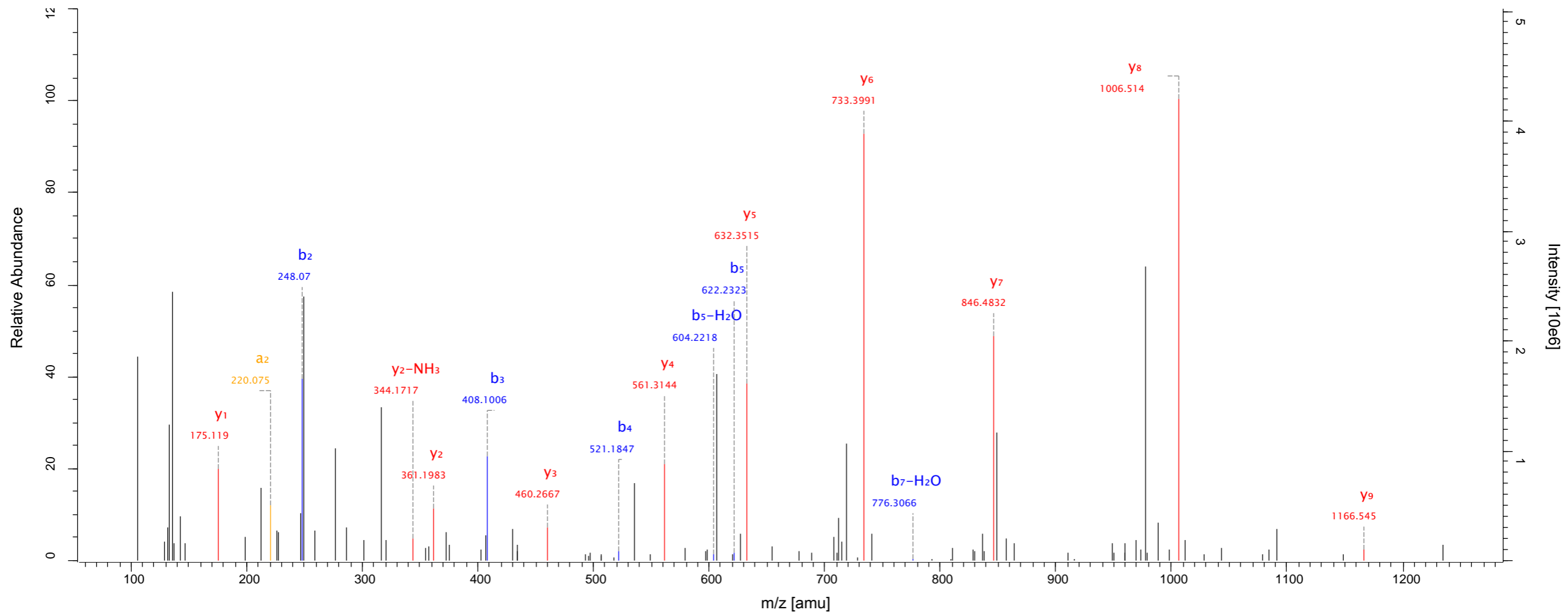
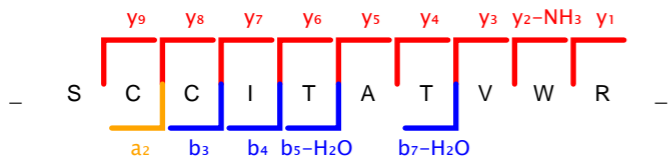
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 24830
 Protein: ORFL159W + NC_006273.2[101475,109052]+ <unknown description>; ORFL159W.iORF1 + NC_006273.2[101526,109052]+ <unknown description>
 Peptide score: 132.32
 Method: FTMS; HCD



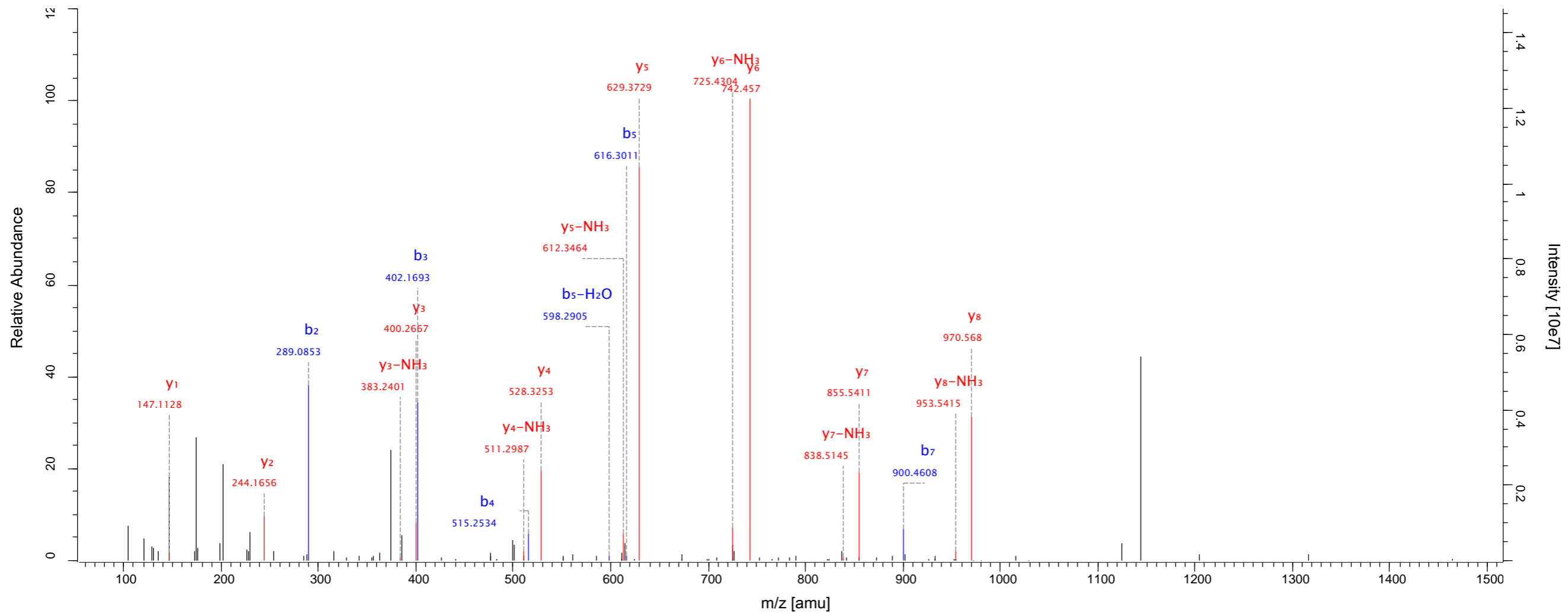
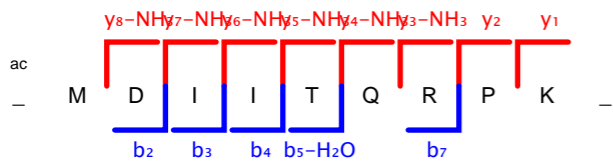
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 26132
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 157.56
 Method: FTMS; HCD



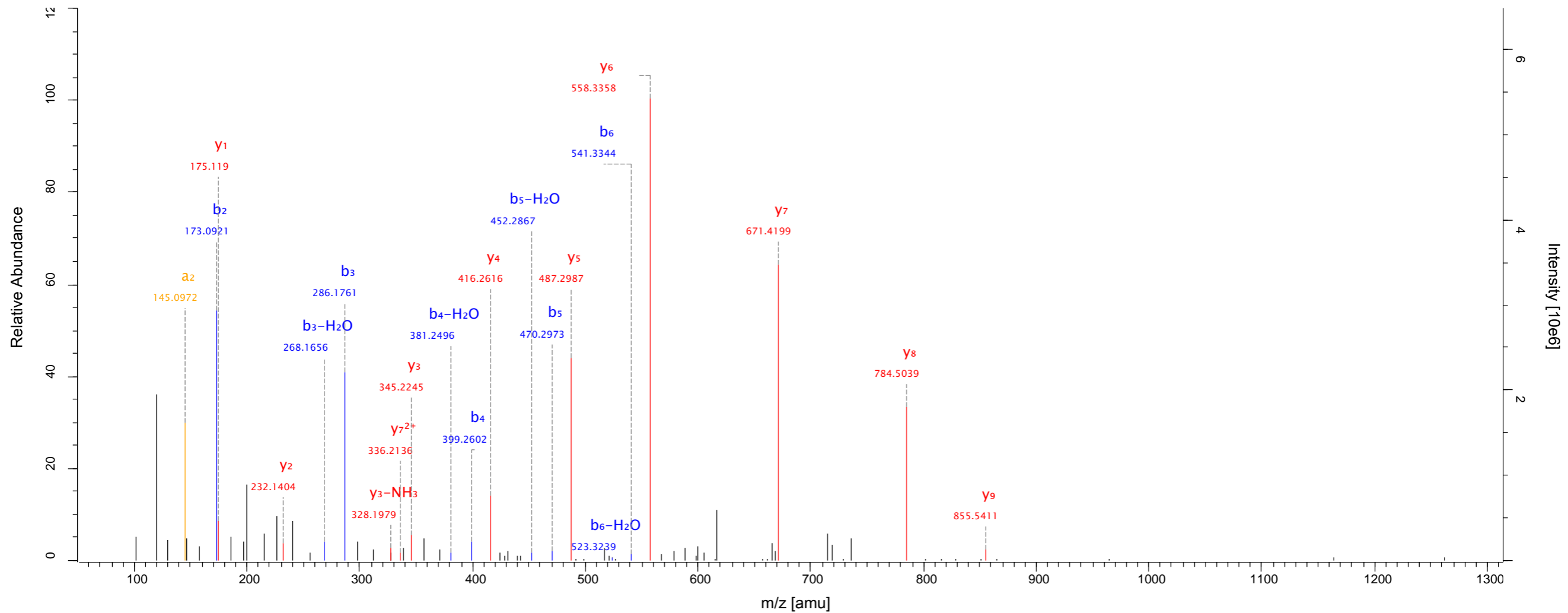
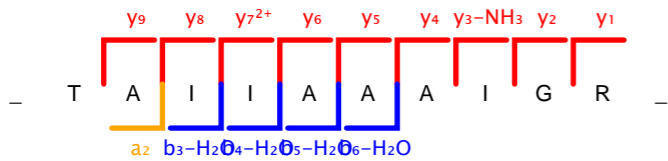
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 28228
Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
Peptide score: 90.11
Method: FTMS; HCD



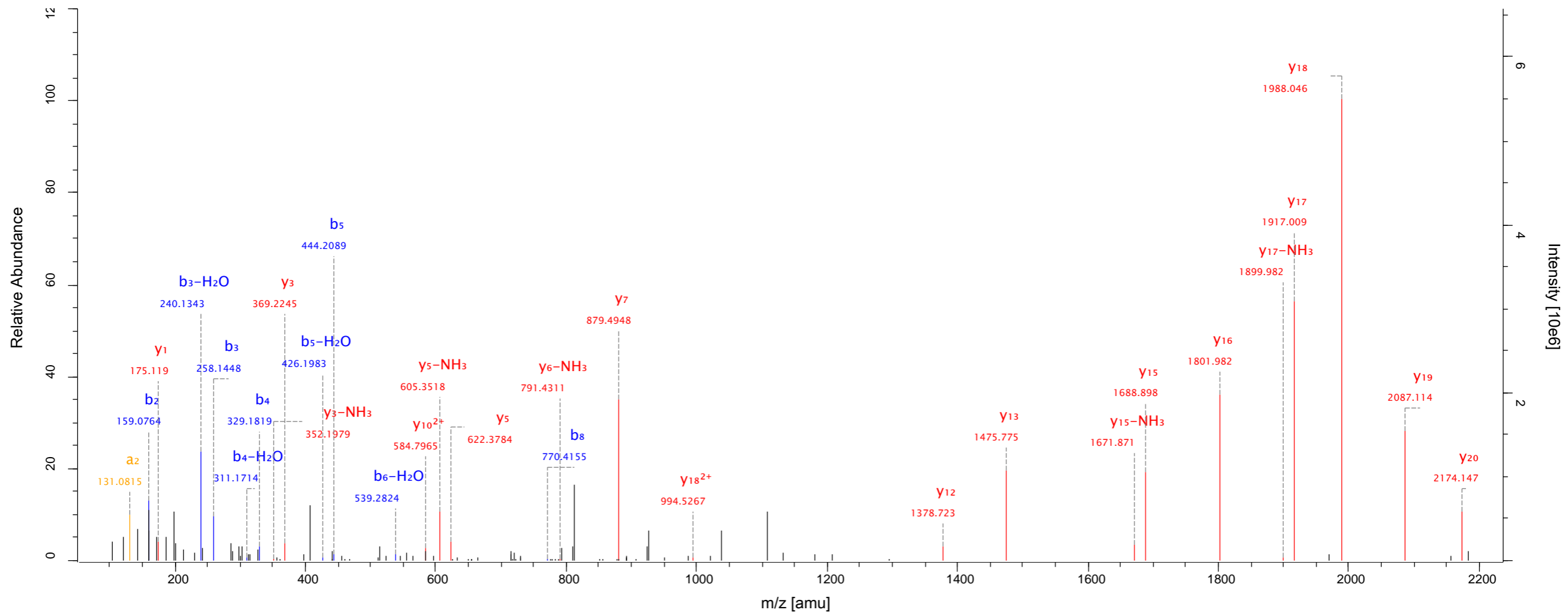
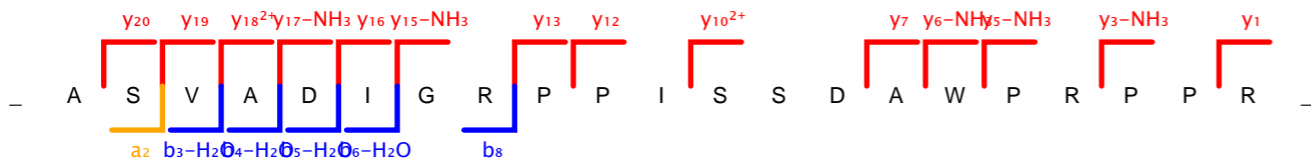
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_18
Scan number: 33776
Protein: ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
Peptide score: 125.82
Method: FTMS; HCD



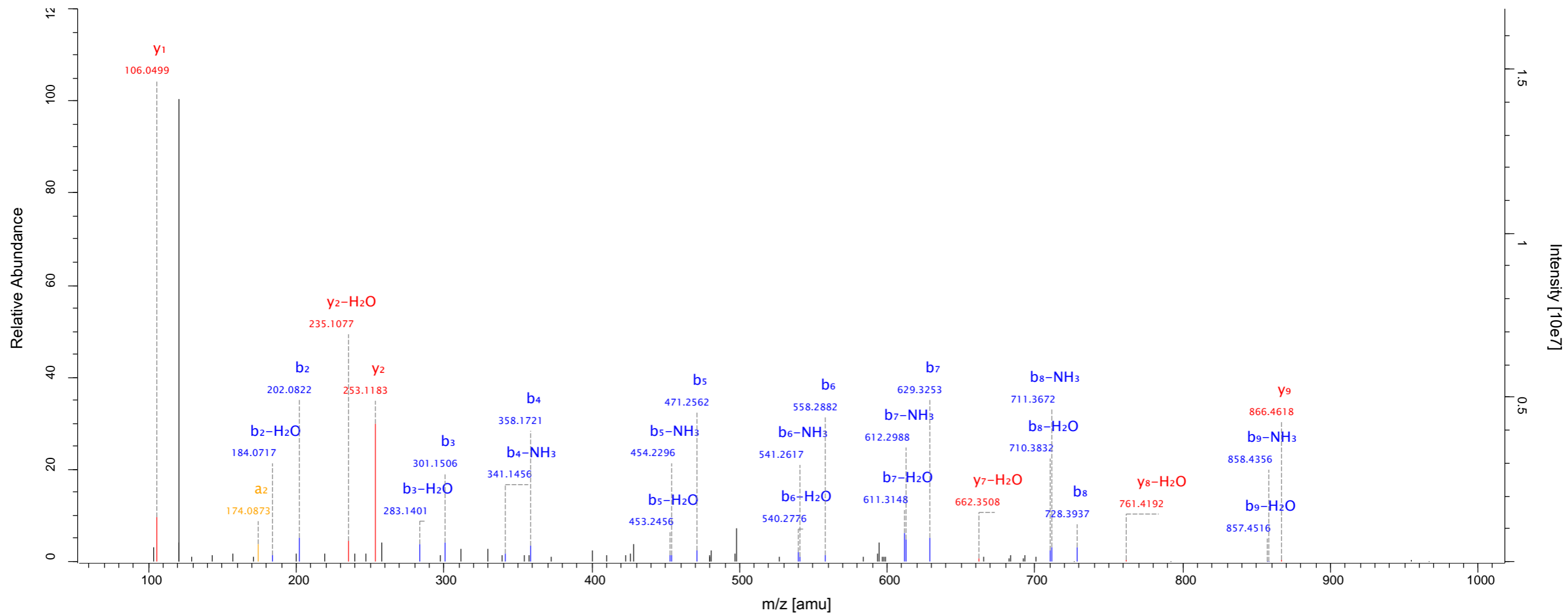
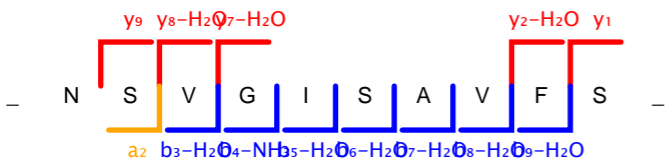
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 34502
Protein: ORFL118C - NC_006273.2[61896,62396]- <unknown description>
Peptide score: 115.82
Method: FTMS; HCD



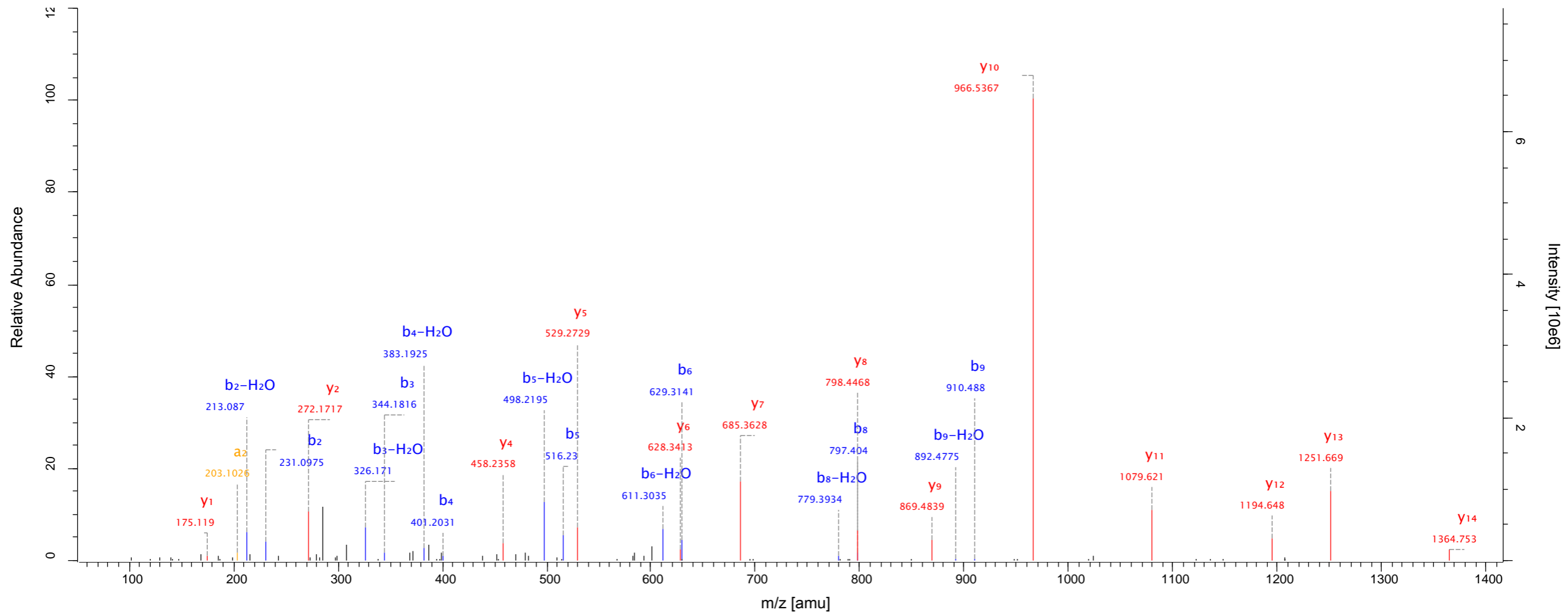
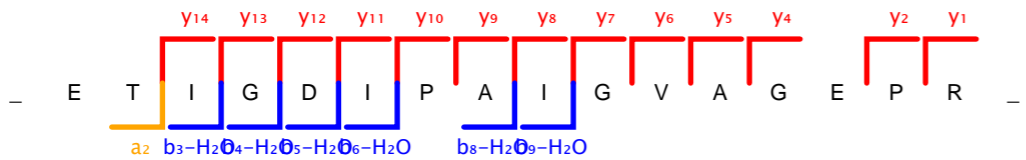
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 35767
 Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
 Peptide score: 127.22
 Method: FTMS; HCD



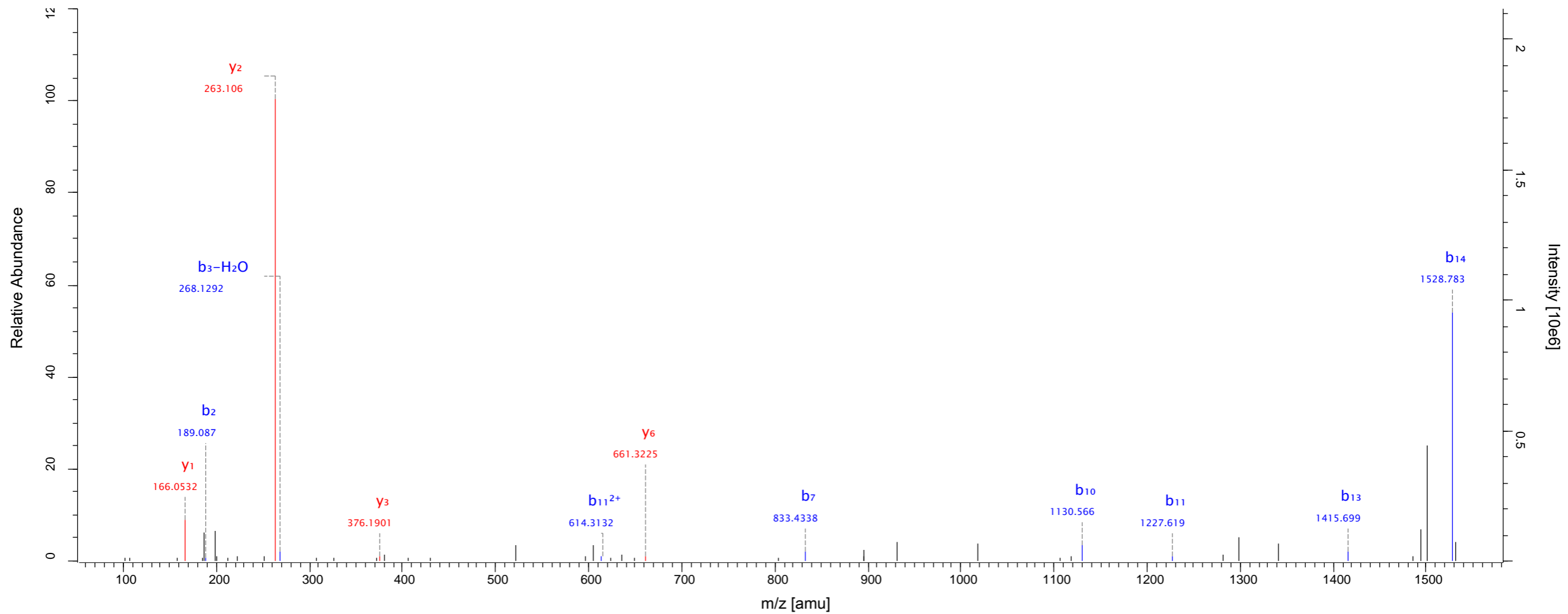
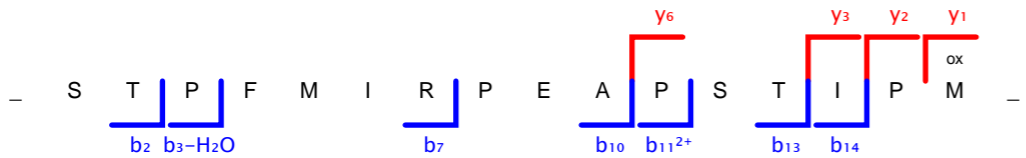
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_18
 Scan number: 39931
 Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>; ORFL27C.iORF1 - NC_006273.2[7912,8184]- <unknown description>
 Peptide score: 140.17
 Method: FTMS; HCD



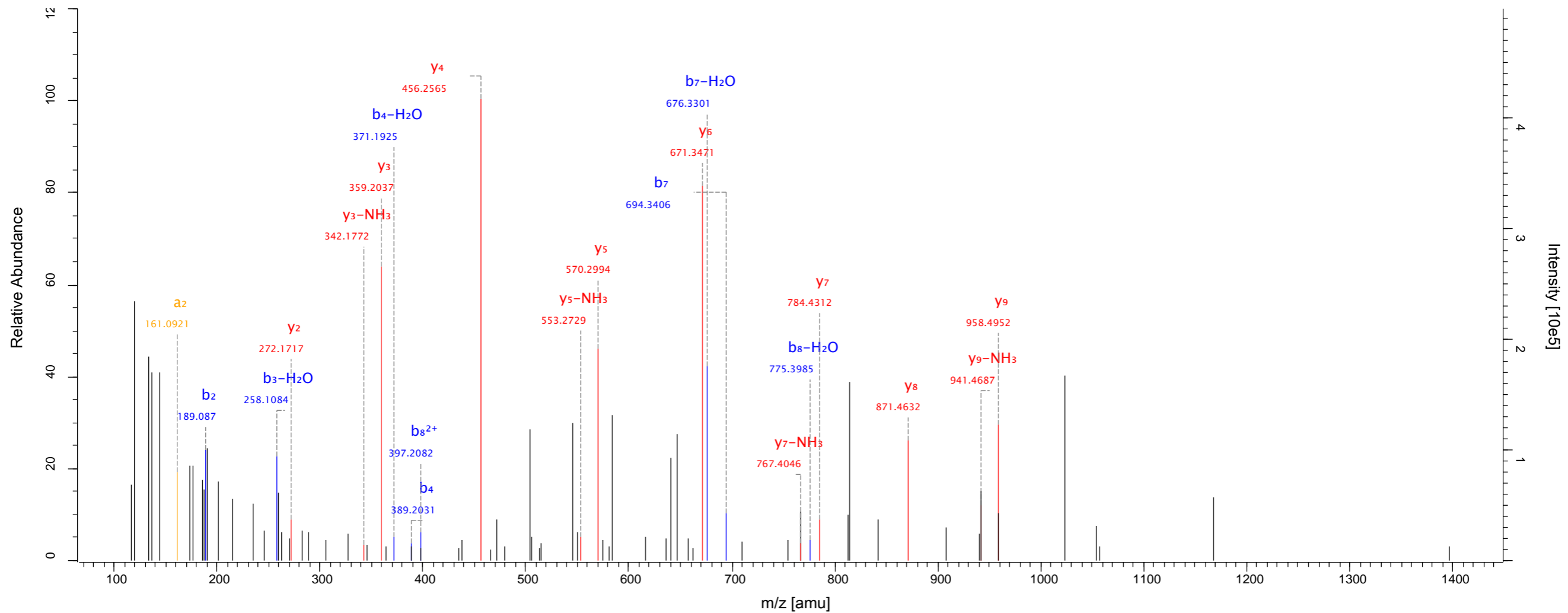
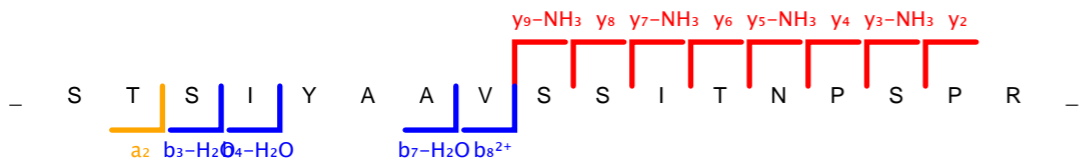
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 41493
Protein: ORFL73C - NC_006273.2[30581,30946]- <unknown description>
Peptide score: 170.56
Method: FTMS; HCD



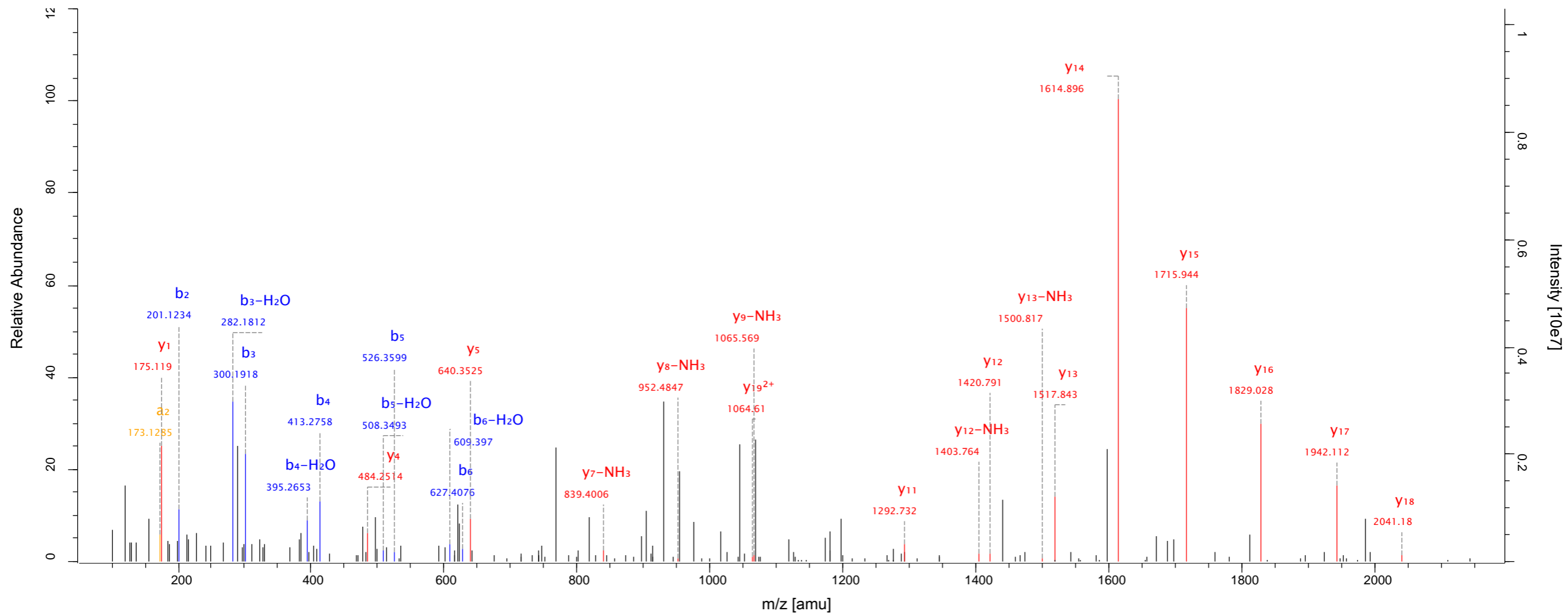
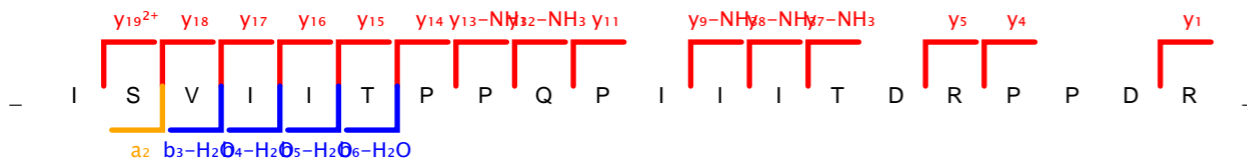
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_18
Scan number: 41766
Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
Peptide score: 54.6
Method: FTMS; HCD



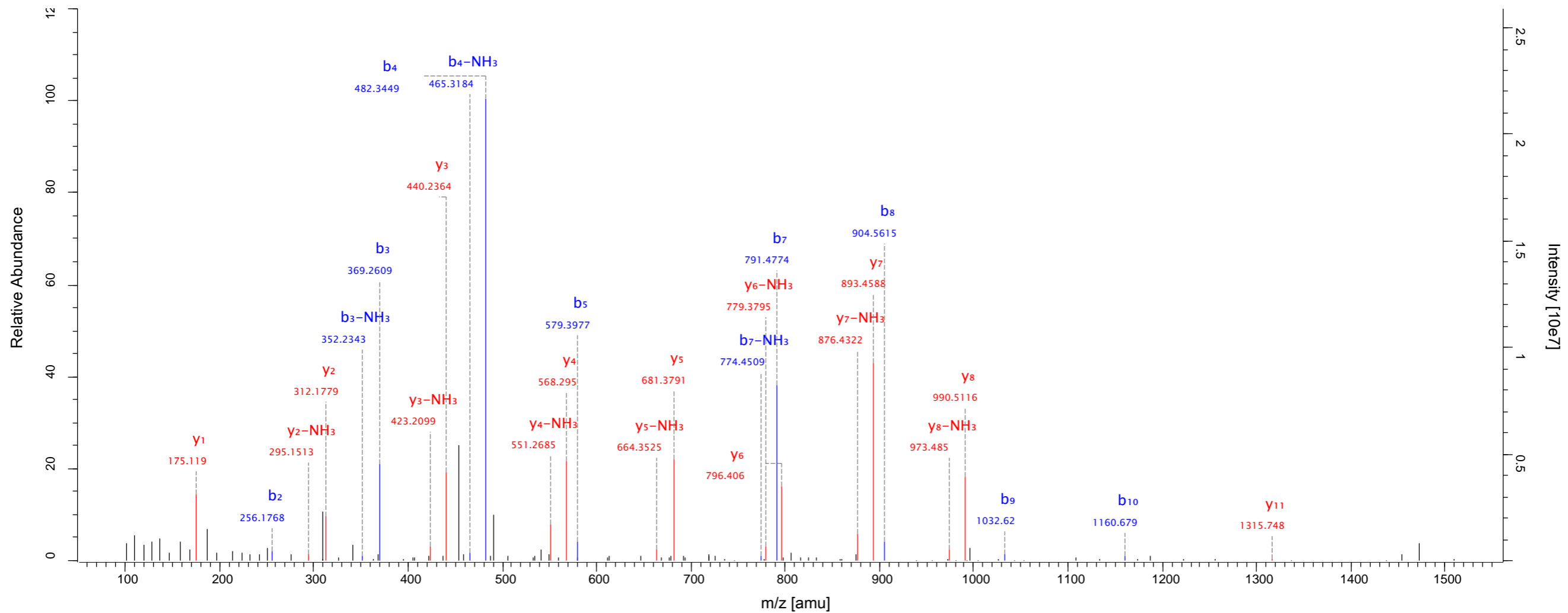
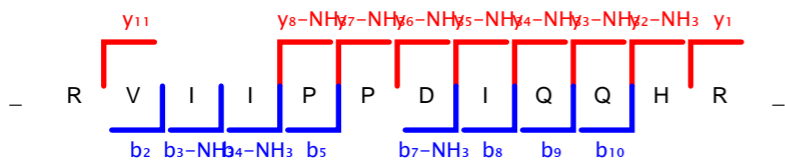
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 45522
 Protein: ORFL253W.IORF3 + NC_006273.2[162059,162286]+ <unknown description>
 Peptide score: 66.25
 Method: FTMS; HCD



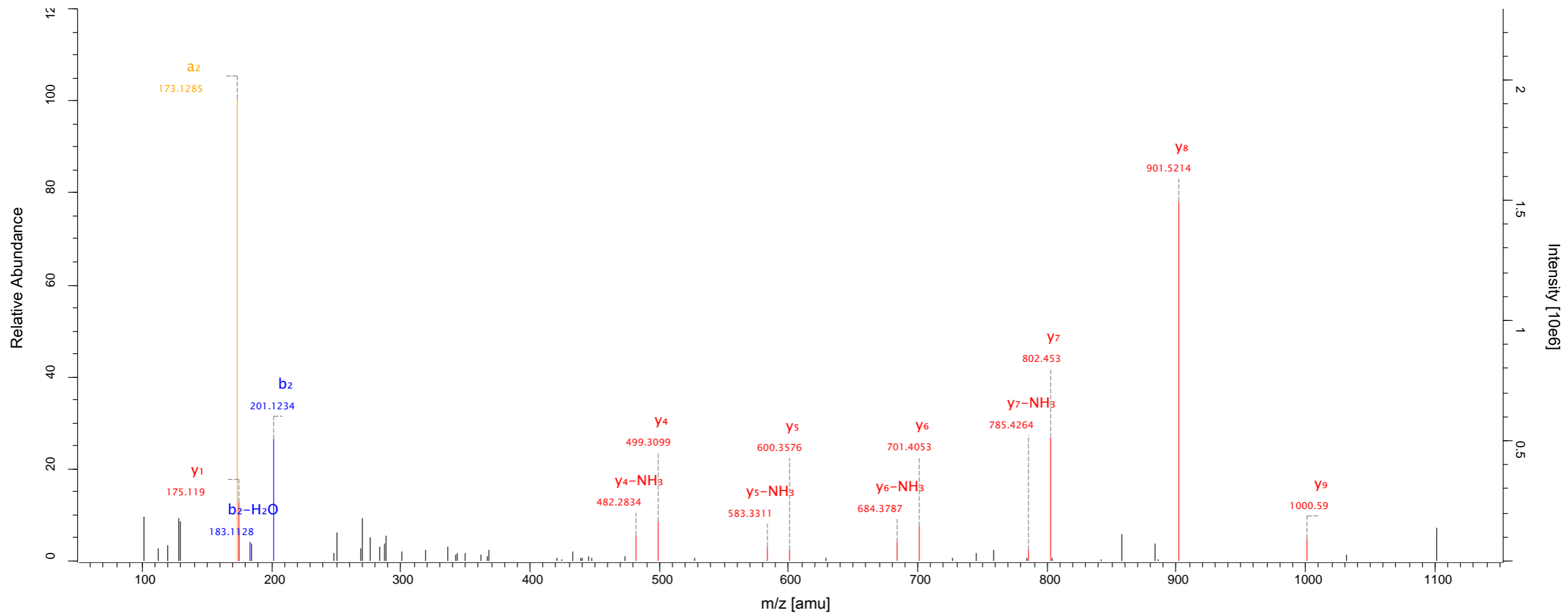
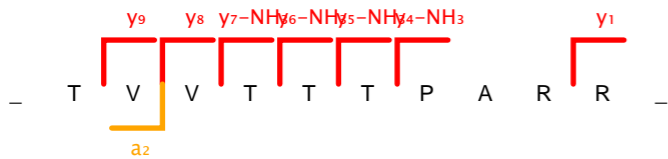
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 52334
 Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
 Peptide score: 94.88
 Method: FTMS; HCD



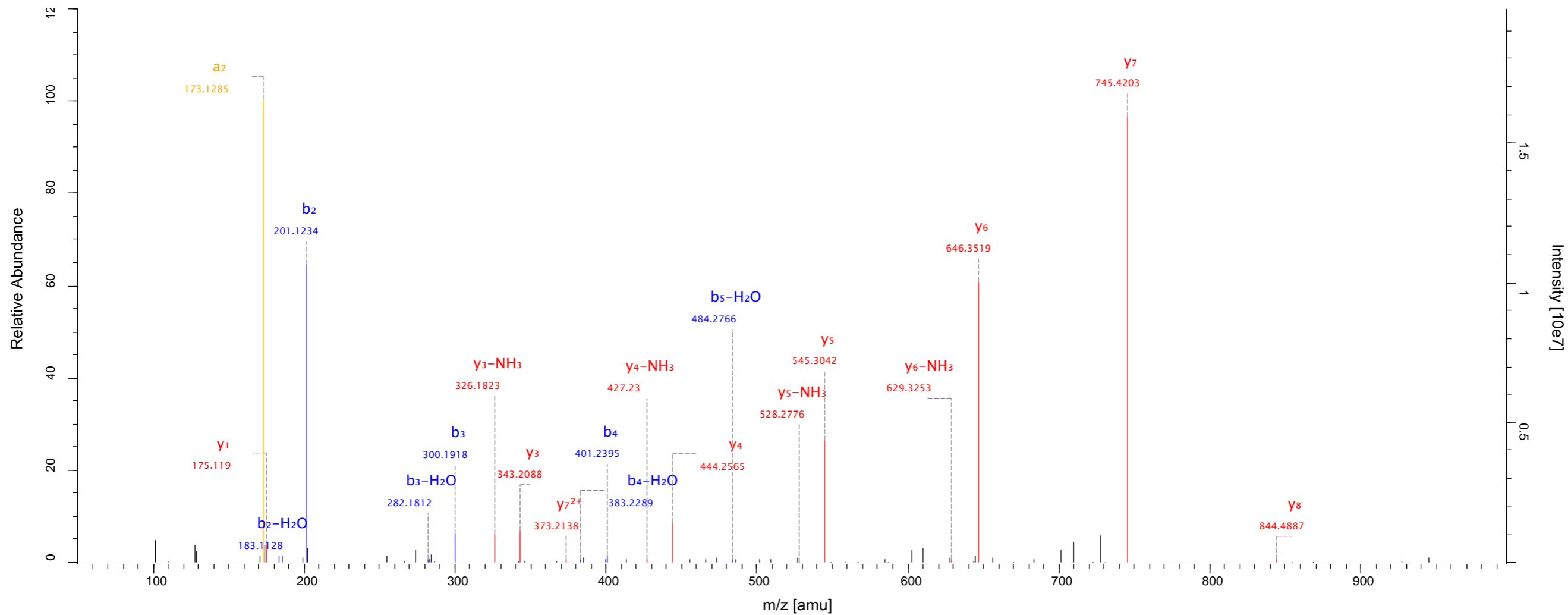
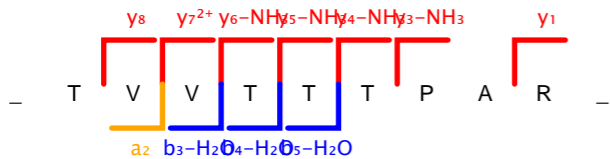
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
 Scan number: 52490
 Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
 Peptide score: 201.6
 Method: FTMS; HCD



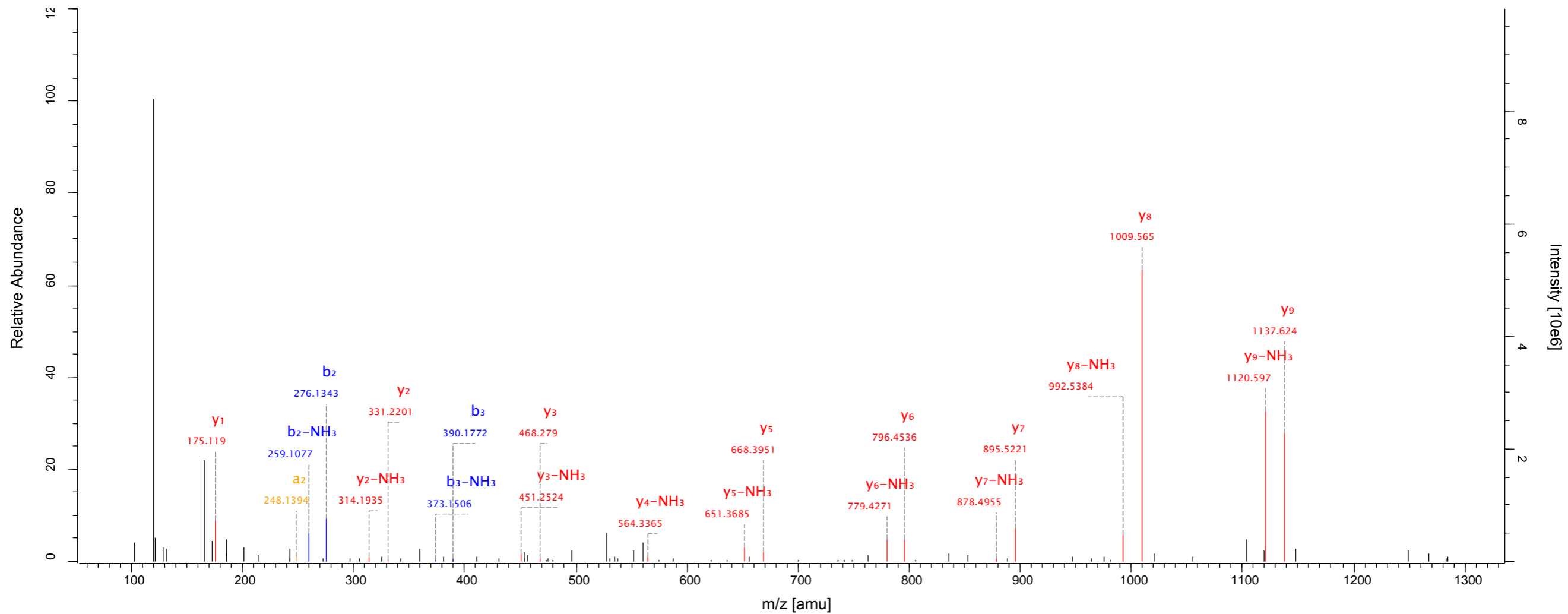
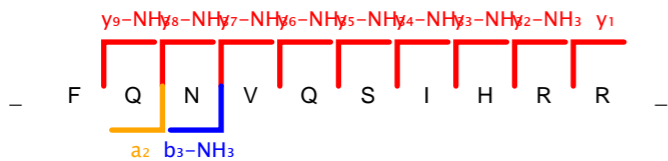
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_18
Scan number: 6478
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 118.26
Method: FTMS; HCD



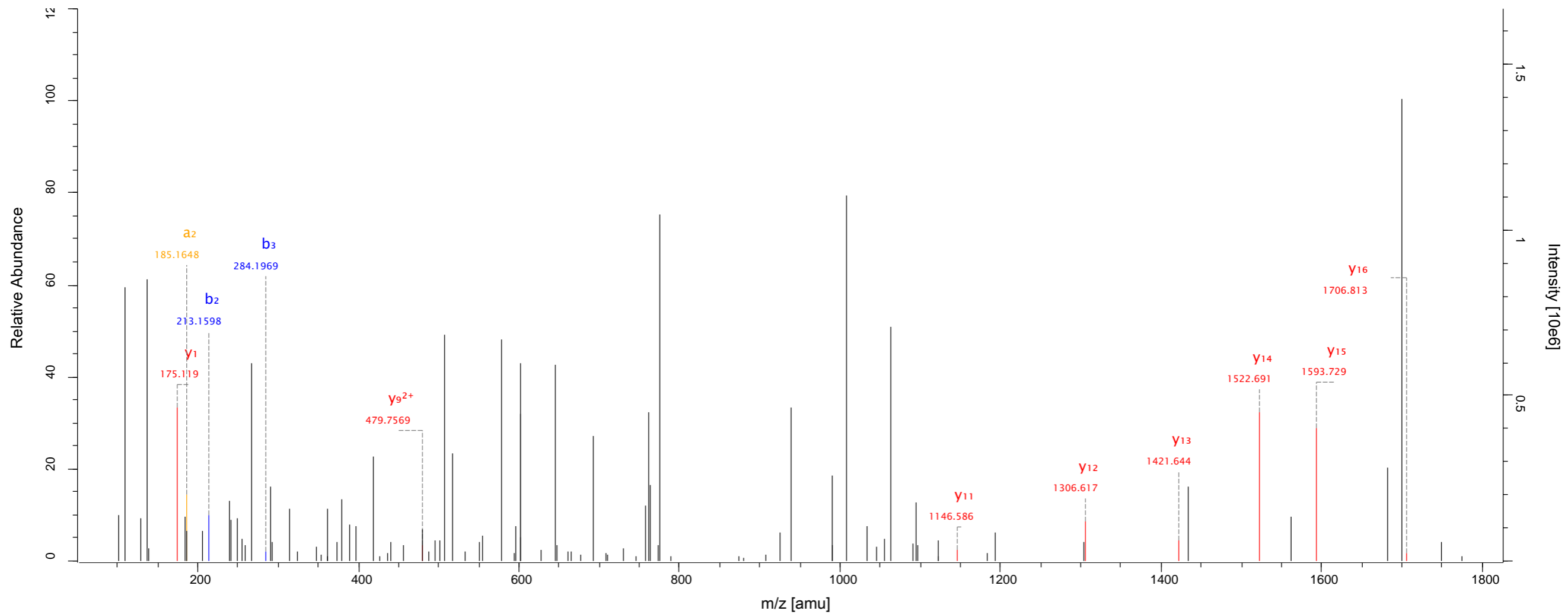
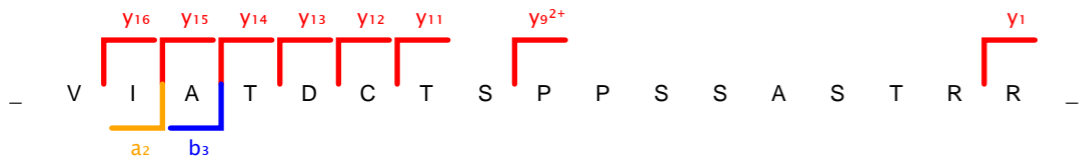
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 7919
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 127.46
Method: FTMS; HCD



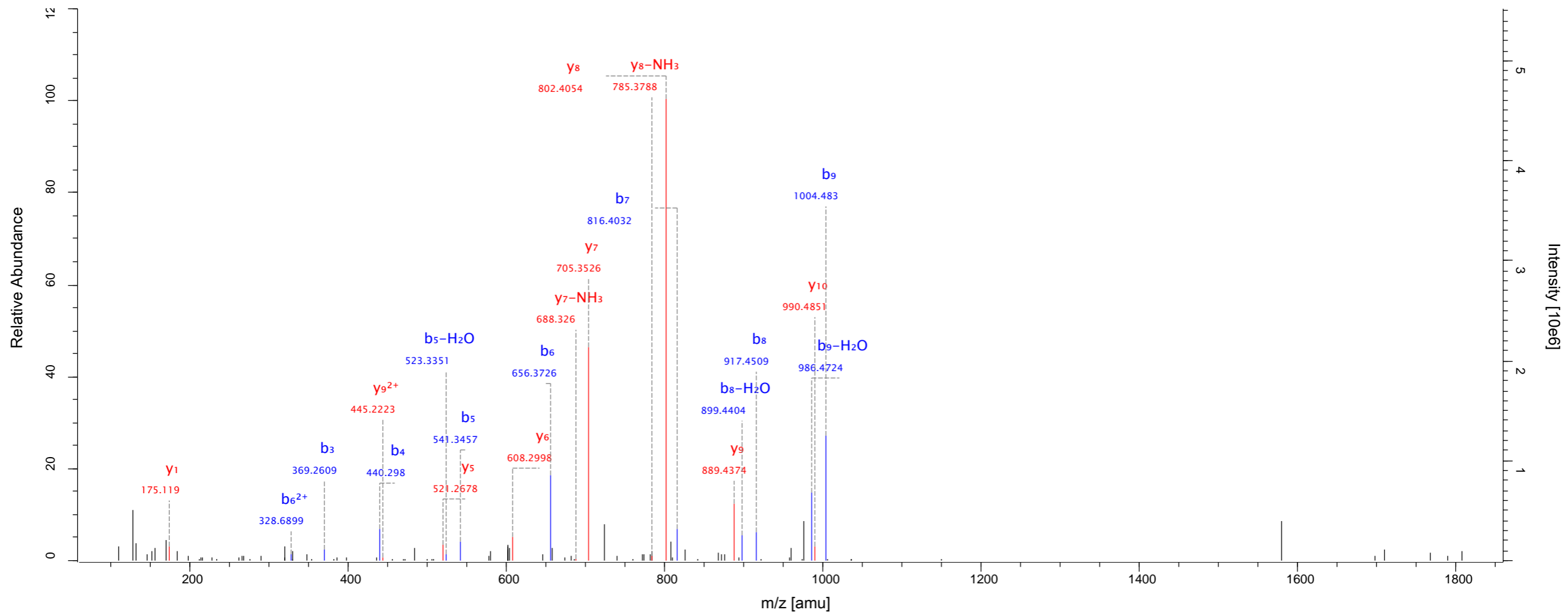
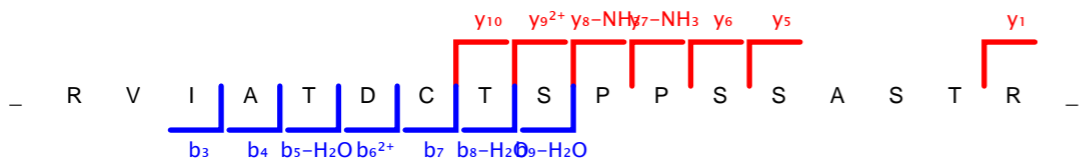
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_18
Scan number: 8693
Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
Peptide score: 124.92
Method: FTMS; HCD



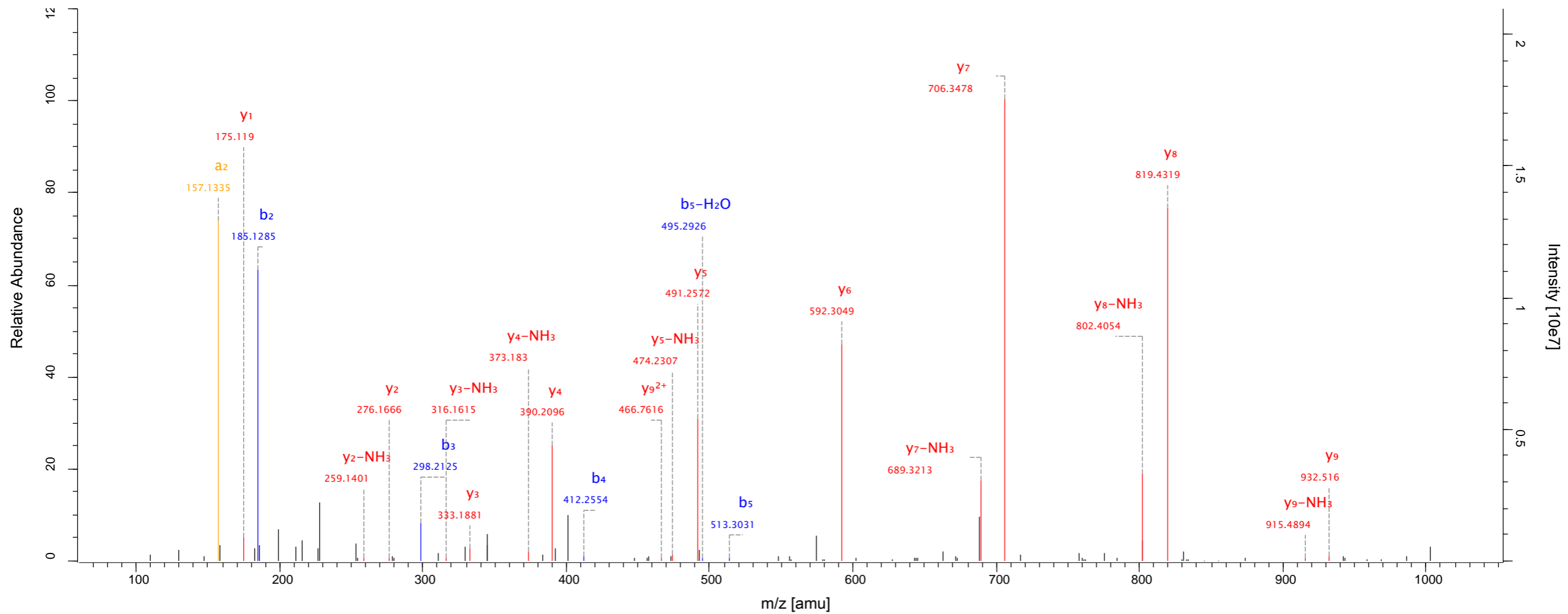
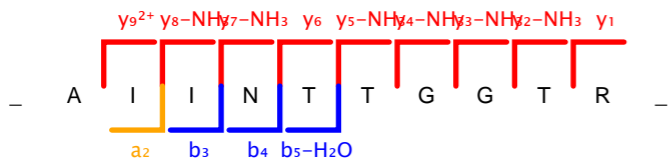
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 10470
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 31.61
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 11193
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 102.65
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
Scan number: 11897
Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
Peptide score: 144.5
Method: FTMS; HCD



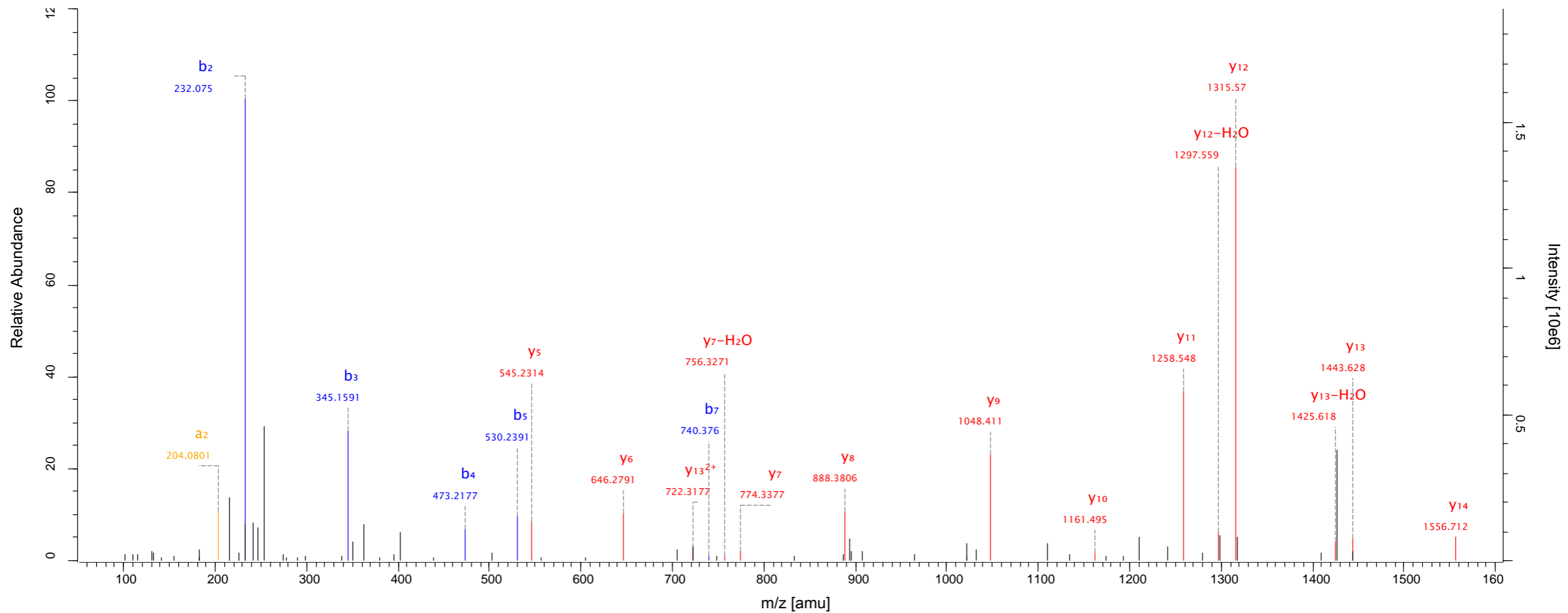
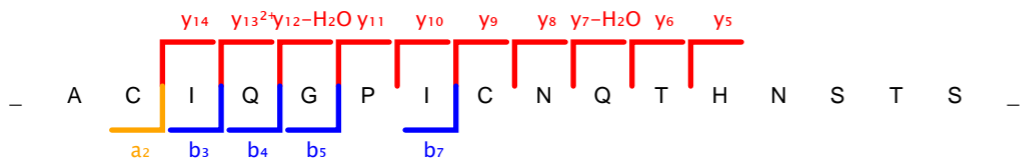
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19

Scan number: 13446

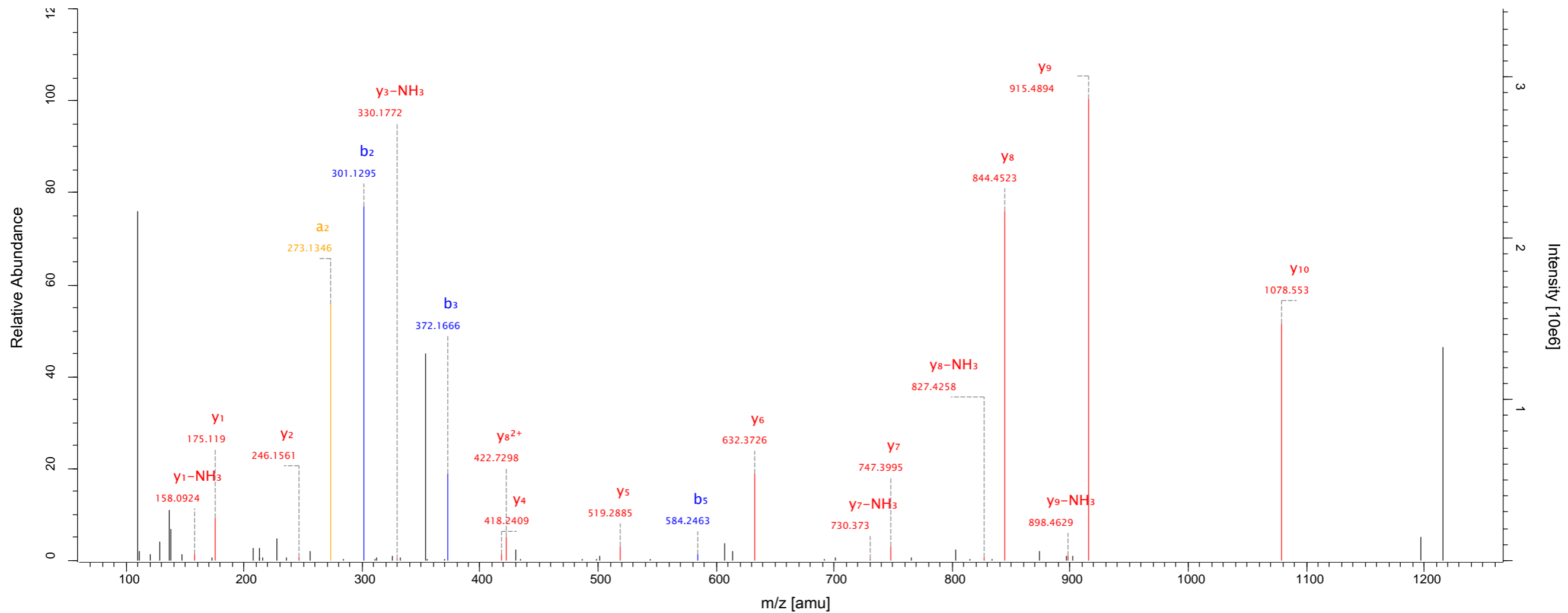
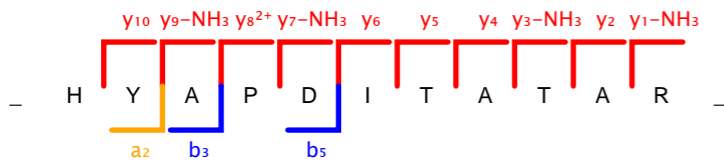
Protein: ORFL159W + NC_006273.2[101475,109052]+ <unknown description>; ORFL159W.iORF1 + NC_006273.2[101526,109052]+ <unknown description>; ORFL178W + NC_006273.2[108142,109052]+ <unknown description>; ORFL163W + NC_006273.2[101475,109052]+ <unknown description>;

Peptide score: 125.61

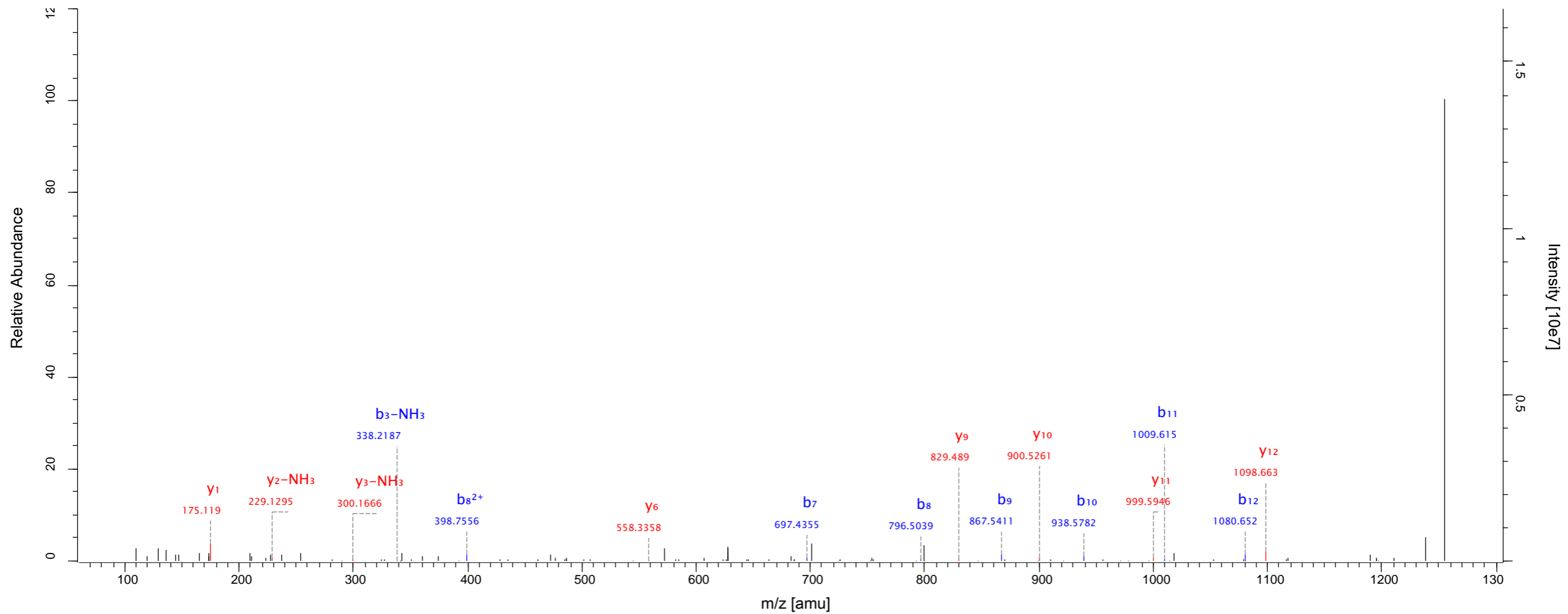
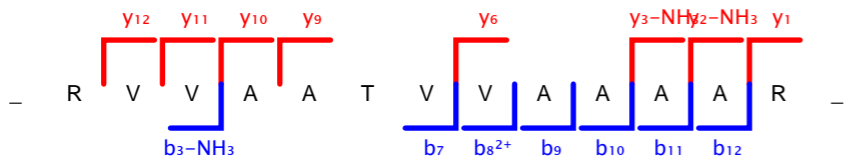
Method: FTMS; HCD



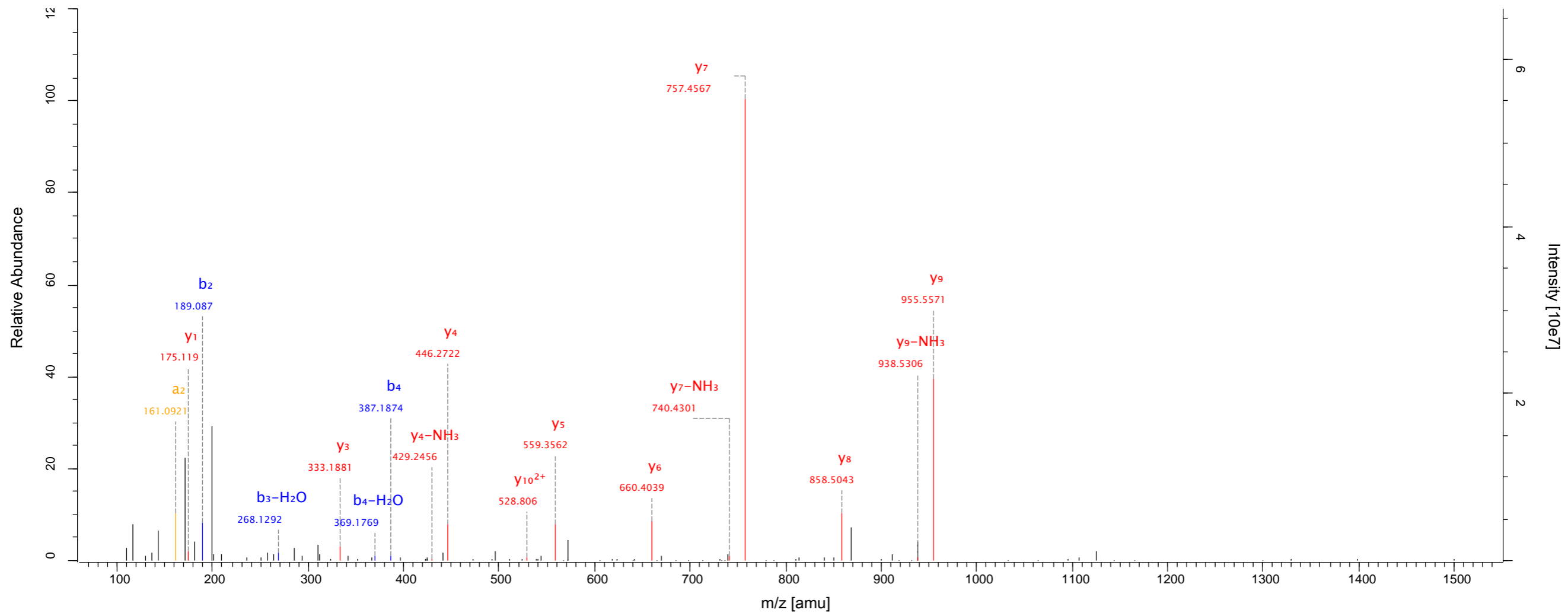
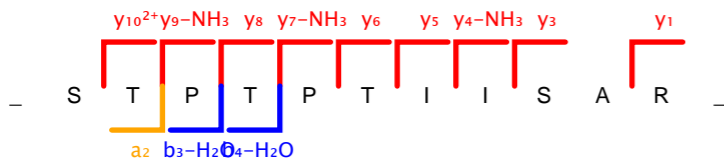
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 13486
Protein: ORFL190C - NC_006273.2[111965,112384]- <unknown description>
Peptide score: 107.79
Method: FTMS; HCD



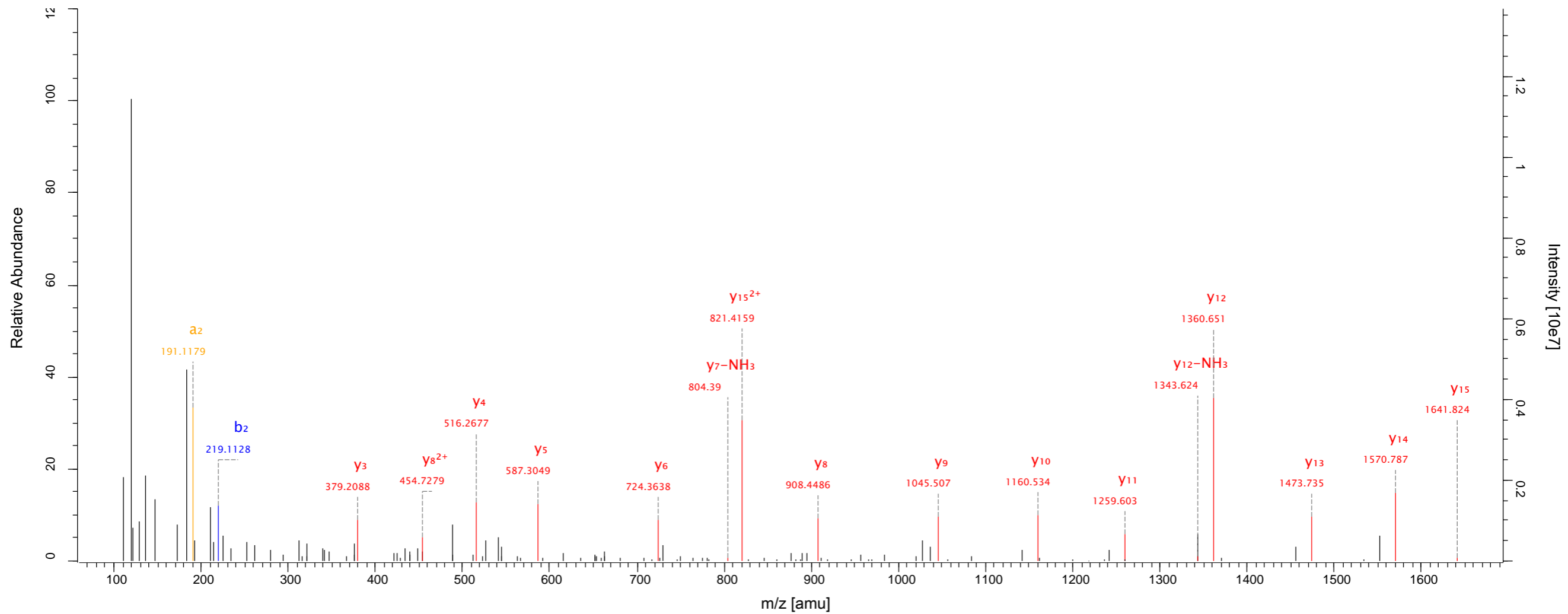
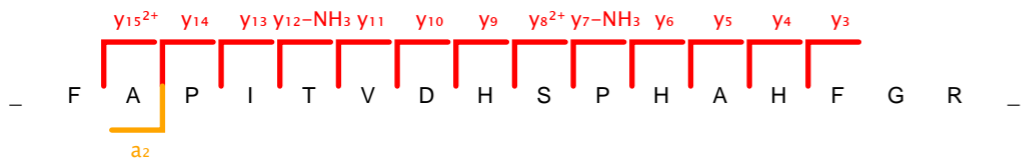
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 18343
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 61.21
Method: FTMS; HCD



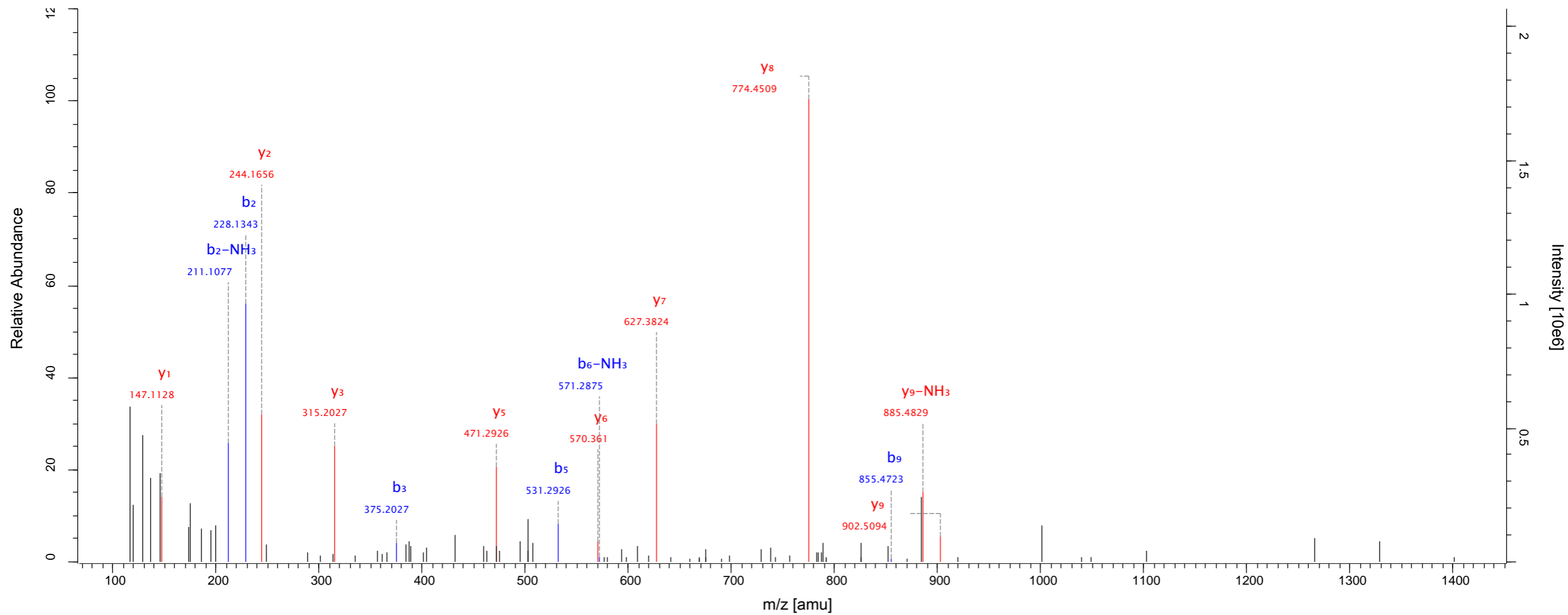
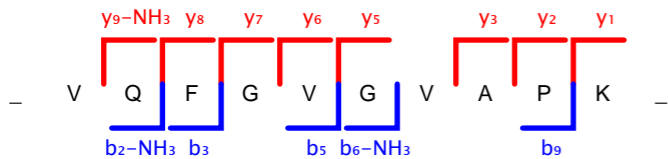
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
Scan number: 19685
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 85.35
Method: FTMS; HCD



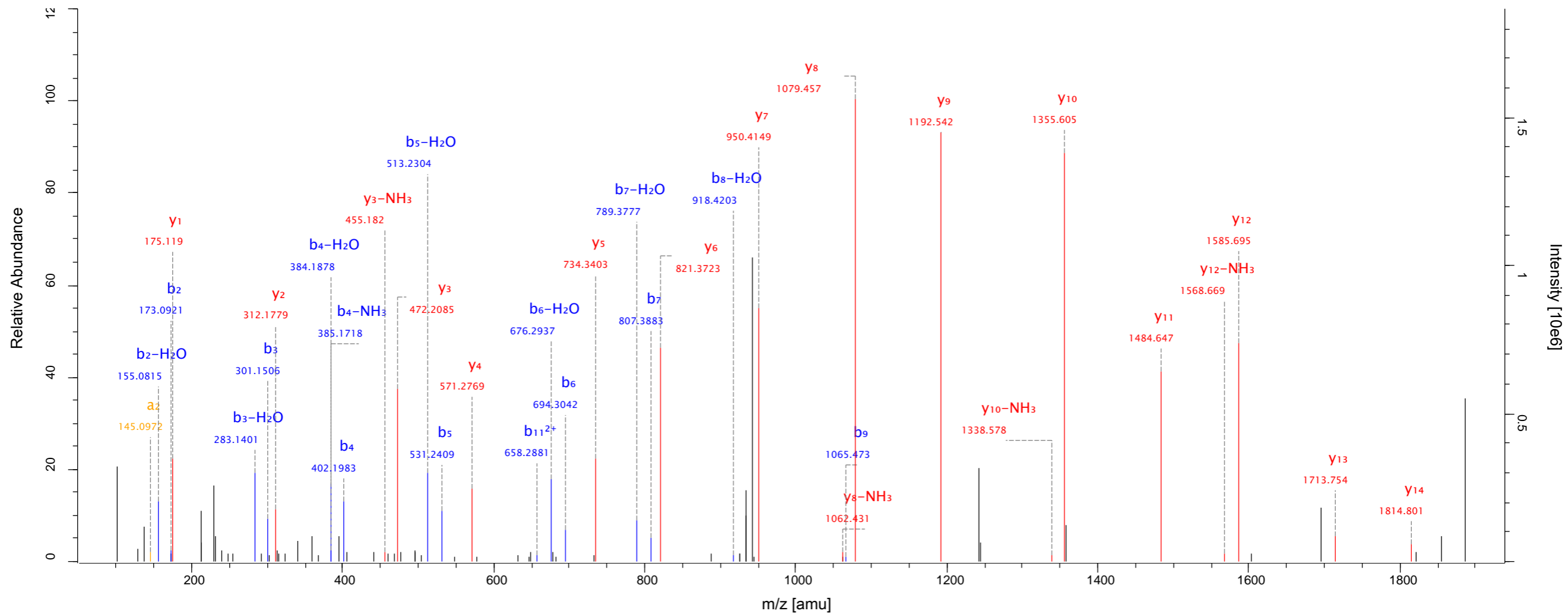
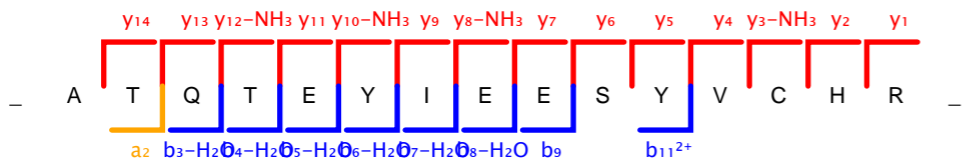
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
Scan number: 20000
Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
Peptide score: 125.61
Method: FTMS; HCD



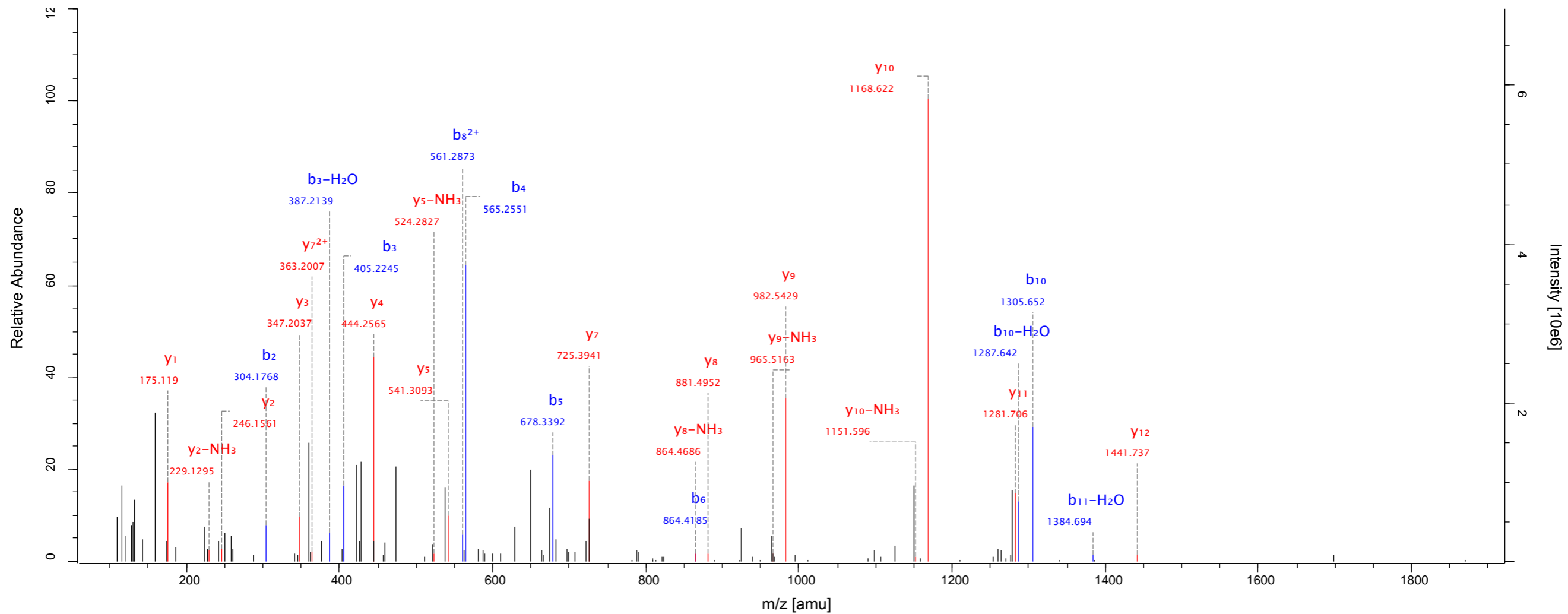
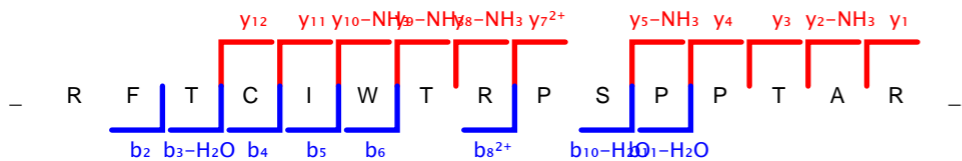
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
Scan number: 22030
Protein: ORFL118C - NC_006273.2[61896,62396]- <unknown description>
Peptide score: 91.58
Method: FTMS; HCD



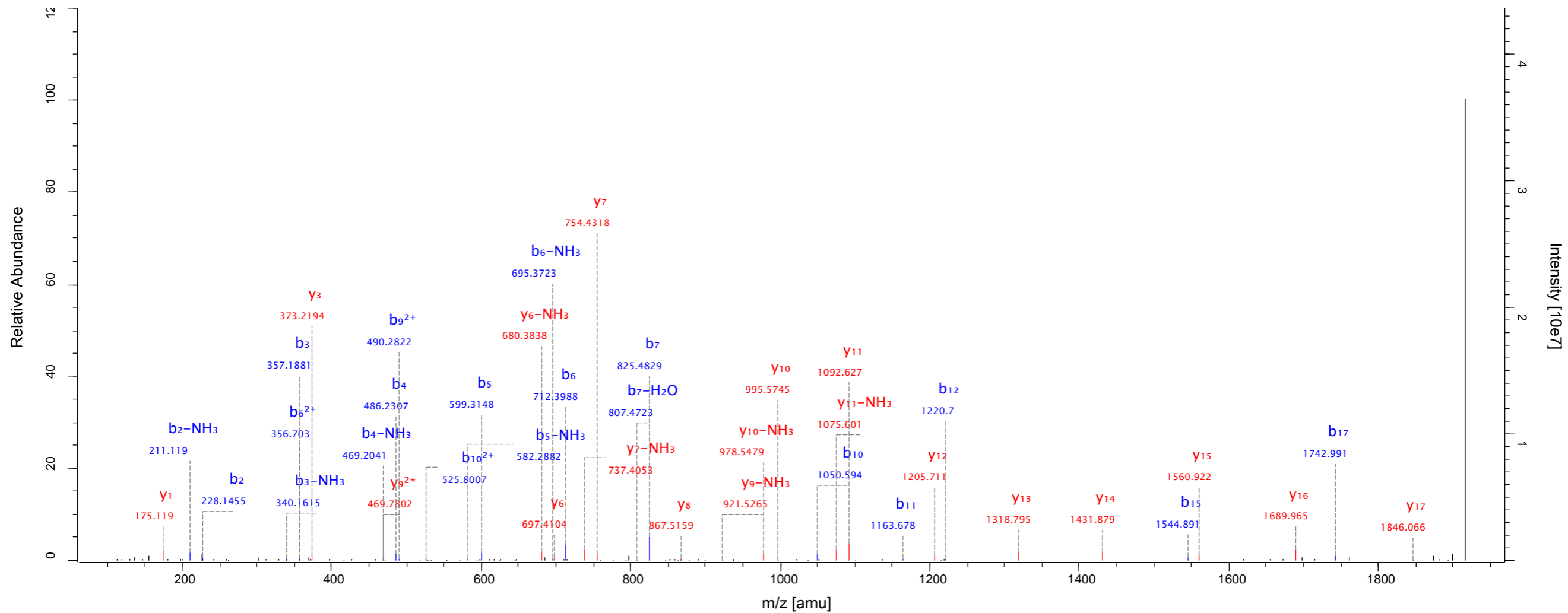
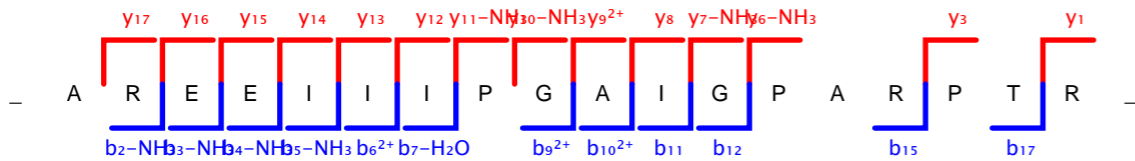
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
 Scan number: 22404
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 231.26
 Method: FTMS; HCD



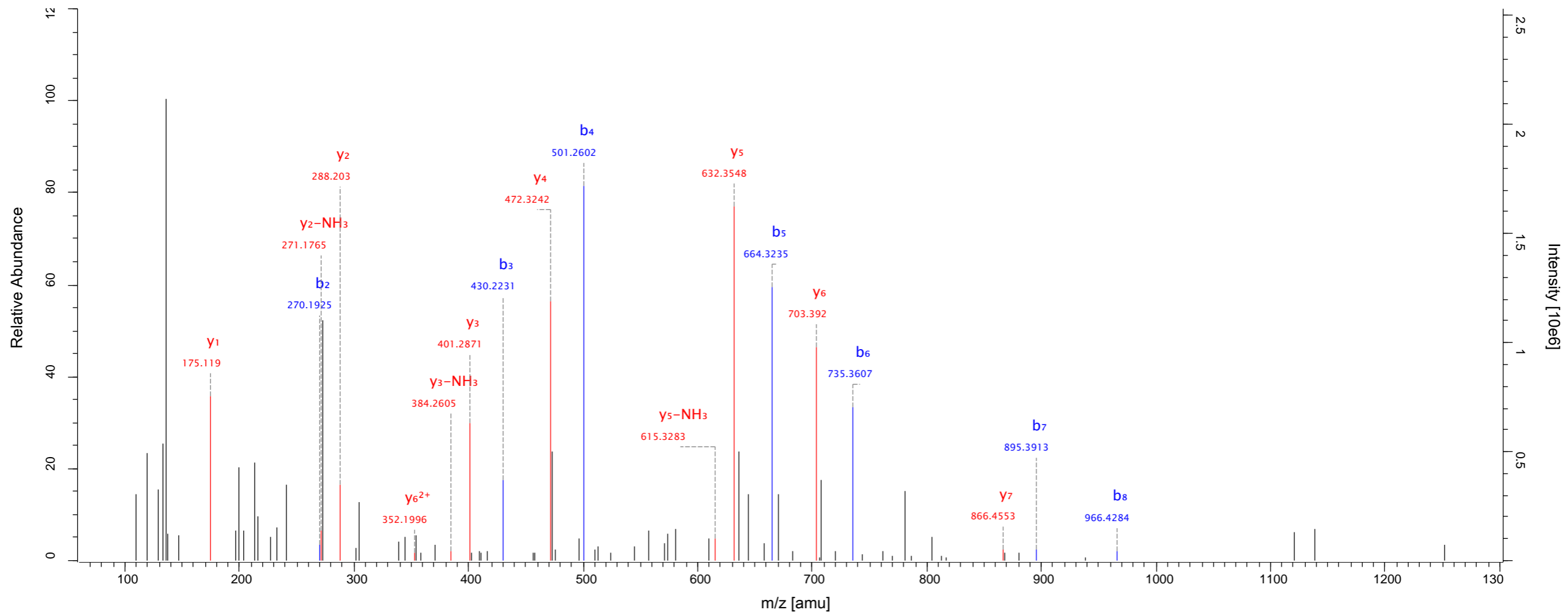
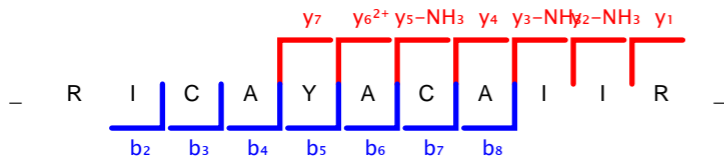
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 24618
 Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
 Peptide score: 119.96
 Method: FTMS; HCD



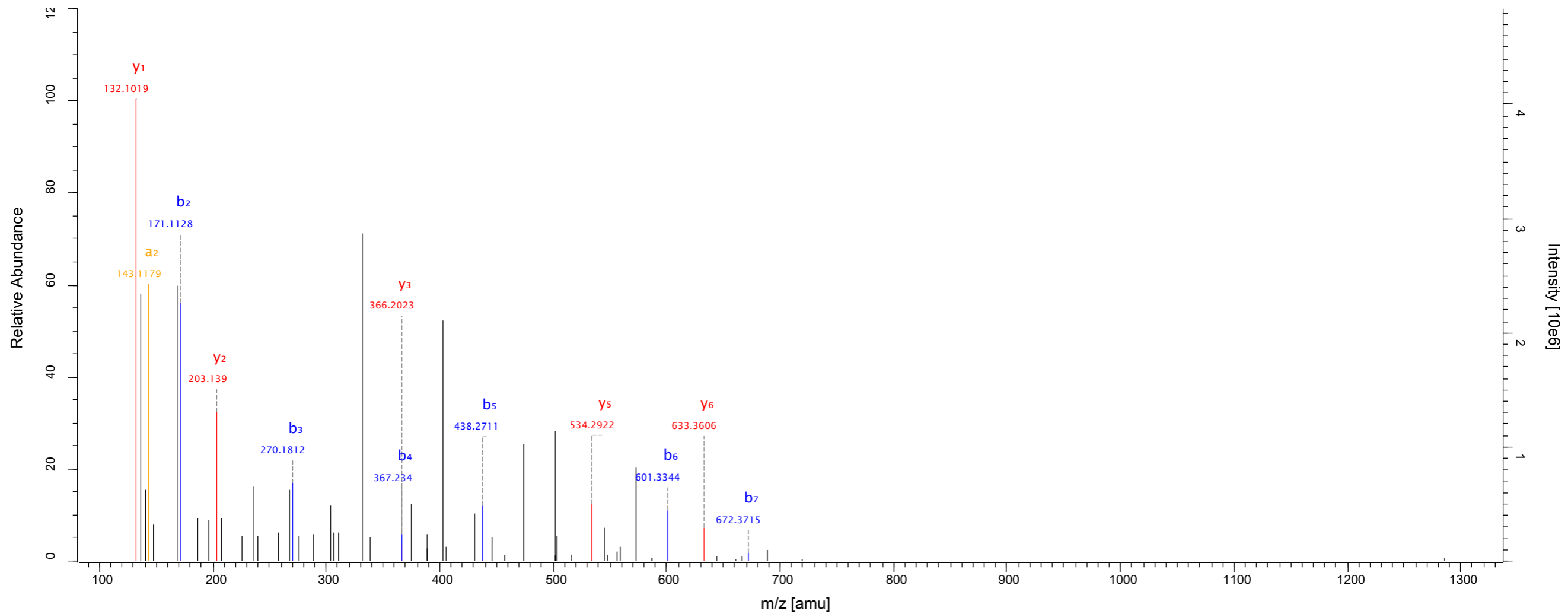
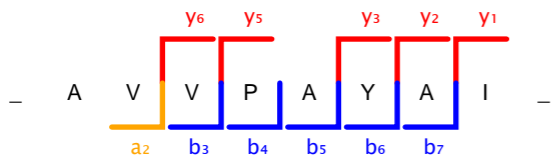
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 29780
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 210.09
 Method: FTMS; HCD



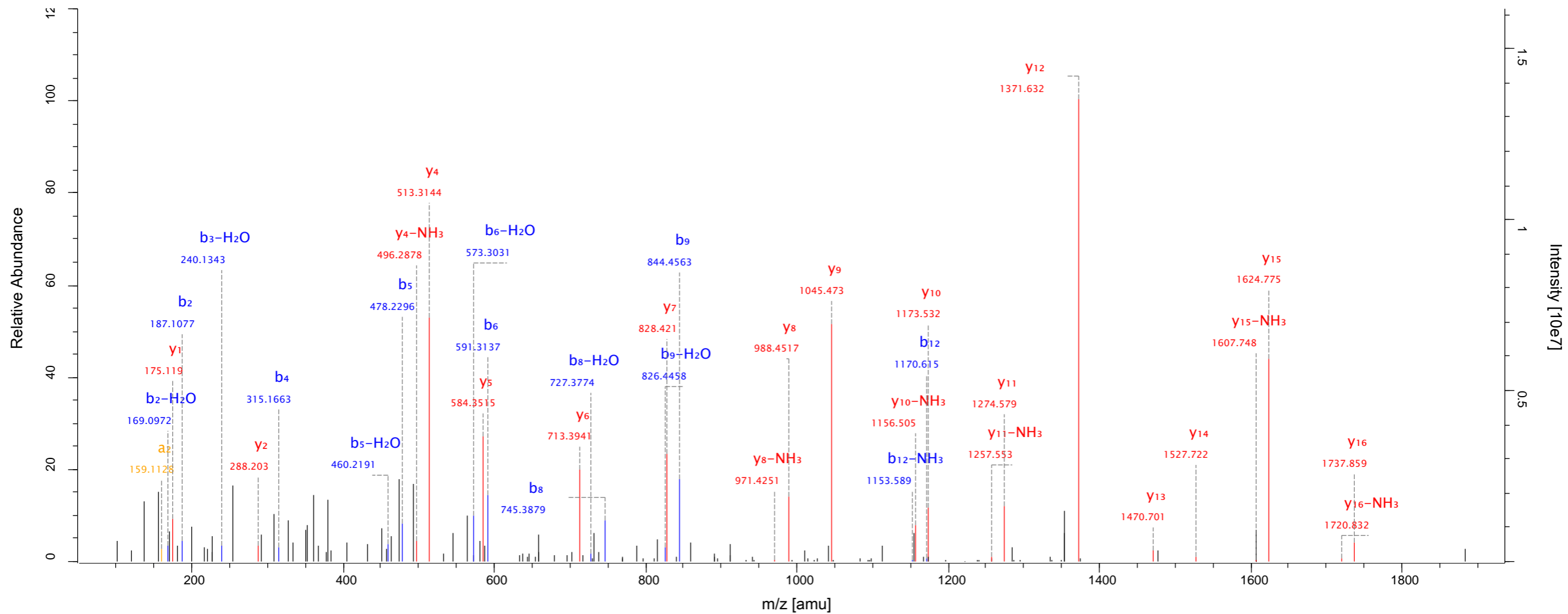
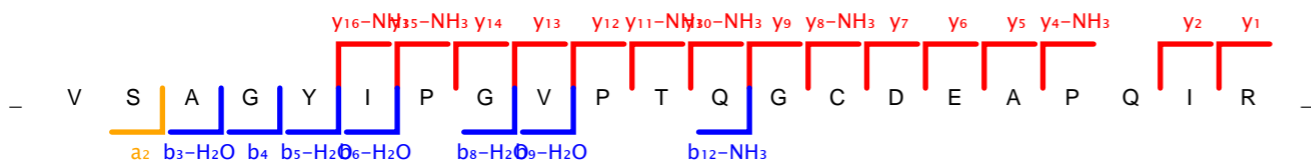
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 30102
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 90.69
Method: FTMS; HCD



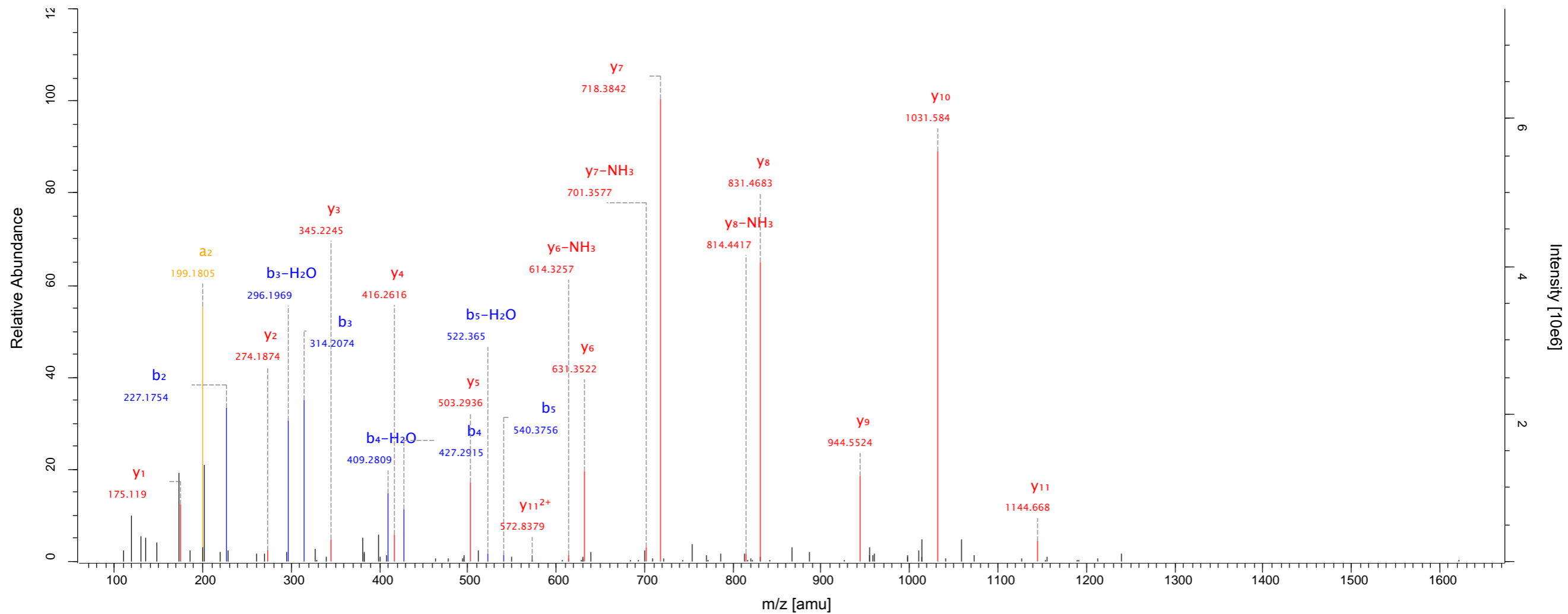
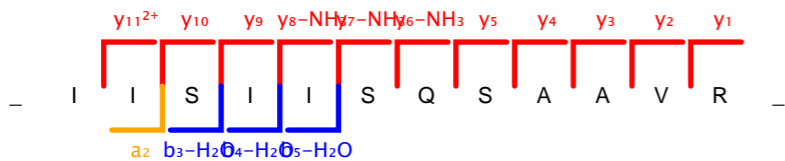
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 35002
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 99.51
Method: FTMS; HCD



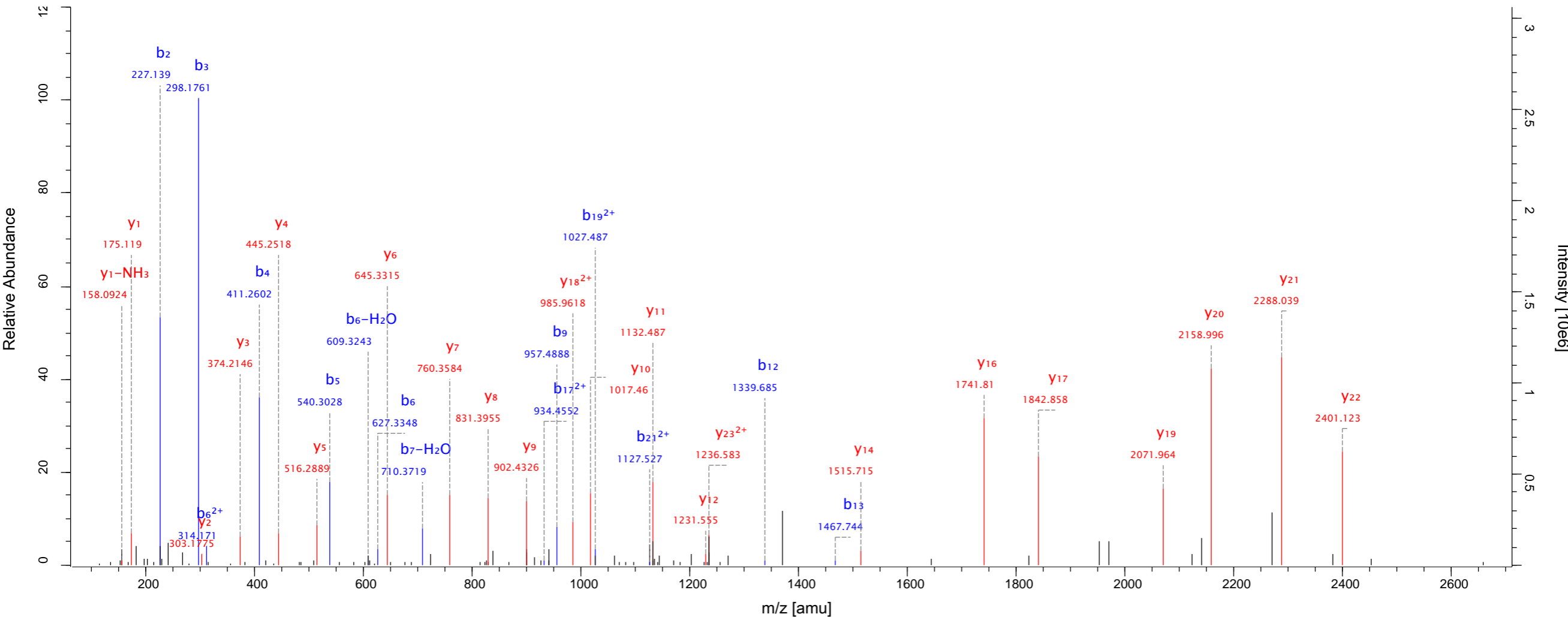
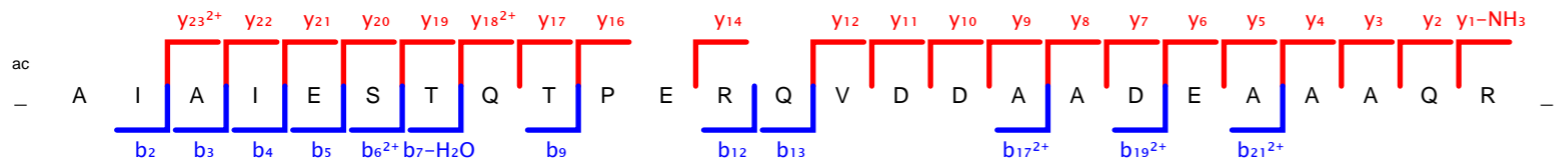
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 35617
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 132.25
 Method: FTMS; HCD



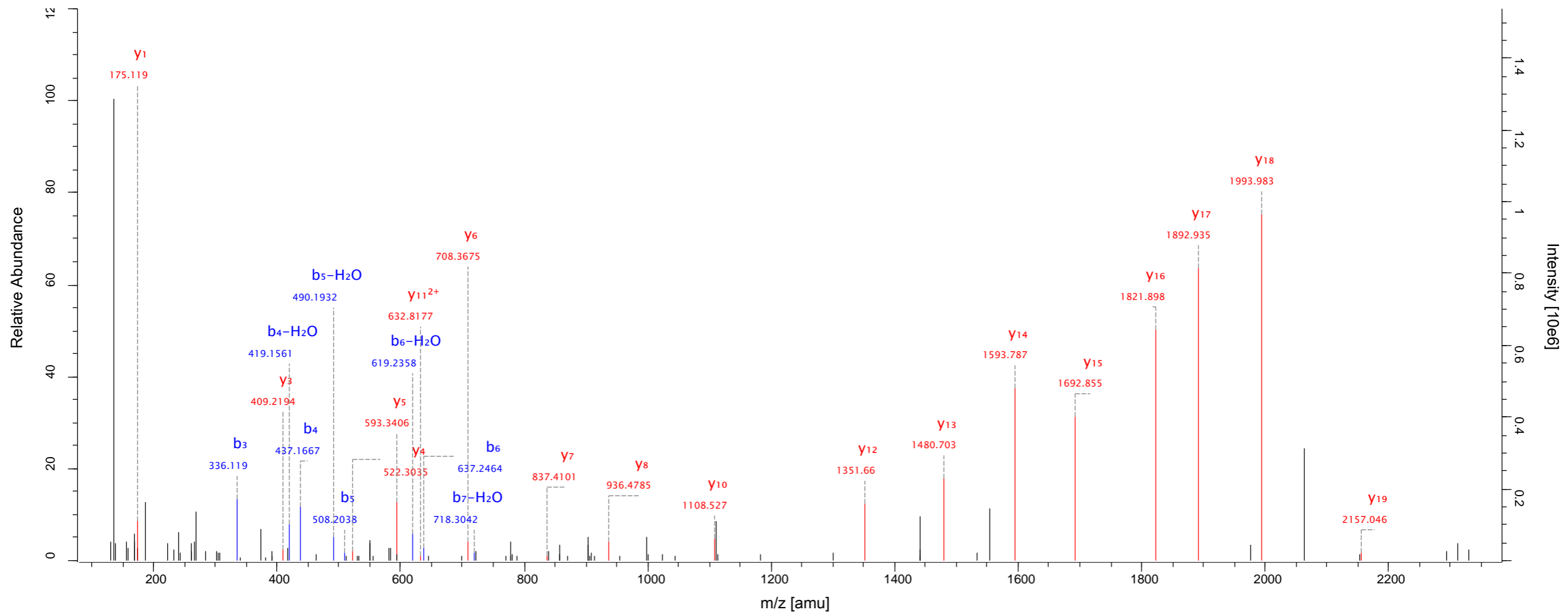
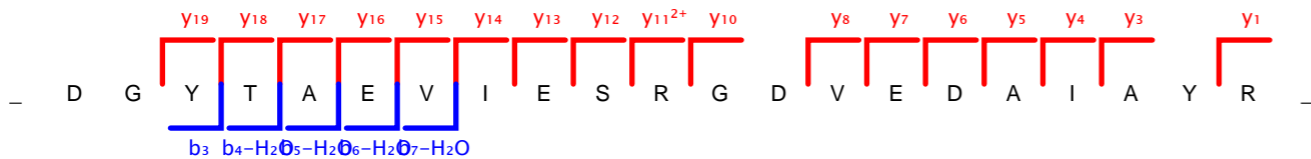
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
Scan number: 36733
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 177.57
Method: FTMS; HCD



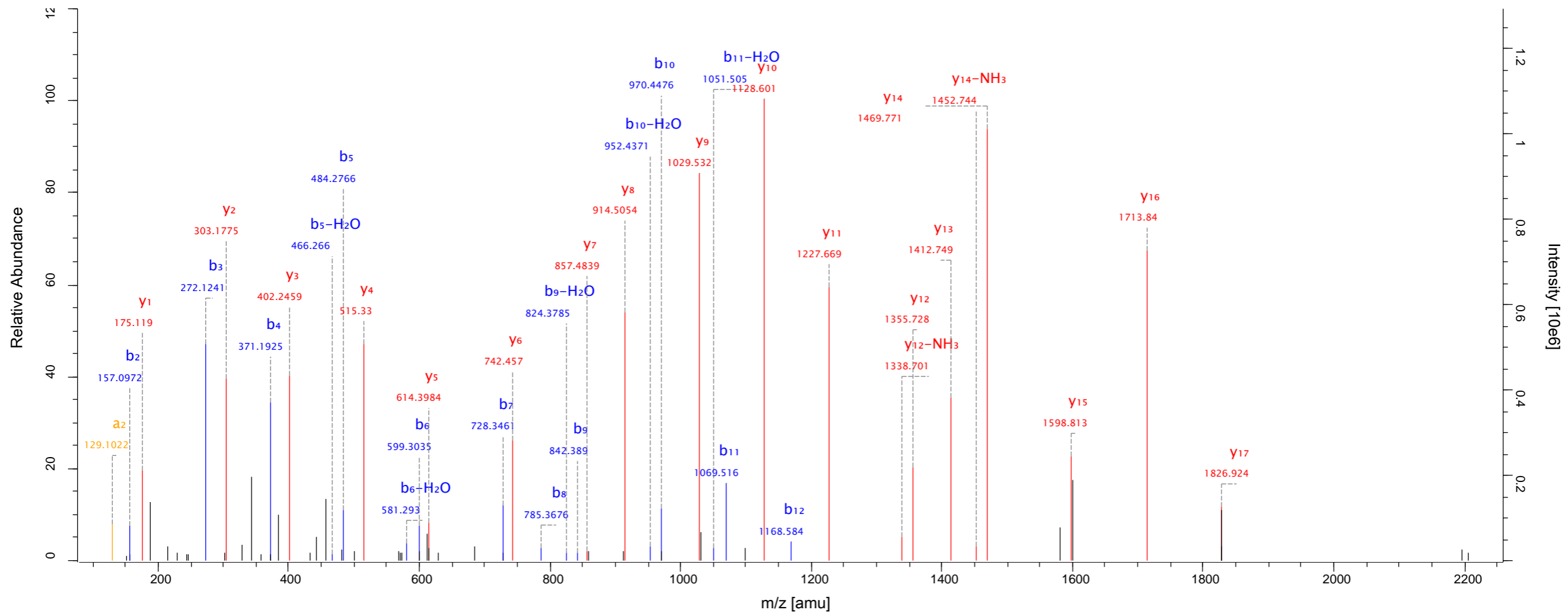
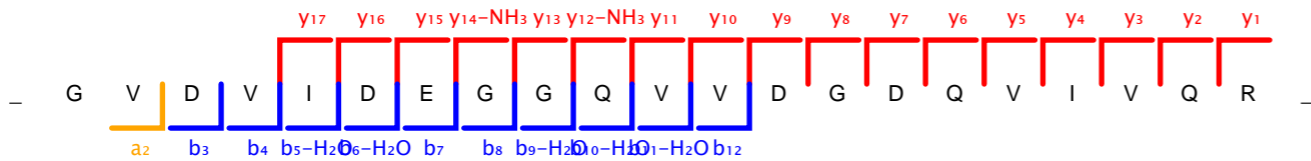
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 37966
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 218.76
 Method: FTMS; HCD



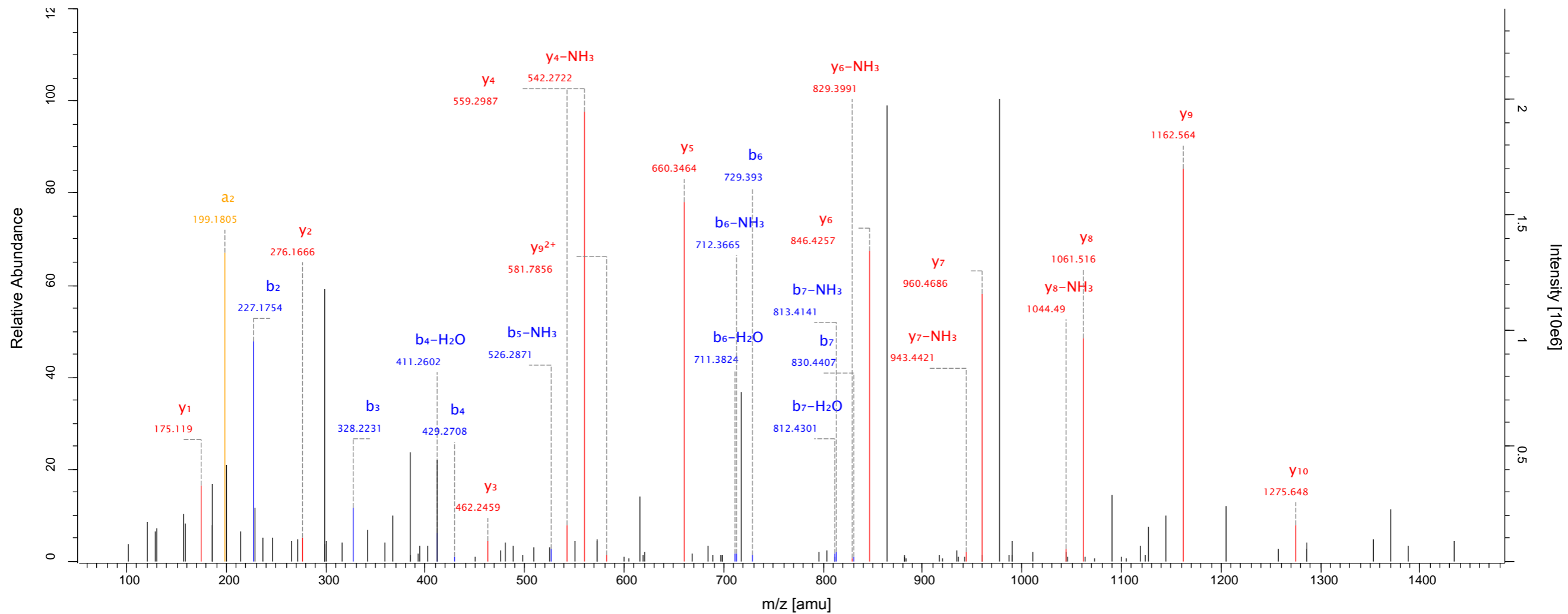
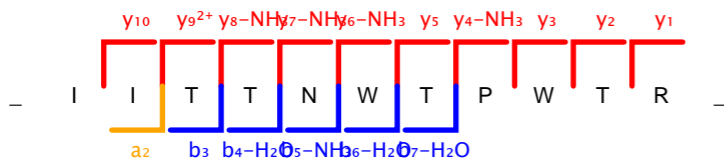
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 39235
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 115.15
 Method: FTMS; HCD



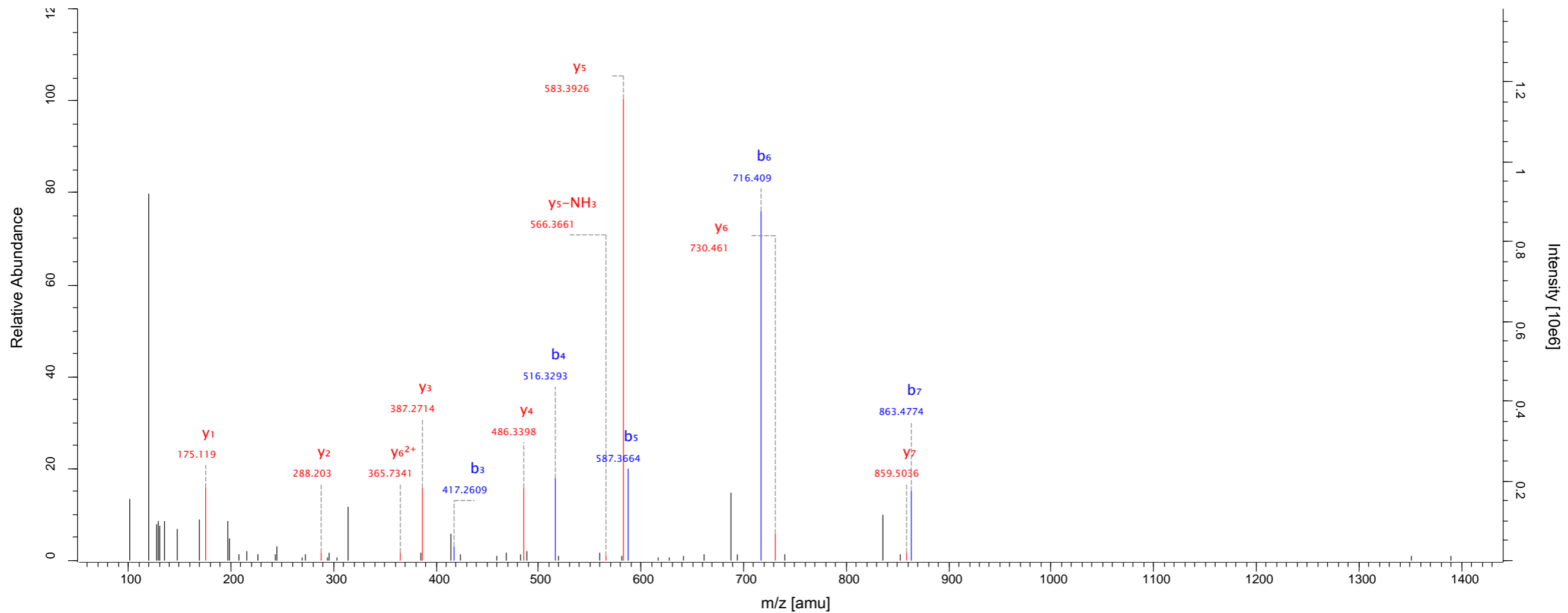
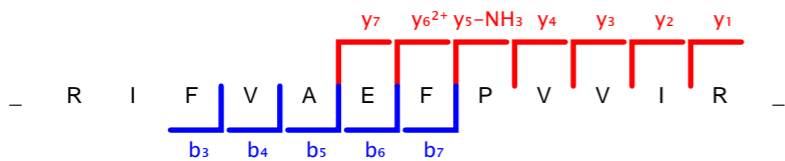
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 39968
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 222.7
 Method: FTMS; HCD



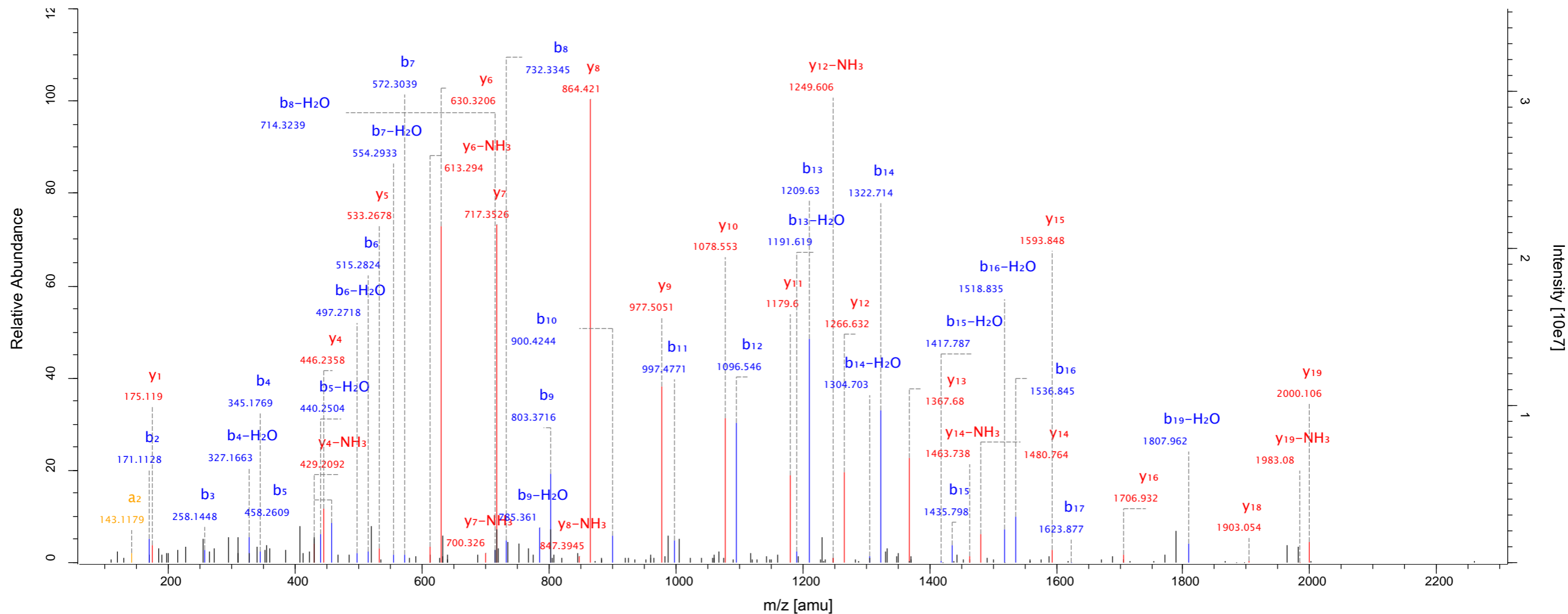
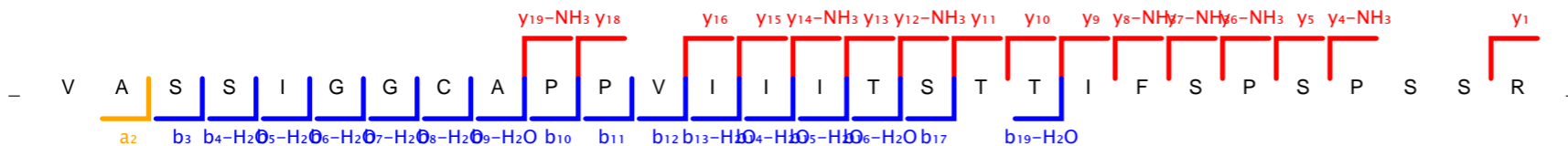
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 44064
 Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
 Peptide score: 150.1
 Method: FTMS; HCD



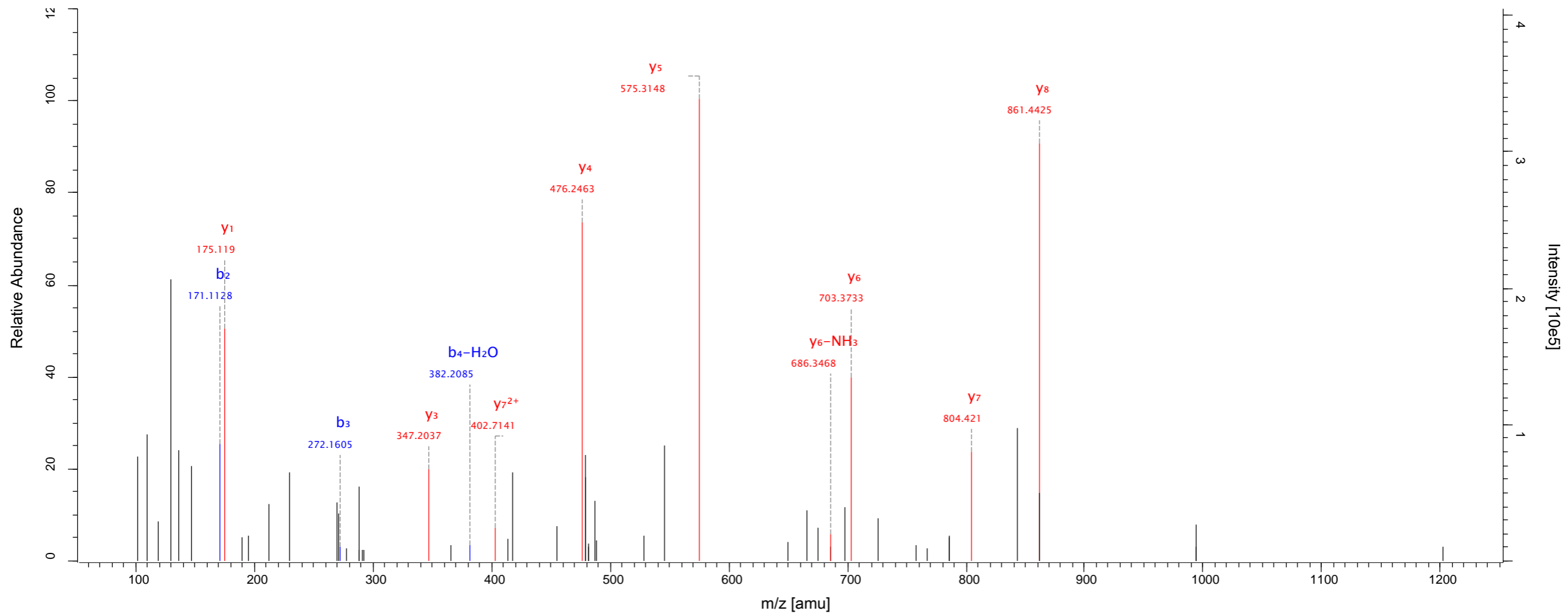
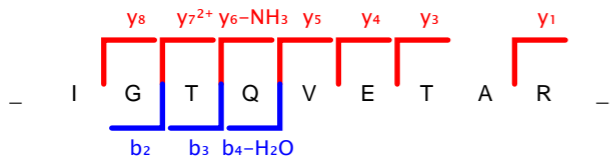
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
Scan number: 48526
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 89.4
Method: FTMS; HCD



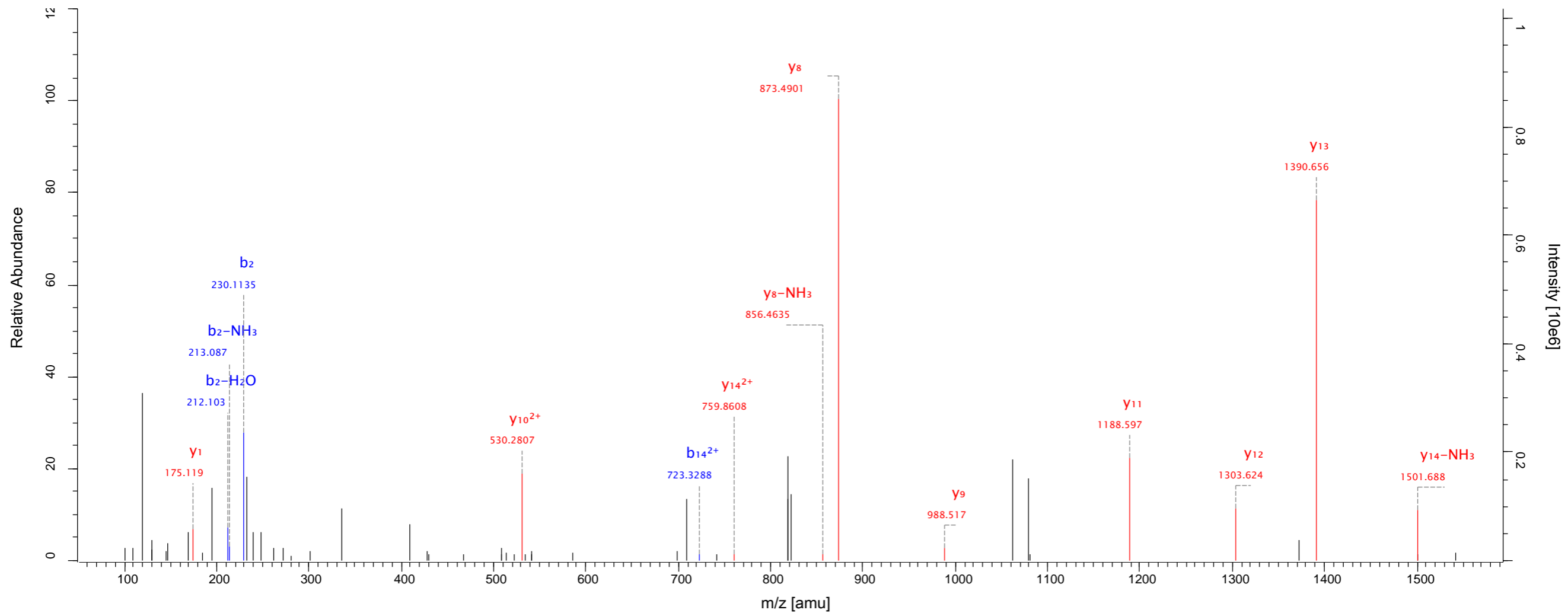
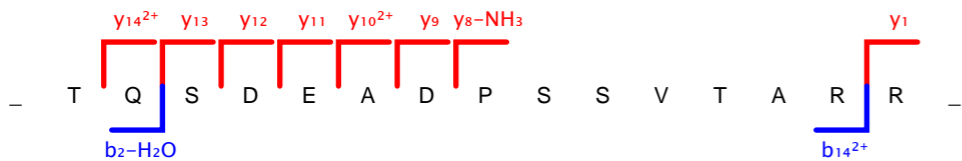
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
 Scan number: 51974
 Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
 Peptide score: 220.88
 Method: FTMS; HCD



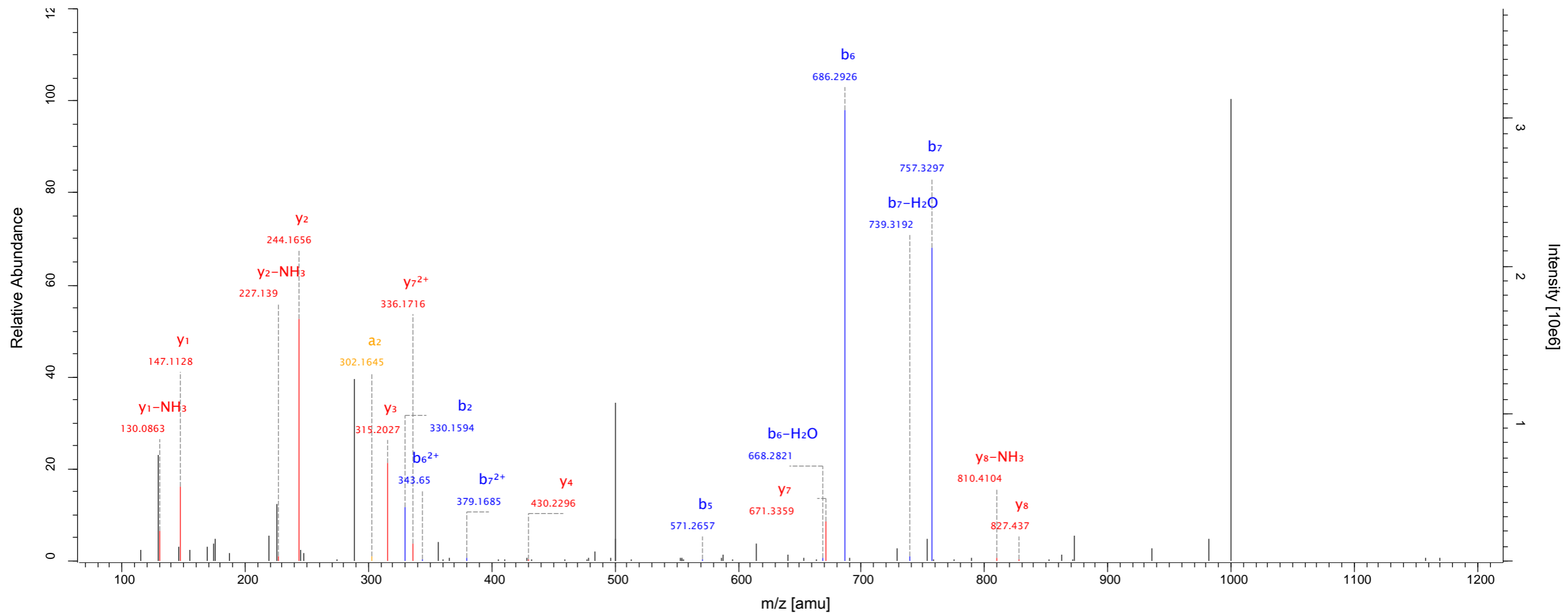
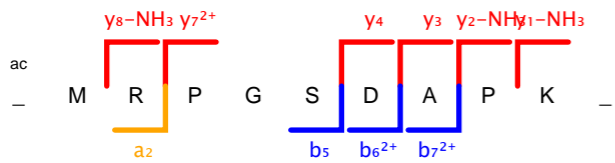
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 8375
Protein: ORFL190C - NC_006273.2[111965,112384]- <unknown description>
Peptide score: 69.95
Method: FTMS; HCD



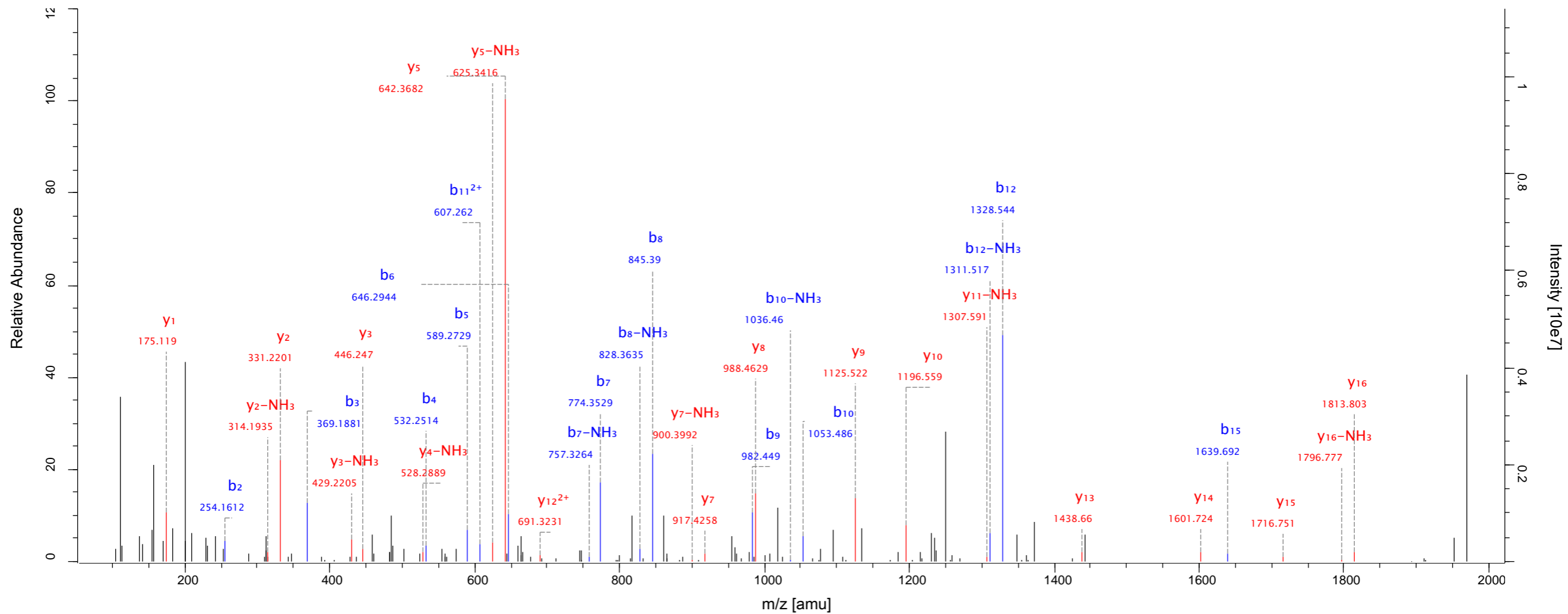
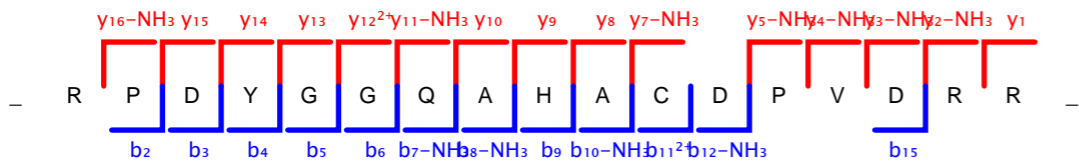
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
Scan number: 8420
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 50.9
Method: FTMS; HCD



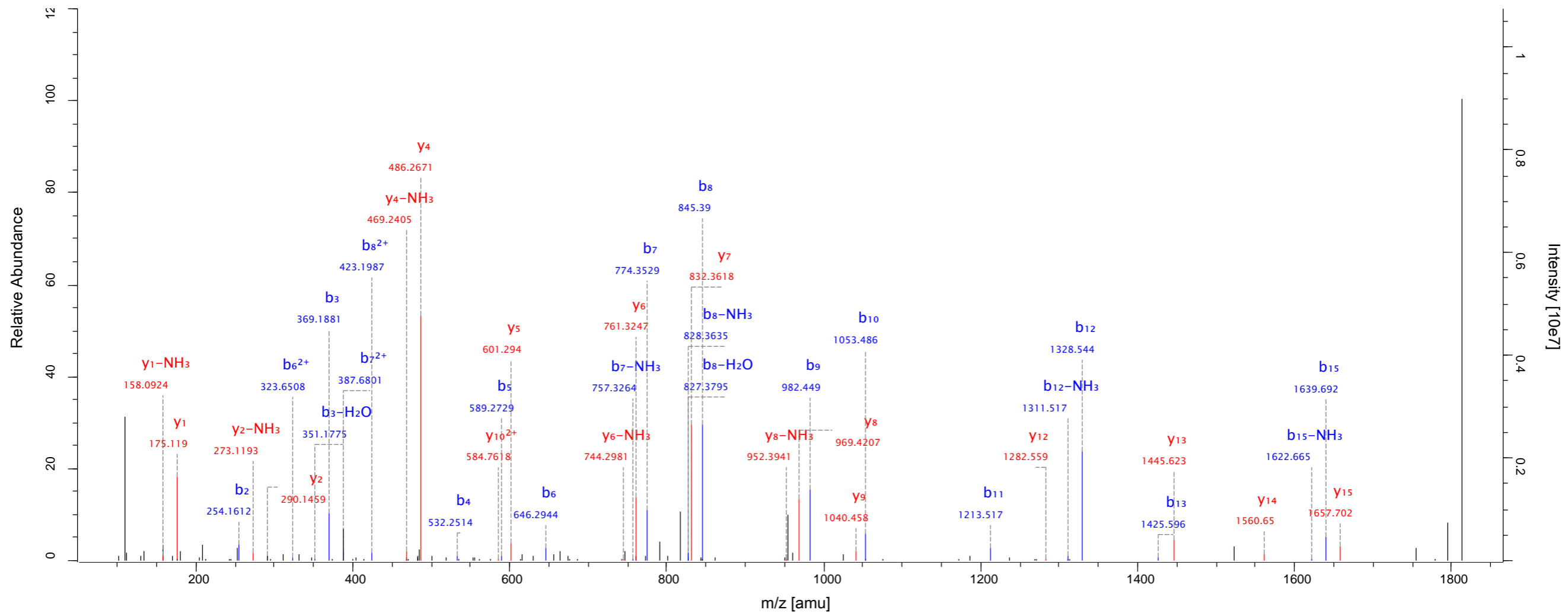
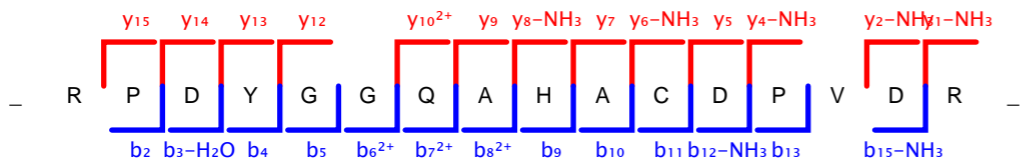
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
Scan number: 8648
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 117.81
Method: FTMS; HCD



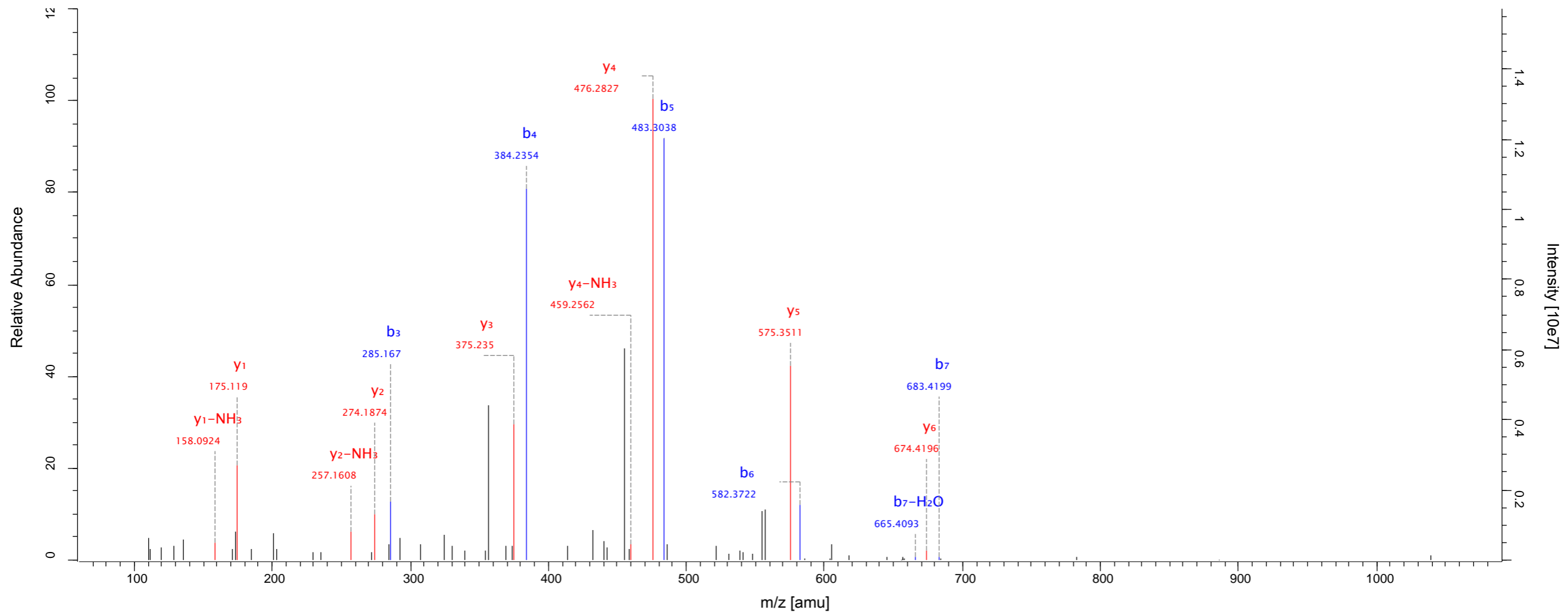
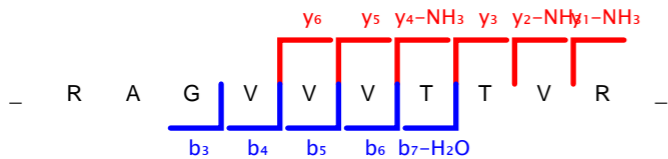
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_19
 Scan number: 9070
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 179.81
 Method: FTMS; HCD



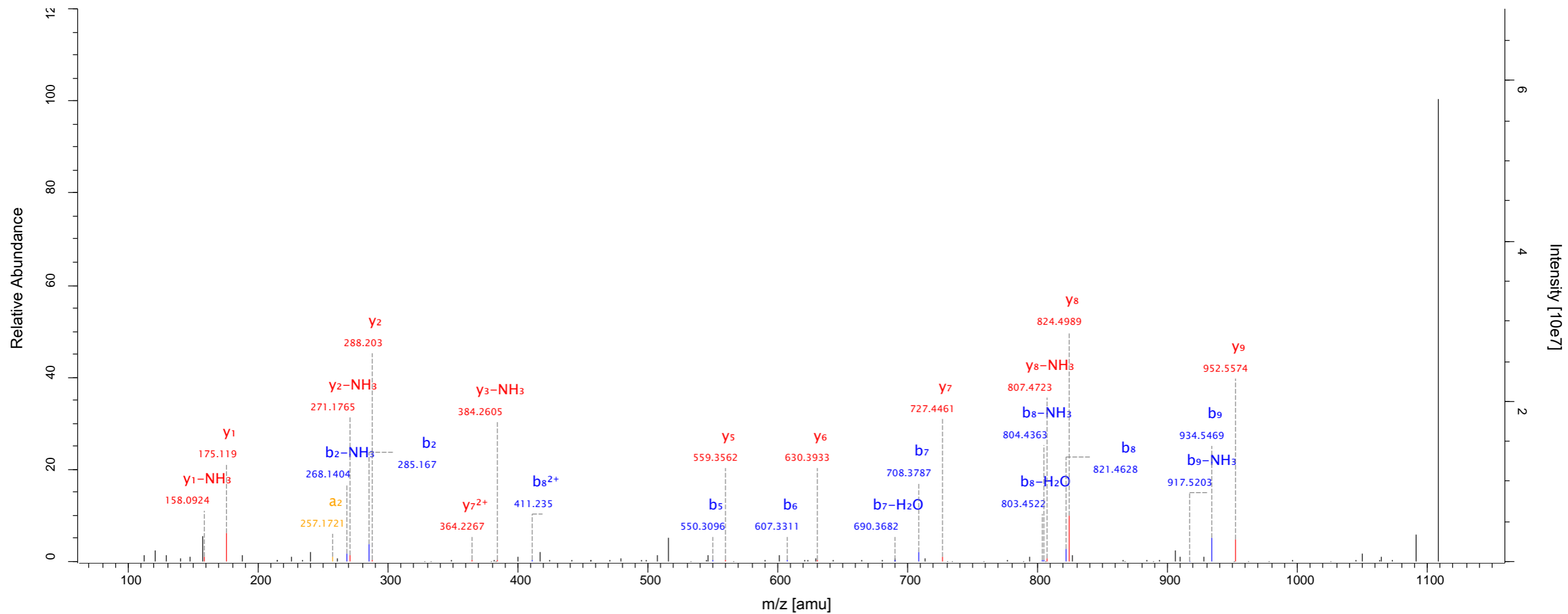
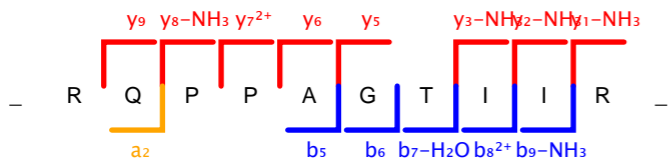
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_19
 Scan number: 9862
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 185.86
 Method: FTMS; HCD



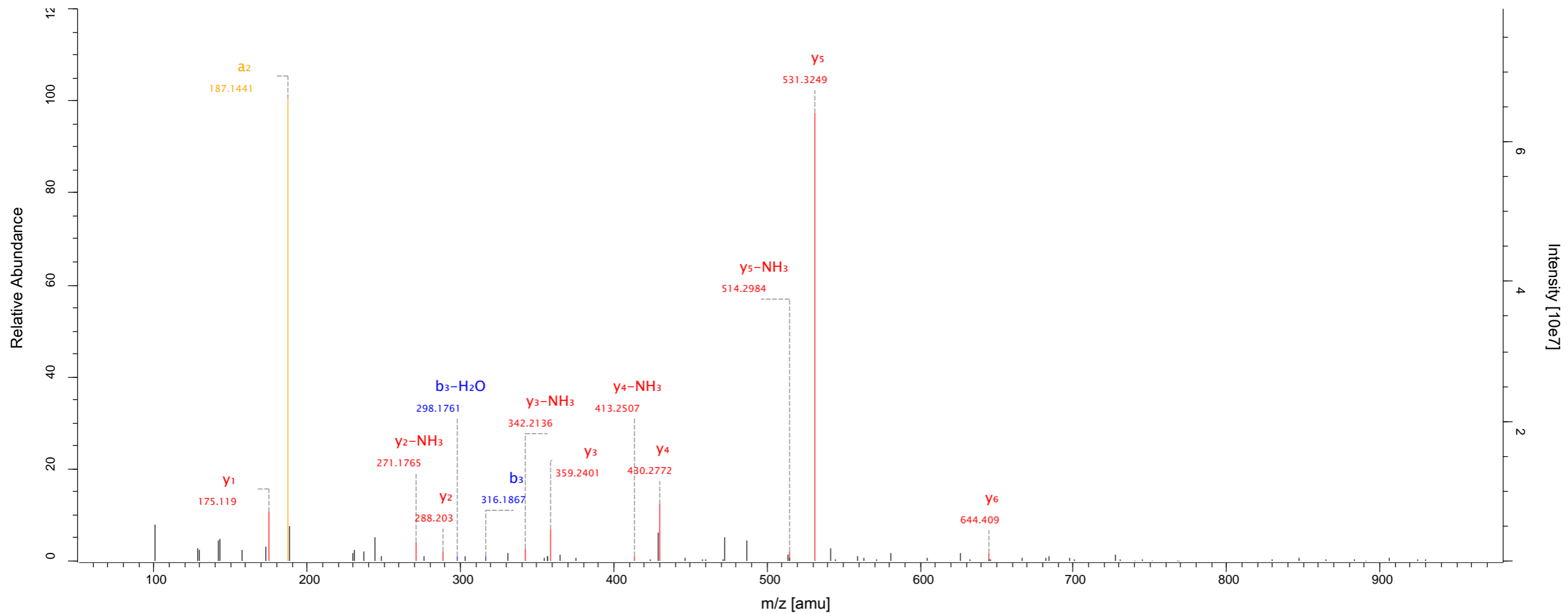
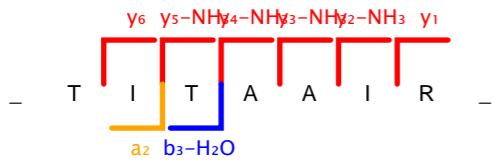
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 10278
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 103.7
Method: FTMS; HCD



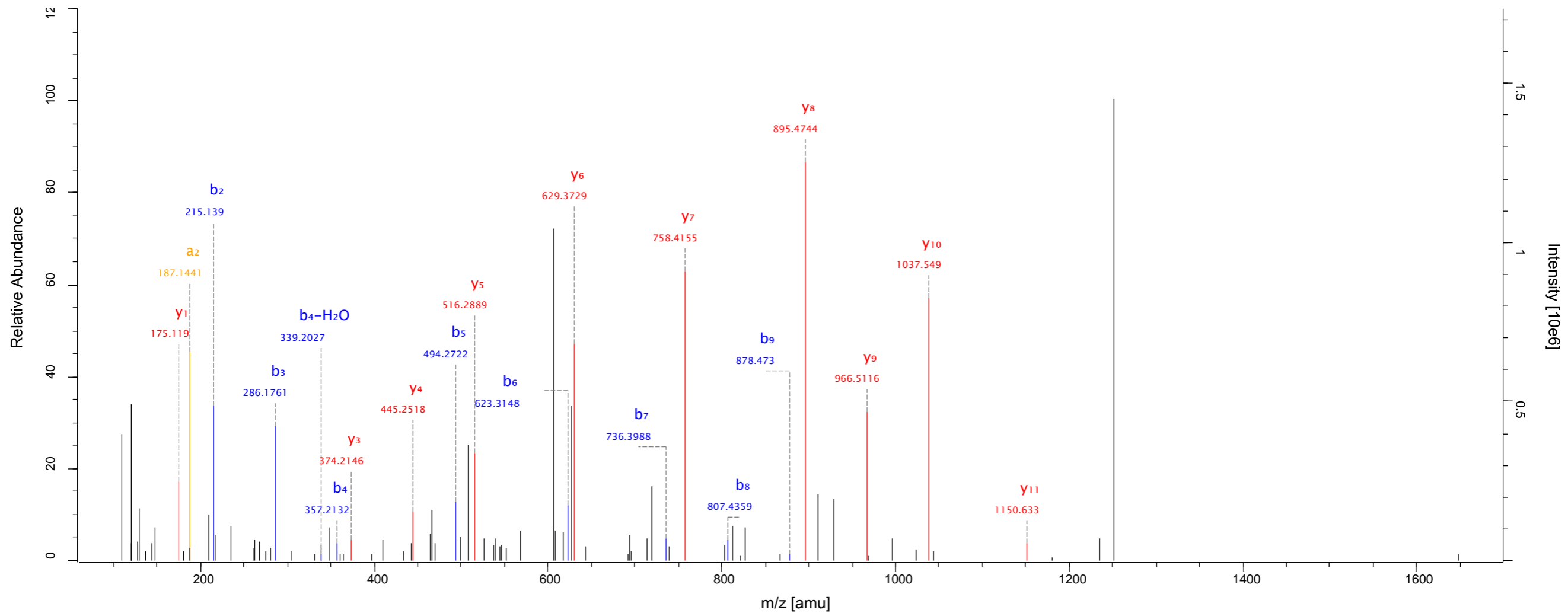
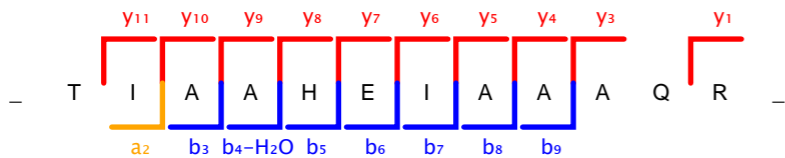
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 11821
Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
Peptide score: 142.36
Method: FTMS; HCD



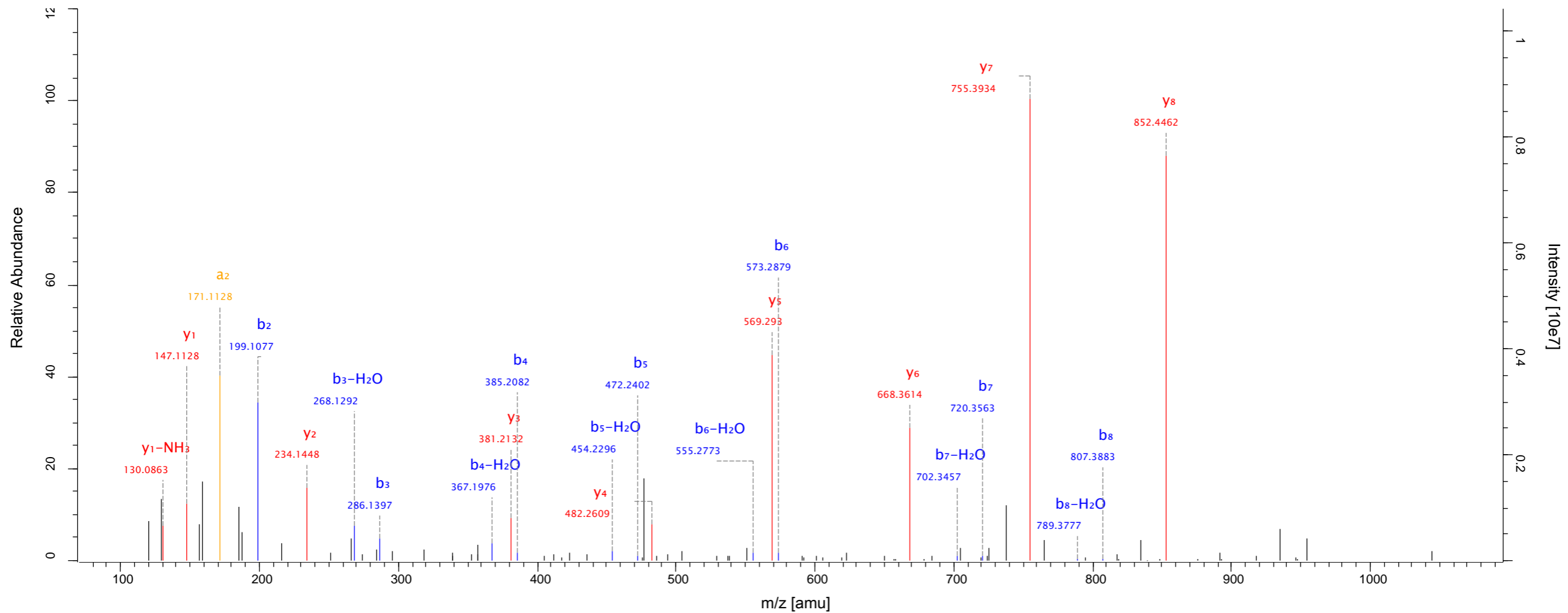
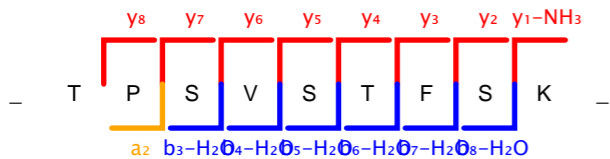
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 11832
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 99.38
Method: FTMS; HCD



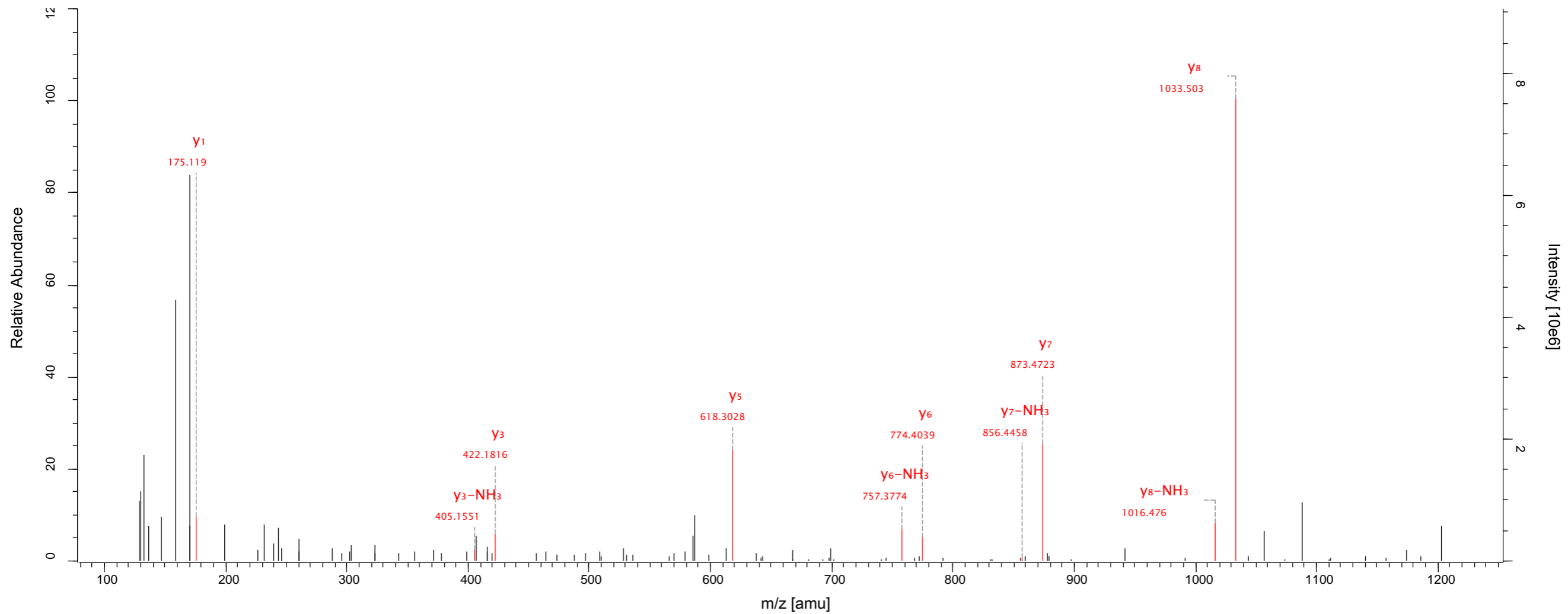
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 12102
Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
Peptide score: 132.32
Method: FTMS; HCD



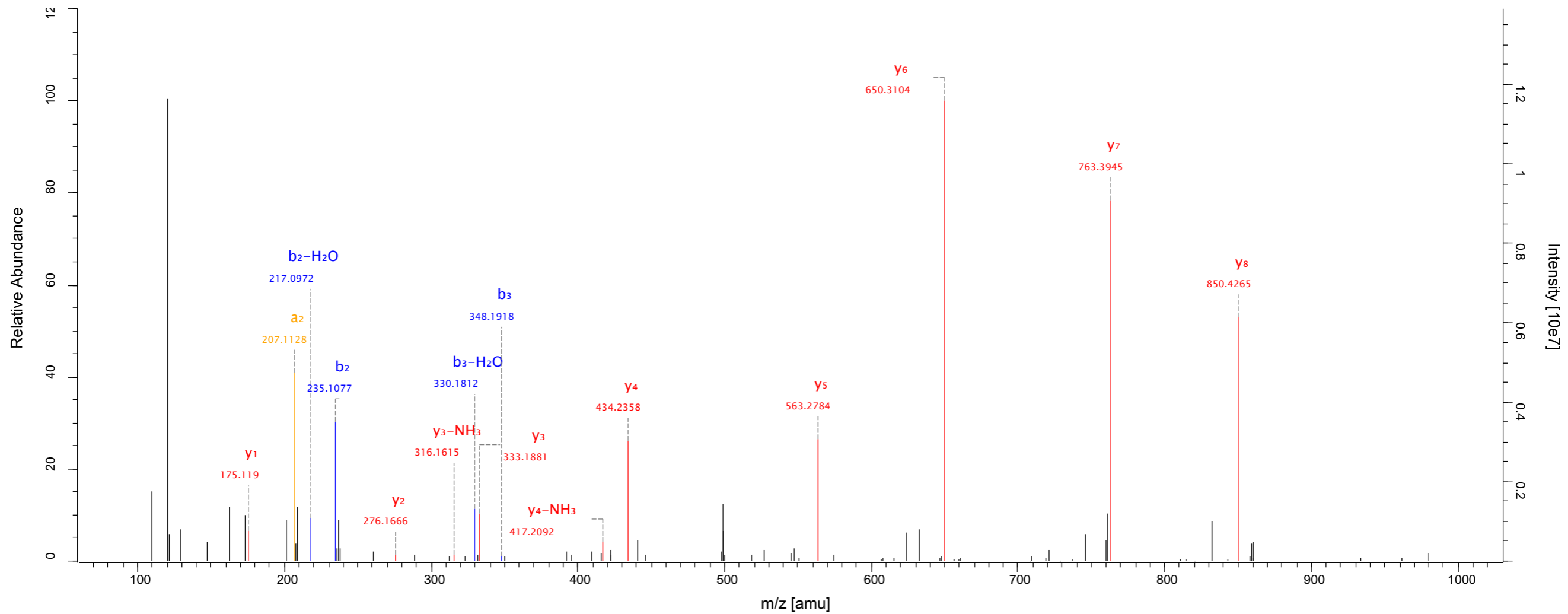
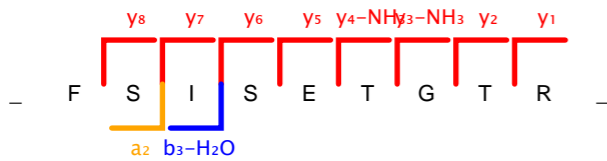
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 13098
Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
Peptide score: 148.14
Method: FTMS; HCD



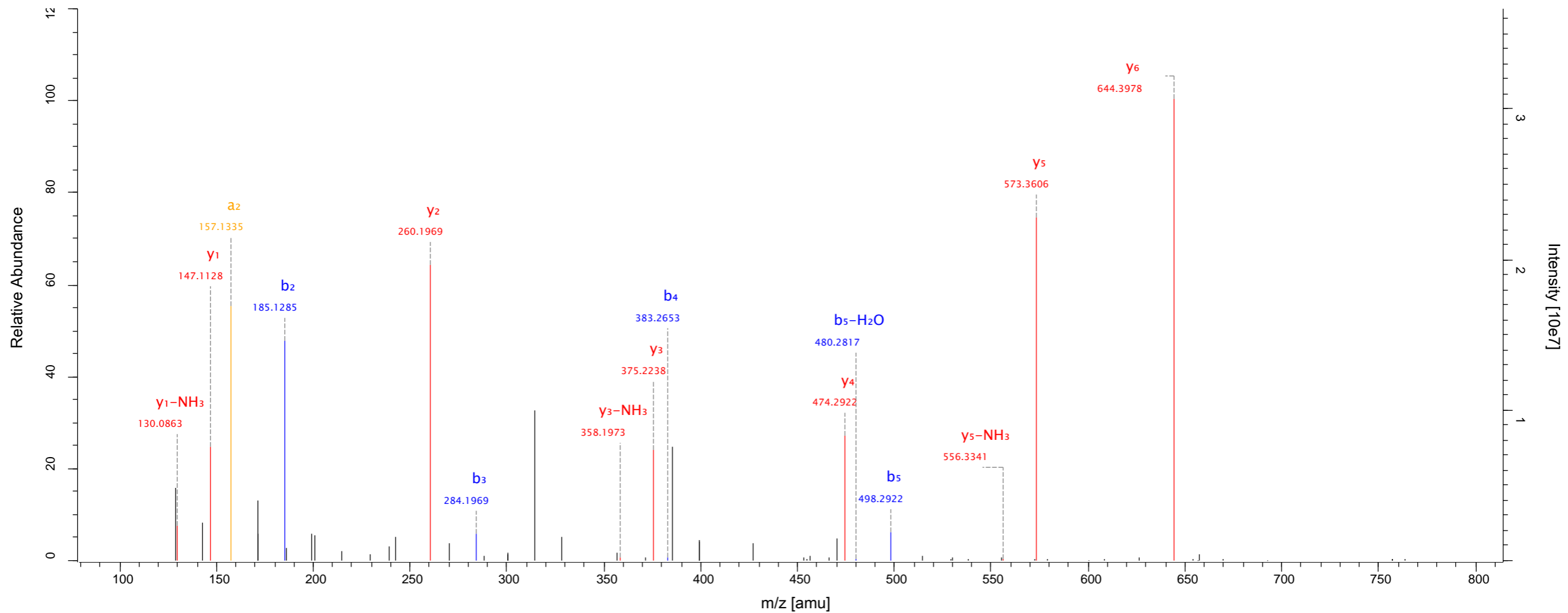
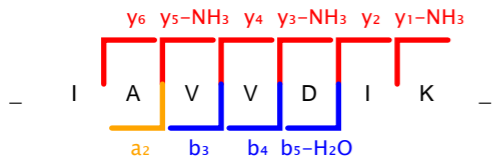
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 14619
Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
Peptide score: 59.8
Method: FTMS; HCD



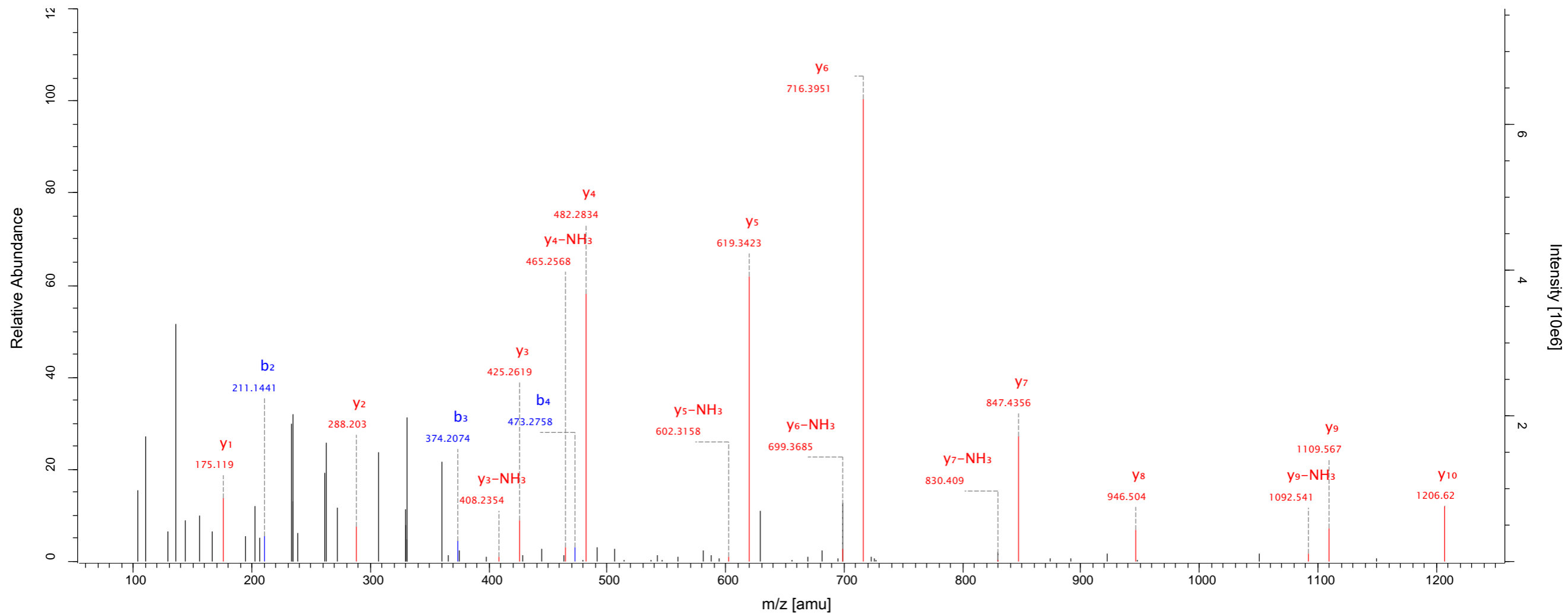
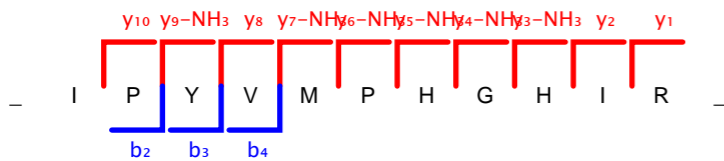
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 16027
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 80.24
Method: FTMS; HCD



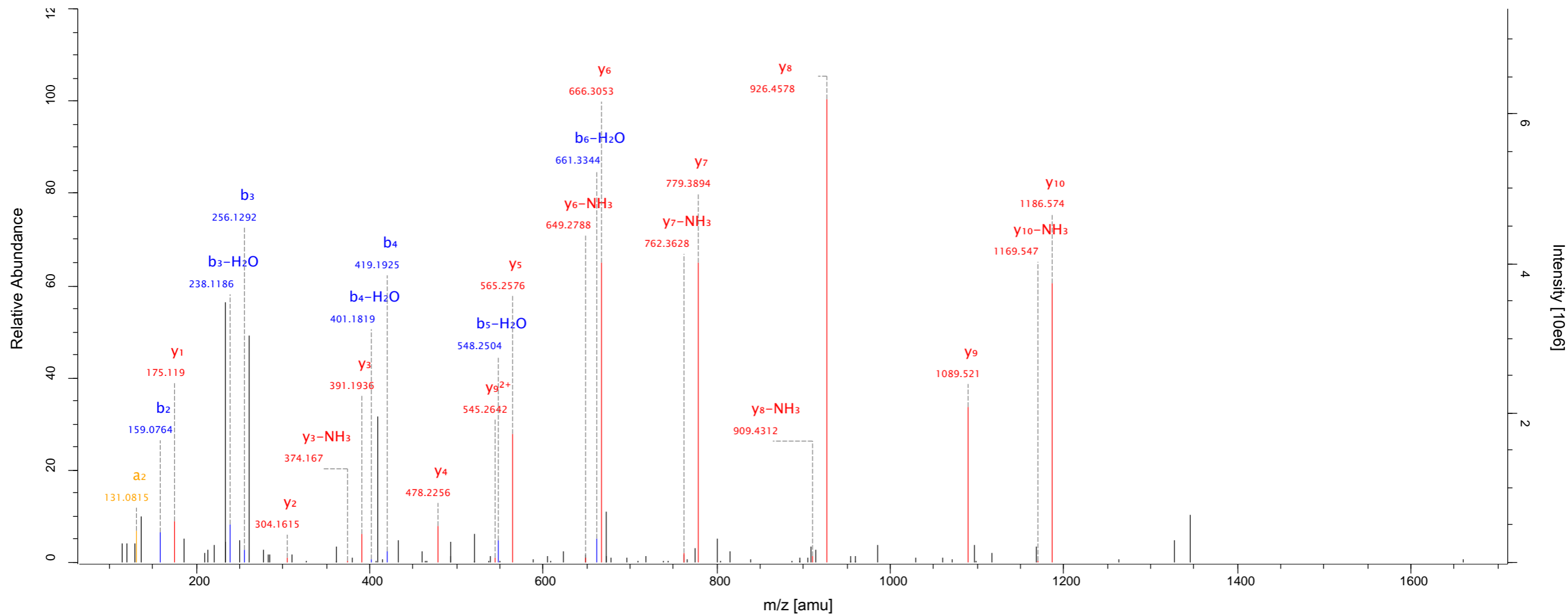
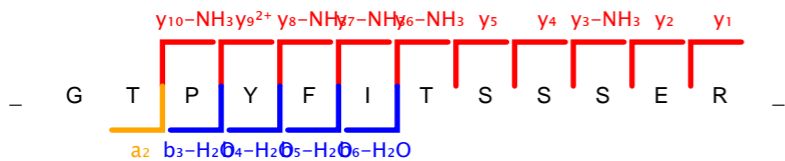
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 19547
Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
Peptide score: 111.81
Method: FTMS; HCD



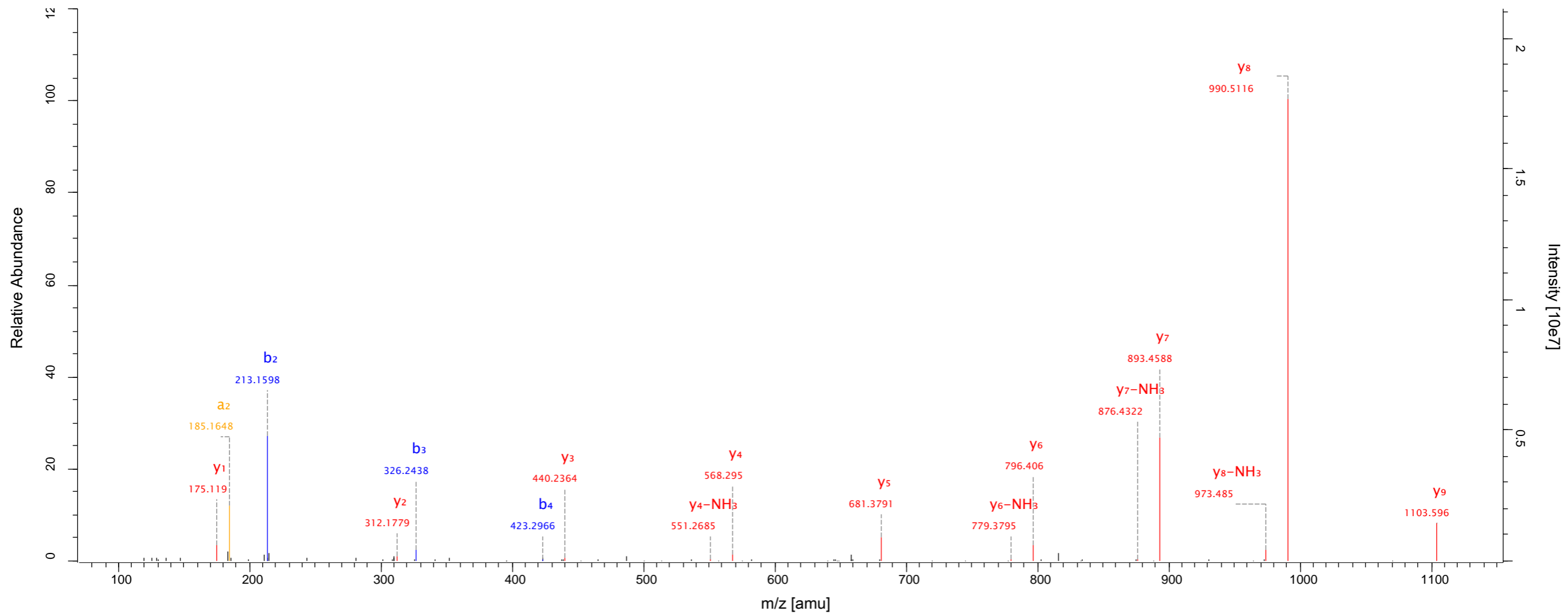
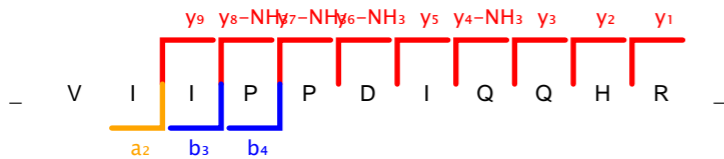
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 19674
Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>
Peptide score: 95.64
Method: FTMS; HCD



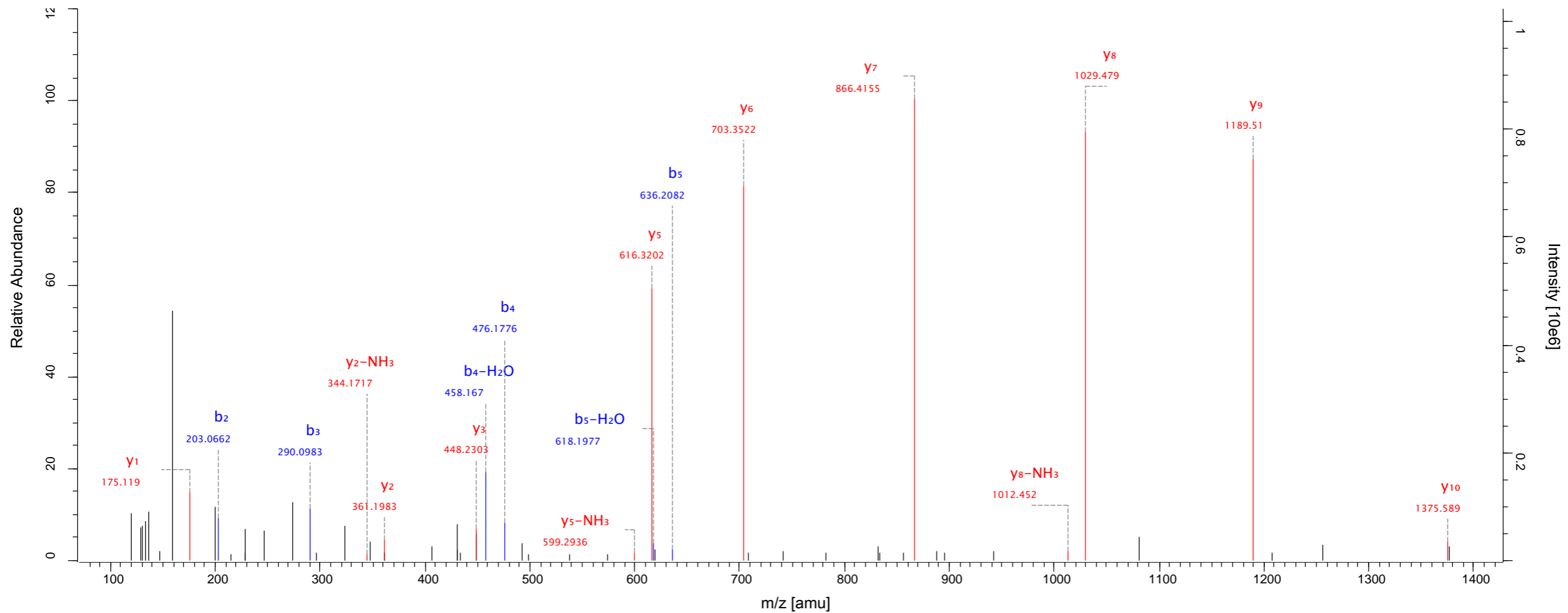
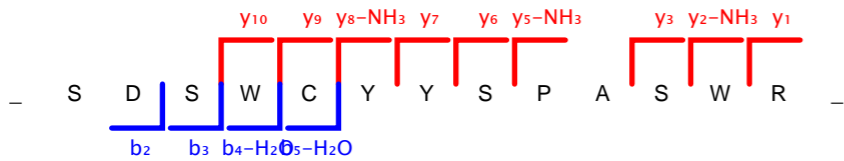
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
 Scan number: 23697
 Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
 Peptide score: 107.06
 Method: FTMS; HCD



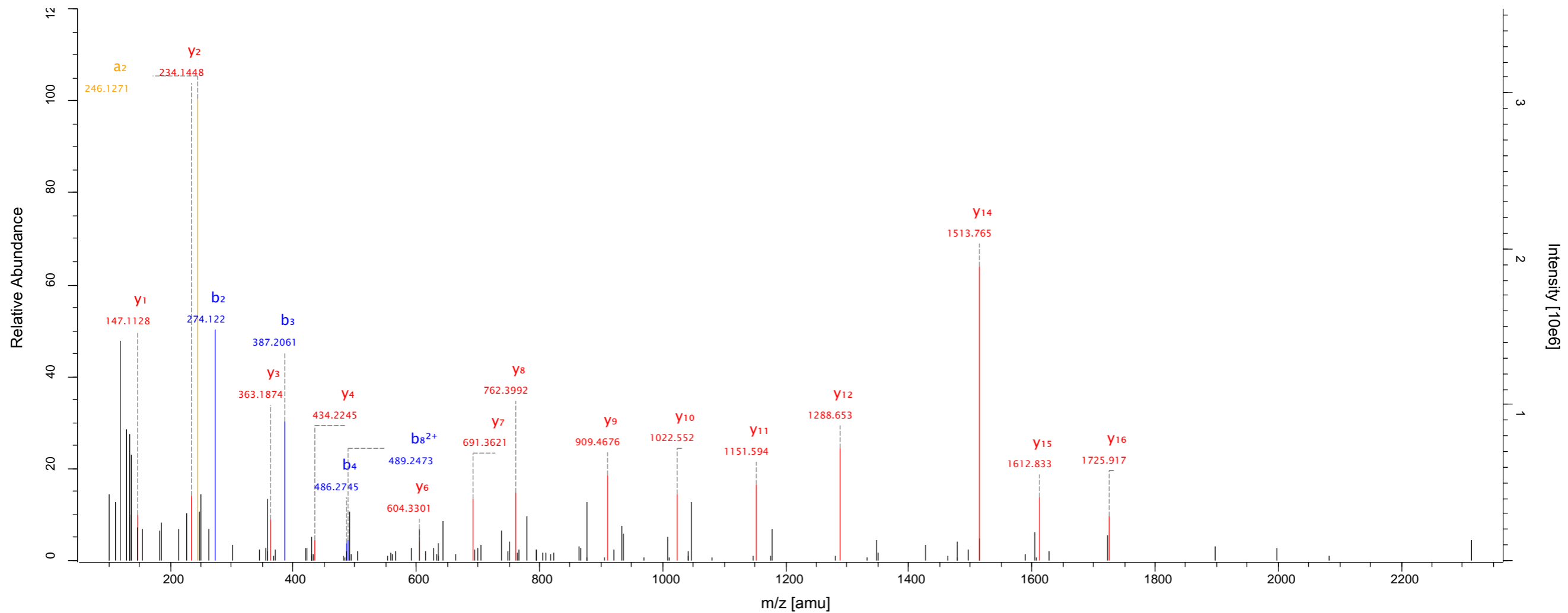
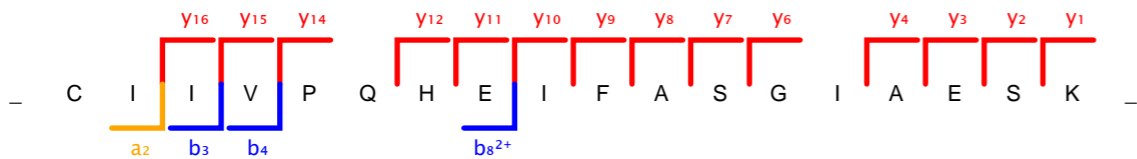
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 23913
Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
Peptide score: 154.56
Method: FTMS; HCD



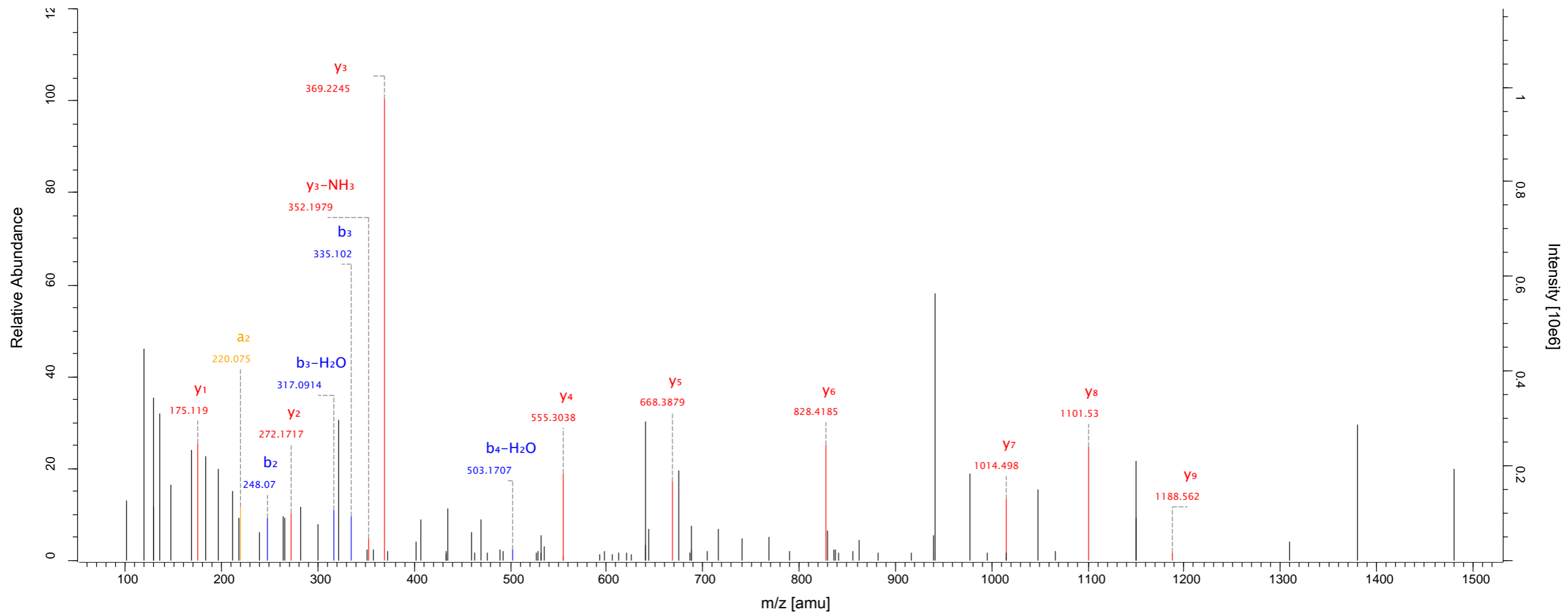
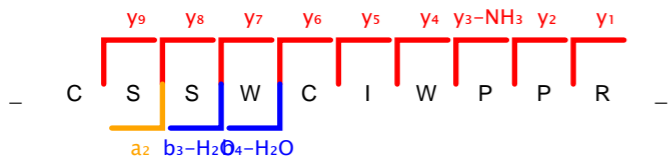
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 36698
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 107.18
Method: FTMS; HCD



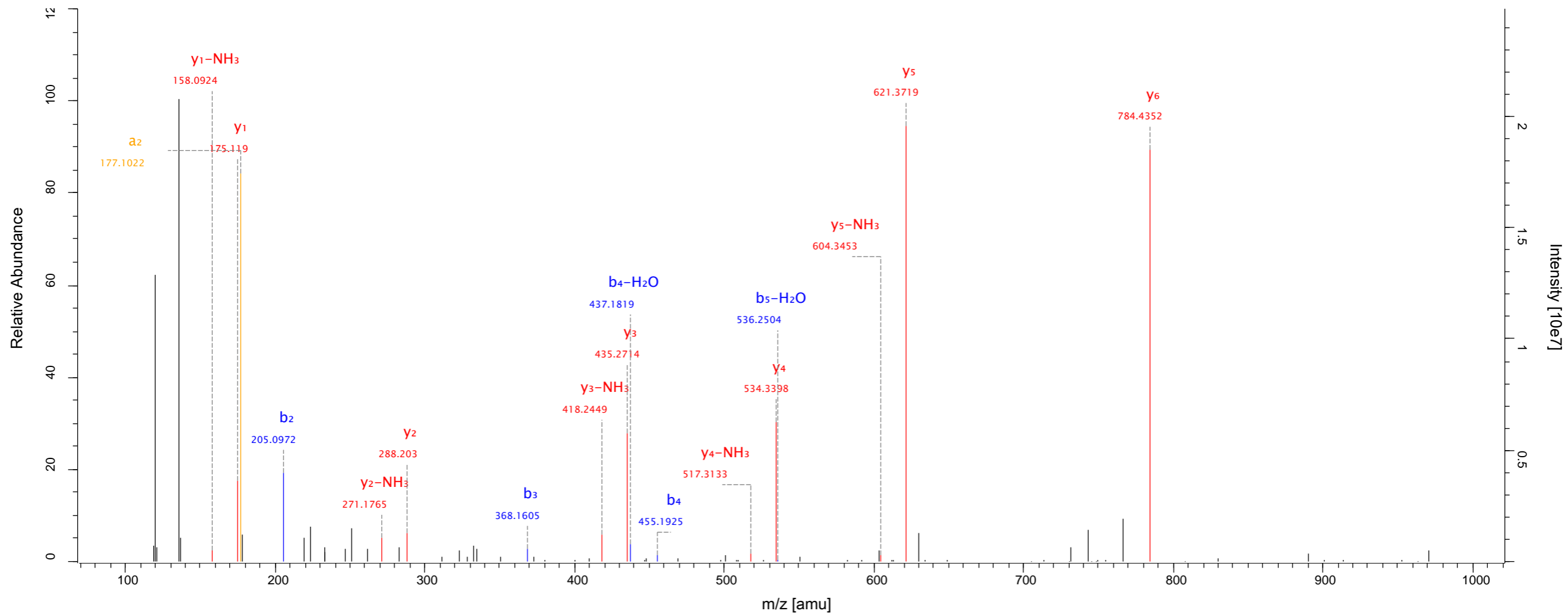
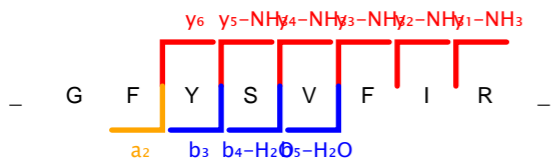
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 37368
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 104.41
Method: FTMS; HCD



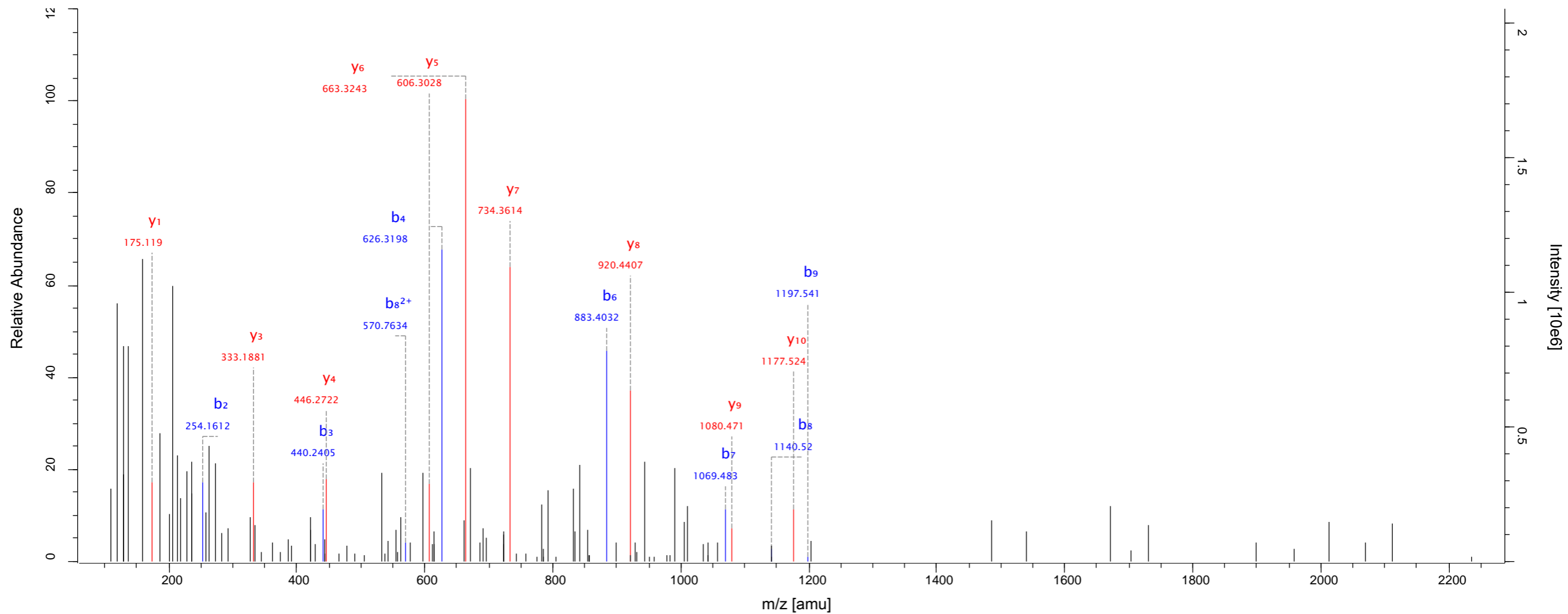
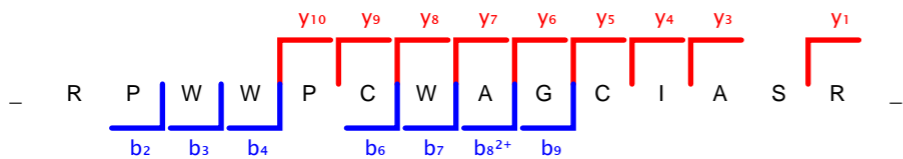
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 39208
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 78.56
Method: FTMS; HCD



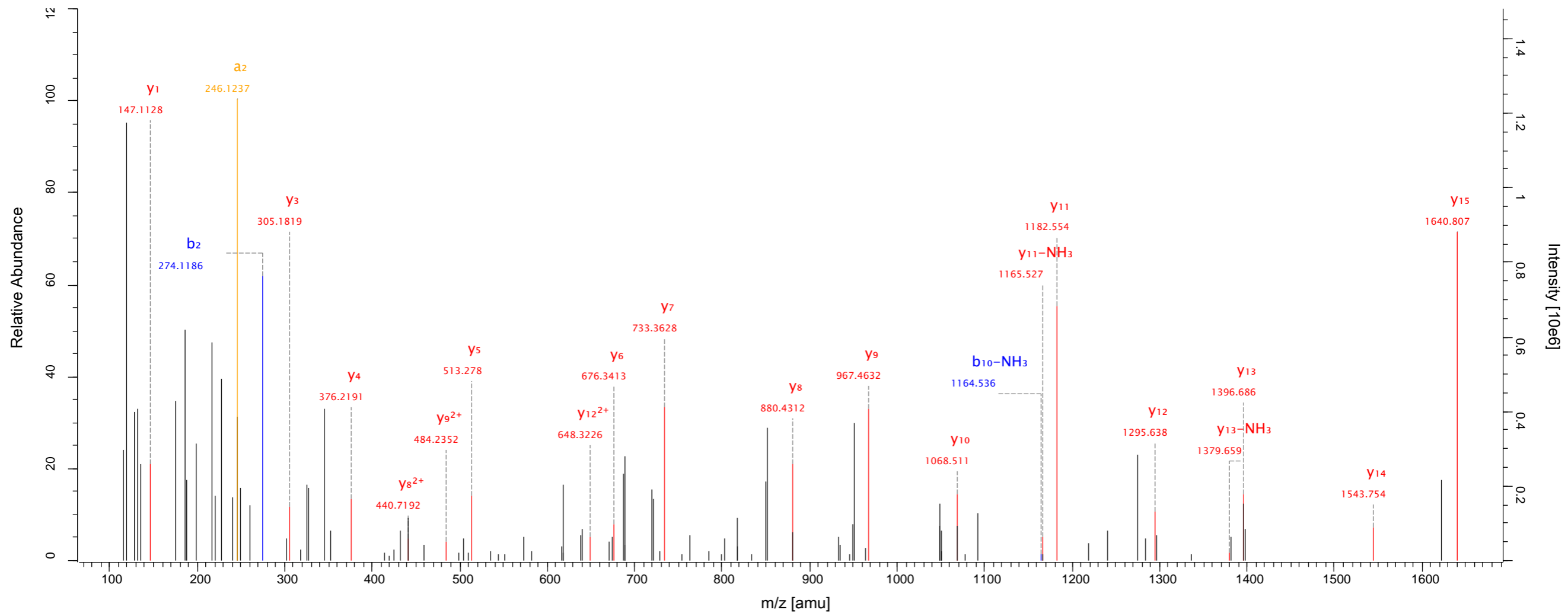
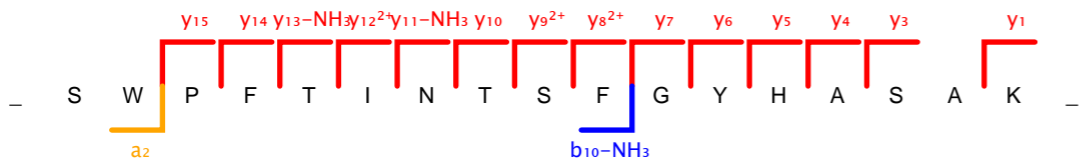
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
 Scan number: 40151
 Protein: ORFL151C - NC_006273.2[92499,92870]- <unknown description>
 Peptide score: 132.09
 Method: FTMS; HCD



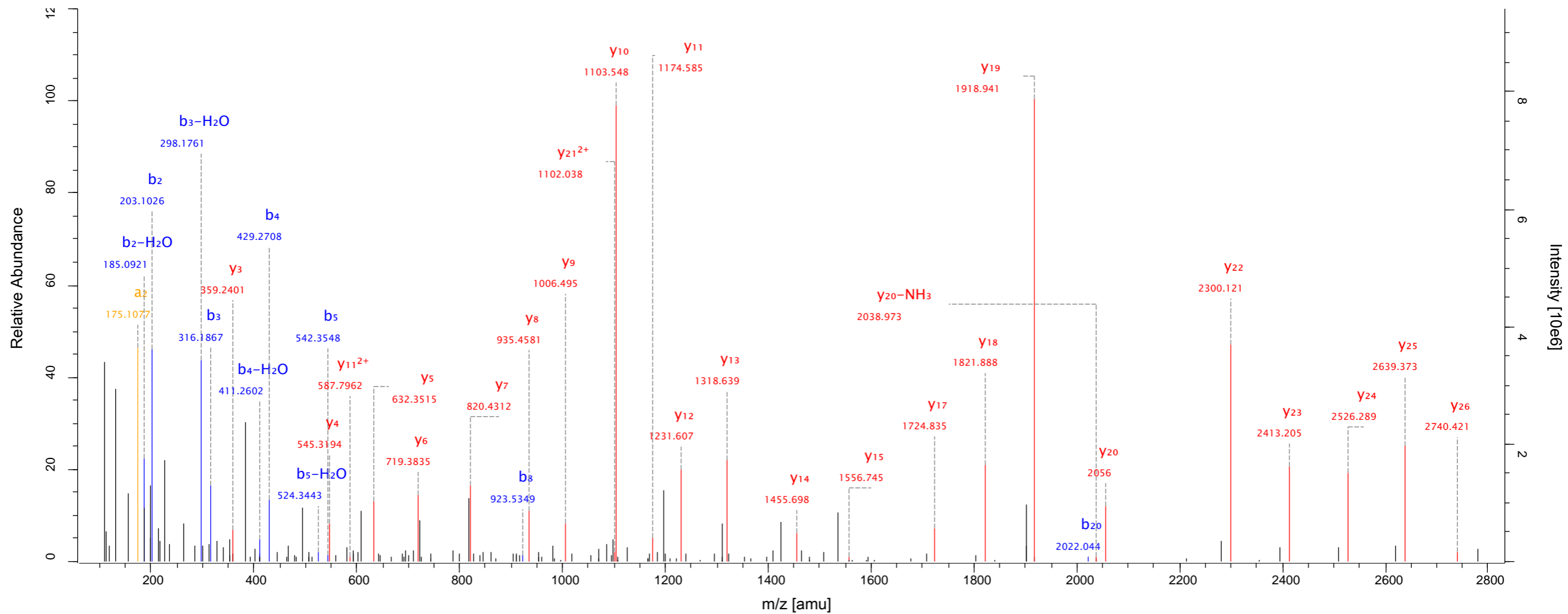
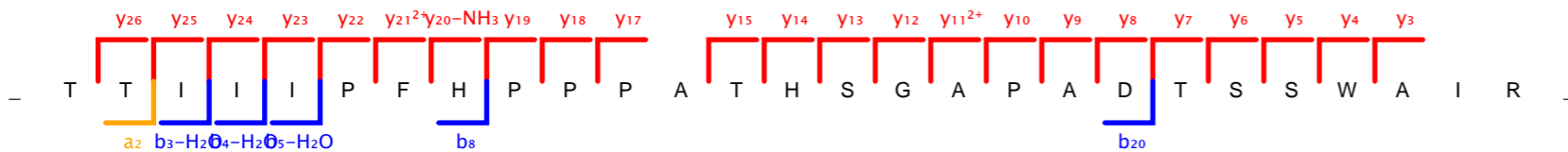
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 42230
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 88.55
Method: FTMS; HCD



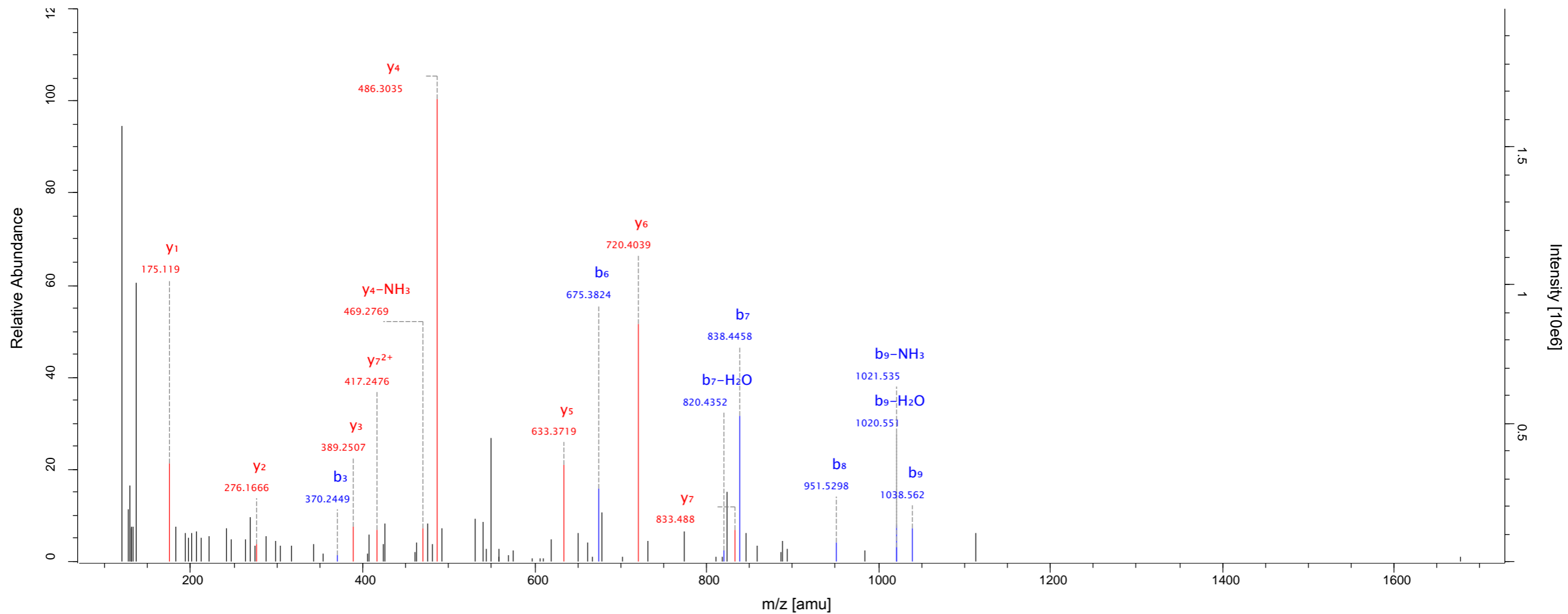
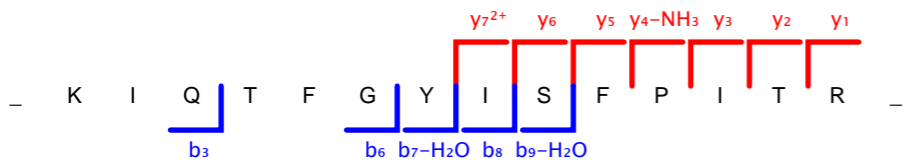
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 42479
Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
Peptide score: 89.54
Method: FTMS; HCD



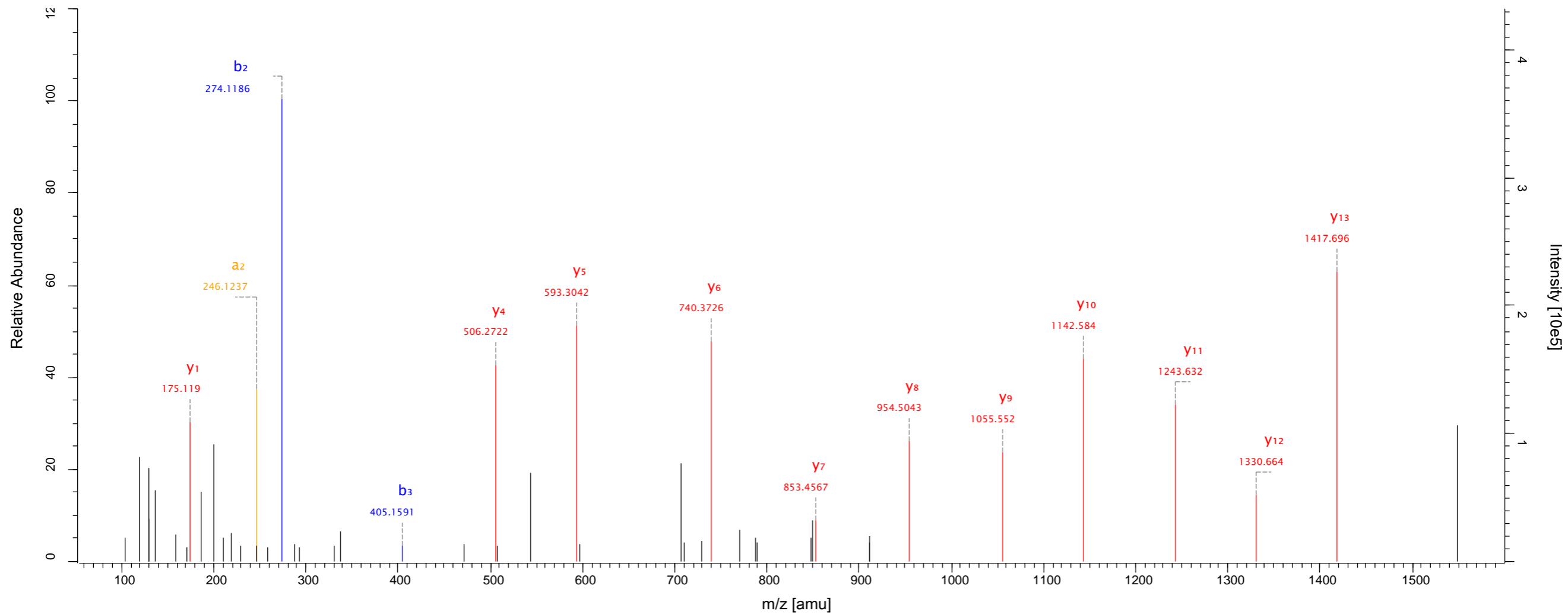
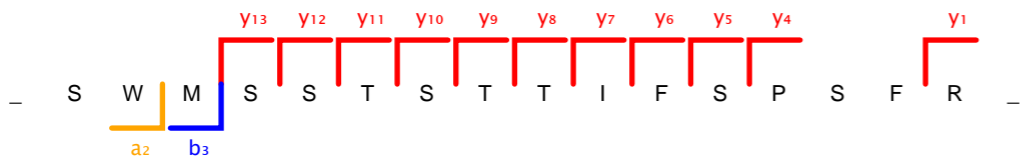
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_2
 Scan number: 43488
 Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
 Peptide score: 194.25
 Method: FTMS; HCD



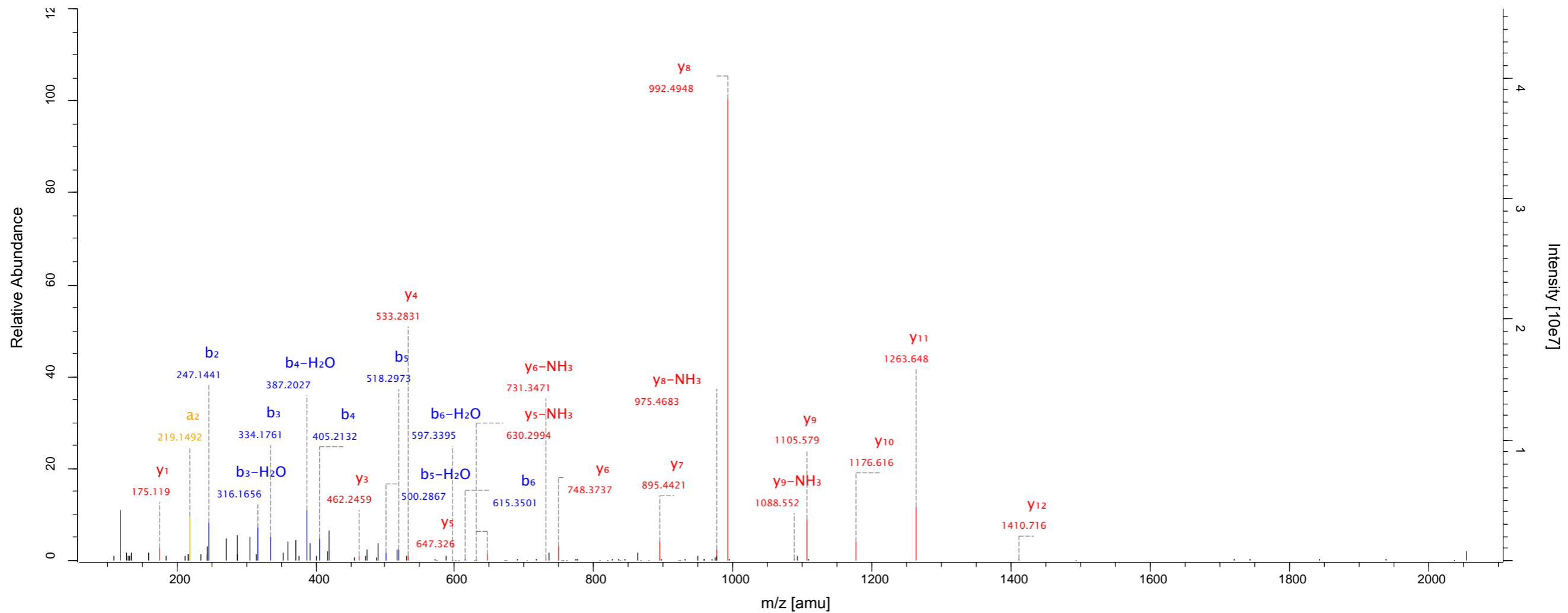
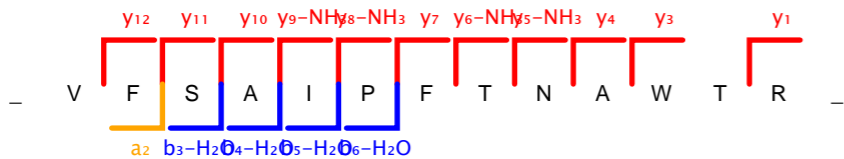
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 43790
Protein: ORFL6C - NC_006273.2[4179,4439]- <unknown description>
Peptide score: 78.49
Method: FTMS; HCD



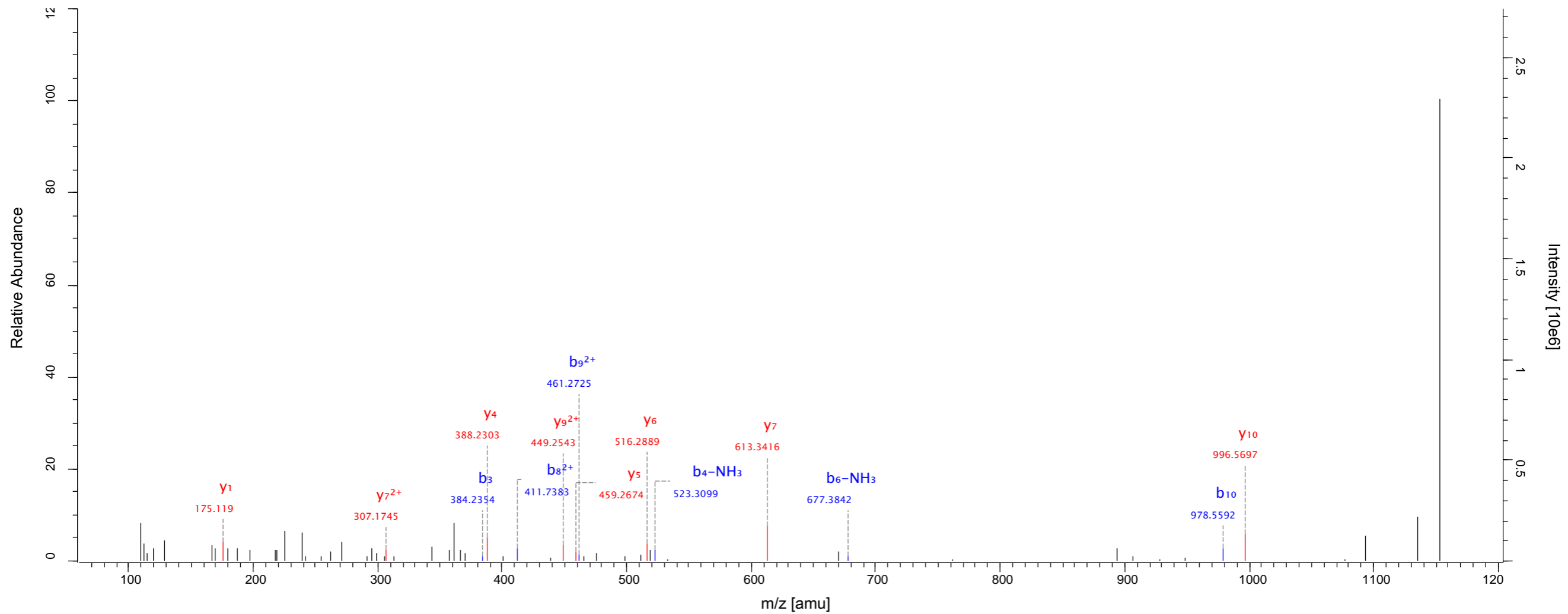
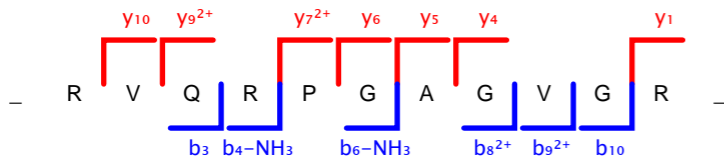
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 45254
Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
Peptide score: 106.13
Method: FTMS; HCD



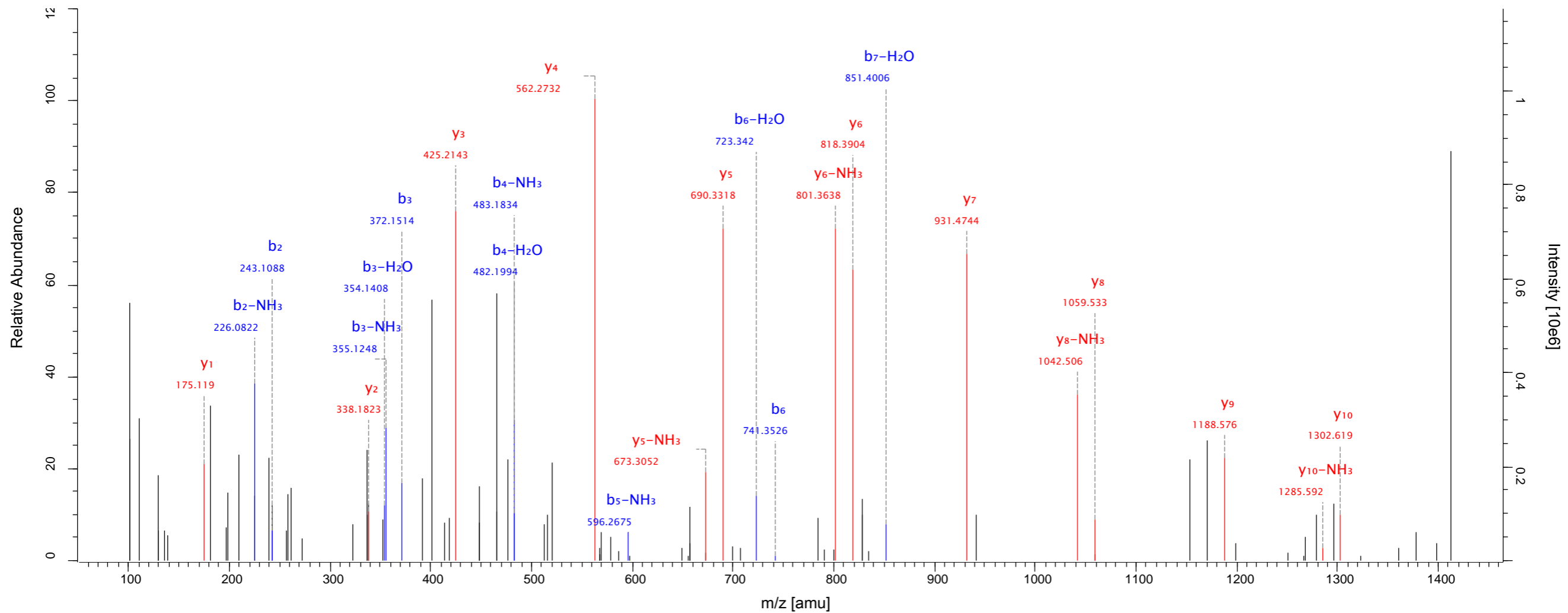
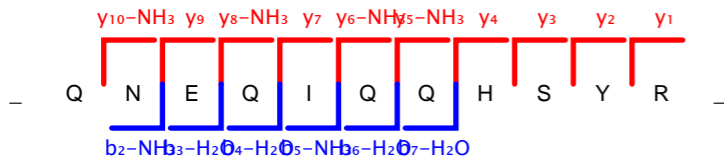
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
 Scan number: 46790
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 151.79
 Method: FTMS; HCD



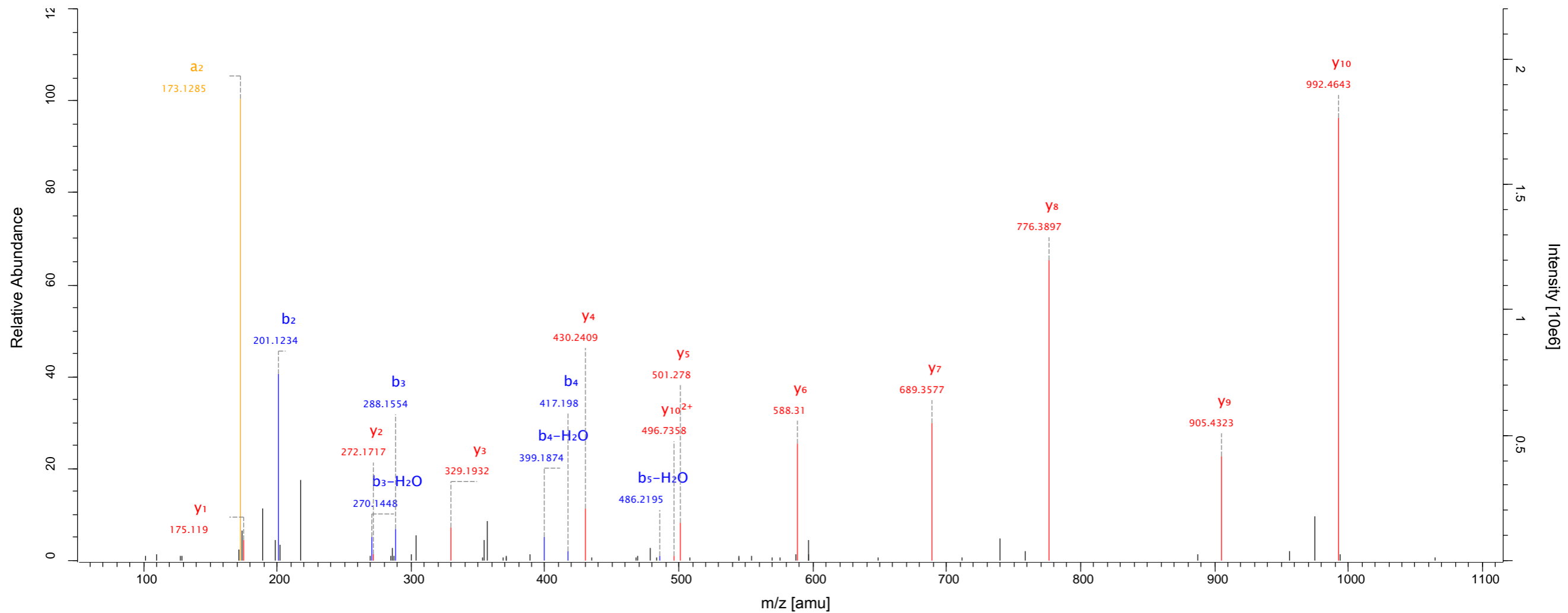
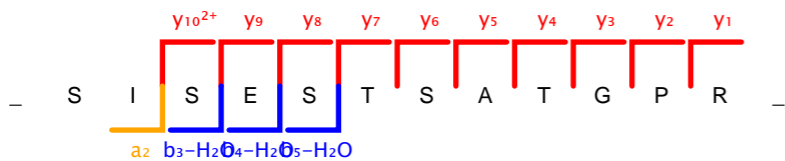
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 4937
Protein: ORFL121C - NC_006273.2[65579,65851]- <unknown description>
Peptide score: 57.79
Method: FTMS; HCD



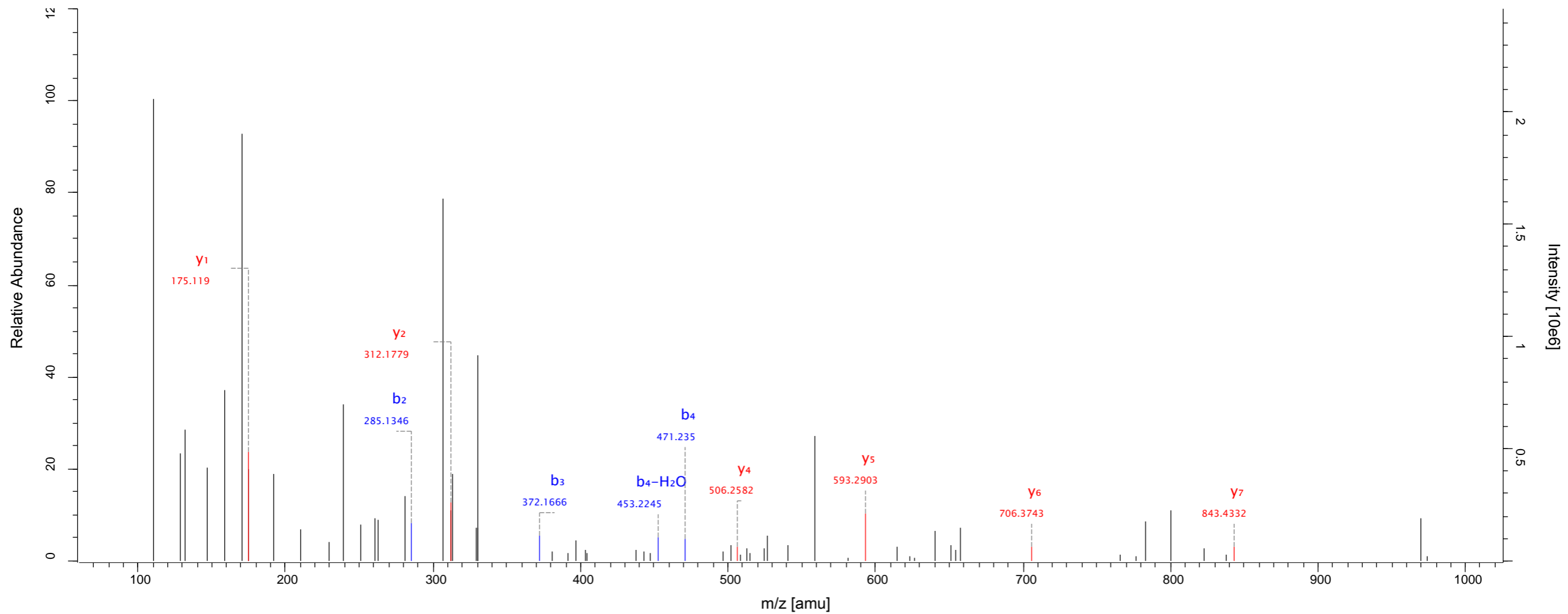
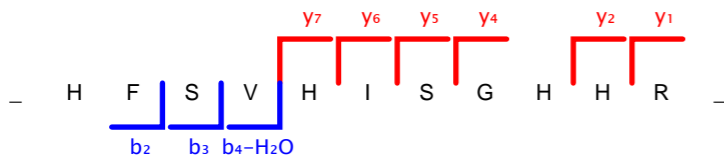
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
 Scan number: 6061
 Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
 Peptide score: 128.17
 Method: FTMS; HCD



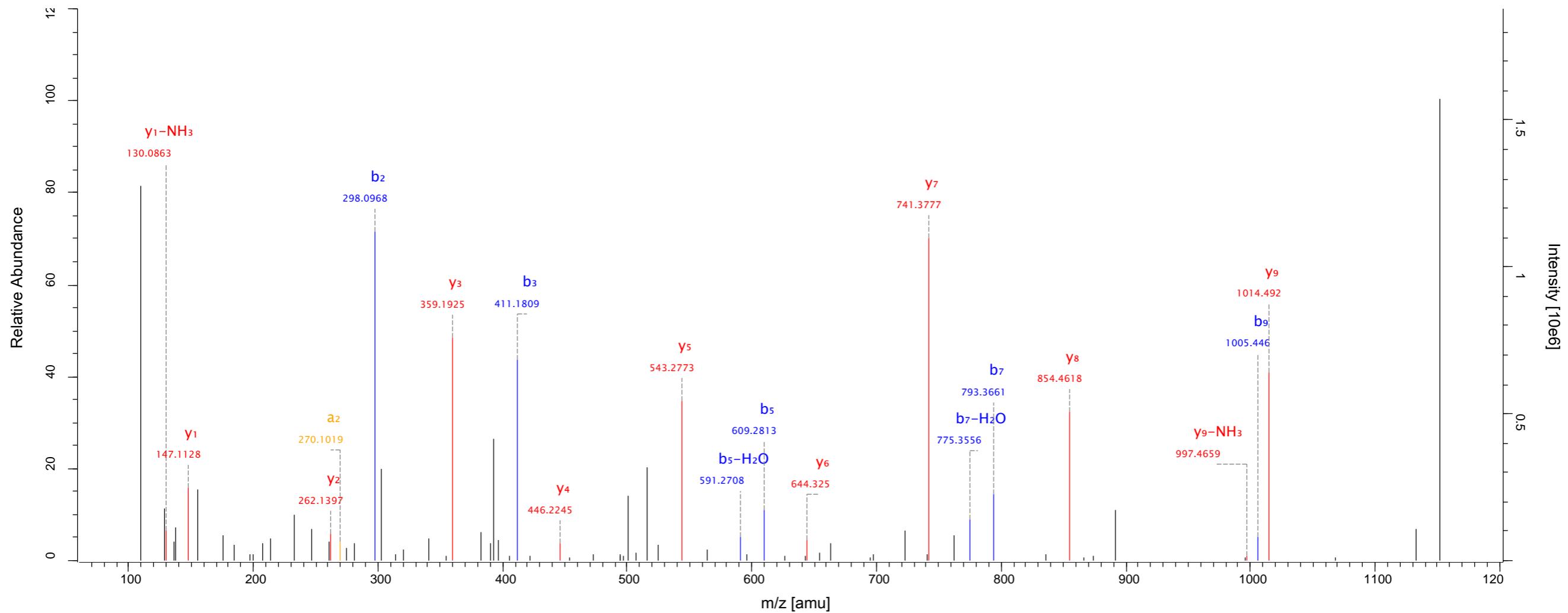
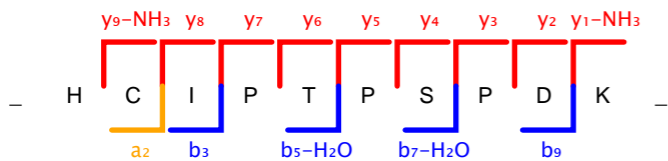
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 7468
Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
Peptide score: 101.93
Method: FTMS; HCD



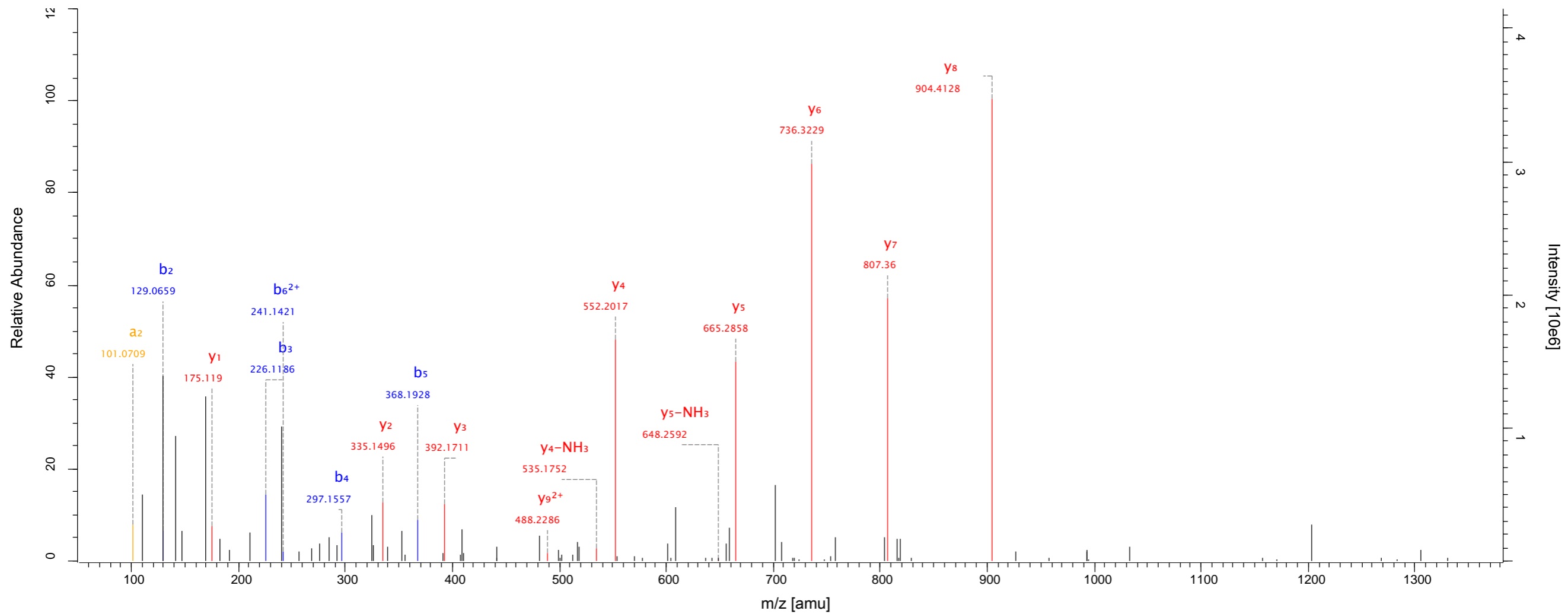
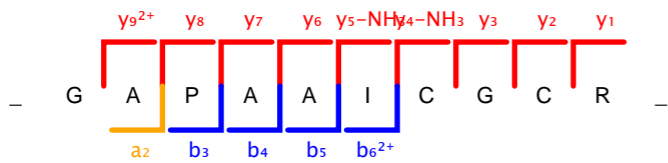
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_2
Scan number: 7754
Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
Peptide score: 40.73
Method: FTMS; HCD



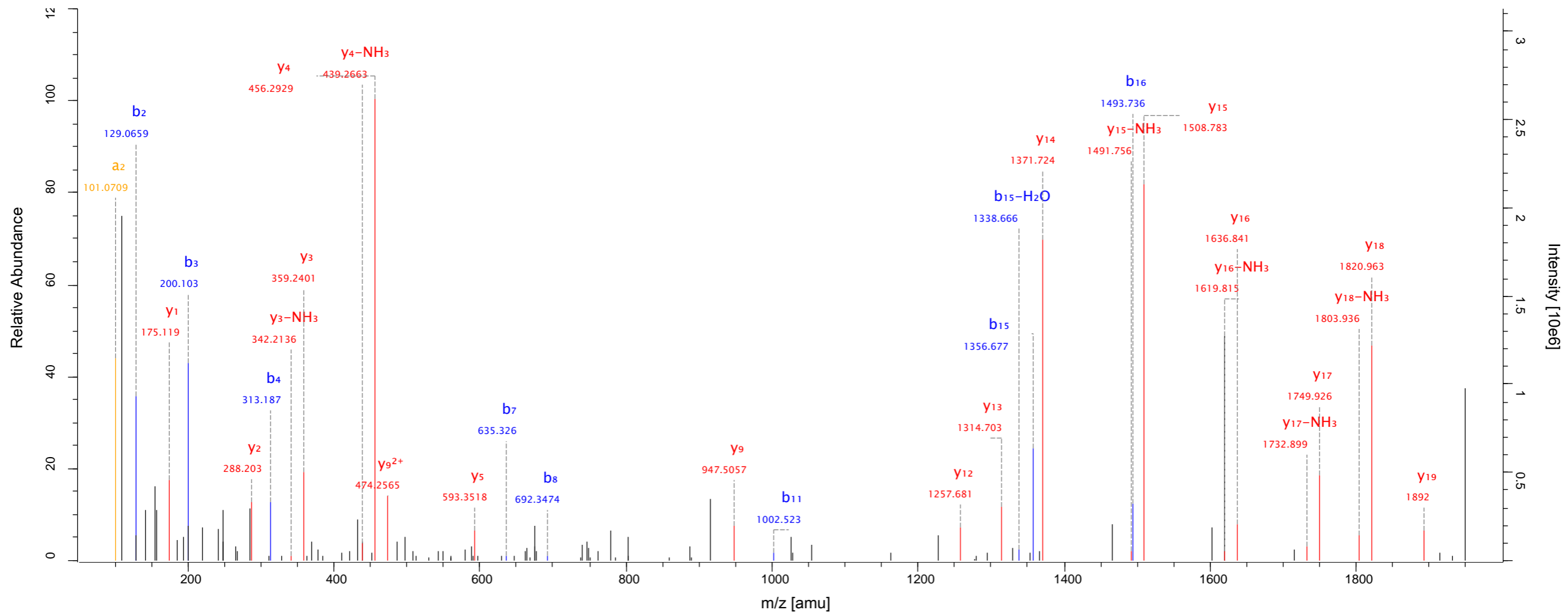
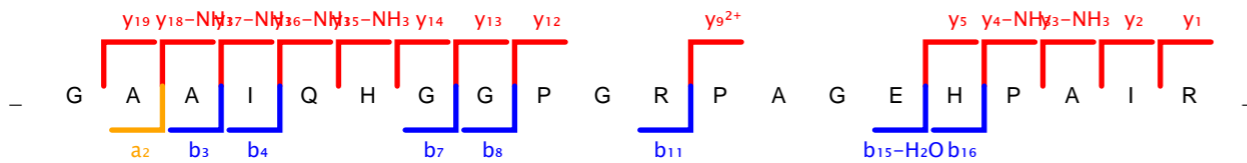
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 8677
Protein: ORFL26W + NC_006273.2[12923,13093]+ <unknown description>
Peptide score: 142.92
Method: FTMS; HCD



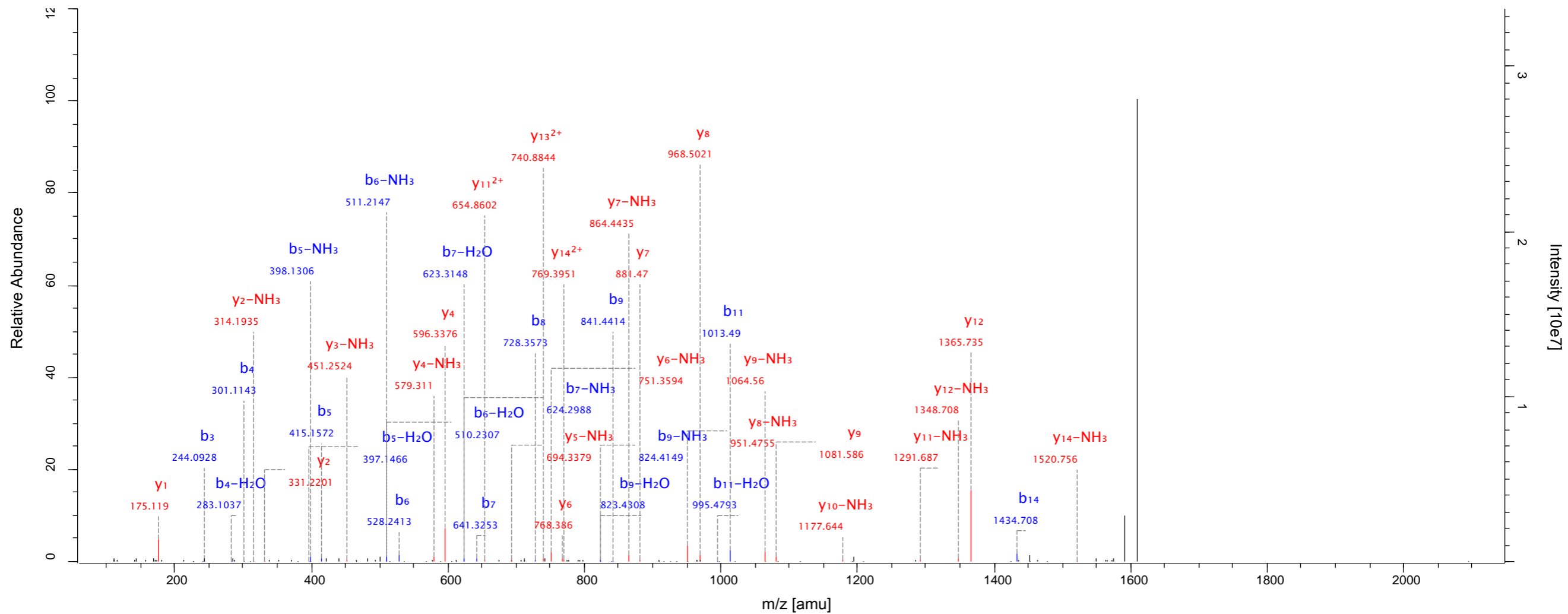
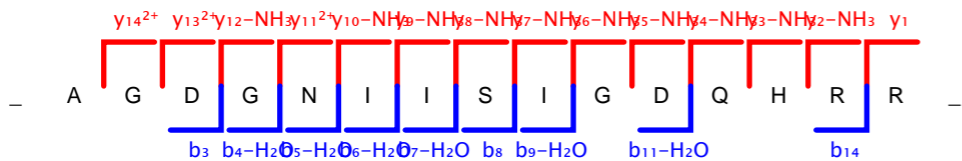
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
Scan number: 8842
Protein: ORFL73C - NC_006273.2[30581,30946]- <unknown description>
Peptide score: 95.5
Method: FTMS; HCD



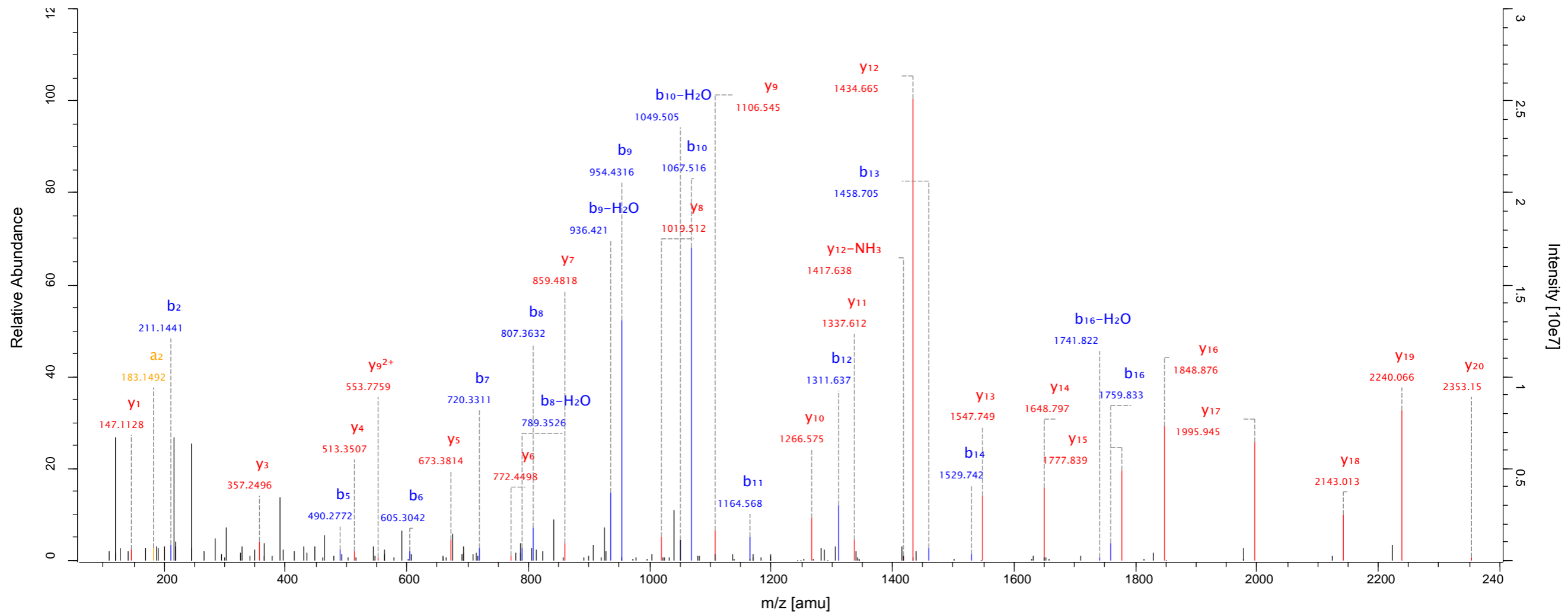
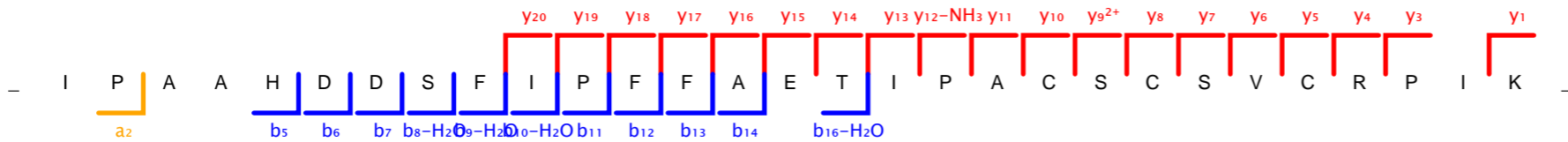
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_2
 Scan number: 9249
 Protein: ORFS359W + NC_006273.2[221986,222303]+ <unknown description>
 Peptide score: 175.65
 Method: FTMS; HCD



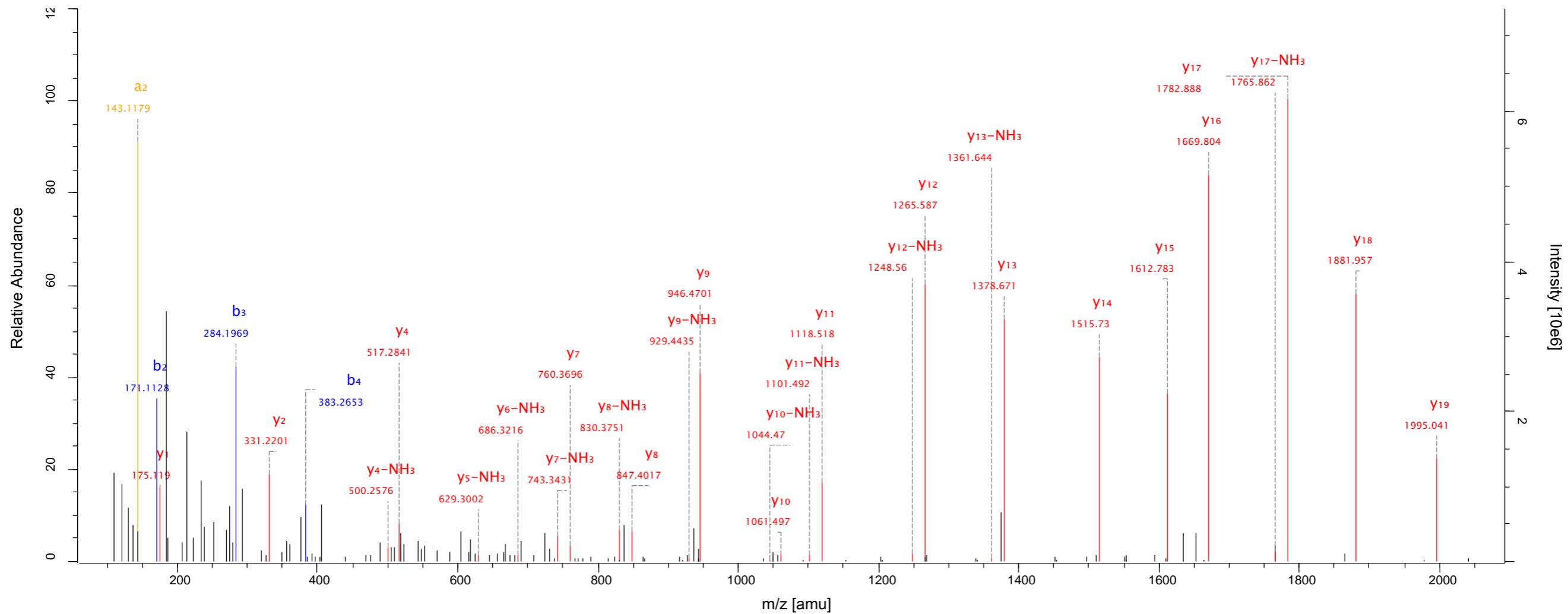
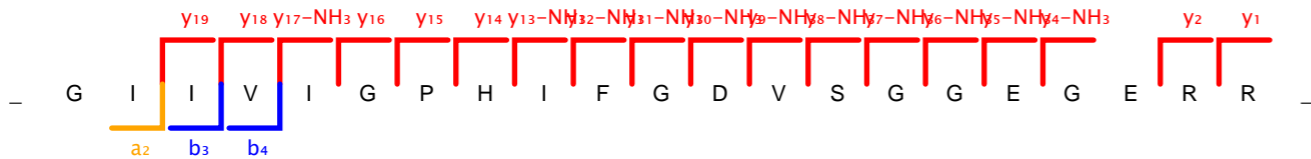
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_20
 Scan number: 17814
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 195.71
 Method: FTMS; HCD



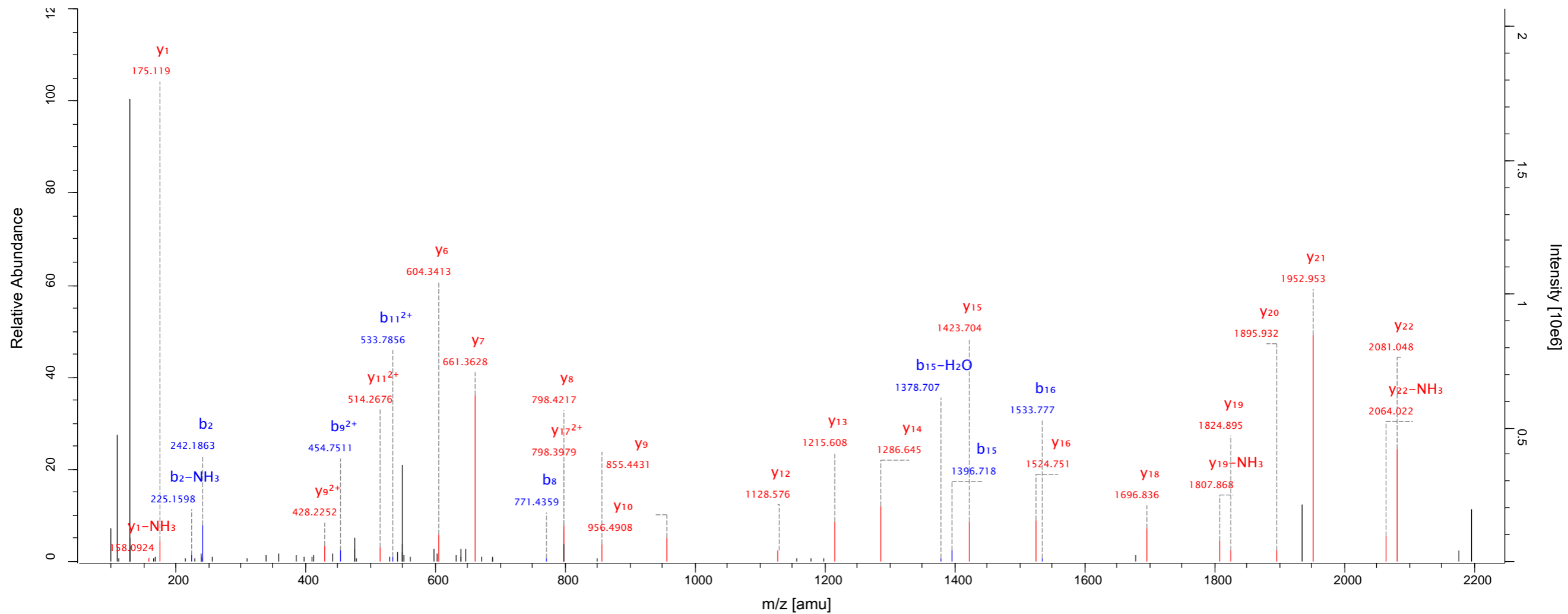
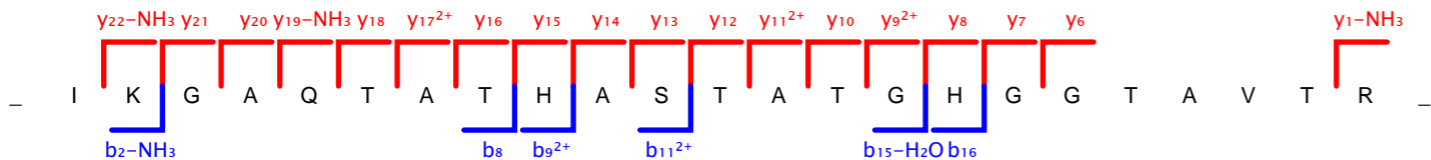
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_20
 Scan number: 55496
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 158.94
 Method: FTMS; HCD



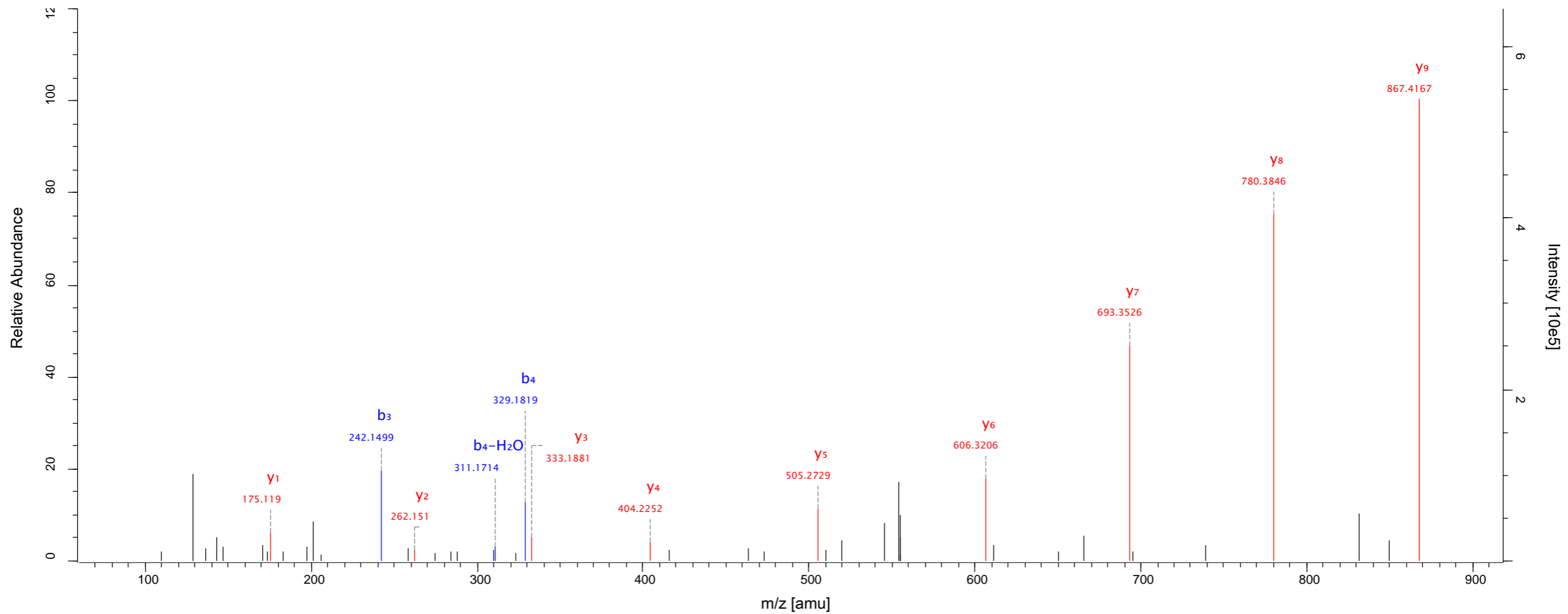
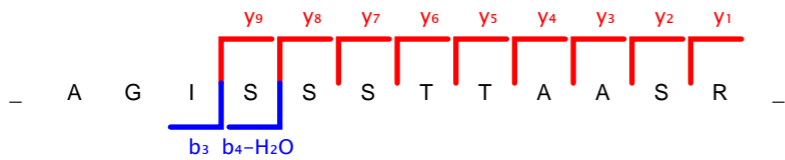
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_20
 Scan number: 55519
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 162.41
 Method: FTMS; HCD



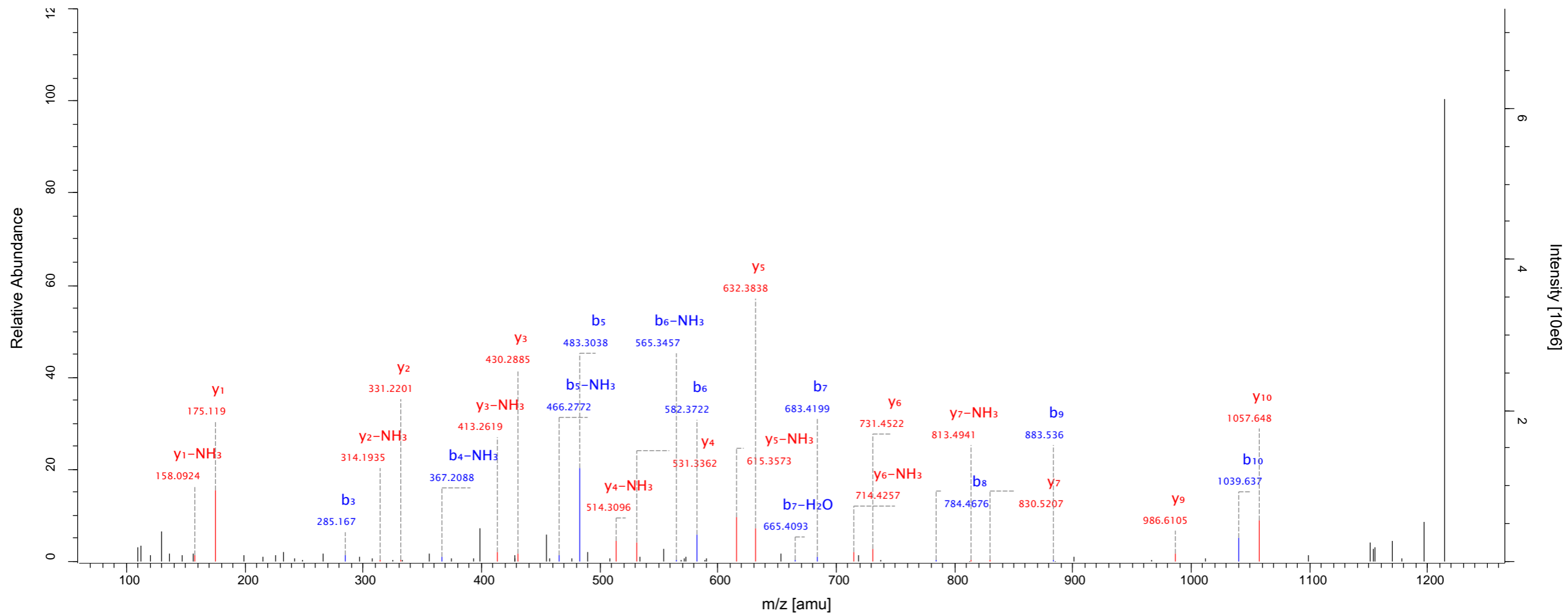
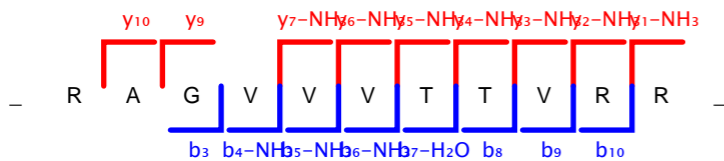
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_20
 Scan number: 6903
 Protein: ORFL114W + NC_006273.2[60151,60708]+ <unknown description>
 Peptide score: 160.67
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_20
Scan number: 7091
Protein: ORFL251W + NC_006273.2[161872,163748]+ <unknown description>; ORFL253W.iORF1 + NC_006273.2[161917,163748]+ <unknown description>
Peptide score: 72.65
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_20
 Scan number: 9061
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 145.55
 Method: FTMS; HCD



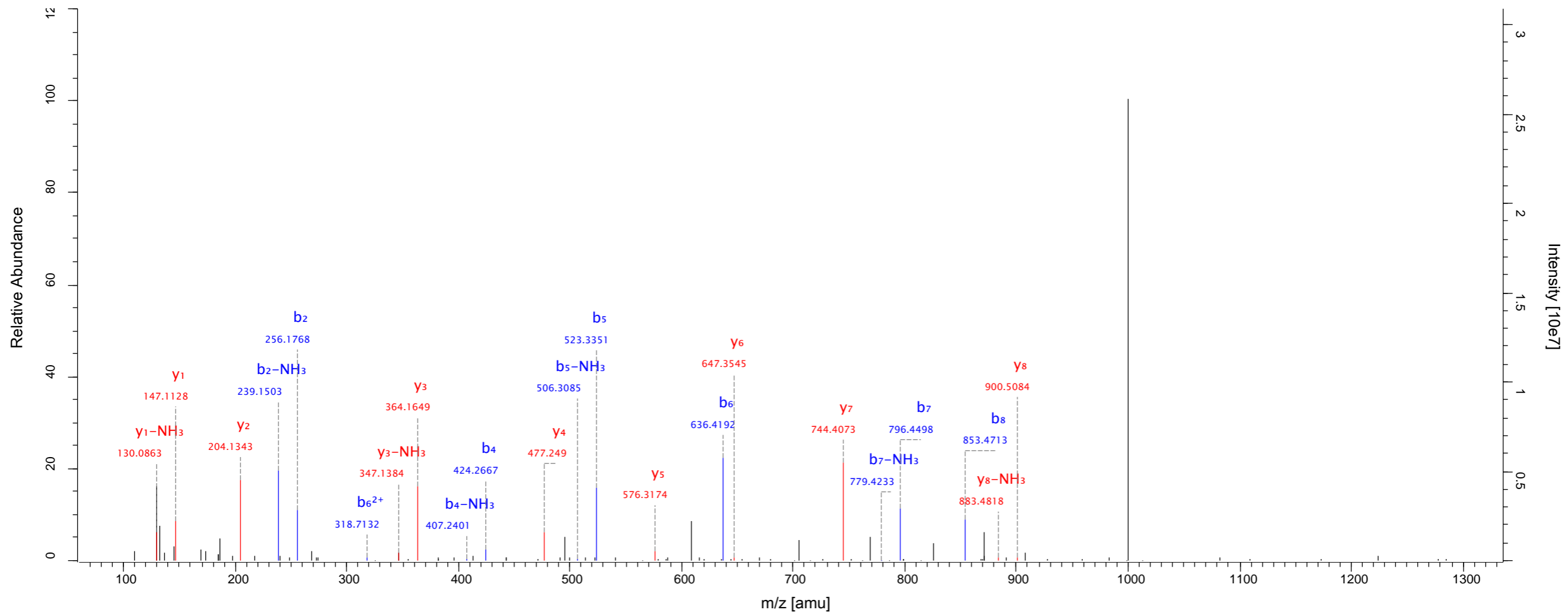
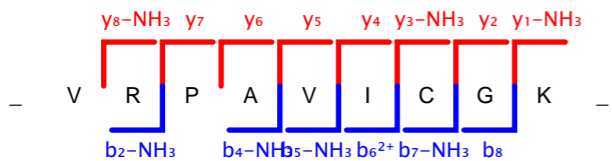
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_21

Scan number: 10367

Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 130.31

Method: FTMS; HCD



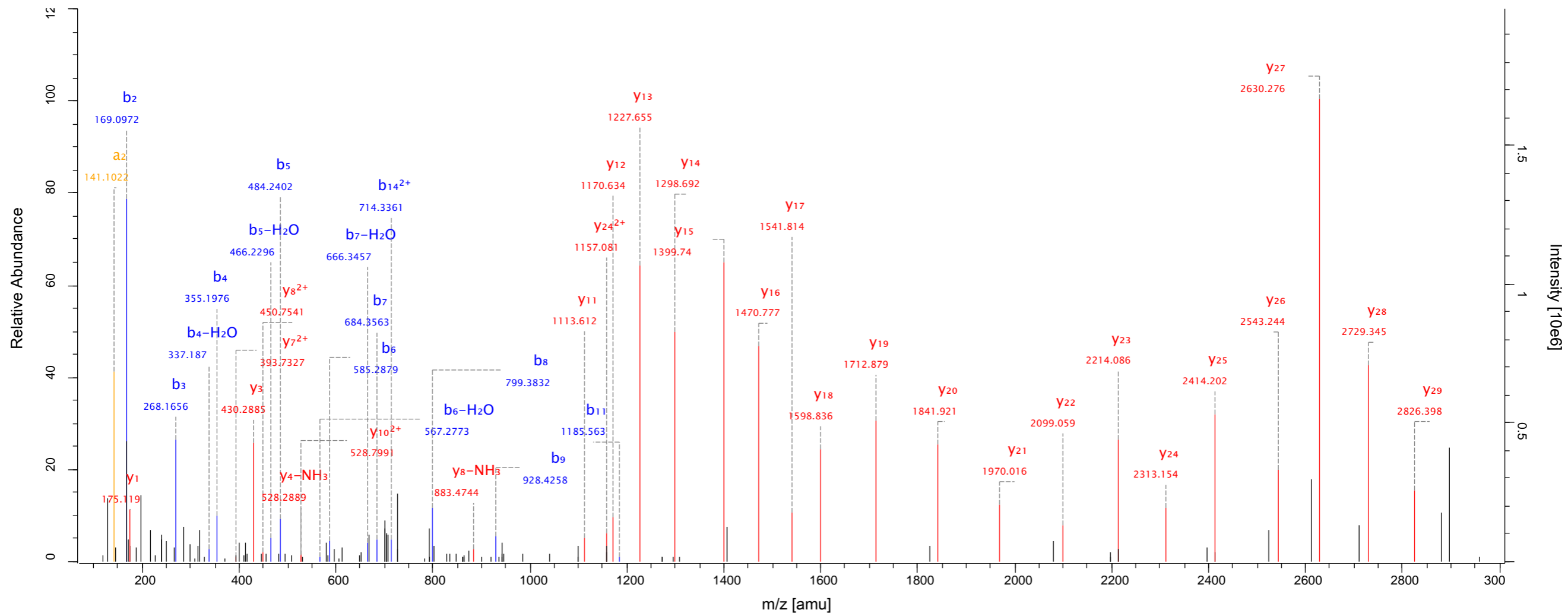
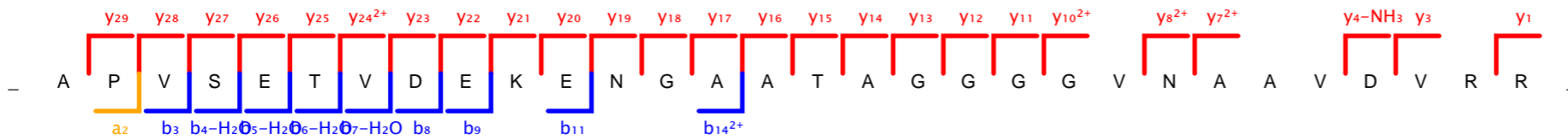
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_21

Scan number: 19969

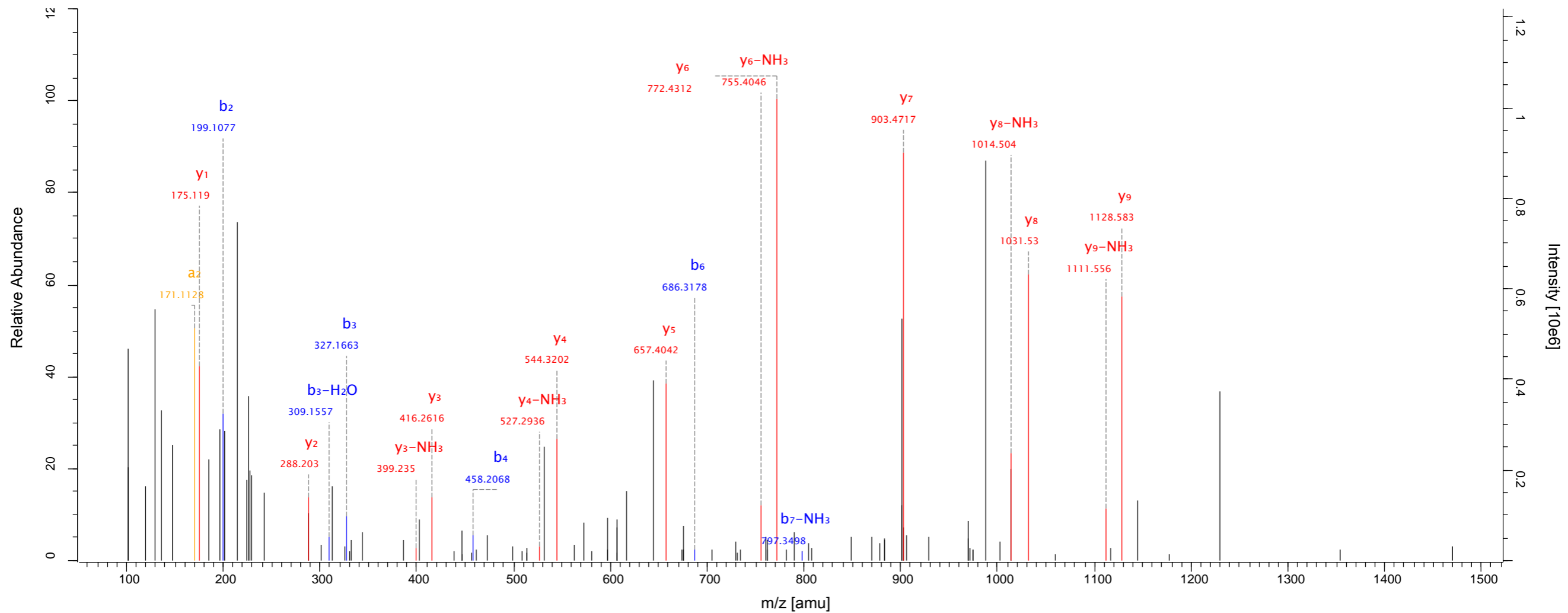
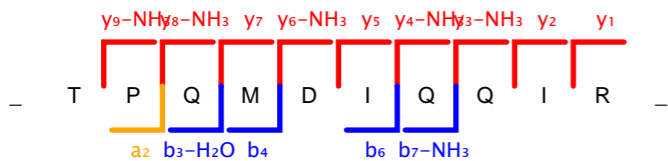
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 207.44

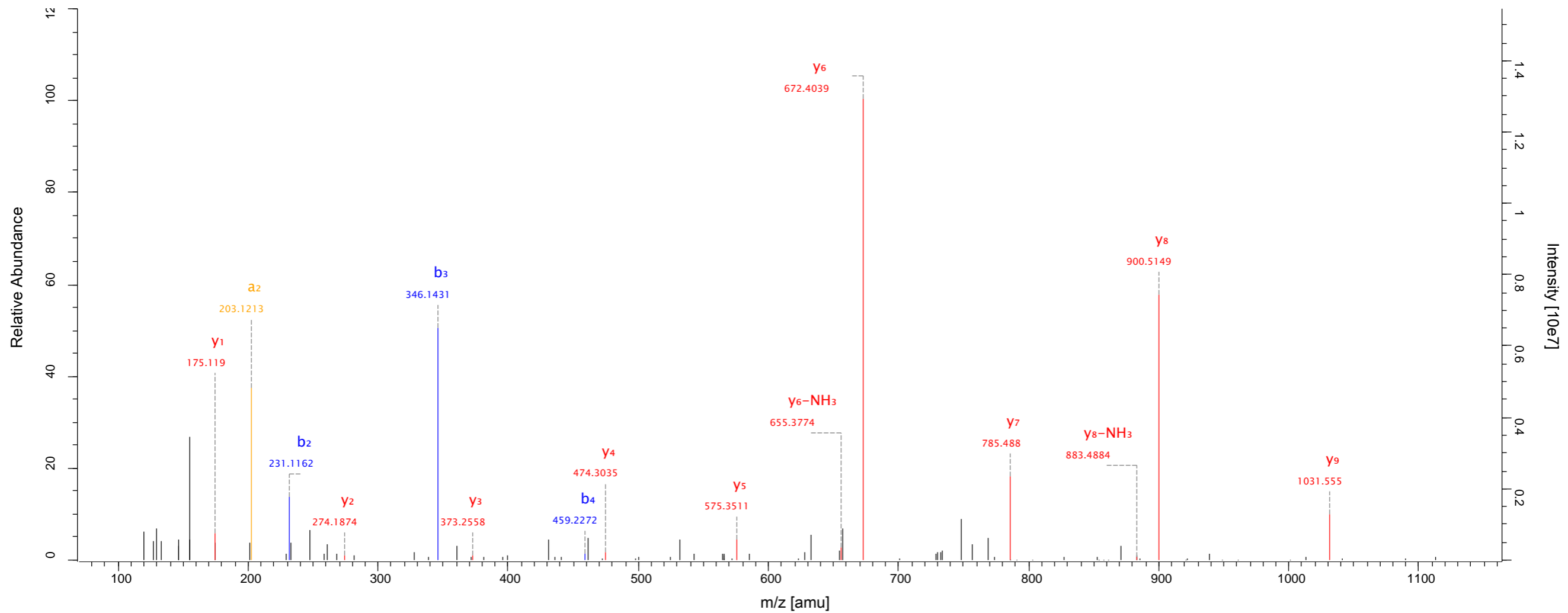
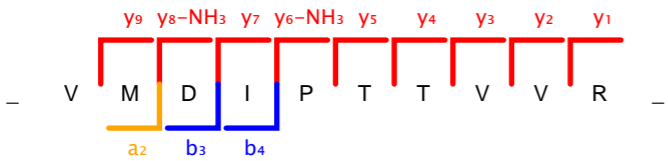
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_21
Scan number: 26548
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 121.5
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_21
Scan number: 32014
Protein: ORFL251W + NC_006273.2[161872,163748]+ <unknown description>
Peptide score: 89.05
Method: FTMS; HCD



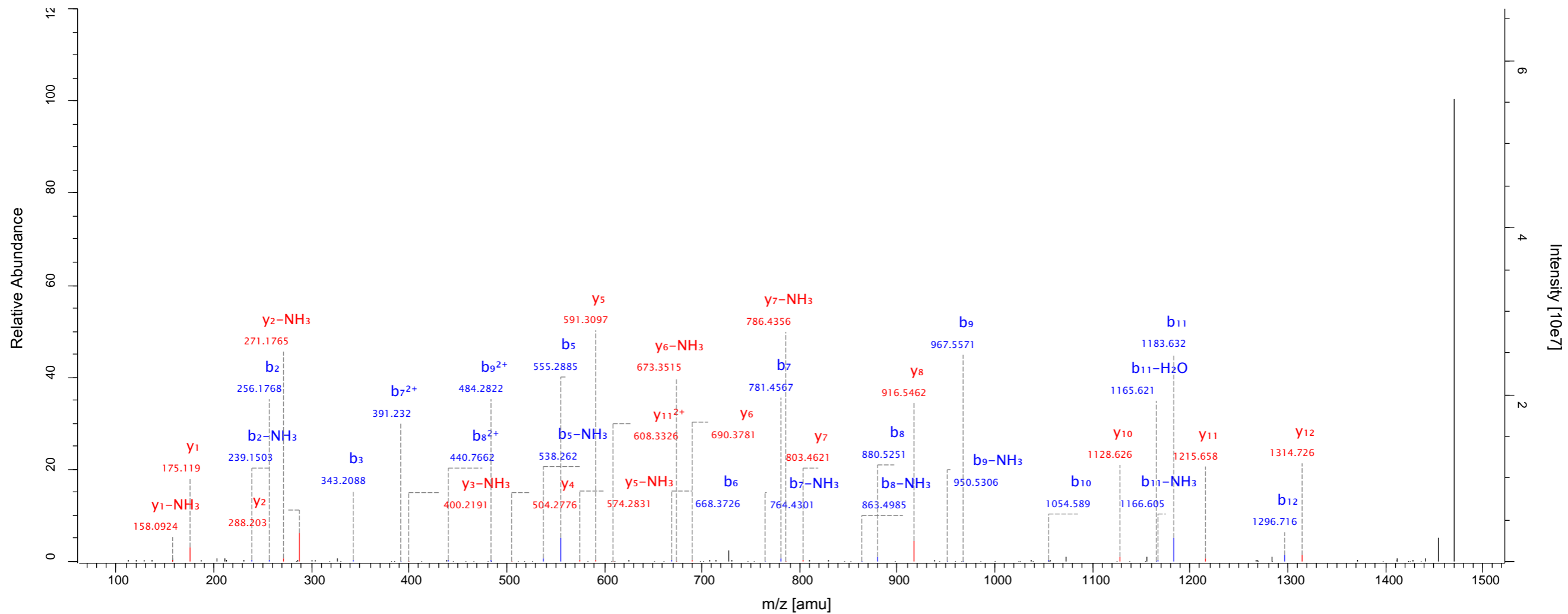
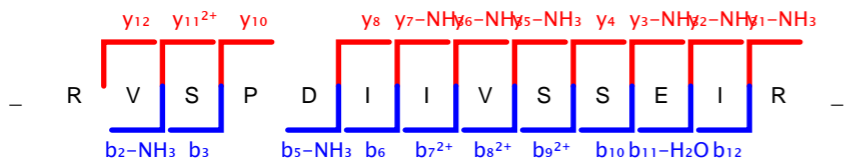
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_21

Scan number: 32634

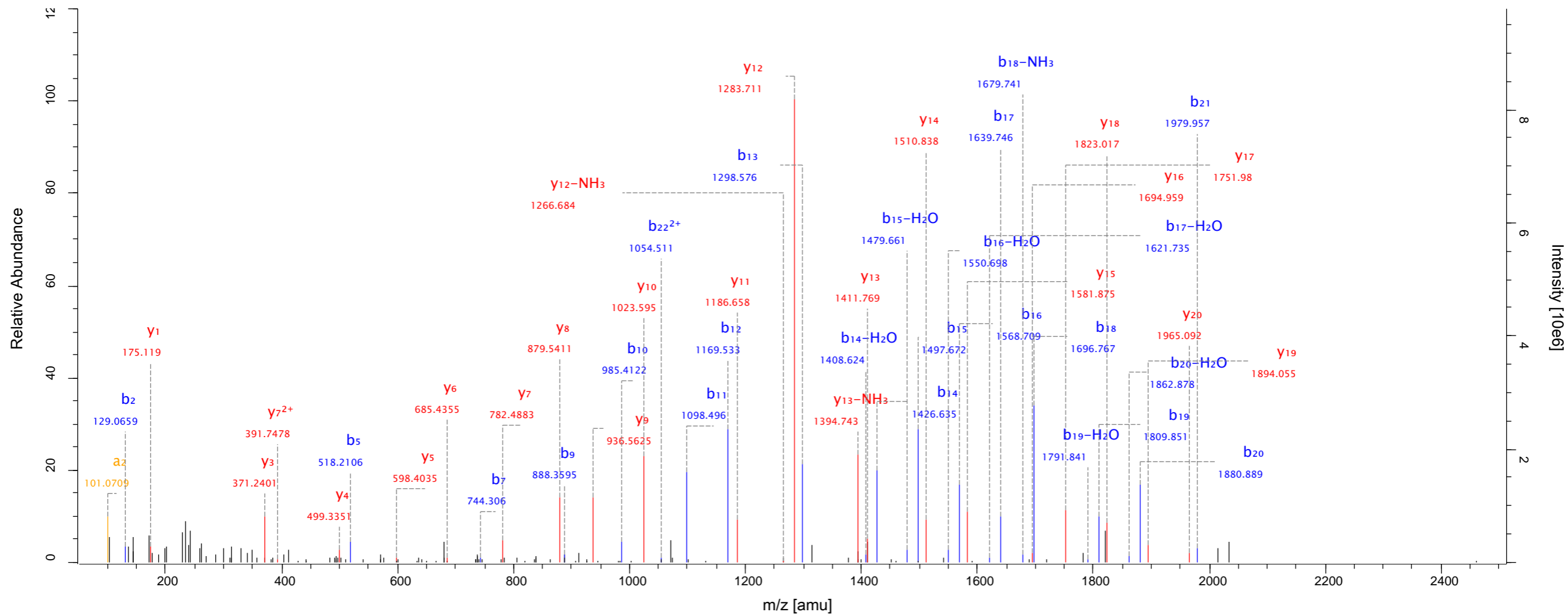
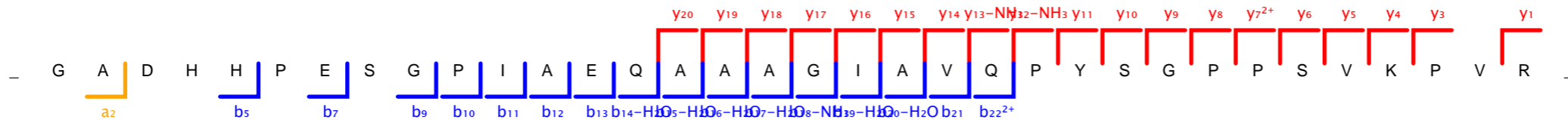
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 199.96

Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_21
 Scan number: 35256
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 193.11
 Method: FTMS; HCD



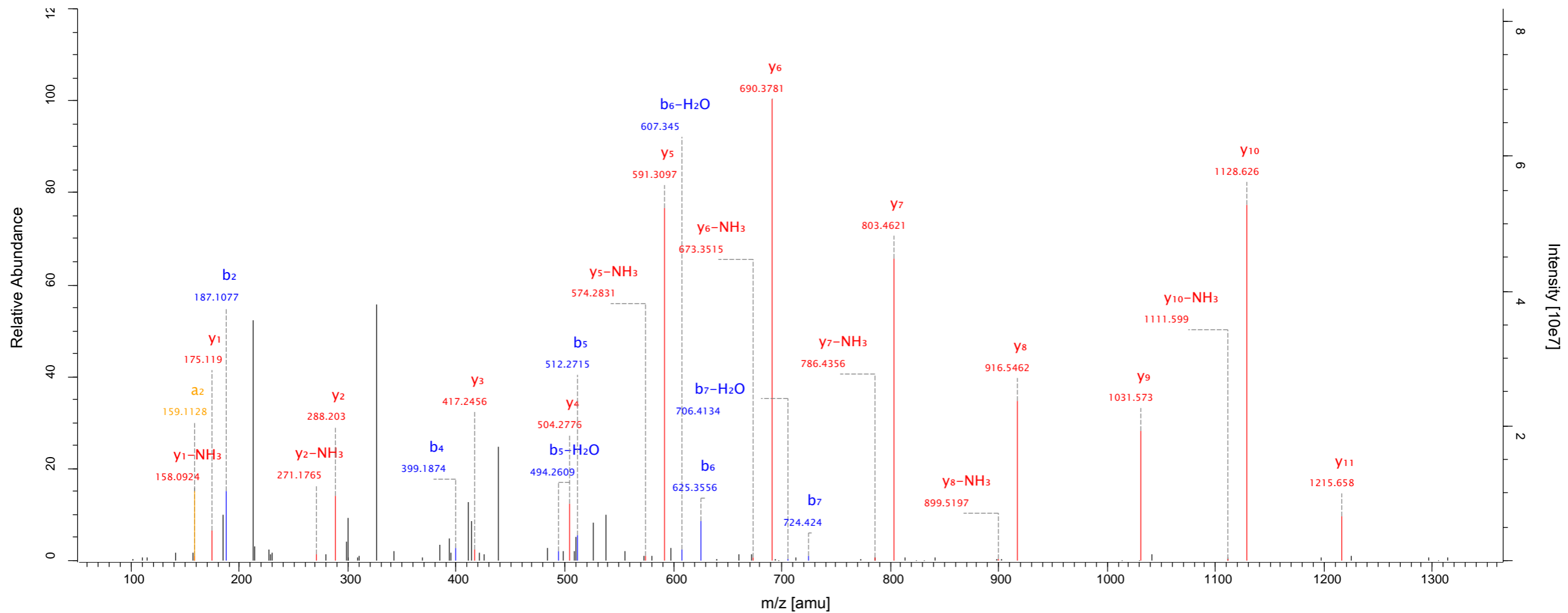
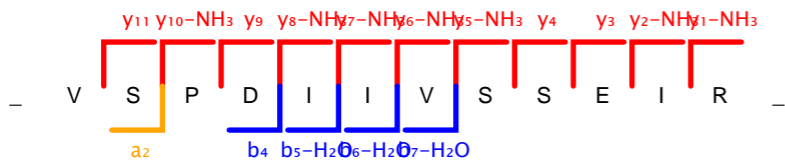
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_21

Scan number: 38690

Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 163.84

Method: FTMS; HCD



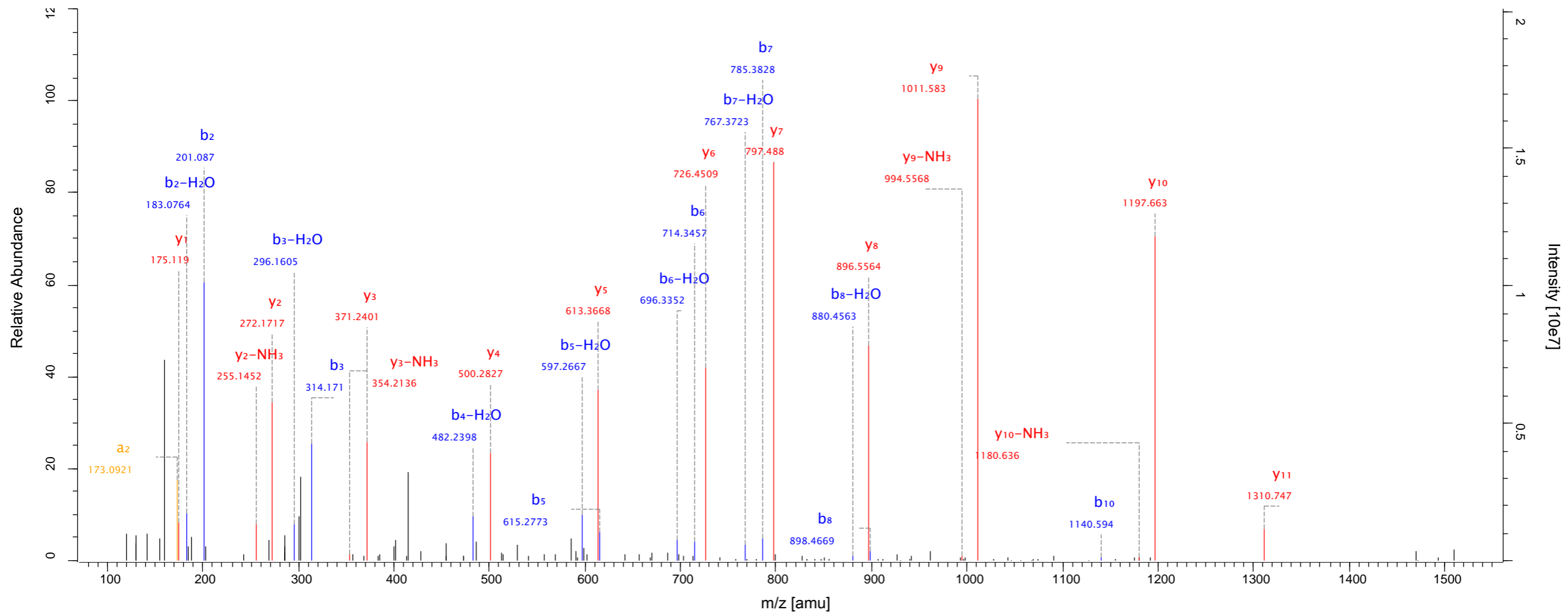
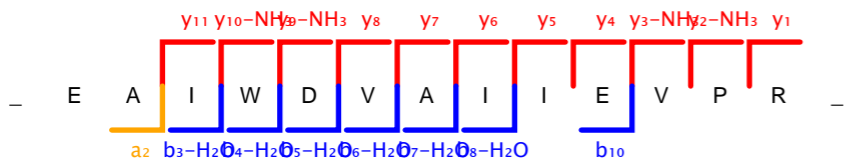
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_21

Scan number: 56117

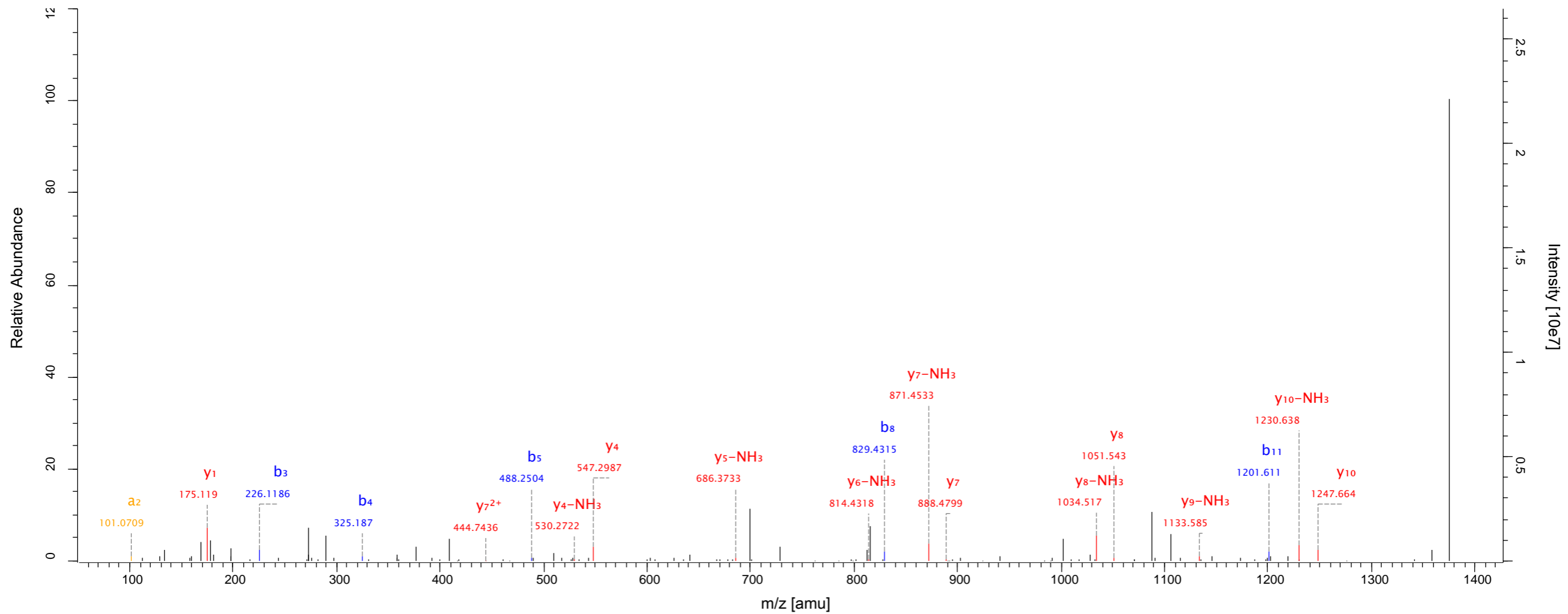
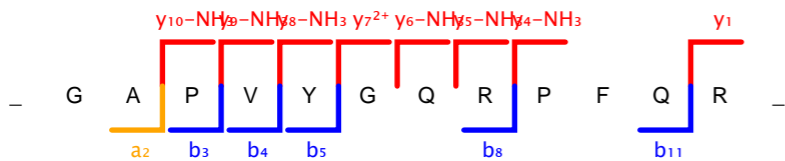
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 211.46

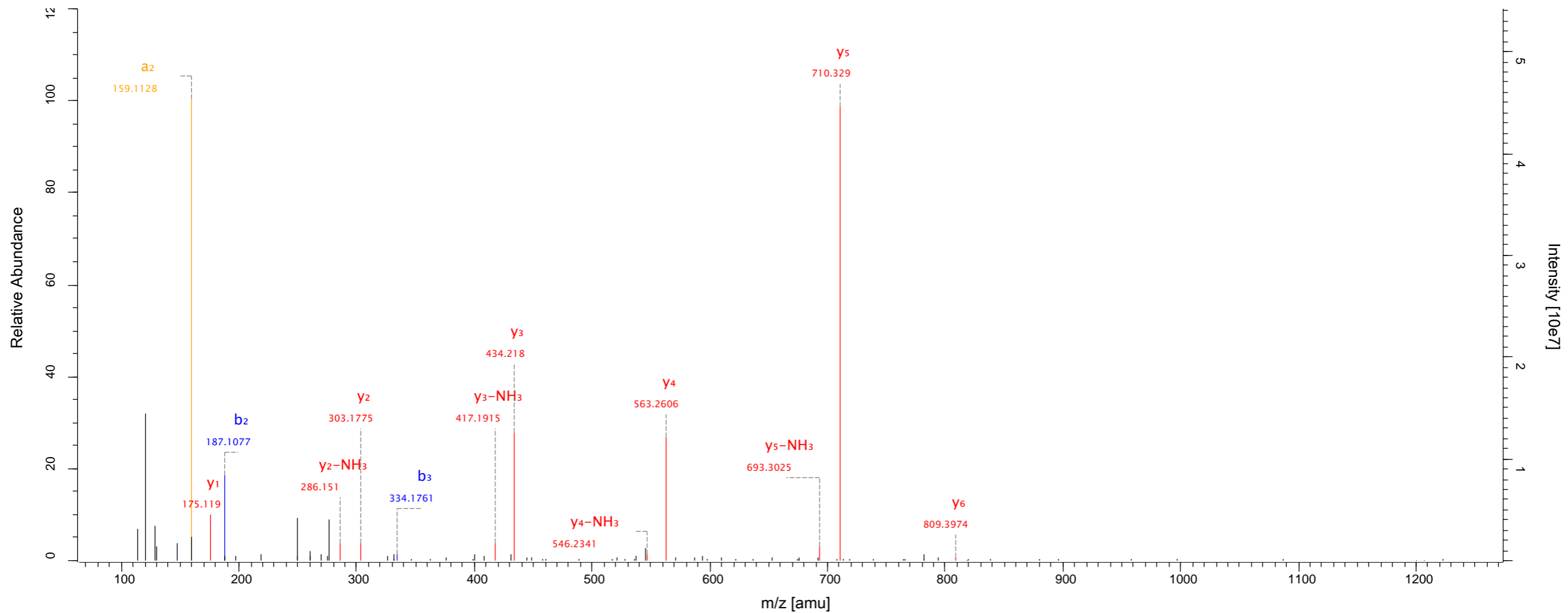
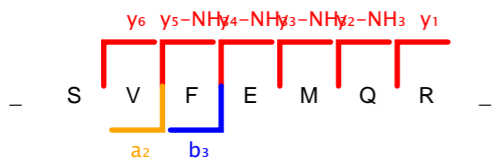
Method: FTMS; HCD



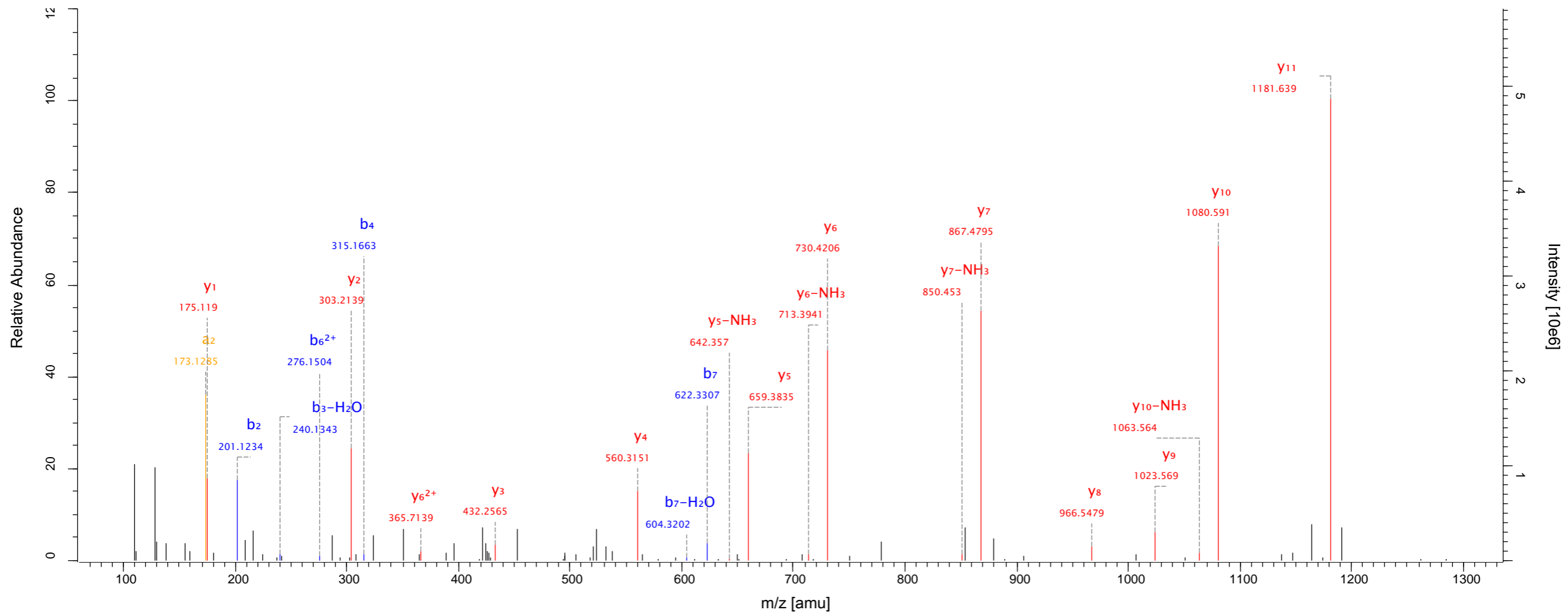
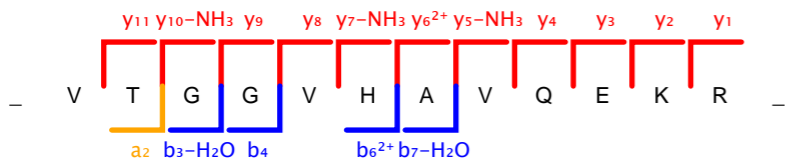
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_23
 Scan number: 16410
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 94.69
 Method: FTMS; HCD



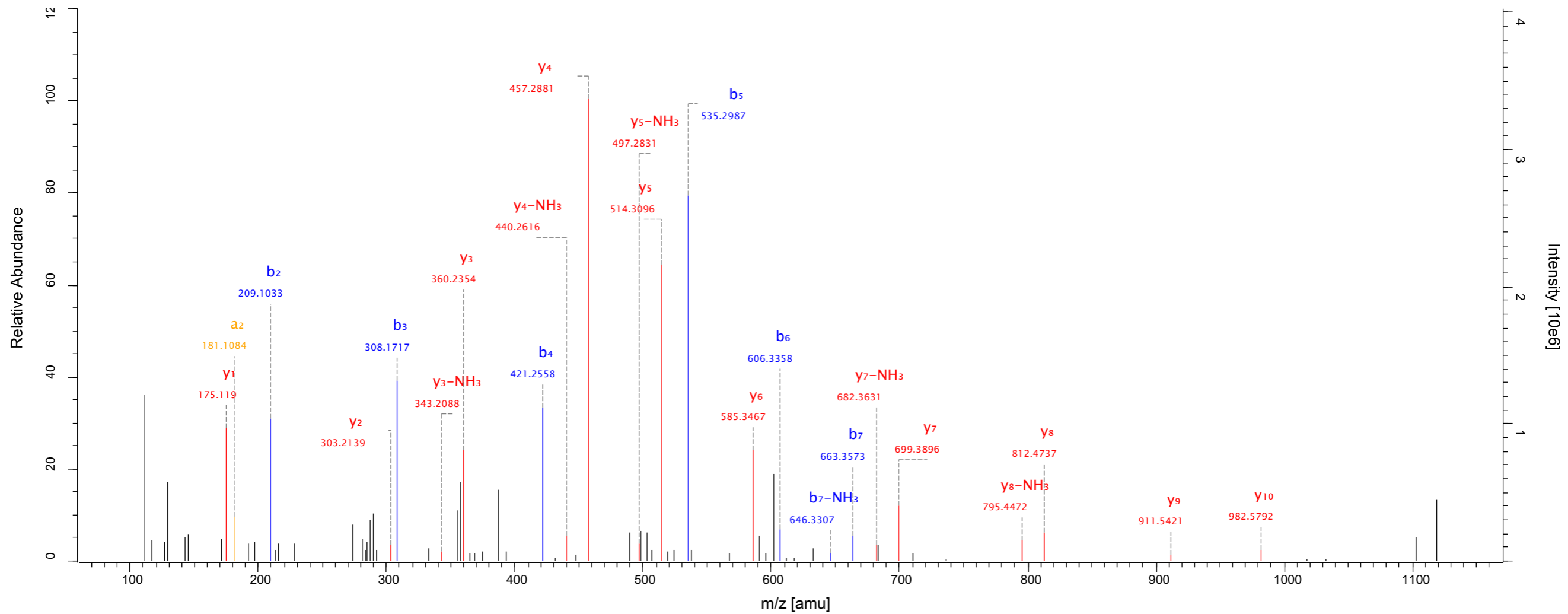
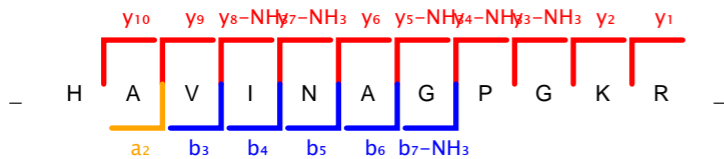
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_23
Scan number: 18985
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 142.11
Method: FTMS; HCD



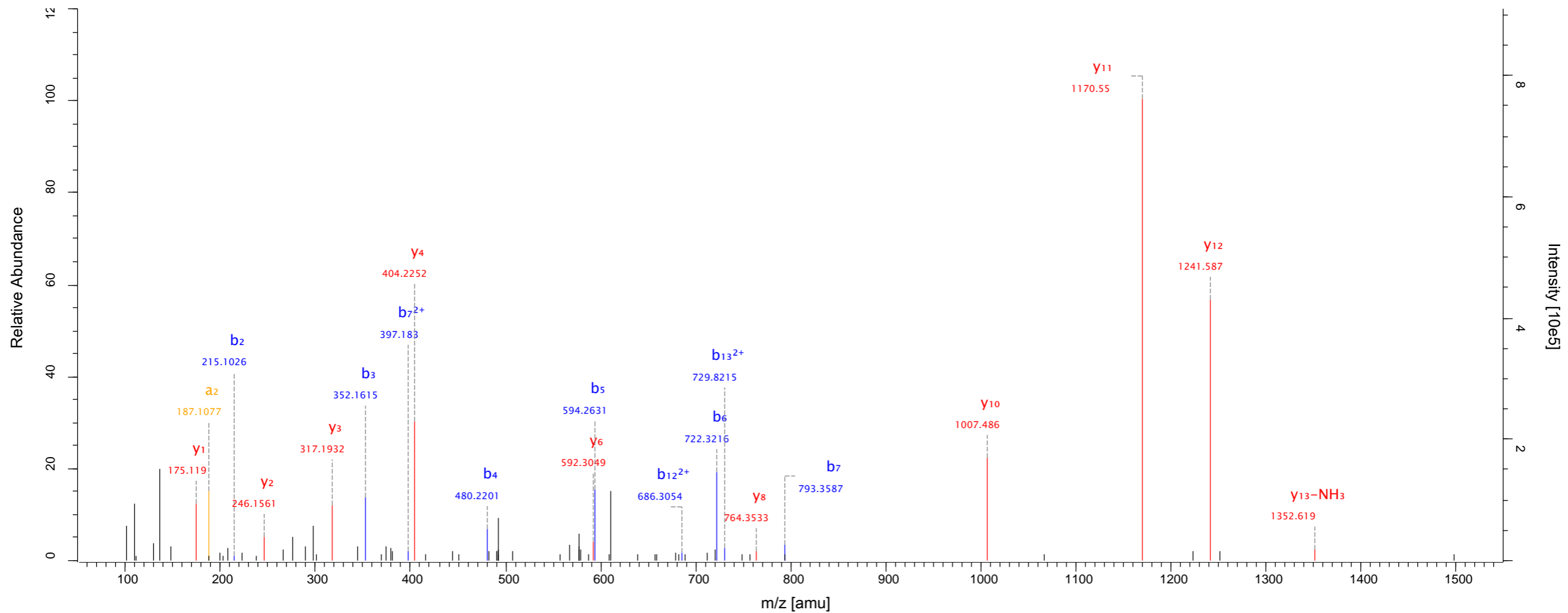
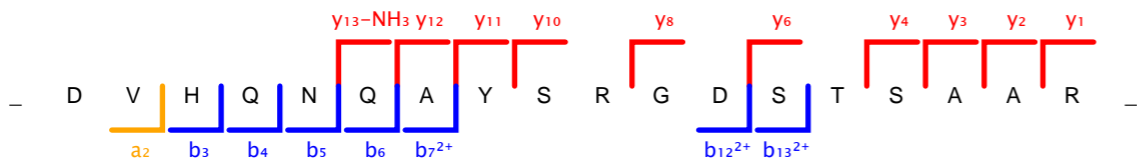
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_23
Scan number: 5525
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 122.69
Method: FTMS; HCD



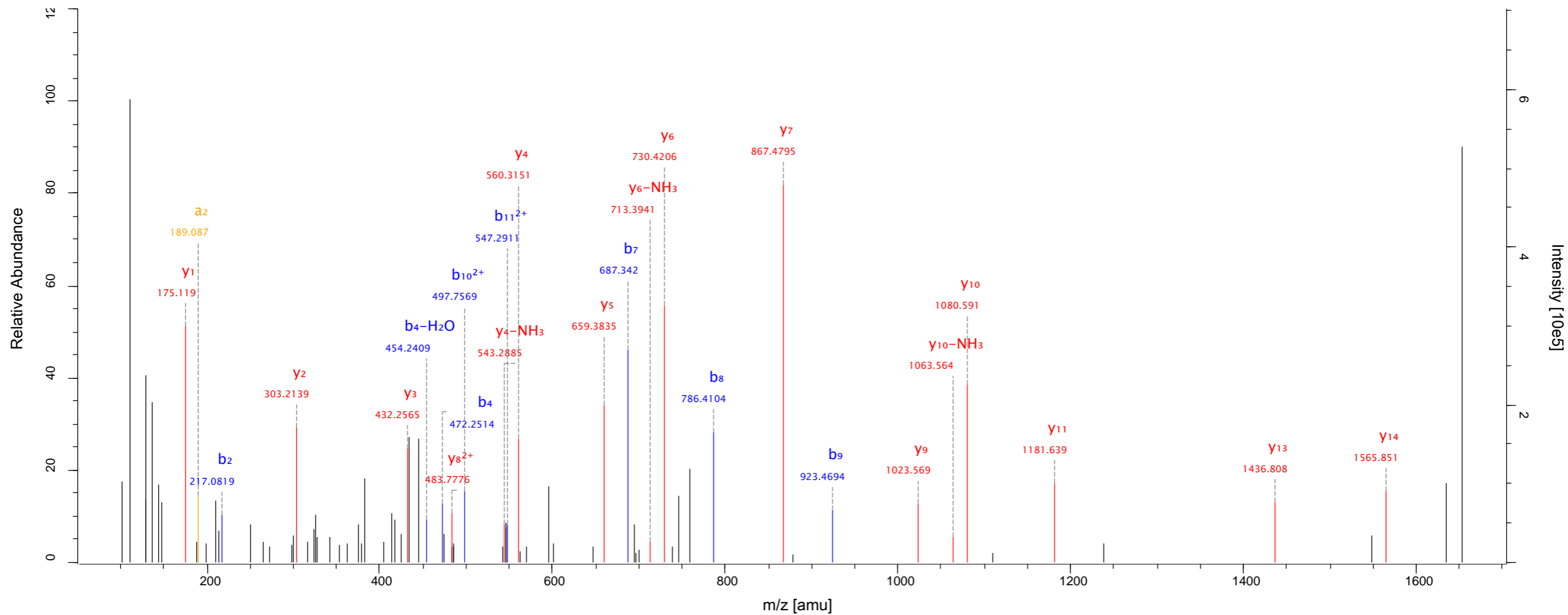
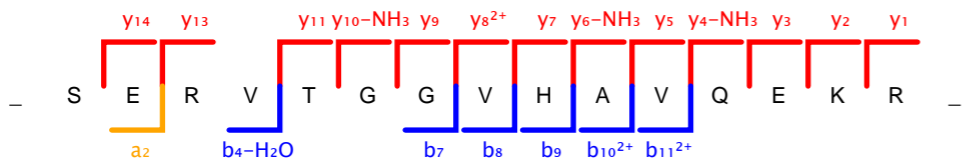
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_23
Scan number: 6153
Protein: ORFL87W + NC_006273.2[38099,40114]+ <unknown description>
Peptide score: 159.4
Method: FTMS; HCD



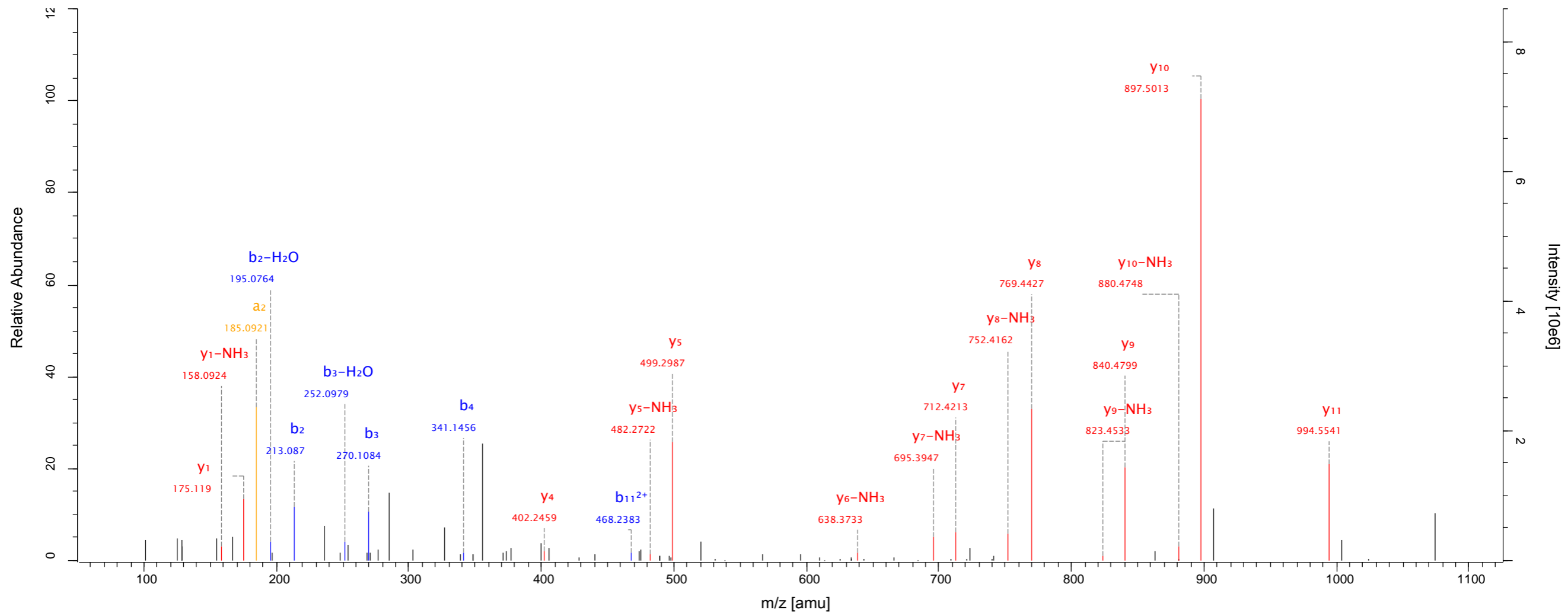
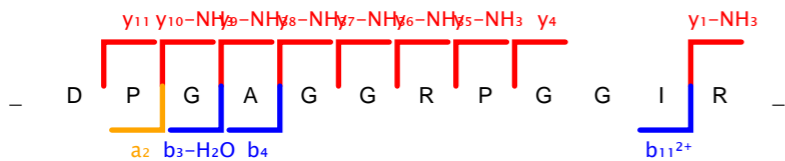
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_23
Scan number: 6232
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 83.42
Method: FTMS; HCD



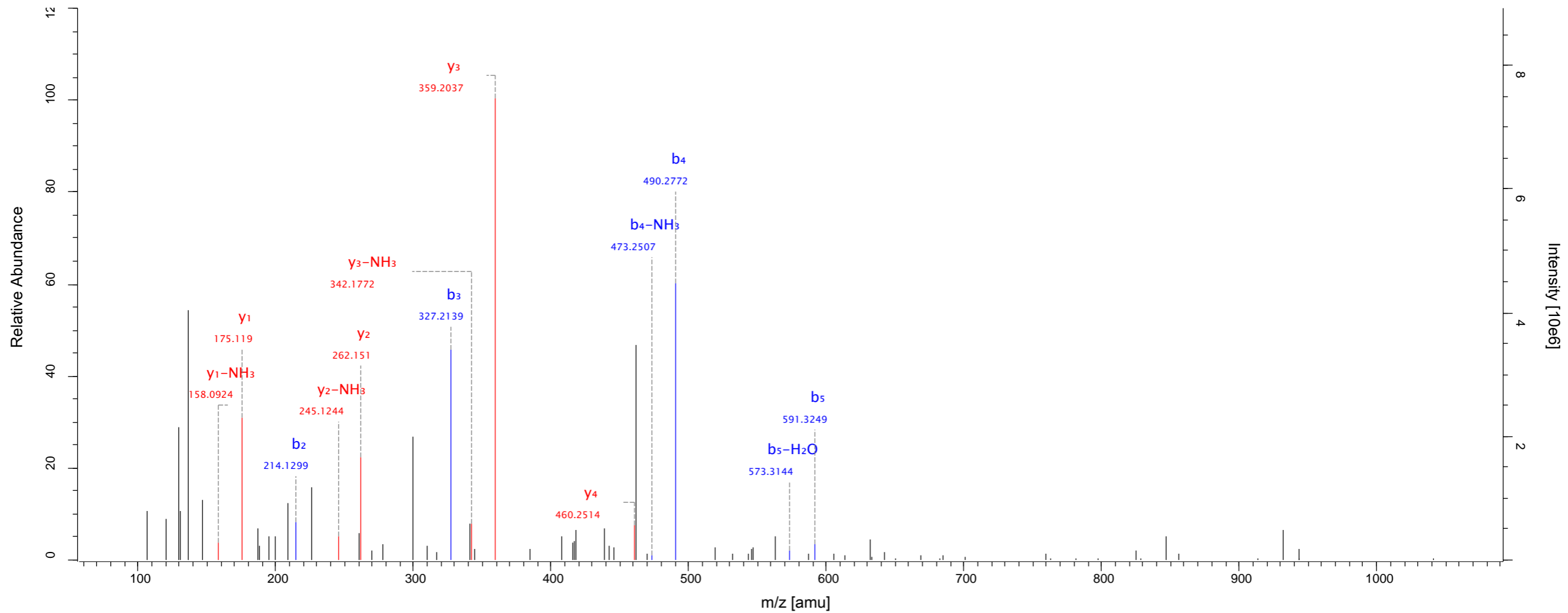
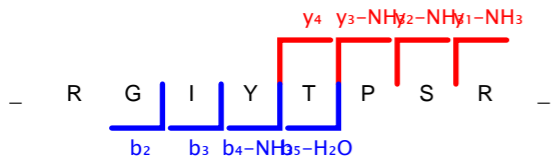
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_23
 Scan number: 7014
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 131.35
 Method: FTMS; HCD



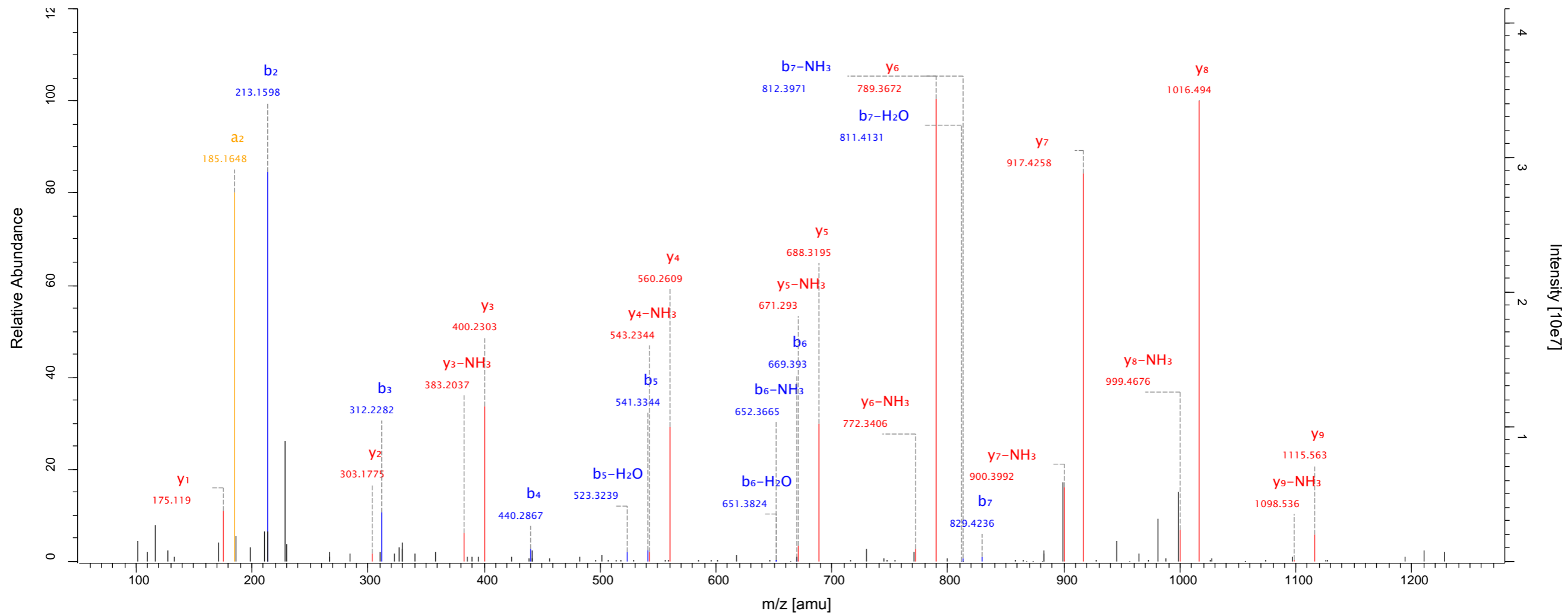
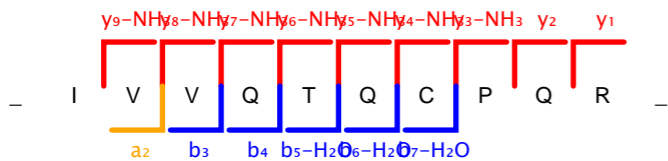
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_23
 Scan number: 7564
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 114.89
 Method: FTMS; HCD



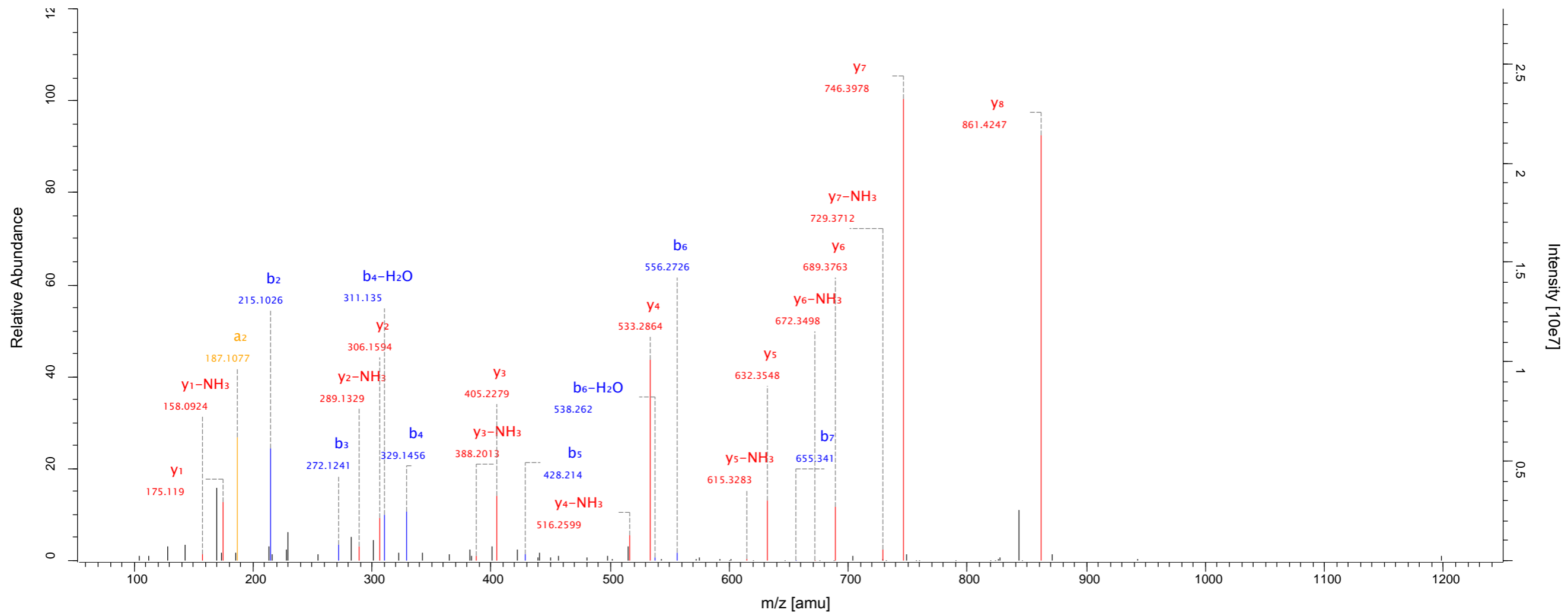
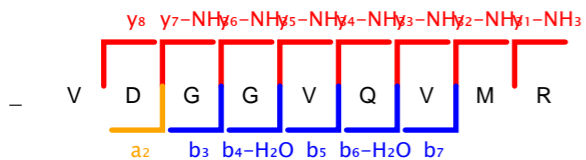
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_23
Scan number: 9504
Protein: ORFL87W + NC_006273.2[38099,40114]+ <unknown description>
Peptide score: 67.17
Method: FTMS; HCD



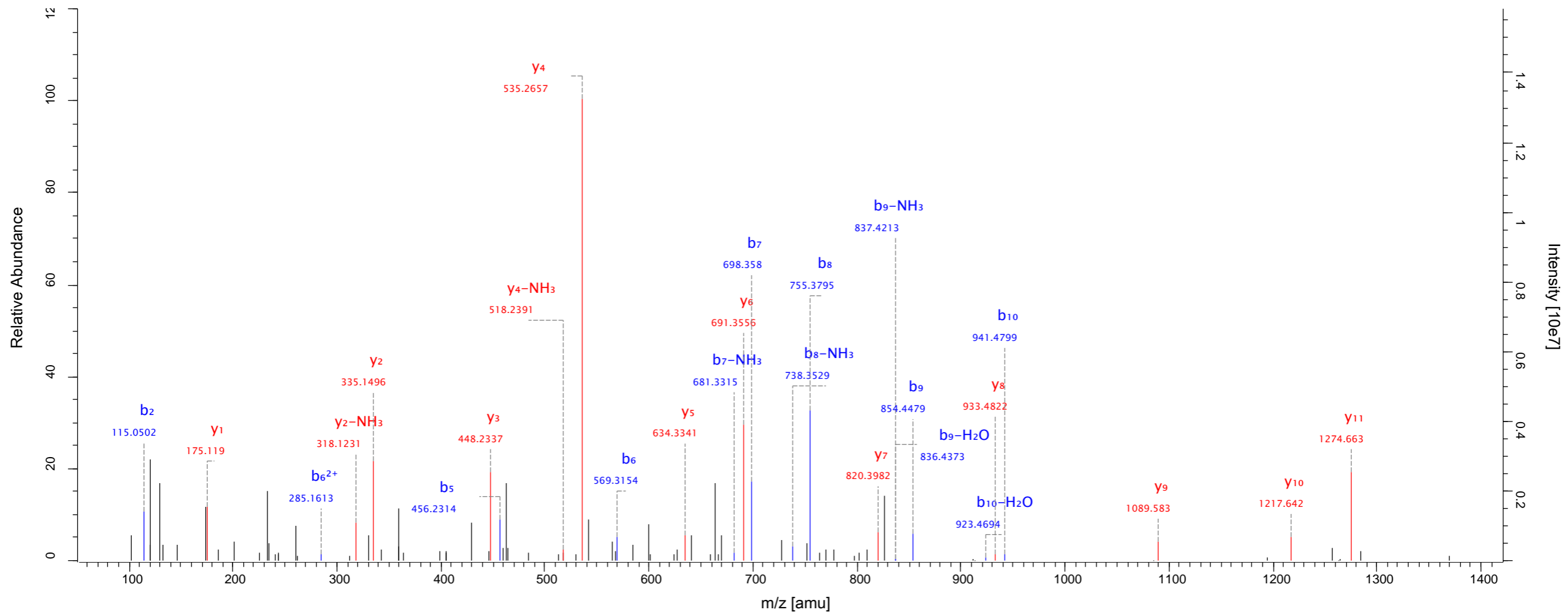
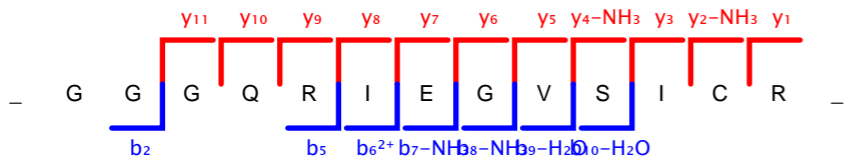
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
 Scan number: 10699
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 208.25
 Method: FTMS; HCD



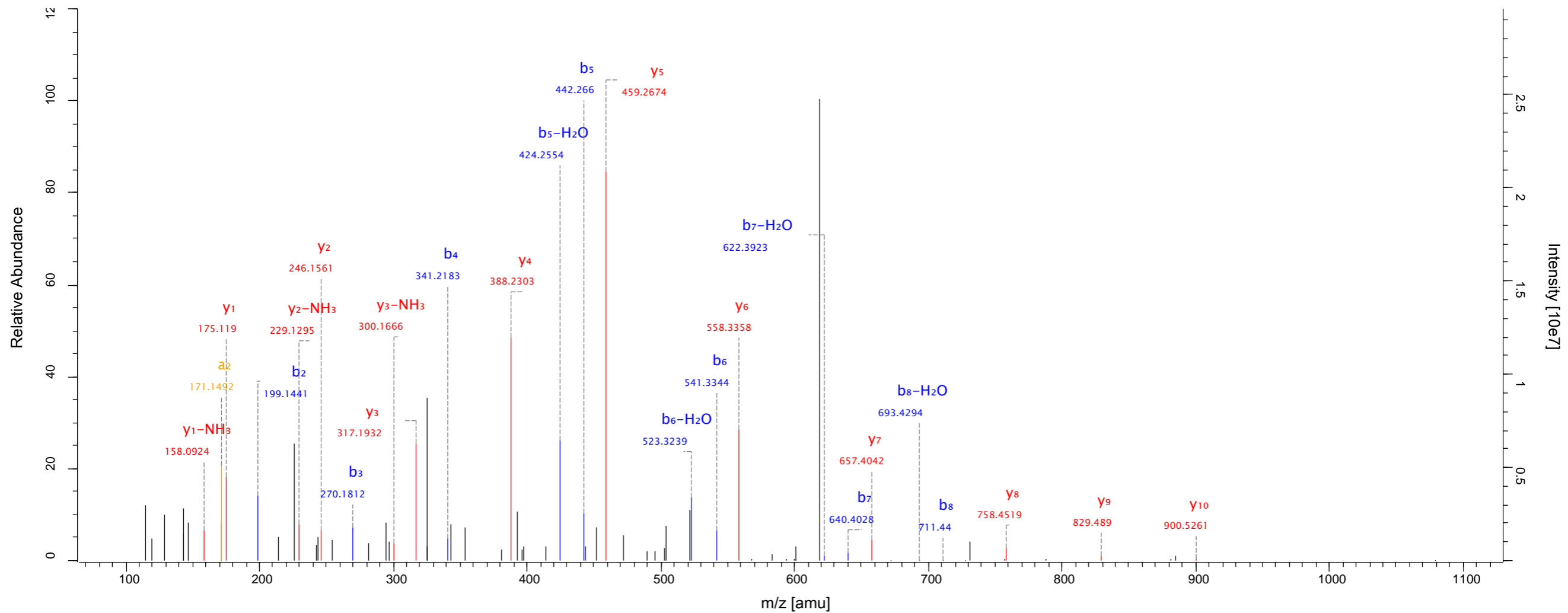
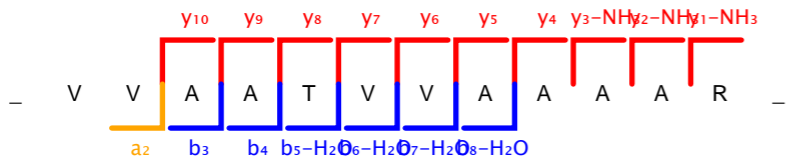
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
 Scan number: 11917
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 168.26
 Method: FTMS; HCD



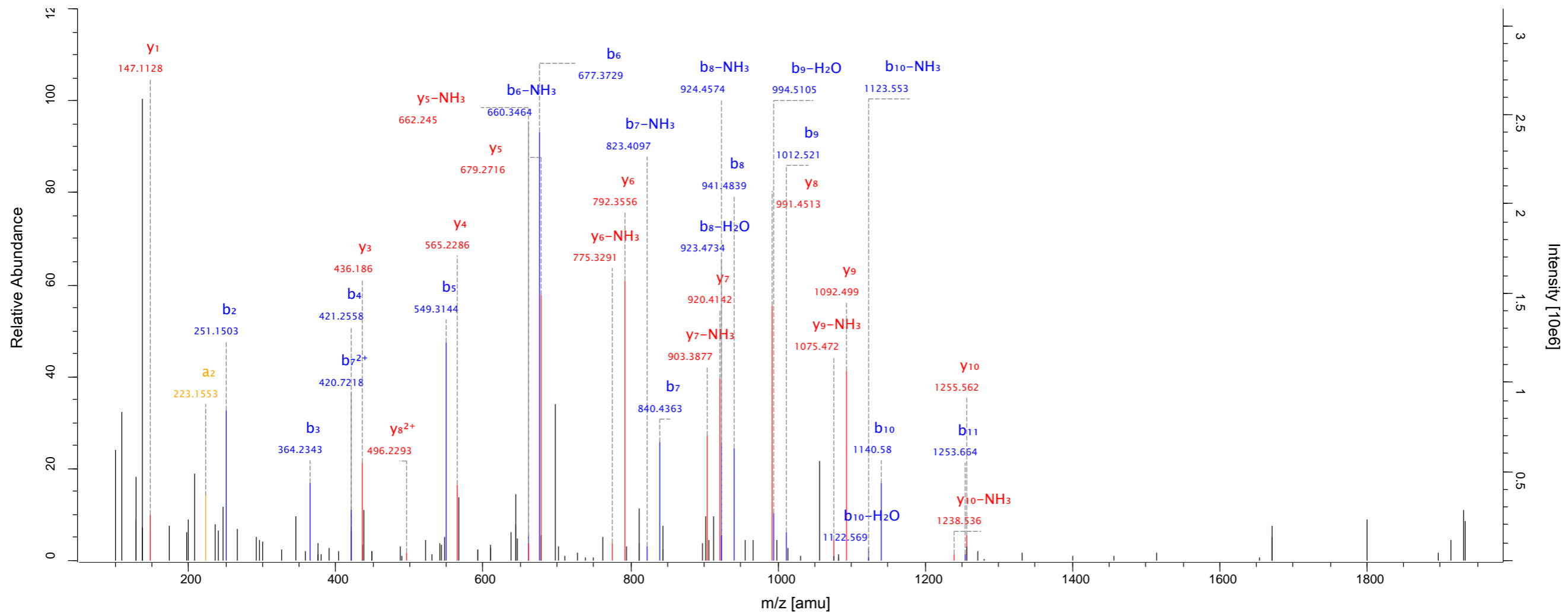
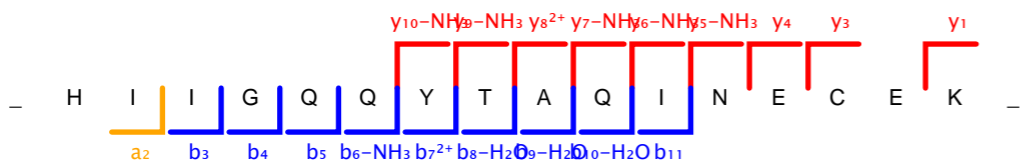
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
 Scan number: 14877
 Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
 Peptide score: 142.91
 Method: FTMS; HCD



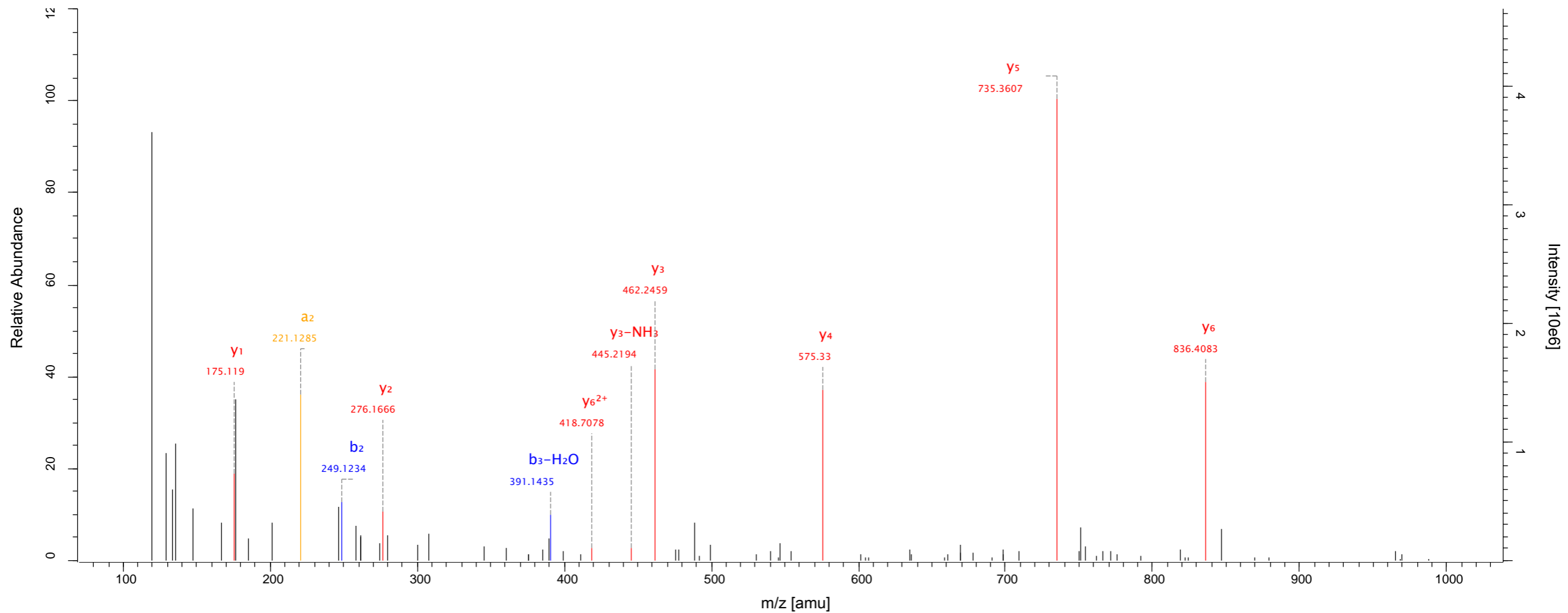
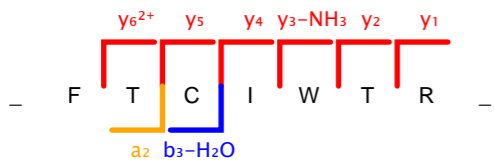
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
 Scan number: 20296
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 138.25
 Method: FTMS; HCD



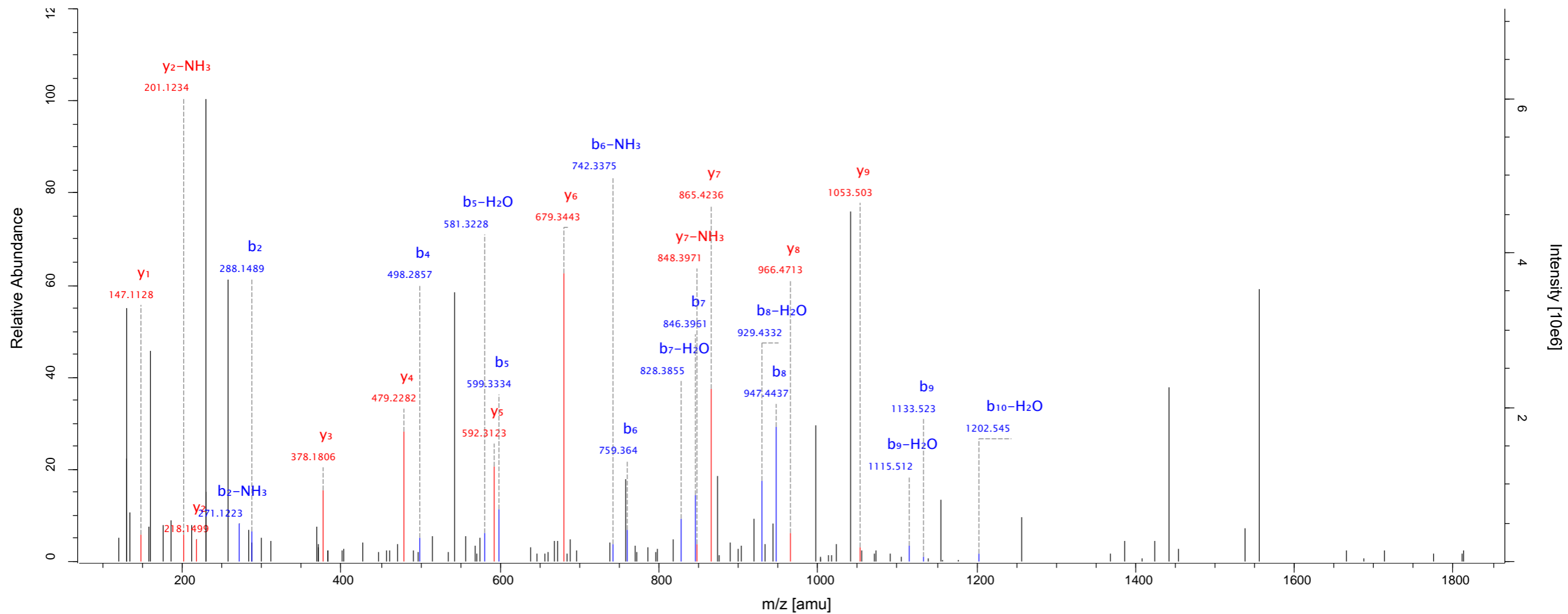
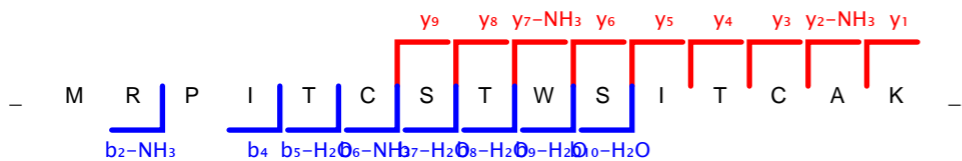
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_3
 Scan number: 21348
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 181.76
 Method: FTMS; HCD



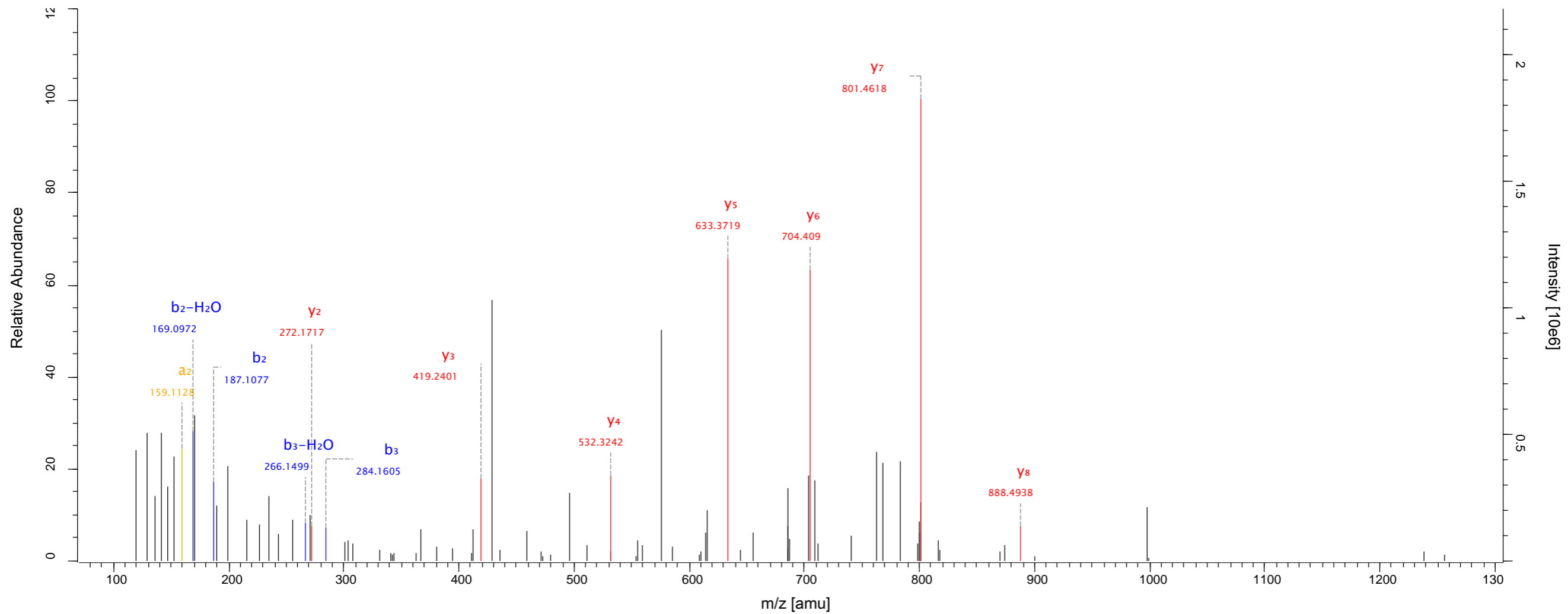
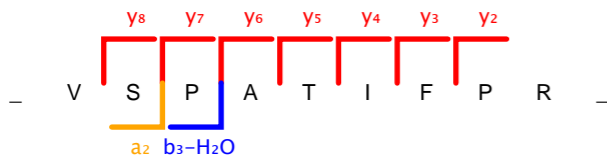
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
Scan number: 25112
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 76.85
Method: FTMS; HCD



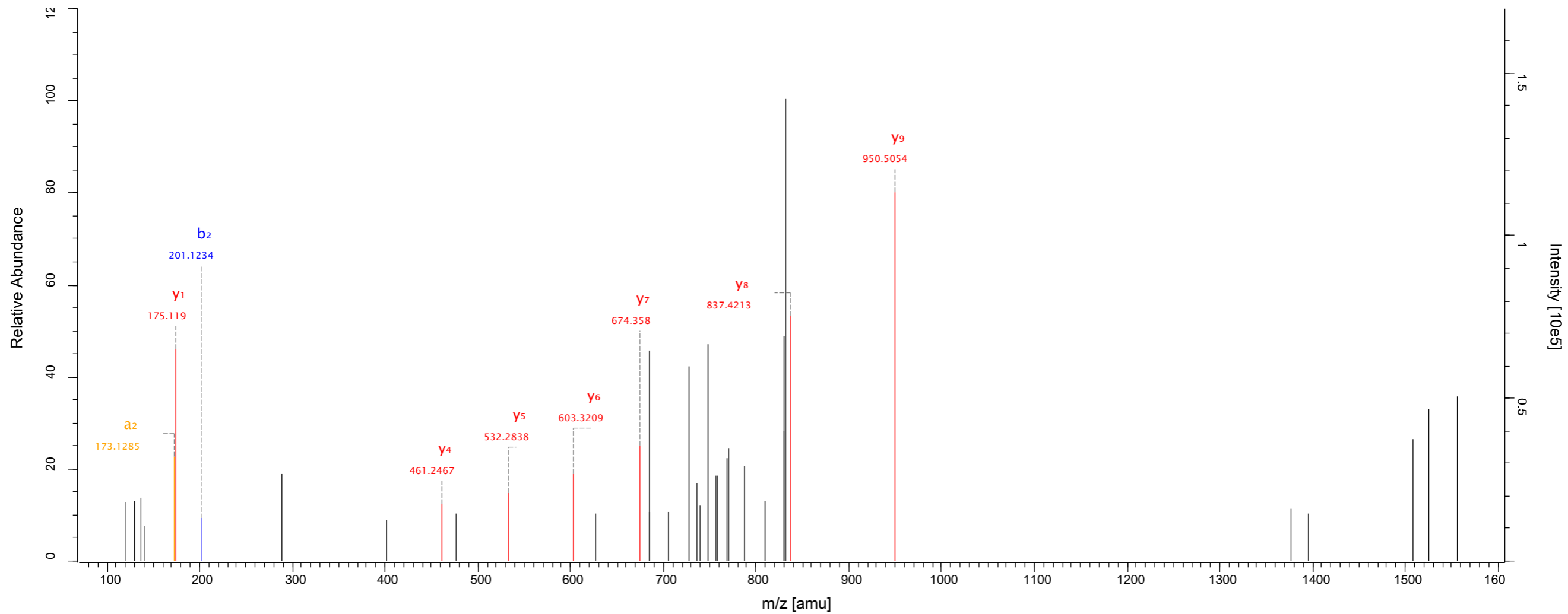
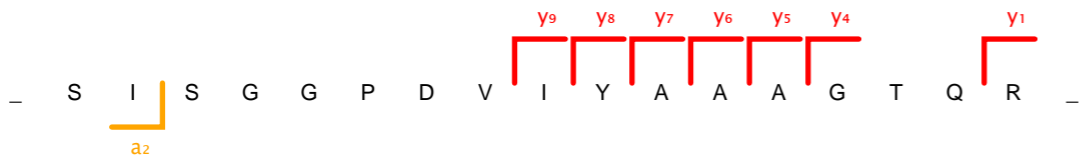
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_3
 Scan number: 29132
 Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
 Peptide score: 109.83
 Method: FTMS; HCD



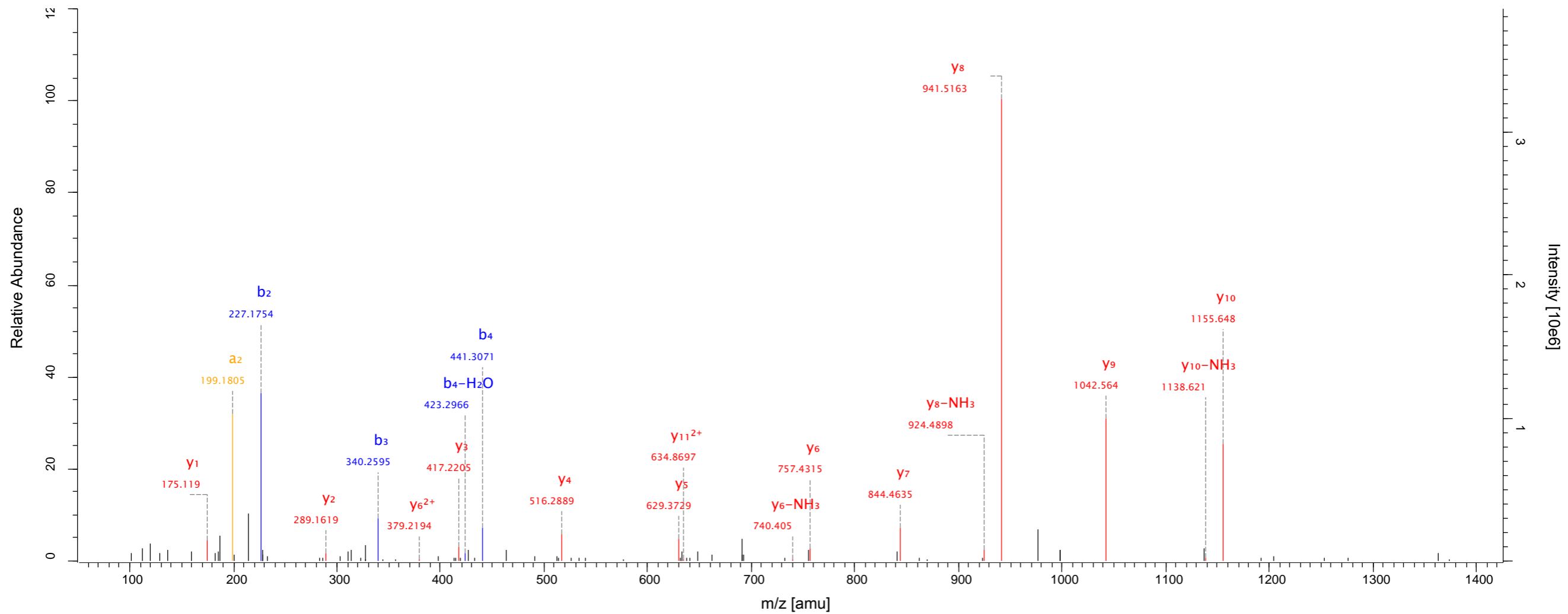
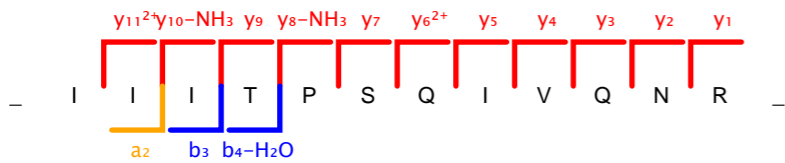
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
Scan number: 29169
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 62.58
Method: FTMS; HCD



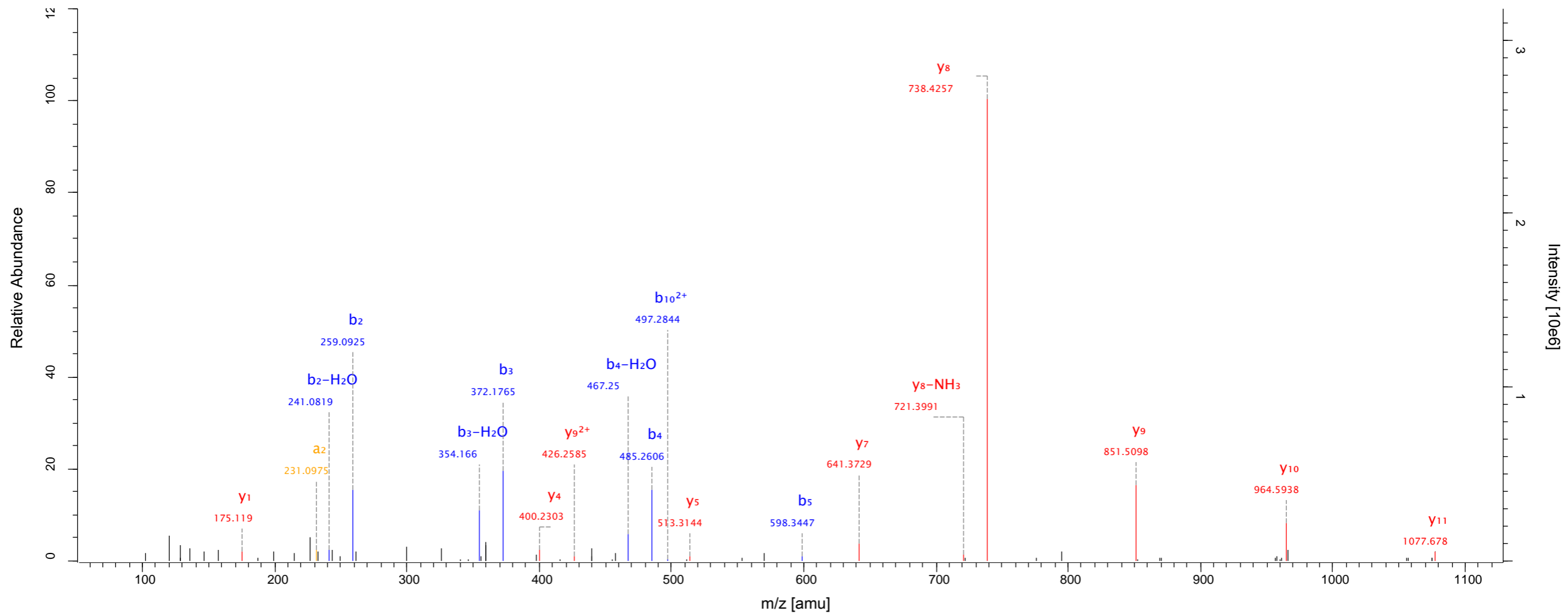
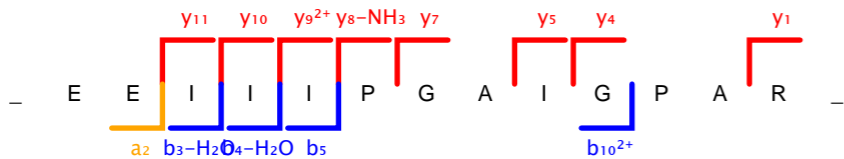
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
Scan number: 29871
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>
Peptide score: 44.81
Method: FTMS; HCD



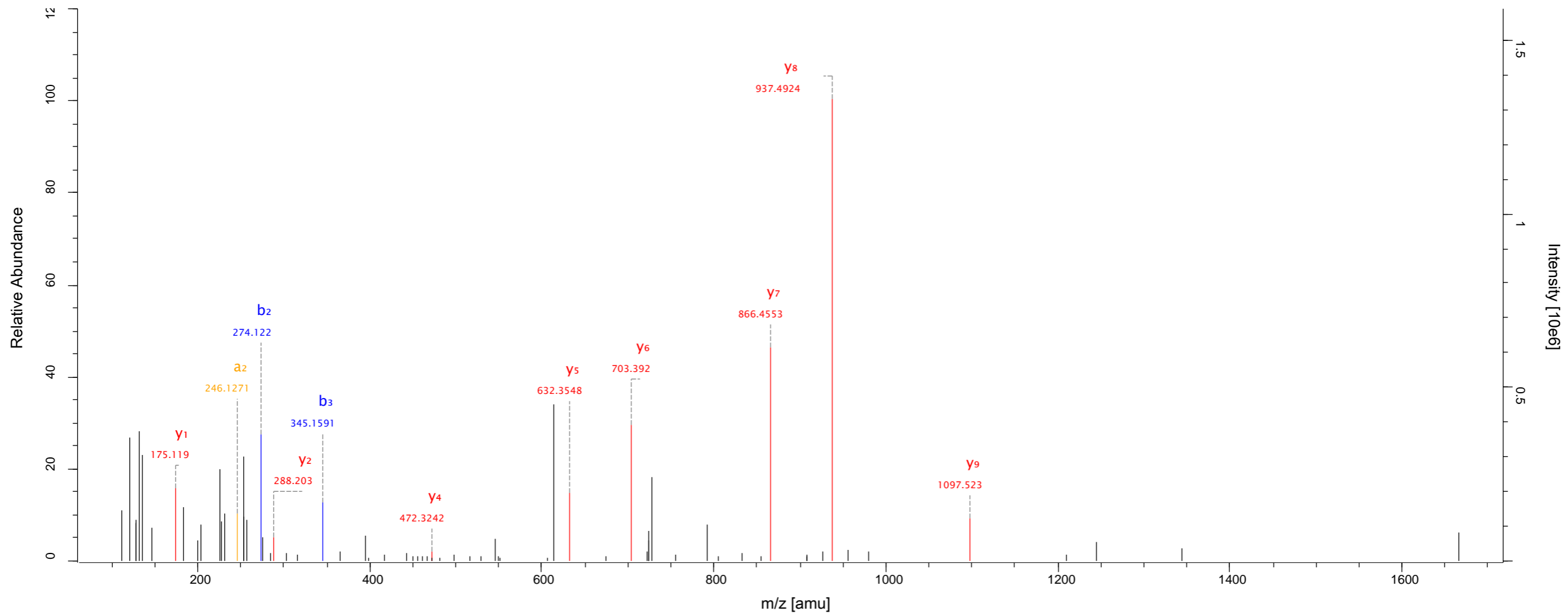
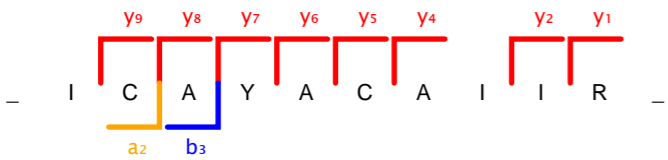
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
Scan number: 33826
Protein: ORFL190C - NC_006273.2[111965,112384]- <unknown description>
Peptide score: 114.83
Method: FTMS; HCD



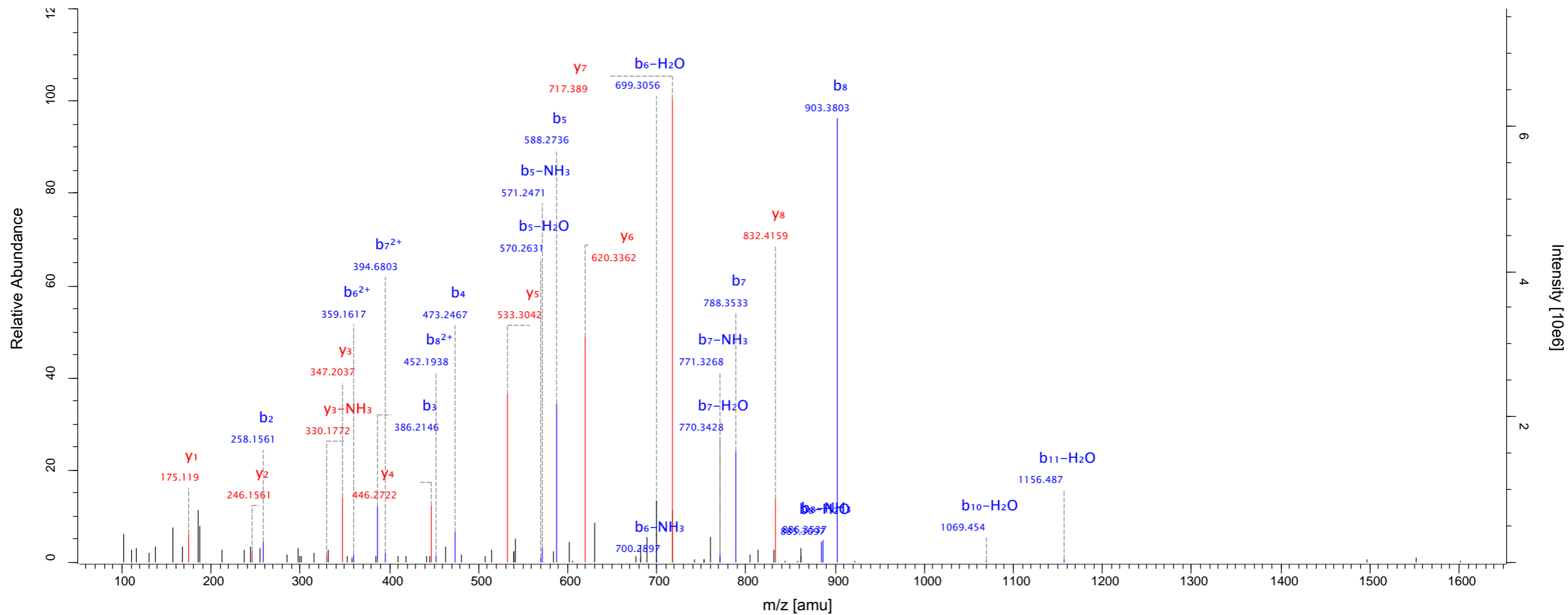
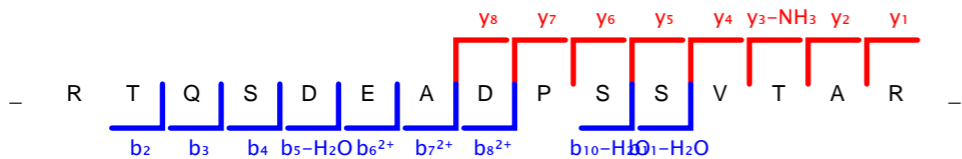
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
Scan number: 36993
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 97.91
Method: FTMS; HCD



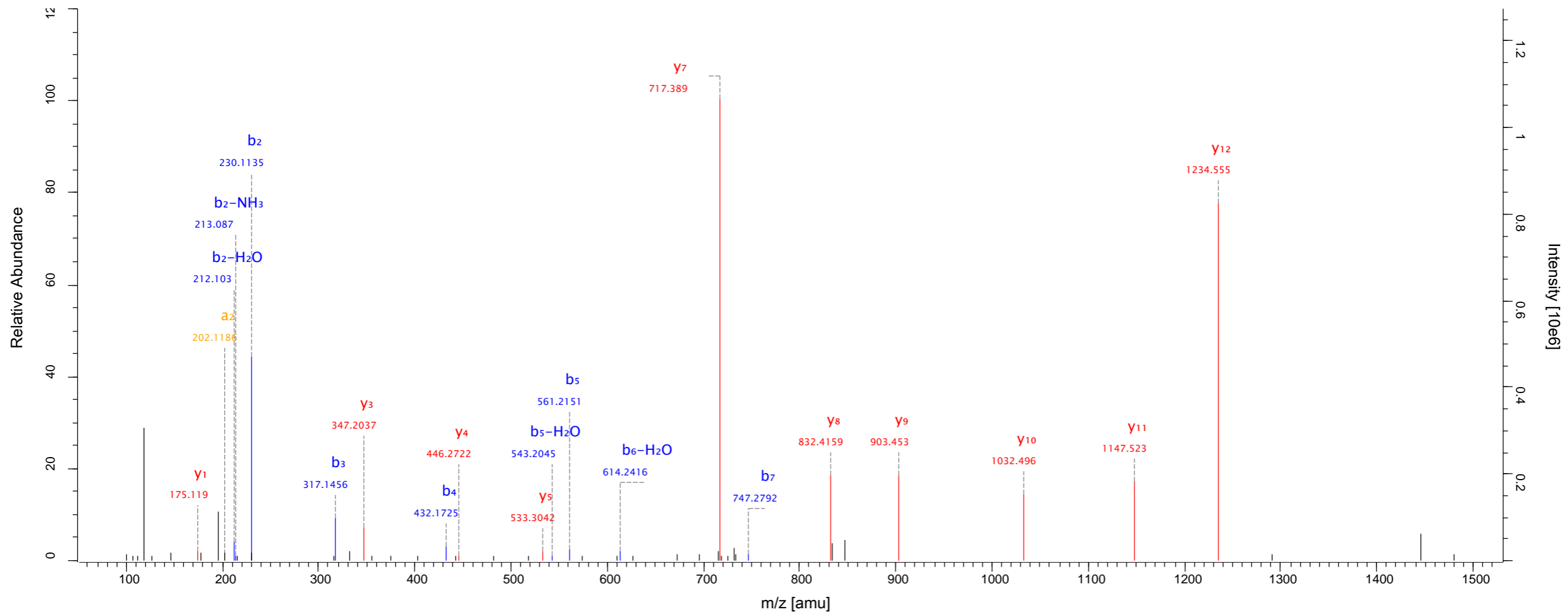
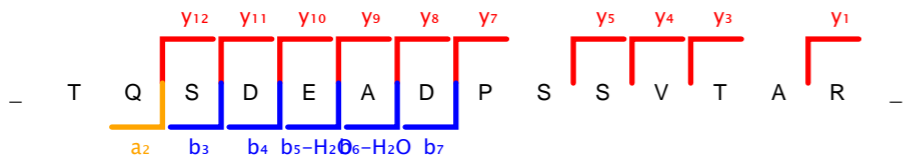
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
Scan number: 44705
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 65.35
Method: FTMS; HCD



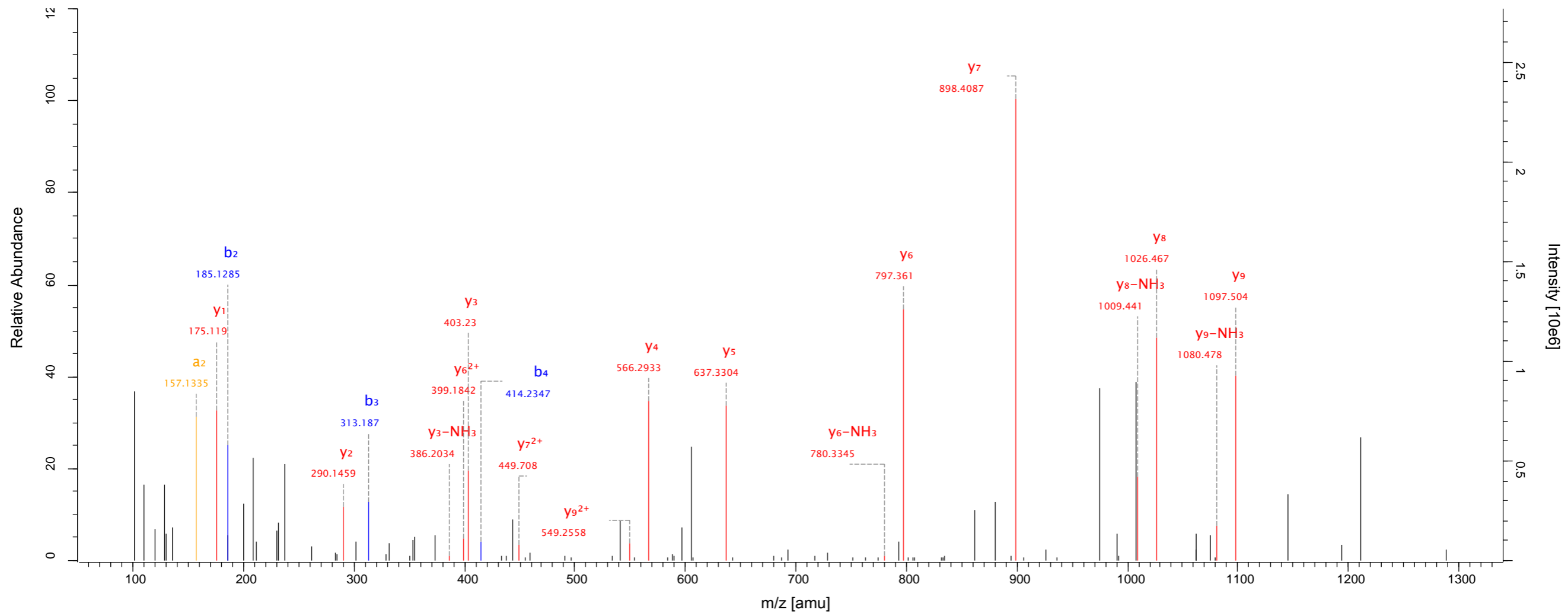
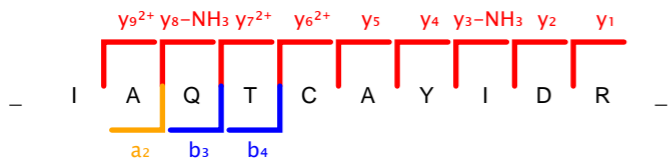
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_3
 Scan number: 6988
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 98.58
 Method: FTMS; HCD



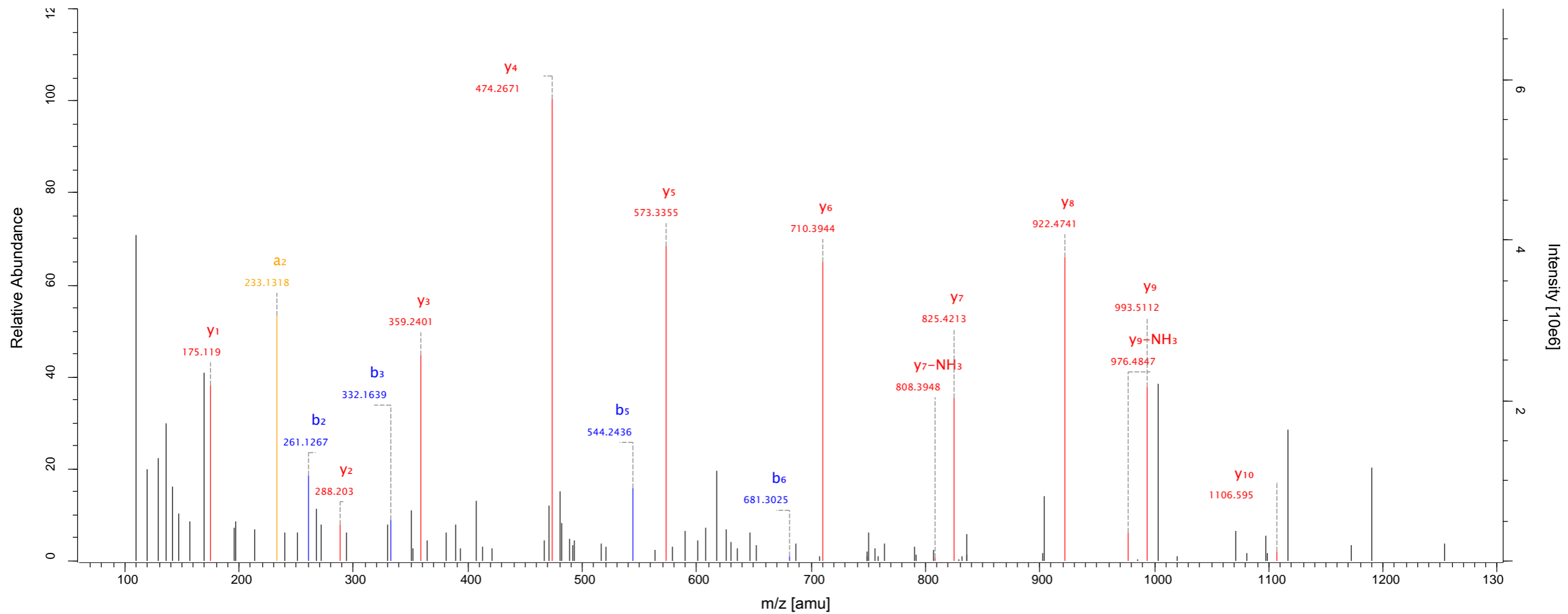
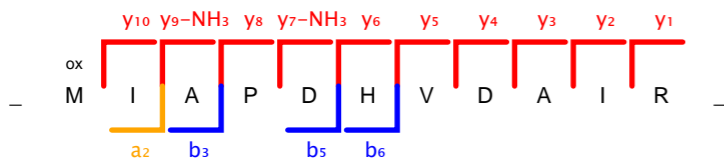
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_3
Scan number: 8334
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 95.65
Method: FTMS; HCD



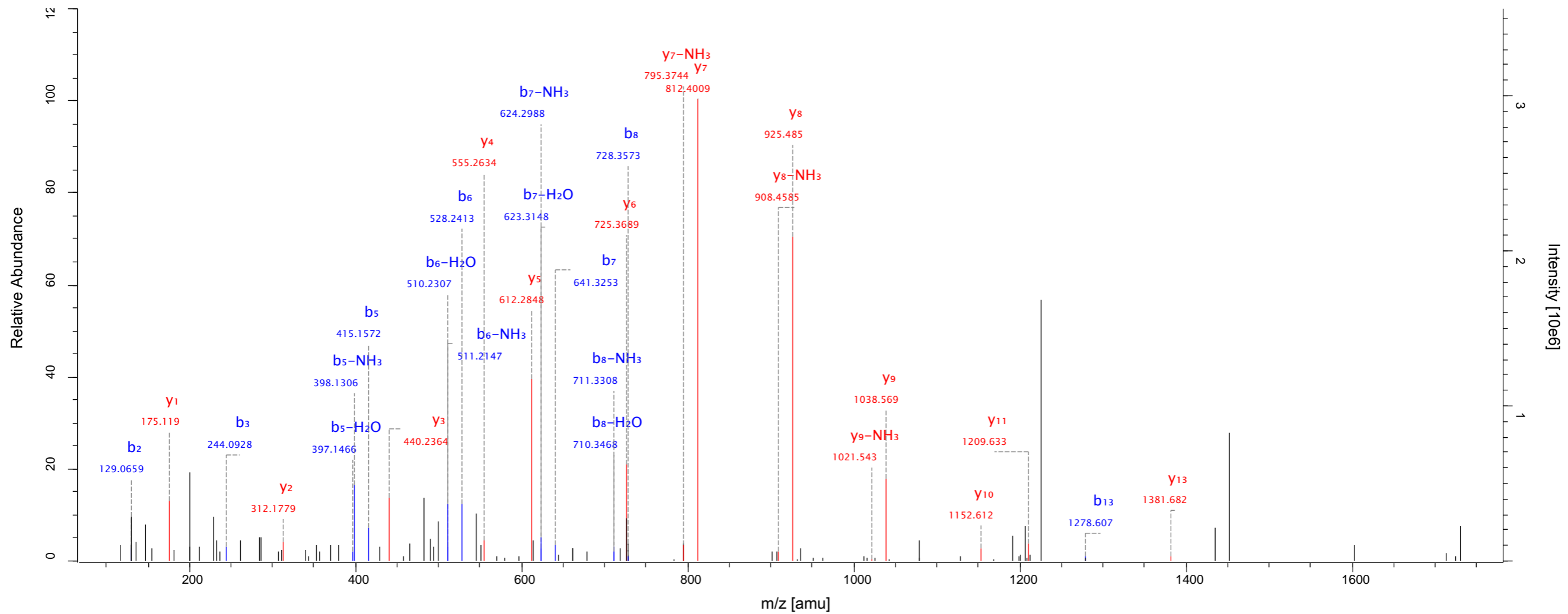
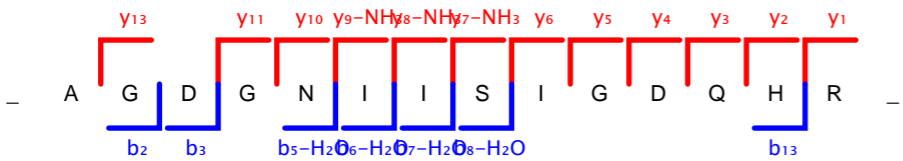
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
Scan number: 16357
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 145.72
Method: FTMS; HCD



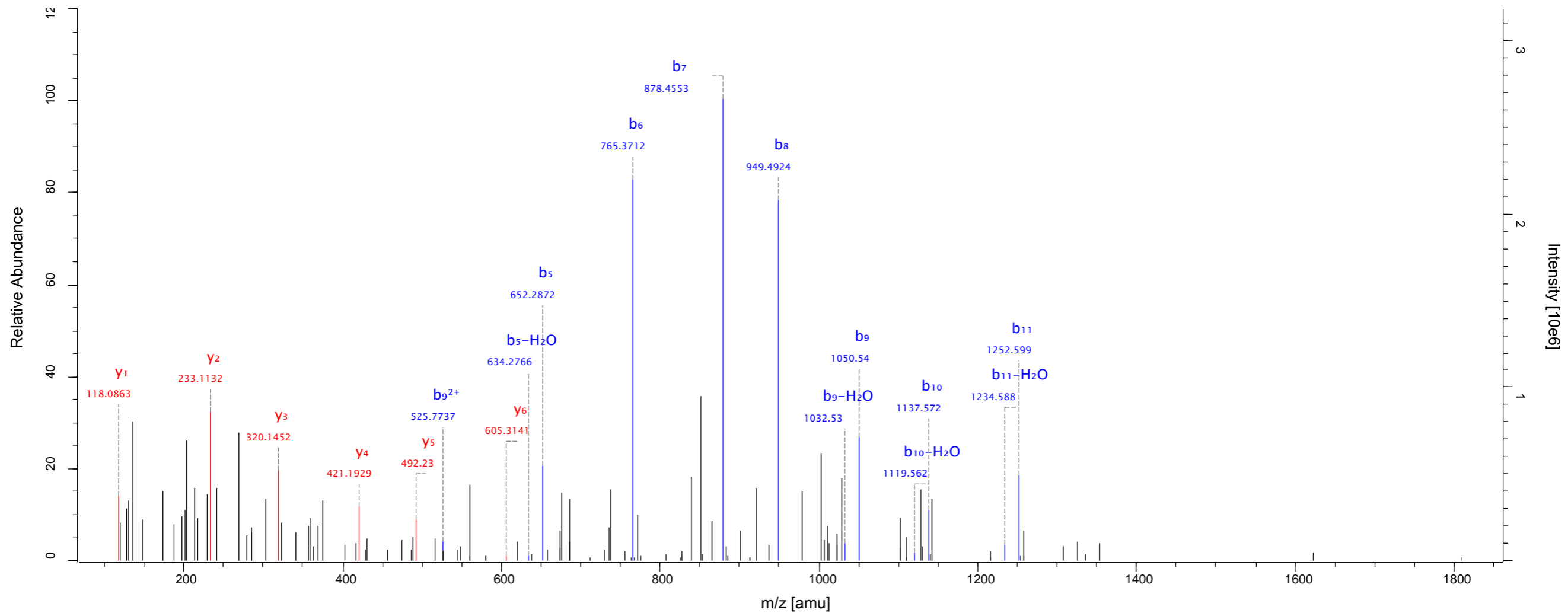
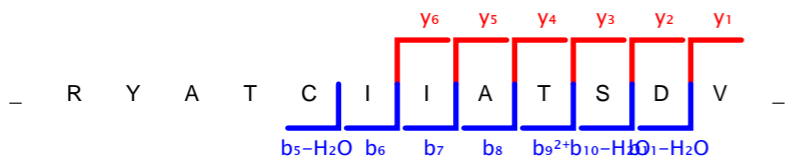
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_4
Scan number: 17002
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 97.46
Method: FTMS; HCD



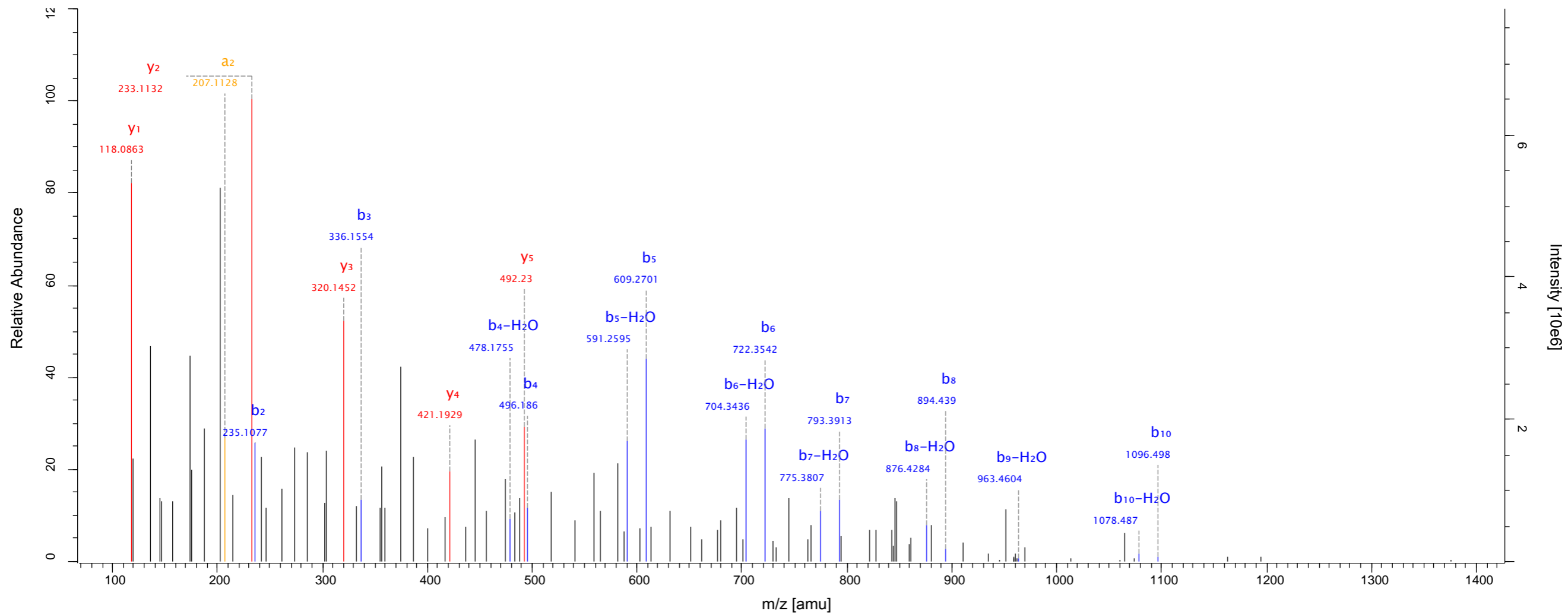
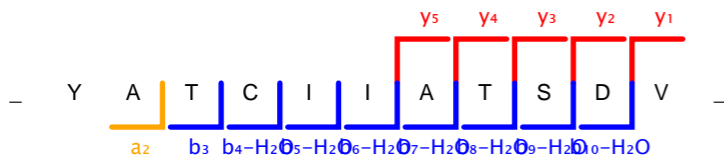
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
 Scan number: 22333
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 145
 Method: FTMS; HCD



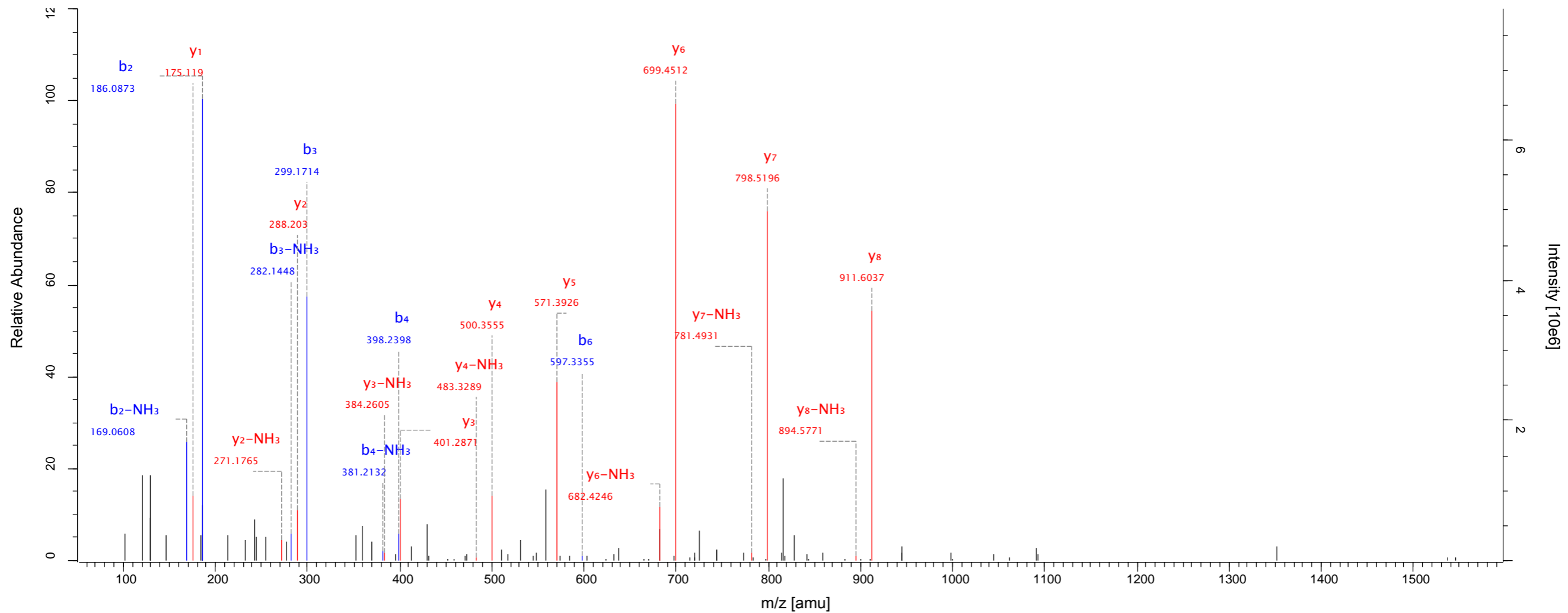
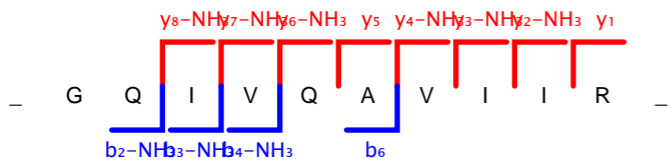
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
Scan number: 29515
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 89.4
Method: FTMS; HCD



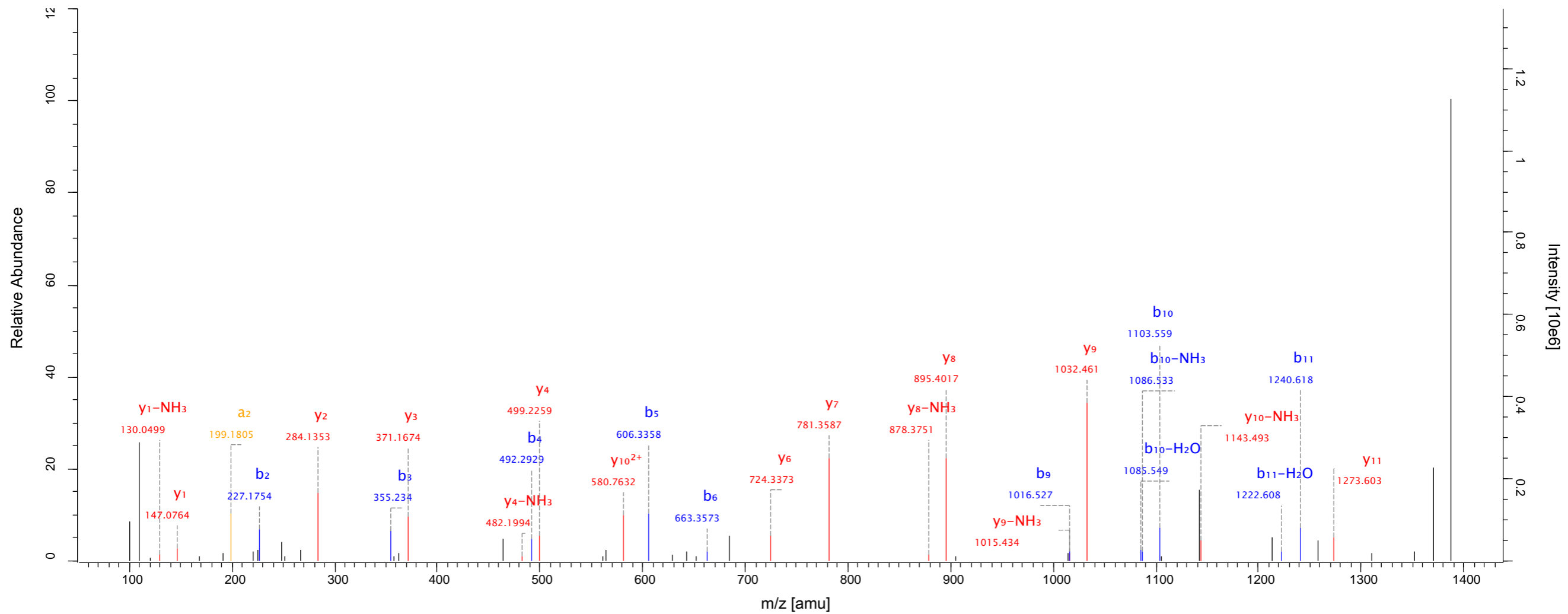
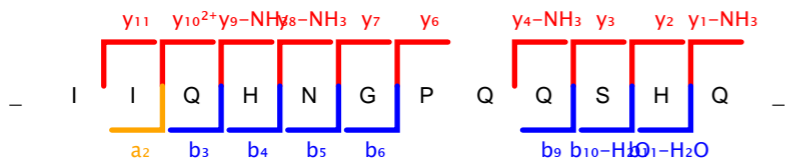
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
 Scan number: 34633
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 123.35
 Method: FTMS; HCD



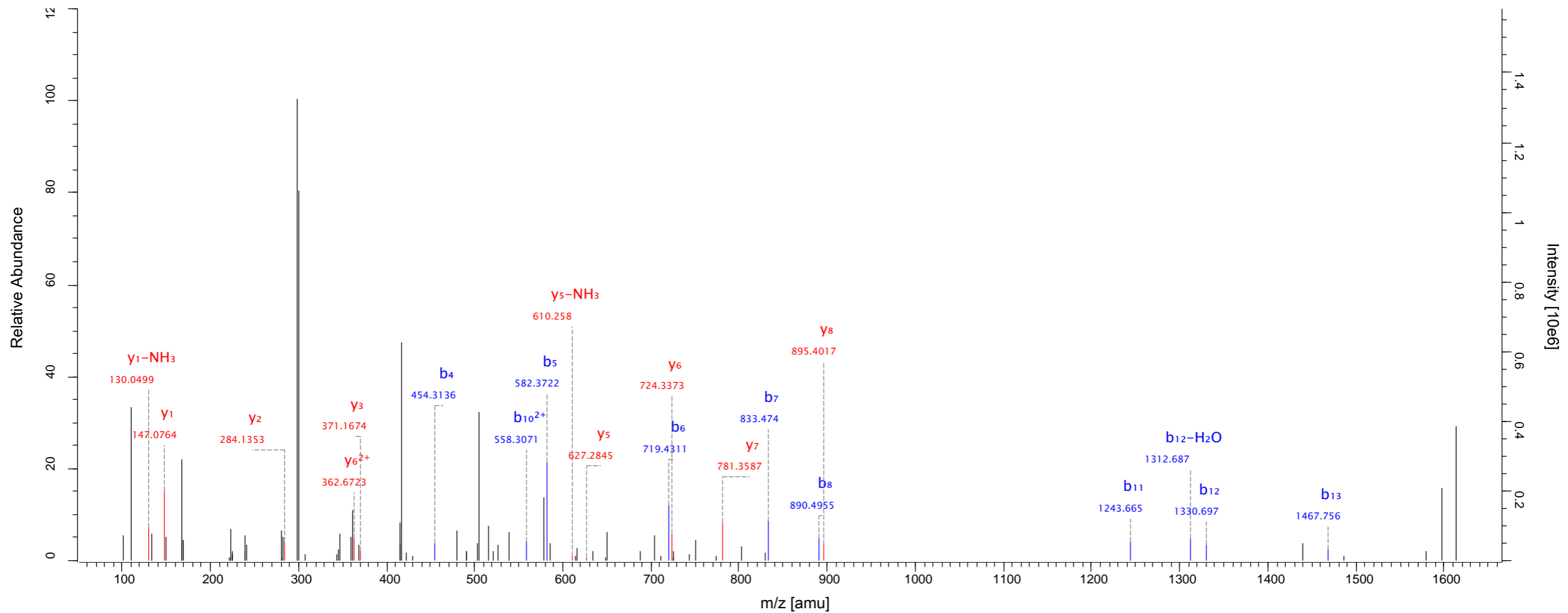
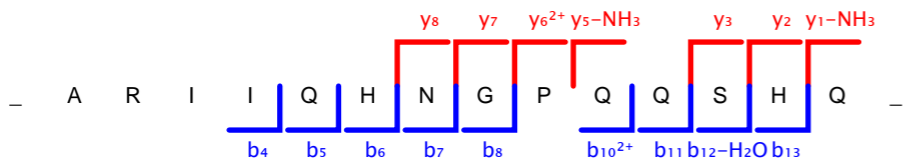
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
 Scan number: 36278
 Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
 Peptide score: 121.64
 Method: FTMS; HCD



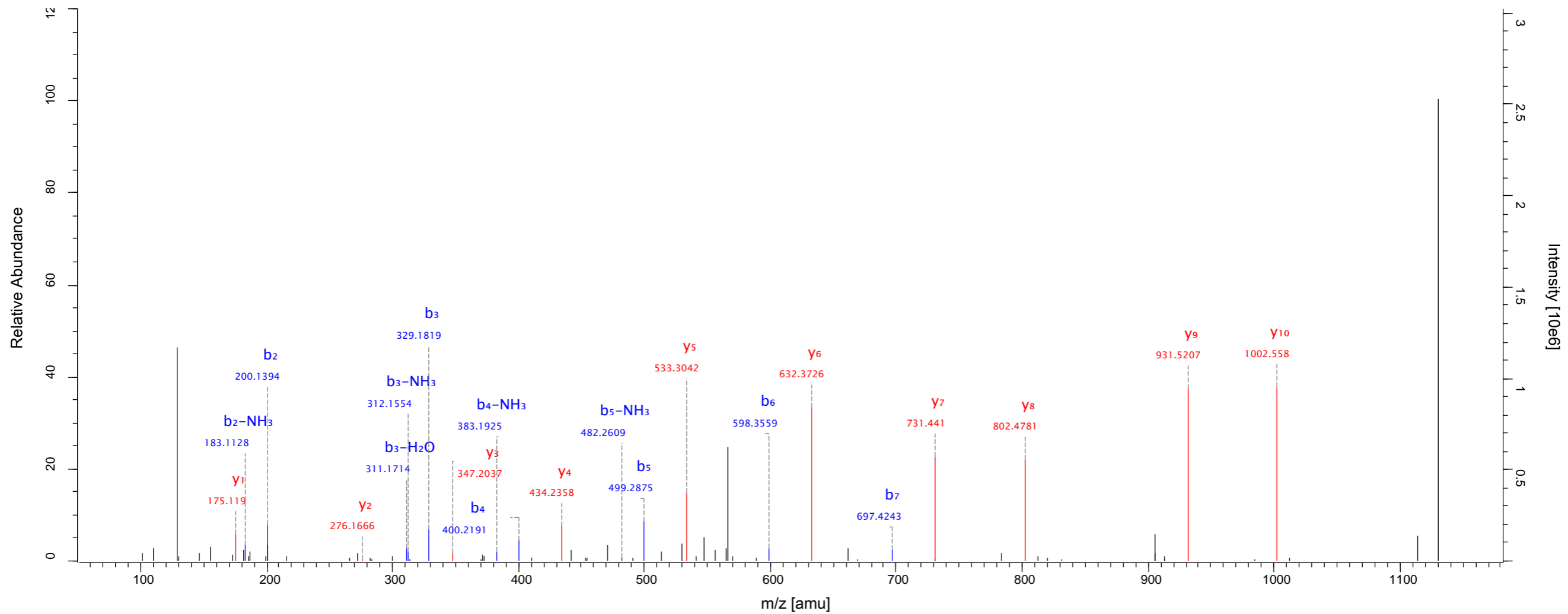
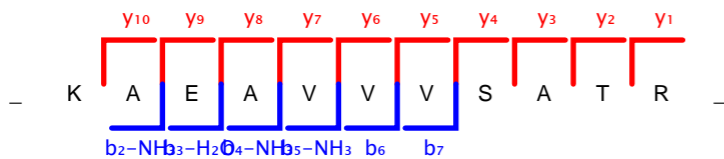
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
Scan number: 5453
Protein: ORFL103C - NC_006273.2[52569,53060]- <unknown description>
Peptide score: 220.94
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
Scan number: 6287
Protein: ORFL103C - NC_006273.2[52569,53060]- <unknown description>
Peptide score: 76.73
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_4
Scan number: 8977
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 143.79
Method: FTMS; HCD



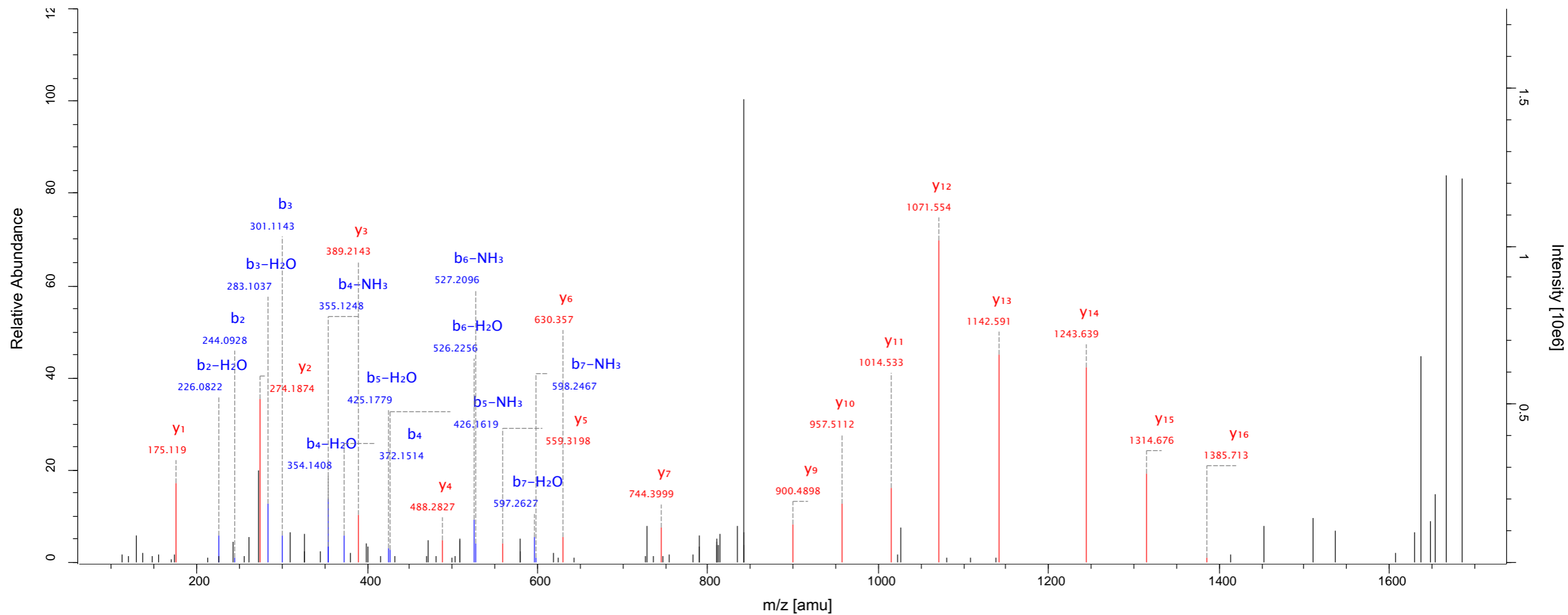
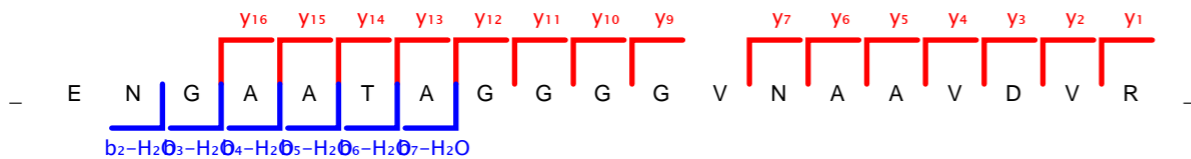
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_5

Scan number: 17119

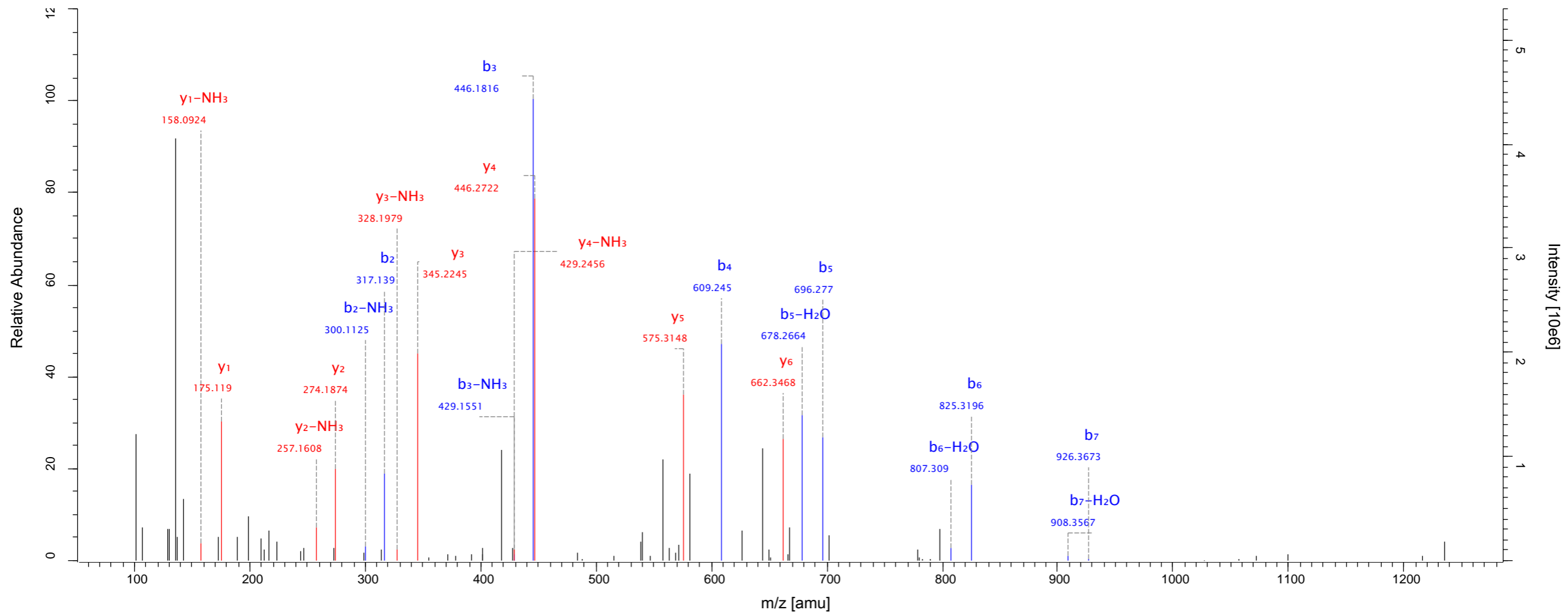
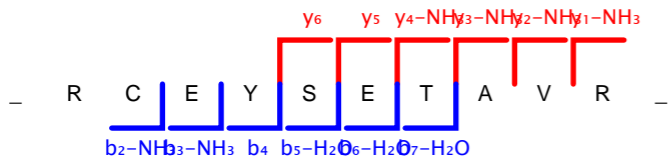
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 94.01

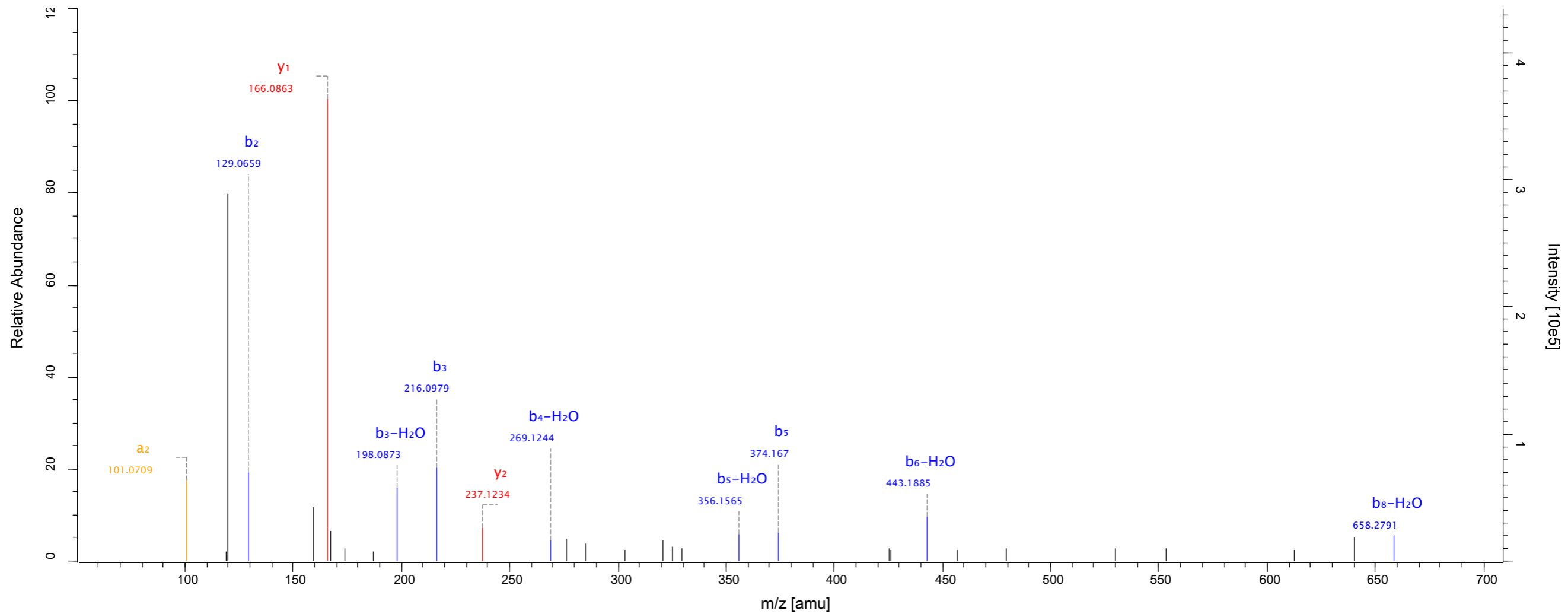
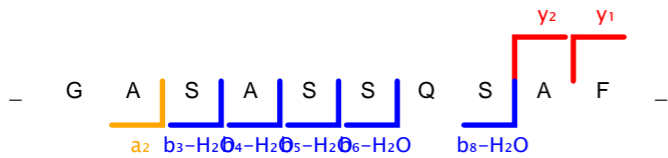
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_5
 Scan number: 8082
 Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>
 Peptide score: 132.78
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_6
Scan number: 10225
Protein: ORFL253W.iORF2 + NC_006273.2[161917,163431]+ <unknown description>
Peptide score: 59.25
Method: FTMS; HCD



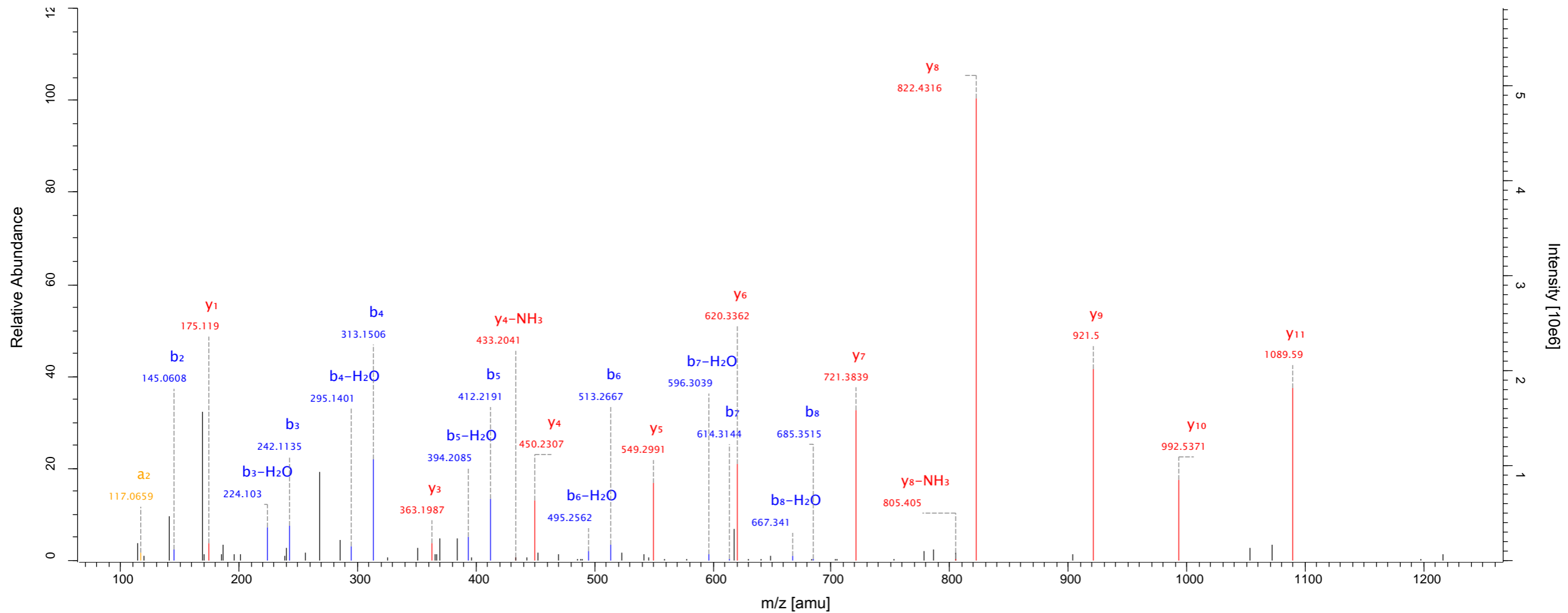
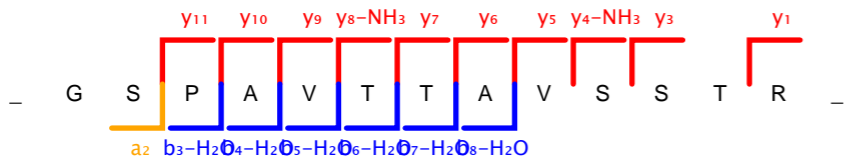
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_6

Scan number: 30601

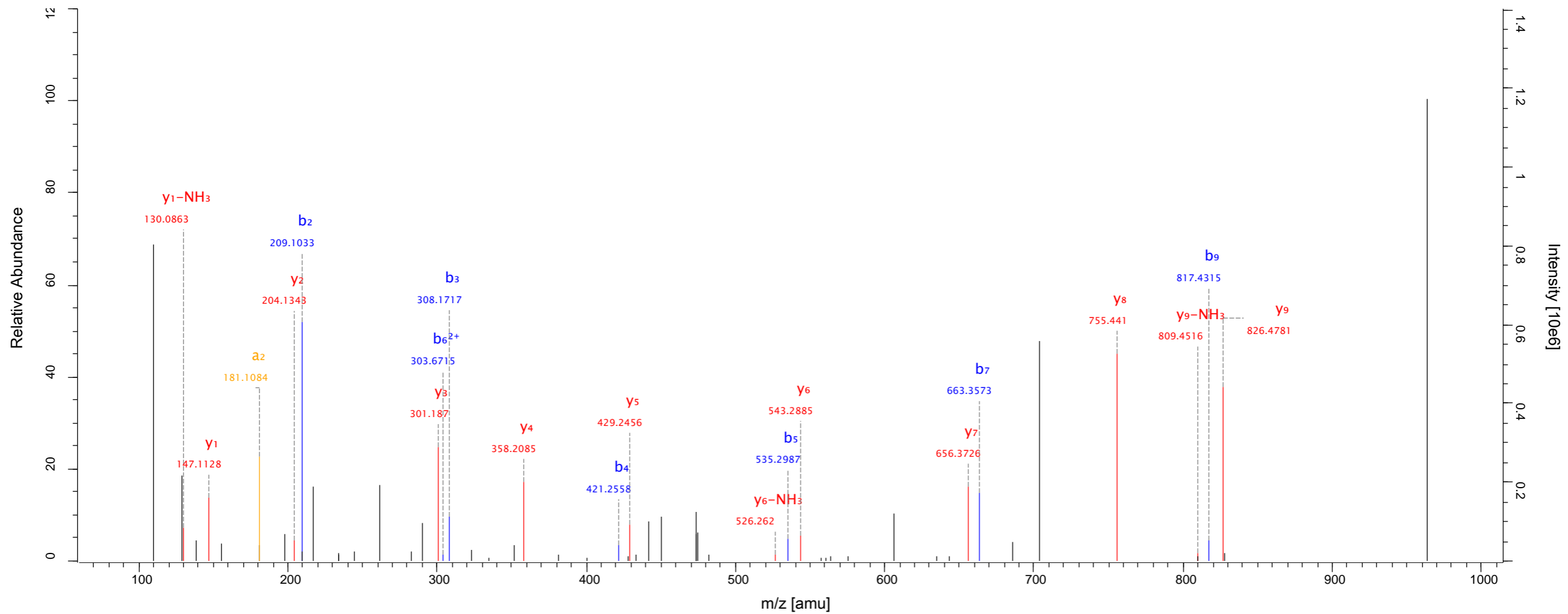
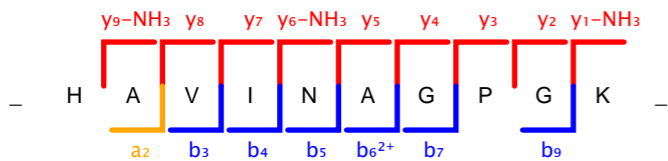
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 147.26

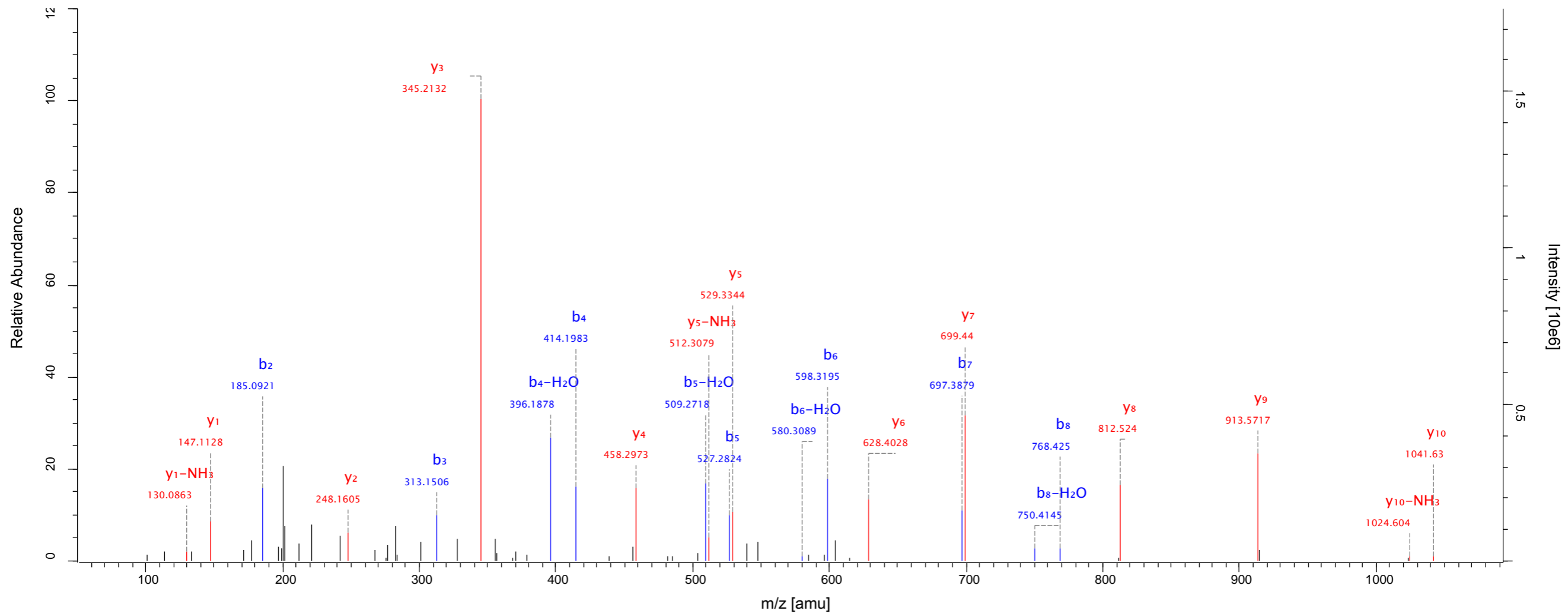
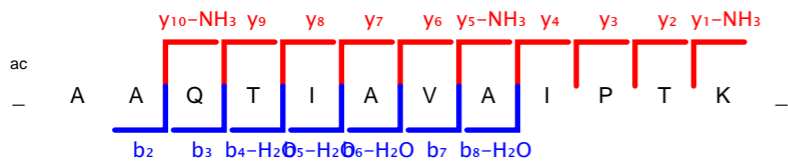
Method: FTMS; HCD



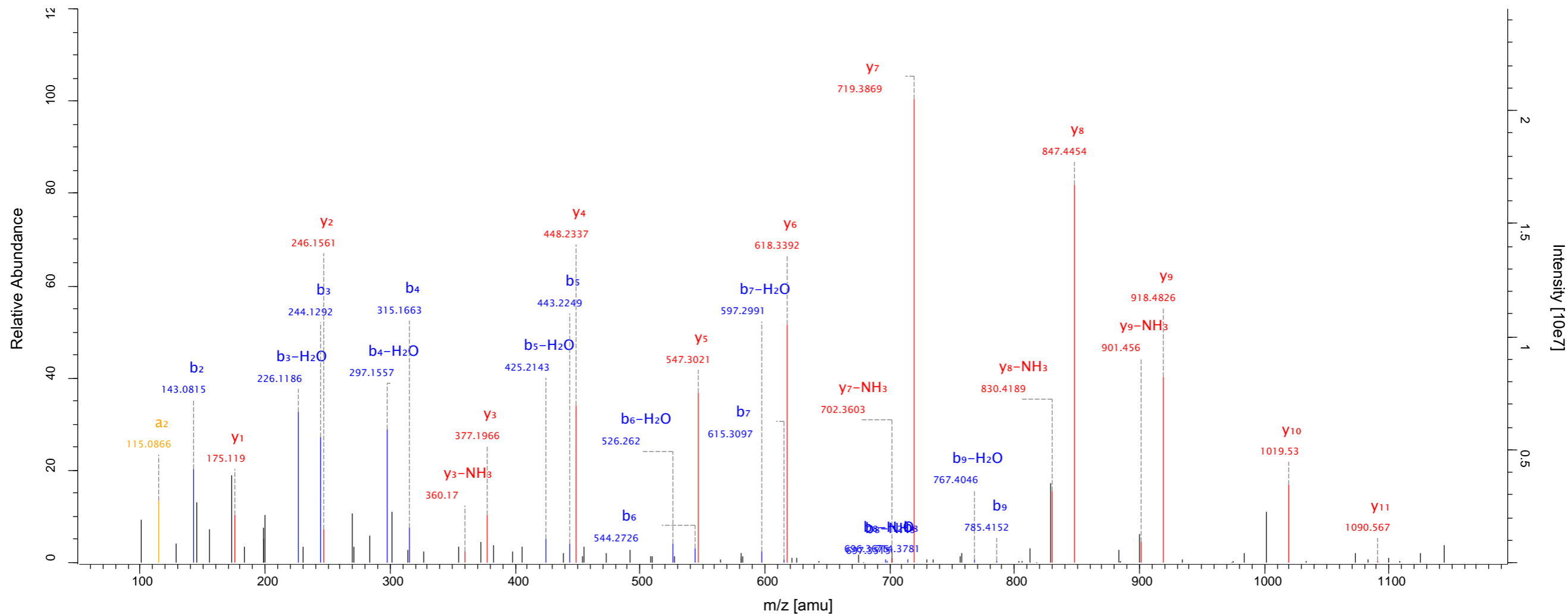
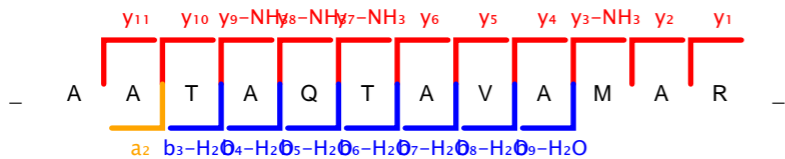
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_8
Scan number: 6529
Protein: ORFL87W + NC_006273.2[38099,40114]+ <unknown description>
Peptide score: 149.72
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_9
 Scan number: 28063
 Protein: ORFS366W + NC_006273.2[227254,227469]+ <unknown description>
 Peptide score: 181.32
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13
 Scan number: 14848
 Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>
 Peptide score: 177.04
 Method: FTMS; HCD



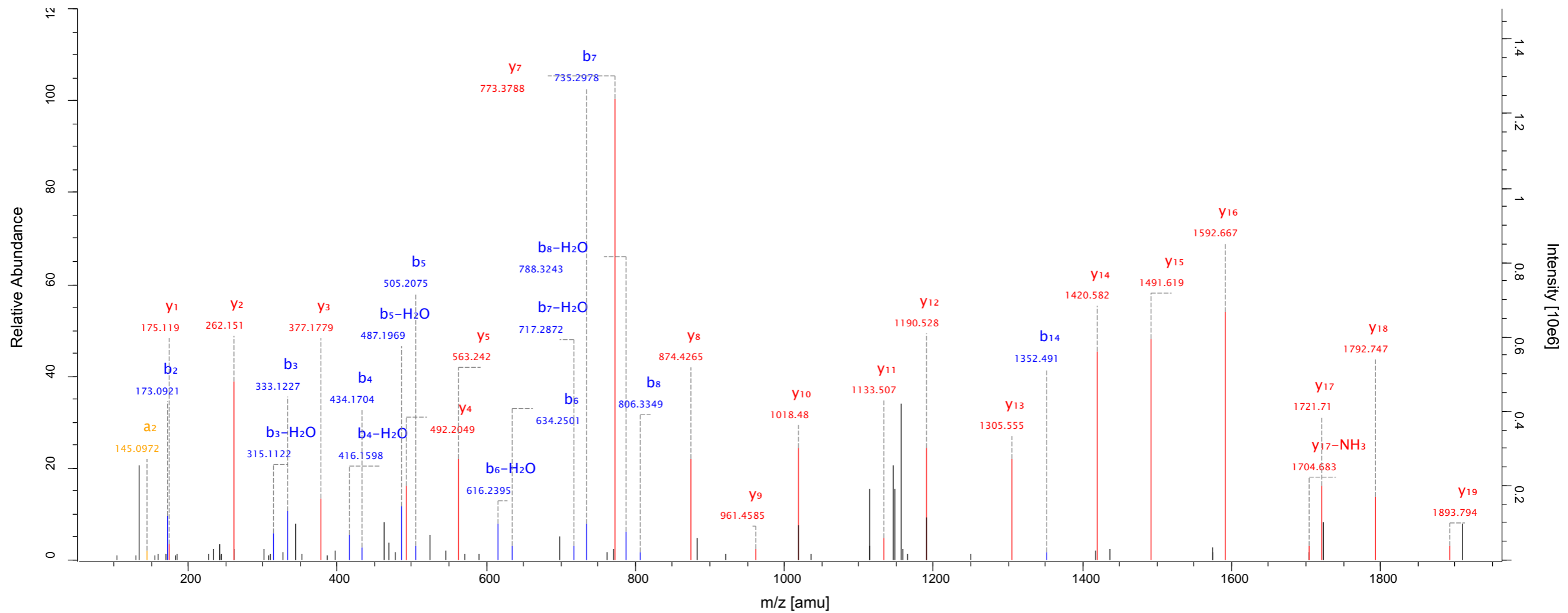
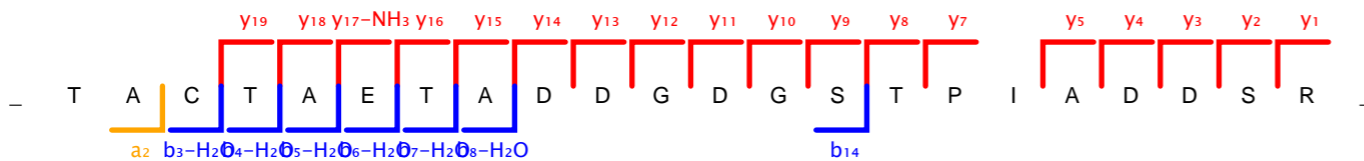
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_13

Scan number: 16148

Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>; ORFL131W.iORF2 + NC_006273.2[73501,73947]+ <unknown description>

Peptide score: 199.76

Method: FTMS; HCD



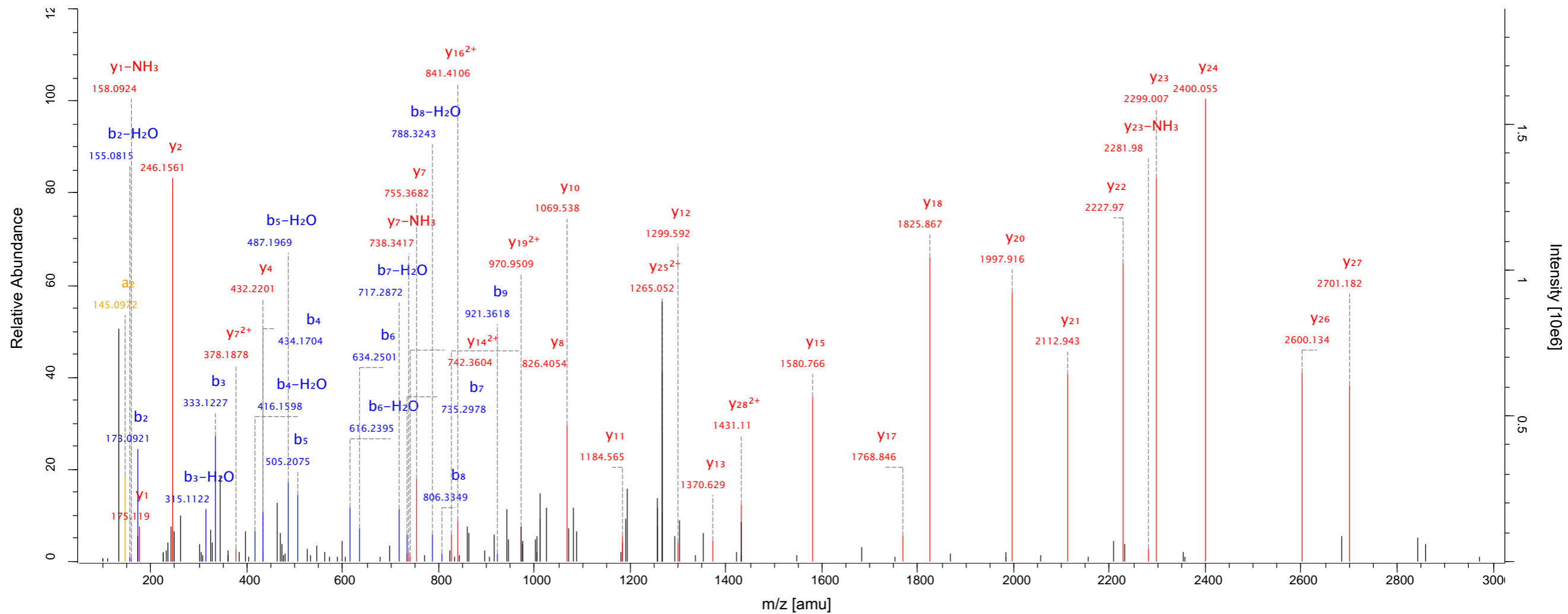
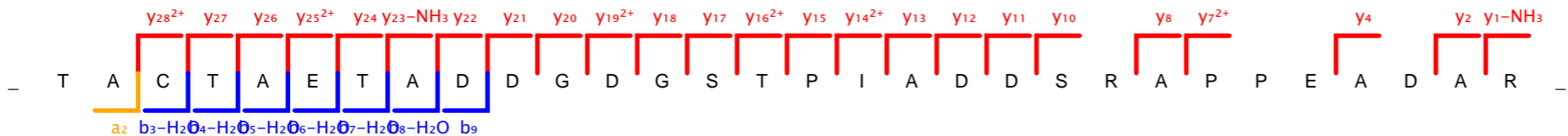
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13

Scan number: 19970

Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>; ORFL131W.iORF2 + NC_006273.2[73501,73947]+ <unknown description>

Peptide score: 192.47

Method: FTMS; HCD



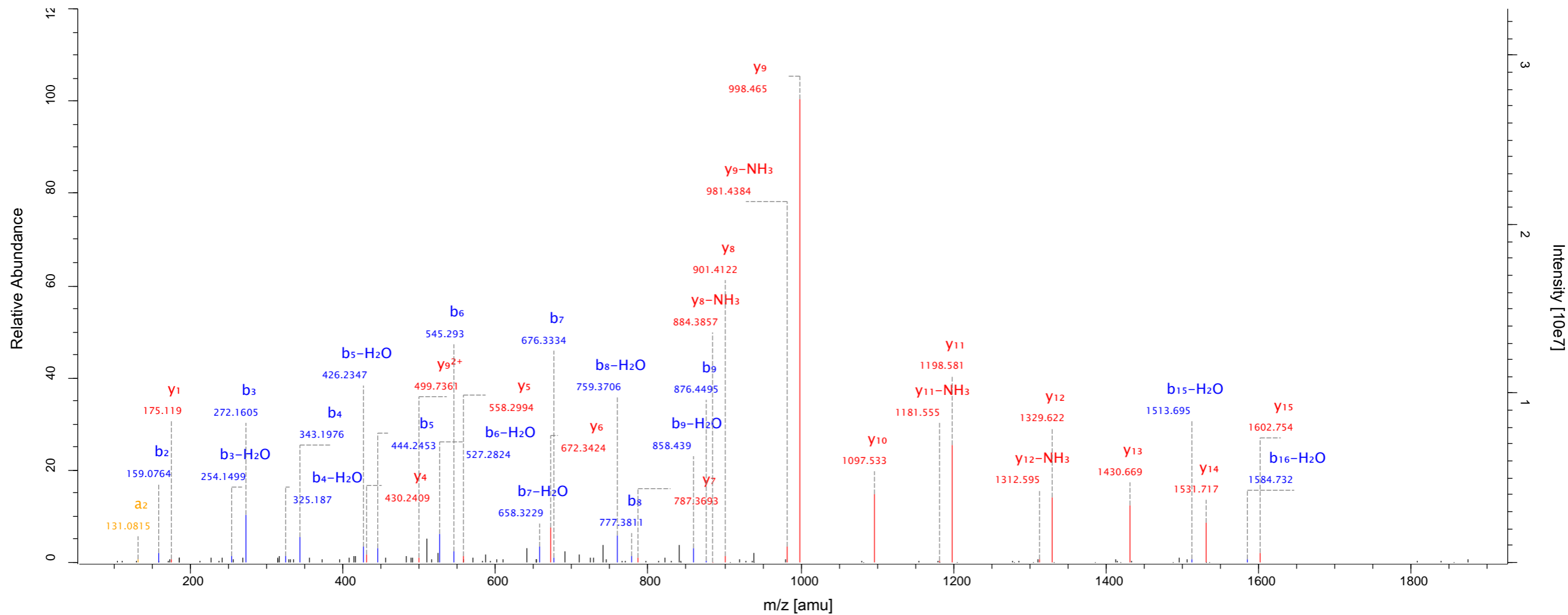
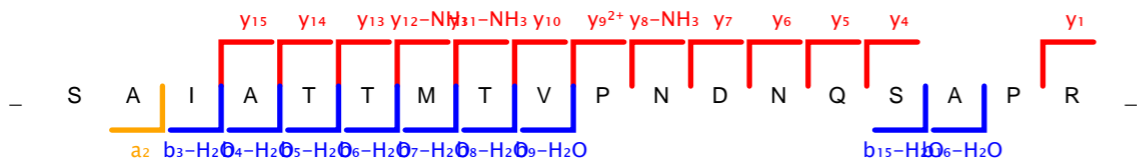
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_13

Scan number: 24694

Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>; ORFL131W.iORF2 + NC_006273.2[73501,73947]+ <unknown description>

Peptide score: 213.34

Method: FTMS; HCD



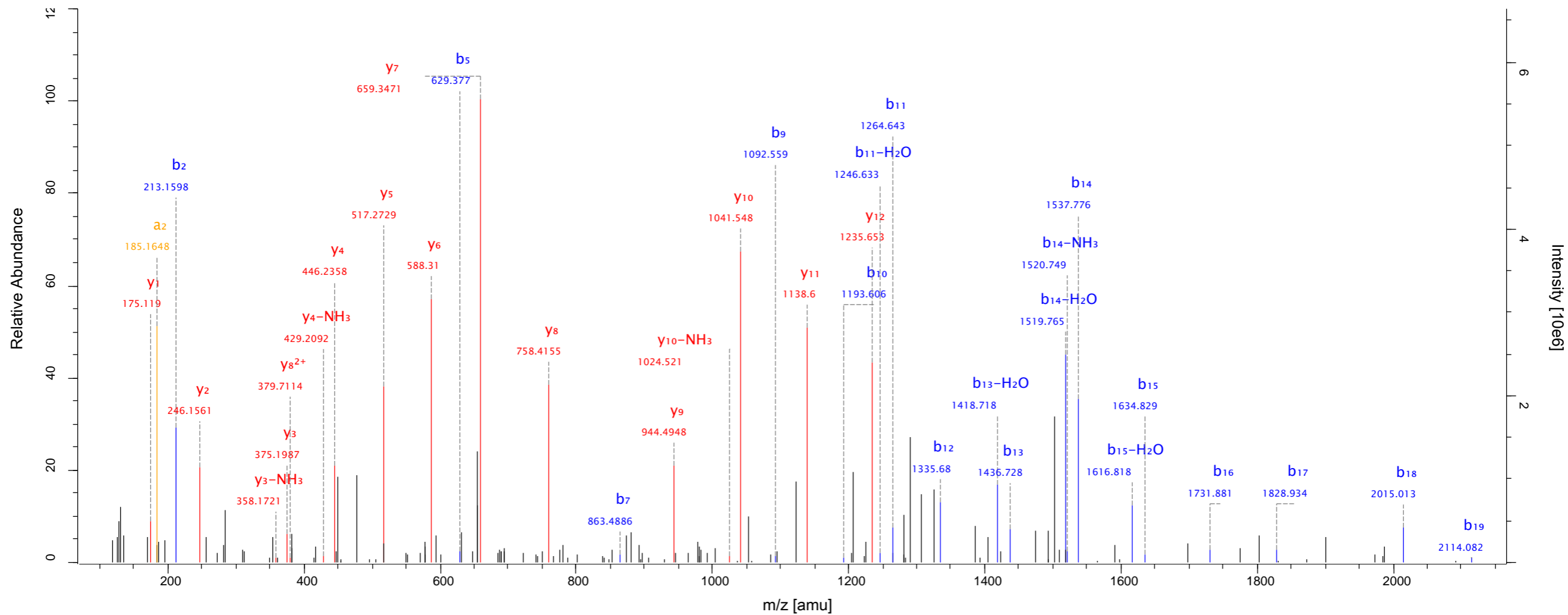
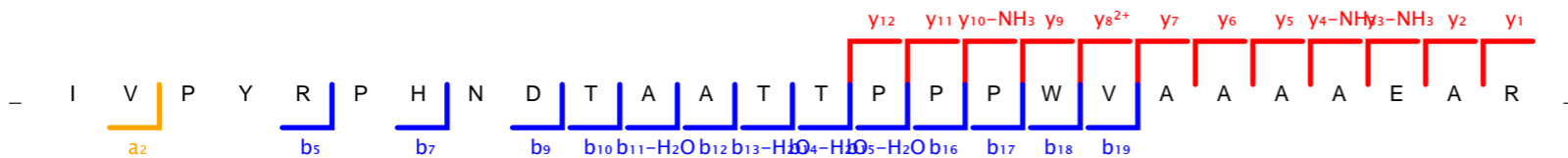
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13

Scan number: 37446

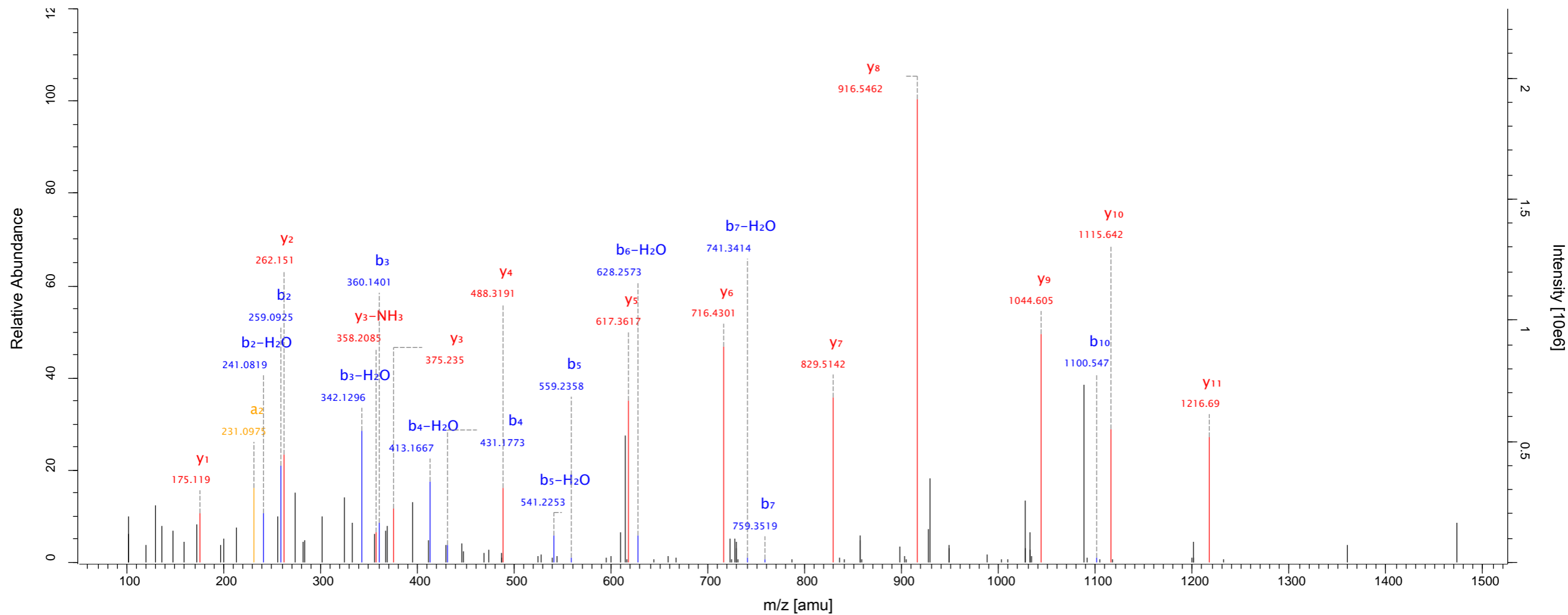
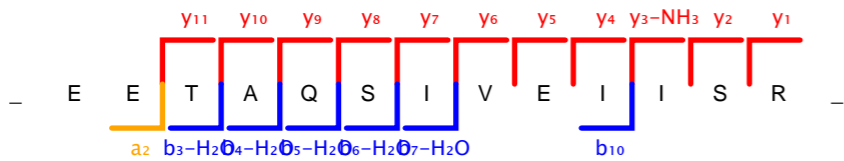
Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>; ORFL131W.iORF2 + NC_006273.2[73501,73947]+ <unknown description>

Peptide score: 129.23

Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13
 Scan number: 51068
 Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>
 Peptide score: 111.39
 Method: FTMS; HCD



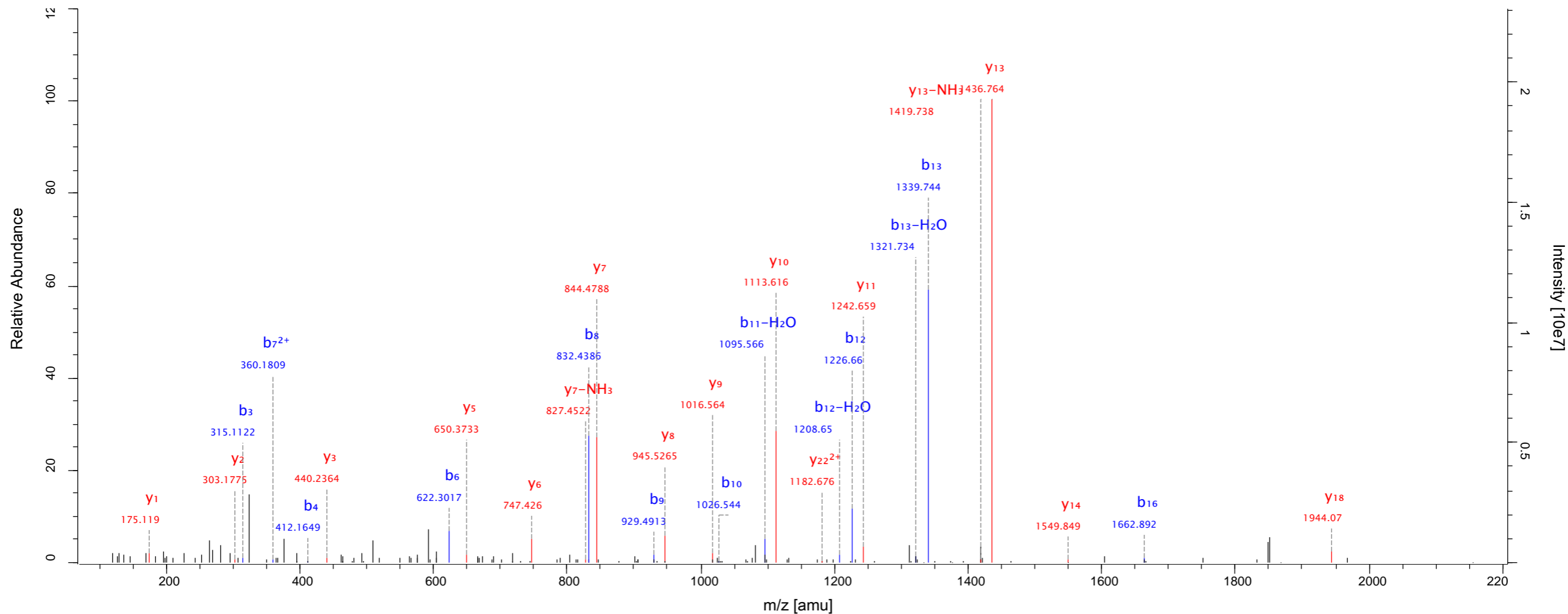
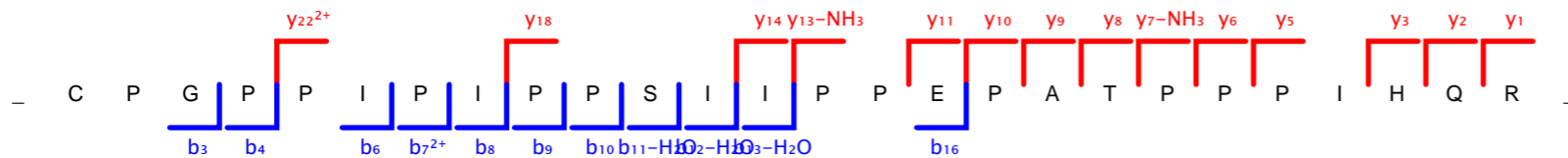
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_13

Scan number: 57325

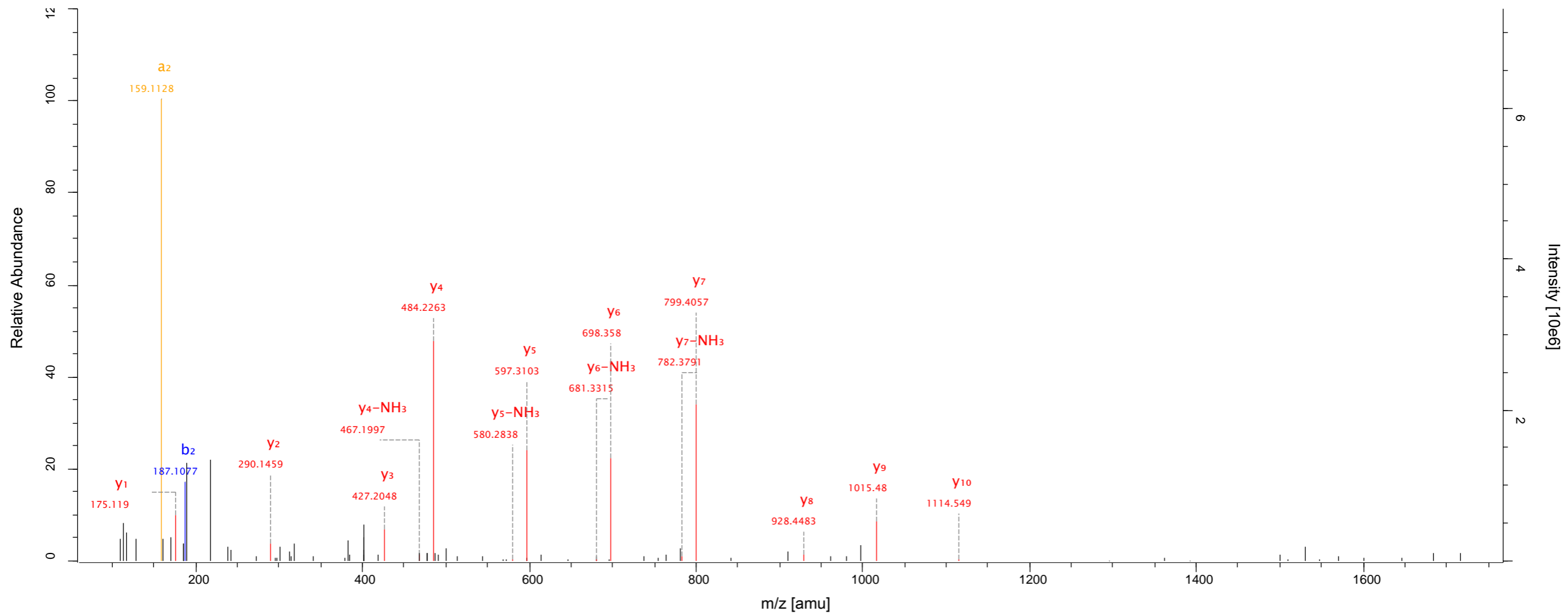
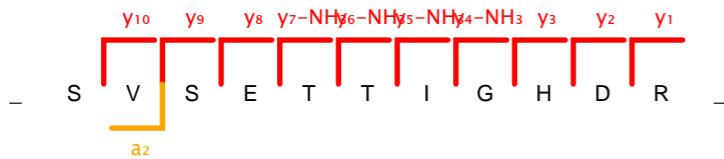
Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>; ORFL131W.iORF2 + NC_006273.2[73501,73947]+ <unknown description>

Peptide score: 84.57

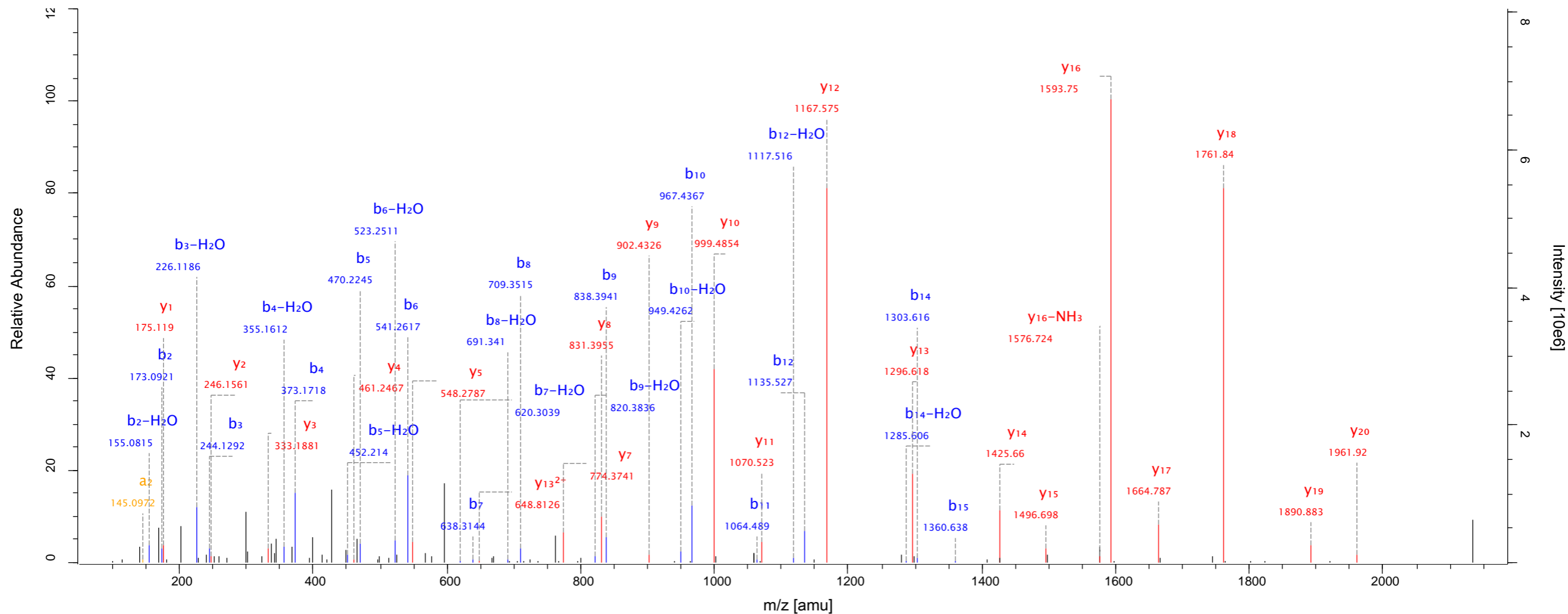
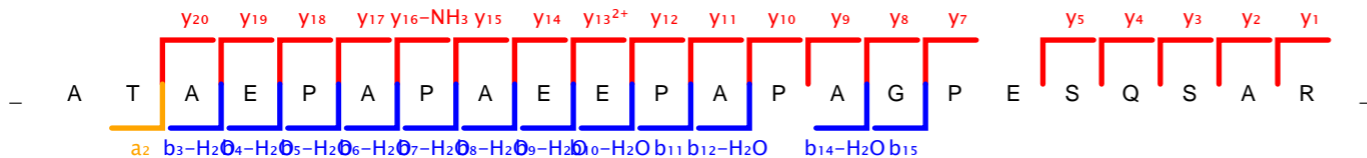
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_13
Scan number: 8060
Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>
Peptide score: 92.54
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_5
 Scan number: 14960
 Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>
 Peptide score: 252.51
 Method: FTMS; HCD



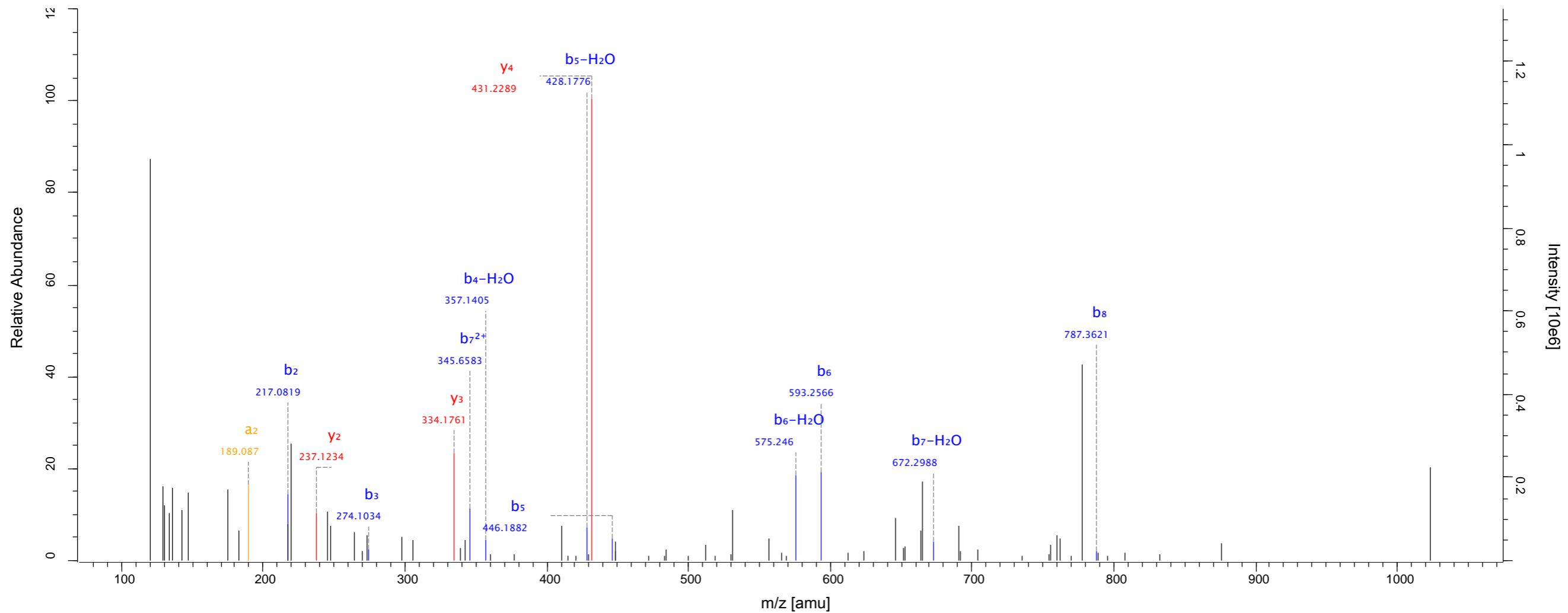
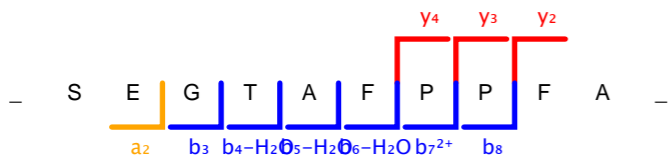
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_5

Scan number: 42435

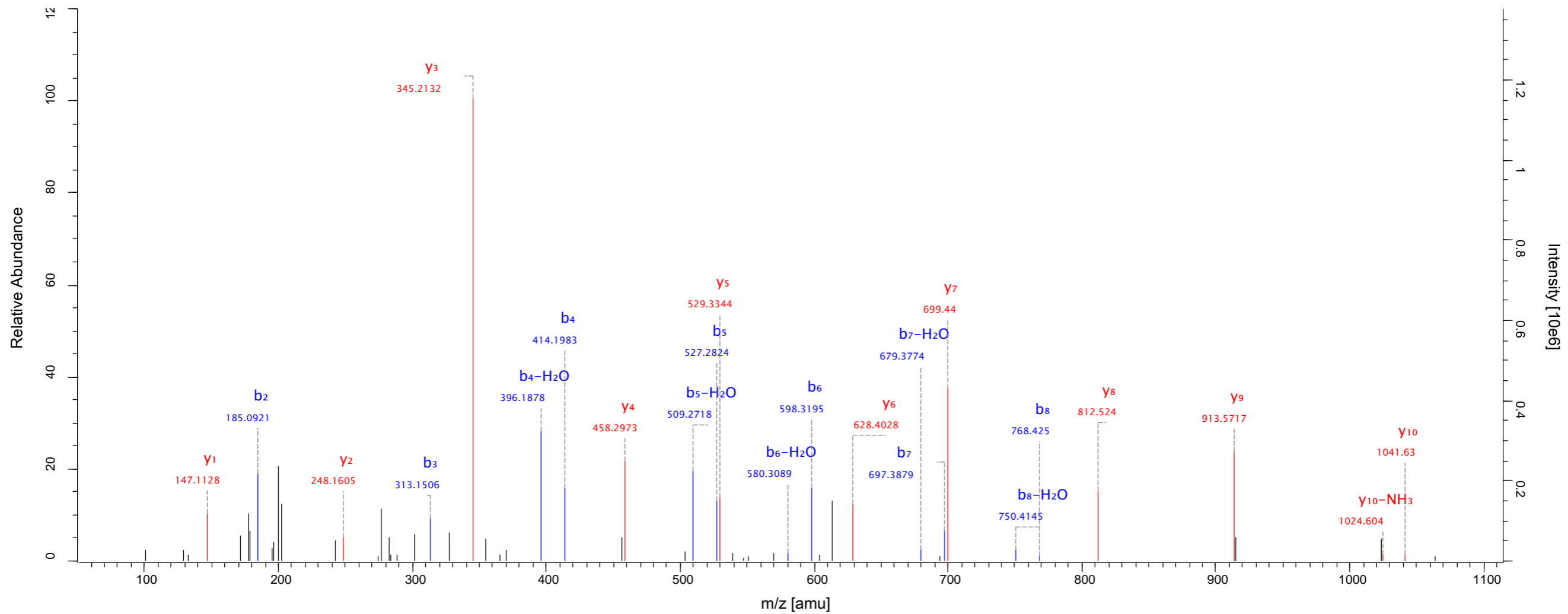
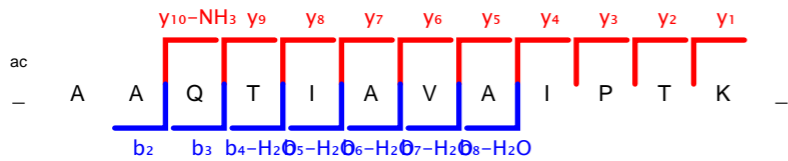
Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>; ORFL131W.iORF2 + NC_006273.2[73501,73947]+ <unknown description>

Peptide score: 67.9

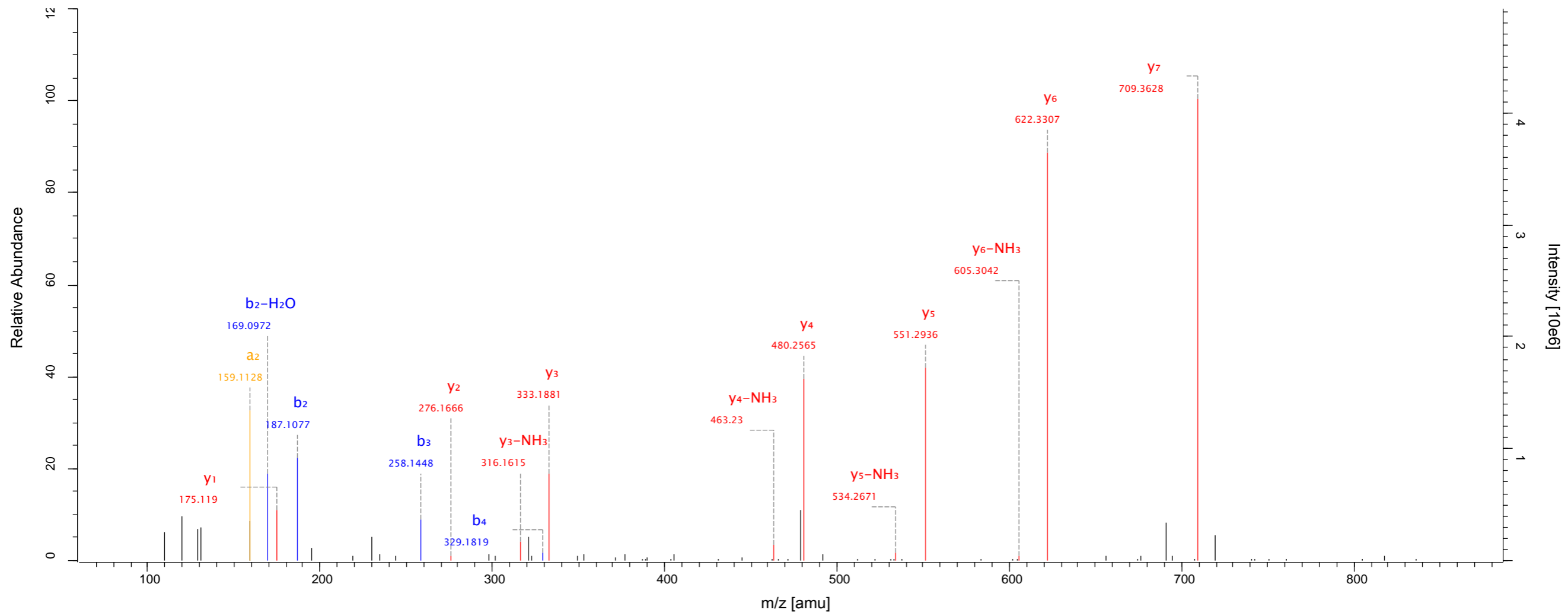
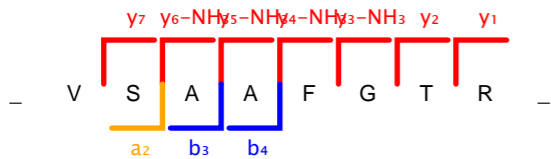
Method: FTMS; HCD



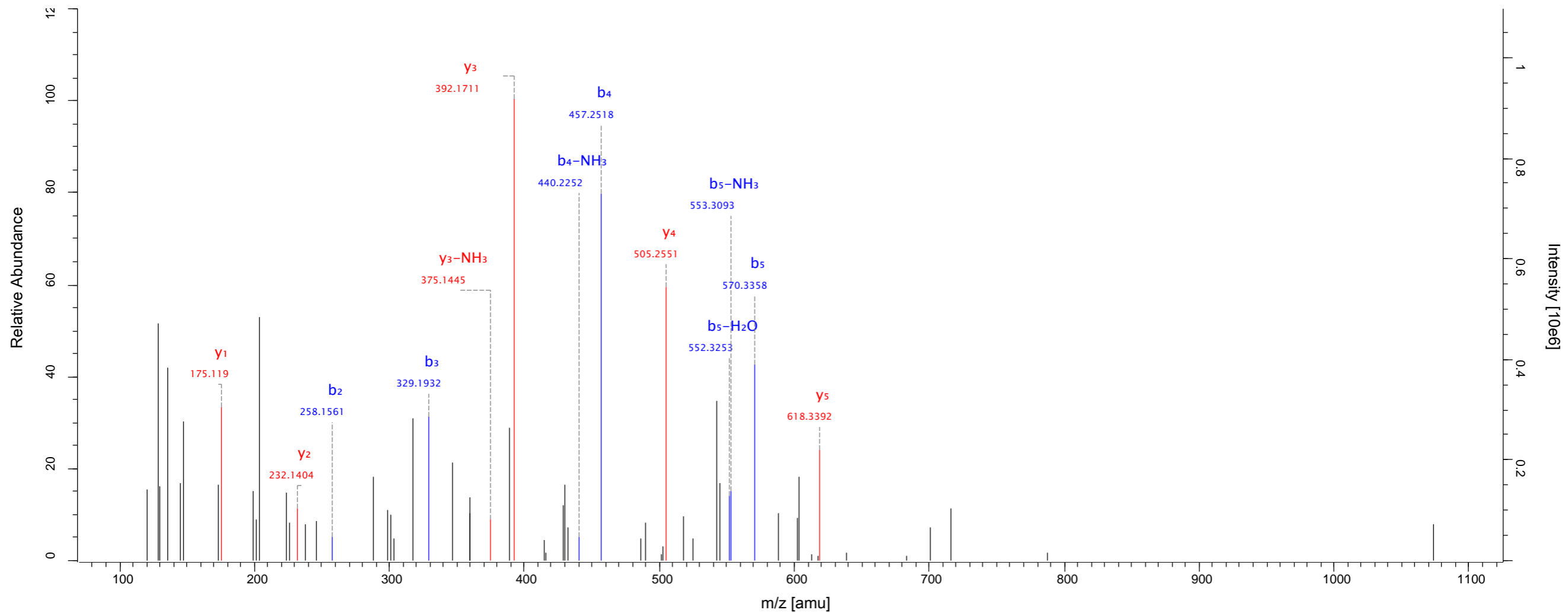
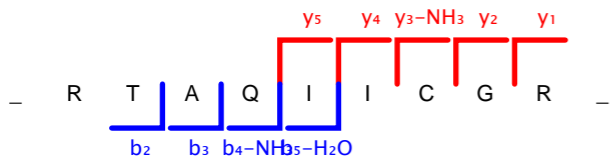
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_25
Scan number: 27270
Protein: ORFS366W + NC_006273.2[227254,227469]+ <unknown description>
Peptide score: 174.84
Method: FTMS; HCD



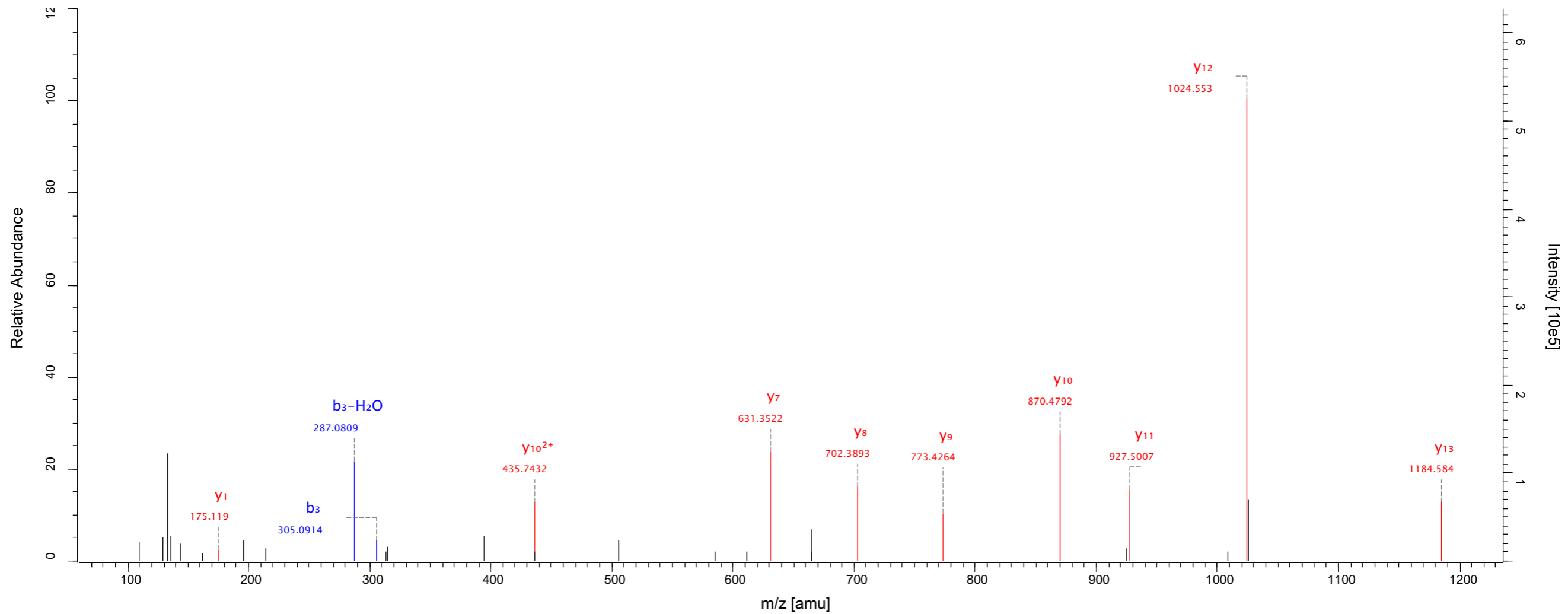
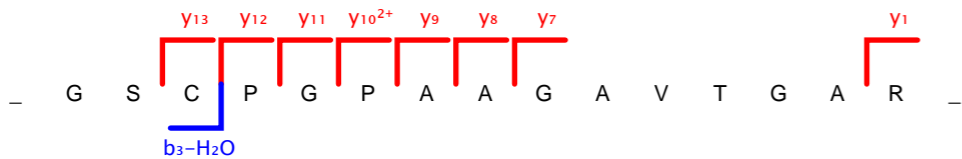
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 10081
Protein: ORFL121C - NC_006273.2[65579,65851]- <unknown description>
Peptide score: 136.11
Method: FTMS; HCD



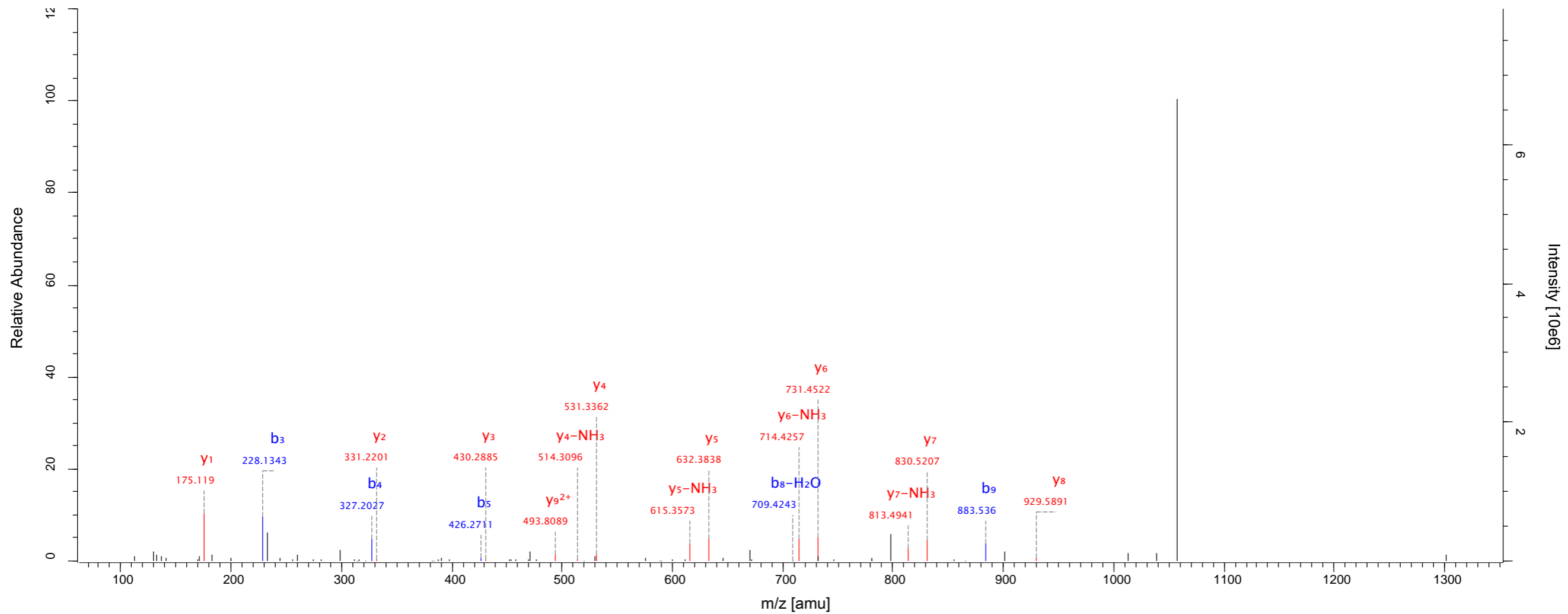
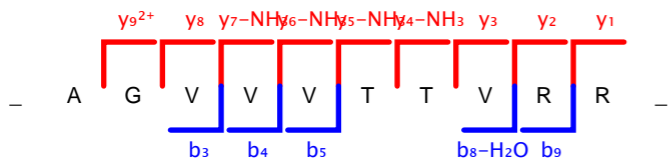
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 10537
Protein: ORFL137W.iORF1 + NC_006273.2[75299,75529]+ <unknown description>
Peptide score: 56.01
Method: FTMS; HCD



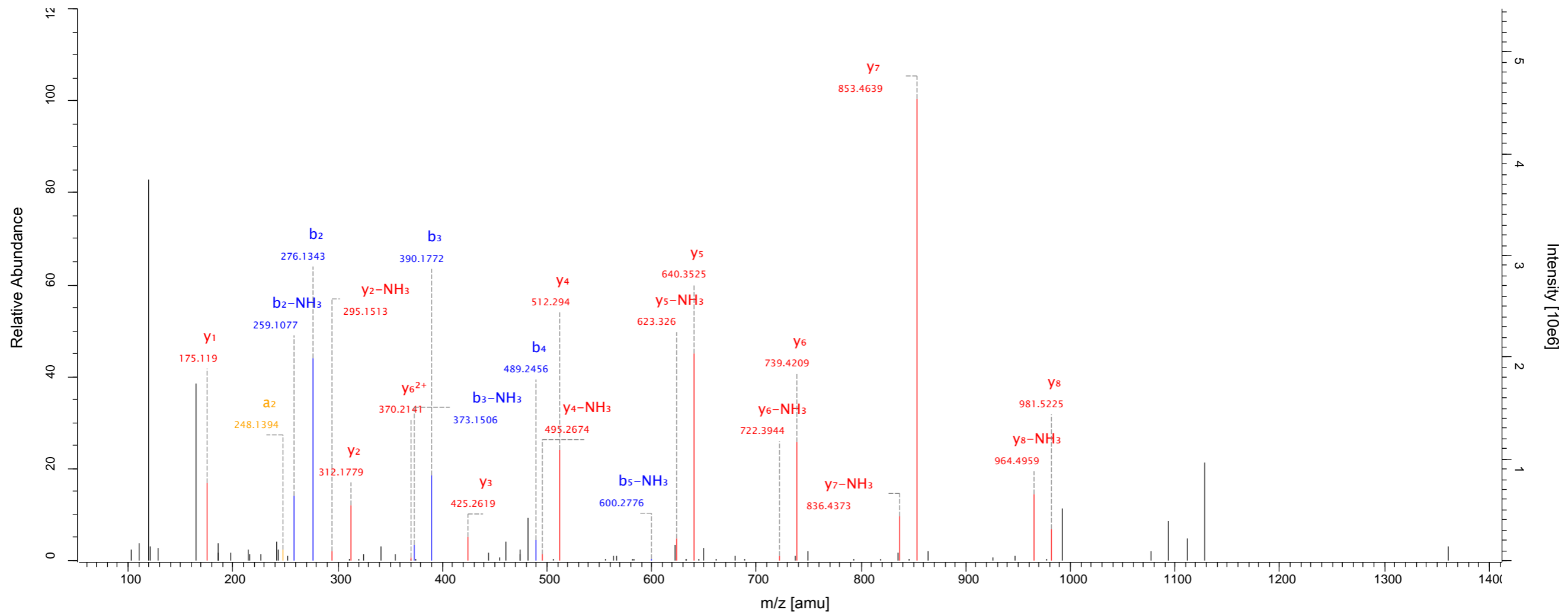
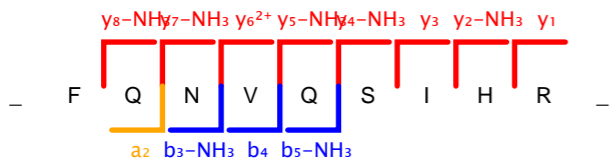
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 10806
Protein: ORFL235W + NC_006273.2[149239,150022]+ <unknown description>
Peptide score: 68.81
Method: FTMS; HCD



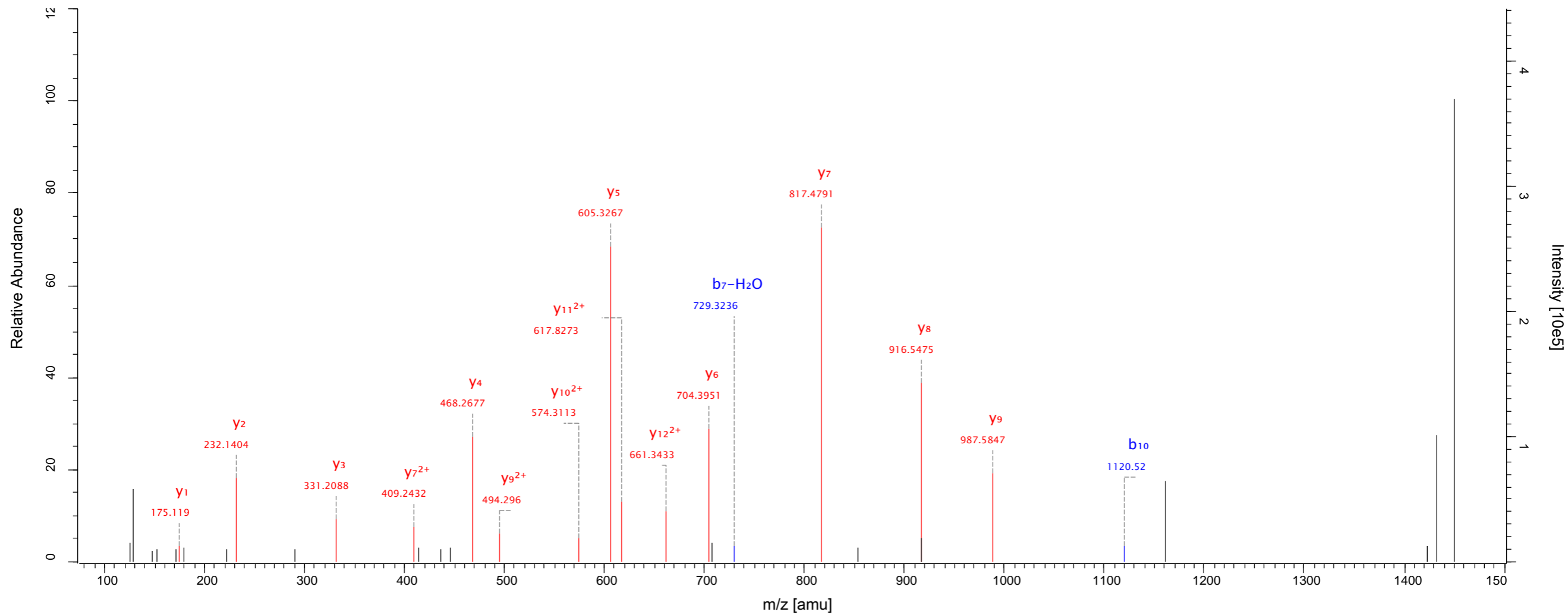
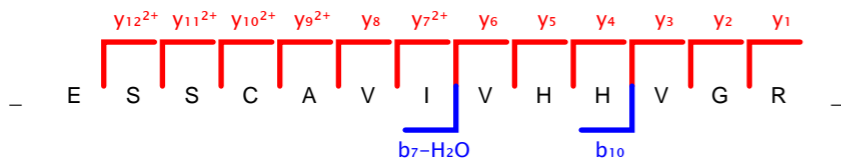
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 10866
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 120.12
Method: FTMS; HCD



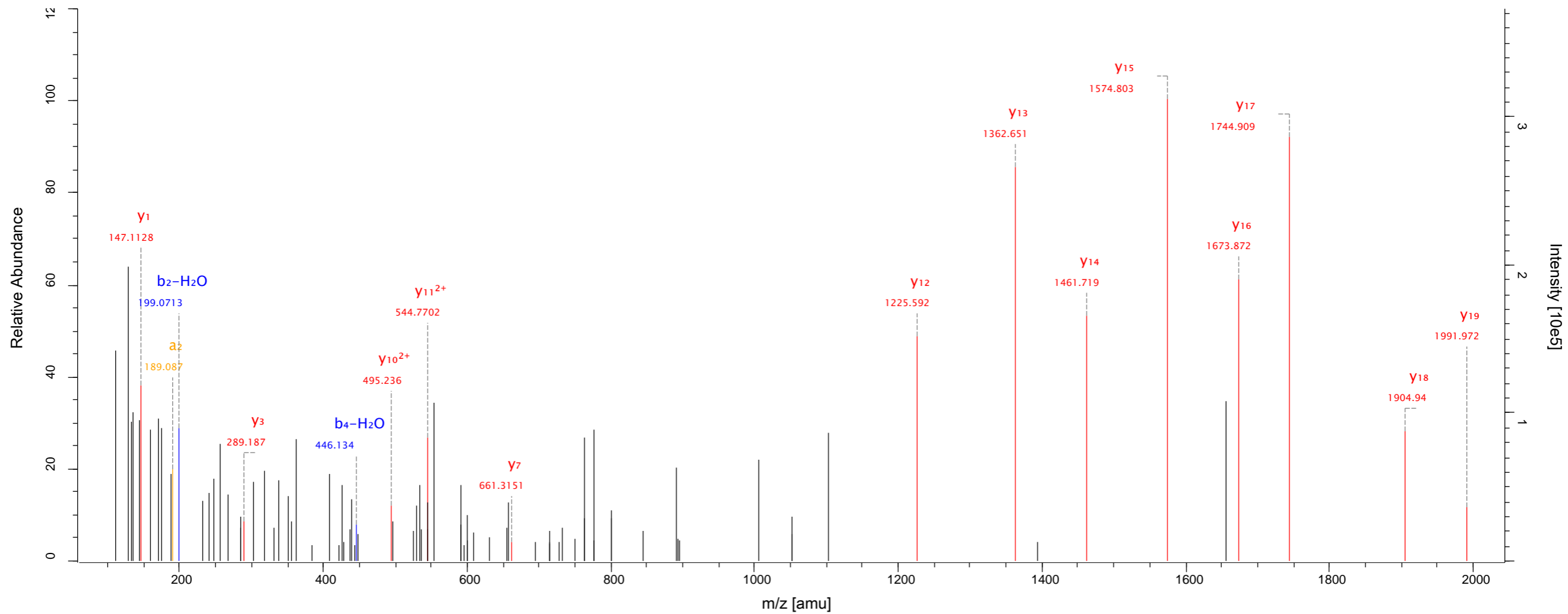
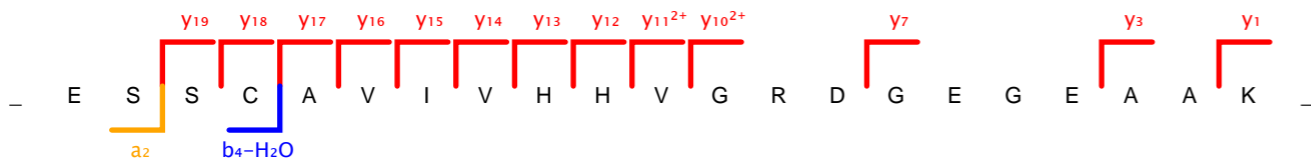
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
 Scan number: 11246
 Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
 Peptide score: 174.16
 Method: FTMS; HCD



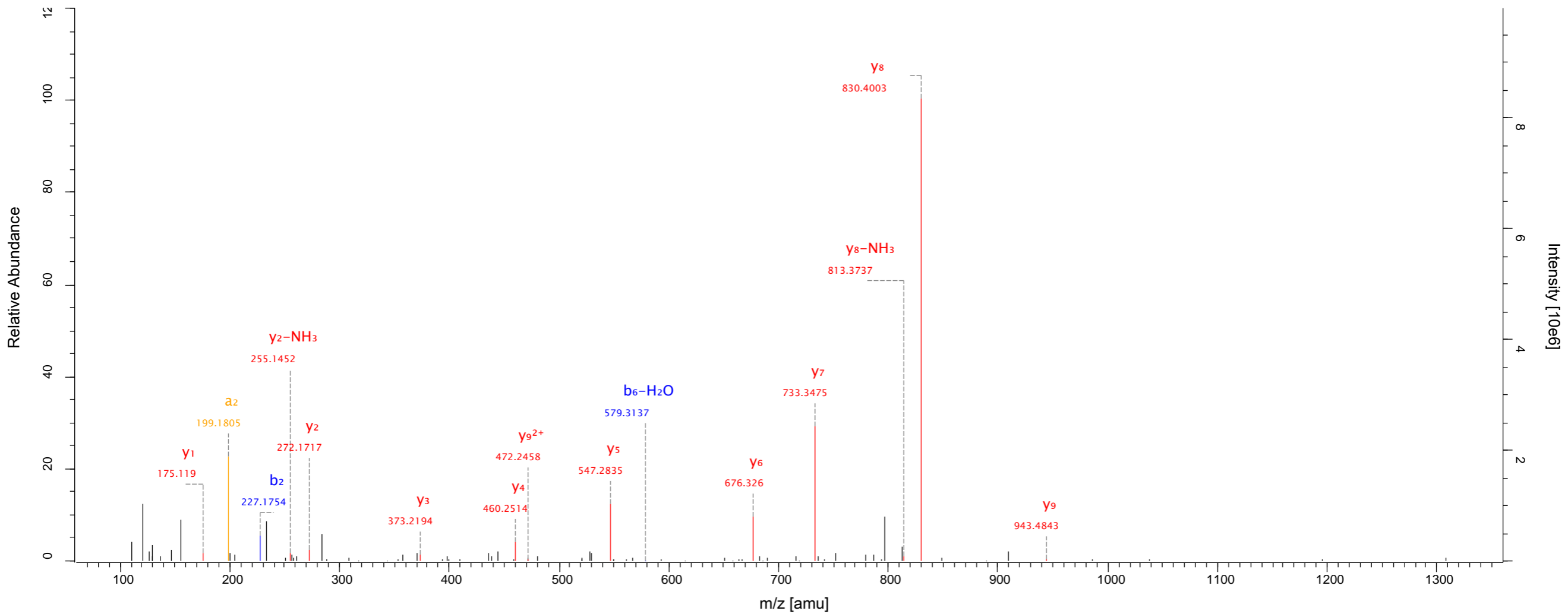
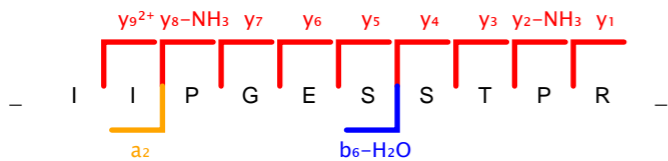
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 11864
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 112.13
Method: FTMS; HCD



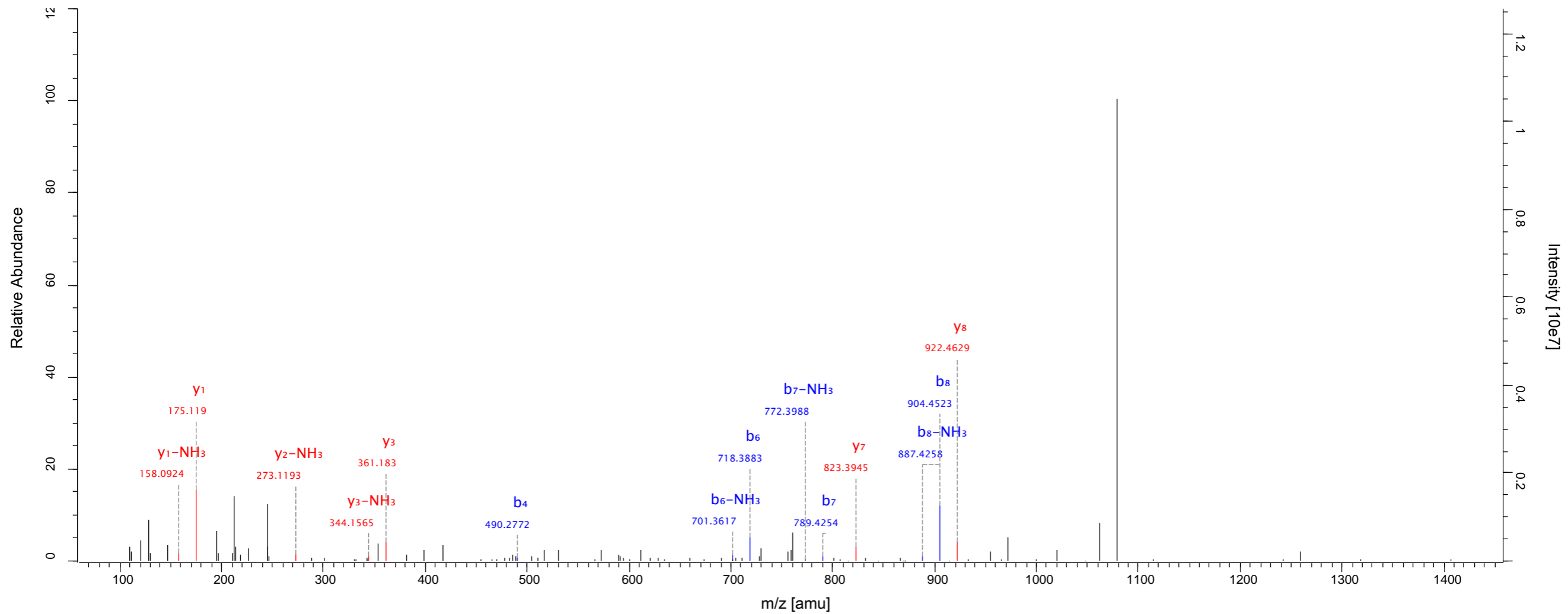
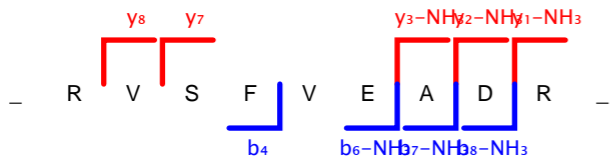
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 11968
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 44.38
Method: FTMS; HCD



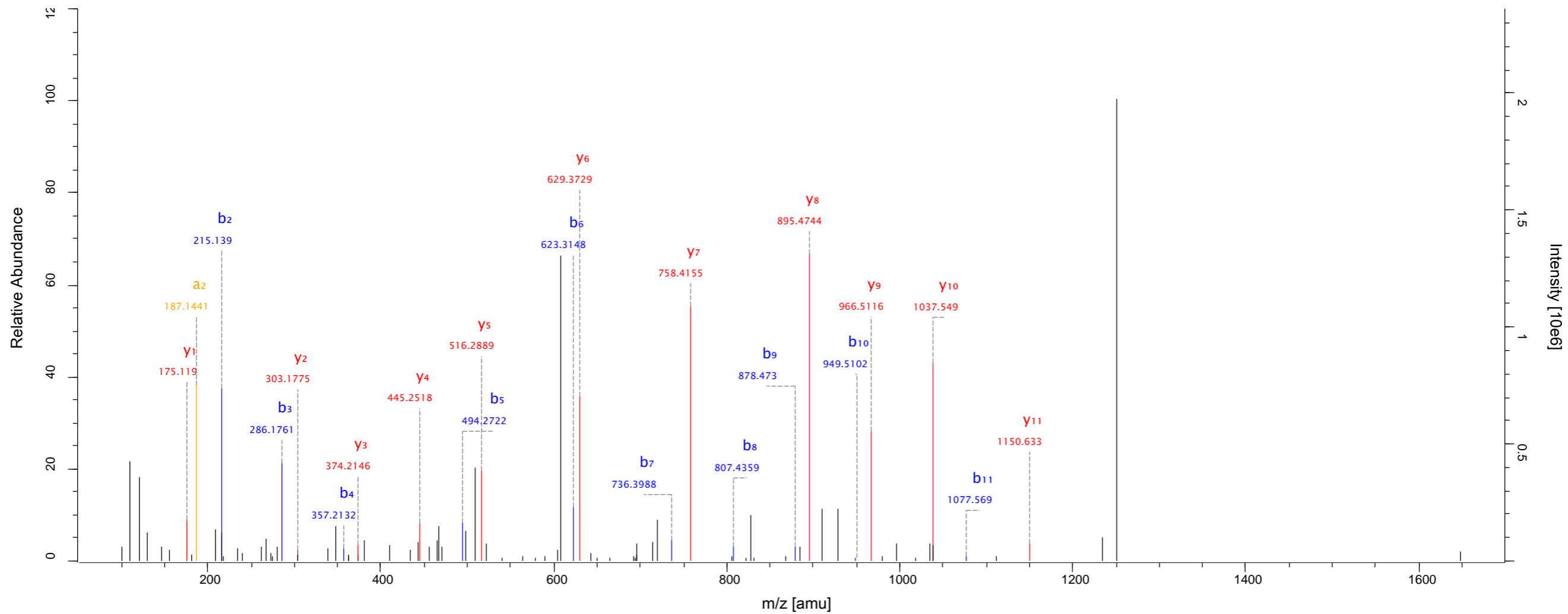
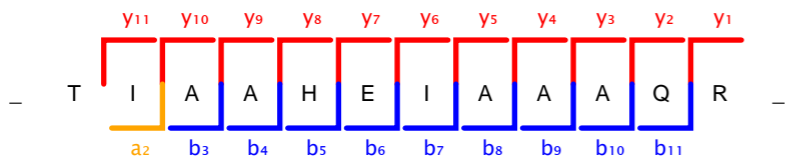
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 12344
Protein: ORFL137W.iORF1 + NC_006273.2[75299,75529]+ <unknown description>
Peptide score: 66.96
Method: FTMS; HCD



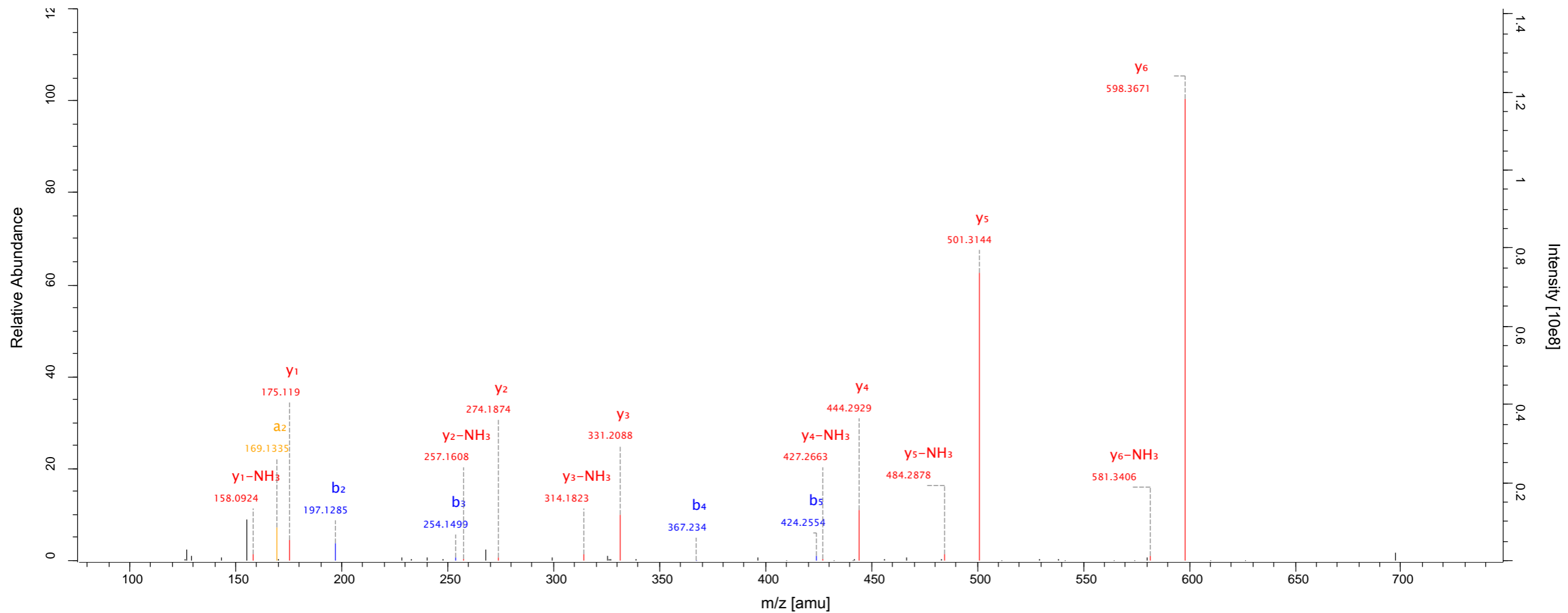
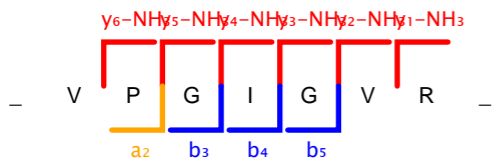
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 12581
Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>; ORFL27C.iORF1 - NC_006273.2[7912,8184]- <unknown description>
Peptide score: 55.46
Method: FTMS; HCD



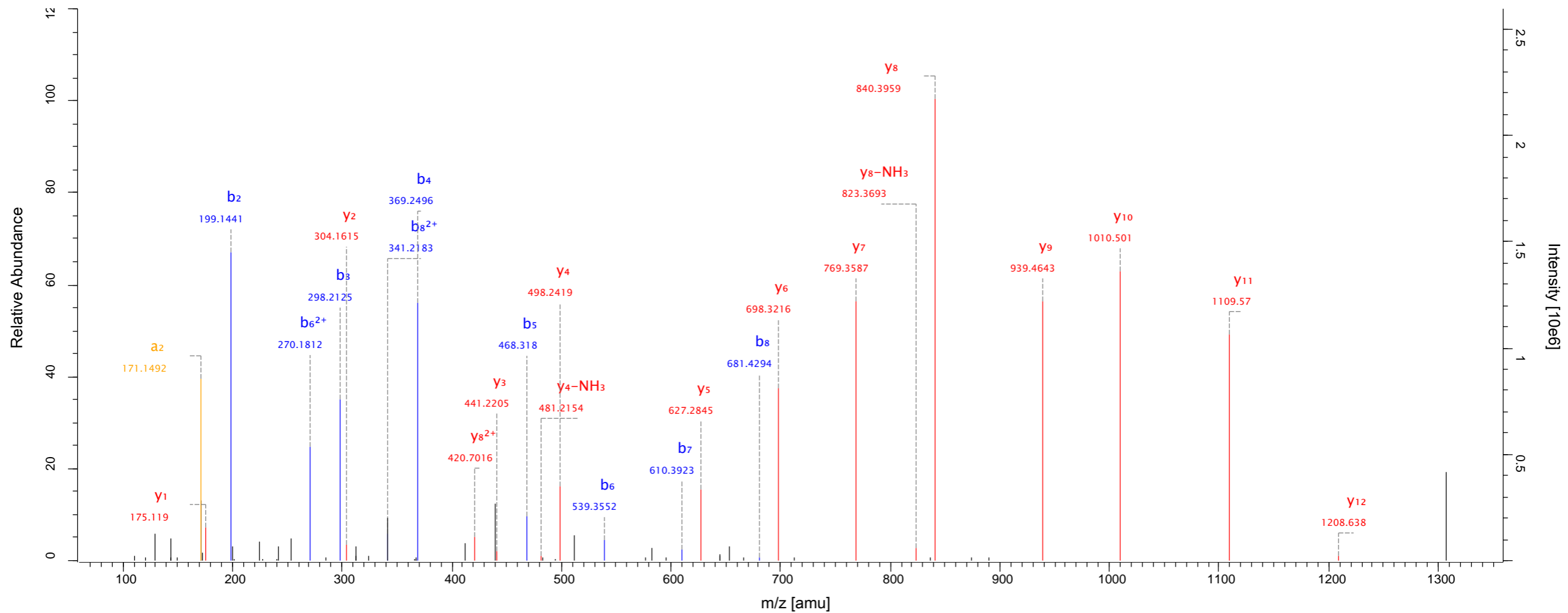
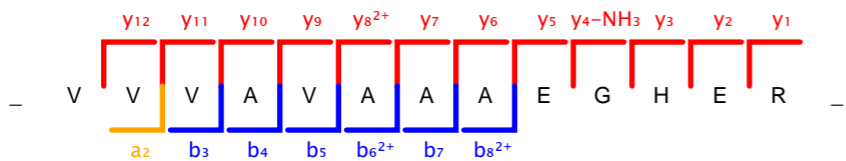
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 13004
Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
Peptide score: 118.87
Method: FTMS; HCD



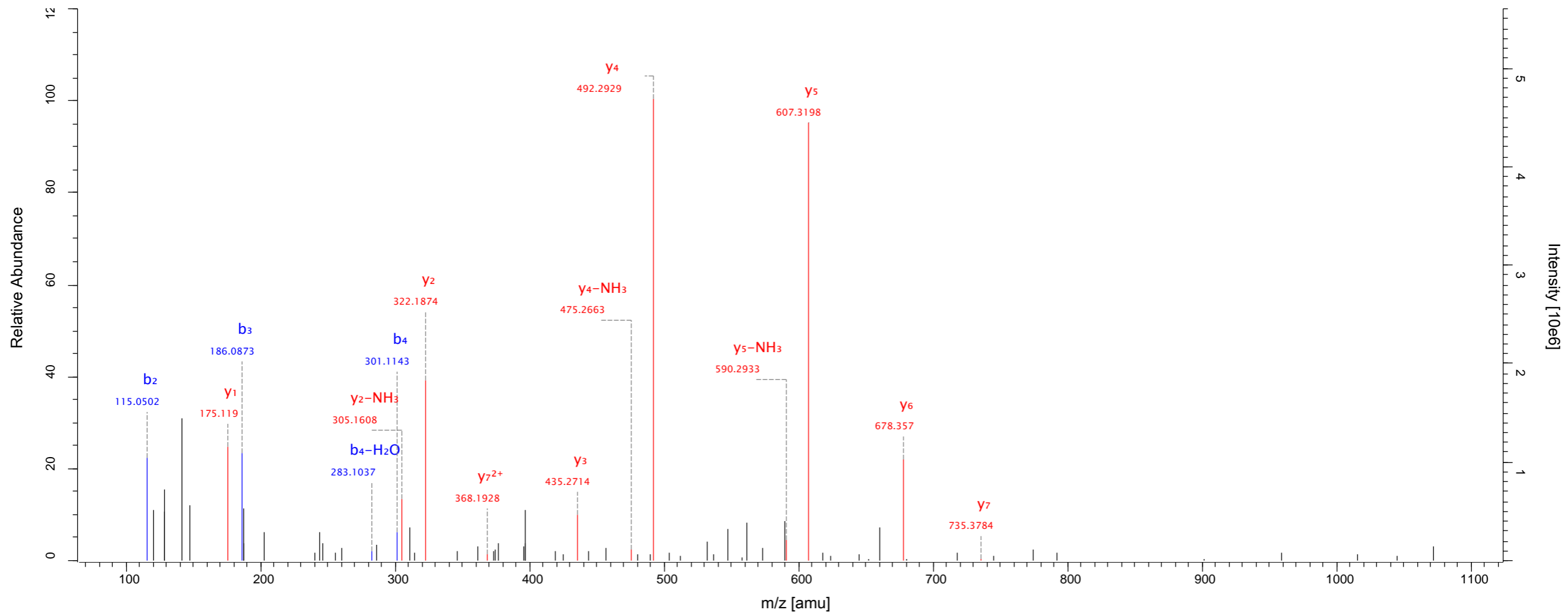
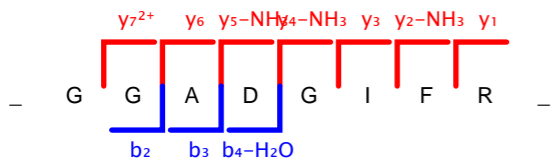
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 14551
Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
Peptide score: 166.66
Method: FTMS; HCD



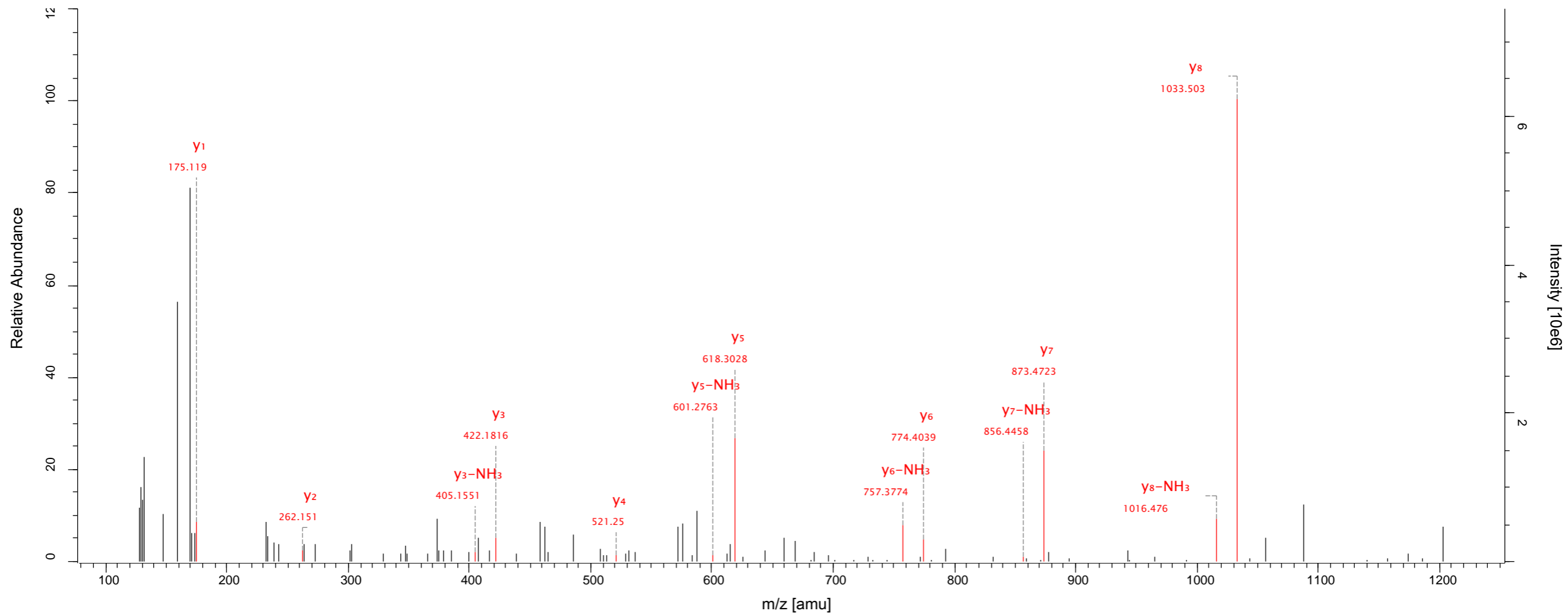
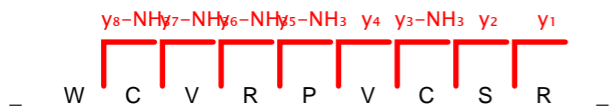
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 14782
Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
Peptide score: 164.65
Method: FTMS; HCD



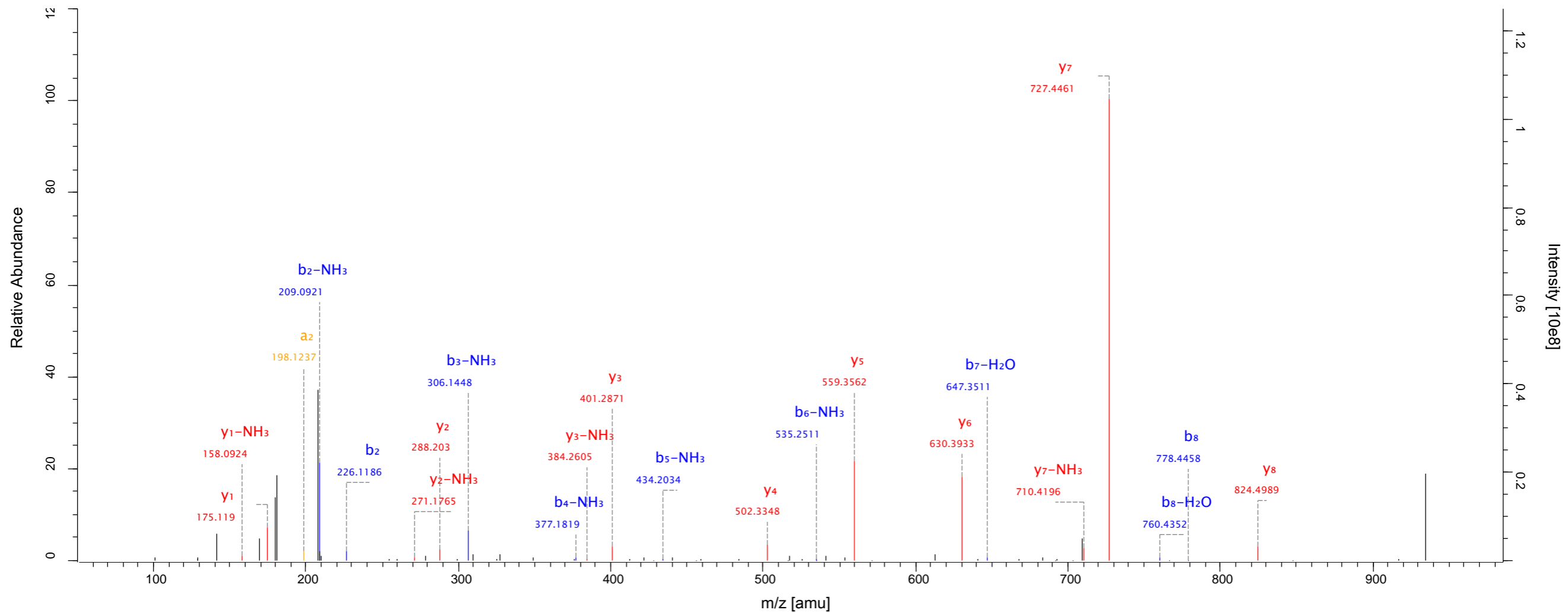
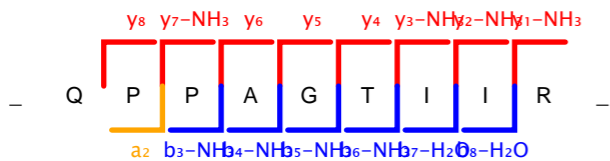
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 15914
Protein: ORFL159W + NC_006273.2[101475,109052]+ <unknown description>; ORFL159W.iORF1 + NC_006273.2[101526,109052]+ <unknown description>
Peptide score: 101.64
Method: FTMS; HCD



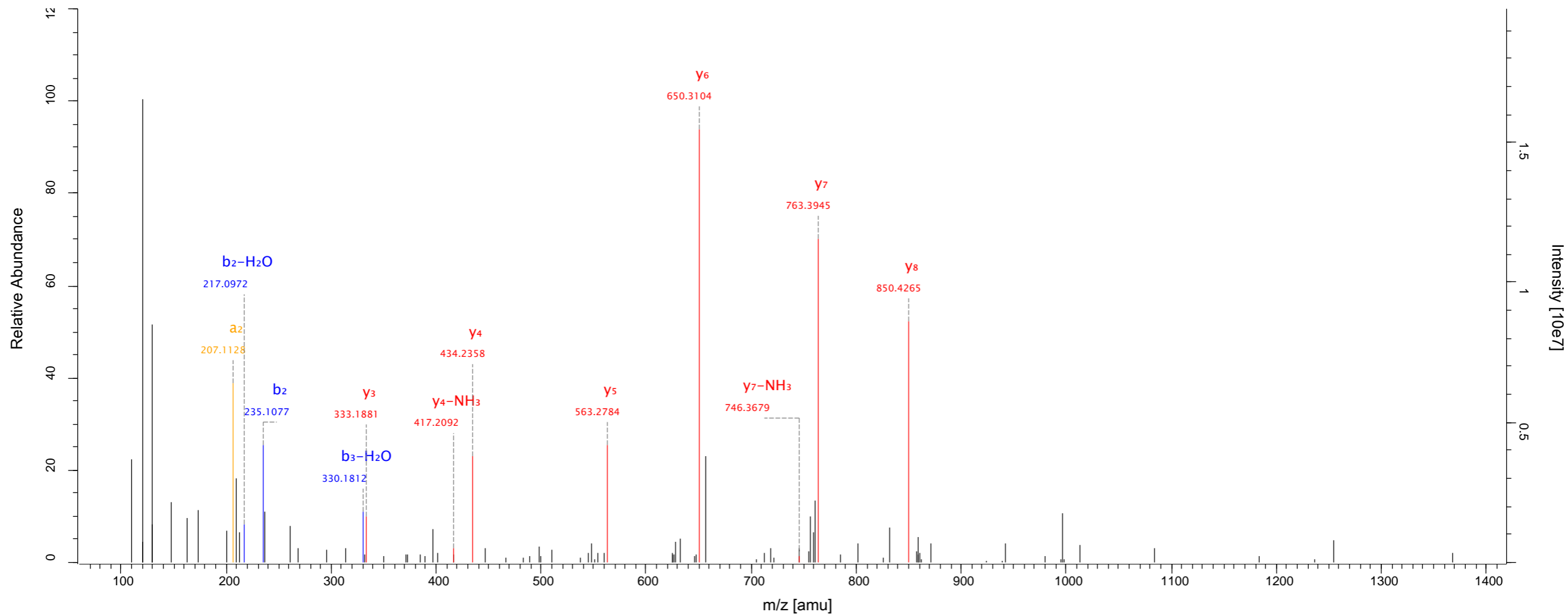
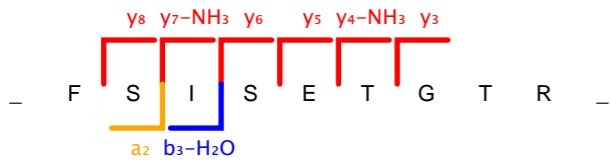
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 15937
Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
Peptide score: 61.58
Method: FTMS; HCD



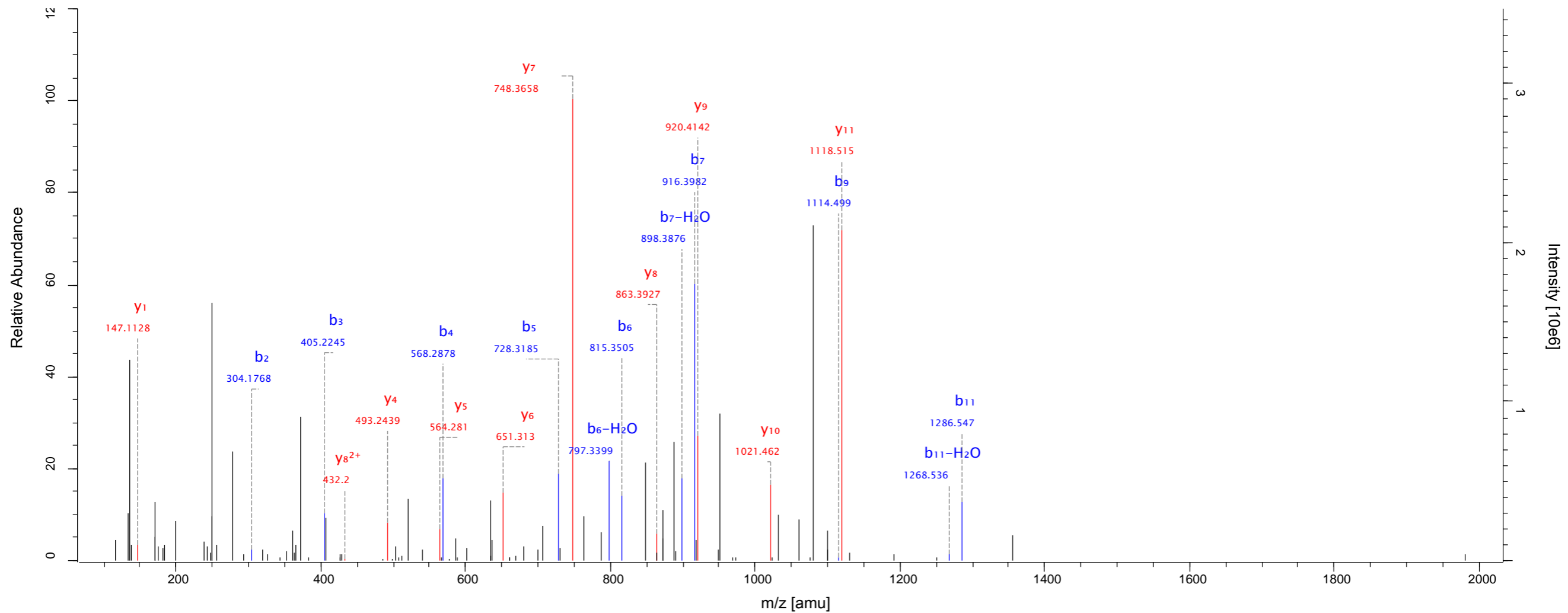
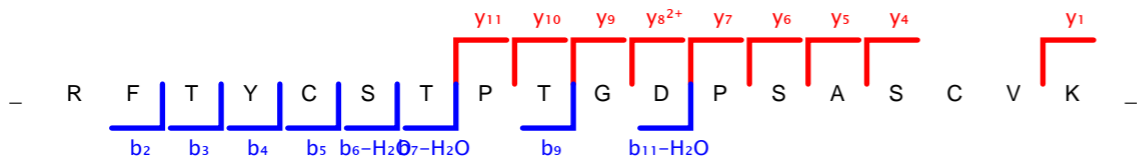
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 16732
Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
Peptide score: 101.25
Method: FTMS; HCD



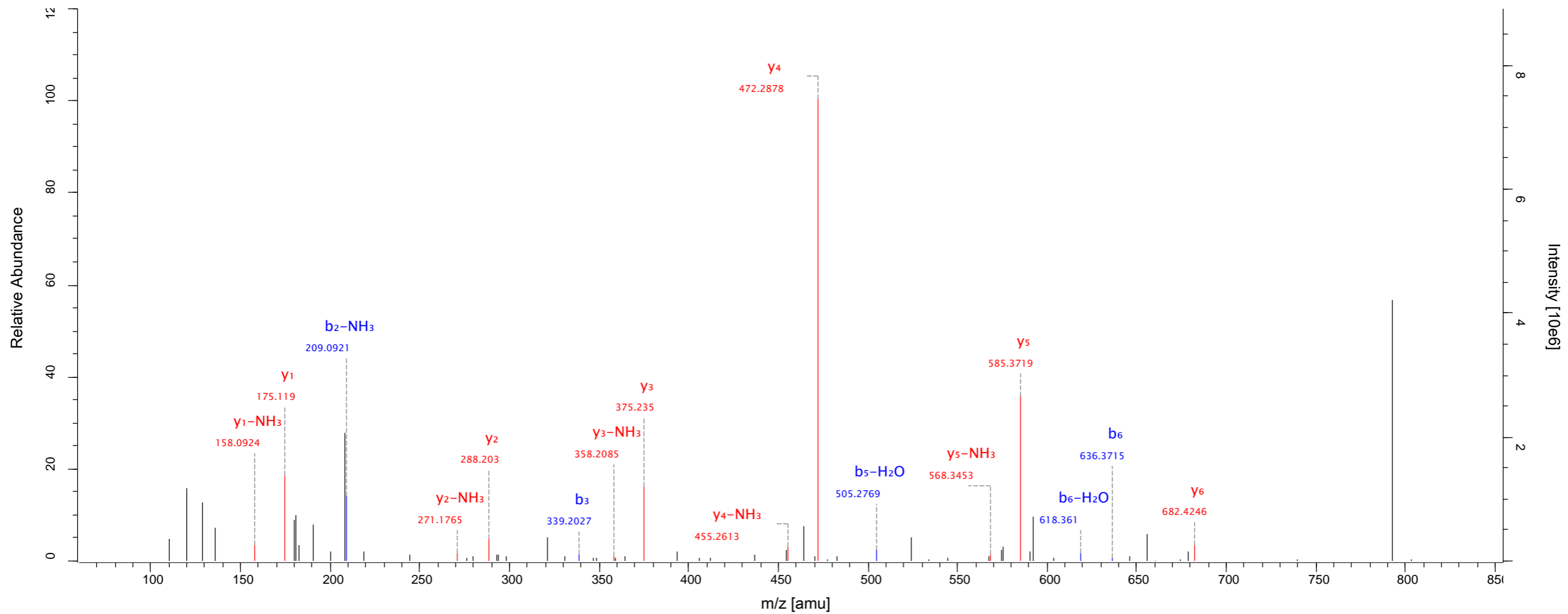
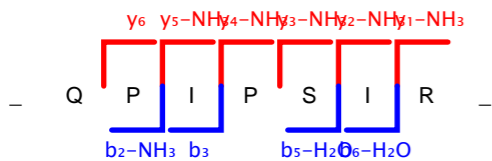
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 16867
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 71.22
Method: FTMS; HCD



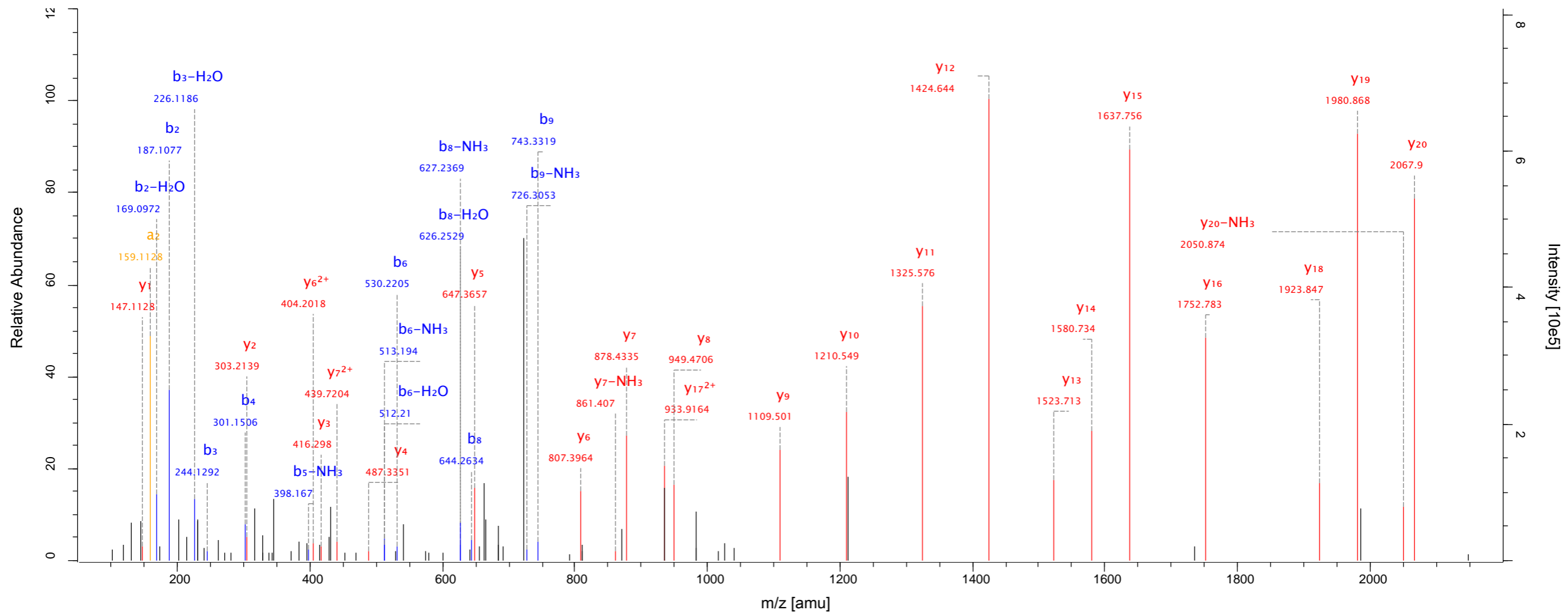
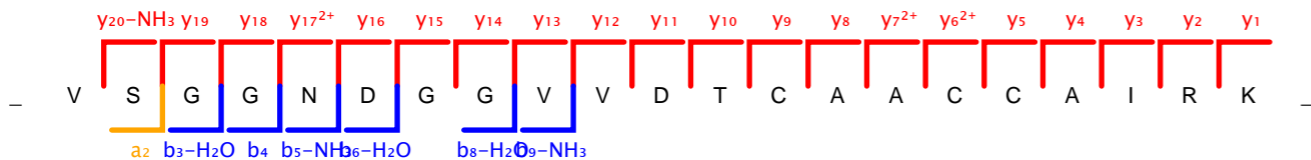
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
 Scan number: 17553
 Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
 Peptide score: 85.64
 Method: FTMS; HCD



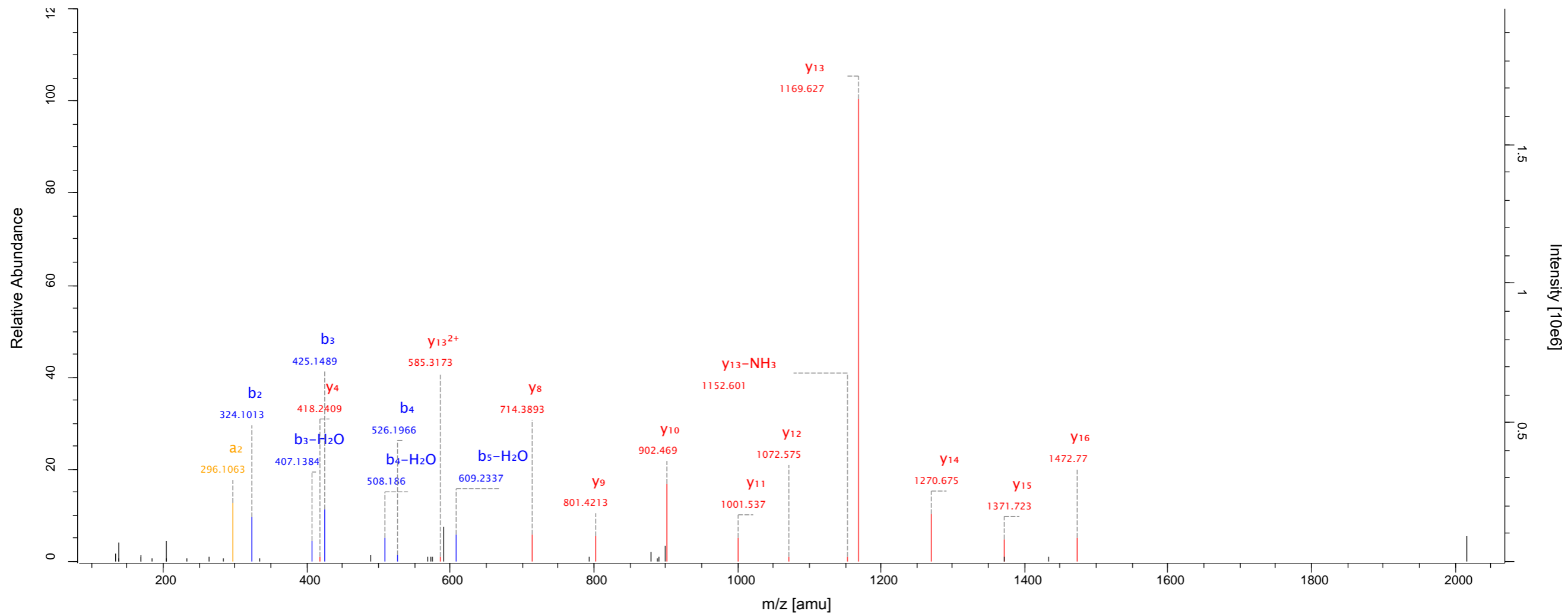
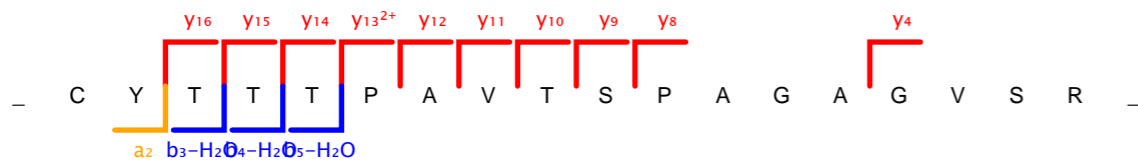
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 17637
Protein: ORFL137W.IORF1 + NC_006273.2[75299,75529]+ <unknown description>
Peptide score: 90.71
Method: FTMS; HCD



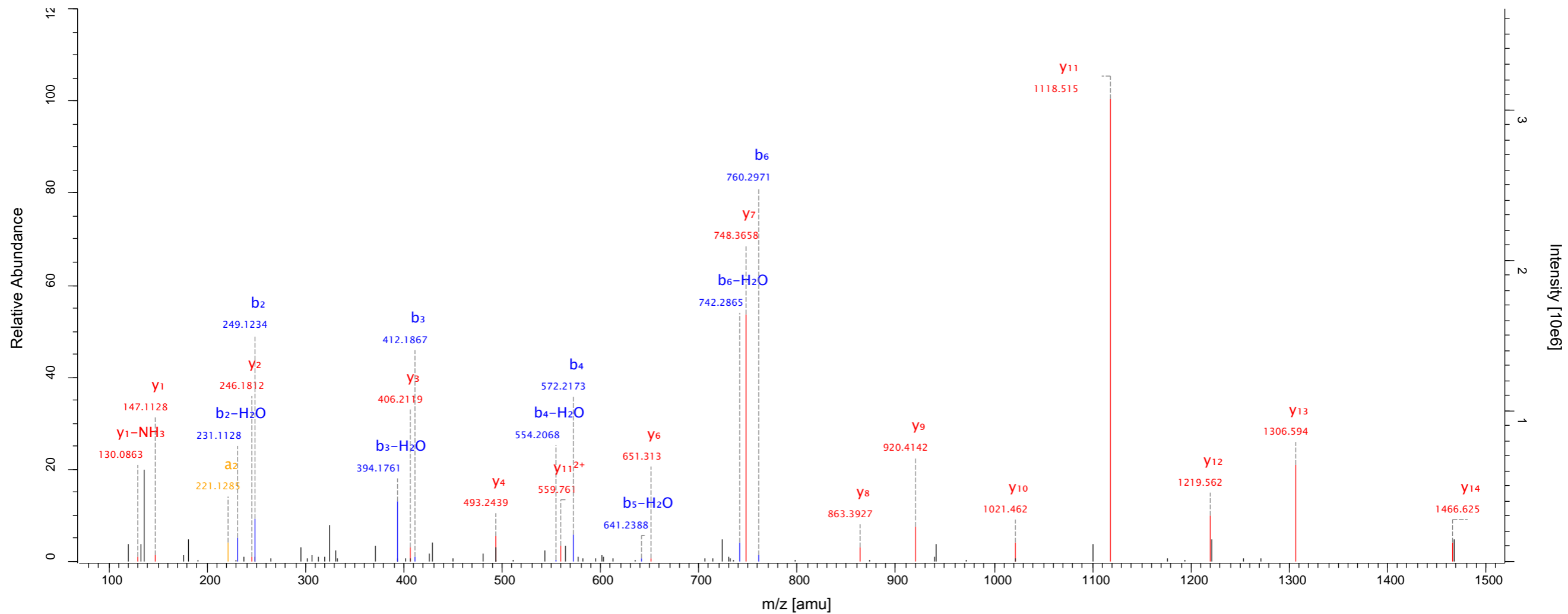
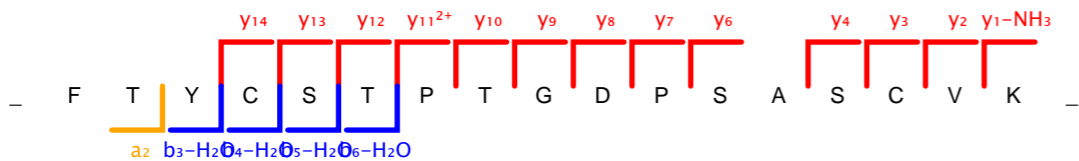
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 18964
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 180.14
 Method: FTMS; HCD



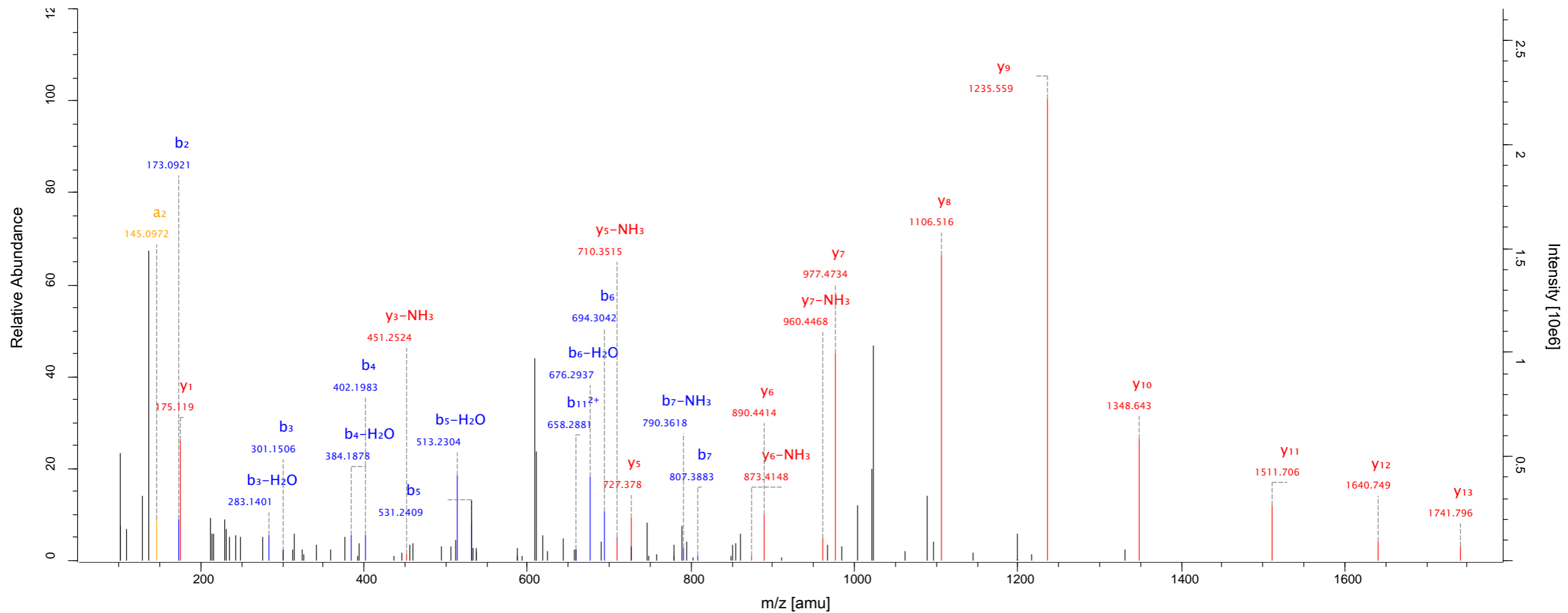
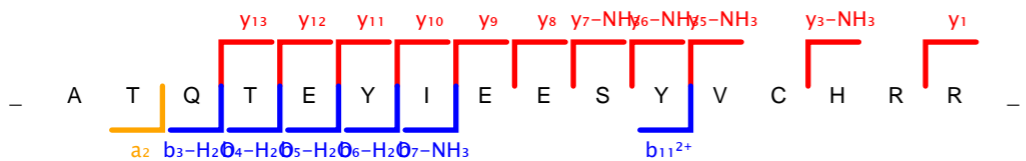
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 19350
Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
Peptide score: 122.34
Method: FTMS; HCD



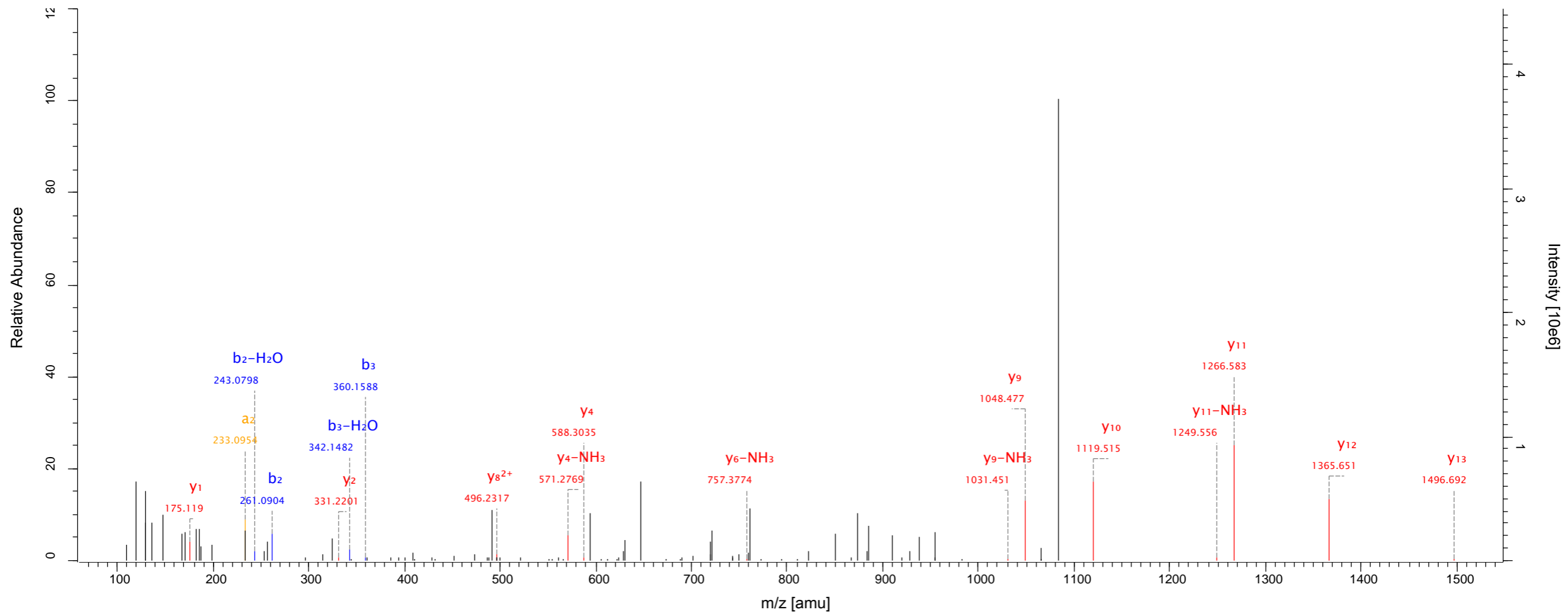
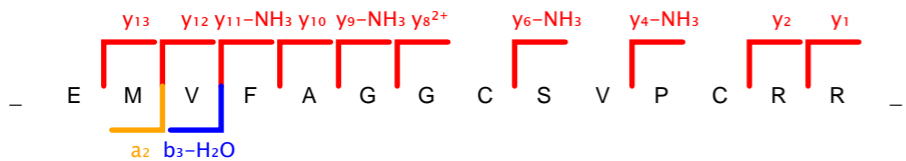
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 19625
 Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
 Peptide score: 109.83
 Method: FTMS; HCD



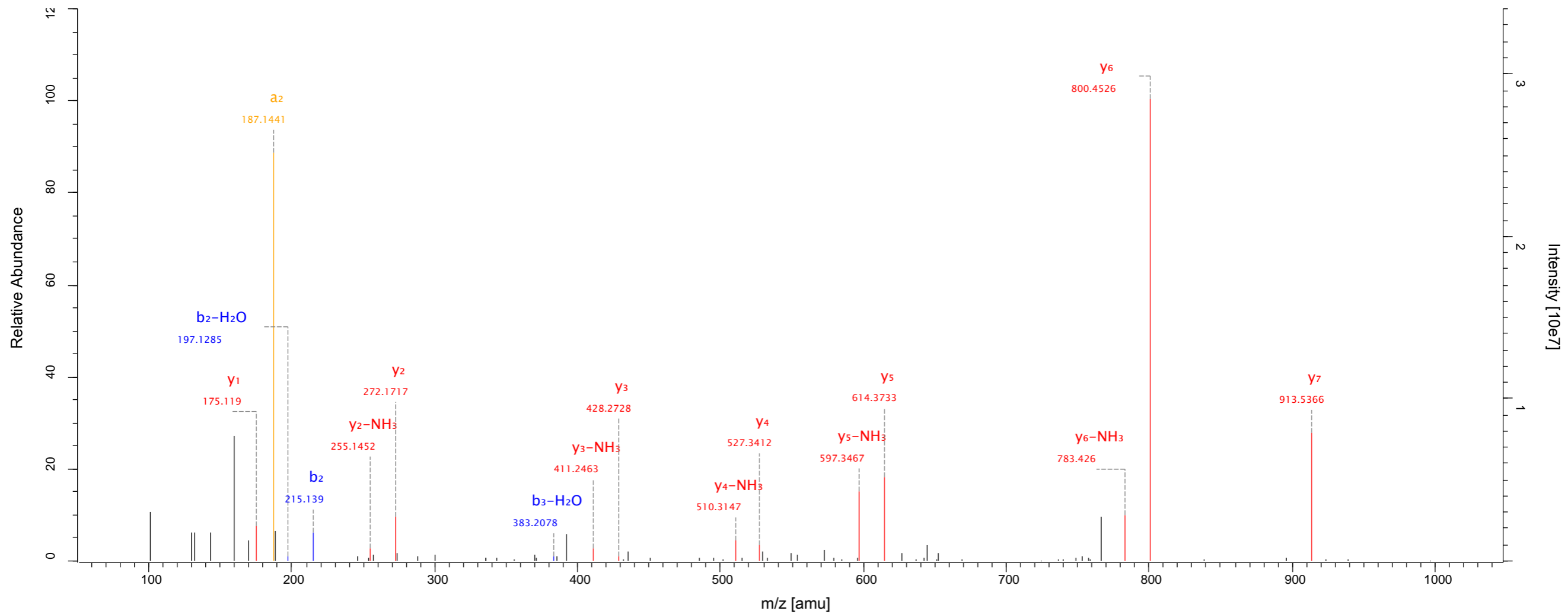
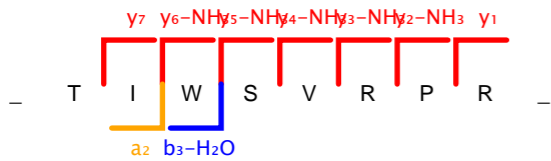
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
 Scan number: 19739
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 121.31
 Method: FTMS; HCD



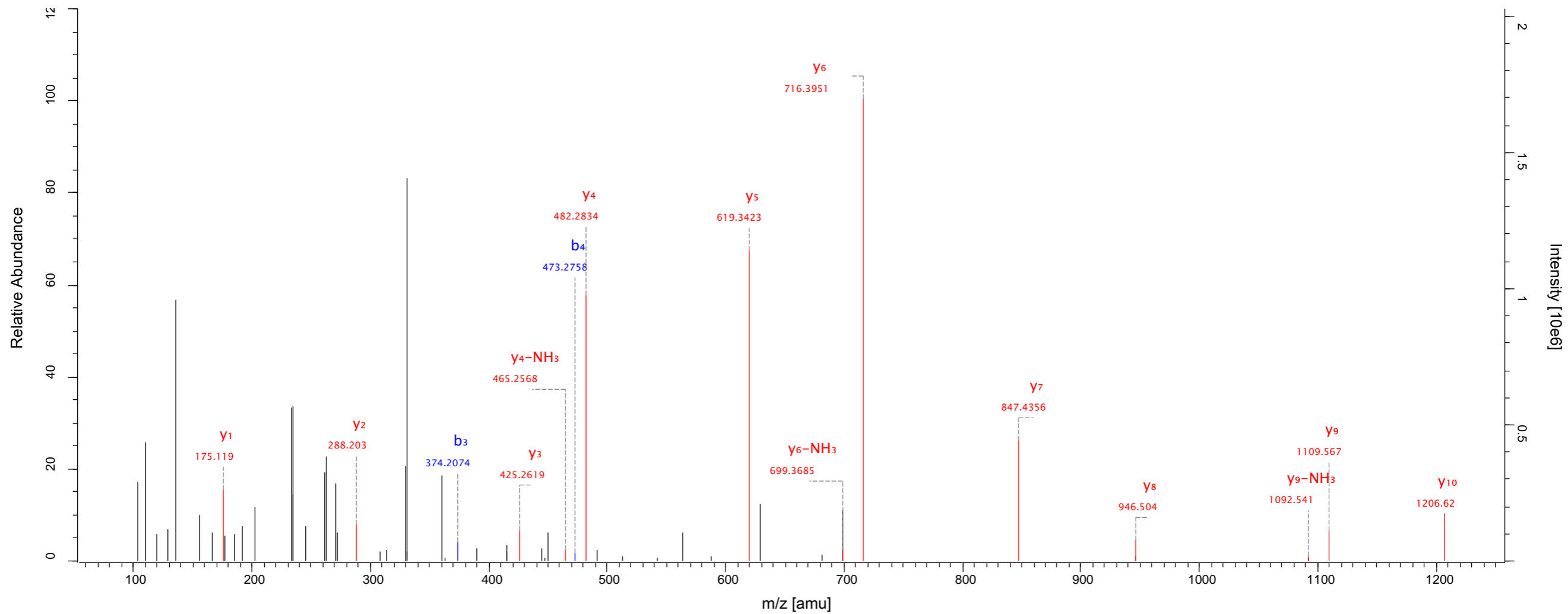
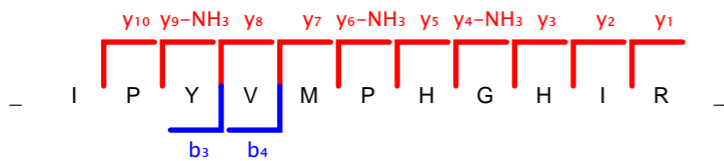
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 20243
 Protein: ORFL159W + NC_006273.2[101475,109052]+ <unknown description>; ORFL159W.iORF1 + NC_006273.2[101526,109052]+ <unknown description>
 Peptide score: 71.24
 Method: FTMS; HCD



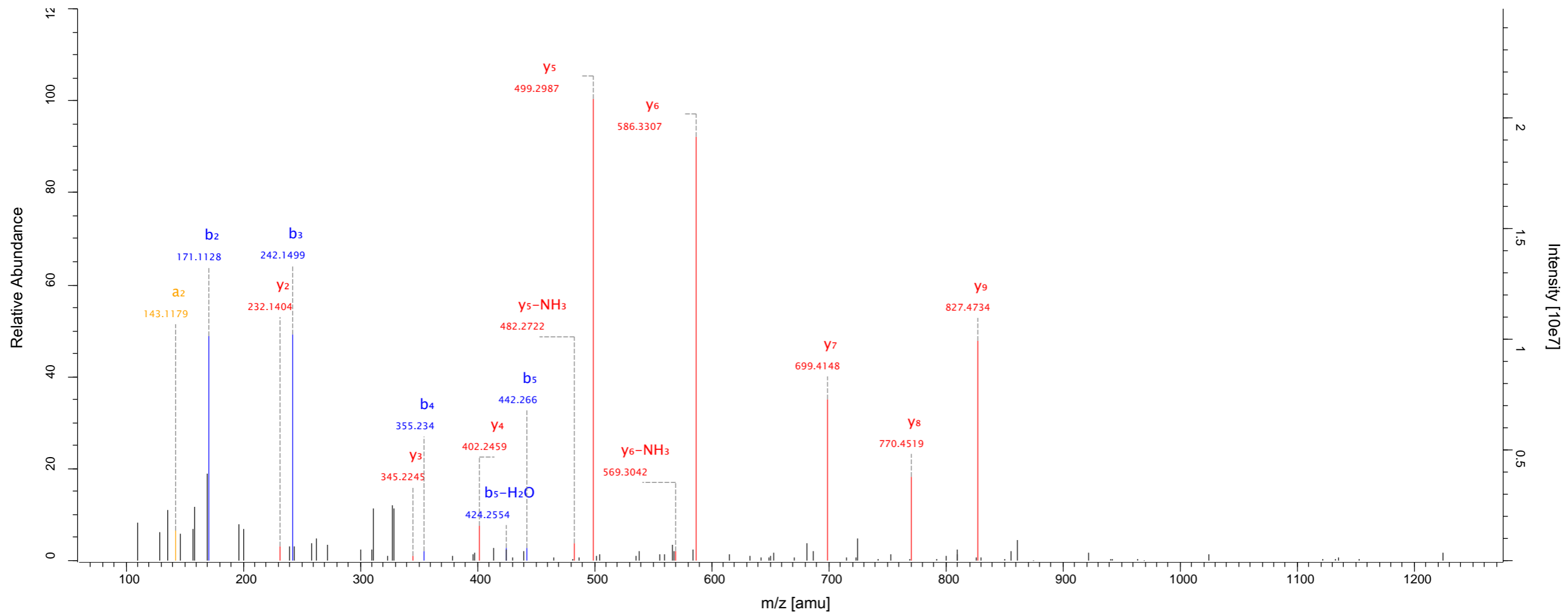
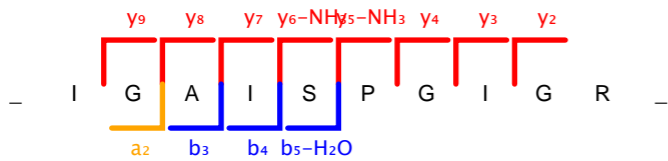
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 20338
Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>
Peptide score: 127.84
Method: FTMS; HCD



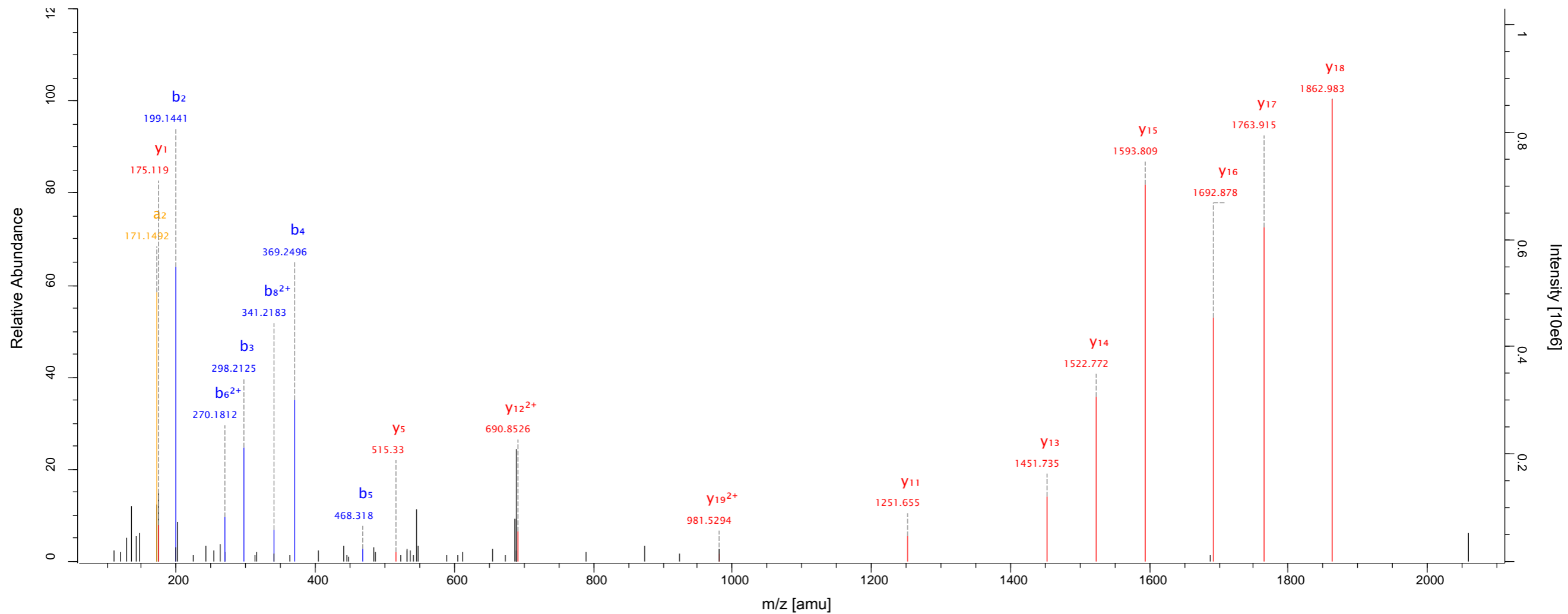
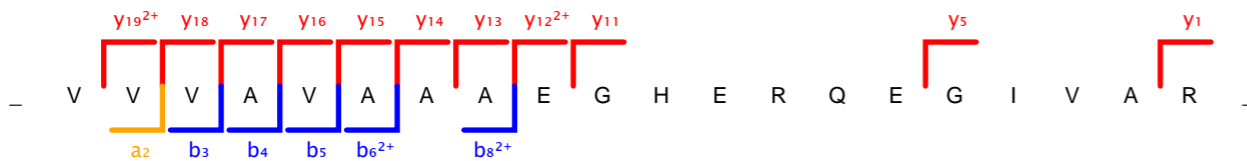
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 21294
Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>
Peptide score: 83
Method: FTMS; HCD



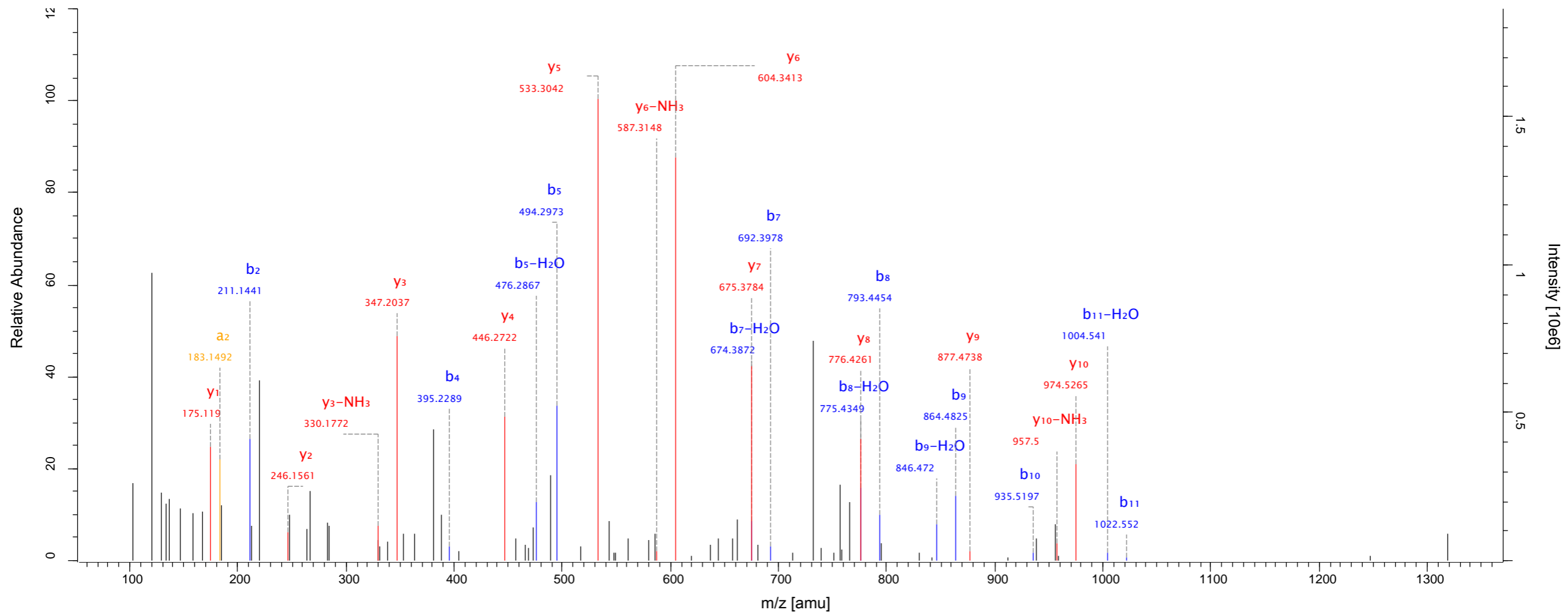
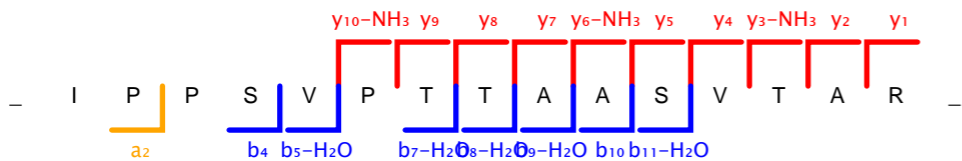
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 21409
Protein: ORFL154W + NC_006273.2[96843,97115]+ <unknown description>; ORFL154W.iORF1 + NC_006273.2[96972,97115]+ <unknown description>
Peptide score: 89.25
Method: FTMS; HCD



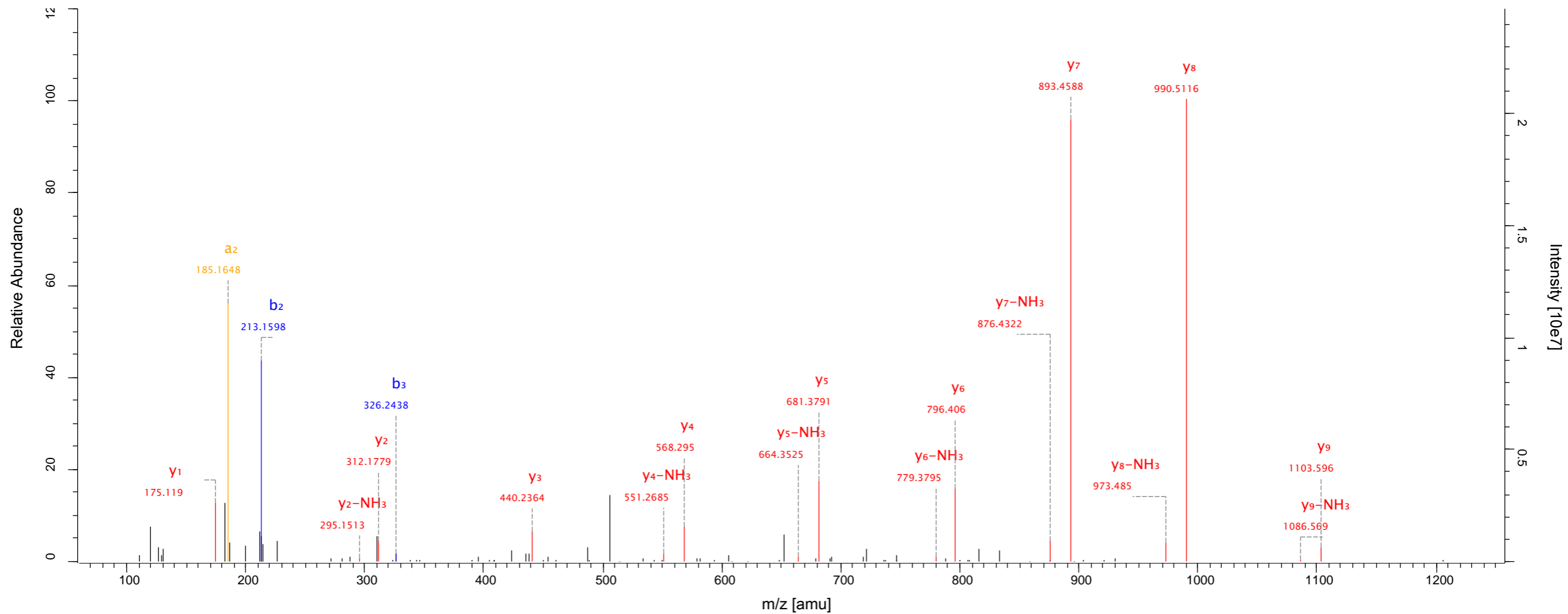
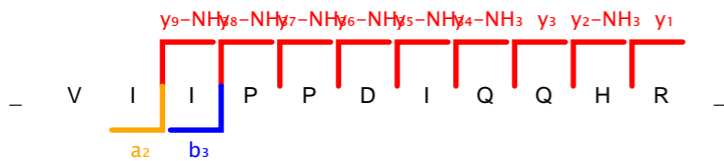
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 22393
 Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
 Peptide score: 84.87
 Method: FTMS; HCD



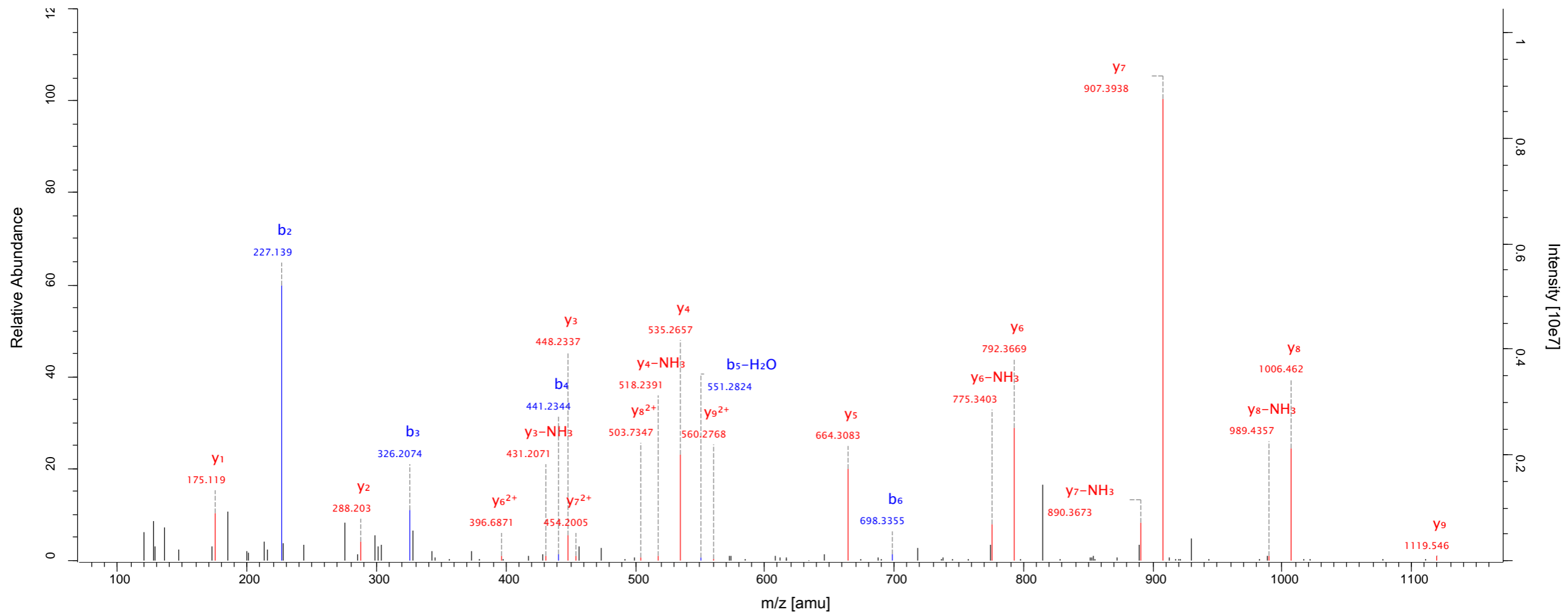
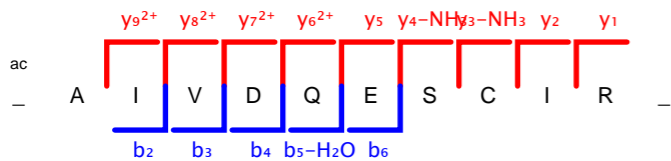
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 23131
 Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
 Peptide score: 138.02
 Method: FTMS; HCD



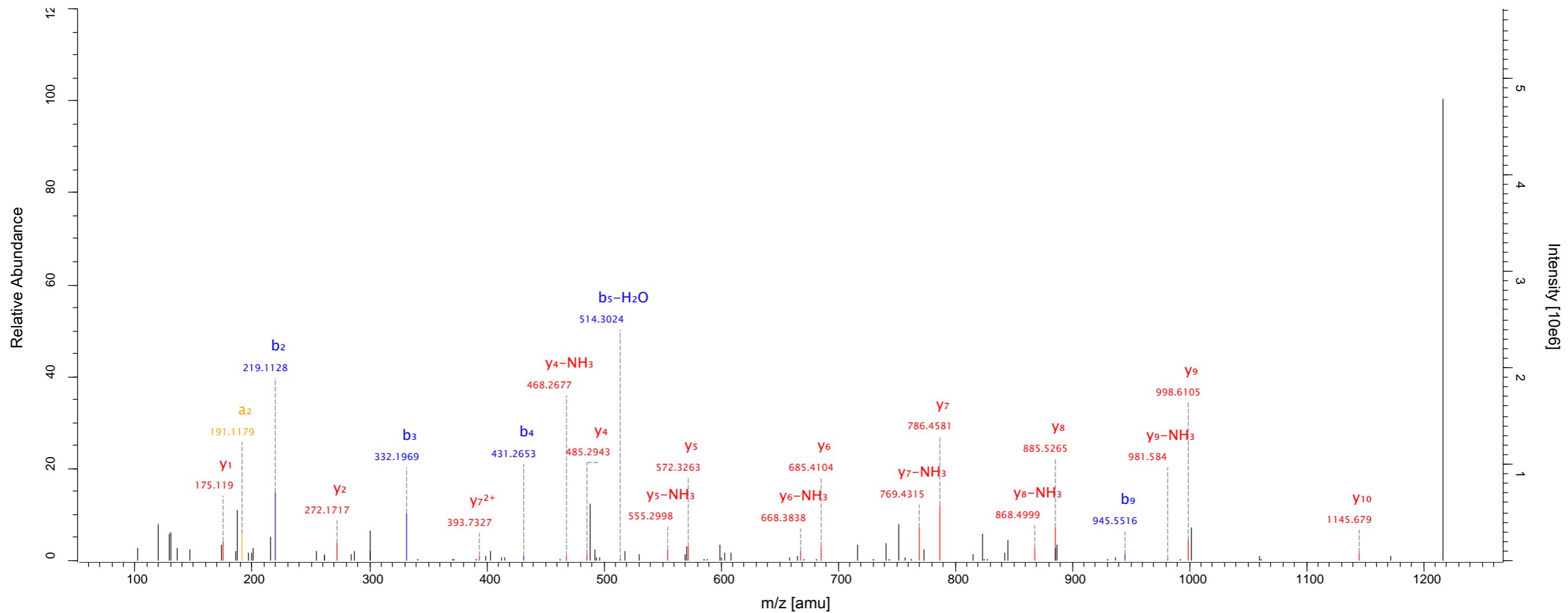
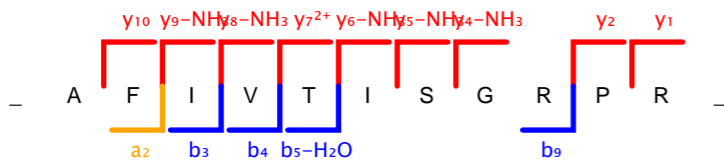
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 25309
Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
Peptide score: 147.2
Method: FTMS; HCD



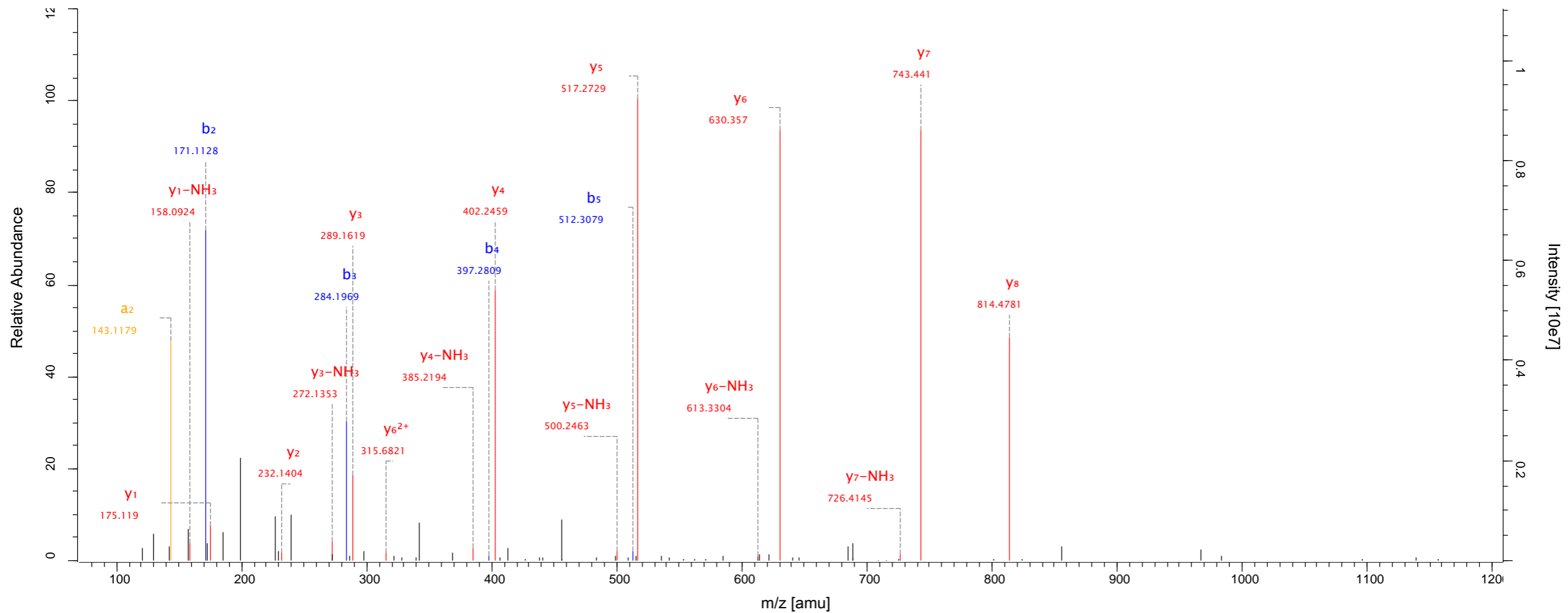
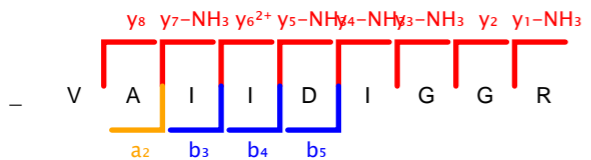
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 27594
Protein: ORFL151C - NC_006273.2[92499,92870]- <unknown description>
Peptide score: 133.13
Method: FTMS; HCD



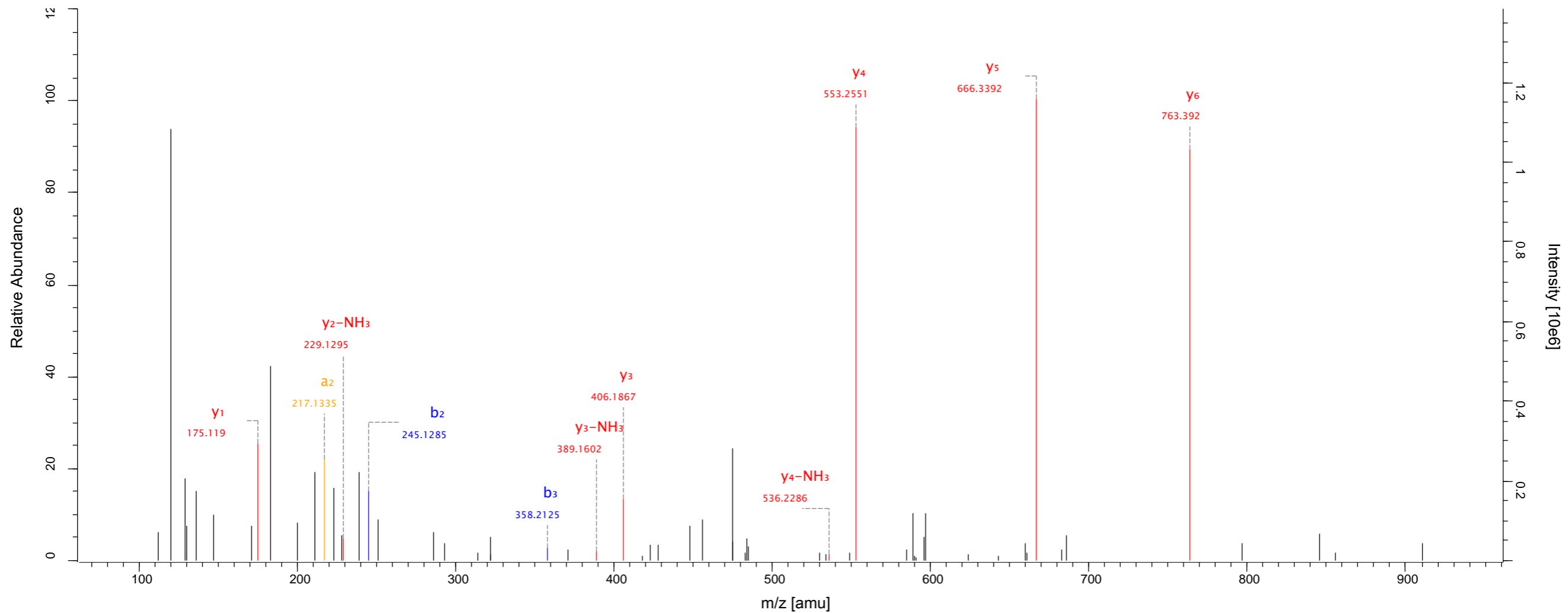
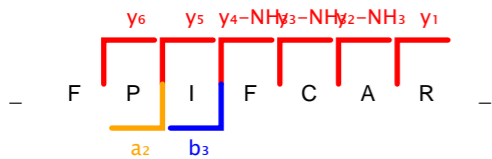
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 28094
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 137.95
Method: FTMS; HCD



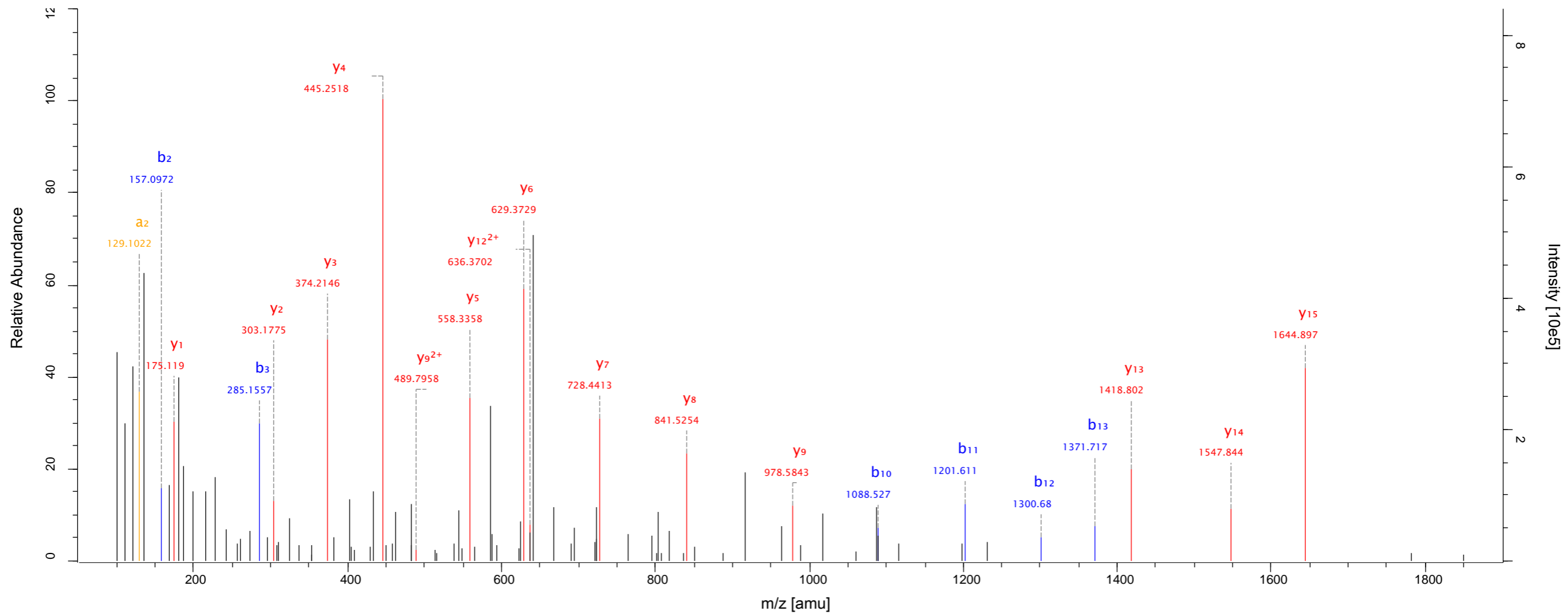
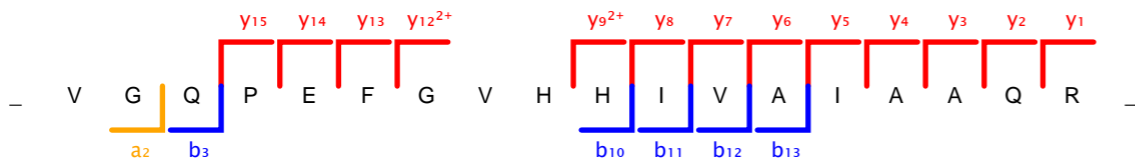
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 32927
Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
Peptide score: 143.89
Method: FTMS; HCD



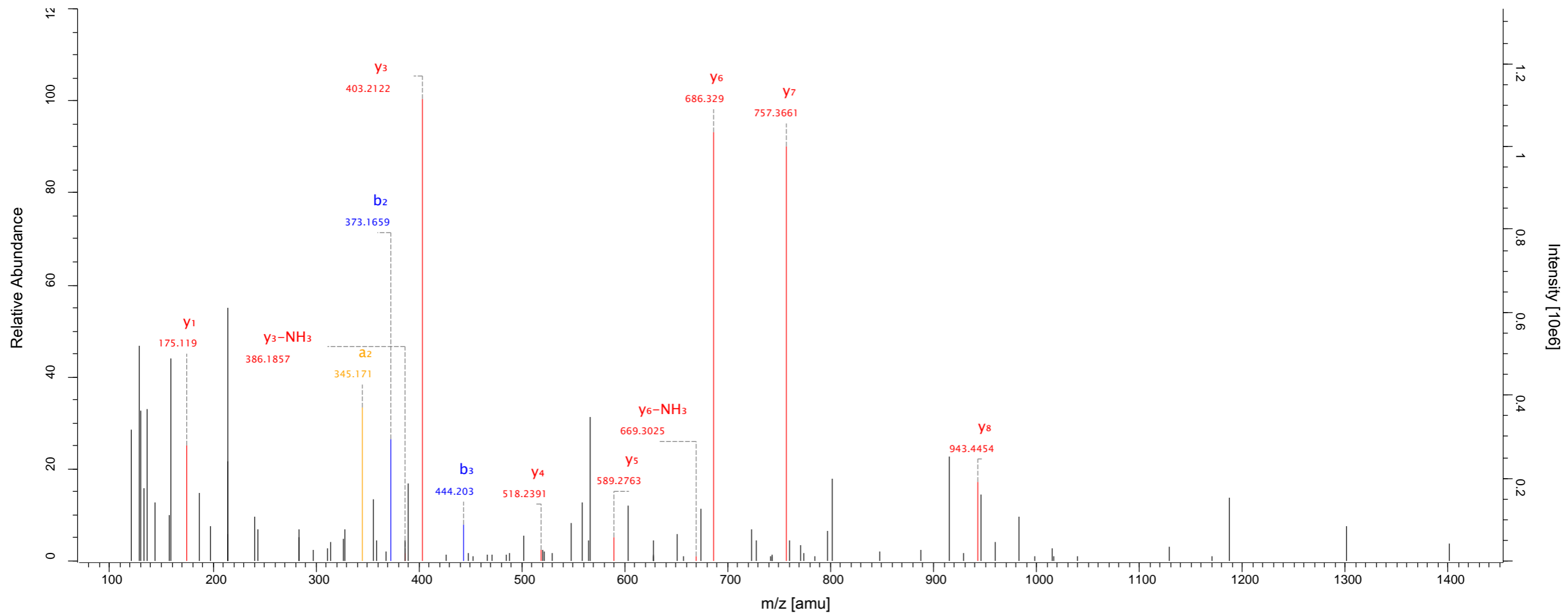
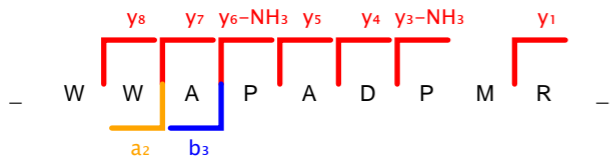
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 33368
Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
Peptide score: 74.72
Method: FTMS; HCD



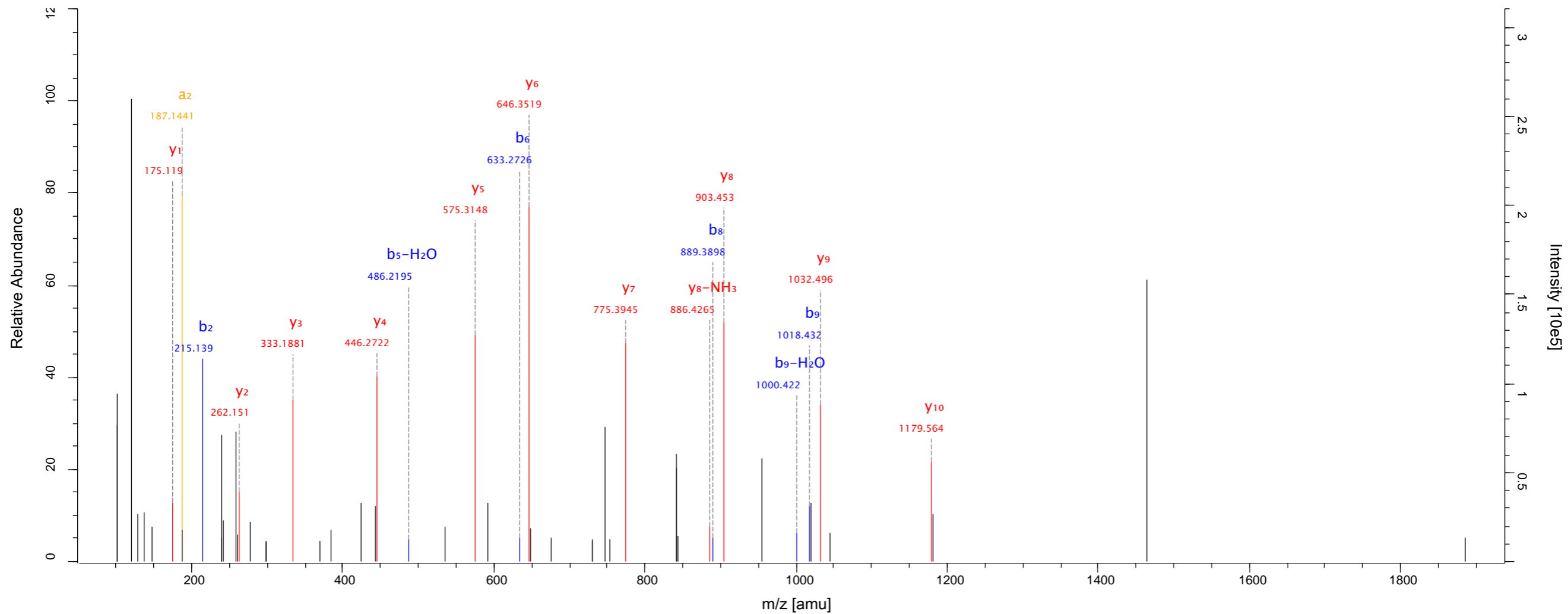
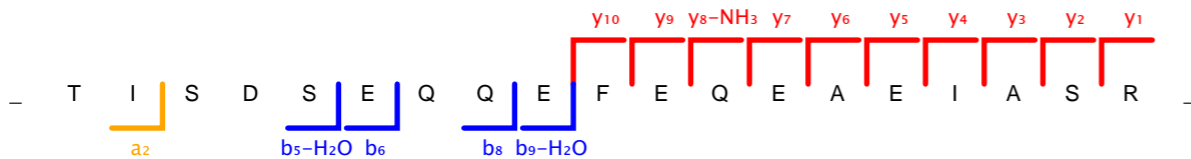
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 34748
Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
Peptide score: 113.69
Method: FTMS; HCD



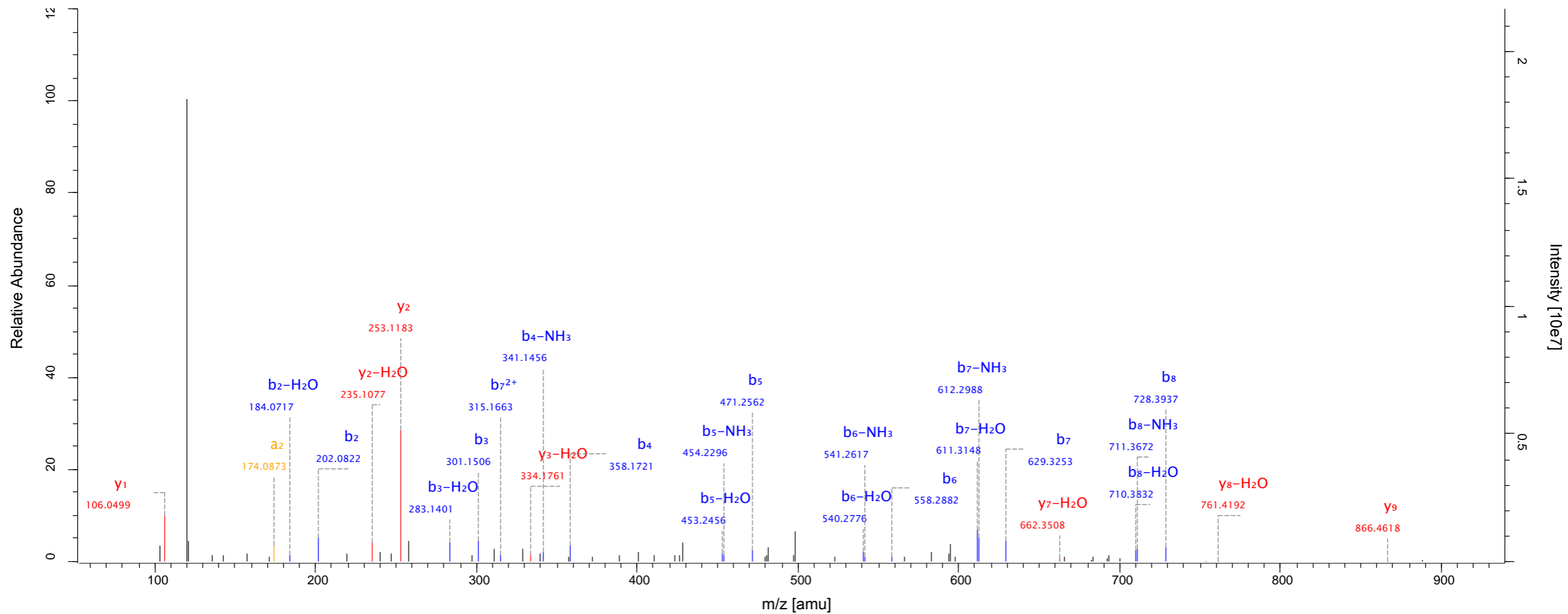
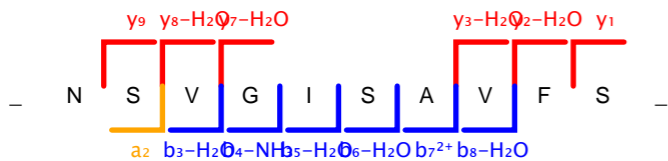
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 35064
Protein: ORFL253W.iORF3 + NC_006273.2[162059,162286]+ <unknown description>
Peptide score: 56.81
Method: FTMS; HCD



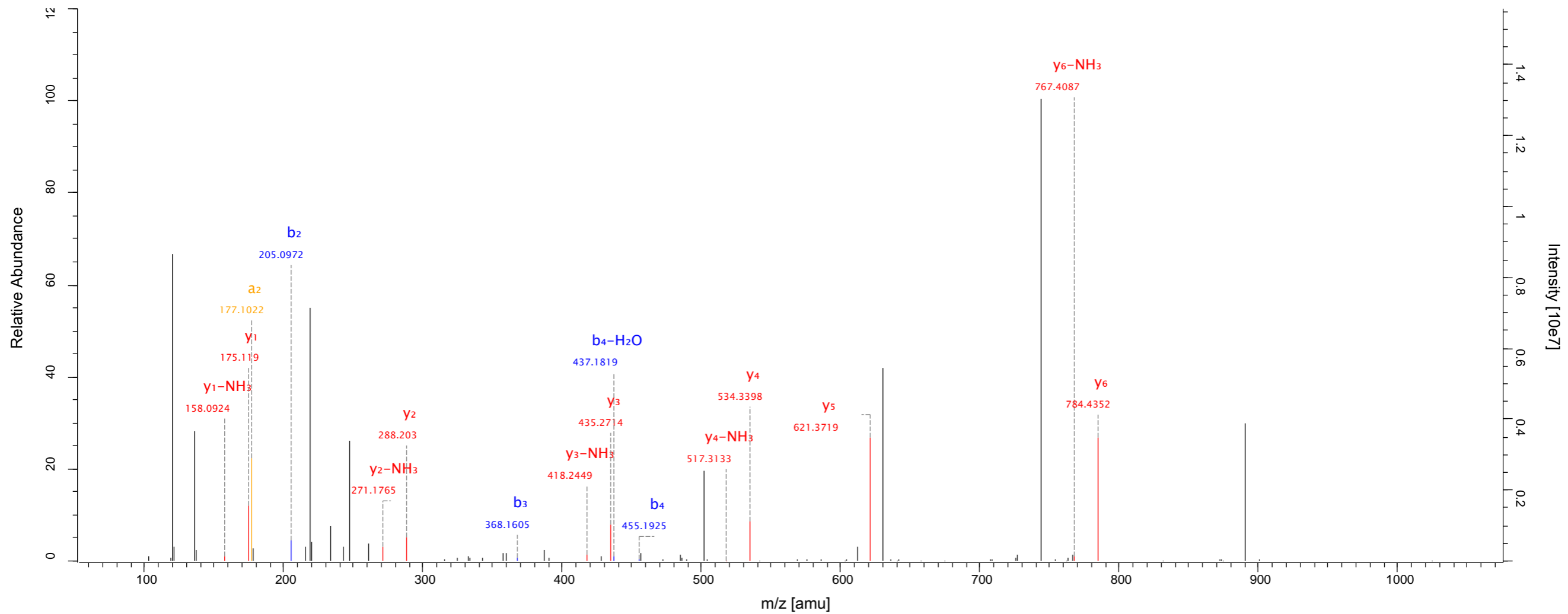
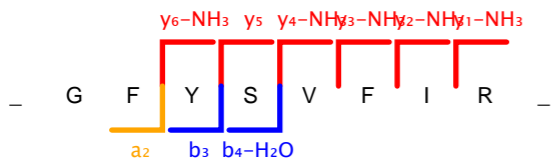
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 36428
 Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>
 Peptide score: 63.69
 Method: FTMS; HCD



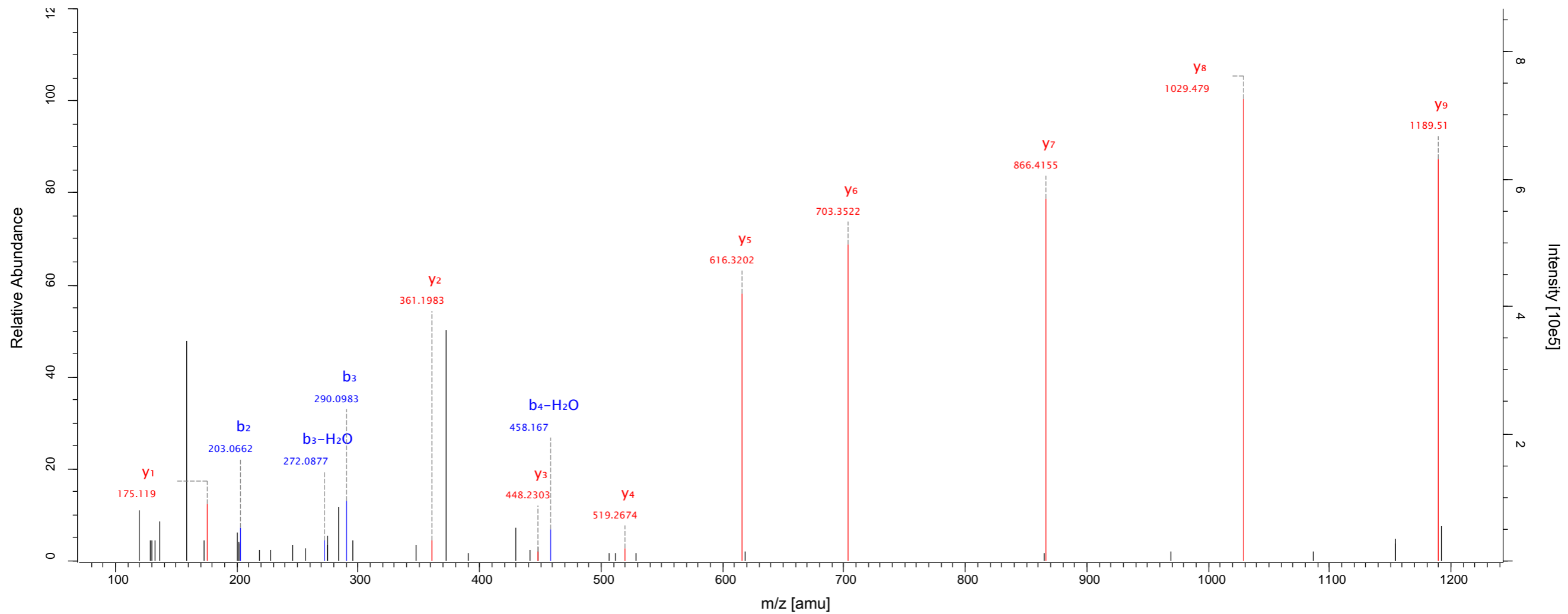
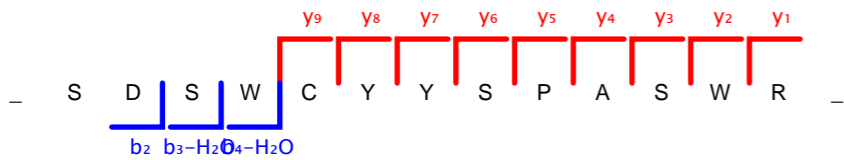
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 38904
 Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>; ORFL27C.iORF1 - NC_006273.2[7912,8184]- <unknown description>
 Peptide score: 166.34
 Method: FTMS; HCD



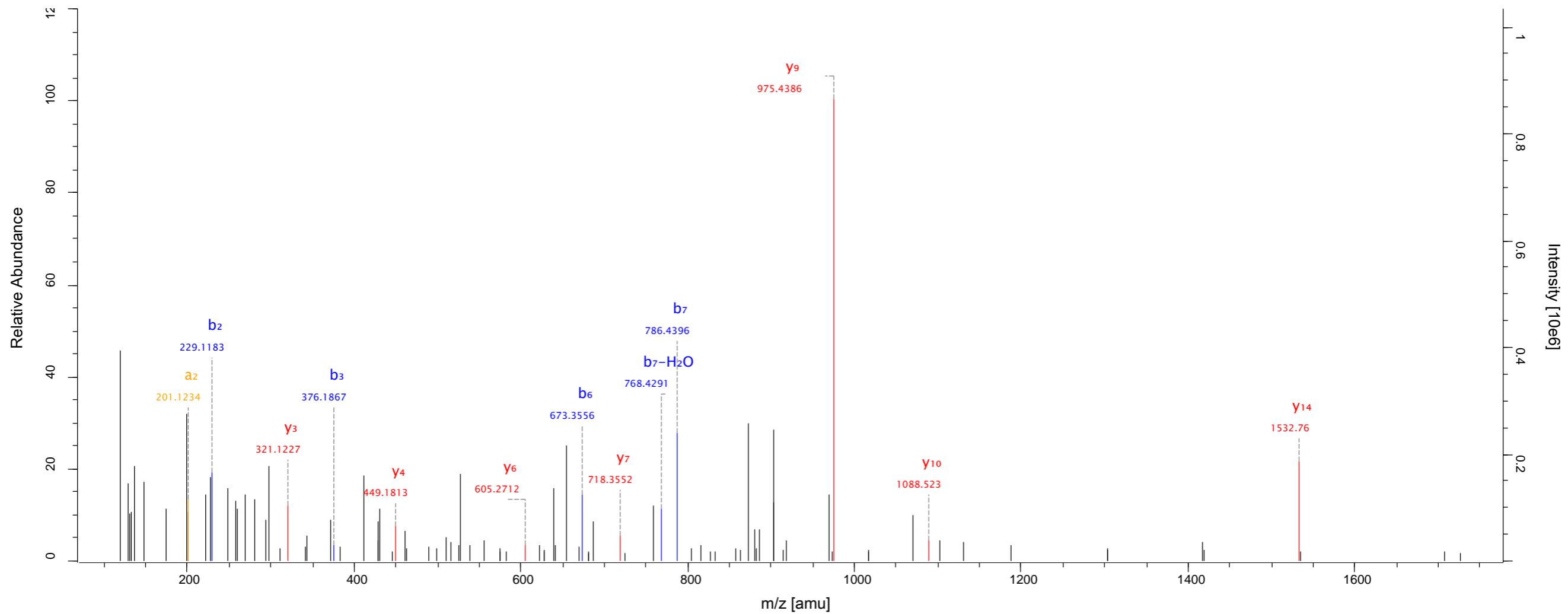
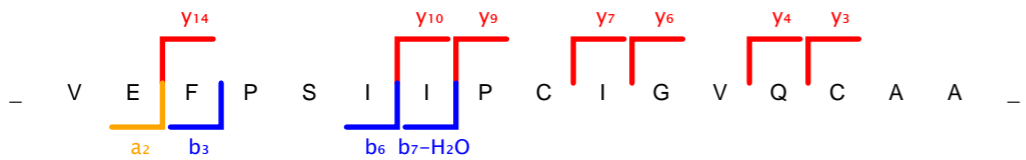
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 42885
Protein: ORFL151C - NC_006273.2[92499,92870]- <unknown description>
Peptide score: 107.67
Method: FTMS; HCD



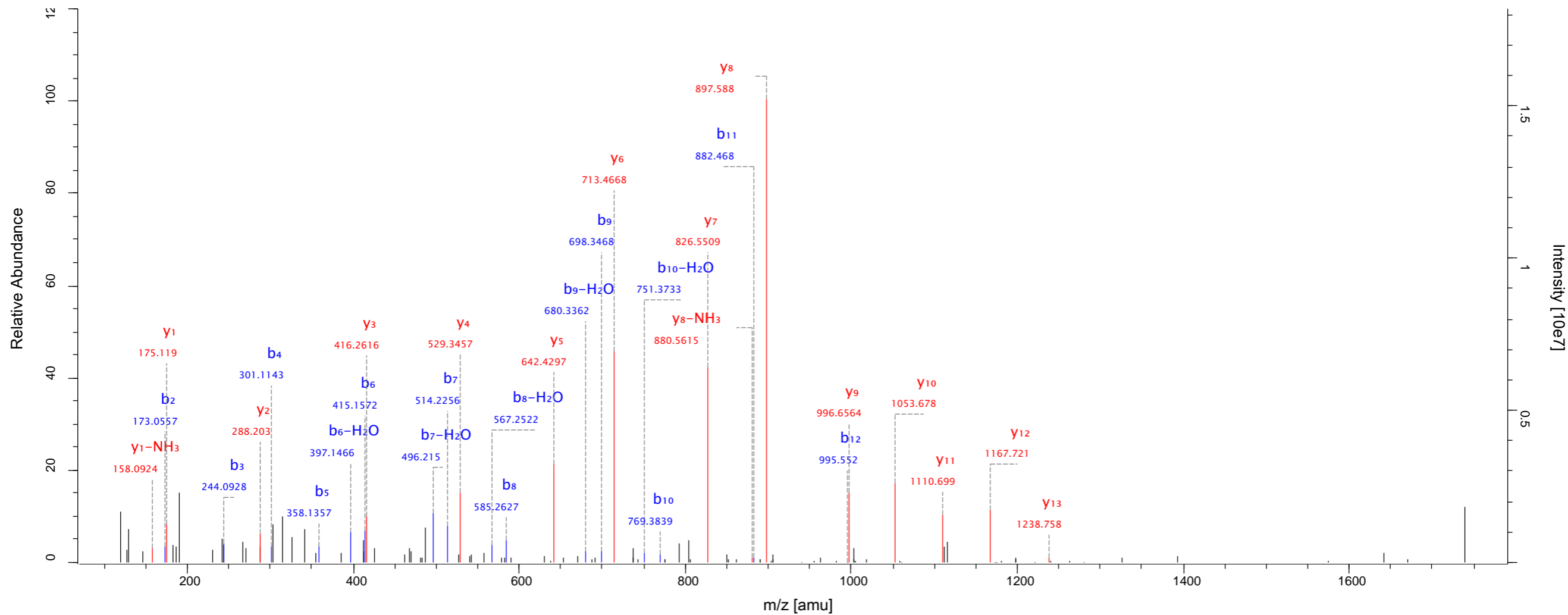
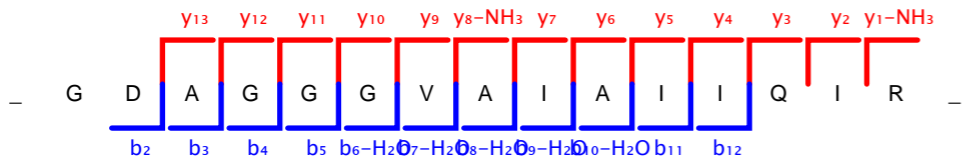
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 43214
Protein: ORFS343C.iORF1 - NC_006273.2[210869,211213]- <unknown description>
Peptide score: 62.92
Method: FTMS; HCD



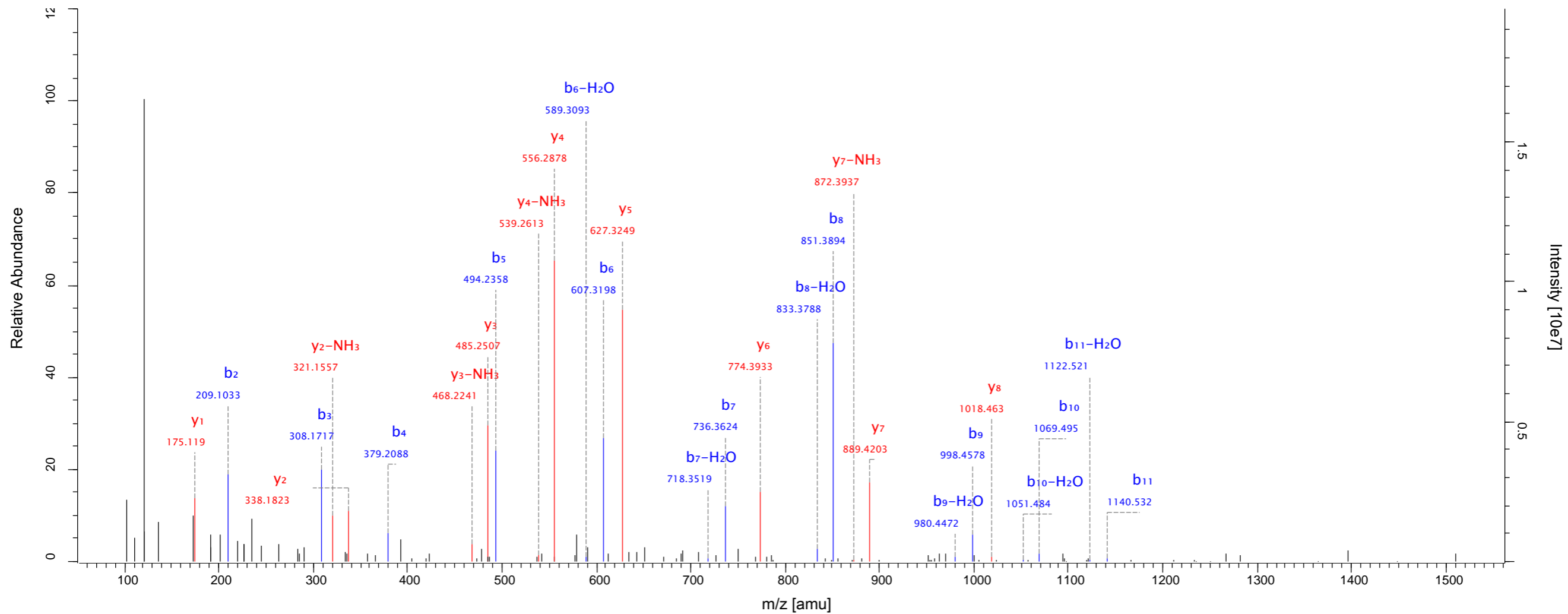
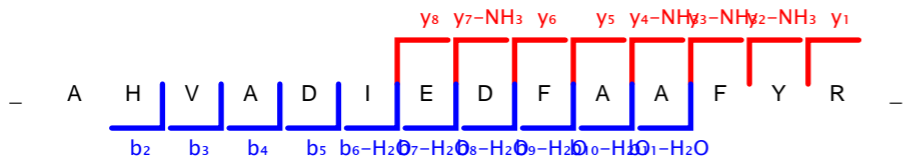
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpI1_26
Scan number: 49562
Protein: ORFL26W + NC_006273.2[12923,13093]+ <unknown description>
Peptide score: 59.23
Method: FTMS; HCD



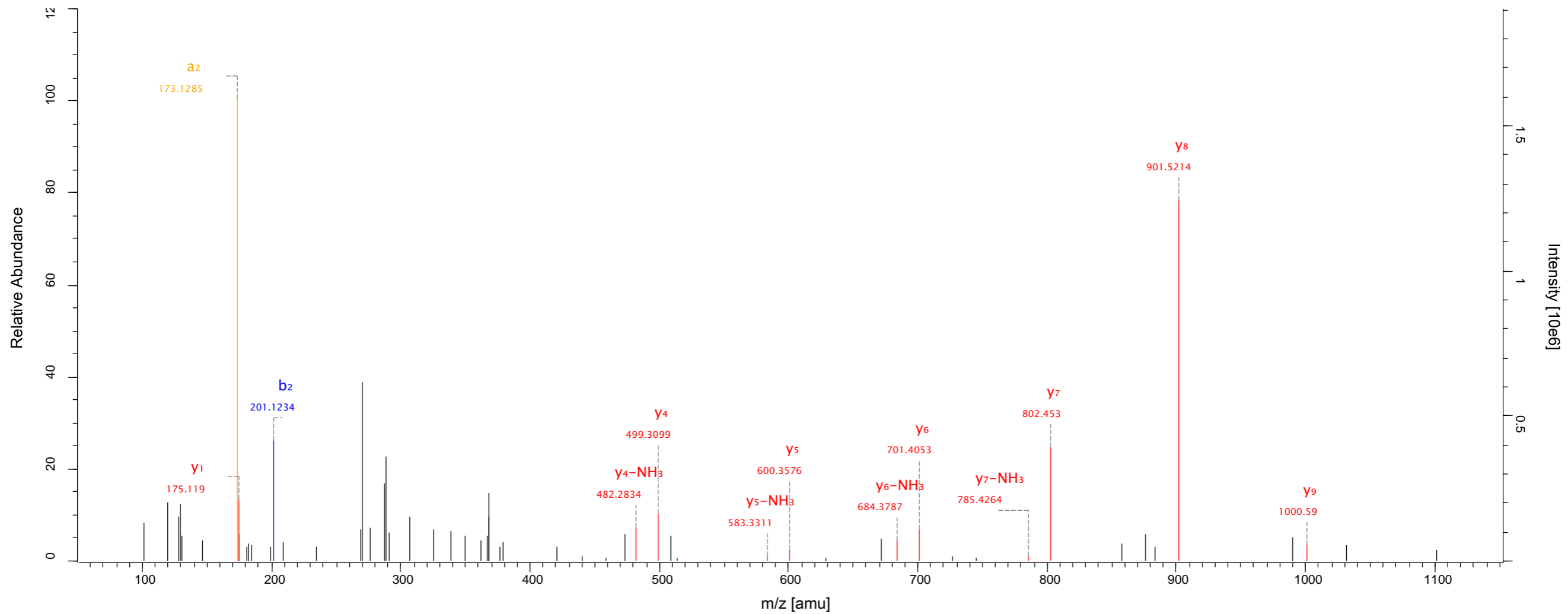
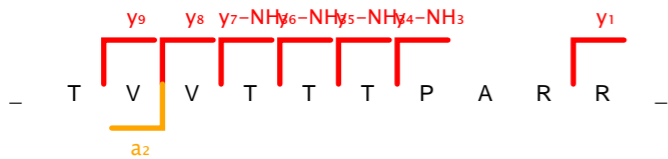
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
 Scan number: 50039
 Protein: ORFL223W + NC_006273.2[139458,139730]+ <unknown description>
 Peptide score: 178.84
 Method: FTMS; HCD



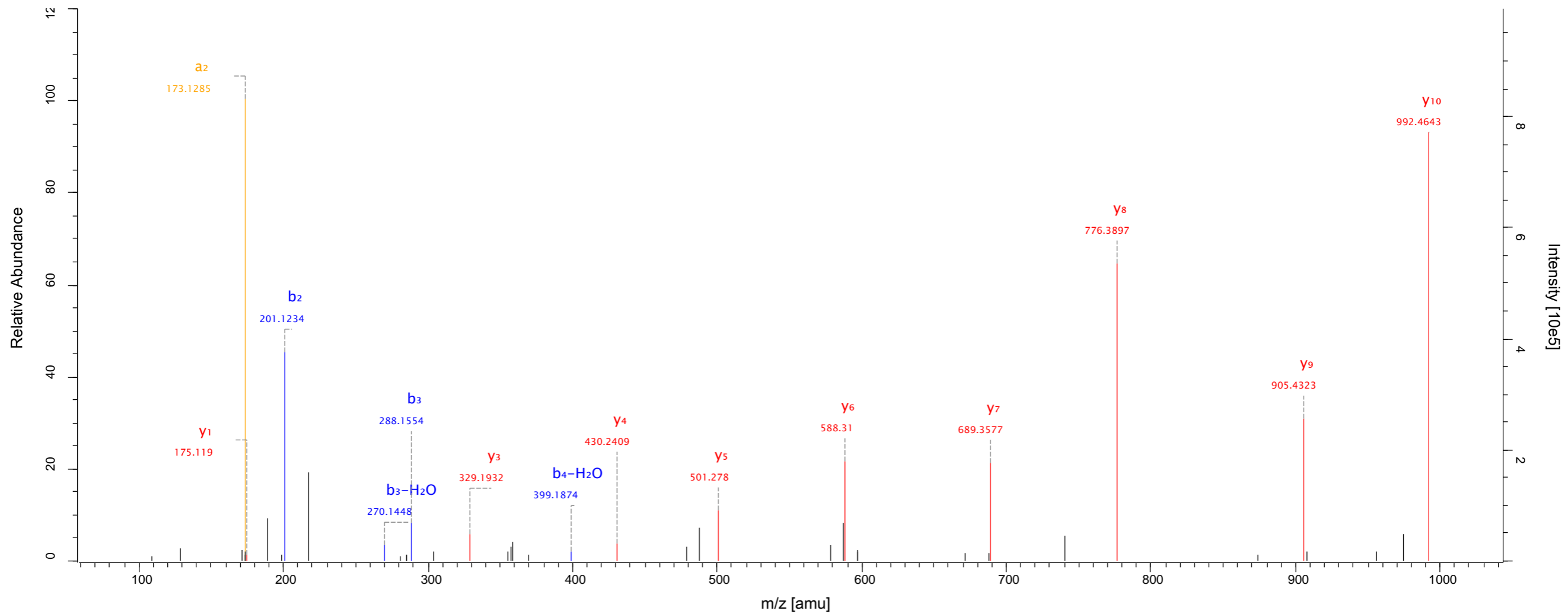
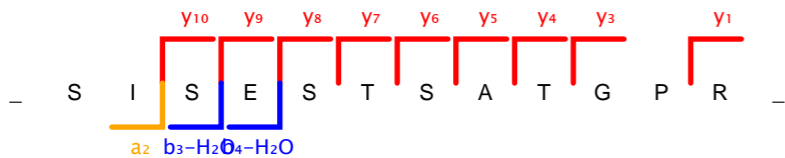
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
 Scan number: 50423
 Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>
 Peptide score: 150.68
 Method: FTMS; HCD



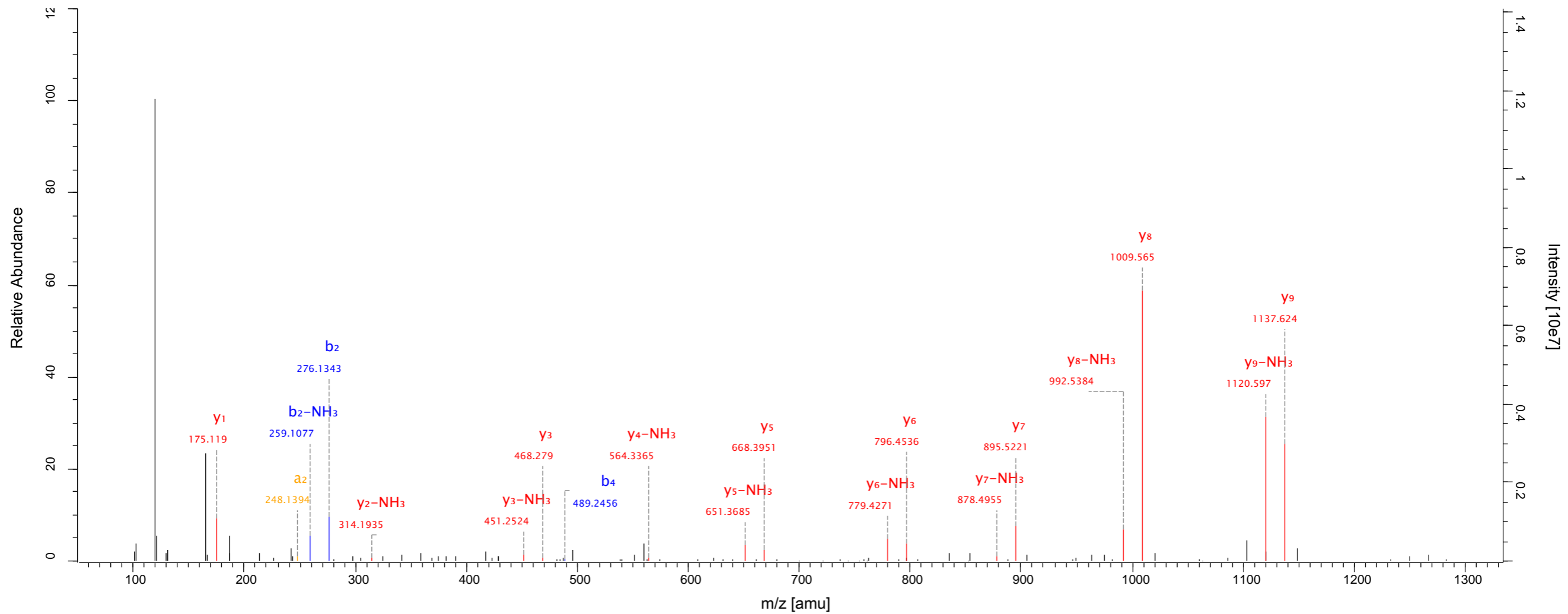
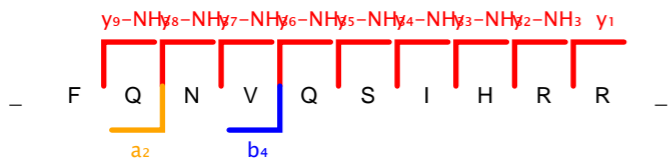
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 6247
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 110.87
Method: FTMS; HCD



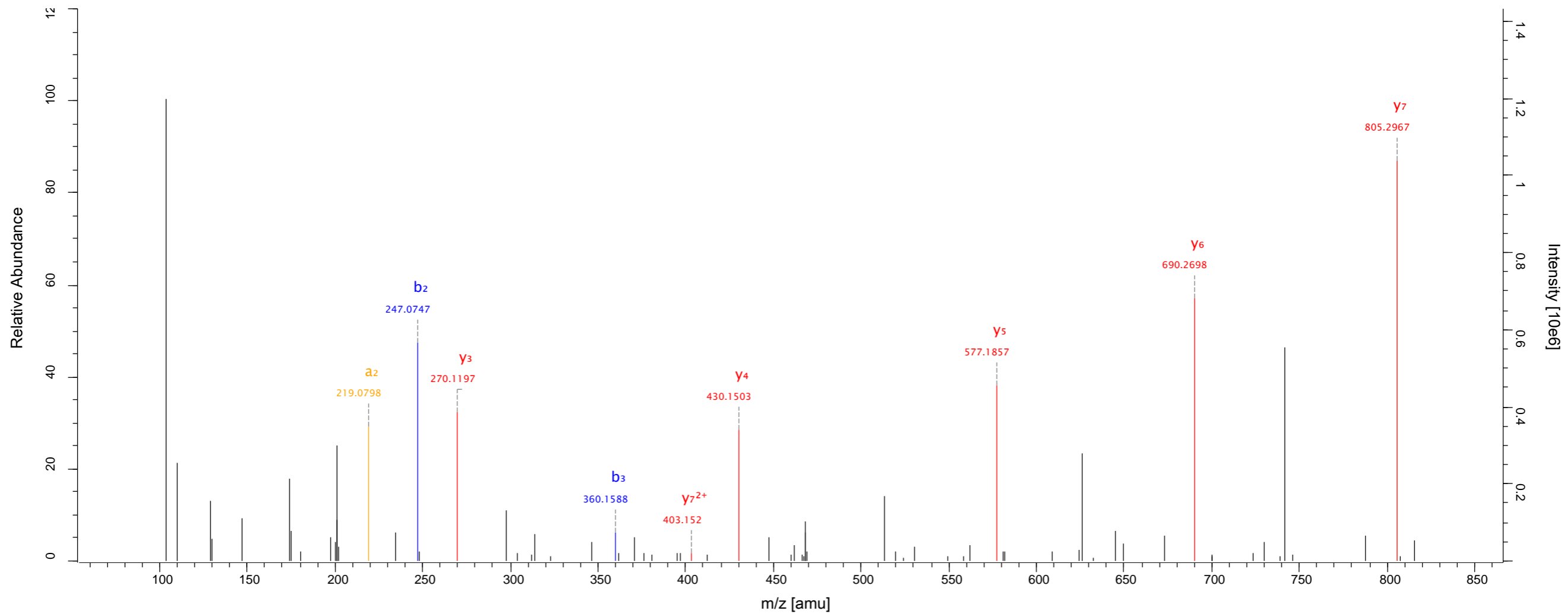
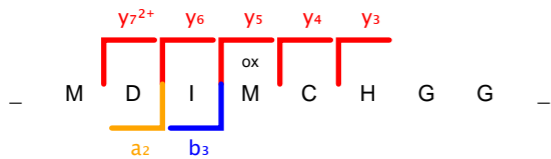
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 7508
Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
Peptide score: 101.95
Method: FTMS; HCD



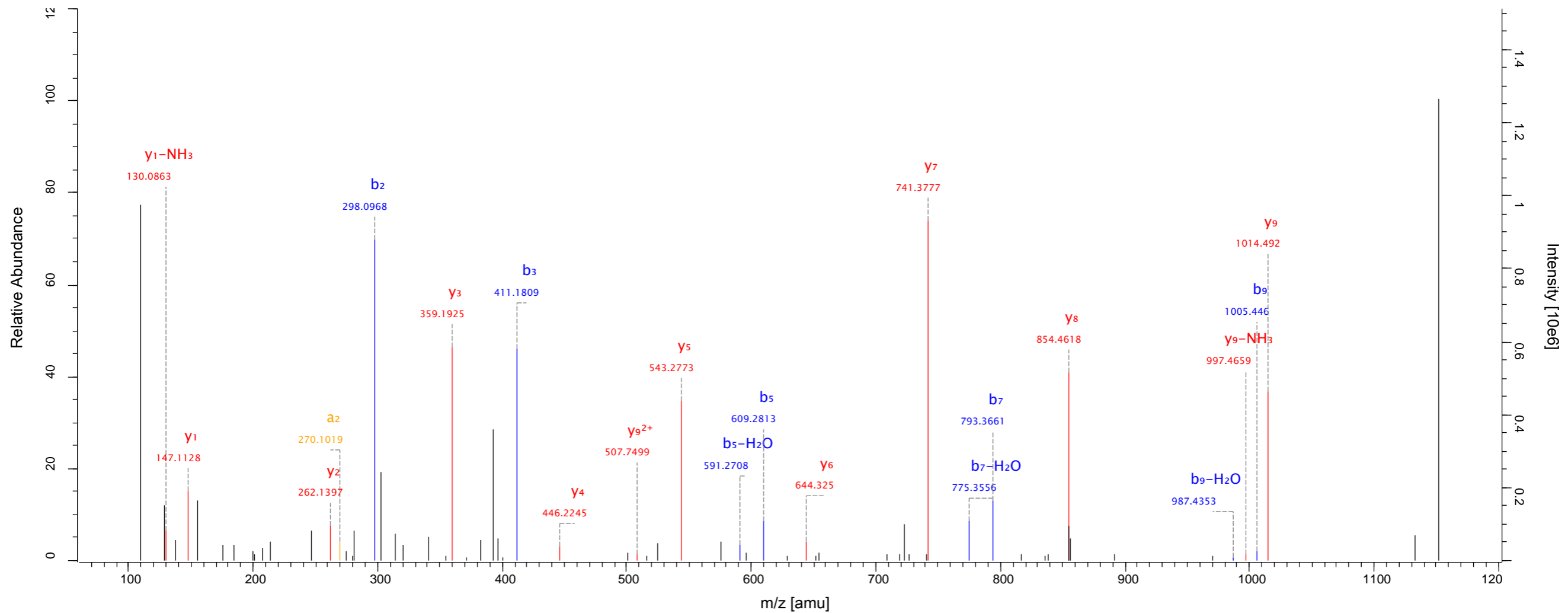
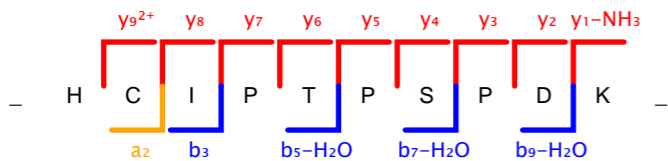
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 8446
Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
Peptide score: 116.52
Method: FTMS; HCD



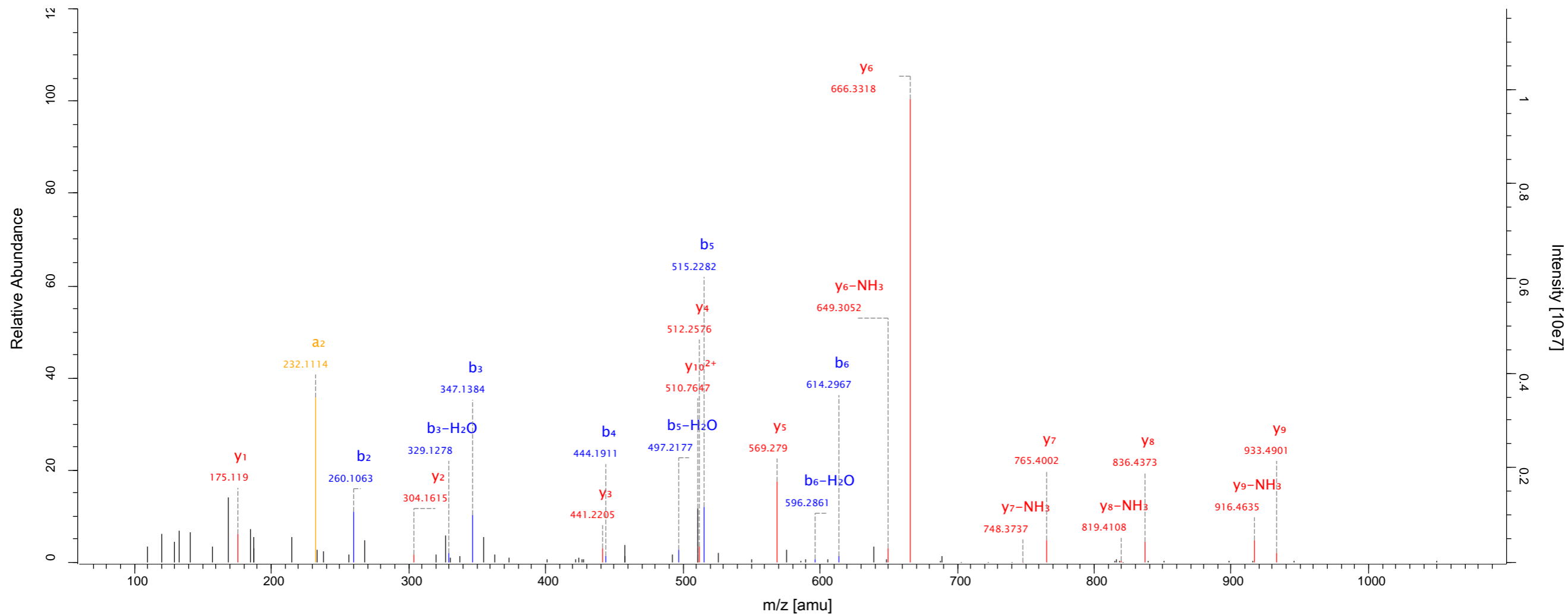
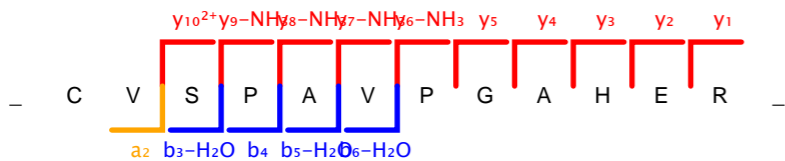
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_26
Scan number: 8766
Protein: ORFL88C - NC_006273.2[37894,38133]- <unknown description>; rORF58 - NC_006273.2[37894,37920]- <unknown description>
Peptide score: 85.85
Method: FTMS; HCD



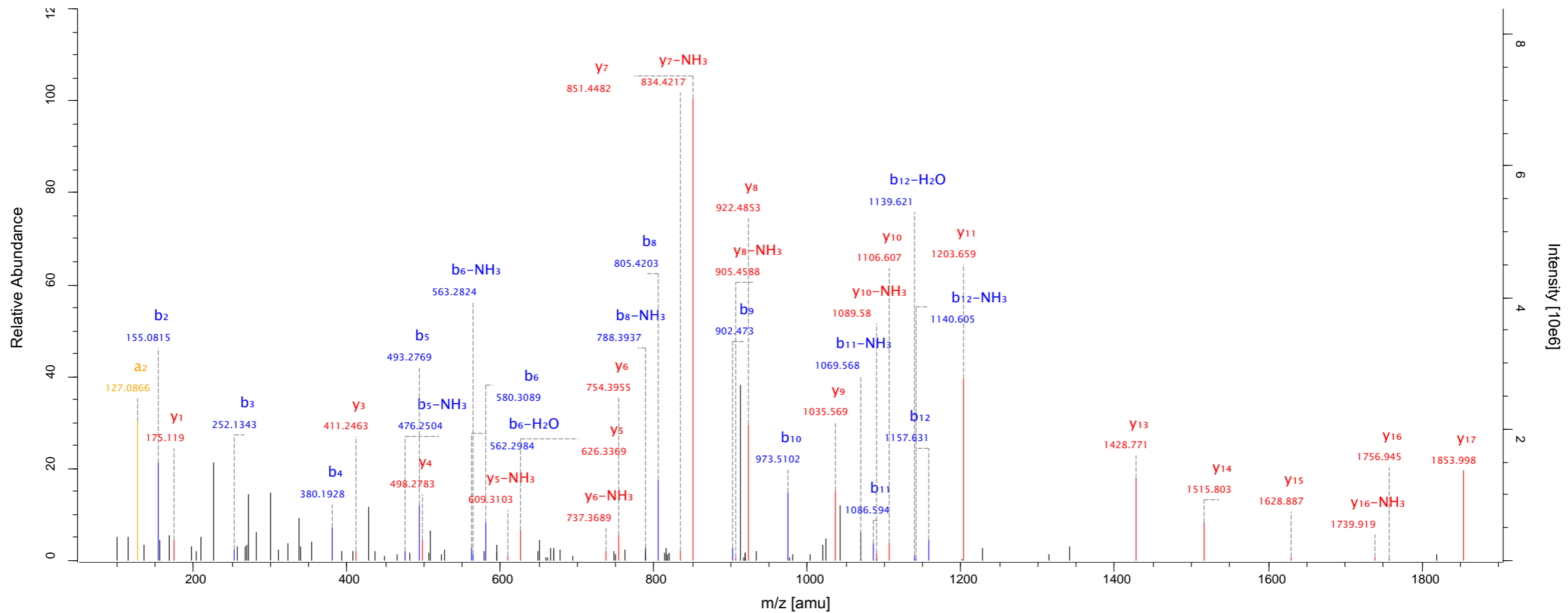
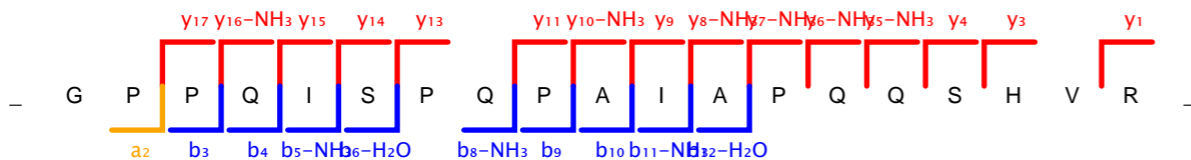
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_26
Scan number: 8954
Protein: ORFL26W + NC_006273.2[12923,13093]+ <unknown description>
Peptide score: 158.76
Method: FTMS; HCD



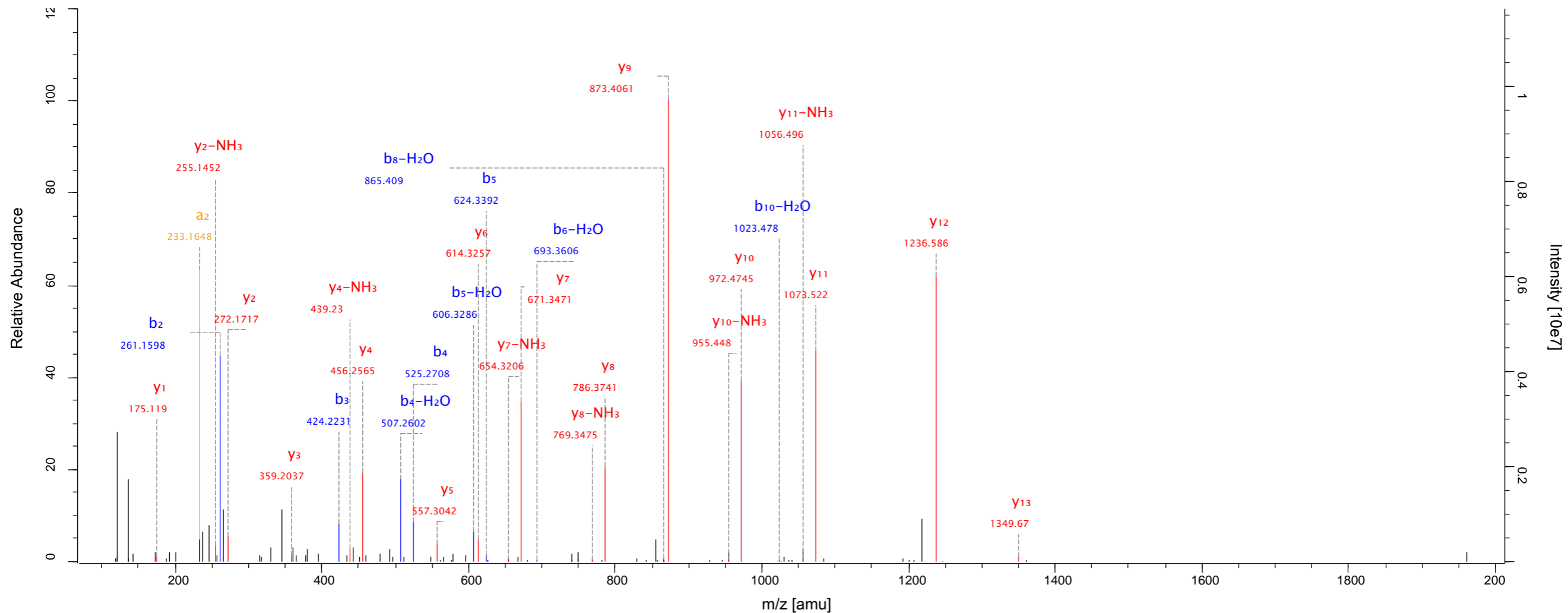
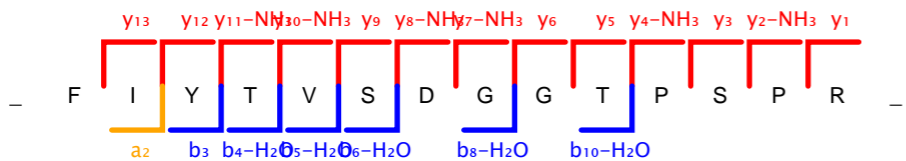
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
 Scan number: 10302
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 147.74
 Method: FTMS; HCD



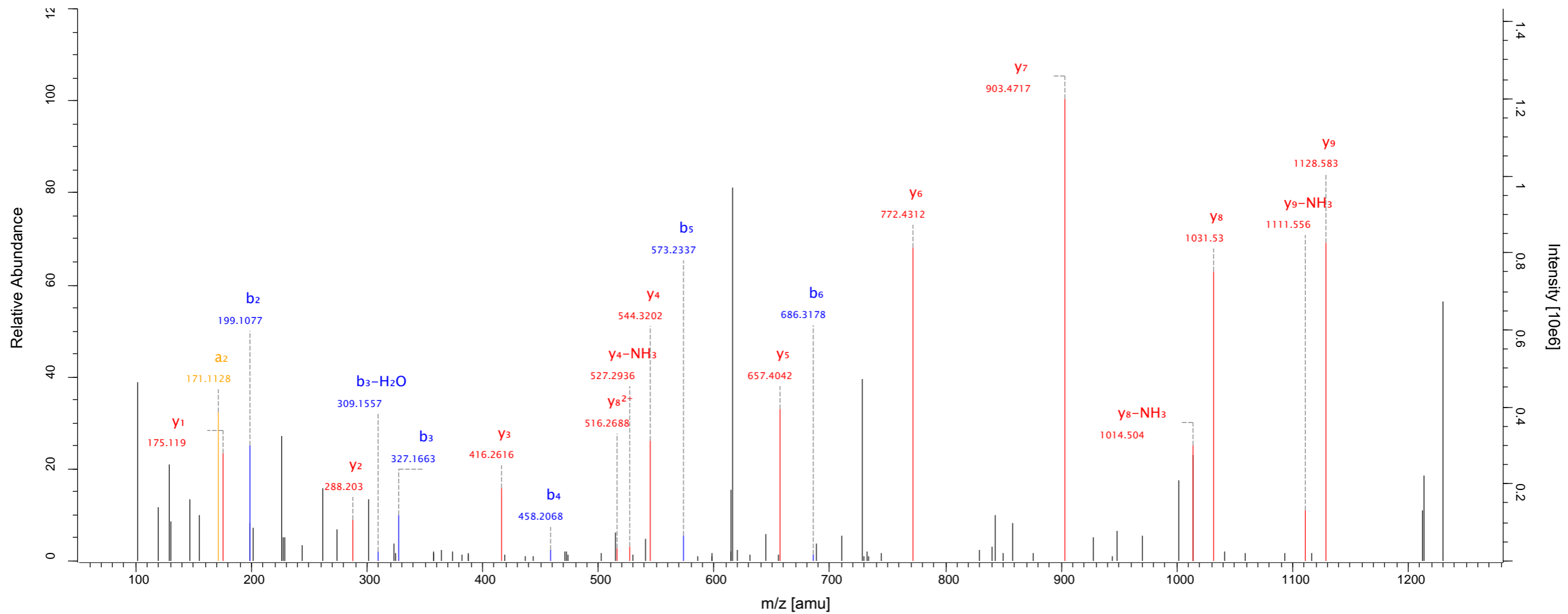
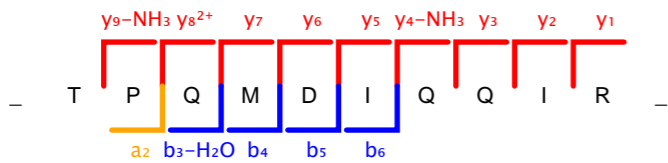
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27_111216192808
 Scan number: 22630
 Protein: ORFL253W.iORF1 + NC_006273.2[161917,163748]+ <unknown description>; ORFL251W + NC_006273.2[161872,163748]+ <unknown description>
 Peptide score: 183.36
 Method: FTMS; HCD



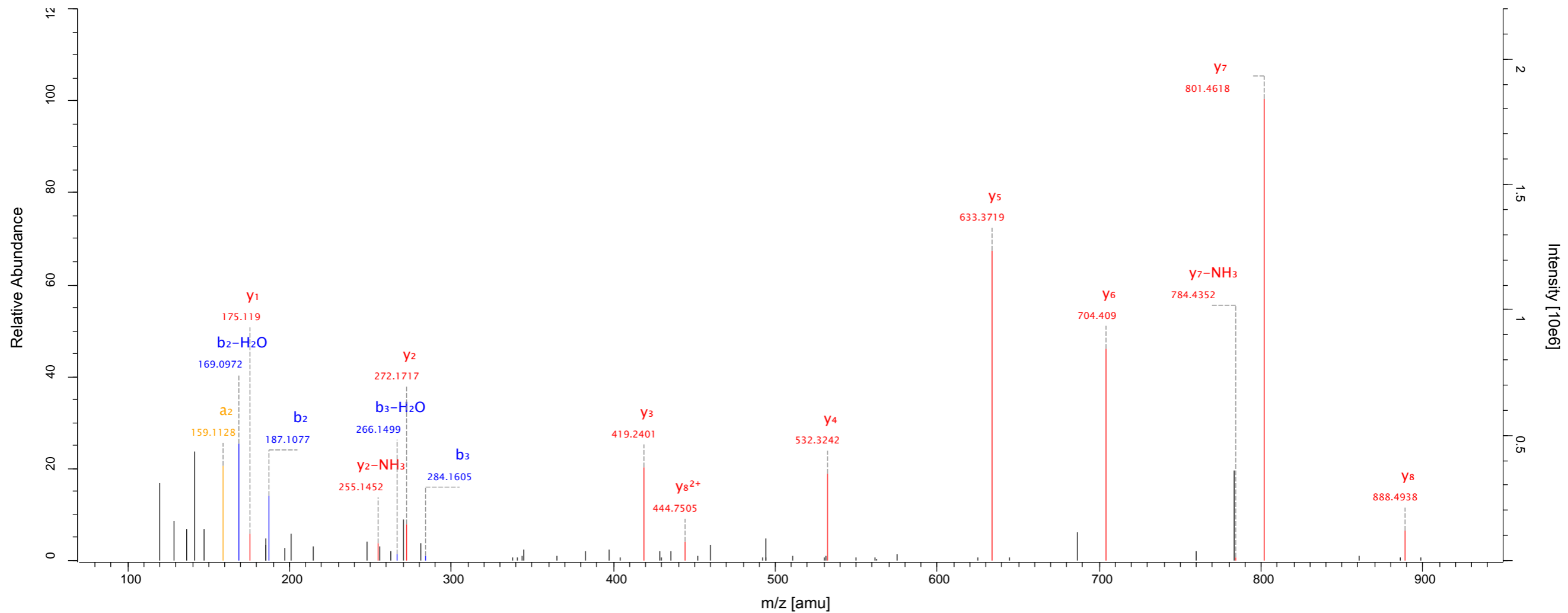
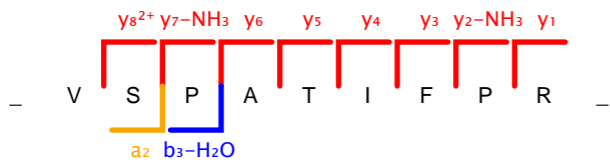
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
 Scan number: 25420
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 135.1
 Method: FTMS; HCD



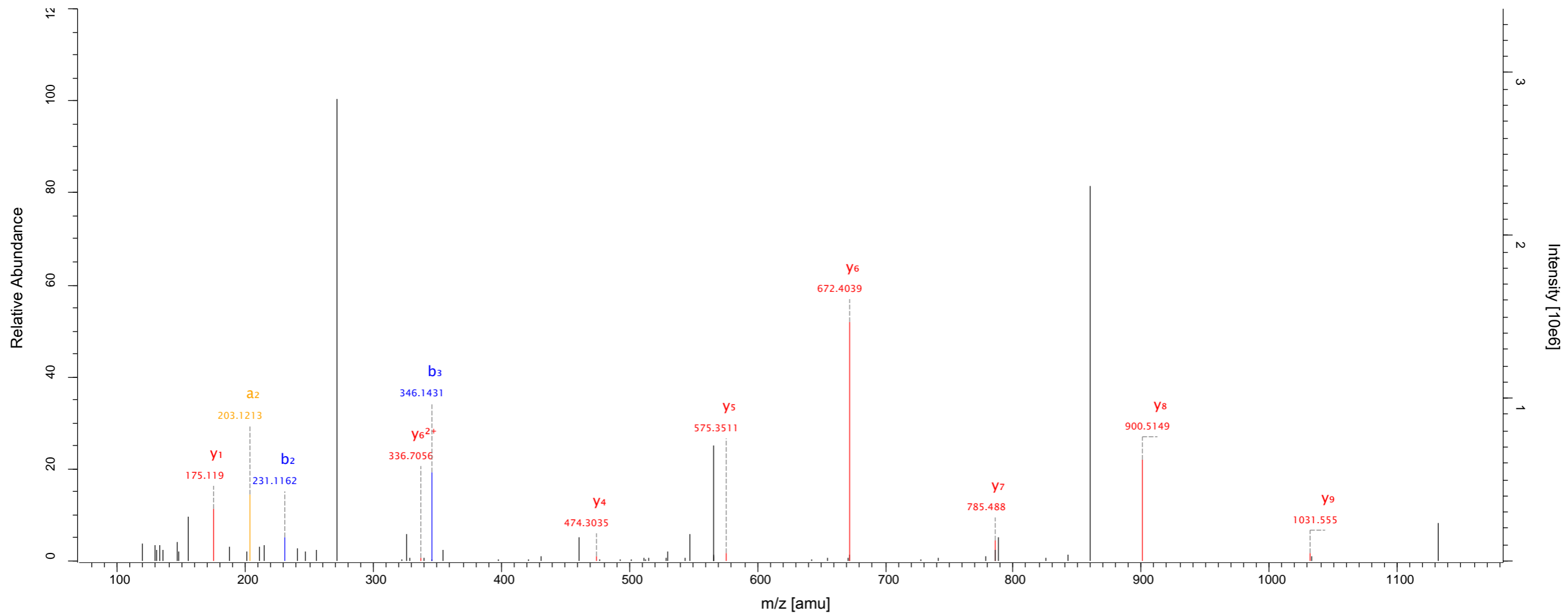
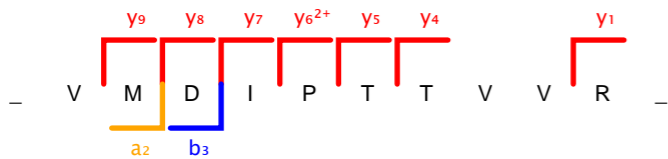
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_27_111216192808
Scan number: 26124
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 119.53
Method: FTMS; HCD



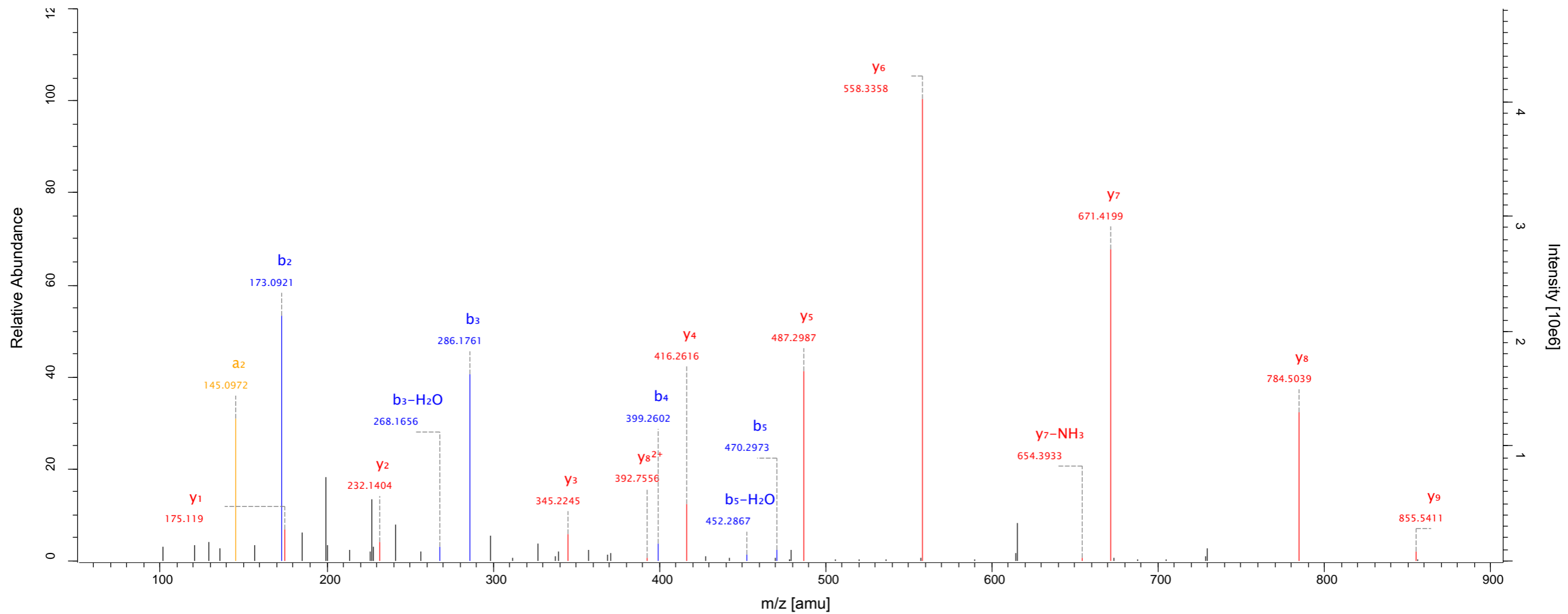
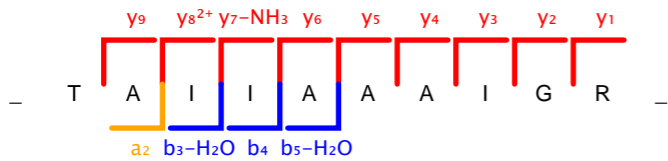
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
Scan number: 30921
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 99.44
Method: FTMS; HCD



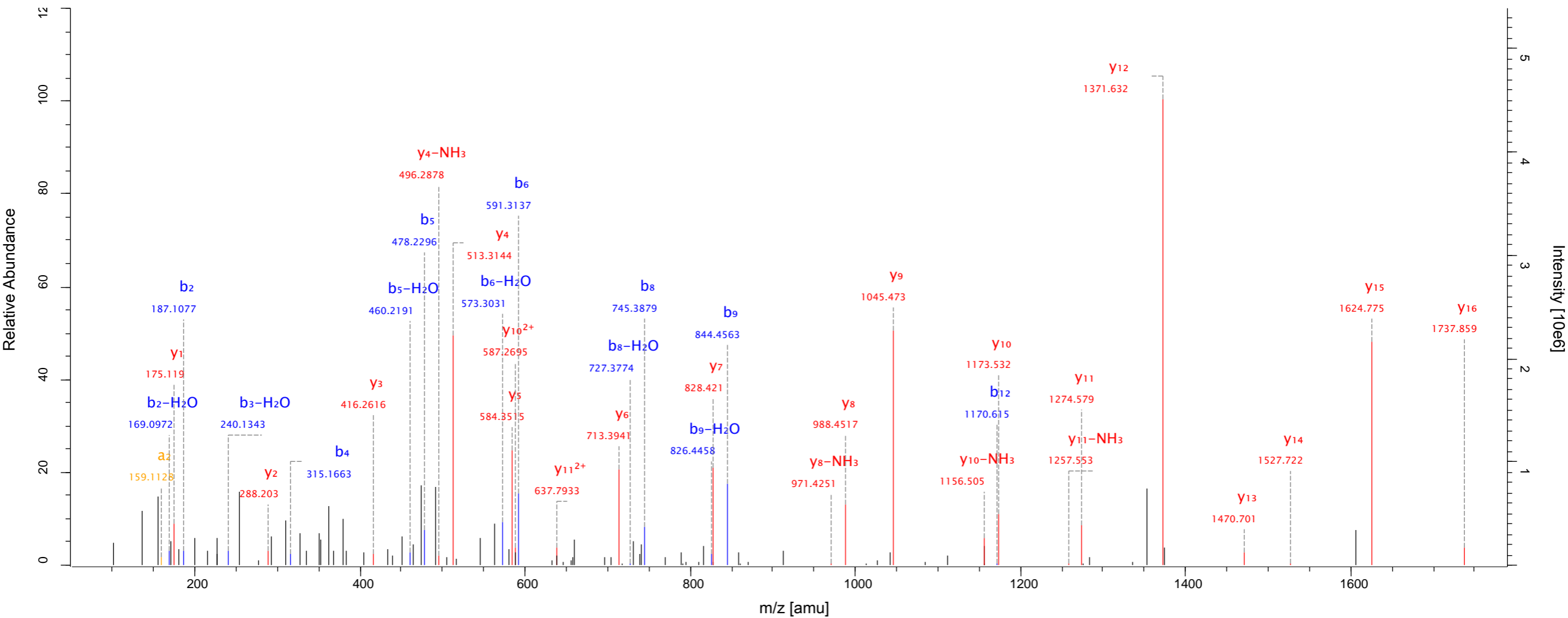
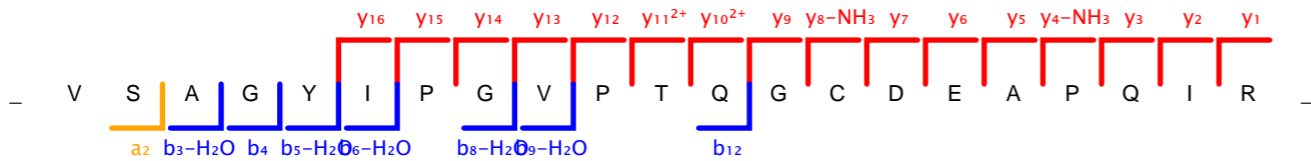
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
Scan number: 31053
Protein: ORFL251W + NC_006273.2[161872,163748]+ <unknown description>
Peptide score: 62.82
Method: FTMS; HCD



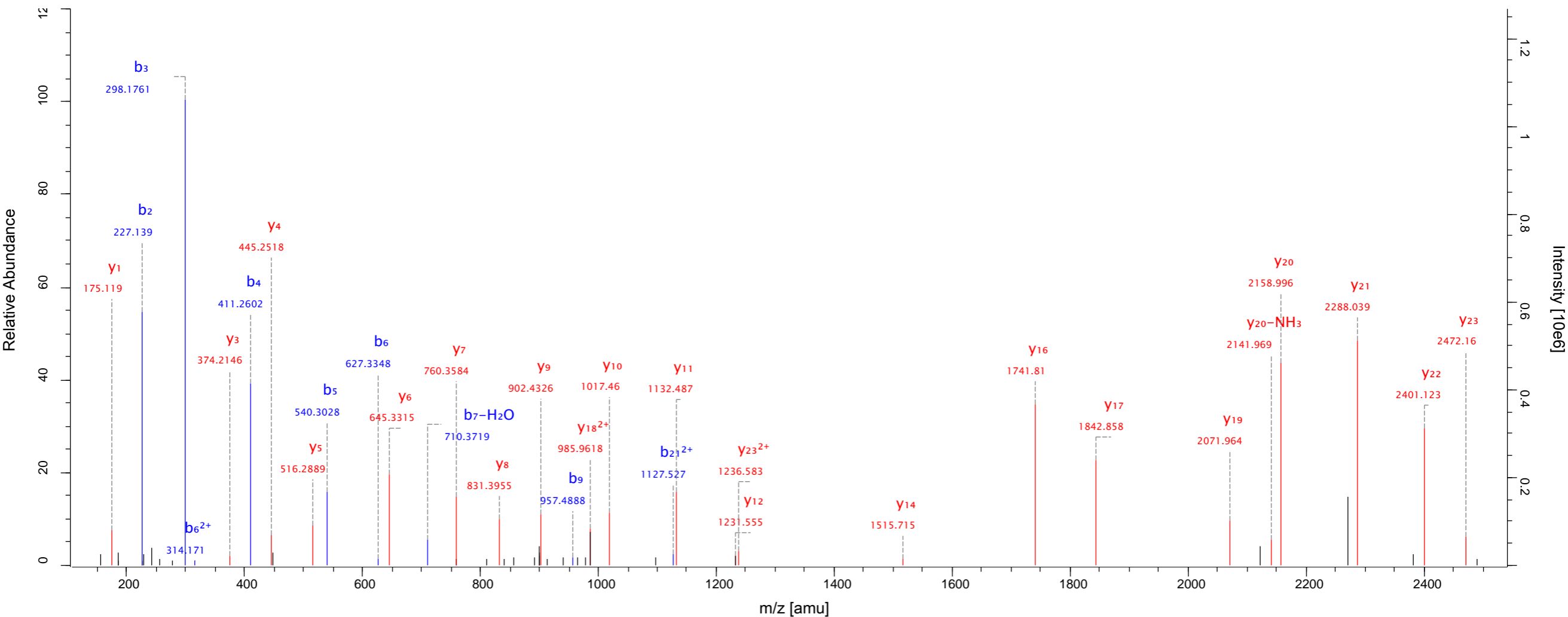
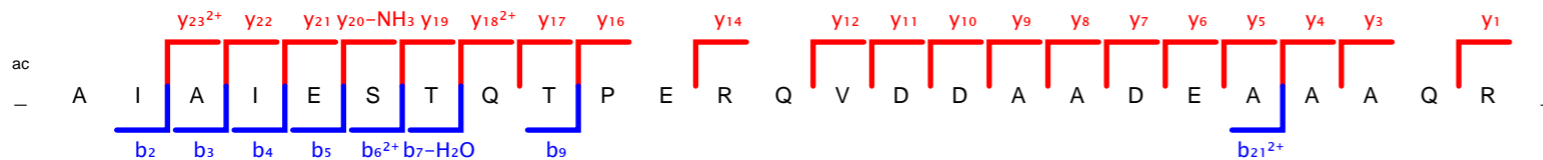
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
Scan number: 33768
Protein: ORFL118C - NC_006273.2[61896,62396]- <unknown description>
Peptide score: 107.21
Method: FTMS; HCD



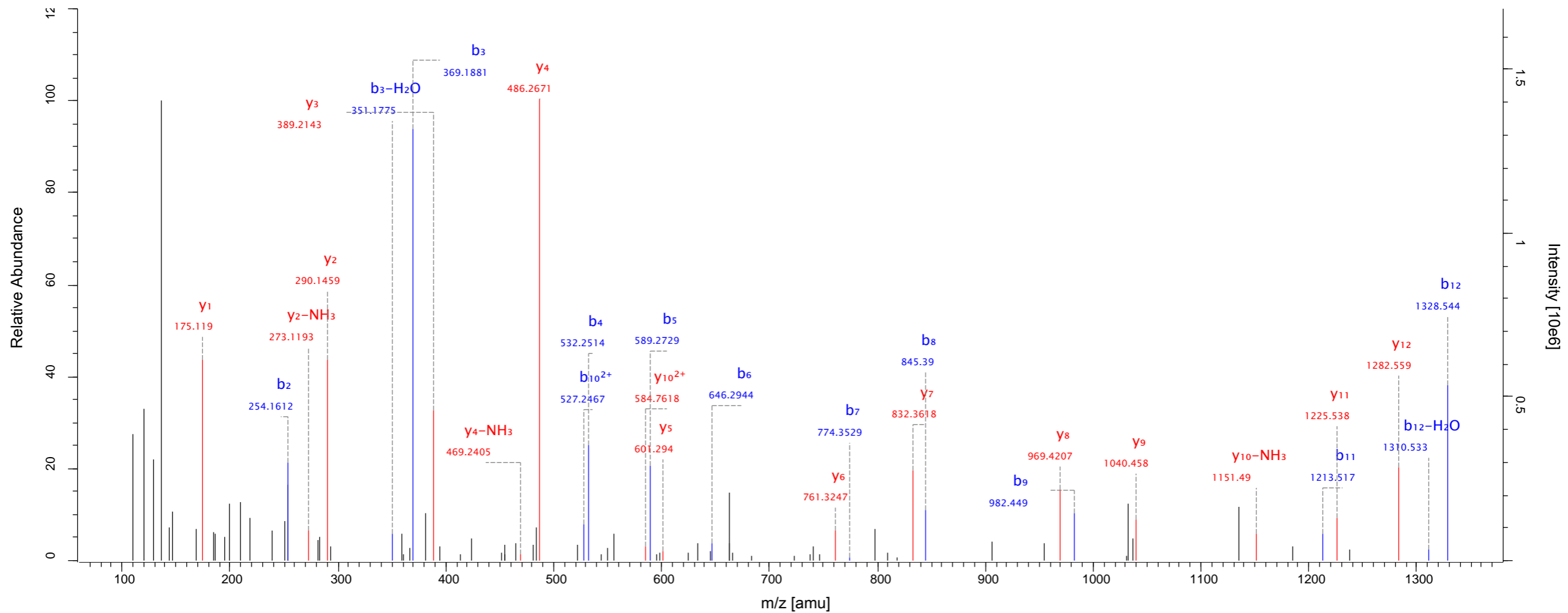
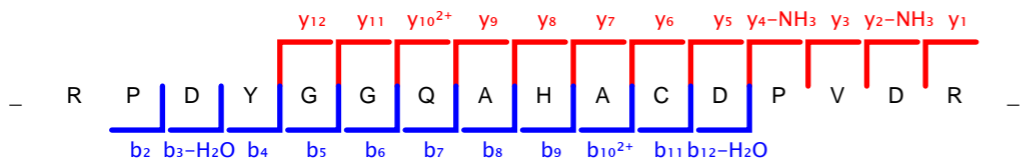
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
 Scan number: 43262
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 132.13
 Method: FTMS; HCD



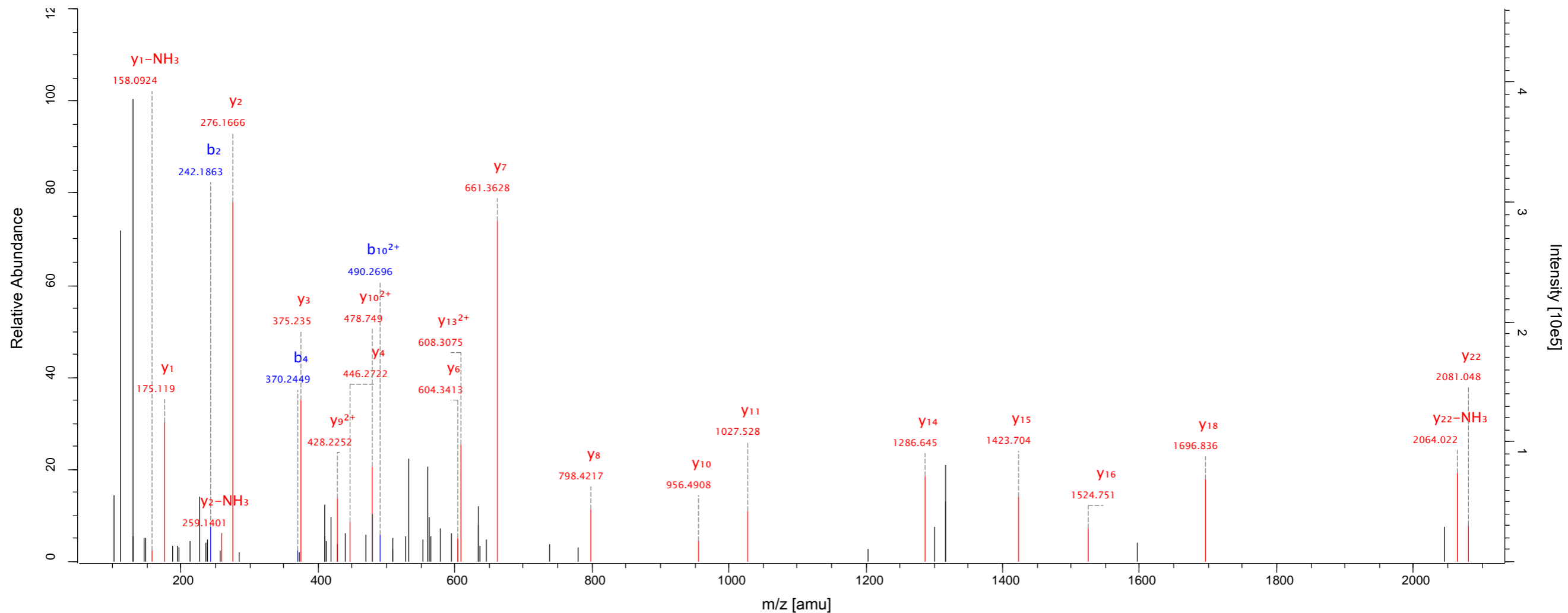
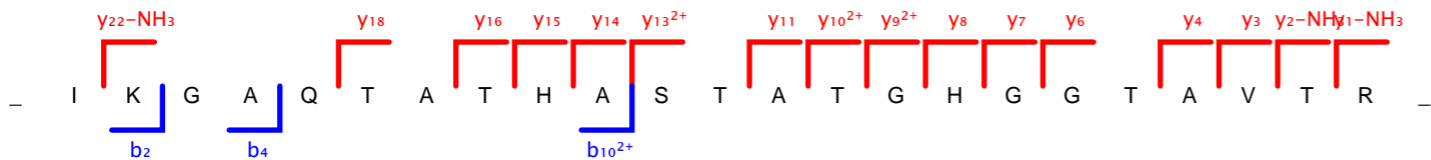
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
 Scan number: 43454
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 235.9
 Method: FTMS; HCD



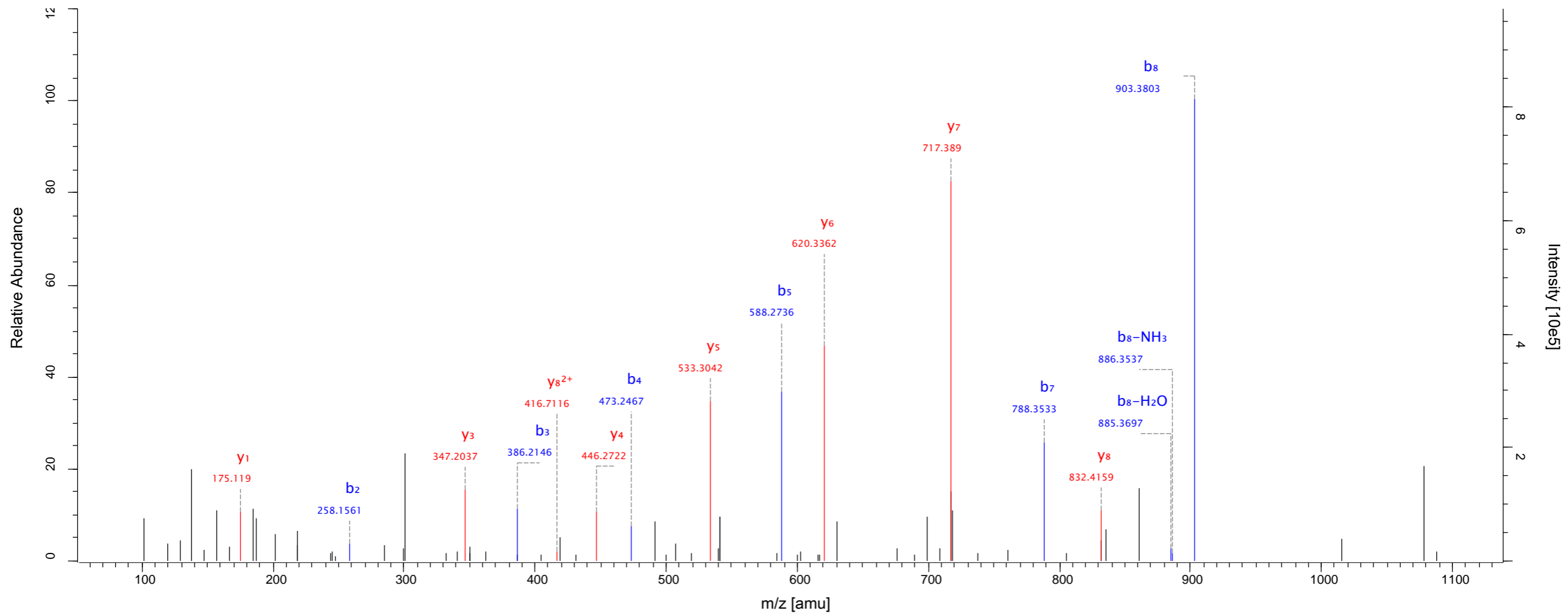
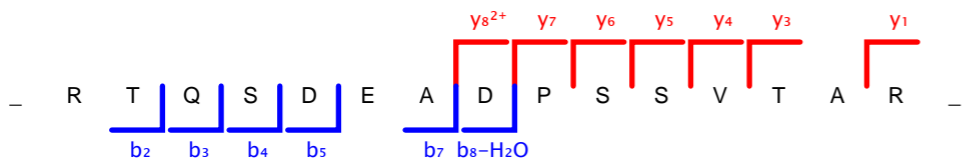
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
Scan number: 43848
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 137.19
Method: FTMS; HCD



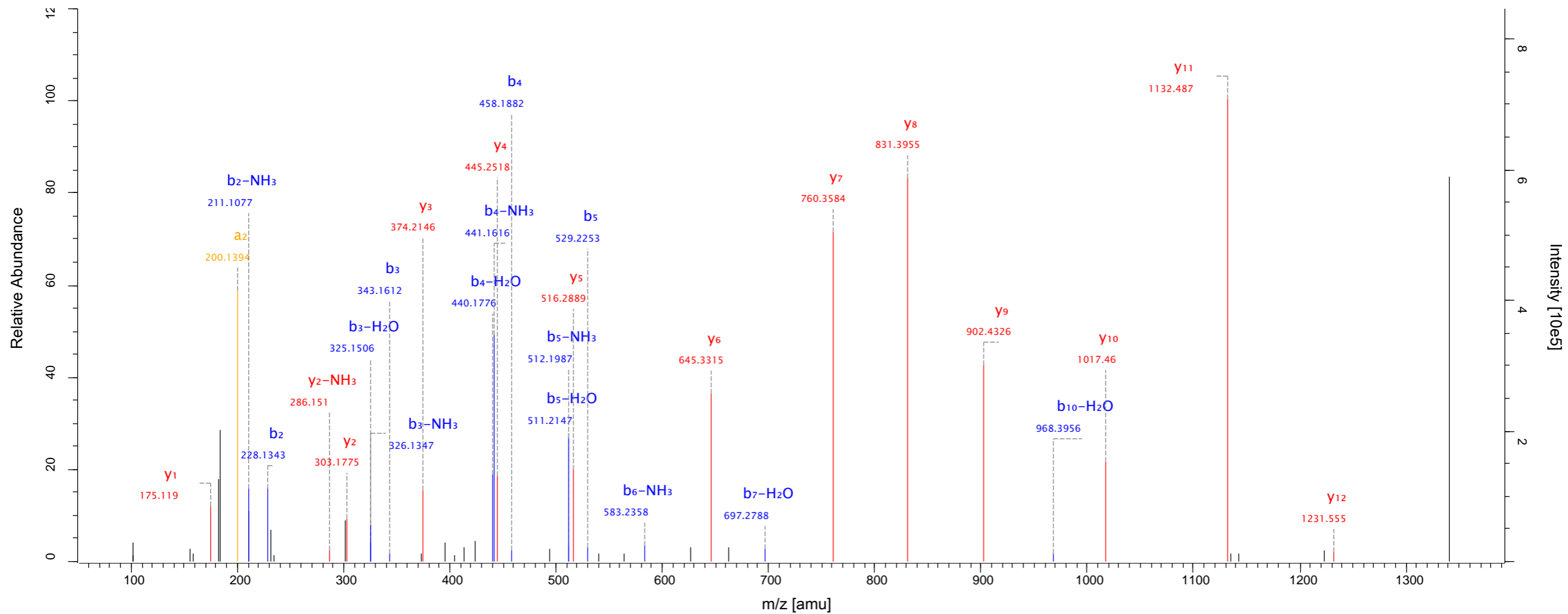
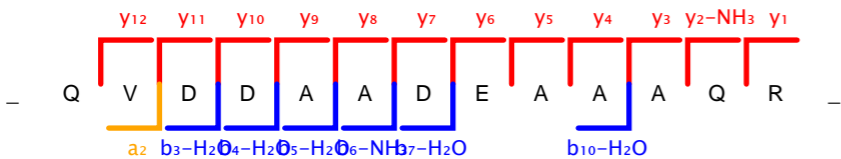
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
 Scan number: 6782
 Protein: ORFL114W + NC_006273.2[60151,60708]+ <unknown description>
 Peptide score: 101.64
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27_111216192808
Scan number: 7426
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 83.69
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27_111216192808
 Scan number: 8745
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 149.23
 Method: FTMS; HCD



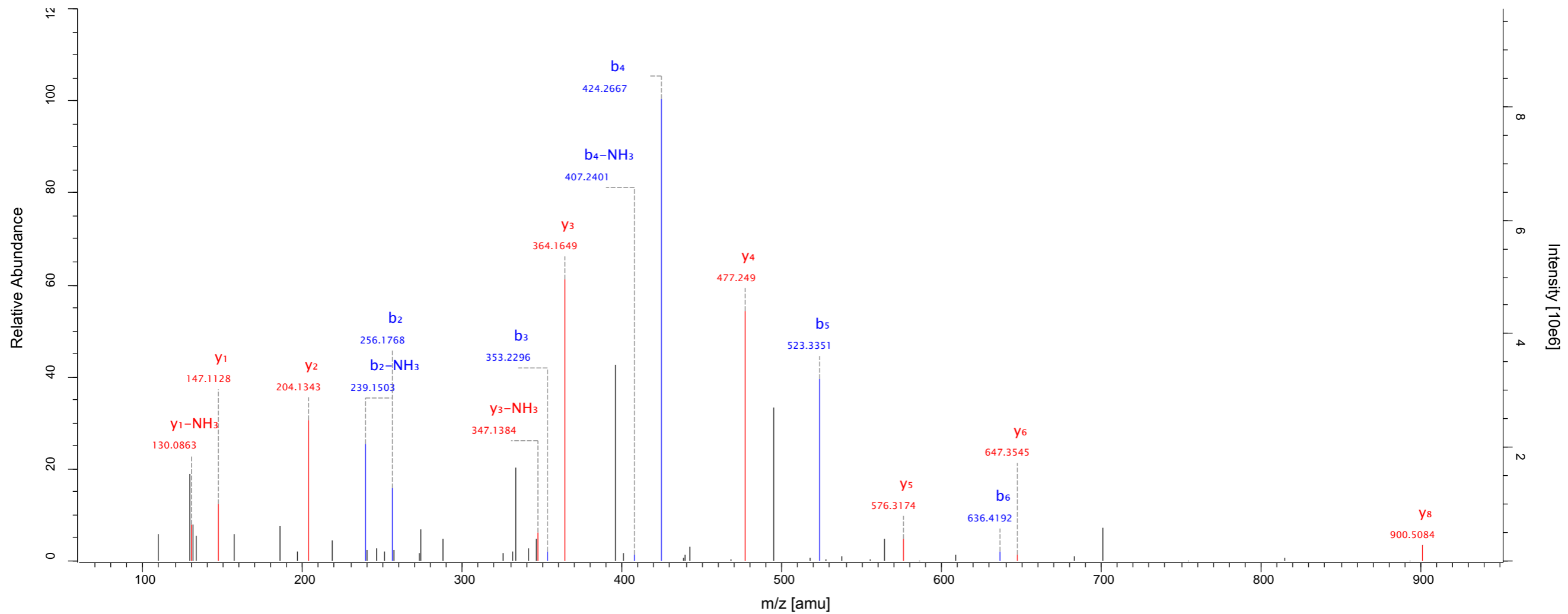
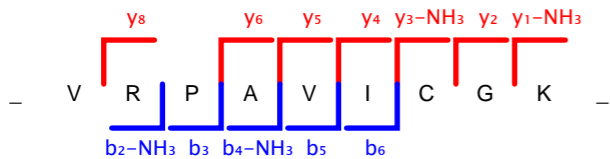
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27

Scan number: 10080

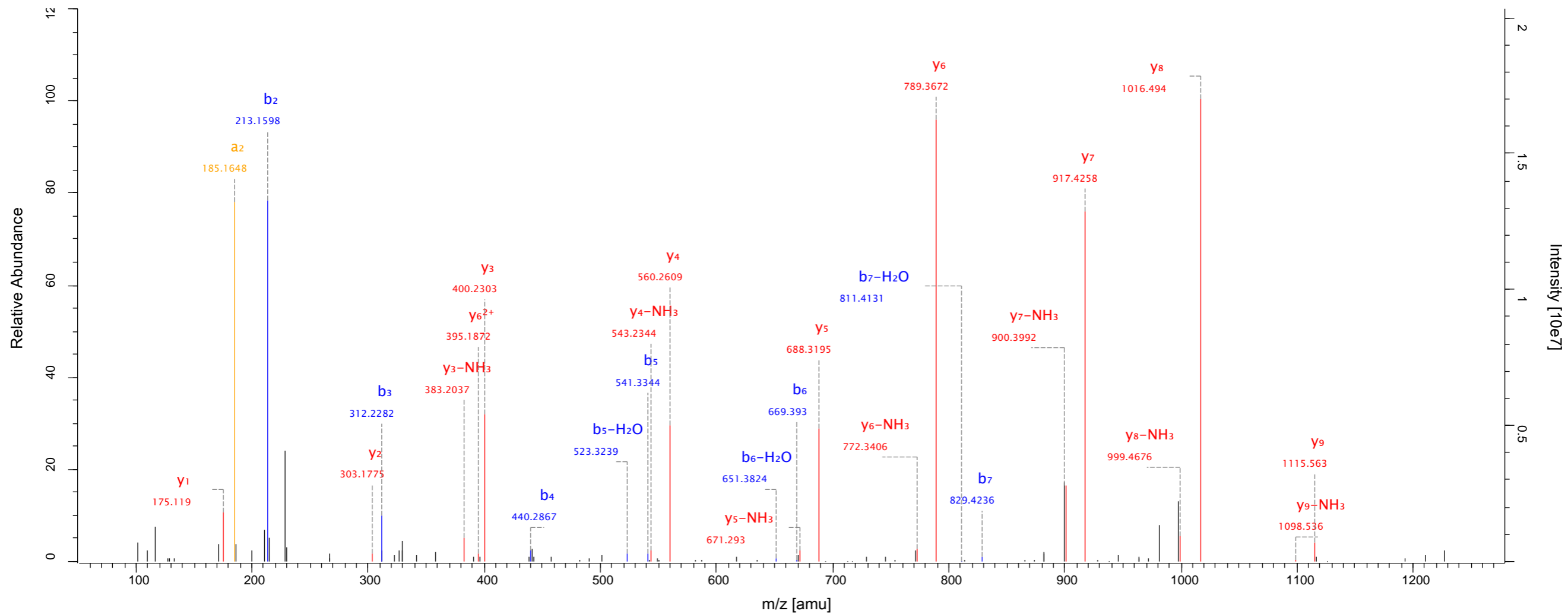
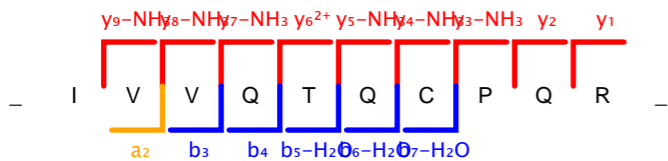
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 118.26

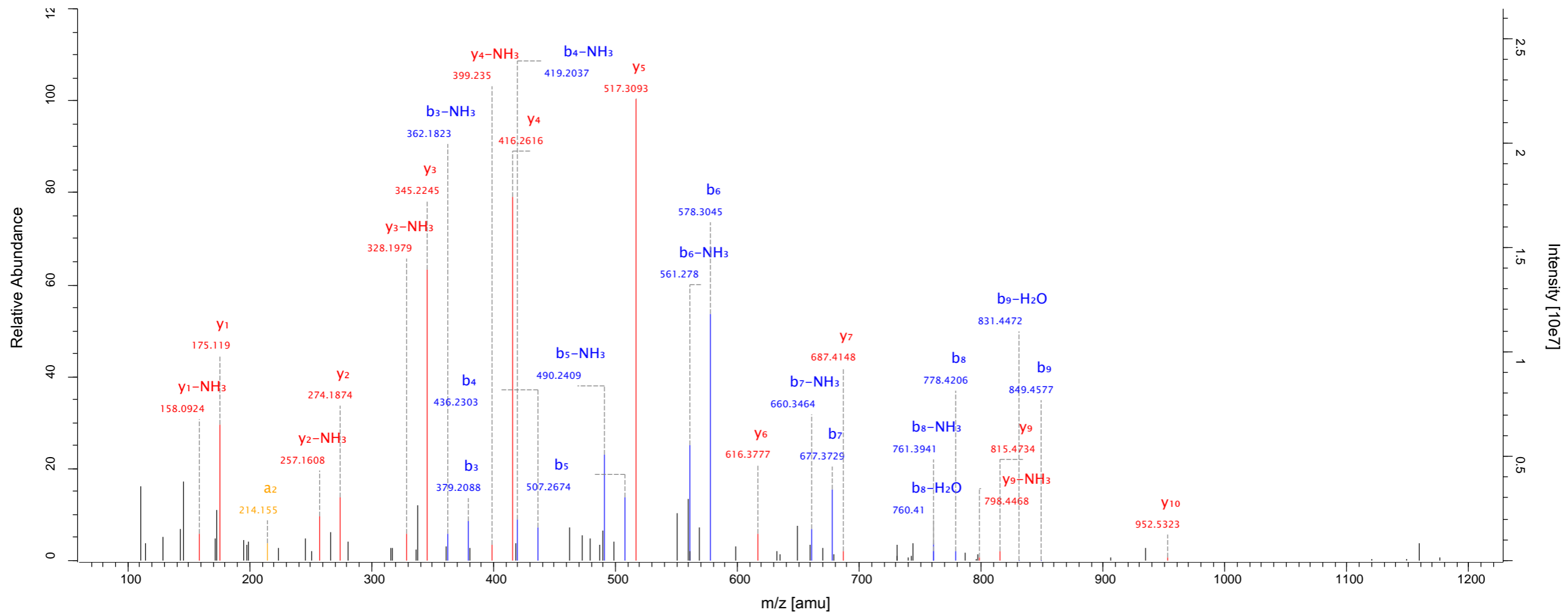
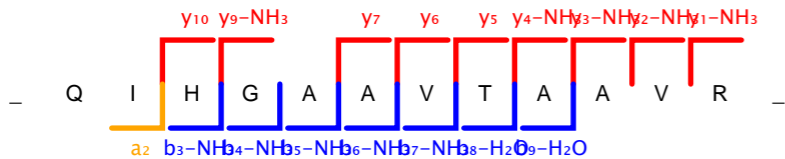
Method: FTMS; HCD



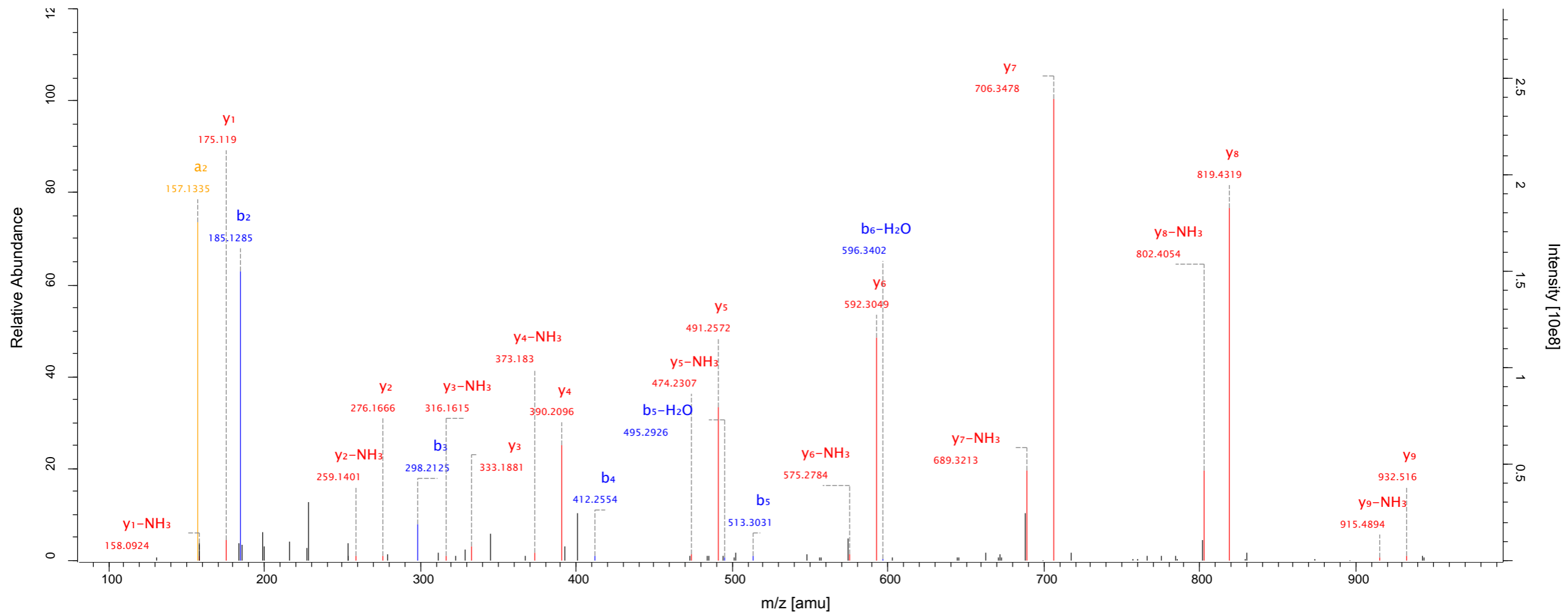
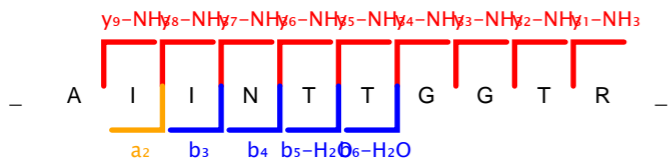
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 11445
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 209.6
 Method: FTMS; HCD



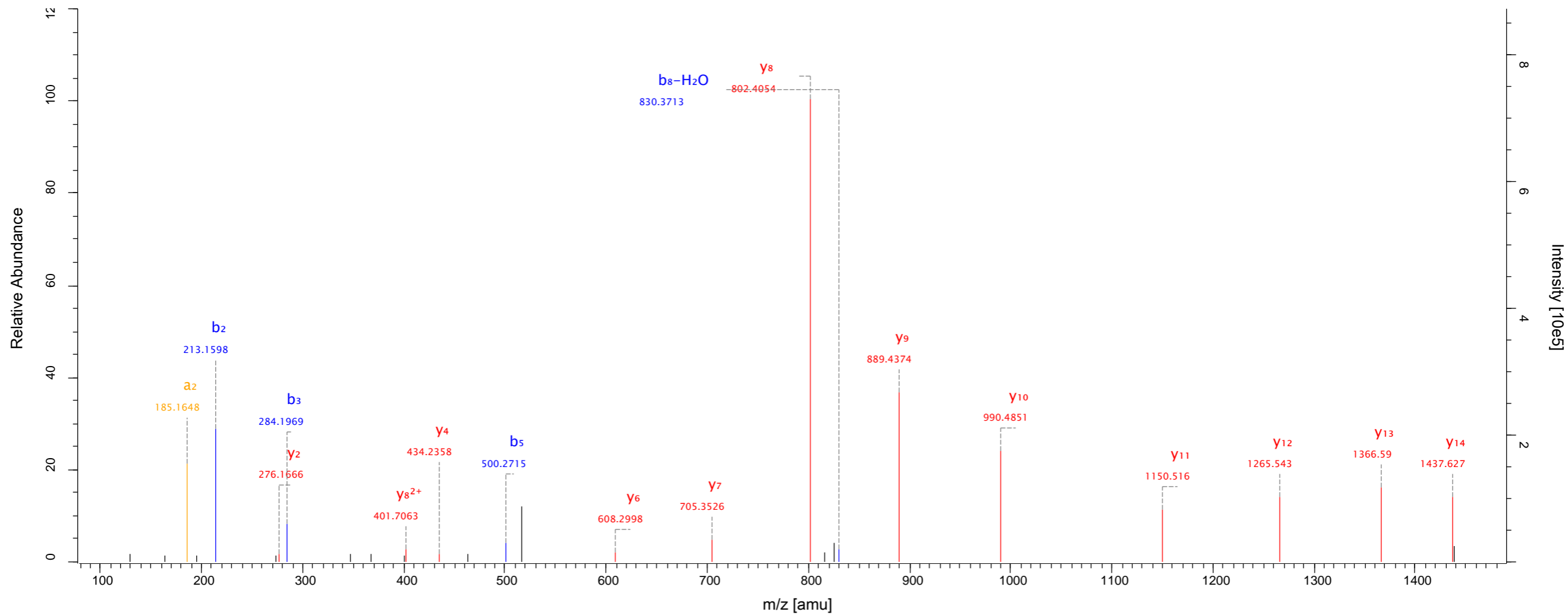
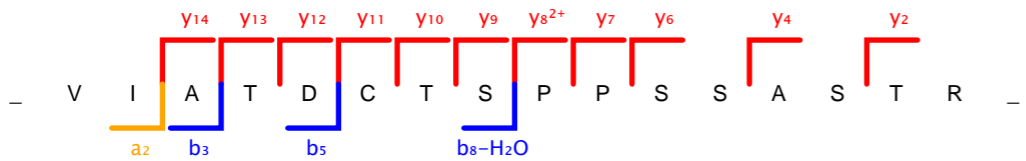
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 11506
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 196.44
 Method: FTMS; HCD



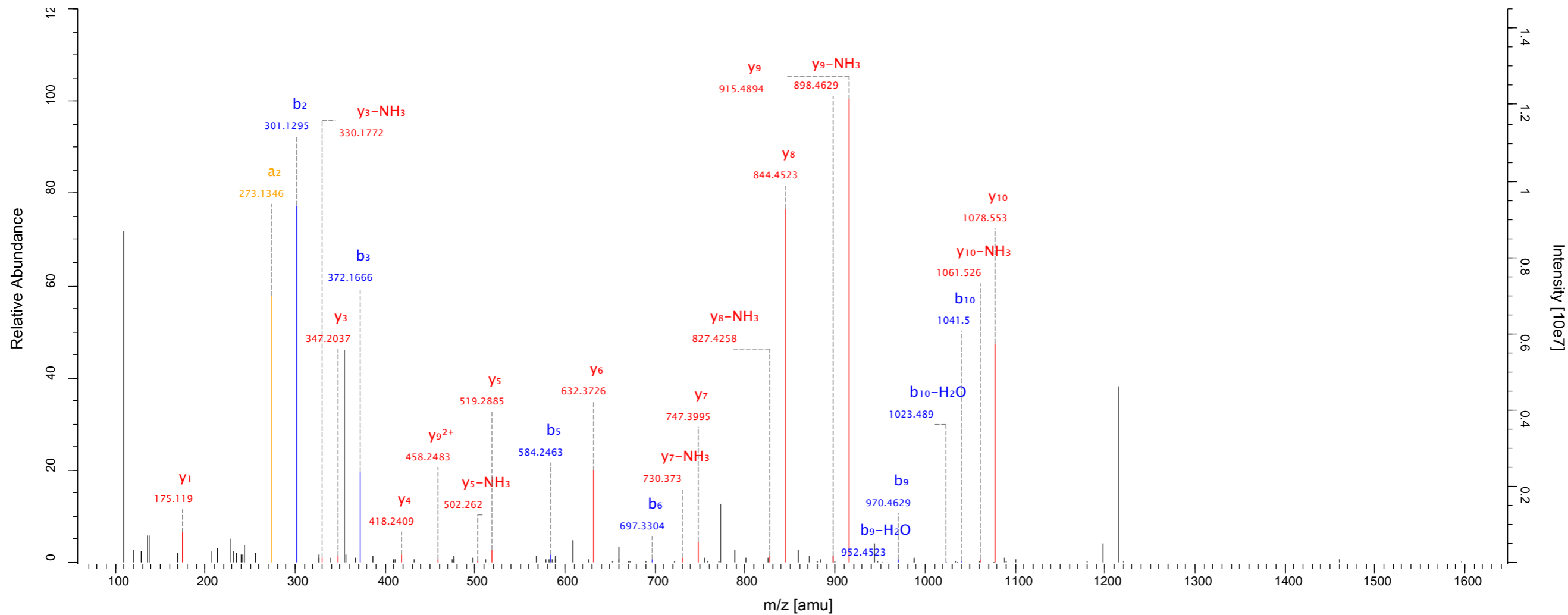
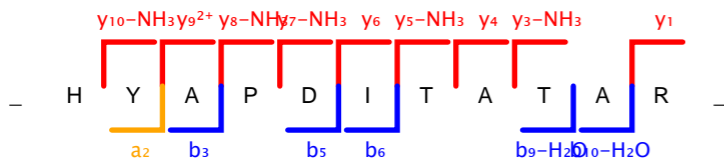
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 11789
 Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
 Peptide score: 146.27
 Method: FTMS; HCD



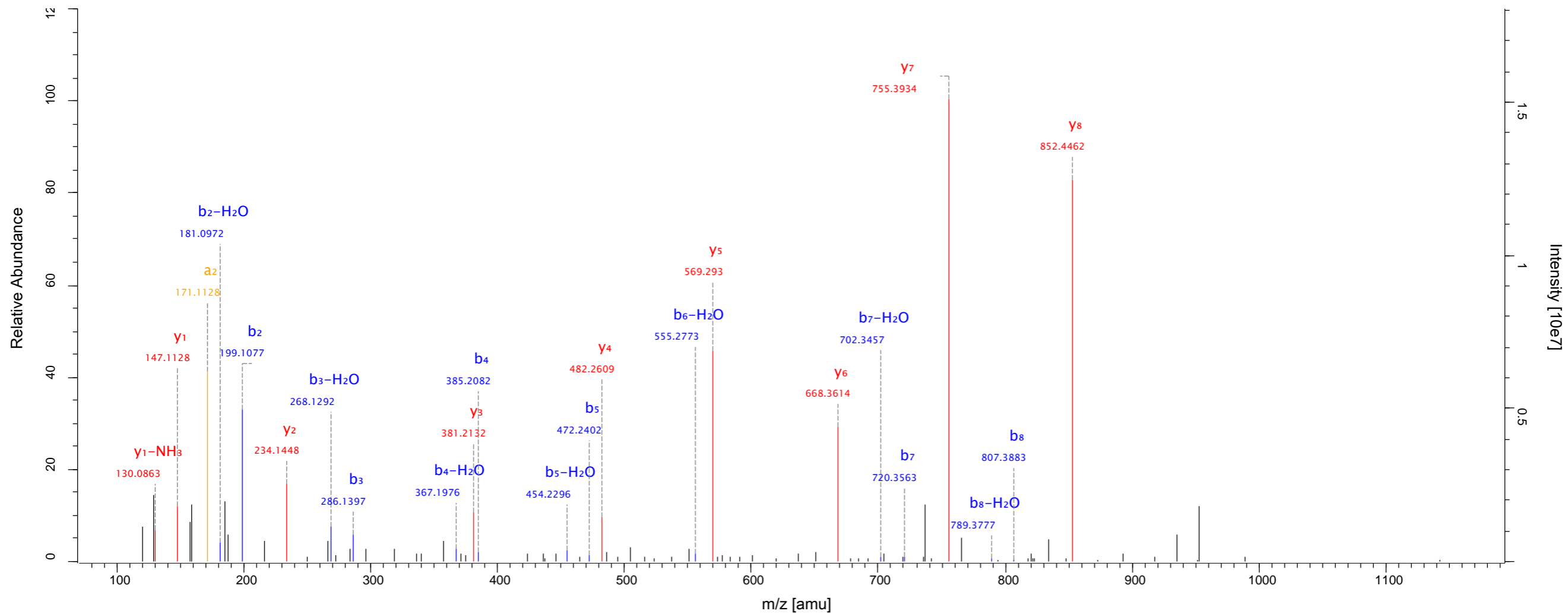
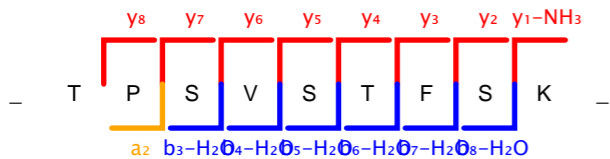
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_27
Scan number: 12291
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 131.26
Method: FTMS; HCD



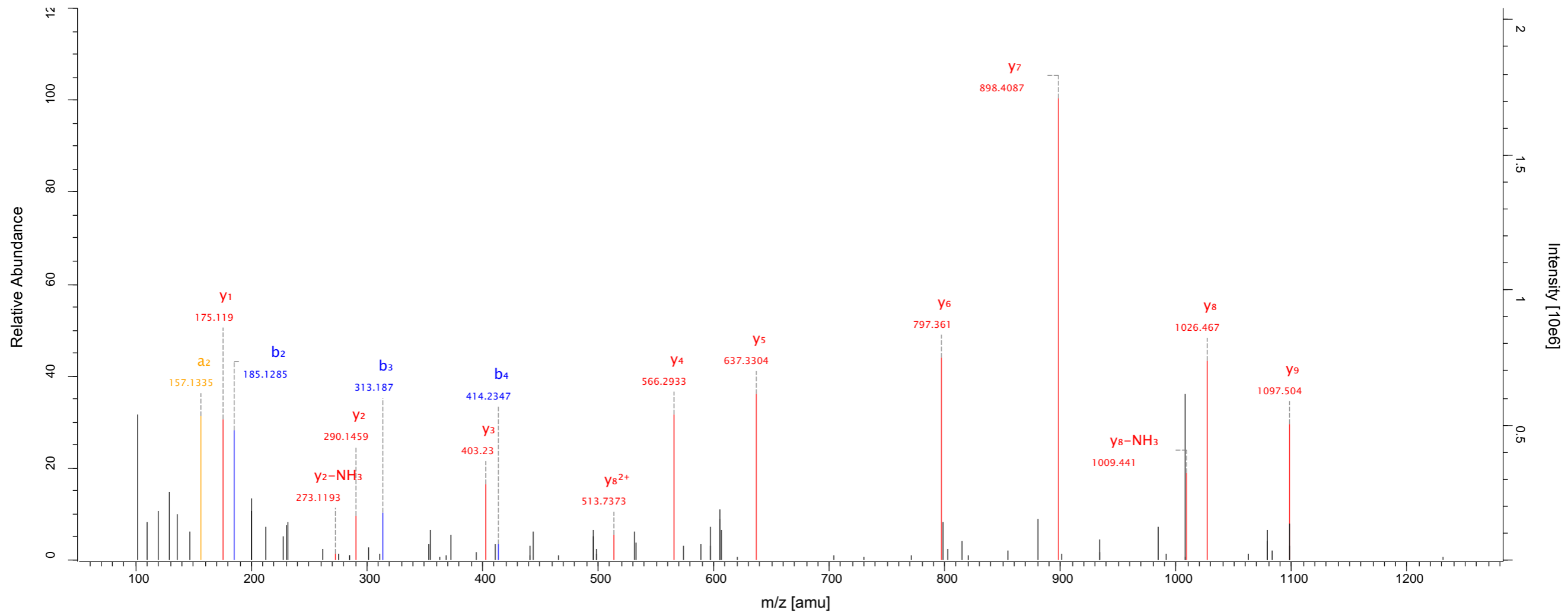
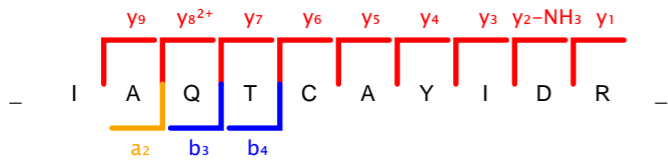
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 13569
 Protein: ORFL190C - NC_006273.2[111965,112384]- <unknown description>
 Peptide score: 162.64
 Method: FTMS; HCD



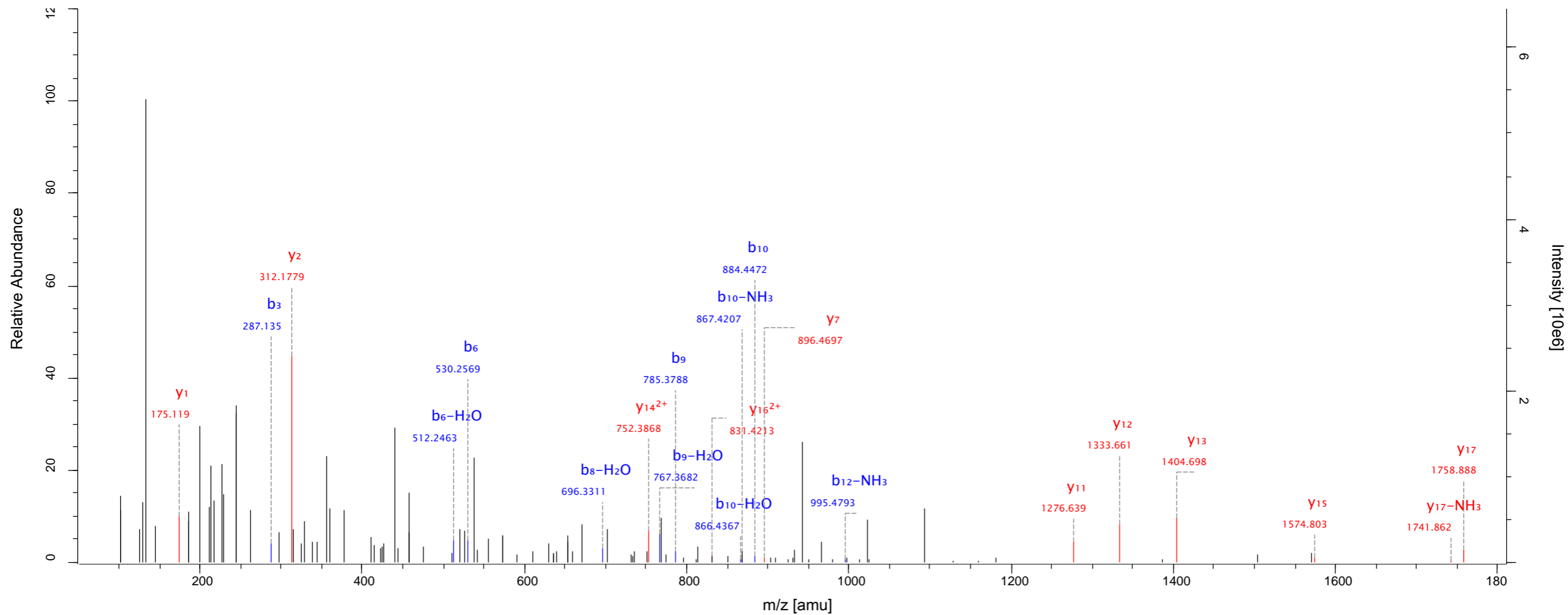
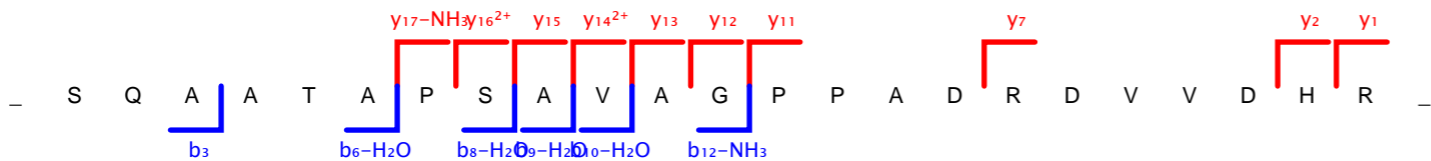
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
 Scan number: 13591
 Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>; ORFL184C.iORF3 - NC_006273.2[111134,111379]- <unknown description>
 Peptide score: 130.31
 Method: FTMS; HCD



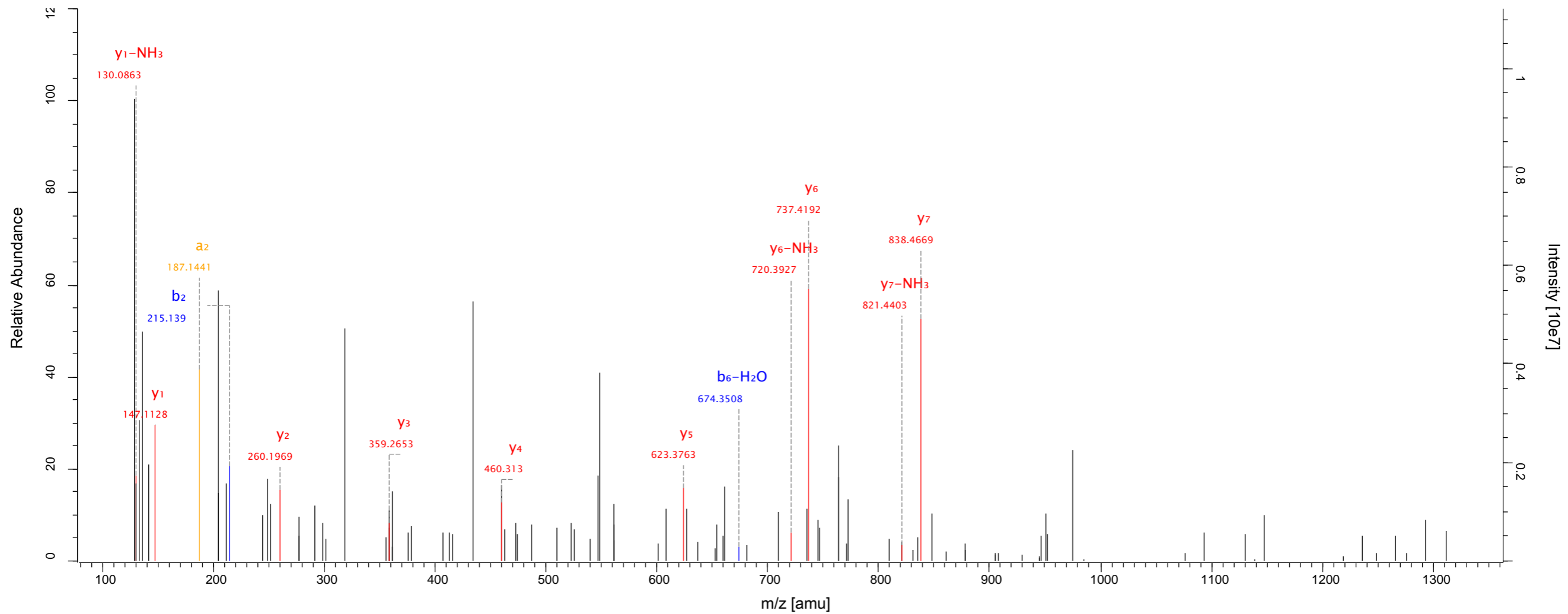
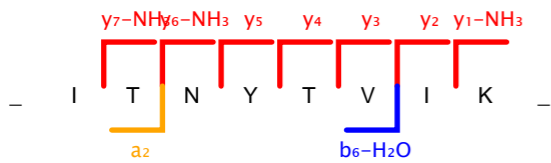
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 15863
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 116.3
Method: FTMS; HCD



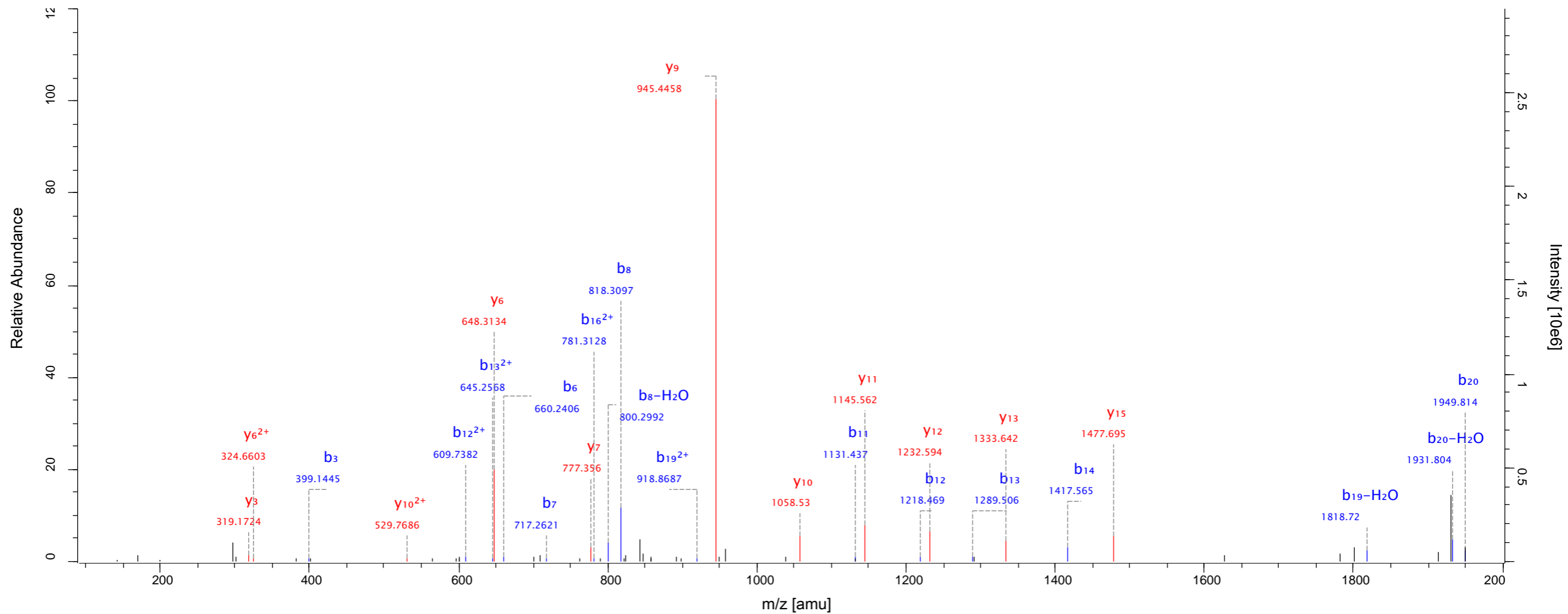
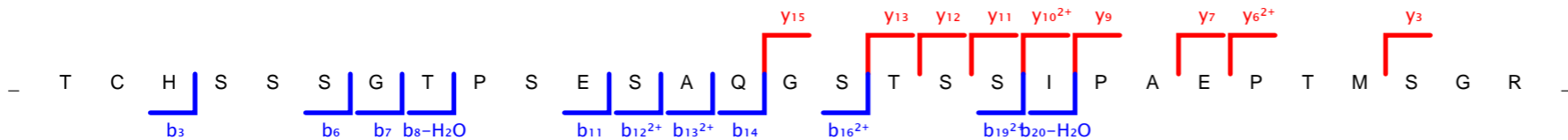
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
 Scan number: 15934
 Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>
 Peptide score: 32.08
 Method: FTMS; HCD



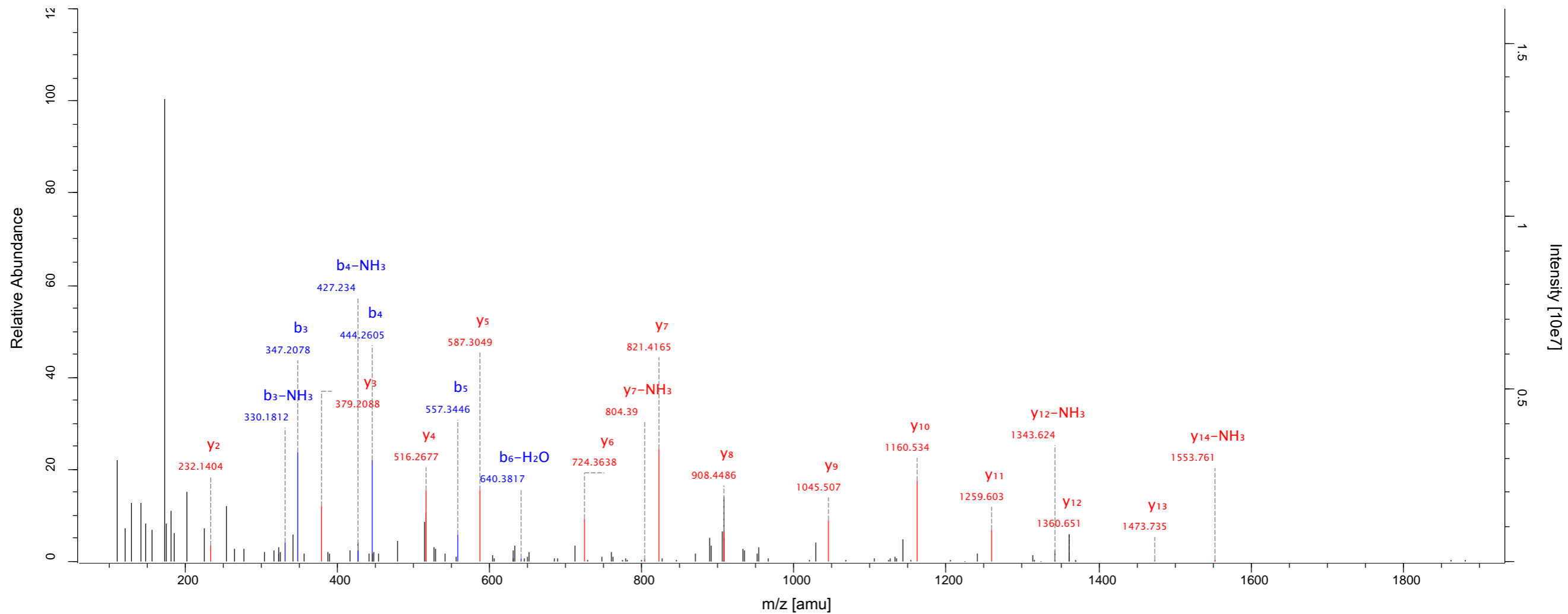
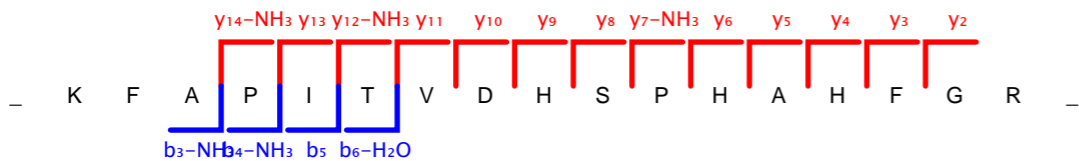
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 17229
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 66.43
Method: FTMS; HCD



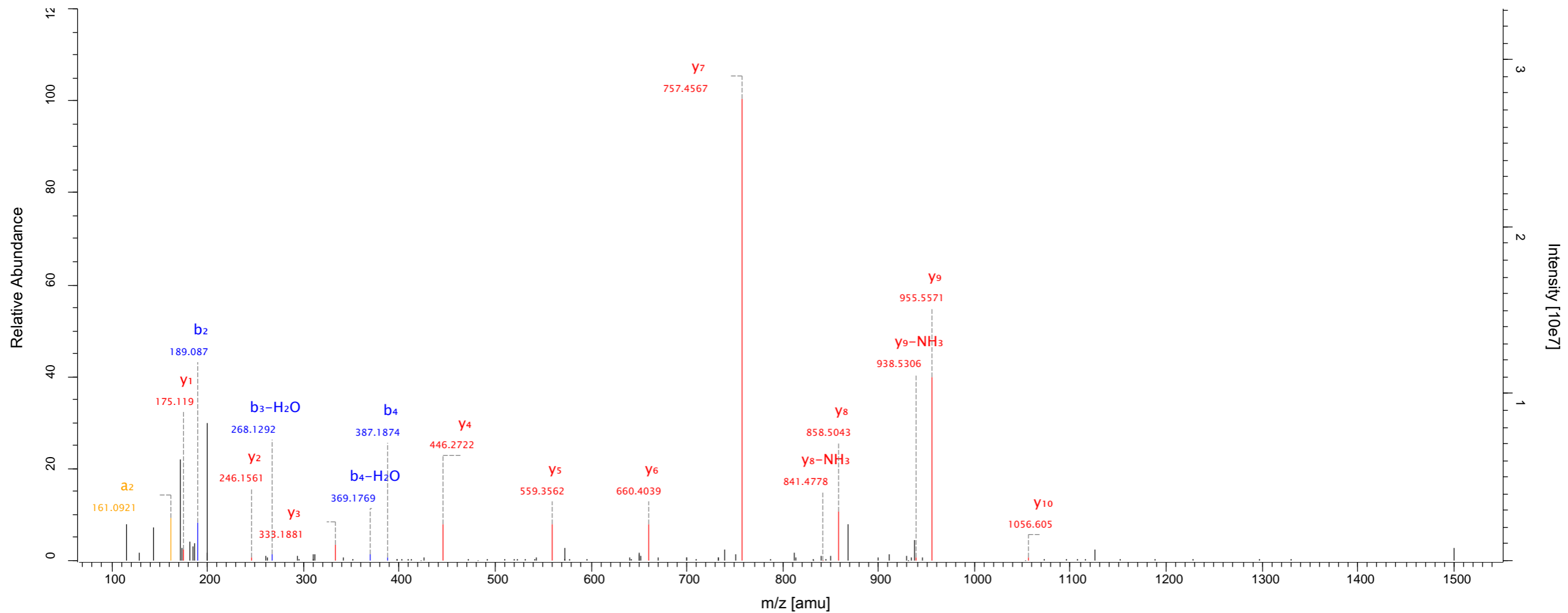
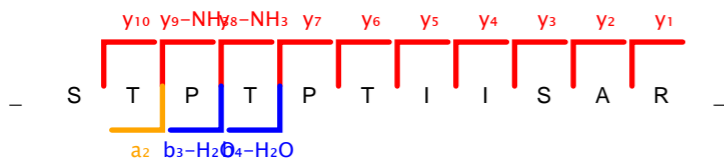
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 17558
 Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
 Peptide score: 92.15
 Method: FTMS; HCD



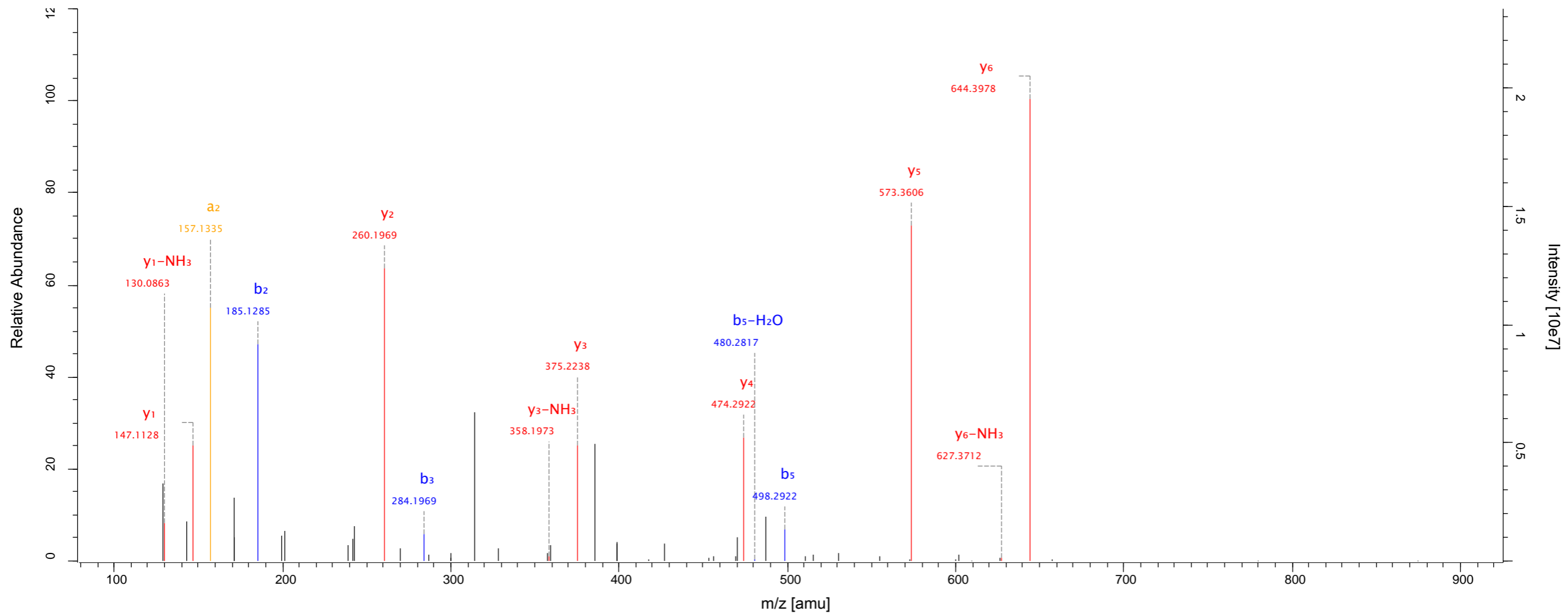
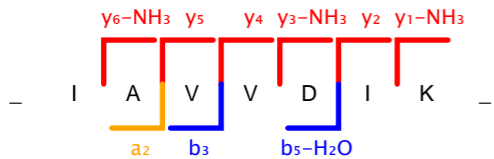
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
 Scan number: 18282
 Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
 Peptide score: 79.02
 Method: FTMS; HCD



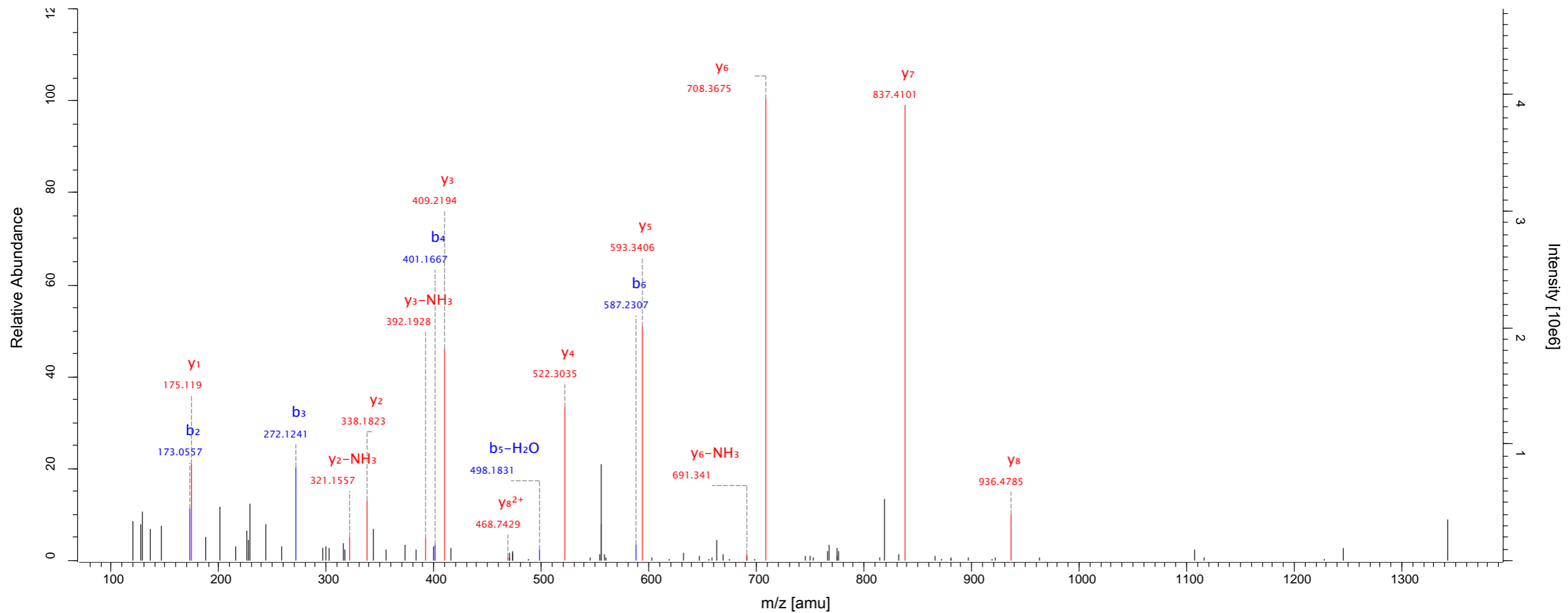
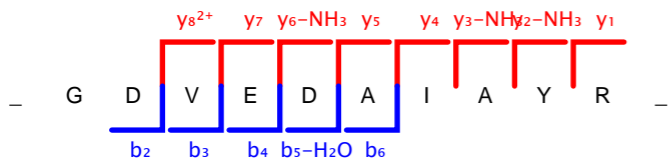
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 19577
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 80.76
Method: FTMS; HCD



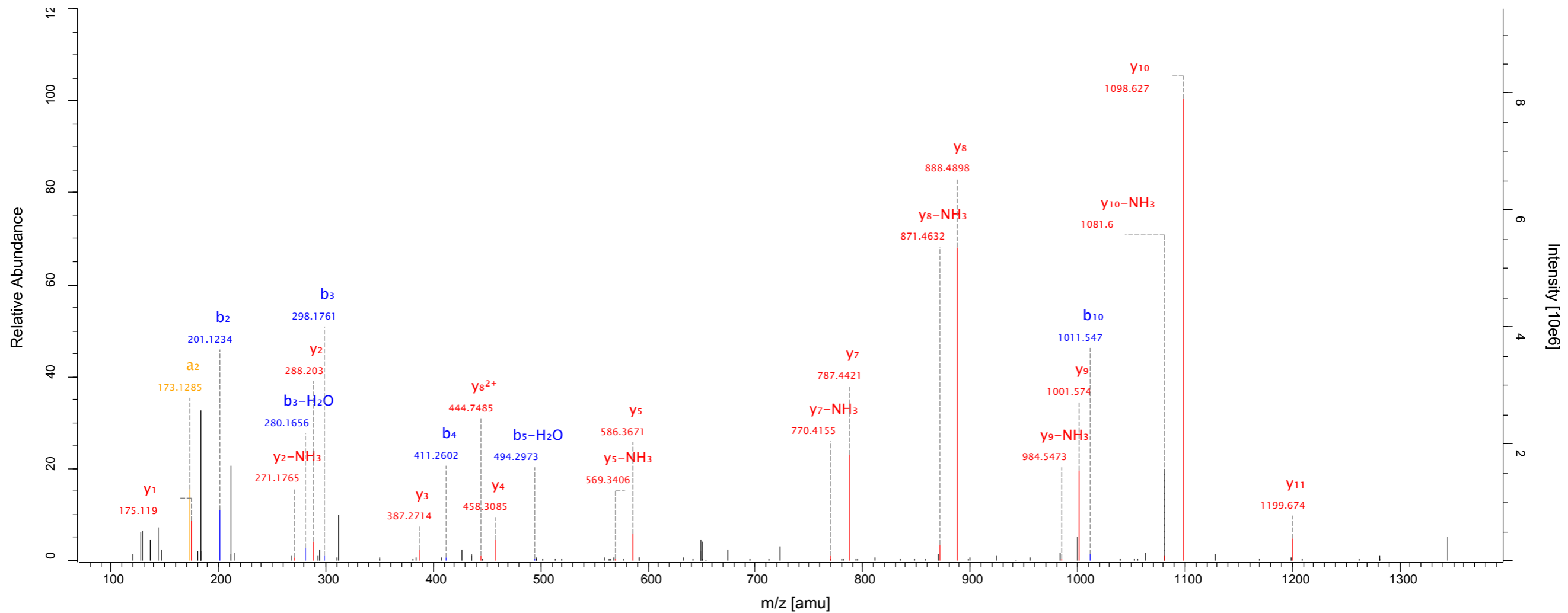
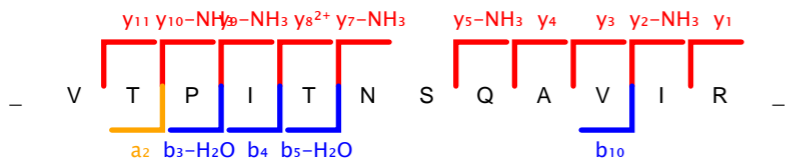
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 19955
Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
Peptide score: 99.92
Method: FTMS; HCD



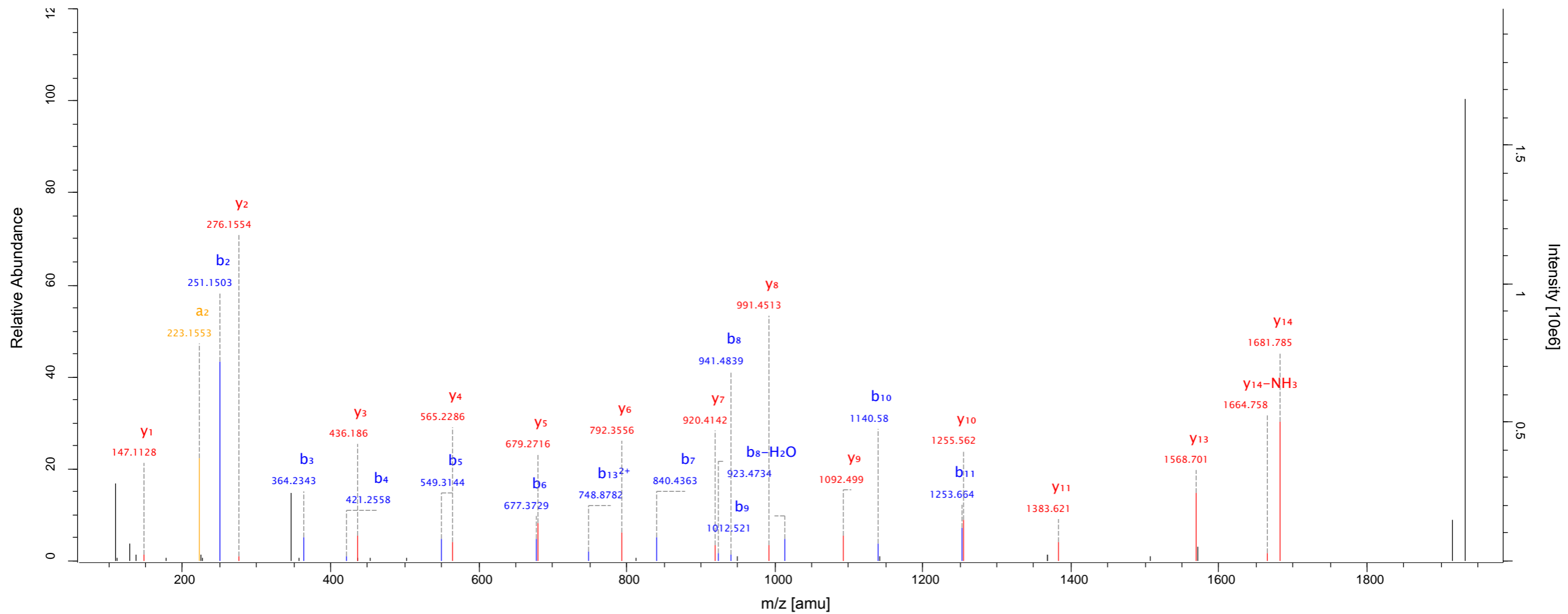
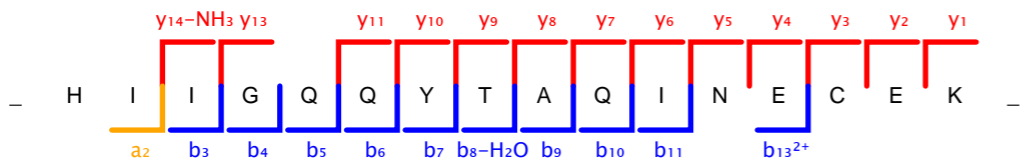
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 20020
Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
Peptide score: 111.61
Method: FTMS; HCD



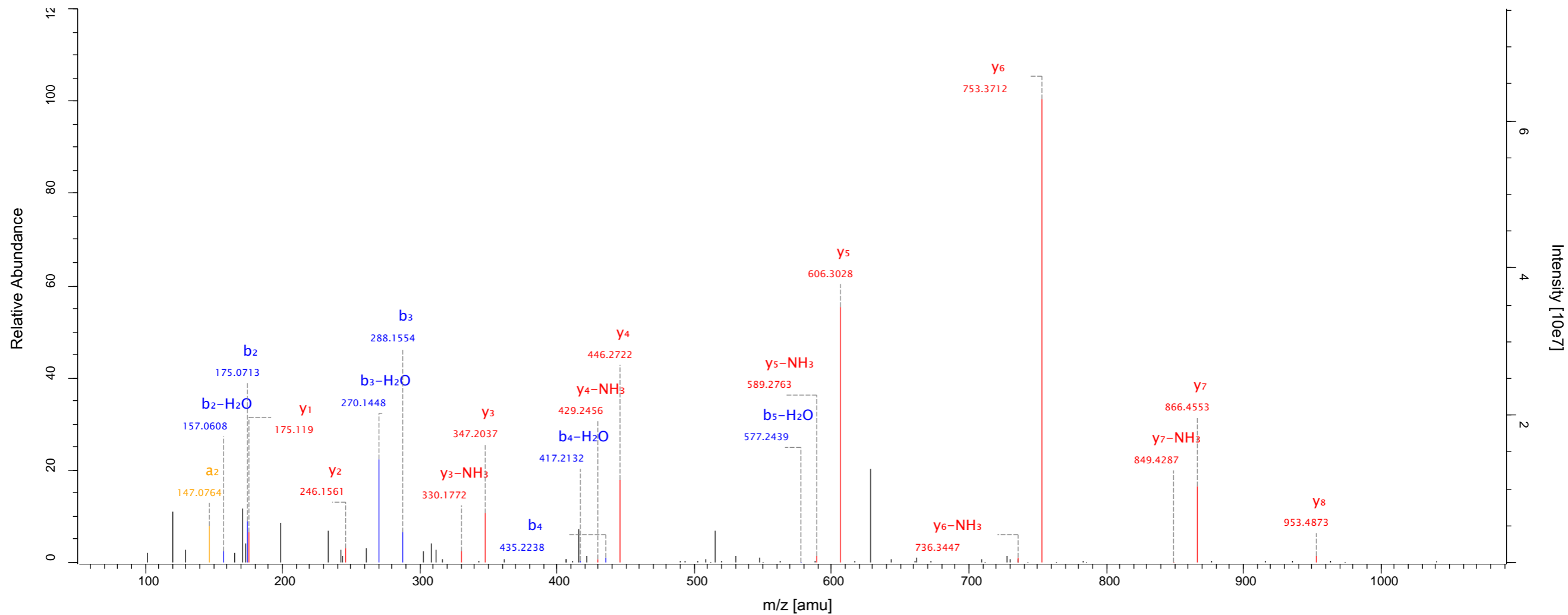
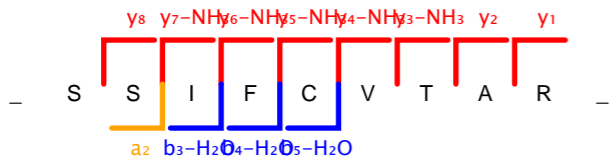
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 22703
 Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
 Peptide score: 125.23
 Method: FTMS; HCD



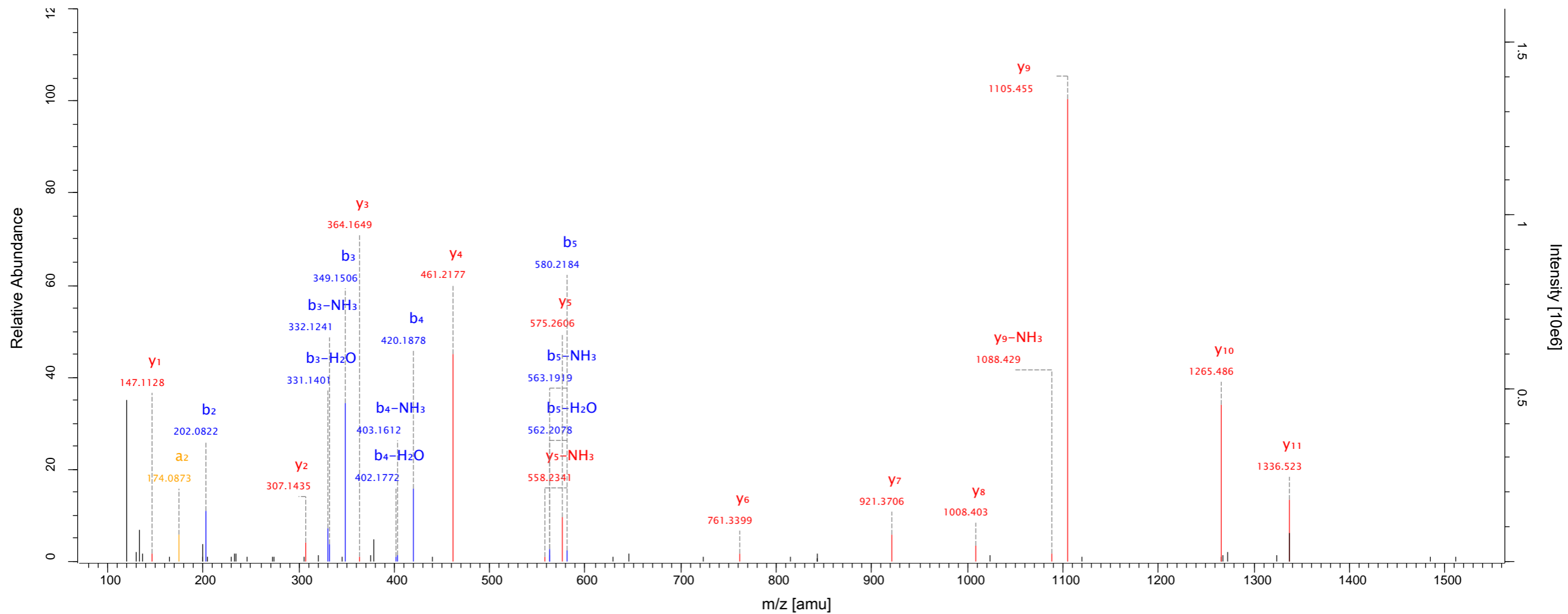
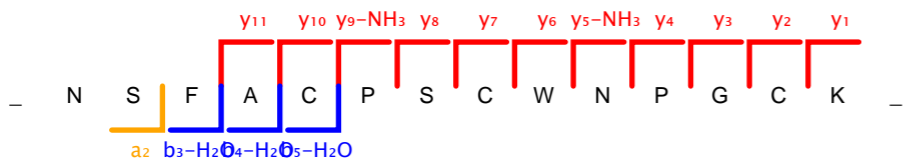
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 22902
Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
Peptide score: 233.42
Method: FTMS; HCD



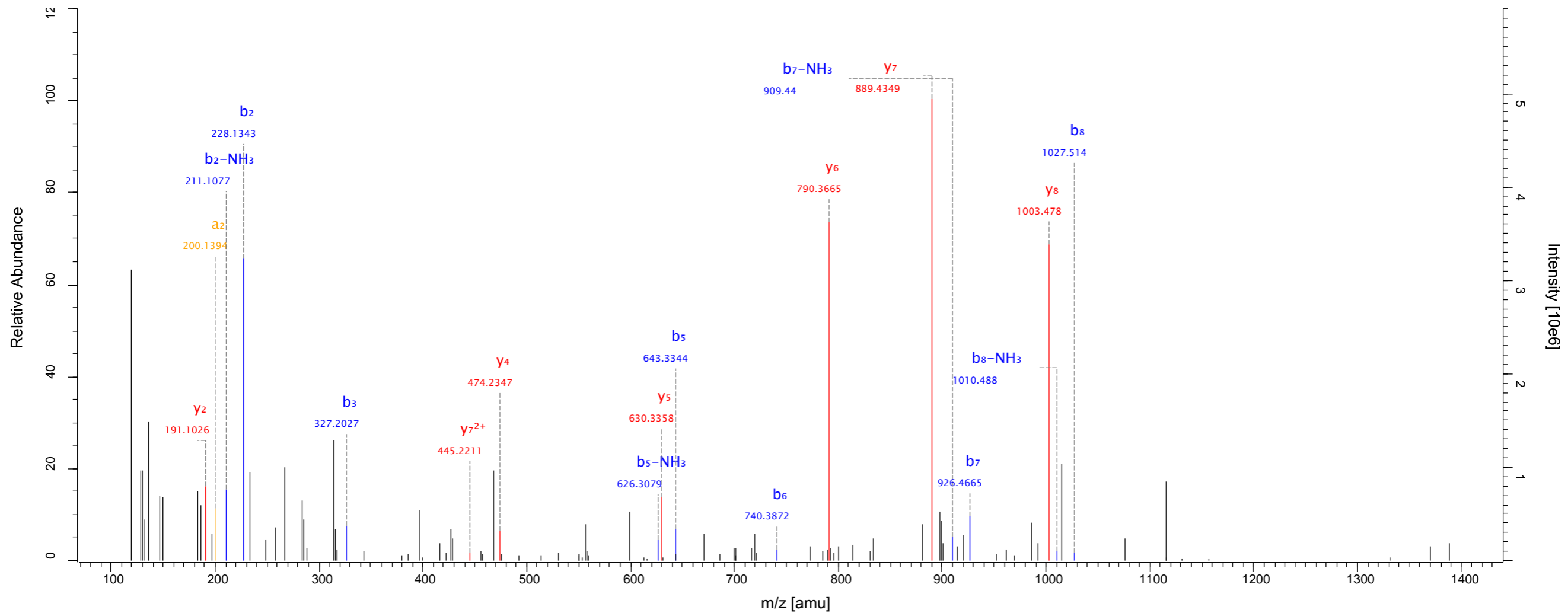
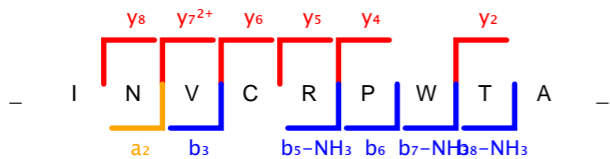
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 24869
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 126.62
Method: FTMS; HCD



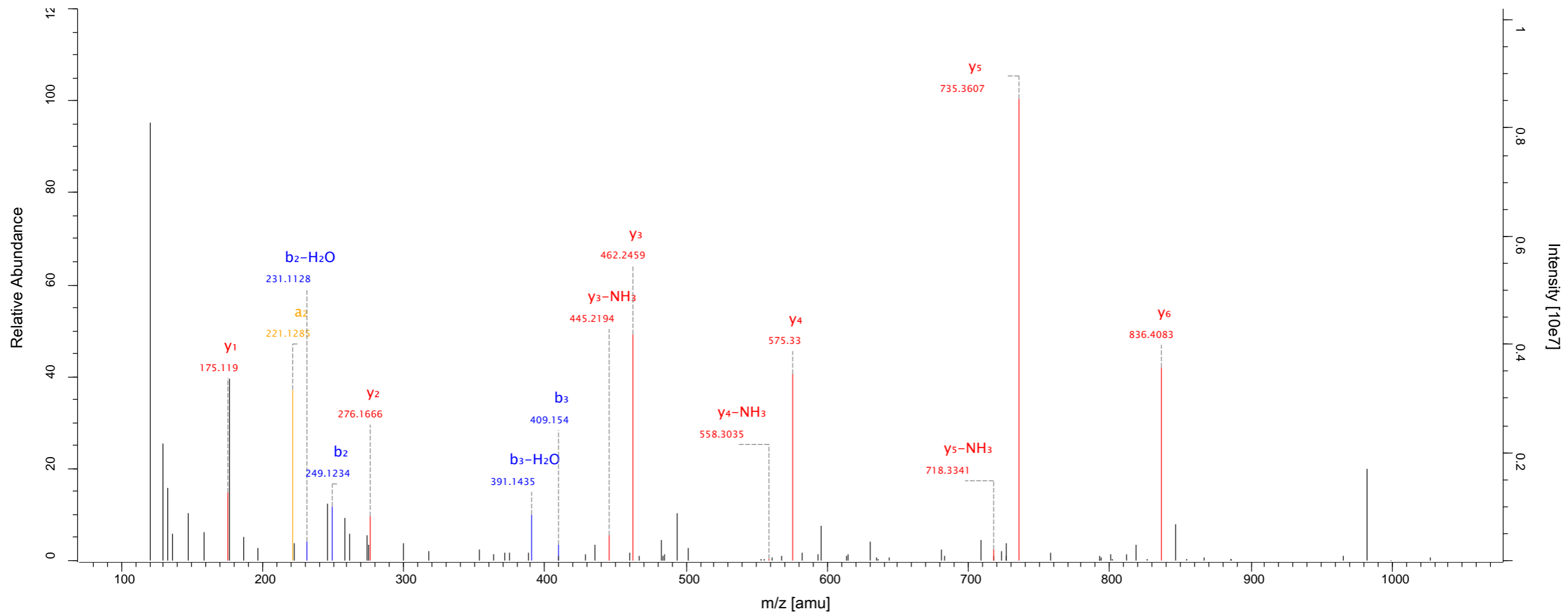
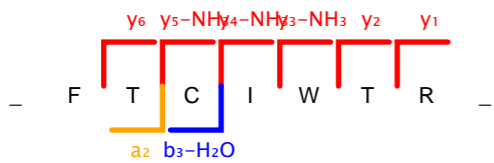
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 25343
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 128.31
 Method: FTMS; HCD



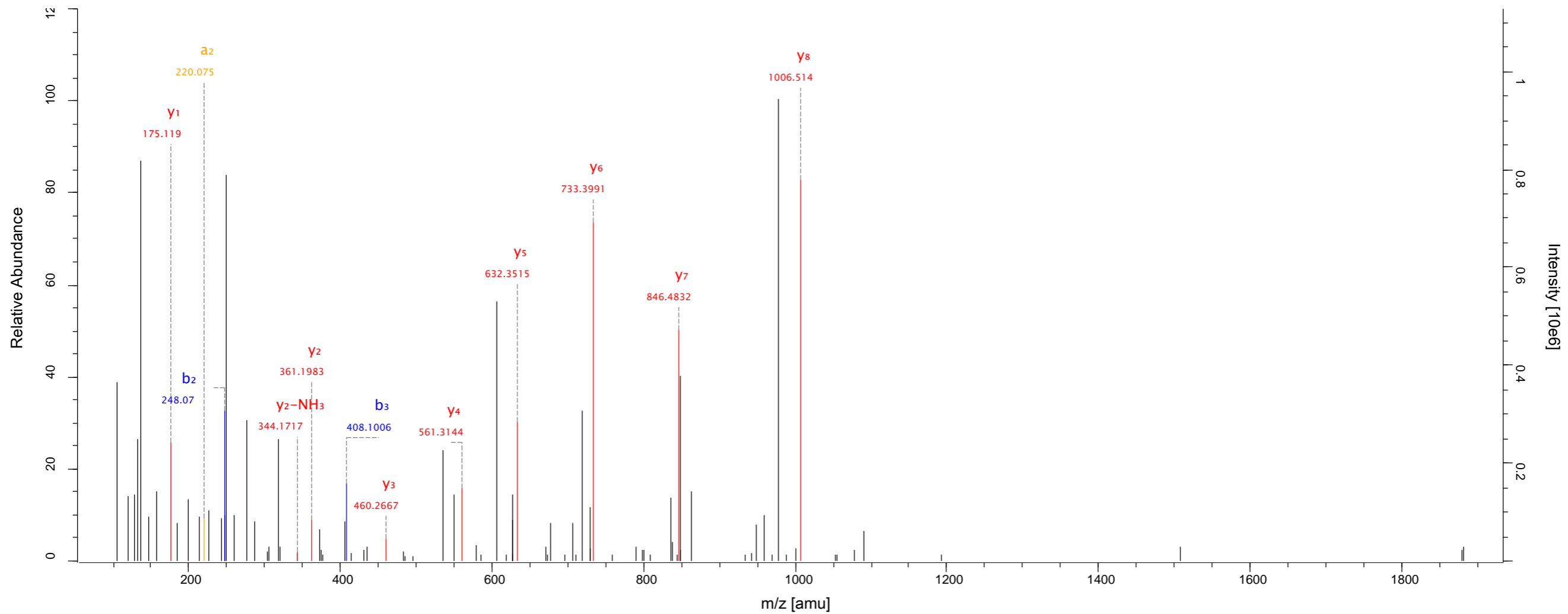
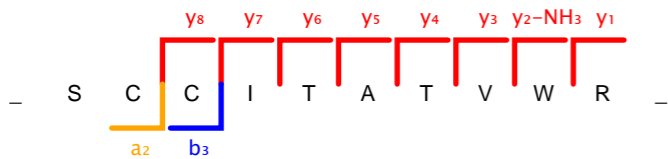
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 25887
Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
Peptide score: 125.81
Method: FTMS; HCD



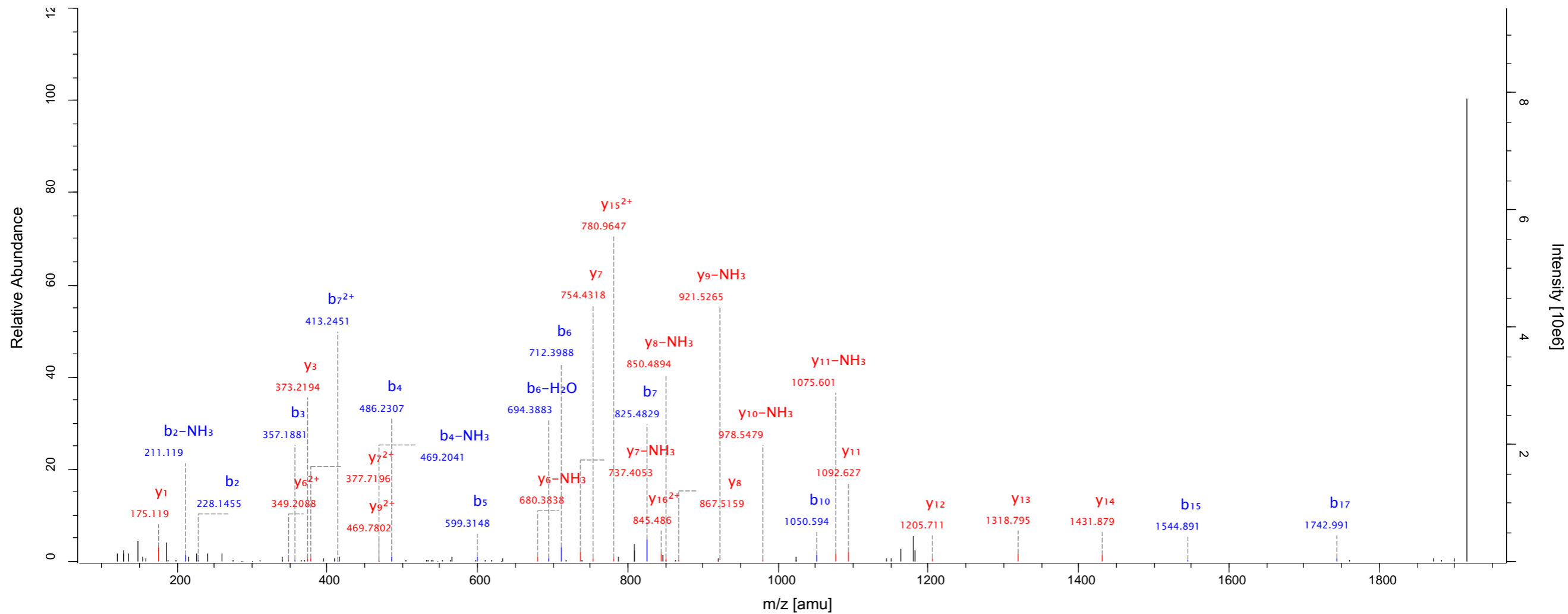
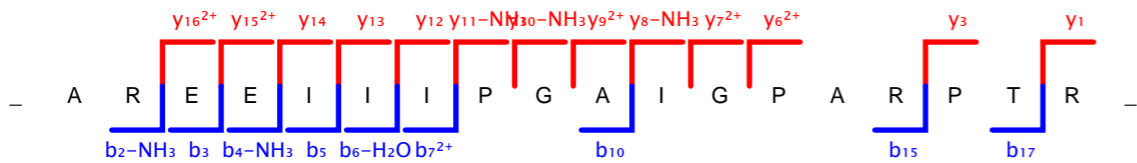
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 26027
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 100.07
Method: FTMS; HCD



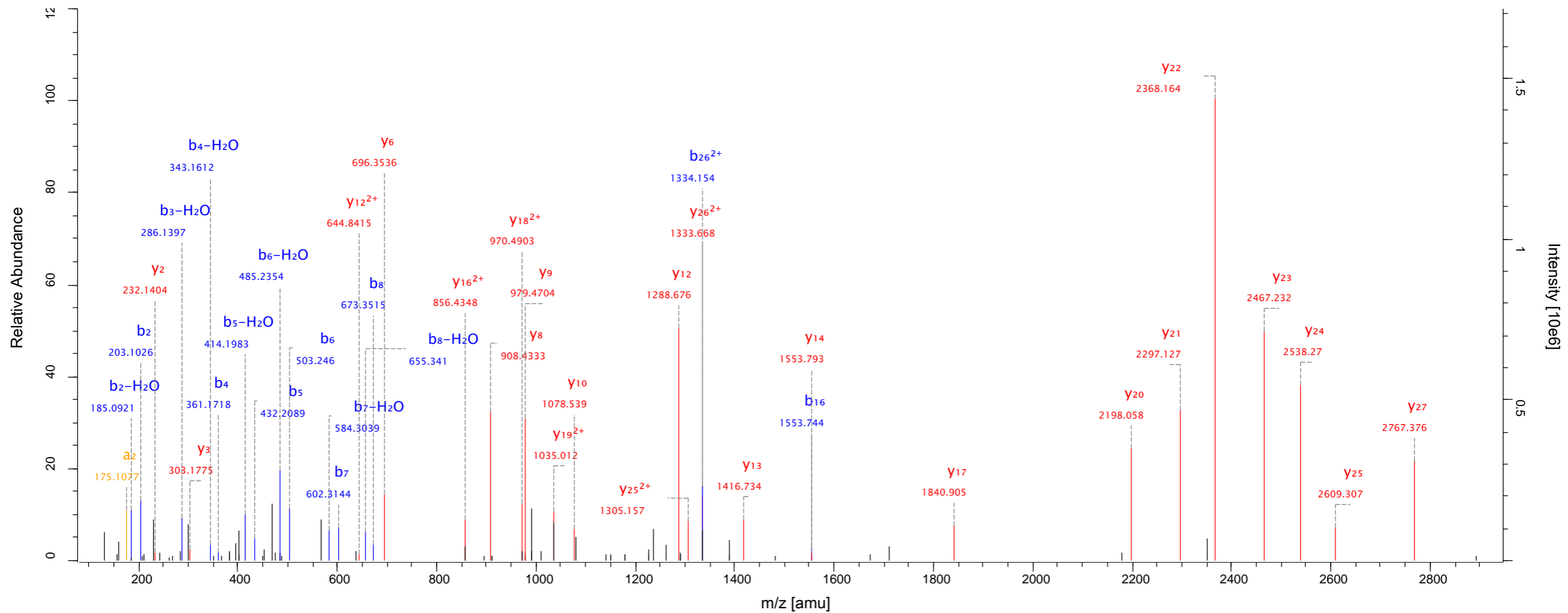
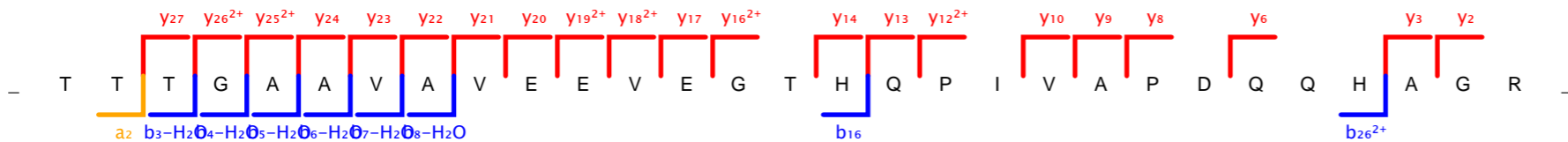
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 27227
Protein: ORFL47W.iORF1 + NC_006273.2[21305,21601]+ <unknown description>
Peptide score: 71.35
Method: FTMS; HCD



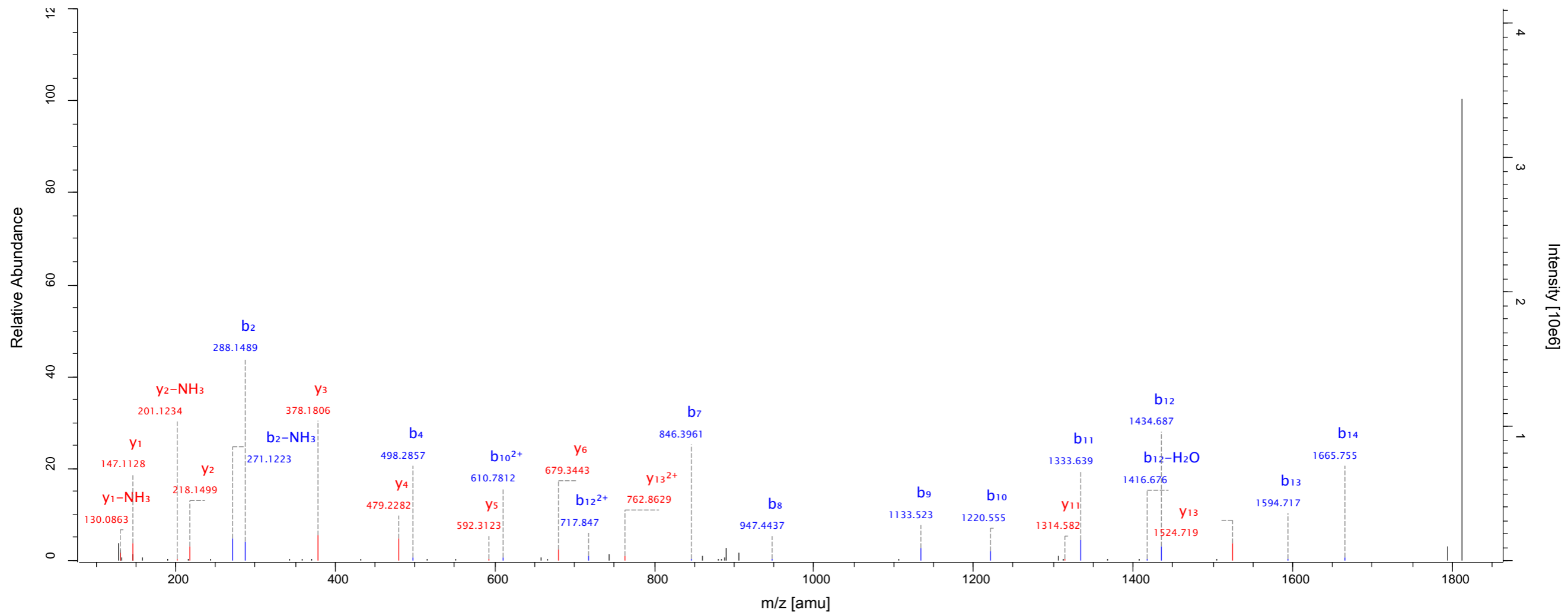
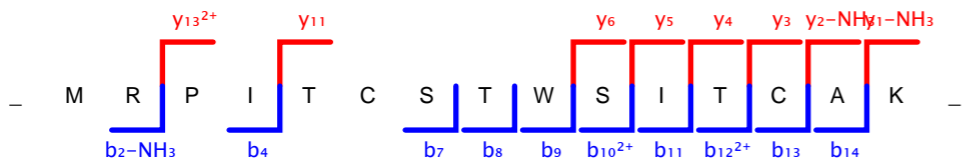
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
 Scan number: 29928
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 103.67
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 29980
 Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
 Peptide score: 201.15
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 30520
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 169.19
Method: FTMS; HCD



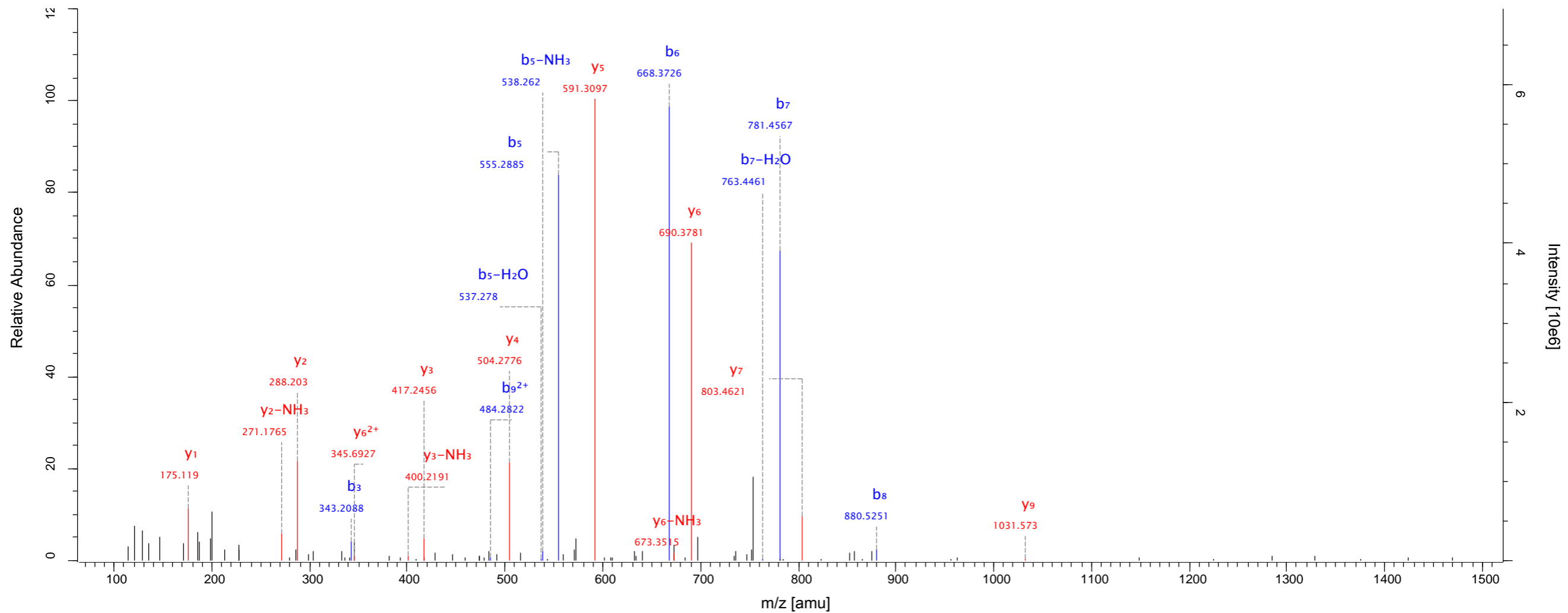
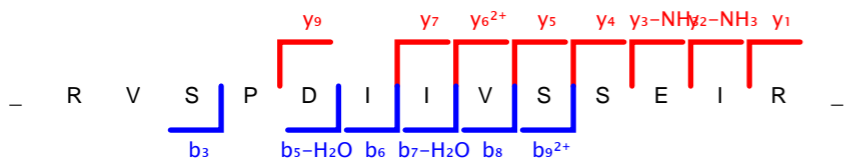
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27

Scan number: 30970

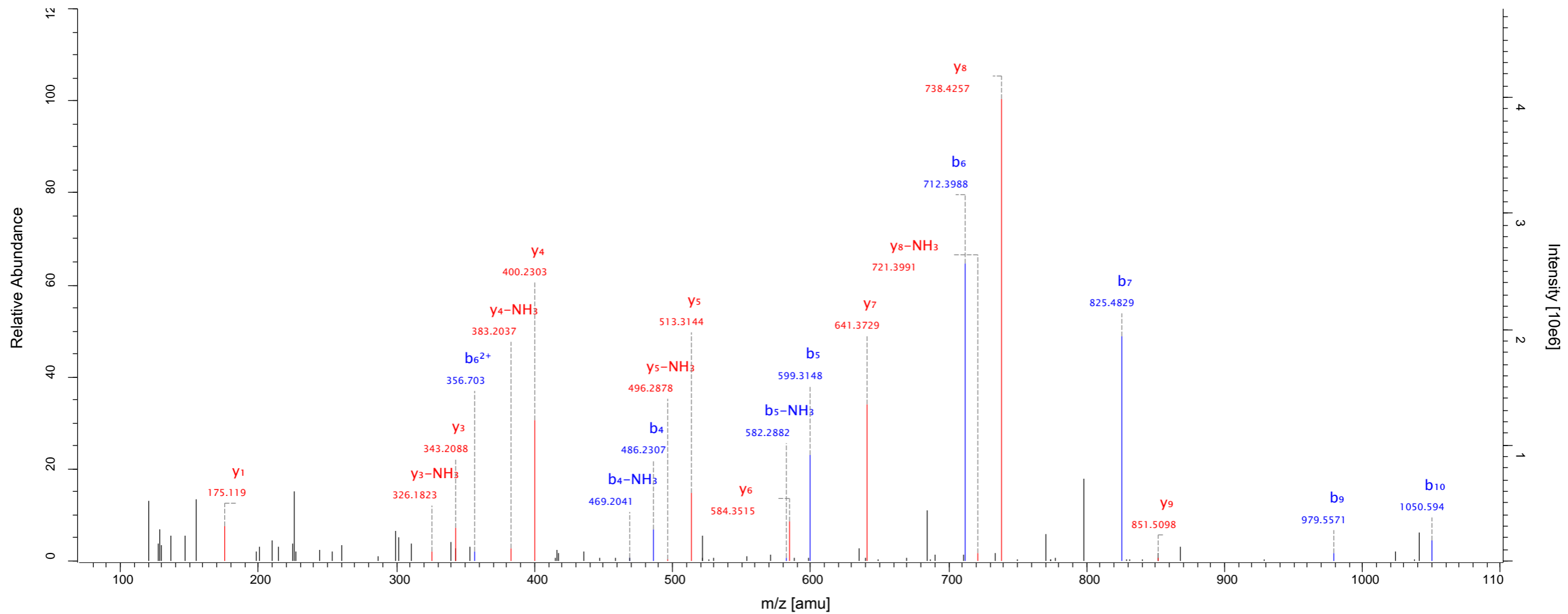
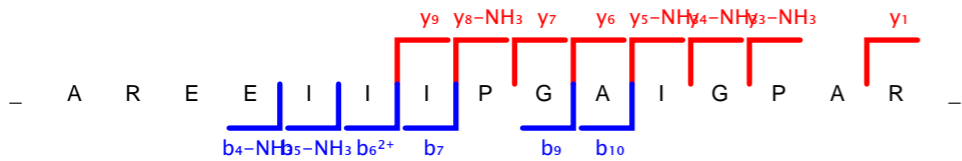
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 113.65

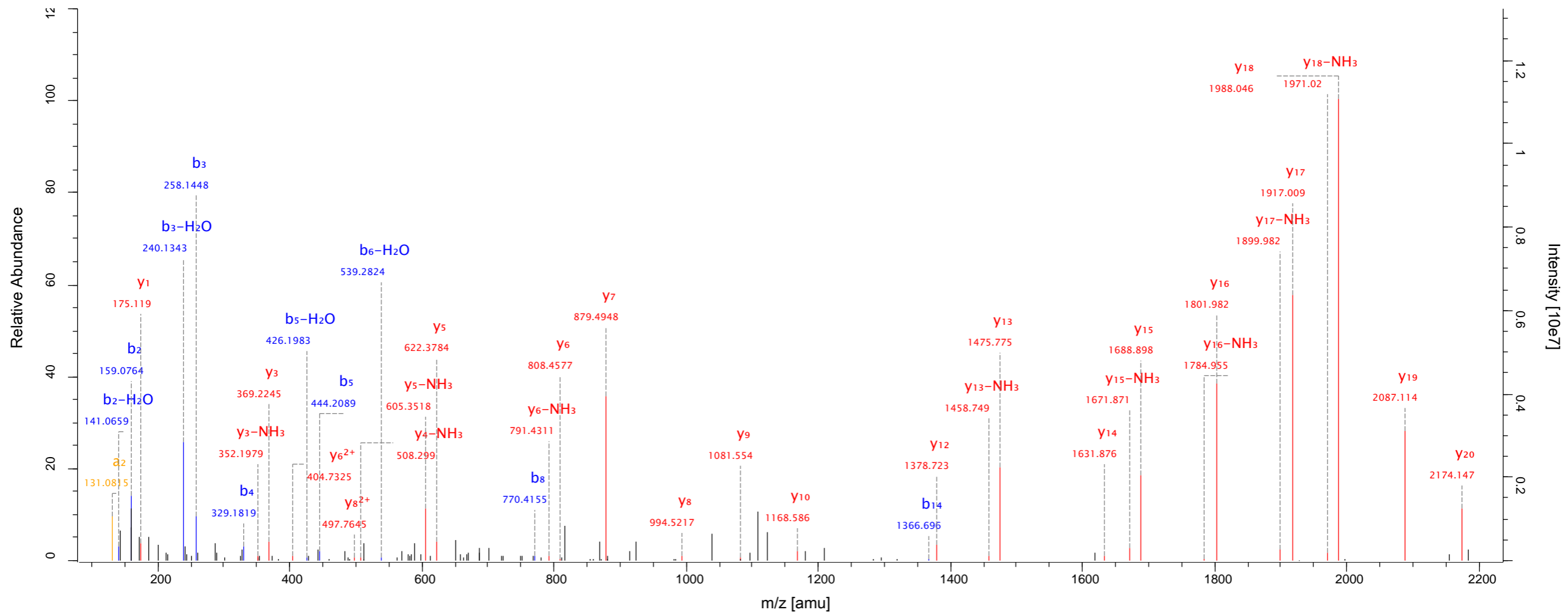
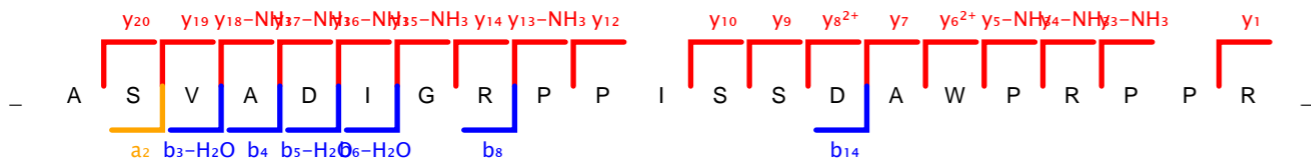
Method: FTMS; HCD



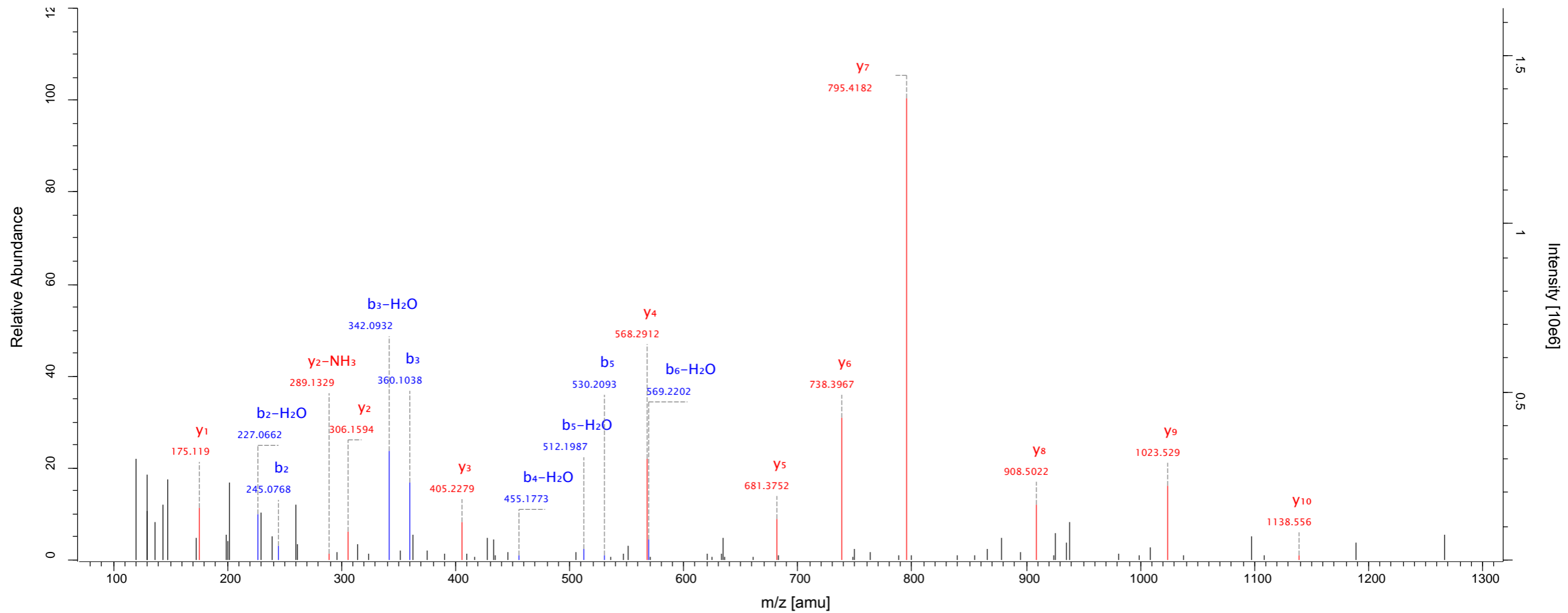
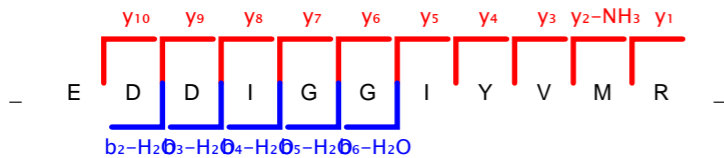
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 33014
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 92.06
Method: FTMS; HCD



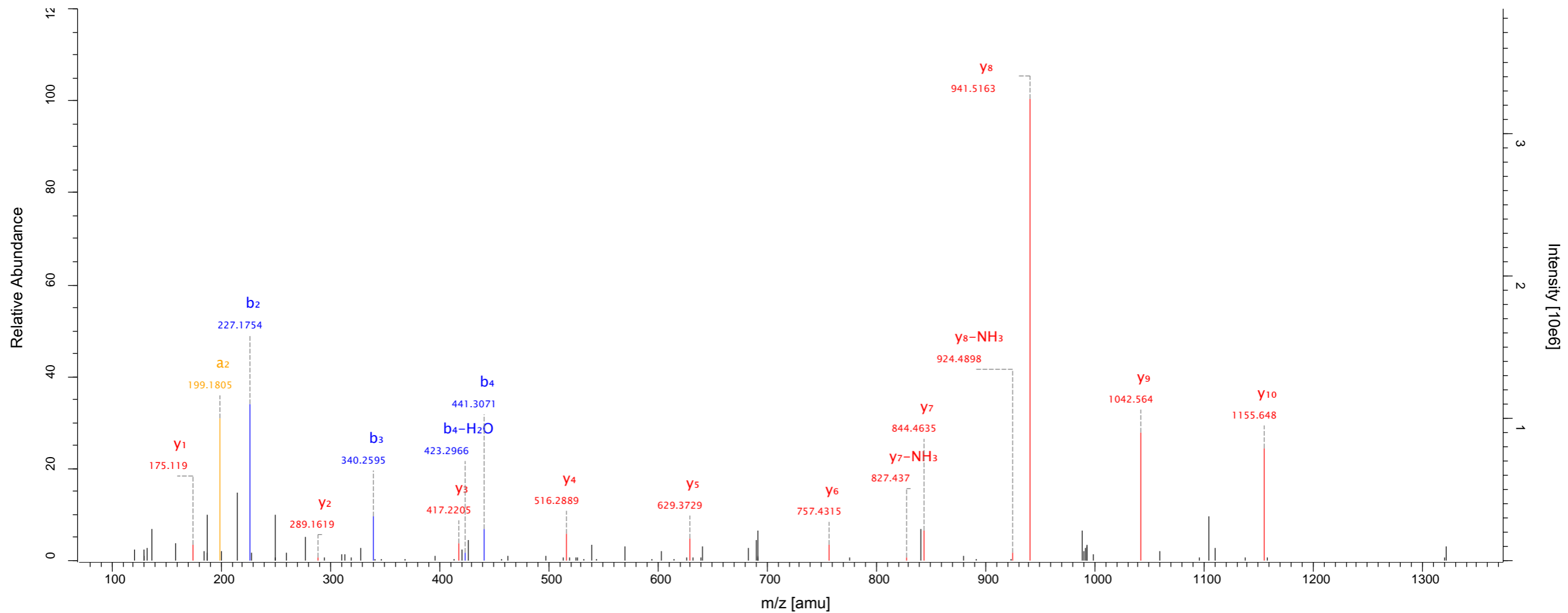
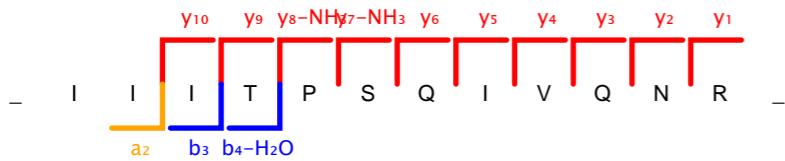
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 34075
 Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
 Peptide score: 195.27
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 34177
Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
Peptide score: 84.66
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVpI1_27
Scan number: 34920
Protein: ORFL190C - NC_006273.2[111965,112384]- <unknown description>
Peptide score: 104.79
Method: FTMS; HCD



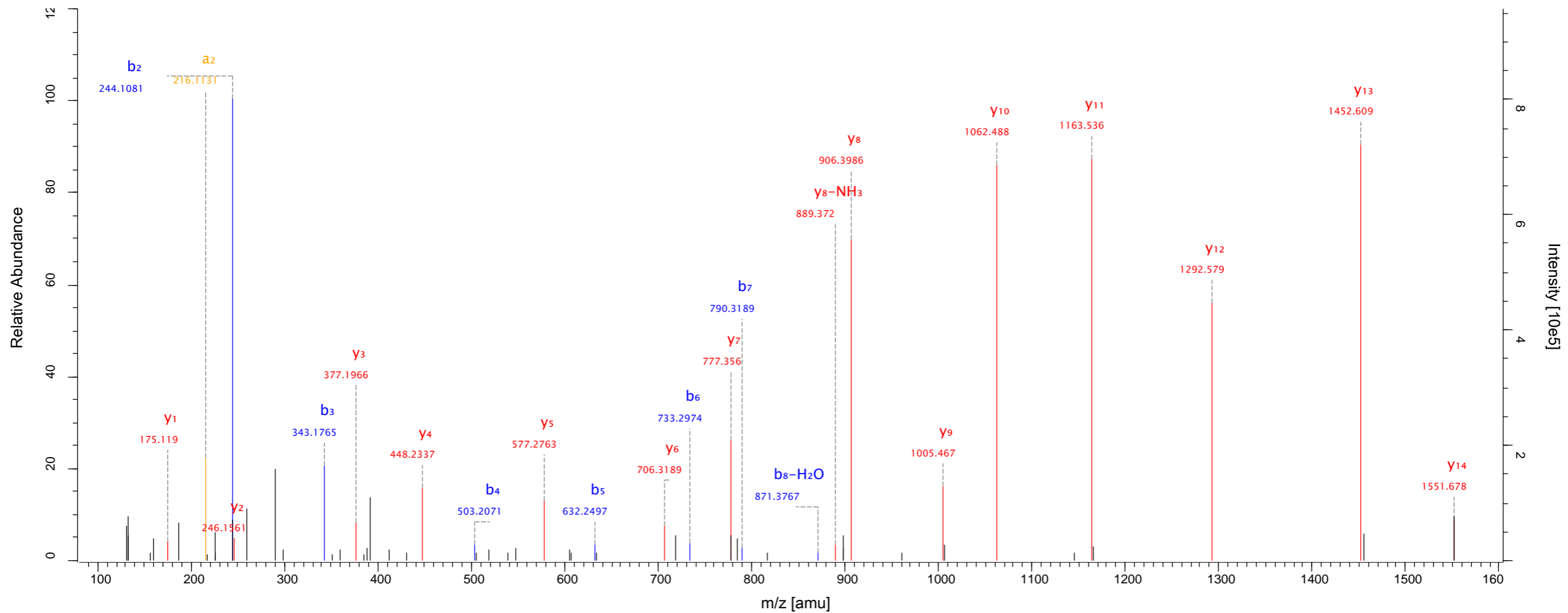
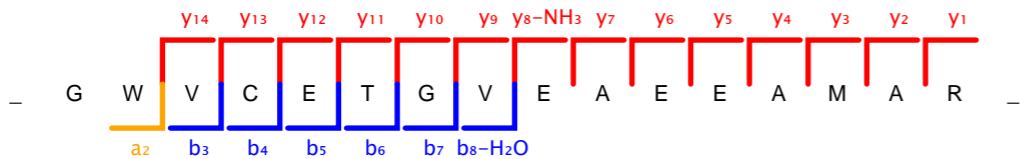
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27

Scan number: 35182

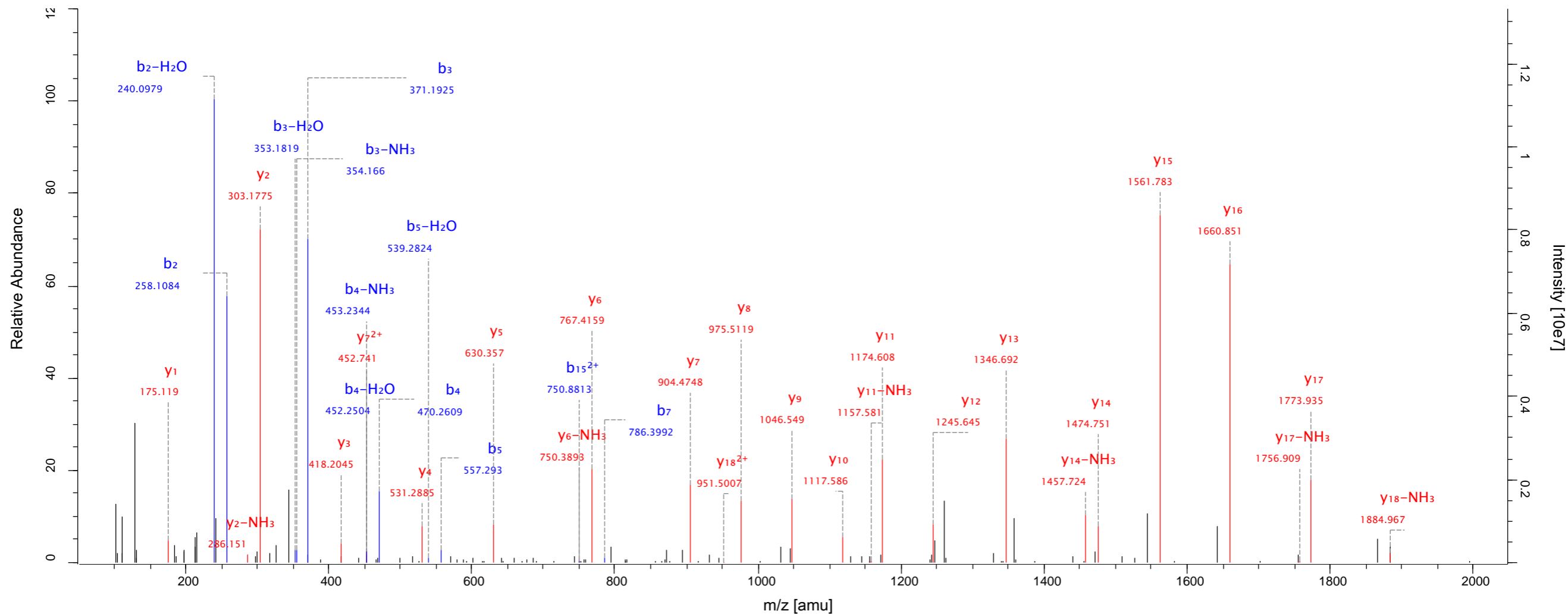
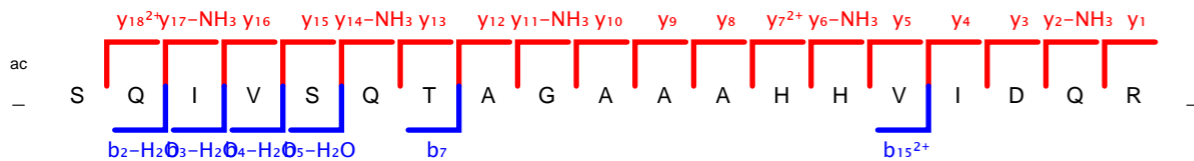
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 124.45

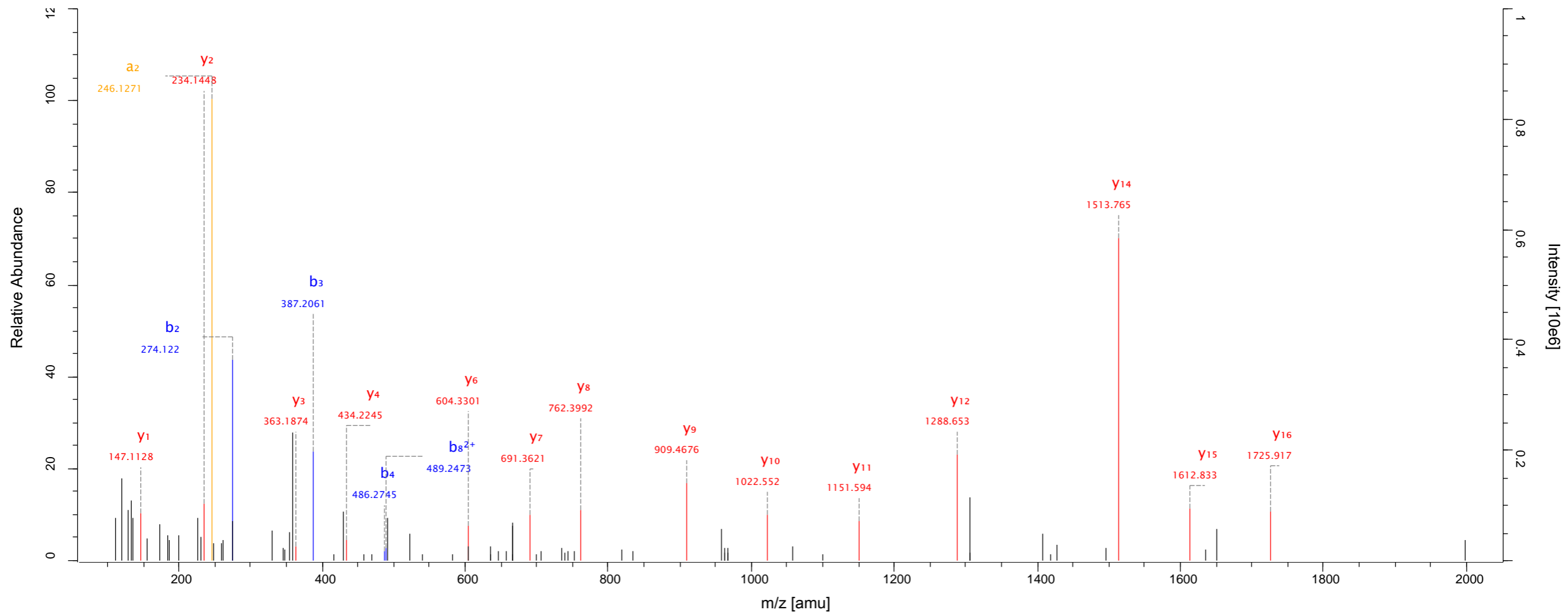
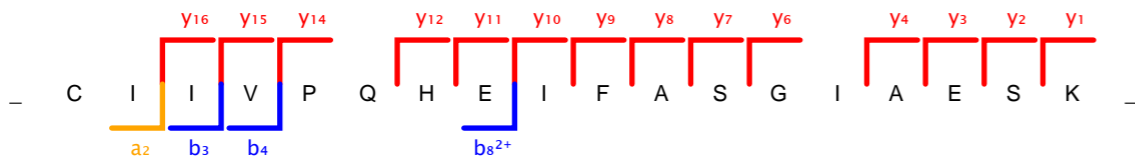
Method: FTMS; HCD



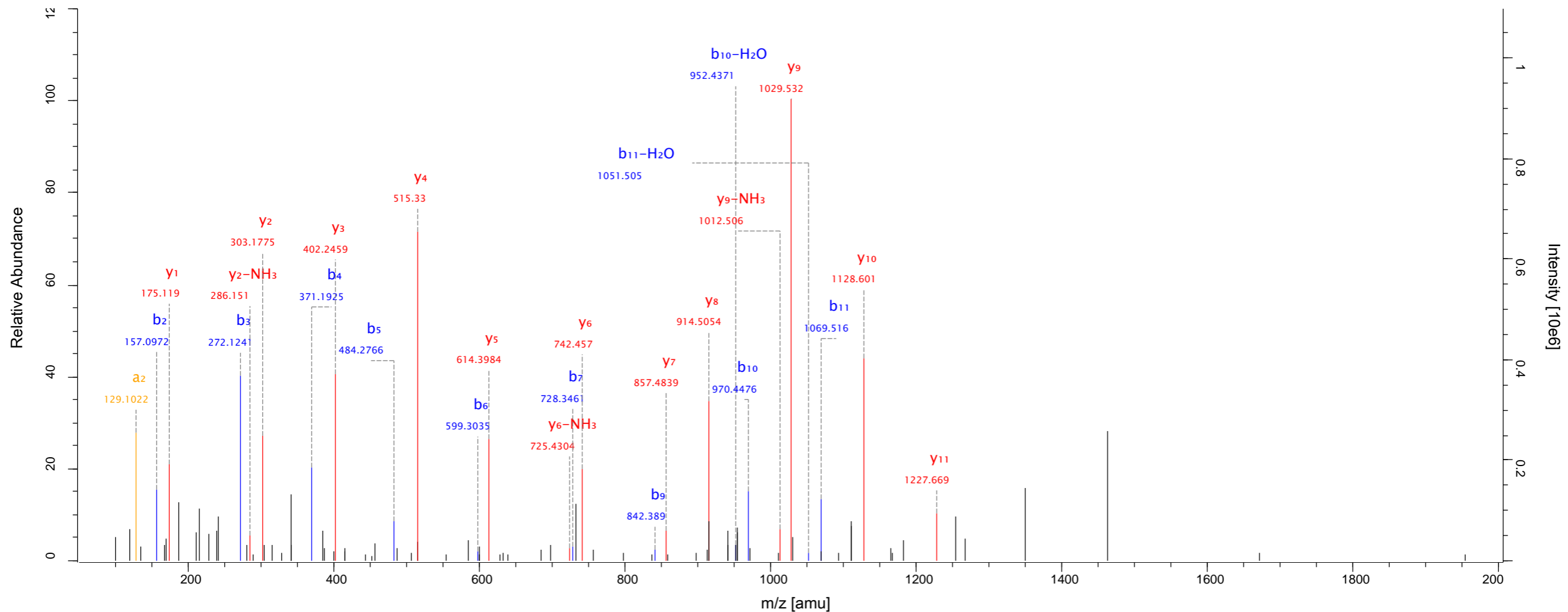
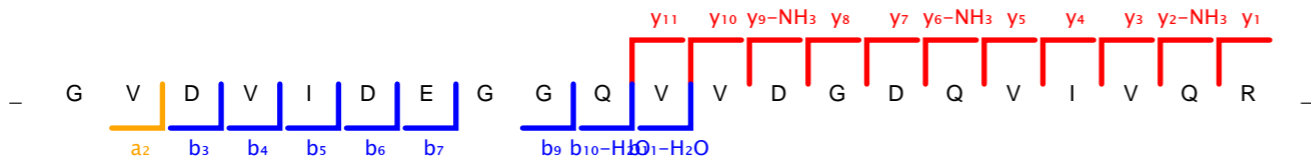
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
 Scan number: 38592
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 171.56
 Method: FTMS; HCD



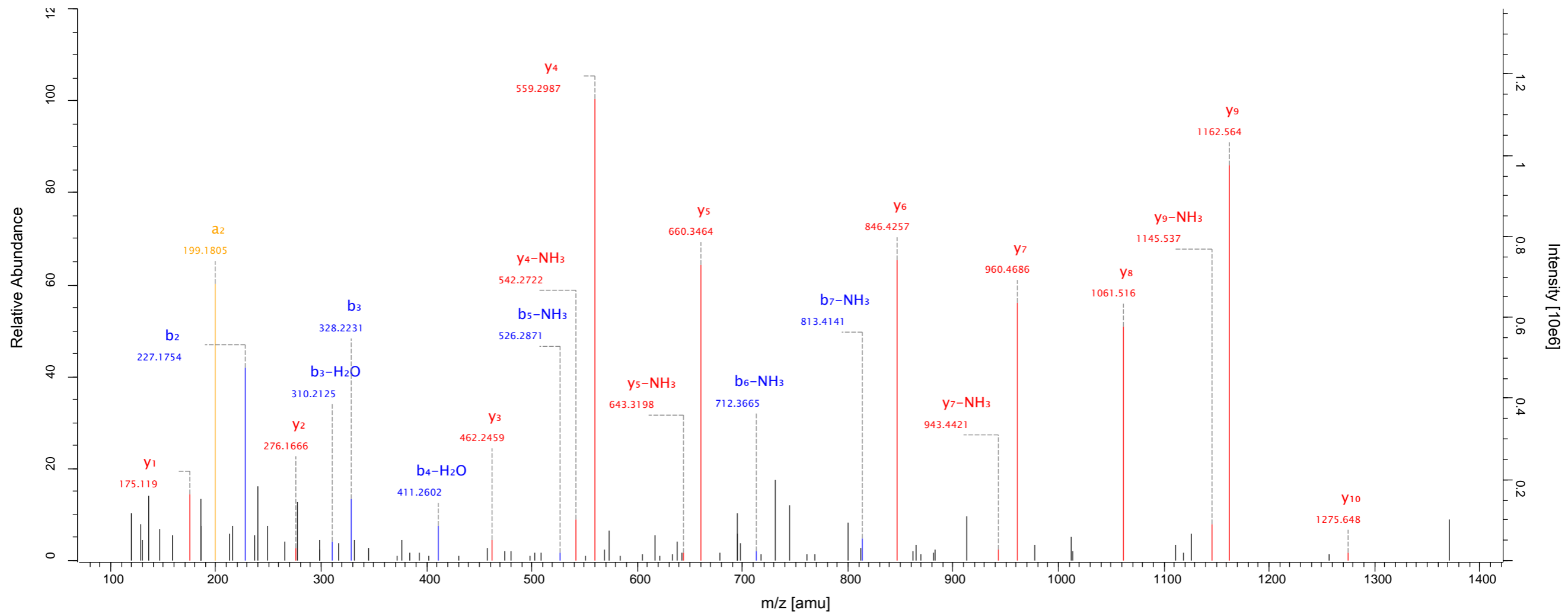
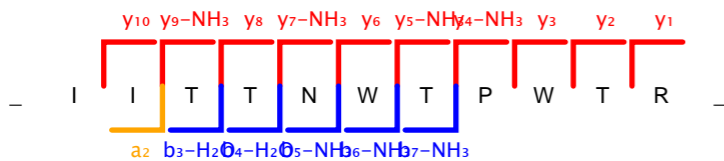
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 39244
Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
Peptide score: 94.49
Method: FTMS; HCD



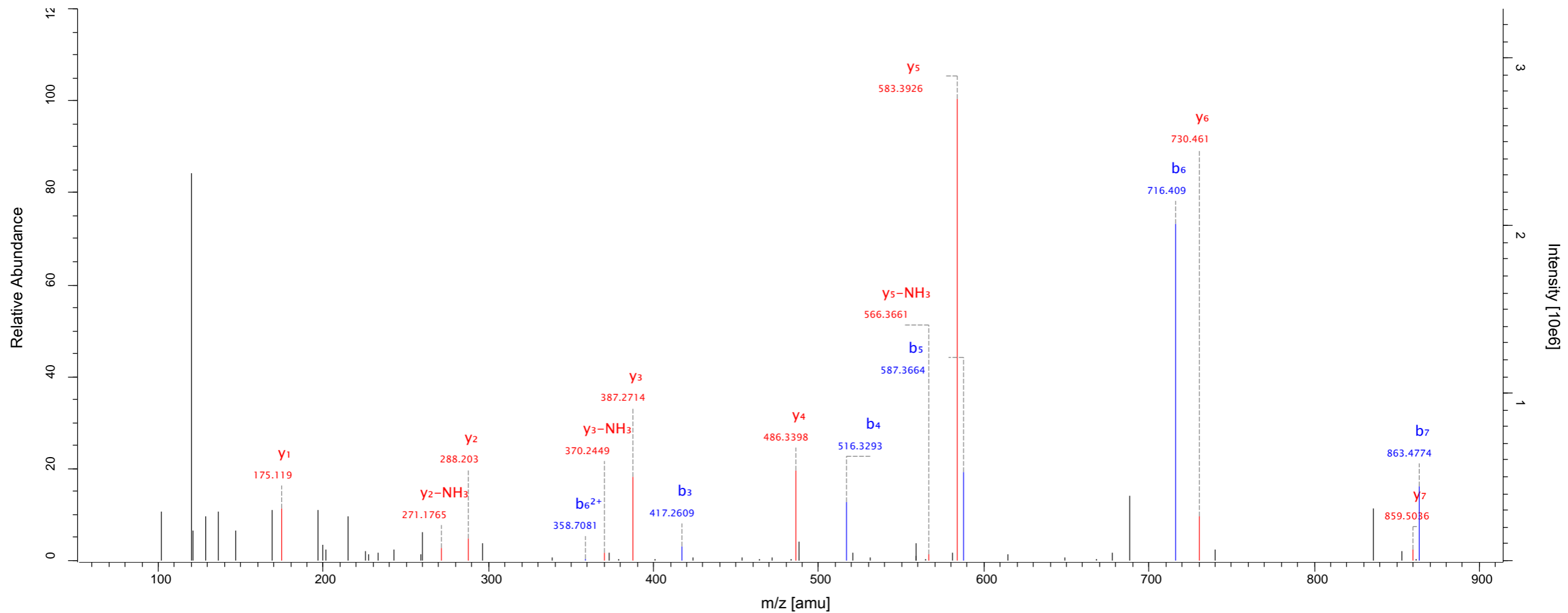
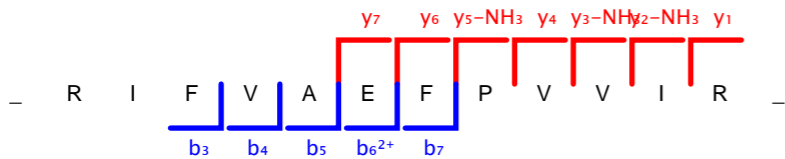
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 40618
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 125.12
 Method: FTMS; HCD



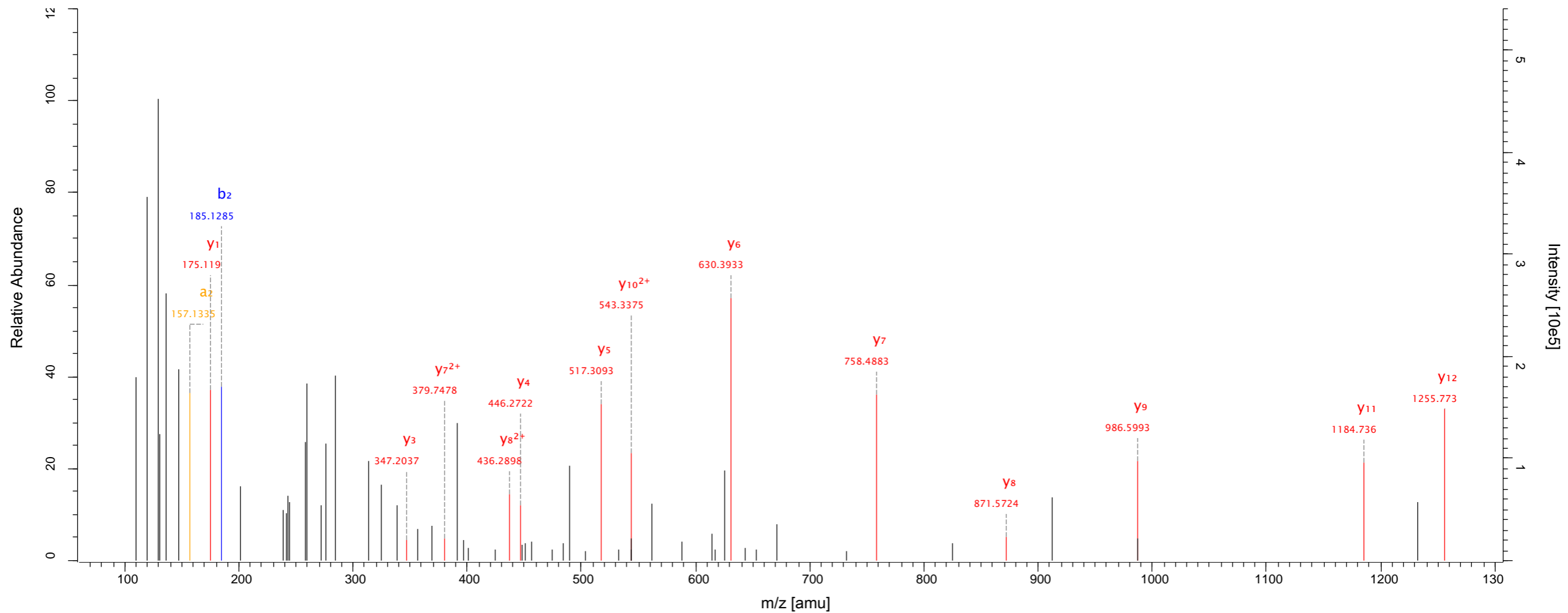
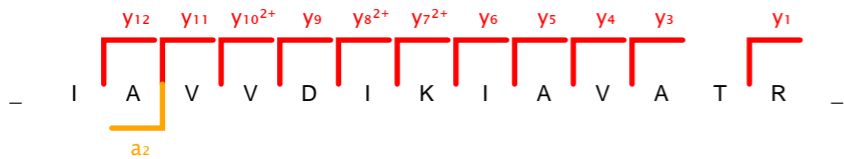
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 43241
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 134.59
Method: FTMS; HCD



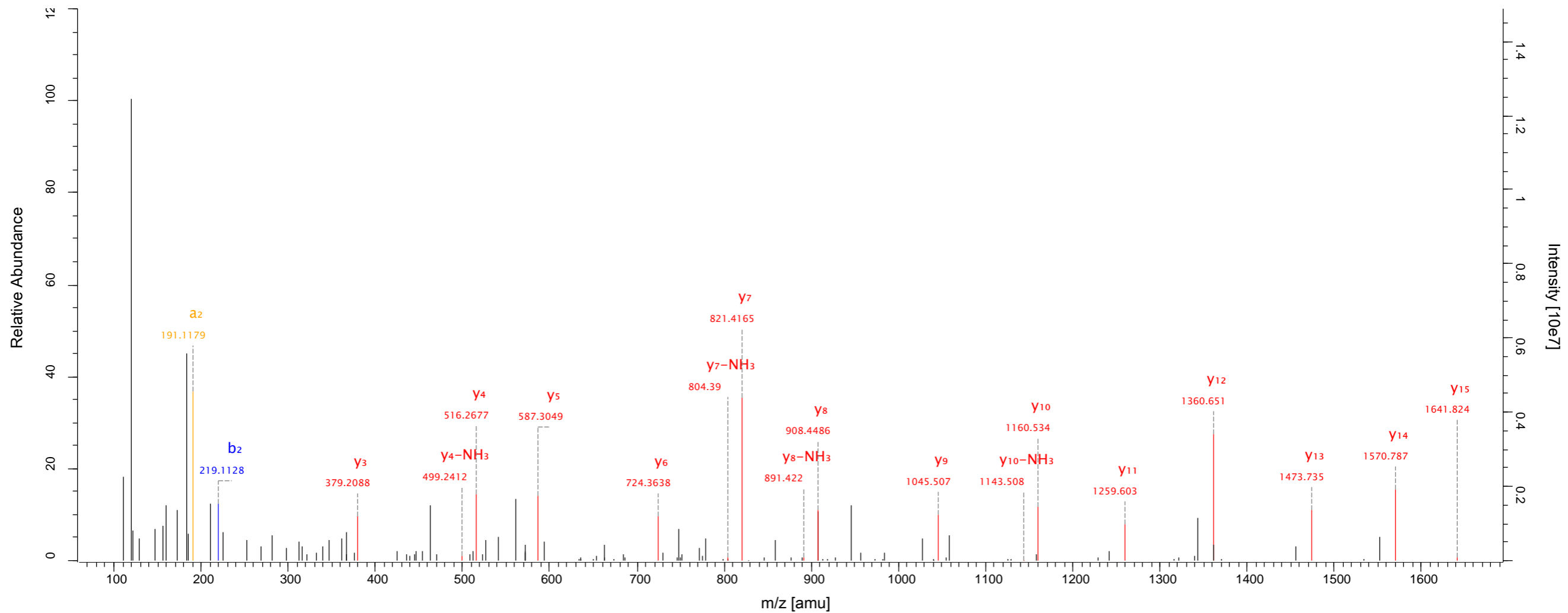
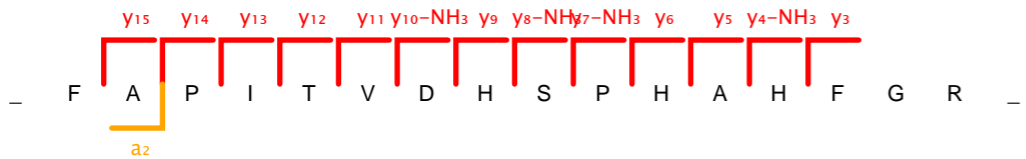
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 47239
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 84.48
Method: FTMS; HCD



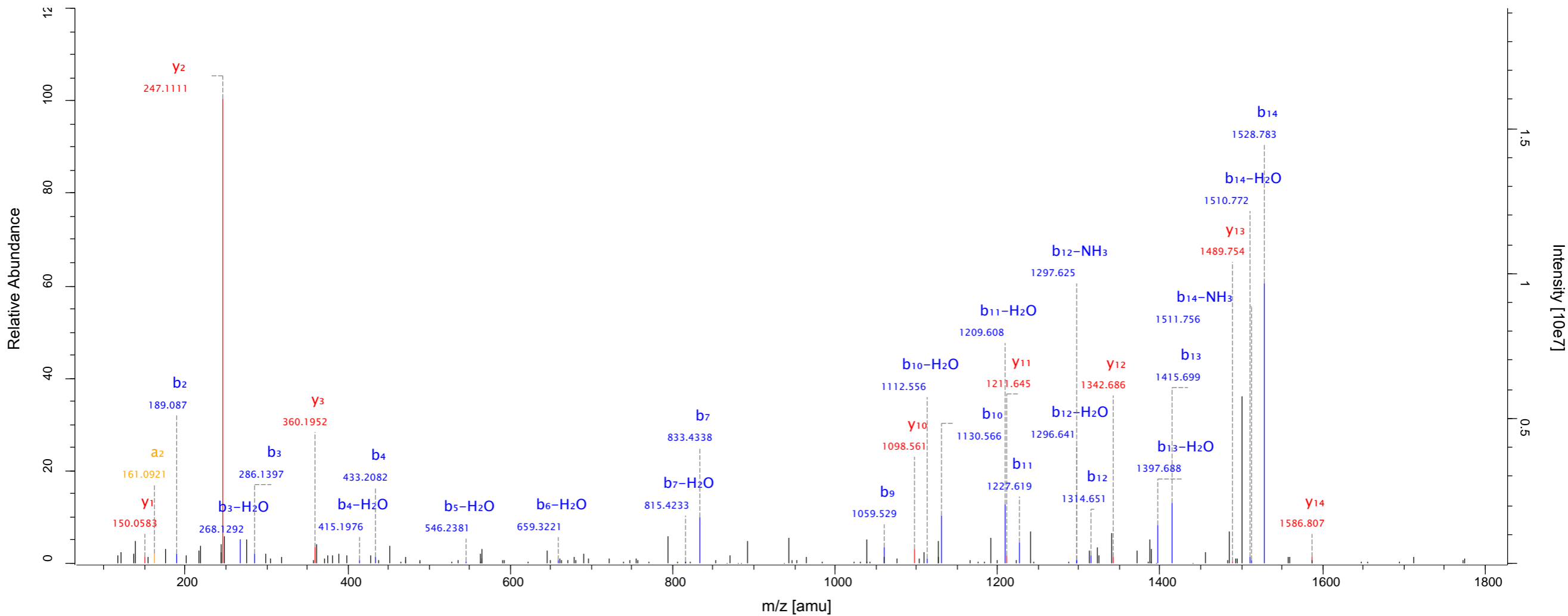
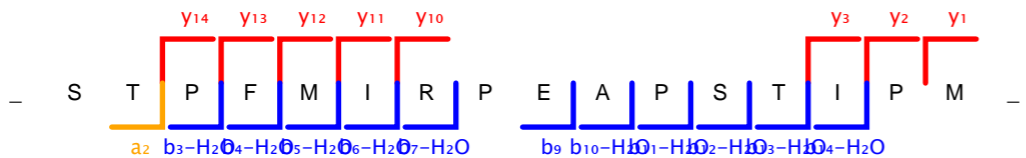
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 48108
Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
Peptide score: 68.44
Method: FTMS; HCD



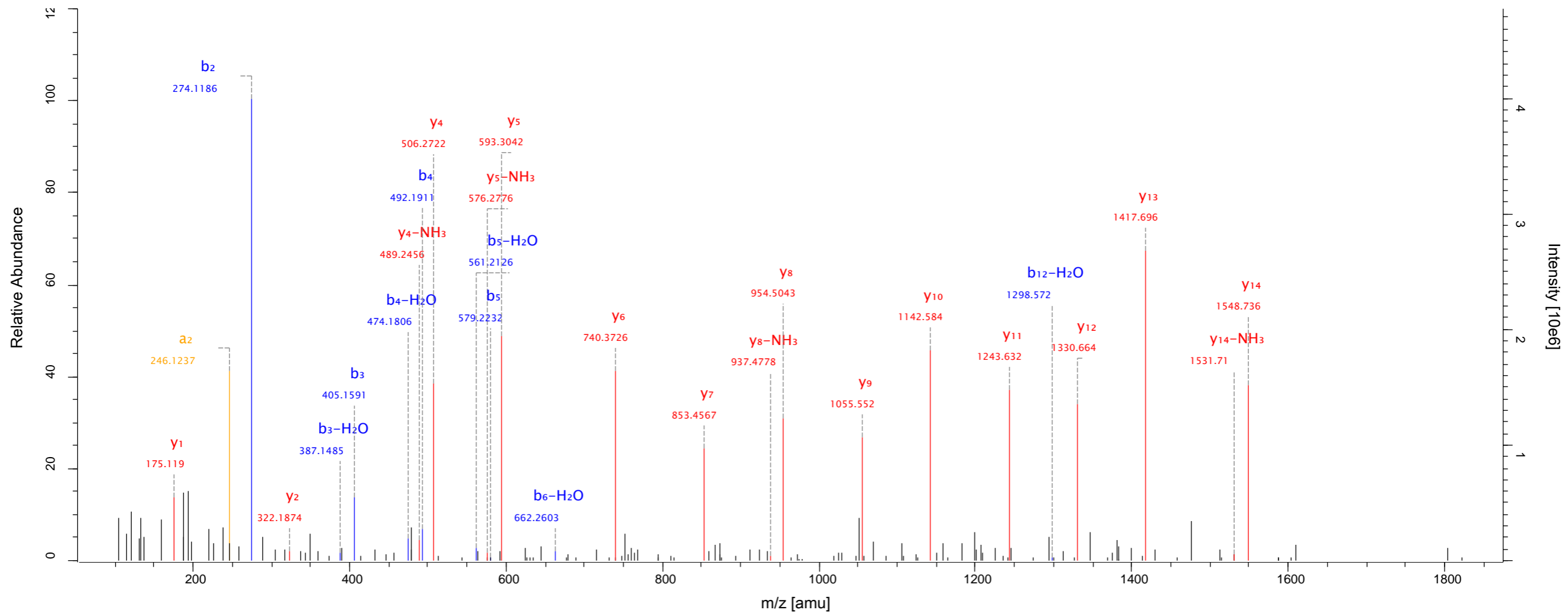
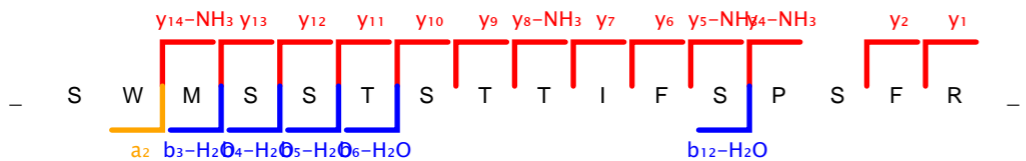
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 48331
 Protein: ORFL237C - NC_006273.2[149340,149699]- <unknown description>
 Peptide score: 106.13
 Method: FTMS; HCD



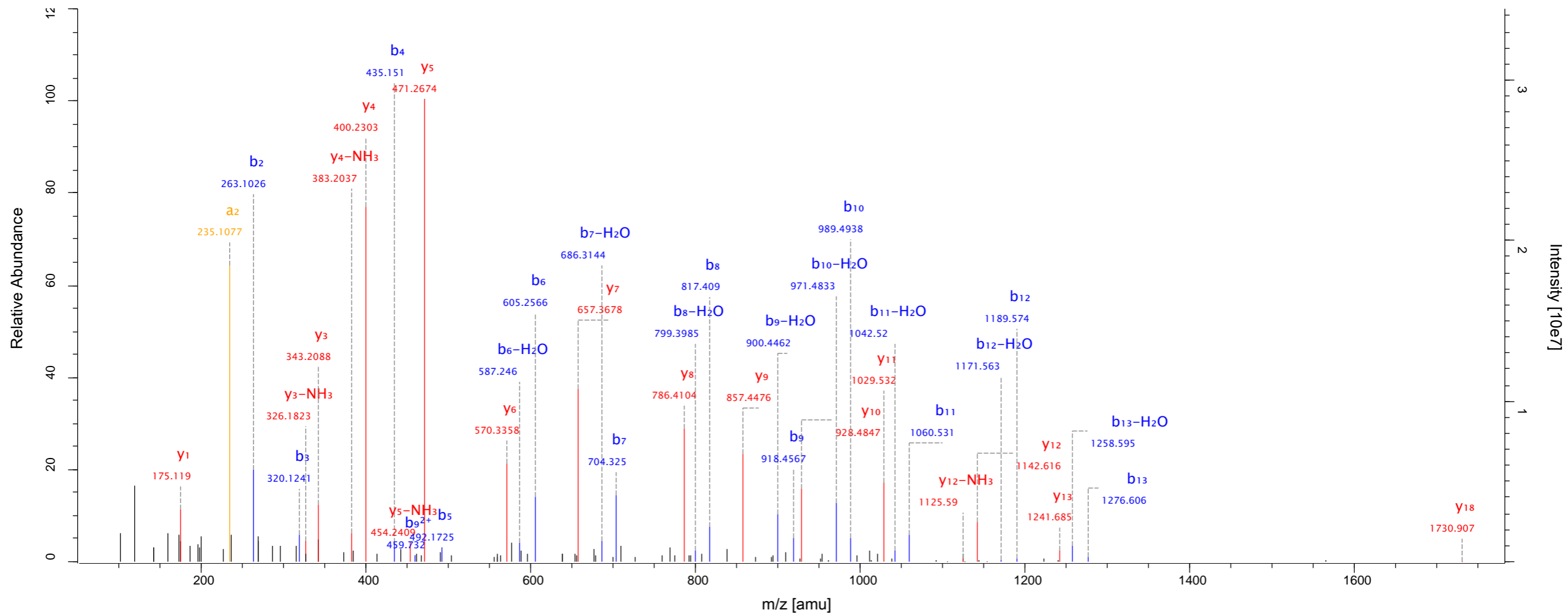
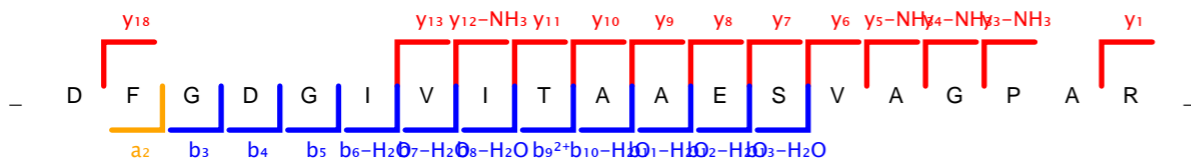
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 48783
 Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
 Peptide score: 117.77
 Method: FTMS; HCD



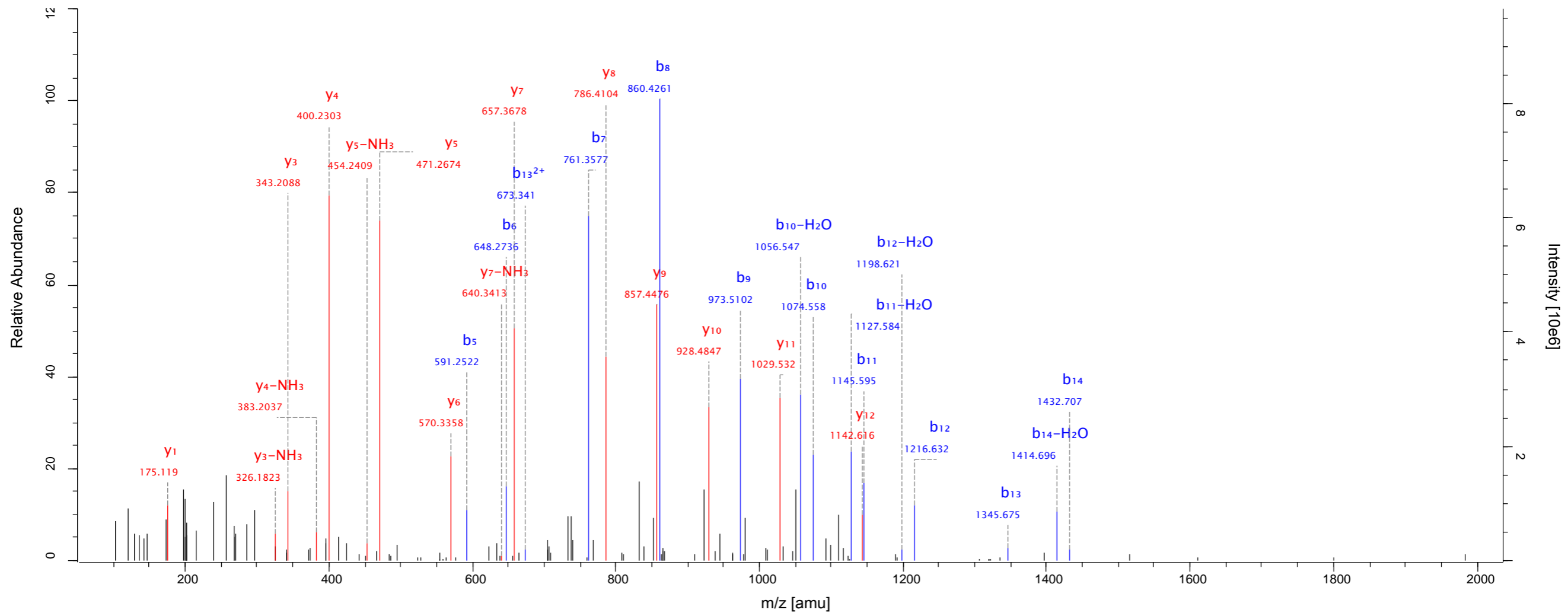
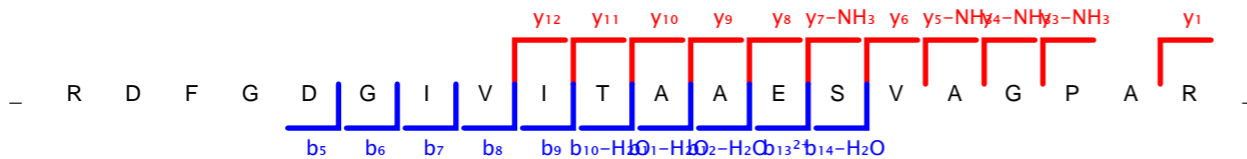
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 48877
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 113.72
 Method: FTMS; HCD



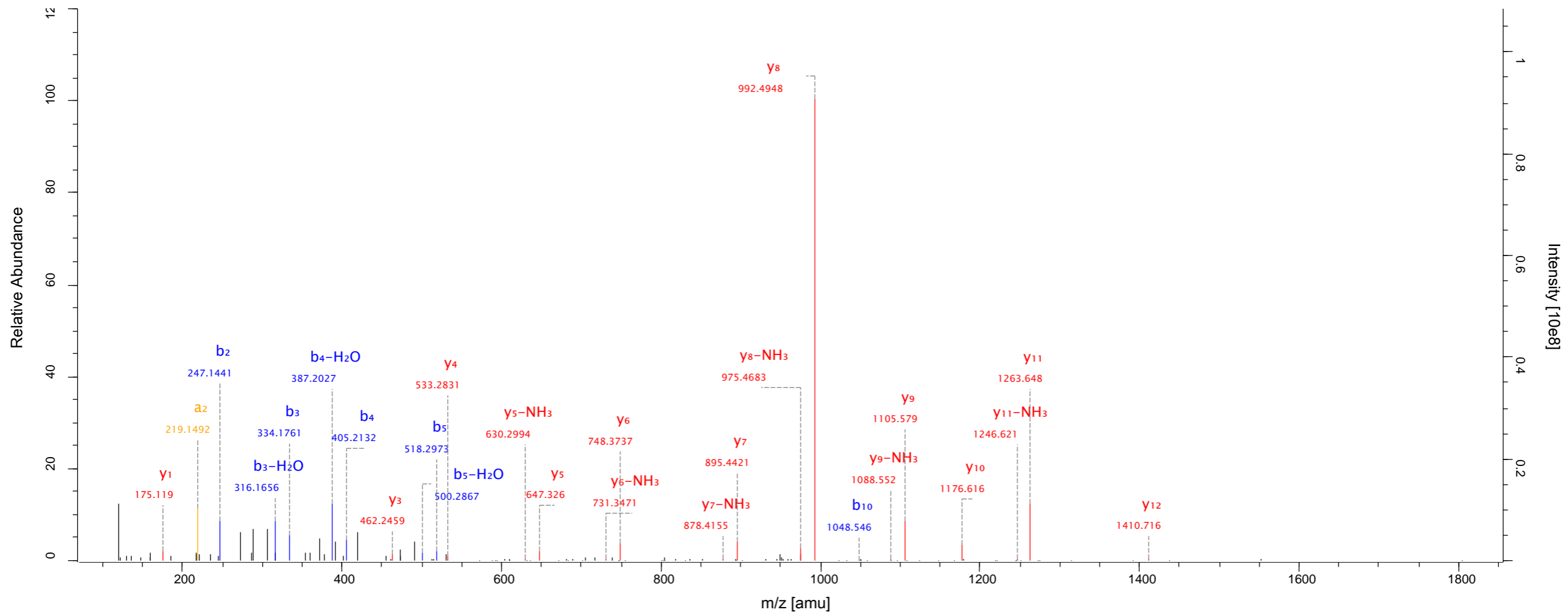
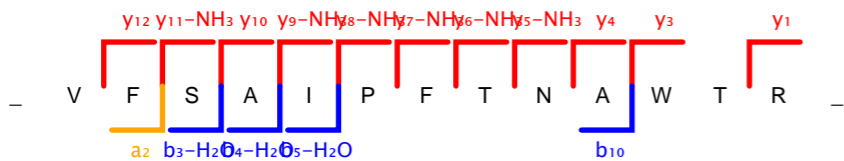
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 49026
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 269.26
 Method: FTMS; HCD



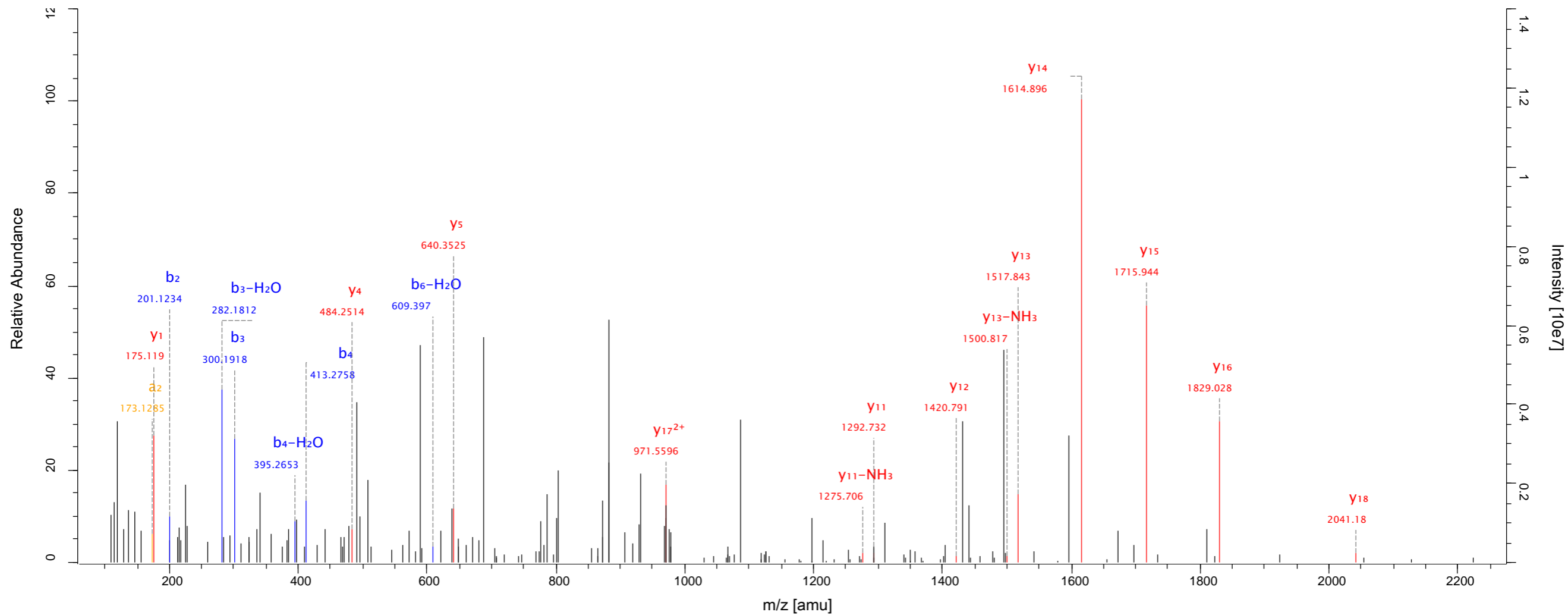
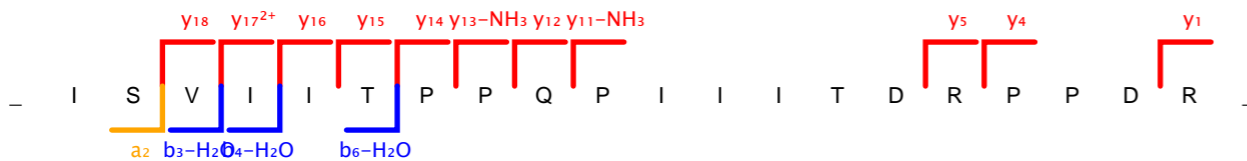
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
 Scan number: 49212
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 181.41
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
 Scan number: 49325
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 167.18
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 49357
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 62.94
Method: FTMS; HCD



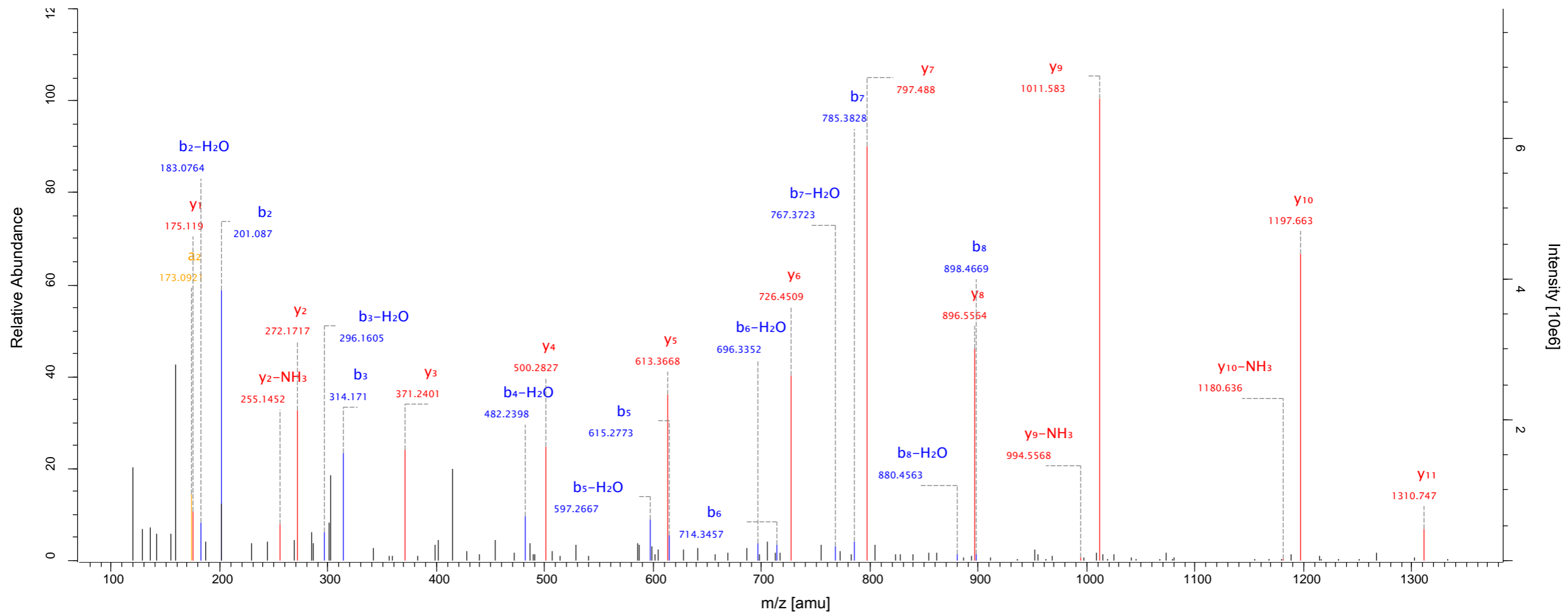
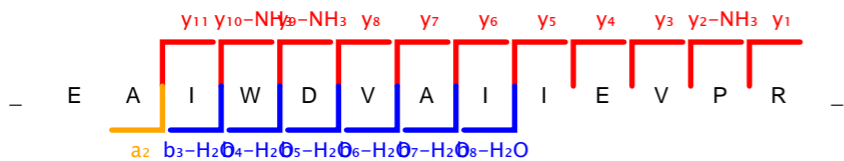
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27

Scan number: 49926

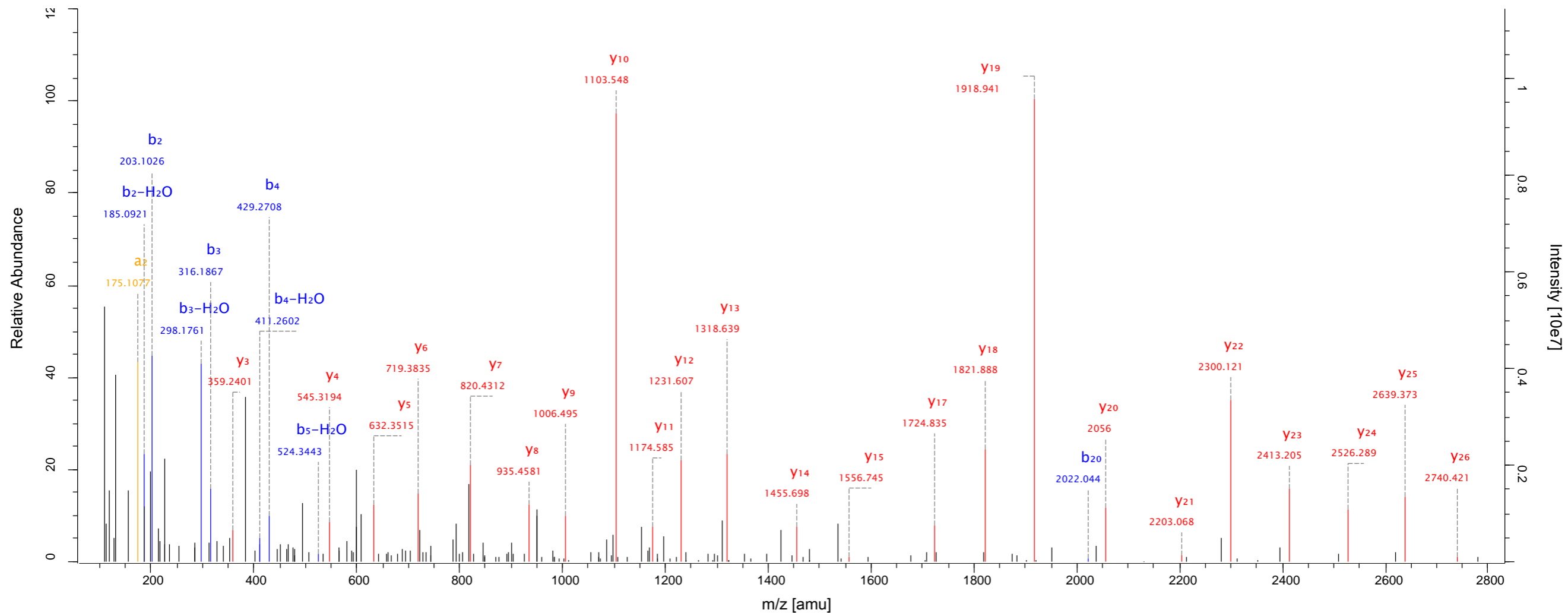
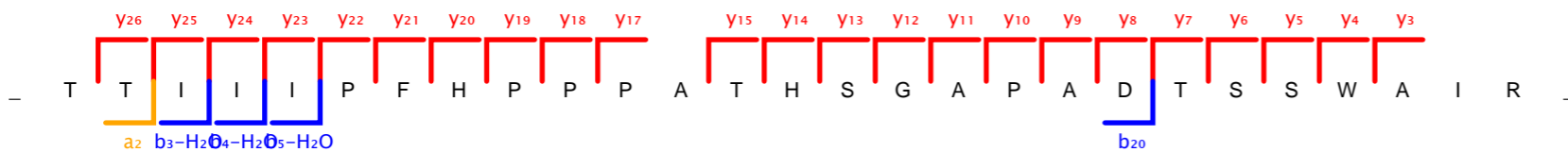
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 158.3

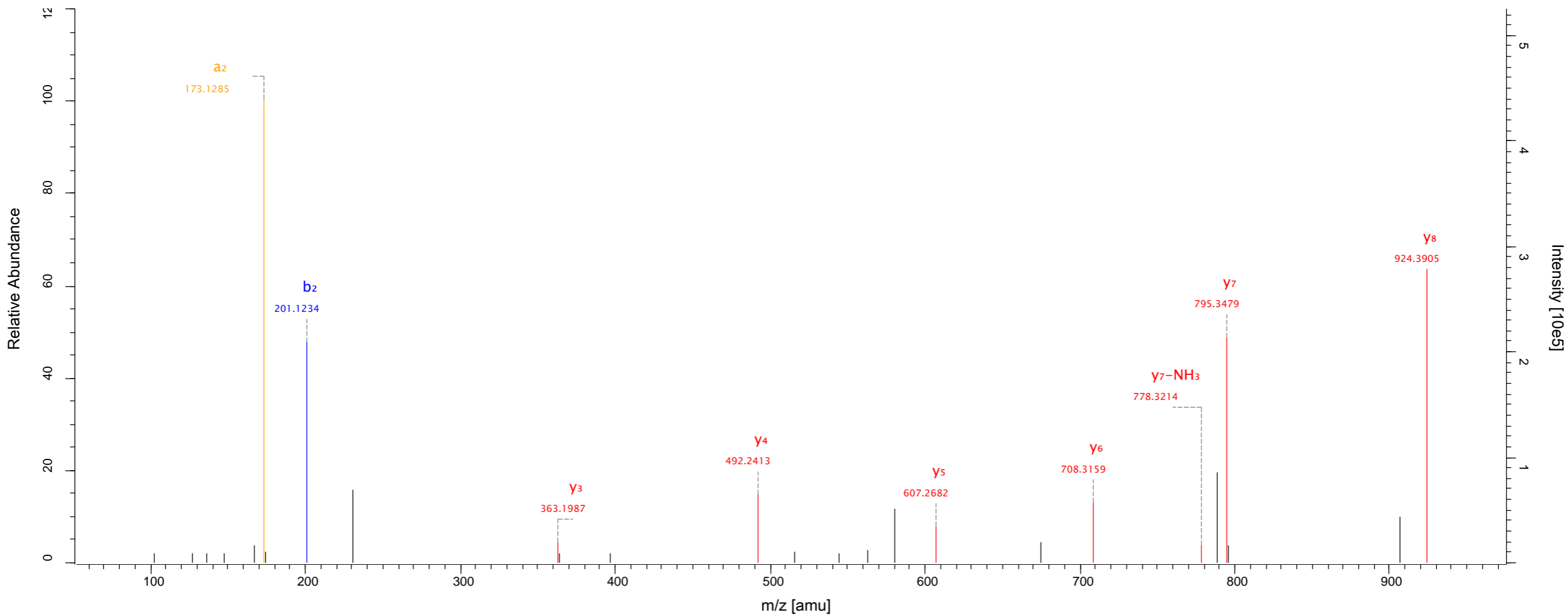
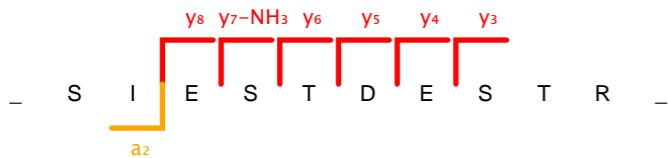
Method: FTMS; HCD



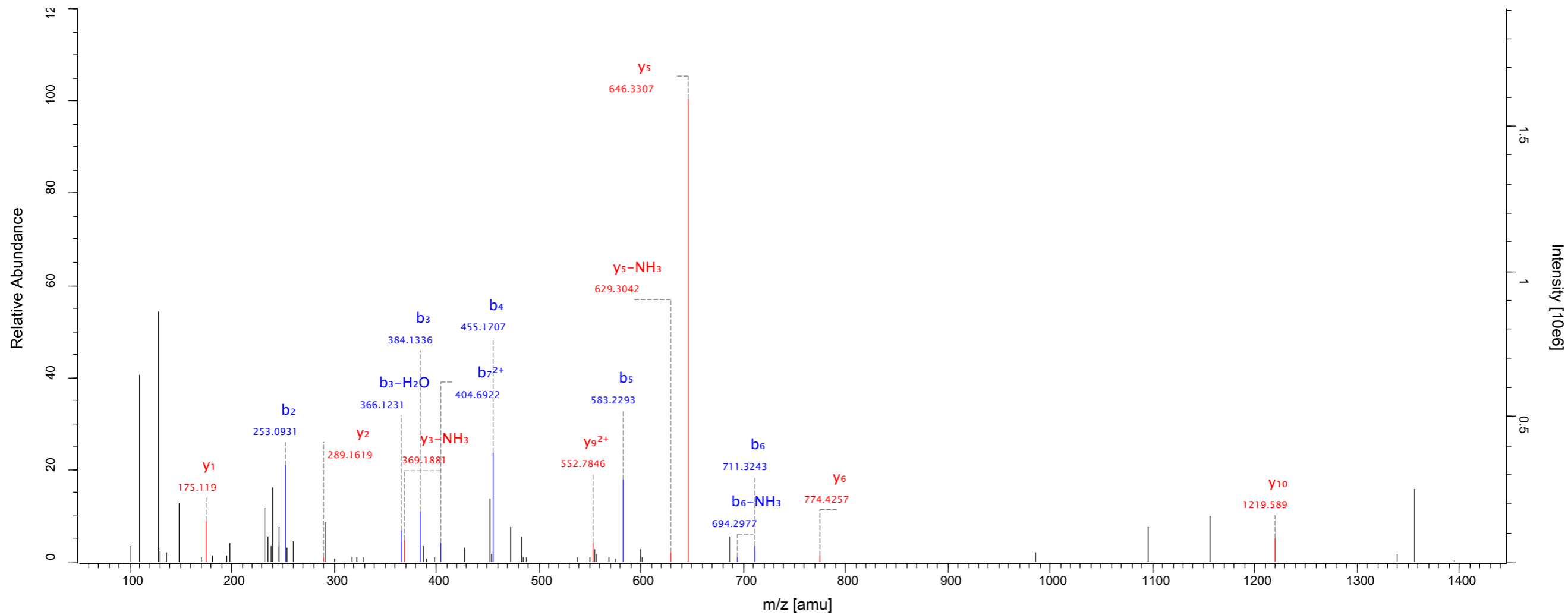
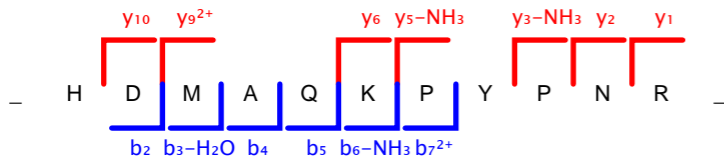
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 50380
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 195.39
Method: FTMS; HCD



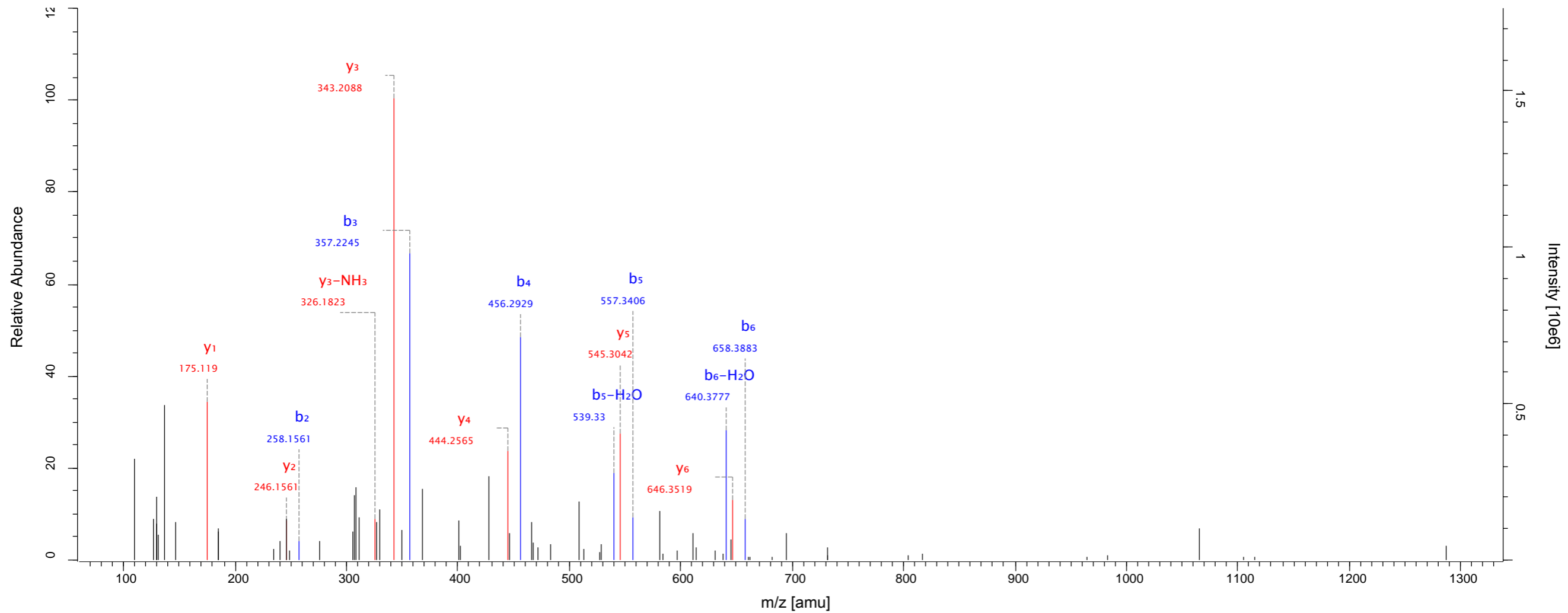
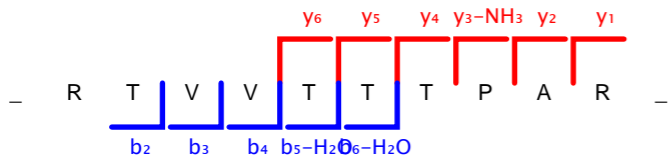
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 5988
Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
Peptide score: 66.43
Method: FTMS; HCD



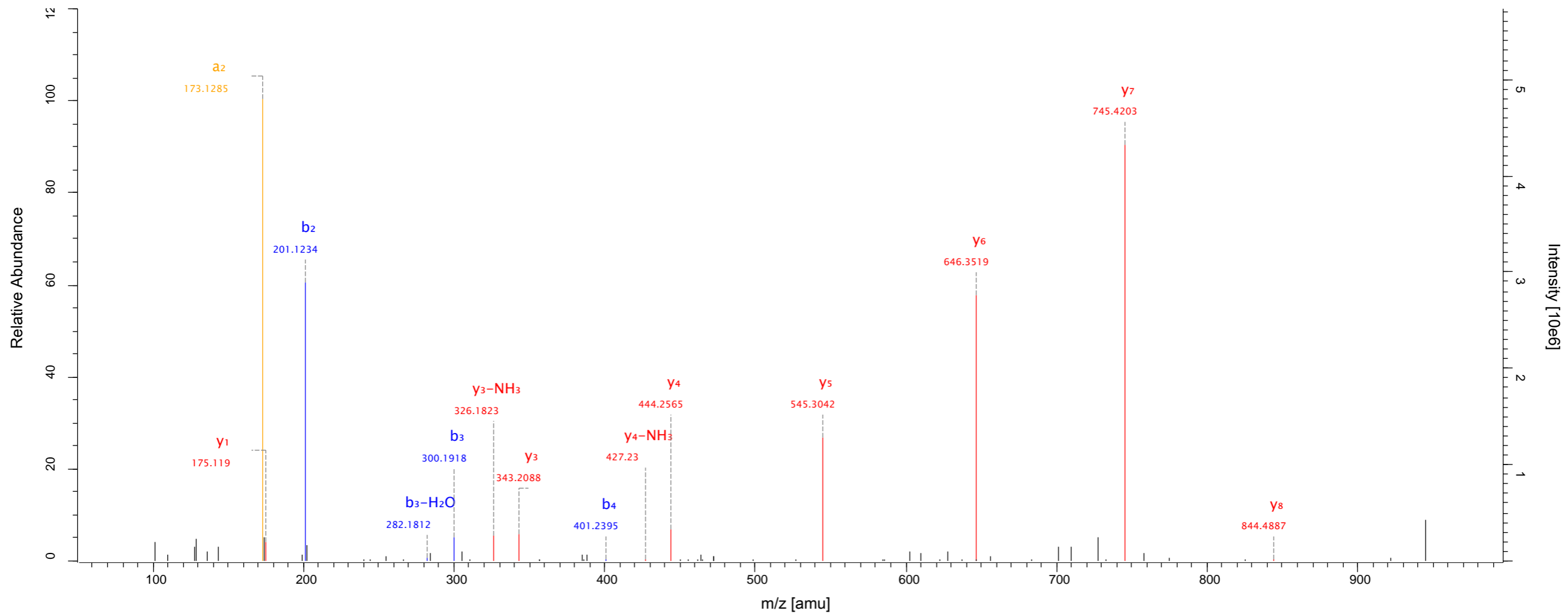
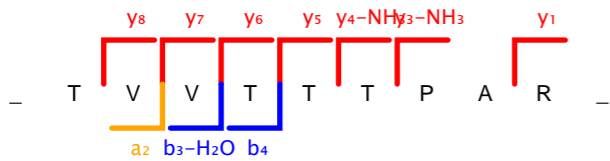
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 6073
Protein: ORFL185C - NC_006273.2[111134,111490]- <unknown description>
Peptide score: 82.42
Method: FTMS; HCD



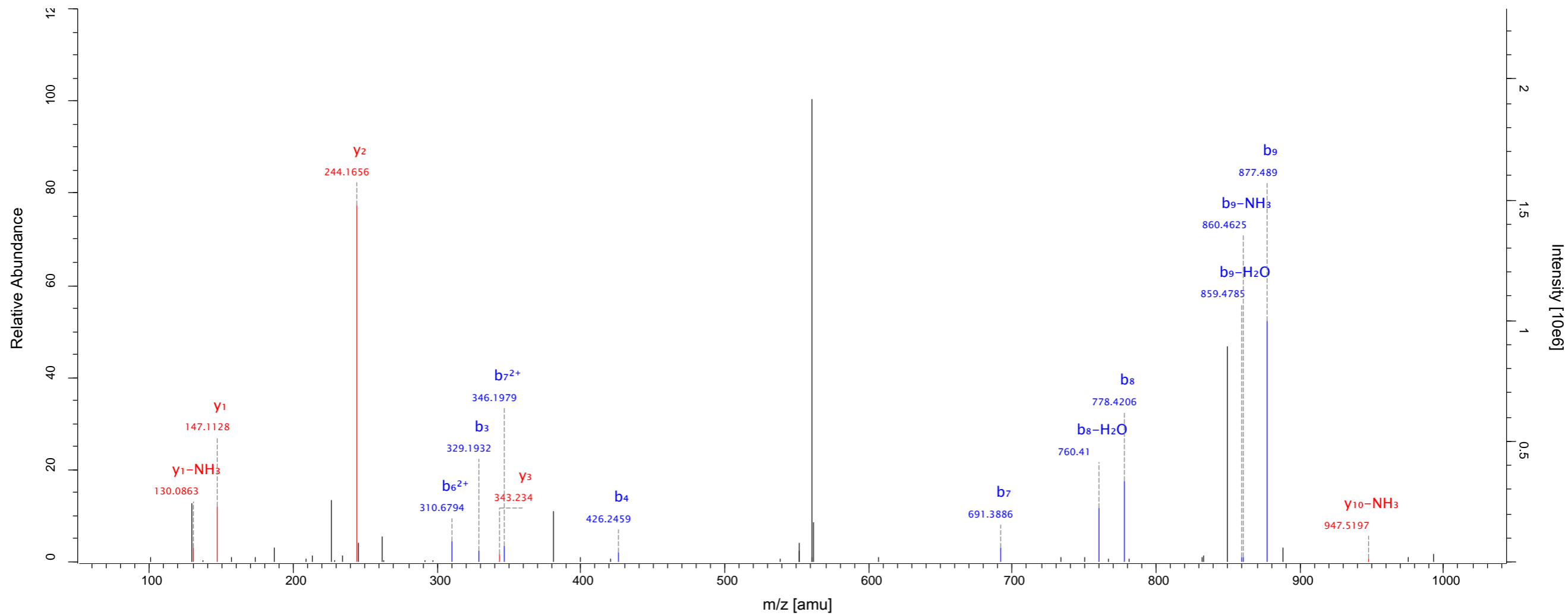
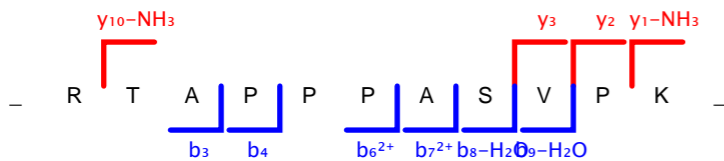
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 6920
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 80.76
Method: FTMS; HCD



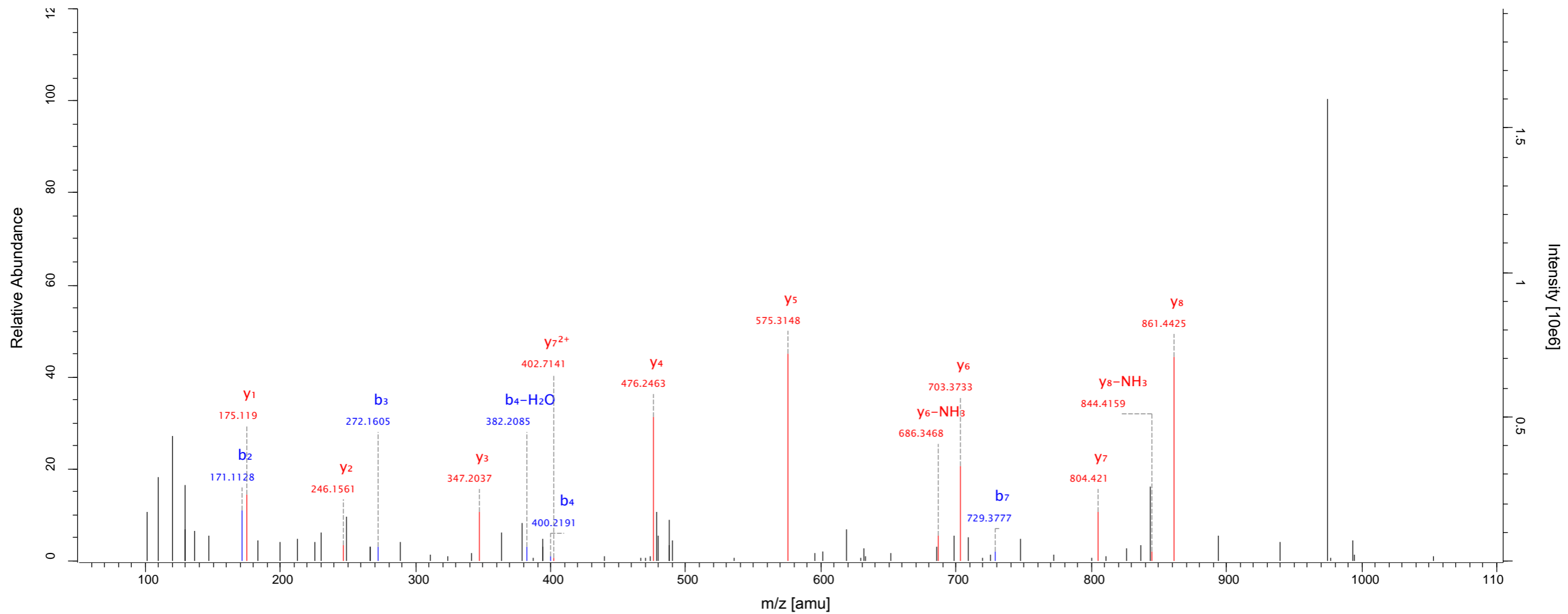
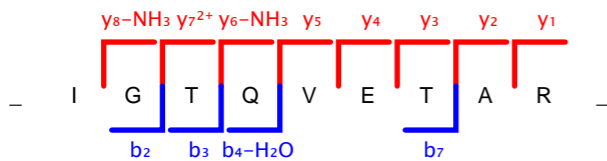
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 8015
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 80.24
Method: FTMS; HCD



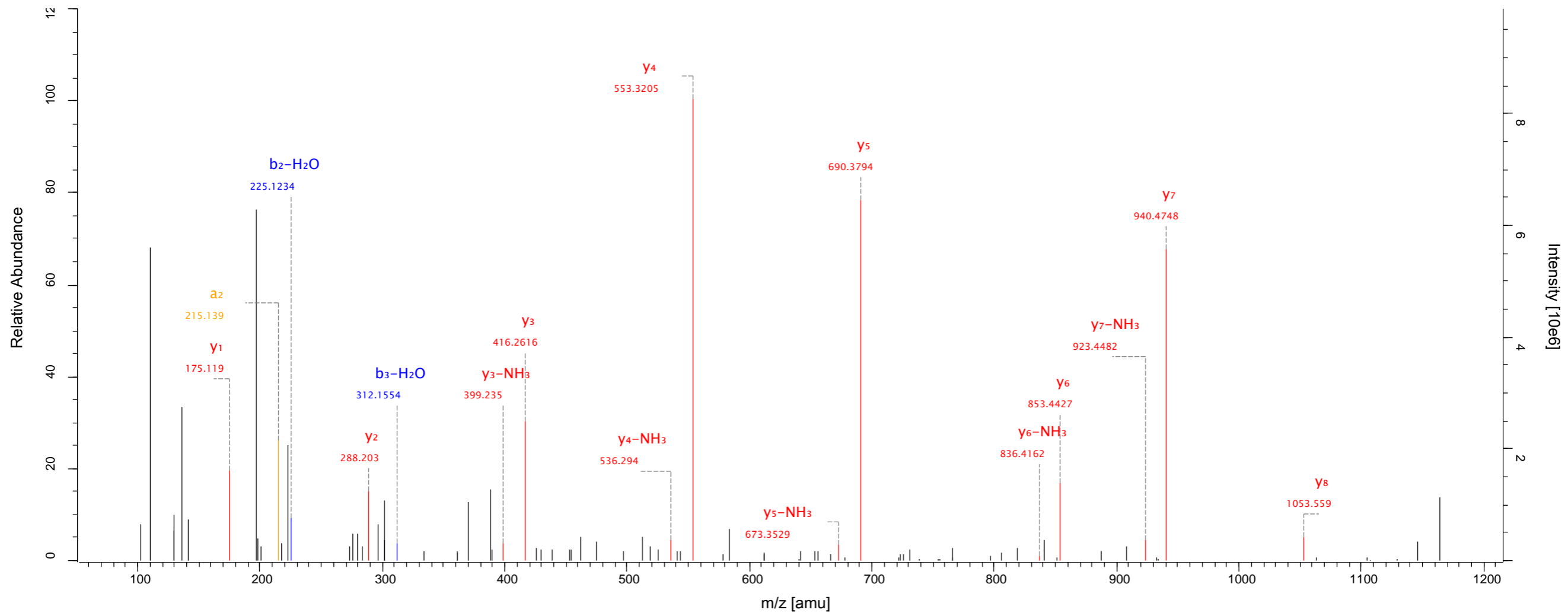
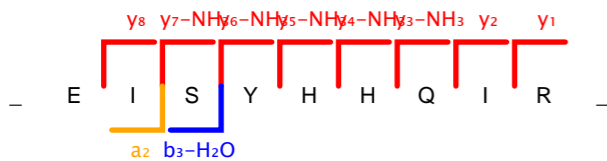
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 8072
Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
Peptide score: 73.78
Method: FTMS; HCD



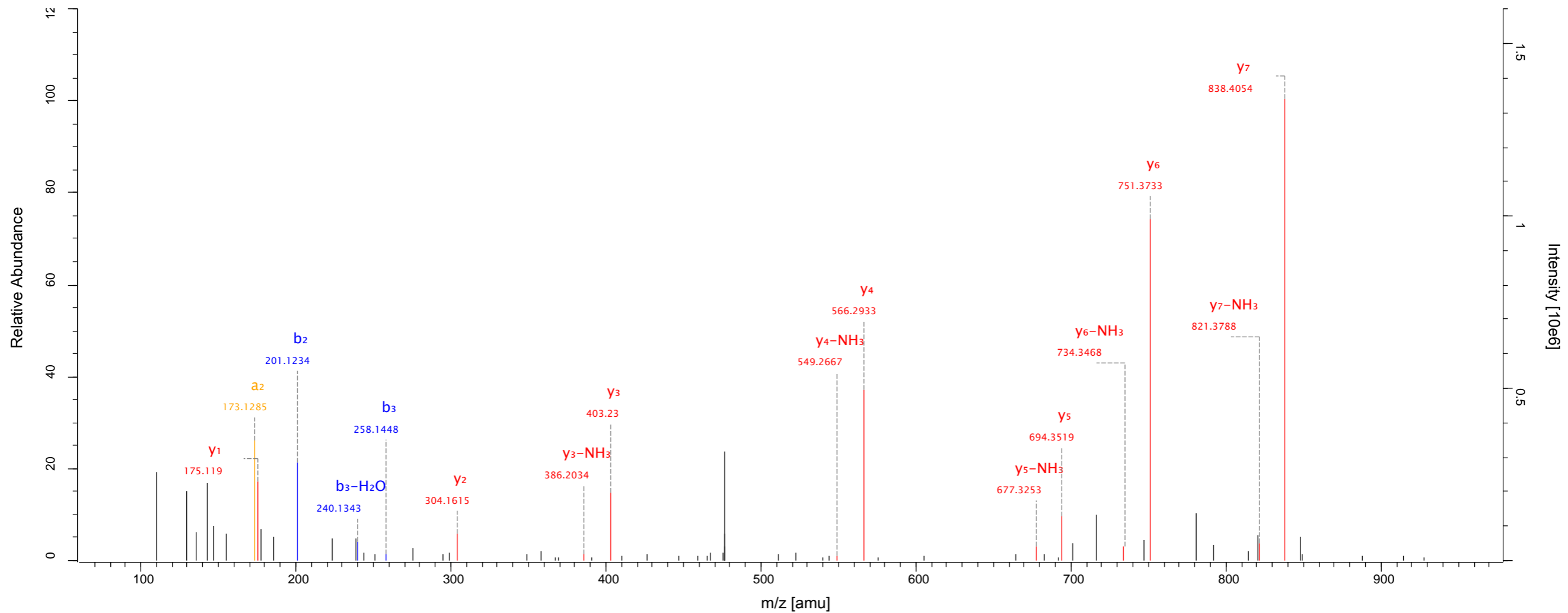
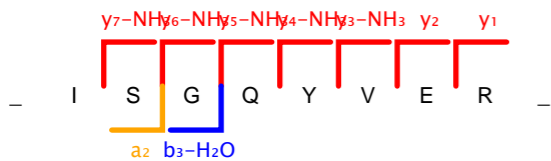
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 8216
Protein: ORFL190C - NC_006273.2[111965,112384]- <unknown description>
Peptide score: 84
Method: FTMS; HCD



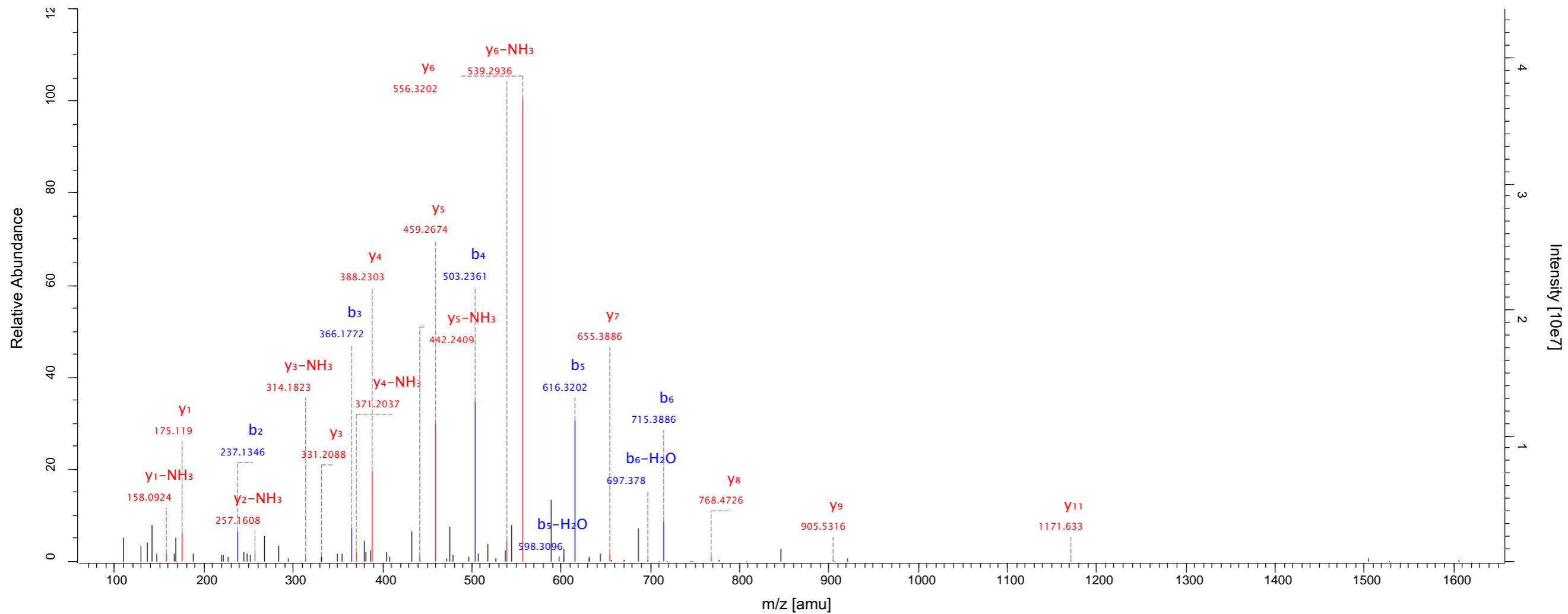
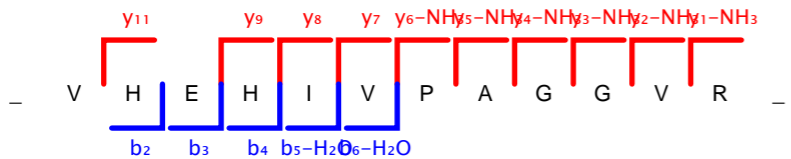
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 8923
Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
Peptide score: 96.71
Method: FTMS; HCD



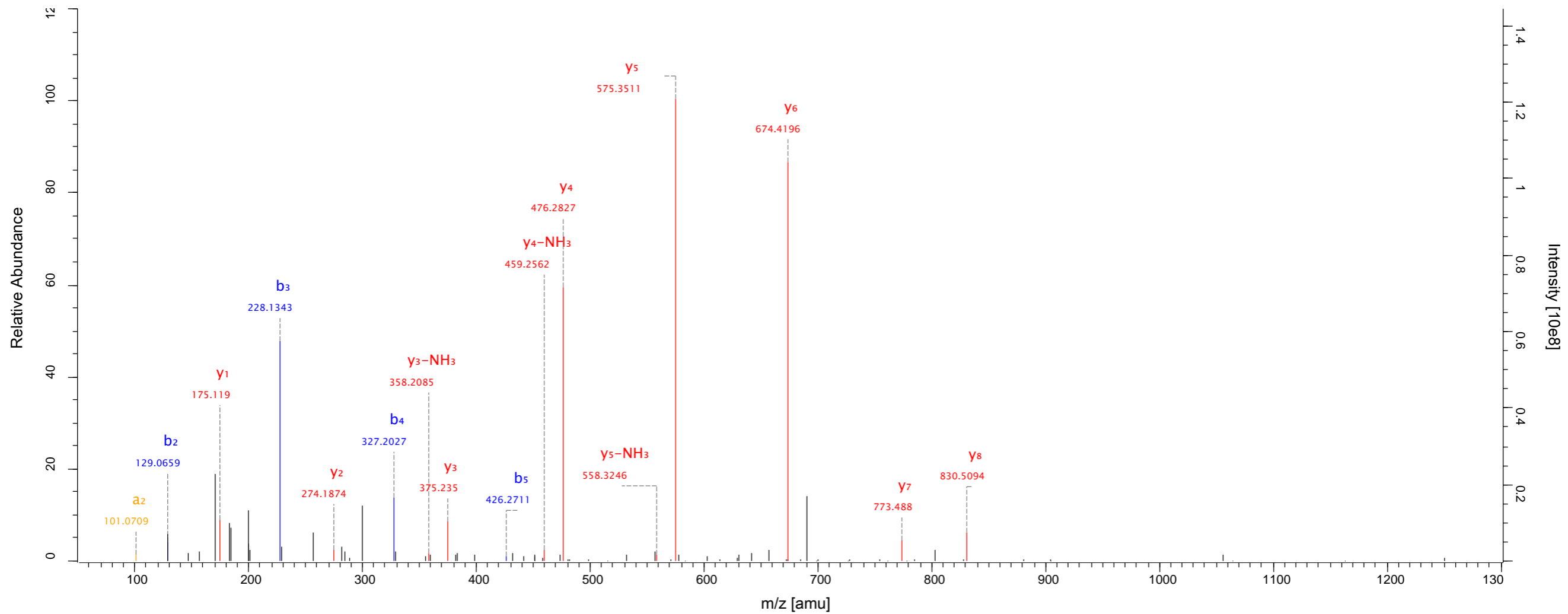
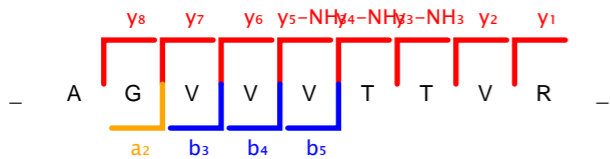
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 9323
Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
Peptide score: 101.43
Method: FTMS; HCD



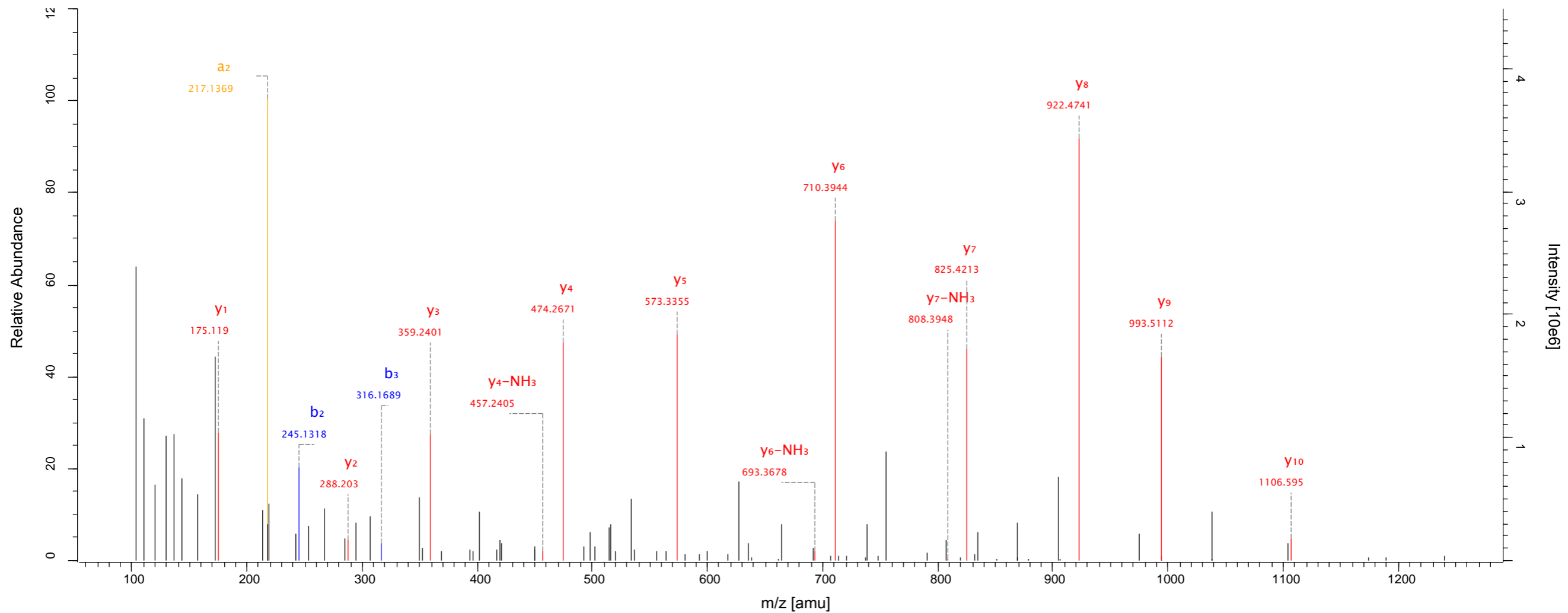
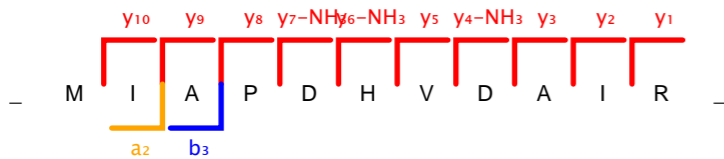
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_29
 Scan number: 11409
 Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
 Peptide score: 108.31
 Method: FTMS; HCD



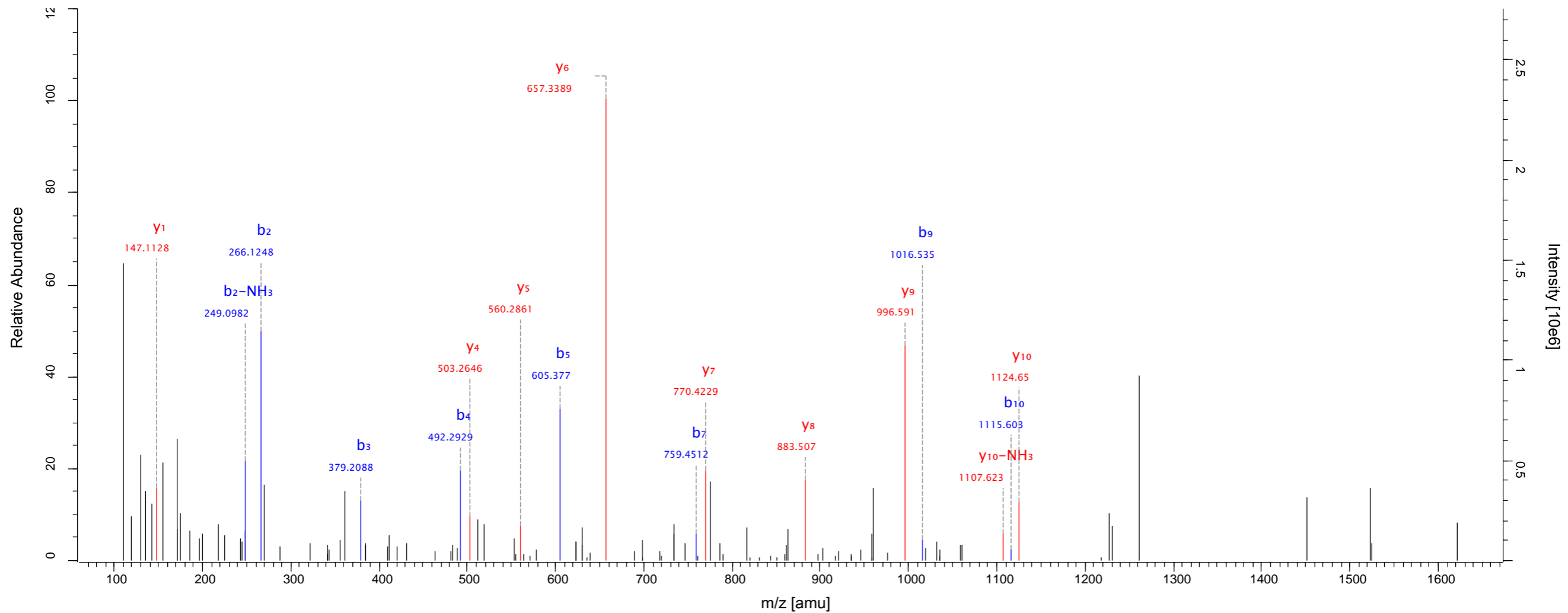
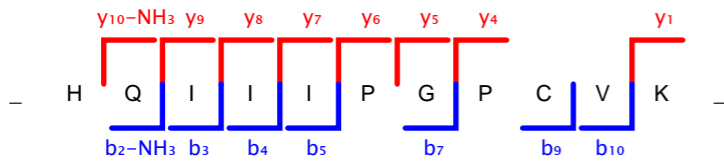
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_29
Scan number: 14263
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 105.14
Method: FTMS; HCD



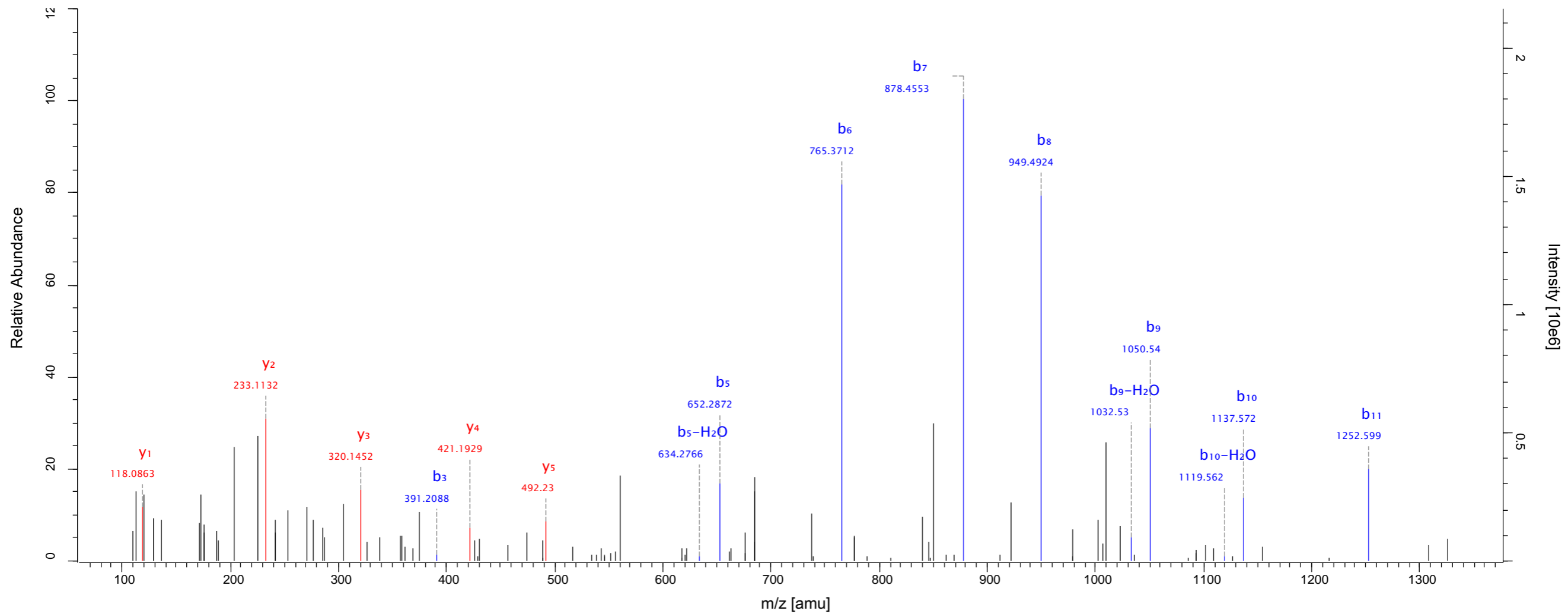
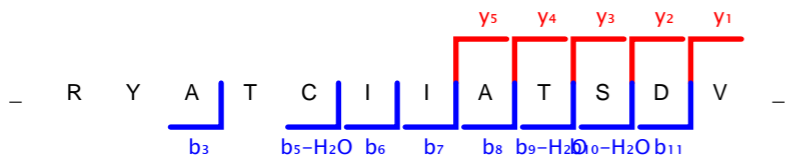
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_29
Scan number: 21970
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 83
Method: FTMS; HCD



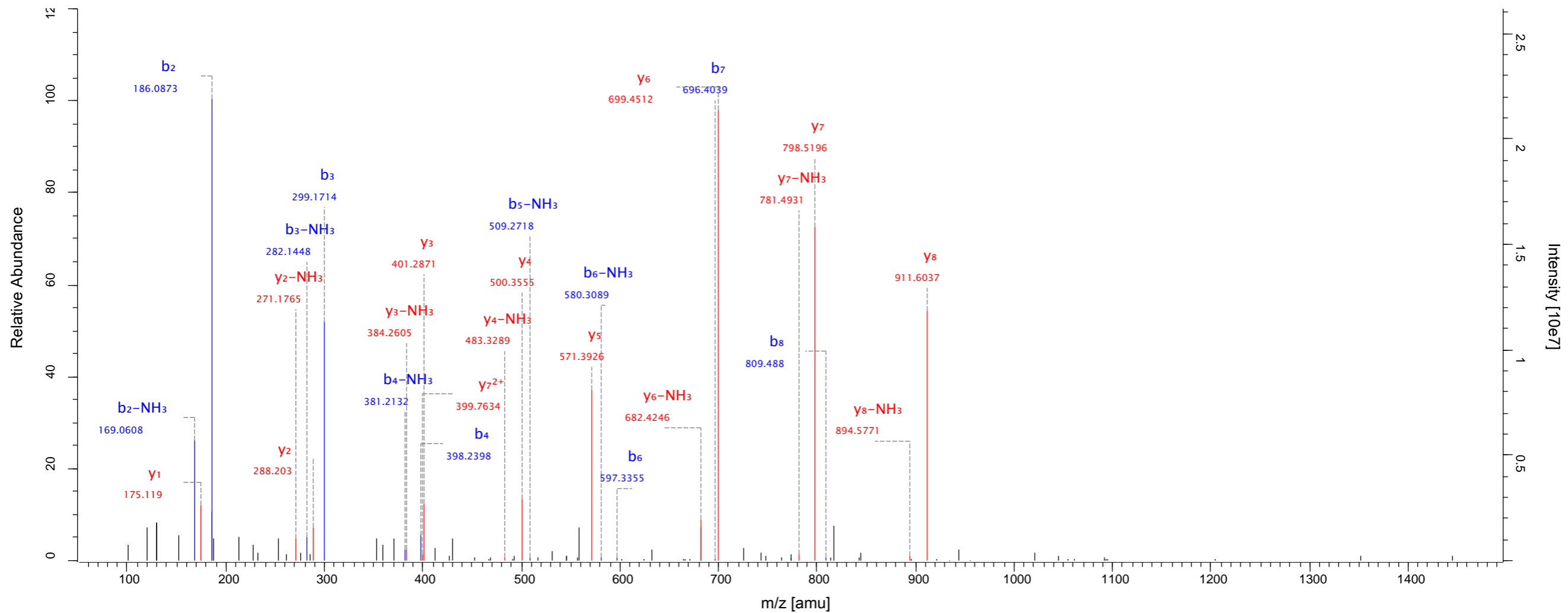
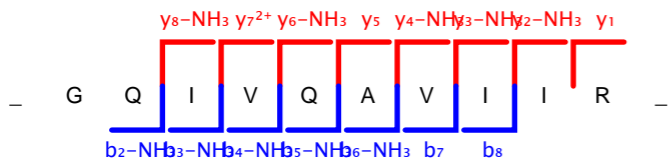
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_29
Scan number: 25809
Protein: ORFL253W.IORF1 + NC_006273.2[161917,163748]+ <unknown description>; ORFL251W + NC_006273.2[161872,163748]+ <unknown description>
Peptide score: 114.83
Method: FTMS; HCD



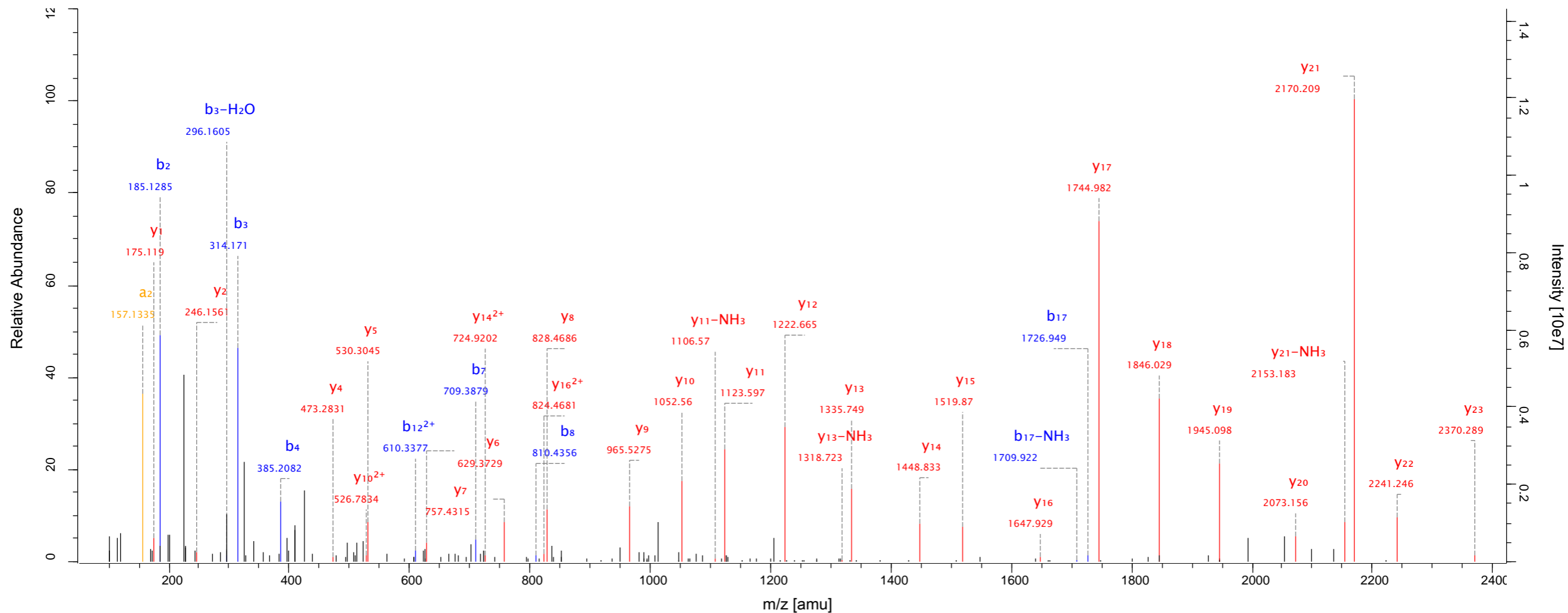
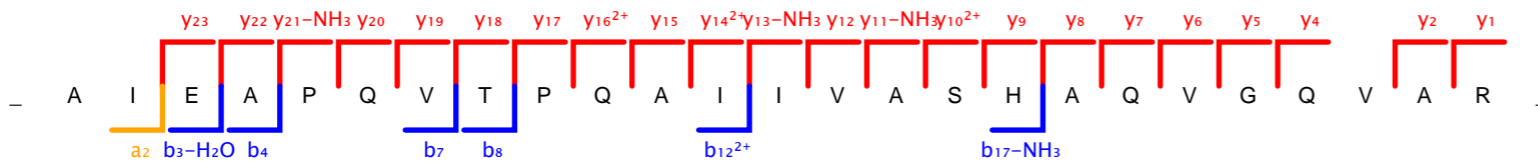
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_29
Scan number: 30479
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 85.29
Method: FTMS; HCD



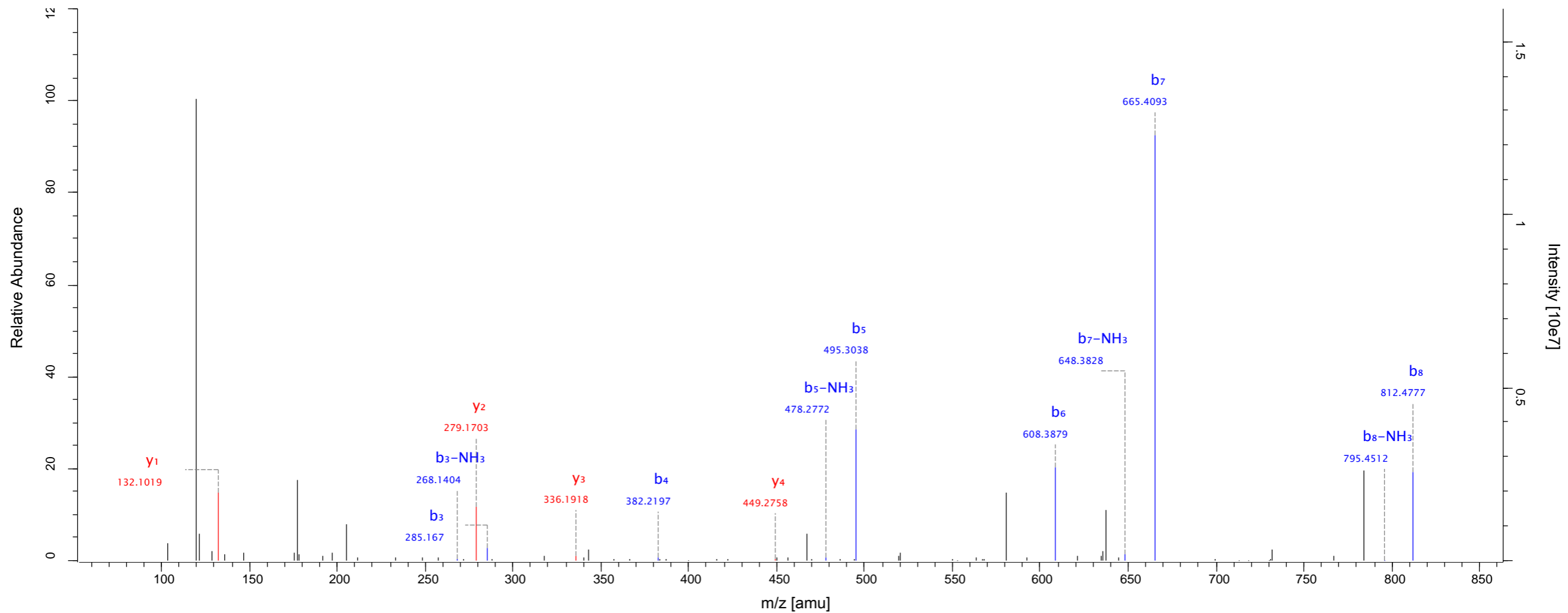
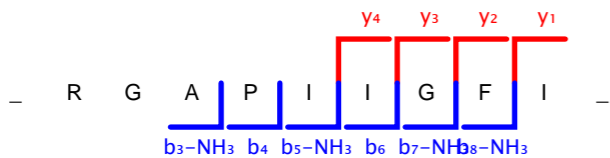
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_29
 Scan number: 37461
 Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
 Peptide score: 150.53
 Method: FTMS; HCD



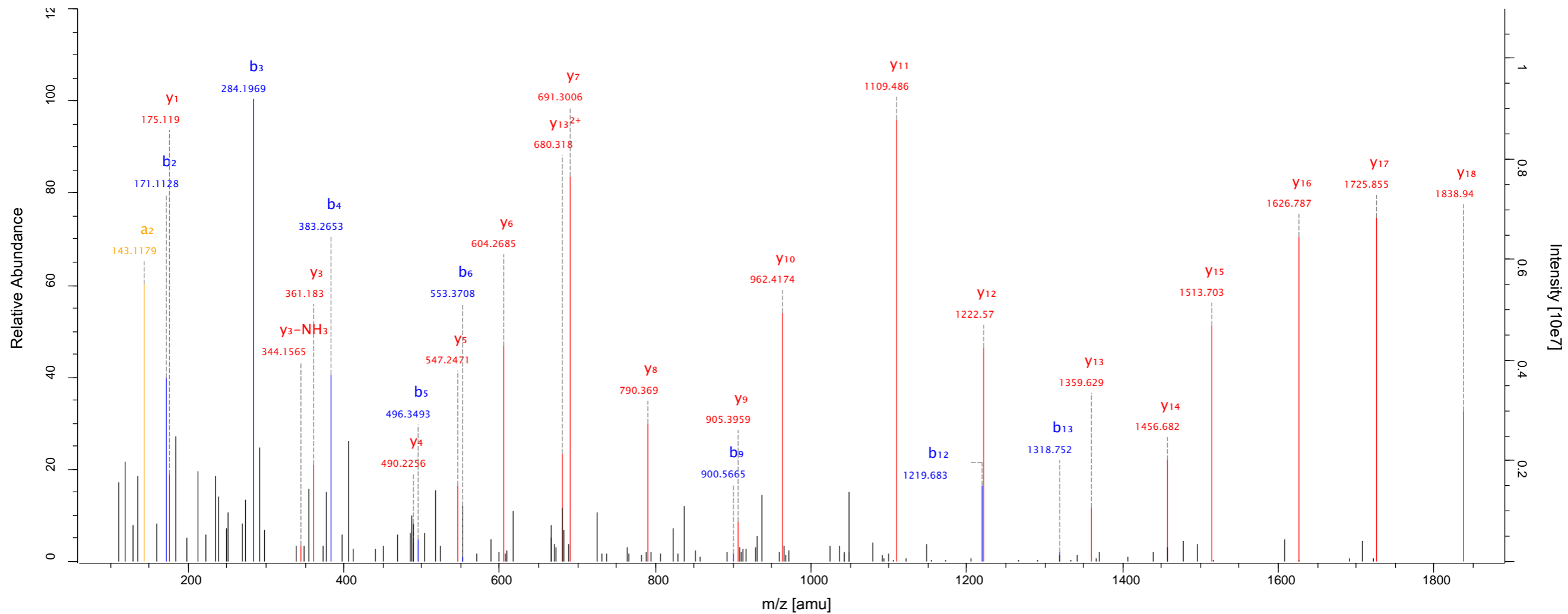
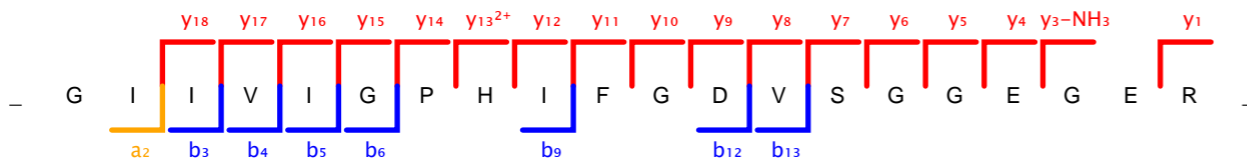
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_29
 Scan number: 53671
 Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
 Peptide score: 189.79
 Method: FTMS; HCD



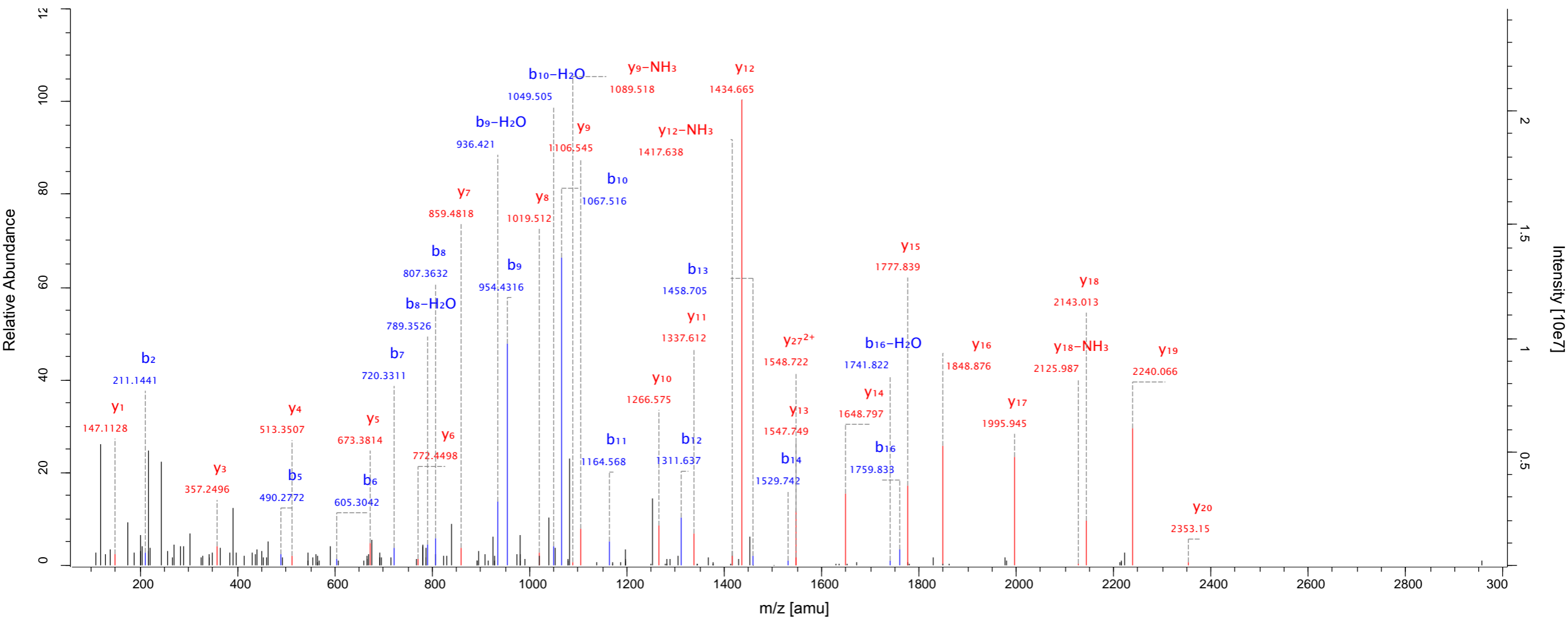
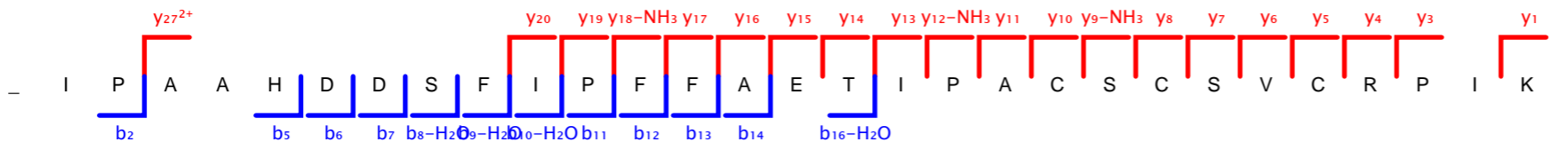
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_29
Scan number: 53735
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 78.77
Method: FTMS; HCD



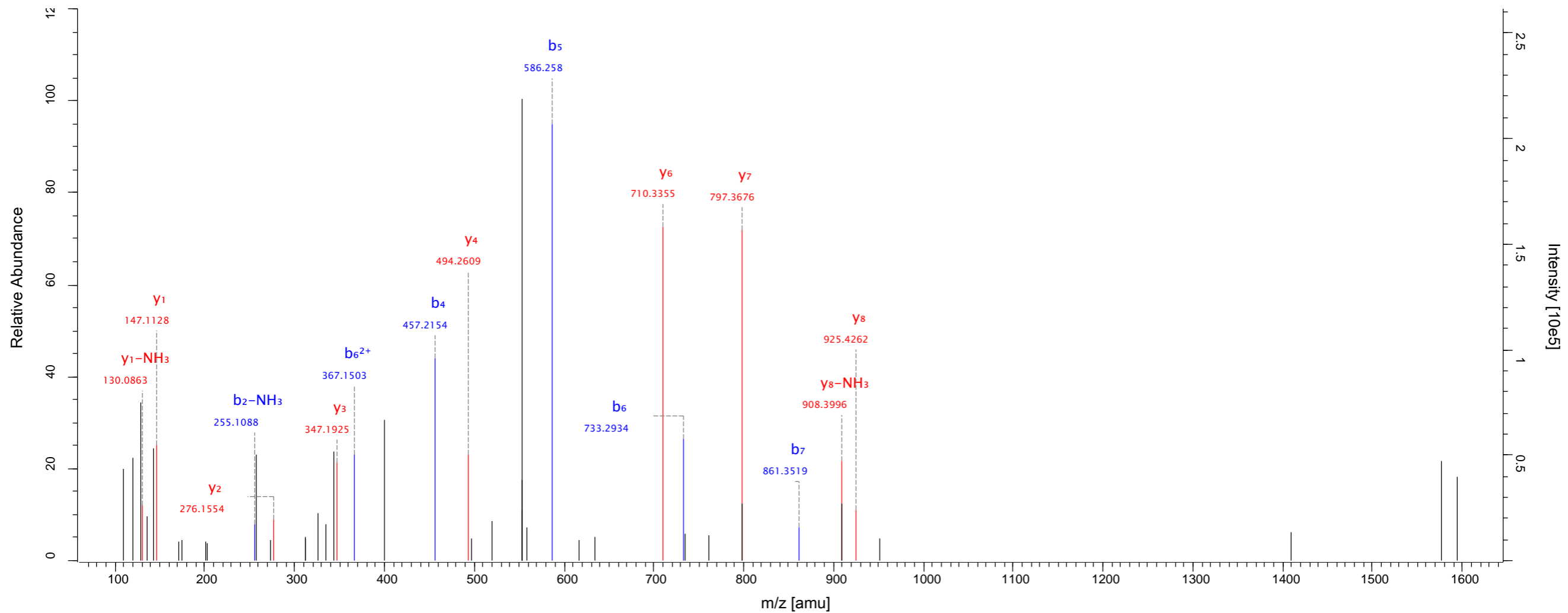
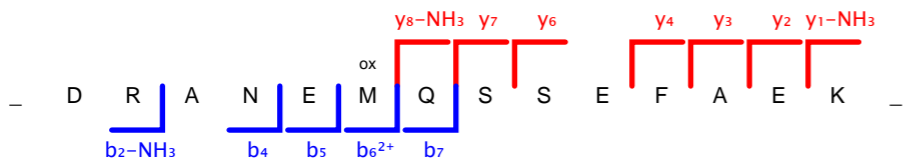
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_29
Scan number: 55174
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 165.75
Method: FTMS; HCD



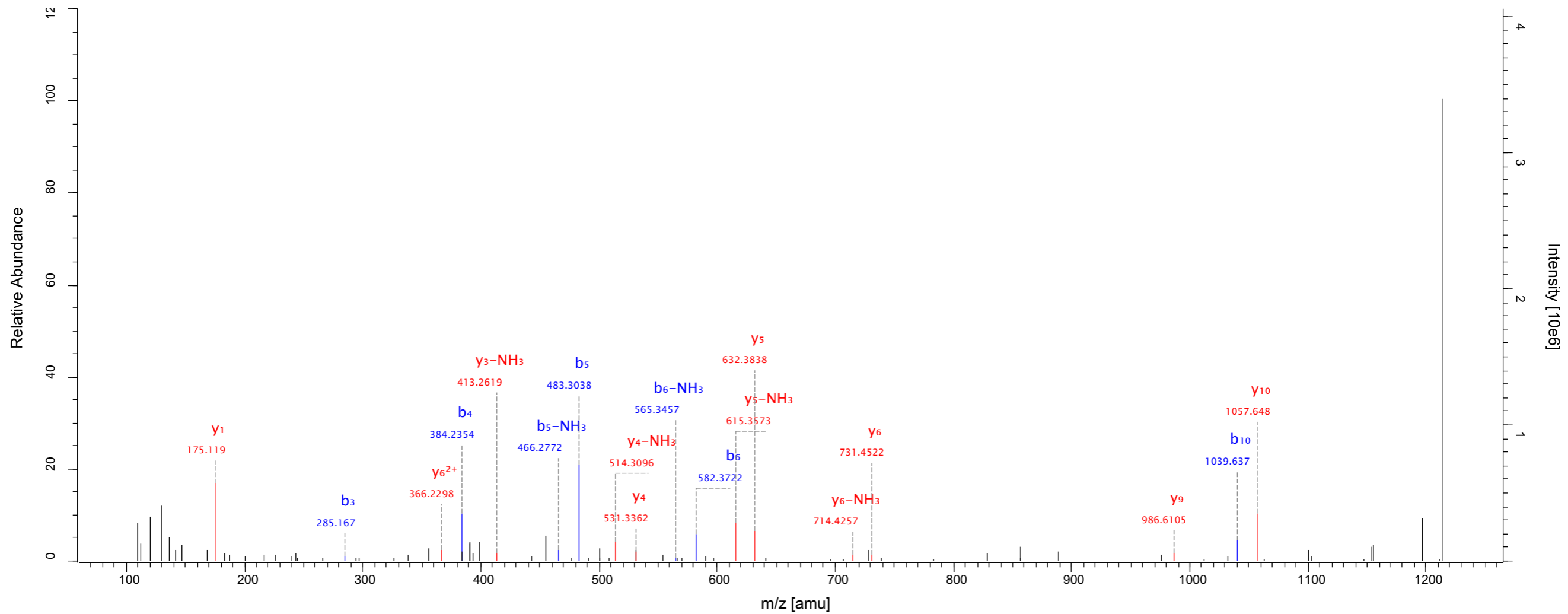
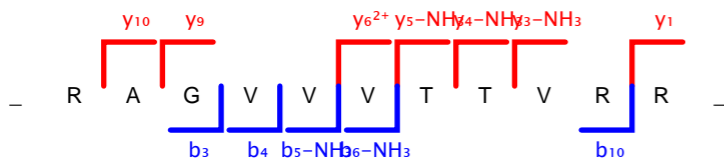
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_29
 Scan number: 55615
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 167.99
 Method: FTMS; HCD



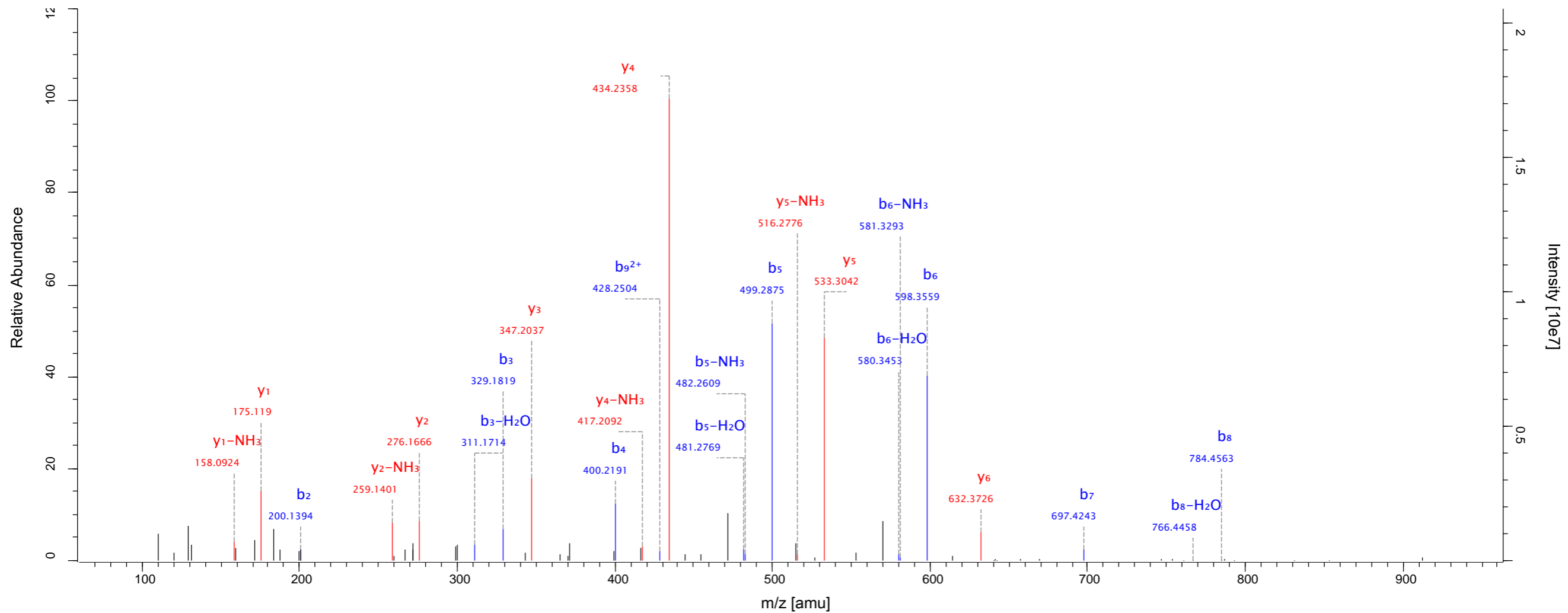
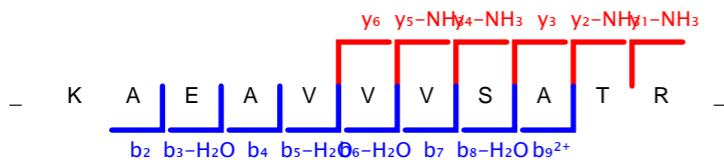
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_29
Scan number: 7963
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 79.47
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_29
Scan number: 9374
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 97.46
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_29
Scan number: 9573
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 133.91
Method: FTMS; HCD



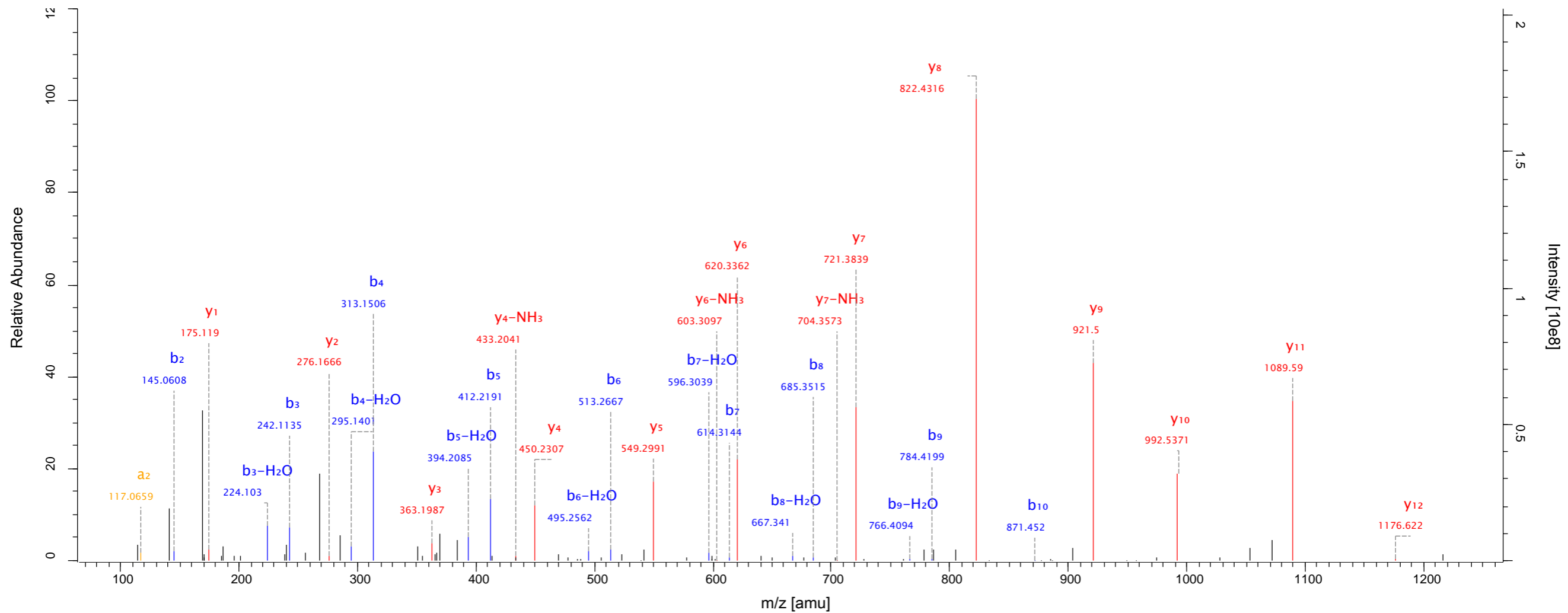
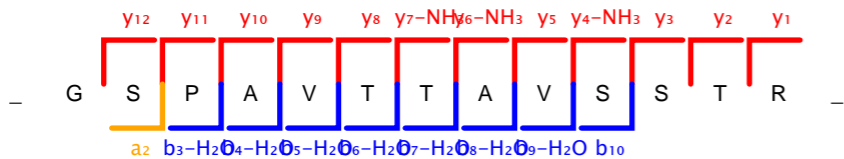
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_30

Scan number: 13072

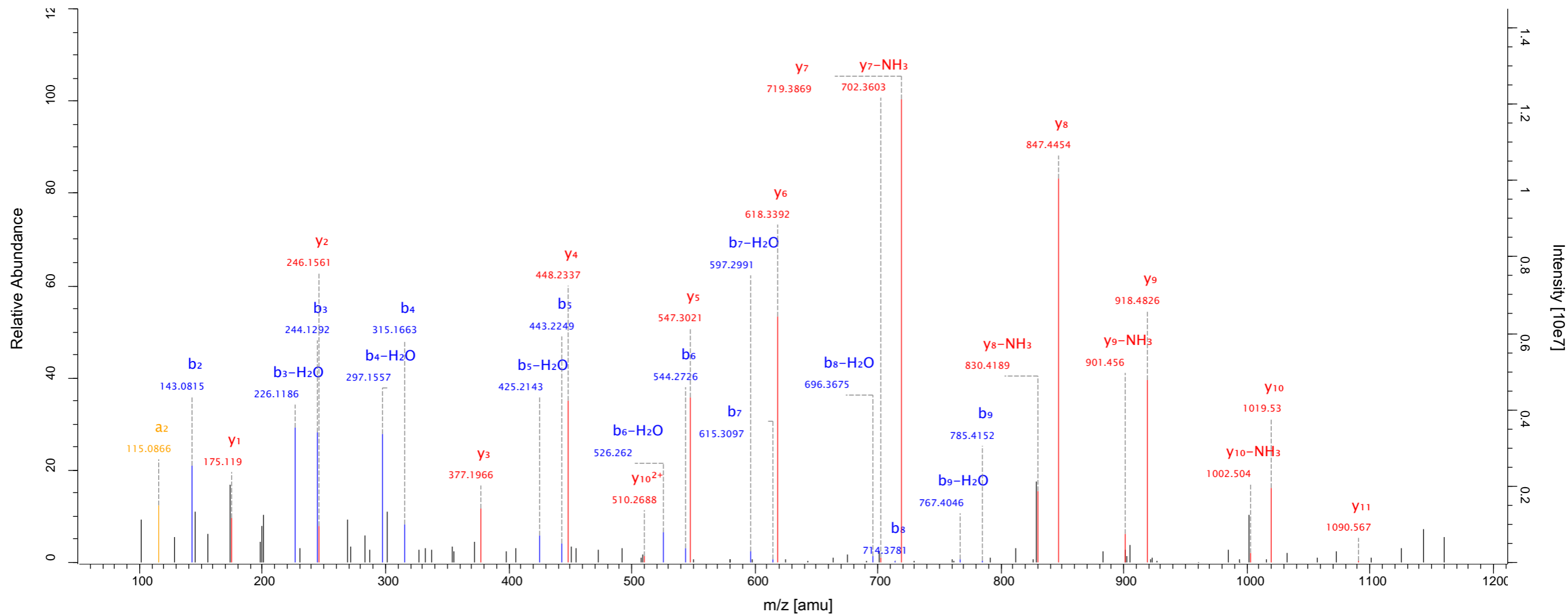
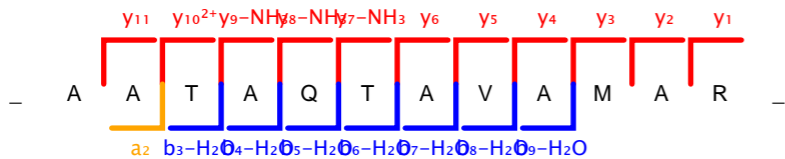
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 187.58

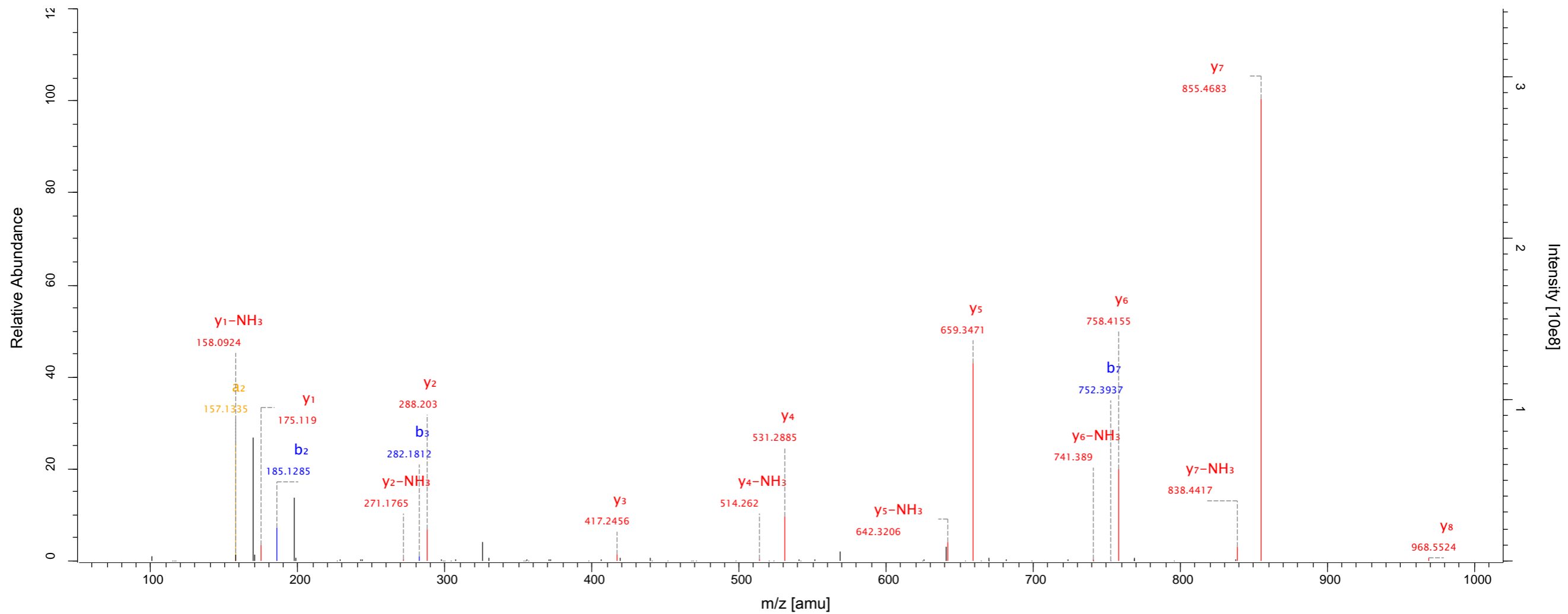
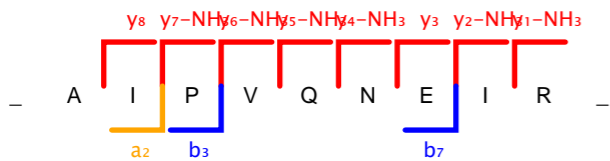
Method: FTMS; HCD



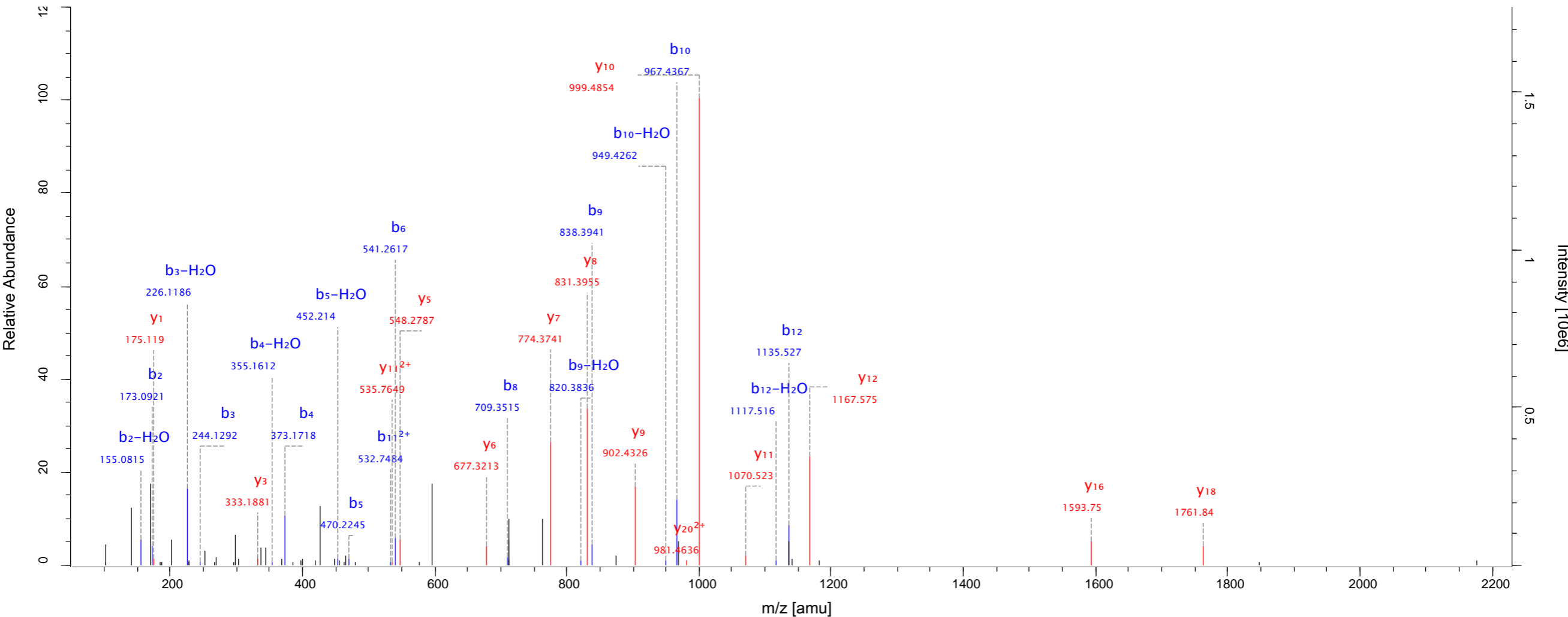
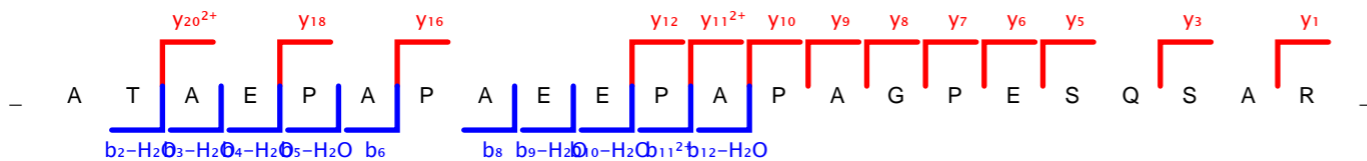
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_30
 Scan number: 14968
 Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>
 Peptide score: 221.55
 Method: FTMS; HCD



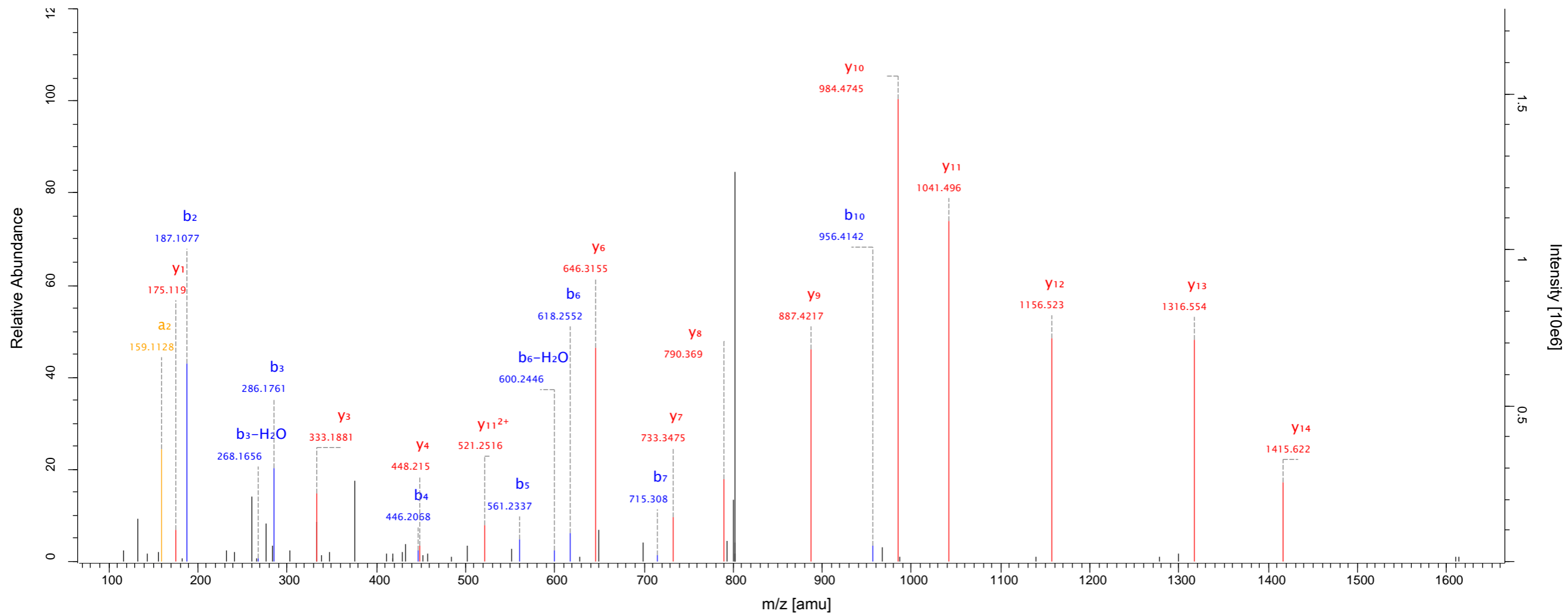
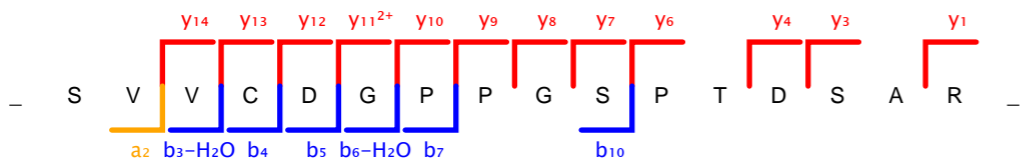
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_30
Scan number: 21371
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 124.98
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_30
 Scan number: 22176
 Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>
 Peptide score: 136.66
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_30
 Scan number: 26222
 Protein: ORFL71C - NC_006273.2[29278,30408]- <unknown description>; ORFL71C.iORF1 - NC_006273.2[29278,30354]- <unknown description>
 Peptide score: 140.1
 Method: FTMS; HCD



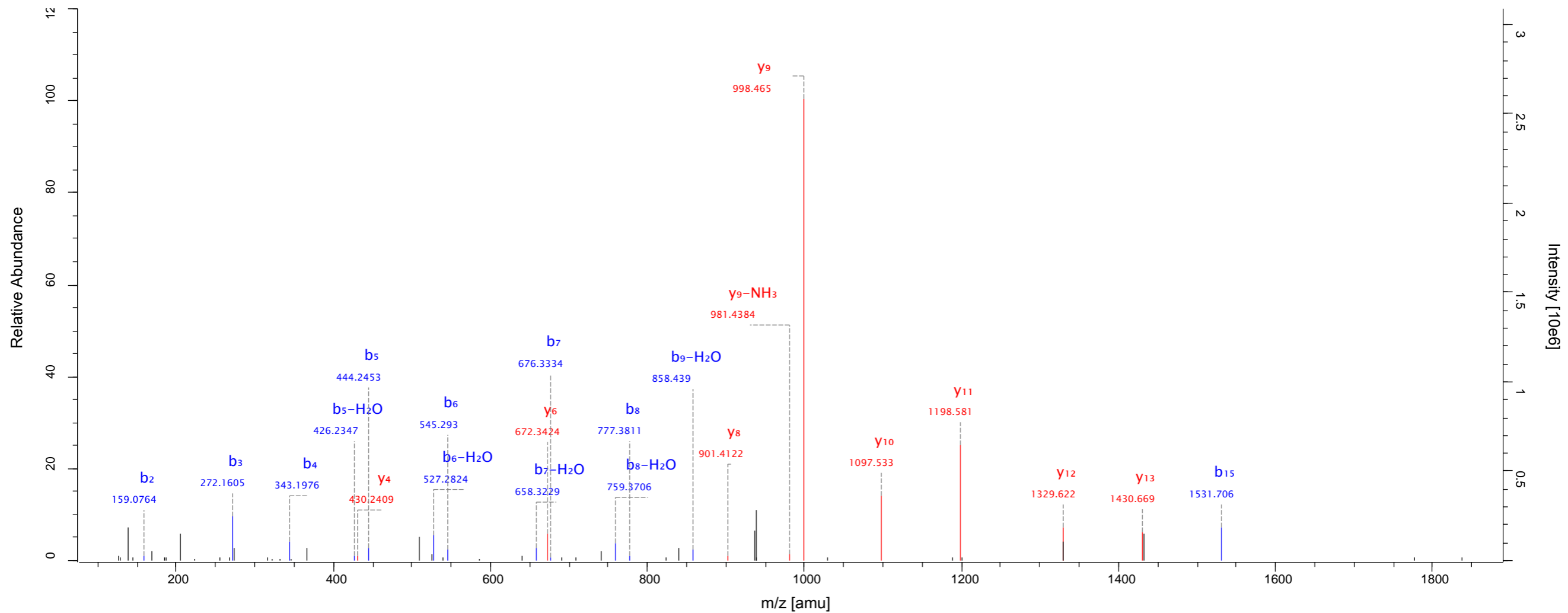
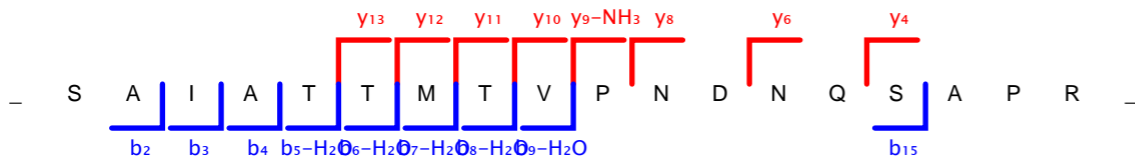
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_30

Scan number: 39700

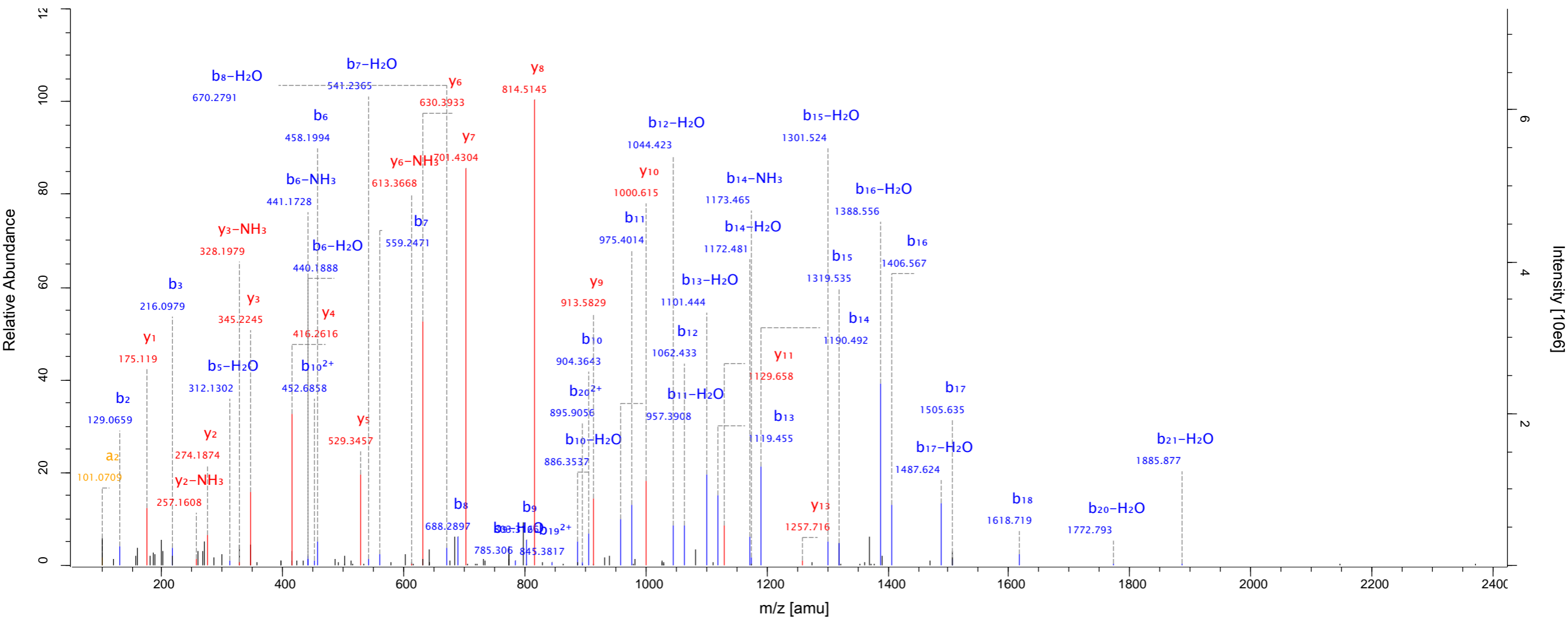
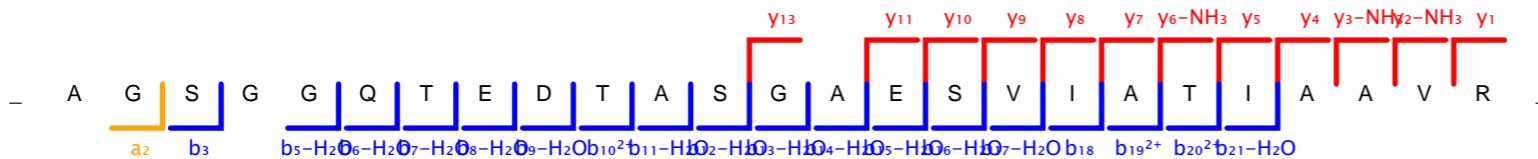
Protein: ORFL131W + NC_006273.2[73147,73947]+ <unknown description>; ORFL131W.iORF1 + NC_006273.2[73294,73947]+ <unknown description>; ORFL131W.iORF2 + NC_006273.2[73501,73947]+ <unknown description>

Peptide score: 118.21

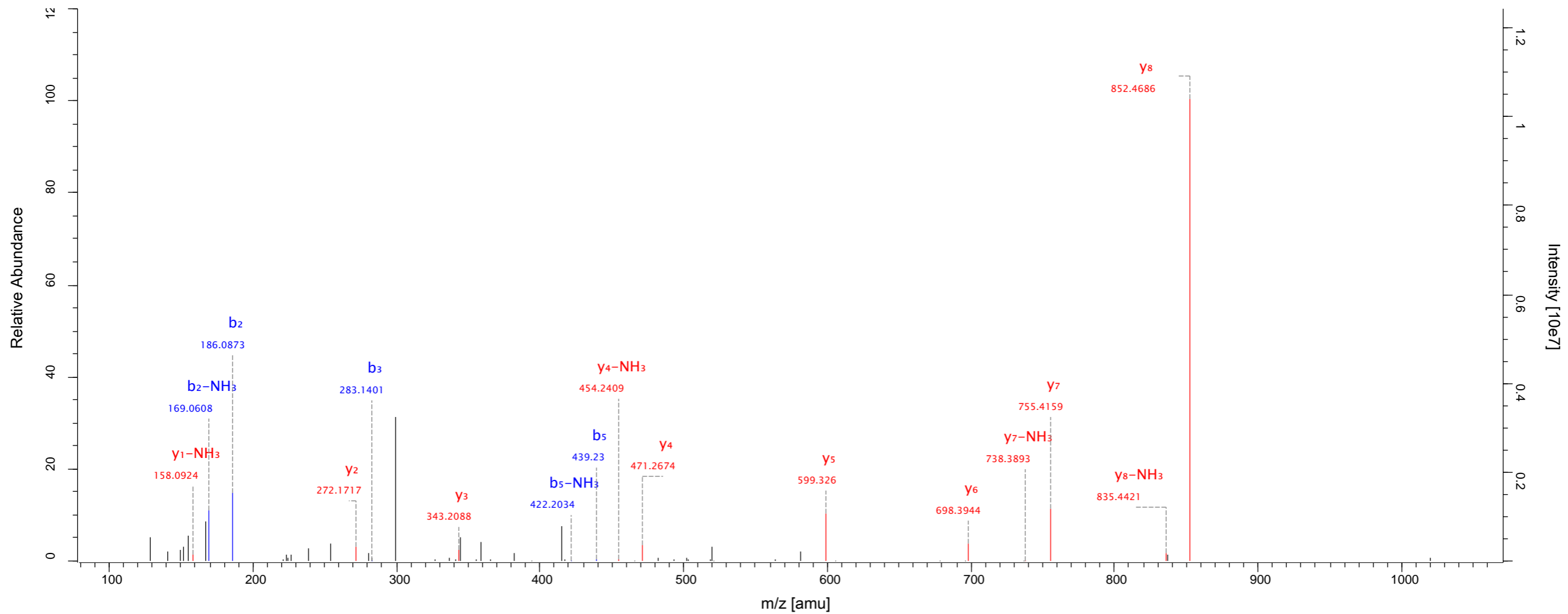
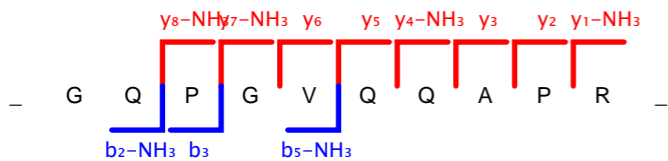
Method: FTMS; HCD



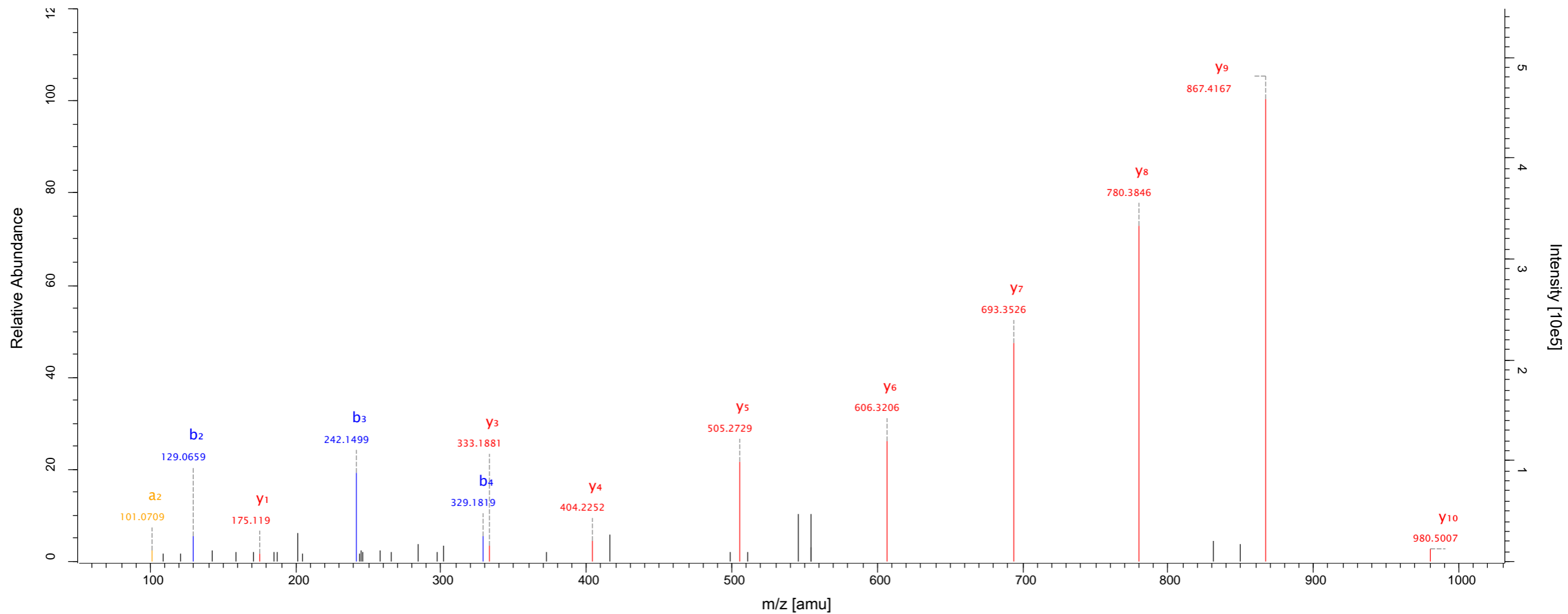
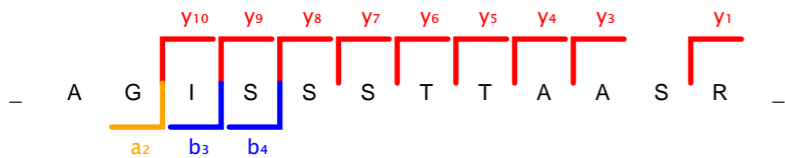
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_30
 Scan number: 49429
 Protein: ORFL71C - NC_006273.2[29278,30408]- <unknown description>; ORFL71C.iORF1 - NC_006273.2[29278,30354]- <unknown description>
 Peptide score: 234.23
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_30
Scan number: 6732
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 80.69
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_30
Scan number: 7069
Protein: ORFL253W.iORF1 + NC_006273.2[161917,163748]+ <unknown description>; ORFL251W + NC_006273.2[161872,163748]+ <unknown description>
Peptide score: 78.65
Method: FTMS; HCD



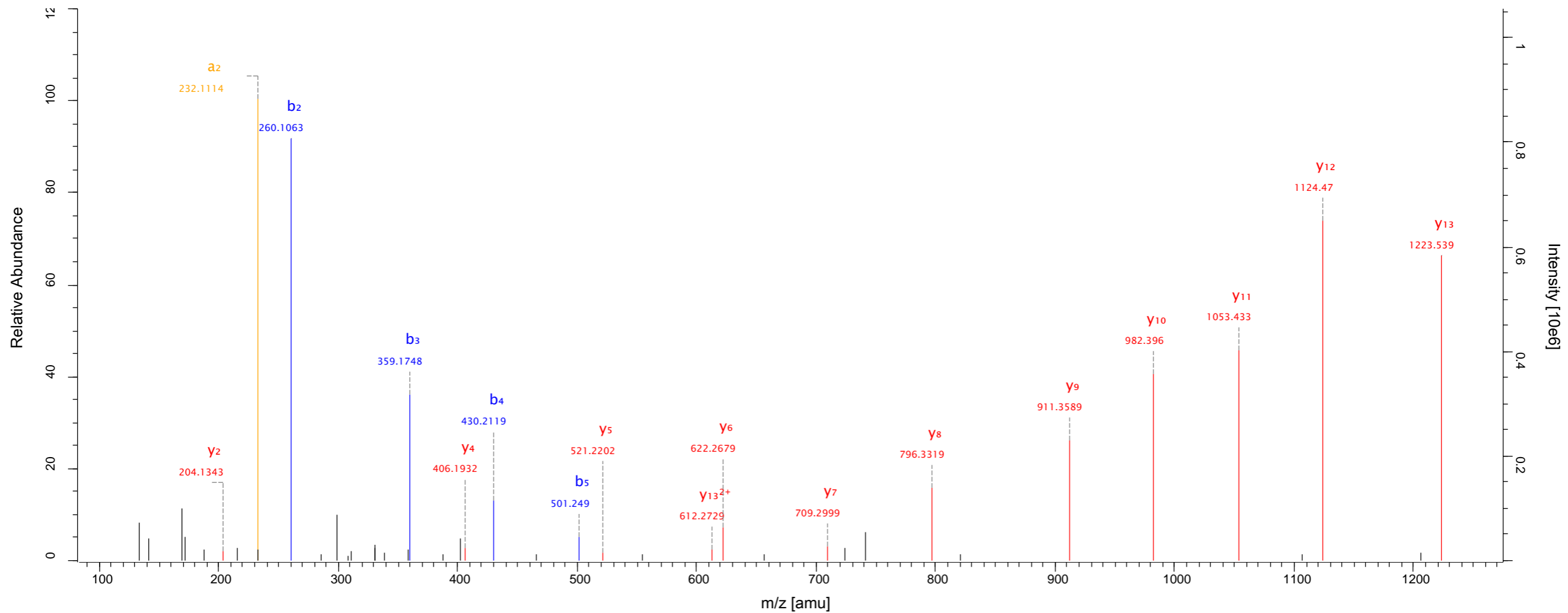
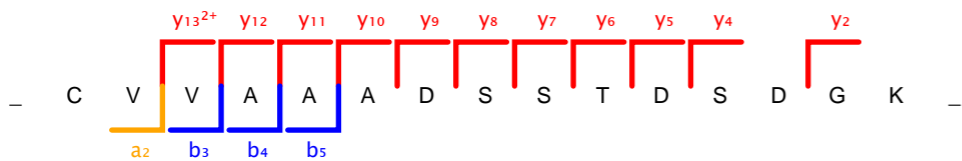
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_30

Scan number: 9263

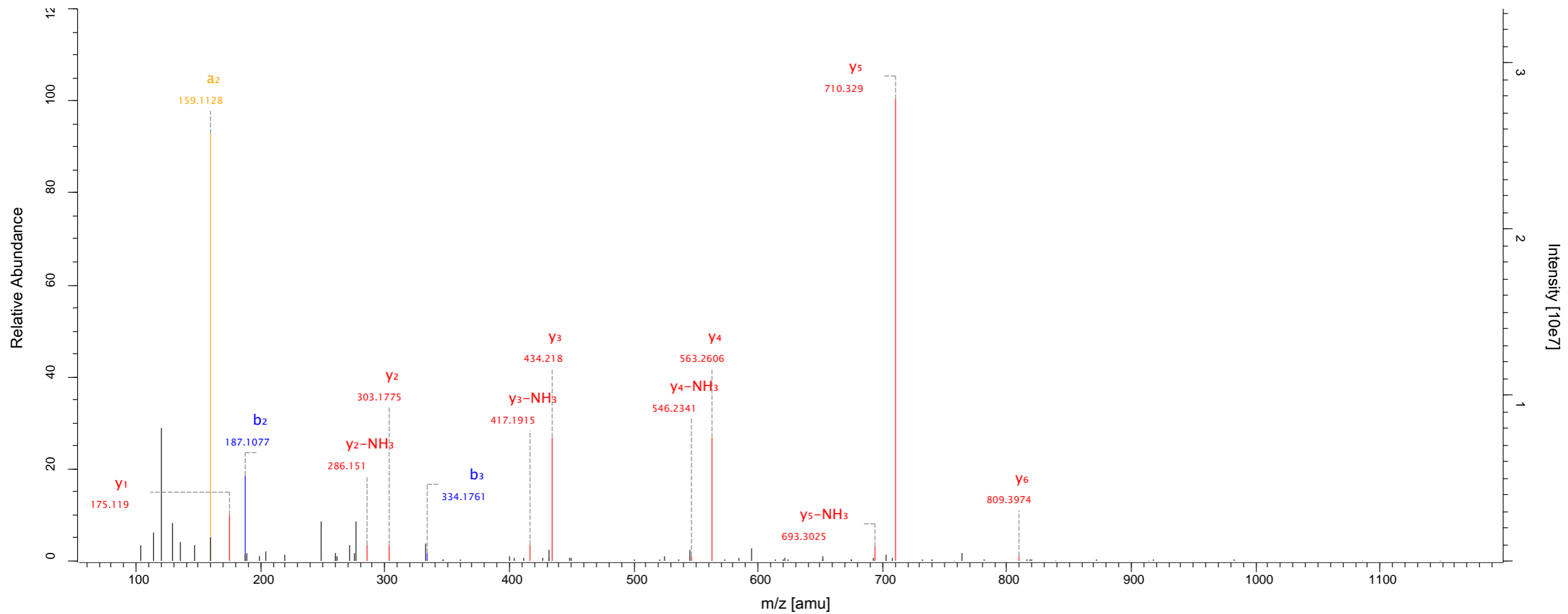
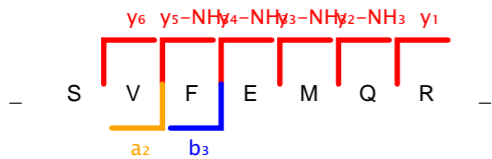
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 124.86

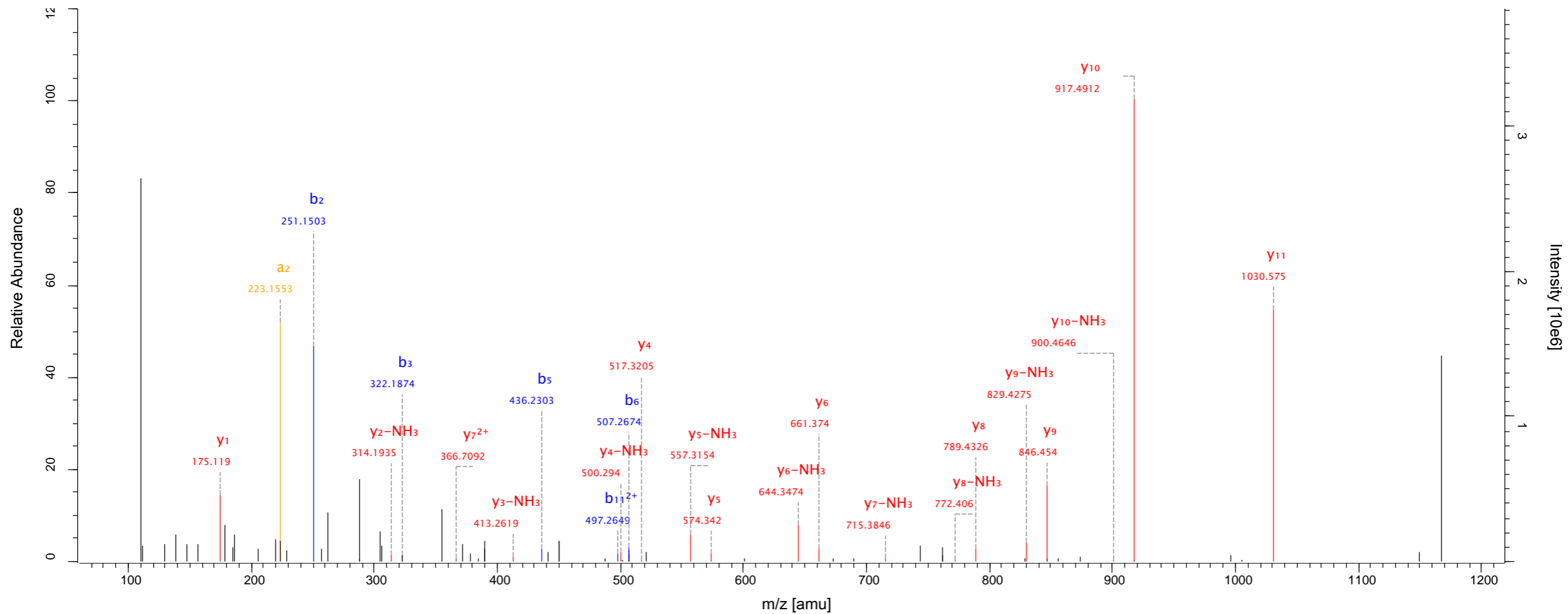
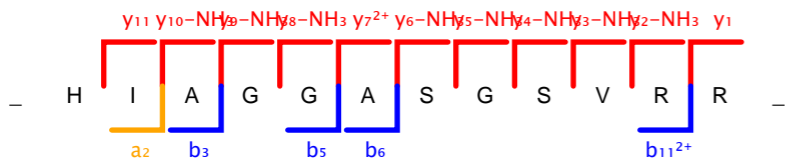
Method: FTMS; HCD



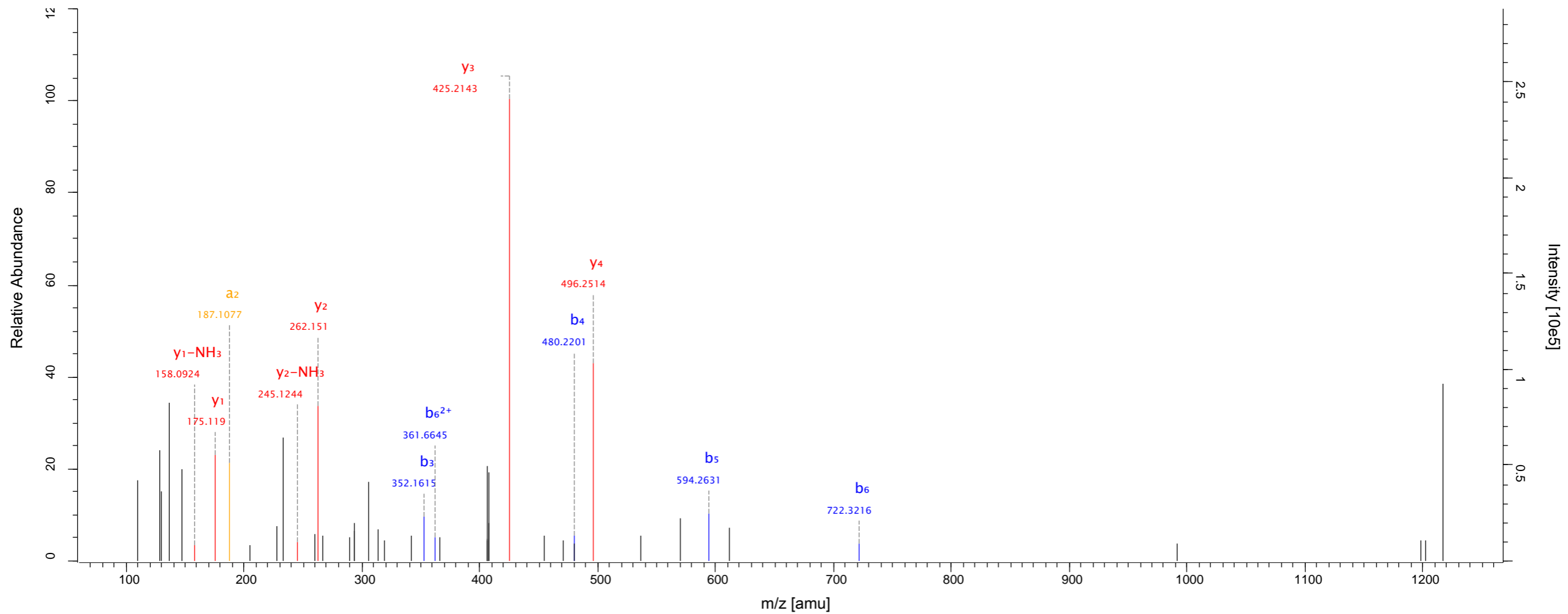
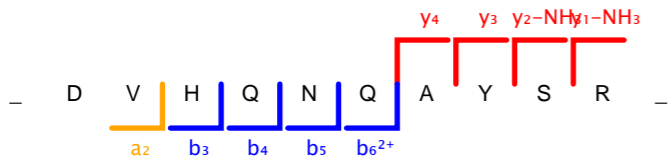
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_31
Scan number: 18351
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 114.02
Method: FTMS; HCD



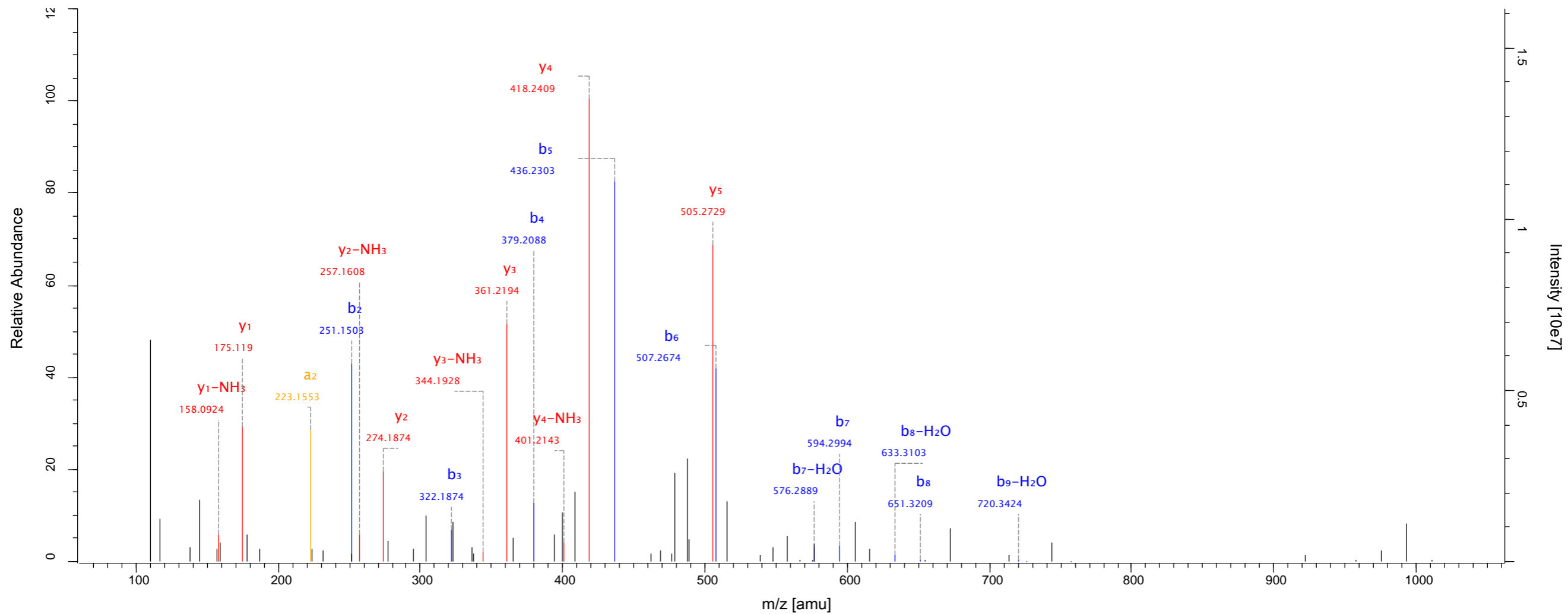
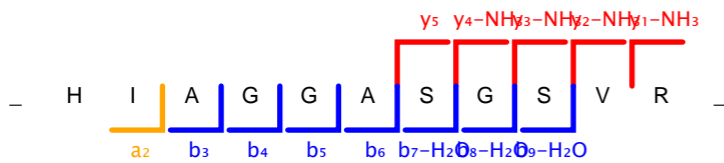
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_31
 Scan number: 4750
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 112.36
 Method: FTMS; HCD



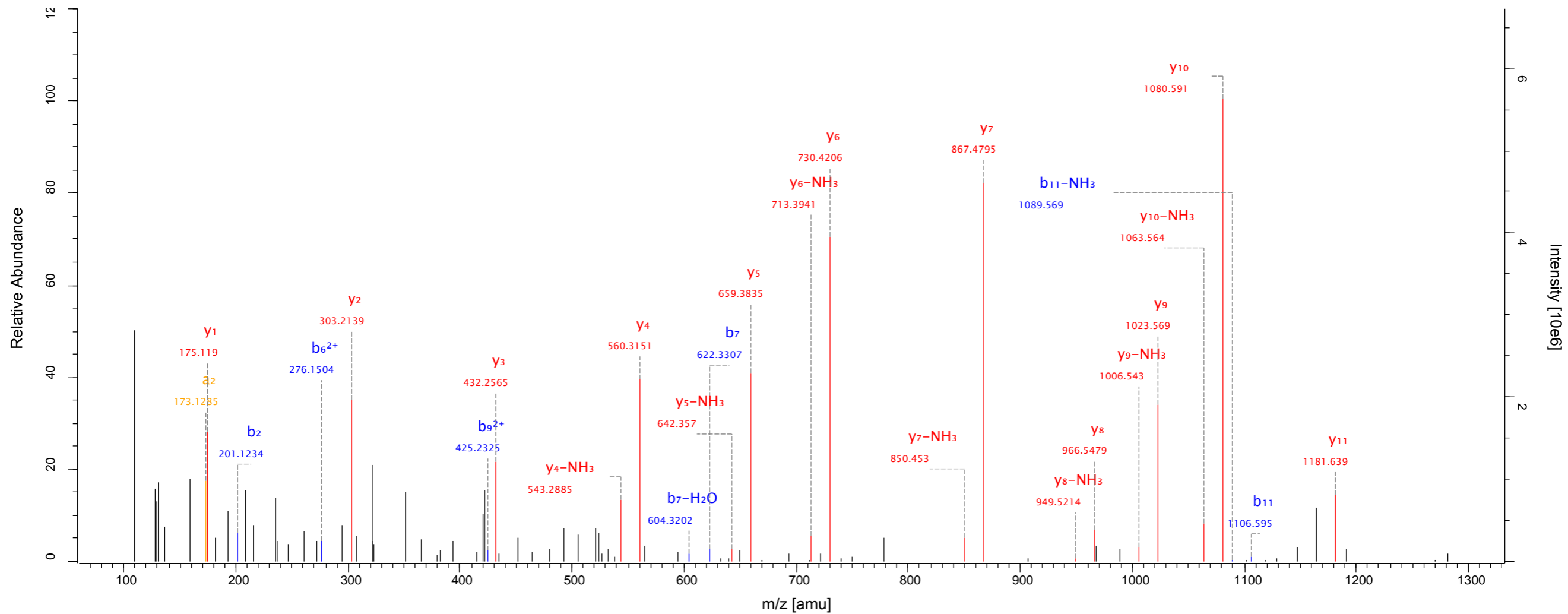
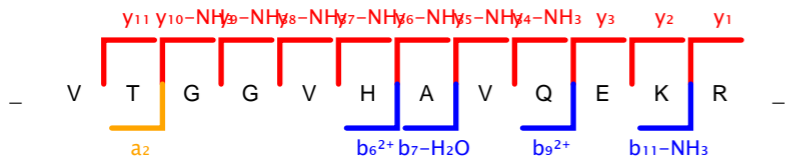
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_31
Scan number: 4775
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 59.43
Method: FTMS; HCD



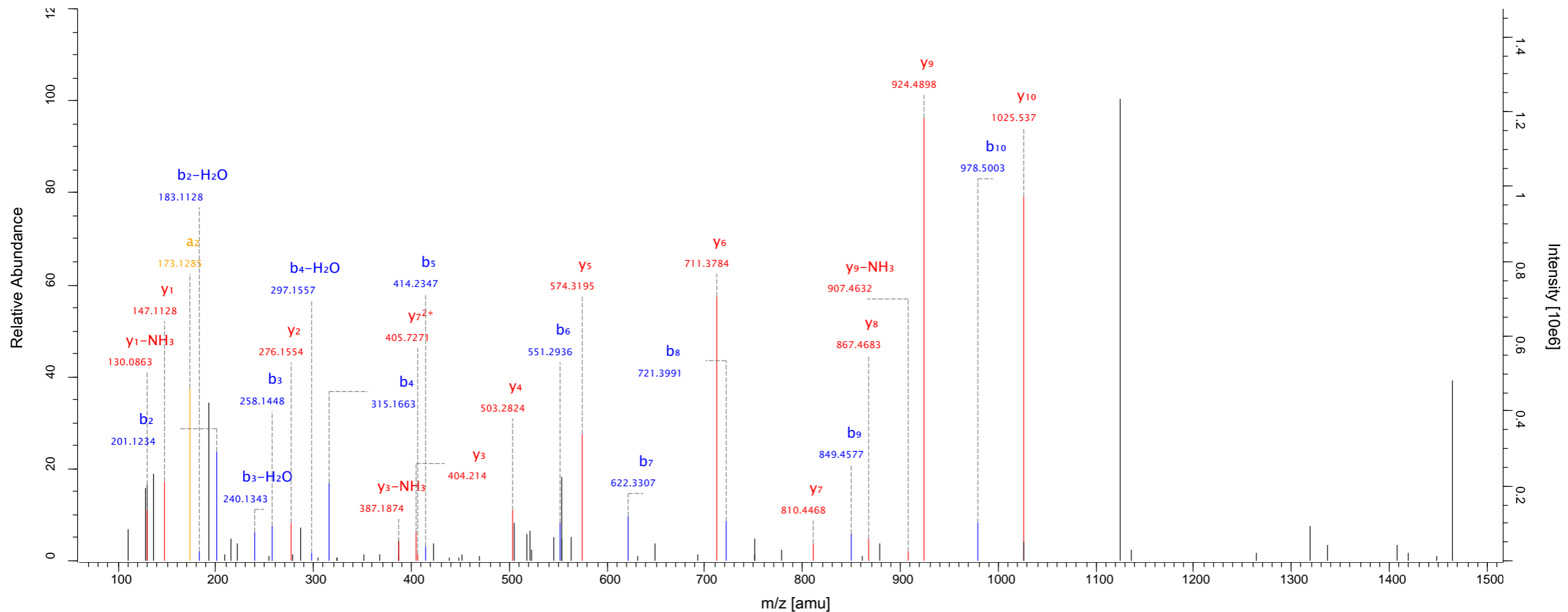
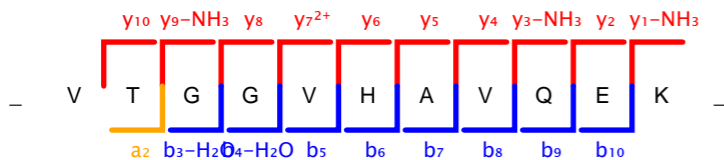
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_31
 Scan number: 5127
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 123.35
 Method: FTMS; HCD



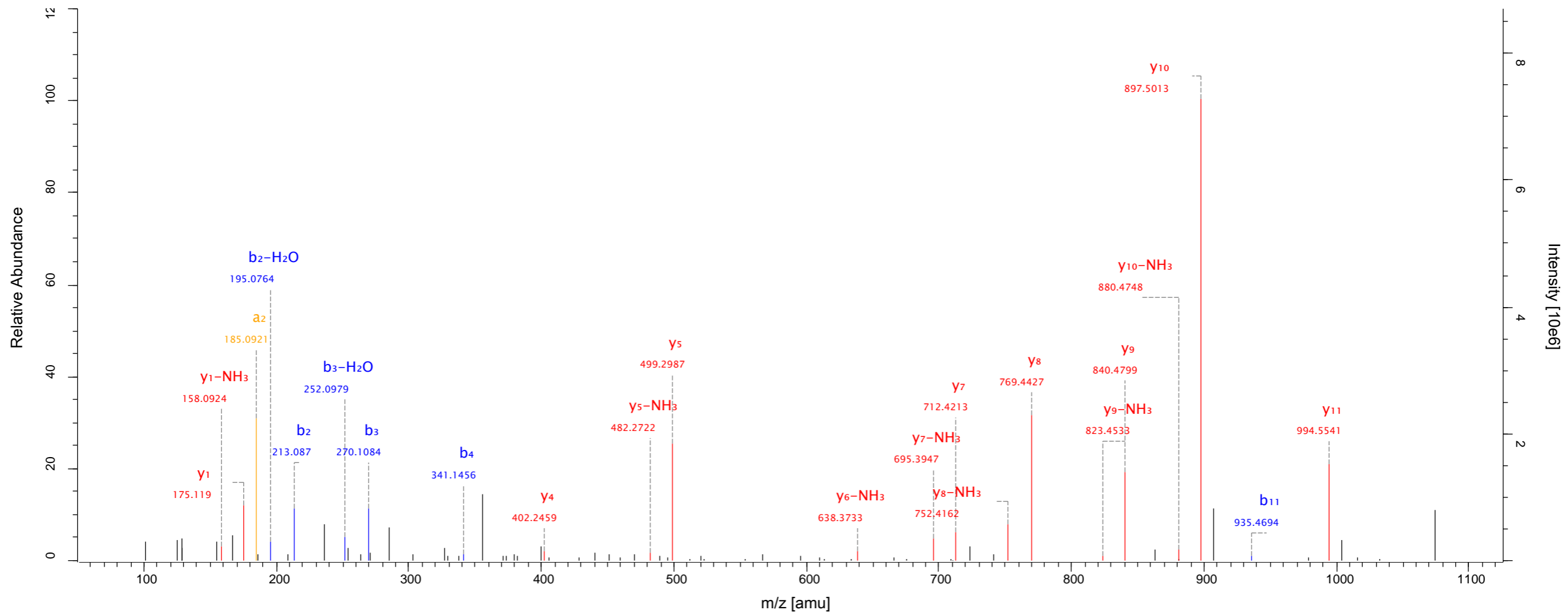
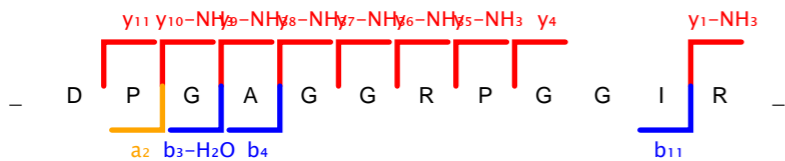
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_31
 Scan number: 5302
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 149.96
 Method: FTMS; HCD



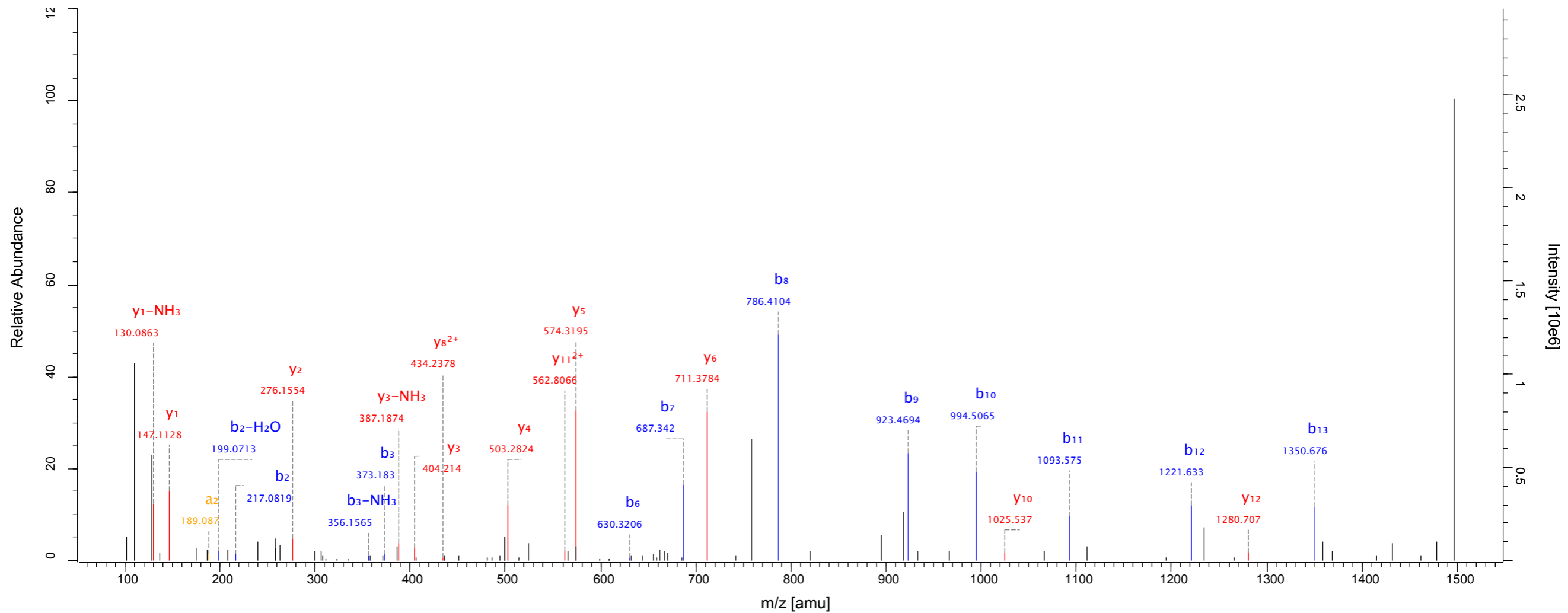
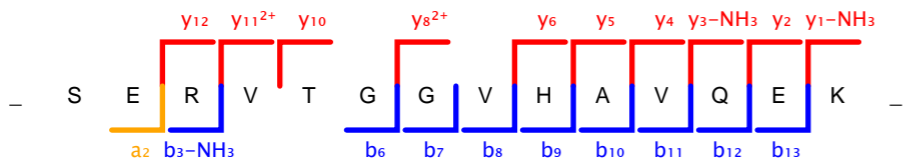
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_31
Scan number: 5972
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 180.19
Method: FTMS; HCD



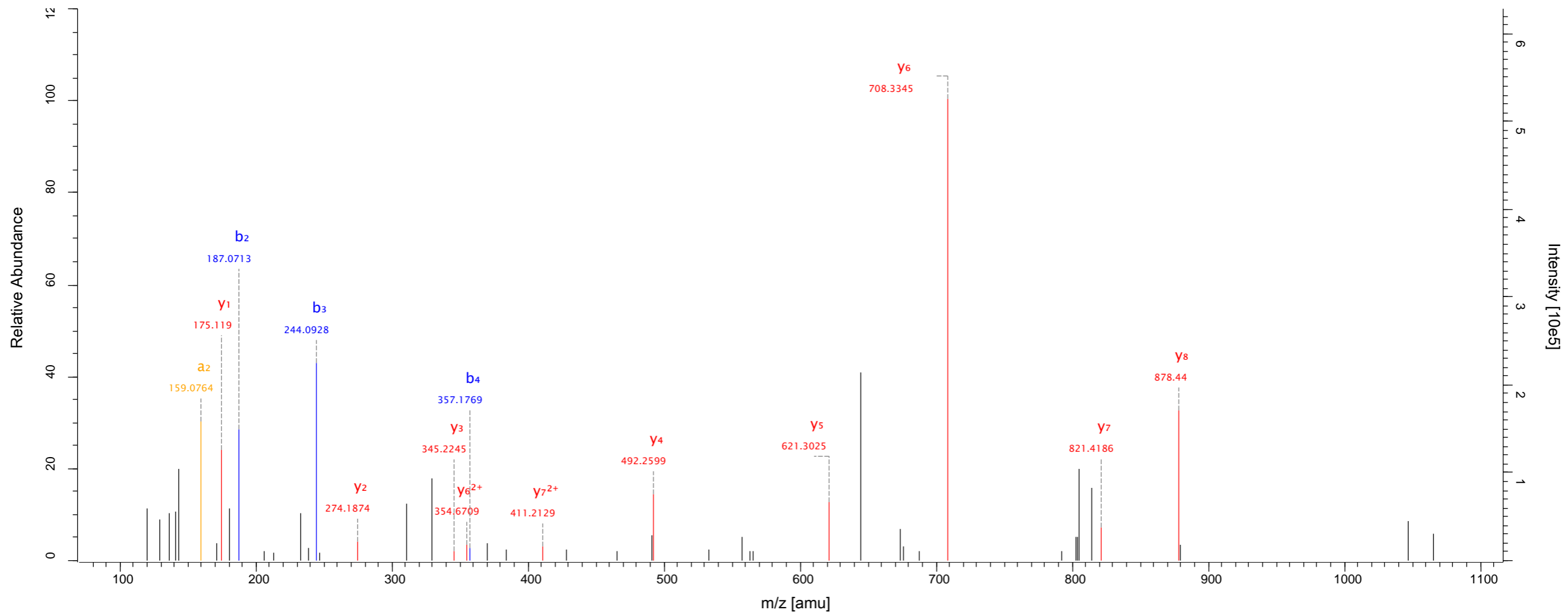
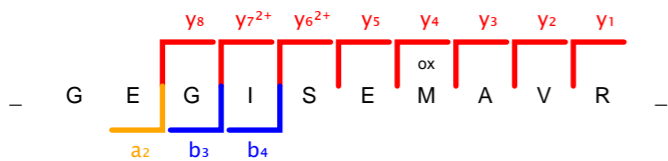
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_31
 Scan number: 7324
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 138.25
 Method: FTMS; HCD



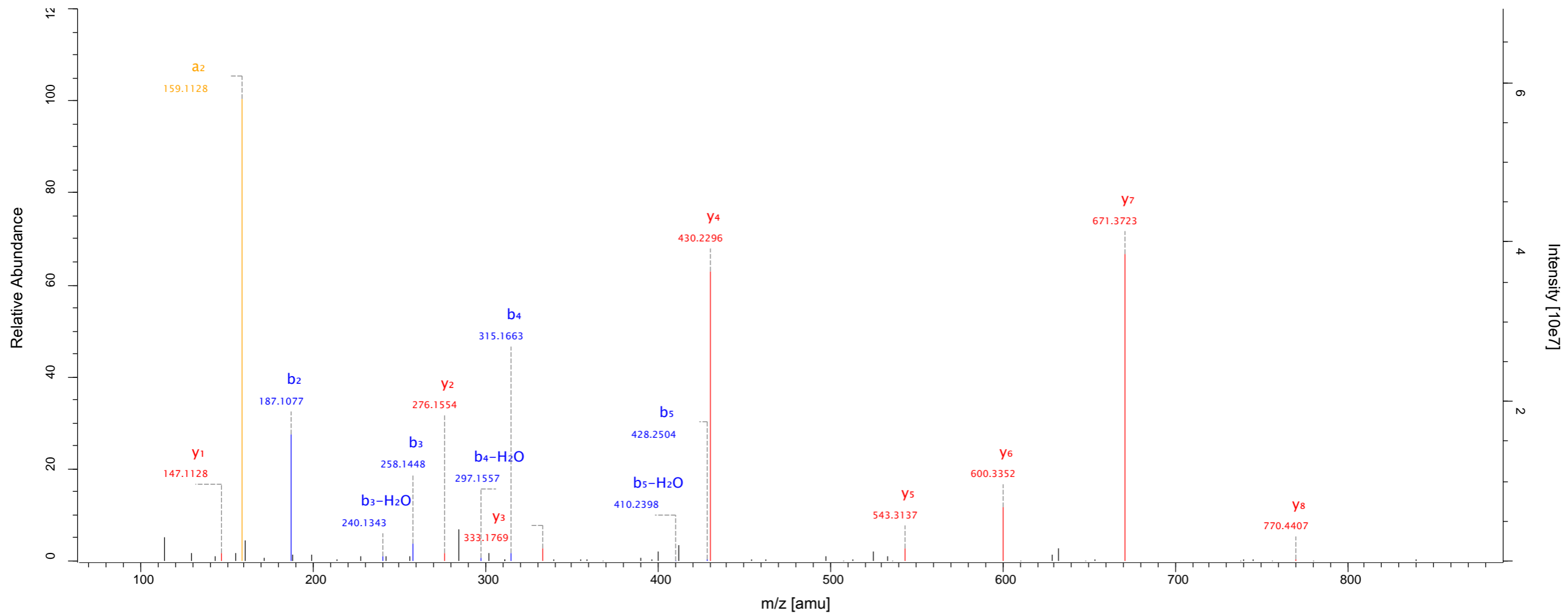
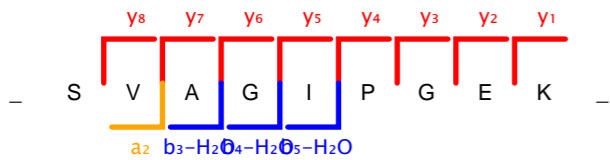
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_31
Scan number: 7421
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 107.35
Method: FTMS; HCD



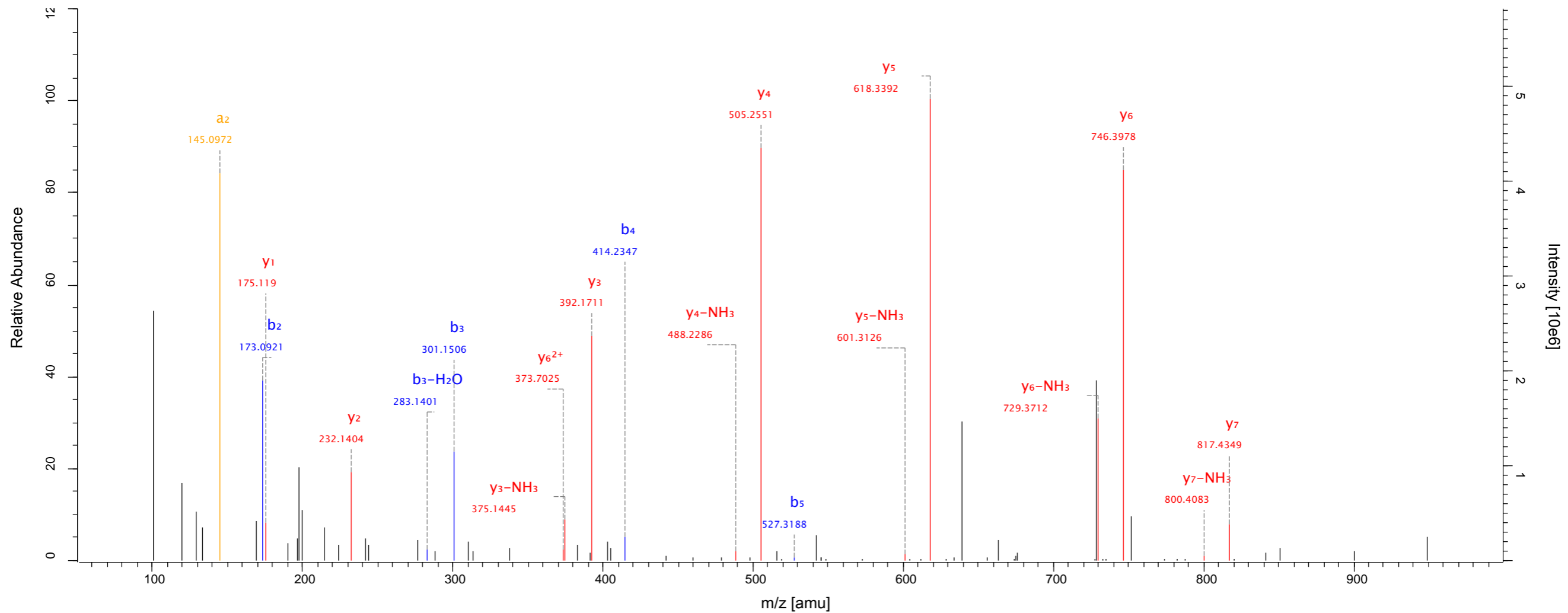
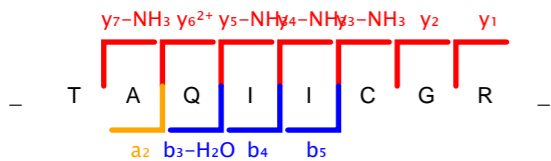
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
Scan number: 10241
Protein: ORFL18W + NC_006273.2[8020,8253]+ <unknown description>
Peptide score: 79.15
Method: FTMS; HCD



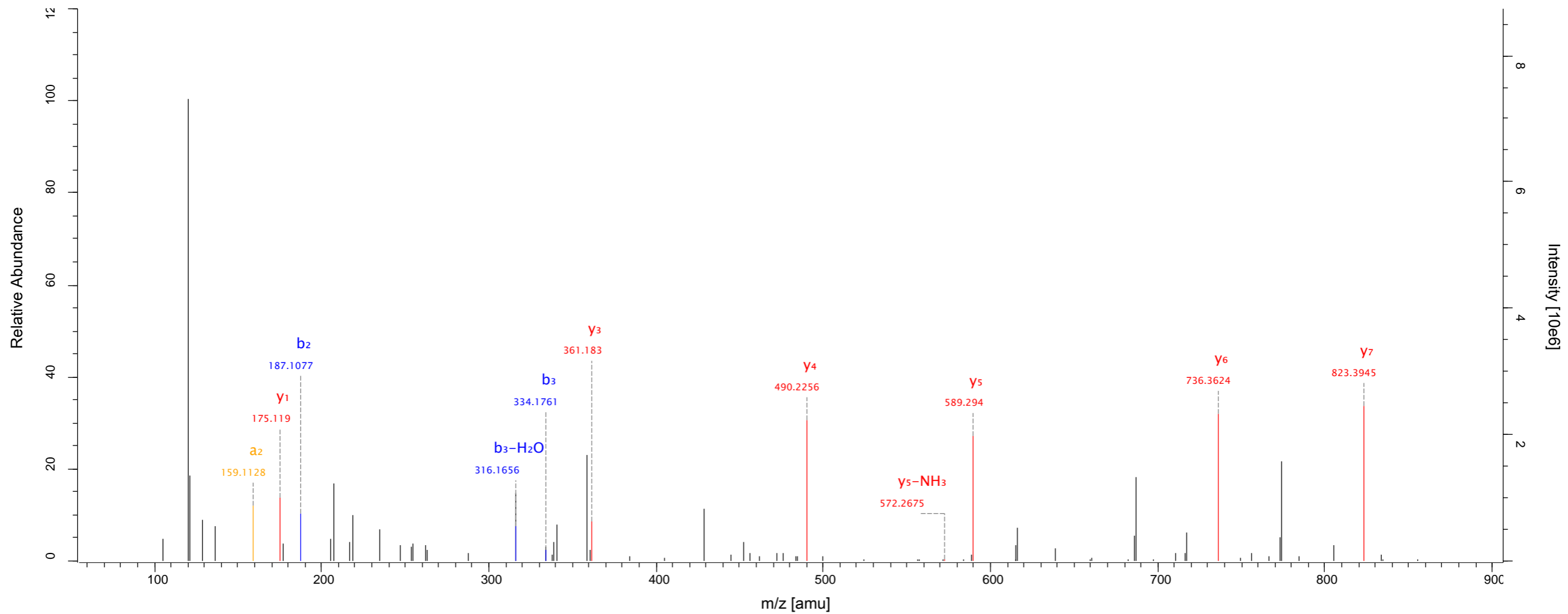
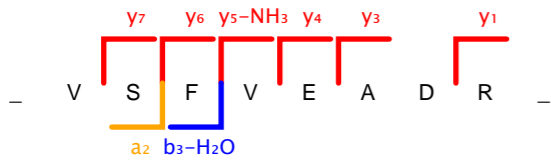
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_34
Scan number: 11522
Protein: ORFL301C - NC_006273.2[185358,185666]- <unknown description>
Peptide score: 88.18
Method: FTMS; HCD



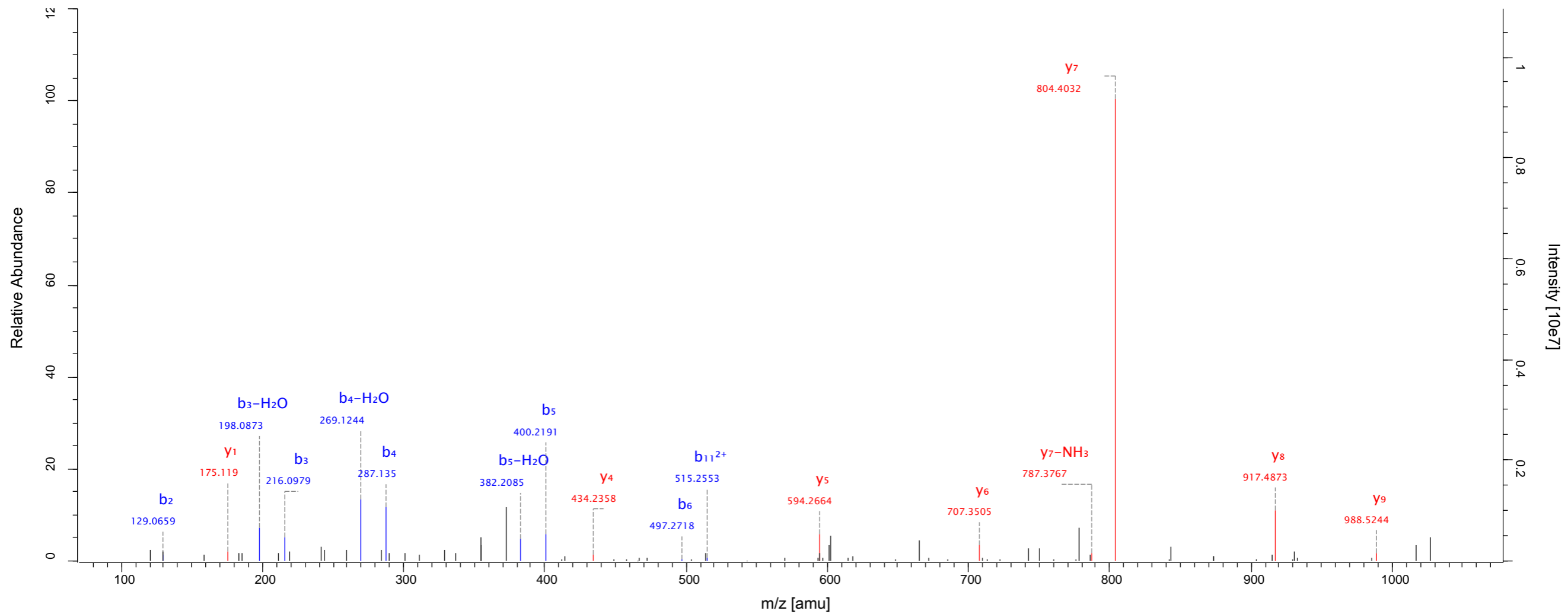
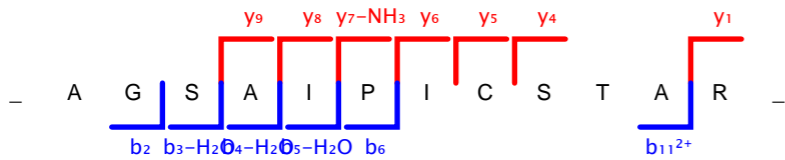
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
Scan number: 12310
Protein: ORFL137W.IORF1 + NC_006273.2[75299,75529]+ <unknown description>
Peptide score: 128.69
Method: FTMS; HCD



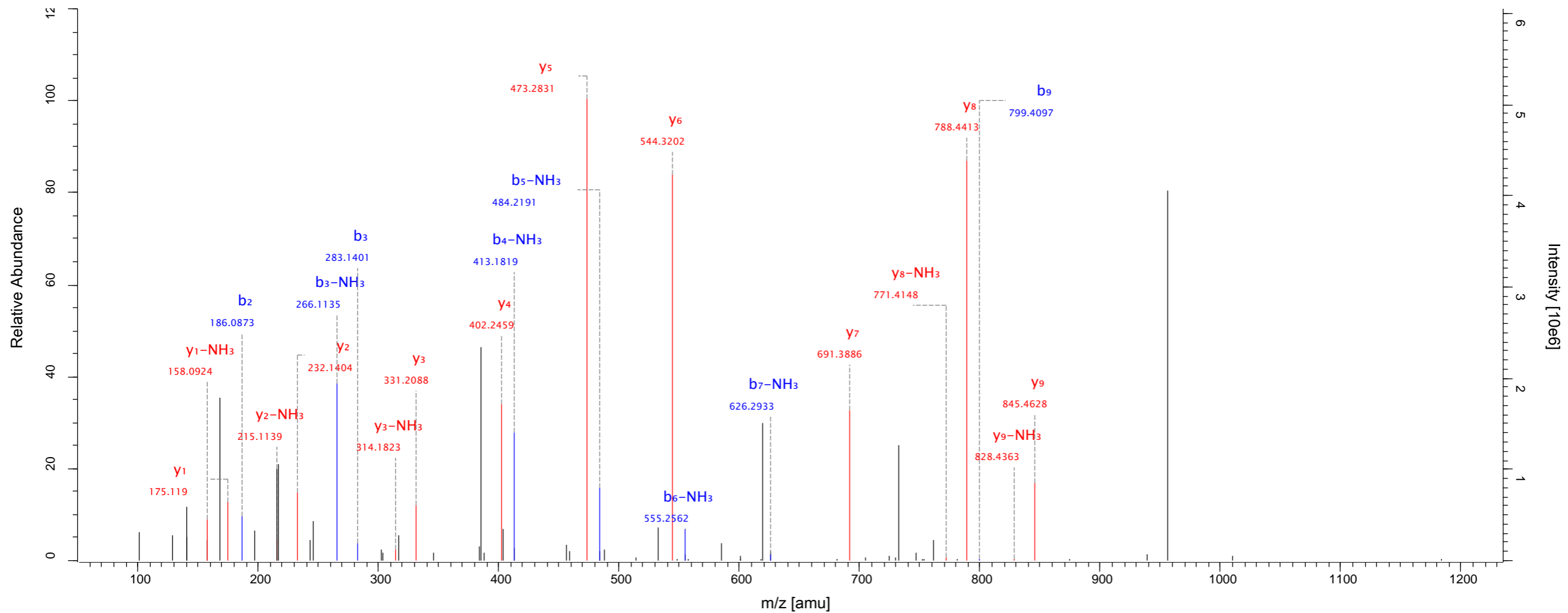
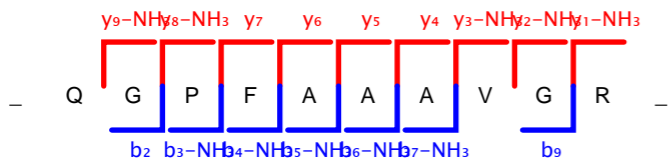
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
Scan number: 14863
Protein: ORFL27C - NC_006273.2[7912,13003]- <unknown description>; ORFL27C.iORF1 - NC_006273.2[7912,8184]- <unknown description>
Peptide score: 56.72
Method: FTMS; HCD



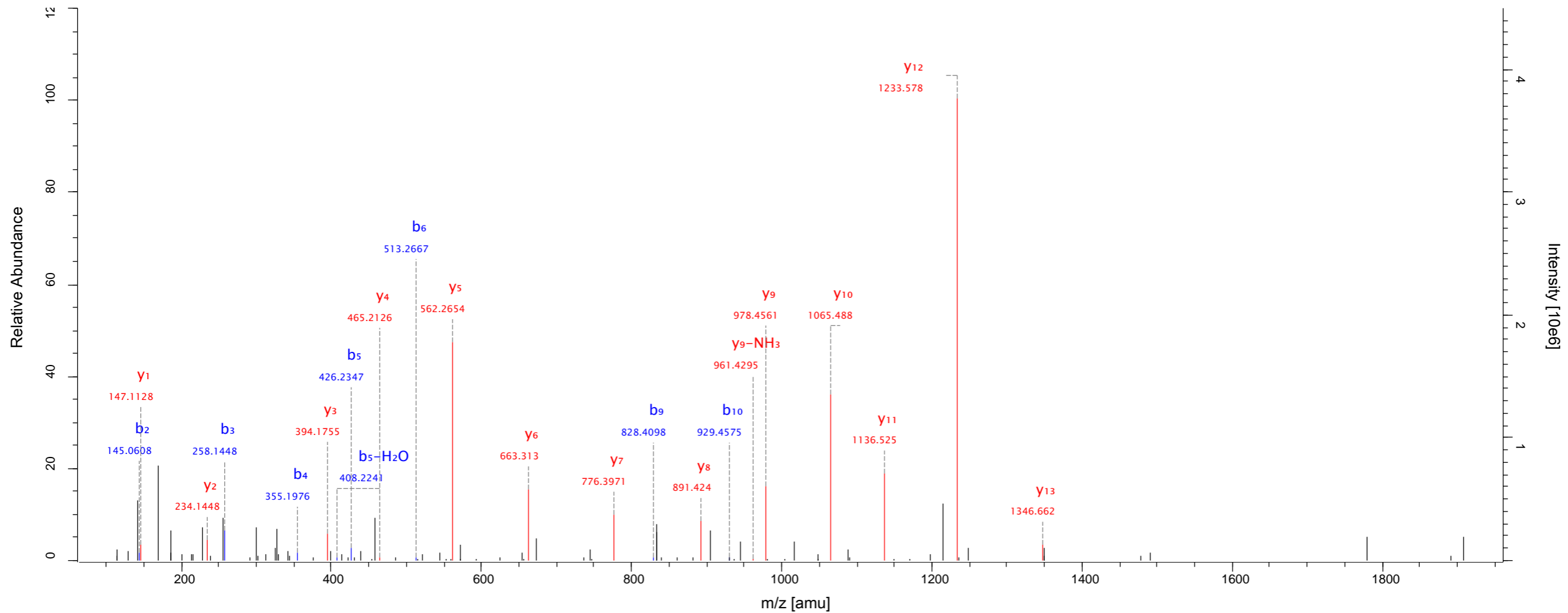
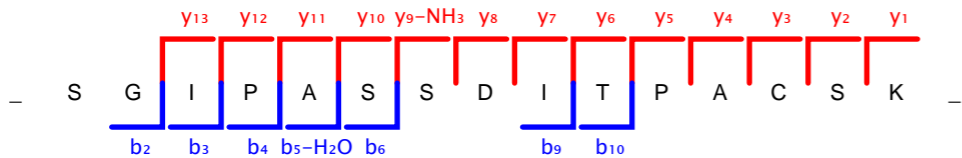
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
Scan number: 17577
Protein: ORFL253W.iORF3 + NC_006273.2[162059,162286]+ <unknown description>
Peptide score: 106.62
Method: FTMS; HCD



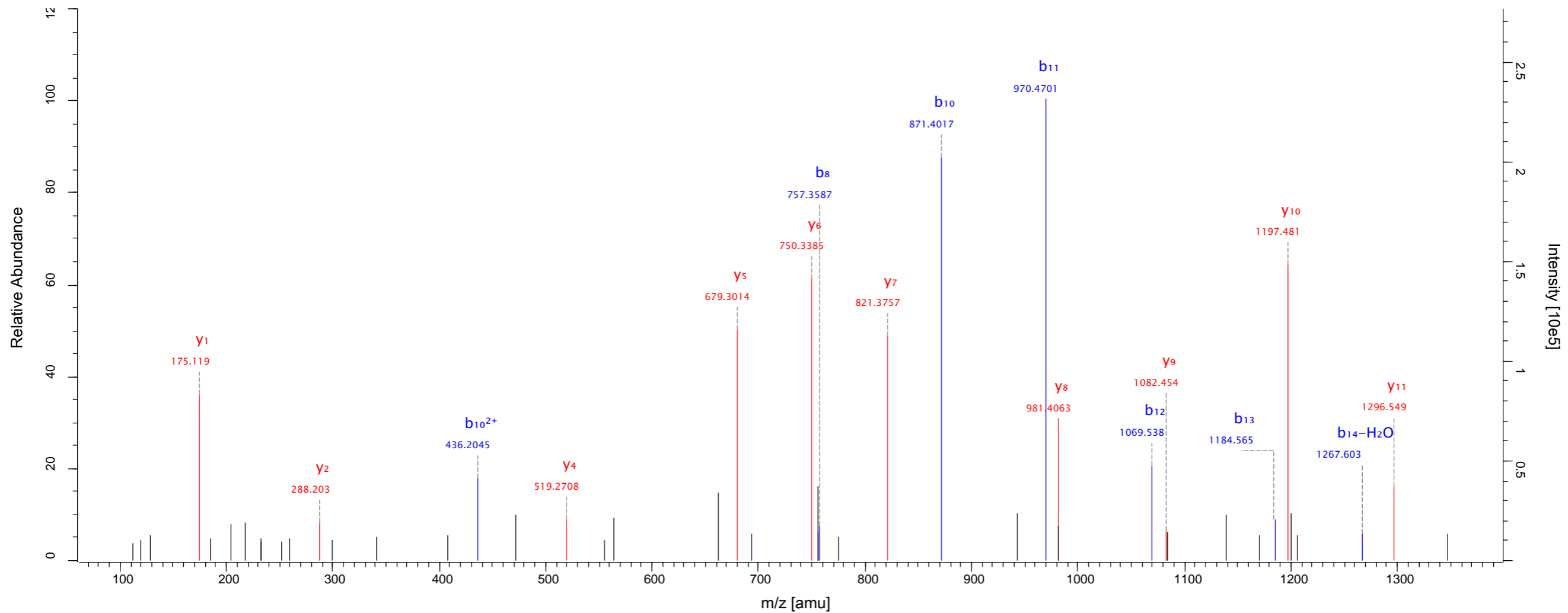
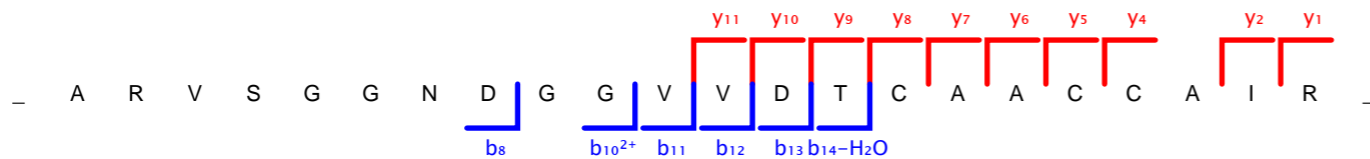
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 18035
Protein: ORFL121C - NC_006273.2[65579,65851]- <unknown description>
Peptide score: 97.97
Method: FTMS; HCD



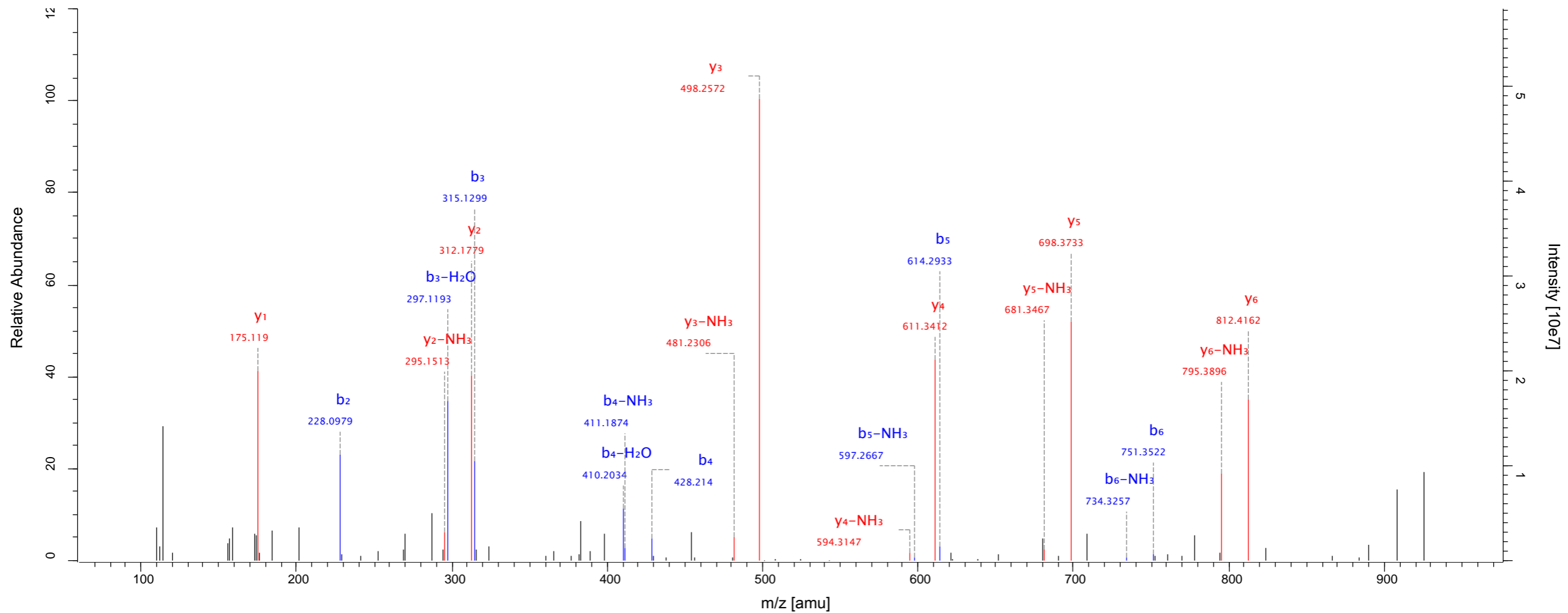
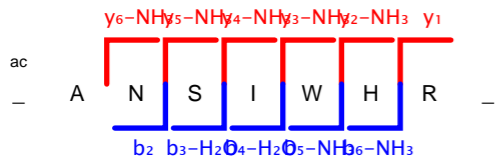
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 19077
Protein: ORFL298W + NC_006273.2[184737,184916]+ <unknown description>
Peptide score: 83.87
Method: FTMS; HCD



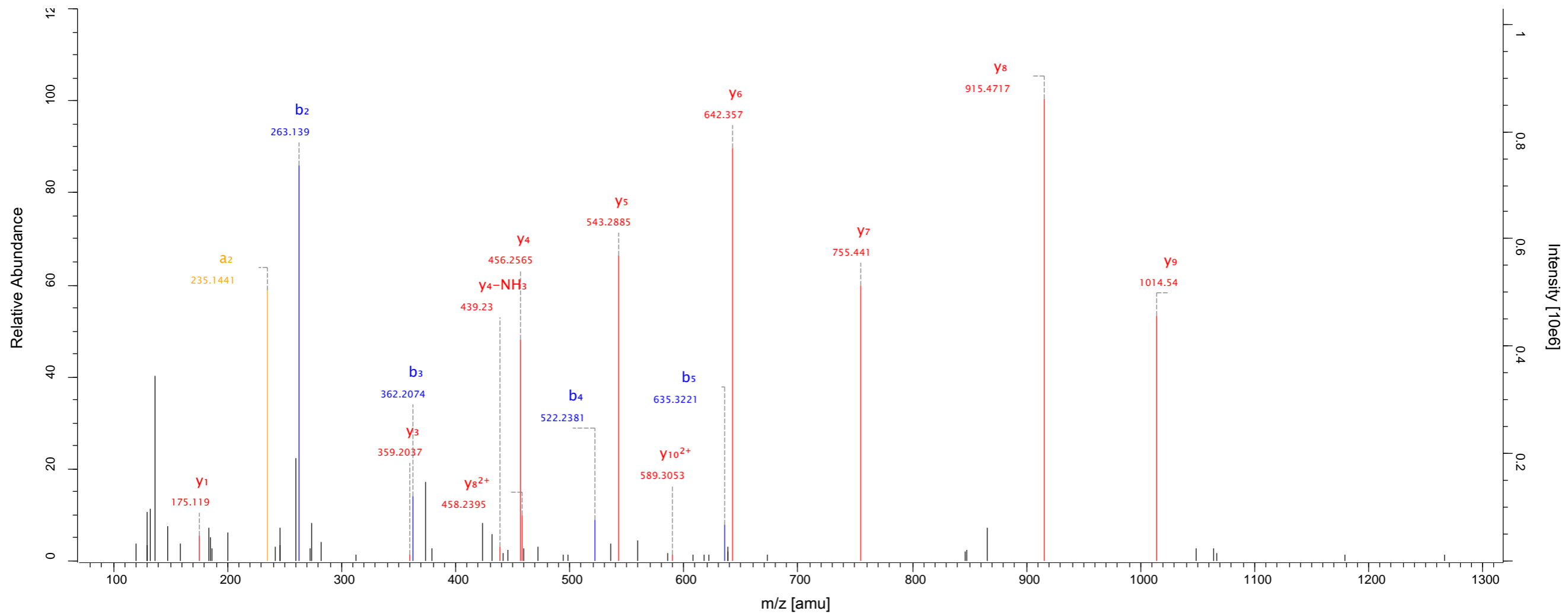
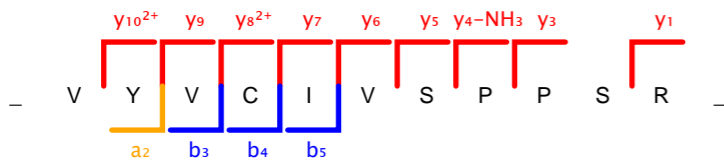
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 19716
Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
Peptide score: 72.64
Method: FTMS; HCD



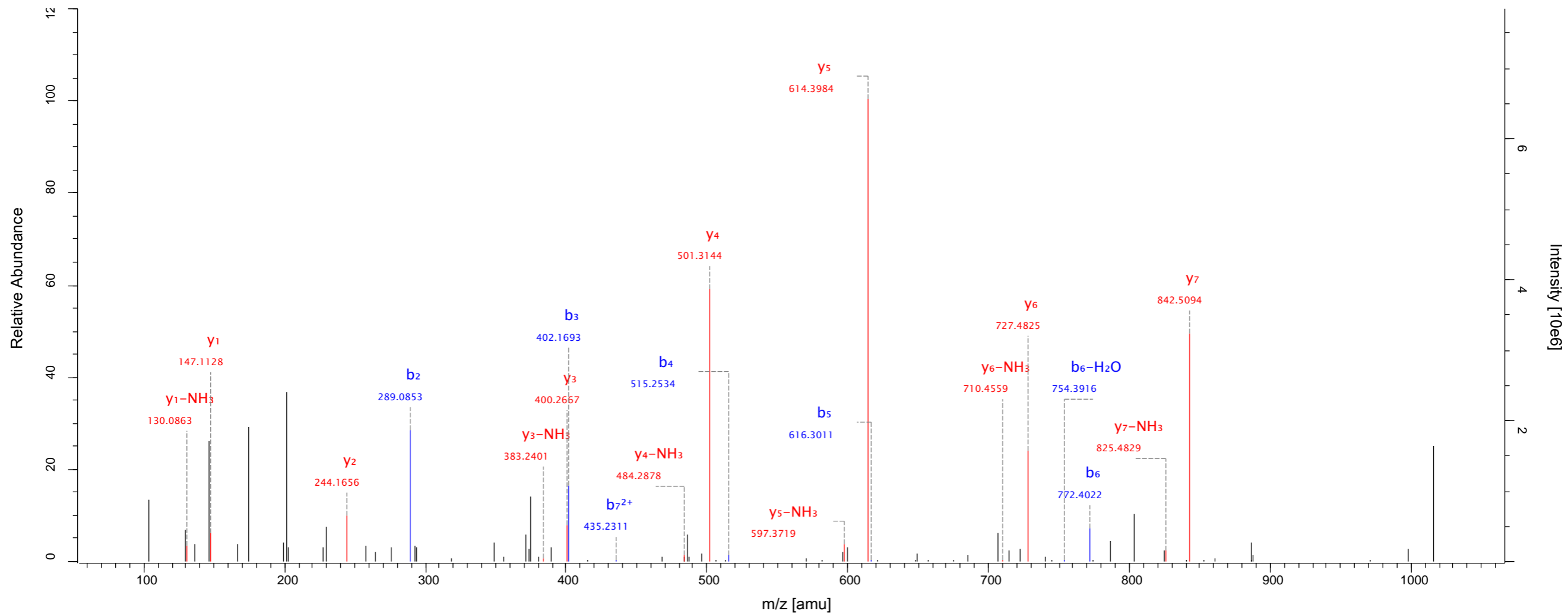
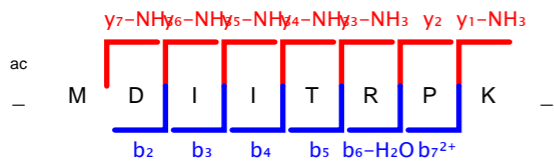
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
 Scan number: 21160
 Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
 Peptide score: 176.66
 Method: FTMS; HCD



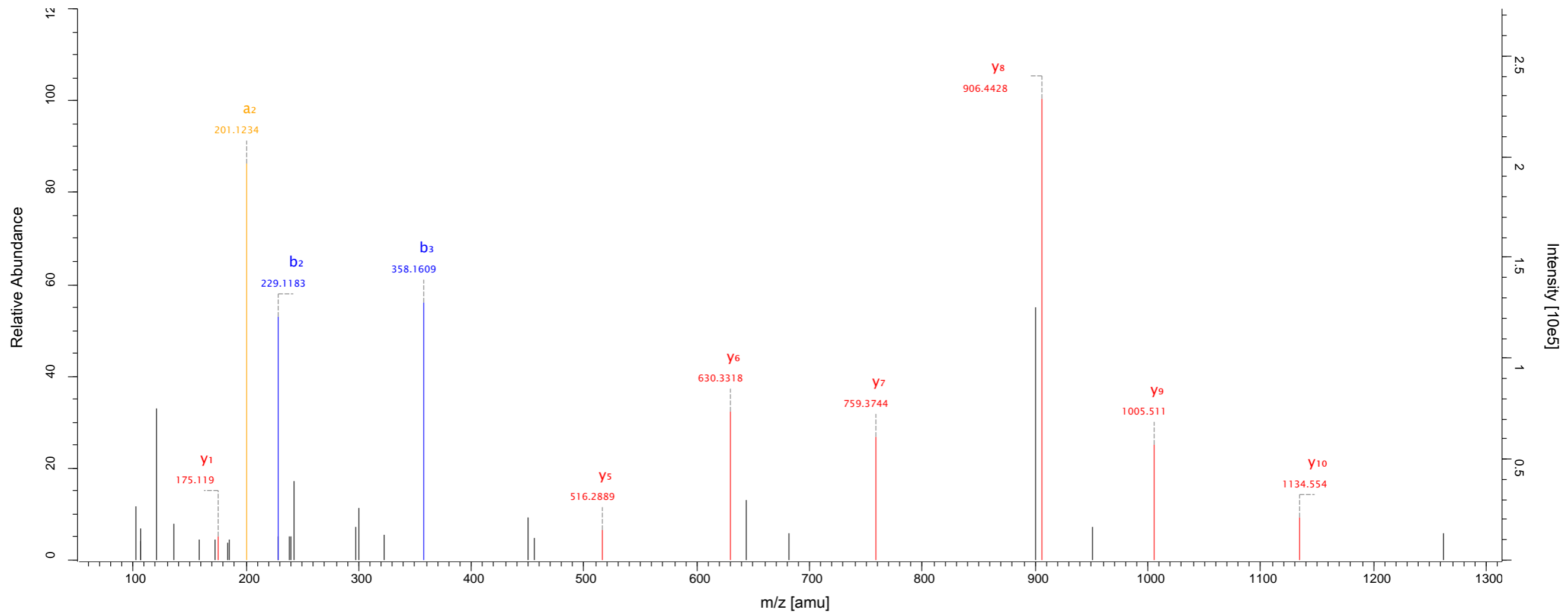
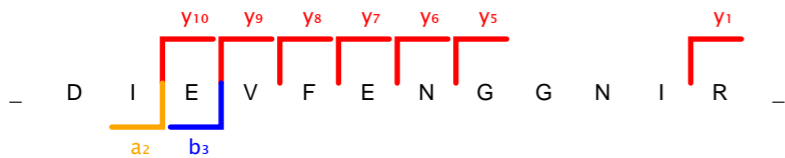
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 27189
Protein: ORFL7C - NC_006273.2[4436,4690]- <unknown description>
Peptide score: 111.65
Method: FTMS; HCD



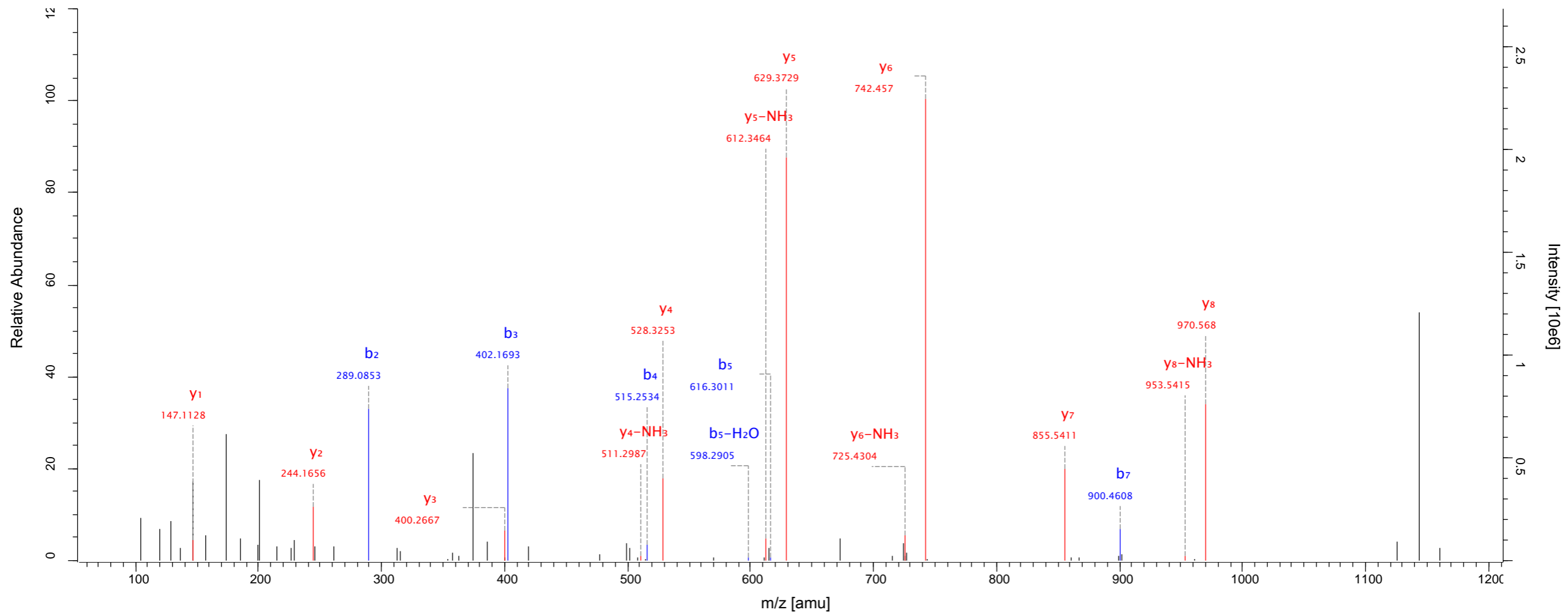
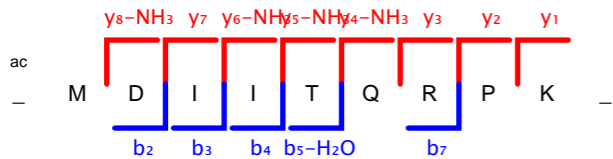
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_34
Scan number: 30477
Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>
Peptide score: 125.5
Method: FTMS; HCD



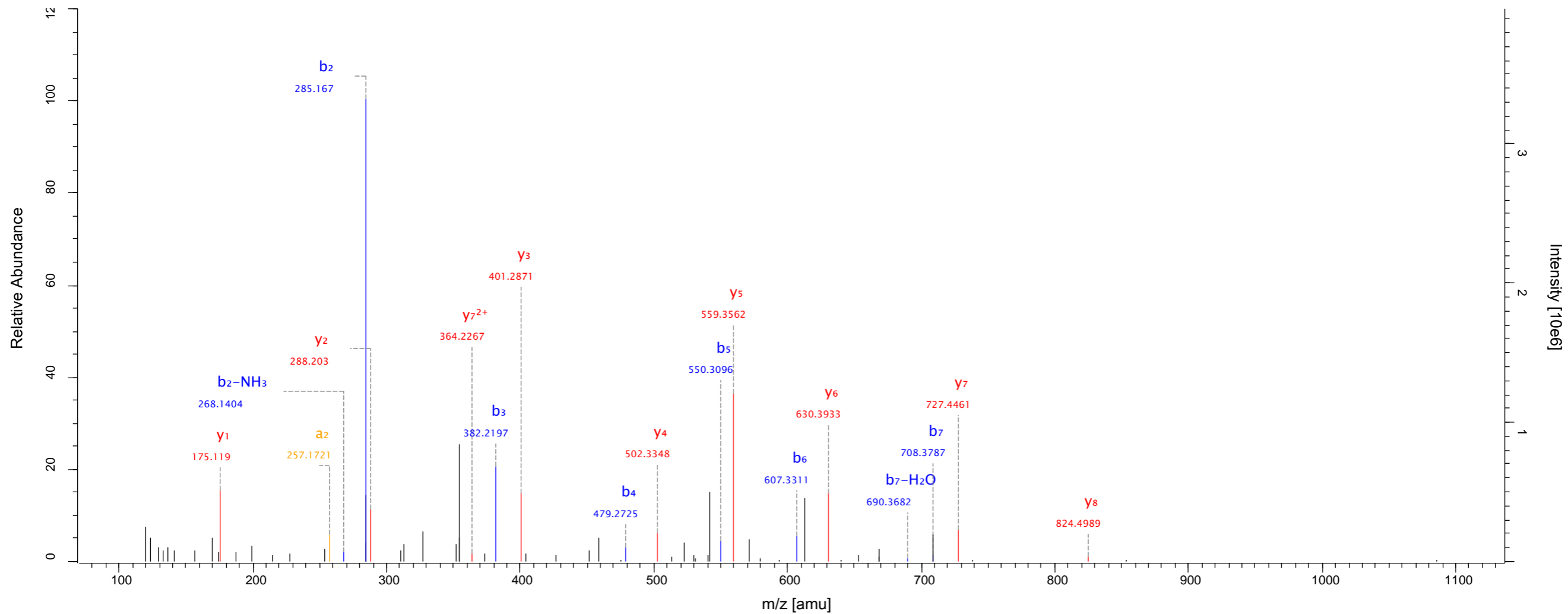
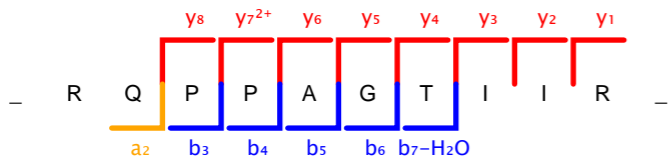
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 31902
Protein: ORFL154W + NC_006273.2[96843,97115]+ <unknown description>; ORFL154W.iORF1 + NC_006273.2[96972,97115]+ <unknown description>
Peptide score: 68.07
Method: FTMS; HCD



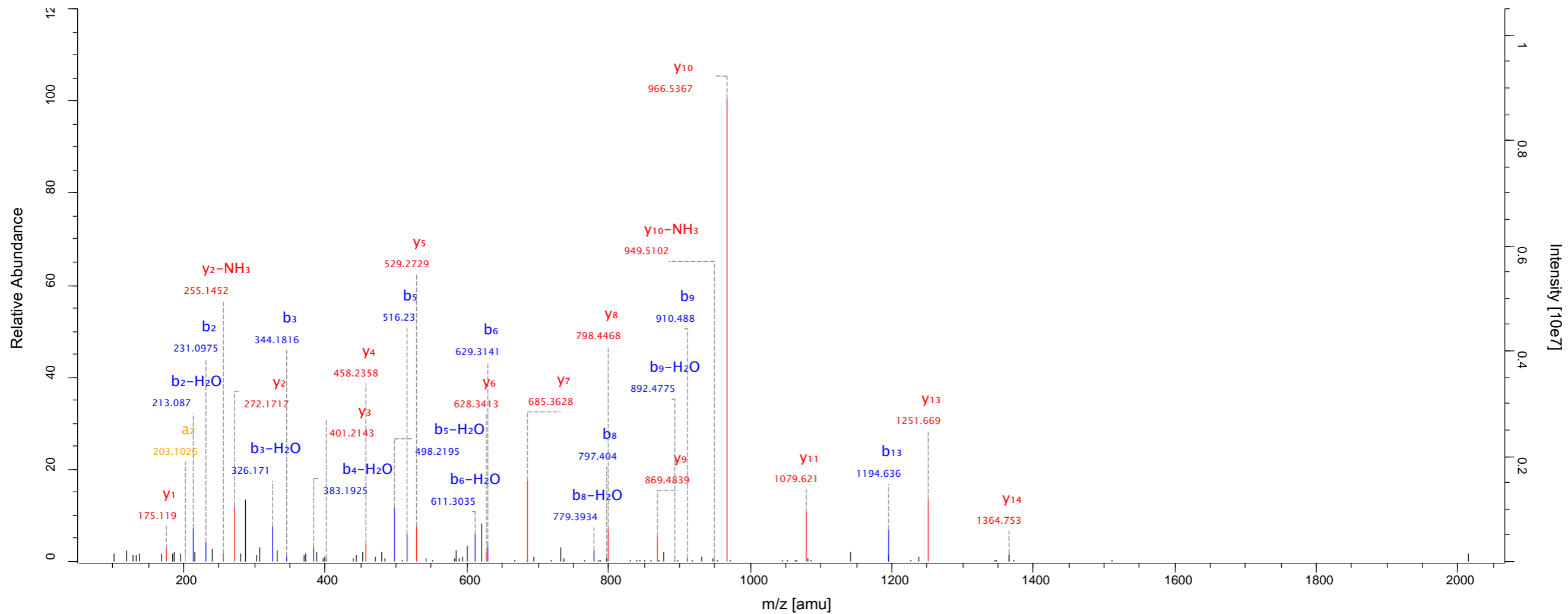
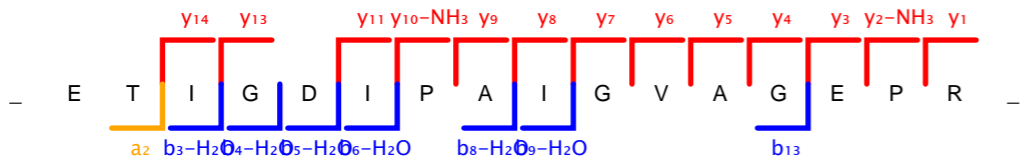
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 32471
Protein: ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
Peptide score: 120.12
Method: FTMS; HCD



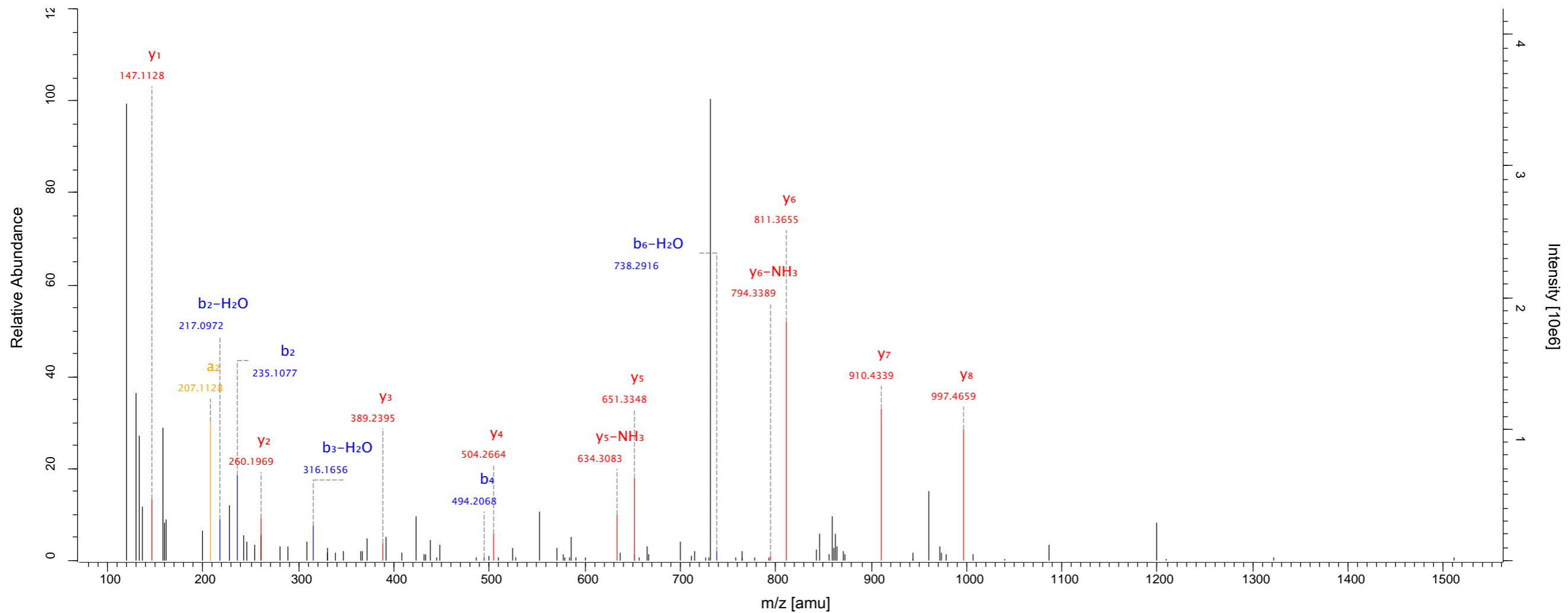
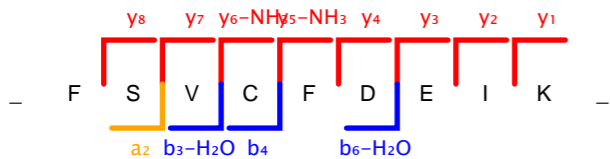
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
Scan number: 41028
Protein: ORFL271C.iORF1 - NC_006273.2[176607,176858]- <unknown description>
Peptide score: 94.69
Method: FTMS; HCD



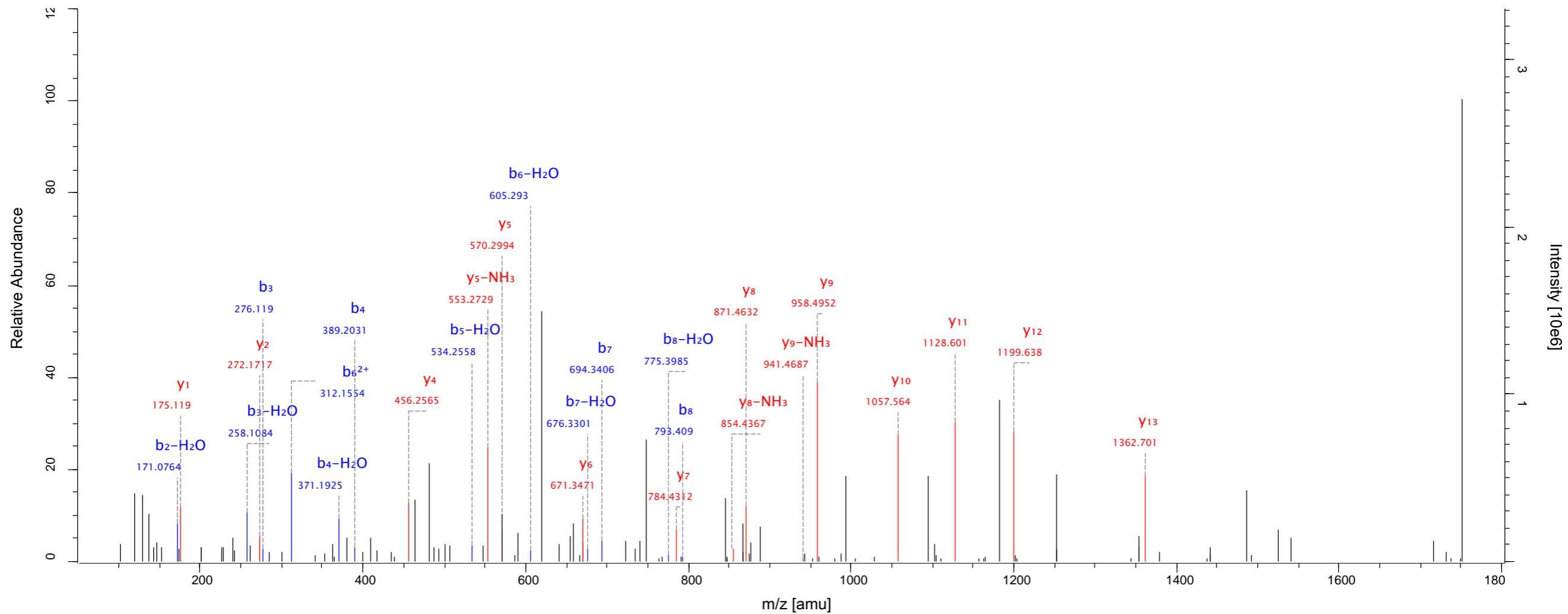
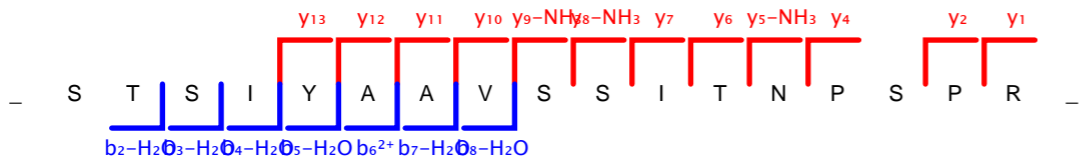
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
 Scan number: 47161
 Protein: ORFL73C - NC_006273.2[30581,30946]- <unknown description>
 Peptide score: 158.81
 Method: FTMS; HCD



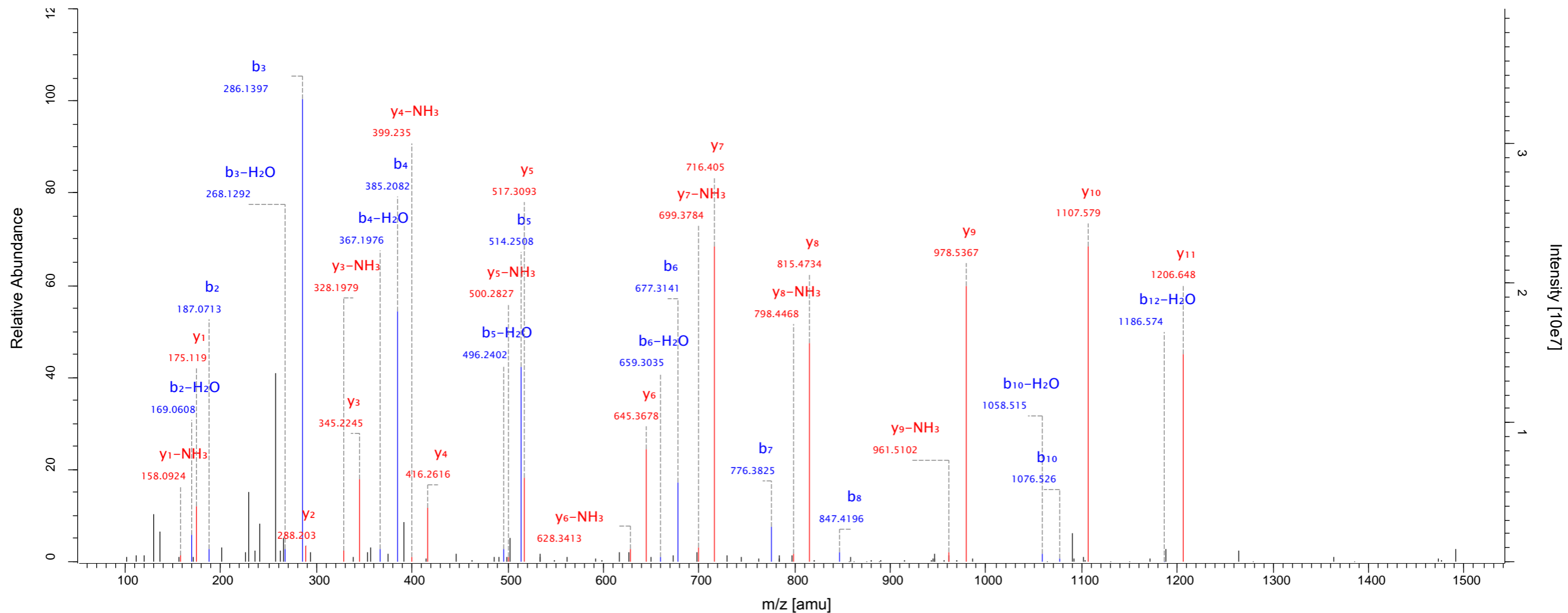
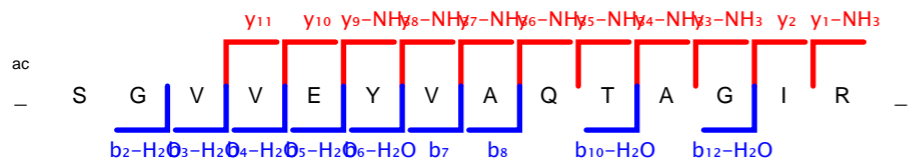
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 47216
Protein: ORFL206W + NC_006273.2[122632,122880]+ <unknown description>
Peptide score: 99.14
Method: FTMS; HCD



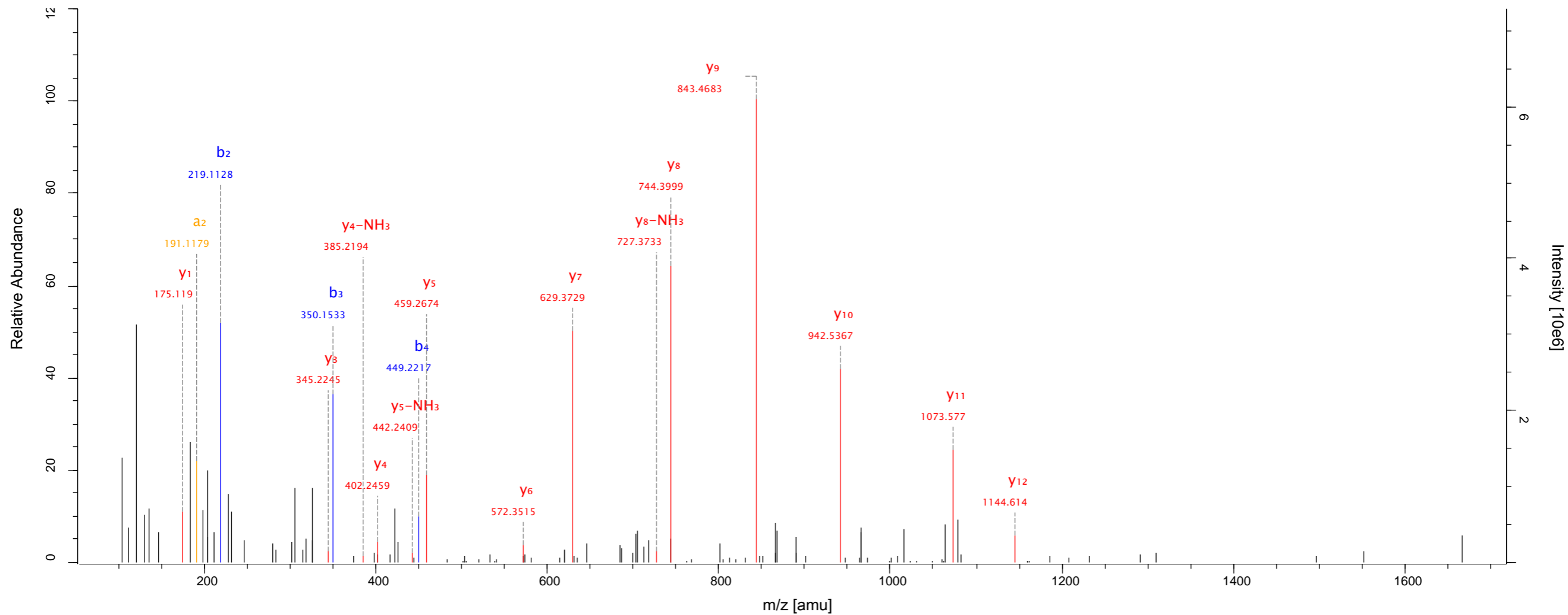
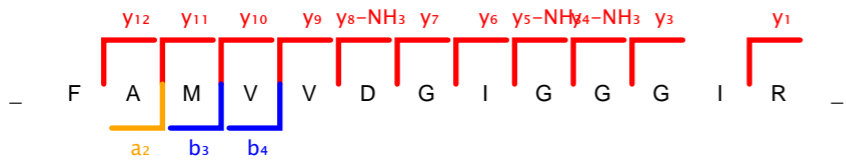
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
 Scan number: 47720
 Protein: ORFL253W.IORF3 + NC_006273.2[162059,162286]+ <unknown description>
 Peptide score: 98.68
 Method: FTMS; HCD



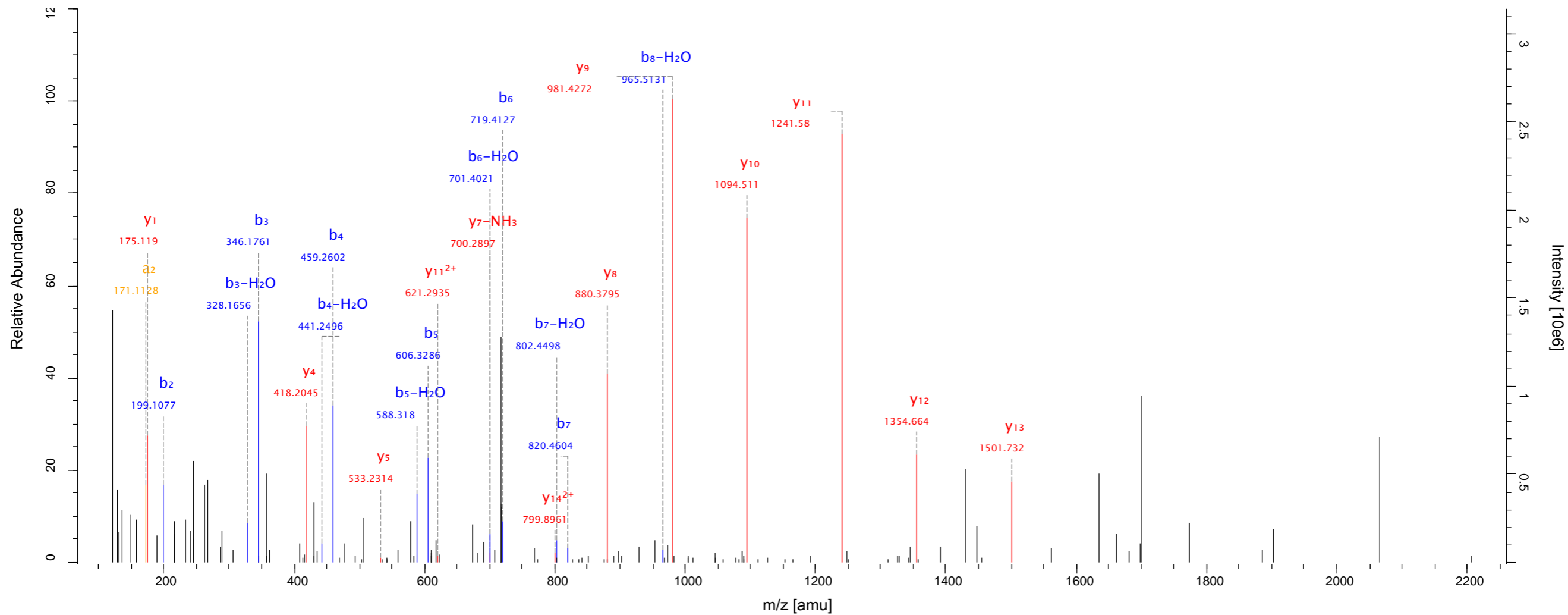
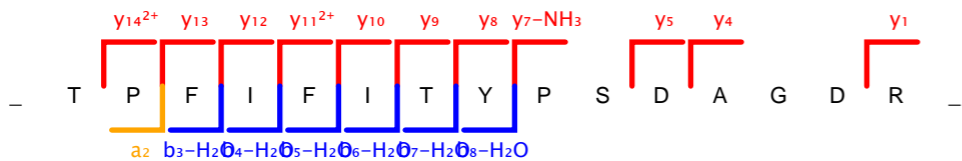
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
Scan number: 47722
Protein: ORFL211W + NC_006273.2[125859,126083]+ <unknown description>
Peptide score: 190.78
Method: FTMS; HCD



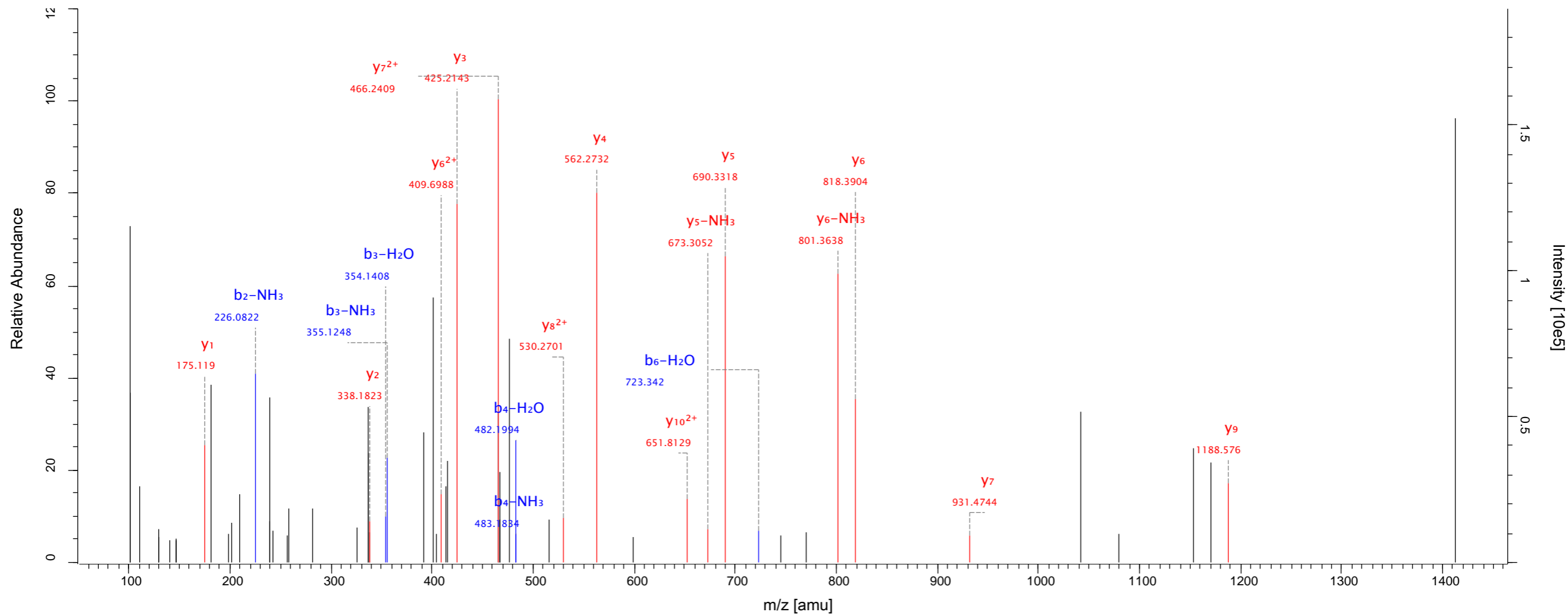
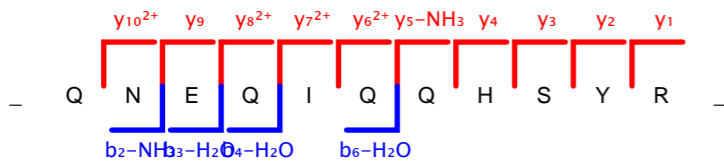
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
Scan number: 48342
Protein: ORFL211W + NC_006273.2[125859,126083]+ <unknown description>
Peptide score: 77.59
Method: FTMS; HCD



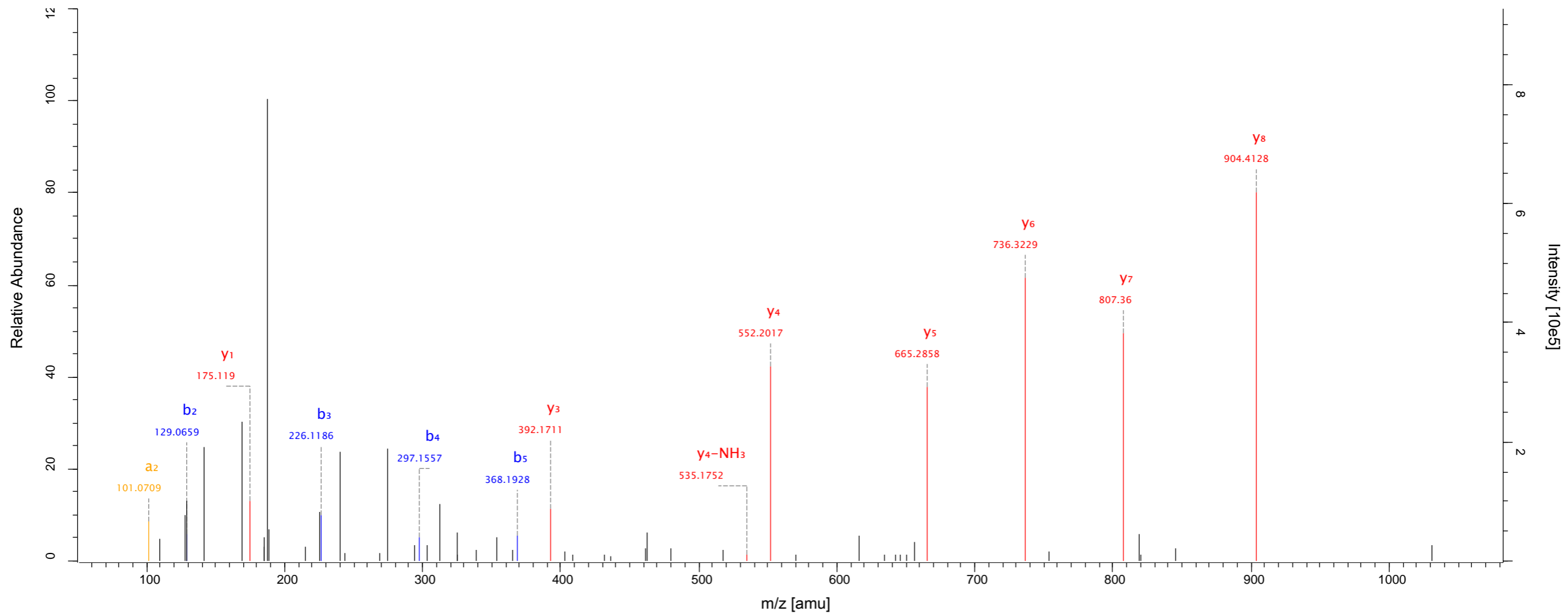
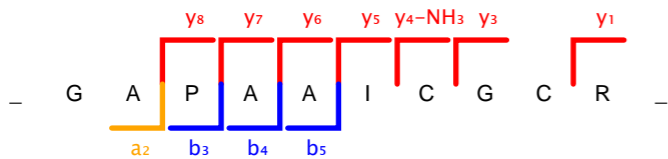
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_34
 Scan number: 48586
 Protein: ORFL174C.iORF1 - NC_006273.2[106302,106568]- <unknown description>
 Peptide score: 120.14
 Method: FTMS; HCD



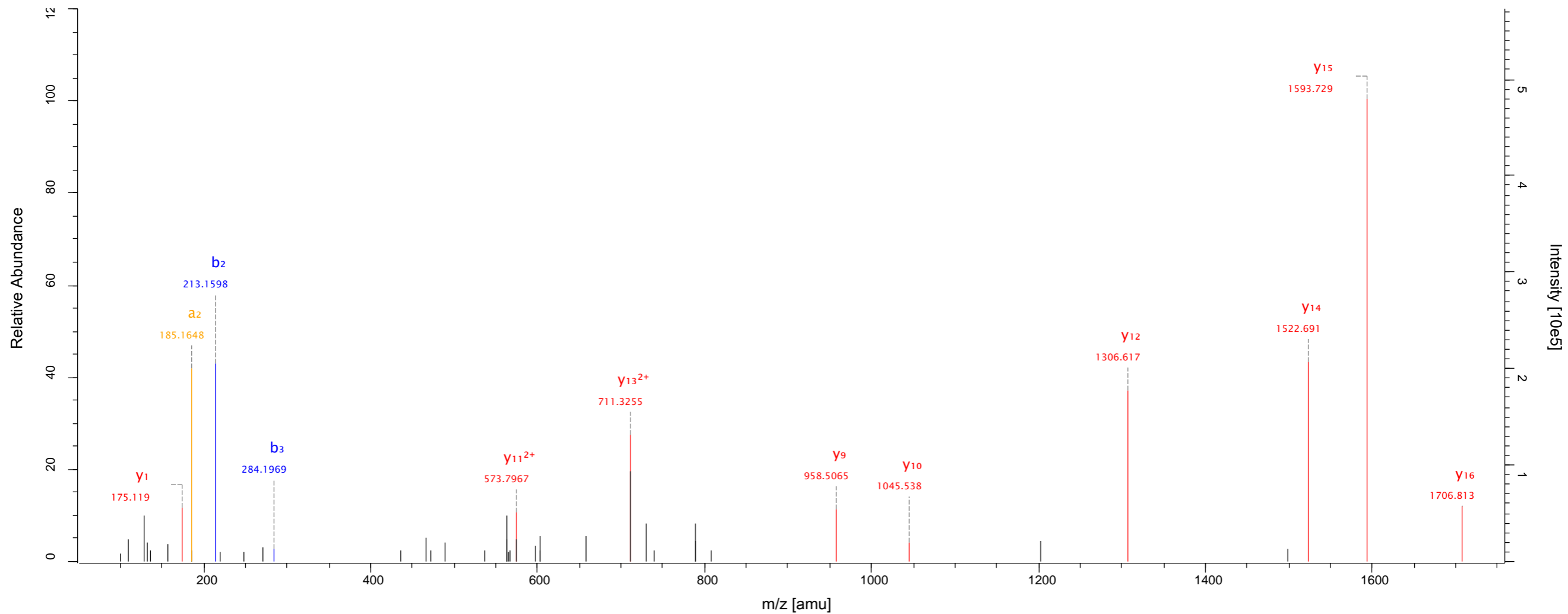
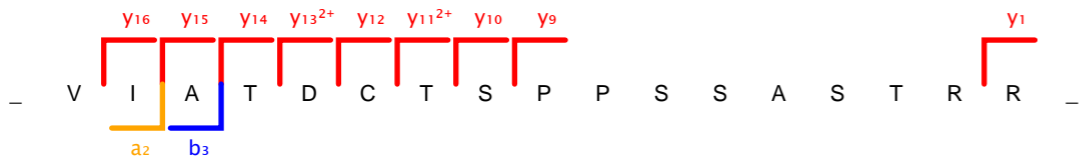
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_34
 Scan number: 6089
 Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
 Peptide score: 91.58
 Method: FTMS; HCD



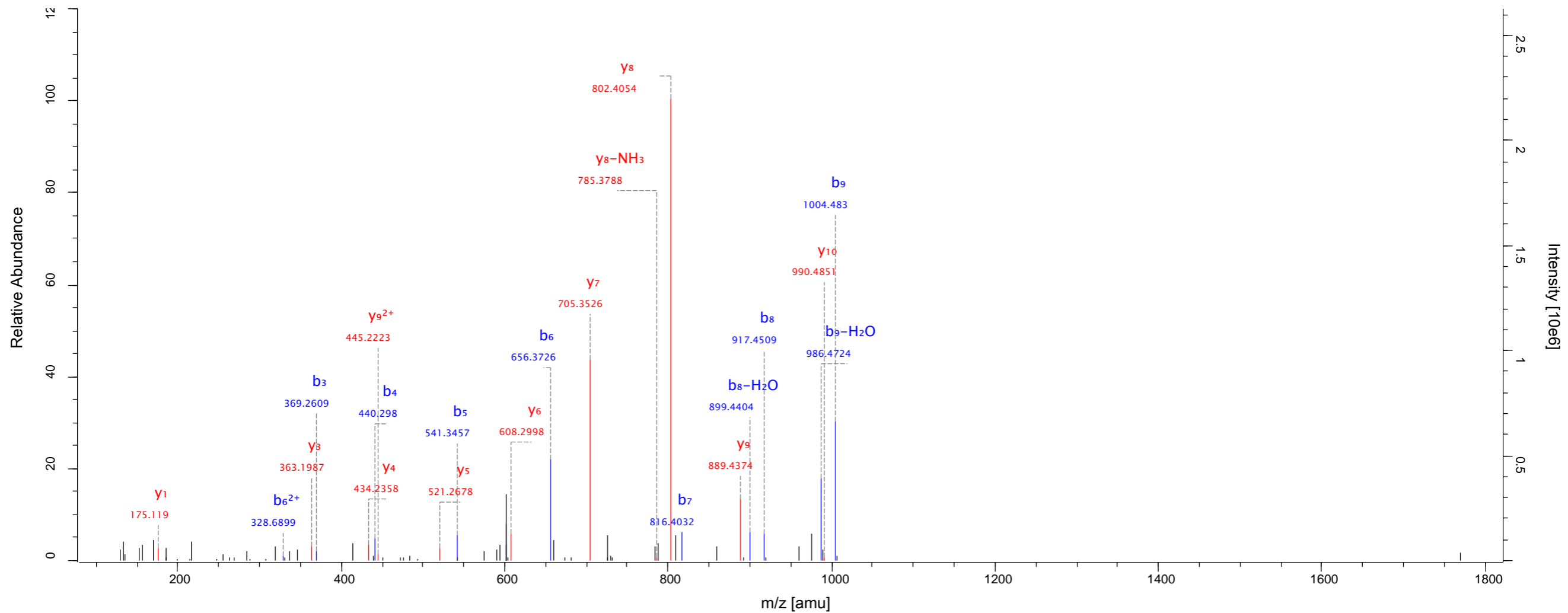
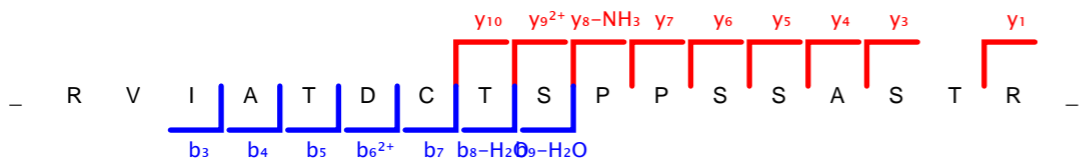
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_34
Scan number: 8955
Protein: ORFL73C - NC_006273.2[30581,30946]- <unknown description>
Peptide score: 74.92
Method: FTMS; HCD



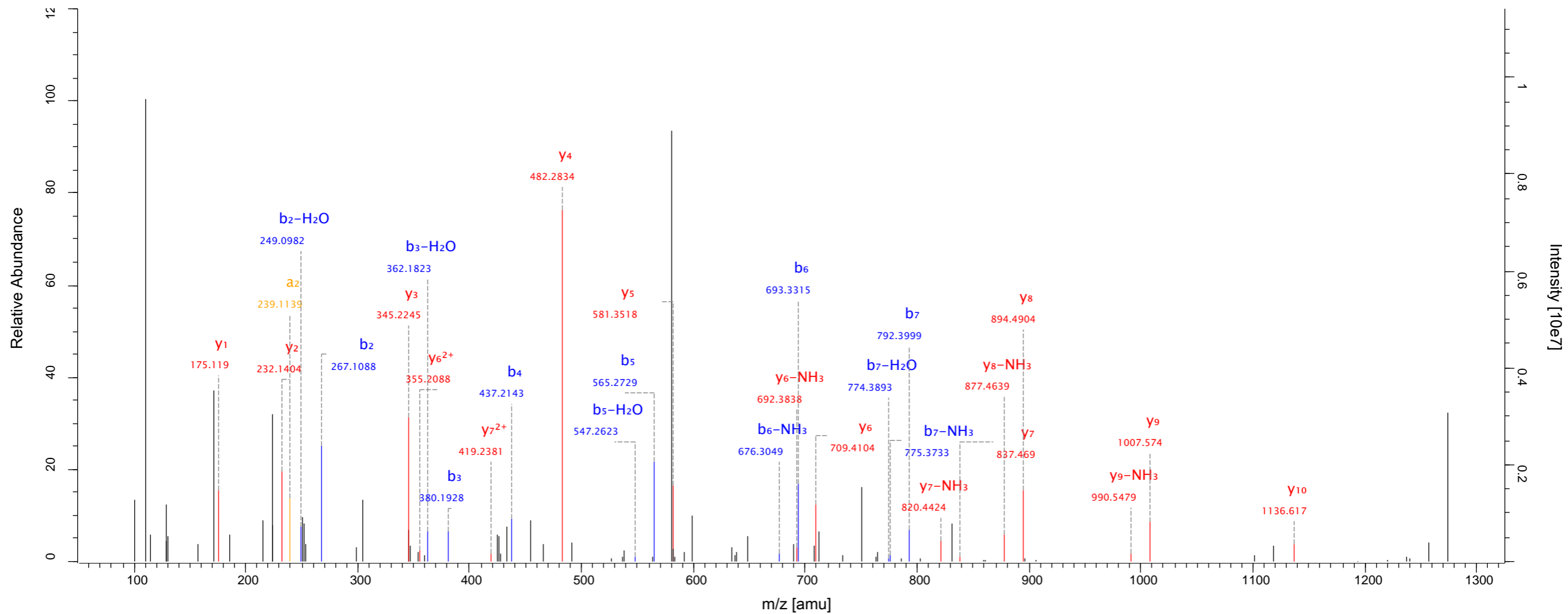
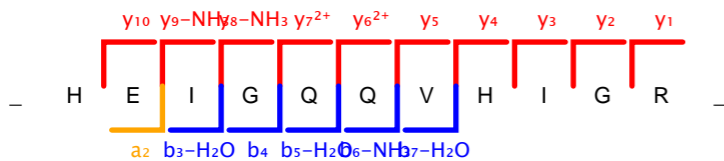
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 10146
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 74.68
Method: FTMS; HCD



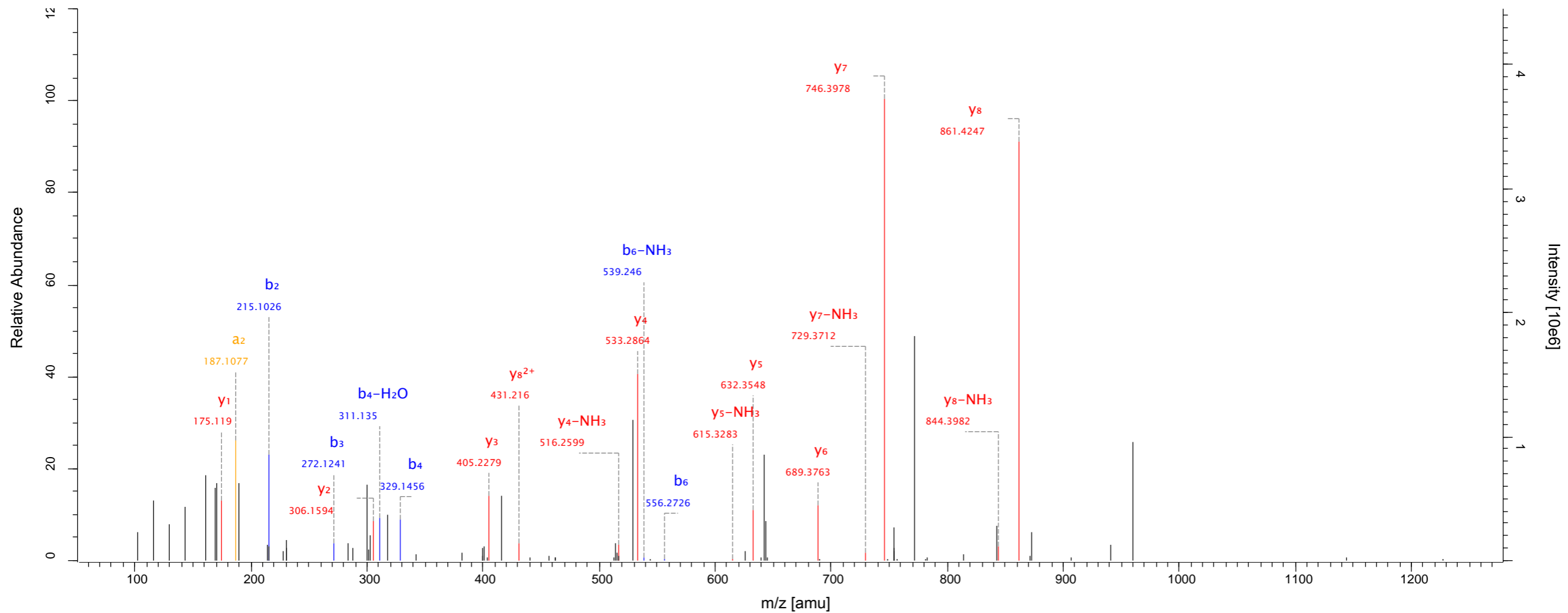
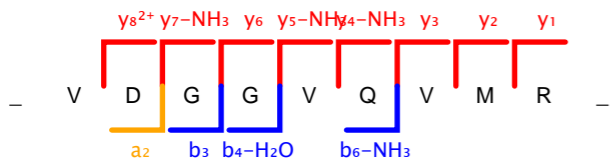
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 10767
 Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
 Peptide score: 75.45
 Method: FTMS; HCD



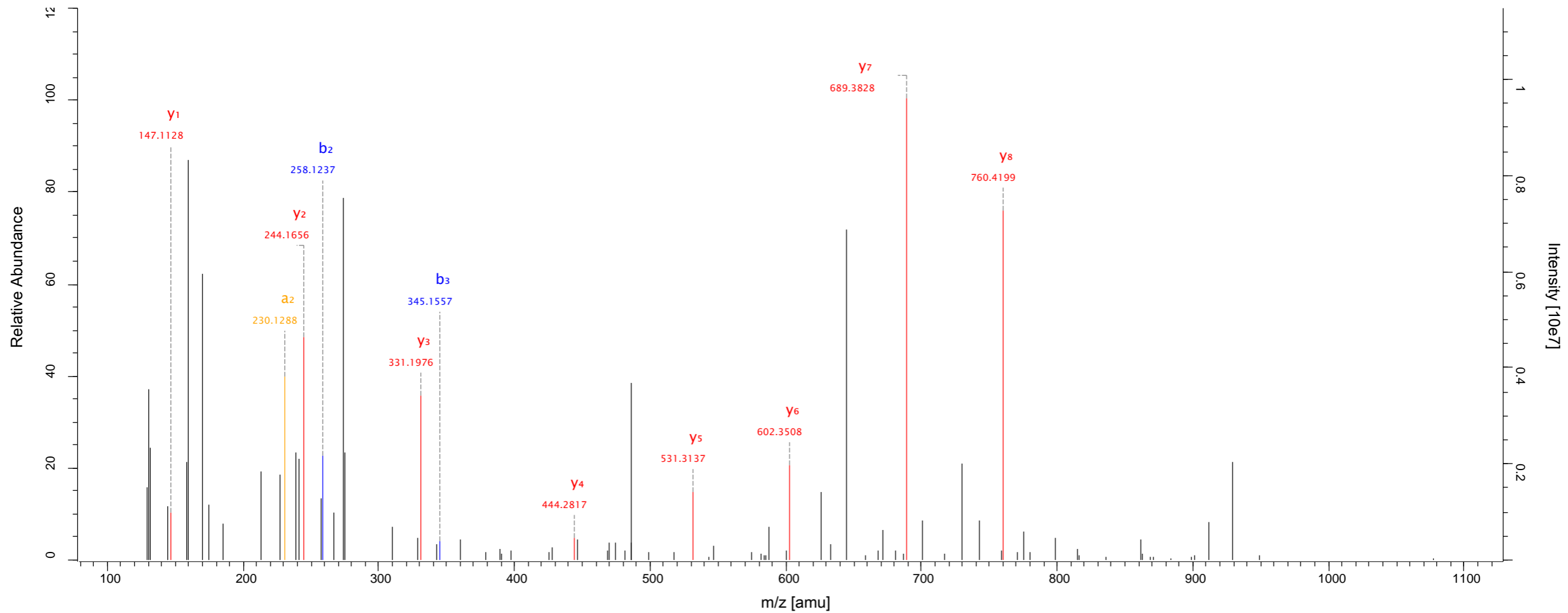
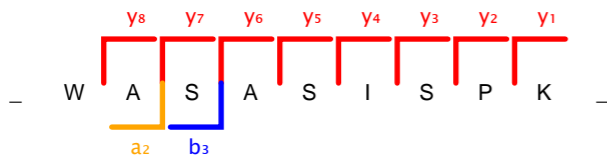
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 11021
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 165.18
 Method: FTMS; HCD



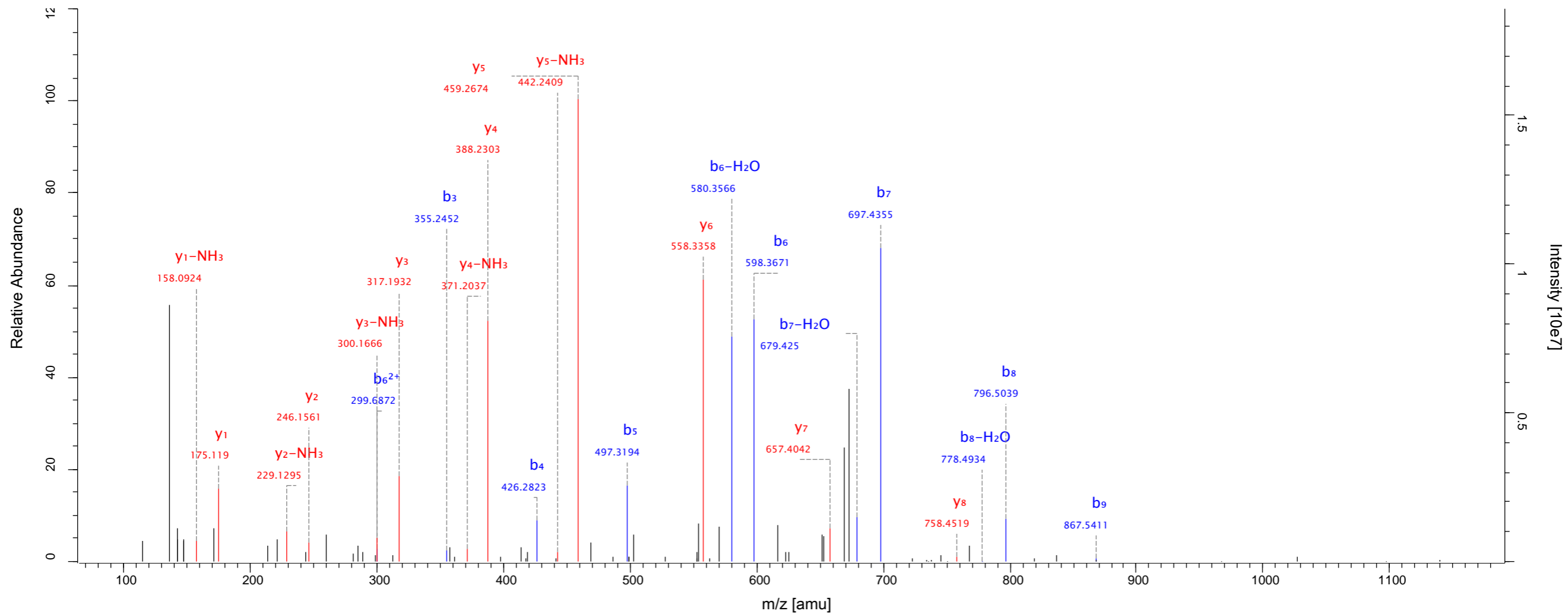
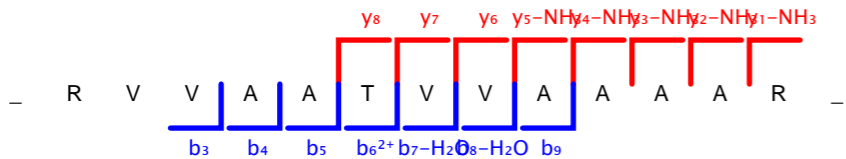
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_35
 Scan number: 12108
 Protein: ORFL119C - NC_006273.2[63363,63800]- <unknown description>
 Peptide score: 117.81
 Method: FTMS; HCD



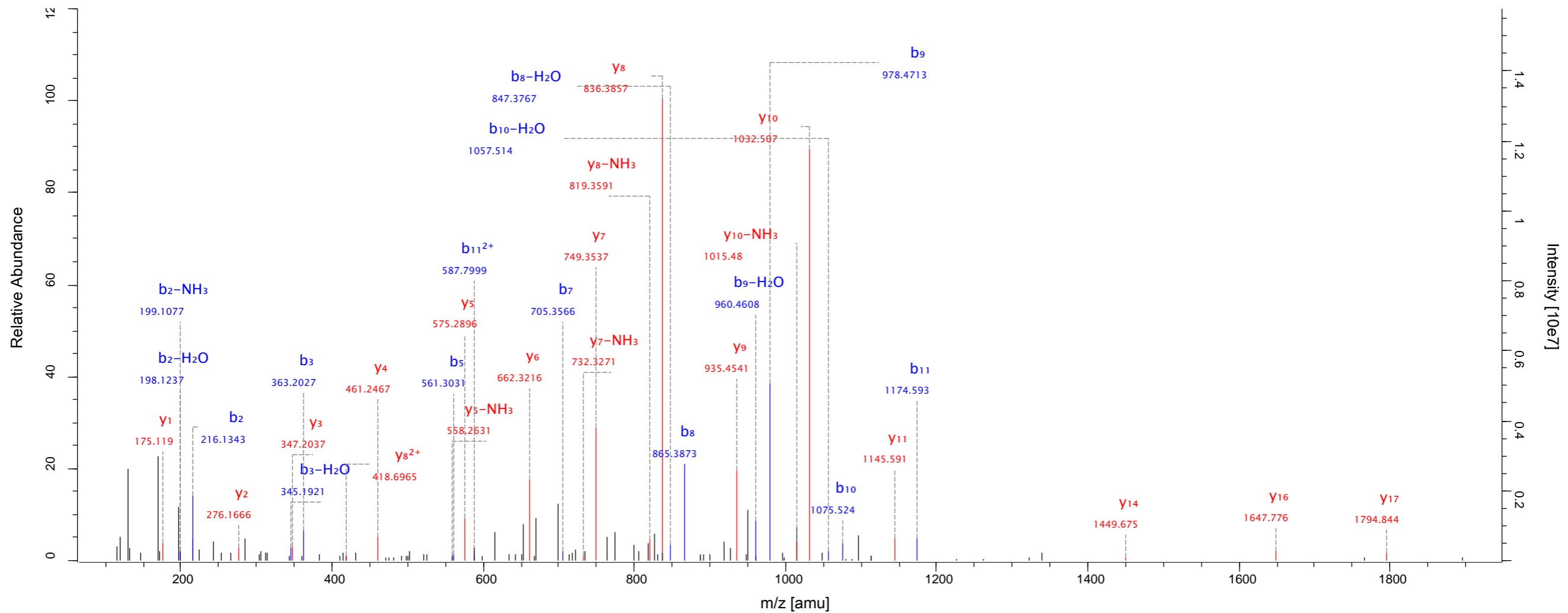
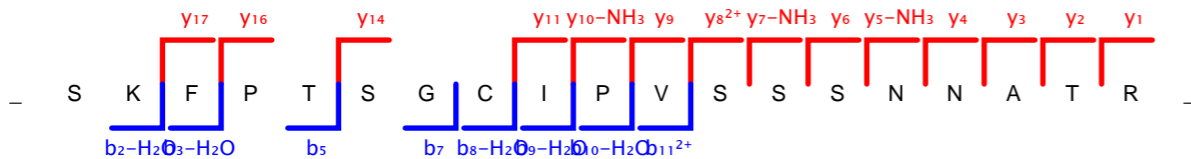
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 14501
Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
Peptide score: 56.56
Method: FTMS; HCD



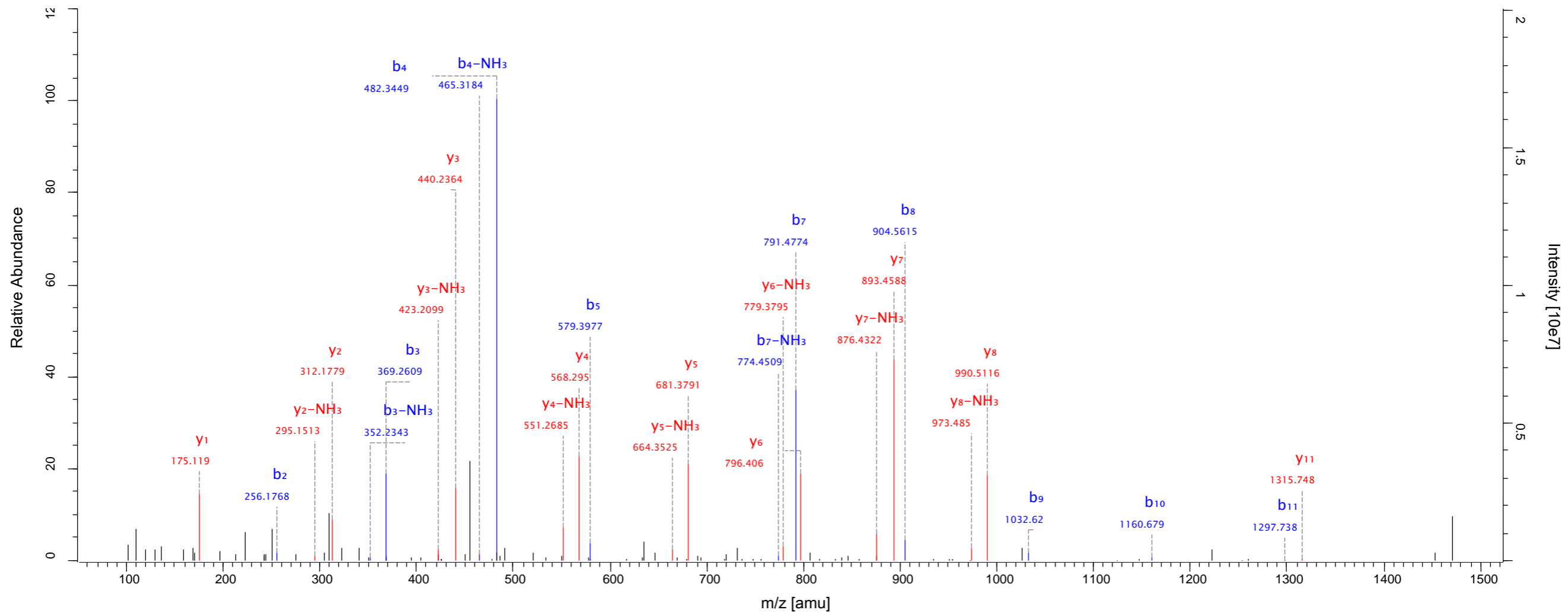
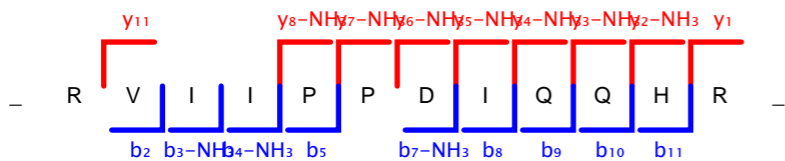
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 18060
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 127.56
 Method: FTMS; HCD



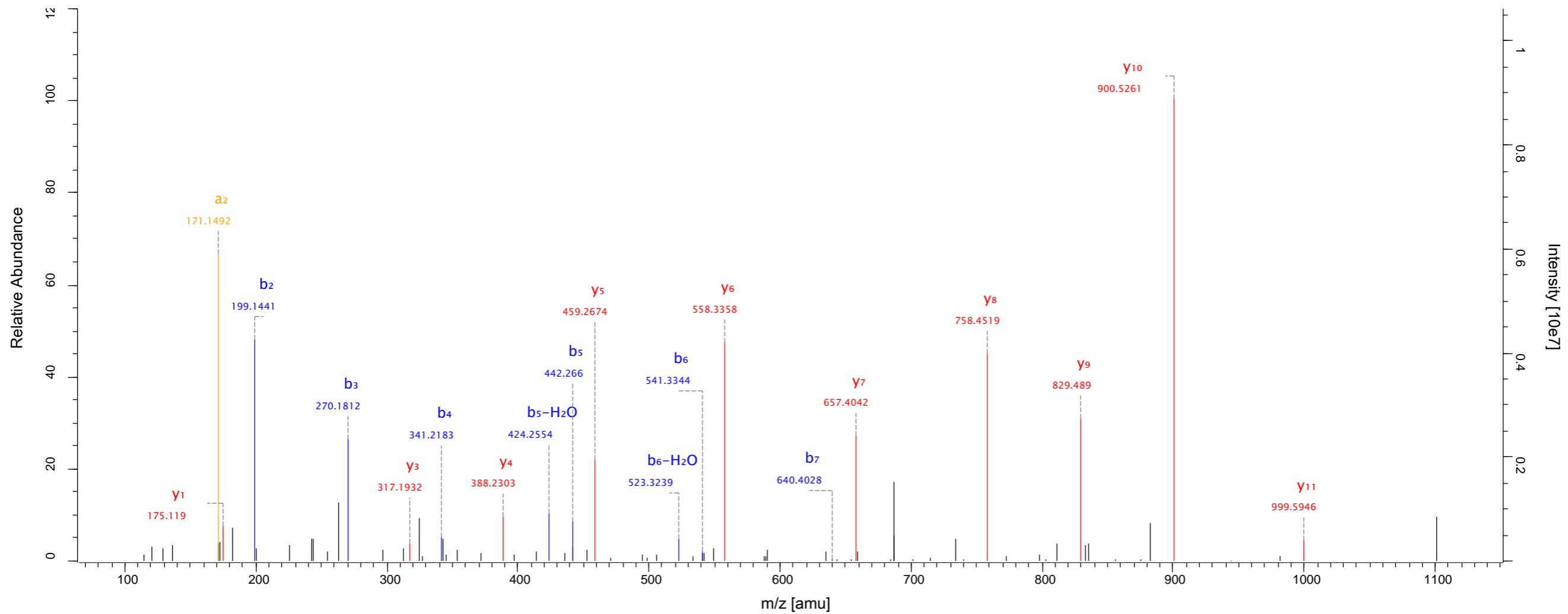
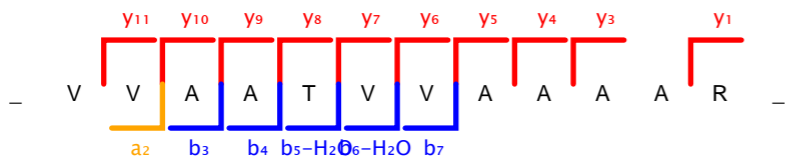
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 19430
 Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
 Peptide score: 132.14
 Method: FTMS; HCD



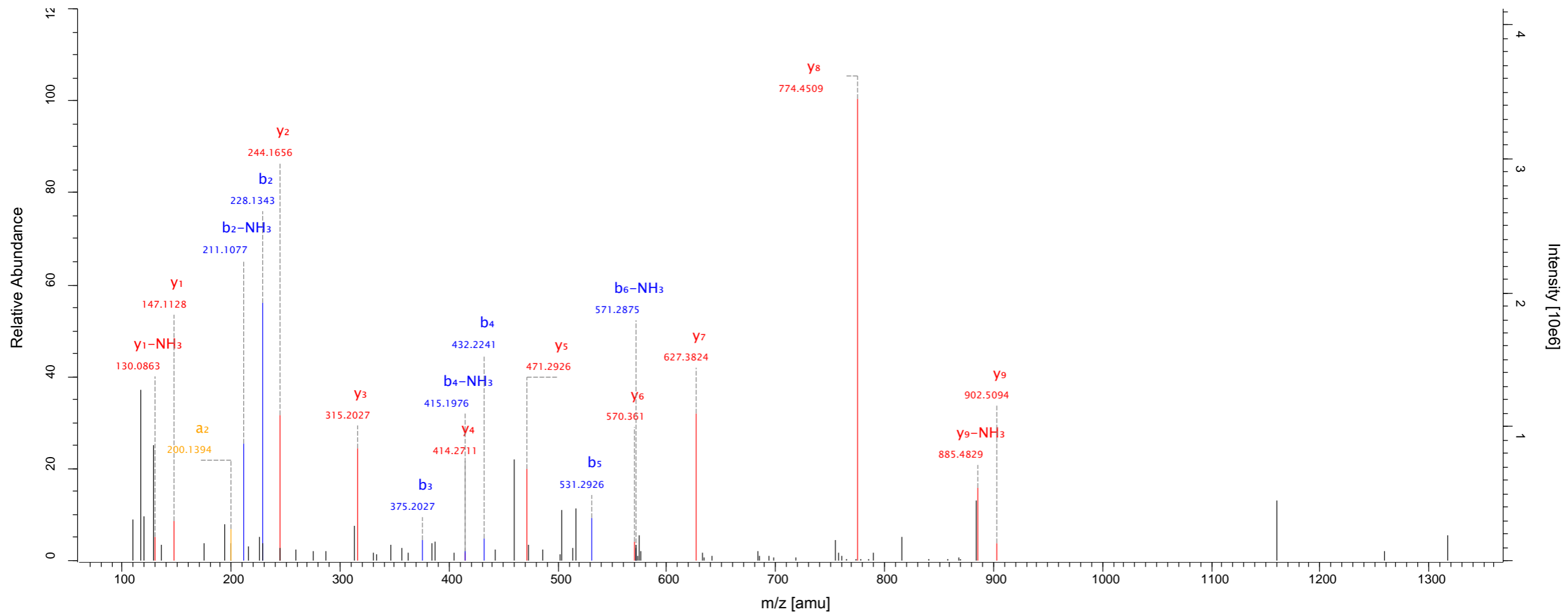
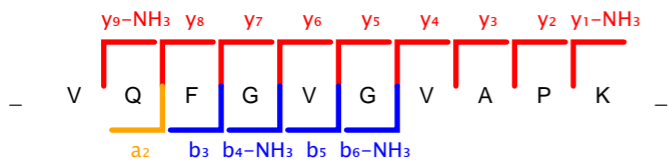
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 20375
 Protein: ORFL106C - NC_006273.2[53935,54294]- <unknown description>
 Peptide score: 210.91
 Method: FTMS; HCD



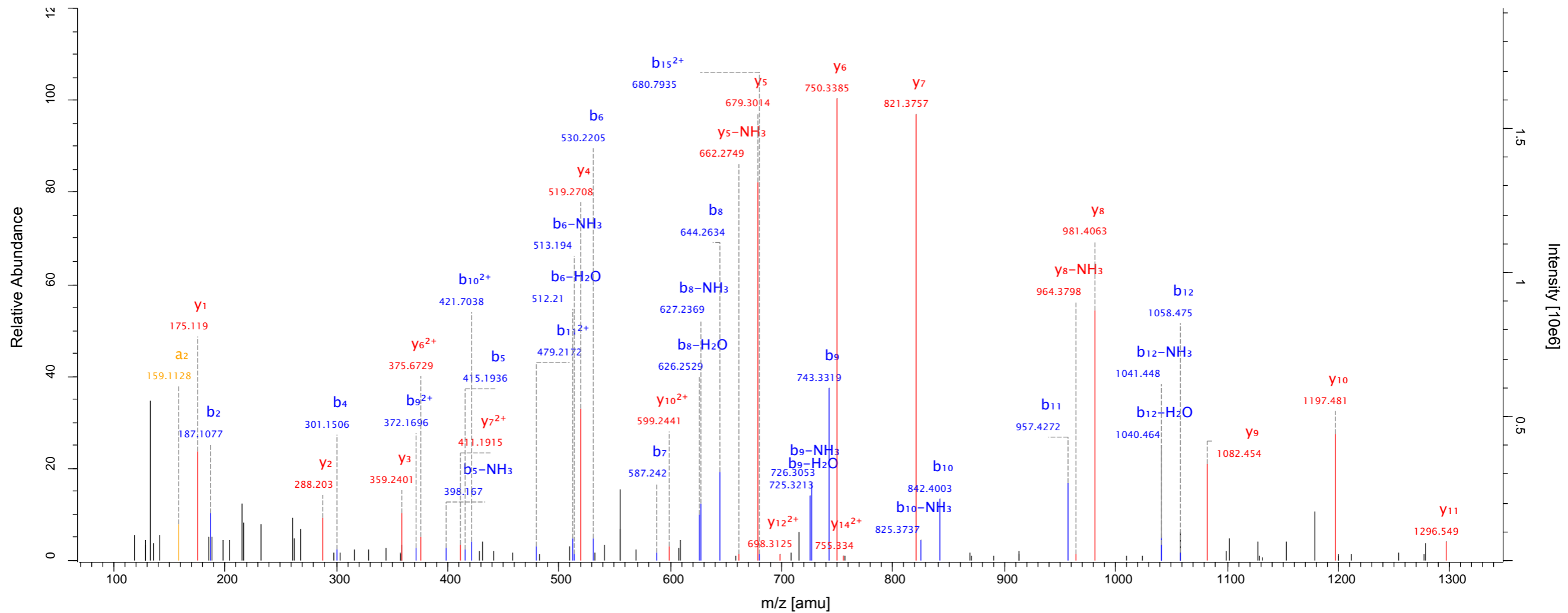
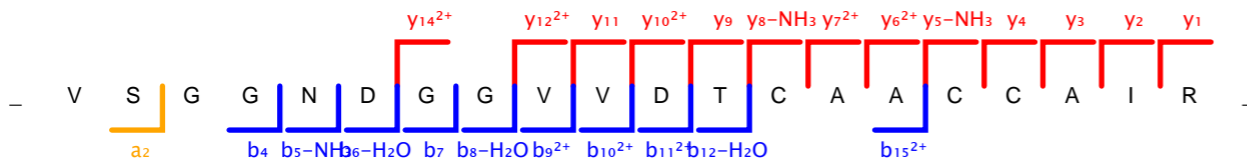
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 20786
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 125.74
Method: FTMS; HCD



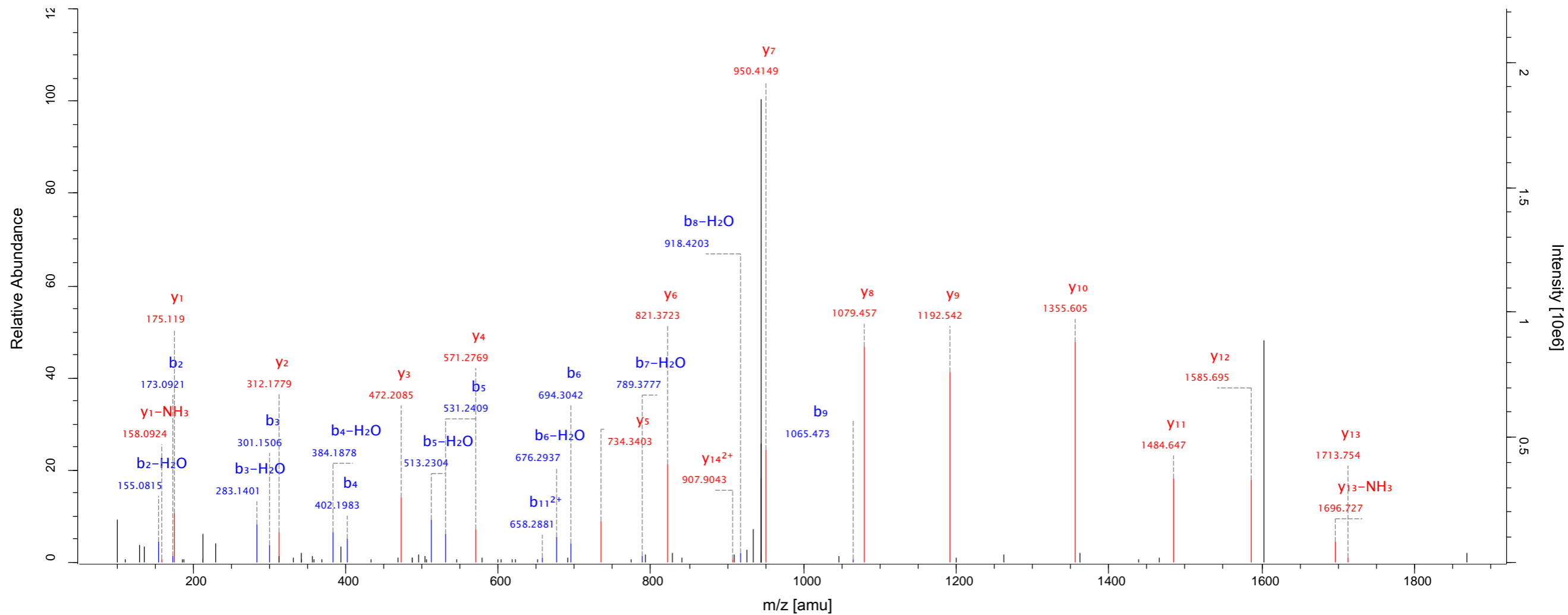
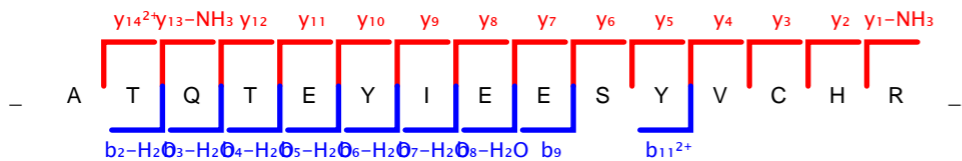
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 21254
Protein: ORFL118C - NC_006273.2[61896,62396]- <unknown description>
Peptide score: 102.06
Method: FTMS; HCD



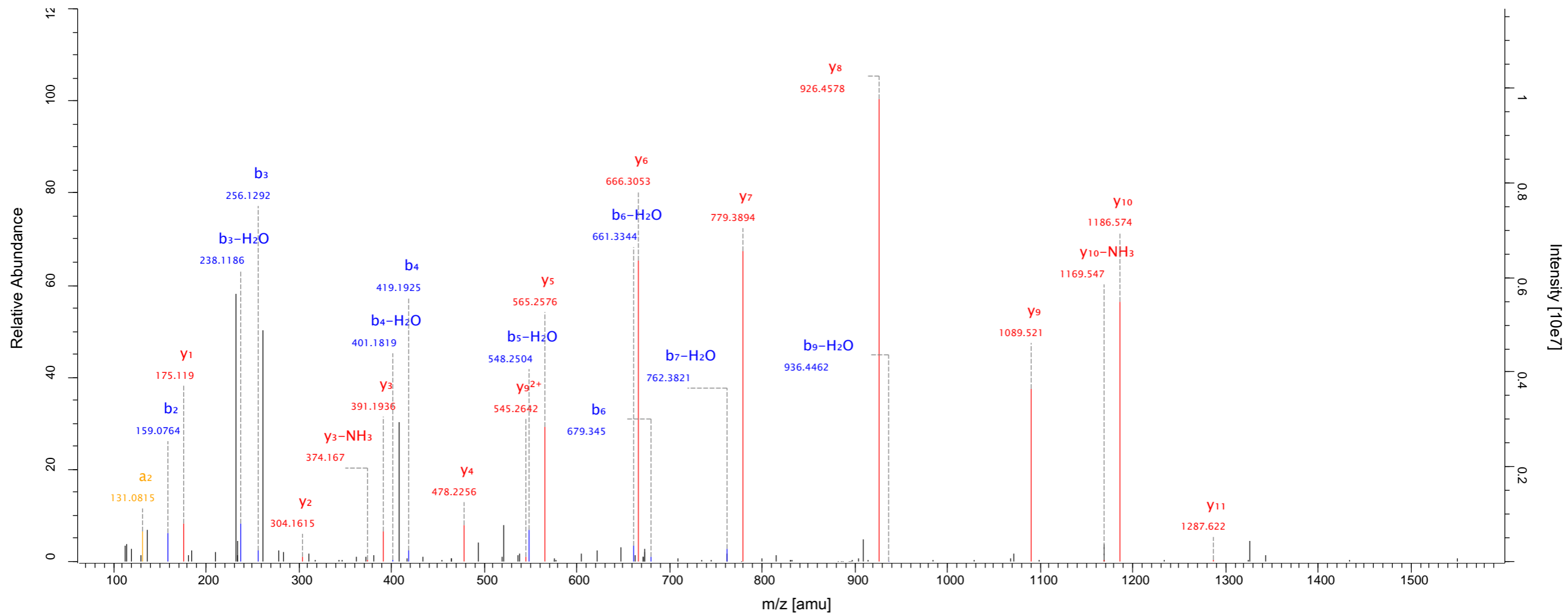
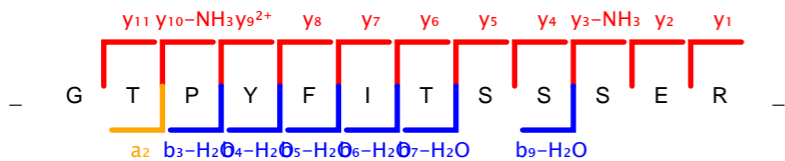
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 21604
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 197.51
 Method: FTMS; HCD



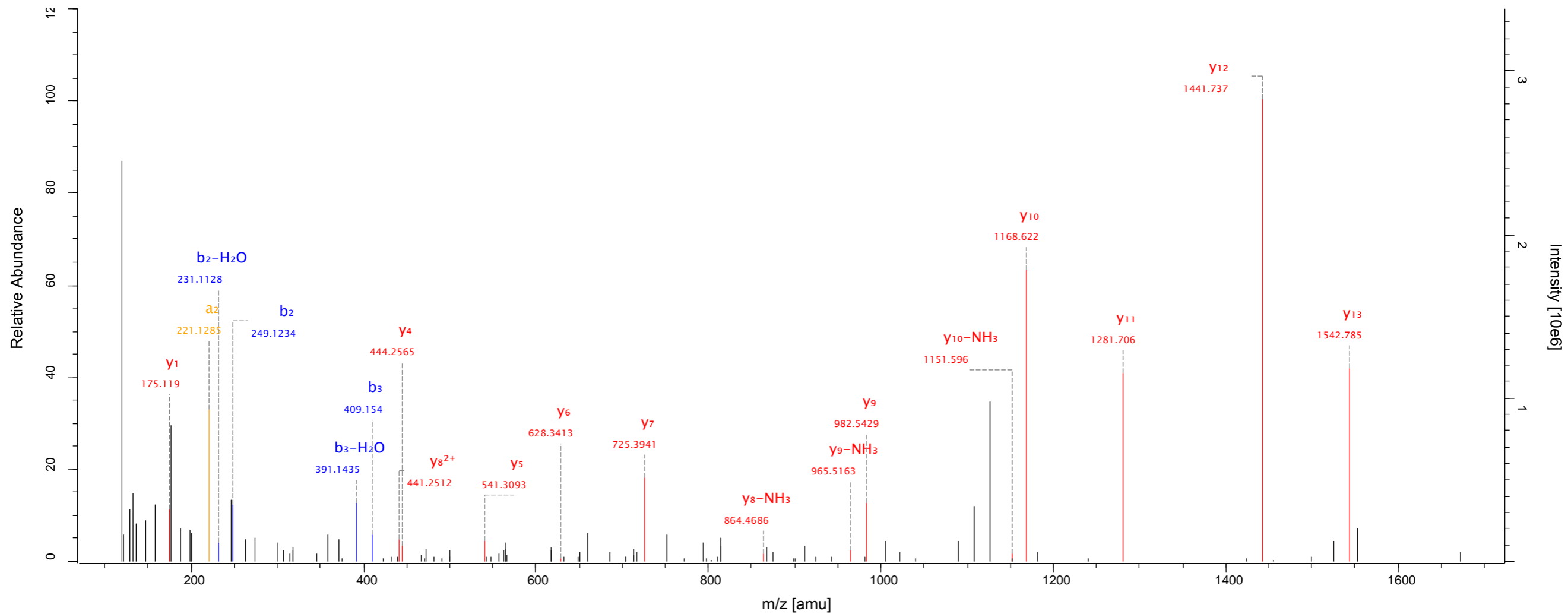
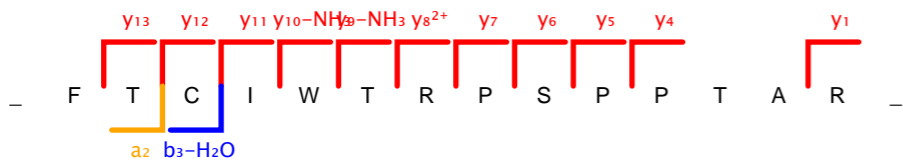
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_35
 Scan number: 22020
 Protein: ORFL294W.iORF1 + NC_006273.2[183230,193441]+ <unknown description>; ORFL294W + NC_006273.2[183230,193441]+ <unknown description>
 Peptide score: 198.33
 Method: FTMS; HCD



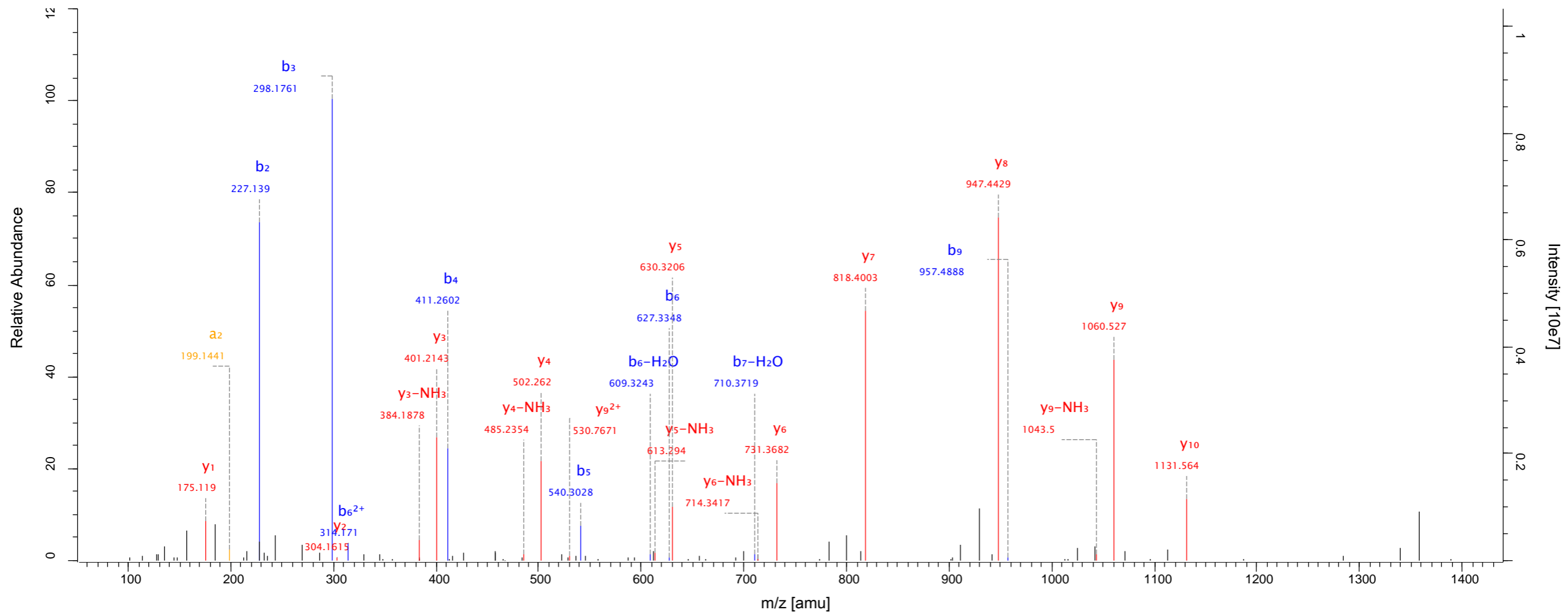
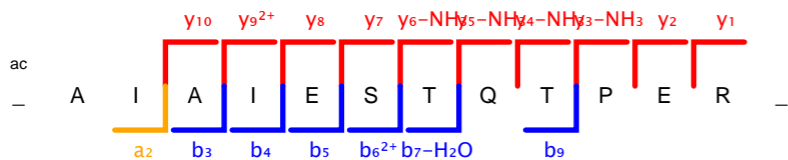
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 24092
Protein: ORFL142W + NC_006273.2[78726,79031]+ <unknown description>
Peptide score: 122.51
Method: FTMS; HCD



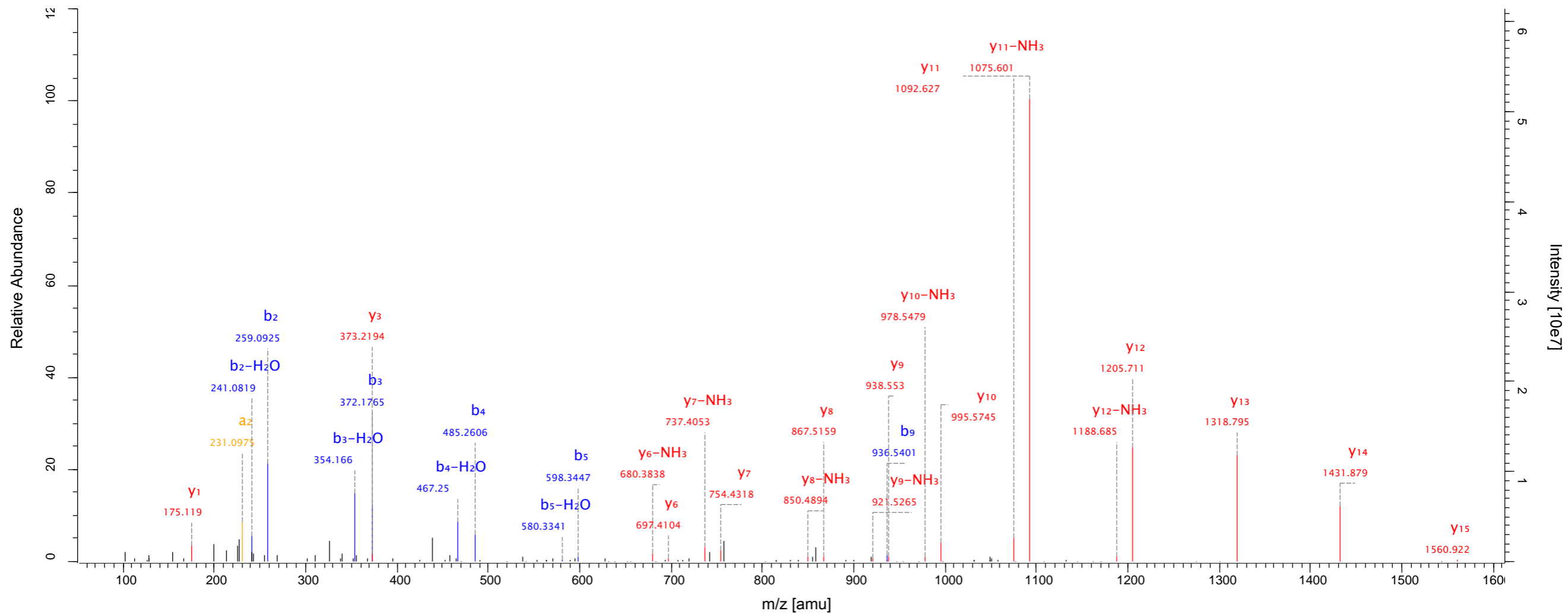
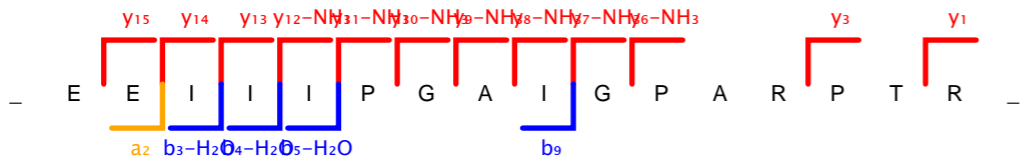
Raw file: 20111212_EXQ5_AnMi_QC_HCMVpl1_35
Scan number: 26775
Protein: ORFL224C.iORF1 - NC_006273.2[134354,134734]- <unknown description>
Peptide score: 93.01
Method: FTMS; HCD



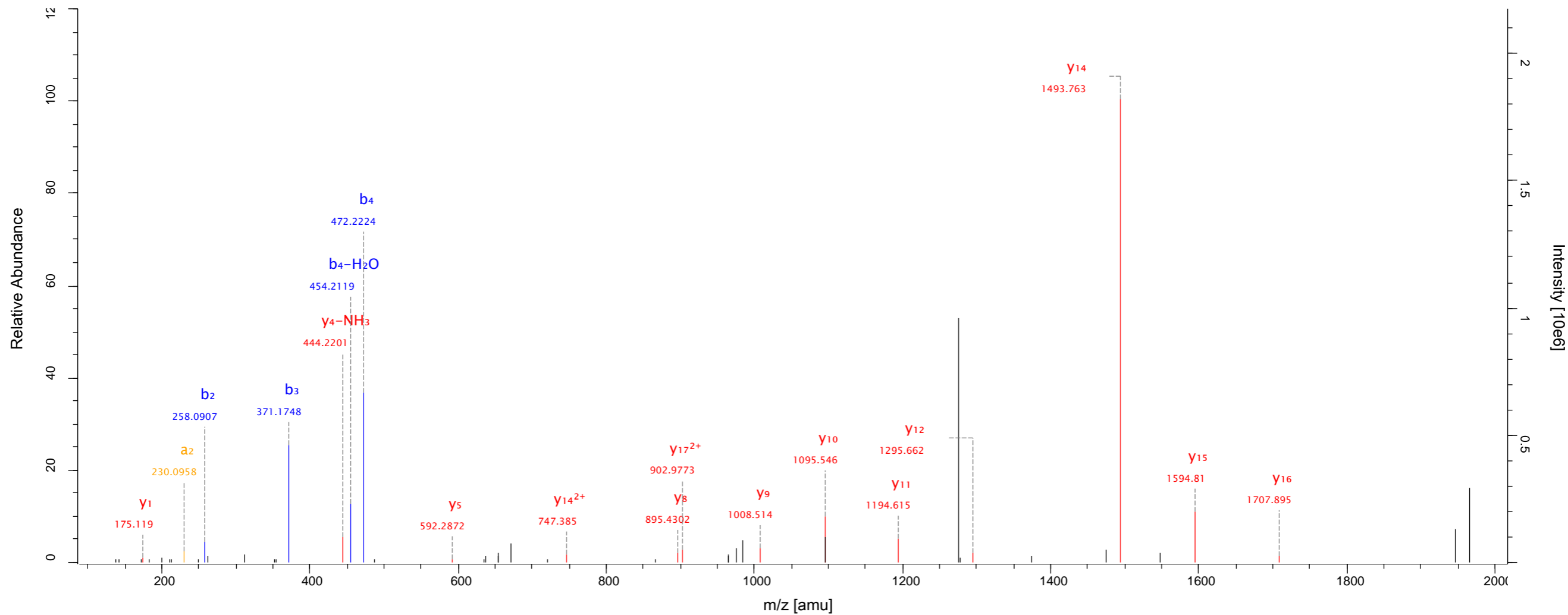
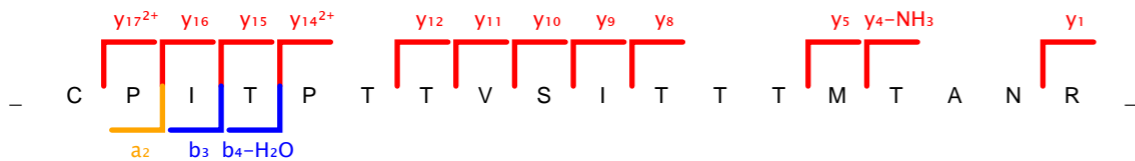
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 28409
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 160.77
 Method: FTMS; HCD



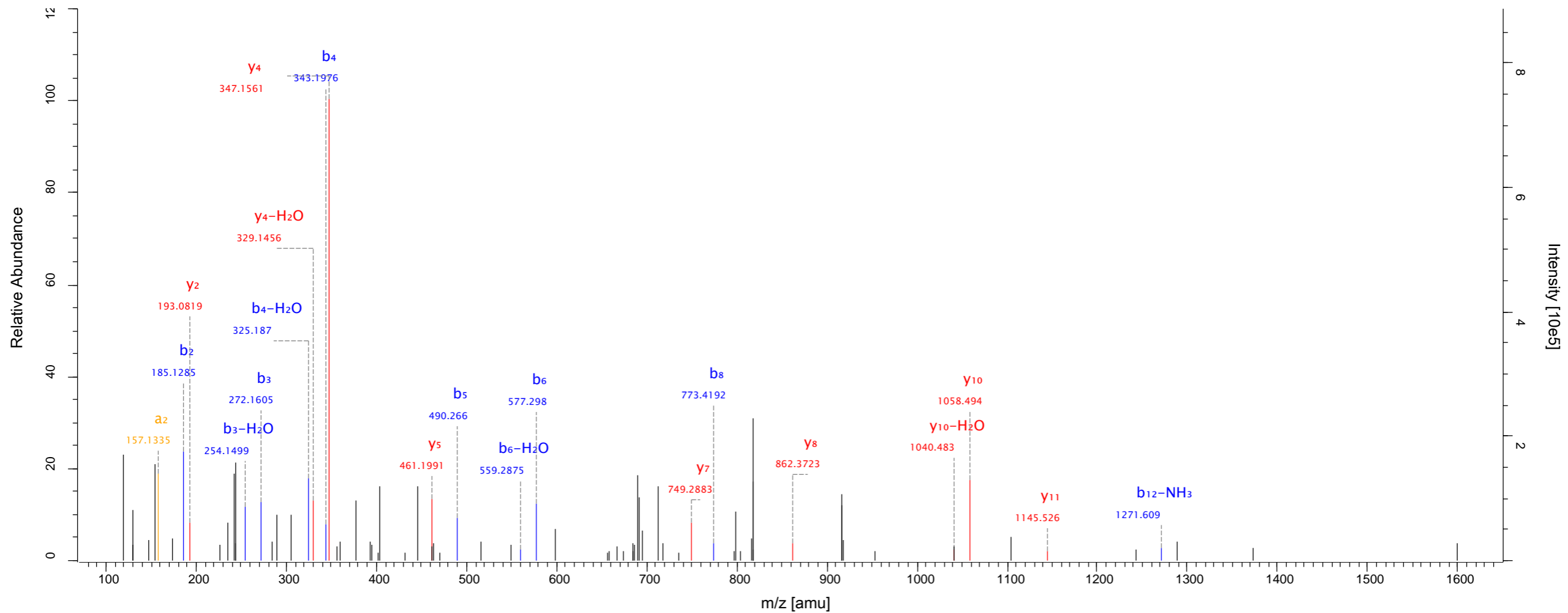
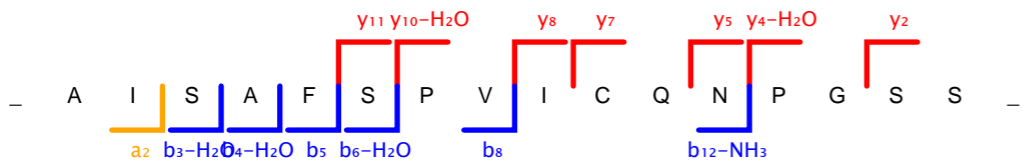
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 33784
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 198.14
 Method: FTMS; HCD



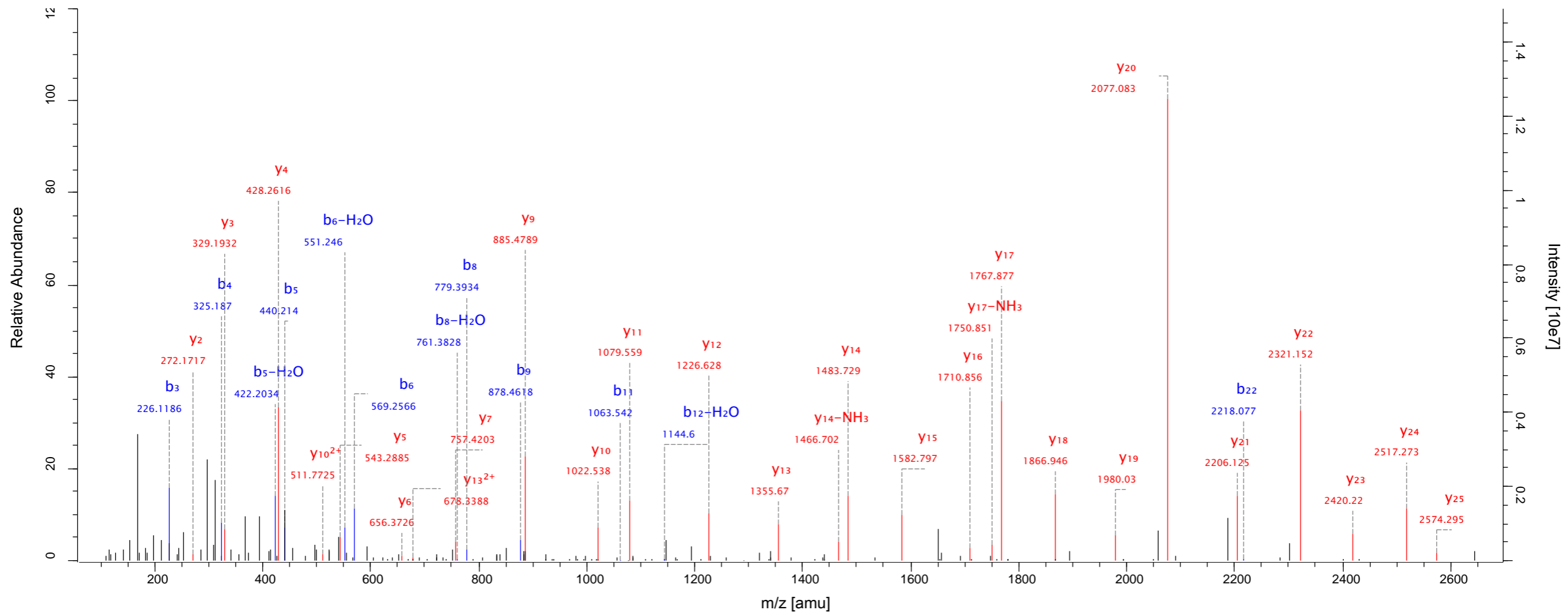
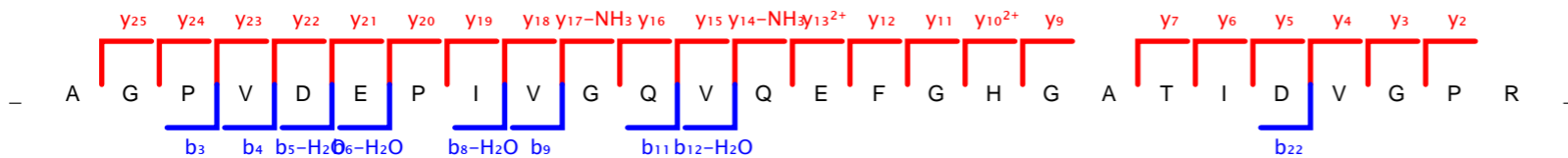
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_35
Scan number: 34491
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 123.38
Method: FTMS; HCD



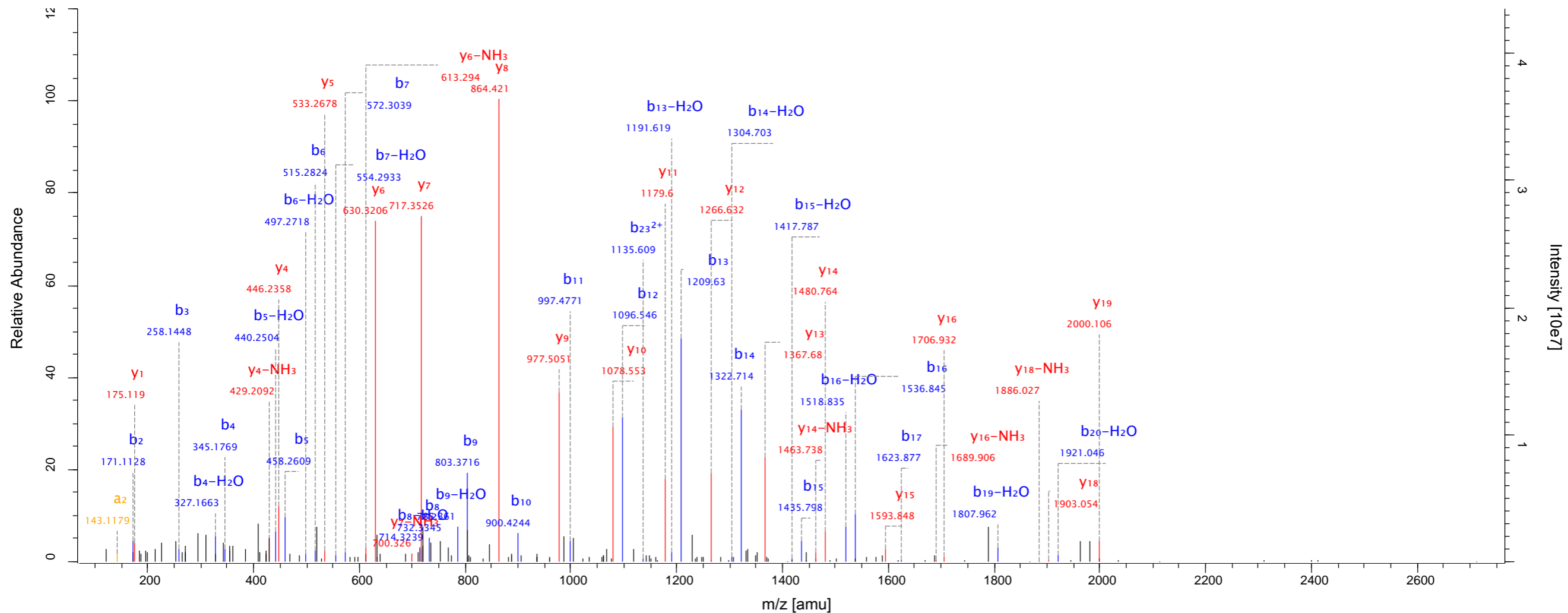
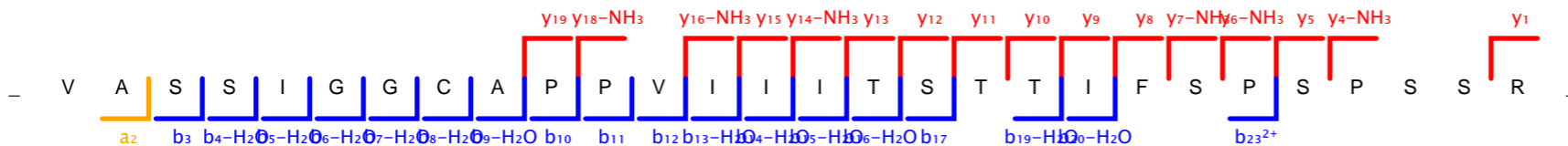
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 36886
Protein: ORFL174C.iORF2 - NC_006273.2[106396,106770]- <unknown description>
Peptide score: 86.76
Method: FTMS; HCD



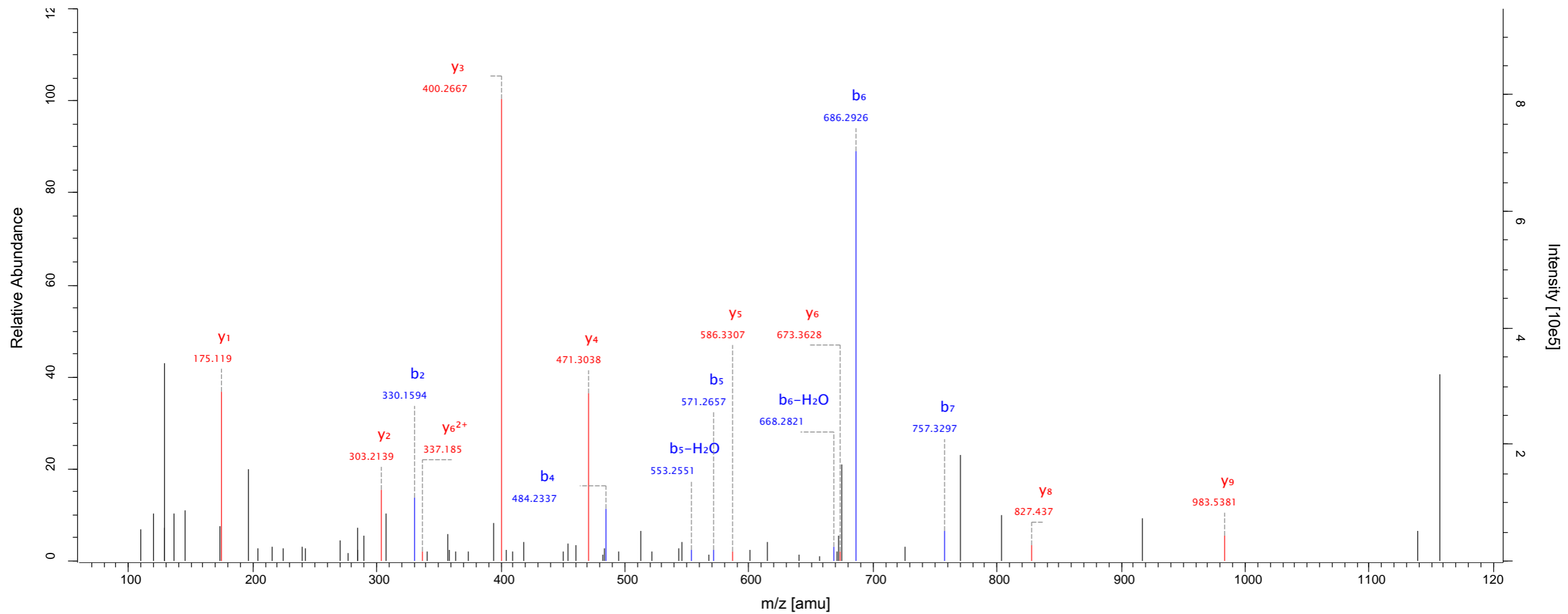
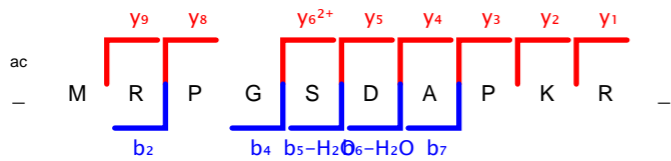
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 47902
Protein: ORFL118C - NC_006273.2[61896,62396]- <unknown description>
Peptide score: 198.31
Method: FTMS; HCD



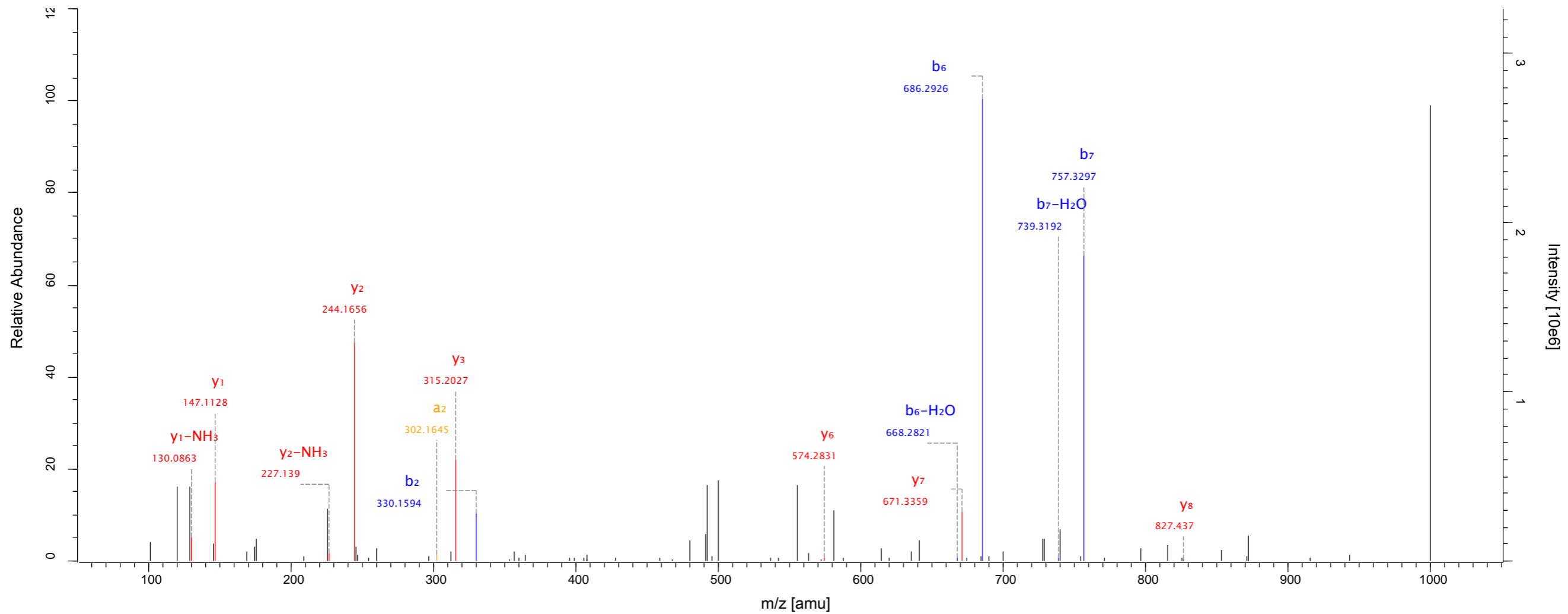
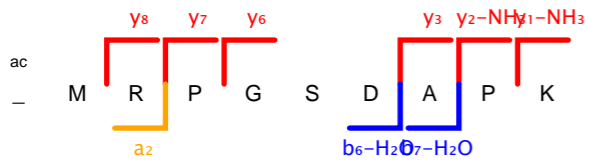
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 49294
 Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
 Peptide score: 220.88
 Method: FTMS; HCD



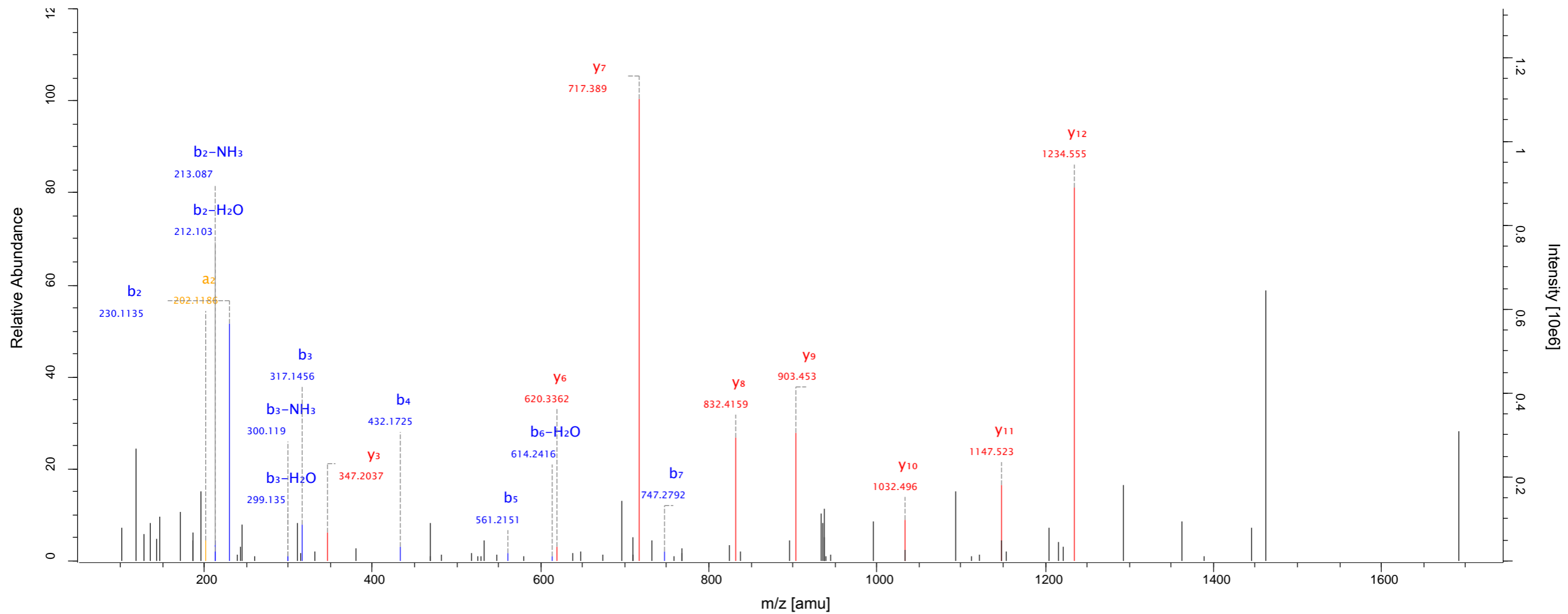
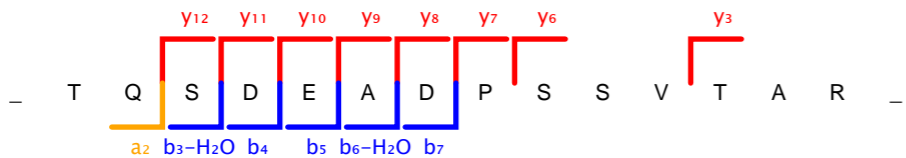
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_35
Scan number: 6724
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 71.22
Method: FTMS; HCD



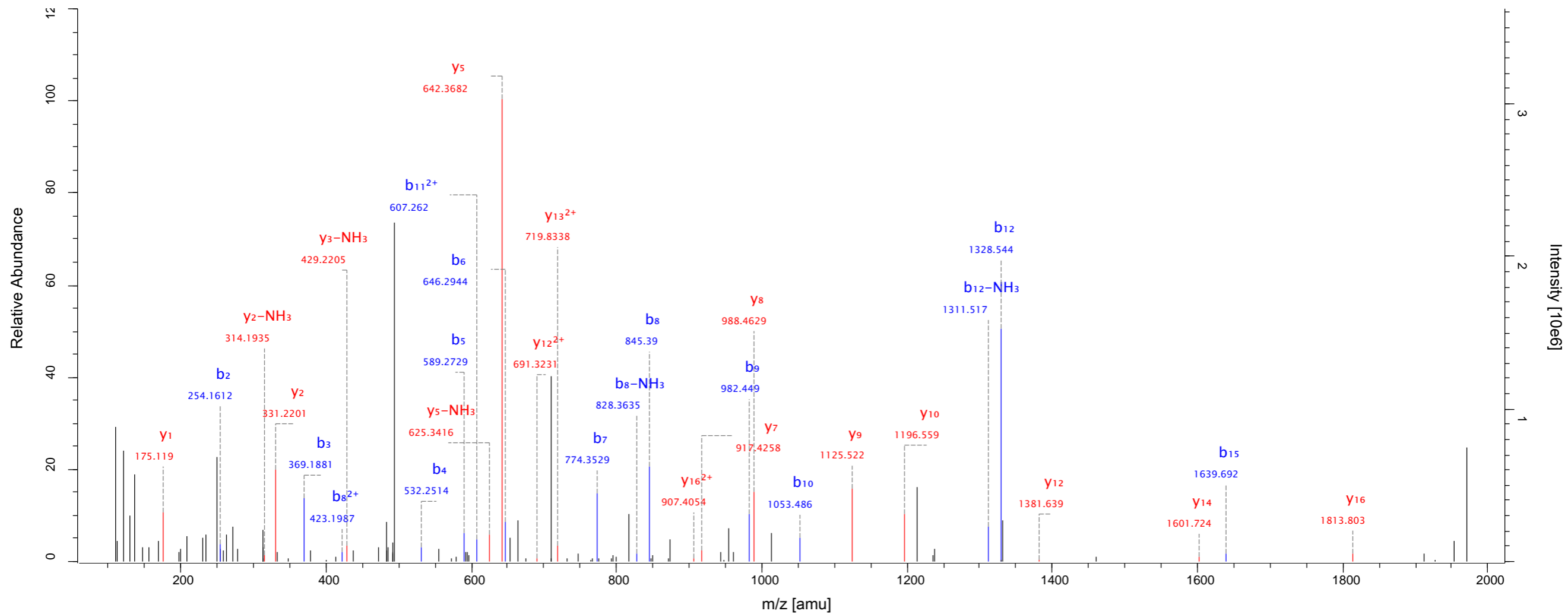
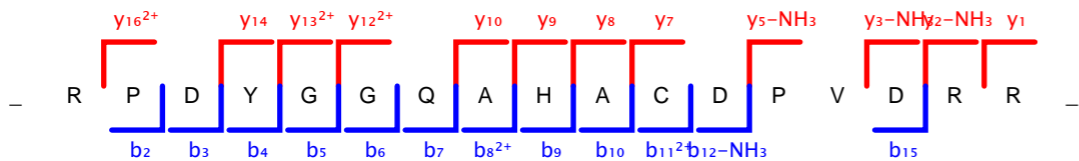
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 8178
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 64.48
Method: FTMS; HCD



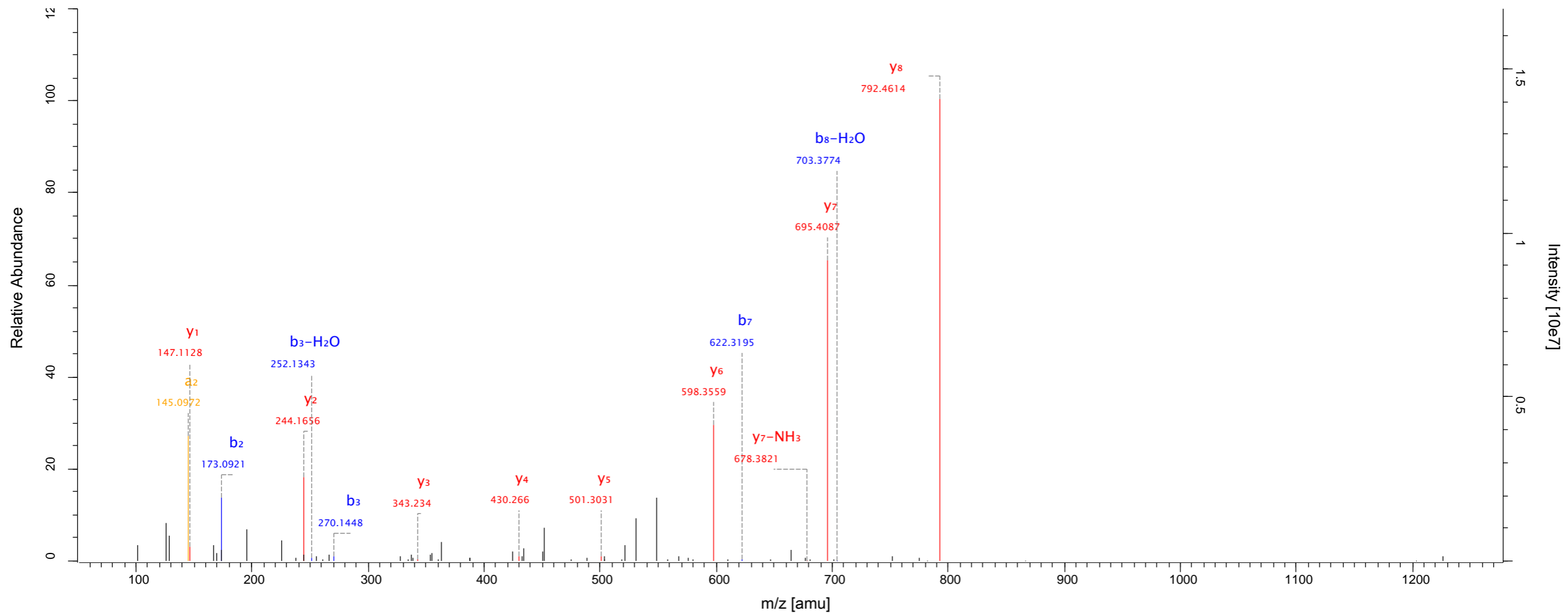
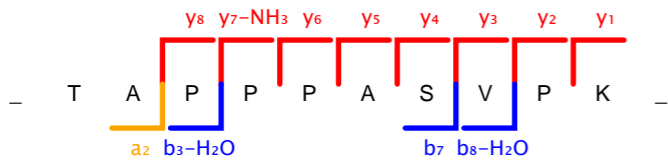
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 8489
Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
Peptide score: 75.46
Method: FTMS; HCD



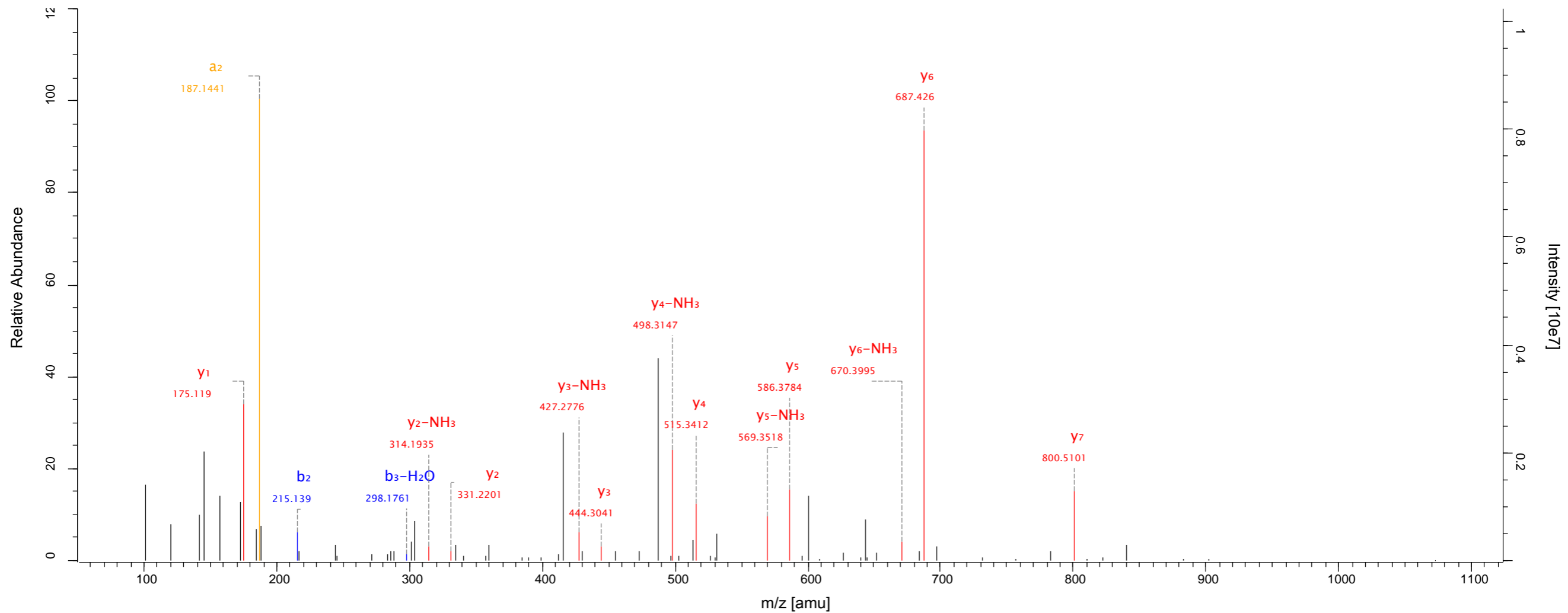
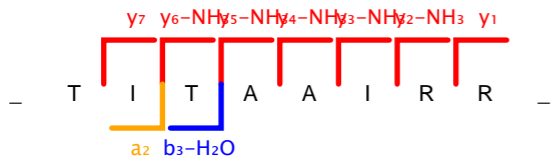
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
 Scan number: 8840
 Protein: ORFL86W + NC_006273.2[38059,38538]+ <unknown description>
 Peptide score: 133.96
 Method: FTMS; HCD



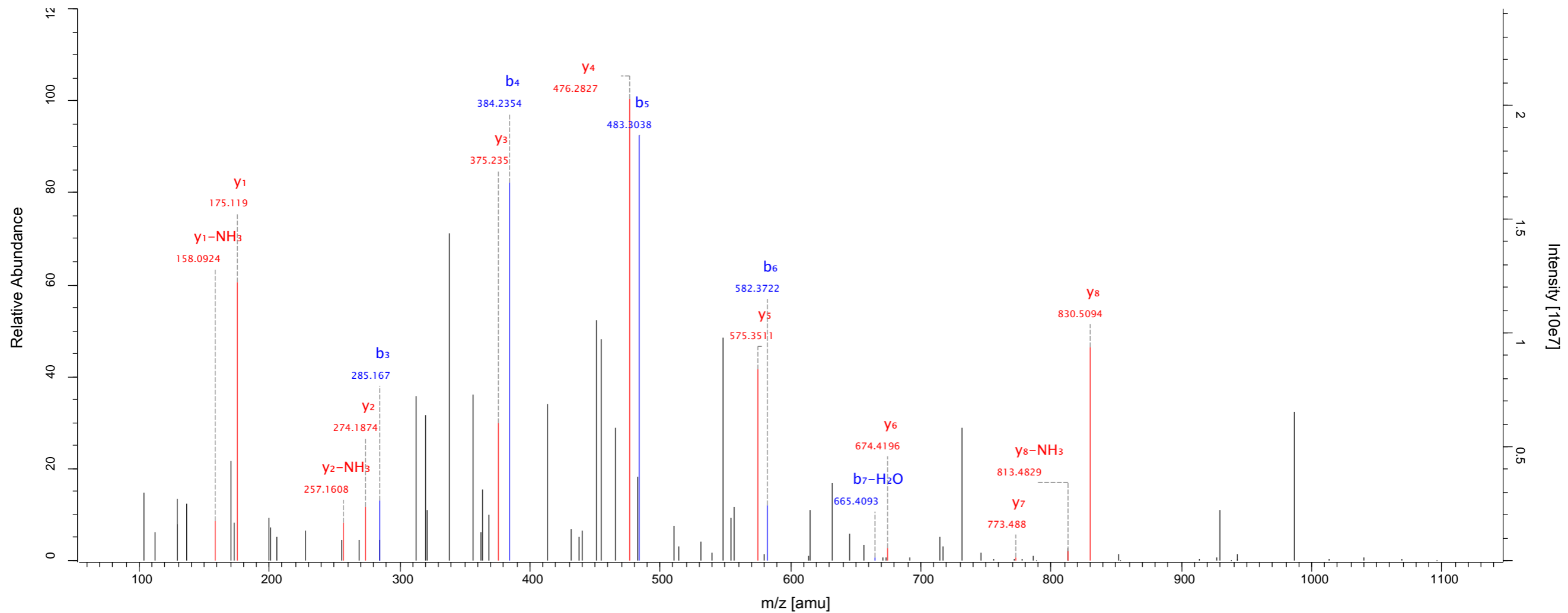
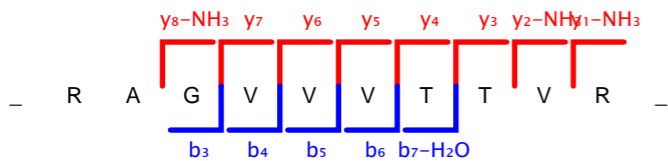
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 9417
Protein: ORFL234C - NC_006273.2[148754,149149]- <unknown description>
Peptide score: 55.08
Method: FTMS; HCD



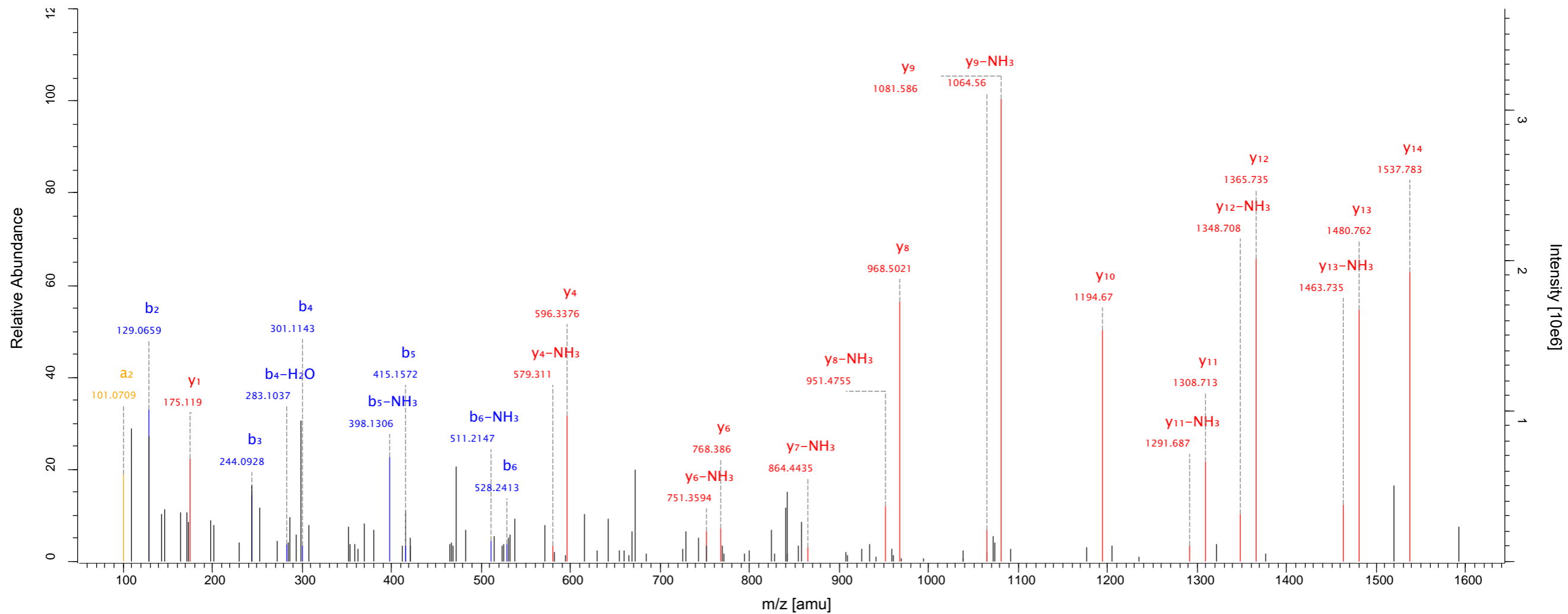
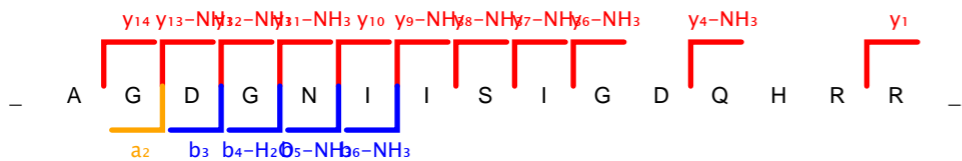
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_35
Scan number: 9876
Protein: ORFL161C.iORF2 - NC_006273.2[101063,101479]- <unknown description>
Peptide score: 104.21
Method: FTMS; HCD



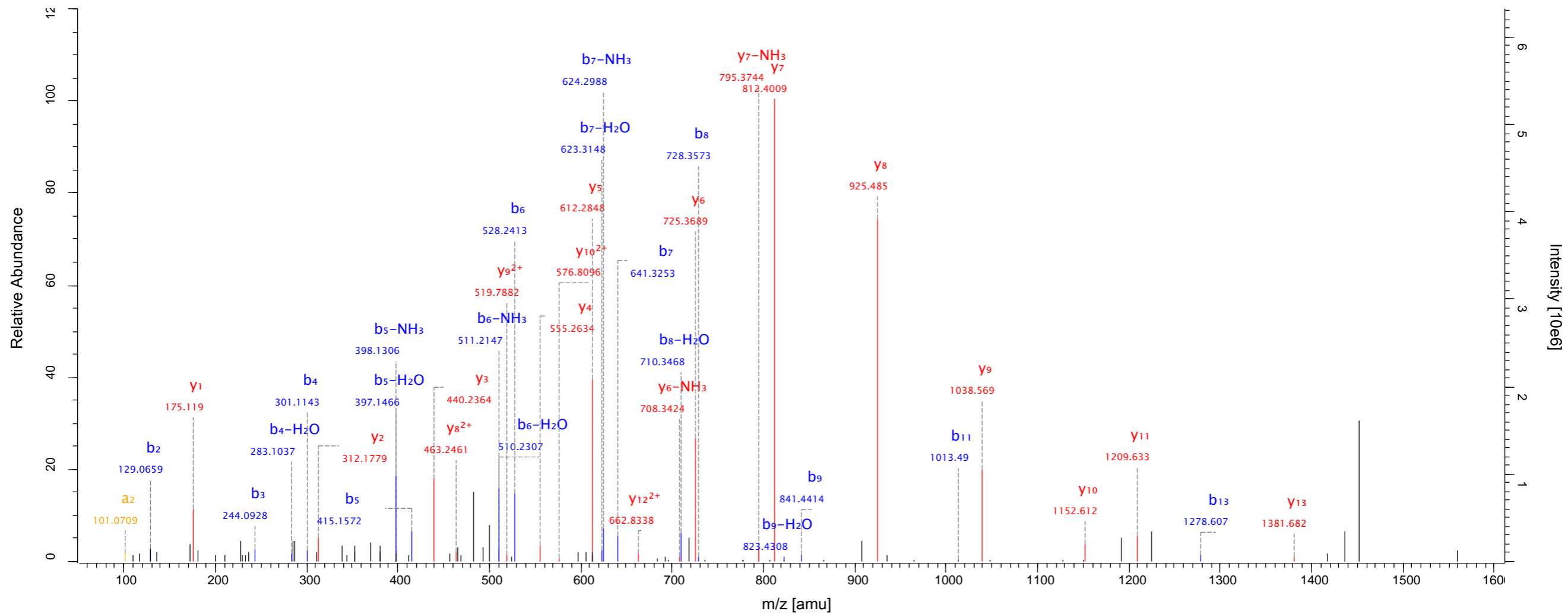
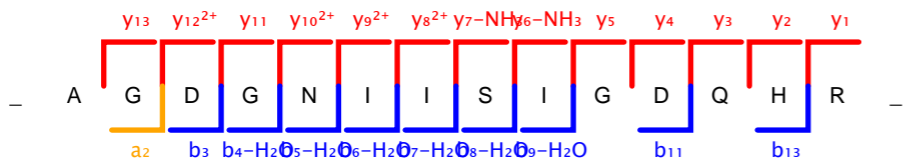
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
Scan number: 10651
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 87.15
Method: FTMS; HCD



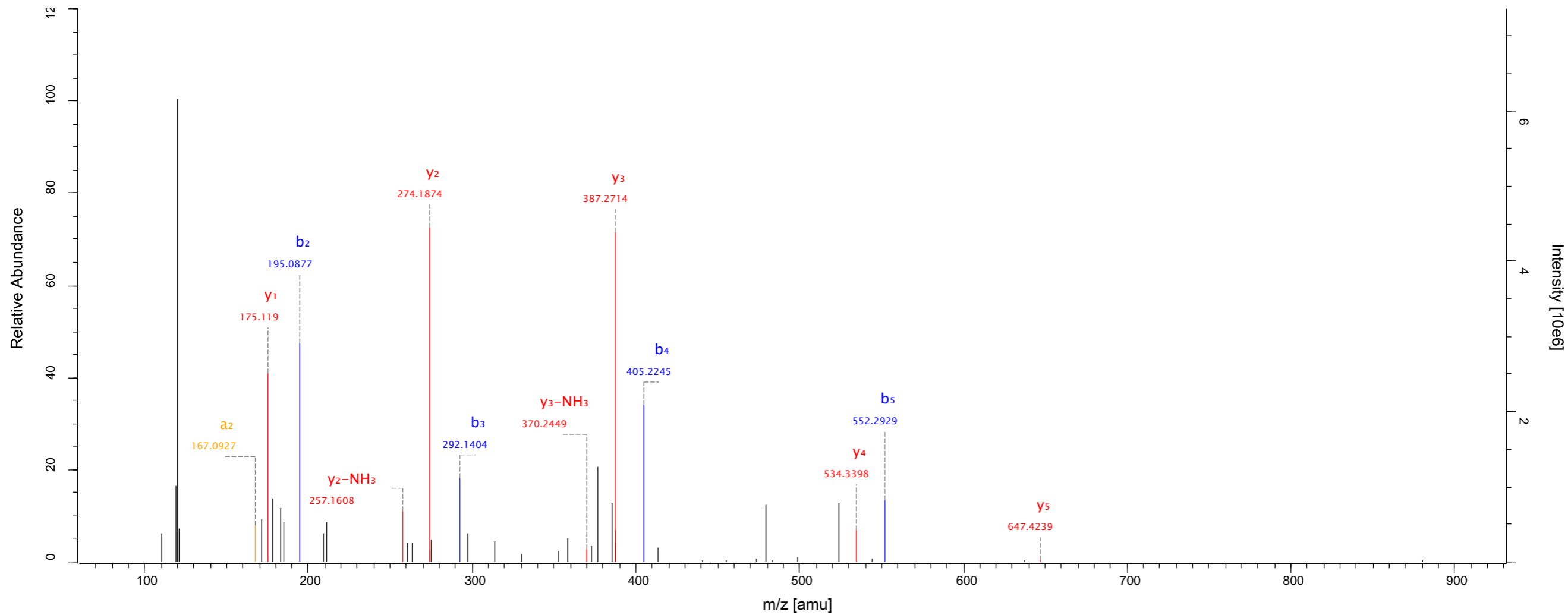
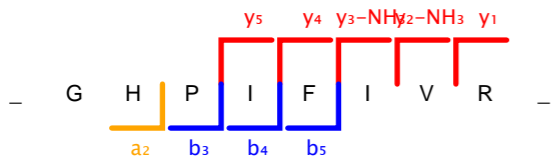
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
 Scan number: 17563
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 138.42
 Method: FTMS; HCD



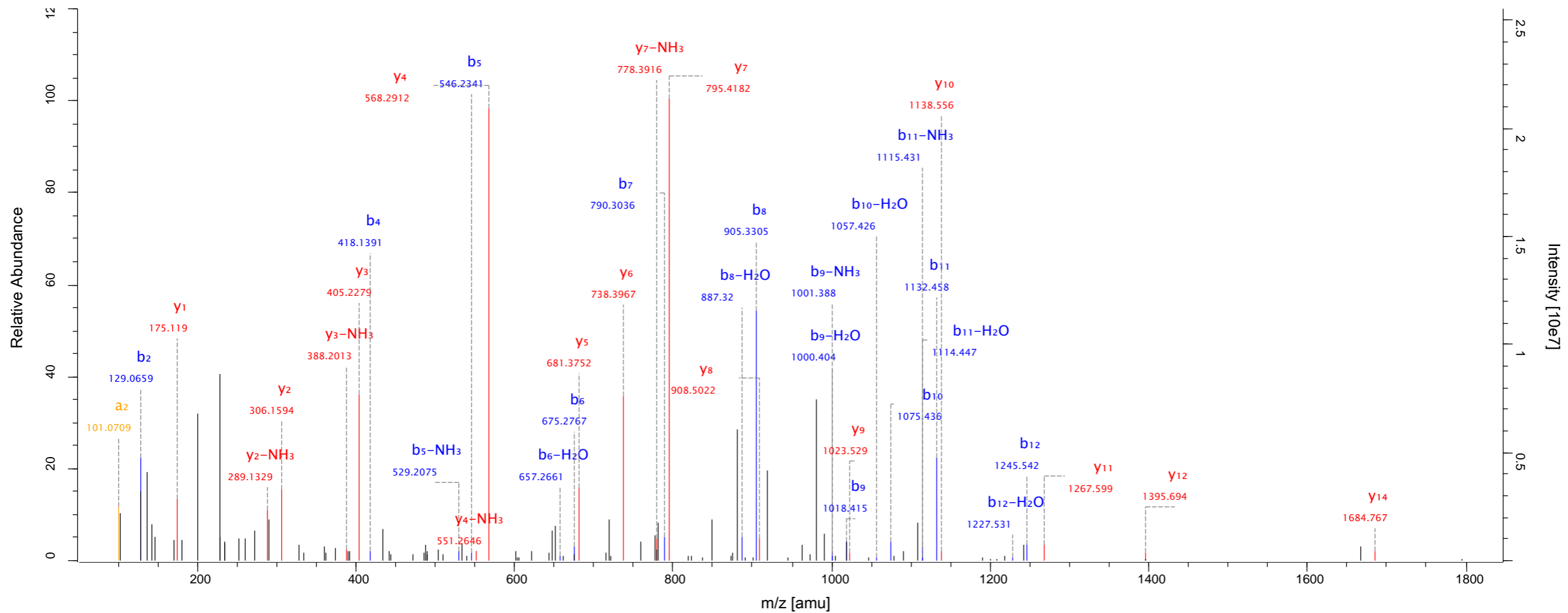
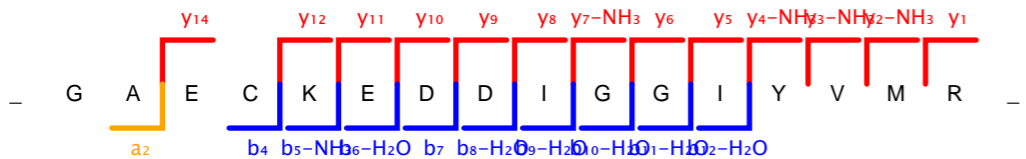
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
 Scan number: 22267
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 213.18
 Method: FTMS; HCD



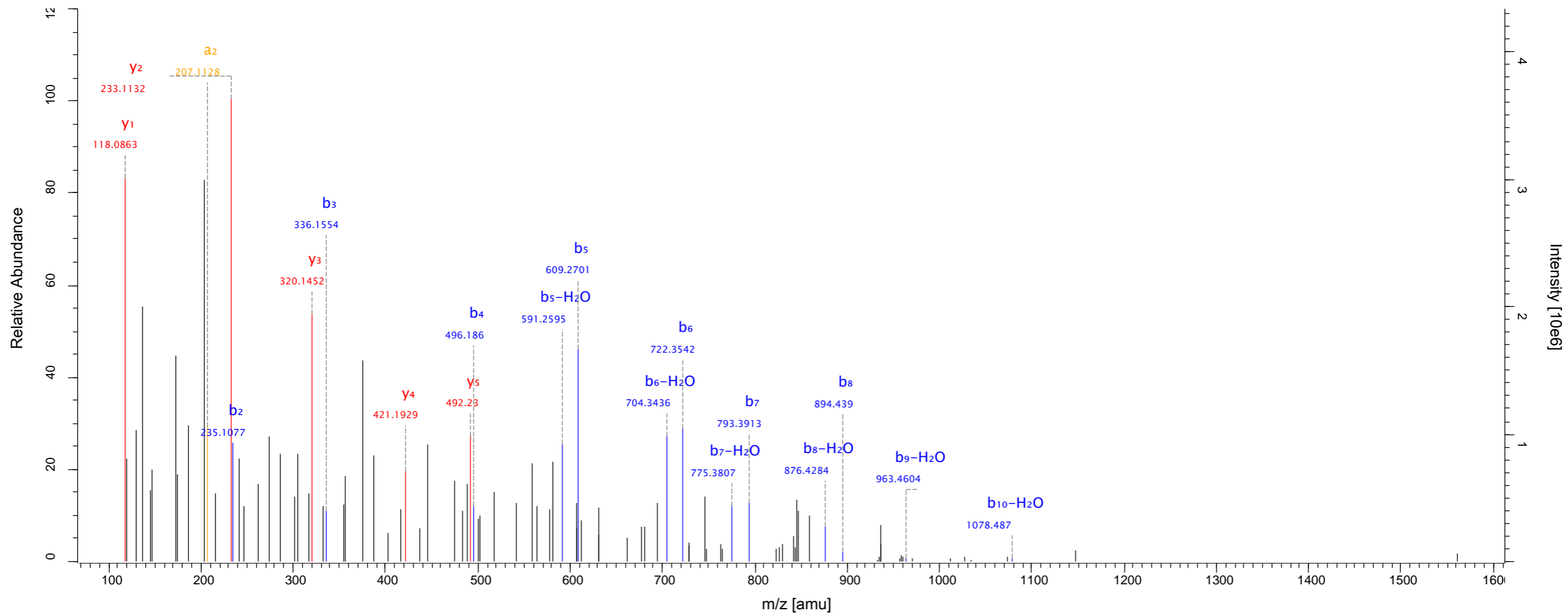
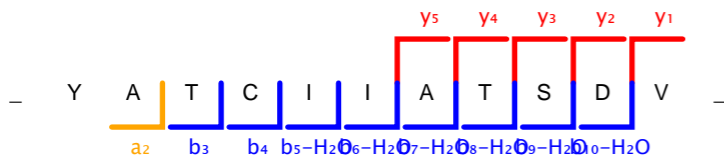
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_36
Scan number: 23427
Protein: ORFL114W + NC_006273.2[60151,60708]+ <unknown description>
Peptide score: 93.26
Method: FTMS; HCD



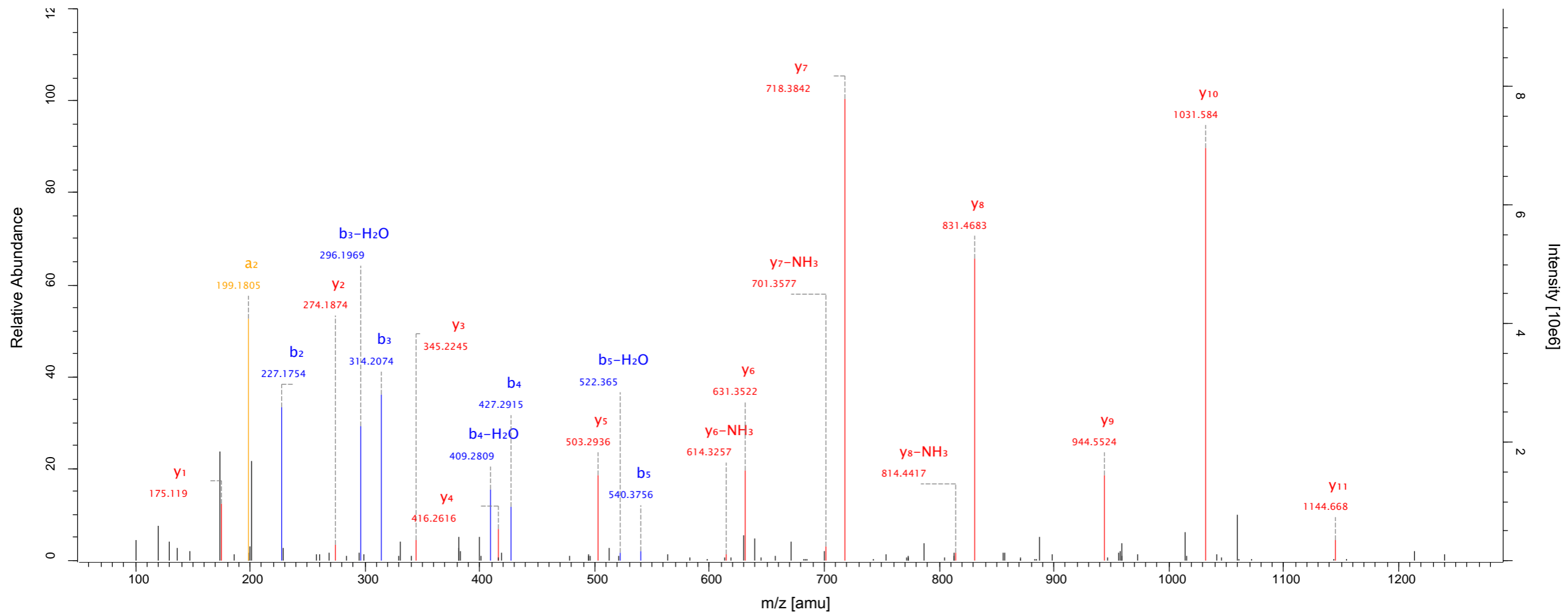
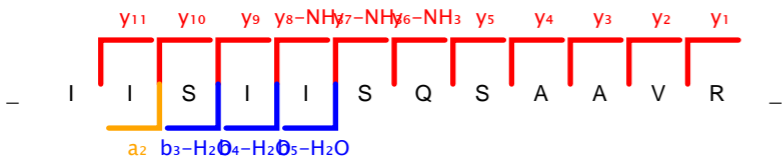
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
 Scan number: 29817
 Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
 Peptide score: 190.41
 Method: FTMS; HCD



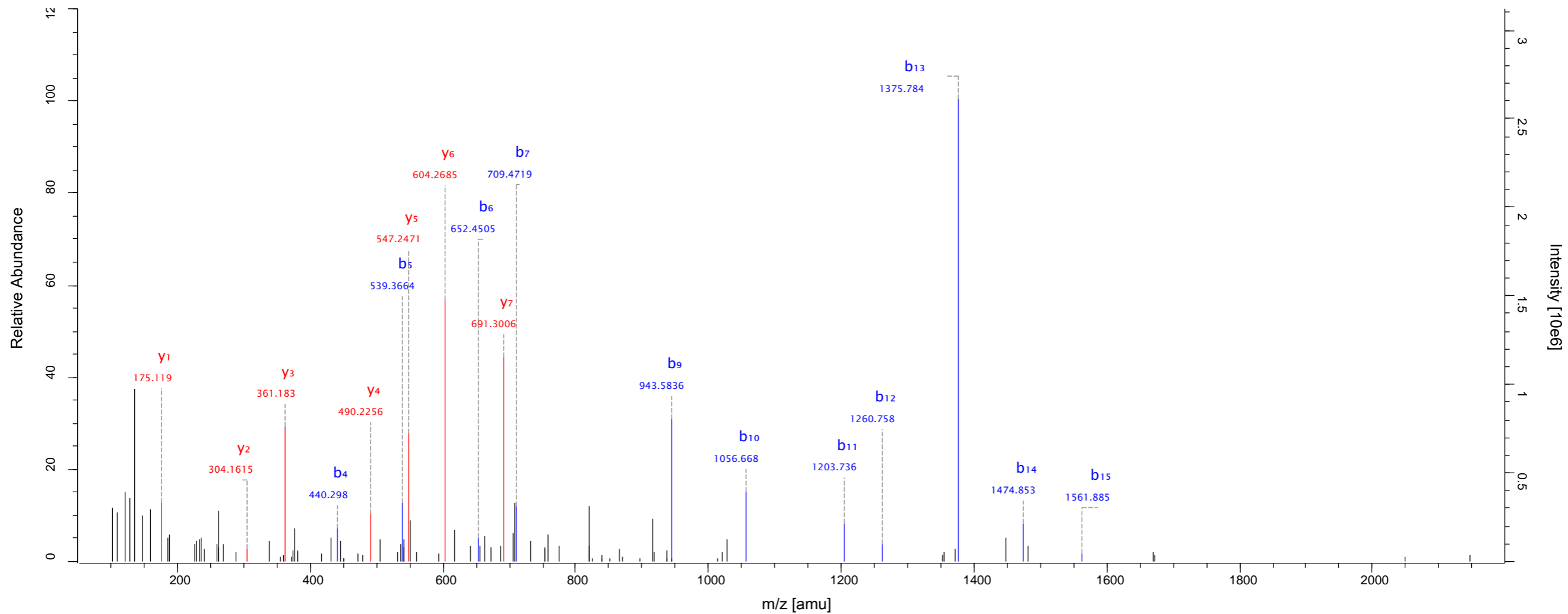
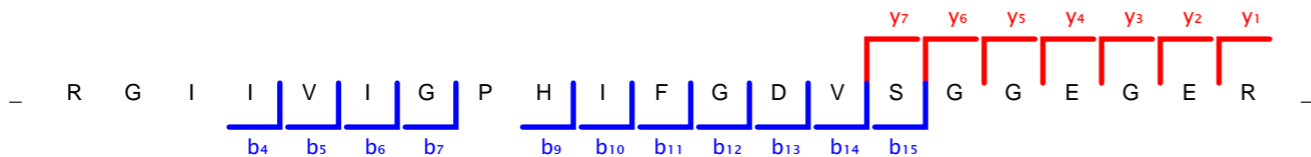
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
 Scan number: 34110
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 92.05
 Method: FTMS; HCD



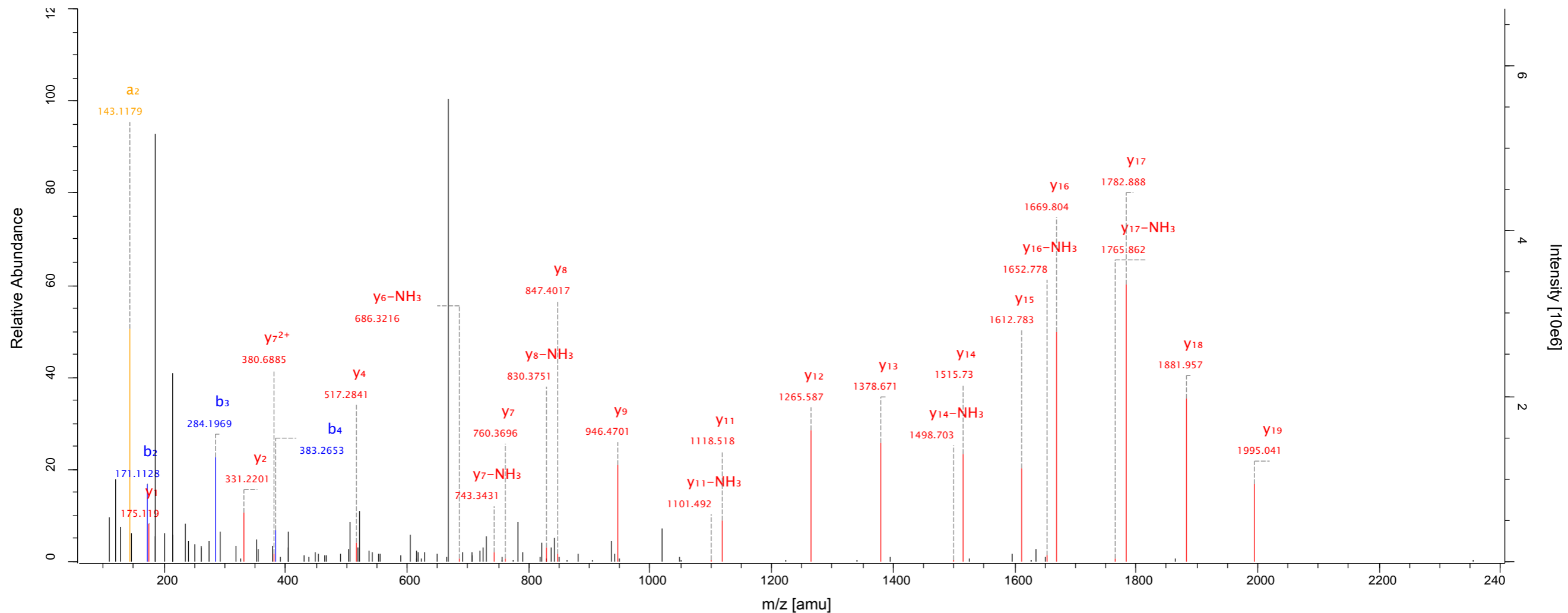
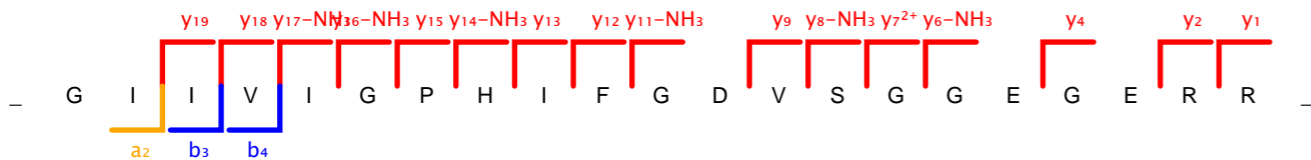
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
 Scan number: 37079
 Protein: ORFS365W + NC_006273.2[225950,226606]+ <unknown description>
 Peptide score: 177.57
 Method: FTMS; HCD



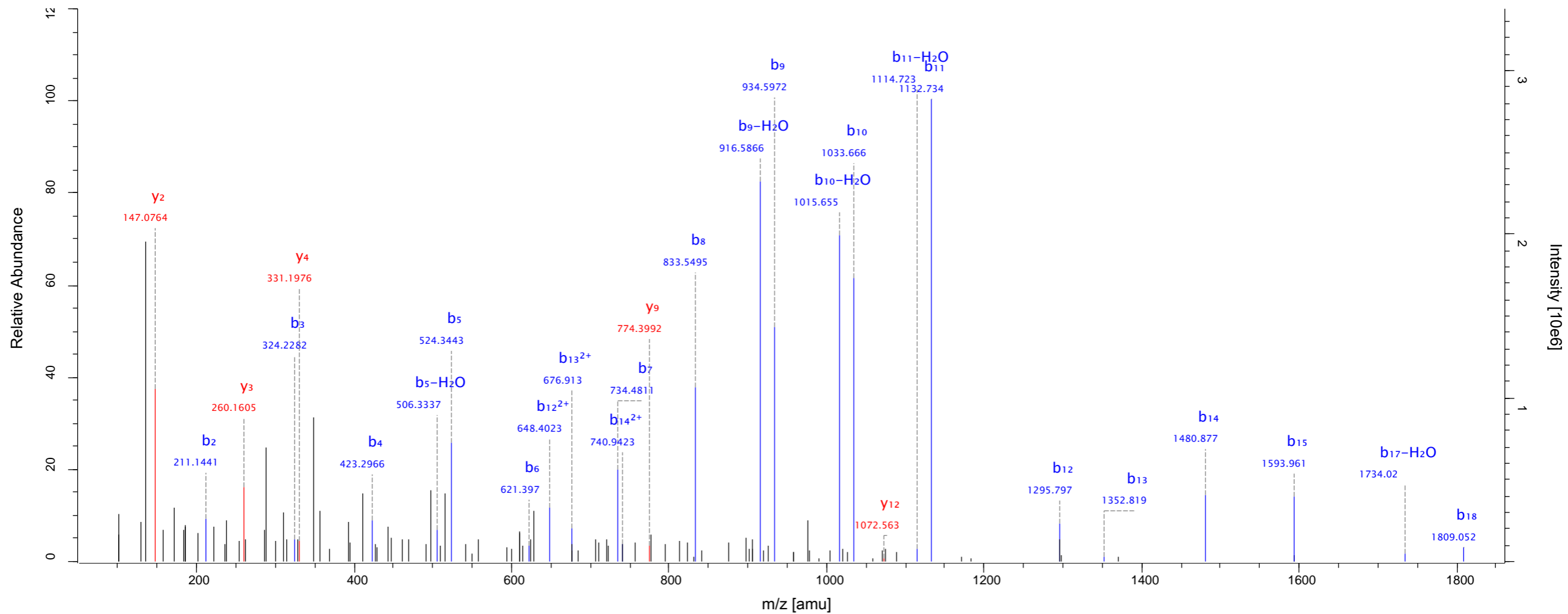
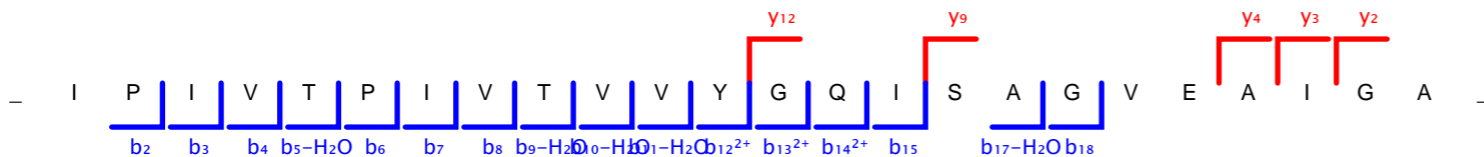
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
Scan number: 44126
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 83.85
Method: FTMS; HCD



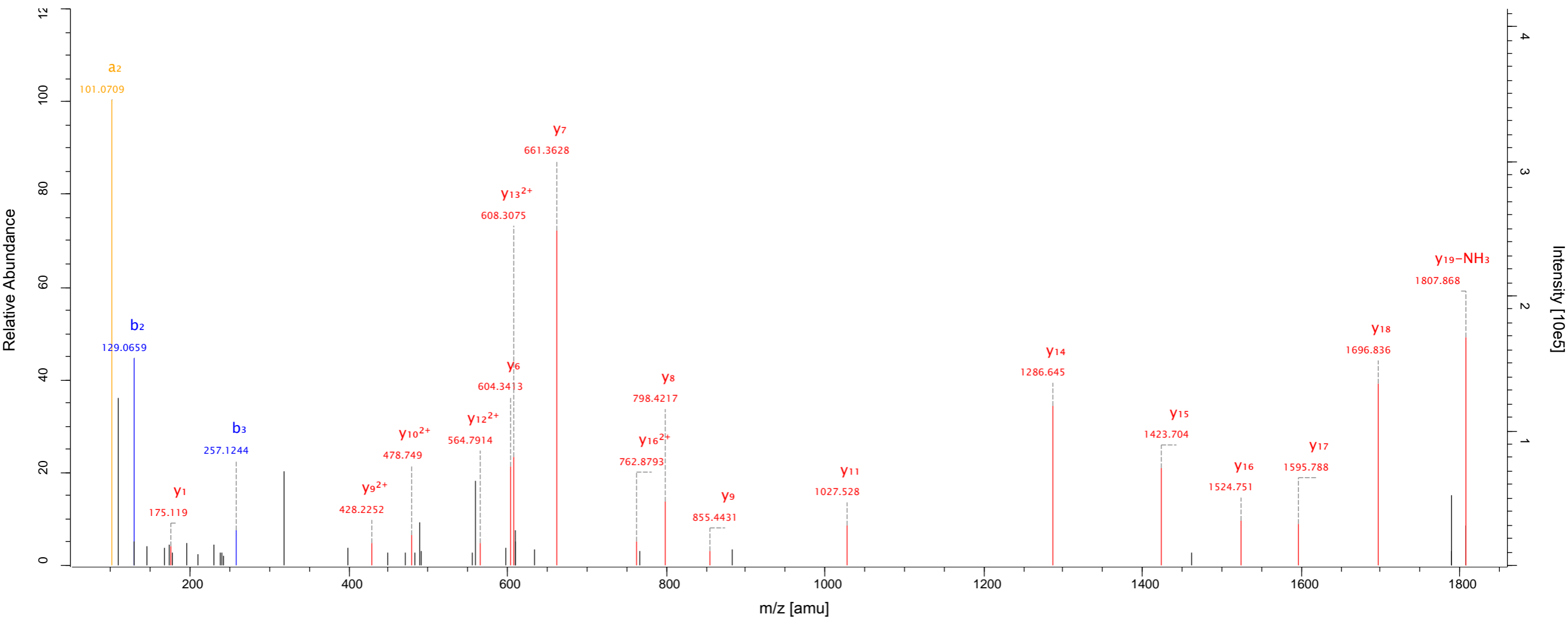
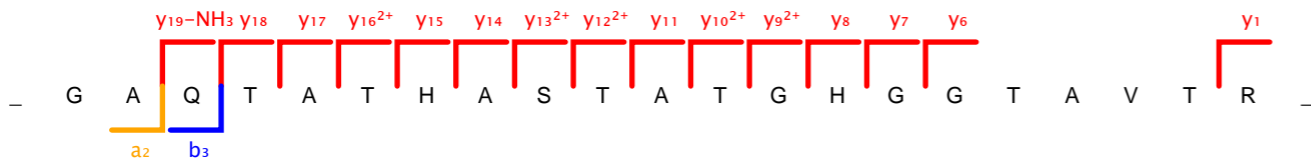
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
 Scan number: 44703
 Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
 Peptide score: 108.56
 Method: FTMS; HCD



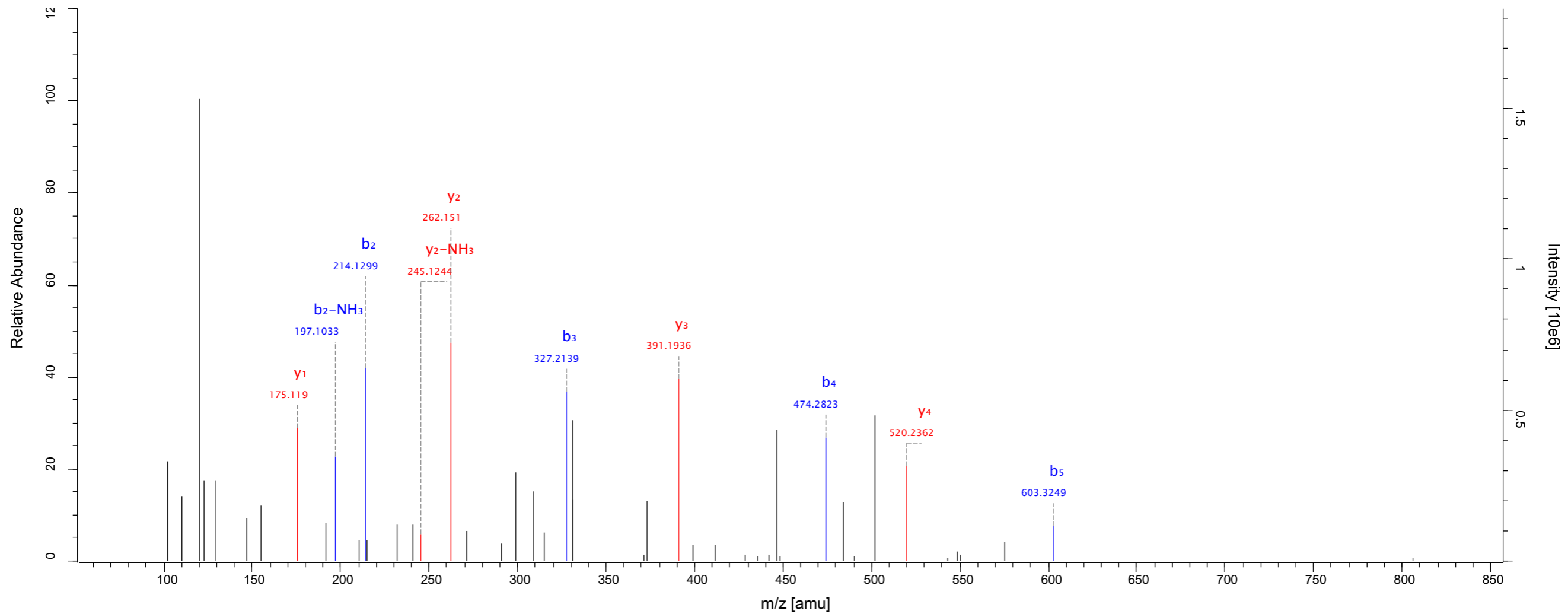
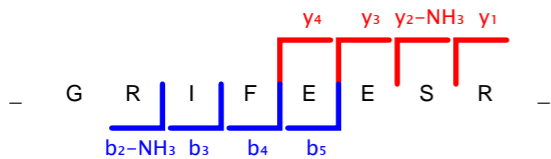
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_36
Scan number: 53871
Protein: ORFL130W + NC_006273.2[72313,72744]+ <unknown description>
Peptide score: 120.11
Method: FTMS; HCD



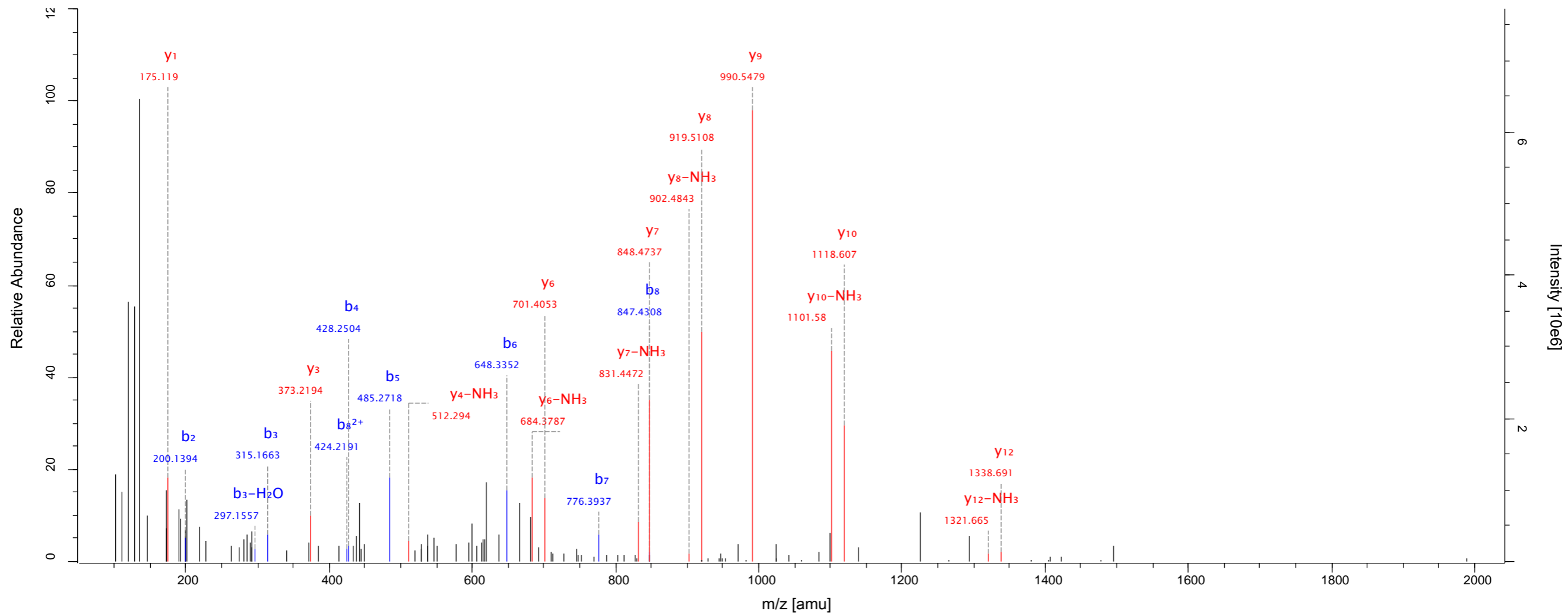
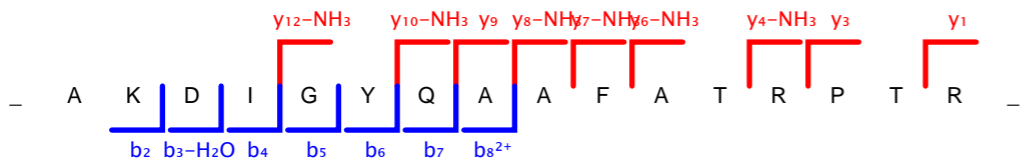
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_36
Scan number: 5634
Protein: ORFL114W + NC_006273.2[60151,60708]+ <unknown description>
Peptide score: 122.1
Method: FTMS; HCD



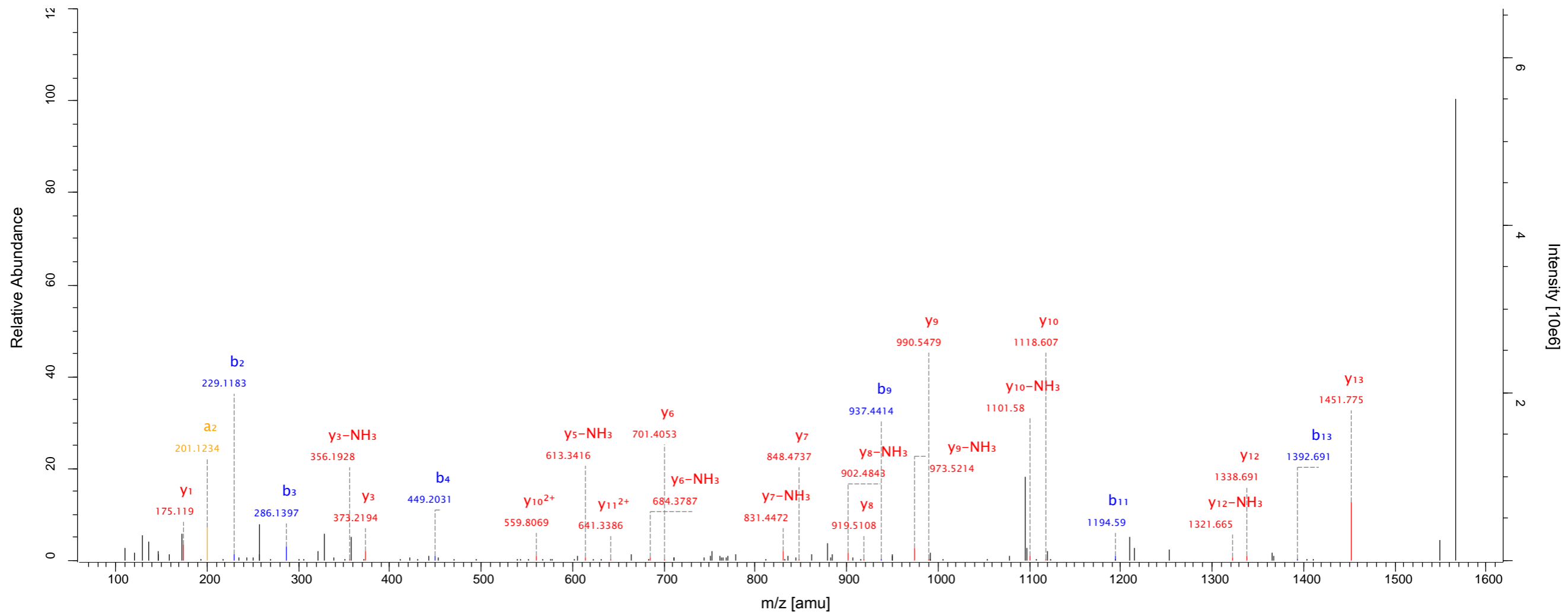
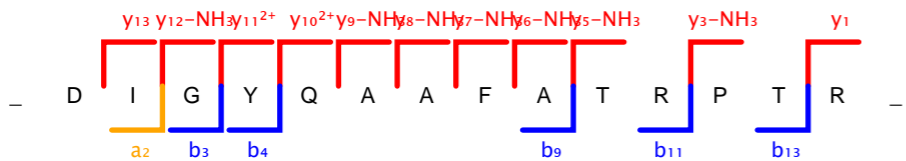
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_36
Scan number: 9408
Protein: ORFL299W + NC_006273.2[185047,185685]+ <unknown description>
Peptide score: 72.48
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_37
 Scan number: 19603
 Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
 Peptide score: 115.82
 Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_37
Scan number: 25814
Protein: ORFL82W + NC_006273.2[36796,37488]+ <unknown description>
Peptide score: 113.07
Method: FTMS; HCD



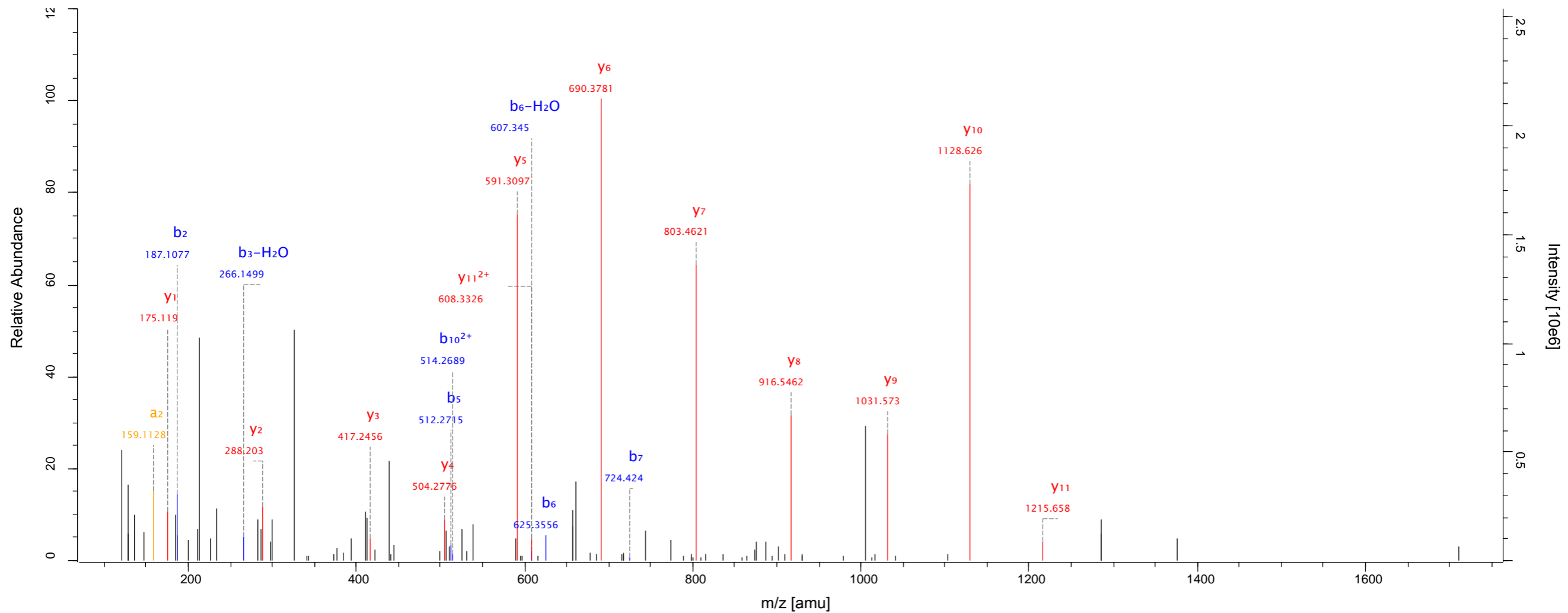
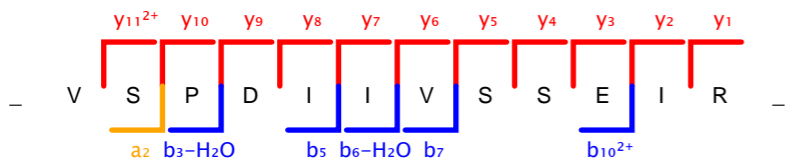
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_37

Scan number: 38008

Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 89.8

Method: FTMS; HCD



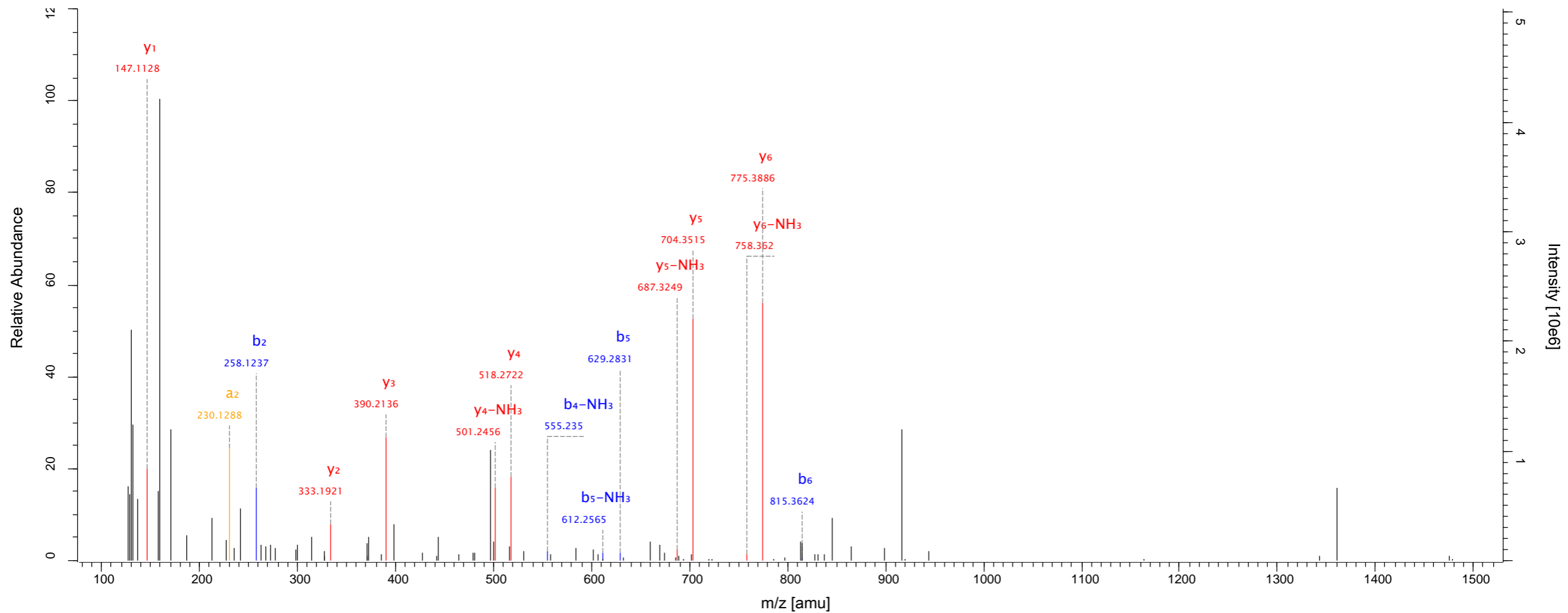
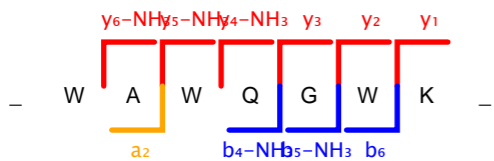
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_37

Scan number: 38766

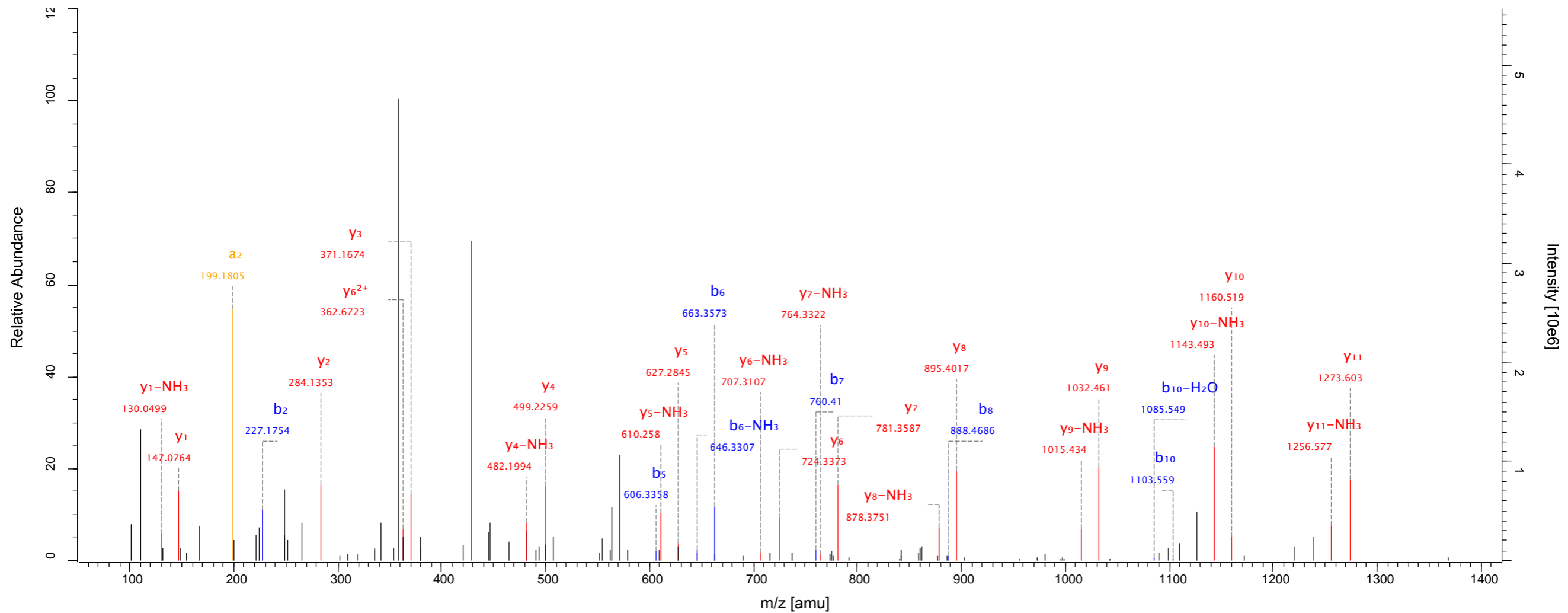
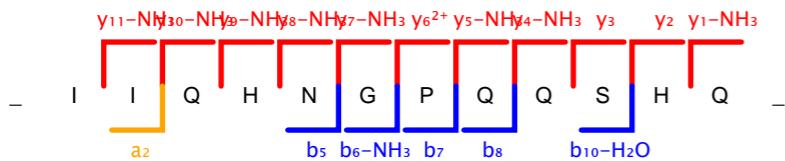
Protein: ORFL321W + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF1 + NC_006273.2[191968,194126]+ <unknown description>; ORFL321W.iORF4 + NC_006273.2[193200,194126]+ <unknown description>

Peptide score: 114.02

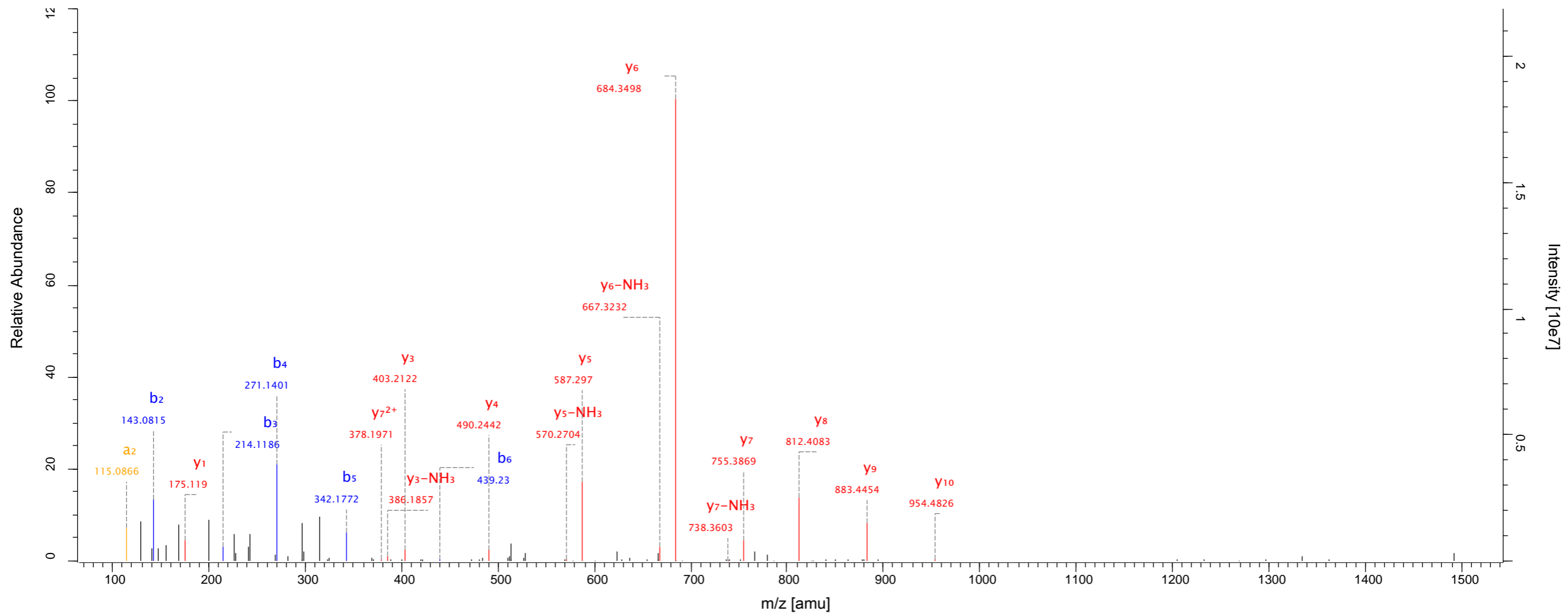
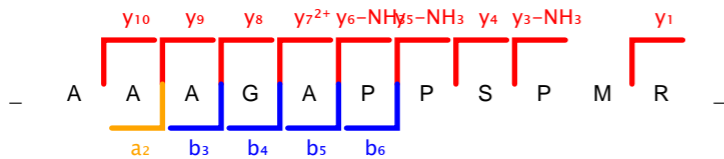
Method: FTMS; HCD



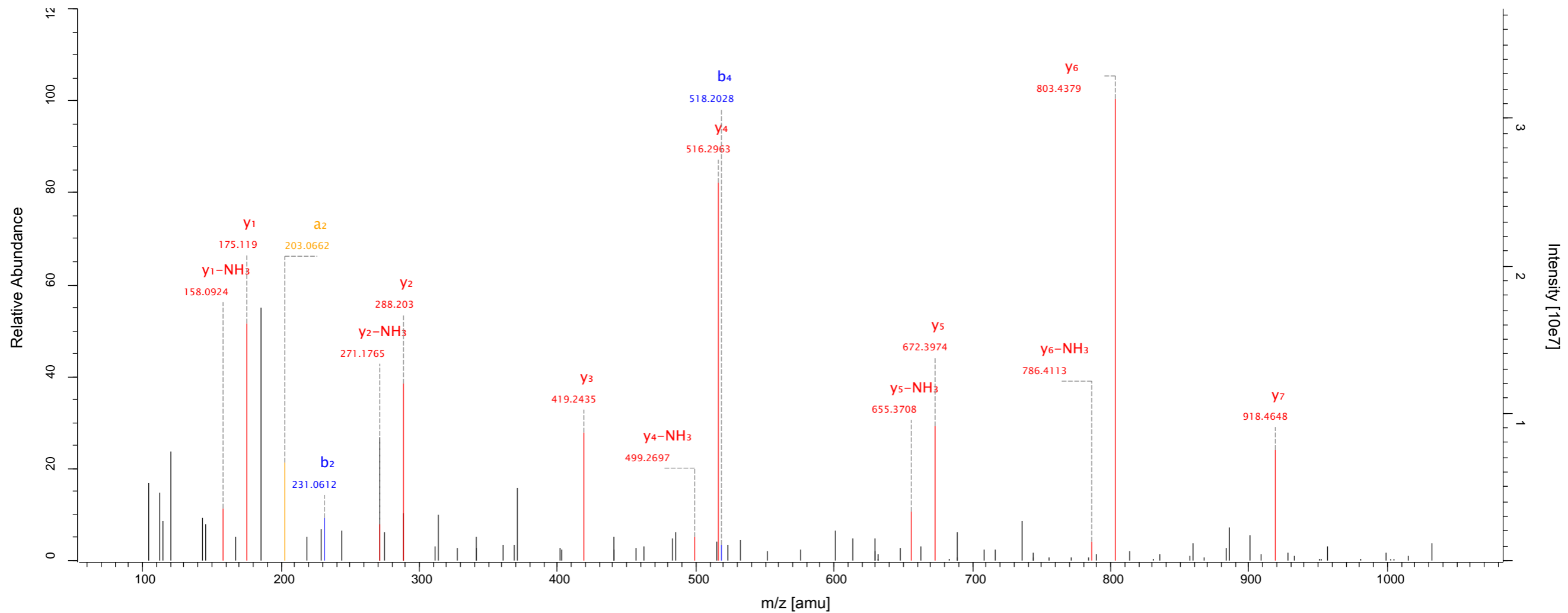
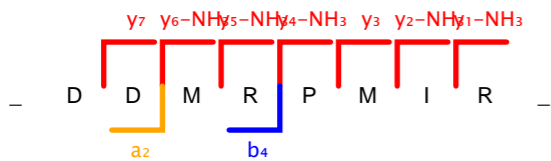
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_37
 Scan number: 5805
 Protein: ORFL103C - NC_006273.2[52569,53060]- <unknown description>
 Peptide score: 199.68
 Method: FTMS; HCD



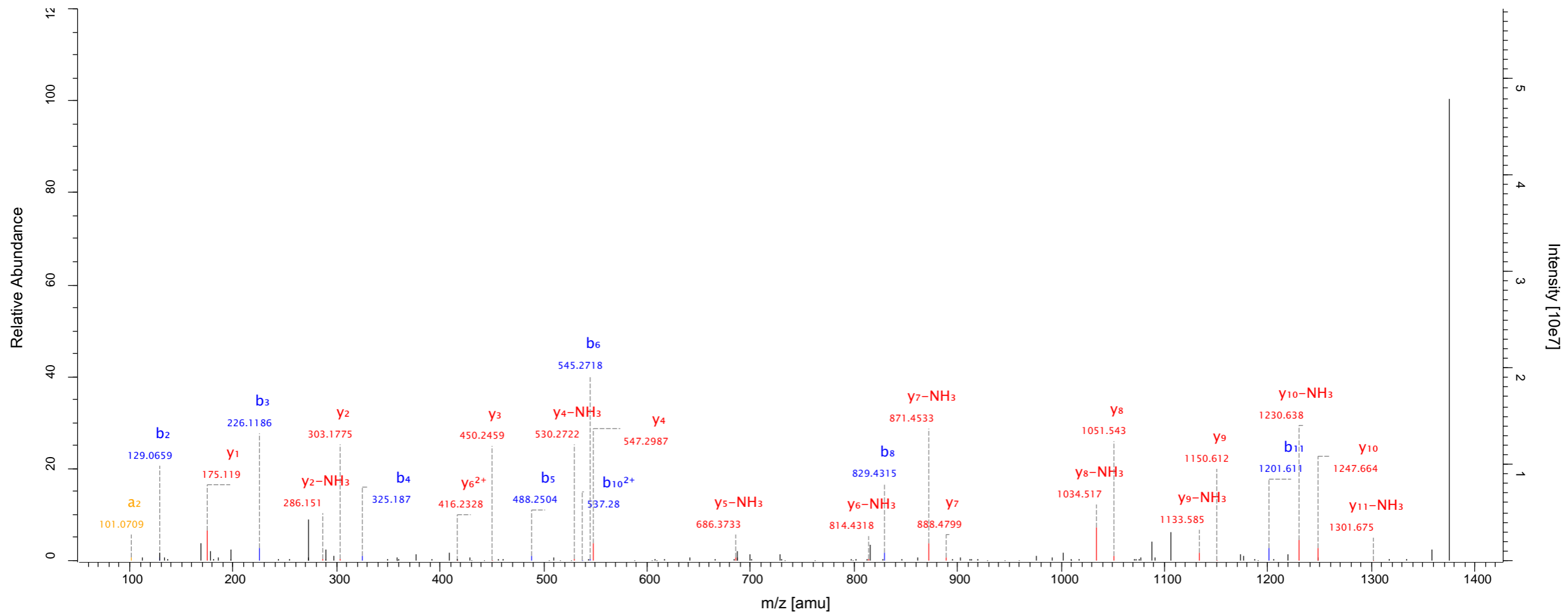
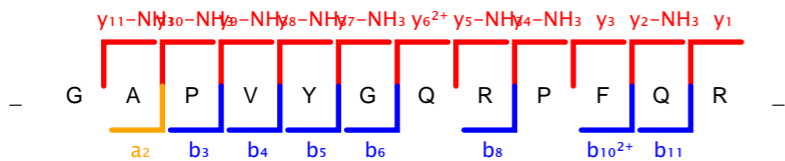
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
Scan number: 10768
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 130.66
Method: FTMS; HCD



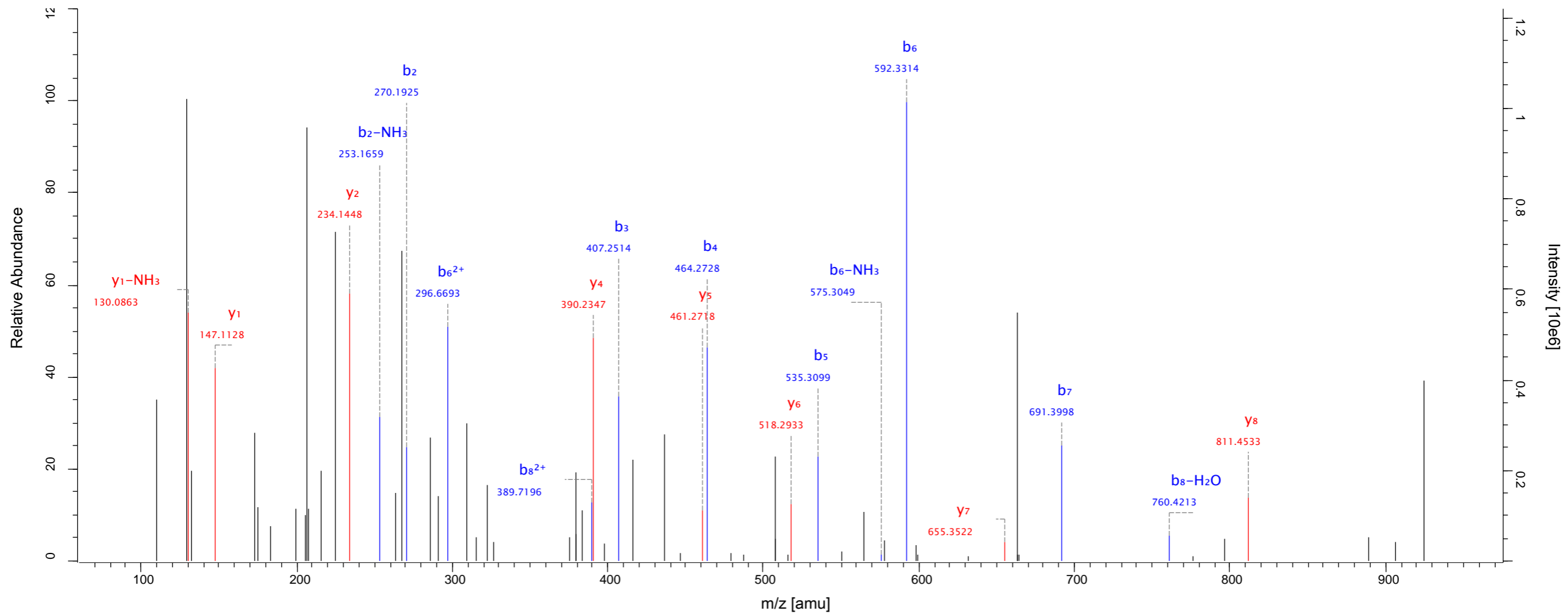
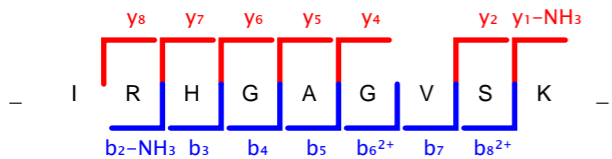
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
Scan number: 14803
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 101.33
Method: FTMS; HCD



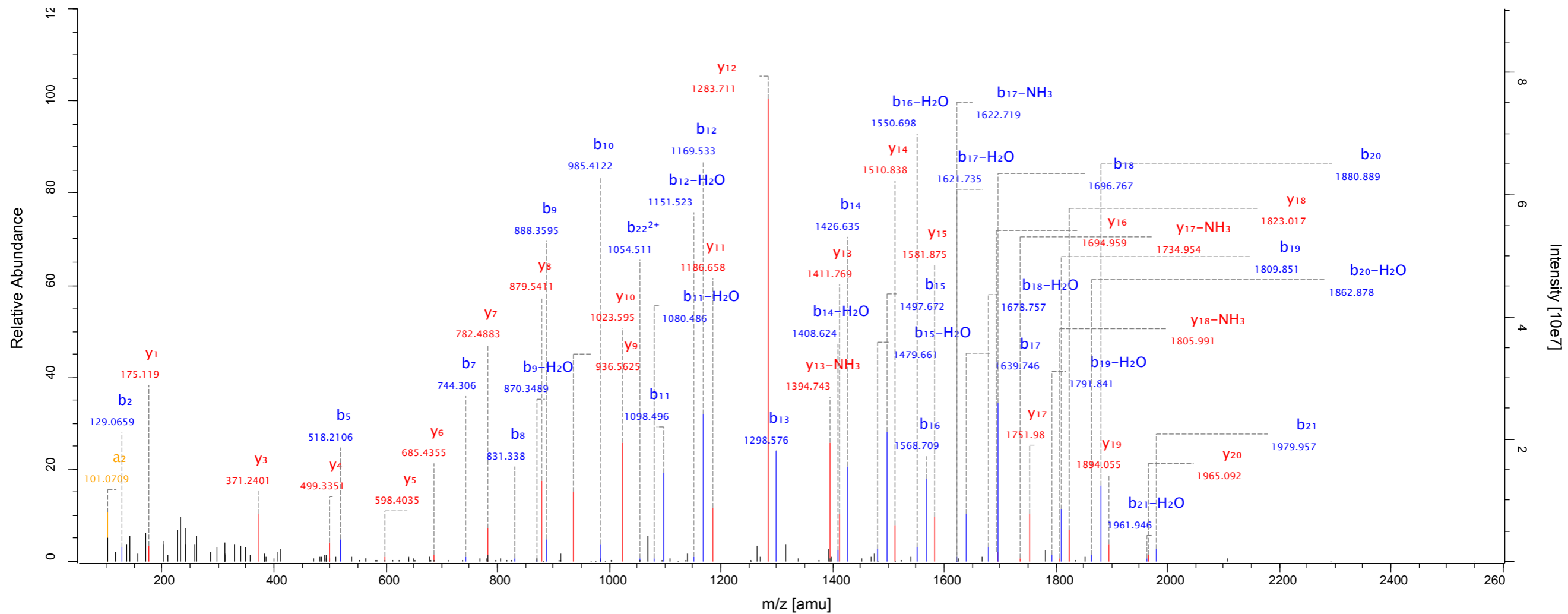
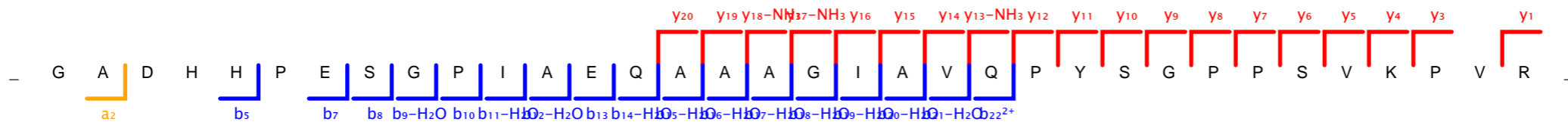
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
 Scan number: 16094
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 126.29
 Method: FTMS; HCD



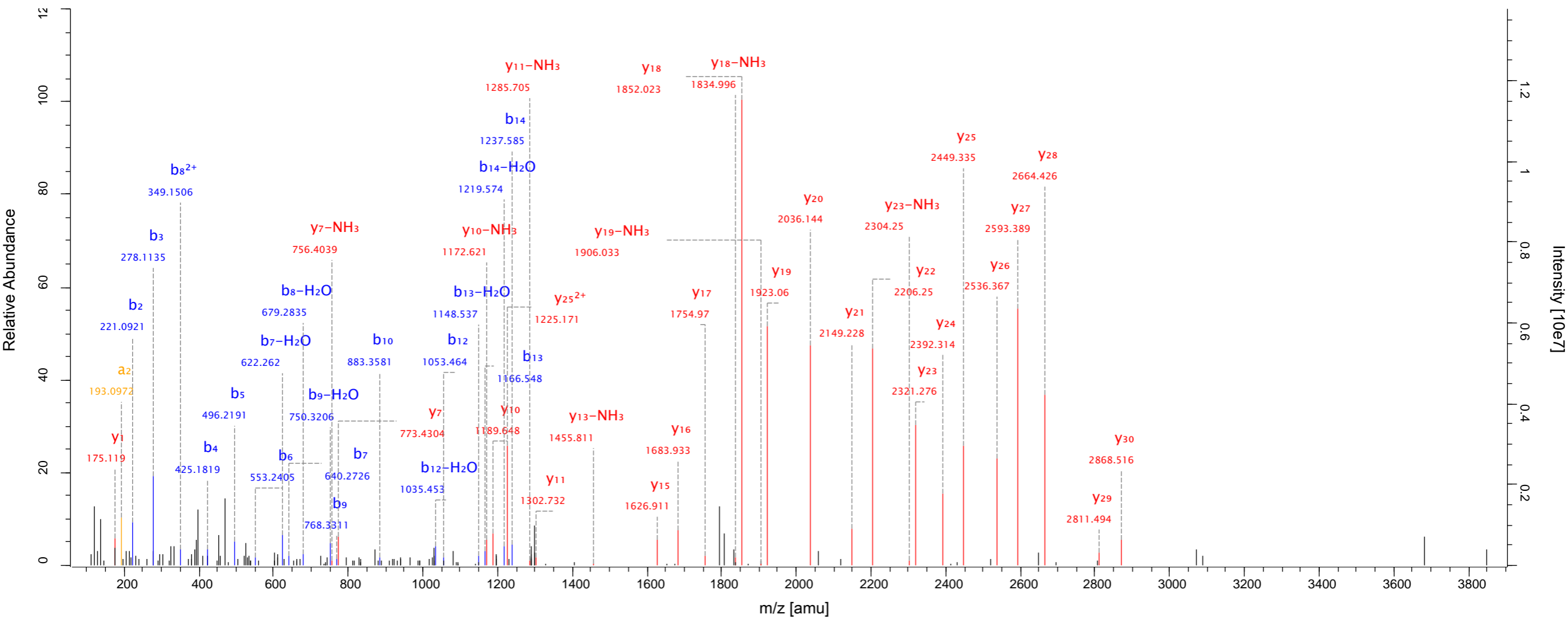
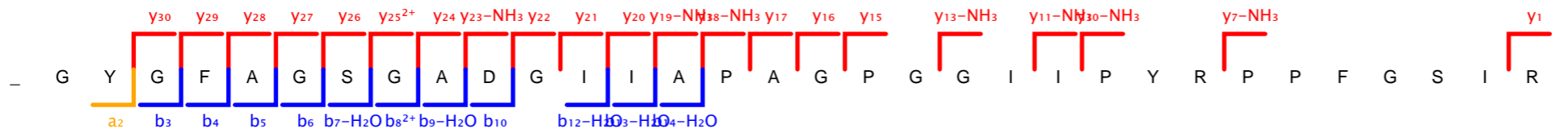
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
Scan number: 3538
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 106.68
Method: FTMS; HCD



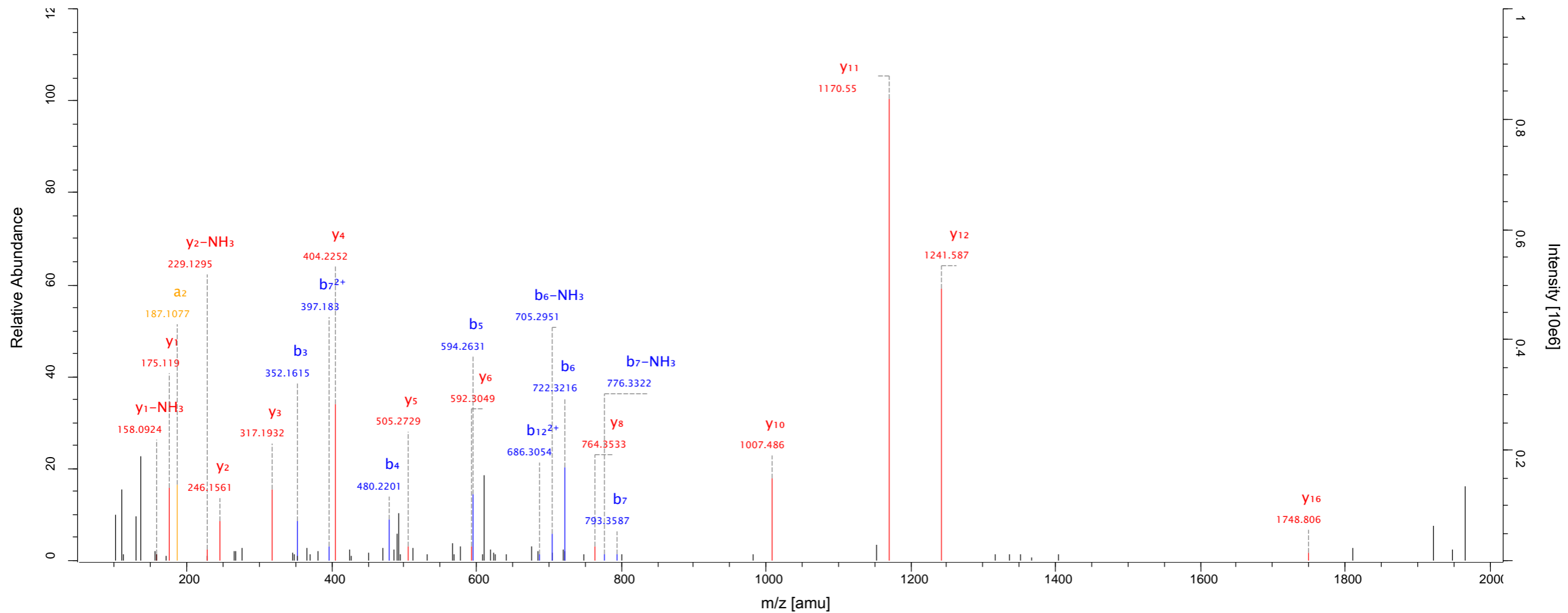
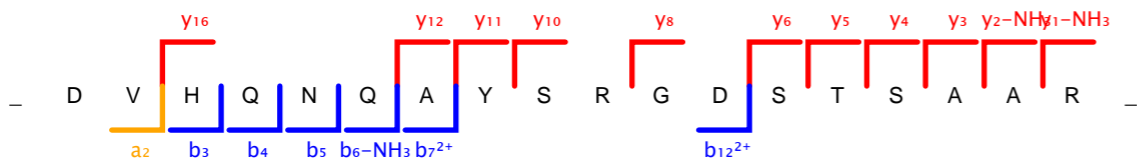
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
 Scan number: 36859
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 229.56
 Method: FTMS; HCD



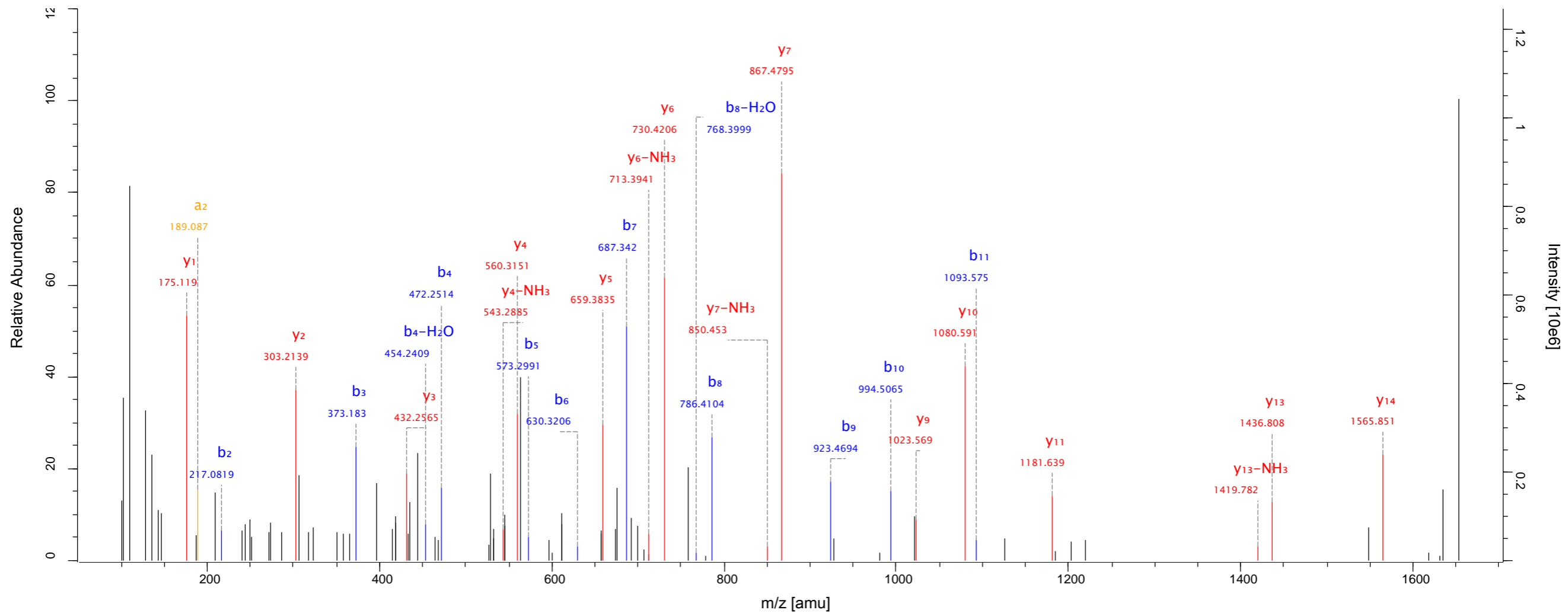
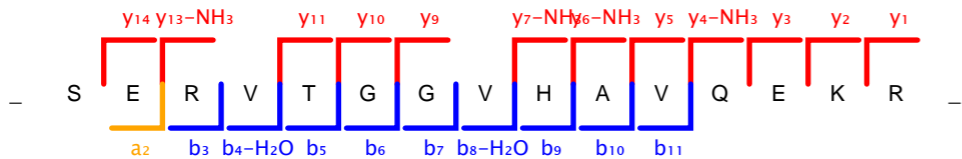
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
 Scan number: 59976
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 198.56
 Method: FTMS; HCD



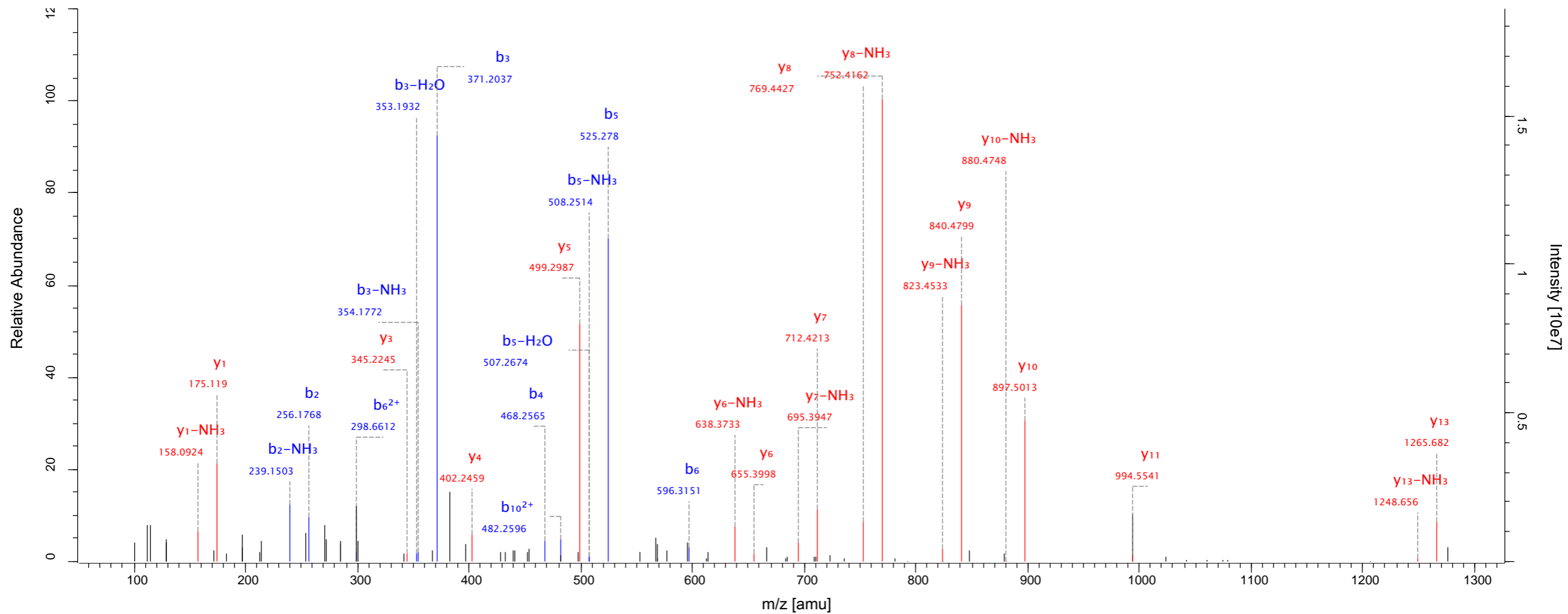
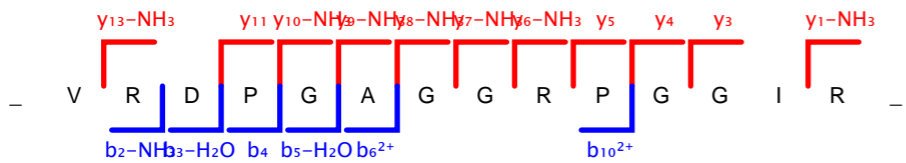
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
 Scan number: 6480
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 106.38
 Method: FTMS; HCD



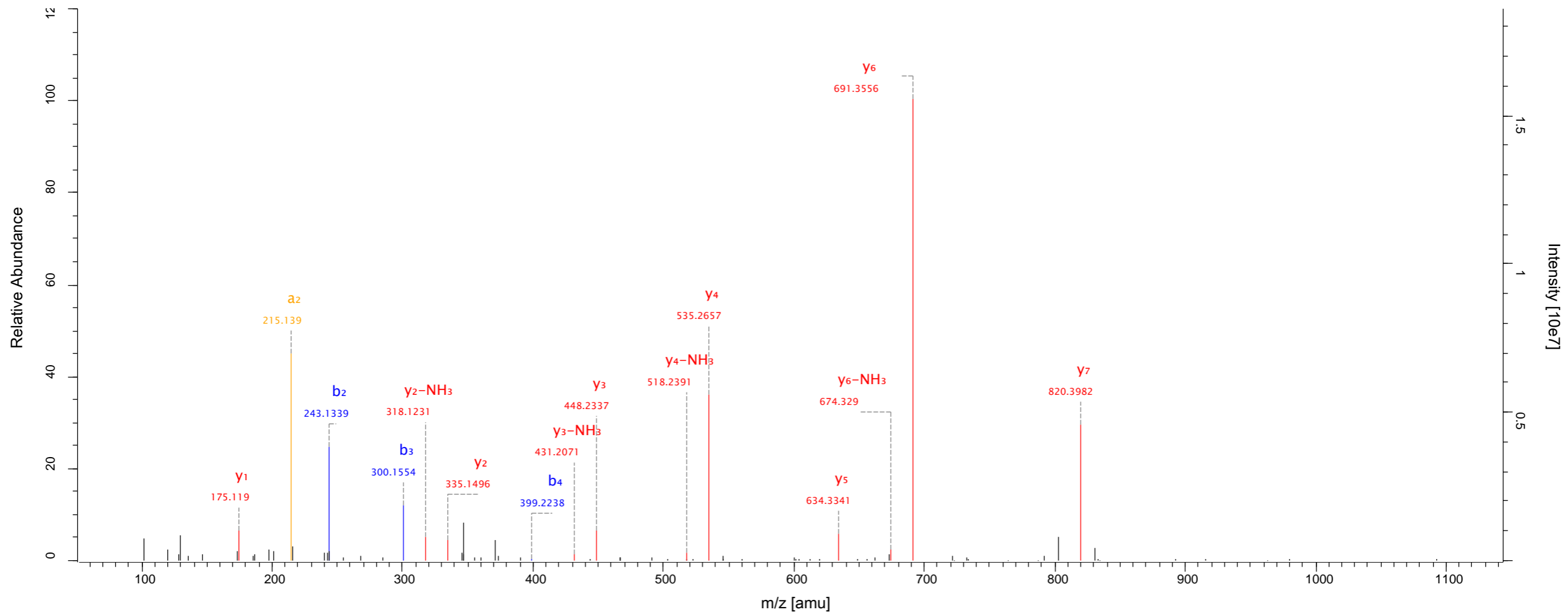
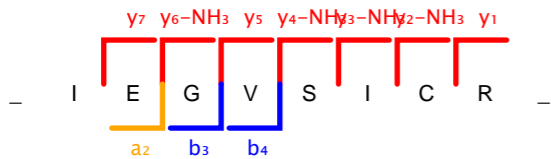
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_39
Scan number: 7226
Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
Peptide score: 151.9
Method: FTMS; HCD



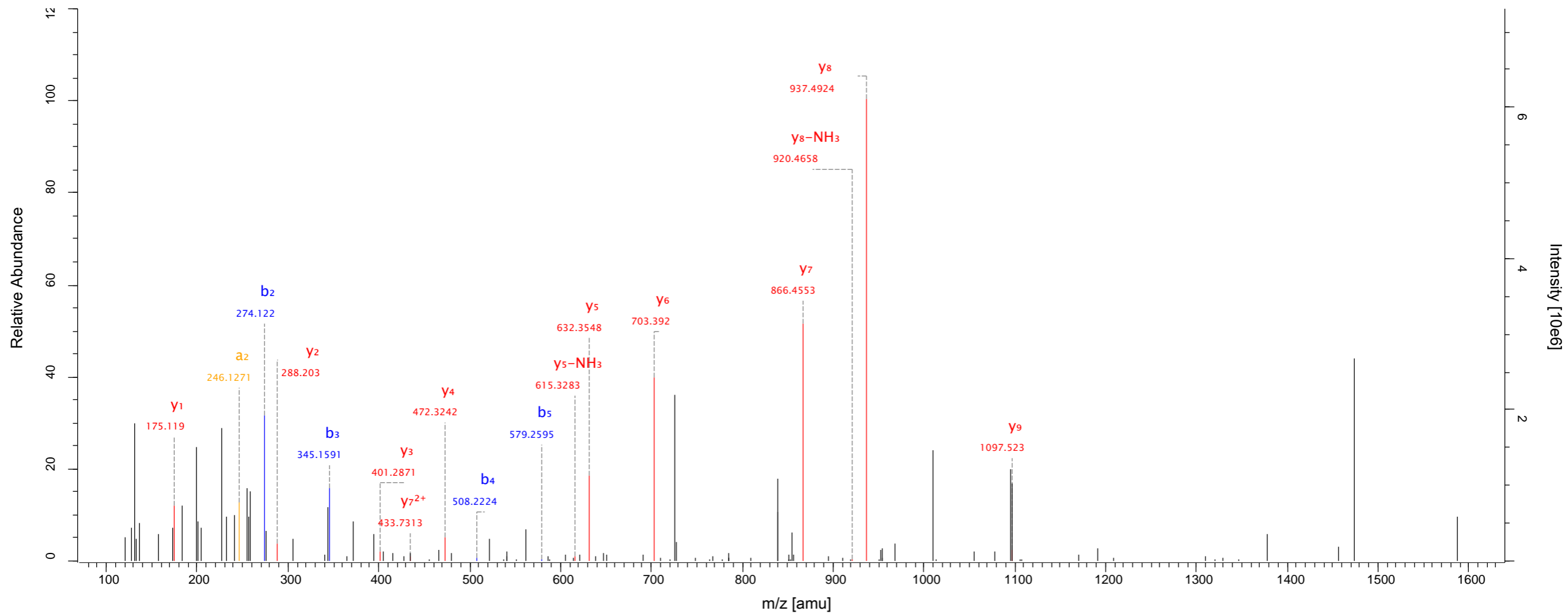
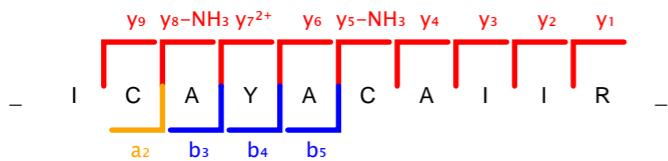
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_40
 Scan number: 8377
 Protein: ORFL147C - NC_006273.2[85932,87362]- <unknown description>
 Peptide score: 150.49
 Method: FTMS; HCD



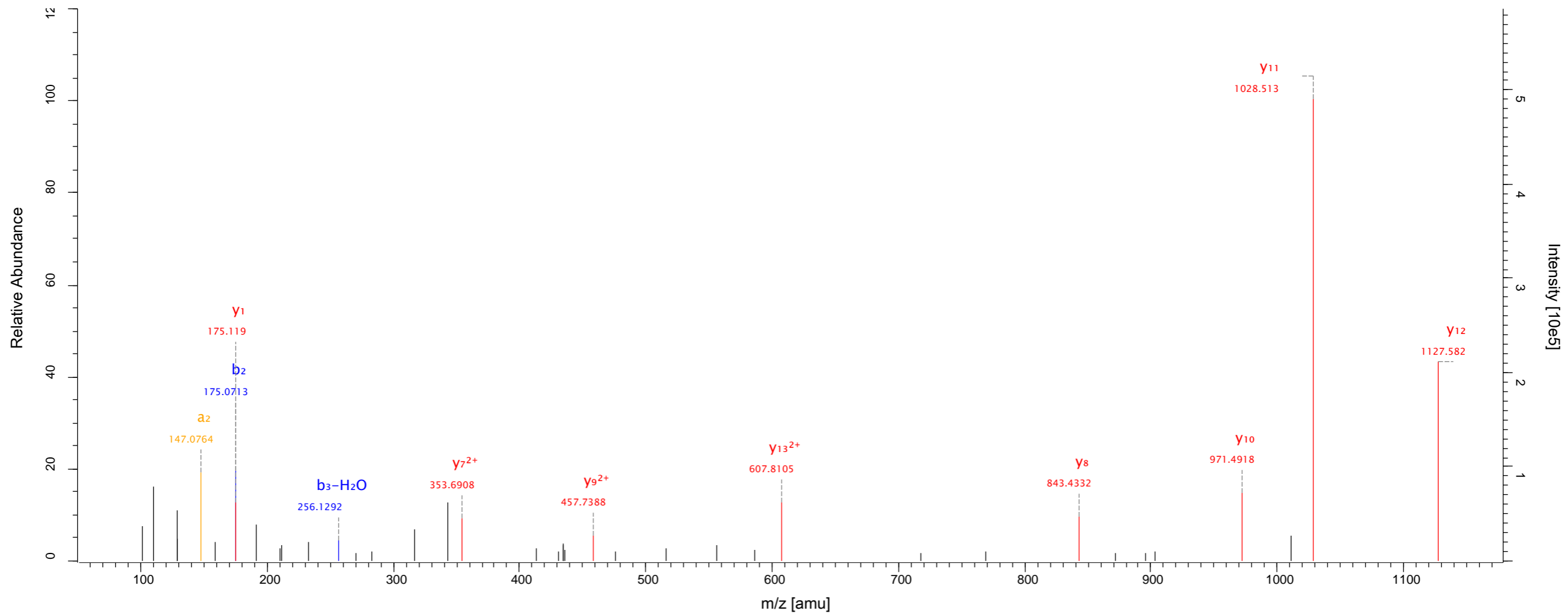
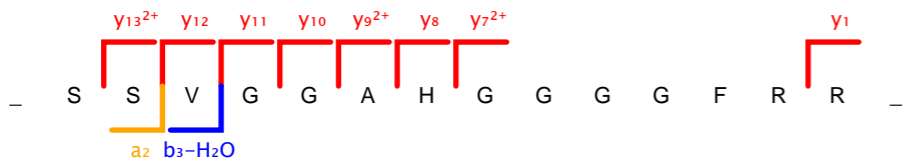
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 17385
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 124.08
Method: FTMS; HCD



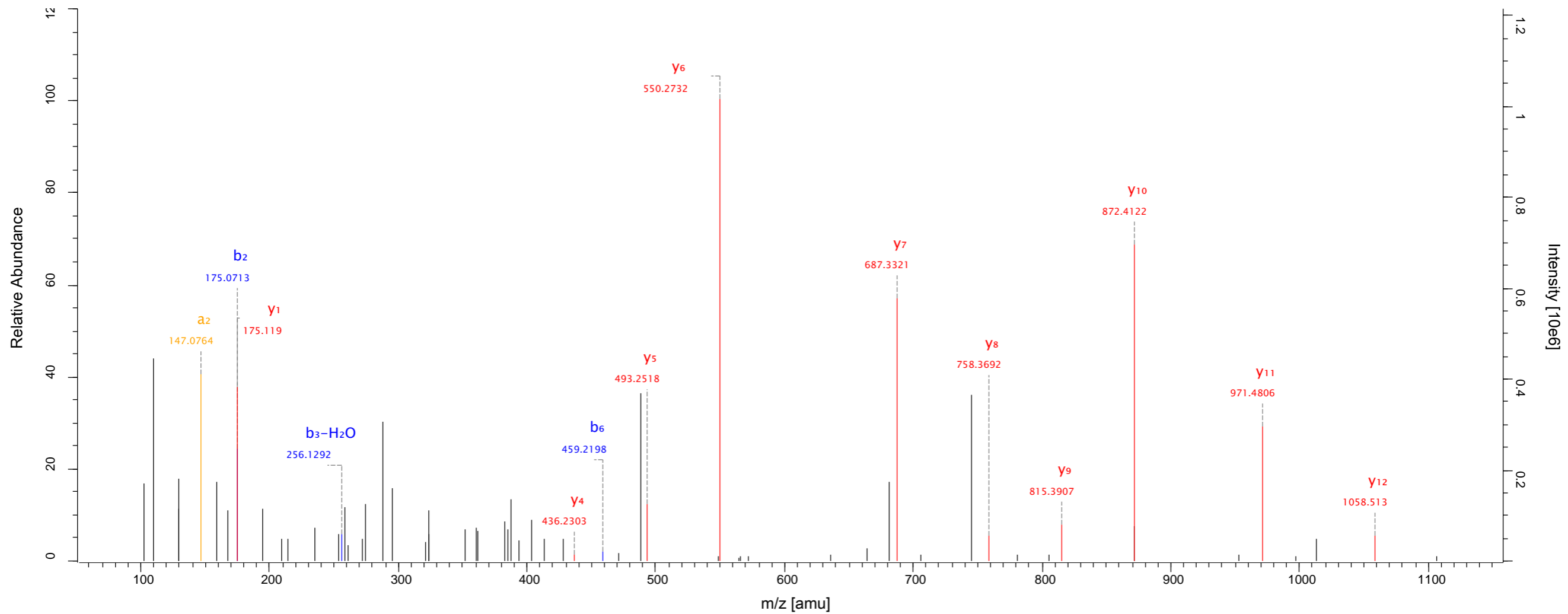
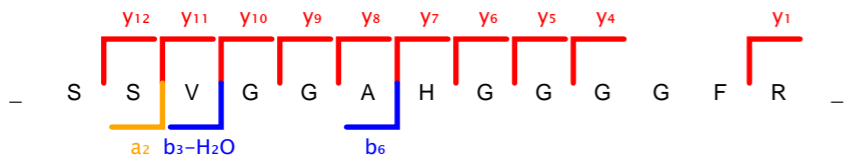
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 32768
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 81.16
Method: FTMS; HCD



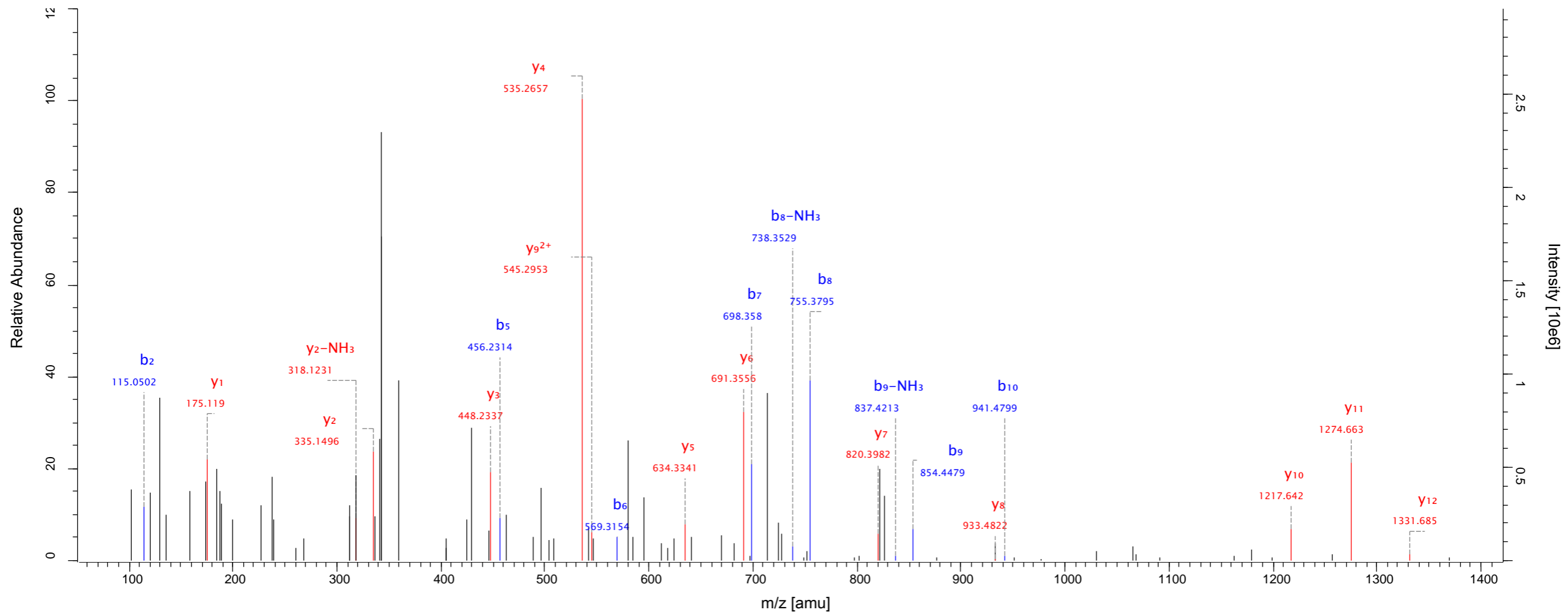
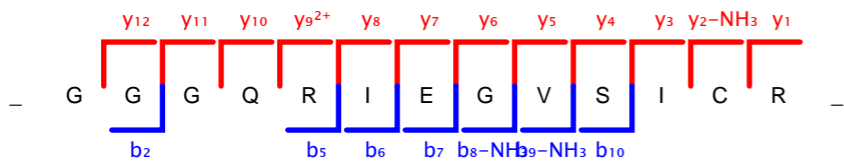
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_27
Scan number: 5509
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 63.73
Method: FTMS; HCD



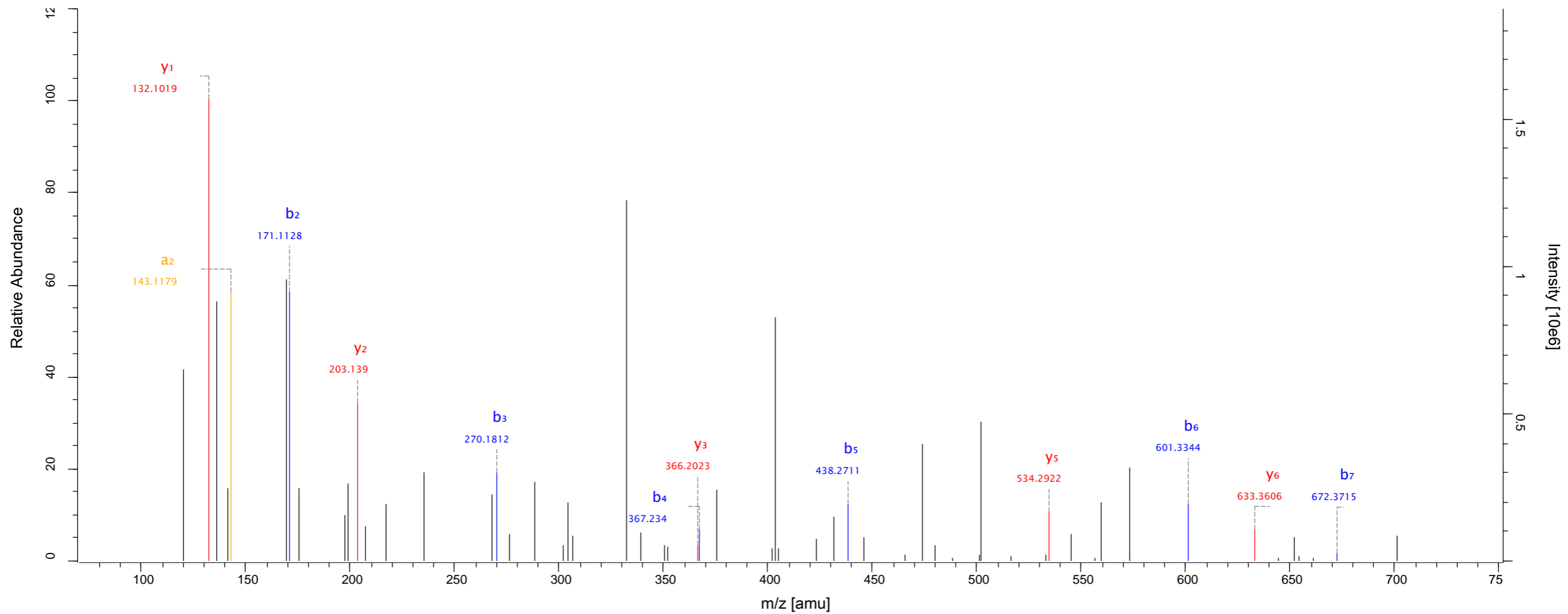
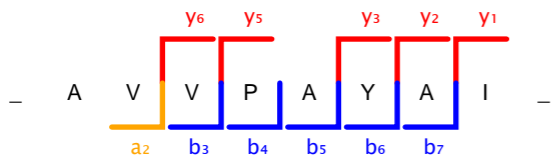
Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_27
Scan number: 6344
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 72.96
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 15524
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 104.17
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp1_35
Scan number: 33235
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 109.88
Method: FTMS; HCD



Raw file: 20111212_EXQ5_AnMi_QC_HCMVp11_35
Scan number: 8220
Protein: ORFL149C.iORF2 - NC_006273.2[91004,91471]- <unknown description>
Peptide score: 103.43
Method: FTMS; HCD

