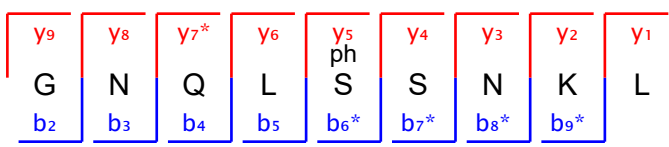
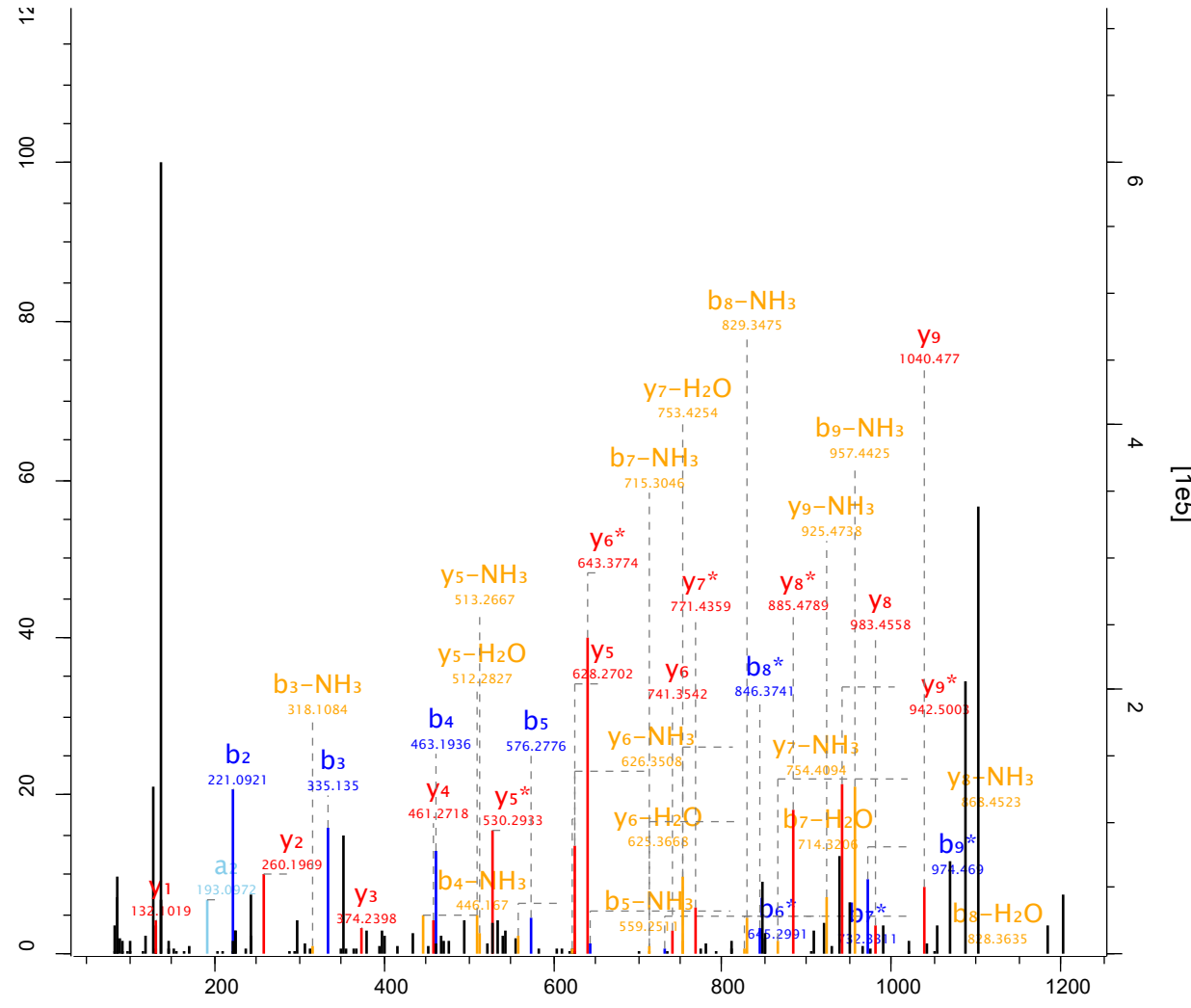
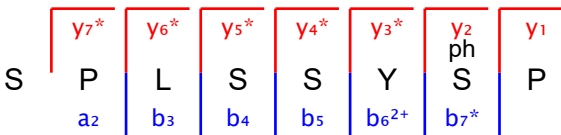
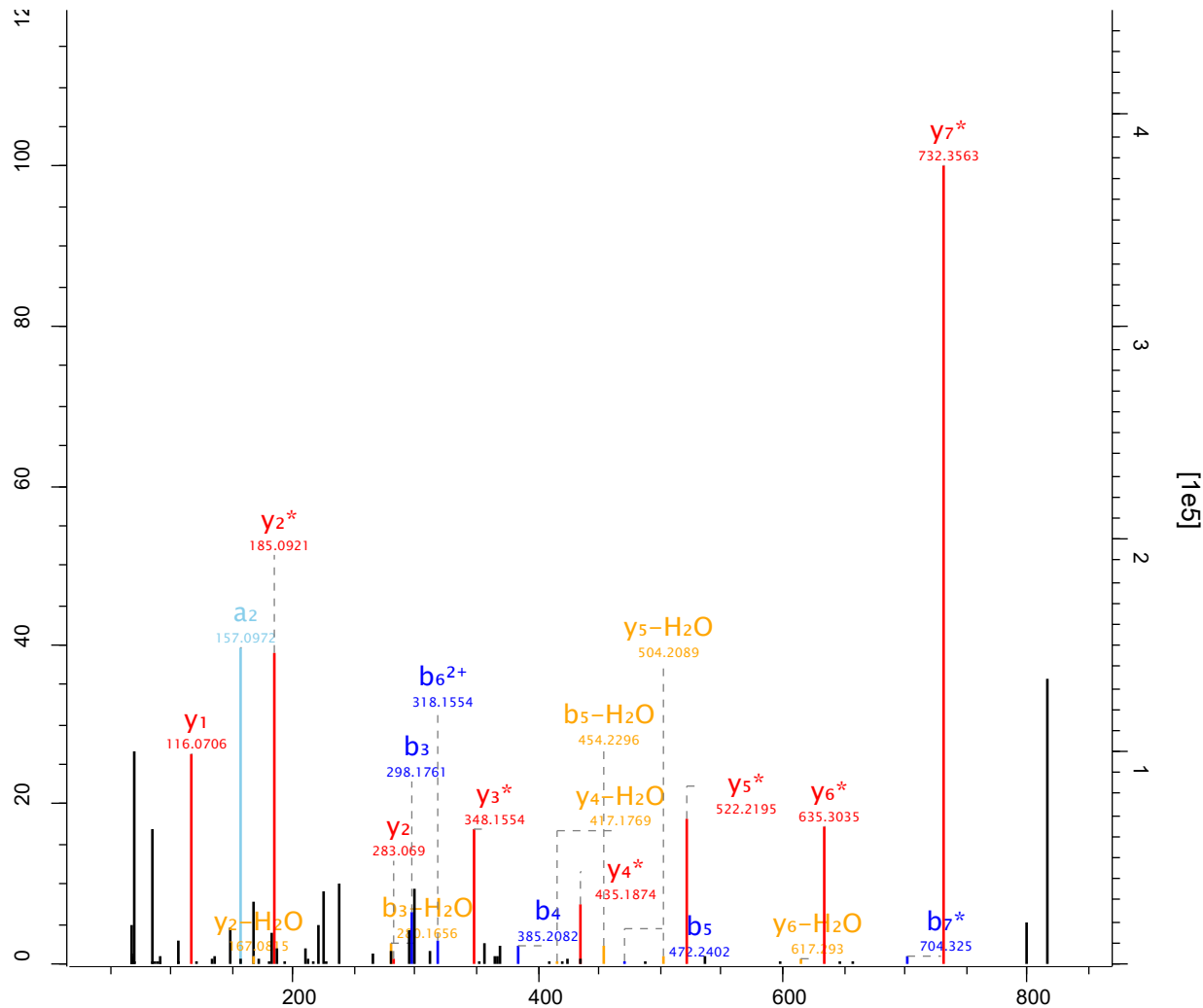


Supplementary Figure 2: Representative annotated spectra of identified phosphopeptides under systemin,A17 and water treatment as exported from MaxQuant.

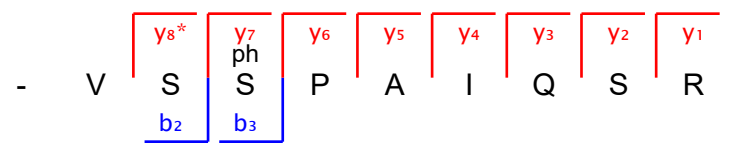
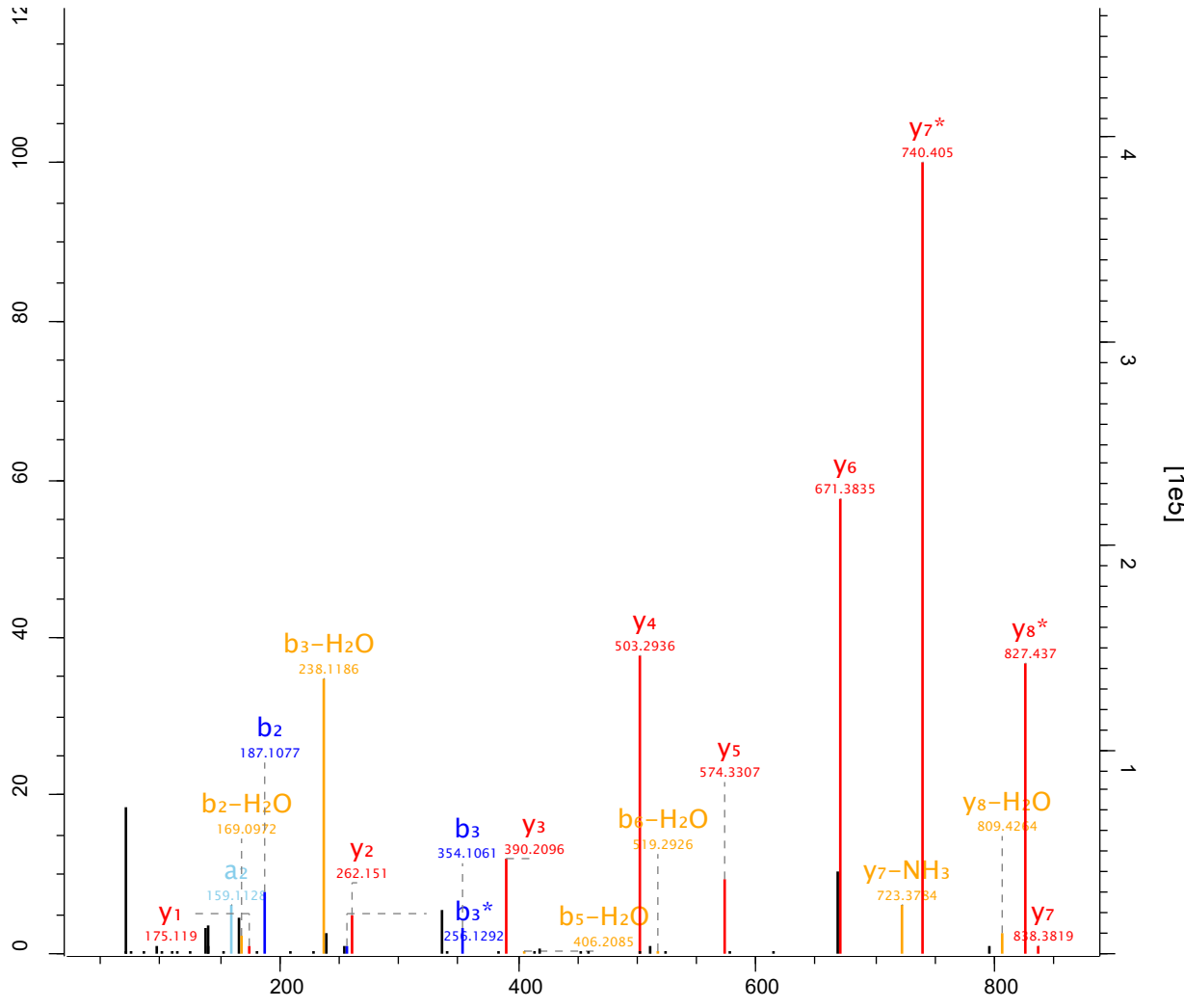
Raw file Scan Method Score m/z
 sys_30_2 10272 FTMS; HCD 218.01 602.27



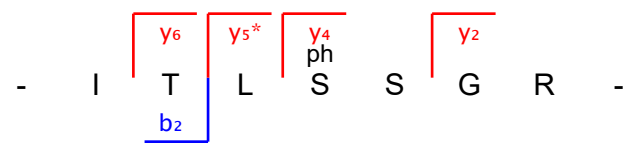
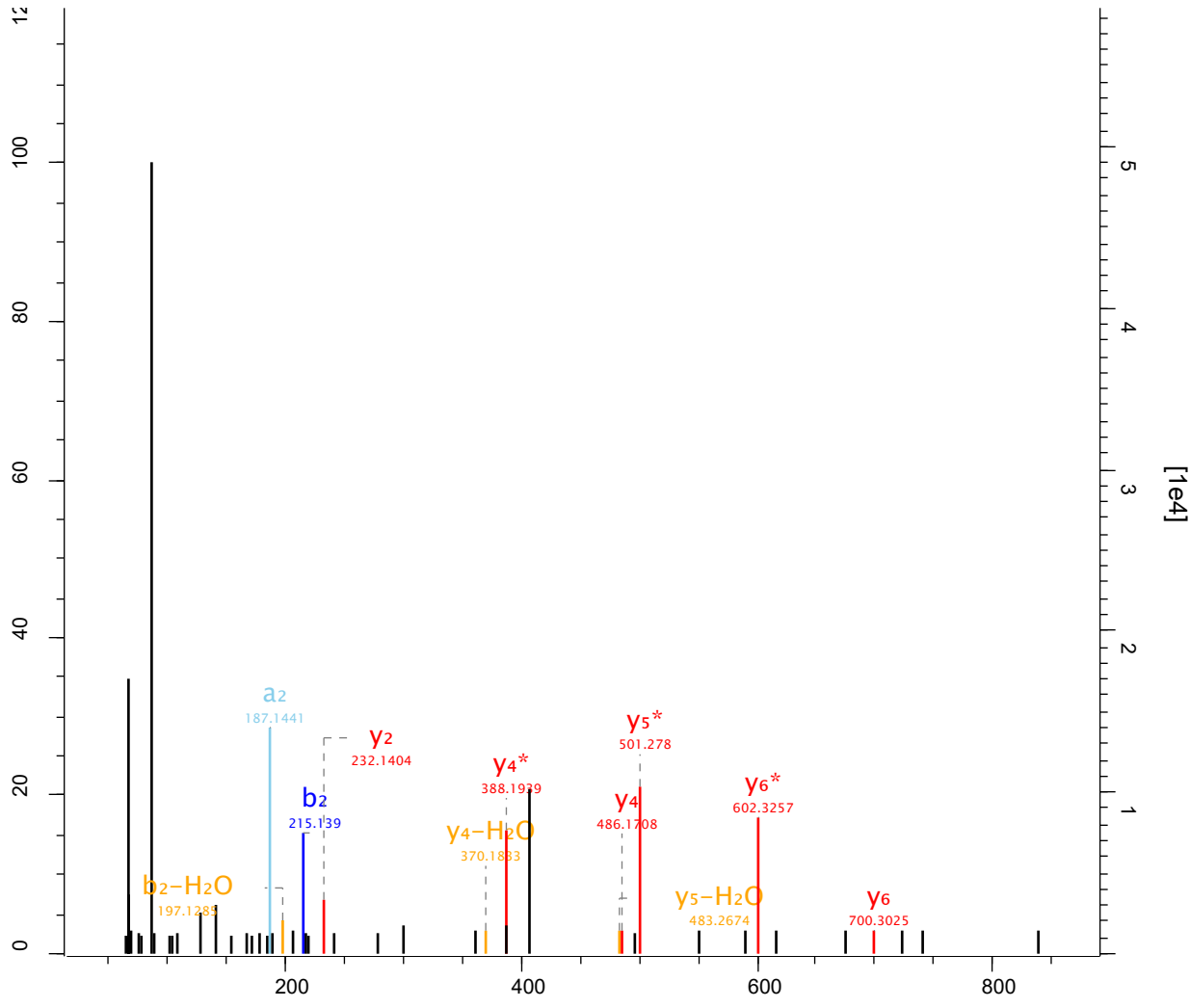
Raw file Scan Method Score m/z
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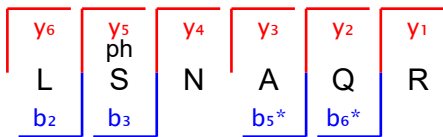
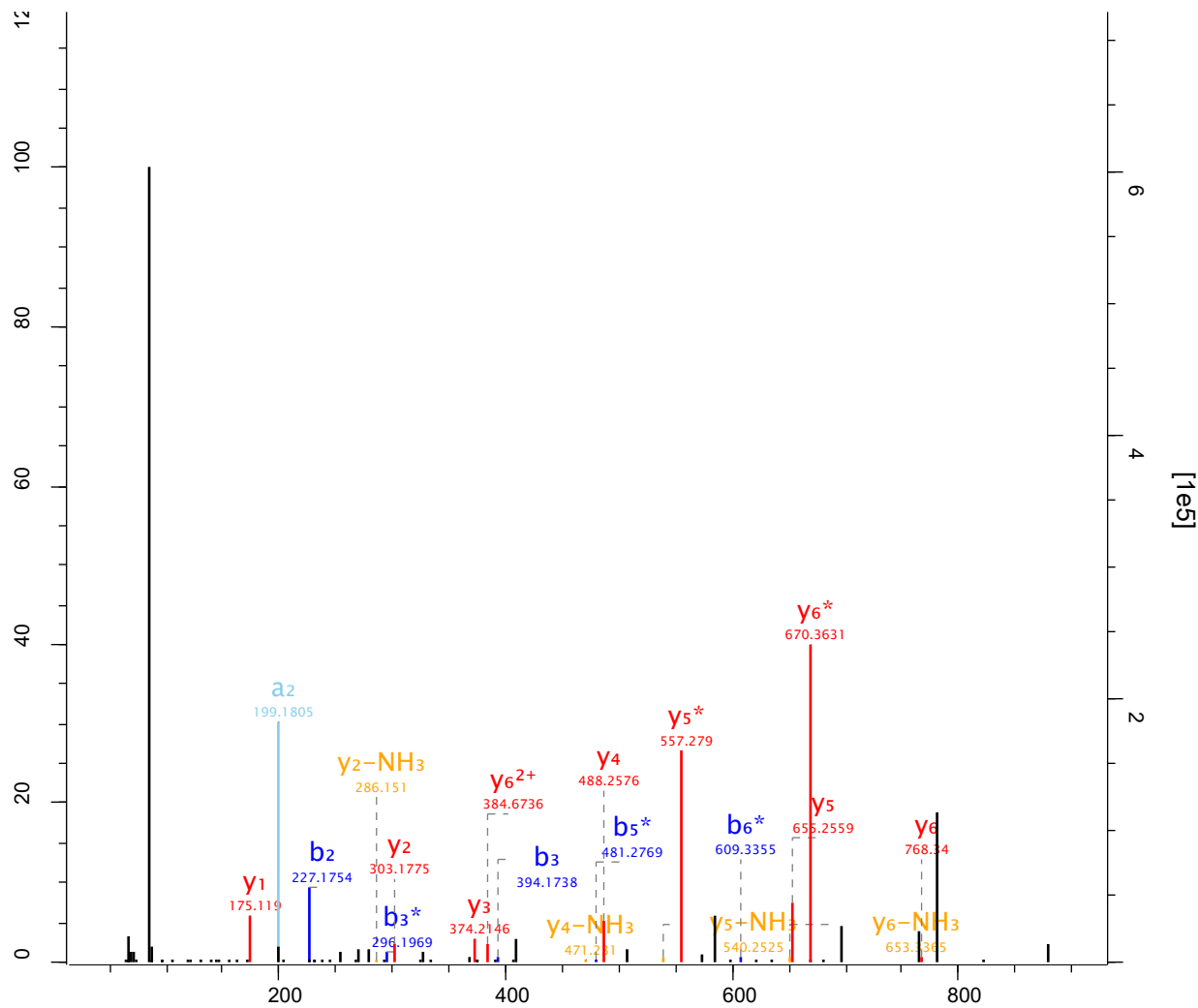
Raw file Scan Method Score m/z
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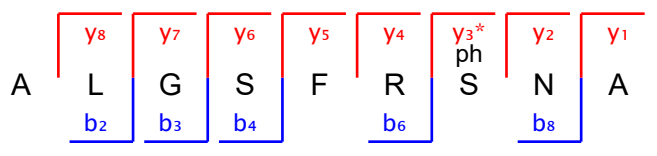
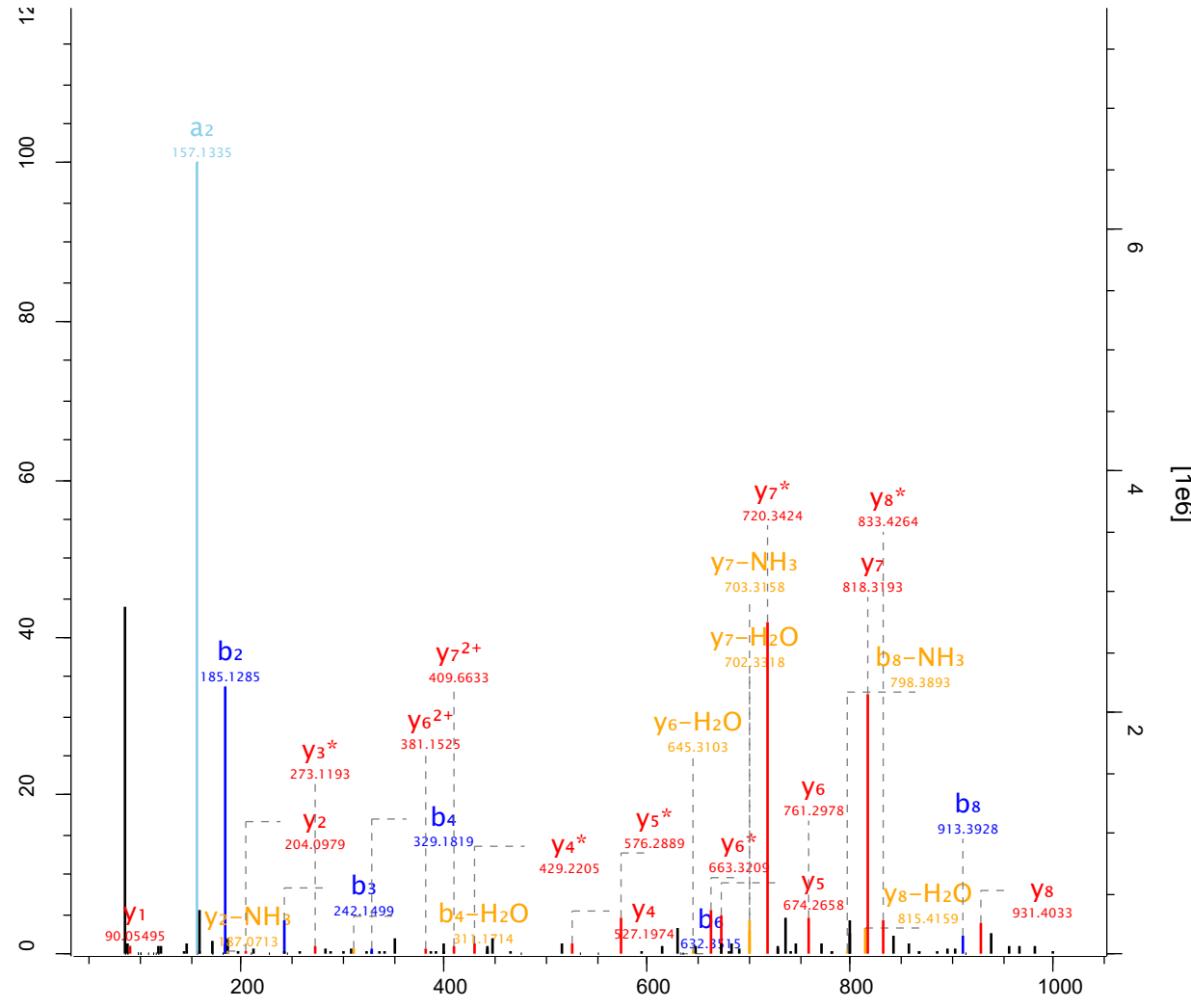
Raw file	Scan	Method	Score	m/z
sys_30_2	5526	FTMS; HCD	66.27	407.2



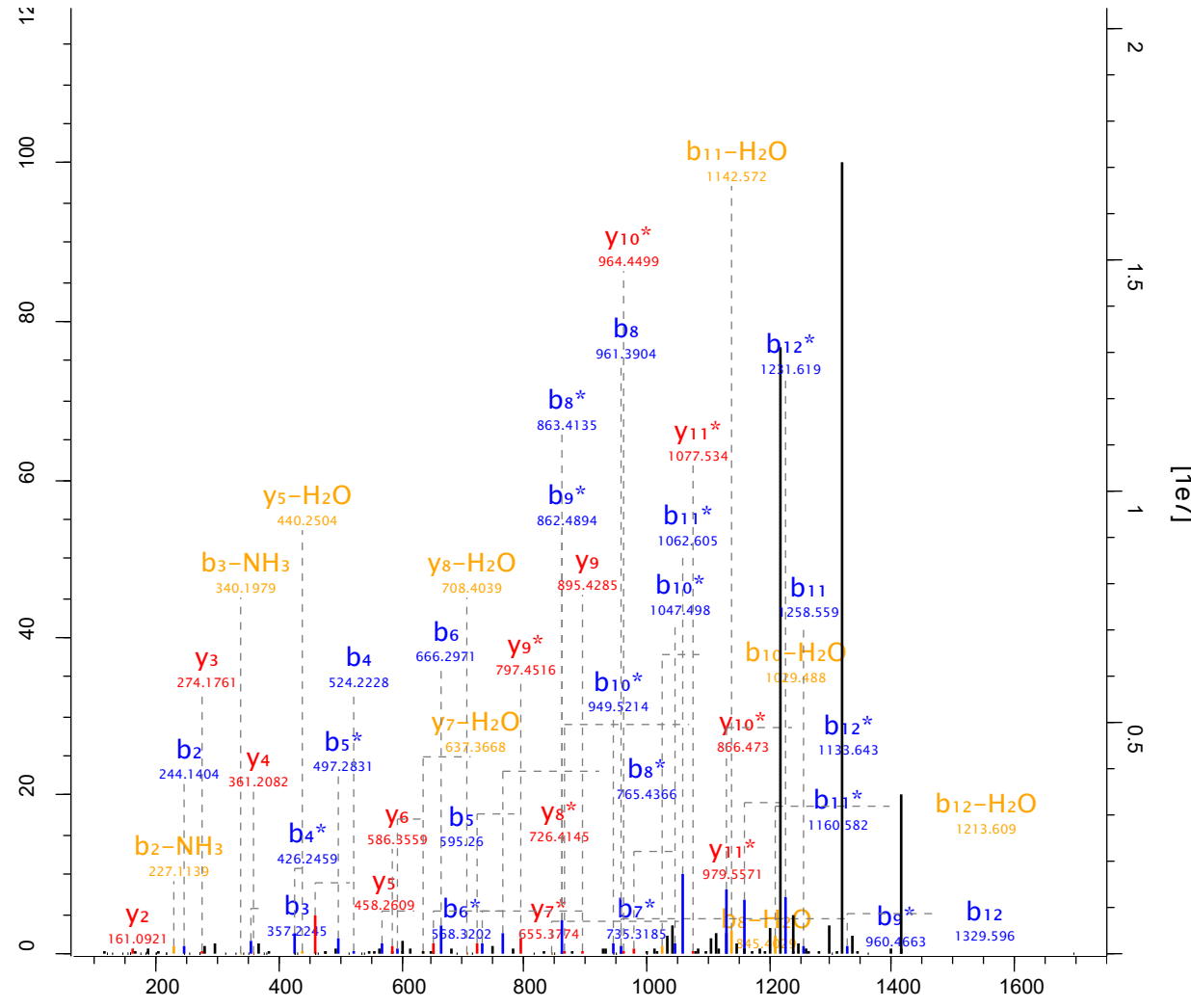
Raw file Scan Method Score m/z
 sys_30_2 6657 FTMS; HCD 119.75 441.22



Raw file Scan Method Score m/z
 sys_30_2 8708 FTMS; HCD 152.23 501.72



Raw file Scan Method Score m/z
 sys_30_3 10802 FTMS; HCD 150.13 709.83

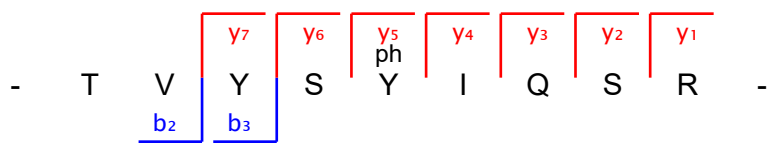
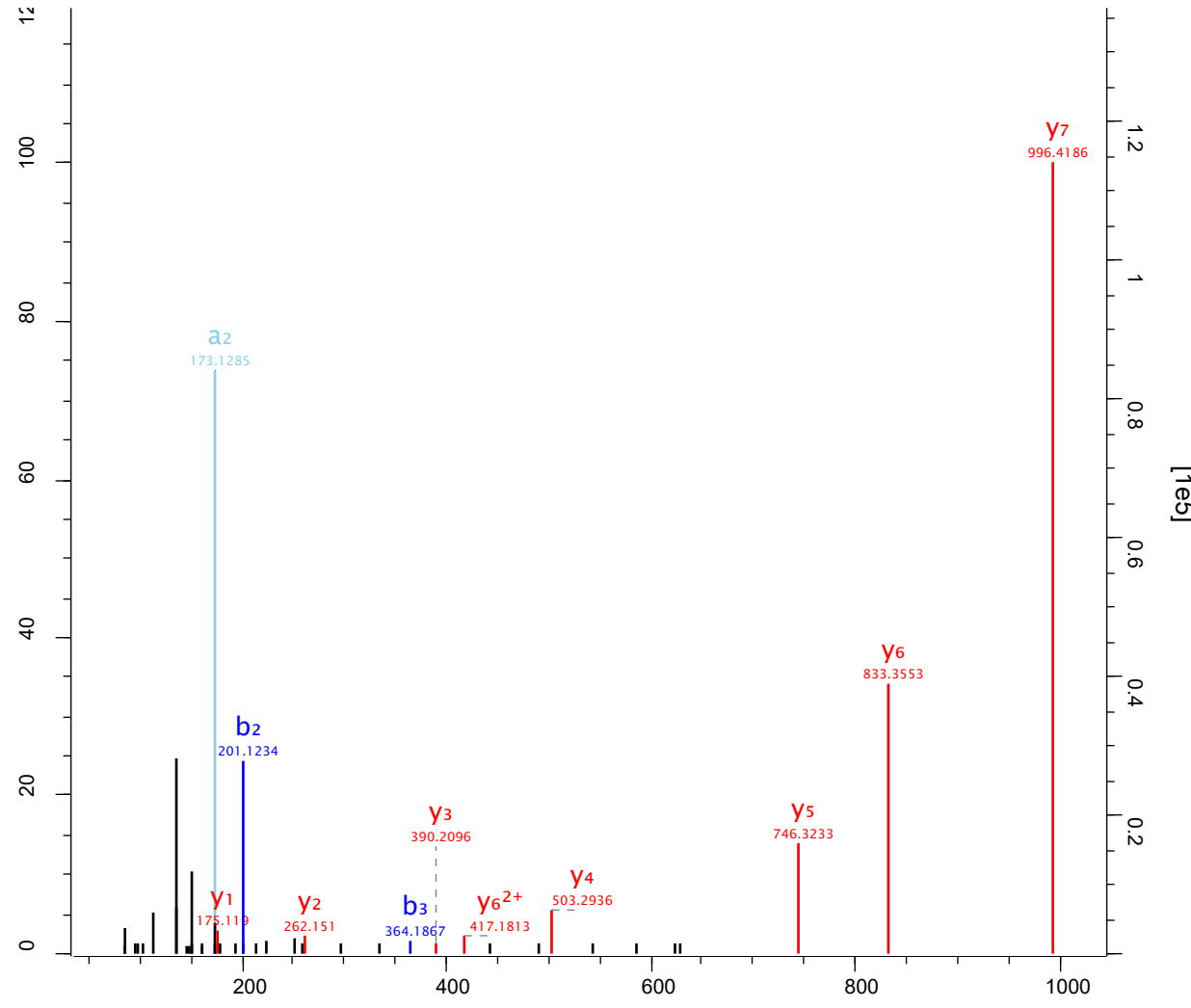


- S R L ph S A A S K P S I A A -

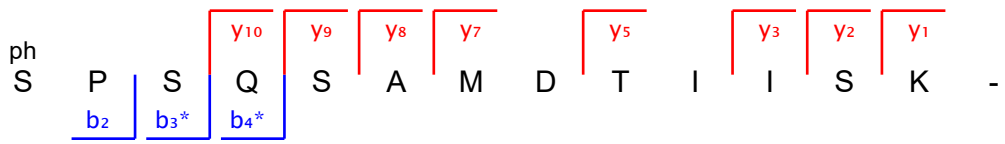
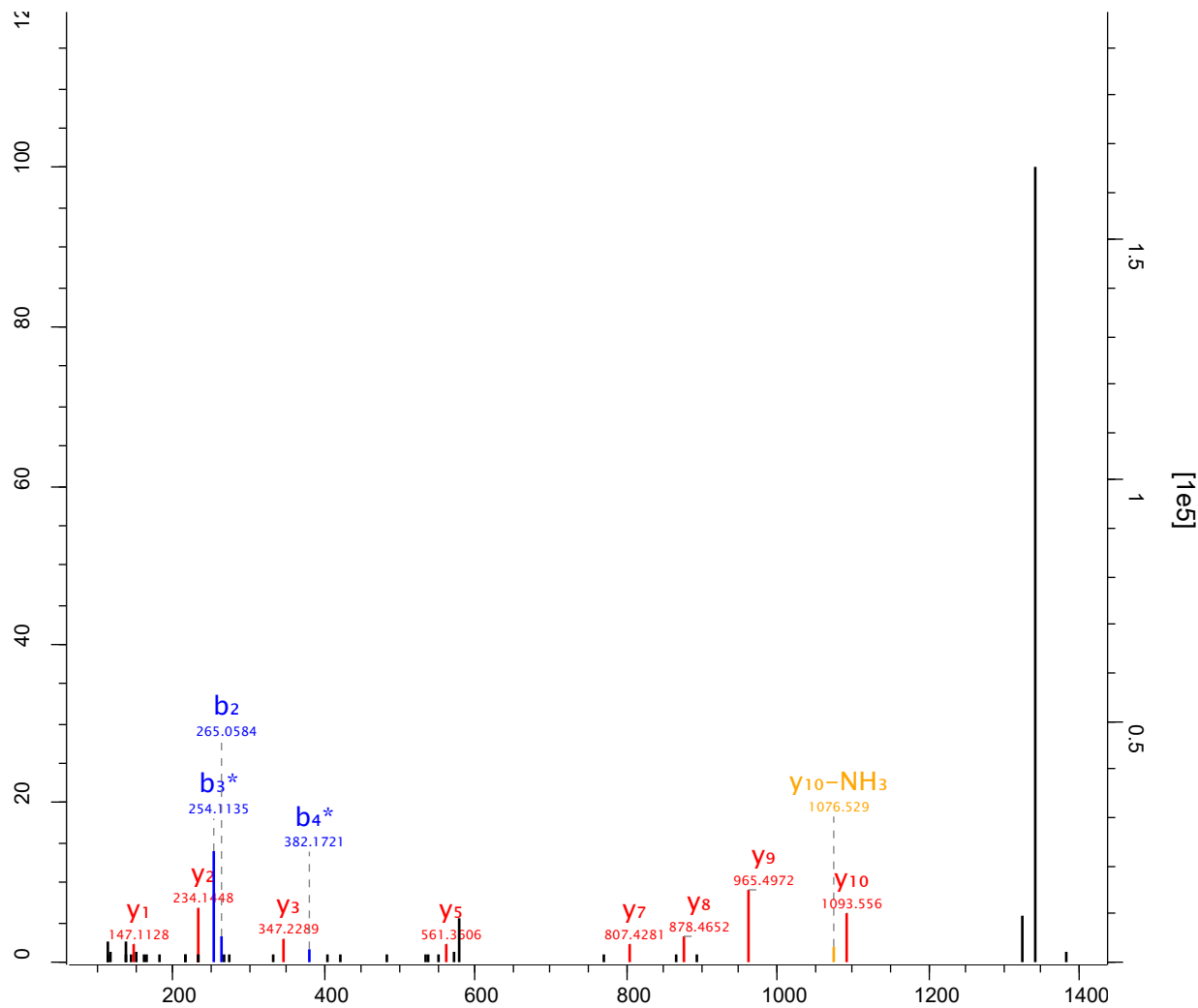
y11*
y10*
y9
y8*
y7*
y6
y5
y4
y3
y2

b2
b3
b4
b5
b6
b7*
b8
b9*
b10*
b11
b12

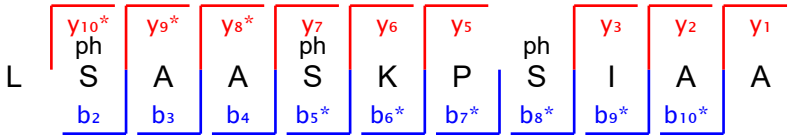
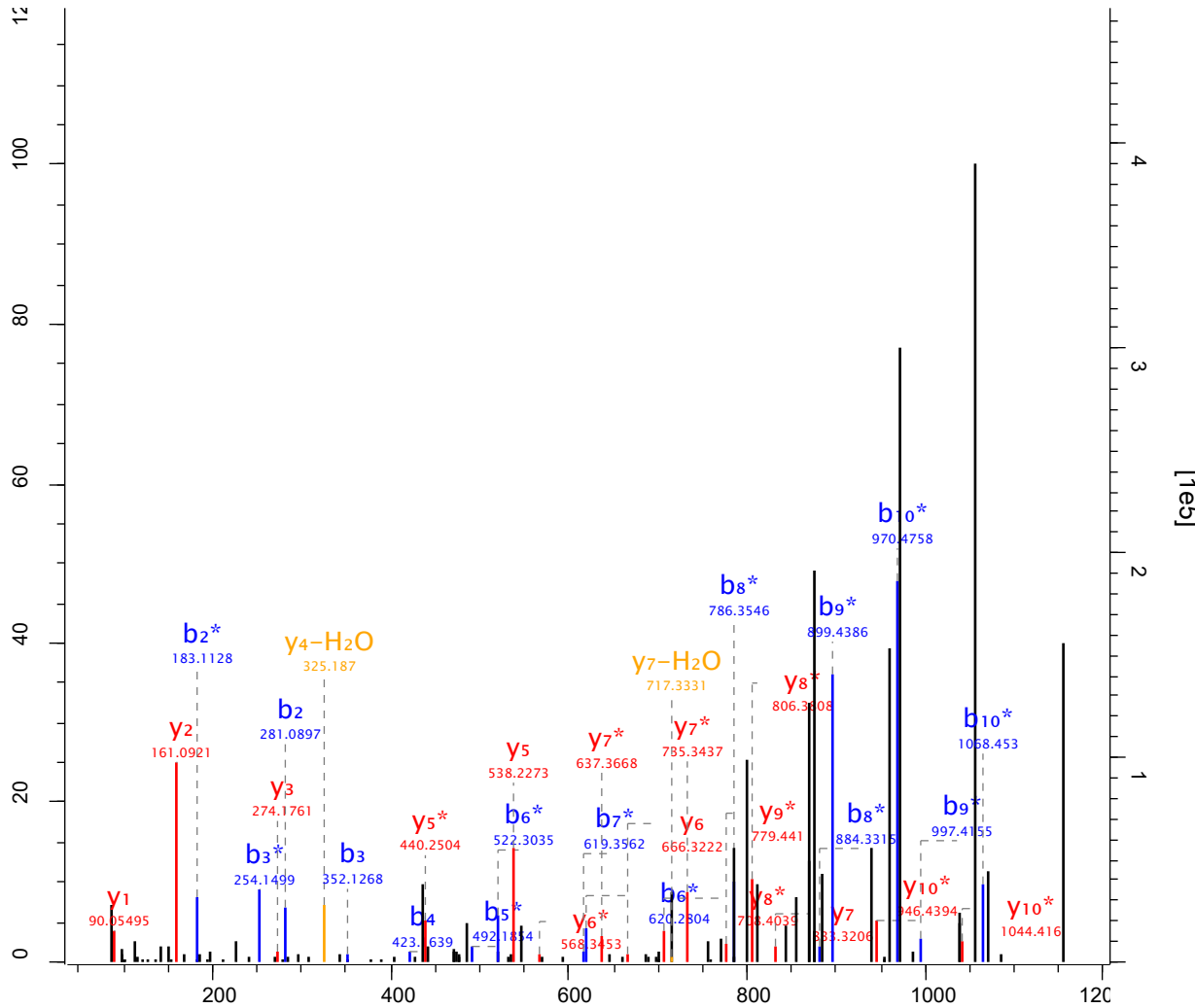
Raw file Scan Method Score m/z
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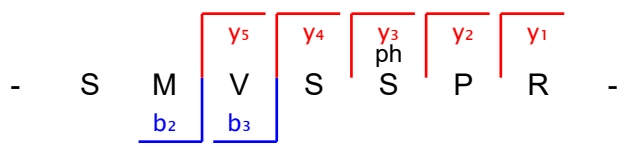
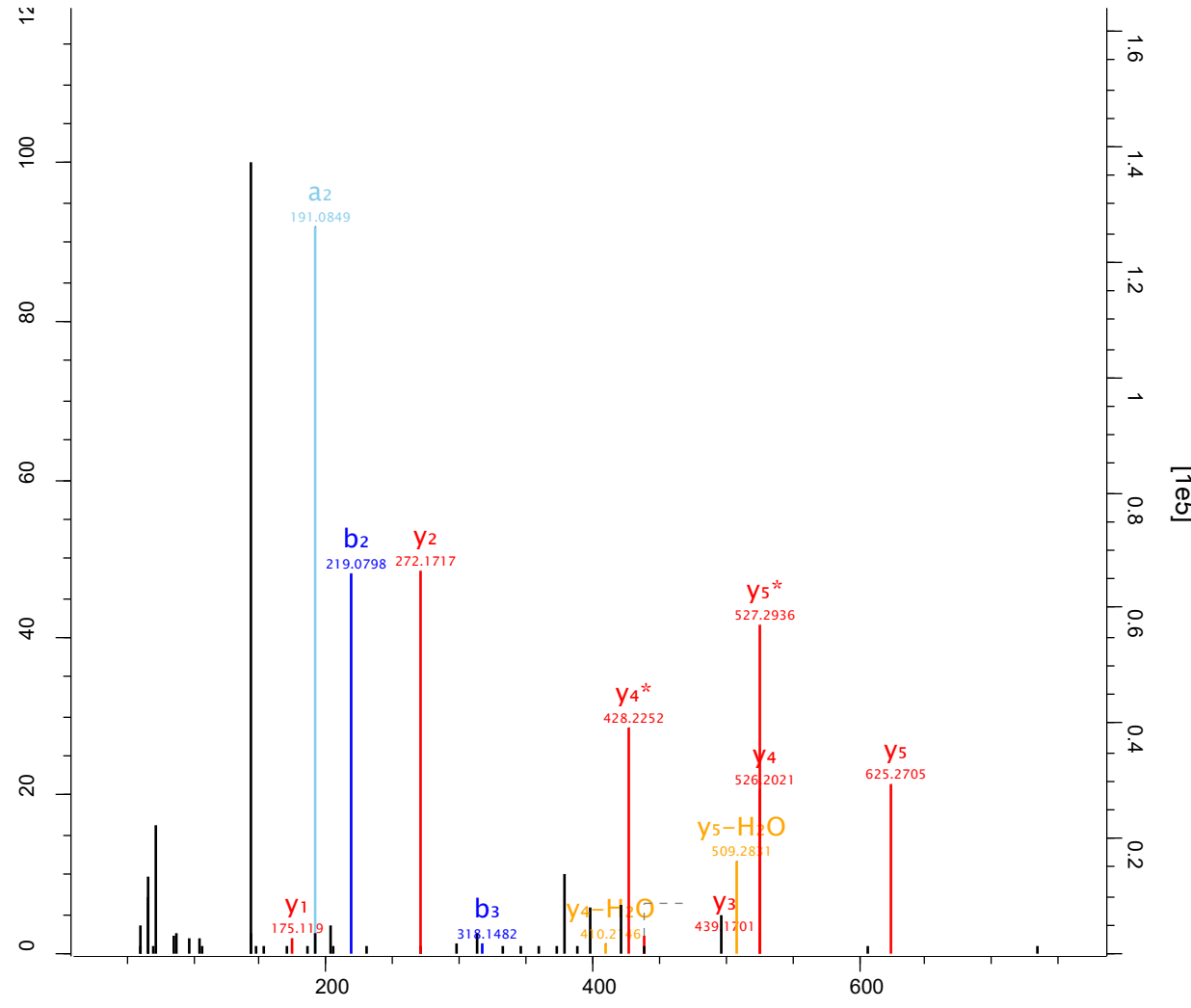
Raw file Scan Method Score m/z
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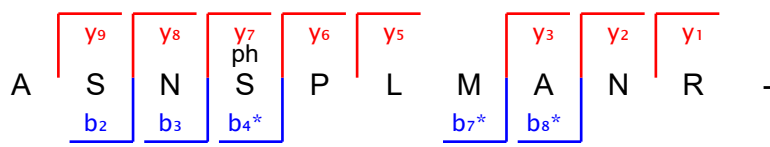
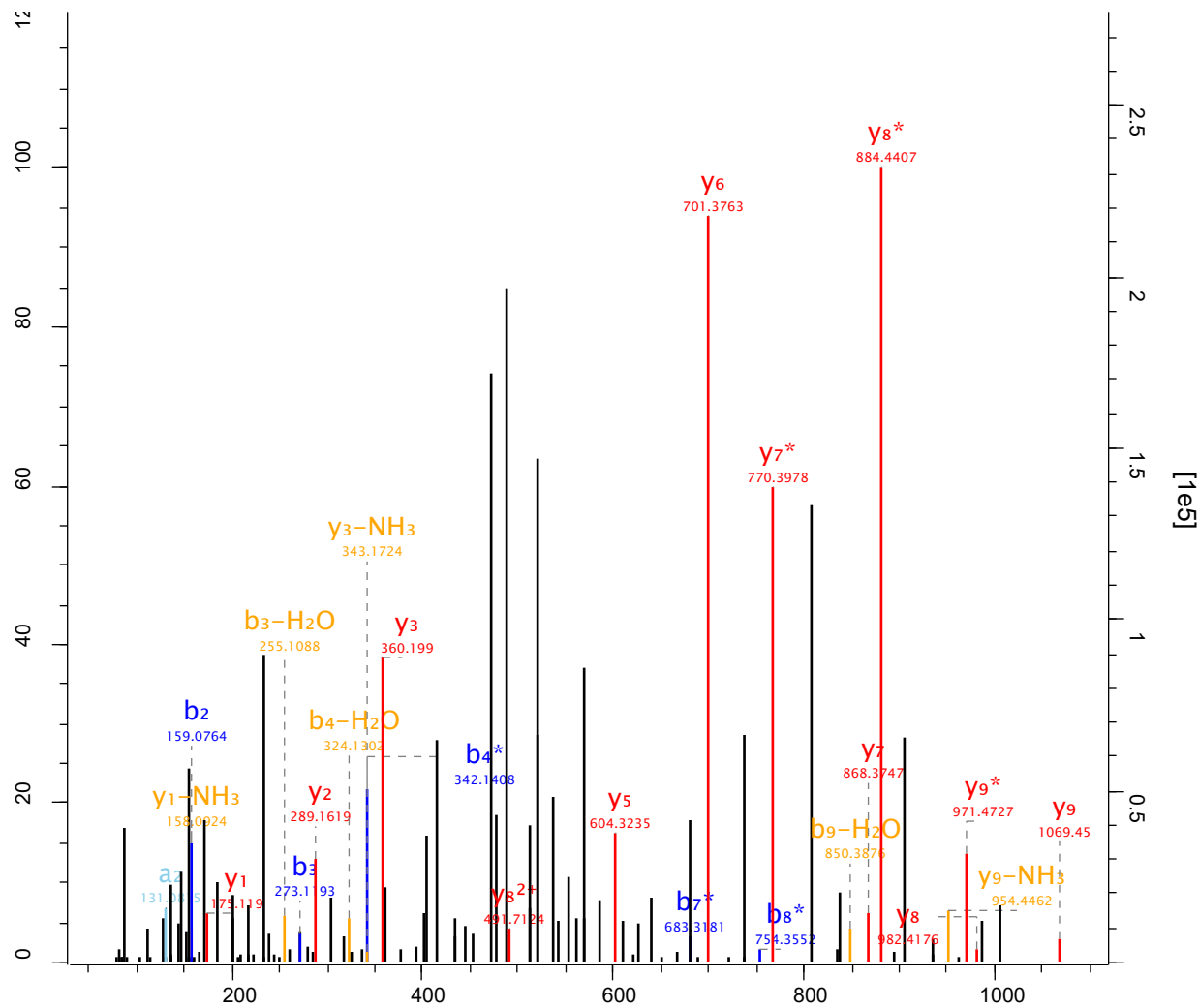
Raw file Scan Method Score m/z
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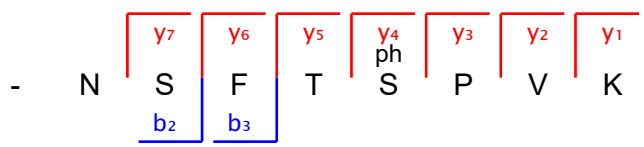
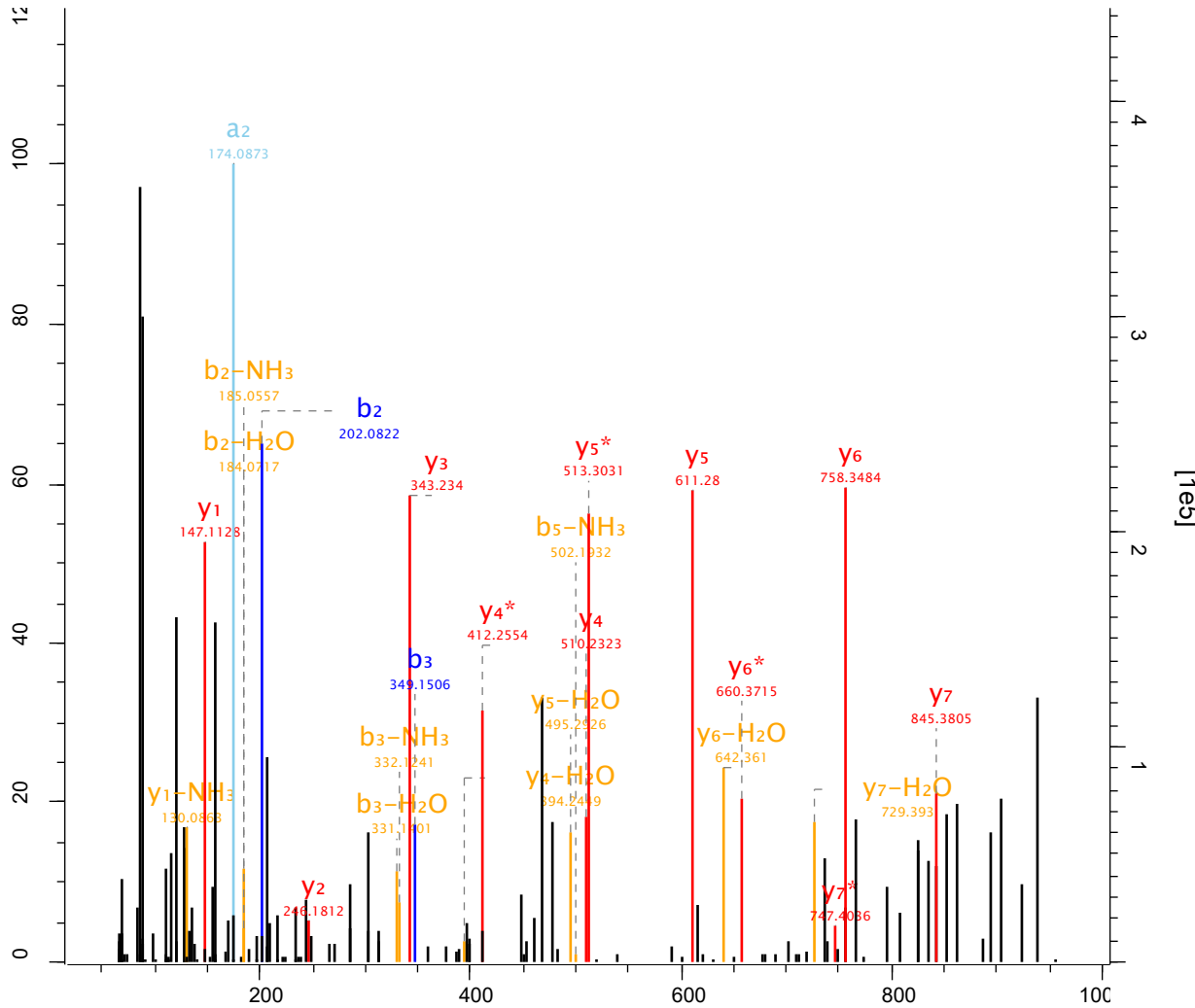
Raw file	Scan	Method	Score	m/z
sys_30_3	3325	FTMS; HCD	66.57	422.17



Raw file Scan Method Score m/z
 sys_30_3 5822 FTMS; HCD 78.81 570.75



Raw file Scan Method Score m/z
 sys_30_3 8493 FTMS; HCD 139.86 480.22



Raw file Scan Method Score m/z
 sys_30_3 9507 FTMS; HCD 140.31 621.28

