



Edit Structural Modification

Name:

OK
Cancel

Amino acid: Terminus: Variable

Chemical formula:

Monoisotopic mass: Average mass:

Loss <<

Neutral losses:

1175.3963 - H73C44N1O35	+
1378.4757 - H86C52N2O40	

Transitions

<input type="checkbox"/>	<input checked="" type="checkbox"/>		precursor [M-1] - 865.3488+++ (i rank 7) 7
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		precursor - 865.6832+++ (i rank 2) 2
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		precursor [M+1] - 866.0175+++ (i rank 1) 1
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		precursor [M+2] - 866.3518+++ (i rank 3) 3
<input type="checkbox"/>	<input checked="" type="checkbox"/>		precursor [M+3] - 866.6861+++ (i rank 4) 4
<input type="checkbox"/>	<input checked="" type="checkbox"/>		precursor [M+4] - 867.0203+++ (i rank 5) 5
<input type="checkbox"/>	<input checked="" type="checkbox"/>		precursor [M+5] - 867.3545+++ (i rank 6) 6
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		precursor -1175.4 - 473.8844+++
<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>		precursor -1378.5 - 406.1913+++
<input type="checkbox"/>	<input checked="" type="checkbox"/>		H [y11 -1175.4] - 1332.6066+
<input type="checkbox"/>	<input checked="" type="checkbox"/>		H [y11 -1378.5] - 1129.5273+
<input type="checkbox"/>	<input checked="" type="checkbox"/>		P [y10 -1175.4] - 1195.5477+

Can we monitor these ions with 1+ or 2+?