

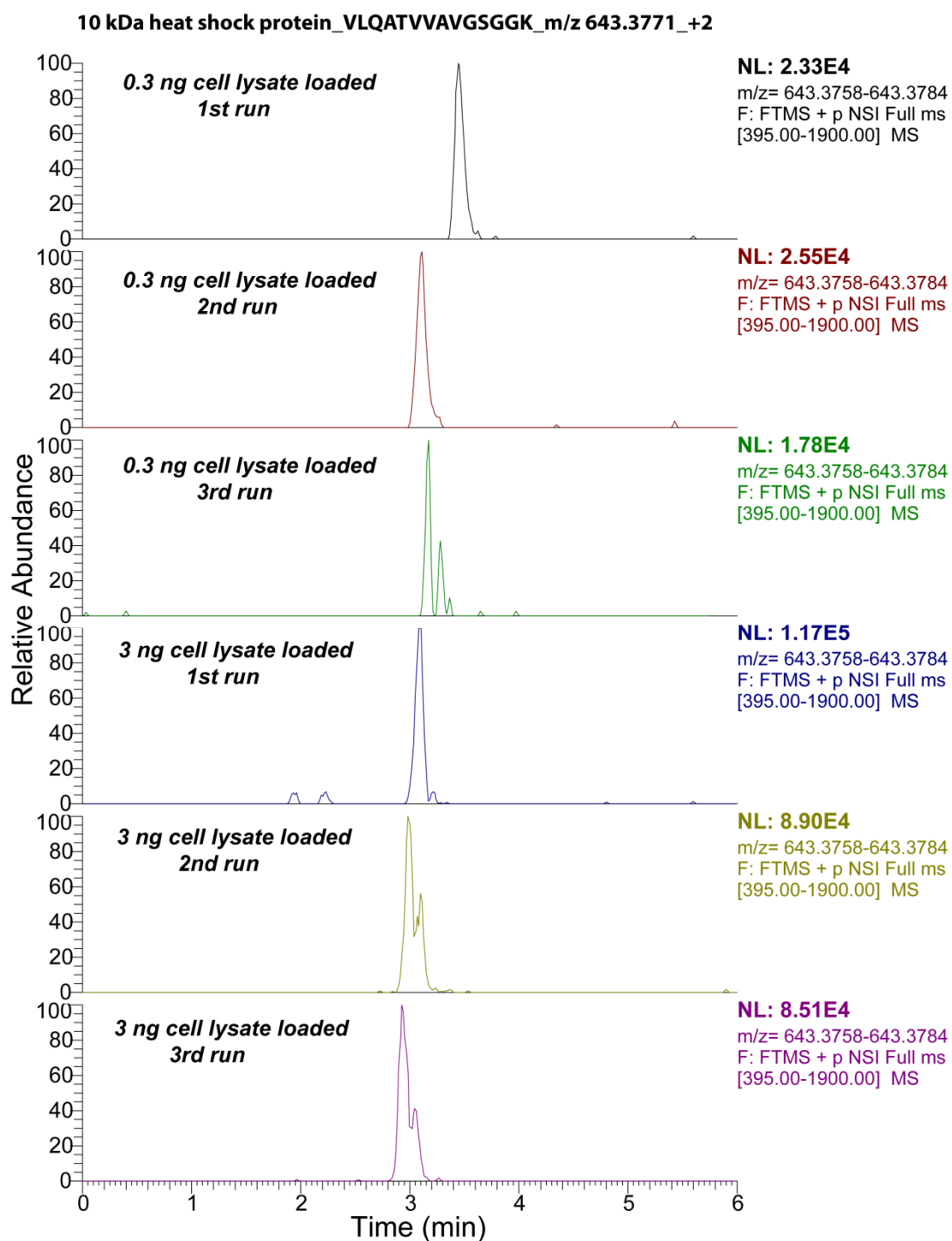
## **Supporting material II**

Integrated CZE-ESI-MS/MS system with an immobilized trypsin  
microreactor for online digestion and analysis of picogram amounts of RAW  
264.7 cell lysate

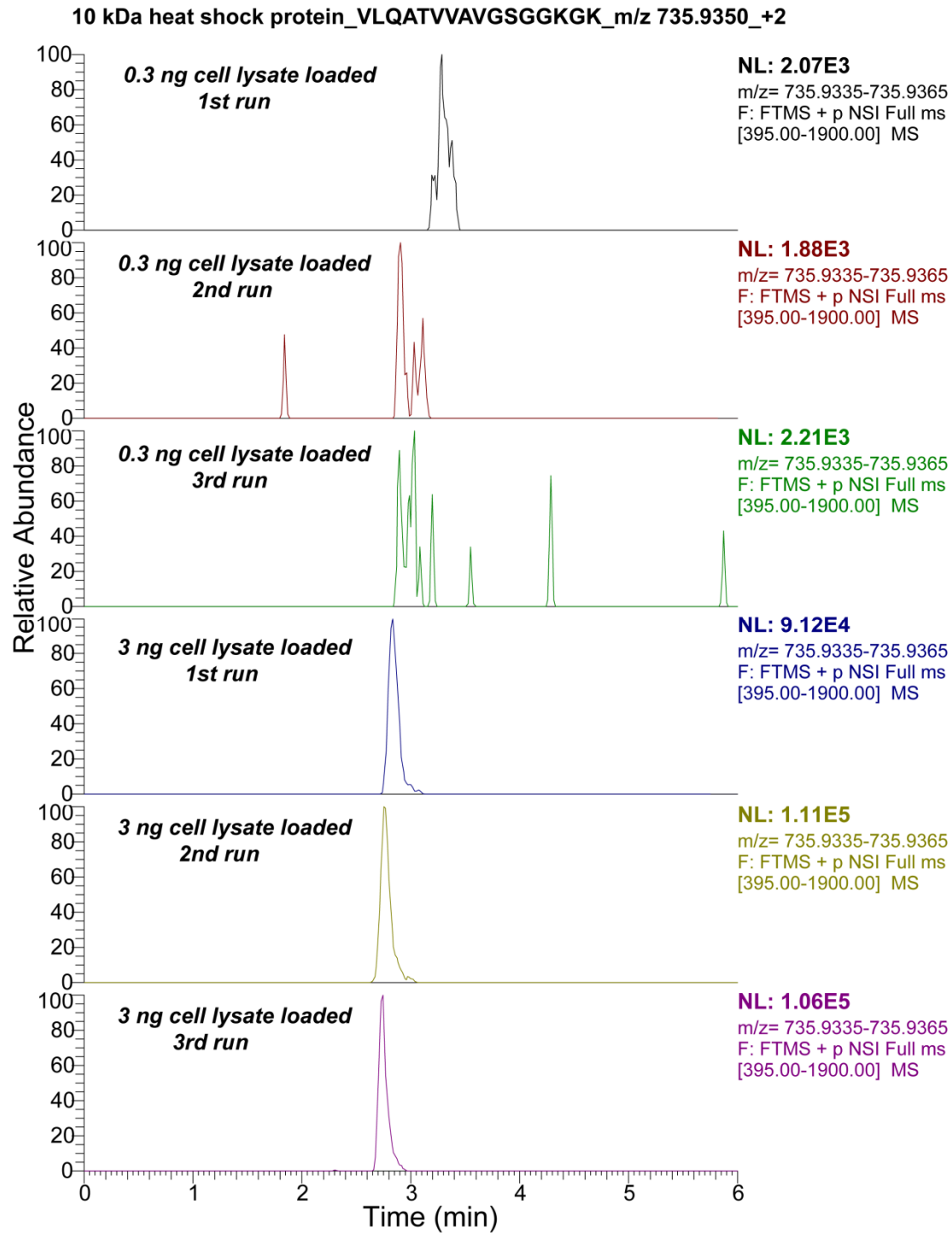
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Department of Chemistry and Biochemistry, University of Notre Dame, Notre Dame, IN  
46556, USA

**Part I S-Figure 1 and S-Figure 2**



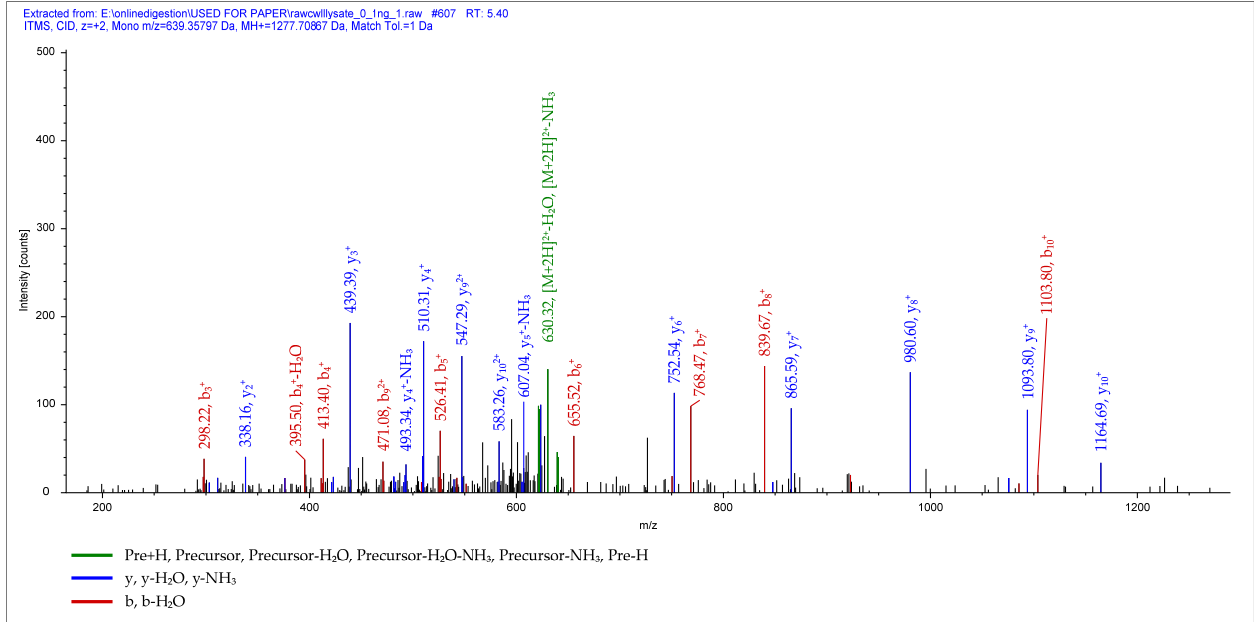
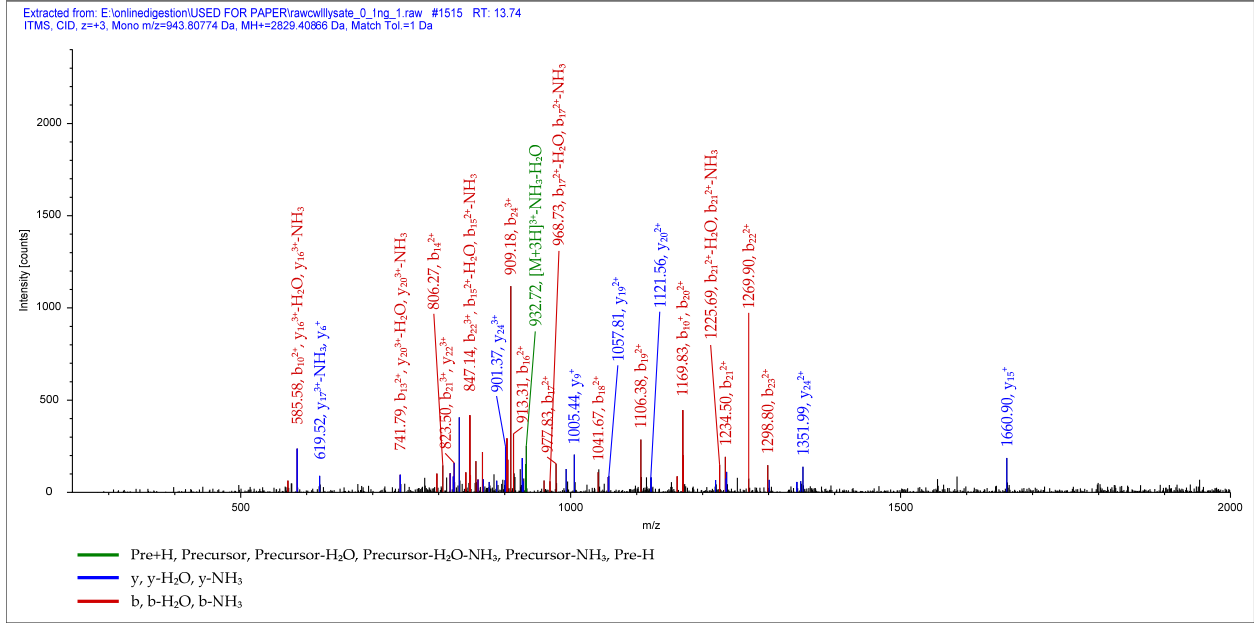
**S-Figure 1** Triplicate extracted ion electropherograms of a peptide (VLQATVVAVGSGGK) that was only identified in the tandem spectra from 0.3 ng cell lysate. The peptide's ion intensity was extracted with mass tolerance as 2 ppm, and Gaussian smoothing with 5 points was applied.

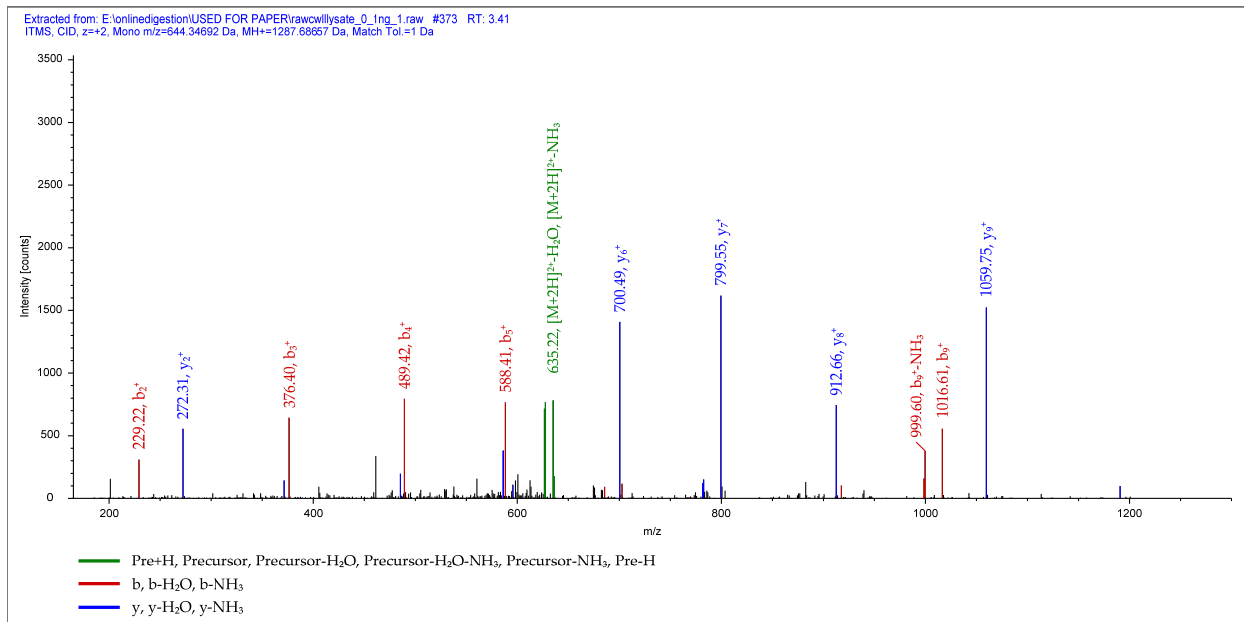


**S-Figure 2** Triplicate extracted ion electropherograms of a peptide (VLQATVVAVGSGGK) that was only identified in the tandem spectra from 3 ng cell lysate. The peptide was extracted with mass tolerance as 2 ppm, and Gaussian smoothing with 5 points was applied.

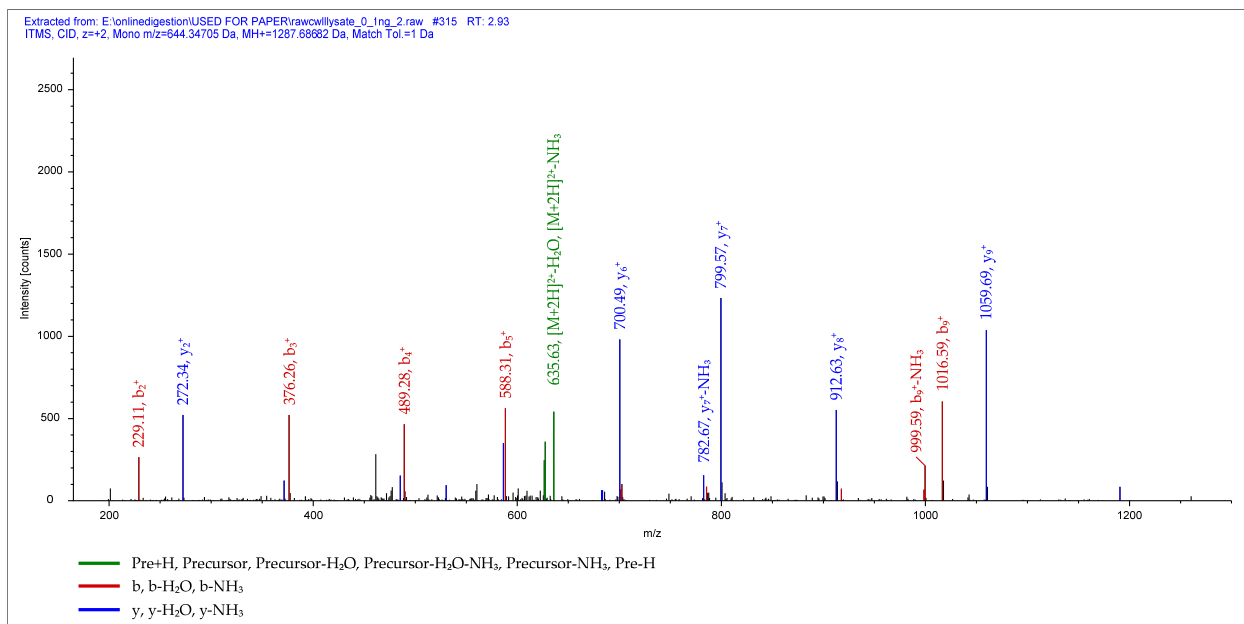
**Part II Annotated tandem spectra of identified peptides from 0.3 ng and 3 ng RAW 264.7 cell lysate analyzed by the integrated CZE-ESI-MS/MS system in triplicate runs**

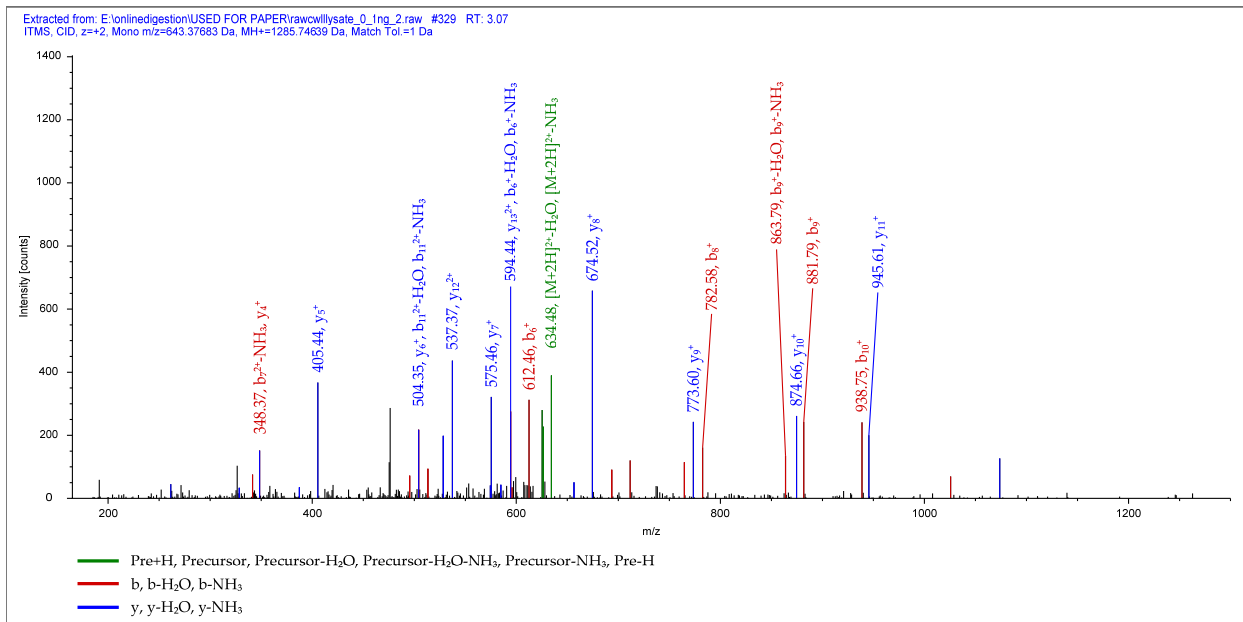
# 0.3 ng RAW 264.7 cell lysate\_1st run



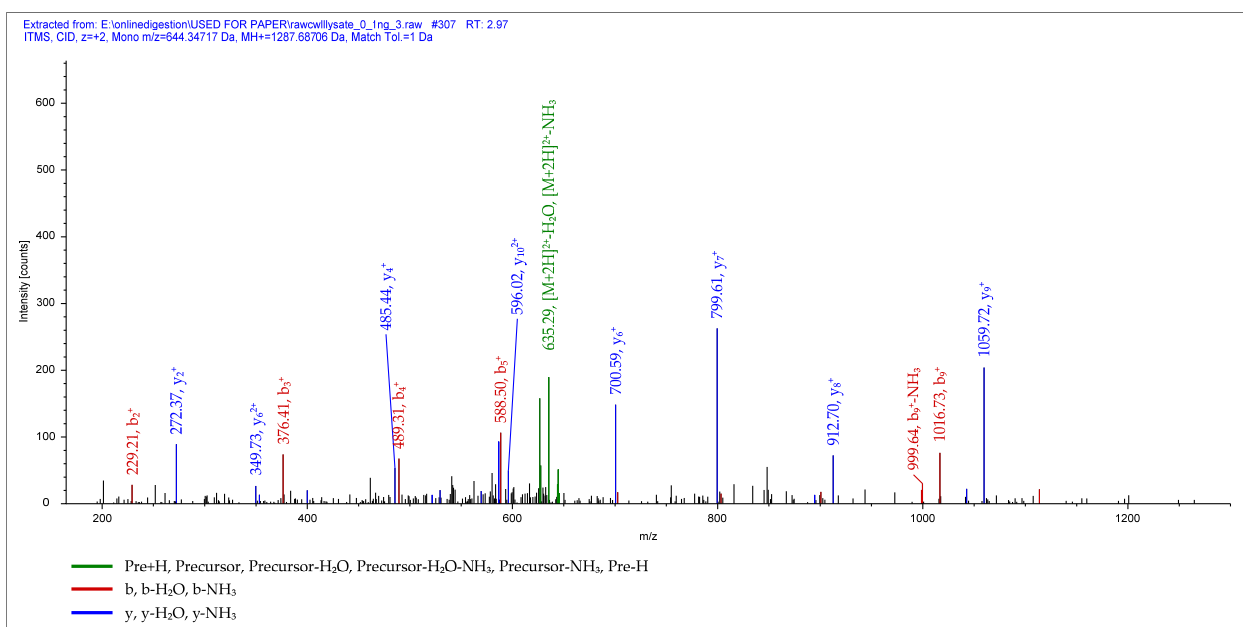


### 0.3 ng RAW 264.7 cell lysate\_2nd run

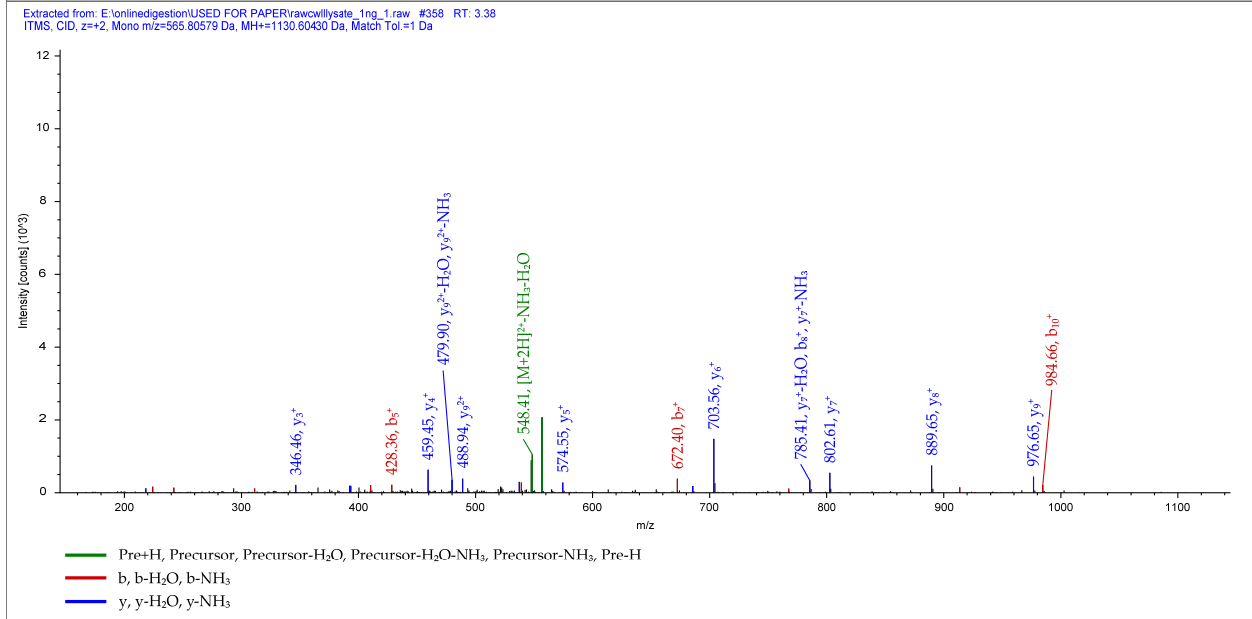
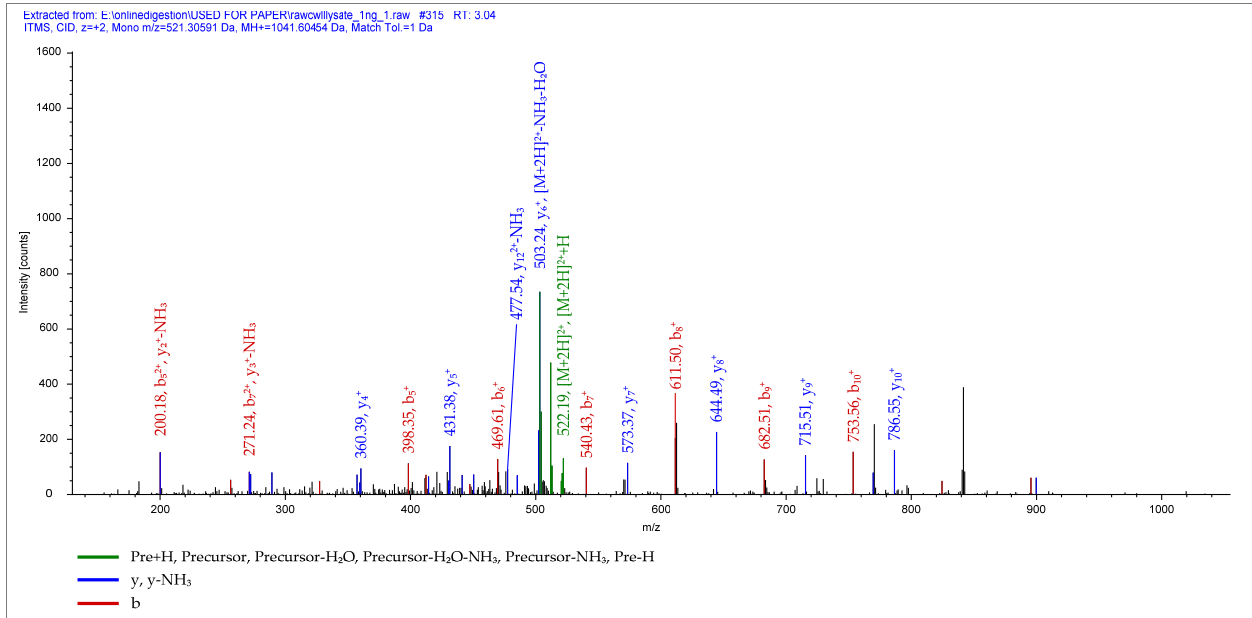




### 0.3 ng RAW 264.7 cell lysate\_3rd run

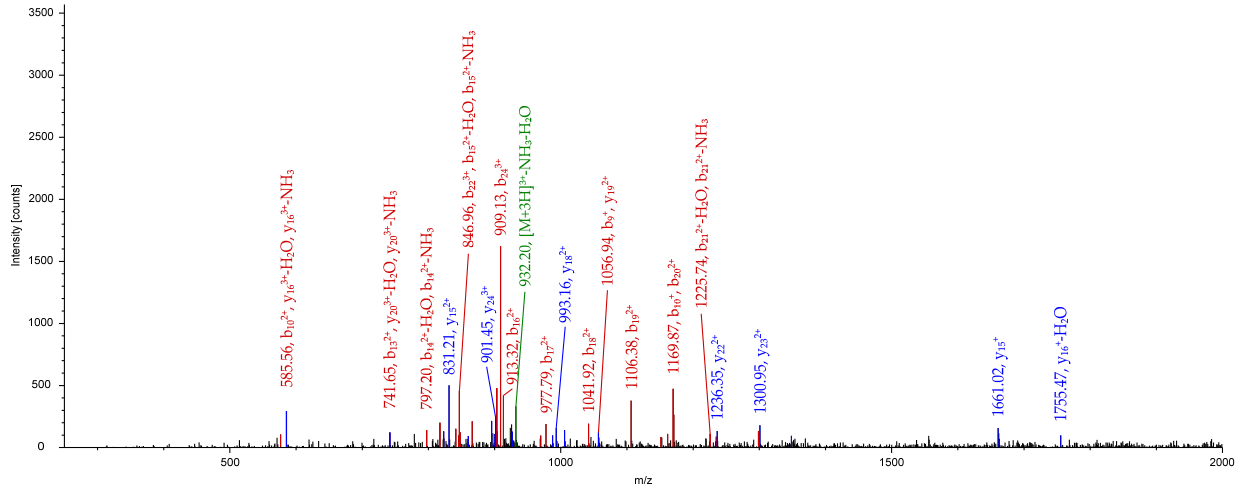


### 3 ng RAW 264.7 cell lysate\_1st run



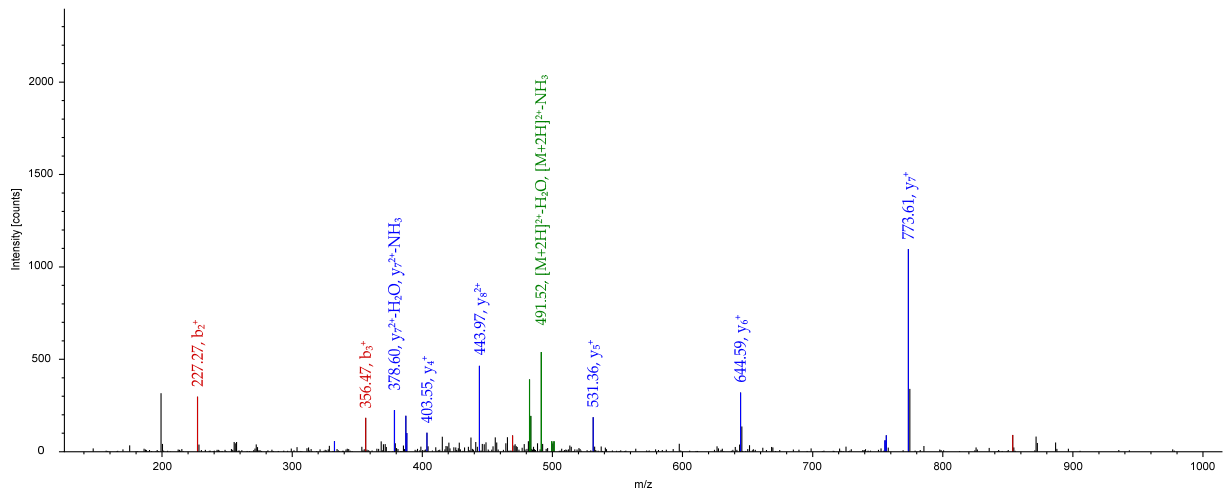


Extracted from: E:\onlinedigestion\USED FOR PAPER\rawcwllylsate\_1ng\_1.raw #481  
ITMS, CID, z=+3, Mono m/z=943.80908 Da, MH+=2829.41269 Da, Match Tol.=1 Da



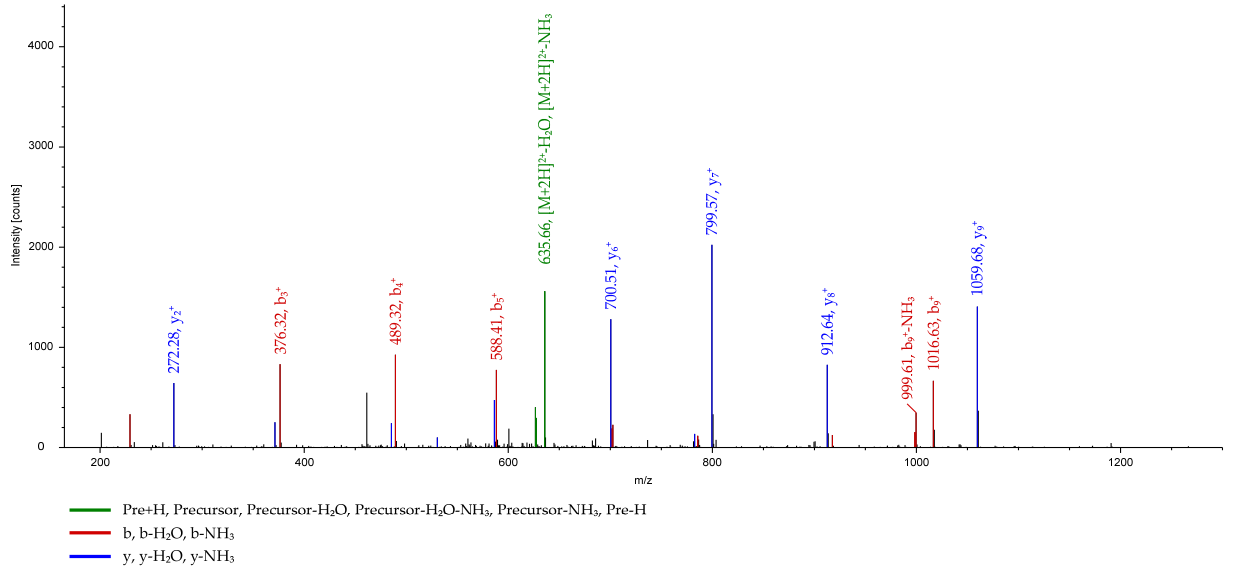
— Pre+H, Precursor, Precursor-H<sub>2</sub>O, Precursor-H<sub>2</sub>O-NH<sub>3</sub>, Precursor-NH<sub>3</sub>, Pre-H  
— y, y-H<sub>2</sub>O, y-NH<sub>3</sub>  
— b, b-H<sub>2</sub>O, b-NH<sub>3</sub>

Extracted from: E:\onlinedigestion\USED FOR PAPER\rawcwllylsate\_1ng\_1.raw #319 RT: 3.06  
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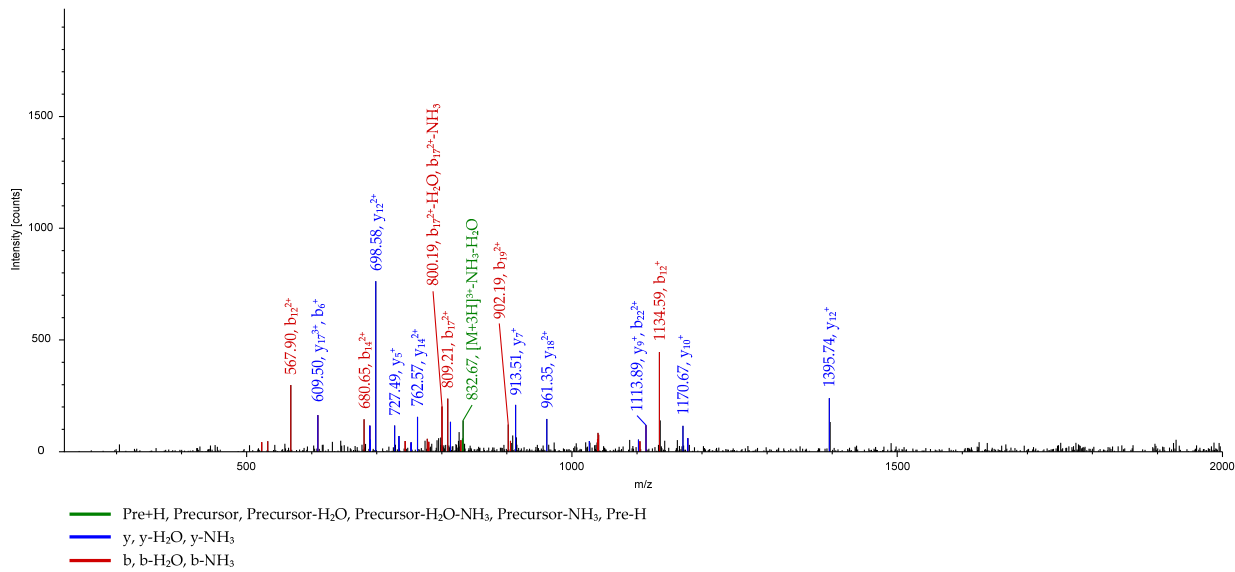


— Pre+H, Precursor, Precursor-H<sub>2</sub>O, Precursor-H<sub>2</sub>O-NH<sub>3</sub>, Precursor-NH<sub>3</sub>, Pre-H  
— b, b-H<sub>2</sub>O, b-NH<sub>3</sub>  
— y, y-H<sub>2</sub>O, y-NH<sub>3</sub>

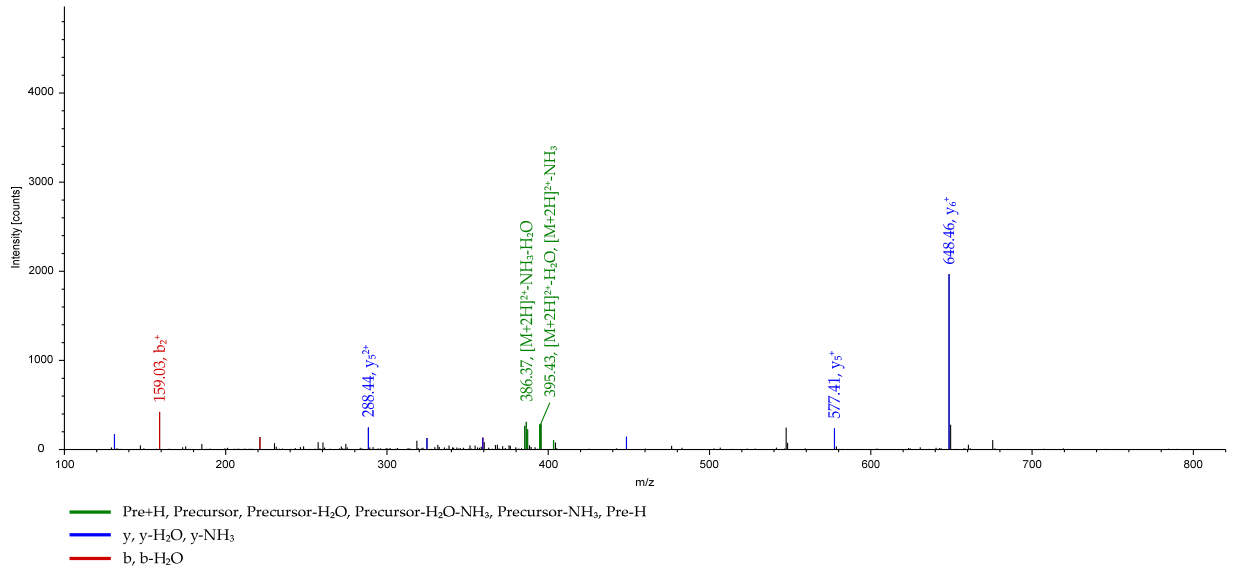
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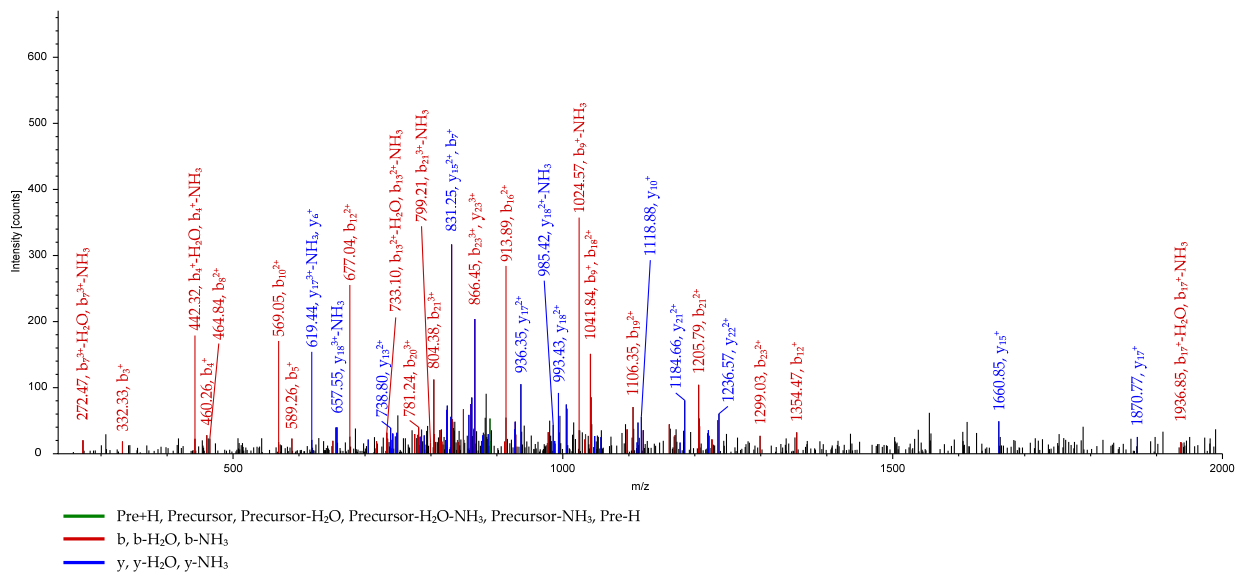
Extracted from: E:\onlinedigestion\USED FOR PAPER\rawcowllylsate\_1ng\_1.raw #383 RT: 3.54  
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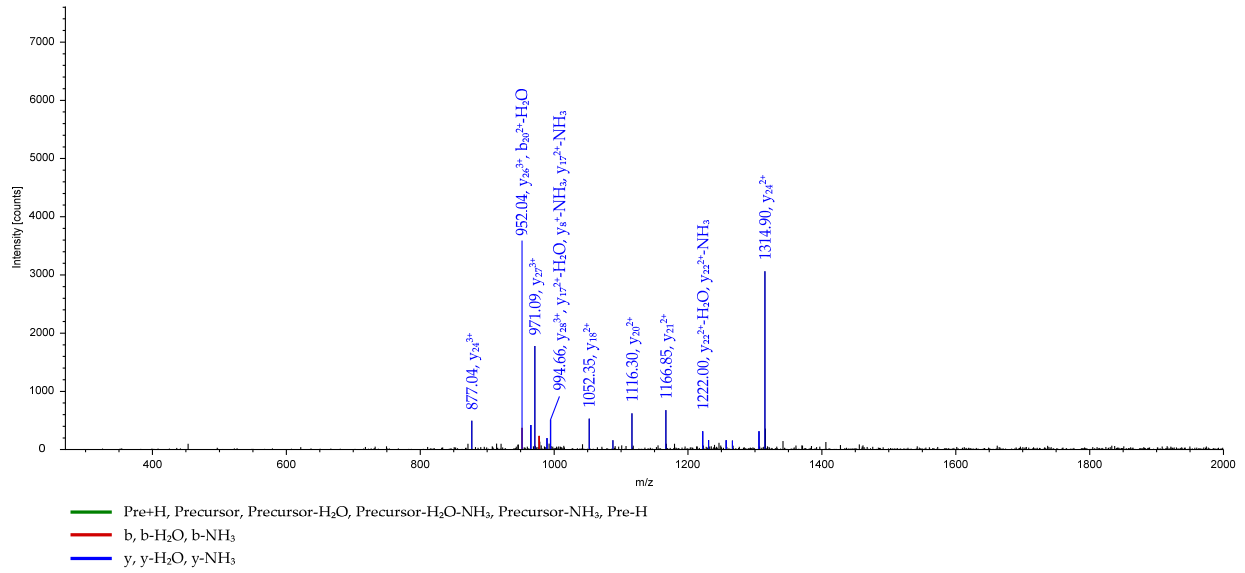
Extracted from: E:\onlinedigestion\USED FOR PAPER\rawcwllylate\_1ng\_1.raw #417 RT: 3.87  
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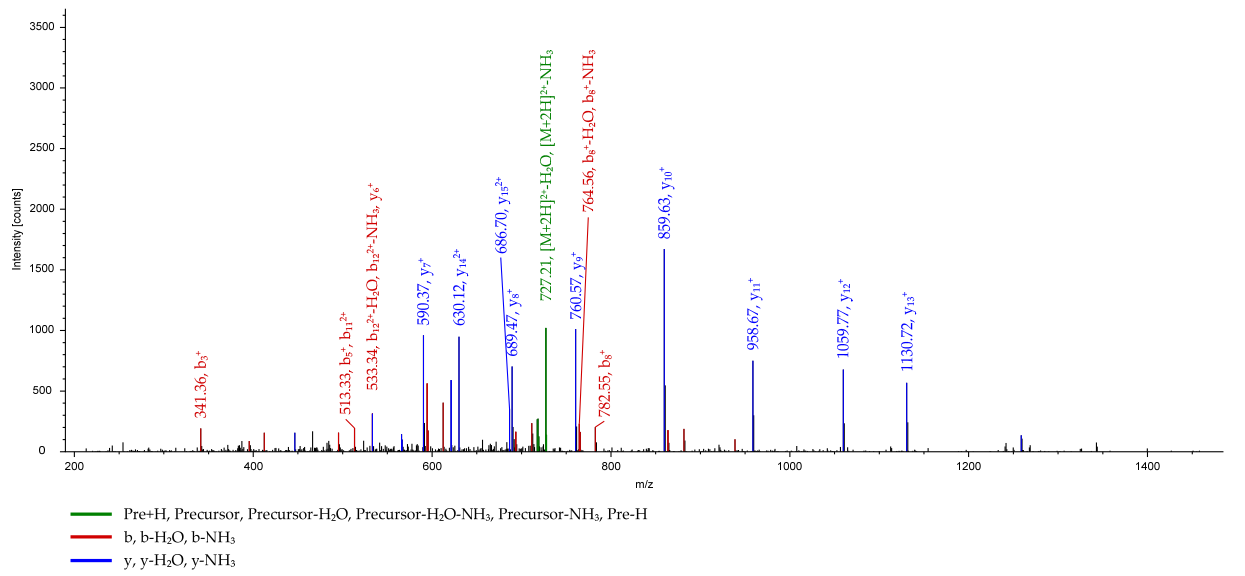
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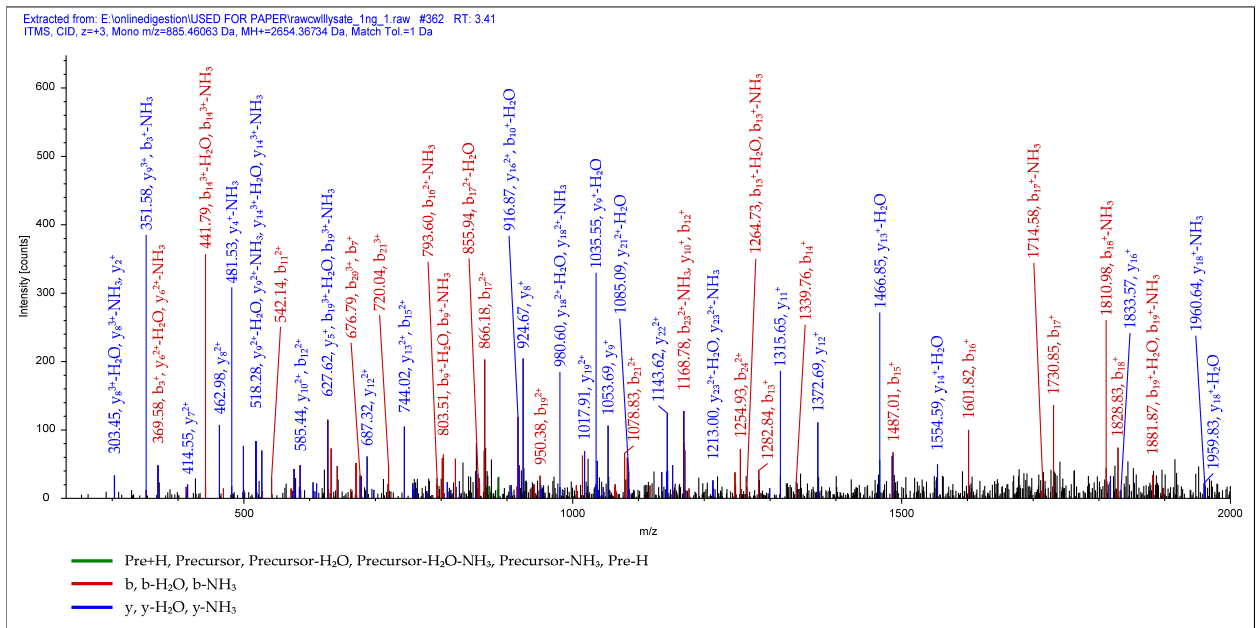


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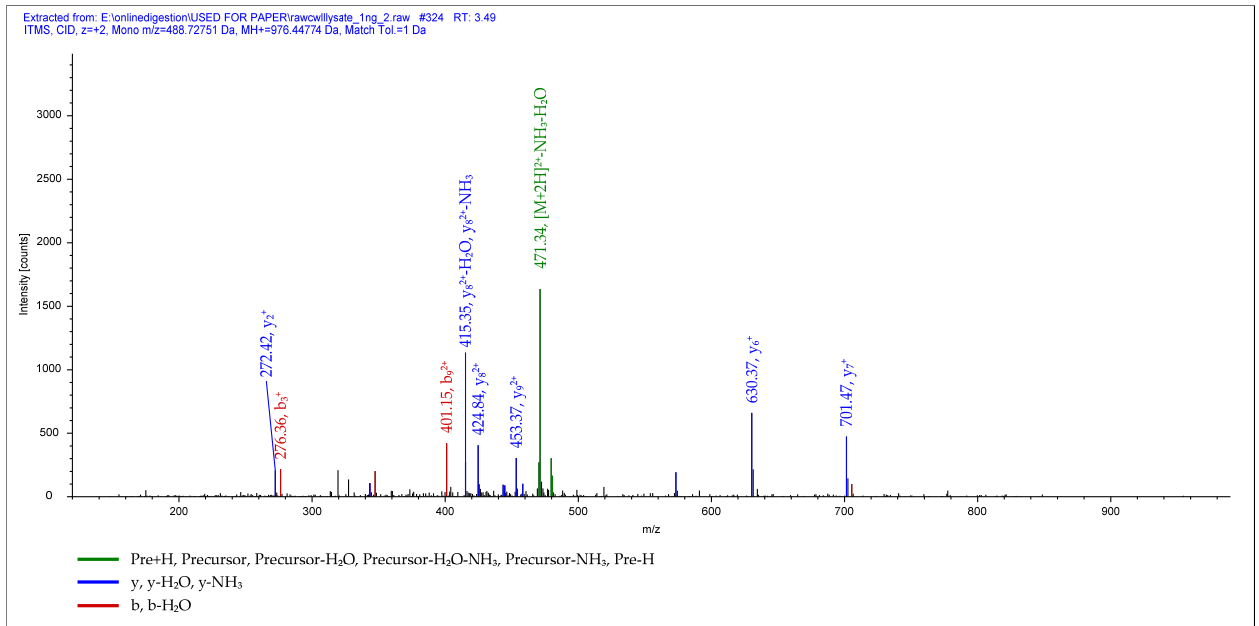


Extracted from: E:\onlinedigestion\USED FOR PAPER\rawcwllylate\_1ng\_1.raw #266 RT: 2.78  
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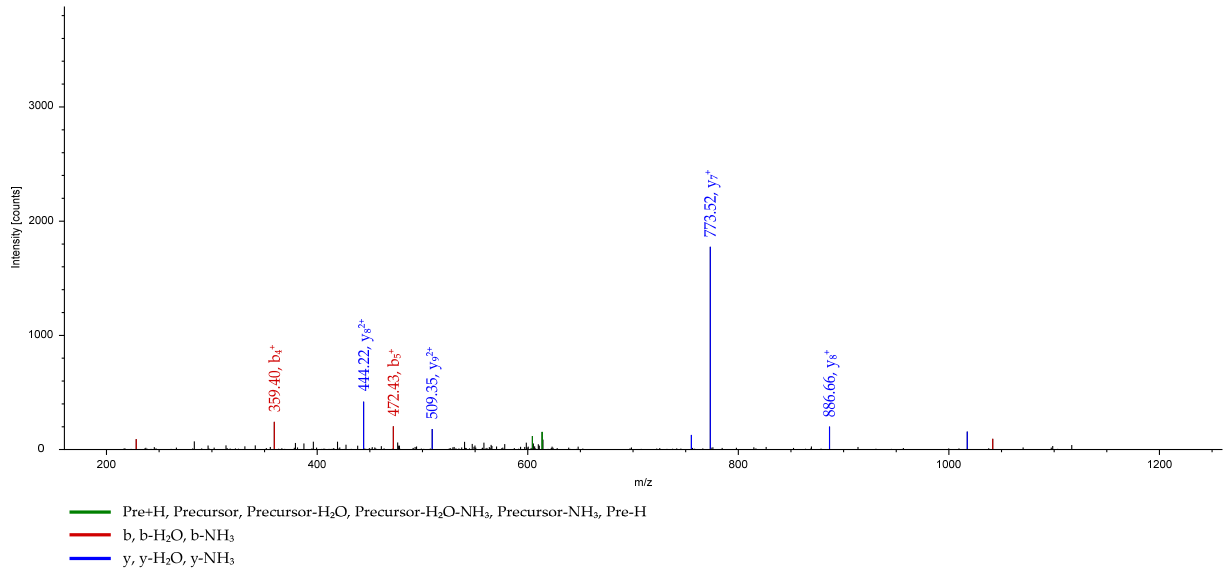




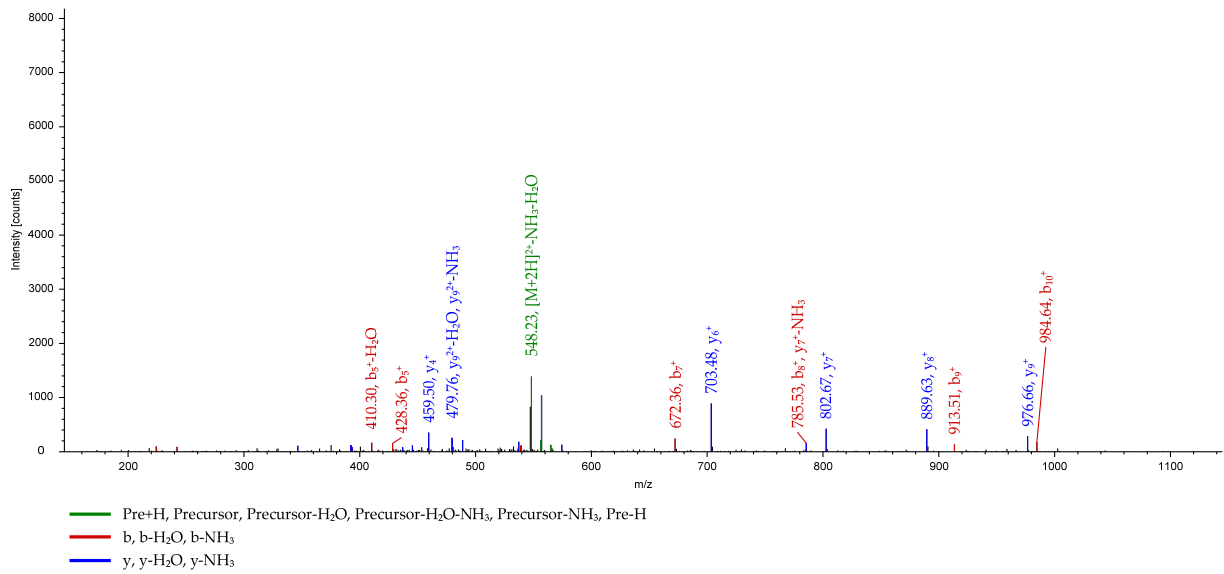
### 3 ng RAW 264.7 cell lysate\_2nd run



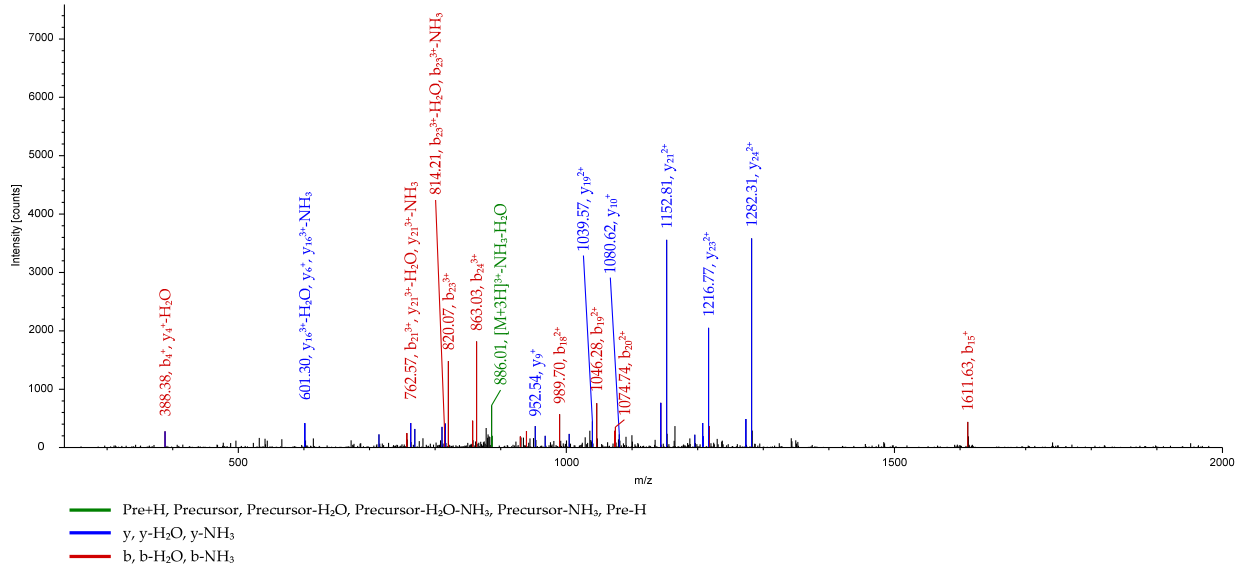
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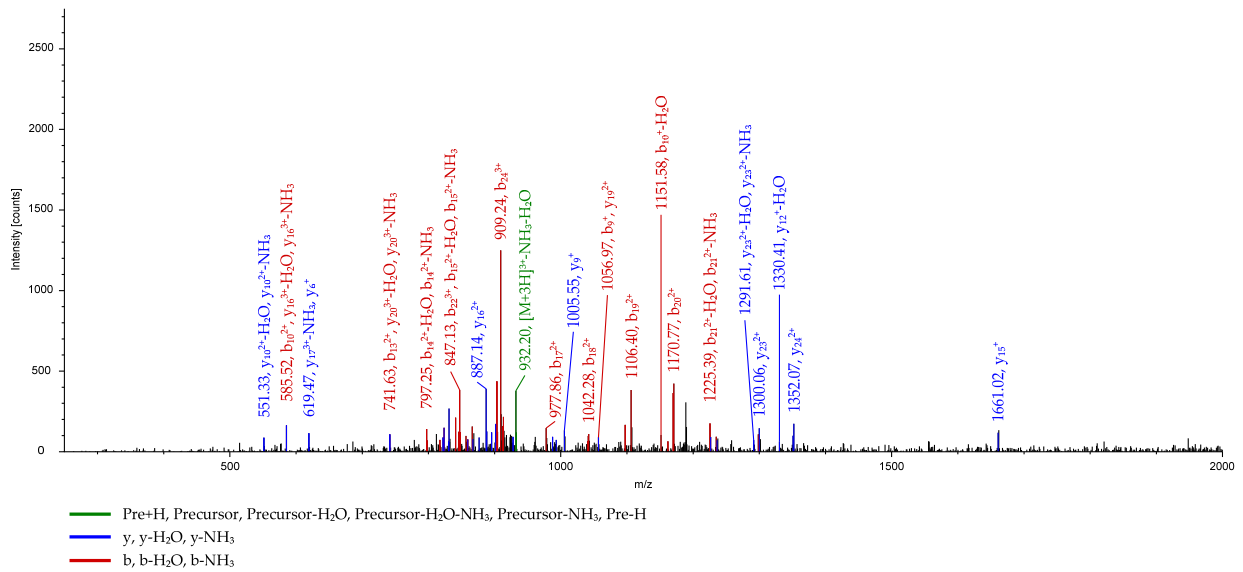
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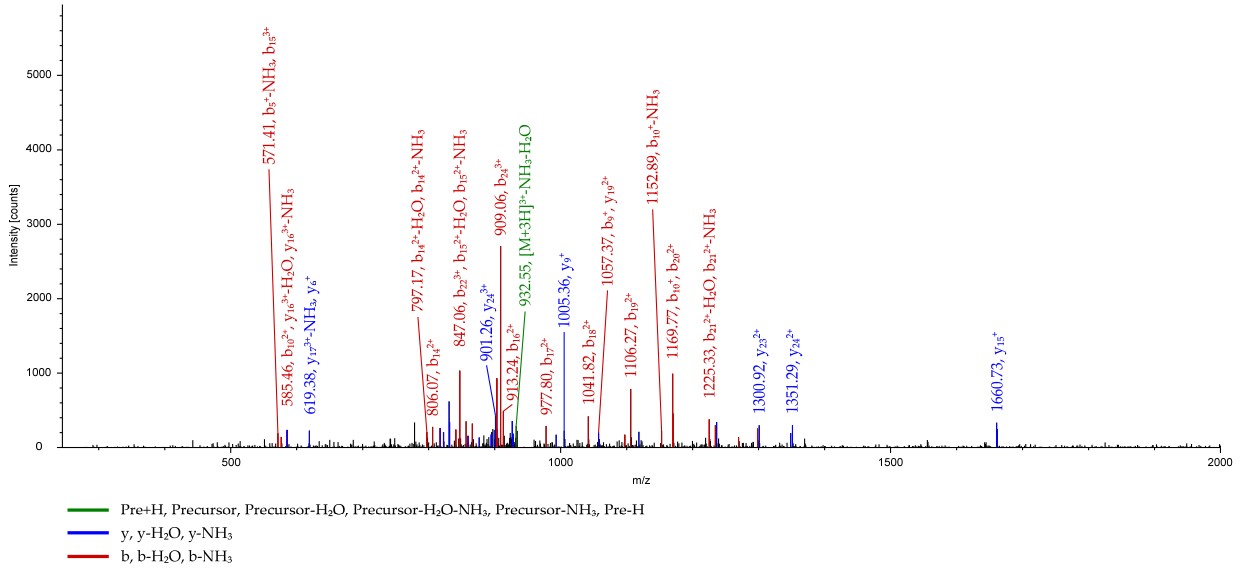
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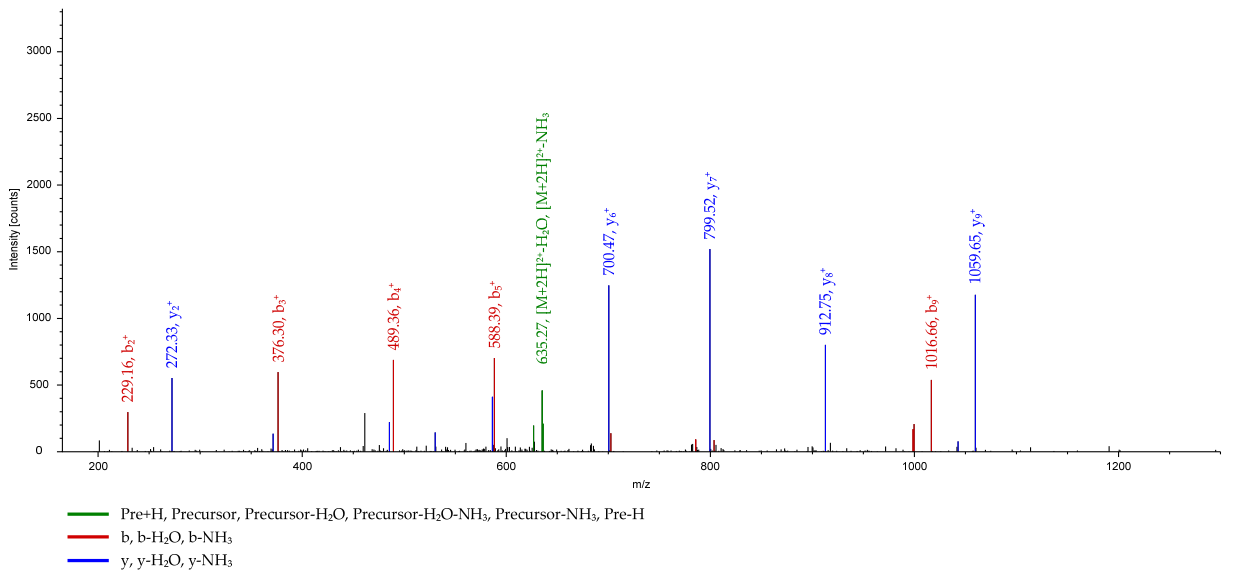
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ITMS, CID, z=+3, Mono m/z=943.80865 Da, MH+=2829.41141 Da, Match Tol.=1 Da

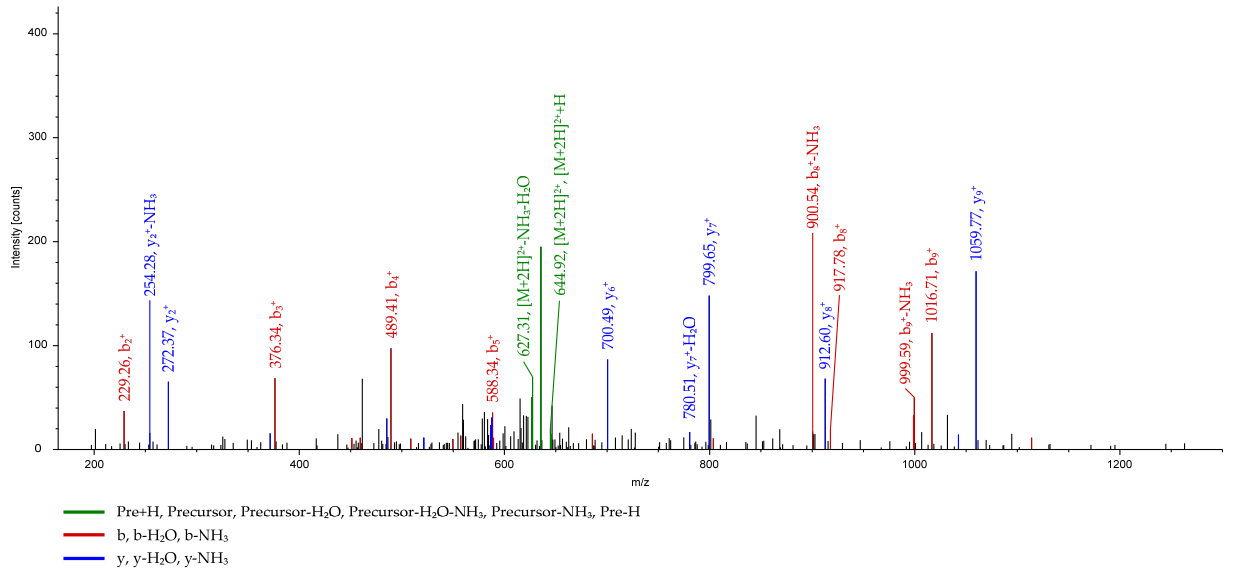


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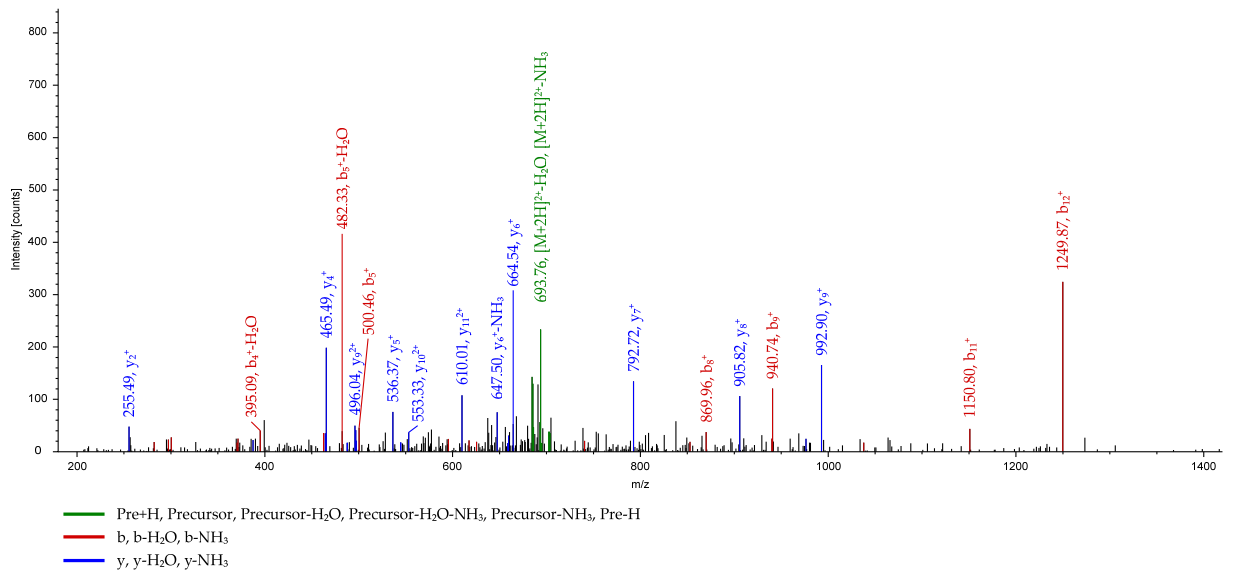




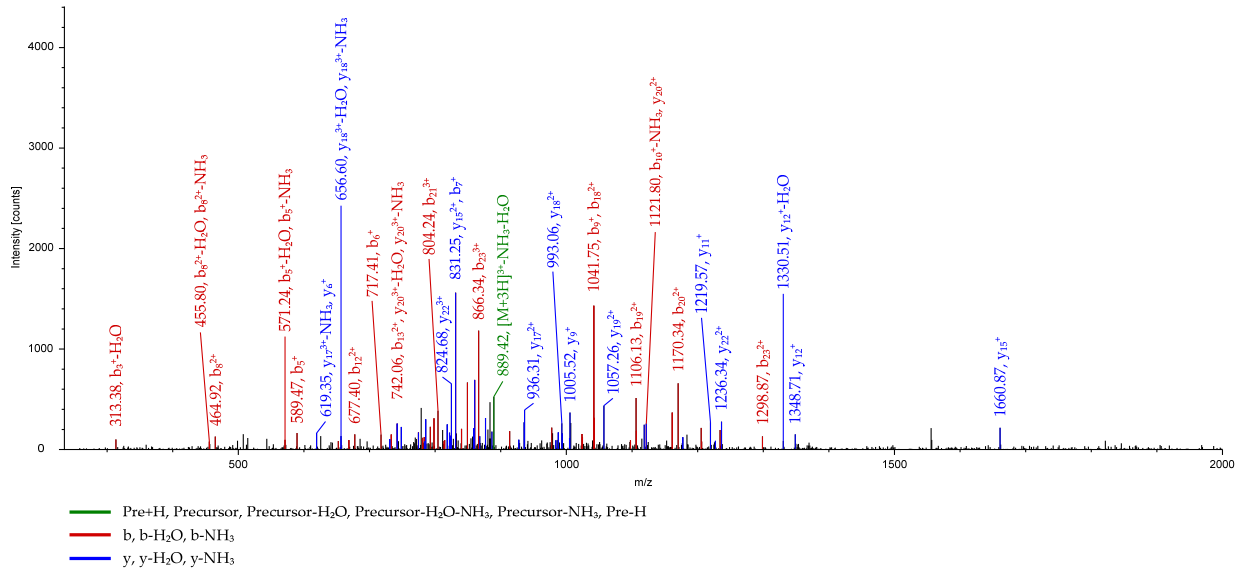
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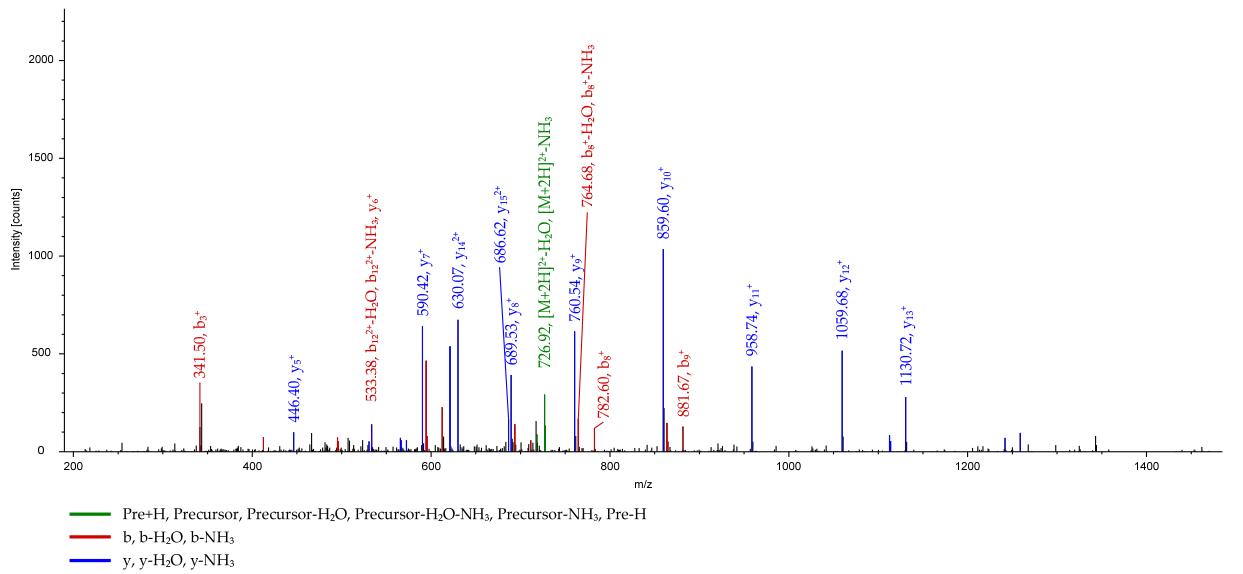
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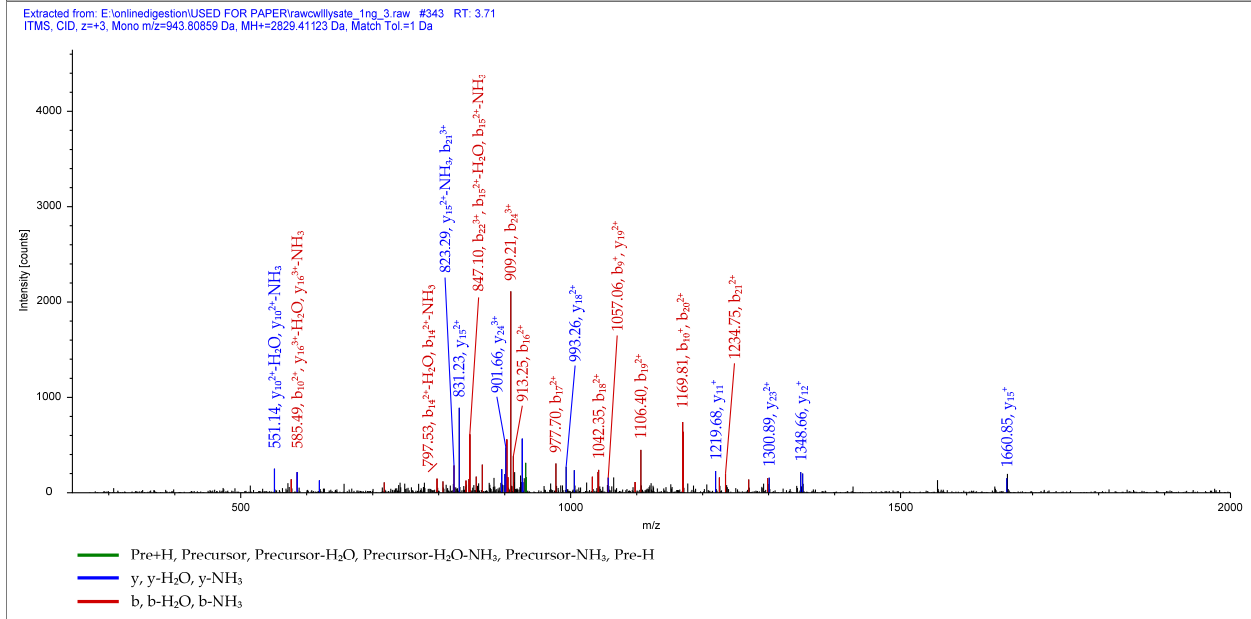
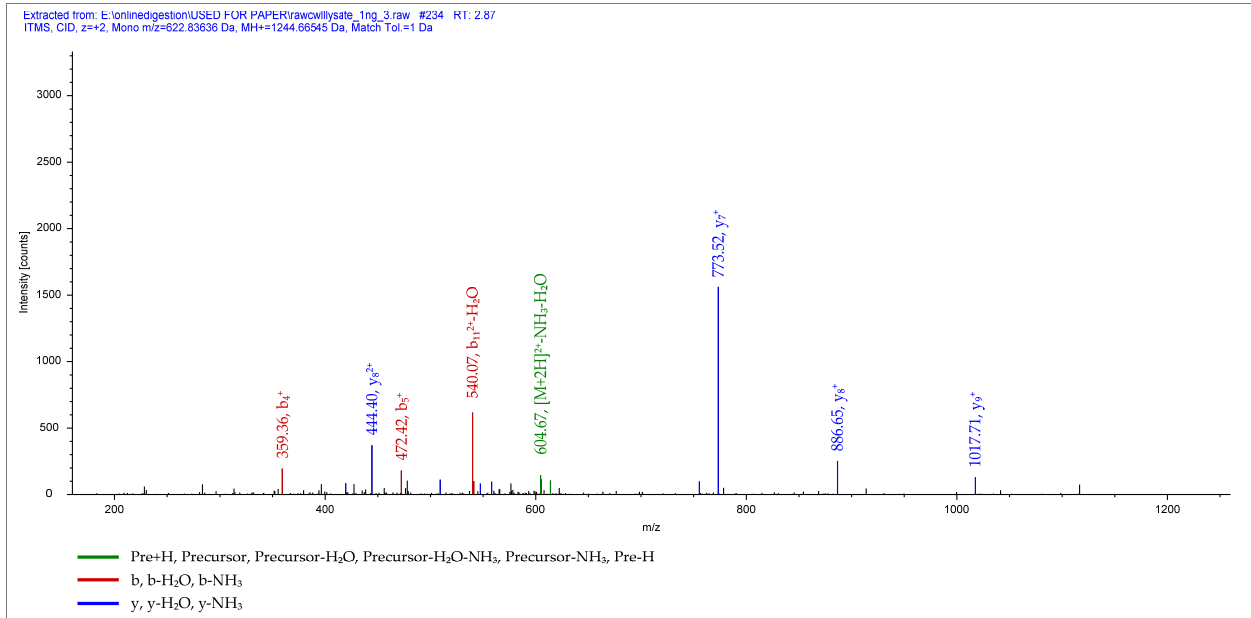
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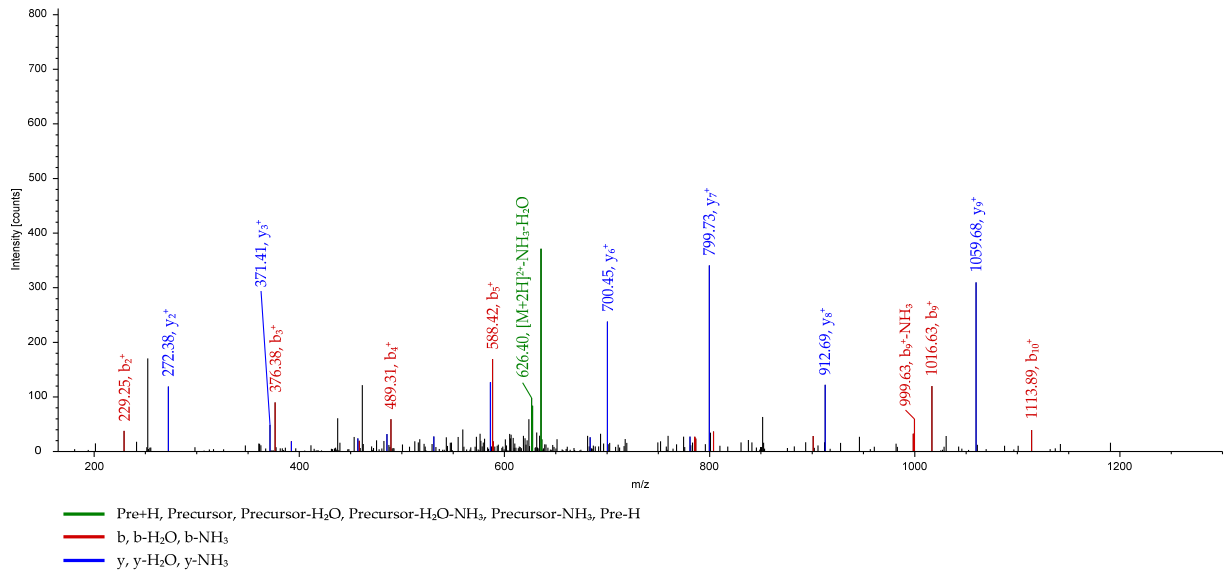
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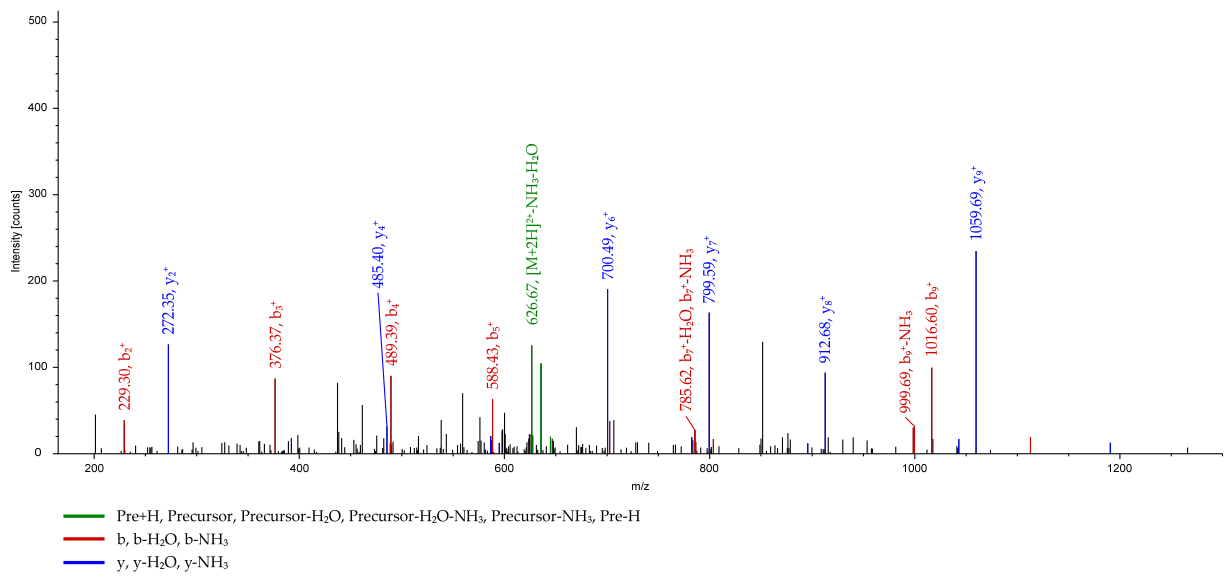
### 3 ng RAW 264.7 cell lysate\_3rd run



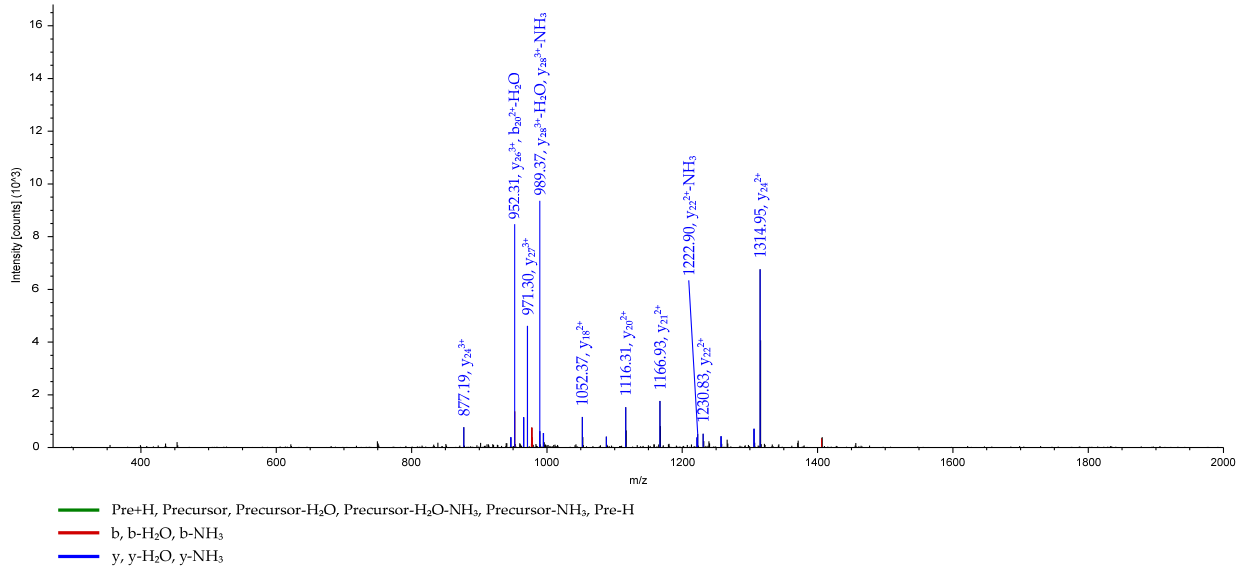
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Extracted from: E:\onlinedigestion\USED FOR PAPER\rawcwllylsate\_1ng\_3.raw #332 RT: 3.61  
ITMS, CID, z=+3, Mono m/z=1027.53162 Da, MH+=3080.58030 Da, Match Tol.=1 Da



Extracted from: E:\onlinedigestion\USED FOR PAPER\rawcwllylsate\_1ng\_3.raw #209 RT: 2.70  
ITMS, CID, z=+2, Mono m/z=735.93506 Da, MH+=1470.86284 Da, Match Tol.=1 Da

