

SUPPORTING INFORMATION

for

Structural characterization of glycerophospholipids by combinations of ozone- and collision-induced dissociation mass spectrometry: The next step towards “top-down” lipidomics

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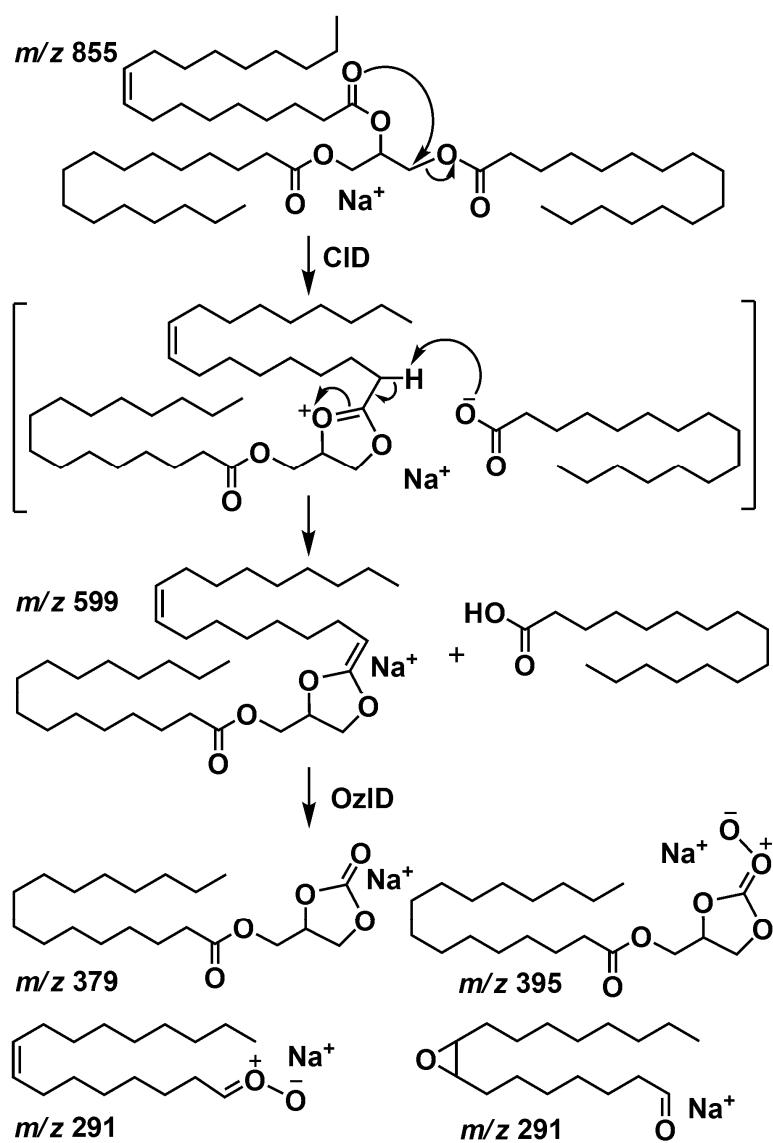
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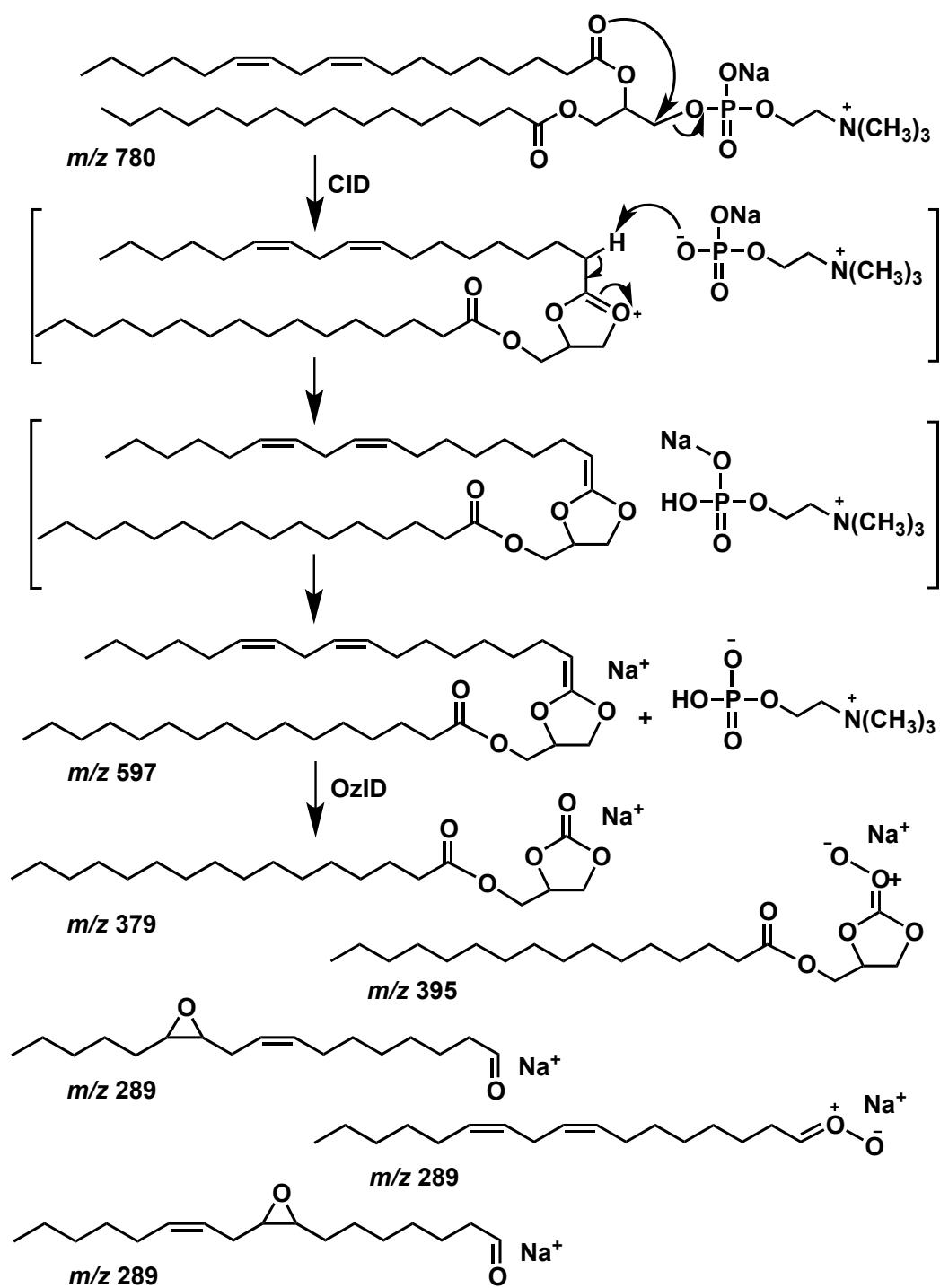
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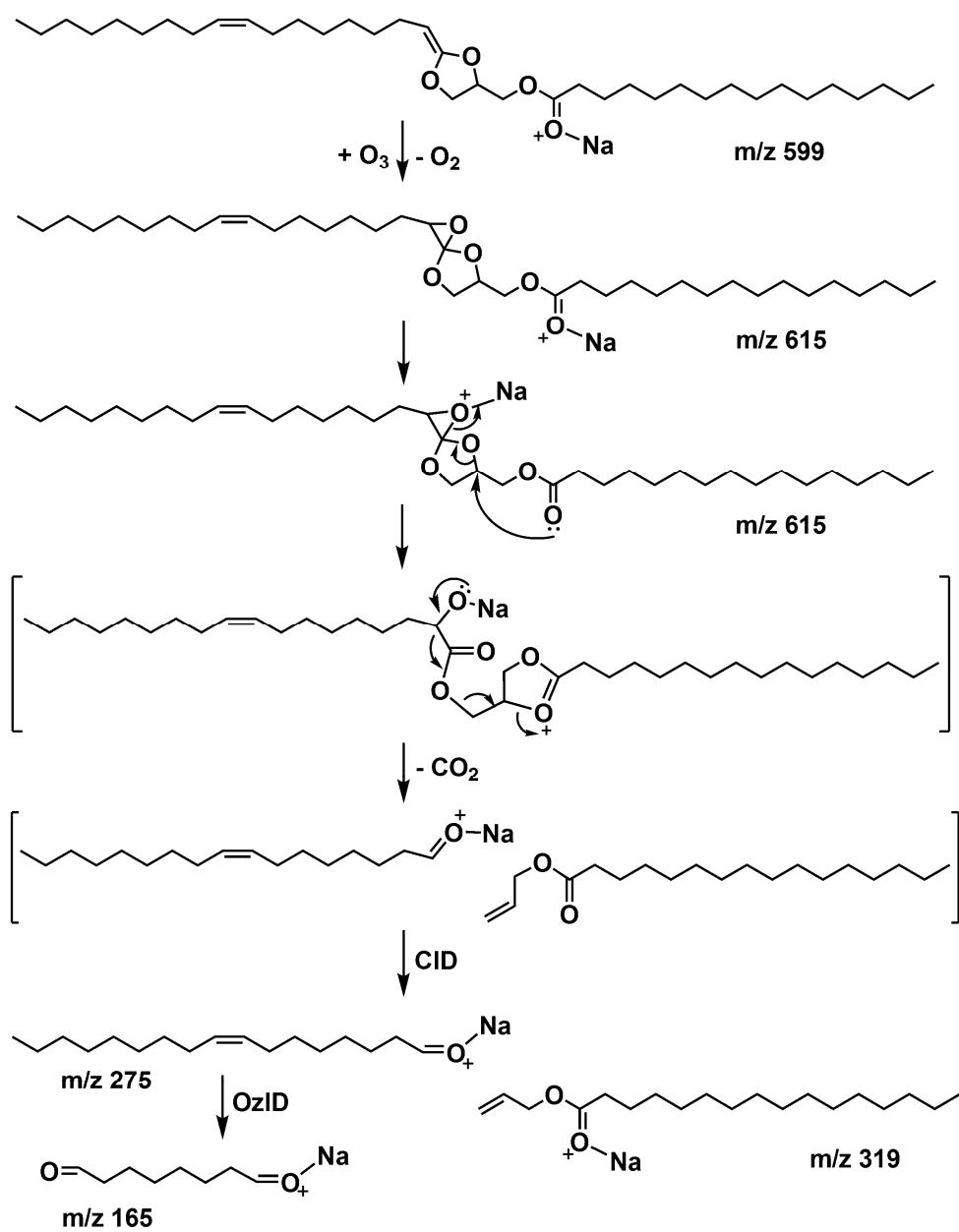
Scheme S-1: The substitution mechanism with a five-membered ring intermediate driven by the *sn*-2 fatty acyl chain during CID/OzID fragmentation of $[TG\ 16:0/18:1/16:0 + Na]^+$.



Scheme S-2: Proposed structures of major product ions forming in CID/OzID from $[PC\text{ }16:0/18:2 + \text{Na}]^+$.



Scheme S-3: A mechanism proposed for the formation and $(\text{CID/OzID})^2$ fragmentation of the m/z 614/615 ion resulting from the PC 16:0/18:1(9Z) standard.



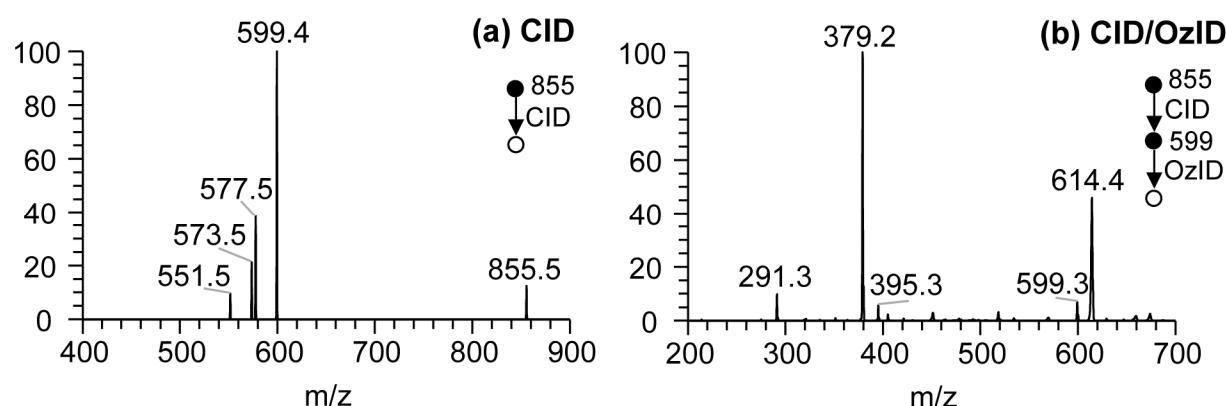


Figure S-1: (a) CID spectrum of $[TG\ 16:0/18:1/16:0 + Na]^+$ precursor ion at m/z 855, (b) CID/OzID spectrum of m/z 599 fragment ion $[TG\ 16:0/18:1/16:0 + Na - 16:0]^+$ formed in (a).

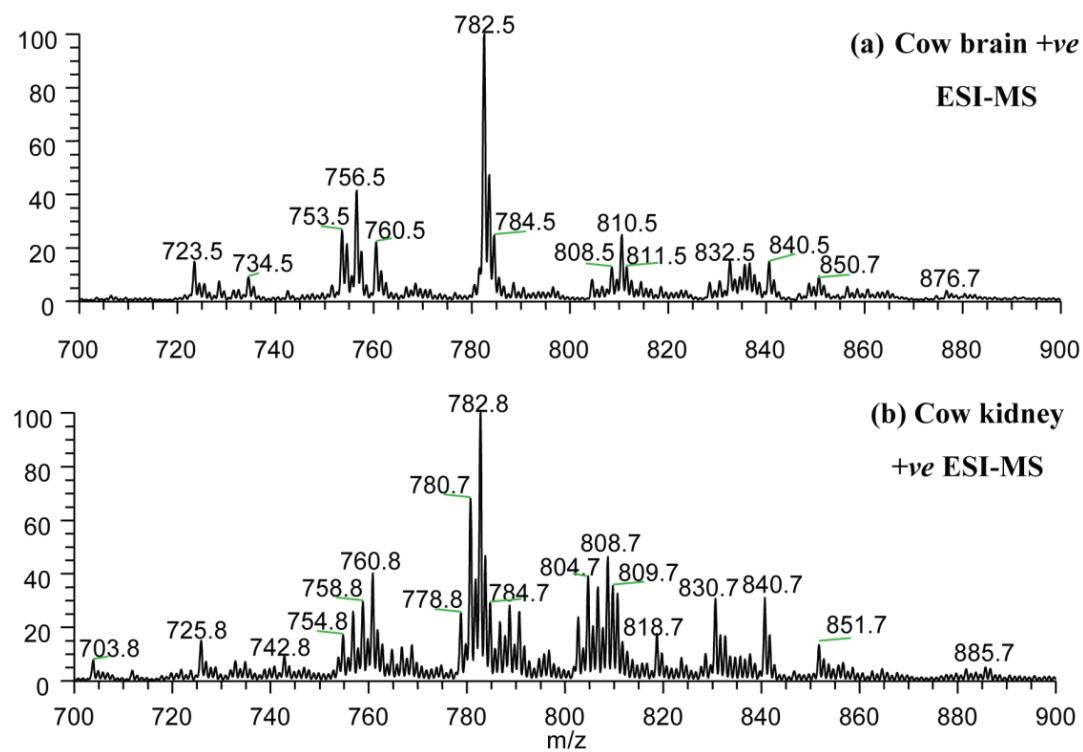


Figure S-2: ESI-MS mass spectra obtained in positive ion mode for lipids extracts from (a) cow brain and (b) cow kidney. The ions observed represent the $[M + Na]^+$ adducts of a wide range of phospholipids present in the samples.

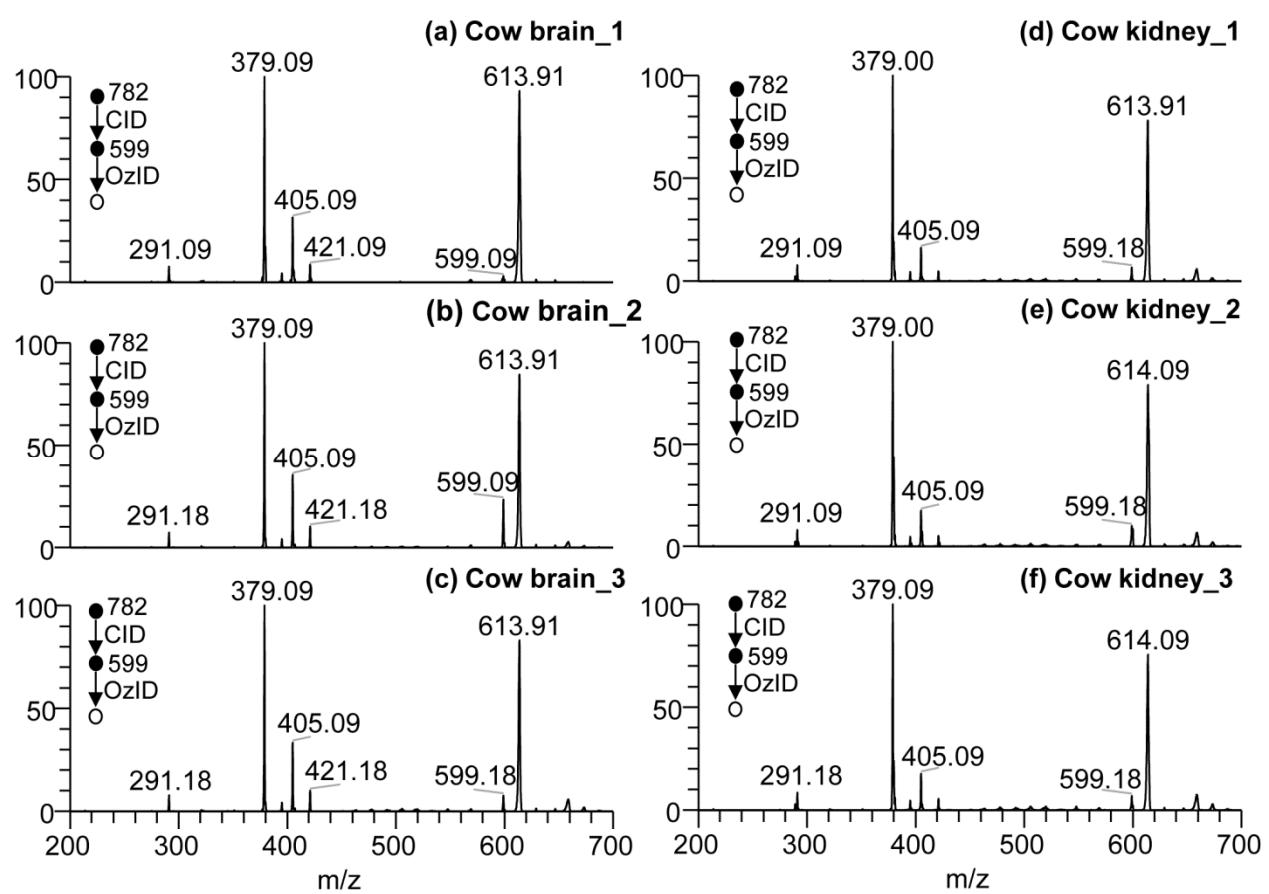


Figure S-3: Duplicate measurement of CID/OzID spectra generated by initially mass-selecting $[M + Na]^+$ adduct ions at m/z 782 from (a), (b) and (c) bovine brain and (d), (e), (f) bovine kidney extracts.

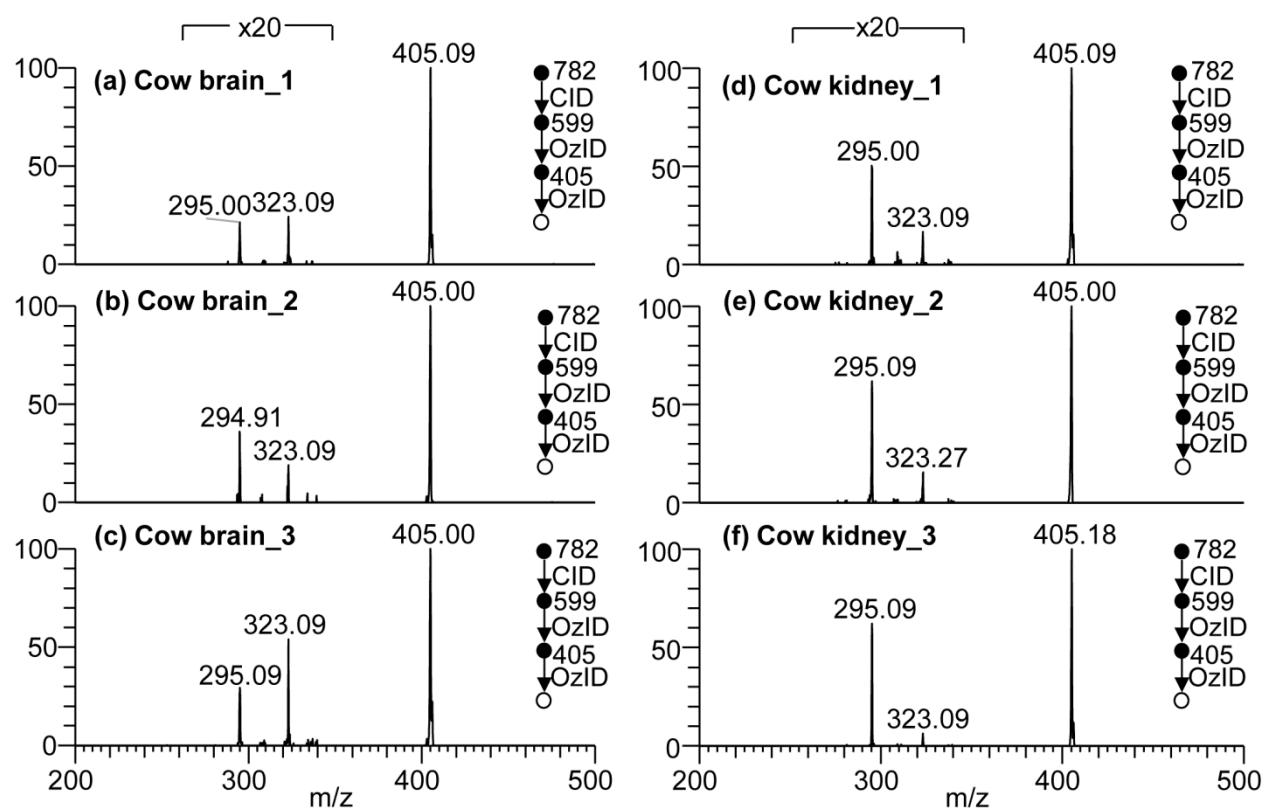


Figure S-4: Duplicate measurement of CID/OzID² spectra generated by initially mass-selecting $[M + Na]^{+}$ adduct ions at m/z 782 from (a), (b) and (c) bovine brain and (d), (e), (f) bovine kidney extracts. The magnifying factor of 20 times is the same for all spectra.

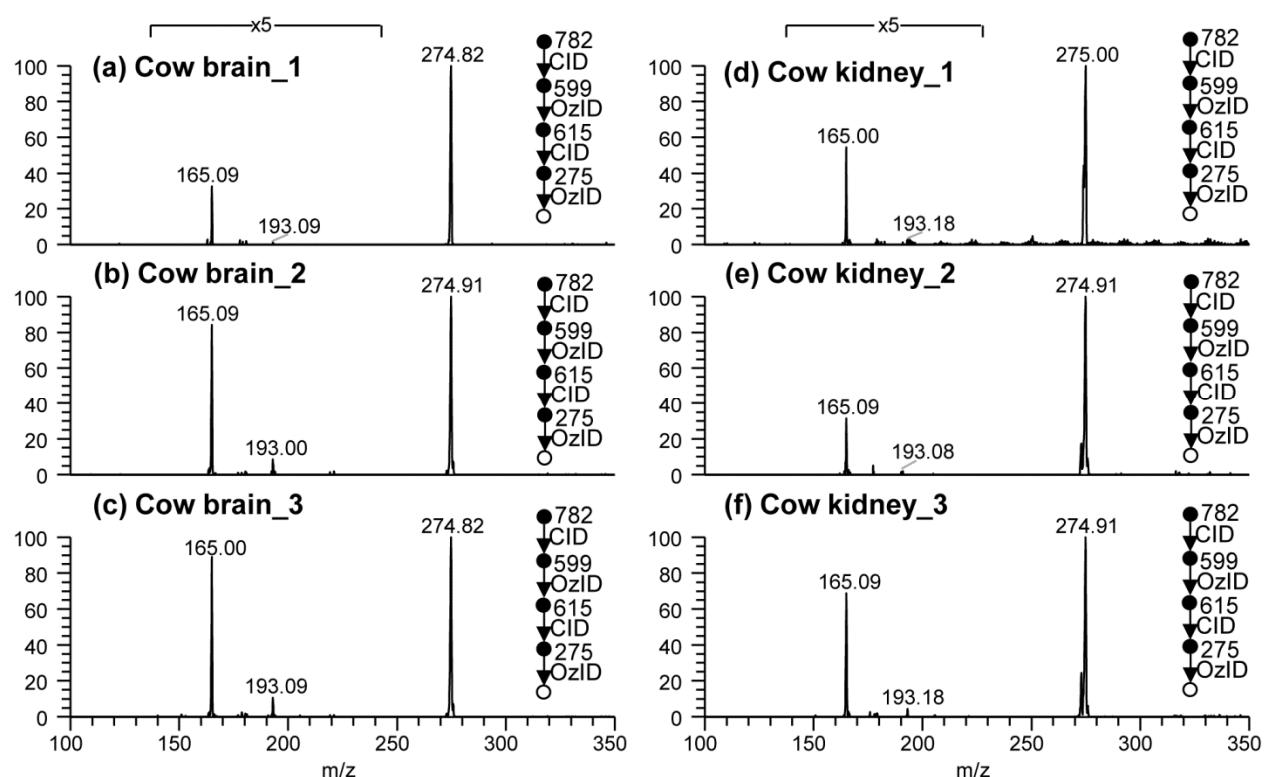


Figure S-5: Duplicate measurement of $(\text{CID}/\text{OzID})^2$ spectra generated by initially mass-selecting $[\text{M} + \text{Na}]^+$ adduct ions at m/z 782 from (a), (b) and (c) bovine brain and (d), (e), (f) bovine kidney extracts. The magnifying factor of 5 times is the same for all spectra.