

## Supporting Information

# Separation of isomeric tau phosphopeptides from Alzheimer's disease brain by cyclic ion mobility mass spectrometry

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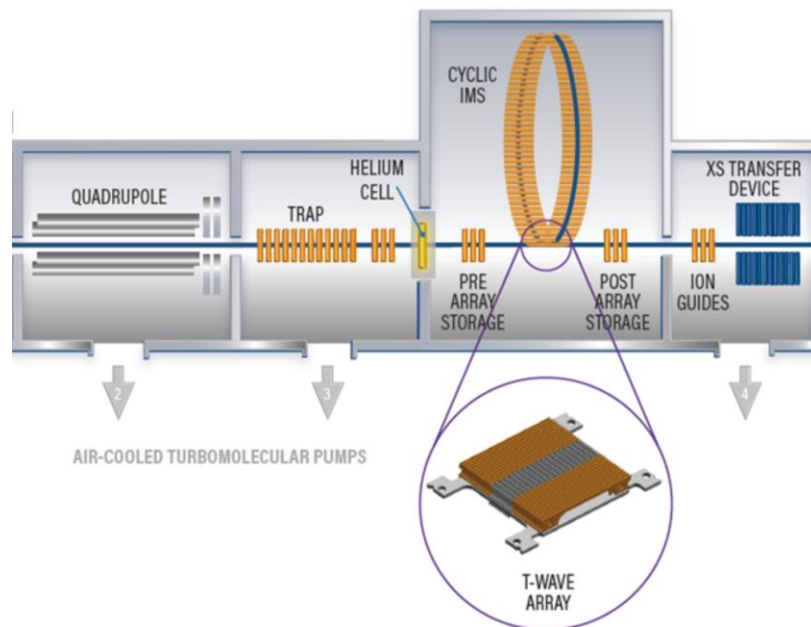
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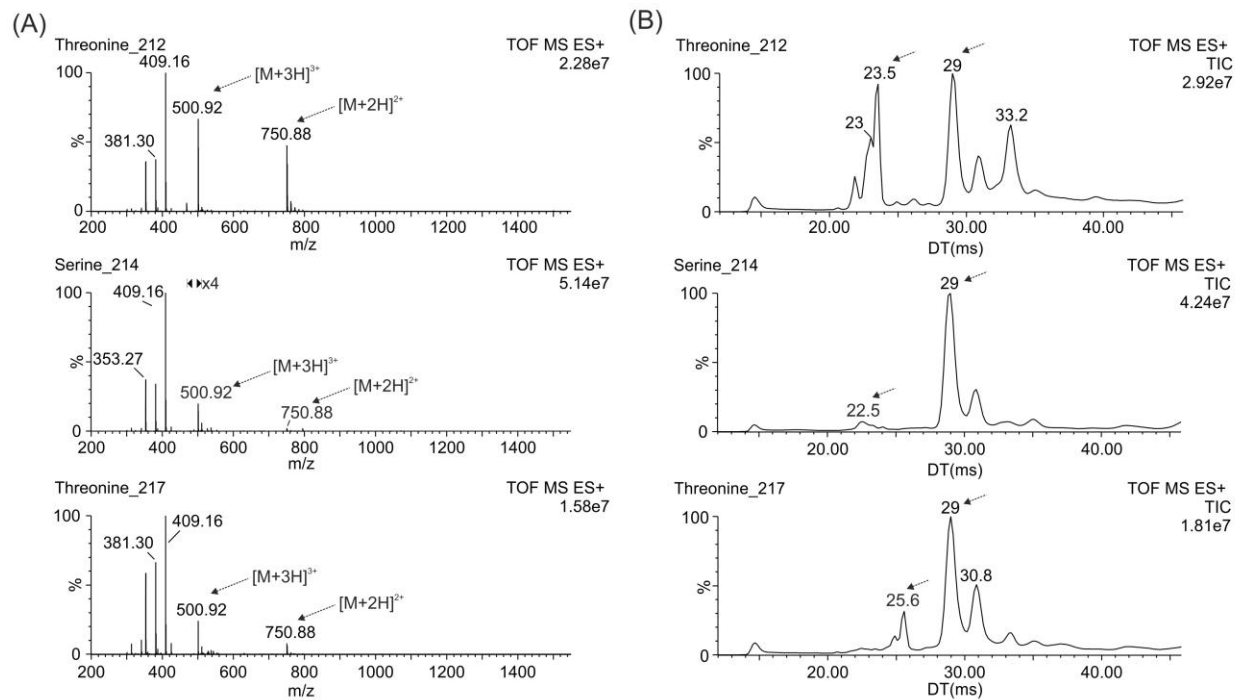
**Table S1.** Characteristic fragment ions for precursor  $[M+3H]^{3+}$



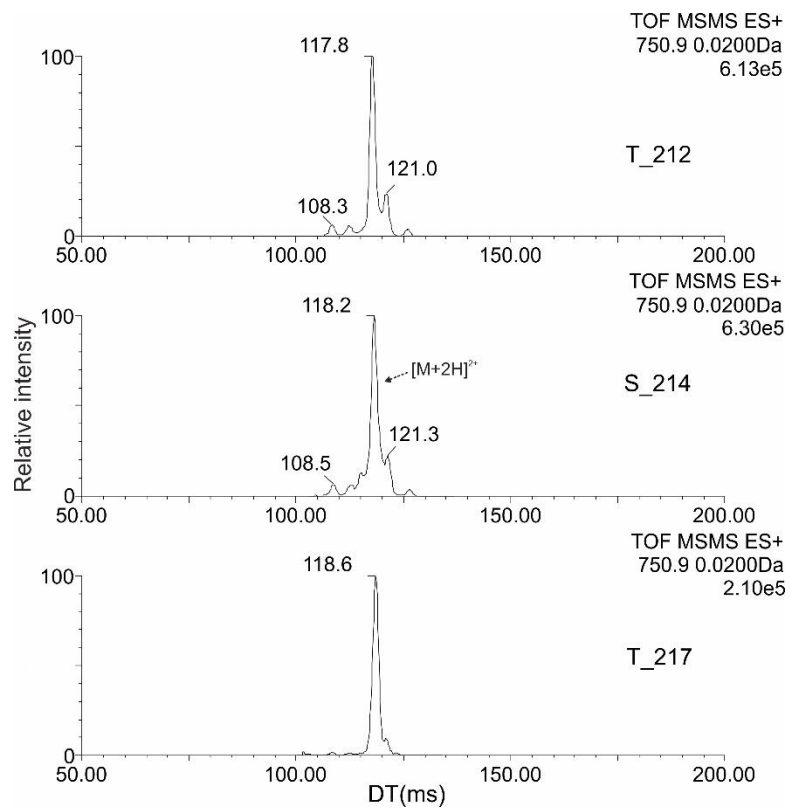
**Figure S1. Western blot analysis of brain extract.** The AD brains eluates after immuno-pulldown were analysed by Western blot using phosphorylation-dependent anti-tau antibodies against pThr212 and pThr217. Total tau was determined with a DC190 anti-tau antibody.



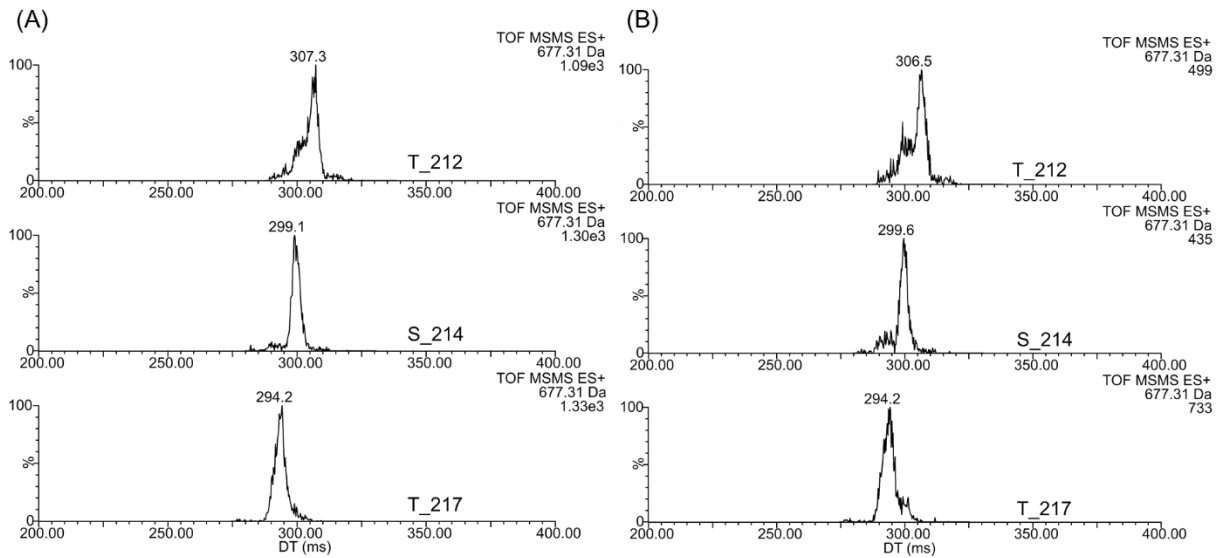
**Figure S2. Schematic of the Waters Select Series Cyclic IM Spectrometer.** After the initial trapping of ions in the first collision cell (trap), they are injected into the multifunctional array (T-wave array) where they are accelerated sidewise by a travelling wave into the cyclic separation IM cell.<sup>1</sup> Reprinted in part with permission of co-author Dale A. Cooper-Shepherd from "Application of cyclic ion mobility coupled to mass spectrometry for high peak capacity analysis of native and deuterated peptide mixtures", Martin Palmer; Malcolm Anderson; Dale A. Cooper-Shepherd; James I. Langridge; Robert Tonge; John R. Engen, poster nr. ThP285, Proceedings of 67th ASMS Conference on Mass Spectrometry and Allied Topics, Atlanta, Georgia, U.S.A., 2019. Copyright 2019 Waters Corporation.



**Figure S3. Separation of studied tau phosphopeptides by cIM.** The full scan MS spectra (A) and single pass total ion mobilograms (B) of infused standards of isomeric phosphopeptides (T212, S214 and T217). Arrows mark mobility peaks related to  $[M+3H]^{3+}$  (around 23 ms) and  $[M+2H]^{2+}$  (cca 29 ms).



**Figure S4. Ion mobilograms of  $[M+2H]^{2+}$  after 5 pass separation.** Extracted ion mobilograms of precursor ion at m/z 750.9.



**Figure S5. Ten pass ion mobilograms of phosphopeptides at low concentrations.** Extracted ion mobilograms of N terminal b6 fragment ion at m/z 677 generated by CID in trap (24 V) from  $[M+3H]^{3+}$  for T212, S214, T217, the concentration of standards: 1 nmol/L (A); 0.1 nmol/L (B); without smoothing, drift time aligned to Figure 3 by subtracting 4 ms.

**Table S1. Characteristic fragment ions for precursor  $[M+3H]^{3+}$ .** A unique fragment ion is in bold; for common fragment ions differing significantly in signal intensity, s stands for stronger, w for weaker.

phosphopeptides	fragment ions (m/z)
<b>T212</b>	279.1 (s); 286.1 (w); 1116.6 (s); 1135.7 (s); 1222.7 (s)
<b>S214</b>	
<b>T217</b>	279.1 (w); 286.1 (s); <b>1102.5</b> ; 1116.6 (w); 1135.7 (w); 1222.7 (w)

## Reference

(1) Giles, K.; Ujma, J.; Wildgoose, J.; Pringle, S.; Richardson, K.; Langridge, D.; Green, M. A Cyclic Ion Mobility-Mass Spectrometry System. *Anal. Chem.* **2019**, *91* (13), 8564-8573.