

Correction to Hydrophobic Contributions to the Membrane Docking of Synaptotagmin 7 C2A Domain: Mechanistic Contrast between Isoforms 1 and 7

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Page 7660. The final sentence of the first Discussion paragraph refers to the incorrect protein isoform. The corrected sentence follows.

(vi) The Syt7 C2A dissociation kinetics are further slowed by kosmotropic solutes, providing evidence that the rate-limiting step in dissociation may involve exposure of hydrophobic residues to solvent.

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