Ucheor B Choi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Architecture of the synaptotagmin–SNARE machinery for neuronal exocytosis. Nature, 2015, 525, 62-67.	27.8	268
2	Single-molecule FRET–derived model of the synaptotagmin 1–SNARE fusion complex. Nature Structural and Molecular Biology, 2010, 17, 318-324.	8.2	194
3	Molecular Mechanisms of Synaptic Vesicle Priming by Munc13 and Munc18. Neuron, 2017, 95, 591-607.e10.	8.1	185
4	Accessory Proteins Stabilize the Acceptor Complex for Synaptobrevin, the 1:1 Syntaxin/SNAP-25 Complex. Structure, 2008, 16, 308-320.	3.3	151
5	Molecular Mechanisms of Fast Neurotransmitter Release. Annual Review of Biophysics, 2018, 47, 469-497.	10.0	133
6	Beyond the Random Coil: Stochastic Conformational Switching in Intrinsically Disordered Proteins. Structure, 2011, 19, 566-576.	3.3	109
7	Optimizing Methods to Recover Absolute FRET Efficiency from Immobilized Single Molecules. Biophysical Journal, 2010, 99, 961-970.	0.5	93
8	Conformational change of syntaxin linker region induced by Munc13s initiates <scp>SNARE</scp> complex formation in synaptic exocytosis. EMBO Journal, 2017, 36, 816-829.	7.8	78
9	The pre-synaptic fusion machinery. Current Opinion in Structural Biology, 2019, 54, 179-188.	5.7	72
10	Structural principles of SNARE complex recognition by the AAA+ protein NSF. ELife, 2018, 7, .	6.0	67
11	C-terminal domain of mammalian complexin-1 localizes to highly curved membranes. Proceedings of the United States of America, 2016, 113, E7590-E7599.	7.1	66
12	Ca2+-Triggered Synaptic Vesicle Fusion Initiated by Release of Inhibition. Trends in Cell Biology, 2018, 28, 631-645.	7.9	46
13	Effect of Src Kinase Phosphorylation on Disordered C-terminal Domain of N-Methyl-d-aspartic Acid (NMDA) Receptor Subunit GluN2B Protein. Journal of Biological Chemistry, 2011, 286, 29904-29912.	3.4	44
14	N-terminal domain of complexin independently activates calcium-triggered fusion. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E4698-E4707.	7.1	44
15	Complexin induces a conformational change at the membrane-proximal C-terminal end of the SNARE complex. ELife, 2016, 5, .	6.0	36
16	NSF-mediated disassembly of on- and off-pathway SNARE complexes and inhibition by complexin. ELife, 2018, 7, .	6.0	34
17	Modulating the Intrinsic Disorder in the Cytoplasmic Domain Alters the Biological Activity of the N-Methyl-d-aspartate-sensitive Glutamate Receptor. Journal of Biological Chemistry, 2013, 288, 22506-22515.	3.4	33
18	Immobilization of Proteins for Single-Molecule Fluorescence Resonance Energy Transfer		26

Measurements of Conformation and Dynamics. , 2012, 896, 3-20.

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19	Conformational dynamics of auto-inhibition in the ER calcium sensor STIM1. ELife, 2021, 10, .	6.0	22
20	Structures of neurexophilin–neurexin complexes reveal a regulatory mechanism of alternative splicing. EMBO Journal, 2019, 38, e101603.	7.8	19
21	Munc18a Does Not Alter Fusion Rates Mediated by Neuronal SNAREs, Synaptotagmin, and Complexin. Journal of Biological Chemistry, 2015, 290, 10518-10534.	3.4	17
22	Reconstitution of Multivalent PDZ Domain Binding to the Scaffold Protein PSD-95 Reveals Ternary-Complex Specificity of Combinatorial Inhibition. Structure, 2014, 22, 1458-1466.	3.3	15
23	Conformational change of Syntaxin-3b in regulating SNARE complex assembly in the ribbon synapses. Scientific Reports, 2022, 12, .	3.3	2
24	Conformational Dynamics of SNARE Proteins during NSFâ€Mediated Disassembly. FASEB Journal, 2021, 35,	0.5	0