

Shyam S Krishnakumar

List of Publications by Year in descending order

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Version: 2024-02-01

41
papers

2,059
citations

218677

26
h-index

265206

42
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56
all docs

56
docs citations

56
times ranked

2227
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular determinants of complexin clamping and activation function. <i>ELife</i> , 2022, 11, .	6.0	16
2	Munc13 binds and recruits SNAP25 to chaperone SNARE complex assembly. <i>FEBS Letters</i> , 2021, 595, 297-309.	2.8	33
3	Vesicle capture by membrane-bound Munc13 requires self-assembly into discrete clusters. <i>FEBS Letters</i> , 2021, 595, 2185-2196.	2.8	15
4	Symmetrical arrangement of proteins under release-ready vesicles in presynaptic terminals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	40
5	Synaptotagmin-1 membrane binding is driven by the C2B domain and assisted cooperatively by the C2A domain. <i>Scientific Reports</i> , 2020, 10, 18011.	3.3	22
6	Synaptotagmin 1 oligomers clamp and regulate different modes of neurotransmitter release. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 3819-3827.	7.1	47
7	Synergistic roles of Synaptotagmin-1 and complexin in calcium-regulated neuronal exocytosis. <i>ELife</i> , 2020, 9, .	6.0	40
8	Structural basis for the clamping and Ca ²⁺ activation of SNARE-mediated fusion by synaptotagmin. <i>Nature Communications</i> , 2019, 10, 2413.	12.8	39
9	Mechanisms of Neurological Dysfunction in GOSR2 Progressive Myoclonus Epilepsy, a Golgi SNAREopathy. <i>Neuroscience</i> , 2019, 420, 41-49.	2.3	10
10	Mutations in the Neuronal Vesicular SNARE VAMP2 Affect Synaptic Membrane Fusion and Impair Human Neurodevelopment. <i>American Journal of Human Genetics</i> , 2019, 104, 721-730.	6.2	88
11	Synaptotagmin oligomers are necessary and can be sufficient to form a Ca ²⁺ -sensitive fusion clamp. <i>FEBS Letters</i> , 2019, 593, 154-162.	2.8	42
12	Symmetrical organization of proteins under docked synaptic vesicles. <i>FEBS Letters</i> , 2019, 593, 144-153.	2.8	34
13	Using Nanodiscs to Probe Ca ²⁺ -Dependent Membrane Interaction of Synaptotagmin-1. <i>Methods in Molecular Biology</i> , 2019, 1860, 221-236.	0.9	1
14	Rearrangements under confinement lead to increased binding energy of Synaptotagmin with anionic membranes in Mg ²⁺ and Ca ²⁺ . <i>FEBS Letters</i> , 2018, 592, 1497-1506.	2.8	13
15	PRRT2 Regulates Synaptic Fusion by Directly Modulating SNARE Complex Assembly. <i>Cell Reports</i> , 2018, 22, 820-831.	6.4	67
16	Two Disease-Causing SNAP-25B Mutations Selectively Impair SNARE C-terminal Assembly. <i>Journal of Molecular Biology</i> , 2018, 430, 479-490.	4.2	21
17	Synaptotagmin oligomerization is essential for calcium control of regulated exocytosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E7624-E7631.	7.1	51
18	Synergistic control of neurotransmitter release by different members of the synaptotagmin family. <i>Current Opinion in Neurobiology</i> , 2018, 51, 154-162.	4.2	34

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19	Homozygous mutations in <i>VAMP1</i> cause a presynaptic congenital myasthenic syndrome. <i>Annals of Neurology</i> , 2017, 81, 597-603.	5.3	48
20	Kv1.1 channelopathy abolishes presynaptic spike width modulation by subthreshold somatic depolarization. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 2395-2400.	7.1	31
21	Mutations in <i>Membrin/ GOSR2</i> Reveal Stringent Secretory Pathway Demands of Dendritic Growth and Synaptic Integrity. <i>Cell Reports</i> , 2017, 21, 97-109.	6.4	29
22	Hypothesis "buttressed rings assemble, clamp, and release SNAREpins for synaptic transmission. <i>FEBS Letters</i> , 2017, 591, 3459-3480.	2.8	76
23	Otoferlin acts as a Ca ²⁺ sensor for vesicle fusion and vesicle pool replenishment at auditory hair cell ribbon synapses. <i>ELife</i> , 2017, 6, .	6.0	108
24	Circular oligomerization is an intrinsic property of synaptotagmin. <i>ELife</i> , 2017, 6, .	6.0	47
25	Dilation of fusion pores by crowding of SNARE proteins. <i>ELife</i> , 2017, 6, .	6.0	57
26	Nanodisc-cell fusion: control of fusion pore nucleation and lifetimes by SNARE protein transmembrane domains. <i>Scientific Reports</i> , 2016, 6, 27287.	3.3	39
27	Using ApoE Nanolipoprotein Particles To Analyze SNARE-Induced Fusion Pores. <i>Langmuir</i> , 2016, 32, 3015-3023.	3.5	22
28	Ring-like oligomers of Synaptotagmins and related C2 domain proteins. <i>ELife</i> , 2016, 5, .	6.0	57
29	Re-visiting the trans insertion model for complexin clamping. <i>ELife</i> , 2015, 4, .	6.0	33
30	Calcium sensitive ring-like oligomers formed by synaptotagmin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 13966-13971.	7.1	76
31	Conformational Dynamics of Calcium-Triggered Activation of Fusion by Synaptotagmin. <i>Biophysical Journal</i> , 2013, 105, 2507-2516.	0.5	39
32	A conformational switch in complexin is required for synaptotagmin to trigger synaptic fusion. <i>Nature Structural and Molecular Biology</i> , 2011, 18, 934-940.	8.2	85
33	Complexin cross-links prefusion SNAREs into a zigzag array. <i>Nature Structural and Molecular Biology</i> , 2011, 18, 927-933.	8.2	149
34	The Effect of Hydrophilic Substitutions and Anionic Lipids upon the Transverse Positioning of the Transmembrane Helix of the ErbB2 (neu) Protein Incorporated into Model Membrane Vesicles. <i>Journal of Molecular Biology</i> , 2010, 396, 209-220.	4.2	18
35	Effect of Sequence Hydrophobicity and Bilayer Width upon the Minimum Length Required for the Formation of Transmembrane Helices in Membranes. <i>Journal of Molecular Biology</i> , 2007, 374, 671-687.	4.2	48
36	The Control of Transmembrane Helix Transverse Position in Membranes by Hydrophilic Residues. <i>Journal of Molecular Biology</i> , 2007, 374, 1251-1269.	4.2	36

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37	Membrane Topography of the Hydrophobic Anchor Sequence of Poliovirus 3A and 3AB Proteins and the Functional Effect of 3A/3AB Membrane Association upon RNA Replication. <i>Biochemistry</i> , 2007, 46, 5185-5199.	2.5	65
38	Multiple-probe analysis of folding and unfolding pathways of human serum albumin. Evidence for a framework mechanism of folding. <i>FEBS Journal</i> , 2004, 271, 1789-1797.	0.2	53
39	Active and Inactive Orientations of the Transmembrane and Cytosolic Domains of the Erythropoietin Receptor Dimer. <i>Molecular Cell</i> , 2003, 12, 1239-1250.	9.7	193
40	Glutamate-induced Assembly of Bacterial Cell Division Protein FtsZ. <i>Journal of Biological Chemistry</i> , 2003, 278, 3735-3741.	3.4	59
41	Spatial Relationship between the Prodan Site, Trp-214, and Cys-34 Residues in Human Serum Albumin and Loss of Structure through Incremental Unfolding. <i>Biochemistry</i> , 2002, 41, 7443-7452.	2.5	63