

Dirk Fasshauer

List of Publications by Year in descending order

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71
papers

10,417
citations

61977

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110368

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79
all docs

79
docs citations

79
times ranked

6640
citing authors

#	ARTICLE	IF	CITATIONS
1	Choanoflagellates and the ancestry of neurosecretory vesicles. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2021, 376, 20190759.	4.0	17
2	A conformational switch driven by phosphorylation regulates the activity of the evolutionarily conserved SNARE Ykt6. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	12
3	Hidden cell diversity in Placozoa: ultrastructural insights from <i>Hoilungia hongkongensis</i> . <i>Cell and Tissue Research</i> , 2021, 385, 623-637.	2.9	22
4	<i>BET1</i> variants establish impaired vesicular transport as a cause for muscular dystrophy with epilepsy. <i>EMBO Molecular Medicine</i> , 2021, 13, e13787.	6.9	9
5	The diversification and lineage-specific expansion of nitric oxide signaling in Placozoa: insights in the evolution of gaseous transmission. <i>Scientific Reports</i> , 2020, 10, 13020.	3.3	37
6	PI(4,5)P ₂ -dependent regulation of exocytosis by amisyn, the vertebrate-specific competitor of synaptobrevin 2. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 13468-13479.	7.1	10
7	Glycine as a signaling molecule and chemoattractant in <i>Trichoplax</i> (Placozoa): insights into the early evolution of neurotransmitters. <i>NeuroReport</i> , 2020, 31, 490-497.	1.2	27
8	Prototypic SNARE Proteins Are Encoded in the Genomes of <i>Heimdallarchaeota</i> , Potentially Bridging the Gap between the Prokaryotes and Eukaryotes. <i>Current Biology</i> , 2020, 30, 2468-2480.e5.	3.9	24
9	Probing the Conformational Flexibility of the Munc18-1/Syntaxin-1A Complex. <i>Biophysical Journal</i> , 2020, 118, 24a.	0.5	0
10	High Cell Diversity and Complex Peptidergic Signaling Underlie Placozoan Behavior. <i>Current Biology</i> , 2018, 28, 3495-3501.e2.	3.9	84
11	Getting Nervous: An Evolutionary Overhaul for Communication. <i>Annual Review of Genetics</i> , 2017, 51, 455-476.	7.6	44
12	Functional assays for the assessment of the pathogenicity of variants in <i>GOSR2</i> , an ER-to-Golgi SNARE involved in progressive myoclonus epilepsies. <i>DMM Disease Models and Mechanisms</i> , 2017, 10, 1391-1398.	2.4	11
13	Evidence for a conserved inhibitory binding mode between the membrane fusion assembly factors Munc18 and syntaxin in animals. <i>Journal of Biological Chemistry</i> , 2017, 292, 20449-20460.	3.4	11
14	Shedding light on the expansion and diversification of the Cdc48 protein family during the rise of the eukaryotic cell. <i>BMC Evolutionary Biology</i> , 2016, 16, 215.	3.2	15
15	Analysis of Scfd2 - A New Member of the SM Protein Family. <i>Biophysical Journal</i> , 2016, 110, 597a.	0.5	0
16	The SM protein Sly1 accelerates assembly of the ER-Golgi SNARE complex. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 13828-13833.	7.1	40
17	Novel Cell Types, Neurosecretory Cells, and Body Plan of the Early-Diverging Metazoan <i>Trichoplax adhaerens</i> . <i>Current Biology</i> , 2014, 24, 1565-1572.	3.9	209
18	Interaction of Munc18C with Syntaxin4 and the Role of Munc18C in Exocytosis. <i>Biophysical Journal</i> , 2014, 106, 311a.	0.5	0

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19	Phosphatidylinositol 4,5-bisphosphate clusters act as molecular beacons for vesicle recruitment. <i>Nature Structural and Molecular Biology</i> , 2013, 20, 679-686.	8.2	246
20	Syntaxin1a variants lacking an N-peptide or bearing the LE mutation bind to Munc18a in a closed conformation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 12637-12642.	7.1	58
21	Munc18-1 mutations that strongly impair SNARE-complex binding support normal synaptic transmission. <i>EMBO Journal</i> , 2012, 31, 2156-2168.	7.8	62
22	Molecular machines governing exocytosis of synaptic vesicles. <i>Nature</i> , 2012, 490, 201-207.	27.8	830
23	Untangling the evolution of Rab G proteins: implications of a comprehensive genomic analysis. <i>BMC Biology</i> , 2012, 10, 71.	3.8	159
24	Primordial neurosecretory apparatus identified in the choanoflagellate <i>Monosiga brevicollis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 15264-15269.	7.1	74
25	A Coiled Coil Trigger Site Is Essential for Rapid Binding of Synaptobrevin to the SNARE Acceptor Complex. <i>Journal of Biological Chemistry</i> , 2010, 285, 21549-21559.	3.4	25
26	Single vesicle millisecond fusion kinetics reveals number of SNARE complexes optimal for fast SNARE-mediated membrane fusion.. <i>Journal of Biological Chemistry</i> , 2010, 285, 11753.	3.4	5
27	Synaptobrevin N-terminally bound to syntaxinâ€“SNAP-25 defines the primed vesicle state in regulated exocytosis. <i>Journal of Cell Biology</i> , 2010, 188, 401-413.	5.2	115
28	Synaptobrevin N-terminally bound to syntaxinâ€“SNAP-25 defines the primed vesicle state in regulated exocytosis. <i>Journal of General Physiology</i> , 2010, 135, i2-i2.	1.9	0
29	The Ca ²⁺ Affinity of Synaptotagmin 1 Is Markedly Increased by a Specific Interaction of Its C2B Domain with Phosphatidylinositol 4,5-Bisphosphate. <i>Journal of Biological Chemistry</i> , 2009, 284, 25749-25760.	3.4	125
30	Single Vesicle Millisecond Fusion Kinetics Reveals Number of SNARE Complexes Optimal for Fast SNARE-mediated Membrane Fusion. <i>Journal of Biological Chemistry</i> , 2009, 284, 32158-32166.	3.4	148
31	Is Assembly of the SNARE Complex Enough to Fuel Membrane Fusion?. <i>Journal of Biological Chemistry</i> , 2009, 284, 13143-13152.	3.4	65
32	A Conserved Membrane Attachment Site in Î±-SNAP Facilitates N-Ethylmaleimide-sensitive Factor (NSF)-driven SNARE Complex Disassembly. <i>Journal of Biological Chemistry</i> , 2009, 284, 31817-31826.	3.4	55
33	Phylogeny of the SNARE vesicle fusion machinery yields insights into the conservation of the secretory pathway in fungi. <i>BMC Evolutionary Biology</i> , 2009, 9, 19.	3.2	51
34	Differences in the SNARE evolution of fungi and metazoa. <i>Biochemical Society Transactions</i> , 2009, 37, 787-791.	3.4	23
35	Insights Into The Energetics Of Neuronal SNARE Complex Formation. <i>Biophysical Journal</i> , 2009, 96, 357a.	0.5	0
36	Munc18a controls SNARE assembly through its interaction with the syntaxin N-peptide. <i>EMBO Journal</i> , 2008, 27, 923-933.	7.8	237

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37	Imaging the assembly and disassembly kinetics of cis-SNARE complexes on native plasma membranes. <i>FEBS Letters</i> , 2008, 582, 3563-3568.	2.8	15
38	SNAREing the Basis of Multicellularity: Consequences of Protein Family Expansion during Evolution. <i>Molecular Biology and Evolution</i> , 2008, 25, 2055-2068.	8.9	73
39	A Novel Site of Action for $\hat{\pm}$ -SNAP in the SNARE Conformational Cycle Controlling Membrane Fusion. <i>Molecular Biology of the Cell</i> , 2008, 19, 776-784.	2.1	41
40	Determinants of Synaptobrevin Regulation in Membranes. <i>Molecular Biology of the Cell</i> , 2007, 18, 2037-2046.	2.1	58
41	An Elaborate Classification of SNARE Proteins Sheds Light on the Conservation of the Eukaryotic Endomembrane System. <i>Molecular Biology of the Cell</i> , 2007, 18, 3463-3471.	2.1	201
42	Early endosomal SNAREs form a structurally conserved SNARE complex and fuse liposomes with multiple topologies. <i>EMBO Journal</i> , 2007, 26, 9-18.	7.8	71
43	Budding insights on cell polarity. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 360-362.	8.2	10
44	Synaptotagmin activates membrane fusion through a Ca ²⁺ -dependent trans interaction with phospholipids. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 904-911.	8.2	152
45	N- to C-Terminal SNARE Complex Assembly Promotes Rapid Membrane Fusion. <i>Science</i> , 2006, 313, 673-676.	12.6	343
46	Identification of SNAP-47, a Novel Qbc-SNARE with Ubiquitous Expression*. <i>Journal of Biological Chemistry</i> , 2006, 281, 17076-17083.	3.4	90
47	Sequential N- to C-terminal SNARE complex assembly drives priming and fusion of secretory vesicles. <i>EMBO Journal</i> , 2006, 25, 955-966.	7.8	251
48	Alternative Splicing of SNAP-25 Regulates Secretion through Nonconservative Substitutions in the SNARE Domain. <i>Molecular Biology of the Cell</i> , 2005, 16, 5675-5685.	2.1	61
49	A Transient N-terminal Interaction of SNAP-25 and Syntaxin Nucleates SNARE Assembly. <i>Journal of Biological Chemistry</i> , 2004, 279, 7613-7621.	3.4	165
50	Structural Basis for the Inhibitory Role of Tomosyn in Exocytosis. <i>Journal of Biological Chemistry</i> , 2004, 279, 47192-47200.	3.4	100
51	Structural insights into the SNARE mechanism. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2003, 1641, 87-97.	4.1	96
52	The Habc Domain and the SNARE Core Complex Are Connected by a Highly Flexible Linker. <i>Biochemistry</i> , 2003, 42, 4009-4014.	2.5	41
53	Crystal Structure of a Complex Between Human Spliceosomal Cyclophilin H and a U4/U6 snRNP-60K Peptide. <i>Journal of Molecular Biology</i> , 2003, 331, 45-56.	4.2	46
54	The R-SNARE Motif of Tomosyn Forms SNARE Core Complexes with Syntaxin 1 and SNAP-25 and Down-regulates Exocytosis. <i>Journal of Biological Chemistry</i> , 2003, 278, 31159-31166.	3.4	122

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55	Single-molecule fluorescence resonance energy transfer reveals a dynamic equilibrium between closed and open conformations of syntaxin 1. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 15516-15521.	7.1	268
56	Rapid and Selective Binding to the Synaptic SNARE Complex Suggests a Modulatory Role of Complexins in Neuroexocytosis. Journal of Biological Chemistry, 2002, 277, 7838-7848.	3.4	121
57	Crystal structure of the endosomal SNARE complex reveals common structural principles of all SNAREs. Nature Structural Biology, 2002, 9, 107-111.	9.7	239
58	SNARE assembly and disassembly exhibit a pronounced hysteresis. Nature Structural Biology, 2002, 9, 144-151.	9.7	141
59	Homo- and Heterooligomeric SNARE Complexes Studied by Site-directed Spin Labeling. Journal of Biological Chemistry, 2001, 276, 13169-13177.	3.4	115
60	A SNARE complex mediating fusion of late endosomes defines conserved properties of SNARE structure and function. EMBO Journal, 2000, 19, 6453-6464.	7.8	245
61	Selective Interaction of Complexin with the Neuronal SNARE Complex. Journal of Biological Chemistry, 2000, 275, 19808-19818.	3.4	162
62	Mixed and Non-cognate SNARE Complexes. Journal of Biological Chemistry, 1999, 274, 15440-15446.	3.4	271
63	Kinetics of Synaptotagmin Responses to Ca ²⁺ and Assembly with the Core SNARE Complex onto Membranes. Neuron, 1999, 24, 363-376.	8.1	258
64	Crystal structure of a SNARE complex involved in synaptic exocytosis at 2.4-Å resolution. Nature, 1998, 395, 347-353.	27.8	2,191
65	Identification of a Minimal Core of the Synaptic SNARE Complex Sufficient for Reversible Assembly and Disassembly. Biochemistry, 1998, 37, 10354-10362.	2.5	239
66	Conserved structural features of the synaptic fusion complex: SNARE proteins reclassified as Q- and R-SNAREs. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 15781-15786.	7.1	860
67	A Structural Change Occurs upon Binding of Syntaxin to SNAP-25. Journal of Biological Chemistry, 1997, 272, 4582-4590.	3.4	167
68	Structural Changes Are Associated with Soluble N-Ethylmaleimide-sensitive Fusion Protein Attachment Protein Receptor Complex Formation. Journal of Biological Chemistry, 1997, 272, 28036-28041.	3.4	308
69	ADP ribosylation factor and a 14-kD polypeptide are associated with heparan sulfate-carrying post-trans-Golgi network secretory vesicles in rat hepatocytes. Journal of Cell Biology, 1994, 125, 721-732.	5.2	35
70	ARF and VAPP14: Two Proteins Involved in the Delivery of Heparan Sulfate Proteoglycan from the trans-Golgi Network to the Plasma Membrane. Annals of the New York Academy of Sciences, 1994, 733, 344-356.	3.8	0
71	Megaviruses contain various genes encoding for eukaryotic vesicle trafficking factors. Traffic, 0, , .	2.7	5