Rajesh Ramachandran

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular characterization of linker and loop-mediated structural modulation and hinge motion in the C4-C5 domains of cMyBPC. Journal of Structural Biology, 2022, 214, 107856.	2.8	5
2	Identification of Phosphorylation and Other Post-Translational Modifications in the Central C4C5 Domains of Murine Cardiac Myosin Binding Protein C. ACS Omega, 2022, 7, 14189-14202.	3.5	4
3	Reply to Roy and Pucadyil: A gain of function by a GTPase-impaired Drp1. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	1
4	Drp1 and the cytoskeleton: mechanistic nexus in mitochondrial division. Current Opinion in Physiology, 2022, 29, 100574.	1.8	2
5	NMR identification of a conserved Drp1 cardiolipin-binding motif essential for stress-induced mitochondrial fission. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	31
6	GFP fluorescence tagging alters dynamin-related proteinÂ1 oligomerization dynamics and creates disassembly-refractory puncta to mediate mitochondrial fission. Scientific Reports, 2020, 10, 14777.	3.3	18
7	Imaging Dynamin-Related Protein 1 (Drp1)-Mediated Mitochondrial Fission in Living Cells. Methods in Molecular Biology, 2020, 2159, 205-217.	0.9	5
8	The cryo-EM structure of the SNX–BAR Mvp1 tetramer. Nature Communications, 2020, 11, 1506.	12.8	22
9	Microscale Thermophoresis (MST) as a Tool for Measuring Dynamin Superfamily Protein (DSP)–Lipid Interactions. Methods in Molecular Biology, 2020, 2159, 85-92.	0.9	2
10	AAV9 gene transfer of cMyBPC N-terminal domains ameliorates cardiomyopathy in cMyBPC-deficient mice. JCI Insight, 2020, 5, .	5.0	18
11	A Single Common Protocol for the Expression and Purification of Soluble Mammalian DSPs from Escherichia coli. Methods in Molecular Biology, 2020, 2159, 31-40.	0.9	2
12	Exploring the links between lipid geometry and mitochondrial fission: Emerging concepts. Mitochondrion, 2019, 49, 305-313.	3.4	18
13	Crystal Structures of the Full-Length Murine and Human Gasdermin D Reveal Mechanisms of Autoinhibition, Lipid Binding, and Oligomerization. Immunity, 2019, 51, 43-49.e4.	14.3	151
14	ATAD3A oligomerization causes neurodegeneration by coupling mitochondrial fragmentation and bioenergetics defects. Nature Communications, 2019, 10, 1371.	12.8	59
15	Geometric instability catalyzes mitochondrial fission. Molecular Biology of the Cell, 2019, 30, 160-168.	2.1	10
16	The dynamin superfamily. Current Biology, 2018, 28, R411-R416.	3.9	93
17	Mitochondrial dynamics: The dynamin superfamily and execution by collusion. Seminars in Cell and Developmental Biology, 2018, 76, 201-212.	5.0	67
18	Steric interference from intrinsically disordered regions controls dynamin-related protein 1 self-assembly during mitochondrial fission. Scientific Reports, 2018, 8, 10879.	3.3	27

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19	Cooperative and independent roles of Drp1 adaptors Mff and MiD49/51 in mitochondrial fission. Journal of Cell Science, 2016, 129, 2170-81.	2.0	234
20	Coincident Phosphatidic Acid Interaction Restrains Drp1 in Mitochondrial Division. Molecular Cell, 2016, 63, 1034-1043.	9.7	150
21	Distinct Splice Variants of Dynamin-related Protein 1 Differentially Utilize Mitochondrial Fission Factor as an Effector of Cooperative GTPase Activity. Journal of Biological Chemistry, 2016, 291, 493-507.	3.4	78
22	Dynamin-related Protein 1 Oligomerization in Solution Impairs Functional Interactions with Membrane-anchored Mitochondrial Fission Factor. Journal of Biological Chemistry, 2016, 291, 478-492.	3.4	78
23	The Mechanoenzymatic Core of Dynamin-related Protein 1 Comprises the Minimal Machinery Required for Membrane Constriction. Journal of Biological Chemistry, 2015, 290, 11692-11703.	3.4	96
24	Cardiolipin's propensity for phase transition and its reorganization by dynamin-related protein 1 form a basis for mitochondrial membrane fission. Molecular Biology of the Cell, 2015, 26, 3104-3116.	2.1	129
25	Syndapin 3 modulates fusion pore expansion in mouse neuroendocrine chromaffin cells. American Journal of Physiology - Cell Physiology, 2014, 306, C831-C843.	4.6	9
26	Alternate pleckstrin homology domain orientations regulate dynamin-catalyzed membrane fission. Molecular Biology of the Cell, 2014, 25, 879-890.	2.1	34
27	A dimeric equilibrium intermediate nucleates Drp1 reassembly on mitochondrial membranes for fission. Molecular Biology of the Cell, 2014, 25, 1905-1915.	2.1	149
28	Vesicle scission: Dynamin. Seminars in Cell and Developmental Biology, 2011, 22, 10-17.	5.0	50
29	Differential curvature sensing and generating activities of dynamin isoforms provide opportunities for tissue-specific regulation. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, E234-42.	7.1	87
30	Membrane Insertion of the Pleckstrin Homology Domain Variable Loop 1 Is Critical for Dynamin-catalyzed Vesicle Scission. Molecular Biology of the Cell, 2009, 20, 4630-4639.	2.1	94
31	Dissecting dynamin's role in clathrin-mediated endocytosis. Biochemical Society Transactions, 2009, 37, 1022-1026.	3.4	169
32	Real-time detection reveals that effectors couple dynamin's GTP-dependent conformational changes to the membrane. EMBO Journal, 2008, 27, 27-37.	7.8	102
33	Supported lipid bilayer array to study clathrin mediated endocytosis in vitro. , 2007, , .		0
34	The dynamin middle domain is critical for tetramerization and higher-order self-assembly. EMBO Journal, 2007, 26, 559-566.	7.8	164
35	Physical and functional connection between auxilin and dynamin during endocytosis. EMBO Journal, 2006, 25, 4163-4174.	7.8	29
36	The domains of a cholesterol-dependent cytolysin undergo a major FRET-detected rearrangement during pore formation. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 7139-7144.	7.1	87

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37	Robust Colorimetric Assays for Dynamin's Basal and Stimulated GTPase Activities. Methods in Enzymology, 2005, 404, 490-503.	1.0	97
38	Membrane-dependent conformational changes initiate cholesterol-dependent cytolysin oligomerization and intersubunit β-strand alignment. Nature Structural and Molecular Biology, 2004, 11, 697-705.	8.2	168
39	Structural insights into the membrane-anchoring mechanism of a cholesterol-dependent cytolysin. Nature Structural Biology, 2002, 9, 823-7.	9.7	116