## Miriam-Rose Ash

List of Publications by Year in descending order

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933447 1199594 12 416 10 12 citations h-index g-index papers 12 12 12 580 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Structure and autoregulation of a P4-ATPase lipid flippase. Nature, 2019, 571, 366-370.	27.8	126
2	High phosphatidylinositol 4-phosphate (PI4P)-dependent ATPase activity for the Drs2p-Cdc50p flippase after removal of its N- and C-terminal extensions. Journal of Biological Chemistry, 2017, 292, 7954-7970.	3.4	29
3	A High-Yield Co-Expression System for the Purification of an Intact Drs2p-Cdc50p Lipid Flippase Complex, Critically Dependent on and Stabilized by Phosphatidylinositol-4-Phosphate. PLoS ONE, 2014, 9, e112176.	2.5	23
4	A GTPase Chimera Illustrates an Uncoupled Nucleotide Affinity and Release Rate, Providing Insight into the Activation Mechanism. Biophysical Journal, 2014, 107, L45-L48.	0.5	4
5	The cationâ€dependent Gâ€proteins: In a class of their own. FEBS Letters, 2012, 586, 2218-2224.	2.8	37
6	Molecular Basis of the Cooperative Binding of Cu(I) and Cu(II) to the CopK Protein from <i>Cupriavidus metallidurans </i>	2.5	18
7	The Initiation of GTP Hydrolysis by the G-Domain of FeoB: Insights from a Transition-State Complex Structure. PLoS ONE, 2011, 6, e23355.	2.5	34
8	A suite of Switch I and Switch II mutant structures from the G-protein domain of FeoB. Acta Crystallographica Section D: Biological Crystallography, 2011, 67, 973-980.	2.5	14
9	The structure of an N11A mutant of the G-protein domain of FeoB. Acta Crystallographica Section F: Structural Biology Communications, 2011, 67, 1511-1515.	0.7	8
10	Conserved Î <sup>2</sup> -Hairpin Recognition by the GYF Domains of Smy2 and GIGYF2 in mRNA Surveillance and Vesicular Transport Complexes. Structure, 2010, 18, 944-954.	3.3	34
11	Potassium-activated GTPase Reaction in the G Protein-coupled Ferrous Iron Transporter B. Journal of Biological Chemistry, 2010, 285, 14594-14602.	3.4	51
12	Unprecedented Binding Cooperativity between Cu <sup>I</sup> and Cu <sup>II</sup> in the Copper Resistance Protein CopK from <i>Cupriavidus metallidurans</i> CH34: Implications from Structural Studies by NMR Spectroscopy and X-Ray Crystallography. Journal of the American Chemical Society, 2009, 131, 3549-3564.	13.7	38