## Rachel M Johnson

List of Publications by Year in descending order

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858243 1181555 14 367 12 14 citations h-index g-index papers 14 14 14 636 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A structural basis for amylin receptor phenotype. Science, 2022, 375, eabm9609.	6.0	28
2	A Structurally Characterized <i>Staphylococcus aureus</i> Evolutionary Escape Route from Treatment with the Antibiotic Linezolid. Microbiology Spectrum, 2022, 10, .	1.2	4
3	Membranes under the Magnetic Lens: A Dive into the Diverse World of Membrane Protein Structures Using Cryo-EM. Chemical Reviews, 2022, 122, 13989-14017.	23.0	17
4	Evolving cryo-EM structural approaches for GPCR drug discovery. Structure, 2021, 29, 963-974.e6.	1.6	29
5	Cryo-EM structure of the dual incretin receptor agonist, peptide-19, in complex with the glucagon-like peptide-1 receptor. Biochemical and Biophysical Research Communications, 2021, 578, 84-90.	1.0	14
6	Human TRPC5 structures reveal interaction of a xanthine-based TRPC1/4/5 inhibitor with a conserved lipid binding site. Communications Biology, 2020, 3, 704.	2.0	36
7	Structure-Based Identification and Characterization of Inhibitors of the Epilepsy-Associated KNa1.1 (KCNT1) Potassium Channel. IScience, 2020, 23, 101100.	1.9	29
8	Potent Tetrahydroquinolone Eliminates Apicomplexan Parasites. Frontiers in Cellular and Infection Microbiology, 2020, 10, 203.	1.8	21
9	Cryo-EM Structure and Molecular Dynamics Analysis of the Fluoroquinolone Resistant Mutant of the AcrB Transporter from Salmonella. Microorganisms, 2020, 8, 943.	1.6	25
10	Emerging Role of Electron Microscopy in Drug Discovery. Trends in Biochemical Sciences, 2019, 44, 897-898.	3.7	3
11	Dimeric structures of quinol-dependent nitric oxide reductases (qNORs) revealed by cryo–electron microscopy. Science Advances, 2019, 5, eaax1803.	4.7	14
12	LAT1 (SLC7A5) and CD98hc (SLC3A2) complex dynamics revealed by single-particle cryo-EM. Acta Crystallographica Section D: Structural Biology, 2019, 75, 660-669.	1.1	16
13	Approaches to altering particle distributions in cryo-electron microscopy sample preparation. Acta Crystallographica Section D: Structural Biology, 2018, 74, 560-571.	1.1	108
14	X-ray and cryo-EM structures of inhibitor-bound cytochrome <i>bc</i> csub>1complexes for structure-based drug discovery. IUCrJ, 2018, 5, 200-210.	1.0	23