

Shanlin Rao

List of Publications by Year in descending order

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papers

740
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686830

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991
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular Simulations of Hydrophobic Gating of Pentameric Ligand Gated Ion Channels: Insights into Water and Ions. <i>Journal of Physical Chemistry B</i> , 2021, 125, 981-994.	1.2	27
2	Norfluoxetine inhibits TREK-2 K2P channels by multiple mechanisms including state-independent effects on the selectivity filter gate. <i>Journal of General Physiology</i> , 2021, 153, .	0.9	17
3	Water Nanoconfined in a Hydrophobic Pore: Molecular Dynamics Simulations of Transmembrane Protein 175 and the Influence of Water Models. <i>ACS Nano</i> , 2021, 15, 19098-19108.	7.3	14
4	Mechanisms of activation and desensitization of full-length glycine receptor in lipid nanodiscs. <i>Nature Communications</i> , 2020, 11, 3752.	5.8	74
5	Water in Nanopores and Biological Channels: A Molecular Simulation Perspective. <i>Chemical Reviews</i> , 2020, 120, 10298-10335.	23.0	121
6	The MscS-like channel YnaI has a gating mechanism based on flexible pore helices. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 28754-28762.	3.3	30
7	Characterizing Membrane Association and Periplasmic Transfer of Bacterial Lipoproteins through Molecular Dynamics Simulations. <i>Structure</i> , 2020, 28, 475-487.e3.	1.6	15
8	Structure and assembly of calcium homeostasis modulator proteins. <i>Nature Structural and Molecular Biology</i> , 2020, 27, 150-159.	3.6	55
9	Induced Polarization in Molecular Dynamics Simulations of the 5-HT ₃ Receptor Channel. <i>Journal of the American Chemical Society</i> , 2020, 142, 9415-9427.	6.6	38
10	A heuristic derived from analysis of the ion channel structural proteome permits the rapid identification of hydrophobic gates. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 13989-13995.	3.3	52
11	CHAP: A Versatile Tool for the Structural and Functional Annotation of Ion Channel Pores. <i>Journal of Molecular Biology</i> , 2019, 431, 3353-3365.	2.0	97
12	A Newly Available Tool for Functional Annotation of Ion Channel Structures Based on Molecular Dynamics Simulations. <i>Biophysical Journal</i> , 2018, 114, 134a.	0.2	1
13	Water and hydrophobic gates in ion channels and nanopores. <i>Faraday Discussions</i> , 2018, 209, 231-247.	1.6	48
14	Cryo-EM reveals two distinct serotonin-bound conformations of full-length 5-HT _{3A} receptor. <i>Nature</i> , 2018, 563, 270-274.	13.7	98
15	A BEST example of channel structure annotation by molecular simulation. <i>Channels</i> , 2017, 11, 347-353.	1.5	26