

Toby W Allen

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

67

papers

4,483

citations

37

h-index

66

g-index

74

ext. papers

4,962

ext. citations

5.6

avg, IF

5.68

L-index

#	Paper	IF	Citations
67	Energetics of ion conduction through the gramicidin channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 117-22	11.5	335
66	Theoretical and computational models of biological ion channels. <i>Quarterly Reviews of Biophysics</i> , 2004 , 37, 15-103	7	321
65	On the thermodynamic stability of a charged arginine side chain in a transmembrane helix. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 4943-8	11.5	221
64	The different interactions of lysine and arginine side chains with lipid membranes. <i>Journal of Physical Chemistry B</i> , 2013 , 117, 11906-20	3.4	176
63	Permeation of ions across the potassium channel: Brownian dynamics studies. <i>Biophysical Journal</i> , 1999 , 77, 2517-33	2.9	176
62	Molecular dynamics - potential of mean force calculations as a tool for understanding ion permeation and selectivity in narrow channels. <i>Biophysical Chemistry</i> , 2006 , 124, 251-67	3.5	155
61	Mechanisms of permeation and selectivity in calcium channels. <i>Biophysical Journal</i> , 2001 , 80, 195-214	2.9	151
60	Molecular dynamics study of the KcsA potassium channel. <i>Biophysical Journal</i> , 1999 , 77, 2502-16	2.9	138
59	Mechanism of potassium-channel selectivity revealed by Na(+) and Li(+) binding sites within the KcsA pore. <i>Nature Structural and Molecular Biology</i> , 2009 , 16, 1317-24	17.6	131
58	The role of tryptophan side chains in membrane protein anchoring and hydrophobic mismatch. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013 , 1828, 864-76	3.8	130
57	On the importance of atomic fluctuations, protein flexibility, and solvent in ion permeation. <i>Journal of General Physiology</i> , 2004 , 124, 679-90	3.4	127
56	Ion permeation through a narrow channel: using gramicidin to ascertain all-atom molecular dynamics potential of mean force methodology and biomolecular force fields. <i>Biophysical Journal</i> , 2006 , 90, 3447-68	2.9	123
55	The potassium channel: Structure, selectivity and diffusion. <i>Journal of Chemical Physics</i> , 2000 , 112, 8191-8204	3.4	122
54	Conducting-state properties of the KcsA potassium channel from molecular and Brownian dynamics simulations. <i>Biophysical Journal</i> , 2002 , 82, 628-45	2.9	120
53	Structure of gramicidin a in a lipid bilayer environment determined using molecular dynamics simulations and solid-state NMR data. <i>Journal of the American Chemical Society</i> , 2003 , 125, 9868-77	16.4	115
52	Potential of mean force and pKa profile calculation for a lipid membrane-exposed arginine side chain. <i>Journal of Physical Chemistry B</i> , 2008 , 112, 9574-87	3.4	102
51	Gramicidin A channel as a test ground for molecular dynamics force fields. <i>Biophysical Journal</i> , 2003 , 84, 2159-68	2.9	102

50	Membrane Protein Structure, Function, and Dynamics: a Perspective from Experiments and Theory. <i>Journal of Membrane Biology</i> , 2015 , 248, 611-40	2.3	101
49	Study of ionic currents across a model membrane channel using Brownian dynamics. <i>Biophysical Journal</i> , 1998 , 75, 793-809	2.9	101
48	Assessing atomistic and coarse-grained force fields for protein-lipid interactions: the formidable challenge of an ionizable side chain in a membrane. <i>Journal of Physical Chemistry B</i> , 2008 , 112, 9588-6023-4	3.4	99
47	Modeling diverse range of potassium channels with Brownian dynamics. <i>Biophysical Journal</i> , 2002 , 83, 263-77	2.9	83
46	The effect of hydrophobic and hydrophilic channel walls on the structure and diffusion of water and ions. <i>Journal of Chemical Physics</i> , 1999 , 111, 7985-7999	3.9	79
45	Ion-induced defect permeation of lipid membranes. <i>Biophysical Journal</i> , 2014 , 106, 586-97	2.9	77
44	Ion conduction and conformational flexibility of a bacterial voltage-gated sodium channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 3454-9	11.5	77
43	Is arginine charged in a membrane?. <i>Biophysical Journal</i> , 2008 , 94, L11-3	2.9	74
42	On the selective ion binding hypothesis for potassium channels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 17963-8	11.5	72
41	Local anesthetic and antiepileptic drug access and binding to a bacterial voltage-gated sodium channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 13057-62	11.5	67
40	Origins of ion selectivity in potassium channels from the perspective of channel block. <i>Journal of General Physiology</i> , 2011 , 137, 405-13	3.4	60
39	Molecular dynamics estimates of ion diffusion in model hydrophobic and KcsA potassium channels. <i>Biophysical Chemistry</i> , 2000 , 86, 1-14	3.5	57
38	The role of membrane thickness in charged protein-lipid interactions. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012 , 1818, 135-45	3.8	55
37	Reservoir boundaries in Brownian dynamics simulations of ion channels. <i>Biophysical Journal</i> , 2002 , 82, 1975-84	2.9	53
36	Atomistic Simulations of Membrane Ion Channel Conduction, Gating, and Modulation. <i>Chemical Reviews</i> , 2019 , 119, 7737-7832	68.1	46
35	The Role of Atomic Polarization in the Thermodynamics of Chloroform Partitioning to Lipid Bilayers. <i>Journal of Chemical Theory and Computation</i> , 2012 , 8, 618-28	6.4	42
34	Electrostatics of deformable lipid membranes. <i>Biophysical Journal</i> , 2010 , 98, 2904-13	2.9	42
33	String method solution of the gating pathways for a pentameric ligand-gated ion channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, E4158-E4167	11.5	37

32	Towards a Structural View of Drug Binding to hERG K Channels. <i>Trends in Pharmacological Sciences</i> , 2017 , 38, 899-907	13.2	37
31	The electrostatics of solvent and membrane interfaces and the role of electronic polarizability. <i>Journal of Chemical Physics</i> , 2010 , 132, 185101	3.9	37
30	The determinants of hydrophobic mismatch response for transmembrane helices. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013 , 1828, 851-63	3.8	35
29	On the role of anionic lipids in charged protein interactions with membranes. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2011 , 1808, 1673-83	3.8	34
28	Evidence for leaflet-dependent redistribution of charged molecules in fluid supported phospholipid bilayers. <i>Langmuir</i> , 2008 , 24, 13250-3	4	34
27	Modeling charged protein side chains in lipid membranes. <i>Journal of General Physiology</i> , 2007 , 130, 237-44	3.4	31
26	Molecular and Brownian dynamics study of ion selectivity and conductivity in the potassium channel. <i>Chemical Physics Letters</i> , 1999 , 313, 358-365	2.5	31
25	A model of calcium channels. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2000 , 1509, 1-6	3.8	29
24	A selectivity filter at the intracellular end of the acid-sensing ion channel pore. <i>ELife</i> , 2017 , 6,	8.9	27
23	The Potassium Ion Channel: Comparison of Linear Scaling Semiempirical and Molecular Mechanics Representations of the Electrostatic Potential. <i>Journal of Physical Chemistry B</i> , 2001 , 105, 12674-12679	3.4	24
22	BennettQ acceptance ratio and histogram analysis methods enhanced by umbrella sampling along a reaction coordinate in configurational space. <i>Journal of Chemical Physics</i> , 2012 , 136, 164103	3.9	21
21	Ball-and-chain inactivation in a calcium-gated potassium channel. <i>Nature</i> , 2020 , 580, 288-293	50.4	20
20	Estimation of Potentials of Mean Force from Nonequilibrium Pulling Simulations Using Both Minh-Adib Estimator and Weighted Histogram Analysis Method. <i>Journal of Chemical Theory and Computation</i> , 2016 , 12, 1000-10	6.4	19
19	Comment on "Free energy simulations of single and double ion occupancy in gramicidin A" [J. Chem. Phys. 126, 105103 (2007)]. <i>Journal of Chemical Physics</i> , 2008 , 128, 227101; author reply 227102	3.9	19
18	Cholesterol depletion inhibits Na,K-ATPase activity in a near-native membrane environment. <i>Journal of Biological Chemistry</i> , 2019 , 294, 5956-5969	5.4	17
17	Identification of electric-field-dependent steps in the Na(+),K(+)-pump cycle. <i>Biophysical Journal</i> , 2014 , 107, 1352-63	2.9	15
16	Computational Models for Predictive Cardiac Ion Channel Pharmacology. <i>Drug Discovery Today: Disease Models</i> , 2014 , 14, 3-10	1.3	11
15	Selectivity filter ion binding affinity determines inactivation in a potassium channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 29968-29978	11.5	11

14	The voltage-sensitive dye RH421 detects a Na,K-ATPase conformational change at the membrane surface. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2017 , 1859, 813-823	3.8	10
13	Selective ion permeation involves complexation with carboxylates and lysine in a model human sodium channel. <i>PLoS Computational Biology</i> , 2018 , 14, e1006398	5	10
12	Positronium states in three-dimensional QED. <i>Physical Review D</i> , 1996 , 53, 5842-5855	4.9	8
11	Polarization Effects in Water-Mediated Selective Cation Transport across a Narrow Transmembrane Channel. <i>Journal of Chemical Theory and Computation</i> , 2021 , 17, 1726-1741	6.4	7
10	Understanding Sodium Channel Function and Modulation Using Atomistic Simulations of Bacterial Channel Structures. <i>Current Topics in Membranes</i> , 2016 , 78, 145-82	2.2	5
9	Determinants of ion selectivity in ASIC1a- and ASIC2a-containing acid-sensing ion channels. <i>Journal of General Physiology</i> , 2020 , 152,	3.4	5
8	Simulating ion channel activation mechanisms using swarms of trajectories. <i>Journal of Computational Chemistry</i> , 2020 , 41, 387-401	3.5	4
7	Comparison of permeation mechanisms in sodium-selective ion channels. <i>Neuroscience Letters</i> , 2019 , 700, 3-8	3.3	3
6	Conformational Dynamics in a Nicotinic Receptor Homologue Probed by Simulations. <i>Biophysical Journal</i> , 2011 , 100, 272a	2.9	3
5	Molecular Dynamics Simulations Based on Polarizable Models Show that Ion Permeation Interconverts between Different Mechanisms as a Function of Membrane Thickness. <i>Journal of Physical Chemistry B</i> , 2021 , 125, 1020-1035	3.4	3
4	Vector positronium states in three-dimensional QED. <i>Physical Review D</i> , 1997 , 55, 4954-4966	4.9	2
3	Chapter 15 Charged Protein Side Chain Movement in Lipid Bilayers Explored with Free Energy Simulation. <i>Current Topics in Membranes</i> , 2008 , 405-459	2.2	2
2	In Honour of Professor Cristobal dos Remedios on behalf of the MAWA Trust. <i>Biophysical Reviews</i> , 2020 , 12, 765-766	3.7	0
1	Membrane protein structure and function. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2012 , 1818, 125	3.8	