

Callum J Dickson

List of Publications by Year in Descending Order

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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.

14
papers

1,276
citations

13
h-index

17
g-index

17
ext. papers

1,620
ext. citations

5.8
avg, IF

4.45
L-index

#	Paper	IF	Citations
14	Inhibitor binding influences the protonation states of histidines in SARS-CoV-2 main protease.. <i>Chemical Science</i> , 2021 , 12, 1513-1527	9.4	29
13	Revealing Molecular Determinants of hERG Blocker and Activator Binding. <i>Journal of Chemical Information and Modeling</i> , 2020 , 60, 192-203	6.1	20
12	Hidden bias in the DUD-E dataset leads to misleading performance of deep learning in structure-based virtual screening. <i>PLoS ONE</i> , 2019 , 14, e0220113	3.7	68
11	Using Membrane Partitioning Simulations To Predict Permeability of Forty-Nine Drug-Like Molecules. <i>Journal of Chemical Information and Modeling</i> , 2019 , 59, 236-244	6.1	15
10	Calculating Kinetic Rates and Membrane Permeability from Biased Simulations. <i>Journal of Physical Chemistry B</i> , 2018 , 122, 11571-11578	3.4	15
9	Structure-Kinetic Relationships of Passive Membrane Permeation from Multiscale Modeling. <i>Journal of the American Chemical Society</i> , 2017 , 139, 442-452	16.4	78
8	Contributions of the membrane dipole potential to the function of voltage-gated cation channels and modulation by small molecule potentiators. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2017 , 1859, 177-194	3.8	18
7	Uncoupling the Structure-Activity Relationships of α Adrenergic Receptor Ligands from Membrane Binding. <i>Journal of Medicinal Chemistry</i> , 2016 , 59, 5780-9	8.3	17
6	Simulation of lipid bilayer self-assembly using all-atom lipid force fields. <i>Physical Chemistry Chemical Physics</i> , 2016 , 18, 10573-84	3.6	32
5	Imaging plasma membrane phase behaviour in live cells using a thiophene-based molecular rotor. <i>Chemical Communications</i> , 2016 , 52, 13269-13272	5.8	29
4	Imaging phase separation in model lipid membranes through the use of BODIPY based molecular rotors. <i>Physical Chemistry Chemical Physics</i> , 2015 , 17, 18393-402	3.6	66
3	Lipid14: The Amber Lipid Force Field. <i>Journal of Chemical Theory and Computation</i> , 2014 , 10, 865-879	6.4	725
2	GAFFlipid: a General Amber Force Field for the accurate molecular dynamics simulation of phospholipid. <i>Soft Matter</i> , 2012 , 8, 9617	3.6	151
1	Further evaluation of quantum chemical methods for the prediction of non-specific binding of positron emission tomography tracers. <i>Physical Chemistry Chemical Physics</i> , 2011 , 13, 21552-7	3.6	7