

# Amanda Buyan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9075225/publications.pdf>

Version: 2024-02-01

11  
papers

191  
citations

1478505

6  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

334  
citing authors

#	ARTICLE	IF	CITATIONS
1	Piezo1 Forms Specific, Functionally Important Interactions with Phosphoinositides and Cholesterol. <i>Biophysical Journal</i> , 2020, 119, 1683-1697.	0.5	60
2	Protonation state of inhibitors determines interaction sites within voltage-gated sodium channels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E3135-E3144.	7.1	38
3	Sidekick for Membrane Simulations: Automated Ensemble Molecular Dynamics Simulations of Transmembrane Helices. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 2165-2175.	5.3	27
4	Primary and Secondary Dimer Interfaces of the Fibroblast Growth Factor Receptor 3 Transmembrane Domain: Characterization via Multiscale Molecular Dynamics Simulations. <i>Biochemistry</i> , 2014, 53, 323-332.	2.5	24
5	Multiscale Simulations Suggest a Mechanism for the Association of the Dok7 PH Domain with PIP-Containing Membranes. <i>PLoS Computational Biology</i> , 2016, 12, e1005028.	3.2	24
6	ReplicOpter: A replicate optimizer for flexible docking. <i>Proteins: Structure, Function and Bioinformatics</i> , 2010, 78, 3156-3165.	2.6	6
7	Voltage-Gated Sodium Channel Pharmacology. <i>Advances in Pharmacology</i> , 2017, 79, 255-285.	2.0	4
8	Differences in local anaesthetic and antiepileptic binding in the inactivated state of human sodium channel Nav1.4. <i>Biophysical Journal</i> , 2021, 120, 5553-5563.	0.5	4
9	Computational studies of Piezo1 yield insights into key lipid-protein interactions, channel activation, and agonist binding. <i>Biophysical Reviews</i> , 2022, 14, 209-219.	3.2	3
10	In pursuit of a selective hepatocellular carcinoma therapeutic agent: Novel thalidomide derivatives with antiproliferative, antimigratory and STAT3 inhibitory properties. <i>European Journal of Medicinal Chemistry</i> , 2021, 217, 113353.	5.5	1
11	Initiating Coarse-Grained MD Simulations for Membrane-Bound Proteins. <i>Methods in Molecular Biology</i> , 2022, 2402, 131-141.	0.9	0