

# Hossein Mohammadiarani

## List of Publications by Year in descending order

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Version: 2024-02-01

12  
papers

128  
citations

1307594

7  
h-index

1474206

9  
g-index

12  
all docs

12  
docs citations

12  
times ranked

211  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular Dynamics Simulation to Uncover the Mechanisms of Protein Instability During Freezing. <i>Journal of Pharmaceutical Sciences</i> , 2021, 110, 2457-2471.	3.3	22
2	Artificial neural networks in tandem with molecular descriptors as predictive tools for continuous liposome manufacturing. <i>International Journal of Pharmaceutics</i> , 2021, 603, 120713.	5.2	16
3	Liposome-based measurement of light-driven chloride transport kinetics of halorhodopsin. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2021, 1863, 183637.	2.6	4
4	Cover Image, Volume 87, Issue 2. <i>Proteins: Structure, Function and Bioinformatics</i> , 2019, 87, C1.	2.6	0
5	Interplay of cysteine exposure and global protein dynamics in small-molecule recognition by a regulator of G-protein signaling protein. <i>Proteins: Structure, Function and Bioinformatics</i> , 2019, 87, 146-156.	2.6	13
6	Differential Protein Dynamics of Regulators of G-Protein Signaling: Role in Specificity of Small-Molecule Inhibitors. <i>Journal of the American Chemical Society</i> , 2018, 140, 3454-3460.	13.7	21
7	Measuring Transport Kinetics of Light Driven Membrane Protein, Halorhodopsin. <i>Biophysical Journal</i> , 2018, 114, 146a.	0.5	1
8	Interpreting Hydrogen-Deuterium Exchange Events in Proteins Using Atomistic Simulations: Case Studies on Regulators of G-Protein Signaling Proteins. <i>Journal of Physical Chemistry B</i> , 2018, 122, 9314-9323.	2.6	30
9	Insulin mimetic peptide S371 folds into a helical structure. <i>Journal of Computational Chemistry</i> , 2017, 38, 1158-1166.	3.3	9
10	Conformational Evolution of Three Regulator of G-Protein Signaling Proteins (RGS4, RGS8, RGS19) in Microsecond-Scale Simulations. <i>Biophysical Journal</i> , 2017, 112, 353a-354a.	0.5	0
11	All-Atom Structural Models of the Transmembrane Domains of Insulin and Type 1 Insulin-Like Growth Factor Receptors. <i>Frontiers in Endocrinology</i> , 2016, 7, 68.	3.5	12
12	All-Atom Structural Models of the Transmembrane Domains of Insulin Receptor and Type-1 Insulin-Like Growth Factor Receptor. <i>Biophysical Journal</i> , 2016, 110, 58a.	0.5	0