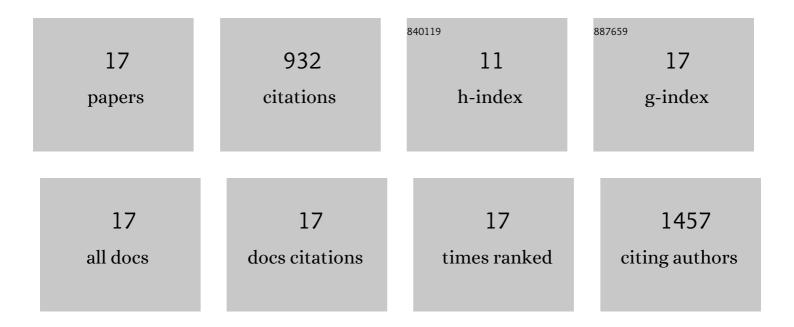
Dhruva K Chakravorty

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Metal Ion Capture Mechanism of a Copper Metallochaperone. Biochemistry, 2016, 55, 501-509.	1.2	9
2	A combined cheminformatic and bioinformatic approach to address the proteolytic stability challenge in peptideâ€based drug discovery. Biopolymers, 2015, 104, 775-789.	1.2	2
3	Role of Substrate Dynamics in Protein Prenylation Reactions. Accounts of Chemical Research, 2015, 48, 439-448.	7.6	11
4	Models for the Metal Transfer Complex of the N-Terminal Region of CusB and CusF. Biochemistry, 2015, 54, 4226-4235.	1.2	10
5	Studying Allosteric Regulation in Metal Sensor Proteins Using Computational Methods. Advances in Protein Chemistry and Structural Biology, 2014, 96, 181-218.	1.0	9
6	Characterizing Solution Surface Loop Conformational Flexibility of the GM2 Activator Protein. Journal of Physical Chemistry B, 2014, 118, 10607-10617.	1.2	12
7	Solution NMR refinement of a metal ion bound protein using metal ion inclusive restrained molecular dynamics methods. Journal of Biomolecular NMR, 2013, 56, 125-137.	1.6	22
8	Structure and Dynamics of the N-Terminal Domain of the Cu(I) Binding Protein CusB. Biochemistry, 2013, 52, 6911-6923.	1.2	26
9	Rational Design of Particle Mesh Ewald Compatible Lennard-Jones Parameters for +2 Metal Cations in Explicit Solvent. Journal of Chemical Theory and Computation, 2013, 9, 2733-2748.	2.3	559
10	Energetics of Zinc-Mediated Interactions in the Allosteric Pathways of Metal Sensor Proteins. Journal of the American Chemical Society, 2013, 135, 30-33.	6.6	24
11	Solution Structure of <i>Mycobacterium tuberculosis</i> NmtR in the Apo State: Insights into Ni(II)-Mediated Allostery. Biochemistry, 2012, 51, 2619-2629.	1.2	42
12	Simulations of Allosteric Motions in the Zinc Sensor CzrA. Journal of the American Chemical Society, 2012, 134, 3367-3376.	6.6	42
13	Insight into the Cationâ~'í€ Interaction at the Metal Binding Site of the Copper Metallochaperone CusF. Journal of the American Chemical Society, 2011, 133, 19330-19333.	6.6	53
14	Finding a Needle in the Haystack: Computational Modeling of Mg ²⁺ Binding in the Active Site of Protein Farnesyltransferase. Biochemistry, 2010, 49, 9658-9666.	1.2	27
15	Impact of Mutation on Proton Transfer Reactions in Ketosteroid Isomerase: Insights from Molecular Dynamics Simulations. Journal of the American Chemical Society, 2010, 132, 7549-7555.	6.6	32
16	Hybrid Quantum/Classical Molecular Dynamics Simulations of the Proton Transfer Reactions Catalyzed by Ketosteroid Isomerase: Analysis of Hydrogen Bonding, Conformational Motions, and Electrostatics. Biochemistry, 2009, 48, 10608-10619.	1.2	37
17	Implementation of Umbrella Integration within the Framework of the Empirical Valence Bond Approach. Journal of Chemical Theory and Computation, 2008, 4, 1974-1980.	2.3	15