

# Soma Ghosh

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5616199/publications.pdf>

Version: 2024-02-01

12  
papers

326  
citations

933447

10  
h-index

1281871

11  
g-index

12  
all docs

12  
docs citations

12  
times ranked

540  
citing authors

#	ARTICLE	IF	CITATIONS
1	Beyond Paralogs: The Multiple Layers of Redundancy in Bacterial Pathogenesis. <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 467.	3.9	84
2	Combinatorial selection in amoebal hosts drives the evolution of the human pathogen <i>Legionella pneumophila</i> . <i>Nature Microbiology</i> , 2020, 5, 599-609.	13.3	78
3	Protein Structure and Function: Looking through the Network of Side-Chain Interactions. <i>Current Protein and Peptide Science</i> , 2015, 17, 4-25.	1.4	51
4	Network properties of decoys and CASP predicted models: a comparison with native protein structures. <i>Molecular BioSystems</i> , 2013, 9, 1774.	2.9	21
5	An analysis and evaluation of the WeFold collaborative for protein structure prediction and its pipelines in CASP11 and CASP12. <i>Scientific Reports</i> , 2018, 8, 9939.	3.3	19
6	A multi-level multi-scale approach to study essential genes in <i>Mycobacterium tuberculosis</i> . <i>BMC Systems Biology</i> , 2013, 7, 132.	3.0	17
7	Iron Limitation Triggers Early Egress by the Intracellular Bacterial Pathogen <i>Legionella pneumophila</i> . <i>Infection and Immunity</i> , 2016, 84, 2185-2197.	2.2	17
8	Rule-based modelling of iron homeostasis in tuberculosis. <i>Molecular BioSystems</i> , 2011, 7, 2750.	2.9	13
9	Mechanism of Iron-Dependent Repressor (IdeR) Activation and DNA Binding: A Molecular Dynamics and Protein Structure Network Study. <i>PLoS Computational Biology</i> , 2015, 11, e1004500.	3.2	13
10	Ranking the quality of protein structure models using sidechain based network properties. <i>F1000Research</i> , 2014, 3, 17.	1.6	11
11	Weighting schemes in metabolic graphs for identifying biochemical routes. <i>Systems and Synthetic Biology</i> , 2014, 8, 47-57.	1.0	2
12	Systems Approaches to Study Infectious Diseases. , 2015, , 151-172.		0