Gayatri Ramakrishnan

List of Publications by Year
in descending order

10	107	1477746 6	1473754 9
papers	citations	h-index	g-index

10	10	10	162
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	In Silico Modeling of FDA-Approved Drugs for Discovery of Anticandida Agents: A Drug-Repurposing Approach. , 2019, , 463-526.		8
2	Repurposing Drugs Based on Evolutionary Relationships Between Targets of Approved Drugs and Proteins of Interest. Methods in Molecular Biology, 2019, 1903, 45-59.	0.4	4
3	Comparison of Leptospira interrogans and Leptospira biflexa genomes: analysis of potential leptospiralâ€“host interactions. Molecular BioSystems, 2017, 13, 883-891.	2.9	6
4	Exploring anti-malarial potential of FDA approved drugs: an in silico approach. Malaria Journal, 2017, 16, 290.	0.8	16
5	Computational recognition and analysis of hitherto uncharacterized nucleotide cyclase-like proteins in bacteria. Biology Direct, 2016, 11, 27.	1.9	3
6	Homology-Based Prediction of Potential Protein-Protein Interactions between Human Erythrocytes and Plasmodium falciparum. Bioinformatics and Biology Insights, 2015, 9, BBI.S31880.	1.0	21
7	SInCReâ€”structural interactome computational resource forMycobacterium tuberculosis. Database: the Journal of Biological Databases and Curation, 2015, 2015, bav060.	1.4	10
8	Enriching the annotation of Mycobacterium tuberculosis H37Rv proteome using remote homology detection approaches: Insights into structure and function. Tuberculosis, 2015, 95, 14-25.	0.8	9
9	Recognizing drug targets using evolutionary information: implications for repurposing FDA-approved drugs against Mycobacterium tuberculosis H37Rv. Molecular BioSystems, 2015, 11, 3316-3331.	2.9	20
10	From workstations to workbenches: Towards predicting physicochemically viable proteinâ€“protein interactions across a host and a pathogen. IUBMB Life, 2014, 66, 759-774.	1.5	10