

Malvika Sharan

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

866
citations

933447

10
h-index

1125743

13
g-index

17
all docs

17
docs citations

17
times ranked

1568
citing authors

#	ARTICLE	IF	CITATIONS
1	An expanded evaluation of protein function prediction methods shows an improvement in accuracy. <i>Genome Biology</i> , 2016, 17, 184.	8.8	308
2	RNA target profiles direct the discovery of virulence functions for the cold-shock proteins CspC and CspE. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 6824-6829.	7.1	110
3	<i>Chlamydia</i> preserves the mitochondrial network necessary for replication via microRNA-dependent inhibition of fission. <i>Journal of Cell Biology</i> , 2017, 216, 1071-1089.	5.2	102
4	Functional high-throughput screening identifies the miR-15 microRNA family as cellular restriction factors for <i>Salmonella</i> infection. <i>Nature Communications</i> , 2014, 5, 4718.	12.8	92
5	Ten Simple Rules for Organizing an Unconference. <i>PLoS Computational Biology</i> , 2015, 11, e1003905.	3.2	69
6	Cell differentiation defines acute and chronic infection cell types in <i>Staphylococcus aureus</i> . <i>ELife</i> , 2017, 6, .	6.0	59
7	A systematic analysis of the RNA-targeting potential of secreted bacterial effector proteins. <i>Scientific Reports</i> , 2017, 7, 9328.	3.3	25
8	Functional screenings reveal different requirements for host microRNAs in <i>Salmonella</i> and <i>Shigella</i> infection. <i>Nature Microbiology</i> , 2020, 5, 192-205.	13.3	25
9	APRICOT: an integrated computational pipeline for the sequence-based identification and characterization of RNA-binding proteins. <i>Nucleic Acids Research</i> , 2017, 45, e96-e96.	14.5	22
10	Analysis of host microRNA function uncovers a role for miR-29b-2-5p in <i>Shigella</i> capture by filopodia. <i>PLoS Pathogens</i> , 2017, 13, e1006327.	4.7	20
11	Centering inclusivity in the design of online conferences—An OHBM—Open Science perspective. <i>GigaScience</i> , 2021, 10, .	6.4	14
12	morFeus: a web-based program to detect remotely conserved orthologs using symmetrical best hits and orthology network scoring. <i>BMC Bioinformatics</i> , 2014, 15, 263.	2.6	9
13	Ten simple rules for establishing a mentorship programme. <i>PLoS Computational Biology</i> , 2022, 18, e1010015.	3.2	4