

# Nikhil Ram-Mohan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6109549/publications.pdf>

Version: 2024-02-01

17  
papers

309  
citations

1163117  
8  
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1125743  
13  
g-index

27  
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docs citations

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times ranked

495  
citing authors

#	ARTICLE	IF	CITATIONS
1	SARS-CoV-2 RNAemia Predicts Clinical Deterioration and Extrapulmonary Complications from COVID-19. <i>Clinical Infectious Diseases</i> , 2022, 74, 218-226.	5.8	51
2	Association Between SARS-CoV-2 RNAemia and Postacute Sequelae of COVID-19. <i>Open Forum Infectious Diseases</i> , 2022, 9, ofab646.	0.9	14
3	Diagnosis of Bloodstream Infections: An Evolution of Technologies towards Accurate and Rapid Identification and Antibiotic Susceptibility Testing. <i>Antibiotics</i> , 2022, 11, 511.	3.7	16
4	Profiling chromatin accessibility responses in human neutrophils with sensitive pathogen detection. <i>Life Science Alliance</i> , 2021, 4, e202000976.	2.8	5
5	A Novel Platform Using RNA Signatures To Accelerate Antimicrobial Susceptibility Testing in <i>Neisseria gonorrhoeae</i> . <i>Journal of Clinical Microbiology</i> , 2020, 58, .	3.9	8
6	Comparative Metatranscriptomics of Periodontitis Supports a Common Polymicrobial Shift in Metabolic Function and Identifies Novel Putative Disease-Associated ncRNAs. <i>Frontiers in Microbiology</i> , 2020, 11, 482.	3.5	16
7	Insights into gene expression changes under conditions that facilitate horizontal gene transfer (mating) of a model archaeon. <i>Scientific Reports</i> , 2020, 10, 22297.	3.3	8
8	Regulatory context drives conservation of glycine riboswitch aptamers. <i>PLoS Computational Biology</i> , 2019, 15, e1007564.	3.2	6
9	Regulatory context drives conservation of glycine riboswitch aptamers. , 2019, 15, e1007564.		0
10	Regulatory context drives conservation of glycine riboswitch aptamers. , 2019, 15, e1007564.		0
11	Regulatory context drives conservation of glycine riboswitch aptamers. , 2019, 15, e1007564.		0
12	Regulatory context drives conservation of glycine riboswitch aptamers. , 2019, 15, e1007564.		0
13	The Transcriptional landscape of <i>Streptococcus pneumoniae</i> TIGR4 reveals a complex operon architecture and abundant riboregulation critical for growth and virulence. <i>PLoS Pathogens</i> , 2018, 14, e1007461.	4.7	37
14	Analysis of the bacteriorhodopsin-producing haloarchaea reveals a core community that is stable over time in the salt crystallizers of Eilat, Israel. <i>Extremophiles</i> , 2016, 20, 747-757.	2.3	8
15	Horizontal Gene Transfer, Dispersal and Haloarchaeal Speciation. <i>Life</i> , 2015, 5, 1405-1426.	2.4	28
16	Evidence from phylogenetic and genome fingerprinting analyses suggests rapidly changing variation in <i>Haloarubrum</i> and <i>Haloarcula</i> populations. <i>Frontiers in Microbiology</i> , 2014, 5, 143.	3.5	25
17	Cell sorting analysis of geographically separated hypersaline environments. <i>Extremophiles</i> , 2013, 17, 265-275.	2.3	46