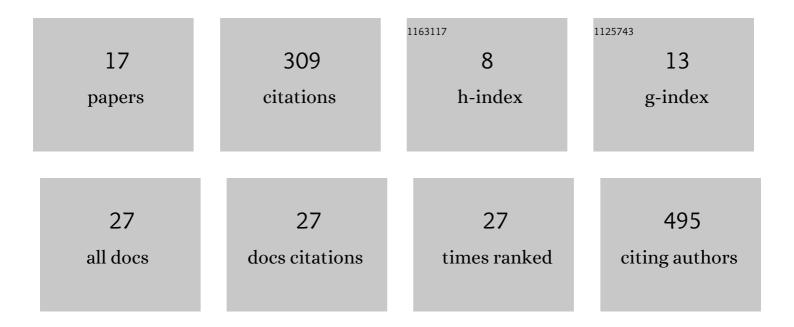
## Nikhil Ram-Mohan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6109549/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	SARS-CoV-2 RNAemia Predicts Clinical Deterioration and Extrapulmonary Complications from COVID-19. Clinical Infectious Diseases, 2022, 74, 218-226.	5.8	51
2	Association Between SARS-CoV-2 RNAemia and Postacute Sequelae of COVID-19. Open Forum Infectious Diseases, 2022, 9, ofab646.	0.9	14
3	Diagnosis of Bloodstream Infections: An Evolution of Technologies towards Accurate and Rapid Identification and Antibiotic Susceptibility Testing. Antibiotics, 2022, 11, 511.	3.7	16
4	Profiling chromatin accessibility responses in human neutrophils with sensitive pathogen detection. Life Science Alliance, 2021, 4, e202000976.	2.8	5
5	A Novel Platform Using RNA Signatures To Accelerate Antimicrobial Susceptibility Testing in Neisseria gonorrhoeae. Journal of Clinical Microbiology, 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. Frontiers in Microbiology, 2020, 11, 482.	3.5	16
7	Insights into gene expression changes under conditions that facilitate horizontal gene transfer (mating) of a model archaeon. Scientific Reports, 2020, 10, 22297.	3.3	8
8	Regulatory context drives conservation of glycine riboswitch aptamers. PLoS Computational Biology, 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 Streptococcus pneumoniae TIGR4 reveals a complex operon architecture and abundant riboregulation critical for growth and virulence. PLoS Pathogens, 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. Extremophiles, 2016, 20, 747-757.	2.3	8
15	Horizontal Gene Transfer, Dispersal and Haloarchaeal Speciation. Life, 2015, 5, 1405-1426.	2.4	28
16	Evidence from phylogenetic and genome fingerprinting analyses suggests rapidly changing variation in Halorubrum and Haloarcula populations. Frontiers in Microbiology, 2014, 5, 143.	3.5	25
17	Cell sorting analysis of geographically separated hypersaline environments. Extremophiles, 2013, 17, 265-275.	2.3	46