## Gaurav Sharma

List of Publications by Year in descending order

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933447 839539 21 396 10 18 citations h-index g-index papers 26 26 26 516 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Harmonizing Prokaryotic Nomenclature: Fixing the Fuss over Phylum Name Flipping. MBio, 2022, 13, e0097022.	4.1	11
2	The lag in SARS-CoV-2 genome submissions to GISAID. Nature Biotechnology, 2021, 39, 1058-1060.	17.5	47
3	Complete genome sequence and identification of polyunsaturated fatty acid biosynthesis genes of the myxobacterium Minicystis rosea DSM 24000T. BMC Genomics, 2021, 22, 655.	2.8	6
4	Global gene expression analysis of the Myxococcus xanthus developmental time course. Genomics, 2021, 113, 120-134.	2.9	14
5	Dynamic tracking of variant frequencies depicts the evolution of mutation sites amongst SARSâ€CoVâ€⊋ genomes from India. Journal of Medical Virology, 2021, 93, 2534-2537.	5.0	10
6	Peripheral rods: a specialized developmental cell type in Myxococcus xanthus. Genomics, 2020, 112, 1588-1597.	2.9	6
7	Draft Genome Sequence of Marinobacter hydrocarbonoclasticus NCT7M, Obtained from the Microbiome of a Marine Copepod. Microbiology Resource Announcements, 2020, 9, .	0.6	O
8	Modulation of bacterial multicellularity via spatio-specific polysaccharide secretion. PLoS Biology, 2020, 18, e3000728.	5.6	37
9	Understanding genomic diversity, pan-genome, and evolution of SARS-CoV-2. PeerJ, 2020, 8, e9576.	2.0	19
10	Draft Genome Sequence of the Caffeine-Degrading Methylotroph Methylorubrum populi Pinkel. Microbiology Resource Announcements, 2019, 8, .	0.6	1
11	Composite genome sequence of Bacillus clausii, a probiotic commercially available as Enterogermina $\hat{A}^{@}$ , and insights into its probiotic properties. BMC Microbiology, 2019, 19, 307.	3.3	41
12	Draft Genome Sequence of Vibrio cyclitrophicus NCT10V, Cultivated from the Microbiome of a Marine Copepod. Microbiology Resource Announcements, 2019, 8, .	0.6	3
13	Comparative Genomics of Myxobacterial Chemosensory Systems. Journal of Bacteriology, 2018, 200, .	2.2	18
14	In silico characterization of a novel putative aerotaxis chemosensory system in the myxobacterium, Corallococcus coralloides. BMC Genomics, 2018, 19, 757.	2.8	3
15	Diversity and Evolution of Myxobacterial Type IV Pilus Systems. Frontiers in Microbiology, 2018, 9, 1630.	3.5	8
16	Unravelling the Complete Genome of Archangium gephyra DSM 2261T and Evolutionary Insights into Myxobacterial Chitinases. Genome Biology and Evolution, 2017, 9, 1304-1311.	2.5	43
17	Complete Genome of the Starch-Degrading Myxobacteria <i>Sandaracinus amylolyticus</i> DSM 53668 <sup>T</sup> . Genome Biology and Evolution, 2016, 8, 2520-2529.	2.5	63
18	Complete Genome Sequence and Comparative Genomics of a Novel Myxobacterium Myxococcus hansupus. PLoS ONE, 2016, 11, e0148593.	2.5	33

#	Article	IF	CITATIONS
19	Draft Genome Sequence of Kocuria palustris PEL. Genome Announcements, 2014, 2, .	0.8	4
20	In silico analysis of ubiquitin/ubiquitin-like modifiers and their conjugating enzymes in Entamoeba species. Parasitology Research, $2012, 111, 37-51$ .	1.6	18
21	Integrin-Like Tethering of Motility Complexes at Bacterial Focal Adhesions. SSRN Electronic Journal, 0,	0.4	3