

Rikky W Purbojati

List of Publications by Year in descending order

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1040056

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47
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47
docs citations

47
times ranked

1440
citing authors

#	ARTICLE	IF	CITATIONS
1	Vertical stratification of the air microbiome in the lower troposphere. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	13
2	The genomic characterisation and comparison of <i>Bacillus cereus</i> strains isolated from indoor air. Gut Pathogens, 2021, 13, 6.	3.4	9
3	Experimental parameters defining ultra-low biomass bioaerosol analysis. Npj Biofilms and Microbiomes, 2021, 7, 37.	6.4	27
4	A compromised developmental trajectory of the infant gut microbiome and metabolome in atopic eczema. Gut Microbes, 2020, 12, 1801964.	9.8	51
5	Taxonomic composition and seasonal dynamics of the air microbiome in West Siberia. Scientific Reports, 2020, 10, 21515.	3.3	10
6	Longitudinal assessment of antibiotic resistance gene profiles in gut microbiomes of infants at risk of eczema. BMC Infectious Diseases, 2020, 20, 312.	2.9	11
7	Whole-Genome Sequencing of <i>Aspergillus terreus</i> Species Complex. Mycopathologia, 2020, 185, 405-408.	3.1	7
8	Complete genome of <i>Enterobacter sichuanensis</i> strain SGAir0282 isolated from air in Singapore. Gut Pathogens, 2020, 12, 12.	3.4	1
9	Metagenomics Reveals a Core Macrolide Resistome Related to Microbiota in Chronic Respiratory Disease. American Journal of Respiratory and Critical Care Medicine, 2020, 202, 433-447.	5.6	58
10	Complete Genome Sequence of <i>Penicillium oxalicum</i> Strain SGAir0226 Isolated from Outdoor Tropical Air in Singapore. Mycopathologia, 2020, 185, 591-594.	3.1	3
11	Complete Genome Sequence of <i>Citricoccus</i> sp. Strain SGAir0253, Isolated from Indoor Air in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.6	1
12	Complete Genome Sequence of <i>Microbacterium</i> sp. Strain SGAir0570, Isolated from Tropical Air Collected in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.6	2
13	Genome Sequence of the Tropical Atmosphere Bacterium <i>Pontibacter</i> sp. Strain SGAir0037. Microbiology Resource Announcements, 2019, 8, .	0.6	0
14	Complete Genome Sequence of <i>Agrococcus</i> sp. Strain SGAir0287, Isolated from Tropical Air Collected in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.6	0
15	Complete Genome Sequence of <i>Micrococcus luteus</i> Strain SGAir0127, Isolated from Indoor Air Samples from Singapore. Microbiology Resource Announcements, 2019, 8, .	0.6	7
16	Whole-Genome Sequence of <i>Bacillus megaterium</i> Strain SGAir0080, Isolated from an Indoor Air Sample. Microbiology Resource Announcements, 2019, 8, .	0.6	3
17	Complete Genome Sequence of <i>Curtobacterium</i> sp. Strain SGAir0471, Isolated from Singapore Air Samples. Microbiology Resource Announcements, 2019, 8, .	0.6	0
18	Neurogenesis and longevity signaling in young germ-free mice transplanted with the gut microbiota of old mice. Science Translational Medicine, 2019, 11, .	12.4	122

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19	Complete Genome Sequence of <i>Brachybacterium</i> sp. Strain SGAir0954, Isolated from Singapore Air. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	4
20	Microbial communities in the tropical air ecosystem follow a precise diel cycle. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 23299-23308.	7.1	99
21	Complete Genome Sequence of <i>Lysinibacillus</i> sp. Strain SGAir0095, Isolated from Tropical Air Samples Collected in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	2
22	Complete Genome Sequence of <i>Streptomyces</i> sp. Strain SGAir0924, an Actinobacterium Isolated from Outdoor Air in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	2
23	Complete Genome Sequence of <i>Nissabacter</i> sp. Strain SGAir0207, Isolated from an Air Sample Collected in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	1
24	Complete Genome Sequence of <i>Pseudomonas</i> sp. Strain SGAir0191, Isolated from Tropical Air Collected in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	0
25	Complete Genome Sequence of <i>Brevundimonas</i> sp. Strain SGAir0440, Isolated from Indoor Air in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	0
26	Complete Genome Sequence of <i>Enterococcus faecalis</i> Strain SGAir0397, Isolated from a Tropical Air Sample Collected in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	0
27	Complete Genome Sequence of <i>Rhodococcus</i> sp. Strain SGAir0479, Isolated from Indoor Air Collected in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	1
28	Complete Genome Sequence of <i>Acinetobacter indicus</i> Type Strain SGAir0564 Isolated from Tropical Air Collected in Singapore. <i>Genome Announcements</i> , 2018, 6, .	0.8	2
29	Genome Sequence of <i>Bacillus velezensis</i> SGAir0473, Isolated from Tropical Air Collected in Singapore. <i>Genome Announcements</i> , 2018, 6, .	0.8	7
30	Complete Genome Sequence of <i>Lelliottia nimipressuralis</i> Type Strain SGAir0187, Isolated from Tropical Air Collected in Singapore. <i>Genome Announcements</i> , 2018, 6, .	0.8	5
31	Complete Genome Sequence of the Bacterium <i>Serratia marcescens</i> SGAir0764, Isolated from Singapore Air. <i>Genome Announcements</i> , 2018, 6, .	0.8	1
32	Complete Genome Sequence of <i>Pseudomonas stutzeri</i> Type Strain SGAir0442, Isolated from Singapore Air Samples. <i>Genome Announcements</i> , 2018, 6, .	0.8	1
33	Complete Genome Sequence of <i>Acinetobacter schindleri</i> SGAir0122 Isolated from Singapore Air. <i>Genome Announcements</i> , 2018, 6, .	0.8	2
34	Complete Genome Sequence of <i>Staphylococcus haemolyticus</i> Type Strain SGAir0252. <i>Genome Announcements</i> , 2018, 6, .	0.8	4
35	Genome Sequence of <i>Geobacillus thermoleovorans</i> SGAir0734, Isolated from Singapore Air. <i>Genome Announcements</i> , 2018, 6, .	0.8	4
36	Genome Sequence of <i>Pantoea ananatis</i> SGAir0210, Isolated from Outdoor Air in Singapore. <i>Genome Announcements</i> , 2018, 6, .	0.8	2

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37	Long-read sequencing uncovers the adaptive topography of a carnivorous plant genome. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E4435-E4441.	7.1	95
38	The microbiomes of blowflies and houseflies as bacterial transmission reservoirs. Scientific Reports, 2017, 7, 16324.	3.3	115
39	Complete Genome Sequence of Bacillus altitudinis Type Strain SGAir0031 Isolated from Tropical Air Collected in Singapore. Genome Announcements, 2017, 5, .	0.8	9
40	Genome Sequence of Prosthecochloris sp. Strain HL-130-GSB, from the Phylum Chlorobi. Genome Announcements, 2017, 5, .	0.8	3
41	Large-scale mitogenomics enables insights into Schizophora (Diptera) radiation and population diversity. Scientific Reports, 2016, 6, 21762.	3.3	66
42	In silico analyses of metagenomes from human atherosclerotic plaque samples. Microbiome, 2015, 3, 38.	11.1	87