

Ana Carolina M Junqueira

List of Publications by Year in descending order

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49
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49
docs citations

49
times ranked

1228
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, .	3.3	13
2	The genomic characterisation and comparison of <i>Bacillus cereus</i> strains isolated from indoor air. Gut Pathogens, 2021, 13, 6.	1.6	9
3	Experimental parameters defining ultra-low biomass bioaerosol analysis. Npj Biofilms and Microbiomes, 2021, 7, 37.	2.9	27
4	Disruption of the odorant coreceptor Orco impairs foraging and host finding behaviors in the New World screwworm fly. Scientific Reports, 2021, 11, 11379.	1.6	7
5	Complete genome of <i>Enterobacter sichuanensis</i> strain SGAir0282 isolated from air in Singapore. Gut Pathogens, 2020, 12, 12.	1.6	1
6	Complete Genome Sequence of <i>Citricoccus</i> sp. Strain SGAir0253, Isolated from Indoor Air in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	1
7	Complete Genome Sequence of <i>Microbacterium</i> sp. Strain SGAir0570, Isolated from Tropical Air Collected in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	2
8	Genome Sequence of the Tropical Atmosphere Bacterium <i>Pontibacter</i> sp. Strain SGAir0037. Microbiology Resource Announcements, 2019, 8, .	0.3	0
9	Complete Genome Sequence of <i>Agrococcus</i> sp. Strain SGAir0287, Isolated from Tropical Air Collected in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	0
10	Complete Genome Sequence of <i>Micrococcus luteus</i> Strain SGAir0127, Isolated from Indoor Air Samples from Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	7
11	Whole-Genome Sequence of <i>Bacillus megaterium</i> Strain SGAir0080, Isolated from an Indoor Air Sample. Microbiology Resource Announcements, 2019, 8, .	0.3	3
12	Complete Genome Sequence of <i>Curtobacterium</i> sp. Strain SGAir0471, Isolated from Singapore Air Samples. Microbiology Resource Announcements, 2019, 8, .	0.3	0
13	Complete Genome Sequence of <i>Brachybacterium</i> sp. Strain SGAir0954, Isolated from Singapore Air. Microbiology Resource Announcements, 2019, 8, .	0.3	4
14	Microbial communities in the tropical air ecosystem follow a precise diel cycle. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 23299-23308.	3.3	99
15	Complete Genome Sequence of <i>Lysinibacillus</i> sp. Strain SGAir0095, Isolated from Tropical Air Samples Collected in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	2
16	Complete Genome Sequence of <i>Streptomyces</i> sp. Strain SGAir0924, an Actinobacterium Isolated from Outdoor Air in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	2
17	Complete Genome Sequence of <i>Nissabacter</i> sp. Strain SGAir0207, Isolated from an Air Sample Collected in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	1
18	Complete Genome Sequence of <i>Pseudomonas</i> sp. Strain SGAir0191, Isolated from Tropical Air Collected in Singapore. Microbiology Resource Announcements, 2019, 8, .	0.3	0

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19	Complete Genome Sequence of <i>Brevundimonas</i> sp. Strain SGAir0440, Isolated from Indoor Air in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	0
20	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, .	0.3	0
21	Complete Genome Sequence of <i>Rhodococcus</i> sp. Strain SGAir0479, Isolated from Indoor Air Collected in Singapore. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	1
22	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
23	Genome Sequence of <i>Bacillus velezensis</i> SGAir0473, Isolated from Tropical Air Collected in Singapore. <i>Genome Announcements</i> , 2018, 6, .	0.8	7
24	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
25	Complete Genome Sequence of the Bacterium <i>Serratia marcescens</i> SGAir0764, Isolated from Singapore Air. <i>Genome Announcements</i> , 2018, 6, .	0.8	1
26	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
27	Complete Genome Sequence of <i>Acinetobacter schindleri</i> SGAir0122 Isolated from Singapore Air. <i>Genome Announcements</i> , 2018, 6, .	0.8	2
28	Complete Genome Sequence of <i>Staphylococcus haemolyticus</i> Type Strain SGAir0252. <i>Genome Announcements</i> , 2018, 6, .	0.8	4
29	Genome Sequence of <i>Geobacillus thermoleovorans</i> SGAir0734, Isolated from Singapore Air. <i>Genome Announcements</i> , 2018, 6, .	0.8	4
30	Genome Sequence of <i>Pantoea ananatis</i> SGAir0210, Isolated from Outdoor Air in Singapore. <i>Genome Announcements</i> , 2018, 6, .	0.8	2
31	The microbiomes of blowflies and houseflies as bacterial transmission reservoirs. <i>Scientific Reports</i> , 2017, 7, 16324.	1.6	115
32	Complete Genome Sequence of <i>Bacillus altitudinis</i> Type Strain SGAir0031 Isolated from Tropical Air Collected in Singapore. <i>Genome Announcements</i> , 2017, 5, .	0.8	9
33	Identification and characterization of microRNAs in the screwworm flies <i>Cochliomyia hominivorax</i> and <i>Cochliomyia macellaria</i> (Diptera: Calliphoridae). <i>Insect Molecular Biology</i> , 2017, 26, 46-57.	1.0	4
34	Large-scale mitogenomics enables insights into Schizophora (Diptera) radiation and population diversity. <i>Scientific Reports</i> , 2016, 6, 21762.	1.6	66
35	Molecular phylogenetics of Oestroidea (Diptera: Calyptratae) with emphasis on Calliphoridae: Insights into the inter-familial relationships and additional evidence for paraphyly among blowflies. <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 840-854.	1.2	72
36	Evaluation of the internal transcribed spacer 2 (ITS2) as a molecular marker for phylogenetic inference using sequence and secondary structure information in blow flies (Diptera: Calliphoridae). <i>Genetica</i> , 2011, 139, 1189-1207.	0.5	22

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37	Molecular phylogeny of the <i>Drosophila tripunctata</i> and closely related species groups (Diptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10	1.2	36
38	Structure and evolution of the mitochondrial genomes of <i>Haematobia irritans</i> and <i>Stomoxys calcitrans</i> : The Muscidae (Diptera: Calyptratae) perspective. <i>Molecular Phylogenetics and Evolution</i> , 2008, 48, 850-857.	1.2	52
39	The Mitochondrial Control Region of Blowflies (Diptera: Calliphoridae): A Hot Spot for Mitochondrial Genome Rearrangements. <i>Journal of Medical Entomology</i> , 2008, 45, 667-676.	0.9	24
40	The Mitochondrial Control Region of Blowflies (Diptera: Calliphoridae): A Hot Spot for Mitochondrial Genome Rearrangements. <i>Journal of Medical Entomology</i> , 2008, 45, 667-676.	0.9	12
41	The mitochondrial genome of the blowfly <i>Chrysomya chloropyga</i> (Diptera: Calliphoridae). <i>Gene</i> , 2004, 339, 7-15.	1.0	151
42	Analysis of a conserved duplicated tRNA gene in the mitochondrial genome of blowflies. <i>Gene</i> , 2004, 339, 1-6.	1.0	31
43	Methods for the recovery of mitochondrial DNA sequences from museum specimens of myiasis-causing flies. <i>Medical and Veterinary Entomology</i> , 2002, 16, 39-45.	0.7	36
44	The mitochondrial genome of the primary screwworm fly <i>Cochliomyia hominivorax</i> (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462 T	1.0	129