Rebecca J Rockett

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9254416/publications.pdf

Version: 2024-02-01

933447 642732 1,129 24 10 23 citations g-index h-index papers 34 34 34 2186 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Assessment of Inter-Laboratory Differences in SARS-CoV-2 Consensus Genome Assemblies between Public Health Laboratories in Australia. Viruses, 2022, 14, 185.	3.3	4
2	Genome-wide networks reveal emergence of epidemic strains of Salmonella Enteritidis. International Journal of Infectious Diseases, 2022, 117, 65-73.	3.3	8
3	Resistance Mutations in SARS-CoV-2 Delta Variant after Sotrovimab Use. New England Journal of Medicine, 2022, 386, 1477-1479.	27.0	134
4	Co-infection with SARS-CoV-2 Omicron and Delta variants revealed by genomic surveillance. Nature Communications, 2022, 13, 2745.	12.8	64
5	SARS-CoV-2 Within-Host and in vitro Genomic Variability and Sub-Genomic RNA Levels Indicate Differences in Viral Expression Between Clinical Cohorts and in vitro Culture. Frontiers in Microbiology, 2022, 13, .	3.5	4
6	Microbial Genomics as a Catalyst for Targeted Antivirulence Therapeutics. Frontiers in Medicine, 2021, 8, 641260.	2.6	4
7	Epidemiologic Evidence for Airborne Transmission of SARS-CoV-2 during Church Singing, Australia, 2020. Emerging Infectious Diseases, 2021, 27, 1677-1680.	4.3	97
8	Risk factors leading to COVIDâ€19 cases in a Sydney restaurant. Australian and New Zealand Journal of Public Health, 2021, 45, 512-516.	1.8	3
9	SARS-CoV-2 neutralizing antibodies: Longevity, breadth, and evasion by emerging viral variants. PLoS Medicine, 2021, 18, e1003656.	8.4	109
10	SARS-CoV-2 Genome Sequencing Methods Differ in Their Abilities To Detect Variants from Low-Viral-Load Samples. Journal of Clinical Microbiology, 2021, 59, e0104621.	3.9	33
11	Bondi and beyond. Lessons from three waves of COVID-19 from 2020. Public Health Research and Practice, 2021, 31, .	1.5	10
12	Documenting elimination of co-circulating COVID-19 clusters using genomics in New South Wales, Australia. BMC Research Notes, 2021, 14, 415.	1.4	4
13	Complete Genome Sequences, Derived by Next-Generation Sequencing, of JC Polyomavirus Strains Isolated from Vietnamese Renal Transplant Recipients. Microbiology Resource Announcements, 2020, 9, .	0.6	1
14	Genomic Surveillance Enables Suitability Assessment of <i>Salmonella</i> Culture-Independent Diagnostic Testing. Journal of Clinical Microbiology, 2020, 58, .	3.9	5
15	Revealing COVID-19 transmission in Australia by SARS-CoV-2 genome sequencing and agent-based modeling. Nature Medicine, 2020, 26, 1398-1404.	30.7	283
16	Deep Sequencing of Urine Specimens Detects Two BK Polyomavirus Genotypes in a Hematopoietic Stem Cell Transplant Recipient. Microbiology Resource Announcements, 2020, 9, .	0.6	0
17	An emergent clade of SARS-CoV-2 linked to returned travellers from Iran. Virus Evolution, 2020, 6, veaa027.	4.9	119
18	Optimization of sample preparation for culture-independent sequencing of Bordetella pertussis. Microbial Genomics, 2020, 6, .	2.0	8

#	Article	IF	CITATIONS
19	Complete microbial genomes for public health in Australia and the Southwest Pacific. Microbial Genomics, 2020, 6, .	2.0	10
20	Genome Sequencing Links Persistent Outbreak of Legionellosis in Sydney (New South Wales, Australia) to an Emerging Clone of Legionella pneumophila Sequence Type 211. Applied and Environmental Microbiology, 2018, 84, .	3.1	10
21	Advances in Clinical Sample Preparation for Identification and Characterization of Bacterial Pathogens Using Metagenomics. Frontiers in Public Health, 2018, 6, 363.	2.7	48
22	Genome-wide analysis of Streptococcus pneumoniae serogroup 19 in the decade after the introduction of pneumococcal conjugate vaccines in Australia. Scientific Reports, 2018, 8, 16969.	3.3	14
23	Multidrug-Resistant <i>Salmonellaenterica</i> 4,[5],12:i- Sequence Type 34, New South Wales, Australia, 2016–2017. Emerging Infectious Diseases, 2018, 24, 751-753.	4.3	52
24	Whole Genome Sequencing of Candida glabrata for Detection of Markers of Antifungal Drug Resistance. Journal of Visualized Experiments, 2017, , .	0.3	24