Mathieu Almeida

List of Publications by Year in descending order

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777949 993246 17,704 17 13 17 citations h-index g-index papers 20 20 20 24445 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Three transmission events of Vibrio cholerae O1 into Lusaka, Zambia. BMC Infectious Diseases, 2021, 21, 570.	1.3	5
2	MSPminer: abundance-based reconstitution of microbial pan-genomes from shotgun metagenomic data. Bioinformatics, 2019, 35, 1544-1552.	1.8	82
3	Microbial ecology perturbation in human IgA deficiency. Science Translational Medicine, 2018, 10, .	5.8	206
4	Molecular characterization of Vibrio cholerae responsible for cholera epidemics in Uganda by PCR, MLVA and WGS. PLoS Neglected Tropical Diseases, 2018, 12, e0006492.	1.3	40
5	Genetic relatedness of Vibrio cholerae isolates within and between households during outbreaks in Dhaka, Bangladesh. BMC Genomics, 2017, 18, 903.	1.2	13
6	Minimal genetic change in Vibrio cholerae in Mozambique over time: Multilocus variable number tandem repeat analysis and whole genome sequencing. PLoS Neglected Tropical Diseases, 2017, 11, e0005671.	1.3	31
7	Comparison of inferred relatedness based on multilocus variable-number tandem-repeat analysis and whole genome sequencing of Vibrio choleraeO1. FEMS Microbiology Letters, 2016, 363, fnw116.	0.7	19
8	Capturing the most wanted taxa through cross-sample correlations. ISME Journal, 2016, 10, 2459-2467.	4.4	9
9	High-Throughput Sequencing as a Tool for Exploring the Human Microbiome. , 2015, , 55-66.		3
10	Construction of a dairy microbial genome catalog opens new perspectives for the metagenomic analysis of dairy fermented products. BMC Genomics, 2014, 15, 1101.	1.2	64
11	Identification and assembly of genomes and genetic elements in complex metagenomic samples without using reference genomes. Nature Biotechnology, 2014, 32, 822-828.	9.4	909
12	Richness of human gut microbiome correlates with metabolic markers. Nature, 2013, 500, 541-546.	13.7	3,641
13	Dietary intervention impact on gut microbial gene richness. Nature, 2013, 500, 585-588.	13.7	1,485
14	A metagenome-wide association study of gut microbiota in type 2 diabetes. Nature, 2012, 490, 55-60.	13.7	5,345
15	Enterotypes of the human gut microbiome. Nature, 2011, 473, 174-180.	13.7	5,800
16	Complete Genome Sequence of the Commensal Streptococcus salivarius Strain JIM8777. Journal of Bacteriology, 2011, 193, 5024-5025.	1.0	19
17	Complete Genome Sequence of the Clinical Streptococcus salivarius Strain CCHSS3. Journal of Bacteriology, 2011, 193, 5041-5042.	1.0	9