

Felipe Cabarcas

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12125622/publications.pdf>

Version: 2024-02-01

10
papers

221
citations

1478505

6
h-index

1474206

9
g-index

10
all docs

10
docs citations

10
times ranked

448
citing authors

#	ARTICLE	IF	CITATIONS
1	Inherited p40phox deficiency differs from classic chronic granulomatous disease. <i>Journal of Clinical Investigation</i> , 2018, 128, 3957-3975.	8.2	99
2	Differential determinants of virulence in two <i>Mycobacterium tuberculosis</i> Colombian clinical isolates of the LAM09 family. <i>Virulence</i> , 2019, 10, 695-710.	4.4	36
3	Metagenomic assessment of the microbial community and methanogenic pathways in biosolids from a municipal wastewater treatment plant in Medellín, Colombia. <i>Science of the Total Environment</i> , 2019, 648, 572-581.	8.0	26
4	Annual microbial community dynamics in a full-scale anaerobic sludge digester from a wastewater treatment plant in Colombia. <i>Science of the Total Environment</i> , 2020, 726, 138479.	8.0	23
5	Transcriptome profiling of the cysticercus stage of the laboratory model <i>Taenia crassiceps</i> , strain ORF. <i>Acta Tropica</i> , 2016, 154, 50-62.	2.0	13
6	Dual RNA Sequencing of <i>Mycobacterium tuberculosis</i> -Infected Human Splenic Macrophages Reveals a Strain-Dependent Host-Pathogen Response to Infection. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1803.	4.1	10
7	Comparative genomic analysis of the principal <i>Cryptosporidium</i> species that infect humans. <i>PeerJ</i> , 2020, 8, e10478.	2.0	7
8	<i>Cryptosporidium hominis</i> Phylogenomic Analysis Reveals Separate Lineages With Continental Segregation. <i>Frontiers in Genetics</i> , 2021, 12, 740940.	2.3	3
9	CLAME: a new alignment-based binning algorithm allows the genomic description of a novel Xanthomonadaceae from the Colombian Andes. <i>BMC Genomics</i> , 2018, 19, 858.	2.8	2
10	DATMA: Distributed AuTomatic Metagenomic Assembly and annotation framework. <i>PeerJ</i> , 2020, 8, e9762.	2.0	2