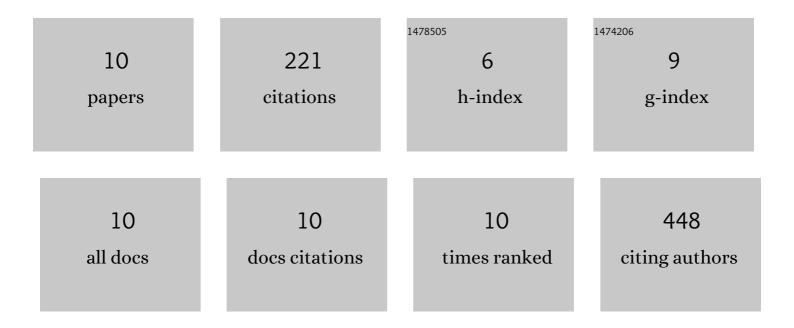
Felipe Cabarcas

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12125622/publications.pdf Version: 2024-02-01



FELIDE CARADCAS

#	Article	IF	CITATIONS
1	Inherited p40phox deficiency differs from classic chronic granulomatous disease. Journal of Clinical Investigation, 2018, 128, 3957-3975.	8.2	99
2	Differential determinants of virulence in two Mycobacterium tuberculosis Colombian clinical isolates of the LAM09 family. Virulence, 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 MedellÃn, Colombia. Science of the Total Environment, 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. Science of the Total Environment, 2020, 726, 138479.	8.0	23
5	Transcriptome profiling of the cysticercus stage of the laboratory model Taenia crassiceps, strain ORF. Acta Tropica, 2016, 154, 50-62.	2.0	13
6	Dual RNA Sequencing of Mycobacterium tuberculosis-Infected Human Splenic Macrophages Reveals a Strain-Dependent Host–Pathogen Response to Infection. International Journal of Molecular Sciences, 2022, 23, 1803.	4.1	10
7	Comparative genomic analysis of the principal <i>Cryptosporidium</i> species that infect humans. PeerJ, 2020, 8, e10478.	2.0	7
8	Cryptosporidium hominis Phylogenomic Analysis Reveals Separate Lineages With Continental Segregation. Frontiers in Genetics, 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. BMC Genomics, 2018, 19, 858.	2.8	2
10	DATMA: Distributed AuTomatic Metagenomic Assembly and annotation framework. PeerJ, 2020, 8, e9762.	2.0	2