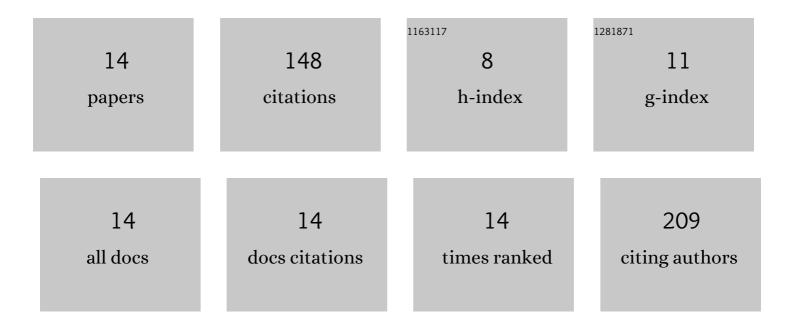
Renata De A B Assis

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3389587/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Identification and analysis of seven effector protein families with different adaptive and evolutionary histories in plant-associated members of the Xanthomonadaceae. Scientific Reports, 2017, 7, 16133.	3.3	35
2	Genome-Wide Profiling and Phylogenetic Analysis of the SWEET Sugar Transporter Gene Family in Walnut and Their Lack of Responsiveness to Xanthomonas arboricola pv. juglandis Infection. International Journal of Molecular Sciences, 2020, 21, 1251.	4.1	17
3	Biotechnological potential of plant growth-promoting bacteria from the roots and rhizospheres of endemic plants in ironstone vegetation in southeastern Brazil. World Journal of Microbiology and Biotechnology, 2018, 34, 156.	3.6	15
4	Analyses of Seven New Genomes of Xanthomonas citri pv. aurantifolii Strains, Causative Agents of Citrus Canker B and C, Show a Reduced Repertoire of Pathogenicity-Related Genes. Frontiers in Microbiology, 2019, 10, 2361.	3.5	14
5	Proteome Analysis of Walnut Bacterial Blight Disease. International Journal of Molecular Sciences, 2020, 21, 7453.	4.1	12
6	Alcaligenes faecalisassociated with Mimosa calodendron rizhosphere assist plant survival in arsenic rich soils. Journal of Soil Science and Plant Nutrition, 2017, 17, 1102-1115.	3.4	11
7	Serratia liquefaciens FG3 isolated from a metallophyte plant sheds light on the evolution and mechanisms of adaptive traits in extreme environments. Scientific Reports, 2019, 9, 18006.	3.3	10
8	Brazilian Ironstone Plant Communities as Reservoirs of Culturable Bacteria With Diverse Biotechnological Potential. Frontiers in Microbiology, 2018, 9, 1638.	3.5	9
9	Complete genome sequence and analysis of Alcaligenes faecalis strain Mc250, a new potential plant bioinoculant. PLoS ONE, 2020, 15, e0241546.	2.5	9
10	Gene Tags Assessment by Comparative Genomics (GTACG): A User-Friendly Framework for Bacterial Comparative Genomics. Frontiers in Genetics, 2019, 10, 725.	2.3	4
11	De Novo Arginine Synthesis Is Required for Full Virulence of <i>Xanthomonas arboricola</i> pv. <i>juglandis</i> During Walnut Bacterial Blight Disease. Phytopathology, 2022, 112, 1500-1512.	2.2	4
12	TabPath: interactive tables for metabolic pathway analysis. Bioinformatics, 2018, 34, 1040-1042.	4.1	3
13	Deep Learning Neural Network Prediction Method Improves Proteome Profiling of Vascular Sap of Grapevines during Pierce's Disease Development. Biology, 2020, 9, 261.	2.8	3
14	A Secreted Chorismate Mutase from Xanthomonas arboricola pv. juglandis Attenuates Virulence and Walnut Blight Symptoms. International Journal of Molecular Sciences, 2021, 22, 10374.	4.1	2