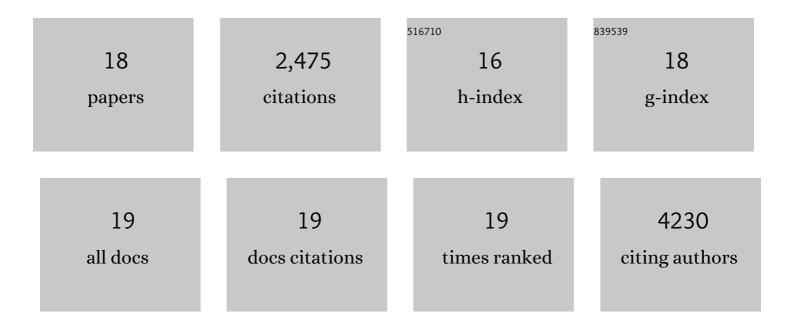
Sharvari Gujja

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11862963/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Identification of driver genes for critical forms of COVID-19 in a deeply phenotyped young patient cohort. Science Translational Medicine, 2022, 14, eabj7521.	12.4	71
2	Quantum processor-inspired machine learning in the biomedical sciences. Patterns, 2021, 2, 100246.	5.9	16
3	Smooth Muscle Cell Reprogramming in Aortic Aneurysms. Cell Stem Cell, 2020, 26, 542-557.e11.	11.1	114
4	Endothelial TGF-β signalling drives vascular inflammation and atherosclerosis. Nature Metabolism, 2019, 1, 912-926.	11.9	172
5	Genome Sequence for Candida albicans Clinical Oral Isolate 529L. Microbiology Resource Announcements, 2019, 8, .	0.6	13
6	Brown Fat AKT2 Is a Cold-Induced Kinase that Stimulates ChREBP-Mediated De Novo Lipogenesis to Optimize Fuel Storage and Thermogenesis. Cell Metabolism, 2018, 27, 195-209.e6.	16.2	151
7	Tracing Genetic Exchange and Biogeography of <i>Cryptococcus neoformans</i> var. <i>grubii</i> at the Global Population Level. Genetics, 2017, 207, 327-346.	2.9	105
8	Endothelial Mitogen-Activated Protein Kinase Kinase Kinase Kinase 4 ls Critical for Lymphatic Vascular Development and Function. Molecular and Cellular Biology, 2016, 36, 1740-1749.	2.3	21
9	Gld2-catalyzed 3′ monoadenylation of miRNAs in the hippocampus has no detectable effect on their stability or on animal behavior. Rna, 2016, 22, 1492-1499.	3.5	29
10	Cholesterol-Independent SREBP-1 Maturation Is Linked to ARF1 Inactivation. Cell Reports, 2016, 16, 9-18.	6.4	40
11	The Dynamic Genome and Transcriptome of the Human Fungal Pathogen Blastomyces and Close Relative Emmonsia. PLoS Genetics, 2015, 11, e1005493.	3.5	57
12	Genome Evolution and Innovation across the Four Major Lineages of Cryptococcus gattii. MBio, 2015, 6, e00868-15.	4.1	101
13	Genetic and phenotypic intra-species variation in <i>Candida albicans</i> . Genome Research, 2015, 25, 413-425.	5.5	305
14	Analysis of the Genome and Transcriptome of Cryptococcus neoformans var. grubii Reveals Complex RNA Expression and Microevolution Leading to Virulence Attenuation. PLoS Genetics, 2014, 10, e1004261.	3.5	336
15	Comparative Genomic and Transcriptomic Analysis of <i>Wangiella dermatitidis</i> , A Major Cause of Phaeohyphomycosis and a Model Black Yeast Human Pathogen. G3: Genes, Genomes, Genetics, 2014, 4, 561-578.	1.8	58
16	Genomic epidemiology of the <i>Escherichia coli</i> O104:H4 outbreaks in Europe, 2011. Proceedings of the United States of America, 2012, 109, 3065-3070.	7.1	262
17	Comparative Functional Genomics of the Fission Yeasts. Science, 2011, 332, 930-936.	12.6	458
18	Comparative Genomic Analysis of Human Fungal Pathogens Causing Paracoccidioidomycosis. PLoS Genetics, 2011, 7, e1002345.	3.5	164