Sharvari Gujja

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11862963/publications.pdf

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	516710	839539
2,475	16	18
citations	h-index	g-index
		4000
19	19	4230
docs citations	times ranked	citing authors
	2,475 citations 19 docs citations	2,475 16 citations h-index 19 19

#	Article	IF	CITATIONS
1	Comparative Functional Genomics of the Fission Yeasts. Science, 2011, 332, 930-936.	12.6	458
2	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
3	Genetic and phenotypic intra-species variation in <i>Candida albicans</i> . Genome Research, 2015, 25, 413-425.	5.5	305
4	Genomic epidemiology of the <i>Escherichia coli</i> O104:H4 outbreaks in Europe, 2011. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 3065-3070.	7.1	262
5	Endothelial TGF- \hat{l}^2 signalling drives vascular inflammation and atherosclerosis. Nature Metabolism, 2019, 1, 912-926.	11.9	172
6	Comparative Genomic Analysis of Human Fungal Pathogens Causing Paracoccidioidomycosis. PLoS Genetics, 2011, 7, e1002345.	3.5	164
7	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
8	Smooth Muscle Cell Reprogramming in Aortic Aneurysms. Cell Stem Cell, 2020, 26, 542-557.e11.	11.1	114
9	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
10	Genome Evolution and Innovation across the Four Major Lineages of Cryptococcus gattii. MBio, 2015, 6, e00868-15.	4.1	101
11	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
12	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
13	The Dynamic Genome and Transcriptome of the Human Fungal Pathogen Blastomyces and Close Relative Emmonsia. PLoS Genetics, 2015, 11, e1005493.	3.5	57
14	Cholesterol-Independent SREBP-1 Maturation Is Linked to ARF1 Inactivation. Cell Reports, 2016, 16, 9-18.	6.4	40
15	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
16	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
17	Quantum processor-inspired machine learning in the biomedical sciences. Patterns, 2021, 2, 100246.	5.9	16
18	Genome Sequence for Candida albicans Clinical Oral Isolate 529L. Microbiology Resource Announcements, 2019, 8, .	0.6	13