Jane Usher

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2683948/publications.pdf

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		1040056	1125743	
16	339	9	13	
papers	citations	h-index	g-index	
17	17	17	580	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Fluconazole resistant Candida auris clinical isolates have increased levels of cell wall chitin and increased susceptibility to a glucosamine-6-phosphate synthase inhibitor. Cell Surface, 2022, 8, 100076.	3.0	11
2	Genetic interaction analysis in microbial pathogens: unravelling networks of pathogenesis, antimicrobial susceptibility and host interactions. FEMS Microbiology Reviews, 2021, 45, .	8.6	8
3	Functional Characterization of a Novel Oxidative Stress Protection Protein in the Pathogenic Yeast Candida glabrata. Frontiers in Genetics, 2020, 11, 530915.	2.3	2
4	Advances in Molecular Tools and In Vivo Models for the Study of Human Fungal Pathogenesis. Microorganisms, 2020, 8, 803.	3.6	12
5	The Mechanisms of Mating in Pathogenic Fungi—A Plastic Trait. Genes, 2019, 10, 831.	2.4	11
6	Functional genomic characterization of metallothioneins in brown trout (Salmo trutta L.). using synthetic genetic analysis. Scientific Reports, 2019, 9, 11827.	3.3	0
7	Attenuating the emergence of anti-fungal drug resistance by harnessing synthetic lethal interactions in a model organism. PLoS Genetics, 2019, 15, e1008259.	3.5	18
8	Using DNA Barcoding to Investigate Patterns of Species Utilisation in UK Shark Products Reveals Threatened Species on Sale. Scientific Reports, 2019, 9, 1028.	3.3	38
9	Functional characterisation of novel oxidative stress protection proteins in the pathogenic yeast Candida glabrata. Access Microbiology, 2019, 1, .	0.5	O
10	Data-driven prediction of genetic interactions in Candida glabrata. Access Microbiology, 2019, $1, .$	0.5	0
11	Utilising established SDL-screening methods as a tool for the functional genomic characterisation of model and non-model organisms. FEMS Yeast Research, 2015, 15, fov091.	2.3	5
12	Combinatorial stresses kill pathogenic <i>Candida</i> species. Medical Mycology, 2012, 50, 699-709.	0.7	79
13	Chemical and Synthetic Genetic Array Analysis Identifies Genes that Suppress Xylose Utilization and Fermentation in <i>Saccharomyces cerevisiae</i> . G3: Genes, Genomes, Genetics, 2011, 1, 247-258.	1.8	22
14	Functional Genomics Analysis of the <i>Saccharomyces cerevisiae </i> Iron Responsive Transcription Factor Aft1 Reveals Iron-Independent Functions. Genetics, 2010, 185, 1111-1128.	2.9	34
15	Recombination between Homoeologous Chromosomes of Lager Yeasts Leads to Loss of Function of the Hybrid GPH1 Gene. Applied and Environmental Microbiology, 2009, 75, 4573-4579.	3.1	22
16	Lager yeasts possess dynamic genomes that undergo rearrangements and gene amplification in response to stress. Current Genetics, 2008, 53, 139-152.	1.7	77