

Yen-Ping Hsueh

List of Publications by Year in descending order

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36
papers

2,037
citations

331670

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31
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43
all docs

43
docs citations

43
times ranked

2047
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#	ARTICLE	IF	CITATIONS
1	Analysis of the Genome and Transcriptome of <i>Cryptococcus neoformans</i> var. <i>grubii</i> Reveals Complex RNA Expression and Microevolution Leading to Virulence Attenuation. <i>PLoS Genetics</i> , 2014, 10, e1004261.	3.5	336
2	Spores as Infectious Propagules of <i>Cryptococcus neoformans</i> . <i>Infection and Immunity</i> , 2009, 77, 4345-4355.	2.2	299
3	Magnificent seven: roles of G protein-coupled receptors in extracellular sensing in fungi. <i>FEMS Microbiology Reviews</i> , 2008, 32, 1010-1032.	8.6	165
4	Nematode-Trapping Fungi Eavesdrop on Nematode Pheromones. <i>Current Biology</i> , 2013, 23, 83-86.	3.9	152
5	Sex-induced silencing defends the genome of <i>Cryptococcus neoformans</i> via RNAi. <i>Genes and Development</i> , 2010, 24, 2566-2582.	5.9	134
6	Nematophagous fungus <i>Arthrobotrys oligospora</i> mimics olfactory cues of sex and food to lure its nematode prey. <i>ELife</i> , 2017, 6, .	6.0	75
7	Recombination Hotspots Flank the <i>Cryptococcus</i> Mating-Type Locus: Implications for the Evolution of a Fungal Sex Chromosome. <i>PLoS Genetics</i> , 2006, 2, e184.	3.5	72
8	Orchestration of sexual reproduction and virulence by the fungal mating-type locus. <i>Current Opinion in Microbiology</i> , 2008, 11, 517-524.	5.1	66
9	G protein signaling governing cell fate decisions involves opposing G β subunits in <i>Cryptococcus neoformans</i> . <i>Molecular Biology of the Cell</i> , 2007, 18, 3237-3249.	2.1	64
10	A constitutively active GPCR governs morphogenic transitions in <i>Cryptococcus neoformans</i> . <i>EMBO Journal</i> , 2009, 28, 1220-1233.	7.8	63
11	Natural diversity in the predatory behavior facilitates the establishment of a robust model strain for nematode-trapping fungi. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 6762-6770.	7.1	59
12	Gene Conversion Occurs within the Mating-Type Locus of <i>Cryptococcus neoformans</i> during Sexual Reproduction. <i>PLoS Genetics</i> , 2012, 8, e1002810.	3.5	54
13	Discovery of a Modified Tetrapolar Sexual Cycle in <i>Cryptococcus amyloletus</i> and the Evolution of MAT in the <i>Cryptococcus</i> Species Complex. <i>PLoS Genetics</i> , 2012, 8, e1002528.	3.5	54
14	The RGS protein Crg2 regulates both pheromone and cAMP signalling in <i>Cryptococcus neoformans</i> . <i>Molecular Microbiology</i> , 2008, 70, 379-395.	2.5	53
15	Predator-prey interactions of nematode-trapping fungi and nematodes: both sides of the coin. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 3939-3949.	3.6	52
16	Transitions in Sexuality: Recapitulation of an Ancestral Tri- and Tetrapolar Mating System in <i>Cryptococcus neoformans</i> . <i>Eukaryotic Cell</i> , 2008, 7, 1847-1855.	3.4	50
17	Transmission of Hypervirulence Traits via Sexual Reproduction within and between Lineages of the Human Fungal Pathogen <i>Cryptococcus gattii</i> . <i>PLoS Genetics</i> , 2013, 9, e1003771.	3.5	45
18	A Homolog of Ste6, the α -Factor Transporter in <i>Saccharomyces cerevisiae</i> , Is Required for Mating but Not for Monokaryotic Fruiting in <i>Cryptococcus neoformans</i> . <i>Eukaryotic Cell</i> , 2005, 4, 147-155.	3.4	33

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19	Prey sensing and response in a nematode-trapping fungus is governed by the MAPK pheromone response pathway. <i>Genetics</i> , 2021, 217, .	2.9	30
20	The High Osmolarity Glycerol (HOG) Pathway Functions in Osmosensing, Trap Morphogenesis and Conidiation of the Nematode-Trapping Fungus <i>Arthrobotrys oligospora</i> . <i>Journal of Fungi (Basel)</i> , 2021, 7, 10.	2.9	10
21	Nematode-Trapping Fungi Produce Diverse Metabolites during Predator-Prey Interaction. <i>Metabolites</i> , 2020, 10, 117.	2.9	25
22	Evolution of the Mating-Type Locus: The Basidiomycetes. , 0, , 19-34.		25
23	Fungal feature tracker (FFT): A tool for quantitatively characterizing the morphology and growth of filamentous fungi. <i>PLoS Computational Biology</i> , 2019, 15, e1007428.	3.2	20
24	Sensory cilia as the Achilles heel of nematodes when attacked by carnivorous mushrooms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 6014-6022.	7.1	20
25	Genome sequence of the oyster mushroom <i>Pleurotus ostreatus</i> strain PC9. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	20
26	The Mating-Type Locus of <i>Cryptococcus</i> : Evolution of Gene Clusters Governing Sex Determination and Sexual Reproduction from the Phylogenomic Perspective. , 0, , 139-149.		7
27	Nematode-free agricultural system of a fungus-growing termite. <i>Scientific Reports</i> , 2019, 9, 8917.	3.3	6
28	Epigenetic Manipulation Induces the Production of Coumarin-Type Secondary Metabolite from <i>Arthrobotrys foliicola</i> . <i>Israel Journal of Chemistry</i> , 2019, 59, 432-438.	2.3	6
29	Genomic analyses of two Italian oyster mushroom <i>Pleurotus pulmonarius</i> strains. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	6
30	Laboratory Maintenance and Culturing of the Nematode-Trapping Fungus <i>Arthrobotrys oligospora</i> . <i>Current Protocols</i> , 2021, 1, e41.	2.9	5
31	Forward genetic screens identified mutants with defects in trap morphogenesis in the nematode-trapping fungus <i>Arthrobotrys oligospora</i> . <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	4
32	Sexual Reproduction of <i>Cryptococcus</i> . , 0, , 81-96.		3
33	Assessment of Constitutive Activity of a G Protein-Coupled Receptor, Cpr2, in <i>Cryptococcus neoformans</i> by Heterologous and Homologous Methods. <i>Methods in Enzymology</i> , 2010, 484, 397-412.	1.0	2
34	The conserved regulator of autophagy and innate immunity <i>hlh-30/TFEB</i> mediates tolerance of enterohemorrhagic <i>Escherichia coli</i> in <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2021, 217, 1-17.	2.9	2
35	Possible impacts of the predominant <i>Bacillus</i> bacteria on the <i>Ophiocordyceps unilateralis</i> s. l. in its infected ant cadavers. <i>Scientific Reports</i> , 2021, 11, 22695.	3.3	2
36	Nematode Communication. , 2014, , 383-407.		0