

Guanghua Huang

List of Publications by Year in descending order

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57
papers

2,679
citations

236925

25
h-index

189892

50
g-index

60
all docs

60
docs citations

60
times ranked

1915
citing authors

#	ARTICLE	IF	CITATIONS
1	Bistable expression of WOR1, a master regulator of white-opaque switching in <i>Candida albicans</i> . Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 12813-12818.	7.1	277
2	<i>Candida auris</i> : Epidemiology, biology, antifungal resistance, and virulence. PLoS Pathogens, 2020, 16, e1008921.	4.7	270
3	N-Acetylglucosamine Induces White to Opaque Switching, a Mating Prerequisite in <i>Candida albicans</i> . PLoS Pathogens, 2010, 6, e1000806.	4.7	180
4	CO ₂ Regulates White-to-Opaque Switching in <i>Candida albicans</i> . Current Biology, 2009, 19, 330-334.	3.9	160
5	Regulation of phenotypic transitions in the fungal pathogen <i>Candida albicans</i> . Virulence, 2012, 3, 251-261.	4.4	130
6	The first isolate of <i>Candida auris</i> in China: clinical and biological aspects. Emerging Microbes and Infections, 2018, 7, 1-9.	6.5	126
7	Discovery of a "White-Gray-Opaque" Tristable Phenotypic Switching System in <i>Candida albicans</i> : Roles of Non-genetic Diversity in Host Adaptation. PLoS Biology, 2014, 12, e1001830.	5.6	122
8	White-Opaque Switching in Natural MTL ^a Isolates of <i>Candida albicans</i> : Evolutionary Implications for Roles in Host Adaptation, Pathogenesis, and Sex. PLoS Biology, 2013, 11, e1001525.	5.6	107
9	Filamentation in <i>Candida auris</i> , an emerging fungal pathogen of humans: passage through the mammalian body induces a heritable phenotypic switch. Emerging Microbes and Infections, 2018, 7, 1-13.	6.5	105
10	Tec1 Mediates the Pheromone Response of the White Phenotype of <i>Candida albicans</i> : Insights into the Evolution of New Signal Transduction Pathways. PLoS Biology, 2010, 8, e1000363.	5.6	85
11	Alternative Mating Type Configurations (a ⁺ versus a/a or ⁺ / ⁺) of <i>Candida albicans</i> Result in Alternative Biofilms Regulated by Different Pathways. PLoS Biology, 2011, 9, e1001117.	5.6	73
12	Multiple roles and diverse regulation of the Ras/cAMP/protein kinase A pathway in <i>Candida albicans</i> . Molecular Microbiology, 2019, 111, 6-16.	2.5	64
13	Roles of <i>Candida albicans</i> Gat2, a GATA-Type Zinc Finger Transcription Factor, in Biofilm Formation, Filamentous Growth and Virulence. PLoS ONE, 2012, 7, e29707.	2.5	61
14	Global regulatory roles of the cAMP/PKA pathway revealed by phenotypic, transcriptomic and phosphoproteomic analyses in a null mutant of the PKA catalytic subunit in <i>Candida albicans</i> . Molecular Microbiology, 2017, 105, 46-64.	2.5	60
15	N-Acetylglucosamine Induces White-to-Opaque Switching and Mating in <i>Candida tropicalis</i> , Providing New Insights into Adaptation and Fungal Sexual Evolution. Eukaryotic Cell, 2012, 11, 773-782.	3.4	58
16	Integration of the tricarboxylic acid (TCA) cycle with cAMP signaling and Sfl2 pathways in the regulation of CO ₂ sensing and hyphal development in <i>Candida albicans</i> . PLoS Genetics, 2017, 13, e1006949.	3.5	58
17	The transcription factor Flo8 mediates CO ₂ sensing in the human fungal pathogen <i>Candida albicans</i> . Molecular Biology of the Cell, 2012, 23, 2692-2701.	2.1	51
18	Experimental Evolution Identifies Adaptive Aneuploidy as a Mechanism of Fluconazole Resistance in <i>Candida auris</i> . Antimicrobial Agents and Chemotherapy, 2020, 65, .	3.2	46

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19	Environmental pH adaption and morphological transitions in <i>Candida albicans</i> . <i>Current Genetics</i> , 2016, 62, 283-286.	1.7	37
20	<i>Bcr1</i> plays a central role in the regulation of opaque cell filamentation in <i>Candida albicans</i> . <i>Molecular Microbiology</i> , 2013, 89, 732-750.	2.5	36
21	<i>N</i> -Acetylglucosamine-Induced Cell Death in <i>Candida albicans</i> and Its Implications for Adaptive Mechanisms of Nutrient Sensing in Yeasts. <i>MBio</i> , 2015, 6, e01376-15.	4.1	35
22	pH Regulates White-Opaque Switching and Sexual Mating in <i>Candida albicans</i> . <i>Eukaryotic Cell</i> , 2015, 14, 1127-1134.	3.4	34
23	Regulation of filamentation in the human fungal pathogen <i>Candida tropicalis</i> . <i>Molecular Microbiology</i> , 2016, 99, 528-545.	2.5	34
24	Beauvericin counteracted multi-drug resistant <i>Candida albicans</i> by blocking ABC transporters. <i>Synthetic and Systems Biotechnology</i> , 2016, 1, 158-168.	3.7	31
25	Innate immune responses against the fungal pathogen <i>Candida auris</i> . <i>Nature Communications</i> , 2022, 13, .	12.8	30
26	Lactic acid bacteria differentially regulate filamentation in two heritable cell types of the human fungal pathogen <i>Candida albicans</i> . <i>Molecular Microbiology</i> , 2016, 102, 506-519.	2.5	29
27	White Cells Facilitate Opposite- and Same-Sex Mating of Opaque Cells in <i>Candida albicans</i> . <i>PLoS Genetics</i> , 2014, 10, e1004737.	3.5	23
28	Self-Induction of <i>a</i> / <i>a</i> or $\hat{1}\pm/\hat{1}\pm$ Biofilms in <i>Candida albicans</i> Is a Pheromone-Based Paracrine System Requiring Switching. <i>Eukaryotic Cell</i> , 2011, 10, 753-760.	3.4	22
29	Genomic epidemiology of <i>Candida auris</i> in a general hospital in Shenyang, China: a three-year surveillance study. <i>Emerging Microbes and Infections</i> , 2021, 10, 1088-1096.	6.5	21
30	The mitochondrial protein <i>Mcu1</i> plays important roles in carbon source utilization, filamentation, and virulence in <i>Candida albicans</i> . <i>Fungal Genetics and Biology</i> , 2015, 81, 150-159.	2.1	20
31	Environmental and genetic regulation of white–opaque switching in <i>Candida tropicalis</i> . <i>Molecular Microbiology</i> , 2017, 106, 999-1017.	2.5	20
32	The Regulatory Subunit of Protein Kinase A (<i>Bcy1</i>) in <i>Candida albicans</i> Plays Critical Roles in Filamentation and White-Opaque Switching but Is Not Essential for Cell Growth. <i>Frontiers in Microbiology</i> , 2016, 7, 2127.	3.5	19
33	Environment-induced same-sex mating in the yeast <i>Candida albicans</i> through the <i>Hsf1</i> “ <i>Hsp90</i> pathway. <i>PLoS Biology</i> , 2019, 17, e2006966.	5.6	19
34	Filamentous growth is a general feature of <i>Candida auris</i> clinical isolates. <i>Medical Mycology</i> , 2021, 59, 734-740.	0.7	19
35	The zinc-finger transcription factor, <i>Ofi1</i> , regulates white–opaque switching and filamentation in the yeast <i>Candida albicans</i> . <i>Acta Biochimica Et Biophysica Sinica</i> , 2015, 47, 335-341.	2.0	16
36	Discovery of the gray phenotype and white-gray-opaque tristable phenotypic transitions in <i>Candida dubliniensis</i> . <i>Virulence</i> , 2016, 7, 230-242.	4.4	15

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37	A coupled process of same- and opposite-sex mating generates polyploidy and genetic diversity in <i>Candida tropicalis</i> . <i>PLoS Genetics</i> , 2018, 14, e1007377.	3.5	14
38	The gray phenotype and tristable phenotypic transitions in the human fungal pathogen <i>Candida tropicalis</i> . <i>Fungal Genetics and Biology</i> , 2016, 93, 10-16.	2.1	13
39	Role of the N-acetylglucosamine kinase (Hxk1) in the regulation of white-gray-opaque tristable phenotypic transitions in <i>C. albicans</i> . <i>Fungal Genetics and Biology</i> , 2016, 92, 26-32.	2.1	11
40	Revision of the medically relevant species of the yeast genus <i>Diutina</i> . <i>Medical Mycology</i> , 2019, 57, 226-233.	0.7	11
41	A biological and genomic comparison of a drug-resistant and a drug-susceptible strain of <i>Candida auris</i> isolated from Beijing, China. <i>Virulence</i> , 2021, 12, 1388-1399.	4.4	11
42	Phenotypic diversity and correlation between white-to-opaque switching and the CAI microsatellite locus in <i>Candida albicans</i> . <i>Current Genetics</i> , 2016, 62, 585-593.	1.7	10
43	Discovery of the Diploid Form of the Emerging Fungal Pathogen <i>Candida auris</i> . <i>ACS Infectious Diseases</i> , 2020, 6, 2641-2646.	3.8	10
44	A case of <i>Candida auris</i> candidemia in Xiamen, China, and a comparative analysis of clinical isolates in China. <i>Mycology</i> , 2022, 13, 68-75.	4.4	10
45	<i>Streptococcus mutans</i> suppresses filamentous growth of <i>Candida albicans</i> through secreting mutanocyclin, an unacylated tetramic acid. <i>Virulence</i> , 2022, 13, 542-557.	4.4	10
46	N-acetylglucosamine-induced white-to-opaque switching in <i>Candida albicans</i> is independent of the Wor2 transcription factor. <i>Fungal Genetics and Biology</i> , 2014, 62, 71-77.	2.1	9
47	N-Acetylglucosamine (GlcNAc) Sensing, Utilization, and Functions in <i>Candida albicans</i> . <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 129.	3.5	9
48	<i>Candida auris</i> infections in China. <i>Virulence</i> , 2022, 13, 589-591.	4.4	9
49	The Als3 Cell Wall Adhesin Plays a Critical Role in Human Serum Amyloid A1-Induced Cell Death and Aggregation in <i>Candida albicans</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, .	3.2	8
50	Antifungal Activity of Mammalian Serum Amyloid A1 against <i>Candida albicans</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 64, .	3.2	7
51	The general transcriptional repressor Tup1 governs filamentous development in. <i>Acta Biochimica Et Biophysica Sinica</i> , 2019, 51, 463-470.	2.0	4
52	The PHO pathway regulates white-to-opaque switching and sexual mating in the human fungal pathogen <i>Candida albicans</i> . <i>Current Genetics</i> , 2020, 66, 1155-1162.	1.7	4
53	Genetic regulation of the development of mating projections in <i>Candida albicans</i> . <i>Emerging Microbes and Infections</i> , 2020, 9, 413-426.	6.5	3
54	Ploidy Variation and Spontaneous Haploid-Diploid Switching of <i>Candida glabrata</i> Clinical Isolates. <i>MSphere</i> , 2022, 7, .	2.9	3

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55	Biological and genomic analyses of a clinical isolate of <i>Yarrowia galli</i> from China. <i>Current Genetics</i> , 2020, 66, 549-559.	1.7	2
56	Epigenetic Switching in the Human Fungal Pathogen <i>Candida albicans</i> . <i>Epigenetics and Human Health</i> , 2017, , 175-187.	0.2	0
57	<i>Candida albicans</i> MTL _{a2} regulates the mating response through both the a-factor and $\hat{\pm}$ -factor sensing pathways. <i>Fungal Genetics and Biology</i> , 2022, 159, 103664.	2.1	0