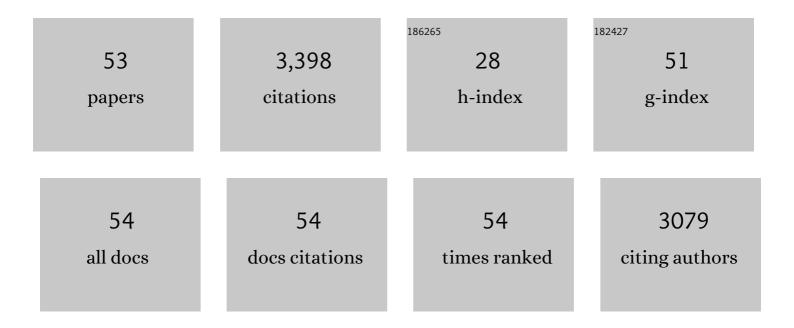
Chaoyang Xue

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Macrophage Mediated Immunomodulation During Cryptococcus Pulmonary Infection. Frontiers in Cellular and Infection Microbiology, 2022, 12, 859049.	3.9	10
2	Development of Antifungal Peptides against Cryptococcus neoformans; Leveraging Knowledge about the <i>cdc50Δ</i> Mutant Susceptibility for Lead Compound Development. Microbiology Spectrum, 2022, 10, e0043922.	3.0	5
3	Inositol Metabolism Regulates Capsule Structure and Virulence in the Human Pathogen Cryptococcus neoformans. MBio, 2021, 12, e0279021.	4.1	10
4	More Than Just Cleaning: Ubiquitin-Mediated Proteolysis in Fungal Pathogenesis. Frontiers in Cellular and Infection Microbiology, 2021, 11, 774613.	3.9	15
5	A spontaneous mutation in DNA polymerase POL3 during in vitro passaging causes a hypermutator phenotype in Cryptococcus species. DNA Repair, 2020, 86, 102751.	2.8	10
6	Activation of Meiotic Genes Mediates Ploidy Reduction during Cryptococcal Infection. Current Biology, 2020, 30, 1387-1396.e5.	3.9	27
7	More than flipping the lid: Cdc50 contributes to echinocandin resistance by regulating calcium homeostasis in Cryptococcus neoformans. Microbial Cell, 2020, 7, 115-118.	3.2	6
8	Identification of Pathogen Genomic Differences That Impact Human Immune Response and Disease during Cryptococcus neoformans Infection. MBio, 2019, 10, .	4.1	39
9	Nutrient and Stress Sensing in Pathogenic Yeasts. Frontiers in Microbiology, 2019, 10, 442.	3.5	41
10	A Mechanosensitive Channel Governs Lipid Flippase-Mediated Echinocandin Resistance in Cryptococcus neoformans. MBio, 2019, 10, .	4.1	28
11	A Heat-Killed <i>Cryptococcus</i> Mutant Strain Induces Host Protection against Multiple Invasive Mycoses in a Murine Vaccine Model. MBio, 2019, 10, .	4.1	36
12	Phosphatidylserine synthesis is essential for viability of the human fungal pathogen Cryptococcus neoformans. Journal of Biological Chemistry, 2019, 294, 2329-2339.	3.4	14
13	Role of the inositol pyrophosphate multikinase Kcs1 in Cryptococcus inositol metabolism. Fungal Genetics and Biology, 2018, 113, 42-51.	2.1	5
14	The F-Box Protein Fbp1 Shapes the Immunogenic Potential of Cryptococcus neoformans. MBio, 2018, 9, .	4.1	28
15	Nutrient Sensing at the Plasma Membrane of Fungal Cells. Microbiology Spectrum, 2017, 5, .	3.0	24
16	Mismatch Repair of DNA Replication Errors Contributes to Microevolution in the Pathogenic Fungus <i>Cryptococcus neoformans</i> . MBio, 2017, 8, .	4.1	76
17	Nutrient Sensing at the Plasma Membrane of Fungal Cells. , 2017, , 417-439.		4
18	Cryptococcus flips its lid - membrane phospholipid asymmetry modulates antifungal drug resistance and virulence. Microbial Cell, 2016, 3, 358-360.	3.2	7

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#	Article	IF	CITATIONS
19	Lipid Flippase Subunit Cdc50 Mediates Drug Resistance and Virulence in Cryptococcus neoformans. MBio, 2016, 7, .	4.1	60
20	Crystal structure of Gib2, a signal-transducing protein scaffold associated with ribosomes in Cryptococcus neoformans. Scientific Reports, 2015, 5, 8688.	3.3	11
21	Finding the Sweet Spot: How Human Fungal Pathogens Acquire and Turn the Sugar Inositol against Their Hosts. MBio, 2015, 6, e00109.	4.1	5
22	Cryptococcus inositol utilization modulates the host protective immune response during brain infection. Cell Communication and Signaling, 2014, 12, 51.	6.5	23
23	Bypassing Both Surface Attachment and Surface Recognition Requirements for Appressorium Formation by Overactive Ras Signaling in <i>Magnaporthe oryzae</i> . Molecular Plant-Microbe Interactions, 2014, 27, 996-1004.	2.6	41
24	Fbp1-Mediated Ubiquitin-Proteasome Pathway Controls Cryptococcus neoformans Virulence by Regulating Fungal Intracellular Growth in Macrophages. Infection and Immunity, 2014, 82, 557-568.	2.2	56
25	How Fungi Sense Sugars, Alcohols, and Amino Acids. , 2014, , 467-479.		0
26	Brain Inositol Is a Novel Stimulator for Promoting Cryptococcus Penetration of the Blood-Brain Barrier. PLoS Pathogens, 2013, 9, e1003247.	4.7	69
27	The Glucose Sensor-Like Protein Hxs1 Is a High-Affinity Glucose Transporter and Required for Virulence in Cryptococcus neoformans. PLoS ONE, 2013, 8, e64239.	2.5	18
28	Cryptococcus and Beyond—Inositol Utilization and Its Implications for the Emergence of Fungal Virulence. PLoS Pathogens, 2012, 8, e1002869.	4.7	29
29	DNA Mutations Mediate Microevolution between Host-Adapted Forms of the Pathogenic Fungus Cryptococcus neoformans. PLoS Pathogens, 2012, 8, e1002936.	4.7	76
30	Molecular mechanisms of cryptococcal meningitis. Virulence, 2012, 3, 173-181.	4.4	105
31	Characterization and Complete Nucleotide Sequence of Two Isolates of <i>Tomato mosaic virus</i> . Journal of Phytopathology, 2012, 160, 115-119.	1.0	4
32	A Pmk1-Interacting Gene Is Involved in Appressorium Differentiation and Plant Infection in <i>Magnaporthe oryzae</i> . Eukaryotic Cell, 2011, 10, 1062-1070.	3.4	31
33	The Casein Kinase I Protein Cck1 Regulates Multiple Signaling Pathways and Is Essential for Cell Integrity and Fungal Virulence in Cryptococcus neoformans. Eukaryotic Cell, 2011, 10, 1455-1464.	3.4	19
34	The Ubiquitin-Proteasome System and F-box Proteins in Pathogenic Fungi. Mycobiology, 2011, 39, 243-248.	1.7	56
35	The F-Box Protein Fbp1 Regulates Sexual Reproduction and Virulence in Cryptococcus neoformans. Eukaryotic Cell, 2011, 10, 791-802.	3.4	61
36	Two Major Inositol Transporters and Their Role in Cryptococcal Virulence. Eukaryotic Cell, 2011, 10, 618-628.	3.4	31

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#	Article	lF	CITATIONS
37	Cryptococcal Titan Cell Formation Is Regulated by G-Protein Signaling in Response to Multiple Stimuli. Eukaryotic Cell, 2011, 10, 1306-1316.	3.4	105
38	Transcription Factors Mat2 and Znf2 Operate Cellular Circuits Orchestrating Opposite- and Same-Sex Mating in Cryptococcus neoformans. PLoS Genetics, 2010, 6, e1000953.	3.5	111
39	Assessment of Constitutive Activity of a G Protein-Coupled Receptor, Cpr2, in Cryptococcus neoformans by Heterologous and Homologous Methods. Methods in Enzymology, 2010, 484, 397-412.	1.0	2
40	Role of an Expanded Inositol Transporter Repertoire in Cryptococcus neoformans Sexual Reproduction and Virulence. MBio, 2010, 1, .	4.1	61
41	A constitutively active GPCR governs morphogenic transitions in Cryptococcus neoformans. EMBO Journal, 2009, 28, 1220-1233.	7.8	63
42	Magnificent seven: roles of G protein-coupled receptors in extracellular sensing in fungi. FEMS Microbiology Reviews, 2008, 32, 1010-1032.	8.6	165
43	The RGS protein Crg2 regulates both pheromone and cAMP signalling in <i>Cryptococcus neoformans</i> . Molecular Microbiology, 2008, 70, 379-395.	2.5	53
44	The Human Fungal Pathogen Cryptococcus Can Complete Its Sexual Cycle during a Pathogenic Association with Plants. Cell Host and Microbe, 2007, 1, 263-273.	11.0	175
45	G protein signaling governing cell fate decisions involves opposing Gα subunits inCryptococcus neoformans. Molecular Biology of the Cell, 2007, 18, 3237-3249.	2.1	64
46	Sensing the environment: lessons from fungi. Nature Reviews Microbiology, 2007, 5, 57-69.	28.6	331
47	Multiple Upstream Signals Converge on the Adaptor Protein Mst50 in Magnaporthe grisea. Plant Cell, 2006, 18, 2822-2835.	6.6	147
48	G Protein-coupled Receptor Gpr4 Senses Amino Acids and Activates the cAMP-PKA Pathway inCryptococcus neoformans. Molecular Biology of the Cell, 2006, 17, 667-679.	2.1	144
49	Two PAK Kinase Genes, CHM1 and MST20, Have Distinct Functions in Magnaporthe grisea. Molecular Plant-Microbe Interactions, 2004, 17, 547-556.	2.6	89
50	A Mitogen-Activated Protein Kinase Gene (MGV1) in Fusarium graminearum Is Required for Female Fertility, Heterokaryon Formation, and Plant Infection. Molecular Plant-Microbe Interactions, 2002, 15, 1119-1127.	2.6	442
51	Two Novel Fungal Virulence Genes Specifically Expressed in Appressoria of the Rice Blast Fungus. Plant Cell, 2002, 14, 2107-2119.	6.6	161
52	MST12 Regulates Infectious Growth But Not Appressorium Formation in the Rice Blast Fungus Magnaporthe grisea. Molecular Plant-Microbe Interactions, 2002, 15, 183-192.	2.6	194
53	Time for a blast: genomics ofMagnaporthe grisea. Molecular Plant Pathology, 2002, 3, 173-176.	4.2	11