

Ana Traven

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/599295/publications.pdf>

Version: 2024-02-01

64
papers

3,421
citations

159525

30
h-index

149623

56
g-index

66
all docs

66
docs citations

66
times ranked

5055
citing authors

#	ARTICLE	IF	CITATIONS
1	Disruption of Iron Homeostasis and Mitochondrial Metabolism Are Promising Targets to Inhibit <i>Candida auris</i> . <i>Microbiology Spectrum</i> , 2022, 10, e0010022.	1.2	9
2	Natural Variation in Clinical Isolates of <i>Candida albicans</i> Modulates Neutrophil Responses. <i>MSphere</i> , 2020, 5, .	1.3	12
3	Metabolic competition between host and pathogen dictates inflammasome responses to fungal infection. <i>PLoS Pathogens</i> , 2020, 16, e1008695.	2.1	28
4	Immunometabolism in fungal infections: the need to eat to compete. <i>Current Opinion in Microbiology</i> , 2020, 58, 32-40.	2.3	23
5	The YEATS Domain Histone Crotonylation Readers Control Virulence-Related Biology of a Major Human Pathogen. <i>Cell Reports</i> , 2020, 31, 107528.	2.9	19
6	Targeting NLRP3 and Staphylococcal pore-forming toxin receptors in human-induced pluripotent stem cell-derived macrophages. <i>Journal of Leukocyte Biology</i> , 2020, 108, 967-981.	1.5	19
7	The RSC (Remodels the Structure of Chromatin) complex of <i>Candida albicans</i> shows compositional divergence with distinct roles in regulating pathogenic traits. <i>PLoS Genetics</i> , 2020, 16, e1009071.	1.5	8
8	Central metabolic interactions of immune cells and microbes: prospects for defeating infections. <i>EMBO Reports</i> , 2019, 20, e47995.	2.0	47
9	Characterization of Key Bioâ€“Nano Interactions between Organosilica Nanoparticles and <i>Candida albicans</i> . <i>ACS Applied Materials & Interfaces</i> , 2019, 11, 34676-34687.	4.0	11
10	Mdivi-1 and mitochondrial fission: recent insights from fungal pathogens. <i>Current Genetics</i> , 2019, 65, 837-845.	0.8	14
11	Mitochondrial Control of Fungal Cell Walls: Models and Relevance in Fungal Pathogens. <i>Current Topics in Microbiology and Immunology</i> , 2019, 425, 277-296.	0.7	20
12	Glucose Homeostasis Is Important for Immune Cell Viability during <i>Candida</i> Challenge and Host Survival of Systemic Fungal Infection. <i>Cell Metabolism</i> , 2018, 27, 988-1006.e7.	7.2	162
13	A Metabolic Checkpoint for the Yeast-to-Hyphae Developmental Switch Regulated by Endogenous Nitric Oxide Signaling. <i>Cell Reports</i> , 2018, 25, 2244-2258.e7.	2.9	37
14	The Antifungal Plant Defensin HsAFP1 Is a Phosphatidic Acid-Interacting Peptide Inducing Membrane Permeabilization. <i>Frontiers in Microbiology</i> , 2017, 8, 2295.	1.5	36
15	The Mitochondrial GTPase Gem1 Contributes to the Cell Wall Stress Response and Invasive Growth of <i>Candida albicans</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 2555.	1.5	15
16	The Endoplasmic Reticulum-Mitochondrion Tether ERMES Orchestrates Fungal Immune Evasion, Illuminating Inflammasome Responses to Hyphal Signals. <i>MSphere</i> , 2016, 1, .	1.3	39
17	Postâ€“transcriptional gene regulation in the biology and virulence of <i>Candida albicans</i> . <i>Cellular Microbiology</i> , 2016, 18, 800-806.	1.1	22
18	Searching for new strategies against polymicrobial biofilm infections: guanylated polymethacrylates kill mixed fungal/bacterial biofilms. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 413-421.	1.3	65

#	ARTICLE	IF	CITATIONS
19	Anti-infective Surface Coatings: Design and Therapeutic Promise against Device-Associated Infections. PLoS Pathogens, 2016, 12, e1005598.	2.1	43
20	Candida and macrophages: a deadly affair. Microbiology Australia, 2015, 36, 53.	0.1	0
21	Integration of Posttranscriptional Gene Networks into Metabolic Adaptation and Biofilm Maturation in <i>Candida albicans</i> . PLoS Genetics, 2015, 11, e1005590.	1.5	31
22	System-level impact of mitochondria on fungal virulence: to metabolism and beyond. FEMS Yeast Research, 2015, 15, fov027.	1.1	93
23	PAT-seq: a method to study the integration of 3' UTR dynamics with gene expression in the eukaryotic transcriptome. Rna, 2015, 21, 1502-1510.	1.6	78
24	Surface coatings with covalently attached caspofungin are effective in eliminating fungal pathogens. Journal of Materials Chemistry B, 2015, 3, 8469-8476.	2.9	31
25	Mitochondrial Biogenesis: Cell-Cycle-Dependent Investment in Making Mitochondria. Current Biology, 2015, 25, R78-R80.	1.8	24
26	Identification of a Class of Protein ADP-Ribosylating Sirtuins in Microbial Pathogens. Molecular Cell, 2015, 59, 309-320.	4.5	79
27	Microbial Egress: A Hitchhiker's Guide to Freedom. PLoS Pathogens, 2014, 10, e1004201.	2.1	19
28	The Pathogen <i>Candida albicans</i> Hijacks Pyroptosis for Escape from Macrophages. MBio, 2014, 5, e00003-14.	1.8	181
29	Activation of stress signalling pathways enhances tolerance of fungi to chemical fungicides and antifungal proteins. Cellular and Molecular Life Sciences, 2014, 71, 2651-2666.	2.4	76
30	Bovine pancreatic trypsin inhibitor is a new antifungal peptide that inhibits cellular magnesium uptake. Molecular Microbiology, 2014, 92, 1188-1197.	1.2	25
31	A Global Virulence Regulator in <i>Acinetobacter baumannii</i> and Its Control of the Phenylacetic Acid Catabolic Pathway. Journal of Infectious Diseases, 2014, 210, 46-55.	1.9	139
32	Solvent-exposed serines in the Gal4 DNA-binding domain are required for promoter occupancy and transcriptional activation <i>in vivo</i> . FEMS Yeast Research, 2014, 14, 302-309.	1.1	3
33	RAFT-derived antimicrobial polymethacrylates: elucidating the impact of end-groups on activity and cytotoxicity. Polymer Chemistry, 2014, 5, 5813-5822.	1.9	68
34	The ins and outs of the intermembrane space: Diverse mechanisms and evolutionary rewiring of mitochondrial protein import routes. Biochimica Et Biophysica Acta - General Subjects, 2014, 1840, 1246-1253.	1.1	12
35	<i>let-1</i> is a broad scale translational repressor required for normal P granule formation in <i>C. elegans</i> . Journal of Cell Science, 2013, 126, 850-9.	1.2	32
36	Identification and Mechanism of Action of the Plant Defensin NaD1 as a New Member of the Antifungal Drug Arsenal against <i>Candida albicans</i> . Antimicrobial Agents and Chemotherapy, 2013, 57, 3667-3675.	1.4	104

#	ARTICLE	IF	CITATIONS
37	Phospholipase C of <i>Cryptococcus neoformans</i> Regulates Homeostasis and Virulence by Providing Inositol Trisphosphate as a Substrate for Arg1 Kinase. <i>Infection and Immunity</i> , 2013, 81, 1245-1255.	1.0	36
38	Guanylated Polymethacrylates: A Class of Potent Antimicrobial Polymers with Low Hemolytic Activity. <i>Biomacromolecules</i> , 2013, 14, 4021-4031.	2.6	174
39	The cellular roles of Ccr4-NOT in model and pathogenic fungi—implications for fungal virulence. <i>Frontiers in Genetics</i> , 2013, 4, 302.	1.1	23
40	Preparation of Mitochondria from <i>Candida albicans</i> . <i>Bio-protocol</i> , 2013, 3, .	0.2	1
41	<i>Candida albicans</i> Mitochondrial Protein Import Assay. <i>Bio-protocol</i> , 2013, 3, .	0.2	1
42	The Functions of Mediator in <i>Candida albicans</i> Support a Role in Shaping Species-Specific Gene Expression. <i>PLoS Genetics</i> , 2012, 8, e1002613.	1.5	50
43	Mitochondrial Sorting and Assembly Machinery Subunit Sam37 in <i>Candida albicans</i> : Insight into the Roles of Mitochondria in Fitness, Cell Wall Integrity, and Virulence. <i>Eukaryotic Cell</i> , 2012, 11, 532-544.	3.4	57
44	A Small Tim Homohexamer in the Relict Mitochondrion of <i>Cryptosporidium</i> . <i>Molecular Biology and Evolution</i> , 2012, 29, 113-122.	3.5	22
45	ePAT: A simple method to tag adenylated RNA to measure poly(A)-tail length and other 3' RACE applications. <i>Rna</i> , 2012, 18, 1289-1295.	1.6	87
46	A model system for mitochondrial biogenesis reveals evolutionary rewiring of protein import and membrane assembly pathways. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E3358-66.	3.3	30
47	Transcriptional Profiling of a Yeast Colony Provides New Insight into the Heterogeneity of Multicellular Fungal Communities. <i>PLoS ONE</i> , 2012, 7, e46243.	1.1	34
48	The mRNA Decay Pathway Regulates the Expression of the Flo11 Adhesin and Biofilm Formation in <i>Saccharomyces cerevisiae</i> . <i>Genetics</i> , 2012, 191, 1387-1391.	1.2	8
49	Cell wall integrity is linked to mitochondria and phospholipid homeostasis in <i>Candida albicans</i> through the activity of the post-transcriptional regulator Ccr4-Pop2. <i>Molecular Microbiology</i> , 2011, 79, 968-989.	1.2	115
50	PUF proteins: repression, activation and mRNA localization. <i>Trends in Cell Biology</i> , 2011, 21, 104-112.	3.6	263
51	Mitochondria and Fungal Pathogenesis: Drug Tolerance, Virulence, and Potential for Antifungal Therapy. <i>Eukaryotic Cell</i> , 2011, 10, 1376-1383.	3.4	198
52	Probing connectivity between transcriptional and post-transcriptional gene networks. <i>Microbiology Australia</i> , 2011, 32, 166.	0.1	0
53	Dual functions of Mdt1 in genome maintenance and cell integrity pathways in <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 2010, 27, 41-52.	0.8	7
54	The Yeast PUF Protein Puf5 Has Pop2-Independent Roles in Response to DNA Replication Stress. <i>PLoS ONE</i> , 2010, 5, e10651.	1.1	11

#	ARTICLE	IF	CITATIONS
55	The Ccr4-Pop2-NOT mRNA Deadenylation Contributes to Septin Organization in <i>Saccharomyces cerevisiae</i> . <i>Genetics</i> , 2009, 182, 955-966.	1.2	23
56	Yeast Gal4: a transcriptional paradigm revisited. <i>EMBO Reports</i> , 2006, 7, 496-499.	2.0	163
57	SQ/TQ cluster domains: concentrated ATM/ATR kinase phosphorylation site regions in DNA-damage-response proteins. <i>BioEssays</i> , 2005, 27, 397-407.	1.2	182
58	Mitochondrial dysfunction enhances Gal4-dependent transcription. <i>FEMS Microbiology Letters</i> , 2005, 253, 207-213.	0.7	2
59	Ccr4-Not Complex mRNA Deadenylation Activity Contributes to DNA Damage Responses in <i>Saccharomyces cerevisiae</i> . <i>Genetics</i> , 2005, 169, 65-75.	1.2	47
60	Patterns that Define the Four Domains Conserved in Known and Novel Isoforms of the Protein Import Receptor Tom20. <i>Journal of Molecular Biology</i> , 2005, 347, 81-93.	2.0	53
61	Protein hijacking. <i>Cancer Cell</i> , 2004, 5, 107-108.	7.7	10
62	The Retinoblastoma Family of Proteins Directly Represses Transcription in <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 8797-8801.	1.6	5
63	The yeast protein Xtc1 functions as a direct transcriptional repressor. <i>Nucleic Acids Research</i> , 2002, 30, 2358-2364.	6.5	4
64	Interorganellar Communication. <i>Journal of Biological Chemistry</i> , 2001, 276, 4020-4027.	1.6	190