

# Alex Andrianopoulos

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8298410/publications.pdf>

Version: 2024-02-01

57  
papers

3,306  
citations

159585

30  
h-index

161849

54  
g-index

59  
all docs

59  
docs citations

59  
times ranked

3548  
citing authors

#	ARTICLE	IF	CITATIONS
1	mpeg1 promoter transgenes direct macrophage-lineage expression in zebrafish. <i>Blood</i> , 2011, 117, e49-e56.	1.4	900
2	Fungal dimorphism: the switch from hyphae to yeast is a specialized morphogenetic adaptation allowing colonization of a host. <i>FEMS Microbiology Reviews</i> , 2015, 39, 797-811.	8.6	186
3	<i>Saccharomyces cerevisiae</i> TEC1 is required for pseudohyphal growth. <i>Molecular Microbiology</i> , 1996, 19, 1255-1263.	2.5	172
4	Characterization of the <i>Aspergillus nidulans nmrA</i> Gene Involved in Nitrogen Metabolite Repression. <i>Journal of Bacteriology</i> , 1998, 180, 1973-1977.	2.2	143
5	Thermally Dimorphic Human Fungal Pathogens—Polyphyletic Pathogens with a Convergent Pathogenicity Trait. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2015, 5, a019794.	6.2	103
6	Control of morphogenesis and actin localization by the <i>Penicillium marneffei</i> RACHomolog. <i>Journal of Cell Science</i> , 2003, 116, 1249-1260.	2.0	97
7	The Ras and Rho GTPases genetically interact to co-ordinately regulate cell polarity during development in <i>Penicillium marneffei</i> . <i>Molecular Microbiology</i> , 2005, 55, 1487-1501.	2.5	96
8	An STE12 Homolog From the Asexual, Dimorphic Fungus <i>Penicillium marneffei</i> Complements the Defect in Sexual Development of an <i>Aspergillus nidulans steA</i> Mutant. <i>Genetics</i> , 2001, 157, 1003-1014.	2.9	94
9	Control of morphogenesis in the human fungal pathogen <i>Penicillium marneffei</i> . <i>International Journal of Medical Microbiology</i> , 2002, 292, 331-347.	3.6	93
10	The <i>abaA</i> homologue of <i>Penicillium marneffei</i> participates in two developmental programmes: conidiation and dimorphic growth. <i>Molecular Microbiology</i> , 2002, 38, 1034-1047.	2.5	81
11	The CDC42 Homolog of the Dimorphic Fungus <i>Penicillium marneffei</i> Is Required for Correct Cell Polarization during Growth but Not Development. <i>Journal of Bacteriology</i> , 2001, 183, 3447-3457.	2.2	79
12	<i>FacB</i> , the <i>Aspergillus nidulans</i> activator of acetate utilization genes, binds dissimilar DNA sequences. <i>EMBO Journal</i> , 1998, 17, 2042-2054.	7.8	77
13	A basic helix-loop-helix protein with similarity to the fungal morphological regulators, <i>Phd1p</i> , <i>Efg1p</i> and <i>StuA</i> , controls conidiation but not dimorphic growth in <i>Penicillium marneffei</i> . <i>Molecular Microbiology</i> , 2002, 44, 621-631.	2.5	60
14	<i>TupA</i> , the <i>Penicillium marneffei</i> <i>Tup1p</i> homologue, represses both yeast and spore development. <i>Molecular Microbiology</i> , 2003, 48, 85-94.	2.5	60
15	The two-component histidine kinases <i>DrkA</i> and <i>SlnA</i> are required for <i>in vivo</i> growth in the human pathogen <i>Penicillium marneffei</i> . <i>Molecular Microbiology</i> , 2011, 82, 1164-1184.	2.5	60
16	Macrophages protect <i>Talaromyces marneffei</i> conidia from myeloperoxidase-dependent neutrophil fungicidal activity during infection establishment in vivo. <i>PLoS Pathogens</i> , 2018, 14, e1007063.	4.7	60
17	A global call for talaromycosis to be recognised as a neglected tropical disease. <i>The Lancet Global Health</i> , 2021, 9, e1618-e1622.	6.3	52
18	The G-Protein $\beta$ -Subunit <i>GasC</i> Plays a Major Role in Germination in the Dimorphic Fungus <i>Penicillium marneffei</i> . <i>Genetics</i> , 2003, 164, 487-499.	2.9	51

#	ARTICLE	IF	CITATIONS
19	G-Protein Signaling Mediates Asexual Development at 25Â°C but Has No Effect on Yeast-Like Growth at 37Â°C in the Dimorphic Fungus <i>Penicillium marneffei</i> . <i>Eukaryotic Cell</i> , 2002, 1, 440-447.	3.4	47
20	A p21-Activated Kinase Is Required for Conidial Germination in <i>Penicillium marneffei</i> . <i>PLoS Pathogens</i> , 2007, 3, e162.	4.7	47
21	Ste20-related kinases: effectors of signaling and morphogenesis in fungi. <i>Trends in Microbiology</i> , 2011, 19, 400-410.	7.7	47
22	Morphogenetic Circuitry Regulating Growth and Development in the Dimorphic Pathogen <i>Penicillium marneffei</i> . <i>Eukaryotic Cell</i> , 2013, 12, 154-160.	3.4	45
23	Clonality Despite Sex: The Evolution of Host-Associated Sexual Neighborhoods in the Pathogenic Fungus <i>Penicillium marneffei</i> . <i>PLoS Pathogens</i> , 2012, 8, e1002851.	4.7	44
24	Intracellular Growth Is Dependent on Tyrosine Catabolism in the Dimorphic Fungal Pathogen <i>Penicillium marneffei</i> . <i>PLoS Pathogens</i> , 2015, 11, e1004790.	4.7	44
25	Developmental regulation of the glyoxylate cycle in the human pathogen <i>Penicillium marneffei</i> . <i>Molecular Microbiology</i> , 2006, 62, 1725-1738.	2.5	43
26	<i>KdmA</i> , a histone <i>H3</i> demethylase with bipartite function, differentially regulates primary and secondary metabolism in <i>Aspergillus nidulans</i> . <i>Molecular Microbiology</i> , 2015, 96, 839-860.	2.5	43
27	Tools for high efficiency genetic manipulation of the human pathogen <i>Penicillium marneffei</i> . <i>Fungal Genetics and Biology</i> , 2012, 49, 772-778.	2.1	42
28	In Vivo Yeast Cell Morphogenesis Is Regulated by a p21-Activated Kinase in the Human Pathogen <i>Penicillium marneffei</i> . <i>PLoS Pathogens</i> , 2009, 5, e1000678.	4.7	35
29	<i>HgrA</i> is necessary and sufficient to drive hyphal growth in the dimorphic pathogen <i>Penicillium marneffei</i> . <i>Molecular Microbiology</i> , 2013, 88, 998-1014.	2.5	35
30	The <i>pbrB</i> Gene Encodes a Laccase Required for DHN-Melanin Synthesis in Conidia of <i>Talaromyces (Penicillium) marneffei</i> . <i>PLoS ONE</i> , 2015, 10, e0122728.	2.5	35
31	Genome Sequence of the AIDS-Associated Pathogen <i>Penicillium marneffei</i> (ATCC18224) and Its Near Taxonomic Relative <i>Talaromyces stipitatus</i> (ATCC10500). <i>Genome Announcements</i> , 2015, 3, .	0.8	29
32	The RFX Protein RfxA Is an Essential Regulator of Growth and Morphogenesis in <i>Penicillium marneffei</i> . <i>Eukaryotic Cell</i> , 2010, 9, 578-591.	3.4	26
33	Cell-Type-Specific Transcriptional Profiles of the Dimorphic Pathogen <i>Penicillium marneffei</i> Reflect Distinct Reproductive, Morphological, and Environmental Demands. <i>G3: Genes, Genomes, Genetics</i> , 2013, 3, 1997-2014.	1.8	25
34	The <i>Aspergillus nidulans rcoA</i> Gene Is Required for <i>veA</i> -Dependent Sexual Development. <i>Genetics</i> , 2006, 174, 1685-1688.	2.9	23
35	<i>AreA</i> controls nitrogen source utilisation during both growth programs of the dimorphic fungus <i>Penicillium marneffei</i> . <i>Fungal Biology</i> , 2012, 116, 145-154.	2.5	21
36	<i>Talaromyces marneffei</i> laccase modifies THP-1 macrophage responses. <i>Virulence</i> , 2016, 7, 702-717.	4.4	20

#	ARTICLE	IF	CITATIONS
37	Î2-glucanâ€“dependent shuttling of conidia from neutrophils to macrophages occurs during fungal infection establishment. <i>PLoS Biology</i> , 2019, 17, e3000113.	5.6	20
38	A genome-wide analysis of carbon catabolite repression in <i>Schizosaccharomyces pombe</i> . <i>BMC Genomics</i> , 2019, 20, 251.	2.8	20
39	Extensive Metabolic Remodeling Differentiates Non-pathogenic and Pathogenic Growth Forms of the Dimorphic Pathogen <i>Talaromyces marneffe</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 368.	3.9	18
40	The Fungal Type II Myosin in <i>Penicillium marneffe</i> , MyoB, Is Essential for Chitin Deposition at Nascent Septation Sites but Not Actin Localization. <i>Eukaryotic Cell</i> , 2011, 10, 302-312.	3.4	17
41	Identification of amdX, a new Cys-2-His-2 (C2H2) zinc-finger gene involved in the regulation of the amdS gene of <i>Aspergillus nidulans</i> . <i>Molecular Microbiology</i> , 1997, 23, 591-602.	2.5	16
42	Conditional lethal disruption of TATA-binding protein gene in <i>Penicillium marneffe</i> . <i>Fungal Genetics and Biology</i> , 2005, 42, 893-903.	2.1	15
43	Two-Component Signaling Regulates Osmotic Stress Adaptation via SskA and the High-Osmolarity Glycerol MAPK Pathway in the Human Pathogen <i>Talaromyces marneffe</i> . <i>MSphere</i> , 2016, 1, .	2.9	14
44	Differentially regulated highâ€“affinity iron assimilation systems support growth of the various cell types in the dimorphic pathogen <i>Talaromyces marneffe</i> . <i>Molecular Microbiology</i> , 2016, 102, 715-737.	2.5	11
45	Reproductive competence: a recurrent logic module in eukaryotic development. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20130819.	2.6	8
46	A unique aspartyl protease gene expansion in <i>Talaromyces marneffe</i> plays a role in growth inside host phagocytes. <i>Virulence</i> , 2019, 10, 277-291.	4.4	8
47	Antifungal Activity and Molecular Mechanisms of Partial Purified Antifungal Proteins from <i>Rhinacanthus nasutus</i> against <i>Talaromyces marneffe</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2020, 6, 333.	3.5	8
48	Laboratory Maintenance and Growth of <i>Talaromyces marneffe</i> . <i>Current Protocols in Microbiology</i> , 2020, 56, e97.	6.5	6
49	Organism-wide studies into pathogenicity and morphogenesis in <i>Talaromyces marneffe</i> . <i>Future Microbiology</i> , 2016, 11, 511-526.	2.0	5
50	Calcineurin A Is Essential in the Regulation of Asexual Development, Stress Responses and Pathogenesis in <i>Talaromyces marneffe</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 3094.	3.5	5
51	The novel Dbl homology/BAR domain protein, MsgA, of <i>Talaromyces marneffe</i> regulates yeast morphogenesis during growth inside host cells. <i>Scientific Reports</i> , 2021, 11, 2334.	3.3	5
52	The Biology of the Thermally Dimorphic Fungal Pathogen <i>Penicillium marneffe</i> . , 2007, , 213-226.		5
53	Signaling Pathways in the Dimorphic Human Fungal Pathogen <i>Penicillium marneffe</i> . , 0, , 441-454.		4
54	A Plastic Vegetative Growth Threshold Governs Reproductive Capacity in <i>Aspergillus nidulans</i> . <i>Genetics</i> , 2016, 204, 1161-1175.	2.9	2

#	ARTICLE	IF	CITATIONS
55	Talaromyces marneffei simA Encodes a Fungal Cytochrome P450 Essential for Survival in Macrophages. <i>MSphere</i> , 2018, 3, .	2.9	2
56	Adaptation to Industrial Stressors Through Genomic and Transcriptional Plasticity in a Bioethanol Producing Fission Yeast Isolate. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 1375-1391.	1.8	1
57	Morphogenesis and pathogenesis: control of cell identity in a dimorphic pathogen. <i>Microbiology Australia</i> , 2015, 36, 95.	0.4	0