

Alex Andrianopoulos

List of Publications by Year in descending order

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57
papers

3,306
citations

159585

30
h-index

161849

54
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59
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59
docs citations

59
times ranked

3548
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | The novel Dbl homology/BAR domain protein, MsgA, of <i>Talaromyces marneffei</i> regulates yeast morphogenesis during growth inside host cells. <i>Scientific Reports</i> , 2021, 11, 2334. | 3.3 | 5 |
| 2 | 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 |
| 3 | Antifungal Activity and Molecular Mechanisms of Partial Purified Antifungal Proteins from <i>Rhinacanthus nasutus</i> against <i>Talaromyces marneffei</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2020, 6, 333. | 3.5 | 8 |
| 4 | Laboratory Maintenance and Growth of <i>Talaromyces marneffei</i> . <i>Current Protocols in Microbiology</i> , 2020, 56, e97. | 6.5 | 6 |
| 5 | 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 |
| 6 | β -glucan α -dependent shuttling of conidia from neutrophils to macrophages occurs during fungal infection establishment. <i>PLoS Biology</i> , 2019, 17, e3000113. | 5.6 | 20 |
| 7 | A unique aspartyl protease gene expansion in <i>Talaromyces marneffei</i> plays a role in growth inside host phagocytes. <i>Virulence</i> , 2019, 10, 277-291. | 4.4 | 8 |
| 8 | A genome-wide analysis of carbon catabolite repression in <i>Schizosaccharomyces pombe</i> . <i>BMC Genomics</i> , 2019, 20, 251. | 2.8 | 20 |
| 9 | Calcineurin A Is Essential in the Regulation of Asexual Development, Stress Responses and Pathogenesis in <i>Talaromyces marneffei</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 3094. | 3.5 | 5 |
| 10 | <i>Talaromyces marneffei</i> simA Encodes a Fungal Cytochrome P450 Essential for Survival in Macrophages. <i>MSphere</i> , 2018, 3, . | 2.9 | 2 |
| 11 | 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 |
| 12 | Extensive Metabolic Remodeling Differentiates Non-pathogenic and Pathogenic Growth Forms of the Dimorphic Pathogen <i>Talaromyces marneffei</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 368. | 3.9 | 18 |
| 13 | <i>Talaromyces marneffei</i> laccase modifies THP-1 macrophage responses. <i>Virulence</i> , 2016, 7, 702-717. | 4.4 | 20 |
| 14 | Organism-wide studies into pathogenicity and morphogenesis in <i>Talaromyces marneffei</i> . <i>Future Microbiology</i> , 2016, 11, 511-526. | 2.0 | 5 |
| 15 | Differentially regulated high-affinity iron assimilation systems support growth of the various cell types in the dimorphic pathogen <i>Talaromyces marneffei</i> . <i>Molecular Microbiology</i> , 2016, 102, 715-737. | 2.5 | 11 |
| 16 | A Plastic Vegetative Growth Threshold Governs Reproductive Capacity in <i>Aspergillus nidulans</i> . <i>Genetics</i> , 2016, 204, 1161-1175. | 2.9 | 2 |
| 17 | Two-Component Signaling Regulates Osmotic Stress Adaptation via SskA and the High-Osmolarity Glycerol MAPK Pathway in the Human Pathogen <i>Talaromyces marneffei</i> . <i>MSphere</i> , 2016, 1, . | 2.9 | 14 |
| 18 | <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 |

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|----|---|-----|-----------|
| 19 | 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 |
| 20 | 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 |
| 21 | 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 |
| 22 | Morphogenesis and pathogenesis: control of cell identity in a dimorphic pathogen. <i>Microbiology Australia</i> , 2015, 36, 95. | 0.4 | 0 |
| 23 | 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 |
| 24 | The <i>pbrB</i> Gene Encodes a Laccase Required for DHN-Melanin Synthesis in Conidia of <i>Talaromyces</i> (<i>Penicillium</i>) <i>marneffei</i> . <i>PLoS ONE</i> , 2015, 10, e0122728. | 2.5 | 35 |
| 25 | 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 |
| 26 | <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 |
| 27 | 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 |
| 28 | 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 |
| 29 | 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 |
| 30 | 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 |
| 31 | <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 |
| 32 | <i>Ste20</i> -related kinases: effectors of signaling and morphogenesis in fungi. <i>Trends in Microbiology</i> , 2011, 19, 400-410. | 7.7 | 47 |
| 33 | 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 |
| 34 | The Fungal Type II Myosin in <i>Penicillium marneffei</i> , <i>MyoB</i> , Is Essential for Chitin Deposition at Nascent Septation Sites but Not Actin Localization. <i>Eukaryotic Cell</i> , 2011, 10, 302-312. | 3.4 | 17 |
| 35 | <i>mpeg1</i> promoter transgenes direct macrophage-lineage expression in zebrafish. <i>Blood</i> , 2011, 117, e49-e56. | 1.4 | 900 |
| 36 | The RFX Protein <i>RfxA</i> Is an Essential Regulator of Growth and Morphogenesis in <i>Penicillium marneffei</i> . <i>Eukaryotic Cell</i> , 2010, 9, 578-591. | 3.4 | 26 |

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|----|---|-----|-----------|
| 37 | 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 |
| 38 | A p21-Activated Kinase Is Required for Conidial Germination in <i>Penicillium marneffei</i> . <i>PLoS Pathogens</i> , 2007, 3, e162. | 4.7 | 47 |
| 39 | The Biology of the Thermally Dimorphic Fungal Pathogen <i>Penicillium marneffei</i> . , 2007, , 213-226. | | 5 |
| 40 | 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 |
| 41 | The <i>Aspergillus nidulans</i> <i>rcoA</i> Gene Is Required for <i>veA</i> -Dependent Sexual Development. <i>Genetics</i> , 2006, 174, 1685-1688. | 2.9 | 23 |
| 42 | 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 |
| 43 | Conditional lethal disruption of TATA-binding protein gene in <i>Penicillium marneffei</i> . <i>Fungal Genetics and Biology</i> , 2005, 42, 893-903. | 2.1 | 15 |
| 44 | <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 |
| 45 | Control of morphogenesis and actin localization by the <i>Penicillium marneffei</i> <i>RAC</i> homologue. <i>Journal of Cell Science</i> , 2003, 116, 1249-1260. | 2.0 | 97 |
| 46 | The G-Protein β -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 |
| 47 | G-Protein Signaling Mediates Asexual Development at 25°C but Has No Effect on Yeast-Like Growth at 37°C in the Dimorphic Fungus <i>Penicillium marneffei</i> . <i>Eukaryotic Cell</i> , 2002, 1, 440-447. | 3.4 | 47 |
| 48 | 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 |
| 49 | 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 |
| 50 | 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 |
| 51 | The <i>CDC42</i> 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 |
| 52 | An <i>STE12</i> Homolog From the Asexual, Dimorphic Fungus <i>Penicillium marneffei</i> Complements the Defect in Sexual Development of an <i>Aspergillus nidulans</i> <i>steA</i> Mutant. <i>Genetics</i> , 2001, 157, 1003-1014. | 2.9 | 94 |
| 53 | <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 |
| 54 | Characterization of the <i>Aspergillus nidulans</i> <i>nmrA</i> Gene Involved in Nitrogen Metabolite Repression. <i>Journal of Bacteriology</i> , 1998, 180, 1973-1977. | 2.2 | 143 |

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|----|---|-----|-----------|
| 55 | 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 |
| 56 | <i>Saccharomyces cerevisiae</i> TEC1 is required for pseudohyphal growth. <i>Molecular Microbiology</i> , 1996, 19, 1255-1263. | 2.5 | 172 |
| 57 | Signaling Pathways in the Dimorphic Human Fungal Pathogen <i>Penicillium marneffei</i> . , 0, , 441-454. | | 4 |