

Alexandra A Soukup

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

603
citations

1051969

10
h-index

939365

18
g-index

21
all docs

21
docs citations

21
times ranked

905
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Gata2</i> +9.5 enhancer regulates adult hematopoietic stem cell self-renewal and T-cell development. <i>Blood Advances</i> , 2022, 6, 1095-1099.	2.5	5
2	GATA2 deficiency elevates interferon regulatory factor-8 to subvert a progenitor cell differentiation program. <i>Blood Advances</i> , 2022, 6, 1464-1473.	2.5	8
3	Conditionally pathogenic genetic variants of a hematopoietic disease-suppressing enhancer. <i>Science Advances</i> , 2021, 7, eabk3521.	4.7	8
4	PRAM: a novel pooling approach for discovering intergenic transcripts from large-scale RNA sequencing experiments. <i>Genome Research</i> , 2020, 30, 1655-1666.	2.4	2
5	GATA2 +9.5 enhancer: from principles of hematopoiesis to genetic diagnosis in precision medicine. <i>Current Opinion in Hematology</i> , 2020, 27, 163-171.	1.2	11
6	GATA2 Enhancer Modules Governing Hematopoietic Regeneration. <i>Blood</i> , 2020, 136, 15-16.	0.6	0
7	Single-nucleotide human disease mutation inactivates a blood-regenerative GATA2 enhancer. <i>Journal of Clinical Investigation</i> , 2019, 129, 1180-1192.	3.9	47
8	CoIN: co-inducible nitrate expression system for secondary metabolites in <i>Aspergillus nidulans</i> . <i>Fungal Biology and Biotechnology</i> , 2018, 5, 6.	2.5	29
9	An <i>LaeA</i> - and <i>BrlA</i> -Dependent Cellular Network Governs Tissue-Specific Secondary Metabolism in the Human Pathogen <i>Aspergillus fumigatus</i> . <i>MSphere</i> , 2018, 3, .	1.3	58
10	Human leukemia mutations corrupt but do not abrogate GATA-2 function. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E10109-E10118.	3.3	34
11	The <i>Aspergillus nidulans</i> <i>Pbp1</i> homolog is required for normal sexual development and secondary metabolism. <i>Fungal Genetics and Biology</i> , 2017, 100, 13-21.	0.9	8
12	Revitalization of a Forward Genetic Screen Identifies Three New Regulators of Fungal Secondary Metabolism in the Genus <i>Aspergillus</i> . <i>MBio</i> , 2017, 8, .	1.8	47
13	New Aspercryptins, Lipopeptide Natural Products, Revealed by HDAC Inhibition in <i>Aspergillus nidulans</i> . <i>ACS Chemical Biology</i> , 2016, 11, 2117-2123.	1.6	56
14	Enhancing Nonribosomal Peptide Biosynthesis in Filamentous Fungi. <i>Methods in Molecular Biology</i> , 2016, 1401, 149-160.	0.4	12
15	Large-Scale Metabolomics Reveals a Complex Response of <i>Aspergillus nidulans</i> to Epigenetic Perturbation. <i>ACS Chemical Biology</i> , 2015, 10, 1535-1541.	1.6	90
16	<i>VeA</i> and <i>MvIA</i> repression of the cryptic orsellinic acid gene cluster in <i>Aspergillus nidulans</i> involves histone 3 acetylation. <i>Molecular Microbiology</i> , 2013, 89, 963-974.	1.2	37
17	Western Analysis of Histone Modifications (<i>Aspergillus nidulans</i>). <i>Bio-protocol</i> , 2013, 3, .	0.2	4
18	Overexpression of the <i>Aspergillus nidulans</i> histone 4 acetyltransferase <i>EsaA</i> increases activation of secondary metabolite production. <i>Molecular Microbiology</i> , 2012, 86, 314-330.	1.2	116

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19	NosA, a transcription factor important in <i>Aspergillus fumigatus</i> stress and developmental response, rescues the germination defect of a <i>laeA</i> deletion. <i>Fungal Genetics and Biology</i> , 2012, 49, 857-865.	0.9	31