

Mårton Miskei

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10817531/publications.pdf>

Version: 2024-02-01

22
papers

1,427
citations

623734

14
h-index

677142

22
g-index

22
all docs

22
docs citations

22
times ranked

2281
citing authors

#	ARTICLE	IF	CITATIONS
1	The phosphatome of opportunistic pathogen <i>Candida</i> species. <i>Fungal Biology Reviews</i> , 2021, 35, 40-51.	4.7	9
2	Genome-wide mapping of binding sites of the transposase-derived SETMAR protein in the human genome. <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 4032-4041.	4.1	3
3	Sequence-based prediction of protein binding mode landscapes. <i>PLoS Computational Biology</i> , 2020, 16, e1007864.	3.2	41
4	Sequence-Based Prediction of Fuzzy Protein Interactions. <i>Journal of Molecular Biology</i> , 2020, 432, 2289-2303.	4.2	77
5	Two Targets, One Hit: new Anticancer Therapeutics to Prevent Tumorigenesis Without Cardiotoxicity. <i>Frontiers in Pharmacology</i> , 2020, 11, 569955.	3.5	1
6	Comparative genomics reveals high biological diversity and specific adaptations in the industrially and medically important fungal genus <i>Aspergillus</i> . <i>Genome Biology</i> , 2017, 18, 28.	8.8	417
7	Study on the glutathione metabolism of the filamentous fungus <i>Aspergillus nidulans</i> . <i>Acta Microbiologica Et Immunologica Hungarica</i> , 2017, 64, 255-272.	0.8	11
8	Fuzziness enables context dependence of protein interactions. <i>FEBS Letters</i> , 2017, 591, 2682-2695.	2.8	60
9	FuzDB: database of fuzzy complexes, a tool to develop stochastic structure-function relationships for protein complexes and higher-order assemblies. <i>Nucleic Acids Research</i> , 2017, 45, D228-D235.	14.5	96
10	Insights into Adaptations to a Near-Obligate Nematode Endoparasitic Lifestyle from the Finished Genome of <i>Drechmeria coniospora</i> . <i>Scientific Reports</i> , 2016, 6, 23122.	3.3	32
11	Stress tolerances of nullmutants of functionâ€unknown genes encoding menadione stressâ€responsive proteins in <i>Aspergillus nidulans</i> . <i>Journal of Basic Microbiology</i> , 2016, 56, 827-833.	3.3	3
12	Fuzzy complexes: Specific binding without complete folding. <i>FEBS Letters</i> , 2015, 589, 2533-2542.	2.8	177
13	Transcriptome changes initiated by carbon starvation in <i>Aspergillus nidulans</i> . <i>Microbiology (United Kingdom)</i> 157, 1843-1854. doi:10.1099/mic/0/000000.0	1.8	72
14	Protein phosphatase Z modulates oxidative stress response in fungi. <i>Fungal Genetics and Biology</i> , 2012, 49, 708-716.	2.1	26
15	Comparison of transcriptional and translational changes caused by long-term menadione exposure in <i>Aspergillus nidulans</i> . <i>Fungal Genetics and Biology</i> , 2011, 48, 92-103.	2.1	38
16	Molecular Evolution of Phosphoprotein Phosphatases in <i>Drosophila</i> . <i>PLoS ONE</i> , 2011, 6, e22218.	2.5	11
17	Conservation of male-specific expression of novel phosphoprotein phosphatases in <i>Drosophila</i> . <i>Development Genes and Evolution</i> , 2010, 220, 123-128.	0.9	8
18	AtfA bZIP-type transcription factor regulates oxidative and osmotic stress responses in <i>Aspergillus nidulans</i> . <i>Molecular Genetics and Genomics</i> , 2010, 283, 289-303.	2.1	78

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19	The polymorphism of protein phosphatase Z1 gene in <i>Candida albicans</i> . Journal of Basic Microbiology, 2010, 50, S74-82.	3.3	13
20	Annotation of stress-response proteins in the aspergilli. Fungal Genetics and Biology, 2009, 46, S105-S120.	2.1	76
21	The 2008 update of the <i>Aspergillus nidulans</i> genome annotation: A community effort. Fungal Genetics and Biology, 2009, 46, S2-S13.	2.1	99
22	Comparison of gene expression signatures of diamide, H ₂ O ₂ and menadione exposed <i>Aspergillus nidulans</i> cultures – linking genome-wide transcriptional changes to cellular physiology. BMC Genomics, 2005, 6, 182.	2.8	79