

David Moi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3671590/publications.pdf>

Version: 2024-02-01

12
papers

335
citations

1307594

7
h-index

1588992

8
g-index

16
all docs

16
docs citations

16
times ranked

413
citing authors

#	ARTICLE	IF	CITATIONS
1	Citrullination Was Introduced into Animals by Horizontal Gene Transfer from Cyanobacteria. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	16
2	Discovery of archaeal fusexins homologous to eukaryotic HAP2/GCS1 gamete fusion proteins. <i>Nature Communications</i> , 2022, 13, .	12.8	17
3	OMA orthology in 2021: website overhaul, conserved isoforms, ancestral gene order and more. <i>Nucleic Acids Research</i> , 2021, 49, D373-D379.	14.5	137
4	Gene Duplication and Gain in the Trematode <i>Atriophallophorus winterbourni</i> Contributes to Adaptation to Parasitism. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	7
5	Scalable phylogenetic profiling using MinHash uncovers likely eukaryotic sexual reproduction genes. <i>PLoS Computational Biology</i> , 2020, 16, e1007553.	3.2	19
6	A putative origin of the insect chemosensory receptor superfamily in the last common eukaryotic ancestor. <i>ELife</i> , 2020, 9, .	6.0	16
7	Scalable phylogenetic profiling using MinHash uncovers likely eukaryotic sexual reproduction genes. , 2020, 16, e1007553.		0
8	Scalable phylogenetic profiling using MinHash uncovers likely eukaryotic sexual reproduction genes. , 2020, 16, e1007553.		0
9	Scalable phylogenetic profiling using MinHash uncovers likely eukaryotic sexual reproduction genes. , 2020, 16, e1007553.		0
10	Scalable phylogenetic profiling using MinHash uncovers likely eukaryotic sexual reproduction genes. , 2020, 16, e1007553.		0
11	Scalable phylogenetic profiling using MinHash uncovers likely eukaryotic sexual reproduction genes. , 2020, 16, e1007553.		0
12	<i>Arabidopsis</i> HAP2/GCS1 is a gamete fusion protein homologous to somatic and viral fusogens. <i>Journal of Cell Biology</i> , 2017, 216, 571-581.	5.2	93