

Yue Hao

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4820874/publications.pdf>

Version: 2024-02-01

12
papers

390
citations

1040056

9
h-index

1199594

12
g-index

15
all docs

15
docs citations

15
times ranked

553
citing authors

#	ARTICLE	IF	CITATIONS
1	Preferential retention of genes from one parental genome after polyploidy illustrates the nature and scope of the genomic conflicts induced by hybridization. <i>PLoS Genetics</i> , 2018, 14, e1007267.	3.5	78
2	Transcriptome and organellar sequencing highlights the complex origin and diversification of allotetraploid <i>Brassica napus</i> . <i>Nature Communications</i> , 2019, 10, 2878.	12.8	78
3	clustifyr: an R package for automated single-cell RNA sequencing cluster classification. <i>F1000Research</i> , 2020, 9, 223.	1.6	71
4	Diet shifts provoke complex and variable changes in the metabolic networks of the ruminal microbiome. <i>Microbiome</i> , 2017, 5, 60.	11.1	38
5	Genes derived from ancient polyploidy have higher genetic diversity and are associated with domestication in <i>Brassica rapa</i> . <i>New Phytologist</i> , 2021, 230, 372-386.	7.3	26
6	Patterns of Population Variation in Two Paleopolyploid Eudicot Lineages Suggest That Dosage-Based Selection on Homeologs Is Long-Lived. <i>Genome Biology and Evolution</i> , 2018, 10, 999-1011.	2.5	21
7	The contributions from the progenitor genomes of the mesopolyploid Brassiceae are evolutionarily distinct but functionally compatible. <i>Genome Research</i> , 2021, 31, 799-810.	5.5	21
8	Convergent evolution of polyploid genomes from across the eukaryotic tree of life. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	16
9	A Single, Shared Triploidy in Three Species of Parasitic Nematodes. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 225-233.	1.8	13
10	Baby Genomics: Tracing the Evolutionary Changes That Gave Rise to Placentation. <i>Genome Biology and Evolution</i> , 2020, 12, 35-47.	2.5	11
11	Functional characterization of SMN evolution in mouse models of SMA. <i>Scientific Reports</i> , 2019, 9, 9472.	3.3	8
12	POInT: A Tool for Modeling Ancient Polyploidies Using Multiple Polyploid Genomes. <i>Methods in Molecular Biology</i> , 2022, , 81-91.	0.9	1