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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6280636/publications.pdf>

Version: 2024-02-01

18
papers

796
citations

623734

14
h-index

888059

17
g-index

20
all docs

20
docs citations

20
times ranked

1591
citing authors

#	ARTICLE	IF	CITATIONS
1	Antarctica's ecological isolation will be broken by storm-driven dispersal and warming. <i>Nature Climate Change</i> , 2018, 8, 704-708.	18.8	220
2	The Eucalyptus terpene synthase gene family. <i>BMC Genomics</i> , 2015, 16, 450.	2.8	125
3	Whole-genome sequencing to detect mutations associated with resistance to insecticides and Bt proteins in <i>Spodoptera frugiperda</i> . <i>Insect Science</i> , 2021, 28, 627-638.	3.0	61
4	The evolution of foliar terpene diversity in Myrtaceae. <i>Phytochemistry Reviews</i> , 2014, 13, 695-716.	6.5	60
5	Differences in gene expression within a striking phenotypic mosaic Eucalyptus tree that varies in susceptibility to herbivory. <i>BMC Plant Biology</i> , 2013, 13, 29.	3.6	43
6	A phylogenomic approach reveals a low somatic mutation rate in a long-lived plant. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2020, 287, 20192364.	2.6	39
7	The molecular basis of host plant selection in <i>Melaleuca quinquenervia</i> by a successful biological control agent. <i>Phytochemistry</i> , 2010, 71, 1237-1244.	2.9	38
8	Transcriptome analysis of terpene chemotypes of <i>Melaleuca alternifolia</i> across different tissues. <i>Plant, Cell and Environment</i> , 2017, 40, 2406-2425.	5.7	34
9	Global population genomic signature of <i>Spodoptera frugiperda</i> (fall armyworm) supports complex introduction events across the Old World. <i>Communications Biology</i> , 2022, 5, 297.	4.4	34
10	Accuracy of Genomic Prediction for Foliar Terpene Traits in <i>Eucalyptus polybractea</i> . <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 2573-2583.	1.8	28
11	High marker density GWAS provides novel insights into the genomic architecture of terpene oil yield in Eucalyptus. <i>New Phytologist</i> , 2019, 223, 1489-1504.	7.3	27
12	Mosaic Eucalypt Trees Suggest Genetic Control at a Point That Influences Several Metabolic Pathways. <i>Journal of Chemical Ecology</i> , 2012, 38, 914-923.	1.8	21
13	Transcriptome Sequencing of Two Phenotypic Mosaic Eucalyptus Trees Reveals Large Scale Transcriptome Re-Modelling. <i>PLoS ONE</i> , 2015, 10, e0123226.	2.5	18
14	Four terpene synthases contribute to the generation of chemotypes in tea tree (<i>Melaleuca</i>). <i>Tree Physiology</i> , 2017, 37, 150-157.	3.8	17
15	Association genetics of essential oil traits in <i>Eucalyptus loxophleba</i> : explaining variation in oil yield. <i>Molecular Breeding</i> , 2017, 37, 1.	2.1	9
16	Genomic analyses suggest strong population connectivity over large spatial scales of the commercially important baitworm, <i>Australonuphis teres</i> (Onuphidae). <i>Marine and Freshwater Research</i> , 2020, 71, 1549.	1.3	3
17	Correction: Differences in gene expression within a striking phenotypic mosaic Eucalyptus tree that varies in susceptibility to herbivory. <i>BMC Plant Biology</i> , 2013, 13, 57.	3.6	1
18	A predicted novel protein isoform of HOXA9. <i>Leukemia Research</i> , 2019, 82, 7-10.	0.8	1