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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4006894/publications.pdf>

Version: 2024-02-01

16
papers

260
citations

1307594

7
h-index

1125743

13
g-index

17
all docs

17
docs citations

17
times ranked

389
citing authors

#	ARTICLE	IF	CITATIONS
1	The draft genome sequence of cork oak. <i>Scientific Data</i> , 2018, 5, 180069.	5.3	98
2	ChIP-Seq reveals that QsMYB1 directly targets genes involved in lignin and suberin biosynthesis pathways in cork oak (<i>Quercus suber</i>). <i>BMC Plant Biology</i> , 2018, 18, 198.	3.6	50
3	Expression Profiling in <i>Pinus pinaster</i> in Response to Infection with the Pine Wood Nematode <i>Bursaphelenchus xylophilus</i> . <i>Forests</i> , 2017, 8, 279.	2.1	22
4	Comparative Transcriptomic Response of Two <i>Pinus</i> Species to Infection with the Pine Wood Nematode <i>Bursaphelenchus xylophilus</i> . <i>Forests</i> , 2020, 11, 204.	2.1	19
5	CheNER: chemical named entity recognizer. <i>Bioinformatics</i> , 2014, 30, 1039-1040.	4.1	15
6	The Crown Pearl: a draft genome assembly of the European freshwater pearl mussel <i>Margaritifera margaritifera</i> (Linnaeus, 1758). <i>DNA Research</i> , 2021, 28, .	3.4	15
7	Methods for and results from the study of design principles in molecular systems. <i>Mathematical Biosciences</i> , 2011, 231, 3-18.	1.9	13
8	Characterization of the cork formation and production transcriptome in <i>Quercus cerris</i> – <i>Q. suber</i> hybrids. <i>Physiology and Molecular Biology of Plants</i> , 2018, 24, 535-549.	3.1	9
9	Comprehensive Analysis of the Cork Oak (<i>Quercus suber</i>) Transcriptome Involved in the Regulation of Bud Sprouting. <i>Forests</i> , 2017, 8, 486.	2.1	6
10	Biblio-MetReS for user-friendly mining of genes and biological processes in scientific documents. <i>PeerJ</i> , 2014, 2, e276.	2.0	4
11	Ovine footrot in Southern Portugal: Detection of <i>Dichelobacter nodosus</i> and <i>Fusobacterium necrophorum</i> in sheep with different lesion scores. <i>Veterinary Microbiology</i> , 2022, 266, 109339.	1.9	3
12	Kinship Analysis and Pedigree Reconstruction of a Natural Regenerated Cork Oak (<i>Quercus suber</i>) Population. <i>Forests</i> , 2022, 13, 226.	2.1	3
13	<i>Quercus suber</i> Transcriptome Analyses: Identification of Genes and SNPs Related to Cork Quality. , 2022, 11, .		2
14	Database Constraints Applied to Metabolic Pathway Reconstruction Tools. <i>Scientific World Journal</i> , The, 2014, 2014, 1-12.	2.1	1
15	P-Biblio-MetReS, a parallel data mining tool for the reconstruction of molecular networks. , 2013, , .		0
16	Transcriptome Characterization of Different Tissues of Stone Pine (<i>Pinus pinea</i> L.): De Novo Assembly. , 2021, 11, .		0