

TimothÃ©e Flutre

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9330928/publications.pdf>

Version: 2024-02-01

21
papers

12,761
citations

840776

11
h-index

752698

20
g-index

25
all docs

25
docs citations

25
times ranked

32692
citing authors

#	ARTICLE	IF	CITATIONS
1	From cultivar mixtures to allelic mixtures: opposite effects of allelic richness between genotypes and genotype richness in wheat. <i>New Phytologist</i> , 2022, 233, 2573-2584.	7.3	14
2	Across-population genomic prediction in grapevine opens up promising prospects for breeding. <i>Horticulture Research</i> , 2022, 9, .	6.3	9
3	A genome-wide association and prediction study in grapevine deciphers the genetic architecture of multiple traits and identifies genes under many new QTLs. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	18
4	Adoption and Optimization of Genomic Selection To Sustain Breeding for Apricot Fruit Quality. <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 4513-4529.	1.8	11
5	VviUCC1 Nucleotide Diversity, Linkage Disequilibrium and Association with Rachis Architecture Traits in Grapevine. <i>Genes</i> , 2020, 11, 598.	2.4	7
6	A novel high-density grapevine (<i>Vitis vinifera</i> L.) integrated linkage map using GBS in a half-diallel population. <i>Theoretical and Applied Genetics</i> , 2019, 132, 2237-2252.	3.6	38
7	PlantBreedGame: A Serious Game that Puts Students in the Breeder's Seat. <i>Crop Science</i> , 2019, 59, 1374-1375.	1.8	1
8	Quantitative Assessment of Grapevine Wood Colonization by the Dieback Fungus <i>Eutypa lata</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2017, 3, 21.	3.5	11
9	Twenty Tips for High-School Students Engaging in Research with Scientists. <i>Frontiers for Young Minds</i> , 2015, 3, .	0.8	4
10	The Genotype-Tissue Expression (GTEx) pilot analysis: Multitissue gene regulation in humans. <i>Science</i> , 2015, 348, 648-660.	12.6	4,659
11	The epigenomic landscape of African rainforest hunter-gatherers and farmers. <i>Nature Communications</i> , 2015, 6, 10047.	12.8	75
12	A Statistical Framework for Joint eQTL Analysis in Multiple Tissues. <i>PLoS Genetics</i> , 2013, 9, e1003486.	3.5	226
13	The Genotype-Tissue Expression (GTEx) project. <i>Nature Genetics</i> , 2013, 45, 580-585.	21.4	6,815
14	Roadmap for Annotating Transposable Elements in Eukaryote Genomes. <i>Methods in Molecular Biology</i> , 2012, 859, 53-68.	0.9	15
15	The ABO blood group is a trans-species polymorphism in primates. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 18493-18498.	7.1	127
16	Transposable Element Annotation in Completely Sequenced Eukaryote Genomes. <i>Topics in Current Genetics</i> , 2012, , 17-39.	0.7	16
17	TriAnnot: A Versatile and High Performance Pipeline for the Automated Annotation of Plant Genomes. <i>Frontiers in Plant Science</i> , 2012, 3, 5.	3.6	73
18	Considering Transposable Element Diversification in De Novo Annotation Approaches. <i>PLoS ONE</i> , 2011, 6, e16526.	2.5	477

#	ARTICLE	IF	CITATIONS
19	In search of lost trajectories. <i>Mobile Genetic Elements</i> , 2011, 1, 151-154.	1.8	4
20	Pilot scheme for misconduct database. <i>Nature</i> , 2011, 478, 37-37.	27.8	0
21	Extensive synteny conservation of holocentric chromosomes in Lepidoptera despite high rates of local genome rearrangements. <i>Proceedings of the National Academy of Sciences of America</i> , 2010, 107, 7680-7685.	7.1	147