

Christopher D Town

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

2,562
citations

687363

13
h-index

1058476

14
g-index

15
all docs

15
docs citations

15
times ranked

4701
citing authors

#	ARTICLE	IF	CITATIONS
1	Araport11: a complete reannotation of the <i>Arabidopsis thaliana</i> reference genome. <i>Plant Journal</i> , 2017, 89, 789-804.	5.7	925
2	An improved genome release (version Mt4.0) for the model legume <i>Medicago truncatula</i> . <i>BMC Genomics</i> , 2014, 15, 312.	2.8	381
3	The pangenome of an agronomically important crop plant <i>Brassica oleracea</i> . <i>Nature Communications</i> , 2016, 7, 13390.	12.8	375
4	A mosaic monoploid reference sequence for the highly complex genome of sugarcane. <i>Nature Communications</i> , 2018, 9, 2638.	12.8	299
5	Transcript Analysis of Early Nodulation Events in <i>Medicago truncatula</i> . <i>Plant Physiology</i> , 2006, 140, 221-234.	4.8	264
6	MTGD: The <i>Medicago truncatula</i> Genome Database. <i>Plant and Cell Physiology</i> , 2015, 56, e1-e1.	3.1	83
7	Annotating the genome of <i>Medicago truncatula</i> . <i>Current Opinion in Plant Biology</i> , 2006, 9, 122-127.	7.1	47
8	ThaleMine: A Warehouse for <i>Arabidopsis</i> Data Integration and Discovery. <i>Plant and Cell Physiology</i> , 2017, 58, pcw200.	3.1	39
9	Population Gene Introgression and High Genome Plasticity for the Zoonotic Pathogen <i>Streptococcus agalactiae</i> . <i>Molecular Biology and Evolution</i> , 2019, 36, 2572-2590.	8.9	36
10	Dominance and Sexual Dimorphism Pervade the <i>Salix purpurea</i> L. Transcriptome. <i>Genome Biology and Evolution</i> , 2017, 9, 2377-2394.	2.5	35
11	Araport Lives: An Updated Framework for <i>Arabidopsis</i> Bioinformatics. <i>Plant Cell</i> , 2020, 32, 2683-2686.	6.6	28
12	Genomic analysis of 16 Colorado human NL63 coronaviruses identifies a new genotype, high sequence diversity in the N-terminal domain of the spike gene and evidence of recombination. <i>Journal of General Virology</i> , 2012, 93, 2387-2398.	2.9	25
13	Reprogramming of Root Cells during Nitrogen-Fixing Symbiosis Involves Dynamic Polysome Association of Coding and Noncoding RNAs. <i>Plant Cell</i> , 2020, 32, 352-373.	6.6	20
14	Nonadditive gene expression is correlated with nonadditive phenotypic expression in interspecific triploid hybrids of willow (<i>Salix</i> spp.). <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	3
15	Large-Scale DNA Sequencing. , 2005, , 337-351.		2