

Li Lei

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/338787/publications.pdf>

Version: 2024-02-01

12
papers

510
citations

1040056

9
h-index

1372567

10
g-index

16
all docs

16
docs citations

16
times ranked

805
citing authors

#	ARTICLE	IF	CITATIONS
1	Plant Pan-Genomics Comes of Age. <i>Annual Review of Plant Biology</i> , 2021, 72, 411-435.	18.7	56
2	Environmental Association Identifies Candidates for Tolerance to Low Temperature and Drought. G3: Genes, Genomes, Genetics, 2019, 9, 3423-3438.	1.8	18
3	Identification of three new resources of resistance to Fusarium head blight in wheat. <i>Czech Journal of Genetics and Plant Breeding</i> , 2019, 55, 15-19.	0.8	0
4	Development of a Multiparent Population for Genetic Mapping and Allele Discovery in Six-Row Barley. <i>Genetics</i> , 2019, 213, 595-613.	2.9	23
5	Sequence-related amplified polymorphisms (SRAPs) reveal genetic diversity and variation regions in upland cotton (<i>Gossypium hirsutum</i> L.) in China. <i>Crop Breeding and Applied Biotechnology</i> , 2018, 18, 155-160.	0.4	0
6	Comparative Genomics Approaches Accurately Predict Deleterious Variants in Plants. G3: Genes, Genomes, Genetics, 2018, 8, 3321-3329.	1.8	36
7	Reevaluation of Two Quantitative Trait Loci for Type II Resistance to Fusarium Head Blight in Wheat Germplasm PI 672538. <i>Phytopathology</i> , 2017, 107, 92-99.	2.2	14
8	Plant organ evolution revealed by phylotranscriptomics in <i>Arabidopsis thaliana</i> . <i>Scientific Reports</i> , 2017, 7, 7567.	3.3	11
9	Temporal fitness fluctuations in experimental <i>Arabidopsis thaliana</i> populations. <i>PLoS ONE</i> , 2017, 12, e0178990.	2.5	9
10	Barcoding the kingdom Plantae: new PCR primers for ITS regions of plants with improved universality and specificity. <i>Molecular Ecology Resources</i> , 2016, 16, 138-149.	4.8	280
11	Collinearity Analysis and High-Density Genetic Mapping of the Wheat Powdery Mildew Resistance Gene Pm40 in PI 672538. <i>PLoS ONE</i> , 2016, 11, e0164815.	2.5	12
12	Expansion and diversification of the SET domain gene family following whole-genome duplications in <i>Populus trichocarpa</i> . <i>BMC Evolutionary Biology</i> , 2012, 12, 51.	3.2	49