

Wei Hua

List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

1,392
citations

394421

19
h-index

361022

35
g-index

35
all docs

35
docs citations

35
times ranked

1612
citing authors

#	ARTICLE	IF	CITATIONS
1	An improved allele-specific PCR primer design method for SNP marker analysis and its application. <i>Plant Methods</i> , 2012, 8, 34.	4.3	192
2	Natural variation in <i>ARF18</i> gene simultaneously affects seed weight and silique length in polyploid rapeseed. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E5123-32.	7.1	185
3	Knockout of two <i>BnaMAX1</i> homologs by CRISPR/Cas9-targeted mutagenesis improves plant architecture and increases yield in rapeseed (<i>Brassica napus</i> L.). <i>Plant Biotechnology Journal</i> , 2020, 18, 644-654.	8.3	117
4	Selection and evaluation of novel reference genes for quantitative reverse transcription PCR (qRT-PCR) based on genome and transcriptome data in <i>Brassica napus</i> L.. <i>Gene</i> , 2014, 538, 113-122.	2.2	111
5	A New Light on Photosystem II Maintenance in Oxygenic Photosynthesis. <i>Frontiers in Plant Science</i> , 2019, 10, 975.	3.6	72
6	Genomic insights into the origin, domestication and diversification of <i>Brassica juncea</i> . <i>Nature Genetics</i> , 2021, 53, 1392-1402.	21.4	66
7	Identification of stable QTLs for seed oil content by combined linkage and association mapping in <i>Brassica napus</i> . <i>Plant Science</i> , 2016, 252, 388-399.	3.6	63
8	Genome-Wide Association Study Reveals Candidate Genes for Control of Plant Height, Branch Initiation Height and Branch Number in Rapeseed (<i>Brassica napus</i> L.). <i>Frontiers in Plant Science</i> , 2017, 8, 1246.	3.6	63
9	Genetic analysis on oil content in rapeseed (<i>Brassica napus</i> L.). <i>Euphytica</i> , 2010, 173, 17-24.	1.2	58
10	PTGBase: an integrated database to study tandem duplicated genes in plants. <i>Database: the Journal of Biological Databases and Curation</i> , 2015, 2015, .	3.0	46
11	Systematic Analysis of Hsf Family Genes in the <i>Brassica napus</i> Genome Reveals Novel Responses to Heat, Drought and High CO ₂ Stresses. <i>Frontiers in Plant Science</i> , 2017, 8, 1174.	3.6	43
12	Complete chloroplast genome sequence of rapeseed (<i>Brassica napus</i> L.) and its evolutionary implications. <i>Genetic Resources and Crop Evolution</i> , 2011, 58, 875-887.	1.6	34
13	A Novel Chimeric Mitochondrial Gene Confers Cytoplasmic Effects on Seed Oil Content in Polyploid Rapeseed (<i>Brassica napus</i>). <i>Molecular Plant</i> , 2019, 12, 582-596.	8.3	26
14	CRISPR/Cas9-targeted mutagenesis of the <i>BnaA03.BP</i> gene confers semi-dwarf and compact architecture to rapeseed (<i>Brassica napus</i> L.). <i>Plant Biotechnology Journal</i> , 2021, 19, 2383-2385.	8.3	26
15	Three <i>BnaIAA7</i> homologs are involved in auxin/brassinosteroid-mediated plant morphogenesis in rapeseed (<i>Brassica napus</i> L.). <i>Plant Cell Reports</i> , 2019, 38, 883-897.	5.6	25
16	Fine-mapping and transcriptome analysis of a candidate gene controlling plant height in <i>Brassica napus</i> L.. <i>Biotechnology for Biofuels</i> , 2020, 13, 42.	6.2	25
17	Genome-wide identification and expression analysis of CaM/CML genes in <i>Brassica napus</i> under abiotic stress. <i>Journal of Plant Physiology</i> , 2020, 255, 153251.	3.5	24
18	Important photosynthetic contribution of silique wall to seed yield-related traits in <i>Arabidopsis thaliana</i> . <i>Photosynthesis Research</i> , 2018, 137, 493-501.	2.9	22

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19	Development and application of single nucleotide polymorphism markers in the polyploid <i>Brassica napus</i> by 454 sequencing of expressed sequence tags. <i>Plant Breeding</i> , 2012, 131, 293-299.	1.9	21
20	Effects of specific organs on seed oil accumulation in <i>Brassica napus</i> L.. <i>Plant Science</i> , 2014, 227, 60-68.	3.6	20
21	Genome-wide screening and analysis of imprinted genes in rapeseed (<i>Brassica napus</i> L.) endosperm. <i>DNA Research</i> , 2018, 25, 629-640.	3.4	18
22	Systematic Analysis of the DNA Methylase and Demethylase Gene Families in Rapeseed (<i>Brassica napus</i> L.) and Their Expression Variations After Salt and Heat stresses. <i>International Journal of Molecular Sciences</i> , 2020, 21, 953.	4.1	16
23	Integrated strategies for increasing rapeseed yield. <i>Trends in Plant Science</i> , 2022, 27, 742-745.	8.8	16
24	Overexpression of CHMP7 from rapeseed and <i>Arabidopsis</i> causes dwarfism and premature senescence in <i>Arabidopsis</i> . <i>Journal of Plant Physiology</i> , 2016, 204, 16-26.	3.5	14
25	Genome-Wide Identification and Characterization of FBA Gene Family in Polyploid Crop <i>Brassica napus</i> . <i>International Journal of Molecular Sciences</i> , 2019, 20, 5749.	4.1	14
26	Integrative analysis of GWAS and transcriptome to reveal novel loci regulation flowering time in semi-winter rapeseed. <i>Plant Science</i> , 2021, 310, 110980.	3.6	14
27	Effective Extraction and Assembly Methods for Simultaneously Obtaining Plastid and Mitochondrial Genomes. <i>PLoS ONE</i> , 2014, 9, e108291.	2.5	13
28	Overexpression of <i>BnaAOX1b</i> Confers Tolerance to Osmotic and Salt Stress in Rapeseed. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 3501-3511.	1.8	11
29	Nitric oxide affects seed oil accumulation and fatty acid composition through protein S-nitrosation. <i>Journal of Experimental Botany</i> , 2021, 72, 385-397.	4.8	7
30	A large-scale population based organelle pan-genomes construction and phylogeny analysis reveal the genetic diversity and the evolutionary origins of chloroplast and mitochondrion in <i>Brassica napus</i> L.. <i>BMC Genomics</i> , 2022, 23, 339.	2.8	7
31	Genome-Wide Identification of the TIFY Gene Family in Brassiceae and Its Potential Association with Heavy Metal Stress in Rapeseed. <i>Plants</i> , 2022, 11, 667.	3.5	6
32	An integrated omics analysis reveals the gene expression profiles of maize, castor bean, and rapeseed for seed oil biosynthesis. <i>BMC Plant Biology</i> , 2022, 22, 153.	3.6	6
33	Abscisic Acid Improves Linoleic Acid Accumulation Possibly by Promoting Expression of <i>EgFAD2</i> and Other Fatty Acid Biosynthesis Genes in Oil Palm Mesocarp. <i>Frontiers in Plant Science</i> , 2021, 12, 748130.	3.6	5
34	Genome-Wide Identification and Evolutionary Analysis of the Fruit-Weight 2.2-Like Gene Family in Polyploid Oilseed Rape (<i>Brassica napus</i> L.). <i>DNA and Cell Biology</i> , 2020, 39, 766-782.	1.9	3
35	Stress-induced higher vein density in the C3-C4 intermediate <i>Moricandia suffruticosa</i> under drought and heat stress. <i>Journal of Experimental Botany</i> , 2022, 73, 6334-6351.	4.8	3