Huolin Shen

List of Publications by Year in descending order

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687363 713466 24 478 13 21 citations h-index g-index papers 26 26 26 422 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	Mapping of CaPP2C35 involved in the formation of light-green immature pepper (Capsicum annuum L.) fruits via GWAS and BSA. Theoretical and Applied Genetics, 2022, 135, 591-604.	3.6	8
2	Identification of the Regulatory Genes of UV-B-Induced Anthocyanin Biosynthesis in Pepper Fruit. International Journal of Molecular Sciences, 2022, 23, 1960.	4.1	14
3	Loci underlying leaf agronomic traits identified by re-sequencing celery accessions based on an assembled genome. IScience, 2022, 25, 104565.	4.1	6
4	Complete Mitochondrial Genome Sequence and Identification of a Candidate Gene Responsible for Cytoplasmic Male Sterility in Celery (Apium graveolens L.). International Journal of Molecular Sciences, 2021, 22, 8584.	4.1	12
5	Hairiness Gene Regulated Multicellular, Non-Glandular Trichome Formation in Pepper Species. Frontiers in Plant Science, 2021, 12, 784755.	3.6	1
6	DNA methylation is involved in the regulation of pepper fruit ripening and interacts with phytohormones. Journal of Experimental Botany, 2020, 71, 1928-1942.	4.8	54
7	Phenotypic, genetic, and molecular function of msc-2, a genic male sterile mutant in pepper (Capsicum) Tj ETQq1	1,0,78431 3.6	4 rgBT /Cwe
8	Fine mapping of the Ca3GT gene controlling anthocyanin biosynthesis in mature unripe fruit of Capsicum annuum L Theoretical and Applied Genetics, 2020, 133, 2729-2742.	3.6	18
9	Complementary Transcriptomic and Proteomic Analysis Reveals a Complex Network Regulating Pollen Abortion in GMS (msc-1) Pepper (Capsicum annuum L.). International Journal of Molecular Sciences, 2019, 20, 1789.	4.1	10
10	Candidate Gene Selection for Cytoplasmic Male Sterility in Pepper (Capsicum annuum L.) through Whole Mitochondrial Genome Sequencing. International Journal of Molecular Sciences, 2019, 20, 578.	4.1	24
11	Genome-Wide Correlation of 36 Agronomic Traits in the 287 Pepper (Capsicum) Accessions Obtained from the SLAF-seq-Based GWAS. International Journal of Molecular Sciences, 2019, 20, 5675.	4.1	40
12	Pedigree-Based Deciphering of Genome-Wide Conserved Patterns in an Elite Potato Parental Line. Frontiers in Plant Science, 2018, 9, 690.	3.6	10
13	The Aborted Microspores (AMS)-Like Gene Is Required for Anther and Microspore Development in Pepper (Capsicum annuum L.). International Journal of Molecular Sciences, 2018, 19, 1341.	4.1	23
14	Identification of candidate genes underlying genic male-sterile msc-1 locus via genome resequencing in Capsicum annuum L Theoretical and Applied Genetics, 2018, 131, 1861-1872.	3.6	26
15	Proteomic analysis reveals strong mitochondrial involvement in cytoplasmic male sterility of pepper () Tj ETQq1 1 (0,784314 2.4	rgBT /Overlo
16	Mutation in the gene encoding <i>1â€aminocyclopropaneâ€1â€carboxylate synthase 4</i> (<i>CitACS4</i>) led to andromonoecy in watermelon. Journal of Integrative Plant Biology, 2016, 58, 762-765.	8.5	34
17	Molecular mapping of a gene conferring resistance to Phytophthora capsici Leonian race 2 in pepper line PI201234 (Capsicum annuum L.). Molecular Breeding, 2016, 36, 1.	2.1	23
18	PAP3 Regulates Stamen but Not Petal Development in Capsicum annuum L Horticultural Plant Journal, 2016, 2, 91-96.	5.0	4

#	Article	IF	CITATION
19	A Truncated F-Box Protein Confers the Dwarfism in Cucumber. Journal of Genetics and Genomics, 2016, 43, 223-226.	3.9	27
20	Identification and Expression Analysis of Candidate Genes Associated with Defense Responses to Phytophthora capsici in Pepper Line "Pl 201234― International Journal of Molecular Sciences, 2015, 16, 11417-11438.	4.1	24
21	Inheritance of sex forms in watermelon (Citrullus lanatus). Scientia Horticulturae, 2015, 193, 367-373.	3.6	30
22	Expression Pattern of Class B Gene PAP3 in Flower Development of Pepper. International Journal of Molecular Sciences, 2013, 14, 24643-24655.	4.1	3
23	Mapping of a Heat-Stable Gene for Resistance to Southern Root-Knot Nematode in Solanum lycopersicum. Plant Molecular Biology Reporter, 2013, 31, 352-362.	1.8	15
24	Genetic variation in tomato populations from four breeding programs revealed by single nucleotide polymorphism and simple sequence repeat markers. Scientia Horticulturae, 2009, 122, 6-16.	3.6	29