

Yunqing Cheng

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

256
citations

933447

10
h-index

940533

16
g-index

20
all docs

20
docs citations

20
times ranked

168
citing authors

#	ARTICLE	IF	CITATIONS
1	Vegetative cells may perform nitrogen fixation function under nitrogen deprivation in <i>Anabaena</i> sp. strain PCC 7120 based on genome-wide differential expression analysis. <i>PLoS ONE</i> , 2021, 16, e0248155.	2.5	5
2	Pollen tube in hazel grows intermittently: Role of Ca ²⁺ and expression of auto-inhibited Ca ²⁺ pump. <i>Scientia Horticulturae</i> , 2021, 282, 110032.	3.6	7
3	Genome-Wide Identification of the ARF Gene Family and ARF3 Target Genes Regulating Ovary Initiation in Hazel via CHIP Sequencing. <i>Frontiers in Plant Science</i> , 2021, 12, 715820.	3.6	10
4	Chromosome-Level Genome Assembly and HazelOmics Database Construction Provides Insights Into Unsaturated Fatty Acid Synthesis and Cold Resistance in Hazelnut (<i>Corylus heterophylla</i>). <i>Frontiers in Plant Science</i> , 2021, 12, 766548.	3.6	7
5	Identification of vital candidate microRNA/mRNA pairs regulating ovule development using high-throughput sequencing in hazel. <i>BMC Developmental Biology</i> , 2020, 20, 13.	2.1	11
6	iTRAQ protein profiling reveals candidate proteins regulating ovary and ovule differentiation in pistillate inflorescences after pollination in hazel. <i>Tree Genetics and Genomes</i> , 2019, 15, 1.	1.6	4
7	Whole-Genome Re-Sequencing of <i>Corylus heterophylla</i> Blank-Nut Mutants Reveals Sequence Variations in Genes Associated With Embryo Abortion. <i>Frontiers in Plant Science</i> , 2019, 10, 1465.	3.6	4
8	New insight into ovary abortion during ovary development of hazelnut through a combined proteomic and transcriptomic analysis. <i>Scientia Horticulturae</i> , 2018, 234, 36-48.	3.6	8
9	Identification of genes regulating ovary differentiation after pollination in hazel by comparative transcriptome analysis. <i>BMC Plant Biology</i> , 2018, 18, 84.	3.6	14
10	Construction of an RNAi vector for knockdown of GM-ACS genes in the cotyledonary nodes of soybean. <i>Electronic Journal of Biotechnology</i> , 2017, 26, 40-45.	2.2	0
11	Comparison of phytohormone biosynthesis and signal transduction pathways in developing and abortive hazelnut ovules. <i>Plant Growth Regulation</i> , 2017, 81, 147-157.	3.4	23
12	Transcriptome Analysis and Gene Expression Profiling of Abortive and Developing Ovules during Fruit Development in Hazelnut. <i>PLoS ONE</i> , 2015, 10, e0122072.	2.5	25
13	Analysis of ovary DNA methylation during delayed fertilization in hazel using the methylation-sensitive amplification technique. <i>Acta Physiologiae Plantarum</i> , 2015, 37, 1.	2.1	10
14	The effects of ethylene on the HCl-extractability of trace elements during soybean seed germination. <i>Electronic Journal of Biotechnology</i> , 2015, 18, 333-337.	2.2	2
15	Comparison of ultrastructure, pollen tube growth pattern and starch content in developing and abortive ovaries during the progamic phase in hazel. <i>Frontiers in Plant Science</i> , 2014, 5, 528.	3.6	24
16	Pistillate flower development and pollen tube growth mode during the delayed fertilization stage in <i>Corylus heterophylla</i> Fisch. <i>Plant Reproduction</i> , 2014, 27, 145-152.	2.2	35
17	Temporal changes of disodium fluorescein transport in hazelnut during fruit development stage. <i>Scientia Horticulturae</i> , 2013, 150, 348-353.	3.6	20
18	Construction of ethylene regulatory network based on the phytohormones related gene transcriptome profiling and prediction of transcription factor activities in soybean. <i>Acta Physiologiae Plantarum</i> , 2013, 35, 1303-1317.	2.1	16

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19	The relationship between reproductive growth and blank fruit formation in <i>Corylus heterophylla</i> Fisch. <i>Scientia Horticulturae</i> , 2012, 136, 128-134.	3.6	28
20	Dual RNA Sequencing Analysis of <i>Bacillus amyloliquefaciens</i> and <i>Sclerotinia sclerotiorum</i> During Infection of Soybean Seedlings by <i>S. sclerotiorum</i> Unveils Antagonistic Interactions. <i>Frontiers in Microbiology</i> , 0, 13, .	3.5	3