

Shutian Tao

List of Publications by Year in descending order

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34
papers

1,790
citations

471371

17
h-index

414303

32
g-index

34
all docs

34
docs citations

34
times ranked

1792
citing authors

#	ARTICLE	IF	CITATIONS
1	Cryptochrome-mediated blue-light signal contributes to lignin biosynthesis in stone cells in pear fruit. <i>Plant Science</i> , 2022, 318, 111211.	1.7	17
2	Genome-wide analysis and expression pattern of the PIN gene family during Korla fragrant pear calyx development. <i>Acta Physiologiae Plantarum</i> , 2022, 44, 1.	1.0	2
3	Transcriptome analysis provides new ideas for studying the regulation of glucose-induced lignin biosynthesis in pear calli. <i>BMC Plant Biology</i> , 2022, 22, .	1.6	2
4	Genome-wide association studies provide insights into the genetic determination of fruit traits of pear. <i>Nature Communications</i> , 2021, 12, 1144.	5.8	44
5	PbCSE1 promotes lignification during stone cell development in pear (<i>Pyrus bretschneideri</i>) fruit. <i>Scientific Reports</i> , 2021, 11, 9450.	1.6	10
6	CAD Genes: Genome-Wide Identification, Evolution, and Their Contribution to Lignin Biosynthesis in Pear (<i>Pyrus bretschneideri</i>). <i>Plants</i> , 2021, 10, 1444.	1.6	6
7	Analysis of PRX Gene Family and Its Function on Cell Lignification in Pears (<i>Pyrus bretschneideri</i>). <i>Plants</i> , 2021, 10, 1874.	1.6	4
8	Transcriptome provides potential insights into how calcium affects the formation of stone cell in <i>Pyrus</i> . <i>BMC Genomics</i> , 2021, 22, 831.	1.2	1
9	Candidate proteins involved in the calyx abscission process of “Kuerlexiangli”™ (<i>Pyrus sinkiangensis</i> Yu) identified by iTRAQ analysis. <i>Acta Physiologiae Plantarum</i> , 2020, 42, 1.	1.0	3
10	PbMC1a/1b regulates lignification during stone cell development in pear (<i>Pyrus bretschneideri</i>) fruit. <i>Horticulture Research</i> , 2020, 7, 59.	2.9	20
11	The Î²-amylase PbrBAM3 from pear (<i>Pyrus betulaefolia</i>) regulates soluble sugar accumulation and ROS homeostasis in response to cold stress. <i>Plant Science</i> , 2019, 287, 110184.	1.7	52
12	Genome-wide analyses and expression patterns under abiotic stress of NAC transcription factors in white pear (<i>Pyrus bretschneideri</i>). <i>BMC Plant Biology</i> , 2019, 19, 161.	1.6	41
13	Characterization and Quantification of Polyphenols and Triterpenoids in Thinned Young Fruits of Ten Pear Varieties by UPLC-Q TRAP-MS/MS. <i>Molecules</i> , 2019, 24, 159.	1.7	62
14	Transcriptomic and evolutionary analyses of white pear (<i>Pyrus bretschneideri</i>) Î²-amylase genes reveals their importance for cold and drought stress responses. <i>Gene</i> , 2019, 689, 102-113.	1.0	22
15	The unique evolutionary pattern of the Hydroxyproline-rich glycoproteins superfamily in Chinese white pear (<i>Pyrus bretschneideri</i>). <i>BMC Plant Biology</i> , 2018, 18, 36.	1.6	6
16	The mining and evolutionary investigation of AP2/ERF genes in pear (<i>Pyrus</i>). <i>BMC Plant Biology</i> , 2018, 18, 46.	1.6	41
17	Physiological and Nutritional Responses of Pear Seedlings to Nitrate Concentrations. <i>Frontiers in Plant Science</i> , 2018, 9, 1679.	1.7	33
18	Diversification and independent domestication of Asian and European pears. <i>Genome Biology</i> , 2018, 19, 77.	3.8	149

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19	Calcium treatments promote the aroma volatiles emission of pear (<i>Pyrus ussuriensis</i> ‘Nanguoli’™) fruit during post-harvest ripening process. <i>Scientia Horticulturae</i> , 2017, 215, 102-111.	1.7	27
20	Genome-wide analysis and characterization of molecular evolution of the HCT gene family in pear (<i>Pyrus bretschneideri</i>). <i>Plant Systematics and Evolution</i> , 2017, 303, 71-90.	0.3	23
21	Transcriptome profiling reveals the candidate genes associated with aroma metabolites and emission of pear (<i>Pyrus ussuriensis</i> cv.). <i>Scientia Horticulturae</i> , 2016, 206, 33-42.	1.7	15
22	Genome-wide Annotation and Comparative Analysis of Long Terminal Repeat Retrotransposons between Pear Species of <i>P. bretschneideri</i> and <i>P. Communis</i> . <i>Scientific Reports</i> , 2015, 5, 17644.	1.6	16
23	Mitochondrial dysfunction mediated by cytoplasmic acidification results in pollen tube growth cessation in <i>Pyrus pyrifolia</i> . <i>Physiologia Plantarum</i> , 2015, 153, 603-615.	2.6	18
24	Cinnamate-4-Hydroxylase Gene Is Involved in the Step of Lignin Biosynthesis in Chinese White Pear. <i>Journal of the American Society for Horticultural Science</i> , 2015, 140, 573-579.	0.5	12
25	Evolution of the Aroma Volatiles of Pear Fruits Supplemented with Fatty Acid Metabolic Precursors. <i>Molecules</i> , 2014, 19, 20183-20196.	1.7	41
26	Long-chain base phosphates modulate pollen tube growth via channel-mediated influx of calcium. <i>Plant Journal</i> , 2014, 79, 507-516.	2.8	17
27	The genome of the pear (<i>Pyrus bretschneideri</i> Rehd.). <i>Genome Research</i> , 2013, 23, 396-408.	2.4	832
28	Evaluation of the volatile profile of 33 <i>Pyrus ussuriensis</i> cultivars by HS-SPME with GC-MS. <i>Food Chemistry</i> , 2012, 134, 2367-2382.	4.2	83
29	Effect of mulching systems on fruit quality and phytochemical composition of newly developed strawberry lines. <i>Agricultural and Food Science</i> , 2012, 21, 132-140.	0.3	10
30	Effects of Drought Stress and Rewatering on Physiological Characteristics of Pear Seedling. <i>Ying Yong Yu Huan Jing Sheng Wu Xue Bao = Chinese Journal of Applied and Environmental Biology</i> , 2012, 18, 218.	0.1	0
31	<i>In vitro</i> antifungal activity and mode of action of selected polyphenolic antioxidants on <i>Botrytis cinerea</i> . <i>Archives of Phytopathology and Plant Protection</i> , 2010, 43, 1564-1578.	0.6	30
32	Anatomy, ultrastructure and lignin distribution of stone cells in two <i>Pyrus</i> species. <i>Plant Science</i> , 2009, 176, 413-419.	1.7	138
33	PROFILE OF ANTIOXIDANT ACTIVITIES OF SELECTED STRAWBERRY GENOTYPES. <i>Acta Horticulturae</i> , 2009, , 551-556.	0.1	7
34	MYB1R1 and MYC2 Regulate Δ^3 Fatty Acid Desaturase Involved in ABA-Mediated Suberization in the Russet Skin of a Mutant of ‘Dangshansuli’™ (<i>Pyrus bretschneideri</i> Rehd.). <i>Frontiers in Plant Science</i> , 0, 13, .	1.7	6