Chunxian Chen

List of Publications by Year in descending order

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53 papers 2,095 citations

394421 19 h-index 243625 44 g-index

53 all docs 53 docs citations

53 times ranked 1980 citing authors

#	Article	IF	CITATIONS
1	Population Structure and Phylogeny of Some U.S. Peach Cultivars. Journal of the American Society for Horticultural Science, 2022, 147, 1-6.	1.0	2
2	Assessment of Prunus Rootstock Accessions Using Chloroplast and Nuclear Microsatellites. Journal of the American Society for Horticultural Science, 2022, 147, 95-103.	1.0	O
3	Mating Type Idiomorphs, Heterothallism, and High Genetic Diversity in <i>Venturia carpophila</i> , Cause of Peach Scab. Phytopathology, 2021, 111, 408-424.	2.2	5
4	Genetic relationship and parentages of historical peaches revealed by microsatellite markers. Tree Genetics and Genomes, $2021, 17, 1$.	1.6	3
5	â€~Rich Joy' Peach. Hortscience: A Publication of the American Society for Hortcultural Science, 2020, 55, 591-592.	1.0	1
6	â€~Liberty Joy' Peach. Hortscience: A Publication of the American Society for Hortcultural Science, 2020, 55, 951-952.	1.0	2
7	â€~UF-1013-1': An Infertile Cultivar of Lantana camara. Hortscience: A Publication of the American Society for Hortcultural Science, 2020, 55, 953-958.	1.0	5
8	â€~Crimson Joy' Peach. Hortscience: A Publication of the American Society for Hortcultural Science, 2020, 55, 972-973.	1.0	1
9	Novel Primers and Sampling for PCR Detection of <i>Xylella fastidiosa </i> in Peach. Phytopathology, 2019, 109, 307-317.	2.2	13
10	Effect of a Late Spring Application of Hydrogen Cyanamide on High-Chill Peaches. Agronomy, 2019, 9, 726.	3.0	8
11	New Somatic Hybrid Mandarin Tetraploids Generated by Optimized Protoplast Fusion and Confirmed by Molecular Marker Analysis and Flow Cytometry. Journal of the American Society for Horticultural Science, 2019, 144, 151-163.	1.0	7
12	Mining and characterization of microsatellites from a genome of Venturia carpophila. Mycological Progress, 2018, 17, 885-895.	1.4	6
13	Comparative analysis of juice volatiles in selected mandarins, mandarin relatives and other citrus genotypes. Journal of the Science of Food and Agriculture, 2018, 98, 1124-1131.	3.5	21
14	Juice volatile composition differences between Valencia orange and its mutant Rohde Red Valencia are associated with carotenoid profile differences. Food Chemistry, 2018, 245, 223-232.	8.2	29
15	Pecan Bacterial Leaf Scorch, Caused by Xylella fastidiosa, Is Endemic in Georgia Pecan Orchards. Plant Health Progress, 2018, 19, 284-287.	1.4	5
16	Genetic Diversity and Population Structure Analysis of Citrus Germplasm with Single Nucleotide Polymorphism Markers. Journal of the American Society for Horticultural Science, 2018, 143, 399-408.	1.0	7
17	Characterization of Polymorphic Chloroplast Microsatellites in Prunus Species and Maternal Lineages in Peach Genotypes. Journal of the American Society for Horticultural Science, 2017, 142, 217-224.	1.0	5
18	Identification of QTLs controlling aroma volatiles using a â€Fortune' x â€Murcott' (Citrus reticulata) population. BMC Genomics, 2017, 18, 646.	2.8	35

#	Article	IF	CITATIONS
19	Draft genome sequence of Venturia carpophila, the causal agent of peach scab. Standards in Genomic Sciences, 2017, 12, 68.	1.5	22
20	Draft genome sequence of Fusicladium effusum, cause of pecan scab. Standards in Genomic Sciences, 2016, 11, 36.	1.5	19
21	Comparison of fruit characters and volatile components in peach-to-nectarine mutants. Euphytica, 2016, 209, 409-418.	1.2	5
22	Peach Fruit Set and Buttoning after Spring Frost. Hortscience: A Publication of the American Society for Hortcultural Science, 2016, 51, 816-821.	1.0	14
23	Novel Peach Flower Types in a Segregating Population from â€~Helen Borchers'. Journal of the American Society for Horticultural Science, 2015, 140, 172-177.	1.0	6
24	Cybridization of Grapefruit with â€~Dancy' Mandarin Leads to Improved Fruit Characteristics. Journal of the American Society for Horticultural Science, 2015, 140, 427-435.	1.0	12
25	Identification of genes associated with low furanocoumarin content in grapefruit. Genome, 2014, 57, 537-545.	2.0	5
26	Novel expression patterns of carotenoid pathway-related genes in citrus leaves and maturing fruits. Tree Genetics and Genomes, 2014, 10, 439-448.	1.6	33
27	Sequencing of diverse mandarin, pummelo and orange genomes reveals complex history of admixture during citrus domestication. Nature Biotechnology, 2014, 32, 656-662.	17.5	572
28	Genome-wide characterization and selection of expressed sequence tag simple sequence repeat primers for optimized marker distribution and reliability in peach. Tree Genetics and Genomes, 2014, 10, 1271-1279.	1.6	10
29	Comparison of carotenoid accumulation and biosynthetic gene expression between Valencia and Rohde Red Valencia sweet oranges. Plant Science, 2014, 227, 28-36.	3.6	48
30	Genetic variability among populations of Fusicladium species from different host trees and geographic locations in the USA. Mycological Progress, 2014, 13, 1179.	1.4	8
31	Cytological and molecular characterization of three gametoclones of Citrus clementina. BMC Plant Biology, 2013, 13, 129.	3.6	18
32	Mining of haplotype-based expressed sequence tag single nucleotide polymorphisms in citrus. BMC Genomics, 2013, 14, 746.	2.8	28
33	Differential anatomical responses of tolerant and susceptible citrus species to the infection of †Candidatus Liberibacter asiaticus'. Physiological and Molecular Plant Pathology, 2013, 83, 69-74.	2.5	42
34	Mechanism-based inhibition of human Cytochrome P450-3A activity by grapefruit hybrids having low furanocoumarin content. Xenobiotica, 2012, 42, 1163-1169.	1.1	14
35	Isolation and characterization of a novel anthocyanin-promoting MYBA gene family in Citrus. Tree Genetics and Genomes, 2012, 8, 675-685.	1.6	8
36	Rectification concerning "lsolation and characterization of a novel anthocyanin-promoting MYBA gene family in Citrus― Tree Genetics and Genomes, 2012, 8, 687-687.	1.6	1

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37	Comparative Transcriptional and Anatomical Analyses of Tolerant Rough Lemon and Susceptible Sweet Orange in Response to $\hat{a} \in \mathbb{Z} \setminus \mathbb{Z}$ Candidatus $ a $ Liberibacter asiaticus $\hat{a} \in \mathbb{Z} \setminus \mathbb{Z}$ Infection. Molecular Plant-Microbe Interactions, 2012, 25, 1396-1407.	2.6	80
38	A reference genetic map of C. clementina hort. ex Tan.; citrus evolution inferences from comparative mapping. BMC Genomics, 2012, 13, 593.	2.8	129
39	Citrus genomics. Tree Genetics and Genomes, 2012, 8, 611-626.	1.6	104
40	Expression and phylogenetic analysis of two new lycopene ⟨i⟩β⟨/i⟩â€cyclases from ⟨i⟩Citrus paradisi⟨/i⟩. Physiologia Plantarum, 2011, 141, 1-10.	5. 2	36
41	Comparative iTRAQ proteome and transcriptome analyses of sweet orange infected by " <i>Candidatus</i> Liberibacter asiaticus― Physiologia Plantarum, 2011, 143, 235-245.	5.2	122
42	Characterization of Furanocoumarin Profile and Inheritance Toward Selection of Low Furanocoumarin Seedless Grapefruit Cultivars. Journal of the American Society for Horticultural Science, 2011, 136, 358-363.	1.0	15
43	Lack of Evidence for Transmission of <i>†Candidatus</i> Liberibacter asiaticus†Through Citrus Seed Taken from Affected Fruit. Plant Disease, 2010, 94, 1200-1205.	1.4	30
44	Identification of novel members in sweet orange carotenoid biosynthesis gene families. Tree Genetics and Genomes, 2010, 6, 905-914.	1.6	27
45	Production of New Allotetraploid and Autotetraploid Citrus Breeding Parents: Focus on Zipperskin Mandarins. Hortscience: A Publication of the American Society for Hortcultural Science, 2010, 45, 1160-1163.	1.0	19
46	EST-SSR genetic maps for Citrus sinensis and Poncirus trifoliata. Tree Genetics and Genomes, 2008, 4, 1-10.	1.6	119
47	Characterization of zygotic and nucellar seedlings from sour orange-like citrus rootstock candidates using RAPD and EST-SSR markers. Tree Genetics and Genomes, 2008, 4, 113-124.	1.6	50
48	Origin and frequency of 2n gametes in Citrus sinensis×Poncirus trifoliata and their reciprocal crosses. Plant Science, 2008, 174, 1-8.	3.6	56
49	Verification of Mandarin and Pummelo Somatic Hybrids by Expressed Sequence Tag–Simple Sequence Repeat Marker Analysis. Journal of the American Society for Horticultural Science, 2008, 133, 794-800.	1.0	11
50	Development of pGreen-derived GFP Binary Vectors for Citrus Transformation. Hortscience: A Publication of the American Society for Hortcultural Science, 2007, 42, 7-10.	1.0	6
51	Mining and characterizing microsatellites from citrus ESTs. Theoretical and Applied Genetics, 2006, 112, 1248-1257.	3.6	216
52	Title is missing!. Plant Cell, Tissue and Organ Culture, 2002, 71, 147-155.	2.3	51
53	Title is missing!. Plant Molecular Biology Reporter, 1999, 17, 231-238.	1.8	4