

# Kunsong Chen

## List of Publications by Year in descending order

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100  
papers

4,923  
citations

87888

38  
h-index

110387

64  
g-index

102  
all docs

102  
docs citations

102  
times ranked

4377  
citing authors

#	ARTICLE	IF	CITATIONS
1	High-resolution spatiotemporal transcriptome mapping of tomato fruit development and ripening. <i>Nature Communications</i> , 2018, 9, 364.	12.8	255
2	Chilling-induced tomato flavor loss is associated with altered volatile synthesis and transient changes in DNA methylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 12580-12585.	7.1	208
3	Global increase in DNA methylation during orange fruit development and ripening. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 1430-1436.	7.1	190
4	Expression of Genes Associated with Aroma Formation Derived from the Fatty Acid Pathway during Peach Fruit Ripening. <i>Journal of Agricultural and Food Chemistry</i> , 2010, 58, 6157-6165.	5.2	184
5	Differential expression within the LOX gene family in ripening kiwifruit. <i>Journal of Experimental Botany</i> , 2006, 57, 3825-3836.	4.8	161
6	Transcription factor CitERF71 activates the terpene synthase gene CitTPS16 involved in the synthesis of E-geraniol in sweet orange fruit. <i>Journal of Experimental Botany</i> , 2017, 68, 4929-4938.	4.8	161
7	Downregulation of RdDM during strawberry fruit ripening. <i>Genome Biology</i> , 2018, 19, 212.	8.8	147
8	Effect of Non-Thermal Plasma-Activated Water on Fruit Decay and Quality in Postharvest Chinese Bayberries. <i>Food and Bioprocess Technology</i> , 2016, 9, 1825-1834.	4.7	142
9	A critical evaluation of the role of ethylene and <i>MADS</i> transcription factors in the network controlling fleshy fruit ripening. <i>New Phytologist</i> , 2019, 221, 1724-1741.	7.3	126
10	Regulatory Mechanisms of Textural Changes in Ripening Fruits. <i>Critical Reviews in Plant Sciences</i> , 2010, 29, 222-243.	5.7	120
11	Low temperature conditioning reduces postharvest chilling injury in loquat fruit. <i>Postharvest Biology and Technology</i> , 2006, 41, 252-259.	6.0	112
12	Transcriptome and metabolome analyses of sugar and organic acid metabolism in Ponkan ( <i>Citrus Tj</i> ) fruit. <i>Journal of Agricultural and Food Chemistry</i> , 2010, 58, 105-115.	2.2	105
13	Bioactive components and antioxidant capacity of Chinese bayberry ( <i>Myrica rubra</i> Sieb. and Zucc.) fruit in relation to fruit maturity and postharvest storage. <i>European Food Research and Technology</i> , 2008, 227, 1091-1097.	3.3	101
14	Integrative analyses of metabolome and genome-wide transcriptome reveal the regulatory network governing flavor formation in kiwifruit ( <i>Actinidia chinensis</i> ). <i>New Phytologist</i> , 2022, 233, 373-389.	7.3	100
15	Three AP2/ERF family members modulate flavonoid synthesis by regulating type IV chalcone isomerase in citrus. <i>Plant Biotechnology Journal</i> , 2021, 19, 671-688.	8.3	99
16	Roles of RIN and ethylene in tomato fruit ripening and ripening-associated traits. <i>New Phytologist</i> , 2020, 226, 460-475.	7.3	98
17	Molecular and Hormonal Mechanisms Regulating Fleshy Fruit Ripening. <i>Cells</i> , 2021, 10, 1136.	4.1	96
18	UV-B irradiation differentially regulates terpene synthases and terpene content of peach. <i>Plant, Cell and Environment</i> , 2017, 40, 2261-2275.	5.7	95

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19	Identification of Proanthocyanidins from Litchi ( <i>Litchi chinensis</i> Sonn.) Pulp by LC-ESI-Q-TOF-MS and Their Antioxidant Activity. <i>PLoS ONE</i> , 2015, 10, e0120480.	2.5	93
20	Phenolic Composition and Antioxidant Properties of Different Peach [ <i>Prunus persica</i> (L.) Batsch] Cultivars in China. <i>International Journal of Molecular Sciences</i> , 2015, 16, 5762-5778.	4.1	85
21	The strawberry transcription factor FaRAV1 positively regulates anthocyanin accumulation by activation of <i>FaMYB10</i> and anthocyanin pathway genes. <i>Plant Biotechnology Journal</i> , 2020, 18, 2267-2279.	8.3	82
22	Genome-Wide Identification, Expression Patterns, and Functional Analysis of UDP Glycosyltransferase Family in Peach ( <i>Prunus persica</i> L. Batsch). <i>Frontiers in Plant Science</i> , 2017, 8, 389.	3.6	76
23	Involvement of multiple phytoene synthase genes in tissue- and cultivar-specific accumulation of carotenoids in loquat. <i>Journal of Experimental Botany</i> , 2013, 65, 4679-4689.	4.8	75
24	Acetylsalicylic acid alleviates chilling injury of postharvest loquat ( <i>Eriobotrya japonica</i> Lindl.) fruit. <i>European Food Research and Technology</i> , 2006, 223, 533-539.	3.3	73
25	Systemic induction of photosynthesis via illumination of the shoot apex is mediated by phytochrome B. <i>Plant Physiology</i> , 2016, 172, pp.01202.2016.	4.8	73
26	An ETHYLENE RESPONSE FACTOR-MYB Transcription Complex Regulates Furaneol Biosynthesis by Activating <i>QUINONE OXIDOREDUCTASE</i> Expression in Strawberry. <i>Plant Physiology</i> , 2018, 178, 189-201.	4.8	70
27	Physicochemical characterisation of four cherry species ( <i>Prunus</i> spp.) grown in China. <i>Food Chemistry</i> , 2015, 173, 855-863.	8.2	66
28	The MrWD40-1 Gene of Chinese Bayberry ( <i>Myrica rubra</i> ) Interacts with MYB and bHLH to Enhance Anthocyanin Accumulation. <i>Plant Molecular Biology Reporter</i> , 2013, 31, 1474-1484.	1.8	65
29	Transcriptional and epigenetic analysis reveals that NAC transcription factors regulate fruit flavor ester biosynthesis. <i>Plant Journal</i> , 2021, 106, 785-800.	5.7	65
30	A NAC transcription factor, EjNAC1, affects lignification of loquat fruit by regulating lignin. <i>Postharvest Biology and Technology</i> , 2015, 102, 25-31.	6.0	64
31	UDP-glucosyltransferase PpUGT85A2 controls volatile glycosylation in peach. <i>Journal of Experimental Botany</i> , 2019, 70, 925-936.	4.8	59
32	CmMYB#7, an R3 MYB transcription factor, acts as a negative regulator of anthocyanin biosynthesis in chrysanthemum. <i>Journal of Experimental Botany</i> , 2019, 70, 3111-3123.	4.8	55
33	ETHYLENE RESPONSE FACTOR39-MYB8 complex regulates low-temperature-induced lignification of loquat fruit. <i>Journal of Experimental Botany</i> , 2020, 71, 3172-3184.	4.8	54
34	Phytochemical Characterization of Chinese Bayberry ( <i>Myrica rubra</i> Sieb. et Zucc.) of 17 Cultivars and Their Antioxidant Properties. <i>International Journal of Molecular Sciences</i> , 2015, 16, 12467-12481.	4.1	52
35	Effect of Ethylene on Cell Wall and Lipid Metabolism during Alleviation of Postharvest Chilling Injury in Peach. <i>Cells</i> , 2019, 8, 1612.	4.1	51
36	Effect of salicylic acid treatment on sensory quality, flavor-related chemicals and gene expression in peach fruit after cold storage. <i>Postharvest Biology and Technology</i> , 2020, 161, 111089.	6.0	48

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37	Phenolic Composition from Different Loquat ( <i>Eriobotrya japonica</i> Lindl.) Cultivars Grown in China and Their Antioxidant Properties. <i>Molecules</i> , 2015, 20, 542-555.	3.8	46
38	Differential Expression of Organic Acid Degradation-Related Genes During Fruit Development of Navel Oranges ( <i>Citrus sinensis</i> ) in Two Habitats. <i>Plant Molecular Biology Reporter</i> , 2013, 31, 1131-1140.	1.8	44
39	Comprehensive structural characterization of phenolics in litchi pulp using tandem mass spectral molecular networking. <i>Food Chemistry</i> , 2019, 282, 9-17.	8.2	41
40	A tomato LATERAL ORGAN BOUNDARIES transcription factor, <i>SlLOB1</i> , predominantly regulates cell wall and softening components of ripening. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	41
41	Genome-Wide Analysis of MYB Gene Family in Chinese Bayberry ( <i>Morella rubra</i> ) and Identification of Members Regulating Flavonoid Biosynthesis. <i>Frontiers in Plant Science</i> , 2021, 12, 691384.	3.6	40
42	Integration of Metabolite Profiling and Transcriptome Analysis Reveals Genes Related to Volatile Terpenoid Metabolism in Finger Citron ( <i>C. medica</i> var. <i>sarcodactylis</i> ). <i>Molecules</i> , 2019, 24, 2564.	3.8	39
43	Effects of flavonoid-rich Chinese bayberry ( <i>Morella rubra</i> Sieb. et Zucc.) fruit extract on regulating glucose and lipid metabolism in diabetic KK-A <sup>y</sup> mice. <i>Food and Function</i> , 2016, 7, 3130-3140.	4.6	38
44	Identification of a lipase gene with a role in tomato fruit short-chain fatty acid-derived flavor volatiles by genome-wide association. <i>Plant Journal</i> , 2020, 104, 631-644.	5.7	37
45	Chitosan/PCL nanofibrous films developed by SBS to encapsulate thymol/HP $\beta$ CD inclusion complexes for fruit packaging. <i>Carbohydrate Polymers</i> , 2022, 286, 119267.	10.2	36
46	Synthesis of flavour-related linalool is regulated by <i>PpHLH1</i> and associated with changes in DNA methylation during peach fruit ripening. <i>Plant Biotechnology Journal</i> , 2021, 19, 2082-2096.	8.3	35
47	E-Nose and GC-MS Reveal a Difference in the Volatile Profiles of White- and Red-Fleshed Peach Fruit. <i>Sensors</i> , 2018, 18, 765.	3.8	34
48	Low Temperature Induced Changes in Citrate Metabolism in Ponkan ( <i>Citrus reticulata</i> Blanco cv.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 3	2.5	33
49	Combination of the biocontrol yeast <i>Cryptococcus laurentii</i> with UV-C treatment for control of postharvest diseases of tomato fruit. <i>BioControl</i> , 2013, 58, 269-281.	2.0	32
50	Peach Carboxylesterase PpCXE1 Is Associated with Catabolism of Volatile Esters. <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 5189-5196.	5.2	32
51	Hydroxylation decoration patterns of flavonoids in horticultural crops: chemistry, bioactivity, and biosynthesis. <i>Horticulture Research</i> , 2022, 9, .	6.3	32
52	Label-free visualization of fruit lignification: Raman molecular imaging of loquat lignified cells. <i>Plant Methods</i> , 2018, 14, 58.	4.3	30
53	Transcriptome analysis provides insights into the regulation of metabolic processes during postharvest cold storage of loquat ( <i>Eriobotrya japonica</i> ) fruit. <i>Horticulture Research</i> , 2019, 6, 49.	6.3	29
54	Effects of cushioning materials and temperature on quality damage of ripe peaches according to the vibration test. <i>Food Packaging and Shelf Life</i> , 2020, 25, 100518.	7.5	28

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55	Genome-Wide Identification and Functional Analysis of Carboxylesterase and Methylsterase Gene Families in Peach ( <i>Prunus persica</i> L. Batsch). <i>Frontiers in Plant Science</i> , 2019, 10, 1511.	3.6	27
56	Involvement of MdUGT75B1 and MdUGT71B1 in flavonol galactoside/glucooside biosynthesis in apple fruit. <i>Food Chemistry</i> , 2020, 312, 126124.	8.2	24
57	Combination Strategy of Reactive and Catalytic Matrices for Qualitative and Quantitative Profiling of N-Glycans in MALDI-MS. <i>Analytical Chemistry</i> , 2019, 91, 9251-9258.	6.5	23
58	Rapid and Non-Destructive Detection of Decay in Peach Fruit at the Cold Environment Using a Self-Developed Handheld Electronic-Nose System. <i>Food Analytical Methods</i> , 2018, 11, 2990-3004.	2.6	22
59	Linalool synthesis related PpTPS1 and PpTPS3 are activated by transcription factor PpERF61 whose expression is associated with DNA methylation during peach fruit ripening. <i>Plant Science</i> , 2022, 317, 111200.	3.6	22
60	Anti-Obesity and Hypoglycemic Effects of <i>Poncirus trifoliata</i> L. Extracts in High-Fat Diet C57BL/6 Mice. <i>Molecules</i> , 2016, 21, 453.	3.8	21
61	EjHAT1 Participates in Heat Alleviation of Loquat Fruit Lignification by Suppressing the Promoter Activity of Key Lignin Monomer Synthesis Gene EjCAD5. <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 5204-5211.	5.2	21
62	Comprehensive Profiling of Phenolic Compounds in White and Red Chinese Bayberries ( <i>Morella</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Networking. <i>Journal of Agricultural and Food Chemistry</i> , 2021, 69, 741-749.	5.2	18
63	Heat shock transcription factors expression during fruit development and under hot air stress in Ponkan ( <i>Citrus reticulata</i> Blanco cv. Ponkan) fruit. <i>Gene</i> , 2015, 559, 129-136.	2.2	17
64	Protective effect of cyanidin-3-O-glucoside on neonatal porcine islets. <i>Journal of Endocrinology</i> , 2017, 235, 237-249.	2.6	17
65	Label-free visualization of lignin deposition in loquats using complementary stimulated and spontaneous Raman microscopy. <i>Horticulture Research</i> , 2019, 6, 72.	6.3	16
66	Transcriptome and DNA methylome analysis reveal new insights into methyl jasmonate-alleviated chilling injury of peach fruit after cold storage. <i>Postharvest Biology and Technology</i> , 2022, 189, 111915.	6.0	16
67	An EjbHLH14-EjHB1-EjPRX12 module is involved in methyl jasmonate alleviation of chilling-induced lignin deposition in loquat fruit. <i>Journal of Experimental Botany</i> , 2022, 73, 1668-1682.	4.8	16
68	Quantitative visualization of pectin distribution maps of peach fruits. <i>Scientific Reports</i> , 2017, 7, 9275.	3.3	15
69	Volatile Profile and Biosynthesis of Post-harvest Apples are Affected by the Mechanical Damage. <i>Journal of Agricultural and Food Chemistry</i> , 2021, 69, 9716-9724.	5.2	15
70	Application of Solution Blow Spinning for Rapid Fabrication of Gelatin/Nylon 66 Nanofibrous Film. <i>Foods</i> , 2021, 10, 2339.	4.3	15
71	Involvement of CitCHX and CitDIC in Developmental-Related and Postharvest-Hot-Air Driven Citrate Degradation in Citrus Fruits. <i>PLoS ONE</i> , 2015, 10, e0119410.	2.5	15
72	Elucidation of myricetin biosynthesis in <i>Morella rubra</i> of the Myricaceae. <i>Plant Journal</i> , 2021, 108, 411-425.	5.7	14

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73	Ternary complex EjbHLH1-EjMYB2-EjAP2-1 retards low temperature-induced flesh lignification in loquat fruit. <i>Plant Physiology and Biochemistry</i> , 2019, 139, 731-737.	5.8	13
74	CircPlant: An Integrated Tool for circRNA Detection and Functional Prediction in Plants. <i>Genomics, Proteomics and Bioinformatics</i> , 2020, 18, 352-358.	6.9	13
75	Unravelling the consecutive glycosylation and methylation of flavonols in peach in response to UV-B irradiation. <i>Plant, Cell and Environment</i> , 2022, 45, 2158-2175.	5.7	13
76	Hybrid Label-Free Molecular Microscopies for Simultaneous Visualization of Changes in Cell Wall Polysaccharides of Peach at Single- and Multiple-Cell Levels during Postharvest Storage. <i>Cells</i> , 2020, 9, 761.	4.1	12
77	Peach fruit PpNAC1 activates <i>PpFAD3-1</i> transcription to provide $\gamma$ -3 fatty acids for the synthesis of short-chain flavor volatiles. <i>Horticulture Research</i> , 2022, 9, .	6.3	12
78	Biosynthetic labeling with 3-O-propargylcaffeyl alcohol reveals in vivo cell-specific patterned lignification in loquat fruits during development and postharvest storage. <i>Horticulture Research</i> , 2021, 8, 61.	6.3	11
79	Feasibility Study on Quantitative Pixel-Level Visualization of Internal Quality at Different Cross Sections Inside Postharvest Loquat Fruit. <i>Food Analytical Methods</i> , 2017, 10, 287-297.	2.6	10
80	Packaging Design to Protect Hongmeiren Orange Fruit from Mechanical Damage during Simulated and Road Transportation. <i>Horticulturae</i> , 2022, 8, 258.	2.8	10
81	Ougan ( <i>Citrus reticulata</i> cv. <i>Suavissima</i> ) flavedo extract suppresses cancer motility by interfering with epithelial-to-mesenchymal transition in SKOV3 cells. <i>Chinese Medicine</i> , 2015, 10, 14.	4.0	9
82	Grafting Hollow Covalent Organic Framework Nanoparticles with Thermal-Responsive Polymers for the Controlled Release of Preservatives. <i>ACS Applied Materials &amp; Interfaces</i> , 2022, 14, 22982-22988.	8.0	9
83	Transcription Factor CitERF16 Is Involved in Citrus Fruit Sucrose Accumulation by Activating CitSWEET11d. <i>Frontiers in Plant Science</i> , 2021, 12, 809619.	3.6	9
84	Identification and Characterization of Transcripts Differentially Expressed in Peel and Juice Vesicles of Immature and Ripe Orange ( <i>Citrus sinensis</i> ) Fruit. <i>Plant Molecular Biology Reporter</i> , 2008, 26, 121-132.	1.8	8
85	Effects of Combined Heat and Preservative Treatment on Storability of Ponkan Fruit ( <i>Citrus</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 2.6 8	2.6	8
86	Glycosylamines-based reactive matrix designed for imaging acidity in Ponkan fruit using matrix assisted laser desorption/ionization mass spectrometry imaging. <i>Analytica Chimica Acta</i> , 2018, 1041, 78-86.	5.4	8
87	Application of electronic nose and GC-MS for detection of strawberries with vibrational damage. <i>Food Quality and Safety</i> , 2020, 4, 181-192.	1.8	8
88	The Interaction Between CitMYB52 and CitbHLH2 Negatively Regulates Citrate Accumulation by Activating CitALMT in Citrus Fruit. <i>Frontiers in Plant Science</i> , 2022, 13, 848869.	3.6	8
89	The Isolation and Identification of Anthocyanin-Related GSTs in <i>Chrysanthemum</i> . <i>Horticulturae</i> , 2021, 7, 231.	2.8	7
90	Role of the tomato fruit ripening regulator MADS-RIN in resistance to <i>Botrytis cinerea</i> infection. <i>Food Quality and Safety</i> , 2021, 5, .	1.8	7

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91	Cyanidin-3-O-Glucoside improves the viability of human islet cells treated with amylin or A $\beta$ 1-42 in vitro. PLoS ONE, 2021, 16, e0258208.	2.5	7
92	Non-Destructive Detection of Damaged Strawberries after Impact Based on Analyzing Volatile Organic Compounds. Sensors, 2022, 22, 427.	3.8	7
93	Two Myricetin-Derived Flavonols from Morella rubra Leaves as Potent $\beta$ -Glucosidase Inhibitors and Structure-Activity Relationship Study by Computational Chemistry. Oxidative Medicine and Cellular Longevity, 2022, 2022, 1-16.	4.0	7
94	Cultivation Conditions Change Aroma Volatiles of Strawberry Fruit. Horticulturae, 2021, 7, 81.	2.8	5
95	Mechanical damages and packaging methods along the fresh fruit supply chain: A review. Critical Reviews in Food Science and Nutrition, 2023, 63, 10283-10302.	10.3	5
96	Comparative Transcriptome Analysis Revealed Two Alternative Splicing bHLHs Account for Flower Color Alteration in Chrysanthemum. International Journal of Molecular Sciences, 2021, 22, 12769.	4.1	4
97	EjMYB4 is a transcriptional activator of 4-Coumarate:coenzyme A ligase involved in lignin biosynthesis in loquat (Eriobotrya japonica). Plant Growth Regulation, 2018, 86, 413-421.	3.4	3
98	Off-flavor and Loss of Aroma in Young Coconut Fruit During Cold Storage are Associated with the Expression of Genes Derived from the LOX Pathway and <i>Badh2</i> . Horticulture Journal, 2022, 91, 209-220.	0.8	3
99	Determination of 9(10H)-Acridone by HPLC with Fluorescence Detection. Journal of Liquid Chromatography and Related Technologies, 2007, 30, 245-254.	1.0	1
100	C-CorA: A Cluster-Based Method for Correlation Analysis of RNA-Seq Data. Horticulturae, 2022, 8, 124.	2.8	1