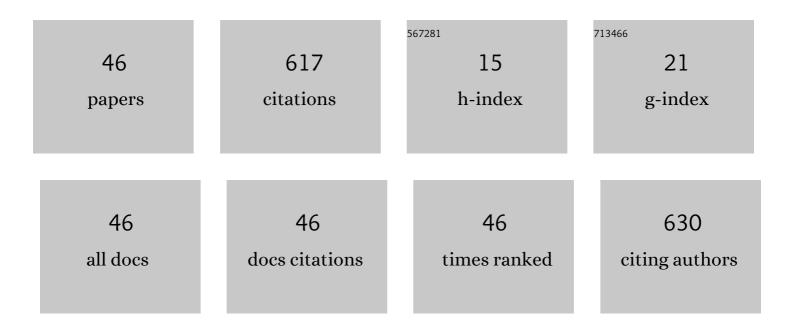
Bai-Hong Chen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5728149/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Anthocyanin accumulation correlates with hormones in the fruit skin of â€~Red Delicious' and its four generation bud sport mutants. BMC Plant Biology, 2018, 18, 363.	3.6	55
2	Whole-genome DNA methylation patterns and complex associations with gene expression associated with anthocyanin biosynthesis in apple fruit skin. Planta, 2019, 250, 1833-1847.	3.2	53
3	Genome-Wide Identification and Expression Analysis of GA2ox, GA3ox, and GA20ox Are Related to Gibberellin Oxidase Genes in Grape (Vitis Vinifera L.). Genes, 2019, 10, 680.	2.4	44
4	Synthesis of light-inducible and light-independent anthocyanins regulated by specific genes in grape â€~Marselan' (<i>V. vinifera</i> L.). PeerJ, 2019, 7, e6521.	2.0	31
5	Transcriptome and Metabolite Conjoint Analysis Reveals that Exogenous Methyl Jasmonate Regulates Monoterpene Synthesis in Grape Berry Skin. Journal of Agricultural and Food Chemistry, 2020, 68, 5270-5281.	5.2	29
6	Elevated CO2 concentration promotes photosynthesis of grape (Vitis vinifera L. cv. â€~Pinot noir') plantlet in vitro by regulating RbcS and Rca revealed by proteomic and transcriptomic profiles. BMC Plant Biology, 2019, 19, 42.	3.6	28
7	RNA Sequencing Reveals that Endoplasmic Reticulum Stress and Disruption of Membrane Integrity Underlie Dimethyl Trisulfide Toxicity against Fusarium oxysporum f. sp. cubense Tropical Race 4. Frontiers in Microbiology, 2017, 8, 1365.	3.5	25
8	Genomeâ€wide identification and characterization of the <i>BES1</i> gene family in apple (<i>Malus) Tj ETQq0 (</i>) 0.rgBT /C	verlock 10
9	Genome-wide annotation and expression responses to biotic stresses of the WALL-ASSOCIATED KINASE - RECEPTOR-LIKE KINASE (WAK-RLK) gene family in Apple (Malus domestica). European Journal of Plant Pathology, 2019, 153, 771-785.	1.7	20
10	Genome-Wide Analysis of the Apple (Malus domestica) Cysteine-Rich Receptor-Like Kinase (CRK) Family: Annotation, Genomic Organization, and Expression Profiles in Response to Fungal Infection. Plant Molecular Biology Reporter, 2020, 38, 14-24.	1.8	20
11	Identification and expression analysis of the small auxin-up RNA (SAUR) gene family in apple by inducing of auxin. Gene, 2020, 750, 144725.	2.2	20
12	The Changes in Color, Soluble Sugars, Organic Acids, Anthocyanins and Aroma Components in "Starkrimson―during the Ripening Period in China. Molecules, 2016, 21, 812.	3.8	18

13 Transcriptome analysis revealed glucose application affects plant hormone signal transduction pathway in "Red Globe―grape plantlets. Plant Growth Regulation, 2018, 84, 45-56.	3.4	18
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Genome-Wide Identification and Expression Analysis of the CrRLK1L Gene Family in Apple (Malus) Tj ETQq0 0 0 rgBT $_{1.8}^{10}$ Overlock 10 Tf 50

15	Recent trends and comprehensive appraisal for the biotechnological production of trans-resveratrol and its derivatives. Phytochemistry Reviews, 2018, 17, 491-508.	6.5	17
16	Different exogenous sugars affect the hormone signal pathway and sugar metabolism in "Red Globe― (Vitis vinifera L.) plantlets grown in vitro as shown by transcriptomic analysis. Planta, 2017, 246, 537-552.	3.2	15
17	Significant and unique changes in phosphorylation levels of four phosphoproteins in two apple rootstock genotypes under drought stress. Molecular Genetics and Genomics, 2017, 292, 1307-1322.	2.1	13
18	Effects of short-term heat stress on PSII and subsequent recovery for senescent leaves of Vitis vinifera L. cv. Red Globe. Journal of Integrative Agriculture, 2018, 17, 2683-2693.	3.5	13

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19	Genome-wide identification of BAM genes in grapevine (Vitis vinifera L.) and ectopic expression of VvBAM1 modulating soluble sugar levels to improve low-temperature tolerance in tomato. BMC Plant Biology, 2021, 21, 156.	3.6	13
20	Effects of paclobutrazol on the physiological characteristics of <i>Malus halliana</i> Koehne Seedlings under drought stress via principal component analysis and membership function analysis. Arid Land Research and Management, 2019, 33, 97-113.	1.6	11
21	Genome-wide characterization and expression analyses of the auxin/indole-3-acetic acid (Aux/IAA) gene family in apple (Malus domestica). Gene, 2021, 768, 145302.	2.2	11
22	Cyclic nucleotide gated channel genes (CNGCs) in Rosaceae: genome-wide annotation, evolution and the roles on Valsa canker resistance. Plant Cell Reports, 2021, 40, 2369-2382.	5.6	10
23	Temperature-phase transcriptomics reveals that hormones and sugars in the phloem of grape participate in tolerance during cold acclimation. Plant Cell Reports, 2022, 41, 1357-1373.	5.6	10
24	Effects of exogenous growth regulators and bud picking on grafting of grapevine hard branches. Scientia Horticulturae, 2020, 264, 109186.	3.6	9
25	Effects of Shading on the Synthesis of Volatile Organic Compounds in â€~Marselan' Grape Berries (Vitis) T	j ETQg1 1 0. 5.1	784314 rg81
26	A Novel Identification Method for Apple (Malus domestica Borkh.) Cultivars Based on a Deep Convolutional Neural Network with Leaf Image Input. Symmetry, 2020, 12, 217.	2.2	9
27	VaAPL1 Promotes Starch Synthesis to Constantly Contribute to Soluble Sugar Accumulation, Improving Low Temperature Tolerance in Arabidopsis and Tomato. Frontiers in Plant Science, 0, 13, .	3.6	8
28	Physical–Chemical Composition and Quality Related Changes in "Ruaner―Pear (Pyrus ussuriensis) During Freezing–Thawing Period. Molecules, 2019, 24, 2611.	3.8	7
29	The mechanism of color fading in sunburned apple peel. Acta Physiologiae Plantarum, 2019, 41, 1.	2.1	7
30	Branch age and angle as crucial drivers of leaf photosynthetic performance and fruiting in high-density planting: A study case in spur-type apple "Vallee Spur―(Malus domestica). Scientia Horticulturae, 2019, 246, 898-906.	3.6	7
31	MYB_SH[AL]QKY[RF] transcription factors <i>MdLUX</i> and <i>MdPCL-like</i> promote anthocyanin accumulation through DNA hypomethylation and <i>MdF3H</i> activation in apple. Tree Physiology, 2021, 41, 836-848.	3.1	7
32	RNA sequencing analysis provides new insights into dynamic molecular responses to Valsa mali pathogenicity in apple †Changfu No. 2'. Tree Genetics and Genomes, 2018, 14, 1.	1.6	6
33	Genome-wide identification and expression analysis of the EXO70 gene family in grape (<i>Vitis) Tj ETQq1 1 0</i>	.784314 rgB 2.0	T /Qverlock
34	Effects of CEPA and 1-MCP on Flower Bud Differentiation of Apple cv. â€~Nagafu No.2' Grafted on Different Rootstocks. Journal of Plant Growth Regulation, 2019, 38, 842-854.	5.1	5
35	Insight into VvGH3 genes evolutional relationship from monocotyledons and dicotyledons reveals that VvGH3-9 negatively regulates the drought tolerance in transgenic Arabidopsis. Plant Physiology and Biochemistry, 2022, 172, 70-86.	5.8	4
36	Exogenous ABA and its inhibitor regulate flower bud induction of apple cv. â€~Nagafu No. 2′ grafted on different rootstocks. Trees - Structure and Function, 2021, 35, 609-620.	1.9	3

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#	Article	IF	CITATIONS
37	Thin layer drying kinetics and quality dynamics of persimmon (Diospyros kaki) treated with preservatives and solar dried under different temperatures. PLoS ONE, 2022, 17, e0265111.	2.5	3
38	Genome-wide Identification and Characterization of the Strawberry (Fragaria Vesca) FvAP2/ERF Gene Family in Abiotic Stress. Plant Molecular Biology Reporter, 2022, 40, 646-660.	1.8	3
39	Petiole hormones act as regulators in the early phototropic leaf movements of grape (Vitis vinifera L.) revealed by comparative transcriptome profiling. Scientia Horticulturae, 2021, 283, 110049.	3.6	2
40	Effects of CO2 on transplantation of grape plantlets cultured in vitro by promoting photosynthesis. Scientia Horticulturae, 2021, 287, 110286.	3.6	2
41	Transcriptomic Analysis Revealed Hormone-Related and Receptor-Like Kinase Genes Involved in Wound Healing of â€~Duli' and its Resistance to Valsa Pyri. Plant Molecular Biology Reporter, 0, , 1.	1.8	1
42	Genome-wide annotation and expression responses to biotic stresses of the WALL-ASSOCIATED KINASE - RECEPTOR-LIKE KINASE (WAK-RLK) gene family in Apple (Malus domestica). , 2019, 153, 771.		1
43	Exogenous carbon promotes plantlet growth by inducing ethylene signaling in grapevine. Scientia Horticulturae, 2022, 293, 110659.	3.6	1

44 Molecular cloning, bioinformatics analysis, and transient expression of MdAux/IAA28 in apple (Malus) Tj ETQq0 0 0 ggBT /Overlock 10 Tf

45	Multivariate Statistical Analyses and Predictive Model of Cold Resistance Associated with Eleven Crabapples and Fuji Apple. Cryo-Letters, 2018, 39, 235-244.	0.3	Ο
46	Transcriptome Profile in a Susceptible Pear â€~Zaosu' (Pyrus bretschneideri Rehd.)–Valsa pyri Interaction. Journal of Plant Growth Regulation, 0, , 1.	5.1	0