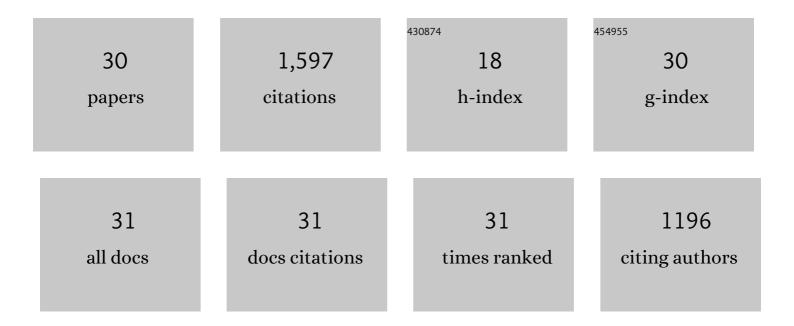
Fangyuan Zhang

List of Publications by Year in descending order

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ΕλΝΟΥΠΑΝ ΖΗΛΝΟ

#	Article	IF	CITATIONS
1	Molecular insights into AabZIP1-mediated regulation on artemisinin biosynthesis and drought tolerance in Artemisia annua. Acta Pharmaceutica Sinica B, 2022, 12, 1500-1513.	12.0	17
2	Engineering Nootkatone Biosynthesis in <i>Artemisia annua</i> . ACS Synthetic Biology, 2021, 10, 957-963.	3.8	9
3	High-Level Patchoulol Biosynthesis in Artemisia annua L Frontiers in Bioengineering and Biotechnology, 2020, 8, 621127.	4.1	3
4	Effect of Thermal Processing on Carotenoids and Folate Changes in Six Varieties of Sweet Potato (Ipomoes batata L.). Foods, 2019, 8, 215.	4.3	18
5	The cold-induced transcription factor bHLH112 promotes artemisinin biosynthesis indirectly via ERF1 in Artemisia annua. Journal of Experimental Botany, 2019, 70, 4835-4848.	4.8	47
6	Metabolic Characterization of Hyoscyamus niger Ornithine Decarboxylase. Frontiers in Plant Science, 2019, 10, 229.	3.6	10
7	Interaction of bZIP transcription factor TGA6 with salicylic acid signaling modulates artemisinin biosynthesis in Artemisia annua. Journal of Experimental Botany, 2019, 70, 3969-3979.	4.8	46
8	AaPP2C1 negatively regulates the expression of genes involved in artemisinin biosynthesis through dephosphorylating AaAPK1. FEBS Letters, 2019, 593, 743-750.	2.8	5
9	Effect of Steaming Processing on Phenolic Profiles and Cellular Antioxidant Activities of Castanea mollissima. Molecules, 2019, 24, 703.	3.8	16
10	The Genome of Artemisia annua Provides Insight into the Evolution of Asteraceae Family and Artemisinin Biosynthesis. Molecular Plant, 2018, 11, 776-788.	8.3	205
11	Comparison of two hyoscyamine 6β-hydroxylases in engineering scopolamine biosynthesis in root cultures of Scopolia lurida. Biochemical and Biophysical Research Communications, 2018, 497, 25-31.	2.1	27
12	ARTEMISININ BIOSYNTHESIS PROMOTING KINASE 1 positively regulates artemisinin biosynthesis through phosphorylating AabZIP1. Journal of Experimental Botany, 2018, 69, 1109-1123.	4.8	40
13	The Artemisia annua FLOWERING LOCUS T Homolog 2, AaFT2, is a key regulator of flowering time. Plant Physiology and Biochemistry, 2018, 126, 197-205.	5.8	5
14	Metabolic characterization of Hyoscyamus niger root-specific putrescine N-methyltransferase. Plant Physiology and Biochemistry, 2018, 127, 47-54.	5.8	17
15	Molecular Characterization of the 1-Deoxy-D-Xylulose 5-Phosphate Synthase Gene Family in Artemisia annua. Frontiers in Plant Science, 2018, 9, 952.	3.6	27
16	Molecular cloning and characterization of the promoter of aldehyde dehydrogenase gene from <i>Artemisia annua</i> . Biotechnology and Applied Biochemistry, 2017, 64, 902-910.	3.1	5
17	Overexpression of a Novel NAC Domain-Containing Transcription Factor Gene (<i>AaNAC1</i>) Enhances the Content of Artemisinin and Increases Tolerance to Drought and <i>Botrytis cinerea</i> in <i>Artemisia annua</i> . Plant and Cell Physiology, 2016, 57, 1961-1971.	3.1	95
18	Genome-wide inference of protein interaction network and its application to the study of crosstalk in Arabidopsis abscisic acid signaling. Plant Physiology, 2016, 171, pp.00057.2016.	4.8	27

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#	ARTICLE	IF	CITATIONS
19	The jasmonateâ€responsive Aa <scp>MYC</scp> 2 transcription factor positively regulates artemisinin biosynthesis in <i>Artemisia annua</i> . New Phytologist, 2016, 210, 1269-1281.	7.3	230
20	Branch Pathway Blocking in <i>Artemisia annua</i> is a Useful Method for Obtaining High Yield Artemisinin. Plant and Cell Physiology, 2016, 57, 588-602.	3.1	70
21	A Basic Leucine Zipper Transcription Factor, AabZIP1, Connects Abscisic Acid Signaling with Artemisinin Biosynthesis in Artemisia annua. Molecular Plant, 2015, 8, 163-175.	8.3	198
22	Overexpression of Allene Oxide Cyclase Improves the Biosynthesis of Artemisinin in Artemisia annua L PLoS ONE, 2014, 9, e91741.	2.5	27
23	Type 2C Phosphatase 1 of <i>Artemisia annua</i> L. Is a Negative Regulator of ABA Signaling. BioMed Research International, 2014, 2014, 1-9.	1.9	14
24	Molecular Cloning and Characterization of a Trichome-Specific Promoter of Artemisinic Aldehyde Δ11(13) Reductase (DBR2) in Artemisia annua. Plant Molecular Biology Reporter, 2014, 32, 82-91.	1.8	35
25	Characterization of the Promoter of Artemisia annua Amorpha-4,11-diene Synthase (ADS) Gene Using Homologous and Heterologous Expression as well as Deletion Analysis. Plant Molecular Biology Reporter, 2014, 32, 406-418.	1.8	20
26	The stacked over-expression of FPS, CYP71AV1 and CPR genes leads to the increase of artemisinin level in Artemisia annua L Plant Biotechnology Reports, 2013, 7, 287-295.	1.5	34
27	<i><scp>A</scp>a<scp>ORA</scp></i> , a trichomeâ€specific <scp>AP</scp> 2/ <scp>ERF</scp> transcription factor of <i><scp>A</scp>rtemisia annua</i> , is a positive regulator in the artemisinin biosynthetic pathway and in disease resistance to <i><scp>B</scp>otrytis cinerea</i> . New Phytologist, 2013, 198, 1191-1202.	7.3	255
28	Overexpression of the Artemisia Orthologue of ABA Receptor, AaPYL9, Enhances ABA Sensitivity and Improves Artemisinin Content in Artemisia annua L. PLoS ONE, 2013, 8, e56697.	2.5	61
29	Identification of Putative Artemisia annua ABCG Transporter Unigenes Related to Artemisinin Yield Following Expression Analysis in Different Plant Tissues and in Response to Methyl Jasmonate and Abscisic Acid Treatments. Plant Molecular Biology Reporter, 2012, 30, 838-847.	1.8	20
30	Characterization of the Jasmonate Biosynthetic Gene Allene Oxide Cyclase in Artemisia annua L., Source of the Antimalarial Drug Artemisinin. Plant Molecular Biology Reporter, 2011, 29, 489-497.	1.8	14