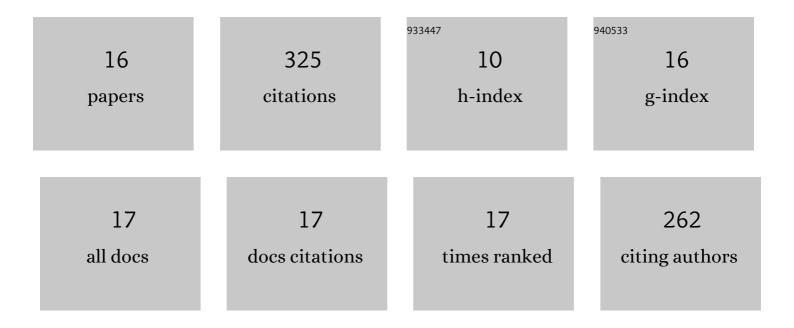
## Anzhi Wei

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3971888/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Extensive Sampling Provides New Insights into Phylogenetic Relationships between Wild and Domesticated Zanthoxylum Species in China. Horticulturae, 2022, 8, 440.	2.8	3
2	ZbAGL11, a class D MADS-box transcription factor of Zanthoxylum bungeanum, is involved in sporophytic apomixis. Horticulture Research, 2021, 8, 23.	6.3	14
3	Transcriptome and metabolite analysis reveals key genes for melanin synthesis during the development of Zanthoxylum bungeanum seeds. Industrial Crops and Products, 2021, 165, 113419.	5.2	8
4	<i>Zanthoxylum-</i> specific whole genome duplication and recent activity of transposable elements in the highly repetitive paleotetraploid <i>Z. bungeanum</i> genome. Horticulture Research, 2021, 8, 205.	6.3	19
5	Genomic analysis reveals the genetic diversity, population structure, evolutionary history and relationships of Chinese pepper. Horticulture Research, 2020, 7, 158.	6.3	25
6	Genetic Diversity and Evolutionary Relationships of Chinese Pepper Based on nrDNA Markers. Forests, 2020, 11, 543.	2.1	5
7	miRNAs and their target genes regulate the antioxidant system of Zanthoxylum bungeanum under drought stress. Plant Physiology and Biochemistry, 2020, 150, 196-203.	5.8	23
8	Geographical variations in the fatty acids of Zanthoxylum seed oils: A chemometric classification based on the random forest algorithm. Industrial Crops and Products, 2019, 134, 146-153.	5.2	33
9	The steps from sexual reproduction to apomixis. Planta, 2019, 249, 1715-1730.	3.2	22
10	Single-Molecule Long-Read Sequencing of Zanthoxylum bungeanum Maxim. Transcriptome: Identification of Aroma-Related Genes. Forests, 2018, 9, 765.	2.1	14
11	Genetic structure of cultivated Zanthoxylum species investigated with SSR markers. Tree Genetics and Genomes, 2018, 14, 1.	1.6	8
12	Expression Stabilities of Ten Candidate Reference Genes for RT-qPCR in Zanthoxylum bungeanum Maxim. Molecules, 2018, 23, 802.	3.8	46
13	De novo transcriptome assembly of Zanthoxylum bungeanum using Illumina sequencing for evolutionary analysis and simple sequence repeat marker development. Scientific Reports, 2017, 7, 16754.	3.3	38
14	Phylogenetic relationships among cultivated Zanthoxylum species in China based on cpDNA markers. Tree Genetics and Genomes, 2016, 12, 1.	1.6	30
15	Genetic relationships of Chinese prickly ash as revealed by ISSR markers. Biologia (Poland), 2015, 70, 45-51.	1.5	11
16	Genetic diversity and relationships of wild and cultivated Zanthoxylum germplasms based on sequence-related amplified polymorphism (SRAP) markers. Genetic Resources and Crop Evolution, 2015, 62, 1193-1204.	1.6	26