

# Xiaoming Yang

## List of Publications by Year in descending order

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13  
papers

615  
citations

1040056

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1125743

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docs citations

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times ranked

729  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic analyses of primitive, wild and cultivated citrus provide insights into asexual reproduction. <i>Nature Genetics</i> , 2017, 49, 765-772.	21.4	316
2	Genome of Wild Mandarin and Domestication History of Mandarin. <i>Molecular Plant</i> , 2018, 11, 1024-1037.	8.3	130
3	Tung Tree ( <i>Vernicia fordii</i> ) Genome Provides A Resource for Understanding Genome Evolution and Improved Oil Production. <i>Genomics, Proteomics and Bioinformatics</i> , 2019, 17, 558-575.	6.9	43
4	Systematic investigation and expression profiles of the GbR2R3-MYB transcription factor family in ginkgo ( <i>Ginkgo biloba</i> L.). <i>International Journal of Biological Macromolecules</i> , 2021, 172, 250-262.	7.5	23
5	Structural characterization and comparative analysis of the chloroplast genome of <i>Ginkgo biloba</i> and other gymnosperms. <i>Journal of Forestry Research</i> , 2021, 32, 765-778.	3.6	19
6	Genetic diversity and population structure of pummelo ( <i>Citrus maxima</i> ) germplasm in China. <i>Tree Genetics and Genomes</i> , 2017, 13, 1.	1.6	16
7	Molecular phylogeography and population evolution analysis of <i>Citrus ichangensis</i> (Rutaceae). <i>Tree Genetics and Genomes</i> , 2017, 13, 1.	1.6	15
8	<i>Taxus yunnanensis</i> genome offers insights into gymnosperm phylogeny and taxol production. <i>Communications Biology</i> , 2021, 4, 1203.	4.4	15
9	The genetic diversity and population structure of <i>Sophora alopecuroides</i> (Faboideae) as determined by microsatellite markers developed from transcriptome. <i>PLoS ONE</i> , 2019, 14, e0226100.	2.5	12
10	Selection of Suitable Reference Genes Based on Transcriptomic Data in <i>Ginkgo biloba</i> under Different Experimental Conditions. <i>Forests</i> , 2020, 11, 1217.	2.1	11
11	Genome-Wide Identification and Coexpression Network Analysis of DNA Methylation Pathway Genes and Their Differentiated Functions in <i>Ginkgo biloba</i> L.. <i>Forests</i> , 2020, 11, 1076.	2.1	6
12	The Diversity of <i>Melia azedarach</i> L. from China Based on Transcriptome-Developed SSR Marker. <i>Forests</i> , 2022, 13, 1011.	2.1	5
13	Molecular cloning and expression analysis of a WRKY transcription factor gene, GbWRKY20, from <i>Ginkgo biloba</i> . <i>Plant Signaling and Behavior</i> , 2021, 16, 1930442.	2.4	4