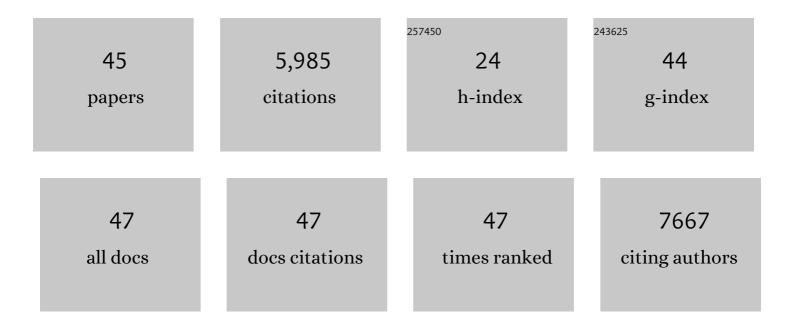
## Yuan-Nian Jiao

List of Publications by Year in descending order

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YUAN-NIAN ILAO

#	Article	lF	CITATIONS
1	WheatCENet: A Database for Comparative Co-Expression Networks Analysis of Allohexaploid Wheat and Its Progenitors. Genomics, Proteomics and Bioinformatics, 2023, 21, 324-336.	6.9	6
2	The <i>Chrysanthemum lavandulifolium</i> genome and the molecular mechanism underlying diverse capitulum types. Horticulture Research, 2022, 9, .	6.3	24
3	Genomes shed light on the evolution of <i>Begonia</i> , a megaâ€diverse genus. New Phytologist, 2022, 234, 295-310.	7.3	18
4	Diversity, phylogeny, and adaptation of bryophytes: insights from genomic and transcriptomic data. Journal of Experimental Botany, 2022, 73, 4306-4322.	4.8	16
5	A recent burst of gene duplications in Triticeae. Plant Communications, 2022, 3, 100268.	7.7	18
6	The Gastrodia menghaiensis (Orchidaceae) genome provides new insights of orchid mycorrhizal interactions. BMC Plant Biology, 2022, 22, 179.	3.6	13
7	The Cycas genome and the early evolution of seed plants. Nature Plants, 2022, 8, 389-401.	9.3	80
8	Chromosome-level and haplotype-resolved genome provides insight into the tetraploid hybrid origin of patchouli. Nature Communications, 2022, 13, .	12.8	20
9	Phylogenomic Analysis Reconstructed the Order Matoniales from Paleopolyploidy Veil. Plants, 2022, 11, 1529.	3.5	3
10	Insights into angiosperm evolution, floral development and chemical biosynthesis from the Aristolochia fimbriata genome. Nature Plants, 2021, 7, 1239-1253.	9.3	51
11	The Physalis floridana genome provides insights into the biochemical and morphological evolution of Physalis fruits. Horticulture Research, 2021, 8, 244.	6.3	15
12	New insights into the dispersion history and adaptive evolution of taxon Aegilops tauschii in China. Journal of Genetics and Genomics, 2021, , .	3.9	3
13	Whole-Genome Sequencing of <i>Acer catalpifolium</i> Reveals Evolutionary History of Endangered Species. Genome Biology and Evolution, 2021, 13, .	2.5	7
14	Genetic Contribution of Paleopolyploidy to Adaptive Evolution in Angiosperms. Molecular Plant, 2020, 13, 59-71.	8.3	178
15	The water lily genome and the early evolution of flowering plants. Nature, 2020, 577, 79-84.	27.8	238
16	The Chromosome-Based Rubber Tree Genome Provides New Insights into Spurge Genome Evolution and Rubber Biosynthesis. Molecular Plant, 2020, 13, 336-350.	8.3	73
17	The ancient wave of polyploidization events in flowering plants and their facilitated adaptation to environmental stress. Plant, Cell and Environment, 2020, 43, 2847-2856.	5.7	71
18	A Collinearity-Incorporating Homology Inference Strategy for Connecting Emerging Assemblies in the Triticeae Tribe as a Pilot Practice in the Plant Pangenomic Era. Molecular Plant, 2020, 13, 1694-1708.	8.3	126

Yuan-Nian Jiao

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19	The Tetracentron genome provides insight into the early evolution of eudicots and the formation of vessel elements. Genome Biology, 2020, 21, 291.	8.8	23
20	A chromosome-scale reference genome of Aquilegia oxysepala var. kansuensis. Horticulture Research, 2020, 7, 113.	6.3	20
21	The hornwort genome and early land plant evolution. Nature Plants, 2020, 6, 107-118.	9.3	203
22	In silico identification and structure function analysis of a putative coclaurine N-methyltransferase from Aristolochia fimbriata. Computational Biology and Chemistry, 2020, 85, 107201.	2.3	1
23	Evolutionary strategies drive a balance of the interacting gene products for the <i>CBL</i> and <i>CIPK</i> gene families. New Phytologist, 2020, 226, 1506-1516.	7.3	52
24	Genome Sequence of Striga asiatica Provides Insight into the Evolution of Plant Parasitism. Current Biology, 2019, 29, 3041-3052.e4.	3.9	109
25	Gene duplication and genetic innovation in cereal genomes. Genome Research, 2019, 29, 261-269.	5.5	29
26	Double the Genome, Double the Fun: Genome Duplications in Angiosperms. Molecular Plant, 2018, 11, 357-358.	8.3	33
27	Clinical cancer genomic profiling by three-platform sequencing of whole genome, whole exome and transcriptome. Nature Communications, 2018, 9, 3962.	12.8	142
28	The Genome of Opium Poppy Reveals Evolutionary History of Morphinan Pathway. Genomics, Proteomics and Bioinformatics, 2018, 16, 460-462.	6.9	5
29	A TRIM insertion in the promoter of Ms2 causes male sterility in wheat. Nature Communications, 2017, 8, 15407.	12.8	79
30	Genome-wide mapping of DNase I hypersensitive sites reveals chromatin accessibility changes in Arabidopsis euchromatin and heterochromatin regions under extended darkness. Scientific Reports, 2017, 7, 4093.	3.3	19
31	The Aegilops tauschii genome reveals multiple impacts of transposons. Nature Plants, 2017, 3, 946-955.	9.3	164
32	The Genome of a Southern Hemisphere Seagrass Species ( <i>Zostera muelleri</i> ). Plant Physiology, 2016, 172, 272-283.	4.8	88
33	Multi-Phase US Spread and Habitat Switching of a Post-Columbian Invasive, Sorghum halepense. PLoS ONE, 2016, 11, e0164584.	2.5	28
34	Evolution of the H9N2 influenza genotype that facilitated the genesis of the novel H7N9 virus. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 548-553.	7.1	287
35	Prehistory of the Angiosperms. Advances in Botanical Research, 2014, , 223-245.	1.1	7
36	Integrated Syntenic and Phylogenomic Analyses Reveal an Ancient Genome Duplication in Monocots. Plant Cell, 2014, 26, 2792-2802.	6.6	220

Yuan-Nian Jiao

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37	Polyploidy-associated genome modifications during land plant evolution. Philosophical Transactions of the Royal Society B: Biological Sciences, 2014, 369, 20130355.	4.0	90
38	Evolution of a horizontally acquired legume gene, albumin 1, in the parasitic plant Phelipanche aegyptiaca and related species. BMC Evolutionary Biology, 2013, 13, 48.	3.2	39
39	Characterization of the basal angiosperm Aristolochia fimbriata: a potential experimental system for genetic studies. BMC Plant Biology, 2013, 13, 13.	3.6	22
40	The <i>Amborella</i> Genome and the Evolution of Flowering Plants. Science, 2013, 342, 1241089.	12.6	743
41	Genome of the long-living sacred lotus (Nelumbo nucifera Gaertn.). Genome Biology, 2013, 14, R41.	9.6	329
42	A genome triplication associated with early diversification of the core eudicots. Genome Biology, 2012, 13, R3.	9.6	389
43	A physical map for the Amborella trichopoda genome sheds light on the evolution of angiosperm genome structure. Genome Biology, 2011, 12, R48.	9.6	28
44	Ancestral polyploidy in seed plants and angiosperms. Nature, 2011, 473, 97-100.	27.8	1,862
45	Generation of a large-scale genomic resource for functional and comparative genomics in Liriodendron tulipifera L Tree Genetics and Genomes, 2011, 7, 941-954.	1.6	13