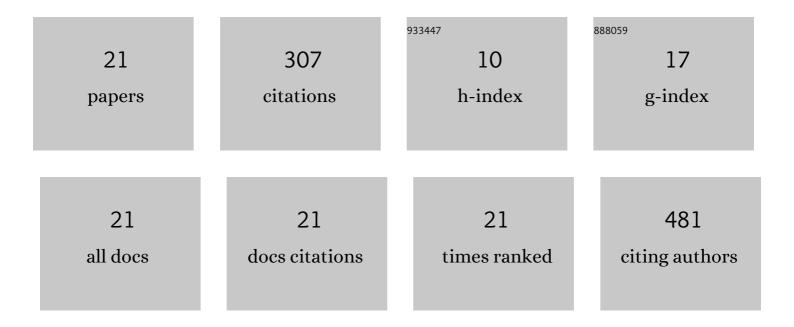
Ying Bao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6659805/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Possible role of the CYC2c gene in the cornflower-like ray floret phenotype of Gaillardia cultivars. Journal of Plant Research, 2022, 135, 465-472.	2.4	5
2	Origins of weedy rice revealed by polymorphisms of chloroplast DNA sequences and nuclear microsatellites. Journal of Systematics and Evolution, 2021, 59, 316-325.	3.1	4
3	Genome-Wide Mining of MYB Transcription Factors in the Anthocyanin Biosynthesis Pathway of Gossypium Hirsutum. Biochemical Genetics, 2021, 59, 678-696.	1.7	12
4	PIF4: Integrator of light and temperature cues in plant growth. Plant Science, 2021, 313, 111086.	3.6	16
5	Changes in the promoter of a defender against apoptotic cell death gene affect its expression in upland cotton. Journal of Systematics and Evolution, 2020, 59, 1018.	3.1	1
6	Unraveling cis and trans regulatory evolution during cotton domestication. Nature Communications, 2019, 10, 5399.	12.8	58
7	Abundant Small Genetic Alterations after Upland Cotton Domestication. BioMed Research International, 2018, 2018, 1-7.	1.9	2
8	PIDA/I ₂ -Mediated α- and β-C(sp ³)–H Bond Dual Functionalization of Tertiary Amines. Journal of Organic Chemistry, 2018, 83, 10166-10174.	3.2	22
9	Synthesis of selective 11 \hat{I}^2 -HSD1 inhibitors based on dammarane scaffold. European Journal of Medicinal Chemistry, 2017, 135, 324-338.	5.5	6
10	Identification and characterization of Rubisco activase genes in <i>Oryza punctata</i> . Journal of Systematics and Evolution, 2017, 55, 200-207.	3.1	2
11	New neo -clerodane diterpenoids with neurotrophic activity from the aerial parts of Salvia tiliifolia. Fìtoterapìâ, 2017, 123, 44-50.	2.2	13
12	Total Synthesis of (±)-Cermizine B. Journal of Organic Chemistry, 2017, 82, 11110-11116.	3.2	9
13	A biomimetic semisynthesis enables structural elucidation of selaginellin U: a tautomeric cyclic alkynylphenol from <i>Selaginella tamariscina</i> . Royal Society Open Science, 2017, 4, 170352.	2.4	9
14	Divergent homoeolog evolution of the anthocyanin regulatory gene <i>R</i> in <i>Oryza</i> allopolyploids. Journal of Systematics and Evolution, 2016, 54, 511-518.	3.1	0
15	Extensive reprogramming of cytosine methylation in Oryza allotetraploids. Genes and Genomics, 2015, 37, 517-524.	1.4	4
16	Population genetic structure of Tamarix chinensis in the Yellow River Delta, China. Plant Systematics and Evolution, 2012, 298, 147-153.	0.9	7
17	Parallel up-regulation of the profilin gene family following independent domestication of diploid and allopolyploid cotton (<i>Gossypium</i>). Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 21152-21157.	7.1	61
18	Multiple patterns of rDNA evolution following polyploidy in Oryza. Molecular Phylogenetics and Evolution, 2010, 55, 136-142.	2.7	32

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#	Article	IF	CITATIONS
19	Effects of copy number of an octopine synthase enhancer element and its distance from the TATA box on heterologous expression in transgenic tobacco. Acta Physiologiae Plantarum, 2009, 31, 705-710.	2.1	11
20	Genetic diversity and evolutionary relationships ofOryza species with the B- and C-genomes as revealed by SSR markers. Journal of Plant Biology, 2006, 49, 339-347.	2.1	18
21	Identification of genomic constitutions of Oryza species with the B and C genomes by the PCR-RFLP method. Genetic Resources and Crop Evolution, 2005, 52, 69-76.	1.6	15