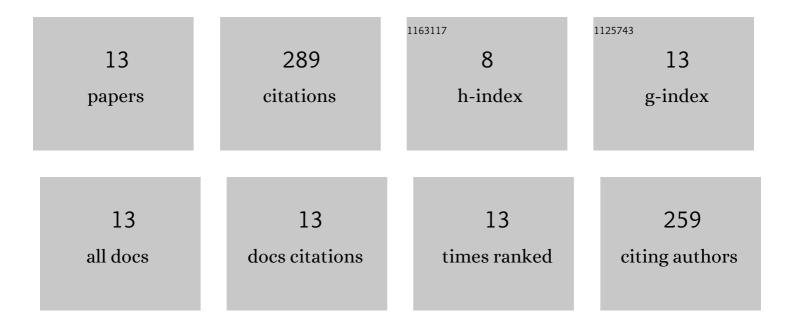
Lixia Zhou

List of Publications by Year in descending order

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Ιιχιλ ΖΗΟΠ

#	Article	IF	CITATIONS
1	The genome draft of coconut (Cocos nucifera). GigaScience, 2017, 6, 1-11.	6.4	96
2	Genome-wide identification of WRKY genes and their expression profiles under different abiotic stresses in Elaeis guineensis. PLoS ONE, 2017, 12, e0189224.	2.5	51
3	Genome-wide identification and expression analysis of MYB gene family in oil palm (Elaeis guineensis) Tj ETQq1 1	0.784314 4.2	• rgBT /Over
4	Coconut genome assembly enables evolutionary analysis of palms and highlights signaling pathways involved in salt tolerance. Communications Biology, 2021, 4, 105.	4.4	26
5	The auxin response factor (ARF) gene family in Oil palm (Elaeis guineensis Jacq.): Genome-wide identification and their expression profiling under abiotic stresses. Protoplasma, 2022, 259, 47-60.	2.1	17
6	Genome-Wide Identification and Characterization of AP2/ERF Transcription Factor Family Genes in Oil Palm under Abiotic Stress Conditions. International Journal of Molecular Sciences, 2021, 22, 2821.	4.1	16
7	miRNAs as key regulators via targeting the phytohormone signaling pathways during somatic embryogenesis of plants. 3 Biotech, 2020, 10, 495.	2.2	13
8	CRISPR/Cas mediated base editing: a practical approach for genome editing in oil palm. 3 Biotech, 2020, 10, 306.	2.2	10
9	Genome-wide identification and expression analysis of bZIP transcription factors in oil palm (Elaeis) Tj ETQq1 1 0.	784314 rg 2.1	gBT_/Overloc
10	The oil palm R2R3-MYB subfamily genes EgMYB111 and EgMYB157 improve multiple abiotic stress tolerance in transgenic Arabidopsis plants. Plant Cell Reports, 2022, 41, 377-393.	5.6	7
11	Sequence-Related Amplified Polymorphism (SRAP) Markers Based Genetic Diversity and Population Structure Analysis of Oil Palm (Elaeis guineensis Jacq.). Tropical Plant Biology, 2021, 14, 63-71.	1.9	5
12	Development of SSR markers based on transcriptome data and association mapping analysis for fruit shell thickness associated traits in oil palm (Elaeis guineensis Jacq.). 3 Biotech, 2020, 10, 280.	2.2	4
13	Identification and expression analysis of histone modification gene (HM) family during somatic embryogenesis of oil palm. BMC Genomics, 2022, 23, 11.	2.8	4