Lixia Zhou

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/295123/publications.pdf

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| | | 1163117 | 1125743 | |
|----------|----------------|--------------|----------------|--|
| 13 | 289 | 8 | 13 | |
| papers | citations | h-index | g-index | |
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| | | | | |
| 13 | 13 | 13 | 259 | |
| all docs | docs citations | times ranked | citing authors | |
| | | | | |

| # | Article | IF | CITATIONS |
|----|--|-----------|----------------|
| 1 | 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 |
| 2 | Genome-wide identification and expression analysis of bZIP transcription factors in oil palm (Elaeis) Tj ETQq0 0 0 | rgBT /Ove | erlock 10 Tf 5 |
| 3 | 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 |
| 4 | Identification and expression analysis of histone modification gene (HM) family during somatic embryogenesis of oil palm. BMC Genomics, 2022, 23, 11. | 2.8 | 4 |
| 5 | 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 |
| 6 | Coconut genome assembly enables evolutionary analysis of palms and highlights signaling pathways involved in salt tolerance. Communications Biology, 2021, 4, 105. | 4.4 | 26 |
| 7 | 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 |
| 8 | miRNAs as key regulators via targeting the phytohormone signaling pathways during somatic embryogenesis of plants. 3 Biotech, 2020, 10, 495. | 2.2 | 13 |
| 9 | Genome-wide identification and expression analysis of MYB gene family in oil palm (Elaeis guineensis) Tj ETQq1 | 0.78431 | 4 rgBT /Overl |
| 10 | 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 |
| 11 | CRISPR/Cas mediated base editing: a practical approach for genome editing in oil palm. 3 Biotech, 2020, 10, 306. | 2.2 | 10 |
| 12 | The genome draft of coconut (Cocos nucifera). GigaScience, 2017, 6, 1-11. | 6.4 | 96 |
| 13 | 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 |