

Mingzhi Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3271608/publications.pdf>

Version: 2024-02-01

11
papers

299
citations

1307594

7
h-index

1281871

11
g-index

11
all docs

11
docs citations

11
times ranked

392
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Complete Chloroplast Genome Sequence and Phylogenetic Analysis of <i>Quercus acutissima</i> . <i>International Journal of Molecular Sciences</i> , 2018, 19, 2443. | 4.1 | 75 |
| 2 | Physiological and transcriptomic analyses reveal a response mechanism to cold stress in <i>Santalum album</i> L. leaves. <i>Scientific Reports</i> , 2017, 7, 42165. | 3.3 | 58 |
| 3 | Involvement of rice histone deacetylase HDA705 in seed germination and in response to ABA and abiotic stresses. <i>Biochemical and Biophysical Research Communications</i> , 2016, 470, 439-444. | 2.1 | 52 |
| 4 | A Genome-Wide Identification of the WRKY Family Genes and a Survey of Potential WRKY Target Genes in <i>Dendrobium officinale</i> . <i>Scientific Reports</i> , 2017, 7, 9200. | 3.3 | 28 |
| 5 | Genome-Wide Characterization, Expression Profile Analysis of WRKY Family Genes in <i>Santalum album</i> and Functional Identification of Their Role in Abiotic Stress. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5676. | 4.1 | 27 |
| 6 | Comparative transcriptome analysis of roots, stems and leaves of <i>Isodon amethystoides</i> reveals candidate genes involved in Wangzaozins biosynthesis. <i>BMC Plant Biology</i> , 2018, 18, 272. | 3.6 | 22 |
| 7 | Selection and Validation of Novel RT-qPCR Reference Genes under Hormonal Stimuli and in Different Tissues of <i>Santalum album</i> . <i>Scientific Reports</i> , 2018, 8, 17511. | 3.3 | 13 |
| 8 | An easy method for identifying 315 categories of commonly-used Chinese herbal medicines based on automated image recognition using AutoML platforms. <i>Informatics in Medicine Unlocked</i> , 2021, 25, 100607. | 3.4 | 8 |
| 9 | Characterization of the complete chloroplast genome sequence of <i>Tsuga longibracteata</i> W. C. Cheng (Pinaceae). <i>Conservation Genetics Resources</i> , 2019, 11, 117-120. | 0.8 | 7 |
| 10 | RNA-Sequencing Analysis Reveals Critical Roles of Hormone Metabolism and Signaling Transduction in Seed Germination of <i>Andrographis paniculata</i> . <i>Journal of Plant Growth Regulation</i> , 2019, 38, 273-282. | 5.1 | 5 |
| 11 | Transcriptomic analysis of seed germination improvement of <i>Andrographis paniculata</i> responding to air plasma treatment. <i>PLoS ONE</i> , 2020, 15, e0240939. | 2.5 | 4 |