

Rasmieh Hamid

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/328183/publications.pdf>

Version: 2024-02-01

9
papers

155
citations

1163117
8
h-index

1474206
9
g-index

9
all docs

9
docs citations

9
times ranked

113
citing authors

#	ARTICLE	IF	CITATIONS
1	Unravelling the treasure trove of drought-responsive genes in wild-type peanut through transcriptomics and physiological analyses of root. <i>Functional and Integrative Genomics</i> , 2022, 22, 215-233.	3.5	10
2	Fertilizer source and chitosan effect on productivity, nutrient accumulation, and phenolic compounds of <i>Thymus daenensis</i> Celak. <i>Agronomy Journal</i> , 2021, 113, 5499-5515.	1.8	3
3	Genic microsatellite marker characterization and development in little millet (<i>Panicum sumatrense</i>) using transcriptome sequencing. <i>Scientific Reports</i> , 2021, 11, 20620.	3.3	18
4	Transcriptome landscaping for gene mining and SSR marker development in Coriander (<i>Coriandrum</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.9	32
5	Comparative RNA-Seq profiling of a resistant and susceptible peanut (<i>Arachis hypogaea</i>) genotypes in response to leaf rust infection caused by <i>Puccinia arachidis</i> . <i>3 Biotech</i> , 2020, 10, 284.	2.2	20
6	Uncloaking lncRNA-mediated gene expression as a potential regulator of CMS in cotton (<i>Gossypium</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.9	20
7	Peanut (<i>Arachis hypogaea</i>) transcriptome revealed the molecular interactions of the defense mechanism in response to early leaf spot fungi (<i>Cercospora arachidicola</i>). <i>Plant Gene</i> , 2020, 23, 100243.	2.3	12
8	Transcriptome analysis identified aberrant gene expression in pollen developmental pathways leading to CGMS in cotton (<i>Gossypium hirsutum</i> L.). <i>PLoS ONE</i> , 2019, 14, e0218381.	2.5	20
9	Transcriptome profiling and cataloging differential gene expression in floral buds of fertile and sterile lines of cotton (<i>Gossypium hirsutum</i> L.). <i>Gene</i> , 2018, 660, 80-91.	2.2	20