

# Allah Ditta

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1794870/publications.pdf>

Version: 2024-02-01

11  
papers

113  
citations

1684129

5  
h-index

1372553

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

100  
citing authors

#	ARTICLE	IF	CITATIONS
1	Dissection of Drought Tolerance in Upland Cotton Through Morpho-Physiological and Biochemical Traits at Seedling Stage. <i>Frontiers in Plant Science</i> , 2021, 12, 627107.	3.6	27
2	Genome-Wide Mining and Identification of Protein Kinase Gene Family Impacts Salinity Stress Tolerance in Highly Dense Genetic Map Developed from Interspecific Cross between <i>G. hirsutum</i> L. and <i>G. darwinii</i> G. Watt. <i>Agronomy</i> , 2019, 9, 560.	3.0	21
3	Assessment of Genetic Diversity, Population Structure, and Evolutionary Relationship of Uncharacterized Genes in a Novel Germplasm Collection of Diploid and Allotetraploid <i>Gossypium</i> Accessions Using EST and Genomic SSR Markers. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2401.	4.1	20
4	Exploring potential of copper and silver nano particles to establish efficient callogenesis and regeneration system for wheat ( <i>Triticum aestivum</i> L.). <i>GM Crops and Food</i> , 2021, 12, 564-585.	3.8	18
5	Identification and characterization of genes related to salt stress tolerance within segregation distortion regions of genetic map in F2 population of upland cotton. <i>PLoS ONE</i> , 2021, 16, e0247593.	2.5	8
6	Genome-Wide Mining and Characterization of SSR Markers for Gene Mapping and Gene Diversity in <i>Gossypium barbadense</i> L. and <i>Gossypium darwinii</i> G. Watt Accessions. <i>Agronomy</i> , 2018, 8, 181.	3.0	6
7	Short communication. Partial resistance of a cotton mutant to Cotton leaf curl Burewala virus. <i>Spanish Journal of Agricultural Research</i> , 2010, 8, 1098.	0.6	4
8	Whole Transcriptome Sequencing Reveals Drought Resistance-Related Genes in Upland Cotton. <i>Genes</i> , 2022, 13, 1159.	2.4	4
9	Genome wide characterization, evolution and expression analysis of FBA gene family under salt stress in <i>Gossypium</i> species. <i>Biologia (Poland)</i> , 2019, 74, 1539-1552.	1.5	3
10	Genome-wide identification and characterization of the CLASP_N gene family in upland cotton ( <i>Gossypium hirsutum</i> L.). <i>PeerJ</i> , 2022, 10, e12733.	2.0	2
11	Genome-wide identification and analysis of the GUB_WAK_bind gene family in <i>Gossypium hirsutum</i> . <i>Molecular Biology Reports</i> , 2022, , 1.	2.3	0