

Hafiz Ghulam Muhu-Din Ahmed

List of Publications by Year in descending order

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13
papers

221
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1307594

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docs citations

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161
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association mapping for high temperature tolerance in wheat through 90k SNP array using physiological and yield traits. PLoS ONE, 2022, 17, e0262569.	2.5	10
2	Genome-wide association mapping of bread wheat genotypes for sustainable food security and yield potential under limited water conditions. PLoS ONE, 2022, 17, e0263263.	2.5	7
3	Molecular characterization and validation of sunflower (<i>Helianthus annuus</i> L.) hybrids through SSR markers. PLoS ONE, 2022, 17, e0267383.	2.5	2
4	Genome-Wide Association Mapping for Stomata and Yield Indices in Bread Wheat under Water Limited Conditions. Agronomy, 2021, 11, 1646.	3.0	13
5	Genome wide association mapping through 90K SNP array against leaf rust pathogen in bread wheat genotypes under field conditions. Journal of King Saud University - Science, 2021, 33, 101628.	3.5	4
6	Genome-Wide Association Mapping through 90K SNP Array for Quality and Yield Attributes in Bread Wheat against Water-Deficit Conditions. Agriculture (Switzerland), 2020, 10, 392.	3.1	29
7	Conferring drought-tolerant wheat genotypes through morpho-physiological and chlorophyll indices at seedling stage. Saudi Journal of Biological Sciences, 2020, 27, 2116-2123.	3.8	38
8	Selection Criteria for Drought-Tolerant Bread Wheat Genotypes at Seedling Stage. Sustainability, 2019, 11, 2584.	3.2	70
9	Early selection of bread wheat genotypes using morphological and photosynthetic attributes conferring drought tolerance. Journal of Integrative Agriculture, 2019, 18, 2483-2491.	3.5	20
10	Genetic analysis of yield and physical traits of spring wheat grain. Journal of the National Science Foundation of Sri Lanka, 2018, 46, 23.	0.2	7
11	Genome wide analysis of heat shock transcription factor (HSF) family in chickpea and its comparison with Arabidopsis. Plant OMICS, 2016, 9, 136-141.	0.4	13
12	Mutation breeding approach to breed drought tolerant maize hybrids. International Journal of Biosciences, 2015, 6, 427-436.	0.1	5
13	Dissection of genetic architecture for tiller angle in rice (<i>Oryza sativa</i> L.) by multiple genome-wide association analyses. PeerJ, 0, 10, e12674.	2.0	3