Mukesh Choudhary

List of Publications by Year in descending order

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840776 552781 33 830 11 26 citations g-index h-index papers 33 33 33 534 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Heavy metal incorporation in foraminiferal calcite under variable environmental and acute level seawater pollution: multi-element culture experiments for Amphisorus hemprichii. Environmental Science and Pollution Research, 2022, 29, 3826-3839.	5.3	2
2	Quantitative Trait Loci for Heat Stress Tolerance in Brassica rapa L. Are Distributed across the Genome and Occur in Diverse Genetic Groups, Flowering Phenologies and Morphotypes. Genes, 2022, 13, 296.	2.4	1
3	How do plants defend themselves against pathogens-Biochemical mechanisms and genetic interventions. Physiology and Molecular Biology of Plants, 2022, 28, 485-504.	3.1	95
4	Impacts, Tolerance, Adaptation, and Mitigation of Heat Stress on Wheat under Changing Climates. International Journal of Molecular Sciences, 2022, 23, 2838.	4.1	55
5	Genetic Diversity, Population Structure and Linkage Disequilibrium Analyses in Tropical Maize Using Genotyping by Sequencing. Plants, 2022, 11, 799.	3.5	11
6	Molecular mechanisms, genetic mapping, and genome editing for insect pest resistance in field crops. Theoretical and Applied Genetics, 2022, 135, 3875-3895.	3.6	12
7	Salinity stress tolerance and omics approaches: revisiting the progress and achievements in major cereal crops. Heredity, 2022, 128, 497-518.	2.6	34
8	Population Structure Analysis and Association Mapping for Turcicum Leaf Blight Resistance in Tropical Maize Using SSR Markers. Genes, 2022, 13, 618.	2.4	9
9	Pangenomics in Microbial and Crop Research: Progress, Applications, and Perspectives. Genes, 2022, 13, 598.	2.4	9
10	Heat Stress during Meiosis Has Lasting Impacts on Plant Growth and Reproduction in Wheat (Triticum) Tj ETQqQ	00 rgBT	/Oyerlock 10
11	Wheat Proteomics for Abiotic Stress Tolerance and Root System Architecture: Current Status and Future Prospects. Proteomes, 2022, 10, 17.	3.5	14
12	Maize microbiome: current insights for the sustainable agriculture., 2021,, 267-297.		10
13	Isolation of genes/quantitative trait loci for drought stress tolerance in maize , 2021, , 267-281.		O
14	Coping with low moisture stress: Remembering and responding. Physiologia Plantarum, 2021, 172, 1162-1169.	5.2	6
15	<i>Heterosis in Genomic Era: Advances in the Molecular Understanding and Techniques for Rapid Exploitation</i> Critical Reviews in Plant Sciences, 2021, 40, 218-242.	5.7	8
16	Skim sequencing: an advanced NGS technology for crop improvement. Journal of Genetics, 2021, 100, 1.	0.7	10
17	Meta-analysis of QTLs associated with popping traits in maize (Zea mays L.). PLoS ONE, 2021, 16, e0256389.	2.5	23
18	Harnessing the Wild Relatives and Landraces for Fe and Zn Biofortification in Wheat through Genetic Interventions—A Review. Sustainability, 2021, 13, 12975.	3.2	3

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19	Nitrogen use efficiency (NUE): elucidated mechanisms, mapped genes and gene networks in maize (Zea) Tj ETQq	1 1.0.7843	14 rgBT /0
20	Enabling technologies for utilization of maize as a bioenergy feedstock. Biofuels, Bioproducts and Biorefining, 2020, 14, 402-416.	3.7	18
21	Genetically modified crops: current status and future prospects. Planta, 2020, 251, 91.	3.2	218
22	Recent Advances in Genomics Assisted Breeding for Drought Stress Tolerance in Major Cereals. Journal of Cereal Research, 2020, 12, .	0.1	12
23	GGE biplot analysis of genotype $\tilde{A}-$ environment interaction and identification of mega-environment for baby corn hybrids evaluation in India. Indian Journal of Genetics and Plant Breeding, 2020, 79, .	0.5	7
24	Green Production Strategies. , 2019, , 492-500.		4
25	QTLian breeding for climate resilience in cereals: progress and prospects. Functional and Integrative Genomics, 2019, 19, 685-701.	3.5	34
26	Marker-Assisted Breeding for Abiotic Stress Tolerance in Crop Plants. , 2018, , 1-23.		8
27	Harnessing Crop Wild Relatives for Crop Improvement. LS International Journal of Life Sciences, 2017, 6, 73.	0.2	17
28	An Overview on Molecular Basis of Genetic Recombination. International Journal of Current Microbiology and Applied Sciences, 2017, 6, 1154-1167.	0.1	1
29	Microsatellite marker-based genetic diversity analyses of novel maize inbreds possessing rare allele of \hat{l}^2 -carotene hydroxylase (crtRB1) for their utilization in \hat{l}^2 -carotene enrichment. Journal of Plant Biochemistry and Biotechnology, 2016, 25, 12-20.	1.7	19
30	DUS characterization and diversity assessment in pearl millet inbreds. Electronic Journal of Plant Breeding, 2016, 7, 925.	0.1	2
31	Popping quality attributes of popcorn hybrids in relation to weevil (Sitophilus oryzae) infestation. Indian Journal of Genetics and Plant Breeding, 2015, 75, 510.	0.5	11
32	Development of \hat{l}^2 -Carotene Rich Maize Hybrids through Marker-Assisted Introgression of \hat{l}^2 -carotene hydroxylase Allele. PLoS ONE, 2014, 9, e113583.	2.5	154
33	Assessment of Genetic Diversity in Rice (Oryza sativa L.) under Irrigated and Drought Stress Condition. Current Journal of Applied Science and Technology, 0, , 112-125.	0.3	2