

Rajesh K Gazara

List of Publications by Year in descending order

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Version: 2024-02-01

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papers

191
citations

1307366

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docs citations

15
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265
citing authors

#	ARTICLE	IF	CITATIONS
1	Reinforcing Synthetic Data for Meticulous Survival Prediction of Patients Suffering From Left Ventricular Systolic Dysfunction. IEEE Access, 2021, 9, 72661-72669.	2.6	6
2	Next-generation sequencing: an expedition from workstation to clinical applications. , 2021, , 29-47.		5
3	Genome-Wide Analysis of the COBRA-Like Gene Family Supports Gene Expansion through Whole-Genome Duplication in Soybean (<i>Glycine max</i>). Plants, 2021, 10, 167.	1.6	10
4	Comparative transcriptome profiling of rice colonized with beneficial endophyte, <i>Piriformospora indica</i> , under high salinity environment. Molecular Biology Reports, 2020, 47, 7655-7673.	1.0	17
5	Connecting the dots: Advances in modern metabolomics and its application in yeast system. Biotechnology Advances, 2020, 44, 107616.	6.0	12
6	Systematic analysis of 1298 RNA-seq samples and construction of a comprehensive soybean (<i>Glycine</i>) Tj ETQq0,0 0 rgBT/Overlock	2.8	54
7	Transcriptional landscape of soybean (<i>Glycine max</i>) embryonic axes during germination in the presence of paclobutrazol, a gibberellin biosynthesis inhibitor. Scientific Reports, 2019, 9, 9601.	1.6	18
8	Expansion and diversification of the gibberellin receptor GIBBERELLIN INSENSITIVE DWARF1 (GID1) family in land plants. Plant Molecular Biology, 2018, 97, 435-449.	2.0	22
9	Transcription Factor Repertoire of Necrotrophic Fungal Phytopathogen <i>Ascochyta rabiei</i> : Predominance of MYB Transcription Factors As Potential Regulators of Secretome. Frontiers in Plant Science, 2017, 8, 1037.	1.7	19
10	Comparative Structural Modeling of Six Old Yellow Enzymes (OYEs) from the Necrotrophic Fungus <i>Ascochyta rabiei</i> : Insight into Novel OYE Classes with Differences in Cofactor Binding, Organization of Active Site Residues and Stereopreferences. PLoS ONE, 2014, 9, e95989.	1.1	22