

# Rajesh K Gazara

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7236233/publications.pdf>

Version: 2024-02-01

10  
papers

191  
citations

1307366

7  
h-index

1474057

9  
g-index

15  
all docs

15  
docs citations

15  
times ranked

265  
citing authors

#	ARTICLE	IF	CITATIONS
1	Systematic analysis of 1298 RNA-seq samples and construction of a comprehensive soybean (<i>Glycine) Tj ETQq1.1 0.784314 rgB	2.8	54
2	Expansion and diversification of the gibberellin receptor GIBBERELLIN INSENSITIVE DWARF1 (GID1) family in land plants. <i>Plant Molecular Biology</i> , 2018, 97, 435-449.	2.0	22
3	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. <i>PLoS ONE</i> , 2014, 9, e95989.	1.1	22
4	Transcription Factor Repertoire of Necrotrophic Fungal Phytopathogen <i>Ascochyta rabiei</i> : Predominance of MYB Transcription Factors As Potential Regulators of Secretome. <i>Frontiers in Plant Science</i> , 2017, 8, 1037.	1.7	19
5	Transcriptional landscape of soybean ( <i>Glycine max</i> ) embryonic axes during germination in the presence of paclobutrazol, a gibberellin biosynthesis inhibitor. <i>Scientific Reports</i> , 2019, 9, 9601.	1.6	18
6	Comparative transcriptome profiling of rice colonized with beneficial endophyte, <i>Piriformospora indica</i> , under high salinity environment. <i>Molecular Biology Reports</i> , 2020, 47, 7655-7673.	1.0	17
7	Connecting the dots: Advances in modern metabolomics and its application in yeast system. <i>Biotechnology Advances</i> , 2020, 44, 107616.	6.0	12
8	Genome-Wide Analysis of the COBRA-Like Gene Family Supports Gene Expansion through Whole-Genome Duplication in Soybean ( <i>Glycine max</i> ). <i>Plants</i> , 2021, 10, 167.	1.6	10
9	Reinforcing Synthetic Data for Meticulous Survival Prediction of Patients Suffering From Left Ventricular Systolic Dysfunction. <i>IEEE Access</i> , 2021, 9, 72661-72669.	2.6	6
10	Next-generation sequencing: an expedition from workstation to clinical applications. , 2021, , 29-47.		5