

# Nibedita Chakraborty

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2798384/publications.pdf>

Version: 2024-02-01

10  
papers

219  
citations

1478505

6  
h-index

1281871

11  
g-index

12  
all docs

12  
docs citations

12  
times ranked

243  
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeting Non-Coding RNAs in Plants with the CRISPR-Cas Technology is a Challenge yet Worth Accepting. <i>Frontiers in Plant Science</i> , 2015, 6, 1001.	3.6	53
2	Genome-wide identification of miRNAs and lncRNAs in <i>Cajanus cajan</i> . <i>BMC Genomics</i> , 2017, 18, 878.	2.8	40
3	Computational prediction of miRNAs and their targets in <i>Phaseolus vulgaris</i> using simple sequence repeat signatures. <i>BMC Plant Biology</i> , 2015, 15, 140.	3.6	38
4	Identification and characterization of differentially expressed <i>Phaseolus vulgaris</i> miRNAs and their targets during mungbean yellow mosaic India virus infection reveals new insight into <i>Phaseolus</i> -MYMIV interaction. <i>Genomics</i> , 2019, 111, 1333-1342.	2.9	26
5	Exogenous application of methyl jasmonate induces defense response and develops tolerance against mungbean yellow mosaic India virus in <i>Vigna mungo</i> . <i>Functional Plant Biology</i> , 2019, 46, 69.	2.1	23
6	Comparative transcriptome profiling of a resistant vs. susceptible <i>Vigna mungo</i> cultivar in response to Mungbean yellow mosaic India virus infection reveals new insight into MYMIV resistance. <i>Current Plant Biology</i> , 2018, 15, 8-24.	4.7	16
7	Molecular and biochemical characterization of mungbean yellow mosaic India virus resistance in leguminous host <i>Vigna mungo</i> . <i>Journal of Plant Biochemistry and Biotechnology</i> , 2018, 27, 318-330.	1.7	8
8	Differential responses of <i>Phaseolus vulgaris</i> cultivars following mungbean yellow mosaic India virus infection. <i>Physiology and Molecular Biology of Plants</i> , 2020, 26, 817-828.	3.1	4
9	Tiny Yet Indispensable Plant MicroRNAs Are Worth to Explore as Key Components for Combating Genotoxic Stresses. <i>Frontiers in Plant Science</i> , 2019, 10, 1197.	3.6	3
10	Genome-wide prediction of cauliflower miRNAs and lncRNAs and their roles in post-transcriptional gene regulation. <i>Planta</i> , 2021, 254, 72.	3.2	2