

List of Publications by Year in descending order

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14
papers

1,365
citations

623574

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h-index

1058333

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14
all docs

14
docs citations

14
times ranked

1614
citing authors

#	ARTICLE	IF	CITATIONS
1	Mechanisms and Functions of Long Non-Coding RNAs at Multiple Regulatory Levels. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5573.	1.8	493
2	Identification of <i>Gossypium hirsutum</i> long non-coding RNAs (lncRNAs) under salt stress. <i>BMC Plant Biology</i> , 2018, 18, 23.	1.6	142
3	Genome-Wide Profiling of miRNAs and Other Small Non-Coding RNAs in the <i>Verticillium dahliae</i> Inoculated Cotton Roots. <i>PLoS ONE</i> , 2012, 7, e35765.	1.1	115
4	Genome-wide characterization and expression analyses of superoxide dismutase (SOD) genes in <i>Gossypium hirsutum</i> . <i>BMC Genomics</i> , 2017, 18, 376.	1.2	101
5	Difference in miRNA expression profiles between two cotton cultivars with distinct salt sensitivity. <i>Molecular Biology Reports</i> , 2012, 39, 4961-4970.	1.0	77
6	Role of plant respiratory burst oxidase homologs in stress responses. <i>Free Radical Research</i> , 2018, 52, 826-839.	1.5	76
7	The long non-coding RNA lncRNA973 is involved in cotton response to salt stress. <i>BMC Plant Biology</i> , 2019, 19, 459.	1.6	70
8	The Catalase Gene Family in Cotton: Genome-Wide Characterization and Bioinformatics Analysis. <i>Cells</i> , 2019, 8, 86.	1.8	57
9	Plant MicroRNAs in Cross-Kingdom Regulation of Gene Expression. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2007.	1.8	53
10	Identification of miRNAs and Their Targets in Cotton Inoculated with <i>Verticillium dahliae</i> by High-Throughput Sequencing and Degradome Analysis. <i>International Journal of Molecular Sciences</i> , 2015, 16, 14749-14768.	1.8	46
11	Long noncoding RNA lncRNA354 functions as a competing endogenous RNA of miR160b to regulate ARF genes in response to salt stress in upland cotton. <i>Plant, Cell and Environment</i> , 2021, 44, 3302-3321.	2.8	46
12	MicroRNA414c affects salt tolerance of cotton by regulating reactive oxygen species metabolism under salinity stress. <i>RNA Biology</i> , 2019, 16, 362-375.	1.5	43
13	Genome-Wide Analysis of the RNA Helicase Gene Family in <i>Gossypium raimondii</i> . <i>International Journal of Molecular Sciences</i> , 2014, 15, 4635-4656.	1.8	24
14	Comprehensive analysis of the <i>Gossypium hirsutum</i> L. respiratory burst oxidase homolog (Ghrboh) gene family. <i>BMC Genomics</i> , 2020, 21, 91.	1.2	22