Deshika Kohli

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9997689/publications.pdf

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		1478505	1372567	
11	251	6	10	
papers	citations	h-index	g-index	
11	11	11	365	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Identification and Characterization of Wilt and Salt Stress-Responsive MicroRNAs in Chickpea through High-Throughput Sequencing. PLoS ONE, 2014, 9, e108851.	2.5	101
2	The CarERF genes in chickpea (Cicer arietinum L.) and the identification of CarERF116 as abiotic stress responsive transcription factor. Functional and Integrative Genomics, 2015, 15, 27-46.	3.5	47
3	Genome-wide analysis identifies chickpea (Cicer arietinum) heat stress transcription factors (Hsfs) responsive to heat stress at the pod development stage. Journal of Plant Research, 2018, 131, 525-542.	2.4	32
4	Conferring root-knot nematode resistance via host-delivered RNAi-mediated silencing of four Mi-msp genes in Arabidopsis. Plant Science, 2020, 298, 110592.	3.6	24
5	Development of nematode resistance in Arabidopsis by HD-RNAi-mediated silencing of the effector gene Mi-msp2. Scientific Reports, 2019, 9, 17404.	3.3	17
6	Characterization of root-knot nematode responsive and root-specific promoter containing PIN domain from <i>Arabidopsis thaliana</i> (L.) Heynh. Indian Journal of Genetics and Plant Breeding, 2016, 76, 75.	0.5	8
7	Host delivered-RNAi of effector genes for imparting resistance against root-knot and cyst nematodes in plants. Physiological and Molecular Plant Pathology, 2022, 118, 101802.	2.5	6
8	Host-mediated RNAi of a Notch-like receptor gene inMeloidogyne incognitainduces nematode resistance. Parasitology, 2018, 145, 1896-1906.	1.5	5
9	RNAi for Resistance Against Biotic Stresses in Crop Plants. , 2018, , 67-112.		5
10	A Polyketide cyclase/dehydrase and lipid transport superfamily gene of Arabidopsis and its orthologue of chickpea exhibit rapid response to wounding. Indian Journal of Genetics and Plant Breeding, 2014, 74, 463.	0.5	5
11	Quantitative detection of pathogen load of Fusarium oxysporum f.sp. ciceris infected wilt resistant and susceptible genotypes of chickpea using intergenic spacer region-based marker. Physiological and Molecular Plant Pathology, 2021, 114, 101622.	2.5	1