

Hai Nian

List of Publications by Year in descending order

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papers

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#	ARTICLE	IF	CITATIONS
1	A Single Nucleotide Deletion in J Encoding GmELF3 Confers Long Juvenility and Is Associated with Adaption of Tropic Soybean. <i>Molecular Plant</i> , 2017, 10, 656-658.	8.3	96
2	Fine mapping of a Phytophthora-resistance gene RpsWY in soybean (<i>Glycine max L.</i>) by high-throughput genome-wide sequencing. <i>Theoretical and Applied Genetics</i> , 2017, 130, 1041-1051.	3.6	80
3	Construction of high-density genetic map and QTL mapping of yield-related and two quality traits in soybean RILs population by RAD-sequencing. <i>BMC Genomics</i> , 2017, 18, 466.	2.8	51
4	Fine-mapping of QTLs for individual and total isoflavone content in soybean (<i>Glycine max L.</i>) using a high-density genetic map. <i>Theoretical and Applied Genetics</i> , 2018, 131, 555-568.	3.6	34
5	Fine mapping of a Phytophthora-resistance locus RpsGZ in soybean using genotyping-by-sequencing. <i>BMC Genomics</i> , 2020, 21, 280.	2.8	20
6	Genetic mapping of powdery mildew resistance genes in soybean by high-throughput genome-wide sequencing. <i>Theoretical and Applied Genetics</i> , 2019, 132, 1833-1845.	3.6	17
7	QTL fine-mapping of soybean (<i>Glycine max L.</i>) leaf type associated traits in two RILs populations. <i>BMC Genomics</i> , 2019, 20, 260.	2.8	17
8	QTL mapping for soybean (<i>Glycine max L.</i>) leaf chlorophyll-content traits in a genotyped RIL population by using RAD-seq based high-density linkage map. <i>BMC Genomics</i> , 2020, 21, 739.	2.8	14
9	Fine mapping of an adult-plant resistance gene to powdery mildew in soybean cultivar Zhonghuang 24. <i>Crop Journal</i> , 2022, 10, 1103-1110.	5.2	7
10	Fine mapping and genetic analysis of resistance genes, Rsc18, against soybean mosaic virus. <i>Journal of Integrative Agriculture</i> , 2022, 21, 644-653.	3.5	4
11	Fine mapping and analyses of the RSC15ZH resistance candidate gene for the soybean mosaic virus. <i>Euphytica</i> , 2020, 216, 1.	1.2	2