

Qingyuan Li

List of Publications by Year in descending order

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Version: 2024-02-01

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papers

562
citations

1040056

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13
times ranked

666
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of FAD2 and FAD3 genes in Brassica napus genome and development of allele-specific markers for high oleic and low linolenic acid contents. Theoretical and Applied Genetics, 2012, 125, 715-729.	3.6	154
2	Comparative transcriptomic analysis uncovers the complex genetic network for resistance to Sclerotinia sclerotiorum in Brassica napus. Scientific Reports, 2016, 6, 19007.	3.3	126
3	Mapping of quantitative trait loci and development of allele-specific markers for seed weight in Brassica napus. Theoretical and Applied Genetics, 2010, 121, 1289-1301.	3.6	99
4	RNA-seq based transcriptomic analysis uncovers $\hat{\pm}$ -linolenic acid and jasmonic acid biosynthesis pathways respond to cold acclimation in Camellia japonica. Scientific Reports, 2016, 6, 36463.	3.3	66
5	Expression of <i>Brassica napus</i> TTG2, a regulator of trichome development, increases plant sensitivity to salt stress by suppressing the expression of auxin biosynthesis genes. Journal of Experimental Botany, 2015, 66, 5821-5836.	4.8	39
6	Prediction of Anticancer Peptides Using a Low-Dimensional Feature Model. Frontiers in Bioengineering and Biotechnology, 2020, 8, 892.	4.1	30
7	SMRT sequencing of a full-length transcriptome reveals transcript variants involved in C18 unsaturated fatty acid biosynthesis and metabolism pathways at chilling temperature in Pennisetum giganteum. BMC Genomics, 2020, 21, 52.	2.8	16
8	Development of genic SSR marker resources from RNA-seq data in Camellia japonica and their application in the genus Camellia. Scientific Reports, 2021, 11, 9919.	3.3	16
9	Identification and Classification of Enhancers Using Dimension Reduction Technique and Recurrent Neural Network. Computational and Mathematical Methods in Medicine, 2020, 2020, 1-9.	1.3	10
10	Identification and classification of promoters using the attention mechanism based on long short-term memory. Frontiers of Computer Science, 2022, 16, .	2.4	6