

Xiaoming Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7749271/publications.pdf>

Version: 2024-02-01

9
papers

728
citations

1163117
8
h-index

1474206
9
g-index

9
all docs

9
docs citations

9
times ranked

829
citing authors

#	ARTICLE	IF	CITATIONS
1	Natural variation at the soybean J locus improves adaptation to the tropics and enhances yield. <i>Nature Genetics</i> , 2017, 49, 773-779.	21.4	341
2	Resequencing of 683 common bean genotypes identifies yield component trait associations across a north-south cline. <i>Nature Genetics</i> , 2020, 52, 118-125.	21.4	81
3	<i>GmCOL1a</i> and <i>GmCOL1b</i> Function as Flowering Repressors in Soybean Under Long-Day Conditions. <i>Plant and Cell Physiology</i> , 2015, 56, 2409-2422.	3.1	73
4	Overexpression of <i>GmFDL19</i> enhances tolerance to drought and salt stresses in soybean. <i>PLoS ONE</i> , 2017, 12, e0179554.	2.5	69
5	Quantitative Trait Locus Mapping of Flowering Time and Maturity in Soybean Using Next-Generation Sequencing-Based Analysis. <i>Frontiers in Plant Science</i> , 2018, 9, 995.	3.6	57
6	<i>GmmiR156b</i> overexpression delays flowering time in soybean. <i>Plant Molecular Biology</i> , 2015, 89, 353-363.	3.9	49
7	Quantitative Trait Locus Mapping of Soybean Maturity Gene <i>E6</i> . <i>Crop Science</i> , 2017, 57, 2547-2554.	1.8	29
8	QTL mapping for flowering time in different latitude in soybean. <i>Euphytica</i> , 2015, 206, 725-736.	1.2	27
9	QTL mapping and identification of genes associated with the resistance to <i>Acanthoscelides obtectus</i> in cultivated common bean using a high-density genetic linkage map. <i>BMC Plant Biology</i> , 2022, 22, .	3.6	2