Zhenhai Cui

List of Publications by Year in descending order

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ΖΗΕΝΗΛΙ ΟΙΙΙ

#	Article	IF	CITATIONS
1	Effect of Trait Heritability, Training Population Size and Marker Density on Genomic Prediction Accuracy Estimation in 22 bi-parental Tropical Maize Populations. Frontiers in Plant Science, 2017, 8, 1916.	3.6	145
2	Genome-wide association study (GWAS) reveals the genetic architecture of four husk traits in maize. BMC Genomics, 2016, 17, 946.	2.8	59
3	Linkage mapping combined with association analysis reveals QTL and candidate genes for three husk traits in maize. Theoretical and Applied Genetics, 2018, 131, 2131-2144.	3.6	19
4	ZmRAD51C Is Essential for Double-Strand Break Repair and Homologous Recombination in Maize Meiosis. International Journal of Molecular Sciences, 2019, 20, 5513.	4.1	17
5	Genome-Wide Association Study Dissects the Genetic Architecture of Maize Husk Tightness. Frontiers in Plant Science, 2020, 11, 861.	3.6	17
6	Assessment of the Potential for Genomic Selection To Improve Husk Traits in Maize. G3: Genes, Genomes, Genetics, 2020, 10, 3741-3749.	1.8	16
7	Denser Markers and Advanced Statistical Method Identified More Genetic Loci Associated with Husk Traits in Maize. Scientific Reports, 2020, 10, 8165.	3.3	12
8	Identification of maize brace-root quantitative trait loci in a recombinant inbred line population. Euphytica, 2018, 214, 1.	1.2	7
9	miRNA expression profiling and zeatin dynamic changes in a new model system of in vivo indirect regeneration of tomato. PLoS ONE, 2020, 15, e0237690.	2.5	6
10	A Genome-Wide Association Study Dissects the Genetic Architecture of the Metaxylem Vessel Number in Maize Brace Roots. Frontiers in Plant Science, 2022, 13, 847234.	3.6	5
11	Genetic analysis of maize shank length by QTL mapping in three recombinant inbred line populations. Plant Science, 2021, 303, 110767.	3.6	4
12	Genetic analysis of three maize husk traits by QTL mapping in a maize-teosinte population. BMC Genomics, 2021, 22, 386.	2.8	4
13	Transcriptional analyses of maize leaves in response to highâ€density planting. Agronomy Journal, 0, ,	1.8	1
14	Genetic basis of maize ear angle revealed by high-density single nucleotide polymorphism markers in four recombinant inbred line populations. Euphytica, 2020, 216, 1.	1.2	0