

Daizhen Sun

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Quantitative trait loci mapping and candidate gene analysis of stoma-related traits in wheat (<i>Triticum aestivum</i> L.) glumes. PeerJ, 2022, 10, e13262.	2.0	1
2	In Silico and Transcription Analysis of Trehalose-6-phosphate Phosphatase Gene Family of Wheat: Trehalose Synthesis Genes Contribute to Salinity, Drought Stress and Leaf Senescence. Genes, 2021, 12, 1652.	2.4	7
3	Genome-wide identification and characterization of long non-coding RNAs related to grain yield in foxtail millet [<i>Setaria italica</i> (L.) P. Beauv.]. BMC Genomics, 2020, 21, 853.	2.8	6
4	QTL mapping and candidate gene analysis of seed vigor-related traits during artificial aging in wheat (<i>Triticum aestivum</i>). Scientific Reports, 2020, 10, 22060.	3.3	18
5	Mining the stable quantitative trait loci for agronomic traits in wheat (<i>Triticum aestivum</i> L.) based on an introgression line population. BMC Plant Biology, 2020, 20, 275.	3.6	6
6	QTL mapping for flag leaf-related traits and genetic effect of QFLW-6A on flag leaf width using two related introgression line populations in wheat. PLoS ONE, 2020, 15, e0229912.	2.5	11
7	Identification of TaPPH-7A haplotypes and development of a molecular marker associated with important agronomic traits in common wheat. BMC Plant Biology, 2019, 19, 296.	3.6	20
8	QTL analysis of wheat kernel traits, and genetic effects of qKW-6A on kernel width. Euphytica, 2019, 215, 1.	1.2	18
9	Association of yield-related traits in founder genotypes and derivatives of common wheat (<i>Triticum</i>) Tj ETQq1 1 0.784314 rgBT /Overl	3.6	28