Shahjahan Ali

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/10751061/publications.pdf

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15	1,101	933447	1125743
papers	citations	10 h-index	13 g-index

15	15	15	2049
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Identification of QTL Associated with Regrowth Vigor Using the Nested Association Mapping Population in Switchgrass. Plants, 2022, 11, 566.	3.5	2
2	Genomic regions associated with salinity tolerance in lowland switchgrass. Crop Science, 2021, 61, 4022-4037.	1.8	3
3	Genomeâ€wide quantitative trait loci detection for biofuel traits in switchgrass (Panicum) Tj ETQq1 1 0.784314 rgBT /Overlock 10	5.6	3
4	Highâ€density linkage map reveals QTL underlying growth traits in AP13Ã—VS16 biparental population of switchgrass. GCB Bioenergy, 2019, 11, 672-690.	5.6	13
5	Viral Metagenomics: Analysis of Begomoviruses by Illumina High-Throughput Sequencing. Viruses, 2014, 6, 1219-1236.	3.3	69
6	Genome-wide analysis of alternative splicing of pre-mRNA under salt stress in Arabidopsis. BMC Genomics, 2014, 15, 431.	2.8	234
7	A KH-Domain RNA-Binding Protein Interacts with FIERY2/CTD Phosphatase-Like 1 and Splicing Factors and Is Important for Pre-mRNA Splicing in Arabidopsis. PLoS Genetics, 2013, 9, e1003875.	3.5	88
8	The genome of the extremophile crucifer Thellungiella parvula. Nature Genetics, 2011, 43, 913-918.	21.4	318
9	Genome Structures and Halophyte-Specific Gene Expression of the Extremophile Thellungiella parvula in Comparison with Thellungiella salsuginea (Thellungiella halophila) and Arabidopsis. Plant Physiology, 2010, 154, 1040-1052.	4.8	97
10	Arabidopsis cytochrome P450s through the looking glass: a window on plant biochemistry. Phytochemistry Reviews, 2006, 5, 205-237.	6.5	76
11	Untranslated Regions from C4 Amaranth AhRbcS1 mRNAs Confer Translational Enhancement and Preferential Bundle Sheath Cell Expression in Transgenic C4Flaveria bidentis. Plant Physiology, 2004, 136, 3550-3561.	4.8	35
12	Isolation of gibberellin metabolic pathway genes from barley and comparative mapping in barley, wheat and rice. Theoretical and Applied Genetics, 2004, 109, 847-855.	3.6	86
13	The 3' non-coding region of a C4 photosynthesis gene increases transgene expression when combined with heterologous promoters. , 2001, 46, 325-334.		37
14	Quantitative regulation of the Flaveria Me1 gene is controlled by the 3'-untranslated region and sequences near the amino terminus. , 2001, 46, 251-261.		25
15	Normalisation of cereal endosperm EST libraries for structural and functional genomic analysis. Plant Molecular Biology Reporter, 2000, 18, 123-132.	1.8	15