

Nestor Kippes

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

585
citations

1040056

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h-index

1281871

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13
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13
docs citations

13
times ranked

785
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of the <i>VERNALIZATION 4</i> gene reveals the origin of spring growth habit in ancient wheats from South Asia. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E5401-10.	7.1	131
2	Genetic and physical mapping of the earliness per se locus Eps-A m 1 in <i>Triticum monococcum</i> identifies EARLY FLOWERING 3 (ELF3) as a candidate gene. Functional and Integrative Genomics, 2016, 16, 365-382.	3.5	102
3	Antitumor effects of hyaluronic acid inhibitor 4-methylumbelliferone in an orthotopic hepatocellular carcinoma model in mice. Glycobiology, 2012, 22, 400-410.	2.5	91
4	RNA-seq studies using wheat PHYTOCHROME B and PHYTOCHROME C mutants reveal shared and specific functions in the regulation of flowering and shade-avoidance pathways. BMC Plant Biology, 2016, 16, 141.	3.6	67
5	Fine mapping and epistatic interactions of the vernalization gene VRN-D4 in hexaploid wheat. Molecular Genetics and Genomics, 2014, 289, 47-62.	2.1	48
6	Lack of the Matricellular Protein SPARC (Secreted Protein, Acidic and Rich in Cysteine) Attenuates Liver Fibrogenesis in Mice. PLoS ONE, 2013, 8, e54962.	2.5	43
7	Single nucleotide polymorphisms in a regulatory site of VRN-A1 first intron are associated with differences in vernalization requirement in winter wheat. Molecular Genetics and Genomics, 2018, 293, 1231-1243.	2.1	37
8	Development and characterization of a spring hexaploid wheat line with no functional VRN2 genes. Theoretical and Applied Genetics, 2016, 129, 1417-1428.	3.6	35
9	Effect of phyB and phyC loss-of-function mutations on the wheat transcriptome under short and long day photoperiods. BMC Plant Biology, 2020, 20, 297.	3.6	27
10	Diploid mint (<i>M. longifolia</i>) can produce spearmint type oil with a high yield potential. Scientific Reports, 2021, 11, 23521.	3.3	2
11	Efficient construction of a linkage map and haplotypes for <i>Mentha suaveolens</i> using sequence capture. G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	1