Chiara Campoli

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1345483/publications.pdf

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		1040056	1372567	
10	825	9	10	
papers	citations	h-index	g-index	
11	11	11	1108	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Mapping-by-Sequencing Identifies <i>HvPHYTOCHROME C</i> as a Candidate Gene for the <i>early maturity 5</i> Locus Modulating the Circadian Clock and Photoperiodic Flowering in Barley. Genetics, 2014, 198, 383-396.	2.9	102
2	Genetic Control of Reproductive Development in Temperate Cereals. Advances in Botanical Research, 2014, 72, 131-158.	1.1	28
3	Genetic Control of Reproductive Development. Biotechnology in Agriculture and Forestry, 2014, , 81-99.	0.2	3
4	<i>Hv<scp>LUX</scp>1</i> is a candidate gene underlying the <i>early maturity 10</i> locus in barley: phylogeny, diversity, and interactions with the circadian clock and photoperiodic pathways. New Phytologist, 2013, 199, 1045-1059.	7.3	110
5	Expression conservation within the circadian clock of a monocot: natural variation at barley Ppd-H1affects circadian expression of flowering time genes, but not clock orthologs. BMC Plant Biology, 2012, 12, 97.	3.6	125
6	Functional characterisation of <i>HvCO1</i> , the barley (<i>Hordeum vulgare</i>) flowering time ortholog of <i>CONSTANS</i> . Plant Journal, 2012, 69, 868-880.	5.7	136
7	Comparative expression of Cbf genes in the Triticeae under different acclimation induction temperatures. Molecular Genetics and Genomics, 2009, 282, 141-152.	2.1	70
8	Parallel pigment and transcriptomic analysis of four barley Albina and Xantha mutants reveals the complex network of the chloroplast-dependent metabolism. Plant Molecular Biology, 2009, 71, 173-191.	3.9	17
9	Photosynthetic Antenna Size in Higher Plants Is Controlled by the Plastoquinone Redox State at the Post-transcriptional Rather than Transcriptional Level. Journal of Biological Chemistry, 2007, 282, 29457-29469.	3.4	69
10	Transcriptome Analysis of Cold Acclimation in Barley Albina and Xantha Mutants. Plant Physiology, 2006, 141, 257-270.	4.8	164