Sergei A Filichkin

List of Publications by Year in descending order

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		932766	1125271	
13	1,775 citations	10	13	
papers	citations	h-index	g-index	
1.4	1.4	1.4	2764	
14	14	14	2764	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Identification of transcription factors from NF-Y, NAC, and SPL families responding to osmotic stress in multiple tomato varieties. Plant Science, 2018, 274, 441-450.	1.7	9
2	Abiotic Stresses Modulate Landscape of Poplar Transcriptome via Alternative Splicing, Differential Intron Retention, and Isoform Ratio Switching. Frontiers in Plant Science, 2018, 9, 5.	1.7	122
3	DNase I SIM: A Simplified In-Nucleus Method for DNase I Hypersensitive Site Sequencing. Methods in Molecular Biology, 2017, 1629, 141-154.	0.4	1
4	Small Genetic Circuits and MicroRNAs: Big Players in Polymerase II Transcriptional Control in Plants. Plant Cell, 2016, 28, 286-303.	3.1	38
5	Improved DNase-seq protocol facilitates high resolution mapping of DNase I hypersensitive sites in roots in Arabidopsis thaliana. Plant Methods, 2015, 11, 42.	1.9	20
6	Environmental Stresses Modulate Abundance andÂTiming of Alternatively Spliced Circadian Transcripts in Arabidopsis. Molecular Plant, 2015, 8, 207-227.	3.9	142
7	The cyclophilin A DIAGEOTROPICA gene affects auxin transport in both root and shoot to control lateral root formation. Development (Cambridge), 2015, 142, 712-21.	1.2	57
8	Alternative splicing in plants: directing traffic at the crossroads of adaptation and environmental stress. Current Opinion in Plant Biology, 2015, 24, 125-135.	3.5	215
9	Environmental Stresses Modulate Abundance and Timing of Alternatively Spliced Circadian Transcripts in Arabidopsis. Molecular Plant, 2014, , .	3.9	9
10	Unproductive alternative splicing and nonsense mRNAs: A widespread phenomenon among plant circadian clock genes. Biology Direct, 2012, 7, 20.	1.9	125
11	Global Profiling of Rice and Poplar Transcriptomes Highlights Key Conserved Circadian-Controlled Pathways and cis-Regulatory Modules. PLoS ONE, 2011, 6, e16907.	1.1	188
12	Genome-wide mapping of alternative splicing in <i>Arabidopsis thaliana</i> . Genome Research, 2010, 20, 45-58.	2.4	825
13	Efficiency of gene silencing in Arabidopsis: direct inverted repeats vs. transitive RNAi vectors. Plant Biotechnology Journal, 2007, 5, 615-626.	4.1	23