

Leonardo Miguel Galindo

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

14
papers

627
citations

10
h-index

17
g-index

17
ext. papers

819
ext. citations

4.7
avg, IF

3.82
L-index

#	Paper	IF	Citations
14	The genome of flax (<i>Linum usitatissimum</i>) assembled de novo from short shotgun sequence reads. <i>Plant Journal</i> , 2012 , 72, 461-73	6.9	315
13	LTR-retrotransposons in plants: Engines of evolution. <i>Gene</i> , 2017 , 626, 14-25	3.8	101
12	RNA-seq Transcriptome Response of Flax (L.) to the Pathogenic Fungus f. sp.. <i>Frontiers in Plant Science</i> , 2016 , 7, 1766	6.2	42
11	Integrated transcriptomic and proteomic profiling of white spruce stems during the transition from active growth to dormancy. <i>Plant, Cell and Environment</i> , 2012 , 35, 682-701	8.4	38
10	Identification, characterization and distribution of transposable elements in the flax (<i>Linum usitatissimum</i> L.) genome. <i>BMC Genomics</i> , 2012 , 13, 644	4.5	25
9	Influence of water deficit on the molecular responses of <i>Pinus contorta</i> [Pinus banksiana mature trees to infection by the mountain pine beetle fungal associate, <i>Grosmannia clavigera</i> . <i>Tree Physiology</i> , 2014 , 34, 1220-39	4.2	22
8	Shaping Plant Adaptability, Genome Structure and Gene Expression through Transposable Element Epigenetic Control: Focus on Methylation. <i>Agronomy</i> , 2018 , 8, 180	3.6	19
7	Ion Torrent sequencing as a tool for mutation discovery in the flax (<i>Linum usitatissimum</i> L.) genome. <i>Plant Methods</i> , 2015 , 11, 19	5.8	16
6	Response of to Involves Salicylic Acid-Mediated Immunity: An RNA-Seq-Based Study. <i>Frontiers in Plant Science</i> , 2020 , 11, 1025	6.2	15
5	Isolation and characterization of RNase LTR sequences of Ty1-copia retrotransposons in common bean (<i>Phaseolus vulgaris</i> L). <i>Genome</i> , 2004 , 47, 84-95	2.4	11
4	Diverse chitinases are invoked during the activity-dormancy transition in spruce. <i>Tree Genetics and Genomes</i> , 2015 , 11, 1	2.1	9
3	Comparative Transcriptome Analysis of Rutabaga () Cultivars Indicates Activation of Salicylic Acid and Ethylene-Mediated Defenses in Response to. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	7
2	Ty1-copia elements reveal diverse insertion sites linked to polymorphisms among flax (<i>Linum usitatissimum</i> L.) accessions. <i>BMC Genomics</i> , 2016 , 17, 1002	4.5	7
1	Candidate Effectors of Pathotype 5X During Infection of Two Genotypes. <i>Frontiers in Microbiology</i> , 2021 , 12, 742268	5.7	0