MÃ³nica Sebastiana

List of Publications by Year
in descending order

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23	897	471509	677142
papers	citations	17	22
		h-index	g-index

23	23	23	1504
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	First Insights into the Effect of Mycorrhizae on the Expression of Pathogen Effectors during the Infection of Grapevine with Plasmopara viticola. Sustainability, 2021, 13, 1226.	3.2	17
2	Pisolithus. , 2020, , 707-726.		1
3	The leaf lipid composition of ectomycorrhizal oak plants shows a drought-tolerance signature. Plant Physiology and Biochemistry, 2019, 144, 157-165.	5.8	29
4	Ectomycorrhizal inoculation with Pisolithus tinctorius reduces stress induced by drought in cork oak. Mycorrhiza, 2018, 28, 247-258.	2.8	40
5	A genomic study on mammary gland acclimatization to tropical environment in the Holstein cattle. Tropical Animal Health and Production, 2018, 50, 187-195.	1.4	3
6	Specific adjustments in grapevine leaf proteome discriminating resistant and susceptible grapevine genotypes to Plasmopara viticola. Journal of Proteomics, 2017, 152, 48-57.	2.4	41
7	Oak protein profile alterations upon root colonization by an ectomycorrhizal fungus. Mycorrhiza, 2017, 27, 109-128.	2.8	25
8	Genomic study of the mammary gland in bovines acclimated to a tropical environment. South African Journal of Animal Sciences, 2016, 46, 1.	0.5	3
9	Metabolite extraction for high-throughput FTICR-MS-based metabolomics of grapevine leaves. EuPA Open Proteomics, 2016, 12, 4-9.	2.5	35
10	Tracking cashew economically important diseases in the West African region using metagenomics. Frontiers in Plant Science, 2015, 6, 482.	3.6	21
11	Castanea root transcriptome in response to Phytophthora cinnamomi challenge. Tree Genetics and Genomes, 2015, 11, 1.	1.6	72
12	First clues on a jasmonic acid role in grapevine resistance against the biotrophic fungus Plasmopara viticola. European Journal of Plant Pathology, 2015, 142, 645-652.	1.7	33
13	Subtilisin-like proteases in plantÃ¢â‚¬â€œpathogen recognition and immune priming: a perspective. Frontiers in Plant Science, 2014, 5, 739.	3.6	135
14	A comprehensive assessment of the transcriptome of cork oak (Quercus suber) through EST sequencing. BMC Genomics, 2014, 15, 371.	2.8	53
15	Oak Root Response to Ectomycorrhizal Symbiosis Establishment: RNA-Seq Derived Transcript Identification and Expression Profiling. PLoS ONE, 2014, 9, e98376.	2.5	45
16	A possible approach for gel-based proteomic studies in recalcitrant woody plants. SpringerPlus, 2013, 2, 210.	1.2	13
17	Ectomycorrhizal inoculation with Pisolithus tinctorius increases the performance of Quercus suber L. (cork oak) nursery and field seedlings. New Forests, 2013, 44, 937-949.	1.7	42
18	Reference Gene Selection and Validation for the Early Responses to Downy Mildew Infection in Susceptible and Resistant Vitis vinifera Cultivars. PLoS ONE, 2013, 8, e72998.	2.5	78

#	ARTICLE	IF	CITATIONS
19	Labellum transcriptome reveals alkene biosynthetic genes involved in orchid sexual deception and pollination-induced senescence. Functional and Integrative Genomics, 2012, 12, 693-703.	3.5	11
20	Identification of plant genes involved on the initial contact between ectomycorrhizal symbionts (Castanea sativa â€“ European chestnut and Pisolithus tinctorius). European Journal of Soil Biology, 2009, 45, 275-282.	3.2	23
21	Fungal Transcript Pattern During the Preinfection Stage (12Âh) of Ectomycorrhiza Formed Between Pisolithus tinctorius and Castanea sativa Roots, Identified Using cDNA Microarrays. Current Microbiology, 2008, 57, 620-625.	2.2	19
22	Organogenic nodule development in hop (Humulus lupulus L.): Transcript and metabolic responses. BMC Genomics, 2008, 9, 445.	2.8	17
23	Transcriptional and metabolic profiling of grape (Vitis vinifera L.) leaves unravel possible innate resistance against pathogenic fungi. Journal of Experimental Botany, 2008, 59, 3371-3381.	4.8	141