ViÂ-ctor M Guerrero-Sanchez

List of Publications by Year
in descending order

14 papers	229 citations	1040056 9 h-index	1199594 12 g-index
14 all docs	14 docs citations	14 times ranked	148 citing authors

#	ARTICLE	IF	CITATIONS
1	Holm Oak (Quercus ilex) Transcriptome. De novo Sequencing and Assembly Analysis. Frontiers in Molecular Biosciences, 2017, 4, 70.	3.5	46
2	A Multi-Omics Analysis Pipeline for the Metabolic Pathway Reconstruction in the Orphan Species Quercus ilex. Frontiers in Plant Science, 2018, 9, 935.	3.6	37
3	Ion Torrent and Illumina, two complementary RNA-seq platforms for constructing the holm oak (Quercus ilex) transcriptome. PLoS ONE, 2019, 14, e0210356.	2.5	28
4	Proteomics, Holm Oak (Quercus ilex L.) and Other Recalcitrant and Orphan Forest Tree Species: How do They See Each Other?. International Journal of Molecular Sciences, 2019, 20, 692.	4.1	20
5	Quercus ilex pollen allergen, Que i 1, responsible for pollen food allergy syndrome caused by fruits in Spanish allergic patients. Clinical and Experimental Allergy, 2020, 50, 815-823.	2.9	20
6	Responses and Differences in Tolerance to Water Shortage under Climatic Dryness Conditions in Seedlings from Quercus spp. and Andalusian Q. ilex Populations. Forests, 2020, 11, 707.	2.1	19
7	Proteomics Data Analysis for the Identification of Proteins and Derived Proteotypic Peptides of Potential Use as Putative Drought Tolerance Markers for Quercus ilex. International Journal of Molecular Sciences, 2021, 22, 3191.	4.1	13
8	Changes in the transcript and protein profiles of Quercus ilex seedlings in response to drought stress. Journal of Proteomics, 2021, 243, 104263.	2.4	13
9	Untargeted MS-Based Metabolomics Analysis of the Responses to Drought Stress in Quercus ilex L. Leaf Seedlings and the Identification of Putative Compounds Related to Tolerance. Forests, 2022, 13, 551.	2.1	13
10	Protein Interaction Networks: Functional and Statistical Approaches. Methods in Molecular Biology, 2020, 2139, 21-56.	0.9	7
11	Recent Advances in MS-Based Plant Proteomics: Proteomics Data Validation Through Integration with Other Classic and -Omics Approaches. Progress in Botany Fortschritte Der Botanik, 2019, , 77-101.	0.3	6
12	Specific Protein Database Creation from Transcriptomics Data in Nonmodel Species: Holm Oak (Quercus ilex L.). Methods in Molecular Biology, 2020, 2139, 57-68.	0.9	3
13	Identification of Proteases and Protease Inhibitors in Seeds of the Recalcitrant Forest Tree Species Quercus ilex. Frontiers in Plant Science, 0, 13, .	3.6	3
14	Proteomics Analysis of Plant Tissues Based on Two-Dimensional Gel Electrophoresis. , 2018, , 309-322.		1