

Margherita Lucchin

List of Publications by Year in descending order

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14
papers

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#	ARTICLE	IF	CITATIONS
1	Molecular Relationships and Genetic Diversity Analysis of Venetian Radicchio (Leaf Chicory), Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	3.7	3
2	Dioecy in Flowering Plants: From the First Observations of Prospero Alpini in the XVI Century to the Most Recent Advances in the Genomics Era. Agriculture (Switzerland), 2022, 12, 364.	3.1	1
3	The grapevine (<i>Vitis vinifera</i> L.) floral transcriptome in Pinot noir variety: identification of tissue-related gene networks and whorl-specific markers in pre- and post-anthesis phases. Horticulture Research, 2021, 8, 200.	6.3	5
4	The Effect of Soil on the Biochemical Plasticity of Berry Skin in Two Italian Grapevine (<i>V. vinifera</i> L.) Cultivars. Frontiers in Plant Science, 2020, 11, 822.	3.6	7
5	Genomics of Flower Identity in Grapevine (<i>Vitis vinifera</i> L.). Frontiers in Plant Science, 2019, 10, 316.	3.6	22
6	Combinatorial Regulation of Stilbene Synthase Genes by WRKY and MYB Transcription Factors in Grapevine (<i>Vitis vinifera</i> L.). Plant and Cell Physiology, 2018, 59, 1043-1059.	3.1	116
7	The leaf transcriptome of fennel (<i>Foeniculum vulgare</i> Mill.) enables characterization of the t-anethole pathway and the discovery of microsatellites and single-nucleotide variants. Scientific Reports, 2018, 8, 10459.	3.3	14
8	Current Advances in Genomics and Breeding of Leaf Chicory (<i>Cichorium intybus</i> L.). Agriculture (Switzerland), 2016, 6, 50.	3.1	27
9	Grapevine Rootstocks Differentially Affect the Rate of Ripening and Modulate Auxin-Related Genes in Cabernet Sauvignon Berries. Frontiers in Plant Science, 2016, 7, 69.	3.6	67
10	Comprehensive transcript profiling of two grapevine rootstock genotypes contrasting in drought susceptibility links the phenylpropanoid pathway to enhanced tolerance. Journal of Experimental Botany, 2015, 66, 5739-5752.	4.8	133
11	Genetic Segregation and Genomic Hybridization Patterns Support an Allotetraploid Structure and Disomic Inheritance for <i>Salix</i> Species. Diversity, 2014, 6, 633-651.	1.7	21
12	The coding region of the UFGT gene is a source of diagnostic SNP markers that allow single-locus DNA genotyping for the assessment of cultivar identity and ancestry in grapevine (<i>Vitis vinifera</i> L.). BMC Research Notes, 2013, 6, 502.	1.4	15
13	Alfalfa Mob1-like Genes are Expressed in Reproductive Organs during Meiosis and Gametogenesis. Plant Molecular Biology, 2005, 58, 789-807.	3.9	27
14	Immature embryos culture in Italian red chicory. Plant Cell, Tissue and Organ Culture, 2000, 62, 75-77.	2.3	10