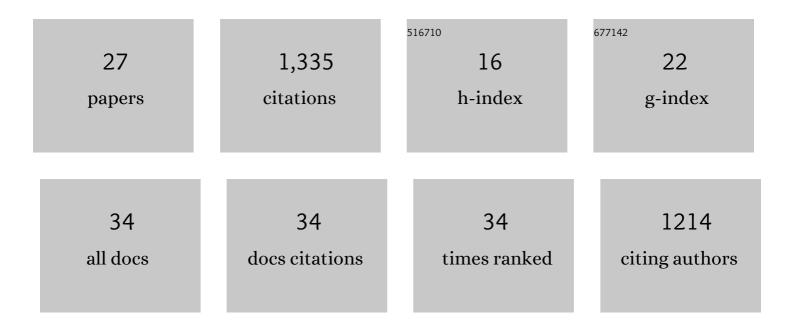
Eva Zyprian

List of Publications by Year in descending order

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Ενά Ζυρριανι

#	Article	IF	CITATIONS
1	High-Throughput Phenotyping of Leaf Discs Infected with Grapevine Downy Mildew Using Shallow Convolutional Neural Networks. Agronomy, 2021, 11, 1768.	3.0	5
2	Confirmation and Fine Mapping of the Resistance Locus Ren9 from the Grapevine Cultivar â€~Regent'. Plants, 2021, 10, 24.	3.5	22
3	Differential expression of transcription factor- and further growth-related genes correlates with contrasting cluster architecture in Vitis vinifera †Pinot Noir' and Vitis spp. genotypes. Theoretical and Applied Genetics, 2020, 133, 3249-3272.	3.6	6
4	Genetic Diversity, Population Structure, and Parentage Analysis of Croatian Grapevine Germplasm. Genes, 2020, 11, 737.	2.4	29
5	Colonization of Different Grapevine Tissues by Plasmopara viticola—A Histological Study. Frontiers in Plant Science, 2019, 10, 951.	3.6	30
6	Transcriptome analysis of early downy mildew (Plasmopara viticola) defense in grapevines carrying the Asian resistance locus Rpv10. Euphytica, 2019, 215, 1.	1.2	20
7	Identification of co-located QTLs and genomic regions affecting grapevine cluster architecture. Theoretical and Applied Genetics, 2019, 132, 1159-1177.	3.6	22
8	Fine mapping of Ren3 reveals two loci mediating hypersensitive response against Erysiphe necator in grapevine. Euphytica, 2017, 213, 1.	1.2	60
9	Genetic analysis of loose cluster architecture in grapevine. BIO Web of Conferences, 2017, 9, 01016.	0.2	3
10	Quantitative trait loci affecting pathogen resistance and ripening of grapevines. Molecular Genetics and Genomics, 2016, 291, 1573-1594.	2.1	124
11	Expression profiling of genes involved in drought stress and leaf senescence in juvenile barley. BMC Plant Biology, 2016, 16, 3.	3.6	57
12	QTL analysis of flowering time and ripening traits suggests an impact of a genomic region on linkage group 1 in Vitis. Theoretical and Applied Genetics, 2014, 127, 1857-1872.	3.6	44
13	Genetic diversity and population structure assessed by SSR and SNP markers in a large germplasm collection of grape. BMC Plant Biology, 2013, 13, 39.	3.6	325
14	Alterations in grapevine leaf metabolism upon inoculation with Plasmopara viticola in different time-points. Plant Science, 2012, 191-192, 100-107.	3.6	51
15	Cultivar-specific kinetics of gene induction during downy mildew early infection in grapevine. Functional and Integrative Genomics, 2012, 12, 379-386.	3.5	54
16	Rpv10: a new locus from the Asian Vitis gene pool for pyramiding downy mildew resistance loci in grapevine. Theoretical and Applied Genetics, 2012, 124, 163-176.	3.6	184
17	Marker Development for Important Grapevine Traits by Genetic Diversity Studies and Investigation of Differential Gene Expression. , 2010, , 375-387.		1
18	Identification, isolation and characterization of a CC-NBS-LRR candidate disease resistance gene family in grapevine. Molecular Breeding, 2008, 22, 421-432.	2.1	55

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#	Article	IF	CITATIONS
19	Development of SCAR markers linked to powdery mildew (Uncinula necator) resistance in grapevine (Vitis vinifera L. and Vitis sp.). Molecular Breeding, 2007, 19, 103-111.	2.1	78
20	Characterization ofPlasmopara-Resistance in grapevine usingin vitroplants. Journal of Plant Physiology, 2003, 160, 1393-1400.	3.5	107
21	COMPARATIVE MOLECULAR MAPPING IN SEGREGATING POPULATIONS OF GRAPEVINE. Acta Horticulturae, 2003, , 73-77.	0.2	6
22	GENOMICS TOOLS FOR MARKER ASSISTED SELECTION IN GRAPEVINE. Acta Horticulturae, 2003, , 511-517.	0.2	1
23	MOLECULAR MAPPING OF [REGENT X LEMBERGER] AND QTL-ANALYSIS OF AGRONOMIC TRAITS. Acta Horticulturae, 2003, , 69-71.	0.2	0
24	Plant Breeding: Genetic Mapping in Woody Crops. Progress in Botany Fortschritte Der Botanik, 1999, , 167-189.	0.3	1
25	Function of Genetic Material Molecular Biology of Environmentally Stressed Plants. , 1997, , 368-385.		0
26	Function: Histones in Higher Plants. , 1995, , 319-331.		0
27	Expression of a chloramphenicol-resistance determinant carried on hybrid plasmids in Gram-positive and Gram-negative bacteria. Gene, 1984, 32, 151-160.	2.2	48