

Gabriella De Lorenzis

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

1,392
citations

279798

23
h-index

361022

35
g-index

58
all docs

58
docs citations

58
times ranked

1416
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic diversity analysis of cultivated and wild grapevine (<i>Vitis vinifera</i> L.) accessions around the Mediterranean basin and Central Asia. <i>BMC Plant Biology</i> , 2018, 18, 137.	3.6	118
2	Grapevine field experiments reveal the contribution of genotype, the influence of environment and the effect of their interaction (G×E) on the berry transcriptome. <i>Plant Journal</i> , 2018, 93, 1143-1159.	5.7	75
3	Study of genetic variability in <i>Vitis vinifera</i> L. germplasm by high-throughput Vitis18kSNP array: the case of Georgian genetic resources. <i>BMC Plant Biology</i> , 2015, 15, 154.	3.6	68
4	Azole-resistant <i>Aspergillus fumigatus</i> in the environment of northern Italy, May 2011 to June 2012. <i>Eurosurveillance</i> , 2014, 19, 20747.	7.0	68
5	From the cradle of grapevine domestication: molecular overview and description of Georgian grapevine (<i>Vitis vinifera</i> L.) germplasm. <i>Tree Genetics and Genomes</i> , 2013, 9, 641-658.	1.6	65
6	The vintage effect overcomes the terroir effect: a three year survey on the wine yeast biodiversity in Franciacorta and Oltrepò Pavese, two northern Italian vine-growing areas. <i>Microbiology (United Kingdom)</i> , 2010, 156, 1010-1019.	1.0	50
7	SNP genotyping elucidates the genetic diversity of Magna Graecia grapevine germplasm and its historical origin and dissemination. <i>BMC Plant Biology</i> , 2019, 19, 7.	3.6	51
8	Unique resistance traits against downy mildew from the center of origin of grapevine (<i>Vitis vinifera</i>). <i>Scientific Reports</i> , 2018, 8, 12523.	3.3	50
9	Retrotransposon-based molecular markers for grapevine species and cultivars identification. <i>Tree Genetics and Genomes</i> , 2010, 6, 451-466.	1.6	49
10	Triazole resistance in <i>Aspergillus fumigatus</i> isolates from patients with cystic fibrosis in Italy. <i>Journal of Cystic Fibrosis</i> , 2017, 16, 64-69.	0.7	42
11	The Influence of Genotype and Environment on Small RNA Profiles in Grapevine Berry. <i>Frontiers in Plant Science</i> , 2016, 7, 1459.	3.6	40
12	Rpv29, Rpv30 and Rpv31: Three Novel Genomic Loci Associated With Resistance to <i>Plasmopara viticola</i> in <i>Vitis vinifera</i> . <i>Frontiers in Plant Science</i> , 2020, 11, 562432.	3.6	38
13	Intraspecific variations of <i>Dekkera/Brettanomyces bruxellensis</i> genome studied by capillary electrophoresis separation of the intron splice site profiles. <i>International Journal of Food Microbiology</i> , 2012, 157, 6-15.	4.7	37
14	Novel Aspects on The Interaction Between Grapevine and <i>Plasmopara viticola</i> : Dual-RNA-Seq Analysis Highlights Gene Expression Dynamics in The Pathogen and The Plant During The Battle For Infection. <i>Genes</i> , 2020, 11, 261.	2.4	37
15	High-throughput 18K SNP array to assess genetic variability of the main grapevine cultivars from Sicily. <i>Tree Genetics and Genomes</i> , 2016, 12, 1.	1.6	35
16	Anthocyanin biosynthesis during berry development in corvina grape. <i>Scientia Horticulturae</i> , 2016, 212, 74-80.	3.6	33
17	Azole Resistance in <i>Aspergillus fumigatus</i> Clinical Isolates from an Italian Culture Collection. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 682-685.	3.2	32
18	From plant resistance response to the discovery of antimicrobial compounds: The role of volatile organic compounds (VOCs) in grapevine downy mildew infection. <i>Plant Physiology and Biochemistry</i> , 2021, 160, 294-305.	5.8	32

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19	Italian wild grapevine (<i>Vitis vinifera</i> L. subsp. <i>sylvestris</i>) population: insights into eco-geographical aspects and genetic structure. <i>Tree Genetics and Genomes</i> , 2014, 10, 1369-1385.	1.6	31
20	Pedigree Reconstruction of the Italian Grapevine Aglianico (<i>Vitis vinifera</i> L.) from Campania. <i>Molecular Biotechnology</i> , 2013, 54, 634-642.	2.4	26
21	Georgian Grapevine Cultivars: Ancient Biodiversity for Future Viticulture. <i>Frontiers in Plant Science</i> , 2021, 12, 630122.	3.6	26
22	Novel and emerging biotechnological crop protection approaches. <i>Plant Biotechnology Journal</i> , 2021, 19, 1495-1510.	8.3	26
23	Genetic structure of Italian population of the grapevine downy mildew agent, <i>Plasmopara viticola</i> . <i>Annals of Applied Biology</i> , 2020, 176, 257-267.	2.5	25
24	RNAi of a Putative Grapevine Susceptibility Gene as a Possible Downy Mildew Control Strategy. <i>Frontiers in Plant Science</i> , 2021, 12, 667319.	3.6	25
25	Back to the Origins: Background and Perspectives of Grapevine Domestication. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4518.	4.1	24
26	Effectiveness of AFLPs and Retrotransposon-Based Markers for the Identification of Portuguese Grapevine Cultivars and Clones. <i>Molecular Biotechnology</i> , 2012, 52, 26-39.	2.4	23
27	Allelic variation in the <i>VvMYBA1</i> and <i>VvMYBA2</i> domestication genes in natural grapevine populations (<i>Vitis vinifera</i> subsp. <i>sylvestris</i>). <i>Plant Systematics and Evolution</i> , 2015, 301, 1613-1624.	0.9	21
28	Genotyping of Sicilian grapevine germplasm resources (<i>V. vinifera</i> L.) and their relationships with Sangiovese. <i>Scientia Horticulturae</i> , 2014, 169, 189-198.	3.6	20
29	Single nucleotide polymorphism profiles reveal an admixture genetic structure of grapevine germplasm from Calabria, Italy, uncovering its key role for the diversification of cultivars in the Mediterranean Basin. <i>Australian Journal of Grape and Wine Research</i> , 2018, 24, 345-359.	2.1	19
30	RETROTRANSPOSON-BASED MOLECULAR MARKERS IN GRAPEVINE SPECIES AND CULTIVARS IDENTIFICATION AND PHYLOGENETIC ANALYSIS. <i>Acta Horticulturae</i> , 2009, , 45-52.	0.2	19
31	Grapevine Non- <i>vinifera</i> Genetic Diversity Assessed by Simple Sequence Repeat Markers as a Starting Point for New Rootstock Breeding Programs. <i>American Journal of Enology and Viticulture</i> , 2019, 70, 390-397.	1.7	18
32	Genetic Diversity and Population Structure in a <i>Vitis</i> spp. Core Collection Investigated by SNP Markers. <i>Diversity</i> , 2020, 12, 103.	1.7	16
33	Study of intra-varietal diversity in biotypes of Aglianico and Muscat of Alexandria (<i>Vitis vinifera</i> L.) cultivars. <i>Australian Journal of Grape and Wine Research</i> , 2017, 23, 132-142.	2.1	15
34	Zibibbo Nero Characterization, a Red-Wine Grape Revertant of Muscat of Alexandria. <i>Molecular Biotechnology</i> , 2015, 57, 265-274.	2.4	14
35	How Do Novel M-Rootstock (<i>Vitis</i> Spp.) Genotypes Cope with Drought?. <i>Plants</i> , 2020, 9, 1385.	3.5	14
36	Climate Change Impacts on Plant Phenology: Grapevine (<i>Vitis vinifera</i>) Bud Break in Wintertime in Southern Italy. <i>Foods</i> , 2021, 10, 2769.	4.3	12

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37	Genetic investigation of grapevine varieties "Ribolla Gialla"™ (Italy), "Rebula"™ (Slovenia) and "Robola"™ (Ionian Islands). <i>Scientia Horticulturae</i> , 2013, 150, 425-431.	3.6	11
38	Iron deficiency stimulates anthocyanin accumulation in grapevine apical leaves. <i>Plant Physiology and Biochemistry</i> , 2017, 119, 286-293.	5.8	11
39	Genomic Designing for Biotic Stress Resistant Grapevine. , 2022, , 87-255.		11
40	Progress for research of grape and wine culture in Georgia, the South Caucasus. <i>BIO Web of Conferences</i> , 2019, 12, 03003.	0.2	10
41	Azole resistance in <i>Aspergillus</i> isolates by different types of patients and correlation with environment "An Italian prospective multicentre study (ARiA study). <i>Mycoses</i> , 2021, 64, 528-536.	4.0	9
42	Integrated Bayesian Approaches Shed Light on the Dissemination Routes of the Eurasian Grapevine Germplasm. <i>Frontiers in Plant Science</i> , 2021, 12, 692661.	3.6	9
43	Renewal of wild grapevine (<i>Vitis vinifera</i> L. subsp. <i>sylvestris</i> (Gmelin) Hegi) populations through sexual pathway: Some Italian case studies. <i>Flora: Morphology, Distribution, Functional Ecology of Plants</i> , 2016, 219, 85-93.	1.2	6
44	Pink berry grape (<i>Vitis vinifera</i> L.) characterization: Reflectance spectroscopy, HPLC and molecular markers. <i>Plant Physiology and Biochemistry</i> , 2016, 98, 138-145.	5.8	6
45	Prospective multicentre study on azole resistance in <i>Aspergillus</i> isolates from surveillance cultures in haematological patients in Italy. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 22, 231-237.	2.2	6
46	MULTIDISCIPLINARY STUDY OF TRADITIONAL GRAPE CULTIVARS FROM KARTLI PROVINCE OF GEORGIA (THE Tj ETQo 0 0 0 rgBT /Overlo	0.2	4
47	Disfunctions in the anthocyanin accumulation of <i>Vitis vinifera</i> L. varieties studied by a targeted resequencing approach. <i>Journal of Berry Research</i> , 2020, , 1-19.	1.4	3
48	'RIBOLLA GIALLA' FROM NORTH EASTERN ITALY, 'REBULA' FROM NORTHERN BALKANS AND 'ROBOLA' FROM IONIAN ISLANDS; DO THEY BELONG TO THE SAME POPULATION VARIETY OR ARE THEY GENETICALLY DIFFERENT?. <i>Acta Horticulturae</i> , 2014, , 645-652.	0.2	2
49	Yeast DNA recovery during the secondary fermentation step of Lombardy sparkling wines produced by Champenoise method. <i>European Food Research and Technology</i> , 2015, 240, 885-895.	3.3	2
50	Dissecting the susceptibility/resistance mechanism of <i>Vitis vinifera</i> for the future control of downy mildew. <i>BIO Web of Conferences</i> , 2022, 44, 04002.	0.2	2
51	MOLECULAR SURVEY OF GEORGIAN TRADITIONAL GRAPEVINE GENETIC RESOURCES. <i>Acta Horticulturae</i> , 2014, , 581-586.	0.2	1
52	Evidence for a Sympatric Origin of Ribolla gialla, Gouais Blanc and Schiava cultivars (<i>V. vinifera</i> L.). <i>South African Journal of Enology and Viticulture</i> , 2016, 35, .	0.4	1
53	PROTEOMIC ANALYSIS AMONG DIFFERENT AGLIANICO ECOTYPES. <i>Acta Horticulturae</i> , 2014, , 653-657.	0.2	0
54	ANALYSIS OF GENETIC STRUCTURE OF TWELVE SICILIAN GRAPEVINE CULTIVARS. <i>Acta Horticulturae</i> , 2014, , 677-680.	0.2	0

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55	ITALIAN WILD GRAPEVINE: A STATE OF THE ART ON GERMPLASM AND CONSERVATION IN 2010; THE YEAR OF BIODIVERSITY. Acta Horticulturae, 2014, , 639-644.	0.2	0
56	MOLECULAR INVESTIGATION OF CAUCASIAN AND EASTERN EUROPEAN GRAPEVINE GENETIC RESOURCES (V. Tj ETQq0 0 0 ggBT /Overl	0.2	0