

Jose Miguel Martinez Zapater

List of Publications by Year in descending order

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129
papers

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22153

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#	ARTICLE	IF	CITATIONS
1	Multiple origins of cultivated grapevine (<i>Vitis vinifera</i> L. ssp. <i>sativa</i>) based on chloroplast DNA polymorphisms. <i>Molecular Ecology</i> , 2006, 15, 3707-3714.	3.9	423
2	Low Temperature Induces the Accumulation of Phenylalanine Ammonia-Lyase and Chalcone Synthase mRNAs of <i>Arabidopsis thaliana</i> in a Light-Dependent Manner. <i>Plant Physiology</i> , 1995, 108, 39-46.	4.8	387
3	Constitutive expression of <i>Arabidopsis</i> LEAFY or APETALA1 genes in citrus reduces their generation time. <i>Nature Biotechnology</i> , 2001, 19, 263-267.	17.5	355
4	Regulation of flowering time by FVE, a retinoblastoma-associated protein. <i>Nature Genetics</i> , 2004, 36, 162-166.	21.4	347
5	Genetic diversity and population structure assessed by SSR and SNP markers in a large germplasm collection of grape. <i>BMC Plant Biology</i> , 2013, 13, 39.	3.6	325
6	Analysis of DNA methylation in <i>Arabidopsis thaliana</i> based on methylation-sensitive AFLP markers. <i>Molecular Genetics and Genomics</i> , 2002, 268, 543-552.	2.1	250
7	A highly repeated DNA sequence in <i>Arabidopsis thaliana</i> . <i>Molecular Genetics and Genomics</i> , 1986, 204, 417-423.	2.4	236
8	High throughput SNP discovery and genotyping in grapevine (<i>Vitis vinifera</i> L.) by combining a re-sequencing approach and SNPlex technology. <i>BMC Genomics</i> , 2007, 8, 424.	2.8	227
9	Isolation and molecular characterization of the <i>Arabidopsis</i> TPS1 gene, encoding trehalose 6-phosphate synthase. <i>Plant Journal</i> , 1998, 13, 685-689.	5.7	215
10	Genome-Wide Analysis of MIKCC-Type MADS Box Genes in Grapevine. <i>Plant Physiology</i> , 2009, 149, 354-369.	4.8	194
11	Different roles of flowering-time genes in the activation of floral initiation genes in <i>Arabidopsis</i> . <i>Plant Cell</i> , 1997, 9, 1921-1934.	6.6	181
12	Comparative analysis of grapevine whole-genome gene predictions, functional annotation, categorization and integration of the predicted gene sequences. <i>BMC Research Notes</i> , 2012, 5, 213.	1.4	176
13	Altitudinal and Climatic Adaptation Is Mediated by Flowering Traits and <i>FRI</i> , <i>FLC</i> , and <i>PHYC</i> Genes in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2011, 157, 1942-1955.	4.8	171
14	Mutations in the Ca ²⁺ /H ⁺ Transporter CAX1 Increase CBF/DREB1 Expression and the Cold-Acclimation Response in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2003, 15, 2940-2951.	6.6	170
15	Effect of Light Quality and Vernalization on Late-Flowering Mutants of <i>Arabidopsis thaliana</i> . <i>Plant Physiology</i> , 1990, 92, 770-776.	4.8	168
16	Sucrose availability on the aerial part of the plant promotes morphogenesis and flowering of <i>Arabidopsis</i> in the dark. <i>Plant Journal</i> , 1999, 20, 581-590.	5.7	168
17	The FT/TFL1 gene family in grapevine. <i>Plant Molecular Biology</i> , 2007, 63, 637-650.	3.9	167
18	Application of AFLPs to the characterization of grapevine <i>Vitis vinifera</i> L. genetic resources. A case study with accessions from Rioja (Spain). <i>Theoretical and Applied Genetics</i> , 1998, 97, 51-59.	3.6	164

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19	Extended diversity analysis of cultivated grapevine <i>Vitis vinifera</i> with 10K genome-wide SNPs. <i>PLoS ONE</i> , 2018, 13, e0192540.	2.5	164
20	Environmental regulation of flowering. <i>International Journal of Developmental Biology</i> , 2005, 49, 689-705.	0.6	149
21	Genetic and Molecular Analyses of Natural Variation Indicate CBF2 as a Candidate Gene for Underlying a Freezing Tolerance Quantitative Trait Locus in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2005, 139, 1304-1312.	4.8	149
22	Ancient asymmetries in the evolution of flowers. <i>Current Biology</i> , 2001, 11, 1050-1052.	3.9	147
23	Synergistic effect of methyljasmonate and cyclodextrin on stilbene biosynthesis pathway gene expression and resveratrol production in Monastrell grapevine cell cultures. <i>BMC Research Notes</i> , 2008, 1, 132.	1.4	147
24	Mutations at the <i>Arabidopsis</i> CHM locus promote rearrangements of the mitochondrial genome.. <i>Plant Cell</i> , 1992, 4, 889-899.	6.6	146
25	Molecular genetics of berry colour variation in table grape. <i>Molecular Genetics and Genomics</i> , 2006, 276, 427-435.	2.1	144
26	EARLY IN SHORT DAYS 1 (ESD1) encodes ACTIN-RELATED PROTEIN 6 (AtARP6), a putative component of chromatin remodelling complexes that positively regulates FLC accumulation in <i>Arabidopsis</i> . <i>Development (Cambridge)</i> , 2006, 133, 1241-1252.	2.5	144
27	Transcriptome and metabolome reprogramming in <i>Vitis vinifera</i> cv. Trincadeira berries upon infection with <i>Botrytis cinerea</i> . <i>Journal of Experimental Botany</i> , 2015, 66, 1769-1785.	4.8	144
28	A genetic analysis of seed and berry weight in grapevine. <i>Genome</i> , 2006, 49, 1572-1585.	2.0	139
29	A molecular genetic perspective of reproductive development in grapevine. <i>Journal of Experimental Botany</i> , 2008, 59, 2579-2596.	4.8	139
30	Transcript and metabolite analysis in Trincadeira cultivar reveals novel information regarding the dynamics of grape ripening. <i>BMC Plant Biology</i> , 2011, 11, 149.	3.6	133
31	Genetic diversity of wild grapevine populations in Spain and their genetic relationships with cultivated grapevines. <i>Molecular Ecology</i> , 2012, 21, 800-816.	3.9	130
32	VFL, the Grapevine FLORICAULA/LEAFY Ortholog, Is Expressed in Meristematic Regions Independently of Their Fate. <i>Plant Physiology</i> , 2002, 130, 68-77.	4.8	127
33	A 48 SNP set for grapevine cultivar identification. <i>BMC Plant Biology</i> , 2011, 11, 153.	3.6	127
34	Thermotolerance Responses in Ripening Berries of <i>Vitis vinifera</i> L. cv Muscat Hamburg. <i>Plant and Cell Physiology</i> , 2013, 54, 1200-1216.	3.1	123
35	A novel cold-inducible gene from <i>Arabidopsis</i> , RCI3, encodes a peroxidase that constitutes a component for stress tolerance. <i>Plant Journal</i> , 2002, 32, 13-24.	5.7	121
36	EARLY BOLTING IN SHORT DAYS Is Related to Chromatin Remodeling Factors and Regulates Flowering in <i>Arabidopsis</i> by Repressing FT. <i>Plant Cell</i> , 2003, 15, 1552-1562.	6.6	121

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37	Transcriptome changes in grapevine (<i>Vitis vinifera</i> L.) cv. Malbec leaves induced by ultraviolet-B radiation. <i>BMC Plant Biology</i> , 2010, 10, 224.	3.6	120
38	Floral Meristem Identity Genes Are Expressed during Tendril Development in Grapevine. <i>Plant Physiology</i> , 2004, 135, 1491-1501.	4.8	118
39	Chloroplast microsatellite polymorphisms in <i>Vitis</i> species. <i>Genome</i> , 2002, 45, 1142-1149.	2.0	117
40	Natural Genetic Variation of <i>Arabidopsis thaliana</i> Is Geographically Structured in the Iberian Peninsula. <i>Genetics</i> , 2008, 180, 1009-1021.	2.9	116
41	Transposon-induced gene activation as a mechanism generating cluster shape somatic variation in grapevine. <i>Plant Journal</i> , 2010, 61, 545-557.	5.7	116
42	Two related low-temperature-inducible genes of <i>Arabidopsis</i> encode proteins showing high homology to 14-3-3 proteins, a family of putative kinase regulators. <i>Plant Molecular Biology</i> , 1994, 25, 693-704.	3.9	110
43	Berry Flesh and Skin Ripening Features in <i>Vitis vinifera</i> as Assessed by Transcriptional Profiling. <i>PLoS ONE</i> , 2012, 7, e39547.	2.5	108
44	Low Temperature Induces the Accumulation of Alcohol Dehydrogenase mRNA in <i>Arabidopsis thaliana</i> , a Chilling-Tolerant Plant. <i>Plant Physiology</i> , 1993, 101, 833-837.	4.8	102
45	The Major Origin of Seedless Grapes Is Associated with a Missense Mutation in the MADS-Box Gene <i>VviAGL11</i> . <i>Plant Physiology</i> , 2018, 177, 1234-1253.	4.8	102
46	Structural and Functional Analysis of the GRAS Gene Family in Grapevine Indicates a Role of GRAS Proteins in the Control of Development and Stress Responses. <i>Frontiers in Plant Science</i> , 2016, 7, 353.	3.6	101
47	Two Homologous Low-Temperature-Inducible Genes from <i>Arabidopsis</i> Encode Highly Hydrophobic Proteins. <i>Plant Physiology</i> , 1997, 115, 569-576.	4.8	100
48	Catastrophic Unbalanced Genome Rearrangements Cause Somatic Loss of Berry Color in Grapevine. <i>Plant Physiology</i> , 2017, 175, 786-801.	4.8	98
49	DNA methylation increases throughout <i>Arabidopsis</i> development. <i>Planta</i> , 2005, 222, 301-306.	3.2	93
50	Chromatin remodeling in plant development. <i>International Journal of Developmental Biology</i> , 2009, 53, 1581-1596.	0.6	91
51	AP2 Gene Determines the Identity of Perianth Organs in Flowers of <i>Arabidopsis thaliana</i> . <i>Plant Cell</i> , 1989, 1, 1195.	6.6	84
52	Transcriptome variation along bud development in grapevine (<i>Vitis vinifera</i> L.). <i>BMC Plant Biology</i> , 2012, 12, 181.	3.6	83
53	Seed gene flow and fine-scale structure in a Mediterranean pine (<i>Pinus pinaster</i> Ait.) using nuclear microsatellite markers. <i>Theoretical and Applied Genetics</i> , 2002, 104, 1290-1297.	3.6	76
54	Genetic relationships among biotypes of <i>Bemisia tabaci</i> (Hemiptera: Aleyrodidae) based on AFLP analysis. <i>Bulletin of Entomological Research</i> , 2000, 90, 391-396.	1.0	75

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55	Arabidopsis late-flowering five mutants are affected in both vegetative and reproductive development. <i>Plant Journal</i> , 1995, 7, 543-551.	5.7	74
56	early bolting in short days: An Arabidopsis Mutation That Causes Early Flowering and Partially Suppresses the Floral Phenotype of leafy. <i>Plant Cell</i> , 2001, 13, 1011-1024.	6.6	71
57	Chromatin-Dependent Repression of the <i>Arabidopsis</i> Floral Integrator Genes Involves Plant Specific PHD-Containing Proteins. <i>Plant Cell</i> , 2014, 26, 3922-3938.	6.6	71
58	<i>EARLY IN SHORT DAYS 7</i> (<i>ESD7</i>) encodes the catalytic subunit of DNA polymerase epsilon and is required for flowering repression through a mechanism involving epigenetic gene silencing. <i>Plant Journal</i> , 2010, 61, 623-636.	5.7	68
59	Structural and functional annotation of the MADS-box transcription factor family in grapevine. <i>BMC Genomics</i> , 2016, 17, 80.	2.8	64
60	Developmental, transcriptome, and genetic alterations associated with parthenocarpy in the grapevine seedless somatic variant Corinto blanco. <i>Journal of Experimental Botany</i> , 2016, 67, 259-273.	4.8	61
61	The regulation of flowering in <i>Arabidopsis thaliana</i> : meristems, morphogenesis, and mutants. <i>Canadian Journal of Botany</i> , 1995, 73, 959-981.	1.1	60
62	A freezing-sensitive mutant of <i>Arabidopsis</i> , <i>frs1</i> , is a new <i>aba3</i> allele. <i>Planta</i> , 2000, 211, 648-655.	3.2	60
63	The Flowering Repressor SVP Underlies a Novel <i>Arabidopsis thaliana</i> QTL Interacting with the Genetic Background. <i>PLoS Genetics</i> , 2013, 9, e1003289.	3.5	58
64	A high-density collection of EMS-induced mutations for TILLING in Landsberg erecta genetic background of <i>Arabidopsis</i> . <i>BMC Plant Biology</i> , 2009, 9, 147.	3.6	57
65	Temporal analysis of natural variation for the rate of leaf production and its relationship with flowering initiation in <i>Arabidopsis thaliana</i> . <i>Journal of Experimental Botany</i> , 2010, 61, 1611-1623.	4.8	56
66	Dissecting the Transcriptional Response to Elicitors in <i>Vitis vinifera</i> Cells. <i>PLoS ONE</i> , 2014, 9, e109777.	2.5	56
67	Genetic relationship among cultivated and wild grapevine accessions from Tunisia. <i>Genome</i> , 2004, 47, 1211-1219.	2.0	55
68	The genetic structure of <i>Arabidopsis thaliana</i> in the south-western Mediterranean range reveals a shared history between North Africa and southern Europe. <i>BMC Plant Biology</i> , 2014, 14, 17.	3.6	53
69	Comparative genome-wide transcriptome analysis of <i>Vitis vinifera</i> responses to adapted and non-adapted strains of two-spotted spider mite, <i>Tetranychus urticae</i> . <i>BMC Genomics</i> , 2016, 17, 74.	2.8	53
70	Marker assisted selection for seedlessness in table grape breeding. <i>Tree Genetics and Genomes</i> , 2012, 8, 1003-1015.	1.6	51
71	The distribution of 5-methylcytosine in the nuclear genome of plants. <i>Nucleic Acids Research</i> , 1992, 20, 3207-3210.	14.5	49
72	Mutations causing defects in the biosynthesis and response to gibberellins, abscisic acid and phytochrome B do not inhibit vernalization in <i>Arabidopsis fca-1</i> . <i>Planta</i> , 2000, 210, 677-682.	3.2	49

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73	Grapevine genetics after the genome sequence: Challenges and limitations. Australian Journal of Grape and Wine Research, 2010, 16, 33-46.	2.1	49
74	Misexpression of a <i>PISTILLATA</i> -like <i>MADS</i> box gene prevents fruit development in grapevine. Plant Journal, 2013, 73, 918-928.	5.7	47
75	Challenges of viticulture adaptation to global change: tackling the issue from the roots. Australian Journal of Grape and Wine Research, 2021, 27, 8-25.	2.1	46
76	Generation of ESTs in <i>Vitis vinifera</i> wine grape (Cabernet Sauvignon) and table grape (Muscat) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 627 402, 40-50.	2.2	45
77	Novel natural alleles at <i>FLC</i> and <i>LVR</i> loci account for enhanced vernalization responses in <i>Arabidopsis thaliana</i> . Plant, Cell and Environment, 2012, 35, 1672-1684.	5.7	45
78	AFLP evaluation of genetic similarity among laurel populations (<i>Laurus</i> L.). Euphytica, 2001, 122, 155-164.	1.2	41
79	Genetic structure of natural populations of the grass endophyte <i>Epichloe festucae</i> in semiarid grasslands. Molecular Ecology, 2002, 11, 355-364.	3.9	41
80	Polymorphisms and minihaplotypes in the <i>VvNAC26</i> gene associate with berry size variation in grapevine. BMC Plant Biology, 2015, 15, 253.	3.6	41
81	Transcriptional Analysis of Tendril and Inflorescence Development in Grapevine (<i>Vitis vinifera</i> L.). PLoS ONE, 2014, 9, e92339.	2.5	40
82	Chloroplast genes transferred to the nuclear plant genome have adjusted to nuclear base composition and codon usage. Nucleic Acids Research, 1990, 18, 65-73.	14.5	39
83	Growing up fast: manipulating the generation time of trees. Current Opinion in Biotechnology, 2002, 13, 151-155.	6.6	38
84	Genetic Analysis of Variegated Mutants in Arabidopsis. Journal of Heredity, 1993, 84, 138-140.	2.4	37
85	Flowering transition in grapevine (<i>Vitis vinifera</i> L.) This review is one of a selection of papers presented at the symposium on <i>Vitis</i> at the XVII International Botanical Congress held in Vienna, Austria, 2005.. Canadian Journal of Botany, 2007, 85, 701-711.	1.1	37
86	Quantitative genetic analysis of flowering time in tomato. Genome, 2007, 50, 303-315.	2.0	36
87	Circadian oscillatory transcriptional programs in grapevine ripening fruits. BMC Plant Biology, 2014, 14, 78.	3.6	35
88	A genetic classification of potato cultivars based on allozyme patterns. Theoretical and Applied Genetics, 1985, 69, 305-311.	3.6	33
89	Characterisation of Iberian pig genotypes using AFLP markers. Animal Genetics, 2000, 31, 117-122.	1.7	33
90	Genetic Origin of the Grapevine Cultivar Tempranillo. American Journal of Enology and Viticulture, 2012, 63, 549-553.	1.7	33

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91	Quantitative genetic analysis of berry firmness in table grape (<i>Vitis vinifera</i> L.). <i>Tree Genetics and Genomes</i> , 2015, 11, 1.	1.6	33
92	Characterization of <i>Vitis vinifera</i> L. somatic variants exhibiting abnormal flower development patterns. <i>Journal of Experimental Botany</i> , 2007, 58, 4107-4118.	4.8	32
93	Composition and biological activity of the Algerian plant <i>Rosa canina</i> L. by HPLC-UV-MS. <i>Arabian Journal of Chemistry</i> , 2020, 13, 1105-1119.	4.9	32
94	Relationships among Gene Expression and Anthocyanin Composition of Malbec Grapevine Clones. <i>Journal of Agricultural and Food Chemistry</i> , 2014, 62, 6716-6725.	5.2	31
95	Low temperature regulates <i>Arabidopsis</i> Lhcb gene expression in a light-independent manner. <i>Plant Journal</i> , 1998, 13, 411-418.	5.7	30
96	Dimerization of <i>Arabidopsis</i> 14-3-3 proteins: structural requirements within the N-terminal domain and effect of calcium. <i>FEBS Letters</i> , 1999, 462, 377-382.	2.8	29
97	Regulation of the <i>rab17</i> gene promoter in transgenic <i>Arabidopsis</i> wild-type, ABA-deficient and ABA-insensitive mutants. <i>Plant Molecular Biology</i> , 1994, 24, 561-569.	3.9	28
98	Combining microsatellite markers and capillary gel electrophoresis with laser-induced fluorescence to identify the grape (<i>Vitis vinifera</i>) variety of musts. <i>European Food Research and Technology</i> , 2006, 223, 625-631.	3.3	27
99	Non-random distribution of transposable elements in the nuclear genome of plants. <i>Nucleic Acids Research</i> , 1993, 21, 2369-2373.	14.5	26
100	Haplotype diversity of <i>VvTFL1A</i> gene and association with cluster traits in grapevine (<i>V. vinifera</i>). <i>BMC Plant Biology</i> , 2014, 14, 209.	3.6	26
101	Population genetic analysis in old Montenegrin vineyards reveals ancient ways currently active to generate diversity in <i>Vitis vinifera</i> . <i>Scientific Reports</i> , 2020, 10, 15000.	3.3	22
102	Expression of <i>Arabidopsis</i> <i>APETALA1</i> in tomato reduces its vegetative cycle without affecting plant production. <i>Molecular Breeding</i> , 2004, 13, 155-163.	2.1	21
103	B-Chromosomes and E-1 isozyme activity in mosaic bulbs of <i>Scilla autumnalis</i> (Liliaceae). <i>Chromosoma</i> , 1982, 85, 399-403.	2.2	20
104	Selfing and sibship structure in a two-cohort stand of maritime pine (<i>Pinus pinaster</i> Ait.) using nuclear SSR markers. <i>Annals of Forest Science</i> , 2003, 60, 115-121.	2.0	17
105	Grapevine breeding and clonal selection programmes in Spain. , 2015, , 183-209.		17
106	Identification of genes specifically expressed in cauliflower reproductive meristems. Molecular characterization of <i>BoREM1</i> . <i>Plant Molecular Biology</i> , 1999, 39, 427-436.	3.9	16
107	Identification by SNP Analysis of a Major Role for Cayetana Blanca in the Genetic Network of Iberian Peninsula Grapevine Varieties. <i>American Journal of Enology and Viticulture</i> , 2012, 63, 121-126.	1.7	16
108	Genetic diversity and parentage of Tunisian wild and cultivated grapevines (<i>Vitis vinifera</i> L.) as revealed by single nucleotide polymorphism (SNP) markers. <i>Tree Genetics and Genomes</i> , 2014, 10, 1103-1112.	1.6	16

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109	Reducing sampling bias in molecular studies of grapevine fruit ripening: transcriptomic assessment of the density sorting method. <i>Theoretical and Experimental Plant Physiology</i> , 2016, 28, 109-129.	2.4	16
110	Isozyme gene duplication in diploid and tetraploid potatoes. <i>Theoretical and Applied Genetics</i> , 1985, 70, 172-177.	3.6	14
111	Expression of grapevine AINTEGUMENTA-like genes is associated with variation in ovary and berry size. <i>Plant Molecular Biology</i> , 2016, 91, 67-80.	3.9	13
112	Somatic Variation and Cultivar Innovation in Grapevine. , 2019, , .		13
113	Whole genome resequencing and custom genotyping unveil clonal lineages in "Malbec"™ grapevines (<i>Vitis vinifera</i> L.). <i>Scientific Reports</i> , 2021, 11, 7775.	3.3	12
114	Drosophila P-element transcripts are incorrectly processed in tobacco. <i>Plant Molecular Biology</i> , 1988, 11, 601-607.	3.9	11
115	Characterization of Tempranillo negro (VN21), a high phenolic content grapevine Tempranillo clone, through UHPLC-QqQ-MS/MS polyphenol profiling. <i>Food Chemistry</i> , 2021, 360, 130049.	8.2	10
116	Comparative ampelographic and genetic analysis of grapevine cultivars from Algeria and Morocco. <i>Australian Journal of Grape and Wine Research</i> , 2014, 20, 324-333.	2.1	9
117	Grapevine Diversity and Genetic Relationships in Northeast Portugal Old Vineyards. <i>Plants</i> , 2021, 10, 2755.	3.5	9
118	A Brassica oleracea Gene Expressed in a Variety-Specific Manner May Encode a Novel Plant Transmembrane Receptor. <i>Plant and Cell Physiology</i> , 2001, 42, 404-413.	3.1	8
119	SSR and SNP genetic profiling of Armenian grape cultivars gives insights into their identity and pedigree relationships. <i>Oeno One</i> , 2021, 55, 101-114.	1.4	8
120	Genetic Identification and Origin of Grapevine Cultivars (<i>Vitis vinifera</i> L.) in Tunisia. <i>American Journal of Enology and Viticulture</i> , 2013, 64, 538-544.	1.7	5
121	Genetic variation and association analyses identify genes linked to fruit set-related traits in grapevine. <i>Plant Science</i> , 2021, 306, 110875.	3.6	5
122	Evaluation of Microsatellite Detection Using Autoradiography and Capillary Electrophoresis in Hops. <i>Journal of the American Society of Brewing Chemists</i> , 2005, 63, 57-62.	1.1	3
123	Molecular Maps, Qtl Mapping & Association Mapping In Grapevine. , 2009, , 535-563.		3
124	Genetic variation for grapevine reproductive development. <i>Acta Horticulturae</i> , 2019, , 319-326.	0.2	3
125	Grape color variation involves genetic and micro-environmental changes that alter berry phenolic and aromatic composition. <i>Acta Horticulturae</i> , 2019, , 471-478.	0.2	2
126	Reduced gamete viability associated to somatic genome rearrangements increases fruit set sensitivity to the environment in Tempranillo Blanco grapevine cultivar. <i>Scientia Horticulturae</i> , 2021, 290, 110497.	3.6	2

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127	Characterization of deletions causing berry-color variation in Garnacha and Tempranillo. Acta Horticulturae, 2019, , 463-470.	0.2	1
128	Is aromatic terpenoid composition of grapes in Northwestern Iberian wine cultivars related to variation in VviDXS1 gene?. Journal of Berry Research, 2021, 11, 187-200.	1.4	1
129	NATO-ASI course on plant molecular biology. Plant Molecular Biology Reporter, 1993, 11, 350-358.	1.8	0