

Amalia Barone

List of Publications by Year in descending order

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76
papers

5,688
citations

147726

31
h-index

79644

73
g-index

77
all docs

77
docs citations

77
times ranked

7414
citing authors

#	ARTICLE	IF	CITATIONS
1	The tomato genome sequence provides insights into fleshy fruit evolution. <i>Nature</i> , 2012, 485, 635-641.	13.7	2,860
2	Enhancing the Health-Promoting Effects of Tomato Fruit for Biofortified Food. <i>Mediators of Inflammation</i> , 2014, 2014, 1-16.	1.4	189
3	RFLP mapping on potato chromosomes of two genes controlling extreme resistance to potato virus X (PVX). <i>Molecular Genetics and Genomics</i> , 1991, 227, 81-85.	2.4	167
4	Localization by restriction fragment length polymorphism mapping in potato of a major dominant gene conferring resistance to the potato cyst nematode <i>Globodera rostocbiensis</i> . <i>Molecular Genetics and Genomics</i> , 1990, 224, 177-182.	2.4	150
5	The ascorbic acid content of tomato fruits is associated with the expression of genes involved in pectin degradation. <i>BMC Plant Biology</i> , 2010, 10, 163.	1.6	103
6	Exploring a Tomato Landraces Collection for Fruit-Related Traits by the Aid of a High-Throughput Genomic Platform. <i>PLoS ONE</i> , 2015, 10, e0137139.	1.1	91
7	Bioactive Compounds in Brassicaceae Vegetables with a Role in the Prevention of Chronic Diseases. <i>Molecules</i> , 2018, 23, 15.	1.7	86
8	Endosperm balance number manipulation for direct in vivo germplasm introgression to potato from a sexually isolated relative (<i>Solanum commersonii</i> Dun.). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1997, 94, 12013-12017.	3.3	85
9	An association mapping approach to identify favourable alleles for tomato fruit quality breeding. <i>BMC Plant Biology</i> , 2014, 14, 337.	1.6	84
10	The Use of a Plant-Based Biostimulant Improves Plant Performances and Fruit Quality in Tomato Plants Grown at Elevated Temperatures. <i>Agronomy</i> , 2020, 10, 363.	1.3	75
11	A Snapshot of the Emerging Tomato Genome Sequence. <i>Plant Genome</i> , 2009, 2, .	1.6	73
12	Molecular marker-assisted selection for potato breeding. <i>American Journal of Potato Research</i> , 2004, 81, 111-117.	0.5	71
13	Ploidy level manipulations in potato through sexual hybridisation. <i>Annals of Applied Biology</i> , 2005, 146, 71-79.	1.3	68
14	Antioxidant bioactive compounds in tomato fruits at different ripening stages and their effects on normal and cancer cells. <i>Journal of Functional Foods</i> , 2015, 18, 83-94.	1.6	67
15	Metabolite Profiling of Italian Tomato Landraces with Different Fruit Types. <i>Frontiers in Plant Science</i> , 2016, 7, 664.	1.7	65
16	Quantitative Trait Loci Pyramiding Can Improve the Nutritional Potential of Tomato (<i>Solanum</i>) Tj ETQq0 0 0 rgBT JOverlock_10 Tf 50	2.4	57
17	Vitamin E Content and Composition in Tomato Fruits: Beneficial Roles and Bio-Fortification. <i>International Journal of Molecular Sciences</i> , 2015, 16, 29250-29264.	1.8	54
18	Heterozygosity in 2n gametes of potato evaluated by RFLP markers. <i>Theoretical and Applied Genetics</i> , 1995, 91, 98-104.	1.8	51

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19	Quantitative trait loci pyramiding for fruit quality traits in tomato. <i>Molecular Breeding</i> , 2013, 31, 217-222.	1.0	51
20	Identification of candidate genes for phenolics accumulation in tomato fruit. <i>Plant Science</i> , 2013, 205-206, 87-96.	1.7	51
21	Unraveling the complexity of transcriptomic, metabolomic and quality environmental response of tomato fruit. <i>BMC Plant Biology</i> , 2017, 17, 66.	1.6	48
22	Structural and Functional Genomics of Tomato. <i>International Journal of Plant Genomics</i> , 2008, 2008, 1-12.	2.2	46
23	An ascorbic acid-enriched tomato genotype to fight UVA-induced oxidative stress in normal human keratinocytes. <i>Journal of Photochemistry and Photobiology B: Biology</i> , 2016, 163, 284-289.	1.7	46
24	Pectic enzymes as potential enhancers of ascorbic acid production through the D-galacturonate pathway in Solanaceae. <i>Plant Science</i> , 2018, 266, 55-63.	1.7	46
25	Exploiting Genetic and Genomic Resources to Enhance Heat-Tolerance in Tomatoes. <i>Agronomy</i> , 2019, 9, 22.	1.3	45
26	Carotenoids in fresh and processed tomato (<i>Solanum lycopersicum</i>) fruits protect cells from oxidative stress injury. <i>Journal of the Science of Food and Agriculture</i> , 2017, 97, 1616-1623.	1.7	42
27	High-Throughput Genomics Enhances Tomato Breeding Efficiency. <i>Current Genomics</i> , 2009, 10, 1-9.	0.7	40
28	Selection for aneuploid potato hybrids combining a low wild genome content and resistance traits from <i>Solanum commersonii</i> . <i>Theoretical and Applied Genetics</i> , 2004, 109, 1139-1146.	1.8	37
29	A Novel Protein Hydrolysate-Based Biostimulant Improves Tomato Performances under Drought Stress. <i>Plants</i> , 2021, 10, 783.	1.6	37
30	Selection of tomato landraces with high fruit yield and nutritional quality under elevated temperatures. <i>Journal of the Science of Food and Agriculture</i> , 2020, 100, 2791-2799.	1.7	35
31	Resistance to <i>Ralstonia solanacearum</i> of Sexual Hybrids Between <i>Solanum commersonii</i> and <i>S. tuberosum</i> . <i>American Journal of Potato Research</i> , 2009, 86, 196-202.	0.5	34
32	Molecular marker-assisted introgression of the wild <i>Solanum commersonii</i> genome into the cultivated <i>S. tuberosum</i> gene pool. <i>Theoretical and Applied Genetics</i> , 2001, 102, 900-907.	1.8	33
33	Dissection of genetic and environmental factors involved in tomato organoleptic quality. <i>BMC Plant Biology</i> , 2011, 11, 58.	1.6	33
34	Patchwork sequencing of tomato San Marzano and Vesuviano varieties highlights genome-wide variations. <i>BMC Genomics</i> , 2014, 15, 138.	1.2	32
35	Biochemical, sensorial and genomic profiling of traditional Italian tomato varieties. <i>Euphytica</i> , 2008, 164, 571-582.	0.6	31
36	Production of Pharmaceutical Proteins in Solanaceae Food Crops. <i>International Journal of Molecular Sciences</i> , 2013, 14, 2753-2773.	1.8	31

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37	Eco-physiological response to water stress of drought-tolerant and drought-sensitive tomato genotypes. <i>Plant Biosystems</i> , 2016, 150, 682-691.	0.8	30
38	Accelerating Tomato Breeding by Exploiting Genomic Selection Approaches. <i>Plants</i> , 2020, 9, 1236.	1.6	30
39	Exploiting Genomics Resources to Identify Candidate Genes Underlying Antioxidants Content in Tomato Fruit. <i>Frontiers in Plant Science</i> , 2016, 7, 397.	1.7	29
40	Evaluation and use of plant biodiversity for food and pharmaceuticals. <i>FÄ-toterapÄ-Äç</i> , 2000, 71, S66-S72.	1.1	28
41	Integrated bioinformatics to decipher the ascorbic acid metabolic network in tomato. <i>Plant Molecular Biology</i> , 2016, 91, 397-412.	2.0	26
42	A novel synthetic peptide from a tomato defensin exhibits antibacterial activities against <i>Helicobacter pylori</i> . <i>Journal of Peptide Science</i> , 2012, 18, 755-762.	0.8	24
43	Metabolic and Molecular Changes of the Phenylpropanoid Pathway in Tomato (<i>Solanum lycopersicum</i>) Lines Carrying Different <i>Solanum pennellii</i> Wild Chromosomal Regions. <i>Frontiers in Plant Science</i> , 2016, 7, 1484.	1.7	23
44	Transcriptional Regulation of Ascorbic Acid During Fruit Ripening in Pepper (<i>Capsicum annum</i>) Varieties with Low and High Antioxidants Content. <i>Plants</i> , 2019, 8, 206.	1.6	23
45	New insights in the control of antioxidants accumulation in tomato by transcriptomic analyses of genotypes exhibiting contrasting levels of fruit metabolites. <i>BMC Genomics</i> , 2019, 20, 43.	1.2	23
46	Eco-Physiological Screening of Different Tomato Genotypes in Response to High Temperatures: A Combined Field-to-Laboratory Approach. <i>Plants</i> , 2020, 9, 508.	1.6	23
47	Cytological evidences of SDR-FDR mixture in the formation of 2n eggs in a potato diploid clone. <i>Theoretical and Applied Genetics</i> , 1991, 81, 59-63.	1.8	21
48	Identification of single nucleotide polymorphisms in Toll-like receptor candidate genes associated with tuberculosis infection in water buffalo (<i>Bubalus bubalis</i>). <i>BMC Genetics</i> , 2014, 15, 139.	2.7	21
49	Phenotyping to dissect the biostimulant action of a protein hydrolysate in tomato plants under combined abiotic stress. <i>Plant Physiology and Biochemistry</i> , 2022, 179, 32-43.	2.8	20
50	Bioactive Compound Content and Cytotoxic Effect on Human Cancer Cells of Fresh and Processed Yellow Tomatoes. <i>Molecules</i> , 2016, 21, 33.	1.7	18
51	Identification of tomato accessions as source of new genes for improving heat tolerance: from controlled experiments to field. <i>BMC Plant Biology</i> , 2021, 21, 345.	1.6	18
52	Evidence for tetrasomic inheritance in a tetraploid <i>Solanum commersonii</i> (+) <i>S. tuberosum</i> somatic hybrid through the use of molecular markers. <i>Theoretical and Applied Genetics</i> , 2002, 104, 539-546.	1.8	17
53	Glycoalkaloids and acclimation capacity of hybrids between <i>Solanum tuberosum</i> and the incongruent hardy species <i>Solanum commersonii</i> . <i>Theoretical and Applied Genetics</i> , 2003, 107, 1187-1194.	1.8	16
54	High-Throughput Genotyping of Resilient Tomato Landraces to Detect Candidate Genes Involved in the Response to High Temperatures. <i>Genes</i> , 2020, 11, 626.	1.0	16

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55	Dissecting a QTL into Candidate Genes Highlighted the Key Role of Pectinesterases in Regulating the Ascorbic Acid Content in Tomato Fruit. <i>Plant Genome</i> , 2015, 8, eplantgenome2014.08.0038.	1.6	16
56	Tuber quality and soft rot resistance of hybrids between <i>Solanum tuberosum</i> and the incongruent wild relative <i>S. commersonii</i> . <i>American Journal of Potato Research</i> , 2002, 79, 345-352.	0.5	14
57	Phenotypic and Molecular Selection of a Superior <i>Solanum pennellii</i> Introgression Sub-Line Suitable for Improving Quality Traits of Cultivated Tomatoes. <i>Frontiers in Plant Science</i> , 2019, 10, 190.	1.7	14
58	Tomato genomic prediction for good performance under high-temperature and identification of loci involved in thermotolerance response. <i>Horticulture Research</i> , 2021, 8, 212.	2.9	14
59	A basic Helix-Loop-Helix (SIARANCIO), identified from a <i>Solanum pennellii</i> introgression line, affects carotenoid accumulation in tomato fruits. <i>Scientific Reports</i> , 2019, 9, 3699.	1.6	13
60	Genetic Diversity within Wild Potato Species (<i>Solanum</i> spp.) Revealed by AFLP and SCAR Markers. <i>American Journal of Plant Sciences</i> , 2010, 01, 95-103.	0.3	12
61	Host and Non-Host Plant Response to Bacterial Wilt in Potato: Role of the Lipopolysaccharide Isolated from <i>Ralstonia solanacearum</i> and Molecular Analysis of Plant-Pathogen Interaction. <i>Chemistry and Biodiversity</i> , 2008, 5, 2662-2675.	1.0	11
62	A comparative study of the physico-chemical properties affecting the organoleptic quality of fresh and thermally treated yellow tomato ecotype fruit. <i>International Journal of Food Science and Technology</i> , 2018, 53, 1219-1226.	1.3	11
63	Use of seedling tubers from TPS in southern Italy. <i>American Potato Journal</i> , 1994, 71, 29-38.	0.4	8
64	Chromosome pairing in <i>Solanum commersonii</i> - <i>S. tuberosum</i> sexual hybrids detected by <i>commersonii</i> -specific RAPDs and cytological analysis. <i>Genome</i> , 1999, 42, 218-224.	0.9	8
65	Exploiting the great potential of Sequence Capture data by a new tool, SUPER-CAP. <i>DNA Research</i> , 2016, 24, dsw050.	1.5	8
66	Comparative Transcriptomic Profiling of Two Tomato Lines with Different Ascorbate Content in the Fruit. <i>Biochemical Genetics</i> , 2012, 50, 908-921.	0.8	7
67	Positive selection in the leucine-rich repeat domain of Gro1 genes in <i>Solanum</i> species. <i>Journal of Genetics</i> , 2014, 93, 755-765.	0.4	7
68	Genomic Dissection of a Wild Region in a Superior <i>Solanum pennellii</i> Introgression Sub-Line with High Ascorbic Acid Accumulation in Tomato Fruit. <i>Genes</i> , 2020, 11, 847.	1.0	7
69	Chromosome pairing in <i>Solanum commersonii</i> - <i>S. tuberosum</i> sexual hybrids detected by <i>commersonii</i> -specific RAPDs and cytological analysis. <i>Genome</i> , 1999, 42, 218-224.	0.9	6
70	Impact of Wild Loci on the Allergenic Potential of Cultivated Tomato Fruits. <i>PLoS ONE</i> , 2016, 11, e0155803.	1.1	4
71	Accelerating the Development of Heat Tolerant Tomato Hybrids through a Multi-Traits Evaluation of Parental Lines Combining Phenotypic and Genotypic Analysis. <i>Plants</i> , 2021, 10, 2168.	1.6	3
72	A Novel Plant-Based Biostimulant Improves Plant Performances under Drought Stress in Tomato. <i>Biology and Life Sciences Forum</i> , 2021, 4, 52.	0.6	2

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73	Genome Analysis of Species of Agricultural Interest. Springer Optimization and Its Applications, 2009, , 385-402.	0.6	1
74	Knowledge on the Genomes of Wild Tomato Species is the Key to Unlocking Their Breeding Potential. Compendium of Plant Genomes, 2021, , 155-166.	0.3	0
75	Higher Yield and Fruit Quality of a Solanum pennellii Introgression Line. Biology and Life Sciences Forum, 2021, 3, 31.	0.6	0
76	One Plant-Based Biostimulant Stimulates Good Performances of Tomato Plants Grown in Open Field. Biology and Life Sciences Forum, 2021, 3, .	0.6	0