

Enrico Francia

List of Publications by Year in descending order

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67
papers

4,800
citations

159585

30
h-index

106344

65
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70
all docs

70
docs citations

70
times ranked

5443
citing authors

#	ARTICLE	IF	CITATIONS
1	Plant Biostimulants in Sustainable Potato Production: an Overview. <i>Potato Research</i> , 2022, 65, 83-104.	2.7	17
2	Agronomic Comparisons of Heirloom and Modern Processing Tomato Genotypes Cultivated in Organic and Conventional Farming Systems. <i>Agronomy</i> , 2021, 11, 349.	3.0	4
3	Genetic and Management Effects on Barley Yield and Phenology in the Mediterranean Basin. <i>Frontiers in Plant Science</i> , 2021, 12, 655406.	3.6	12
4	Effects of Biostimulants on the Chemical Composition of Essential Oil and Hydrosol of Lavandin (<i>Lavandula x intermedia</i> Emeric ex Loisel.) Cultivated in Tuscan-Emilian Apennines. <i>Molecules</i> , 2021, 26, 6157.	3.8	10
5	Extensive allele mining discovers novel genetic diversity in the loci controlling frost tolerance in barley. <i>Theoretical and Applied Genetics</i> , 2021, , 1.	3.6	9
6	Influence of CNV on transcript levels of HvCBF genes at Fr-H2 locus revealed by resequencing in resistant barley cv. "Nure"™ and expression analysis. <i>Plant Science</i> , 2020, 290, 110305.	3.6	5
7	In Silico Identification of MYB and bHLH Families Reveals Candidate Transcription Factors for Secondary Metabolic Pathways in <i>Cannabis sativa</i> L.. <i>Plants</i> , 2020, 9, 1540.	3.5	14
8	Characterization of Celiac Disease-Related Epitopes and Gluten Fractions, and Identification of Associated Loci in Durum Wheat. <i>Agronomy</i> , 2020, 10, 1231.	3.0	6
9	Interspecific rootstock can enhance yield of processing tomatoes (<i>Solanum lycopersicum</i> L.) in organic farming. <i>Biological Agriculture and Horticulture</i> , 2020, 36, 156-171.	1.0	2
10	Influence of environmental and genetic factors on content of toxic and immunogenic wheat gluten peptides. <i>European Journal of Agronomy</i> , 2020, 118, 126091.	4.1	10
11	Bioplastic Film from Black Soldier Fly Prepupae Proteins Used as Mulch: Preliminary Results. <i>Agronomy</i> , 2020, 10, 933.	3.0	12
12	Using Digestate and Biochar as Fertilizers to Improve Processing Tomato Production Sustainability. <i>Agronomy</i> , 2020, 10, 138.	3.0	53
13	Changes in yield components, morphological, physiological and fruit quality traits in processing tomato cultivated in Italy since the 1930s. <i>Scientia Horticulturae</i> , 2019, 257, 108726.	3.6	32
14	Nitrogen Fertilizers Shape the Composition and Predicted Functions of the Microbiota of Field-Grown Tomato Plants. <i>Phytobiomes Journal</i> , 2019, 3, 315-325.	2.7	26
15	Valorization of Vineyard By-Products to Obtain Composted Digestate and Biochar Suitable for Nursery Grapevine (<i>Vitis vinifera</i> L.) Production. <i>Agronomy</i> , 2019, 9, 420.	3.0	27
16	Blossom end-rot in tomato (<i>Solanum lycopersicum</i> L.): A multi-disciplinary overview of inducing factors and control strategies. <i>Scientia Horticulturae</i> , 2019, 249, 49-58.	3.6	65
17	Use of black soldier fly (<i>Hermetia illucens</i> (L.), Diptera: Stratiomyidae) larvae processing residue in peat-based growing media. <i>Waste Management</i> , 2019, 95, 278-288.	7.4	88
18	Arbuscular Mycorrhizal Fungi and Plant Growth Promoting Rhizobacteria Avoid Processing Tomato Leaf Damage during Chilling Stress. <i>Agronomy</i> , 2019, 9, 299.	3.0	32

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19	The impact of climate change on barley yield in the Mediterranean basin. <i>European Journal of Agronomy</i> , 2019, 106, 1-11.	4.1	93
20	Carbon footprint and energetic analysis of tomato production in the organic vs the conventional cropping systems in Southern Italy. <i>Journal of Cleaner Production</i> , 2019, 220, 836-845.	9.3	49
21	Technological Quality and Nutritional Value of Two Durum Wheat Varieties Depend on Both Genetic and Environmental Factors. <i>Journal of Agricultural and Food Chemistry</i> , 2019, 67, 2384-2395.	5.2	29
22	Interaction of Tomato Genotypes and Arbuscular Mycorrhizal Fungi under Reduced Irrigation. <i>Horticulturae</i> , 2019, 5, 79.	2.8	13
23	A Meta-Analysis of Comparative Transcriptomic Data Reveals a Set of Key Genes Involved in the Tolerance to Abiotic Stresses in Rice. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5662.	4.1	24
24	Combined Effect of Cadmium and Lead on Durum Wheat. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5891.	4.1	21
25	Effects of solid and liquid digestate for hydroponic baby leaf lettuce (<i>Lactuca sativa</i> L.) cultivation. <i>Scientia Horticulturae</i> , 2019, 244, 172-181.	3.6	66
26	Physiological responses to chilling in cultivars of processing tomato released and cultivated over the past decades in Southern Europe. <i>Scientia Horticulturae</i> , 2018, 231, 118-125.	3.6	26
27	<i>Panicum</i> spikelets from the Early Holocene Takarkori rockshelter (SW Libya): Archaeo-molecular and -botanical investigations. <i>Plant Biosystems</i> , 2018, 152, 1-13.	1.6	13
28	Testing the influence of digestate from biogas on growth and volatile compounds of basil (<i>Ocimum</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 Medicinal and Aromatic Plants, 2018, 11, 18-26.	1.5	20
29	Tracking celiac disease-triggering peptides and whole wheat flour quality as function of germination kinetics. <i>Food Research International</i> , 2018, 112, 345-352.	6.2	6
30	Transcriptome profiling of short-term response to chilling stress in tolerant and sensitive <i>Oryza sativa</i> ssp. Japonica seedlings. <i>Functional and Integrative Genomics</i> , 2018, 18, 627-644.	3.5	34
31	Biomass production and dry matter partitioning of processing tomato under organic vs conventional cropping systems in a Mediterranean environment. <i>Scientia Horticulturae</i> , 2017, 224, 163-170.	3.6	52
32	Marker characterization of vernalization and low-temperature tolerance loci in barley genotypes adapted to semi-arid environments. <i>Czech Journal of Genetics and Plant Breeding</i> , 2016, 52, 157-162.	0.8	4
33	Agronomic and molecular evaluation of cocksfoot and tall fescue cultivars for adaptation to an Algerian drought-prone environment. <i>Euphytica</i> , 2016, 212, 371-386.	1.2	8
34	Evaluation of <i>Cucurbita pepo</i> germplasm for staminate flower production and adaptation to the frozen food industry. <i>Scientia Horticulturae</i> , 2016, 213, 321-330.	3.6	3
35	Copy number variation at the HvCBF4-HvCBF2 genomic segment is a major component of frost resistance in barley. <i>Plant Molecular Biology</i> , 2016, 92, 161-175.	3.9	45
36	Physiological responses of processing tomato in organic and conventional Mediterranean cropping systems. <i>Scientia Horticulturae</i> , 2015, 190, 161-172.	3.6	39

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37	CNV and Structural Variation in Plants: Prospects of NGS Approaches. , 2015, , 211-232.		8
38	The barley Frost resistance-H2 locus. Functional and Integrative Genomics, 2014, 14, 85-100.	3.5	19
39	Candidate gene expression profiling in two contrasting tomato cultivars under chilling stress. <i>Biologia Plantarum</i> , 2014, 58, 283-295.	1.9	26
40	QTLs for barley yield adaptation to Mediterranean environments in the "Nure"–"Tremois" biparental population. <i>Euphytica</i> , 2014, 197, 73-86.	1.2	74
41	Genome-wide association mapping of frost tolerance in barley (<i>Hordeum vulgare</i> L.). <i>BMC Genomics</i> , 2013, 14, 424.	2.8	101
42	Determinants of barley grain yield in drought-prone Mediterranean environments. <i>Italian Journal of Agronomy</i> , 2013, 8, 1.	1.0	17
43	QTLs for resistance to the false brome rust <i>Puccinia brachypodii</i> in the model grass <i>Brachypodium distachyon</i> L.. <i>Genome</i> , 2012, 55, 152-163.	2.0	28
44	Natural variation in a homolog of <i>Antirrhinum CENTRORADIALIS</i> contributed to spring growth habit and environmental adaptation in cultivated barley. <i>Nature Genetics</i> , 2012, 44, 1388-1392.	21.4	477
45	Determinants of barley grain yield in a wide range of Mediterranean environments. <i>Field Crops Research</i> , 2011, 120, 169-178.	5.1	73
46	Inside the CBF locus in Poaceae. <i>Plant Science</i> , 2011, 180, 39-45.	3.6	60
47	Diversity in the Response to Low Temperature in Representative Barley Genotypes Cultivated in Europe. <i>Crop Science</i> , 2011, 51, 2759-2779.	1.8	42
48	Epigenetic chromatin modifiers in barley: IV. The study of barley Polycomb group (PcG) genes during seed development and in response to external ABA. <i>BMC Plant Biology</i> , 2010, 10, 73.	3.6	63
49	Marker-assisted characterization of frost tolerance in barley (<i>Hordeum vulgare</i> L.). <i>Plant Breeding</i> , 2009, 128, 381-386.	1.9	29
50	QTL alleles from a winter feed type can improve malting quality in barley. <i>Plant Breeding</i> , 2009, 128, 598-605.	1.9	19
51	Epigenetic chromatin modifiers in barley: I. Cloning, mapping and expression analysis of the plant specific HD2 family of histone deacetylases from barley, during seed development and after hormonal treatment. <i>Physiologia Plantarum</i> , 2009, 136, 358-368.	5.2	65
52	Gene expression in grapevine cultivars in response to Bois Noir phytoplasma infection. <i>Plant Science</i> , 2009, 176, 792-804.	3.6	94
53	Barley adaptation and improvement in the Mediterranean basin. <i>Plant Breeding</i> , 2008, 127, 554-560.	1.9	40
54	Drought tolerance improvement in crop plants: An integrated view from breeding to genomics. <i>Field Crops Research</i> , 2008, 105, 1-14.	5.1	1,122

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55	Expression levels of barley <i>Cbf</i> genes at the Frost resistance <i>H2</i> locus are dependent upon alleles at <i>Fr-H1</i> and <i>Fr-H2</i> . <i>Plant Journal</i> , 2007, 51, 308-321.	5.7	121
56	Haplotype structure around the nud locus in barley and its association with resistance to leaf stripe (<i>Pyrenophora graminea</i>). <i>Plant Breeding</i> , 2007, 126, 24-29.	1.9	2
57	Fine mapping of a HvCBF gene cluster at the frost resistance locus Fr-H2 in barley. <i>Theoretical and Applied Genetics</i> , 2007, 115, 1083-1091.	3.6	145
58	Dual-purpose barley and oat in a Mediterranean environment. <i>Field Crops Research</i> , 2006, 99, 158-166.	5.1	48
59	Mapping regulatory genes as candidates for cold and drought stress tolerance in barley. <i>Theoretical and Applied Genetics</i> , 2006, 112, 445-454.	3.6	128
60	Marker assisted selection in crop plants. <i>Plant Cell, Tissue and Organ Culture</i> , 2005, 82, 317-342.	2.3	176
61	Molecular and Structural Characterization of Barley Vernalization Genes. <i>Plant Molecular Biology</i> , 2005, 59, 449-467.	3.9	258
62	Development of PCR-based markers on chromosome 5H for assisted selection of frost-tolerant genotypes in barley. <i>Molecular Breeding</i> , 2004, 14, 265-273.	2.1	21
63	Hv-WRKY38: a new transcription factor involved in cold- and drought-response in barley. <i>Plant Molecular Biology</i> , 2004, 55, 399-416.	3.9	273
64	Two loci on chromosome 5H determine low-temperature tolerance in a 'Nure' (winter) – 'Tremois' (spring) barley map. <i>Theoretical and Applied Genetics</i> , 2004, 108, 670-680.	3.6	199
65	Isolate-specific QTLs of resistance to leaf stripe (<i>Pyrenophora graminea</i>) in the 'Steptoe' – 'Morex' spring barley cross. <i>Theoretical and Applied Genetics</i> , 2003, 106, 668-675.	3.6	68
66	Genomic regions determining resistance to leaf stripe (<i>Pyrenophora graminea</i>) in barley. <i>Genome</i> , 2002, 45, 460-466.	2.0	24
67	Bioestimulants and cherry rootstock increased tomato fruit yield and quality in sustainable farming systems. <i>Italian Journal of Agronomy</i> , 0, , .	1.0	5