

Erica Lumini

List of Publications by Year in descending order

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63
papers

4,918
citations

172457

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133252

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all docs

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docs citations

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times ranked

4923
citing authors

#	ARTICLE	IF	CITATIONS
1	Alpine constructed wetlands: A metagenomic analysis reveals microbial complementary structure. <i>Science of the Total Environment</i> , 2022, 822, 153640.	8.0	3
2	Diversity of Arbuscular Mycorrhizal Fungi in olive orchard soils in arid regions of Southern Tunisia. <i>Arid Land Research and Management</i> , 2022, 36, 411-427.	1.6	5
3	Metabarcoding of Soil Fungal Communities Associated with Alpine Field-Grown Saffron (<i>Crocus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock	3.5	13
4	Strategies to Modulate Specialized Metabolism in Mediterranean Crops: From Molecular Aspects to Field. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2887.	4.1	29
5	Arbuscular Mycorrhizal Fungi from Argentinean Highland Puna Soils Unveiled by Propagule Multiplication. <i>Plants</i> , 2021, 10, 1803.	3.5	9
6	Impact of land use history on the arbuscular mycorrhizal fungal diversity in arid soils of Argentinean farming fields. <i>FEMS Microbiology Letters</i> , 2020, 367, .	1.8	9
7	Water management and phenology influence the root-associated rice field microbiota. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	2.7	28
8	Mining the Microbiome of Key Species from African Savanna Woodlands: Potential for Soil Health Improvement and Plant Growth Promotion. <i>Microorganisms</i> , 2020, 8, 1291.	3.6	11
9	Impact of an arbuscular mycorrhizal fungal inoculum and exogenous MeJA on fenugreek secondary metabolite production under water deficit. <i>Environmental and Experimental Botany</i> , 2020, 176, 104096.	4.2	23
10	Native Arbuscular Mycorrhizal Fungi Characterization from Saline Lands in Arid Oases, Northwest China. <i>Journal of Fungi (Basel, Switzerland)</i> , 2020, 6, 80.	3.5	8
11	Effects of Different Microbial Inocula on Tomato Tolerance to Water Deficit. <i>Agronomy</i> , 2020, 10, 170.	3.0	36
12	High-Throughput DNA Sequence-Based Analysis of AMF Communities. <i>Methods in Molecular Biology</i> , 2020, 2146, 99-116.	0.9	2
13	Arbuscular Mycorrhizal Fungi Modulate the Crop Performance and Metabolic Profile of Saffron in Soilless Cultivation. <i>Agronomy</i> , 2019, 9, 232.	3.0	48
14	Glomalin gene as molecular marker for functional diversity of arbuscular mycorrhizal fungi in soil. <i>Biology and Fertility of Soils</i> , 2019, 55, 411-417.	4.3	21
15	Saffron Cultivation in Marginal Alpine Environments: How AMF Inoculation Modulates Yield and Bioactive Compounds. <i>Agronomy</i> , 2019, 9, 12.	3.0	35
16	Focus on mycorrhizal symbioses. <i>Applied Soil Ecology</i> , 2018, 123, 299-304.	4.3	43
17	Impact of two arbuscular mycorrhizal fungi on <i>Arundo donax</i> L. response to salt stress. <i>Planta</i> , 2018, 247, 573-585.	3.2	62
18	Seasonal variation in winter wheat field soil arbuscular mycorrhizal fungus communities after non-mycorrhizal crop cultivation. <i>Mycorrhiza</i> , 2018, 28, 535-548.	2.8	16

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19	Soil bacterial networks are less stable under drought than fungal networks. <i>Nature Communications</i> , 2018, 9, 3033.	12.8	992
20	The last 50 years of climate-induced melting of the Maliy Aktru glacier (Altai Mountains, Russia) revealed in a primary ecological succession. <i>Ecology and Evolution</i> , 2018, 8, 7401-7420.	1.9	18
21	Differential biodiversity responses between kingdoms (plants, fungi, bacteria and metazoa) along an Alpine succession gradient. <i>Molecular Ecology</i> , 2018, 27, 3671-3685.	3.9	33
22	Wild <i>Camellia japonica</i> specimens in the Shimane prefecture (Japan) host previously undescribed AMF diversity. <i>Applied Soil Ecology</i> , 2017, 115, 10-18.	4.3	11
23	Arbuscular mycorrhizal fungal community differences among European long-term observatories. <i>Mycorrhiza</i> , 2017, 27, 331-343.	2.8	14
24	AMF components from a microbial inoculum fail to colonize roots and lack soil persistence in an arable maize field. <i>Symbiosis</i> , 2017, 72, 73-80.	2.3	25
25	The abundance and diversity of arbuscular mycorrhizal fungi are linked to the soil chemistry of screes and to slope in the Alpic paleo-endemic <i>Berardia subacaulis</i> . <i>PLoS ONE</i> , 2017, 12, e0171866.	2.5	39
26	Insights On the Impact of Arbuscular Mycorrhizal Symbiosis On Tomato Tolerance to Water Stress. <i>Plant Physiology</i> , 2016, 171, pp.00307.2016.	4.8	227
27	Transfiguring biodegradation of frescoes in the Beata Vergine del Pilone Sanctuary (Italy): Microbial analysis and minero-chemical aspects. <i>International Biodeterioration and Biodegradation</i> , 2015, 98, 6-18.	3.9	10
28	Arbuscular mycorrhizal fungal diversity in the Tuber <i>melanosporum</i> brÃ»lÃ©. <i>Fungal Biology</i> , 2015, 119, 518-527.	2.5	20
29	Edaphic factors trigger diverse AM fungal communities associated to exotic camellias in closely located Lake Maggiore (Italy) sites. <i>Mycorrhiza</i> , 2015, 25, 253-265.	2.8	25
30	Arbuscular Mycorrhizal Fungi as Natural Biofertilizers: Let's Benefit from Past Successes. <i>Frontiers in Microbiology</i> , 2015, 6, 1559.	3.5	543
31	Arbuscular Mycorrhizal Fungi and their Value for Ecosystem Management. , 2014, , .		22
32	Sequencing and comparison of the mitochondrial COI gene from isolates of Arbuscular Mycorrhizal Fungi belonging to Gigasporaceae and Glomeraceae families. <i>Molecular Phylogenetics and Evolution</i> , 2014, 75, 1-10.	2.7	13
33	Application of laser microdissection to identify the mycorrhizal fungi that establish arbuscules inside root cells. <i>Frontiers in Plant Science</i> , 2013, 4, 135.	3.6	33
34	454 Pyrosequencing Analysis of Fungal Assemblages from Geographically Distant, Disparate Soils Reveals Spatial Patterning and a Core Mycobiome. <i>Diversity</i> , 2013, 5, 73-98.	1.7	82
35	The genome of the obligate endobacterium of an AM fungus reveals an interphylum network of nutritional interactions. <i>ISME Journal</i> , 2012, 6, 136-145.	9.8	176
36	Discrimination of <i>Gigaspora</i> species by PCR specific primers and phylogenetic analysis. <i>Mycotaxon</i> , 2012, 118, 17-26.	0.3	1

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37	Effects of different management practices on arbuscular mycorrhizal fungal diversity in maize fields by a molecular approach. <i>Biology and Fertility of Soils</i> , 2012, 48, 911-922.	4.3	95
38	Unravelling Soil Fungal Communities from Different Mediterranean Land-Use Backgrounds. <i>PLoS ONE</i> , 2012, 7, e34847.	2.5	194
39	SELECTION OF ARBUSCULAR MYCORRHIZAL FUNGAL ISOLATES FOR SUSTAINABLE FLORICULTURE. <i>Acta Horticulturae</i> , 2011, , 319-324.	0.2	1
40	Different farming and water regimes in Italian rice fields affect arbuscular mycorrhizal fungal soil communities. , 2011, 21, 1696-1707.		99
41	Unique arbuscular mycorrhizal fungal communities uncovered in date palm plantations and surrounding desert habitats of Southern Arabia. <i>Mycorrhiza</i> , 2011, 21, 195-209.	2.8	55
42	Disclosing arbuscular mycorrhizal fungal biodiversity in soil through a landscape gradient using a pyrosequencing approach. <i>Environmental Microbiology</i> , 2010, 12, 2165-2179.	3.8	313
43	Cohorts of arbuscular mycorrhizal fungi (AMF) in <i>Vitis vinifera</i> , a typical Mediterranean fruit crop. <i>Environmental Microbiology Reports</i> , 2010, 2, 594-604.	2.4	77
44	The <i>ftsZ</i> Gene of the Endocellular Bacterium <i>Candidatus</i> <i>Glomeribacter gigasporarum</i> Is Preferentially Expressed During the Symbiotic Phases of Its Host Mycorrhizal Fungus. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 302-310.	2.6	31
45	Simultaneous detection and quantification of the unculturable microbe <i>Candidatus</i> <i>Glomeribacter gigasporarum</i> inside its fungal host <i>Gigaspora margarita</i> . <i>New Phytologist</i> , 2008, 180, 248-257.	7.3	31
46	THE IMPACT OF TILLAGE PRACTICES ON ARBUSCULAR MYCORRHIZAL FUNGAL DIVERSITY IN SUBTROPICAL CROPS. , 2008, 18, 527-536.		172
47	Glomeromycotean associations in liverworts: a molecular, cellular, and taxonomic analysis. <i>American Journal of Botany</i> , 2007, 94, 1756-1777.	1.7	141
48	Presymbiotic growth and spore morphology are affected in the arbuscular mycorrhizal fungus <i>Gigaspora margarita</i> cured of its endobacteria. <i>Cellular Microbiology</i> , 2007, 9, 1716-1729.	2.1	140
49	Assessment of arbuscular mycorrhizal fungal diversity in roots of <i>Solidago gigantea</i> growing in a polluted soil in Northern Italy. <i>Environmental Microbiology</i> , 2006, 8, 971-983.	3.8	109
50	Endobacteria or bacterial endosymbionts? To be or not to be. <i>New Phytologist</i> , 2006, 170, 205-208.	7.3	32
51	Phylogenetic analysis of Glomeromycota by partial LSU rDNA sequences. <i>Mycorrhiza</i> , 2006, 16, 183-189.	2.8	57
52	Vertical Transmission of Endobacteria in the Arbuscular Mycorrhizal Fungus <i>Gigaspora margarita</i> through Generation of Vegetative Spores. <i>Applied and Environmental Microbiology</i> , 2004, 70, 3600-3608.	3.1	126
53	<i>Candidatus</i> <i>Glomeribacter gigasporarum</i> gen. nov., sp. nov., an endosymbiont of arbuscular mycorrhizal fungi. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2003, 53, 121-124.	1.7	188
54	A combined morphological and molecular approach to characterize isolates of arbuscular mycorrhizal fungi in <i>Gigaspora</i> (Glomales). <i>New Phytologist</i> , 2001, 152, 169-179.	7.3	25

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55	Detection and Identification of Bacterial Endosymbionts in Arbuscular Mycorrhizal Fungi Belonging to the Family Gigasporaceae. Applied and Environmental Microbiology, 2000, 66, 4503-4509.	3.1	156
56	Polymerase chain reaction - restriction fragment length polymorphisms for assessing and increasing biodiversity of <i>Frankia</i> culture collections. Canadian Journal of Botany, 1999, 77, 1261-1269.	1.1	6
57	The Nuclear Ribosomal DNA Intergenic Spacer as a Target Sequence To Study Intraspecific Diversity of the Ectomycorrhizal Basidiomycete <i>Hebeloma cylindrosporum</i> Directly on <i>Pinus</i> Root Systems. Applied and Environmental Microbiology, 1999, 65, 903-909.	3.1	51
58	Polymerase chain reaction - restriction fragment length polymorphisms for assessing and increasing biodiversity of <i>Frankia</i> culture collections. Canadian Journal of Botany, 1999, 77, 1261-1269.	1.1	19
59	PCR-RFLP and total DNA homology revealed three related genomic species among broad-host-range <i>Frankia</i> strains. FEMS Microbiology Ecology, 1996, 21, 303-311.	2.7	34
60	PCR-restriction fragment length polymorphism identification and host range of single-spore isolates of the flexible <i>Frankia</i> sp. strain UFI 132715. Applied and Environmental Microbiology, 1996, 62, 3026-3029.	3.1	29
61	PCR-RFLP and total DNA homology revealed three related genomic species among broad-host-range <i>Frankia</i> strains. FEMS Microbiology Ecology, 1996, 21, 303-311.	2.7	1
62	Field performance of <i>Alnus cordata</i> loisel (Italian alder) inoculated with <i>Frankia</i> and VA-mycorrhizal strains in mine-spoil afforestation plots. Soil Biology and Biochemistry, 1994, 26, 659-661.	8.8	51
63	Botanica Applicata. Giornale Botanico Italiano (Florence, Italy: 1962), 1993, 127, 521-530.	0.0	0