

# Erica Lumini

## List of Publications by Year in descending order

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63  
papers

4,918  
citations

172457

29  
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133252

59  
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63  
all docs

63  
docs citations

63  
times ranked

4923  
citing authors

#	ARTICLE	IF	CITATIONS
1	Soil bacterial networks are less stable under drought than fungal networks. <i>Nature Communications</i> , 2018, 9, 3033.	12.8	992
2	Arbuscular Mycorrhizal Fungi as Natural Biofertilizers: Let's Benefit from Past Successes. <i>Frontiers in Microbiology</i> , 2015, 6, 1559.	3.5	543
3	Disclosing arbuscular mycorrhizal fungal biodiversity in soil through a land-use gradient using a pyrosequencing approach. <i>Environmental Microbiology</i> , 2010, 12, 2165-2179.	3.8	313
4	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
5	Unravelling Soil Fungal Communities from Different Mediterranean Land-Use Backgrounds. <i>PLoS ONE</i> , 2012, 7, e34847.	2.5	194
6	â€Candidatus <i>Glomeribacter gigasporarum</i> â€™ 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
7	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
8	THE IMPACT OF TILLAGE PRACTICES ON ARBUSCULAR MYCORRHIZAL FUNGAL DIVERSITY IN SUBTROPICAL CROPS. , 2008, 18, 527-536.		172
9	Detection and Identification of Bacterial Endosymbionts in Arbuscular Mycorrhizal Fungi Belonging to the Family Gigasporaceae. <i>Applied and Environmental Microbiology</i> , 2000, 66, 4503-4509.	3.1	156
10	Glomeromycotean associations in liverworts: a molecular, cellular, and taxonomic analysis. <i>American Journal of Botany</i> , 2007, 94, 1756-1777.	1.7	141
11	Presymbiotic growth and sporal 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
12	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
13	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
14	Different farming and water regimes in Italian rice fields affect arbuscular mycorrhizal fungal soil communities. , 2011, 21, 1696-1707.		99
15	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
16	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
17	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
18	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

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19	Phylogenetic analysis of Glomeromycota by partial LSU rDNA sequences. <i>Mycorrhiza</i> , 2006, 16, 183-189.	2.8	57
20	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
21	Field performance of <i>Alnus cordata</i> loisel (Italian alder) inoculated with Frankia and VA-mycorrhizal strains in mine-spoil afforestation plots. <i>Soil Biology and Biochemistry</i> , 1994, 26, 659-661.	8.8	51
22	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. <i>Applied and Environmental Microbiology</i> , 1999, 65, 903-909.	3.1	51
23	Arbuscular Mycorrhizal Fungi Modulate the Crop Performance and Metabolic Profile of Saffron in Soilless Cultivation. <i>Agronomy</i> , 2019, 9, 232.	3.0	48
24	Focus on mycorrhizal symbioses. <i>Applied Soil Ecology</i> , 2018, 123, 299-304.	4.3	43
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	Effects of Different Microbial Inocula on Tomato Tolerance to Water Deficit. <i>Agronomy</i> , 2020, 10, 170.	3.0	36
27	Saffron Cultivation in Marginal Alpine Environments: How AMF Inoculation Modulates Yield and Bioactive Compounds. <i>Agronomy</i> , 2019, 9, 12.	3.0	35
28	PCR-RFLP and total DNA homology revealed three related genomic species among broad-host-range Frankia strains. <i>FEMS Microbiology Ecology</i> , 1996, 21, 303-311.	2.7	34
29	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
30	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
31	Endobacteria or bacterial endosymbionts? To be or not to be. <i>New Phytologist</i> , 2006, 170, 205-208.	7.3	32
32	Simultaneous detection and quantification of the unculturable microbe <i>Candidatus</i> Glomeribacter gigasporarum inside its fungal host <i>Gigaspora margarita</i> . <i>New Phytologist</i> , 2008, 180, 248-257.	7.3	31
33	The <i>ftsZ</i> Gene of the Endocellular Bacterium <i>Candidatus</i> Glomeribacter gigasporarum <sup>TM</sup> 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
34	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
35	PCR-restriction fragment length polymorphism identification and host range of single-spore isolates of the flexible Frankia sp. strain UFI 132715. <i>Applied and Environmental Microbiology</i> , 1996, 62, 3026-3029.	3.1	29
36	Water management and phenology influence the root-associated rice field microbiota. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	2.7	28

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37	A combined morphological and molecular approach to characterize isolates of arbuscular mycorrhizal fungi in Gigaspora (Glomales). <i>New Phytologist</i> , 2001, 152, 169-179.	7.3	25
38	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
39	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
40	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
41	Arbuscular Mycorrhizal Fungi and their Value for Ecosystem Management. , 2014, , .		22
42	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
43	Arbuscular mycorrhizal fungal diversity in the Tuber melanosporum brÃ»lÃ©. <i>Fungal Biology</i> , 2015, 119, 518-527.	2.5	20
44	Polymerase chain reaction - restriction fragment length polymorphisms for assessing and increasing biodiversity of <i>Frankia</i> culture collections. <i>Canadian Journal of Botany</i> , 1999, 77, 1261-1269.	1.1	19
45	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
46	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
47	Arbuscular mycorrhizal fungal community differences among European long-term observatories. <i>Mycorrhiza</i> , 2017, 27, 331-343.	2.8	14
48	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
49	Metabarcoding of Soil Fungal Communities Associated with Alpine Field-Grown Saffron ( <i>Crocus</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock	3.5	13
50	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
51	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
52	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
53	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
54	Arbuscular Mycorrhizal Fungi from Argentinean Highland Puna Soils Unveiled by Propagule Multiplication. <i>Plants</i> , 2021, 10, 1803.	3.5	9

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55	Native Arbuscular Mycorrhizal Fungi Characterization from Saline Lands in Arid Oases, Northwest China. <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 80.	3.5	8
56	Polymerase chain reaction - restriction fragment length polymorphisms for assessing and increasing biodiversity of <i>Frankia</i> culture collections. <i>Canadian Journal of Botany</i> , 1999, 77, 1261-1269.	1.1	6
57	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
58	Alpine constructed wetlands: A metagenomic analysis reveals microbial complementary structure. <i>Science of the Total Environment</i> , 2022, 822, 153640.	8.0	3
59	High-Throughput DNA Sequence-Based Analysis of AMF Communities. <i>Methods in Molecular Biology</i> , 2020, 2146, 99-116.	0.9	2
60	SELECTION OF ARBUSCULAR MYCORRHIZAL FUNGAL ISOLATES FOR SUSTAINABLE FLORICULTURE. <i>Acta Horticulturae</i> , 2011, , 319-324.	0.2	1
61	Discrimination of <i>Gigaspora</i> species by PCR specific primers and phylogenetic analysis. <i>Mycotaxon</i> , 2012, 118, 17-26.	0.3	1
62	PCR-RFLP and total DNA homology revealed three related genomic species among broad-host-range <i>Frankia</i> strains. <i>FEMS Microbiology Ecology</i> , 1996, 21, 303-311.	2.7	1
63	<i>Botanica Applicata</i> . <i>Giornale Botanico Italiano</i> (Florence, Italy: 1962), 1993, 127, 521-530.	0.0	0