

François Lutzoni

List of Publications by Year in descending order

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Version: 2024-02-01

59
papers

5,046
citations

126907

33
h-index

128289

60
g-index

60
all docs

60
docs citations

60
times ranked

4013
citing authors

#	ARTICLE	IF	CITATIONS
1	Ecological generalism drives hyperdiversity of secondary metabolite gene clusters in xylarialean endophytes. <i>New Phytologist</i> , 2022, 233, 1317-1330.	7.3	23
2	Community dynamics of soil-borne fungal communities along elevation gradients in neotropical and palaeotropical forests. <i>Molecular Ecology</i> , 2022, 31, 2044-2060.	3.9	11
3	Comparative transcriptomics of fungal endophytes in co-culture with their moss host <i>Dicranum scoparium</i> reveals fungal trophic lability and moss unchanged to slightly increased growth rates. <i>New Phytologist</i> , 2022, 234, 1832-1847.	7.3	5
4	Phylogenetic structure of specialization: A new approach that integrates partner availability and phylogenetic diversity to quantify biotic specialization in ecological networks. <i>Ecology and Evolution</i> , 2022, 12, e8649.	1.9	6
5	Climate and seasonality drive the richness and composition of tropical fungal endophytes at a landscape scale. <i>Communications Biology</i> , 2021, 4, 313.	4.4	45
6	Exploring syntenic conservation across genomes for phylogenetic studies of organisms subjected to horizontal gene transfers: A case study with Cyanobacteria and cyanolichens. <i>Molecular Phylogenetics and Evolution</i> , 2021, 162, 107100.	2.7	8
7	Cyanolichen microbiome contains novel viruses that encode genes to promote microbial metabolism. <i>ISME Communications</i> , 2021, 1, .	4.2	3
8	Turnover of Lecanoroid Mycobionts and Their Trebouxia Photobionts Along an Elevation Gradient in Bolivia Highlights the Role of Environment in Structuring the Lichen Symbiosis. <i>Frontiers in Microbiology</i> , 2021, 12, 774839.	3.5	16
9	T-BAS Version 2.1: Tree-Based Alignment Selector Toolkit for Evolutionary Placement of DNA Sequences and Viewing Alignments and Specimen Metadata on Curated and Custom Trees. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	35
10	The lichen symbiosis re-viewed through the genomes of <i>Cladonia grayi</i> and its algal partner <i>Asterochloris glomerata</i> . <i>BMC Genomics</i> , 2019, 20, 605.	2.8	98
11	Host availability drives distributions of fungal endophytes in the imperilled boreal realm. <i>Nature Ecology and Evolution</i> , 2019, 3, 1430-1437.	7.8	91
12	Species diversification and phylogenetically constrained symbiont switching generated high modularity in the lichen genus <i>Peltigera</i> . <i>Journal of Ecology</i> , 2019, 107, 1645-1661.	4.0	20
13	Differential gene expression associated with fungal trophic shifts along the senescence gradient of the moss <i>Dicranum scoparium</i> . <i>Environmental Microbiology</i> , 2019, 21, 2273-2289.	3.8	11
14	Molybdenum threshold for ecosystem scale alternative vanadium nitrogenase activity in boreal forests. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 24682-24688.	7.1	60
15	Biodiversity assessment of ascomycetes inhabiting <i>Lobariella</i> lichens in Andean cloud forests led to one new family, three new genera and 13 new species of lichenicolous fungi. <i>Plant and Fungal Systematics</i> , 2019, 64, 283-344.	0.5	30
16	scRNA-based analyses reveal fungal communities structured by a senescence gradient in the moss <i>Dicranum scoparium</i> and the presence of putative multi-trophic fungi. <i>New Phytologist</i> , 2018, 218, 1597-1611.	7.3	44
17	Contemporaneous radiations of fungi and plants linked to symbiosis. <i>Nature Communications</i> , 2018, 9, 5451.	12.8	189
18	Contrasting Symbiotic Patterns in Two Closely Related Lineages of Trimembered Lichens of the Genus <i>Peltigera</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 2770.	3.5	25

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19	Stable isotope analyses reveal previously unknown trophic mode diversity in the Hymenochaetales. <i>American Journal of Botany</i> , 2018, 105, 1869-1887.	1.7	19
20	Species delimitation at a global scale reveals high species richness with complex biogeography and patterns of symbiont association in <i>Peltigera</i> section <i>Peltigera</i> (lichenized Ascomycota): Tj ETQq0 0 OrgBT /Overclock 10 T		
21	Bioclimatic factors at an intrabiome scale are more limiting than cyanobiont availability for the lichen-forming genus <i>Peltigera</i> . <i>American Journal of Botany</i> , 2018, 105, 1198-1211.	1.7	19
22	Molecular data favours a monogeneric <i>Peltulaceae</i> (Lichinomycetes). <i>Lichenologist</i> , 2018, 50, 313-327.	0.8	9
23	Species in section <i>Peltidea</i> (aphthosa group) of the genus <i>Peltigera</i> remain cryptic after molecular phylogenetic revision. <i>Plant and Fungal Systematics</i> , 2018, 63, 45-64.	0.5	12
24	T-BAS: Tree-Based Alignment Selector toolkit for phylogenetic-based placement, alignment downloads and metadata visualization: an example with the Pezizomycotina tree of life. <i>Bioinformatics</i> , 2017, 33, 1160-1168.	4.1	55
25	Conserved genomic collinearity as a source of broadly applicable, fast evolving, markers to resolve species complexes: A case study using the lichen-forming genus <i>Peltigera</i> section <i>Polydactylon</i> . <i>Molecular Phylogenetics and Evolution</i> , 2017, 117, 10-29.	2.7	30
26	Biological nitrogen fixation by alternative nitrogenases in boreal cyanolichens: importance of molybdenum availability and implications for current biological nitrogen fixation estimates. <i>New Phytologist</i> , 2017, 213, 680-689.	7.3	54
27	<i>A Liber Amicorum: Irwin Brodo</i> . <i>Lichenologist</i> , 2016, 48, 343-346.	0.8	1
28	Photobiont associations in co-occurring umbilicate lichens with contrasting modes of reproduction in coastal Norway. <i>Lichenologist</i> , 2016, 48, 545-557.	0.8	13
29	Contributions of North American endophytes to the phylogeny, ecology, and taxonomy of <i>Xylariaceae</i> (Sordariomycetes, Ascomycota). <i>Molecular Phylogenetics and Evolution</i> , 2016, 98, 210-232.	2.7	110
30	Interaction type influences ecological network structure more than local abiotic conditions: evidence from endophytic and endolichenic fungi at a continental scale. <i>Oecologia</i> , 2016, 180, 181-191.	2.0	50
31	Phylogenetic analyses of eurotiomycetous endophytes reveal their close affinities to Chaetothiales, Eurotiales, and a new order "Phaeomoniellales. <i>Molecular Phylogenetics and Evolution</i> , 2015, 85, 117-130.	2.7	66
32	Determination of elemental baseline using peltigerelean lichens from Northeastern Canada (Québec): Initial data collection for long term monitoring of the impact of global climate change on boreal and subarctic area in Canada. <i>Science of the Total Environment</i> , 2015, 533, 1-7.	8.0	26
33	A comparison of the community diversity of foliar fungal endophytes between seedling and adult loblolly pines (<i>Pinus taeda</i>). <i>Fungal Biology</i> , 2015, 119, 917-928.	2.5	79
34	The adaptive radiation of lichen-forming Teloschistaceae is associated with sunscreens pigments and a bark-to-rock substrate shift. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 11600-11605.	7.1	77
35	Is vanadium a biometal for boreal cyanolichens?. <i>New Phytologist</i> , 2014, 202, 765-771.	7.3	34
36	A multigene phylogenetic synthesis for the class Lecanoromycetes (Ascomycota): 1307 fungi representing 1139 infrageneric taxa, 317 genera and 66 families. <i>Molecular Phylogenetics and Evolution</i> , 2014, 79, 132-168.	2.7	248

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37	Phylogenetic placement, species delimitation, and cyanobiont identity of endangered aquatic <i>Peltigera</i> species (lichen-forming Ascomycota, Lecanoromycetes). <i>American Journal of Botany</i> , 2014, 101, 1141-1156.	1.7	37
38	Lichen-symbiotic cyanobacteria associated with <i>Peltigera</i> have an alternative vanadium-dependent nitrogen fixation system. <i>European Journal of Phycology</i> , 2014, 49, 11-19.	2.0	50
39	Assessing population structure and host specialization in lichenized cyanobacteria. <i>New Phytologist</i> , 2013, 198, 557-566.	7.3	57
40	Twenty-five cultures of lichenizing fungi available for experimental studies on symbiotic systems. <i>Symbiosis</i> , 2013, 59, 165-171.	2.3	31
41	Phylogenetic study of <i>Diploschistes</i> (lichen-forming Ascomycota: Ostropales: Graphidaceae), based on morphological, chemical, and molecular data. <i>Taxon</i> , 2013, 62, 267-280.	0.7	16
42	Host and geographic structure of endophytic and endolichenic fungi at a continental scale. <i>American Journal of Botany</i> , 2012, 99, 898-914.	1.7	304
43	Hypogymnia phylogeny, including Cavernularia, reveals biogeographic structure. <i>Bryologist</i> , 2011, 114, 392.	0.6	27
44	Revisiting photobiont diversity in the lichen family Verrucariaceae (Ascomycota). <i>European Journal of Phycology</i> , 2011, 46, 399-415.	2.0	148
45	Community Analysis Reveals Close Affinities Between Endophytic and Endolichenic Fungi in Mosses and Lichens. <i>Microbial Ecology</i> , 2010, 60, 340-353.	2.8	191
46	Phylogenetic generic classification of parmelioid lichens (Parmeliaceae, Ascomycota) based on molecular, morphological and chemical evidence. <i>Taxon</i> , 2010, 59, 1735-1753.	0.7	178
47	Generic classification of the Verrucariaceae (Ascomycota) based on molecular and morphological evidence: recent progress and remaining challenges. <i>Taxon</i> , 2009, 58, 184-208.	0.7	88
48	Lichens. <i>Current Biology</i> , 2009, 19, R502-R503.	3.9	62
49	Compatibility and thigmotropism in the lichen symbiosis: A reappraisal. <i>Symbiosis</i> , 2009, 47, 109-115.	2.3	41
50	A microbiotic survey of lichen-associated bacteria reveals a new lineage from the Rhizobiales. <i>Symbiosis</i> , 2009, 49, 163-180.	2.3	201
51	A Phylogenetic Estimation of Trophic Transition Networks for Ascomycetous Fungi: Are Lichens Cradles of Symbiotrophic Fungal Diversification?. <i>Systematic Biology</i> , 2009, 58, 283-297.	5.6	321
52	Phylogenetic relationships and taxonomy of the <i>Leptogium lichenoides</i> group (Collemataceae, Tj ETQq0 0 0 rgBT /Overlock 10 T	0.7	28
53	Phylogenetic comparison of protein-coding versus ribosomal RNA-coding sequence data: A case study of the Lecanoromycetes (Ascomycota). <i>Molecular Phylogenetics and Evolution</i> , 2007, 44, 412-426.	2.7	144
54	New insights into classification and evolution of the Lecanoromycetes (Pezizomycotina, Ascomycota) from phylogenetic analyses of three ribosomal RNA- and two protein-coding genes. <i>Mycologia</i> , 2006, 98, 1088-1103.	1.9	140

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55	New insights into classification and evolution of the Lecanoromycetes (Pezizomycotina, Ascomycota) from phylogenetic analyses of three ribosomal RNA- and two protein-coding genes. <i>Mycologia</i> , 2006, 98, 1088-1103.	1.9	227
56	Assessing host specialization in symbiotic cyanobacteria associated with four closely related species of the lichen fungus <i>Peltigera</i> . <i>European Journal of Phycology</i> , 2005, 40, 363-378.	2.0	117
57	Assembling the fungal tree of life: progress, classification, and evolution of subcellular traits. <i>American Journal of Botany</i> , 2004, 91, 1446-1480.	1.7	718
58	New Approach to an Old Problem: Incorporating Signal from Gap-Rich Regions of ITS and rDNA Large Subunit into Phylogenetic Analyses to Resolve the <i>Peltigera canina</i> Species Complex. <i>Mycologia</i> , 2003, 95, 1181.	1.9	19
59	Phylogeny of the Gyalectales and Ostropales (Ascomycota, Fungi): among and within order relationships based on nuclear ribosomal RNA small and large subunits. <i>Molecular Phylogenetics and Evolution</i> , 2002, 25, 138-156.	2.7	237