Francois Buscot

List of Publications by Year
in descending order

249 papers	18,036 citations	12330 69 h-index	18130 120 g-index
258 all docs	258 docs citations	258 times ranked	17678 citing authors

#	ARTICLE	IF	CITATIONS
1	Convergent losses of decay mechanisms and rapid turnover of symbiosis genes in mycorrhizal mutualists. Nature Genetics, 2015, 47, 410-415.	21.4	870
2	Bottom-up effects of plant diversity on multitrophic interactions in a biodiversity experiment. Nature, 2010, 468, 553-556.	27.8	786
3	Choosing and using diversity indices: insights for ecological applications from the German Biodiversity Exploratories. Ecology and Evolution, 2014, 4, 3514-3524.	1.9	697
4	Implementing large-scale and long-term functional biodiversity research: The Biodiversity Exploratories. Basic and Applied Ecology, 2010, 11, 473-485.	2.7	649
5	Biodiversity at multiple trophic levels is needed for ecosystem multifunctionality. Nature, 2016, 536, 456-459.	27.8	526
6	Mineral vs. Organic Amendments: Microbial Community Structure, Activity and Abundance of Agriculturally Relevant Microbes Are Driven by Long-Term Fertilization Strategies. Frontiers in Microbiology, 2016, 7, 1446.	3.5	462
7	Impacts of species richness on productivity in a large-scale subtropical forest experiment. Science, 2018, 362, 80-83.	12.6	433
8	Land-use intensification causes multitrophic homogenization of grassland communities. Nature, 2016, 540, 266-269.	27.8	404
9	A quantitative index of land-use intensity in grasslands: Integrating mowing, grazing and fertilization. Basic and Applied Ecology, 2012, 13, 207-220.	2.7	325
10	Horizon-Specific Bacterial Community Composition of German Grassland Soils, as Revealed by Pyrosequencing-Based Analysis of 16S rRNA Genes. Applied and Environmental Microbiology, 2010, 76, 6751-6759.	3.1	312
11	Biodiversity effects on ecosystem functioning in a 15-year grassland experiment: Patterns, mechanisms, and open questions. Basic and Applied Ecology, 2017, 23, 1-73.	2.7	307
12	Life in leaf litter: novel insights into community dynamics of bacteria and fungi during litter decomposition. Molecular Ecology, 2016, 25, 4059-4074.	3.9	297
13	Interannual variation in land-use intensity enhances grassland multidiversity. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 308-313.	7.1	243
14	Designing forest biodiversity experiments: general considerations illustrated by a new large experiment in subtropical <scp>C</scp>hina. Methods in Ecology and Evolution, 2014, 5, 74-89.	5.2	232
15	Community assembly during secondary forest succession in a Chinese subtropical forest. Ecological Monographs, 2011, 81, 25-41.	5.4	222
16	Differences in the species composition of arbuscular mycorrhizal fungi in spore, root and soil communities in a grassland ecosystem. Environmental Microbiology, 2007, 9, 1930-1938.	3.8	218
17	Blind spots in global soil biodiversity and ecosystem function research. Nature Communications, 2020, 11, 3870.	12.8	192
18	The impact of evenâ€aged and unevenâ€aged forest management on regional biodiversity of multiple taxa in European beech forests. Journal of Applied Ecology, 2018, 55, 267-278.	4.0	188

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19	Multiple forest attributes underpin the supply of multiple ecosystem services. Nature Communications, 2018, 9, 4839.	12.8	182
20	Interacting effects of fertilization, mowing and grazing on plant species diversity of 1500 grasslands in Germany differ between regions. Basic and Applied Ecology, 2013, 14, 126-136.	2.7	177
21	Biodiversity across trophic levels drives multifunctionality in highly diverse forests. Nature Communications, 2018, 9, 2989.	12.8	169
22	Land-use intensity alters networks between biodiversity, ecosystem functions, and services. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 28140-28149.	7.1	164
23	Effects of longâ€term differential fertilization on eukaryotic microbial communities in an arable soil: a multiple barcoding approach. Molecular Ecology, 2014, 23, 3341-3355.	3.9	163
24	Soilâ€carbon preservation through habitat constraints and biological limitations on decomposer activity. Journal of Plant Nutrition and Soil Science, 2008, 171, 27-35.	1.9	156
25	Molecular diversity of arbuscular mycorrhizal fungi in relation to soil chemical properties and heavy metal contamination. Environmental Pollution, 2010, 158, 2757-2765.	7.5	152
26	Wood decay rates of 13 temperate tree species in relation to wood properties, enzyme activities and organismic diversities. Forest Ecology and Management, 2017, 391, 86-95.	3.2	151
27	Tracking, targeting, and conserving soil biodiversity. Science, 2021, 371, 239-241.	12.6	151
28	Resource Partitioning between Bacteria, Fungi, and Protists in the Detritusphere of an Agricultural Soil. Frontiers in Microbiology, 2016, 7, 1524.	3.5	143
29	General Relationships between Abiotic Soil Properties and Soil Biota across Spatial Scales and Different Land-Use Types. PLoS ONE, 2012, 7, e43292.	2.5	142
30	Linking molecular deadwood-inhabiting fungal diversity and community dynamics to ecosystem functions and processes in Central European forests. Fungal Diversity, 2016, 77, 367-379.	12.3	140
31	Diversity of laccase genes from basidiomycetes in a forest soil. Soil Biology and Biochemistry, 2004, 36, 1025-1036.	8.8	136
32	Forest Management Type Influences Diversity and Community Composition of Soil Fungi across Temperate Forest Ecosystems. Frontiers in Microbiology, 2015, 6, 1300.	3.5	136
33	Host plant genusâ€level diversity is the best predictor of ectomycorrhizal fungal diversity in a Chinese subtropical forest. Molecular Ecology, 2013, 22, 3403-3414.	3.9	133
34	Network Analysis Reveals Ecological Links between N-Fixing Bacteria and Wood-Decaying Fungi. PLoS ONE, 2014, 9, e88141.	2.5	129
35	Nitrogen supply affects arbuscular mycorrhizal colonization of Artemisia vulgaris in a phosphateâ€polluted field site. New Phytologist, 2005, 166, 981-992.	7.3	117
36	Locally rare species influence grassland ecosystem multifunctionality. Philosophical Transactions of the Royal Society B: Biological Sciences, 2016, 371, 20150269.	4.0	117

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37	Protein-SIP enables time-resolved analysis of the carbon flux in a sulfate-reducing, benzene-degrading microbial consortium. ISME Journal, 2012, 6, 2291-2301.	9.8	109
38	Community assembly of ectomycorrhizal fungi along a subtropical secondary forest succession. New Phytologist, 2015, 205, 771-785.	7.3	107
39	Diversity of bacterial laccase-like multicopper oxidase genes in forest and grassland Cambisol soil samples. Soil Biology and Biochemistry, 2008, 40, 638-648.	8.8	104
40	High plant species richness indicates management-related disturbances rather than the conservation status of forests. Basic and Applied Ecology, 2013, 14, 496-505.	2.7	102
41	On the combined effect of soil fertility and topography on tree growth in subtropical forest ecosystemsâ€"a study from SE China. Journal of Plant Ecology, 2017, 10, 111-127.	2.3	102
42	<i>Streptomyces</i>-Induced Resistance Against Oak Powdery Mildew Involves Host Plant Responses in Defense, Photosynthesis, and Secondary Metabolism Pathways. Molecular Plant-Microbe Interactions, 2014, 27, 891-900.	2.6	101
43	A pyrosequencing insight into sprawling bacterial diversity and community dynamics in decaying deadwood logs of Fagus sylvatica and Picea abies. Scientific Reports, 2015, 5, 9456.	3.3	101
44	Recent advances in exploring physiology and biodiversity of ectomycorrhizas highlight the functioning of these symbioses in ecosystems. FEMS Microbiology Reviews, 2000, 24, 601-614.	8.6	98
45	OakContig<scp>DF</scp>159.1, a reference library for studying differential gene expression in <i>Quercus robur</i> during controlled biotic interactions: use for quantitative transcriptomic profiling of oak roots in ectomycorrhizal symbiosis. New Phytologist, 2013, 199, 529-540.	7.3	97
46	Diversity Promotes Temporal Stability across Levels of Ecosystem Organization in Experimental Grasslands. PLoS ONE, 2010, 5, e13382.	2.5	95
47	pH as a Driver for Ammonia-Oxidizing Archaea in Forest Soils. Microbial Ecology, 2015, 69, 879-883.	2.8	95
48	A multitrophic perspective on biodiversityâ€"ecosystem functioning research. Advances in Ecological Research, 2019, 61, 1-54.	2.7	95
49	Patchiness and Spatial Distribution of Laccase Genes of Ectomycorrhizal, Saprotrophic, and Unknown Basidiomycetes in the Upper Horizons of a Mixed Forest Cambisol. Microbial Ecology, 2005, 50, 570-579.	2.8	94
50	Differences in Soil Fungal Communities between European Beech (Fagus sylvatica L.) Dominated Forests Are Related to Soil and Understory Vegetation. PLoS ONE, 2012, 7, e47500.	2.5	93
51	Specialisation and diversity of multiple trophic groups are promoted by different forest features. Ecology Letters, 2019, 22, 170-180.	6.4	92
52	Molecular evidence strongly supports deadwood-inhabiting fungi exhibiting unexpected tree species preferences in temperate forests. ISME Journal, 2018, 12, 289-295.	9.8	90
53	Community structure of arbuscular mycorrhizal fungi associated to Veronica rechingeri at the Anguran zinc and lead mining region. Environmental Pollution, 2008, 156, 1277-1283.	7.5	89
54	Laccases: toward disentangling their diversity and functions in relation to soil organic matter cycling. Biology and Fertility of Soils, 2010, 46, 215-225.	4.3	87

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55	Combining nested PCR and restriction digest of the internal transcribed spacer region to characterize arbuscular mycorrhizal fungi on roots from the field. Mycorrhiza, 2003, 13, 191-198.	2.8	86
56	Specific bottomâ€"up effects of arbuscular mycorrhizal fungi across a plantâ€"herbivoreâ€"parasitoid system. Oecologia, 2009, 160, 267-277.	2.0	86
57	Investigating the consequences of climate change under different landâ€use regimes: a novel experimental infrastructure. Ecosphere, 2019, 10, e02635.	2.2	85
58	Divergent habitat filtering of root and soil fungal communities in temperate beech forests. Scientific Reports, 2016, 6, 31439.	3.3	84
59	Characterization and identification of black alder ectomycorrhizas by PCR/RFLP analyses of the rDNA internal transcribed spacer (ITS). New Phytologist, 1997, 137, 357-369.	7.3	82
60	Taxonomic and phylogenetic contributions to fungi associated with the invasive weed Chromolaena odorata (Siam weed). Fungal Diversity, 2020, 101, 1-175.	12.3	82
61	Contrasting responses of above- and belowground diversity to multiple components of land-use intensity. Nature Communications, 2021, 12, 3918.	12.8	81
62	Shifts in physiological capabilities of the microbiota during the decomposition of leaf litter in a black alder (Alnus glutinosa (Gaertn.) L.) forest. Soil Biology and Biochemistry, 2001, 33, 921-930.	8.8	80
63	Functionally and phylogenetically diverse plant communities key to soil biota. Ecology, 2013, 94, 1878-1885.	3.2	80
64	Soil and tree species traits both shape soil microbial communities during early growth of Chinese subtropical forests. Soil Biology and Biochemistry, 2016, 96, 180-190.	8.8	80
65	Towards a universally adaptable method for quantitative extraction of high-purity nucleic acids from soil. Journal of Microbiological Methods, 2008, 75, 19-24.	1.6	77
66	Insights into organohalide respiration and the versatile catabolism of <scp><i>S</i></scp><i>ulfurospirillum multivorans</i> gained from comparative genomics and physiological studies. Environmental Microbiology, 2014, 16, 3562-3580.	3.8	76
67	Grassland management intensification weakens the associations among the diversities of multiple plant and animal taxa. Ecology, 2015, 96, 1492-1501.	3.2	75
68	Oribatid Mites as Potential Vectors for Soil Microfungi: Study of Mite-Associated Fungal Species. Microbial Ecology, 2005, 50, 518-528.	2.8	74
69	Impacts of earthworms and arbuscular mycorrhizal fungi (Glomus intraradices) on plant performance are not interrelated. Soil Biology and Biochemistry, 2009, 41, 561-567.	8.8	74
70	Impact of the plant community composition on labile soil organic carbon, soil microbial activity and community structure in semi-natural grassland ecosystems of different productivity. Plant and Soil, 2012, 352, 253-265.	3.7	74
71	Fine Spatial Scale Variation of Soil Microbial Communities under European Beech and Norway Spruce. Frontiers in Microbiology, 2016, 7, 2067.	3.5	74
72	Genome sequences of two dehalogenation specialists<i>- Dehalococcoides mccartyi</i>strains BTF08 and DCMB5 enriched from the highly polluted Bitterfeld region. FEMS Microbiology Letters, 2013, 343, 101-104.	1.8	73

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73	DNA polymorphism in morels: PCR/RFLP analysis of the ribosomal DNA spacers and microsatellite-primed PCR. Mycological Research, 1996, 100, 63-71.	2.5	72
74	Species composition of arbuscular mycorrhizal fungi in two mountain meadows with differing management types and levels of plant biodiversity. Biology and Fertility of Soils, 2006, 42, 286-298.	4.3	72
75	TaqMan Real-Time PCR Assays To Assess Arbuscular Mycorrhizal Responses to Field Manipulation of Grassland Biodiversity: Effects of Soil Characteristics, Plant Species Richness, and Functional Traits. Applied and Environmental Microbiology, 2010, 76, 3765-3775.	3.1	72
76	A molecular method to evaluate basidiomycete laccase gene expression in forest soils. Geoderma, 2005, 128, 18-27.	5.1	71
77	Diversity of laccase-like multicopper oxidase genes in Morchellaceae: identification of genes potentially involved in extracellular activities related to plant litter decay. FEMS Microbiology Ecology, 2007, 61, 153-163.	2.7	65
78	Uncoupling of microbial community structure and function in decomposing litter across beech forest ecosystems in Central Europe. Scientific Reports, 2014, 4, 7014.	3.3	65
79	Influence of Different Forest System Management Practices on Leaf Litter Decomposition Rates, Nutrient Dynamics and the Activity of Ligninolytic Enzymes: A Case Study from Central European Forests. PLoS ONE, 2014, 9, e93700.	2.5	65
80	Characterization and spatial distribution of ectomycorrhizas colonizing aspen clones released in an experimental field. Mycorrhiza, 2004, 14, 295-306.	2.8	64
81	Spatial Distribution of Fungal Communities in an Arable Soil. PLoS ONE, 2016, 11, e0148130.	2.5	63
82	Land-Use Intensity Rather Than Plant Functional Identity Shapes Bacterial and Fungal Rhizosphere Communities. Frontiers in Microbiology, 2018, 9, 2711.	3.5	62
83	Mycorrhizal colonization of transgenic aspen in a field trial. Planta, 2002, 214, 653-660.	3.2	61
84	Manipulation of the onset of ectomycorrhiza formation by indole-3-acetic acid, activated charcoal or relative humidity in the association between oak microcuttings and Piloderma croceum: influence on plant development and photosynthesis. Journal of Plant Physiology, 2004, 161, 509-517.	3.5	61
85	Changes within a single land-use category alter microbial diversity and community structure: Molecular evidence from wood-inhabiting fungi in forest ecosystems. Journal of Environmental Management, 2014, 139, 109-119.	7.8	61
86	Diversity of and yeasts (Basidiomycota) inhabiting arbuscular mycorrhizal roots or spores. FEMS Yeast Research, 2004, 4, 597-603.	2.3	60
87	Effects of resource availability and quality on the structure of the micro-food web of an arable soil across depth. Soil Biology and Biochemistry, 2012, 50, 1-11.	8.8	60
88	Diversity of arbuscular mycorrhizal fungi in grassland spontaneously developed on area polluted by a fertilizer plant. Environmental Pollution, 2005, 135, 255-266.	7.5	57
89	Soil nematodes associated with the mammal pathogenic fungal genus Malassezia (Basidiomycota:) Tj ETQq1 1 0.784314 rgBT /Overlo	4.3	56
90	Effects of Forest Management Practices in Temperate Beech Forests on Bacterial and Fungal Communities Involved in Leaf Litter Degradation. Microbial Ecology, 2015, 69, 905-913.	2.8	56

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91	Molecular fungal community and its decomposition activity in sapwood and heartwood of 13 temperate European tree species. PLoS ONE, 2019, 14, e0212120.	2.5	55
92	Forest Age and Plant Species Composition Determine the Soil Fungal Community Composition in a Chinese Subtropical Forest. PLoS ONE, 2013, 8, e66829.	2.5	53
93	Short-term bioavailability of carbon in soil organic matter fractions of different particle sizes and densities in grassland ecosystems. Science of the Total Environment, 2014, 497-498, 29-37.	8.0	53
94	A gnotobiotic culture system with oak microcuttings to study specific effects of mycobionts on plant morphology before, and in the early phase of, ectomycorrhiza formation by Paxillus involutus and Piloderma croceum. New Phytologist, 1998, 138, 203-212.	7.3	52
95	Metal uptake and detoxification mechanisms in Erica andevalensis growing in a pyrite mine tailing. Environmental and Experimental Botany, 2007, 61, 117-123.	4.2	52
96	Arbuscular mycorrhizal abundance in contaminated soils around a zinc and lead deposit. European Journal of Soil Biology, 2008, 44, 381-391.	3.2	51
97	Temporal changes in diversity and expression patterns of fungal laccase genes within the organic horizon of a brown forest soil. Soil Biology and Biochemistry, 2009, 41, 1380-1389.	8.8	51
98	Growing Research Networks on Mycorrhizae for Mutual Benefits. Trends in Plant Science, 2018, 23, 975-984.	8.8	51
99	The arbuscular mycorrhizal <i>Paraglomus majewskii</i> sp. nov. represents a distinct basal lineage in Glomeromycota. Mycologia, 2012, 104, 148-156.	1.9	50
100	The oak gene expression atlas: insights into Fagaceae genome evolution and the discovery of genes regulated during bud dormancy release. BMC Genomics, 2015, 16, 112.	2.8	49
101	Mycorrhiza in tree diversityâ€“ecosystem function relationships: conceptual framework and experimental implementation. Ecosphere, 2018, 9, e02226.	2.2	49
102	Are correlations between deadwood fungal community structure, wood physico-chemical properties and lignin-modifying enzymes stable across different geographical regions?. Fungal Ecology, 2016, 22, 98-105.	1.6	47
103	Rationalizing molecular analysis of field-collected roots for assessing diversity of arbuscular mycorrhizal fungi: to pool, or not to pool, that is the question. Mycorrhiza, 2006, 16, 525-531.	2.8	46
104	Implication of evolution and diversity in arbuscular and ectomycorrhizal symbioses. Journal of Plant Physiology, 2015, 172, 55-61.	3.5	46
105	Correlations between the composition of modular fungal communities and litter decomposition-associated ecosystem functions. Fungal Ecology, 2016, 22, 106-114.	1.6	46
106	Bacteria inhabiting deadwood of 13 tree species are heterogeneously distributed between sapwood and heartwood. Environmental Microbiology, 2018, 20, 3744-3756.	3.8	44
107	Morphological and anatomical characterisation of black alder Alnus glutinosa (L.) Gaertn. ectomycorrhizas. Mycorrhiza, 1997, 7, 201-216.	2.8	43
108	Determinants of Deadwood-Inhabiting Fungal Communities in Temperate Forests: Molecular Evidence From a Large Scale Deadwood Decomposition Experiment. Frontiers in Microbiology, 2018, 9, 2120.	3.5	43

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109	Relationships Between Soil Microorganisms, Plant Communities, and Soil Characteristics in Chinese Subtropical Forests. Ecosystems, 2012, 15, 624-636.	3.4	42
110	Dynamics of Soil Bacterial Communities Over a Vegetation Season Relate to Both Soil Nutrient Status and Plant Growth Phenology. Microbial Ecology, 2018, 75, 216-227.	2.8	42
111	Toward a methodical framework for comprehensively assessing forest multifunctionality. Ecology and Evolution, 2017, 7, 10652-10674.	1.9	41
112	Above- and belowground biodiversity jointly tighten the P cycle in agricultural grasslands. Nature Communications, 2021, 12, 4431.	12.8	40
113	Detection and quantification of a mycorrhization helper bacterium and a mycorrhizal fungus in plant-soil microcosms at different levels of complexity. BMC Microbiology, 2013, 13, 205.	3.3	39
114	Belowground top-down and aboveground bottom-up effects structure multitrophic community relationships in a biodiverse forest. Scientific Reports, 2017, 7, 4222.	3.3	38
115	Can multiâ€taxa diversity in European beech forest landscapes be increased by combining different management systems?. Journal of Applied Ecology, 2020, 57, 1363-1375.	4.0	38
116	Land use and host neighbor identity effects on arbuscular mycorrhizal fungal community composition in focal plant rhizosphere. Biodiversity and Conservation, 2013, 22, 2193-2205.	2.6	37
117	Multitrophic diversity in a biodiverse forest is highly nonlinear across spatial scales. Nature Communications, 2015, 6, 10169.	12.8	37
118	Carbon input and crop-related changes in microbial biomarker levels strongly affect the turnover and composition of soil organic carbon. Soil Biology and Biochemistry, 2015, 85, 39-50.	8.8	37
119	Leaf litter diversity alters microbial activity, microbial abundances, and nutrient cycling in a subtropical forest ecosystem. Biogeochemistry, 2017, 134, 163-181.	3.5	36
120	Preservation of nucleic acids by freeze-drying for next generation sequencing analyses of soil microbial communities. Journal of Plant Ecology, 2017, 10, 81-90.	2.3	36
121	Host Phylogeny Is a Major Determinant of Fagaceae-Associated Ectomycorrhizal Fungal Community Assembly at a Regional Scale. Frontiers in Microbiology, 2018, 9, 2409.	3.5	36
122	Wood decomposition is more strongly controlled by temperature than by tree species and decomposer diversity in highly species rich subtropical forests. Oikos, 2019, 128, 701-715.	2.7	36
123	Interactions between soil properties, agricultural management and cultivar type drive structural and functional adaptations of the wheat rhizosphere microbiome to drought. Environmental Microbiology, 2021, 23, 5866-5882.	3.8	36
124	Characterization of Unexplored Deadwood Mycobiome in Highly Diverse Subtropical Forests Using Culture-independent Molecular Technique. Frontiers in Microbiology, 2017, 8, 574.	3.5	35
125	The association of Morchella rotunda (Pers.) Boudier with roots of Picea abies (L.) Karst. New Phytologist, 1990, 116, 425-430.	7.3	33
126	Interactive effects of mycorrhizae and a root hemiparasite on plant community productivity and diversity. Oecologia, 2009, 159, 191-205.	2.0	33

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127	Tree species richness and fungi in freshly fallen leaf litter: Unique patterns of fungal species composition and their implications for enzymatic decomposition. Soil Biology and Biochemistry, 2018, 127, 120-126.	8.8	33
128	The Dark Side of Animal Phenology. Trends in Ecology and Evolution, 2018, 33, 898-901.	8.7	33
129	Association between living roots and ascocarps of Morchella rotunda. Transactions of the British Mycological Society, 1987, 89, 249-252.	0.6	32
130	Resource Type and Availability Regulate Fungal Communities Along Arable Soil Profiles. Microbial Ecology, 2015, 70, 390-399.	2.8	32
131	Largeâ€scale drivers of relationships between soil microbial properties and organic carbon across Europe. Global Ecology and Biogeography, 2021, 30, 2070-2083.	5.8	32
132	Back to the Future: Decomposability of a Biobased and Biodegradable Plastic in Field Soil Environments and Its Microbiome under Ambient and Future Climates. Environmental Science & Technology, 2021, 55, 12337-12351.	10.0	32
133	Identification of premycorrhizaâ€related plant genes in the association between Quercus robur and Piloderma croceum. New Phytologist, 2004, 163, 149-157.	7.3	31
134	What Are Soils?. , 2005, , 3-17.		31
135	Glomus drummondii and G. walkeri, two new species of arbuscular mycorrhizal fungi (Glomeromycota). Mycological Research, 2006, 110, 555-566.	2.5	31
136	Glomus africanumandG. iranicum, two new species of arbuscular mycorrhizal fungi (Glomeromycota). Mycologia, 2010, 102, 1450-1462.	1.9	31
137	Arbuscular mycorrhizas in phosphate-polluted soil: interrelations between root colonization and nitrogen. Plant and Soil, 2011, 343, 379-392.	3.7	31
138	Comparing fungal richness and community composition in coarse woody debris in Central European beech forests under three types of management. Mycological Progress, 2014, 13, 959-964.	1.4	31
139	Fungal guilds and soil functionality respond to tree community traits rather than to tree diversity in European forests. Molecular Ecology, 2021, 30, 572-591.	3.9	31
140	Cross talks at the morphogenetic, physiological and gene regulation levels between the mycobiont Piloderma croceum and oak microcuttings (Quercus robur) during formation of ectomycorrhizas. Phytochemistry, 2007, 68, 52-67.	2.9	29
141	Diversity of the ectomycorrhiza community at a uranium mining heap. Biology and Fertility of Soils, 2005, 41, 439-446.	4.3	28
142	Transcriptional changes in two types of pre-mycorrhizal roots and in ectomycorrhizas of oak microcuttings inoculated with Piloderma croceum. Planta, 2006, 225, 331-340.	3.2	28
143	Large scale transcriptome analysis reveals interplay between development of forest trees and a beneficial mycorrhiza helper bacterium. BMC Genomics, 2015, 16, 658.	2.8	28
144	Archaeal Diversity and CO₂Fixers in Carbonate-/Siliciclastic-Rock Groundwater Ecosystems. Archaea, 2017, 2017, 1-13.	2.3	28

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145	Experimental Evidence of Functional Group-Dependent Effects of Tree Diversity on Soil Fungi in Subtropical Forests. Frontiers in Microbiology, 2018, 9, 2312.	3.5	28
146	Ectomycorrhizal types and endobacteria associated with ectomycorrhizas of Morchella elata (Fr.) Boudier with Picea abies (L.) Karst. Mycorrhiza, 1994, 4, 223-232.	2.8	27
147	Characteristics and energetic strategies of the rhizosphere in ecosystems of the BornhÃ¶ved Lake district. Applied Soil Ecology, 2000, 15, 201-210.	4.3	27
148	Actinobacteria may influence white truffle (Tuber magnatum Pico) nutrition, ascocarp degradation and interactions with other soil fungi. Fungal Ecology, 2013, 6, 527-538.	1.6	27
149	Septoglomus fuscum and S. furcatum, two new species of arbuscular mycorrhizal fungi (Glomeromycota). Mycologia, 2013, 105, 670-680.	1.9	27
150	Endogenous rhythmic growth in oak trees is regulated by internal clocks rather than resource availability. Journal of Experimental Botany, 2015, 66, 7113-7127.	4.8	27
151	Tree species, tree genotypes and tree genotypic diversity levels affect microbe-mediated soil ecosystem functions in a subtropical forest. Scientific Reports, 2016, 6, 36672.	3.3	27
152	Unraveling spatiotemporal variability of arbuscular mycorrhizal fungi in a temperate grassland plot. Environmental Microbiology, 2020, 22, 873-888.	3.8	27
153	Synthesis of two types of association between Morchella esculenta and Picea abies under controlled culture conditions. Journal of Plant Physiology, 1993, 141, 12-17.	3.5	26
154	A molecular phylogeny of Hebeloma species from Europe. Mycological Research, 2006, 110, 369-380.	2.5	26
155	<i>Glomus indicum,</i> a new arbuscular mycorrhizal fungus. Botany, 2010, 88, 132-143.	1.0	26
156	Contrasting effects of grassland management modes on species-abundance distributions of multiple groups. Agriculture, Ecosystems and Environment, 2017, 237, 143-153.	5.3	26
157	Increasing N deposition impacts neither diversity nor functions of deadwoodâ€inhabiting fungal communities, but adaptation and functional redundancy ensure ecosystem function. Environmental Microbiology, 2018, 20, 1693-1710.	3.8	26
158	First insights into the living groundwater mycobiome of the terrestrial biogeosphere. Water Research, 2018, 145, 50-61.	11.3	26
159	Shifts Between and Among Populations of Wheat Rhizosphere Pseudomonas, Streptomyces and Phyllobacterium Suggest Consistent Phosphate Mobilization at Different Wheat Growth Stages Under Abiotic Stress. Frontiers in Microbiology, 2019, 10, 3109.	3.5	25
160	Species diversity within the group (Ascomycota: Morchellaceae) in Germany and France. Organisms Diversity and Evolution, 2005, 5, 101-107.	1.6	24
161	Superimposed Pristine Limestone Aquifers with Marked Hydrochemical Differences Exhibit Distinct Fungal Communities. Frontiers in Microbiology, 2016, 7, 666.	3.5	24
162	Transcriptome analysis in oak uncovers a strong impact of endogenous rhythmic growth on the interaction with plant-parasitic nematodes. BMC Genomics, 2016, 17, 627.	2.8	24

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163	Home-Field Advantage in Wood Decomposition Is Mainly Mediated by Fungal Community Shifts at â€œHomeâ€•Versus â€œAwayâ€• Microbial Ecology, 2019, 78, 725-736.	2.8	24
164	Organic agricultural practice enhances arbuscular mycorrhizal symbiosis in correspondence to soil warming and altered precipitation patterns. Environmental Microbiology, 2021, 23, 6163-6176.	3.8	24
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