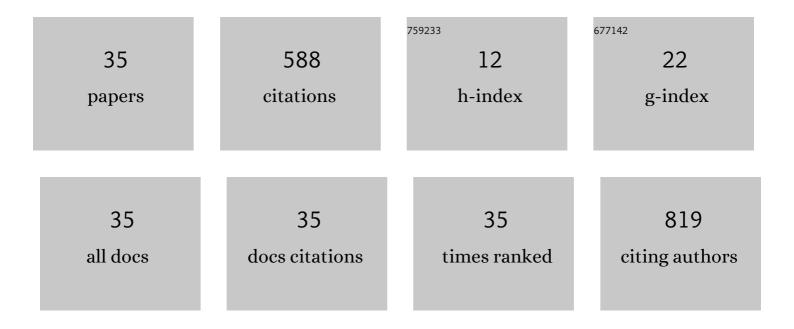
## Dong Liu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4440657/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Soil physicochemical and microbial characteristics of contrasting land-use types along soil depth gradients. Catena, 2018, 162, 345-353.	5.0	67
2	The restoration age of Robinia pseudoacacia plantation impacts soil microbial biomass and microbial community structure in the Loess Plateau. Catena, 2018, 165, 192-200.	5.0	56
3	Microbial functionality as affected by experimental warming of a temperate mountain forest soil—A metaproteomics survey. Applied Soil Ecology, 2017, 117-118, 196-202.	4.3	48
4	Effects of Revegetation on Soil Microbial Biomass, Enzyme Activities, and Nutrient Cycling on the Loess Plateau in China. Restoration Ecology, 2013, 21, 600-607.	2.9	38
5	Geographic distance and soil microbial biomass carbon drive biogeographical distribution of fungal communities in Chinese Loess Plateau soils. Science of the Total Environment, 2019, 660, 1058-1069.	8.0	36
6	The Biogeographical Distribution of Soil Bacterial Communities in the Loess Plateau as Revealed by High-Throughput Sequencing. Frontiers in Microbiology, 2018, 9, 2456.	3.5	35
7	Response of Microbial Communities and Their Metabolic Functions to Drying–Rewetting Stress in a Temperate Forest Soil. Microorganisms, 2019, 7, 129.	3.6	35
8	Is there a convergence of deciduous leaf litter stoichiometry, biochemistry and microbial population during decay?. Geoderma, 2016, 272, 93-100.	5.1	33
9	Effects of nitrogen addition on rhizospheric soil microbial communities of poplar plantations at different ages. Forest Ecology and Management, 2021, 494, 119328.	3.2	28
10	Decoupled diversity patterns in microbial geographic distributions on the arid area (the Loess) Tj ETQq0 0 0 rgBT	/Overlock	10 Tf 50 38
	Relationship between maturity and microbial communities during pig manure composting by		

11	Relationship between maturity and microbial communities during pig manure composting by phospholipid fatty acid (PLFA) and correlation analysis. Journal of Environmental Management, 2018, 206, 532-539.	7.8	20
12	Passive and active ecological restoration strategies for abandoned farmland leads to shifts in potential soil nitrogen loss by denitrification and soil denitrifying microbes. Land Degradation and Development, 2020, 31, 1086-1098.	3.9	20
13	Diversity patterns and drivers of soil bacterial and fungal communities along elevational gradients in the Southern Himalayas, China. Applied Soil Ecology, 2022, 178, 104563.	4.3	16
14	Long-Term Nitrogen Deposition Alters Ectomycorrhizal Community Composition and Function in a Poplar Plantation. Journal of Fungi (Basel, Switzerland), 2021, 7, 791.	3.5	15
15	Variability in Soil Microbial Biomass and Diversity Among Different Aggregate-Size Fractions of Different Land Use Types. Soil Science, 2014, 179, 242-249.	0.9	13
16	Dynamics of soil nitrogen fractions and their relationship with soil microbial communities in two forest species of northern China. PLoS ONE, 2018, 13, e0196567.	2.5	12
17	Circumscription and phylogeny of the Lepidostromatales ( <i>Lichenized Basidiomycota</i> ) following discovery of new species from China and Africa. Mycologia, 2017, 109, 730-748.	1.9	10
18	Amino acid substitutions in antigenic region B of hemagglutinin play a critical role in the antigenic drift of subclade 2.3.4.4 highly pathogenic H5NX influenza viruses. Transboundary and Emerging Diseases, 2020, 67, 263-275.	3.0	9

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19	Microbiome Community Structure and Functional Gene Partitioning in Different Micro-Niches Within a Sporocarp-Forming Fungus. Frontiers in Microbiology, 2021, 12, 629352.	3.5	9
20	Tuber pseudohimalayense ascomata-compartments strongly select their associated bacterial microbiome from nearby pine forest soils independently of their maturation stage. Pedobiologia, 2021, 87-88, 150743.	1.2	9
21	Taxonomic study of the genus <i>Anzia</i> ( <i>Lecanorales</i> , lichenized Ascomycota) from Hengduan Mountains, China. Lichenologist, 2015, 47, 99-115.	0.8	8
22	Macrofungi Cultivation in Shady Forest Areas Significantly Increases Microbiome Diversity, Abundance and Functional Capacity in Soil Furrows. Journal of Fungi (Basel, Switzerland), 2021, 7, 775.	3.5	7
23	Land rehabilitation improves edaphic conditions and increases soil microbial biomass and abundance. Soil Ecology Letters, 2020, 2, 145-156.	4.5	6
24	Anemochore Seeds Harbor Distinct Fungal and Bacterial Abundance, Composition, and Functional Profiles. Journal of Fungi (Basel, Switzerland), 2022, 8, 89.	3.5	6
25	Provenances originate morphological and microbiome variation of Tuber pseudobrumale in southwestern China despite strong genetic consistency. Mycological Progress, 2020, 19, 1545-1558.	1.4	5
26	New species and records of Pyxine (Caliciaceae) in China. MycoKeys, 2019, 45, 93-109.	1.9	5
27	Distinct Compartmentalization of Microbial Community and Potential Metabolic Function in the Fruiting Body of Tricholoma matsutake. Journal of Fungi (Basel, Switzerland), 2021, 7, 586.	3.5	4
28	Soil Rehabilitation Promotes Resilient Microbiome with Enriched Keystone Taxa than Agricultural Infestation in Barren Soils on the Loess Plateau. Biology, 2021, 10, 1261.	2.8	4
29	The Genus <i>Letrouitia</i> (Letrouitiaceae: Lichenized Ascomycota) New to Cambodia. Mycobiology, 2015, 43, 163-165.	1.7	2
30	The genus Bulbothrix (Parmeliaceae) in China. Lichenologist, 2016, 48, 121-133.	0.8	2
31	Three new species and one new combination of Gypsoplaca (lichenized Ascomycota) from the Hengduan Mountains in China. Mycological Progress, 2018, 17, 781-790.	1.4	2
32	A design of self-service speech explaining system based on RFID. , 2012, , .		1
33	New species and new records of Ophioparmaceae (lichenized Ascomycota) from China. Lichenologist, 2018, 50, 89-99.	0.8	1
34	Truffle species strongly shape their surrounding soil mycobiota in a Pinus armandii forest. Archives of Microbiology, 2021, 203, 6303-6314.	2.2	1
35	Taxonomic Study of Hypotrachyna Subg. Everniastrum (Hale Ex Sipman) Divakar, A.Crespo, Sipman, Elix & Lumbsch (Ascomycota) from China. Cryptogamie, Mycologie, 2020, 41, .	1.0	0