

# Anna Maria Fiore-Donno

## List of Publications by Year in descending order

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Version: 2024-02-01

44  
papers

2,677  
citations

236833

25  
h-index

254106

43  
g-index

50  
all docs

50  
docs citations

50  
times ranked

2848  
citing authors

#	ARTICLE	IF	CITATIONS
1	Soil compartments (bulk soil, litter, root and rhizosphere) as main drivers of soil protistan communities distribution in forests with different nitrogen deposition. <i>Soil Biology and Biochemistry</i> , 2022, 168, 108628.	4.2	19
2	Ecological clusters of soil taxa within bipartite networks are highly sensitive to climatic conditions in global drylands. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2022, 377, .	1.8	4
3	Different community compositions between obligate and facultative oomycete plant parasites in a landscape-scale metabarcoding survey. <i>Biology and Fertility of Soils</i> , 2021, 57, 245-256.	2.3	21
4	A Parasite's Paradise: Biotrophic Species Prevail Oomycete Community Composition in Tree Canopies. <i>Frontiers in Forests and Global Change</i> , 2021, 4, .	1.0	2
5	On the phenology of protists: recurrent patterns reveal seasonal variation of protistan (Rhizaria:) Tj ETQq1 1 0.784314 rgBT /Overlock 1.3 9	1.3	9
6	Contrasting responses of above- and belowground diversity to multiple components of land-use intensity. <i>Nature Communications</i> , 2021, 12, 3918.	5.8	81
7	Making sense of environmental sequencing data: Ecologically important functional traits of the protistan groups Cercozoa and Endomyxa (Rhizaria). <i>Molecular Ecology Resources</i> , 2020, 20, 398-403.	2.2	66
8	Phylogeny of Physarida (Amoebozoa, Myxogastria) Based on the Small Subunit Ribosomal RNA Gene, Redefinition of <i>Physarum pusillum</i> s. str. and Reinstatement of <i>P. Agravidum</i> Morgan. <i>Journal of Eukaryotic Microbiology</i> , 2020, 67, 327-336.	0.8	10
9	Land-use intensity alters networks between biodiversity, ecosystem functions, and services. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 28140-28149.	3.3	164
10	Contrasting Responses of Protistan Plant Parasites and Phagotrophs to Ecosystems, Land Management and Soil Properties. <i>Frontiers in Microbiology</i> , 2020, 11, 1823.	1.5	27
11	From Forest Soil to the Canopy: Increased Habitat Diversity Does Not Increase Species Richness of Cercozoa and Oomycota in Tree Canopies. <i>Frontiers in Microbiology</i> , 2020, 11, 592189.	1.5	7
12	Multitrophic interactions in the rhizosphere microbiome of wheat: from bacteria and fungi to protists. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	77
13	New insights into the phylogeny of the dark-spored Myxomycetes (Amoebozoa: Conosa: Myxogastria:) Tj ETQq1 1 0.784314 rgBT /Overlock 0.5 7	0.5	7
14	Functional Traits and Spatio-Temporal Structure of a Major Group of Soil Protists (Rhizaria:) Tj ETQq0 0 0 rgBT /Overlock 1.5 10 Tf 50 222 T	1.5	82
15	Metatranscriptomics reveals unsuspected protistan diversity in leaf litter across temperate beech forests, with Amoebozoa the dominating lineage. <i>FEMS Microbiology Ecology</i> , 2019, 95, .	1.3	23
16	Distinct communities of Cercozoa at different soil depths in a temperate agricultural field. <i>FEMS Microbiology Ecology</i> , 2019, 95, .	1.3	21
17	A Non-flagellated Member of the Myxogastria and Expansion of the Echinosteliida. <i>Journal of Eukaryotic Microbiology</i> , 2019, 66, 538-544.	0.8	19
18	The Protists in Soil: A Token of Untold Eukaryotic Diversity. , 2019, , 125-140.		15

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19	New barcoded primers for efficient retrieval of cercozoan sequences in high-throughput environmental diversity surveys, with emphasis on worldwide biological soil crusts. <i>Molecular Ecology Resources</i> , 2018, 18, 229-239.	2.2	71
20	Protists are an integral part of the <i>Arabidopsis thaliana</i> microbiome. <i>Environmental Microbiology</i> , 2018, 20, 30-43.	1.8	85
21	Soil protistology rebooted: 30 fundamental questions to start with. <i>Soil Biology and Biochemistry</i> , 2017, 111, 94-103.	4.2	130
22	Inferring interactions in complex microbial communities from nucleotide sequence data and environmental parameters. <i>PLoS ONE</i> , 2017, 12, e0173765.	1.1	15
23	Phylogeny of the Highly Divergent Echinosteliales (Amoebozoa). <i>Journal of Eukaryotic Microbiology</i> , 2016, 63, 453-459.	0.8	19
24	Metacommunity analysis of amoeboid protists in grassland soils. <i>Scientific Reports</i> , 2016, 6, 19068.	1.6	82
25	Expansion of the molecular and morphological diversity of Acanthamoebidae (Centramoebida.) <i>Tj ETQq1 1 0.784314 rgBT / Overlock 10 Tf 50 3</i>	1.9	58
26	First insight into dead wood protistan diversity: a molecular sampling of bright-spored Myxomycetes (Amoebozoa, slime-moulds) in decaying beech logs. <i>FEMS Microbiology Ecology</i> , 2015, 91, .	1.3	23
27	Multigene phylogeny resolves deep branching of Amoebozoa. <i>Molecular Phylogenetics and Evolution</i> , 2015, 83, 293-304.	1.2	84
28	Using environmental niche models to test the "everything is everywhere"™ hypothesis for <i>Badhamia</i> . <i>ISME Journal</i> , 2014, 8, 737-745.	4.4	55
29	Acanthamoeba everywhere: high diversity of Acanthamoeba in soils. <i>Parasitology Research</i> , 2014, 113, 3151-3158.	0.6	75
30	Multigene eukaryote phylogeny reveals the likely protozoan ancestors of opisthokonts (animals.) <i>Tj ETQq0 0 0 rgBT / Overlock 10 Tf 50 3</i>	1.2	97
31	Phylogenetic position of the enigmatic myxomycete genus <i>Kelleromyxa</i> revealed by SSU rDNA sequences. <i>Mycological Progress</i> , 2013, 12, 599-608.	0.5	24
32	Exploring slime mould diversity in high-altitude forests and grasslands by environmental RNA analysis. <i>FEMS Microbiology Ecology</i> , 2013, 84, 98-109.	1.3	34
33	Two-Gene Phylogeny of Bright-Spored Myxomycetes (Slime Moulds, Superorder Lucisporidia). <i>PLoS ONE</i> , 2013, 8, e62586.	1.1	58
34	CBOL Protist Working Group: Barcoding Eukaryotic Richness beyond the Animal, Plant, and Fungal Kingdoms. <i>PLoS Biology</i> , 2012, 10, e1001419.	2.6	488
35	18S rDNA Phylogeny of Lamproderma and Allied Genera (Stemonitales, Myxomycetes, Amoebozoa). <i>PLoS ONE</i> , 2012, 7, e35359.	1.1	75
36	Genetic Structure of Two Protist Species (Myxogastria, Amoebozoa) Suggests Asexual Reproduction in Sexual Amoebae. <i>PLoS ONE</i> , 2011, 6, e22872.	1.1	47

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37	Myxomycetes in soil. <i>Soil Biology and Biochemistry</i> , 2011, 43, 2237-2242.	4.2	52
38	Deep Phylogeny and Evolution of Slime Moulds (Mycetozoa). <i>Protist</i> , 2010, 161, 55-70.	0.6	122
39	Invalidation of <i>Hyperamoeba</i> by Transferring its Species to Other Genera of Myxogastria. <i>Journal of Eukaryotic Microbiology</i> , 2010, 57, 189-196.	0.8	25
40	Ecology of sandstone ravine myxomycetes from Saxonian Switzerland (Germany). <i>Nova Hedwigia</i> , 2010, 90, 277-302.	0.2	15
41	<i>Semimorula liquescens</i> is a modified echinostelid myxomycete (Mycetozoa). <i>Mycologia</i> , 2009, 101, 773-776.	0.8	20
42	Evolution of dark-spored Myxomycetes (slime-molds): Molecules versus morphology. <i>Molecular Phylogenetics and Evolution</i> , 2008, 46, 878-889.	1.2	96
43	Higher Order Phylogeny of Plasmodial Slime Molds (Myxogastria) Based on Elongation Factor 1 and Small Subunit rRNA Gene Sequences. <i>Journal of Eukaryotic Microbiology</i> , 2005, 52, 201-210.	0.8	84
44	Populations of ectomycorrhizal <i>Laccaria amethystina</i> and <i>Xerocomus</i> spp. show contrasting colonization patterns in a mixed forest. <i>New Phytologist</i> , 2001, 152, 533-542.	3.5	85