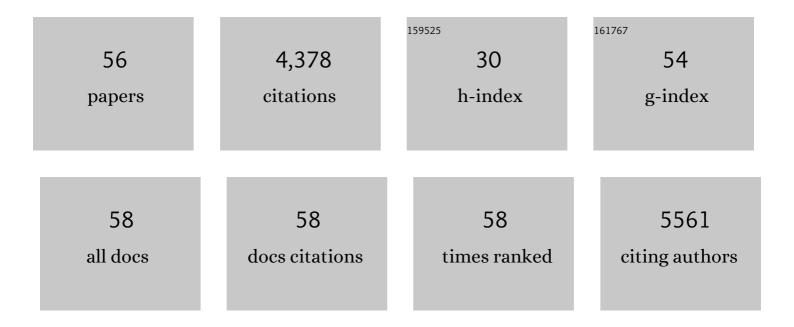
Maria Teresa Ceccherini

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Microbial diversity and soil functions. European Journal of Soil Science, 2003, 54, 655-670.	1.8	1,496
2	Extracellular DNA in soil and sediment: fate and ecological relevance. Biology and Fertility of Soils, 2009, 45, 219-235.	2.3	408
3	Distribution of microbial communities in a forest soil profile investigated by microbial biomass, soil respiration and DGGE of total and extracellular DNA. Soil Biology and Biochemistry, 2004, 36, 859-868.	4.2	272
4	Microbial diversity and soil functions. European Journal of Soil Science, 2017, 68, 12-26.	1.8	268
5	Direct molecular biological analysis of ammonia oxidising bacteria populations in cultivated soil plots treated with swine manure. FEMS Microbiology Ecology, 2006, 23, 45-54.	1.3	105
6	Hydrolase activity, microbial biomass and community structure in long-term Cd-contaminated soils. Soil Biology and Biochemistry, 2004, 36, 443-451.	4.2	100
7	Degradation and Transformability of DNA from Transgenic Leaves. Applied and Environmental Microbiology, 2003, 69, 673-678.	1.4	92
8	Beyond microbial diversity for predicting soil functions: A mini review. Pedosphere, 2020, 30, 5-17.	2.1	85
9	A simplified rapid, low-cost and versatile DNA-based assessment of soil microbial biomass. Ecological Indicators, 2014, 45, 75-82.	2.6	79
10	Fate and transport of antibiotic resistance genes in saturated soil columns. European Journal of Soil Biology, 2003, 39, 65-71.	1.4	75
11	Sequential extraction and genetic fingerprinting of a forest soil metagenome. Applied Soil Ecology, 2009, 42, 176-181.	2.1	74
12	Effect of Mediterranean Diet Enriched in High Quality Extra Virgin Olive Oil on Oxidative Stress, Inflammation and Gut Microbiota in Obese and Normal Weight Adult Subjects. Frontiers in Pharmacology, 2019, 10, 1366.	1.6	72
13	Soil Pollution from Micro- and Nanoplastic Debris: A Hidden and Unknown Biohazard. Sustainability, 2020, 12, 7255.	1.6	70
14	Long-term effects of aided phytostabilisation of trace elements on microbial biomass and activity, enzyme activities, and composition of microbial community in the Jales contaminated mine spoils. Environmental Pollution, 2008, 152, 702-712.	3.7	66
15	Adsorption of pure and dirty bacterial DNA on clay minerals and their transformation frequency. Biology and Fertility of Soils, 2007, 43, 731-739.	2.3	63
16	Composition, biomass and activity of microflora, and leaf yields and foliar elemental concentrations of lettuce, after in situ stabilization of an arsenic-contaminated soil. Applied Soil Ecology, 2009, 41, 351-359.	2.1	63
17	Evaluation of the Performances of Ribosomal Database Project (RDP) Classifier for Taxonomic Assignment of 16S rRNA Metabarcoding Sequences Generated from Illumina-Solexa NGS. Journal of Genomics, 2015, 3, 36-39.	0.6	59
18	Are humus forms, mesofauna and microflora in subalpine forest soils sensitive to thermal conditions?. Biology and Fertility of Soils, 2012, 48, 709-725.	2.3	57

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19	Amino Acid: Its Dual Role as Nutrient and Scavenger of Free Radicals in Soil. Sustainability, 2017, 9, 1402.	1.6	55
20	Phosphomonoesterase production and persistence and composition of bacterial communities during plant material decomposition in soils with different pH values. Soil Biology and Biochemistry, 2006, 38, 795-802.	4.2	54
21	Microbial community development and unseen diversity recovery in inoculated sterile soil. Biology and Fertility of Soils, 2014, 50, 1069-1076.	2.3	53
22	Protease encoding microbial communities and protease activity of the rhizosphere and bulk soils of two maize lines with different N uptake efficiency. Soil Biology and Biochemistry, 2016, 96, 176-179.	4.2	49
23	Seasonal variation and distribution of total and active microbial community of β-glucosidase encoding genes in coniferous forest soil. Soil Biology and Biochemistry, 2017, 105, 71-80.	4.2	46
24	Assessment of some cultural experimental methods to study the effects of antibiotics on microbial activities in a soil: An incubation study. PLoS ONE, 2017, 12, e0180663.	1.1	44
25	Maize lines with different nitrogen use efficiency select bacterial communities with different β-glucosidase-encoding genes and glucosidase activity in the rhizosphere. Biology and Fertility of Soils, 2015, 51, 995-1004.	2.3	40
26	Impact of chlortetracycline and sulfapyridine antibiotics on soil enzyme activities. International Agrophysics, 2017, 31, 499-505.	0.7	38
27	Physico-chemical and microbiological evidence of exposure effects on Picea abies – Coarse woody debris at different stages of decay. Forest Ecology and Management, 2017, 391, 376-389.	1.4	37
28	Enzyme activity and microbial community structure in the rhizosphere of two maize lines differing in N use efficiency. Plant and Soil, 2015, 387, 413-424.	1.8	36
29	Experimental discrimination and molecular characterization of the extracellular soil DNA fraction. Antonie Van Leeuwenhoek, 2009, 96, 653-657.	0.7	35
30	Purification and isotopic signatures (δ13C, δ15N, Δ14C) of soil extracellular DNA. Biology and Fertility of Soils, 2007, 44, 353-361.	2.3	33
31	Vertical advection of extracellular DNA by water capillarity in soil columns. Soil Biology and Biochemistry, 2007, 39, 158-163.	4.2	31
32	Effects of swine manure fertilization on autotrophic ammonia oxidizing bacteria in soil. Applied Soil Ecology, 1998, 7, 149-157.	2.1	26
33	Microbial dynamics in Mediterranean Moder humus. Biology and Fertility of Soils, 2012, 48, 259-270.	2.3	26
34	Rhizosphere as Hotspot for Plant-Soil-Microbe Interaction. , 2020, , 17-43.		26
35	Phylogeny of the genus based on 16S rDNA sequence. FEMS Microbiology Letters, 1995, 129, 195-200.	0.7	19
36	Leaching and transformability of transgenic DNA in unsaturated soil columns. Ecotoxicology and Environmental Safety, 2010, 73, 67-72.	2.9	19

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37	Nannipieri, P., Ascher, J., Ceccherini, M.T., Landi, L., Pietramellara, G. & Renella, G. 2003. Microbial diversity and soil functions. <i>European Journal of Soil Science</i> , 54, 655–670 European Journal of Soil Science, 2017, 68, 2-5.	1.8	19
38	The effect of pharmaceutical waste-fungal biomass, treated to degrade DNA, on the composition of eubacterial and ammonia oxidizing populations of soil. Biology and Fertility of Soils, 2007, 44, 299-306.	2.3	18
39	Use of random amplified polymorphic DNA markers for the detection of Azospirillum strains in soil microcosms. Applied Microbiology and Biotechnology, 1998, 49, 221-225.	1.7	15
40	Evaluation of the denaturing gradient gel electrophoresis-apparatus as a parameter influencing soil microbial community fingerprinting. World Journal of Microbiology and Biotechnology, 2010, 26, 1721-1726.	1.7	15
41	Cattle impact on composition of archaeal, bacterial, and fungal communities by comparative fingerprinting of total and extracellular DNA. Biology and Fertility of Soils, 2013, 49, 351-361.	2.3	15
42	Immediate- and Short-term Wildfire Impact on Soil Microbial Diversity and Activity in a Mediterranean Forest Soil. Soil Science, 2019, 184, 35-42.	0.9	15
43	Long-term persistence and bacterial transformation potential of transplastomic plant DNA in soil. Research in Microbiology, 2010, 161, 326-334.	1.0	12
44	Physical protection of extracellular and intracellular DNA in soil aggregates against simulated natural oxidative processes. Applied Soil Ecology, 2021, 165, 104002.	2.1	12
45	Molecular discrimination of bacteria (organic versus mineral soil layers) of dry woodlands of Argentina. Journal of Arid Environments, 2012, 85, 18-26.	1.2	11
46	<i>Inâ€field</i> detection and quantification of extracellular DNA. Journal of Plant Nutrition and Soil Science, 2009, 172, 626-629.	1.1	9
47	Response of Soil Bacterial Community to Application of Organic and Inorganic Phosphate Based Fertilizers under Vicia faba L. Cultivation at Two Different Phenological Stages. Sustainability, 2020, 12, 9706.	1.6	9
48	Exploring the dynamics of bacterial community composition in soil: the pan-bacteriome approach. Antonie Van Leeuwenhoek, 2015, 107, 785-797.	0.7	8
49	Preliminary evidences of the presence of extracellular DNA single stranded forms in soil. PLoS ONE, 2020, 15, e0227296.	1.1	7
50	Persistence of transgenic and not transgenic extracellular DNA in soil and bacterial transformation. Theoretical Biology Forum, 2006, 99, 37-68.	0.2	7
51	Chemical and microbiological changes in Norway spruce deadwood during the early stage of decomposition as a function of exposure in an alpine setting. Arctic, Antarctic, and Alpine Research, 2018, 50, .	0.4	6
52	Soil microbiome biomass, activity, composition and <scp>CO₂</scp> emissions in a longâ€ŧerm organic and conventional farming systems. Soil Use and Management, 2023, 39, 588-605.	2.6	6
53	Upward movement of Verticillium dahliae from soil to olive plants detected by qPCR. World Journal of Microbiology and Biotechnology, 2013, 29, 1961-1967.	1.7	5
54	Oral Lactobacillus Species in Systemic Sclerosis. Microorganisms, 2021, 9, 1298.	1.6	4

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#	ARTICLE	IF	CITATIONS
55	The extracellular DNA can baffle the assessment of soil bacterial community, but the effect varies with microscale spatial distribution. FEMS Microbiology Letters, 2021, 368, .	0.7	4

56 Soil carbon dioxide emission flux from organic and conventional farming in a long term experiment in Tuscany. , 2019, , .