

Going back to the roots: the microbial ecology of the rhizosphere

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Syringic acid inhibited cucumber seedling growth and changed rhizosphere microbial communities. <i>Plant, Soil and Environment</i> , 2014, 60, 158-164.	1.0	18
2	Selective Microbial Genomic DNA Isolation Using Restriction Endonucleases. <i>PLoS ONE</i> , 2014, 9, e109061.	1.1	19
3	Managing biotic interactions for ecological intensification of agroecosystems. <i>Frontiers in Ecology and Evolution</i> , 2014, 2, .	1.1	42
4	Microbial dispersal in unsaturated porous media: Characteristics of motile bacterial cell motions in unsaturated angular pore networks. <i>Water Resources Research</i> , 2014, 50, 7406-7429.	1.7	73
5	Contribution of Arbuscular Mycorrhizal Fungi to Soil Carbon Sequestration. <i>Soil Biology</i> , 2014, , 287-296.	0.6	12
6	Mycorrhizal Fungi: Use in Sustainable Agriculture and Land Restoration. <i>Soil Biology</i> , 2014, , .	0.6	21
7	Interactions between <i>Bacillus anthracis</i> and Plants May Promote Anthrax Transmission. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e2903.	1.3	40
8	The rhizosphere microbiota of plant invaders: an overview of recent advances in the microbiomics of invasive plants. <i>Frontiers in Microbiology</i> , 2014, 5, 368.	1.5	145
9	Ecological dynamics and complex interactions of <i>Agrobacterium</i> megaplasmids. <i>Frontiers in Plant Science</i> , 2014, 5, 635.	1.7	36
10	Comparative Metagenomic Analysis of Human Gut Microbiome Composition Using Two Different Bioinformatic Pipelines. <i>BioMed Research International</i> , 2014, 2014, 1-10.	0.9	68
11	Differences between the rhizosphere microbiome of <i>Beta vulgaris</i> ssp. <i>maritima</i> ancestor of all beet crops and modern sugar beets. <i>Frontiers in Microbiology</i> , 2014, 5, 415.	1.5	124
12	Draft Genome Sequence of <i>Paenibacillus pini</i> JCM 16418 <sup>T</sup> , Isolated from the Rhizosphere of Pine Tree. <i>Genome Announcements</i> , 2014, 2, .	0.8	5
13	Role of 2-hexyl, 5-propyl resorcinol production by <i>Pseudomonas chlororaphis</i> PCL1606 in the multitrophic interactions in the avocado rhizosphere during the biocontrol process. <i>FEMS Microbiology Ecology</i> , 2014, 89, 20-31.	1.3	50
14	The <i>Sphagnum</i> microbiome supports bog ecosystem functioning under extreme conditions. <i>Molecular Ecology</i> , 2014, 23, 4498-4510.	2.0	98
15	Belowground biodiversity and ecosystem functioning. <i>Nature</i> , 2014, 515, 505-511.	13.7	2,371
16	Invasive plant species set up their own niche. <i>New Phytologist</i> , 2014, 204, 435-437.	3.5	4
17	Belowground opportunities in vegetation science. <i>Journal of Vegetation Science</i> , 2014, 25, 1117-1125.	1.1	30
18	An increasing opine carbon bias in artificial exudation systems and genetically modified plant rhizospheres leads to an increasing reshaping of bacterial populations. <i>Molecular Ecology</i> , 2014, 23, 4846-4861.	2.0	33

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19	An introduction to the analysis of shotgun metagenomic data. <i>Frontiers in Plant Science</i> , 2014, 5, 209.	1.7	446
20	Building the crops of tomorrow: advantages of symbiont-based approaches to improving abiotic stress tolerance. <i>Frontiers in Microbiology</i> , 2014, 5, 283.	1.5	196
21	Rhizosphere microbial communities associated with Rhizoctonia damage at the field and disease patch scale. <i>Applied Soil Ecology</i> , 2014, 78, 37-47.	2.1	42
22	Phyllosphere Microbiota Composition and Microbial Community Transplantation on Lettuce Plants Grown Indoors. <i>MBio</i> , 2014, 5, .	1.8	84
23	Multitrophic microbial interactions for eco- and agro-biotechnological processes: theory and practice. <i>Trends in Biotechnology</i> , 2014, 32, 529-537.	4.9	63
24	The pathogen <i>Batrachochytrium dendrobatidis</i> disturbs the frog skin microbiome during a natural epidemic and experimental infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E5049-58.	3.3	264
25	Soybean Plants Modify Metal Oxide Nanoparticle Effects on Soil Bacterial Communities. <i>Environmental Science &amp; Technology</i> , 2014, 48, 13489-13496.	4.6	99
26	Functional Soil Microbiome: Belowground Solutions to an Aboveground Problem. <i>Plant Physiology</i> , 2014, 166, 689-700.	2.3	299
27	How may biochar influence severity of diseases caused by soilborne pathogens?. <i>Carbon Management</i> , 2014, 5, 169-183.	1.2	117
28	Bacterial community characterization in the soils of native and restored rainforest fragments. <i>Antonie Van Leeuwenhoek</i> , 2014, 106, 947-957.	0.7	2
29	Impacts of vegetation, tidal process, and depth on the activities, abundances, and community compositions of denitrifiers in mangrove sediment. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 9375-9387.	1.7	45
30	Root exudates mediated interactions belowground. <i>Soil Biology and Biochemistry</i> , 2014, 77, 69-80.	4.2	671
31	Phytoextraction and dissipation of lindane by <i>Spinacia oleracea</i> L. <i>Ecotoxicology and Environmental Safety</i> , 2014, 109, 22-26.	2.9	33
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34	Insect herbivore-associated organisms affect plant responses to herbivory. <i>New Phytologist</i> , 2014, 204, 315-321.	3.5	78
35	Induced Systemic Resistance by Beneficial Microbes. <i>Annual Review of Phytopathology</i> , 2014, 52, 347-375.	3.5	2,193
36	Taxonomical and functional microbial community selection in soybean rhizosphere. <i>ISME Journal</i> , 2014, 8, 1577-1587.	4.4	633

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37	Exploring interactions of plant microbiomes. <i>Scientia Agricola</i> , 2014, 71, 528-539.	0.6	122
38	<i>Populus trichocarpa</i> and <i>Populus deltoides</i> Exhibit Different Metabolomic Responses to Colonization by the Symbiotic Fungus <i>Laccaria bicolor</i> . <i>Molecular Plant-Microbe Interactions</i> , 2014, 27, 546-556.	1.4	69
39	Soil microbial community responses to heat wave components: drought and high temperature. <i>Climate Research</i> , 2015, 66, 243-264.	0.4	72
40	Planting increases the abundance and structure complexity of soil core functional genes relevant to carbon and nitrogen cycling. <i>Scientific Reports</i> , 2015, 5, 14345.	1.6	26
42	Revealing crosstalk of plant and fungi in the symbiotic roots of sewage-cleaning <i>Eichhornia crassipes</i> using direct de novo metatranscriptomic analysis. <i>Scientific Reports</i> , 2015, 5, 15407.	1.6	15
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48	Isolation and genetic characterization of endophytic and rhizospheric microorganisms from <i>Butia purpurascens</i> Glassman. <i>African Journal of Microbiology Research</i> , 2015, 9, 1907-1916.	0.4	3
49	Biotic interactions in the rhizosphere in relation to plant and soil nutrient dynamics. <i>Journal of Soil Science and Plant Nutrition</i> , 2015, , 0-0.	1.7	11
50	THE RESURRECTION PLANT <i>TRIOGON SPICATUS</i> (POACEAE) HARBORS A DIVERSITY OF PLANT GROWTH PROMOTING BACTERIA IN NORTHEASTERN BRAZILIAN CAATINGA. <i>Revista Brasileira De Ciencia Do Solo</i> , 2015, 39, 993-1002.	0.5	25
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53	Metabolic transition in mycorrhizal tomato roots. <i>Frontiers in Microbiology</i> , 2015, 6, 598.	1.5	111
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57	Assessment of Culturable Tea Rhizobacteria Isolated from Tea Estates of Assam, India for Growth Promotion in Commercial Tea Cultivars. <i>Frontiers in Microbiology</i> , 2015, 6, 1252.	1.5	52
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59	Cross-Kingdom Similarities in Microbiome Ecology and Biocontrol of Pathogens. <i>Frontiers in Microbiology</i> , 2015, 6, 1311.	1.5	24
60	Optimizing Polychlorinated Biphenyl Degradation by Flavonoid-Induced Cells of the Rhizobacterium <i>Rhodococcus erythropolis</i> U23A. <i>PLoS ONE</i> , 2015, 10, e0126033.	1.1	28
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62	Root Microbiome Assemblage is Modulated by Plant Host Factors. <i>Advances in Botanical Research</i> , 2015, 75, 57-79.	0.5	28
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65	Beyond borders: investigating microbiome interactivity and diversity for advanced biocontrol technologies. <i>Microbial Biotechnology</i> , 2015, 8, 5-7.	2.0	33
66	Bioprospecting glacial ice for plant growth promoting bacteria. <i>Microbiological Research</i> , 2015, 177, 1-7.	2.5	38
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78	Cropland soil-plant systems control production and consumption of methane and nitrous oxide and their emissions to the atmosphere. <i>Soil Science and Plant Nutrition</i> , 2015, 61, 2-33.	0.8	40
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81	Metagenomic Analysis of the Airborne Environment in Urban Spaces. <i>Microbial Ecology</i> , 2015, 69, 346-355.	1.4	76
82	Insights into the role of plant on ammonia-oxidizing bacteria and archaea in the mangrove ecosystem. <i>Journal of Soils and Sediments</i> , 2015, 15, 1212-1223.	1.5	31
83	Diverse bacterial communities are recruited on spores of different arbuscular mycorrhizal fungal isolates. <i>Biology and Fertility of Soils</i> , 2015, 51, 379-389.	2.3	111
84	Bacterial communities in the rhizosphere of <i>Vitis vinifera</i> L. cultivated under distinct agricultural practices in Argentina. <i>Antonie Van Leeuwenhoek</i> , 2015, 107, 575-588.	0.7	61
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92	The root cap: a short story of life and death. <i>Journal of Experimental Botany</i> , 2015, 66, 5651-5662.	2.4	128
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97	Bacterial diversity and community structure of Western Indian Himalayan red kidney bean ( <i>Phaseolus</i> ) Tj ETQq1 1 0,784314 rgBT /Over 0.8 46	0.8	46
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114	Composition of soil microbial communities in the rhizosphere of cucumber cultivars with differing nitrogen acquisition efficiency. <i>Applied Soil Ecology</i> , 2015, 95, 90-98.	2.1	13
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130	Structured Heterogeneity in a Marine Terrace Chronosequence: Upland Mottling. <i>Vadose Zone Journal</i> , 2016, 15, 1-14.	1.3	25
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133	Impact of Microalgae-Bacteria Interactions on the Production of Algal Biomass and Associated Compounds. <i>Marine Drugs</i> , 2016, 14, 100.	2.2	293
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146	Potential of Endophytic Bacterium <i>Paenibacillus</i> sp. PHE-3 Isolated from <i>Plantago asiatica</i> L. for Reduction of PAH Contamination in Plant Tissues. <i>International Journal of Environmental Research and Public Health</i> , 2016, 13, 633.	1.2	21
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148	Specific Microbial Communities Associate with the Rhizosphere of <i>Welwitschia mirabilis</i> , a Living Fossil. <i>PLoS ONE</i> , 2016, 11, e0153353.	1.1	41
149	Impact of Reed Canary Grass Cultivation and Mineral Fertilisation on the Microbial Abundance and Genetic Potential for Methane Production in Residual Peat of an Abandoned Peat Extraction Area. <i>PLoS ONE</i> , 2016, 11, e0163864.	1.1	11

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636	Environmental filtering: A case of bacterial community assembly in soil. <i>Soil Biology and Biochemistry</i> , 2019, 136, 107531.	4.2	23
637	Drought modulates interactions between arbuscular mycorrhizal fungal diversity and barley genotype diversity. <i>Scientific Reports</i> , 2019, 9, 9650.	1.6	42
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1349	Omics technologies used in pesticide residue detection and mitigation in crop. <i>Journal of Hazardous Materials</i> , 2021, 420, 126624.	6.5	19
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2006	Root traits and soil nutrient and carbon availability drive soil microbial diversity and composition in a northern temperate forest. <i>Plant and Soil</i> , 2022, 479, 281-299.	1.8	7
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2030	Response of rhizosphere microbiomes to climate change. , 2022, , 259-274.		0
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2037	<sc>Rhizosphere</sc> <sc>bacterial communities differ among traditional maize landraces</sc>. <i>Environmental DNA</i> , 2022, 4, 1241-1249.	3.1	5
2038	The influence of biostimulants on the qualitative composition of carrots. <i>IOP Conference Series: Earth and Environmental Science</i> , 2022, 1045, 012087.	0.2	1
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2053	Impact of PaGLK transgenic poplar on microbial community and soil enzyme activity in rhizosphere soil. <i>Frontiers in Bioengineering and Biotechnology</i> , 0, 10, .	2.0	2
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2094	<i>Trichoderma harzianum</i> sensu lato TSM39: A wheat microbiome fungus that mitigates spot blotch disease of wheat ( <i>Triticum turgidum</i> L. subsp. <i>durum</i> ) caused by <i>Bipolaris sorokiniana</i> . <i>Biological Control</i> , 2022, 175, 105055.	1.4	5
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2097	The Use of Biopesticides for Sustainable Farming: Way Forward toward Sustainable Development Goals (SDGs). , 2022, , 571-596.		1
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2104	Characterization of the microbial community response to replant diseases in peach orchards. Journal of Integrative Agriculture, 2023, 22, 1082-1092.	1.7	1
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2108	Biostimulant-induced Improvement of Soil Health and Water-use Efficiency in Plants. , 2022, , 72-84.		0
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2115	Effects of Rhizosphere Bacteria on Strawberry Plants (Fragaria Ã— ananassa Duch.) under Water Deficit. International Journal of Molecular Sciences, 2022, 23, 10449.	1.8	7
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2118	Semi-Arid-Habitat-Adapted Plant-Growth-Promoting Rhizobacteria Allows Efficient Wheat Growth Promotion. <i>Agronomy</i> , 2022, 12, 2221.	1.3	5
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2130	Phytomicrobiome communications: Novel implications for stress resistance in plants. <i>Frontiers in Microbiology</i> , 0, 13, .	1.5	9
2131	Rhizosphere melatonin application reprograms nitrogen-cycling related microorganisms to modulate low temperature response in barley. <i>Frontiers in Plant Science</i> , 0, 13, .	1.7	4
2132	Biogeography and ecological functions of root-associated and soil fungi of <i>Pinus sylvestris</i> var. <i>mongolica</i> across different afforestation areas in desertified Northern China. <i>Land Degradation and Development</i> , 2023, 34, 313-326.	1.8	4
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