

J Peter W Young

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166
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115
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178
ext. papers

15,782
ext. citations

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L-index

#	Paper	IF	Citations
166	Ploughing up the wood-wide web?. <i>Nature</i> , 1998 , 394, 431	50.4	732
165	The role of ecological theory in microbial ecology. <i>Nature Reviews Microbiology</i> , 2007 , 5, 384-92	22.2	643
164	Genome of an arbuscular mycorrhizal fungus provides insight into the oldest plant symbiosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 20117-22	11.5	499
163	Improved PCR primers for the detection and identification of arbuscular mycorrhizal fungi. <i>FEMS Microbiology Ecology</i> , 2008 , 65, 339-49	4.3	463
162	The genome of <i>Rhizobium leguminosarum</i> has recognizable core and accessory components. <i>Genome Biology</i> , 2006 , 7, R34	18.3	421
161	Molecular diversity of arbuscular mycorrhizal fungi colonising arable crops. <i>FEMS Microbiology Ecology</i> , 2001 , 36, 203-209	4.3	398
160	Selectivity and functional diversity in arbuscular mycorrhizas of co-occurring fungi and plants from a temperate deciduous woodland. <i>Journal of Ecology</i> , 2002 , 90, 371-384	6	362
159	Co-existing grass species have distinctive arbuscular mycorrhizal communities. <i>Molecular Ecology</i> , 2003 , 12, 3085-95	5.7	353
158	Arbuscular mycorrhizal community composition associated with two plant species in a grassland ecosystem. <i>Molecular Ecology</i> , 2002 , 11, 1555-64	5.7	350
157	Diversity of fungal symbionts in arbuscular mycorrhizas from a natural community. <i>New Phytologist</i> , 1995 , 130, 259-265	9.8	336
156	Extensive fungal diversity in plant roots. <i>Science</i> , 2002 , 295, 2051	33.3	319
155	Molecular diversity of arbuscular mycorrhizal fungi and patterns of host association over time and space in a tropical forest. <i>Molecular Ecology</i> , 2002 , 11, 2669-78	5.7	296
154	Plant communities affect arbuscular mycorrhizal fungal diversity and community composition in grassland microcosms. <i>New Phytologist</i> , 2004 , 161, 503-515	9.8	287
153	Legume-nodulating betaproteobacteria: diversity, host range, and future prospects. <i>Molecular Plant-Microbe Interactions</i> , 2011 , 24, 1276-88	3.6	269
152	The transcriptome of the arbuscular mycorrhizal fungus <i>Glomus intraradices</i> (DAOM 197198) reveals functional tradeoffs in an obligate symbiont. <i>New Phytologist</i> , 2012 , 193, 755-769	9.8	262
151	Introducing the bacterial χ romidS not a chromosome, not a plasmid. <i>Trends in Microbiology</i> , 2010 , 18, 141-8	12.4	249
150	Diversity and phylogeny of rhizobia. <i>New Phytologist</i> , 1996 , 133, 87-94	9.8	244

149	Three phylogenetic groups of nodA and nifH genes in Sinorhizobium and Mesorhizobium isolates from leguminous trees growing in Africa and Latin America. <i>Applied and Environmental Microbiology</i> , 1998 , 64, 419-26	4.8	240
148	Nonlegumes, legumes, and root nodules harbor different arbuscular mycorrhizal fungal communities. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 6240-6	4.8	215
147	Burkholderia species are ancient symbionts of legumes. <i>Molecular Ecology</i> , 2010 , 19, 44-52	5.7	185
146	The glutamine synthetases of rhizobia: phylogenetics and evolutionary implications. <i>Molecular Biology and Evolution</i> , 2000 , 17, 309-19	8.3	180
145	Active root-inhabiting microbes identified by rapid incorporation of plant-derived carbon into RNA. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 16970-5	11.5	177
144	Symbiosis within Symbiosis: Evolving Nitrogen-Fixing Legume Symbionts. <i>Trends in Microbiology</i> , 2016 , 24, 63-75	12.4	173
143	Molecular diversity of arbuscular mycorrhizal fungi colonising Hyacinthoides non-scripta (bluebell) in a seminatural woodland. <i>Molecular Ecology</i> , 1999 , 8, 659-666	5.7	172
142	Diversity of the ribosomal internal transcribed spacers within and among isolates of Glomus mosseae and related mycorrhizal fungi. <i>New Phytologist</i> , 1996 , 133, 103-111	9.8	150
141	Differentiation of Pseudomonas solanacearum, Pseudomonas syzygii, Pseudomonas pickettii and the Blood Disease Bacterium by partial 16S rRNA sequencing: construction of oligonucleotide primers for sensitive detection by polymerase chain reaction. <i>Journal of General Microbiology</i> , 1993 , 139, 1587-94		148
140	Diversity and specificity of Rhizobium leguminosarum biovar viciae on wild and cultivated legumes. <i>Molecular Ecology</i> , 2004 , 13, 2435-44	5.7	141
139	Proof that Burkholderia strains form effective symbioses with legumes: a study of novel Mimosa-nodulating strains from South America. <i>Applied and Environmental Microbiology</i> , 2005 , 71, 7461-71	4.8	139
138	Nodulation and nitrogen fixation by Mimosa spp. in the Cerrado and Caatinga biomes of Brazil. <i>New Phytologist</i> , 2010 , 186, 934-946	9.8	133
137	Nodulation of Cyclopia spp. (Leguminosae, Papilionoideae) by Burkholderia tuberum. <i>Annals of Botany</i> , 2007 , 100, 1403-11	4.1	131
136	Specificity and resilience in the arbuscular mycorrhizal fungi of a natural woodland community. <i>Journal of Ecology</i> , 2007 , 95, 623-630	6	124
135	Impact of soil warming and shading on colonization and community structure of arbuscular mycorrhizal fungi in roots of a native grassland community. <i>Global Change Biology</i> , 2004 , 10, 52-64	11.4	113
134	Burkholderia spp. are the most competitive symbionts of Mimosa, particularly under N-limited conditions. <i>Environmental Microbiology</i> , 2009 , 11, 762-78	5.2	107
133	The evolution of specificity in the legume-rhizobium symbiosis. <i>Trends in Ecology and Evolution</i> , 1989 , 4, 341-9	10.9	101
132	Temporal variation in the arbuscular mycorrhizal communities colonising seedlings in a tropical forest. <i>FEMS Microbiology Ecology</i> , 2002 , 42, 131-6	4.3	99

131	Biodiversity of rhizobia isolated from a wide range of forest legumes in Brazil. <i>Molecular Ecology</i> , 1998 , 7, 889-95	5.7	98
130	Higher diversity of <i>Rhizobium leguminosarum</i> biovar <i>viciae</i> populations in arable soils than in grass soils. <i>Applied and Environmental Microbiology</i> , 2000 , 66, 2445-50	4.8	97
129	<i>Burkholderia sabiae</i> sp. nov., isolated from root nodules of <i>Mimosa caesalpiniiifolia</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008 , 58, 2174-9	2.2	94
128	Minimal standards for the description of new genera and species of rhizobia and agrobacteria. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019 , 69, 1852-1863	2.2	94
127	Bacterial genospecies that are not ecologically coherent: population genomics of <i>Rhizobium leguminosarum</i> . <i>Open Biology</i> , 2015 , 5, 140133	7	93
126	Ribosomal small subunit sequence variation within spores of an arbuscular mycorrhizal fungus, <i>Scutellospora</i> sp. <i>Molecular Ecology</i> , 1999 , 8, 915-21	5.7	93
125	Establishment, persistence and effectiveness of arbuscular mycorrhizal fungal inoculants in the field revealed using molecular genetic tracing and measurement of yield components. <i>New Phytologist</i> , 2012 , 194, 810-822	9.8	87
124	The mitochondrial genome sequence of the arbuscular mycorrhizal fungus <i>Glomus intraradices</i> isolate 494 and implications for the phylogenetic placement of <i>Glomus</i> . <i>New Phytologist</i> , 2009 , 183, 200-211	9.8	82
123	Nodulation of <i>Sesbania</i> species by <i>Rhizobium</i> (<i>Agrobacterium</i>) strain IRBG74 and other rhizobia. <i>Environmental Microbiology</i> , 2009 , 11, 2510-25	5.2	81
122	Substrate induction and glucose repression of maltose utilization by <i>Streptomyces coelicolor</i> A3(2) is controlled by <i>malR</i> , a member of the <i>lacI-galR</i> family of regulatory genes. <i>Molecular Microbiology</i> , 1997 , 23, 537-49	4.1	76
121	<i>Burkholderia diazotrophica</i> sp. nov., isolated from root nodules of <i>Mimosa</i> spp. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013 , 63, 435-441	2.2	75
120	Sib competition can favour sex in two ways. <i>Journal of Theoretical Biology</i> , 1981 , 88, 755-6	2.3	75
119	Relationship between assemblages of mycorrhizal fungi and bacteria on grass roots. <i>Environmental Microbiology</i> , 2008 , 10, 534-41	5.2	73
118	<i>Mesorhizobium septentrionale</i> sp. nov. and <i>Mesorhizobium temperatum</i> sp. nov., isolated from <i>Astragalus adsurgens</i> growing in the northern regions of China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2004 , 54, 2003-2012	2.2	71
117	Genetic and symbiotic characterization of rhizobia isolated from tree and herbaceous legumes grown in soils from ecologically diverse sites in Kenya. <i>Soil Biology and Biochemistry</i> , 2002 , 34, 801-811	7.5	71
116	Horizontal Transfer of Symbiosis Genes within and Between Rhizobial Genera: Occurrence and Importance. <i>Genes</i> , 2018 , 9,	4.2	70
115	An invasive <i>Mimosa</i> in India does not adopt the symbionts of its native relatives. <i>Annals of Botany</i> , 2013 , 112, 179-96	4.1	70
114	Effects of long-term fertilization on AM fungal community structure and Glomalin-related soil protein in the Loess Plateau of China. <i>Plant and Soil</i> , 2011 , 342, 233-247	4.2	70

113	Phylogeny of the Glomerales and Diversisporales (Fungi: Glomeromycota) from actin and elongation factor 1-alpha sequences. <i>FEMS Microbiology Letters</i> , 2003 , 229, 127-32	2.9	68
112	<i>Burkholderia symbiotica</i> sp. nov., isolated from root nodules of <i>Mimosa</i> spp. native to north-east Brazil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2012 , 62, 2272-2278	2.2	67
111	Real-time PCR and microscopy: are the two methods measuring the same unit of arbuscular mycorrhizal fungal abundance?. <i>Fungal Genetics and Biology</i> , 2008 , 45, 581-96	3.9	67
110	Chickpea rhizobia symbiosis genes are highly conserved across multiple <i>Mesorhizobium</i> species. <i>FEMS Microbiology Ecology</i> , 2008 , 66, 391-400	4.3	63
109	Population mixing of <i>Rhizobium leguminosarum</i> bv. <i>viciae</i> nodulating <i>Vicia faba</i> : the role of recombination and lateral gene transfer. <i>FEMS Microbiology Ecology</i> , 2010 , 73, 563-76	4.3	62
108	Interactions between <i>Pseudomonas fluorescens</i> biocontrol agents and <i>Glomus mosseae</i> , an arbuscular mycorrhizal fungus, within the rhizosphere. <i>FEMS Microbiology Letters</i> , 1998 , 166, 297-303	2.9	62
107	Quantification of an arbuscular mycorrhizal fungus, <i>Glomus mosseae</i> , within plant roots by competitive polymerase chain reaction. <i>Mycological Research</i> , 1997 , 101, 1440-1444		57
106	A common genomic framework for a diverse assembly of plasmids in the symbiotic nitrogen fixing bacteria. <i>PLoS ONE</i> , 2008 , 3, e2567	3.7	57
105	Rhizobium Population Genetics: Enzyme Polymorphism in <i>Rhizobium leguminosarum</i> from Plants and Soil in a Pea Crop. <i>Applied and Environmental Microbiology</i> , 1987 , 53, 397-402	4.8	56
104	Identification of roots from grass swards using PCR-RFLP and FFLP of the plastid trnL (UAA) intron. <i>BMC Ecology</i> , 2003 , 3, 8	2.7	55
103	Endemic <i>Mimosa</i> species from Mexico prefer alphaproteobacterial rhizobial symbionts. <i>New Phytologist</i> , 2016 , 209, 319-33	9.8	55
102	A diverse population of introns in the nuclear ribosomal genes of ericoid mycorrhizal fungi includes elements with sequence similarity to endonuclease-coding genes. <i>Molecular Biology and Evolution</i> , 2000 , 17, 44-59	8.3	54
101	Invasive <i>Robinia pseudoacacia</i> in China is nodulated by <i>Mesorhizobium</i> and <i>Sinorhizobium</i> species that share similar nodulation genes with native American symbionts. <i>FEMS Microbiology Ecology</i> , 2009 , 68, 320-8	4.3	53
100	<i>Rhizobium anhuiense</i> sp. nov., isolated from effective nodules of <i>Vicia faba</i> and <i>Pisum sativum</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015 , 65, 2960-2967	2.2	52
99	<i>Azorhizobium doebereineriae</i> sp. Nov. Microsymbiont of <i>Sesbania virgata</i> (Caz.) Pers. <i>Systematic and Applied Microbiology</i> , 2006 , 29, 197-206	4.2	51
98	High diversity of chickpea <i>Mesorhizobium</i> species isolated in a Portuguese agricultural region. <i>FEMS Microbiology Ecology</i> , 2004 , 48, 101-7	4.3	50
97	The common nodulation genes of <i>Astragalus sinicus</i> rhizobia are conserved despite chromosomal diversity. <i>Applied and Environmental Microbiology</i> , 2000 , 66, 2988-95	4.8	50
96	Multilocus sequence analysis reveals multiple symbiovars within <i>Mesorhizobium</i> species. <i>Systematic and Applied Microbiology</i> , 2012 , 35, 359-67	4.2	48

95	Modafinil in the treatment of idiopathic hypersomnia without long sleep time--a randomized, double-blind, placebo-controlled study. <i>Journal of Sleep Research</i> , 2015 , 24, 74-81	5.8	47
94	Morphogenesis of the compound leaf in three genotypes of the pea, <i>Pisum sativum</i> . <i>Canadian Journal of Botany</i> , 1986 , 64, 1268-1276		47
93	<i>Mesorhizobium alhagi</i> sp. nov., isolated from wild <i>Alhagi sparsifolia</i> in north-western China. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010 , 60, 958-962	2.2	45
92	Diversity of <i>Sinorhizobium meliloti</i> from the Central Asian Alfalfa Gene Center. <i>Applied and Environmental Microbiology</i> , 2002 , 68, 4694-7	4.8	44
91	T-RFLP analysis of bacterial communities in the midguts of <i>Apis mellifera</i> and <i>Apis cerana</i> honey bees in Thailand. <i>FEMS Microbiology Ecology</i> , 2012 , 79, 273-81	4.3	43
90	Distribution of repC plasmid-replication sequences among plasmids and isolates of <i>Rhizobium leguminosarum</i> bv. <i>viciae</i> from field populations. <i>Microbiology (United Kingdom)</i> , 1998 , 144, 771-780	2.9	42
89	<i>Bradyrhizobium guangdongense</i> sp. nov. and <i>Bradyrhizobium guangxiense</i> sp. nov., isolated from effective nodules of peanut. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015 , 65, 4655-4661	2.2	42
88	Average nucleotide identity of genome sequences supports the description of <i>Rhizobium lentis</i> sp. nov., <i>Rhizobium bangladeshense</i> sp. nov. and <i>Rhizobium binae</i> sp. nov. from lentil (<i>Lens culinaris</i>) nodules. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015 , 65, 3037-3045	2.2	41
87	Biochemical characterization of "LAP," a polymorphic aminopeptidase from the blue mussel, <i>Mytilus edulis</i> . <i>Biochemical Genetics</i> , 1979 , 17, 305-23	2.4	40
86	Increased sequencing depth does not increase captured diversity of arbuscular mycorrhizal fungi. <i>Mycorrhiza</i> , 2017 , 27, 761-773	3.9	39
85	Effect of rice cultivation systems on indigenous arbuscular mycorrhizal fungal community structure. <i>Microbes and Environments</i> , 2013 , 28, 316-24	2.6	39
84	<i>Burkholderia</i> sp. induces functional nodules on the South African invasive legume <i>Dipogon lignosus</i> (Phaseoleae) in New Zealand soils. <i>Microbial Ecology</i> , 2014 , 68, 542-55	4.4	38
83	Population genomics of <i>Sinorhizobium medicae</i> based on low-coverage sequencing of sympatric isolates. <i>ISME Journal</i> , 2011 , 5, 1722-34	11.9	37
82	Direct amplification of nodD from community DNA reveals the genetic diversity of <i>Rhizobium leguminosarum</i> in soil. <i>Environmental Microbiology</i> , 2001 , 3, 363-70	5.2	37
81	Complete Genome sequence of <i>Burkholderia phymatum</i> STM815(T), a broad host range and efficient nitrogen-fixing symbiont of <i>Mimosa</i> species. <i>Standards in Genomic Sciences</i> , 2014 , 9, 763-74		36
80	What does a bacterial genome sequence represent? Mis-assignment of MAFF 303099 to the genospecies <i>Mesorhizobium loti</i> . <i>Microbiology (United Kingdom)</i> , 2002 , 148, 3330-3331	2.9	36
79	Symbiotic and genetic diversity of <i>Rhizobium galegae</i> isolates collected from the <i>Galega orientalis</i> gene center in the Caucasus. <i>Applied and Environmental Microbiology</i> , 2003 , 69, 1067-74	4.8	35
78	The replicator region of the <i>Rhizobium leguminosarum</i> cryptic plasmid pRL8JI. <i>FEMS Microbiology Letters</i> , 1995 , 133, 53-8	2.9	35

77	Slipins: ancient origin, duplication and diversification of the stomatin protein family. <i>BMC Evolutionary Biology</i> , 2008 , 8, 44	3	33
76	dnaJ is a useful phylogenetic marker for alphaproteobacteria. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008 , 58, 2839-49	2.2	33
75	A genetic discontinuity in root-nodulating bacteria of cultivated pea in the Indian trans-Himalayas. <i>Molecular Ecology</i> , 2012 , 21, 145-59	5.7	32
74	Mesorhizobium camelthorni sp. nov., isolated from Alhagi sparsifolia. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011 , 61, 574-579	2.2	31
73	Defining functional diversity for lignocellulose degradation in a microbial community using multi-omics studies. <i>Biotechnology for Biofuels</i> , 2018 , 11, 166	7.8	29
72	Genetic and genomic glimpses of the elusive arbuscular mycorrhizal fungi. <i>Current Opinion in Plant Biology</i> , 2012 , 15, 454-61	9.9	29
71	A distinct class of peas (Pisum sativum L.) from Afghanistan that show strain specificity for symbiotic Rhizobium. <i>Heredity</i> , 1982 , 48, 203-210	3.6	29
70	Phylogeny of bethylid wasps (Hymenoptera: Bethyridae) inferred from 28S and 16S rRNA genes. <i>Insect Systematics and Evolution</i> , 2010 , 41, 55-73	0.6	28
69	A search for peas (Pisum sativum L.) showing strain specificity for symbiotic Rhizobium leguminosarum. <i>Heredity</i> , 1982 , 48, 197-201	3.6	28
68	Sequence diversity of the plasmid replication gene repC in the Rhizobiaceae. <i>Plasmid</i> , 2000 , 44, 209-19	3.3	27
67	A typing scheme for the honeybee pathogen Melissococcus plutonius allows detection of disease transmission events and a study of the distribution of variants. <i>Environmental Microbiology Reports</i> , 2013 , 5, 525-9	3.7	26
66	Characterisation of rhizobia from African acacias and other tropical woody legumes using Biolog [®] and partial 16S rRNA sequencing. <i>FEMS Microbiology Letters</i> , 1999 , 170, 111-117	2.9	26
65	Coordinated regulation of core and accessory genes in the multipartite genome of Sinorhizobium fredii. <i>PLoS Genetics</i> , 2018 , 14, e1007428	6	25
64	Genotypic characterisation of rhizobia nodulating Vicia faba from the soils of Jordan: a comparison with UK isolates. <i>Soil Biology and Biochemistry</i> , 2003 , 35, 709-714	7.5	25
63	How many fungi does it take to change a plant community?. <i>Trends in Plant Science</i> , 1999 , 4, 81-82	13.1	25
62	Evolutionary dynamics of insertion sequences in relation to the evolutionary histories of the chromosome and symbiotic plasmid genes of Rhizobium etli populations. <i>Applied and Environmental Microbiology</i> , 2010 , 76, 6504-13	4.8	24
61	Sexual swarms in Daphnia magna, a cyclic parthenogen. <i>Freshwater Biology</i> , 1978 , 8, 279-281	3.1	24
60	Host-specific competitiveness to form nodules in Rhizobium leguminosarum symbiovar viciae. <i>New Phytologist</i> , 2020 , 226, 555-568	9.8	24

59	Genome diversity in arbuscular mycorrhizal fungi. <i>Current Opinion in Plant Biology</i> , 2015 , 26, 113-9	9.9	23
58	Revealing the insoluble metasecretome of lignocellulose-degrading microbial communities. <i>Scientific Reports</i> , 2017 , 7, 2356	4.9	23
57	DNA-based Identification of Goose Species from Two Archaeological Sites in Lincolnshire. <i>Journal of Archaeological Science</i> , 2000 , 27, 91-100	2.9	23
56	Bacteria Are Smartphones and Mobile Genes Are Apps. <i>Trends in Microbiology</i> , 2016 , 24, 931-932	12.4	23
55	Rhizobium leguminosarum is the symbiont of lentils in the Middle East and Europe but not in Bangladesh. <i>FEMS Microbiology Ecology</i> , 2014 , 87, 64-77	4.3	22
54	Genome sequencing of two Neorhizobium galegae strains reveals a noeT gene responsible for the unusual acetylation of the nodulation factors. <i>BMC Genomics</i> , 2014 , 15, 500	4.5	22
53	Diversity and persistence of arbuscular mycorrhizas in a low-Arctic meadow habitat. <i>New Phytologist</i> , 2007 , 176, 691-698	9.8	22
52	A molecular guide to the taxonomy of arbuscular mycorrhizal fungi. <i>New Phytologist</i> , 2012 , 193, 823-826	9.8	20
51	Acquisition of an Agrobacterium Ri plasmid and pathogenicity by other alpha-Proteobacteria in cucumber and tomato crops affected by root mat. <i>Applied and Environmental Microbiology</i> , 2004 , 70, 2779-85	4.8	20
50	Molecular diversity of Frankia in root nodules of Alnus incana grown with inoculum from polluted urban soils. <i>FEMS Microbiology Ecology</i> , 2004 , 50, 255-63	4.3	20
49	Rhizobium etli is the dominant common bean nodulating rhizobia in cultivated soils from different locations in Jordan. <i>Applied Soil Ecology</i> , 2004 , 26, 193-200	5	20
48	Rhizobium population genetics: Effect of clover variety and inoculum dilution on the genetic diversity sampled from natural populations. <i>Plant and Soil</i> , 1987 , 103, 147-150	4.2	20
47	International Committee on Systematics of Prokaryotes Subcommittee for the Taxonomy of Rhizobium and Agrobacterium Minutes of the meeting, Budapest, 25 August 2016. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017 , 67, 2485-2494	2.2	20
46	Defining the Species Complex. <i>Genes</i> , 2021 , 12,	4.2	19
45	Rhizobia with 16S rRNA and nifH similar to Mesorhizobium huakuii but Novel recA, glnII, nodA and nodC genes are symbionts of New Zealand Carmichaelinae. <i>PLoS ONE</i> , 2012 , 7, e47677	3.7	17
44	Rhizobium population genetics: Host preference and strain competition effects on the range of Rhizobium leguminosarum biovar Trifolii genotypes isolated from natural populations. <i>Soil Biology and Biochemistry</i> , 1989 , 21, 981-986	7.5	16
43	Symbiosis genes show a unique pattern of introgression and selection within a species complex. <i>Microbial Genomics</i> , 2020 , 6,	4.4	15
42	A new clade of Mesorhizobium nodulating Alhagi sparsifolia. <i>Systematic and Applied Microbiology</i> , 2009 , 32, 8-16	4.2	14

41	The genetic diversity of intraterrestrial aliens. <i>New Phytologist</i> , 2008 , 178, 465-8	9.8	14
40	Recurrent outbreaks of root mat in cucumber and tomato are associated with a monomorphic, cucumopine, Ri-plasmid harboured by various Alphaproteobacteria. <i>FEMS Microbiology Letters</i> , 2006 , 258, 136-43	2.9	14
39	Arbuscular mycorrhizal communities associated with maples (<i>Acer</i> spp.) in a common garden are influenced by season and host plant. <i>Botany</i> , 2014 , 92, 321-326	1.3	13
38	Modification of Pea Leaf Morphology by 2,3,5-Triiodobenzoic Acid. <i>Botanical Gazette</i> , 1991 , 152, 133-138		13
37	Identification and analysis of rhizobial plasmid origins of transfer. <i>FEMS Microbiology Ecology</i> , 2002 , 42, 227-34	4.3	11
36	L-System Analysis of Compound Leaf Development in <i>Pisum sativum</i> L. <i>Annals of Botany</i> , 1992 , 70, 189-196	1.6	11
35	Characterization of arbuscular mycorrhizal fungus communities of <i>Aquilaria crassna</i> and <i>Tectona grandis</i> roots and soils in Thailand plantations. <i>PLoS ONE</i> , 2014 , 9, e112591	3.7	11
34	Ecology and Evolution of Rhizobia 2019 ,		11
33	The NfeD protein family and its conserved gene neighbours throughout prokaryotes: functional implications for stomatin-like proteins. <i>Journal of Molecular Evolution</i> , 2009 , 69, 657-67	3.1	10
32	Kissing cousins: mycorrhizal fungi get together. <i>New Phytologist</i> , 2009 , 181, 751-753	9.8	10
31	The determination of pea leaves, leaflets, and tendrils. <i>American Journal of Botany</i> , 1994 , 81, 352-360	2.7	10
30	Linkage of <i>sym-2</i> , the symbiotic specificity locus of <i>Pisum sativum</i> . <i>Journal of Heredity</i> , 1985 , 76, 207-208	2.4	9
29	International Committee on Systematics of Prokaryotes Subcommittee on the taxonomy of rhizobia and agrobacteria Minutes of the closed meeting, Granada, 4 September 2017. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018 , 68, 3363-3368	2.2	9
28	User-friendly bioinformatics pipeline gDAT (graphical downstream analysis tool) for analysing rDNA sequences. <i>Molecular Ecology Resources</i> , 2021 , 21, 1380-1392	8.4	8
27	Induction of root-mat symptoms on cucumber plants by <i>Rhizobium</i> , but not by <i>Ochrobactrum</i> or <i>Sinorhizobium</i> , harbouring a cucumopine Ri plasmid. <i>Plant Pathology</i> , 2005 , 54, 799-805	2.8	7
26	Does growth rate determine leaf form in <i>Pisum sativum</i> ?. <i>Canadian Journal of Botany</i> , 1989 , 67, 2590-2595		7
25	The determination of pea leaves, leaflets, and tendrils 1994 , 81, 352		7
24	Maximizing the Adjacent Possible in Automata Chemistries. <i>Artificial Life</i> , 2016 , 22, 49-75	1.4	6

23	The molecular palaeoecology of geese: identification of archaeological goose remains using ancient DNA analysis. <i>International Journal of Osteoarchaeology</i> , 1998 , 8, 280-287	1.1	5
22	International Committee on Systematics of Prokaryotes Subcommittee on the Taxonomy of Rhizobia and Agrobacteria Minutes of the meeting by video conference, 11 July 2018. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019 , 69, 1835-1840	2.2	5
21	International Committee on Systematics of Prokaryotes Subcommittee on the Taxonomy of Rhizobia and Agrobacteria Minutes of the closed meeting by videoconference, 17 July 2019. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020 , 70, 3563-3571	2.2	4
20	The genomic architecture of introgression among sibling species of bacteria		4
19	PLAZZMID: An Evolutionary Agent-Based Architecture Inspired by Bacteria and Bees 2007 , 1151-1160		3
18	History of Rhizobial Taxonomy 2019 , 23-39		3
17	MAUI-seq: Metabarcoding using amplicons with unique molecular identifiers to improve error correction		2
16	Molecular Microprograms. <i>Lecture Notes in Computer Science</i> , 2011 , 297-304	0.9	2
15	Symbiosis Genes: Organisation and Diversity 2019 , 123-144		2
14	Interactions between <i>Pseudomonas fluorescens</i> biocontrol agents and <i>Glomus mosseae</i> , an arbuscular mycorrhizal fungus, within the rhizosphere		2
13	Amplicons and isolates: Rhizobium diversity in fields under conventional and organic management		2
12	International Committee on Systematics of Prokaryotes Subcommittee on the Taxonomy of Rhizobia and Agrobacteria Minutes of the closed meeting by videoconference, 6 July 2020. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021 , 71,	2.2	2
11	MAUI-seq: Metabarcoding using amplicons with unique molecular identifiers to improve error correction. <i>Molecular Ecology Resources</i> , 2021 , 21, 703-720	8.4	2
10	Genetic variation is associated with differences in facilitative and competitive interactions in the <i>Rhizobium leguminosarum</i> species complex. <i>Environmental Microbiology</i> , 2021 ,	5.2	2
9	Gene regulation in a particle metabolome 2009 ,		1
8	Why are rhizobial symbiosis genes mobile?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2022 , 377, 20200471	5.8	1
7	Introducing a Novel, Broad Host Range Temperate Phage Family Infecting and Beyond. <i>Frontiers in Microbiology</i> , 2021 , 12, 765271	5.7	1
6	Genomics and Evolution of Rhizobia 2019 , 103-119		1

- 5 Fields with no recent legume cultivation have sufficient nitrogen-fixing rhizobia for crops of faba bean (*Vicia faba* L.). *Plant and Soil*, **2022**, 472, 345-368 4.2 1
- 4 Genetic Variation in Host-Specific Competitiveness of the Symbiont Symbiovar. *Frontiers in Plant Science*, **2021**, 12, 719987 6.2 0
- 3 Molecular biology of the Rhizobiaceae. *New Phytologist*, **2001**, 149, 17-17 9.8
- 2 Novel Mimosa-Nodulating Strains of Burkholderia from South America. *Current Plant Science and Biotechnology in Agriculture*, **2005**, 391-393
- 1 Evolution of Symbiosis Genes: Vertical and Horizontal Gene Transfer **2019**, 145-152