## Ramon A Rosselló-Móra

List of Publications by Year in descending order

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209 papers 27,495 citations

59 h-index 159

224 all docs

224 docs citations

times ranked

224

19804 citing authors

g-index

#	Article	lF	CITATIONS
1	Solar salterns as model systems to study the units of bacterial diversity that matter for ecosystem functioning. Current Opinion in Biotechnology, 2022, 73, 151-157.	6.6	8
2	Judicial Opinions 103–111. International Journal of Systematic and Evolutionary Microbiology, 2022, 72, .	1.7	8
3	Toward quantifying the adaptive role of bacterial pangenomes during environmental perturbations. ISME Journal, 2022, 16, 1222-1234.	9.8	13
4	Litter Management Strategies and Their Impact on the Environmental and Respiratory Microbiome Might Influence Health in Poultry. Microorganisms, 2022, 10, 878.	3.6	1
5	Distinct ecotypes within a natural haloarchaeal population enable adaptation to changing environmental conditions without causing population sweeps. ISME Journal, 2021, 15, 1178-1191.	9.8	14
6	Preparing a revision of the International Code of Nomenclature of Prokaryotes. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	17
7	Emendation of Rules 5b, 8, 15 and 22 of the International Code of Nomenclature of Prokaryotes to include the rank of phylum. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	27
8	Public discussion on a proposed revision of the International Code of Nomenclature of Prokaryotes. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	19
9	Evidence for the existence of a new genus Chlamydiifrater gen. nov. inside the family Chlamydiaceae with two new species isolated from flamingo (Phoenicopterus roseus): Chlamydiifrater phoenicopteri sp. nov. and Chlamydiifrater volucris sp. nov Systematic and Applied Microbiology, 2021, 44, 126200.	2.8	24
10	Description of three new Alteromonas species Alteromonas antoniana sp. nov., Alteromonas lipotrueae sp. nov. and Alteromonas lipotrueiana sp. nov. isolated from marine environments, and proposal for reclassification of the genus Salinimonas as Alteromonas. Systematic and Applied Microbiology, 2021, 44, 126226.	2.8	39
11	Ancient saltern metagenomics: tracking changes in microbes and their viruses from the underground to the surface. Environmental Microbiology, 2021, 23, 3477-3498.	3.8	6
12	Release LTP_12_2020, featuring a new ARB alignment and improved 16S rRNA tree for prokaryotic type strains. Systematic and Applied Microbiology, 2021, 44, 126218.	2.8	44
13	Emendation of General Consideration 5 and Rules 18a, 24a and 30 of the International Code of Nomenclature of Prokaryotes to resolve the status of the Cyanobacteria in the prokaryotic nomenclature. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	1.7	15
14	Addressing the sublime scale of the microbial world: reconciling an appreciation of microbial diversity with the need to describe species. New Microbes and New Infections, 2021, 43, 100931.	1.6	14
15	Cultivable $\langle i \rangle$ Winogradskyella $\langle  i \rangle$ species are genomically distinct from the sympatric abundant candidate species. ISME Communications, 2021, 1, .	4.2	10
16	Landscapes and bacterial signatures of mucosa-associated intestinal microbiota in Chilean and Spanish patients with inflammatory bowel disease. Microbial Cell, 2021, 8, 223-238.	3.2	11
17	Inverted microbial community stratification and spatial–temporal stability in hypersaline anaerobic sediments from the S'Avall solar salterns. Systematic and Applied Microbiology, 2021, 44, 126231.	2.8	5
18	Comparative genome analysis of the genus Hydrotalea and proposal of the novel species Hydrotalea lipotrueae sp. nov., isolated from a groundwater aquifer in the south of Mallorca Island, Spain. Systematic and Applied Microbiology, 2021, 44, 126277.	2.8	7

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19	Profiling the Bladder Microbiota in Patients With Bladder Cancer. Frontiers in Microbiology, 2021, 12, 718776.	3.5	13
20	Taxonomic study of nine new Winogradskyella species occurring in the shallow waters of Helgoland Roads, North Sea. Proposal of Winogradskyella schleiferi sp. nov., Winogradskyella costae sp. nov., Winogradskyella helgolandensis sp. nov., Winogradskyella vidalii sp. nov., Winogradskyella forsetii sp. nov., Winogradskyella ludwigii sp. nov., Winogradskyella ursingii sp. nov., Winogradskyella wichelsiae sp. nov., and Candidatus "Winogradskyella atlantica―sp. nov Systematic and Applied	2.8	38
21	Microbiology, 2020, 43, 126128. Species-Level Analysis of Human Gut Microbiota With Metataxonomics. Frontiers in Microbiology, 2020, 11, 2029.	3.5	50
22	Roadmap for naming uncultivated Archaea and Bacteria. Nature Microbiology, 2020, 5, 987-994.	13.3	115
23	Complete mitochondrial genome of the pearly razorfish <i>Xyrichtys novacula:</i> phylogenetic analysis of its placement within the Labridae family. Mitochondrial DNA Part B: Resources, 2020, 5, 644-645.	0.4	2
24	Opinion: Response to concerns about the use of DNA sequences as types in the nomenclature of prokaryotes. Systematic and Applied Microbiology, 2020, 43, 126070.	2.8	8
25	High-throughput cultivation of heterotrophic bacteria during a spring phytoplankton bloom in the North Sea. Systematic and Applied Microbiology, 2020, 43, 126066.	2.8	21
26	Advantages outweigh concerns about using genome sequence as type material for prokaryotic taxonomy. Environmental Microbiology, 2020, 22, 819-822.	3.8	12
27	Ancestry and adaptive radiation of Bacteroidetes as assessed by comparative genomics. Systematic and Applied Microbiology, 2020, 43, 126065.	2.8	17
28	<scp>The importance of naturally attenuated SARSâ€CoV</scp> â€2 <scp>in the fight against COVID</scp> â€19. Environmental Microbiology, 2020, 22, 1997-2000.	3.8	54
29	Novel Genes Involved in Resistance to Both Ultraviolet Radiation and Perchlorate From the Metagenomes of Hypersaline Environments. Frontiers in Microbiology, 2020, 11, 453.	3.5	10
30	Glycine Betaine and Ectoine Are the Major Compatible Solutes Used by Four Different Halophilic Heterotrophic Ciliates. Microbial Ecology, 2019, 77, 317-331.	2.8	12
31	Dialogue on the nomenclature and classification of prokaryotes. Systematic and Applied Microbiology, 2019, 42, 5-14.	2.8	41
32	Predominance of deterministic microbial community dynamics in salterns exposed to different light intensities. Environmental Microbiology, 2019, 21, 4300-4315.	3.8	20
33	Reflections on the introduction of the Digital Protologue Database $\hat{a} \in \text{``A partial success?. Systematic}$ and Applied Microbiology, 2019, 42, 1-2.	2.8	4
34	Toward unrestricted use of public genomic data. Science, 2019, 363, 350-352.	12.6	45
35	Reflections on the introduction of the Digital Protologue Database – a partial success?. Antonie Van Leeuwenhoek, 2019, 112, 141-143.	1.7	5
36	Prokaryotic and viral community of the sulfateâ€rich crust from Peñahueca ephemeral lake, an astrobiology analogue. Environmental Microbiology, 2019, 21, 3577-3600.	3.8	9

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37	Consent insufficient for data release—Response. Science, 2019, 364, 446-446.	12.6	5
38	First description of two moderately halophilic and psychrotolerant Mycoplasma species isolated from cephalopods and proposal of Mycoplasma marinum sp. nov. and Mycoplasma todarodis sp. nov. Systematic and Applied Microbiology, 2019, 42, 457-467.	2.8	22
39	" <i>Candidatus</i> Macondimonas diazotrophicaâ€, a novel gammaproteobacterial genus dominating crude-oil-contaminated coastal sediments. ISME Journal, 2019, 13, 2129-2134.	9.8	46
40	Moving the cataloguing of the "uncultivated majorityâ€forward. Systematic and Applied Microbiology, 2019, 42, 3-4.	2.8	3
41	Ming et al. 2016, Meiothermus terrae Yu et al. 2014 and Meiothermus timidus Pires et al. 2005, to Calidithermus gen. nov., as Calidithermus chliarophilus comb. nov., Calidithermus roseus comb. nov., Calidithermus terrae comb. nov. and Calidithermus timidus comb. nov., respectively, and emended description of the genus Meiothermus. International Journal of Systematic and Evolutionary	1.7	28
42	Proposal for changes in the International Code of Nomenclature of Prokaryotes: granting priority to Candidatus names. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 2174-2175.	1.7	27
43	Genomic comparison between members of the Salinibacteraceae family, and description of a new species of Salinibacter (Salinibacter altiplanensis sp. nov.) isolated from high altitude hypersaline environments of the Argentinian Altiplano. Systematic and Applied Microbiology, 2018, 41, 198-212.	2.8	29
44	Biogeographical patterns of bacterial and archaeal communities from distant hypersaline environments. Systematic and Applied Microbiology, 2018, 41, 139-150.	2.8	39
45	Vibrio communities in scleractinian corals differ according to health status and geographic location in the Mediterranean Sea. Systematic and Applied Microbiology, 2018, 41, 131-138.	2.8	23
46	Reply to the commentary "Uncultivated microbes—in need of their own nomenclature?― ISME Journal, 2018, 12, 653-654.	9.8	8
47	Characterization of ecologically diverse viruses infecting co-occurring strains of cosmopolitan hyperhalophilic <i>Bacteroidetes</i> . ISME Journal, 2018, 12, 424-437.	9.8	29
48	Precise Fecal Microbiome of the Herbivorous Tibetan Antelope Inhabiting High-Altitude Alpine Plateau. Frontiers in Microbiology, 2018, 9, 2321.	3 <b>.</b> 5	33
49	Temperature modulates Fischerella thermalis ecotypes in Porcelana Hot Spring. Systematic and Applied Microbiology, 2018, 41, 531-543.	2.8	32
50	Effect of the natural arsenic gradient on the diversity and arsenic resistance of bacterial communities of the sediments of Camarones River (Atacama Desert, Chile). PLoS ONE, 2018, 13, e0195080.	<b>2.</b> 5	16
51	Non-halophilic endophytes associated with the euhalophyte Arthrocnemum macrostachyum and their plant growth promoting activity potential. FEMS Microbiology Letters, 2018, 365, .	1.8	9
52	The Microbial Genomes Atlas (MiGA) webserver: taxonomic and gene diversity analysis of Archaea and Bacteria at the whole genome level. Nucleic Acids Research, 2018, 46, W282-W288.	14.5	458
53	Proposal of the suffix –ota to denote phyla. Addendum to â€ <sup>*</sup> Proposal to include the rank of phylum in the International Code of Nomenclature of Prokaryotes'. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 967-969.	1.7	136
54	Metataxonomics reveal vultures as a reservoir for Clostridium perfringens. Emerging Microbes and Infections, 2017, 6, 1-8.	6.5	40

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55	Introducing a Digital Protologue: A timely move towards a database-driven systematics of Archaea and Bacteria. Systematic and Applied Microbiology, 2017, 40, 121-122.	2.8	40
56	The low diverse gastric microbiome of the jellyfish <i>Cotylorhiza tuberculata </i> is dominated by four novel taxa. Environmental Microbiology, 2017, 19, 3039-3058.	3.8	62
57	Transition boundaries for protistan species turnover in hypersaline waters of different biogeographic regions. Environmental Microbiology, 2017, 19, 3186-3200.	3.8	27
58	Uncultivated microbes in need of their own taxonomy. ISME Journal, 2017, 11, 2399-2406.	9.8	572
59	Introducing a digital protologue: a timely move towards a database-driven systematics of archaea and bacteria. Antonie Van Leeuwenhoek, 2017, 110, 455-456.	1.7	85
60	Distinctive Gut Microbiota Is Associated with Diarrheagenic Escherichia coli Infections in Chilean Children. Frontiers in Cellular and Infection Microbiology, 2017, 7, 424.	3.9	26
61	Genetic Diversity and Virulence Determinants of Escherichia coli Strains Isolated from Patients with Crohn's Disease in Spain and Chile. Frontiers in Microbiology, 2017, 8, 639.	3.5	62
62	Endophytic microbial diversity of the halophyte <i>Arthrocnemum macrostachyum</i> compartments. FEMS Microbiology Ecology, 2016, 92, fiw145.	2.7	56
63	Revised phylogeny of Bacteroidetes and proposal of sixteen new taxa and two new combinations including Rhodothermaeota phyl. nov Systematic and Applied Microbiology, 2016, 39, 281-296.	2.8	214
64	After All, Only Millions?. MBio, 2016, 7, .	4.1	38
65	Reply to "The Underestimation of Global Microbial Diversity― MBio, 2016, 7, .	4.1	6
66	Effects of the 2015 heat wave on benthic invertebrates inÂtheÂTabarcaÂMarine Protected Area (southeast) Tj ET	Qg0500 rs	gBT/Overlock
67	Meeting report: GenBank microbial genomic taxonomy workshop ( $12\hat{a}$ $\in$ "13 May, 2015). Standards in Genomic Sciences, 2016, 11, .	1.5	81
68	JSpeciesWS: a web server for prokaryotic species circumscription based on pairwise genome comparison. Bioinformatics, 2016, 32, 929-931.	4.1	2,023
69	Halorhabdus rudnickae sp. nov., a halophilic archaeon isolated from a salt mine borehole in Poland. Systematic and Applied Microbiology, 2016, 39, 100-105.	2.8	23
70	Filimonas aurantiibacter sp. nov., an orange-pigmented bacterium isolated from lake water and emended description of the genus Filimonas. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4027-4032.	1.7	13
71	Salt resistance genes revealed by functional metagenomics from brines and moderate-salinity rhizosphere within a hypersaline environment. Frontiers in Microbiology, 2015, 6, 1121.	3.5	45
72	Diversity of extremely halophilic cultivable prokaryotes in Mediterranean, Atlantic and Pacific solar salterns: Evidence that unexplored sites constitute sources of cultivable novelty. Systematic and Applied Microbiology, 2015, 38, 266-275.	2.8	46

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73	Classifying the uncultivated microbial majority: A place for metagenomic data in the Candidatus proposal. Systematic and Applied Microbiology, 2015, 38, 223-230.	2.8	61
74	Emendation of the family Chlamydiaceae: Proposal of a single genus, Chlamydia, to include all currently recognized species. Systematic and Applied Microbiology, 2015, 38, 99-103.	2.8	156
75	Past and future species definitions for Bacteria and Archaea. Systematic and Applied Microbiology, 2015, 38, 209-216.	2.8	470
76	Crohn associated microbial communities associated to colonic mucosal biopsies in patients of the western Mediterranean. Systematic and Applied Microbiology, 2015, 38, 442-452.	2.8	37
77	Moderate halophilic bacteria colonizing the phylloplane of halophytes of the subfamily Salicornioideae (Amaranthaceae). Systematic and Applied Microbiology, 2015, 38, 406-416.	2.8	58
78	The effect of oil spills on the bacterial diversity and catabolic function in coastal sediments: a case study on the Prestige oil spill. Environmental Science and Pollution Research, 2015, 22, 15200-15214.	5.3	80
79	Taxonomy in the age of genomics. Systematic and Applied Microbiology, 2015, 38, 207-208.	2.8	12
80	Prokaryotic microbiota in the digestive cavity of the jellyfish Cotylorhiza tuberculata. Systematic and Applied Microbiology, 2015, 38, 494-500.	2.8	21
81	Microbial diversity and dynamics of a groundwater and a still bottled natural mineral water. Environmental Microbiology, 2015, 17, 577-593.	3.8	40
82	From community approaches to single-cell genomics: the discovery of ubiquitous hyperhalophilic <i>Bacteroidetes</i> generalists. ISME Journal, 2015, 9, 16-31.	9.8	51
83	Proposal to include the rank of phylum in the International Code of Nomenclature of Prokaryotes. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 4284-4287.	1.7	84
84	Species. , 2015, , 2304-2306.		0
85	Defining Microbial Diversity-the Species Concept for Prokaryotic and Eukaryotic Microorganisms. , 2014, , 29-39.		11
86	Genomic Encyclopedia of Bacteria and Archaea: Sequencing a Myriad of Type Strains. PLoS Biology, 2014, 12, e1001920.	5.6	190
87	Harmonized Phylogenetic Trees for The Prokaryotes. , 2014, , 1-3.		10
88	Evidence for the existence of two new members of the family Chlamydiaceae and proposal of Chlamydia avium sp. nov. and Chlamydia gallinacea sp. nov Systematic and Applied Microbiology, 2014, 37, 79-88.	2.8	154
89	Uniting the classification of cultured and uncultured bacteria and archaea using 16S rRNA gene sequences. Nature Reviews Microbiology, 2014, 12, 635-645.	28.6	2,000
90	Technical note: Comparison of automated ribosomal intergenic spacer analysis and denaturing gradient gel electrophoresis to assess bacterial diversity in the rumen of sheep1. Journal of Animal Science, 2014, 92, 1083-1088.	0.5	10

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91	"Candidatus Haloectosymbiotes riaformosensis―(Halobacteriaceae), an archaeal ectosymbiont of the hypersaline ciliate Platynematum salinarum. Systematic and Applied Microbiology, 2014, 37, 244-251.	2.8	21
92	Composición de la comunidad procariota involucrada en la producción de nitrógeno en sedimentos de la bahÃa Mejillones. Revista De Biologia Marina Y Oceanografia, 2014, 49, 225-241.	0.2	1
93	The Family Rhodobiaceae. , 2014, , 513-531.		3
94	Species., 2014,, 1-2.		0
95	Characterization of the anaerobic microbial community in oilâ€polluted subtidal sediments: aromatic biodegradation potential after the ⟨i⟩Prestige⟨ i⟩ oil spill. Environmental Microbiology, 2013, 15, 77-92.	3.8	132
96	New insights into the archaeal diversity of a hypersaline microbial mat obtained by a metagenomic approach. Systematic and Applied Microbiology, 2013, 36, 205-214.	2.8	31
97	Description of Bacillus toyonensis sp. nov., a novel species of the Bacillus cereus group, and pairwise genome comparisons of the species of the group by means of ANI calculations. Systematic and Applied Microbiology, 2013, 36, 383-391.	2.8	217
98	Sphingobacterium psychroaquaticum sp. nov., a psychrophilic bacterium isolated from Lake Michigan water. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 952-958.	1.7	40
99	Sequencing orphan species initiative (SOS): Filling the gaps in the 16S rRNA gene sequence database for all species with validly published names. Systematic and Applied Microbiology, 2013, 36, 69-73.	2.8	98
100	Complete Genome Sequence of Bacillus toyonensis BCT-7112 $\langle \sup >T \langle   \sup > \rangle$ , the Active Ingredient of the Feed Additive Preparation Toyocerin. Genome Announcements, 2013, 1, .	0.8	30
101	Diversity of Benzylsuccinate Synthase-Like ( <i>bssA</i> ) Genes in Hydrocarbon-Polluted Marine Sediments Suggests Substrate-Dependent Clustering. Applied and Environmental Microbiology, 2013, 79, 3667-3676.	3.1	52
102	Metaproteogenomic insights beyond bacterial response to naphthalene exposure and bio-stimulation. ISME Journal, 2013, 7, 122-136.	9.8	124
103	High Metabolomic Microdiversity within Co-Occurring Isolates of the Extremely Halophilic Bacterium Salinibacter ruber. PLoS ONE, 2013, 8, e64701.	2.5	48
104	All-Species Living Tree Project. , 2013, , 1-11.		0
105	Culture-Independent Approaches for Studying Viruses from Hypersaline Environments. Applied and Environmental Microbiology, 2012, 78, 1635-1643.	3.1	70
106	Mycoplasma neophronis sp. nov., isolated from the upper respiratory tract of Canarian Egyptian vultures (Neophron percnopterus majorensis). International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 1321-1325.	1.7	13
107	Pseudarcicella hirudinis gen. nov., sp. nov., isolated from the skin of the medical leech Hirudo medicinalis. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 2247-2251.	1.7	27
108	Nevskia aquatilis sp. nov. and Nevskia persephonica sp. nov., isolated from a mineral water aquifer and the emended description of the genus Nevskia. Systematic and Applied Microbiology, 2012, 35, 297-301.	2.8	25

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109	Neoscardovia arbecensis gen. nov., sp. nov., isolated from porcine slurries. Systematic and Applied Microbiology, 2012, 35, 374-379.	2.8	19
110	On the fitness of microbial taxonomy. Trends in Microbiology, 2012, 20, 514-516.	7.7	25
111	Sphingomicrobium lutaoense gen. nov., sp. nov., isolated from a coastal hot spring. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 1326-1330.	1.7	29
112	Towards a taxonomy of <i> Bacteria &lt; /i &gt; and <i> Archaea &lt; /i &gt; based on interactive and cumulative data repositories. Environmental Microbiology, 2012, 14, 318-334.</i></i>	3.8	64
113	DNA–DNA Hybridization. Methods in Microbiology, 2011, 38, 325-347.	0.8	19
114	Response of sulfateâ€reducing bacteria to an artificial oilâ€spill in a coastal marine sediment. Environmental Microbiology, 2011, 13, 1488-1499.	3.8	55
115	Metatranscriptomic analysis of extremely halophilic viral communities. ISME Journal, 2011, 5, 1621-1633.	9.8	36
116	Evaluation of matrix-assisted laser desorption ionization-time of flight whole cell profiles for assessing the cultivable diversity of aerobic and moderately halophilic prokaryotes thriving in solar saltern sediments. Systematic and Applied Microbiology, 2011, 34, 69-75.	2.8	47
117	MALDI-TOF MS: A return to phenotyping in microbial identification?. Systematic and Applied Microbiology, 2011, 34, 1.	2.8	13
118	Release LTPs104 of the All-Species Living Tree. Systematic and Applied Microbiology, 2011, 34, 169-170.	2.8	146
119	The genus Allochromatium (Chromatiales Chromatiaceae) revisited: A study on its intragenic structure based on multilocus sequence analysis (MLSA) and DNA–DNA hybridization (DDH). Systematic and Applied Microbiology, 2011, 34, 590-594.	2.8	4
120	Response to adverse conditions in two strains of the extremely halophilic species Salinibacter ruber. Extremophiles, 2011, 15, 379-389.	2.3	22
121	Taxonomic and Functional Metagenomic Profiling of the Microbial Community in the Anoxic Sediment of a Sub-saline Shallow Lake (Laguna de Carrizo, Central Spain). Microbial Ecology, 2011, 62, 824-837.	2.8	51
122	Determination of cobalamins (hydroxo-, cyano-, adenosyl- and methyl-cobalamins) in seawater using reversed-phase liquid chromatography with diode-array detection. Analytica Chimica Acta, 2011, 701, 81-85.	5.4	20
123	From Genomics to Microevolution and Ecology: The Case of Salinibacter ruber. , 2011, , 109-122.		2
124	Evaluation of the use of multilocus sequence analysis (MLSA) to resolve taxonomic conflicts within the genus Marichromatium. Systematic and Applied Microbiology, 2010, 33, 116-121.	2.8	23
125	Pseudomonas arsenicoxydans sp nov., an arsenite-oxidizing strain isolated from the Atacama desert. Systematic and Applied Microbiology, 2010, 33, 193-197.	2.8	54
126	Breoghania corrubedonensis gen. nov. sp. nov., a novel alphaproteobacterium isolated from a Galician beach (NW Spain) after the Prestige fuel oil spill, and emended description of the family Cohaesibacteraceae and the species Cohaesibacter gelatinilyticus. Systematic and Applied Microbiology, 2010, 33, 316-321.	2.8	25

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127	Update of the All-Species Living Tree Project based on 16S and 23S rRNA sequence analyses. Systematic and Applied Microbiology, 2010, 33, 291-299.	2.8	441
128	Bioaugmentation with Pseudomonas $\hat{A} \in \hat{A} \in \hat{A} $ sp. strain MHP41 promotes simazine attenuation and bacterial community changes in agricultural soils. FEMS Microbiology Ecology, 2010, 71, 114-126.	2.7	56
129	Bioaugmentation with <i>Pseudomonas</i> sp. strain MHP41 promotes simazine attenuation and bacterial community changes in agricultural soils. FEMS Microbiology Ecology, 2010, 72, 152-152.	2.7	2
130	Fine-scale evolution: genomic, phenotypic and ecological differentiation in two coexisting <i>Salinibacter ruber</i> strains. ISME Journal, 2010, 4, 882-895.	9.8	81
131	Labrys wisconsinensis sp. nov., a budding bacterium isolated from Lake Michigan water, and emended description of the genus Labrys. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 1570-1576.	1.7	17
132	Determination of the Diversity of <i>Rhodopirellula</i> Isolates from European Seas by Multilocus Sequence Analysis. Applied and Environmental Microbiology, 2010, 76, 776-785.	3.1	32
133	Notes on the characterization of prokaryote strains for taxonomic purposes. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 249-266.	1.7	1,232
134	Extremely halophilic microbial communities in anaerobic sediments from a solar saltern. Environmental Microbiology Reports, 2010, 2, 258-271.	2.4	44
135	Shifting the genomic gold standard for the prokaryotic species definition. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 19126-19131.	7.1	5,234
136	Lactobacillus oeni sp. nov., from wine. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 2010-2014.	1.7	31
137	Occurrence of Halococcus spp. in the nostrils salt glands of the seabird Calonectris diomedea. Extremophiles, 2009, 13, 557-565.	2.3	33
138	Automated microextraction sample preparation coupled on-line to FT-ICR-MS: application to desalting and concentration of river and marine dissolved organic matter. Analytical and Bioanalytical Chemistry, 2009, 395, 797-807.	3.7	34
139	Evaluation of the 18S rRNA clone library approach to study the diversity of the macroeukaryotic leaf-epiphytic community of the seagrass Posidonia oceanica (L.) Delile. Marine Biology, 2009, 156, 1963-1976.	1.5	6
140	Combining chip-ESI with APLI (cESILI) as a multimode source for analysis of complex mixtures with ultrahigh-resolution mass spectrometry. Analytical and Bioanalytical Chemistry, 2008, 391, 2803-2809.	3.7	45
141	Application of temperature gradient gel electrophoresis technique to monitor changes in the structure of the eukaryotic leaf-epiphytic community of Posidonia oceanica. Marine Biology, 2008, 155, 451-460.	1.5	2
142	Identification of a bacterial strain isolated from the liver of a laboratory mouse as Microbacterium paraoxydans and emended description of the species Microbacterium paraoxydans Laffineur et al 2003. Indian Journal of Microbiology, 2008, 48, 243-251.	2.7	19
143	Metabolic evidence for biogeographic isolation of the extremophilic bacterium <i>Salinibacter ruber</i> . ISME Journal, 2008, 2, 242-253.	9.8	108
144	The All-Species Living Tree project: A 16S rRNA-based phylogenetic tree of all sequenced type strains. Systematic and Applied Microbiology, 2008, 31, 241-250.	2.8	884

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145	Reclassification of Rhodobium marinum and Rhodobium pfennigii as Afifella marina gen. nov. comb. nov. and Afifella pfennigii comb. nov., a new genus of photoheterotrophic Alphaproteobacteria and emended descriptions of Rhodobium, Rhodobium orientis and Rhodobium gokarnense. Systematic and Applied Microbiology, 2008, 31, 339-351.	2.8	111
146	Lactobacillus uvarum sp. nov. – A new lactic acid bacterium isolated from Spanish Bobal grape must. Systematic and Applied Microbiology, 2008, 31, 425-433.	2.8	26
147	Distribution, abundance and diversity of the extremely halophilic bacterium Salinibacter ruber. Saline Systems, 2008, 4, 15.	2.0	54
148	Undibacterium pigrum gen. nov., sp. nov., isolated from drinking water. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 1510-1515.	1.7	73
149	Proposal of Viridibacillus gen. nov. and reclassification of Bacillus arvi, Bacillus arenosi and Bacillus neidei as Viridibacillus arvi gen. nov., comb. nov., Viridibacillus arenosi comb. nov. and Viridibacillus neidei comb. nov International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 2729-2737.	1.7	69
150	Metagenomic approach to the study of halophages: the environmental halophage 1. Environmental Microbiology, 2007, 9, 1711-1723.	3.8	59
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