

Peter Kampfer

List of Publications by Year in descending order

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432
papers

20,820
citations

17776

65
h-index

23173

116
g-index

487
all docs

487
docs citations

487
times ranked

15134
citing authors

#	ARTICLE	IF	CITATIONS
1	A Preliminary Comparison on Faecal Microbiomes of Free-Ranging Large Baleen (Balaenoptera) Tj ETQq1 1 0.784314 rgBT /Overlock 10 2022, 83, 18-33.	1.4	3
2	Isolation of <i>Hermetia illucens</i> larvae core gut microbiota by two different cultivation strategies. <i>Antonie Van Leeuwenhoek</i> , 2022, 115, 821-837.	0.7	4
3	<i>Pseudoneobacillus rhizosphaerae</i> gen. nov., sp. nov., isolated from maize root rhizosphere. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	0.8	10
4	<i>Ottowia testudinis</i> sp. nov., isolated from the cloaca of a giant Asian pond turtle (<i>Heosemys grandis</i>). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	0.8	5
5	<i>Erysipelothrix anatis</i> sp. nov., <i>Erysipelothrix aquatica</i> sp. nov. and <i>Erysipelothrix urinaevulpis</i> sp. nov., three novel species of the genus, and emended description of <i>Erysipelothrix</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	0.8	20
6	<i>Bacillus rhizoplanae</i> sp. nov. from maize roots. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2022, 72, .	0.8	4
7	<i>Acinetobacter stercoris</i> sp. nov. isolated from output source of a mesophilic german biogas plant with anaerobic operating conditions. <i>Antonie Van Leeuwenhoek</i> , 2021, 114, 235-251.	0.7	12
8	<i>Streptomyces bathyalis</i> sp. nov., an actinobacterium isolated from the sponge in a deep sea. <i>Antonie Van Leeuwenhoek</i> , 2021, 114, 425-435.	0.7	22
9	<i>Pseudomonas carbonaria</i> sp. nov., isolated from charcoal. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	6
10	Description and comparison of the skin and ear canal microbiota of non-allergic and allergic German shepherd dogs using next generation sequencing. <i>PLoS ONE</i> , 2021, 16, e0250695.	1.1	9
11	Description of three new <i>Alteromonas</i> species <i>Alteromonas antoniana</i> sp. nov., <i>Alteromonas lipotrueae</i> sp. nov. and <i>Alteromonas lipotrueiana</i> sp. nov. isolated from marine environments, and proposal for reclassification of the genus <i>Salinimonas</i> as <i>Alteromonas</i> . <i>Systematic and Applied Microbiology</i> , 2021, 44, 126226.	1.2	39
12	<i>Streptomonospora litoralis</i> sp. nov., a halophilic thiopeptides producer isolated from sand collected at Cuxhaven beach. <i>Antonie Van Leeuwenhoek</i> , 2021, 114, 1483-1496.	0.7	6
13	<i>Entomomonas asaccharolytica</i> sp. nov., isolated from <i>Acheta domesticus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	6
14	<i>Flavobacterium bizetiae</i> sp. nov., isolated from diseased freshwater fish in Canada at the end of the 1970s. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	9
15	<i>Paenibacillus allorhizosphaerae</i> sp. nov., from soil of the rhizosphere of <i>Zea mays</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	11
16	<i>Luteolibacter ambystomatis</i> sp. nov., isolated from the skin of an Anderson's salamander (<i>Ambystoma</i>) Tj ETQq0 0 0 rgBT /Overlock 9	0.8	9
17	Comparative genome analysis of the genus <i>Hydrotalea</i> and proposal of the novel species <i>Hydrotalea lipotrueae</i> sp. nov., isolated from a groundwater aquifer in the south of Mallorca Island, Spain. <i>Systematic and Applied Microbiology</i> , 2021, 44, 126277.	1.2	7
18	<i>Devosia equisanguinis</i> sp. nov., isolated from horse blood. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	9

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19	<i>Paralysiella testudinis</i> gen. nov., sp. nov., isolated from the cloaca of a toad-headed turtle (<i>Mesoclemmys nasuta</i>). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	9
20	<i>Leucobacter soli</i> sp. nov., from soil amended with humic acid. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	0.8	6
21	<i>Zooshikella harenae</i> sp. nov., Isolated from Pacific Oyster <i>Crassostrea gigas</i> , and Establishment of <i>Zooshikella ganghwensis</i> subsp. marina subsp. nov. and <i>Zooshikella ganghwensis</i> subsp. ganghwensis subsp. nov.. <i>Diversity</i> , 2021, 13, 641.	0.7	15
22	Epidemiological analysis of <i>Trueperella abortus</i> isolated from cases of pig abortion of a single farm. <i>Folia Microbiologica</i> , 2020, 65, 491-496.	1.1	10
23	Endophytic bacterial communities of oilseed rape associate with genotype-specific resistance against <i>Verticillium longisporum</i> . <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	8
24	High diversity of <i>Vibrio</i> spp. associated with different ecological niches in a marine aquaria system and description of <i>Vibrio aquimaris</i> sp. nov. <i>Systematic and Applied Microbiology</i> , 2020, 43, 126123.	1.2	14
25	<i>Taxonomic study of nine new Winogradskyella species occurring in the shallow waters of Helgoland Roads, North Sea. Proposal of Winogradskyella schleiferi</i> sp. nov., <i>Winogradskyella costae</i> sp. nov., <i>Winogradskyella helgolandensis</i> sp. nov., <i>Winogradskyella vidalii</i> sp. nov., <i>Winogradskyella forsetii</i> sp. nov., <i>Winogradskyella ludwigii</i> sp. nov., <i>Winogradskyella ursingii</i> sp. nov., <i>Winogradskyella wichelsiae</i> sp. nov., and <i>Candidatus Winogradskyella atlantica</i> sp. nov.. <i>Systematic and Applied Microbiology</i> , 2020, 43, 126128.	1.2	38
26	<i>Acinetobacter baumannii</i> in manure and anaerobic digestates of German biogas plants. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	19
27	Global warming shifts the composition of the abundant bacterial phyllosphere microbiota as indicated by a cultivation-dependent and -independent study of the grassland phyllosphere of a long-term warming field experiment. <i>FEMS Microbiology Ecology</i> , 2020, 96, .	1.3	21
28	Roadmap for naming uncultivated Archaea and Bacteria. <i>Nature Microbiology</i> , 2020, 5, 987-994.	5.9	115
29	The Bacterial Microbiome of the Long-Term Aquarium Cultured High-Microbial Abundance Sponge <i>Haliclona nidata</i> Sustained Bioactivity Despite Community Shifts Under Detrimental Conditions. <i>Frontiers in Marine Science</i> , 2020, 7, .	1.2	12
30	The gut and feed residue microbiota changing during the rearing of <i>Hermetia illucens</i> larvae. <i>Antonie Van Leeuwenhoek</i> , 2020, 113, 1323-1344.	0.7	33
31	<i>Corynebacterium endometrii</i> sp. nov., isolated from the uterus of a cow with endometritis. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 146-152.	0.8	12
32	<i>Psychrobacter pygoscelis</i> sp. nov. isolated from the penguin <i>Pygoscelis papua</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 211-219.	0.8	14
33	Taxonomic reassessment of the genus <i>Pseudocitrobacter</i> using whole genome sequencing: <i>Pseudocitrobacter anthropi</i> is a later heterotypic synonym of <i>Pseudocitrobacter faecalis</i> and description of <i>Pseudocitrobacter vendiensis</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 4432-4450.	0.8	14
34	Observation of discontinuities in amino acid identity values, a possible consequence of major extinction events, guides transfer of nine species to the genus <i>Epilithonimonas</i> , eleven species to the genus <i>Kaistella</i> , and three species to the genus <i>Halpernia</i> gen. nov., with description of <i>Kaistella daneshvariae</i> sp. nov. and <i>Epilithonimonas vandammei</i> sp. nov. derived from clinical specimens. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 4432-4450.	0.8	215
35	Proposal to reclassify <i>Leptotrichia goodfellowii</i> into a novel genus as <i>Pseudoleptotrichia goodfellowii</i> gen. nov., comb. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 2084-2088.	0.8	11
36	Proposal to reclassify <i>Streptobacillus hongkongensis</i> into a novel genus as <i>Pseudostreptobacillus hongkongensis</i> gen. nov., comb. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 2366-2368.	0.8	12

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37	<i>Streptobacillus canis</i> sp. nov. isolated from a dog. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 2648-2656.	0.8	14
38	<i>Corynebacterium silvaticum</i> sp. nov., a unique group of NTTB corynebacteria in wild boar and roe deer. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 3614-3624.	0.8	36
39	<i>Corynebacterium urogenitale</i> sp. nov. isolated from the genital tract of a cow. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 3625-3632.	0.8	9
40	The status of the genus <i>Prolinoborus</i> (Pot et al. 1992) and the species <i>Prolinoborus fasciculus</i> (Pot et) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54	0.8	8
41	<i>Flavobacterium salmonis</i> sp. nov. isolated from Atlantic salmon (<i>Salmo salar</i>) and formal proposal to reclassify <i>Flavobacterium spartansii</i> as a later heterotypic synonym of <i>Flavobacterium tractae</i> . International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 6147-6154.	0.8	15
42	<i>Arthrobacter ulcerisalmonis</i> sp. nov., isolated from an ulcer of a farmed Atlantic salmon (<i>Salmo</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54 Systematic and Evolutionary Microbiology, 2020, 70, 1963-1968.	0.8	11
43	<i>Arcanobacterium bovis</i> sp. nov., isolated from the milk of a cow with mastitis. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 4105-4110.	0.8	9
44	<i>Flavobacterium panici</i> sp. nov. isolated from the rhizosphere of the switchgrass <i>Panicum virgatum</i> . International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 5824-5831.	0.8	7
45	Proposal of <i>Pedobacter nototheniae</i> sp. nov., isolated from the spleen of a black rock cod (<i>Notothenia coriiceps</i> , Richardson 1844) from the Chilean Antarctica. Antonie Van Leeuwenhoek, 2019, 112, 1465-1475.	0.7	10
46	Karamomycins Aâ€“C: 2-Naphthalen-2-yl-thiazoles from <i>Nonomuraea endophytica</i> . Journal of Natural Products, 2019, 82, 870-877.	1.5	18
47	Proposal of <i>Lysobacter pythonis</i> sp. nov. isolated from royal pythons (<i>Python regius</i>). Systematic and Applied Microbiology, 2019, 42, 326-333.	1.2	10
48	The Precursor Hypothesis of Sponge Kleptocnidism: Development of Nematocysts in <i>Haliclona cnidata</i> sp. nov. (Porifera, Demospongiae, Haplosclerida). Frontiers in Marine Science, 2019, 5, .	1.2	7
49	<i>Aromatoleum</i> gen. nov., a novel genus accommodating the phylogenetic lineage including <i>Azoarcus evansii</i> and related species, and proposal of <i>Aromatoleum aromaticum</i> sp. nov., <i>Aromatoleum petrolei</i> sp. nov., <i>Aromatoleum bremense</i> sp. nov., <i>Aromatoleum toluolicum</i> sp. nov. and <i>Aromatoleum diolicum</i> sp. nov.. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 982-997.	0.8	88
50	Classification of three corynebacterial strains isolated from the Northern Bald Ibis (<i>Geronticus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54 <i>Corynebacterium gerontici</i> sp. nov.. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 2928-2935.	0.8	21
51	<i>Sneathia vaginalis</i> sp. nov. (Fusobacteriales, Leptotrichiaceae) as a replacement of the species â€“ <i>Sneathia amnii</i> â€™ Harwich et al. 2012 and â€“ <i>Leptotrichia amnionii</i> â€™ Shukla et al. 2002, and emended description of <i>Sneathia</i> Collins et al. 2001. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	0.8	17
52	<i>Filibacter tadaridae</i> sp. nov., isolated from within a guano pile from a colony of Mexican free-tailed bats <i>Tadarida brasiliensis</i> . International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1438-1442.	0.8	6
53	<i>Paracoccus haematequi</i> sp. nov., isolated from horse blood. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1682-1688.	0.8	9
54	<i>Jeotgalicoccus meleagridis</i> sp. nov. isolated from bioaerosol from emissions of a turkey fattening plant and reclassification of <i>Jeotgalicoccus halophilus</i> Liu et al. 2011 as a later heterotypic synonym of <i>Jeotgalicoccus aerolatus</i> Martin et al. 2011. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	0.8	12

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55	<i>Streptomyces dysideae</i> sp. nov., isolated from a marine Mediterranean sponge <i>Dysidea tupa</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 71, .	0.8	6
56	<i>Pigmentiphaga humi</i> sp. nov., isolated from soil amended with humic acid. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 1573-1578.	0.8	4
57	<i>Xinfangfangia humi</i> sp. nov., isolated from soil amended with humic acid. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 2070-2075.	0.8	10
58	<i>Paracoccus nototherniae</i> sp. nov., isolated from a black rock cod fish (<i>Notothernia coriiceps</i>) from the Chilean Antarctic. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 2794-2800.	0.8	9
59	<i>Parendozoicomonas haliclona</i> gen. nov. sp. nov. isolated from a marine sponge of the genus <i>Haliclona</i> and description of the family <i>Endozoicomonadaceae</i> fam. nov. comprising the genera <i>Endozoicomonas</i> , <i>Parendozoicomonas</i> , and <i>Kistimonas</i> . <i>Systematic and Applied Microbiology</i> , 2018, 41, 73-84.	1.2	48
60	Long-Term Warming Shifts the Composition of Bacterial Communities in the Phyllosphere of <i>Galium album</i> in a Permanent Grassland Field-Experiment. <i>Frontiers in Microbiology</i> , 2018, 9, 144.	1.5	76
61	Taxonomic analyses of members of the <i>Streptomyces cinnabarinus</i> cluster, description of <i>Streptomyces cinnabarinus</i> sp. nov. and <i>Streptomyces davaonensis</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 382-393.	0.8	26
62	<i>Nannocystis konarekensis</i> sp. nov., a novel myxobacterium from an Iranian desert. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 721-729.	0.8	21
63	Proposal of <i>Litorimonas haliclona</i> sp. nov., isolated from a marine sponge of the genus <i>Haliclona</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 835-843.	0.8	22
64	<i>Marinicrinis lubricantis</i> sp. nov., isolated from a coolant lubricant. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1018-1022.	0.8	6
65	<i>Halomonas malpeensis</i> sp. nov., isolated from rhizosphere sand of a coastal sand dune plant. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1037-1046.	0.8	19
66	<i>Comamonas aquatilis</i> sp. nov., isolated from a garden pond. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1210-1214.	0.8	15
67	<i>Novosphingobium lubricantis</i> sp. nov., isolated from a coolant lubricant emulsion. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1560-1564.	0.8	18
68	Proposal for a new classification of a deep branching bacterial phylogenetic lineage: transfer of <i>Coprothermobacter proteolyticus</i> and <i>Coprothermobacter platensis</i> to <i>Coprothermobacteraceae</i> fam. nov., within <i>Coprothermobacterales</i> ord. nov., <i>Coprothermobacteria classis</i> nov. and <i>Coprothermobacterota</i> phyl. nov. and emended description of the family <i>Thermodesulfobiaceae</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1627-1632.	0.8	46
69	<i>Winogradskyella pocilloporae</i> sp. nov. isolated from healthy tissue of the coral <i>Pocillopora damicornis</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1689-1696.	0.8	17
70	<i>Stenotrophomonas lactitubi</i> sp. nov. and <i>Stenotrophomonas indicatrix</i> sp. nov., isolated from surfaces with food contact. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1830-1838.	0.8	28
71	<i>Lelliottia aquatilis</i> sp. nov., isolated from drinking water. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 2454-2461.	0.8	23
72	A polyphasic approach leads to seven new species of the cellulose-decomposing genus <i>Sorangium</i> , <i>Sorangium ambruticinum</i> sp. nov., <i>Sorangium arenae</i> sp. nov., <i>Sorangium bulgaricum</i> sp. nov., <i>Sorangium dawidii</i> sp. nov., <i>Sorangium kenyense</i> sp. nov., <i>Sorangium orientale</i> sp. nov. and <i>Sorangium reichenbachii</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3576-3586.	0.8	46

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73	<i>Streptomyces ciscaucasicus</i> Sveshnikova et al. 1983 is a later subjective synonym of <i>Streptomyces canus</i> Heinemann et al. 1953. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 42-46.	0.8	4
74	<i>Qingshengfania soli</i> Zhang et al. 2015 is a later heterotypic synonym of <i>Pseudochelatococcus lubricantis</i> Kämpfer et al. 2015. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 2240-2241.	0.8	2
75	<i>Chromobacterium pseudoviolaceum</i> Kämpfer et al. 2009 is a later heterotypic synonym of <i>Chromobacterium violaceum</i> Bergonzini 1880. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 2967-2968.	0.8	6
76	Detection and Characterization of Endobacteria in the Fungal Endophyte <i>Piriformospora indica</i> . , 2017, , 237-250.		1
77	Genome Sequence of <i>Paracoccus contaminans</i> LMG 29738 T , Isolated from a Water Microcosm. <i>Genome Announcements</i> , 2017, 5, .	0.8	12
78	The Abundance of Endofungal Bacterium <i>Rhizobium radiobacter</i> (syn. <i>Agrobacterium tumefaciens</i>) Increases in Its Fungal Host <i>Piriformospora indica</i> during the Tripartite Sebacinalean Symbiosis with Higher Plants. <i>Frontiers in Microbiology</i> , 2017, 8, 629.	1.5	54
79	Fatal infection in three Grey Slender Lorises (<i>Loris lydekkerianus nordicus</i>) caused by clonally related <i>Trueperella pyogenes</i> . <i>BMC Veterinary Research</i> , 2017, 13, 273.	0.7	12
80	“ <i>Streptomyces caelicus</i> ”™, an antibiotic-producing species of the genus <i>Streptomyces</i> , and <i>Streptomyces canchipurensis</i> Li et al. 2015 are later heterotypic synonyms of <i>Streptomyces muensis</i> Ningthoujam et al. 2014. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 548-556.	0.8	14
81	<i>Cohnella lubricantis</i> sp. nov., isolated from a coolant lubricant solution. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 466-471.	0.8	16
82	<i>Racemicystis persica</i> sp. nov., a myxobacterium from soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 472-478.	0.8	10
83	<i>Photorhabdus luminescens</i> subsp. <i>namnaonensis</i> subsp. nov., isolated from <i>Heterorhabditis baujardi</i> nematodes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1046-1051.	0.8	24
84	<i>Xenorhabdus thuongxuanensis</i> sp. nov. and <i>Xenorhabdus eapokensis</i> sp. nov., isolated from <i>Steinernema</i> species. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1107-1114.	0.8	26
85	<i>Paenibacillus rhizoplanae</i> sp. nov., isolated from the rhizosphere of <i>Zea mays</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1058-1063.	0.8	23
86	<i>Arcanobacterium wilhelmae</i> sp. nov., isolated from the genital tract of a rhinoceros (<i>Rhinoceros</i>) Tj ETQq0 0 0 rgBT /Qverlock 10 Tf 50 2.	0.8	12
87	<i>Bacillus zeae</i> sp. nov., isolated from the rhizosphere of <i>Zea mays</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1241-1246.	0.8	9
88	<i>Psychromonas aquatilis</i> sp. nov., isolated from seawater samples obtained in the Chilean Antarctica. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1306-1311.	0.8	11
89	Proposal of <i>Mucilagibacter galii</i> sp. nov. isolated from leaves of <i>Galium album</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1318-1326.	0.8	11
90	<i>Paracandidimonas soli</i> gen. nov., sp. nov., isolated from soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 1740-1745.	0.8	14

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91	<i>Stenotrophomonas bentonitica</i> sp. nov., isolated from bentonite formations. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2779-2786.	0.8	31
92	<i>Streptomyces phaeopurpureus</i> Shinobu 1957 (Approved Lists 1980) and <i>Streptomyces griseorubiginosus</i> (Ryabova and Preobrazhenskaya 1957) Pridham et al. 1958 (Approved Lists 1980) are heterotypic subjective synonyms. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3111-3116.	0.8	5
93	<i>Flavobacterium gossypii</i> sp. nov. isolated from the root tissue of field-grown cotton. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3345-3350.	0.8	7
94	<i>Winogradskyella haliclona</i> sp. nov., isolated from a marine sponge of the genus <i>Haliclona</i> . International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 4902-4910.	0.8	22
95	<i>Paenibacillus nebraskensis</i> sp. nov., isolated from the root surface of field-grown maize. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 4956-4961.	0.8	12
96	Cultivation of vancomycin-resistant enterococci and methicillin-resistant staphylococci from input and output samples of German biogas plants. FEMS Microbiology Ecology, 2016, 92, fiw010.	1.3	9
97	<i>Varibaculum anthropi</i> sp. nov. represented by three genetically different genomovars isolated from clinical material and emended description of the genus <i>Varibaculum</i> . Systematic and Applied Microbiology, 2016, 39, 546-552.	1.2	15
98	Bacterial diversity and antibiotic resistances of abundant aerobic culturable bacteria in input and output samples of 15 German biogas plants. Journal of Applied Microbiology, 2016, 121, 1673-1684.	1.4	15
99	Circulation of clonal populations of fluoroquinolone-resistant CTX-M-15-producing <i>Escherichia coli</i> ST410 in humans and animals in Germany. International Journal of Antimicrobial Agents, 2016, 47, 457-465.	1.1	107
100	Draft genome of the <i>Arabidopsis thaliana</i> phyllosphere bacterium, <i>Williamsia</i> sp. ARP1. Standards in Genomic Sciences, 2016, 11, 8.	1.5	10
101	<i>Kosakonia pseudosacchari</i> sp. nov., an endophyte of <i>Zea mays</i> . Systematic and Applied Microbiology, 2016, 39, 1-7.	1.2	30
102	Non-pathogenic <i>Rhizobium radiobacter</i> F4 deploys plant beneficial activity independent of its host <i>Pisiformospora indica</i> . ISME Journal, 2016, 10, 871-884.	4.4	93
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104	The status of the species <i>Enterobacter siamensis</i> Khunthongpan et al. 2014 . Request for an Opinion. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 524-525.	0.8	8
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122	<i>Pseudoduganella danionis</i> sp. nov., isolated from zebrafish (<i>Danio rerio</i>). International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4671-4675.	0.8	17
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221	<i>Novosphingobium soli</i> sp. nov., isolated from soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 259-263.	0.8	39
222	<i>Chryseobacterium chaponense</i> sp. nov., isolated from farmed Atlantic salmon (<i>Salmo salar</i>). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 497-501.	0.8	50
223	<i>Ochrobactrum pecoris</i> sp. nov., isolated from farm animals. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2011, 61, 2278-2283.	0.8	49
224	Detection of <i>Saccharopolyspora rectivirgula</i> by Quantitative Real-Time PCR. <i>Annals of Occupational Hygiene</i> , 2011, 55, 612-9.	1.9	13
225	The characterization of prokaryote strains for taxonomic purposes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 7-7.	0.8	8
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228	Detection of <i>Jeotgalicoccus</i> spp. in poultry house air. <i>Systematic and Applied Microbiology</i> , 2010, 33, 188-192.	1.2	19
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232	<i>Chryseobacterium arthrosphaerae</i> sp. nov., isolated from the faeces of the pill millipede <i>Arthrosphaera magna</i> Attems. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1765-1769.	0.8	42
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236	<i>Ochrobactrum pituitosum</i> sp. nov., isolated from an industrial environment. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 321-326.	0.8	48
237	<i>Citricoccus parietis</i> sp. nov., isolated from a mould-colonized wall and emended description of <i>Citricoccus alkalitolerans</i> Li et al. 2005. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 271-274.	0.8	17
238	<i>Brucella inopinata</i> sp. nov., isolated from a breast implant infection. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 801-808.	0.8	276
239	<i>Paenochrobactrum gallinarii</i> gen. nov., sp. nov., isolated from air of a duck barn, and reclassification of <i>Pseudochrobactrum glaciei</i> as <i>Paenochrobactrum glaciei</i> comb. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2010, 60, 1493-1498.	0.8	30
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251	<i>Algoriphagus olei</i> sp. nov., isolated from oil-contaminated soil. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 2909-2915.	0.8	42
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254	<i>Corynebacterium lubricantis</i> sp. nov., isolated from a coolant lubricant. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 1112-1115.	0.8	23
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257	Description of <i>Chryseobacterium anthropi</i> sp. nov. to accommodate clinical isolates biochemically similar to <i>Kaistella koreensis</i> and <i>Chryseobacterium haifense</i> , proposal to reclassify <i>Kaistella koreensis</i> as <i>Chryseobacterium koreense</i> comb. nov. and emended description of the genus <i>Chryseobacterium</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 2421-2428.	0.8	95
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282	<i>Deinococcus aquatilis</i> sp. nov., isolated from water. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 2803-2806.	0.8	53
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