Peter A Vandamme

List of Publications by Year
in descending order

613 papers	52,996 citations	1981 104 h-index	3037 194 g-index
635 all docs	635 docs citations	635 times ranked	32782 citing authors

#	Article	IF	Citations
1	Outbreak of Burkholderia contaminans endophthalmitis traced to a clinic ventilation system. Infection Control and Hospital Epidemiology, 2022, 43, 1705-1707.	1.0	2
2	Co-cultivation enhanced microbial protein production based on autotrophic nitrogen-fixing hydrogen-oxidizing bacteria. Chemical Engineering Journal, 2022, 429, 132535.	6.6	16
3	Dominance of honey bees is negatively associated with wild bee diversity in commercial apple orchards regardless of management practices. Agriculture, Ecosystems and Environment, 2022, 323, 107697.	2.5	25
4	Description of Pseudoclavibacter triregionum sp. nov. from human blood and Pseudoclavibacter albus comb. nov., and revised classification of the genus Pseudoclavibacter: proposal of Caespitibacter gen. nov., with Caespitibacter soli comb. nov. and Caespitibacter caeni comb. nov. Antonie Van Leeuwenhoek, 2022, 115, 461-472.	0.7	23
5	A comparative analysis of crop pollinator survey methods along a large-scale climatic gradient. Agriculture, Ecosystems and Environment, 2022, 329, 107871.	2.5	10
6	Accurate prediction of metagenome-assembled genome completeness by MAGISTA, a random forest model built on alignment-free intra-bin statistics. Environmental Microbiomes, 2022, 17, 9.	2.2	3
7	Paraburkholderia gardini sp. nov. and Paraburkholderia saeva sp. nov.: Novel aromatic compound degrading bacteria isolated from garden and forest soil samples. Systematic and Applied Microbiology, 2022, 45, 126318.	1.2	11
8	Backslopping Time, Rinsing of the Grains During Backslopping, and Incubation Temperature Influence the Water Kefir Fermentation Process. Frontiers in Microbiology, 2022, 13, .	1.5	4
9	Ecological, environmental, and management data indicate apple production is driven by wild bee diversity and management practices. Ecological Indicators, 2022, 139, 108880.	2.6	13
10	Phylogenomic Analyses of Snodgrassella Isolates from Honeybees and Bumblebees Reveal Taxonomic and Functional Diversity. MSystems, 2022, 7, .	1.7	19
11	A comprehensive investigation into the production of gamma-aminobutyric acid by Limosilactobacillus fermentum NG16, a tuna gut isolate. Acta Alimentaria, 2022, 51, 302-311.	0.3	0
12	Introduction to the principles and methods underlying the recovery of metagenomeâ€assembled genomes from metagenomic data. MicrobiologyOpen, 2022, 11, .	1.2	8
13	Microbial diversity and antimicrobial susceptibility in endotracheal tube biofilms recovered from mechanically ventilated COVID-19 patients. Biofilm, 2022, 4, 100079.	1.5	9
14	Development of a database for the rapid and accurate routine identification of Achromobacter species by matrix-assisted laser desorption/ionizationâ€“time-of-flight mass spectrometry (MALDI-TOF MS). Clinical Microbiology and Infection, 2021, 27, 126.e1-126.e5.	2.8	15
15	gcType: a high-quality type strain genome database for microbial phylogenetic and functional research. Nucleic Acids Research, 2021, 49, D694-D705.	6.5	53
16	Diagnostic approach for detection and identification of emerging enteric pathogens revisited: the (Ali)arcobacter lanthieri case. New Microbes and New Infections, 2021, 39, 100829.	0.8	11
17	Induction of antibiotic specialized metabolism by coâ€culturing in a collection of phyllosphere bacteria. Environmental Microbiology, 2021, 23, 2132-2151.	1.8	12
18	Network Analysis Based on Unique Spectral Features Enables an Efficient Selection of Genomically Diverse Operational Isolation Units. Microorganisms, 2021, 9, 416.	1.6	4

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19	Burkholderia Bacteria Produce Multiple Potentially Novel Molecules that Inhibit Carbapenem-Resistant Gram-Negative Bacterial Pathogens. Antibiotics, 2021, 10, 147.	1.5	12
20	Presence of the Weakly Pathogenic Fusarium poae in the Fusarium Head Blight Disease Complex Hampers Biocontrol and Chemical Control of the Virulent Fusarium graminearum Pathogen. Frontiers in Plant Science, 2021, 12, 641890.	1.7	17
21	The Type and Concentration of Inoculum and Substrate as Well as the Presence of Oxygen Impact the Water Kefir Fermentation Process. Frontiers in Microbiology, 2021, 12, 628599.	1.5	17
22	Genomics of an endemic cystic fibrosis Burkholderia multivorans strain reveals low within-patient evolution but high between-patient diversity. PLoS Pathogens, 2021, 17, e1009418.	2.1	11
23	The bacterial diversity of raw Moroccon camel milk. International Journal of Food Microbiology, 2021, 341, 109050.	2.1	16
24	Paenibacillus foliorum sp. nov., Paenibacillus phytohabitans sp. nov., Paenibacillus plantarum sp. nov., Paenibacillus planticolens sp. nov., Paenibacillus phytorum sp. nov. and Paenibacillus germinis sp. nov., isolated from the Arabidopsis thaliana phyllosphere. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	31
25	Clinical course, treatment and visual outcome of an outbreak of Burkholderia contaminans endophthalmitis following cataract surgery. Journal of Ophthalmic Inflammation and Infection, 2021, 11, 12.	1.2	6
26	MALDI-TOF MS insight into the biodiversity of Staka, the artisanal Cretan soured cream. International Dairy Journal, 2021, 116, 104969.	1.5	8
27	Cutibacterium modestum and â€œPropionibacterium humerusiiâ€•represent the same species that is commonly misidentified as Cutibacterium acnes. Antonie Van Leeuwenhoek, 2021, 114, 1315-1320.	0.7	4
28	Bumblebee resilience to climate change, through plastic and adaptive responses. Global Change Biology, 2021, 27, 4223-4237.	4.2	49
29	Genomic Aromatic Compound Degradation Potential of Novel Paraburkholderia Species: Paraburkholderia domus sp. nov., Paraburkholderia haematera sp. nov. and Paraburkholderia nemoris sp. nov.. International Journal of Molecular Sciences, 2021, 22, 7003.	1.8	22
30	Impact of intraspecific variation on measurements of thermal tolerance in bumble bees. Journal of Thermal Biology, 2021, 99, 103002.	1.1	17
31	Burkholderia perseverans sp. nov., a bacterium isolated from the Restinga ecosystem, is a producer of volatile and diffusible compounds that inhibit plant pathogens. Brazilian Journal of Microbiology, 2021, 52, 2145-2152.	0.8	4
32	The Ever-Expanding Pseudomonas Genus: Description of 43 New Species and Partition of the Pseudomonas putida Group. Microorganisms, 2021, 9, 1766.	1.6	206
33	Paraburkholderia dioscoreae sp. nov., a novel plant associated growth promotor. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	11
34	Gulosibacter hominis sp. nov.: a novel human microbiome bacterium that may cause opportunistic infections. Antonie Van Leeuwenhoek, 2021, 114, 1841-1854.	0.7	10
35	Biosynthesis of Ditropolonyl Sulfide, an Antibacterial Compound Produced by Burkholderia cepacia Complex Strain R-12632. Applied and Environmental Microbiology, 2021, 87, e0116921.	1.4	4
36	Matrix-Assisted Laser Desorption Ionizationâ€“Time of Flight Mass Spectrometry for Rapid Detection of Isolates Belonging to the Epidemic Clones Achromobacter xylosoxidans ST137 and Achromobacter ruhlandii DES from Cystic Fibrosis Patients. Journal of Clinical Microbiology, 2021, 59, e0094621.	1.8	4

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37	Identification and characterization of acid-tolerant spore-forming spoilage bacteria from acidified and low-acid pasteurized sauces. LWT - Food Science and Technology, 2021, 152, 112378.	2.5	8
38	Gluconacetobacter dulcium sp. nov., a novel Gluconacetobacter species from sugar-rich environments. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	5
39	Optimization of culture conditions for gamma-aminobutyric acid production by newly identified Pediococcus pentosaceus MN12 isolated from â€˜mam nemâ€™, a fermented fish sauce. Bioengineered, 2021, 12, 54-62.	1.4	14
40	The Holobiont as a Key to the Adaptation and Conservation of Wild Bees in the Anthropocene. Frontiers in Ecology and Evolution, 2021, 9, .	1.1	12
41	Out with the old and in with the new: time to rethink twentieth century chemotaxonomic practices in bacterial taxonomy. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	17
42	Aliarcobacter, Halarcobacter, Malaciobacter, Pseudarcobacter and Poseidonibacter are later synonyms of Arcobacter: transfer of Poseidonibacter parvus, Poseidonibacter antarcticus, â€˜Halarcobacter arenosusâ€™, and â€˜Aliarcobacter vitoriensisâ€™ to Arcobacter as Arcobacter parvus comb. nov., Arcobacter antarcticus comb. nov., Arcobacter arenosus comb. nov. and Arcobacter vitoriensis comb. nov.. International Journal of Systematic and Evolutionary Microbiology, 2021, 71, .	0.8	30
43	Bacterial species identification using MALDI-TOF mass spectrometry and machine learning techniques: A large-scale benchmarking study. Computational and Structural Biotechnology Journal, 2021, 19, 6157-6168.	1.9	20
44	PaSiT: a novel approach based on short-oligonucleotide frequencies for efficient bacterial identification and typing. Bioinformatics, 2020, 36, 2337-2344.	1.8	5
45	Description of Komagataeibacter melaceti sp. nov. and Komagataeibacter melomenusus sp. nov. Isolated from Apple Cider Vinegar. Microorganisms, 2020, 8, 1178.	1.6	34
46	Burkholderia cepacia Complex Taxon K: Where to Split?. Frontiers in Microbiology, 2020, 11, 1594.	1.5	35
47	Comparative Microbiomics of Tephritid Frugivorous Pests (Diptera: Tephritidae) From the Field: A Tale of High Variability Across and Within Species. Frontiers in Microbiology, 2020, 11, 1890.	1.5	24
48	Genome sequence-based curation of PubMLST data challenges interspecies recombination in the Burkholderia cepacia complex. Future Microbiology, 2020, 15, 1091-1093.	1.0	2
49	Decoding the Capability of Lactobacillus plantarum W1 Isolated from Soybean Whey in Producing an Exopolysaccharide. ACS Omega, 2020, 5, 33387-33394.	1.6	7
50	Microbial enrichment, functional characterization and isolation from a cold seep yield piezotolerant obligate hydrocarbon degraders. FEMS Microbiology Ecology, 2020, 96, .	1.3	5
51	Roadmap for naming uncultivated Archaea and Bacteria. Nature Microbiology, 2020, 5, 987-994.	5.9	115
52	MALDI-TOF MS profiling of non-starter lactic acid bacteria from artisanal cheeses of the Greek island of Naxos. International Journal of Food Microbiology, 2020, 323, 108586.	2.1	33
53	A critical rebuttal of the proposed division of the genus Arcobacter into six genera using comparative genomic, phylogenetic, and phenotypic criteria. Systematic and Applied Microbiology, 2020, 43, 126108.	1.2	33
54	Achromobacter veterisilvae sp. nov., from a mixed hydrogen-oxidizing bacteria enrichment reactor for microbial protein production. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 530-536.	0.8	21

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55	Pectobacterium parvum sp. nov., having a Salmonella SPI-1-like Type III secretion system and low virulence. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 2440-2448.	0.8	51
56	A taxonomic note on the genus Lactobacillus: Description of 23 novel genera, emended description of the genus Lactobacillus Beijerinck 1901, and union of Lactobacillaceae and Leuconostocaceae. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 2782-2858.	0.8	2,775
57	An emended description of Arcobacter anaerophilus Sasi Jyothsna et al. 2013: genomic and phenotypic insights. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 3921-3923.	0.8	5
58	Micromonospora fluminis sp. nov., isolated from mountain river sediment. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 6428-6436.	0.8	8
59	Novel acetic acid bacteria from cider fermentations: Acetobacter conturbans sp. nov. and Acetobacter fallax sp. nov. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 6163-6171.	0.8	25
60	Nocardiopsis dassonvillei subsp. crassaminis subsp. nov., isolated from freshwater sediment, and reappraisal of Nocardiopsis alborubida Grund and Kroppenstedt 1990 emend. Nouioui et al. 2018. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 6172-6179.	0.8	7
61	Introducing SPeDE: High-Throughput Dereplication and Accurate Determination of Microbial Diversity from Matrix-Assisted Laser Desorptionâ€“Ionization Time of Flight Mass Spectrometry Data. MSystems, 2019, 4, .	1.7	53
62	Comparative Genomics of Pandoraea, a Genus Enriched in Xenobiotic Biodegradation and Metabolism. Frontiers in Microbiology, 2019, 10, 2556.	1.5	85
63	Enriched hydrogen-oxidizing microbiomes show a high diversity of co-existing hydrogen-oxidizing bacteria. Applied Microbiology and Biotechnology, 2019, 103, 8241-8253.	1.7	24
64	A Method for Comprehensive Proteomic Analysis of Human Faecal Samples to Investigate Gut Dysbiosis in Patients with Cystic Fibrosis. Advances in Experimental Medicine and Biology, 2019, 1073, 137-160.	0.8	5
65	Gene Expansion and Positive Selection as Bacterial Adaptations to Oligotrophic Conditions. MSphere, 2019, 4, .	1.3	28
66	Impact of Sample Preservation and Manipulation on Insect Gut Microbiome Profiling. A Test Case With Fruit Flies (Diptera, Tephritidae). Frontiers in Microbiology, 2019, 10, 2833.	1.5	38
67	The Buffer Capacity and Calcium Concentration of Water Influence the Microbial Species Diversity, Grain Growth, and Metabolite Production During Water Kefir Fermentation. Frontiers in Microbiology, 2019, 10, 2876.	1.5	29
68	Influence of microbiota in the susceptibility of parasitic wasps to abamectin insecticide: deep sequencing, esterase and toxicity tests. Pest Management Science, 2019, 75, 79-86.	1.7	16
69	Lactococci dominate the bacterial communities of fermented maize, sorghum and millet slurries in Zimbabwe. International Journal of Food Microbiology, 2019, 289, 77-87.	2.1	15
70	Actinomadura roseirufa sp. nov., producer of semduramicin, a polyether ionophore. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 3068-3073.	0.8	8
71	Characterization of novel Gluconobacter species from fruits and fermented food products: Gluconobacter cadivus sp. nov., Gluconobacter vitians sp. nov. and Gluconobacter potus sp. nov. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	0.8	20
72	Pedobacter gandavensis sp. nov., Pedobacter foliorum sp. nov. and Pedobacter planticolens sp. nov., isolated from leaves of Arabidopsis thaliana. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	0.8	16

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73	Entomobacter blattae gen. nov., sp. nov., a new member of the Acetobacteraceae isolated from the gut of the cockroach Gromphadorhina portentosa. International Journal of Systematic and Evolutionary Microbiology, 2019, 71, .	0.8	19
74	Oxygen and diverse nutrients influence the water kefir fermentation process. Food Microbiology, 2018, 73, 351-361.	2.1	59
75	Baobab fruit pulp and mopane worm as potential functional ingredients to improve the iron and zinc content and bioaccessibility of fermented cereals. Innovative Food Science and Emerging Technologies, 2018, 47, 390-398.	2.7	30
76	Temporal and Spatial Distribution of the Acetic Acid Bacterium Communities throughout the Wooden Casks Used for the Fermentation and Maturation of Lambic Beer Underlines Their Functional Role. Applied and Environmental Microbiology, 2018, 84, .	1.4	32
77	Flow cytometric fingerprinting for microbial strain discrimination and physiological characterization. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2018, 93, 201-212.	1.1	43
78	High amorphous poly-beta-hydroxybutyrate (PHB) content in a probiotic Bacillus strain displays better protective effects in Vibrio -challenged gnotobiotic Artemia. Aquaculture, 2018, 487, 15-21.	1.7	13
79	A different gut microbial community between larvae and adults of a wild bumblebee nest (Bombus) Tj ETQq1 1 0.784314 rgBT /Over	1.5	23
80	Traditional fermentation and cooking of finger millet: Implications on mineral binders and subsequent bioaccessibility. Journal of Food Composition and Analysis, 2018, 68, 87-94.	1.9	15
81	Largeâ€scale cultivation of the bumblebee gut microbiota reveals an underestimated bacterial species diversity capable of pathogen inhibition. Environmental Microbiology, 2018, 20, 214-227.	1.8	40
82	Iron and zinc bioaccessibility of fermented maize, sorghum and millets from five locations in Zimbabwe. Food Research International, 2018, 103, 361-370.	2.9	27
83	Wort Substrate Consumption and Metabolite Production During Lambic Beer Fermentation and Maturation Explain the Successive Growth of Specific Bacterial and Yeast Species. Frontiers in Microbiology, 2018, 9, 2763.	1.5	35
84	Genome sequences and description of novel exopolysaccharides producing species Komagataeibacter pomaceti sp. nov. and reclassification of Komagataeibacter kombuchae (Dutta and Gachhui 2007) Yamada et al., 2013 as a later heterotypic synonym of Komagataeibacter hansenii (GosselÃ© et al. 1983) Yamada et al., 2013. Systematic and Applied Microbiology, 2018, 41, 581-592.	1.2	46
85	Enzymatic degradation of mineral binders in cereals: Impact on iron and zinc bioaccessibility. Journal of Cereal Science, 2018, 82, 223-229.	1.8	7
86	A prokaryoticâ€“eukaryotic relation in the fat body of Bombus terrestris. Environmental Microbiology Reports, 2018, 10, 644-650.	1.0	6
87	Filling the gaps in clinical proteomics: a do-it-yourself guide for the identification of the emerging pathogen Arcobacter by matrix-assisted laser desorption ionization-time of flight mass spectrometry. Journal of Microbiological Methods, 2018, 152, 92-97.	0.7	10
88	In Vitro Susceptibility of Burkholderia cepacia Complex Isolated from Cystic Fibrosis Patients to Ceftazidime-Avibactam and Ceftolozane-Tazobactam. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	34
89	Halomonas nigrificans sp. nov., isolated from cheese. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 371-376.	0.8	17
90	Burkholderia insecticola sp. nov., a gut symbiotic bacterium of the bean bug Riptortus pedestris. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2370-2374.	0.8	27

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91	Symbiotic and non-symbiotic Paraburkholderia isolated from South African Lebeckia ambigua root nodules and the description of Paraburkholderia fynbosensis sp. nov.. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2607-2614.	0.8	28
92	Antimicrobial activity against Mycobacterium tuberculosis under in vitro lipid-rich dormancy conditions. Journal of Medical Microbiology, 2018, 67, 282-285.	0.7	12
93	Can fermentation be used as a sustainable strategy to reduce iron and zinc binders in traditional African fermented cereal porridges or gruels?. Food Reviews International, 2017, 33, 561-586.	4.3	24
94	Effect of oral administration of lactic acid bacteria on colony performance and gut microbiota in indoor-reared bumblebees (Bombus terrestris). Apidologie, 2017, 48, 41-50.	0.9	22
95	Identification of acetic acid bacteria through matrix-assisted laser desorption/ionization time-of-flight mass spectrometry and report of Gluconobacter nephelii Kommanee et al. 2011 and Gluconobacter uchimurae Tanasupawat et al. 2012 as later heterotypic synonyms of Gluconobacter japonicus Malimas et al. 2009 and Gluconobacter oxydans (Henneberg 1897) De Ley 1961 (Approved Lists) Tj ETQq1 1 0.784314 rgB	1.2	21
96	Gilliamella intestini sp. nov., Gilliamella bombicola sp. nov., Gilliamella bombi sp. nov. and Gilliamella mensalis sp. nov.: Four novel Gilliamella species isolated from the bumblebee gut. Systematic and Applied Microbiology, 2017, 40, 199-204.	1.2	44
97	Paenibacillus aquistagni sp. nov., isolated from an artificial lake accumulating industrial wastewater. Antonie Van Leeuwenhoek, 2017, 110, 1189-1197.	0.7	13
98	Draft genome and description of Orrella dioscoreae gen. nov. sp. nov., a new species of Alcaligenaceae isolated from leaf acumens of Dioscorea sansibarensis. Systematic and Applied Microbiology, 2017, 40, 11-21.	1.2	42
99	Clinical and microbiological profile of chronic Burkholderia cepacia complex infections in a cystic fibrosis reference hospital in Brazil. European Journal of Clinical Microbiology and Infectious Diseases, 2017, 36, 2263-2271.	1.3	5
100	Detection, isolation and characterization of Fusobacterium gastrosuis sp. nov. colonizing the stomach of pigs. Systematic and Applied Microbiology, 2017, 40, 42-50.	1.2	40
101	â€œGenetic regulation of Mycobacterium tuberculosis in a lipid-rich environmentâ€. Infection, Genetics and Evolution, 2017, 55, 392-402.	1.0	18
102	The transcriptome of Mycobacterium tuberculosis in a lipid-rich dormancy model through RNAseq analysis. Scientific Reports, 2017, 7, 17665.	1.6	88
103	Reconciliation between operational taxonomic units and species boundaries. FEMS Microbiology Ecology, 2017, 93, .	1.3	71
104	Characterization of Mycobacterium chelonae-Like Strains by Comparative Genomics. Frontiers in Microbiology, 2017, 8, 789.	1.5	6
105	Comparative Genomics of Burkholderia singularis sp. nov., a Low G+C Content, Free-Living Bacterium That Defies Taxonomic Dissection of the Genus Burkholderia. Frontiers in Microbiology, 2017, 8, 1679.	1.5	36
106	In vitro activity of bedaquiline against rapidly growing nontuberculous mycobacteria. Journal of Medical Microbiology, 2017, 66, 1140-1143.	0.7	38
107	Aeromicrobium choanae sp. nov., an actinobacterium isolated from the choana of a garden warbler. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 357-361.	0.8	13
108	Paraburkholderia piptadeniae sp. nov. and Paraburkholderia ribeironis sp. nov., two root-nodulating symbiotic species of Piptadenia gonoacantha in Brazil. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 432-440.	0.8	50

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109	Pedobacter jamesrossensis sp. nov., Pedobacter lithocola sp. nov., Pedobacter mendelii sp. nov. and Pedobacter petrophilus sp. nov., isolated from the Antarctic environment. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 1499-1507.	0.8	32
110	Arcobacter haliotis sp. nov., isolated from abalone species Haliotis gigantea. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3050-3056.	0.8	26
111	Oryzisolibacter propanilivorax gen. nov., sp. nov., a propanil-degrading bacterium. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3752-3758.	0.8	12
112	Minimal standards for describing new species belonging to the families Campylobacteraceae and Helicobacteraceae: Campylobacter, Arcobacter, Helicobacter and Wolinella spp.. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 5296-5311.	0.8	84
113	Comparative genomics of Burkholderia multivorans, a ubiquitous pathogen with a highly conserved genomic structure. PLoS ONE, 2017, 12, e0176191.	1.1	17
114	Microbial Ecology of Traditional Beer Fermentations. , 2017, , .		9
115	Members of the genus Burkholderia: good and bad guys. F1000Research, 2016, 5, 1007.	0.8	280
116	Biochemical properties of three lactic acid bacteria strains isolated from traditional cassava starters used for attieke preparation. African Journal of Food Science, 2016, 10, 271-277.	0.4	3
117	Phylogenomic Study of Burkholderia glathei-like Organisms, Proposal of 13 Novel Burkholderia Species and Emended Descriptions of Burkholderia sordidicola, Burkholderia zhejiangensis, and Burkholderia grimmiae. Frontiers in Microbiology, 2016, 7, 877.	1.5	120
118	PCR detection of Burkholderia multivorans in water and soil samples. BMC Microbiology, 2016, 16, 184.	1.3	6
119	Phylogenomic Analysis Reveals an Asian Origin for African Burkholderia pseudomallei and Further Supports Melioidosis Endemicity in Africa. MSphere, 2016, 1, .	1.3	57
120	Unipept web services for metaproteomics analysis. Bioinformatics, 2016, 32, 1746-1748.	1.8	44
121	Extensive cultivation of soil and water samples yields various pathogens in patients with cystic fibrosis but not Burkholderia multivorans. Journal of Cystic Fibrosis, 2016, 15, 769-775.	0.3	30
122	Burkholderia: an update on taxonomy and biotechnological potential as antibiotic producers. Applied Microbiology and Biotechnology, 2016, 100, 5215-5229.	1.7	222
123	Combined approach to the identification of clinically infrequent non-tuberculous mycobacteria in Argentina. International Journal of Tuberculosis and Lung Disease, 2016, 20, 1257-1262.	0.6	5
124	New Methyloceanibacter diversity from North Sea sediments includes methanotroph containing solely the soluble methane monooxygenase. Environmental Microbiology, 2016, 18, 4523-4536.	1.8	81
125	Effect of Fermentation and Cooking on Soluble and Bound Phenolic Profiles of Finger Millet Sour Porridge. Journal of Agricultural and Food Chemistry, 2016, 64, 7615-7621.	2.4	23
126	The unique peptidome: Taxon-specific tryptic peptides as biomarkers for targeted metaproteomics. Proteomics, 2016, 16, 2313-2318.	1.3	28

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127	Optimizing of a protein extraction method for Mycobacterium tuberculosis proteome analysis using mass spectrometry. Journal of Microbiological Methods, 2016, 131, 144-147.	0.7	20
128	Timely Identification of the Emerging Zoonotic Enteric Pathogen Arcobacter: Thank Heaven for Matrix-Assisted Laser Desorption/Ionization?. Open Forum Infectious Diseases, 2016, 3, .	0.4	0
129	Selected Lactobacillus strains isolated from sugary and milk kefir reduce Salmonella infection of epithelial cells in vitro. Beneficial Microbes, 2016, 7, 585-595.	1.0	29
130	Proposal of Helicobacter canicola sp. nov., previously identified as Helicobacter cinaedi, isolated from canines. Systematic and Applied Microbiology, 2016, 39, 307-312.	1.2	20
131	Acetic Acid Bacteria in Fermented Food and Beverage Ecosystems. , 2016, , 73-99.		8
132	Helicobacter saguini, a Novel Helicobacter Isolated from Cotton-Top Tamarins with Ulcerative Colitis, Has Proinflammatory Properties and Induces Typhlocolitis and Dysplasia in Gnotobiotic IL-10 ^{â^'/â^'} Mice. Infection and Immunity, 2016, 84, 2307-2316.	1.0	25
133	MALDI-TOF MS as a Novel Tool for Dereplication and Characterization of Microbiota in Bacterial Diversity Studies. , 2016, , 235-256.		9
134	Microbial diversity and metabolite composition of Belgian red-brown acidic ales. International Journal of Food Microbiology, 2016, 221, 1-11.	2.1	38
135	Divergence between the Highly Virulent Zoonotic Pathogen Helicobacter heilmannii and Its Closest Relative, the Low-Virulence â€œHelicobacter ailurogastricusâ€•sp. nov. Infection and Immunity, 2016, 84, 293-306.	1.0	37
136	Faecal proteomics: A tool to investigate dysbiosis and inflammation in patients with cystic fibrosis. Journal of Cystic Fibrosis, 2016, 15, 242-250.	0.3	72
137	Reclassification of Achromobacter spiritinus Vandamme et al. 2013 as a later heterotypic synonym of Achromobacter marplatensis Gomila et al. 2011. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 1641-1644.	0.8	14
138	Bifidobacterium aquikefiri sp. nov., isolated from water kefir. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 1281-1286.	0.8	53
139	Apibacter mensalis sp. nov.: a rare member of the bumblebee gut microbiota. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 1645-1651.	0.8	29
140	Corynebacterium lowii sp. nov. and Corynebacterium oculi sp. nov., derived from human clinical disease and an emended description of Corynebacterium mastitidis. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 2803-2812.	0.8	40
141	Notes on the use of Greek word roots in genus and species names of prokaryotes. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 2129-2140.	0.8	11
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