## Linda L. Blackall

List of Publications by Year in descending order

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20759 20900 14,761 179 60 115 citations h-index g-index papers 186 186 186 10548 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Advances in enhanced biological phosphorus removal: From micro to macro scale. Water Research, 2007, 41, 2271-2300.	5.3	998
2	Identification of Polyphosphate-Accumulating Organisms and Design of 16S rRNA-Directed Probes for Their Detection and Quantitation. Applied and Environmental Microbiology, 2000, 66, 1175-1182.	1.4	691
3	Metagenomic analysis of two enhanced biological phosphorus removal (EBPR) sludge communities. Nature Biotechnology, 2006, 24, 1263-1269.	9.4	634
4	rRNA Sequences and Evolutionary Relationships among Toxic and Nontoxic Cyanobacteria of the Genus Microcystis. International Journal of Systematic Bacteriology, 1997, 47, 693-697.	2.8	439
5	Microbial ecology meets electrochemistry: electricity-driven and driving communities. ISME Journal, 2007, 1, 9-18.	4.4	433
6	Bacterial community structures of phosphate-removing and non-phosphate-removing activated sludges from sequencing batch reactors. Applied and Environmental Microbiology, 1995, 61, 1910-1916.	1.4	429
7	Glycogen-accumulating organisms in laboratory-scale and full-scale wastewater treatment processes b bThe GenBank accession numbers for the sequences reported in this paper are given in Methods Microbiology (United Kingdom), 2002, 148, 3353-3364.	0.7	377
8	Shifting paradigms in restoration of the world's coral reefs. Global Change Biology, 2017, 23, 3437-3448.	4.2	351
9	Phylogenetic Diversity of Bacteria Associated with the Marine Sponge Rhopaloeides odorabile. Applied and Environmental Microbiology, 2001, 67, 434-444.	1.4	322
10	Investigation of Candidate Division TM7, a Recently Recognized Major Lineage of the Domain Bacteria with No Known Pure-Culture Representatives. Applied and Environmental Microbiology, 2001, 67, 411-419.	1.4	311
11	Multiple Lateral Transfers of Dissimilatory Sulfite Reductase Genes between Major Lineages of Sulfate-Reducing Prokaryotes. Journal of Bacteriology, 2001, 183, 6028-6035.	1.0	309
12	Metamorphosis of a Scleractinian Coral in Response to Microbial Biofilms. Applied and Environmental Microbiology, 2004, 70, 1213-1221.	1.4	287
13	Cathodic oxygen reduction catalyzed by bacteria in microbial fuel cells. ISME Journal, 2008, 2, 519-527.	4.4	268
14	Coral microbiome dynamics, functions and design in a changing world. Nature Reviews Microbiology, 2019, 17, 557-567.	13.6	267
15	Filamentous Chloroflexi (green non-sulfur bacteria) are abundant in wastewater treatment processes with biological nutrient removal c cThe EMBL accession numbers for the sequences reported in this paper are X84472 (strain SBR1029 16S rDNA), X84474 (strain SBR1031 16S rDNA), X84498 (strain SBR1064) T	j <b>ΕΦQ</b> q1 1	0. <b>28</b> \$314 rgl
16	Comparison of acetate and propionate uptake by polyphosphate accumulating organisms and glycogen accumulating organisms. Biotechnology and Bioengineering, 2005, 91, 162-168.	1.7	233
17	Use of Stable-Isotope Probing, Full-Cycle rRNA Analysis, and Fluorescence In Situ Hybridization-Microautoradiography To Study a Methanol-Fed Denitrifying Microbial Community. Applied and Environmental Microbiology, 2004, 70, 588-596.	1.4	213
18	Identification of Some of the Major Groups of Bacteria in Efficient and Nonefficient Biological Phosphorus Removal Activated Sludge Systems. Applied and Environmental Microbiology, 1999, 65, 4077-4084.	1.4	202

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19	Hydrolysis and microbial community analyses in two-stage anaerobic digestion of energy crops. Journal of Applied Microbiology, 2007, 103, 516-527.	1.4	186
20	Coralâ€"the world's most diverse symbiotic ecosystem. Molecular Ecology, 2015, 24, 5330-5347.	2.0	184
21	Induction of Larval Metamorphosis of the Coral Acropora millepora by Tetrabromopyrrole Isolated from a Pseudoalteromonas Bacterium. PLoS ONE, 2011, 6, e19082.	1.1	184
22	Identification and comparison of aerobic and denitrifying polyphosphate-accumulating organisms. Biotechnology and Bioengineering, 2003, 83, 140-148.	1.7	162
23	A review and update of the microbiology of enhanced biological phosphorus removal in wastewater treatment plants. Antonie Van Leeuwenhoek, 2002, 81, 681-691.	0.7	161
24	Investigation of an Acetate-Fed Denitrifying Microbial Community by Stable Isotope Probing, Full-Cycle rRNA Analysis, and Fluorescent In Situ Hybridization-Microautoradiography. Applied and Environmental Microbiology, 2005, 71, 8683-8691.	1.4	160
25	Development and Use of Fluorescent In Situ Hybridization Probes for the Detection and Identification of "Microthrix parvicella―in Activated Sludge. Systematic and Applied Microbiology, 1997, 20, 310-318.	1.2	158
26	Enhanced biological phosphorus removal in a sequencing batch reactor using propionate as the sole carbon source. Biotechnology and Bioengineering, 2004, 85, 56-67.	1.7	158
27	Putative glycogen-accumulating organisms belonging to the Alphaproteobacteria identified through rRNA-based stable isotope probing. Microbiology (United Kingdom), 2006, 152, 419-429.	0.7	156
28	The influence of substrate kinetics on the microbial community structure in granular anaerobic biomass. Water Research, 2004, 38, 1390-1404.	<b>5.</b> 3	155
29	Microbiology of a Nitrite-Oxidizing Bioreactor. Applied and Environmental Microbiology, 1998, 64, 1878-1883.	1.4	154
30	Minimizing errors in RT-PCR detection and quantification of SARS-CoV-2 RNA for wastewater surveillance. Science of the Total Environment, 2022, 805, 149877.	3.9	153
31	Reclassification of Pasteurella gallinarum, [Haemophilus] paragallinarum, Pasteurella avium and Pasteurella volantium as Avibacterium gallinarum gen. nov., comb. nov., Avibacterium paragallinarum comb. nov., Avibacterium avium comb. nov. and Avibacterium volantium comb. nov International lournal of Systematic and Evolutionary Microbiology, 2005, 55, 353-362.	0.8	141
32	A bacterial metapopulation adapts locally to phage predation despite global dispersal. Genome Research, 2008, 18, 293-297.	2.4	135
33	Streptococcus caprinus sp.nov., a tannin-resistant ruminal bacterium from feral goats. Letters in Applied Microbiology, 1994, 18, 313-318.	1.0	130
34	The use of second derivative plots for the determination of mol% guanine plus cytosine of DNA by the thermal denaturation method. Journal of Microbiological Methods, 1986, 5, 139-156.	0.7	129
35	Dynamics of a temperature-related coral disease outbreak. Marine Ecology - Progress Series, 2004, 281, 63-77.	0.9	120
36	Design and Evaluation of 16S rRNA-Targeted Oligonucleotide Probes for Fluorescence In Situ Hybridization., 2002, 179, 029-042.		116

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37	Identification, Detection, and Spatial Resolution of Clostridium Populations Responsible for Cellulose Degradation in a Methanogenic Landfill Leachate Bioreactor. Applied and Environmental Microbiology, 2004, 70, 2414-2419.	1.4	113
38	Changes in equine hindgut bacterial populations during oligofructose-induced laminitis. Environmental Microbiology, 2006, 8, 885-898.	1.8	113
39	Phylogenetic relationships of filamentous sulfur bacteria (Thiothrix spp. and Eikelboom type 021N) Tj ETQq1 1 C Thiothrix unzii sp. nov., Thiothrix fructosivorans sp. nov. and Thiothrix defluvii sp. nov International Journal of Systematic and Evolutionary Microbiology, 1999, 49, 1817-1827.	0.784314 i 0.8	gBT /Overloc 112
40	Phylogenetic positions of phytoplasmas associated with dieback, yellow crinkle and mosaic diseases of papaya, and their proposed inclusion in 'Candidatus Phytoplasma australiense' and a new taxon, 'Candidatus Phytoplasma australasia'. International Journal of Systematic Bacteriology, 1998, 48, 941-951.	2.8	111
41	Challenges for simultaneous nitrification, denitrification, and phosphorus removal in microbial aggregates: mass transfer limitation and nitrous oxide production. FEMS Microbiology Ecology, 2005, 52, 329-338.	1.3	108
42	Anaerobic and aerobic metabolism of glycogen-accumulating organisms selected with propionate as the sole carbon source. Microbiology (United Kingdom), 2006, 152, 2767-2778.	0.7	108
43	Abundance and ecophysiology of Defluviicoccus spp., glycogen-accumulating organisms in full-scale wastewater treatment processes. Microbiology (United Kingdom), 2007, 153, 178-185.	0.7	106
44	The contribution of microbial biotechnology to mitigating coral reef degradation. Microbial Biotechnology, 2017, 10, 1236-1243.	2.0	101
45	Phylogeny of the filamentous bacterium Eikelboom Type 1851, and design and application of a 16S rRNA targeted oligonucleotide probe for its fluorescence in situ identification in activated sludge. FEMS Microbiology Letters, 2002, 207, 179-183.	0.7	100
46	Microplastic pollution alters forest soil microbiome. Journal of Hazardous Materials, 2021, 409, 124606.	6.5	100
47	Sludge population optimisation: a new dimension for the control of biological wastewater treatment systems. Water Research, 2002, 36, 482-490.	5.3	98
48	Functionally Relevant Microorganisms to Enhanced Biological Phosphorus Removal Performance at Fullâ€Scale Wastewater Treatment Plants in the United States. Water Environment Research, 2008, 80, 688-698.	1.3	95
49	What do we really know about sponge-microbial symbioses?. ISME Journal, 2009, 3, 1-3.	4.4	92
50	A Proposal To Reclassify Nocardia pinensis Blackall et al. as Skermania piniformis gen. nov., comb. nov International Journal of Systematic Bacteriology, 1997, 47, 127-131.	2.8	87
51	Microbial distribution of <i>Accumulibacter</i> spp. and <i>Competibacter</i> spp. in aerobic granules from a labâ€scale biological nutrient removal system. Environmental Microbiology, 2008, 10, 354-363.	1.8	86
52	Molecular investigation of a microbial mat associated with the Great Artesian Basin. FEMS Microbiology Ecology, 1998, 25, 391-403.	1.3	84
53	Fluorescence In Situ Hybridization and Spectral Imaging of Coral-Associated Bacterial Communities. Applied and Environmental Microbiology, 2006, 72, 3016-3020.	1.4	83
54	Microbial ecology of the equine hindgut during oligofructose-induced laminitis. ISME Journal, 2008, 2, 1089-1100.	4.4	82

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55	Disease and cell death in white syndrome of Acroporid corals on the Great Barrier Reef. Marine Biology, 2007, 151, 19-29.	0.7	81
56	Development of a multi-locus sequence typing scheme for avian isolates of Pasteurella multocida. Veterinary Microbiology, 2010, 141, 354-361.	0.8	81
57	'Candidatus Nostocoida limicola', a filamentous bacterium from activated sludge International Journal of Systematic and Evolutionary Microbiology, 2000, 50, 703-709.	0.8	77
58	"Microthrix parvicella―is a Novel, Deep Branching Member of the Actinomycetes Subphylum. Systematic and Applied Microbiology, 1995, 17, 513-518.	1.2	76
59	Structure of a cellulose degrading bacterial community during anaerobic digestion. Biotechnology and Bioengineering, 2005, 92, 871-878.	1.7	<b>7</b> 5
60	"Candidatus Microthrix parvicella," a Filamentous Bacterium from Activated Sludge Sewage Treatment Plants. International Journal of Systematic Bacteriology, 1996, 46, 344-346.	2.8	73
61	Concurrent microscopic observations and activity measurements of cellulose hydrolyzing and methanogenic populations during the batch anaerobic digestion of crystalline cellulose. Biotechnology and Bioengineering, 2005, 91, 369-378.	1.7	70
62	Beneath the surface: community assembly and functions of the coral skeleton microbiome. Microbiome, 2019, 7, 159.	4.9	67
63	16S rRNA Analysis of Isolates Obtained from Gram-Negative, Filamentous Bacteria Micromanipulated from Activated Sludge. Systematic and Applied Microbiology, 1996, 19, 334-343.	1.2	61
64	Analysis of the microbial community structure and function of a laboratory scale enhanced biological phosphorus removal reactor. Environmental Microbiology, 2002, 4, 559-569.	1.8	61
65	Intracellular bacteria are common and taxonomically diverse in cultured and <i>in hospite</i> endosymbionts of coral reefs. ISME Journal, 2021, 15, 2028-2042.	4.4	61
66	Metabolic transformations and characterisation of the sludge community in an enhanced biological phosphorus removal system. Applied Microbiology and Biotechnology, 1998, 49, 226-234.	1.7	56
67	Proton motive force generation from stored polymers for the uptake of acetate under anaerobic conditions. FEMS Microbiology Letters, 2007, 274, 245-251.	0.7	56
68	Experimental Inoculation of Coral Recruits With Marine Bacteria Indicates Scope for Microbiome Manipulation in Acropora tenuis and Platygyra daedalea. Frontiers in Microbiology, 2019, 10, 1702.	1.5	55
69	Cellulolytic and dextranolytic Gramâ€negative bacteria: revival of the genus <i>Cellvibrio</i> . Journal of Applied Bacteriology, 1985, 59, 81-97.	1.1	53
70	Monitoring associations between clade-level variation, overall community structure and ecosystem function in enhanced biological phosphorus removal (EBPR) systems using terminal-restriction fragment length polymorphism (T-RFLP). Water Research, 2010, 44, 4908-4923.	5.3	51
71	Molecular Identification of Activated Sludge Foaming Bacteria. Water Science and Technology, 1994, 29, 35-42.	1.2	49
72	A multiple-outgroup approach to resolving division-level phylogenetic relationships using 16S rDNA data International Journal of Systematic and Evolutionary Microbiology, 2001, 51, 385-391.	0.8	46

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73	Cinnamaldehyde disrupts biofilm formation and swarming motility of Pseudomonas aeruginosa. Microbiology (United Kingdom), 2018, 164, 1087-1097.	0.7	46
74	Bioenergetic models for acetate and phosphate transport in bacteria important in enhanced biological phosphorus removal. Environmental Microbiology, 2008, 10, 87-98.	1.8	45
75	Anaerobic phosphate release from activated sludge with enhanced biological phosphorus removal. A possible mechanism of intracellular pH control. , 1999, 63, 507-515.		44
76	Anaerobic glyoxylate cycle activity during simultaneous utilization of glycogen and acetate in uncultured <i>Accumulibacter</i> enriched in enhanced biological phosphorus removal communities. ISME Journal, 2008, 2, 1040-1051.	4.4	44
77	Foaming in activated sludge plants: A survey in Queensland, Australia and an evaluation of some control strategies. Water Research, 1991, 25, 313-317.	5.3	42
78	PCR detection of Clostridium perfringens producing different toxins in faeces of goats. Letters in Applied Microbiology, 1997, 25, 339-344.	1.0	42
79	Kinetic and phylogenetic characterization of an anaerobic dechlorinating microbial community. Microbiology (United Kingdom), 2003, 149, 459-469.	0.7	40
80	Phylogeny of the filamentous bacterium 'Nostocoida limicola' III from activated sludge International Journal of Systematic and Evolutionary Microbiology, 2001, 51, 195-202.	0.8	40
81	The use of 16S rDNA clone libraries to describe the microbial diversity of activated sludge communities. Water Science and Technology, 1998, 37, 451-454.	1.2	39
82	Microscale structure and function of anaerobic $\hat{A}$ $\hat{A}$ $\hat{A}$ $\hat{A}$ aerobic granules containing glycogen accumulating organisms. FEMS Microbiology Ecology, 2003, 45, 253-261.	1.3	39
83	Limitations of the widely used GAM42a and BET42a probes targeting bacteria in the Gammaproteobacteria radiation. Microbiology (United Kingdom), 2003, 149, 1239-1247.	0.7	39
84	Meganema perideroedes gen. nov., sp. nov., a filamentous alphaproteobacterium from activated sludge. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 1865-1868.	0.8	39
85	The filamentous morphotype Eikelboom Type 1863 is not a single genetic entity. Journal of Applied Microbiology, 1997, 82, 411-421.	1.4	38
86	Activity of Cinnamaldehyde on Quorum Sensing and Biofilm Susceptibility to Antibiotics in Pseudomonas aeruginosa. Microorganisms, 2020, 8, 455.	1.6	38
87	The mechanism of stabilization of actinomycete foams and the prevention of foaming under laboratory conditions. Journal of Industrial Microbiology, 1989, 4, 181-187.	0.9	36
88	Actinomycete Scum Problems in Australian Activated Sludge Plants. Water Science and Technology, 1988, 20, 493-495.	1.2	36
89	Activated sludge foams: Effects of environmental variables on organism growth and foam formation. Environmental Technology (United Kingdom), 1991, 12, 241-248.	1.2	35
90	The Filamentous Bacterial Morphotype â€~Nostocoida limicola' I Contains at least Two Previously Described Genera in the Low G+C Gram Positive Bacteria. Systematic and Applied Microbiology, 2000, 23, 528-534.	1.2	33

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91	Exaiptasia diaphana from the great barrier reef: a valuable resource for coral symbiosis research. Symbiosis, 2020, 80, 195-206.	1.2	33
92	The characterization and description of representatives of  G' bacteria from activated sludge plants. Letters in Applied Microbiology, 1997, 25, 63-69.	1.0	32
93	Using Bacterial Extract along with Differential Gene Expression in Acropora millepora Larvae to Decouple the Processes of Attachment and Metamorphosis. PLoS ONE, 2012, 7, e37774.	1.1	32
94	Phylogenetic and physiological characterization of a heterotrophic, chemolithoautotrophic Thiothrix strain isolated from activated sludge. International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 1271-1276.	0.8	31
95	Mixedâ€mode bacterial transmission in the common brooding coral <i>Pocillopora acuta</i> Environmental Microbiology, 2020, 22, 397-412.	1.8	31
96	Phenotypic and phylogenetic description of an Italian isolate of " Microthrix parvicella ― Journal of Applied Microbiology, 1997, 82, 405-410.	1.4	30
97	Variability of type 021N in activated sludge as determined by substrate uptake pattern and hybridization with fluorescent rRNA targeted probes. Water Science and Technology, 1998, 37, 423.	1.2	30
98	Early Life Stages of a Common Broadcast Spawning Coral Associate with Specific Bacterial Communities Despite Lack of Internalized Bacteria. Microbial Ecology, 2020, 79, 706-719.	1.4	30
99	Development of a free radical scavenging bacterial consortium to mitigate oxidative stress in cnidarians. Microbial Biotechnology, 2021, 14, 2025-2040.	2.0	30
100	Biological nutrient removal efficiency in treatment of saline wastewater. Water Science and Technology, 1999, 39, 183.	1.2	28
101	Assessment of bacterial community composition within and among Acropora loripes colonies in the wild and in captivity. Coral Reefs, 2020, 39, 1245-1255.	0.9	28
102	The use of 16S rDNA clone libraries to describe the microbial diversity of activated sludge communities. Water Science and Technology, 1998, 37, 451.	1.2	27
103	Nitrifying bacterial communities in an aquaculture wastewater treatment system using fluorescence in situ hybridization (FISH), 16S rRNA gene cloning, and phylogenetic analysis. Biotechnology and Bioengineering, 2007, 97, 985-990.	1.7	27
104	Some physiological properties of an Italian isolate of ". Water Science and Technology, 1998, 37, 1.	1.2	26
105	Comparison of cellulose solubilisation rates in rumen and landfill leachate inoculated reactors. Bioresource Technology, 2006, 97, 2356-2363.	4.8	26
106	A feral goat rumen fluid inoculum improves nitrogen retention in sheep consuming a mulga (Acacia) Tj ETQq0 0	0 rgBT /O	verlock 10 Tf 5
107	Characterisation of a novel <i>Mannheimia</i> sp from Australian feedlot cattle. Australian Veterinary Journal, 2001, 79, 634-639.	0.5	24
108	PCR detection of Clostridium chauvoei in pure cultures and in formalin-fixed, paraffin-embedded tissues. Veterinary Microbiology, 2003, 91, 239-248.	0.8	23

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109	The effect of biomass density on cellulose solubilisation rates. Bioresource Technology, 2008, 99, 4723-4731.	4.8	23
110	Microbiota characterization of Exaiptasia diaphana from the Great Barrier Reef. Animal Microbiome, 2020, 2, 10.	1.5	23
111	Host Traits and Phylogeny Contribute to Shaping Coral-Bacterial Symbioses. MSystems, 2022, 7, e0004422.	1.7	22
112	Multiple techniques point to oxygenic phototrophs dominating the Isopora palifera skeletal microbiome. Coral Reefs, 2021, 40, 275-282.	0.9	21
113	Some physiological properties of an Italian isolate of "microthrix parvicella― Water Science and Technology, 1998, 37, 1-8.	1.2	20
114	Aerobic nitrate respiration in a nitrite-oxidising bioreactor. FEMS Microbiology Letters, 2000, 184, 113-118.	0.7	20
115	DNA extraction from coral reef sediment bacteria for the polymerase chain reaction. Journal of Microbiological Methods, 2000, 43, 73-80.	0.7	20
116	Phylogenetic analysis of <i>Porphyromonas</i> species isolated from the oral cavity of Australian marsupials. Environmental Microbiology, 2008, 10, 2425-2432.	1.8	20
117	Anaerobic central metabolic pathways active during polyhydroxyalkanoate production in uncultured cluster $1 < i > Defluviicoccus < / i > enriched in activated sludge communities. FEMS Microbiology Letters, 2009, 298, 79-84.$	0.7	20
118	Intracellular Bacterial Symbionts in Corals: Challenges and Future Directions. Microorganisms, 2021, 9, 2209.	1.6	20
119	The opportunistic pathogen Nocardia farcinica is a foam-producing bacterium in activated sludge plants. Letters in Applied Microbiology, 1996, 22, 342-346.	1.0	19
120	A review of anti-nutritive factors limiting potential use of Acacia angustissima as a ruminant feed. Animal Feed Science and Technology, 2008, 147, 158-171.	1.1	19
121	Application of flowcell technology for monitoring biofilm development and cellulose degradation in leachate and rumen systems. Bioresource Technology, 2009, 100, 492-496.	4.8	19
122	Characterisation of enhanced biological phosphorus removal activated sludges with dissimilar phosphorus removal performances. Water Science and Technology, 1998, 37, 567-571.	1.2	19
123	Isolation and identification of an Eikelboom type 1863 strain as Acinetobacter johnsonii. Water Research, 1997, 31, 657-660.	5.3	18
124	Isolation and characterization of a Clostridium sp. with cinnamoyl esterase activity and unusual cell envelope ultrastructure. Archives of Microbiology, 1999, 172, 139-149.	1.0	18
125	Oral Disease in Animals: The Australian Perspective. Isolation and Characterisation of Black-Pigmented Bacteria from the Oral Cavity of Marsupials. Anaerobe, 2002, 8, 79-87.	1.0	18
126	The Effect of Thermal Stress on the Bacterial Microbiome of Exaiptasia diaphana. Microorganisms, 2020, 8, 20.	1.6	18

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127	An in vitro cultured rumen inoculum improves nitrogen digestion in mulga-fed sheep. Australian Journal of Agricultural Research, 1997, 48, 403.	1.5	18
128	Phylogenetic Analysis and Taxonomic History of Nocardia pinensis and Nocardia amarae. Systematic and Applied Microbiology, 1995, 17, 519-525.	1.2	16
129	Bio-P and non-bio-P bacteria identification by a novel microbial approach. Water Science and Technology, 1999, 39, 13.	1.2	16
130	Discrepancies in the widely applied GAM42a fluorescence in situ hybridisation probe for Gamma proteobacteria. FEMS Microbiology Letters, 2005, 242, 367-373.	0.7	16
131	Applicability of experience from laboratory reactors with biological phosphorus removal in full-scale plants. Water Science and Technology, 2006, 54, 267-275.	1.2	16
132	Microbes orchestrate life on Earth. ISME Journal, 2008, 2, 795-796.	4.4	16
133	Streptococcus caprinus is ineffective as a rumen inoculum to improve digestion of mulga (Acacia) Tj ETQq $1\ 1\ 0.7$	784314 rg 1.5	$BT_{15}^{/O}$ verlock
134	Characterisation of enhanced biological phosphorus removal activated sludges with dissimilar phosphorus removal performances. Water Science and Technology, 1998, 37, 567.	1.2	15
135	Integrating process engineering and microbiology tools to advance activated sludge wastewater treatment research and development. Reviews in Environmental Science and Biotechnology, 2002, $1$ , 83-97.	3.9	15
136	Bacterial and algal symbiont dynamics in early recruits exposed to two adult coral species. Coral Reefs, 2020, 39, 189-202.	0.9	15
137	Detection by polymerase chain reaction of Clostridium perfringens producing epsilon toxin in faeces and in gastrointestinal contents of goats. Letters in Applied Microbiology, 1996, 23, 13-17.	1.0	14
138	A survey of the relative abundance of specific groups of cellulose degrading bacteria in anaerobic environments using fluorescencein situhybridization. Journal of Applied Microbiology, 2007, 103, 1332-1343.	1.4	14
139	Induction of membrane permeability in <i>Escherichia coli</i> $\hat{A}$ \$\hat{\text{A}}\$\hat{\text{A}}\$\hat{\text{f}}\$ mediated by lysis protein of the ColE7 operon. FEMS Microbiology Letters, 2009, 298, 85-92.	0.7	14
140	Microbiome characterization of defensive tissues in the model anemone Exaiptasia diaphana. BMC Microbiology, 2021, 21, 152.	1.3	14
141	Towards understanding the taxonomy of some of the filamentous bacteria causing bulking and foaming in activated sludge plants. Water Science and Technology, 1996, 34, 137-144.	1.2	14
142	Exploring microbiome engineering as a strategy for improved thermal tolerance in <i>Exaiptasia diaphana</i> . Journal of Applied Microbiology, 2022, 132, 2940-2956.	1.4	14
143	Lack of evidence for the oxidative stress theory of bleaching in the sea anemone, Exaiptasia diaphana, under elevated temperature. Coral Reefs, 2022, 41, 1161-1172.	0.9	14
144	New foam-forming nocardioforms found in activated sludge. Water Science and Technology, 1998, 37, 495.	1.2	12

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145	Further limitations of phylogenetic group-specific probes used for detection of bacteria in environmental samples. ISME Journal, 2010, 4, 959-961.	4.4	12
146	New foam-forming nocardioforms found in activated sludge. Water Science and Technology, 1998, 37, 495-502.	1.2	12
147	PCR detection ofClostridium perfringenstype D in formalinâ€fixed, paraffinâ€embedded tissues of goats and sheep. Letters in Applied Microbiology, 1999, 29, 15-19.	1.0	11
148	Enrichment, isolation and characterisation of ruminal bacteria that degrade non-protein amino acids from the tropical legume Acacia angustissima. Animal Feed Science and Technology, 2005, 121, 191-204.	1.1	11
149	Short-Term Exposure to Sterile Seawater Reduces Bacterial Community Diversity in the Sea Anemone, Exaiptasia diaphana. Frontiers in Marine Science, 2021, 7, .	1.2	11
150	Effect of Multispecies Microbial Consortia on Microbially Influenced Corrosion of Carbon Steel. Corrosion and Materials Degradation, 2021, 2, 133-149.	1.0	11
151	Porphyromonas loveana sp. nov., isolated from the oral cavity of Australian marsupials. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 3771-3778.	0.8	11
152	Towards understanding the taxonomy of some of the filamentous bacteria causing bulking and foaming in activated sludge plants. Water Science and Technology, 1996, 34, 137.	1.2	10
153	Characterisation of the bacterial consortium involved in nitrite oxidation in activated sludge. Water Science and Technology, 1999, 39, 45.	1.2	10
154	Isolation and characterization of an unusual bacterium, allied to the soil bacteriumBacillus benzoevorans, from feedlot manure pads in Australia. Letters in Applied Microbiology, 1999, 29, 71-75.	1.0	9
155	Eco-physiological characterization of fluorescence in situ hybridization probe-targeted denitrifiers in activated sludge using culture-independent methods. Letters in Applied Microbiology, 2007, 44, 399-405.	1.0	9
156	Is marine sediment the source of microbes associated with accelerated low water corrosion?. Applied Microbiology and Biotechnology, 2019, 103, 449-459.	1.7	9
157	Probiotics for corals. Microbiology Australia, 2020, 41, 100.	0.1	9
158	Characterisation of the bacterial consortium involved in nitrite oxidation in activated sludge. Water Science and Technology, 1999, 39, 45-52.	1.2	9
159	Bio-P and non-bio-P bacteria identification by a novel microbial approach. Water Science and Technology, 1999, 39, 13-20.	1.2	8
160	Microbial Communities of Orange Tubercles in Accelerated Low-Water Corrosion. Applied and Environmental Microbiology, 2020, 86, .	1.4	5
161	A comparison of biological nutrient removal in intermittent cyclic and continuous activated sludge systems. Resources, Conservation and Recycling, 1994, 11, 149-159.	5.3	4
162	Molecular biological methods to detect "―and to determine its abundance in activated sludge. Water Science and Technology, 1998, 37, 37.	1.2	4

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163	Bacterial growth kinetics estimation by fluorescence in situ hybridization and spectrofluorometric quantification. Letters in Applied Microbiology, 2007, 44, 643-648.	1.0	4
164	Antibiotics reduce bacterial load in Exaiptasia diaphana, but biofilms hinder its development as a gnotobiotic coral model. Access Microbiology, 2022, 4, 000314.	0.2	4
165	Identification of microbes isolated with test kits through culture-dependent and metabarcoding techniques for assessment of microbial corrosion. Current Research in Biotechnology, 2022, 4, 129-137.	1.9	4
166	Construction and analysis of a metagenomic library from an enhanced biological phosphorus removal biomass. Water Science and Technology, 2006, 54, 277-284.	1.2	2
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