

# Brendan P Burns

## List of Publications by Year in descending order

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80  
papers

4,425  
citations

94433

37  
h-index

110387

64  
g-index

84  
all docs

84  
docs citations

84  
times ranked

4863  
citing authors

#	ARTICLE	IF	CITATIONS
1	Advances in on-line drinking water quality monitoring and early warning systems. <i>Water Research</i> , 2011, 45, 741-747.	11.3	286
2	Diversity within cyanobacterial mat communities in variable salinity meltwater ponds of McMurdo Ice Shelf, Antarctica. <i>Environmental Microbiology</i> , 2005, 7, 519-529.	3.8	252
3	Identification and Characterization of <i>Helicobacter pylori</i> Genes Essential for Gastric Colonization. <i>Journal of Experimental Medicine</i> , 2003, 197, 813-822.	8.5	246
4	Microbial diversity of extant stromatolites in the hypersaline marine environment of Shark Bay, Australia. <i>Environmental Microbiology</i> , 2004, 6, 1096-1101.	3.8	225
5	Bacterial, archaeal and eukaryotic diversity of smooth and pustular microbial mat communities in the hypersaline lagoon of Shark Bay. <i>Geobiology</i> , 2009, 7, 82-96.	2.4	164
6	Microgravity Induces Pelvic Bone Loss through Osteoclastic Activity, Osteocytic Osteolysis, and Osteoblastic Cell Cycle Inhibition by CDKN1a/p21. <i>PLoS ONE</i> , 2013, 8, e61372.	2.5	148
7	Unravelling core microbial metabolisms in the hypersaline microbial mats of Shark Bay using high-throughput metagenomics. <i>ISME Journal</i> , 2016, 10, 183-196.	9.8	147
8	Phylogeography of the invasive cyanobacterium <i>Cylindrospermopsis raciborskii</i> . <i>Molecular Ecology</i> , 2002, 12, 133-140.	3.9	138
9	Detection and sequencing of the microcystin LR-degrading gene, <i>mlrA</i> , from new bacteria isolated from Japanese lakes. <i>FEMS Microbiology Letters</i> , 2003, 229, 271-276.	1.8	137
10	Niche differentiation of bacterial communities at a millimeter scale in Shark Bay microbial mats. <i>Scientific Reports</i> , 2015, 5, 15607.	3.3	137
11	Carotenoid Analysis of Halophilic Archaea by Resonance Raman Spectroscopy. <i>Astrobiology</i> , 2007, 7, 631-643.	3.0	132
12	Determining the specific microbial populations and their spatial distribution within the stromatolite ecosystem of Shark Bay. <i>ISME Journal</i> , 2009, 3, 383-396.	9.8	125
13	Bioastronautics: The Influence of Microgravity on Astronaut Health. <i>Astrobiology</i> , 2010, 10, 463-473.	3.0	115
14	Microgravity Reduces the Differentiation and Regenerative Potential of Embryonic Stem Cells. <i>Stem Cells and Development</i> , 2015, 24, 2605-2621.	2.1	94
15	Disentangling the drivers of functional complexity at the metagenomic level in Shark Bay microbial mat microbiomes. <i>ISME Journal</i> , 2018, 12, 2619-2639.	9.8	94
16	Dynamics of archaea at fine spatial scales in Shark Bay mat microbiomes. <i>Scientific Reports</i> , 2017, 7, 46160.	3.3	87
17	Quorum Sensing in Extreme Environments. <i>Life</i> , 2013, 3, 131-148.	2.4	80
18	<i>Halococcus hamelinensis</i> sp. nov., a novel halophilic archaeon isolated from stromatolites in Shark Bay, Australia. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2006, 56, 1323-1329.	1.7	73

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19	Mechanical unloading of bone in microgravity reduces mesenchymal and hematopoietic stem cell-mediated tissue regeneration. <i>Stem Cell Research</i> , 2014, 13, 181-201.	0.7	68
20	Microbial Diversity of Browning Peninsula, Eastern Antarctica Revealed Using Molecular and Cultivation Methods. <i>Frontiers in Microbiology</i> , 2017, 8, 591.	3.5	66
21	Asgard archaea: Diversity, function, and evolutionary implications in a range of microbiomes. <i>AIMS Microbiology</i> , 2019, 5, 48-61.	2.2	65
22	Interactions between intracellular Na <sup>+</sup> levels and saxitoxin production in <i>Cylindrospermopsis raciborskii</i> T3. <i>Microbiology (United Kingdom)</i> , 2004, 150, 455-461.	1.8	61
23	Molecular Identification of Cyanobacteria Associated with Stromatolites from Distinct Geographical Locations. <i>Astrobiology</i> , 2002, 2, 271-280.	3.0	59
24	Lipid biomarkers in Hamelin Pool microbial mats and stromatolites. <i>Organic Geochemistry</i> , 2010, 41, 1207-1218.	1.8	57
25	Optimized Rapid Amplification of cDNA Ends (RACE) for Mapping Bacterial mRNA Transcripts. <i>BioTechniques</i> , 2000, 28, 448-456.	1.8	56
26	Lipid biomarker analysis of cyanobacteria-dominated microbial mats in meltwater ponds on the McMurdo Ice Shelf, Antarctica. <i>Organic Geochemistry</i> , 2009, 40, 258-269.	1.8	52
27	The Entner-Doudoroff Pathway in <i>Helicobacter pylori</i> . <i>Archives of Biochemistry and Biophysics</i> , 1994, 312, 349-356.	3.0	51
28	Novel homologs of the multiple resistance regulator <i>marA</i> in antibiotic-contaminated environments. <i>Water Research</i> , 2008, 42, 4271-4280.	11.3	50
29	Untapped Resources: Biotechnological Potential of Peptides and Secondary Metabolites in Archaea. <i>Archaea</i> , 2015, 2015, 1-7.	2.3	50
30	Host specificity and phylogeography of the prochlorophyte <i>Prochloron</i> sp., an obligate symbiont in didemnid ascidians. <i>Environmental Microbiology</i> , 2007, 9, 890-899.	3.8	49
31	<i>Haloferax elongans</i> sp. nov. and <i>Haloferax mucosum</i> sp. nov., isolated from microbial mats from Hamelin Pool, Shark Bay, Australia. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 798-802.	1.7	46
32	Osmoadaptive Strategies of the Archaeon <i>Halococcus hamelinensis</i> Isolated from a Hypersaline Stromatolite Environment. <i>Astrobiology</i> , 2011, 11, 529-536.	3.0	46
33	Lysis efficiency of standard DNA extraction methods for <i>Halococcus</i> spp. in an organic rich environment. <i>Extremophiles</i> , 2008, 12, 301-308.	2.3	43
34	Molecular Ecology of Hypersaline Microbial Mats: Current Insights and New Directions. <i>Microorganisms</i> , 2016, 4, 6.	3.6	43
35	Global Protein-Level Responses of <i>Halobacterium salinarum</i> NRC-1 to Prolonged Changes in External Sodium Chloride Concentrations. <i>Journal of Proteome Research</i> , 2009, 8, 2218-2225.	3.7	42
36	Discovery of an Abundance of Biosynthetic Gene Clusters in Shark Bay Microbial Mats. <i>Frontiers in Microbiology</i> , 2020, 11, 1950.	3.5	39

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37	Genetic potential for secondary metabolite production in stromatolite communities. <i>FEMS Microbiology Letters</i> , 2005, 243, 293-301.	1.8	38
38	Analysis of intergenic spacer region length polymorphisms to investigate the halophilic archaeal diversity of stromatolites and microbial mats. <i>Extremophiles</i> , 2007, 11, 203-210.	2.3	38
39	Modern analogues and the early history of microbial life. <i>Precambrian Research</i> , 2009, 173, 10-18.	2.7	38
40	A review of analytical methods for assessing the public health risk from microcystin in the aquatic environment. <i>Journal of Water Supply: Research and Technology - AQUA</i> , 2005, 54, 509-518.	1.4	36
41	Microbial dark matter filling the niche in hypersaline microbial mats. <i>Microbiome</i> , 2020, 8, 135.	11.1	35
42	Extremophilic adaptations and biotechnological applications in diverse environments. <i>AIMS Microbiology</i> , 2016, 2, 251-261.	2.2	34
43	Identification of an Na <sup>+</sup> -Dependent Transporter Associated with Saxitoxin-Producing Strains of the Cyanobacterium <i>Anabaena circinalis</i> . <i>Applied and Environmental Microbiology</i> , 2004, 70, 4711-4719.	3.1	33
44	Investigations into the taxonomy, toxicity and ecology of benthic cyanobacterial accumulations in Myall Lake, Australia. <i>Marine and Freshwater Research</i> , 2005, 56, 45.	1.3	33
45	Characterisation of glucose transport in <i>Helicobacter pylori</i> . <i>Biochimica Et Biophysica Acta - General Subjects</i> , 1995, 1244, 269-276.	2.4	32
46	Viral Communities of Shark Bay Modern Stromatolites. <i>Frontiers in Microbiology</i> , 2018, 9, 1223.	3.5	32
47	New Approaches to Detect Biosynthetic Gene Clusters in the Environment. <i>Medicines (Basel)</i> , 2021, 10, 1430.	1.4	30
48	Identification and regulation of novel compatible solutes from hypersaline stromatolite-associated cyanobacteria. <i>Archives of Microbiology</i> , 2010, 192, 1031-1038.	2.2	29
49	Bioinformatic, phylogenetic and chemical analysis of the UV-absorbing compounds scytonemin and mycosporine-like amino acids from the microbial mat communities of Shark Bay, Australia. <i>Environmental Microbiology</i> , 2019, 21, 702-715.	3.8	27
50	Between a Rock and a Soft Place: The Role of Viruses in Lithification of Modern Microbial Mats. <i>Trends in Microbiology</i> , 2021, 29, 204-213.	7.7	26
51	Modern arsenotrophic microbial mats provide an analogue for life in the anoxic Archean. <i>Communications Earth &amp; Environment</i> , 2020, 1, .	6.8	24
52	Genome Sequence of the Halophilic Archaeon <i>Halococcus hamelinensis</i> . <i>Journal of Bacteriology</i> , 2012, 194, 2100-2101.	2.2	23
53	Adaptation, Ecology, and Evolution of the Halophilic Stromatolite Archaeon <i>Halococcus hamelinensis</i> Inferred through Genome Analyses. <i>Archaea</i> , 2015, 2015, 1-11.	2.3	23
54	Detection and characterization of N-acyl-L-homoserine lactones using GFP-based biosensors in conjunction with thin-layer chromatography. <i>Journal of Microbiological Methods</i> , 2015, 118, 164-167.	1.6	20

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55	Functional Gene Expression in Shark Bay Hypersaline Microbial Mats: Adaptive Responses. <i>Frontiers in Microbiology</i> , 2020, 11, 560336.	3.5	20
56	Molecular assessment of UVC radiation-induced DNA damage repair in the stromatolitic halophilic archaeon, <i>Halococcus hamelinensis</i> . <i>Journal of Photochemistry and Photobiology B: Biology</i> , 2011, 102, 140-145.	3.8	18
57	Correlation of bio-optical properties with photosynthetic pigment and microorganism distribution in microbial mats from Hamelin Pool, Australia. <i>FEMS Microbiology Ecology</i> , 2019, 95, .	2.7	18
58	The Vulnerability of Microbial Ecosystems in A Changing Climate: Potential Impact in Shark Bay. <i>Life</i> , 2019, 9, 71.	2.4	16
59	Comparative gene expression of PSP-toxin producing and non-toxic <i>Anabaena circinalis</i> strains. <i>Environment International</i> , 2006, 32, 743-748.	10.0	15
60	On the Response of Halophilic Archaea to Space Conditions. <i>Life</i> , 2014, 4, 66-76.	2.4	15
61	The <i>Helicobacter pylori</i> pyrB Gene Encoding Aspartate Carbamoyltransferase Is Essential for Bacterial Survival. <i>Archives of Biochemistry and Biophysics</i> , 2000, 380, 78-84.	3.0	14
62	In Situ Properties of <i>Helicobacter pylori</i> Aspartate Carbamoyltransferase. <i>Archives of Biochemistry and Biophysics</i> , 1997, 347, 119-125.	3.0	13
63	Molecular Detection of Genes Responsible for Cyanobacterial Toxin Production in the Genera <i>Microcystis</i> , <i>Nodularia</i> , and <i>Cylindrospermopsis</i> . , 2004, 268, 213-222.		13
64	A Novel Mechanism for Resistance to the Antimetabolite N -Phosphonoacetyl- Aspartate by <i>Helicobacter pylori</i> . <i>Journal of Bacteriology</i> , 1998, 180, 5574-5579.	2.2	13
65	Isolation of novel quorum-sensing active bacteria from microbial mats in Shark Bay Australia. <i>FEMS Microbiology Ecology</i> , 2019, 95, .	2.7	12
66	Methods for the measurement of a bacterial enzyme activity in cell lysates and extracts. <i>Biological Procedures Online</i> , 1998, 1, 17-26.	2.9	11
67	Quorum Sensing in Archaea: Recent Advances and Emerging Directions. , 2017, , 119-132.		11
68	Communication within East Antarctic Soil Bacteria. <i>Applied and Environmental Microbiology</i> , 2019, 86, .	3.1	11
69	Archaea join the conversation: detection of AHL-like activity across a range of archaeal isolates. <i>FEMS Microbiology Letters</i> , 2020, 367, .	1.8	11
70	Absence of detectable levels of the cyanobacterial toxin (microcystin-LR) carry-over into milk. <i>Toxicon</i> , 2002, 40, 1173-1180.	1.6	10
71	Genome-resolved metagenomics provides insights into the functional complexity of microbial mats in Blue Holes, Shark Bay. <i>FEMS Microbiology Ecology</i> , 2022, 98, .	2.7	10
72	Use of ion-channel modulating agents to study cyanobacterial Na <sup>+</sup> -K <sup>+</sup> fluxes. <i>Biological Procedures Online</i> , 2004, 6, 137-143.	2.9	9

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73	Eukarya the chimera: eukaryotes, a secondary innovation of the two domains of life?. Trends in Microbiology, 2021, , .	7.7	6
74	Characterization of Arginine Transport in Helicobacter pylori. Helicobacter, 2003, 8, 245-251.	3.5	3
75	A Cyanobacteria Enriched Layer of Shark Bay Stromatolites Reveals a New Acaryochloris Strain Living in Near Infrared Light. Microorganisms, 2022, 10, 1035.	3.6	1
76	Stromatolites as a Resource for Novel Natural Products. Origins of Life and Evolution of Biospheres, 2007, 36, 623-624.	1.9	0
77	Session 18. Functional Complexity of Modern Stromatolites and Microbial Mats. Astrobiology, 2008, 8, 378-383.	3.0	0
78	Adsorption and Biodegradation Characteristics of Musty Odorous Compounds, 2-Methylisoborneol and Geosmin. Japanese Journal of Water Treatment Biology, 2006, 42, 85-91.	0.1	0
79	Molecular Approaches to Studying Living Stromatolites. Lecture Notes in Earth Sciences, 2011, , 91-100.	0.5	0
80	Metabolite Transport. , 0, , 207-217.		0