## Catherine Anne Biggs

List of Publications by Year in descending order

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44 papers

1,924 citations

304602 22 h-index 42 g-index

45 all docs

45 docs citations

45 times ranked

3277 citing authors

#	Article	IF	CITATIONS
1	An insight into iTRAQ: where do we stand now?. Analytical and Bioanalytical Chemistry, 2012, 404, 1011-1027.	1.9	293
2	Methodological approaches for studying the microbial ecology of drinking water distribution systems. Water Research, 2014, 65, 134-156.	5.3	215
3	Characterization of the Extracellular Polymeric Substances Produced by <i>Escherichia coli</i> Using Infrared Spectroscopic, Proteomic, and Aggregation Studies. Biomacromolecules, 2008, 9, 686-695.	2.6	188
4	Proteomics with a pinch of salt: A cyanobacterial perspective. Saline Systems, 2008, 4, 1.	2.0	114
5	Mechanisms of Bacillus cereus biofilm formation: an investigation of the physicochemical characteristics of cell surfaces and extracellular proteins. Applied Microbiology and Biotechnology, 2011, 89, 1161-1175.	1.7	98
6	"Biofilmology― a multidisciplinary review of the study of microbial biofilms. Applied Microbiology and Biotechnology, 2011, 90, 1869-1881.	1.7	96
7	Magnetic-Silk Core–Shell Nanoparticles as Potential Carriers for Targeted Delivery of Curcumin into Human Breast Cancer Cells. ACS Biomaterials Science and Engineering, 2017, 3, 1027-1038.	2.6	75
8	A new coupon design for simultaneous analysis of in situ microbial biofilm formation and community structure in drinking water distribution systems. Applied Microbiology and Biotechnology, 2010, 87, 749-756.	1.7	67
9	The polymer physics and chemistry of microbial cell attachment and adhesion. Faraday Discussions, 2008, 139, 85.	1.6	59
10	Physicochemical analysis of initial adhesion and biofilm formation of Methanosarcina barkeri on polymer support material. Colloids and Surfaces B: Biointerfaces, 2016, 143, 518-525.	2.5	58
11	Comparative Proteomics Study of Salt Tolerance between a Nonsequenced Extremely Halotolerant Cyanobacterium and Its Mildly Halotolerant Relative Using <i>in vivo</i> Metabolic Labeling and <i>in vitro</i> Isobaric Labeling. Journal of Proteome Research, 2009, 8, 818-828.	1.8	51
12	Methods in Quantitative Proteomics: Setting iTRAQ on the Right Track. Current Proteomics, 2011, 8, 17-30.	0.1	42
13	Macromolecular Fingerprinting of <i>Sulfolobus</i> Species in Biofilm: A Transcriptomic and Proteomic Approach Combined with Spectroscopic Analysis. Journal of Proteome Research, 2011, 10, 4105-4119.	1.8	41
14	Investigating the Surface Properties of Escherichia coli under Glucose Controlled Conditions and Its Effect on Aggregation. Langmuir, 2007, 23, 6691-6697.	1.6	40
15	A Quantitative Proteomic Analysis of Light Adaptation in a Globally Significant Marine Cyanobacterium Prochlorococcus marinus MED4. Journal of Proteome Research, 2007, 6, 996-1005.	1.8	37
16	Influence of fermentation conditions on the surface properties and adhesion of Lactobacillus rhamnosus GG. Microbial Cell Factories, 2012, 11, 116.	1.9	36
17	A crossâ€species quantitative proteomic study of salt adaptation in a halotolerant environmental isolate using <sup>15</sup> N metabolic labelling. Proteomics, 2008, 8, 2266-2284.	1.3	35
18	Biodesalination: A Case Study for Applications of Photosynthetic Bacteria in Water Treatment Â. Plant Physiology, 2014, 164, 1661-1676.	2.3	33

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19	Simbiotics: A Multiscale Integrative Platform for 3D Modeling of Bacterial Populations. ACS Synthetic Biology, 2017, 6, 1194-1210.	1.9	33
20	Bacterial growth dynamics in activated sludge batch assays. Water Research, 1998, 32, 587-596.	5.3	32
21	Harvesting and dewatering yeast by microflotation. Biochemical Engineering Journal, 2014, 82, 174-182.	1.8	26
22	Using a multi-faceted approach to determine the changes in bacterial cell surface properties influenced by a biofilm lifestyle. Biofouling, 2012, 28, 1-14.	0.8	24
23	Effect of temperature on the substrate utilization profiles of microbial communities in different sewer sediments. Environmental Technology (United Kingdom), 2011, 32, 133-144.	1.2	22
24	The quantitative proteomic response of Synechocystis sp. PCC6803 to phosphate acclimation. Aquatic Biosystems, 2013, 9, 5.	1.8	22
25	Deciphering the unique cellulose degradation mechanism of the ruminal bacterium Fibrobacter succinogenes S85. Scientific Reports, 2019, 9, 16542.	1.6	22
26	Investigating the effect of patulin, penicillic acid and EDTA on biofilm formation of isolates from dental unit water lines. Applied Microbiology and Biotechnology, 2008, 81, 349-358.	1.7	20
27	Quantitative protein expression and cell surface characteristics of <i>Escherichia coli</i> MG1655 biofilms. Proteomics, 2011, 11, 339-351.	1.3	20
28	A systems biology approach to investigate the response of Synechocystis sp. PCC6803 to a high salt environment. Saline Systems, 2009, 5, 8.	2.0	19
29	The importance of sewer biofilms. Wiley Interdisciplinary Reviews: Water, 2016, 3, 487-494.	2.8	18
30	Influence of Substrates on the Surface Characteristics and Membrane Proteome of Fibrobacter succinogenes S85. PLoS ONE, 2015, 10, e0141197.	1.1	13
31	Comparative quantitative proteomics of prochlorococcus ecotypes to a decrease in environmental phosphate concentrations. Aquatic Biosystems, 2012, 8, 7.	1.8	12
32	Cellular acclimation strategies of a minimal picocyanobacterium to phosphate stress. FEMS Microbiology Letters, 2010, 306, 127-134.	0.7	11
33	Enumeration of sulphate-reducing bacteria for assessing potential for hydrogen sulphide production in urban drainage systems. Water Science and Technology, 2016, 73, 3087-3094.	1.2	11
34	Application of enhanced assimilable organic carbon method across operational drinking water systems. PLoS ONE, 2019, 14, e0225477.	1.1	11
35	Understanding the costs of investigating coliform and <i>E.Âcoli</i> detections during routine drinking water quality monitoring. Urban Water Journal, 2018, 15, 101-108.	1.0	6
36	Hydraulic conditioning to manage potable water discolouration. Water Management, 2019, 172, 3-13.	0.4	6

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37	Proteomic analysis of the impact of static culturing on the expansion of rat bone marrow mesenchymal stem cells. Biotechnology Letters, 2012, 34, 1589-1596.	1.1	4
38	Bacteriological water quality compliance and root cause analysis: an industry case study. Water Science and Technology: Water Supply, 2013, 13, 1034-1045.	1.0	4
39	Spatial and temporal variability of bacterial communities within a combined sewer system. MicrobiologyOpen, 2016, 5, 616-625.	1.2	4
40	Comparative study of in vitro expansion of bone marrow-derived mesenchymal stem cells. Biotechnology Letters, 2013, 35, 463-469.	1,1	2
41	Data mining T-RFLP profiles from urban water system sampling using self-organizing maps. , 2012, , .		1
42	Detecting, monitoring and controlling biofilm formation. Membrane Technology, 2014, 2014, 9-10.	0.5	1
43	Looking through the FOG: microbiome characterization and lipolytic bacteria isolation from a fatberg site. Microbiology (United Kingdom), 2021, 167, .	0.7	1
44	'Biodesalination': a synthetic biology approach for the use of photosynthetic bacteria in water treatment. New Biotechnology, 2014, 31, S140-S141.	2.4	0