## Paul Baker

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Electrochemical quantification of d-glucose during the production of bioethanol from thermo-mechanically pre-treated wheat straw. Electrochemistry Communications, 2021, 124, 106942.	2.3	8
2	HPLC-DAD-qTOF Compositional Analysis of the Phenolic Compounds Present in Crude Tomato Protein Extracts Derived from Food Processing. Molecules, 2021, 26, 6403.	1.7	6
3	Developing an Olive Biorefinery in Slovenia: Analysis of Phenolic Compounds Found in Olive Mill Pomace and Wastewater. Molecules, 2021, 26, 7.	1.7	15
4	Antimicrobial Potential of Plastic Films Incorporated with Sage Extract on Chicken Meat. Foods, 2021, 10, 2812.	1.9	15
5	A comparison in protein extraction from four major crop residues in Europe using chemical and enzymatic processes-a review. Innovative Food Science and Emerging Technologies, 2020, 59, 102239.	2.7	26
6	Pressurised disc refining of wheat straw as a pre-treatment approach for agricultural residues: A preliminary assessment of energy consumption and fibre composition. Bioresource Technology, 2020, 304, 122976.	4.8	12
7	Fibre degradation of wheat straw by <i>Pleurotus erygnii</i> under low moisture conditions during solid-state fermentation. Letters in Applied Microbiology, 2019, 68, 182-187.	1.0	8
8	Fungal pre-treatment of forestry biomass with a focus on biorefining: A comparison of biomass degradation and enzyme activities by wood rot fungi across three tree species. Biomass and Bioenergy, 2017, 107, 20-28.	2.9	10
9	The patterns of bacterial community and relationships between sulfate-reducing bacteria and hydrochemistry in sulfate-polluted groundwater of Baogang rare earth tailings. Environmental Science and Pollution Research, 2016, 23, 21766-21779.	2.7	10
10	Biodegradation of Different Genotypes of Miscanthus by Wood Rot Fungi. BioResources, 2016, 11, .	0.5	7
11	Increased delignification by white rot fungi after pressure refining Miscanthus. Bioresource Technology, 2015, 189, 81-86.	4.8	26
12	Immobilization of Shewanella oneidensis MR-1 in diffusive gradients in thin films for determining metal bioavailability. Chemosphere, 2015, 138, 309-315.	4.2	2
13	In situ extraction of RNA from marine-derived fungi associated with the marine sponge, Haliclona simulans. Mycological Progress, 2012, 11, 953-956.	0.5	0
14	Endoglucanase activities and growth of marine-derived fungi isolated from the sponge <i>Haliclona simulans</i> . Journal of Applied Microbiology, 2010, 108, 1668-1675.	1.4	17
15	Isolation and Analysis of Bacteria with Antimicrobial Activities from the Marine Sponge Haliclona simulans Collected from Irish Waters. Marine Biotechnology, 2009, 11, 384-396.	1.1	168
16	Phylogenetic Diversity and Antimicrobial Activities of Fungi Associated with Haliclona simulans Isolated from Irish Coastal Waters. Marine Biotechnology, 2009, 11, 540-547.	1.1	72
17	Mir space station bacteria responses to modeled reduced gravity under starvation conditions. Advances in Space Research, 2006, 38, 1152-1158.	1.2	5
18	Intraspecific differences in bacterial responses to modelled reduced gravity. Journal of Applied Microbiology, 2005, 98, 1239-1246.	1.4	12

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19	Attachment to stainless steel by Mir Space Station bacteria growing under modeled reduced gravity at varying nutrient concentrations. Biofilms, 2005, 2, 1-7.	0.6	33
20	Seasonal patterns of abundance of viruses and bacteria in a Northeast Ohio (USA) stream. Archiv Für Hydrobiologie, 2004, 161, 225-233.	1.1	10
21	The effect of simulated microgravity on bacteria from the mir space station. Microgravity Science and Technology, 2004, 15, 35-41.	0.7	37
22	Escherichia coli growth under modeled reduced gravity. Microgravity Science and Technology, 2004, 15, 39-44.	0.7	39
23	Marine prosthecate bacteria involved in the ennoblement of stainless steel. Environmental Microbiology, 2003, 5, 925-932.	1.8	22
24	Bacterial populations occurring in a trichloroethylene-contaminated aquifer during methane injection. Environmental Microbiology, 2001, 3, 187-193.	1.8	10
25	Molecular diversity of pMMO and sMMO in a TCE-contaminated aquifer during bioremediation. FEMS Microbiology Ecology, 2001, 38, 161-167.	1.3	23
26	Environmentally relevant microorganisms. Journal of Bioscience and Bioengineering, 2000, 89, 1-11.	1.1	67
27	Analysis of actinomycete communities by specific amplification of genes encoding 16S rRNA and gel-electrophoretic separation in denaturing gradients. Applied and Environmental Microbiology, 1997, 63, 3233-3241.	1.4	1,298
28	Pre-processing Waste Tomatoes into Separated Streams with the Intention of Recovering Protein: Towards an Integrated Fruit and Vegetable Biorefinery Approach to Waste Minimization. Waste and Biomass Valorization, 0, , 1.	1.8	1