

# John W Moreau

## List of Publications by Year in descending order

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Version: 2024-02-01

47  
papers

2,448  
citations

257450

24  
h-index

223800

46  
g-index

55  
all docs

55  
docs citations

55  
times ranked

3777  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mercury sources, distribution, and bioavailability in the North Pacific Ocean: Insights from data and models. <i>Global Biogeochemical Cycles</i> , 2009, 23, .	4.9	378
2	Uranium mobility in organic matter-rich sediments: A review of geological and geochemical processes. <i>Earth-Science Reviews</i> , 2016, 159, 160-185.	9.1	283
3	Extracellular Proteins Limit the Dispersal of Biogenic Nanoparticles. <i>Science</i> , 2007, 316, 1600-1603.	12.6	254
4	Microbial mercury methylation in Antarctic sea ice. <i>Nature Microbiology</i> , 2016, 1, 16127.	13.3	158
5	Mineralogical Biosignatures and the Search for Life on Mars. <i>Astrobiology</i> , 2001, 1, 447-465.	3.0	139
6	Ultrastructure, aggregation-state, and crystal growth of biogenic nanocrystalline sphalerite and wurtzite. <i>American Mineralogist</i> , 2004, 89, 950-960.	1.9	102
7	Impact Features on Europa: Results of the Galileo Europa Mission (GEM). <i>Icarus</i> , 2001, 151, 93-111.	2.5	92
8	Ancient DNA from marine sediments: Precautions and considerations for seafloor coring, sample handling and data generation. <i>Earth-Science Reviews</i> , 2019, 196, 102887.	9.1	90
9	Mercury methylation by metabolically versatile and cosmopolitan marine bacteria. <i>ISME Journal</i> , 2021, 15, 1810-1825.	9.8	74
10	A Transmission Electron Microscopy Study of Silica and Kerogen Biosignatures in ~1.9 Ga Gunflint Microfossils. <i>Astrobiology</i> , 2004, 4, 196-210.	3.0	54
11	Model biomimetic studies of templated growth and assembly of nanocrystalline FeOOH. <i>Geochimica Et Cosmochimica Acta</i> , 2003, 67, 1185-1195.	3.9	50
12	Diversity of Dissimilatory Sulfite Reductase Genes ( <i>dsrAB</i> ) in a Salt Marsh Impacted by Long-Term Acid Mine Drainage. <i>Applied and Environmental Microbiology</i> , 2010, 76, 4819-4828.	3.1	48
13	Changes in the deep subsurface microbial biosphere resulting from a field-scale CO <sub>2</sub> geosequestration experiment. <i>Frontiers in Microbiology</i> , 2014, 5, 209.	3.5	44
14	Quantifying Heavy Metals Sequestration by Sulfate-Reducing Bacteria in an Acid Mine Drainage-Contaminated Natural Wetland. <i>Frontiers in Microbiology</i> , 2013, 4, 43.	3.5	43
15	Thio arsenic species measurements in marine organisms and geothermal waters. <i>Microchemical Journal</i> , 2013, 111, 82-90.	4.5	42
16	The Effect of Natural Organic Matter on Mercury Methylation by <i>Desulfobulbus propionicus</i> 1pr3. <i>Frontiers in Microbiology</i> , 2015, 6, 1389.	3.5	42
17	Characterization of uranium redox state in organic-rich Eocene sediments. <i>Chemosphere</i> , 2018, 194, 602-613.	8.2	40
18	<i>Thermoflavifilum aggregans</i> gen. nov., sp. nov., a thermophilic and slightly halophilic filamentous bacterium from the phylum Bacteroidetes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 1264-1270.	1.7	39

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19	Environmental considerations for subseabed geological storage of CO <sub>2</sub> : A review. <i>Continental Shelf Research</i> , 2014, 83, 116-128.	1.8	37
20	Metagenomic and lipid analyses reveal a diel cycle in a hypersaline microbial ecosystem. <i>ISME Journal</i> , 2015, 9, 2697-2711.	9.8	35
21	Microbial contributions to coupled arsenic and sulfur cycling in the acid-sulfide hot spring Champagne Pool, New Zealand. <i>Frontiers in Microbiology</i> , 2014, 5, 569.	3.5	32
22	<i>Thermorudis pharmacophila</i> sp. nov., a novel member of the class Thermomicrobia isolated from geothermal soil, and emended descriptions of <i>Thermomicrobium roseum</i> , <i>Thermomicrobium carboxidum</i> , <i>Thermorudis peleae</i> and <i>Sphaerobacter thermophilus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 4479-4487.	1.7	32
23	New insights into the genetic and metabolic diversity of thiocyanate-degrading microbial consortia. <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 1101-1108.	3.6	29
24	Crustal fluid and ash alteration impacts on the biosphere of Shikoku Basin sediments, Nankai Trough, Japan. <i>Geobiology</i> , 2015, 13, 562-580.	2.4	28
25	Distribution of iron- and sulfate-reducing bacteria across a coastal acid sulfate soil (CASS) environment: implications for passive bioremediation by tidal inundation. <i>Frontiers in Microbiology</i> , 2015, 6, 624.	3.5	27
26	Thiocyanate adsorption on ferrihydrite and its fate during ferrihydrite transformation to hematite and goethite. <i>Chemosphere</i> , 2015, 119, 987-993.	8.2	27
27	Characterization of an autotrophic bioreactor microbial consortium degrading thiocyanate. <i>Applied Microbiology and Biotechnology</i> , 2017, 101, 5889-5901.	3.6	23
28	Biodegradation of thiocyanate by a novel strain <i>Burkholderia phytofirmans</i> from soil contaminated by gold mine tailings. <i>Letters in Applied Microbiology</i> , 2013, 57, 368-372.	2.2	21
29	<i>Limisphaera ngatamarikiensis</i> gen. nov., sp. nov., a thermophilic, pink-pigmented coccus isolated from subaqueous mud of a geothermal hot spring. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 1114-1121.	1.7	20
30	<i>In Situ</i> Stimulation of Thiocyanate Biodegradation through Phosphate Amendment in Gold Mine Tailings Water. <i>Environmental Science &amp; Technology</i> , 2017, 51, 13353-13362.	10.0	20
31	Genome-Resolved Metagenomics and Detailed Geochemical Speciation Analyses Yield New Insights into Microbial Mercury Cycling in Geothermal Springs. <i>Applied and Environmental Microbiology</i> , 2020, 86, .	3.1	19
32	The geomicrobiology of CO <sub>2</sub> geosequestration: a focused review on prokaryotic community responses to field-scale CO <sub>2</sub> injection. <i>Frontiers in Microbiology</i> , 2015, 6, 263.	3.5	17
33	Experimental evaluation of sampling, storage and analytical protocols for measuring arsenic speciation in sulphidic hot spring waters. <i>Microchemical Journal</i> , 2017, 130, 162-167.	4.5	16
34	Genome-resolved metagenomics of an autotrophic thiocyanate-remediating microbial bioreactor consortium. <i>Water Research</i> , 2019, 158, 106-117.	11.3	11
35	Biodegradation of thiocyanate by a native groundwater microbial consortium. <i>PeerJ</i> , 2019, 7, e6498.	2.0	10
36	Manganese-reducing <i>Pseudomonas fluorescens</i> -group bacteria control arsenic mobility in gold mining-contaminated groundwater. <i>Environmental Earth Sciences</i> , 2014, 71, 4187-4198.	2.7	9

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37	Thiocyanate biodegradation: harnessing microbial metabolism for mine remediation. <i>Microbiology Australia</i> , 2018, 39, 157.	0.4	8
38	Mercury distribution and mobility at the abandoned Puhipuhi mercury mine, Northland, New Zealand. <i>New Zealand Journal of Geology, and Geophysics</i> , 2015, 58, 78-87.	1.8	7
39	The effect of heavy metals on thiocyanate biodegradation by an autotrophic microbial consortium enriched from mine tailings. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 417-427.	3.6	6
40	Time-resolved microbial guild responses to tidal cycling in a coastal acid-sulfate system. <i>Environmental Chemistry</i> , 2018, 15, 2.	1.5	5
41	Seawater recirculation through subducting sediments sustains a deeply buried population of sulfate-reducing bacteria. <i>Geobiology</i> , 2019, 17, 172-184.	2.4	5
42	Characterisation of uranium-pyrite associations within organic-rich Eocene sediments using EM, XFM- $\mu$ XANES and $\mu$ XRD. <i>Ore Geology Reviews</i> , 2021, 133, 104051.	2.7	5
43	Effects of Environmental Parameters on Thiocyanate Biodegradation by <i>Burkholderia phytofirmans</i> Candidate Strain ST01hv. <i>Environmental Engineering Science</i> , 2018, 35, 62-66.	1.6	4
44	Subsurface carbon monoxide oxidation capacity revealed through genome-resolved metagenomics of a carboxydotroph. <i>Environmental Microbiology Reports</i> , 2020, 12, 525-533.	2.4	3
45	COUPLED NITROGEN AND OXYGEN ISOTOPE STUDY OF NITRATE AT A RURAL UNLINED LANDFILL IN WESTERN VICTORIA, AUSTRALIA. <i>American Journal of Environmental Sciences</i> , 2014, 10, 383-390.	0.5	2
46	Bacterial predation limits microbial sulfate-reduction in a coastal acid sulfate soil (CASS) ecosystem. <i>Soil Biology and Biochemistry</i> , 2020, 148, 107930.	8.8	2
47	Can Bacteria Living Underground Help Fight Climate Change?. <i>Frontiers for Young Minds</i> , 0, 6, .	0.8	0