## Erin K Field

## List of Publications by Year in descending order

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687363 839539 19 807 13 18 h-index citations g-index papers 19 19 19 1321 citing authors docs citations times ranked all docs

#	Article	IF	Citations
1	Orange leads to black: evaluating the efficacy of coâ€culturing ironâ€oxidizing and sulfateâ€reducing bacteria to discern ecological relationships. Environmental Microbiology Reports, 2021, 13, 317-324.	2.4	O
2	Introducing a "core steel microbiome―and community functional analysis associated with microbially influenced corrosion. FEMS Microbiology Ecology, 2020, 97, .	2.7	8
3	A Shallow Water Ferrous-Hulled Shipwreck Reveals a Distinct Microbial Community. Frontiers in Microbiology, 2020, 11, 1897.	3.5	11
4	Iron Flocs and the Three Domains: Microbial Interactions in Freshwater Iron Mats. MBio, 2020, 11, .	4.1	5
5	The effects of temperature on Bosmina longirostris susceptibility to microcystin-LR acute toxicity. PLoS ONE, 2019, 14, e0219342.	2.5	3
6	Hydrologic Shifts Create Complex Transient Distributions of Particulate Organic Carbon and Biogeochemical Responses in Beach Aquifers. Journal of Geophysical Research G: Biogeosciences, 2019, 124, 3024-3038.	3.0	29
7	Environmental Evidence for and Genomic Insight into the Preference of Iron-Oxidizing Bacteria for More-Corrosion-Resistant Stainless Steel at Higher Salinities. Applied and Environmental Microbiology, 2019, 85, .	3.1	11
8	Carbon-dependent chromate toxicity mechanism in an environmental Arthrobacter isolate. Journal of Hazardous Materials, 2018, 355, 162-169.	12.4	20
9	Genomic exploration of individual giant ocean viruses. ISME Journal, 2017, 11, 1736-1745.	9.8	40
10	Novel Pelagic Iron-Oxidizing Zetaproteobacteria from the Chesapeake Bay Oxic–Anoxic Transition Zone. Frontiers in Microbiology, 2017, 8, 1280.	3.5	72
11	Single cell genomics indicates horizontal gene transfer and viral infections in a deep subsurface Firmicutes population. Frontiers in Microbiology, 2015, 6, 349.	3.5	61
12	Nanoarchaeota, Their Sulfolobales Host, and Nanoarchaeota Virus Distribution across Yellowstone National Park Hot Springs. Applied and Environmental Microbiology, 2015, 81, 7860-7868.	3.1	63
13	Genomic insights into the uncultivated marine <i>Zetaproteobacteria</i> at Loihi Seamount. ISME Journal, 2015, 9, 857-870.	9.8	69
14	Genomic and Metabolic Diversity of Marine Group I Thaumarchaeota in the Mesopelagic of Two Subtropical Gyres. PLoS ONE, 2014, 9, e95380.	2.5	95
15	Hexavalent chromium reduction by Cellulomonas sp. strain ES6: the influence of carbon source, iron minerals, and electron shuttling compounds. Biodegradation, 2013, 24, 437-450.	3.0	44
16	Comparative genomics of freshwater Fe-oxidizing bacteria: implications for physiology, ecology, and systematics. Frontiers in Microbiology, 2013, 4, 254.	3.5	188
17	Influence of carbon sources and electron shuttles on ferric iron reduction by Cellulomonas sp. strain ES6. Biodegradation, 2011, 22, 983-995.	3.0	18
18	UO speciation determines uranium toxicity and bioaccumulation in an environmental <i>Pseudomonas</i> sp. isolate. Environmental Toxicology and Chemistry, 2010, 29, 763-769.	4.3	31

#	Article	IF	CITATIONS
19	Application of Molecular Techniques To Elucidate the Influence of Cellulosic Waste on the Bacterial Community Structure at a Simulated Low-Level-Radioactive-Waste Site. Applied and Environmental Microbiology, 2010, 76, 3106-3115.	3.1	39