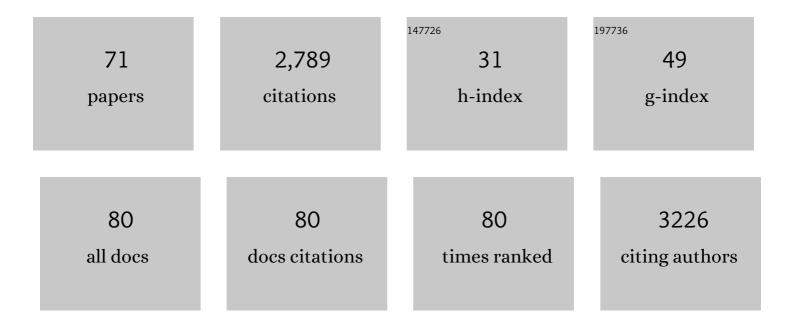
## **Trinity Hamilton**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3488194/publications.pdf Version: 2024-02-01



Τρινιτν Ηλμιτον

#	Article	IF	CITATIONS
1	Meet Me in the Middle: Median Temperatures Impact Cyanobacteria and Photoautotrophy in Eruptive Yellowstone Hot Springs. MSystems, 2022, 7, e0145021.	1.7	6
2	Diversity and distribution of sediment bacteria across an ecological and trophic gradient. PLoS ONE, 2022, 17, e0258079.	1.1	3
3	Characterization of diverse bacteriohopanepolyols in a permanently stratified, hyper-euxinic lake. Organic Geochemistry, 2022, 168, 104431.	0.9	3
4	Temperature and Geographic Location Impact the Distribution and Diversity of Photoautotrophic Gene Variants in Alkaline Yellowstone Hot Springs. Microbiology Spectrum, 2022, 10, e0146521.	1.2	7
5	Recharge from glacial meltwater is critical for alpine springs and their microbiomes. Environmental Research Letters, 2021, 16, 064012.	2.2	8
6	Genomics, Exometabolomics, and Metabolic Probing Reveal Conserved Proteolytic Metabolism of Thermoflexus hugenholtzii and Three Candidate Species From China and Japan. Frontiers in Microbiology, 2021, 12, 632731.	1.5	8
7	Environmental DNA reveals arboreal cityscapes at the Ancient Maya Center of Tikal. Scientific Reports, 2021, 11, 12725.	1.6	16
8	Metagenome-Assembled Genomes of Novel Taxa from an Acid Mine Drainage Environment. Applied and Environmental Microbiology, 2021, 87, e0077221.	1.4	9
9	Biological albedo reduction on ice sheets, glaciers, and snowfields. Earth-Science Reviews, 2021, 220, 103728.	4.0	30
10	Hot Spring Microbial Community Elemental Composition: Hot Spring and Soil Inputs, and the Transition from Biocumulus to Siliceous Sinter. Astrobiology, 2021, 21, 1526-1546.	1.5	6
11	Inorganic carbon addition stimulates snow algae primary productivity. ISME Journal, 2020, 14, 857-860.	4.4	19
12	Carbon and nitrogen recycling during cyanoHABs in dreissenid-invaded and non-invaded US midwestern lakes and reservoirs. Hydrobiologia, 2020, 847, 939-965.	1.0	8
13	Trace Element Concentrations in Hydrothermal Silica Deposits as a Potential Biosignature. Astrobiology, 2020, 20, 525-536.	1.5	10
14	Metabolic diversity and co-occurrence of multiple Ferrovum species at an acid mine drainage site. BMC Microbiology, 2020, 20, 119.	1.3	9
15	Molecular genetic and geochemical assays reveal severe contamination of drinking water reservoirs at the ancient Maya city of Tikal. Scientific Reports, 2020, 10, 10316.	1.6	19
16	Temperature impacts community structure and function of phototrophic Chloroflexi and Cyanobacteria in two alkaline hot springs in Yellowstone National Park. Environmental Microbiology Reports, 2020, 12, 503-513.	1.0	52
17	The effect of woodchip bioreactors on microbial concentration in subsurface drainage water and the associated risk of antibiotic resistance dissemination. Ecological Engineering: X, 2020, 143, 100017.	3.5	4
18	Productivity and Community Composition of Low Biomass/High Silica Precipitation Hot Springs: A Possible Window to Earth's Early Biosphere?. Life, 2019, 9, 64.	1.1	18

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19	Matrotrophic viviparity constrains microbiome acquisition during gestation in a liveâ€bearing cockroach, Diploptera punctata. Ecology and Evolution, 2019, 9, 10601-10614.	0.8	6
20	The Antarctic mite, Alaskozetes antarcticus, shares bacterial microbiome community membership but not abundance between adults and tritonymphs. Polar Biology, 2019, 42, 2075-2085.	0.5	2
21	Hypolithic Photosynthesis in Hydrothermal Areas and Implications for Cryptic Oxygen Oases on Archean Continental Surfaces. Frontiers in Earth Science, 2019, 7, .	0.8	9
22	The trouble with oxygen: The ecophysiology of extant phototrophs and implications for the evolution of oxygenic photosynthesis. Free Radical Biology and Medicine, 2019, 140, 233-249.	1.3	38
23	Anoxygenic Phototrophs Span Geochemical Gradients and Diverse Morphologies in Terrestrial Geothermal Springs. MSystems, 2019, 4, .	1.7	24
24	Snow algae drive productivity and weathering at volcanic rock-hosted glaciers. Geochimica Et Cosmochimica Acta, 2019, 247, 220-242.	1.6	26
25	Cyanobacterial photosynthesis under sulfidic conditions: insights from the isolate <i>Leptolyngbya</i> sp. strain hensonii. ISME Journal, 2018, 12, 568-584.	4.4	50
26	Geobiological feedbacks and the evolution of thermoacidophiles. ISME Journal, 2018, 12, 225-236.	4.4	70
27	Water column and sediment stable carbon isotope biogeochemistry of permanently redoxâ€stratified Fayetteville Green Lake, New York, U.S.A Limnology and Oceanography, 2018, 63, 570-587.	1.6	26
28	Low-Light Anoxygenic Photosynthesis and Fe-S-Biogeochemistry in a Microbial Mat. Frontiers in Microbiology, 2018, 9, 858.	1.5	19
29	Silica Dissolution and Precipitation in Glaciated Volcanic Environments and Implications for Mars. Geophysical Research Letters, 2018, 45, 7371-7381.	1.5	22
30	Effect of salinity on mercury methylating benthic microbes and their activities in Great Salt Lake, Utah. Science of the Total Environment, 2017, 581-582, 495-506.	3.9	40
31	Primary productivity of snow algae communities on stratovolcanoes of the Pacific Northwest. Geobiology, 2017, 15, 280-295.	1.1	54
32	Microbial ecology of mountain glacier ecosystems: biodiversity, ecological connections and implications of a warming climate. Environmental Microbiology, 2017, 19, 2935-2948.	1.8	130
33	Oxygenic and anoxygenic photosynthesis in a microbial mat from an anoxic and sulfidic spring. Environmental Microbiology, 2017, 19, 1251-1265.	1.8	18
34	Microbial communities and organic biomarkers in a Proterozoicâ€analog sinkhole. Geobiology, 2017, 15, 784-797.	1.1	14
35	Draft Genome Sequence of Anoxybacillus ayderensis Strain MT-Cab ( Firmicutes ). Genome Announcements, 2017, 5, .	0.8	3
36	Sulfur and carbon isotopic evidence for metabolic pathway evolution and a four-stepped Earth system progression across the Archean and Paleoproterozoic. Earth-Science Reviews, 2017, 174, 1-21.	4.0	58

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37	Hot Spring Microbial Community Composition, Morphology, and Carbon Fixation: Implications for Interpreting the Ancient Rock Record. Frontiers in Earth Science, 2017, 5, .	0.8	50
38	The Physiological Functions and Structural Determinants of Catalytic Bias in the [FeFe]-Hydrogenases Cpl and CpII of Clostridium pasteurianum Strain W5. Frontiers in Microbiology, 2017, 8, 1305.	1.5	30
39	Geochemistry and microbial community composition across a range of acid mine drainage impact and implications for the Neoarcheanâ&Paleoproterozoic transition. Journal of Geophysical Research G: Biogeosciences, 2017, 122, 1404-1422.	1.3	12
40	Carbon and Sulfur Cycling below the Chemocline in a Meromictic Lake and the Identification of a Novel Taxonomic Lineage in the FCB Superphylum, Candidatus Aegiribacteria. Frontiers in Microbiology, 2016, 7, 598.	1.5	51
41	Substrate preference, uptake kinetics and bioenergetics in a facultatively autotrophic, thermoacidophilic crenarchaeote. FEMS Microbiology Ecology, 2016, 92, fiw069.	1.3	10
42	The role of biology in planetary evolution: cyanobacterial primary production in lowâ€oxygen Proterozoic oceans. Environmental Microbiology, 2016, 18, 325-340.	1.8	151
43	Aerobic and Anaerobic Thiosulfate Oxidation by a Cold-Adapted, Subglacial Chemoautotroph. Applied and Environmental Microbiology, 2016, 82, 1486-1495.	1.4	62
44	The behavior of biologically important trace elements across the oxic/euxinic transition of meromictic Fayetteville Green Lake, New York, USA. Geochimica Et Cosmochimica Acta, 2015, 165, 389-406.	1.6	52
45	Metabolic diversity and ecological niches of Achromatium populations revealed with single-cell genomic sequencing. Frontiers in Microbiology, 2015, 6, 822.	1.5	20
46	Evolution of Molybdenum Nitrogenase during the Transition from Anaerobic to Aerobic Metabolism. Journal of Bacteriology, 2015, 197, 1690-1699.	1.0	97
47	Rock comminution as a source of hydrogen for subglacial ecosystems. Nature Geoscience, 2015, 8, 851-855.	5.4	82
48	[FeFe]-Hydrogenase Abundance and Diversity along a Vertical Redox Gradient in Great Salt Lake, USA. International Journal of Molecular Sciences, 2014, 15, 21947-21966.	1.8	17
49	Draft Genome Sequence of the Moderately Thermophilic Bacterium Schleiferia thermophila Strain Yellowstone ( <i>Bacteroidetes</i> ). Genome Announcements, 2014, 2, .	0.8	13
50	Draft Genome Sequence of a Sulfide-Oxidizing, Autotrophic Filamentous Anoxygenic Phototrophic Bacterium, <i>Chloroflexus</i> sp. Strain MS-G ( <i>Chloroflexi</i> ). Genome Announcements, 2014, 2,	0.8	18
51	Coupled reductive and oxidative sulfur cycling in the phototrophic plate of a meromictic lake. Geobiology, 2014, 12, 451-468.	1.1	45
52	Chemolithotrophic Primary Production in a Subglacial Ecosystem. Applied and Environmental Microbiology, 2014, 80, 6146-6153.	1.4	92
53	Competition for Ammonia Influences the Structure of Chemotrophic Communities in Geothermal Springs. Applied and Environmental Microbiology, 2014, 80, 653-661.	1.4	46
54	Metagenomic insights into S(0) precipitation in a terrestrial subsurface lithoautotrophic ecosystem. Frontiers in Microbiology, 2014, 5, 756.	1.5	75

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55	Energy, ecology and the distribution of microbial life. Philosophical Transactions of the Royal Society B: Biological Sciences, 2013, 368, 20120383.	1.8	28
56	Molecular evidence for an active endogenous microbiome beneath glacial ice. ISME Journal, 2013, 7, 1402-1412.	4.4	116
57	The Role of Tetraether Lipid Composition in the Adaptation of Thermophilic Archaea to Acidity. Frontiers in Microbiology, 2013, 4, 62.	1.5	69
58	Environmental constraints defining the distribution, composition, and evolution of chlorophototrophs in thermal features of Yellowstone National Park. Geobiology, 2012, 10, 236-249.	1.1	42
59	Radical AdoMet enzymes in complex metal cluster biosynthesis. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2012, 1824, 1254-1263.	1.1	25
60	Transcriptional Profiling of Nitrogen Fixation in Azotobacter vinelandii. Journal of Bacteriology, 2011, 193, 4477-4486.	1.0	99
61	An Alternative Path for the Evolution of Biological Nitrogen Fixation. Frontiers in Microbiology, 2011, 2, 205.	1.5	105
62	Biological nitrogen fixation in acidic highâ€ŧemperature geothermal springs in Yellowstone National Park, Wyoming. Environmental Microbiology, 2011, 13, 2204-2215.	1.8	45
63	A late methanogen origin for molybdenumâ€dependent nitrogenase. Geobiology, 2011, 9, 221-232.	1.1	141
64	Environmental Constraints Underpin the Distribution and Phylogenetic Diversity of nifH in the Yellowstone Geothermal Complex. Microbial Ecology, 2011, 61, 860-870.	1.4	40
65	Diversity, Abundance, and Potential Activity of Nitrifying and Nitrate-Reducing Microbial Assemblages in a Subglacial Ecosystem. Applied and Environmental Microbiology, 2011, 77, 4778-4787.	1.4	119
66	FAD Binding by ApbE Protein from <i>Salmonella enterica</i> : a New Class of FAD-Binding Proteins. Journal of Bacteriology, 2011, 193, 887-895.	1.0	36
67	Differential Accumulation of <i>nif</i> Structural Gene mRNA in Azotobacter vinelandii. Journal of Bacteriology, 2011, 193, 4534-4536.	1.0	11
68	[FeFe]-hydrogenase in Yellowstone National Park: evidence for dispersal limitation and phylogenetic niche conservatism. ISME Journal, 2010, 4, 1485-1495.	4.4	63
69	Crystal Structure of the L Protein of <i>Rhodobacter sphaeroides</i> Light-Independent Protochlorophyllide Reductase with MgADP Bound: A Homologue of the Nitrogenase Fe Protein. Biochemistry, 2008, 47, 13004-13015.	1.2	66
70	Cloning, sequence analysis and confirmation of derived gene sequences for three epitope-mapped monoclonal antibodies against human phagocyte flavocytochrome b. Molecular Immunology, 2007, 44, 625-637.	1.0	6
71	Paleoecological Studies at the Ancient Maya Center of Yaxnohcah Using Analyses of Pollen, Environmental DNA, and Plant Macroremains. Frontiers in Ecology and Evolution, 0, 10, .	1.1	4