

Yuki Morono

List of Publications by Year in descending order

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Version: 2024-02-01

111
papers

5,132
citations

117625

34
h-index

106344

65
g-index

118
all docs

118
docs citations

118
times ranked

4812
citing authors

#	ARTICLE	IF	CITATIONS
1	Significant contribution of Archaea to extant biomass in marine subsurface sediments. <i>Nature</i> , 2008, 454, 991-994.	27.8	583
2	Isolation of an archaeon at the prokaryote–eukaryote interface. <i>Nature</i> , 2020, 577, 519-525.	27.8	449
3	Exploring deep microbial life in coal-bearing sediment down to ~2.5 km below the ocean floor. <i>Science</i> , 2015, 349, 420-424.	12.6	376
4	Presence of oxygen and aerobic communities from sea floor to basement in deep-sea sediments. <i>Nature Geoscience</i> , 2015, 8, 299-304.	12.9	226
5	Carbon and nitrogen assimilation in deep subseafloor microbial cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 18295-18300.	7.1	205
6	Global diversity of microbial communities in marine sediment. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 27587-27597.	7.1	174
7	Discriminative detection and enumeration of microbial life in marine subsurface sediments. <i>ISME Journal</i> , 2009, 3, 503-511.	9.8	140
8	An improved cell separation technique for marine subsurface sediments: applications for high-throughput analysis using flow cytometry and cell sorting. <i>Environmental Microbiology</i> , 2013, 15, 2841-2849.	3.8	119
9	Cultivation of methanogenic community from subseafloor sediments using a continuous-flow bioreactor. <i>ISME Journal</i> , 2011, 5, 1913-1925.	9.8	108
10	Sedimentary membrane lipids recycled by deep-sea benthic archaea. <i>Nature Geoscience</i> , 2010, 3, 858-861.	12.9	103
11	Dehalogenation Activities and Distribution of Reductive Dehalogenase Homologous Genes in Marine Subsurface Sediments. <i>Applied and Environmental Microbiology</i> , 2009, 75, 6905-6909.	3.1	95
12	Methyl-compound use and slow growth characterize microbial life in 2-km-deep subseafloor coal and shale beds. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E9206-E9215.	7.1	94
13	Phylogenetic and enzymatic diversity of deep subseafloor aerobic microorganisms in organics- and methane-rich sediments off Shimokita Peninsula. <i>Extremophiles</i> , 2008, 12, 519-527.	2.3	93
14	Acetogenesis in Deep Subseafloor Sediments of The Juan de Fuca Ridge Flank: A Synthesis of Geochemical, Thermodynamic, and Gene-based Evidence. <i>Geomicrobiology Journal</i> , 2010, 27, 183-211.	2.0	89
15	A Modified SDS-Based DNA Extraction Method for High Quality Environmental DNA from Seafloor Environments. <i>Frontiers in Microbiology</i> , 2016, 07, 986.	3.5	80
16	Deep-biosphere methane production stimulated by geofluids in the Nankai accretionary complex. <i>Science Advances</i> , 2018, 4, eaao4631.	10.3	79
17	Microbial Diversity in Sediments from the Bottom of the Challenger Deep, the Mariana Trench. <i>Microbes and Environments</i> , 2018, 33, 186-194.	1.6	75
18	Bioturbation as a key driver behind the dominance of Bacteria over Archaea in near-surface sediment. <i>Scientific Reports</i> , 2017, 7, 2400.	3.3	73

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19	Endospore abundance and d:l-amino acid modeling of bacterial turnover in holocene marine sediment (Aarhus Bay). <i>Geochimica Et Cosmochimica Acta</i> , 2012, 99, 87-99.	3.9	72
20	Aerobic microbial life persists in oxic marine sediment as old as 101.5 million years. <i>Nature Communications</i> , 2020, 11, 3626.	12.8	72
21	Temperature limits to deep seafloor life in the Nankai Trough subduction zone. <i>Science</i> , 2020, 370, 1230-1234.	12.6	65
22	Microbial dormancy in the marine subsurface: Global endospore abundance and response to burial. <i>Science Advances</i> , 2019, 5, eaav1024.	10.3	64
23	Niche Separation of Methanotrophic Archaea (ANME-1 and -2) in Methane-Seep Sediments of the Eastern Japan Sea Offshore Joetsu. <i>Geomicrobiology Journal</i> , 2011, 28, 118-129.	2.0	61
24	High frequency of phylogenetically diverse reductive dehalogenase-homologous genes in deep seafloor sedimentary metagenomes. <i>Frontiers in Microbiology</i> , 2014, 5, 80.	3.5	61
25	Magmatism, serpentinization and life: Insights through drilling the Atlantis Massif (IODP Expedition) Tj ETQq1 1 0.784314 rgBT /Overl	1.4	58
26	Atribacteria from the Subseafloor Sedimentary Biosphere Disperse to the Hydrosphere through Submarine Mud Volcanoes. <i>Frontiers in Microbiology</i> , 2017, 8, 1135.	3.5	55
27	Persistent organic matter in oxic seafloor sediment. <i>Nature Geoscience</i> , 2019, 12, 126-131.	12.9	53
28	Microbiological Assessment of Circulation Mud Fluids During the First Operation of Riser Drilling by the Deep-Earth Research Vessel Chikyū. <i>Geomicrobiology Journal</i> , 2008, 25, 274-282.	2.0	51
29	Metabolically active microbial communities in marine sediment under high-CO ₂ and low-pH extremes. <i>ISME Journal</i> , 2013, 7, 555-567.	9.8	51
30	Hot-Alkaline DNA Extraction Method for Deep-Subseafloor Archaeal Communities. <i>Applied and Environmental Microbiology</i> , 2014, 80, 1985-1994.	3.1	49
31	Variance and potential niche separation of microbial communities in seafloor sediments off Shimokita Peninsula, Japan. <i>Environmental Microbiology</i> , 2016, 18, 1889-1906.	3.8	48
32	Bacterial dominance in seafloor sediments characterized by methane hydrates. <i>FEMS Microbiology Ecology</i> , 2012, 81, 88-98.	2.7	46
33	Dense microbial community on a ferromanganese nodule from the ultra-oligotrophic South Pacific Gyre: Implications for biogeochemical cycles. <i>Earth and Planetary Science Letters</i> , 2016, 447, 10-20.	4.4	41
34	Characterization of Metabolically Active Bacterial Populations in Subseafloor Nankai Trough Sediments above, within, and below the Sulfate-Methane Transition Zone. <i>Frontiers in Microbiology</i> , 2012, 3, 113.	3.5	39
35	Origins of lithium in submarine mud volcano fluid in the Nankai accretionary wedge. <i>Earth and Planetary Science Letters</i> , 2015, 414, 144-155.	4.4	37
36	Naturally occurring, microbially induced smectite-to-illite reaction. <i>Geology</i> , 2019, 47, 535-539.	4.4	37

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37	Distribution of dehalogenation activity in subseafloor sediments of the Nankai Trough subduction zone. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2013, 368, 20120249.	4.0	35
38	Application of glutaraldehyde for the staining of esterase-active cells with carboxyfluorescein diacetate. <i>Biotechnology Letters</i> , 2004, 26, 379-383.	2.2	32
39	Addition of Aromatic Substrates Restores Trichloroethylene Degradation Activity in <i>Pseudomonas putida</i> F1. <i>Applied and Environmental Microbiology</i> , 2004, 70, 2830-2835.	3.1	30
40	Predominance of Viable Spore-Forming Piezophilic Bacteria in High-Pressure Enrichment Cultures from ~1.5 to 2.4 km-Deep Coal-Bearing Sediments below the Ocean Floor. <i>Frontiers in Microbiology</i> , 2017, 8, 137.	3.5	30
41	Deep microbial proliferation at the basalt interface in 33.5±104 million-year-old oceanic crust. <i>Communications Biology</i> , 2020, 3, 136.	4.4	29
42	A new DNA extraction method by controlled alkaline treatments from consolidated subsurface sediments. <i>FEMS Microbiology Letters</i> , 2012, 326, 47-54.	1.8	28
43	Analysis of Low-Biomass Microbial Communities in the Deep Biosphere. <i>Advances in Applied Microbiology</i> , 2016, 95, 149-178.	2.4	28
44	Accessing the Subsurface Biosphere Within Rocks Undergoing Active Low-Temperature Serpentinization in the Samail Ophiolite (Oman Drilling Project). <i>Journal of Geophysical Research G: Biogeosciences</i> , 2021, 126, e2021JG006315.	3.0	27
45	Variability of subseafloor viral abundance at the geographically and geologically distinct continental margins. <i>FEMS Microbiology Ecology</i> , 2014, 88, 60-68.	2.7	26
46	Correlation of TCE cometabolism with growth characteristics on aromatic substrates in toluene-degrading bacteria. <i>Biochemical Engineering Journal</i> , 2006, 31, 173-179.	3.6	25
47	Comparative Study of Subseafloor Microbial Community Structures in Deeply Buried Coral Fossils and Sediment Matrices from the Challenger Mound in the Porcupine Seabight. <i>Frontiers in Microbiology</i> , 2011, 2, 231.	3.5	25
48	Optimization of distinction between viable and dead cells by fluorescent staining method and its application to bacterial consortia. <i>Biochemical Engineering Journal</i> , 2007, 37, 56-61.	3.6	24
49	Biological CO ₂ conversion to acetate in subsurface coal-sand formation using a high-pressure reactor system. <i>Frontiers in Microbiology</i> , 2013, 4, 361.	3.5	24
50	Size and Carbon Content of Sub-seafloor Microbial Cells at Landsort Deep, Baltic Sea. <i>Frontiers in Microbiology</i> , 2016, 7, 1375.	3.5	24
51	Cell-Specific Thioautotrophic Productivity of Epsilon-Proteobacterial Epibionts Associated with <i>Shinkaia crosnieri</i> . <i>PLoS ONE</i> , 2012, 7, e46282.	2.5	23
52	D:L-Amino Acid Modeling Reveals Fast Microbial Turnover of Days to Months in the Subsurface Hydrothermal Sediment of Guaymas Basin. <i>Frontiers in Microbiology</i> , 2018, 9, 967.	3.5	23
53	Intact preservation of environmental samples by freezing under an alternating magnetic field. <i>Environmental Microbiology Reports</i> , 2015, 7, 243-251.	2.4	22
54	Significant contribution of subseafloor microparticles to the global manganese budget. <i>Nature Communications</i> , 2019, 10, 400.	12.8	22

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55	An Improved Method for Extracting Viruses From Sediment: Detection of Far More Viruses in the Subseafloor Than Previously Reported. <i>Frontiers in Microbiology</i> , 2019, 10, 878.	3.5	21
56	An improved method to identify osmium-stained organic matter within soil aggregate structure by electron microscopy and synchrotron X-ray micro-computed tomography. <i>Soil and Tillage Research</i> , 2019, 191, 275-281.	5.6	21
57	Rapid metabolism fosters microbial survival in the deep, hot subseafloor biosphere. <i>Nature Communications</i> , 2022, 13, 312.	12.8	21
58	Ecophysiology of Zetaproteobacteria Associated with Shallow Hydrothermal Iron-Oxyhydroxide Deposits in Nagahama Bay of Satsuma Iwo-Jima, Japan. <i>Frontiers in Microbiology</i> , 2015, 6, 1554.	3.5	20
59	Cellular content of biomolecules in sub-seafloor microbial communities. <i>Geochimica Et Cosmochimica Acta</i> , 2016, 188, 330-351.	3.9	20
60	Automatic Slide-Loader Fluorescence Microscope for Discriminative Enumeration of Subseafloor Life. <i>Scientific Drilling</i> , 0, 9, 32-36.	0.6	20
61	Cool, alkaline serpentinite formation fluid regime with scarce microbial habitability and possible abiotic synthesis beneath the South Chamorro Seamount. <i>Progress in Earth and Planetary Science</i> , 2018, 5, .	3.0	19
62	Kinetic analyses of trichloroethylene cometabolism by toluene-degrading bacteria harboring a tod homologous gene. <i>Biochemical Engineering Journal</i> , 2005, 26, 59-64.	3.6	18
63	Aerobic and Anaerobic Methanotrophic Communities Associated with Methane Hydrates Exposed on the Seafloor: A High-Pressure Sampling and Stable Isotope-Incubation Experiment. <i>Frontiers in Microbiology</i> , 2017, 8, 2569.	3.5	18
64	Gold-ISH: A nano-size gold particle-based phylogenetic identification compatible with NanoSIMS. <i>Systematic and Applied Microbiology</i> , 2014, 37, 261-266.	2.8	17
65	Cultivable microbial community in 2-km-deep, 20-million-year-old subseafloor coalbeds through ~1000 days anaerobic bioreactor cultivation. <i>Scientific Reports</i> , 2019, 9, 2305.	3.3	17
66	Expedition 357 summary. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	16
67	Assessment of Capacity to Capture DNA Aerosols by Clean Filters for Molecular Biology Experiments. <i>Microbes and Environments</i> , 2018, 33, 222-226.	1.6	14
68	Expedition 385 methods. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	14
69	Microbial Metabolism and Community Dynamics in Hydraulic Fracturing Fluids Recovered From Deep Hydrocarbon-Rich Shale. <i>Frontiers in Microbiology</i> , 2019, 10, 376.	3.5	13
70	Site U1545. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	13
71	An improved sample preparation method for imaging microstructures of fine-grained marine sediment using microfocuss X-ray computed tomography and scanning electron microscopy. <i>Limnology and Oceanography: Methods</i> , 2014, 12, 469-483.	2.0	12
72	Geophysical constraints on microbial biomass in subseafloor sediments and coal seams down to 2.5 km off Shimokita Peninsula, Japan. <i>Progress in Earth and Planetary Science</i> , 2018, 5, .	3.0	12

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73	Origin of Short-Chain Organic Acids in Serpentinite Mud Volcanoes of the Mariana Convergent Margin. <i>Frontiers in Microbiology</i> , 2019, 10, 1729.	3.5	11
74	High Fluid-Pressure Patches Beneath the DÃ©collement: A Potential Source of Slow Earthquakes in the Nankai Trough off Cape Muroto. <i>Journal of Geophysical Research: Solid Earth</i> , 2021, 126, e2021JB021831.	3.4	11
75	Evolution of (Bio-Geochemical Processes and Diagenetic Alteration of Sediments Along the Tectonic Migration of Ocean Floor in the Shikoku Basin off Japan. <i>Geochemistry, Geophysics, Geosystems</i> , 2021, 22, e2020GC009585.	2.5	11
76	Site U1546. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	11
77	Expedition 357 methods. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	11
78	Bio-Archive Core Storage and Subsampling Procedure for Subseafloor Molecular Biological Research. <i>Scientific Drilling</i> , 0, 8, 35-37.	0.6	11
79	A New Method for Quality Control of Geological Cores by X-Ray Computed Tomography: Application in IODP Expedition 370. <i>Frontiers in Earth Science</i> , 2019, 7, .	1.8	10
80	The Limits of Life and the Biosphere in Earth's Interior. <i>Oceanography</i> , 2019, 32, 208-211.	1.0	10
81	Expedition 385 summary. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	10
82	Sites U1547 and U1548. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	9
83	Hot fluids, burial metamorphism and thermal histories in the underthrust sediments at IODP 370 site C0023, Nankai Accretionary Complex. <i>Marine and Petroleum Geology</i> , 2020, 112, 104080.	3.3	8
84	Expedition 370 methods. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	8
85	Site U1549. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	7
86	Biomass, Diversity, and Metabolic Functions of Subseafloor Life. <i>Developments in Marine Geology</i> , 2014, 7, 65-83.	0.4	6
87	Site U1550. <i>Proceedings of the International Ocean Discovery Program</i> , 0, , .	0.0	6
88	Archaeal MutS5 tightly binds to Holliday junction similarly to eukaryotic MutS ³ . <i>FEBS Journal</i> , 2017, 284, 3470-3483.	4.7	5
89	In-situ mechanical weakness of subducting sediments beneath a plate boundary dÃ©collement in the Nankai Trough. <i>Progress in Earth and Planetary Science</i> , 2018, 5, .	3.0	5
90	EDTA-FISH: A Simple and Effective Approach to Reduce Non-specific Adsorption of Probes in Fluorescence &in situ Hybridization (FISH) for Environmental Samples. <i>Microbes and Environments</i> , 2020, 35, n/a.	1.6	5

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91	Site C0023. Proceedings of the International Ocean Discovery Program, 0, , .	0.0	5
92	Preliminary experiment for cell count using flow cytometry. Proceedings of the Integrated Ocean Drilling Program Integrated Ocean Drilling Program, 0, , .	1.0	5
93	Site U1552. Proceedings of the International Ocean Discovery Program, 0, , .	0.0	4
94	Expedition 370 summary. Proceedings of the International Ocean Discovery Program, 0, , .	0.0	4
95	Radical Gas-Based DNA Decontamination for Ultra-Sensitive Molecular Experiments. Microbes and Environments, 2012, 27, 512-514.	1.6	3
96	Simple In-liquid Staining of Microbial Cells for Flow Cytometry Quantification of the Microbial Population in Marine Subseafloor Sediments. Microbes and Environments, 2021, 36, n/a.	1.6	3
97	Site U1551. Proceedings of the International Ocean Discovery Program, 0, , .	0.0	3
98	CO ₂ emission and shallow-type methane hydrate decomposition experiment on deep-sea floor. JAMSTEC Report of Research and Development, 2015, 20, 61-71.	0.2	3
99	5. Detecting slow metabolism in the subseafloor: analysis of single cells using NanoSIMS. , 2014, , 101-120.		2
100	Exploration of the deep-subseafloor-biosphere frontiers: Achievements and perspectives. Journal of the Geological Society of Japan, 2018, 124, 77-92.	0.6	2
101	Metal-ion-induced expression of gene fragments from subseafloor micro-organisms in the Kumano forearc basin, Nankai Trough. Journal of Applied Microbiology, 2018, 125, 1396-1407.	3.1	2
102	Crucial Scientific Issues in Earth Science Revealed Only by Mantle Drilling: Understanding the Current State of the Oceanic Plates of a Life-bearing Planet. Journal of Geography (Chigaku Zasshi), 2021, 130, 483-506.	0.3	2
103	Purification of Disc-Shaped Diatoms from the Southern Ocean Sediment by a Cell Sorter to Obtain an Accurate Oxygen Isotope Record. ACS Earth and Space Chemistry, 0, , .	2.7	2
104	Developing community-based scientific priorities and new drilling proposals in the southern Indian and southwestern Pacific oceans. Scientific Drilling, 0, 24, 61-70.	0.6	2
105	Increase in acetate concentrations during sediment sample onboard storage: a caution for pore-water geochemical analyses. Geochemical Journal, 2013, 47, 567-571.	1.0	1
106	Osmium Plasma Coating for Observation of Microfossils, Using Optical and Scanning Electron Microscopes. Paleontological Research, 2016, 20, 296-301.	1.0	1
107	Modelling the Shimokita deep coalbed biosphere over deep geological time: Starvation, stimulation, material balance and population models. Basin Research, 2020, 32, 804-829.	2.7	1
108	Data report: water activity of the deep coal-bearing basin off Shimokita from IODP Expedition 337. Proceedings of the Integrated Ocean Drilling Program Integrated Ocean Drilling Program, 0, , .	1.0	1

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109	“Mark the Gene”: a Method for Nondestructive Introduction of Marker Sequences Inside the Gene Frame of Transgenes. Applied and Environmental Microbiology, 2007, 73, 4915-4921.	3.1	0
110	Forging Partnerships with Other Federal Programs: NASA and the National Science Foundation (NSF) through Scientific Ocean Drilling. , 2021, 53, .		0
111	Construction of Aerobic/Anaerobic-Substrate-Induced Gene Expression Procedure for Exploration of Metagenomes From Subseafloor Sediments. Frontiers in Microbiology, 2021, 12, 726024.	3.5	0