

Nicolas Tsesmetzis

List of Publications by Year in descending order

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Version: 2024-02-01

28
papers

1,085
citations

516215

16
h-index

610482

24
g-index

29
all docs

29
docs citations

29
times ranked

1770
citing authors

#	ARTICLE	IF	CITATIONS
1	IMG/VR: a database of cultured and uncultured DNA Viruses and retroviruses. <i>Nucleic Acids Research</i> , 2016, 45, D457-D465.	6.5	177
2	Succession in the petroleum reservoir microbiome through an oil field production lifecycle. <i>ISME Journal</i> , 2017, 11, 2141-2154.	4.4	136
3	Complementary Microorganisms in Highly Corrosive Biofilms from an Offshore Oil Production Facility. <i>Applied and Environmental Microbiology</i> , 2016, 82, 2545-2554.	1.4	135
4	Damage to offshore production facilities by corrosive microbial biofilms. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 2525-2533.	1.7	70
5	Beyond the tip of the iceberg; a new view of the diversity of sulfite- and sulfate-reducing microorganisms. <i>ISME Journal</i> , 2018, 12, 2096-2099.	4.4	67
6	Molecular cloning and characterization of an almond 9-hydroperoxide lyase, a new CYP74 targeted to lipid bodies*. <i>Journal of Experimental Botany</i> , 2005, 56, 2321-2333.	2.4	54
7	Arabidopsis Reactome: A Foundation Knowledgebase for Plant Systems Biology. <i>Plant Cell</i> , 2008, 20, 1426-1436.	3.1	52
8	Comparative metagenomics of hydrocarbon and methane seeps of the Gulf of Mexico. <i>Scientific Reports</i> , 2017, 7, 16015.	1.6	52
9	Will Membranes Break Barriers on Volatile Fatty Acid Recovery from Anaerobic Digestion?. <i>ACS ES&T Engineering</i> , 2021, 1, 141-153.	3.7	39
10	Microbial redox processes in deep subsurface environments and the potential application of (per)chlorate in oil reservoirs. <i>Frontiers in Microbiology</i> , 2014, 5, 428.	1.5	37
11	Dynamic bacterial and fungal microbiomes during sweet sorghum ensiling impact bioethanol production. <i>Bioresource Technology</i> , 2018, 264, 163-173.	4.8	37
12	Bioelectrochemical remediation of phenanthrene in a microbial fuel cell using an anaerobic consortium enriched from a hydrocarbon-contaminated site. <i>Journal of Hazardous Materials</i> , 2020, 389, 121845.	6.5	32
13	Subcellular localisation of <i>Medicago truncatula</i> 9/13-hydroperoxide lyase reveals a new localisation pattern and activation mechanism for CYP74C enzymes. <i>BMC Plant Biology</i> , 2007, 7, 58.	1.6	30
14	Perchlorate and chlorate reduction by the <i>Crenarchaeon</i> <i>Aeropyrum pernix</i> and two thermophilic <i>Ferromicrobium</i> .	1.0	30
15	Microbial and Isotopic Evidence for Methane Cycling in Hydrocarbon-Containing Groundwater from the Pennsylvania Region. <i>Frontiers in Microbiology</i> , 2017, 8, 593.	1.5	30
16	Contrasting Pathways for Anaerobic Methane Oxidation in Gulf of Mexico Cold Seep Sediments. <i>MSystems</i> , 2019, 4, .	1.7	27
17	Microbial community analysis of three hydrocarbon reservoir cores provides valuable insights for the assessment of reservoir souring potential. <i>International Biodeterioration and Biodegradation</i> , 2018, 126, 177-188.	1.9	15
18	Preservation of ancestral Cretaceous microflora recovered from a hypersaline oil reservoir. <i>Scientific Reports</i> , 2016, 6, 22960.	1.6	14

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19	Syngas mediated microbial electrosynthesis for CO ₂ to acetate conversion using <i>Clostridium ljungdahlii</i> . <i>Resources, Conservation and Recycling</i> , 2022, 184, 106395.	5.3	14
20	Proposal of Improved Biomonitoring Standard for Purpose of Microbiologically Influenced Corrosion Risk Assessment. , 2016, , .		8
21	Syntrophic Hydrocarbon Degradation in a Decommissioned Off-Shore Subsea Oil Storage Structure. <i>Microorganisms</i> , 2021, 9, 356.	1.6	7
22	Diagnosing microbiologically influenced corrosion at a crude oil pipeline facility leak site â€” A multiple lines of evidence approach. <i>International Biodeterioration and Biodegradation</i> , 2022, 172, 105438.	1.9	7
23	Electrocatalytic Membranes for Tunable Syngas Production and High-Efficiency Delivery to Biocompatible Electrolytes. <i>ACS Sustainable Chemistry and Engineering</i> , 2021, 9, 6012-6022.	3.2	6
24	The gateway pDEST17 expression vector encodes a âˆ²1 ribosomal frameshifting sequence. <i>Nucleic Acids Research</i> , 2007, 35, 1322-1332.	6.5	5
25	MlxS-HCR: a MlxS extension defining a minimal information standard for sequence data from environments pertaining to hydrocarbon resources. <i>Standards in Genomic Sciences</i> , 2016, 11, 78.	1.5	2
26	MetaHCR: a web-enabled metagenome data management system for hydrocarbon resources. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, 1-10.	1.4	1
27	TheArabidopsis Localizome: Subcellular Protein Localization and Interactions inARABIDOPSIS. , 0, , 61-81.		0
28	Using genomics for environmental monitoring in the oil and gas industry. <i>Integrated Environmental Assessment and Management</i> , 2017, 13, 797-799.	1.6	0