Carlo R Carere

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/610224/publications.pdf

Version: 2024-02-01

21 papers 1,995 citations

15 h-index 713332 21 g-index

29 all docs 29 docs citations

times ranked

29

2539 citing authors

#	Article	IF	CITATIONS
1	RNA stable isotope probing and highâ€throughput sequencing to identify active microbial community members in a methaneâ€driven denitrifying biofilm. Journal of Applied Microbiology, 2022, 132, 1526-1542.	1.4	4
2	Growth on Formic Acid Is Dependent on Intracellular pH Homeostasis for the Thermoacidophilic Methanotroph Methylacidiphilum sp. RTK17.1. Frontiers in Microbiology, 2021, 12, 651744.	1.5	12
3	Draft Genome Sequence of Limisphaera ngatamarikiensis NGM72.4 ^T , a Moderately Alkaliphilic Thermophile Belonging to the Class <i>Verrucomicrobiae</i> Announcements, 2020, 9, .	0.3	2
4	Thermophilic methanotrophs: in hot pursuit. FEMS Microbiology Ecology, 2019, 95, .	1.3	18
5	Hydrogen Oxidation Influences Glycogen Accumulation in a Verrucomicrobial Methanotroph. Frontiers in Microbiology, 2019, 10, 1873.	1.5	15
6	Two Chloroflexi classes independently evolved the ability to persist on atmospheric hydrogen and carbon monoxide. ISME Journal, 2019, 13, 1801-1813.	4.4	129
7	Mixed culture polyhydroxyalkanoate (PHA) synthesis from nutrient rich wet oxidation liquors. Water Research, 2018, 140, 1-11.	5.3	47
8	Microbial biogeography of 925 geothermal springs in New Zealand. Nature Communications, 2018, 9, 2876.	5.8	163
9	Interaction between ferruginous clay sediment and an iron-reducing hyperthermophilic Pyrobaculum sp. in a terrestrial hot spring. FEMS Microbiology Ecology, 2018, 94, .	1.3	2
10	Mixotrophy drives niche expansion of verrucomicrobial methanotrophs. ISME Journal, 2017, 11, 2599-2610.	4.4	107
11	Atmospheric trace gases support primary production in Antarctic desert surface soil. Nature, 2017, 552, 400-403.	13.7	290
12	The methanogenic redox cofactor F420 is widely synthesized by aerobic soil bacteria. ISME Journal, 2017, 11, 125-137.	4.4	66
13	Cofactor Tail Length Modulates Catalysis of Bacterial F420-Dependent Oxidoreductases. Frontiers in Microbiology, 2017, 8, 1902.	1.5	15
14	Genomic and metagenomic surveys of hydrogenase distribution indicate H2 is a widely utilised energy source for microbial growth and survival. ISME Journal, 2016, 10, 761-777.	4.4	503
15	Persistence of the dominant soil phylum <i>Acidobacteria</i> by trace gas scavenging. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 10497-10502.	3.3	117
16	Role of transcription and enzyme activities in redistribution of carbon and electron flux in response to N2 and H2 sparging of open-batch cultures of Clostridium thermocellum ATCC 27405. Applied Microbiology and Biotechnology, 2014, 98, 2829-2840.	1.7	16
17	Linking genome content to biofuel production yields: a meta-analysis of major catabolic pathways among select H2and ethanol-producing bacteria. BMC Microbiology, 2012, 12, 295.	1.3	58
18	Challenges for biohydrogen production via direct lignocellulose fermentation. International Journal of Hydrogen Energy, 2009, 34, 7390-7403.	3.8	85

#	Article	IF	CITATIONS
19	Pyruvate catabolism and hydrogen synthesis pathway genes of Clostridium thermocellum ATCC 27405. Indian Journal of Microbiology, 2008, 48, 252-266.	1.5	40
20	Third Generation Biofuels via Direct Cellulose Fermentation. International Journal of Molecular Sciences, 2008, 9, 1342-1360.	1.8	252
21	Formate synthesis byClostridium thermocellumduring anaerobic fermentation. Canadian Journal of Microbiology, 2006, 52, 681-688.	0.8	47