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List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/12058893/publications.pdf

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8	201	1162889 8	1588896 8
papers	citations	h-index	g-index

8	8	8	255
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Biogeochemical N signatures from rate-yield trade-offs during in vitro chemosynthetic NO3â˜ reduction by deep-sea vent Îµ-Proteobacteria and Aquificae growing at different temperatures. Geochimica Et Cosmochimica Acta, 2017, 211, 214-227.	1.6	10
2	From deep-sea volcanoes to human pathogens: a conserved quorum-sensing signal in Epsilonproteobacteria. ISME Journal, 2015, 9, 1222-1234.	4.4	55
3	Detection and phylogenetic analysis of the membrane-bound nitrate reductase (Nar) in pure cultures and microbial communities from deep-sea hydrothermal vents. FEMS Microbiology Ecology, 2013, 86, 256-267.	1.3	17
4	Complete genome sequence of Thermovibrio ammonificans HB-1T, a thermophilic, chemolithoautotrophic bacterium isolated from a deep-sea hydrothermal vent. Standards in Genomic Sciences, 2012, 7, 82-90.	1.5	11
5	Phorcysia thermohydrogeniphila gen. nov., sp. nov., a thermophilic, chemolithoautotrophic, nitrate-ammonifying bacterium from a deep-sea hydrothermal vent. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 2388-2394.	0.8	20
6	Draft genome sequence of Caminibacter mediatlanticus strain TB-2T, an epsilonproteobacterium isolated from a deep-sea hydrothermal vent. Standards in Genomic Sciences, 2011, 5, 135-143.	1.5	17
7	Nautilia nitratireducens sp. nov., a thermophilic, anaerobic, chemosynthetic, nitrate-ammonifying bacterium isolated from a deep-sea hydrothermal vent. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 1182-1186.	0.8	33
8	Salinisphaera hydrothermalis sp. nov., a mesophilic, halotolerant, facultatively autotrophic, thiosulfate-oxidizing gammaproteobacterium from deep-sea hydrothermal vents, and emended description of the genus Salinisphaera. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 1497-1503.	0.8	38