

Deciphering the evolution and metabolism of an anammox genome

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Citation Report

#	ARTICLE	IF	CITATIONS
2	Microfluidic Digital PCR Enables Multigene Analysis of Individual Environmental Bacteria. <i>Science</i> , 2006, 314, 1464-1467.	6.0	694
3	Population genomics in natural microbial communities. <i>Trends in Ecology and Evolution</i> , 2006, 21, 508-516.	4.2	94
5	The molecular ecologist's guide to expressed sequence tags. <i>Molecular Ecology</i> , 2006, 16, 907-924.	2.0	326
8	The Planctomycetes, Verrucomicrobia, Chlamydiae and sister phyla comprise a superphylum with biotechnological and medical relevance. <i>Current Opinion in Biotechnology</i> , 2006, 17, 241-249.	3.3	405
9	A new approach to non-destructive analysis of biofilms by confocal Raman microscopy. <i>Analytical and Bioanalytical Chemistry</i> , 2006, 386, 286-292.	1.9	64
10	Environmental Genomics: Exploring Ecological Sequence Space. <i>Current Biology</i> , 2006, 16, R499-R501.	1.8	9
11	Wastewater treatment: a model system for microbial ecology. <i>Trends in Biotechnology</i> , 2006, 24, 483-489.	4.9	216
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13	Whole-Genome Reciprocal BLAST Analysis Reveals that Planctomycetes Do Not Share an Unusually Large Number of Genes with Eukarya and Archaea. <i>Applied and Environmental Microbiology</i> , 2006, 72, 6841-6844.	1.4	33
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22	Effects of Specific Inhibitors on Anammox and Denitrification in Marine Sediments. <i>Applied and Environmental Microbiology</i> , 2007, 73, 3151-3158.	1.4	113

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24	Have we overemphasized the role of denitrification in aquatic ecosystems? A review of nitrate removal pathways. <i>Frontiers in Ecology and the Environment</i> , 2007, 5, 89-96.	1.9	906
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