

HGT-DB: a database of putative horizontally transferred genomes

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Speciation in Chlamydia : Genomewide Phylogenetic Analyses Identified a Reliable Set of Acquired Genes. <i>Journal of Molecular Evolution</i> , 2003, 57, 672-680.	1.8	20
2	The tripartite tricarboxylate transporter (TTT) family. <i>Research in Microbiology</i> , 2003, 154, 457-465.	2.1	85
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4	Analysis of DNA Sequences through Segmentation: Exploring the Mosaic via Statistical Measures. <i>Physica Scripta</i> , 2003, T106, 21.	2.5	0
5	Successful Lateral Transfer Requires Codon Usage Compatibility Between Foreign Genes and Recipient Genomes. <i>Molecular Biology and Evolution</i> , 2004, 21, 1884-1894.	8.9	75
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17	Exploration of phylogenetic data using a global sequence analysis method. <i>BMC Evolutionary Biology</i> , 2005, 5, 63.	3.2	38
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