

A complete bacterial genome assembled de novo using c

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Rapid metagenomic identification of viral pathogens in clinical samples by real-time nanopore sequencing analysis. <i>Genome Medicine</i> , 2015, 7, 99.	3.6	456
2	Scaffolding of a bacterial genome using MinION nanopore sequencing. <i>Scientific Reports</i> , 2015, 5, 11996.	1.6	70
3	MinIONâ„¢: New, Long Read, Portable Nucleic Acid Sequencing Device. <i>Journal of Bacteriology and Virology</i> , 2015, 45, 285.	0.0	6
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5	Transposable element detection from whole genome sequence data. <i>Mobile DNA</i> , 2015, 6, 24.	1.3	139
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9	Gating-like Motions and Wall Porosity in a DNA Nanopore Scaffold Revealed by Molecular Simulations. <i>ACS Nano</i> , 2015, 9, 11209-11217.	7.3	51
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