

Whole genome comparison of a large collection of myco continuum of phage genetic diversity

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Global insights into acetic acid resistance mechanisms and genetic stability of <i>Acetobacter pasteurianus</i> strains by comparative genomics. <i>Scientific Reports</i> , 2015, 5, 18330.	3.3	47
2	An Unusual Phage Repressor Encoded by Mycobacteriophage BPs. <i>PLoS ONE</i> , 2015, 10, e0137187.	2.5	12
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4	Draft Genome Sequence of <i>Mycobacterium heraklionense</i> Strain Davo. <i>Genome Announcements</i> , 2015, 3, .	0.8	4
5	Adding pieces to the puzzle: New insights into bacteriophage diversity from integrated research-education programs. <i>Bacteriophage</i> , 2015, 5, e1084073.	1.9	12
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9	A century of the phage: past, present and future. <i>Nature Reviews Microbiology</i> , 2015, 13, 777-786.	28.6	537
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11	Genome Sequences of <i>Gordonia terrae</i> Phages Benczkowski14 and Katyusha. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
12	Genome Sequences of <i>Gordonia</i> Phages BaxterFox, Kita, Nymphadora, and Yeezy. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
13	Genome Sequence of <i>Gordonia</i> Phage Emalyn. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
14	Genome Sequences of <i>Gordonia</i> Phages Hotorobo, Woes, and Monty. <i>Genome Announcements</i> , 2016, 4, .	0.8	1
15	Genome Sequence of Mycobacteriophage ErnieJ. <i>Genome Announcements</i> , 2016, 4, .	0.8	0
16	The Global Reciprocal Reprogramming between Mycobacteriophage SWU1 and <i>Mycobacterium</i> Reveals the Molecular Strategy of Subversion and Promotion of Phage Infection. <i>Frontiers in Microbiology</i> , 2016, 7, 41.	3.5	8
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