

Host–pathogen coevolution in human tuberculosis

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Bovine TB and badger culling. <i>Veterinary Record</i> , 2012, 171, 353-354.	0.2	0
2	The Importance of Understanding the Human-Animal Interface. <i>Current Topics in Microbiology and Immunology</i> , 2012, 365, 49-81.	0.7	25
3	Importance of the Genetic Diversity within the Mycobacterium tuberculosis Complex for the Development of Novel Antibiotics and Diagnostic Tests of Drug Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 6080-6087.	1.4	55
4	Comprehensive analysis of methods used for the evaluation of compounds against Mycobacterium tuberculosis. <i>Tuberculosis</i> , 2012, 92, 453-488.	0.8	193
5	Diverse approaches to analysing the history of human and pathogen evolution: how to tell the story of the past 70 000 years. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2012, 367, 765-769.	1.8	3
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7	Co-infection with <i>Mycobacterium tuberculosis</i> and human immunodeficiency virus: an overview and motivation for systems approaches. <i>Pathogens and Disease</i> , 2013, 69, 101-113.	0.8	23
8	Long-term dominance of Mycobacterium tuberculosis Uganda family in peri-urban Kampala-Uganda is not associated with cavitory disease. <i>BMC Infectious Diseases</i> , 2013, 13, 484.	1.3	36
9	Single nucleotide polymorphisms in Mycobacterium tuberculosis and the need for a curated database. <i>Tuberculosis</i> , 2013, 93, 30-39.	0.8	43
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15	Phylogenetic lineages of tuberculosis isolates in New Zealand and their association with patient demographics. <i>International Journal of Tuberculosis and Lung Disease</i> , 2013, 17, 892-897.	0.6	21
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18	Mycobacterium tuberculosis: Evolution, Host-Pathogen Interactions, and Implications for Tuberculosis Control. , 2013, , 111-146.		0

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116	Mathematical Models for the Epidemiology and Evolution of <i>Mycobacterium tuberculosis</i> . <i>Advances in Experimental Medicine and Biology</i> , 2017, 1019, 281-307.	0.8	1
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