

Mapping of a Novel Susceptibility Locus Suggests a Role Tuberculosis

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Genetic studies of African populations: an overview on disease susceptibility and response to vaccines and therapeutics. <i>Human Genetics</i> , 2008, 123, 557-598.	3.8	79
2	South Africa: from species cradle to genomic applications. <i>Nature Reviews Genetics</i> , 2008, 9, S19-S23.	16.3	39
3	Identification of Tuberculosis Susceptibility Genes with Human Macrophage Gene Expression Profiles. <i>PLoS Pathogens</i> , 2008, 4, e1000229.	4.7	134
4	Genome Scan of <i>M. tuberculosis</i> Infection and Disease in Ugandans. <i>PLoS ONE</i> , 2008, 3, e4094.	2.5	113
5	Polymorphic Exact Tandem Repeat A (PETRA): a Newly Defined Lineage of <i>Mycobacterium tuberculosis</i> in Israel Originating Predominantly in Sub-Saharan Africa. <i>Journal of Clinical Microbiology</i> , 2009, 47, 4006-4020.	3.9	11
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18	Genetic susceptibility to tuberculosis associated with cathepsin Z haplotype in a Ugandan household contact study. <i>Human Immunology</i> , 2011, 72, 426-430.	2.4	22
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20	A Preliminary Study of Genetic Factors That Influence Susceptibility to Bovine Tuberculosis in the British Cattle Herd. <i>PLoS ONE</i> , 2011, 6, e18806.	2.5	28
21	Polymorphisms in MC3R promoter and CTSZ 3'UTR are associated with tuberculosis susceptibility. <i>European Journal of Human Genetics</i> , 2011, 19, 676-681.	2.8	38

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23	Knowledge gaining by human genetic studies on tuberculosis susceptibility. <i>Journal of Human Genetics</i> , 2011, 56, 177-182.	2.3	33
24	Genetic Epidemiology of Tuberculosis Susceptibility: Impact of Study Design. <i>PLoS Pathogens</i> , 2011, 7, e1001189.	4.7	77
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33	Evolution, revolution and heresy in the genetics of infectious disease susceptibility. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2012, 367, 840-849.	4.0	117
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37	Host-Pathogen Specificity in Tuberculosis. <i>Advances in Experimental Medicine and Biology</i> , 2013, 783, 33-44.	1.6	24
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59	Human genetics of mycobacterial disease. <i>Mammalian Genome</i> , 2018, 29, 523-538.	2.2	27
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63	rs6127698 polymorphism in the MC3R gene and susceptibility to multifocal tuberculosis in southern Chinese Han population. <i>Infection, Genetics and Evolution</i> , 2020, 82, 104292.	2.3	4
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