

CITATION REPORT

List of articles citing

Discovery of susceptibility loci associated with tuberculosis in Han Chinese

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Human Molecular Genetics, 2017, 26, 4752-4763.

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| # | Paper | IF | Citations |
|----|--|------|-----------|
| 46 | Highlights on the Application of Genomics and Bioinformatics in the Fight Against Infectious Diseases: Challenges and Opportunities in Africa. <i>Frontiers in Genetics</i> , 2018 , 9, 575 | 4.5 | 12 |
| 45 | Genetic Resistance to Infection and Disease. <i>Frontiers in Immunology</i> , 2018 , 9, 2219 | 8.4 | 17 |
| 44 | Genetic variants in IFNG and IFNGR1 and tuberculosis susceptibility. <i>Cytokine</i> , 2019 , 123, 154775 | 4 | 6 |
| 43 | Homozygosity for P1104A underlies tuberculosis in about 1% of patients in a cohort of European ancestry. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019 , 116, 10430-10434 | 11.5 | 46 |
| 42 | An exome wide association study of pulmonary tuberculosis patients and their asymptomatic household contacts. <i>Infection, Genetics and Evolution</i> , 2019 , 71, 76-81 | 4.5 | 9 |
| 41 | A Sex-Stratified Genome-Wide Association Study of Tuberculosis Using a Multi-Ethnic Genotyping Array. <i>Frontiers in Genetics</i> , 2018 , 9, 678 | 4.5 | 14 |
| 40 | Tuberculosis infection and lung adenocarcinoma: Mendelian randomization and pathway analysis of genome-wide association study data from never-smoking Asian women. <i>Genomics</i> , 2020 , 112, 1223-1232 | 4.3 | 8 |
| 39 | Interleukin 1 α and 1 β gene variations are associated with tuberculosis in silica exposed subjects. <i>American Journal of Industrial Medicine</i> , 2020 , 63, 74-84 | 2.7 | 3 |
| 38 | Mycobacterium tuberculosis infection up-regulates MFN2 expression to promote NLRP3 inflammasome formation. <i>Journal of Biological Chemistry</i> , 2020 , 295, 17684-17697 | 5.4 | 11 |
| 37 | Mitofusin 2 in Macrophages Links Mitochondrial ROS Production, Cytokine Release, Phagocytosis, Autophagy, and Bactericidal Activity. <i>Cell Reports</i> , 2020 , 32, 108079 | 10.6 | 33 |
| 36 | Functional nucleotide polymorphisms up-regulating transforming growth factor β expression are associated with increased tuberculosis susceptibility. <i>Journal of Infectious Diseases</i> , 2020 , | 7 | 2 |
| 35 | RGS12 Is a Novel Critical NF- κ B Activator in Inflammatory Arthritis. <i>iScience</i> , 2020 , 23, 101172 | 6.1 | 14 |
| 34 | The dominant model analysis of Sirt3 genetic variants is associated with susceptibility to tuberculosis in a Chinese Han population. <i>Molecular Genetics and Genomics</i> , 2020 , 295, 1155-1162 | 3.1 | 1 |
| 33 | Genome-wide association study of Buruli ulcer in rural Benin highlights role of two LncRNAs and the autophagy pathway. <i>Communications Biology</i> , 2020 , 3, 177 | 6.7 | 13 |
| 32 | Evaluation of the Host Genetic Effects of Tuberculosis-Associated Variants Among Patients With Type 1 and Type 2 Diabetes Mellitus. <i>Open Forum Infectious Diseases</i> , 2020 , 7, ofaa106 | 1 | 0 |
| 31 | Host genetics and infectious disease: new tools, insights and translational opportunities. <i>Nature Reviews Genetics</i> , 2021 , 22, 137-153 | 30.1 | 27 |
| 30 | Mitochondria: Powering the Innate Immune Response to Mycobacterium tuberculosis Infection. <i>Infection and Immunity</i> , 2021 , 89, | 3.7 | 1 |

| | | | |
|----|---|------|----|
| 29 | Tuberculosis Exposure With Risk of Behçet Disease Among Patients With Uveitis. <i>JAMA Ophthalmology</i> , 2021 , 139, 415-422 | 3.9 | 0 |
| 28 | Sequencing of over 100,000 individuals identifies multiple genes and rare variants associated with Crohns disease susceptibility. | | 2 |
| 27 | Mitochondrial Dynamics Related Genes - and Polymorphisms are Associated with Risk of Lung Cancer. <i>Pharmacogenomics and Personalized Medicine</i> , 2021 , 14, 695-703 | 2.1 | |
| 26 | A next generation sequencing combined genome-wide association study identifies novel tuberculosis susceptibility loci in Chinese population. <i>Genomics</i> , 2021 , 113, 2377-2384 | 4.3 | 1 |
| 25 | A cross-population atlas of genetic associations for 220 human phenotypes. <i>Nature Genetics</i> , 2021 , 53, 1415-1424 | 36.3 | 40 |
| 24 | A global atlas of genetic associations of 220 deep phenotypes. | | 14 |
| 23 | A sex-stratified genome-wide association study of tuberculosis using a multi-ethnic genotyping array. | | 2 |
| 22 | Local Ancestry Adjusted Allelic Association Analysis Robustly Captures Tuberculosis Susceptibility Loci. <i>Frontiers in Genetics</i> , 2021 , 12, 716558 | 4.5 | 3 |
| 21 | Association between Tuberculosis Case and CD44 Gene Polymorphism. <i>Korean Journal of Clinical Laboratory Science</i> , 2019 , 51, 323-328 | 0.4 | 1 |
| 20 | Genome-wide association study of Buruli ulcer in rural Benin. | | |
| 19 | Replicated Association Study between Tuberculosis and CLCN6, DOK7, HLA-DRA in Korean. <i>Biomedical Science Letters</i> , 2020 , 26, 238-243 | 0.3 | 0 |
| 18 | Integrative genomics analysis identifies promising SNPs and genes implicated in tuberculosis risk based on multiple omics datasets. <i>Aging</i> , 2020 , 12, 19173-19220 | 5.6 | 2 |
| 17 | Integrative genomics analysis identifies promising SNPs and genes implicated in tuberculosis risk based on multiple omics datasets. <i>Aging</i> , 2020 , 12, 19173-19220 | 5.6 | 2 |
| 16 | Large Registry Based Analysis of Genetic Predisposition to Tuberculosis Identifies Genetic Risk Factors at HLA. | | |
| 15 | Association between human leukocyte antigen class II (HLA-DRB and -DQB) alleles and outcome of exposure to : a cross-sectional study in Nairobi, Kenya.. <i>Pan African Medical Journal</i> , 2022 , 41, 149 | 1.2 | |
| 14 | Genetic Analysis of TB Susceptibility Variants in Ghana Reveals Candidate Protective Loci in and Genes.. <i>Frontiers in Genetics</i> , 2021 , 12, 729737 | 4.5 | 0 |
| 13 | Deciphering Genetic Susceptibility to Tuberculous Meningitis.. <i>Frontiers in Neurology</i> , 2022 , 13, 820168 | 4.1 | |
| 12 | Data_Sheet_1.docx. 2019 , | | |

11 Data_Sheet_2.xlsx. **2019**,

| | | | |
|----|--|------|---|
| 10 | Human Leukocyte Antigen (HLA) System: Genetics and Association with Bacterial and Viral Infections. <i>Journal of Immunology Research</i> , 2022 , 2022, 1-15 | 4.5 | 4 |
| 9 | The Awesome Power of Human Genetics of Infectious Disease. <i>Annual Review of Genetics</i> , 2022 , 56, | 14.5 | 0 |
| 8 | Large registry based analysis of genetic predisposition to tuberculosis identifies genetic risk factors at HLA. | | |
| 7 | Genetic architecture of tuberculosis susceptibility: A comprehensive research synopsis, meta-analyses, and epidemiological evidence. 2022 , 104, 105352 | | 1 |
| 6 | Multi-ancestry meta-analysis of host genetic susceptibility to tuberculosis identifies shared genetic architecture. | | 0 |
| 5 | Decoding the spatial chromatin organization and dynamic epigenetic landscapes of macrophage cells during differentiation and immune activation. 2022 , 13, | | 1 |
| 4 | The immunogenetics of tuberculosis (TB) susceptibility. | | 0 |
| 3 | Genome-wide association study of tuberculosis in the western Chinese Han and Tibetan population. 2023 , 4, | | 0 |
| 2 | Fulminant pulmonary tuberculosis in a previously healthy young woman from the Marshall Islands: Potential risk factors. 2023 , 31, 100351 | | 0 |
| 1 | Altered IL-6 signalling and risk of tuberculosis disease: a meta-analysis and Mendelian randomisation study. | | 0 |