Liping Pan

List of Publications by Year
in descending order

20 papers	372 citations	840776 11 h-index	839539 18 g-index
22 all docs	22 docs citations	22 times ranked	575 citing authors

#	Article	IF	Citations
1	Rapid Detection of Mycobacterium tuberculosis in Pleural Fluid Using Resuscitation-Promoting Factor-Based Thin Layer Agar Culture Method. Frontiers in Microbiology, 2022, 13, 803521.	3.5	0
2	Diagnostic Performance of a Novel CXCL10 mRNA Release Assay for Mycobacterium tuberculosis Infection. Frontiers in Microbiology, 2022, 13, 825413.	3.5	2
3	Analysis of drug resistance and mutation profiles in <i>Mycobacterium tuberculosis</i> isolates in a surveillance site in Beijing, China. Journal of International Medical Research, 2021, 49, 030006052098493.	1.0	3
4	Hsp16.3 of mycobacterium tuberculosis in exosomes as a biomarker of tuberculosis. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 2427-2430.	2.9	10
5	Use of T-SPOT.TB for the diagnosis of unconventional pleural tuberculosis is superior to ADA in high prevalence areas: aÂprospective analysis of 601 cases. BMC Infectious Diseases, 2021, 21, 4.	2.9	16
6	Labelâ€Free Quantitative Proteomics Identifies Novel Biomarkers for Distinguishing Tuberculosis Pleural Effusion from Malignant Pleural Effusion. Proteomics - Clinical Applications, 2020, 14, 1900001.	1.6	4
7	Evaluation of digital PCR assay in detection of M.tuberculosis IS6110 and IS1081 in tuberculosis patients plasma. BMC Infectious Diseases, 2020, 20, 657.	2.9	14
8	Application of the CRISPRi system to repress sepF expression in Mycobacterium smegmatis. Infection, Genetics and Evolution, 2019, 72, 183-190.	2.3	13
9	Histone deacetylase inhibitors impair the host immune response against Mycobacterium tuberculosis infection. Tuberculosis, 2019, 118, 101861.	1.9	13
10	Genome-Wide miRNA Analysis Identifies Potential Biomarkers in Distinguishing Tuberculous and Viral Meningitis. Frontiers in Cellular and Infection Microbiology, 2019, 9, 323.	3.9	21
11	Small RNA Profiles of Serum Exosomes Derived From Individuals With Latent and Active Tuberculosis. Frontiers in Microbiology, 2019, 10, 1174.	3.5	62
12	A Two-Way Proteome Microarray Strategy to Identify Novel Mycobacterium tuberculosis-Human Interactors. Frontiers in Cellular and Infection Microbiology, 2019, 9, 65.	3.9	4
13	Proteomic profiling for plasma biomarkers of tuberculosis progression. Molecular Medicine Reports, 2018, 18, 1551-1559.	2.4	16
14	Label-Free Quantitative Proteomics Identifies Novel Plasma Biomarkers for Distinguishing Pulmonary Tuberculosis and Latent Infection. Frontiers in Microbiology, 2018, 9, 1267.	3.5	31
15	Interferon-Gamma Release Assay Performance of Cerebrospinal Fluid and Peripheral Blood in Tuberculous Meningitis in China. BioMed Research International, 2017, 2017, 1-10.	1.9	16
16	Genome-wide transcriptional profiling identifies potential signatures in discriminating active tuberculosis from latent infection. Oncotarget, 2017, 8, 112907-112916.	1.8	19
17	Diagnostic performance of interferon-Î³ release assay for lymph node tuberculosis. Diagnostic Microbiology and Infectious Disease, 2016, 85, 56-60.	1.8	13
18	Risk factors for false-negative T-SPOT.TB assay results in patients with pulmonary and extra-pulmonary TB. Journal of Infection, 2015, 70, 367-380.	3.3	56

#	Article	IF	Citations
19	A proteomics approach to the identification of plasma biomarkers for latent tuberculosis infection. Diagnostic Microbiology and Infectious Disease, 2014, 79, 432-437.	1.8	12
20	Prevalence and Risk Factors for Latent Tuberculosis Infection among Health Care Workers in China: A Cross-Sectional Study. PLoS ONE, 2013, 8, e66412.	2.5	46