

Bouke Catherine de Jong

List of Publications by Year in descending order

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147
papers

7,498
citations

87723

38
h-index

64668

79
g-index

160
all docs

160
docs citations

160
times ranked

6952
citing authors

#	ARTICLE	IF	CITATIONS
1	Definitive outcomes in patients with rifampicin-resistant tuberculosis treated in Niger from 2012 to 2019: A retrospective cohort study. <i>International Health</i> , 2023, 15, 258-264.	0.8	1
2	Multidrug-resistant tuberculosis control in Rwanda overcomes a successful clone that causes most disease over a quarter century. <i>Journal of Clinical Tuberculosis and Other Mycobacterial Diseases</i> , 2022, 27, 100299.	0.6	2
3	High yield of retrospective active case finding for leprosy in Comoros. <i>PLoS Neglected Tropical Diseases</i> , 2022, 16, e0010158.	1.3	3
4	Acquired rifampicin resistance during first TB treatment: magnitude, relative importance, risk factors and keys to control in low-income settings. <i>JAC-Antimicrobial Resistance</i> , 2022, 4, dlac037.	0.9	0
5	Investigating drug resistance of <i>Mycobacterium leprae</i> in the Comoros: an observational deep-sequencing study. <i>Lancet Microbe</i> , The, 2022, 3, e693-e700.	3.4	9
6	Second-line injectable drugs for rifampicin-resistant tuberculosis: better the devil we know?. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 831-835.	1.3	9
7	Genetic diversity of the <i>Mycobacterium tuberculosis</i> complex strains from newly diagnosed tuberculosis patients in Northwest Ethiopia reveals a predominance of East-African-Indian and Euro-American lineages. <i>International Journal of Infectious Diseases</i> , 2021, 103, 72-80.	1.5	7
8	High rifampicin-resistant TB cure rates and prevention of severe ototoxicity after replacing the injectable by linezolid in early stage of hearing loss. <i>European Respiratory Journal</i> , 2021, 57, 2002250.	3.1	12
9	Deep amplicon sequencing for culture-free prediction of susceptibility or resistance to 13 anti-tuberculous drugs. <i>European Respiratory Journal</i> , 2021, 57, 2002338.	3.1	58
10	Low Cycle Threshold Value in Xpert MTB/RIF Assay May Herald False Detection of Tuberculosis and Rifampicin Resistance: A Study of Two Cases. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofab034.	0.4	2
11	Phylogenomics of <i>Mycobacterium africanum</i> reveals a new lineage and a complex evolutionary history. <i>Microbial Genomics</i> , 2021, 7, .	1.0	71
12	<i>Mycobacterium tuberculosis</i> <i>rpoB</i> mutations: emerging from the unknown. <i>European Respiratory Journal</i> , 2021, 58, 2100783.	3.1	9
13	<i>Mycobacterium tuberculosis</i> precursor rRNA as a measure of treatment-shortening activity of drugs and regimens. <i>Nature Communications</i> , 2021, 12, 2899.	5.8	38
14	<i>Mycobacterium tuberculosis</i> complex lineage 5 exhibits high levels of within-lineage genomic diversity and differing gene content compared to the type strain H37Rv. <i>Microbial Genomics</i> , 2021, 7, .	1.0	9
15	Exploring clustering of leprosy in the Comoros and Madagascar: A geospatial analysis. <i>International Journal of Infectious Diseases</i> , 2021, 108, 96-101.	1.5	7
16	Bedaquiline can act as core drug in a standardized treatment regimen for fluoroquinolone-resistant rifampicin-resistant tuberculosis. <i>European Respiratory Journal</i> , 2021, , 2102124.	3.1	2
17	Quantifying transmission fitness costs of multi-drug resistant tuberculosis. <i>Epidemics</i> , 2021, 36, 100471.	1.5	13
18	Minimally invasive sampling to identify leprosy patients with a high bacterial burden in the Union of the Comoros. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009924.	1.3	3

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19	World Health Organization 2018 treatment guidelines for rifampicin-resistant tuberculosis: uncertainty, potential risks and the way forward. <i>International Journal of Antimicrobial Agents</i> , 2020, 55, 105822.	1.1	19
20	Management of multidrug-resistant tuberculosis with shorter treatment regimen in Niger: Nationwide programmatic achievements. <i>Respiratory Medicine</i> , 2020, 161, 105844.	1.3	26
21	MDR M. tuberculosis outbreak clone in Eswatini missed by Xpert has elevated bedaquiline resistance dated to the pre-treatment era. <i>Genome Medicine</i> , 2020, 12, 104.	3.6	50
22	Characterization of <i>Mycobacterium tuberculosis</i> var. <i>africanum</i> isolated from a patient with pulmonary tuberculosis in Brazil. <i>Infection, Genetics and Evolution</i> , 2020, 85, 104550.	1.0	6
23	Injectables™ key role in rifampicin-resistant tuberculosis shorter treatment regimen outcomes. <i>PLoS ONE</i> , 2020, 15, e0238016.	1.1	6
24	Introduction of <i>Mycobacterium ulcerans</i> disease in the Bankim Health District of Cameroon follows damming of the Mapo River. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008501.	1.3	5
25	Initial resistance to companion drugs should not be considered an exclusion criterion for the shorter multidrug-resistant tuberculosis treatment regimen. <i>International Journal of Infectious Diseases</i> , 2020, 100, 357-365.	1.5	12
26	Management of falsepositive rifampicin resistant Xpert MTB/RIF – Authors' reply. <i>Lancet Microbe</i> , The, 2020, 1, e239.	3.4	0
27	The perceived impact of isoniazid resistance on outcome of first-line rifampicin-throughout regimens is largely due to missed rifampicin resistance. <i>PLoS ONE</i> , 2020, 15, e0233500.	1.1	16
28	Prevalence and drivers of false-positive rifampicin-resistant Xpert MTB/RIF results: a prospective observational study in Rwanda. <i>Lancet Microbe</i> , The, 2020, 1, e74-e83.	3.4	35
29	A sister lineage of the <i>Mycobacterium tuberculosis</i> complex discovered in the African Great Lakes region. <i>Nature Communications</i> , 2020, 11, 2917.	5.8	136
30	High-Dose First-Line Treatment Regimen for Recurrent Rifampicin-Susceptible Tuberculosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 201, 1578-1579.	2.5	8
31	False Rifampicin Resistance in Xpert Ultra Applied to Lymph Node Aspirate: A Case Report. <i>Open Forum Infectious Diseases</i> , 2020, 7, ofaa204.	0.4	1
32	Targeted next-generation sequencing of sputum for diagnosis of drug-resistant TB: results of a national survey in Democratic Republic of the Congo. <i>Scientific Reports</i> , 2020, 10, 10786.	1.6	13
33	Tuberculosis treatment: one-shot approach or cascade of regimens?. <i>Lancet Respiratory Medicine</i> , the, 2020, 8, e4-e5.	5.2	7
34	Evolution of <i>Mycobacterium tuberculosis</i> complex lineages and their role in an emerging threat of multidrug resistant tuberculosis in Bamako, Mali. <i>Scientific Reports</i> , 2020, 10, 327.	1.6	23
35	Comparative genomics shows differences in the electron transport and carbon metabolic pathways of <i>Mycobacterium africanum</i> relative to <i>Mycobacterium tuberculosis</i> and suggests an adaptation to low oxygen tension. <i>Tuberculosis</i> , 2020, 120, 101899.	0.8	15
36	Multidrug-resistant patients receiving treatment in Niger who are infected with <i>M. tuberculosis</i> Cameroon family convert faster in smear and culture than those with <i>M. tuberculosis</i> Ghana family. <i>Tuberculosis</i> , 2020, 122, 101922.	0.8	6

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37	Case Report: Dynamics of Acquired Fluoroquinolone Resistance under Standardized Short-Course Treatment of Multidrug-Resistant Tuberculosis. <i>American Journal of Tropical Medicine and Hygiene</i> , 2020, 103, 1443-1446.	0.6	6
38	How Well Do Routine Molecular Diagnostics Detect Rifampin Heteroresistance in <i>Mycobacterium tuberculosis</i> ?. <i>Journal of Clinical Microbiology</i> , 2019, 57, .	1.8	36
39	Variable ability of rapid tests to detect <i>Mycobacterium tuberculosis</i> rpoB mutations conferring phenotypically occult rifampicin resistance. <i>Scientific Reports</i> , 2019, 9, 11826.	1.6	38
40	Clustering of leprosy beyond the household level in a highly endemic setting on the Comoros, an observational study. <i>BMC Infectious Diseases</i> , 2019, 19, 501.	1.3	22
41	Whole genome sequencing of <i>Mycobacterium tuberculosis</i> : current standards and open issues. <i>Nature Reviews Microbiology</i> , 2019, 17, 533-545.	13.6	237
42	Reference set of <i>Mycobacterium tuberculosis</i> clinical strains: A tool for research and product development. <i>PLoS ONE</i> , 2019, 14, e0214088.	1.1	56
43	<i>Mycobacterium ulcerans</i> Population Genomics To Inform on the Spread of Buruli Ulcer across Central Africa. <i>MSphere</i> , 2019, 4, .	1.3	14
44	Systematic screening for tuberculosis among hospital outpatients in Cameroon: The role of screening and testing algorithms to improve case detection. <i>Journal of Clinical Tuberculosis and Other Mycobacterial Diseases</i> , 2019, 15, 100095.	0.6	15
45	Protocol, rationale and design of PEOPLE (Post ExpOsure Prophylaxis for LEprosy in the Comoros and) Tj ETQq1 1 0.784314 rgBT /Overd post-exposure prophylaxis of leprosy contacts. <i>BMC Infectious Diseases</i> , 2019, 19, 1033.	1.3	18
46	Multidrug-resistant tuberculosis outbreak in South Africa – Authors' reply. <i>Lancet Infectious Diseases</i> , The, 2019, 19, 135-136.	4.6	0
47	Bridging the TB data gap: <i>in silico</i> extraction of rifampicin-resistant tuberculosis diagnostic test results from whole genome sequence data. <i>PeerJ</i> , 2019, 7, e7564.	0.9	4
48	Isoniazid resistance levels of <i>Mycobacterium tuberculosis</i> can largely be predicted by high-confidence resistance-conferring mutations. <i>Scientific Reports</i> , 2018, 8, 3246.	1.6	87
49	Comparative Genomics Shows That <i>Mycobacterium ulcerans</i> Migration and Expansion Preceded the Rise of Buruli Ulcer in Southeastern Australia. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	1.4	32
50	Diagnostic Accuracy of Clinical and Microbiological Signs in Patients With Skin Lesions Resembling Buruli Ulcer in an Endemic Region. <i>Clinical Infectious Diseases</i> , 2018, 67, 827-834.	2.9	27
51	<i>Mycobacterium africanum</i> (Lineage 6) shows slower sputum smear conversion on tuberculosis treatment than <i>Mycobacterium tuberculosis</i> (Lineage 4) in Bamako, Mali. <i>PLoS ONE</i> , 2018, 13, e0208603.	1.1	17
52	Effect of efavirenz-based antiretroviral therapy and high-dose rifampicin on the pharmacokinetics of isoniazid and acetyl-isoniazid. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 74, 139-148.	1.3	21
53	Outbreak of multidrug-resistant tuberculosis in South Africa undetected by WHO-endorsed commercial tests: an observational study. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 1350-1359.	4.6	118
54	Outbreak of Tuberculosis and Multidrug-Resistant Tuberculosis, Mbuji-Mayi Central Prison, Democratic Republic of the Congo. <i>Emerging Infectious Diseases</i> , 2018, 24, 2029-2035.	2.0	21

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55	The relationship between transmission time and clustering methods in Mycobacterium tuberculosis epidemiology. <i>EBioMedicine</i> , 2018, 37, 410-416.	2.7	106
56	Comparative genomics of Mycobacterium africanum Lineage 5 and Lineage 6 from Ghana suggests distinct ecological niches. <i>Scientific Reports</i> , 2018, 8, 11269.	1.6	34
57	Xpert Ultra Can Unambiguously Identify Specific Rifampin Resistance-Confering Mutations. <i>Journal of Clinical Microbiology</i> , 2018, 56, .	1.8	18
58	Improving clinical and epidemiological predictors of Buruli ulcer. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006713.	1.3	3
59	Performance of OMNIgene [®] SPUTUM (DNA Genotek) and cetylpyridinium chloride for sputum storage prior to mycobacterial culture. <i>Journal of Medical Microbiology</i> , 2018, 67, 798-805.	0.7	3
60	Low sensitivity of the MPT64 identification test to detect lineage 5 of the Mycobacterium tuberculosis complex. <i>Journal of Medical Microbiology</i> , 2018, 67, 1718-1727.	0.7	13
61	Potential Application of Digitally Linked Tuberculosis Diagnostics for Real-Time Surveillance of Drug-Resistant Tuberculosis Transmission: Validation and Analysis of Test Results. <i>JMIR Medical Informatics</i> , 2018, 6, e12.	1.3	11
62	Unexpected high prevalence of resistance-associated <i>Rv0678</i> variants in MDR-TB patients without documented prior use of clofazimine or bedaquiline. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, dkw502.	1.3	134
63	Some Synonymous and Nonsynonymous <i>gyrA</i> Mutations in Mycobacterium tuberculosis Lead to Systematic False-Positive Fluoroquinolone Resistance Results with the Hain GenoType MTBDR <i>sls</i> Assays. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	1.4	29
64	Significant under expression of the DosR regulon in M. tuberculosis complex lineage 6 in sputum. <i>Tuberculosis</i> , 2017, 104, 58-64.	0.8	15
65	The predominance of Ethiopian specific Mycobacterium tuberculosis families and minimal contribution of Mycobacterium bovis in tuberculous lymphadenitis patients in Southwest Ethiopia. <i>Infection, Genetics and Evolution</i> , 2017, 55, 251-259.	1.0	28
66	The Biology and Epidemiology of Mycobacterium africanum. <i>Advances in Experimental Medicine and Biology</i> , 2017, 1019, 117-133.	0.8	24
67	Whole-genome sequencing illuminates the evolution and spread of multidrug-resistant tuberculosis in Southwest Nigeria. <i>PLoS ONE</i> , 2017, 12, e0184510.	1.1	27
68	Extensively drug resistant tuberculosis in Mali: a case report. <i>BMC Research Notes</i> , 2017, 10, 561.	0.6	6
69	Multiple introductions and recent spread of the emerging human pathogen <i>Mycobacterium ulcerans</i> across Africa. <i>Genome Biology and Evolution</i> , 2017, 9, evx003.	1.1	32
70	Genotypic characterization directly applied to sputum improves the detection of Mycobacterium africanum West African 1, under-represented in positive cultures. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005900.	1.3	24
71	Bacterial diversity in Buruli ulcer skin lesions: Challenges in the clinical microbiome analysis of a skin disease. <i>PLoS ONE</i> , 2017, 12, e0181994.	1.1	13
72	Half of rifampicin-resistant Mycobacterium tuberculosis complex isolated from tuberculosis patients in Sub-Saharan Africa have concomitant resistance to pyrazinamide. <i>PLoS ONE</i> , 2017, 12, e0187211.	1.1	8

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73	Whole genome sequencing reveals mycobacterial microevolution among concurrent isolates from sputum and blood in HIV infected TB patients. <i>BMC Infectious Diseases</i> , 2016, 16, 371.	1.3	11
74	Do Xpert MTB/RIF Cycle Threshold Values Provide Information about Patient Delays for Tuberculosis Diagnosis?. <i>PLoS ONE</i> , 2016, 11, e0162833.	1.1	13
75	Whole genome sequencing to complement tuberculosis drug resistance surveys in Uganda. <i>Infection, Genetics and Evolution</i> , 2016, 40, 8-16.	1.0	28
76	Correlation of different phenotypic drug susceptibility testing methods for four fluoroquinolones in <i>Mycobacterium tuberculosis</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 1233-1240.	1.3	32
77	Sputum is a surrogate for bronchoalveolar lavage for monitoring <i>Mycobacterium tuberculosis</i> transcriptional profiles in TB patients. <i>Tuberculosis</i> , 2016, 100, 89-94.	0.8	27
78	Adaptation of <i>Mycobacterium tuberculosis</i> to Impaired Host Immunity in HIV-Infected Patients. <i>Journal of Infectious Diseases</i> , 2016, 214, 1205-1211.	1.9	19
79	<i>Mycobacterium tuberculosis</i> lineage 4 comprises globally distributed and geographically restricted sublineages. <i>Nature Genetics</i> , 2016, 48, 1535-1543.	9.4	326
80	Use of RODAC plates to measure containment of <i>Mycobacterium tuberculosis</i> in a Class IIB biosafety cabinet during routine operations. <i>International Journal of Mycobacteriology</i> , 2016, 5, 148-154.	0.3	1
81	Specific <i>gyrA</i> gene mutations predict poor treatment outcome in MDR-TB. <i>Journal of Antimicrobial Chemotherapy</i> , 2016, 71, 314-323.	1.3	86
82	A <i>Mycobacterium tuberculosis</i> Perspective on Tuberculosis in West Africa: Significant Geographical Variation of <i>M. africanum</i> and Other <i>M. tuberculosis</i> Complex Lineages. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004408.	1.3	35
83	Impact of the <i>Mycobacterium africanum</i> West Africa 2 Lineage on TB Diagnostics in West Africa: Decreased Sensitivity of Rapid Identification Tests in The Gambia. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004801.	1.3	20
84	A tuberculosis nationwide prevalence survey in Gambia, 2012. <i>Bulletin of the World Health Organization</i> , 2016, 94, 433-441.	1.5	17
85	Buruli Ulcer in Traveler from Suriname, South America, to the Netherlands. <i>Emerging Infectious Diseases</i> , 2015, 21, 497-499.	2.0	7
86	A Genomic Approach to Resolving Relapse versus Reinfection among Four Cases of Buruli Ulcer. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0004158.	1.3	14
87	High Genotypic Discordance of Concurrent <i>Mycobacterium tuberculosis</i> Isolates from Sputum and Blood of HIV-Infected Individuals. <i>PLoS ONE</i> , 2015, 10, e0132581.	1.1	15
88	GeneXpert MTB/RIF Assay for the Diagnosis of Tuberculous Lymphadenitis on Concentrated Fine Needle Aspirates in High Tuberculosis Burden Settings. <i>PLoS ONE</i> , 2015, 10, e0137471.	1.1	47
89	First insights into circulating <i>Mycobacterium tuberculosis</i> complex lineages and drug resistance in Guinea. <i>Infection, Genetics and Evolution</i> , 2015, 33, 314-319.	1.0	14
90	Transcriptional Adaptation of Drug-tolerant <i>Mycobacterium tuberculosis</i> During Treatment of Human Tuberculosis. <i>Journal of Infectious Diseases</i> , 2015, 212, 990-998.	1.9	82

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91	Whole Genome Comparisons Suggest Random Distribution of <i>Mycobacterium ulcerans</i> Genotypes in a Buruli Ulcer Endemic Region of Ghana. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003681.	1.3	23
92	Bedaquiline susceptibility testing of <i>Mycobacterium tuberculosis</i> in an automated liquid culture system. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 2300-2305.	1.3	41
93	Predominant <i>Mycobacterium tuberculosis</i> Families and High Rates of Recent Transmission among New Cases Are Not Associated with Primary Multidrug Resistance in Lima, Peru. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1854-1863.	1.8	22
94	Disputed <i>rpoB</i> mutations can frequently cause important rifampicin resistance among new tuberculosis patients. <i>International Journal of Tuberculosis and Lung Disease</i> , 2015, 19, 185-190.	0.6	90
95	<i>M. tuberculosis</i> T Cell Epitope Analysis Reveals Paucity of Antigenic Variation and Identifies Rare Variable TB Antigens. <i>Cell Host and Microbe</i> , 2015, 18, 538-548.	5.1	142
96	The thin-layer agar method for direct phenotypic detection of multi- and extensively drug-resistant tuberculosis. <i>International Journal of Tuberculosis and Lung Disease</i> , 2015, 19, 1547-1552.	0.6	9
97	Acquired Resistance of <i>Mycobacterium tuberculosis</i> to Bedaquiline. <i>PLoS ONE</i> , 2014, 9, e102135.	1.1	320
98	Insertion Sequence Element Single Nucleotide Polymorphism Typing Provides Insights into the Population Structure and Evolution of <i>Mycobacterium ulcerans</i> across Africa. <i>Applied and Environmental Microbiology</i> , 2014, 80, 1197-1209.	1.4	18
99	A Four-Month Gatifloxacin-Containing Regimen for Treating Tuberculosis. <i>New England Journal of Medicine</i> , 2014, 371, 1588-1598.	13.9	352
100	Differences in T cell responses between <i>Mycobacterium tuberculosis</i> and <i>Mycobacterium africanum</i> infected patients. <i>European Journal of Immunology</i> , 2014, 44, 1387-1398.	1.6	21
101	Pseudo-Outbreak of Pre-Extensively Drug-Resistant (Pre-XDR) Tuberculosis in Kinshasa: Collateral Damage Caused by False Detection of Fluoroquinolone Resistance by GenoType MTBDRsl. <i>Journal of Clinical Microbiology</i> , 2014, 52, 2876-2880.	1.8	18
102	Decontamination methods for samples preserved in cetylpyridinium chloride and cultured on thin-layer agar. <i>International Journal of Tuberculosis and Lung Disease</i> , 2014, 18, 972-977.	0.6	3
103	An operational study comparing microscopes and staining variations for tuberculosis LED FM. <i>International Journal of Tuberculosis and Lung Disease</i> , 2014, 18, 964-971.	0.6	3
104	Multicenter External Quality Assessment Program for PCR Detection of <i>Mycobacterium ulcerans</i> in Clinical and Environmental Specimens. <i>PLoS ONE</i> , 2014, 9, e89407.	1.1	29
105	Shifts in <i>Mycobacterial</i> Populations and Emerging Drug-Resistance in West and Central Africa. <i>PLoS ONE</i> , 2014, 9, e110393.	1.1	8
106	Rifampin Drug Resistance Tests for Tuberculosis: Challenging the Gold Standard. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2633-2640.	1.8	216
107	Genetic variability of <i>Mycobacterium tuberculosis</i> complex in patients with no known risk factors for MDR-TB in the North-eastern part of Lima, Peru. <i>BMC Infectious Diseases</i> , 2013, 13, 397.	1.3	12
108	Targeting multidrug-resistant tuberculosis (MDR-TB) by therapeutic vaccines. <i>Medical Microbiology and Immunology</i> , 2013, 202, 95-104.	2.6	63

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109	Immunogenic <i>Mycobacterium africanum</i> Strains Associated with Ongoing Transmission in The Gambia. <i>Emerging Infectious Diseases</i> , 2013, 19, 1598-1604.	2.0	12
110	Deciphering the Growth Behaviour of <i>Mycobacterium africanum</i> . <i>PLoS Neglected Tropical Diseases</i> , 2013, 7, e2220.	1.3	36
111	ADVISER-PYRO: Amplicon Identification using SparsE Representation of PYROsequencing signal. <i>Bioinformatics</i> , 2013, 29, 1963-1969.	1.8	8
112	Burden of <i>Mycobacterium ulcerans</i> Disease (Buruli Ulcer) and the Underreporting Ratio in the Territory of Songololo, Democratic Republic of Congo. <i>PLoS Neglected Tropical Diseases</i> , 2013, 7, e2563.	1.3	22
113	Rifampin Resistance Missed in Automated Liquid Culture System for <i>Mycobacterium tuberculosis</i> Isolates with Specific <i>rpoB</i> Mutations. <i>Journal of Clinical Microbiology</i> , 2013, 51, 2641-2645.	1.8	186
114	Skewing of the CD4+ T-Cell Pool Toward Monofunctional Antigen-Specific Responses in Patients With Immune Reconstitution Inflammatory Syndrome in The Gambia. <i>Clinical Infectious Diseases</i> , 2013, 57, 594-603.	2.9	3
115	The First Phylogeographic Population Structure and Analysis of Transmission Dynamics of <i>M. africanum</i> West African "Combining Molecular Data from Benin, Nigeria and Sierra Leone. <i>PLoS ONE</i> , 2013, 8, e77000.	1.1	26
116	The Genome of <i>Mycobacterium Africanum</i> West African 2 Reveals a Lineage-Specific Locus and Genome Erosion Common to the <i>M. tuberculosis</i> Complex. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1552.	1.3	69
117	Impaired Fitness of <i>Mycobacterium africanum</i> Despite Secretion of ESAT-6. <i>Journal of Infectious Diseases</i> , 2012, 205, 984-990.	1.9	39
118	Fluorescein diacetate vital staining allows earlier diagnosis of rifampicin-resistant tuberculosis. <i>International Journal of Tuberculosis and Lung Disease</i> , 2012, 16, 1174-1179.	0.6	23
119	The TDR Tuberculosis Strain Bank: a resource for basic science, tool development and diagnostic services. <i>International Journal of Tuberculosis and Lung Disease</i> , 2012, 16, 24-31.	0.6	36
120	Immune Reconstitution Inflammatory Syndrome and the Influence of T Regulatory Cells: A Cohort Study in the Gambia. <i>PLoS ONE</i> , 2012, 7, e39213.	1.1	12
121	Factors associated with mortality in patients with drug-susceptible pulmonary tuberculosis. <i>BMC Infectious Diseases</i> , 2011, 11, 1.	1.3	204
122	Effect of a Control Project on Clinical Profiles and Outcomes in Buruli Ulcer: A Before/After Study in Bas-Congo, Democratic Republic of Congo. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1402.	1.3	19
123	Identification of Probable Early-Onset Biomarkers for Tuberculosis Disease Progression. <i>PLoS ONE</i> , 2011, 6, e25230.	1.1	39
124	Immunogenicity of antigens from the TbD1 region present in <i>M. africanum</i> and missing from "modern" <i>M. tuberculosis</i> : a cross-sectional study. <i>BMC Infectious Diseases</i> , 2010, 10, 11.	1.3	2
125	Differences between tuberculosis cases infected with <i>Mycobacterium africanum</i> , West African type 2, relative to Euro-American <i>Mycobacterium tuberculosis</i> : an update. <i>FEMS Immunology and Medical Microbiology</i> , 2010, 58, 102-105.	2.7	61
126	<i>Mycobacterium africanum</i> "Review of an Important Cause of Human Tuberculosis in West Africa. <i>PLoS Neglected Tropical Diseases</i> , 2010, 4, e744.	1.3	221

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127	Infection with <i>Helicobacter pylori</i> Is Associated with Protection against Tuberculosis. <i>PLoS ONE</i> , 2010, 5, e8804.	1.1	133
128	Production of TNF- α , IL-12(p40) and IL-17 Can Discriminate between Active TB Disease and Latent Infection in a West African Cohort. <i>PLoS ONE</i> , 2010, 5, e12365.	1.1	90
129	Use of Spoligotyping and Large Sequence Polymorphisms To Study the Population Structure of the <i>Mycobacterium tuberculosis</i> Complex in a Cohort Study of Consecutive Smear-Positive Tuberculosis Cases in The Gambia. <i>Journal of Clinical Microbiology</i> , 2009, 47, 994-1001.	1.8	53
130	Supervised learning for the automated transcription of spacer classification from spoligotype films. <i>BMC Bioinformatics</i> , 2009, 10, 248.	1.2	4
131	Analysis of flow cytometry data using an automatic processing tool. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2008, 73A, 857-867.	1.1	16
132	Effects of cryopreservation on CD4+ CD25+ T cells of HIV-1 infected individuals. <i>Journal of Clinical Laboratory Analysis</i> , 2008, 22, 153-158.	0.9	22
133	Catch Them While You Can!. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2008, 178, 5-6.	2.5	2
134	Progression to Active Tuberculosis, but Not Transmission, Varies by <i>Mycobacterium tuberculosis</i> Lineage in The Gambia. <i>Journal of Infectious Diseases</i> , 2008, 198, 1037-1043.	1.9	269
135	On Treatment Outcomes of Patients with HIV and Tuberculosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2008, 177, 121-122.	2.5	0
136	Incidence of Tuberculosis and the Predictive Value of ELISPOT and Mantoux Tests in Gambian Case Contacts. <i>PLoS ONE</i> , 2008, 3, e1379.	1.1	116
137	Comparative evaluation of BACTEC MGIT 960 with BACTEC 9000 MB and LJ for isolation of mycobacteria in The Gambia. <i>Journal of Infection in Developing Countries</i> , 2008, 2, 200-5.	0.5	17
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