

Theodore Cohen

List of Publications by Year in descending order

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Version: 2024-02-01

99
papers

3,245
citations

186254

28
h-index

189881

50
g-index

105
all docs

105
docs citations

105
times ranked

4497
citing authors

#	ARTICLE	IF	CITATIONS
1	Estimation of Excess Deaths Associated With the COVID-19 Pandemic in the United States, March to May 2020. <i>JAMA Internal Medicine</i> , 2020, 180, 1336.	5.1	374
2	Origin and Proliferation of Multiple-Drug Resistance in Bacterial Pathogens. <i>Microbiology and Molecular Biology Reviews</i> , 2015, 79, 101-116.	6.6	183
3	The transmission of <i>Mycobacterium tuberculosis</i> in high burden settings. <i>Lancet Infectious Diseases</i> , The, 2016, 16, 227-238.	9.1	149
4	Feasibility of achieving the 2025 WHO global tuberculosis targets in South Africa, China, and India: a combined analysis of 11 mathematical models. <i>The Lancet Global Health</i> , 2016, 4, e806-e815.	6.3	138
5	Genomic diversity in autopsy samples reveals within-host dissemination of HIV-associated <i>Mycobacterium tuberculosis</i> . <i>Nature Medicine</i> , 2016, 22, 1470-1474.	30.7	133
6	Drivers of Tuberculosis Transmission. <i>Journal of Infectious Diseases</i> , 2017, 216, S644-S653.	4.0	123
7	Beyond the SNP Threshold: Identifying Outbreak Clusters Using Inferred Transmissions. <i>Molecular Biology and Evolution</i> , 2019, 36, 587-603.	8.9	121
8	Data for action: collection and use of local data to end tuberculosis. <i>Lancet</i> , The, 2015, 386, 2324-2333.	13.7	89
9	Internal migration and transmission dynamics of tuberculosis in Shanghai, China: an epidemiological, spatial, genomic analysis. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 788-795.	9.1	85
10	Progression from latent infection to active disease in dynamic tuberculosis transmission models: a systematic review of the validity of modelling assumptions. <i>Lancet Infectious Diseases</i> , The, 2018, 18, e228-e238.	9.1	79
11	Rapid Drug Susceptibility Testing of Drug-Resistant <i>Mycobacterium tuberculosis</i> Isolates Directly from Clinical Samples by Use of Amplicon Sequencing: a Proof-of-Concept Study. <i>Journal of Clinical Microbiology</i> , 2016, 54, 2058-2067.	3.9	76
12	Lifetime burden of disease due to incident tuberculosis: a global reappraisal including post-tuberculosis sequelae. <i>The Lancet Global Health</i> , 2021, 9, e1679-e1687.	6.3	74
13	Cost-effectiveness and resource implications of aggressive action on tuberculosis in China, India, and South Africa: a combined analysis of nine models. <i>The Lancet Global Health</i> , 2016, 4, e816-e826.	6.3	69
14	Classic reaction kinetics can explain complex patterns of antibiotic action. <i>Science Translational Medicine</i> , 2015, 7, 287ra73.	12.4	67
15	Identifying Hotspots of Multidrug-Resistant Tuberculosis Transmission Using Spatial and Molecular Genetic Data. <i>Journal of Infectious Diseases</i> , 2016, 213, 287-294.	4.0	62
16	Prospects for Tuberculosis Elimination in the United States: Results of a Transmission Dynamic Model. <i>American Journal of Epidemiology</i> , 2018, 187, 2011-2020.	3.4	58
17	Smear positivity in paediatric and adult tuberculosis: systematic review and meta-analysis. <i>BMC Infectious Diseases</i> , 2016, 16, 282.	2.9	57
18	Evaluating strategies for control of tuberculosis in prisons and prevention of spillover into communities: An observational and modeling study from Brazil. <i>PLoS Medicine</i> , 2019, 16, e1002737.	8.4	55

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19	Incidence and prevalence of tuberculosis in incarcerated populations: a systematic review and meta-analysis. <i>Lancet Public Health</i> , The, 2021, 6, e300-e308.	10.0	54
20	Post-tuberculosis mortality and morbidity: valuing the hidden epidemic. <i>Lancet Respiratory Medicine</i> , the, 2020, 8, 332-333.	10.7	50
21	Within-Host Heterogeneity of <i>Mycobacterium tuberculosis</i> Infection Is Associated With Poor Early Treatment Response: A Prospective Cohort Study. <i>Journal of Infectious Diseases</i> , 2016, 213, 1796-1799.	4.0	45
22	Catastrophic costs potentially averted by tuberculosis control in India and South Africa: a modelling study. <i>The Lancet Global Health</i> , 2017, 5, e1123-e1132.	6.3	41
23	Spatially targeted screening to reduce tuberculosis transmission in high-incidence settings. <i>Lancet Infectious Diseases</i> , The, 2019, 19, e89-e95.	9.1	41
24	Transmissibility and potential for disease progression of drug resistant <i>Mycobacterium tuberculosis</i> : prospective cohort study. <i>BMJ: British Medical Journal</i> , 2019, 367, l5894.	2.3	38
25	The escalating tuberculosis crisis in central and South American prisons. <i>Lancet</i> , The, 2021, 397, 1591-1596.	13.7	38
26	Comparison of Estimated Effectiveness of Case-Based and Population-Based Interventions on COVID-19 Containment in Taiwan. <i>JAMA Internal Medicine</i> , 2021, 181, 913-921.	5.1	37
27	The potential impact of coinfection on antimicrobial chemotherapy and drug resistance. <i>Trends in Microbiology</i> , 2015, 23, 537-544.	7.7	36
28	Tuberculosis control interventions targeted to previously treated people in a high-incidence setting: a modelling study. <i>The Lancet Global Health</i> , 2018, 6, e426-e435.	6.3	34
29	Disparities in access to diagnosis and care in Blantyre, Malawi, identified through enhanced tuberculosis surveillance and spatial analysis. <i>BMC Medicine</i> , 2019, 17, 21.	5.5	34
30	High burden of prevalent tuberculosis among previously treated people in Southern Africa suggests potential for targeted control interventions. <i>European Respiratory Journal</i> , 2016, 48, 1227-1230.	6.7	33
31	Multidrug-resistant tuberculosis treatment failure detection depends on monitoring interval and microbiological method. <i>European Respiratory Journal</i> , 2016, 48, 1160-1170.	6.7	27
32	Yield, Efficiency, and Costs of Mass Screening Algorithms for Tuberculosis in Brazilian Prisons. <i>Clinical Infectious Diseases</i> , 2021, 72, 771-777.	5.8	27
33	Fitness Costs of Drug Resistance Mutations in Multidrug-Resistant <i>Mycobacterium tuberculosis</i> : A Household-Based Case-Control Study. <i>Journal of Infectious Diseases</i> , 2016, 213, 149-155.	4.0	25
34	Genomic variant-identification methods may alter <i>Mycobacterium tuberculosis</i> transmission inferences. <i>Microbial Genomics</i> , 2020, 6, .	2.0	24
35	Global estimates of paediatric tuberculosis incidence in 2013â€“19: a mathematical modelling analysis. <i>The Lancet Global Health</i> , 2022, 10, e207-e215.	6.3	23
36	Racial/Ethnic Segregation and Access to COVID-19 Testing: Spatial Distribution of COVID-19 Testing Sites in the Four Largest Highly Segregated Cities in the United States. <i>American Journal of Public Health</i> , 2022, 112, 518-526.	2.7	23

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37	Priority-Setting for Novel Drug Regimens to Treat Tuberculosis: An Epidemiologic Model. <i>PLoS Medicine</i> , 2017, 14, e1002202.	8.4	20
38	How could preventive therapy affect the prevalence of drug resistance? Causes and consequences. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015, 370, 20140306.	4.0	19
39	Evaluation of Tuberculosis Treatment Response With Serial C-Reactive Protein Measurements. <i>Open Forum Infectious Diseases</i> , 2018, 5, ofy253.	0.9	19
40	Development of a Treatment-decision Algorithm for Human Immunodeficiency Virus-uninfected Children Evaluated for Pulmonary Tuberculosis. <i>Clinical Infectious Diseases</i> , 2021, 73, e904-e912.	5.8	19
41	Evaluating the reliability of mobility metrics from aggregated mobile phone data as proxies for SARS-CoV-2 transmission in the USA: a population-based study. <i>The Lancet Digital Health</i> , 2022, 4, e27-e36.	12.3	19
42	Where is tuberculosis transmission happening? Insights from the literature, new tools to study transmission and implications for the elimination of tuberculosis. <i>Respirology</i> , 2018, 23, 807-817.	2.3	17
43	Using Chemical Reaction Kinetics to Predict Optimal Antibiotic Treatment Strategies. <i>PLoS Computational Biology</i> , 2017, 13, e1005321.	3.2	16
44	Risk ratios for contagious outcomes. <i>Journal of the Royal Society Interface</i> , 2018, 15, 20170696.	3.4	16
45	Household studies provide key insights on the transmission of, and susceptibility to, SARS-CoV-2. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 1103-1104.	9.1	16
46	Time Since Infection and Risks of Future Disease for Individuals with Mycobacterium tuberculosis Infection in the United States. <i>Epidemiology</i> , 2021, 32, 70-78.	2.7	16
47	Phylogeography and transmission of M. tuberculosis in Moldova: A prospective genomic analysis. <i>PLoS Medicine</i> , 2022, 19, e1003933.	8.4	16
48	Cost-effectiveness of expanding the capacity of opioid agonist treatment in Ukraine: dynamic modeling analysis. <i>Addiction</i> , 2020, 115, 437-450.	3.3	15
49	Cost-effectiveness of post-treatment follow-up examinations and secondary prevention of tuberculosis in a high-incidence setting: a model-based analysis. <i>The Lancet Global Health</i> , 2020, 8, e1223-e1233.	6.3	15
50	Trends in C-Reactive Protein, D-Dimer, and Fibrinogen during Therapy for HIV-Associated Multidrug-Resistant Tuberculosis. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018, 99, 1336-1341.	1.4	15
51	ClassTR: Classifying Within-Host Heterogeneity Based on Tandem Repeats with Application to Mycobacterium tuberculosis Infections. <i>PLoS Computational Biology</i> , 2016, 12, e1004475.	3.2	14
52	Second line drug susceptibility testing to inform the treatment of rifampin-resistant tuberculosis: a quantitative perspective. <i>International Journal of Infectious Diseases</i> , 2017, 56, 185-189.	3.3	14
53	Impact of Effective Global Tuberculosis Control on Health and Economic Outcomes in the United States. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 202, 1567-1575.	5.6	14
54	Whole-genome sequencing of Mycobacterium tuberculosis for rapid diagnostics and beyond. <i>Lancet Respiratory Medicine</i> , the, 2016, 4, 6-8.	10.7	13

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55	A Likelihood Approach for Real-Time Calibration of Stochastic Compartmental Epidemic Models. <i>PLoS Computational Biology</i> , 2017, 13, e1005257.	3.2	13
56	Comparative Modeling of Tuberculosis Epidemiology and Policy Outcomes in California. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 201, 356-365.	5.6	13
57	Effect of empirical treatment on outcomes of clinical trials of diagnostic assays for tuberculosis. <i>Lancet Infectious Diseases</i> , The, 2015, 15, 16-17.	9.1	12
58	Identifying cost-effective dynamic policies to control epidemics. <i>Statistics in Medicine</i> , 2016, 35, 5189-5209.	1.6	12
59	The cost-effectiveness of alternative vaccination strategies for polyvalent meningococcal vaccines in Burkina Faso: A transmission dynamic modeling study. <i>PLoS Medicine</i> , 2018, 15, e1002495.	8.4	12
60	Use of daily Internet search query data improves real-time projections of influenza epidemics. <i>Journal of the Royal Society Interface</i> , 2018, 15, 20180220.	3.4	11
61	Drug-target binding quantitatively predicts optimal antibiotic dose levels in quinolones. <i>PLoS Computational Biology</i> , 2020, 16, e1008106.	3.2	11
62	Risk factors for recurrent tuberculosis after successful treatment in a high burden setting: a cohort study. <i>BMC Infectious Diseases</i> , 2020, 20, 789.	2.9	10
63	Evaluating the potential impact of enhancing HIV treatment and tuberculosis control programmes on the burden of tuberculosis. <i>Journal of the Royal Society Interface</i> , 2015, 12, 20150146.	3.4	9
64	Tradeoffs in Introduction Policies for the Anti-Tuberculosis Drug Bedaquiline: A Model-Based Analysis. <i>PLoS Medicine</i> , 2016, 13, e1002142.	8.4	9
65	Population implications of the use of bedaquiline in people with extensively drug-resistant tuberculosis: are fears of resistance justified?. <i>Lancet Infectious Diseases</i> , The, 2017, 17, e429-e433.	9.1	9
66	Towards better prediction of Mycobacterium tuberculosis lineages from MIRU-VNTR data. <i>Infection, Genetics and Evolution</i> , 2019, 72, 59-66.	2.3	9
67	Accurate quantification of uncertainty in epidemic parameter estimates and predictions using stochastic compartmental models. <i>Statistical Methods in Medical Research</i> , 2019, 28, 3591-3608.	1.5	9
68	Trends, Mechanisms, and Racial/Ethnic Differences of Tuberculosis Incidence in the US-Born Population Aged 50 Years or Older in the United States. <i>Clinical Infectious Diseases</i> , 2022, 74, 1594-1603.	5.8	9
69	Population Immunity to Pre-Omicron and Omicron Severe Acute Respiratory Syndrome Coronavirus 2 Variants in US States and Counties Through 1 December 2021. <i>Clinical Infectious Diseases</i> , 2023, 76, e350-e359.	5.8	9
70	Protective effects of household-based TB interventions are robust to neighbourhood-level variation in exposure risk in Lima, Peru: a model-based analysis. <i>International Journal of Epidemiology</i> , 2018, 47, 185-192.	1.9	8
71	Tracking and predicting U.S. influenza activity with a real-time surveillance network. <i>PLoS Computational Biology</i> , 2020, 16, e1008180.	3.2	8
72	Adaptive Policies to Balance Health Benefits and Economic Costs of Physical Distancing Interventions during the COVID-19 Pandemic. <i>Medical Decision Making</i> , 2021, 41, 386-392.	2.4	7

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73	Bayesian evidence synthesis to estimate subnational TB incidence: An application in Brazil. <i>Epidemics</i> , 2021, 35, 100443.	3.0	7
74	High-resolution estimates of tuberculosis incidence among non-U.S.-born persons residing in the United States, 2000–2016. <i>Epidemics</i> , 2020, 33, 100419.	3.0	6
75	Adaptive guidelines for the treatment of gonorrhea to increase the effective life span of antibiotics among men who have sex with men in the United States: A mathematical modeling study. <i>PLoS Medicine</i> , 2020, 17, e1003077.	8.4	6
76	Test, trace, and isolate in the UK. <i>BMJ</i> , The, 2021, 372, n822.	6.0	6
77	Use of Lot Quality Assurance Sampling to Ascertain Levels of Drug Resistant Tuberculosis in Western Kenya. <i>PLoS ONE</i> , 2016, 11, e0154142.	2.5	6
78	Neighbourhood prevalence-to-notification ratios for adult bacteriologically-confirmed tuberculosis reveals hotspots of underdiagnosis in Blantyre, Malawi. <i>PLoS ONE</i> , 2022, 17, e0268749.	2.5	6
79	Evolution and emergence of multidrug-resistant <i>Mycobacterium tuberculosis</i> in Chisinau, Moldova. <i>Microbial Genomics</i> , 2021, 7, .	2.0	5
80	Cost-Effectiveness of Alternative Uses of Polyvalent Meningococcal Vaccines in Niger: An Agent-Based Transmission Modeling Study. <i>Medical Decision Making</i> , 2019, 39, 553-567.	2.4	4
81	Ongoing challenges to understanding multidrug- and rifampicin-resistant tuberculosis in children versus adults. <i>European Respiratory Journal</i> , 2021, 57, 2002504.	6.7	4
82	The positive externalities of migrant-based TB control strategy in a Chinese urban population with internal migration: a transmission-dynamic modeling study. <i>BMC Medicine</i> , 2021, 19, 95.	5.5	4
83	Evaluation of 6-Month Versus Continuous Isoniazid Preventive Therapy for <i>Mycobacterium tuberculosis</i> in Adults Living With HIV/AIDS in Malawi. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2020, 85, 643-650.	2.1	4
84	Evidence sources on the natural history of latent tuberculosis infection. <i>Lancet Infectious Diseases</i> , The, 2018, 18, 834-835.	9.1	2
85	Assessing Local Risk of Rifampicin-Resistant Tuberculosis in KwaZulu-Natal, South Africa Using Lot Quality Assurance Sampling. <i>PLoS ONE</i> , 2016, 11, e0153143.	2.5	2
86	The Health and Economic Benefits of Tests That Predict Future Progression to Tuberculosis Disease. <i>Epidemiology</i> , 2022, 33, 75-83.	2.7	2
87	vCOMBAT: a novel tool to create and visualize a computational model of bacterial antibiotic target-binding. <i>BMC Bioinformatics</i> , 2022, 23, 22.	2.6	2
88	Antimicrobial Resistance Risks of Cholera Prophylaxis for United Nations Peacekeepers. <i>Antimicrobial Agents and Chemotherapy</i> , 2017, 61, .	3.2	1
89	Transmission Modeling with Regression Adjustment for Analyzing Household-based Studies of Infectious Disease. <i>Epidemiology</i> , 2020, 31, 238-247.	2.7	1
90	Children as sentinels of tuberculosis transmission: disease mapping of programmatic data. <i>BMC Medicine</i> , 2020, 18, 234.	5.5	1

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91	Evaluating spatially adaptive guidelines for the treatment of gonorrhea to reduce the incidence of gonococcal infection and increase the effective lifespan of antibiotics. <i>PLoS Computational Biology</i> , 2022, 18, e1009842.	3.2	1
92	The contributions of Steve Lawn to the science, advocacy and policy of HIV-associated TB. <i>International Journal of Tuberculosis and Lung Disease</i> , 2016, 20, 1563-1564.	1.2	0
93	Reply to Chen et al. <i>Journal of Infectious Diseases</i> , 2016, 214, 1287-1288.	4.0	0
94	82125 Multiple epidemics of multidrug-resistant tuberculosis revealed by spatial disease mapping and whole-genome sequencing analysis in urban China. <i>Journal of Clinical and Translational Science</i> , 2021, 5, 5-6.	0.6	0
95	Spatially targeted digital chest radiography to reduce tuberculosis in high-burden settings: A study of adaptive decision making. <i>Epidemics</i> , 2022, 38, 100540.	3.0	0
96	Title is missing!. , 2020, 17, e1003077.		0
97	Title is missing!. , 2020, 17, e1003077.		0
98	Title is missing!. , 2020, 17, e1003077.		0
99	Title is missing!. , 2020, 17, e1003077.		0