



COVID-19 and Tuberculosis: Findings from an ecological study

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7 **COVID-19 and Tuberculosis: Findings from an ecological analysis**

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31 **Abstract:**

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34 Coronavirus disease 2019 (COVID-19) represents an unprecedented challenge in modern public health
35 practice. As this pandemic grows in size and scope, understanding the associations between COVID-19
36 and other respiratory illnesses like tuberculosis (TB), the largest infectious disease killer globally, becomes
37 essential. In this ecological analysis, we demonstrate an association between COVID-19 case positivity and
38 increasing gross domestic product (GDP) per capita, in which case positivity increases by 45.10 per
39 100,000 people tested for a 10% increase in GDP per capita ($p=0.001$). We further examine this
40 relationship among countries designated as high-TB burden countries by the World Health Organization
41 (WHO) and observe a stronger association between COVID-19 case positivity and increasing GDP per
42 capita (79.41 per 100,000 people tested). Even after controlling for TB incidence, this association holds at
43 the ecological level (77.66 per 100,000 people tested) among high-TB burden countries. That these
44 findings are of a higher magnitude among high-TB burden countries suggests that these countries may
45 require additional support in responding to the COVID-19 pandemic. In a pandemic setting, we must
46 prioritize existing health system functions to avoid resurgence or exacerbation of infectious disease
47 threats like TB.
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Introduction

Coronavirus disease 2019 (COVID-19) represents an unprecedented challenge in modern public health practice. Having spread to over 180 countries and having affected over 2.7 million individuals, the COVID-19 pandemic requires a coordinated, effective response without sacrifice to quality or availability of other essential medical services.¹ This rapidly-moving pandemic has laid bare the importance of effective surveillance, quarantine, testing and diagnosis, contact tracing and hospital infection prevention and control measures. Certain countries, including Singapore, have effectively contained community spread of the virus through early and broad quarantine, testing and contact tracing measures.²⁻⁴ In countries like Italy and the United States – where responses were slower, narrower in focus and less consistently implemented – COVID-19 transmission has spread widely in the community, and responses have shifted from containment to mitigation.⁴⁻⁶ In these countries where containment measures have failed, health systems must plan to ration life-saving medical equipment like mechanical ventilators.⁷⁻⁹

Tuberculosis (TB), from which more people die than any other single infectious agent, represents a similar public health challenge in that it is transmitted through respiratory droplet spread, causes great economic harm (either directly through treatment costs or indirectly through lost wages and family support) and requires frequent testing and contact tracing to reduce spread in the community.^{10,11} TB disproportionately affects people with lower income, which is explained through structural factors like housing insecurity and more frequent exposure to congregate settings, economic factors which prevent access to high-quality preventive or curative health services except in dire emergencies, and social factors like delaying reporting or care to prevent loss of wages or social capital.¹²⁻¹⁶

As noted by the WHO, the Stop TB partnership and other organizations, we must ensure the provision of essential TB services while responding to the COVID-19 pandemic. As the Stop TB partnership noted in April through a rapid assessment of the impact of the COVID-19 pandemic on the TB response in high-burden countries, there are significant disruptions to TB services in high-burden countries.¹⁷ Based on these results, we must understand the relationships between COVID-19 incidence, TB burden and gross domestic product (GDP) per capita and generate hypotheses for creating interventions which maintain TB services in the context of increasing COVID-19 caseloads.

In this ecological analysis, we seek to understand possible associations between COVID-19 cases and GDP per capita and TB and GDP per capita. We also aim to elucidate whether these relationships differ when restricted to high-burden TB countries alone. Finally, we seek to compare these individual disease analyses to understand any similarities in the trends observed between the individual diseases and GDP per capita.

Methods

COVID-19 case data were retrieved from the Johns Hopkins Center for Systems Science and Engineering (JHU CSSE) Coronavirus Resource Center and include confirmed COVID-19 cases through 4-April 2020.¹⁸ We retrieved all confirmed COVID-19 cases at the national level for all available countries. We also retrieved COVID-19 testing data from Worldometer, in which we collected number of tests reported at the national level. Testing data were available for 148 countries.¹⁹ Tuberculosis incidence data were retrieved from available WHO data as of 4-April 2020, in which we included data from 2018 for all countries with available data.²⁰ A total of 142 countries had data for COVID-19 cases, COVID-19 testing and TB incidence. GDP per capita and population data were retrieved from World Bank data sources, for which we retained 2018 values from each available country.²¹ A total of 133 countries had available data for COVID-19 cases, TB incidence and GDP per capita.

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3 We collated and examined data using Stata/SE 15.1 (StataCorp, College Station, TX). We computed crude
4 incidence of TB per 100,000 people using WHO estimated incident TB cases in 2018 and World Bank
5 population estimates for 2018. We then considered creating a COVID-19 incidence measure based on
6 confirmed COVID-19 cases and World Bank population statistics, but found this to be insufficient in
7 describing incidence due to wide variation in testing coverage globally. We decided to use COVID-19
8 positivity, in which we calculated the number of confirmed COVID-19 cases per 100,000 people tested for
9 SARS-CoV-2 (the viral agent that causes COVID-19) using JHU CSSE confirmed case data to date and
10 Worldometer testing data as a proxy for COVID-19 incidence. Given that testing rates vary in many
11 countries, that many COVID-19 cases are mild, and that many cases are treated based on clinical suspicion
12 versus a confirmed test, we decided to use case positivity as the most robust comparative measure given
13 the available data.
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17 We began by examining the distribution of incident TB cases per 100,000 people, COVID-19 case positivity
18 per 100,000 people tested and GDP per capita. We reviewed box plots for each distribution and noted
19 potential outliers. We noted that high values of GDP per capita (above \$50,000 USD) were potential
20 outliers and that these values skewed the distribution, so we transformed GDP per capita to the natural
21 logarithm of GDP to normalize the distribution. We chose this transformation in favor of a square-root
22 transformation. We did not transform COVID-19 case positivity per 100,000 people tested nor TB
23 incidence per 100,000 people.
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26 We plotted both COVID-19 case positivity per 100,000 people tested and TB incidence per 100,000 people
27 against the natural logarithm of GDP using Tableau Desktop Professional Edition 10.5 (Tableau Software,
28 Inc., Seattle, WA). We also plotted these values among the 30 highest TB burden countries (of which 20
29 had fully available data) according to the latest WHO Global TB Report (2019) to understand any
30 differences in these plots compared to the general country plots. We performed a simple linear regression
31 (SLR) of COVID-19 case positivity per 100,000 people tested on the natural logarithm of GDP per capita
32 and an SLR of TB incidence per 100,000 people on the natural logarithm of GDP to understand possible
33 crude associations at the ecological level. Finally, we performed a multiple linear regression of COVID-19
34 case positivity per 100,000 people tested and TB incidence per 100,000 people on the natural logarithm
35 of GDP to understand possible remaining associations between COVID-19 and GDP per capita after
36 controlling for TB incidence per 100,000 people. All statistical analyses were performed using Stata/SE
37 15.1 (College Station, TX).
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41 **Results**

42 TB incidence varied from 0 to 549.2 per 100,000 with an interquartile range (IQR) of 8.7 to 143.6 per
43 100,000. COVID-19 case positivity per 100,000 people tested varied from 57.7 to 30,614.7 per 100,000
44 with an IQR of 1,184.2 to 5,281.2 per 100,000. GDP per capita ranged from \$389.40 USD to \$87,208.54
45 with an IQR of \$2,028.90 to \$17,277.97. The natural logarithm of GDP per capita ranged from 5.751 to
46 11.667 with an IQR of 7.667 to 9.867. Figure 1 describes the change in distribution observed after
47 transforming GDP per capita to the natural logarithm of GDP per capita.
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50 The distributions of incident TB cases per 100,000 people and natural logarithm of GDP per capita and of
51 COVID-19 case positivity per 100,000 people tested and natural logarithm of GDP per capita are presented
52 in Figures 2a and 2b, respectively. Figure 2a describes a slightly positive association between proxy COVID-
53 19 incidence per 100,000 people tested and the natural logarithm of GDP, whereas figure 2b describes a
54 negative association between TB incidence per 100,000 people and the natural logarithm of GDP. Figures
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3a and 3b present these same findings, respectively, after having restricted to the 30 highest TB burden countries (among which 20 had complete data).

The results of our simple linear regression analyses are presented in Table 1. Considering our use of a log-transformed predictor on a linear regression, we present some methodological details to guide interpretation along with our results.²²

In the first SLR (GDP on natural logarithm of COVID-19 case positivity), we use the following model:

$$COVID19\ Case\ Positivity = \beta_0 + \beta_1 \times \ln(GDP)$$

Using a 10 percent change in GDP, we arrive at the following:

$$COVID19(GDP_2) - COVID19(GDP_1) = \beta_1 \times \left[\ln \left(\frac{GDP_2}{GDP_1} \right) \right] = \beta_1 \times \ln(1.1)$$

To interpret these results, we understand that a unit increase in GDP will be the same, regardless of baseline. In linear terms, we take the log of the percentage increase (10%, for the purposes of this analysis) and multiply by the predictor (GDP) coefficient. We then interpret the result as the change in COVID-19 case positivity per 100,000 people tested after a 10% increase in GDP per capita. We present original coefficients, change in COVID-19 case positivity per 100,000 tested per 10% increase in GDP, p-values and 95% confidence intervals for the original coefficient. The same interpretation can be applied to the SLR results for TB incidence, where we used the following model:

$$TB\ Incidence = \beta_0 + \beta_1 \times \ln(GDP)$$

Table 1. Results of three simple linear regressions examining change in COVID-19 case positivity with change in GDP per capita in \$USD. Outcome variable is COVID-19 case positivity and change in GDP is 10%.

	Log(GDP) all countries	Log(GDP), high-burden countries	TB incidence per 100,000 people
Coefficient	1,089.68	1,918.45	-51.05
Change in COVID-19 case positivity per 100,000 per 10% increase in GDP per capita	45.10	79.41	-2.11
p-value	0.001	0.065	<0.001
95% Confidence Interval	(426.64 - 1752.71)	(-130.46 - 3,967.37)	(-62.88 - -39.22)

In the multiple linear regression model, we noted similar results as in the individual SLRs in Table 1. Using the following model, we sought to understand the change in COVID-19 case positivity per 100,000 people tested with a 10% increase in GDP after controlling for TB incidence per 100,000 people:

$$COVID19\ Case\ Positivity = \beta_0 + \beta_1 \times TB\ Incidence + \beta_2 \times \ln(GDP)$$

After controlling for TB incidence per 100,000 people, we found a significant relationship between COVID-19 case positivity per 100,000 people tested and GDP. With 10% increase in GDP, we observed a 35.97 per 100,000 people tested (95% CI: 67.62 – 1,760.57; p=0.034) increase in COVID-19 case positivity per

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3 100,000 people tested after controlling for TB incidence. Among high-TB burden countries, we observe an
4 increase in case positivity of 77.66 per 100,000 people tested (95% CI: -294.86 – 4,047.15; p=0.086).

6 **Discussion**

7 These findings confirm the negative association between TB and GDP per capita, and further elucidate
8 associations between COVID-19 and GDP per capita, even after controlling for TB incidence. We also note
9 that among high-burden TB countries specifically, the increase in COVID-19 case positivity per 100,000
10 people tested is of a larger magnitude (79.41, 95%CI: -130.46 – 3,967.37, p=0.065) than the general
11 relationship when including all countries. This finding raises important concerns for COVID-19 control in
12 high-burden TB countries.
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15 This study suffers from certain limitations, chief among these is the ecological design. The authors have
16 taken care in ensuring standardized data sources and calculations while remaining focused in the findings
17 and analysis from this study. We recognize that these results are only applicable at the population level
18 and do not represent individual level data or findings. This analysis remains useful in that it helps identify
19 possible associations between COVID-19 and GDP per capita among all countries affected by COVID-19
20 and specifically among the highest-burden TB countries.
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23 Despite these limitations, this study remains as one of the first to examine the relationship between
24 COVID-19 and GDP per capita in the context of TB incidence and GDP per capita, and one of the first to
25 discuss these results in the context of high-burden TB countries. These countries are at especially high risk
26 for COVID-19 transmission due to the extant population with lung disease and the possibly enabling
27 environments which drive TB transmission, including congregate settings, migrant workers, smokers and
28 other people at especially high risk for TB. We must leverage existing TB control programs in these
29 countries to effectively respond to COVID-19 and to prevent an increase in incident TB cases during a
30 demand surge for emergency services, when regular TB services may be disrupted or unavailable.
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33 Areas for collaboration are numerous and include prevention, diagnosis, treatment and care, digital health
34 technology, procurement, supply chain, risk management, human resources and capacity building efforts.
35 We present some examples for a collaborative, TB-aware COVID-19 response below:
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38 *Prevention:* While modes of transmission differ between these diseases, similar measures can be taken to
39 prevent infection in congregate settings and in health care facilities. These include application of
40 administrative controls, environmental controls and personal protective equipment, which are hallmarks
41 of successful TB control programs. These existing efforts should be intensified in light of COVID-19 and
42 leveraged to ensure a seamless response.
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45 *Testing and diagnosis:* Accurate diagnostic tests remain essential for TB and for COVID-19. While testing
46 methods differ, there are extensive TB sample transport and testing mechanisms in place in high-TB
47 burden countries which should be leveraged for COVID-19 testing and diagnosis.
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49 *Human resources and capacity building:* Physicians, nurses, pulmonologists and TB specialists, along with
50 nurses and health care workers at the primary care level should be trained on COVID-19 diagnosis and
51 treatment protocols including updated case definitions and care protocols. TB specialists serve as an entry
52 point for buy-in and training as this cadre has similar training in infection prevention and control and in
53 decentralized patient management. COVID-19 responses can benefit from the extensive capacity building
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3 network in high-TB burden countries, in which infection prevention and control, contact tracing,
4 household and community-based care and surveillance and monitoring systems are robust and functional.
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6 National TB control programs, especially in high-burden TB countries, are well-suited for respiratory
7 disease control and should be consulted and funded to aid in the COVID-19 response. In an unprecedented
8 public health emergency, we must leverage every possible opportunity to collaborate and to build on
9 existing systems, rather than building new and untested parallel structures.
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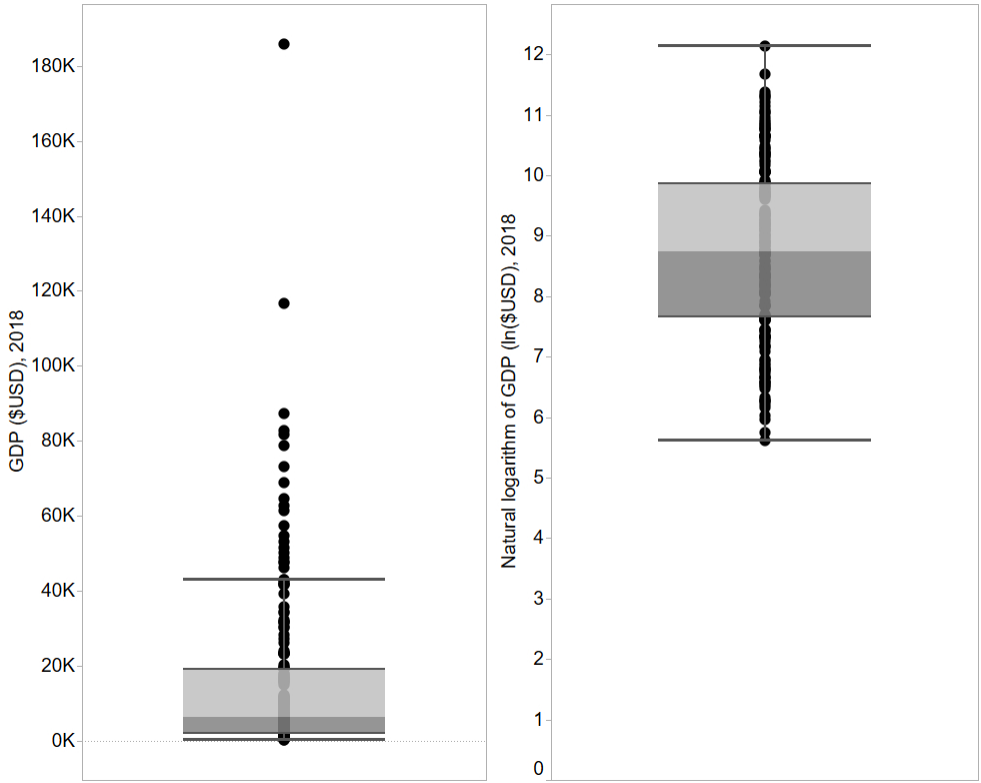
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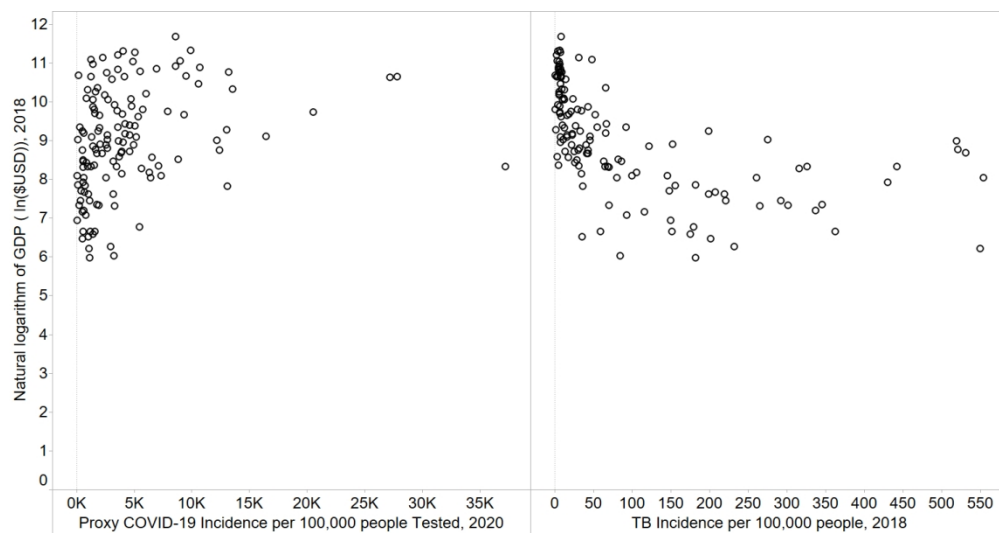
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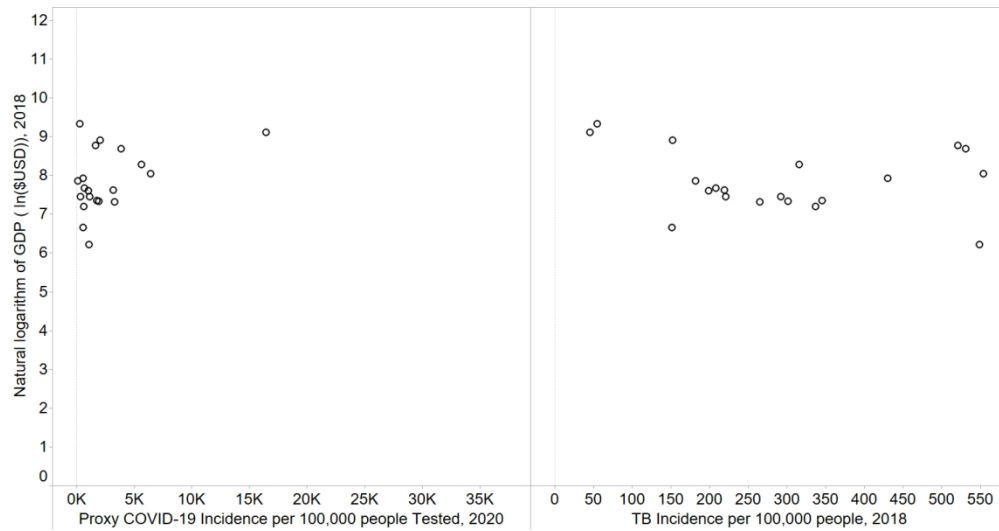
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Left: Distribution of GDP per capita, 2018. Right: Distribution of natural logarithm of GDP per capita, 2018



a (left): COVID-19 case positivity per 100,000 people tested, 2020 (to date) vs. the natural logarithm of GDP per capita, 2018. b (right): TB incidence per 100,000 people, 2018 vs. the natural logarithm of GDP per capita, 2018.



a (left): COVID-19 case positivity per 100,000 people tested, 2020 (to date) vs. the natural logarithm of GDP per capita, 2018 among high burden TB countries. b (right): TB incidence per 100,000 people, 2018 vs. the natural logarithm of GDP per capita, 2018 among high burden TB countries.