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Modelling the social and structural determinants of tuberculosis: opportunities and challenges

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SUMMARY

INTRODUCTION: Despite the close link between tuberculosis (TB) and poverty, most mathematical models of TB have not addressed underlying social and structural determinants.

OBJECTIVE: To review studies employing mathematical modelling to evaluate the epidemiological impact of the structural determinants of TB.

METHODS: We systematically searched PubMed and personal libraries to identify eligible articles. We extracted data on the modelling techniques employed, research question, types of structural determinants modelled and setting.

RESULTS: From 232 records identified, we included eight articles published between 2008 and 2015; six employed population-based dynamic TB transmission models and two non-dynamic analytic models. Seven studies focused on proximal TB determinants (four on nutritional status, one on wealth, one on indoor air pollution, and one examined overcrowding, socio-economic and nutritional status), and one focused on macro-economic influences.

CONCLUSIONS: Few modelling studies have attempted to evaluate structural determinants of TB, resulting in key knowledge gaps. Despite the challenges of modelling such a complex system, models must broaden their scope to remain useful for policy making. Given the intersectoral nature of the interrelations between structural determinants and TB outcomes, this work will require multidisciplinary collaborations. A useful starting point would be to focus on developing relatively simple models that can strengthen our knowledge regarding the potential effect of the structural determinants on TB outcomes.

KEY WORDS: mathematical modelling; tuberculosis; social determinants

TUBERCULOSIS (TB) is widely recognised as a disease of poverty,1–3 with disproportionate disease burden falling on the poorest in society and the most vulnerable communities. The need to design and implement comprehensive strategies to achieve TB elimination through universal health coverage and interventions to address the underlying social determinants of TB is a key element of the World Health Organization’s (WHO’s) End TB strategy for 2015–2035.4,5

The targets and indicators of this new TB action framework are anchored in the 17 Sustainable Development Goals (SDGs) adopted by the United Nations and which mark the global development agenda that began on 1 January 2016. By placing their emphasis on the interdependence and synergies between socio-economic development and health, these offer unique entry points for addressing the social determinants (SDs) of TB.

In the present article, we follow the definition of SDs of health of the WHO Commission on Social Determinants of Health:7 ‘The structural determinants of TB are those conditions that generate or reinforce social stratification (e.g., socio-economic inequalities, population growth, urbanisation), and therefore give rise to an unequal distribution of key social determinants of TB epidemiology, such as poor housing, poverty and malnutrition, which in turn influence exposure to risk, vulnerability and ability to recover after developing the disease.’8 These definitions are shown in Table 1.

Quantitative analytical tools such as mathematical modelling can play an important role in informing the End TB Strategy, evaluating the impact of novel
poverty-reduction interventions nested in its vision (including in combination with existing biomedical tools), and exploring the contribution of socio-economic drivers to the epidemic. However, to do so, TB models will inevitably need to expand their focus beyond diagnosis and treatment to incorporate SDs, but the potential of modelling as well as its main limitations in supporting this research agenda remain unclear.

In the present paper, we report findings from a systematic review of the literature carried out with the aim to provide an overview of the current state of knowledge in the mathematical modelling of SDs of TB. We then go on to discuss key methodological challenges and gaps in empirical evidence that existing mathematical models need to overcome to be able to incorporate SDs to remain relevant to policy-making.

**METHODS**

*Search strategy and selection criteria*

For the purposes of this review, ‘mathematical model’ was defined based on that envisaged by Garnett et al. as a mechanistic representation of how disease burden is established. This includes both dynamic transmission and decision (non-dynamic) analytic models. We searched PubMed for any relevant article on modelling and the socio-economic determinants of TB (e.g., nutrition, crowding, poverty). The full search string is included in Table 2. Titles and abstracts were screened for eligibility. Articles were eligible for full-text review if they were written in English (due to limited resources), the target population was human individuals and mathematical modelling assessed the epidemiological impact of the SDs of TB.

We excluded systematic reviews, epidemiological studies that did not use mathematical modelling techniques and ecological analyses looking at the SDs of TB. The search focused on socio-economic factors (i.e., the intervention or exposure involving a socio-economic factor), and excluded studies focusing only on diabetes mellitus (DM), the human immunodeficiency virus (HIV) and behavioural risk factors such as alcohol consumption and smoking unless their association with socio-economic factors were also considered. We applied no restrictions as to the year.

Additional relevant articles were identified in the authors’ personal libraries and are included in the review. DP selected the articles with support from RMGJH, DB and KL; data were extracted by DP and RMGJH.

*Data abstraction and synthesis*

Figure 1 presents the details of the selection process. The aim of the study, first author and publication dates, type and feature of the model, the socio-economic factor, the setting and the main findings were extracted into a pre-designed form. We focused on a qualitative synthesis of the methods employed in the articles we identified.

**RESULTS**

A total of 229 unique records were found in the literature search, and four additional articles were added from the authors’ personal libraries. Of these, 54 underwent full-text evaluation. After full-text screening, we included eight articles published between 2008 and 2015, with four articles published...
in 2015 only. Table 3 gives the main features of the selected studies.12–19

**Socio-economic factors investigated**

The study by Reeves et al. was the only article that looked at the impact of distal determinants, i.e., government expenditure per capita on public health services, gross domestic product (GDP) and cumulative decline in GDP, as a measure of the severity of an economic recession on TB control.12 The remainder modelled proximal TB determinants: four focused on nutritional status (body mass index [BMI] and undernutrition),14,16,18,19 one on wealth,15 one on smoking and indoor air pollution,13 and one on nutritional status, overcrowding and socio-economic status.17 All studies looked at one factor at a time, with the exception of the study by Dye et al.,19 which also explored the combined effect of nutritional status and demographic changes, including age structure and urbanisation, on TB incidence.

**Modelling methods, structure and parameters**

Compartmental population-based dynamic TB transmission models were the most common simulation approach employed in the selected articles (6/8, 75%); two studies used non-dynamic analytical models and both investigated the effect of both DM and nutritional status on TB epidemics. Most studies included a conceptual framework to illustrate the mechanics of the models and the hypotheses behind their research questions.

Transmission models employed standard SLIR (susceptible-latent-infectious-recovered) models that were adapted to explore the research question set in each study: the model by Oxlade et al., for example, was stratified by levels of undernutrition by wealth quartile.15 Andrews et al. implemented a parallel structure for two wealth groups to a standard TB model to explore the benefit of assortative mixing to interventions targeting the poor, highlighting the potential importance of including mixing parameters in TB models even if data are currently not available to inform these models.13

With regard to model parameters, Ackley et al. explored changes in differences in susceptibility to infection and progression to disease in hypothetical scenarios.16 Different levels of BMI drove changes in reactivation and progression parameters in the model used by Oxlade et al.15 The study by Reeves et al. used an econometric analysis to estimate changes in relevant model parameters controlling case detection.12 Bhunu et al. divided the population into ‘rich’ and ‘poor’ communities, and compared the reproduction numbers for these two strata (Appendix Table A).17*

Data on the different exposures were mainly drawn from the literature,16,18 national population-based surveys13–15,19 or publicly available databases.12 Very few data employed in these studies were local or regional. The majority of the studies were calibrated to TB data (e.g., incidence trends or point estimates) from WHO estimates.

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* The appendix is available in the online version of this article, at http://www.ingentaconnect.com/content/iuatld/ijtld/2017/00000021/00000009/art00005
<table>
<thead>
<tr>
<th>Author, year, reference</th>
<th>Aim of the study</th>
<th>Key socio-economic factors investigated</th>
<th>Mathematical modelling methods/type and features of the simulation model</th>
<th>Setting</th>
<th>Conclusion(s)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reeves, 2015</td>
<td>To project the potential influence of the economic recession on TB epidemiology in Europe until 2030</td>
<td>Government expenditure per capita on public health services, GDP and cumulative decline in GDP during the recession period as a measure of the severity of the recession</td>
<td>Dynamic model: SLIR model. Authors applied the findings from the preceding econometric models to dynamic mathematical models of TB transmission and mortality. The mathematical models simulated longitudinal TB rates in each country—given the data on case detection observed before, during and after the financial crisis—as well as a counterfactual scenario in which case detection was unaffected by either the recession or the related austerity.</td>
<td>Europe</td>
<td>Recession can lead to short-term reductions in the financial support of programmes for TB control. The associated decrease in the detection of TB is projected to result in sustained, long-term rises in TB incidence, prevalence and mortality.</td>
</tr>
<tr>
<td>Andrews, 2015</td>
<td>To illustrate the role of social mixing in shaping disparities in the distribution of TB, and demonstrate how the concentration of disease risk and transmission among the poor presents challenges and opportunities for TB control</td>
<td>Wealth</td>
<td>Dynamic: deterministic, compartmental model with parallel structure for two wealth groups with varying parameters, contact rates and social mixing</td>
<td>India</td>
<td>TB control efforts may benefit from preferential targeting toward the poor.</td>
</tr>
<tr>
<td>Lin, 2008</td>
<td>To predict the effects of risk factors trends on COPD, lung cancer and TB</td>
<td>Smoking, solid fuel use</td>
<td>Dynamic: dynamic TB transmission model: deterministic compartmental (SLIR)</td>
<td>China</td>
<td>Reducing smoking and solid fuel use can substantially reduce predictions of COPD and lung cancer burden and would contribute to effective TB control in China (even when DOTS implementation is less effective).</td>
</tr>
<tr>
<td>Oxlade, 2015</td>
<td>To project future trends in TB-related outcomes under different scenarios for reducing undernutrition in the adult population in the Central Eastern states of India</td>
<td>Undernutrition</td>
<td>Dynamic: compartmental TB transmission model stratified by BMI parameterised using national and regional data from India (model population is stratified into four exposure levels defined by the mean BMI for each quartile).</td>
<td>India</td>
<td>Intervening on under-nutrition could have a substantial impact on TB incidence and mortality in areas with high prevalence of under nutrition.</td>
</tr>
<tr>
<td>Ackley, 2015</td>
<td>To explore the population-level effects of malnutrition and genetic heterogeneity in TB susceptibility on TB epidemics</td>
<td>Malnutrition, genetic heterogeneity</td>
<td>Dynamic: dynamic TB transmission model: deterministic compartmental SLIR model</td>
<td>First Nations community in Canada</td>
<td>1) Changes in a population's nutritional status can have significant effects on TB dynamics. 2) Inclusion of heterogeneity in susceptibility to M. tuberculosis infection or risk of TB disease yields improved fit to data.</td>
</tr>
<tr>
<td>Bhunu, 2012</td>
<td>To assess the impact of socio-economic conditions on TB transmission, taking into account heterogeneous mixing patterns</td>
<td>Socio-economic conditions (overcrowding, increased endogenous reactivation, reduced socio-economic status, reduced treatment uptake and poor nutrition on TB dynamics)</td>
<td>Dynamic: dynamic TB transmission model: deterministic compartmental SLIR model</td>
<td>Zimbabwe</td>
<td>Poverty enhances TB transmission as overcrowding, poor nutrition, reduced treatment uptake and lower socio-economic status worsen TB; TB transmission rates are therefore higher in poor communities than in rich ones.</td>
</tr>
</tbody>
</table>
Key findings of the modelling studies

The studies in our review support the notion that TB control is linked to and would benefit from action on TB social determinants. Reeves et al. found that a decrease in funding to control TB due to an economic recession (distal factor) can lead to a decline in TB case detection, and consequently to higher TB rates. Lin et al. showed that interventions on smoking and indoor air pollution (proximal factors) can accelerate TB decline. The studies that focused on nutritional status (proximal factor) found that reducing undernutrition would substantially reduce TB incidence. Andrews et al. showed that preferential targeting of the poor can benefit TB control (wealth as a proximal factor). From the analysis of reproduction numbers for the poor and rich communities, Bhunu et al. found that overcrowding, poor nutrition, lower socio-economic status (proximal factors) and reduced TB treatment uptake worsened TB transmission. Finally, the study by Dye et al. concluded that a combination of nutritional and demographic changes (proximal factors) operating over the decade from 1998 tended to increase TB incidence per capita in high-burden India but reduce it in lower-burden Korea.

DISCUSSION

This review has highlighted the paucity of mathematical modelling studies looking at the effects of socio-economic factors on TB pathogenesis and epidemiology, but has also shown that, although fairly recent, work in this field seems to be growing as the number of articles published has increased from 2011 onwards. This is possibly a reflection of changing policy priorities that are now part of the End TB Strategy.

Our findings point to the need, at this stage, to develop relatively simple models that improve and expand the current body of work to incorporate available evidence and strengthen our knowledge of the potential effect of SDs on TB outcomes. For instance, most models focused on one or two factors only, and those that considered two factors did not account for possible interactions between them. It is to be noted that most mathematical modelling studies focussed on assessing the effect of nutritional status and changes in BMI on TB epidemiology. This is not surprising, as undernutrition has long been acknowledged as a key socially determined TB risk factor. We found no modelling work looking at the impact of improved socio-economic macro-indicators on TB outcomes, or of social protection interventions targeting TB patients and their households. With respect to proximal risk factors, only one model assessed the effect of crowding on TB epidemiology, possibly a reflection of the fact that data on crowding...
and TB are not rich enough to unpick causality for a model.

**Challenges in translating from determinant to model**

The narrow focus of past global health and development policies and TB control strategies only partly explains why TB modelling has so far shown some reluctance to include SDs. This has also been due to the real and perceived weaknesses in the empirical evidence needed to populate models and quantify the pathways from socio-economic factors to changes in the natural history of TB in a population. Figure 2 provides a conceptual framework that outlines how distal/structural determinants (such as macro-economic policies) work through a potential array of more proximal determinants (e.g., crowding and nutrition), which in turn affects the dynamics of a standard mechanistic TB model at multiple points,²¹,²² such as the intensity of transmission (through crowding) or the rate of progression after recent and/or latent infection (e.g., through nutrition) (Figure 2).

While there are some data to inform parts of, for example, the pathway from macro-economic policies (e.g., GDP) to TB incidence,¹² our ability to quantify the exact relationship of each step remains limited. However, it should be noted that the same limitations apply to current TB models, ranging from capturing the impact of HIV or, when models look to evaluate the potential impact of interventions, including current approaches to improving case detection and reducing patient delay, or future hypothetical tools.²³,²⁴

When translating the effect of changing a socio-economic determinant into a mechanistic model, it does not suffice to have an estimate of the magnitude of the effect (see examples in Table 4).²⁰,²⁵–³⁴ One needs to know, or make assumptions about, the model parameters that should be changed to achieve the estimated impact. As shown in Figure 2, changes in disease risk may be due to influences at one or several of the stages on the pathway between exposure and disease that are captured by transmission models. As direct evidence is often still lacking, this means that choices need to be made based on the likely biological mechanism.

The range of these potential model parameters includes those that directly capture the intensity of transmission, e.g., social mixing or crowding in households, but also parameters guiding progression to disease after infection, which can be affected, for example, by nutritional status. It is also plausible that different paths to progression (primary, reactivation, reinfection) are affected at different stages of the pathway. In addition, any interventions that reduce barriers to care and treatment completion will change model parameters capturing the time to diagnosis as well as retention in care (e.g., alcohol and drug abuse).

In addition to effects on incidence, SDs may alter the natural history of disease (e.g., reduced infec-

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**Figure 2** Framework for proximate risk factors, upstream determinants and TB mechanics. Source: Adapted from Lonnroth et al.² This framework provides an example of the complexity when considering SDs in TB models: it illustrates the complicated cascade of parameters from distal to downstream determinants affecting the development of disease, and care and prevention. SD = structural determinants; SES = socio-economic status; HIV = human immunodeficiency virus; LTBI = latent tuberculous infection.
To further our knowledge, projects are urgently needed that advance the field while avoiding the pitfall of developing overly complex models that include population or pathway structures not adequately supported by empirical evidence or fully understood. In addition, the complexity of the pathways involved and the multisectoral nature of new approaches to end TB evidently require collaborations from different disciplines, including social scientists, epidemiologists, economists, policy makers as well as mathematical modellers. While recognising the importance of such projects but at the same time the struggle to identify suitable funding opportunities for such cross-disciplinary collaborative work, the TB Modelling and Analysis Consortium organised a meeting at the end of 2015 to discuss existing experiences and the potential path forward. A range of projects was developed that would both advance the field and be feasible, given current data. Two of these projects have been funded, preliminary results were produced at the end of 2016 and publications are under review: an interdisciplinary project looking at how social protection interventions can accelerate TB elimination (the Social Protection to Enhance the Control of TB Consortium, S-PROTECT), and a project assessing the relative contribution of TB programme (DOTS) expansion and improvements in socio-economic indicators on TB epidemiology in China.

In this article, we highlighted that the literature on mathematical modelling of social determinants of TB remains limited. We argue that to maintain its key role in policy discussions in the era of the SDGs and End TB Strategy, the TB modelling community needs to embrace the technical challenges to adequately represent the interplay between TB and its socio-economic drivers. While some work is underway, more funding, data and capacity are urgently needed to ensure TB modelling remains a useful tool for the ultimate goal of TB elimination.

**Acknowledgements**

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Conflicts of interest: none declared.

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**References**


**CONCLUSIONS AND RECOMMENDATIONS**

Mathematical modelling is a powerful and flexible tool to inform policy discussions and estimate the potential impact of various interventions relative to one another. However, to be useful, models need to be able to reflect the relevant aspects of the epidemic and address the questions faced by policy makers. In the SDGs and End TB Strategy era, this means that mathematical models of TB must translate the impact of socio-economic determinants into their mechanistic components. As a starting point, the TB modelling community should use the existing scientific evidence to construct relatively simple mechanistic models that add to our understanding of the effect of SDs on TB, and help improve specific policy decisions.

As shown in this article, there exists a scarcity of TB models that include SDs, but also a small but increasing body of work that has explored initial ideas. Some modelling of proximal risk factors and related public health interventions has been done but, for example, this has never moved upstream. TB models can leverage the existing data, and highlight the value of collecting those that are missing, such as the exact link between changes in nutritional status and changes in progression to disease, or the relationship between transmission intensity and living environments (e.g., urban slums compared with rural settings).

<table>
<thead>
<tr>
<th>Proximal determinant</th>
<th>Relative risk of TB disease</th>
<th>Author, year, reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>HIV infection</td>
<td>2–20</td>
<td>Corbett, 2013</td>
</tr>
<tr>
<td></td>
<td>1.4 per 100 cells/mm³</td>
<td>Sonnenberg, 2005</td>
</tr>
<tr>
<td></td>
<td>1.14 per decrement in CD4</td>
<td>Williams, 2005</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Lönneborn, 2010</td>
</tr>
<tr>
<td>Low BMI</td>
<td>2–4</td>
<td>Jeon, 2008</td>
</tr>
<tr>
<td>Diabetes mellitus</td>
<td>2–4</td>
<td>Stevenson, 2007</td>
</tr>
<tr>
<td>Alcohol use (&gt;=40 g/day)</td>
<td>2–5</td>
<td>Rehm, 2009</td>
</tr>
<tr>
<td>Smoking</td>
<td>1–5</td>
<td>Bates, 2007</td>
</tr>
<tr>
<td>Indoor air pollution</td>
<td>1–6</td>
<td>Lin, 2007</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Sumpter, 2013</td>
</tr>
</tbody>
</table>

TB = tuberculosis; HIV = human immunodeficiency virus; BMI = body mass index.


## APPENDIX

### Table A  Summary of model structure and parameters employed in the studies included in the review

<table>
<thead>
<tr>
<th>Study</th>
<th>Dynamic Non-dynamic Description</th>
<th>Parameters employed to capture the effect of socio-economic factors</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reeves, 2015[^12]</td>
<td>SLIR deterministic compartmental model</td>
<td>Parameter: diagnostic rate (the rate in %/year at which TB cases are diagnosed per year) Quantitative relationship: authors used the cumulative fall in GDP during the recession associated with falling case detection rates (from regression analysis, –0.22%) and applied it to dynamic models as a reduction in diagnosis rate</td>
</tr>
<tr>
<td>Oxlade, 2015[^15]</td>
<td>SLIR deterministic model</td>
<td>Parameter: rapid progression and reactivation rates by BMI stratum Quantitative relationship: relative risks of TB disease by BMI status directly applied to rapid progression and reactivation parameter values in each BMI stratum, i.e., relative risk of two for disease implemented as double the value for rapid progression and reactivation parameter values</td>
</tr>
<tr>
<td>Lin, 2008[^14]</td>
<td>SLIR deterministic compartmental model</td>
<td>Parameter: transmission and progression to disease Quantitative relationship: relative risks from systematic reviews, applied to specific strata. Effect on prevalence of latent infection applied as change in transmission</td>
</tr>
<tr>
<td>Ackley, 2015[^16]</td>
<td>SLIR deterministic compartmental model for historical TB epidemics among First Nation populations in Canada</td>
<td>Parameters: rapid progression to disease, reactivation, TB-specific mortality, immunity Quantitative relationship: model sampled from a range of relative risks of 1–3 to find fit to data</td>
</tr>
<tr>
<td>Bhunu, 2011[^17]</td>
<td>SLIR deterministic compartmental model</td>
<td>Parameters: contact rate, transmission upon contact, progression to disease, treatment access, death due to TB Quantitative relationship: theoretical scenarios where being poor leads to a higher probability of TB or death, and lower probability of accessing treatment</td>
</tr>
<tr>
<td>Odone, 2014[^18]</td>
<td>Analytic model where change in TB incidence is directly estimated based on prevalence of diabetes and undernutrition, and relative risk of disease given that risk factor. Authors estimate the effect of diabetes and undernutrition on TB incidence per person per year in different age groups, WHO regions, and over time in various scenarios</td>
<td>Parameter: prevalence of diabetes and/or undernutrition Quantitative relationship: relative risk of TB disease for diabetes and undernutrition</td>
</tr>
</tbody>
</table>

SLIR = susceptible-latent-infectious-recovered; TB = tuberculosis; GDP = gross domestic product; BMI = body mass index; WHO = World Health Organization
INTRODUCTION : En dépit du lien étroit entre tuberculose (TB) et la pauvreté, la majorité des modèles mathématiques de TB n’ont pas examiné les déterminants sous-jacents sociaux et structurels (SD). Dans cet article, nous revoyons les études employant une modélisation mathématique afin d’évaluer l’impact épidémiologique des SD de la TB.

MÉTHODE : Nous avons fait une recherche systématique sur PubMed et dans des bibliothèques personnelles pour identifier les articles éligibles. Nous avons extrait des données relatives aux techniques de modélisation employées, aux sujets de recherche, le type de SD modélisés et le contexte.

RÉSULTATS : A partir de 232 documents identifiés, nous avons inclus huit articles publiés entre 2008 et 2015 ; six ont employé des modèles dynamiques basés en population de la transmission de la TB. Sept études se sont focalisées sur les déterminants proximaux de la TB (quatre sur le statut nutritionnel, une sur le niveau économique, une sur la pollution de l’air intérieur et une a examiné la surpopulation et le statut socioéconomique et nutritionnel) et une s’est concentrée sur les influences macroéconomiques.

CONCLUSION : Peu d’études de modélisation ont tenté d’évaluer les SD de la TB, ce qui a abouti à des lacunes majeures en termes de connaissances. En dépit des défis de la modélisation d’un système aussi complexe, les modèles doivent élargir leur portée pour rester utiles aux décisions de politique. Étant donné la nature intersectorielle des interrelations entre SD et évolution de la TB, ce travail va requérir une collaboration multidisciplinaire. Un point de départ utile consisterait à se concentrer sur l’élaboration de modèles relativement simples qui peuvent renforcer nos connaissances relatives à l’effet potentiel des SD sur l’évolution de la TB.