# Modeling Intervention Strategies for United States TB Control

Jessica Ginepro, Emma Hartman, Ryo Kimura, Matthew McDermott, Colin Pawlowski, & Dylan Shepardson Mathematical Modeling Group, Mount Holyoke College, South Hadley, MA, USA

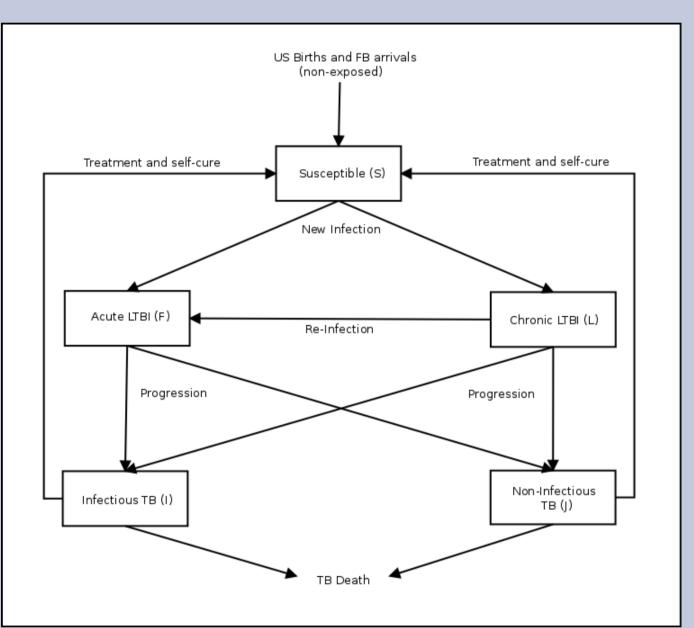




#### Introduction

Epidemiological models offer insight into the structure of disease outbreaks and the merits of various interventions. Compartmental differential equation models are a common model in which populations move between various health states, or compartments, according to predetermined rates. This work is an extension of the Hill Model, a complex compartmental model of tuberculosis (TB) in the United States.

#### The Basic Hill Model



Populations:

- ► US Born (USB)
- ► Foreign Born (FB)
  Individuals also leave the model due to natural death.

Figure: The Hill Model schematic

- ► USB TB incidence rates are declining
- ► FB latent TB infection (LTBI) arrivals remain high
- ► TB elimination in total population not projected by 2100

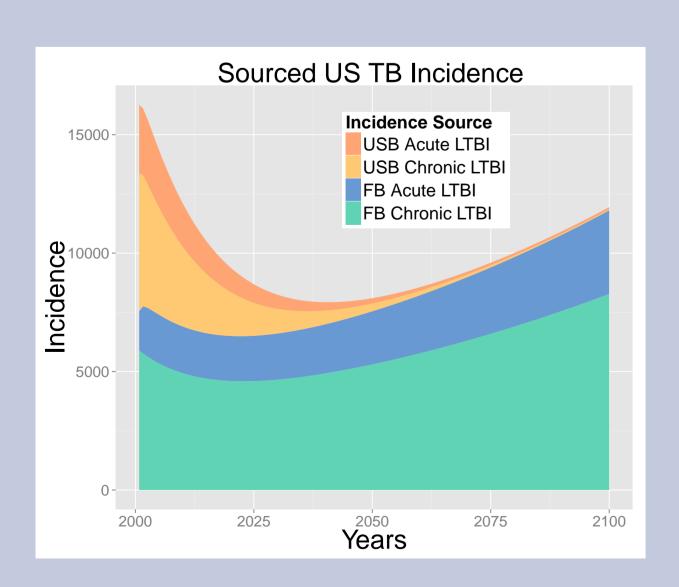


Figure : The source population of US TB incidence

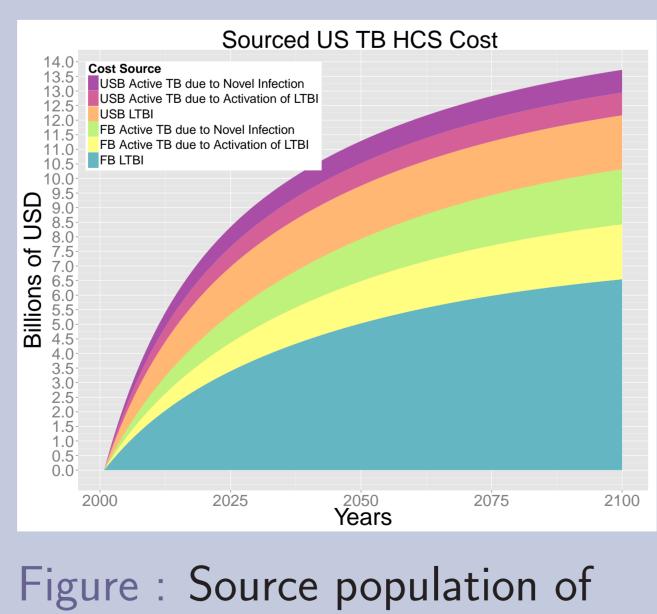


Figure : Source population of US HCS TB cost

## Analyzing US TB Reduction Strategies

- Implemented in R, with various numerical DE solvers
- ► Tracks US Health Care System (HCS) cost
- ► Tracks statistics about various health states

#### Intervention Analysis

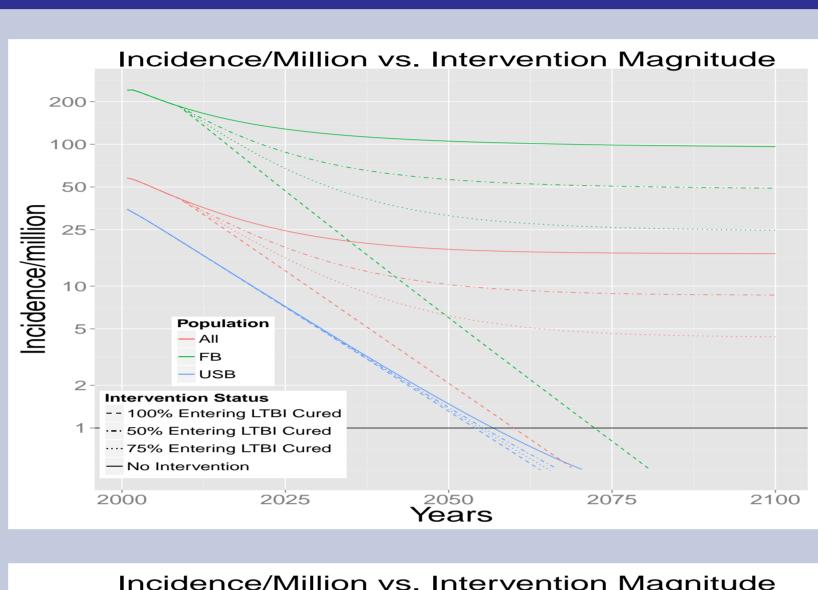


Figure:
Incidence/million in
USB, FB, and total
populations, given 0%,
50%, 75%, or 100%
treatment of incoming
LTBI

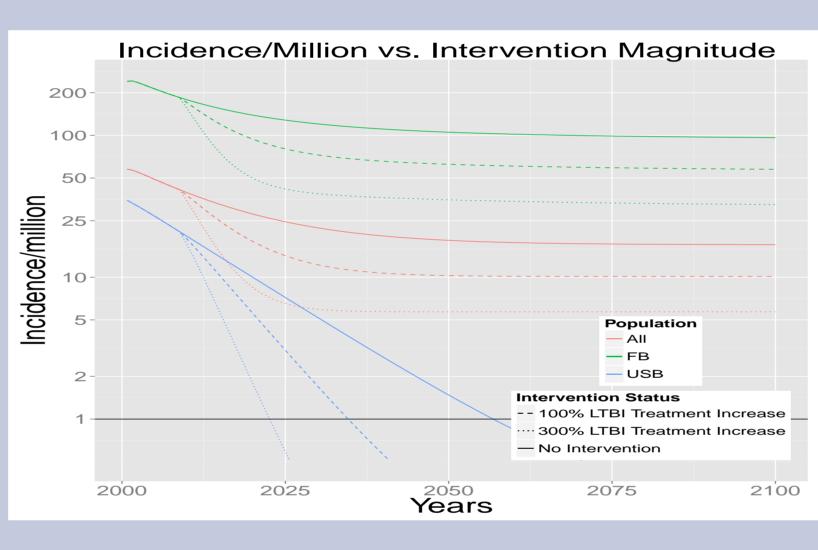


Figure:
Incidence/million in
USB, FB, and total
populations, given 0%,
100%, or 300% LTBI
treatment increase

# **Economic Modeling**

- ► Tracks treatment costs for various disease states
- Estimates implementation cost of intervention

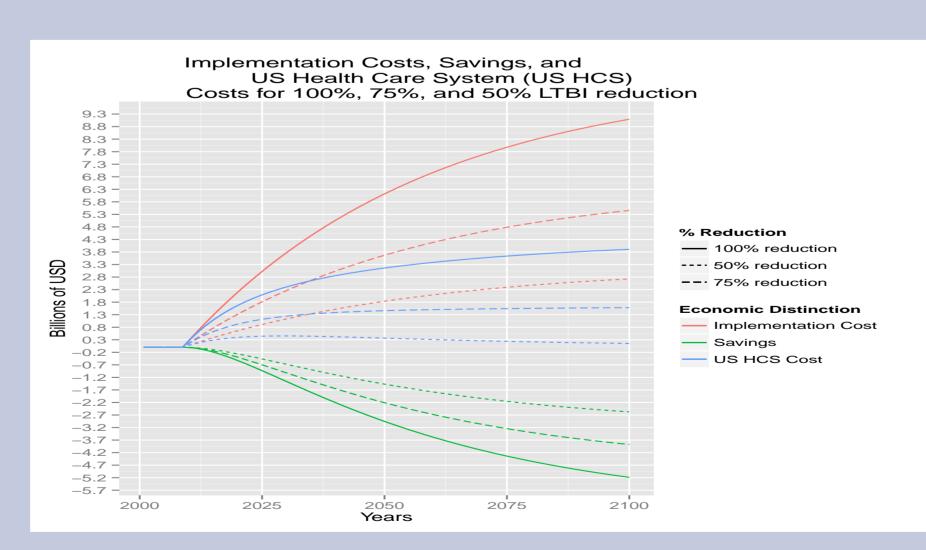
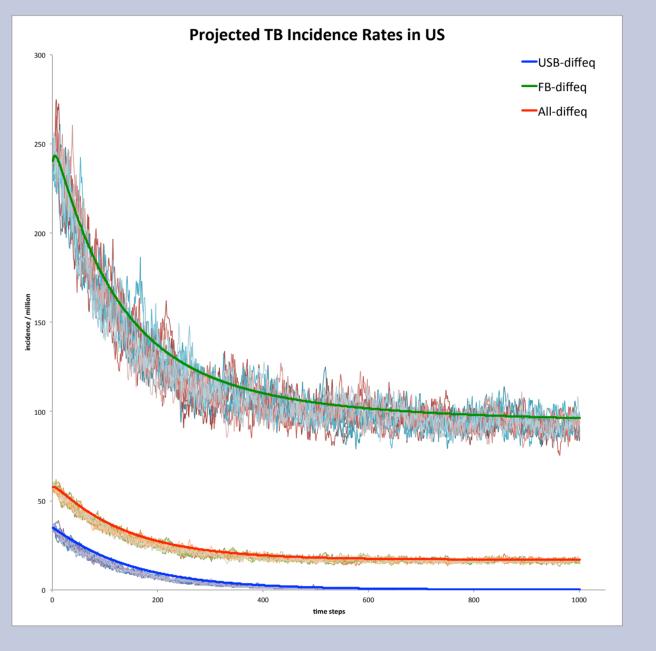


Figure: Cumulative implementation costs, US HCS savings, and net US costs of LTBI arrival cure rates. Cost/case cured was \$600, \$800, and \$1000 for 50%, 75%, and 100% cured

Base HCS
Costs:
Active TB:
\$14,014.90
LTBI:
\$403.45

# An Agent-Based Implementation

Agent-based models capture disease dynamics on the individual level and reflect stochasticity and granularity lost in compartmental models. Agent-based counterparts to the Hill model were implemented in Netlogo and C++.



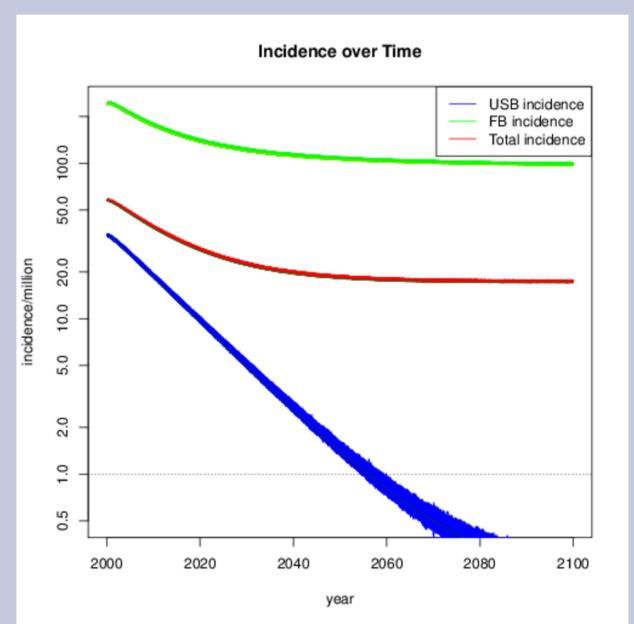
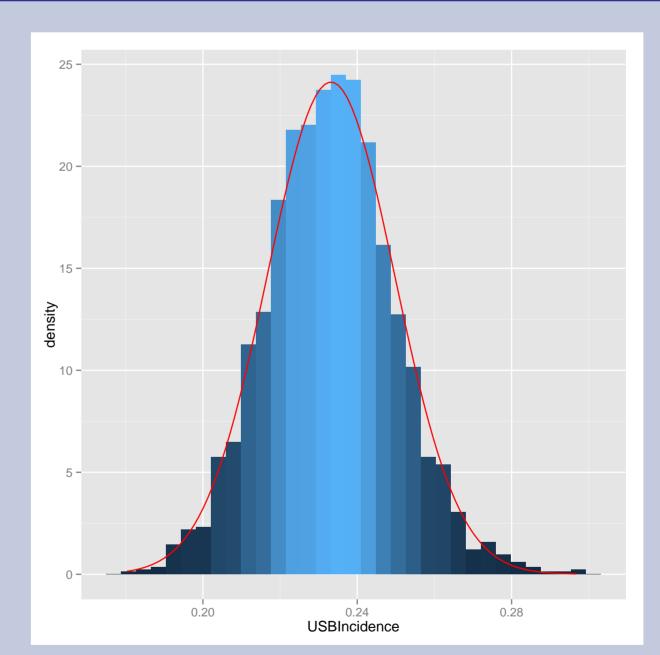
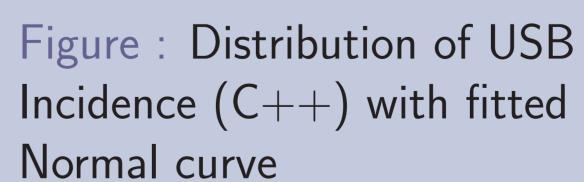


Figure : Incidence/million for R and NetLogo models (12 runs,  $\Delta t = 0.1$ , popConst = 100)

Figure : Incidence/million for R and C++ models (2100 runs,  $\Delta t = 0.01$ , popConst = 1)

### Stochastic Models as a Measure of Variability





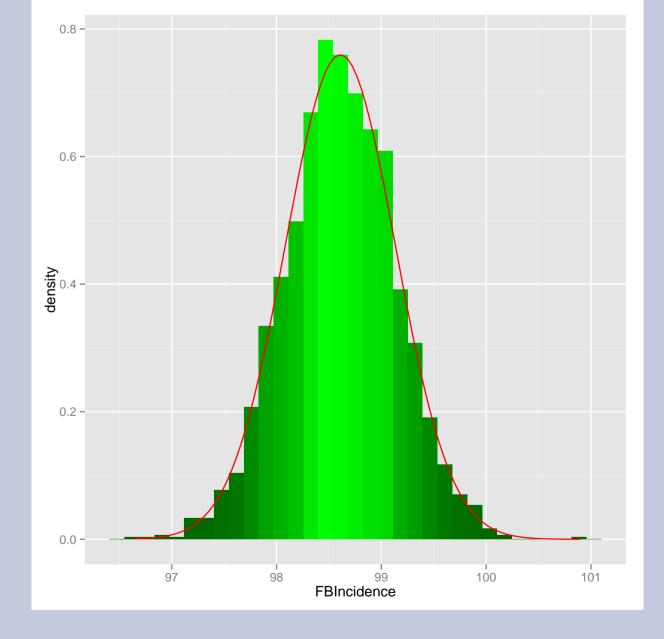


Figure: Distribution of FB Incidence (C++) with fitted Normal curve

#### References

Hill, A. N., Becerra, J. E., & Castro, K. G. (2012). Modelling tuberculosis trends in the USA. Epidemiology and infection, 140(10), 1862.