# 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

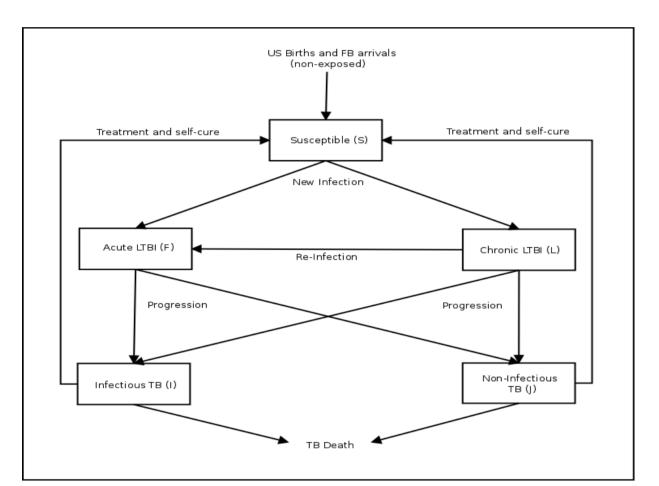
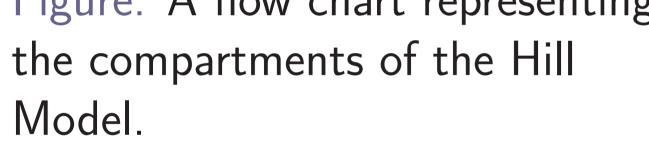


Figure: A flow chart representing the compartments of the Hill

# Populations:

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



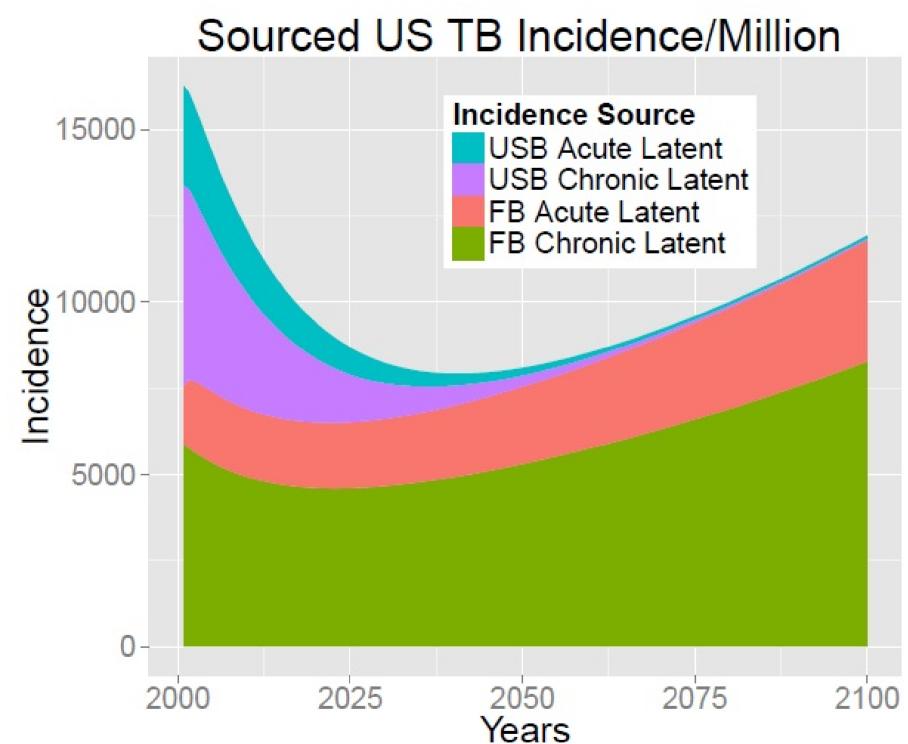


Figure: The source population of US TB incidence Sourced US TB HCS Cost USB Active TB due to Novel Infection USB Active TB due to Activation of LTBI FB Active TB due to Novel Infection FB Active TB due to Activation of LTBI

## **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**

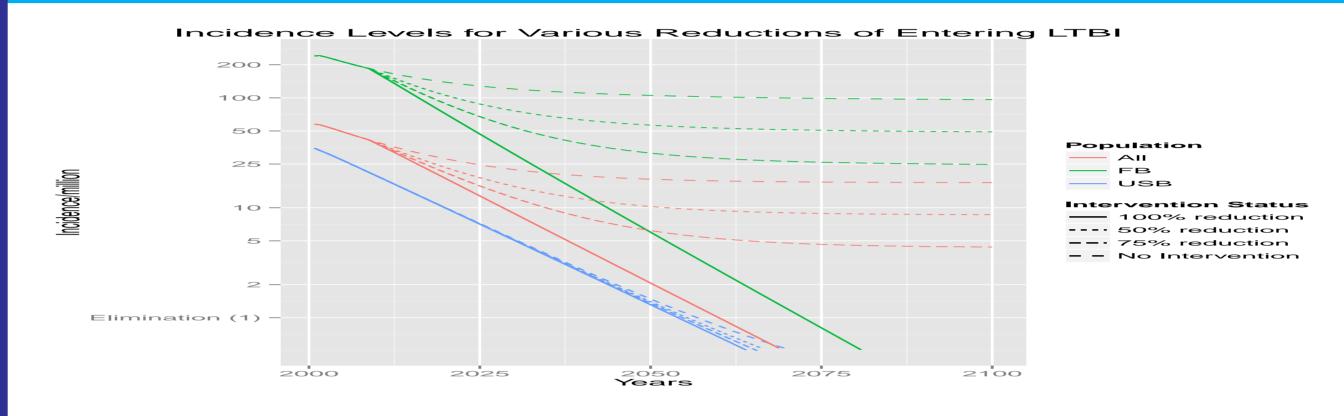


Figure: Incidence/million in USB, FB, and total populations, given 0%, 50%, 75%, or 100% treatment of incoming LTBI.

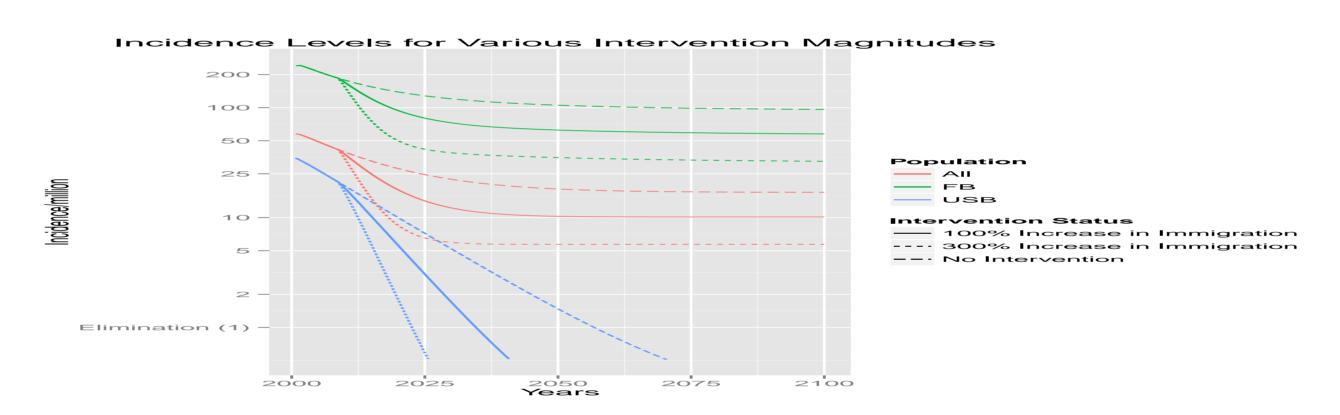


Figure: Incidence/million in USB, FB, and total populations, given 0%, 100%, or 300% LTBI treatment increase.

## **Economic Modeling**

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

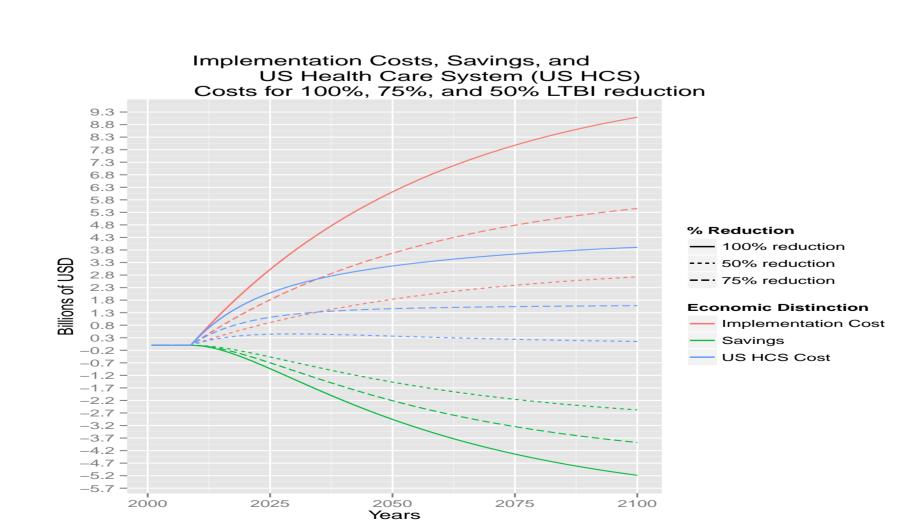


Figure: Cumulative implementation costs, US HCS savings, and net US costs of LTBI arrival cure rates. Cost/case cured was \$600, \$800, and \$1000 for 50%, 75%, and 100% cured.

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++.

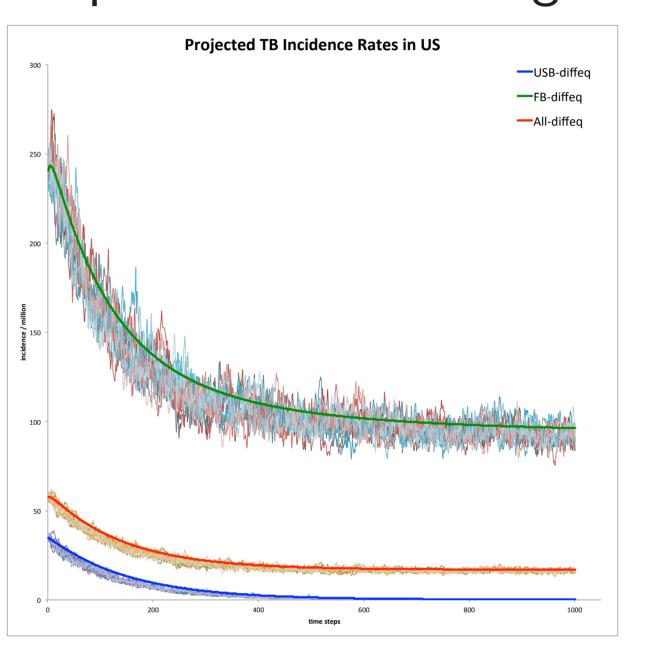


Figure: Incidence/million for R and NetLogo models (12 runs,  $\Delta t = 0.1$ , popConst = 100)

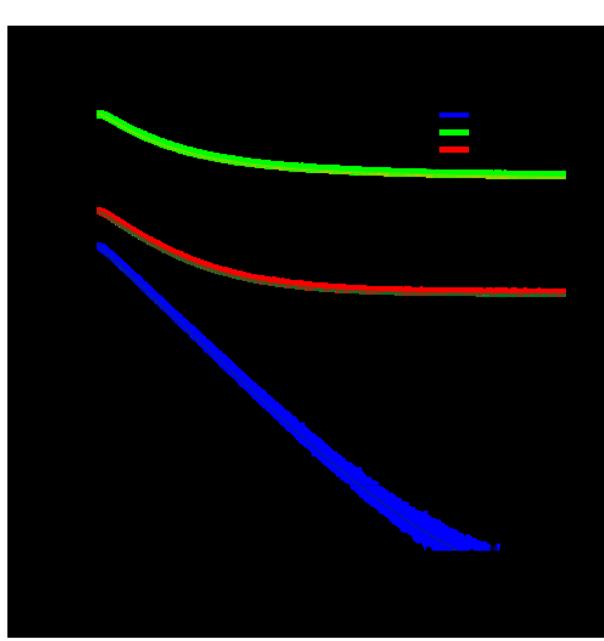


Figure: Incidence/million for R and C++ models (2100 runs,  $\Delta t = 0.01$ , popConst = 1)

## Stochastic Models as a Measure of Variability

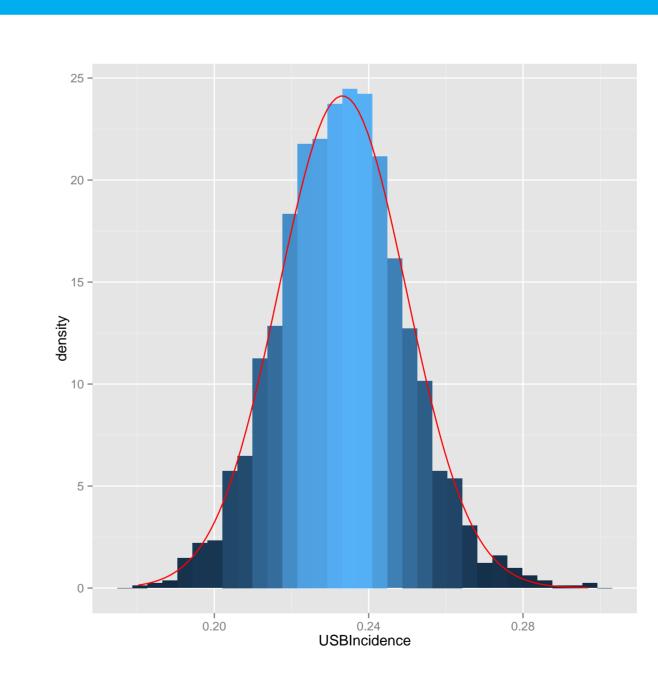


Figure: Distribution of USB Incidence (C++) with fitted Normal curve

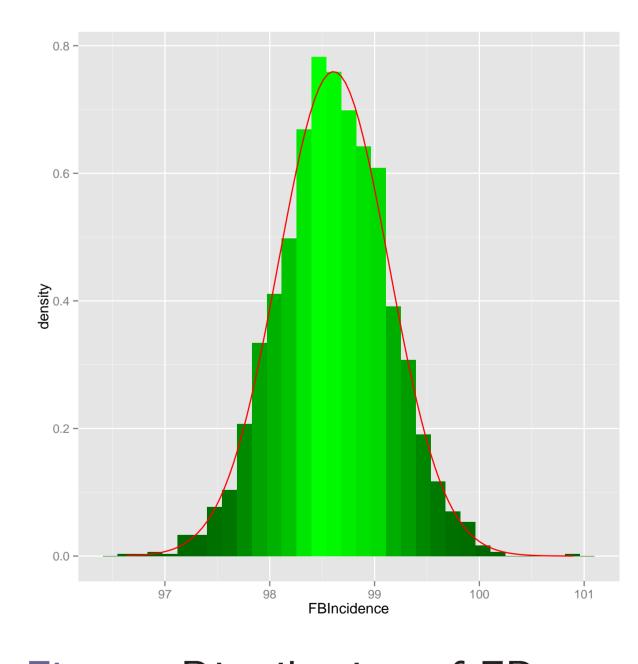


Figure: Distribution of FB Incidence (C++) with fitted Normal curve

#### References

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