

# Network-Informed Intervention Strategies for Pandemic Response

Lucas Laird



18 May, 2021





# Our Team



Tina  
Eliassi-Rad



Leo  
Torres



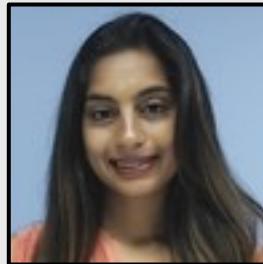
Jan-Willem  
van de Meent



Rajmonda  
Caceres



Lucas  
Laird



Neela  
Kaushik



Mykel  
Kochenderfer



Ross  
Alexander



Robin  
Walters



Niklas  
Smedemark-Margulies



Heiko  
Zimmermann



Chris  
Vanderloo



Peter  
Morales



Shushman  
Choudhury



Kunal  
Menda



Northeastern  
University



LINCOLN LABORATORY  
MASSACHUSETTS INSTITUTE OF TECHNOLOGY

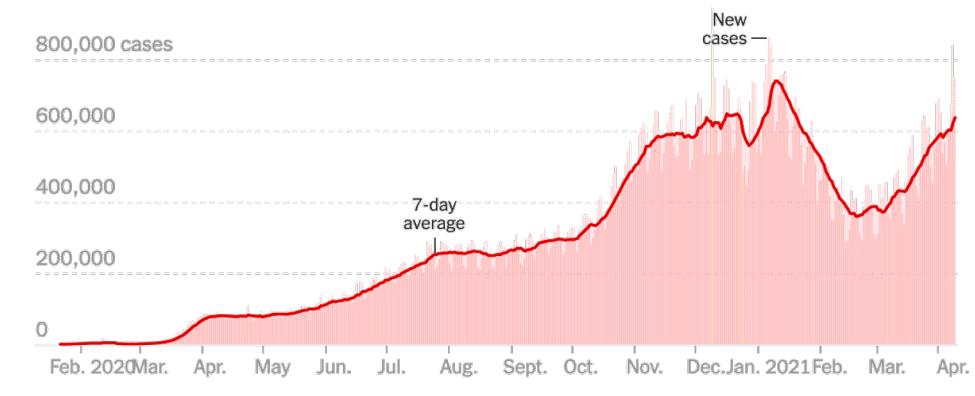
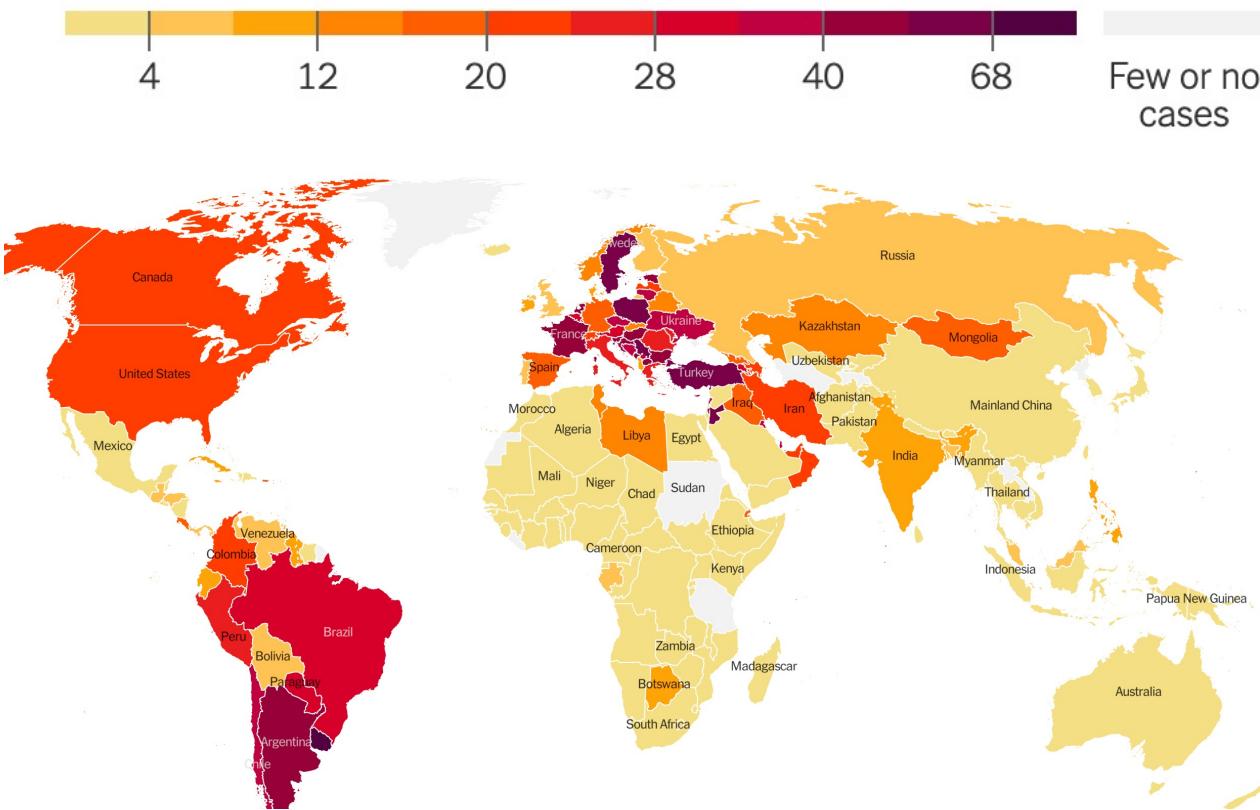


Stanford  
University



# COVID-19 Pandemic

Average daily cases per 100,000 people in past week



	TOTAL REPORTED	ON APRIL 9	14-DAY CHANGE
Cases	134.6 million+	749,008	+19% →
Deaths	2.9 million+	13,498	+19% →

**COVID-19 is causing devastating human and economic losses at global scale**

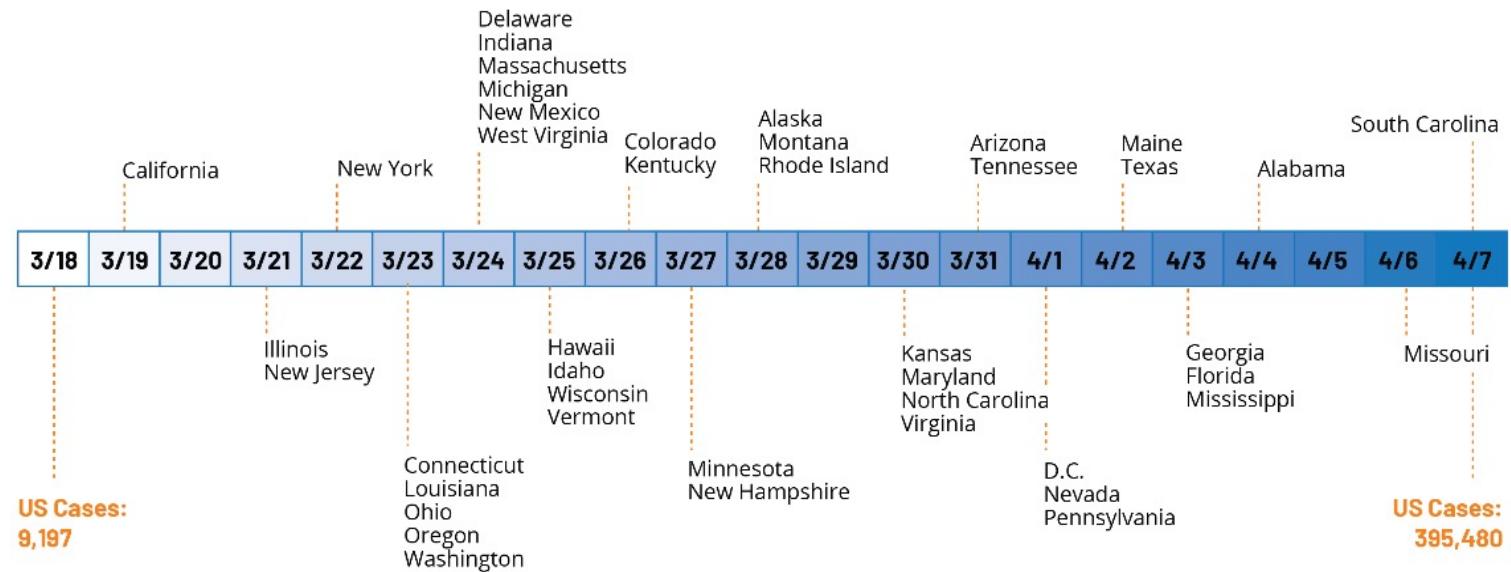


# Why is Control of Pandemics Difficult?

**COVID revealed that we don't have good NPI strategies**

- Uncoordinated state responses
- Severe, one-size-fits-all strategies do not respect discrepancies in local resources
- Responses do not adapt to changing conditions (new strains, new outbreaks, vaccination)

## Timeline of state stay-at-home orders

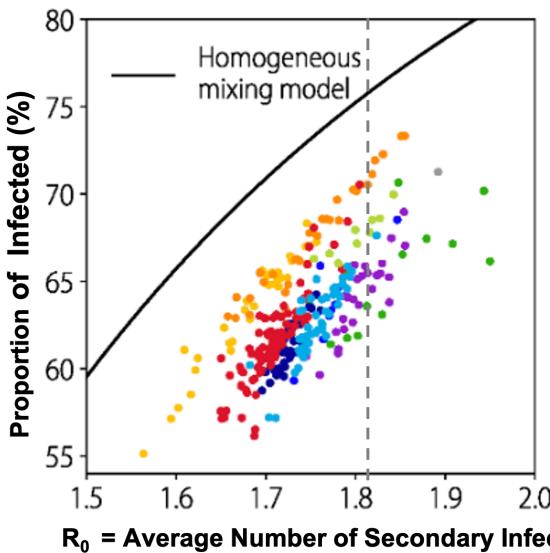


**We lack systems that identify contextual, effective, lower-cost pandemic mitigation strategies**



# Importance of Complex Network Modeling

## Complex Network Effects



Same local effects lead to heterogeneous meso-scale and global scale effects

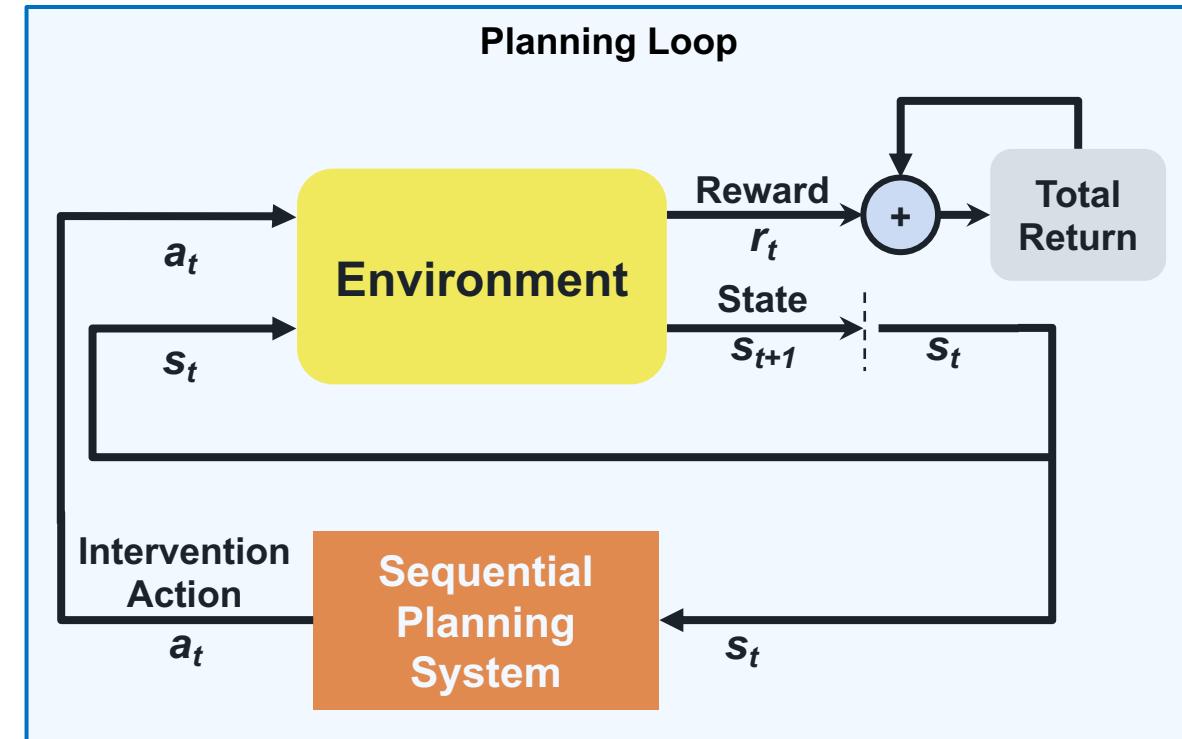
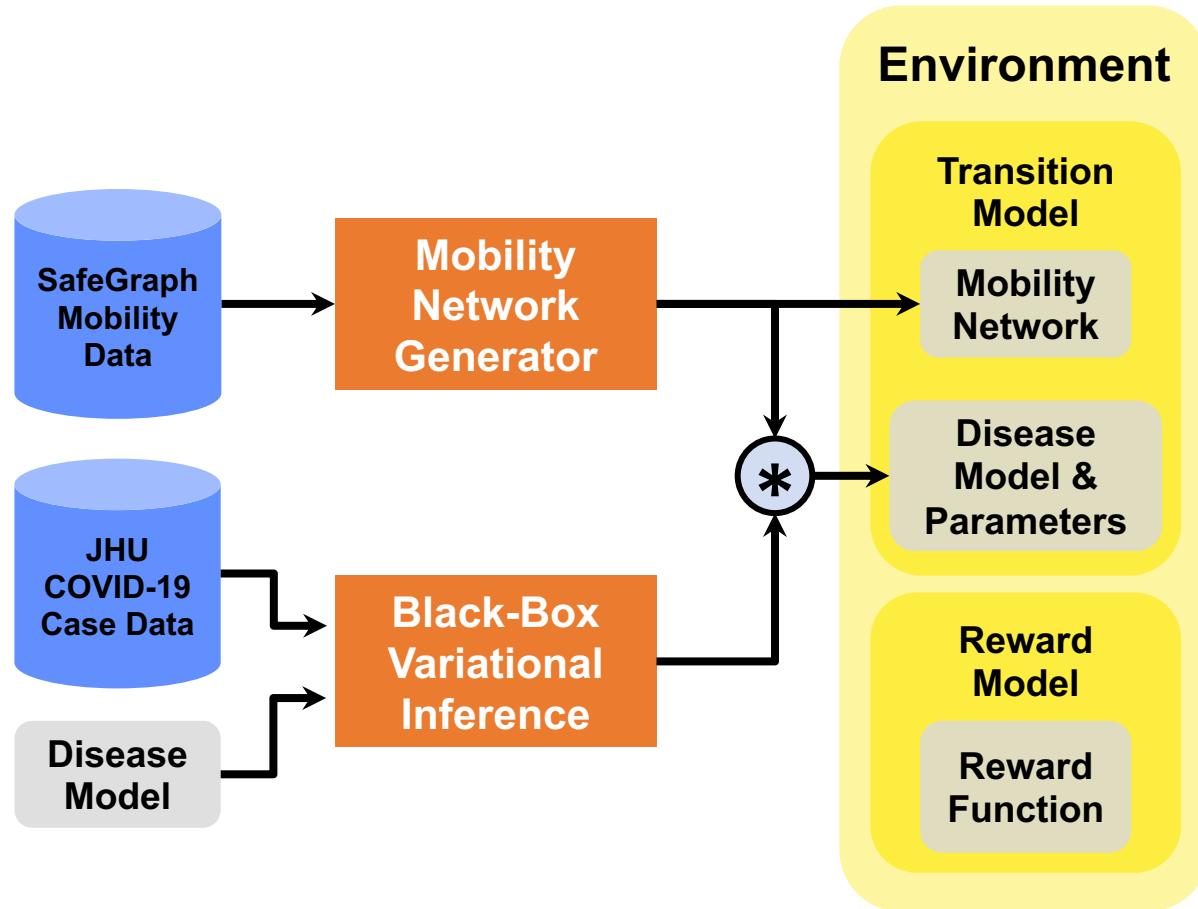
## Incorporating network effects is crucial for realistic modeling of diseases

- Models that ignore underlying networks produce inaccurate predictions leading to poor evaluation and public distrust
- Evaluating intervention strategies with inaccurate models is insufficient at best, dangerous at worst
- Realistic simulations allow adaptively finding new strategies through simulated trial and error

High-resolution models are necessary for realistic simulation and evaluating intervention efficacy



# Courses of Actions over Incomplete Networks (COANET)





# Outline



- Network SEIR model
- Sequential Planning Algorithm
- Results and Discussion

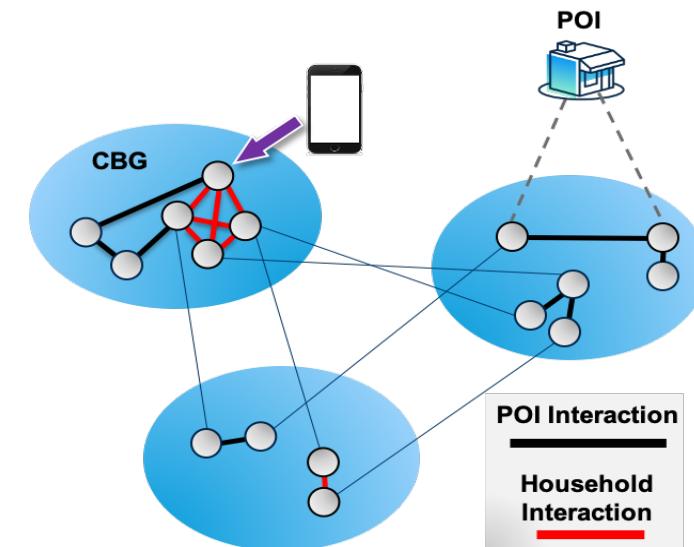
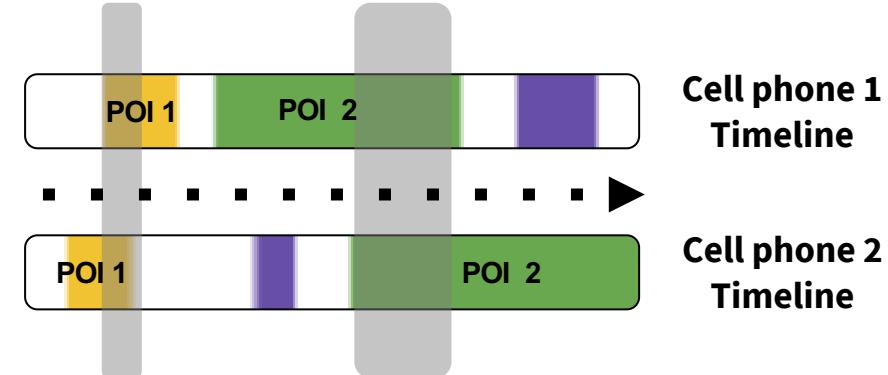


# Contact Networks as Stochastic Block Models



SAFEGRAPH

- Device to points-of-interests (POI) location data from opt-in smartphone devices<sup>1</sup>
- Each smartphone device is mapped to a “home” Census Block Group (CBG)
- Leverage a DCSBM<sup>2</sup> network model to simulate geographically structured contact patterns
  - Nodes are individuals; stochastic blocks are CBGs
  - Edges represent different types interactions (household, POI co-location)
  - POI interaction probability is based on shared (correlation) POI visit counts, household interaction probability is set to 1
  - Weight of interactions captures the duration of co-location

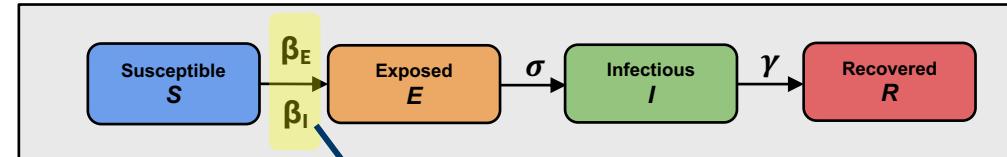




# Incorporating Networks into SEIR Modeling

## Compartmental Models:

$$\frac{dS}{dt} = -\frac{\beta SI}{N} \quad \frac{dE}{dt} = \frac{\beta SI}{N} - \sigma E$$



## Network-based Model: Network-SEIR

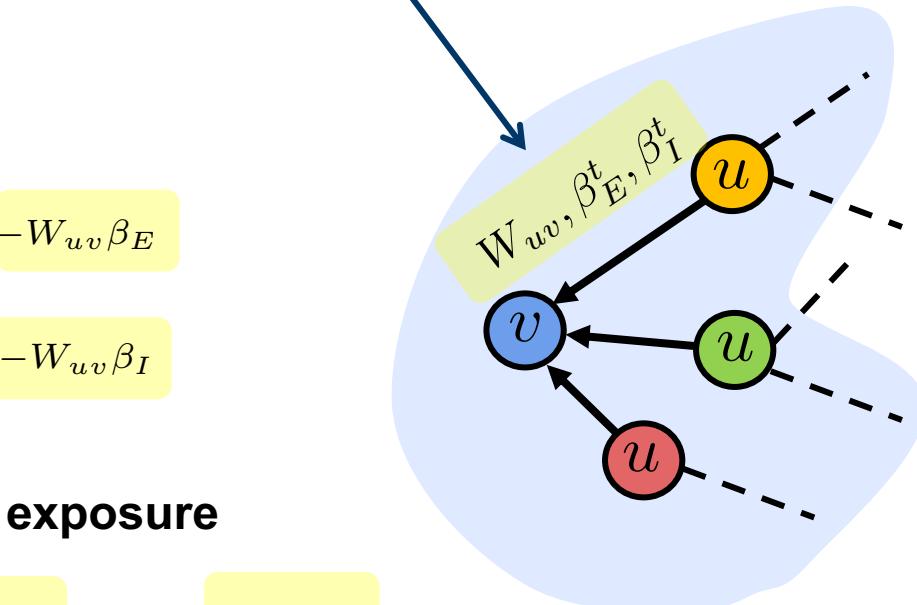
- Exponential model of node interactions

$$P(v \text{ becomes exposed} \mid u \text{ is exposed}) = 1 - e^{-W_{uv}\beta_E}$$

$$P(v \text{ becomes exposed} \mid u \text{ is infected}) = 1 - e^{-W_{uv}\beta_I}$$

- Local neighborhood determines probability of exposure

$$P(v \text{ becomes exposed}) = 1 - e^{-\sum_{u \in N_E} W_{uv}\beta_E - \sum_{u \in N_I} W_{uv}\beta_I}$$

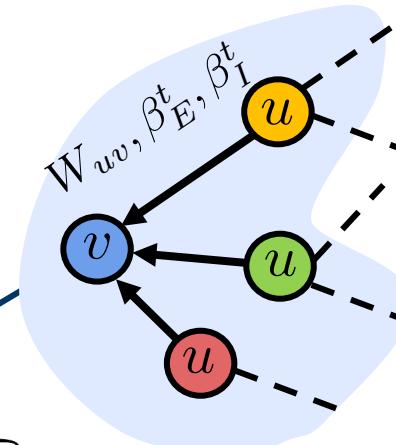
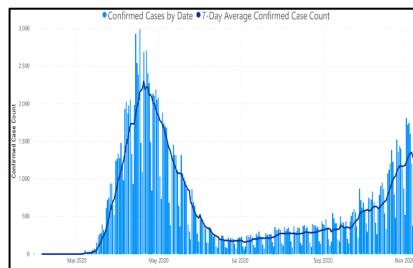




# Estimating GraphSEIR Parameters from Data

Fit network-level parameters directly to data using probabilistic programming

JHU Reported Infection Counts



$$p(\beta \mid \text{data}) = \frac{\overbrace{p(\text{data} \mid \beta) p(\beta)}^{}_{}}{p(\text{data})}$$

Black-box  
Variational  
Inference  
(BBVI)

$$\begin{aligned}\phi^* &= \arg \min_{\phi} \text{KL}(q_{\phi}(\cdot) \mid \underbrace{| p(\cdot \mid \text{data})}_{\text{intractable}}) \\ &= \arg \max_{\phi} \mathcal{L}(\phi) \leftarrow \text{tractable surrogate objective (ELBO)}\end{aligned}$$



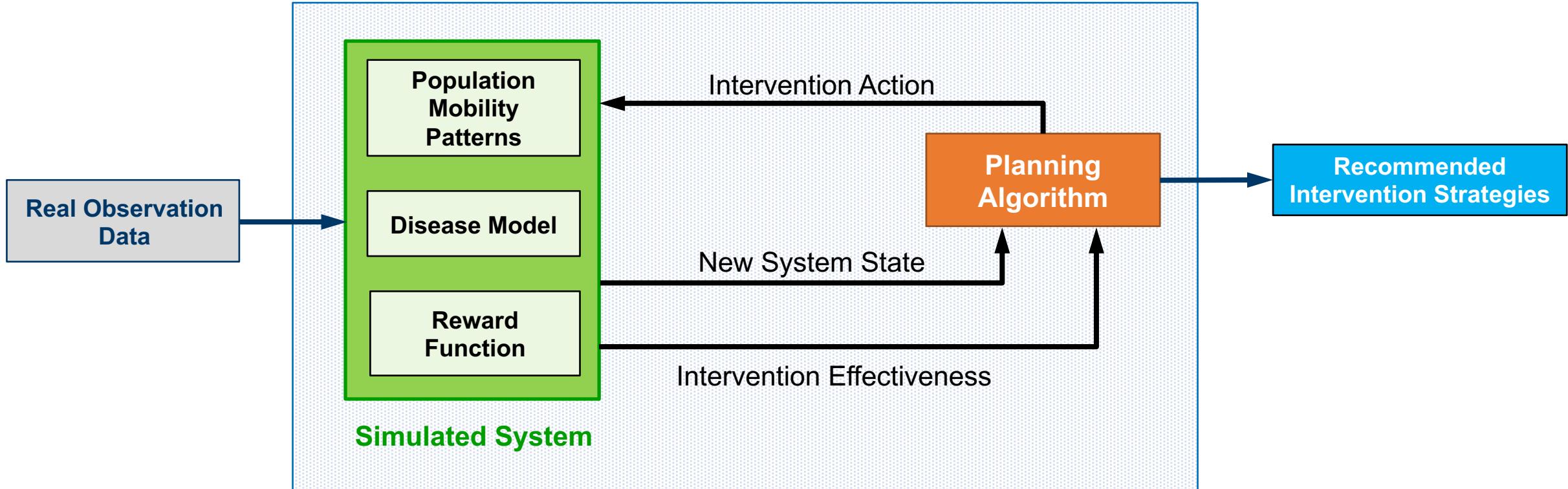
# Outline

- Network SEIR model
- Sequential Planning Algorithm
- Results and Discussion





# Pandemic Control via Reinforcement Learning

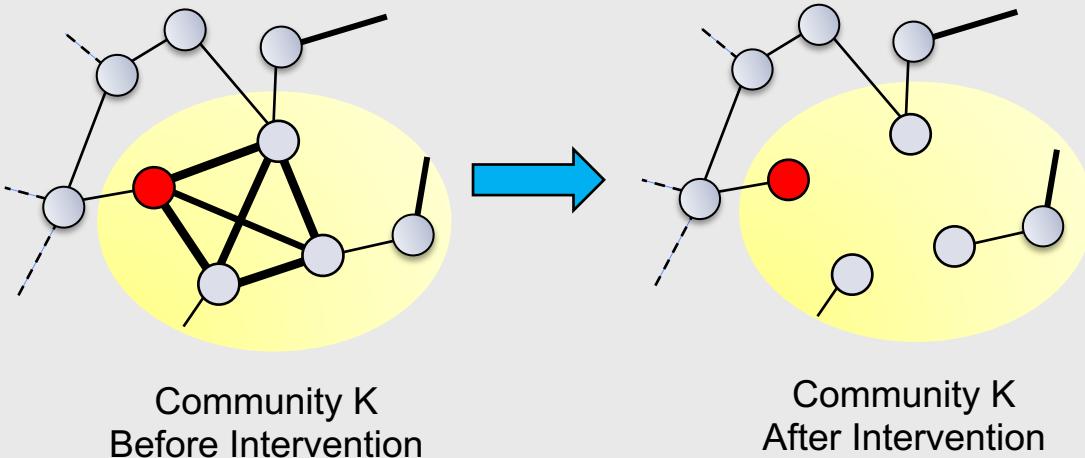


## COANET System Overview



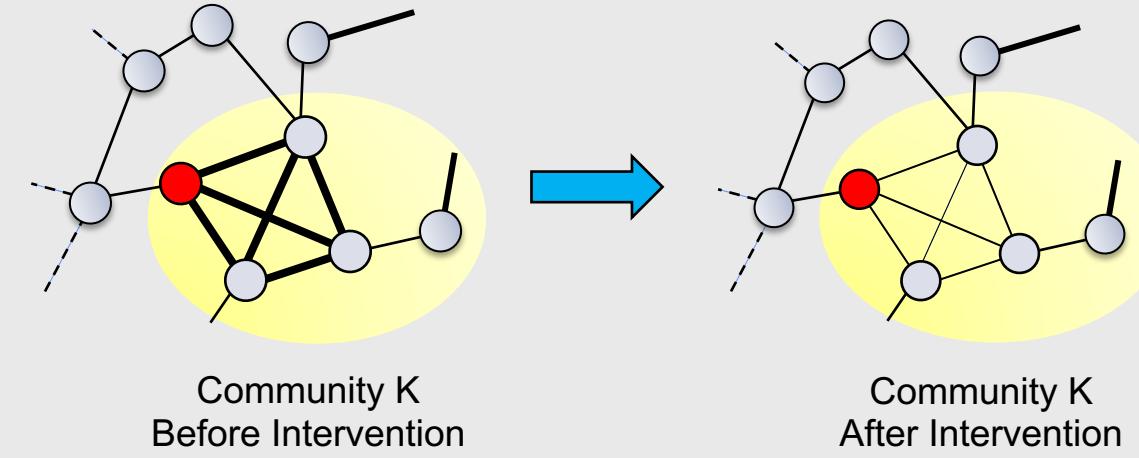
# Action Space

Action 1: Enforce stay-at-home action in community K with a given intensity strength



**Network Sparsification**

Action 2: Enforce mask-wearing in community K with a given effectiveness strength



**Reduction of disease spread**

- **Jitter Suppression:** Actions must have a minimum active duration
- **Prioritization:** Limit actions to top ranked communities by network centrality and disease prevalence
- **Feasibility:** Allow only one, single community action per timestep



# Reward Function

- **Reward function intuitions**

- Maximizing susceptible individuals rewards limiting disease spread
- Policy scale represents relative disruptiveness of intervention
- Total policy cost is proportional to number of impacted individuals
- Policy change cost penalizes changes in interventions to capture public “policy fatigue”

$$r = \frac{1}{N} \left( S(G) - \sum_{i=1}^{|C|} (\alpha_{sh} s_{sh,i} |c_i| + \alpha_{mw} s_{mw,i} |c_i|) + \alpha_{a \neq 0} |c_i| \right)$$



$N$ : Population Size

$c_i$ : Community being intervened

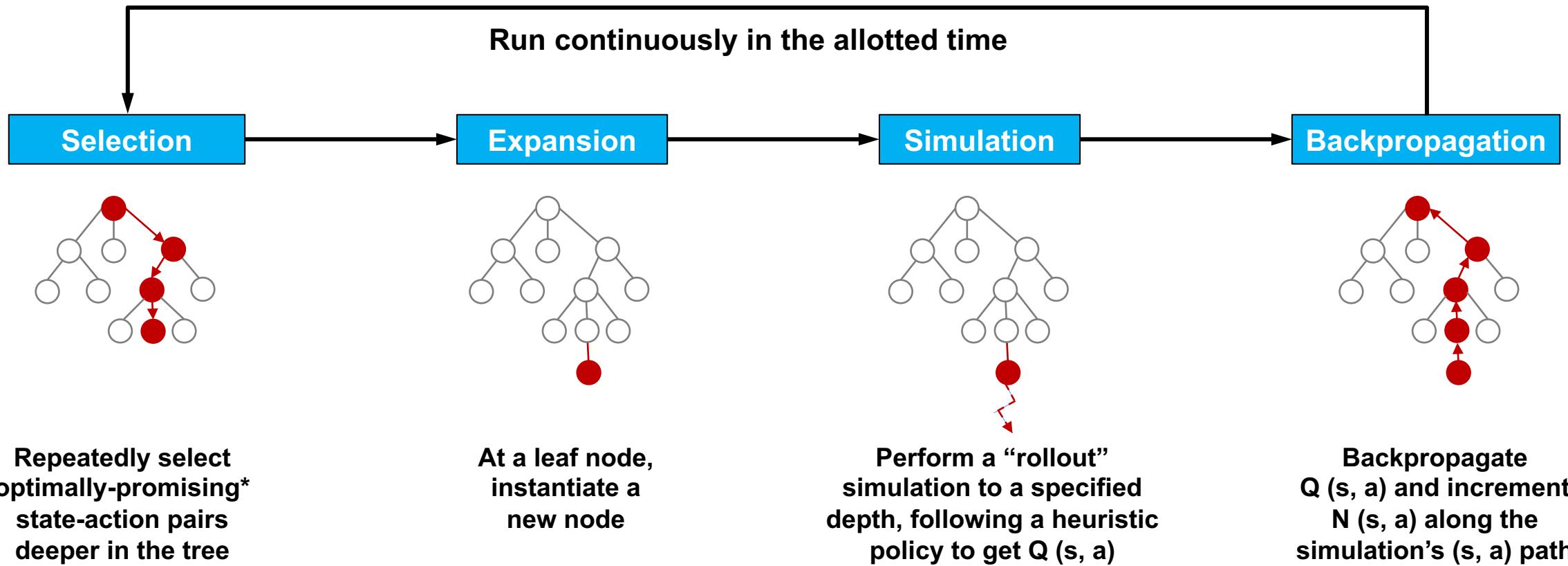


# Monte Carlo Tree Search (MCTS) Algorithm

## State-action tree with

$Q(s, a)$  - the estimated expected return of taking action  $a$  from state  $s$

$N(s, a)$  - how often we have visited state-action pairs in the tree





# Outline

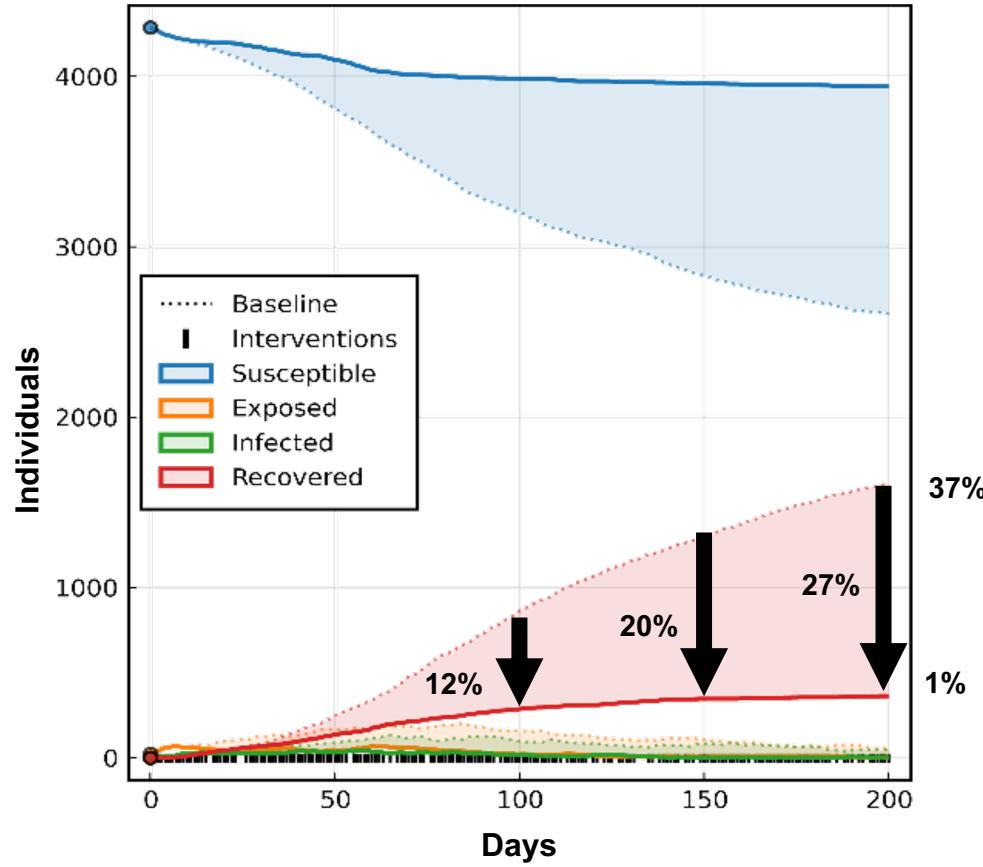
- Network SEIR model
- Sequential Planning Algorithm
- Results and Discussion





# Controllability of COVID-19

## Uniform initial infection spread

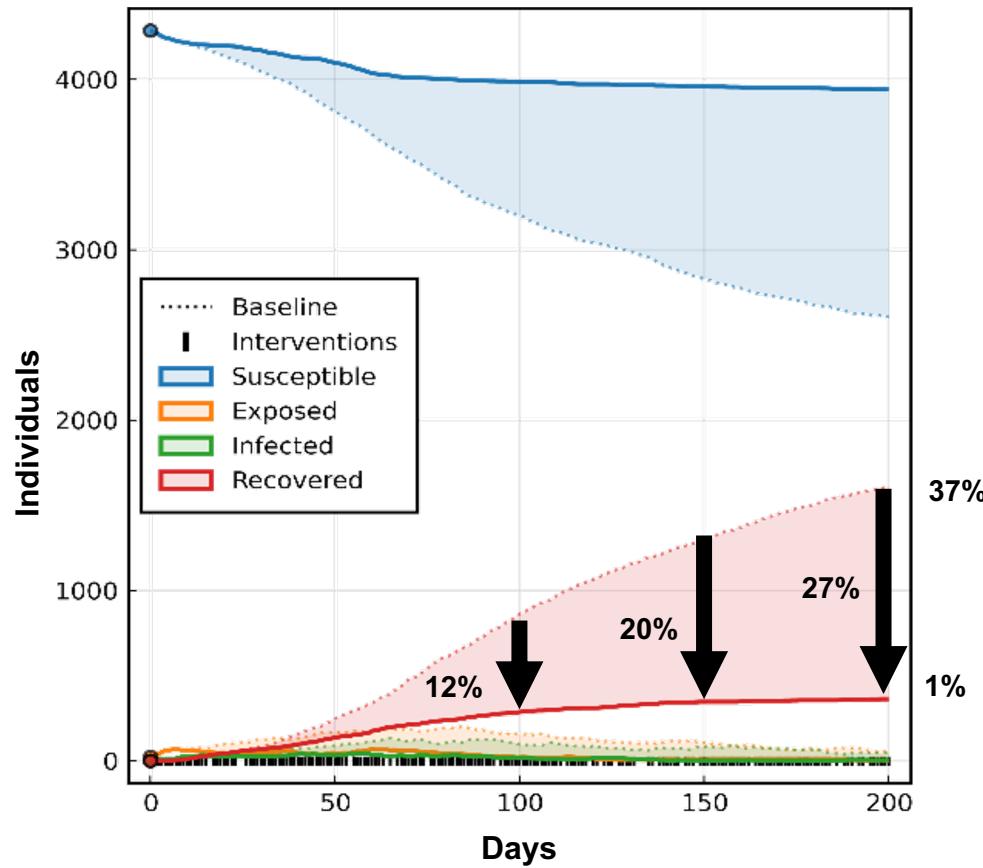


Planning leads to more effective and targeted control

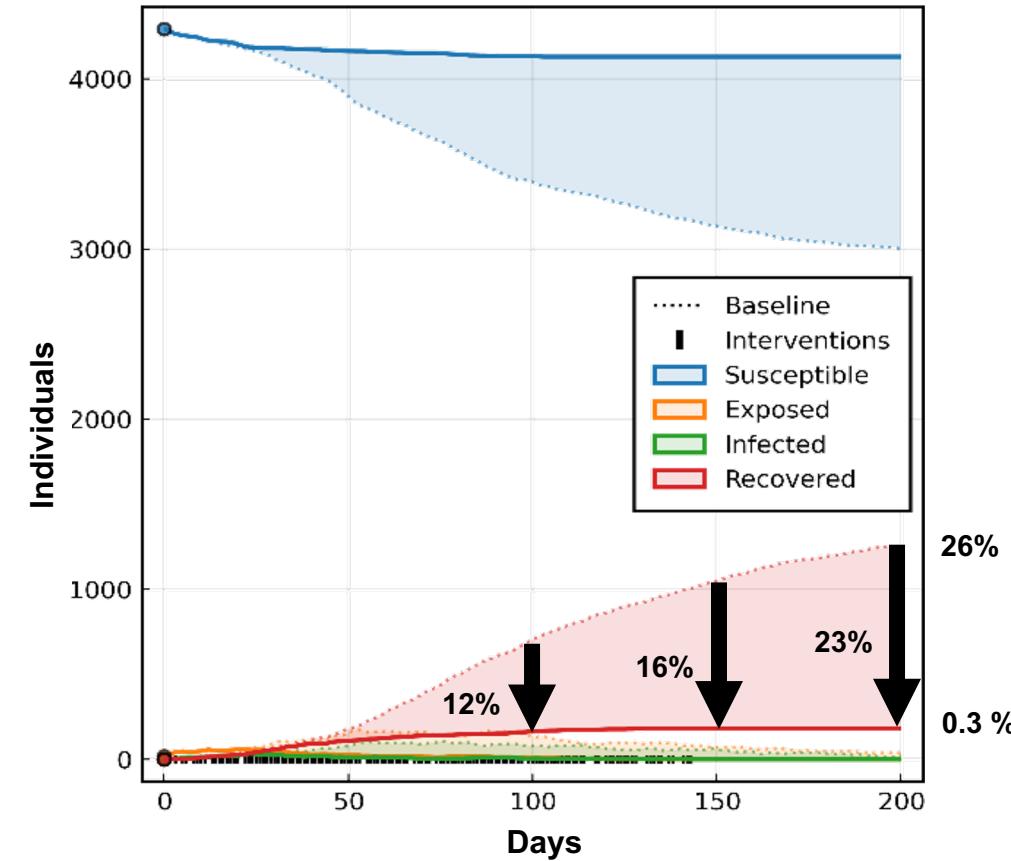


# Controllability of COVID-19

Uniform initial infection spread



Initial infection spread in 2 census blocks

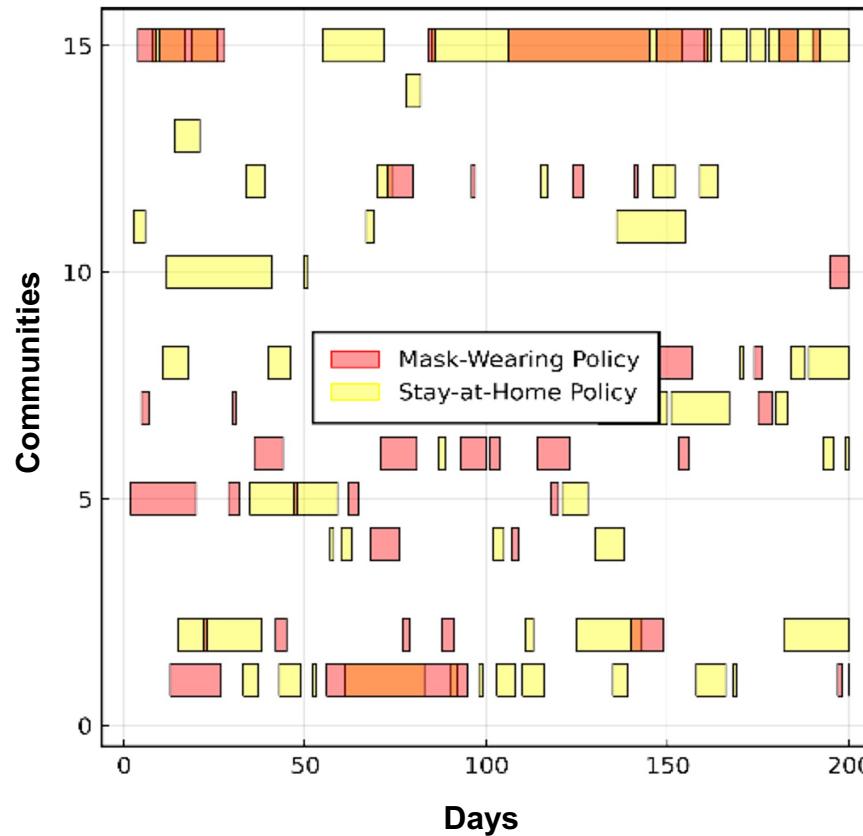


The more localized the initial conditions the more effective the control

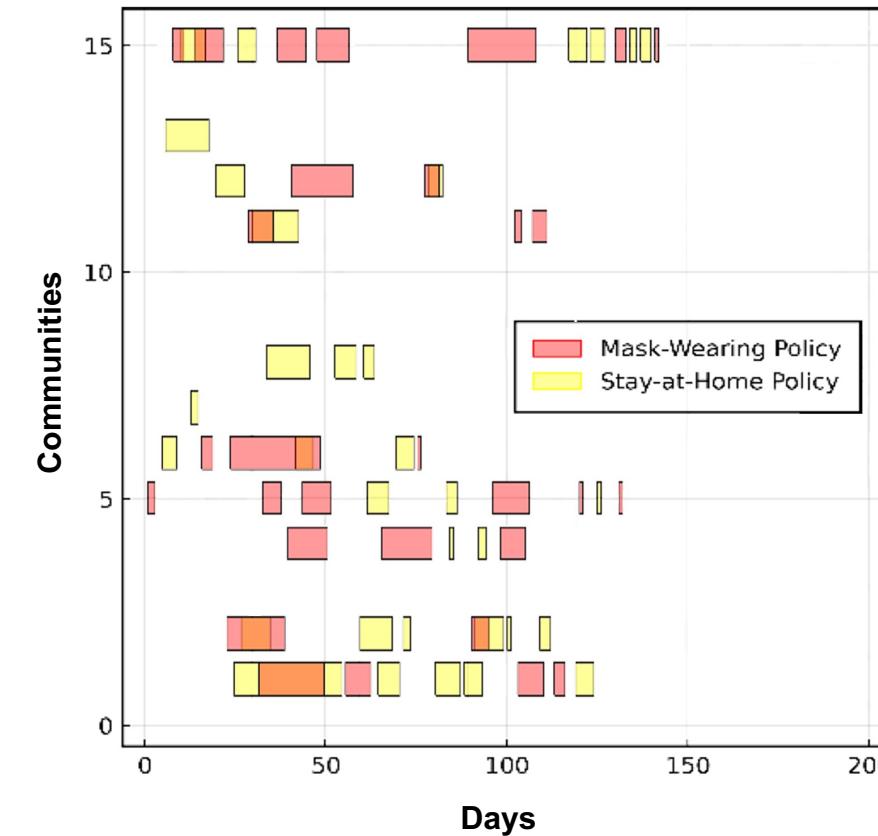


# Controllability of COVID-19

Uniform initial infection spread



Initial infection spread in 2 census blocks

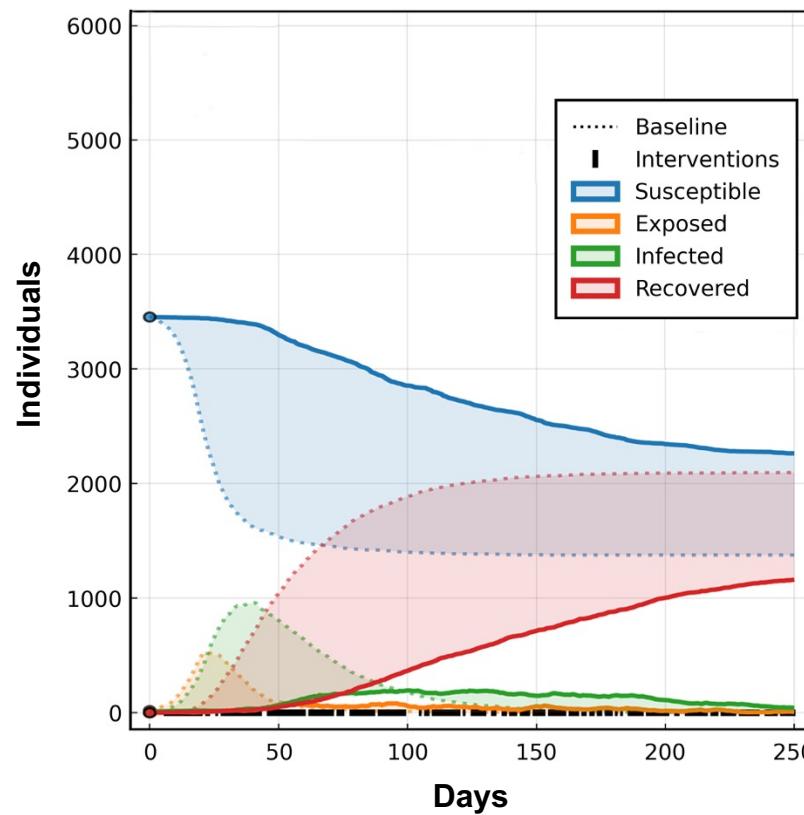


Adaptive strategies are non-trivial – COANET finds no universal lockdown required

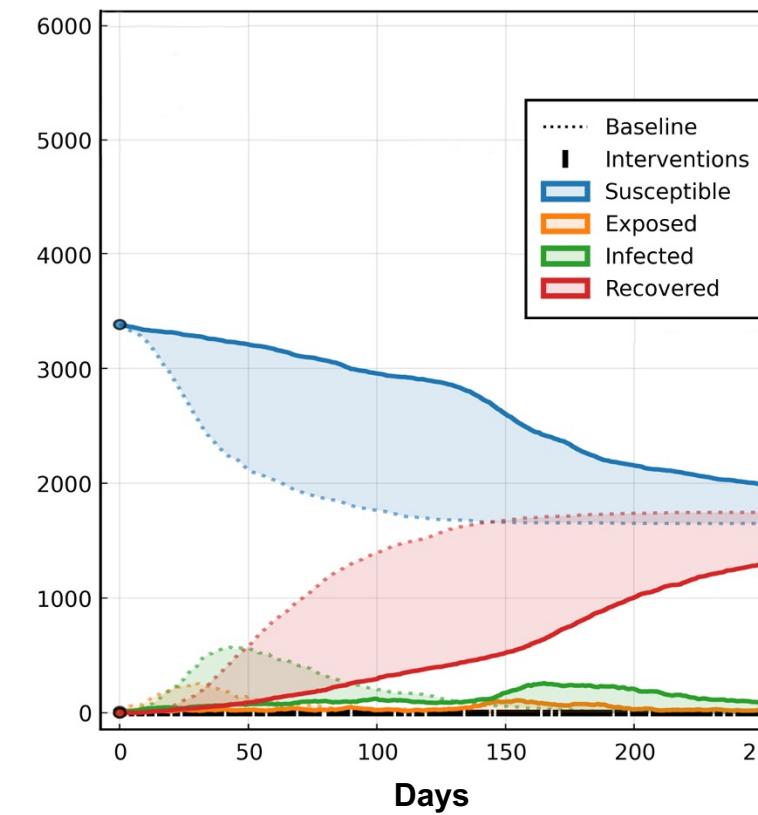


# Control Effectiveness in Different Geographic Regions

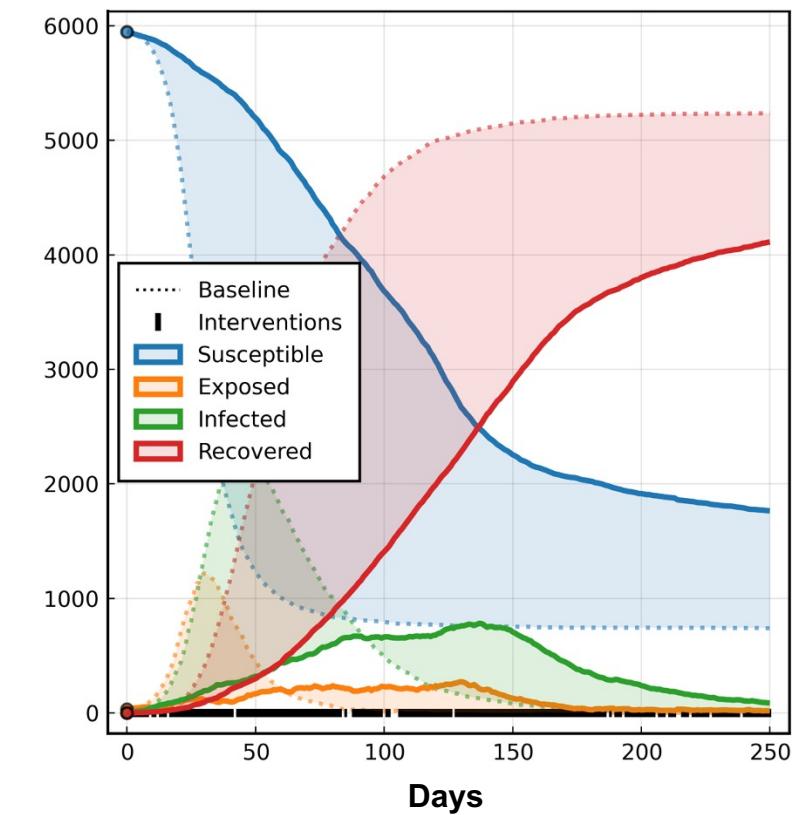
Miami-Dade County, FL



Middlesex County, MA



Los Angeles County, CA

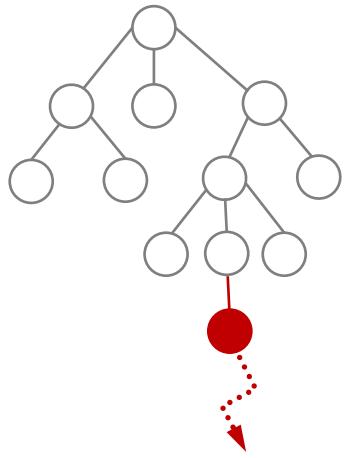


Our system customizes control strategies to different geographic regions and maintain effectiveness



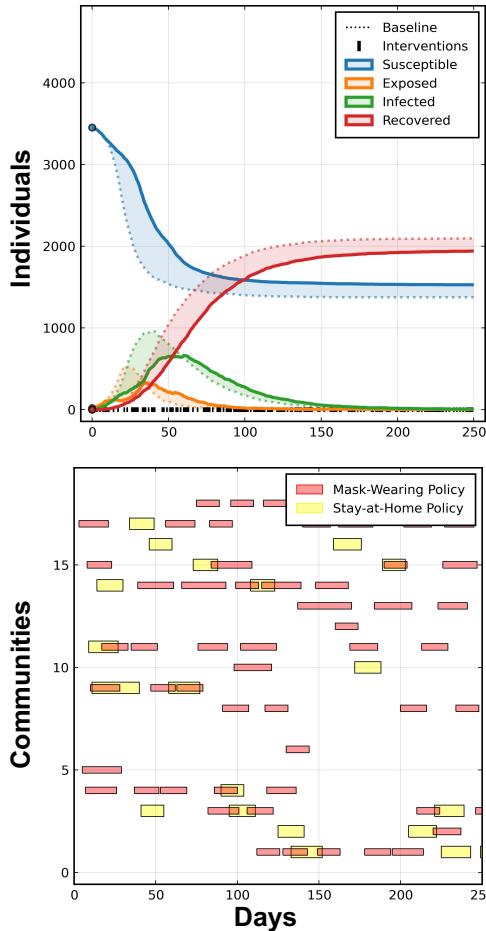
# Effects of Longer Lookaheads

## Simulation



**Actions are evaluated by simulating disease forward in time**

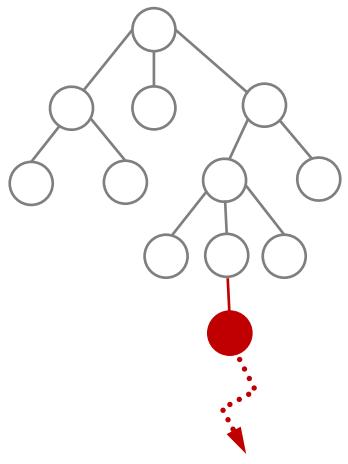
2 week lookahead





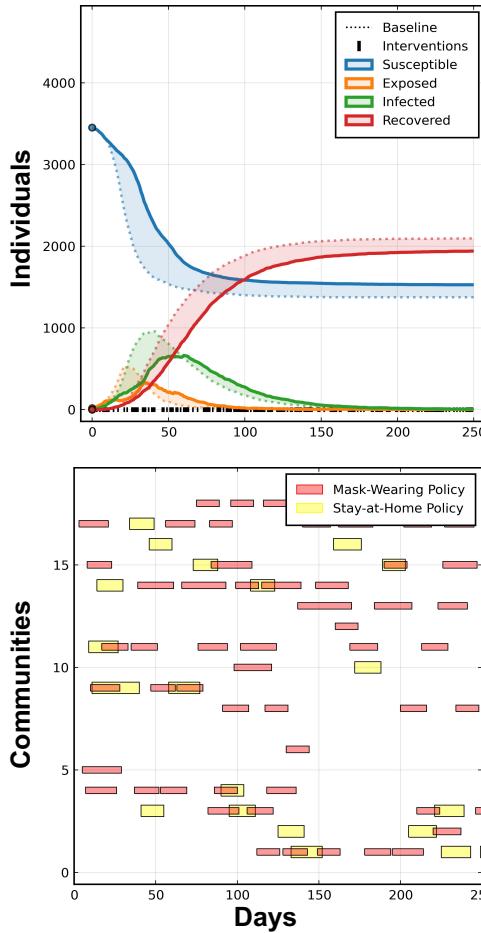
# Effects of Longer Lookaheads

## Simulation

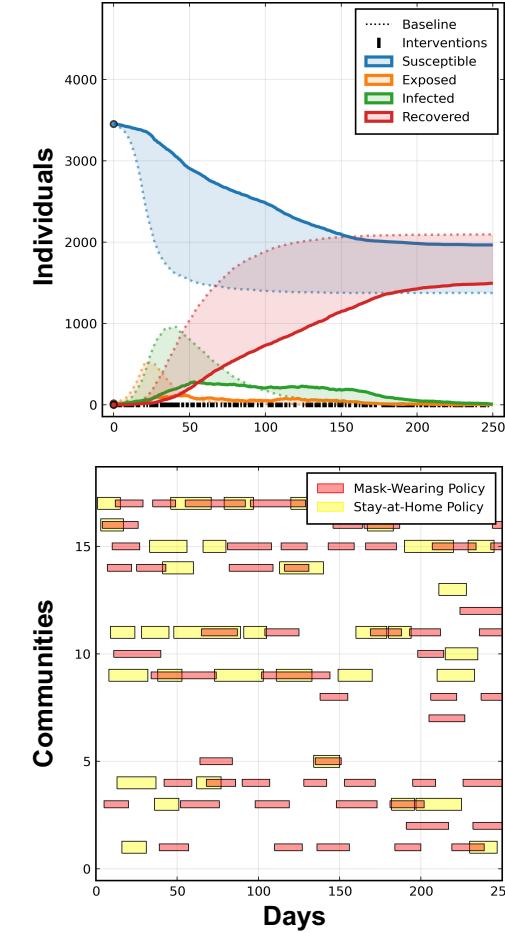


**Actions are evaluated by simulating disease forward in time**

2 week lookahead



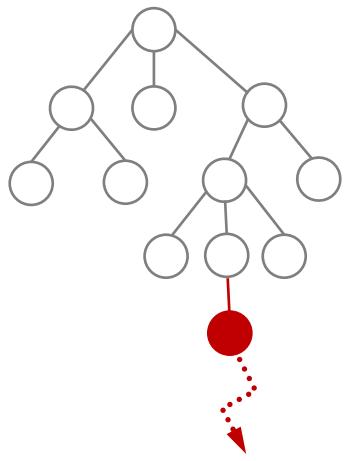
4 week lookahead





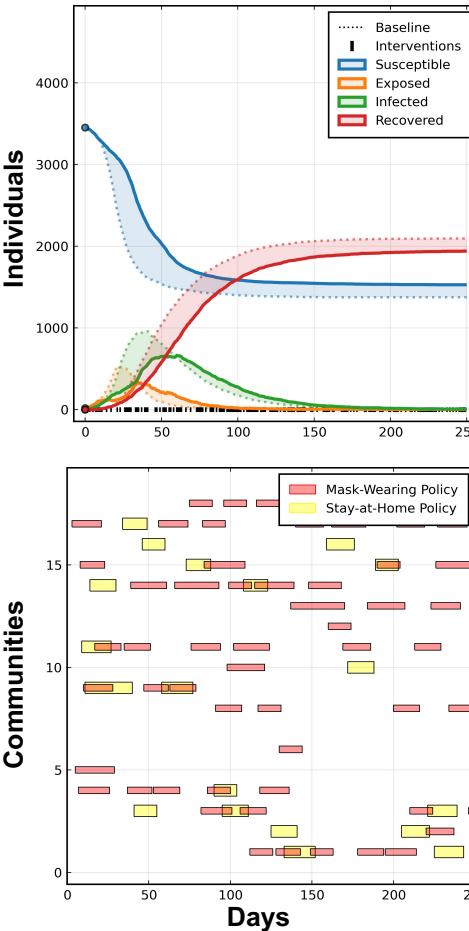
# Effects of Longer Lookaheads

## Simulation

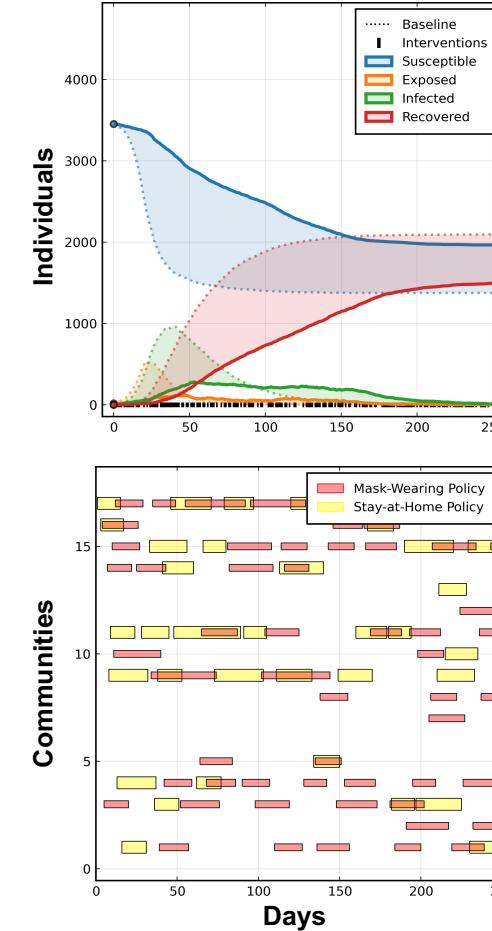


**Actions are evaluated by simulating disease forward in time**

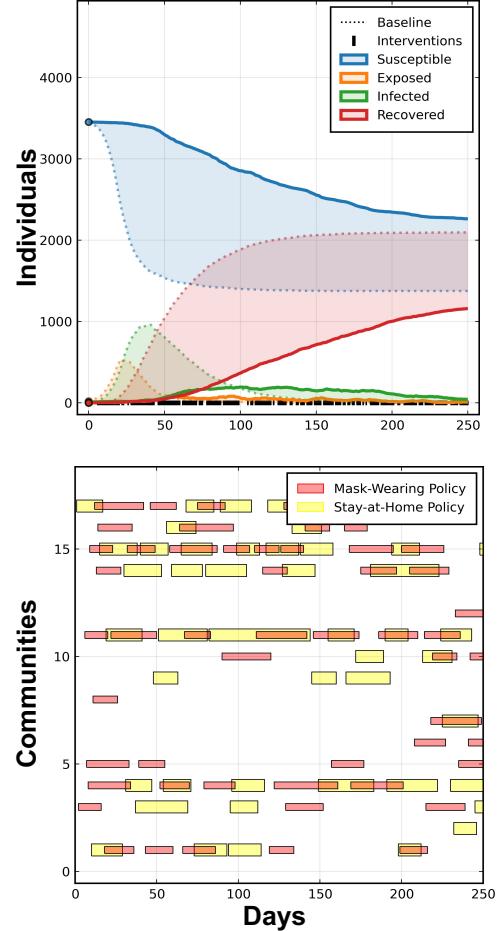
**2 week lookahead**



**4 week lookahead**



**6 week lookahead**

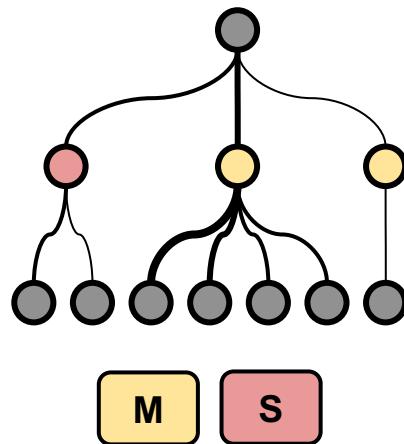


**COANET identifies a minimum, required 3-4 week lookahead time for effective planning**



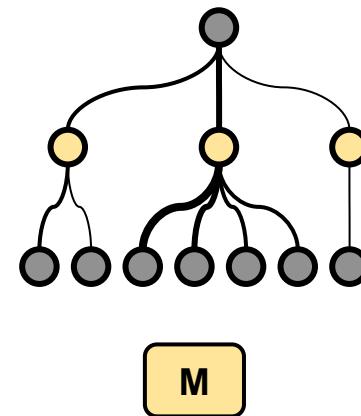
# Comparison With Baseline Approaches

MCTS Planner with  
Mask-Wearing and  
Stay-at-Home Actions

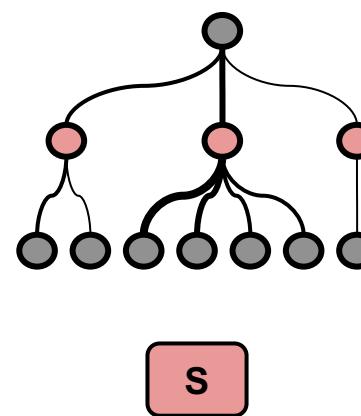


Our Approach

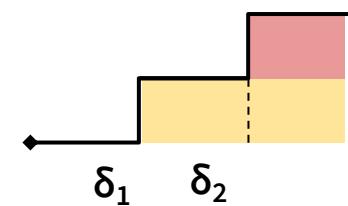
MCTS Planner with  
Mask-Wearing  
Actions Only



MCTS Planner with  
Stay-at-Home  
Actions



Global  
Threshold  
Policy



Random  
Policy

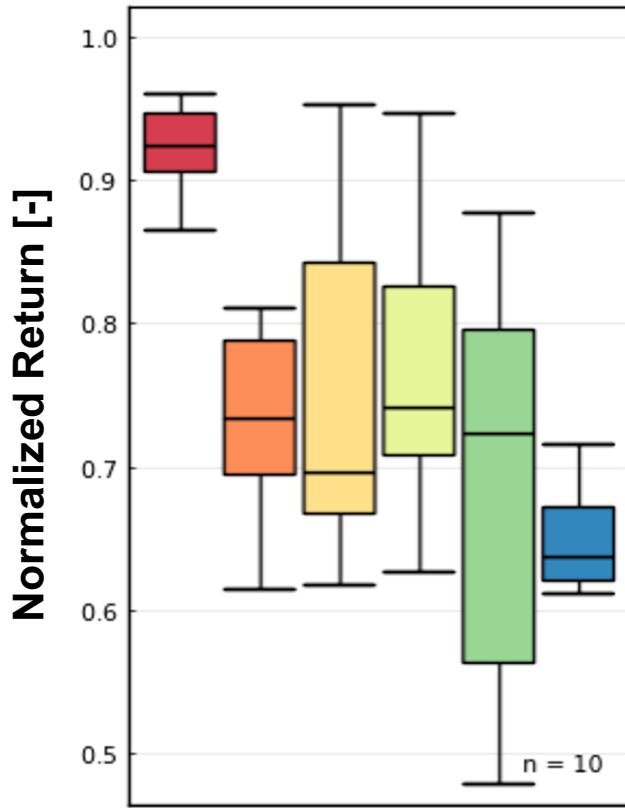


Evaluate the performance of our adaptive planner against several baseline heuristics

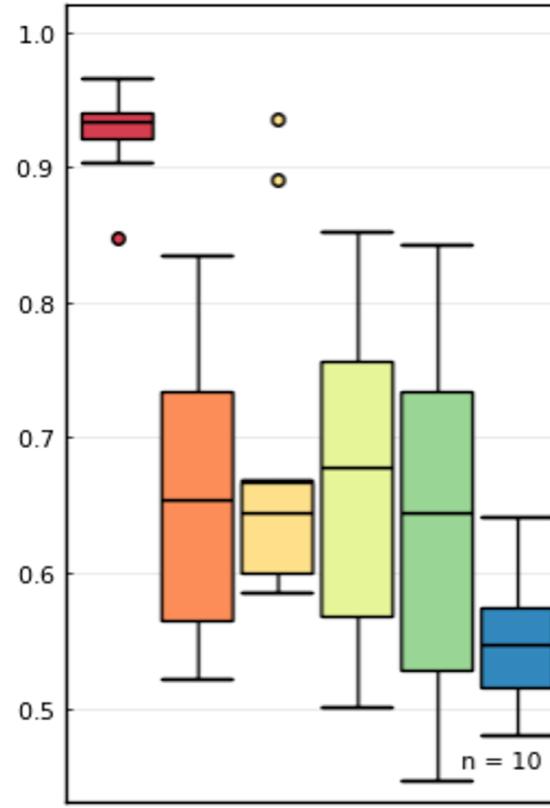


# Performance Comparison With Baselines

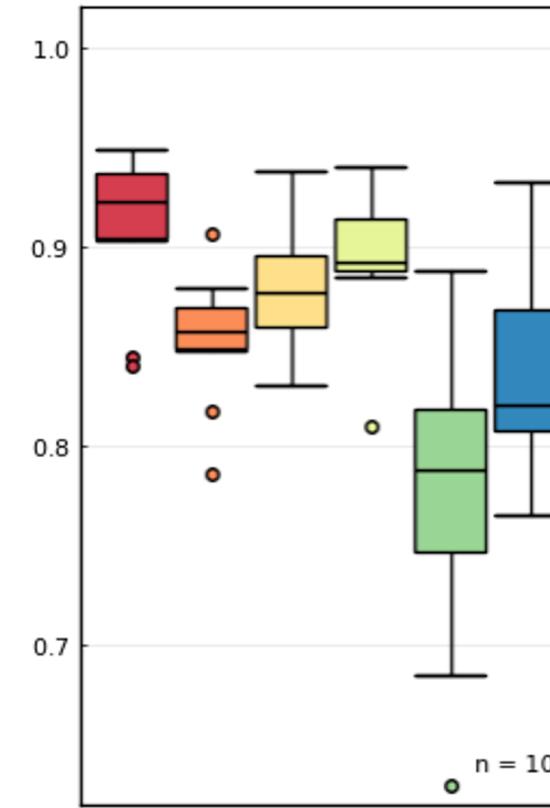
Miami-Dade County, FL



Middlesex County, MA



Los Angeles County, CA

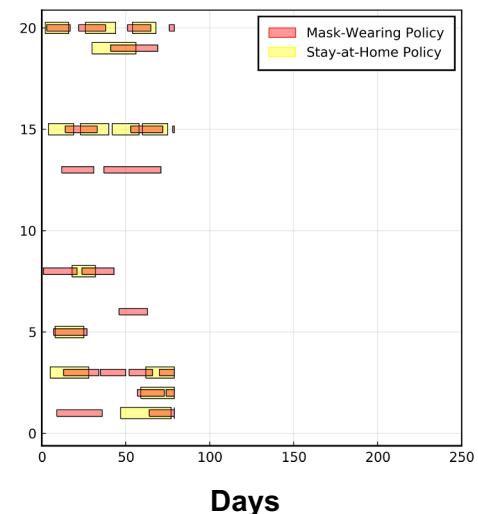


Our planner achieves 1) higher cumulative return and 2) lower variance

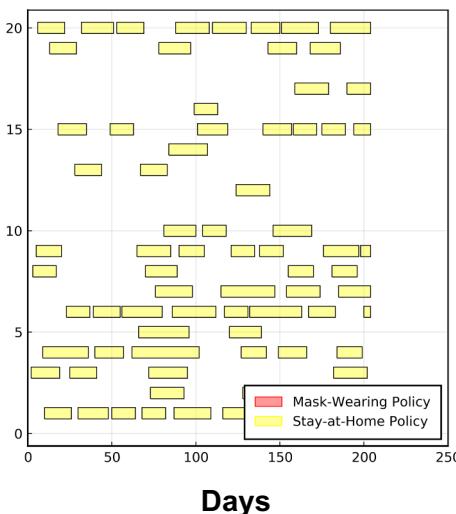


# Middlesex County, MA Results

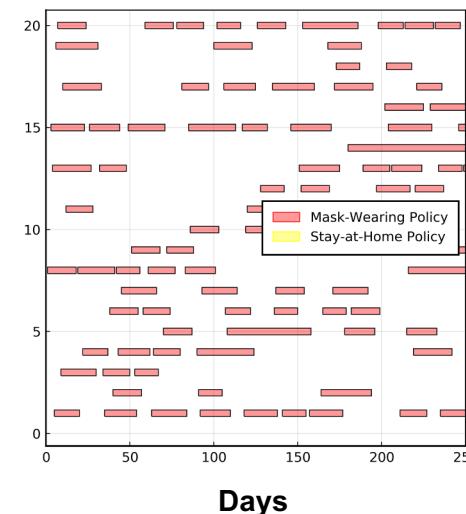
## Our Approach



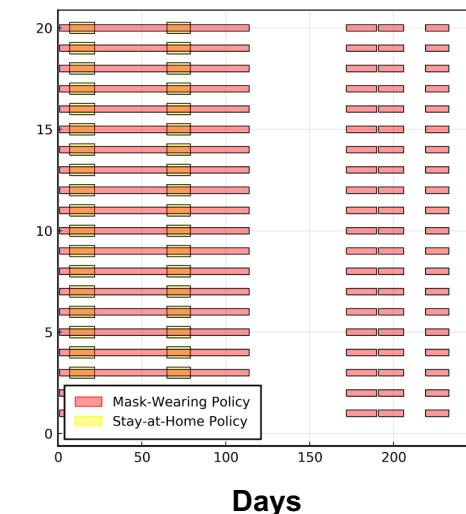
## Stay home-Only



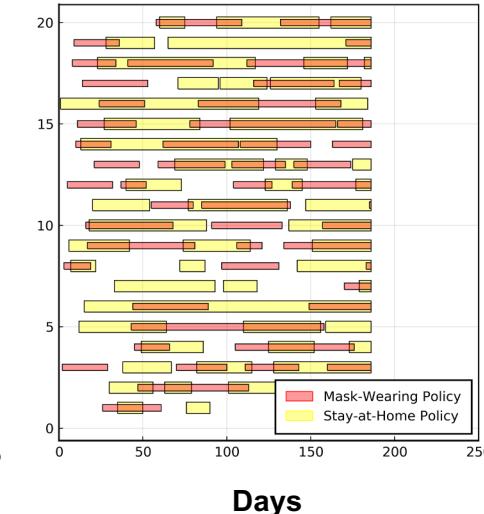
## Mask-Only



## Global Threshold



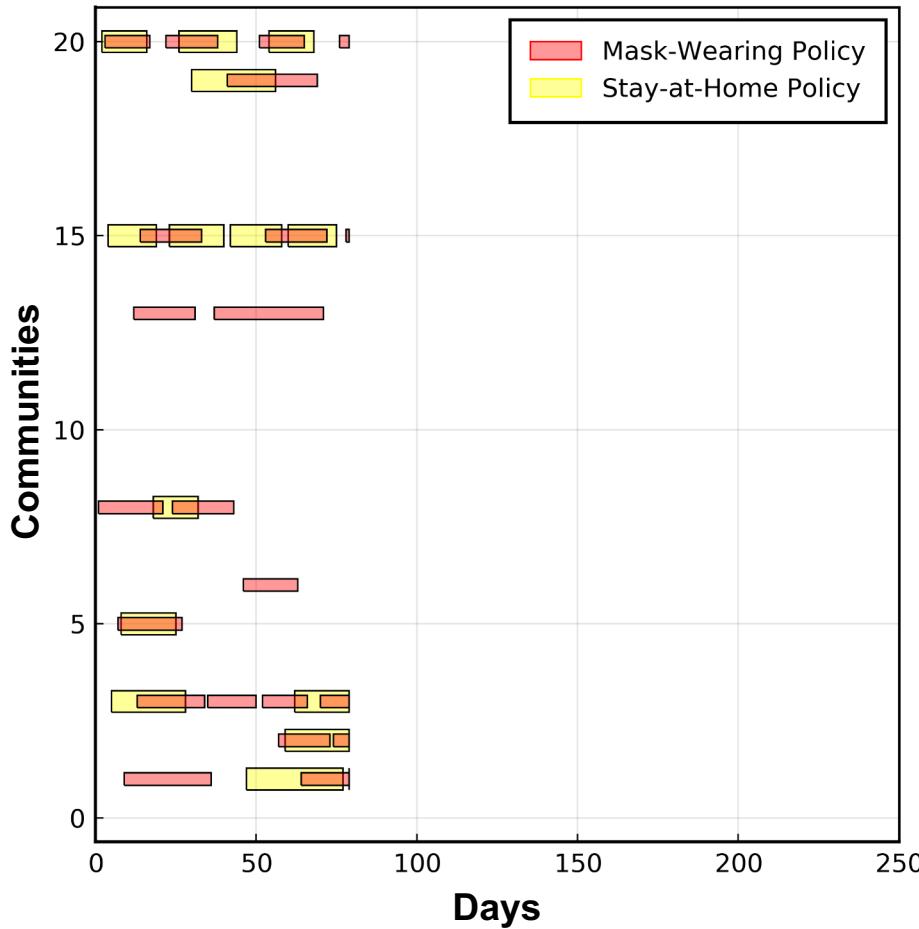
## Random Action



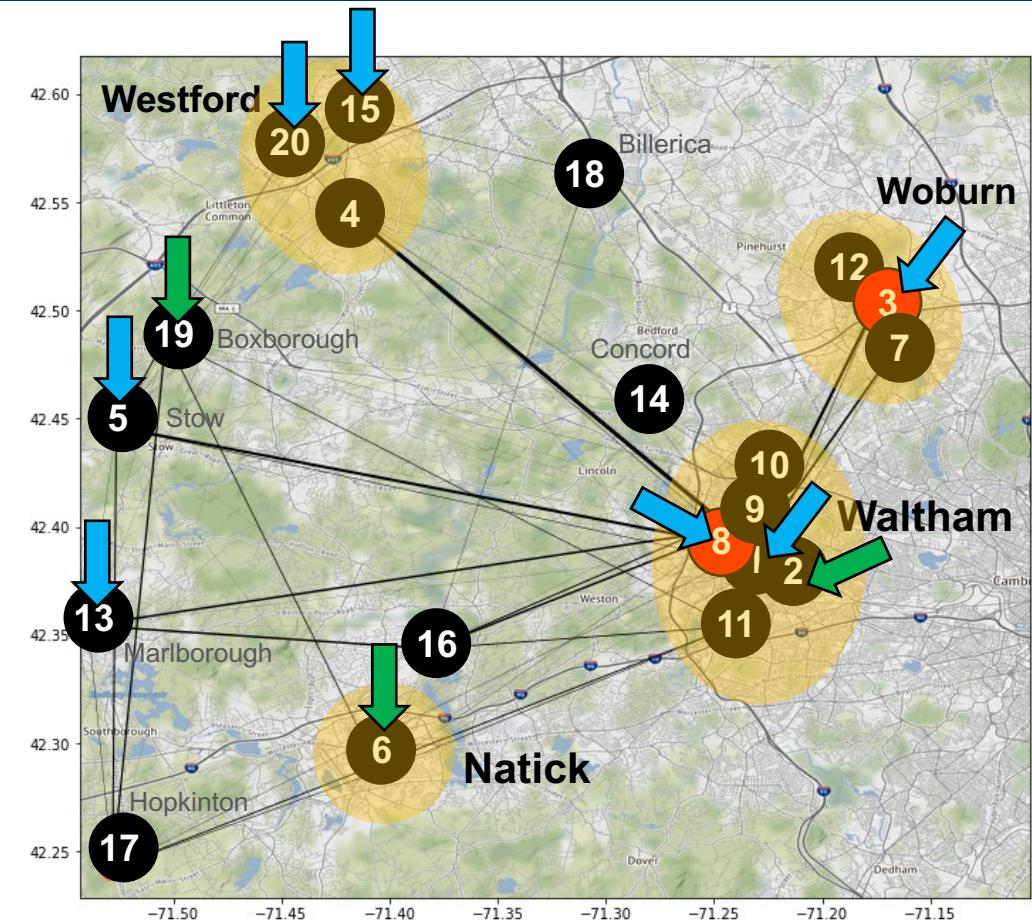
Our planner identifies much more targeted interventions while achieving higher cumulative reward



# Exploring Learned Intervention Policies



# Middlesex County, MA



**Our planner offers a sandbox environment for exploring and inferring canonical intervention strategies**



# Ongoing Work and Future Directions

## Interpretability and Usability

- Analyze planner output to identify canonical, simple strategies
- Characterization of topological effects on planning decisions
- Develop usable dashboard that assists with analyzing planner outputs

## Scalability

- Efficient and specialized network heuristics for reducing state/action spaces
- Engineering cloud computing and GPU acceleration for scalability
- Distributed hierarchical planning over complex networks

## Realism and Performance Improvement

- Designing and fusing offline and online training environments
- Probabilistic programming applied to various system components (topology, control)
- Planning with partial observability and lack of compliance



# References

---

- K. Menda, L. Laird, M. J. Kochenderfer, R. S. Caceres, “**Explaining COVID-19 Outbreaks with Reactive SEIRD Models**”, Scientific Reports 2021, under review, <https://doi.org/10.1101/2021.02.09.21251440>.
- N. Smedemark-Margulies, R. Walters, H. Zimmermann, L. Laird, N. Kaushik, R. S. Caceres and J. van de Meent, “**Inference in Network-based Epidemiological models with probabilistic programming**”, ICLR Workshop on AI for Public Health, 2021.
- R. B. Alexander, L. Laird, N. Kaushik. C. Vanderloo, N. Smedemark-Margulies, R. Walters, H. Zimmermann, L. Torres, R. S. Caceres, J. van de Meent, T. Eliassi-Rad, M. J. Kochenderfer, “**Adaptive Intervention Strategies for Epidemic Control on Population Topologies**”, in preparation.
- N. Kaushik, R. S. Caceres, L. Laird, C. Vanderloo, L. Torres, T. Eliassi-Rad, “**Network-based modeling of mobility patterns and SEIR dynamics to analyze COVID 19**”, in preparation.



# Questions

## Contact Me

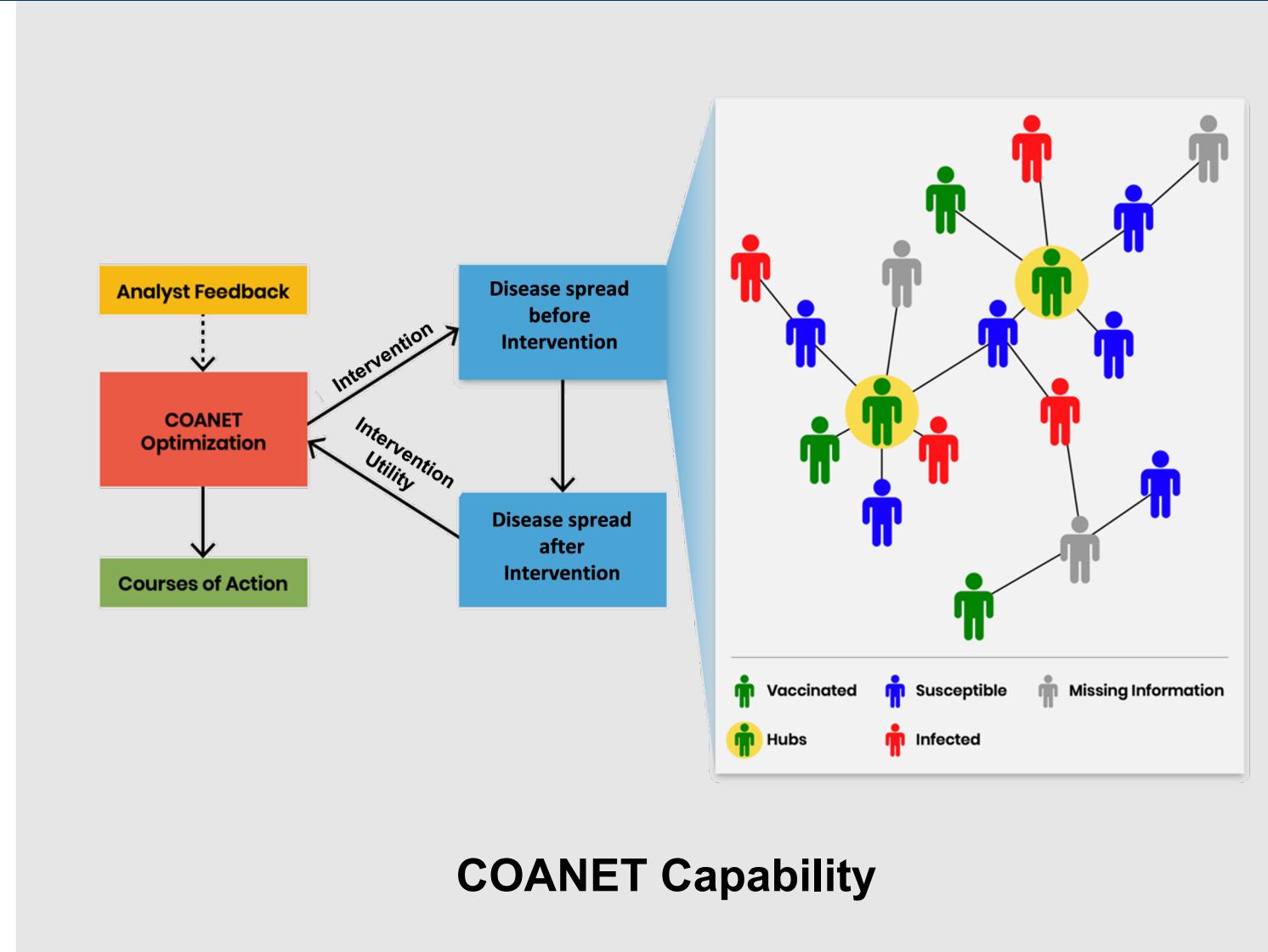
Lucas Laird

[lucas.laird@ll.mit.edu](mailto:lucas.laird@ll.mit.edu)

## Contact the Team

Rajmonda Caceres

[rajmonda.caceres@ll.mit.edu](mailto:rajmonda.caceres@ll.mit.edu)



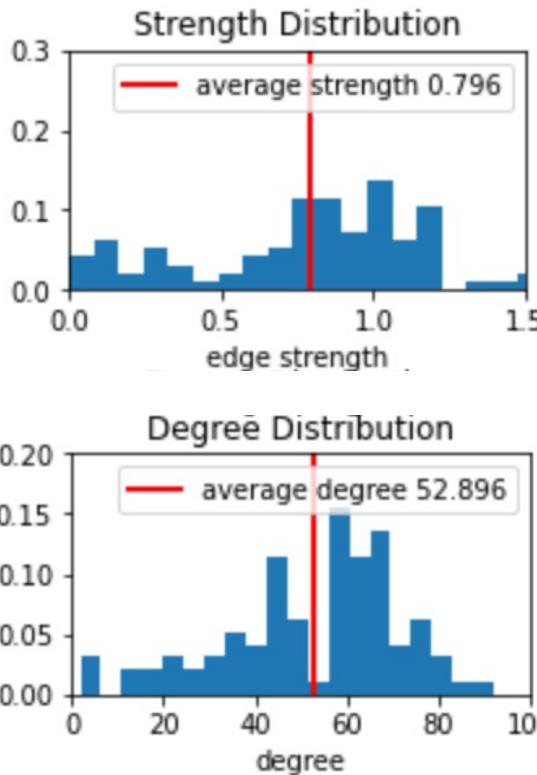


# Backup

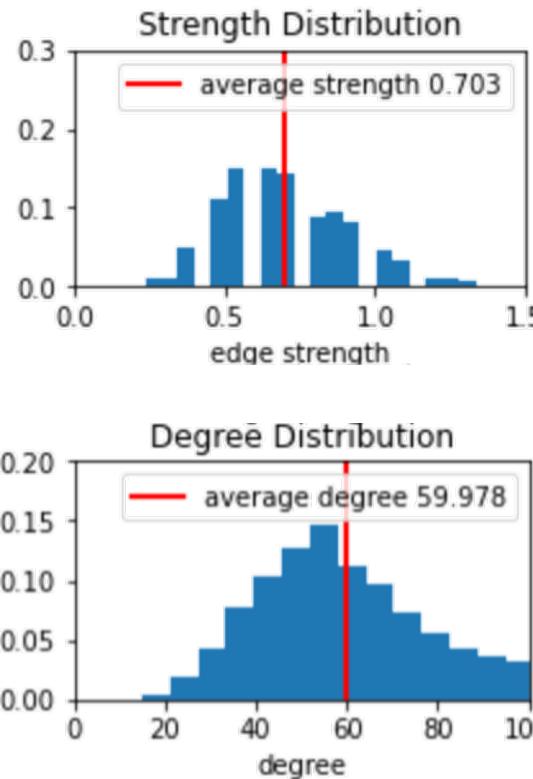


# Mobility Networks as Stochastic Block Models

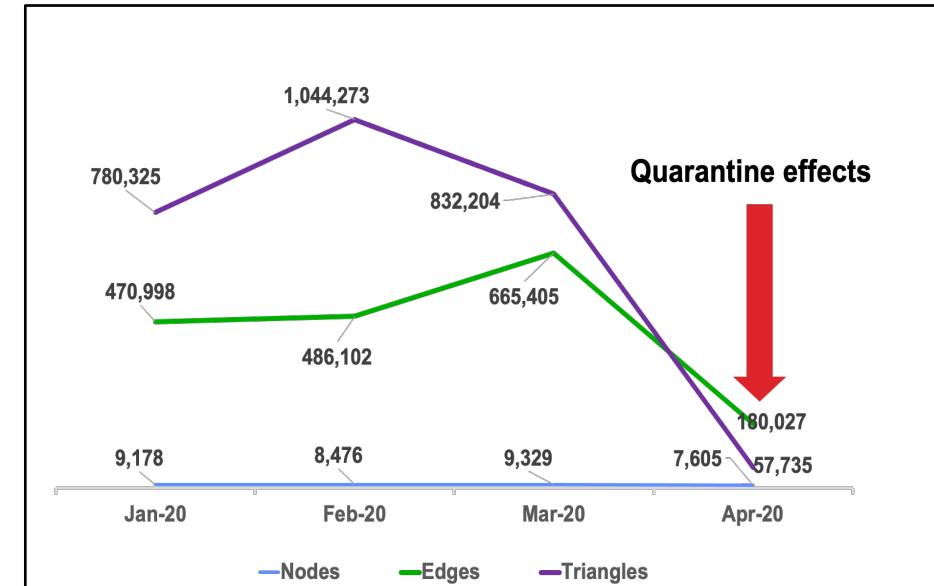
MIT Reality Mining Network<sup>1</sup>



MIT LL Network Snapshot<sup>2</sup>



MIT LL Network over Time



Our model captures contact patterns also observed in other social and mobility networks<sup>1,3,4,5</sup> and reflects observed intervention effects

<sup>3</sup> <https://www.medrxiv.org/content/10.1101/2020.06.15.20131979v1.full.pdf>

<sup>4</sup> <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7273304/>

<sup>5</sup> <https://www.networkscienceinstitute.org/publications/assessing-changes-in-commuting-and-individual-mobility-in-major-metropolitan-areas-in-the-united-states-during-the-covid-19-outbreak>

<sup>1</sup> MIT Reality Mining: <http://realitycommons.media.mit.edu/realitymining.html>

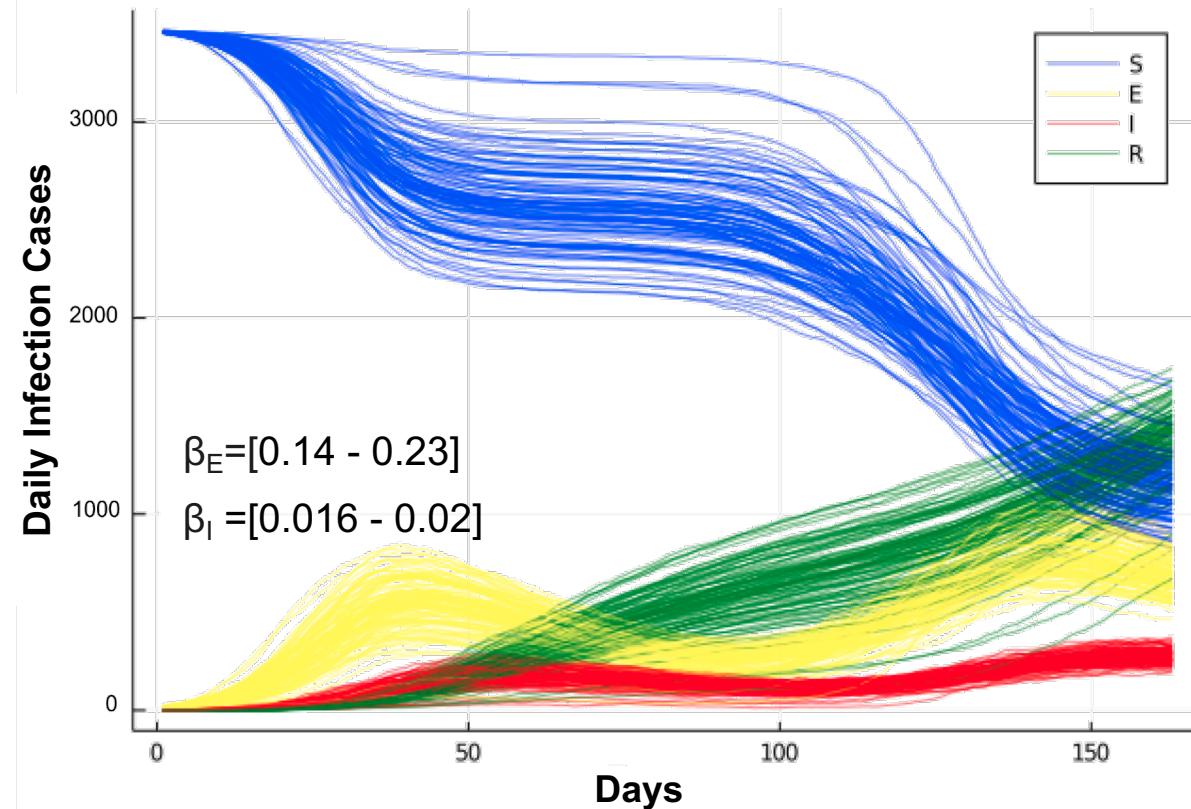
<sup>2</sup> Middlesex Network Snapshot



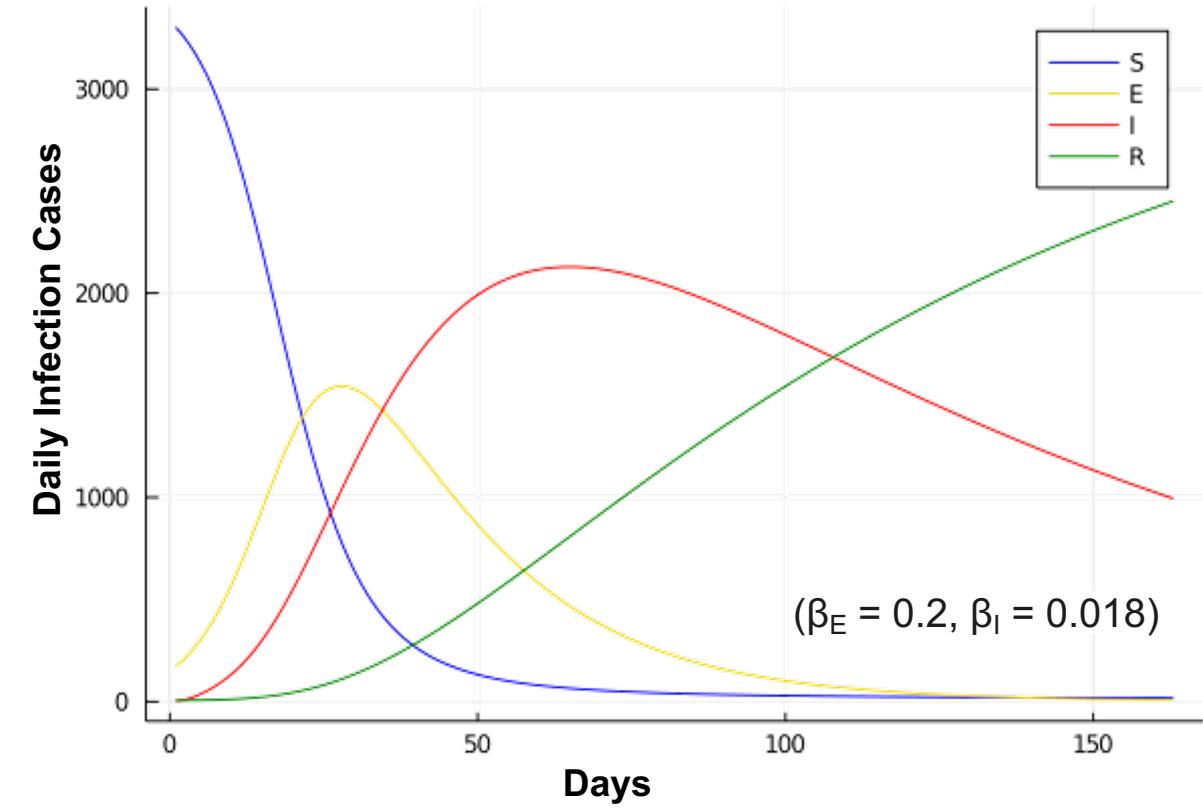
# Estimating Network SEIR Parameters from Data

(Probabilistic Programming Approach)

Our Probabilistic Programming Network-SEIR Model



Vanilla SEIR Model



Plots reflect 100 simulated disease trajectories

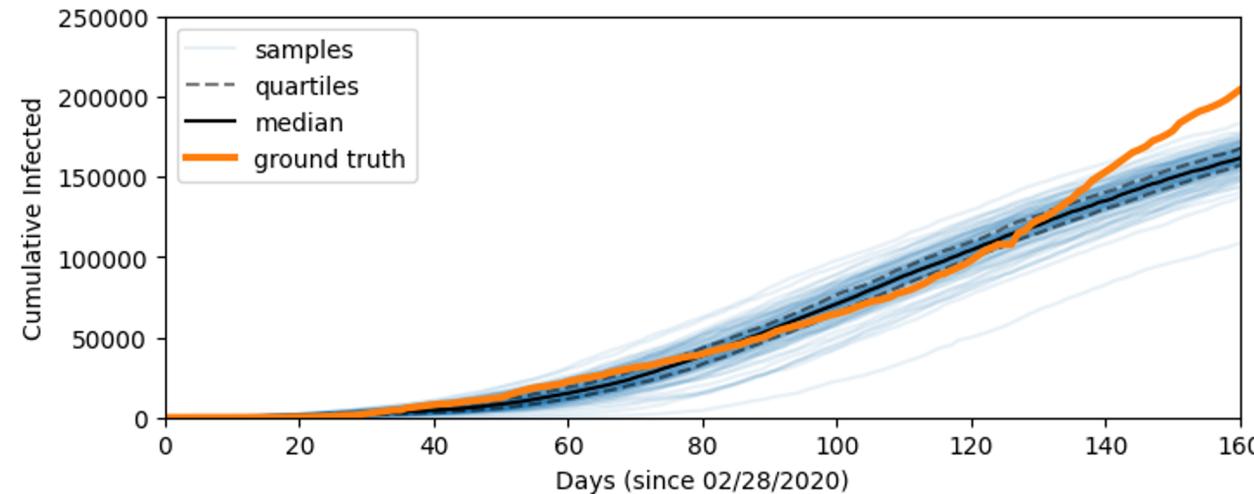
Using probabilistic programming and network SEIR, we can model more realistic disease progression



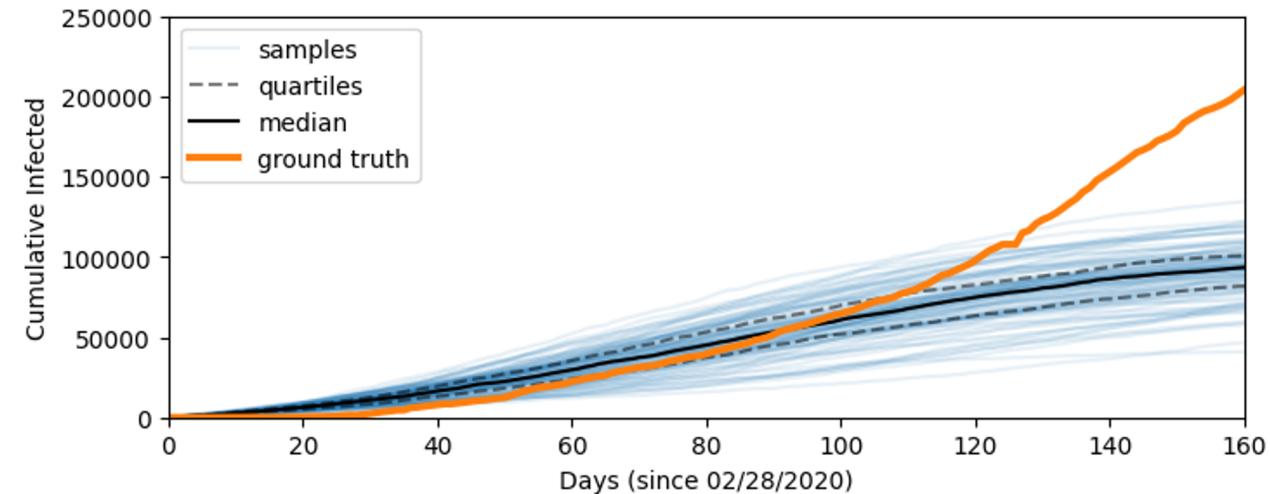
# Estimating Network SEIR Parameters from Data

## (Probabilistic Programming Approach)

### Our Method



### $R_t$ -Analytic Baseline



Disease Model	Fitting Method	LA-MDAE	Miami-Dade-MDAE
Compartmental	CE-EM	0.0251	0.0161
Network	$R_t$ -Analytic	0.0075	0.0086
Network	BBVI	<b>0.0029</b>	<b>0.0053</b>

Using probabilistic programming and network SEIR model allows for a closer fit to data

\* Network  $R_t$  Analytic estimator assumes the network has average constant degree

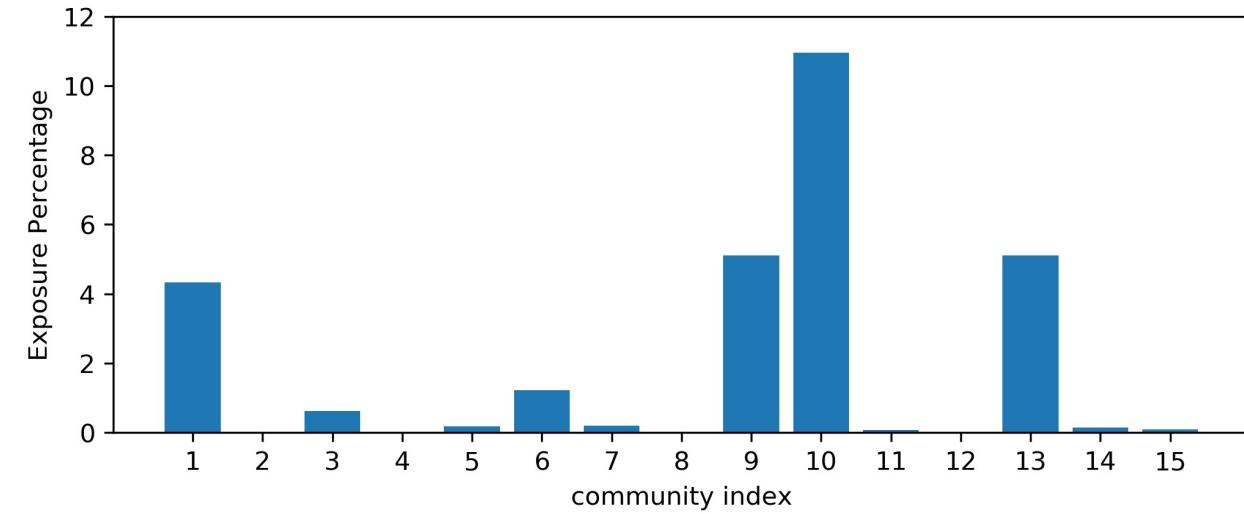
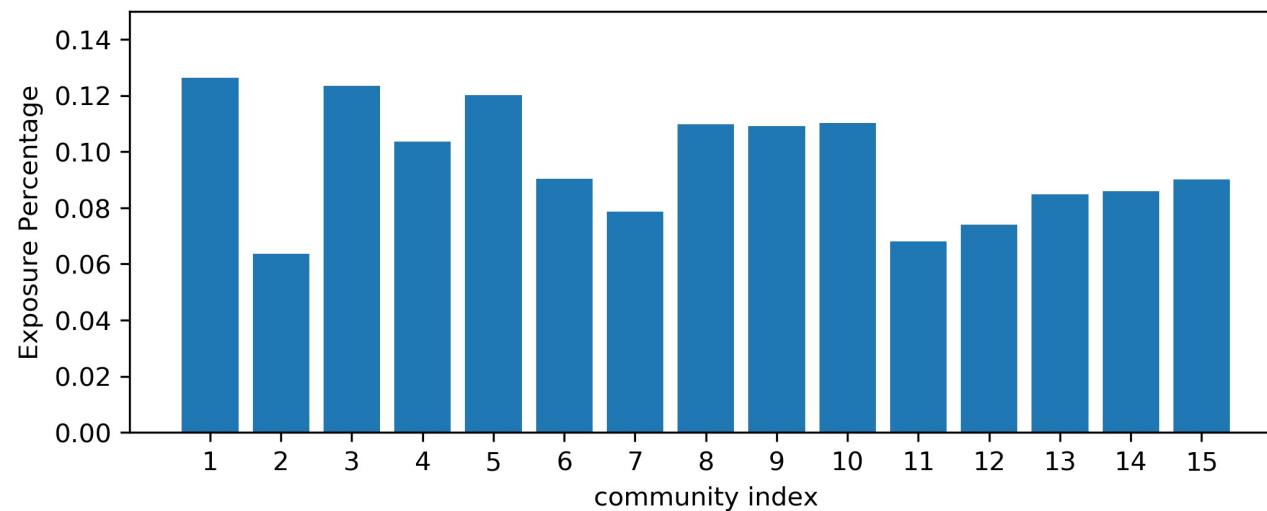
Plots reflect 100 simulated disease trajectories

MDAE: Mean Daily Absolute Error



# Inferring Starting Communities

Miami-Dade County, FL

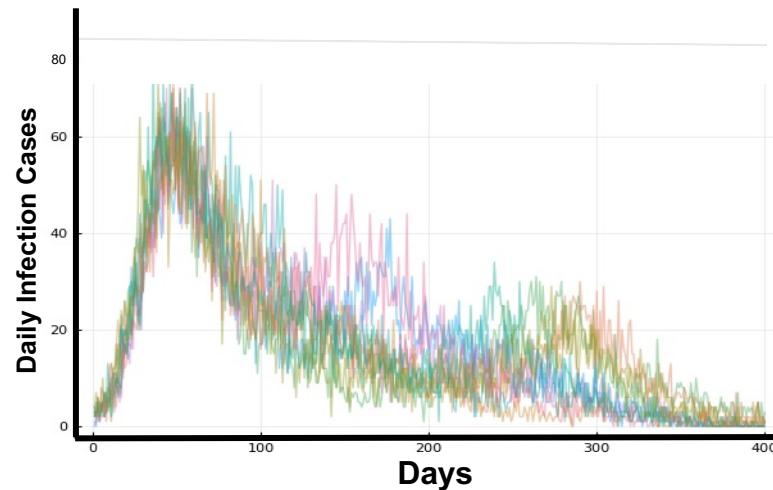


We can identify likely communities where disease could have spread from given observed data

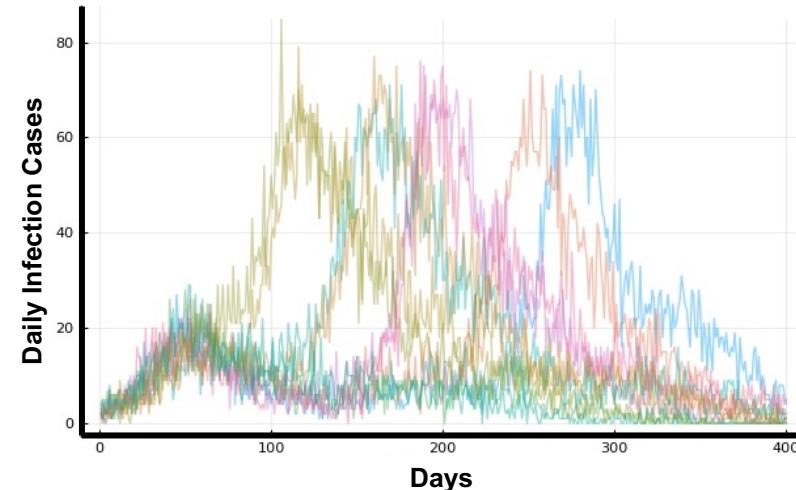


# Patterns of Disease Spread

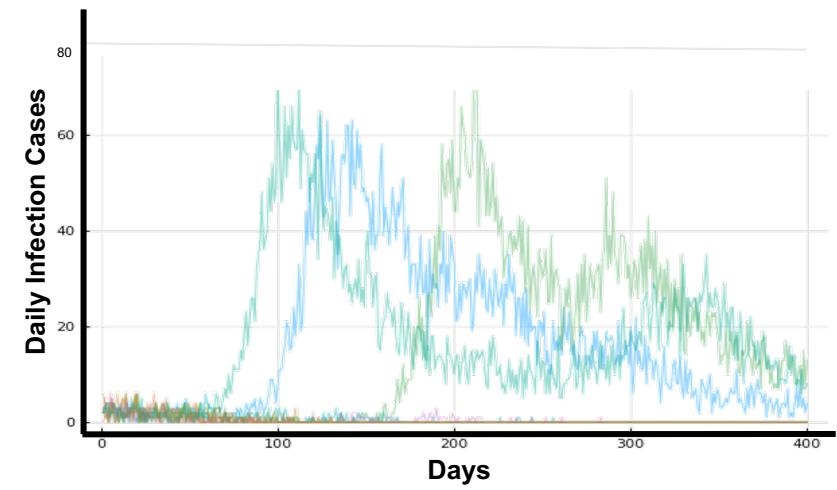
Same geographic area, same initial infections that start in different communities



Consistent Peak & Timing



Consistent Peak & Delayed Timing



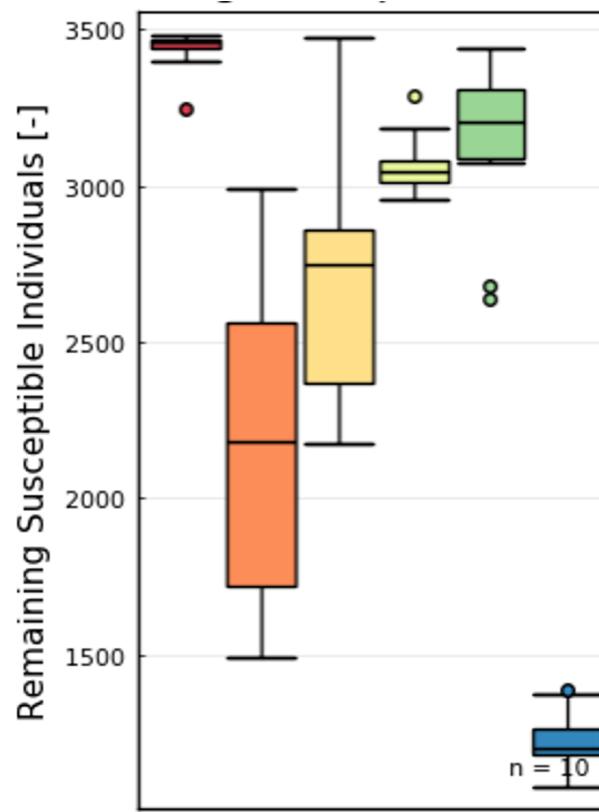
Inconsistent Peak & Timing

- Our network-SEIR model can simulate super-spreader events (sharper peaks) and recurring outbreaks (multiple peaks) that traditional SEIR models cannot capture
- We observe non-trivial, heterogeneous network effects on the patterns of disease spread

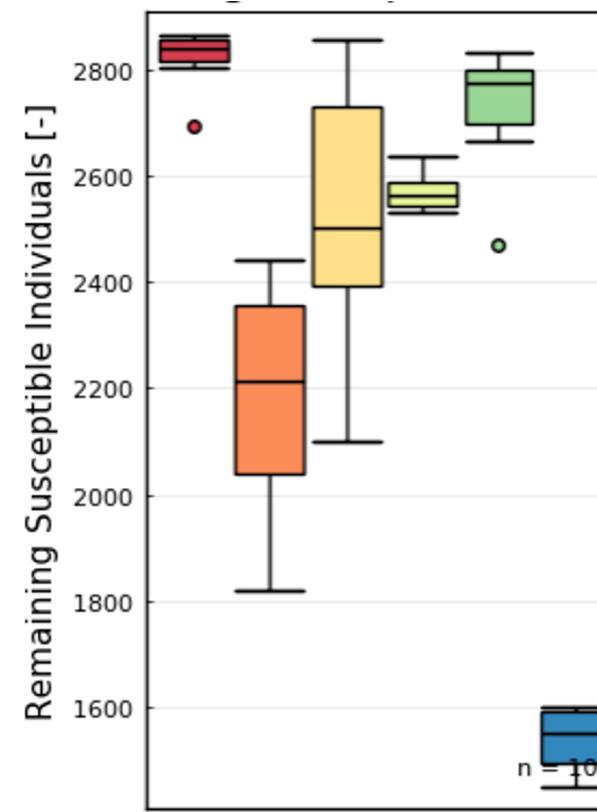


# Comparison with Non-Adaptive Heuristics

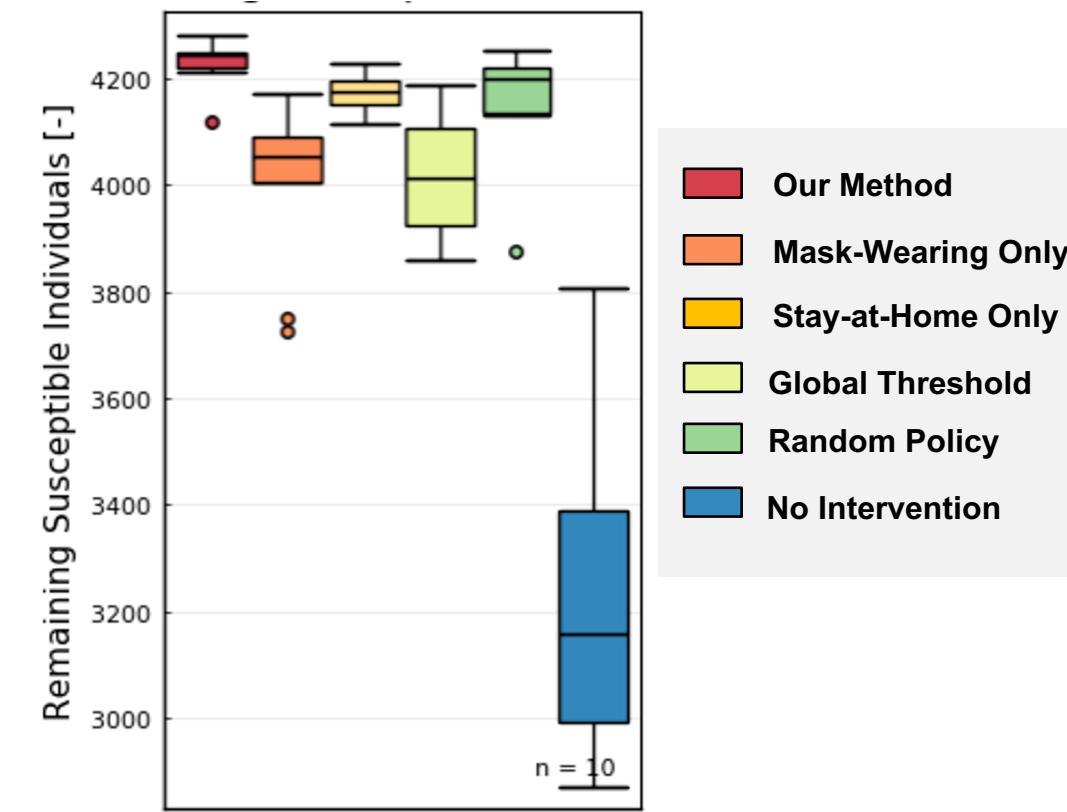
Middlesex County, MA



Miami-Dade County, FL



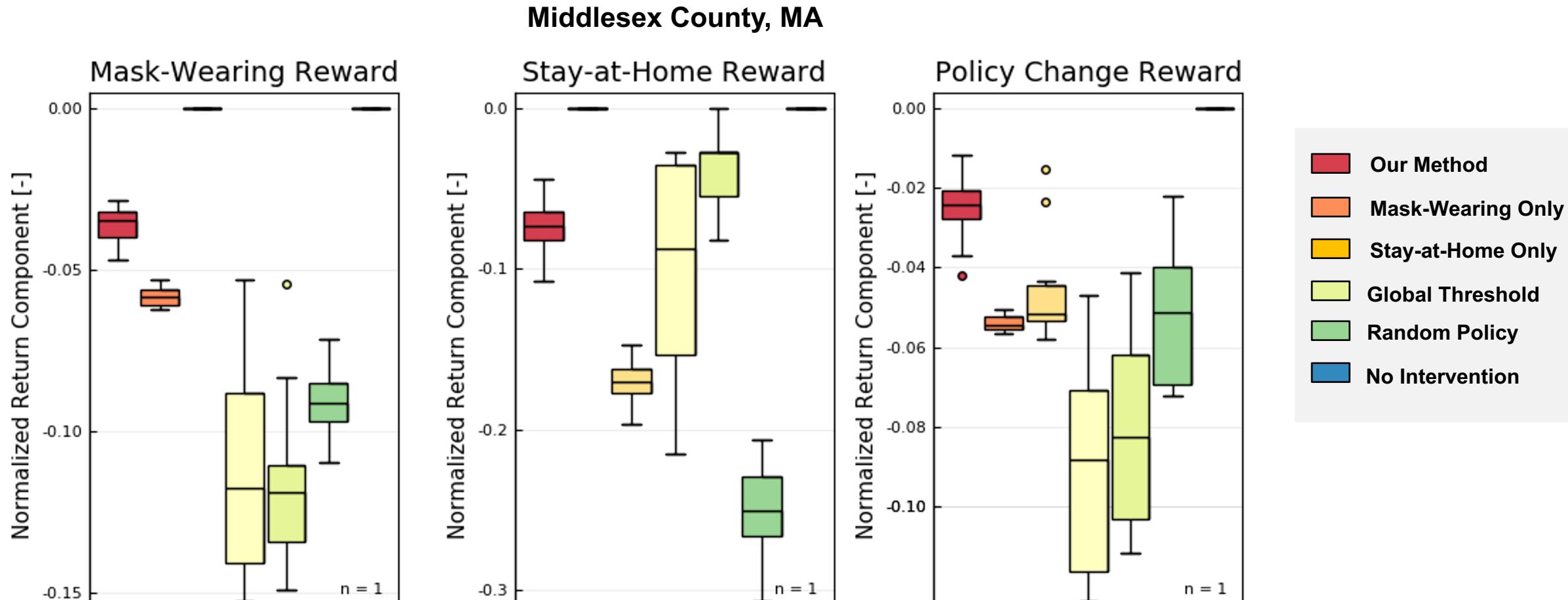
Los Angeles County, CA



Our planner leads to a smaller fraction of population being infected



# Comparison with Non-Adaptive Heuristics



Our planner finds a good trade-off between the various reward components