

# Don't Simulate Twice!

## One-shot Sensitivity Analyses via Automatic Differentiation

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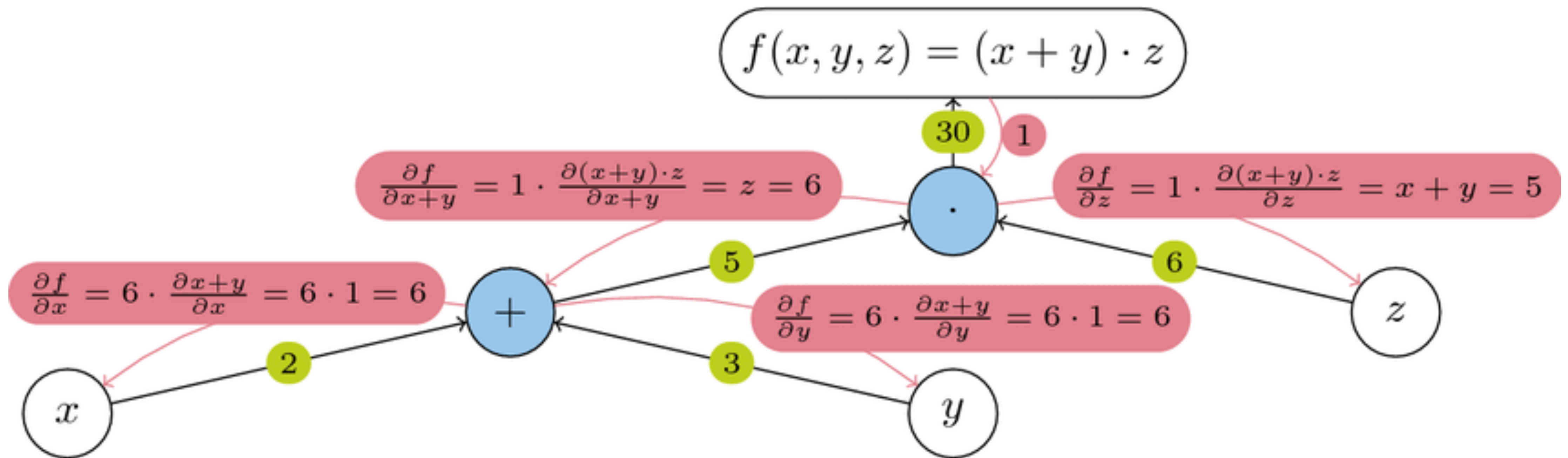


# Agent-Based Models

- ABMs promising tool to model complex systems “bottom-up”
- Wide adoption hindered by (not exhaustive):
  - Scalability
  - Robustness
  - Data availability

# Differentiable Agent-Based Models

- Idea: Use [Automatic Differentiation](#) in ABMs



# Case study: the JUNE epidemiological model

- JUNE is a 1:1 epi model of England (56 million agents)
- GradABM-JUNE is its differentiable implementation (PyTorch).

	Simulation
JUNE	50 hours
GRADABM-JUNE (GPU)	5 seconds

**Tensorisation enables scalability to millions (billions?) of agents**

**Ref: Ayush Chopra previous talk**

# Case study: the JUNE epidemiological model

- We can use gradient descent / variational inference for calibration

	Simulation	Calibration (No UQ)	Bayesian Calibration
JUNE	50 hours	-	100k hours
GRADABM-JUNE (GPU)	5 seconds	20 minutes	8 hours

Paper coming soon...

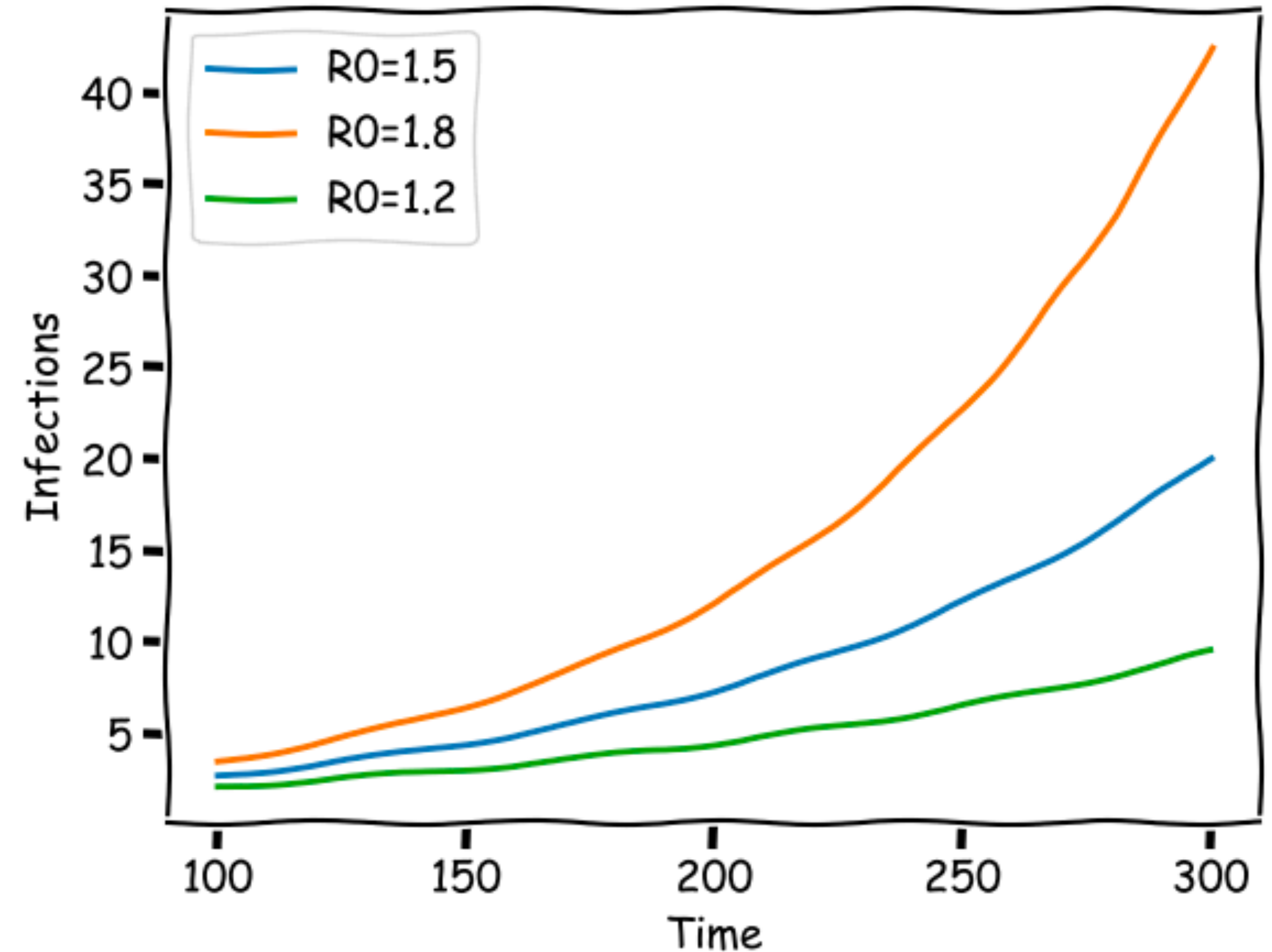
**Differentiability enables fast  
and accurate model calibration**

# Sensitivity Analyses

Why is it necessary?

## 1. Robustness

- Example: Epidemiological ABM
- One parameter:  $R_0$ , an expert measures to be  $R_0 = 1.5 \pm 0.3$



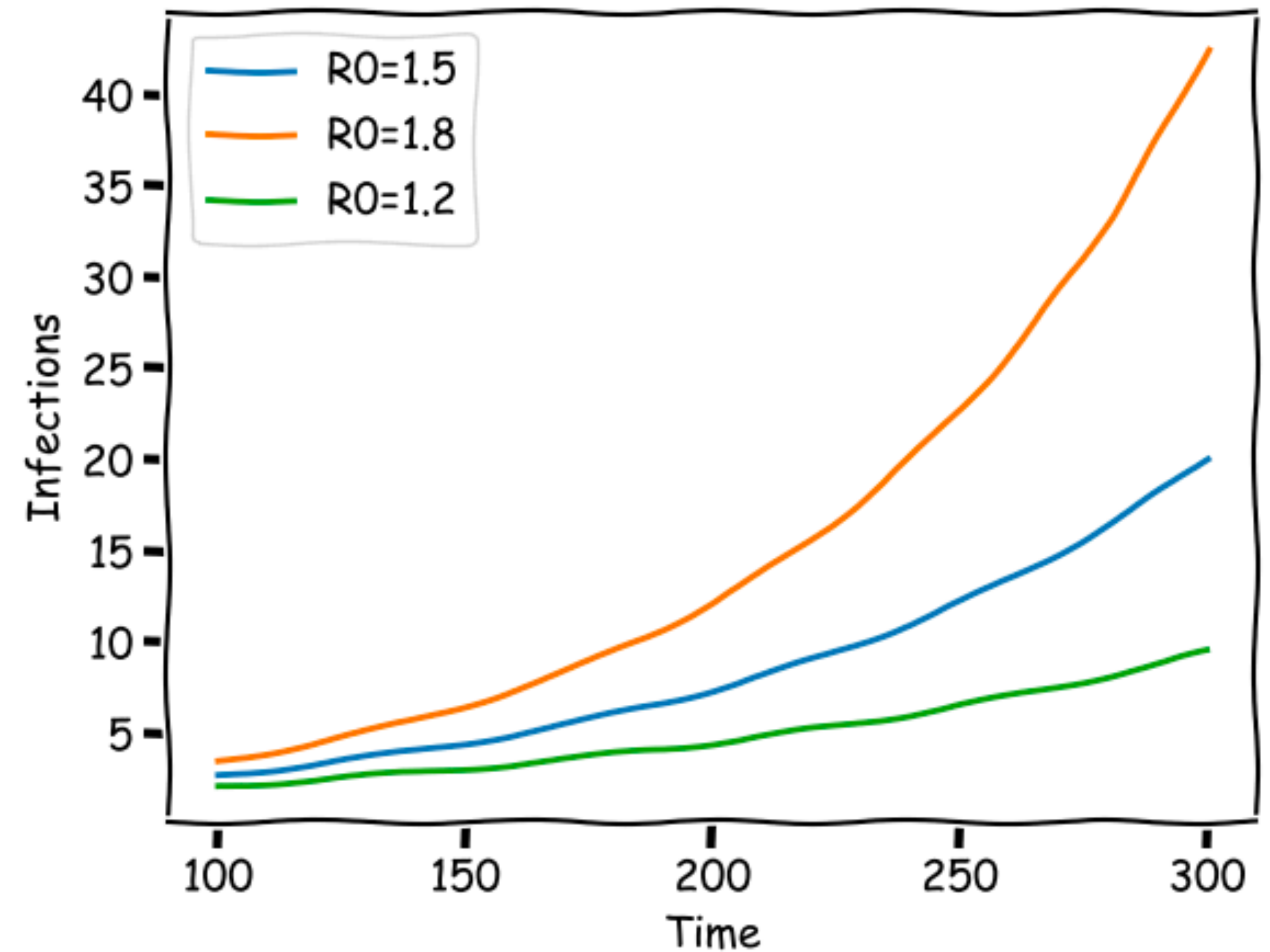
**Sensitivity Analyses  
crucial for policy evaluation**

# Sensitivity Analyses

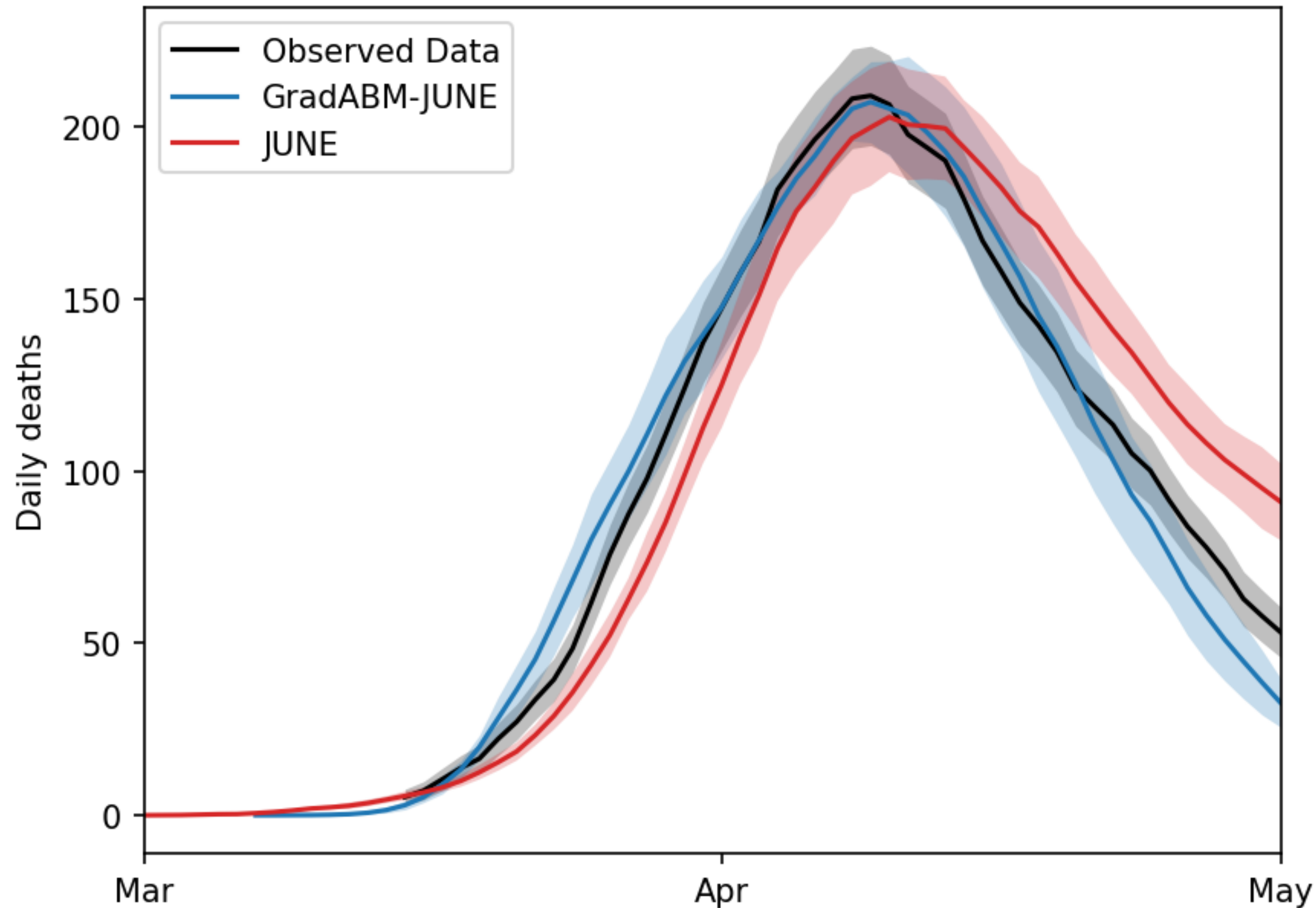
Why is it necessary?

## 2. Interpretability

- Sensitive parameters tells us what's important in the model.



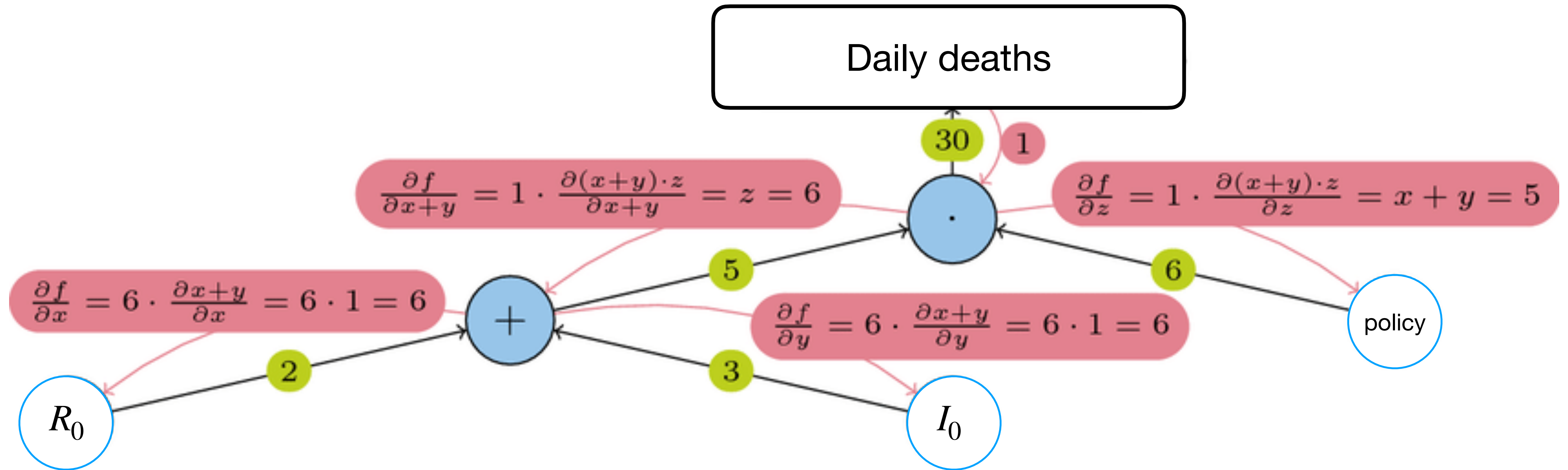
# JUNE: London case study



11 Free parameters:  
10 contact intensity locations  
(schools, companies, etc.)  
+  
Initial number of cases



# Sensitivity Analysis via Automatic Differentiation



	Simulation	Calibration	Regression	Sensitivity Analysis
JUNE	<b>Reverse-mode AD enables almost instant SA</b>			5k hours
GRADABM-JUNE (GPU)				10 seconds

# The impact of uncertainty on predictions of the CovidSim epidemiological code

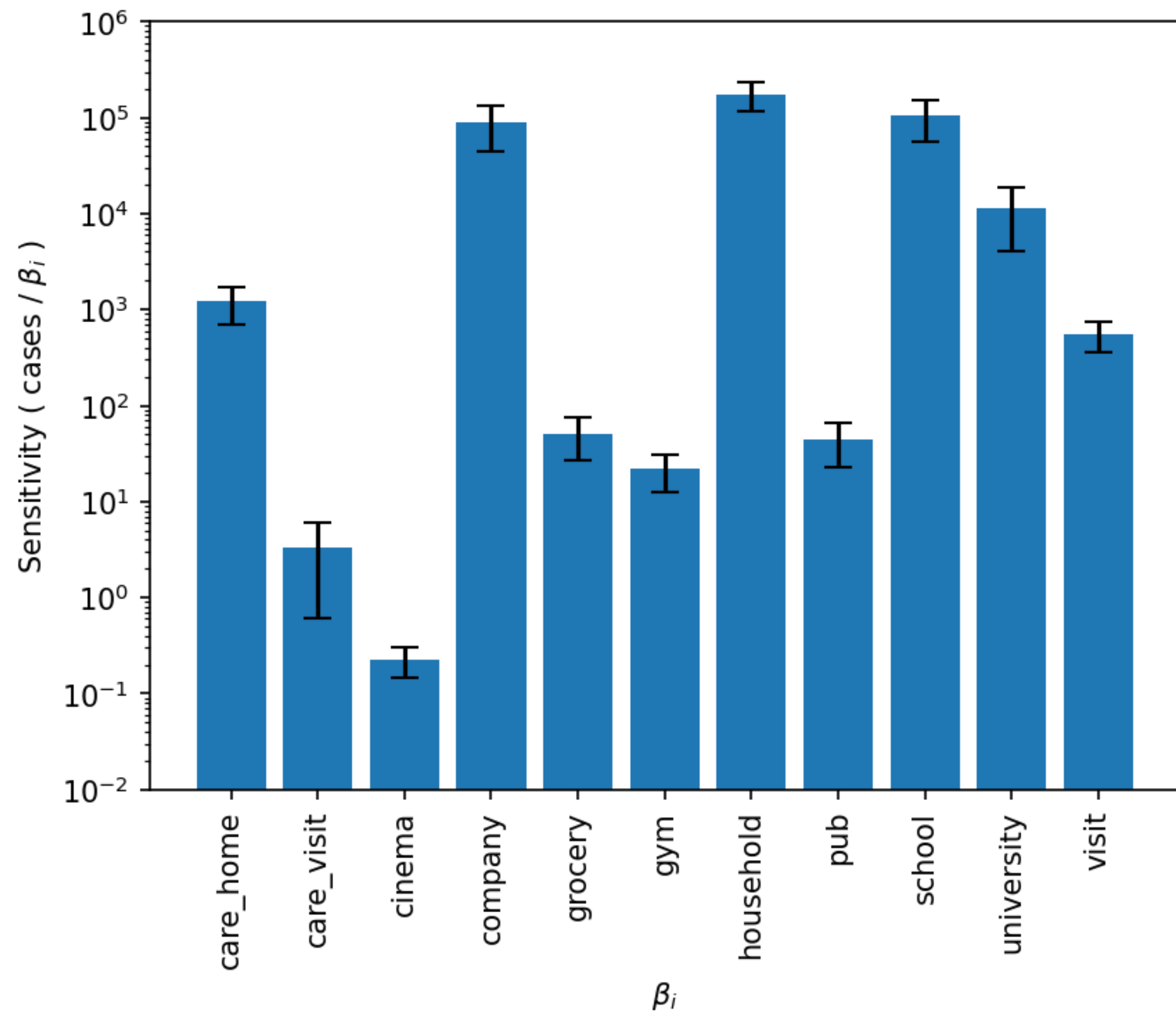
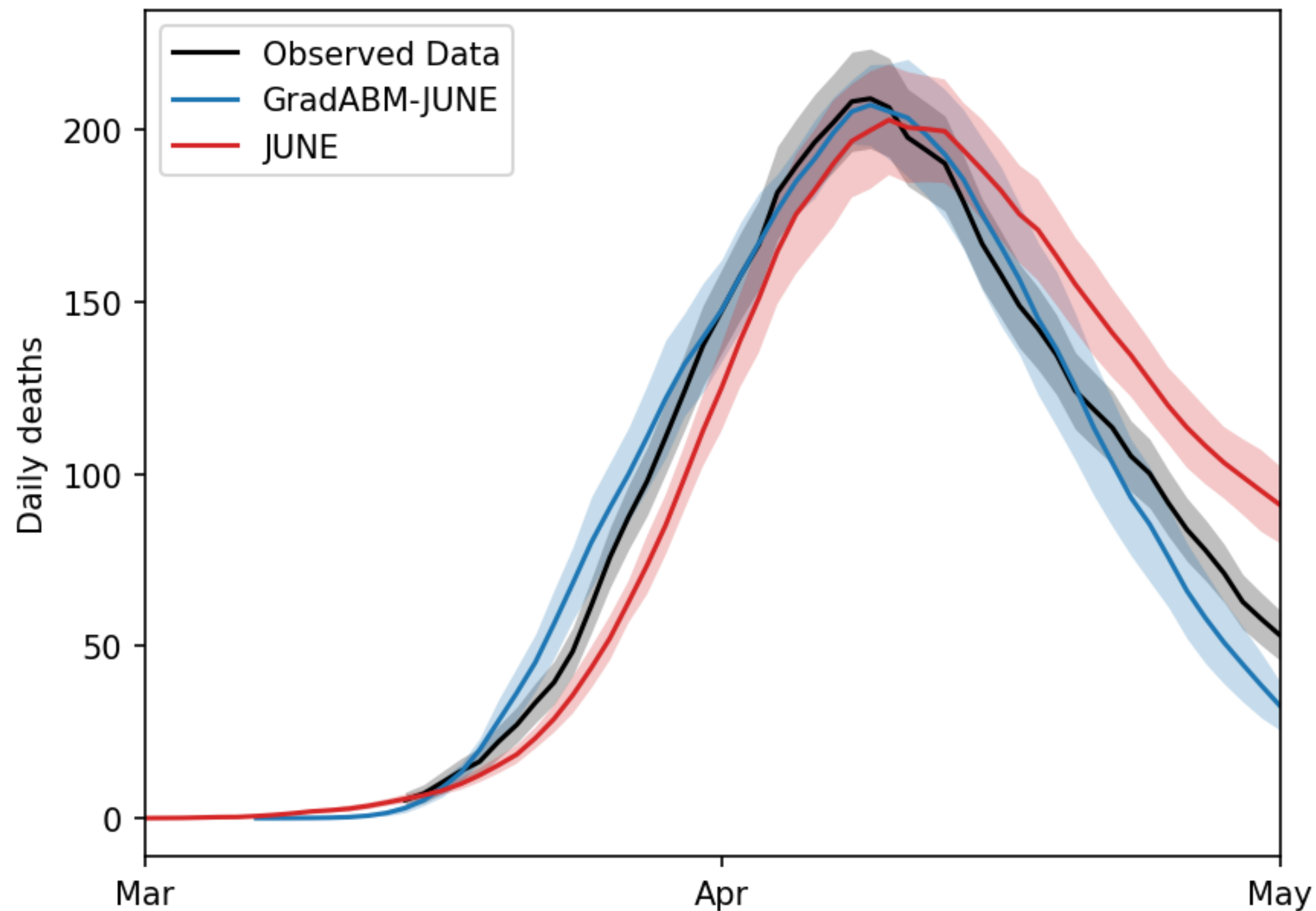
Wouter Edeling<sup>1</sup>, Hamid Arabnejad<sup>2</sup>, Robbie Sinclair<sup>3</sup>, Diana Suleimenova<sup>2</sup>,  
Krishnakumar Gopalakrishnan<sup>3</sup>, Bartosz Bosak<sup>4</sup>, Derek Groen<sup>2</sup>, Imran Mahmood<sup>2</sup>,  
Daan Crommelin<sup>1,5</sup> and Peter V. Coveney<sup>3,6</sup> ✉

Epidemiological modelling has assisted in identifying interventions that reduce the impact of COVID-19. The UK government relied, in part, on the CovidSim model to guide its policy to contain the rapid spread of the COVID-19 pandemic during March and April 2020; however, CovidSim contains several sources of uncertainty that affect the quality of its predictions: parametric uncertainty, model structure uncertainty and scenario uncertainty. **Here we report on parametric sensitivity analysis and uncertainty quantification of the code. From the 940 parameters used as input into CovidSim, we find a subset of 19 to which the code output is most sensitive**—imperfect knowledge of these inputs is magnified in the outputs by up to 300%. The model displays substantial bias with respect to observed data, failing to describe validation data well. Quantifying parametric input uncertainty is therefore not sufficient: the effect of model structure and scenario uncertainty must also be properly understood.

*Ensemble execution.* Consequently, through the use of adaptive methods we make the uncertainty analysis of CovidSim tractable, but our analysis nevertheless required us to perform thousands of runs, each with its own unique set of input parameters. Specifically, we used the Eagle supercomputer at the Posnan

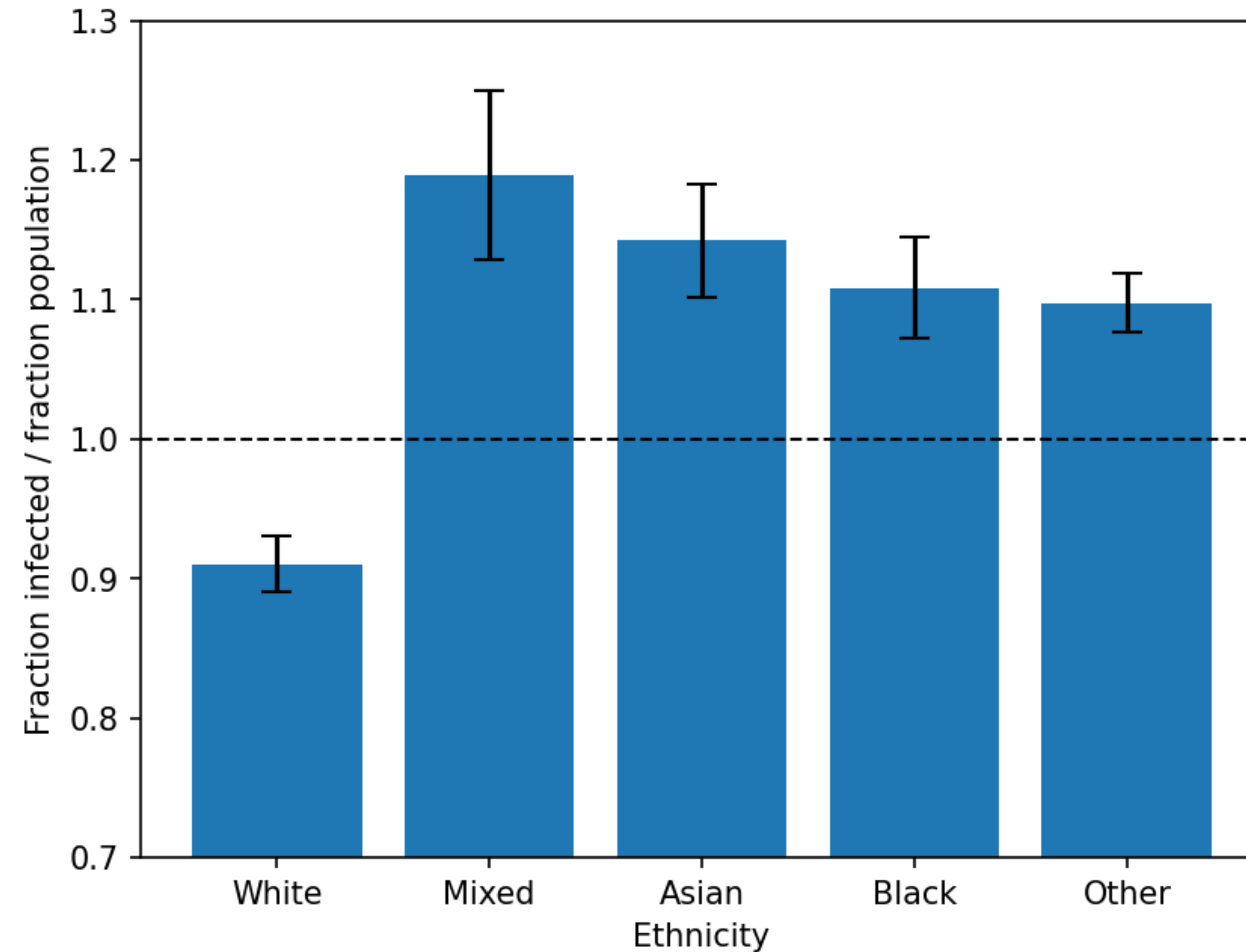
**Reverse-mode AD  
independent of number of  
parameters!**

# Sensitivity Analysis



# Interrogating the model

JUNE recovers infection inequalities across demographic groups **without** explicit calibration



**What causes this imbalance?**

# Analysing the sensitivity of each demographic group

Infected population in group d

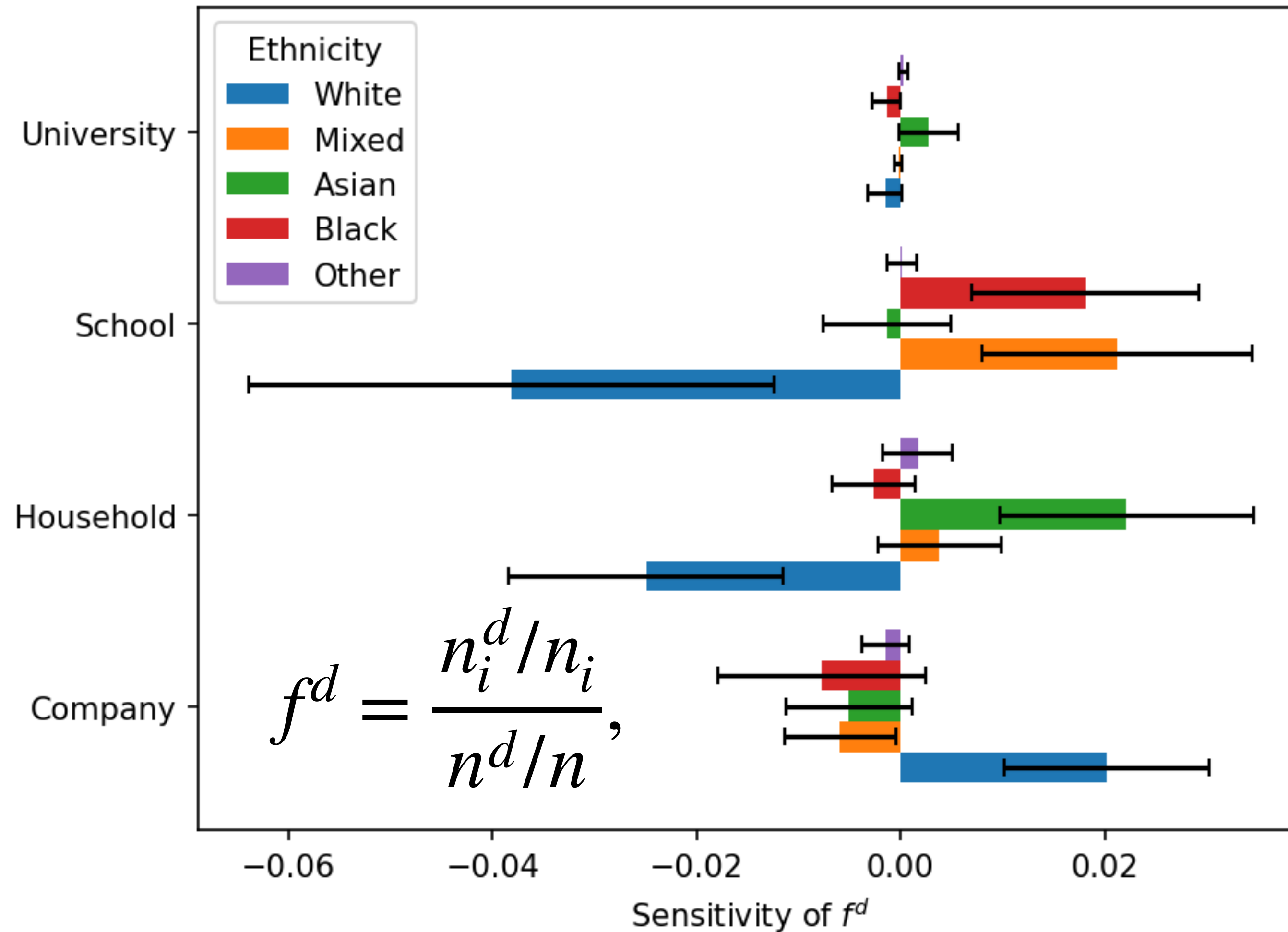
Infected population

$$f^d = \frac{n_i^d / n_i}{n^d / n},$$

Population in group d

Total population

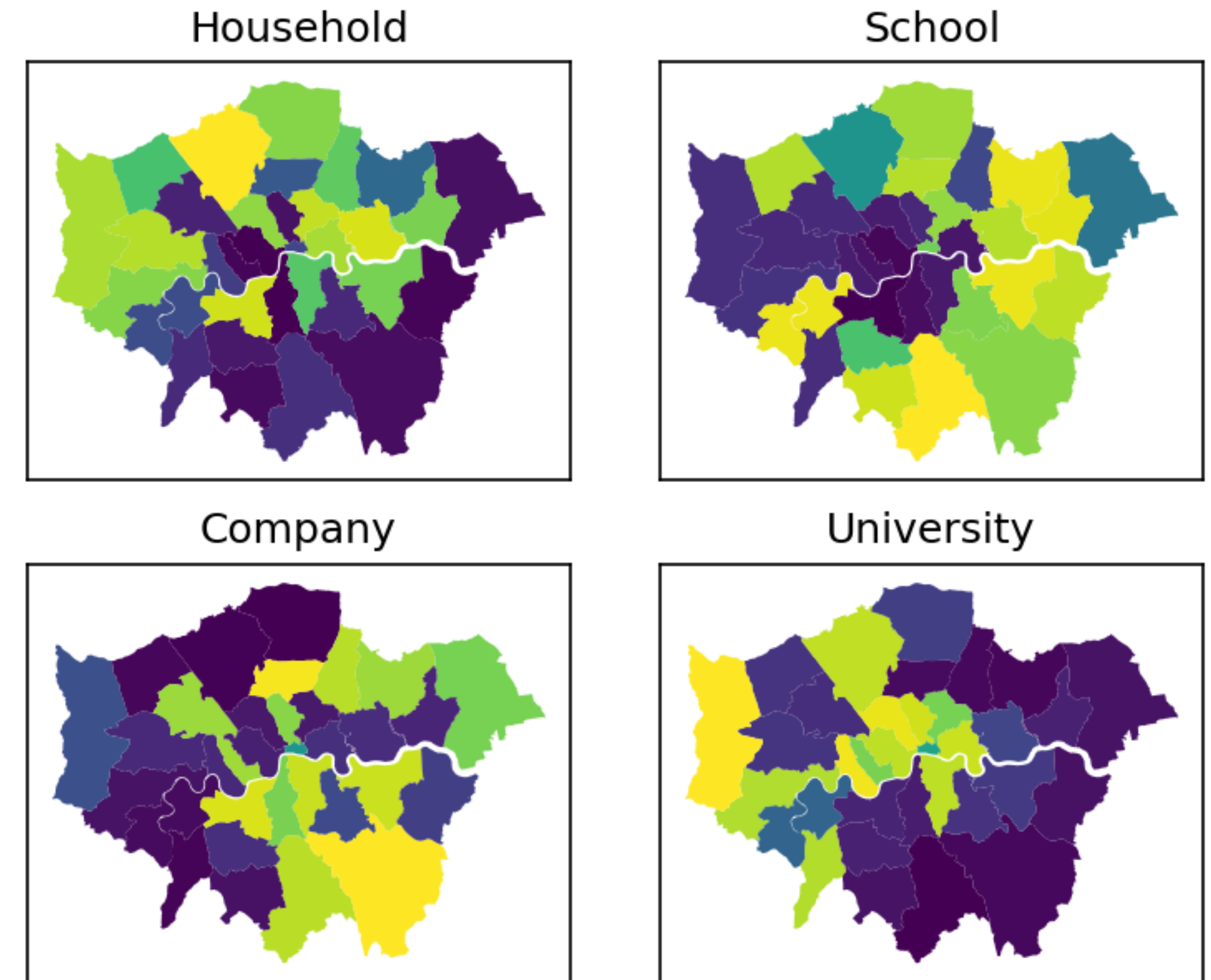
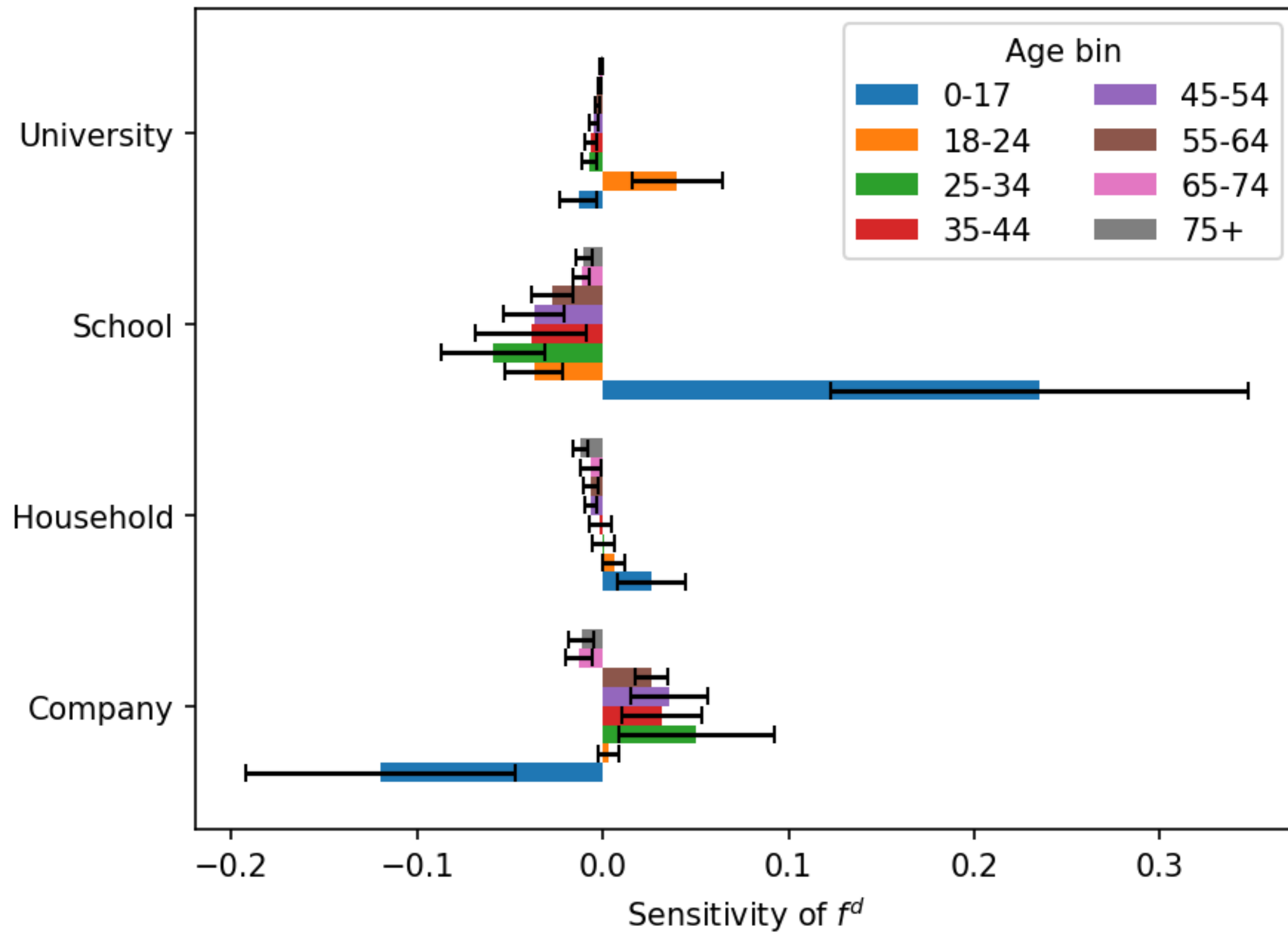
# Analysing the sensitivity of each demographic group



Some Ethnic groups are more vulnerable to infections in certain locations.

This is due to household size, work sector, family structure, etc.

# Analysing the sensitivity of each demographic group



# Conclusions

Differentiable agent-based models enable:

1. Fast **simulation** via tensorisation.
2. Fast and accurate Bayesian **calibration** via gradients.
3. Fast and accurate **sensitivity analyses** via gradients.

Paper + slides: [www.arnau.ai/aamas23](http://www.arnau.ai/aamas23)