

# **The Impact of Uncertainty on the CovidSim Pandemic Code**

INI Event: The Role of Uncertainty in Mathematical Modelling of Pandemics

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# CovidSim: model



- CovidSim: an individual-based epidemiological code (Imperial College).
- Modified from an earlier influenza version.
- March 2020: predict effect of [Non-Pharmaceutical Interventions \(NPIs\)](#) to reduce spread of COVID19 in UK.
  
- Creates a network of individuals based on population density data.
- Individual interact within places:
  - Household
  - Schools
  - Universities
  - Workplaces
  
- Influential model: key paper (Report 9 [1]) was (in part) responsible for reorienting UK policy from herd immunity to suppression.
  
- We performed uncertainty analysis on behalf of UK RAMP team.

- CovidSim is uncertain:
  - 1) **Parametric uncertainty**: input parameter values  $\xi \in \mathbb{R}^d$  are not known exactly.
  - 2) **Model-form uncertainty**: what is the appropriate mathematical structure of the model? For instance: missing epidemiological processes or missing places types.
  - 3) **Scenario uncertainty**: uncertainty in the scenario  $\mathcal{S}$  under which the model  $\mathcal{M}$  is applied. For instance: initial conditions or selected NPI measures.
- The output Quantity of Interest (QoI):  $q = q(\xi, \mathcal{M}, \mathcal{S})$
- We quantified the uncertainty in  $q$  due to parametric uncertainty, and indirectly due to scenario uncertainty.

- Specifically, we looked at **robustness**:
  - How much is parametric uncertainty amplified from the input to the output?
- Involves computing mean and variance of  $q$ :

$$\mathbb{E}[q \mid \mathcal{M}, \mathcal{S}] := \int_{\Omega_{\xi}} q(\xi, \mathcal{M}, \mathcal{S}) p(\xi) d\xi,$$

$$\mathbb{V}[q \mid \mathcal{M}, \mathcal{S}] = \int_{\Omega_{\xi}} (q - \mathbb{E}[q \mid \mathcal{M}, \mathcal{S}])^2 p(\xi) d\xi$$

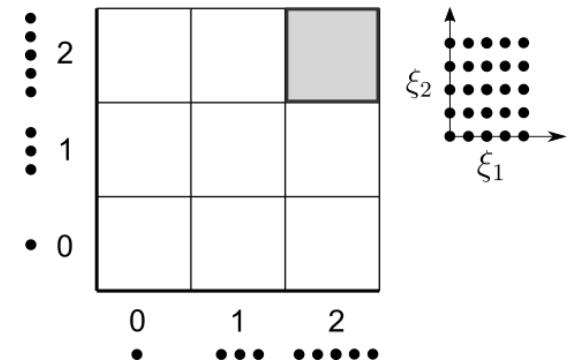
- Important: results are conditional on the model and application scenario.
- Problem 1:  $\xi \in \mathbb{R}^d$  with  $d = 940$ .
- Problem 2:  $p(\xi)$  is not given, but must be chosen.

# CovidSim: input dimension

- Problem 1:  $\xi \in \mathbb{R}^d$  with  $d = 940$ .
  - Many are not relevant, manual selection: 60 inputs left
  - Divided these into 3 groups:
    - 1) **Intervention parameters**, e.g. “Household compliance w/ quarantine”
    - 2) **Biological disease parameters**, e.g. “Latent period”.
    - 3) **Spatial / Geographic parameters**, e.g. “Relative place contact rates”.
  - Perform preliminary UQ on each group & identify important inputs.
  - Final UQ campaign: include all identified important inputs: 19 inputs.
- Problem 2:  $p(\xi)$  is not given, but must be chosen.
  - We used expert opinion: CovidSim developers.
  - We used uniform inputs due to lack of knowledge.

# CovidSim: input dimension

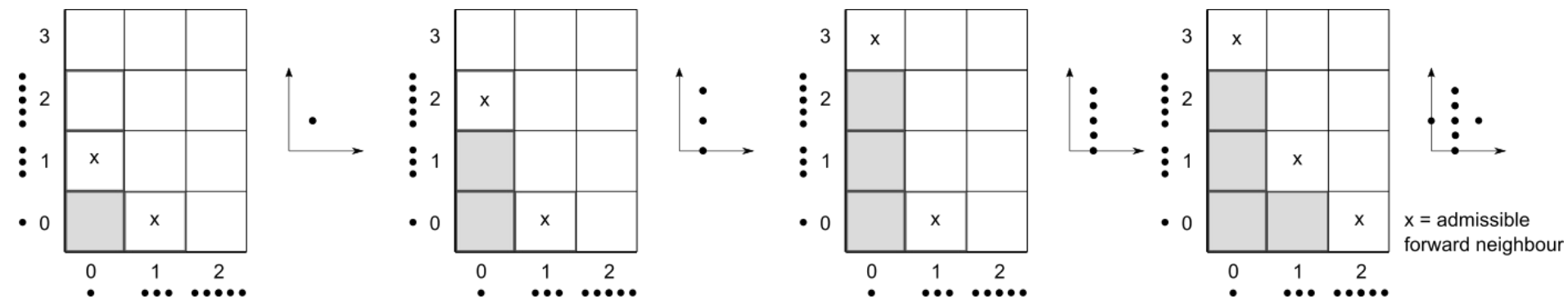
- Final campaign: 19 inputs.
  - This is still quite high!
  - EasyVVUQ: (Quasi) Monte Carlo, Polynomial Chaos, Stochastic Collocation.
  - None are ideal: we implemented a **dimension-adaptive SC sampler [2]**.
  
- Stochastic Collocation:
  - Polynomial expansion
  - Basic building blocks: 1D quadrature rules of  $m$  points.
  
  - Standard SC: extend to  $d$  dimensions
  - via a single tensor product.
  - Computational cost:  $m^d$



$$q(\boldsymbol{\xi}) \approx \tilde{q}(\boldsymbol{\xi}) = \sum_{j_1=1}^{m_1} \cdots \sum_{j_d=1}^{m_d} q(\xi_{j_1}, \cdots, \xi_{j_d}) a_{j_1}(\xi_1) \otimes \cdots \otimes a_{j_d}(\xi_d)$$

# Curse of dimensionality

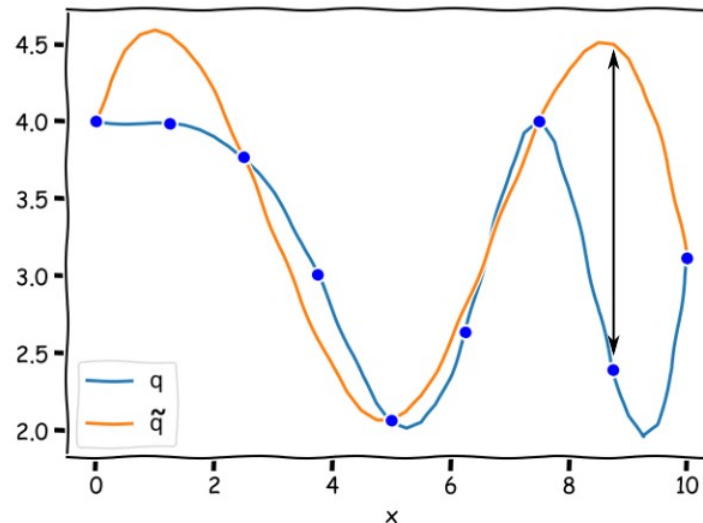
- For  $d = 19$ ,  $m^d$  is far too expensive.
- Dimension-adaptive sampling: postpone the curse of dimensionality
- Basic idea:
  - **Initialize:** start with a single sample.
  - **Look ahead:** evaluate code in ‘candidate directions’
  - **Rank order:** compute error metric for all directions
  - **Adapt:** only add direction with highest error to sampling plan



# Error measure

- **Rank order:** compute error metric for all directions
- Common error choice: **hierarchical surplus**
  - Difference between code  $q$  and SC interpolation  $\tilde{q}$  at new candidate inputs

$$\Delta \left( \boldsymbol{\xi}_j^{(l)} \right) := q \left( \boldsymbol{\xi}_j^{(l)} \right) - \tilde{q}_\Lambda \left( \boldsymbol{\xi}_j^{(l)} \right), \quad \boldsymbol{\xi}_j^{(l)} \in X_l \setminus X_\Lambda.$$

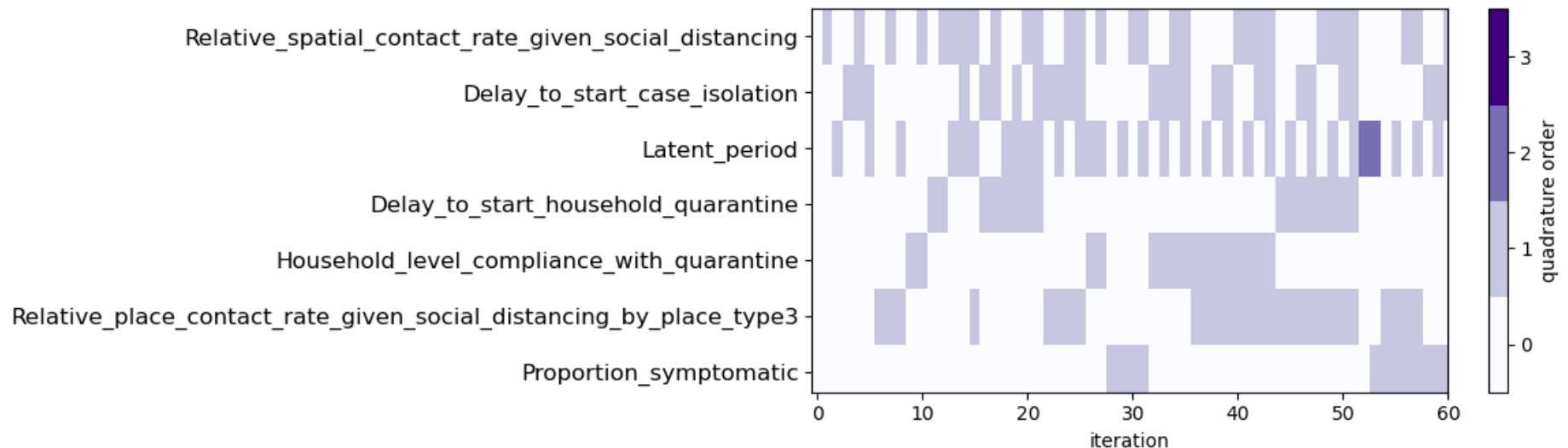




# Adaptation



- Visualization of adaptation for CovidSim:

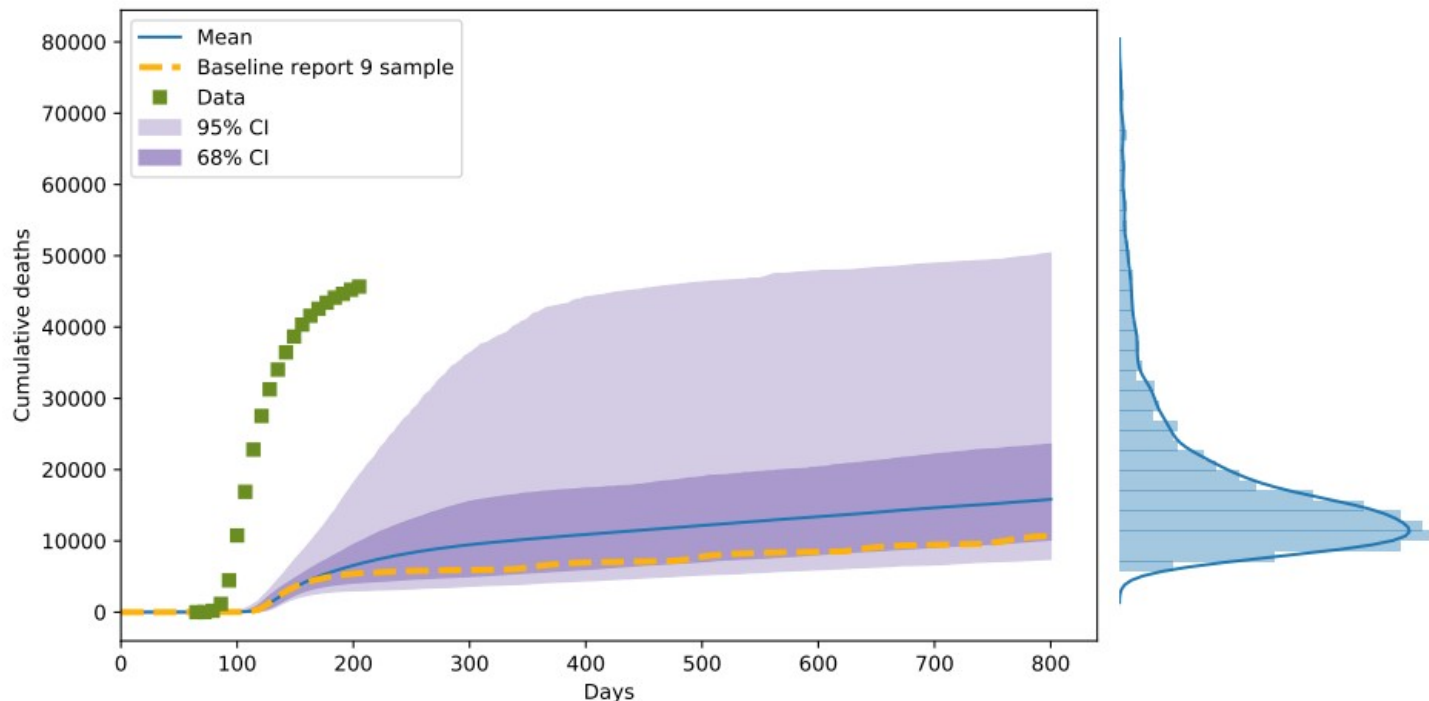


- After 60 iterations, 7 / 19 inputs are refined to quadrature order  $> 0$ .
- Each iteration: new EasyVVUQ ensemble of CovidSim samples.
  - Used **FabSim3** to submit **EasyVVUQ** jobs to the **PSNC Eagle machine**.
  - Each CovidSim sample used 28 cores.

# Results: confidence intervals



- Confidence intervals cumulative deaths, for a given scenario  $\mathcal{S}_1$  from Report 9:
  - $R_0 = 2.4$ , turn NPIs on: weekly new ICU patients  $> 60$ , turn NPIs off  $< 15$ .

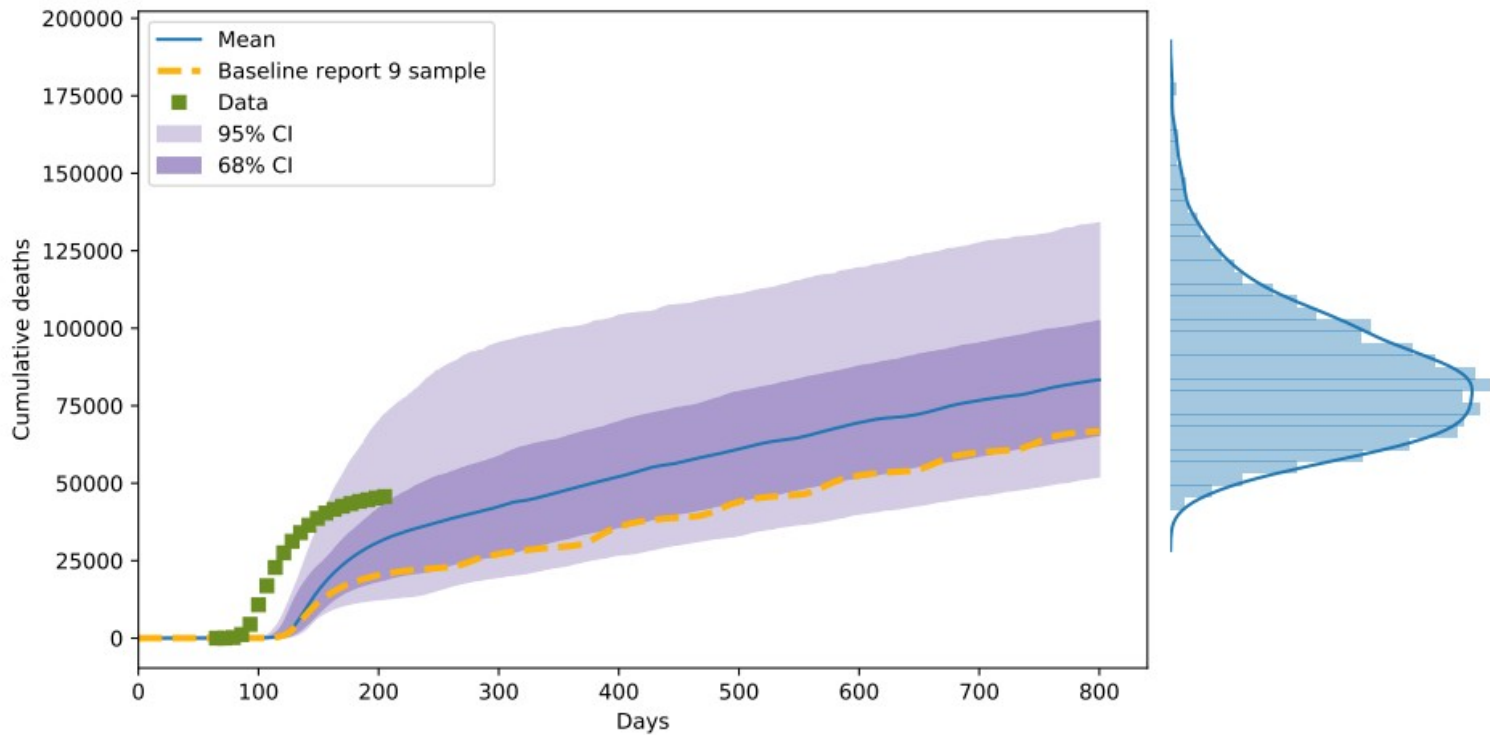


- Despite large variance, does not capture data: uncertainty in initial condition is missing. With post-hoc tuning bias can be fixed [4].

# Results: confidence intervals

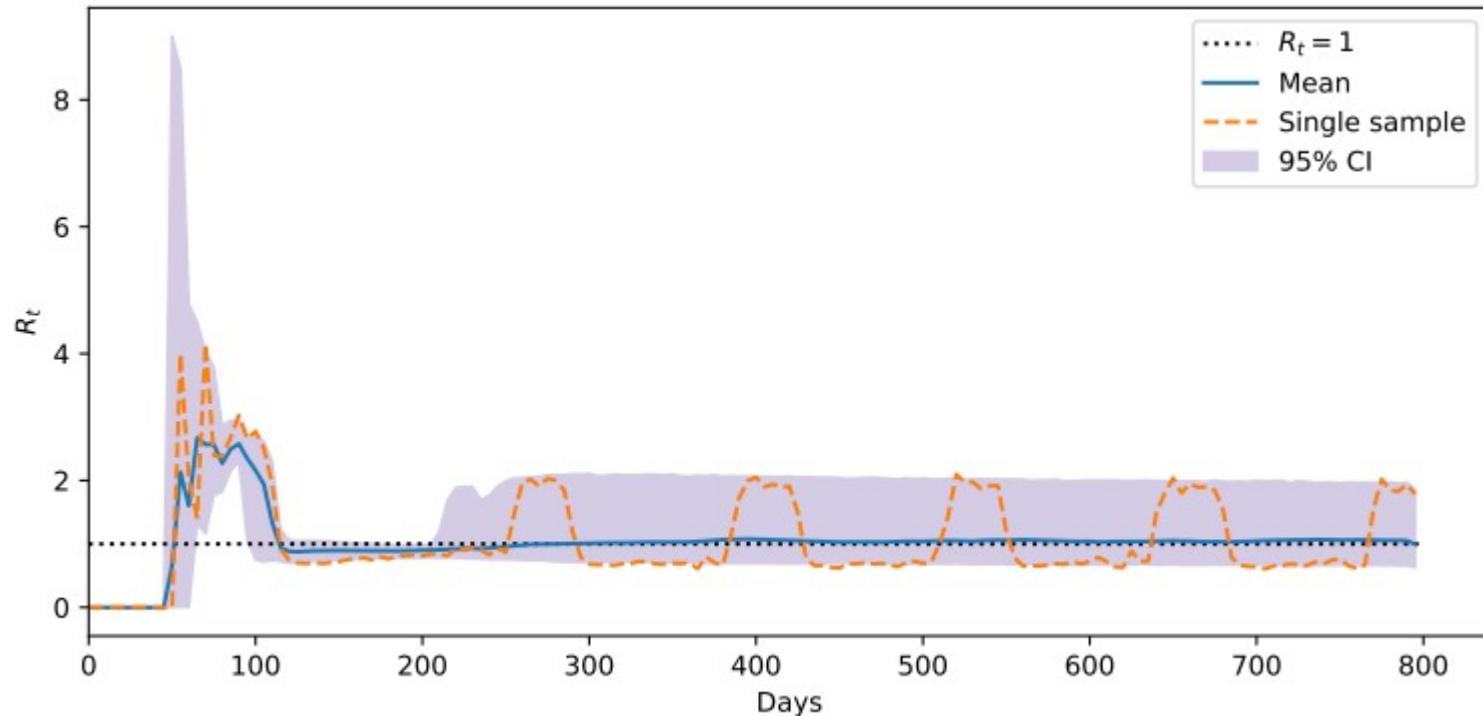


- Confidence intervals, for a **different** scenario  $\mathcal{S}_2$ :
  - $R_0 = 2.6$  , turn NPIs on: weekly new ICU patients  $> 400$ , turn NPIs off  $< 300$ .



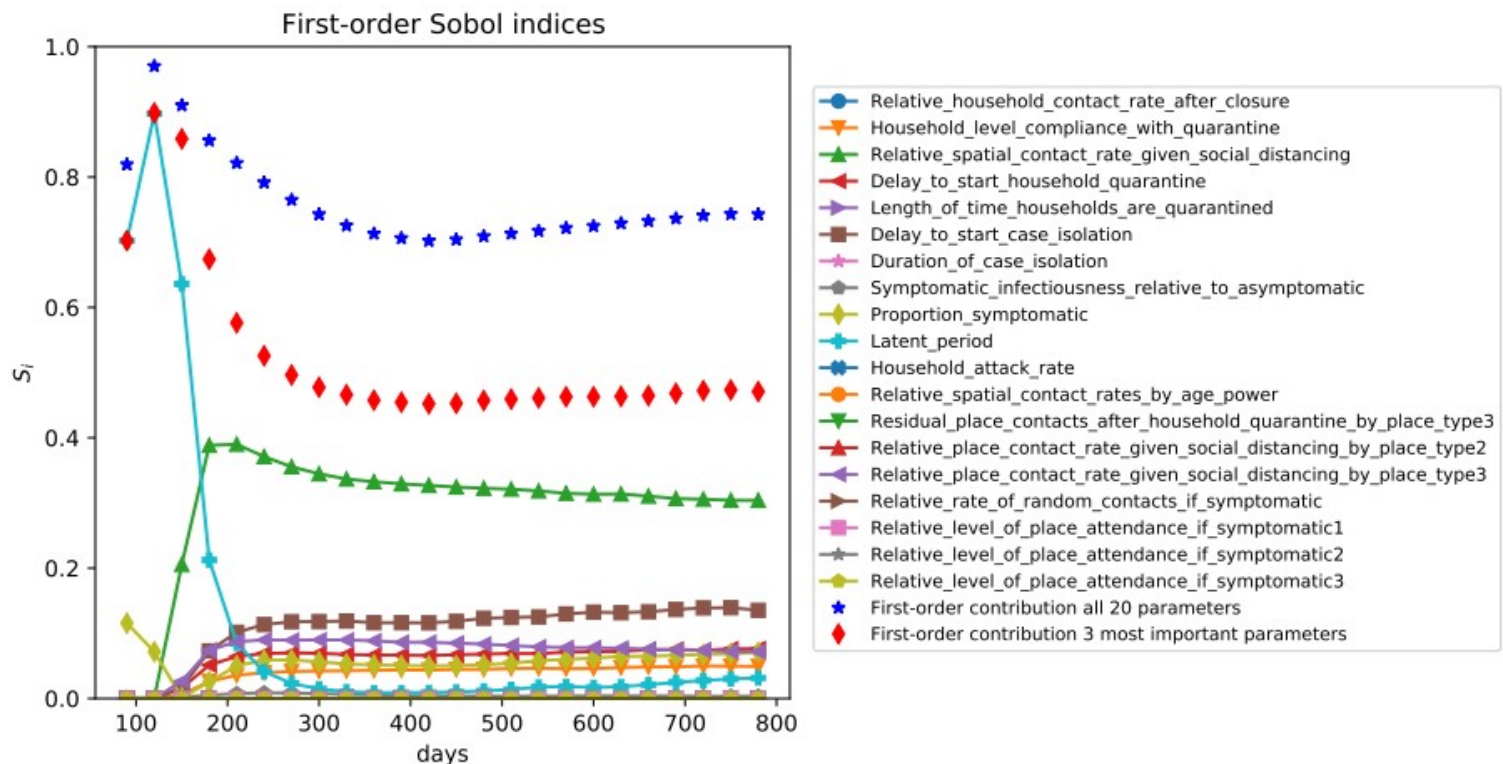
# Results: confidence intervals

- Confidence intervals, for a different QoI:  $R_t$



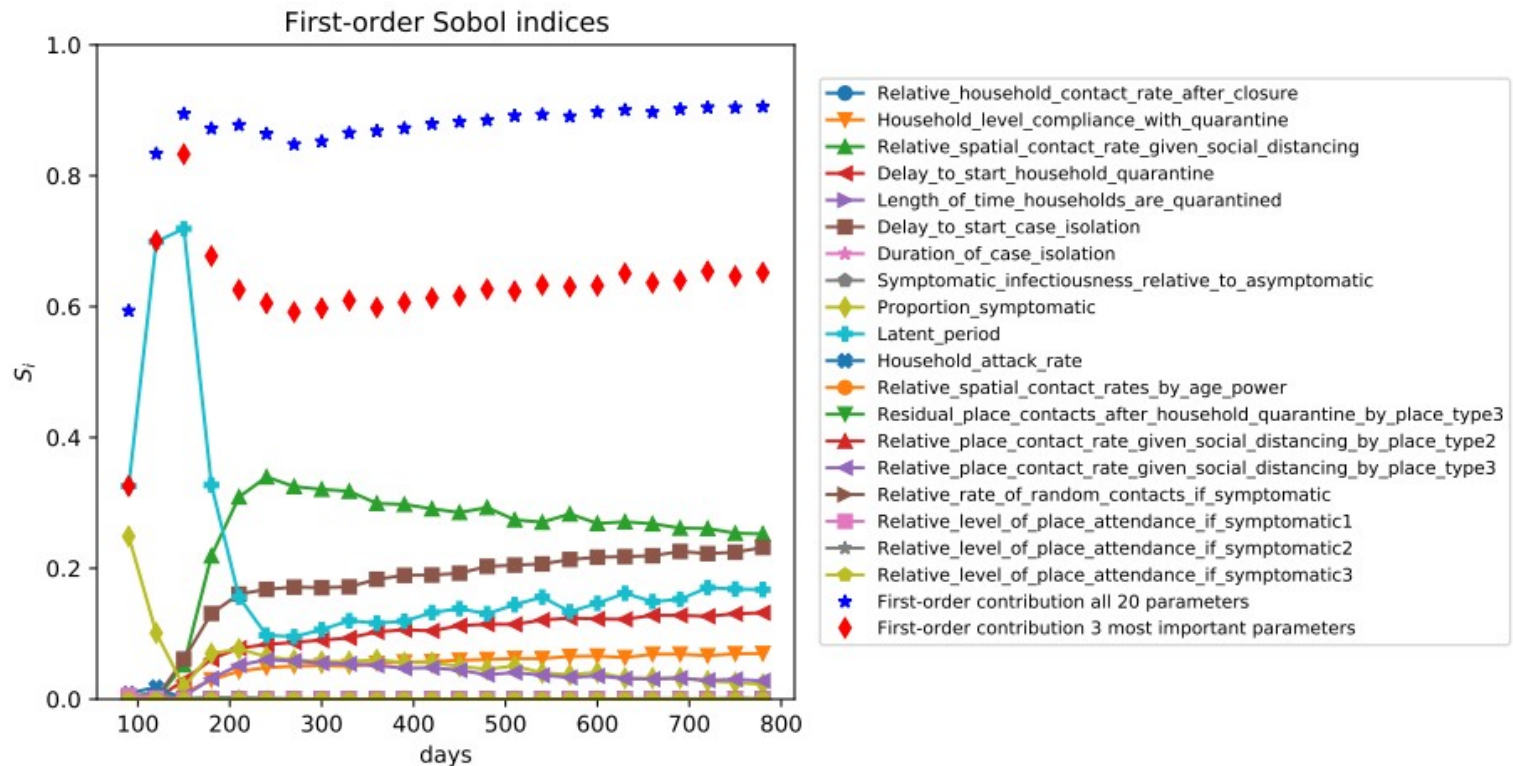
# Results: Sobol indices

- First-order Sobol indices for scenario  $\mathcal{S}_1$  :
  - First-order index: measures **fraction of output variance attributed to each input parameter**. Computed using method of [3].



# Results: Sobol indices

- First-order Sobol indices for scenario  $\mathcal{S}_2$ : qualitatively similar.
  - Overall, NPI-related parameters are most important
  - Note that **just 3 inputs** already cover > 60% of the variance.



## Results: robustness

- Robustness: examine the amplification (or damping) of uncertainty.
- General question: *is the observed output variance large compared to the “amount” of uncertainty injected at the input?*
- Sobol indices do not measure this, we look at the Ratio of the average input – output Coefficients of Variation (CVR)
- CV = standard deviation / mean, a dimensionless measure of variability.

$$CVR := CV(\bar{q}) / CV(\bar{\xi}) = \left( \frac{1}{N} \sum_{n=1}^N \frac{\sigma_{q_n}}{\mu_{q_n}} \right) / \left( \frac{1}{d} \sum_{i=1}^d \frac{\sigma_{\xi_i}}{\mu_{\xi_i}} \right).$$

- CVR > 1: amplification of uncertainty, CVR < 1: damping.

## Results: robustness

- CovidSim results, for scenario  $\mathcal{S}_1$  and  $\mathcal{S}_2$  and cumulative death QoI:

scenario	$CV(\bar{\xi})$	$CV(\bar{q})$	$CVR$
$\mathcal{S}_1$	0.1950	0.6097	3.13
$\mathcal{S}_2$	0.1950	0.3872	1.99

- Both scenarios amplify uncertainty, by a factor of 3 and 2.



# Conclusion



- We performed adaptive parametric UQ on CovidSim.
- It has 940 parameters, which we iteratively brought down to 19.
- Of the 19, just 3 already cover  $> 60\%$  of the output variance.
- It is also important to:
  - Quantify the scenario uncertainty as well.
  - Possibly, use an ensemble of models to estimate the model-choice uncertainty.
  - Investigate calibration / data assimilation.

In case you wish to read the paper:

Edeling, Wouter and Arabnejad, Hamid and Sinclair, Robbie and Suleimenova, Diana and Gopalakrishnan, Krishnakumar and Bosak, Bartosz and Groen, Derek and Mahmood, Imran and Crommelin, Daan and Coveney, Peter V. (2021). *The impact of uncertainty on predictions of the CovidSim epidemiological code*. Nature Computational Science, 1(2), 128-135.

[www.nature.com/articles/s43588-021-00028-9](https://www.nature.com/articles/s43588-021-00028-9)

**nature computational science**

Research also featured in:

- Nature news & views: Kathy Leung, *Quantifying the uncertainty of CovidSim*
- Nature news: David Adam, *Simulating the pandemic: What COVID forecasters can learn from climate models*
- UK science museum: Roger Highfield, *Corona virus: virtual pandemics*

Thank you

