

# Detecting Model Misspecification in Amortized Bayesian Inference with Neural Networks

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Marvin Schmitt<sup>1</sup>, Paul-Christian Bürkner<sup>1,2</sup>, Ullrich Köthe<sup>3</sup>, & Stefan T. Radev<sup>3</sup>

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<sup>1</sup> University of Stuttgart

<sup>2</sup> TU Dortmund University

<sup>3</sup> Heidelberg University

For Jovita

**Do you have a moment to talk about our  
lord and savior**

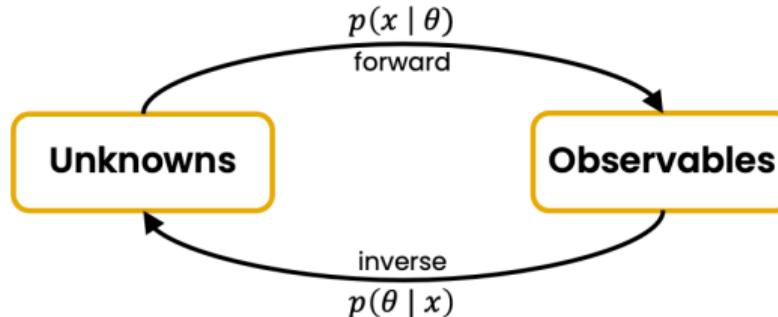
**Do you have a moment to talk about our**

**lord and savior**

**Sir Thomas Bayes?**



# Inverse problems



**Statistical modeling:** **Parameters  $\theta$**

**Data  $x$**

**Epidemiology:** Virus attributes

Infection curve (time series)

**Image processing:** Crisp image

Blurry image

**Physics:** Physical attributes

Gravitation wave measurements

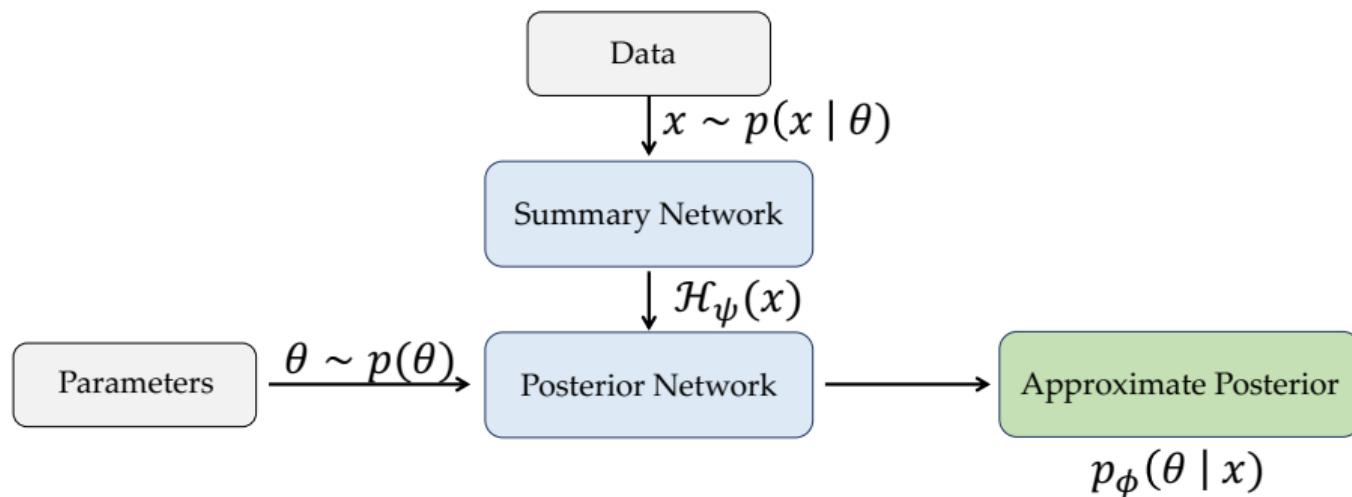
# Bayesian inference

$$\widehat{p(\theta | x)}^{\text{inverse}} = \frac{\overbrace{p(x | \theta) p(\theta)}^{\text{forward}}}{\underbrace{\int p(x, \theta) d\theta}_{= p(x)}}$$



Figure 1: Maybe Thomas Bayes

# Neural posterior estimation (NPE)



# NPE: Optimization objective

The analytic posterior  $p(\boldsymbol{\theta} | \mathbf{x})$  and the approximated posterior  $p_{\phi}(\boldsymbol{\theta} | \mathcal{H}_{\psi}(\mathbf{x}))$  on learned summary statistics  $\mathcal{H}_{\psi}(\mathbf{x})$  shall match:

$$\begin{aligned} (\boldsymbol{\phi}^*, \boldsymbol{\psi}^*) &= \underset{\boldsymbol{\phi}, \boldsymbol{\psi}}{\operatorname{argmin}} \mathbb{E}_{p^*(\mathbf{x})} \left[ \mathbb{KL} \left( p(\boldsymbol{\theta} | \mathbf{x}) \middle\| p_{\phi}(\boldsymbol{\theta} | \mathcal{H}_{\psi}(\mathbf{x})) \right) \right] \\ &= \underset{\boldsymbol{\phi}, \boldsymbol{\psi}}{\operatorname{argmin}} \mathbb{E}_{p^*(\mathbf{x})} \left[ \mathbb{E}_{p(\boldsymbol{\theta} | \mathbf{x})} \left[ -\log p_{\phi}(\boldsymbol{\theta} | \mathcal{H}_{\psi}(\mathbf{x})) \right] \right] \end{aligned}$$

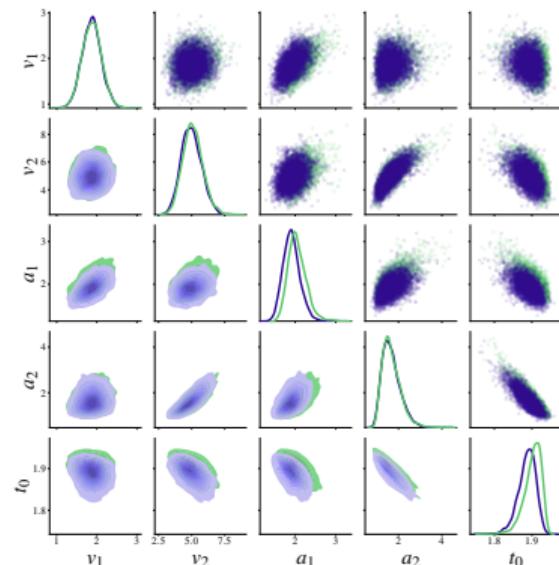
# Out-of-distribution in NPE

Assume that the true data distribution  $p^*(x)$  equals the simulated  $p(x)$ :

$$\begin{aligned}(\phi^*, \psi^*) &= \operatorname{argmin}_{\phi, \psi} \mathbb{E}_{p(\mathbf{x})} \left[ \mathbb{E}_{p(\boldsymbol{\theta} | \mathbf{x})} \left[ -\log p_\phi(\boldsymbol{\theta} | \mathcal{H}_\psi(\mathbf{x})) \right] \right] \\&= \operatorname{argmin}_{\phi, \psi} \mathbb{E}_{p(\mathbf{x}, \boldsymbol{\theta})} \left[ -\log p_\phi(\boldsymbol{\theta} | \mathcal{H}_\psi(\mathbf{x})) \right]\end{aligned}$$

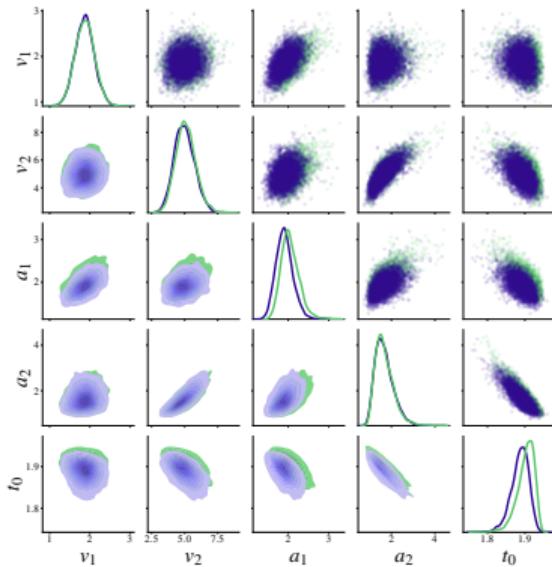
If  $p^*(x) \neq p(x)$ , we optimize with respect to the wrong distribution.

# What happens when MCMC and NPE encounter OOD data?

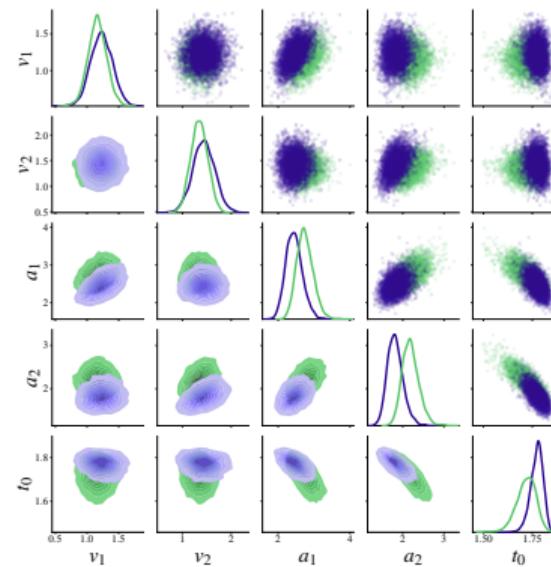


**(a)** well-specified:  $\text{MCMC} \approx \text{NPE}$

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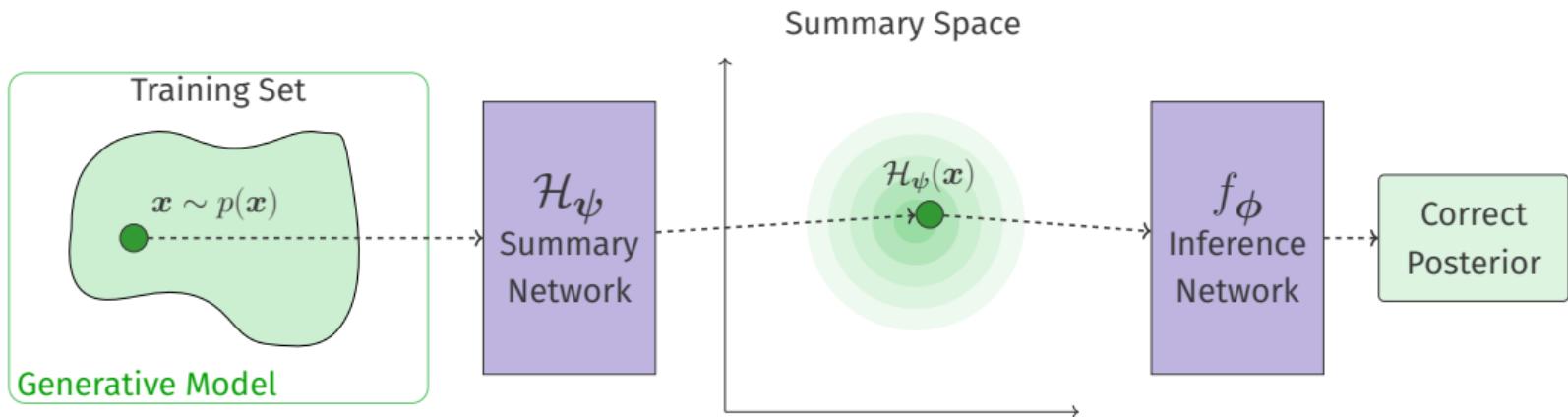


**(a)** well-specified:  $\text{MCMC} \approx \text{NPE}$



**(b)** misspecified:  $\text{MCMC} \neq \text{NPE}$

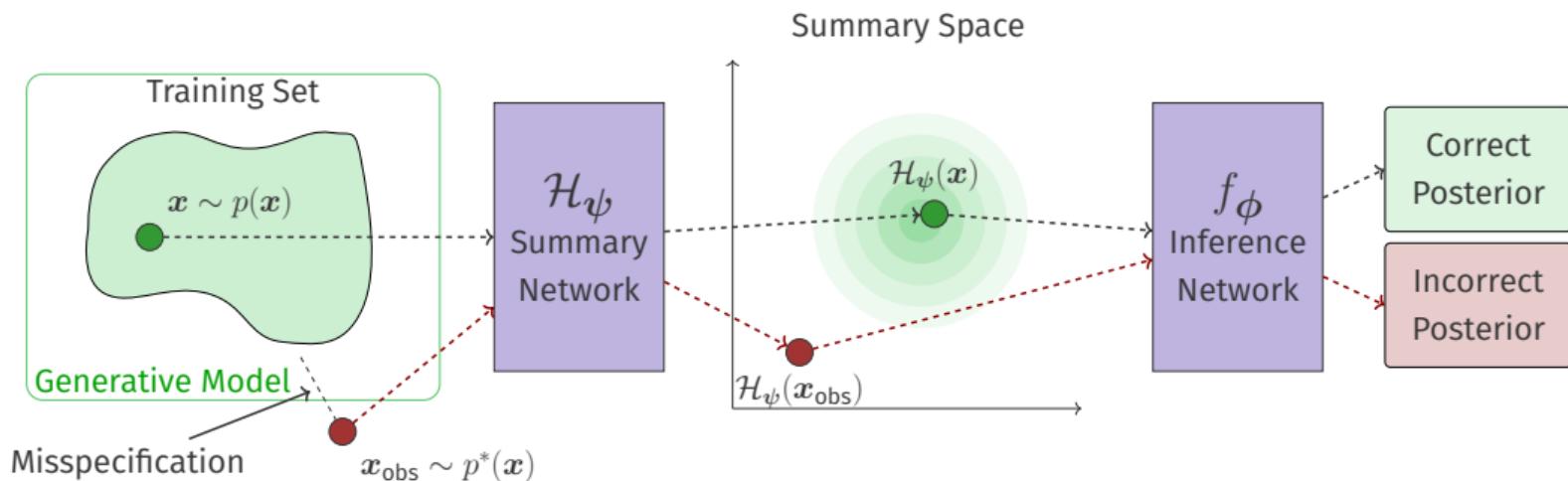
# Structured summary statistics



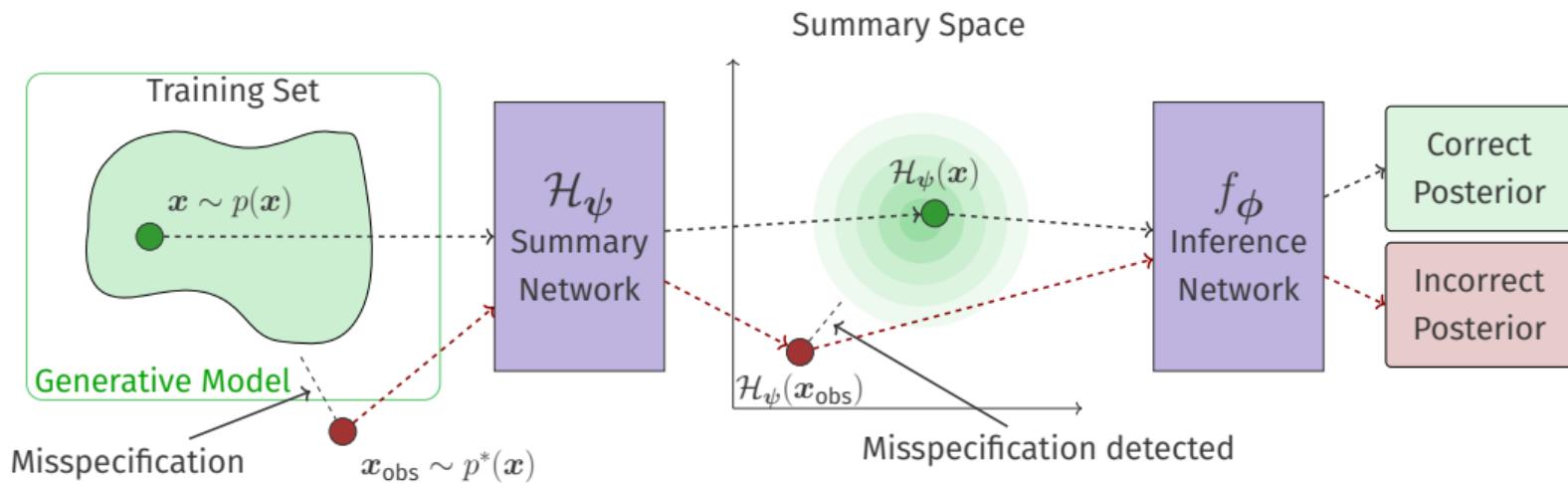
Optimize the summary network's output  $\mathcal{H}_\psi(x)$  towards a unit Gaussian:

$$p(\mathcal{H}_\psi(x)) \approx \mathcal{N}(z | 0, \mathbb{I})$$

# Detecting out-of-distribution data



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Detect via MMD between simulated  $\mathcal{H}_\psi(x)$  and observed  $\mathcal{H}_\psi(x_{\text{obs}})$

# Experiment 1: Gaussian toy model

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# Gaussian: Setup

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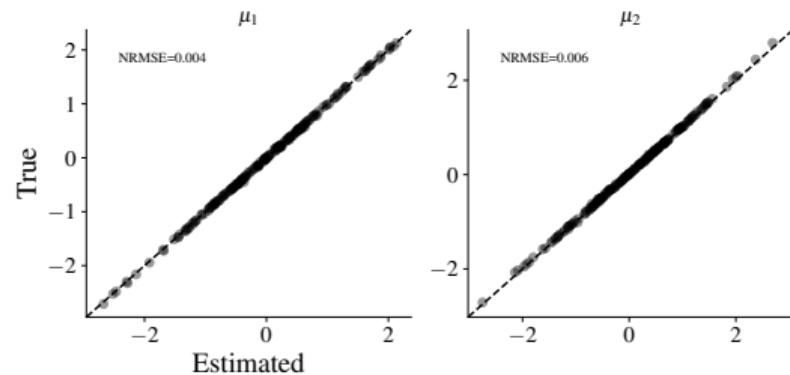
Recover mean vector  $\mu$  of a 2–dimensional spherical Gaussian:

$$\begin{aligned}\mu &\sim \mathcal{N}(\mu | \mu_0, \tau_0 \mathbb{I}) \\ x_k &\sim \mathcal{N}(x | \mu, \tau \mathbb{I}) \quad \text{for } k = 1, \dots, K.\end{aligned}\tag{1}$$

## Potential misspecifications:

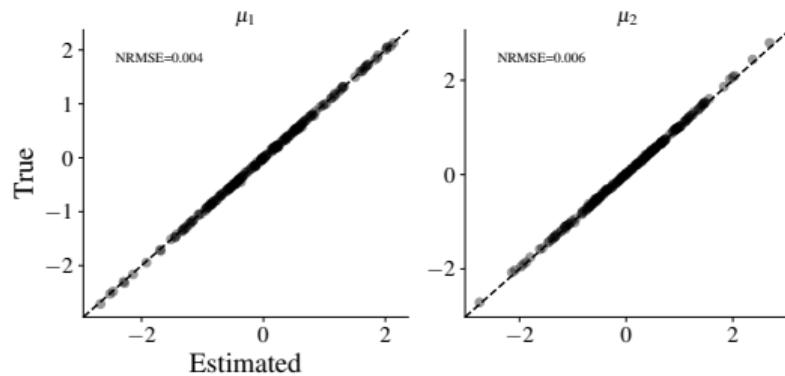
- Prior location  $\mu_0$  and scale  $\tau_0$
- Likelihood scale  $\tau$
- Unmodeled noise

# Gaussian: Perfect performance for well-specified model

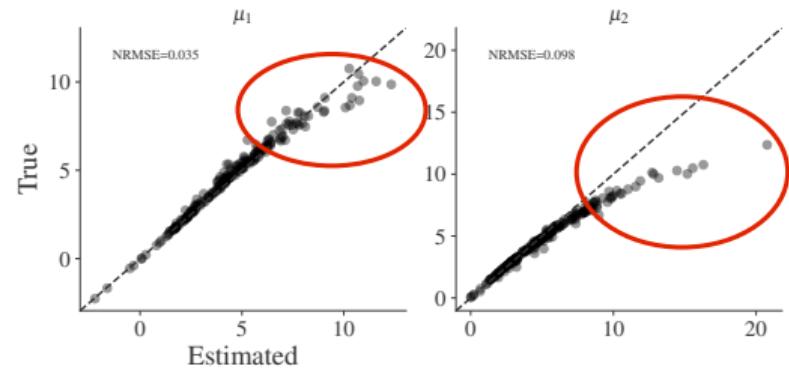


**Figure 3:** Well-specified case

# Gaussian: Perfect performance for well-specified model

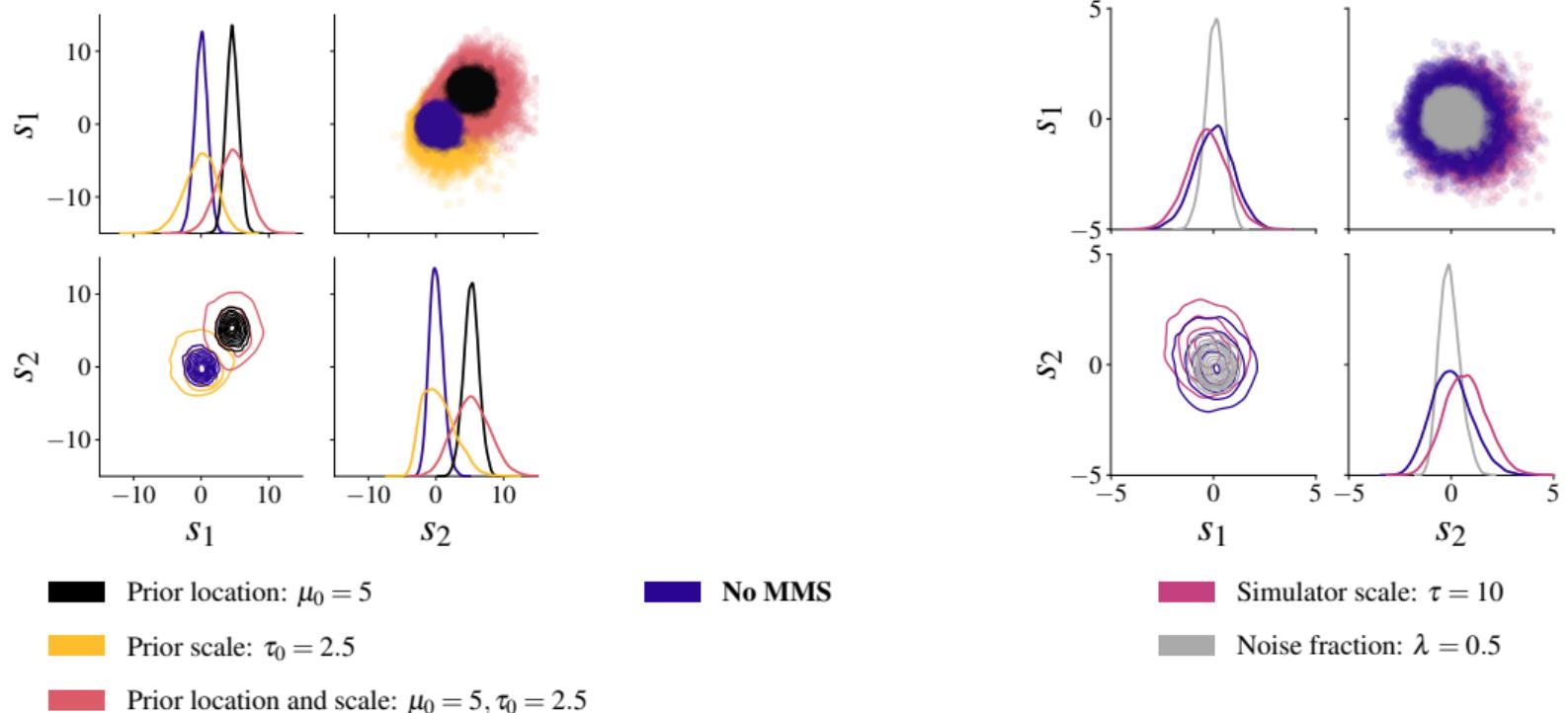


**Figure 3:** Well-specified case

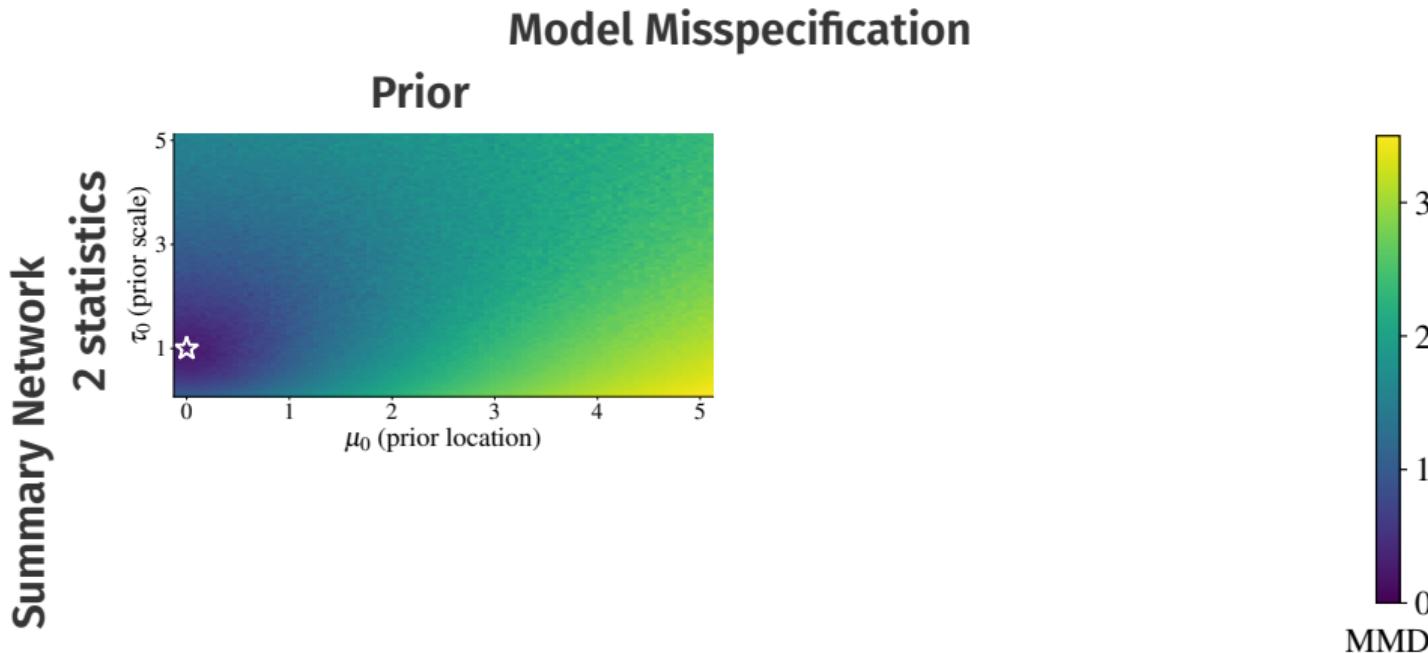


**Figure 4:** Prior misspecification:  $\mu_0 = 2.5$

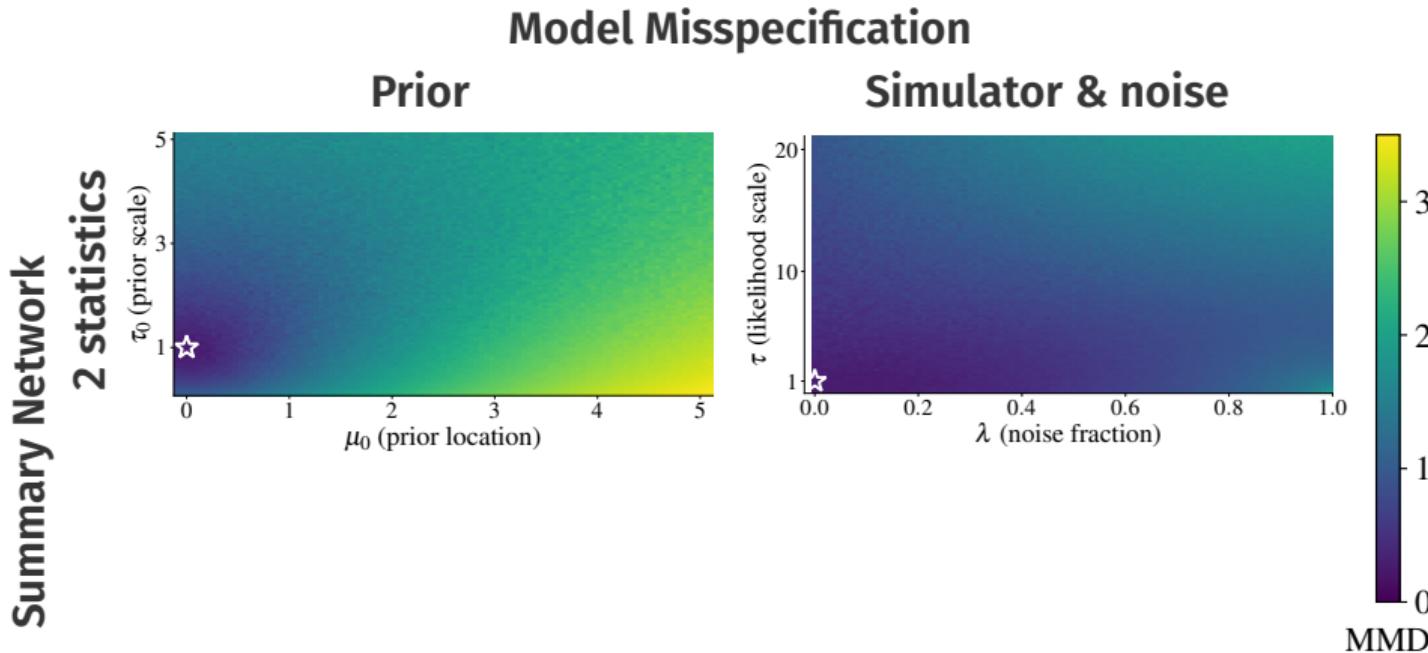
# Gaussian: Inspecting the summary space



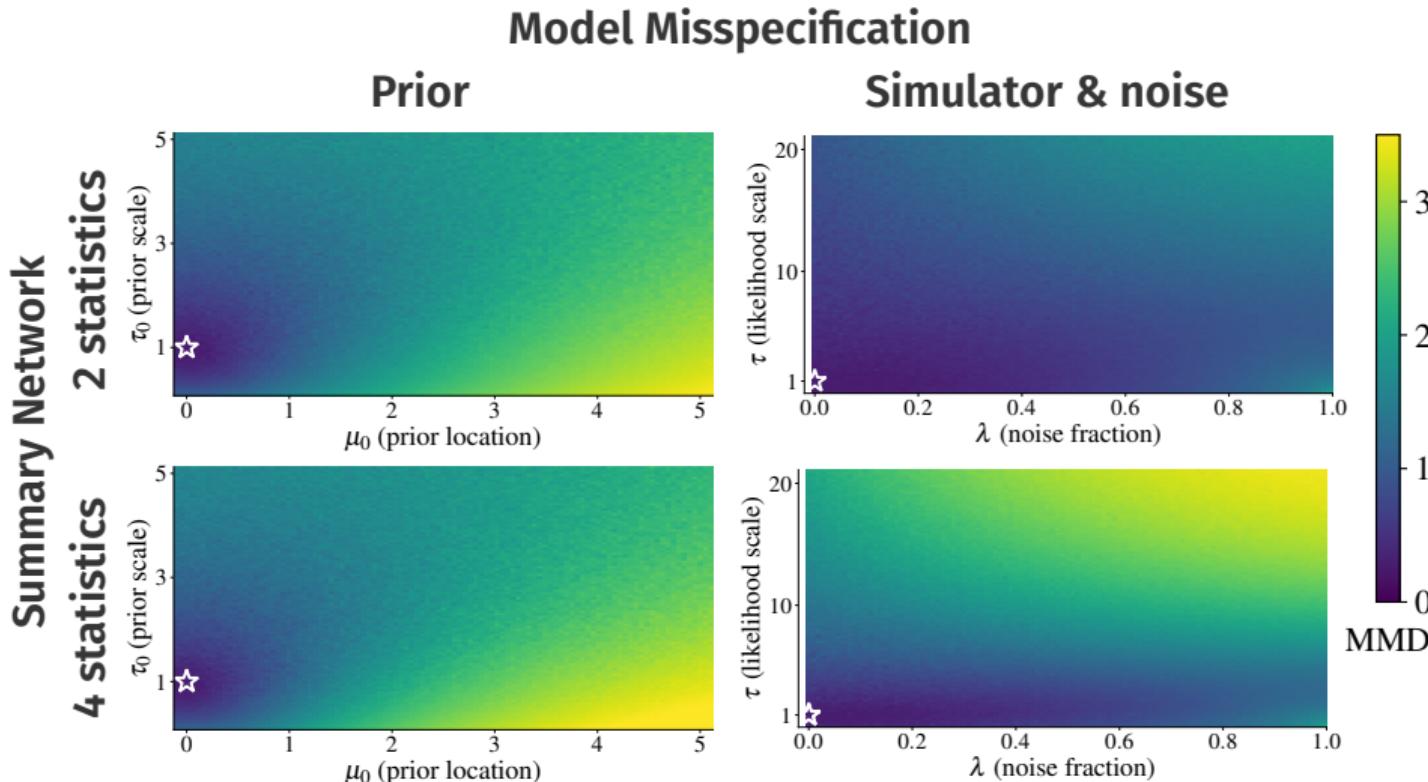
# Gaussian: How many summary statistics?



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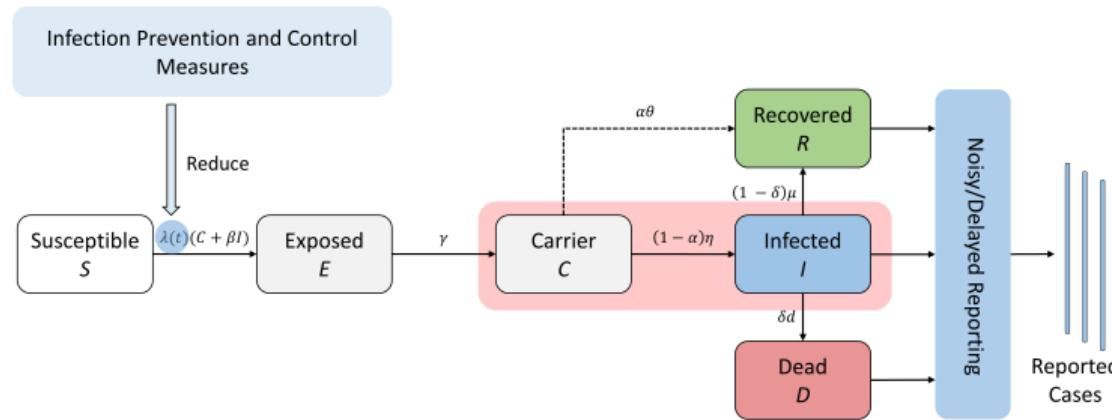
## **Experiment 3: COVID-19 modeling**

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# COVID-19: Motivation

## Compartmental Models for disease outbreaks (Radev et al., 2021)

1. Inference is based on posteriors → must be trustworthy
2. Are initially well-specified models misspecified at some point?



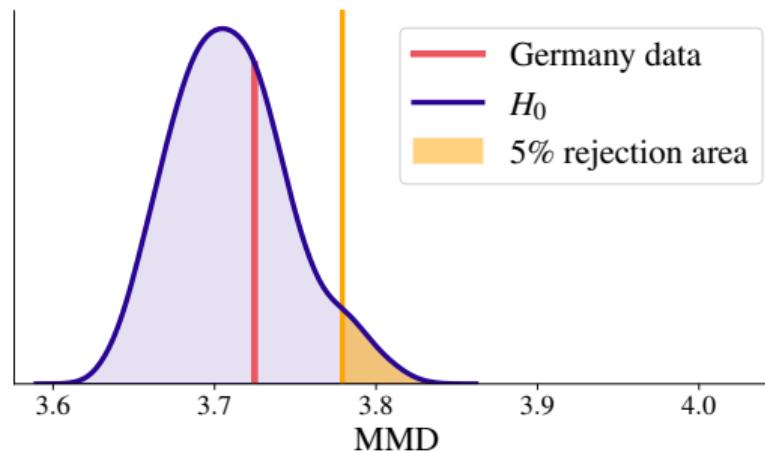
# COVID-19: Setup

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- Train the network on data from the full model  $\mathcal{M}^*$
- Simulate 1000 time series each from
  - $\mathcal{M}^*$ : full model
  - $\mathcal{M}_1$ : no intervention sub-model
  - $\mathcal{M}_2$ : no observation sub-model
- Find discrepancies in the latent summary space

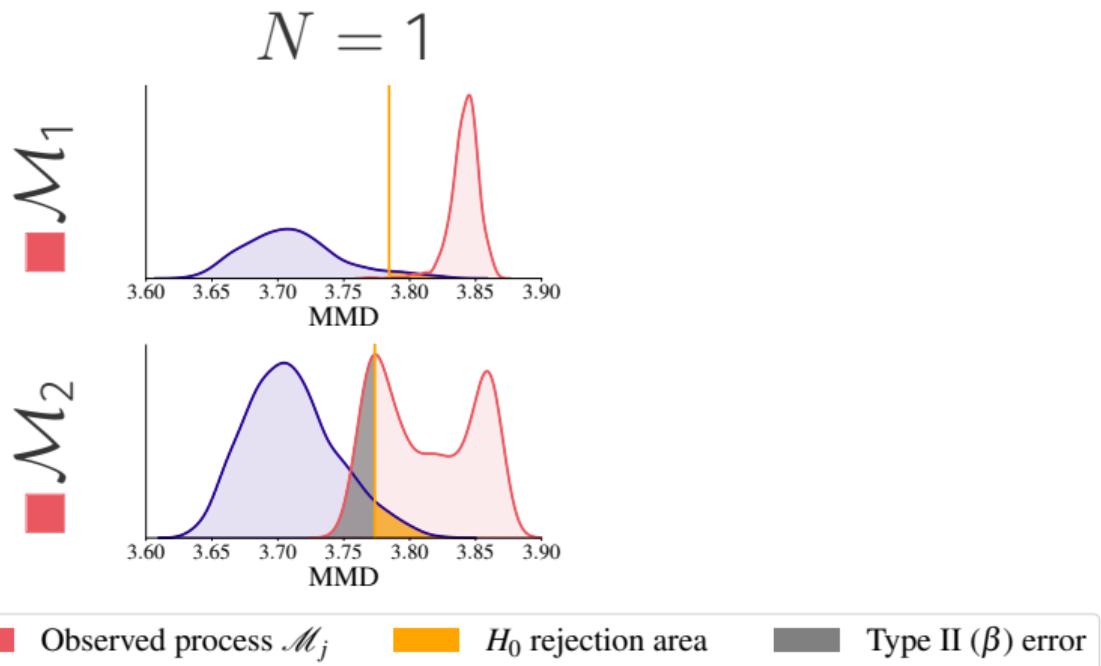
# COVID-19: Is the model well-specified for German data?

Frequentist hypothesis test:  $H_0 : p^*(x) = p(x)$     $H_1 : p^*(x) \neq p(x)$

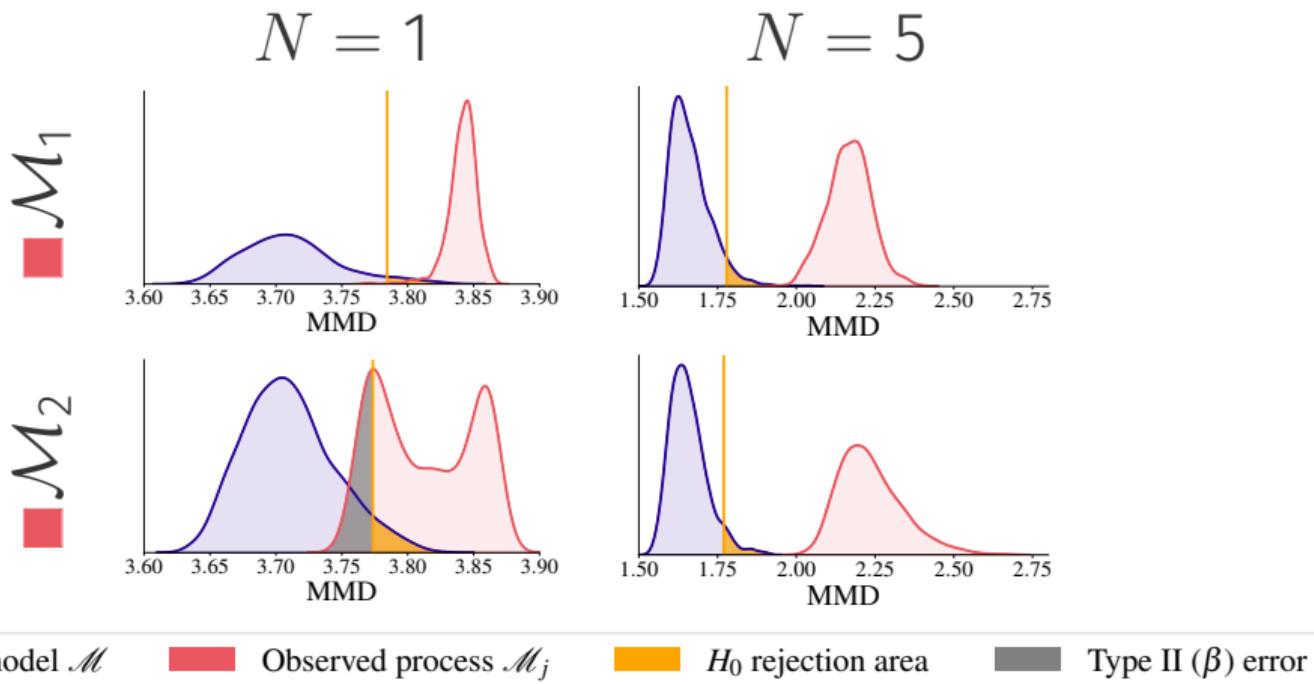


**Conclusion:** Don't reject the null hypothesis → model is well-specified.

# Power of a frequentist hypothesis test on summary space MMD



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  - Bring your own distance metric
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- Unhappy with MMD or the frequentist hypothesis test?
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- All implementations in the *BayesFlow* library: [bayesflow.org](http://bayesflow.org)

# Contact



**Marvin Schmitt**  
 @MarvinSchmittML



**Paul Bürkner**  
 @paulbuerkner



**Ullrich Köthe**  
R<sup>G</sup> Ullrich Köthe



**Stefan Radev**  
 @stefanradev13

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