

On the identifiability of kernels for Population Balance Modelling

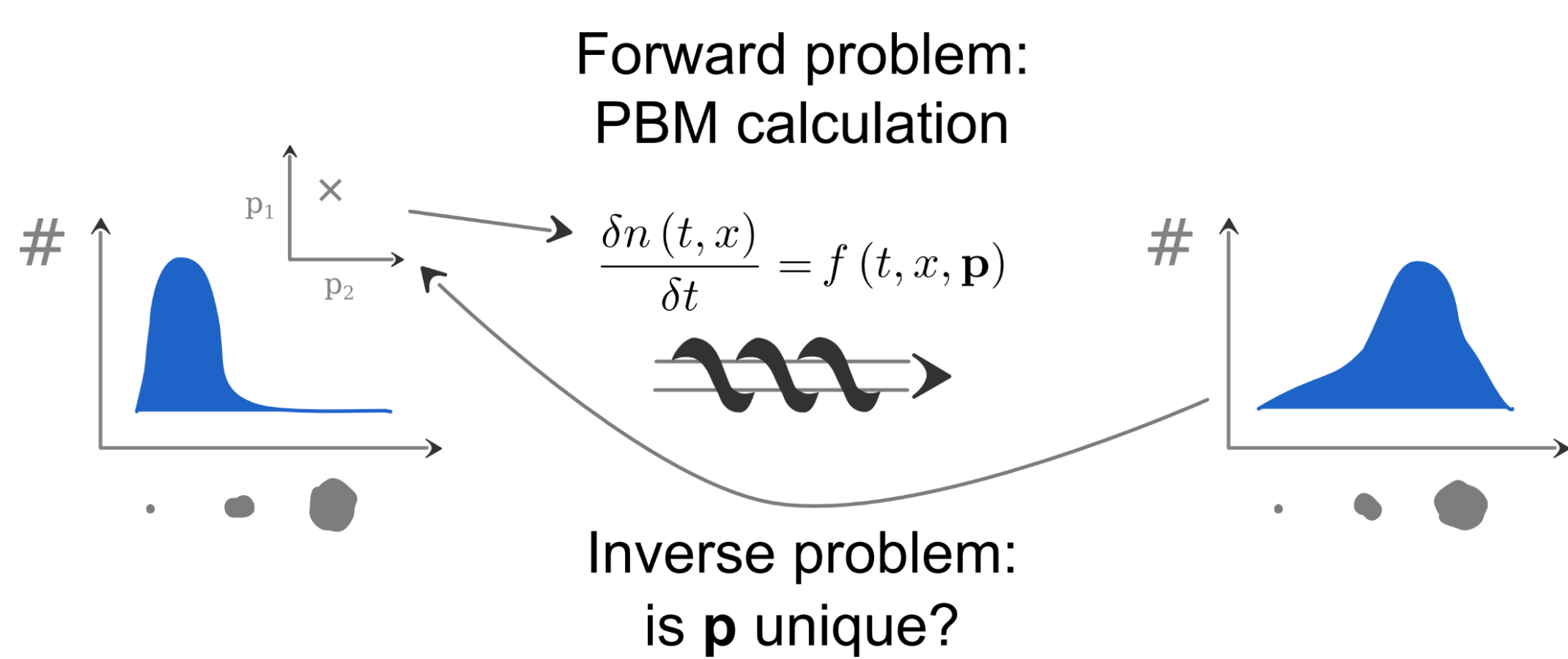
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1. Why identifiability?



- Identifiability is the inverse modelling problem:

$$f(t, x, \mathbf{p}) = f(t, x, \mathbf{q}) \implies \mathbf{p} = \mathbf{q}$$

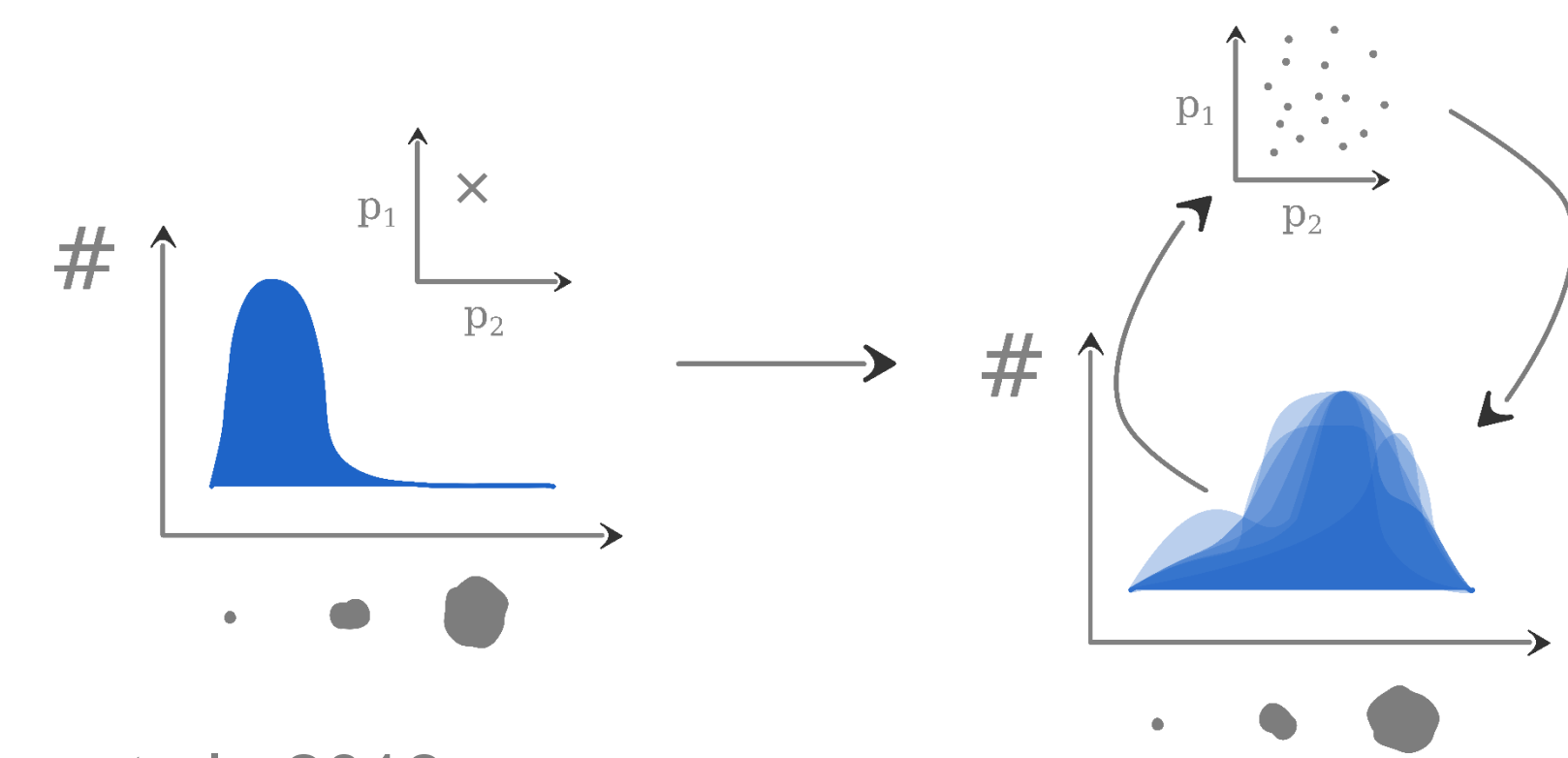
- Important when parameters are linked to physical mechanisms (e.g. machine/operational settings)
- Important for broader application of model

2. How is it calculated for PBM?

- Analytical methods: not applicable
- Most suitable method: repeated numerical parameter estimations *

$$\hat{\mathbf{p}} = \underset{\mathbf{p}}{\operatorname{argmin}} \operatorname{RMSE}(y, f(t, x, \mathbf{p}))$$

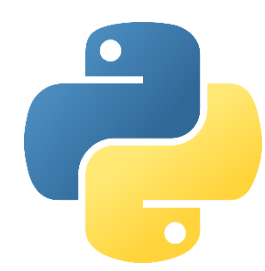
- Practical identifiability: synthetic data with relative Gaussian noise



* reference: Tuncer et al., 2018

3. Focus of this poster

$$\frac{\delta n(t, x)}{\delta t} = \frac{1}{2} \int_0^x \beta(t, x - \varepsilon, \varepsilon) n(t, x - \varepsilon) n(t, \varepsilon) d\varepsilon - n(t, x) \int_0^\infty \beta(t, x, \varepsilon) n(t, \varepsilon) d\varepsilon + \int_x^\infty b(t, x, \varepsilon) S(t, \varepsilon) n(t, \varepsilon) d\varepsilon - S(t, x) n(t, x)$$



- Implementation in Python using Cell Average Technique
- This poster: pure aggregation process, two kernels are assessed:

$$\beta = \beta_0 \cdot (x - \varepsilon)^{\frac{1}{3}} \cdot \varepsilon^{\frac{1}{3}}$$

$$\beta = \beta_0 \cdot (x - \varepsilon)^{\frac{1}{3}} \cdot \varepsilon^{\frac{1}{3}} \cdot (1 + 0.5(\text{step} - 1) \cdot (1 + \tanh(R - \sqrt{(x - \varepsilon)^2 + \varepsilon^2})))$$

4. Results (part 1)

- Analysis of optimization results: average relative error of estimated parameters:

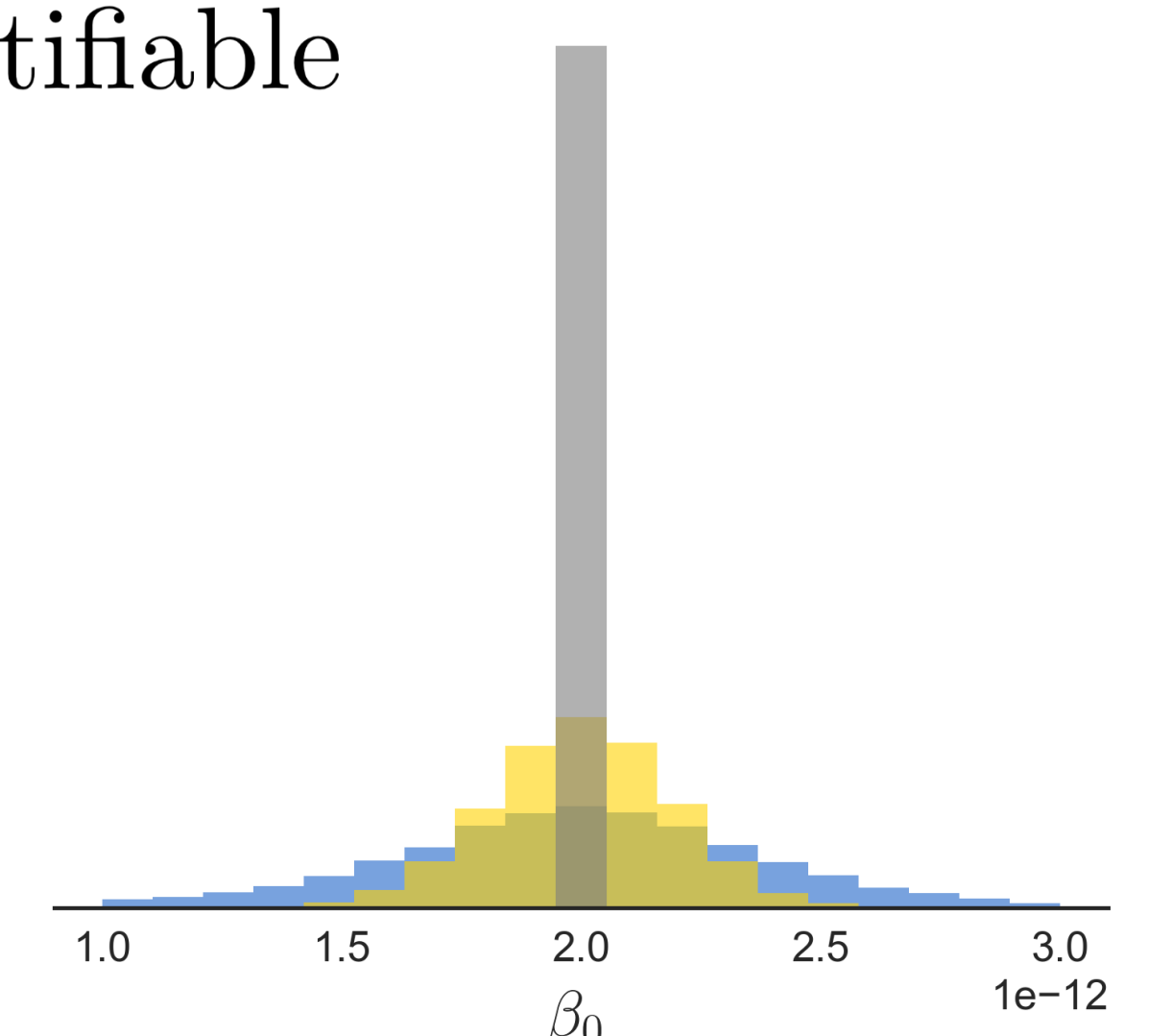
$$\operatorname{ARE}(\hat{p}^k) = 100\% \frac{1}{M} \sum_{j=1}^M \frac{|p^k - \hat{p}_j^k|}{p^k}$$

$$\operatorname{ARE} < \sigma_{\text{noise}} \implies \text{identifiable}$$

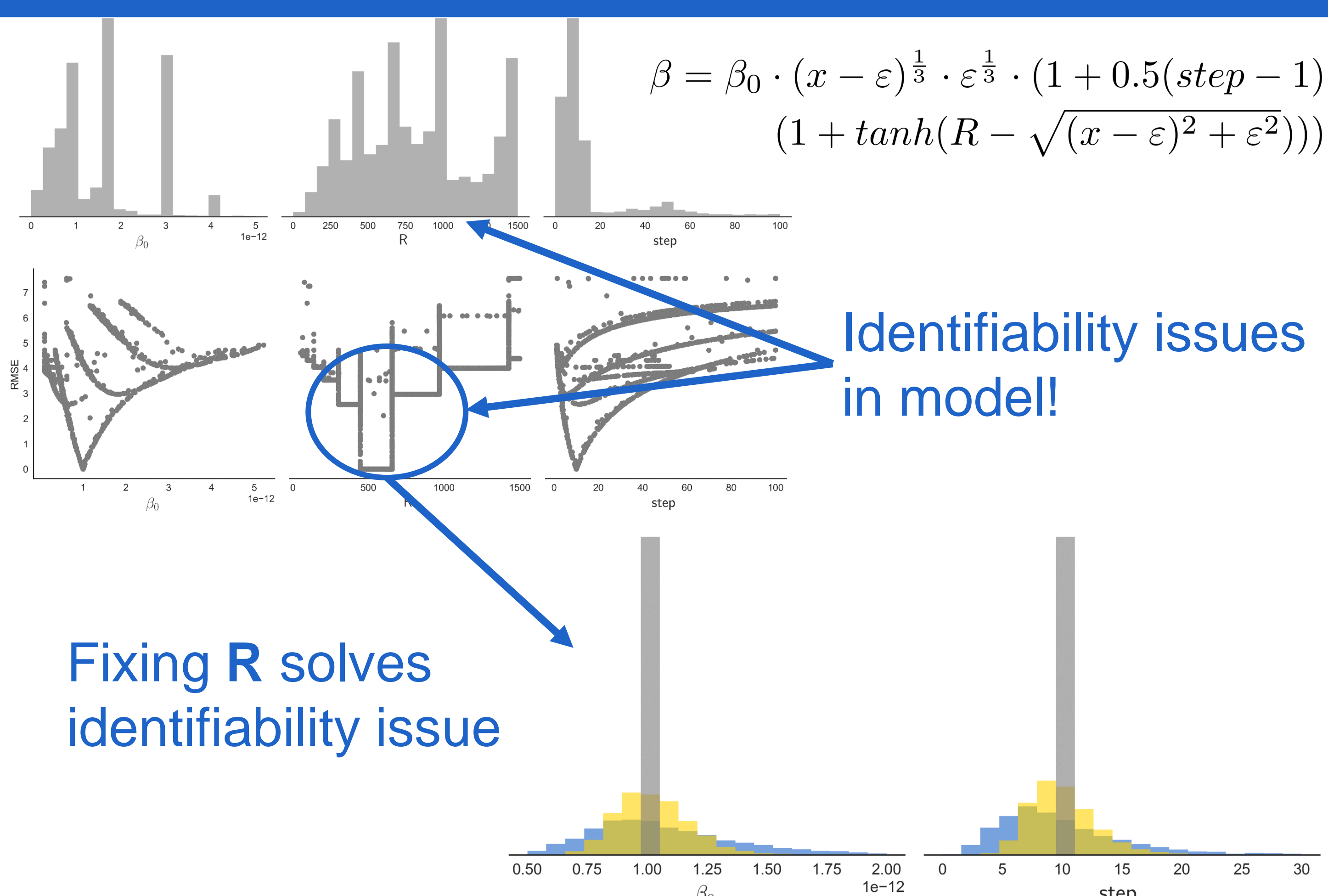
- Product kernel: structurally and practically identifiable

$$\beta = \beta_0 \cdot (x - \varepsilon)^{\frac{1}{3}} \cdot \varepsilon^{\frac{1}{3}}$$

- structural
- practical (25% relative error)
- practical (50% relative error)



5. Results (part 2)



6. Conclusion and prospects

- The used method is elegant, widely applicable, and yields good results
- Analysis for new kernels can be performed in a short timeframe
- It is possible to assess and improve kernel structures
- Future work will include more kernels, and combinations of mechanisms: aggregation and breakage
- Other identifiability techniques can be explored (e.g. profile likelihood)