Self-supervised learning for Bayesian experimental design

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### Contents

Research objective

Two main goals: inference and experimental design Tasks are computationally intractable for simulator models

Self-supervised learning to deal with intractability Link to logistic regression and Jensen-Shannon divergence Technical challenge: the density-chasm problem

Application to Bayesian experimental design Via self-supervised learning of density ratios Exploiting bounds to increase computational efficiency

# DALL·E's visual summary of the talk



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# Overall goal

- Goal: Understanding properties of some data source
- Enables predictions, decision making under uncertainty, ....



### Two fundamental tasks

- Inference task : Given x<sub>o</sub>, what can we robustly say about the properties of the source?
- Experimental design task : How to obtain a x<sub>o</sub> that is maximally useful for learning about the properties?



### Example: stochastic SIR model

- Stochastic model describing the population of susceptibles  $S(\tau)$ , infected  $I(\tau)$  and recovered  $R(\tau)$  as a function of time.
- Parameters  $\theta$ : rate of infection  $\beta$  and the rate of recovery  $\gamma$ .
- Inference task : determine plausible values of β and γ given some measurements of the population sizes.
- Exp design task : find the optimal times at which to perform the measurements to most accurately estimate  $\beta$  and  $\gamma$ .



<sup>(</sup>Figure by Steven Kleinegesse)

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- **•** Bayesian inference of  $\theta$  for data  $\mathbf{x}_o$  obtained with design  $\mathbf{d}_o$ :

$$p(\boldsymbol{\theta}|\mathbf{x}, \mathbf{d}) = \frac{p(\mathbf{x}|\boldsymbol{\theta}, \mathbf{d})}{p(\mathbf{x}|\mathbf{d})} p(\boldsymbol{\theta}|\mathbf{d})$$
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with  $\mathbf{x}$  fixed to  $\mathbf{x}_o$  and  $\mathbf{d}$  to  $\mathbf{d}_o$ .

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Experimental design by maximising mutual information (MI) between data x and parameters θ:

$$\hat{\mathbf{d}} = \operatorname*{argmax}_{\mathbf{d}} \mathsf{MI}_{\mathbf{d}}(\mathbf{x}, \boldsymbol{\theta}) \tag{2}$$

$$\mathsf{MI}_{\mathbf{d}}(\mathbf{x}, \boldsymbol{\theta}) = \mathbb{E}_{p(\mathbf{x}|\mathbf{d})} \mathsf{KL}\left(p(\boldsymbol{\theta}|\mathbf{x}, \mathbf{d}) || p(\boldsymbol{\theta}|\mathbf{d})\right)$$
(3)

$$= \mathbb{E}_{\rho(\mathbf{x},\theta|\mathbf{d})} \log \left[ \frac{\rho(\mathbf{x}|\theta,\mathbf{d})}{\rho(\mathbf{x}|\mathbf{d})} \right]$$
(4)

## Simulator models

- Not all models are specified as family of pdfs.
- We consider here the important class of simulator models: models that are specified via a parameterised stochastic mechanism for generating data



DALL·E's view on simulator models

$$\mathbf{x} = g(\boldsymbol{\theta}, \mathbf{d}, \boldsymbol{\omega}), \quad \boldsymbol{\omega} \sim p(\boldsymbol{\omega})$$
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Maps params heta, design variables **d**, and "noise"  $\omega$  to data **x** 

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- Function g is not known in closed form but implemented as a (complex) computer programme.
- No closed form expression for  $p(\mathbf{x}|\boldsymbol{\theta}, \mathbf{d})$  available
- Sampling data  $\mathbf{x}|\boldsymbol{\theta}, \mathbf{d} \sim p(\mathbf{x}|\boldsymbol{\theta}, \mathbf{d})$  is possible

# Simulator models are widely used in the natural sciences

- Evolutionary biology: to model evolution
- Biochemistry: to model gene expression
- Neuroscience: to model neural processing
- Cognitive sciences: to model human decision making
- Epidemiology: to model the spread of an infectious disease



Simulated neural activity in rat somatosensory cortex (Figure from https://bbp.epfl.ch/nmc-portal)

Simulator models have great modelling power.

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#### Research objective:

Develop efficient tools for Bayesian inference and experimental design with simulator models.

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### Self-supervised learning to deal with intractability Link to logistic regression and Jensen-Shannon divergence Technical challenge: the density-chasm problem

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Link to (log) density ratio estimation

$$\underbrace{\log p_b}_{\text{reference}} + \underbrace{\log p_a - \log p_b}_{\text{difference}} \Rightarrow \underbrace{\log p_a}_{\text{interest}}$$

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- ▶ Let  $\{\mathbf{x}_1, \ldots, \mathbf{x}_n\}$  be the data of interest,  $\mathbf{x}_i \sim p$  (iid), and  $\{\mathbf{y}_1, \ldots, \mathbf{y}_m\}$  be the reference data,  $\mathbf{y}_i \sim q$  (iid).

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- Label the data: (x<sub>i</sub>, 1), (y<sub>i</sub>, 0) and minimise the (rescaled) logistic loss J(h)

$$(h) = \frac{1}{n} \sum_{i=1}^{n} \log \left[ 1 + \nu \exp(-h(\mathbf{x}_i)) \right] + \frac{\nu}{m} \sum_{i=1}^{m} \log \left[ 1 + \frac{1}{\nu} \exp(h(\mathbf{y}_i)) \right]$$
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where  $\nu = n/m$  and *h* is a nonlinearity (e.g. neural network) that we learn.

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where  $\nu = n/m$  and h is a nonlinearity (e.g. neural network) that we learn.

For large sample sizes n and m (and fixed ratio ν), the optimal h is

$$h^* = \log p - \log q \tag{11}$$

Two key points:

1. The optimisation is done without any constraints (e.g. normalisation). The optimal *h* is automagically the ratio between two *densities* 

$$h^* = \log p - \log q \tag{12}$$

2. We only need samples from p and q; we do not need their densities or a model of them (but we do need an appropriate model for the ratio)

▶ For large sample sizes *n* and *m*,  $J(h) \rightarrow \overline{J}(h)$  and the corresponding minimal loss is

$$\begin{split} \bar{J}(h^*) &= \mathbb{E}_{\mathbf{x} \sim p} \log \left[ 1 + \nu \frac{q(\mathbf{x})}{p(\mathbf{x})} \right] + \nu \mathbb{E}_{\mathbf{y} \sim q} \log \left[ 1 + \frac{p(\mathbf{y})}{\nu q(\mathbf{y})} \right] \\ &= -\mathbb{E}_{\mathbf{x} \sim p} \log \left[ \frac{p(\mathbf{x})}{p(\mathbf{x}) + \nu q(\mathbf{x})} \right] - \nu \mathbb{E}_{\mathbf{y} \sim q} \log \left[ \frac{\nu q(\mathbf{y})}{p(\mathbf{y}) + \nu q(\mathbf{y})} \right] \end{split}$$
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► For 
$$\nu = 1$$
 and introducing  $m = (p + q)/2$   
 $\overline{J}(h^*) = -KL(p||m) - KL(q||m) + 2\log 2$  (14)  
 $= -2JSD(p,q) + 2\log 2$  (15)
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Since we are minimising the loss  $\overline{J}(h)$ , we have

$$\overline{J}(h) \ge -2JSD(p,q) + 2\log 2 \tag{16}$$

Rearranging, we obtain

$$JSD(p,q) \ge \log 2 - \frac{1}{2}\bar{J}(h)$$
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- Contrastive learning via classification with the logistic loss corresponds to estimating the Jensen-Shannon divergence (JSD) between p and q.
- For a review paper on statistical applications of contrastive learning, see Gutmann, Kleinegesse, and Rhodes, Behaviormetrika, 2022.

In the following, I will focus on the logistic loss as done in our early work on contrastive learning for the estimation of unnormalised models, "Noise-contrastive estimation (NCE)" (Gutmann and Hyvärinen, AISTATS 2010).

- In the following, I will focus on the logistic loss as done in our early work on contrastive learning for the estimation of unnormalised models, "Noise-contrastive estimation (NCE)" (Gutmann and Hyvärinen, AISTATS 2010).
- But other loss functions can be used:
  - multinomial logistic loss (Srivastava, et al, TMLR 2023)
  - Bregman divergences (Gutmann and Hirayama, UAI 2011)
  - f-divergences (e.g. Rhodes and Gutmann, AISTATS, 2019)
  - ▶ ...

# Constructing reference data

Choice depends on the specific application of contrastive learning.

- Fit a preliminary model and keep it fixed (as often done in NCE)
- Iterative approach: fitted model becomes reference in the next iteration (as also done in our original work on NCE)
- Use other segments for time series data (Hyvärinen and Morioka, NeurIPS 2016)
- For Bayesian inference, use prior predictive distribution (Thomas et al, 2016; Thomas et al, Bayesian Analysis, 2020)
- Generate it conditionally on observed data (Ceylan and Gutmann, ICML 2018)
- Iterative adaptive approach with generative models: results into GANs (Goodfellow et al, NeurIPS 2014)
- Iterative adaptive approach with flexible density model such as flows ("Flow-contrastive estimation", Gao et al, NeurIPS 2019)

#### Is there an optimal reference ("noise") distribution?

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For parameter estimation, see the paper *The Optimal Noise in Noise-Contrastive Learning Is Not What You Think* by Omar Chehab, Alex Gramfort, Aapo Hyvarinen, at UAI, 2022.

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- Problem: Single ratio methods are sample inefficient if the two distributions are very different ("density chasm")
- Consider ratio between two zero-mean Gaussians. 10'000 samples from each distribution. Ratio parametrised by θ ∈ ℝ.
- Solution in red bridges the "gap" using telescopic ratio estimation (TRE) (Rhodes, Xu, and Gutmann, NeurIPS 2020)



A single density-ratio fails to "bridge" the density-chasm.

Let us thus use multiple bridges.



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(relabel  $p \equiv p_0$  and  $q \equiv p_4$ ) and compute *telescoping* product

$$\frac{p_0(\mathbf{x})}{p_4(\mathbf{x})} = \frac{p_0(\mathbf{x})}{p_1(\mathbf{x})} \frac{p_1(\mathbf{x})}{p_2(\mathbf{x})} \frac{p_2(\mathbf{x})}{p_3(\mathbf{x})} \frac{p_3(\mathbf{x})}{p_4(\mathbf{x})}.$$
(19)

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Sample efficiency curves for the 1d peaked ratio experiment.

More results in the paper!



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For further improvements, see "Estimating the Density Ratio between Distributions with High Discrepancy using Multinomial Logistic Regression", Srivastava et al, TMLR 2023.

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(Kleinegesse and Gutmann, AISTATS 2019; ICML 2020; arXiv:2105.04379) (Kleinegesse, Drovandi and Gutmann, Bayesian Analysis 2020)

Example: Stochastic SIR model with noisy observations Latent process: Susceptibles  $\rightarrow$  Infected  $I(t) \rightarrow$  Recovered Observation model:  $y(t)|\theta \sim \text{Poisson}(y;\phi I(t))$ 

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- Parameters  $\boldsymbol{\theta} = (\beta, \gamma)$  (infection rate and recovery rate)

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Experimental design by maximising mutual information (MI)

$$\hat{\mathbf{d}} = \underset{\mathbf{d}}{\operatorname{argmax}} \mathbb{E}_{p(\mathbf{x},\theta|\mathbf{d})} \log \left[ \frac{p(\mathbf{x}|\theta, \mathbf{d})}{p(\mathbf{x}|\mathbf{d})} \right]$$
(20)

Use contrastive self-supervised learning to estimate

$$h_{\mathbf{d}}(\mathbf{x}, \boldsymbol{\theta}) = \log p(\mathbf{x}|\boldsymbol{\theta}, \mathbf{d}) - \log p(\mathbf{x}|\mathbf{d}), \qquad (21)$$

and maximise sample average of  $h_{d}(\mathbf{x}, \theta)$  with respect to **d** 

- Static setting: Kleinegesse and Gutmann, AISTATS 2019
- Sequential setting where we update our belief about θ as we sequentially acquire the data: Kleinegesse, Drovandi and Gutmann, Bayesian Analysis 2020

$$\hat{\mathbf{d}} = \operatorname{argmax}_{\mathbf{d}} \mathbb{E}_{p(\mathbf{x}, \boldsymbol{\theta} | \mathbf{d})} \log \left[ \frac{p(\mathbf{x} | \boldsymbol{\theta}, \mathbf{d})}{p(\mathbf{x} | \mathbf{d})} \right]$$

# Learning the ratio h<sub>d</sub>(x, θ) and approximating the MI is computationally costly.

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- But we do not need to estimate the MI accurately everywhere! Only around it's maximum.

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- Let us use lower bounds on the MI (or proxy) where we concurrently tighten the bound and maximise the (proxy) MI!

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We can (again!) leverage logistic regression.

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- Logistic regression results in replacing the KL divergence with the JSD when measuring the MI.

$$\mathsf{JSD}(p,q;\mathbf{d}) \ge \log 2 - \frac{1}{2}\bar{J}(h;\mathbf{d}) \tag{22}$$

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where *h* is the regression function and  $\overline{J}$  the logistic loss.

Perform experimental design by maximising the negative logistic loss jointly with respect to h and d.

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- ▶ For sequential setting: Ivanova et al, NeurIPS, 2021

#### SIR example: static case (Kleinegesse and Gutmann, ICML 2020)



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# SIR example: sequential case (Ivanova et al, NeurIPS, 2021)



# Conclusions

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## Thank you for your attention!



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