Statistical applications of contrastive learning

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Research presented here is the result of joint work with Aapo Hyvärinen, Jukka Corander, Jun-ichiro Hirayama, Chris Drovandi, and my PhD students Ben Rhodes and Steven Kleinegesse.

- 1. The likelihood function is computationally intractable for energy-based and simulator-based models.
- 2. Contrastive learning is an intuitive and computationally feasible alternative to likelihood-based learning.
- 3. We used it in a broad range of tasks: (1) parameter estimation, (2) Bayesian inference, and (3) Bayesian experimental design.

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Overall goal

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

Two fundamental tasks

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

The likelihood function L(*θ*)

- \blacktriangleright Probability that the model generates data like x° when using parameter value *θ*
- \triangleright Classically, the main workhorse to solve the inference and design task.

The likelihood function L(*θ*)

For models expressed as a family of pdfs $\{p(\mathbf{x}|\theta)\}\)$ indexed by θ : $L(\theta) = p(\mathbf{x}|\theta)$ where **x** is fixed.

 \blacktriangleright Inference:

$$
\hat{\theta} = \underset{\theta}{\arg\max} p(\mathbf{x}|\theta) \quad \text{or} \quad p(\theta|\mathbf{x}) = \frac{p(\mathbf{x}|\theta)}{p(\mathbf{x})} p(\theta) \quad (1)
$$

with **x** fixed to **x**^o.

 \triangleright Experimental design via mutual information: expand model to include (deterministic) design variable **d**, $\{p(\mathbf{x}|\boldsymbol{\theta},\mathbf{d})\}$

$$
\hat{\mathbf{d}} = \underset{\mathbf{d}}{\text{argmax}} \, \text{MI}_{\mathbf{d}}(\mathbf{x}, \theta) \tag{2}
$$

$$
MI_{\mathbf{d}}(\mathbf{x},\boldsymbol{\theta}) = KL\left(p(\boldsymbol{\theta},\mathbf{x}|\mathbf{d})||p(\boldsymbol{\theta}|\mathbf{d})p(\mathbf{x}|\mathbf{d})\right) \tag{3}
$$

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

Energy and simulator-based models

- \triangleright Not all models are specified as family of pdfs.
- \blacktriangleright Two important classes considered here
	- 1. Energy-based (unnormalised) models
	- 2. Simulator-based (implicit) models
- \blacktriangleright The models are rather different, common point:

Multiple integrals needed to be solved to represent the models in terms of pdfs.

- \triangleright Solving the integrals exactly is computationally impossible (curse of dimensionality)
	- \Rightarrow No model pdfs
	- \Rightarrow No standard likelihood-based inference or experimental design

Energy-based models

 \blacktriangleright Widely used:

 \triangleright computer vision and modelling of images \blacktriangleright natural language processing and machine translation \triangleright modelling social or biological networks \blacktriangleright ...

If Specified via an energy function $E(\mathbf{x}; \theta)$ so that $\phi(\mathbf{x}|\boldsymbol{\theta}) = \exp(-E(\mathbf{x}; \boldsymbol{\theta})) \propto p(\mathbf{x}|\boldsymbol{\theta}),$

$$
\int \cdots \int \phi(\mathbf{x}|\boldsymbol{\theta}) d\mathbf{x} = Z(\boldsymbol{\theta}) \neq 1 \qquad \qquad p(\mathbf{x}|\boldsymbol{\theta}) = \frac{\phi(\mathbf{x}|\boldsymbol{\theta})}{Z(\boldsymbol{\theta})}
$$

- Advantage: Specifying an energy $E(\mathbf{x}; \theta)$ is often easier than specifying normalised models
- \triangleright Disadvantage: Integral defining the partition function $Z(\theta)$ can generally not be computed. Model pdf and likelihood function are intractable.

We cannot just ignore the partition function

Consider
$$
p(x; \theta) = \frac{\phi(x; \theta)}{Z(\theta)} = \frac{\exp(-\theta \frac{x^2}{2})}{\sqrt{2\pi/\theta}}
$$

I Log-likelihood function for precision $\theta > 0$

$$
\ell(\theta) = -n \log \sqrt{\frac{2\pi}{\theta}} - \theta \sum_{i=1}^{n} \frac{x_i^2}{2}
$$
 (5)

- \blacktriangleright Data-dependent (blue) and independent part (red) balance each other.
- **Ignoring** $Z(\theta)$ **leads to** meaningless estimates.

Simulator-based models

- \blacktriangleright Widely used:
	- \triangleright computer models/simulators in the natural sciences
	- \blacktriangleright evolutionary biology to model evolution
	- \blacktriangleright epidemiology to model the spread of an infectious disease \blacktriangleright . . .
- \triangleright Specified via a measurable function g, typically not known in closed form but implemented as a computer programme.

$$
\mathbf{x} = g(\boldsymbol{\theta}, \boldsymbol{\omega}), \quad \boldsymbol{\omega} \sim p(\boldsymbol{\omega}) \tag{6}
$$

Maps parameters *θ* and "noise" *ω* to data **x**

- \blacktriangleright Advantage: connects statistics to the natural sciences
- \triangleright Disadvantage: Model pdf and lik function are intractable.

$$
\mathsf{Pr}(\mathbf{x} \in \mathcal{A}|\boldsymbol{\theta}) = \mathsf{Pr}(\{\omega : g(\boldsymbol{\theta}, \boldsymbol{\omega}) \in \mathcal{A}\})
$$

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 \blacktriangleright Link to Bayes' rule

$$
\underbrace{\log p(\theta)}_{\text{reference}} + \underbrace{\log p(\mathbf{x}|\theta) - \log p(\mathbf{x})}_{\text{difference}} \Rightarrow \underbrace{\log p(\theta|\mathbf{x})}_{\text{interest}} \qquad (9)
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- **I** Let $\{x_1, \ldots, x_n\}$ be the data of interest, $x_i \sim p$ (iid), and {**y**1*, . . .* **y**m} be reference data, **y**ⁱ ∼ q (iid).

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- label the data: $(x_i, 1)$, $(y_i, 0)$ and minimise the (rescaled) logistic loss $J(h)$

$$
J(h) = \frac{1}{n} \sum_{i=1}^{n} \log \left[1 + \nu \exp(-h(\mathbf{x}_i)) \right] +
$$

$$
\frac{\nu}{m} \sum_{i=1}^{m} \log \left[1 + \frac{1}{\nu} \exp(h(\mathbf{y}_i)) \right]
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where $\nu = m/n$

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where $\nu = m/n$

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 model of them (but we do need an appropriate model for the ratio)

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

$$
\overline{J}(h^*) = \mathbb{E}_{\mathbf{x} \sim p} \log \left[1 + \nu \frac{q(\mathbf{x})}{p(\mathbf{x})} \right] + \nu \mathbb{E}_{\mathbf{y} \sim q} \log \left[1 + \frac{p(\mathbf{y})}{\nu q(\mathbf{y})} \right]
$$
\n
$$
= \dots \tag{13}
$$

 $= -KL(p||M_{\nu}) - \nu KL(q||M_{\nu}) + 2 \log 2$ (14)

with $M_{\nu} = (p + \nu q)/2$

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= *. . .* = −KL(p||M*ν*) − *ν*KL(q||M*ν*) + 2 log 2 (14)

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\nFor $\nu = 1$, $\overline{J}(h^*) = -2JSD(p, q) + 2 \log 2$, and hence
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 \triangleright Contrastive learning via classification with the logistic loss estimates the JSD.

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).

- \blacktriangleright 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).
- \triangleright But other loss functions can be used:
	- \triangleright multinomial logistic loss when we contrast more than two data points.
	- \blacktriangleright Bregman divergences
	- \blacktriangleright f-divergences

Constructing reference data

Choice depends on the specific application of contrastive learning.

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

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- \triangleright 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.
- \triangleright Solution in red bridges the "gap" using telescopic ratio estimation (TRE) (Rhodes, Xu, and Gutmann, NeurIPS 2020)

Telescoping density-ratio estimation (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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Let us thus use multiple bridges.

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

$$
\frac{p_0(\mathbf{x})}{p_4(\mathbf{x})} = \frac{p_0(\mathbf{x})}{p_1(\mathbf{x})} \frac{p_1(\mathbf{x})}{p_2(\mathbf{x})} \frac{p_2(\mathbf{x})}{p_3(\mathbf{x})} \frac{p_3(\mathbf{x})}{p_4(\mathbf{x})}.
$$
(16)

Telescoping density-ratio estimation (Rhodes, Xu, and Gutmann, NeurIPS 2020)

Sample efficiency curves for the 1d peaked ratio experiment.

More results in the paper!

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(Gutmann and Hyvärinen, AISTATS 2010; JMLR 2012) (Pihlaja, Gutmann, and Hyvärinen, UAI2010; Gutmann and Hirayama, UAI 2011) (Rhodes, Xu, and Gutmann, NeurIPS 2020)

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	- Either parametrise $h(x; \theta)$ in terms of an energy-based model if provided, i.e. $h(\mathbf{x}; \theta) = \log \phi(\mathbf{x}|\theta) - \log q(\mathbf{x}) + \text{const}$

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• Set
$$
\log p(\mathbf{x}|\hat{\theta}) = \underbrace{\log q(\mathbf{x})}_{\sim} + \underbrace{h(\mathbf{x}; \hat{\theta})}_{\sim}
$$

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Gaussian Copula Flow

(Figure from Rhodes, Xu, and Gutmann, NeurIPS 2020)

Bayesian inference for simulator-based models

(Likelihood-Free Inference by Ratio Estimation, Thomas et al, 2016; 2020) (Dinev and Gutmann, arXiv:1810.09899, 2018)

- \triangleright Consider simulator-based model **x** = $g(\theta, \omega)$, $\omega \sim p(\omega)$
- **I** Task: estimate the posterior $p(\theta|\mathbf{x}^{\circ})$
- \triangleright Contrastive interpretation of Bayes' rule:

$$
\underbrace{\log p(\boldsymbol{\theta})}_{\text{reference}} + \underbrace{\log p(\mathbf{x}|\boldsymbol{\theta}) - \log p(\mathbf{x})}_{\text{difference}} \Rightarrow \underbrace{\log p(\boldsymbol{\theta}|\mathbf{x})}_{\text{interest}} \qquad (17)
$$

- \triangleright Use simulator to generate data from $p(\mathbf{x}|\theta)$ and from $p(\mathbf{x})$.
- \blacktriangleright Learning the difference provides an estimate of the desired $h(\mathbf{x}, \theta) = \log p(\mathbf{x}|\theta) - \log p(\mathbf{x})$ and hence an estimate of the posterior.

(Kleinegesse and Gutmann, AISTATS 2019; ICML 2020; arXiv:2105.04379) (Kleinegesse, Drovandi and Gutmann, Bayesian Analysis 2020) (Ivanova, Foster, Kleinegesse, Gutmann and Rainforth, NeurIPS 2021)

 \blacktriangleright 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** $\theta = (\beta, \gamma)$ **: infection rate and recovery rate**

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- **Parameters** $\boldsymbol{\theta} = (\beta, \gamma)$ **: infection rate and recovery rate**
- \blacktriangleright Task: find the optimal times at which to take measurements to most accurately estimate *θ*.

 \triangleright Experimental design by maximising mutual information (MI)

$$
\hat{\mathbf{d}} = \underset{\mathbf{d}}{\text{argmax}} \mathbb{E}_{p(\mathbf{x}, \theta | \mathbf{d})} \log \left[\frac{p(\mathbf{x} | \theta, \mathbf{d})}{p(\mathbf{x} | \mathbf{d})} \right] \tag{18}
$$

 \triangleright Use contrastive learning to estimate

$$
h_{\mathbf{d}}(\mathbf{x}, \boldsymbol{\theta}) = \log p(\mathbf{x}|\boldsymbol{\theta}, \mathbf{d}) - \log p(\mathbf{x}|\mathbf{d}), \tag{19}
$$

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

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

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

I Learning the ratio $h_d(x, \theta)$ and approximating the MI is computationally costly.

$$
\hat{\textbf{d}} = \text{argmax}_{\textbf{d}} \mathbb{E}_{\rho(\textbf{x}, \theta | \textbf{d})} \log \left[\tfrac{\rho(\textbf{x} | \theta, \textbf{d})}{\rho(\textbf{x} | \textbf{d})} \right]
$$

- Example 1 Learning the ratio $h_d(x, \theta)$ and approximating the MI is computationally costly.
- \triangleright But we do not need to estimate the MI accurately everywhere! Only around it's maximum.

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- In Suggests approach using lower bounds on the MI (or proxy quantities) where we concurrently tighten the bound and maximise the (proxy) MI.

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(Kleinegesse and Gutmann, ICML 2020; arXiv:2105.04379)

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

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- \blacktriangleright For more details and other loss functions: Kleinegesse and Gutmann, ICML 2020; arXiv:2105.04379

SIR example

Conclusions

- Introduced energy-based (unnormalised) and simulator-based (implicit) models.
- \triangleright Pointed out that their likelihood function is computationally intractable.
- Introduced contrastive learning as an intuitive and computationally feasible alternative to likelihood-based learning.
- \triangleright Contrastive learning is closely related to classification, logistic regression, and ratio estimation.
- \triangleright We can use it to solve a range of difficult statistical problems:
	- 1. Parameter estimation for energy-based models
	- 2. Bayesian inference for simulator-based models
	- 3. Bayesian experimental design for simulator-based models
- For papers, see <https://michaelgutmann.github.io>