Neural Likelihood-free Inference

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The probability of ending in bin x corresponds to the total probability of all the paths z from start to x,

$$p(x| heta) = \int p(x,z| heta) dz = inom{n}{x} heta^x (1- heta)^{n-x}.$$



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But what if we shift or remove some of the pins?

The Galton board is a metaphore of simulation-based science:

Galton board device	\rightarrow	Computer simulation
Parameters $ heta$	\rightarrow	Model parameters $ heta$
Buckets x	\rightarrow	Observables x
Random paths <i>z</i>	\rightarrow	Latent variables <i>z</i> (stochastic execution traces through simulator)

Inference in this context requires likelihood-free algorithms.



Credits: Johann Brehmer.



A thriving field of research



Particle physics







Astrophysics



Epidemiology



Climatology

(... and many others!)

Particle physics



SM with parameters θ

$$\begin{split} & \int (f_{1}(x,y_{1},y_{2},x,y_{2},y_{2$$

Simulated observables $oldsymbol{x}$



Real observations $x_{ m obs}$







Credits: Johann Brehmer.







$$p(x| heta) = igstarrow ec{\int} p(z_p| heta) p(z_s|z_p) p(z_d|z_s) p(x|z_d) dz_p dz_s dz_d$$
intractable

Likelihood ratio

The likelihood ratio

$$r(x| heta_0, heta_1)=rac{p(x| heta_0)}{p(x| heta_1)}$$

is the quantity that is central to many statistical inference procedures.

Examples

- Frequentist hypothesis testing
- Supervised learning
- Bayesian posterior sampling with MCMC
- Bayesian posterior inference through Variational Inference
- Generative adversarial networks
- Empirical Bayes with Adversarial Variational Optimization
- Optimal compression

When solving a problem of interest, do not solve a more general problem as an intermediate step. – Vladimir Vapnik



Direct likelihood ratio estimation is simpler than density estimation.

(This is fortunate, we are in the likelihood-free scenario!)

Frequentist inference

The frequentist (physicist's) way

The Neyman-Pearson lemma states that the likelihood ratio

$$r(x| heta_0, heta_1) = rac{p(x| heta_0)}{p(x| heta_1)}$$



is the most powerful test statistic to discriminate between a null hypothesis θ_0 and an alternative θ_1 .





Define a projection function $s:\mathcal{X} o\mathbb{R}$ mapping observables x to a summary statistic x'=s(x).

Then, approximate the likelihood p(x| heta) with the surrogate $\hat{p}(x| heta) = p(x'| heta)$.

From this it comes

$$rac{p(x| heta_0)}{p(x| heta_1)}pproxrac{\hat{p}\left(x| heta_0
ight)}{\hat{p}\left(x| heta_1
ight)}=\hat{r}(x| heta_0, heta_1).$$

Wilks theorem

Consider the test statistic

$$q(heta) = -2\sum_x \log rac{p(x| heta)}{p(x|\hat{ heta})} = -2\sum_x \log r(x| heta, \hat{ heta})$$

for a fixed number N of observations $\{x\}$ and where $\hat{\theta}$ is the maximum likelihood estimator.

When $N \to \infty$, $q(\theta) \sim \chi_2$. Therefore, an observed value $q_{\rm obs}(\theta)$ translates directly to a p-value that measures the confidence with which θ can be excluded:

$$p_{ heta} \equiv \int_{q_{
m obs}(heta)}^{\infty} \mathrm{d}q \, p(q| heta) = 1 - F_{\chi_2}(q_{
m obs}(heta)).$$



Discovery of the Higgs boson at 5- σ (p-value cutoff at $3 imes 10^{-7}$)

- Choosing the projection *s* is difficult and problem-dependent.
- Often there is no single good variable: compressing to any x' loses information.
- Ideally, analyze high-dimensional x', including all correlations.
- Unfortunately, filling highdimensional histograms is not tractable.





Supervised learning provides a way to automatically construct s:

- Let us consider a neural network classifier \hat{s} tasked to distinguish $x\sim p(x| heta_0)$ from $x\sim p(x| heta_1).$
- Train \hat{s} by minimizing the cross-entropy loss

$$egin{aligned} L_{XE}[\,\hat{s}\,] &= -\mathbb{E}_{p(x| heta)\pi(heta)}[1(heta= heta_0)\log\,\hat{s}(x) + \ &1(heta= heta_1)\log(1-\hat{s}(x))] \end{aligned}$$



The solution \hat{s} found after training approximates the optimal classifier

$$\hat{s}(x)pprox s^*(x)=rac{p(x| heta_1)}{p(x| heta_0)+p(x| heta_1)}.$$

Therefore,

$$r(x| heta_0, heta_1)pprox \hat{r}(x| heta_0, heta_1)=rac{1-\hat{s}(x)}{\hat{s}(x)}$$

That is, supervised classification is equivalent to likelihood ratio estimation.

Cranmer, Pavez and Louppe, 2015 [arXiv: 1506.02169].



To avoid retraining a classifier \hat{s} for every (θ_0, θ_1) pair, fix θ_1 to θ_{ref} and train a single parameterized classifier $\hat{s}(x|\theta_0, \theta_{ref})$ where θ_0 is also given as input.

Therefore, we have

$$\hat{r}\left(x| heta_{0}, heta_{ ext{ref}}
ight)=rac{1-\hat{s}\left(x| heta, heta_{ ext{ref}}
ight)}{\hat{s}\left(x| heta_{0}, heta_{ ext{ref}}
ight)}$$

such that for any (θ_0, θ_1) ,

$$r(x| heta_0, heta_1)pprox rac{\hat{r}\left(x| heta_0, heta_{ ext{ref}}
ight)}{\hat{r}\left(x| heta_1, heta_{ ext{ref}}
ight)}.$$

Cranmer, Pavez and Louppe, 2015 [arXiv:1506.02169].

Opening the black box





Traditional likelihood-free inference treats the simulator as a generative black box: parameters in, samples out. But in most real-life problems, we have access to the simulator code and some understanding of the microscopic processes.



 $p(x|\theta)$ is usually intractable. What about $p(x, z|\theta)$?

As the trajectory $z_1, ..., z_T$ and the observable x are emitted, it is often possible:

- to calculate the joint likelihood $p(x, z | \theta)$;
- to calculate the joint likelihood ratio $r(x,z| heta_0, heta_1);$
- to calculate the joint score $t(x,z| heta_0) =
 abla_ heta\log p(x,z| heta) \Big|_{ heta_0}$.

We call this process mining gold from your simulator!

Extracting the joint likelihood ratio

- Computer simulations typically evolve along a tree-like structure of successive random branchings.
- The probabilities of each branching $p(z_i|z_{i-1}, \theta)$ are often clearly defined in the code:

```
if random() > 0.1+2.5+model_parameter:
    do_one_thing()
else:
    do_another_thing()
```

• For each run, we can calculate the probability of the chosen path for different values of the parameters and the joint likelihood-ratio:

$$r(x,z| heta_0, heta_1) = rac{p(x,z| heta_0)}{p(x,z| heta_1)} = \prod_i rac{p(z_i|z_{i-1}, heta_0)}{p(z_i|z_{i-1}, heta_1)}$$



Regressing the likelihood ratio

Observe that the joint likelihood ratios

 $r(x,z| heta_0, heta_1)=rac{p(x,z| heta_0)}{p(x,z| heta_1)}$

are scattered around $r(x| heta_0, heta_1)$.

Can we use them to approximate $r(x| heta_0, heta_1)$?



Consider the squared error of a function $\hat{g}(x)$ that only depends on x, but is trying to approximate a function g(x, z) that also depends on the latent z:

$$L_{MSE} = \mathbb{E}_{p(x,z| heta)} \left[(g(x,z) - \hat{g}(x))^2
ight].$$

Via calculus of variations, we find that the function $g^*(x)$ that extremizes $L_{MSE}[g]$ is given by

$$egin{aligned} g^*(x) &= rac{1}{p(x| heta)}\int p(x,z| heta)g(x,z)dz \ &= \mathbb{E}_{p(z|x, heta)}\left[g(x,z)
ight] \end{aligned}$$

Therefore, by identifying the g(x,z) with the joint likelihood ratio $r(x,z|\theta_0,\theta_1)$ and θ with θ_1 , we define

$$L_r = \mathbb{E}_{p(x,z| heta_1)}\left[(r(x,z| heta_0, heta_1) - \hat{r}(x))^2
ight],$$

which is minimized by

$$egin{aligned} r^*(x) &= rac{1}{p(x| heta_1)} \int p(x,z| heta_1) rac{p(x,z| heta_0)}{p(x,z| heta_1)} dz \ &= rac{p(x| heta_0)}{p(x| heta_1)} \ &= r(x| heta_0, heta_1). \end{aligned}$$



 $r^*(x| heta_0, heta_1) = rg\min_{\hat{r}} L_r[\hat{r}]$

Regressing the score

Similarly, we can mine the simulator to extract the joint score

$$t(x,z| heta_0) =
abla_ heta \log p(x,z| heta)igert_{ heta_0},$$

which indicates how much more or less likely x, z would be if one changed θ_0 .



Using the same trick, by identifying g(x,z) with the joint score $t(x,z| heta_0)$ and heta with $heta_0$, we define

$$L_t = \mathbb{E}_{p(x,z| heta_0)}\left[(t(x,z| heta_0) - \,\hat{t}\,(x))^2
ight],$$

which is minimized by

$$egin{aligned} t^*(x) &= rac{1}{p(x| heta_0)} \int p(x,z| heta_0) (
abla_ heta \log p(x,z| heta)ig|_{ heta_0}) dz \ &= rac{1}{p(x| heta_0)} \int p(x,z| heta_0) rac{
abla_ heta p(x,z| heta)ig|_{ heta_0}}{p(x,z| heta_0)} dz \ &= rac{
abla_ heta p(x| heta)ig|_{ heta_0}}{p(x| heta_0)} \ &=
abla_ heta \log p(x| heta)ig|_{ heta_0} \ &= t(x| heta_0). \end{aligned}$$



$L_{RASCAL} = L_r + L_t$





$L_{RASCAL} = L_r + L_t$



SALLY (= optimal compression)

The local model

In the neighborhood of $heta_{
m ref}$, the Taylor expansion of $\log p(x| heta)$ is

$$\log p(x| heta) = \log p(x| heta_{ ext{ref}}) + \underbrace{
abla_ heta \log p(x| heta)}_{t(x| heta_{ ext{ref}})} \cdot (heta - heta_{ ext{ref}}) + O((heta - heta_{ ext{ref}})^2)$$



This results in the exponential model

$$p_{ ext{local}}(x| heta) = rac{1}{Z(heta)} p(t(x| heta_{ ext{ref}})| heta_{ ext{ref}}) \exp(t(x| heta_{ ext{ref}}) \cdot (heta - heta_{ ext{ref}}))$$

where the score $t(x| heta_{ ext{ref}})$ are its sufficient statistics.

That is,

- knowing $t(x| heta_{
 m ref})$ is just as powerful as knowing the full function $\log p(x| heta)$.
- x can be compressed into a single scalar $t(x| heta_{
 m ref})$ without loss of power.

Brehmer, Louppe, Pavez and Cranmer, 2018 [arXiv: 1805.12244].

SALLY



There is more...

Method	Simulate	Extr $r(x,z)$	act t(x,z)	NN estimates	Asympt. exact	Generative
ROLR	$ heta_0 \sim \pi(heta)$, $ heta_1$	\checkmark		$\hat{r}(x heta_0, heta_1)$	\checkmark	
CASCAL	$ heta_0 \sim \pi(heta)$, $ heta_1$		\checkmark	$\hat{r}(x heta_0, heta_1)$	\checkmark	
ALICE	$ heta_0 \sim \pi(heta)$, $ heta_1$		\checkmark	$\hat{r}(x heta_0, heta_1)$	\checkmark	
RASCAL	$ heta_0 \sim \pi(heta)$, $ heta_1$	\checkmark	\checkmark	$\hat{r}(x heta_0, heta_1)$	\checkmark	
ALICES	$ heta_0 \sim \pi(heta)$, $ heta_1$	\checkmark	\checkmark	$\hat{r}(x heta_0, heta_1)$	\checkmark	
SCANDAL	$\theta \sim \pi(\theta)$		\checkmark	$\hat{p}(x heta)$	\checkmark	\checkmark
SALLY	$ heta_{ref}$		\checkmark	$\hat{t}(x \theta_{ref})$	in local approx.	
SALLINO	$ heta_{ref}$		\checkmark	$\hat{t}(x heta_{ref})$	in local approx.	

Examples

① Hunting new physics at particle colliders

The goal is to constrain two EFT parameters and compare against traditional histogram analysis.



Exciting new physics might hide here! We parameterize it with two EFT coefficients:





at least 16-dimensional

observable space

② Dark matter substructure from gravitational lensing



Number of dark matter subhalos and their mass and location lead to complex latent space of each image. The goal is the **inference of population parameters**.



Bayesian inference

Bayesian inference = computing the posterior

 $p(heta|x) = rac{p(x| heta)p(heta)}{p(x)}.$



Doubly intractable in the likelihood-free scenario:

- Cannot evaluate the likelihood $p(x| heta) = \int p(x,z| heta) dz.$
- Cannot evaluate the evidence $p(x) = \int p(x|\theta)p(\theta)d\theta$.

Approximate Bayesian Computation (ABC)



Issues

- How to choose x'? ϵ ? $|| \cdot ||$?
- No tractable posterior.
- Need to run new simulations for new data or new prior.

Amortizing Bayes

The Bayes rule can be rewritten as

$$p(heta|x) = rac{p(x| heta)p(heta)}{p(x)} = r(x| heta)p(heta) pprox \hat{r}(x| heta)p(heta),$$

where $r(x| heta) = rac{p(x| heta)}{p(x)}$ is the likelihood-to-evidence ratio.

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where $r(x| heta) = rac{p(x| heta)}{p(x)}$ is the likelihood-to-evidence ratio.

As before, the likelihood-to-evidence ratio can be approximated e.g. from a neural network classifier trained to distinguish $x \sim p(x|\theta)$ from $x \sim p(x)$, hence enabling direct and amortized posterior evaluation.



Bayesian inference of dark matter subhalo population parameters



Brehmer, Mishra-Sharma, Hermans, Louppe, and Cranmer, 2019 [arXiv:1909.02005].

MCMC posterior sampling



Likelihood-free MCMC

MCMC samplers require the evaluation of the posterior ratios:

$$egin{aligned} &rac{p(heta_{ ext{new}}|x)}{p(heta_{t-1}|x)} &= rac{p(x| heta_{ ext{new}})p(heta_{ ext{new}})/p(x)}{p(x| heta_{t-1})p(heta_{t-1})/p(x)} \ &= rac{p(x| heta_{ ext{new}})p(heta_{ ext{new}})}{p(x| heta_{t-1})p(heta_{t-1})} \ &= r(x| heta_{ ext{new}}, heta_{t-1})rac{p(heta_{ ext{new}})/p(heta_{ ext{new}})}{p(heta_{ ext{new}})} \end{aligned}$$

Again, MCMC samplers can be made likelihood-free by plugging a learned approximation $\hat{r}(x|\theta_{\text{new}}, \theta_{t-1})$ of the likelihood ratio.

For MCMC, best results are obtained when using ratios of likelihood-to-evidence ratios:

$$\hat{r}(x| heta_{ ext{new}}, heta_{t-1}) = rac{\hat{r}(x| heta_{ ext{new}})}{\hat{r}(x| heta_{t-1})}$$

Hermans, Begy and Louppe, 2019 [arXiv:1903.04057].



Figure 3: Posteriors from the tractable benchmark. The experiments are repeated 25 times and the approximate posteriors are subsampled from those runs. AALR-MCMC shares the same structure with the MCMC truth, demonstrating its accuracy. Some runs of the other methods were not consistent, contributing to the variance observed in Table 2.

Algorithm	MMD	ROC AUC
AALR-MCMC (ours)	0.05 ± 0.005	0.59 ± 0.0010
ABC ($\epsilon = 32$)	0.51 ± 0.001	0.99 ± 0.0001
ABC ($\epsilon = 16$)	0.50 ± 0.003	0.99 ± 0.0002
ABC $(\epsilon = 8)$	0.39 ± 0.001	0.99 ± 0.0003
ABC ($\epsilon = 4$)	0.29 ± 0.004	0.98 ± 0.0007
APT	0.17 ± 0.036	0.86 ± 0.0008
AALR-MCMC (LRT)	0.53 ± 0.004	0.99 ± 0.0001
SNPE-A	0.21 ± 0.070	0.97 ± 0.0098
SNPE-B	0.20 ± 0.061	0.92 ± 0.0181

Table 2: AALR-MCMC outperforms all other methods. Numerical errors introduced by MCMC might have contributed to these results. A comparison of the PDFs between the true posterior and our ratio estimator are shown in Figure 11 (Appendix D.2). The MMD scores are in agreement with [41].

Probabilistic programming



A probabilistic program defines a joint distribution of unobserved x and observed y variables p(x, y).

Probabilistic programming extends ordinary programming with two added constructs:

- Sampling from distributions
- Conditioning random variables by specifying observed values

Inference engines give us distributions over unobserved variables, given observed variables (data)

$$p(x|y) = rac{p(y|x)p(x)}{p(y)}$$

Probabilistic programming languages

- Anglican (Clojure)
- Church (Scheme)
- Edward, TensorFlow Probability (Python, TensorFlow)
- Pyro (Python, PyTorch)
- Figaro (Scala)
- Infer.NET (C#)
- LibBi (C++ template library)
- PyMC3 (Python)
- Stan (C++)
- WebPPL (JavaScript)

A stochastic simulator implicitly defines a probability distribution by sampling pseudo-random numbers.

Scientific simulators are probabilistic programs!



Key idea

Let a neural network take full control of the internals of the simulation program by hijacking all calls to the random number generator.



Le et al, 2016 [arXiv:1610.09900]; Baydin et al, 2018 [arXiv:1807.07706]; Baydin et al, 2019 [arXiv:1907.03382].



③ Taking control of Sherpa (particle physics simulator)

- τ decay in Sherpa, 38 decay channels, coupled with an approximate calorimeter simulation in C++.
- Observations are 3D calorimeter depositions.
- Latent variables (Monte Carlo truth) of interest: decay channel, px, py, pz momenta, final state momenta and IDs.





Inference results



MCMC true posterior (7.7M single node)

IC proposal from trained NN

IC posterior after importance weighting





We obtain posteriors over the whole Sherpa address space, 1000s of addresses.

Interpretability

Latent probabilistic structure of the 250 most frequent trace types:







Summary

- Much of modern science is based on "likelihood-free" simulations.
- The likelihood-ratio is central to many statistical inference procedures, regardless of your religion.
- Supervised learning enables likelihood-ratio estimation.
- Better likelihood-ratio estimates can be achieved by mining simulators.
- Probabilistic programming enables posterior inference in scientific simulators.



Collaborators



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The end.