#### **Simulation-based Inference**

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#### **Abstract**

Simulators are used across different scientific domains where serve as valuable tools to encode empirical knowledge about a system of interest. Inference in this setting often boils down to finding parameters of the simulator for a given, real-world observation, which typically cannot be computed analytically. Papamakarios and Murray [2016] propose to learn a posterior over a simulators parameters by using neural density estimators. Here, we give an overview of their work and SBI in general.

### **Problem Setting and Related Work**

Scientific fields, such as population genetics, particle physics, epidemiology, astrophysics among others make use of sophisticated simulators to model observed systems [Brehmer and Cranmer, 2020, de Witt et al., 2020, Delaunoy et al., 2020, Cranmer et al., 2020, Pritchard et al., 1999]. These simulators are able to encode prior knowledge like causal relations, hierarchies of variables, ... Performing inference in this setting, however, would require a finding a posterior over parameters  $\theta$  for a given observation  $x_0$ . This can be stated in closed form as

$$p(\theta|x=x_0) = \frac{p(x|\theta)p(\theta)}{p(x)} = \frac{\int p(x,z|\theta)dz \, p(\theta)}{\int p(x|\theta)p(\theta)d\theta},$$

$$intractable$$
(1)

where z is a nuisance parameter. If likelihood and evidence are intractable, which is typically the case, this cannot be computed analytically. Intuitively, it can be thought of as inverting the simulator.

A collective term for methods to solve this inverse problem is Approximate Bayesian Computation (ABC). The most simple approach is rejection ABC [Pritchard et al., 1999] where samples produced by a simulator are discarded if notis within a  $\epsilon$ -ball around the observation of interest. For small values of  $\epsilon$  this method may take many steps to even produce a single matching parameter set. For large  $\epsilon$  it is not precise. Moreover, this method does not produce a full posterior over the parameter but merely yields point estimates with confident intervals. With Markov-Chain-Monte-Carlo ABC and Sequential ABC improvements over rejection ABC have been proposed. These methods produce samples more efficiently. While this can lead to faster convergence, it does not produce a full posterior over parameters that is conditioned on the actual observation, but rather a sample that is close to it.

## **Neural Approach**

Papamakarios and Murray [2016] build on recent advances in deep learning to learn a posterior over parameters directly from sample produced by a simulator. They use Deep Neural Networks (DNN) to parameterize a Gaussian Mixture Model (GMM). They use the simulator to perform n simulations and storing pairs of samples x and corresponding parameters  $\theta$ . This gives the set  $\{(x_i, \theta_i)\}_{i \in n}$  which they use as training data. DNNs are fitted to parameterize a GMM

$$\sum_{k} \alpha_{k} \mathcal{N}(\theta | m_{k}, S_{k}) = \sum_{k} f_{1}(x)_{k} \mathcal{N}(\theta | f_{2}(x)_{k}, diag(f_{3}(x))_{k})$$
(2)

, where k is the number of components, m the mean and S the covariance matrix of a Gaussian, and  $f_{\{1,2,3\}}$  are MLPs.

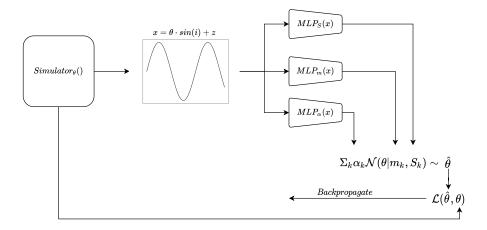


Figure 1: Given samples by the simulator, Multilayer perceptrons (MLP) are trained to parameterize a distribution over parameters generating the sample. A training signal is created by sampling from this distribution and comparing the sample to the actual parameters which are known.

### Possible Enhancements/Open Ends

#### **Conclusion**

Use only the provided maths commands use align for embedded maths (not align\*):

$$\mathscr{F}, \mathbb{R}, \mathbb{N}$$
 (3)

The diff command is for proper typesetting in integrals:  $\int f(x) dx$ , not  $\int f(x) dx$ . If you need conditional probabilities, use mid for *A* conditioned on *B*:  $p(A \mid B)$ .

Write clearly; pay attention to clarity, simplicity, and – above all – correctness of your statements. Look up how to write a scientific document.

# References

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