#### Bayesian Inference via Classification

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4th October 2016

M.U. Gutmann, R. Dutta, S. Kaski, and J. Corander Statistical Inference of Intractable Generative Models via Classification, *arXiv:1407.4981*.

J. Lintusaari, M.U. Gutmann, R. Dutta, S. Kaski, and J. Corander Fundamentals and Recent Developments in Approximate Bayesian Computation, *Systematic Biology*, in press, 2016. By using classification, we can measure the distance between two data sets and infer the parameters of intractable generative models.

#### ► Given:

- Observed data y<sup>o</sup>
- Intractable generative model, parametrised by θ
- (Possibly) A prior probability density function (pdf) for  $\theta$ ,  $p_{\theta}$
- Task: Perform inference about heta
  - Maximum likelihood estimate
  - Posterior distribution of heta

# Generative model

Model that specifies a mechanism for generating data y

- e.g. stochastic dynamical system
- computer model / simulator of some complex biological process
- Such models are widely used
  - Evolutionary biology: Simulating evolution
  - Neuroscience: Simulating neural circuits
  - Ecology: Simulating species migration
  - Health science: Simulating the spread of an infectious disease



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

## Generative model

- Let  $(\Omega, \mathcal{F}, \mathcal{P})$  be a probability space.
- A generative model is a collection of (measurable) functions g(., θ) parametrized by θ,

$$\boldsymbol{\omega} \in \Omega \mapsto \mathbf{y} = g(\boldsymbol{\omega}, \boldsymbol{\theta}) \in \mathcal{Y}$$
 (1)

For any fixed  $\theta$ ,  $\mathbf{y} = g(., \theta)$  is a random variable.



- Generally impossible to write down the pdf p(y|θ) of the generated data.
- Likelihood function L(θ) = p(y<sup>o</sup>|θ) not known well defined, but computation not tractable

supplemental

Exact inference not possible

# Approximate inference for generative models

- Inference is approximate due to approximation of  $p(\mathbf{y}^o|\boldsymbol{\theta})$
- Rough classification:
  - Parametric approximation: e.g. synthetic likelihood (Wood, Nature, 2010)
  - Nonparametric approximation: e.g approximate Bayesian computation

(Recent review: Gutmann et al, Systematic Biology, 2016)

General idea: Identify the values of the parameters θ for which simulated data resemble the observed data

# Meta algorithm for approximate Bayesian computation

- Let y<sup>o</sup> be the observed data.
- Iterate many times:
  - 1. Sample  $\theta$  from a proposal distribution  $q(\theta)$
  - 2. Sample  $\mathbf{y}|\boldsymbol{\theta}$  according to the model
  - Compute the distance d(y, y<sup>o</sup>) between simulated and observed data
  - 4. Retain  $\boldsymbol{\theta}$  if  $d(\mathbf{y}, \mathbf{y}^{o}) \leq \epsilon$
- Different choices for  $q(\theta)$  give different algorithms
- ► Produces samples from the (approximate) posterior when the bandwidth *\epsilon* is small.

- 1. How to handle the computational cost?
- 2. How to measure the distance *d* between simulated and observed data?

## Two major difficulties

- 1. How to handle the computational cost?
  - → Use Bayesian optimization
    M.U. Gutmann and J. Corander
    Bayesian optimization for likelihood-free inference of simulator-based statistical models
     Journal of Machine Learning Research, 17(125): 1–47, 2016
- 2. How to measure the distance *d* between simulated and observed data?
  - $\rightarrow~$  Use classification

M.U. Gutmann, R. Dutta, S. Kaski, and J. Corander Statistical Inference of Intractable Generative Models via Classification, *arXiv:1407.4981*  Correctly classifying data into two categories is usually easier if the two data sets were generated with very different values of  $\theta$  (left) than with similar values (right).





#### Using classification to measure the distance

- Classification accuracy (discriminability) serves as distance measure.
- ▶ Value of 1/2: close; Value of 1: far
- Complete arsenal of classification methods becomes available to inference.



### Using the distance for posterior inference

- Approximate Bayesian computation (ABC) with classification accuracy as distance measure
- Results for toy models:



### Application to epidemiology of infectious diseases

- Data: Colonization states of sampled attendees of 29 child day care centres (DCCs).
- DCC example: Each square indicates an attendee colonized with a strain of the bacterium *Streptococcus pneumoniae*.



# Application to epidemiology of infectious diseases

- Generative model: latent continuous-time Markov chain for the transmission dynamics in a DCC and an observation model (Numminen et, Biometrics, 2013).
- The model has three parameters:
  - $\beta$ : rate of infections within a DCC
  - Λ: rate of infections outside a DCC
  - $\theta$ : possibility to be infected with multiple strains
- Likelihood is intractable (data at a single time point are available only).

## Application to epidemiology of infectious diseases

- Our classification-based distance measure does not use domain/expert knowledge.
- Performs as well as a distance measure based on domain knowledge (Numminen et, Biometrics, 2013).



- Topic: Inference for generative models where the likelihood function is intractable
- Inference principle: Find parameter values for which the distance between simulated and observed data is small
- Showed that, by using classification, we can
  - measure the distance between two data sets, and
  - infer the parameters of intractable generative models.
- ▶ More (results, math, related work, ...) in the reference paper!

## Likelihood function is implicitly defined

