

# ABC: a review and some recent developments

---

Dennis Prangle

November 2019

Newcastle University, UK

# Traditional modelling and inference

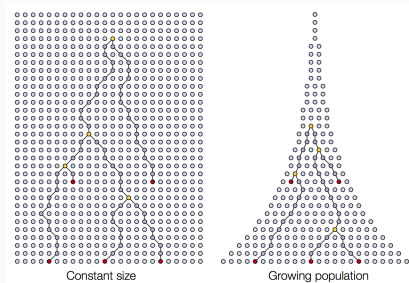
- Observe some data  $y_{\text{obs}}$
- Define **model** with density  $p(y|\theta)$  and parameters  $\theta$
- **Likelihood function** is  $L(\theta) = p(y_{\text{obs}}|\theta)$
- Use  $L(\theta)$  to learn parameters  
e.g. maximum likelihood or Bayesian inference by MCMC
- Inference output + model allows:
  - **Prediction** of future behaviour
  - **Explanation** of phenomena
  - **Control** by understanding effect of interventions
  - ...

# Generative models

- Often hard to directly define probability models for complex systems!
- Easier to describe how system **evolves**
- Can then simulate data given parameters easily
- i.e. **generative** model
- Likelihood calculations for generative models often very hard!
- Goal: **likelihood-free inference** methods
- Inference without making use of likelihood function
- One such method is ABC: *approximate Bayesian computation*

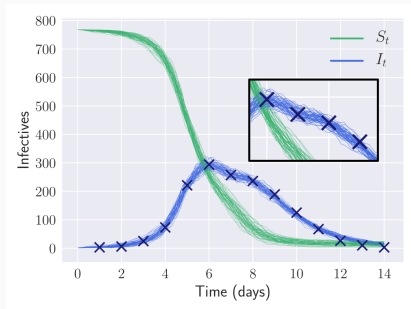
# Applications - population genetics

- Population genetics was first ABC application (late 90s)
- Data are genetic sequences from a population
- Parameters include mutation rates etc
- And also demographic history of population (migration, growth etc)
- Easy to simulate data from parameters
- Likelihood not available



# Applications - infectious disease epidemiology

- Can model infectious disease by population counts of susceptible, infectious and recovered
- Parameters control rates of change between populations
- Model can be ODE, SDE, jump process etc
- All easy to simulate from (at least approximately)



## Applications - infectious disease epidemiology

- Can add extra structure e.g. exposed stage, immigration
- Or move from population level model to **individual based model**
- Many other applications for ODEs, SDEs, jump processes, IBMs!

THE ASTROPHYSICAL JOURNAL, 764:116 (15pp), 2013 February 20  
© 2013. The American Astronomical Society. All rights reserved. Printed in the U.S.A.

doi:[10.1088/0004-637X/764/2/116](https://doi.org/10.1088/0004-637X/764/2/116)

## LIKELIHOOD-FREE COSMOLOGICAL INFERENCE WITH TYPE Ia SUPERNOVAE: APPROXIMATE BAYESIAN COMPUTATION FOR A COMPLETE TREATMENT OF UNCERTAINTY

ANJA WEYANT<sup>1</sup>, CHAD SCHAFER<sup>2</sup>, AND W. MICHAEL WOOD-VASEY<sup>1</sup>

<sup>1</sup> Pittsburgh Particle Physics, Astrophysics, and Cosmology Center (PITT PACC), Physics and Astronomy Department,  
University of Pittsburgh, Pittsburgh, PA 15260, USA; [amw19@pitt.edu](mailto:amw19@pitt.edu)

<sup>2</sup> Department of Statistics, Carnegie Mellon University, Pittsburgh, PA 15213, USA  
*Received 2012 June 12; accepted 2012 December 10; published 2013 January 30*

### ABSTRACT

Cosmological inference becomes increasingly difficult when complex data-generating processes cannot be modeled by simple probability distributions. With the ever-increasing size of data sets in cosmology, there is an increasing

## LIKELIHOOD-FREE COSMOLOGICAL INFERENCE WITH TYPE Ia SUPERNOVAE: APPROXIMATE BAYESIAN COMPUTATION FOR A COMPLETE TREATMENT OF UNCERTAINTY

ANJA WEYANT<sup>1</sup>, CHAD SCHAFER<sup>2</sup>, AND W. MICHAEL WOOD-VASEY<sup>1</sup>

<sup>1</sup> Pittsburgh Particle Physics, Astrophysics, and Cosmology Center (PITT PACC), Physics and Astronomy Department,  
University of Pittsburgh, Pittsburgh, PA 15260, USA; [amw19@pitt.edu](mailto:amw19@pitt.edu)

<sup>2</sup> Department of Statistics, Carnegie Mellon University, Pittsburgh, PA 15213, USA  
*Received 2012 June 12; accepted 2012 December 10; published 2013 January 30*

### ABSTRACT

Cosmological inference becomes increasingly difficult when complex data-generating processes cannot be modeled by simple probability distributions. With the ever-increasing size of data sets in cosmology, there is an increasing

## Distance-learning For Approximate Bayesian Computation To Model a Volcanic Eruption

Lorenzo Pacchiardi<sup>1</sup>, Pierre Kunzli<sup>2\*</sup>, Marcel Schoengens<sup>3</sup>, Bastien Chopard<sup>2</sup>,  
Ritabrata Dutta<sup>4†</sup>



## Approximate Bayesian computation for estimating number concentrations of monodisperse nanoparticles in suspension by optical microscopy

Magnus Röding\*

*SP Food and Bioscience, Soft Materials Science, Göteborg, Sweden and  
School of Energy and Resources, UCL Australia, University College London, Adelaide, Australia*

Elisa Zagato, Katrien Remaut, and Kevin Braeckmans

*Laboratory of General Biochemistry and Physical Pharmacy, Ghent University, Ghent, Belgium and  
Center for Nano- and Biophotonics, Ghent University, Ghent, Belgium*

(Dated: June 3, 2016)

We present an approximate Bayesian computation (ABC) scheme for estimating number concentrations of monodisperse diffusing nanoparticles in suspension by optical particle tracking mi-

### Approximate Bayesian computation for estimating number concentrations of monodisperse nanoparticles in suspension by optical microscopy

Magnus Röding\*

*SP Food and Bioscience, Soft Materials Science, Göteborg, Sweden and  
School of Energy and Resources, UCL Australia, University College London, Adelaide, Australia*

Elisa Zagato, Katrien Remaut, and Kevin Braeckmans

*Laboratory of General Biochemistry and Physical Pharmacy, Ghent University, Ghent, Belgium and  
Center for Nano- and Biophotonics, Ghent University, Ghent, Belgium*

(Dated: June 3, 2016)

We present an approximate Bayesian computation (ABC) scheme for estimating number concentrations of monodisperse diffusing nanoparticles in suspension by optical particle tracking mi-

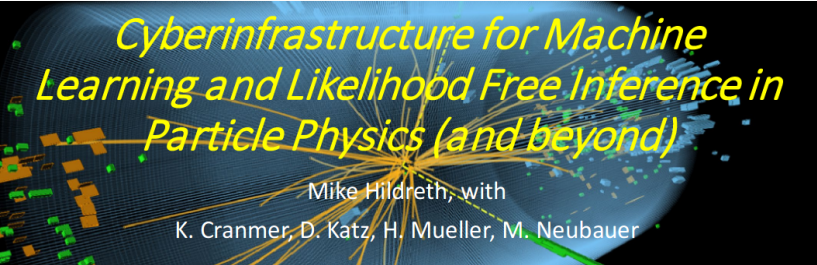
## Bayesian semi-individual based model with approximate Bayesian computation for parameters calibration: Modelling Crown-of-Thorns populations on the Great Barrier Reef

C.C.-M. Chen <sup>a, b</sup> ✉, C.C. Drovandi <sup>b, c</sup> ✉, J.M. Keith <sup>d</sup> ✉, K. Anthony <sup>a</sup> ✉, M.J. Caley <sup>b, c</sup> ✉, K.L. Mengersen <sup>b, c</sup> ✉

✉ [Show more](#)

<https://doi.org/10.1016/j.ecolmodel.2017.09.006>

[Get rights and content](#)



*Cyberinfrastructure for Machine Learning and Likelihood Free Inference in Particle Physics (and beyond)*

Mike Hildreth, with  
K. Cranmer, D. Katz, H. Mueller, M. Neubauer

## Queueing example

- Example used throughout talk
- A single queue
- Gaps between arrivals at back are  $Exp(\theta_1)$
- Service times on reaching front are  $U(\theta_2, \theta_3)$
- Observations are times between queue departures
- Easy to simulate data
- But probability calculations for observations challenging!
- Exact MCMC is possible (Shestopaloff and Neal 2014)
- More complex **queueing network** models useful in applications
  - Passenger flow in airport (Ebert et al 2018)
  - Computing jobs in data centre (Sutton and Jordan 2011)

# Overview of talk

- Standard ABC approaches (approximate Bayesian computation)
- DE methods (density estimation)

## Overview of talk

- Standard ABC approaches (approximate Bayesian computation)
- DE methods (density estimation) – including my recent work!

# Approximate Bayesian Computation

---

- General idea:
  - Simulate data  $y$  from various parameter values  $\theta$
  - Consider **closest matches** of  $y$  to  $y_{\text{obs}}$
  - Use corresponding parameters for inference
- Can be implemented in many different ways
- Many approaches suggested in various fields over last 50 years
- ABC puts this idea into a **Bayesian** framework



# Bayesian inference

- Specify **prior density**  $\pi(\theta)$ 
  - Beliefs about parameters before data observed
- Ideal inference goal is **posterior density**  $p(\theta|y_{\text{obs}})$ 
  - Beliefs updated to take data into account
- Posterior depends on prior and likelihood through **Bayes theorem**:

$$p(\theta|y_{\text{obs}}) \propto \pi(\theta)L(\theta)$$

i.e. posterior  $\propto$  prior  $\times$  likelihood

- Posterior is parameters **conditioned** on observing  $y_{\text{obs}}$

# Basic ABC

Input:

- Observed data  $y_{\text{obs}}$
- Threshold  $\epsilon \geq 0$
- Distance function  $d(y, y')$

Loop over  $i = 1, 2, \dots, N$ :

1. Sample  $\theta_i$  from prior  $\pi(\theta)$
2. Simulate  $y$  from model  $p(y|\theta_i)$
3. If  $d(y, y_{\text{obs}}) \leq \epsilon$  accept  $\theta_i$

Output: accepted  $\theta_i$  values

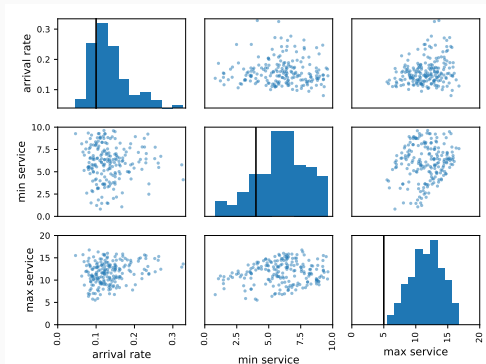
Popular variant – do  $N$  simulations then accept  $M$  with smallest distance

# ABC target distribution

- ABC output is sample from **approximate** posterior
- Ideal Bayesian inference: sample from prior + model and condition on exact match to  $y_{\text{obs}}$
- ABC: sample from prior + model and condition on **approximate** match to  $y_{\text{obs}}$
- Reducing  $\epsilon$  gives closer matches
  - more accurate output
  - but acceptances rarer
- $\epsilon$  controls trade-off between accuracy and cost

# Example - queueing model

- Data: 20 inter-departure times
- Uniform priors for:
  - Service rate  $\theta_1$  on  $[0, 1/3]$
  - Min service time  $\theta_2$  on  $[0, 10]$
  - Service time range  $\theta_3 - \theta_2$  on  $[0, 10]$
- 1 million ABC simulations, best 200 accepted



## ABC distances

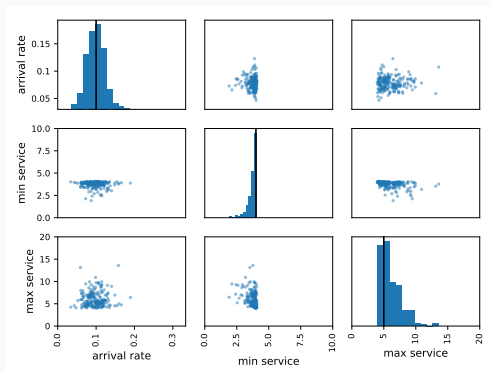
- Choice of distance function crucial
- Simplest setting:  $y$  is fixed-length vector
- Could take  $d$  as Euclidean distance
- Usually sensible to use weighted version
- i.e. normalise  $y$  components so on similar scales
- Many more sophisticated distances proposed:
  - Application specific
  - Based on theory (Wasserstein, MMD)
  - Using machine learning (kernel methods, deep distance learning)

## Summary statistics

- ABC typically poor with high dimensional data
- So lots of data dimensions must match
- $V$  unlikely to get good matches in all!
- So ABC usually reduces data to low dim **summary statistics**
- Then computes distance e.g. Euclidean distance
- Theory: rate of convergence as  $\epsilon \rightarrow 0$  worsens quickly with high  $\text{dim}(y)$ 
  - See e.g. “The rate of convergence for approximate Bayesian computation” (Barber et al 2015)
- **Dimension reduction** using summary statistics helps

## Example - queueing model

- Same data (dimension 20) and simulations as before
- 5 summary statistics: min, lower quartile, median, upper quartile, max (all normalised)



## Other ABC algorithms

- Always sampling  $\theta$  from prior is inefficient
- Posterior usually more concentrated than prior
- Lots of prior  $\theta$ s samples will be poor
- Simulating data from these **wastes time!**
- More efficient to propose promising  $\theta$ s based on previous results
- Can use **MCMC** (Markov chain Monte Carlo)
- Or **SMC** (sequential Monte Carlo)



## Other ABC algorithms

- MCMC
  - Propose next  $\theta$  near to previous accepted one
- SMC
  - Propose a group of  $\theta$ s close to previous accepted sample
- In ABC, SMC is more popular
- Can reduce  $\epsilon$  adaptively during algorithm
- Several versions of ABC-SMC algorithm
- $\epsilon$  tuning harder for MCMC (but see Vihola and Franks 2019)

## More ABC contributions

- Lots more in the literature!
- Diagnostics
- Asymptotic theory
- ABC for model choice
- Choosing summary statistics
- Interpreting approximate results
- Lots of variants on ABC algorithms

- R
  - abc
  - easyABC
- python
  - ABCPy
  - PyAbc
  - ELFI
- julia
  - GpABC
- (not an exhaustive list!)
- Also, basic ABC v easy to code yourself

- Handbook of ABC
- Review papers
  - “Approximate Bayesian Computation for infectious disease modelling” - Minter and Retkute (2019)
  - “Fundamentals and Recent Developments in Approximate Bayesian Computation” - Lintusaari et al (2017)
  - Wikipedia article (commissioned by PLOS comp bio - 2013)
  - “Approximate Bayesian computational methods” - Marin et al (2012)
  - And many more

# Pros and cons

- Pros
  - Basic idea very simple to understand
  - Easy to implement
  - Widely applicable
  - Easy to get rough parameter estimates
- Cons
  - Difficult tuning choices – distance / summary statistics
  - Approximate results – hard to quantify error
  - Doesn't scale up well to expensive simulators
  - Doesn't scale up easily to high dimensional data

# Density estimation approaches

---

# Density estimation for likelihood-free inference

- Lots of powerful density estimation methods developed in recent decade(s)
  - Mixture models
  - Copulas
  - Gaussian processes
  - Invertible neural networks (aka normalising flows)
- Can be applied to likelihood-free inference
- Very broad idea:
  1. Simulate parameters + data
  2. Estimate some relevant density
  3. Use to approximate posterior
- n.b. ABC uses nearest neighbours density estimation!

# Conditional density estimation

- General idea:
  - Simulate many  $(\theta, y)$  pairs from prior and model
  - Estimate joint density  $\hat{\pi}(\theta, y)$
  - Condition on  $y_{\text{obs}}$  and output  $\hat{\pi}(\theta|y_{\text{obs}})$
- Variation:
  - Directly estimate  $\hat{\pi}(\theta|y)$
  - Output  $\hat{\pi}(\theta|y_{\text{obs}})$



## Examples

- Mixture of Gaussians (Bonassi, You and West 2011)
- Random forest (Pudlo et al 2016)
- Extreme gradient boosted tree (Lamperti et al 2017)
- Mixture density neural networks (Papamakarios and Murray 2016, Lueckmann et al 2017)

- General idea:
  - Simulate many  $(\theta, y)$  pairs from prior and model
  - Estimate data density  $\hat{p}(y|\theta)$
  - Likelihood estimate is  $\hat{L}(\theta) = \hat{p}(y_{\text{obs}}|\theta)$
- Examples
  - Kernel density estimation (Diggle and Gratton 1984)
  - Gaussian processes (Wilkinson 2014, Gutmann and Corander 2016)
  - Normalising flows (Papamakarios, Sterrat and Murray 2018)

## Other related methods

- ABC regression correction (Beaumont et al 2002)
  - Run ABC
  - Fit regression to output (predict  $\theta$  given  $y$ )
  - Use to **correct** ABC sample
- Likelihood ratio estimation (Cranmer, Pavez and Louppe 2015)
  - Fit a neural network to estimate **likelihood ratio** given  $\theta_1, \theta_2$
  - (Equivalent to a classification problem)
  - Use in inference e.g. MCMC

- “Learning in implicit generative models” (Mohamed and Lakshminarayanan, 2016)
- “A review of approximate Bayesian computation methods via density estimation: Inference for simulator models” (Grazian and Fan 2019)
- “The frontier of simulation-based inference” (Cranmer Brehmer and Louppe 2019)

# Pros and cons

- Pros
  - Scales up to bigger problems better than ABC
  - Most methods don't need  $\epsilon$  (less tuning!)
  - Reduces need for summary statistics
- Cons
  - Less transparent than ABC
  - Unclear what best method is!
  - Often still needs some dimension reduction by summary statistics
  - Still hard to quantify approx error
  - Little general purpose software yet

# Distilled importance sampling

---

# Motivation

- **ABC** relies on simulator producing close matches to data
- Rare even under best parameter values!
- **Density estimation** methods must estimate densities for wide range of  $y$ s
- Hard modelling task!
- We learn to **control simulator** to always output data close to  $y_{obs}$

- Parameters are  $\theta$
- Let  $u$  be all random draws used in simulator
- Then simulator is function  $y(\theta, u)$
- We try to learn density  $p(\theta, u|y_{\text{obs}})$
- (Tricky as often a near-singular density)



# Main idea

- **Iteration 1**

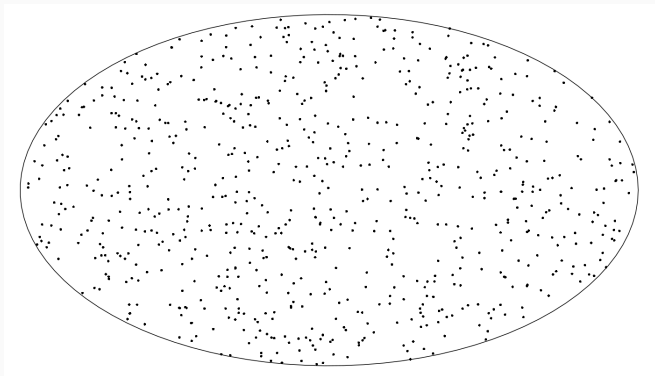
- Sample  $\theta, u$  from prior distributions
- Weight sample depending closeness of  $y(\theta, u)$  to  $y_{\text{obs}}$   
Give moderate weights even to poor matches
- Use weighted sample to train density estimate

- **Iteration 2**

- Sample  $\theta, u$  from density estimate
- Weight sample depending closeness of  $y(\theta, u)$  to  $y_{\text{obs}}$   
Require slightly close matches to get moderate weights
- Use weighted sample to train density estimate
- ...

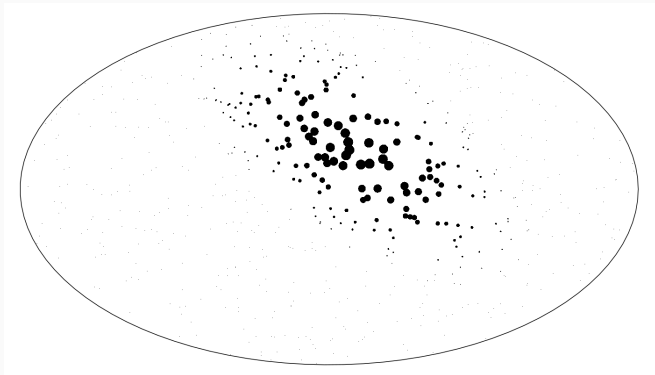
# Illustration

Sample from approximate density



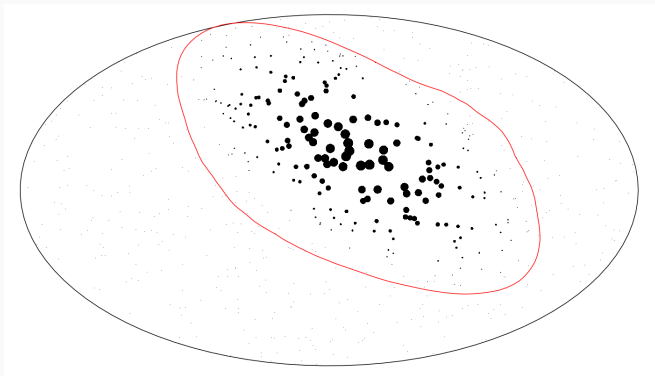
# Illustration

Weight based on closeness

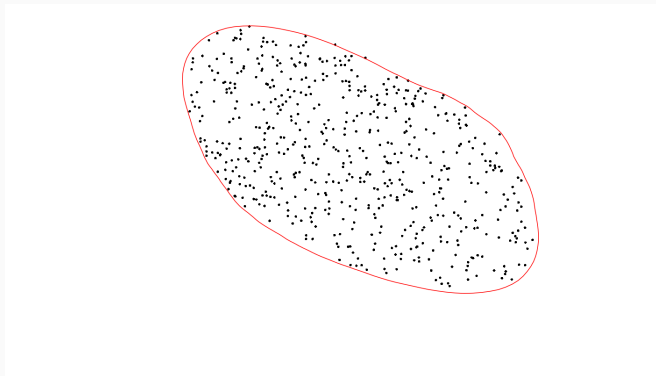


# Illustration

Apply density estimation

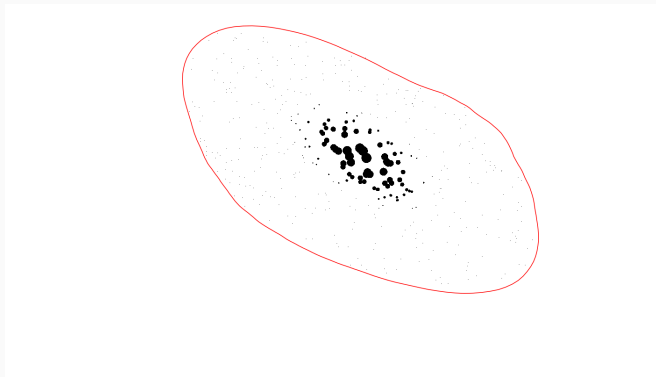


Sample from new approximate density



# Illustration

Weight based on closeness



- Each iteration uses **importance sampling**
- Then uses output in density estimation
- “Distilled importance sampling” (preprint on arxiv)
- Final results still approximate posterior
- Related to:
  - Adaptive importance sampling
  - Sequential importance sampling
  - Cross-entropy method
- We use **normalising flows** for density estimation
- Algorithm reduces to stochastic gradient optimisation
- Similar to variational inference

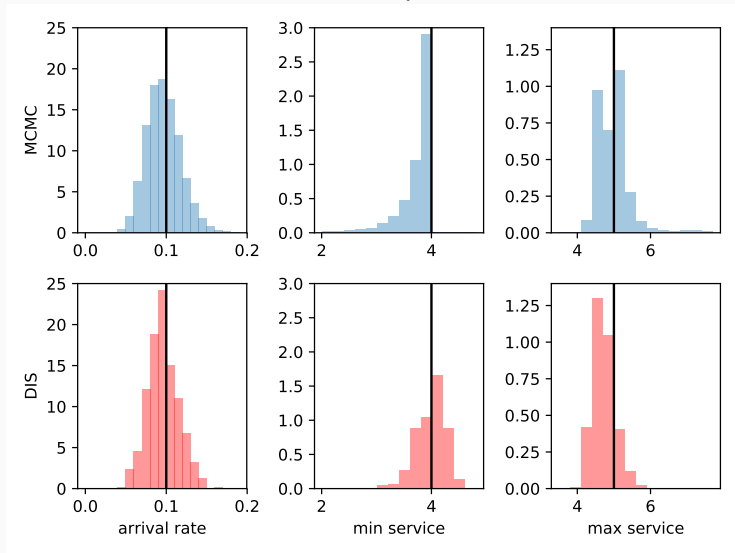
## Queue example

- Need two  $u$  variables for individual in queue
- Seed for arrival time
- Seed for departure time
- Overall: 40  $u$  variables and 3  $\theta$  variables



# MG1 queue results

- Inference time: 3 hours on desktop PC, final  $\epsilon = 0.283$



## Conclusion

---

# Summary

- **Generative models** allow data simulation but not easy likelihood calculation
- **Likelihood-free inference** does **approximate** inference just using simulations
- **ABC** simulates data under many parameter values looking for good matches
- Alternative: use simulations in modern **density estimation** methods
- My recent work improves efficiency by learning to **control simulator**