Bayesian parameter synthesis for Markov population models

Phung Nhat-Huy

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Bayesian parameter synthesis for Markov population models.

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May 17, 2021

Universität Konstanz



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Motivation

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Motivation

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- We study the population dynamics of a system of interest. For example:
 - Number of online nodes in a computer network.
 - Number of surviving individuals in an epidemic model.
- We study the population in a grey-box setup
 - Estimating the model's unknown attributes with experiment data of the population at its steady state.

Motivation

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- ▶ Modeling population using a stochastic process (Markov population model [12])
 - Discrete-time Markov chain
- Parameterization: encoding the unknown attributes of a system by model parameters
 - Parametric discrete-time Markov chain [10]).

Motivation

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As parameters represent unknown features of the system, it gives the following research questions

- (Parameter inference): Given a set of data collected by observing the system, what can we know about its parameters?
- (Parameter synthesis): Which values of parameters instantiate a model that satisfies a specific property of interest?

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Probabilistic model checking

Discrete-time Markov chain

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Definition (Discrete Time Markov Chain [1])

A Discrete-time Markov chain (or DTMC in short) \mathcal{M} is a tuple $(S, P, \iota_{init}, AP, L)$, in which

- \triangleright S is a countable, non-emty set of states
- ▶ $P: S \times S \rightarrow [0,1]$ is the *transition probability* function such that

$$\forall s \in S : \sum_{s' \in S} \mathbf{P}(s, s') = 1$$

 \triangleright $\iota_{init}: S \rightarrow [0,1]$ is the *initial distribution* such that

$$\sum_{s \in S} \iota_{init}(s) = 1$$

- ► *AP* is a set of *atomic propositions*.
- ▶ $L: S \to 2^{AP}$ is the labelling function on states.

Bottom Strongly Connected Components

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Definition (Strongly Connected Component)

Let $\mathcal{M} = (S, P, \iota_{init}, AP, L)$ be a DTMC. A subset $S' \subset S$ is strongly connected if and only if for every pair $s_1, s_2 \in S'$ there is a path between s_1 and s_2 which consists of only states in S'. If S' has no superset $S'' \subseteq S$, such that S'' is strongly connected, then S' is a Strongly Connected Component, or SCC in short.

Definition (Bottom Strongly Connected Component)

Let $\mathcal{M} = (S, \mathbf{P}, \iota_{init}, AP, L)$ be a DTMC and $S' \in S$ a Strongly Connected Component. S' is also a Bottom Strongly Connected Component (or BSCC in short), if and only if there exist no state $s \in S \setminus S'$ that is reachable from any state in S'. If |S'| = 1 then S' is a trivial BSCC. We denote $BSCC(\mathcal{M}) \in S$ is the set of all BSCCs of \mathcal{M} .

Example of DTMC

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Algorithm by Knuth and Yao [13] to model a fair dice by a fair coin.

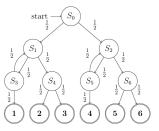


Figure: DTMC model of Knuth-Yao die. There are six BSCCs labbeled "1" to "6"

Probabilistic Computational Tree Logic

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Definition (PCTL [1])

The syntax of PCTL consists of state formulas and path formulas.

State formulas are defined over AP

$$\Phi ::= \mathrm{true} \mid a \mid \Phi \mid \Phi_1 \wedge \Phi_2 \mid \Phi_1 \vee \Phi_2 \mid P_J(\phi)$$

where $a \in AP$, ϕ is a path formula, and $J \subseteq [0,1]$ is an interval.

▶ Path formulas

$$\phi ::= \bigcirc \Phi \mid \Phi_1 \mathsf{U} \Phi_2 \mid \Phi_1 \mathsf{U}^{\leq n} \Phi_2$$

where Φ, Φ_1, Φ_2 are state formulas, and $n \in \mathbb{N}$.

Example of DTMC

Bayesian parameter synthesis for Markov population models

Preliminaries

Algorithm by Knuth and Yao [13] to model a fair dice by a fair coin.

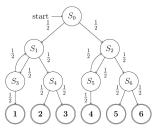


Figure: DTMC model of Knuth-Yao die. There are six BSCCs labbeled "1" to "6"

The probability that the simulation eventually ends with the outcome "one dot" is equal to $\frac{1}{6}$:

$$P_{=\frac{1}{6}}(\text{TrueU"1"})$$

Parametric discrete-time Markov chain

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Reference

Definition (Parametric discrete-time Markov chain [8])

A parametric discrete-time Markov chain \mathcal{M}_{θ} is a tuple $(S, \theta, \mathbf{P}, \iota_{init}, AP, L)$ where

- \triangleright S is a countable, non-emty set of states
- ▶ $\theta \in \mathbb{R}^n$, $n \in \mathbb{N}$ as the set of parameters.
- $ightharpoonup P: S \times S \to \mathbb{Q}(x)$ is the transition probability function such that

$$\forall s \in S : \sum_{s' \in S} \mathbf{P}(s, s') = 1$$

 $\iota_{init}: S \to [0,1]$ is the initial distribution such that

$$\sum_{s \in S} \iota_{init}(s) = 1$$

- ► AP is a set of atomic propositions
- ▶ $L: S \to 2^{AP}$ is the labelling function on states.

Example of DTMC

Bayesian parameter synthesis for Markov population models.

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Algorithm by Knuth and Yao [13] to model a possibly unfair dice by two possibly unfair coins.

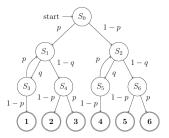


Figure: DTMC model of Knuth-Yao die with two possibly unfair coins. There are six BSCCs labbeled "1" to "6"

Parameter synthesis

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Definition (Parameter synthesis (Katoen [10]))

Given a finite-state parametric Markov model, find the parameter values for which a given reachability property exceeds (or is below) a given threshold β .

Parameter synthesis

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Katoen [9] summarizes the following methods on parameter synthesis of parametric DTMC:

- **1** Computing symbolic reachability probabilities (Daws [4], Hahn [7]): symbolic solving system of linear equations.
- 2 Candidate region generation and checking (Kwiatkowska [14]): partition the parameter space into safe and unsafe regions.
- 3 Parameter lifting ([15]) replace parametric transition system by a non-parametric one with transition labels are bounds from given intervals.

Example of DTMC

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Given DTMC model $\mathcal{M}_{(p,q)}$ as in 3 and a property

$$\Phi = \textit{P}_{\geq 0.2}(\texttt{TrueU"1"})$$

A Monte Carlo sampling using $p, q \sim \textit{Uniform}(0, 1)$ gives the following satisfying parameter values

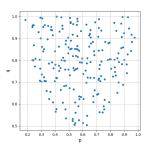


Figure: Samples of (p, q) that instantiate $\mathcal{M}_{(p,q)} \models \Phi$.

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Bayesian inference

Bayes theorem

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Definition (Bayes theorem)

$$\pi(\theta|D_{obs}) = rac{P(D_{obs}|\theta)\pi(\theta)}{\int_{ heta} P(D_{obs}|\theta)\pi(\theta)d\theta}$$

where

- $\triangleright \pi(\theta)$ is the prior distribution.
- $ightharpoonup P(D_{obs}|\theta)$ is the *likelihood*.
- $ightharpoonup \int_{\theta} P(D_{obs}|\theta)\pi(\theta)d\theta$ is the marginal distribution.
- $\blacktriangleright \pi(\theta|D_{obs})$ is the posterior distribution

Bayesian parameter estimation

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With posterior distribution $\pi(\theta|D_{obs})$ we estimate the parameter $\hat{\theta}$ using Bayesian posterior mean.

$$\hat{ heta} = \mathbf{E}[heta] = \int_{ heta} heta \pi(heta|D_{obs}) d heta$$

In case we have samples from posterior distribution, for example a set of N parameter values $(\theta_1, \ldots, \theta_N)$ sampled from the posterior distribution $\pi(\theta|D_{obs})$, the discrete form of posterior mean is used:

$$\hat{ heta} pprox \mathbf{E}[heta] pprox \sum_{a} heta \pi(heta|D_{obs})$$

Approximation of posterior distribution

Bayesian parameter synthesis for Markov population models

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Reference

- Usually posterior distribution $\pi(\theta|D_{obs})$ has no analytical to evaluate
- We use sampling algorithms to draw samples from posterior distribution
 - Metropolis-Hastings
 - Sequential Monte-Carlo
 - Approximate Bayesian computation

Metropolis-Hastings

Bayesian parameter synthesis for Markov population models.

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Algorithm Metropolis-Hastings Algorithm

```
procedure Metropolis-Hastings
 2:
          Init empty trace
 3:
          Init \theta_1 from \pi(\theta)
 4:
         while i < N_{MH} do
 5:
              Draw \theta_{cand} from Q(\theta'|\theta_{i-1})
              Compute \xi = \min(0, \ln(P(D_{obs}|\theta_{cand})) - \ln(P(D_{obs}|\theta_{i-1})))
 6:
 7:
              if \xi > 0 then
 8:
                   Accept \theta_{cand}, append to trace.
 9:
              else
                   Draw a random number u from Uniform(0,1)
10:
11:
                  if u < \exp(\xi) then
12:
                       Accept \theta_{cand}, append to trace.
13:
          Return (\theta_1, \ldots, \theta_N), (w_1, \ldots, w_N)
```

Metropolis-Hastings

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Reference

Advantages of Metropolis-Hastings are:

- + Parameter transition only needs the computation of the likelihood function.
- Computationally efficient, as marginal distribution is canceled out, and likelihood can be replaced by log-likelihood.
- + Simple to implement.

Disadvantages of Metropolis-Hastings are

- Highly probable to be stuck in a local maximum or minimum.
- Not parallelizable

Sequential Monte-Carlo

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```
Algorithm Sequential Monte Carlo Algorithm
```

```
1: procedure Sequential-Monte Carlo
 2:
            Draw (\theta_1, \ldots, \theta_N) from \pi(\theta)

▷ SMC initialization

 3:
           t \leftarrow 1
 4:
           while t \leq M do
 5:
                 Normalize (w_1^{t-1}, \ldots, w_N^{t-1})

⊳ SMC correction step

                 Sample with replacement (\theta_1^t, \dots, \theta_N^t) > SMC selection step
 6:
                       from (\theta_1^{t-1}, \dots, \theta_N^{t-1}) with probabilities (w_1^{t-1}, \dots, w_N^{t-1})
 7:
                 i \leftarrow 1
 8:
                 while i < N do

⊳ SMC perturbation step

                      Draw \hat{\theta}_i^t from F_t(\theta^t | \theta_1^{t-1}, \dots, \theta_N^{t-1}), 1 < t < M
 9:
                      Mutate (\theta_1^*, \dots, \theta_{N_{MU}}^*), (w_1^*, \dots, w_{N_{MU}}^*) \leftarrow MH(\hat{\theta}_i^t)
10:
                      \theta_i \leftarrow \theta_{N_{MH}}^*
11:
                      w_i \leftarrow w_{N_{MH}}^*
12:
13:
            Return (\theta_1, \ldots, \theta_N), (w_1, \ldots, w_N)
```

Sequential Monte-Carlo

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Sequential Monte Carlo algorithm has several advantages compared to Metropolis-Hastings algorithm.

- + Approximate multimodal distributions: *N* particles moving independently.
- + Parallelizable.

However, Sequential Monte Carlo also has disadvantages:

- Selection of perturbation and transition kernel is not trivial (Filippi [6], and Silk [16]).
- More difficult to implement.

Approximate Bayesian Computation

Bayesian parameter synthesis for Markov population models

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A likelihood-free method

- used when likelihood has no analytical form, or the analytical form is expensive to be evaluated.
- estimates the likelihood $P(D_{obs}|\theta)$, or replace it by other measures.

Approximate Bayesian Computation

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References

Algorithm Approximate Bayesian Computation

```
procedure Approximate-Bayesian-Computation
 2:
          Select a proposal distribution \pi(\theta)
 3:
          i \leftarrow 1
 4:
          while i < N do
 5:
               Draw a random particle \theta from \pi(\theta)
 6:
               Simulate data D_{sim} from \mathcal{M}_{\theta}
 7:
               if d = \delta(D_{sim}, D_{obs}) < \epsilon then
                    \theta_i \leftarrow \theta
 8:
 9:
                    w_i = d
           Return (\theta_1, \ldots, \theta_N), (w_1, \ldots, w_N)
10:
```

Approximate Bayesian Computation

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Reference

Advantages of Approximate Bayesian Computation are:

- + Likelihood-free: applicable when the likelihood has no analytical form or there is no likelihood.
- + Easy to implement.

However, Approximate Bayesian Computation has drawbacks:

- How to select a distance threshold ϵ so that the posterior is closely approximated? [17]
- How to choose a summary statistic to capture sufficient information? [3]

Parameter synthesis and Parameter inference

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Reference

What are the differences between parameter inference and parameter synthesis?

	Input	Ouput
Parameter	Model $\mathcal{M}_{ heta}$	Parameter estimation
inference	Observed data D_{obs}	$\mid \hat{ heta} \mid$
Parameter synthesis	Model \mathcal{M}_{θ} Reachability property Φ	$egin{aligned} (heta_1,\ldots, heta_N)\ orall heta_i \in (heta_1,\ldots, heta_N):\ \mathcal{M}_{ heta_i} \models \Phi \end{aligned}$

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Framework design

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- In this thesis, we combine parameter synthesis and parameter inference into a data-informed parameter synthesis framework.
 - Sample a set of parameter values which satisfy a property of interest (parameter synthesis) and use it to estimate a parameter value which the observed data is likely to be simulate from (parameter inference).

Framework design

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Reference

- Given experiment data of a system at its steady state and a property of interest
 - Bayesian parameter inference: apply different sampling algorithm to approximate the posterior distribution of parameter.
 - Parameter synthesis: only accept the sampled points which satisfy the property of interest.
 - Paramter estimation: from sampled point, compute an estimation of model parameter.

Challenges

Bayesian parameter synthesis for Markov population models.

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State explosion

The state-explosion problem occurs when the size of a model state space grows exponentially as the number of state variables in the system increases [2].

The state-space explosion problem renders probabilistic model checking computationally expensive. We cope this problem using 2 different strategies:

- Pre-compute symbolic reachability probability (rational function).
- Estimate reachability probability statistically (statistical model checking).

Challenges

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Approximation of posterior distribution

In Bayesian parameter inference, the posterior distribution usually does not have an analytical form. Hence, we approximate the posterior distribution using different sampling algorithms.

Challenges are:

- ► How to select a sampling algorithm?
- ▶ How to select parameters for the sampling algorithm?

Contribution

Bayesian parameter synthesis for Markov population models

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Reference

- Designed and implemented a data-informed, Bayesian framework on parameter synthesis of parametric Discrete-time Markov chain.
 - when the exact likelihood function of the property of interest is available RF-SMC, and
 - when it has to be approximated by simulations SMC-ABC-SMC.

Contribution

Bayesian parameter synthesis for Markov population models

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References

- Designed and implemented a data-informed, Bayesian framework on parameter synthesis of parametric Discrete-time Markov chain.
 - ▶ when the exact likelihood function of the property of interest is available *RF-SMC*, and
 - when it has to be approximated by simulations SMC-ABC-SMC.
- Compared the performances (accuracy and scalability) of proposed frameworks on different case studies and different model state-space sizes.

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Frameworks

Frameworks

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Framework

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Conclusion

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- ► Generic framework
- ► RF-SMC
- ► SMC-ABC-SMC

Generic framework

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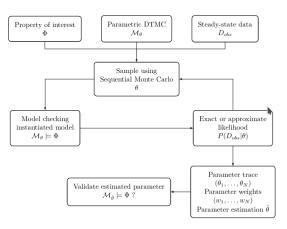


Figure: Generic framework for Bayesian parameter synthesis of parametric DTMC.

Generic framework

Bayesian parameter synthesis for Markov population models.

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References

Input:

- $ightharpoonup \mathcal{M}_{ heta}$: parametric DTMC of parameter heta
- Φ: bounded reachability property of interest.
- $ightharpoonup D_{obs}$: observed data.
- N: number of particles.

Output:

- \bullet $(\theta_1, \ldots, \theta_{N_{MH}})$, sampled particles
- $(w_1, \ldots, w_{N_{MH}})$ particle corresponding weights.
- $\hat{p} = P(\mathcal{M}_{\hat{\theta}} \models \Phi)$

Generic framework

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```
Algorithm Generic framework for Bayesian parameter synthesis
```

```
procedure Generic-Bayesian-Monte-Carlo
 2:
           i \leftarrow 1
 3:
          while i < N do
 4:
                Sample \theta with corresponding weight w
                                by Sequential Monte Carlo sampling algorithm.
 5:
               Verify instantiated model \mathcal{M}_{\theta} against \Phi
 6:
               if \mathcal{M}_{\theta} \models \Phi then
 7:
                    \theta : \leftarrow \theta
                    Estimate w_i as exact or approximated likelihood P(D_{obs}|\theta)
 8:
           Estimate \hat{\theta} using posterior mean.
 9:
           Compute \hat{p} = P(\mathcal{M}_{\hat{a}} \models \Phi)
10:
           Return (\theta_1, \ldots, \theta_N), (w_1, \ldots, w_N), \hat{\theta}, \hat{p}
11:
```

RF-SMC

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- Based on Sequential Monte-Carlo
- Model checking using precomputed symbolic reachability probability at each SMC perturbation step.
 - At each MH transition, a particle is accepted only if it instantiates a satisfying model.

RF-SMC

Bayesian parameter synthesis for Markov population models.

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Algorithm Metropolis-Hastings with rational functions

```
1: procedure RF-MH
 2:
           Init empty trace T
 3:
           Draw \theta_{cand} from \pi(\theta) s.t \mathcal{M}_{\theta_{cand}} \models \Phi (evaluating RF_{\Phi}(\theta))
 4:
           Append \theta_{cand} to trace T
 5:
           i \leftarrow 2
 6:
           while i < N_{MH} do
 7:
                Draw \theta_{cand} from Q(\theta'|\theta_{i-1}) s.t \mathcal{M}_{\theta_{cand}} \models \Phi (evaluating RF_{\Phi}(\theta))
 8:
                Evaluate \xi = \min(0, \ln(P(D_{obs}|\theta_{cand})) - \ln(P(D_{obs}|\theta_{i-1}))) > 0
 9:
                if \xi > 0 then
10:
                     Append \theta_{cand} to trace T
11:
                else
12:
                     Draw a random number u from Uniform(0,1)
13:
                     if u < \exp(\xi) then
14:
                          Append \theta_{cand} to trace T
           Return (\theta_1, \ldots, \theta_{N_{MH}}), (w_1, \ldots, w_{N_{MH}})
15:
```

RF-SMC

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Algorithm Sequential Monte Carlo with rational functions

```
1: procedure RF-SMC
            Draw (\theta_1, \ldots, \theta_N) from \pi(\theta) s.t \mathcal{M}_{\theta_i} \models \Phi (evaluating RF_{\Phi}(\theta))
 2:
 3:
            t \leftarrow 1
 4:
           while t \leq M do
                                                                ▷ SMC correction step
 5:
                 Normalize (w_1^{t-1}, \ldots, w_N^{t-1})
 6:
                 Sample with replacement (\theta_1^t, \dots, \theta_N^t) > SMC selection step
                        from (\theta_1^{t-1}, \dots, \theta_N^{t-1}) with probabilities (w_1^{t-1}, \dots, w_N^{t-1})
 7:
                 i \leftarrow 1
 8:
                 while i \leq N do
                                                                            Draw \hat{\theta}_i^t from F_t(\theta^t | \theta_1^{t-1}, \dots, \theta_N^{t-1}), 1 < t < M
 9.
                       Mutate (\theta_1^*, \dots, \theta_{N_{MU}}^*), (w_1^*, \dots, w_{N_{MU}}^*) \leftarrow RF - MH(\hat{\theta}_i^t)
10:
                       \theta_i, w_i \leftarrow \theta^*_{N_{MH}}, w^*_{N_{MH}}
11:
12:
            Estimate \hat{\theta} using posterior mean, compute \hat{p} = P(\mathcal{M}_{\hat{a}} \models \Phi)
            Return (\theta_1, \ldots, \theta_N), (w_1, \ldots, w_N), \hat{\theta}, \hat{p}
13:
```

SMC-ABC-SMC

Bayesian parameter synthesis for Markov population models

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Reference

Without the availability of analytical form to evaluate the steady-state distribution and the property of interest, we face the following obstacles:

- Absence of likelihood functions
 - no analytical form of likelihood.
 - → likelihood-free methods (Approximate Bayesian Computation)
- Absence of rational functions for evaluation of property of interest
 - ▶ ⇒ Statistical model checking

SMC-ABC-SMC

Bayesian parameter synthesis for Markov population models.

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Algorithm Sequential Monte-Carlo with simulations

```
procedure SMC-ABC-SMC
 2:
            Draw (\theta_1, \ldots, \theta_N) from \pi(\theta) s.t \mathcal{M}_{\theta_i} \models \Phi (Statistical MC)
 3:
            t \leftarrow 1
 4:
            while t \leq M do

⊳ SMC correction step

 5:
                  Normalize (w_1^{t-1}, \ldots, w_N^{t-1})
                  Sample with replacement (\theta_1^t, \dots, \theta_N^t) \triangleright SMC selection step
 6:
                         from (\theta_1^{t-1}, \dots, \theta_N^{t-1}) with probabilities (w_1^{t-1}, \dots, w_N^{t-1})
 7:
                 i \leftarrow 1
 8:
                 while i < N do

▷ SMC perturbation step

                       Draw \hat{\theta}_i^t from F_t(\theta^t | \theta_1^{t-1}, \dots, \theta_N^{t-1}), 1 < t < M
 9:
                       if Statistical Model Checking \mathcal{M}_{\hat{\theta}^t} \models \Phi then
10:
11:
                             Simulate D_{sim} from \mathcal{M}_{\hat{\theta}^t}
12:
                             if \delta = Distance(D_{sim}, D_{obs}) < \epsilon) then
                                  \theta: w: \leftarrow \hat{\theta}^t: \delta^{-1}
13:
            Estimate \hat{\theta} using posterior mean, compute \hat{p} = P(\mathcal{M}_{\hat{\theta}} \models \Phi)
14:
15:
            Return (\theta_1, \ldots, \theta_N), (w_1, \ldots, w_N), theta, \hat{p}
```

Comparison

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```
Algorithm 7 Sequential Monte-Carlo with simulations
                                                                                                            Algorithm 6 Sequential Monte Carlo with rational functions
  1: procedure SMC-ABC-SMC
                                                                                                              1: procedure RF-SMC
           Draw (\theta_1, ..., \theta_N) from \pi(\theta) s.t \mathcal{M}_{\theta_i} \models \Phi (Statistical MC)
                                                                                                                         Draw (\theta_1, \dots, \theta_N) from \pi(\theta) s.t \mathcal{M}_{\theta_i} \models \Phi (evaluating RF_{\Phi}(\theta))
           while t \leq M do
 4
                                                                                                                        while t \leq M do
                Normalize (w_1^{t-1}, \dots, w_N^{t-1})

⊳ SMC correction step

                                                                                                                               Normalize (w_1^{t-1}, \dots, w_N^{t-1})

⊳ SMC correction ster

                Sample with replacement (\theta_1^t, \dots, \theta_N^t) \triangleright SMC selection step
                                                                                                                               Sample with replacement (\theta_1^t, \dots, \theta_N^t) \triangleright SMC selection step
                      from (\theta_1^{t-1}, \dots, \theta_n^{t-1}) with probabilities (w_1^{t-1}, \dots, w_n^{t-1})
                                                                                                                                     from (\theta_1^{t-1}, \dots, \theta_N^{t-1}) with probabilities (w_1^{t-1}, \dots, w_N^{t-1})
                i \leftarrow 1
 8.
                while i \le N do

⊳ SMC perturbation step

                                                                                                                               i \leftarrow 1
                     Draw \hat{\theta}_i^t from F_t(\theta^t | \theta_1^{t-1}, \dots, \theta_N^{t-1}), 1 \leq t \leq M
 9:
                                                                                                              8:
                                                                                                                               while i \le N do

⇒ SMC perturbation ster

                                                                                                                                    Draw \hat{\theta}_t^i from F_t(\theta^t | \theta_1^{t-1}, \dots, \theta_N^{t-1}), 1 \le t \le M
                     if Statistical Model Checking \mathcal{M}_{\hat{a}t} \models \Phi then
10:
11:
                          Simulate D_{sim} from M_{\tilde{a}t}
                                                                                                                                    Mutate (\theta_1^*, \dots, \theta_{N_{MH}}^*), (w_1^*, \dots, w_{N_{MH}}^*) \leftarrow RF - MH(\hat{\theta}_i^t)
                                                                                                             10:
12:
                          if \delta = Distance(D_{sim}, D_{obs}) < \epsilon) then
                                                                                                                                    \theta_i, w_i \leftarrow \theta^*_{Nuu}, w^*_{Nuu}
                                                                                                             11:
13:
                                \theta_i, w_i \leftarrow \hat{\theta}^t, \delta^{-1}
                                                                                                            12:
                                                                                                                         Estimate \hat{\theta} using posterior mean, compute \hat{p} = P(\mathcal{M}_{\hat{\theta}} \models \Phi)
14:
           Estimate \hat{\theta} using posterior mean, compute \hat{p} = P(\mathcal{M}_{\hat{a}} \models \Phi)
                                                                                                             13-
                                                                                                                         Return (\theta_1, \dots, \theta_N), (w_1, \dots, w_N), \hat{\theta}, \hat{\rho}
           Return (\theta_1, \dots, \theta_N), (w_1, \dots, w_N), theta, \hat{p}
15:
```

Figure: Comparison between RF-SMC and SMC-ABC-SMC.

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Case studies:

- Zeroconf
- Social feedback in honeybee colonies
- ► SIR

Evaluation environment:

- Hardware: Intel Xeon W-2135, 64GB RAM
- ► OS: OpenSUSE 15.2
- ▶ Libraries: Storm @stable, PRISM 4.6, Python 3.8.8

Zeroconf

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Zero-configuration protocol (zeroconf for short) [5] is a protocol used in IPv4 network to allocate newly attached device an unique IP address without any intervention from network operators.

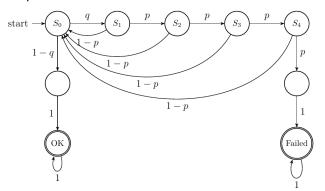


Figure: Example of an IPv4 Zeroconf model with 4 probes

Zeroconf

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work

We select a true parameter (p, q) arbitrarily random for Zeroconf model of 4 and 10 states.

Model ${\mathcal M}$	Zeroconf, 4 probes	Zeroconf, 10 probes
Number of BSCCs	2	2
Number of states	9	14
True parameter $\theta = (p, q)$	(0.105547, 0.449658)	(0.197779, 0.621824)
Number of samples	10000	10000
Synthetic data D_{obs}	(41, 9959)	(22, 9978)
Property of interest Φ	$P_{\geq 0.75}(\mathtt{trueU}^{\leq 4}(\mathtt{"OK"}))$	$P_{\geq 0.75}({ t trueU}^{\leq 10}("0K"))$
Satisfaction property $P(\mathcal{M}_{\theta} \models \Phi)$	0.946409	0.952067

Table: Synthetic data for Zeroconf model of 4 and 10 probes.

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Method	RF-SMC	SMC-ABC-SMC
Estimated parameter $\hat{ heta}$	(0.188956, 0.460554)	(0.176469, 0.355322)
True parameter θ	(0.105547, 0.449658)	(0.105547, 0.449658)
L2 distance $\ \theta,\hat{\theta}\ _2$	0.084117	0.118023
$P(\mathcal{M}_{\hat{\theta}} \models \Phi)$	0.893715	0.918133

Table: Parameter estimation results for Zeroconf model of 4 probes.

Zeroconf 4 results

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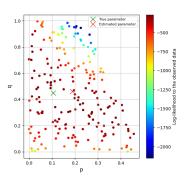
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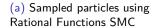
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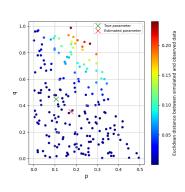
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(b) Sampled particles using Statiscal Model Checking ABC-SMC

Figure: Parameter synthesis results for Zeroconf model of 4 probes.

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Runtime

Method	RF-SMC	SMC-ABC-SMC
Total runtime (minutes)	6.083	54.867
Average perturbation runtime (minutes)	0.32	2.88

Table: Physical runtime on Zeroconf model with 4 probes.

Zeroconf 10 results

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Method	RF-SMC	SMC-ABC-SMC
True parameter θ	(0.197779, 0.621824)	(0.197779, 0.621824)
Estimated parameter $\hat{ heta}$	(0.301807, 0.457090)	(0.378774, 0.405870)
L2 distance $\ heta,\hat{ heta}\ _2$	0.194831	0.281772
$P(\mathcal{M}_{\hat{ heta}} \models \Phi)$	0.952067	0.966142

Table: Parameter estimation results for Zeroconf model of 10 probes.

Zeroconf 10 results

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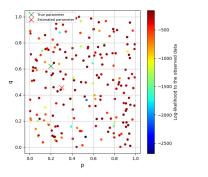
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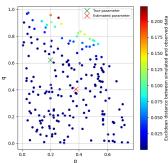
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(a) Sampled particles using RF-SMC

(b) Sampled particles using SMC-ABC-SMC

Figure: Parameter synthesis results for Zeroconf model of 10 probes.

Zeroconf 10 results

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Runtime

Method	RF-SMC	SMC-ABC-SMC
Total runtime (minutes)	9.50	37.93
Average perturbation runtime (minutes)	0.501	1.978

Table: Physical runtime on Zeroconf model with 10 probes.

Zeroconf results discussion

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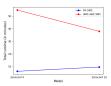
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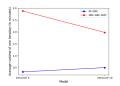
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(a) Total runtime



(b) Average perturbation runtime

Figure: Physical runtime on Zeroconf model of different sizes.

- ▶ RF-SMC and SMC-ABC-SMC have similar accuracy.
- ▶ RF-SMC is much faster

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Each bee in a colony possibly stings after observing a threat in the surrounding environment and warns other bees by releasing a special substance, pheromone. There are 3 assumptions on the system:

- Each bee releases a unit amount of pheromone immediately after stinging.
- A bee dies after stinging and releasing a pheromone unit. In other words, no bee can sting more than once.
- **3** Stinging behavior only depends on the concentration of pheromone in the environment.

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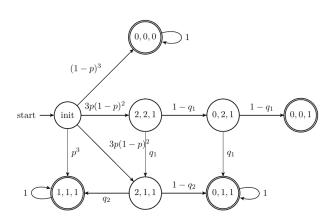


Figure: Parametric DTMC model of 3 bees with 3 parameters p, q_1 , q_2

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True parameters and synthetic data.

Model M	3 bees	5 bees	10 bees
Number of states	13	24	69
Number of BSCCs	4	6	11
True parameter $ heta$	$ \begin{array}{l} p = 0.665623 \\ q_1 = 0.830401 \\ q_2 = 0.839778 \end{array} $	$p = 0.278370$ $q_1 = 0.305994$ $q_2 = 0.489792$ $q_3 = 0.737252$ $q_4 = 0.766581$	$\begin{array}{l} p = 0.222169 \\ q_1 = 0.246993 \\ q_2 = 0.281934 \\ q_3 = 0.446384 \\ q_4 = 0.491612 \\ q_5 = 0.534611 \\ q_6 = 0.569409 \\ q_7 = 0.684651 \\ q_8 = 0.717139 \\ q_9 = 0.800987 \end{array}$
Synthetic data D_{obs}	(344, 54, 1390, 8212)	(1940, 11, 216, 2682, 4200, 951)	(769, 0, 1, 10, 187, 972, 2494, 2982, 2133, 419, 33)
Target property Φ	$P_{\geq 0.25}(\text{trueU}(S > 3))$	$P_{\geq 0.25}(\text{trueU}(S > 5))$	$P_{\geq 0.25}(\text{trueU}(S > 8))$
$P(\mathcal{M}_{\theta} \models \Phi)$	0.819666	0.780172	0.737244

Table: True parameter and synthetic data for 3, 5, and 10 bees models.

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	RF-SMC	SMC-ABC-SMC
	p = 0.665623	p = 0.665623
True parameter $ heta$	$q_1 = 0.830401$	$q_1 = 0.830401$
	$q_2 = 0.839778$	$q_2 = 0.839778$
_	p = 0.671388	p = 0.811651
Estimated parameter $\hat{ heta}$	$q_1 = 0.575026$	$q_1 = 0.621073$
	$q_2 = 0.525502$	$q_2 = 0.544130$
L2 distance $\ \theta, \hat{\theta}\ _2$	0.404992	0.390576
$P(\mathcal{M}_{\hat{\theta}} \models \Phi)$	0.623889	0.595083

Table: Parameter synthesis result for 3 bees model.

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Method	RF-SMC	SMC-ABC-SMC
Total runtime (minutes)	5.917	68.783
Average perturbation runtime (minutes)	0.312	3.614

Table: Physical runtime on 3 bees model.

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	RF-SMC	SMC-ABC-SMC
	p = 0.278370	p = 0.278370
	$q_1 = 0.305994$	$q_1 = 0.305994$
True parameter $ heta$	$q_2 = 0.489792$	$q_2 = 0.489792$
	$q_3 = 0.737252$	$q_3 = 0.737252$
	$q_4 = 0.766581$	$q_4 = 0.766581$
	p = 0.576565	p = 0.361220
_	$q_1 = 0.589724$	$q_1 = 0.316007$
Estimated parameter $\hat{ heta}$	$q_2 = 0.490334$	$q_2 = 0.545691$
	$q_3 = 0.554397$	$q_3 = 0.643962$
	$q_4 = 0.524433$	$q_4 = 0.591206$
L2 distance $\ \theta, \hat{\theta}\ _2$	0.511366	0.222594
$P(\mathcal{M}_{\hat{\theta}} \models \Phi)$	0.623889	0.595083

Table: Parameter synthesis result for 5 bees model

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Method	RF-SMC	SMC-ABC-SMC
Total runtime (minutes)	29.517	352.083
Average perturbation runtime (minutes)	1.553	18.518

Table: Physical runtime on 5 bees model.

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	RF-SMC	SMC-ABC-SMC
	p = 0.222169	p = 0.222169
	$q_1 = 0.246993$	$q_1 = 0.246993$
	$q_2 = 0.281934$	$q_2 = 0.281934$
	$q_3 = 0.446384$	$q_3 = 0.446384$
T	$q_4 = 0.491612$	$q_4 = 0.491612$
True parameter θ	$q_5 = 0.534611$	$q_5 = 0.534611$
	$q_6 = 0.569409$	$q_6 = 0.569409$
	$q_7 = 0.684651$	$q_7 = 0.684651$
	$q_8 = 0.717139$	$q_8 = 0.717139$
	$q_9 = 0.800987$	$q_9 = 0.800987$
	p = 0.604881	p = 0.391313
	$q_1 = 0.472557$	$q_1 = 0.485688$
	$q_2 = 0.281484$	$q_2 = 0.424056$
	$q_3 = 0.500706$	$q_3 = 0.381489$
Estimated parameter $\hat{ heta}$	$q_4 = 0.49340$	$q_4 = 0.440681$
Estillated parameter 0	$q_5 = 0.495508$	$q_5 = 0.578865$
	$q_6 = 0.466596$	$q_6 = 0.594232$
	$q_7 = 0.510167$	$q_7 = 0.564557$
	$q_8 = 0.474153$	$q_8 = 0.547804$
	$q_9 = 0.484061$	$q_9 = 0.520006$
L2 distance $\ \theta, \hat{\theta}\ _2$	0.665837	0.487042
$P(\mathcal{M}_{\hat{\theta}} \models \Phi)$	0.933287	0.907478

Table: Parameter synthesis result for 10 bees model

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Method	RF-SMC	SMC-ABC-SMC
Total runtime (minutes)	3976.117	581.833
Average perturbation runtime (minutes)	209.237	30.592

Table: Physical runtime on 10 bees model.

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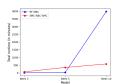
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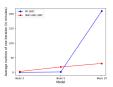
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(a) Total runtime



(b) Average perturbation runtime

Figure: Physical runtime on bees model of different sizes.

Results discussion

- ► SMC-ABC-SMC delivers results with higher accuracy and comparable satisfaction probability.
- ► From 10 bees model, RF-SMC becomes much slower than SMC-ABC-SMC.



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SIR model (Kermack [11]) is a model widely used in modeling epidemics. In a SIR model, each individual is of one among three types:

- Susceptible (S)
- ► Infected (I)
- ► Recovered (R)

SIR is a stochastic system modeled by reactions between S, I and R. In this thesis, we use only two reactions.

$$R_1: S+I \xrightarrow{\alpha} 2I$$

$$R_2:I\xrightarrow{\beta}R$$

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Reactions R_1 , R_2 generate continuous-time Markov chain (CTMC) depends on the initial population

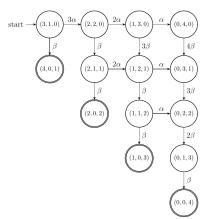


Figure: SIR(3,1,0) CTMC model with parameters(α, β).

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We uniformize CTMC into DTMC

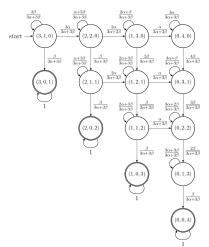


Figure: SIR(3,1,0) uniformized DTMC model with uniformization

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As uniformization of a CTMC preserves bounded until properties, we conduct parameter synthesis experiments on uniformized DTMC. We verify the following property

SIR property of interest

"With the probability of at least 25 percent, the number of infected individuals does not exceed half of the population until the system is in its steady-state. Let $N = S_0 + I_0 + R_0$. In the PCTL formula we have:

$$P_{\geq 0.25}(!(i > N/2) \quad U^{\leq N} \quad (i = 0))$$

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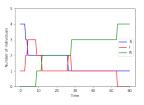
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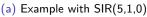
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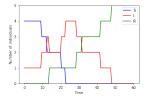
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(b) Counter-example with SIR(5,1,0)

Figure: Example and counter-example on SIR(5,1,0) CTMC with $(\alpha, \beta) = (0.034055, 0.087735)$.

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SIR(5,1,0)	SIR(10,1,0)	SIR(15,1,0)
	1 (,-,-)	311(13,1,0)
6	11	16
27	77	152
(0.034055, 0.087735)	(0.025490, 0.069298)	(0.011499, 0.062111)
(1098, 1377, 1296, 1312, 1466, 3451)	(1002, 1258, 1123, 902, 770, 651, 497, 420, 496, 685, 2196)	(50, 181, 302, 455, 539, 567, 582, 566, 541, 553, 574, 528, 512, 586, 875, 2589)
$P_{>0.25}(i \le 3 \ U^{\le 6} \ i = 0)$	$P_{>0.25} (i \le 5 \ U^{\le 11} \ i = 0)$	$P_{>0.25} (i \le 8 \text{ U}^{\le 16} i$
0.3474444	0.265815	0.327446
	6 27 $(0.034055, 0.087735)$ $(1098, 1377, 1296, 1312, 1466, 3451)$ $P_{>0.25}(i \le 3 \ U^{\le 6} \ i = 0)$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$

Table: True parameters and synthetic data for SIR(5,1,0), SIR(10,1,0), SIR(15,1,0)

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- [1] Christel Baier and Joost-Pieter Katoen. *Principles of model checking*. MIT press, 2008.
- [2] Edmund M Clarke et al. "Model checking and the state explosion problem". In: *LASER Summer School on Software Engineering*. Springer. 2011, pp. 1–30.
- [3] Katalin Csilléry et al. "Approximate Bayesian computation (ABC) in practice". In: *Trends in ecology & evolution* 25.7 (2010), pp. 410–418.
- [4] Conrado Daws. "Symbolic and parametric model checking of discrete-time Markov chains". In:

 International Colloquium on Theoretical Aspects of Computing. Springer. 2004, pp. 280–294.

References II

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- [5] Dynamic Configuration of IPv4 Link-Local Addresses. Accessed: 2021-03-30. URL: https://tools.ietf.org/html/rfc3927.
- [6] Sarah Filippi et al. "On optimality of kernels for approximate Bayesian computation using sequential Monte Carlo". In: Statistical applications in genetics and molecular biology 12.1 (2013), pp. 87–107.
- [7] Ernst Moritz Hahn, Holger Hermanns, and Lijun Zhang. "Probabilistic reachability for parametric Markov models". In: *International Journal on Software Tools for Technology Transfer* 13.1 (2011), pp. 3–19.
- [8] Sebastian Junges et al. "Parameter synthesis for Markov models". In: arXiv preprint arXiv:1903.07993 (2019).

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- [9] Joost-Pieter Katoen. "Model Checking Meets Probability: A Gentle Introduction.". In: Engineering dependable software systems 34 (2013), pp. 177–205.
- [10] Joost-Pieter Katoen. "The probabilistic model checking landscape". In: Proceedings of the 31st Annual ACM/IEEE Symposium on Logic in Computer Science. 2016, pp. 31–45.
- [11] William Ogilvy Kermack and Anderson G McKendrick. "A contribution to the mathematical theory of epidemics". In: *Proceedings of the royal society of london. Series A, Containing papers of a mathematical and physical character* 115.772 (1927), pp. 700–721.
- [12] John FC Kingman. "Markov population processes". In: Journal of Applied Probability (1969), pp. 1–18.

References IV

Bayesian parameter synthesis for Markov population models.

Phung Nhat-Hu

Preliminarie

Contribution

Evaluatio

Conclusion and future work

- [13] Donald E Knuth, KNUTH DE, and YAO AC. "THE COMPLEXITY OF NONUNIFORM RANDOM NUMBER GENERATION.". In: (1976).
- [14] Marta Kwiatkowska, Gethin Norman, and David Parker.
 "Symmetry reduction for probabilistic model checking".
 In: International Conference on Computer Aided Verification. Springer. 2006, pp. 234–248.
- [15] Tim Quatmann et al. "Parameter synthesis for Markov models: Faster than ever". In: *International Symposium on Automated Technology for Verification and Analysis*. Springer. 2016, pp. 50–67.

References V

Bayesian parameter synthesis for Markov population models

Phung Nhat-Hu

Motivatio

Contribution

Framework

Evaluatio

Conclusion and future work

- [16] Daniel Silk, Saran Filippi, and Michael PH Stumpf. "Optimizing threshold-schedules for approximate Bayesian computation sequential Monte Carlo samplers: applications to molecular systems". In: arXiv preprint arXiv:1210.3296 (2012).
- [17] Scott A Sisson, Yanan Fan, and Mark M Tanaka. "Sequential monte carlo without likelihoods". In: Proceedings of the National Academy of Sciences 104.6 (2007), pp. 1760–1765.

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Evaluatio

and future



Figure: Thank you for your attention.