

Model fitting and inference for infectious disease dynamics

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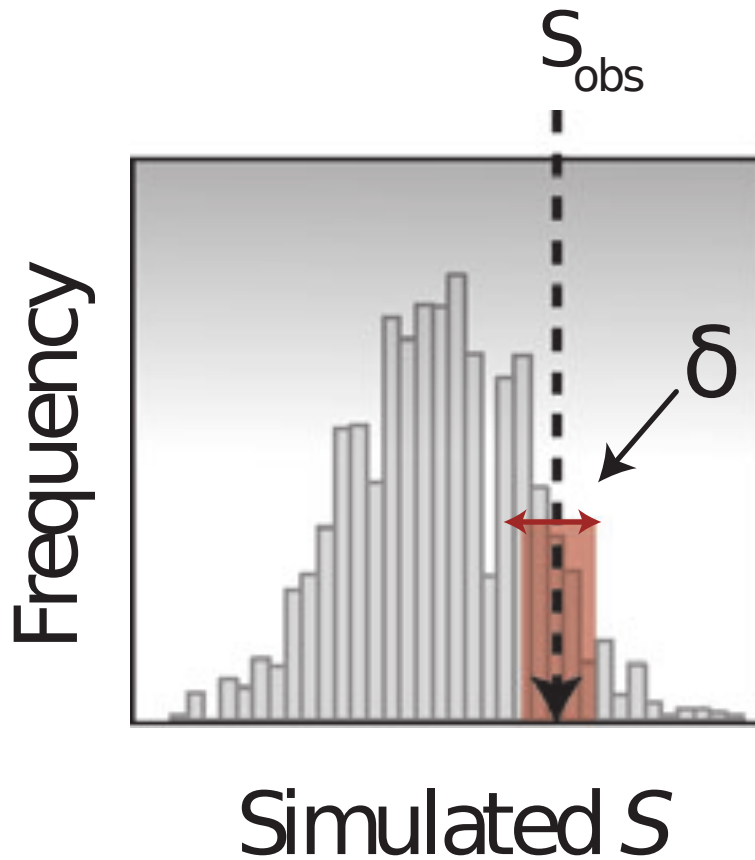
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1 Approximate Bayesian Computation

1.1 Motivation

- we want to evaluate $p(\theta|y)$ (posterior) in a stochastic model
- PMCMC:
 - calculate likelihoods $p(y|\theta, x)$ and marginalise over x
- ABC:
 - *approximate* the posterior using summary statistics S
- Summary statistics
 - Something that is *easy* to calculate and approximates the likelihood
 - e.g., final size, number of peaks, height/timing of peaks
 - idea of *sufficient* summary statistics: to be exact, I need summary statistics that give *exactly* the same result as the likelihood

1.2 Idea of ABC



Frequency that S_{sim} and S_{obs}
closer than δ

1.3 ABC

1. Choose a summary statistic S , positive number ϵ , distance function $d(S_1, S_2)$
2. Calculate summary statistic for data
3. repeat:
 - (a) sample θ from prior $p(\theta)$
 - (b) simulate trajectory
 - (c) calculate S_{sim}

- (d) calculate difference $d(S_{\text{sim}}, S_{\text{data}})$
 - (e) if $d(S_{\text{sim}}, S_{\text{data}}) < \epsilon$ accept, else reject
4. proportion of accepted runs for any θ approximates $p(\theta|y)$

1.4 ABC-MCMC

1. Choose a summary statistic S , positive number ϵ , distance function $d(S_1, S_2)$ and initial θ
2. Calculate summary statistic for data
3. repeat:
 - (a) sample θ^* using transition kernel $q(\theta^*, \theta)$
 - (b) simulate trajectory + observation
 - (c) calculate S_{sim}
 - (d) calculate difference $d(S_{\text{sim}}, S_{\text{data}})$
 - (e) if $d(S_{\text{sim}}, S_{\text{data}}) < \epsilon$ accept with probability

$$\min \left(1, \frac{p(\theta^*)}{p(\theta)} \frac{q(\dots)}{q(\dots)} \right)$$

else reject

4. time spent at any θ approximates $p(\theta|y)$