

Efficient and accurate approximate Bayesian computation

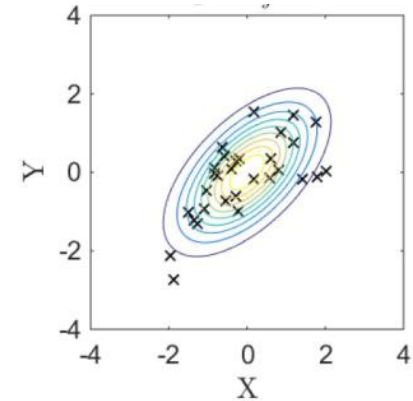
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Marttinen

Outline

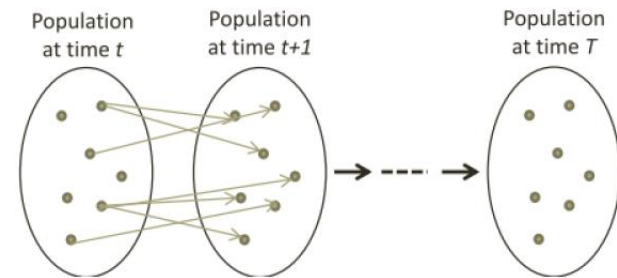
- Simulation-based modeling
- Model-based approximate Bayesian computation (ABC)
- Efficient ABC using Bayesian sequential experimental design
- Results
- Conclusion

Simulation-based modeling

- Statistical inference, the common way
 - Assume some likelihood:
 $p(\text{data}/\text{parameters})$
 - Learn *parameters* that fit the *data*



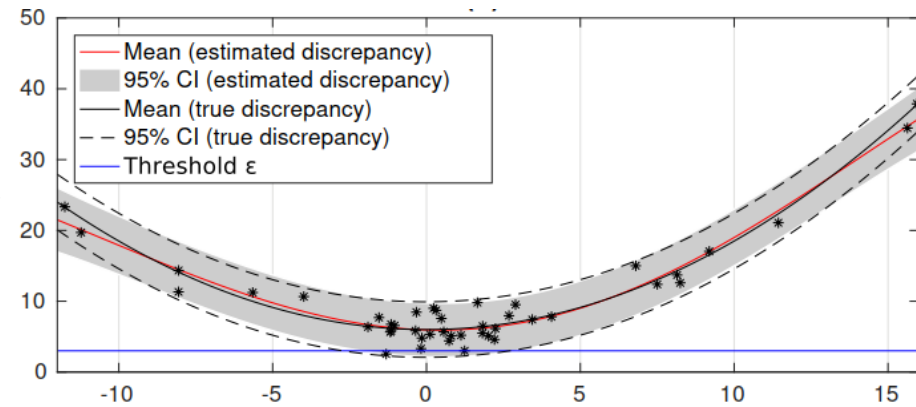
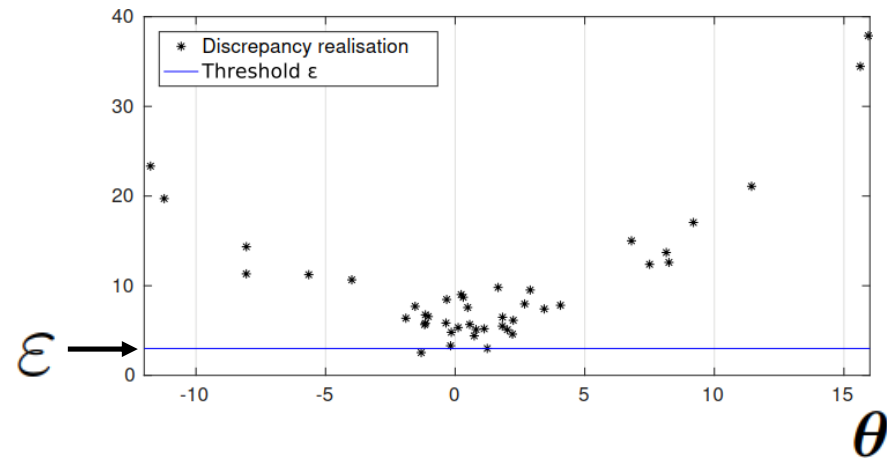
- Sometimes the likelihood can not be computed, but simulating data from the model is possible
 - Example: population genetics



- Applications: economics, material physics, biology, UI design, ...

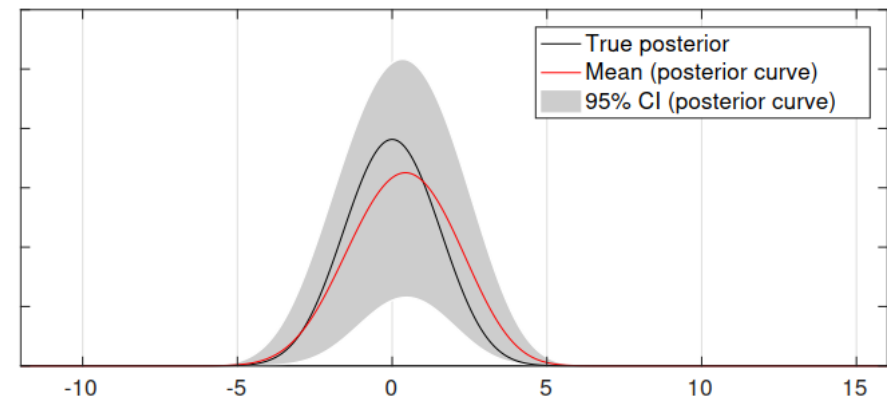
Model-based ABC

$$\Delta_{\theta} = \Delta(\mathbf{x}_{obs}, \mathbf{x}_{\theta})$$



$$\tilde{\pi}_{ABC}(\theta) = \pi(\theta)p_a(\theta)$$

$$p_a(\theta) = \hat{\mathbb{P}}(\Delta_{\theta} \leq \epsilon)$$

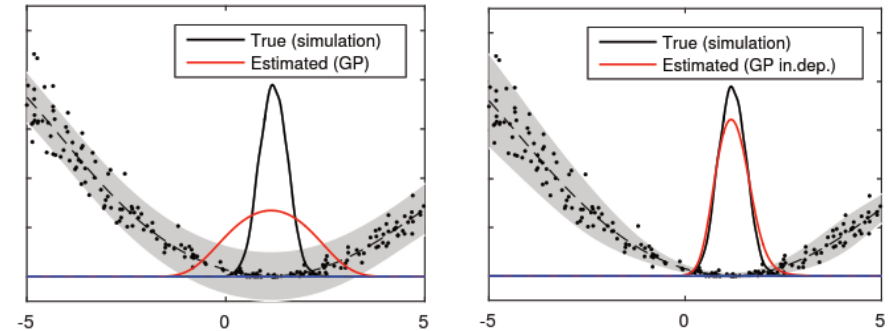


see e.g. Wilkinson (2014); Gutmann and Corander (2016); Järvenpää et al. (2017a)

Efficient and accurate ABC

- How to model the discrepancies

Järvenpää et al. (2017a), submitted.

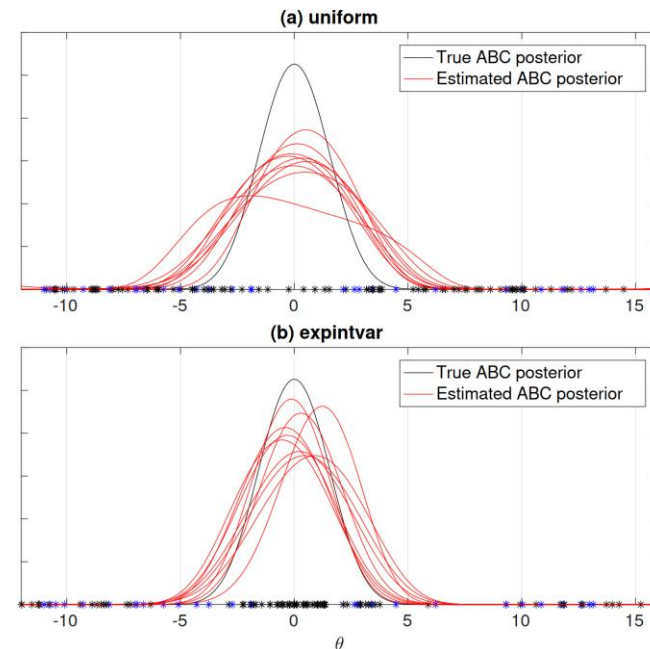


- Where to simulate the model

Järvenpää et al. (2017a), submitted.

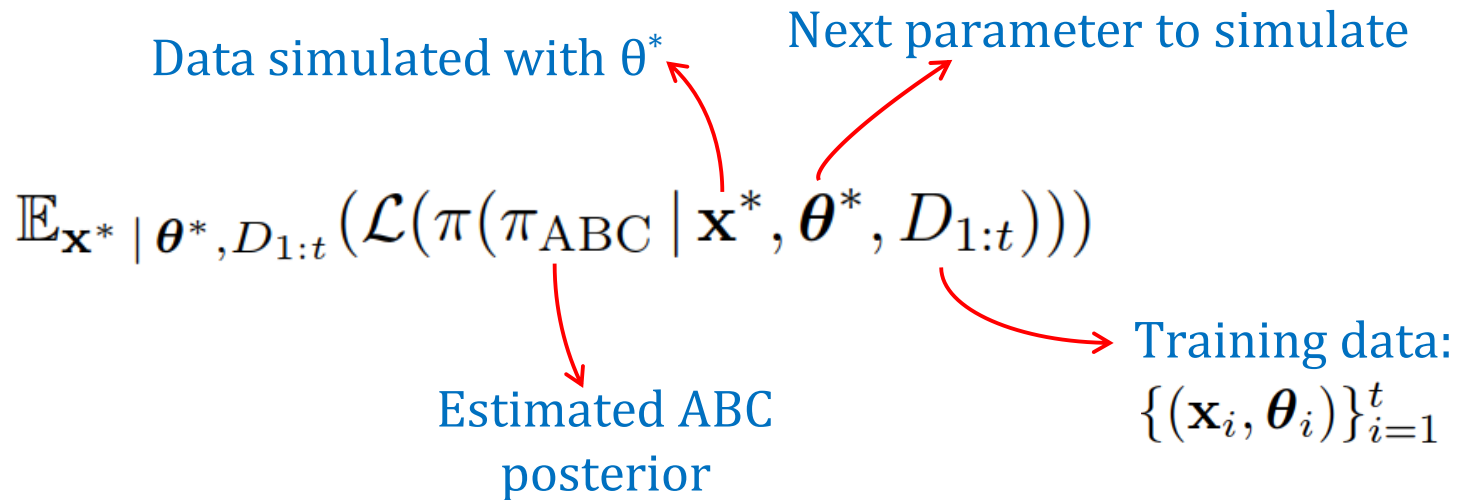
- Implementation: 'ELFI'

Lintusaari, et al. (2017), submitted.



Where to simulate next?

- Bayesian experimental design
- Simulate with θ^* that minimizes the **expected loss**

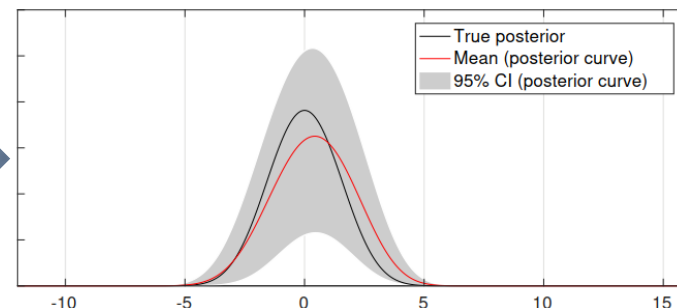
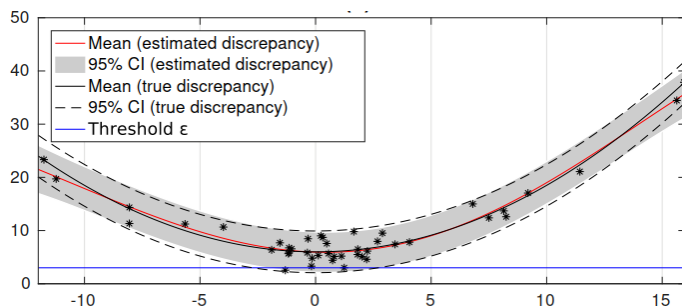


Expected loss

Model for posterior pdf, follows from: $\Delta\theta \sim \mathcal{N}(f(\theta), \sigma_n^2)$
 $f \sim \mathcal{GP}$

$$\mathbb{E}_{\mathbf{x}^* \mid \theta^*, D_{1:t}} (\mathcal{L}(\pi(\pi_{\text{ABC}} \mid \mathbf{x}^*, \theta^*, D_{1:t})))$$

Integrated variance: $\int_{\Theta} \mathbb{V}(\tilde{\pi}_{\text{ABC}}(\theta) \mid D_{1:t}) d\theta$

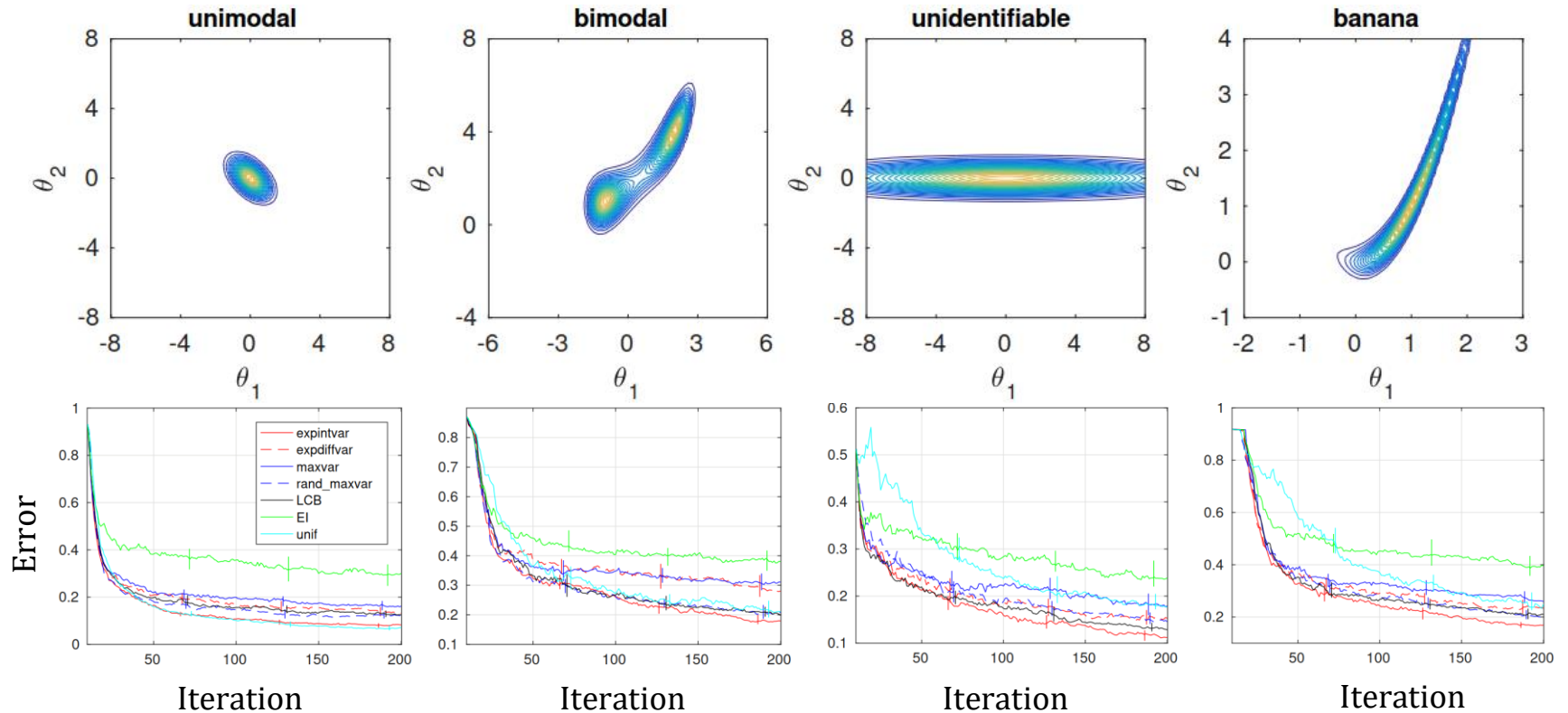


Loss optimization

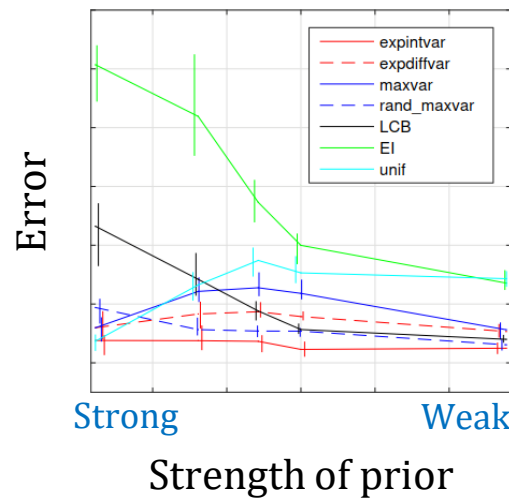
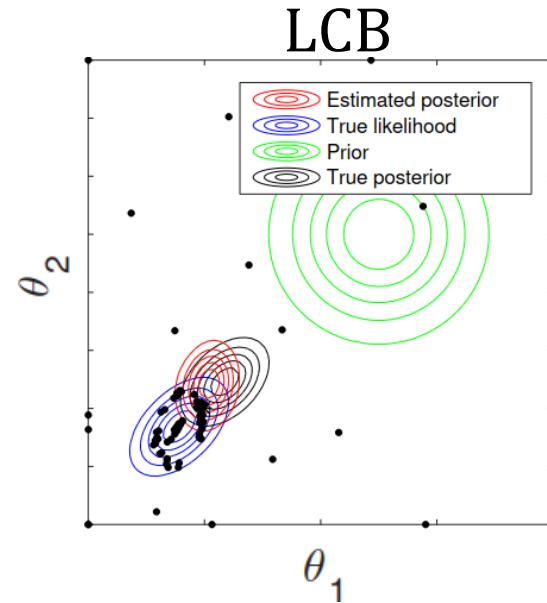
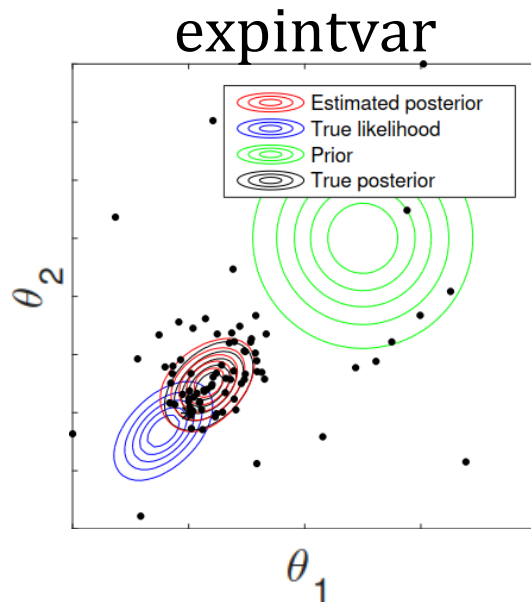
$$L_{1:t}(\boldsymbol{\theta}^*) = 2 \int_{\Theta} \pi^2(\boldsymbol{\theta}) \left[T \left(\frac{\varepsilon - m_{1:t}(\boldsymbol{\theta})}{\sqrt{\sigma_n^2 + v_{1:t}^2(\boldsymbol{\theta})}}, \sqrt{\frac{\sigma_n^2 + v_{1:t}^2(\boldsymbol{\theta}) - \tau_{1:t}^2(\boldsymbol{\theta}, \boldsymbol{\theta}^*)}{\sigma_n^2 + v_{1:t}^2(\boldsymbol{\theta}) + \tau_{1:t}^2(\boldsymbol{\theta}, \boldsymbol{\theta}^*)}} \right) - T \left(\frac{\varepsilon - m_{1:t}(\boldsymbol{\theta})}{\sqrt{\sigma_n^2 + v_{1:t}^2(\boldsymbol{\theta})}}, \frac{\sigma_n}{\sqrt{\sigma_n^2 + 2v_{1:t}^2(\boldsymbol{\theta})}} \right) \right] d\boldsymbol{\theta},$$

- For integration -> **Importance sampling**
- For optimization -> **Gradient descent**

Simulations

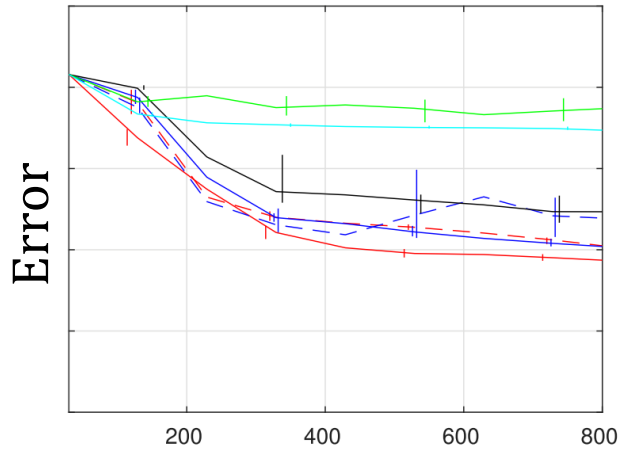


expintvar vs. LCB

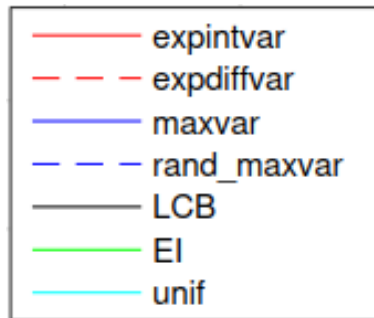
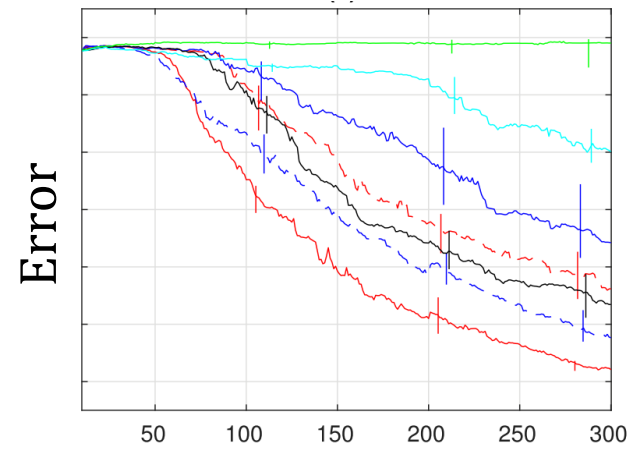


Other examples

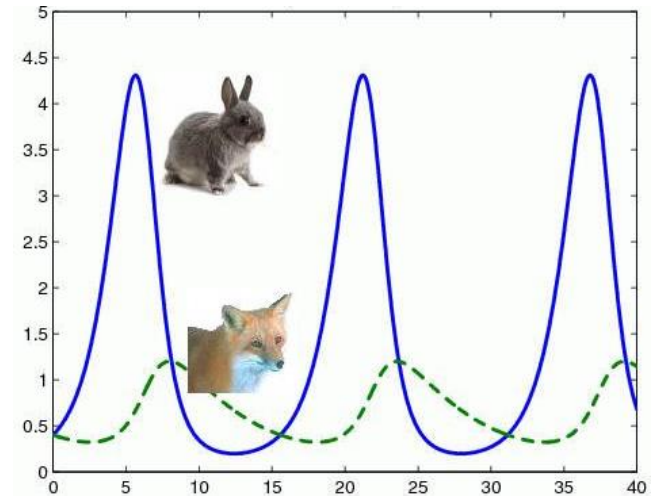
10D Gaussian



Lotka-Volterra

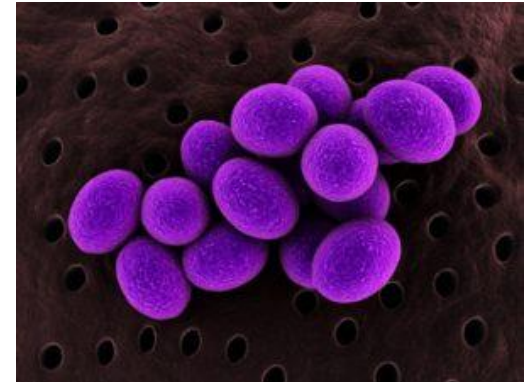


Prey-Predator Cycles



Conclusion

- New principled approach for efficient model-based ABC
- Overall best performance in the experiments, especially when
 - dimension is high
 - there is strong prior information available
- Future:
 - Multi-point proposals for parallelization
 - Models tailored for novel applications



References

Gutmann and Corander (2016). Bayesian optimization for likelihood-free inference of simulator-based statistical models, JMLR.

Järvenpää, Gutmann, Vehtari, Marttinen (2017a). Gaussian process modeling in approximate Bayesian computation to estimate horizontal gene transfer in bacteria , submitted. (arxiv.org/pdf/1610.06462.pdf)

Järvenpää, Gutmann, Pleska, Vehtari, Marttinen (2017b). Efficient acquisition rules for approximate Bayesian computation, submitted. (arxiv.org/pdf/1704.00520.pdf)

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Wilkinson (2014). Accelerating ABC methods using Gaussian processes. In AISTATS.

