

K2-ABC: Approximate Bayesian Computation with Kernel Embeddings

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Approximate Bayesian Computation (ABC)

- Given: prior $p(\boldsymbol{\theta})$, **intractable** likelihood $p(\mathbf{Y}|\boldsymbol{\theta})$, observations \mathbf{Y} .
- Goal: Sample from $p(\boldsymbol{\theta}|\mathbf{Y}) \propto p(\boldsymbol{\theta})p(\mathbf{Y}|\boldsymbol{\theta})$.
- Problem: Cannot evaluate $p(\mathbf{Y}|\boldsymbol{\theta})$. Can sample $\mathbf{X} \sim p(\cdot|\boldsymbol{\theta})$ easily.

Example: a complicated dynamical system for blow fly population

$$N_{t+1} = PN_{t-\tau} \exp\left(-\frac{N_{t-\tau}}{N_0}\right) e_t + N_t \exp(-\delta\epsilon_t)$$

where $e_t \sim \text{Gamma}\left(\frac{1}{\sigma_p^2}, \sigma_p^2\right)$ and $\epsilon_t \sim \text{Gamma}\left(\frac{1}{\sigma_d^2}, \sigma_d^2\right)$.

- $\boldsymbol{\theta} := \{P, N_0, \sigma_d, \sigma_p, \tau, \delta\}$
- Given $\mathbf{Y} = \{N_1, \dots, N_T\}$, want to sample from $p(\boldsymbol{\theta}|\mathbf{Y})$.

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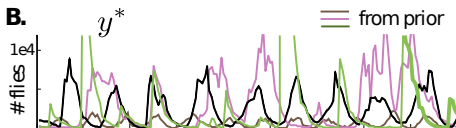
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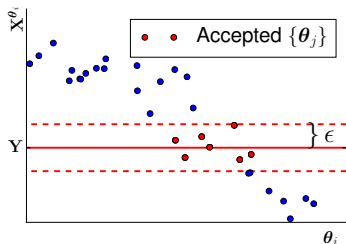
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ABC Likelihood $p_{\epsilon}(\mathbf{Y}|\boldsymbol{\theta})$

- Observe a dataset \mathbf{Y} ,

$$\begin{aligned} p(\mathbf{Y}|\boldsymbol{\theta}) &= \int p(\mathbf{X}|\boldsymbol{\theta})\delta(\mathbf{X} - \mathbf{Y}) d\mathbf{X} \\ &\approx \int p(\mathbf{X}|\boldsymbol{\theta})\kappa_{\epsilon}(\mathbf{X}, \mathbf{Y}) d\mathbf{X} := p_{\epsilon}(\mathbf{Y}|\boldsymbol{\theta}) \\ &\approx \kappa_{\epsilon}(\mathbf{X}^{\boldsymbol{\theta}}, \mathbf{Y}) \text{ where } \mathbf{X}^{\boldsymbol{\theta}} \sim p(\cdot|\boldsymbol{\theta}), \end{aligned}$$

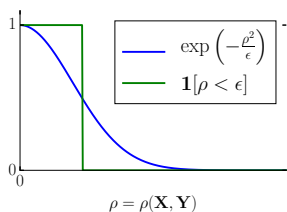


where $\kappa_{\epsilon}(\mathbf{X}, \mathbf{Y})$ defines similarity between \mathbf{X} and \mathbf{Y} .

- Commonly used **rejection ABC** sets

$$\kappa_{\epsilon}(\mathbf{X}, \mathbf{Y}) := \mathbf{1}[\rho(\mathbf{X}, \mathbf{Y}) < \epsilon],$$

- Distance $\rho(\mathbf{X}, \mathbf{Y}) := \|s(\mathbf{X}) - s(\mathbf{Y})\|_2$
- $\mathbf{1}[\cdot] \in \{0, 1\}$: indicator function
- s : function to compute summary statistics



Summary Statistics $s(\cdot)$

- Difficult to choose summary statistics $s(\cdot)$ in

$$\rho(\mathbf{X}, \mathbf{Y}) = \|s(\mathbf{X}) - s(\mathbf{Y})\|_2.$$

- More statistics give high sufficiency.
- But, higher rejection rate.
- Insufficient $s(\cdot)$ will lead to an incorrect posterior.

Contribution:

- Use a kernel distance **MMD** to define ρ . No need to design $s(\cdot)$.

Rejection ABC:

$$\rho(\mathbf{X}, \mathbf{Y}) = \|s(\mathbf{X}) - s(\mathbf{Y})\|_2$$

K2-ABC (proposed):

$$\rho(\mathbf{X}, \mathbf{Y}) = \widehat{\text{MMD}}(\mathbf{X}, \mathbf{Y})$$

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K2-ABC (proposed):

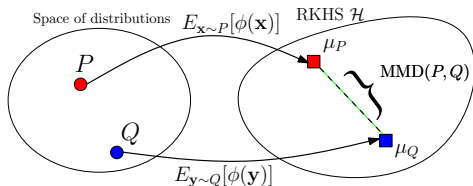
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Maximum Mean Discrepancy (MMD) [Gretton et al., 2006]

$$\begin{aligned} \text{MMD}^2(P, Q) &= \|\mathbb{E}_{\mathbf{x} \sim P}[\phi(\mathbf{x})] - \mathbb{E}_{\mathbf{y} \sim Q}[\phi(\mathbf{y})]\|_{\mathcal{H}}^2 \approx \widehat{\text{MMD}}^2(\mathbf{X}, \mathbf{Y}) \\ &:= \frac{1}{n(n-1)} \sum_{i \neq j} k(\mathbf{x}_i, \mathbf{x}_j) + \frac{1}{n(n-1)} \sum_{i \neq j} k(\mathbf{y}_i, \mathbf{y}_j) - \frac{2}{n^2} \sum_{i, j=1}^n k(\mathbf{x}_i, \mathbf{y}_j) \end{aligned}$$

- If kernel k is characteristic (e.g., Gaussian kernel), $\mu_P = \mathbb{E}_{\mathbf{x} \sim P}[\phi(\mathbf{x})]$ is sufficient for P .

- $k(\mathbf{x}, \mathbf{y}) = \langle \phi(\mathbf{x}), \phi(\mathbf{y}) \rangle_{\mathcal{H}}$
- Intuitively, μ_P contains all moments of P .



K2-ABC (Proposed Method)

- To sample $\{\boldsymbol{\theta}_i\}_{i=1}^M \sim p_\epsilon(\boldsymbol{\theta}|\mathbf{Y})$, do

Output: Approximate posterior $\sum_{i=1}^M \delta_{\boldsymbol{\theta}_i} w_i$

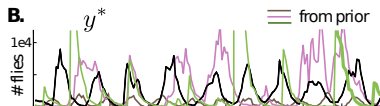
- 1: **for** $i = 1, \dots, M$ **do**
- 2: Sample $\boldsymbol{\theta}_i \sim p(\boldsymbol{\theta})$
- 3: Sample pseudo dataset $\mathbf{X}_i \sim p(\cdot|\boldsymbol{\theta}_i)$
- 4: $\tilde{w}_i = \kappa_\epsilon(\mathbf{X}_i, \mathbf{Y}) = \exp\left(-\frac{\widehat{\text{MMD}}^2(\mathbf{X}_i, \mathbf{Y})}{\epsilon}\right)$
- 5: **end for**
- 6: $w_i = \tilde{w}_i / \sum_{j=1}^M \tilde{w}_j$ for $i = 1, \dots, M$
- 7: **return** $\{\boldsymbol{\theta}_i\}_{i=1}^M$ with weights $\{w_i\}_{i=1}^M$

- Given a function g ,

$$\mathbb{E}_{\boldsymbol{\theta} \sim p(\boldsymbol{\theta}|\mathbf{Y})}[g(\boldsymbol{\theta})] \approx \frac{1}{M} \sum_{i=1}^M w_i g(\boldsymbol{\theta}_i).$$

Blow Fly Population Modelling

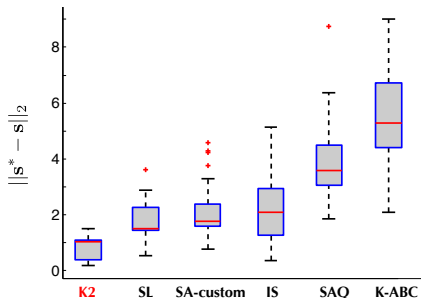
$$N_{t+1} = PN_{t-\tau} \exp\left(-\frac{N_{t-\tau}}{N_0}\right) e_t + N_t \exp(-\delta\epsilon_t)$$



- $e_t \sim \text{Gam}\left(\frac{1}{\sigma_P^2}, \sigma_P^2\right)$ and $\epsilon_t \sim \text{Gam}\left(\frac{1}{\sigma_d^2}, \sigma_d^2\right)$.
- Observe \mathbf{Y} (black solid line).
- Want posterior of $\boldsymbol{\theta} := \{P, N_0, \sigma_d, \sigma_p, \tau, \delta\}$.

-
- Compare 6 ABC methods.
 - 5 other methods use handcrafted 10-dim. summary statistics [Meeds and Welling, 2014].
 - quantiles of the marginal distribution
 - quantiles of first-order differences
 - maximal peaks

Errors on Summary Statistics



- $\tilde{\theta}$:= posterior mean.
- Simulate $\mathbf{X} \sim p(\cdot | \tilde{\theta})$ 100 times.
- $\mathbf{s} = s(\mathbf{X})$ and $\mathbf{s}^* = s(\mathbf{Y})$.

- $\tilde{\theta}$ inferred by K2-ABC gives lowest error on \mathbf{s} .
- Recall that K2-ABC does not use \mathbf{s} , unlike others.

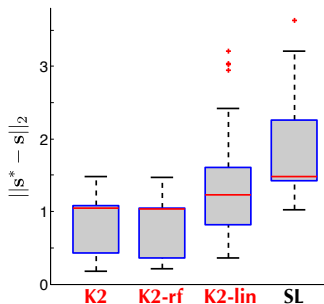
K2-ABC can infer the generative parameters without the need for handcrafted summary statistics.

Linear-Time K2-ABC

- $\widehat{\text{MMD}}^2(\mathbf{X}, \mathbf{Y})$ costs $O(n^2)$ where $n =$ sample size. Expensive.

Solutions:

- 1 Linear-time unbiased estimator. Costs $O(n)$.
- 2 Random Fourier features $\hat{\phi}(\mathbf{x}) \in \mathbb{R}^D$ such that $k(\mathbf{x}, \mathbf{y}) \approx \hat{\phi}(\mathbf{x})^\top \hat{\phi}(\mathbf{y})$. Costs $O(Dn)$. We set $D = 50$.



K2-ABC with random features performs equally well with a much cheaper cost.

Summary

ABC problem:

- Goal: Sample from $p(\boldsymbol{\theta}|\mathbf{Y})$ where the likelihood is intractable.
- Can only sample from the likelihood.

Solution:

- Idea: Keep $\boldsymbol{\theta}$ such that $\mathbf{X} \sim p(\cdot|\boldsymbol{\theta})$ is “similar” to \mathbf{Y} .
- **Contribution**: K2-ABC uses kernel MMD to define the similarity.
 - No need to design summary statistics.
 - Capture all information of $p(\cdot|\boldsymbol{\theta})$.
- Code: <https://github.com/wittawatj/k2abc>



Tue May 10. Poster 6.

Questions?

Thank you

References I

- K2-ABC on Arxiv: <http://arxiv.org/abs/1502.02558>

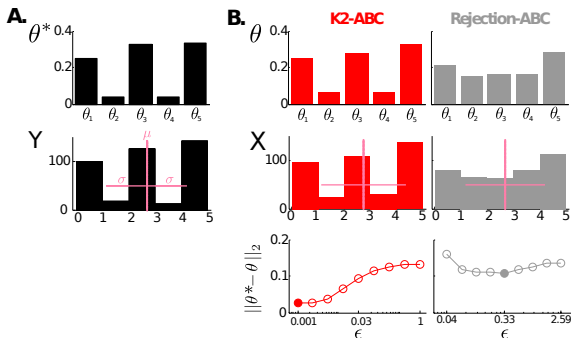
-  Gretton, A., Borgwardt, K. M., Rasch, M., Schölkopf, B., and Smola, A. J. (2006).
A kernel method for the two-sample-problem.
In *Advances in neural information processing systems*, pages 513–520.
-  Meeds, E. and Welling, M. (2014).
GPS-ABC: Gaussian Process Surrogate Approximate Bayesian Computation.
In *UAI*, volume 30, pages 593–601.

Toy Problem: Failure of Insufficient Statistics

$$p(y|\theta) = \sum_{i=1}^5 \theta_i \text{Uniform}(y; [i-1, i])$$

$$\pi(\theta) = \text{Dirichlet}(\theta; \mathbf{1})$$

$$\theta^* = (\text{see figure A})$$



- $s(\mathbf{X}) = (\hat{\mathbb{E}}[x], \hat{\mathbb{V}}[x])^\top$ for Rejection and Soft ABC.
- Insufficient to represent $p(y|\theta)$.

Rejection ABC Algorithm

- **Input:** observed dataset \mathbf{Y} , distance ρ , threshold ϵ
- **Output:** posterior sample $\{\boldsymbol{\theta}_i\}_{i=1}^M$ from approximate posterior $p_\epsilon(\boldsymbol{\theta}|\mathbf{Y}) \propto p(\boldsymbol{\theta})p_\epsilon(\mathbf{Y}|\boldsymbol{\theta})$

```
1: repeat  
2:   Sample  $\boldsymbol{\theta} \sim p(\boldsymbol{\theta})$   
3:   Sample a pseudo dataset  $\mathbf{X} \sim p(\cdot|\boldsymbol{\theta})$   
4:   if  $\rho(\mathbf{X}, \mathbf{Y}) < \epsilon$  then  
5:     Keep  $\boldsymbol{\theta}$   
6:   end if  
7: until we have  $M$  points
```

- **Notation:** \mathbf{Y} = observed set. \mathbf{X} = pseudo (generated) dataset.

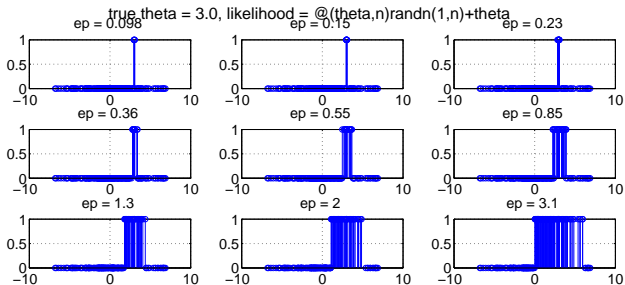
Rejection ABC Example

$$p(y|\theta) = \mathcal{N}(y; \theta, 1)$$

$$p(\theta) = \mathcal{N}(\theta, 0, 8)$$

$$\theta^* = 3.0$$

$$\rho(\mathbf{X}, \mathbf{Y}) = \left| \hat{\mathbb{E}}_{\mathbf{X}}[x] - \hat{\mathbb{E}}_{\mathbf{Y}}[y] \right|$$



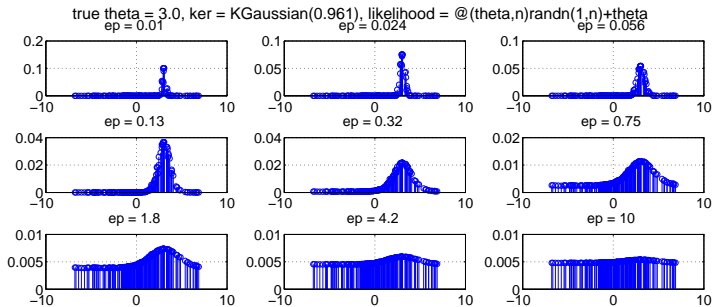
- Low $\epsilon \Rightarrow$ sample closely follows true posterior. High rejection rate.
- High $\epsilon \Rightarrow$ get θ sample from prior.

1D Gaussian Example with K2-ABC

$$p(y|\theta) = \mathcal{N}(y; \theta, 1)$$

$$\pi(\theta) = \mathcal{N}(\theta, 0, 8)$$

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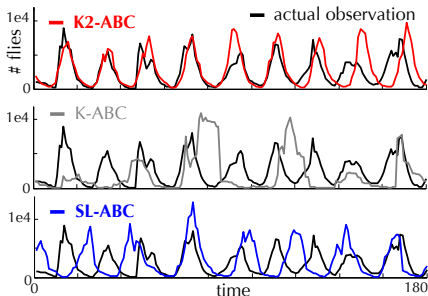
- High $\epsilon \Rightarrow$ get θ sample from prior
- Low $\epsilon \Rightarrow$ sample closely follows true posterior.

Simulated Trajectories

Number of blow flies over time

$$N_{t+1} = PN_{t-\tau} \exp\left(-\frac{N_{t-\tau}}{N_0}\right) e_t + N_t \exp(-\delta\epsilon_t)$$

- $e_t \sim \text{Gam}\left(\frac{1}{\sigma_P^2}, \sigma_P^2\right)$ and $\epsilon_t \sim \text{Gam}\left(\frac{1}{\sigma_d^2}, \sigma_d^2\right)$.
- Want $\theta := \{P, N_0, \sigma_d, \sigma_p, \tau, \delta\}$.



- ← Simulated trajectories with inferred posterior mean of θ
- Other methods use handcrafted 10-dim. summary statistics [Meeds and Welling, 2014].
 - quantiles of the marginal distribution
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