

# How approximate is ABC?

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# ABC Rejection Sampling Algorithm

### Inputs

- N > 0 integer
- $\pi(\theta|y_{obs}) \propto p(y_{obs}|\theta)\pi(\theta)$  target posterior density
- $g(\theta)$  proposal density
- $K_h(u)$  kernel function where h > 0 scale parameter
- s = S(y) summary statistic

### Sampling

For i=1:N

- 1. Generate  $\theta^{(i)} \sim g(\theta)$
- 2. Generate  $y \sim p(y|\hat{\theta}^{(i)})$
- 3. Compute summary statistic s = S(y)
- 4. Accept  $\theta^{(i)}$  with probability  $\frac{K_h(\|s-s_{obs}\|)\pi(\theta^{(i)})}{K*g(\theta^{(i)})}$ , where  $K \geq K_h(0) \max_{\theta} \frac{\pi(\theta)}{\sigma(\theta)}$ , otherwise go to 1.

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- ► Acceptance rate and tolerance
- Summary statistics
- ► Kernel
- ► Proposal

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### Dataset REACH

The Rotterdam Early Arthritis Cohort dataset contains **681 observations**.

The study's aim is to point out which of the following **12 factors** are associated with the development of rheumatoid arthritis:

### Model specification:

$$egin{aligned} & Y_i | \mathsf{X}_i, eta \sim \mathit{Bern}(\phi(\mathsf{X}_i^Teta)), \ \mathsf{i}{=}1,\dots,\mathsf{n} \\ & eta_\gamma | \gamma \sim (1-\gamma_j)\delta_{\{0\}} + \gamma_j \mathsf{N}(0,\sigma_{\beta_j}^2) \\ & \gamma_j | heta_j \sim \mathit{Bern}( heta_j), \ \mathsf{j}{=}1,\dots,\mathsf{K} \\ & heta_i \sim \mathcal{U}(0,1) \end{aligned}$$

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# Acceptance Rate and Tolerance

- Simulate 200 000 observations
- Collect differences in the vector  $d = ||y y_{obs}||$
- ullet Set tolerance=1% percentile of d
- Acceptance rate  $\simeq 1\%$

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# Results

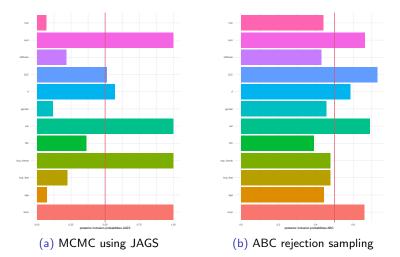


Figure: Comparison of covariates

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### Dataset GENUS

The dataset consists in **341 observations** divided in **12 genus** and, for each observation, we have:

#### BoW SVL BrW

Feature variable: 
$$R = log \left( \frac{BrW/BoW}{1 - BrW/BoW} \right)$$

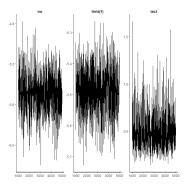
#### Hierarchical model:

$$R_j = R_1, \dots, R_{n_j} | \theta_j, \sigma^2 \sim N(\theta_j, \sigma^2), j=1, \dots, M=12 \text{ (genus)}$$
 $\theta_1, \dots, \theta_M | \mu, \tau^2 \sim N(\mu, \tau^2)$ 
 $(\mu, \tau^2) \sim N(m_0, s_0^2) * InvGamma(\alpha_0, \beta_0)$ 
 $\sigma^2 \sim InvGamma(a_0, b_0)$ 

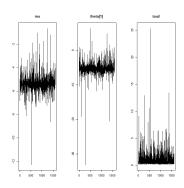
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- ► Acceptance rate and tolerance
- **▶** Summary statistics
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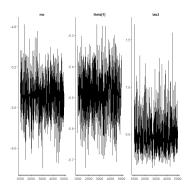


(a) MCMC using STAN

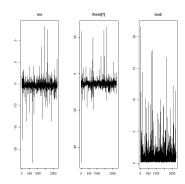


(b) ABC Rejection Sampling using the **mean** as summary statistic

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(c) MCMC using STAN



(d) ABC Rejection Sampling using the **median** as summary statistic

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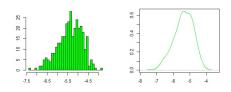


Figure: True value of R

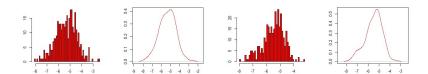


Figure: Estimated values of R: mean and median respectively

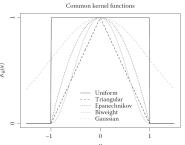
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- ► Acceptance rate and tolerance
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- ► Proposal

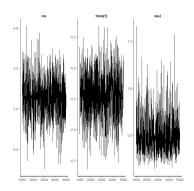
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# Kernel

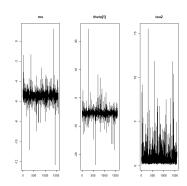
Kernel	K(u)	-	1
Uniform	$\frac{1}{2}\mathbb{1}_{ u \leq 1}$		
Triangular	$(1- u )\mathbb{1}_{ u \leq 1}$	$\langle V_h(u) \rangle$	
Epanechnikov	$\frac{3}{4}(1-u^2)\mathbb{1}_{ u \leq 1}$	$K_h$	
Biweight	$\frac{15}{16}(1-u^2)^3\mathbb{1}_{ u \leq 1}$		
Gaussian	$\frac{1}{\sqrt{(2\pi)}}e^{-\frac{1}{2}u^2}$	c	- <u>-</u>



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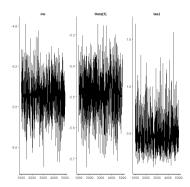


(a) MCMC using STAN

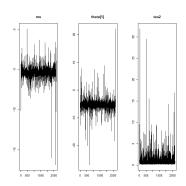


(b) ABC Rejection Sampling using the **mean** as summary statistic **and uniform kernel** 

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(c) MCMC using STAN



(d) ABC Rejection Sampling using the **mean** as summary statistic **and gaussian kernel** 

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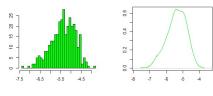


Figure: True value of R

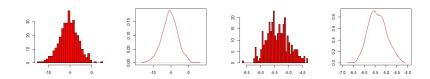


Figure: Estimated values of R: mean with **uniform** and **gaussian kernel** respectively

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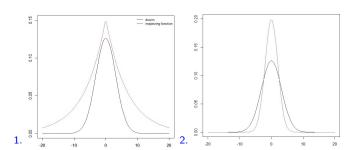
# Proposal

### 1. Majorizing function

It can be obtained by simply rescaling a proposal density

#### 2. Importance weights density

The proposal density is large in parameter ranges that are preferred by the data.



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### FURTHER DEVELOPMENTS

# ABC MCMC

#### REFERENCES

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