

# Bayesian Analysis of Massive Datasets Via Particle Filters

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# Bayesian Analysis

- Use Bayes' theorem to learn about model parameters from data

$$f(\theta \mid \text{data}) \propto f(\text{data} \mid \theta) f(\theta)$$

- Examples:
  - Clustered data: hospitals, schools
  - Spatial models: public health
  - Support vector machines
  - Model based clustering

# Metropolis-Hastings algorithm

1. Initialize  $\theta_1$
2. For  $i$  in  $2, \dots, M$ 
  - a. Draw a proposal  $\theta'$  from  $q(\theta' | \theta_{i-1})$
  - b. Compute the acceptance probability

$$\alpha(\theta', \theta_{i-1}) = \min\left(1, \frac{f(\theta' | \mathbf{x})q(\theta_{i-1} | \theta')}{f(\theta_{i-1} | \mathbf{x})q(\theta' | \theta_{i-1})}\right)$$

- c. Set  $\theta_i = \theta'$  with probability  $\alpha$

Otherwise  $\theta_i = \theta_{i-1}$

# Important ideas

- Metropolis makes Bayesian analysis practical
- Metropolis often requires an enormous number of laps through the dataset
- Given a  $\theta$  drawn from  $f(\theta | \mathbf{x})$ , the Metropolis algorithm produces a new draw having the same distribution
- Using particle filtering we reverse the inner and outer for-loops of Metropolis

# Importance sampling

- Target distribution is  $f(\theta | \mathbf{x})$
- Sampling distribution is  $g(\theta)$

$$\int \theta f(\theta | x_1, \dots, x_N) d\theta = \int \theta \frac{f(\theta | \mathbf{x})}{g(\theta)} g(\theta) d\theta$$

$$= \lim_{M \rightarrow \infty} \frac{\sum_{i=1}^M w_i \theta_i}{\sum_{i=1}^M w_i}$$

- $\theta_i$  has density  $g(\theta)$  and
- $w_i = f(\theta_i | \mathbf{x})/g(\theta_i)$

# Important ideas

- We cannot sample from  $f(\theta | \mathbf{x})$  directly because the model is complex and  $\mathbf{x}$  is massive
- Importance sampling allows us to sample from difficult to sample distributions
- For efficiency,  $g(\theta)$  and  $f(\theta | \mathbf{x})$  should be similar

# Importance sampling for massive datasets

- Set the sampling distribution as

$$g(\theta) = f(\theta \mid x_1, \dots, x_n)$$

where  $n \ll N$

- The importance weights greatly simplify

$$w_i = \frac{f(\theta_i \mid x_1, \dots, x_N)}{f(\theta_i \mid x_1, \dots, x_n)} \propto \prod_{j=n+1}^N f(x_j \mid \theta_i)$$

- Use Metropolis to sample from  $g(\theta)$  and reweight the draws to look like a sample from  $f(\theta \mid \mathbf{x})$

# The algorithm

- Load as much data into memory as possible to form  $D_1$
- Draw  $M$  times from  $f(\theta|D_1)$  via Metropolis
- Purge  $D_1$  from memory
- Set  $w_i = 0, i = 1, \dots, M$

For  $j = n+1, \dots, N$

{

    for  $i = 1, \dots, M$

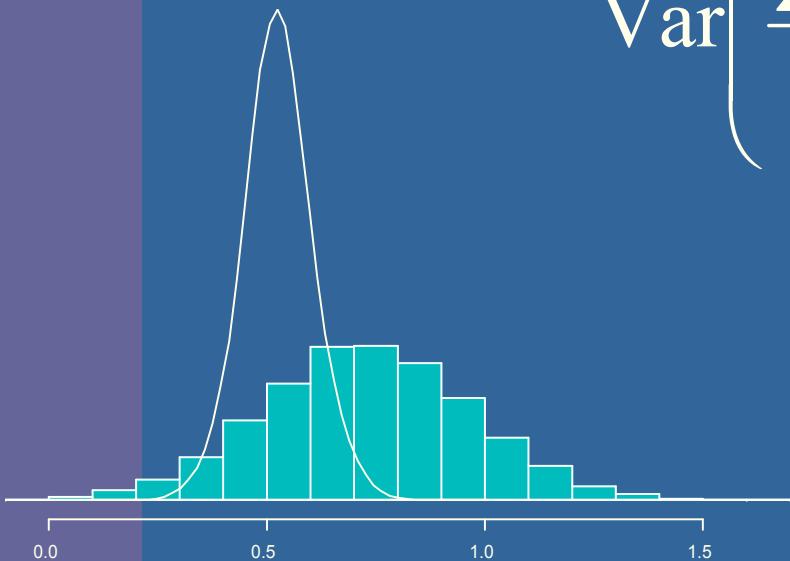
$$\log w_i = \log w_i + \log f(\mathbf{x}_j | \theta_i)$$

}

Transform and rescale to obtain the weights

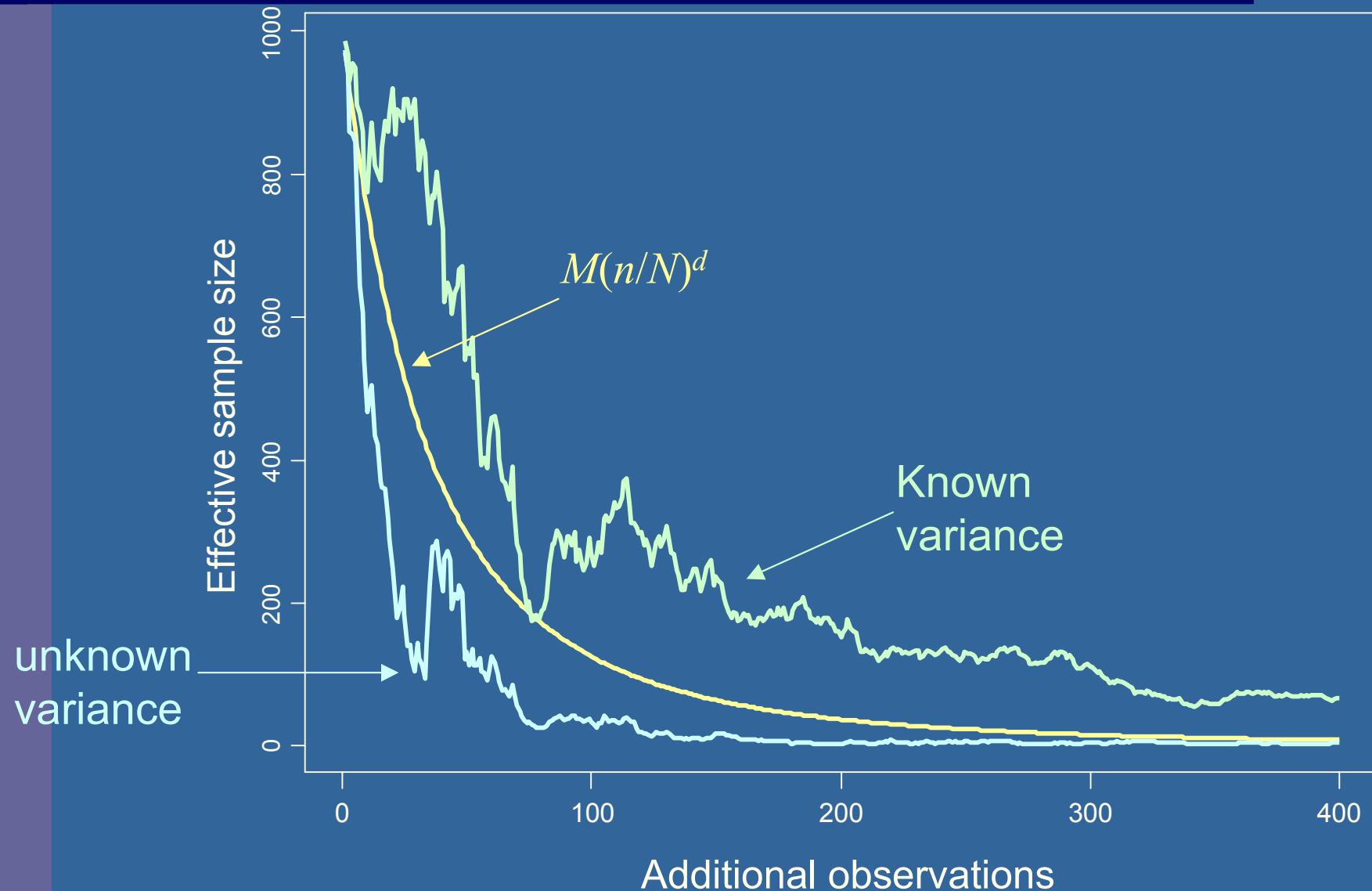
# Effective sample size

$$\text{Var}\left(\frac{\sum_{i=1}^M w_i \theta_i}{\sum_{i=1}^M w_i}\right) = \text{Var}\left(\frac{1}{ESS} \sum_{i=1}^{ESS} \theta_i\right)$$



$$ESS = \frac{\left(\sum_{i=1}^M w_i\right)^2}{\sum_{i=1}^M w_i^2} \approx M \left(\frac{n}{N}\right)^d$$

# ESS deterioration

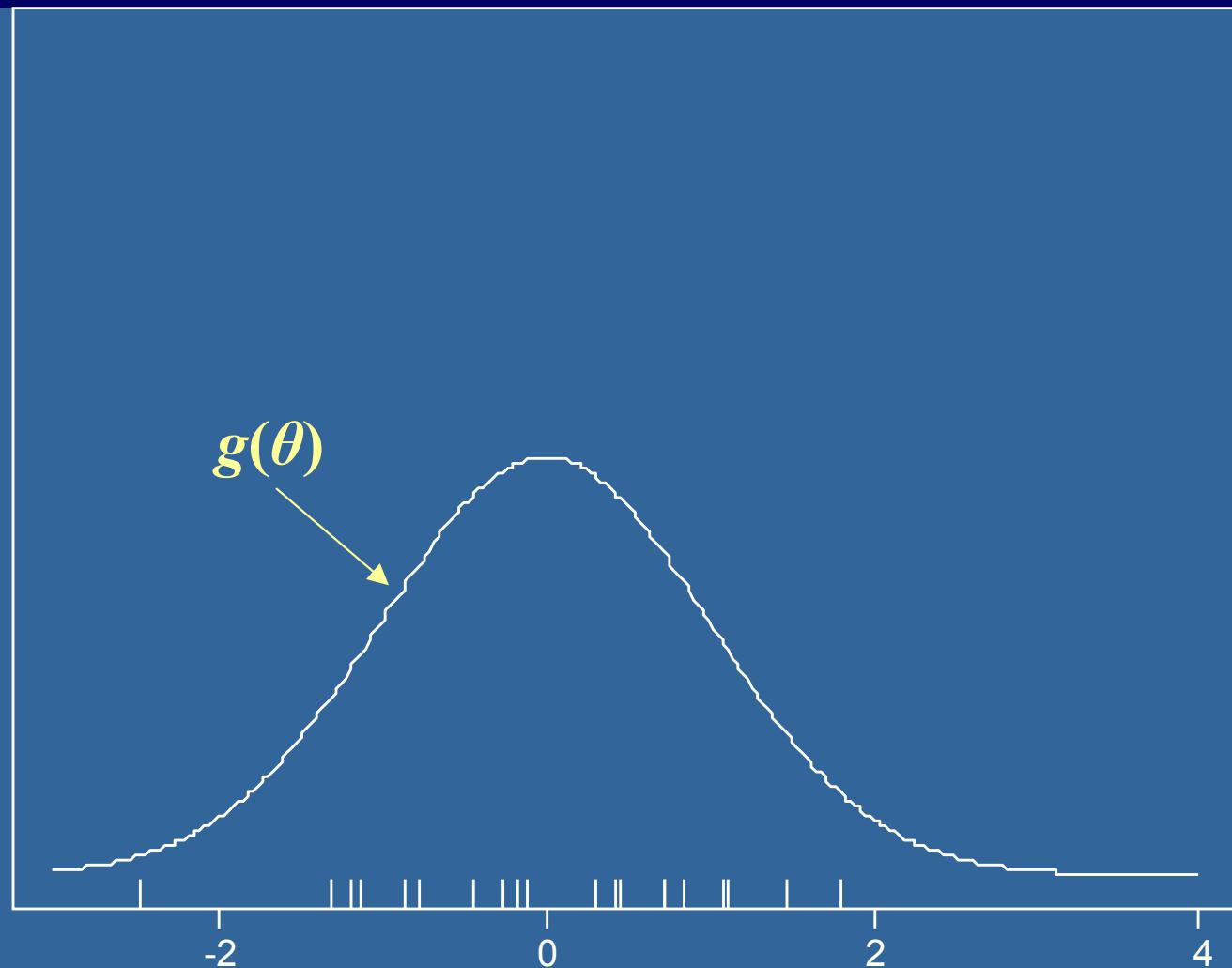


# Gilks and Berzuini rejuvenation

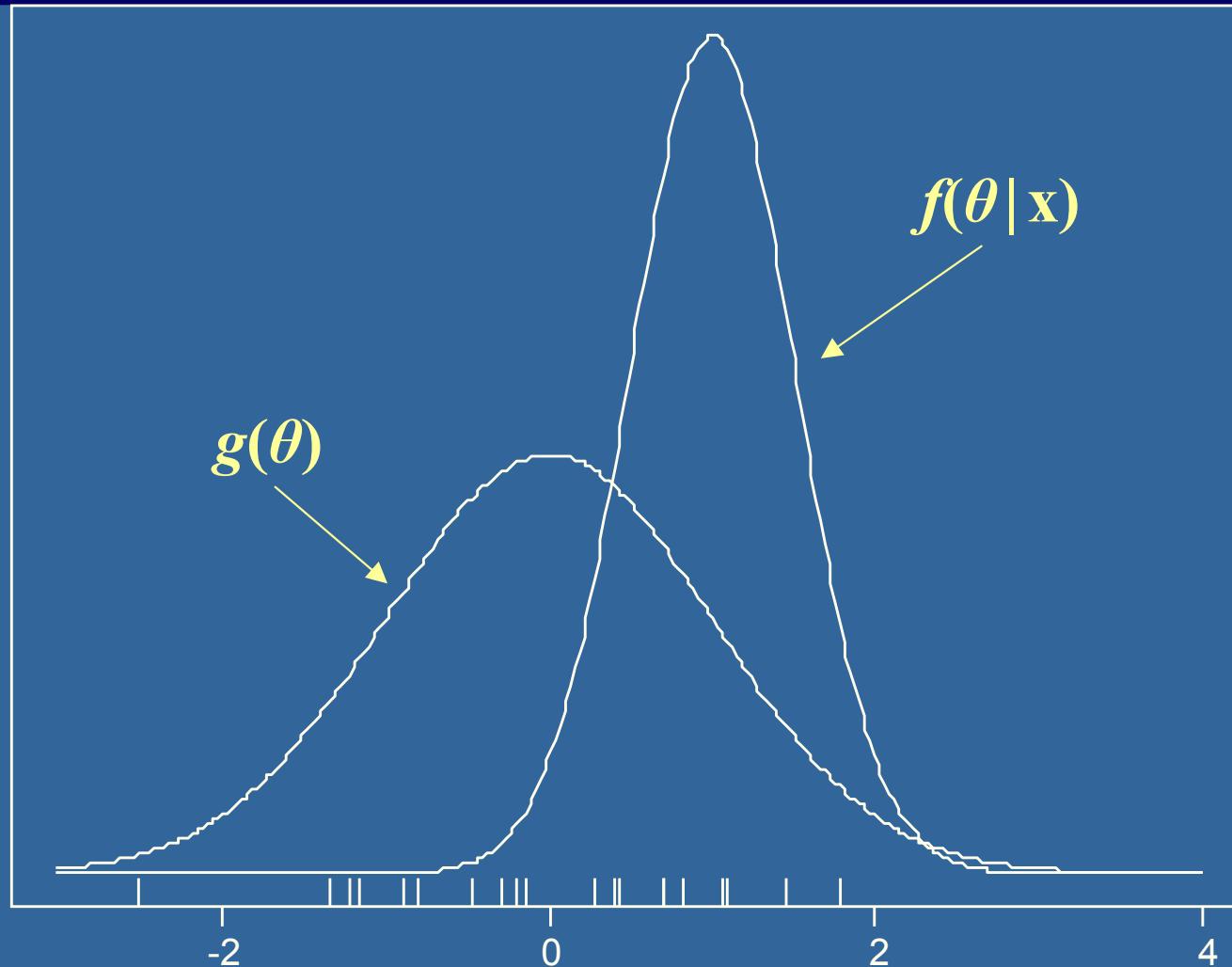
$\theta_1, \dots, \theta_M$  are *particles* and the weights *filter* the  $\theta_i$  with little posterior mass

- Get initial sample from  $f(\theta | x_1, \dots, x_n)$
- While ESS is large enough incorporate new observations using importance reweighting
- Sample with replacement from  $\theta_1, \dots, \theta_M$  with probability proportional to  $w_i$
- Rejuvenate: For each  $\theta_i$  do a single Metropolis step

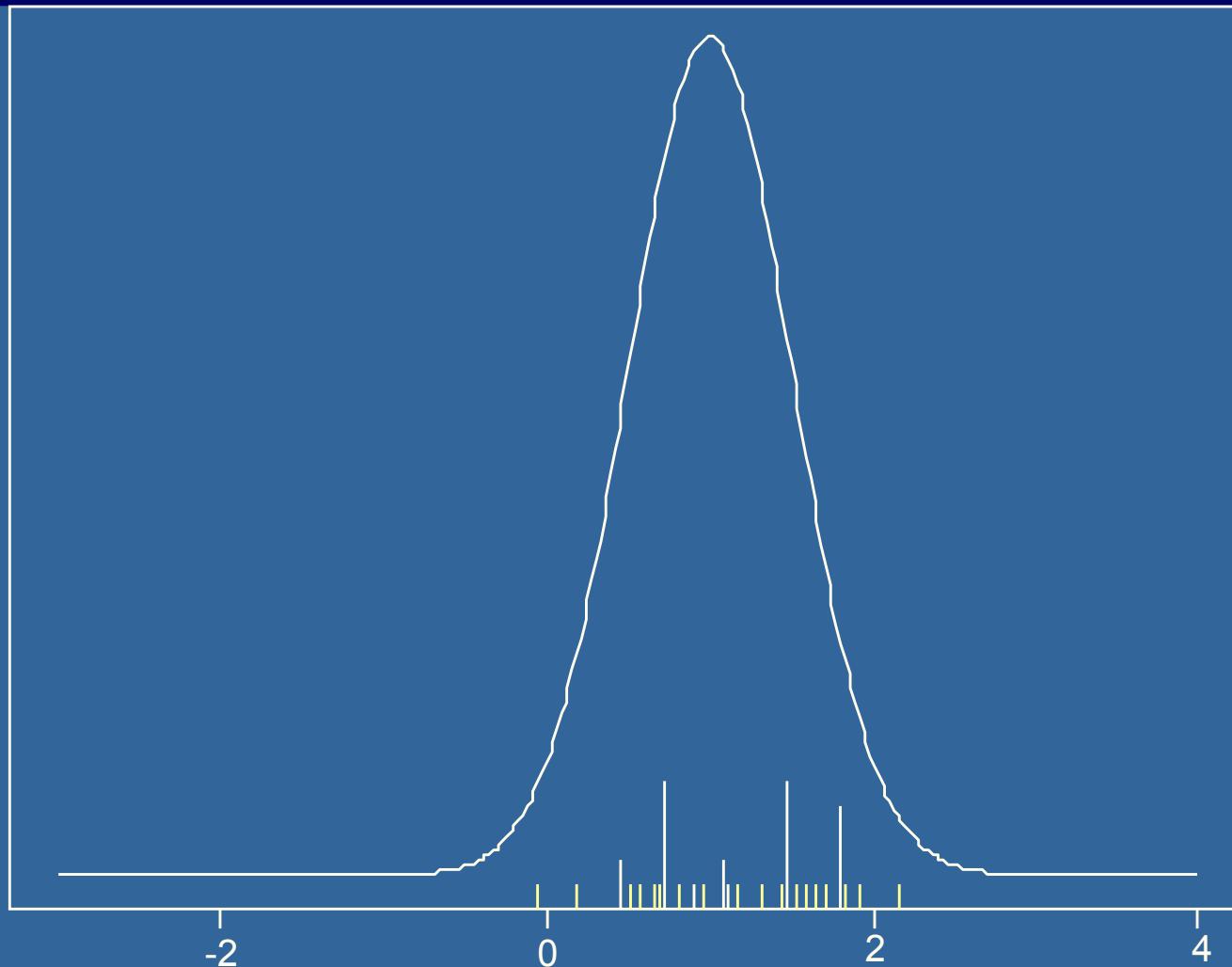
# Sample from $g(\theta)$



# Reweighting, resampling to get $f(\theta | x)$



# Rejuvenate



# Frequency of rejuvenation

Observations absorbed at refresh  $k$

$$= n \left( \frac{1}{p} \right)^{\frac{k}{d}}$$

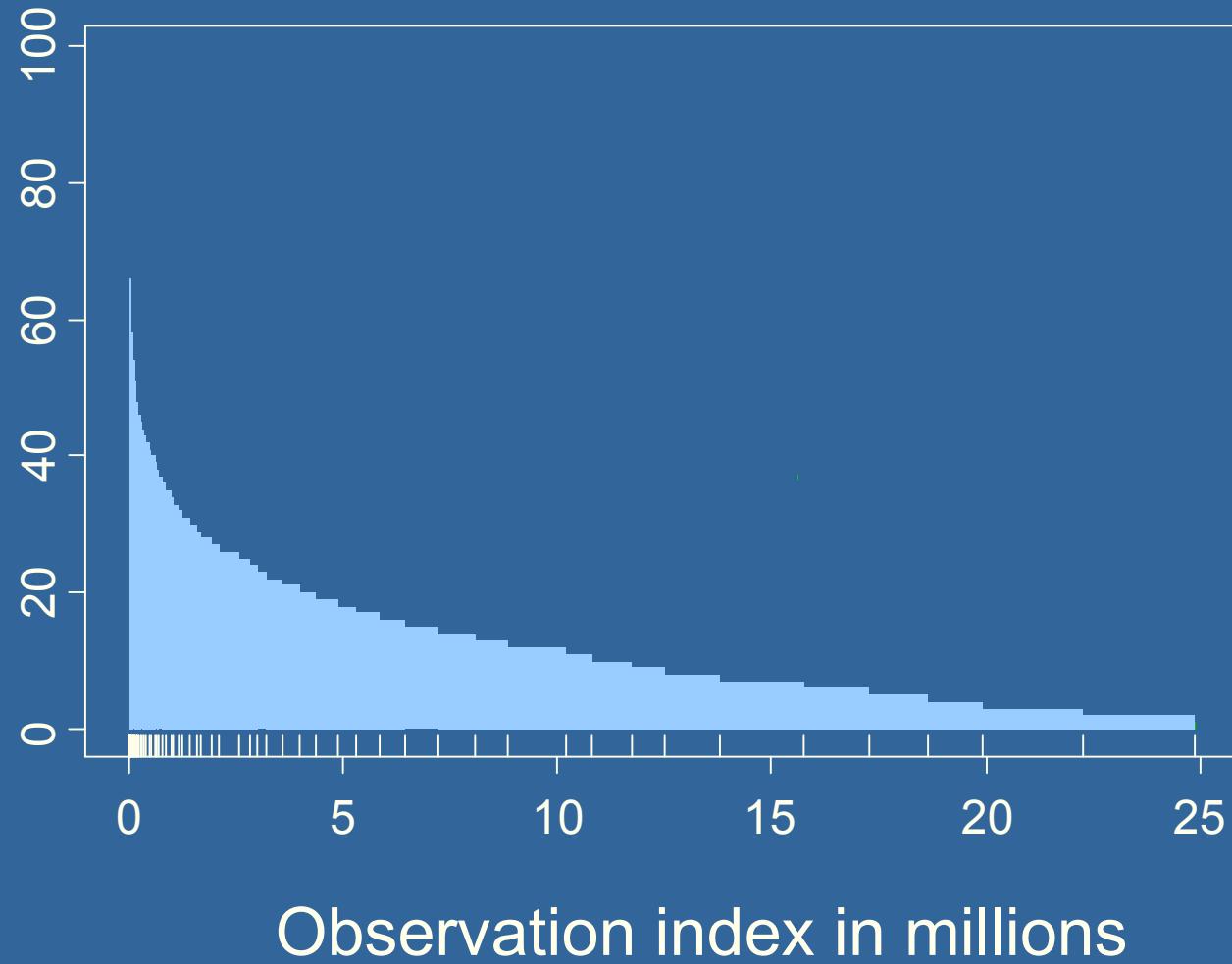
$$= \text{Initial dataset size} \times \left( \frac{1}{\text{percent decrease in ESS}} \right)^{\frac{k}{d}}$$

# Example: Mixture of transition models

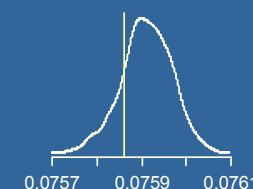
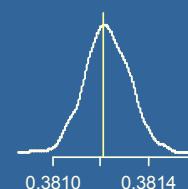
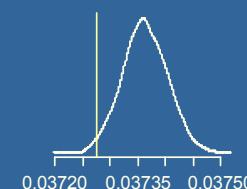
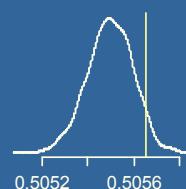
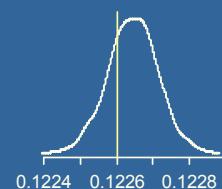
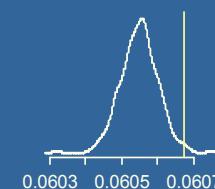
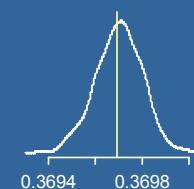
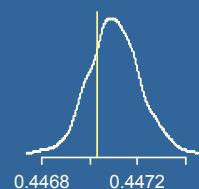
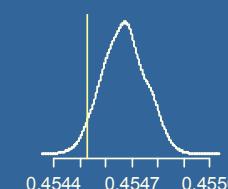
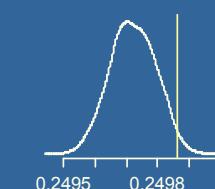
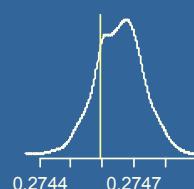
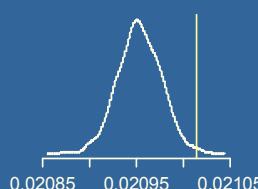
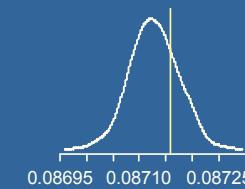
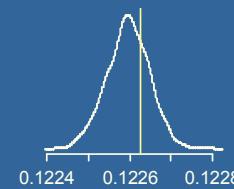
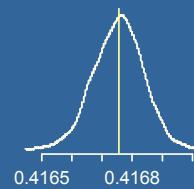
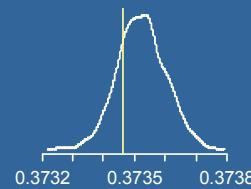
- Set of sequences of about 5 to 20 states visited by each observation
- Each sequence was generated by one of two first order probability transition matrices
- We do not know the transition probabilities nor the cluster assignments
- Properties
  - 25 million observations
  - 1 Gb of data
  - allowed only 1,000 sequences in memory

# Number of accesses

Number of Access



# Cluster 1 transition matrix



# Conclusions

- Requires one good Metropolis-Hastings run up front with a small dataset
- Greatly reduces data access requirements
- Number of data accesses does not depend on  $M$
- Chopin (2002) Biometrika article offers a similar strategy with interesting measures of sample quality

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