

MCMC

Numerical methods for Bayes



SPOILER

HANG THE DJ

ALERT

NETFLIX

BLACK
MIRROR



1000

SIMULATIONS
COMPLETED

998

REBELLIONS
DETECTED

99.8%

MATCH



What are the chances that a solitaire laid out with 52 cards will come out successfully?



Stanislaw Ulam, 1946, LANL

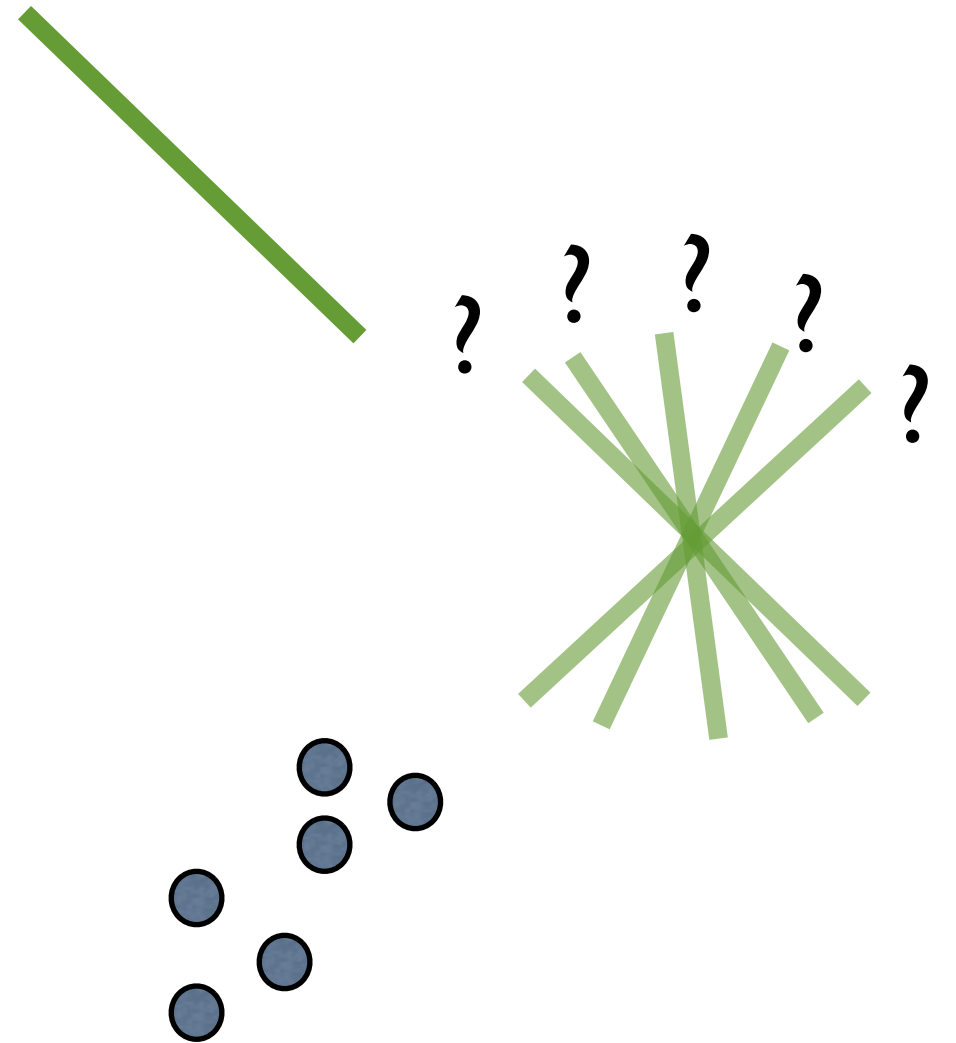
Goal: Find the laws that govern the functioning and the interactions among nature's living organisms.

How? Through a model, $M(\theta)$

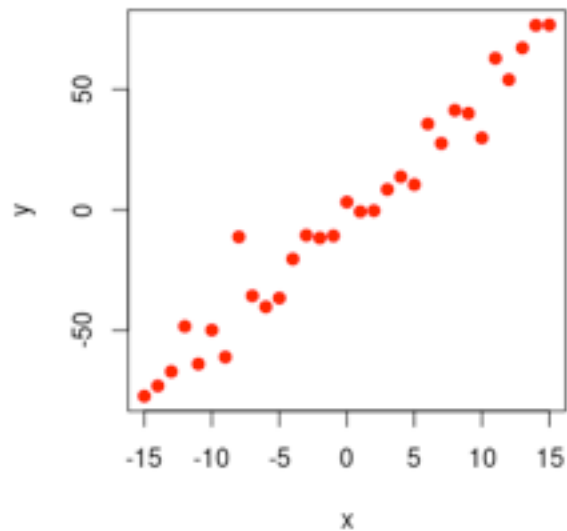
What do we know? $P(\theta)$

What do we have? Some data (Y)

What do we want to know? $P(\theta|Y) = \frac{P(Y|\theta) P(\theta)}{P(Y)}$



Rejection sampling



Model:

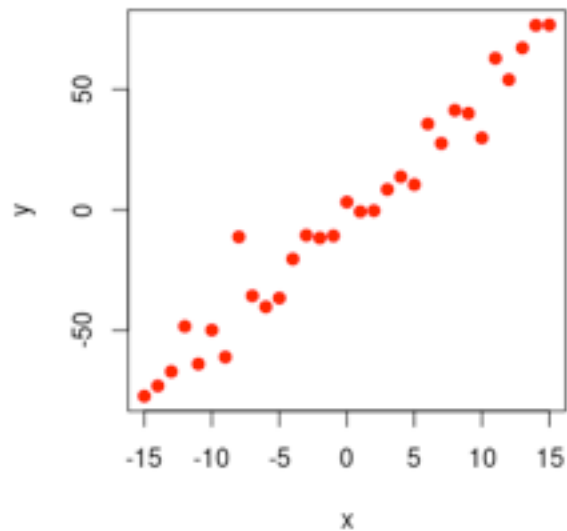
$$y = b + a * x$$

Priors:

$$P(a) \sim U[a_{\min}, a_{\max}]$$

$$P(b) \sim U[b_{\min}, b_{\max}]$$

Rejection sampling



Model:

$$y = b + a \cdot x + \epsilon$$

$$y \sim N(b + a \cdot x, sd)$$

$N(0, sd)$



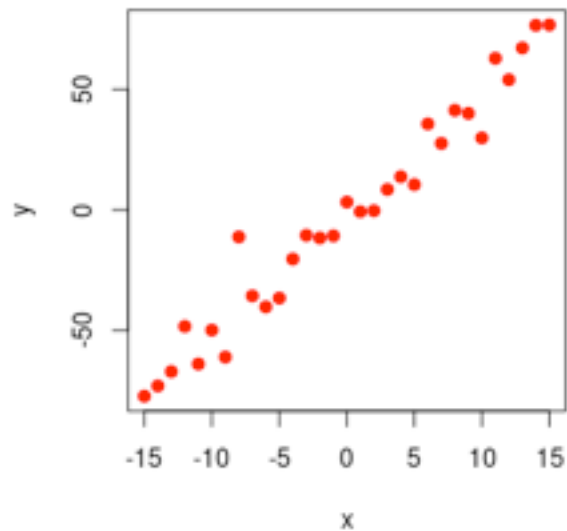
Priors:

$$P(a) \sim U[a_{\min}, a_{\max}]$$

$$P(b) \sim U[b_{\min}, b_{\max}]$$

$$P(sd) \sim U[sd_{\min}, sd_{\max}]$$

Rejection sampling



Model:

$$y \sim N(a \cdot x, 10)$$

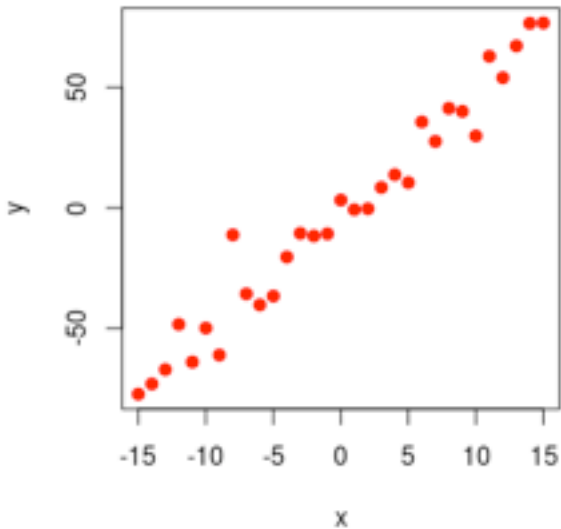
Priors:

$$P(a) \sim U[a_{\min}, a_{\max}]$$

$$b = 0$$

$$sd = 10$$

Rejection sampling



1. draw random values from prior

Model:

$$y \sim N(a \cdot x, 10)$$

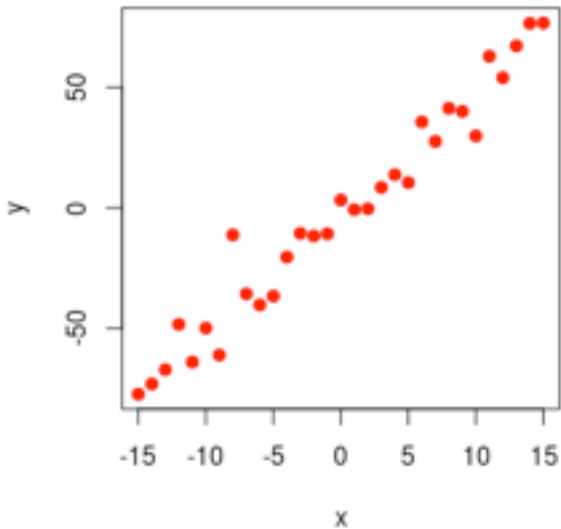
Priors:

$$P(a) \sim U[a_{\min}, a_{\max}] \longrightarrow a_1$$

$$b = 0$$

$$sd = 10$$

Rejection sampling



Model:

$$y \sim N(a*x, 10)$$

Priors:

$$P(a) \sim U[a_{\min}, a_{\max}]$$

$$b = 0$$

$$sd = 10$$

1. draw random values from priors

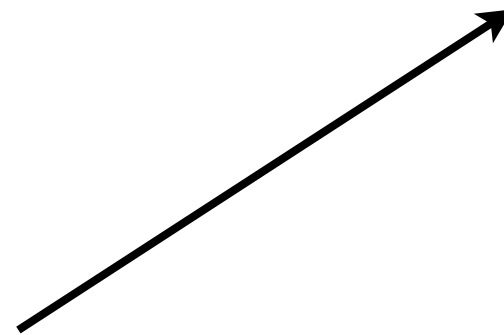
2. calculate likelihood

i.e. the probability of observing data y given the model and drawn parameter values:

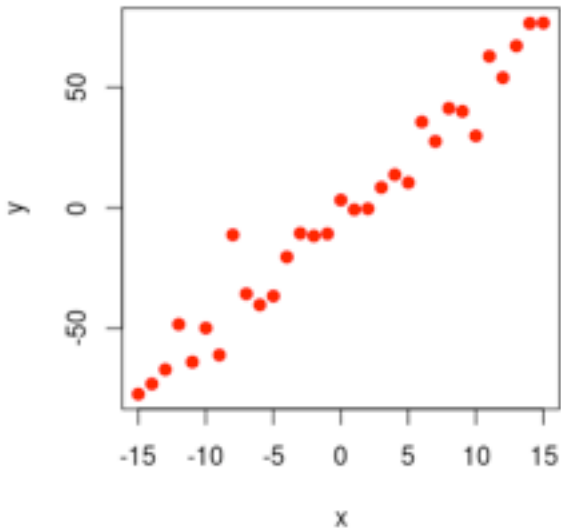
$$y \sim N(a_1 * x, 10)$$



a_1



Rejection sampling



1. draw random values from priors
2. calculate likelihood

i.e. the probability of observing data y given the model and drawn parameter values:

Model:

$$y \sim N(a * x, 10)$$

Priors:

$$P(a) \sim U[a_{\min}, a_{\max}]$$

$$b = 0$$

$$sd = 10$$

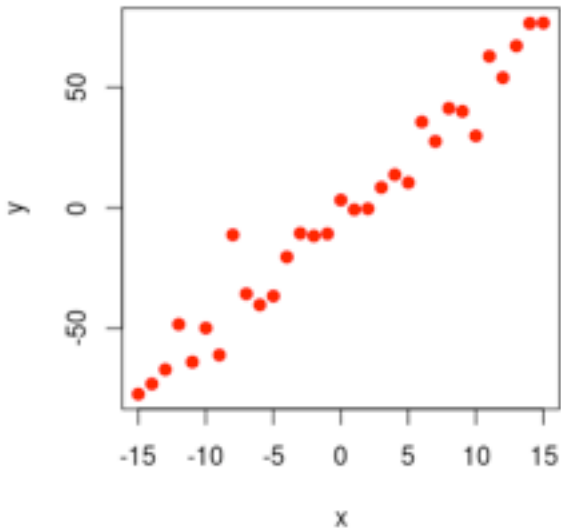
$$y \sim N(a_1 * x, 10)$$

probability density function
for normal distribution

$$\frac{1}{sd\sqrt{2\pi}} e^{-\frac{(y - (b + ax))^2}{2sd^2}}$$

in R: `sum(dnorm(y, mean = 0 + a1*x, sd = 10))`

Rejection sampling



1. draw random values from priors
2. calculate likelihood
3. accept value proportional to likelihood

Model:

$$y \sim N(a * x, 10)$$

$$P(\theta | Y) = \frac{P(Y | \theta) P(\theta)}{P(Y)}$$

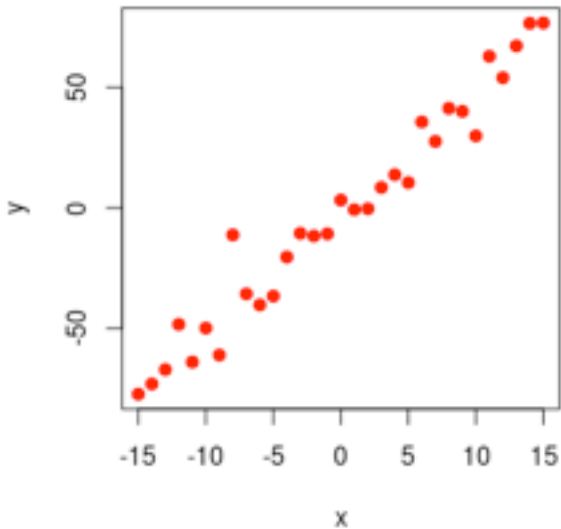
Priors:

$$P(a) \sim U[a_{\min}, a_{\max}] \longrightarrow a_1$$

$$b = 0$$

$$sd = 10$$

Rejection sampling



1. draw random values from priors
2. calculate likelihood
3. accept value proportional to likelihood

Model:

$$y \sim N(a \cdot x, 10)$$

Priors:

$$P(a) \sim U[a_{\min}, a_{\max}]$$

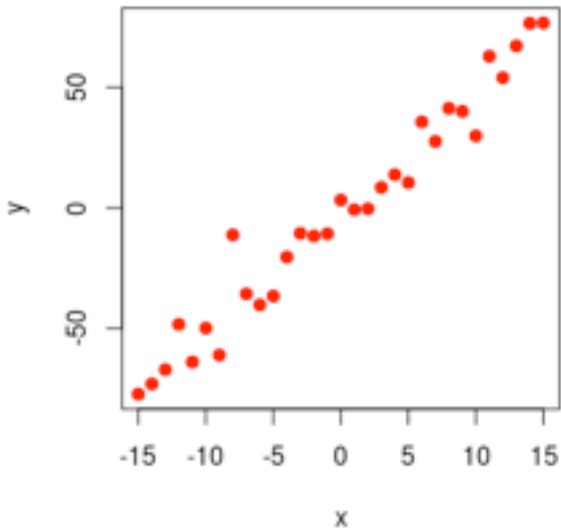
$$b = 0$$

$$sd = 10$$

$$P(a | y) \propto P(y | a) P(a)$$

unnormalized
posterior PDF

Rejection sampling



Model:

$$y \sim N(a \cdot x, 10)$$

Priors:

$$P(a) \sim U[a_{\min}, a_{\max}]$$

$$b = 0$$

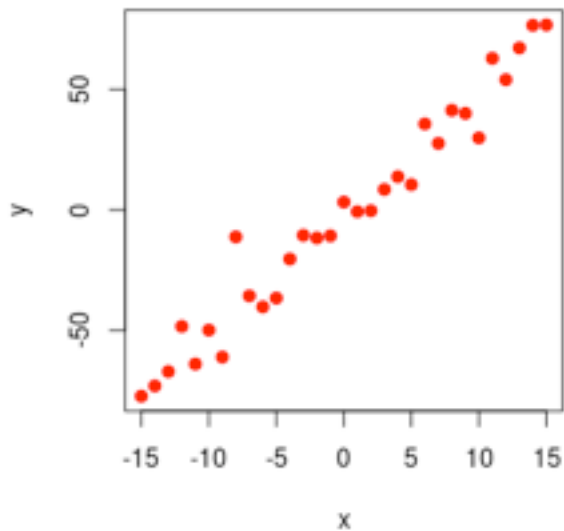
$$sd = 10$$

- ~~1. draw random values from priors~~
- ~~2. calculate likelihood~~
- ~~3. accept value proportional to likelihood~~

a. Sample many values
from prior

```
Asamples <- runif(1000, a_min, a_max)
```

Rejection sampling



- ~~1. draw random values from priors~~
- ~~2. calculate likelihood~~
- ~~3. accept value proportional to likelihood~~

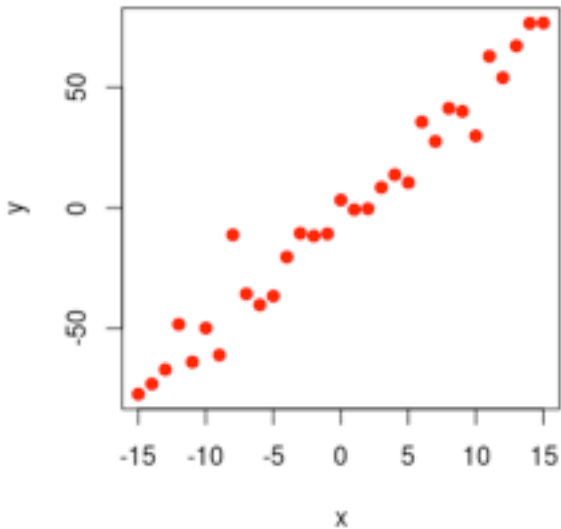
Model:

$$y \sim N(a*x, 10)$$

- a. Sample many values from prior
- b. Calculate unnormalized posterior values

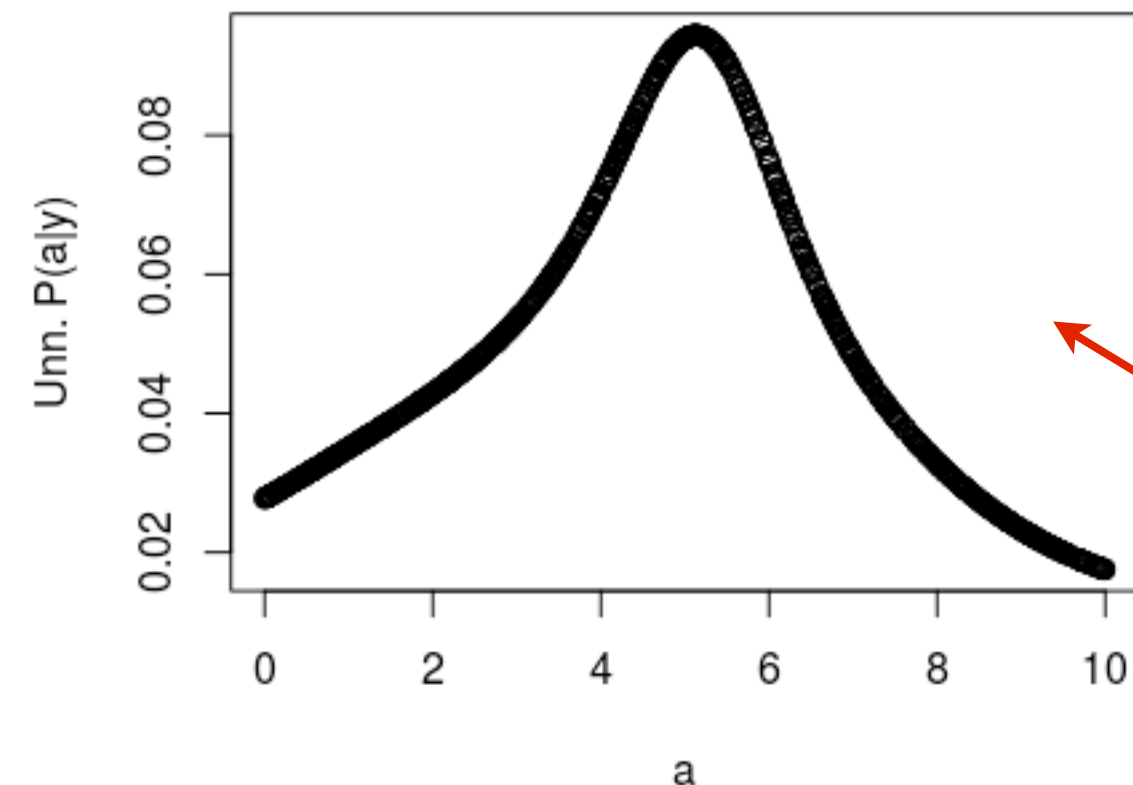
```
for(i in 1:1000)
  LL      <- sum(dnorm(y, mean = Asamples[i]*x, sd =10))
  prior   <- dunif(Asamples[i], a_min, a_max)
  Psamples <- LL * prior
```

Rejection sampling



- ~~1. draw random values from priors~~
- ~~2. calculate likelihood~~
- ~~3. accept value proportional to likelihood~~

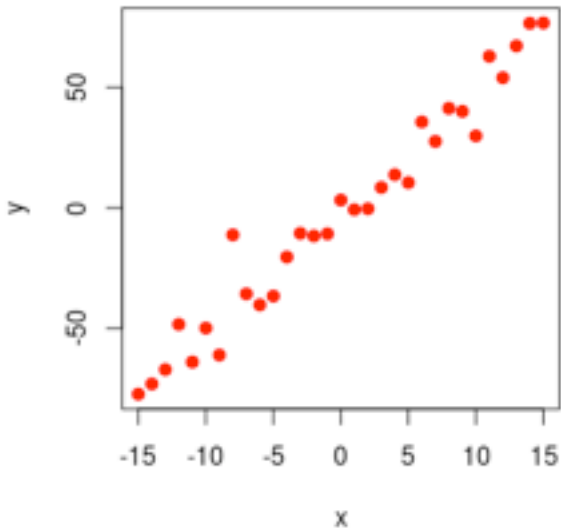
Unnormalized Posterior



- a. Sample many values from prior
- b. Calculate unnormalized posterior values

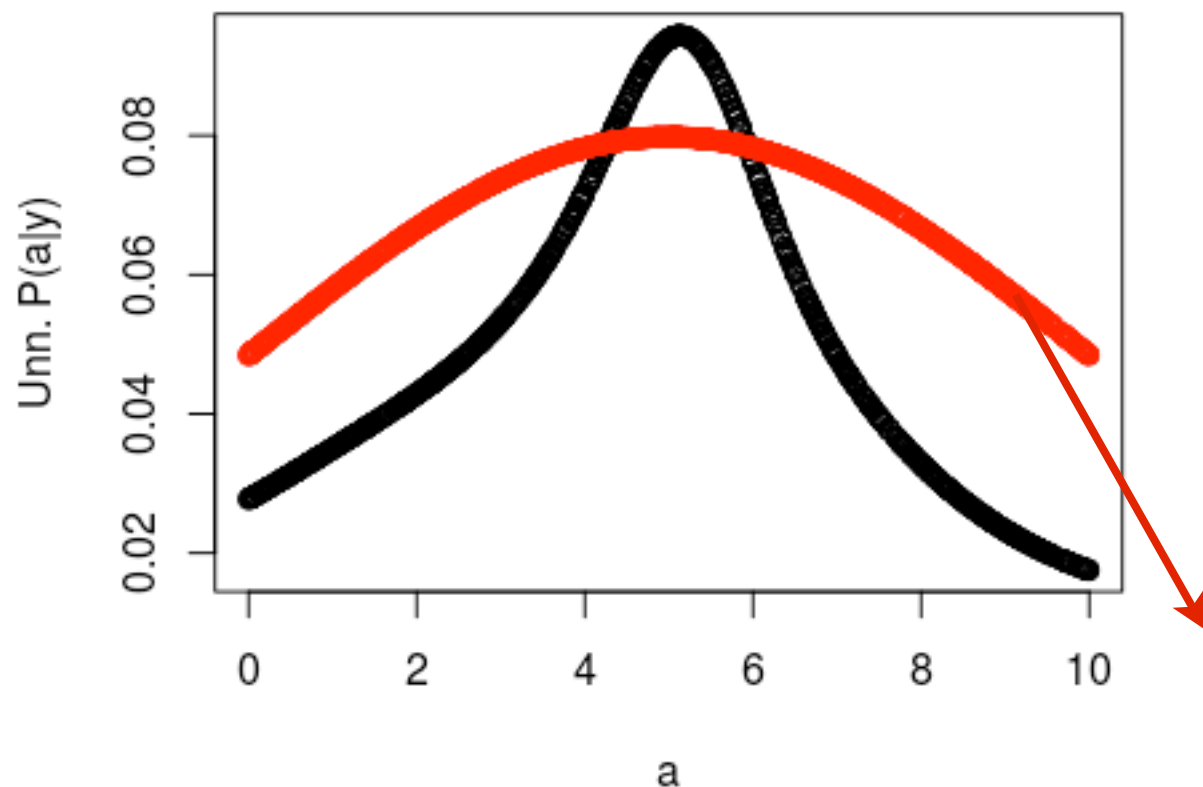
This is proportional to what I want to sample from

Rejection sampling



- ~~1. draw random values from priors~~
- ~~2. calculate likelihood~~
- ~~3. accept value proportional to likelihood~~

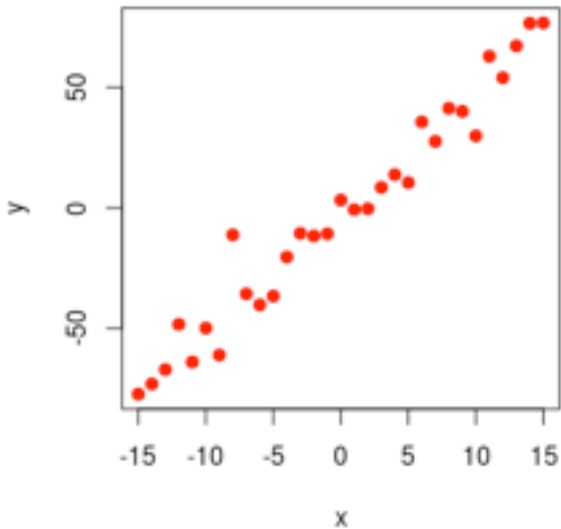
Unnormalized Posterior



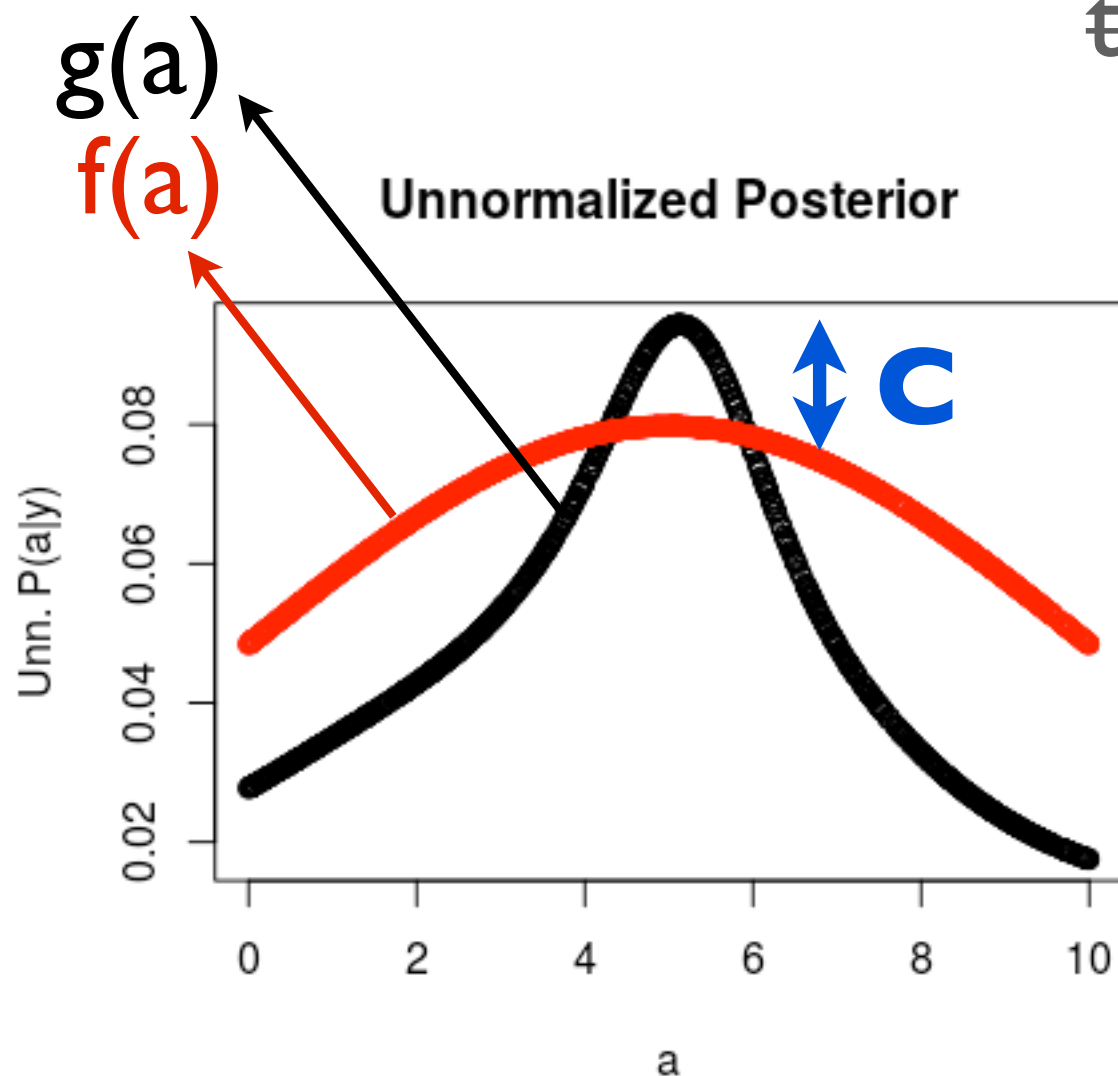
- a. Sample many values from prior
- b. Calculate unnormalized posterior values
- c. Choose a distribution from which sampling is easy

N(5,5) : proposal or envelope distribution

Rejection sampling

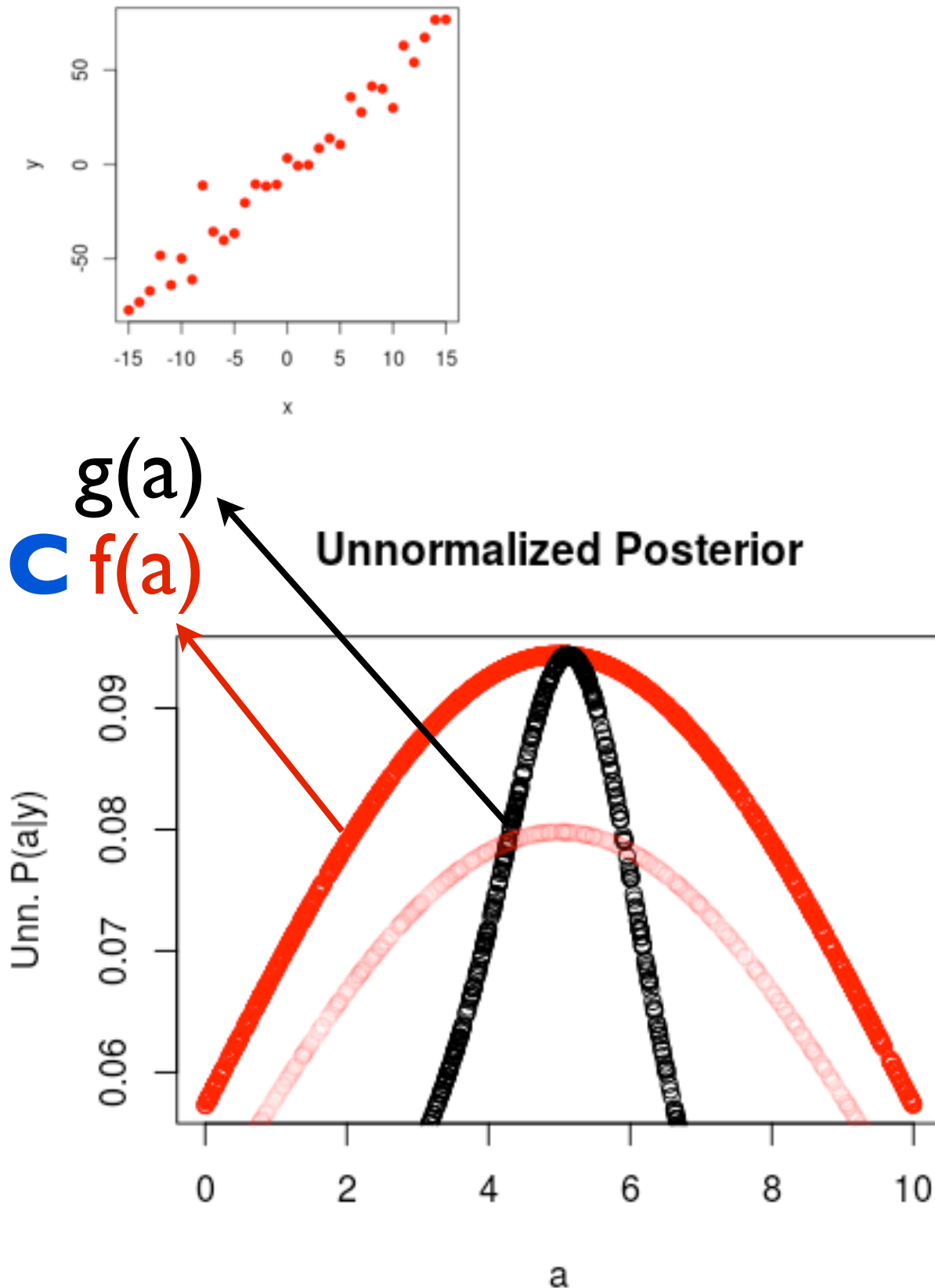


- ~~1. draw random values from priors~~
- ~~2. calculate likelihood~~
- ~~3. accept value proportional to likelihood~~



- Sample many values from prior
- Calculate unnormalized posterior values
- Choose a distribution from which sampling is easy
- Determine scaling constant : $C = \max(g(a)/f(a))$

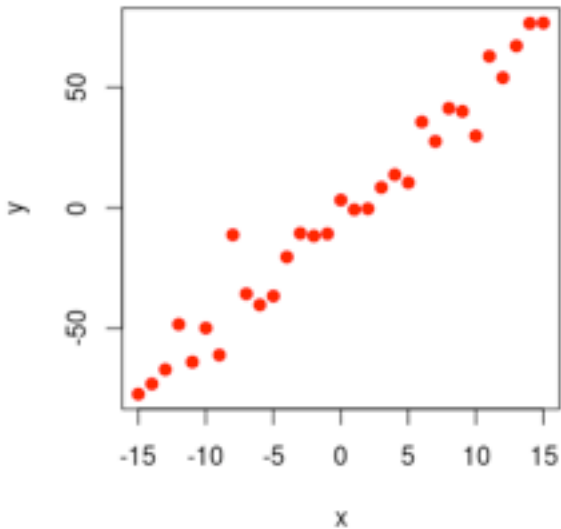
Rejection sampling



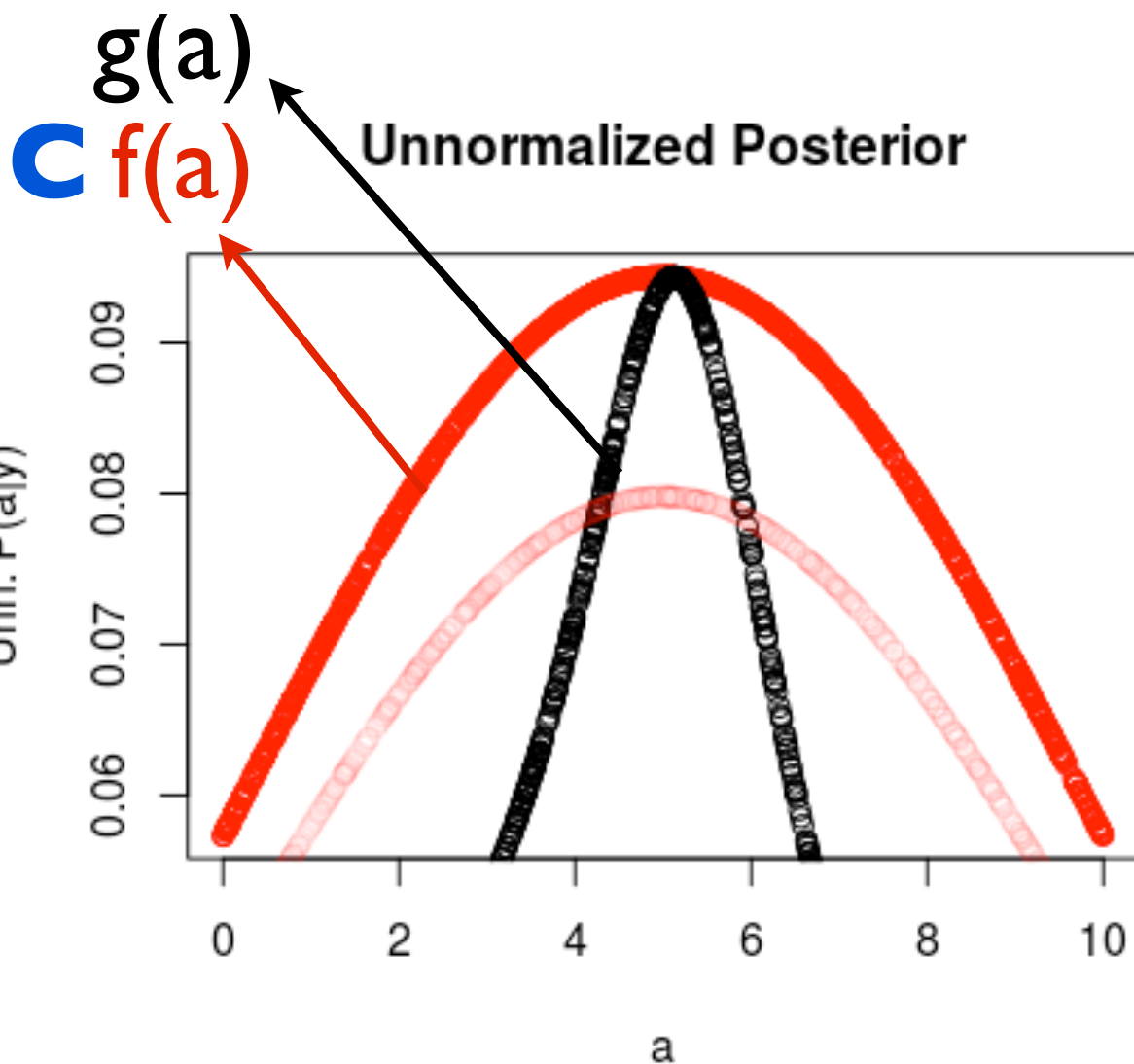
- Sample many values from prior
- Calculate unnormalized posterior values
- Choose a distribution from which sampling is easy
- Determine scaling constant : $C = \max(g(a)/f(a))$
- Scale envelope distribution

You only need to determine “ $C*f(a)$ ” once!

Rejection sampling



1. draw random value from $f(a)$
2. calculate $g(a)$ and $f(a)$
3. calculate u



$$P(a_1 | y) \propto P(y | a_1) P(a_1)$$

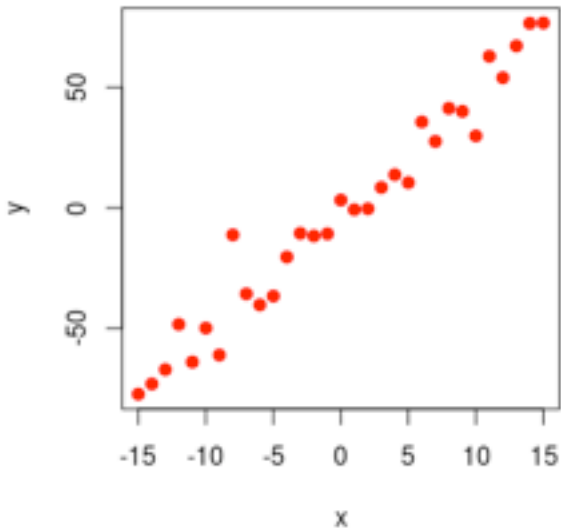
$$g(a_1) =$$

$$\text{sum}(\text{dnorm}(y, \text{mean} = 0 + a_1 * x, \text{sd} = 10)) * \text{dunif}(a_1, a_{\min}, a_{\max})$$

$$f(a_1) = \text{dnorm}(a_1, \text{mean} = 5, \text{sd} = 5)$$

$$u = \frac{g(a_1)}{C * f(a_1)}$$

Rejection sampling

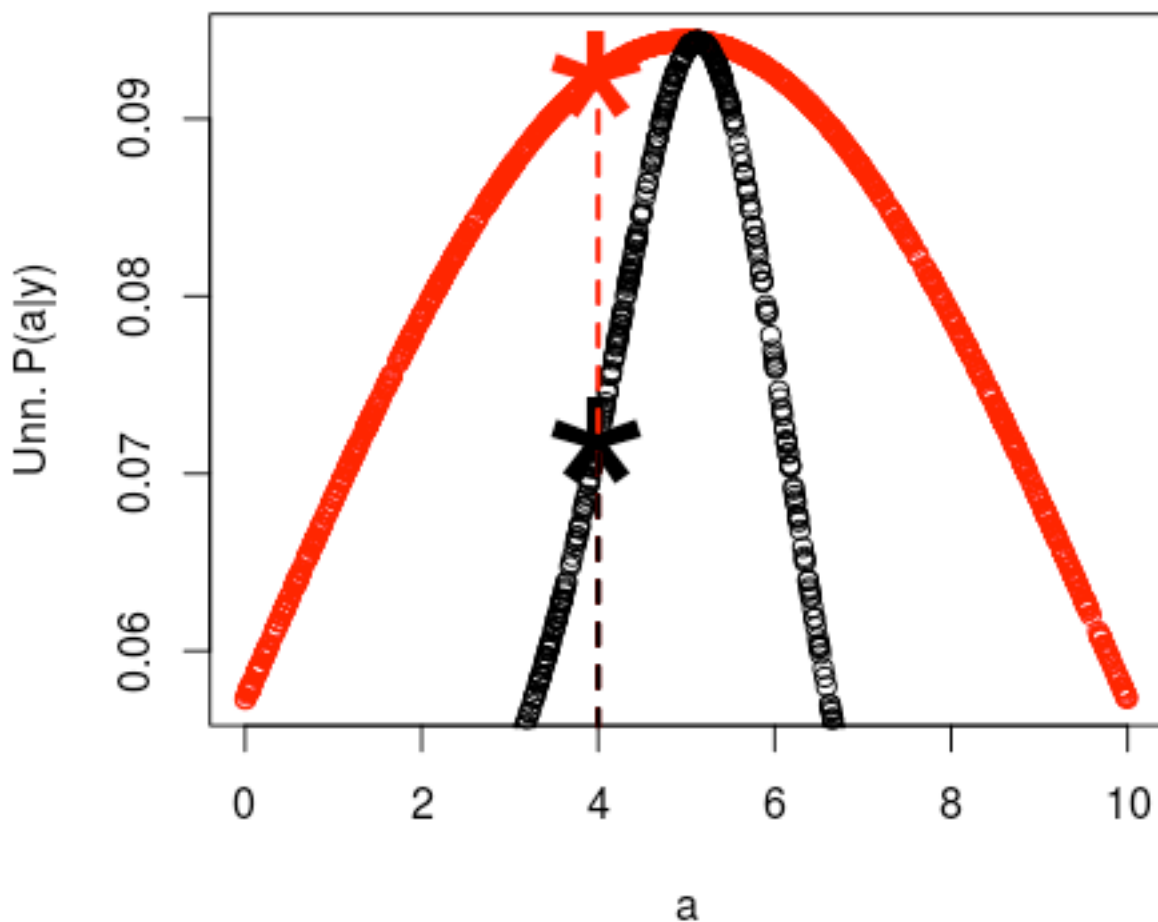


Draw random value: $a_1 = 4$

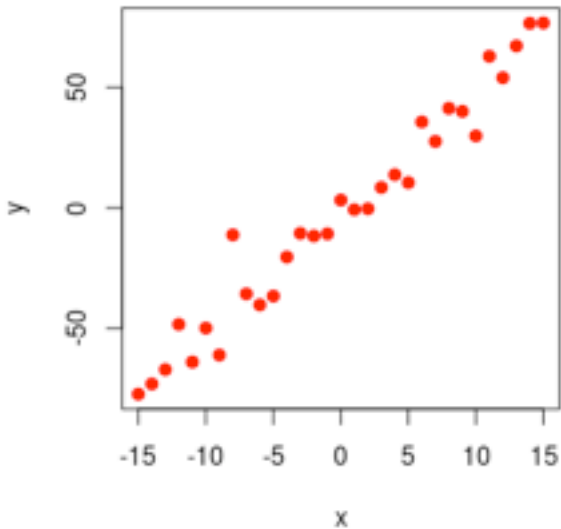
Calculate $f(a)$ and $g(a)$ $g(a_1), f(a_1)$

Calculate u : $u = \frac{g(a_1)}{C * f(a_1)}$

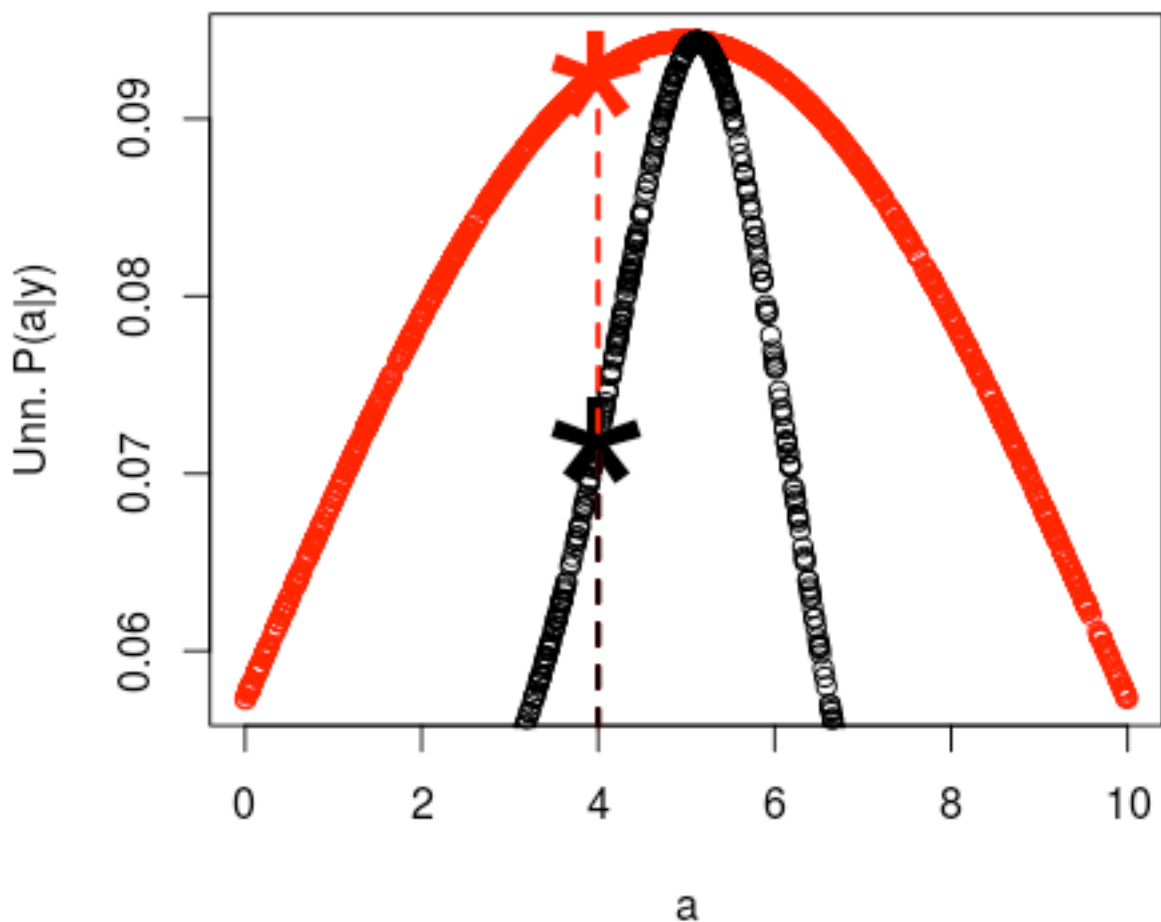
Unnormalized Posterior



Rejection sampling



Unnormalized Posterior



Draw random value: $a_1 = 4$

Calculate $f(a)$ and $g(a)$ $g(a_1), f(a_1)$

Calculate u : $u = \frac{g(a_1)}{C * f(a_1)}$

Accept the proposed a_1 with probability u

based on a Bernoulli trial:

if

$\text{runif}(1,0,1) < u$

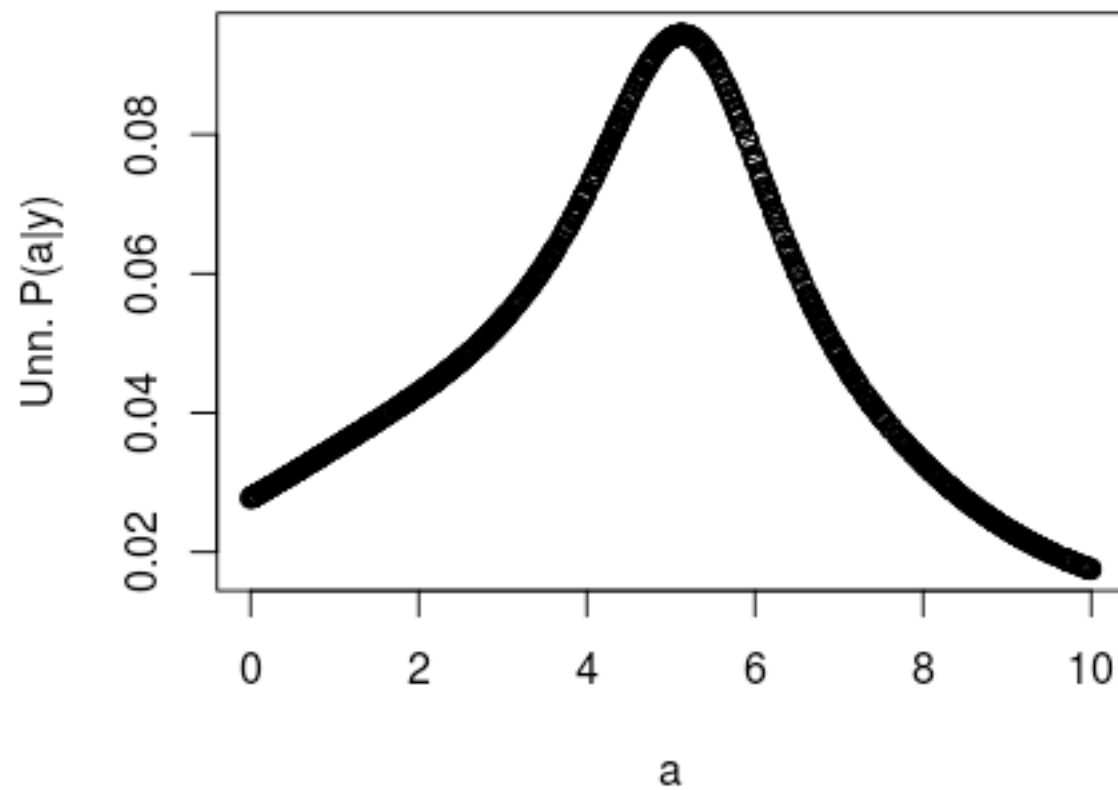
accept

else

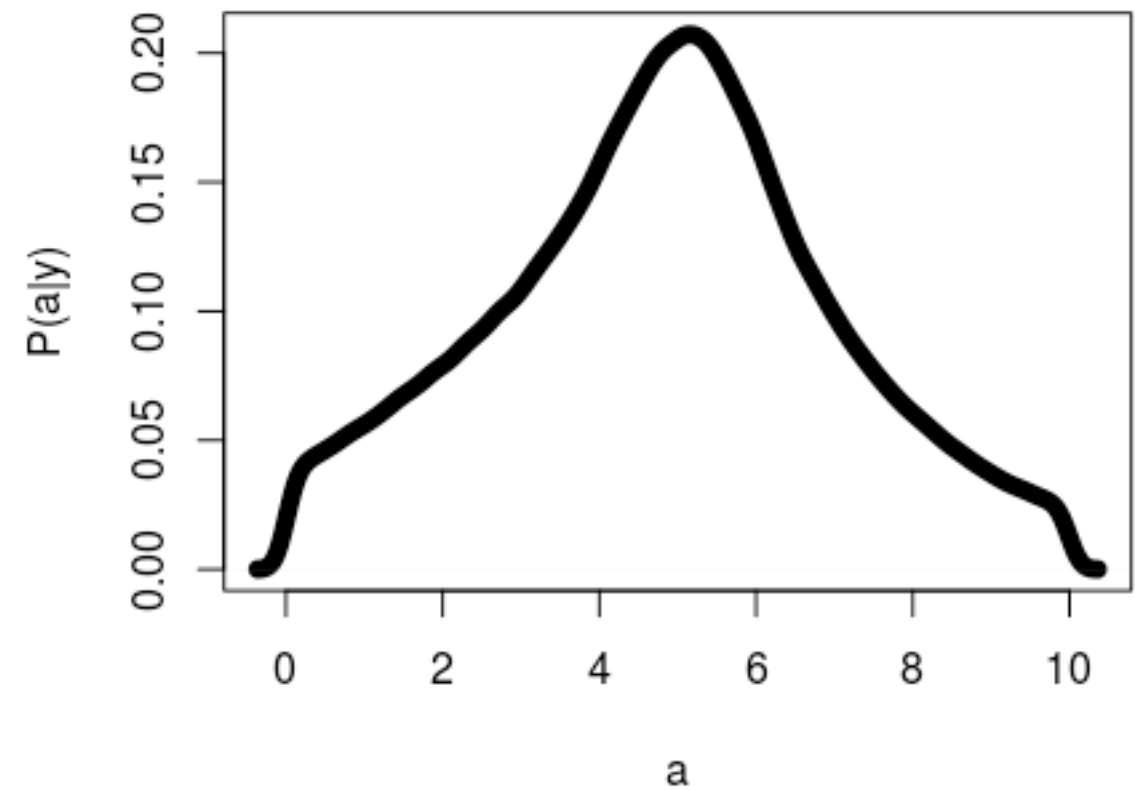
reject

Rejection sampling

Unnormalized Posterior

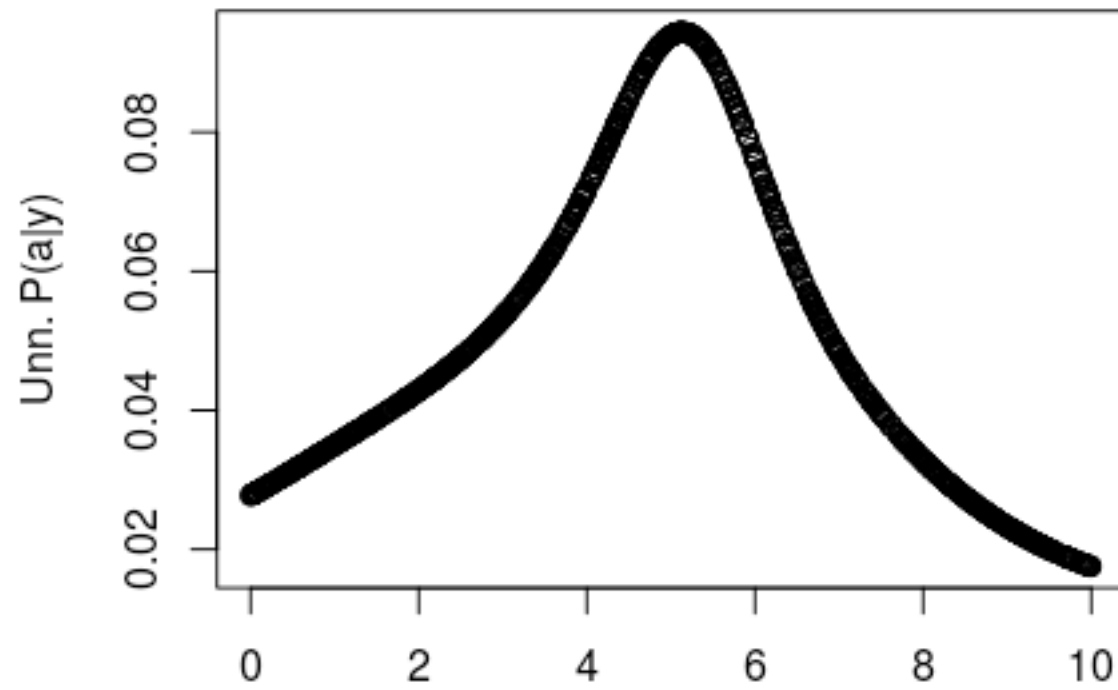


Posterior PDF



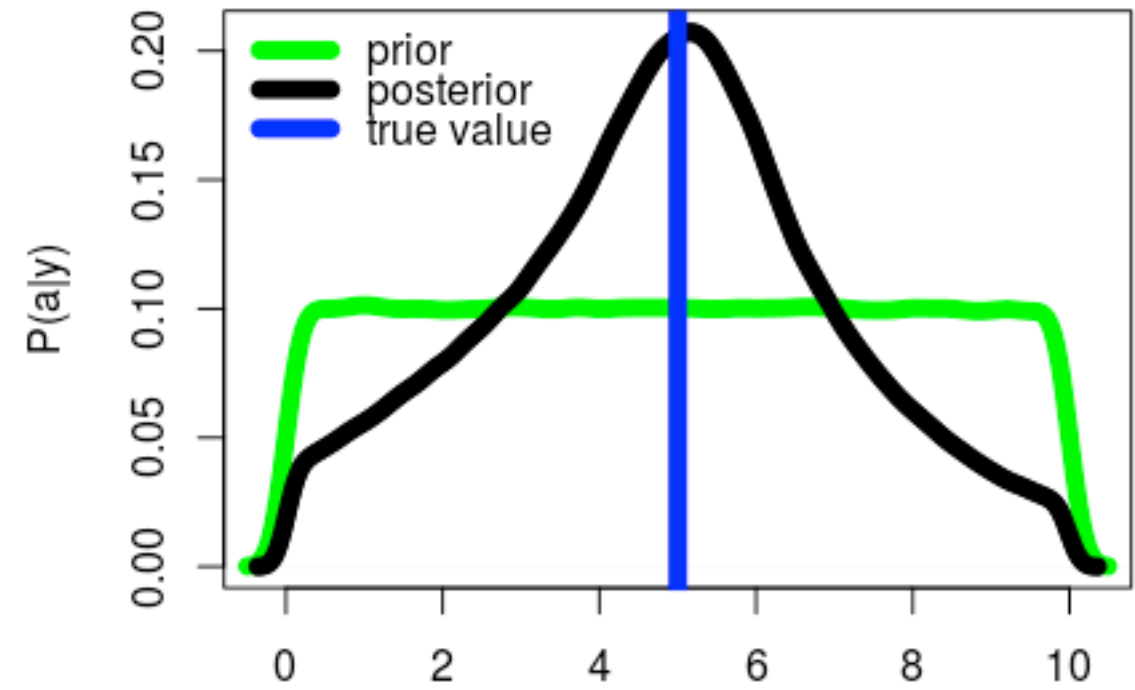
Rejection sampling

Unnormalized Posterior

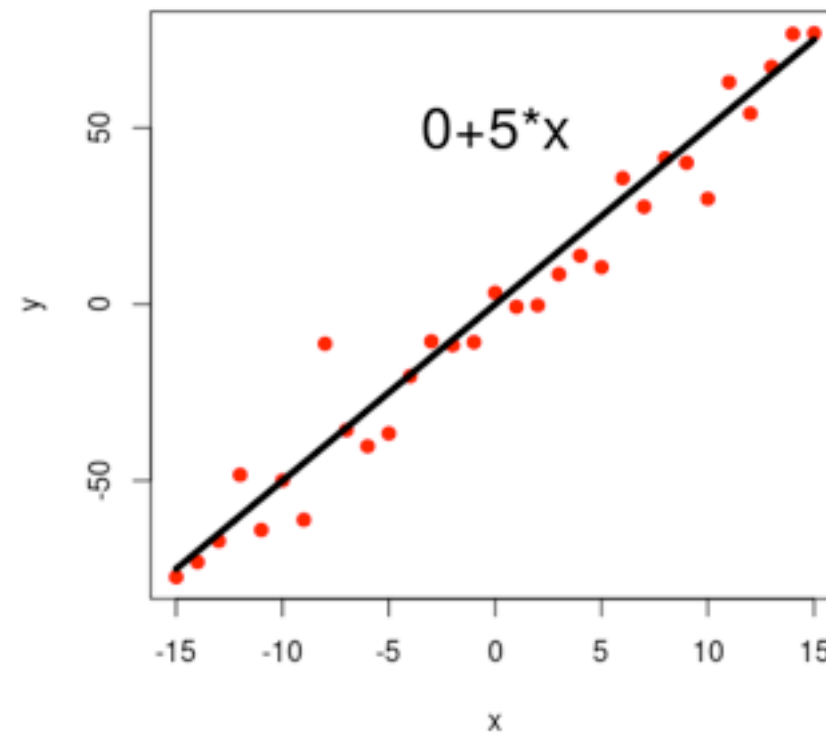


a

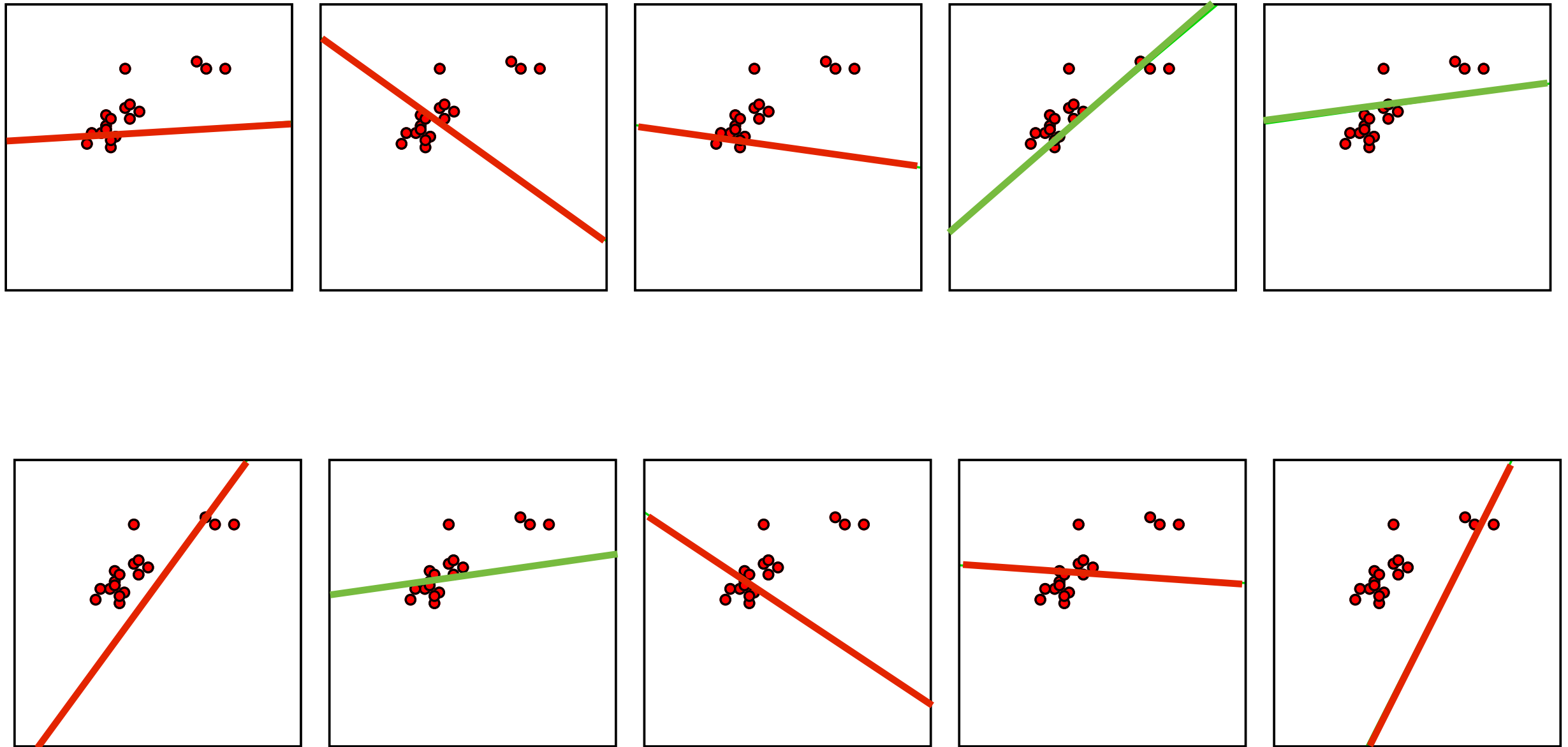
Posterior PDF



a



Rejection sampling



Rejection sampling

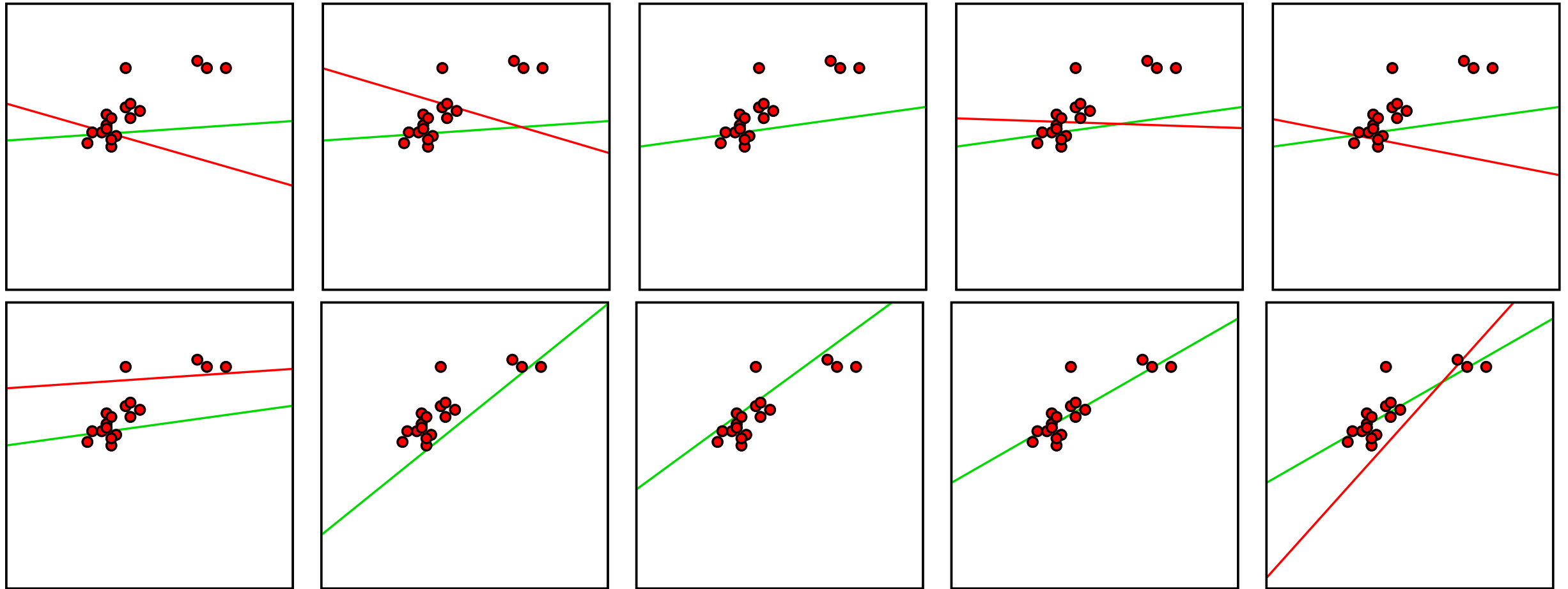
Pros:

- Parallelizable
- Easy to implement

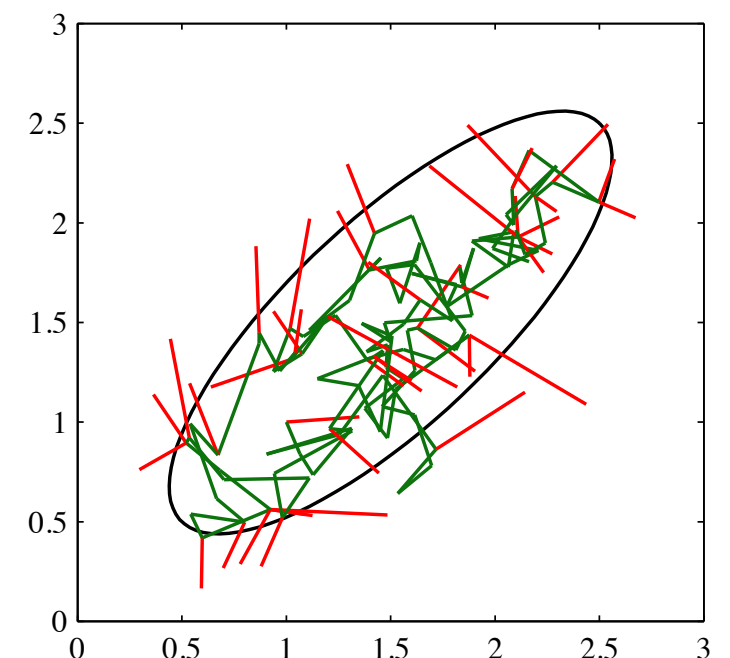
Cons:

- Suffers from curse of dimensionality

Markov Chain Monte Carlo



- Perturb parameters
- Accept if new params are supported more (*also accept sometimes even if not)
- Otherwise keep old parameters



Markov Chain Monte Carlo

- 1) Start from some initial parameter value
- 2) Evaluate the unnormalized posterior
- 3) Propose a new parameter value
- 4) Evaluate the new unnormalized posterior
- 5) Decide whether or not to accept the new value
- 6) Repeat 3-5

Markov Chain Monte Carlo

- 1) Start from some initial parameter value
- 2) Evaluate the unnormalized posterior
- 3) Propose a new parameter value  **dependent on
step 1/previous step**
- 4) Evaluate the new unnormalized posterior
- 5) Decide whether or not to accept the new value
- 6) Repeat 3-5

- Advantages

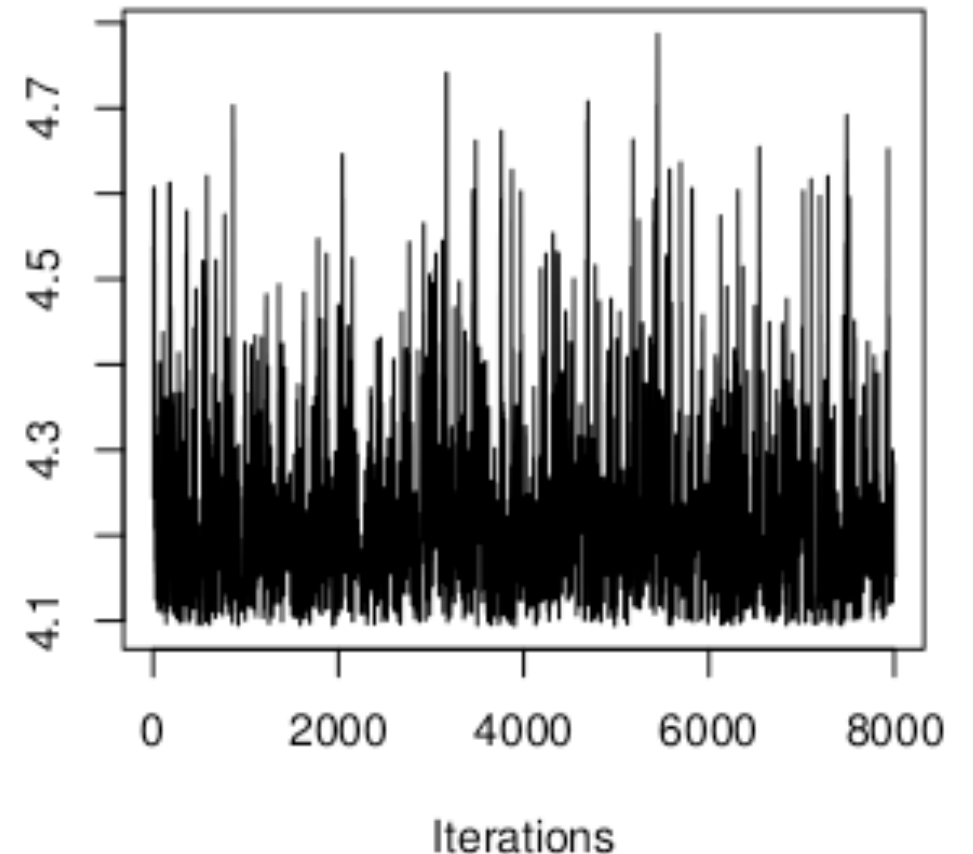
- Multi-dimensional
- Can be applied to
 - Whole joint PDF
 - Each dimension iteratively
 - Groups of parameters
- Simple
- Robust

- Disadvantages

- Sequential samples not independent
- Computationally intensive
- Discard “Burn – in” period before convergence
- Assessing convergence

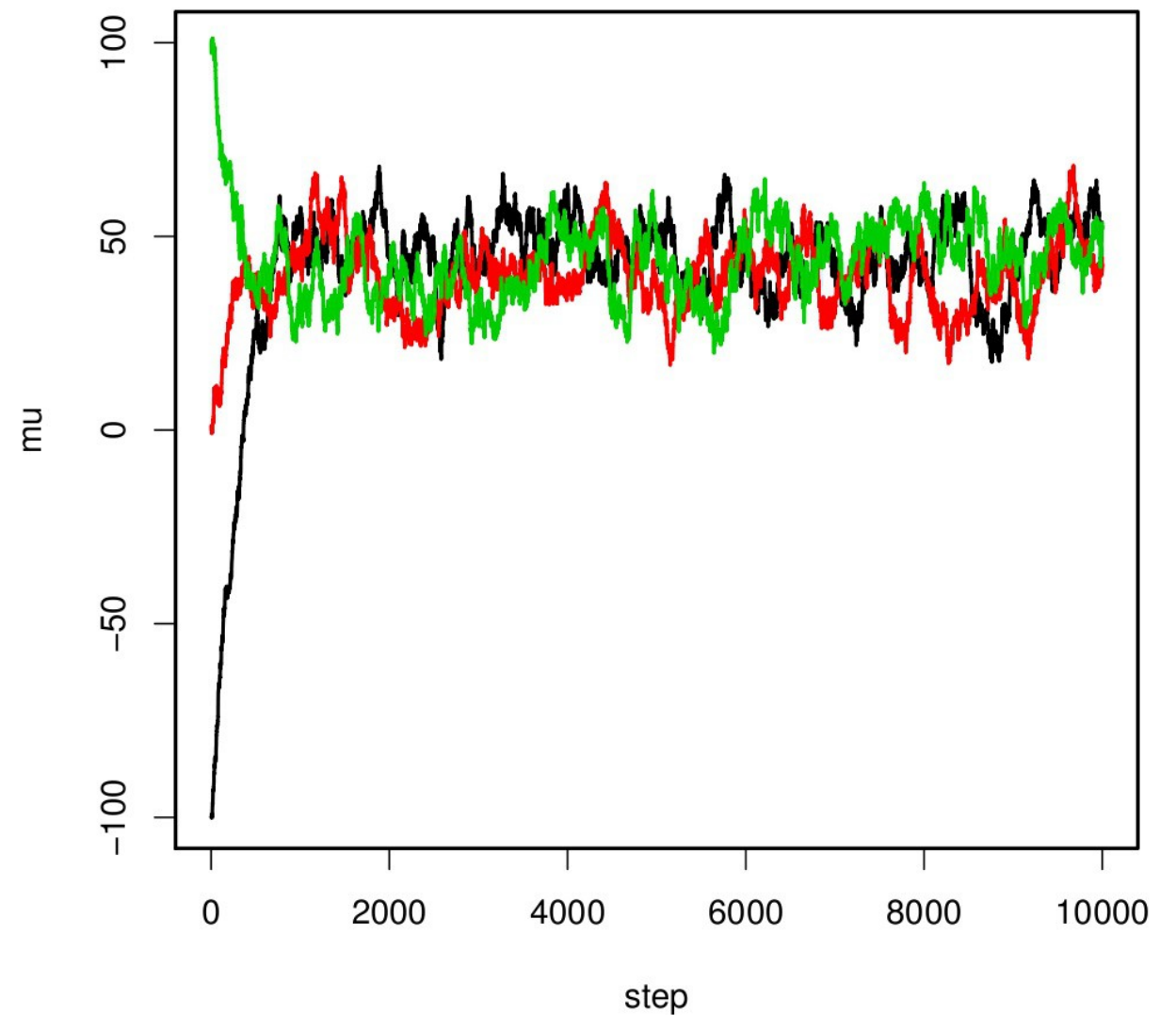
How to assess convergence?

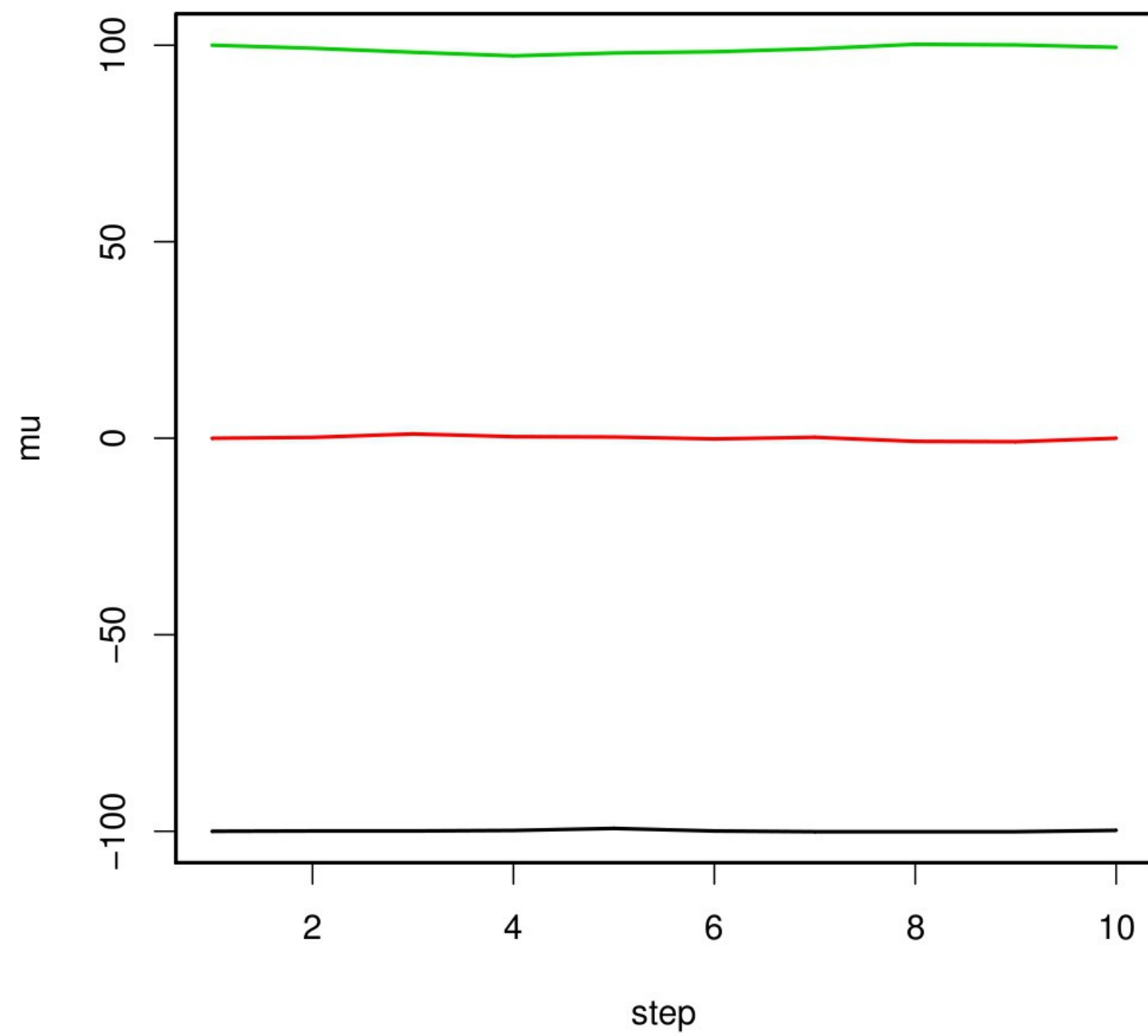
- Visual inspection

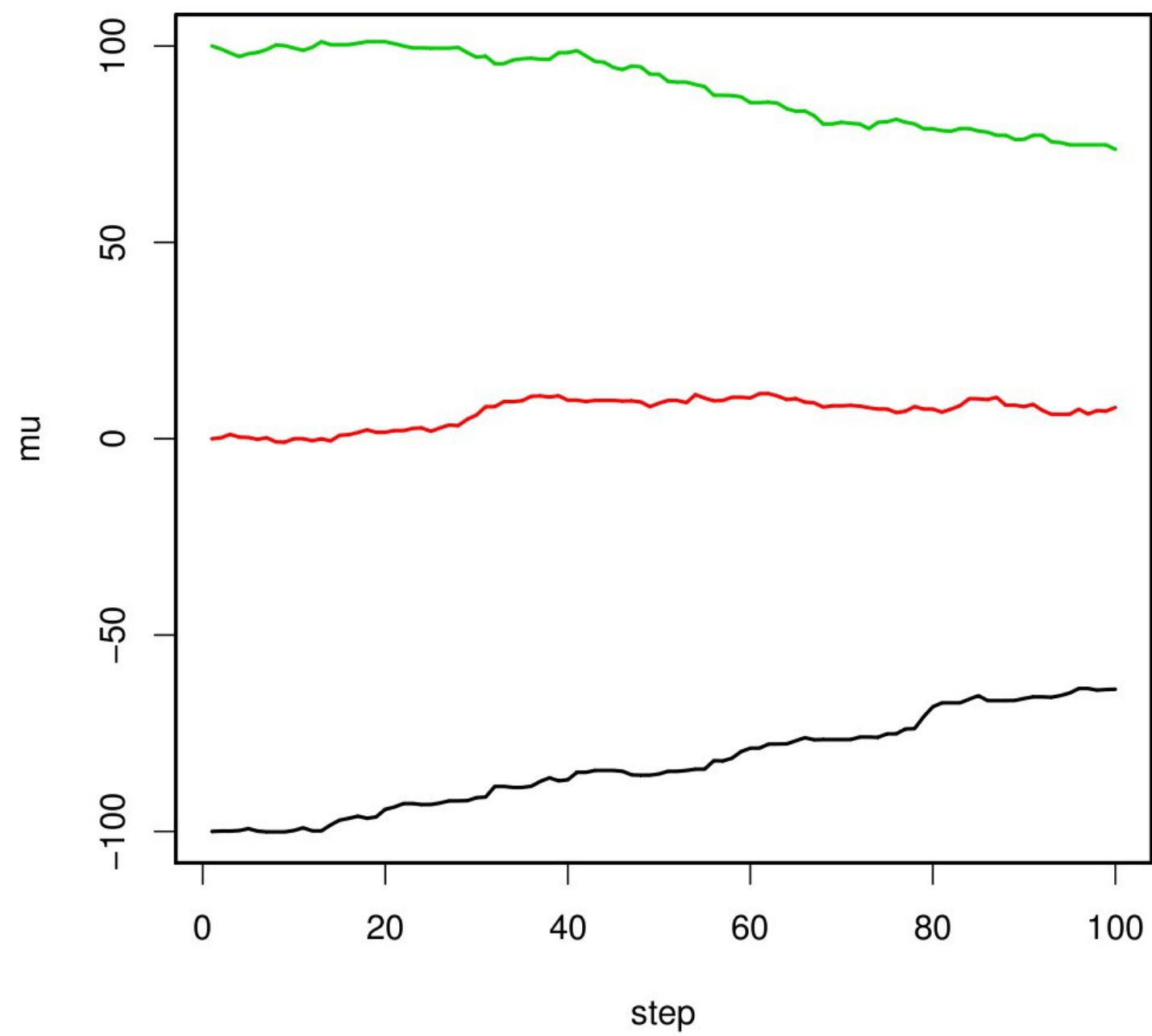


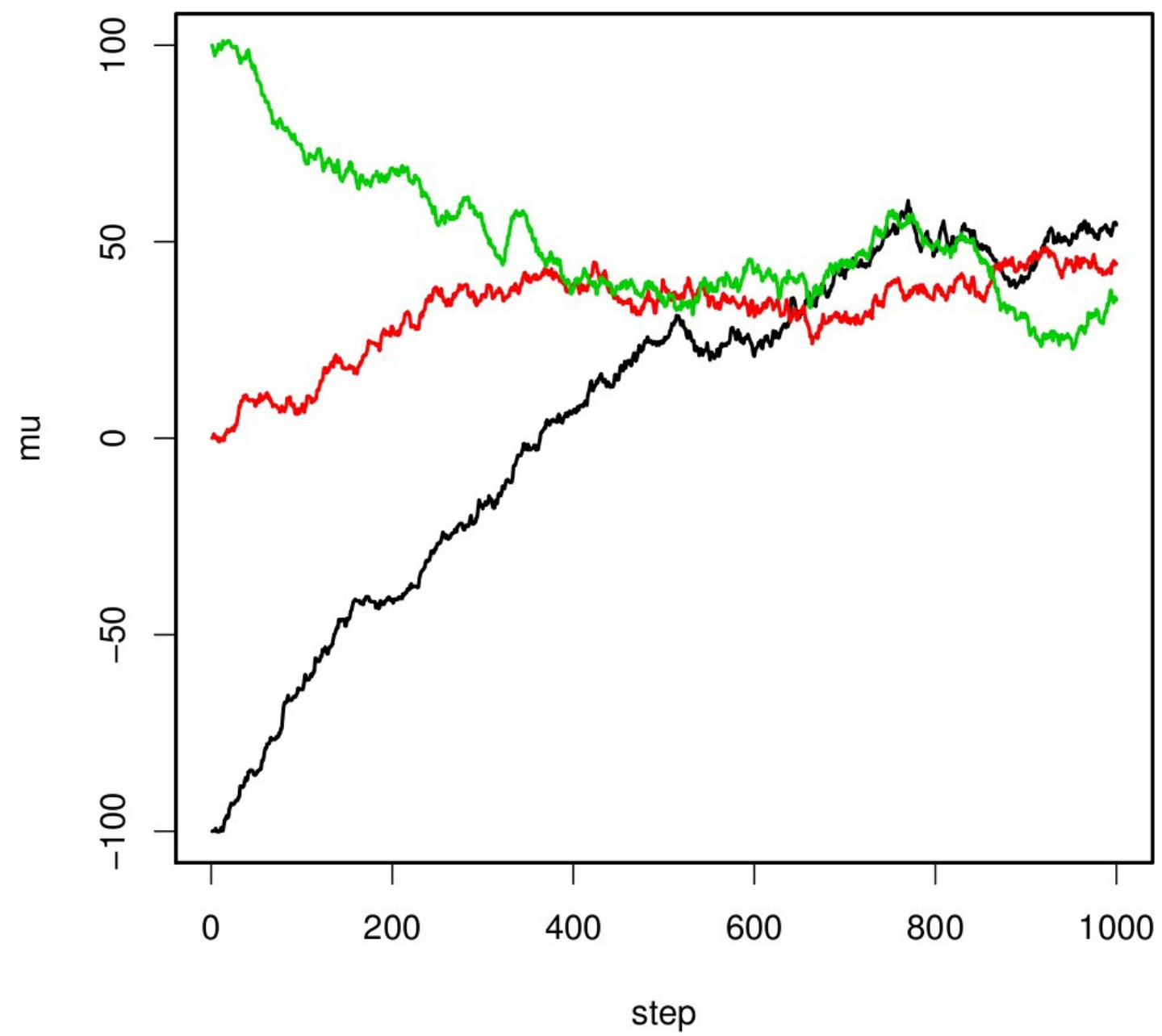
How to assess convergence?

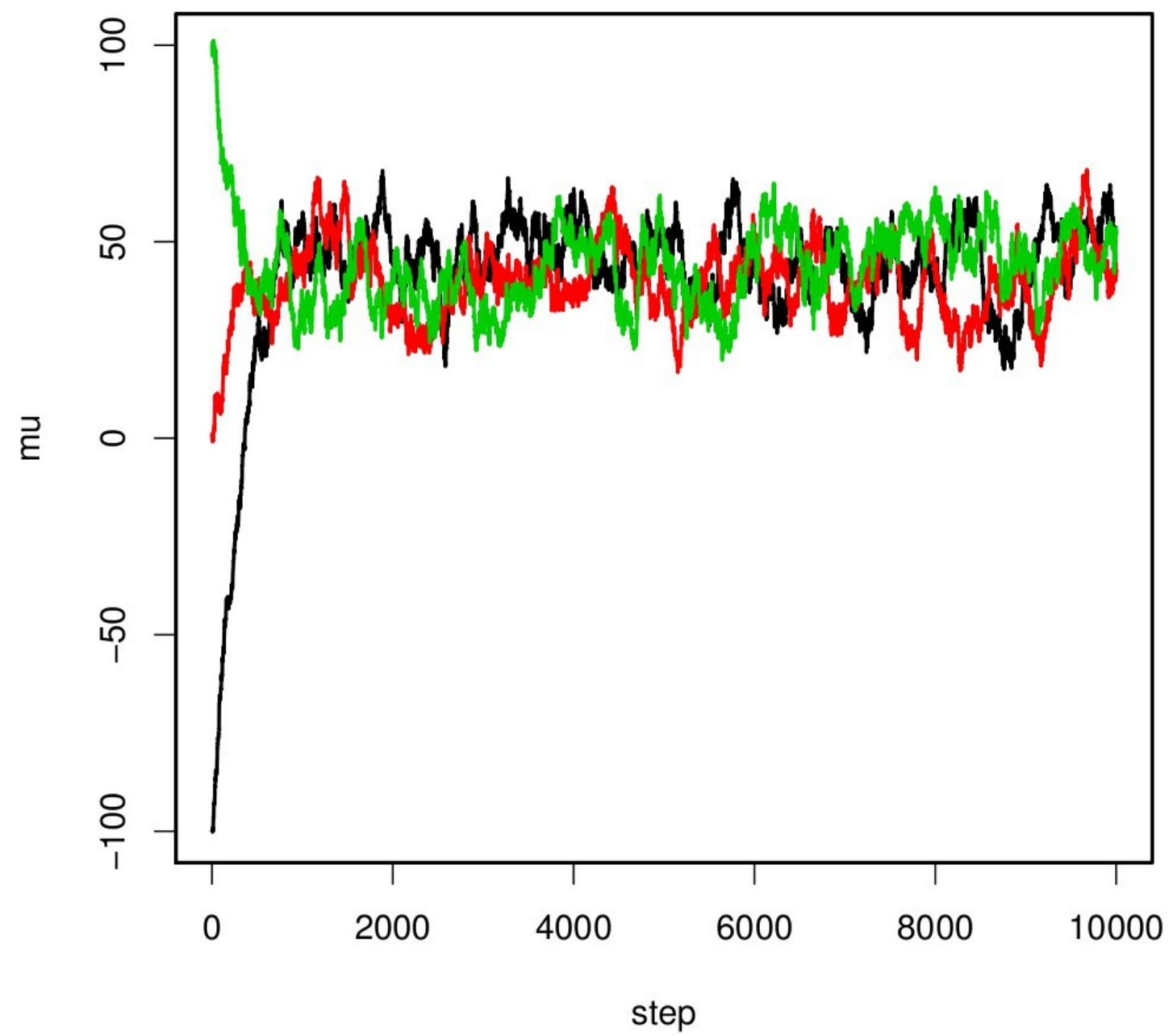
- Visual inspection
- Multiple chains



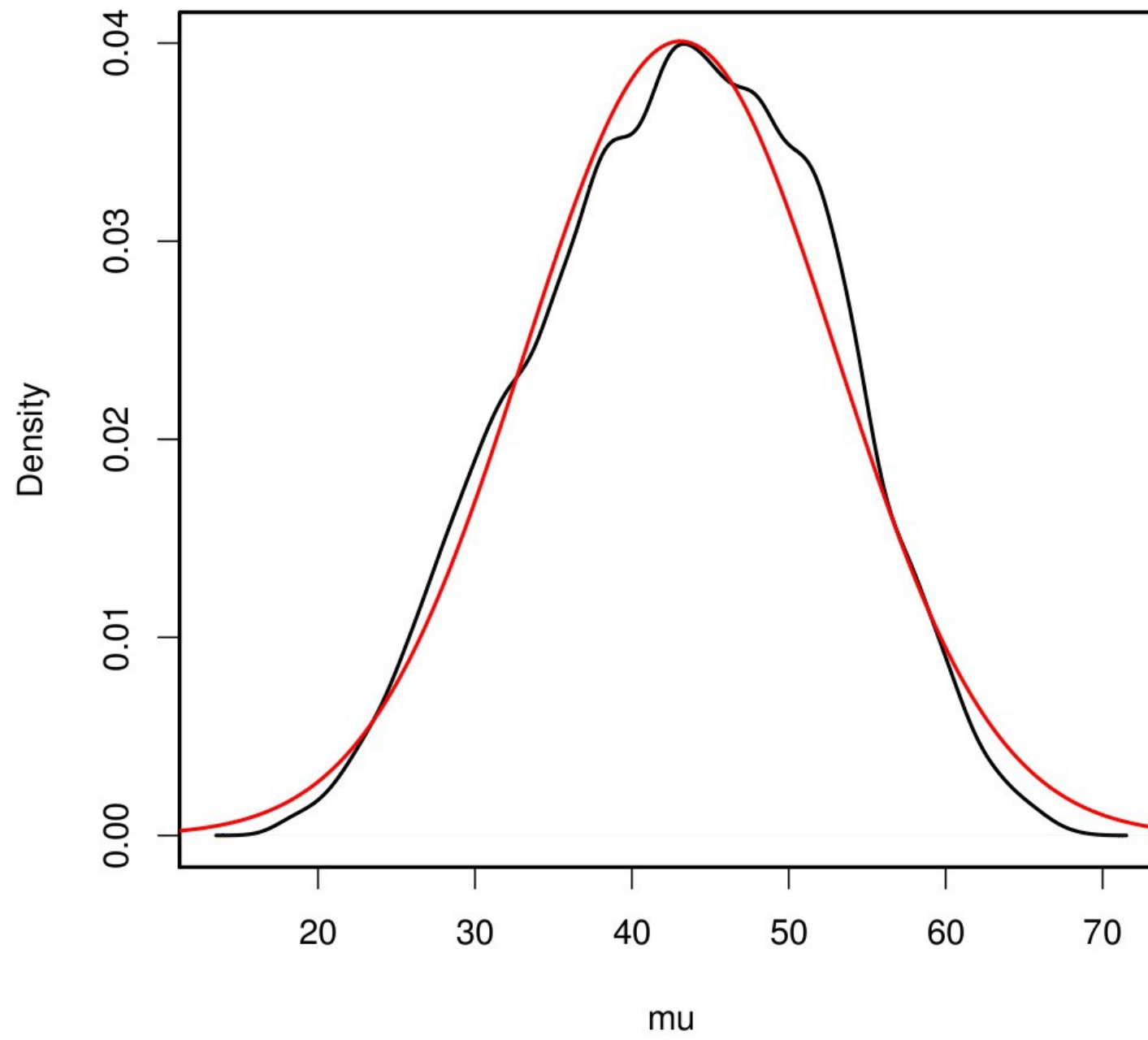








MCMC Posterior Density



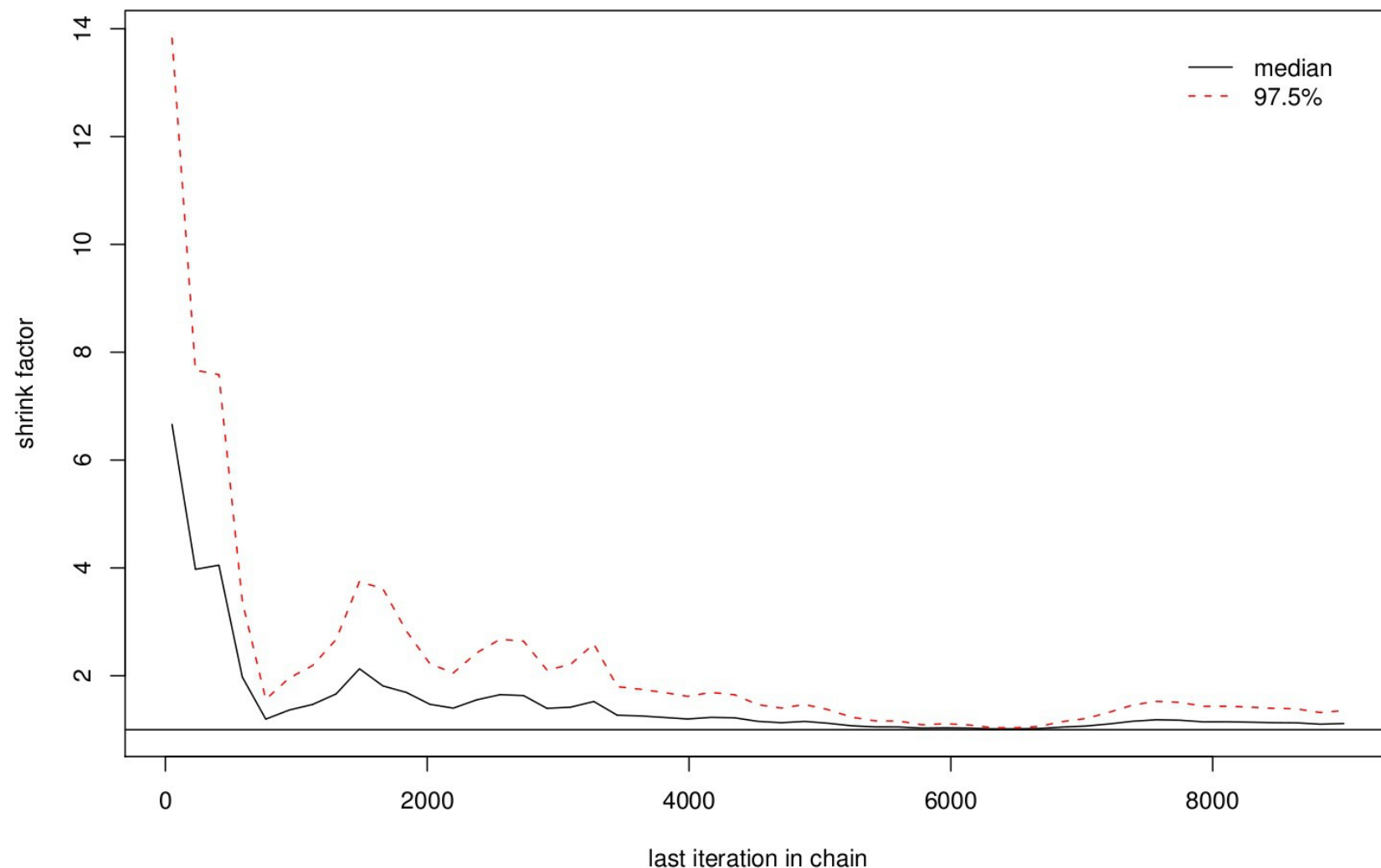
How to assess convergence?

- Visual inspection
- Multiple chains
- Convergence stats

Convergence Statistics

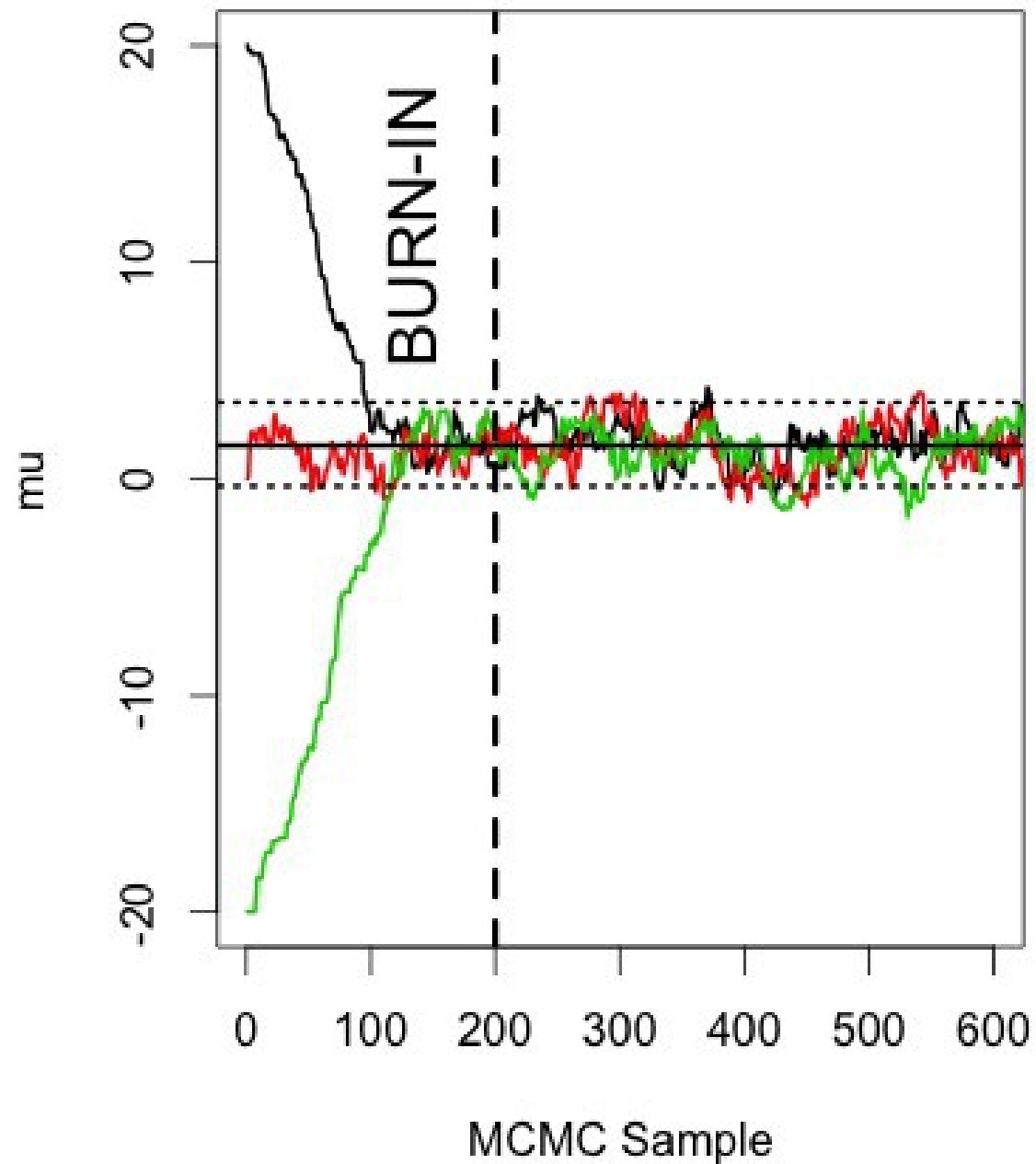
- Brooks Gelman Rubin
 - Within vs among chain variance
 - Should converge to 1

$$\hat{R} = \frac{B}{W}$$

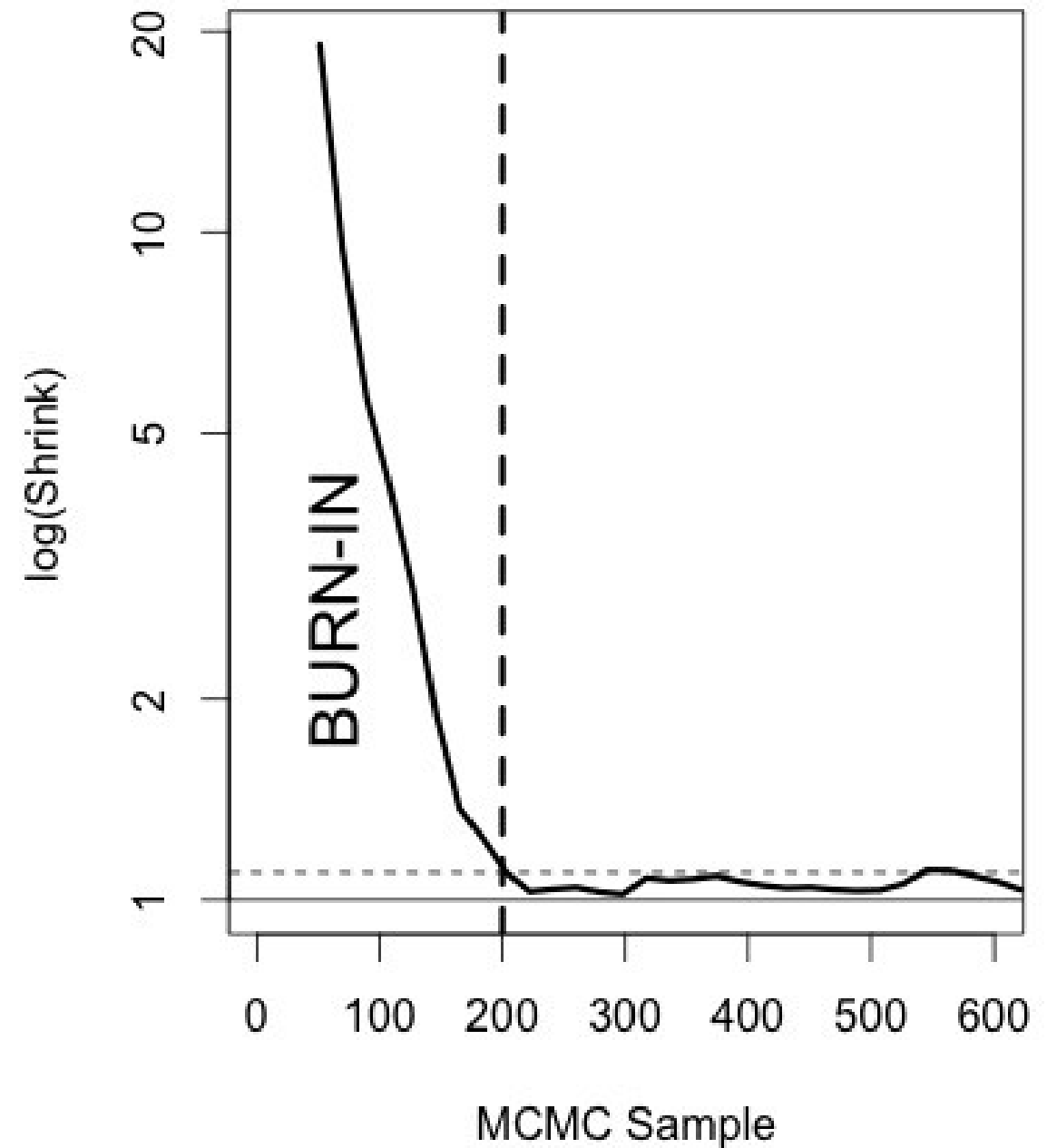


Convergence Statistics

Trace Plot



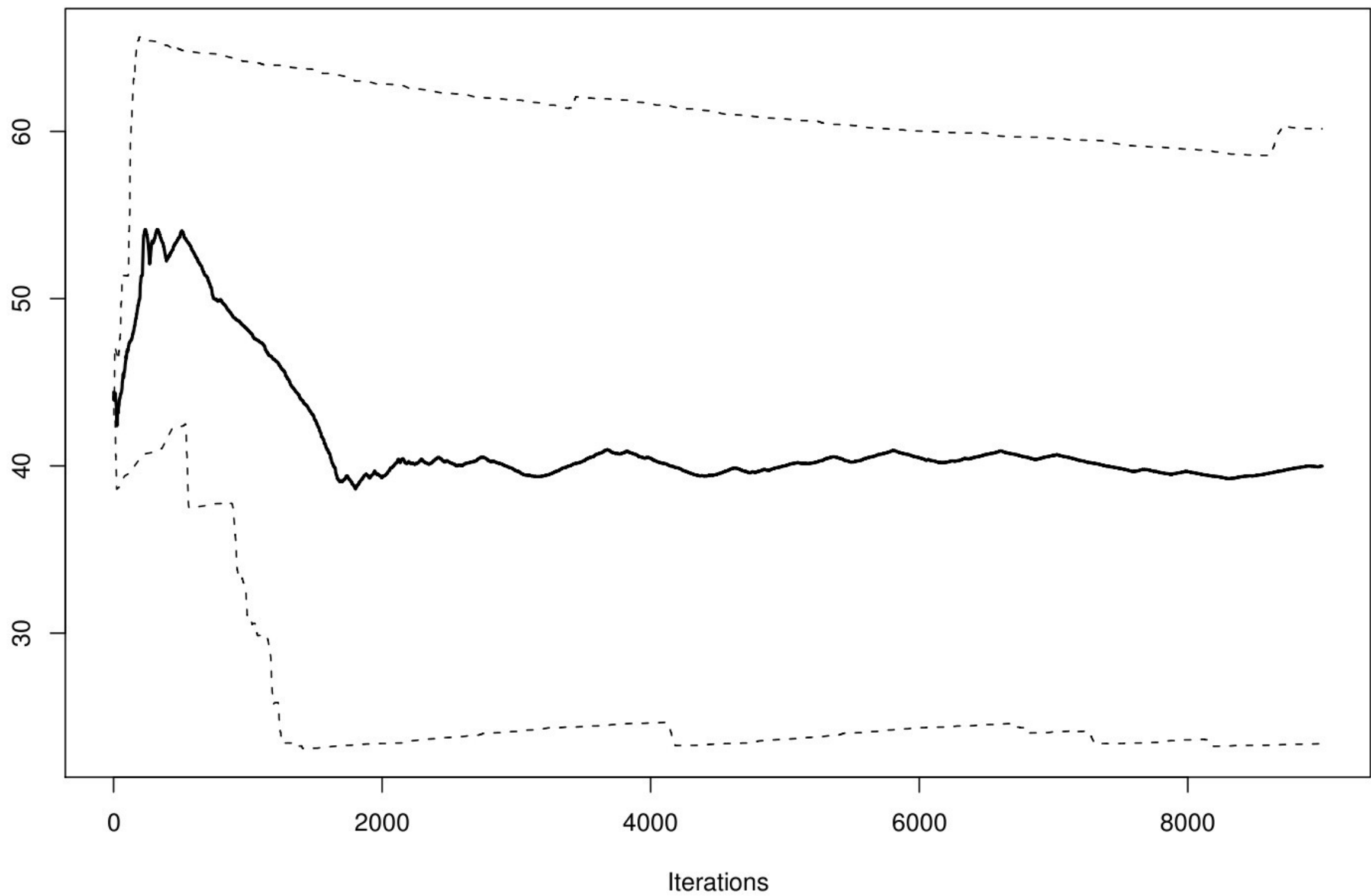
GBR Diagnostic



How to assess convergence?

- Visual inspection
- Multiple chains
- Convergence stats
- Quantiles

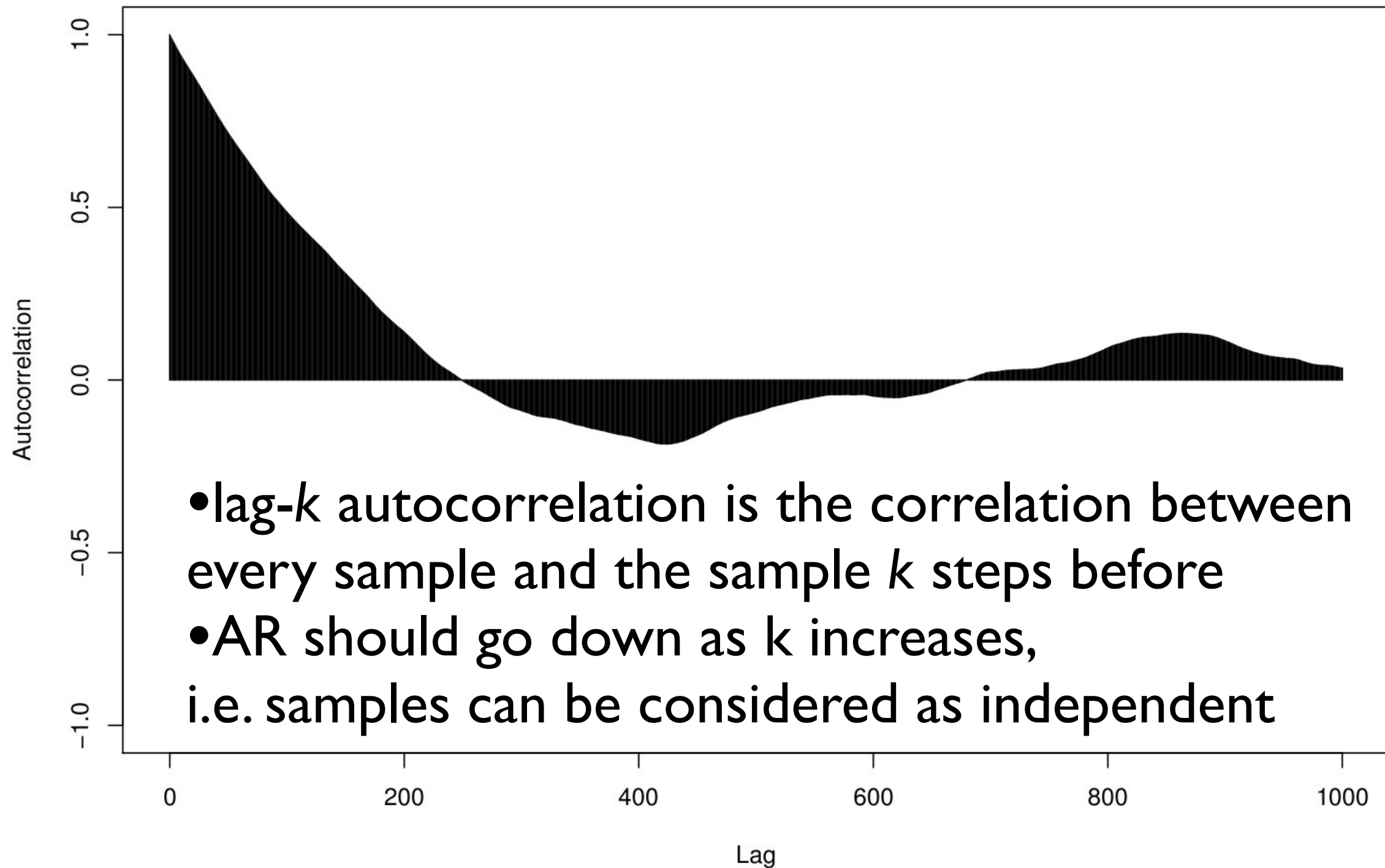
Quantiles



How to assess convergence?

- Visual inspection
- Multiple chains
- Convergence stats
- Quantiles
- Autocorrelation

Autocorrelation



How to assess convergence?

- Visual inspection
- Multiple chains
- Convergence stats
- Quantiles
- Autocorrelation
- Summary statistics

Summary Statistics

Analytical:

Mean	SD
43.09901	9.95037

MCMC:

Mean	SD	Naive SE	Time-series
43.05504	9.28108	0.05648	0.74503

Quantiles:

2.5%	25%	50%	75%	97.5%
24.98	36.46	43.39	49.99	60.01

How to assess convergence?

- Visual inspection
- Multiple chains
- Convergence stats
- Quantiles
- Autocorrelation
- Summary statistics
- Effective sample size
- Acceptance rate