

Approximating posterior distribution and some samplers

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Bayes' theorem

- 1) prior: $p(\text{parameter values}) - p(\theta)$
- 2) likelihood: $p(\text{data values} \mid \text{parameter values}) - p(D|\theta)$
- 3) posterior: $p(\text{parameter values} \mid \text{data values}) - p(\theta|D)$

Bayes' theorem Cont.

$$\begin{aligned}p(D, \theta) &= p(\theta) \cdot p(D|\theta) \\ &= p(D) \cdot p(\theta|D)\end{aligned}$$

$$p(D) \cdot p(\theta|D) = p(\theta) \cdot p(D|\theta)$$

We can conclude that:

$$p(\theta|D) = \frac{p(\theta) \cdot p(D|\theta)}{p(D)}$$

$$p(\theta|D) \propto p(\theta) \cdot p(D|\theta)$$

(Shape of) posterior = prior · likelihood

Grid approximation

1. Define a discrete grid of possible θ values.
2. Evaluate the prior pdf $f(\theta)$ and likelihood function $L(\theta|D)$ at each θ grid value.
3. Obtain a discrete approximation of posterior pdf $f(\theta|D)$ by:
 - ▶ calculating the product $f(\theta)L(\theta|D)$ at each θ grid value
 - ▶ normalizing the products so that they sum to 1 across all θ .
4. Randomly sample n θ grid values with respect to their corresponding normalized posterior probabilities.

Example



Figure 1: rainbow

2) Rejection Sampling

Sample data from a complicated distribution

- ▶ Target (distribution) function $f(x)$ — The “difficult to sample from” distribution. Our distribution of interest!
- ▶ Proposal (distribution) function $g(x)$ — The proxy distribution from which we can sample.

1)

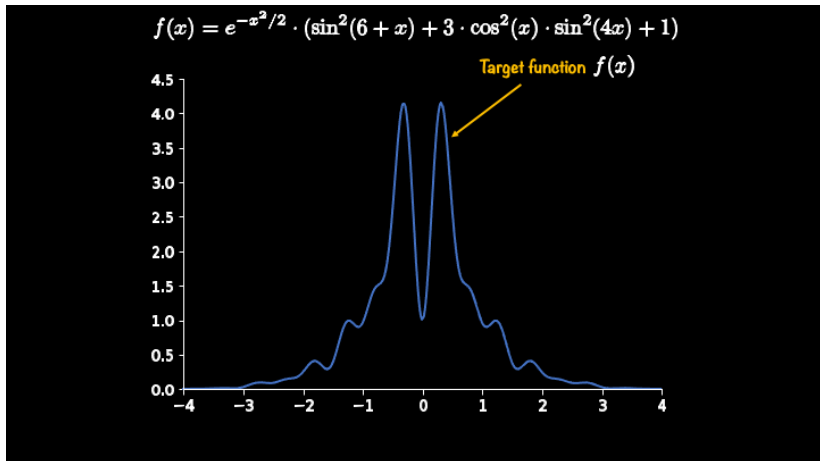


Figure 2: Target Function

2)

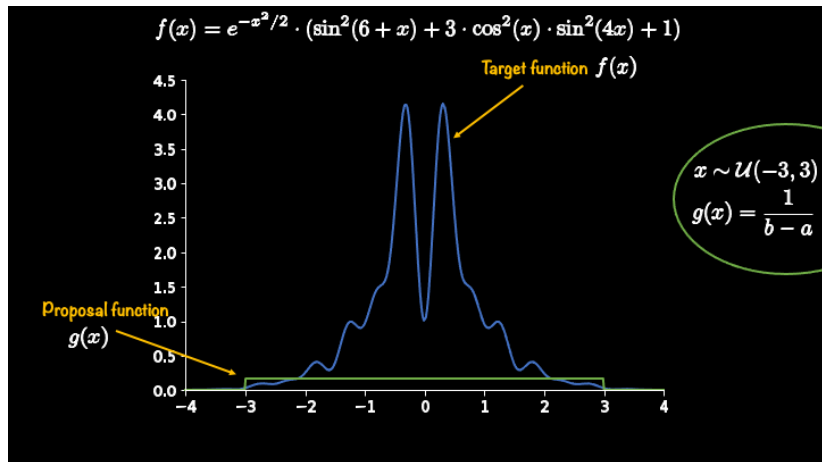


Figure 3: Proposal Function

3)

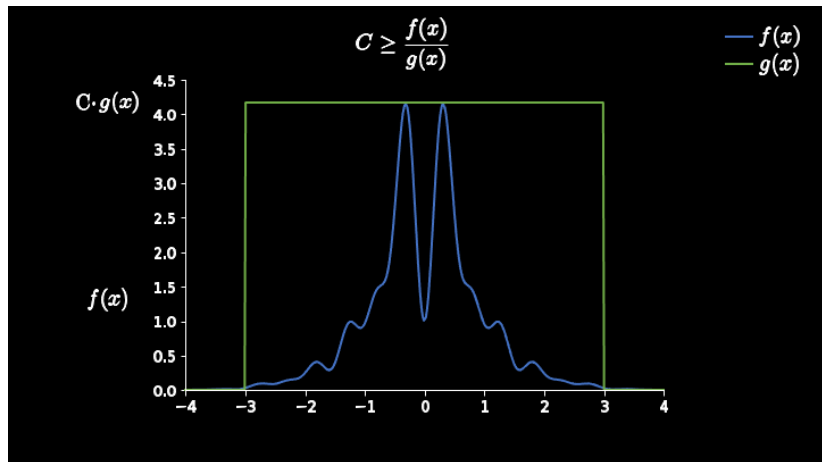
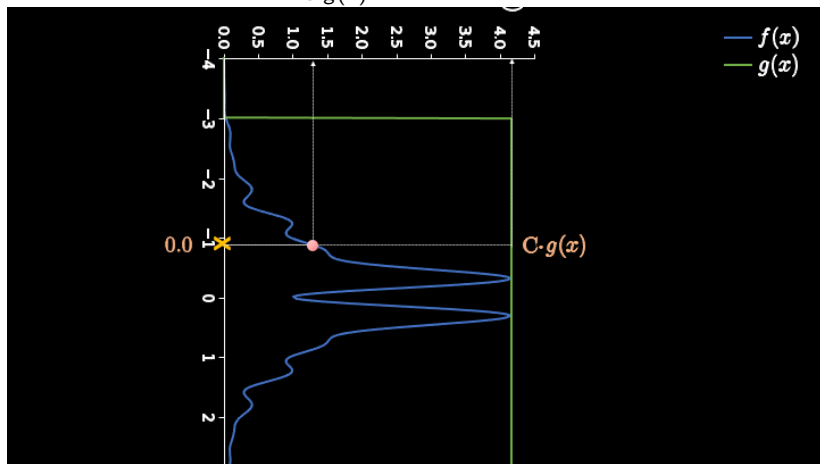


Figure 4: Constant

4)

Accept with probability $\frac{f(x)}{C \cdot g(x)}$



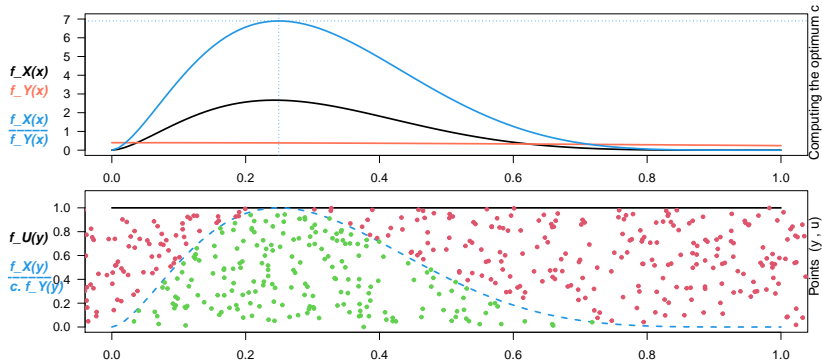
Example

```
blue is ratio
library(AR)

simulation = AR.Sim( n = 200,
  f_X = function(y){dbeta(y,2.7,6.3)},
  Y.dist = "norm", Y.dist.par = c(0,1),
  Rej.Num = TRUE,
  Rej.Rate = TRUE,
  Acc.Rate = FALSE
)
```

```
## Optimal c = 6.898
```

Graphical Presentation to Acceptance-Rejection Method



Output

n

The number/length of data which must be generated/simulated from $(f_X)(\text{TARGET})$ density.

Optimal $c = 6.898$

The numbers of Rejections = 1295

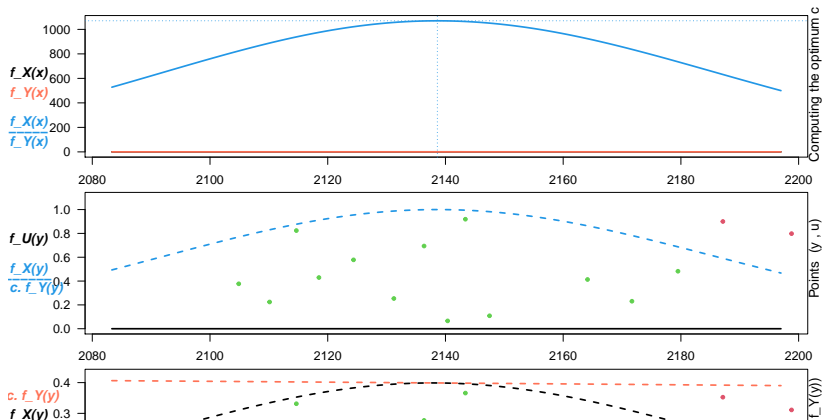
Ratio of Rejections = 0.866

Example_DNA

```
start.time<-Sys.time()
lr_rs = AR.Sim( n = 20,
  f_X = function(y){(posterior(y,as.numeric(gammaPrior_Cont[[sam]]$kij[taxa]), al_c,be_c))},
  Y.dist = "gamma", Y.dist.par = c((k/mean(lc))*al_c,be_c), xlim=c(k-2*mean(lc),k),
  Rej.Num = TRUE,
  Rej.Rate = TRUE,
  Acc.Rate = FALSE
)
```

Optimal c = 1069.962

Graphical Presentation to Acceptance-Rejection Method



Limitation

- ▶ Selecting the appropriate proposal function & finding its scaling constant
- ▶ Requires that the PDF of the target function is known
- ▶ Generally inefficient especially in higher dimensions

2.5) Adaptive Rejection Sampling

Define our proposal distribution in log space

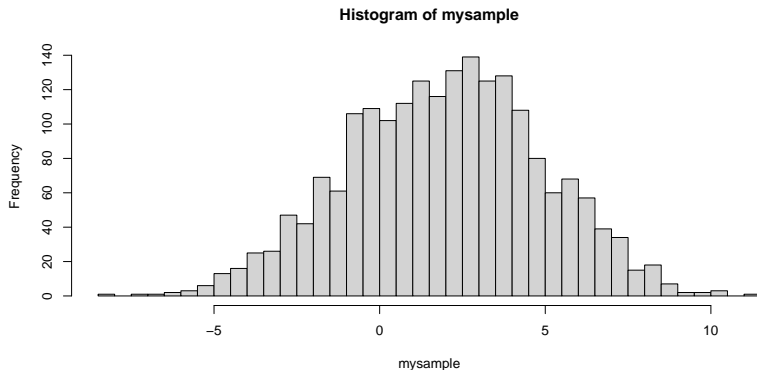
1

¹Gilks, W. R., & Wild, P. (1992). Adaptive rejection sampling for Gibbs sampling. *Journal of the Royal Statistical Society: Series C (Applied Statistics)*, 41(2), 337-348.

Example

sample 2000 values from the normal distribution $N(2,3)$

```
library("ars")  
  
f<-function(x,mu=0,sigma=1){-1/(2*sigma^2)*(x-mu)^2}  
fprima<-function(x,mu=0,sigma=1){-1/sigma^2*(x-mu)}  
mysample<-ars(2000,f,fprima,mu=2,sigma=3)  
hist(mysample, breaks=30)
```

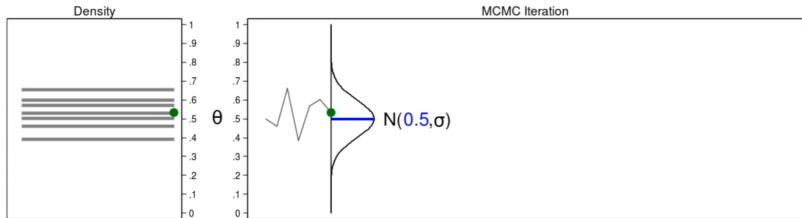


Monte Carlo

Relies on repeated random sampling to obtain numerical result

Ex) $\theta_t \sim \text{Normal}(0.5, \sigma)$

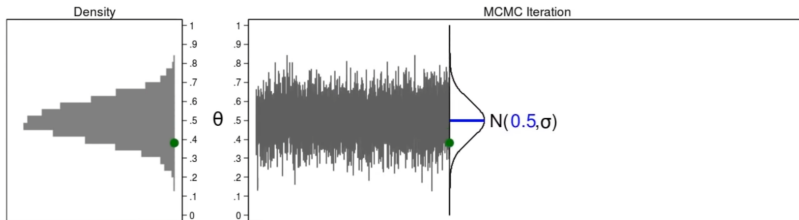
Monte Carlo Trace Plot



Draw $\theta_t \sim \text{Normal}(0.5, \sigma) = 0.534$

Figure 5: Trace Plot

50000 iterations



Draw $\theta_t \sim \text{Normal}(0.5, \sigma) = 0.380$

Figure 6: 50000

Markov property

Given the present, the future does not depend on the past.

$$P(X_n = x_n \mid X_{n-1} = x_{n-1}, \dots, X_0 = x_0) = P(X_n = x_n \mid X_{n-1} = x_{n-1}).$$

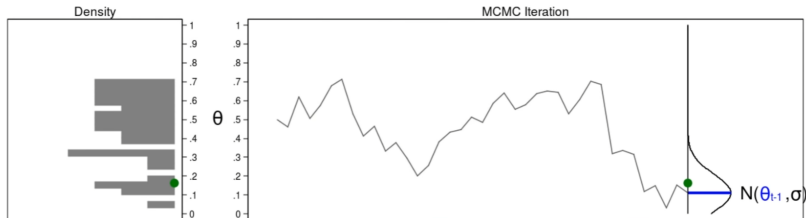
Figure 7: markovproperty

Example

Ex) $\theta_t \sim \text{Normal}(\theta_{t-1}, \sigma)$

Depends on the previous number on a sequence

Trace Plot

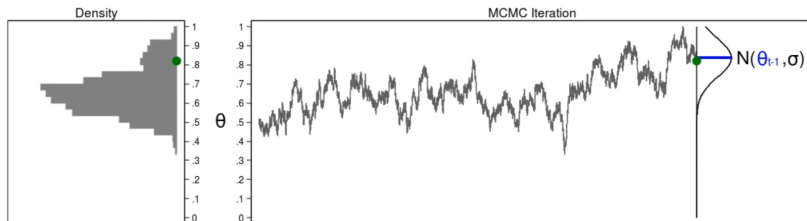


Draw $\theta_t \sim \text{Normal}(\theta_{t-1}, \sigma)$

$\text{Normal}(0.111, \sigma) = 0.164$

Figure 8: Trace Plot

50000 entries



Draw $\theta_t \sim \text{Normal}(\theta_{t-1}, \sigma)$

$\text{Normal}(0.836, \sigma) = 0.820$

3) Metropolis Hasting

The Metropolis–Hastings algorithm can draw samples from any probability distribution $f(x)$, provided that we know a function $q(x)$ proportional to the density of f and the values of $q(x)$ can be calculated. The requirement that $q(x)$ must only be proportional to the density

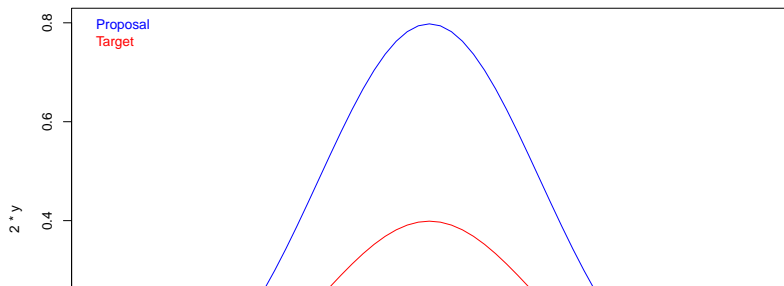
1. Compute $\rho(x, y) = \min \left\{ \frac{f(y)}{f(x)} \times \frac{q(x|y)}{q(y|x)}, 1 \right\}$
2. If $\rho(x, y) < 1$ then generate $U_t \sim \text{Uniform}(0, 1)$.
3. Set $X^{(t+1)} = \begin{cases} y & \text{if } \rho(x, y) = 1 \text{ or } U_t < \rho(x, y) \\ x & \text{if } U_t \geq \rho(x, Y_t) \end{cases}$

Figure 9: mh

Intuition

$$\alpha = \min\left\{\frac{f(b)}{f(a)}, 1\right\}$$

```
x= seq(-3,3, by=0.1)
y=dnorm(x)
plot(x,2*y, type="l", col="blue")
lines(x,y, col="red")
legend("topleft", legend = c("Proposal", "Target"),
text.col = c('blue','red'), bty = "n")
```



Limitation

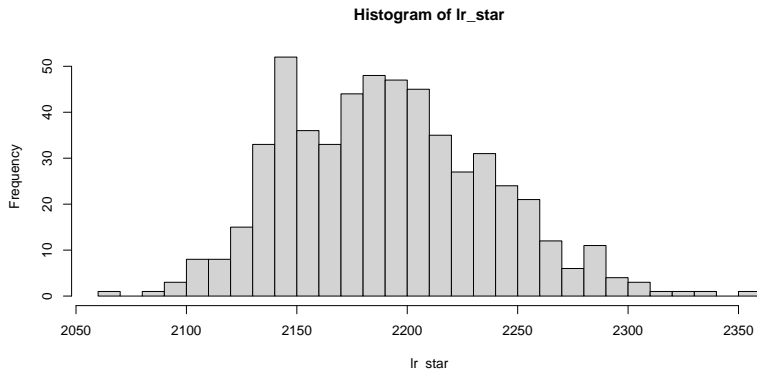
- ▶ Dependence on starting value
 - ▶ Burn-in period
- ▶ Autocorrelation due to the Markov Chain properties

Example

```
start.time<-Sys.time()
chain <- MH_MCMC(itera = 600,
  k = as.numeric(gammaPrior_Cont[[sam]]$kij[taxa]),
  al_c = gammaPrior_Cont[[sam]]$alpha_ij_c[taxa],
  be_c = gammaPrior_Cont[[sam]]$beta_ij_c[taxa],
  startvalue_lamda_r = 0)

end.time<-Sys.time()
time.taken_mh <- end.time-start.time

lr_star <- chain[50:length(chain)]
hist(lr_star, breaks=30)
```



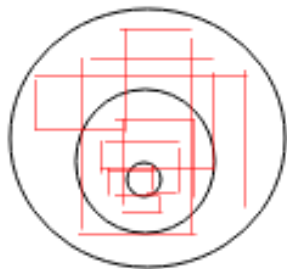
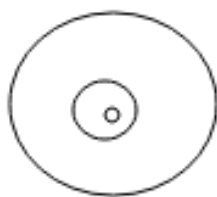
```
initialize  $Y^0, X^0$   
for  $j = 1, 2, 3, \dots$  do  
    sample  $X^j \sim p(X|Y^{j-1})$   
    sample  $Y^j \sim p(Y|X^j)$   
end for
```

Figure 10: Gibbs

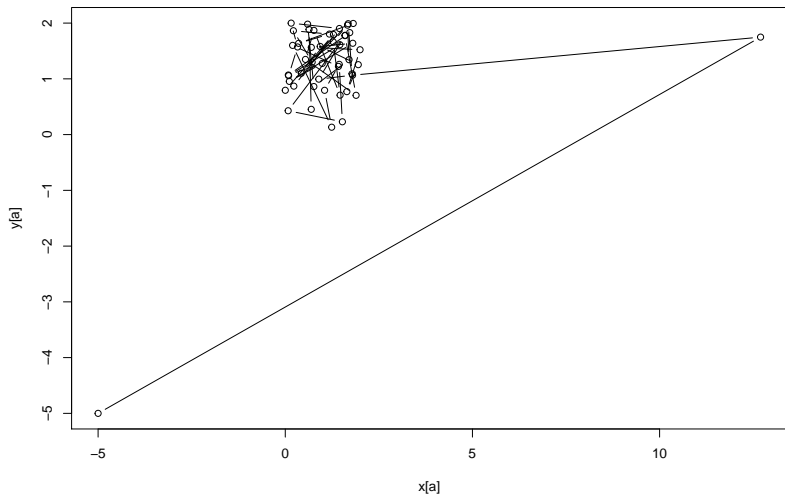
Limitation

	0	1
1	0	$\frac{1}{2}$
0	$\frac{1}{2}$	0

Limitation



Example



Example

```
#library(plotly)  
#plot_ly(x=den$x, y=den$y, z=den$z) %>% add_surface
```


Goodness of fit

```
goodness_of_fit <- function(lc,lr){  
  kij_c_star <- rpois(length(lc),mean(lc))  
  kij_r_star <- rpois(length(lc),mean(lr))  
  kij_star <- kij_r_star+kij_c_star  
  hist(kij_star,  
       main=paste("Goodness of fit for",deparse(substitute(lr))) , breaks =30, xlab="counts")  
  abline(v=as.numeric(gammaPrior_Cont[[sam]]$kij[taxa]))  
}
```

Grid Approximation

```
#goodness_of_fit(lc,post_lr)
```

```
# actual observed counts  
k
```

```
## [1] 2197
```

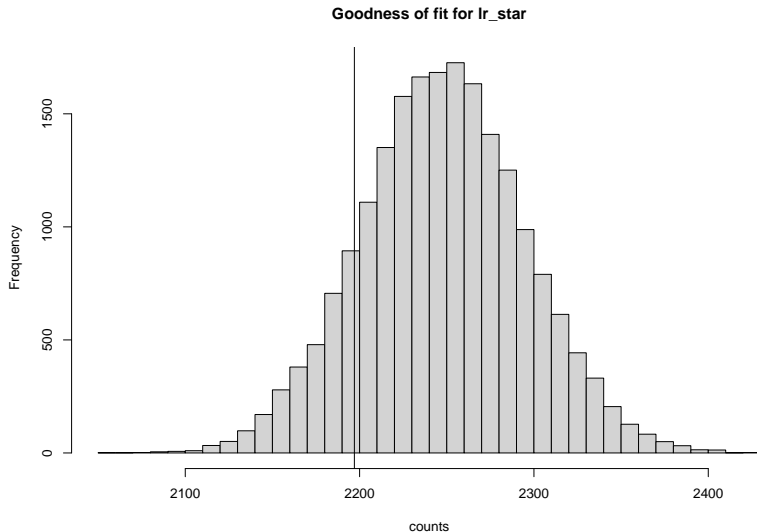
```
# point estimation
```

```
mean(lc) + lr_ga
```

```
## [1] 2192.639
```

Metropolis-Hasting

```
goodness_of_fit(lc,lr_star)
```



Rejection Sampling

```
goodness_of_fit(lc,lr_rs)
```

