Importance Sampling

```
n=1000
b = 1/2
inv\_cdf\_mh\_p < -function(x)\{
 if(x >= 0.5){
   w = -log(2-2*x)
 if(x < 0.5){
   w = log(2*x)
 x1_sample_list<-c(x1_sample_list,-w)
x1_sample<-c()
for (i in 1:n) \{
 x1_sample<-c(x1_sample,inv_cdf_mh_p(runif(1)))
x2\_sample < -rnorm(n, mean = 0, sd = b)
f\_muti < -function(x1,x2)\{
  exponentional <- (1/2*exp(-abs(x1)))
  normal_dist <- (1/sqrt(2*pi*b**2)*exp(-x2^2/(2*b**2)))
  normal_dist*exponentional
h\_muti < -function(x1,x2)\{
 A =as.numeric(x1^2+x2>=0)
  cos(x1+x2)*exp(-(abs(x1)+2*x2^2))*A
mean_a<-mean(h_muti(x1_sample,x2_sample)/f_muti(x1_sample,x2_sample))</pre>
> mean_a
[1] 0.6902987
```

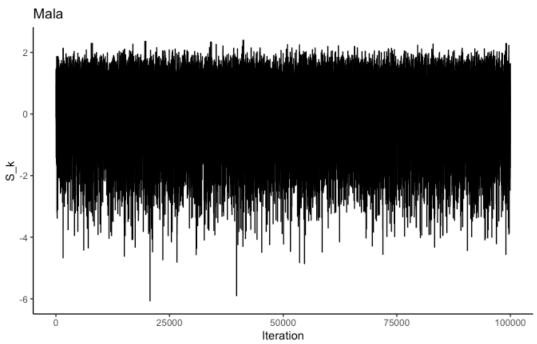
MCMC - Random Walker

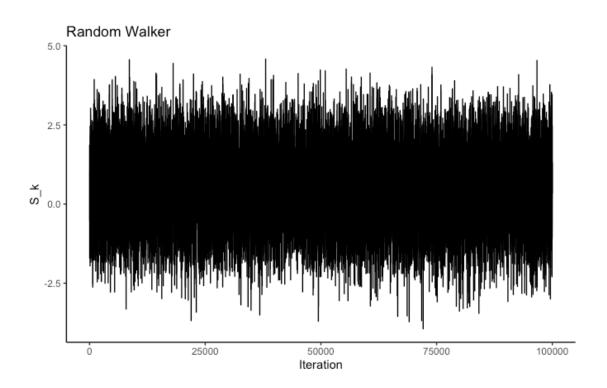
```
aux_p<-function(x1, x2){</pre>
  part1<- (-abs(x1))
  part2<-(-2*x2^2)
  exp(part1+part2)
q1_mcmc<-function(x1_init,x2_init,step_size, iteration){
  x1<-x1_init
  x2<-x2_init
  x1_all < -c(x1)
  x2_all < -c(x2)
  p_1<-aux_p(x1,x2)
  for (i in 1:iteration) {
    y1<-rnorm(1,x1,step_size)
    y2<-rnorm(1,x2,step_size)
    p_2<-aux_p(y1,y2)
    accept_prob < -min(1, p_2/p_1)
    u = runif(1)
    if( accept_prob > u){
      x1<- y1
      x2<- y2
      p_1<-p_2
      x1_all <-c(x1_all, x1)
      x2_all <-c(x2_all, x2)
    }
    if( accept_prob <= u){</pre>
      x1<- x1
      x2<- x2
      p_1<-p_1
      x1_all < -c(x1_all, x1)
      x2_all <-c(x2_all, x2)
    }
  }
  return(cbind(x1_all,x2_all))
expect<-function(x1,x2){
  cos(x1+x2)*as.numeric(x1^2+x2>=0)
x_input1<-1
x_input2<-2
n=1000
a<-q1_mcmc(x_input1,x_input2,1,n)</pre>
z= 2 * sqrt(pi/2)
mean(expect(a[,'x1_all'],a[,'x2_all']))*z
> mean(expect(a[,'x1_all'],a[,'x2_all']))*z
[1] 0.7169394
```

```
q2_target<-function(x,s1,s2,g1){
  part1 < -exp(-x^2/(2*s1^2))
  part2 < -exp(-(x-g)^2/(2*s2^2))
  part1 + part2
gradient_p<-function(x,s1,s2,g1){
  # outter dervative for log
  part1 \leftarrow 1/q2\_target(x,s1,s2,g1)
  # inner dervative
  part2_1 \leftarrow (-x/s1^2)*exp(-x^2/(2*s1^2))
  part2_2 \leftarrow exp(-(x-g1)^2/(2*s2^2))*(-(x-g1)/s2^2)
  part2<- part2_1 + part2_2
  #together
  part1*part2
propose_a_given_b<- function(a,b,step_size,s1,s2,g1){</pre>
  part1<-1/sqrt(2*pi*2*step_size^2)
  inner\_part2\_1 <- \ (a-(b+step\_size^2*gradient\_p(b,s1,s2,g1)))^2
  inner_part2_2<- 2*2*step_size^2
  part2<- exp(-inner_part2_1/inner_part2_2)</pre>
  part1*part2
q2_mala<-function(x1_init,s1,s2,g1,step_size,iteration){
 x1_all_2<-c(x1_init)
 x1<-x1_init
 p_1<-q2_target(x1,s1,s2,g1)
 accept_mala = 0
 for (i in 1:iteration) {
   y1<-rnorm(1,x1+gradient_p(x1,s1,s2,g1)*step_size^2, 2*step_size^2)
   p_2<-q2_target(y1,s1,s2,g1)
   accept\_prob < -min(1,(p\_2*propose\_a\_given\_b(x1,y1,step\_size,s1,s2,g1))/(p\_1*propose\_a\_given\_b(y1,x1,step\_size,s1,s2,g1)))
   u<-runif(1)
   if(accept\_prob >= u){}
     accept_mala = accept_mala+1
     print(accept_mala)
     x1 <- y1
     p_1 <-p_2
     x1_all_2<-c(x1_all_2, x1)
   if( accept_prob < u){</pre>
     x1 <- x1
     p_1<-p_1
     x1_all_2<-c(x1_all_2, x1)
 }
 return(x1_all_2)
```

```
q2_random_walker<-function(x1_init,s1,s2,g1,step_size,iteration){
  x1_all_2<-c(x1_init)
  x1<-x1_init
  p_1<-q2_target(x1,s1,s2,g1)
  for (i in 1:iteration) {
    y1<-rnorm(1,x1, step_size)</pre>
    p_2<-q2_target(y1,s1,s2,g1)
    accept\_prob < -min(1,(p_2)/(p_1))
    u<-runif(1)
    if( accept_prob >= u){
      accept_random_walker = accept_random_walker+1
      print(accept_random_walker)
      x1<- y1
      p_1<-p_2
      x1_all_2<-c(x1_all_2, x1)
    if( accept_prob < u){</pre>
      x1<- x1
      p_1<-p_1
      x1_all_2<-c(x1_all_2, x1)
    }}
  return(x1_all_2)}
Functions to graphs
 b<-q2_mala(init_point,s1,s2,g,step_size, n2)
 c<-q2_random_walker(init_point,s1,s2,g,step_size, n2)</pre>
 b<-as.data.frame(b)
 c<-as.data.frame(c)
library(ggplot2)
ggplot(data=b, aes(x=1:length(b), y=b, group=1)) +
  geom_line()+ theme_classic()+
  labs(title="Mala",x="Iteration", y = "S_k")
hist(b$b,main = "Mala",breaks = 50)
ggplot(data=c, aes(x=1:length(c), y=c, group=1)) +
  geom_line()+ theme_classic()+
  labs(title="Random Walker",x="Iteration", y = "S_k")
hist(c$c,main = "Random Walker",breaks = 50)
Setting 1
 s1<-1
 s2<-1
 g<-1
 step_size<-1
n2<-10^5
init_point<-1
b<-q2_mala(init_point,s1,s2,g,step_size, n2)
 # accpted points: 33452
 c<-q2_random_walker(init_point,s1,s2,g,step_size, n2)</pre>
 # accpted points: 73306
```

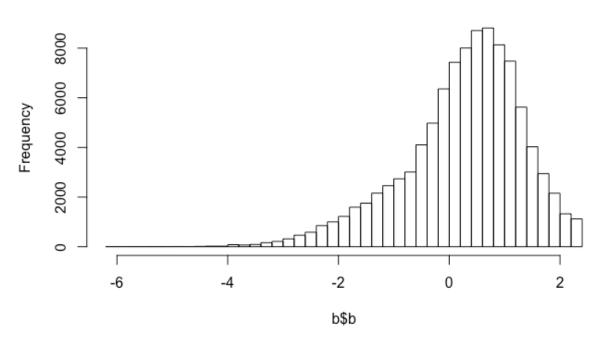
Trace plot



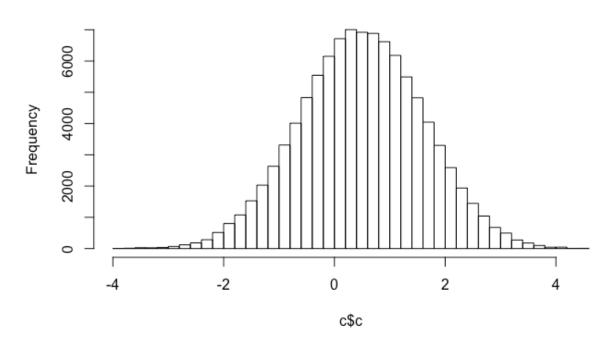


Histogram





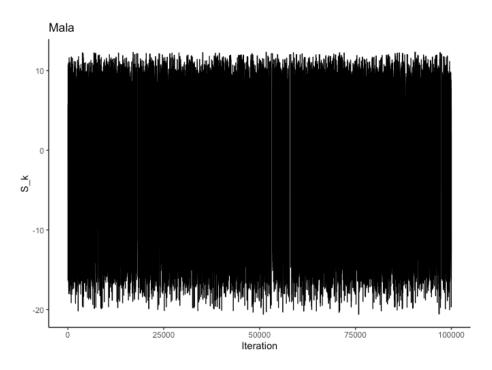
Random Walker

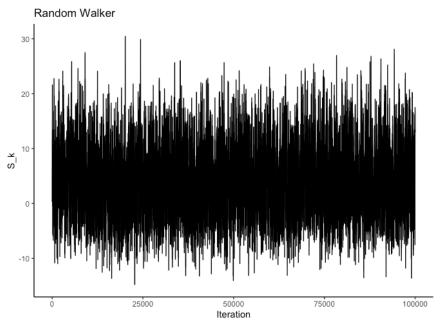


Setting 2

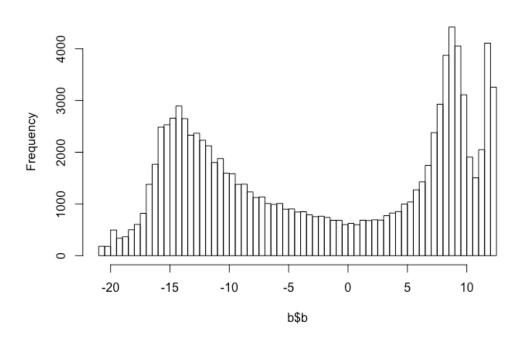
```
s1<-4
s2<-6
g<-7
step_size<-2
n2<-10^5
init_point<-0.3
b<-q2_mala(init_point,s1,s2,g,step_size, n2)
# accpted points: 41342
c<-q2_random_walker(init_point,s1,s2,g,step_size, n2)
# accpted points: 89875</pre>
```

Trace plot

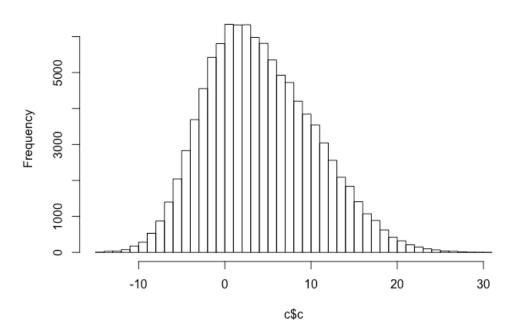








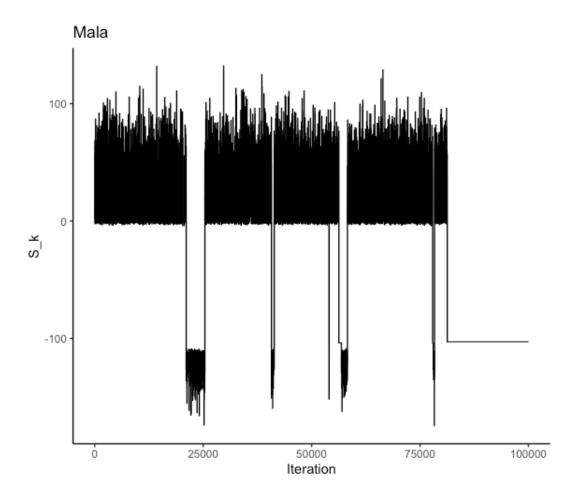
Random Walker

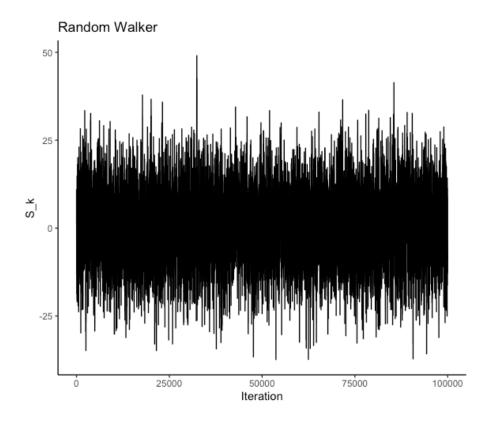


Setting 3

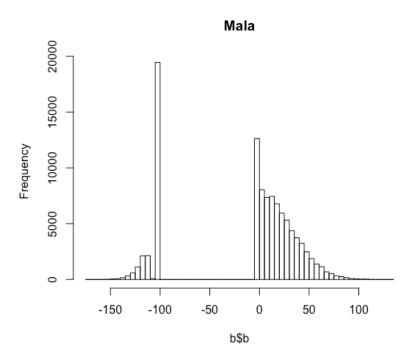
```
s1<-10
s2<-1
g<-7
step_size<-4
n2<-10^5
init_point<-2
b<-q2_mala(init_point,s1,s2,g,step_size, n2)
# accpted points: 31761
c<-q2_random_walker(init_point,s1,s2,g,step_size, n2)
# accpted points: 83864</pre>
```

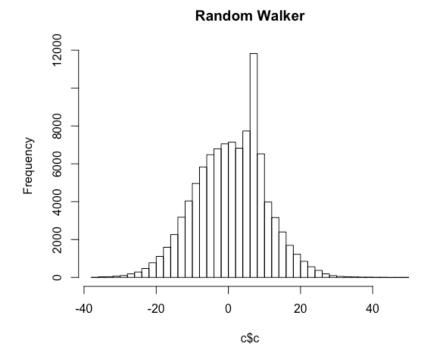
Trace plot





Histogram





How should your histograms look like?

There should be two peaks. S1 and S2 are similar to variance and g is how separate those two peaks should be.

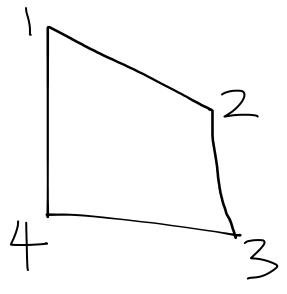
Which algorithms seem to be mixing faster?

It seems like Mala mixing faster because the acceptance rate is lower.

Part 1

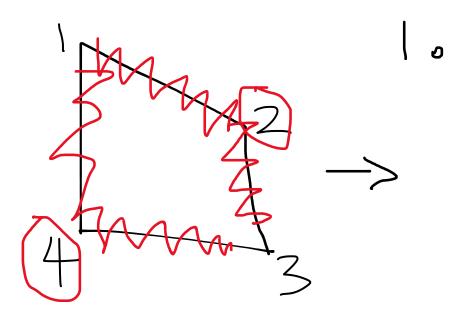
```
total = c()
for (x1 in 1:2) {
    for (x2 in 1:2) {
        for (x3 in 1:2) {
            for (x4 in 1:2) {
                t<-((x1**x2)*(x2**x3)*(x3**x4)*(x4**x1))
                total<-c(total,t)
            }
        }
    }
}
sum(total)
> sum(total)
[1] 433
Z = 433; The purpose of Z is for 1/Z*PMF = 1
```

Part 2



Part 3

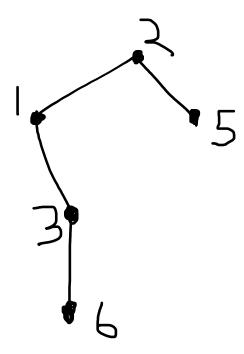
Yes, they are independent as we can use global Markov property to deduce this result that X2 and X4 forms a separating set which allows conditional independence among X1 and X3. Please refers to the picture below for visual.



Part 4

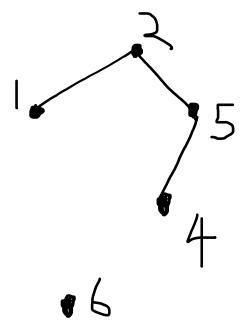
Part A

No, X5 and X6 is not independent given X4 as X4 is not a separating set for X5 and X6.



Part B

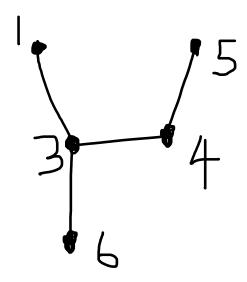
Yes, X5 and X6 is independent given X3 according to the Global Markov property. As X3 is a separating set for X5 and X6.



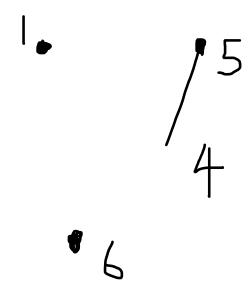
Part C

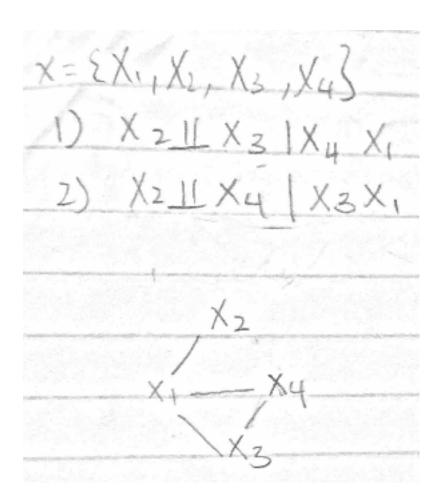
No, we cannot make statement regarding conditionally dependence with the Global Markov property.

,



Part D
Yes, as X2 and X3 is a separating set for X1 and X4. We can use Global Markov property.





Appendix

Calculation for gradient log(p)

