

# MATH 598: TOPICS IN STATISTICS

## THE LANGEVIN ALGORITHM

Denote the target pdf  $\pi(x)$ . Consider the Langevin SDE, and the diffusion process  $\{X_t, t > 0\}$

$$dX_t = \frac{1}{2}S(X_t)dt + dW_t$$

where  $\{W_t\}$  is Brownian motion, and

$$S(x) = \frac{d}{dx} \log \pi(x) = \nabla \log \pi(x).$$

It can be shown that the invariant measure of this continuous time process is  $\pi(x)$ . A standard scheme for simulating this diffusion is given by an Euler approximation based on the time-discretization

$$\{X_{k\delta}, k = 1, 2, \dots\}$$

for some time-step  $\delta > 0$  where  $X_0 = x_0$ , and then for  $k = 0, 1, 2, \dots$

$$X_{k+1} = X_k + \frac{\delta}{2}S(X_k) + \sqrt{\delta}Z_k$$

where  $\{Z_k, k = 1, 2, \dots\}$  are an iid  $Normal_d(0, \mathbf{I}_d)$  sequence. The approximation induced by time-discretization does not quite preserve  $\pi(x)$  as the invariant distribution. This can be easily corrected using a Metropolis-Hastings step to yield the Metropolis-adjusted Langevin algorithm where the proposal

$$X_{k+1}^* \sim Normal_d(X_k + \delta S(X_k)/2, \delta \mathbf{I}_d)$$

is accepted with probability

$$\min \left\{ 1, \frac{\pi(X_{k+1}^*) q(X_k, X_{k+1}^*)}{\pi(X_k) q(X_{k+1}^*, X_k)} \right\}$$

where

$$q(\mathbf{x}, \mathbf{y}) \propto \exp \left\{ -\frac{1}{2\delta} (\mathbf{y} - \mathbf{x} - \delta S(\mathbf{x})/2)^\top (\mathbf{y} - \mathbf{x} - \delta S(\mathbf{x})/2) \right\}$$

**Example:** Suppose  $Y \equiv \text{Gamma}(\alpha, 1)$ , and consider the pdf of  $X = \log Y$ . Then for  $x \in \mathbb{R}$

$$f_X(x) = e^x f_Y(e^x) = \frac{1}{\Gamma(\alpha)} \exp\{\alpha x - e^x\}.$$

We set  $\pi(x) \equiv f_X(x)$  so that

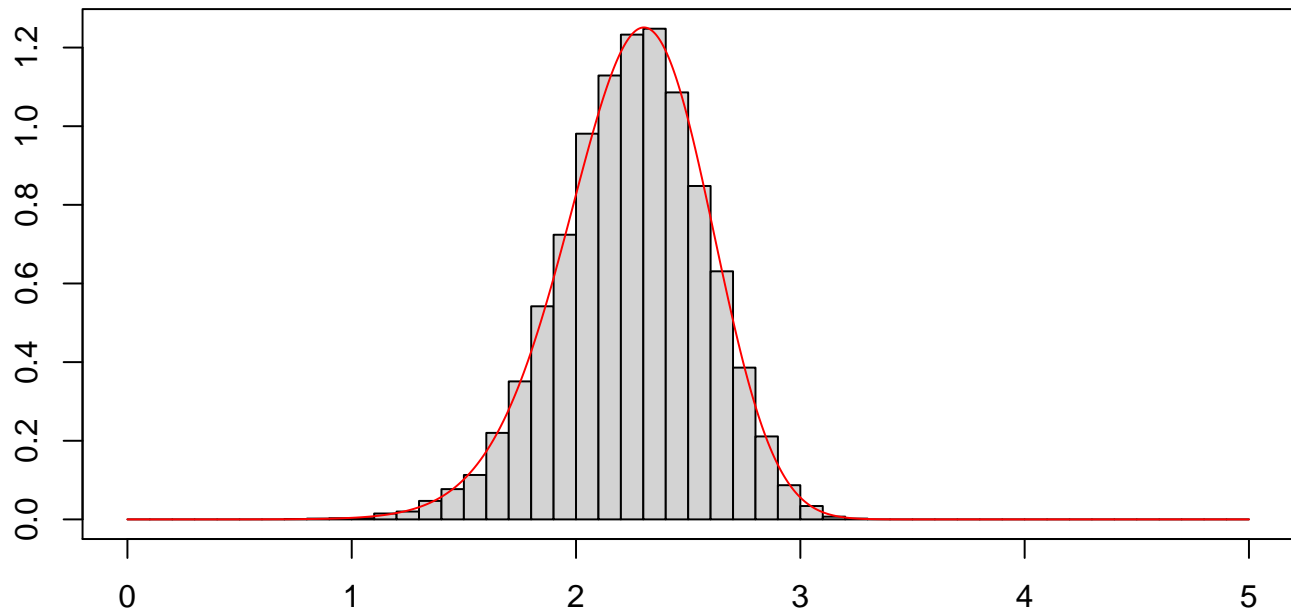
$$\log \pi(x) = \alpha x - e^x - \log \Gamma(\alpha)$$

$$S(x) = \frac{d}{dx} \log \pi(x) = \alpha - e^x$$

It is straightforward to simulate this pdf directly

```
set.seed(23984)
a1<-10
N<-10001
Y<-rgamma(N, a1, 1)
X<-log(Y)
par(mar=c(3,3,3,0))
hist(X, freq=FALSE, breaks=seq(0,5, by=0.1), main='Direct sampling');box()
xvec<-seq(0,5, by=0.001)
yvec<-exp(a1*xvec-exp(xvec))/gamma(a1)
lines(xvec, yvec, col='red')
```

## Direct sampling



We may also simulate the Langevin dynamics directly using the Euler approximation. We use  $N = 10000$  steps, with a time step of  $\delta = 0.01$ , taking  $x_0 = 2$ . We can compute the effective sample size for the sampling approach.

```
del<-0.01
x<-rep(0,N)
x[1]<-2
Z<-rnorm(N)
for(k in 1:N){
  x[k+1] <- x[k] + del*(a1 - exp(x[k]))/2 + sqrt(del)*Z[k]
}
library(coda)
ESS<-effectiveSize(x)
x.1<-x
```

The low effective sample size of 262.942 indicates a high degree of autocorrelation. The effective sample size can be increased by increasing  $\delta$  to 0.1.

```
del<-0.1
x<-rep(0,N)
x[1]<-2
Z<-rnorm(N)
for(k in 1:N){
  x[k+1] <- x[k] + del*(a1 - exp(x[k]))/2 + sqrt(del)*Z[k]
}
(ESS<-effectiveSize(x))

+ var1
+ 2988.588

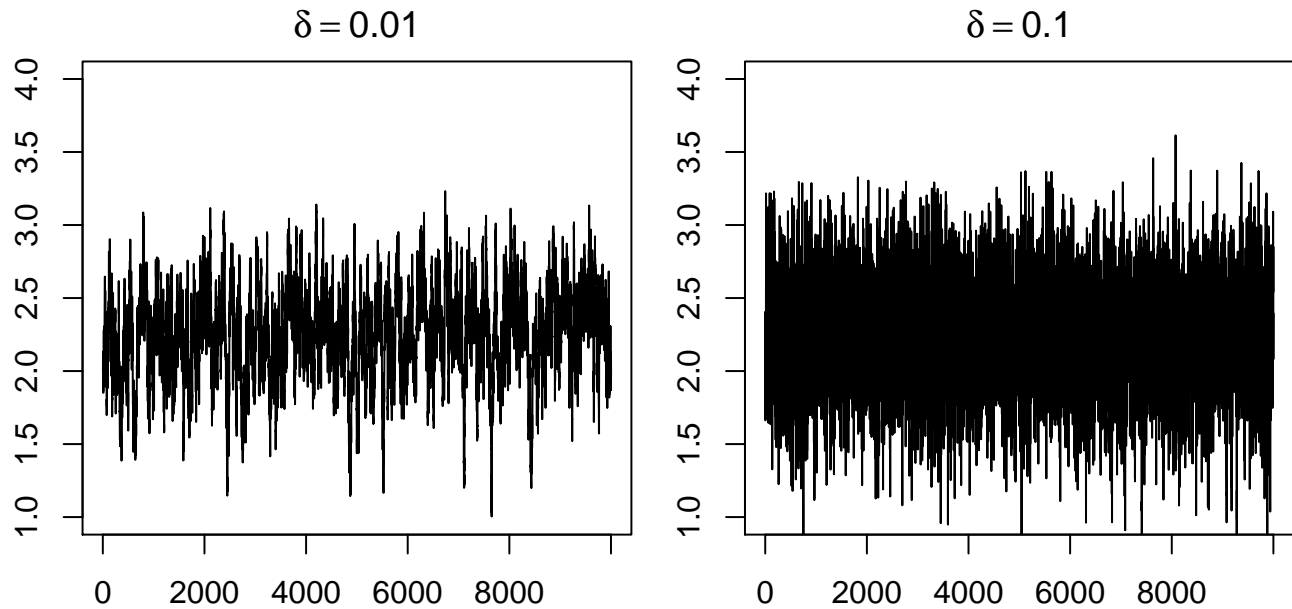
x.2<-x
```

The effective sample size is now 2988.588 so performance seems to be better. We now see how the two schemes with different  $\delta$  values compare.

```

par(mar=c(3,3,2,0),mfrow=c(1,2))
plot(x.1,type='l',main=expression(delta==0.01),ylim=range(1,4))
plot(x.2,type='l',main=expression(delta==0.1),ylim=range(1,4))

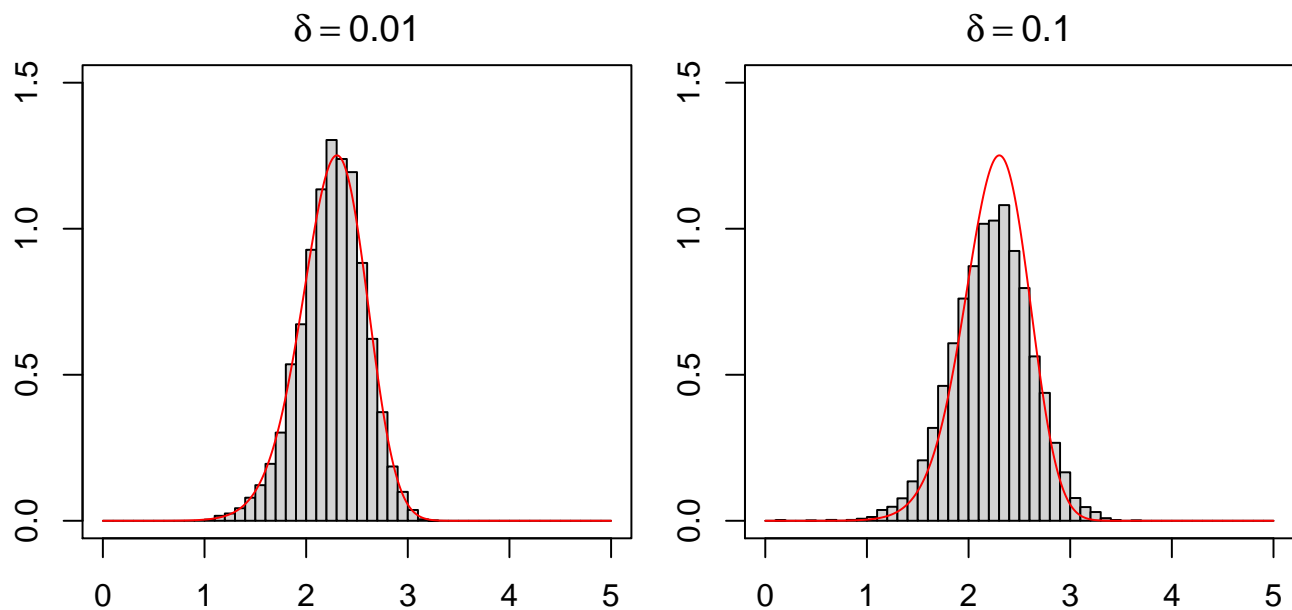
```



```

hist(x.1,freq=FALSE,breaks=seq(0,5,by=0.1),main=expression(delta==0.01),ylim=range(0,1.5));box()
lines(xvec,yvec,col='red')
hist(x.2,freq=FALSE,breaks=seq(0,5,by=0.1),main=expression(delta==0.1),ylim=range(0,1.5));box()
lines(xvec,yvec,col='red')

```



The approximation is poorer when  $\delta = 0.1$ .  
 We now implement the MALA enhancement:

```

log.pi<-function(xv,av){
  return(av*xv - exp(xv))
}
dlog.pi<-function(xv,av){
  return(av- exp(xv))
}
q.func<-function(xv,yv,av,dv){
  m<-xv+0.5*dv*dlog.pi(xv,av)
  return(dnorm(yv,m,sqrt(dv),log=T))
}

del<-0.01
x<-rep(0,N)
x[1]<-2
Z<-rnorm(N)
irej<-0
for(k in 1:N){
  x[k+1] <- x[k] + del*(al - exp(x[k]))/2 + sqrt(del)*Z[k]
  acc<-log.pi(x[k+1],al)-log.pi(x[k],al) + q.func(x[k+1],x[k],al,del) - q.func(x[k],x[k+1],al,del)
  if(log(runif(1)) > acc){
    x[k+1]<-x[k] #Reject !
    irej<-irej+1
  }
}
(ESS<-effectiveSize(x))

+ var1
+ 255.998

x.3<-x
acc.rate.3<-(N-irej)/N

```

The low effective sample size of 255.998 again indicates a high degree of autocorrelation. The effective sample size can again be increased by increasing  $\delta$  to 0.1.

```

del<-0.1
x<-rep(0,N)
x[1]<-2
Z<-rnorm(N)
irej<-0
for(k in 1:N){
  x[k+1] <- x[k] + del*(al - exp(x[k]))/2 + sqrt(del)*Z[k]
  acc<-log.pi(x[k+1],al)-log.pi(x[k],al) + q.func(x[k+1],x[k],al,del) - q.func(x[k],x[k+1],al,del)
  if(log(runif(1)) > acc){
    x[k+1]<-x[k] #Reject !
    irej<-irej+1
  }
}
(ESS<-effectiveSize(x))

+ var1
+ 3223.515

x.4<-x
acc.rate.4<-(N-irej)/N
print(c(acc.rate.3,acc.rate.4))

+ [1] 0.9976002 0.9200080

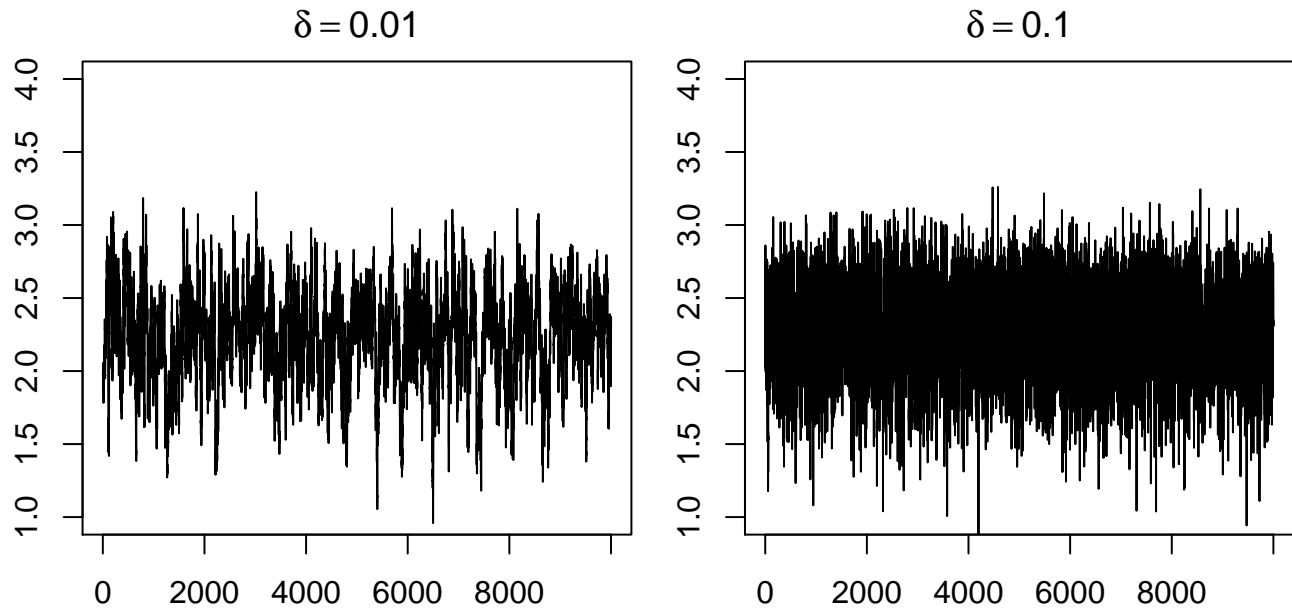
```

The effective sample size is now 3223.515. The acceptance rates for the Metropolis step are very high in both algorithms.

```

par(mar=c(3,3,2,0),mfrow=c(1,2))
plot(x.3,type='l',main=expression(delta==0.01),ylim=range(1,4))
plot(x.4,type='l',main=expression(delta==0.1),ylim=range(1,4))

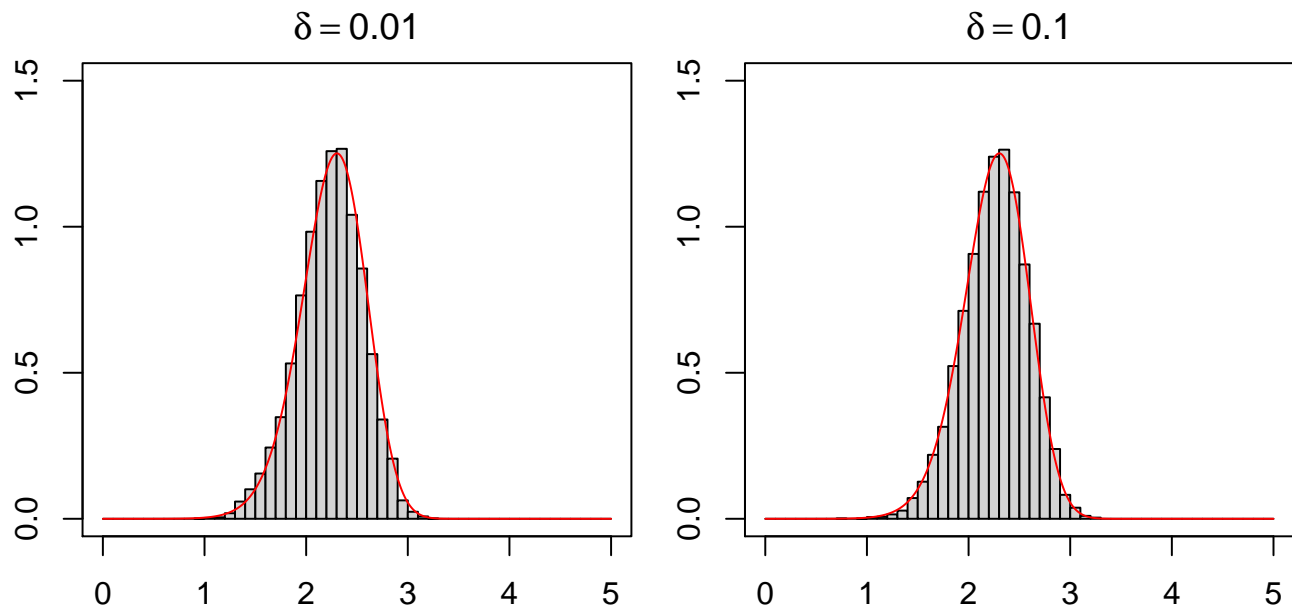
```



```

hist(x.3,freq=FALSE,breaks=seq(0,5,by=0.1),main=expression(delta==0.01),ylim=range(0,1.5));box()
lines(xvec,yvec,col='red')
hist(x.4,freq=FALSE,breaks=seq(0,5,by=0.1),main=expression(delta==0.1),ylim=range(0,1.5));box()
lines(xvec,yvec,col='red')

```



The sampling from the target is good in both cases, the inaccuracy for  $\delta = 0.1$  has been corrected.

We now assess results when  $\delta$  is increased to  $\delta = 0.5$  and  $\delta = 1.0$ .

```

del<-0.5
x<-rep(0,N)
x[1]<-2
Z<-rnorm(N)
irej<-0
for(k in 1:N){
  x[k+1] <- x[k] + del*(al - exp(x[k]))/2 + sqrt(del)*Z[k]
  acc<-log.pi(x[k+1],al)-log.pi(x[k],al) + q.func(x[k+1],x[k],al,del) - q.func(x[k],x[k+1],al,del)
  if(log(runif(1)) > acc){
    x[k+1]<-x[k] #Reject !
    irej<-irej+1
  }
}
(ESS<-effectiveSize(x))

+      var1
+ 3864.617

x.5<-x
acc.rate.5<-(N-irej)/N

del<-1
x<-rep(0,N)
x[1]<-2
Z<-rnorm(N)
irej<-0
for(k in 1:N){
  x[k+1] <- x[k] + del*(al - exp(x[k]))/2 + sqrt(del)*Z[k]
  acc<-log.pi(x[k+1],al)-log.pi(x[k],al) + q.func(x[k+1],x[k],al,del) - q.func(x[k],x[k+1],al,del)
  if(log(runif(1)) > acc){
    x[k+1]<-x[k] #Reject !
    irej<-irej+1
  }
}
(ESS<-effectiveSize(x))

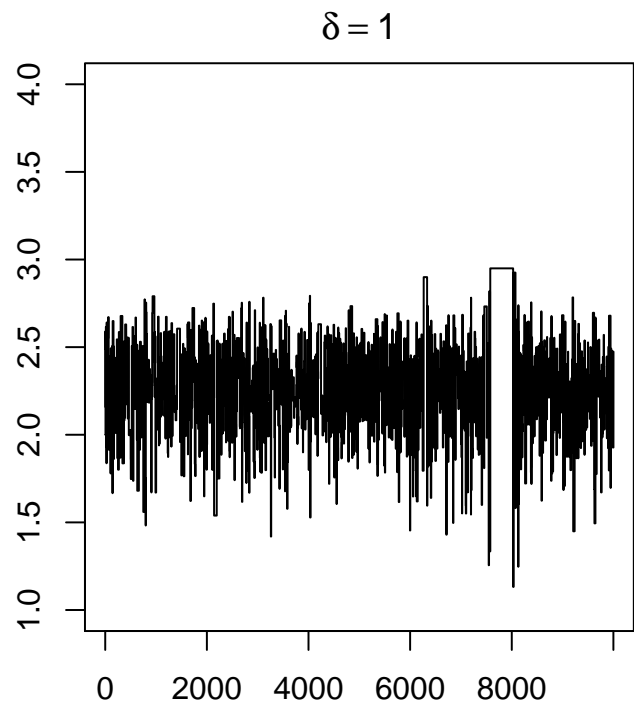
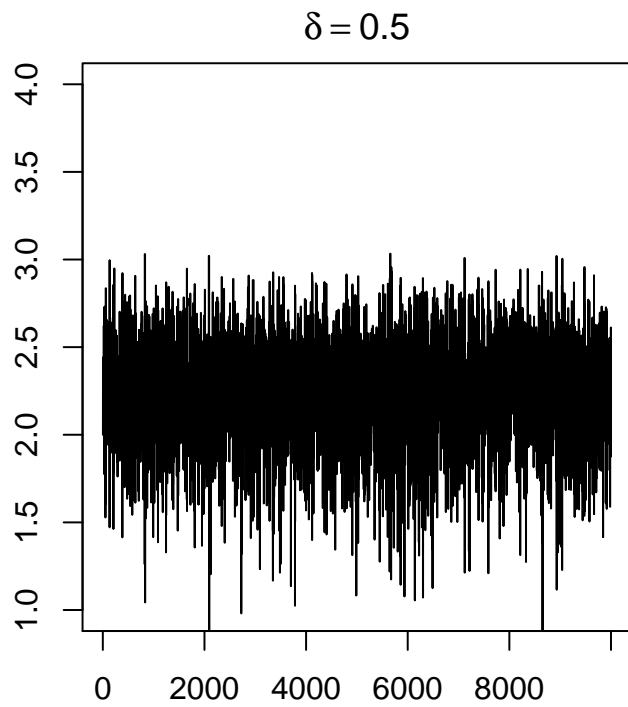
+      var1
+ 321.7487

x.6<-x
acc.rate.6<-(N-irej)/N
print(c(acc.rate.5,acc.rate.6))

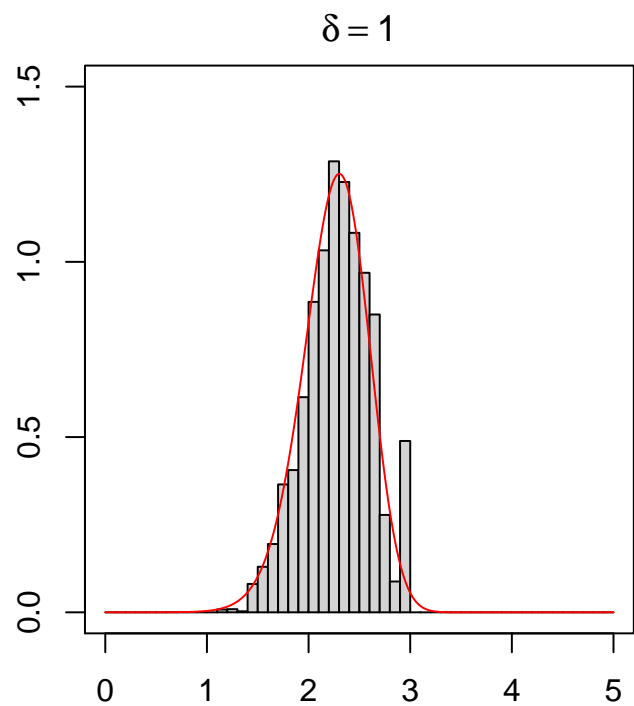
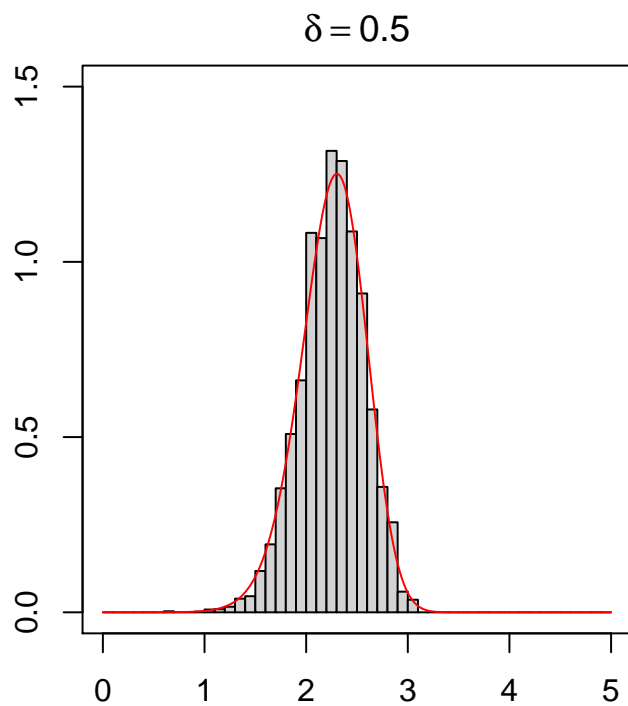
+ [1] 0.4166583 0.1618838

par(mar=c(3,3,2,0),mfrow=c(1,2))
plot(x.5,type='l',main=expression(delta==0.5),ylim=range(1,4))
plot(x.6,type='l',main=expression(delta==1.0),ylim=range(1,4))

```



```
hist(x.5,freq=FALSE,breaks=seq(0,5,by=0.1),main=expression(delta==0.5),ylim=range(0,1.5));box()
lines(xvec,yvec,col='red')
hist(x.6,freq=FALSE,breaks=seq(0,5,by=0.1),main=expression(delta==1.0),ylim=range(0,1.5));box()
lines(xvec,yvec,col='red')
```



```
var(X)
+ [1] 0.1036018

apply(cbind(x.1,x.2,x.3,x.4,x.5,x.6),2,var)
+      x.1      x.2      x.3      x.4      x.5      x.6
+ 0.10031906 0.14562766 0.10238010 0.10187117 0.09821227 0.10574323
```

There is some theory to suggest that the optimal acceptance rate for the MALA algorithm is around 0.57. We examine the variation in the ESS and acceptance rate as  $\delta$  varies from 0.1 to 0.5.

```
N<-100001
del.vec<-acc.vec<-seq(0.1,0.5,by=0.01)
xd<-matrix(0,nrow=N+1,ncol=length(del.vec))
x<-rep(0,N)
for(j in 1:length(del.vec)){
  del<-del.vec[j]
  x[1]<-2
  Z<-rnorm(N)
  irej<-0
  for(k in 1:N){
    x[k+1] <- x[k] + del*(al - exp(x[k]))/2 + sqrt(del)*Z[k]
    acc<-log.pi(x[k+1],al)-log.pi(x[k],al) + q.func(x[k+1],x[k],al,del) - q.func(x[k],x[k+1],al,del)
    if(log(runif(1)) > acc){
      x[k+1]<-x[k] #Reject !
      irej<-irej+1
    }
  }
  xd[,j]<-x
  acc.vec[j]<-(N-irej)/N
}
ESSd<-effectiveSize(xd)
```

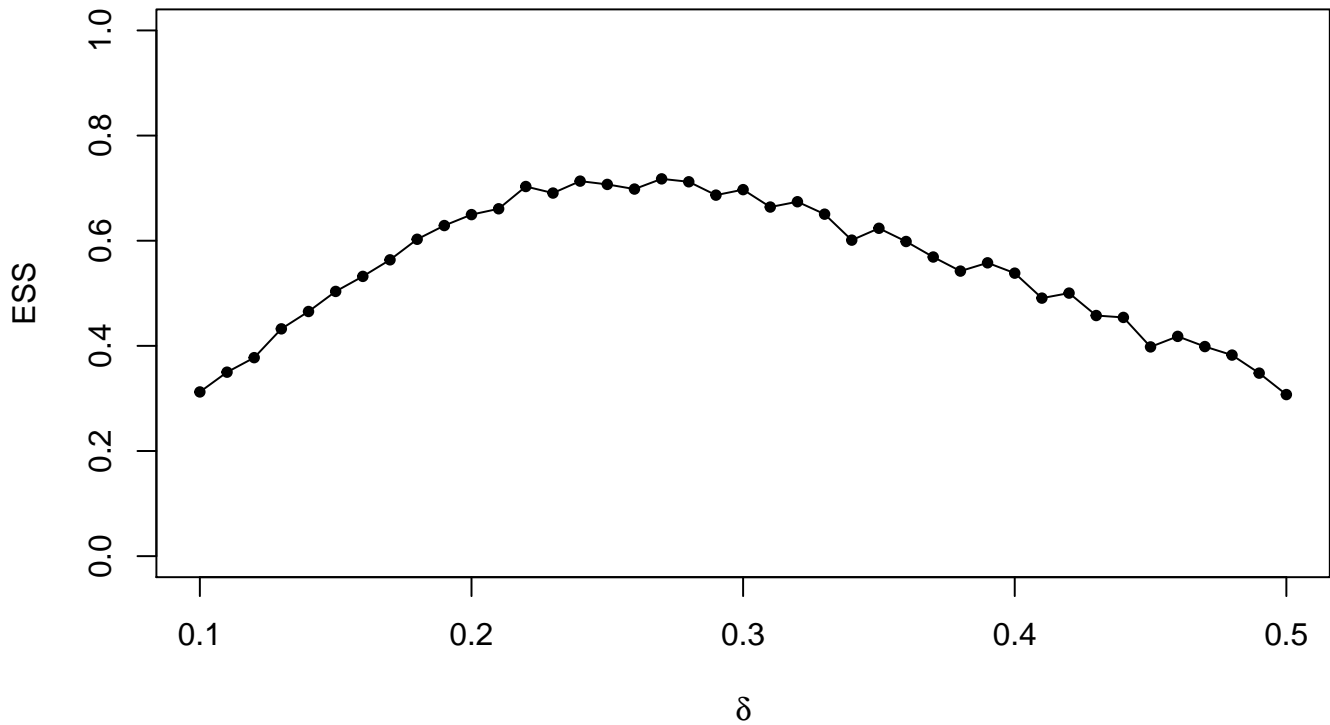
In the plots below, we compare

- Effective sample size vs  $\delta$ ;
- Effective sample size vs Metropolis acceptance probability.

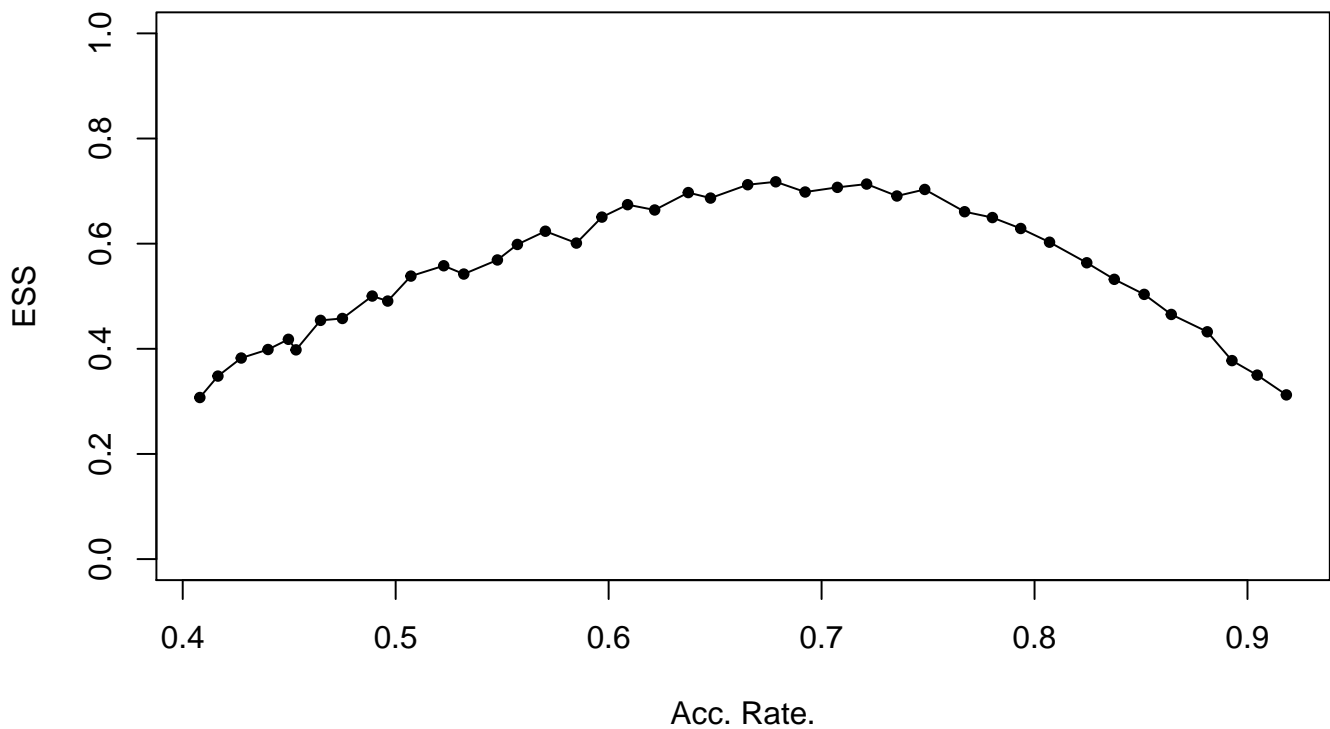
The effective sample sizes are plotted as proportions of the run-length  $N = 100001$ .

```
#cbind(del.vec,ESSd,acc.vec)
par(mar=c(4,4,1,0))
plot(del.vec,ESSd/N,ylim=range(0,1),type='l',xlab=expression(delta),ylab='ESS')
points(del.vec,ESSd/N,pch=20)
```





```
plot(acc.vec, ESSd/N, pch=20, ylim=range(0,1), type='l', xlab='Acc. Rate.', ylab='ESS')
points(acc.vec, ESSd/N, pch=20)
```

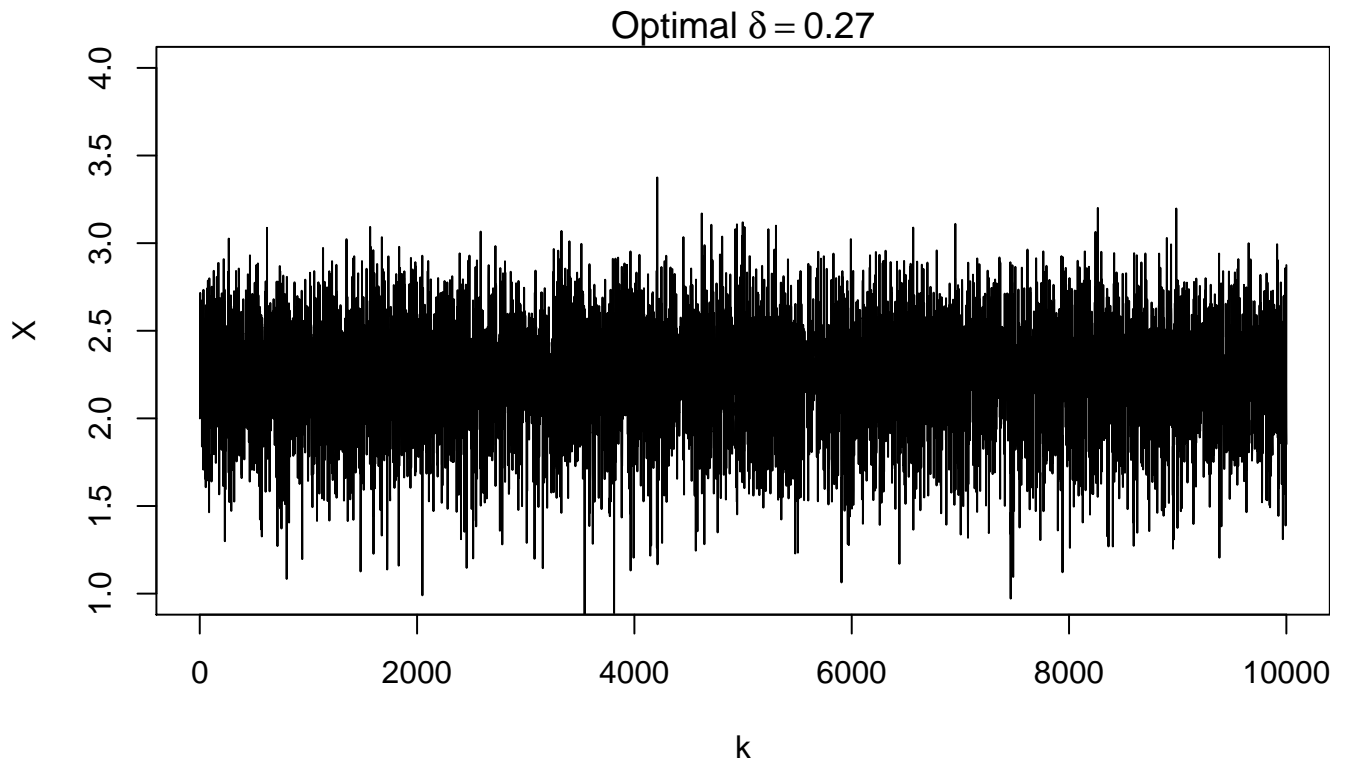


From the plots we deduce that the optimal  $\delta$  in this problem is approximately 0.270, which returns an acceptance rate of 0.678 and an effective sample size of 71763.932.

```

par(mar=c(4,4,1,0))
tval<-substitute(paste('Optimal ',delta)==d,list(d = del.vec[which.max(ESSd)]))
plot(xd[1:10000,which.max(ESSd)],type='l',xlab='k',ylab=expression(X),
     main=tval,ylim=range(1,4))

```



```

hist(xd[1:10000,which.max(ESSd)],freq=FALSE,xlab=expression(X),breaks=seq(0,5,by=0.1),
     main=tval,ylim=range(0,1.5));box();lines(xvec,yvec,col='red')

```

