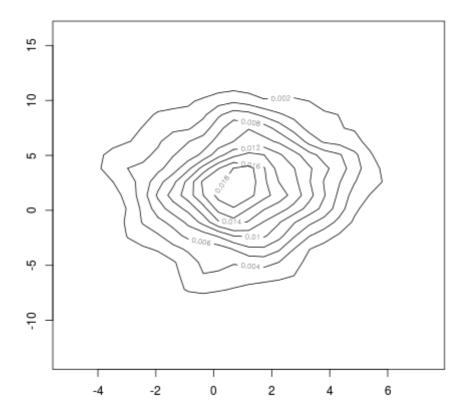
MONTE CARLO ENDSEM REPORT – Vaibhav Saxena (140123041)

Question 1

Algorithm for generating bivariate normal distribution (X,Y) using Cholesky's decomposition:

```
1. Generate Z1, Z2 \sim N(0,1)
    2. Set X = \mu 1 + \sigma 1 \times Z1
    3. Set Y = \mu 2 + \sigma 2 p^* Z 1 + \sigma 2 \sqrt{(1 - \rho^2)} Z 2
Here \mu 1 = 1, \sigma 1 = 2, \mu 2 = 2, \sigma 2 = 4
Code for R:
require(MASS)
# For X
mu1 <- 1
sig1 < -2
# For Y
mu2 <- 2
sig2 <- 4
rho <- 0.05
normvar <- c(0.9597, -1.3404, 1.2238, 0.2551)
# Using Cholesky's decomposition method
noofrand <- 1000
z1 <- rnorm(noofrand, mean=0, sd=1)
z2 <- rnorm(noofrand, mean=0, sd=1)</pre>
X < -mu1 + sig1*z1
Y \le mu^2 + sig^2 rho^2 z^1 + sig^2 sqrt(1 - rho^2)^2 z^2
g \le kde2d(X,Y)
png(file="contour.png")
contour(g)
dev.off()
```

The contour graph for the bivariate distribution (X,Y) can be shown in the figure below:



Question 2

```
Code for R:
```

```
alpha <- 0.4
noofrand <- 200

# Exponential distribution with mean 2
exp2 <- -2*log(runif(noofrand, min=0, max=1))

# Exponential distribution with mean 3
exp3 <- -3*log(runif(noofrand, min=0, max=1))

mixed <- vector('numeric')

for(i in 1:noofrand) {
        if(runif(1,min=0,max=1) < alpha) {
            mixed <- c(mixed, exp2[i])
        }
        else {
            mixed <- c(mixed, exp3[i])
        }
}</pre>
```

```
png(file="exponential_2")
hist(exp2, col="red", breaks=20)
dev.off()

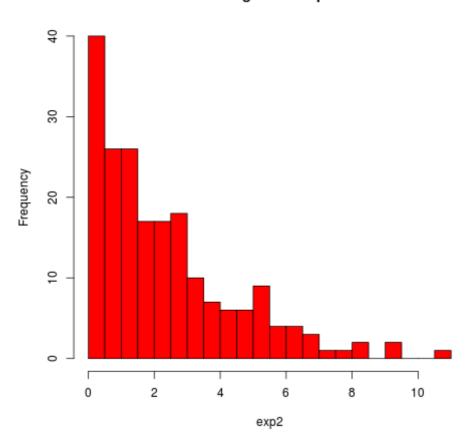
png(file="exponential_3")
hist(exp3, col="red", breaks=20)
dev.off()

png(file="mixed_distribution")
hist(mixed, col="red", breaks=20)
dev.off()
```

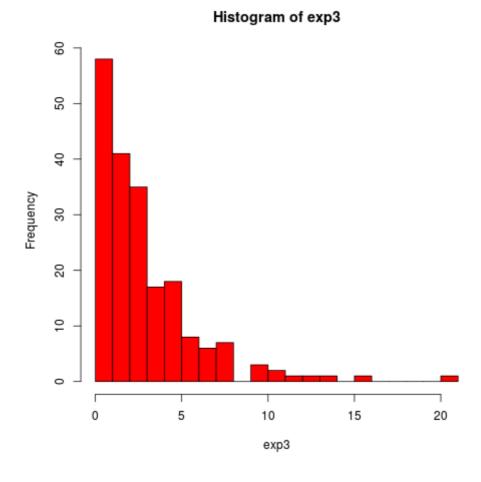
The plotted histrograms are as follows:

For exponential with mean = 2

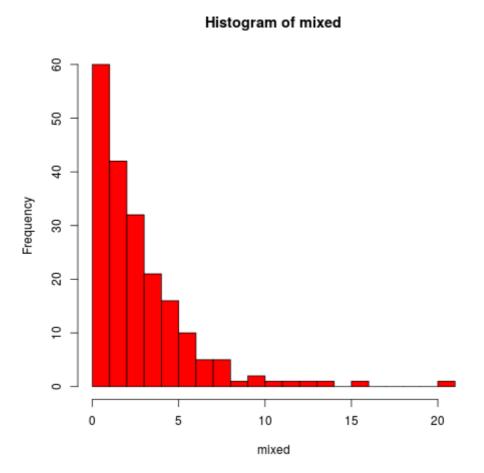
Histogram of exp2



For exponential with mean = 3



For mixed distribution:



Question 3

Code for R:

```
# Naive Monte Carlo method
montecarlo <- function(n, num_estimates) {</pre>
       estimates <- vector(length=num_estimates)</pre>
       for(i in 1:num_estimates) {
               u <- runif(n, min=0, max=1)
               y <- 2*exp(-4*u*u)
               estimates[i] <- mean(y)
       return(c(mean(estimates), var(estimates)))
}
# Antithetic estimate method
antithetic <- function(n, num_estimates) {</pre>
       estimates <- vector(length=num_estimates)</pre>
       for(i in 1:num_estimates) {
               u <- runif(n, min=0, max=1)
               v <- 1-u
               y1 < -2*exp(-4*u*u)
               y2 < -2*exp(-4*v*v)
               y < -(y1+y2)/2
               estimates[i] <- mean(y)
       return(c(mean(estimates), var(estimates)))
}
# Control variate method
controlvariate <- function(n, num_estimates) {</pre>
       estimates <- vector(length=num_estimates)</pre>
       for(i in 1:num_estimates) {
               u <- runif(n,min=0,max=1)</pre>
               x < -2*exp(-4*u*u)
               y < -2*exp(-2*runif(n,min=0,max=1))
               c < -1*cov(x,y)/var(y)
               w < -x + c*(y - mean(y))
               estimates[i] <- mean(w)
       return(c(mean(estimates), var(estimates)))
}
N < -c(100,1000,10000)
num estimates <- 100
naive_mean <- vector(length=length(N))</pre>
naive_var <- vector(length=length(N))</pre>
anti mean <- vector(length=length(N))
anti_var <- vector(length=length(N))</pre>
```

```
control_mean <- vector(length=length(N))</pre>
control_var <- vector(length=length(N))</pre>
for(i in 1:length(N)) {
       naive_mean[i] <- montecarlo(N[i], num_estimates)[1]</pre>
       naive_var[i] <- montecarlo(N[i], num_estimates)[2]</pre>
       anti_mean[i] <- antithetic(N[i], num_estimates)[1]</pre>
       anti_var[i] <- antithetic(N[i], num_estimates)[2]</pre>
       control_mean[i] <- controlvariate(N[i], num_estimates)[1]</pre>
       control_var[i] <- controlvariate(N[i], num_estimates)[2]</pre>
}
for(i in 1:length(N)) {
       cat("For N = ", N[i], "\n")
       cat("Naive Monte Carlo: \n")
       cat("mean =",naive_mean[i],"\n")
       cat("variance =",naive_var[i],"\n")
       cat("Antithetic Estimate: \n")
       cat("mean =",anti_mean[i],"\n")
       cat("variance =",anti_var[i],"\n")
       red1 <- 100*(naive_var[i] - anti_var[i])/naive_var[i]</pre>
       cat("Variance reduction =",red1,"\n")
       cat("Control Variate: \n")
       cat("mean =",control_mean[i],"\n")
       cat("variance =",control var[i],"\n")
       red2 <- 100*(naive_var[i] - control_var[i])/naive_var[i]</pre>
       cat("Variance reduction =",red2,"\n")
       cat("\n")
}
Output:
For N = 100
Naive Monte Carlo:
mean = 0.8839491
variance = 0.005563168
Antithetic Estimate:
mean = 0.8807665
variance = 8.24897e-05
Variance reduction = 98.51722
Control Variate:
mean = 0.8750775
variance = 0.004606603
For N = 1000
Naive Monte Carlo:
mean = 0.8893941
```

variance = 0.000378082

Antithetic Estimate: mean = 0.8824278 variance = 8.552757e-06 Variance reduction = 97.73786 Control Variate: mean = 0.8794938 variance = 0.0004289711

For N = 10000 Naive Monte Carlo: mean = 0.8820661 variance = 4.520528e-05 Antithetic Estimate: mean = 0.8820634 variance = 1.030306e-06 Variance reduction = 97.72083 Control Variate: mean = 0.8820974 variance = 4.503716e-05

N = 100	Mean	Variance
Naive	0.8839491	0.005563168
Antithetic	0.8807665	8.24897e-05
Control	0.8750775	0.004606603

Variance reduction = 98.51722 %

N = 1000	Mean	Variance
Naive	0.8893941	0.000378082
Antithetic	0.8824278	8.552757e-06
Control	0.8794938	0.0004289711

Variance reduction = 97.73786 %

N = 10000	Mean	Variance
Naive	0.8820661	4.520528e-05
Antithetic	0.8820634	1.030306e-06
Control	0.8820974	4.503716e-05

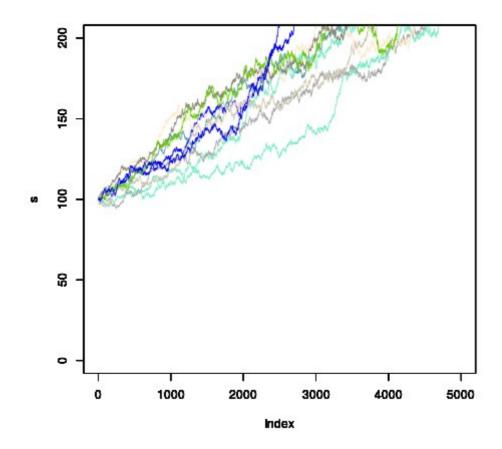
Variance reduction = 97.72083 %

Question 4

```
# Box-Mulle
```

```
# Box-Muller to generate normal distribution
boxmuller <- function(n) {</pre>
       R < -2*log(runif(n,min=0,max=1))
       u < -runif(n,min=0,max=1)
       u <- 2*pi*u
       x \leq - sqrt(R)*cos(u)
       return(x)
}
t <- seq(from=0,to=5,length.out=5000)
mu <- 0.2
sig < -0.1
s5 <- vector(length=10)
#Generating Geometric Brownian Motion
png(file="gbm.png")
for(n in 1:10) {
       s <- vector(length=5000)
       s[1] < -100
       for(i in 2:5000) {
              s[i] <- s[i-1]*exp((mu - 0.5*sig*sig)*(t[i]-t[i-1]) + sig*sqrt(t[i]-t[i-1])*boxmuller(1))
       s5[n] <- s[5000]
       plot(s, type="l", col=colors()[as.integer(runif(1)*100)], ylim=c(100-100,100+100))
       par(new=TRUE)
}
s5meanth <- s[1]*exp(mu*5)
s5varth <- s[1]*s[1]*exp(mu*5)*exp(mu*5)*(exp(sig*sig*5)-1)
cat("Simulated E(S(5)) =", mean(s5), "\n")
cat("Simulated Var(S(5)) = ", var(s5), "\n")
cat("Theoretical E(S(5)) = ", s5meanth, "\n")
cat("Theoretical Var(S(5)) = ", s5varth, "\n")
Output:
Simulated E(S(5)) = 275.367
Simulated Var(S(5)) = 2421.426
Theoretical E(S(5)) = 271.8282
Theoretical Var(S(5)) = 3788.45
```

The generated Geometric Brownian is plotted as below:



Observation:

The simulated and theoretical values of E(S(5)) and Var(S(5)) are in vicinity. Hence the simulated brownian motion is an acceptable approximation.