

# DATA 604 HW 4

Dan Fanelli

10/23/2016

```
library(ggplot2)
```

```
## Warning: package 'ggplot2' was built under R version 3.2.4
```

```
library(knitr)
```

```
# x is a D-dimensional random variable (i.e., a Dx1 column vector) and each component of x is uniformly
```

```
cost_func <- function(x,D=1){  
  denominator <- ((2*pi)^(D/2))  
  return ((1/denominator)*(exp(1)^(-0.5)*(t(x) %*% x)))  
}
```

```
truExp <- function(D=1){  
  return ((1/10)^D)  
}
```

```
truExp()
```

```
## [1] 0.1
```

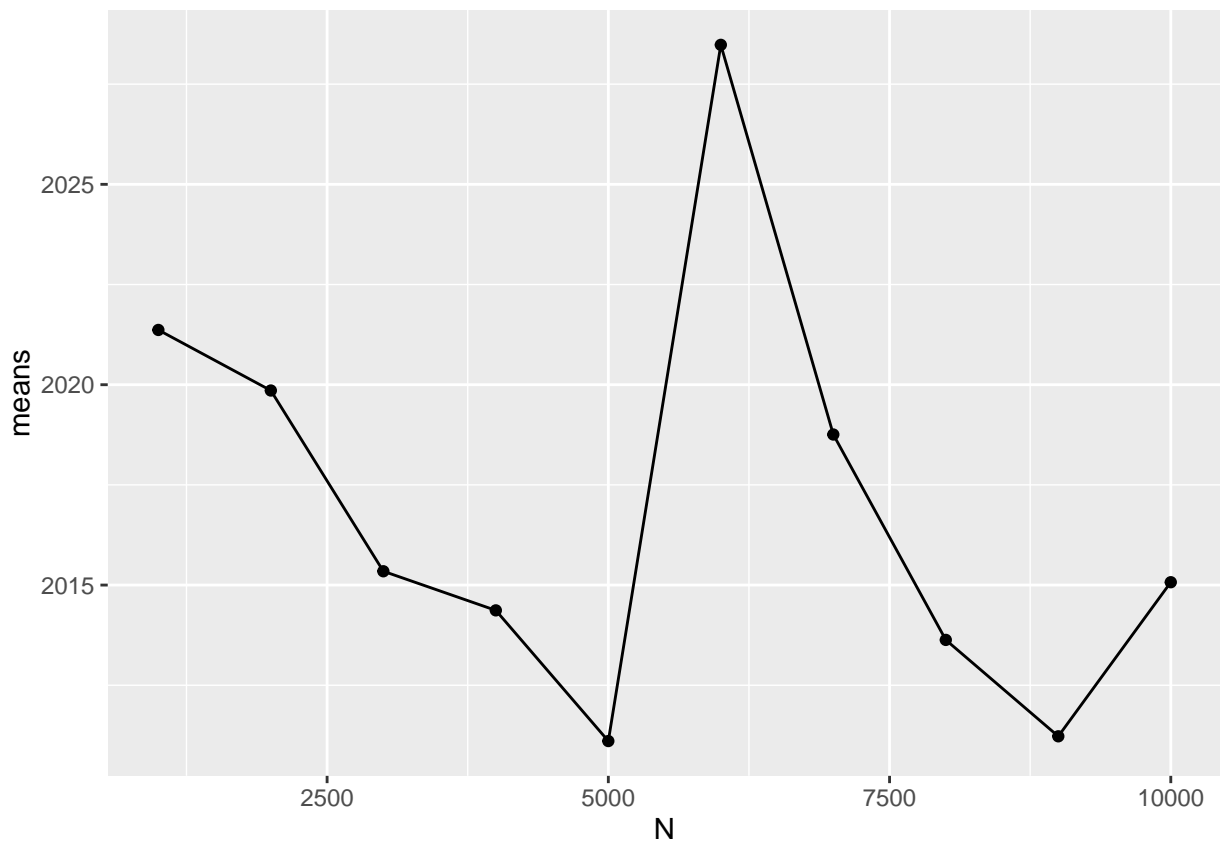
## 1a - Crude Monte Carlo

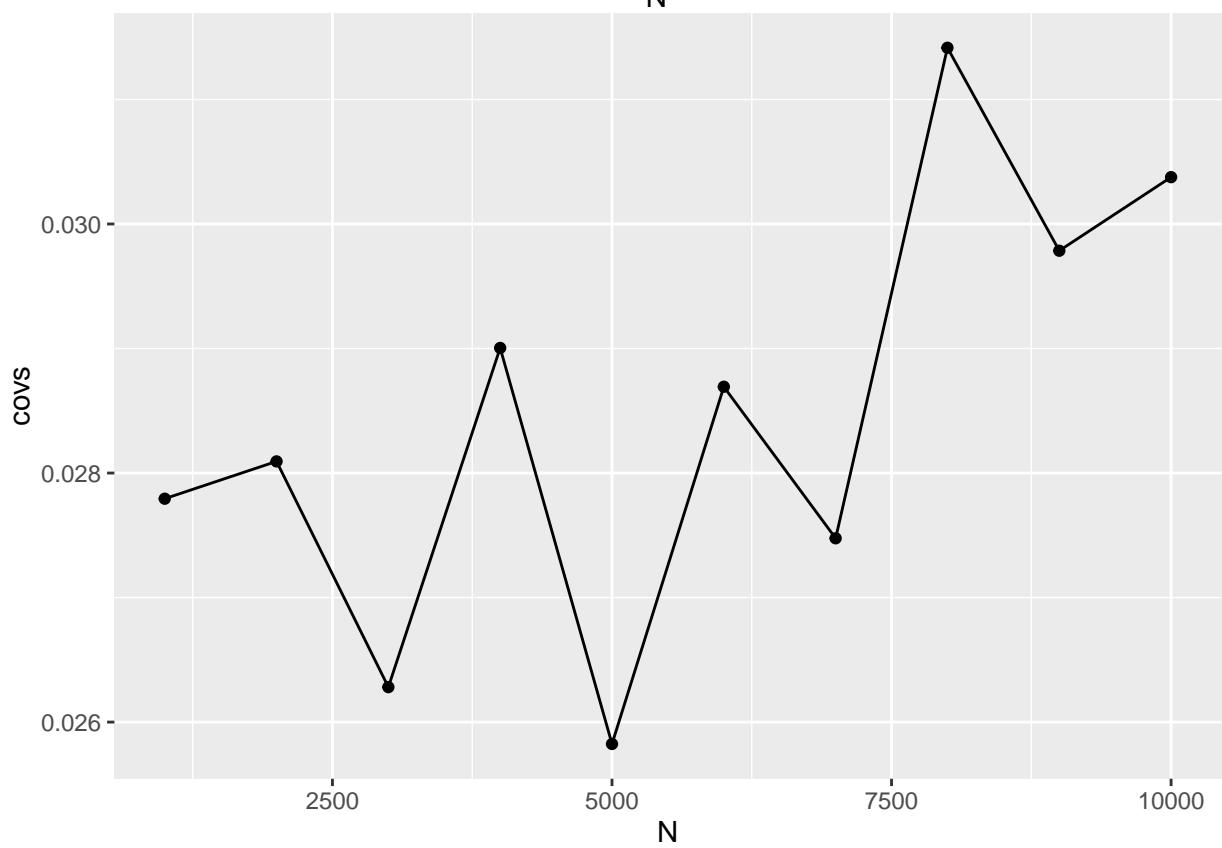
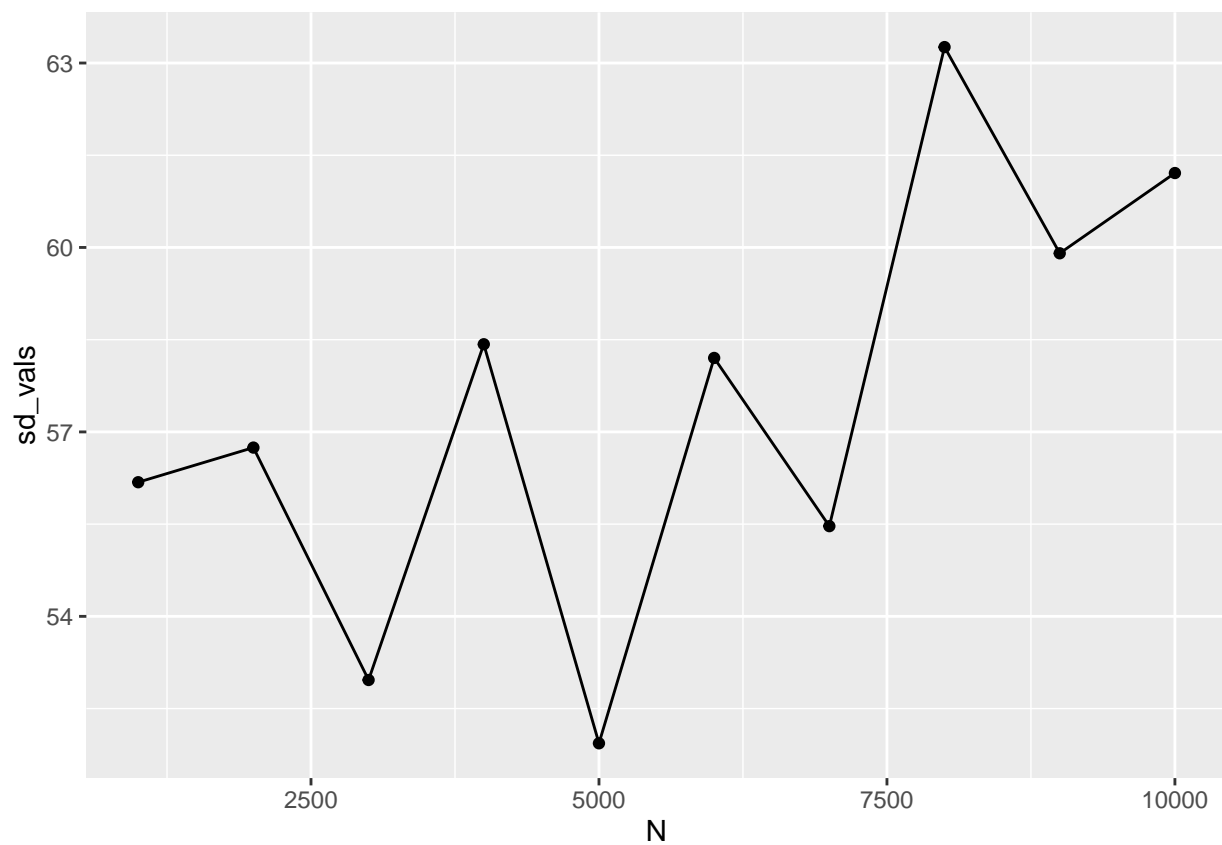
```
go_1a <- function(D=1){  
  N <- 1000 * c(1:10)  
  means <- c()  
  sd_vals <- c()  
  covs <- c()  
  
  for(n in seq(1000,10000,1000)){  
    estimates <- c()  
    for(k in 1:100){  
      estimates <- c(estimates, cost_func(runif(1000, min = -5, max = 5)))  
    }  
    #print(estimates)  
    means <- c(means, mean(estimates))  
    sd_vals <- c(sd_vals, sd(estimates))  
    covs <- c(covs, (sd(estimates) / mean(estimates)))  
  }  
  results <- data.frame(N,means,sd_vals,covs)  
  print(results)  
  
  print(ggplot(data=results, aes(x=N, y=means)) + geom_line() + geom_point())  
  print(ggplot(data=results, aes(x=N, y=sd_vals)) + geom_line() + geom_point())
```

```
print(ggplot(data=results, aes(x=N, y=covs)) + geom_line() + geom_point())
}

go_1a()
```

##	N	means	sd_vals	covs
## 1	1000	2021.367	56.18168	0.02779391
## 2	2000	2019.855	56.74489	0.02809355
## 3	3000	2015.345	52.96503	0.02628088
## 4	4000	2014.368	58.42590	0.02900457
## 5	5000	2011.108	51.93551	0.02582433
## 6	6000	2028.481	58.20300	0.02869290
## 7	7000	2018.757	55.46734	0.02747599
## 8	8000	2013.631	63.25906	0.03141541
## 9	9000	2011.226	59.90533	0.02978548
## 10	10000	2015.071	61.21023	0.03037621





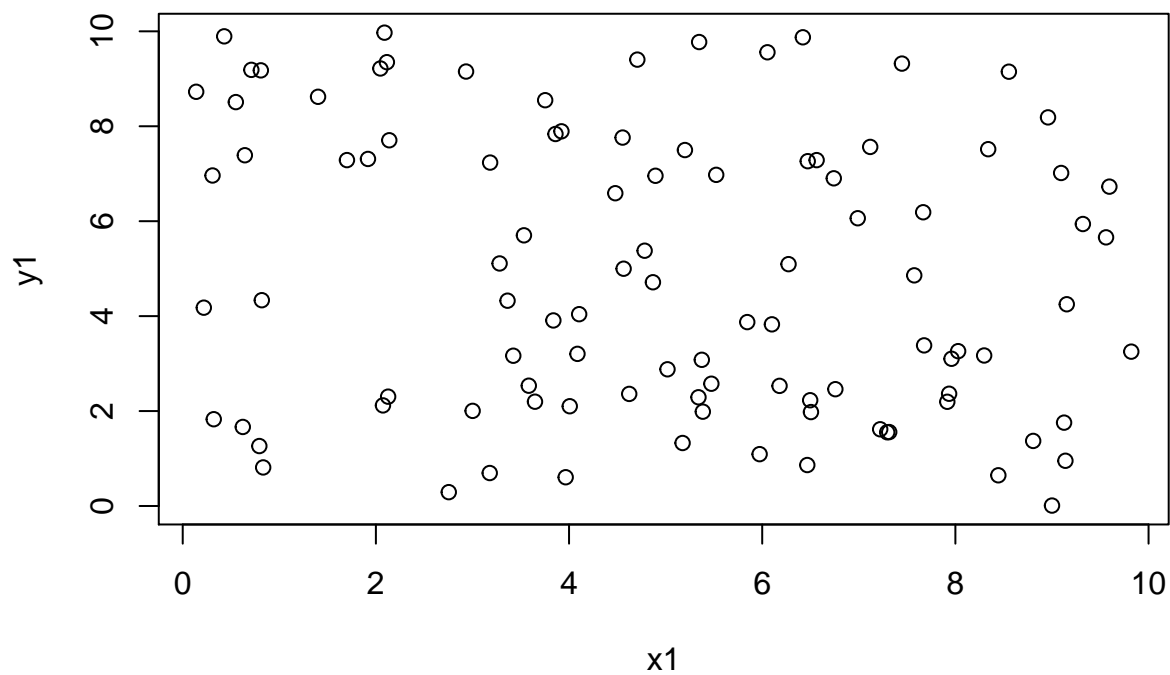
## 1b - Quasi Random Numbers

```
library(qrng)
```

```
## Warning: package 'qrng' was built under R version 3.2.5
```

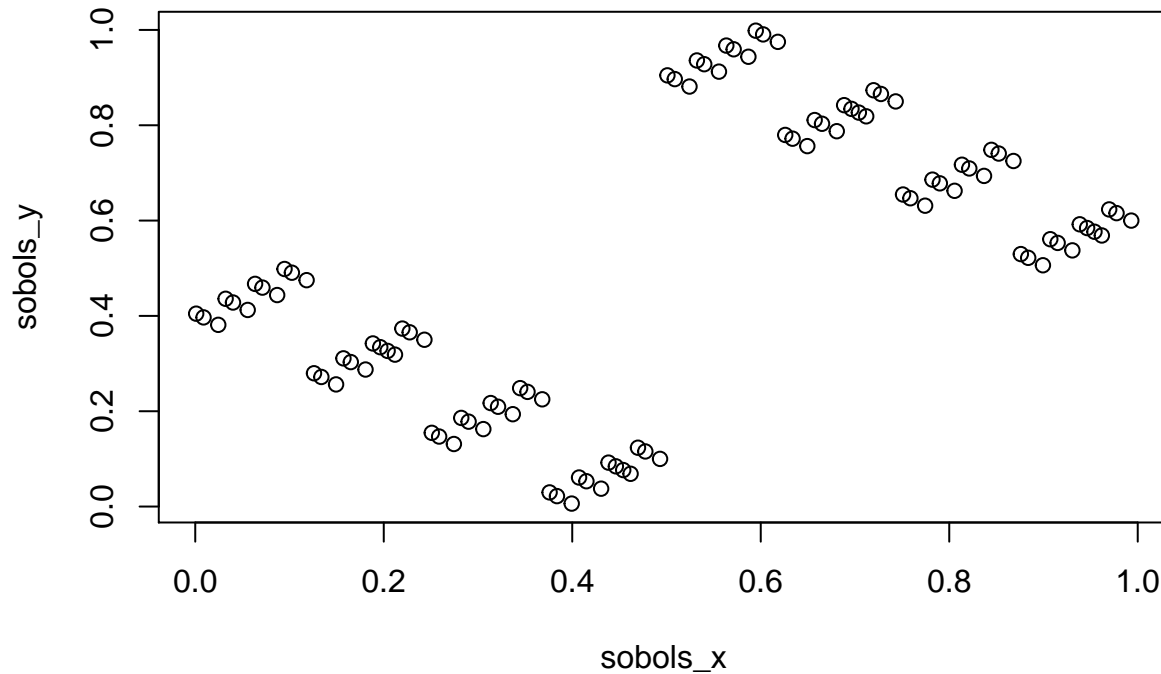
```
x1 <- runif(100,0,10)  
y1 <- runif(100,0,10)
```

```
plot(x1, y1)
```



```
sobols_x <- sobol(n=100, d=1, randomize = TRUE)  
sobols_y <- sobol(n=100, d=1, randomize = TRUE)
```

```
plot(sobols_x, sobols_y)
```



## 1c - Antithetic Variates

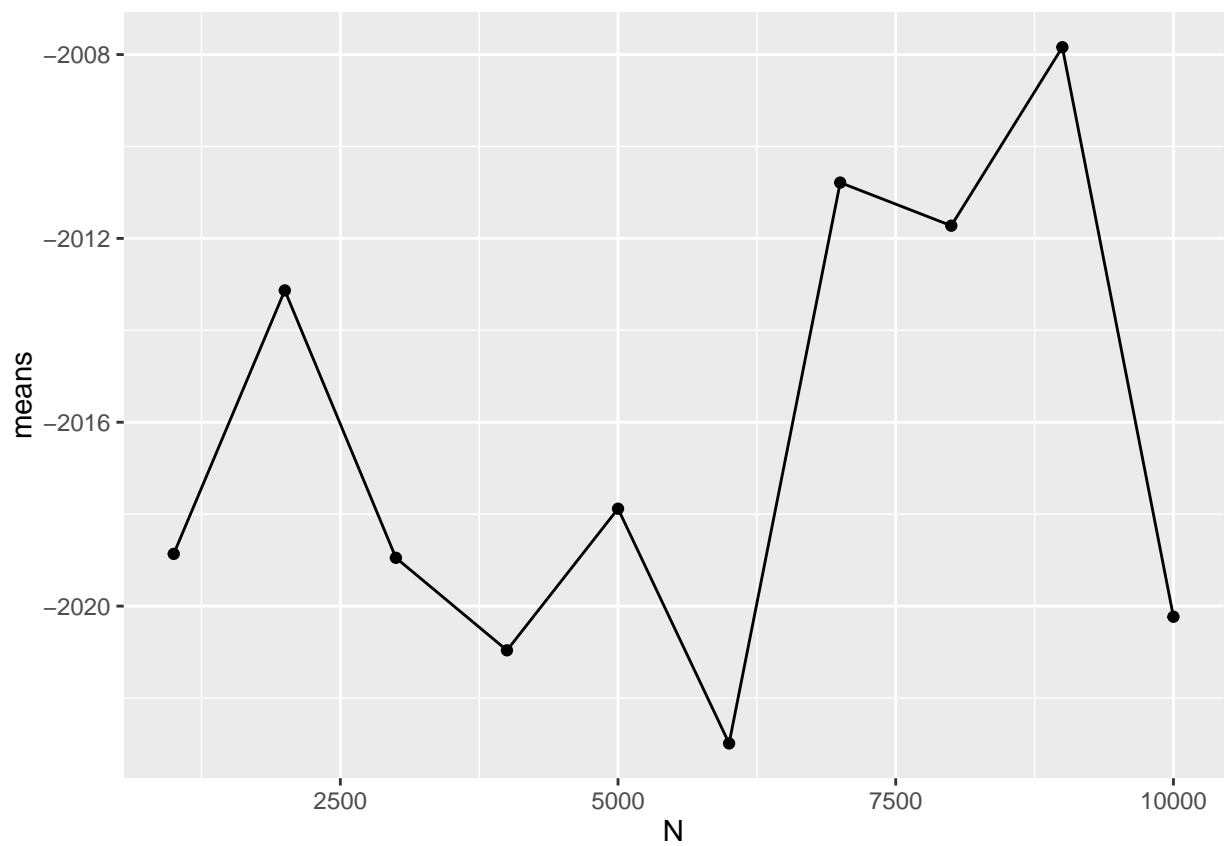
```
go_1c <- function(D=1){
  N <- 1000 * c(1:10)
  means <- c()
  sd_vals <- c()
  covs <- c()

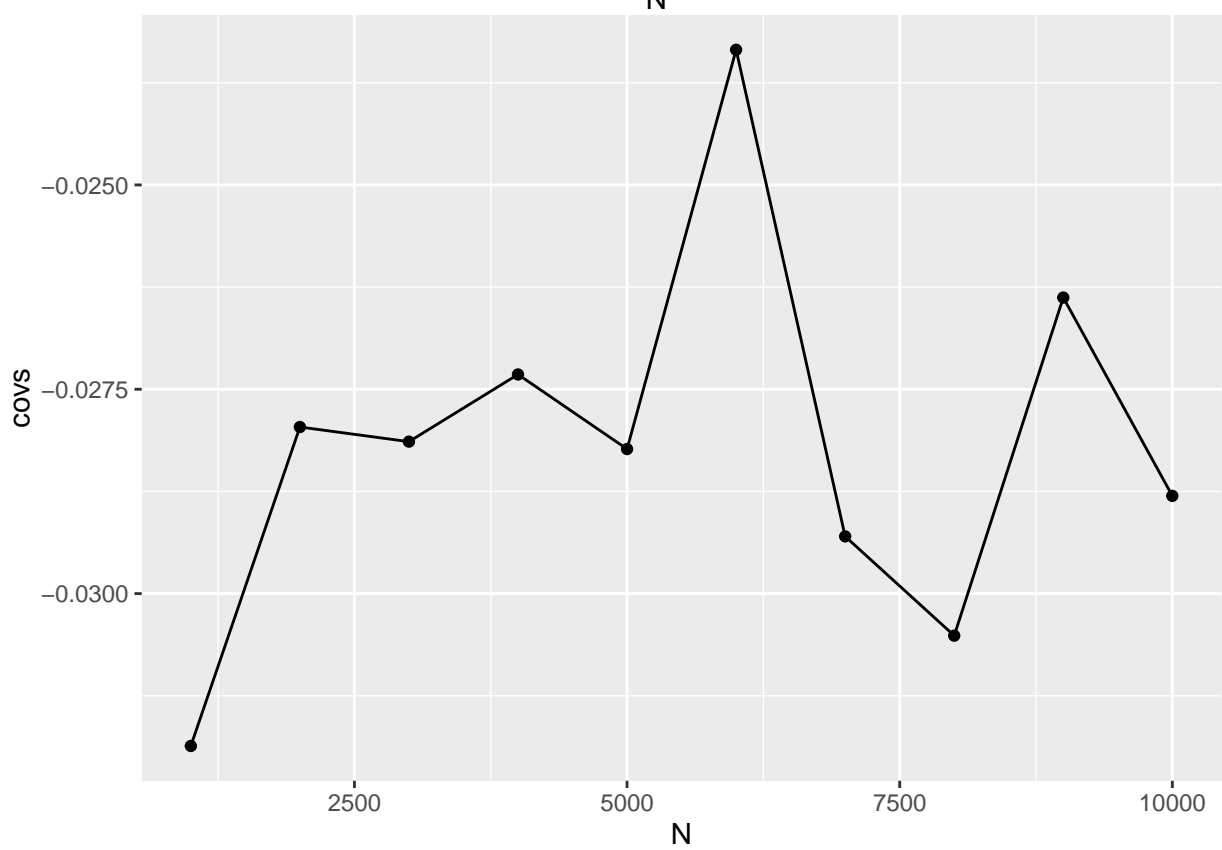
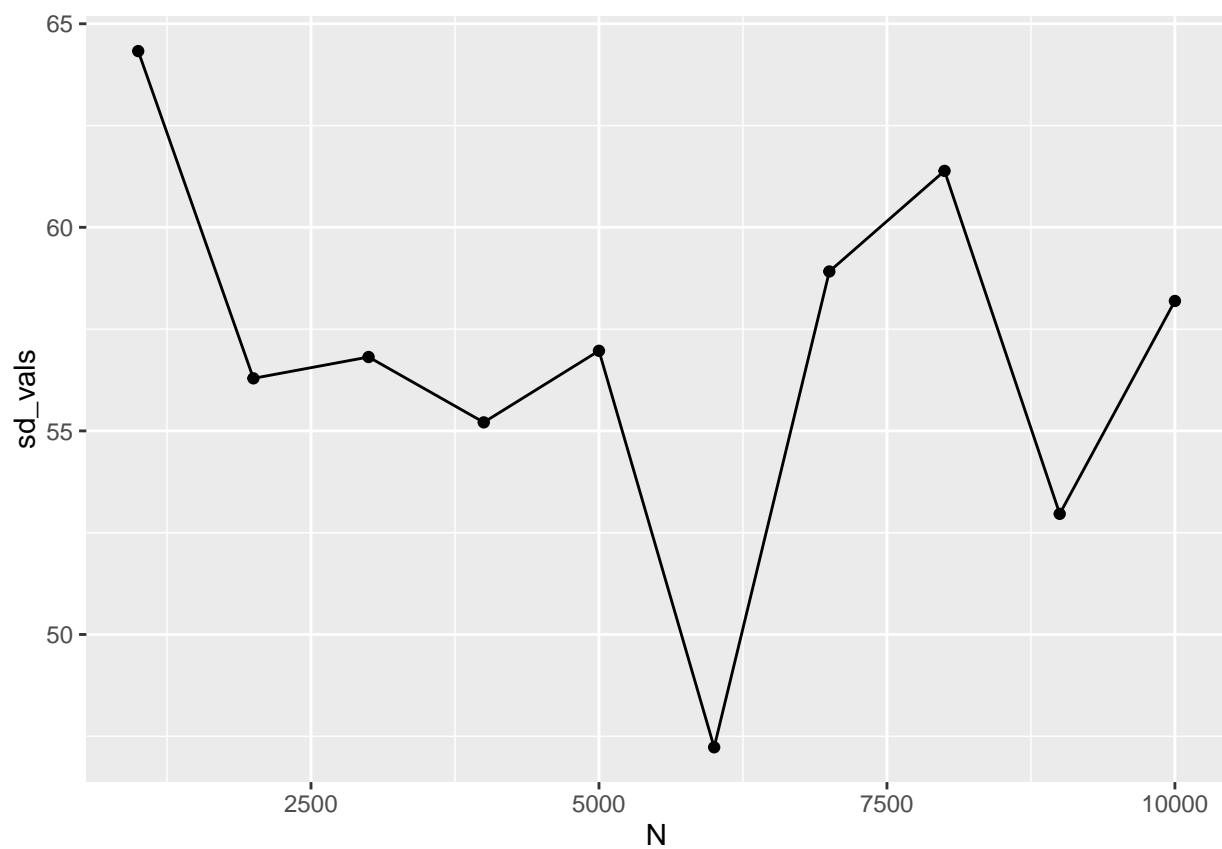
  for(n in seq(1000,10000,1000)){
    estimates <- c()
    for(k in 1:100){
      positives <- cost_func(runif(1000, min = -5, max = 5))
      negatives <- positives * (-1)
      #estimates <- c(estimates, positives, positives * (-1))
      estimates <- c(estimates, negatives)
    }
    #print(estimates)
    means <- c(means, mean(estimates))
    sd_vals <- c(sd_vals, sd(estimates))
    covs <- c(covs, (sd(estimates) / mean(estimates)))
  }
  results <- data.frame(N,means,sd_vals,covs)
  print(results)

  print(ggplot(data=results, aes(x=N, y=means)) + geom_line() + geom_point())
  print(ggplot(data=results, aes(x=N, y=sd_vals)) + geom_line() + geom_point())
  print(ggplot(data=results, aes(x=N, y=covs)) + geom_line() + geom_point())
}
```

```
go_1c()
```

##	N	means	sd_vals	covs
## 1	1000	-2018.863	64.32897	-0.03186395
## 2	2000	-2013.132	56.29086	-0.02796184
## 3	3000	-2018.949	56.81487	-0.02814081
## 4	4000	-2020.962	55.21023	-0.02731879
## 5	5000	-2017.881	56.96759	-0.02823140
## 6	6000	-2022.987	47.22925	-0.02334630
## 7	7000	-2010.786	58.91629	-0.02930013
## 8	8000	-2011.724	61.38651	-0.03051439
## 9	9000	-2007.841	52.96255	-0.02637786
## 10	10000	-2020.231	58.19164	-0.02880445





## 1d - Latin Hypercube Sampling

```
library(lhs)
```

```
## Warning: package 'lhs' was built under R version 3.2.5
```

```
go_1d <- function(D=1){
  N <- 1000 * c(1:10)
  means <- c()
  sd_vals <- c()
  covs <- c()

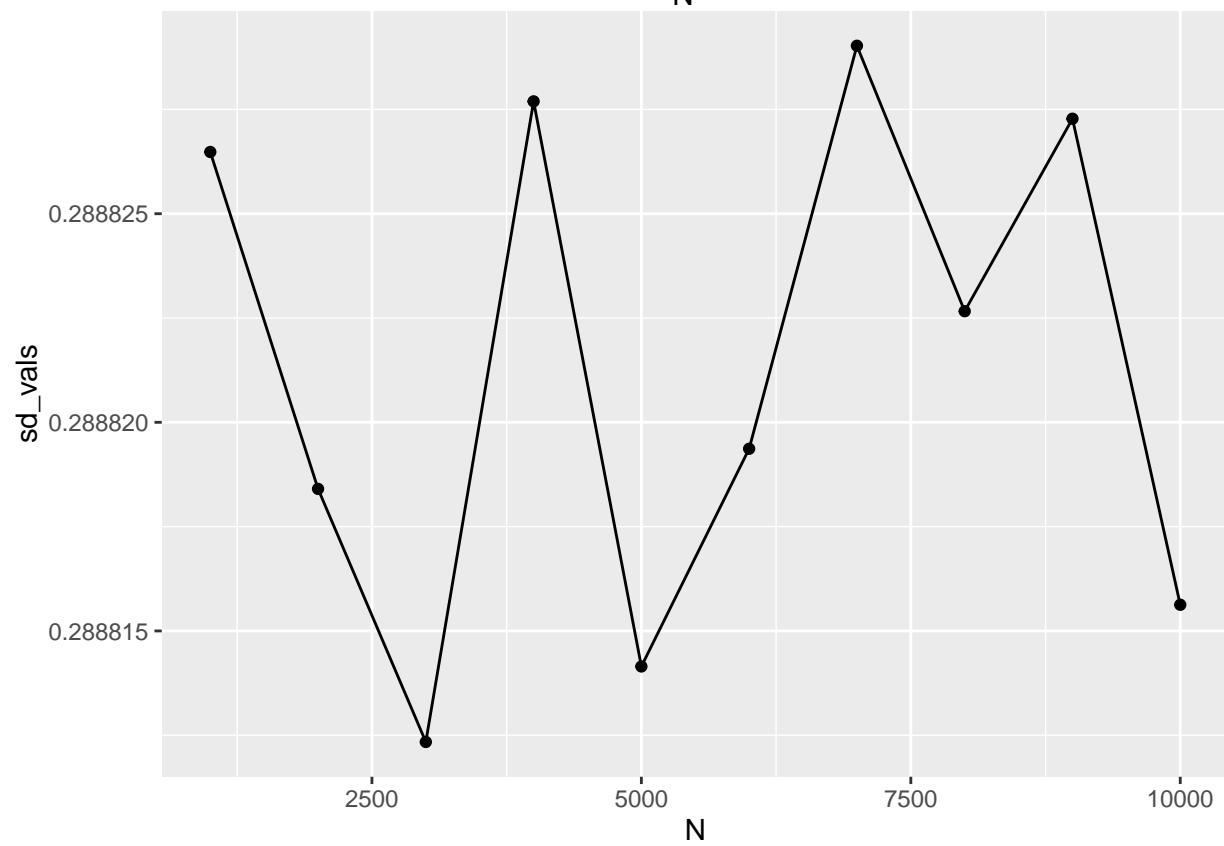
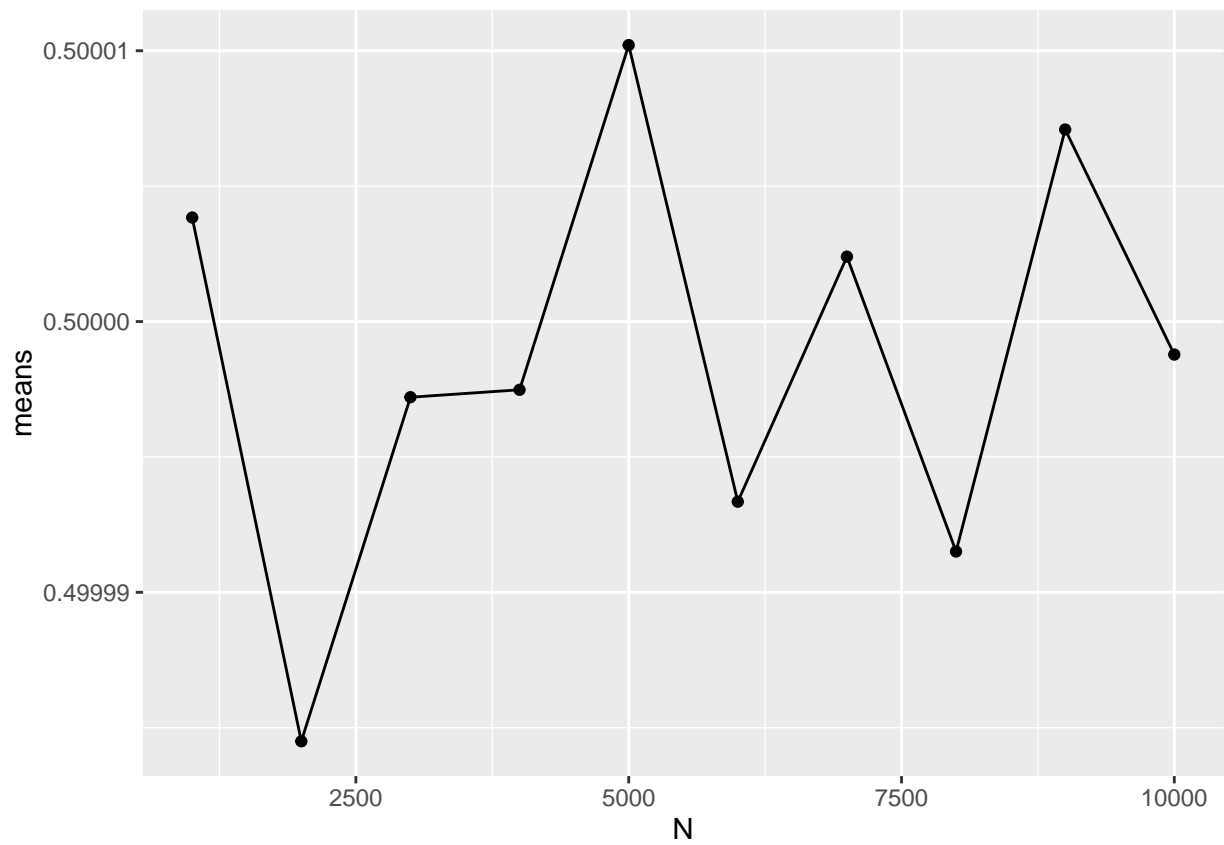
  for(n in seq(1000,10000,1000)){
    estimates <- c()
    for(k in 1:100){
      estimates <- randomLHS(1000, 1)
      #estimates <- c(estimates, cost_func(runif(1000, min = -5, max = 5)))
    }
    #print(estimates)
    means <- c(means, mean(estimates))
    sd_vals <- c(sd_vals, sd(estimates))
    covs <- c(covs, (sd(estimates) / mean(estimates)))
  }
  results <- data.frame(N,means,sd_vals,covs)
  print(results)

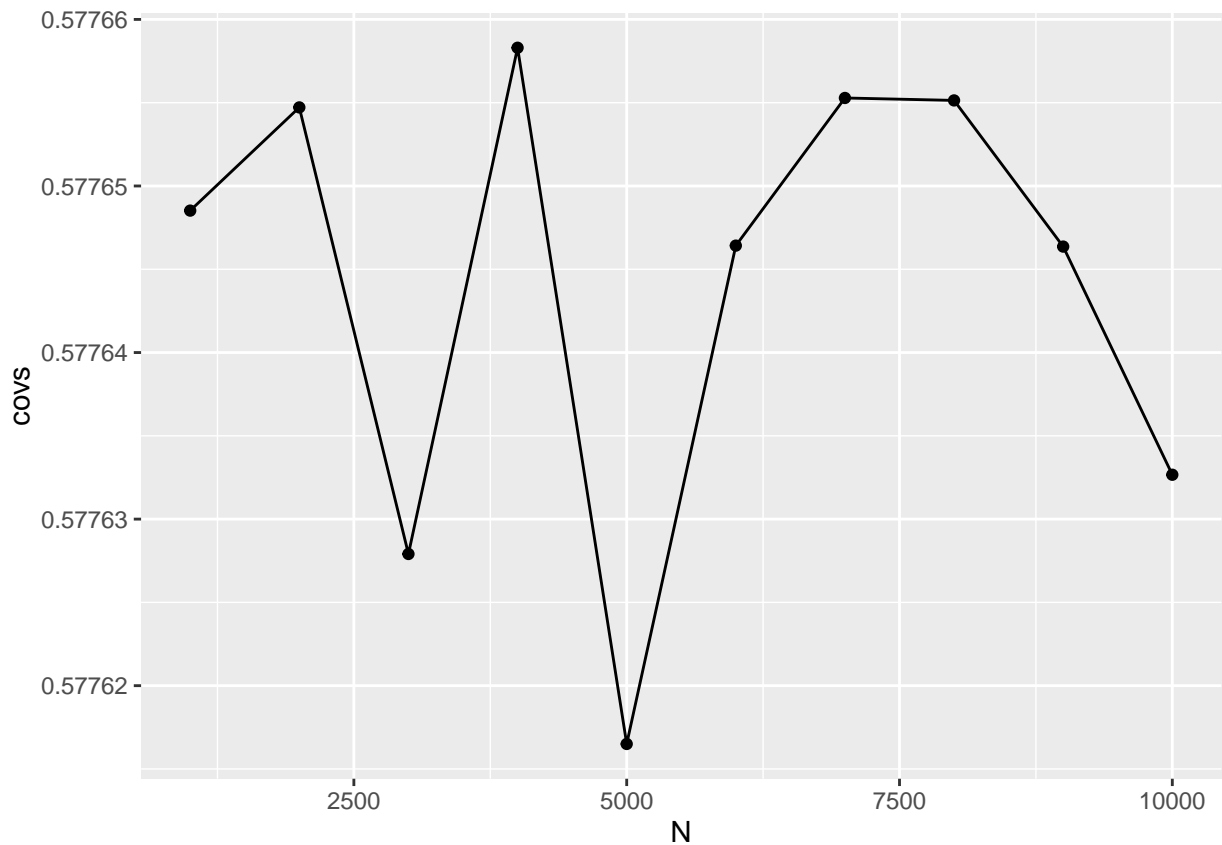
  print(ggplot(data=results, aes(x=N, y=means)) + geom_line() + geom_point())
  print(ggplot(data=results, aes(x=N, y=sd_vals)) + geom_line() + geom_point())
  print(ggplot(data=results, aes(x=N, y=covs)) + geom_line() + geom_point())
}

go_1d()
```

```
##           N      means   sd_vals    covs
## 1    1000 0.5000038 0.2888265 0.5776485
## 2    2000 0.4999845 0.2888184 0.5776547
## 3    3000 0.4999972 0.2888123 0.5776279
## 4    4000 0.4999975 0.2888277 0.5776583
## 5    5000 0.5000102 0.2888141 0.5776165
## 6    6000 0.4999933 0.2888194 0.5776464
## 7    7000 0.5000024 0.2888290 0.5776553
## 8    8000 0.4999915 0.2888227 0.5776551
## 9    9000 0.5000071 0.2888273 0.5776464
## 10 10000 0.4999988 0.2888156 0.5776327
```







## 1e - Importance Sampling

## 1f - Summary

All Code:

```
library(ggplot2)
library(knitr)

# x is a D-dimensional random variable (i.e., a Dx1 column vector) and each component of x is uniformly
cost_func <- function(x,D=1){
  denominator <- ((2*pi)^(D/2))
  return ((1/denominator)*(exp(1)^(-0.5)*(t(x) %*% x)))
}

truExp <- function(D=1){
  return ((1/10)^D)
}

truExp()
go_1a <- function(D=1){
  N <- 1000 * c(1:10)
  means <- c()
```

```

sd_vals <- c()
covs <- c()

for(n in seq(1000,10000,1000)){
  estimates <- c()
  for(k in 1:100){
    estimates <- c(estimates, cost_func(runif(1000, min = -5, max = 5)))
  }
  #print(estimates)
  means <- c(means, mean(estimates))
  sd_vals <- c(sd_vals, sd(estimates))
  covs <- c(covs, (sd(estimates) / mean(estimates)))
}
results <- data.frame(N,means,sd_vals,covs)
print(results)

print(ggplot(data=results, aes(x=N, y=means)) + geom_line() + geom_point())
print(ggplot(data=results, aes(x=N, y=sd_vals)) + geom_line() + geom_point())
print(ggplot(data=results, aes(x=N, y=covs)) + geom_line() + geom_point())
}

go_1a()

library(qrng)

x1 <- runif(100,0,10)
y1 <- runif(100,0,10)

plot(x1, y1)

sobols_x <- sobol(n=100, d=1, randomize = TRUE)
sobols_y <- sobol(n=100, d=1, randomize = TRUE)

plot(sobols_x, sobols_y)
go_1c <- function(D=1){
  N <- 1000 * c(1:10)
  means <- c()
  sd_vals <- c()
  covs <- c()

  for(n in seq(1000,10000,1000)){
    estimates <- c()
    for(k in 1:100){
      positives <- cost_func(runif(1000, min = -5, max = 5))
      negatives <- positives * (-1)
      #estimates <- c(estimates, positives, positives * (-1))
      estimates <- c(estimates, negatives)
    }
    #print(estimates)
    means <- c(means, mean(estimates))
    sd_vals <- c(sd_vals, sd(estimates))
    covs <- c(covs, (sd(estimates) / mean(estimates)))
  }
}

```

```

results <- data.frame(N,means,sd_vals,covs)
print(results)

print(ggplot(data=results, aes(x=N, y=means)) + geom_line() + geom_point())
print(ggplot(data=results, aes(x=N, y=sd_vals)) + geom_line() + geom_point())
print(ggplot(data=results, aes(x=N, y=covs)) + geom_line() + geom_point())
}

go_1c()

library(lhs)
go_1d <- function(D=1){
  N <- 1000 * c(1:10)
  means <- c()
  sd_vals <- c()
  covs <- c()

  for(n in seq(1000,10000,1000)){
    estimates <- c()
    for(k in 1:100){
      estimates <- randomLHS(1000, 1)
      #estimates <- c(estimates, cost_func(runif(1000, min = -5, max = 5)))
    }
    #print(estimates)
    means <- c(means, mean(estimates))
    sd_vals <- c(sd_vals, sd(estimates))
    covs <- c(covs, (sd(estimates) / mean(estimates)))
  }
  results <- data.frame(N,means,sd_vals,covs)
  print(results)

  print(ggplot(data=results, aes(x=N, y=means)) + geom_line() + geom_point())
  print(ggplot(data=results, aes(x=N, y=sd_vals)) + geom_line() + geom_point())
  print(ggplot(data=results, aes(x=N, y=covs)) + geom_line() + geom_point())
}

go_1d()
##

```