

# exercise1

*Alexande Van Roijen*

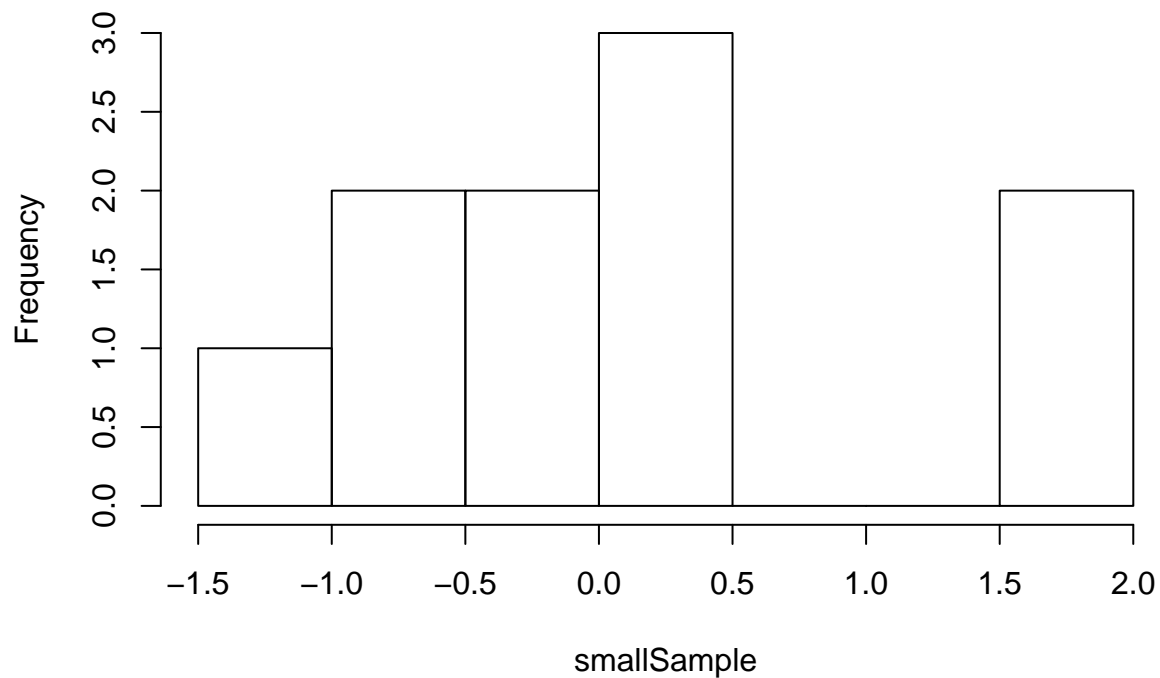
*January 9, 2019*

## R Markdown

### Problem 1

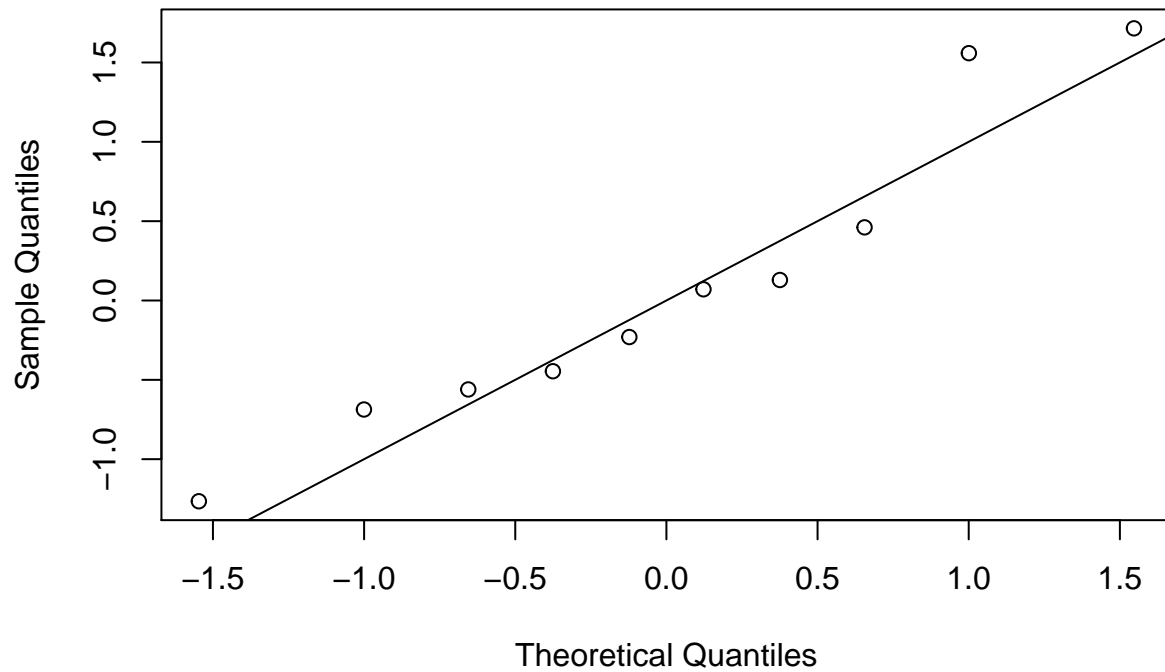
```
set.seed(123)
smallSample = (rnorm(10,0,1))
hist(smallSample)
```

**Histogram of smallSample**



```
qqnorm(smallSample)
abline(0,1)
```

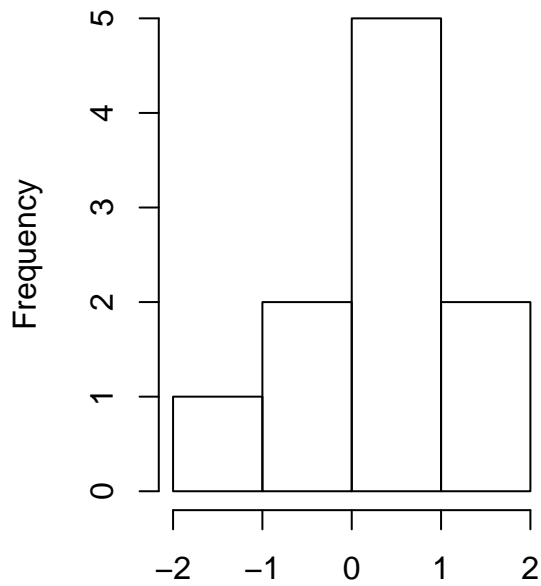
## Normal Q-Q Plot



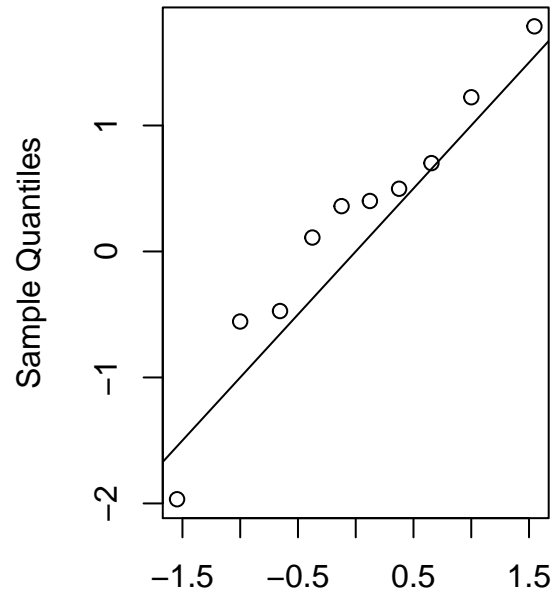
surprisingly, this does look rather normal! or at least the qqplot says so. The histogram does not look normal, more uniform if anything.

```
for(i in 1:20){  
  smallSample = (rnorm(10,0,1))  
  par(mfrow=c(1,2))  
  hist(smallSample)  
  qqnorm(smallSample)  
  abline(0,1)  
}
```

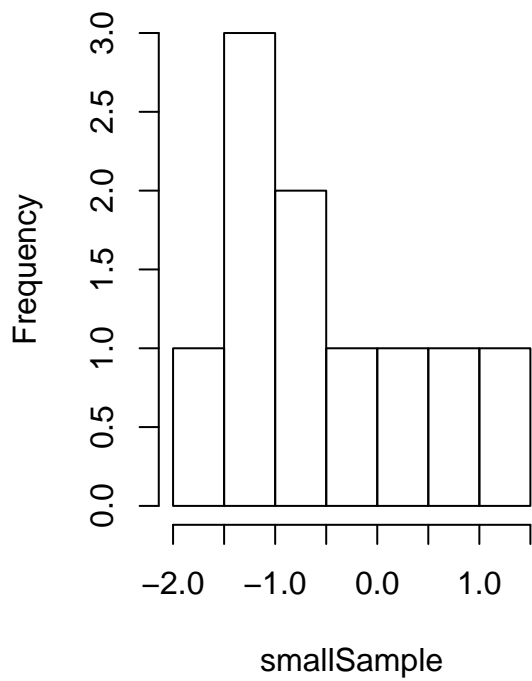
**Histogram of smallSample**



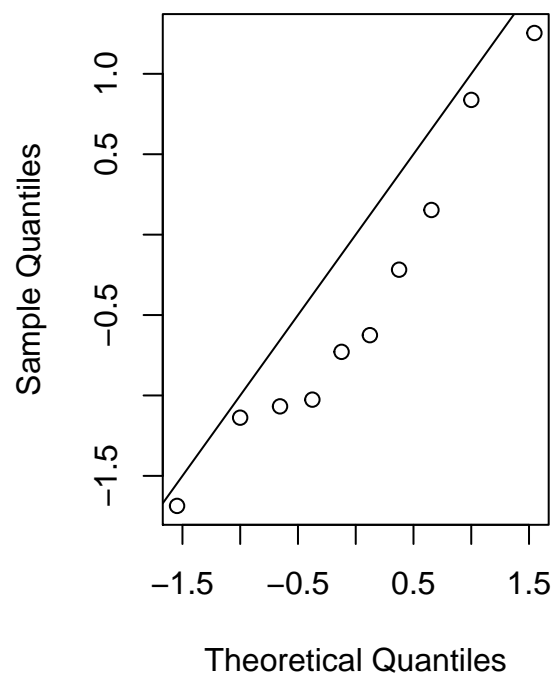
**Normal Q-Q Plot**



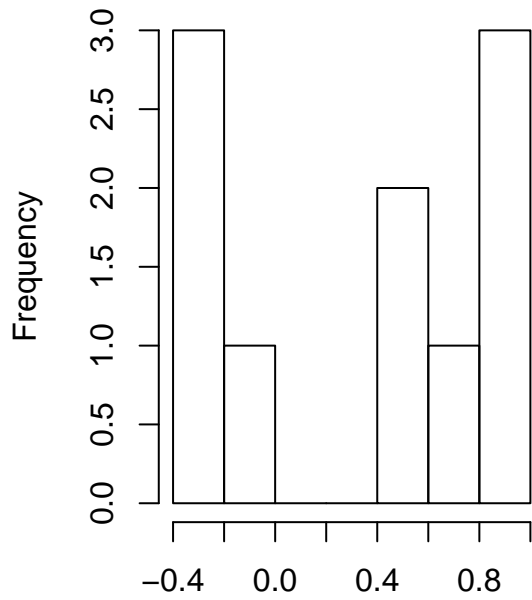
**Histogram of smallSample**



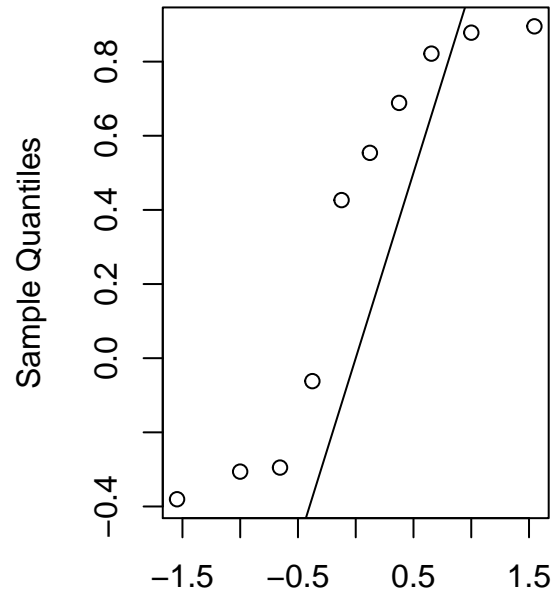
**Normal Q-Q Plot**



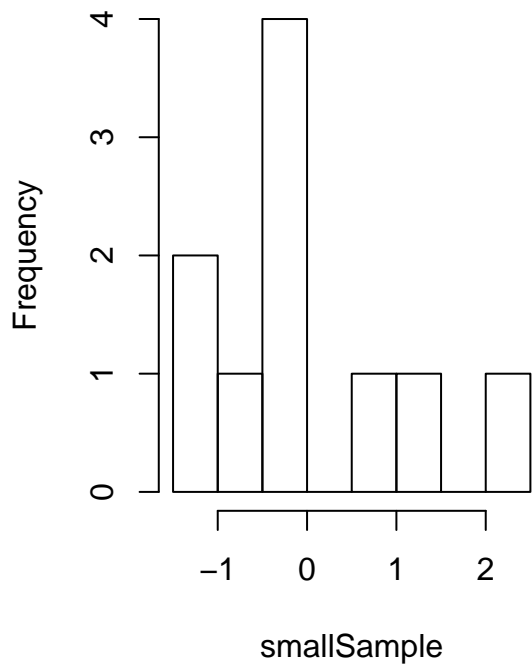
**Histogram of smallSample**



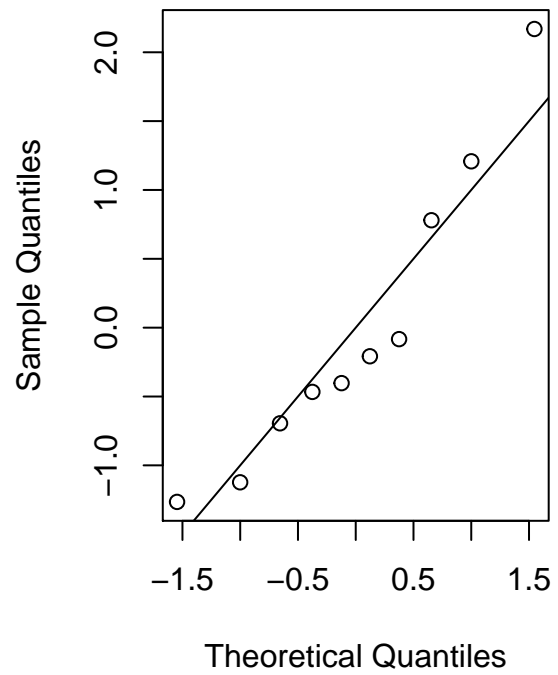
**Normal Q-Q Plot**



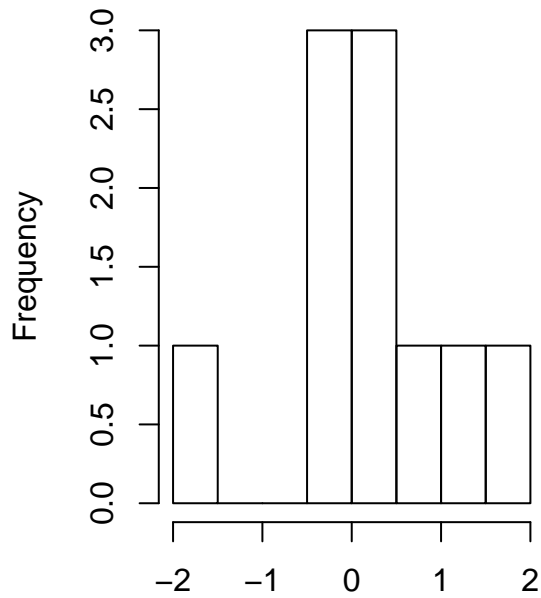
smallSample  
**Histogram of smallSample**



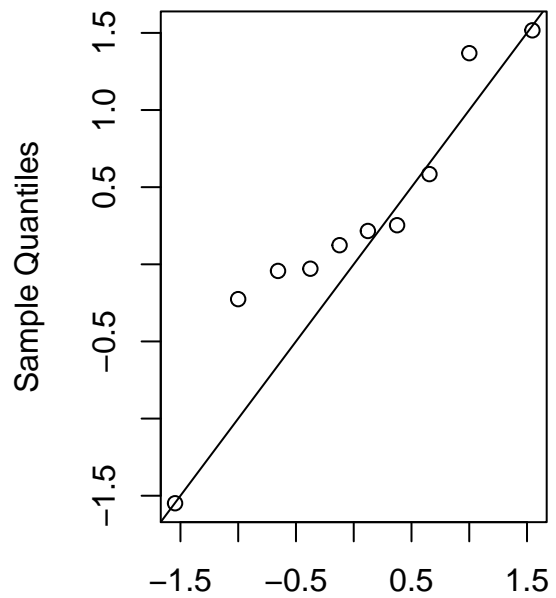
Theoretical Quantiles  
**Normal Q-Q Plot**



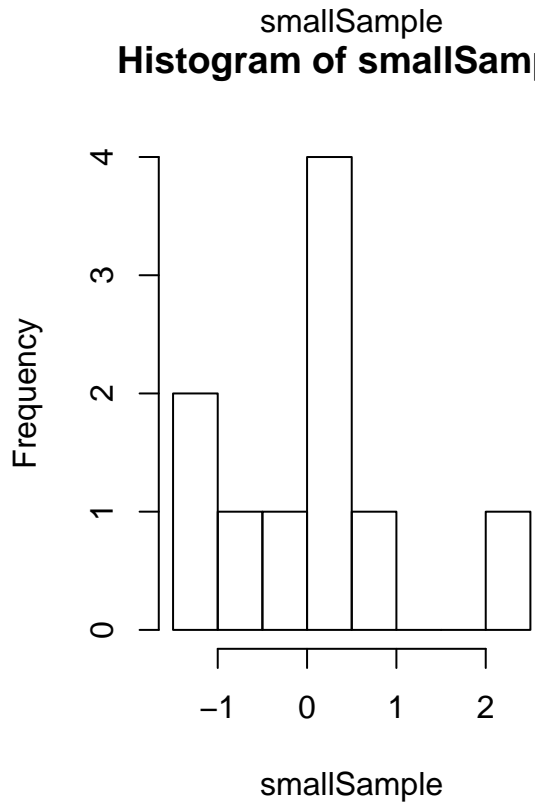
**Histogram of smallSample**



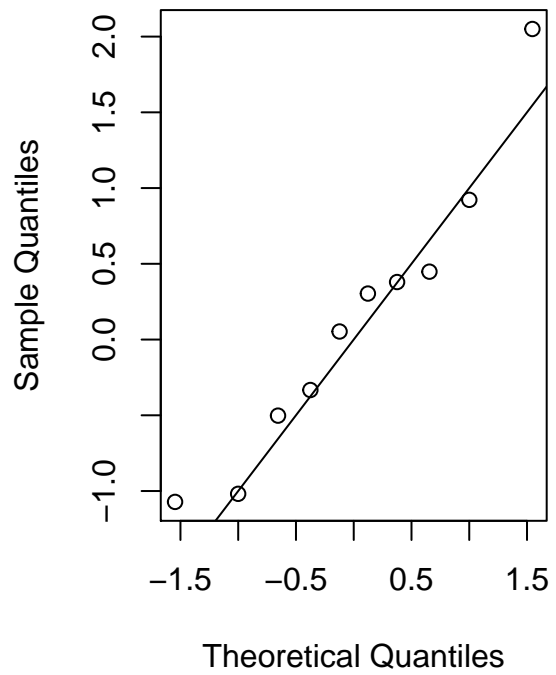
**Normal Q-Q Plot**



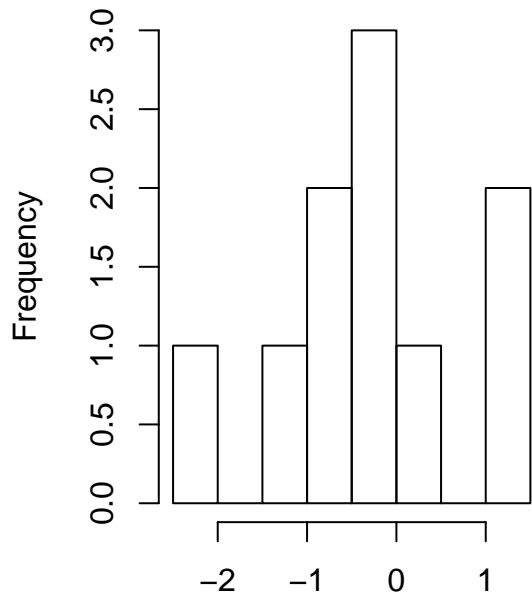
**Histogram of smallSample**



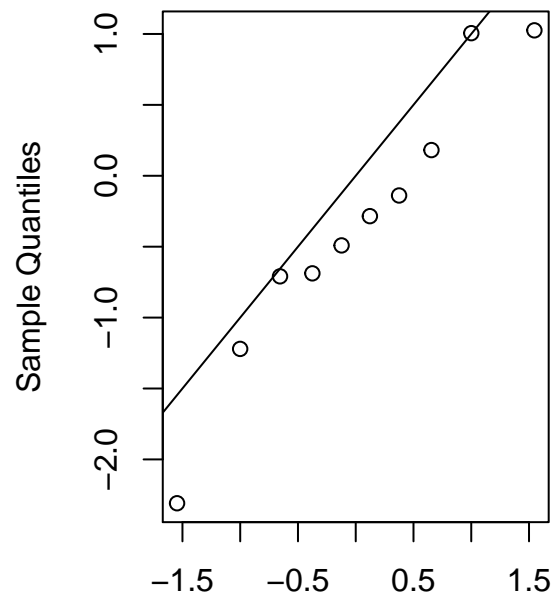
**Normal Q-Q Plot**



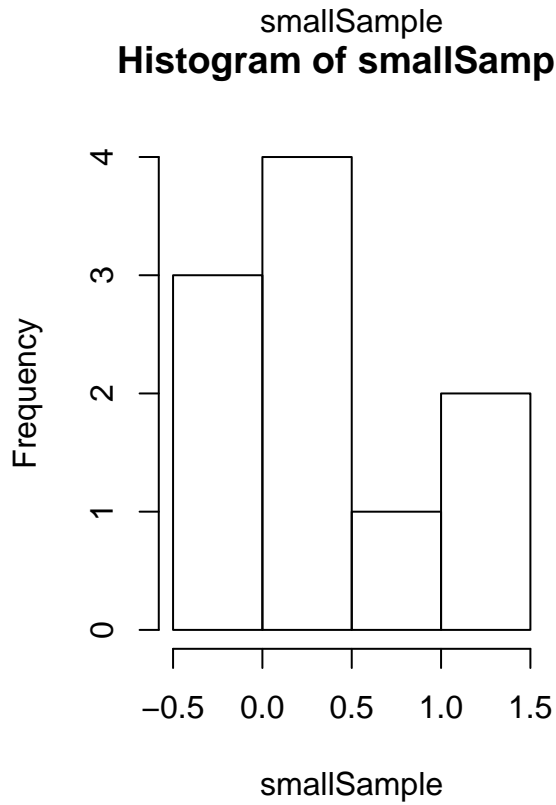
**Histogram of smallSample**



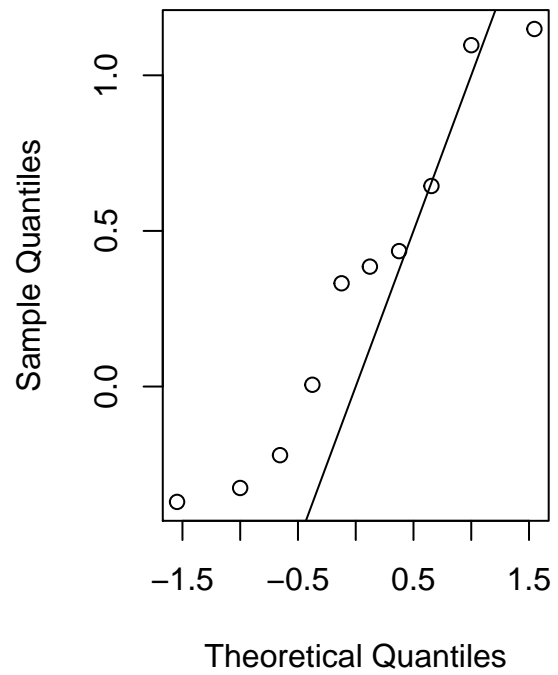
**Normal Q-Q Plot**



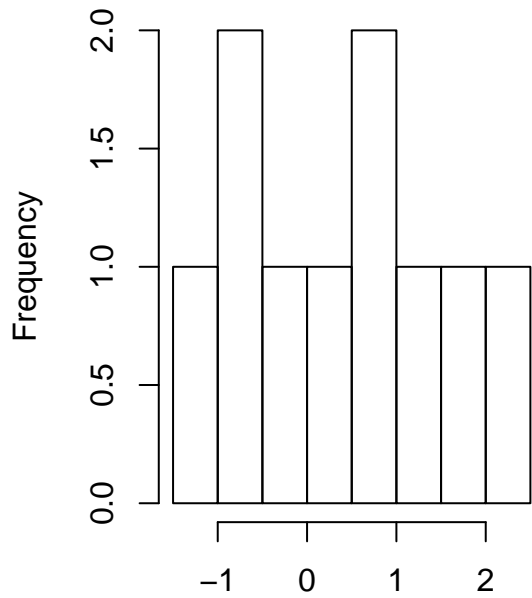
**Histogram of smallSample**



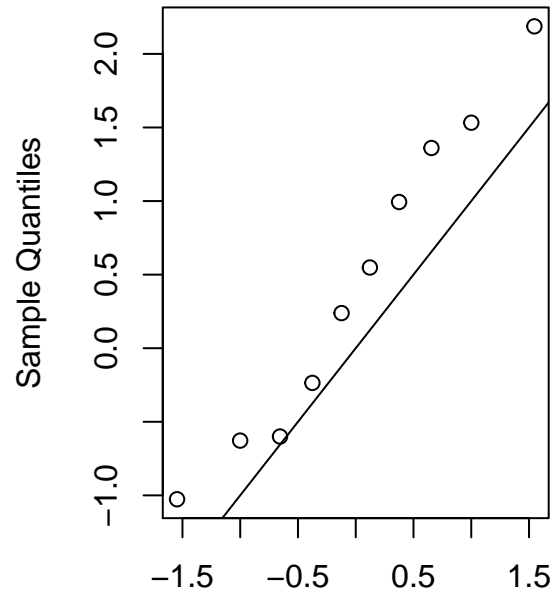
**Normal Q-Q Plot**



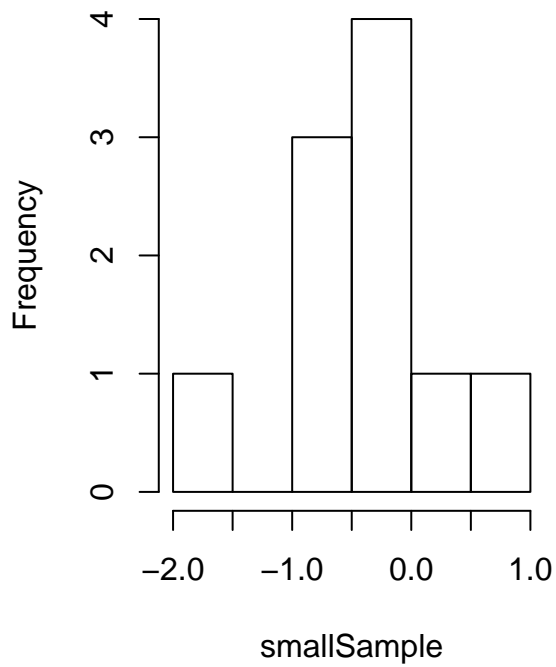
**Histogram of smallSample**



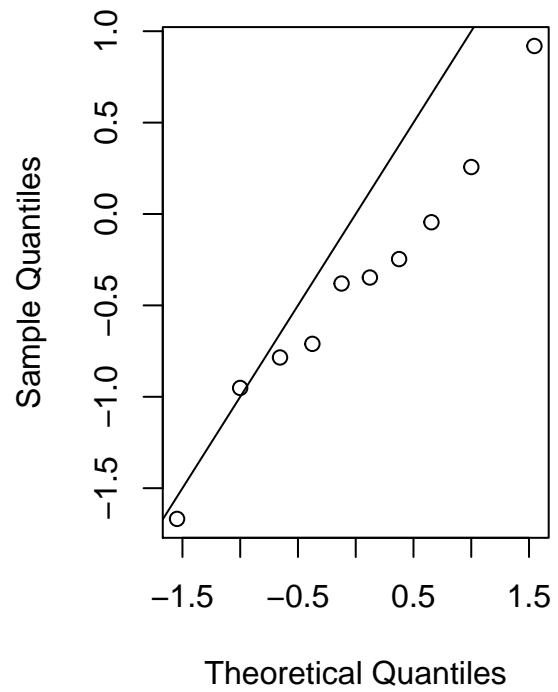
**Normal Q-Q Plot**



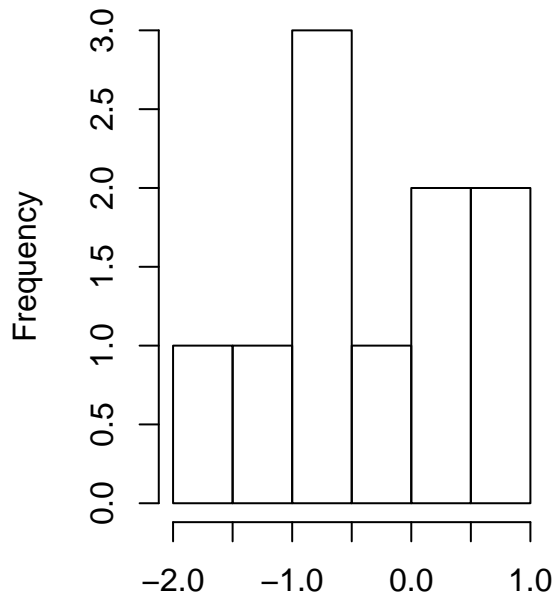
smallSample  
**Histogram of smallSample**



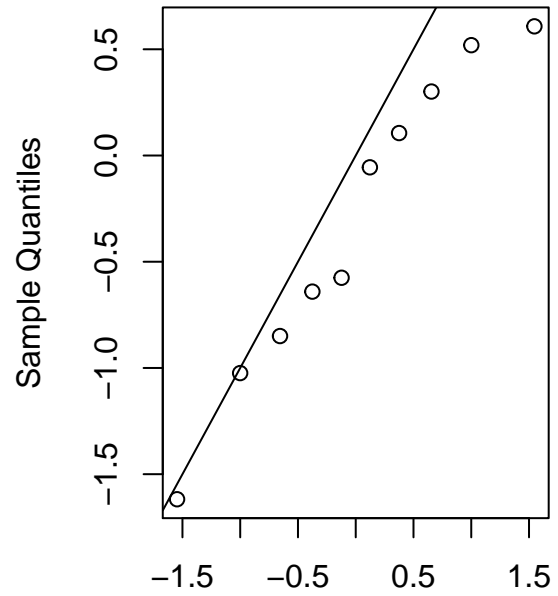
Theoretical Quantiles  
**Normal Q-Q Plot**



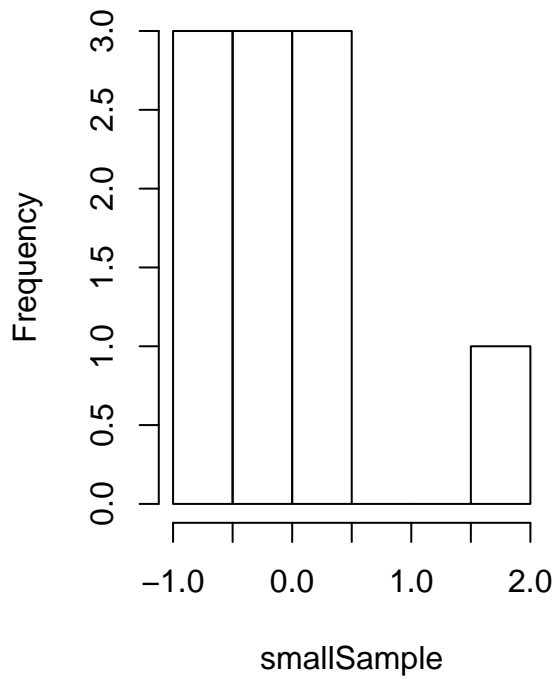
**Histogram of smallSample**



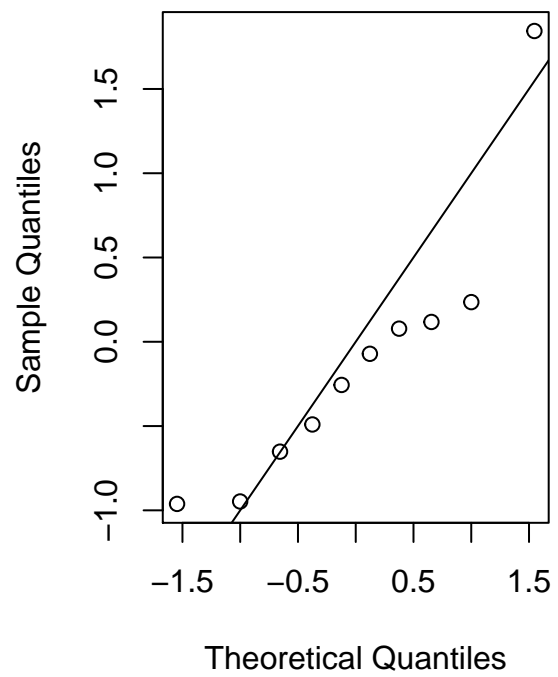
**Normal Q-Q Plot**



smallSample  
**Histogram of smallSample**

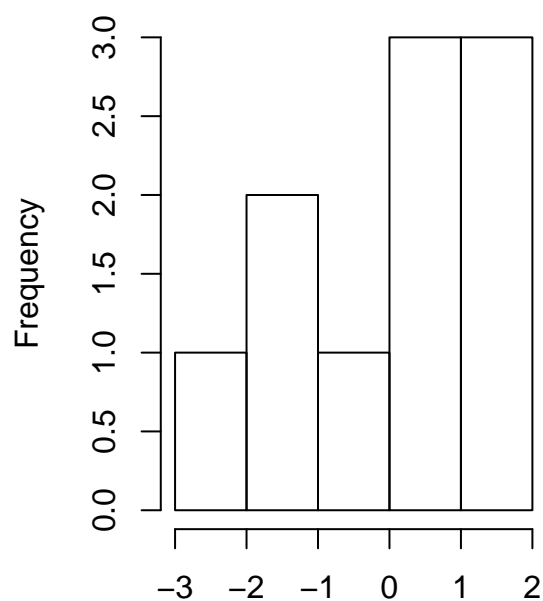


Theoretical Quantiles  
**Normal Q-Q Plot**

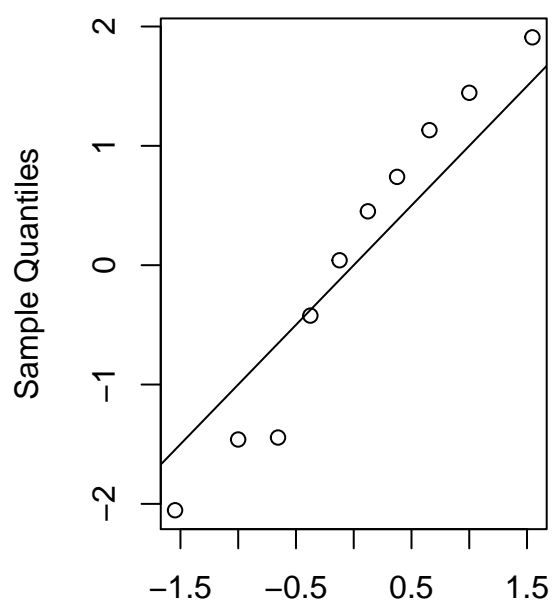




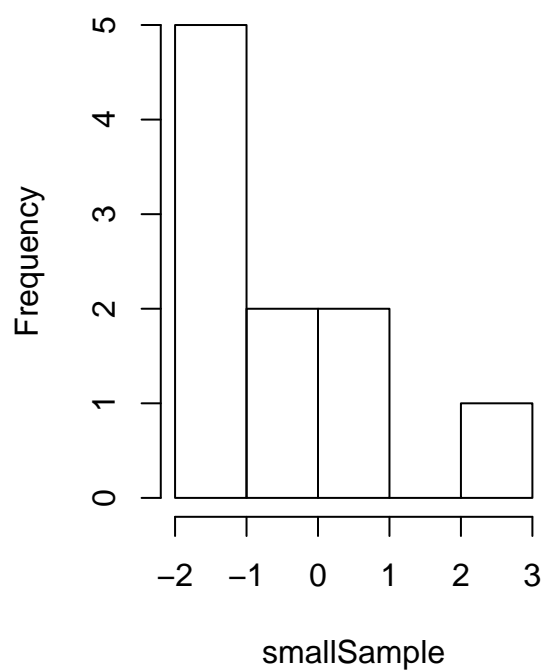
### Histogram of smallSample



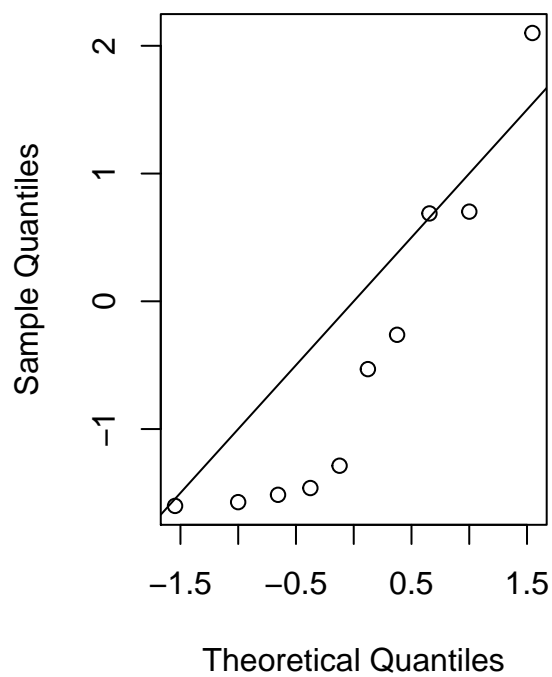
### Normal Q-Q Plot



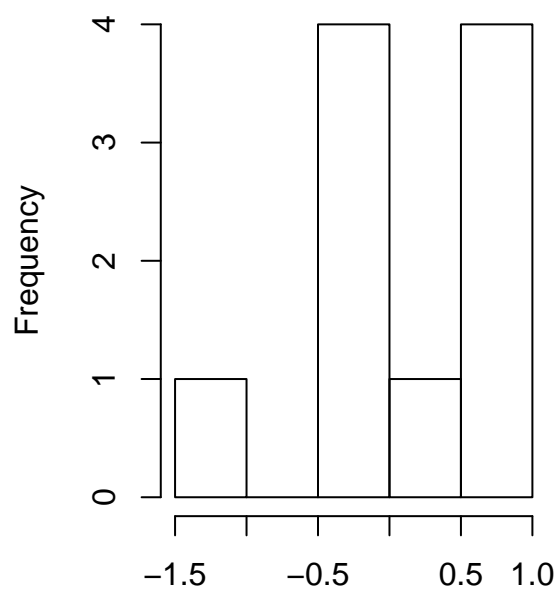
### Histogram of smallSample



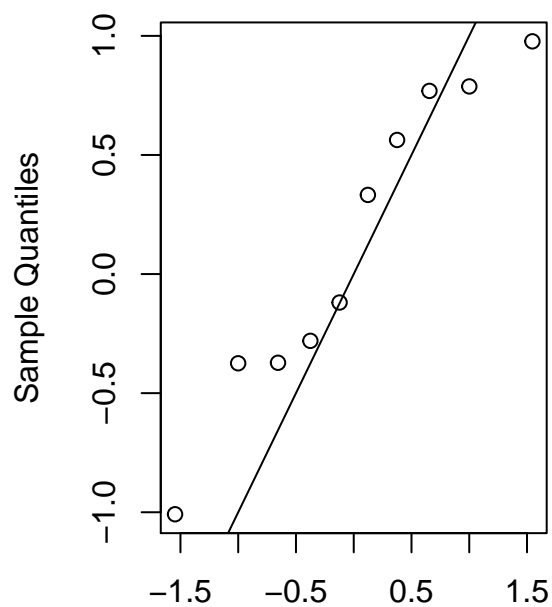
### Normal Q-Q Plot



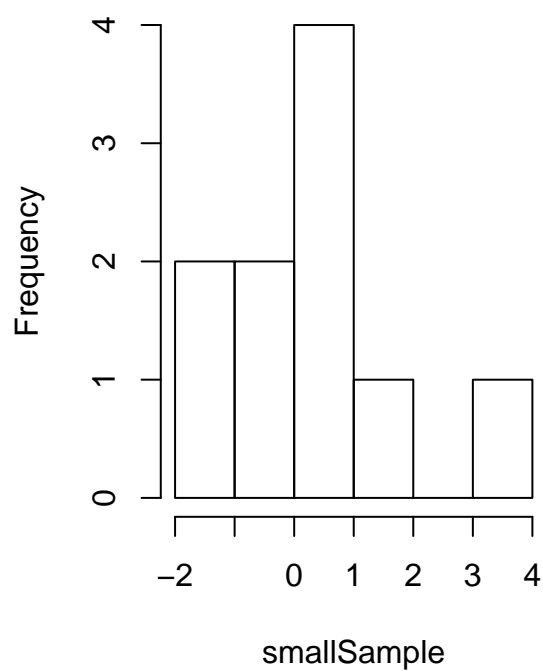
### Histogram of smallSample



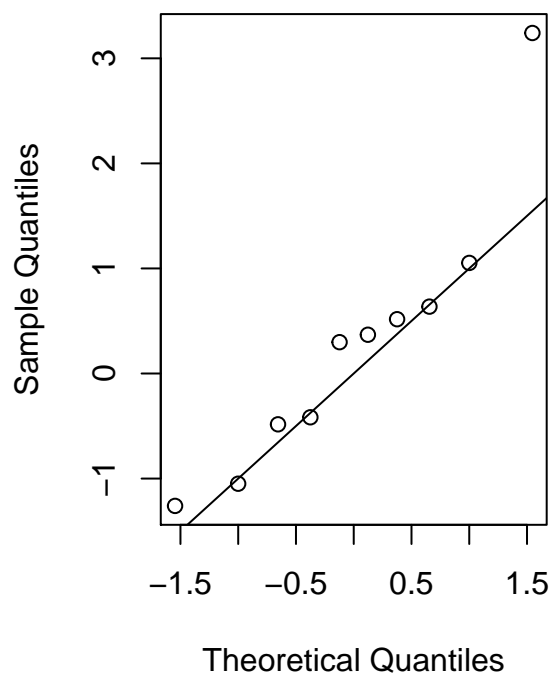
### Normal Q-Q Plot



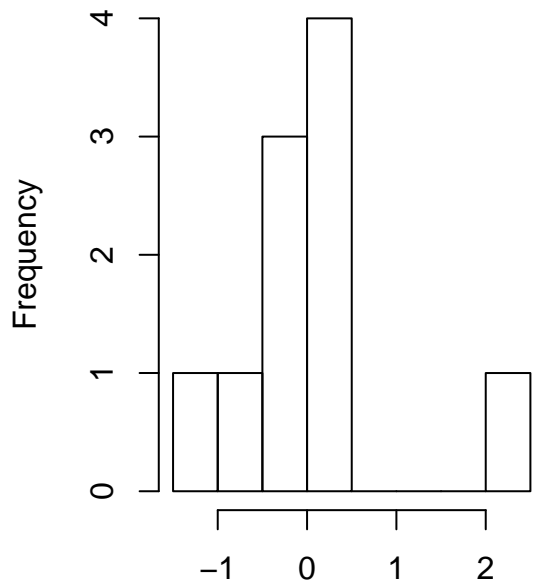
### Histogram of smallSample



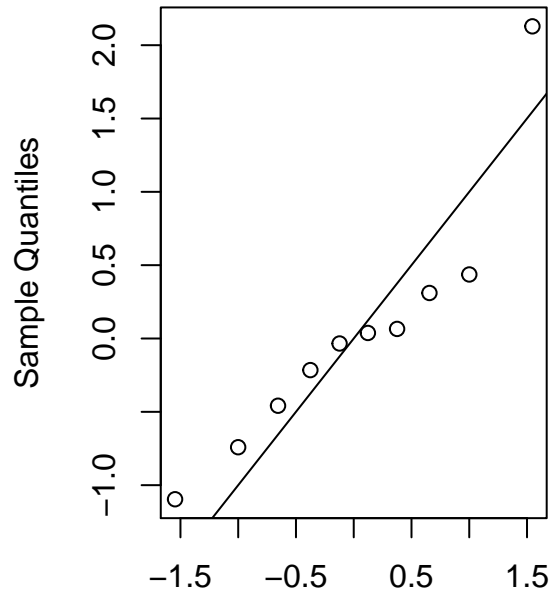
### Normal Q-Q Plot



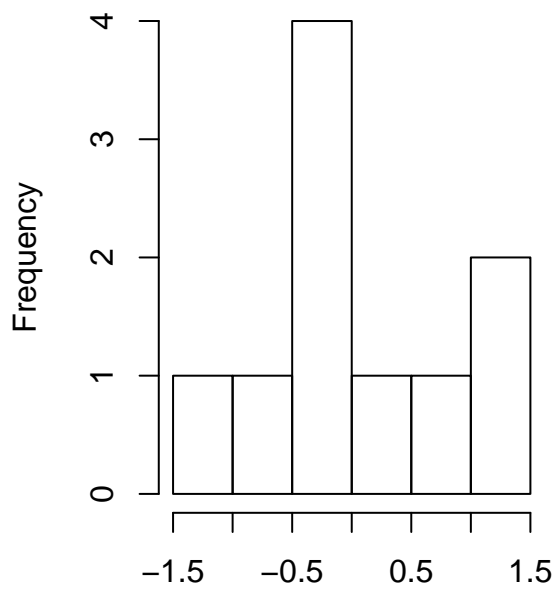
**Histogram of smallSample**



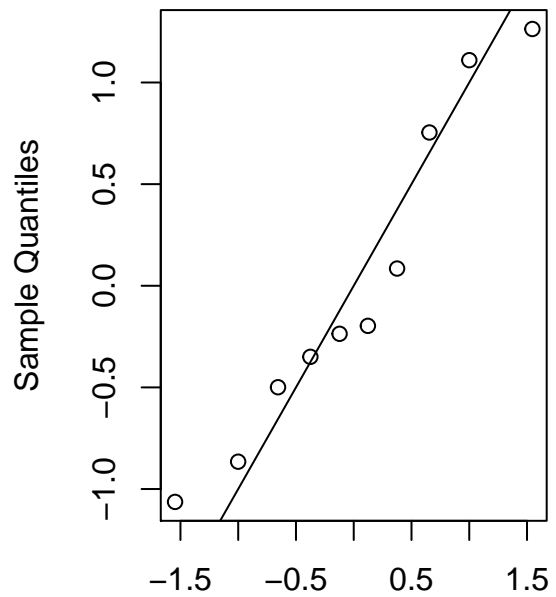
**Normal Q-Q Plot**



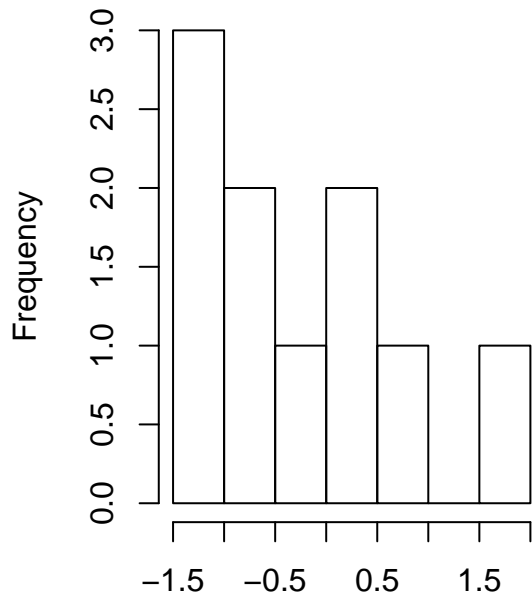
smallSample  
**Histogram of smallSample**



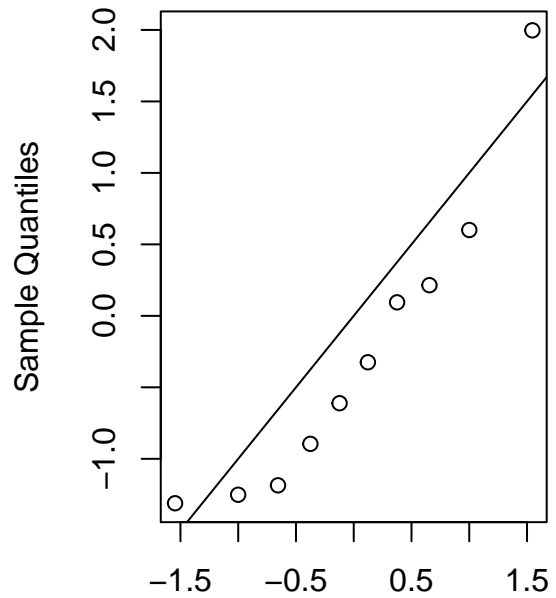
Theoretical Quantiles  
**Normal Q-Q Plot**



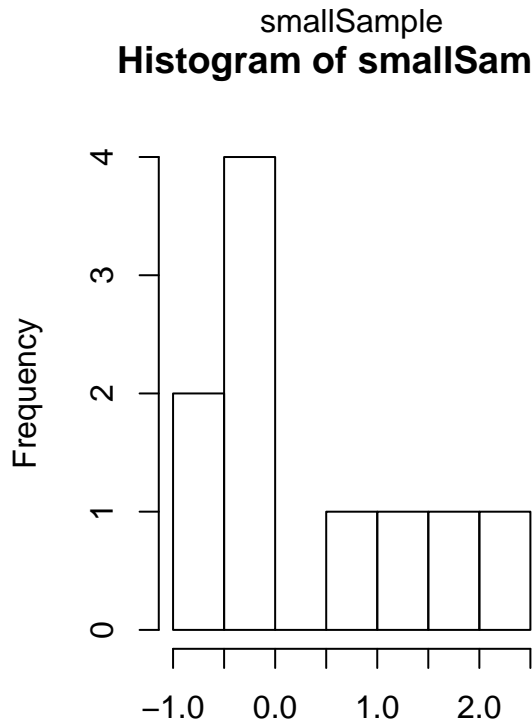
**Histogram of smallSample**



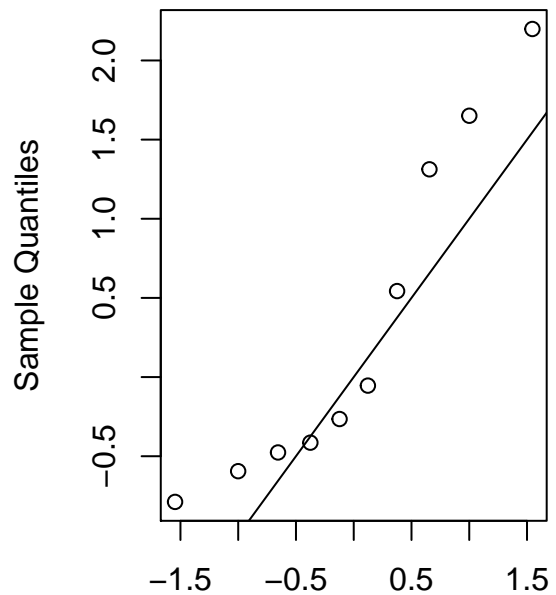
**Normal Q-Q Plot**



**Histogram of smallSample**



**Normal Q-Q Plot**

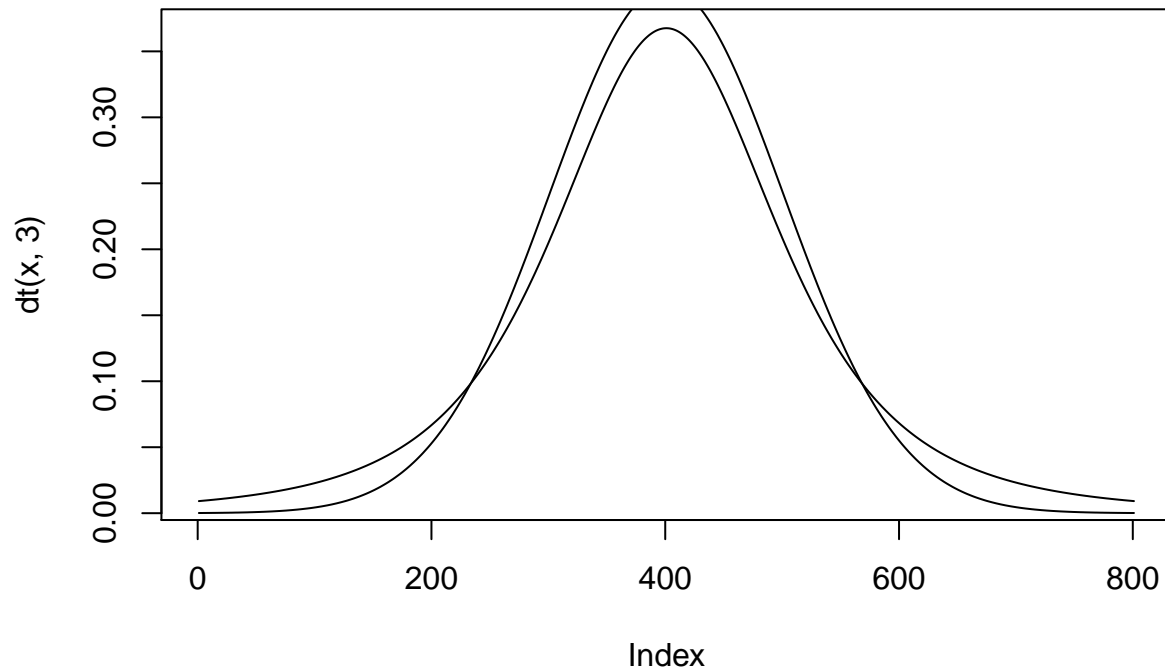


After generating 20 separate instances, we see we were a bit lucky with the qqplot and it doesn't really come up to snuff.

Problem 2

```
x = seq(-4,4,.01)
plot(dt(x,3),type='l')
```

```
lines(dnorm(x,0,1),type='l')
```

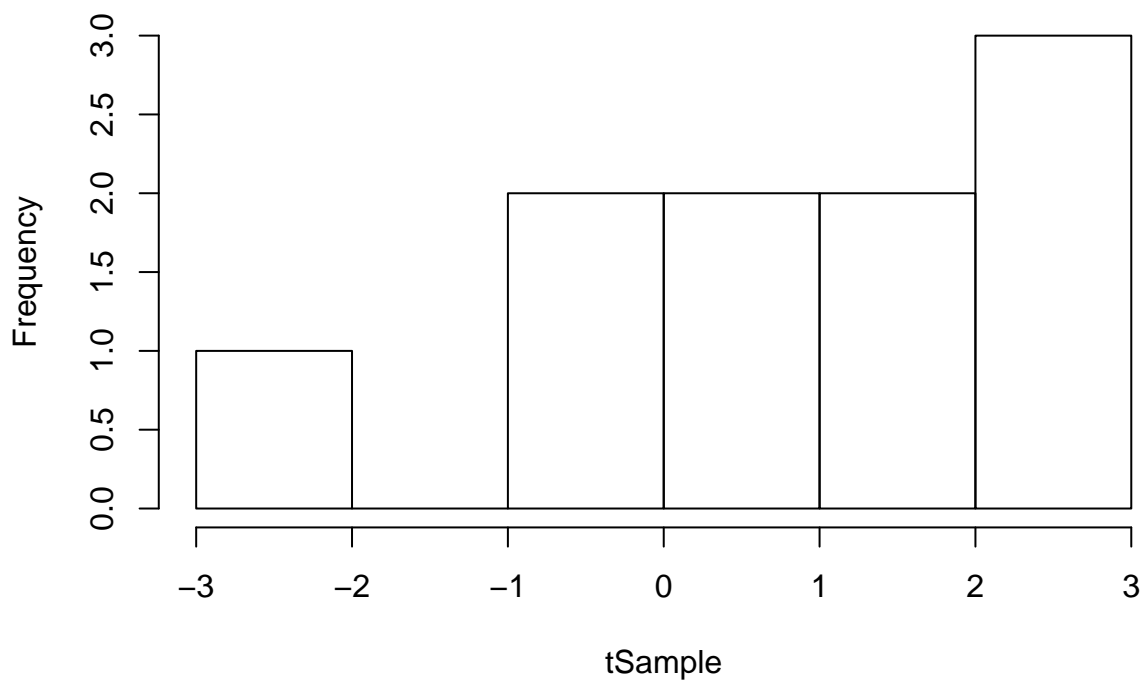


can see, there is a heavier tail and smaller and flatter peak for the t distribution

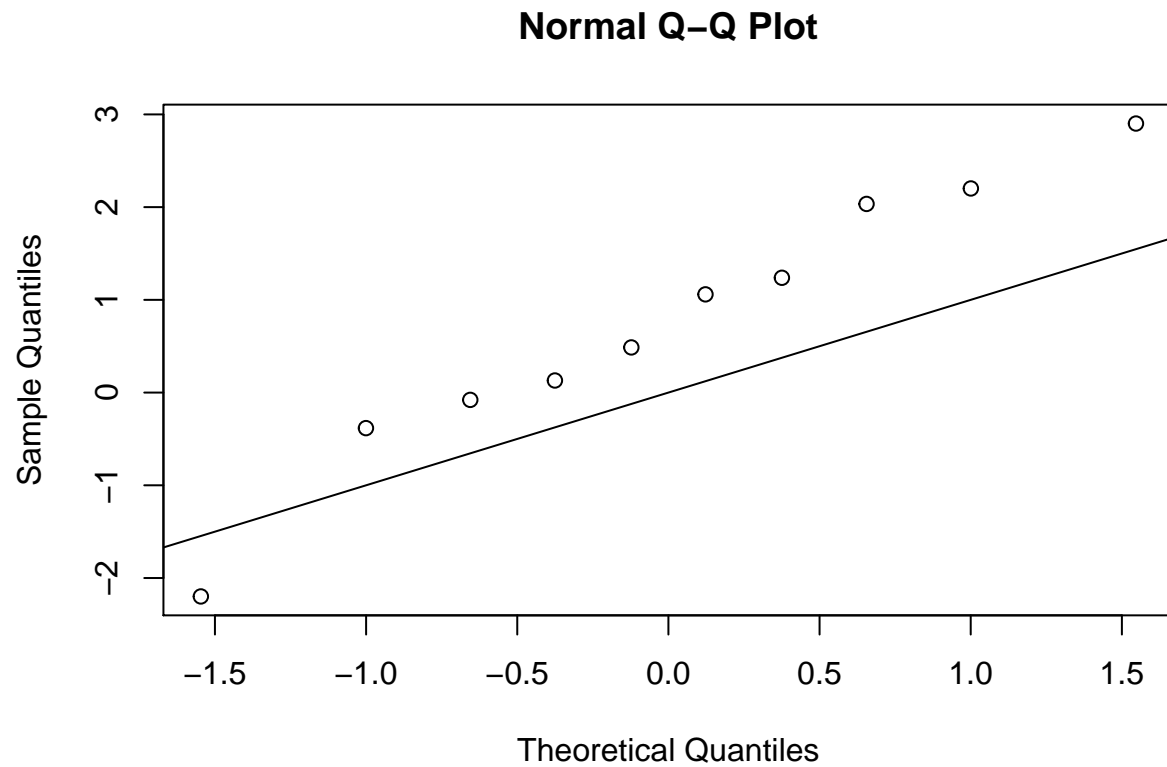
Problem 3

```
tSample = rt(10,3)  
hist(tSample)
```

**Histogram of tSample**



```
qqnorm(tSample)
abline(0,1)
```

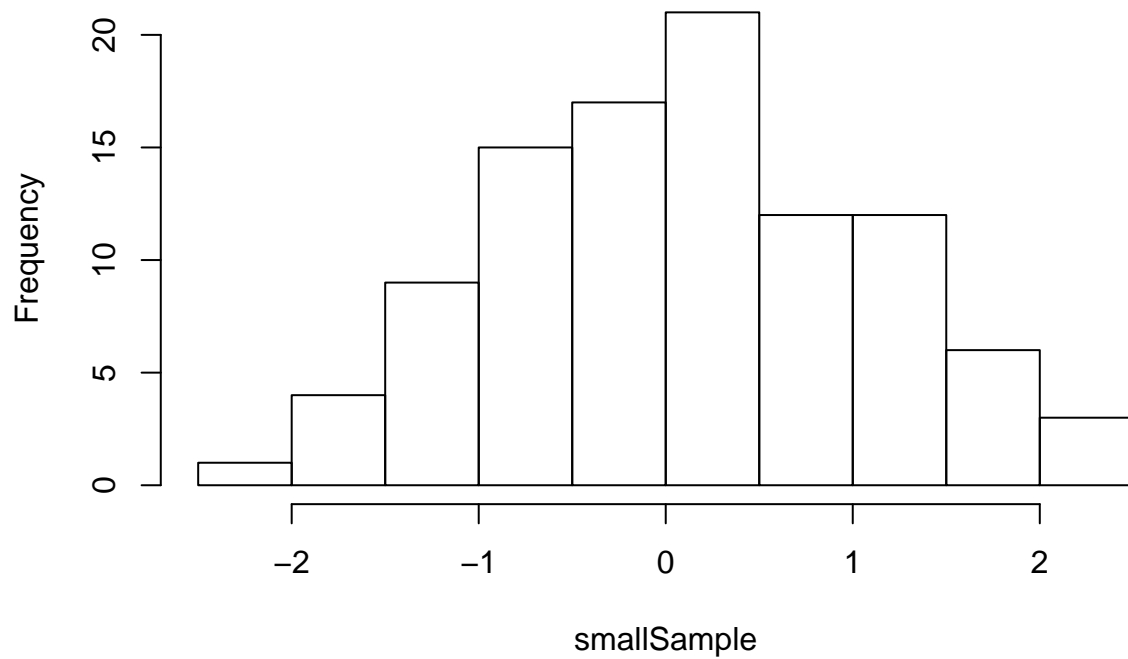


Surprisingly, it looks pretty normal via the qqplot, but not in the histogram, just like before

Problem 4

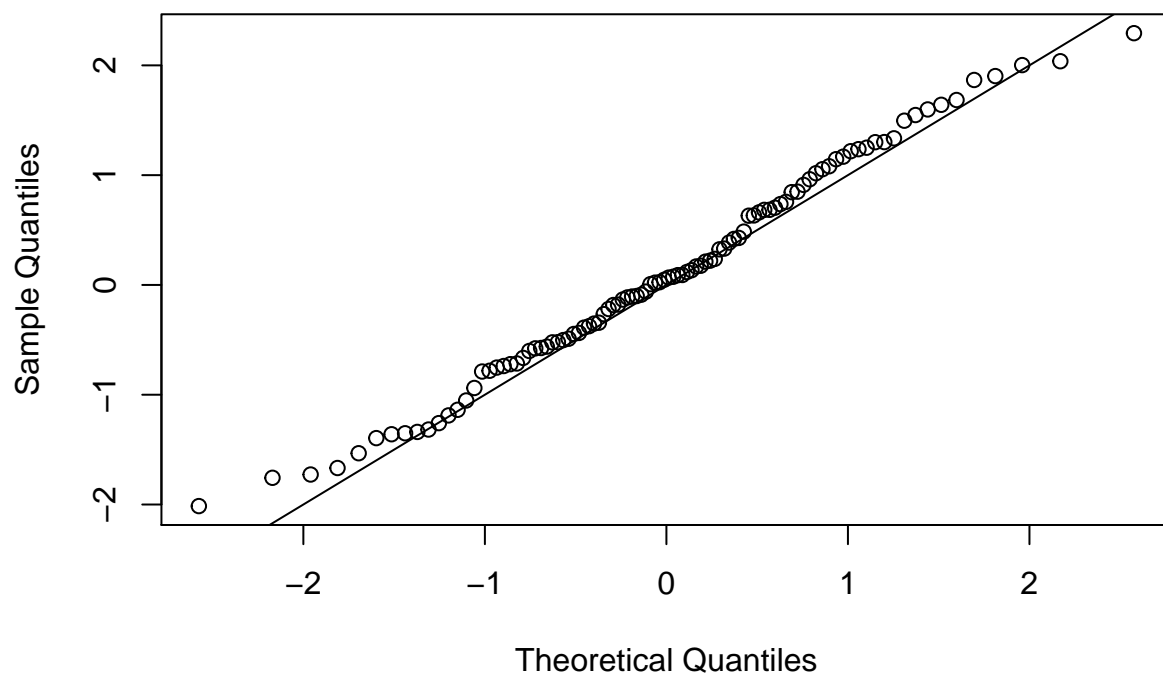
```
n=100
smallSample = (rnorm(n,0,1))
tSample = rt(n,3)
hist(smallSample)
```

### Histogram of smallSample



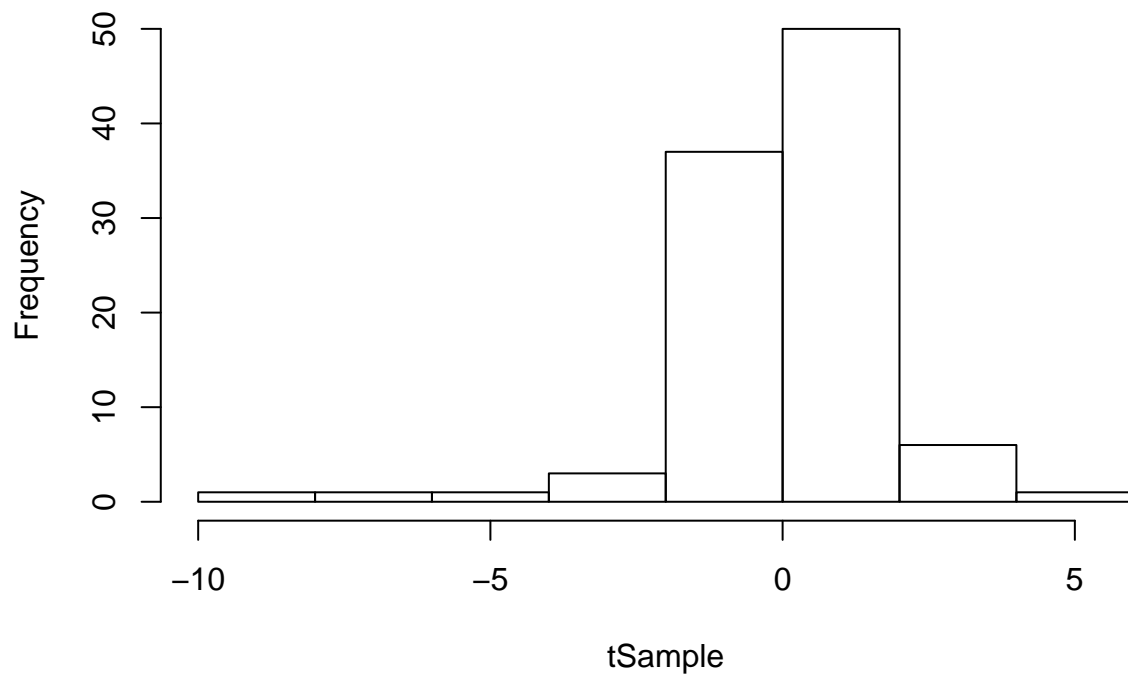
```
qqnorm(smallSample)  
abline(0,1)
```

### Normal Q-Q Plot



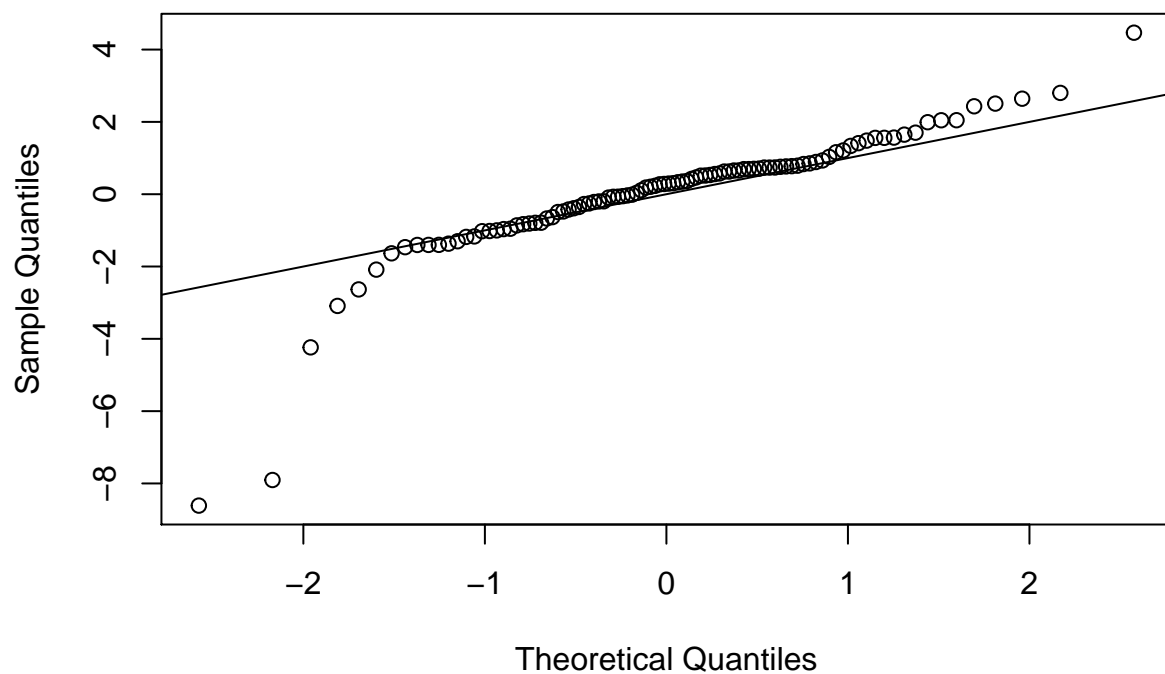
```
hist(tSample)
```

**Histogram of tSample**



```
qqnorm(tSample)  
abline(0,1)
```

**Normal Q-Q Plot**



We seem to have found a normal looking noramally generated distrib, but not very true to the actual curve is the

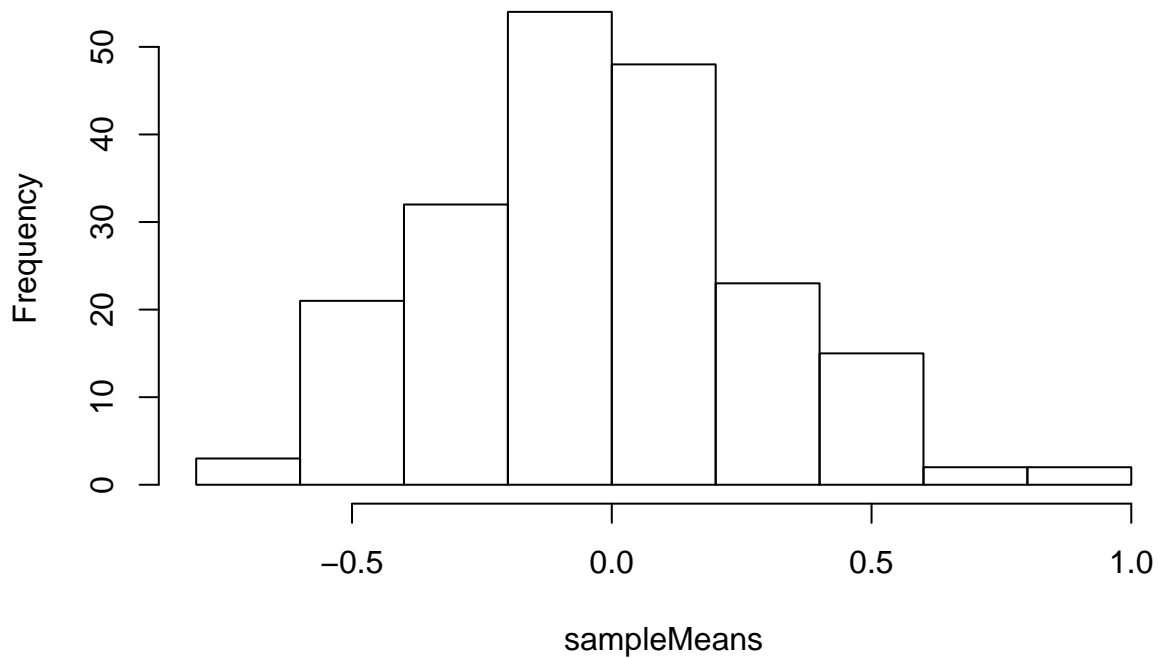


t distributed sample.

Problem 5

```
sampleMeans = numeric(200)
sampleSize = 10
for(i in 1:200)
{
  sampleMeans[i]=mean(rnorm(sampleSize))
}
hist(sampleMeans)
```

**Histogram of sampleMeans**



```
print(mean(sampleMeans))
```

```
## [1] -0.02552663
```

```
print((var(sampleMeans)))
```

```
## [1] 0.09160695
```

```
print(sqrt(var(sampleMeans)))
```

```
## [1] 0.3026664
```

```
print(1/sqrt(sampleSize))
```

```
## [1] 0.3162278
```

Problem 6

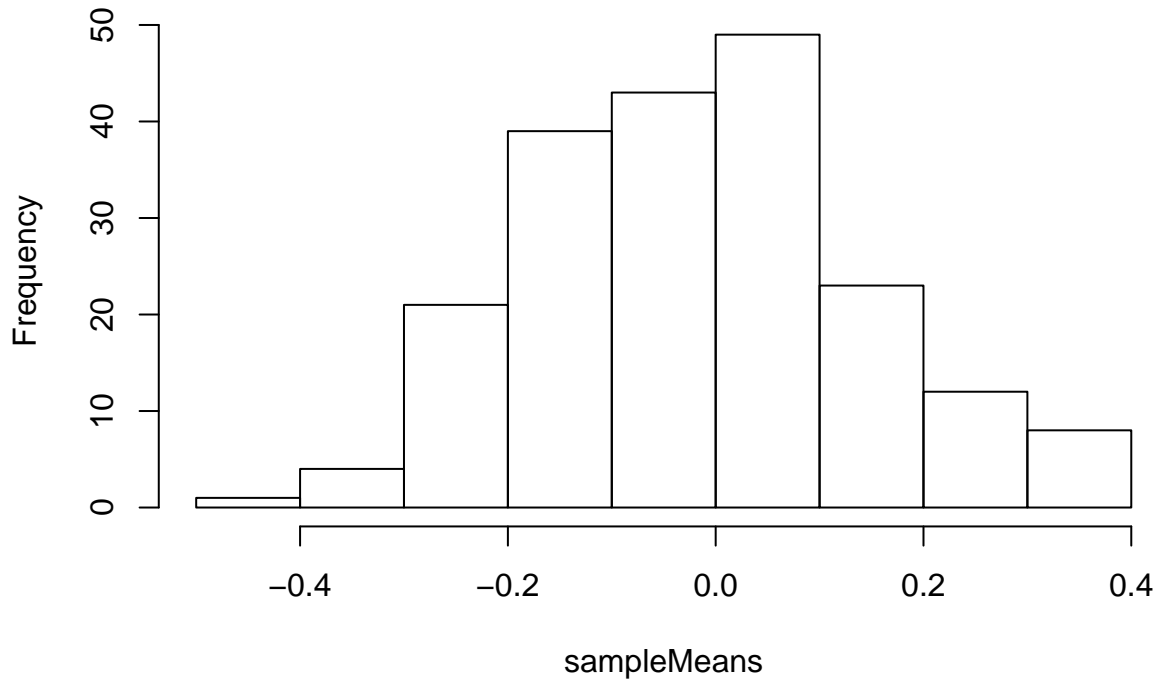
```
sampleMeans = numeric(200)
sampleSize = 40
for(i in 1:200)
{
```

```

sampleMeans[i]=mean(rnorm(sampleSize))
}
hist(sampleMeans)

```

## Histogram of sampleMeans



```
print(mean(sampleMeans))
```

```
## [1] -0.01851286
```

```
print((var(sampleMeans)))
```

```
## [1] 0.02508546
```

```
print(sqrt(var(sampleMeans)))
```

```
## [1] 0.1583839
```

```
print(1/sqrt(sampleSize))
```

```
## [1] 0.1581139
```

Problem 7

```
print(1/.01)
```

```
## [1] 100
```

by dividing the true variance by the standard error we desire, we see that 100 sample size is needed theoretically to achieve it.

Problem 8

```
bernoulSample = rbinom(10,5,0.1)
```

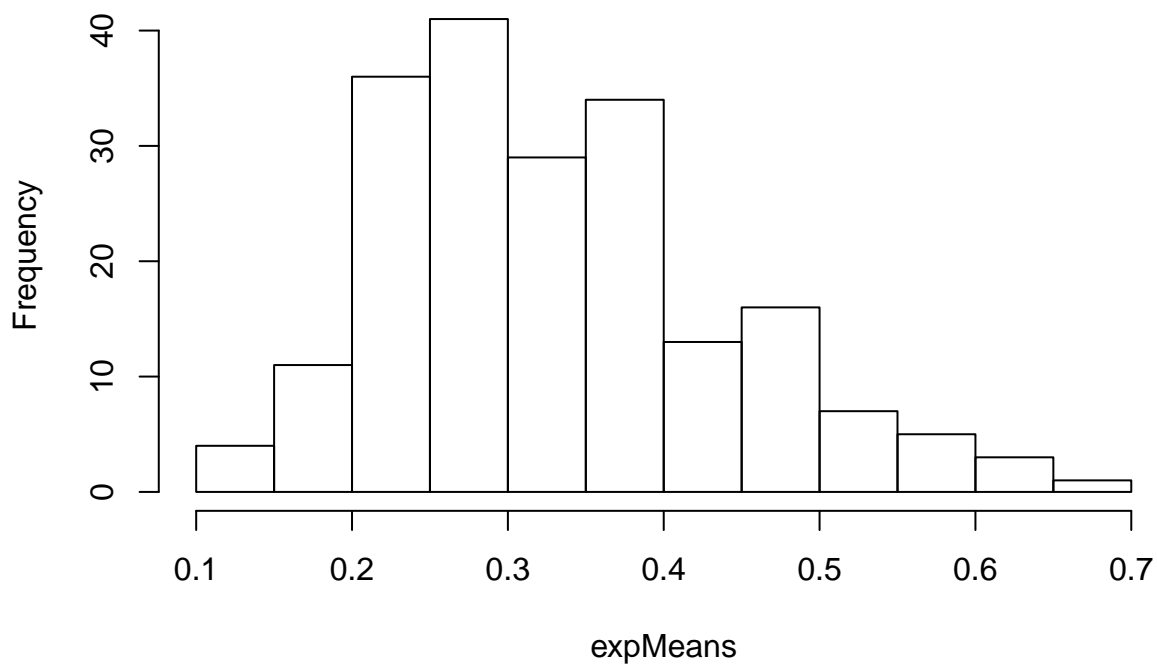
```
print((.1*.9)/(0.01**2))
```

```
## [1] 900
```

In the above, we knew the standard error, which is the SD of the sample proportion, is 0.01, and then knew the true variance follows  $p \cdot q$ . We simply invert, square, and solve for  $n$  Problem 9

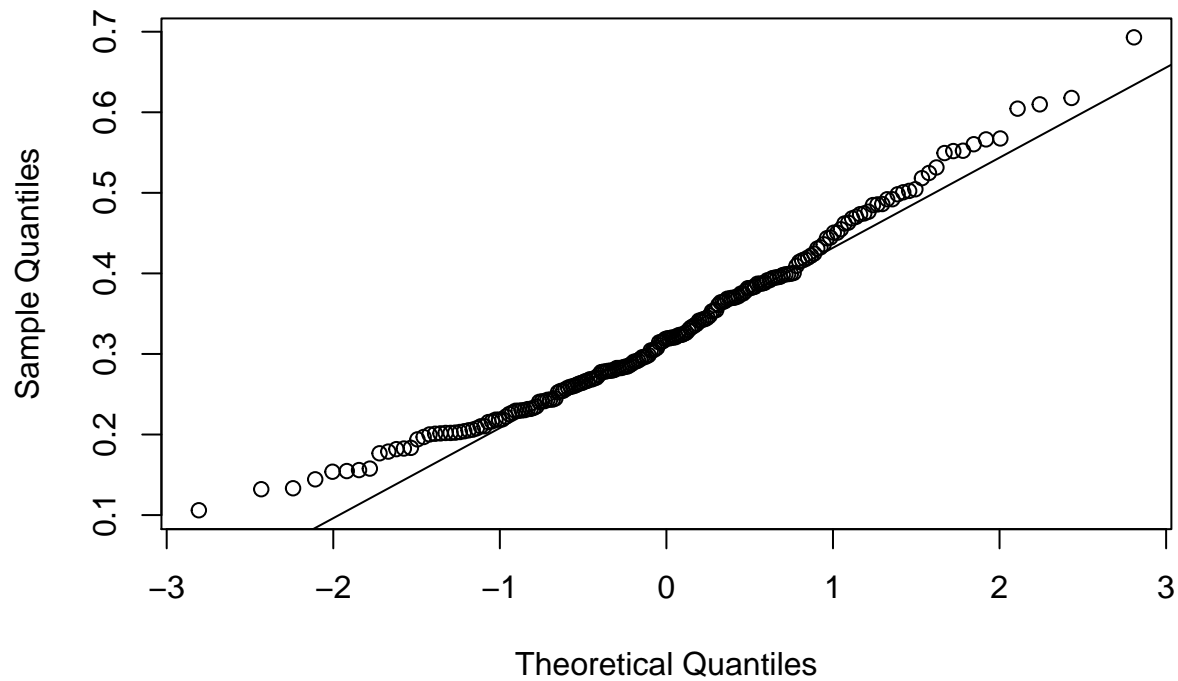
```
expMeans = numeric(200)
for(i in 1:200)
{
  expMeans[i] = mean(rexp(10,3))
}
hist(expMeans)
```

**Histogram of expMeans**



```
qqnorm(expMeans)
qqline(expMeans)
```

## Normal Q-Q Plot



problem 10

```
print(mean(expMeans))
```

```
## [1] 0.3304988
```

```
print(var(expMeans))
```

```
## [1] 0.01217654
```

```
print(sqrt(var(expMeans)))
```

```
## [1] 0.1103474
```

```
print("versus theoreticals")
```

```
## [1] "versus theoreticals"
```

```
print((1/3))
```

```
## [1] 0.3333333
```

```
print((1/9)/10)
```

```
## [1] 0.01111111
```

```
print(sqrt((1/9)/10))
```

```
## [1] 0.1054093
```

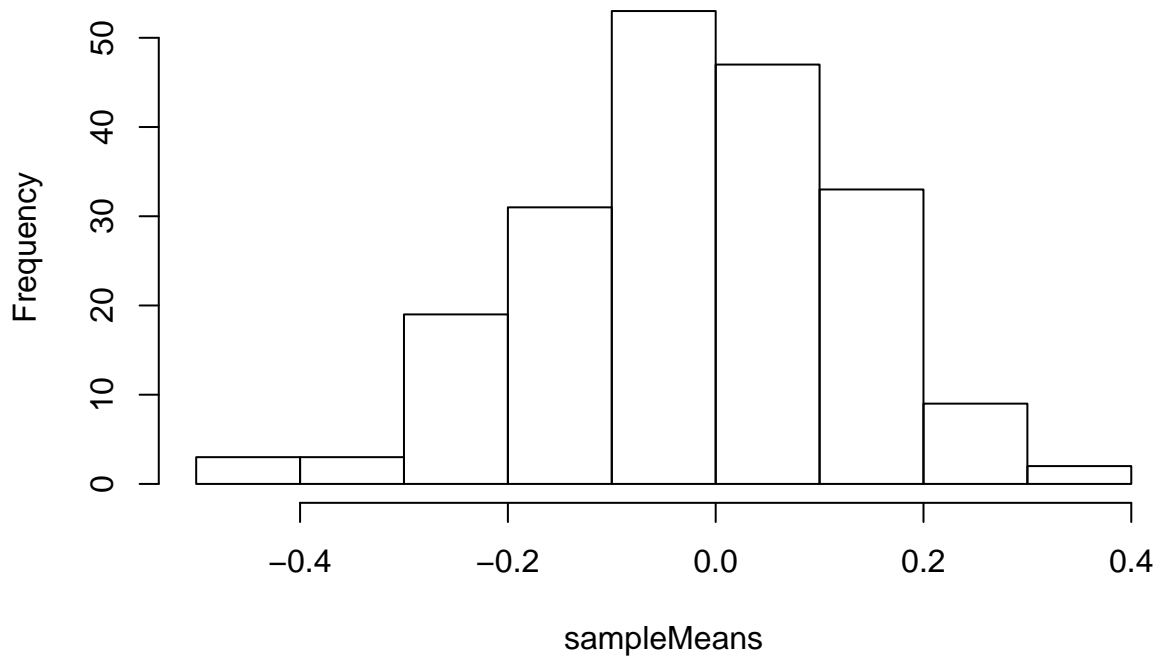
They are pretty close!

problem 11

```
print("problem number 5 repeated")
```

```
## [1] "problem number 5 repeated"
sampleMeans = numeric(200)
sampleSize = 40
for(i in 1:200)
{
  sampleMeans[i]=mean(rnorm(sampleSize))
}
hist(sampleMeans)
```

**Histogram of sampleMeans**



```
print(mean(sampleMeans))
```

```
## [1] -0.02022599
```

```
print((var(sampleMeans)))
```

```
## [1] 0.02224429
```

```
print("versus theoreticals")
```

```
## [1] "versus theoreticals"
```

```
print(sqrt(var(sampleMeans)))
```

```
## [1] 0.1491452
```

```
print(1/sqrt(sampleSize))
```

```
## [1] 0.1581139
```

```
problem 12
```

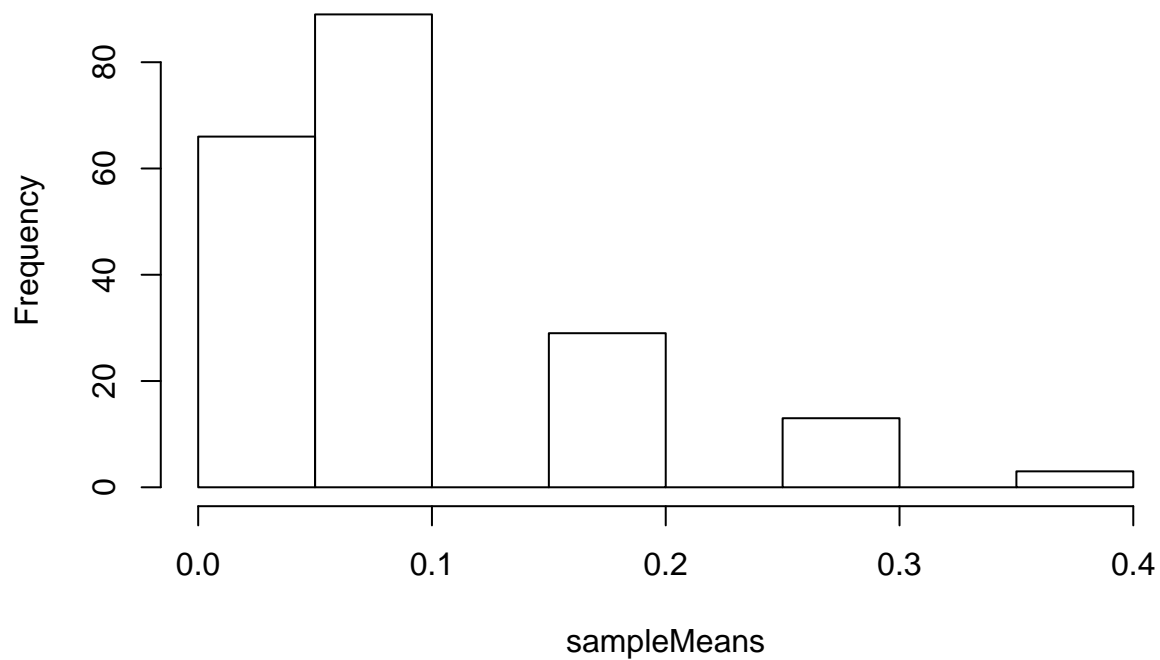
```
sampleMeans = numeric(200)
sampleSize = 10
```

```

for(i in 1:200)
{
  sampleMeans[i]=mean(rbinom(sampleSize,1,0.1))
}
hist(sampleMeans)

```

**Histogram of sampleMeans**

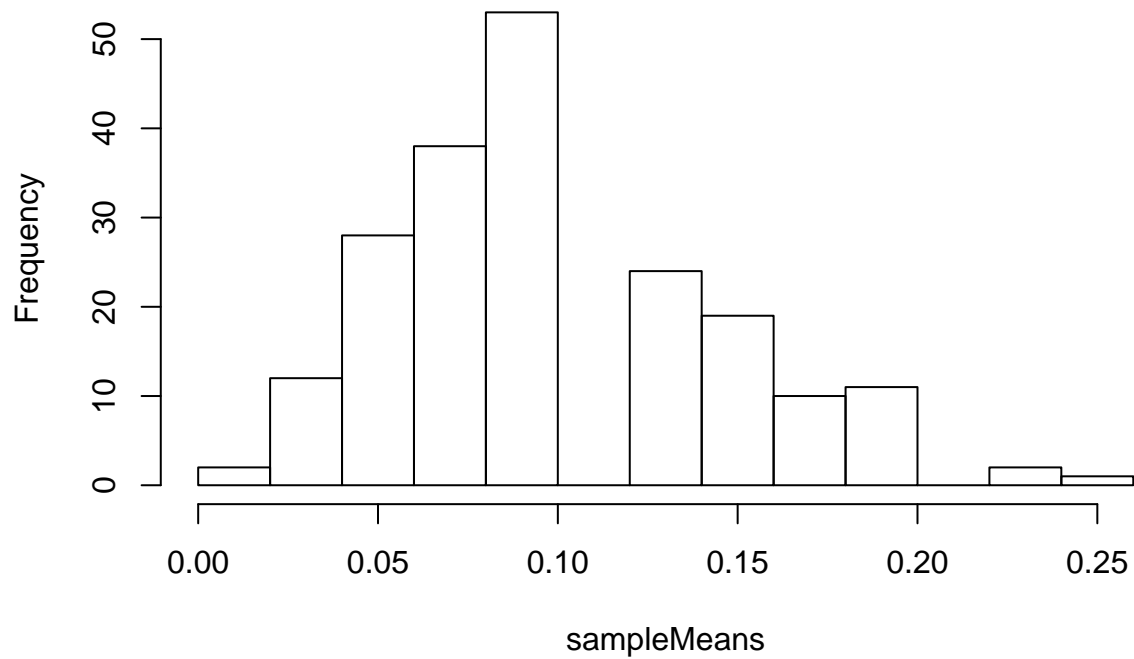


```

sampleMeans = numeric(200)
sampleSize = 40
for(i in 1:200)
{
  sampleMeans[i]=mean(rbinom(n=sampleSize,size=1,prob=0.1))
}
hist(sampleMeans)

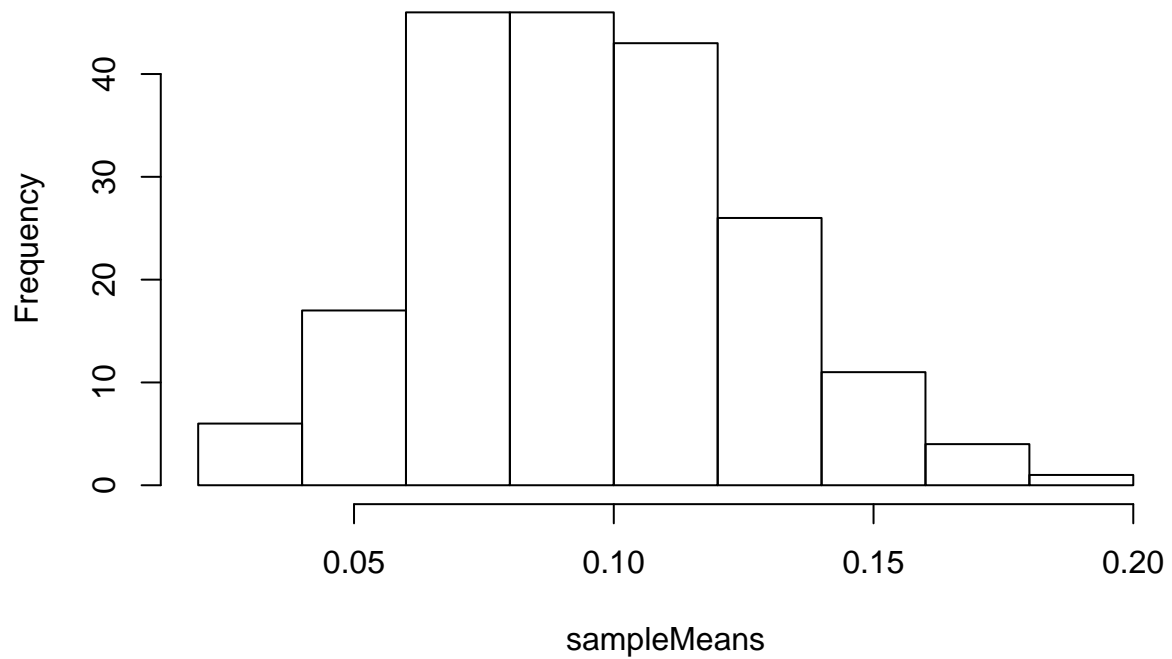
```

## Histogram of sampleMeans



```
sampleMeans = numeric(200)
sampleSize = 100
for(i in 1:200)
{
  sampleMeans[i]=mean(rbinom(n=sampleSize,size=1,prob=0.1))
}
hist(sampleMeans)
```

## Histogram of sampleMeans

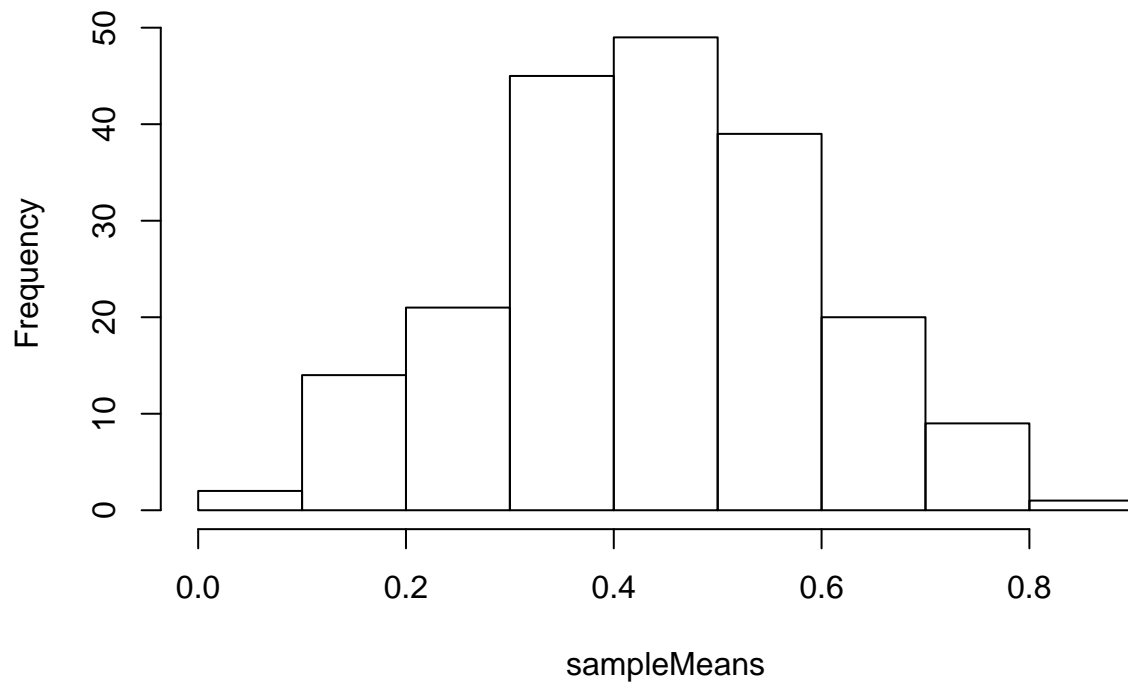


Problem 13

```
sampleMeans = numeric(200)
sampleSize = 10
for(i in 1:200)
{
  sampleMeans[i]=mean(rbinom(sampleSize,1,0.5))
}
hist(sampleMeans)
```

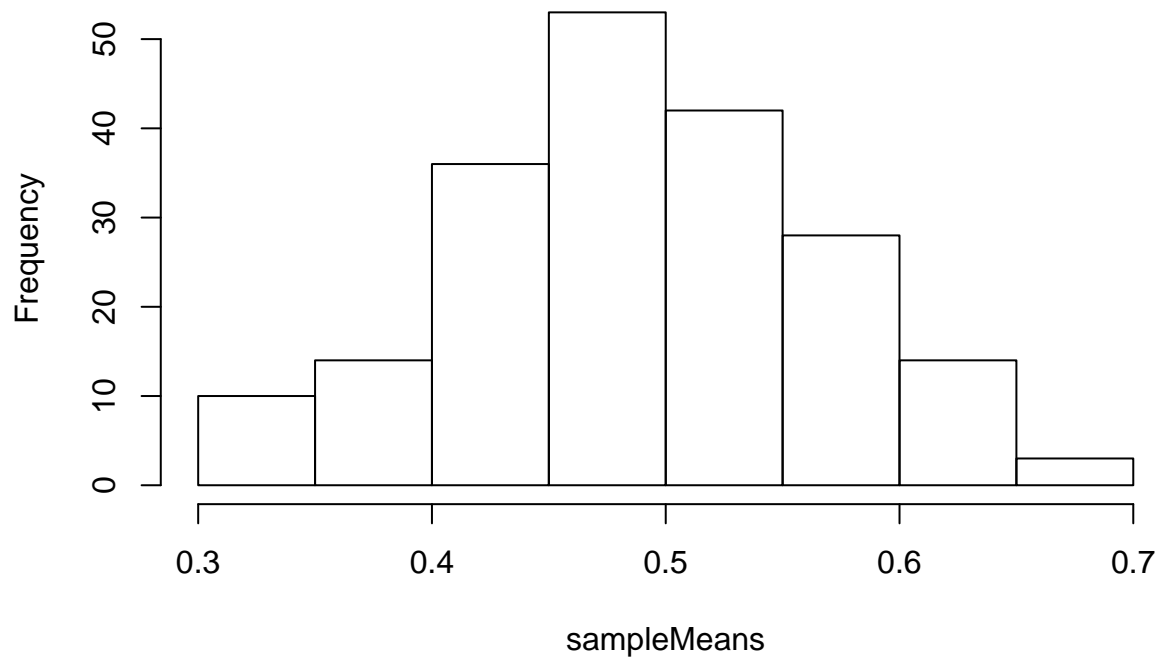


## Histogram of sampleMeans



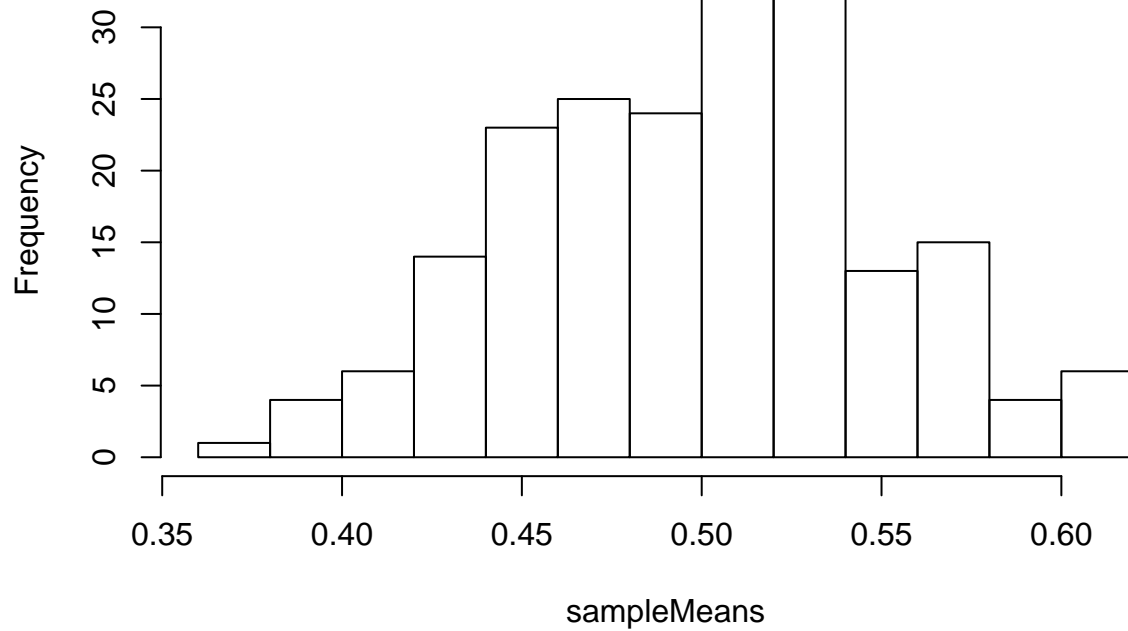
```
sampleMeans = numeric(200)
sampleSize = 40
for(i in 1:200)
{
  sampleMeans[i]=mean(rbinom(n=sampleSize,size=1,prob=0.5))
}
hist(sampleMeans)
```

## Histogram of sampleMeans



```
sampleMeans = numeric(200)
sampleSize = 100
for(i in 1:200)
{
  sampleMeans[i]=mean(rbinom(n=sampleSize,size=1,prob=0.5))
}
hist(sampleMeans)
```

## Histogram of sampleMeans



Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.