

Lab 1 Probability Distributions

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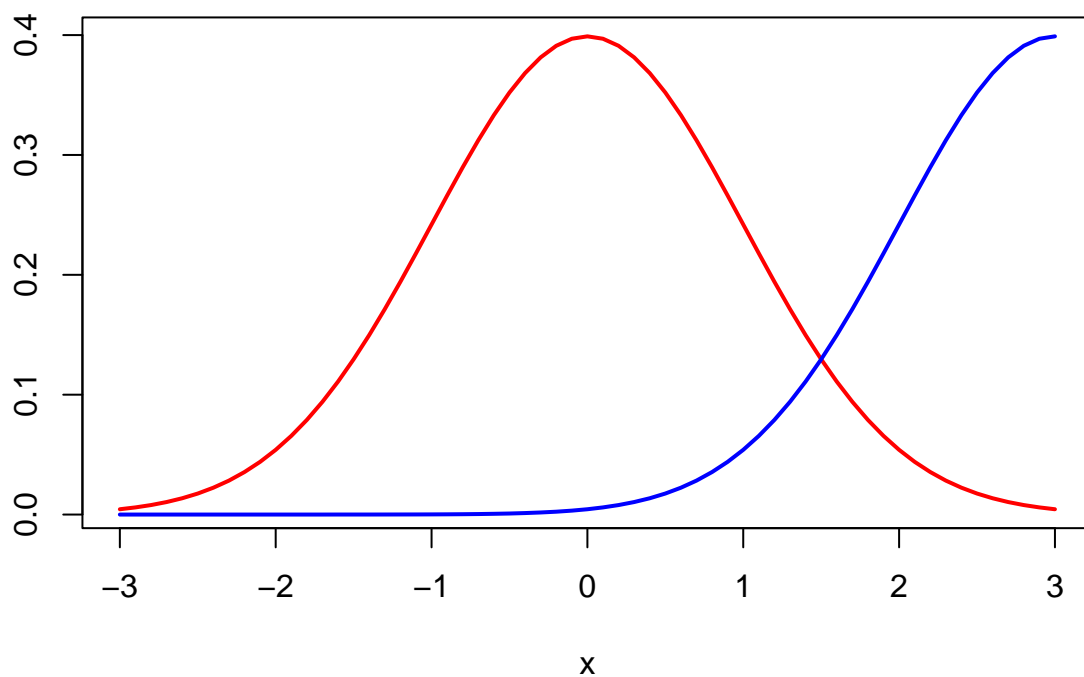
Lab 1 Lab Manual Exercise

copy and paste your work by following each example from the lab manual for this exercise

```
rm(list = setdiff(ls(), lsf.str()))

# Plot Normal Distributions with
#-----
# Same standard deviation, different mean
#-----
# Mean 1, sd 1
# Grid of X-axis values
x <- seq(-3, 3, 0.1)

plot(x, dnorm(x, mean = 0, sd = 1), type = "l",
      ylab = "", lwd = 2, col = "red")
# Mean 3, sd 1
lines(x, dnorm(x, mean = 3, sd = 1), col = "blue", lty = 1, lwd = 2)
```



```
# # Function Syntax
#
# function_name <- function(arg_1, arg_2, ...) {
#   Function body
# }
```

```
# Calculate the 60th %ile of the standard normal.
qnorm(0.6,0,1)
```

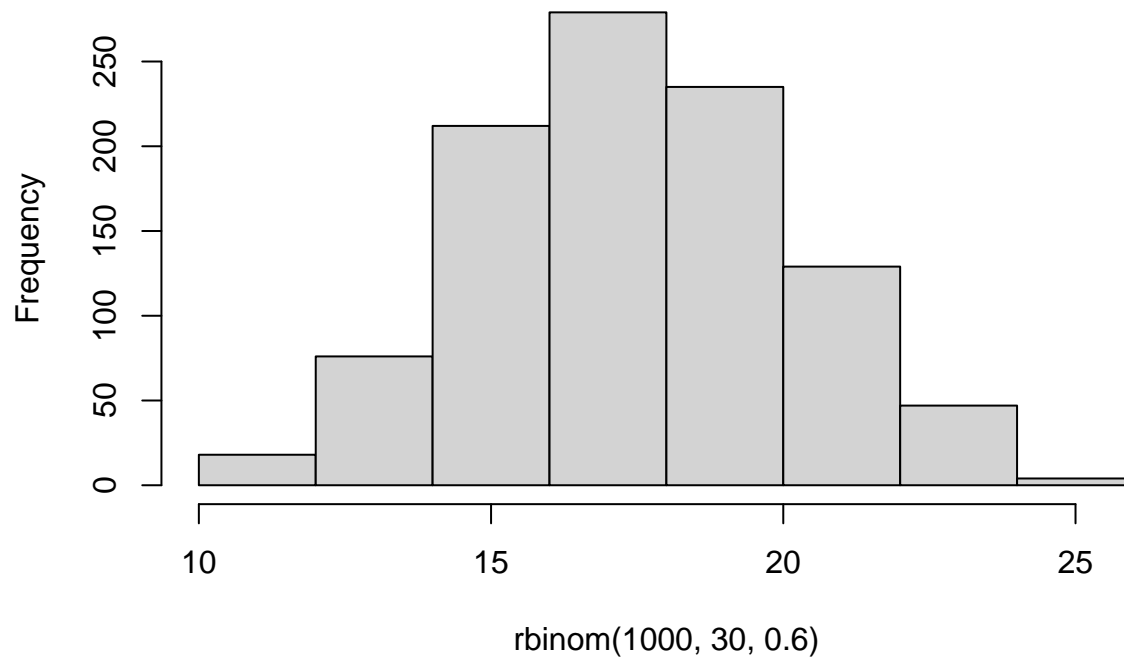
```
## [1] 0.2533471
```

```
# Calculate the probability that a value lies below 0.8 in the standard normal distribution
pnorm(0.8,0,1)
```

```
## [1] 0.7881446
```

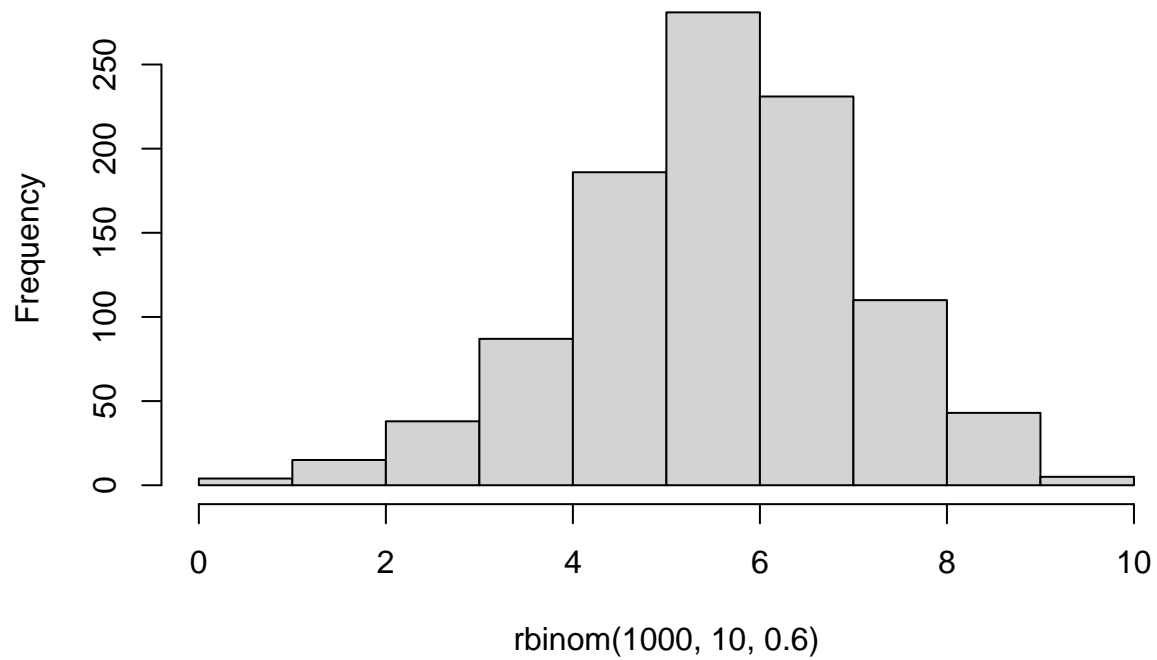
```
# Draw 1000 samples of 30 coin tosses with p(heads) = 0.6 # and plot the distribution
# Syntax: rbinom (# observations, # trials per observation, probability of success )
hist(rbinom(1000,30,0.6))
```

Histogram of rbinom(1000, 30, 0.6)



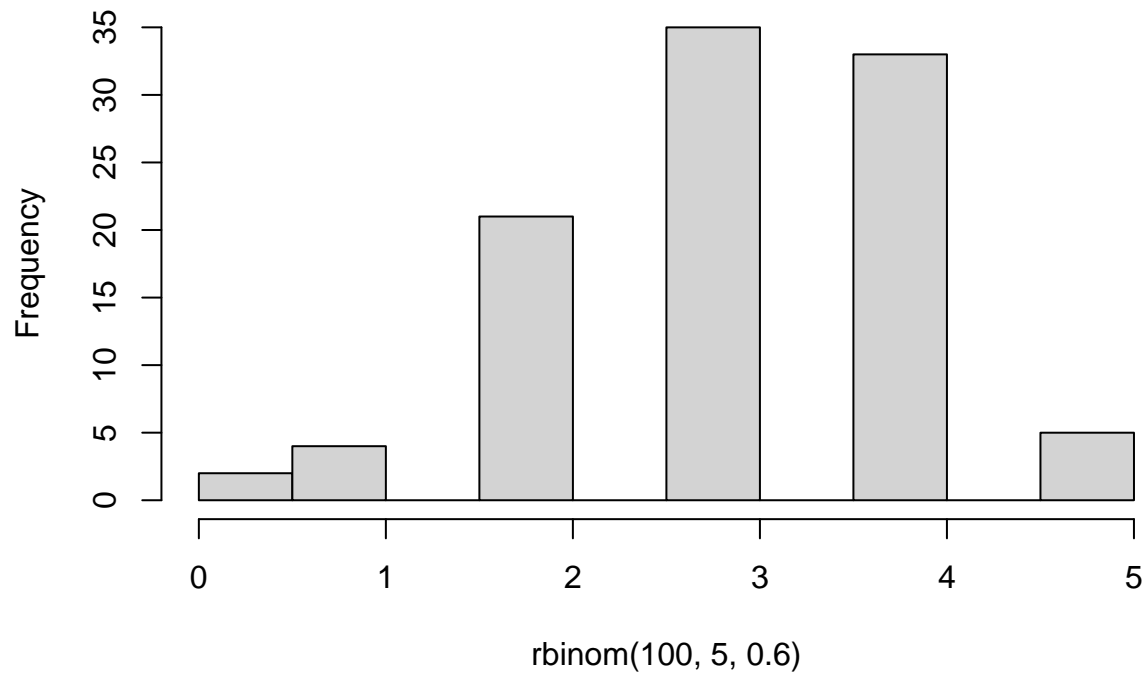
```
# Do the above with only 10 trials per observation  
hist(rbinom(1000,10,0.6))
```

Histogram of rbinom(1000, 10, 0.6)



```
# Do the above with 100 observations and 5 trials per observation  
hist(rbinom(100,5,0.6))
```

Histogram of rbinom(100, 5, 0.6)



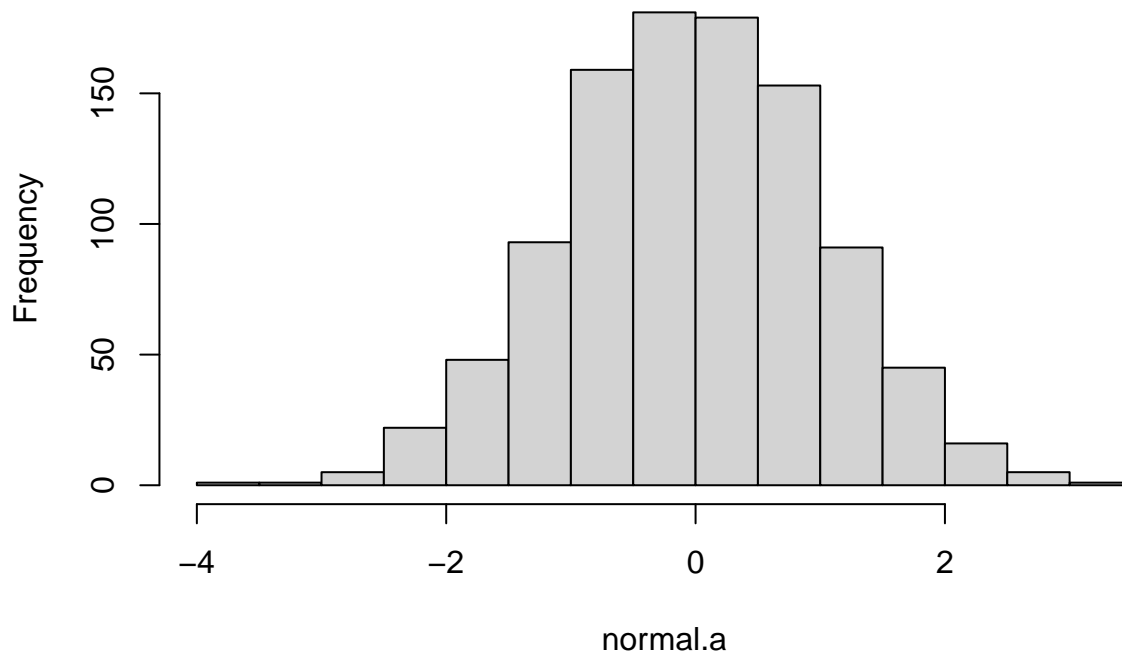
```
# Transformations between probability distributions
```

```
# generate 1000 trials from a normal distribution
```

```
normal.a <- rnorm( n=1000, mean=0, sd=1 )
```

```
hist( normal.a )
```

Histogram of normal.a



#next, we generate a chi-square distribution with 3 #degrees of freedom:

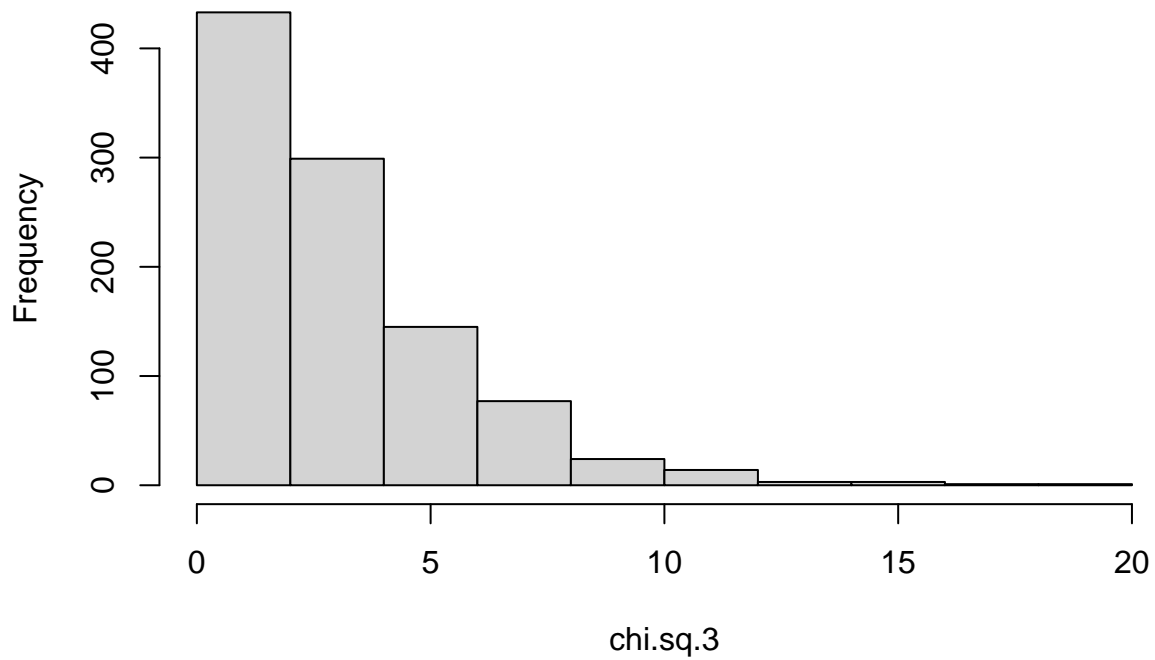
```
normal.b <- rnorm( n=1000 ) # another set of normally distributed data  
normal.c <- rnorm( n=1000 ) # and another!
```

Take the SUM of SQUARES of the above 3 normally distributed variables a, b, and c
`chi.sq.3 <- (normal.a)^2 + (normal.b)^2 + (normal.c)^2`

and the resulting chi.sq.3 variable should contain 1000 observations that follow a chi-square distrib

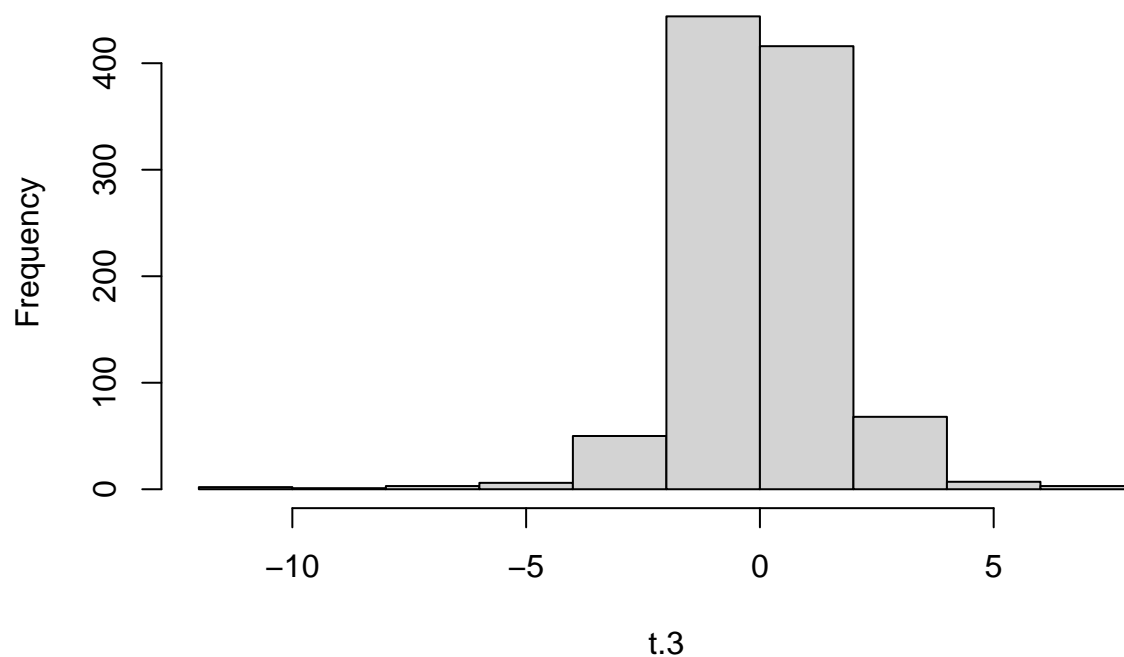
```
hist(chi.sq.3)
```

Histogram of chi.sq.3



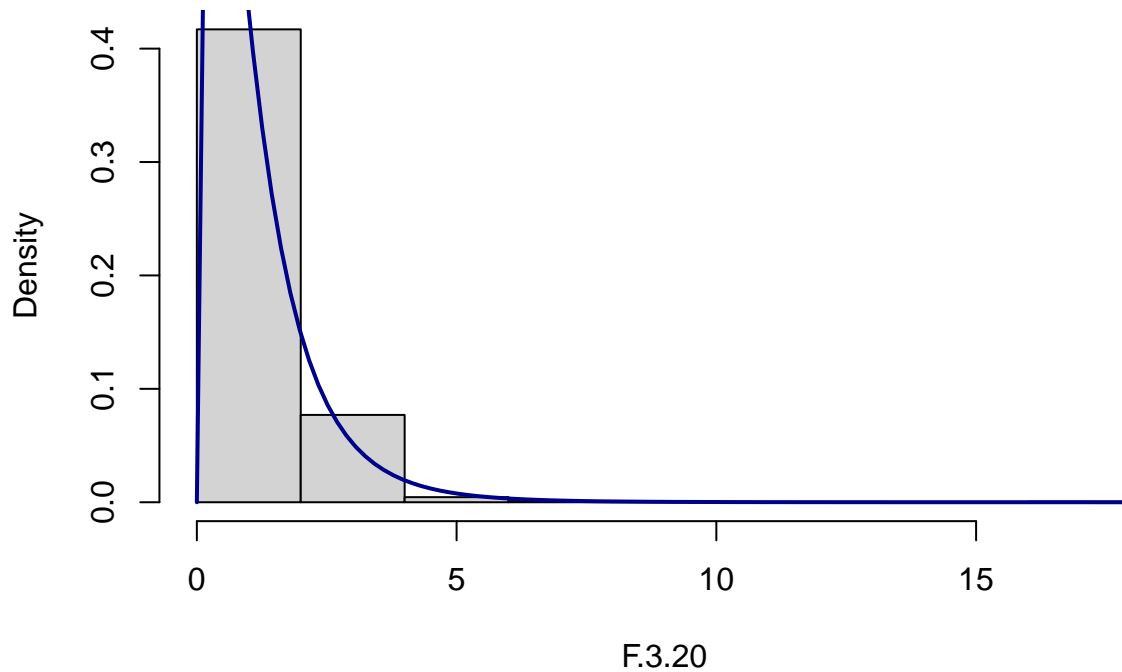
```
## Now how do we get to a t-distribution from Normal and chi-sq distributions?  
# First, take a scaled chi-sq by dividing it by the degrees of freedom  
scaled.chi.sq.3 <- chi.sq.3 / 3  
# Then take a normally distributed variable and divide them by the square root of the scaled chi-sq var  
  
normal.d <- rnorm( n=1000 ) # yet another #set of normally distributed data  
t.3 <- normal.d / sqrt( scaled.chi.sq.3 ) # divide by #square root of scaled chi-square to get t  
hist( t.3 )
```

Histogram of t.3



```
## To get to an F distribution, take the ratio between two scaled chi-sq distributions.  
# F distribution with 3 and 20 degrees of freedom:  
# first take two chi-sq variables, with 3 dof and 20 dof respectively, and take the ratio:  
  
chi.sq.20 <- rchisq( 1000, 20)           # generate chi square data with df = 20...  
scaled.chi.sq.20 <- chi.sq.20 / 20       # scale the chi square variable...  
F.3.20 <- scaled.chi.sq.3 / scaled.chi.sq.20 # take the ratio of the two chi squares...  
hist( F.3.20, freq = FALSE)              # ... and draw a picture  
curve(df(x, 3, 20),  
      col="darkblue", lwd=2, add=TRUE, yaxt="n")
```


Histogram of F.3.20



The curve above confirms this looks similar if you use the R built-in function `df` (just like `dnorm`, `pnorm`, `qnorm`, `pnorm`)

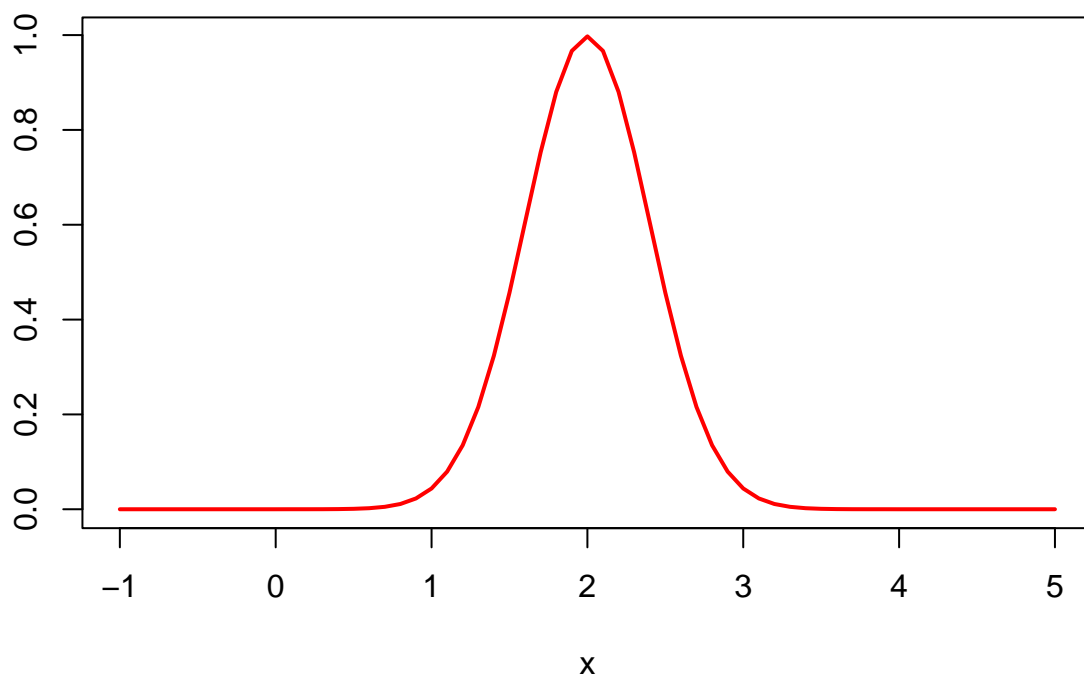
Lab 1 Generalization exercises

use the code from above to attempt to solve the extra things we ask you do for this assignment

Q1 Plot a normal distribution with mean = 2, s.d. = 0.4

```
x <- seq(-1, 5, 0.1)
```

```
plot(x, dnorm(x, mean = 2, sd = 0.4), type = "l",  
     ylab = "", lwd = 2, col = "red")
```



```
# Q2 Calculate the 85th %ile of the above distribution.  
q85 = qnorm(0.85,2,0.4)  
print(q85)
```

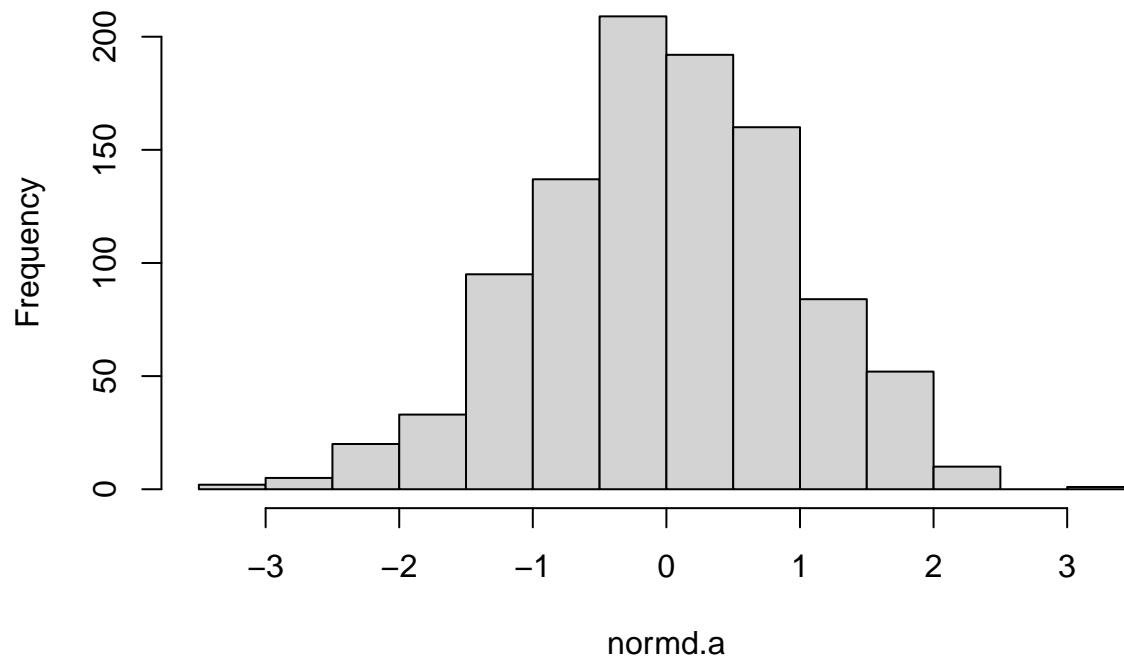
```
## [1] 2.414573
```

```
# Q3 Calculate the probability that a value lies in between 1 and 2 given the above distribution  
prob = pnorm(2,2,0.4) - pnorm(1,2,0.4)  
print(prob)
```

```
## [1] 0.4937903
```

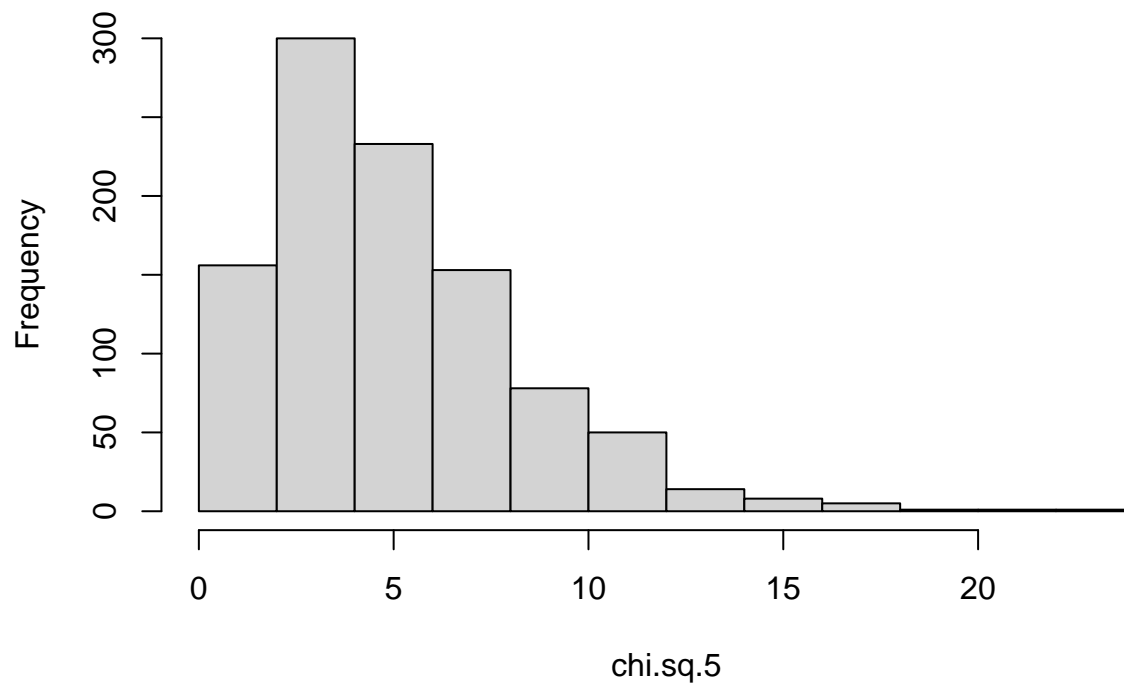
```
# Q4 Plot a simulated t-distribution with 5 degrees of freedom.  
normd.a = rnorm(1000)  
hist(normd.a)
```

Histogram of normd.a

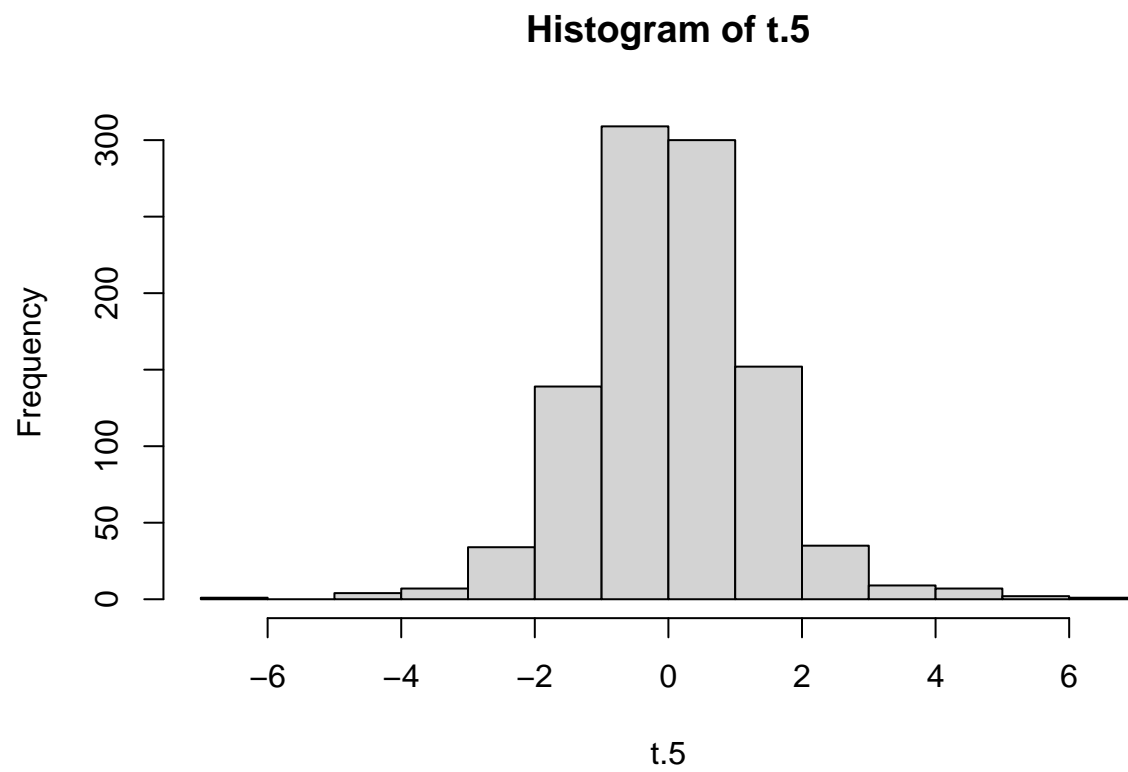


```
normd.b = rnorm(1000)
normd.c = rnorm(1000)
normd.d = rnorm(1000)
normd.e = rnorm(1000)
chi.sq.5 <- (normd.a)^2 + (normd.b)^2 + (normd.c)^2 + (normd.d)^2 + (normd.e)^2
hist(chi.sq.5)
```

Histogram of chi.sq.5



```
scaled.chi.sq.5 = chi.sq.5 / 5  
  
normd.f <- rnorm( n=1000 )           # yet another #set of normally distributed data  
t.5 <- normd.f / sqrt( scaled.chi.sq.5 ) # divide by #square root of scaled chi-square to get t  
hist(t.5)
```



Lab 1 Written answer question
Write your answer here.