# CLT with R

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### Central Limit Theorem

Recall:  $X_i \sim (\mu, \sigma^2), i = 1, 2, 3, \dots, n$  By CLT,

$$\frac{\bar{X}_n - \mu}{\sqrt{\frac{\sigma^2}{n}}} \to^d N(0, 1)$$

as  $n \to \infty$  where  $\bar{X_n} \sim (\mu, \frac{\sigma^2}{n})$ 

Let's do the simulation and see how  $\bar{X_n}$  behaves.

## Generating pseudo data from known distributions

#### Main character : Sample Mean X\_bar

Say  $X_i \sim Binomial(n = 5, p = 0.1), i = 1, 2, ..., 10000 E(X) = 5 \times 0.1 = 0.5, Var(X) = 5 \times 0.1 \times (1 - 0.1) = 0.45$ 

```
\#X \sim Bin(n = 5, p = 0.1)

x1 = rbinom(10000, size = 5, prob = 0.1)

mean(x1)
```

```
## [1] 0.5048
var(x1)
```

## [1] 0.4612231

But this is what only one  $\bar{X}$  behaves. If I want to see the how  $\bar{X}$  distributed, I need a bunch of realizations of  $\bar{X}$ .

#### Writing a function to generate many realizations

I can simply repeat the process above and get many X.

Define a function  $x_bar_bin(n)$  which helps me to get n realizations from Bin(5,0.1).

```
x_bar_bin = function(n){ #input: the number of realization of x_bar
x_bar = c()
for(i in 1:n){
    x1 = rbinom(n, size = 5, prob = 0.1)
    x_bar = rbind(x_bar, mean(x1))
}
return(x_bar)
}
```

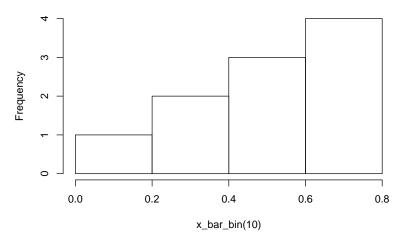
When input is 10, then it gives me 10 realizations.

```
t(x_bar_bin(10)) #10 realization
```

```
## [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
## [1,] 0.7 0.5 0.5 0.1 0.5 0.7 0.7 0.3 0.9 0.3
```

hist(x\_bar\_bin(10)) #see the distribution of 10 realizations

#### Histogram of x\_bar\_bin(10)

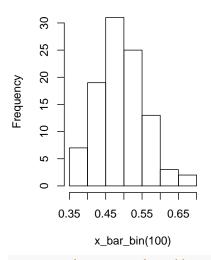


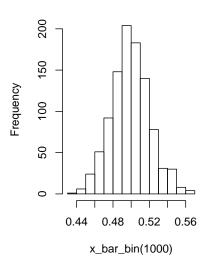
The followings are 100 realizations and 1000 realizations.

```
par(mfrow = c(1,2))
hist(x_bar_bin(100))
hist(x_bar_bin(1000))
```

#### Histogram of x\_bar\_bin(100)

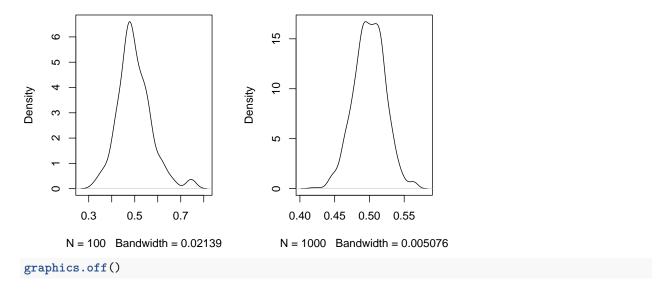
### Histogram of x\_bar\_bin(1000)





```
#density(x_bar_bin(1000)) #check`density()` if you're interested in it
plot(density(x_bar_bin(100)))
plot(density(x_bar_bin(1000)))
```

#### density.default( $x = x_bar_bin(100)$ density.default( $x = x_bar_bin(100)$



#### Adding Normal Distribution Curve

x\_bar\_bin(10)

We just define a function that helps us find  $\bar{X}_n$  under different n. Let's add the normal curve and see how  $\bar{X}_n$  fits normal distribution under different n.

```
set.seed(12345)
par(mfrow = c(1,3))
hist(x_bar_bin(10), freq = F)
curve(dnorm(x, mean = mean(x_bar_bin(10)), sd = sd(x_bar_bin(10))), add = T, col = 'blue')
hist(x_bar_bin(100), freq = F)
curve(dnorm(x, mean = mean(x_bar_bin(100)), sd = sd(x_bar_bin(100))), add = T, col = 'blue')
hist(x_bar_bin(1000), freq = F)
curve(dnorm(x, mean = mean(x_bar_bin(1000)), sd = sd(x_bar_bin(1000))), add = T, col = 'blue')
   Histogram of x_bar_bin(10)
                             Histogram of x_bar_bin(100)
                                                        Histogram of x_bar_bin(1000)
   2.5
   2.0
   5.
                                                     Density
                                                         10
   1.0
                              2
   0.5
                                                             0.45
     0.2 0.4 0.6 0.8 1.0 1.2
                                0.35
                                     0.45
                                          0.55
                                                                  0.50
                                                                       0.55
```

x\_bar\_bin(1000)

x\_bar\_bin(100)

## Normalization

We've seen how  $\bar{X_n}$  behaves. Now let's see how normalized z behaves. Let

$$z = \frac{\bar{X_n} - \mu}{\sqrt{\frac{\sigma^2}{n}}}$$

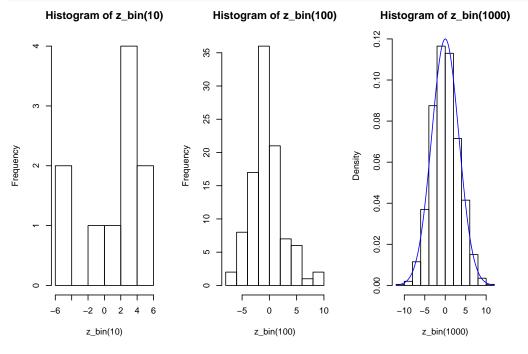
We also define a function that returns the normalized realization of the r.v.

```
#See what the normalized r.v. behaves
#Z = sqrt(n)*(X_bar - mu)/sigma^2
z_bin = function(n){
    z = c()
    for(i in 1:n){
        x1 = rbinom(n, size = 5, prob = 0.1) #Bin(5, 0.1)
        #sqrt(n)*(mean(x1) - 5*0.1)/(5*0.1*0.9)^2 #normalization
        z = rbind(z, sqrt(n)*(mean(x1) - 5*0.1)/(5*0.1*0.9)^2)
}
return(z)
}
```

Similarly,  $z_bin(n)$  gives us n realizations of the normalized r.v.

```
t(z_bin(10)) #10 normalized realization
```

```
##
              [,1]
                        [,2]
                                  [,3]
                                             [,4] [,5]
                                                            [,6]
                                                                     [,7]
## [1,] -6.246474 -1.561619 -1.561619 -3.123237
                                                     0 7.808093 4.684856
##
            [,8]
                      [,9]
                              [,10]
## [1,] 1.561619 4.684856 1.561619
par(mfrow = c(1,3))
hist(z bin(10))
hist(z_bin(100))
hist(z_bin(1000), freq = F)
curve(dnorm(x, mean=mean(z_bin(1000)),sd=sd(z_bin(1000))),add=T, col="blue")
```

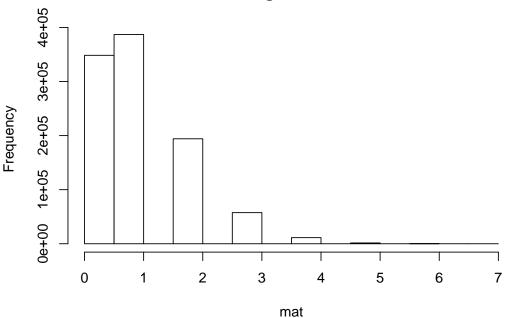


## Large Matrix Approach

a more intuitive approach to look at the random realizations

```
head(rbinom(1000, size = 10, p = 0.1)) #generate 1000 random numbers from Bin(10,0.1)
## [1] 1 1 0 0 0 3
mat = replicate(1000, rbinom(1000, size = 10, p = 0.1)) #generate 1000 sets of the above dim(mat)
## [1] 1000 1000
## [1] 1000 1000
If I just apply hist() on a matrix...
hist(mat) #overall dist. is Bin
```

## Histogram of mat



#this is element-wise operation

The column of the matrix is one of 1000 sets of 1000 realizations. We can get 1000 realizations of  $\bar{X}_n$  from 1000 columns.

```
#vertically sum up and divide by 1000 => get 1000 x_bars!
mean(mat[,1])

## [1] 1.052

mean(mat[,2])# on and on and on...

## [1] 0.98

With the for loop.

x_bar = c()
for(i in 1:1000){
 x_bar = rbind(x_bar, mean(mat[,i]))
}
t(head(x_bar))
```

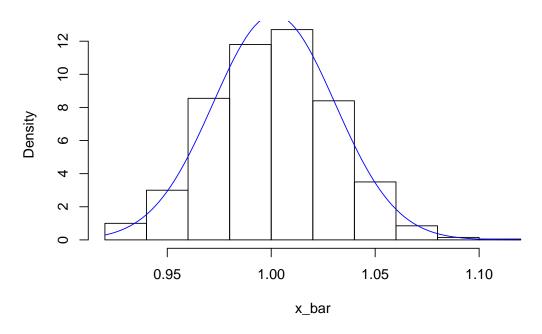
```
## [,1] [,2] [,3] [,4] [,5] [,6]
## [1,] 1.052 0.98 0.991 1.006 1.011 1.035
```

#### **Density Plot**

If we look at the "density" and compare it to normal density.

```
hist(x_bar, freq = F)
curve(expr = dnorm(x, mean = mean(x_bar), sd = sd(x_bar)), add = T, col = 'blue')
```

### Histogram of x\_bar



#### Alternative Expression (Optional)

x\_bar.hist = hist(x\_bar)

We can also draw the "Frequency Plot" instead of density plot. But we have to adjust the "height" of normal curve.

```
multiplier = x_bar.hist$counts/x_bar.hist$density
multiplier[1]

## [1] 20

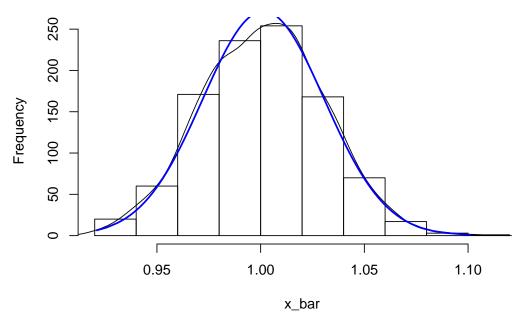
x_bar.den = density(x_bar)

x_bar.den$y = x_bar.den$y*multiplier[1]

#If we look ar the "frequency"...compare to normal
hist(x_bar)
lines(x_bar.den)
```

lines(sort(x\_bar), dnorm(x = sort(x\_bar), mean = mean(x\_bar), sd = sqrt(var(x\_bar))) \* multiplier[1], c





Also, for the normalized r.v., it should fits normal curve.

```
#normalization
z = (x_bar - 10*0.1)/(sqrt(10*0.1*0.9)/sqrt(1000))
h = hist(z, freq = F)
curve(expr = dnorm(x, mean = mean(z), sd = sd(z)), add = T, col = 'red')
```

# Histogram of z

