

week3script

Summer Heschong

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```
#load packages
```

```
library(here)
```

```
## here() starts at /Users/summerheschong/Documents/GitHub/stats_spring25
```

```
library(tidyverse)
```

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
```

```
## v dplyr      1.1.4      v readr      2.1.5
```

```
## v forcats    1.0.0      v stringr    1.5.1
```

```
## v ggplot2    3.5.1      v tibble     3.2.1
```

```
## v lubridate  1.9.4      v tidyr      1.3.1
```

```
## v purrr      1.0.2
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
```

```
## x dplyr::lag()     masks stats::lag()
```

```
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
```

```
library(moments)
```

```
#Discrete Distributions (Bernoulli Trial and Poisson Distribution) ###Bernoulli Trial
```

```
#probability of three successes
```

```
dbinom(x = 3, size = 16, prob = 0.48)
```

```
## [1] 0.01258796
```

```
#probability of 0-16 successes
```

```
prob <- dbinom(x = 0:16, size = 16, prob = 0.48)
```

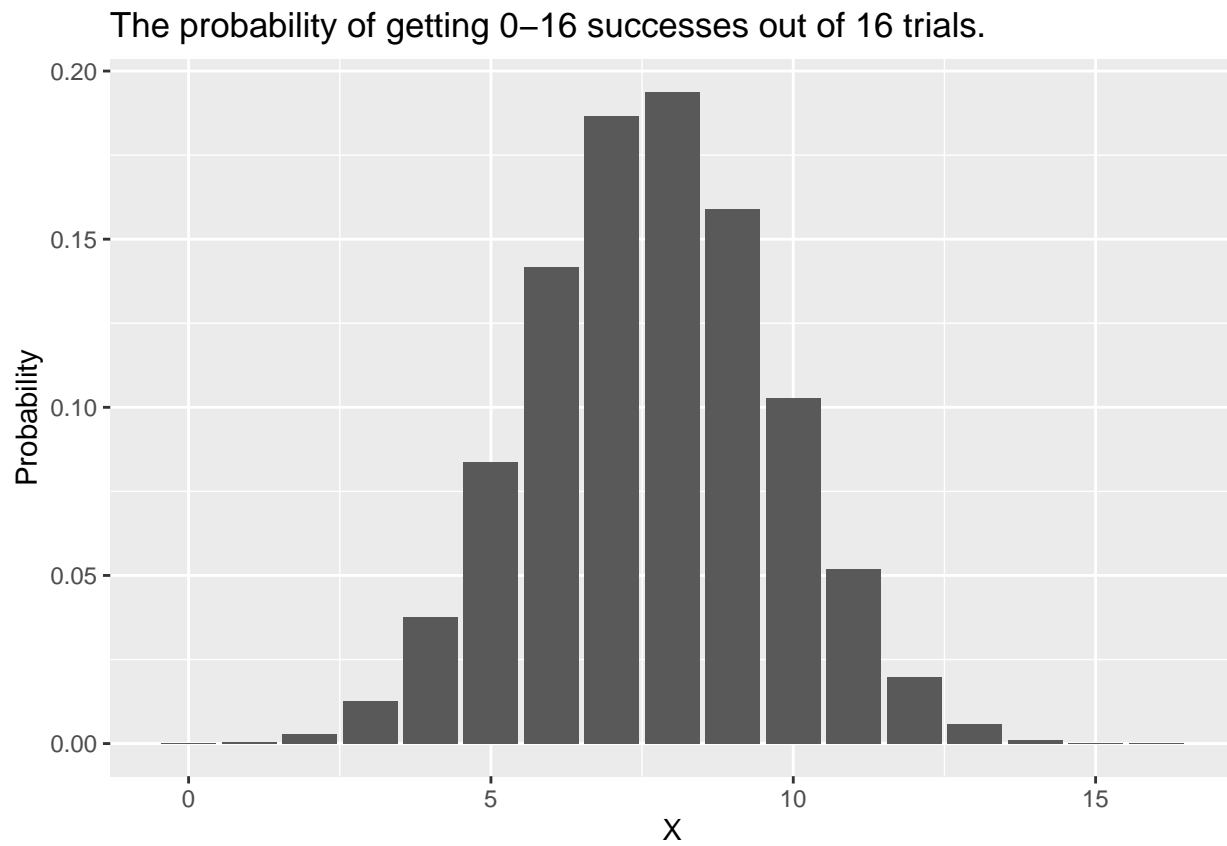
```
#create a list of numbers of successes
```

```
success <- 0:16
```

```
#bind two columns together to create a dataframe
```

```
binom_df <- cbind(prob, success)
```

```
#create a figure
ggplot(binom_df, aes(x = success,
                     y = prob)) +
  geom_bar(stat = "identity") +
  labs(x = 'X',
       y = 'Probability',
       title = 'The probability of getting 0-16 successes out of 16 trials.')
```



```
#Cumulative probability using dbinom
sum(dbinom(x = 0:3, size = 16, prob = 0.48))
```

```
## [1] 0.01596084
```

```
#cumulative probability using generated data
#adding together rows 1 through 4
sum(prob[1:4])
```

```
## [1] 0.01596084
```

```
#Probability of three or fewer successes
pbinom(q = 3, size = 16, prob = 0.48)
```

```
## [1] 0.01596084
```

####Poisson distribution models counts of outcomes rather than modeling successes and failures

```
#probability of 10 occurrences  
dpois(x = 10, lambda = 3)
```

```
## [1] 0.0008101512
```

```
#Probability of 3 or fewer occurrences  
ppois(q = 3, lambda = 3)
```

```
## [1] 0.6472319
```

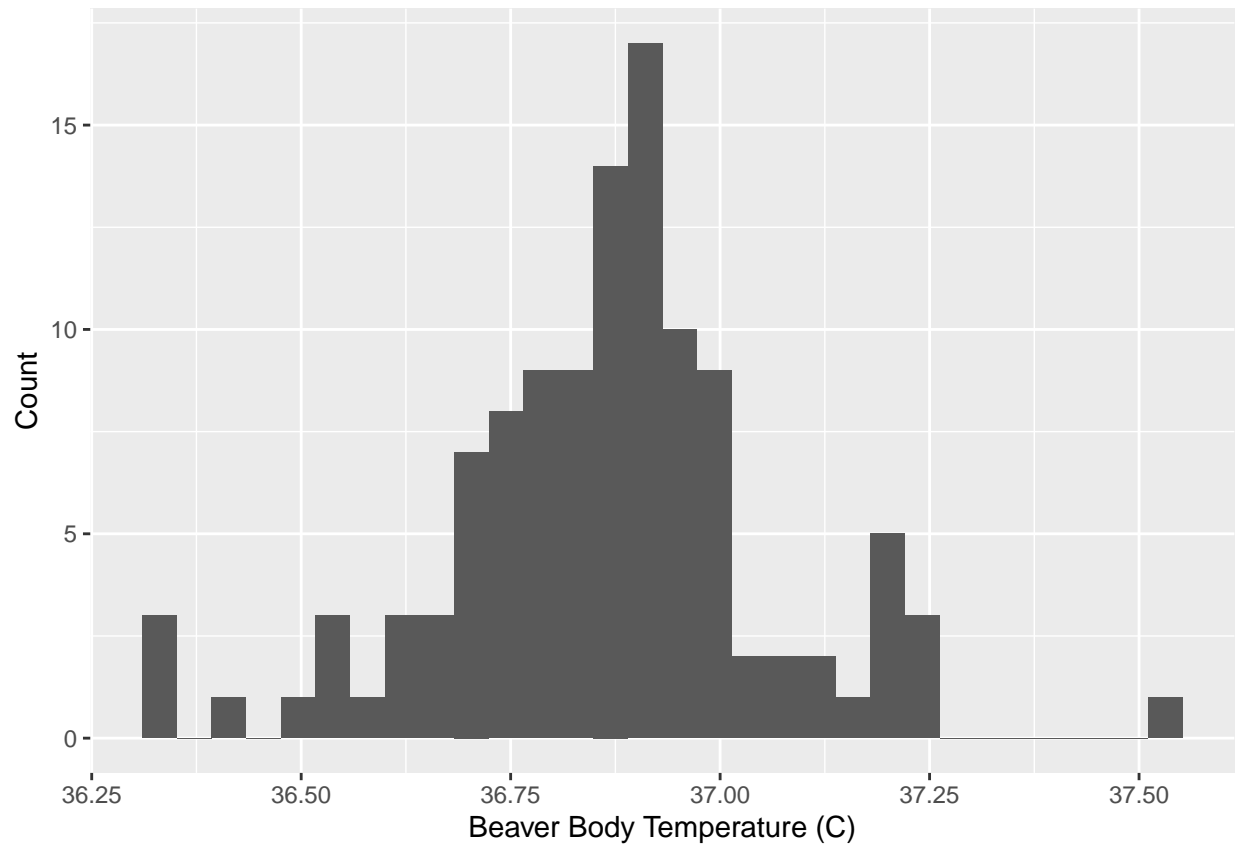
#Assessing Normality

```
#consider information on how this data was collected  
#?beaver1()
```

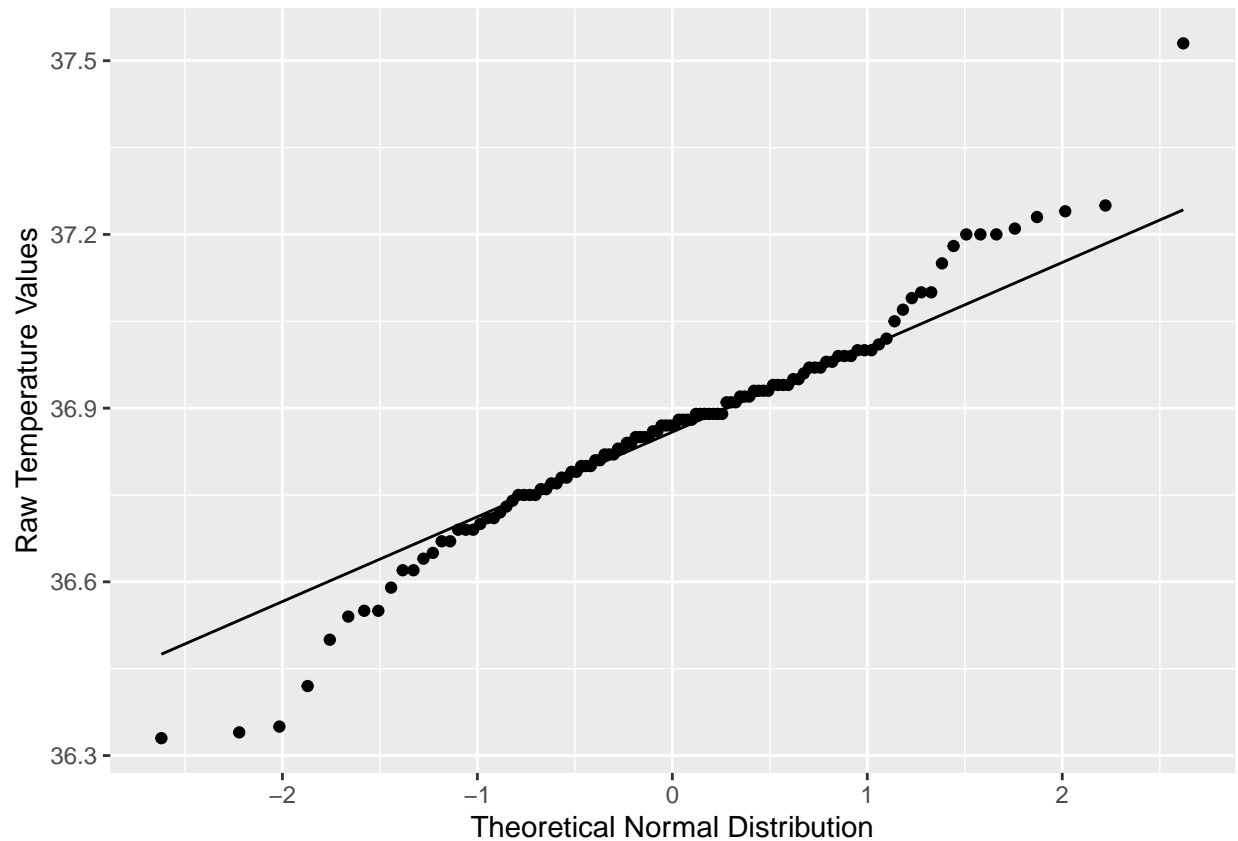
```
#visually inspect data  
#View(beaver1)
```

```
#create a histogram of beaver body temperature  
ggplot(beaver1, aes(x = temp)) +  
  geom_histogram() +  
  labs(x = 'Beaver Body Temperature (C)',  
       y = 'Count')
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```



```
#create a Q-Q plot of beaver body temperature  
ggplot(beaver1, aes(sample = temp)) +  
  geom_qq() +  
  geom_qq_line() +  
  labs(x = 'Theoretical Normal Distribution',  
       y = 'Raw Temperature Values')
```



```
#calculate skewness of beaver body temperature
skewness(beaver1$temp)
```

```
## [1] -0.02782567
```

```
#calculate kurtosis of beaver body temperature
kurtosis(beaver1$temp)
```

```
## [1] 4.351118
```

the absolute value of skewness for beaver body temperature is > 0 and < 0.5 therefore it is approximately symmetrical

kurtosis for beaver body temperature is > 3 therefore it is leptokurtic

#Continuous Distributions pnorm() calculates area under the probability density function of a normal distribution to the left of a specified value of x

```
#probability of value < 1
pnorm(q = 1, mean = 2, sd = 3)
```

```
## [1] 0.3694413
```

```
#prob of value < 1  
pnorm(q = 1, mean = 2, sd = 3)
```

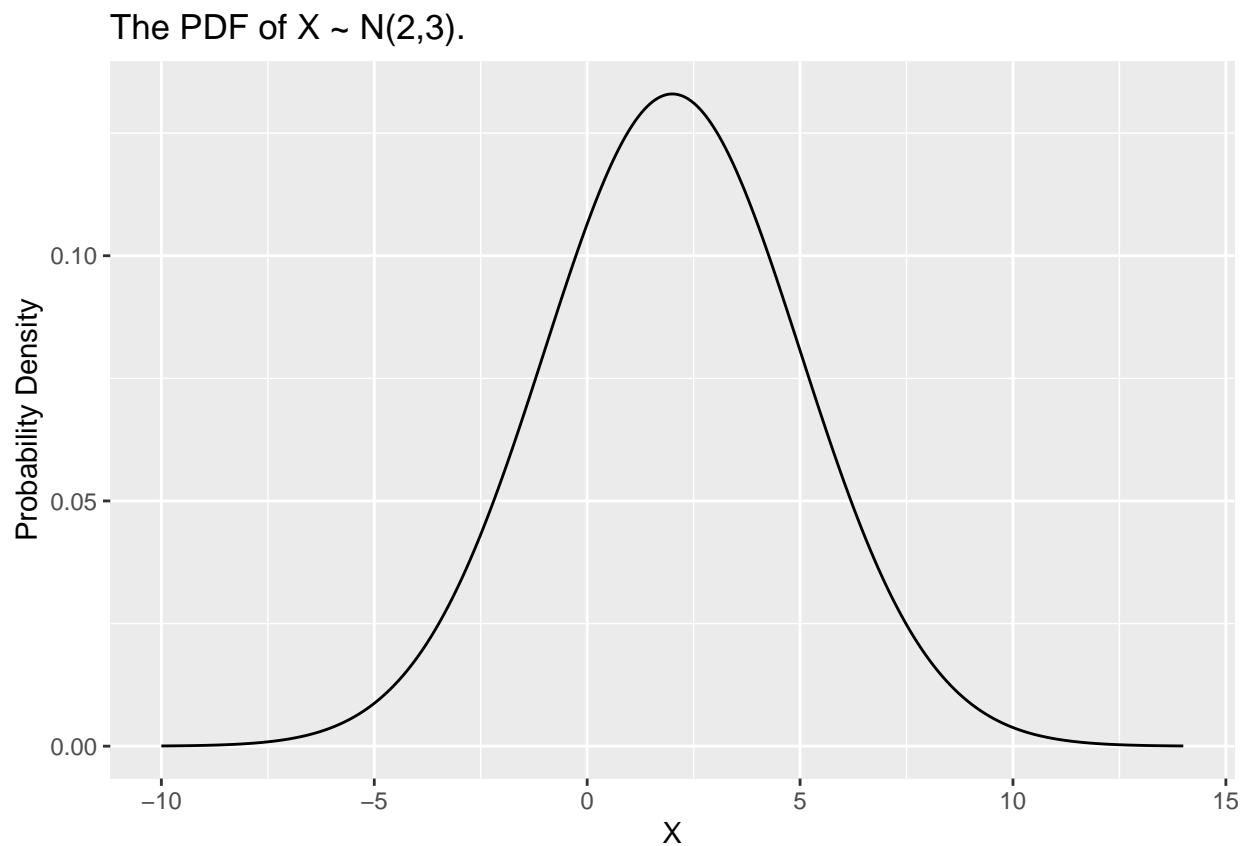
```
## [1] 0.3694413
```

```
#create a list of values with lots of values  
#in between whole numbers (so our curve looks nice and smooth)  
values <- seq(-10, 14, length = 1000)
```

```
#probability densities for values -10 to 14  
prob_norm <- dnorm(x = values, mean = 2, sd = 3)
```

```
#bind two columns together to create a dataframe  
norm_df <- cbind(prob_norm, values)
```

```
#create a figure  
ggplot(norm_df, aes(x = values,  
                    y = prob_norm)) +  
  geom_line() +  
  labs (x= "X",  
        y = 'Probability Density',  
        title = 'The PDF of X ~ N(2,3).')
```



qnorm() returns the value of x that is the upper bound of a particular probability

```
#value of 'x' for a given probability  
qnorm(p = 0.80, mean = 2, sd = 3)
```

```
## [1] 4.524864
```