

Stat123_MT#1

#VNUM: V01003221

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```
#a)
hours <- c(25,35,10,20,40,50)
sum_hours <- sum(hours)
avg_hours <- mean(hours)
len_hours <- length(hours)
len_hours

[1] 6

sum_hours

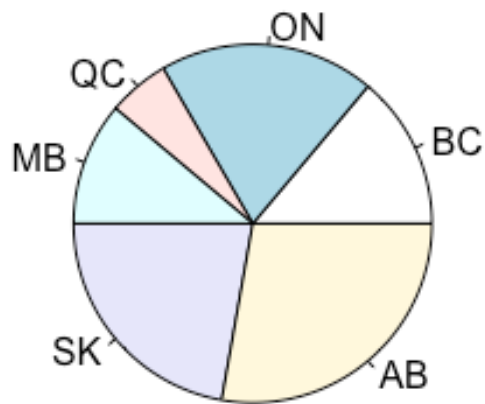
[1] 180

avg_hours

[1] 30

#b)
df <- data.frame(hours, c("BC", "ON", "QC", "MB", "SK", "AB"))
names(df[,1]) <- df[,2]

#c)
hours <- df[,1]
names(hours) <- df[,2]
pie(hours)
```



The length of this sample is 6

The total in of the hours is 180

The average hours from these numbers is 30

```
#2
#a)
Protein <- read.csv("protein.csv")
#b)
colnames(Protein)

[1] "Country" "RedMeat" "WhiteMeat" "Eggs" "Milk" "Fish"
[7] "Cereals" "Starch" "Nuts" "Fr.Veg"

#c)
class(Protein)

[1] "data.frame"

#d)
ProteinMatrix <- as.matrix(Protein[,c(2,3)])
ProteinMatrix
```

	RedMeat	WhiteMeat
[1,]	10.1	1.4
[2,]	8.9	14.0
[3,]	13.5	9.3
[4,]	7.8	6.0
[5,]	9.7	11.4
[6,]	10.6	10.8
[7,]	8.4	11.6
[8,]	9.5	4.9
[9,]	18.0	9.9
[10,]	10.2	3.0
[11,]	5.3	12.4
[12,]	13.9	10.0
[13,]	9.0	5.1
[14,]	9.5	13.6
[15,]	9.4	4.7
[16,]	6.9	10.2
[17,]	6.2	3.7
[18,]	6.2	6.3
[19,]	7.1	3.4
[20,]	9.9	7.8
[21,]	13.1	10.1
[22,]	17.4	5.7
[23,]	9.3	4.6
[24,]	11.4	12.5
[25,]	4.4	5.0

```
#e)
names(ProteinMatrix) <- c("Beef", "Chicken")
ProteinMatrix
```

	RedMeat	WhiteMeat
[1,]	10.1	1.4
[2,]	8.9	14.0
[3,]	13.5	9.3
[4,]	7.8	6.0
[5,]	9.7	11.4
[6,]	10.6	10.8
[7,]	8.4	11.6
[8,]	9.5	4.9
[9,]	18.0	9.9
[10,]	10.2	3.0
[11,]	5.3	12.4
[12,]	13.9	10.0
[13,]	9.0	5.1
[14,]	9.5	13.6
[15,]	9.4	4.7
[16,]	6.9	10.2
[17,]	6.2	3.7
[18,]	6.2	6.3

```

[19,]      7.1      3.4
[20,]      9.9      7.8
[21,]     13.1     10.1
[22,]     17.4      5.7
[23,]      9.3      4.6
[24,]     11.4     12.5
[25,]      4.4      5.0
attr(,"names")
 [1] "Beef"      "Chicken" NA      NA      NA      NA      NA
 [8] NA          NA      NA      NA      NA      NA      NA
[15] NA          NA      NA      NA      NA      NA      NA
[22] NA          NA      NA      NA      NA      NA      NA
[29] NA          NA      NA      NA      NA      NA      NA
[36] NA          NA      NA      NA      NA      NA      NA
[43] NA          NA      NA      NA      NA      NA      NA
[50] NA

```

The names are: Country, RedMeat, WhiteMeat, Eggs, Milk, Fish, Cereals, Starch, Nutes, Fr.Veg

Protein is a data frame

```

#3a)
head(airquality)

  Ozone Solar.R Wind Temp Month Day
1    41    190  7.4   67     5   1
2    36    118  8.0   72     5   2
3    12    149 12.6   74     5   3
4    18    313 11.5   62     5   4
5    NA     NA 14.3   56     5   5
6    28     NA 14.9   66     5   6

hist(airquality$Temp, main = "Maximum daily temperature at La Guardia
Airport", xlab = "Temperature in degrees Fahrenheit", breaks = seq(50,100),
col = "darkmagenta", fill = "darkmagenta")

Warning in plot.window(xlim, ylim, "", ...): "fill" is not a graphical
parameter

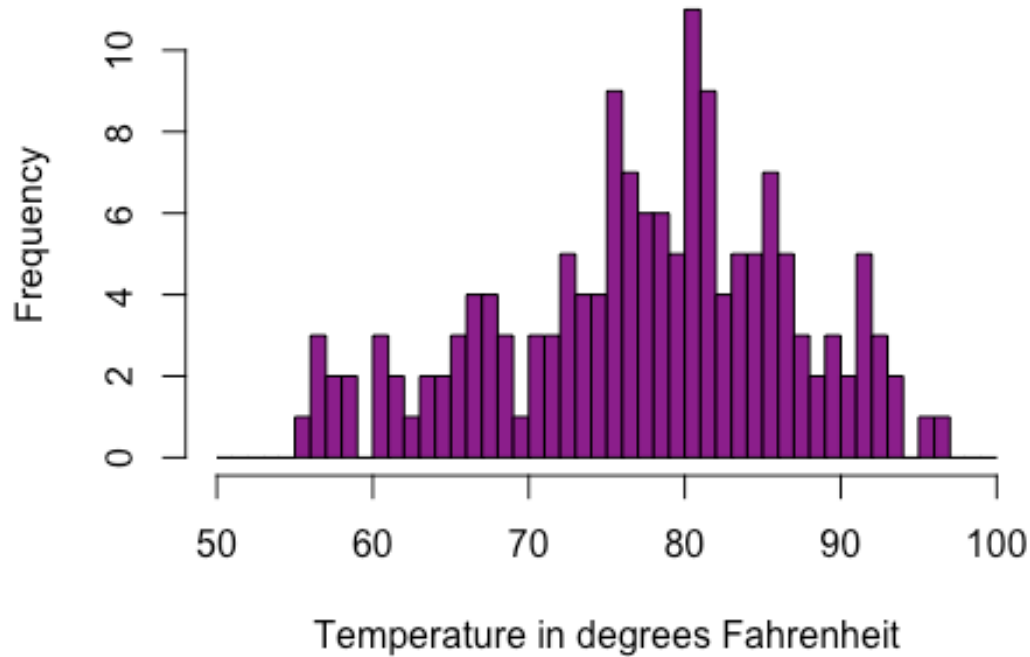
Warning in title(main = main, sub = sub, xlab = xlab, ylab = ylab, ...):
"fill"
is not a graphical parameter

Warning in axis(1, ...): "fill" is not a graphical parameter

Warning in axis(2, at = yt, ...): "fill" is not a graphical parameter

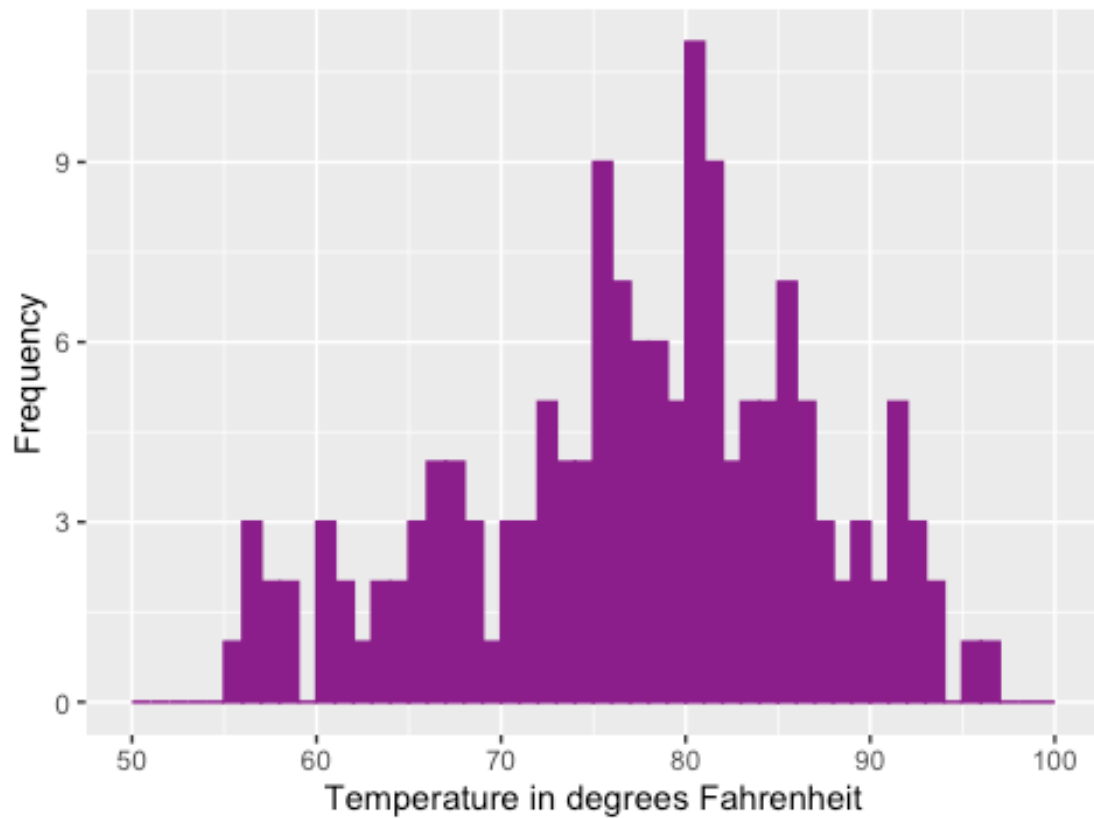
```

Maximum daily temperature at La Guardia Airport



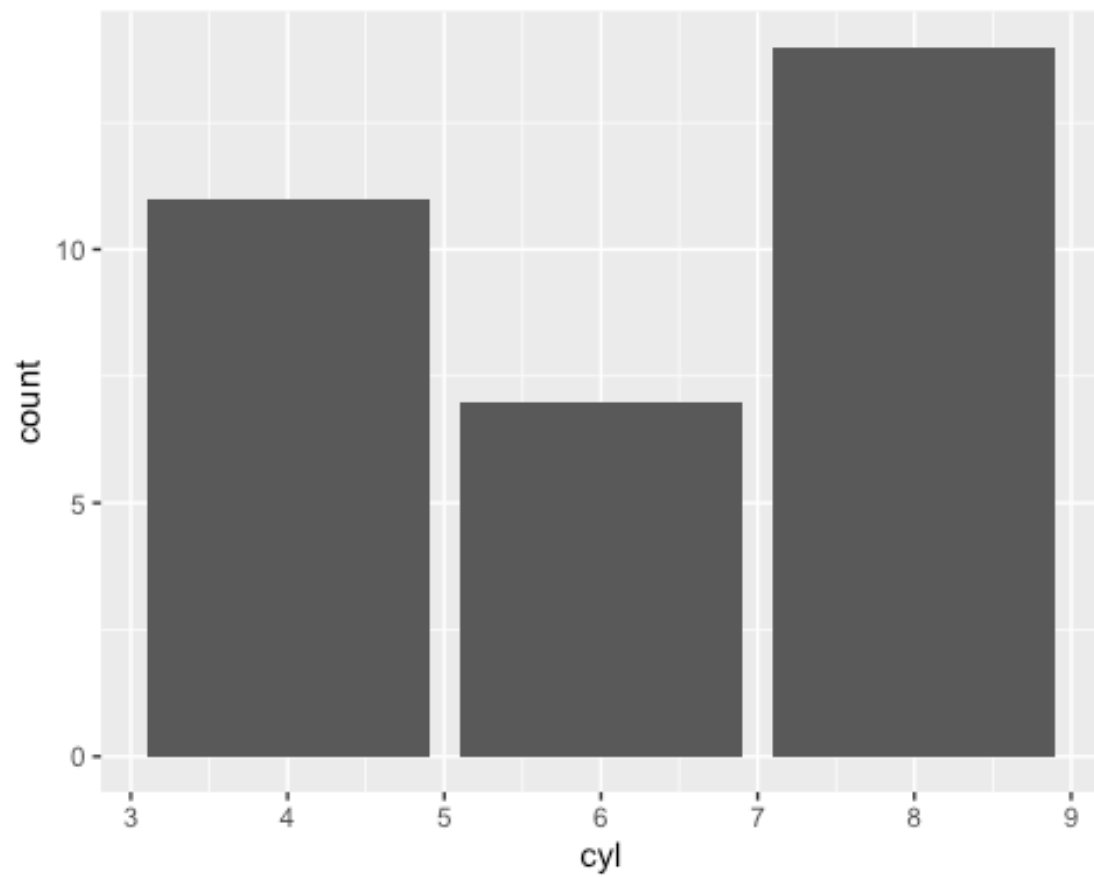
```
#b)
library(ggplot2)
ggplot(airquality, aes(x = Temp)) + geom_histogram(breaks = seq(50,100), col = "darkmagenta", fill = "darkmagenta") + labs(x = "Temperature in degrees Fahrenheit", title = "Maximum daily temperature at La Guardia Airport", y = "Frequency")
```

Maximum daily temperature at La Guardia Airport

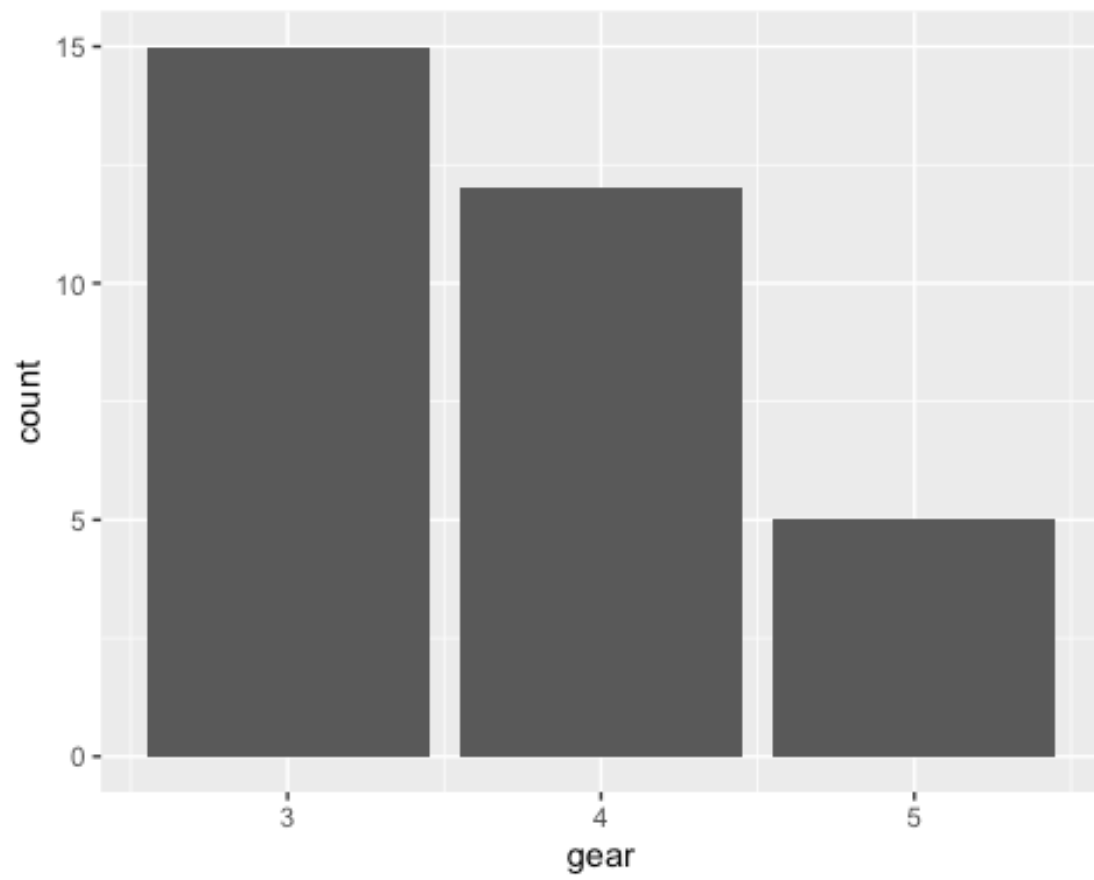


```
?par
gear=as.factor(mtcars$gear)
disp=as.factor(mtcars$disp)

library(ggplot2)
ggplot(mtcars, aes(x = cyl)) + geom_bar()
```

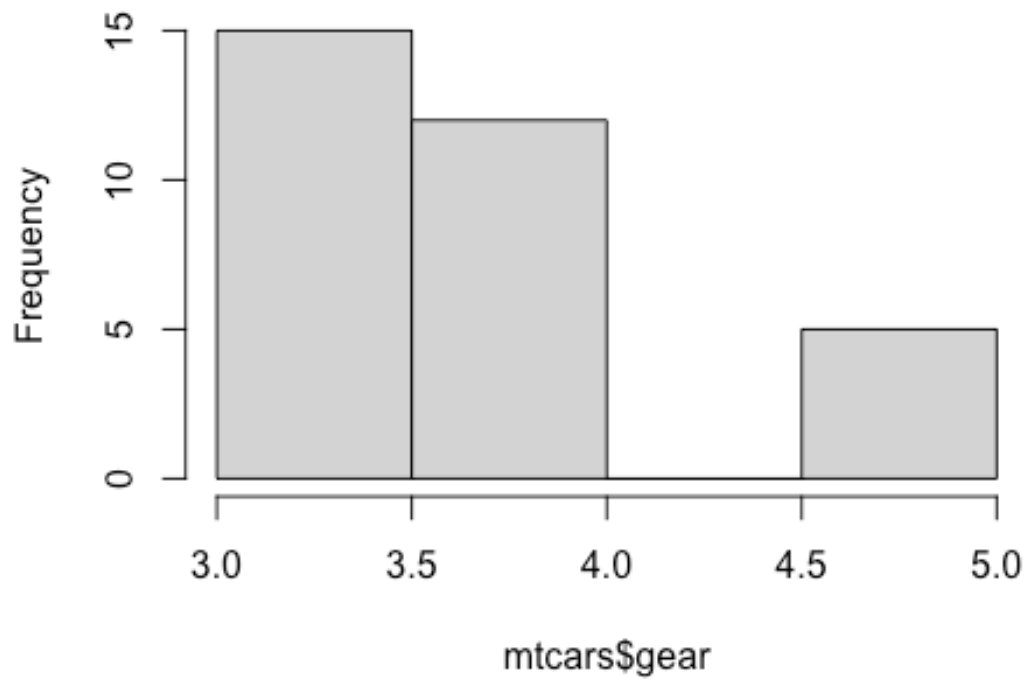


```
ggplot(mtcars, aes(x = gear)) + geom_bar()
```



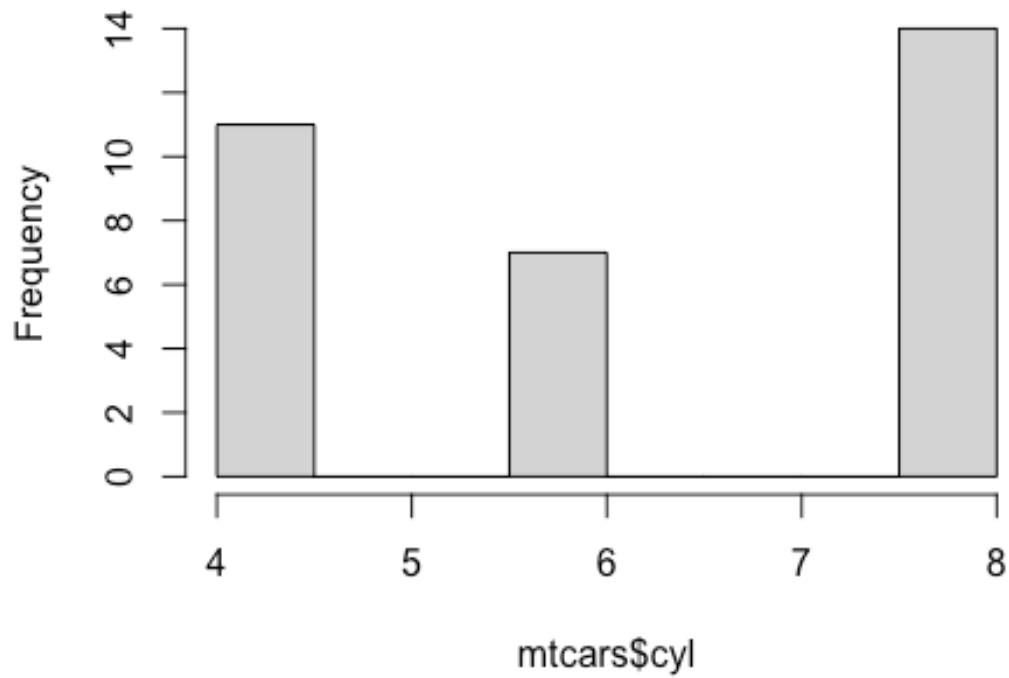
```
hist(mtcars$gear)
```


Histogram of mtcars\$gear



```
hist(mtcars$cyl)
```

Histogram of mtcars\$cyl



?hist