

## Activity 6 - R Programming - If condition and Nested loop

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1. Let's write if and else statement and nested for loop.
  - a. We have a vector of area codes and we want to recode them as either in New York or outside New York. Show a frequency table displaying how many are in NY and outside NY.

```
area.codes <-  
c(865,865,423,615,615,206,626,514,308,308,514,931,425,212,917,585,607,718,347,  
929) #data  
ny.codes <-  
c(212,332,646,917,315,516,518,585,607,631,716,718,845,914,680,838,934,347,929  
) #area codes in NY  
  
in.newyork <- c()  
for (i in 1:length(area.codes)) {  
  if( area.codes[i] %in% ny.codes ) { in.newyork[i] <- "yes" }  
  else { in.newyork[i] <- "no" }  
}  
in.newyork <- factor(in.newyork)  
table(in.newyork)  
## in.newyork  
## no yes  
## 13 7
```

- b. In the CUSTLOYALTY data in regclass, the lifetime values and total transactions of customers are recorded along with some of their characteristics. Let's find the average total transactions for each of the 6 income groups for married, then do the same again for single. Show a matrix that reports the average total transactions for each income group and marital status.

```
library(regclass)  
## Loading required package: bestglm  
## Loading required package: leaps  
## Loading required package: VGAM  
## Loading required package: stats4  
## Loading required package: splines  
## Loading required package: rpart  
## Loading required package: randomForest  
## randomForest 4.7-1.1  
## Type rfNews() to see new features/changes/bug fixes.
```

```

## Important regclass change from 1.3:
## ALL functions that had a . in the name now have an _
## all.correlations -> all_correlations, cor.demo -> cor_demo, etc.
data(CUSTLOYALTY)
levels(CUSTLOYALTY$Married)
## [1] "Married" "Single"
levels(CUSTLOYALTY$Income)
## [1] "f0t30" "f30t45" "f45t60" "f60t75" "f75t90" "f90toINF"

VALUE <- matrix(0, nrow = nlevels(CUSTLOYALTY$Married), ncol =
nlevels(CUSTLOYALTY$Income))
rownames(VALUE) <- levels(CUSTLOYALTY$Married)
colnames(VALUE) <- levels(CUSTLOYALTY$Income)

for (marital_status in 1:length(levels(CUSTLOYALTY$Married))) {
  for (incomegroup in 1:length(levels(CUSTLOYALTY$Income))) {
    SUB <- subset(CUSTLOYALTY, Married ==
levels(CUSTLOYALTY$Married)[marital_status] & Income ==
levels(CUSTLOYALTY$Income)[incomegroup])
    VALUE[marital_status, incomegroup] <- mean(SUB$TotTransactions)
  }
}
VALUE
##           f0t30  f30t45  f45t60  f60t75  f75t90 f90toINF
## Married 2.909091 5.241379 5.600000 8.52381 7.125000 9.250000
## Single  3.611111 5.044118 6.313433 6.80000 7.402778 7.955556

```

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## Activity 7

1. We shall generate several figures by using generic plot function.
  - a. Let's load women data which is saved in dataset R package. And check what variables are included in this dataset by using names Weightfunction.

```

data(women, package="datasets")
names(women)
## [1] "height" "weight"

```

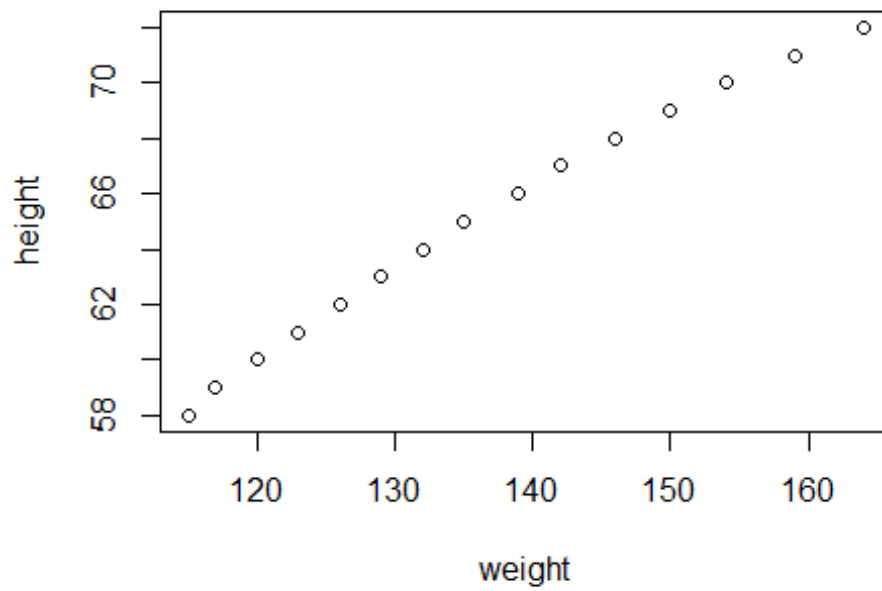
- b. Let's generate a basic plot by displaying x=height, y=weight.

```

plot(height~weight, data=women)

```

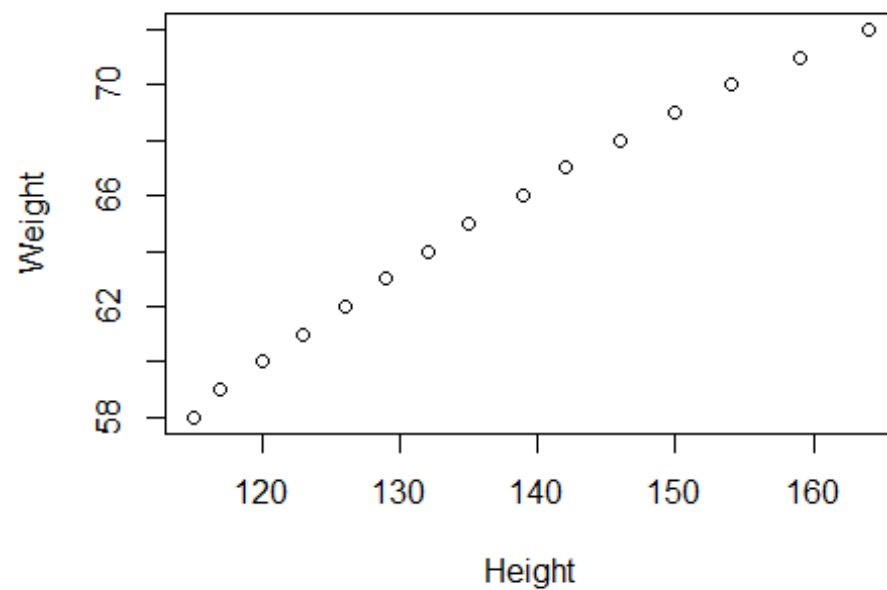
---



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c. Let's add labels for x and y, e.g., x-label = Height, y-label = Weight.

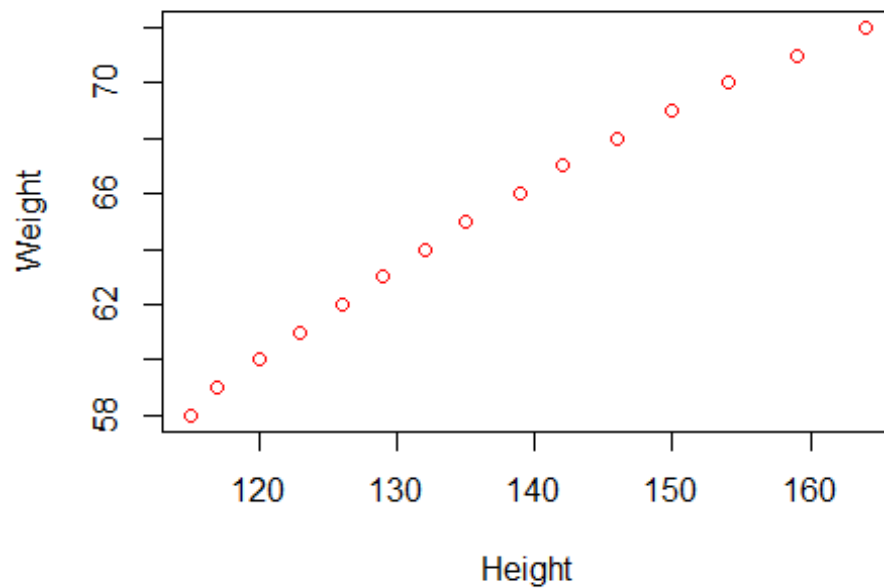
```
plot(height~weight, data=women, xlab="Height", ylab="Weight")
```



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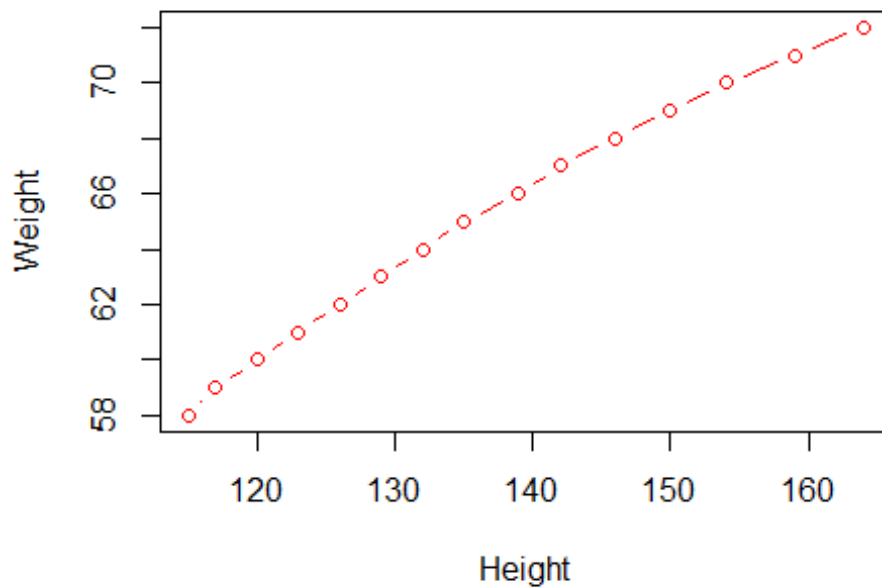
d. Let's change the color of the points to be red.

```
plot(height~weight, data=women, xlab="Height", ylab="Weight", col="red")
```



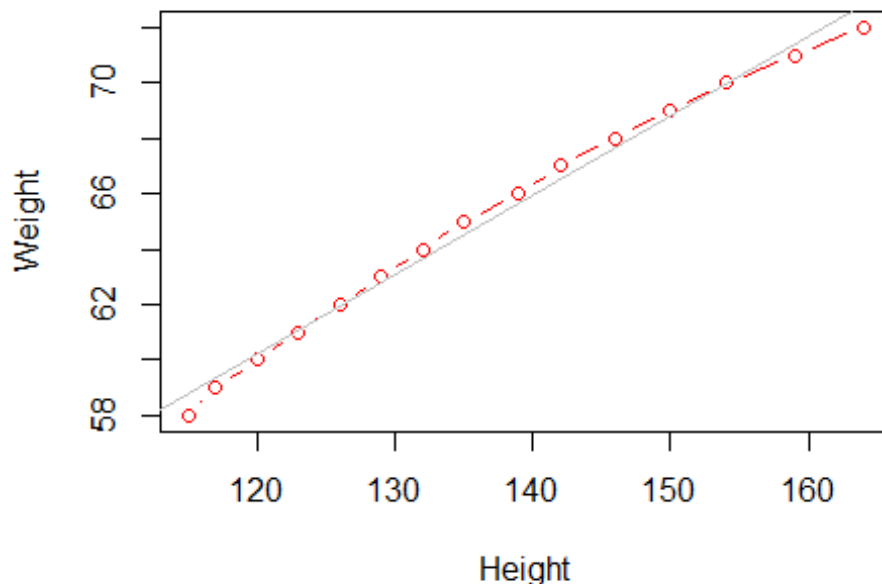
- 
- e. Let's generate a different type of a plot which displays both the points and lines.  
Hint: turn on type argument in a generic `plot()` function.

```
plot(height~weight, data=women, xlab="Height", ylab="Weight", col="red",  
type='b')
```



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- f. Let's add a regression line to the previous plot and let this regression line has gray color.

```
plot(height~weight, data=women, xlab="Height", ylab="Weight", col="red",  
type='b')  
abline(lm(height~weight, data=women), col="gray")
```

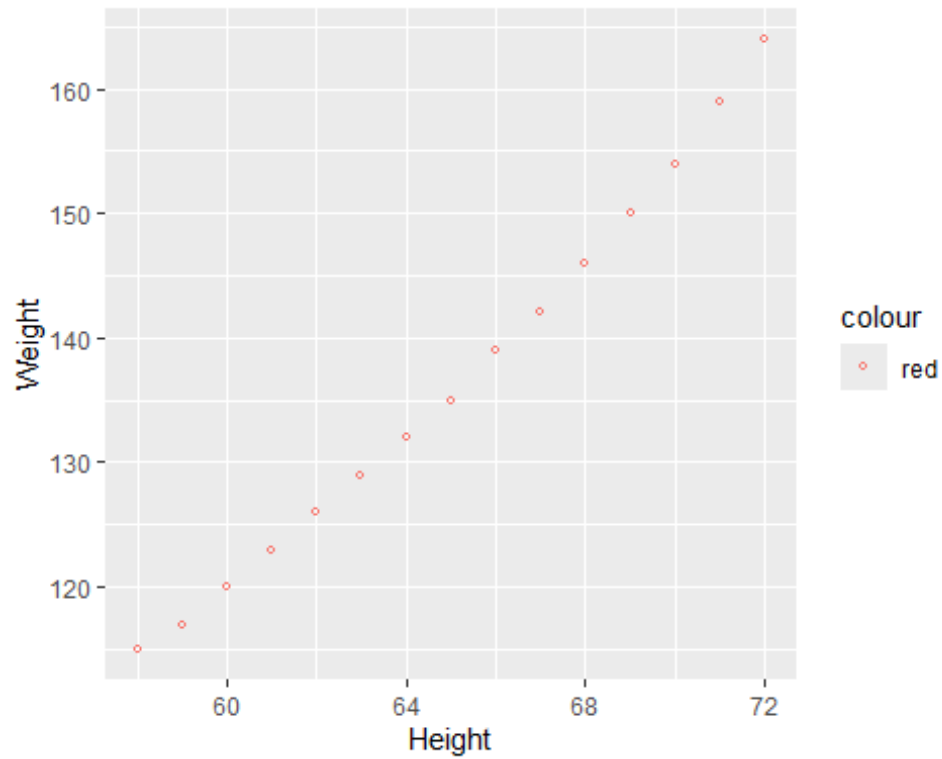


2. We will use ggplot2 R package to generate similar plots. Load ggplot2 R package.

```
#install.packages("ggplot2")
library(ggplot2)
## Warning: package 'ggplot2' was built under R version 4.3.3
##
## Attaching package: 'ggplot2'
## The following object is masked from 'package:randomForest':
##
##     margin
```

- a. Use ggplot() function to generate a (point) plot with x=Height, y=Weight. Add labels for x and y like the graph in Question 1 (c). And let the points be red. Also, let pch=1 and size=1 for the points. Hint: Use ggplot(), geom\_point(), ylab, and xlab (See Visualization slides)

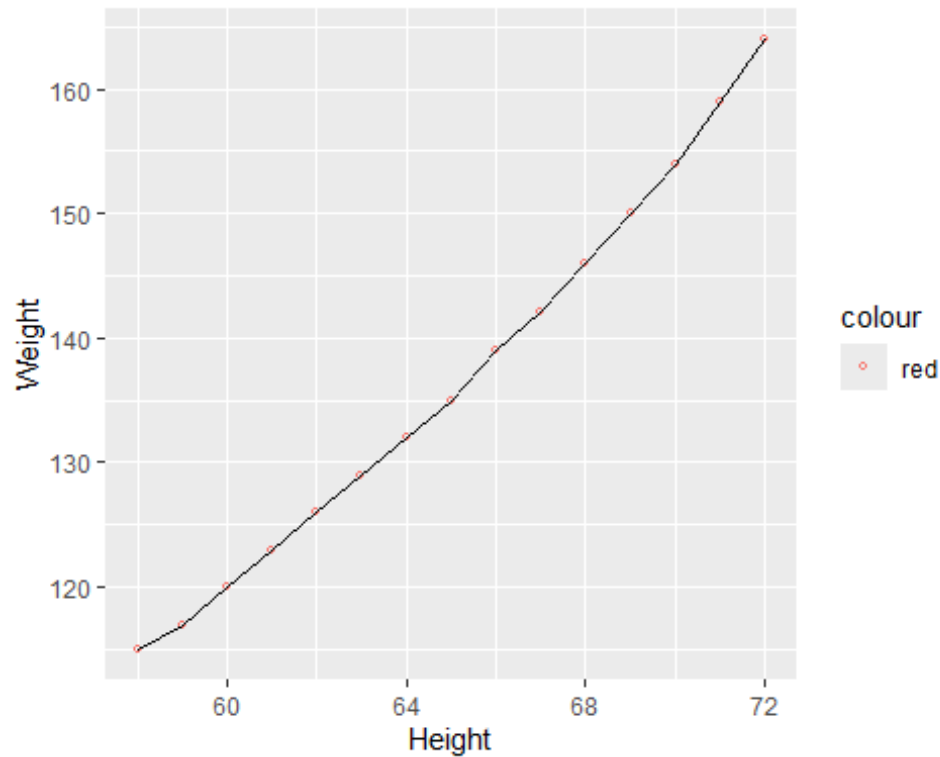
```
ggplot(women, aes(x = height, y = weight, color='red'))+
geom_point(size = 1, pch = 1)+
labs(y="Weight", x="Height")
```



- b. Let's add lines to the previous graph and connect the points. Let the lines be black color. Hint: Use `geom_line()` (See Visualization slides)

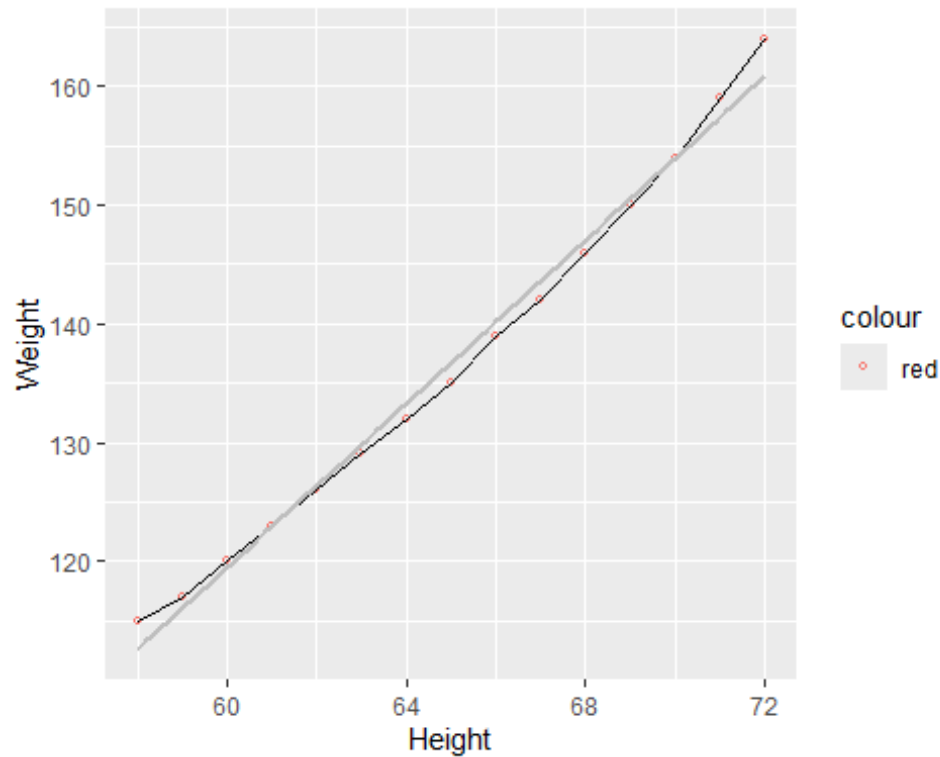
```
ggplot(women, aes(x = height, y = weight, color='red'))+  
geom_point(size = 1, pch = 1)+  
labs(y="Weight", x="Height")+  
geom_line(color="black")
```





- c. Let's add a regression line (without the confidence intervals) to the previous graph and let this regression line has gray color. Hint: Use `geom_smooth()` (See Visualization slides)

```
ggplot(women, aes(x = height, y = weight, color='red'))+  
geom_point(size = 1, pch = 1)+  
labs(y="Weight", x="Height")+  
geom_line(color="black")+  
geom_smooth(method="lm", color="gray", se=FALSE)  
## `geom_smooth()` using formula = 'y ~ x'
```



- d. Let's add a smooth line using loess (without the confidence intervals) to the previous graph and let this regression line has blue color. Hint: Use `geom_smooth()` (See Visualization slides)

```
ggplot(women, aes(x = height, y = weight, color='red'))+  
geom_point(size = 1, pch = 1)+  
labs(y="Weight", x="Height")+  
geom_line(color="black")+  
geom_smooth(method="lm", color="gray", se=FALSE)+  
geom_smooth(method="loess", color="blue", se=FALSE)  
## `geom_smooth()` using formula = 'y ~ x'  
## `geom_smooth()` using formula = 'y ~ x'
```

