

R Basics

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May 12, 2016

1 R Basics

R is a *programming language*.

Let's review some of its basic characteristics

- Basic operations
- Common functions
- Data types

1.1 Basic operations: Assignment

One of the main differences is the operator "<-"

```
#Let's create variables of different types
#numeric
my_num1<- 15
print (my_num1)
```

```
>>> [1] 15
```

```
#character
my_string<- "Hello world"
print (my_string)
```

```
>>> [1] "Hello world"
```

```
#logical
my_bool<- TRUE
print (my_bool)
```

```
>>> [1] TRUE
```

1.2 Basic operations: Aritmetics

```
# We can do operations with the variables
my_num1<-15
my_num2<-5
#subtraction
my_num1-my_num2
```

```
>>> [1] 10
```

```
#addition  
my_num1+my_num2
```

```
>>> [1] 20
```

```
#multiplicatoin  
my_num1*my_num2
```

```
>>> [1] 75
```

```
#division  
my_num1/my_num2
```

```
>>> [1] 3
```

```
# power  
my_num1^2
```

```
>>> [1] 225
```

```
#square root  
sqrt(my_num1 *my_num2)
```

```
>>> [1] 8.660254
```

1.3 Data types

```
# Let's review the internal representation of the variables  
  
# Numeric: default number (float)  
class(my_num1)
```

```
>>> [1] "numeric"
```

```
#Character: strings are characters  
class(my_string)
```

```
>>> [1] "character"
```

```
#Logicals: booleans  
class(my_bool)
```

```
>>> [1] "logical"
```

1.4 Lists

```
#You can create a list of variables with the **c()** command
ranks<-c(1,2,3,4)
print(ranks)
```

```
>>> [1] 1 2 3 4
```

```
# Lists preserve the class of its elements
class(ranks)
```

```
>>> [1] "numeric"
```

```
#You can access the specific elements **INDEX 1**
print(ranks[1] )
```

```
>>> [1] 1
```

```
# You can perform element-wise ops in lists
weights<-c(78.0,66.0,90.0,55.5)
heights <-c(181.0,160.0,190.0,170.0)
```

```
#e.g. division
ratios<-weights/heights
```

```
# You can have lists of characters
names<-c("John","Mary","Bob","Anna")
affiliations<-c("QF","QP","QP","QF")
# Or logicals
is_fit<-c(TRUE,FALSE,TRUE,FALSE)
```

1.5 Common functions

```
# **str**: Structure: tells you about the variable(s) class, size and values
str(my_num1)
```

```
>>> num 15
```

```
str(my_string)
```

```
>>> chr "Hello world"
```

```
str(weights)
```

```
>>> num [1:4] 78 66 90 55.5
```

1.6 Summary

```
# **summary**: Gives you a summary of a variable.  
# Specially useful for lists/vectors
```

```
summary(weights)
```

```
>>>   Min. 1st Qu.  Median    Mean 3rd Qu.    Max.  
>>>  55.50  63.38   72.00   72.38  81.00   90.00
```

1.7 Factors

Factors are a useful data type in R.

They represent categories, with a numerical correspondence (level), and a string representation (label)

```
#Let's factorize the affiliation list  
f_affiliations<-factor(affiliations)  
print(f_affiliations)
```

```
>>> [1] QF QP QP QF  
>>> Levels: QF QP
```

```
#Now let's compare  
str(f_affiliations)
```

```
>>> Factor w/ 2 levels "QF","QP": 1 2 2 1
```

```
str(affiliations)
```

```
>>> chr [1:4] "QF" "QP" "QP" "QF"
```

```
#Even summaries are different  
summary(f_affiliations)
```

```
>>> QF QP  
>>>  2  2
```

```
summary(affiliations)
```

```
>>>   Length    Class    Mode  
>>>      4 character character
```

1.8 Data Frames (DF)

They are collections of variables (columns) with different measurements(rows)

Let's create a list of people!

```
my_df <- data.frame(name=names,affiliation=affiliations,weight=weights,height=heights,is_fit=is_fit)
# The command head shows us a preview of the dataset
head(my_df)
```

```
>>>  name affiliation weight height is_fit
>>> 1 John          QF   78.0   181   TRUE
>>> 2 Mary          QP   66.0   160  FALSE
>>> 3 Bob           QP   90.0   190   TRUE
>>> 4 Anna          QF   55.5   170  FALSE
```

```
# We can see its structure
str(my_df)
```

```
>>> 'data.frame':  4 obs. of  5 variables:
>>> $ name          : Factor w/ 4 levels "Anna","Bob","John",...: 3 4 2 1
>>> $ affiliation: Factor w/ 2 levels "QF","QP": 1 2 2 1
>>> $ weight       : num  78 66 90 55.5
>>> $ height       : num  181 160 190 170
>>> $ is_fit       : logi  TRUE FALSE TRUE FALSE
```

```
# Or a complete summary
summary(my_df)
```

```
>>>  name    affiliation    weight      height      is_fit
>>> Anna:1    QF:2        Min.    :55.50    Min.    :160.0    Mode :logical
>>> Bob :1    QP:2        1st Qu.:63.38    1st Qu.:167.5    FALSE:2
>>> John:1           Median :72.00    Median :175.5    TRUE :2
>>> Mary:1           Mean   :72.38    Mean   :175.2    NA's :0
>>>           3rd Qu.:81.00    3rd Qu.:183.2
>>>           Max.    :90.00    Max.    :190.0
```

1.9 Variables in a DF

```
#print the column name
my_df$name
```

```
>>> [1] John Mary Bob  Anna
>>> Levels: Anna Bob John Mary
```

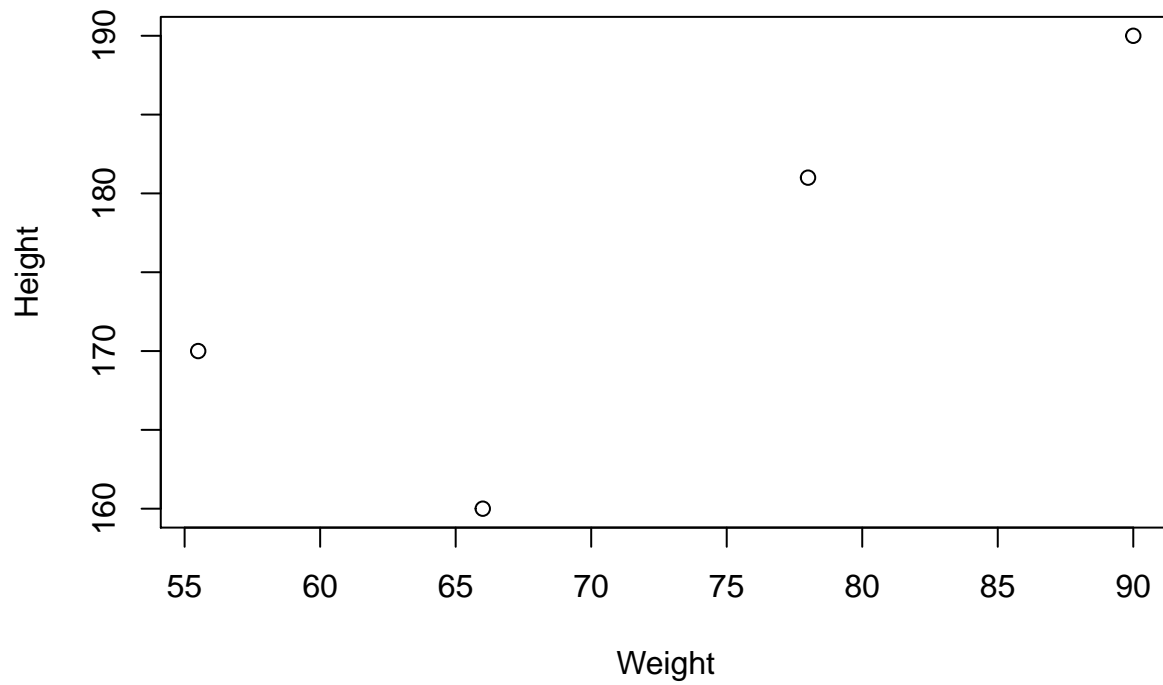
```
my_df$weight
```

```
>>> [1] 78.0 66.0 90.0 55.5
```

1.10 Basic plots

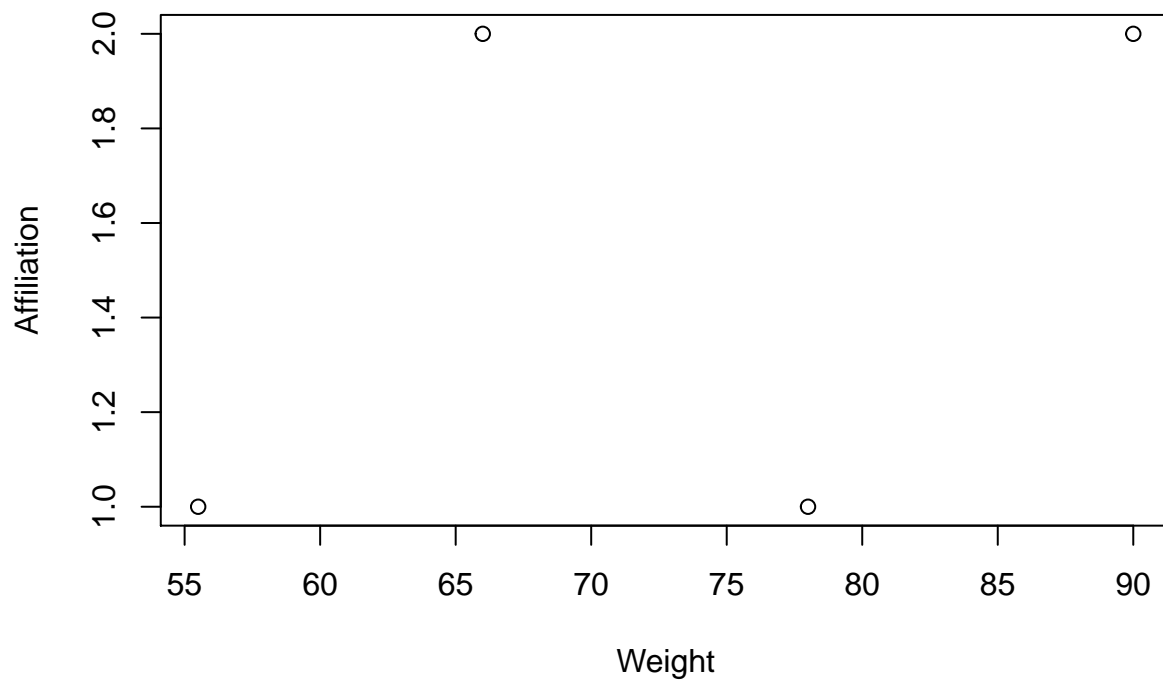
- Two numerics

```
plot(my_df$weight,my_df$height,xlab="Weight",ylab="Height")
```



- One numeric, one factor

```
#This is the wrong way because affiliation is taken as a numeric  
plot(my_df$weight,my_df$affiliation,xlab="Weight", ylab="Affiliation")
```



```
# Instead, we use the "by" operator "~".  
#Note that this operator works as y~x. So now the axes are reversed  
plot(my_df$weight~my_df$affiliation,ylab="Weight",xlab="Affiliatin")
```

