## Class 7

R – Data frame and

**Import and Export Datasets** 

## **Data Frame**

- It is a 2D array like structure
- It's a tabular data referred as a data frame in R
- Each column name considered as variable
- Used to store data table
- It is the special case of the list in which each component has equal length

## **Data Frame - Characteristics**

- Column names should not be empty
- Row names should be unique
- Values can be a numeric, factor or character type
- Each column should contain same number of data items
- The data.frame() function used for creating data frame in R

## **Data Frame - Example**

```
> #Creation of Data frame
> mydata <- data.frame(seq_id=c("Seq_1","Seq_2","Seq_3")</pre>
                       seq_name = c("DREB", "ALS", "WRKY");
                       seq_bp = c("ATCG", "TGCA", "CATG"),
                        seq_length = c(100, 200, 300)
  mydata
  seq_id seq_name seq_bp seq_length
   Seq_1
                                 100
             DREB
                    ATCG
   Seq_2
                                 200
         ALS TGCA
   Seq_3
                                 300
             WRKY
                    CATG
```

## Data Frame - Structure

```
> mydata
 seq_id seq_name seq_bp seq_length
                           100
 Seq_1 DREB
               ATCG
2 Seq_2 ALS TGCA
                           200
3 Seq_3 WRKY CATG
                           300
> str(mydata)
'data.frame': 3 obs. of 4 variables:
$ seq_id : chr "Seq_1" "Seq_2" "Seq_3"
$ seq_name : chr "DREB" "ALS" "WRKY"
$ seq_bp : chr "ATCG" "TGCA" "CATG"
 $ seq_length: num 100 200 300
```

Extract specific column from a data frame using the column name

- Extract the specific rows from data frame
- Extract data from specific rows and columns

Extract specific column from a data frame using the column name

```
mydata
seq_id seq_name seq_bp seq_length
Seq_1
      DREB ATCG
                            100
Seq_2 ALS TGCA
                            200
Seq_3 WRKY CATG
                           300
#Retrieve Based on the column
ext_data <- data.frame(mydata$seq_id,mydata$seq_name)</pre>
print(ext_data)
mydata.seq_id mydata.seq_name
       Seq_1
                       DREB
       Seq_2
                        ALS
       Seq_3
                       WRKY
```

#### Extract the specific rows from data frame

```
> mydata
 seq_id seq_name seq_bp seq_length
 Seq_1
                            100
           DREB ATCG
 Seq_2
                            200
          ALS TGCA
 Seq_3 WRKY CATG
                            300
 #Extract Based on the row
> print(mydata[1:2,])
 seq_id seq_name seq_bp seq_length
  Seq_1
                            100
           DREB
                ATCG
  Seq_2
                            200
            ALS
                 TGCA
```

#### Extract data from specific rows and columns

## Data Frame – How to check?

#### is.data.frame()

```
> mydata
 seq_id seq_name seq_bp seq_length
                                100
  Seq_1
                    ATCG
             DREB
 Seq_2
                                200
            ALS
                   TGCA
  Seq_3
                                300
             WRKY
                    CATG
> #Check Data frame
> is.data.frame(mydata)
   TRUE
```

# **Import Datasets**

## **Import Functions**

```
read.table()
read.csv()
read.csv2()
read.delim()
```

# **Export Functions**

```
write.table()
write.csv()
write.csv2()
```

## **Subset Functions**

Useful for extracting data based on the expression value.

**Basic Syntax:-**

subset(x, subset, ...)

# **Handling Excel Files**

R can read excel files using some of specific packages.

```
Few Packages:-
```

xlsx

readxl

readxlsb

install.packages("xlsx")

# Reading Excel File

```
read.xlsx(
  file,
  sheetIndex,
  sheetName = NULL,
  rowIndex = NULL,
  startRow = NULL,
  endRow = NULL,
  colIndex = NULL,
  as.data.frame = TRUE,
 header = TRUE,
  colClasses = NA,
  keepFormulas = FALSE,
  encoding = "unknown",
 password = NULL,
```