WQD7004 Programming for Data Science Lab 6 Package (dplyr)

- 1. Write R scripts for each of the following
 - a. Create a data frame using the data below.

```
id={7,11,8,14,17,13,5,9,10,1}
read={47,44,57,60,73,44,68,34,52,60}
write={59,33,44,46,57,46,65,60,52,55}
```

- b. Return a subset of rows, where read more than or equal to 55 and order the rows by read column.
- c. Return the id (descending) in vector, where write more than or equal to 50.
- d. Return a subset of rows, where duplicate value in read.
- e. Create a new column RW that measure the different between read and write.
- f. Create a new column Status that return R if RW is positive, E if RW is 0 and W if RW is negative.
- 2. Write R scripts for each of the following
 - a. Create a data frame using the data below.

```
matric={"S111", "S112", "S115", "S124", "S245", "S331"}
name={"Ahmad", "Chong", "Mutu", "Paul", "Siti", "Zaidi"}
test={26, 15, 21, 24, 24, 19}
asgn={17, 11, 13, 15, 14, 15}
post={18, 15, 17, 12, 17, 16}
final={29, 10, 19, 14, 28, 27}
```

- b. Return the name in vector, where matric begin with S1.
- c. Create a new column CA after the asgn column that add the value of test and asgn.
- d. Create a new column SA after the final column that add the value of post and final
- e. Create a new column Total that add the value SA and CA.
- f. Create a new column Grade that return the grade based on the Total value using the table below

90 – 100	A+
80 – 89	Α
75 – 79	A-
70 – 74	B+
65 – 69	В
60 – 64	B-

55 – 59	C+
50 – 54	С
45 – 49	C-
40 - 44	D+
35 – 39	D
0 - 34	F

- g. Calculate the mean and standard deviation of the subject.
- h. Get the matric and name of student with highest score and lowest score.
- i. Get the number of students that passed and failed the subject given the passing mark is at least 65.
- 3. Write R scripts for each of the following using **iris** dataset.
 - a. Create a data frame from the dataset and then display the data frame in a spreadsheet-style.
 - b. Select the versicolor species from the dataset and remove the unnecessary columns.
 - c. Sort the data frame by sepal length and width.
 - d. Sort the data frame by petal length and width.
 - e. Get the summary values for each columns.
- 4. Write R scripts for each of the following using **ToothGrowth** dataset.
 - a. Create a data frame from the dataset and then display the data frame in a spreadsheet-style.
 - b. Split the data frame into two, one with supplement type VC and one with OJ. Display both the data frame in a spreadsheet-style.
 - c. Select the tooth length and dose from the VC data frame and then sort the data frame by tooth length.
 - d. Select the tooth length and dose from the OJ data frame and then sort the data frame by tooth length (descending).
 - e. Get the minimum, maximum and average tooth length by dose for VC and OJ. Show the results in a spreadsheet-style.