

# Biostatistics: Theory and Applications in R (Virtual)

## Week8\_Session2\_R\_training8

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```
#Set the working directory- getwd()/ setwd("Y:/")
getwd()
setwd("C:/Users/Fahmida Sultana/Desktop/R training/R_training_Class_08")
```

```
#install openxlsx package or xlsx package
library(openxlsx)
library(readxl)
```

```
#####import data set from xlsx file
study_1 <- read.xlsx("Tree_height.xlsx")
str(study_1)
```

```
study_2 <- read_excel("Tree_height.xlsx", sheet = "study_2")
str(study_2)
```

## #Add plot together

#The usage of patchwork is simple: just add plots together!

```
library(ggplot2)
library(patchwork)
```

```
str(study_1)
```

```
p1<- ggplot(study_1, aes(x=treatment, y=biomass, fill=treatment)) +
  geom_bar(position="dodge", stat="identity")
```

```
p2<-ggplot(study_1, aes(x = plant, y = biomass)) +
  stat_boxplot(geom = "errorbar", # Error bars
    width = 0.2) + # Bars width
  geom_boxplot()
```

```
p3<-ggplot(study_1, aes(x = treatment, y = biomass, fill= treatment)) +
  stat_boxplot(geom = "errorbar", # Error bars
    width = 0.2) + # Bars width
  geom_violin()
```

```
p4<- ggplot(study_1, aes(x=year, y=biomass)) +
  geom_point() +
  geom_smooth(method=lm , color="red",lwd=1, se=FALSE) +
```

```
theme_gray()
```

### **#simple addition**

```
p1 + p2
```

### **#nested patchwork**

```
patch <- p1 + p2
```

```
p3 + patch
```

```
patch - p3
```

```
#or
```

```
wrap_plots(patch, p3)
```

```
wrap_plots(p3, patch)
```

### **#Adding text: position right side**

```
p1 + grid::textGrob('Some really important text')
```

```
#Adding text: position left side
```

```
# This won't do anything  
grid::textGrob('Text on left side') + p1  
#> NULL
```

```
# This will work  
wrap_elements(grid::textGrob('Text on left side')) + p1
```

### **#Stacking and packing**

```
#putting plots besides each other or on top of each other  
#patchwork provides 2 shortcut operators.
```

# "|" will place plots next to each other  
# "/" will place them on top of each other.

### **#plots next to each other**

p1 + p2  
#or  
p1 | p2

### **#on top of each other**

p1 / p2

### **#complex addition: nested plot and normal plot**

p1 / (p2 | p3)

(p1 | p2) / p3

### **#Modifying the theme of patchwork**

patchwork <- p1 / (p2 | p3)  
patchwork  
  
patchwork & theme\_minimal()

### **#Controlling Layouts**

### **#Adding an empty area**

p1 + plot\_spacer() + p2 + plot\_spacer() + p3 + plot\_spacer()

## #Controlling the grid

```
p1 + p2 + p3 + p4 +  
  plot_layout(ncol = 3)
```

```
p1 + p2 + p3 + p4 +  
  plot_layout(widths = c(1, 1)) # all figure same size
```

```
p1 + p2 + p3 + p4 +  
  plot_layout(widths = c(2, 1)) # 1st column double size and 2nd column single size
```

## #Insets

```
p4 + inset_element(p2, left = 0.6, bottom = 0.6, right = 1, top = 1)
```

## #Loading Files from Other Programs

#File format	- Function	- Library
#ERSI ArcGIS	-read.shapefile	-shapefiles
#Matlab	-readMat	-R.matlab
#minitab	-read.mtp	-foreign
#SAS (permanent data set)	-read.ssd	-foreign
#SAS (XPORT format)	-read.xport	-foreign
#SPSS	-read.spss	-foreign
#Stata	-read.dta	-foreign
#Systat	-read.systat	-foreign

## #How to get data from website

#Importing Data from Web

```
#library(dplyr)
```

```
#URL(https://raw.githubusercontent.com/ds4stats/r-tutorials/master/intro-maps/data/StatePopulation.csv)
```

```
#go to File - Import dataset - from text (readr)...- use the URL
```

```
#or
```

```
web_data <- read.csv("https://raw.githubusercontent.com/ds4stats/r-tutorials/master/intro-maps/data/StatePopulation.csv", as.is = TRUE)  
str(web_data)
```

## #Export a DataFrame to Excel File in R

```
#install.packages("writexl")
```

```
library("writexl")
```

```
#the DataFrame
```

```
web_data <- read.csv("https://raw.githubusercontent.com/ds4stats/r-tutorials/master/intro-  
maps/data/StatePopulation.csv", as.is = TRUE)  
print(web_data)
```

```
#The DataFrame name is: web_data
```

```
#get the path of wd
```

```
getwd()
```

```
#"C:/Users/Fahmida Sultana/Desktop/R training/R_training_Class_08"
```

```
#"C:\\Users\\Fahmida Sultana\\Desktop\\R training\\R_training_Class_08"
```

```
write_xlsx(web_data, "C:\\Users\\Fahmida Sultana\\Desktop\\R  
training\\R_training_Class_08\\week8.xlsx")
```

```
#or
```

```
write_xlsx(web_data, "week8_1.xlsx")
```

```
#another example
```

```
library(openxlsx)
```

```
library(readxl)
```

```
study_1 <- read.xlsx("Tree_height.xlsx")
```

```
str(study_1)
```

```
#summary statistics
```

```
library(Rmisc)
```

```
#export the sum within a data set
```

```
sum1<-aggregate(biomass~treatment+year,data=study_1,FUN=sum)  
print(sum1)
```

```
write_xlsx(sum1, "C:\\Users\\Fahmida Sultana\\Desktop\\R training\\R_training_Class_08\\sum2.xlsx")
```

```
#or  
write_xlsx(sum1, "sum2_1.xlsx")
```

```
#Export the mean, sd, se, ci within a data set
```

```
mean1 <- summarySE(study_1, measurevar="biomass", groupvars=c("treatment"), na.rm=FALSE)  
print(mean1)
```

```
write_xlsx(mean1, "C:\\Users\\Fahmida Sultana\\Desktop\\R  
training\\R_training_Class_08\\treatment.xlsx")
```

```
#or
```

```
write_xlsx(mean1, "treatment_1.xlsx")
```

```
#Export csv file
```

```
write.csv(mean1, "csvfile.csv", row.names = FALSE)
```

```
#Export data with xlsx package
```

```
library(xlsx)
```

```
mean1 <- summarySE(study_1, measurevar="biomass", groupvars=c("treatment"), na.rm=FALSE)  
print(mean1)
```

```
write.xlsx(mean1, "excelfile.xlsx", rowNames = FALSE)
```

## **#Import simple graph and make it better in PowerPoint**

```
str(study_1)
```

```
p1<- ggplot(study_1, aes(x=treatment, y=biomass)) +  
  geom_bar(position="dodge", stat="identity")
```

```
p1
```

```
p2<-ggplot(study_1, aes(x = plant, y = biomass)) +  
  stat_boxplot(geom = "errorbar", # Error bars  
              width = 0.2) + # Bars width  
  geom_boxplot()+  
  theme_bw()
```

```
p2
```