

```
##-----##
```

```
## required libraries
```

```
##
```

```
install.packages("tidyverse") #loads packages from the book
```

```
install.packages("lubridate") #functions useful for working with dates
```

```
install.packages("readxl")
```

```
library(tidyverse)
```

```
library(lubridate)
```

```
library(readxl)
```

```
##-----##
```

```
## data import
```

```
##
```

```
## before you start set your working directory to "source file location"
```

```
##
```

```
## import the "early" worksheet
```

```
early <- read_excel("mass_data.xlsx",sheet="early",skip=2) #skip first two lines
```

```
obs.date <- read_excel("mass_data.xlsx",
```

```
  sheet="early",
```

```
  col_names=FALSE,n_max=1)[[2]] #n_max> hor many lines we're reading
```

```
obs.date[[1]] #obs.date would give the same thing because it has only one element. If i had more i  
would need to specify
```

```

obs <- read_excel("mass_data.xlsx",
  sheet="early",
  col_names=FALSE,skip=1,n_max=1)[[2]] #skip one, read one (me da un elemento con la
segunda linea)
obs[[1]]

## add the date and the observation label and date as columns to the data frame

early$obs.date <- ymd(obs.date[[1]]) ## ymd() strips the time-stamp from this entry (removes time from
dates)
# compare obs.date[[1]] to ymd(obs.date[[1]])

early$obs <- obs[[1]]

early ## print the early data.frame

## import the "late" worksheet

late <- read_excel("mass_data.xlsx",sheet="late",skip=2) ## read in the 1st sheet of .xlsx
obs.date <- read_excel("mass_data.xlsx",sheet="late",col_names=FALSE,n_max=1)[2]
obs <- read_excel("mass_data.xlsx",sheet="late",col_names=FALSE,skip=1,n_max=1)[2]

late$obs.date <- ymd(obs.date[[1]])
late$obs <- obs[[1]]

late ## print the late data.frame

```

```
##-----##
```

```
## Lets put these two data frames together!
```

```
mass_data <- bind_rows(early,late) ## to use bind_rows, all column names must be the same
```

```
mass_data ## open from the global enviroment and examine
```

```
## bind_rows will only line up rows which share column names
```

```
colnames(early)
```

```
colnames(early)[1] ## print the frist element of the colnames vector
```

```
colnames(early)[2] ## the second element
```

```
colnames(early)[1]<-"plant_number" #rename column
```

```
colnames(early)
```

```
data2<-bind_rows(early,late) ## open from the global enviroment or run the line below
```

```
data2[116:124,]
```

```
##-----##
```

```
## library(tidyverse)
```

```
## library(dplyr) - it is part of tidyverse
```

```
##-----##
```

```
## filter() - select variables by a value
```

```
##
```

```
## logical operators
```

```
##
```

```
## >, <, >=, <=
```

```
## & and
```

```
## | or
```

```
## ! not
```

```
##
```

```
filter(mass_data, treatment == "control")
```

```
control_data <- filter(mass_data, treatment == "control") ## pulls out the control data
```

```
control_data
```

```
control_high_data <- filter(mass_data, treatment == "control" | treatment == "high" ) ## pulls data  
which treatment = control or high
```

```
control_high_data
```

```
early_control_block_1and3 <- filter(mass_data, treatment == "control" & obs == "early" & block != 2 ) ##  
pulls, control and early data from block 1 and 3
```

```
early_control_block_1and3
```

```
mean(early_control_block_1and3$mass) ## mean of mass from early observation control data, in blocks  
1 and 3
```

```
filter(early_control_block_1and3, mass >= 20.17) ## pulls all data greater than the mean
```

```
##-----##
```

```
## library(tidyverse)
```

```
## library(dplyr)
```

```
##-----##
```

```
## select() - pick variables by their names
```

```
##
```

mass_data ## has 8 variables (plant_id, treatment, block, rep, notes, obs.date, obs)

```
select(mass_data, plant_id,treatment,mass) ## selects 3 variables, plant_id, treatment, mass
```

```
mass_data
```

```
select(mass_data, obs, plant_id, everything()) ## selects obs and date, then puts all other variables after  
it, useful for organizing data
```

```
#pone todo el resto al final
```

```
mass_data$notes ## useless column, since it has no data lets get rid of it
```

```
mass_data <- select(mass_data, -notes) ## selects everything but the notes columbn
```

```
mass_data
```

```
##-----##
```

```
## library(tidyverse)
```

```
## library(dplyr)
```

```
##-----##
```

```
## summarize() - summarize many values
```

```
##
```

```
summarize(mass_data, mean_mass = mean(mass)) ## computes the mean of the mass, same as  
mean(data$mass) - y la llama mean_mass
```

```
mean(mass_data$mass)
```

```
## not very useful by itself, but when combined with group_by() it really shines
```

```
grouped_data <- group_by(mass_data, treatment, block, obs) ## creates a groups based on  
treatment,block,observation
```

```
grouped_data ## view this output and see -el output dice "groups"
```

```
summarize(grouped_data, mean_mass = mean(mass)) ## calculates the mean of each group
```

```
summarize(grouped_data, se=sd(mass)/sqrt(length(mass))) ## calculates the SE of each group
```

```
#SE esta calculado en lugar de ser una funcion
```

```
summarize(grouped_data, mean_mass = mean(mass), se=sd(mass)/sqrt(length(mass))) ## or put both  
statements together!
```

```
##-----##
```

```
## library(tidyverse)
```

```
## library(tidyr)
```

```
##-----##
```

```
## spread() - two columns to multiple columns
```

```
##
```

```
mass_data ## suppose we want to calculate the difference in mass between the early and late  
observation
```

```
filter(mass_data, plant_id == 1)
```

```
spread(mass_data, key = obs, value = mass) ## something here isn't right... we need to drop date or obs
```

```
spread_data <- select(mass_data, -obs.date)
```

```
spread_data <- spread(spread_data, key = obs, value = mass)
```

```
spread_data
```

```
filter(mass_data, plant_id == 1) ## check these values against the output of spread_data
```

```
filter(mass_data, plant_id == 2) ## check these values against the output of spread_data
```

```
spread_data$diff <- spread_data$late - spread_data$early ## calculate the difference between the late  
and early observation
```

```
spread_data
```

```
ggplot(spread_data, aes(treatment, diff)) + geom_boxplot() ## lets visual the differences between early  
and late observations with a boxplot
```

```
## spread() is also useful for producing readable tables
```

```
grouped_data ## recall the grouped_data
```

```
means <- summarize(grouped_data, mean_mass = mean(mass), se = sd(mass)/sqrt(length(mass)))
```

```
means$mean_mass <- round(means$mean_mass, 2) ## rounds the number to 2 sig digits
```

```
means$se <- round(means$se, 2) ## rounds the number to 2 sig digits
```

means

```
table.data <- unite(means, mean_mass, se, sep=" + ",col="mean + SE")
```

table.data

```
spread(table.data, key= obs, value = 'mean + SE') ##almost ready for the thesis
```

```
table <- spread(table.data, key= obs, value = 'mean + SE')
```

table

```
write_csv(table,"table_one.csv")
```

```
##-----##
```

```
## library(tidyverse)
```

```
## library(tidyr)
```

```
##-----##
```

```
## gather() - column names are values not variables
```

```
##
```

```
spread_data ## is this data.frame tidy? ... no because early and late are values not a variable
```

```
tidy_data <- gather(spread_data, "early", "late", key = "observation", value = "mass")
```

tidy_data

```
nrow(tidy_data) ## 240 rows.. as expected
```