

# Group Activity

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```
rm(list=ls())
```

```
library(tidyverse)
```

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.4      v readr      2.1.5
## v forcats    1.0.0      v stringr   1.5.1
## v ggplot2    3.4.4      v tibble    3.2.1
## v lubridate  1.9.3      v tidyr     1.3.1
## v purrr      1.0.2
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
```

```
## x dplyr::lag()      masks stats::lag()
```

```
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
```

```
library(tidyr)
```

```
library(readxl)
```

```
library(dplyr)
```

```
# Group Activity 01
```

```
# Question 1. Use the piggy data on d2l.
```

```
Piggy <- read_excel("~/Desktop/Data211/Week 10/Piggy.xlsx")
```

```
Piggy
```

```
## # A tibble: 16 x 6
```

```
##   pigID week0 week4 week8 week12 week16
```

```
##   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
```

```
## 1     1     1     4.8  12.6   16    21
```

```
## 2     2     1     4.2    7     10    14
```

```
## 3     3     0.8    4     6      9     10
```

```
## 4     4     0.8    4     6      9     13
```

```
## 5     5     0.8    5     9.4    11    14
```

```
## 6     6     0.8    3.2    7     10    15
```

```
## 7     7     0.8    3.2    5.5    9.6    12
```

```
## 8     8     0.8    3.4    7     8.7   12.4
```

```
## 9     9     1     5.4   10     13   17.4
```

```
## 10    10    1.2    4.8   12.6   16     20
```

```
## 11    11     1     4.6   11     18     22
```

```
## 12    12    0.8    4.2    8     11     13
```

```
## 13    13    0.8    3.8    7     8.9     12
```

```
## 14    14    1    5.4  11    14    19
## 15    15    1    6    5.4  10    17
## 16    16    1    3.4  7.8  10    13
```

```
# a) Is that data a long format or a wide format?
```

```
# - wide because you can see that the ids have single instances of each one,  
# and it is split apart into weeks.
```

```
# columns: "pigID", "week0", "week4", "week8", "week12", "week16"
```

```
Piggy_long <- pivot_longer(Piggy,  
  cols = -pigID, # Columns to pivot (excluding pigID)  
  names_to = "Time", # New column name for time  
  values_to = "Weight") # New column name for values
```

```
Piggy_long
```

```
## # A tibble: 80 x 3  
##   pigID Time   Weight  
##   <dbl> <chr>   <dbl>  
## 1     1 1 week0     1  
## 2     1 1 week4    4.8  
## 3     1 1 week8   12.6  
## 4     1 1 week12   16  
## 5     1 1 week16   21  
## 6     2 2 week0     1  
## 7     2 2 week4    4.2  
## 8     2 2 week8     7  
## 9     2 2 week12   10  
## 10    2 2 week16   14  
## # i 70 more rows
```

```
# Now Piggy_long will have a long format where each row represents one observation for each pig at a sp
```

```
# Question 2. Use the Diabetes data on d2l.
```

```
# library(readxl)
```

```
Diabetes <- read_excel("~/Desktop/Data211/Week 10/Diabetes.xlsx")
```

```
# View(Diabetes)
```

```
# a) Change the format with a variable SBP showing the systolic blood pressure at different  
# times.
```

```
# Note: it is in long format.
```

```
# Columns: "ID", "Time", "SBP"
```

```
# Pivot from long to wide format
```

```
Diabetes_wide <- pivot_wider(Diabetes,  
  id_cols = ID, # Column(s) to keep as identifier(s)  
  names_from = Time, # Values from "Time" will become column names  
  values_from = SBP) # Values from "SBP" will populate the wide format column
```

```
Diabetes_wide
```

```
## # A tibble: 18 x 4  
##   ID Baseline followone followtwo  
##   <dbl>   <dbl>   <dbl>   <dbl>  
## 1     1     130     120     115  
## 2     2     140     130     124
```

```
## 3      3      145      130      125
## 4      4      140      120      110
## 5      5      150      125      113
## 6      6      145      120      115
## 7      7      140      110      106
## 8      8      145      112      109
## 9      9      155      115      110
## 10     10     140      106      100
## 11     11     136      109      101
## 12     12     139      112      101
## 13     13     150      130      121
## 14     14     145      130      120
## 15     15     155      135      126
## 16     16     140      110      110
## 17     17     142      115      114
## 18     18     145      119      112
```

```
# Group Activity 02:
# Use the Tree data from d2l, find the mean for each variable of group, branch
# density, branch thickness, and tree heights.
# (Hint: use for loop to identify the Group, and filter() to filter each group before you
# calculate mean of each group)
```

```
Tree_data <- read_excel("~/Desktop/Data211/Week 10/Tree data.xlsx")
# View(Tree_data)

means_list <- list() # Initialize a list to store means for each group

# Loop through each group
for (i in 1:3) {
  # Filter data for the current group using filter() function
  group_data <- filter(Tree_data, Group == i)
  # Calculate mean for each variable within the group
  means <- sapply(group_data[,2:4], mean)
  # Store means in the list
  means_list[[paste0("Group_", i)]] <- means
} # end for

# Combine means from the list into a data frame
means_df <- do.call(rbind, means_list)
# Rename rows and columns
row.names(means_df) <- paste0("Group_", 1:3)
# Rename columns
colnames(means_df) <- c("Meandensity", "Meanthickness", "Meanheight")
# Print the means for each group
means_df
```

```
##           Meandensity Meanthickness Meanheight
## Group_1      3.775           3.40         13.5
## Group_2      4.600           2.85         12.5
## Group_3      2.225           6.55         12.5
```