

# R Scripting

## Lab for unit 4 - Structured programming

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Please solve the following problems!

1. The `CO2` dataset (included in R) contains information about the uptake of carbon dioxide by different types of plants exposed to different treatments.
  - a. Suppose we wish to subtract the mean value of the `uptake` variable in the `CO2` data frame, where the mean is calculated separately for each `Type/Treatment` combination using `tapply()`. Save these means in an object called `means`.
  - b. Next, we need an index vector that specifies which mean has to be subtracted from which observation. Use the *same* (identical) call to `tapply()` you used for the computation of the group means above, but without specifying a function to be applied. Save the resulting vector as `idx`.
  - c. Compute the group mean-adjusted CO2 uptake for each plant.
  - d. We used two `tapply()` calls to reach our goal, but this can be simplified using the `ave()` function that combines the two operations instead (i.e., `ave()` replaces `tapply()`). Change the code accordingly and check the result.

```
# a.
means <- tapply(CO2$uptake, CO2[c("Type", "Treatment")], mean)
means
```

```
##           Treatment
## Type      nonchilled chilled
##  Quebec      35.33333 31.75238
##  Mississippi 25.95238 15.81429
```

```
# b.
idx <- tapply(CO2$uptake, CO2[c("Type", "Treatment")])
idx
```

```
## [1] 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 3 3 3 3 3 3 3 3 3 3 3 3 3
## [39] 3 3 3 3 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 4 4 4 4 4 4 4 4 4 4 4
## [77] 4 4 4 4 4 4 4 4 4
```

```
# c.
adj.uptake <- CO2$uptake - means[idx]
adj.uptake
```

```
## [1] -19.33333333 -4.93333333 -0.53333333 1.86666667 -0.03333333
## [6] 3.86666667 4.36666667 -21.73333333 -8.03333333 1.76666667
## [11] 6.46666667 5.26666667 6.06666667 8.96666667 -19.13333333
## [16] -2.93333333 4.96666667 6.76666667 7.56666667 8.56666667
## [21] 10.16666667 -17.55238095 -7.65238095 -1.45238095 2.84761905
## [26] 0.74761905 3.64761905 6.94761905 -22.45238095 -4.45238095
## [31] 3.24761905 7.04761905 6.84761905 5.74761905 10.64761905
## [36] -16.65238095 -10.75238095 6.34761905 2.24761905 7.14761905
## [41] 7.84761905 9.64761905 -15.35238095 -6.75238095 0.24761905
## [46] 4.04761905 4.94761905 6.44761905 9.54761905 -13.95238095
## [51] -3.95238095 4.64761905 5.84761905 6.44761905 5.14761905
## [56] 5.54761905 -14.65238095 -6.55238095 -0.15238095 1.94761905
## [61] 2.54761905 2.14761905 1.84761905 -5.31428571 -0.91428571
## [66] 2.28571429 3.08571429 3.68571429 6.38571429 6.08571429
## [71] -8.11428571 -4.41428571 -3.51428571 -2.81428571 -3.31428571
## [76] -2.11428571 -1.41428571 -5.21428571 2.18571429 2.08571429
## [81] 2.08571429 2.08571429 3.08571429 4.08571429
```

```
# d.
adj.uptake <- C02$uptake - ave(C02$uptake, C02$Type, C02$Treatment, FUN = mean)
adj.uptake
```

```
## [1] -19.33333333 -4.93333333 -0.53333333 1.86666667 -0.03333333
## [6] 3.86666667 4.36666667 -21.73333333 -8.03333333 1.76666667
## [11] 6.46666667 5.26666667 6.06666667 8.96666667 -19.13333333
## [16] -2.93333333 4.96666667 6.76666667 7.56666667 8.56666667
## [21] 10.16666667 -17.55238095 -7.65238095 -1.45238095 2.84761905
## [26] 0.74761905 3.64761905 6.94761905 -22.45238095 -4.45238095
## [31] 3.24761905 7.04761905 6.84761905 5.74761905 10.64761905
## [36] -16.65238095 -10.75238095 6.34761905 2.24761905 7.14761905
## [41] 7.84761905 9.64761905 -15.35238095 -6.75238095 0.24761905
## [46] 4.04761905 4.94761905 6.44761905 9.54761905 -13.95238095
## [51] -3.95238095 4.64761905 5.84761905 6.44761905 5.14761905
## [56] 5.54761905 -14.65238095 -6.55238095 -0.15238095 1.94761905
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## [66] 2.28571429 3.08571429 3.68571429 6.38571429 6.08571429
## [71] -8.11428571 -4.41428571 -3.51428571 -2.81428571 -3.31428571
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## [81] 2.08571429 2.08571429 3.08571429 4.08571429
```