# Assignment 2

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```
{r setup, include=FALSE} # install.packages("ggplot2") # #
library(ggplot2) # # knitr::opts_chunk$set(echo = FALSE) #
```

#### Exercise 2.1

a) The data is read in with the following command:

```
data_sat = read.table("./sat.txt", header=TRUE)
expend = data_sat$expend
ratio = data_sat$ratio
salary = data_sat$salary
takers = data_sat$takers
total = data_sat$total
```

The first technique is the step-up method. This method examines the statistics for each variable and selects the option with the highest r-squared value, given the p-value is significant. This procedure is repeated until the added variable's p-value is no longer statistically significant. The outcomes are displayed below.

```
# Step-up method
# First round
summary(lm(total~expend)) # multipled R^2 = 0,1448, p-value = 0.00641
summary(lm(total~ratio)) # multipled R^2 = 0.006602, p-value = 0.575
summary(lm(total~salary)) # multipled R^2 = 0.1935, p-value = 0.00139
summary(lm(total~takers)) # multipled R^2 = 0.787, p-value = <2e-16
# Second round
summary(lm(total~takers+expend)) # multipled R^2 = 0.8195, p-value = 0.00553
summary(lm(total~takers+ratio)) # multipled R^2 = 0.7991, p-value = 0.0982
summary(lm(total~takers+salary)) # multipled R^2 = 0.8056, p-value = 0.0394
# Third round
summary(lm(total~takers+expend+ratio)) # multipled R^2 = 0.8227, p-value = 0.3629
summary(lm(total~takers+expend+salary)) # multipled R^2 = 0.8196, p-value = 0.8527</pre>
```

The variable 'takers' will be chosen in the first round and the variable 'expend' in the second round. We stop after the second round because the p-values for both the ratio and the salary are not

statistically significant. Consequently, the step-up method model will be lm(total~takers+expend). The step-down method does not add variables, but instead begins with a model that contains all variables and then removes options. These are selected using the highest non-significant p-value. The calculation is displayed below.

```
# Step-down method
# First round
summary(lm(total~takers+expend+ratio+salary))
# Second round
summary(lm(total~takers+ratio+salary))
```

The variable 'expend' is removed in the first round because it has the highest non-significant p-value. There is no further removal of variables as all variables have significant p-values. The statistics for the step-up method are as follows: multiple r-squared = 0.8195 and p-value = < 2.2e-16. For the step-down method it is: multiple-r^2 = 0.8239 and p-value = < 2.2e-16 Given these numbers, the model with the greatest r-squared multiple would be preferred. However, as the difference is very small, we also consider the model's complexity. The step-up method utilizes only two variables, whereas the step-down method has three. Therefore, the former is easier. In this situation, we would prefer the step-up method due to the fact that their performance is nearly identical.

b) The variable takers 2 is the square of the takers values. This is followed by the step-up method.

```
takers2=(data_sat$takers)^2
takers; takers2
# Step-up method
# First round
summary(lm(total~expend)) # multipled R^2 = 0.1448, p-value = 0.00641
summary(lm(total~ratio)) # multipled R^2 = 0.006602, p-value = 0.575
summary(lm(total~salary)) # multipled R^2 = 0.1935, p-value = 0.00139
summary(lm(total~takers)) # multipled R^2 = 0.787, p-value = <2e-16
summary(lm(total~takers2)) # multipled R^2 = 0.6578, p-value = 9.28e-13
# Second round
summary(lm(total~takers+expend)) # multipled R^2 = 0.8195, p-value = 0.00553
summary(lm(total~takers+ratio)) # multipled R^2 = 0.7991, p-value = 0.0982
summary(lm(total~takers+salary)) # multipled R^2 = 0.8056, p-value = 0.0394
summary(lm(total~takers+takers2)) # multipled R^2 = 0.8732, p-value = 8.96e-07
# Third round
summary(lm(total~takers+takers2+expend)) # multipled R^2 = 0.8859, p-value = 0.0285
summary(lm(total~takers+takers2+ratio)) # multipled R^2 = 0.8738, p-value = 0.634
summary(lm(total~takers+takers2+salary)) # multipled R^2 = 0.8858, p-value = 0.029
# Fourth round
summary(lm(total~takers+takers2+expend+ratio)) # multipled R^2 = 0.8887, p-value = 0.2936
summary(lm(total~takers+takers2+expend+salary)) # multipled R^2 = 0.8873, p-value = 0.466
```

In the first round, the variable 'takers' is selected, followed by the variable 'takers2' and finally the variable 'expend'. We stop after the third round because the variables added in the fourth round do not have a significant p-value.

The step-down method is shown below:

```
# First round
summary(lm(total~takers+takers2+expend+ratio+salary)) # remove salary since it has the highest
# Second round
summary(lm(total~takers+takers2+expend+ratio)) # remove ratio as it has the highest p-value th
# Third round
summary(lm(total~takers+takers2+expend)) # all variables have a significant p-value, so stop r
```

The variable takers 2 is included in both the step-up and step-down models, making its addition useful. Both have got three variables so they are equally complex. Since both models are identical, it makes no difference which method is chosen, as both will result in the same model.

c) The model in exercise a has an r-squared multiple of 0.8194 and a p-value less than 2.2e-16. Multiple r-squared value of 0.8859 and a p-value less than 2.2e-16 are the statistical results for the model selected in exercise b. However, the first model uses only two variables, while the second model utilises three. We prefer the multiple r-squared value over the complexity of the model and thus select the second model.

d)

```
summary(lm(total~takers+takers2+expend))
chosen_model = lm(total~takers+takers2+expend)
# model = 1052 - 6.381 * takers + 0.04741 * takers2 + 7.914 * expend
newxdata = data.frame(expend=5, takers2=625, salary=36000, takers=25)
predict(chosen_model,newxdata,interval="prediction",level=0.95)
# Fit = 961.5703; lwr = 907.6003; upr = 1015.54;
```

The fit is 961.57, the lowerbound 907.6 and the upperbound 1015.54.

## Exercise 2.2

```
df = read.table("./treeVolume.txt", header=TRUE)
```

**a**)

We assume that there is no hidden relationship between measurements. We also need to know if the data is normalized.

```
shapiro.test(df$diameter);shapiro.test(df$volume);shapiro.test(df$height);
```

```
##
## Shapiro-Wilk normality test
##
## data: df$diameter
## W = 0.97858, p-value = 0.3825
```

```
##
##
     Shapiro-Wilk normality test
##
## data:
            df$volume
## W = 0.94646, p-value = 0.01156
##
##
     Shapiro-Wilk normality test
##
## data: df$height
## W = 0.97876, p-value = 0.3895
par(mfrow=c(3,3));qqnorm(df$diameter, main="QQ-plot diameter");qqnorm(df$volume, main="QQ-plot
          QQ-plot diameter
                                              QQ-plot volume
                                                                                  QQ-plot height
Sample Quantiles
                                   Sample Quantiles
                                                                       Sample Quantiles
                                                                           82
                                        70
    18
                                        5 =
                                                                           92
                           2
                                                      0
                                                                                                  2
           Theoretical Quantiles
                                              Theoretical Quantiles
                                                                                 Theoretical Quantiles
         Histogram diameter
                                                                                 Histogram height
                                             Histogram volume
                                   Frequency
Frequency
                                                                       Frequency
                                                                           15
                        20
           10
                 15
                                            10
                                                  30
                                                       50
                                                             70
                                                                               60
                                                                                      70
                                                                                             80
                                                                                                    90
               df$diameter
                                                   df$volume
                                                                                       df$height
          Boxplot diameter
                                              Boxplot volume
                                                                                  Boxplot height
                                                                           65
```

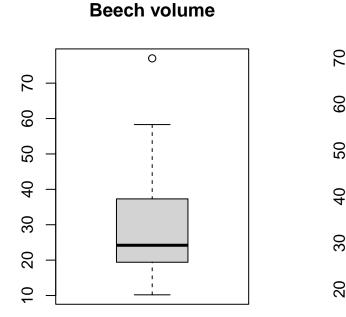
The data follows a fairly normal distribution, as far as we can know with the limited dataset. #{r} #plot(df\$diameter, df\$height,col=factor(df\$type,labels=c("blue","red"))) # Next, we will model the differences in the mean of the volume, as a function of the type of tree:

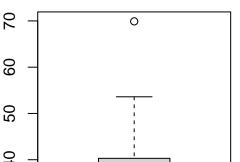
```
# Does the type affect the volume?
fit = aov(volume ~ type, data=df)
summary(fit)
```

```
## Df Sum Sq Mean Sq F value Pr(>F)
## type 1 380 379.5 1.898 0.174
## Residuals 57 11395 199.9
```

The F value is not high, which means there is little variation between groups. Furthermore, the p value is not below 0.05. Therefore it seems (in this test) that the tree type does not have an significant effect on the volume.

```
par(mfrow=c(1,2));boxplot(df[df$type == "beech",]$volume, main="Beech volume");boxplot(df[df$type
```





Oak volume

However, graphing the data shows that an oak does in fact seem to have a higher mean volume compared to a beech, just not significant enough. The variance looks about the same.

b) Now, we'll include diameter and height into the analysis, performing an ANCOVA test:

```
lm2 = lm(volume ~ diameter + height + type, data=df)
fit2 = anova(lm2)
fit2
```

```
1 10826.5 10826.5 1029.5139 < 2.2e-16 ***
## diameter
                          346.2
                                  32.9192 4.254e-07 ***
## height
              1
                  346.2
## type
                   23.2
                           23.2
                                   2.2083
                                              0.143
              1
## Residuals 55
                  578.4
                           10.5
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
```

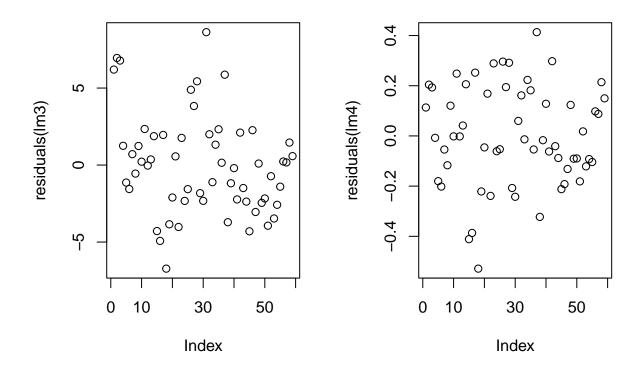
Now, we see that the for the diameter, and to a certain point the height, the F values are high, indicating large differences between groups. Furthermore, the p-values are very small, noting statistical significance.

```
predict(lm2, newdata=data.frame(height=mean(df$height), diameter=mean(df$diameter), type="beed
##
          1
## 33.20049
predict(lm2, newdata=data.frame(height=mean(df$height), diameter=mean(df$diameter), type="oak"
##
          1
## 31.89589
lm2
##
## lm(formula = volume ~ diameter + height + type, data = df)
##
## Coefficients:
## (Intercept)
                                  height
                                               typeoak
                   diameter
                                   0.4172
                                               -1.3046
##
      -63.7814
                     4.6981
```

c) We can square the volume, as the diameter and height can be seen similar to a multiplication.

```
lm3 = lm(volume ~ diameter + height, data=df)
lm4 = lm(sqrt(volume) ~ diameter + height, data=df)

par(mfrow=c(1,2))
plot(residuals(lm3))
plot(residuals(lm4))
```

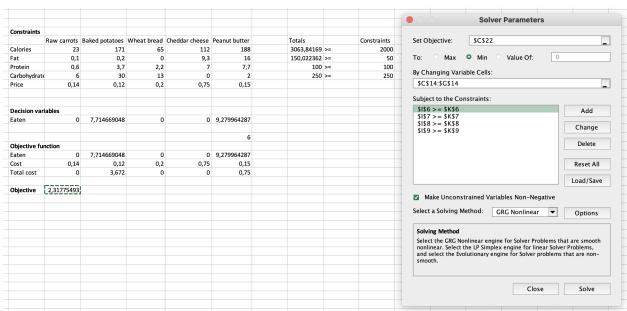


By doing so we can see the residuals going down a lot, which would improve the fit.

## Exercise 2.3

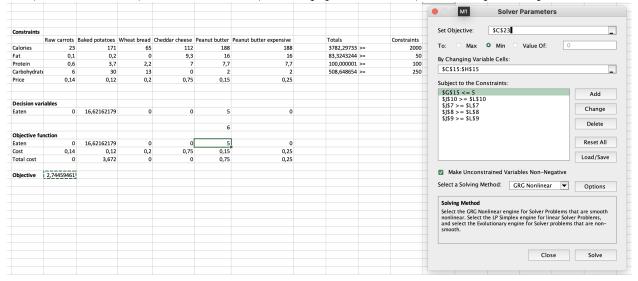
a) The calculation made in cell C23 is '=SUMPRODUCT(C19:G19;C20:G20)'. For cell I7 it is '=SUMPRODUCT(C7:G7;C19:G19)', cell I8 '=SUMPRODUCT(C8:G8;C19:G19)', cell I9 '=SUMPRODUCT(C9:G9;C19:G19)' and cell I10 '=SUMPRODUCT(C10:G10;C19:G19)'. The image depicts the calculations performed in Excel solver.

The products and nutrients are swapped because it was simpler to construct the Excel solver in this manner. The solution is 0 raw carrots, 7,71 baked potatoes, 0 wheat bread, 0 cheddar cheese, and 9,28

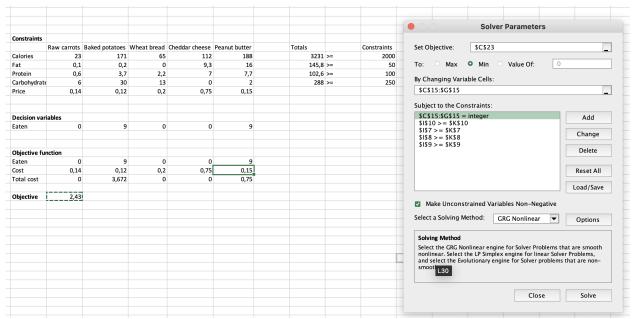


peanut butter.

b) The same calculations in cells I7, I8, I9, and I10 are made as was done in exercise a), but cell C23 is now '=SUMPRODUCT(C19:H19;C20:H20)'. The calculation has been modified slightly because the price of peanut butter increases after the fifth unit. Therefore, a new column ('Peanut butter expensive') containing the same nutrients but at a higher price is added. Additionally, a new constraint has been added so that the quantity of inexpensive peanut butter units cannot exceed five units. The solution is 0 raw carrots, 16,62 baked potatoes, 0 wheat bread, 0 cheddar cheese, 5 cheap peanut butter, and 0 expensive peanut butter.



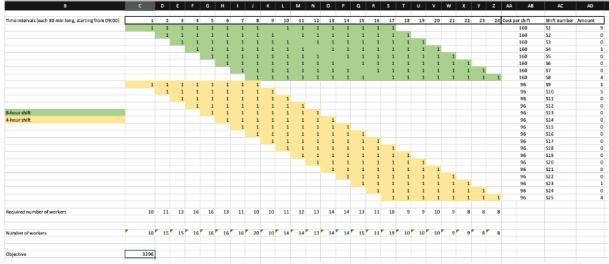
c) The data used in this exercise is the same as the one in exercise a). The calculations made in this exercise are also the same, but with the additional constraint that the decision variables must be integers. The solution is 0 raw carrots, 9 baked potatoes, 0 wheat bread, 0 cheddar cheese, and 9 peanut



butter.

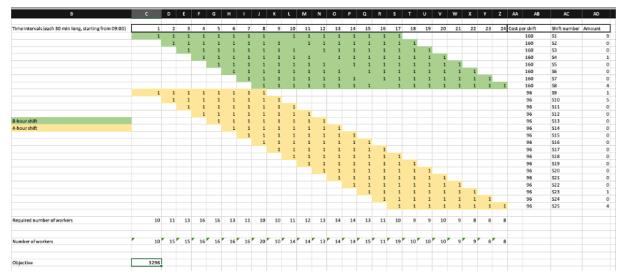
### Exercise 2.5

a) The problem is formulated as an Integer Linear Programming (ILP) problem, using the following



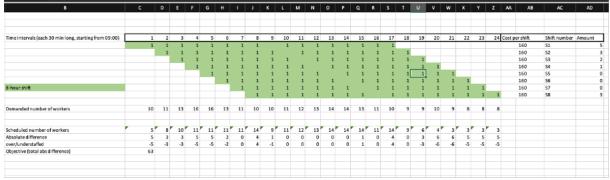
setup:

In this setup, all possible shifts (8-hour and 4-hour shifts starting between mentioned times) are presented in a matrix format. The cost per shift (# hours \* hourly wage) and the required number of workers can also be found in this setup. The amount of shifts of a specific type can be found in the column "amount". The number of workers per interval is corresponding to the amount of shifts per type. Lastly, our objective is the total costs (sum of (amount per shift type\* cost per shift)). By adjusting the column "amount" (decision variables) with the constraints that the amounts should be integers and that the number of workers satisfy the minimum requirement, we are minimizing our objective (total costs). This is done by using the integrated solver of Excel, as follows:



The resulting objective value is 3296. This denotes that the minimum cost to cover all intervals with the required number of workers is equal to 3296 euro.

b) To answer this question, the same formulation as above is used with some adjustments: -4-hour shifts are removed. - Instead of a required number of workers, there is a demanded number of workers. This changes the problem in the way that it is also possible that in some cases the demanded number of workers is not reached (understaffed). In part a, it was only possible to have more scheduled workers than required (overstaffed), because the required number was a hard minimum. Now both situations are possible. - The objective value is now the total absolute difference between the demanded number of workers and the scheduled number of workers. By solving this ILP, we are minimizing this difference. A low objective value corresponds to reliability in scheduling the demanded amount of workers.



As before, the integrated solver of excel (with the adjustments above) is used to solve this problem. As can be seen, the resulting objective value is 63. So the schedule that minimizes the sum of absolute differences between the demanded and scheduled amount of workers has an outcome of 63.