

Homework 4

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Perpare R envrioment

```
library(leaps)
library(knitr)
library(reshape2)
library(ggplot2)
library(grid)
library(plyr)
opts_chunk$set(fig.width = 8, message = FALSE, warning = FALSE)
```

Data preparation

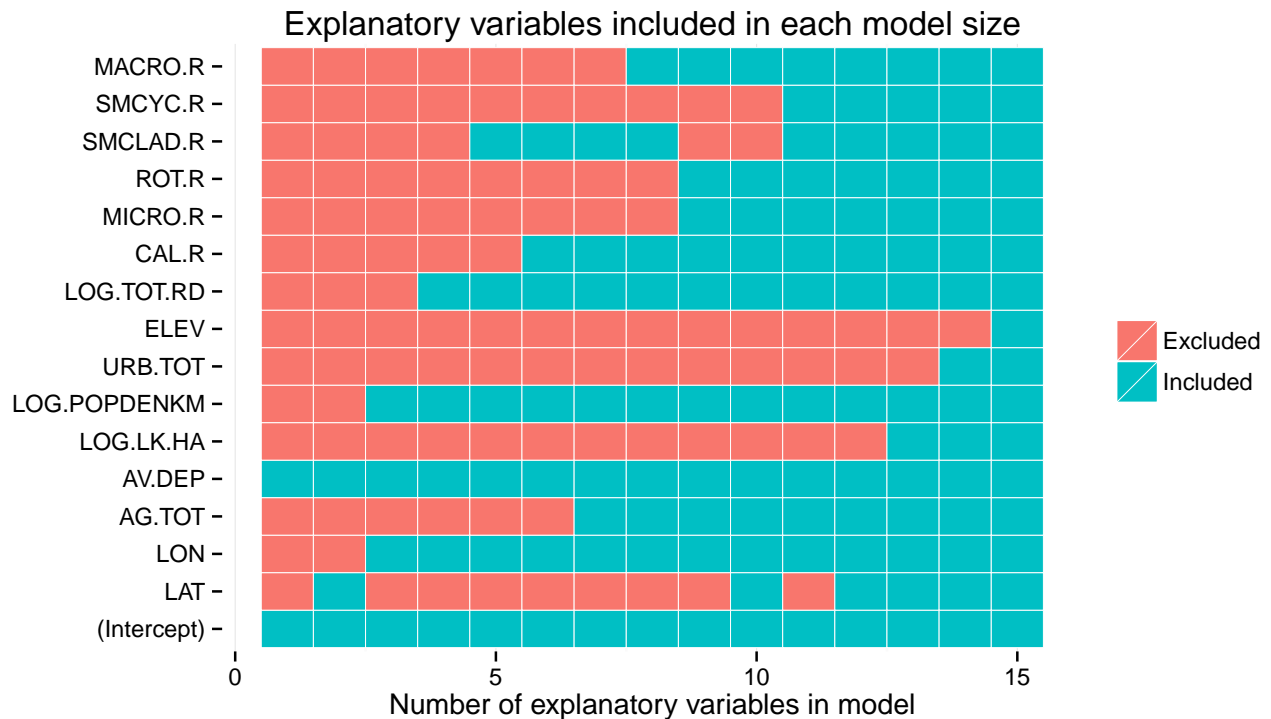
```
data <- read.table("emap.build08.txt", header = TRUE)
vars_to_log <- c("LK.HA", "POPDENKM", "TOT.RD")
data[vars_to_log] <- lapply(data[vars_to_log], function(x) log(x + 1))
names(data) <- ifelse(names(data) %in% vars_to_log,
                      paste("LOG", names(data), sep = "."),
                      names(data))
explanatory <- data[, -3]
response <- data[, 3]
```

Naive all subsets selction without interactions

The `regsubsets` function from the `leaps` package provides a simple way of doing all subsets selection.

```
# Conduct all subsets model evaluation -----
naive_raw <- regsubsets(explanatory, response, nvmax = length(explanatory))
naive_results <- summary(naive_raw)

# Graph all subsets results -----
models <- melt(naive_results$which,
              varnames = c("size", "variables"),
              value.name = "included")
models$included = ifelse(models$included, "Included", "Excluded")
ggplot(models, aes(y = variables, x = size)) +
  geom_tile(aes(fill = included), color = "white") +
  labs(y = "",
       x = "Number of explanatory variables in model",
       fill = "",
       title = "Explanatory variables included in each model size") +
  theme_minimal() + theme(panel.grid.major.y = element_blank())
```



```
# Graph information criteria scores -----
scores <- data.frame(naive_results[c("rsq", "rss", "adjr2", "cp", "bic")])
scores$size <- 1:nrow(scores)
calculate_aic <- function(included) {
  vars <- explanatory[included]
  model <- as.formula(paste("response ~", paste("explanatory", names(vars),
                                                collapse = " + ", sep = "$")))

  AIC(lm(model))
}
scores$aic <- apply(naive_results$which[, -1], 1, calculate_aic)
scores <- melt(scores, variable.name = "method", id.vars = "size")
is_best <- function(x) {
  if (x$method %in% c("rsq", "adjr2"))
    x$value == max(x$value)
  else x$value == min(x$value)
}
scores$best <- as.numeric(unlist(dplyr(scores, "method", is_best)))
scores$size <- ordered(scores$size)
ggplot(scores, aes(x = size, y = value, color = best, size = best)) +
  geom_point(stat = "identity") +
  facet_grid(method ~ ., scales = "free_y") +
  scale_size(range = c(5, 8)) +
  scale_y_continuous(expand = c(.2, 0)) +
  labs(y = "Score",
       x = "Number of explanatory variables in model",
       title = "Model selection criteria scores for best model of each size") +
  theme(legend.position = "none",
        panel.grid.minor = element_blank())
```

