CompStat/R - Paper 3

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Part I: Linear regression

In this first part of the paper, we will program a function which estimates the unknown parameters β and σ of a (ordinary) linear regression model

$$y = X\beta + \varepsilon, \qquad \varepsilon \sim N(0, \sigma^2 I)$$

by the ordinary least squares (OLS) method. For a given design matrix X and response vector y the OLS estimator is given by

$$\hat{\beta} = (X'X)^{-1}X'y \tag{1}$$

with covariance matrix

$$Var(\hat{\beta}) = \sigma^2 (X'X)^{-1} \tag{2}$$

where σ^2 has to be estimated via the sum of squared residuals (SSR):

$$\hat{\sigma}^2 = \frac{SSR}{df} = \frac{\sum_i (y_i - x_i'\hat{\beta})^2}{n - k}.$$
 (3)

The term df refers to the degrees of freedom, i.e. the difference between the number of observations n and the number of coefficients k.

Raw implementation

The function linModEst is a raw implementation of the OLS estimator. The function takes the response vector y(y) and design matrix X(x) as arguments and returns a list with the following named elements:

- coefficients: the estimated coefficients $\hat{\beta}$
- vcov: the estimated covariance matrix $\operatorname{Var}(\hat{\beta})$
- sigma: the square root of the estimated scale parameter $\hat{\sigma}^2$
- df: the degrees of freedom df

We use equations (1), (2), and (3) for the implementation and compute the inverse of X'X using the solve function, which numerically solves the equation

$$(X'X) A = I$$

for the matrix $A = (X'X)^{-1}$. To efficiently compute X'X and X'y, we use the crossprod function.

```
linModEst <- function(x, y) {</pre>
  # Computes the OLS estimator and sample variance assuming a (ordinary) linear
  # regression model.
  # Args:
    x: design matrix x
     y: response vector y
  # Returns:
  #
     A list with the following named elements:
        $coefficients: the estimated coefficients
  #
  #
        $vcov: the estimated covariance matrix
        $sigma: the square root of the estimated variance
  #
        $df: the degrees of freedom in the model, i.e. the difference between
             the number of rows and columns of x
  # Compute the inverse of (x'x) using the solve- and crossprod-function
  inv <- solve(crossprod(x), diag(nrow = ncol(x)))</pre>
  # Compute beta hat, i.e. the estimated coefficients
  coefficients <- inv %*% crossprod(x, y)</pre>
  # Compute the degrees of freedom
  df \leftarrow nrow(x) - ncol(x)
  # Compute the sample variance via the sum of squared residuals (SSR)
  SSR <- sum((y - x %*% coefficients)^2)
  sigmaSquared <- SSR / df
  # Compute the covariance matrix
  vcov <- sigmaSquared * inv
  # Create named results list to be returned
  results <- list(coefficients, vcov, sqrt(sigmaSquared), df)
  names(results) <- c("coefficients", "vcov", "sigma", "df")</pre>
  # Return results
  results
}
```

We test our implementation by computing the linear relationship between heart weight, body weight and sex for the cats dataset contained in the package MASS. In the following piece of code, cbind combines its arguments by columns into a matrix with the number of columns given by the number of arguments and the number of rows given by the greatest length of the given arguments. Shorter arguments are repeated, as long as the matrix number of rows is a multiple of the shorter vector lengths. Hence, cbind(1, cats\$Bwt, as.numeric(cats\$Sex) - 1) creates a design matrix with an intercept column, the variable body weight (bwt), and the variable sex (Sex), which is converted from a factor into a dummy variable using as.numeric. We subtract 1 to receive dummy variable values of 0 and 1, rather than 1 and 2 from the original data. Thus, cbind is used to build a proper design matrix of object type matrix with an intercept and dummy variable, such that our implementation of linModEst works correctly.

```
# Load cats dataset
data(cats, package = "MASS")
# Compute OLS using our implementation
linModEst(
 x = cbind(1, cats$Bwt, as.numeric(cats$Sex) - 1),
 y = cats$Hwt
## $coefficients
               [,1]
## [1,] -0.41495263
## [2,] 4.07576892
## [3,] -0.08209684
##
## $vcov
##
                           [,2]
                                        [,3]
               [,1]
## [1,] 0.52900070 -0.20504763 0.06563743
## [2,] -0.20504763  0.08690026 -0.04696312
## [3,] 0.06563743 -0.04696312 0.09244480
##
## $sigma
## [1] 1.457138
##
## $df
## [1] 141
```

We verify our results by comparing them to the output of R's lm function:

```
summary(lm(Hwt ~ Bwt + Sex, data = cats))
```

```
##
## Call:
## lm(formula = Hwt ~ Bwt + Sex, data = cats)
##
## Residuals:
      Min
               1Q Median
                               3Q
                                      Max
## -3.5833 -0.9700 -0.0948 1.0432 5.1016
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.4149
                          0.7273 -0.571
                                             0.569
                4.0758
## Bwt
                           0.2948 13.826
                                            <2e-16 ***
               -0.0821
## SexM
                           0.3040 -0.270
                                             0.788
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.457 on 141 degrees of freedom
## Multiple R-squared: 0.6468, Adjusted R-squared: 0.6418
## F-statistic: 129.1 on 2 and 141 DF, p-value: < 2.2e-16
```

As we can see, our implementation is correct.

Extend implementation

In this section, we write a new function linMod(formula, data), which estimates a linear regression model specified by formula and uses our linModEst function defined above to estimate the model parameters again by the OLS method. linMod returns a list with the following named elements:

• coefficients: named vector of the estimated coefficients $\hat{\beta}$ • vcov: named estimated covariance matrix $\widehat{\mathrm{Var}}(\hat{\beta})$ • sigma: the square root of the estimated scale parameter $\hat{\sigma}^2$ • df: the degrees of freedom df• formula: the formula that represents the model equation
• call: the arguments with which linMod was called

Below, we use the model.frame, model.extract, and model.matrix functions, which are very convenient for working with objects of the class formula. model.frame returns a data.frame containing only the variables from its passed data argument, which are used in the formula expression given. The returned data.frame from the model.frame function has additional attributes, but these are not needed in our application. With model.extract, we are able to extract the response variable from the data.frame created by model.frame. Moreover, using model.matrix, we can create the design matrix (of object class matrix) again only from formula and data arguments. By default, the matrix returned by model.matrix includes an intercept and converts factor variables into proper dummy variables (i.e. a factor variable with L levels results in L-1 dummy variables). More precisely, the default intercept is taken over from the formula object, which then by default adds an intercept term to the model equation, if not specified otherwise. Finally, we use match.call to return the call of our function with all the specified arguments by their full names.

```
linMod <- function(formula, data) {</pre>
  # Computes the OLS estimator and sample variance assuming a (ordinary) linear
  # regression model with model equation specified by the formula-argument.
  #
  # Args:
      formula: a formula specifying the linear model equation
  #
      data: a data.frame, list or environment, containing the variables used in
            formula
  #
  # Returns:
     A list with the following named elements:
        $coefficients: named vector of the estimated coefficients
  #
        $vcov: named estimated covariance matrix
  #
        $sigma: the square root of the estimated variance
        $df: the degrees of freedom in the model
        $formula: the formula that represents the model equation
        $call: the arguments with which the function was called
  # Extract the response variable using the model.extract function on the
  # data.frame returned by model.frame
  y <- model.extract(model.frame(formula, data = data), "response")
  # Create the design matrix using model.matrix, which overtakes an intercept
  # specified in the formula argument by default and converts factor variables into proper
  # dummy variables
  x <- model.matrix(formula, data = data)
  # Use previously defined linModEst for estimation
```

```
tmp <- linModEst(x, y)

# Prepare the output
rownames(tmp$coefficients) <- colnames(x)
colnames(tmp$vcov) <- colnames(x)
rownames(tmp$vcov) <- colnames(x)

# Create results list to be returned
results <- c(tmp, formula, match.call())
names(results) <- c("coefficients", "vcov", "sigma", "df", "formula", "call")

# Return results
results
}</pre>
```

Let's again test our implementation:

```
linMod(Hwt ~ Bwt + Sex, data = cats)
```

```
## $coefficients
                    [,1]
## (Intercept) -0.41495263
## Bwt
             4.07576892
             -0.08209684
## SexM
##
## $vcov
##
             (Intercept)
                                Bwt
                                          SexM
## (Intercept) 0.52900070 -0.20504763 0.06563743
             ## SexM
              0.06563743 -0.04696312 0.09244480
##
## $sigma
## [1] 1.457138
##
## $df
## [1] 141
##
## $formula
## Hwt ~ Bwt + Sex
##
## $call
## linMod(formula = Hwt ~ Bwt + Sex, data = cats)
```

As we can see, the output has the desired format and the correct results.

Part II: S3 for linear models

In this section we will expand upon our linear regression function using one of R's object oriented systems: S3. Our goal is to improve the function by returning a more concise output and ultimately replicating the results of the 1m function. We'll start by redefining linMod from Part I and assigning it the class linMod.

```
# Begin with the same linMod function as in Part I
linMod <- function(formula, data) {
    y <- model.extract(model.frame(formula, data = data), "response")
    x <- model.matrix(formula, data = data)
    tmp <- linModEst(x, y)

    rownames(tmp$coefficients) <- colnames(x)
    colnames(tmp$cov) <- colnames(x)
    rownames(tmp$vcov) <- colnames(x)

    results <- c(tmp, formula, match.call())
    names(results) <- c("coefficients", "vcov", "sigma", "df", "formula", "call")

# Use the class function to redefine the results with class "linMod"
    class(results) <- "linMod"

# Return results (of class "linMod")
    results
}</pre>
```

Now we'd like to define a printing method for all objects of class linMod so that the function returns a more readable and concise output.

Let's check the structure of linMod:

```
# Define the model to be estimated from Part I
modelFit <- linMod(Hwt ~ Bwt + Sex, data = cats)

# Verify the class of modelFit as well as the objects it contains
str(modelFit)</pre>
```

```
## List of 6
  $ coefficients: num [1:3, 1] -0.415 4.0758 -0.0821
    ..- attr(*, "dimnames")=List of 2
    ....$ : chr [1:3] "(Intercept)" "Bwt" "SexM"
##
##
    .. ..$ : NULL
##
  $ vcov
                 : num [1:3, 1:3] 0.529 -0.205 0.0656 -0.205 0.0869 ...
    ..- attr(*, "dimnames")=List of 2
    ....$ : chr [1:3] "(Intercept)" "Bwt" "SexM"
##
##
    ....$ : chr [1:3] "(Intercept)" "Bwt" "SexM"
                : num 1.46
##
   $ sigma
## $ df
                 : int 141
## $ formula
                 :Class 'formula' length 3 Hwt ~ Bwt + Sex
   ....- attr(*, ".Environment")=<environment: R_GlobalEnv>
                 : language linMod(formula = Hwt ~ Bwt + Sex, data = cats)
  - attr(*, "class")= chr "linMod"
```

Lastly, we simply want to print the model to see that the output is now clear and easy to read. We shouldn't have to explicitly use print either. R will recognize the newly defined linMod class and then locate the linMod method that we have written for the generic print function.

```
# Print modelFit
modelFit

## Call:
## linMod(formula = Hwt ~ Bwt + Sex, data = cats)
##
```

The output is now much clearer and user-friendly.

Coefficients:

(Intercept) Bwt SexM ## -0.415 4.076 -0.082