

DATA 603 - Project

TEAM 5

3/15/2021

Life Expectancy in age

```
library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

library(ggplot2)
library(GGally)

## Warning: package 'GGally' was built under R version 4.0.4

## Registered S3 method overwritten by 'GGally':
##   method from
##   +.gg      ggplot2

library(leaps)
library(lmtest)

## Warning: package 'lmtest' was built under R version 4.0.4

## Loading required package: zoo

## Warning: package 'zoo' was built under R version 4.0.4

##
## Attaching package: 'zoo'

## The following objects are masked from 'package:base':
##
##   as.Date, as.Date.numeric

library(mctest) #for VIF
library(MASS)

##
## Attaching package: 'MASS'
```

```

## The following object is masked from 'package:dplyr':
##
##      select

library(car)

## Loading required package: carData

##
## Attaching package: 'car'

## The following object is masked from 'package:dplyr':
##
##      recode

library(olsrr)

## Warning: package 'olsrr' was built under R version 4.0.4

##
## Attaching package: 'olsrr'

## The following object is masked from 'package:MASS':
##
##      cement

## The following object is masked from 'package:datasets':
##
##      rivers

#life <- read.csv("Life Expectancy Data.csv", header = TRUE)
life <- read.csv("LifeExpectancyData.csv", header = TRUE)

#life

life2015 <- filter(life, life$Year == 2015)

#life2015

# Removing columns with considerable missing values
life2015$Country <- NULL
life2015$Year <- NULL
life2015$Alcohol <- NULL
life2015$percentage.expenditure <- NULL
life2015$Total.expenditure <- NULL
#select(life2015, Life.expectancy, everything())

#Removing rows with Missing Data
life2015 = na.omit(life2015)

#life2015

```

```

# Dummy Variables
#Status
contrasts(factor(life2015$Status))

##           Developing
## Developed           0
## Developing          1

##### MODEL A - ALL Variables
#####
#ggpairs(life2015, lower = list(continuous = "smooth_loess", combo =
"facethist", discrete = "facetbar",
#                               na = "na"))

life2015_full_modelA <- lm(Life.expectancy ~ factor(Status) + Adult.Mortality
+ infant.deaths + Hepatitis.B +
    Measles + BMI + under.five.deaths + Polio + Diphtheria
+ HIV.AIDS + GDP + Population +
    thinness..1.19.years + thinness.5.9.years +
Income.composition.of.resources +
    Schooling, data = life2015)

summary(life2015_full_modelA)

##
## Call:
## lm(formula = Life.expectancy ~ factor(Status) + Adult.Mortality +
##     infant.deaths + Hepatitis.B + Measles + BMI + under.five.deaths +
##     Polio + Diphtheria + HIV.AIDS + GDP + Population +
thinness..1.19.years +
##     thinness.5.9.years + Income.composition.of.resources + Schooling,
##     data = life2015)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -7.3416 -1.4379  0.0359  1.5459  7.9402
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    5.040e+01  2.511e+00  20.074 < 2e-16 ***
## factor(Status)Developing -3.424e-01  8.299e-01  -0.413  0.6807
## Adult.Mortality -2.092e-02  3.607e-03  -5.799 6.15e-08 ***
## infant.deaths    6.601e-02  3.285e-02   2.010  0.0469 *
## Hepatitis.B      4.333e-02  2.260e-02   1.917  0.0577 .
## Measles          -5.119e-05  5.729e-05  -0.893  0.3735
## BMI              -8.580e-03  1.550e-02  -0.554  0.5809
## under.five.deaths -4.783e-02  2.354e-02  -2.032  0.0445 *

```

```

## Polio                1.147e-02  1.267e-02   0.905   0.3676
## Diphtheria           -1.106e-02  2.630e-02  -0.420   0.6750
## HIV.AIDS             -4.847e-01  2.239e-01  -2.165   0.0325 *
## GDP                  5.064e-06  3.003e-05   0.169   0.8664
## Population           -1.010e-09  9.564e-09  -0.106   0.9161
## thinness..1.19.years -1.228e-01  2.338e-01  -0.525   0.6004
## thinness.5.9.years   -1.735e-02  2.287e-01  -0.076   0.9396
## Income.composition.of.resources 3.325e+01  4.981e+00   6.676  9.59e-10 ***
## Schooling            -4.796e-02  2.402e-01  -0.200   0.8421
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.695 on 113 degrees of freedom
## Multiple R-squared:  0.9005, Adjusted R-squared:  0.8865
## F-statistic: 63.95 on 16 and 113 DF,  p-value: < 2.2e-16

paste("Based on the preliminary t-test with all the variables, only
Adult.Mortality, Infant.deaths, under.five.defths, HIV.AIDS
andIncome.composition.of.resources presented p-values lower than 0.05 that
indicates they have significant impact in Life Expectancy at 5% level.")

## [1] "Based on the preliminary t-test with all the variables, only
Adult.Mortality, Infant.deaths, under.five.defths, HIV.AIDS
andIncome.composition.of.resources presented p-values lower than 0.05 that
indicates they have significant impact in Life Expectancy at 5% level."

# (A.1) F test to verify if at least one variable is related to Life
Expectancy
#The Analysis of Variance for Multiple Linear Regression
# Ho : Beta1 = Beta2 = ... BetaP = 0
# Ha : at least one BEta is NOT Zero

#Model with only one intercept
reg1 <- lm(Life.expectancy ~ 1, data = life2015)
summary(reg1)

##
## Call:
## lm(formula = Life.expectancy ~ 1, data = life2015)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -19.741  -5.016   1.409   5.284  14.258
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  70.7415      0.7014   100.9  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 7.997 on 129 degrees of freedom

```

#Comparing the NULL model with the full model

```
anova(reg1, life2015_full_modelA)
```

```
## Analysis of Variance Table
```

```
##
```

```
## Model 1: Life.expectancy ~ 1
```

```
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +  
infant.deaths +
```

```
## Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
```

```
## Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
```

```
## thinness.5.9.years + Income.composition.of.resources + Schooling
```

```
## Res.Df RSS Df Sum of Sq F Pr(>F)
```

```
## 1 129 8250.6
```

```
## 2 113 820.6 16 7430 63.948 < 2.2e-16 ***
```

```
## ---
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
paste("The output shows that Fcal = 63.95 with df 16, 113 (p-value < 2.2e-16  
< alpha = 0.05), indicating that we should clearly REJECT the null  
hypothesis. In other words, the large F-test suggests that at least one of  
the parameters must be related to the life expectancy. Based on the p-value,  
we also have extremely strong evidence that at least one of them is  
associated with the response variable.")
```

```
## [1] "The output shows that Fcal = 63.95 with df 16, 113 (p-value < 2.2e-16  
< alpha = 0.05), indicating that we should clearly REJECT the null  
hypothesis. In other words, the large F-test suggests that at least one of  
the parameters must be related to the life expectancy. Based on the p-value,  
we also have extremely strong evidence that at least one of them is  
associated with the response variable."
```

(A.2) PARTIAL TEST - Individual Coefficients Test (t-test)

Ho : Beta(i) = 0

Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

```
life2015_full_modelA <- lm(Life.expectancy ~ factor(Status) + Adult.Mortality  
+ infant.deaths + Hepatitis.B +  
Measles + BMI + under.five.deaths + Polio + Diphtheria  
+ HIV.AIDS + GDP + Population +  
thinness..1.19.years + thinness.5.9.years +  
Income.composition.of.resources +  
Schooling, data = life2015)
```

```
summary(life2015_full_modelA)
```

```
##
```

```
## Call:
```

```
## lm(formula = Life.expectancy ~ factor(Status) + Adult.Mortality +
```

```
## infant.deaths + Hepatitis.B + Measles + BMI + under.five.deaths +
```

```
## Polio + Diphtheria + HIV.AIDS + GDP + Population +
```

```
thinness..1.19.years +
```

```
## thinness.5.9.years + Income.composition.of.resources + Schooling,
```

```

##      data = life2015)
##
## Residuals:
##      Min        1Q    Median        3Q        Max
## -7.3416 -1.4379  0.0359  1.5459  7.9402
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    5.040e+01  2.511e+00  20.074 < 2e-16 ***
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## Measles         -5.119e-05  5.729e-05  -0.893  0.3735
## BMI             -8.580e-03  1.550e-02  -0.554  0.5809
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## Polio           1.147e-02  1.267e-02   0.905  0.3676
## Diphtheria      -1.106e-02  2.630e-02  -0.420  0.6750
## HIV.AIDS        -4.847e-01  2.239e-01  -2.165  0.0325 *
## GDP             5.064e-06  3.003e-05   0.169  0.8664
## Population      -1.010e-09  9.564e-09  -0.106  0.9161
## thinness..1.19.years -1.228e-01  2.338e-01  -0.525  0.6004
## thinness.5.9.years -1.735e-02  2.287e-01  -0.076  0.9396
## Income.composition.of.resources 3.325e+01  4.981e+00   6.676 9.59e-10 ***
## Schooling       -4.796e-02  2.402e-01  -0.200  0.8421
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.695 on 113 degrees of freedom
## Multiple R-squared:  0.9005, Adjusted R-squared:  0.8865
## F-statistic: 63.95 on 16 and 113 DF,  p-value: < 2.2e-16

paste("Based on the preliminary t-test with all the variables, only
Adult.Mortality, Infant.deaths, under.five.defths, HIV.AIDS
andIncome.composition.of.resources presented p-values lower than 0.05 that
indicates they have significant impact in Life Expectancy at 5%
level.Considering all the variables in the model, it was obtained a linear
model with 88.65% of variability in Life Expectancy represented by this full
model.")

## [1] "Based on the preliminary t-test with all the variables, only
Adult.Mortality, Infant.deaths, under.five.defths, HIV.AIDS
andIncome.composition.of.resources presented p-values lower than 0.05 that
indicates they have significant impact in Life Expectancy at 5%
level.Considering all the variables in the model, it was obtained a linear
model with 88.65% of variability in Life Expectancy represented by this full
model."

#confidence interval for the coefficients
confint(life2015_full_modelA, level = 0.95)

```

	2.5 %	97.5 %
## (Intercept)	4.542957e+01	5.537865e+01
## factor(Status)Developing	-1.986578e+00	1.301834e+00
## Adult.Mortality	-2.806309e-02	-1.377159e-02
## infant.deaths	9.328536e-04	1.310822e-01
## Hepatitis.B	-1.445415e-03	8.810163e-02
## Measles	-1.646942e-04	6.231932e-05
## BMI	-3.928192e-02	2.212157e-02
## under.five.deaths	-9.446317e-02	-1.193444e-03
## Polio	-1.364461e-02	3.657825e-02
## Diphtheria	-6.316618e-02	4.104998e-02
## HIV.AIDS	-9.282519e-01	-4.121939e-02
## GDP	-5.442526e-05	6.455425e-05
## Population	-1.995771e-08	1.793779e-08
## thinness..1.19.years	-5.861402e-01	3.404554e-01
## thinness.5.9.years	-4.704077e-01	4.357038e-01
## Income.composition.of.resources	2.338398e+01	4.312074e+01
## Schooling	-5.238588e-01	4.279341e-01

```
paste("For a 95% confidence interval, only the coefficients Adult.Mortality,
infant.deaths, under.five.deaths, HIV.AIDS and
Income.composition.of.resources do not contain Zero between upper and lower
values! Maybe, the other variables can be removed from the model.")
```

```
## [1] "For a 95% confidence interval, only the coefficients Adult.Mortality,
infant.deaths, under.five.deaths, HIV.AIDS and
Income.composition.of.resources do not contain Zero between upper and lower
values! Maybe, the other variables can be removed from the model."
```

(B.3) STEPWISE REGRESSION PROCEDURE

(B.3.1) BOTH

```
life2015_bothmodel = ols_step_both_p(life2015_full_modelA, pent = 0.05, prem
= 0.05, details = TRUE)
```

```
## Stepwise Selection Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1. factor(Status)
```

```
## 2. Adult.Mortality
```

```
## 3. infant.deaths
```

```
## 4. Hepatitis.B
```

```
## 5. Measles
```

```
## 6. BMI
```

```
## 7. under.five.deaths
```

```
## 8. Polio
```

```
## 9. Diphtheria
```

```
## 10. HIV.AIDS
```

```
## 11. GDP
```

```
## 12. Population
```

```

## 13. thinness..1.19.years
## 14. thinness.5.9.years
## 15. Income.composition.of.resources
## 16. Schooling
##
## We are selecting variables based on p value...
##
## Stepwise Selection: Step 1
##
## - Income.composition.of.resources added
##
##                               Model Summary
## -----
## R                               0.898          RMSE                3.532
## R-Squared                       0.807          Coef. Var          4.992
## Adj. R-Squared                  0.805          MSE                12.472
## Pred R-Squared                  0.801          MAE                2.737
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression          6654.186              1          6654.186          533.532          0.0000
## Residual            1596.410             128              12.472
## Total               8250.596             129
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               model          Beta          Std. Error          Std. Beta
## t          Sig          lower          upper
## -----
##                               (Intercept)          38.692          1.422
## 27.215          0.000          35.879          41.505
## Income.composition.of.resources          47.496          2.056          0.898
## 23.098          0.000          43.427          51.564
## -----
##
##
##
## Stepwise Selection: Step 2

```



```

##
## - Adult.Mortality added
##
##                               Model Summary
## -----
## R                               0.933          RMSE                2.910
## R-Squared                       0.870          Coef. Var          4.113
## Adj. R-Squared                  0.868          MSE                8.465
## Pred R-Squared                  0.862          MAE                2.137
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7175.479           2          3587.739      423.808      0.0000
## Residual        1075.117          127           8.465
## Total           8250.596          129
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               t      Sig      lower      model      Beta      Std. Error      Std. Beta
##                               upper
## -----
##                               (Intercept)      49.172          1.776
## 27.681      0.000      45.657      52.687
## ## Income.composition.of.resources      37.838          2.094          0.715
## 18.070      0.000      33.694      41.981
## ## Adult.Mortality      -0.025          0.003          -0.311      -
## 7.847      0.000      -0.031      -0.019
## -----
##
##
##
##                               Model Summary
## -----
## R                               0.933          RMSE                2.910
## R-Squared                       0.870          Coef. Var          4.113
## Adj. R-Squared                  0.868          MSE                8.465
## Pred R-Squared                  0.862          MAE                2.137
## -----
## RMSE: Root Mean Square Error

```

```

## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##
## ANOVA
## -----
##           Sum of          DF      Mean Square      F      Sig.
##           Squares
## -----
## Regression    7175.479         2      3587.739    423.808    0.0000
## Residual      1075.117        127         8.465
## Total         8250.596        129
## -----
##
## Parameter Estimates
## -----
##           Sig      lower      model      Beta      Std. Error      Std. Beta
##           t
## -----
##           (Intercept)      49.172      1.776
## 27.681    0.000      45.657      52.687
## Income.composition.of.resources      37.838      2.094      0.715
## 18.070    0.000      33.694      41.981
## Adult.Mortality      -0.025      0.003      -0.311      -
## 7.847    0.000      -0.031      -0.019
## -----
##
##
##
## Stepwise Selection: Step 3
##
## - Hepatitis.B added
##
## Model Summary
## -----
## R      0.942      RMSE      2.707
## R-Squared      0.888      Coef. Var      3.826
## Adj. R-Squared      0.885      MSE      7.326
## Pred R-Squared      0.877      MAE      2.093
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
##           Sum of          DF      Mean Square      F      Sig.
##           Squares

```

```

## -----
## Regression      7327.551      3      2442.517      333.415      0.0000
## Residual        923.045     126        7.326
## Total           8250.596     129
## -----
##
##                                     Parameter Estimates
## -----
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##                                     (Intercept)      47.150      1.711
27.555      0.000      43.763      50.536
## Income.composition.of.resources      35.544      2.012      0.672
17.667      0.000      31.563      39.526
## Adult.Mortality      -0.025      0.003      -0.317      -
8.604      0.000      -0.031      -0.020
## Hepatitis.B      0.045      0.010      0.142
4.556      0.000      0.026      0.065
## -----
## -----
##
##
##
##                                     Model Summary
## -----
## R      0.942      RMSE      2.707
## R-Squared      0.888      Coef. Var      3.826
## Adj. R-Squared      0.885      MSE      7.326
## Pred R-Squared      0.877      MAE      2.093
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of
##          Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7327.551      3      2442.517      333.415      0.0000
## Residual        923.045     126        7.326
## Total           8250.596     129
## -----
##
##                                     Parameter Estimates
## -----
## -----

```

```

##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##          (Intercept)      47.150      1.711
27.555      0.000      43.763      50.536
## Income.composition.of.resources      35.544      2.012      0.672
17.667      0.000      31.563      39.526
## Adult.Mortality      -0.025      0.003      -0.317      -
8.604      0.000      -0.031      -0.020
## Hepatitis.B      0.045      0.010      0.142
4.556      0.000      0.026      0.065
## -----
##
##
##
## Stepwise Selection: Step 4
##
## - HIV.AIDS added
##
##          Model Summary
## -----
## R      0.945      RMSE      2.654
## R-Squared      0.893      Coef. Var      3.752
## Adj. R-Squared      0.890      MSE      7.043
## Pred R-Squared      0.878      MAE      2.016
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##          ANOVA
## -----
##          Sum of      DF      Mean Square      F      Sig.
##          Squares
## -----
## Regression      7370.167      4      1842.542      261.597      0.0000
## Residual      880.429      125      7.043
## Total      8250.596      129
## -----
##
##          Parameter Estimates
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##          (Intercept)      47.782      1.697

```

```

28.150    0.000    44.423    51.141
## Income.composition.of.resources    35.112        1.981        0.664
17.728    0.000    31.192    39.032
##                               Adult.Mortality    -0.021        0.003        -0.263    -
6.199    0.000    -0.028    -0.014
##                               Hepatitis.B        0.038        0.010        0.117
3.662    0.000    0.017    0.058
##                               HIV.AIDS        -0.520        0.211        -0.099    -
2.460    0.015    -0.939    -0.102
## -----
##
##
##
##                               Model Summary
## -----
## R                                0.945        RMSE                2.654
## R-Squared                        0.893        Coef. Var              3.752
## Adj. R-Squared                   0.890        MSE                   7.043
## Pred R-Squared                   0.878        MAE                   2.016
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
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##
##                               ANOVA
## -----
##                               Sum of
##                               Squares        DF        Mean Square        F        Sig.
## -----
## Regression    7370.167            4            1842.542        261.597    0.0000
## Residual      880.429           125              7.043
## Total        8250.596           129
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model
##                               upper        Beta        Std. Error        Std. Beta
##                               t        Sig        lower
## -----
##                               (Intercept)    47.782        1.697
28.150    0.000    44.423    51.141
## Income.composition.of.resources    35.112        1.981        0.664
17.728    0.000    31.192    39.032
##                               Adult.Mortality    -0.021        0.003        -0.263    -
6.199    0.000    -0.028    -0.014
##                               Hepatitis.B        0.038        0.010        0.117
3.662    0.000    0.017    0.058

```

```

##              HIV.AIDS    -0.520          0.211        -0.099    -
2.460      0.015      -0.939      -0.102
## -----
##
##
##
## No more variables to be added/removed.
##
##
## Final Model Output
## -----
##
##              Model Summary
## -----
## R              0.945      RMSE              2.654
## R-Squared      0.893      Coef. Var          3.752
## Adj. R-Squared 0.890      MSE              7.043
## Pred R-Squared 0.878      MAE              2.016
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7370.167         4      1842.542      261.597      0.0000
## Residual        880.429        125         7.043
## Total          8250.596        129
## -----
##
##              Parameter Estimates
## -----
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)      47.782      1.697
## 28.150      0.000      44.423      51.141
## Income.composition.of.resources      35.112      1.981      0.664
## 17.728      0.000      31.192      39.032
## Adult.Mortality      -0.021      0.003      -0.263      -
## 6.199      0.000      -0.028      -0.014
## Hepatitis.B      0.038      0.010      0.117
## 3.662      0.000      0.017      0.058
## HIV.AIDS      -0.520      0.211      -0.099      -

```

```
2.460    0.015    -0.939    -0.102
```

```
## -----  
-----
```

```
life2015_bothmodel
```

```
##
```

```
## Stepwise Selection Summary
```

```
## -----
```

```
##  
## Step          Variable      Added/  
Square          C(p)          AIC      RMSE      Removed      R-Square      Adj.  
## -----  
-----
```

```
## 1      Income.composition.of.resources      addition      0.807
```

```
0.805      93.8370      700.9612      3.5316
```

```
## 2      Adult.Mortality      addition      0.870
```

```
0.868      24.0510      651.5686      2.9096
```

```
## 3      Hepatitis.B      addition      0.888
```

```
0.885      5.1100      633.7427      2.7066
```

```
## 4      HIV.AIDS      addition      0.893
```

```
0.890      1.2410      629.5978      2.6539
```

```
## -----  
-----
```

```
#life2015_bothmodel$mallows_cp
```

```
paste("Stepwise procedure with BOTH indicated to consider only the variables:  
Income.composition.of.resources, Adult.Mortality and HIV.AIDS.")
```

```
## [1] "Stepwise procedure with BOTH indicated to consider only the  
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS."
```

```
# (B.3.2) FORWARD
```

```
life2015_forwardmodel = ols_step_forward_p(life2015_full_modelA, pent = 0.05,  
details = TRUE)
```

```
## Forward Selection Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1. factor(Status)
```

```
## 2. Adult.Mortality
```

```
## 3. infant.deaths
```

```
## 4. Hepatitis.B
```

```
## 5. Measles
```

```
## 6. BMI
```

```
## 7. under.five.deaths
```

```
## 8. Polio
```

```
## 9. Diphtheria
```

```

## 10. HIV.AIDS
## 11. GDP
## 12. Population
## 13. thinness..1.19.years
## 14. thinness.5.9.years
## 15. Income.composition.of.resources
## 16. Schooling
##
## We are selecting variables based on p value...
##
##
## Forward Selection: Step 1
##
## - Income.composition.of.resources
##
##
##                               Model Summary
## -----
## R                               0.898          RMSE                3.532
## R-Squared                       0.807          Coef. Var          4.992
## Adj. R-Squared                   0.805          MSE                12.472
## Pred R-Squared                   0.801          MAE                2.737
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      6654.186           1      6654.186      533.532      0.0000
## Residual        1596.410          128      12.472
## Total           8250.596          129
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model      Beta      Std. Error      Std. Beta
## t      Sig      lower      upper
## -----
##                               (Intercept)      38.692      1.422
## 27.215      0.000      35.879      41.505
## Income.composition.of.resources      47.496      2.056      0.898
## 23.098      0.000      43.427      51.564
## -----
## -----
##

```



```

##
##
## Forward Selection: Step 2
##
## - Adult.Mortality
##
##                               Model Summary
## -----
## R                               0.933          RMSE          2.910
## R-Squared                     0.870          Coef. Var      4.113
## Adj. R-Squared                0.868          MSE           8.465
## Pred R-Squared                0.862          MAE           2.137
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      7175.479              2          3587.739          423.808      0.0000
## Residual        1075.117             127              8.465
## Total           8250.596             129
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               t          Sig          lower          model          Beta          Std. Error          Std. Beta
##                               upper
## -----
##                               (Intercept)          49.172          1.776
## 27.681      0.000          45.657          52.687
## ## Income.composition.of.resources          37.838          2.094          0.715
## 18.070      0.000          33.694          41.981
## ##                               Adult.Mortality          -0.025          0.003          -0.311      -
## 7.847      0.000          -0.031          -0.019
## -----
##
##
##
## Forward Selection: Step 3
##
## - Hepatitis.B
##
##                               Model Summary

```

```

## -----
## R                0.942      RMSE                2.707
## R-Squared        0.888      Coef. Var            3.826
## Adj. R-Squared   0.885      MSE                 7.326
## Pred R-Squared   0.877      MAE                 2.093
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##              Sum of          DF      Mean Square      F      Sig.
##              Squares
## -----
## Regression      7327.551         3      2442.517    333.415    0.0000
## Residual        923.045        126         7.326
## Total          8250.596        129
## -----
##
##                               Parameter Estimates
## -----
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t
## -----
##              (Intercept)      47.150      1.711
## 27.555      0.000      43.763      50.536
## Income.composition.of.resources      35.544      2.012      0.672
## 17.667      0.000      31.563      39.526
## Adult.Mortality      -0.025      0.003      -0.317      -
## 8.604      0.000      -0.031      -0.020
## Hepatitis.B      0.045      0.010      0.142
## 4.556      0.000      0.026      0.065
## -----
## -----
##
##
##
## Forward Selection: Step 4
##
## - HIV.AIDS
##
##                               Model Summary
## -----
## R                0.945      RMSE                2.654
## R-Squared        0.893      Coef. Var            3.752
## Adj. R-Squared   0.890      MSE                 7.043
## Pred R-Squared   0.878      MAE                 2.016

```

```

## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7370.167           4      1842.542      261.597      0.0000
## Residual        880.429          125           7.043
## Total          8250.596          129
## -----
##
##                               Parameter Estimates
## -----
##                               Sig.      lower      model      Beta      Std. Error      Std. Beta
##                               t      upper
## -----
##                               (Intercept)      47.782      1.697
## 28.150      0.000      44.423      51.141
## Income.composition.of.resources      35.112      1.981      0.664
## 17.728      0.000      31.192      39.032
## Adult.Mortality      -0.021      0.003      -0.263      -
## 6.199      0.000      -0.028      -0.014
## Hepatitis.B      0.038      0.010      0.117
## 3.662      0.000      0.017      0.058
## HIV.AIDS      -0.520      0.211      -0.099      -
## 2.460      0.015      -0.939      -0.102
## -----
##
##
##
## No more variables to be added.
##
## Variables Entered:
##
## + Income.composition.of.resources
## + Adult.Mortality
## + Hepatitis.B
## + HIV.AIDS
##
##
## Final Model Output
## -----
##

```

```

##                                     Model Summary
## -----
## R                                0.945      RMSE                2.654
## R-Squared                       0.893      Coef. Var          3.752
## Adj. R-Squared                   0.890      MSE                 7.043
## Pred R-Squared                   0.878      MAE                 2.016
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7370.167           4      1842.542      261.597      0.0000
## Residual        880.429          125           7.043
## Total          8250.596          129
## -----
##
##                                     Parameter Estimates
## -----
## -----
##                               model      Beta      Std. Error      Std. Beta
## t          Sig      lower      upper
## -----
##                               (Intercept)      47.782           1.697
## 28.150      0.000      44.423      51.141
## Income.composition.of.resources      35.112           1.981           0.664
## 17.728      0.000      31.192      39.032
## Adult.Mortality      -0.021           0.003      -0.263      -
## 6.199      0.000      -0.028      -0.014
## Hepatitis.B           0.038           0.010           0.117
## 3.662      0.000           0.017           0.058
## HIV.AIDS           -0.520           0.211      -0.099      -
## 2.460      0.015      -0.939      -0.102
## -----
## -----
life2015_forwardmodel

##
##                                     Selection Summary
## -----
## -----
##                               Variable
## Step      Entered      R-Square      Adj.      C(p)
## AIC      RMSE

```

```
## -----
##      1      Income.composition.of.resources      0.8065      0.8050      93.8368
700.9612      3.5316
##      2      Adult.Mortality      0.8697      0.8676      24.0511
651.5686      2.9096
##      3      Hepatitis.B      0.8881      0.8855      5.1097
633.7427      2.7066
##      4      HIV.AIDS      0.8933      0.8899      1.2413
629.5978      2.6539
## -----
```

```
#life2015_forwardmodel$mallows_cp
```

```
paste("Stepwise procedure with FORWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS.")
```

```
## [1] "Stepwise procedure with FORWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS."
```

```
# (B.3.3) BACKWARD
```

```
life2015_backwardmodel = ols_step_backward_p(life2015_full_modelA, prem =
0.05, details = TRUE)
```

```
## Backward Elimination Method
```

```
## -----
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1 . factor(Status)
```

```
## 2 . Adult.Mortality
```

```
## 3 . infant.deaths
```

```
## 4 . Hepatitis.B
```

```
## 5 . Measles
```

```
## 6 . BMI
```

```
## 7 . under.five.deaths
```

```
## 8 . Polio
```

```
## 9 . Diphtheria
```

```
## 10 . HIV.AIDS
```

```
## 11 . GDP
```

```
## 12 . Population
```

```
## 13 . thinness..1.19.years
```

```
## 14 . thinness.5.9.years
```

```
## 15 . Income.composition.of.resources
```

```
## 16 . Schooling
```

```
##
```

```
## We are eliminating variables based on p value...
```

```
##
```

```
## - thinness.5.9.years
```

```
##
```

```
## Backward Elimination: Step 1
```

```

##
## Variable thinness.5.9.years Removed
##
##                               Model Summary
## -----
## R                0.949          RMSE                2.683
## R-Squared        0.901          Coef. Var            3.793
## Adj. R-Squared   0.887          MSE                 7.198
## Pred R-Squared   0.856          MAE                 1.948
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                Sum of          DF      Mean Square      F      Sig.
##                Squares
## -----
## Regression      7429.971         15        495.331      68.811    0.0000
## Residual        820.625         114         7.198
## Total          8250.596         129
## -----
##
##                               Parameter Estimates
## -----
##
##                Sig      lower      model      Beta      Std. Error      Std. Beta
##                t                upper
## -----
##                (Intercept)      50.392      2.495
## 20.198      0.000      45.450      55.334
## factor(Status)Developing      -0.345      0.826      -0.015      -
## 0.417      0.677      -1.980      1.291
## Adult.Mortality      -0.021      0.004      -0.261      -
## 5.880      0.000      -0.028      -0.014
## infant.deaths      0.066      0.032      0.792
## 2.025      0.045      0.001      0.130
## Hepatitis.B      0.043      0.023      0.135
## 1.926      0.057      -0.001      0.088
## Measles      0.000      0.000      -0.058      -
## 0.895      0.373      0.000      0.000
## BMI      -0.008      0.015      -0.022      -
## 0.552      0.582      -0.038      0.022
## under.five.deaths      -0.048      0.023      -0.738      -
## 2.045      0.043      -0.094      -0.001
## Polio      0.011      0.013      0.036
## 0.905      0.367      -0.014      0.036
## Diphtheria      -0.011      0.026      -0.031      -

```

```

0.416    0.678    -0.062    0.041
##              HIV.AIDS    -0.482    0.219    -0.092    -
2.200    0.030    -0.915    -0.048
##              GDP    0.000    0.000    0.006
0.171    0.865    0.000    0.000
##              Population    0.000    0.000    -0.004    -
0.104    0.918    0.000    0.000
##              thinness..1.19.years    -0.139    0.084    -0.076    -
1.657    0.100    -0.306    0.027
## Income.composition.of.resources    33.245    4.958    0.629
6.705    0.000    23.423    43.068
##              Schooling    -0.049    0.239    -0.017    -
0.205    0.838    -0.522    0.424
## -----
##
##
## - Population
##
## Backward Elimination: Step 2
##
## Variable Population Removed
##
##
##              Model Summary
## -----
## R    0.949    RMSE    2.671
## R-Squared    0.901    Coef. Var    3.776
## Adj. R-Squared    0.888    MSE    7.137
## Pred R-Squared    0.862    MAE    1.950
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares    DF    Mean Square    F    Sig.
## -----
## Regression    7429.894    14    530.707    74.365    0.0000
## Residual    820.702    115    7.137
## Total    8250.596    129
## -----
##
##
##              Parameter Estimates
## -----
##
##              model
##              upper
##              lower
##              Sig
##              Beta    Std. Error    Std. Beta
## -----

```

```

-----
##              (Intercept)  50.392      2.484
20.285    0.000    45.472    55.313
##      factor(Status)Developing  -0.348      0.822    -0.015    -
0.423    0.673    -1.975    1.280
##      Adult.Mortality  -0.021      0.004    -0.261    -
5.905    0.000    -0.028    -0.014
##      infant.deaths    0.067      0.031      0.804
2.149    0.034      0.005      0.128
##      Hepatitis.B      0.043      0.022      0.135
1.932    0.056    -0.001      0.088
##      Measles          0.000      0.000    -0.058    -
0.896    0.372      0.000      0.000
##      BMI              -0.008      0.015    -0.022    -
0.549    0.584    -0.038      0.022
##      under.five.deaths  -0.048      0.022    -0.751    -
2.226    0.028    -0.092    -0.005
##      Polio            0.012      0.012      0.037
0.986    0.326    -0.012      0.035
##      Diphtheria       -0.011      0.026    -0.032    -
0.422    0.674    -0.062      0.040
##      HIV.AIDS         -0.480      0.218    -0.091    -
2.207    0.029    -0.911    -0.049
##      GDP              0.000      0.000      0.006
0.167    0.868      0.000      0.000
##      thinness..1.19.years  -0.138      0.083    -0.076    -
1.663    0.099    -0.303      0.026
## Income.composition.of.resources  33.220      4.931      0.628
6.737    0.000    23.452    42.987
##      Schooling       -0.050      0.238    -0.017    -
0.210    0.834    -0.521      0.421
## -----
##
##
## - GDP
##
## Backward Elimination: Step 3
##
## Variable GDP Removed
##
##              Model Summary
## -----
## R              0.949      RMSE              2.660
## R-Squared       0.901      Coef. Var          3.760
## Adj. R-Squared  0.889      MSE              7.077
## Pred R-Squared  0.864      MAE              1.951
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error

```


MAE: Mean Absolute Error

##

##

ANOVA

	Sum of				
	Squares	DF	Mean Square	F	Sig.
## Regression	7429.695	13	571.515	80.76	0.0000
## Residual	820.901	116	7.077		
## Total	8250.596	129			

##

##

Parameter Estimates

##

			model	Beta	Std. Error	Std. Beta	
t	Sig	lower	upper				
##			(Intercept)	50.327	2.442		
20.607	0.000	45.489	55.164				
##			factor(Status)Developing	-0.373	0.804	-0.017	-
0.464	0.644	-1.966	1.220				
##			Adult.Mortality	-0.021	0.004	-0.261	-
5.940	0.000	-0.028	-0.014				
##			infant.deaths	0.066	0.031	0.801	
2.153	0.033	0.005	0.128				
##			Hepatitis.B	0.043	0.022	0.134	
1.936	0.055	-0.001	0.086				
##			Measles	0.000	0.000	-0.058	-
0.905	0.367	0.000	0.000				
##			BMI	-0.008	0.015	-0.021	-
0.541	0.589	-0.038	0.022				
##			under.five.deaths	-0.048	0.022	-0.749	-
2.230	0.028	-0.091	-0.005				
##			Polio	0.012	0.012	0.037	
0.990	0.324	-0.012	0.035				
##			Diphtheria	-0.010	0.025	-0.030	-
0.407	0.685	-0.061	0.040				
##			HIV.AIDS	-0.478	0.216	-0.091	-
2.210	0.029	-0.906	-0.050				
##			thinness..1.19.years	-0.138	0.083	-0.075	-
1.664	0.099	-0.302	0.026				
##			Income.composition.of.resources	33.278	4.898	0.629	
6.795	0.000	23.578	42.979				
##			Schooling	-0.044	0.234	-0.015	-
0.190	0.850	-0.508	0.420				

##

##

```

##
## - Schooling
##
## Backward Elimination: Step 4
##
## Variable Schooling Removed
##
##                               Model Summary
## -----
## R                               0.949          RMSE          2.649
## R-Squared                     0.900          Coef. Var      3.745
## Adj. R-Squared                 0.890          MSE           7.018
## Pred R-Squared                 0.866          MAE           1.955
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      7429.441             12             619.120          88.214          0.0000
## Residual         821.155             117              7.018
## Total           8250.596             129
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model          Beta          Std. Error          Std. Beta
## t          Sig          lower          upper
## -----
##                               (Intercept)          50.283          2.422
## 20.765      0.000          45.488          55.079
## factor(Status)Developing          -0.352          0.793          -0.016          -
## 0.444      0.658          -1.922          1.219
## Adult.Mortality          -0.021          0.003          -0.263          -
## 6.122      0.000          -0.028          -0.014
## infant.deaths          0.067          0.031          0.806
## 2.178      0.031          0.006          0.128
## Hepatitis.B          0.042          0.022          0.132
## 1.937      0.055          -0.001          0.085
## Measles          0.000          0.000          -0.059          -
## 0.939      0.349          0.000          0.000
## BMI          -0.008          0.015          -0.022          -
## 0.567      0.572          -0.038          0.021
## under.five.deaths          -0.048          0.022          -0.751          -

```

```

2.249      0.026      -0.091      -0.006
##                               Polio          0.012          0.012          0.038
1.014      0.312      -0.011      0.035
##                               Diphtheria     -0.010          0.025         -0.029   -
0.395      0.693      -0.060      0.040
##                               HIV.AIDS       -0.478          0.215         -0.091   -
2.223      0.028      -0.905      -0.052
##               thinness..1.19.years        -0.138          0.082         -0.076   -
1.674      0.097      -0.301      0.025
## Income.composition.of.resources    32.529          2.880          0.615
11.294      0.000      26.825      38.233
## -----
##
##
## - Diphtheria
##
## Backward Elimination: Step 5
##
## Variable Diphtheria Removed
##
##                                Model Summary
## -----
## R                                0.949           RMSE                2.640
## R-Squared                       0.900           Coef. Var            3.732
## Adj. R-Squared                  0.891           MSE                 6.968
## Pred R-Squared                  0.869           MAE                 1.951
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                ANOVA
## -----
##              Sum of Squares             DF      Mean Square           F           Sig.
## -----
## Regression      7428.343                11         675.304        96.912      0.0000
## Residual         822.253                118          6.968
## Total           8250.596                129
## -----
##
##                                Parameter Estimates
## -----
##              model Beta      Std. Error      Std. Beta
## t      Sig      lower upper
## -----
##              (Intercept)    50.247          2.411

```

```

20.839    0.000    45.472    55.021
##      factor(Status)Developing    -0.324        0.787        -0.014    -
0.412    0.681    -1.883    1.234
##      Adult.Mortality    -0.021        0.003        -0.263    -
6.155    0.000    -0.028    -0.014
##      infant.deaths    0.067        0.031        0.812
2.206    0.029    0.007    0.128
##      Hepatitis.B    0.035        0.011        0.109
3.059    0.003    0.012    0.057
##      Measles    0.000        0.000        -0.061    -
0.965    0.336    0.000    0.000
##      BMI    -0.008        0.015        -0.020    -
0.523    0.602    -0.037    0.021
##      under.five.deaths    -0.049        0.021        -0.755    -
2.271    0.025    -0.091    -0.006
##      Polio    0.011        0.012        0.035
0.957    0.341    -0.012    0.034
##      HIV.AIDS    -0.471        0.214        -0.090    -
2.205    0.029    -0.894    -0.048
##      thinness..1.19.years    -0.142        0.082        -0.078    -
1.733    0.086    -0.303    0.020
## Income.composition.of.resources    32.285        2.803        0.610
11.517    0.000    26.734    37.836
## -----
##
##
## - factor(Status)
##
## Backward Elimination: Step 6
##
## Variable factor(Status) Removed
##
##                               Model Summary
## -----
## R                               0.949          RMSE                2.631
## R-Squared                       0.900          Coef. Var          3.718
## Adj. R-Squared                   0.892          MSE                6.920
## Pred R-Squared                   0.872          MAE                1.952
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF      Mean Square          F          Sig.
## -----
## Regression          7427.159              10          742.716          107.334          0.0000

```

```

## Residual      823.437      119      6.920
## Total        8250.596      129
## -----
##
##                                     Parameter Estimates
## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      49.701      2.008
24.749      0.000      45.724      53.677
##          Adult.Mortality      -0.021      0.003      -0.264      -
6.210      0.000      -0.028      -0.014
##          infant.deaths      0.067      0.030      0.808
2.203      0.030      0.007      0.127
##          Hepatitis.B      0.035      0.011      0.109
3.081      0.003      0.012      0.057
##          Measles      0.000      0.000      -0.062      -
0.986      0.326      0.000      0.000
##          BMI      -0.008      0.015      -0.021      -
0.541      0.589      -0.037      0.021
##          under.five.deaths      -0.048      0.021      -0.749      -
2.263      0.025      -0.091      -0.006
##          Polio      0.011      0.011      0.035
0.960      0.339      -0.012      0.034
##          HIV.AIDS      -0.463      0.212      -0.088      -
2.184      0.031      -0.883      -0.043
##          thinness..1.19.years      -0.144      0.081      -0.079      -
1.769      0.079      -0.305      0.017
## Income.composition.of.resources      32.705      2.602      0.618
12.570      0.000      27.554      37.857
## -----
##
##
## - BMI
##
## Backward Elimination: Step 7
##
## Variable BMI Removed
##
##                                     Model Summary
## -----
## R      0.949      RMSE      2.623
## R-Squared      0.900      Coef. Var      3.708
## Adj. R-Squared      0.892      MSE      6.879
## Pred R-Squared      0.873      MAE      1.950
## -----

```

##

ANOVA

##	Sum of				
##	Squares	DF	Mean Square	F	Sig.

## Regression	7425.132	9	825.015	119.935	0.0000
---------------	----------	---	---------	---------	--------

## Residual	825.464	120	6.879
-------------	---------	-----	-------

```
## Total      8250.596      129
```

##

```
##                                     Parameter Estimates
```

##	model	Beta	Std. Error	Std. Beta
----	-------	------	------------	-----------

t	Sig	lower	upper
---	-----	-------	-------

##	(Intercept)	49.696	2.002
----	-------------	--------	-------

24.821	0.000	45.732	53.661
--------	-------	--------	--------

##	Adult.Mortality	-0.021	0.003	-0.264	-
----	-----------------	--------	-------	--------	---

		Adjusted for currency	07/02/11	07/03/11	07/04/11
6.221	0.0000	-0.028	-0.014		

##	infant.deaths	0.066	0.030	0.794
----	---------------	-------	-------	-------

Year	Infant deaths	Infant deaths	Infant deaths	Infant deaths
2017	0.031	0.006	0.126	

##	Hepatitis B	0.035	0.011	0.108
----	-------------	-------	-------	-------

3.064	0.003	0.012	0.057
-------	-------	-------	-------

0.004	0.005	0.012	0.037				
##		Measles	0.000	0.000	-0.059	-	

0.947	0.346	0.000	0.000	0.000	0.000	0.000	0.000
-------	-------	-------	-------	-------	-------	-------	-------

0.547	0.548	0.000	0.000			
##		under five deaths	-0.018	0.021	-0.739	-

##	under-five.deaths	-0.048	0.021	-0.755
2 242	0.027	-0.090	-0.006	

2.242	0.027	-0.050	-0.000			
##			Polio	0.013	0.011	0.037

1	0.034	0.308	-0.011	0.034	0.012	0.011	0.037
---	-------	-------	--------	-------	-------	-------	-------

1.024	0.508	-0.011	0.054			
##		HTV	AIDS	0.165	0.311	0.088

##	HIV.AIDS	-0.405	0.211	-0.088	-
2 200 0 030 -0 883 -0 016					

2.200	0.030	-0.885	-0.040				
##	thickness	1	10 years	0.133	0.070	0.073	

##	thickness...1.19.years	-0.133	0.079	-0.073	-
1.604	0.003	0.380	0.033		

1.694	0.093	-0.289	0.023
## Income composition of resources	22 120	2 367	0 607

##	Income.composition.of.Resources	32.129	2.367	0.607
13	536 0 000 37 443 36 815			

```
13.576      0.000      27.443      36.815
##
```

.....

||||

##

##

```
###
###      Message

```

```
## - Measles
##
```

```
##
## Backward Elimination: Step 2
```

```
## Backward Elimination: Step 8
##
```

```
##
## Model 1: Mean 1 - D
##
```

```
## Variable Measles Removed
```

```

##
##                               Model Summary
## -----
## R                               0.948           RMSE           2.622
## R-Squared                       0.899           Coef. Var       3.706
## Adj. R-Squared                   0.893           MSE            6.873
## Pred R-Squared                   0.872           MAE            1.952
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7418.964           8           927.371      134.93      0.0000
## Residual        831.632          121           6.873
## Total          8250.596          129
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model      Beta      Std. Error      Std. Beta
## t      Sig      lower      upper
## -----
##                               (Intercept)      49.538           1.994
## 24.839      0.000      45.589      53.486
##                               Adult.Mortality      -0.021           0.003      -0.261      -
## 6.172      0.000      -0.028      -0.014
##                               infant.deaths           0.048           0.024           0.577
## 2.035      0.044           0.001           0.094
##                               Hepatitis.B           0.033           0.011           0.104
## 2.976      0.004           0.011           0.056
##                               under.five.deaths      -0.037           0.018      -0.576      -
## 2.051      0.042      -0.073      -0.001
##                               Polio           0.012           0.011           0.037
## 1.013      0.313      -0.011           0.034
##                               HIV.AIDS      -0.471           0.211      -0.090      -
## 2.232      0.027      -0.889      -0.053
##                               thinness..1.19.years      -0.113           0.076      -0.062      -
## 1.496      0.137      -0.264           0.037
## Income.composition.of.resources      32.371           2.352           0.612
## 13.765      0.000      27.715      37.027
## -----
## -----
##

```

```

##
## - Polio
##
## Backward Elimination: Step 9
##
## Variable Polio Removed
##
##
## Model Summary
## -----
## R                0.948          RMSE          2.622
## R-Squared         0.898          Coef. Var      3.706
## Adj. R-Squared    0.893          MSE           6.874
## Pred R-Squared    0.876          MAE           1.954
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of Squares      DF      Mean Square      F      Sig.
## -----
## Regression    7411.911      7      1058.844    154.026    0.0000
## Residual      838.685     122        6.874
## Total        8250.596     129
## -----
##
## Parameter Estimates
## -----
## t      Sig      lower      upper      Beta      Std. Error      Std. Beta
## -----
## (Intercept)  25.111    0.000    45.842    53.689    49.765      1.982
## Adult.Mortality  6.191    0.000   -0.028   -0.014   -0.021      0.003    -0.262    -
## infant.deaths  2.049    0.043    0.002    0.095    0.048      0.024    0.581
## Hepatitis.B    3.685    0.000    0.018    0.058    0.038      0.010    0.118
## under.five.deaths  2.071    0.040   -0.073   -0.002   -0.038      0.018   -0.581    -
## HIV.AIDS      2.326    0.022   -0.906   -0.073   -0.489      0.210   -0.093    -
## thinness..1.19.years  1.476    0.143   -0.262    0.038   -0.112      0.076   -0.061    -
## Income.composition.of.resources  32.909      2.291      0.622

```



```

14.362    0.000    28.373    37.445
## -----
##
##
## - thinness..1.19.years
##
## Backward Elimination: Step 10
##
## Variable thinness..1.19.years Removed
##
##                               Model Summary
## -----
## R                               0.947          RMSE                2.634
## R-Squared                       0.897          Coef. Var          3.724
## Adj. R-Squared                  0.891          MSE                6.940
## Pred R-Squared                  0.877          MAE                1.972
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression          7396.941              6          1232.824          177.633          0.0000
## Residual             853.654             123              6.940
## Total              8250.596             129
## -----
##
##                               Parameter Estimates
## -----
##                               t          Sig          lower          model          Beta          Std. Error          Std. Beta
##                               t          Sig          lower          upper
## -----
##                               (Intercept)          48.269          1.711
28.215    0.000          44.882          51.655
##                               Adult.Mortality          -0.021          0.003          -0.256          -
6.053    0.000          -0.027          -0.014
##                               infant.deaths          0.039          0.023          0.471
1.714    0.089          -0.006          0.084
##                               Hepatitis.B          0.036          0.010          0.111
3.488    0.001          0.015          0.056
##                               under.five.deaths          -0.032          0.018          -0.500          -
1.807    0.073          -0.068          0.003
##                               HIV.AIDS          -0.496          0.211          -0.094          -

```

```

2.348    0.020    -0.914    -0.078
## Income.composition.of.resources    34.610    1.990    0.654
17.394    0.000    30.671    38.548
## -----
##
##
## - infant.deaths
##
## Backward Elimination: Step 11
##
## Variable infant.deaths Removed
##
##                               Model Summary
## -----
## R                               0.946    RMSE                               2.655
## R-Squared                       0.894    Coef. Var                       3.753
## Adj. R-Squared                  0.890    MSE                               7.049
## Pred R-Squared                  0.876    MAE                               2.017
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares    DF    Mean Square    F    Sig.
## -----
## Regression    7376.562         5    1475.312    209.304    0.0000
## Residual      874.034        124         7.049
## Total        8250.596        129
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model
##                               upper    Beta    Std. Error    Std. Beta
## t    Sig    lower
## -----
##                               (Intercept)    48.034    1.719
27.951    0.000    44.633    51.436
##                               Adult.Mortality    -0.021    0.003    -0.259    -
6.094    0.000    -0.028    -0.014
##                               Hepatitis.B    0.037    0.010    0.116
3.605    0.000    0.017    0.057
##                               under.five.deaths    -0.002    0.002    -0.029    -
0.953    0.343    -0.006    0.002
##                               HIV.AIDS    -0.532    0.212    -0.101    -

```

```

2.511    0.013    -0.952    -0.113
## Income.composition.of.resources    34.850    2.000    0.659
17.423    0.000    30.891    38.809
## -----
##
##
## - under.five.deaths
##
## Backward Elimination: Step 12
##
## Variable under.five.deaths Removed
##
##                               Model Summary
## -----
## R                               0.945    RMSE                               2.654
## R-Squared                       0.893    Coef. Var                       3.752
## Adj. R-Squared                   0.890    MSE                               7.043
## Pred R-Squared                   0.878    MAE                               2.016
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares    DF    Mean Square    F    Sig.
## -----
## Regression    7370.167         4    1842.542    261.597    0.0000
## Residual      880.429        125         7.043
## Total        8250.596        129
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model
##                               upper    Beta    Std. Error    Std. Beta
## t    Sig    lower
## -----
##                               (Intercept)    47.782    1.697
## 28.150    0.000    44.423    51.141
##                               Adult.Mortality    -0.021    0.003    -0.263    -
## 6.199    0.000    -0.028    -0.014
##                               Hepatitis.B    0.038    0.010    0.117
## 3.662    0.000    0.017    0.058
##                               HIV.AIDS    -0.520    0.211    -0.099    -
## 2.460    0.015    -0.939    -0.102
## Income.composition.of.resources    35.112    1.981    0.664

```

17.728 0.000 31.192 39.032

##

##

##

No more variables satisfy the condition of p value = 0.05

##

##

Variables Removed:

##

- thinness.5.9.years

- Population

- GDP

- Schooling

- Diphtheria

- factor(Status)

- BMI

- Measles

- Polio

- thinness..1.19.years

- infant.deaths

- under.five.deaths

##

##

Final Model Output

##

Model Summary

## R	0.945	RMSE	2.654
------	-------	------	-------

## R-Squared	0.893	Coef. Var	3.752
--------------	-------	-----------	-------

## Adj. R-Squared	0.890	MSE	7.043
-------------------	-------	-----	-------

## Pred R-Squared	0.878	MAE	2.016
-------------------	-------	-----	-------

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

##	Sum of				
##	Squares	DF	Mean Square	F	Sig.

## Regression	7370.167	4	1842.542	261.597	0.0000
---------------	----------	---	----------	---------	--------

## Residual	880.429	125	7.043		
-------------	---------	-----	-------	--	--

## Total	8250.596	129			
----------	----------	-----	--	--	--

##

##

Parameter Estimates

```

## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      47.782      1.697
28.150      0.000      44.423      51.141
##
##          Adult.Mortality      -0.021      0.003      -0.263      -
6.199      0.000      -0.028      -0.014
##
##          Hepatitis.B      0.038      0.010      0.117
3.662      0.000      0.017      0.058
##
##          HIV.AIDS      -0.520      0.211      -0.099      -
2.460      0.015      -0.939      -0.102
## Income.composition.of.resources      35.112      1.981      0.664
17.728      0.000      31.192      39.032
## -----
## -----
life2015_backwardmodel
##
##
##
##          Elimination Summary
## -----
## -----
##          Variable      R-Square      Adj.      C(p)      AIC
## Step      Removed      R-Square
RMSE
## -----
## -----
##          1      thinness.5.9.years      0.9005      0.8875      15.0058
642.4531      2.6830
##          2      Population      0.9005      0.8884      13.0164
640.4653      2.6714
##          3      GDP      0.9005      0.8894      11.0438
638.4968      2.6602
##          4      Schooling      0.9005      0.8903      9.0788
636.5371      2.6492
##          5      Diphtheria      0.9003      0.891      7.2299
634.7107      2.6397
##          6      factor(Status)      0.9002      0.8918      5.3930
632.8979      2.6305
##          7      BMI      0.900      0.8924      3.6722
631.2175      2.6228
##          8      Measles      0.8992      0.8925      2.5215
630.1852      2.6216
##          9      Polio      0.8983      0.8925      1.4928
629.2831      2.6219
##          10      thinness..1.19.years      0.8965      0.8915      1.5542

```

```

629.5830    2.6344
## 11    infant.deaths            0.8941    0.8898    2.3606
630.6500    2.6549
## 12    under.five.deaths        0.8933    0.8899    1.2413
629.5978    2.6539
## -----
-----

#life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS.")

## [1] "Stepwise procedure with BACKWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS."

# (B.4) BEST SUBSET
#OPTION 1
ks = ols_step_best_subset(life2015_full_modelA, details = TRUE)

#Summary of Selected model based on cp, aic, AdjustedR2
rsquare <- c(ks$rsq)
cp <- c(ks$cp)
aic <- c(ks$aic)
AdjustedR2 <- c(ks$adjr)
cbind(rsquare, cp, aic, AdjustedR2)

##          rsquare          cp          aic AdjustedR2
## [1,] 0.8065098 93.836791 700.9612 0.8049981
## [2,] 0.8696922 24.051144 651.5686 0.8676401
## [3,] 0.8881239 5.109743 633.7427 0.8854601
## [4,] 0.8932890 1.241262 629.5978 0.8898743
## [5,] 0.8947742 1.553878 629.7758 0.8905312
## [6,] 0.8965342 1.554191 629.5830 0.8914871
## [7,] 0.8983486 1.492754 629.2831 0.8925161
## [8,] 0.8992035 2.521490 630.1852 0.8925392
## [9,] 0.8999510 3.672170 631.2175 0.8924473
## [10,] 0.9001967 5.393042 632.8979 0.8918098
## [11,] 0.9003402 7.229938 634.7107 0.8910499
## [12,] 0.9004732 9.078801 636.5371 0.8902654
## [13,] 0.9005041 11.043766 638.4968 0.8893537
## [14,] 0.9005282 13.016405 640.4653 0.8884185
## [15,] 0.9005375 15.005758 642.4531 0.8874504
## [16,] 0.9005426 17.000000 644.4465 0.8864601

#OPTION 2
best.subset <- regsubsets(Life.expectancy ~ factor(Status) + Adult.Mortality
+ infant.deaths + Hepatitis.B +
Measles + BMI + under.five.deaths + Polio + Diphtheria
+ HIV.AIDS + GDP + Population +
thinness..1.19.years + thinness.5.9.years +
Income.composition.of.resources +

```

```

Schooling, data = life2015, nv = 16)
summary(best.subset)

## Subset selection object
## Call: regsubsets.formula(Life.expectancy ~ factor(Status) +
Adult.Mortality +
## infant.deaths + Hepatitis.B + Measles + BMI + under.five.deaths +
## Polio + Diphtheria + HIV.AIDS + GDP + Population +
thinness..1.19.years +
## thinness.5.9.years + Income.composition.of.resources + Schooling,
## data = life2015, nv = 16)
## 16 Variables (and intercept)
##
##               Forced in Forced out
## factor(Status)Developing      FALSE      FALSE
## Adult.Mortality                FALSE      FALSE
## infant.deaths                  FALSE      FALSE
## Hepatitis.B                    FALSE      FALSE
## Measles                        FALSE      FALSE
## BMI                            FALSE      FALSE
## under.five.deaths              FALSE      FALSE
## Polio                          FALSE      FALSE
## Diphtheria                     FALSE      FALSE
## HIV.AIDS                       FALSE      FALSE
## GDP                            FALSE      FALSE
## Population                     FALSE      FALSE
## thinness..1.19.years           FALSE      FALSE
## thinness.5.9.years             FALSE      FALSE
## Income.composition.of.resources FALSE      FALSE
## Schooling                      FALSE      FALSE
## 1 subsets of each size up to 16
## Selection Algorithm: exhaustive
##               factor(Status)Developing Adult.Mortality infant.deaths
Hepatitis.B
## 1 ( 1 ) " " " " " "
## 2 ( 1 ) " " "*" " " "
## 3 ( 1 ) " " "*" " "*"
## 4 ( 1 ) " " "*" " "*"
## 5 ( 1 ) " " "*" " "*"
## 6 ( 1 ) " " "*" "*" "*"
## 7 ( 1 ) " " "*" "*" "*"
## 8 ( 1 ) " " "*" "*" "*"
## 9 ( 1 ) " " "*" "*" "*"
## 10 ( 1 ) " " "*" "*" "*"
## 11 ( 1 ) "*" "*" "*" "*"
## 12 ( 1 ) "*" "*" "*" "*"
## 13 ( 1 ) "*" "*" "*" "*"
## 14 ( 1 ) "*" "*" "*" "*"
## 15 ( 1 ) "*" "*" "*" "*"
## 16 ( 1 ) "*" "*" "*" "*"
## Measles BMI under.five.deaths Polio Diphtheria HIV.AIDS GDP

```

## 1	(1)	" "	" "	" "	" "	" "	" "	" "
## 2	(1)	" "	" "	" "	" "	" "	" "	" "
## 3	(1)	" "	" "	" "	" "	" "	" "	" "
## 4	(1)	" "	" "	" "	" "	" "	" "	" "
## 5	(1)	" "	" "	" "	" "	" "	" "	" "
## 6	(1)	" "	" "	" "	" "	" "	" "	" "
## 7	(1)	" "	" "	" "	" "	" "	" "	" "
## 8	(1)	" "	" "	" "	" "	" "	" "	" "
## 9	(1)	" "	" "	" "	" "	" "	" "	" "
## 10	(1)	" "	" "	" "	" "	" "	" "	" "
## 11	(1)	" "	" "	" "	" "	" "	" "	" "
## 12	(1)	" "	" "	" "	" "	" "	" "	" "
## 13	(1)	" "	" "	" "	" "	" "	" "	" "
## 14	(1)	" "	" "	" "	" "	" "	" "	" "
## 15	(1)	" "	" "	" "	" "	" "	" "	" "
## 16	(1)	" "	" "	" "	" "	" "	" "	" "

Population thinness..1.19.years thinness.5.9.years

## 1	(1)	" "	" "	" "
## 2	(1)	" "	" "	" "
## 3	(1)	" "	" "	" "
## 4	(1)	" "	" "	" "
## 5	(1)	" "	" "	" "
## 6	(1)	" "	" "	" "
## 7	(1)	" "	" "	" "
## 8	(1)	" "	" "	" "
## 9	(1)	" "	" "	" "
## 10	(1)	" "	" "	" "
## 11	(1)	" "	" "	" "
## 12	(1)	" "	" "	" "
## 13	(1)	" "	" "	" "
## 14	(1)	" "	" "	" "
## 15	(1)	" "	" "	" "
## 16	(1)	" "	" "	" "

Income.composition.of.resources Schooling

## 1	(1)	" "	" "
## 2	(1)	" "	" "
## 3	(1)	" "	" "
## 4	(1)	" "	" "
## 5	(1)	" "	" "
## 6	(1)	" "	" "
## 7	(1)	" "	" "
## 8	(1)	" "	" "
## 9	(1)	" "	" "
## 10	(1)	" "	" "
## 11	(1)	" "	" "
## 12	(1)	" "	" "
## 13	(1)	" "	" "
## 14	(1)	" "	" "
## 15	(1)	" "	" "
## 16	(1)	" "	" "


```
reg.summary <- summary(best.subset)
```

```
#Plotting
```

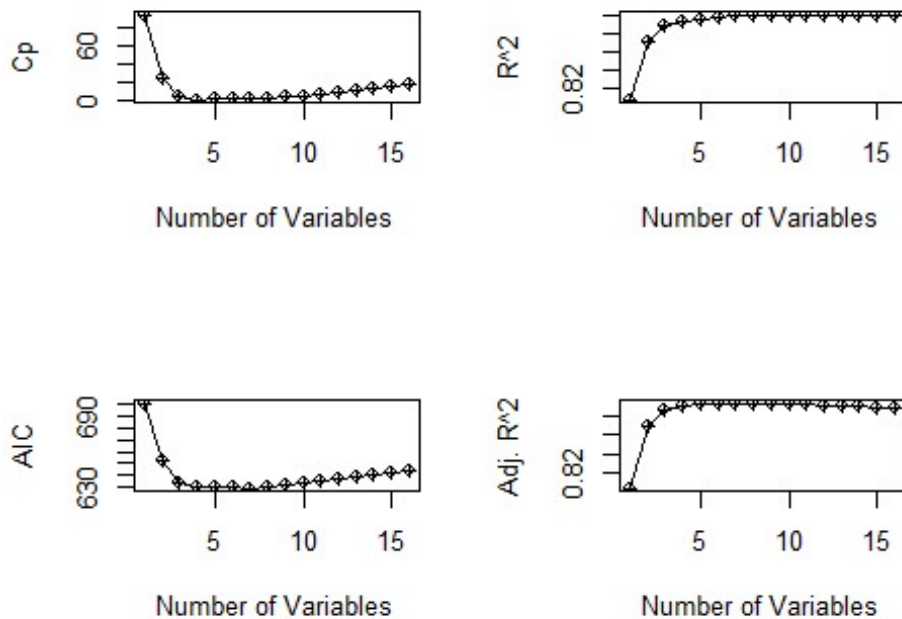
```
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
```

```
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
```

```
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab = "R^2")
```

```
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab = "AIC")
```

```
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Adj. R^2")
```



```
paste("Based on Marlow's Cp the minimum value was obtained with all the  
variables, but not significant reduction if the variables infant.deaths and  
under.five.deaths are considered.")
```

```
## [1] "Based on Marlow's Cp the minimum value was obtained with all the  
variables, but not significant reduction if the variables infant.deaths and  
under.five.deaths are considered."
```

```
#MODEL DIAGNOSTICS
```

```
# (A.3) CHECKING LINEARITY
```

```
#Residuals plot
```

```
plot(life2015_full_modelA, which = 1)
```

```
# a Scale Location Plot
```

```
plot(life2015_full_modelA, which = 3)
```

```
paste("It is verified some slightly change in the patter for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the patter for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected."
```

```
# (A.4) Heteroscedasticity Test - the Breush-Pagan test  
# Ho : heteroscedasticity is NOT presented (homoscedasticity)  
# Ha: heteroscedasticity is presented  
bptest(life2015_full_modelA)
```

```
##  
## studentized Breusch-Pagan test  
##  
## data: life2015_full_modelA  
## BP = 17.841, df = 16, p-value = 0.3333
```

```
paste("The output displays the Breush-Pagan test that from the model  
presented the p-value 0.333 > 0.05, indicating the we should NOT REJECT the  
null hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity does not exist.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model  
presented the p-value 0.333 > 0.05, indicating the we should NOT REJECT the  
null hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity does not exist."
```

```
# (A.5) Testing for normality  
#NORMALITY - Shapiro-Wilk Test  
# Ho : the sample data is significantly normally distributed  
# Ha : the sample data is NOT significantly normally distributed  
shapiro.test(residuals(life2015_full_modelA))
```

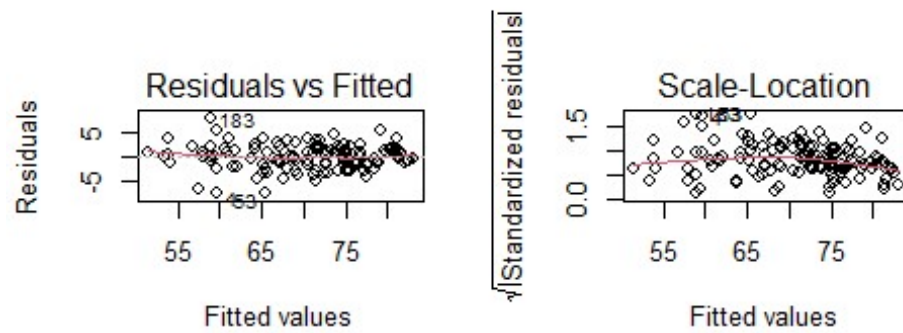
```
##  
## Shapiro-Wilk normality test  
##  
## data: residuals(life2015_full_modelA)  
## W = 0.98759, p-value = 0.2907
```

```
paste("Shapiro-Wilk normality test presented the p-value = 0.2907 > 0.05  
that indicated that we should NOT REJECT the null hypothesis and conclude  
that the residuals are normally distributed at 5 % level.")
```

```
## [1] "Shapiro-Wilk normality test presented the p-value = 0.2907 > 0.05  
that indicated that we should NOT REJECT the null hypothesis and conclude  
that the residuals are normally distributed at 5 % level."
```

```
#Plots
```

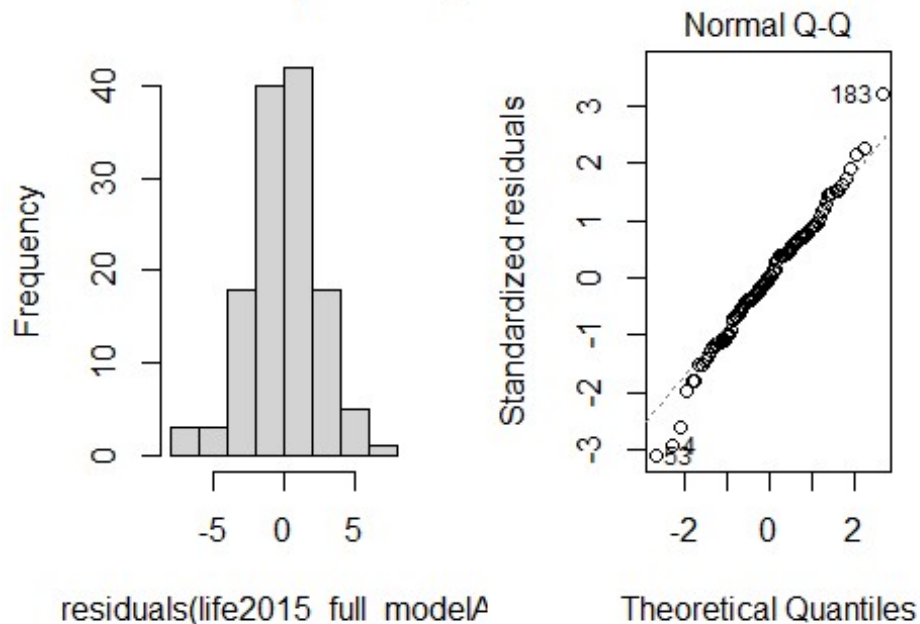
```
par(mfrow = c(1,2))
```



```
hist(residuals(life2015_full_modelA))
```

```
plot(life2015_full_modelA, which = 2)
```

am of residuals(life2015_fi



```
# (A.6) Multicollinearity test
#VIF
imcdiag(life2015_full_modelA, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_full_modelA, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
## VIF detection
## factor(Status)Developing 1.5387 0
## Adult.Mortality 2.2872 0
## infant.deaths 178.2035 1
## Hepatitis.B 5.6673 0
## Measles 4.8613 0
## BMI 1.9167 0
## under.five.deaths 151.2257 1
## Polio 1.8463 0
## Diphtheria 6.6654 0
## HIV.AIDS 2.0574 0
## GDP 1.6345 0
## Population 1.4796 0
## thinness..1.19.years 18.6420 1
## thinness.5.9.years 18.6210 1
## Income.composition.of.resources 10.0782 1
```

```

## Schooling                                7.8089                0
##
## Multicollinearity may be due to infant.deaths under.five.deaths
thinness..1.19.years thinness.5.9.years Income.composition.of.resources
regressors
##
## 1 --> COLLINEARITY is detected by the test
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("The preliminary Multicollinearity test indicated that some variables
with VIF higher than 10 that indicates severe multicollinearity problem. The
variables infant.deaths and under.five.deaths are redundant variables as well
as the thinness.1.19.years and thinness.5.9.years. ")

## [1] "The preliminary Multicollinearity test indicated that some variables
with VIF higher than 10 that indicates severe multicollinearity problem. The
variables infant.deaths and under.five.deaths are redundant variables as well
as the thinness.1.19.years and thinness.5.9.years. "

# (A.7) CHECKING FOR INFLUENTIAL OUTLIERS
#Plotting Cook's distance versus
plot(life2015_full_modelA, which = 5)

paste("It is possible to verify in the standardized residuals plot that none
case beyond the Cook's distance lines. However, points #119, #76 and #53 are
near these lines.")

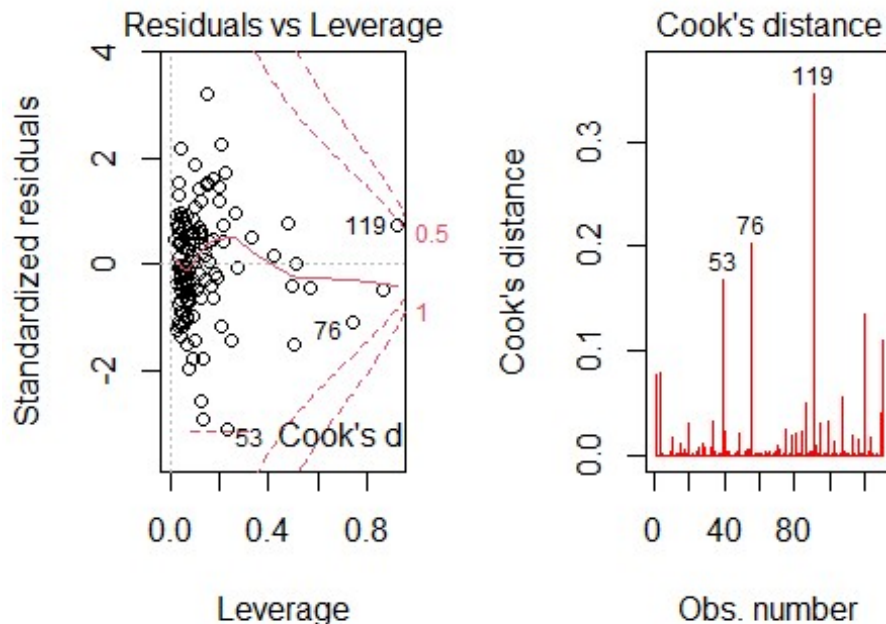
## [1] "It is possible to verify in the standardized residuals plot that none
case beyond the Cook's distance lines. However, points #119, #76 and #53 are
near these lines."

# Cook's Distance
#Checking if there is Cook statistics larger than 0.5
life2015[cooks.distance(life2015_full_modelA) > 0.5, ]

## [1] Status                                Life.expectancy
## [3] Adult.Mortality                        infant.deaths
## [5] Hepatitis.B                            Measles
## [7] BMI                                  under.five.deaths
## [9] Polio                                Diphtheria
## [11] HIV.AIDS                             GDP
## [13] Population                           thinness..1.19.years
## [15] thinness.5.9.years                   Income.composition.of.resources
## [17] Schooling
## <0 rows> (or 0-length row.names)

plot(life2015_full_modelA, pch = 18, col = "red", which = c(4))

```



```
paste("Based on the first model, we can remove some not significant variables
indicated bby t-test and re-valuate everthing considering some more
sophisticated model selection (Stepwise Procedure).")
```

```
## [1] "Based on the first model, we can remove some not significant
variables indicated bby t-test and re-valuate everthing considering some more
sophisticated model selection (Stepwise Procedure)."
```

```
##### MODEL B - First order model with only significant variables
indicated by t-test #####
```

```
paste("Based on the results above (t-test and 95% conf. interval for the
coefficients), we decided just to keep the significant variables:
Income.composition.of.resources, Adult.Mortality, infant.deaths,
under.five.deaths and HIV.AIDS. Some of them are redundant but we will be
removing later.")
```

```
## [1] "Based on the results above (t-test and 95% conf. interval for the
coefficients), we decided just to keep the significant variables:
Income.composition.of.resources, Adult.Mortality, infant.deaths,
under.five.deaths and HIV.AIDS. Some of them are redundant but we will be
removing later."
```

```
# (B.1) F test to verify if at least one variable is relataed to Life
Expectancy
```

```
#The Analysis of Variance for Multiple Linear Regression
```

```
# Ho : Reduced model
```

```
# Ha : Full model
```

#Reduced mode

```
life2015_reduced_modelB <- lm(Life.expectancy ~  
Income.composition.of.resources + Adult.Mortality + infant.deaths +  
under.five.deaths + HIV.AIDS, data = life2015)
```

#Comparing the NULL model with the full model

```
anova(life2015_reduced_modelB, life2015_full_modelA)
```

```
## Analysis of Variance Table
```

```
##
```

```
## Model 1: Life.expectancy ~ Income.composition.of.resources +  
Adult.Mortality +
```

```
## infant.deaths + under.five.deaths + HIV.AIDS
```

```
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +  
infant.deaths +
```

```
## Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
```

```
## Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
```

```
## thinness.5.9.years + Income.composition.of.resources + Schooling
```

```
## Res.Df RSS Df Sum of Sq F Pr(>F)
```

```
## 1 124 938.09
```

```
## 2 113 820.58 11 117.51 1.4711 0.1521
```

```
paste("The output shows that Fcal = 1.4711 with df 11, 113 (p-value = 0.1521  
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in  
other words, the reduced model presents better prediction of Life Expectancy  
compared with full.")
```

```
## [1] "The output shows that Fcal = 1.4711 with df 11, 113 (p-value = 0.1521  
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in  
other words, the reduced model presents better prediction of Life Expectancy  
compared with full."
```

(B.2) PARTIAL TEST - Individual Coefficients Test (t-test)

Ho : Beta(i) = 0

Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

```
life2015_reduced_modelB <- lm(Life.expectancy ~  
Income.composition.of.resources + Adult.Mortality + infant.deaths +  
under.five.deaths + HIV.AIDS, data = life2015)
```

```
summary(life2015_reduced_modelB)
```

```
##
```

```
## Call:
```

```
## lm(formula = Life.expectancy ~ Income.composition.of.resources +
```

```
## Adult.Mortality + infant.deaths + under.five.deaths + HIV.AIDS,
```

```
## data = life2015)
```

```
##
```

```
## Residuals:
```

```
## Min 1Q Median 3Q Max
```

```
## -9.1696 -1.4447 0.2028 1.7311 9.4861
##
## Coefficients:
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      50.054831   1.704210  29.371 < 2e-16 ***
## Income.composition.of.resources 35.952500   2.038205  17.639 < 2e-16 ***
## Adult.Mortality    -0.018214   0.003477  -5.239 6.72e-07 ***
## infant.deaths       0.045318   0.023739   1.909 0.058578 .
## under.five.deaths  -0.037446   0.018562  -2.017 0.045817 *
## HIV.AIDS           -0.717362   0.210455  -3.409 0.000881 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.751 on 124 degrees of freedom
## Multiple R-squared:  0.8863, Adjusted R-squared:  0.8817
## F-statistic: 193.3 on 5 and 124 DF,  p-value: < 2.2e-16

paste("For the reduced model, it is possible to verify a parameter
(infant.deaths) with p-values = 0.058576 > 0.05, that they might be removed
because this is not significant at 5 % level.")

## [1] "For the reduced model, it is possible to verify a parameter
(infant.deaths) with p-values = 0.058576 > 0.05, that they might be removed
because this is not significant at 5 % level."

#confidence interval for the coefficients
confint(life2015_reduced_modelB, level = 0.95)

##               2.5 %           97.5 %
## (Intercept)      46.681722306  53.4279398651
## Income.composition.of.resources 31.918321318  39.9866780097
## Adult.Mortality    -0.025095579  -0.0113321950
## infant.deaths       -0.001669247   0.0923047776
## under.five.deaths  -0.074185695  -0.0007068138
## HIV.AIDS           -1.133912071  -0.3008124332

paste("For a 95% confidence interval, only the coefficients infant.deaths
contains Zero between upper and lower values!")

## [1] "For a 95% confidence interval, only the coefficients infant.deaths
contains Zero between upper and lower values!"

# (B.3) STEPWISE REGRESSION PROCEDURE
# (B.3.1) BOTH
life2015_bothmodel = ols_step_both_p(life2015_reduced_modelB, pent = 0.05,
prem = 0.05, details = TRUE)

## Stepwise Selection Method
## -----
##
## Candidate Terms:
##
```



```

## 1. Income.composition.of.resources
## 2. Adult.Mortality
## 3. infant.deaths
## 4. under.five.deaths
## 5. HIV.AIDS
##
## We are selecting variables based on p value...
##
##
## Stepwise Selection: Step 1
##
## - Income.composition.of.resources added
##
##
##                               Model Summary
## -----
## R                               0.898          RMSE          3.532
## R-Squared                       0.807          Coef. Var      4.992
## Adj. R-Squared                  0.805          MSE           12.472
## Pred R-Squared                  0.801          MAE           2.737
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      6654.186              1          6654.186          533.532      0.0000
## Residual        1596.410             128           12.472
## Total           8250.596             129
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model          Beta          Std. Error          Std. Beta
## t          Sig          lower          upper
## -----
##                               (Intercept)          38.692          1.422
## 27.215      0.000          35.879          41.505
## Income.composition.of.resources          47.496          2.056          0.898
## 23.098      0.000          43.427          51.564
## -----
##
##
##
##

```

Stepwise Selection: Step 2

##

- Adult.Mortality added

##

Model Summary

##	-----
## R	0.933 RMSE 2.910
## R-Squared	0.870 Coef. Var 4.113
## Adj. R-Squared	0.868 MSE 8.465
## Pred R-Squared	0.862 MAE 2.137
##	-----

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

##	-----				
##	Sum of				
##	Squares	DF	Mean Square	F	Sig.
##	-----	-----	-----	-----	-----
## Regression	7175.479	2	3587.739	423.808	0.0000
## Residual	1075.117	127	8.465		
## Total	8250.596	129			
##	-----	-----	-----	-----	-----

##

Parameter Estimates

##	-----						
##							
##			model	Beta	Std. Error	Std. Beta	
t	Sig	lower	upper				
##	-----	-----	-----	-----	-----	-----	-----
##			(Intercept)	49.172	1.776		
27.681	0.000	45.657	52.687				
## Income.composition.of.resources				37.838	2.094	0.715	
18.070	0.000	33.694	41.981				
##			Adult.Mortality	-0.025	0.003	-0.311	-
7.847	0.000	-0.031	-0.019				
##	-----	-----	-----	-----	-----	-----	-----

##

##

##

Model Summary

##	-----
## R	0.933 RMSE 2.910
## R-Squared	0.870 Coef. Var 4.113
## Adj. R-Squared	0.868 MSE 8.465
## Pred R-Squared	0.862 MAE 2.137
##	-----

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

	Sum of Squares	DF	Mean Square	F	Sig.
Regression	7175.479	2	3587.739	423.808	0.0000
Residual	1075.117	127	8.465		
Total	8250.596	129			

##

##

Parameter Estimates

##

	Sig	lower	model upper	Beta	Std. Error	Std. Beta
--	-----	-------	----------------	------	------------	-----------

##

(Intercept)				49.172	1.776	
27.681	0.000	45.657	52.687			
Income.composition.of.resources				37.838	2.094	0.715
18.070	0.000	33.694	41.981			
Adult.Mortality				-0.025	0.003	-0.311
7.847	0.000	-0.031	-0.019			

##

##

##

##

##

Stepwise Selection: Step 3

##

- HIV.AIDS added

##

Model Summary

R	0.939	RMSE	2.782
R-Squared	0.882	Coef. Var	3.932
Adj. R-Squared	0.879	MSE	7.737
Pred R-Squared	0.867	MAE	2.090

##

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

##

Sum of

		Squares	DF	Mean Square	F	Sig.

Regression		7275.694	3	2425.231	313.446	0.0000
Residual		974.902	126	7.737		
Total		8250.596	129			

Parameter Estimates						

		model	Beta	Std. Error	Std. Beta	
t	Sig	lower upper				

		(Intercept)	49.589	1.702		
29.131	0.000	46.220 52.957				
		Income.composition.of.resources	36.633	2.030	0.693	
18.048	0.000	32.616 40.649				
		Adult.Mortality	-0.019	0.004	-0.233	-
5.343	0.000	-0.026 -0.012				
		HIV.AIDS	-0.759	0.211	-0.144	-
3.599	0.000	-1.176 -0.342				

Model Summary						

R		0.939	RMSE	2.782		
R-Squared		0.882	Coef. Var	3.932		
Adj. R-Squared		0.879	MSE	7.737		
Pred R-Squared		0.867	MAE	2.090		

RMSE: Root Mean Square Error						
MSE: Mean Square Error						
MAE: Mean Absolute Error						

ANOVA						

		Sum of	DF	Mean Square	F	Sig.
		Squares				

Regression		7275.694	3	2425.231	313.446	0.0000
Residual		974.902	126	7.737		
Total		8250.596	129			

Parameter Estimates						

```

-----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
t          Sig      lower      upper
## -----
##
##          (Intercept)      49.589      1.702
29.131      0.000      46.220      52.957
## Income.composition.of.resources      36.633      2.030      0.693
18.048      0.000      32.616      40.649
##          Adult.Mortality      -0.019      0.004      -0.233      -
5.343      0.000      -0.026      -0.012
##          HIV.AIDS      -0.759      0.211      -0.144      -
3.599      0.000      -1.176      -0.342
## -----
##
##
##
## No more variables to be added/removed.
##
##
## Final Model Output
## -----
##
##          Model Summary
## -----
## R          0.939      RMSE          2.782
## R-Squared      0.882      Coef. Var      3.932
## Adj. R-Squared      0.879      MSE          7.737
## Pred R-Squared      0.867      MAE          2.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##          ANOVA
## -----
##          Sum of      DF      Mean Square      F      Sig.
##          Squares
## -----
## Regression      7275.694      3      2425.231      313.446      0.0000
## Residual          974.902      126      7.737
## Total          8250.596      129
## -----
##
##          Parameter Estimates
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
t          Sig      lower      upper

```

```
## -----
##               (Intercept)    49.589      1.702
29.131    0.000    46.220    52.957
## Income.composition.of.resources    36.633      2.030      0.693
18.048    0.000    32.616    40.649
##               Adult.Mortality    -0.019      0.004      -0.233    -
5.343    0.000    -0.026    -0.012
##               HIV.AIDS    -0.759      0.211      -0.144    -
3.599    0.000    -1.176    -0.342
## -----
```

life2015_bothmodel

```
##
##                               Stepwise Selection Summary
## -----
##                               Added/      Adj.
## Step      Variable      Removed      R-Square      R-
## Square      C(p)      AIC      RMSE
## -----
##    1    Income.composition.of.resources    addition      0.807
0.805    85.0180    700.9612    3.5316
##    2      Adult.Mortality    addition      0.870
0.868    18.1120    651.5686    2.9096
##    3      HIV.AIDS    addition      0.882
0.879     6.8650    640.8483    2.7816
## -----
```

#life2015_bothmodel\$mallows_cp

paste("Stepwise procedure with BOTH indicated to consider only the variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS.")

[1] "Stepwise procedure with BOTH indicated to consider only the variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS."

(B.3.2) FORWARD

life2015_forwardmodel = **ols_step_forward_p**(life2015_reduced_modelB, **pent** = 0.05, **details** = **TRUE**)

Forward Selection Method

##

Candidate Terms:

##

1. Income.composition.of.resources

2. Adult.Mortality

```

## 3. infant.deaths
## 4. under.five.deaths
## 5. HIV.AIDS
##
## We are selecting variables based on p value...
##
## Forward Selection: Step 1
##
## - Income.composition.of.resources
##
##                               Model Summary
## -----
## R                               0.898           RMSE                3.532
## R-Squared                       0.807           Coef. Var          4.992
## Adj. R-Squared                   0.805           MSE                12.472
## Pred R-Squared                   0.801           MAE                2.737
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      6654.186           1      6654.186      533.532      0.0000
## Residual        1596.410          128           12.472
## Total           8250.596          129
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               model      Beta      Std. Error      Std. Beta
##                               t      Sig      lower      upper
## -----
##                               (Intercept)      38.692           1.422
## 27.215      0.000      35.879      41.505
## Income.composition.of.resources      47.496           2.056           0.898
## 23.098      0.000      43.427      51.564
## -----
##
##
##
## Forward Selection: Step 2
##

```

- Adult.Mortality

##

Model Summary

```
## -----
## R                0.933      RMSE                2.910
## R-Squared        0.870      Coef. Var            4.113
## Adj. R-Squared   0.868      MSE                 8.465
## Pred R-Squared   0.862      MAE                 2.137
## -----
```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

```
## -----
##              Sum of          DF      Mean Square      F      Sig.
##              Squares
## -----
## Regression    7175.479         2      3587.739    423.808    0.0000
## Residual      1075.117        127         8.465
## Total         8250.596        129
## -----
```

##

Parameter Estimates

```
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t
## -----
##              (Intercept)      49.172      1.776
## 27.681      0.000      45.657      52.687
## Income.composition.of.resources      37.838      2.094      0.715
## 18.070      0.000      33.694      41.981
## Adult.Mortality      -0.025      0.003      -0.311      -
## 7.847      0.000      -0.031      -0.019
## -----
```

##

##

##

Forward Selection: Step 3

##

- HIV.AIDS

##

Model Summary

```
## -----
## R                0.939      RMSE                2.782
## R-Squared        0.882      Coef. Var            3.932
## Adj. R-Squared   0.879      MSE                 7.737
```



```
## Pred R-Squared      0.867      MAE      2.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of
## Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7275.694      3      2425.231      313.446      0.0000
## Residual      974.902      126      7.737
## Total      8250.596      129
## -----
##
## Parameter Estimates
## -----
##
## Sig.      lower      upper      Beta      Std. Error      Std. Beta
## -----
## (Intercept)      49.589      1.702
29.131      0.000      46.220      52.957
## Income.composition.of.resources      36.633      2.030      0.693
18.048      0.000      32.616      40.649
## Adult.Mortality      -0.019      0.004      -0.233      -
5.343      0.000      -0.026      -0.012
## HIV.AIDS      -0.759      0.211      -0.144      -
3.599      0.000      -1.176      -0.342
## -----
##
##
## No more variables to be added.
##
## Variables Entered:
##
## + Income.composition.of.resources
## + Adult.Mortality
## + HIV.AIDS
##
## Final Model Output
## -----
##
## Model Summary
## -----
```

```

## R                0.939      RMSE                2.782
## R-Squared        0.882      Coef. Var            3.932
## Adj. R-Squared   0.879      MSE                 7.737
## Pred R-Squared   0.867      MAE                 2.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##              Sum of      DF      Mean Square      F      Sig.
##              Squares
## -----
## Regression    7275.694      3      2425.231    313.446    0.0000
## Residual      974.902     126        7.737
## Total        8250.596     129
## -----
##
##                               Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t
## -----
##              (Intercept)      49.589      1.702
## 29.131      0.000      46.220      52.957
## Income.composition.of.resources      36.633      2.030      0.693
## 18.048      0.000      32.616      40.649
## Adult.Mortality      -0.019      0.004      -0.233      -
## 5.343      0.000      -0.026      -0.012
## HIV.AIDS      -0.759      0.211      -0.144      -
## 3.599      0.000      -1.176      -0.342
## -----
## -----
life2015_forwardmodel
##
##                               Selection Summary
## -----
## -----
##              Variable      Adj.
##              Entered      R-Square      R-Square      C(p)
##              RMSE
## -----
## 1      Income.composition.of.resources      0.8065      0.8050      85.0180
## 700.9612      3.5316

```

```

##      2      Adult.Mortality          0.8697      0.8676      18.1121
651.5686      2.9096
##      3      HIV.AIDS              0.8818      0.8790      6.8653
640.8483      2.7816
## -----
-----

#life2015_forwardmodel$mallows_cp
paste("Stepwise procedure with FORWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS.")

## [1] "Stepwise procedure with FORWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS."

# (B.3.3) BACKWARD
life2015_backwardmodel = ols_step_backward_p(life2015_reduced_modelB, prem =
0.05, details = TRUE)

## Backward Elimination Method
## -----
##
## Candidate Terms:
##
## 1 . Income.composition.of.resources
## 2 . Adult.Mortality
## 3 . infant.deaths
## 4 . under.five.deaths
## 5 . HIV.AIDS
##
## We are eliminating variables based on p value...
##
## - infant.deaths
##
## Backward Elimination: Step 1
##
## Variable infant.deaths Removed
##
##
##                               Model Summary
## -----
## R              0.940      RMSE              2.779
## R-Squared      0.883      Coef. Var          3.929
## Adj. R-Squared 0.879      MSE              7.725
## Pred R-Squared 0.864      MAE              2.094
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of

```

```

##              Squares          DF    Mean Square          F          Sig.
## -----
## Regression    7284.932           4        1821.233    235.749    0.0000
## Residual      965.663          125          7.725
## Total        8250.596          129
## -----
##
##                               Parameter Estimates
## -----
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
## t              Sig      lower      upper
## -----
##              (Intercept)    49.861          1.719
29.004    0.000    46.459    53.264
## Income.composition.of.resources    36.293          2.052          0.686
17.689    0.000    32.233    40.354
##      Adult.Mortality    -0.018          0.004    -0.229    -
5.248    0.000    -0.025    -0.011
##      under.five.deaths    -0.002          0.002    -0.034    -
1.094    0.276    -0.006    0.002
##      HIV.AIDS    -0.769          0.211    -0.146    -
3.648    0.000    -1.187    -0.352
## -----
##
##
## - under.five.deaths
##
## Backward Elimination: Step 2
##
##   Variable under.five.deaths Removed
##
##                               Model Summary
## -----
## R              0.939          RMSE              2.782
## R-Squared      0.882          Coef. Var          3.932
## Adj. R-Squared 0.879          MSE              7.737
## Pred R-Squared 0.867          MAE              2.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##              Sum of
##              Squares          DF    Mean Square          F          Sig.
## -----

```

```

## Regression      7275.694          3      2425.231      313.446      0.0000
## Residual        974.902         126          7.737
## Total           8250.596         129
## -----
##
##                                     Parameter Estimates
## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      49.589          1.702
29.131      0.000      46.220      52.957
## Income.composition.of.resources      36.633          2.030          0.693
18.048      0.000      32.616      40.649
## Adult.Mortality      -0.019          0.004      -0.233      -
5.343      0.000      -0.026      -0.012
## HIV.AIDS      -0.759          0.211      -0.144      -
3.599      0.000      -1.176      -0.342
## -----
##
##
##
## No more variables satisfy the condition of p value = 0.05
##
##
## Variables Removed:
##
## - infant.deaths
## - under.five.deaths
##
## Final Model Output
## -----
##
##                                     Model Summary
## -----
## R          0.939          RMSE          2.782
## R-Squared    0.882          Coef. Var    3.932
## Adj. R-Squared    0.879          MSE          7.737
## Pred R-Squared    0.867          MAE          2.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----

```

```

##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    7275.694        3      2425.231    313.446    0.0000
## Residual      974.902       126        7.737
## Total        8250.596       129
## -----
##
##              Parameter Estimates
## -----
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)      49.589      1.702
29.131    0.000    46.220    52.957
## Income.composition.of.resources      36.633      2.030      0.693
18.048    0.000    32.616    40.649
##      Adult.Mortality      -0.019      0.004      -0.233    -
5.343    0.000    -0.026    -0.012
##      HIV.AIDS      -0.759      0.211      -0.144    -
3.599    0.000    -1.176    -0.342
## -----
## -----
life2015_backwardmodel

##
##
##              Elimination Summary
## -----
## -----
##              Variable      R-Square      Adj.      C(p)      AIC
##              Removed      R-Square
##              RMSE
## -----
##      1      infant.deaths      0.883      0.8792      7.6441      641.6105
2.7794
##      2      under.five.deaths      0.8818      0.879      6.8653      640.8483
2.7816
## -----
## -----
#Life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS.")

## [1] "Stepwise procedure with BACKWARD also indicated to consider only the
variables: Income.composition.of.resources, Adult.Mortality and HIV.AIDS."

```

```
# (B.4) BEST SUBSET
```

```
#OPTION 1
```

```
ks = ols_step_best_subset(life2015_reduced_modelB, details = TRUE)
```

```
#Summary of Selected model based on cp, aic, AdjustedR2
```

```
rsquare <- c(ks$rsq)
```

```
cp <- c(ks$cp)
```

```
aic <- c(ks$aic)
```

```
AdjustedR2 <- c(ks$adjr)
```

```
cbind(rsquare, cp, aic, AdjustedR2)
```

```
##           rsquare           cp           aic AdjustedR2
## [1,] 0.8065098 85.018009 700.9612 0.8049981
## [2,] 0.8696922 18.112052 651.5686 0.8676401
## [3,] 0.8818386 6.865340 640.8483 0.8790252
## [4,] 0.8829584 7.644146 641.6105 0.8792130
## [5,] 0.8862998 6.000000 639.8451 0.8817151
```

```
#OPTION 2
```

```
best.subset <- regsubsets(Life.expectancy ~ Income.composition.of.resources +  
Adult.Mortality + infant.deaths +  
under.five.deaths + HIV.AIDS, data = life2015, nv =  
5)
```

```
summary(best.subset)
```

```
## Subset selection object
```

```
## Call: regsubsets.formula(Life.expectancy ~ Income.composition.of.resources  
+
```

```
## Adult.Mortality + infant.deaths + under.five.deaths + HIV.AIDS,
```

```
## data = life2015, nv = 5)
```

```
## 5 Variables (and intercept)
```

```
##                               Forced in Forced out
## Income.composition.of.resources FALSE FALSE
## Adult.Mortality                 FALSE FALSE
## infant.deaths                   FALSE FALSE
## under.five.deaths               FALSE FALSE
## HIV.AIDS                        FALSE FALSE
```

```
## 1 subsets of each size up to 5
```

```
## Selection Algorithm: exhaustive
```

```
##           Income.composition.of.resources Adult.Mortality infant.deaths
## 1  ( 1 ) "*"                " "                " "
## 2  ( 1 ) "*"                " "                " "
## 3  ( 1 ) "*"                " "                " "
## 4  ( 1 ) "*"                " "                " "
## 5  ( 1 ) "*"                " "                " "
```

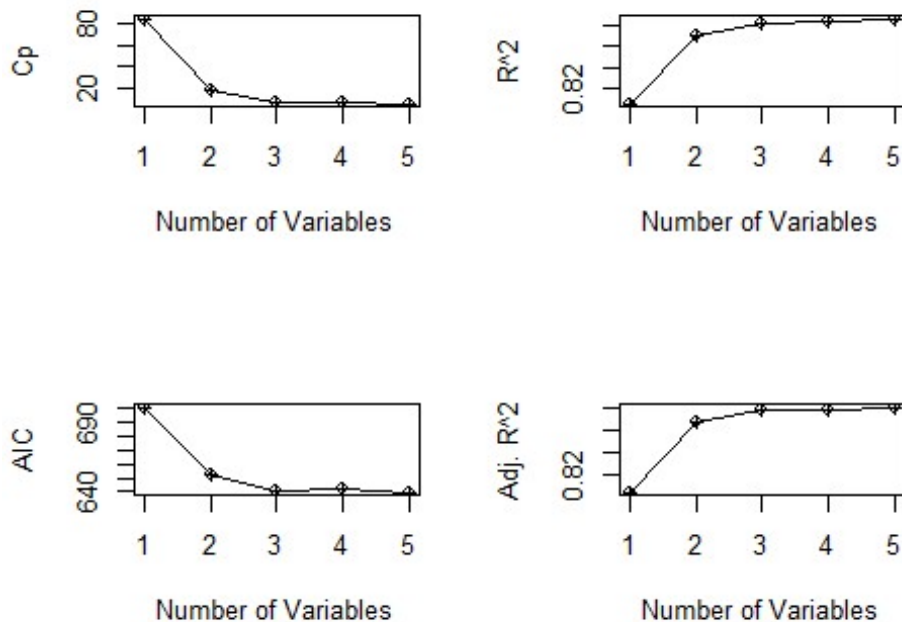
```
##           under.five.deaths HIV.AIDS
```

```
## 1  ( 1 ) " "                " "
## 2  ( 1 ) " "                " "
## 3  ( 1 ) " "                " "
```

```
## 4 ( 1 ) "*"      "*"
## 5 ( 1 ) "*"      "*"

reg.summary <- summary(best.subset)

#Plotting
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab = "R^2")
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab = "AIC")
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Adj. R^2")
```



```
paste("Based on Marlow's Cp the minimum value was obtained with all the
variables, but not significant reduction if the variables infant.deaths and
under.five.deaths are considered.")
```

```
## [1] "Based on Marlow's Cp the minimum value was obtained with all the
variables, but not significant reduction if the variables infant.deaths and
under.five.deaths are considered."
```

```
#MODEL DIAGNOSTICS
# (B.5) CHECKING LINEARITY
#Residuals plot
plot(life2015_reduced_modelB, which = 1)
```



```
# a Scale Location Plot
```

```
plot(life2015_reduced_modelB, which = 3)
```

```
paste("It is verified some slightly change in the patter for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the patter for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected."
```

```
# (B.6) Heteroscedasticity Test - the Breush-Pagan test
```

```
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
```

```
# Ha: heteroscedasticity is presented
```

```
bptest(life2015_reduced_modelB)
```

```
##
```

```
## studentized Breusch-Pagan test
```

```
##
```

```
## data: life2015_reduced_modelB
```

```
## BP = 13.986, df = 5, p-value = 0.0157
```

```
paste("The output displays the Breush-Pagan test that from the model  
presented the p-value 0.0157 < 0.05, indicating the we should REJECT the null  
hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity exists.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model  
presented the p-value 0.0157 < 0.05, indicating the we should REJECT the null  
hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity exists."
```

```
# (B.7) Testing for normality
```

```
#NORMALITY - Shapiro-Wilk Test
```

```
# Ho : the sample data is significantly normally distributed
```

```
# Ha : the sample data is NOT significantly normally distributed
```

```
shapiro.test(residuals(life2015_reduced_modelB))
```

```
##
```

```
## Shapiro-Wilk normality test
```

```
##
```

```
## data: residuals(life2015_reduced_modelB)
```

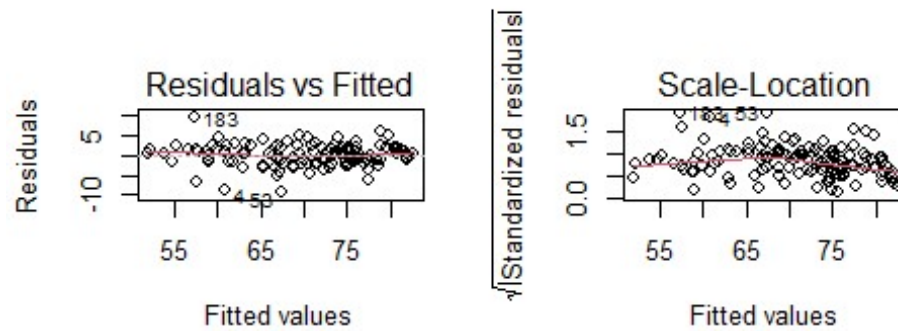
```
## W = 0.9737, p-value = 0.01248
```

```
paste("Shapiro-Wilk normality test presented the p-value = 0.01248 < 0.05  
that indicated that we should REJECT the null hypothesis and conclude that  
the residuals are NOT normally distributed at 5 % level.")
```

```
## [1] "Shapiro-Wilk normality test presented the p-value = 0.01248 < 0.05  
that indicated that we should REJECT the null hypothesis and conclude that  
the residuals are NOT normally distributed at 5 % level."
```

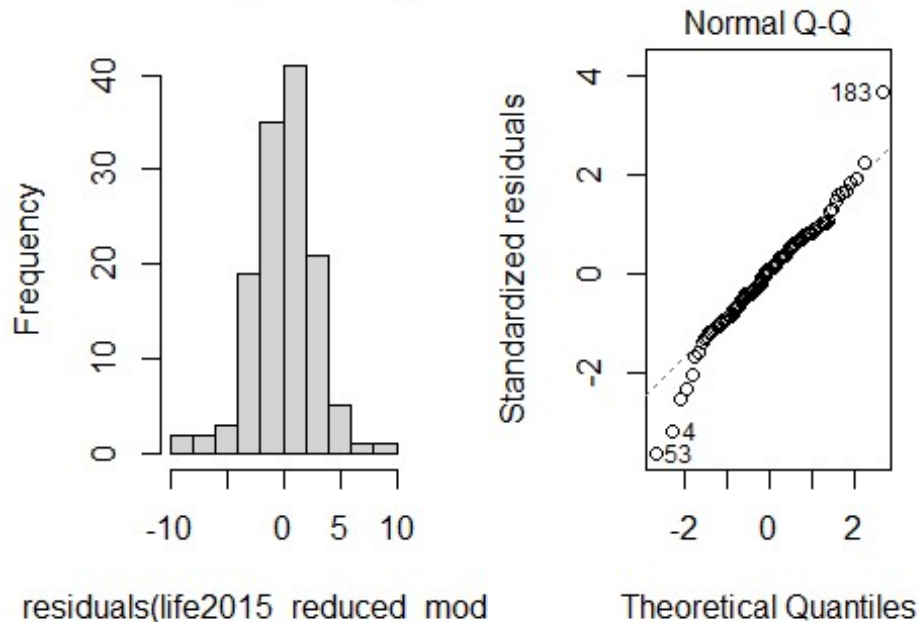
```
#Plots
```

```
par(mfrow = c(1,2))
```



```
hist(residuals(life2015_reduced_modelB))  
plot(life2015_reduced_modelB, which = 2)
```

of residuals(life2015_reduced_modelB)



```
paste("Q-Q plot indicates that data points #53 and #4 have significant
discrepancy what is expected.")
```

```
## [1] "Q-Q plot indicates that data points #53 and #4 have significant
discrepancy what is expected."
```

```
# (B.8) Multicollinearity test
```

```
#VIF
```

```
imcdiag(life2015_reduced_modelB, method = "VIF")
```

```
##
```

```
## Call:
```

```
## imcdiag(mod = life2015_reduced_modelB, method = "VIF")
```

```
##
```

```
##
```

```
## VIF Multicollinearity Diagnostics
```

```
##
```

```
##
```

	VIF	detection
Income.composition.of.resources	1.6198	0
Adult.Mortality	2.0401	0
infant.deaths	89.3513	1
under.five.deaths	90.2655	1
HIV.AIDS	1.7454	0

```
##
```

```
##
```

```
##
```

```
##
```

```
##
```

```
##
```

```
## Multicollinearity may be due to infant.deaths under.five.deaths regressors
```

```
##
```

```
## 1 --> COLLINEARITY is detected by the test
```

```
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("The preliminary Multicollinearity test indicated that infant.deaths and
under.five.deaths are redundant variables. ")

## [1] "The preliminary Multicollinearity test indicated that infant.deaths
and under.five.deaths are redundant variables. "

# (B.9) CHECKING FOR INFLUENTIAL OUTLIERS
#Plotting Cook's distance versus
plot(life2015_reduced_modelB, which = 5)

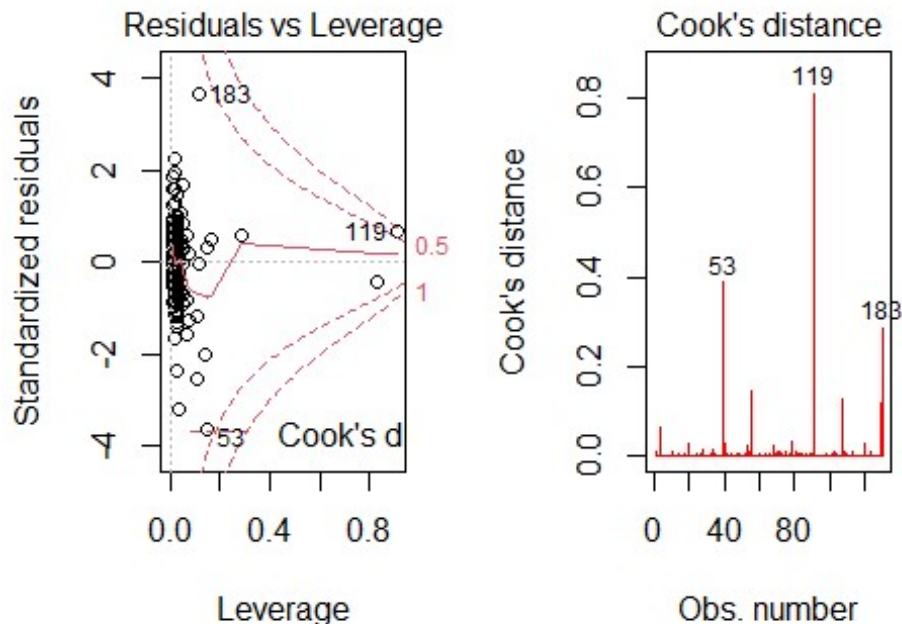
paste("It is possible to verify in the standardized residuals plot that the
data point #119 is beyond Cook's distance lines. However, points #183 and #53
are near these lines.")

## [1] "It is possible to verify in the standardized residuals plot that the
data point #119 is beyond Cook's distance lines. However, points #183 and #53
are near these lines."

# Cook's Distance
#Checking if there is Cook statistics Larger than 0.5
life2015[cooks.distance(life2015_reduced_modelB) > 0.5, ]

##           Status Life.expectancy Adult.Mortality infant.deaths Hepatitis.B
## 119 Developing              54.5              344             483             49
##      Measles  BMI under.five.deaths Polio Diphtheria HIV.AIDS      GDP
## 119   12423 25.4              747    49             49       3.7 2655.158
##      Population thinness..1.19.years thinness.5.9.years
## 119  181181744              9.8              9.7
##      Income.composition.of.resources Schooling
## 119              0.525              10

plot(life2015_reduced_modelB, pch = 18, col = "red", which = c(4))
```



```
paste("Based on the first model, we can remove some not significant variables
indicated bby t-test and re-valuate everthing considering some more
sophisticated model selection (Stepwise Procedure).")
```

```
## [1] "Based on the first model, we can remove some not significant
variables indicated bby t-test and re-valuate everthing considering some more
sophisticated model selection (Stepwise Procedure)."
```

```
##### MODEL C - Previous model without infant.deaths and without Data
point #119 #####
```

```
paste("Stepwise indicated to remove infant.deaths and under.five.deaths, but
as they are redundant might this have influenced in the tests, so we decided
to remove only one per time")
```

```
## [1] "Stepwise indicated to remove infant.deaths and under.five.deaths, but
as they are redundant might this have influenced in the tests, so we decided
to remove only one per time"
```

```
# (C.1) F test to verify if at least one variable is related to Life
Expectancy
```

```
#The Analysis of Variance for Multiple Linear Regression
```

```
# Ho : Reduced model B without infant.deaths
```

```
# Ha : Full model
```

```
#Reduced mode
```

```
life2015_reduced_modelC <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality +
```

```

                                under.five.deaths + HIV.AIDS, data = life2015)

#Comparing the NULL model with the full model
anova(life2015_reduced_modelC, life2015_full_modelA)

## Analysis of Variance Table
##
## Model 1: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
##      under.five.deaths + HIV.AIDS
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +
infant.deaths +
##      Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
##      Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
##      thinness.5.9.years + Income.composition.of.resources + Schooling
##   Res.Df    RSS Df Sum of Sq    F    Pr(>F)
## 1      125 965.66
## 2      113 820.58 12    145.08 1.6649 0.08401 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

paste("The output shows that Fcal = 1.6649 with df 12, 113 (p-value = 0.08401
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in
other words, the reduced model presents better prediction of Life Expectancy
compared with full model.")

## [1] "The output shows that Fcal = 1.6649 with df 12, 113 (p-value =
0.08401 > alpha = 0.05), indicating that we should NOT REJECT the null
hypothesis, in other words, the reduced model presents better prediction of
Life Expectancy compared with full model."

# (C.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_reduced_modelC <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality +
                                under.five.deaths + HIV.AIDS, data = life2015[-
c(119), ])

summary(life2015_reduced_modelC)

##
## Call:
## lm(formula = Life.expectancy ~ Income.composition.of.resources +
##      Adult.Mortality + under.five.deaths + HIV.AIDS, data = life2015[-
c(119),
##      ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max

```

```
## -8.9546 -1.7928 0.2002 1.7862 9.7707
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    49.860771    1.726071   28.887 < 2e-16 ***
## Income.composition.of.resources 36.292788    2.060015   17.618 < 2e-16 ***
## Adult.Mortality    -0.018431    0.003526   -5.228 7.06e-07 ***
## under.five.deaths    -0.002218    0.002039   -1.088 0.278674
## HIV.AIDS          -0.769029    0.211840   -3.630 0.000413 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.791 on 124 degrees of freedom
## Multiple R-squared:  0.8829, Adjusted R-squared:  0.8791
## F-statistic: 233.6 on 4 and 124 DF,  p-value: < 2.2e-16
```

```
paste("For the reduced model, it is possible to verify a parameter
(under.five.deaths) with p-values = 0.2786 > 0.05, that this variable should
be removed because this is not significant at 5 % level.")
```

```
## [1] "For the reduced model, it is possible to verify a parameter
(under.five.deaths) with p-values = 0.2786 > 0.05, that this variable should
be removed because this is not significant at 5 % level."
```

```
#confidence interval for the coefficients
confint(life2015_reduced_modelC, level = 0.95)
```

```
##              2.5 %      97.5 %
## (Intercept)    46.444393713 53.277148081
## Income.composition.of.resources 32.215441432 40.370134940
## Adult.Mortality    -0.025409931 -0.011452783
## under.five.deaths    -0.006253899 0.001817052
## HIV.AIDS          -1.188319764 -0.349738453
```

```
paste("For a 95% confidence interval, the coefficients under.five.deaths
contains ZERO between upper and lower values!")
```

```
## [1] "For a 95% confidence interval, the coefficients under.five.deaths
contains ZERO between upper and lower values!"
```

```
##### MODEL D - Model B without under.five.deaths and without Data
point #119 #####
```

```
paste("Stepwise indicated to remove infant.deaths and under.five.deaths, but
as they are redundant might this have influenced in the tests, so we decided
to remove only one per time")
```

```
## [1] "Stepwise indicated to remove infant.deaths and under.five.deaths, but
as they are redundant might this have influenced in the tests, so we decided
to remove only one per time"
```

```
# (D.1) F test to verify if at least one variable is related to Life
Expectancy
```

```

#The Analysis of Variance for Multiple Linear Regression
# Ho : Reduced model B without under.five.deaths
# Ha : Full model

#Reduced mode
life2015_reduced_modelD <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality +
                                infant.deaths + HIV.AIDS, data = life2015)

#Comparing the NULL model with the full model
anova(life2015_reduced_modelD, life2015_full_modelA)

## Analysis of Variance Table
##
## Model 1: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
##      infant.deaths + HIV.AIDS
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +
infant.deaths +
##      Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
##      Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
##      thinness.5.9.years + Income.composition.of.resources + Schooling
##   Res.Df    RSS Df Sum of Sq    F  Pr(>F)
## 1      125 968.88
## 2      113 820.58 12      148.3 1.7018 0.07539 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

paste("The output shows that Fcal = 1.6649 with df 12, 113 (p-value = 0.08401
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in
other words, the reduced model presents better prediction of Life Expectancy
compared with full model.")

## [1] "The output shows that Fcal = 1.6649 with df 12, 113 (p-value =
0.08401 > alpha = 0.05), indicating that we should NOT REJECT the null
hypothesis, in other words, the reduced model presents better prediction of
Life Expectancy compared with full model."

# (D.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_reduced_modelD <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality +
                                infant.deaths + HIV.AIDS, data = life2015[-c(119),
])

summary(life2015_reduced_modelD)

##
## Call:

```



```

## lm(formula = Life.expectancy ~ Income.composition.of.resources +
##     Adult.Mortality + infant.deaths + HIV.AIDS, data = life2015[-c(119),
##     ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -8.9375 -1.7939  0.1915  1.7940  9.7973
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    49.797057   1.727156  28.832 < 2e-16 ***
## Income.composition.of.resources 36.377013   2.060308  17.656 < 2e-16 ***
## Adult.Mortality    -0.018500   0.003531  -5.240 6.69e-07 ***
## infant.deaths     -0.002289   0.002612  -0.877 0.382451
## HIV.AIDS         -0.769541   0.212349  -3.624 0.000422 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.795 on 124 degrees of freedom
## Multiple R-squared:  0.8825, Adjusted R-squared:  0.8787
## F-statistic: 232.7 on 4 and 124 DF,  p-value: < 2.2e-16

paste("For the reduced model, it is possible to verify a parameter
(infant.deaths) with p-values = 0.3824 > 0.05, that this variable should be
removed because this is not significant at 5 % level.")

## [1] "For the reduced model, it is possible to verify a parameter
(infant.deaths) with p-values = 0.3824 > 0.05, that this variable should be
removed because this is not significant at 5 % level."

#confidence interval for the coefficients
confint(life2015_reduced_modelD, level = 0.95)

##              2.5 %       97.5 %
## (Intercept)    46.378531512  53.215583343
## Income.composition.of.resources 32.299087402  40.454939123
## Adult.Mortality    -0.025488178 -0.011511921
## infant.deaths     -0.007459032  0.002880345
## HIV.AIDS         -1.189838655 -0.349242385

paste("For a 95% confidence interval, the coefficients infant.deaths contains
ZERO between upper and lower values!")

## [1] "For a 95% confidence interval, the coefficients infant.deaths
contains ZERO between upper and lower values!"

##### MODEL E - Model B without under.five.deaths and infant.deaths
and without Data point #119 #####
paste("Stepwise indicated to remove infant.deaths and under.five.deaths.")

## [1] "Stepwise indicated to remove infant.deaths and under.five.deaths."

```

```

# (E.1) F test to verify if at least one variable is related to Life
Expectancy
#The Analysis of Variance for Multiple Linear Regression
# Ho : Reduced model B without under.five.deaths
# Ha : Full model

#Reduced mode
life2015_reduced_modelE <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality +
HIV.AIDS, data = life2015)

#Comparing the NULL model with the full model
anova(life2015_reduced_modelE, life2015_full_modelA)

## Analysis of Variance Table
##
## Model 1: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
## HIV.AIDS
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +
infant.deaths +
## Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
## Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
## thinness.5.9.years + Income.composition.of.resources + Schooling
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 126 974.90
## 2 113 820.58 13 154.32 1.6347 0.08575 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

paste("The output shows that Fcal = 1.6347 with df 13, 113 (p-value = 0.08575
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in
other words, the reduced model presents better prediction of Life Expectancy
compared with full model.")

## [1] "The output shows that Fcal = 1.6347 with df 13, 113 (p-value =
0.08575 > alpha = 0.05), indicating that we should NOT REJECT the null
hypothesis, in other words, the reduced model presents better prediction of
Life Expectancy compared with full model."

# (E.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_reduced_modelE <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality +
HIV.AIDS, data = life2015[-c(119), ])

summary(life2015_reduced_modelE)

```

```
##
## Call:
## lm(formula = Life.expectancy ~ Income.composition.of.resources +
##     Adult.Mortality + HIV.AIDS, data = life2015[-c(119), ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -8.9262 -1.7759  0.1187  1.8593  9.8325
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    49.588104    1.709038   29.015 < 2e-16 ***
## Income.composition.of.resources 36.631473    2.037860   17.975 < 2e-16 ***
## Adult.Mortality    -0.018723    0.003518   -5.322 4.59e-07 ***
## HIV.AIDS         -0.758344    0.211768   -3.581 0.000488 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.793 on 125 degrees of freedom
## Multiple R-squared:  0.8817, Adjusted R-squared:  0.8789
## F-statistic: 310.6 on 3 and 125 DF,  p-value: < 2.2e-16

paste("For the reduced model without infants.deaths and under.five.deaths the
t-test indicated all the coefficients with p-value lower than 0.05, and
consequently the remained variables are significant at 5 % level.")

## [1] "For the reduced model without infants.deaths and under.five.deaths
the t-test indicated all the coefficients with p-value lower than 0.05, and
consequently the remained variables are significant at 5 % level."

#confidence interval for the coefficients
confint(life2015_reduced_modelE, level = 0.95)

##              2.5 %      97.5 %
## (Intercept)    46.20570493  52.97050223
## Income.composition.of.resources 32.59829485  40.66465030
## Adult.Mortality    -0.02568627 -0.01176048
## HIV.AIDS         -1.17745844 -0.33922883

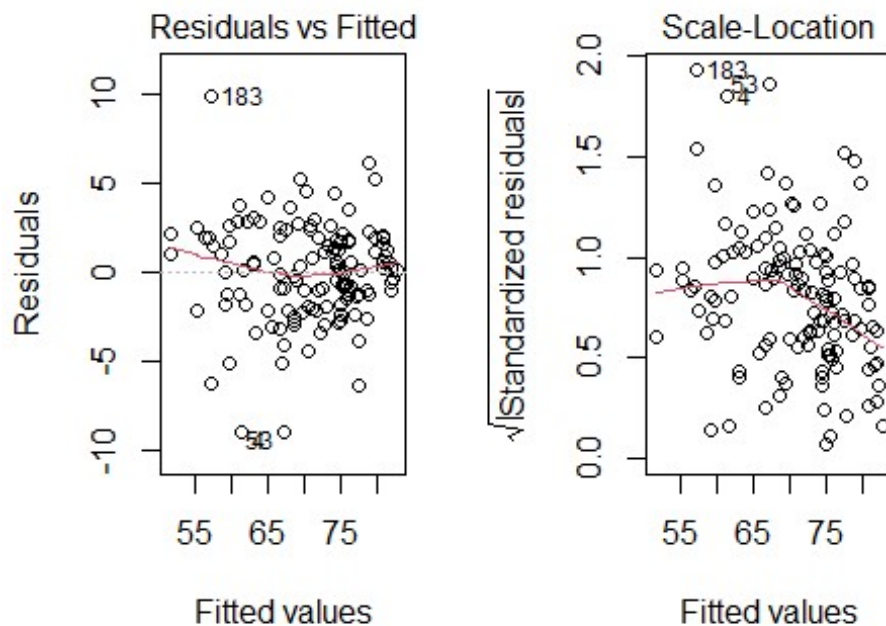
paste("For a 95% confidence interval, none coefficients with Zero between
upper and lower values!")

## [1] "For a 95% confidence interval, none coefficients with Zero between
upper and lower values!"

#MODEL DIAGNOSTICS
# (E.5) CHECKING LINEARITY
#Residuals plot
plot(life2015_reduced_modelE, which = 1)
```

```
# a Scale Location Plot
```

```
plot(life2015_reduced_modelE, which = 3)
```



```
paste("It is verified a pattern at the last plot that might indicates that  
the linearity assumption is not respected.")
```

```
## [1] "It is verified a pattern at the last plot that might indicates that  
the linearity assumption is not respected."
```

```
# (E.6) Heteroscedasticity Test - the Breush-Pagan test
```

```
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
```

```
# Ha: heteroscedasticity is presented
```

```
bptest(life2015_reduced_modelE)
```

```
##
```

```
## studentized Breusch-Pagan test
```

```
##
```

```
## data: life2015_reduced_modelE
```

```
## BP = 15.978, df = 3, p-value = 0.001146
```

```
paste("The output displays the Breush-Pagan test that from the model  
presented the p-value 0.0001146 < 0.05, indicating the we should REJECT the  
null hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity exists.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model  
presented the p-value 0.0001146 < 0.05, indicating the we should REJECT the
```

null hypothesis at 5 % level and consequently the test provides evidence that heteroscedasticity exists."

```
# (E.7) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_reduced_modelE))

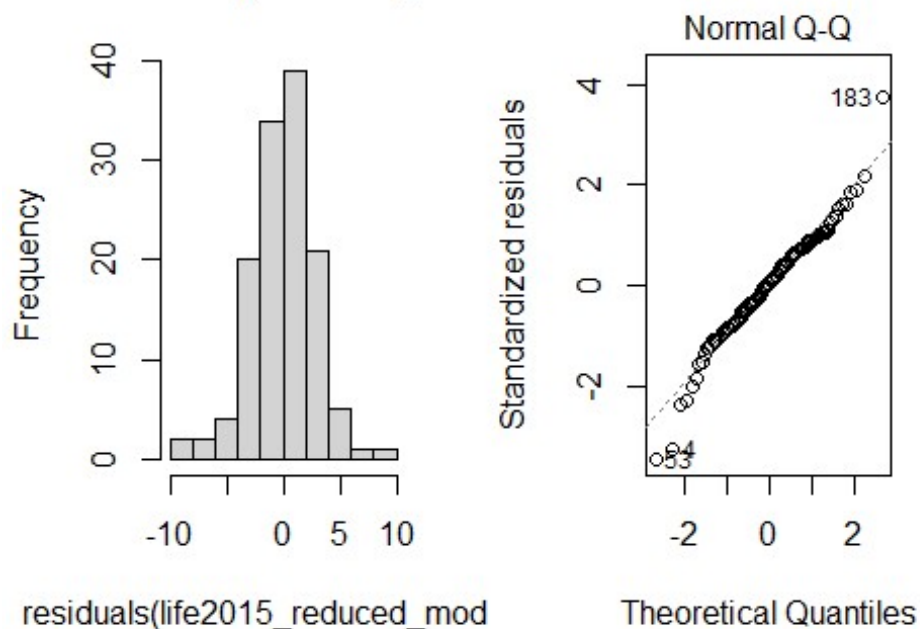
##
##  Shapiro-Wilk normality test
##
## data:  residuals(life2015_reduced_modelE)
## W = 0.97569, p-value = 0.02022

paste("Shapiro-Wilk normality test presented the p-value = 0.02022 < 0.05
that indicated that we should REJECT the null hypothesis and conclude that
the residuals are NOT normally distributed at 5 % level.")

## [1] "Shapiro-Wilk normality test presented the p-value = 0.02022 < 0.05
that indicated that we should REJECT the null hypothesis and conclude that
the residuals are NOT normally distributed at 5 % level."

#Plots
par(mfrow = c(1,2))
hist(residuals(life2015_reduced_modelE))
plot(life2015_reduced_modelE, which = 2)
```

l of residuals(life2015_redi



```

paste("Q-Q plot indicates that data points #53 and #4 have significant
discrepancy what is expected.")

## [1] "Q-Q plot indicates that data points #53 and #4 have significant
discrepancy what is expected."

# (E.8) Multicollinearity test
#VIF
imcdiag(life2015_reduced_modelE, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_reduced_modelE, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##                               VIF detection
## Income.composition.of.resources 1.5697      0
## Adult.Mortality                 2.0252      0
## HIV.AIDS                        1.7115      0
##
## NOTE: VIF Method Failed to detect multicollinearity
##
##
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("The preliminary Multicollinearity test indicated that none of the
selected variables are redudant (VIF < 5). ")

## [1] "The preliminary Multicollinearity test indicated that none of the
selected variables are redudant (VIF < 5). "

# (E.9) CHECKING FOR INFLUENTIAL OUTLIERS
#Plotting Cook's distance versus
plot(life2015_reduced_modelE, which = 5)

paste("It is possible to verify in the standardized residuals plot that the
data point #53 is beyond Cook's distance lines. However, points #183 are near
these lines.")

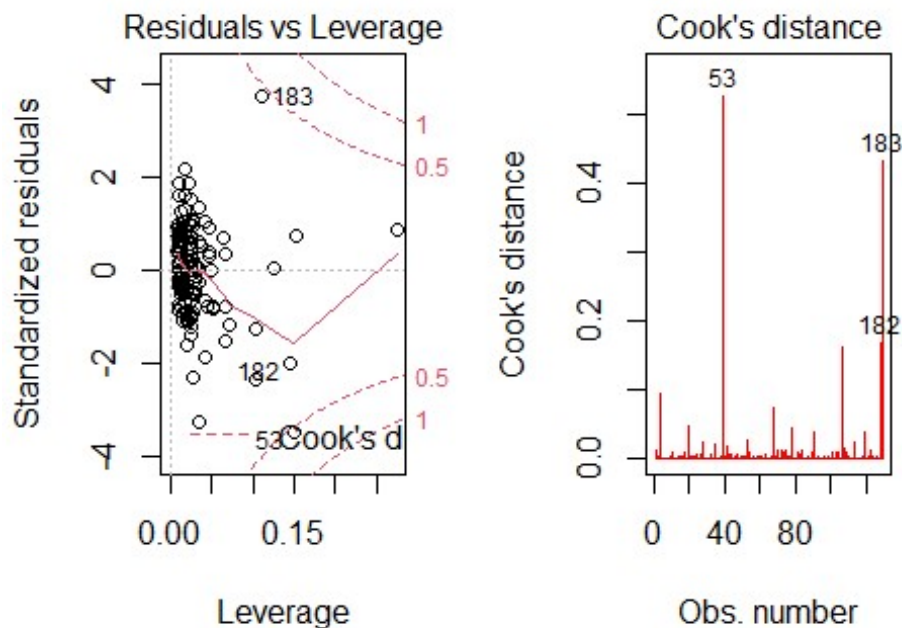
## [1] "It is possible to verify in the standardized residuals plot that the
data point #53 is beyond Cook's distance lines. However, points #183 are near
these lines."

# Coook's Distance
#Checking if there is Cook statistics Larger than 0.5
life2015[cooks.distance(life2015_reduced_modelE) > 0.5, ]

```

```
##      Status Life.expectancy Adult.Mortality infant.deaths Hepatitis.B
Measles
## 53 Developing                58.2                32                3                16
1250
##      BMI under.five.deaths Polio Diphtheria HIV.AIDS      GDP Population
## 53 24.5                    4      17          16      4.2 1347.313    1175389
##      thinness..1.19.years thinness.5.9.years Income.composition.of.resources
## 53                        8.4                8.3                        0.582
##      Schooling
## 53              9.2

plot(life2015_reduced_modelE, pch = 18, col = "red", which = c(4))
```



```
##### MODEL F - Model E with all interaction terms and without Data
points #119 and #53 #####
paste("Stepwise indicated to remove infant.deaths and under.five.deaths.")
## [1] "Stepwise indicated to remove infant.deaths and under.five.deaths."

# (F.1) F test to verify if at least one variable is related to Life
Expectancy
#The Analysis of Variance for Multiple Linear Regression
# Ho : Reduced model B without under.five.deaths
# Ha : Full model

#Reduced mode
life2015_reduced_modelF <- lm(Life.expectancy ~
```

```

(Income.composition.of.resources + Adult.Mortality +
  HIV.AIDS) ^ 2, data = life2015)

#Comparing the NULL model with the full model
anova(life2015_reduced_modelF, life2015_full_modelA)

## Analysis of Variance Table
##
## Model 1: Life.expectancy ~ (Income.composition.of.resources +
Adult.Mortality +
##   HIV.AIDS)^2
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +
infant.deaths +
##   Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
##   Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
##   thinness.5.9.years + Income.composition.of.resources + Schooling
## Res.Df    RSS Df Sum of Sq    F Pr(>F)
## 1      123 835.36
## 2      113 820.58 10      14.777 0.2035 0.9956

paste("The output shows that Fcal = 0.2035 with df 10, 113 (p-value = 0.9956
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in
other words, the reduced model presents better prediction of Life Expectancy
compared with full model.")

## [1] "The output shows that Fcal = 0.2035 with df 10, 113 (p-value = 0.9956
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in
other words, the reduced model presents better prediction of Life Expectancy
compared with full model."

# (F.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_reduced_modelF <- lm(Life.expectancy ~
(Income.composition.of.resources + Adult.Mortality +
  HIV.AIDS) ^ 2, data = life2015[-c(119, 53), ])

summary(life2015_reduced_modelF)

##
## Call:
## lm(formula = Life.expectancy ~ (Income.composition.of.resources +
##   Adult.Mortality + HIV.AIDS)^2, data = life2015[-c(119, 53),
##   ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -8.2749 -1.3370 -0.0992  1.5158 10.0969
##
## Coefficients:

```



```

##                                Estimate Std. Error t
value
## (Intercept)                   51.227651    2.356066
21.743
## Income.composition.of.resources 36.567617    3.148579
11.614
## Adult.Mortality                -0.014770    0.013332  -
1.108
## HIV.AIDS                      -1.679896    1.711485  -
0.982
## Income.composition.of.resources:Adult.Mortality -0.020165    0.021891  -
0.921
## Income.composition.of.resources:HIV.AIDS        -1.255603    2.728092  -
0.460
## Adult.Mortality:HIV.AIDS          0.004750    0.001408
3.374
##                                Pr(>|t|)
## (Intercept)                   < 2e-16 ***
## Income.composition.of.resources < 2e-16 ***
## Adult.Mortality                0.270096
## HIV.AIDS                      0.328283
## Income.composition.of.resources:Adult.Mortality 0.358792
## Income.composition.of.resources:HIV.AIDS        0.646163
## Adult.Mortality:HIV.AIDS        0.000997 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.605 on 121 degrees of freedom
## Multiple R-squared:  0.8998, Adjusted R-squared:  0.8948
## F-statistic: 181 on 6 and 121 DF,  p-value: < 2.2e-16

paste("For the interaction model the t-test indicated only the interaction
term Adult.Mortality * HIV.AIDS with p-value < 0.05 that indicates that only
this interaction term has significant influence in Life Expectancy at 5 %
level.")

## [1] "For the interaction model the t-test indicated only the interaction
term Adult.Mortality * HIV.AIDS with p-value < 0.05 that indicates that only
this interaction term has significant influence in Life Expectancy at 5 %
level."

#confidence interval for the coefficients
confint(life2015_reduced_modelF, level = 0.95)

##                                2.5 %      97.5 %
## (Intercept)                   46.563195609 55.892105765
## Income.composition.of.resources 30.334174183 42.801060260
## Adult.Mortality                -0.041164108  0.011623260
## HIV.AIDS                      -5.068231925  1.708439258
## Income.composition.of.resources:Adult.Mortality -0.063503967  0.023173393

```

```
## Income.composition.of.resources:HIV.AIDS      -6.656579993   4.145374454
## Adult.Mortality:HIV.AIDS                      0.001962741   0.007537992
```

```
paste("For a 95% confidence interval, only the interaction term
Adult.Mortality * HIV.AIDS without Zero between upper and lower values!")
```

```
## [1] "For a 95% confidence interval, only the interaction term
Adult.Mortality * HIV.AIDS without Zero between upper and lower values!"
```

```
# (F.3) STEPWISE REGRESSION PROCEDURE
```

```
# (F.3.1) BOTH
```

```
life2015_bothmodel = ols_step_both_p(life2015_reduced_modelF, pent = 0.05,
prem = 0.05, details = TRUE)
```

```
## Stepwise Selection Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1. Income.composition.of.resources
```

```
## 2. Adult.Mortality
```

```
## 3. HIV.AIDS
```

```
## 4. Income.composition.of.resources:Adult.Mortality
```

```
## 5. Income.composition.of.resources:HIV.AIDS
```

```
## 6. Adult.Mortality:HIV.AIDS
```

```
##
```

```
## We are selecting variables based on p value...
```

```
##
```

```
##
```

```
## Stepwise Selection: Step 1
```

```
##
```

```
## - Income.composition.of.resources added
```

```
##
```

```
## Model Summary
```

```
## -----
## R              0.897      RMSE              3.556
## R-Squared      0.805      Coef. Var        5.025
## Adj. R-Squared 0.804      MSE              12.647
## Pred R-Squared 0.799      MAE              2.765
## -----
```

```
## RMSE: Root Mean Square Error
```

```
## MSE: Mean Square Error
```

```
## MAE: Mean Absolute Error
```

```
##
```

```
## ANOVA
```

```
## -----
## Sum of
## Squares      DF      Mean Square      F      Sig.
## -----
## Regression    6596.898      1      6596.898    521.627    0.0000
## Residual      1593.492     126      12.647
```

```

## Total          8190.390          127
## -----
##
##                                     Parameter Estimates
## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      38.615          1.443
26.763      0.000      35.760      41.470
## Income.composition.of.resources      47.584          2.083          0.897
22.839      0.000      43.461      51.707
## -----
##
##
##
## Stepwise Selection: Step 2
##
## - Income.composition.of.resources:Adult.Mortality added
##
##                                     Model Summary
## -----
## R          0.933          RMSE          2.907
## R-Squared    0.871          Coef. Var    4.107
## Adj. R-Squared    0.869          MSE          8.450
## Pred R-Squared    0.863          MAE          2.104
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##
##          Sum of          DF      Mean Square          F          Sig.
##          Squares
## -----
## Regression    7134.199          2          3567.099    422.166    0.0000
## Residual      1056.191          125           8.450
## Total         8190.390          127
## -----
##
##
##                                     Parameter Estimates
## -----
##
##          Std. Beta      t          Sig      lower      model      Beta      Std. Error
##          t          Sig      lower      upper
## -----

```

```

-----
##                               (Intercept)    45.919    1.493
30.751    0.000    42.963    48.874
##                               Income.composition.of.resources    43.358    1.783
0.818    24.311    0.000    39.828    46.888
## Income.composition.of.resources:Adult.Mortality    -0.045    0.006
-0.268    -7.974    0.000    -0.056    -0.034
## -----

```

```

##
##
##

```

```

##                               Model Summary
## -----
## R                               0.933    RMSE                               2.907
## R-Squared                       0.871    Coef. Var                       4.107
## Adj. R-Squared                   0.869    MSE                               8.450
## Pred R-Squared                   0.863    MAE                               2.104
## -----

```

```

## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##

```

```

##                               ANOVA
## -----
##                               Sum of
##                               Squares    DF    Mean Square    F    Sig.
## -----
## Regression    7134.199    2    3567.099    422.166    0.0000
## Residual      1056.191    125    8.450
## Total         8190.390    127
## -----

```

```

##                               Parameter Estimates
## -----

```

```

##                               model
##                               Beta    Std. Error
Std. Beta    t    Sig    lower    upper
## -----
##                               (Intercept)    45.919    1.493
30.751    0.000    42.963    48.874
##                               Income.composition.of.resources    43.358    1.783
0.818    24.311    0.000    39.828    46.888
## Income.composition.of.resources:Adult.Mortality    -0.045    0.006
-0.268    -7.974    0.000    -0.056    -0.034
## -----

```

```

##
##

```

```

##
## Stepwise Selection: Step 3
##
## - HIV.AIDS added
##
##
##              Model Summary
## -----
## R              0.942          RMSE              2.732
## R-Squared      0.887          Coef. Var         3.860
## Adj. R-Squared 0.884          MSE              7.465
## Pred R-Squared 0.874          MAE              2.009
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##
##              ANOVA
## -----
##              Sum of
##              Squares          DF          Mean Square          F          Sig.
## -----
## Regression    7264.780           3          2421.593        324.411      0.0000
## Residual      925.609          124           7.465
## Total        8190.390          127
## -----
##
##
##              Parameter Estimates
## -----
## -----
##              Std. Beta          t          Sig          lower          model          Beta          Std. Error
##              upper
## -----
##              (Intercept)          47.662          1.464
## 32.554          0.000          44.764          50.560
##              Income.composition.of.resources          40.288          1.830
## 0.760          22.015          0.000          36.665          43.910
##              HIV.AIDS          -0.828          0.198
## -0.158          -4.183          0.000          -1.220          -0.436
##              Income.composition.of.resources:Adult.Mortality          -0.035          0.006
## -0.209          -6.016          0.000          -0.046          -0.023
## -----
##
##
##
##              Model Summary
## -----
## R              0.942          RMSE              2.732
## R-Squared      0.887          Coef. Var         3.860

```

```

## Adj. R-Squared      0.884      MSE      7.465
## Pred R-Squared     0.874      MAE      2.009
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##              Sum of      DF      Mean Square      F      Sig.
##              Squares
## -----
## Regression      7264.780        3      2421.593      324.411      0.0000
## Residual        925.609       124        7.465
## Total          8190.390       127
## -----
##
##                               Parameter Estimates
## -----
##              Std. Beta      t      Sig      lower      upper      model      Beta      Std. Error
## -----
##              (Intercept)      47.662      1.464
## 32.554      0.000      44.764      50.560
##              Income.composition.of.resources      40.288      1.830
## 0.760      22.015      0.000      36.665      43.910
##              HIV.AIDS      -0.828      0.198
## -0.158      -4.183      0.000      -1.220      -0.436
## Income.composition.of.resources:Adult.Mortality      -0.035      0.006
## -0.209      -6.016      0.000      -0.046      -0.023
## -----
##
##
##
## Stepwise Selection: Step 4
##
## - Adult.Mortality:HIV.AIDS added
##
##                               Model Summary
## -----
## R      0.948      RMSE      2.597
## R-Squared      0.899      Coef. Var      3.669
## Adj. R-Squared      0.895      MSE      6.744
## Pred R-Squared      0.881      MAE      1.902
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error

```

MAE: Mean Absolute Error

##

ANOVA

	Sum of Squares	DF	Mean Square	F	Sig.
Regression	7360.922	4	1840.231	272.884	0.0000
Residual	829.468	123	6.744		
Total	8190.390	127			

##

##

Parameter Estimates

##

	Std. Beta	t	Sig.	lower	upper	Beta	Std. Error
--	-----------	---	------	-------	-------	------	------------

##

					(Intercept)	49.290	1.457
--	--	--	--	--	-------------	--------	-------

33.833	0.000	46.406	52.174		Income.composition.of.resources	39.384	1.756
--------	-------	--------	--------	--	---------------------------------	--------	-------

0.743	22.430	0.000	35.908	42.859			
-------	--------	-------	--------	--------	--	--	--

					HIV.AIDS	-2.359	0.447
--	--	--	--	--	----------	--------	-------

-0.450	-5.277	0.000	-3.244	-1.474			
--------	--------	-------	--------	--------	--	--	--

					Income.composition.of.resources:Adult.Mortality	-0.043	0.006
--	--	--	--	--	---	--------	-------

-0.258	-7.272	0.000	-0.055	-0.031			
--------	--------	-------	--------	--------	--	--	--

					Adult.Mortality:HIV.AIDS	0.005	0.001
--	--	--	--	--	--------------------------	-------	-------

0.329	3.776	0.000	0.002	0.007			
-------	-------	-------	-------	-------	--	--	--

##

##

##

##

##

##

Model Summary

##

R	0.948	RMSE	2.597
---	-------	------	-------

R-Squared	0.899	Coef. Var	3.669
-----------	-------	-----------	-------

Adj. R-Squared	0.895	MSE	6.744
----------------	-------	-----	-------

Pred R-Squared	0.881	MAE	1.902
----------------	-------	-----	-------

##

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

##

ANOVA

##

	Sum of Squares	DF	Mean Square	F	Sig.
--	-------------------	----	-------------	---	------

##

```

## Regression      7360.922          4      1840.231      272.884      0.0000
## Residual        829.468         123          6.744
## Total           8190.390         127
## -----
##
##                                     Parameter Estimates
## -----
##
##                               model      Beta      Std. Error
## Std. Beta      t          Sig      lower      upper
## -----
##
##                               (Intercept)      49.290          1.457
## 33.833      0.000      46.406      52.174
##                               Income.composition.of.resources      39.384          1.756
## 0.743      22.430      0.000      35.908      42.859
##                               HIV.AIDS      -2.359          0.447
## -0.450      -5.277      0.000      -3.244      -1.474
## Income.composition.of.resources:Adult.Mortality      -0.043          0.006
## -0.258      -7.272      0.000      -0.055      -0.031
##                               Adult.Mortality:HIV.AIDS      0.005          0.001
## 0.329      3.776      0.000      0.002      0.007
## -----
##
##
##
## Stepwise Selection: Step 5
##
## - Adult.Mortality added
##
##                               Model Summary
## -----
## R                               0.948      RMSE                               2.596
## R-Squared                       0.900      Coef. Var                       3.668
## Adj. R-Squared                  0.895      MSE                               6.741
## Pred R-Squared                  0.875      MAE                               1.919
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7367.949          5      1473.590      218.591      0.0000
## Residual        822.441         122          6.741
## Total           8190.390         127

```


Parameter Estimates						
Beta	Std. Error					
t	Sig.	lower	upper			
(Intercept)						
51.166	2.345					
Income.composition.of.resources	36.763					
HIV.AIDS	-2.439					
Adult.Mortality	-0.012					
Income.composition.of.resources:Adult.Mortality	-0.026					
Adult.Mortality:HIV.AIDS	0.005					
Model Summary						
R	0.948	RMSE	2.596			
R-Squared	0.900	Coef. Var	3.668			
Adj. R-Squared	0.895	MSE	6.741			
Pred R-Squared	0.875	MAE	1.919			
ANOVA						
Sum of Squares	DF	Mean Square	F	Sig.		
Regression	7367.949	5	1473.590	218.591	0.0000	
Residual	822.441	122	6.741			
Total	8190.390	127				
Parameter Estimates						

```

##
Std. Beta      t          Sig      lower      upper      model      Beta      Std. Error
## -----
##
## (Intercept)      51.166      2.345
21.822      0.000      46.525      55.807
## Income.composition.of.resources      36.763      3.110
0.693      11.822      0.000      30.607      42.919
## HIV.AIDS      -2.439      0.454
-0.465      -5.375      0.000      -3.338      -1.541
## Adult.Mortality      -0.012      0.011
-0.143      -1.021      0.309      -0.034      0.011
## Income.composition.of.resources:Adult.Mortality      -0.026      0.018
-0.155      -1.456      0.148      -0.061      0.009
## Adult.Mortality:HIV.AIDS      0.005      0.001
0.364      3.889      0.000      0.002      0.008
## -----
##
##
##
## Stepwise Selection: Step 6
##
## - Adult.Mortality:HIV.AIDS added
##
##
## Model Summary
## -----
## R      0.942      RMSE      2.741
## R-Squared      0.887      Coef. Var      3.873
## Adj. R-Squared      0.883      MSE      7.515
## Pred R-Squared      0.869      MAE      2.005
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of      DF      Mean Square      F      Sig.
## Squares
## -----
## Regression      7266.004      4      1816.501      241.706      0.0000
## Residual      924.386      123      7.515
## Total      8190.390      127
## -----
##
##
## Parameter Estimates
## -----
##
## model      Beta      Std. Error

```

Std. Beta	t	Sig	lower	upper
## -----				
## (Intercept)				
21.343	0.000	42.642	51.360	47.001 2.202
## Income.composition.of.resources				
0.778	13.544	0.000	35.237	47.300 41.268 3.047
## HIV.AIDS				
-0.164	-4.023	0.000	-1.283	-0.437 -0.860 0.214
## Adult.Mortality				
0.056	0.403	0.687	-0.018	0.027 0.004 0.011
## Income.composition.of.resources:Adult.Mortality				
-0.250	-2.289	0.024	-0.078	-0.006 -0.042 0.018
## -----				
##				
##				
##				
## No more variables to be added/removed.				
##				
##				
## Final Model Output				
## -----				
##				
## Model Summary				
## -----				
## R		0.942	RMSE	2.741
## R-Squared		0.887	Coef. Var	3.873
## Adj. R-Squared		0.883	MSE	7.515
## Pred R-Squared		0.869	MAE	2.005
## -----				
## RMSE: Root Mean Square Error				
## MSE: Mean Square Error				
## MAE: Mean Absolute Error				
##				
## ANOVA				
## -----				
##	Sum of			
##	Squares	DF	Mean Square	F Sig.
## -----				
## Regression	7266.004	4	1816.501	241.706 0.0000
## Residual	924.386	123	7.515	
## Total	8190.390	127		
## -----				
##				
##				
## Parameter Estimates				
## -----				
##			model	Beta Std. Error
Std. Beta	t	Sig	lower upper	

```
## -----
##                               (Intercept)    47.001    2.202
21.343    0.000    42.642    51.360
##                               Income.composition.of.resources    41.268    3.047
0.778    13.544    0.000    35.237    47.300
##                               HIV.AIDS    -0.860    0.214
-0.164    -4.023    0.000    -1.283    -0.437
##                               Adult.Mortality    0.004    0.011
0.056    0.403    0.687    -0.018    0.027
## Income.composition.of.resources:Adult.Mortality    -0.042    0.018
-0.250    -2.289    0.024    -0.078    -0.006
## -----
```

life2015_bothmodel

```
##
##                               Stepwise Selection
Summary
## -----
##                               Added/
Adj.                               Removed    R-
## Step                               R-Square    C(p)    Variable    AIC    RMSE
## -----
##    1                               Income.composition.of.resources    addition
0.805    0.804    110.8500    692.0198    3.5562
##    2    Income.composition.of.resources:Adult.Mortality    addition
0.871    0.869    33.6620    641.3787    2.9068
##    3                               HIV.AIDS    addition
0.887    0.884    16.4170    626.4863    2.7321
##    4    Adult.Mortality:HIV.AIDS    addition
0.899    0.895    4.2470    614.4488    2.5969
##    5    Adult.Mortality    addition
0.900    0.895    5.2120    615.3598    2.5964
##    6    Adult.Mortality:HIV.AIDS    removal
0.887    0.883    18.2370    628.3170    2.7414
## -----
```

#Life2015_bothmodel\$mallows_cp

```
paste("Stepwise procedure with BOTH indicated to consider only the
interaction terms: Income.composition.of.resources * Adult.Mortality and
Adult.Mortality * HIV.AIDS.")
```

```
## [1] "Stepwise procedure with BOTH indicated to consider only the
interaction terms: Income.composition.of.resources * Adult.Mortality and
Adult.Mortality * HIV.AIDS."
```

```
# (F.3.2) FORWARD
life2015_forwardmodel = ols_step_forward_p(life2015_reduced_modelF, pent =
0.05, details = TRUE)
```

```
## Forward Selection Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1. Income.composition.of.resources
```

```
## 2. Adult.Mortality
```

```
## 3. HIV.AIDS
```

```
## 4. Income.composition.of.resources:Adult.Mortality
```

```
## 5. Income.composition.of.resources:HIV.AIDS
```

```
## 6. Adult.Mortality:HIV.AIDS
```

```
##
```

```
## We are selecting variables based on p value...
```

```
##
```

```
##
```

```
## Forward Selection: Step 1
```

```
##
```

```
## - Income.composition.of.resources
```

```
##
```

```
## Model Summary
```

```
## -----
## R                0.897      RMSE                3.556
## R-Squared        0.805      Coef. Var            5.025
## Adj. R-Squared   0.804      MSE                12.647
## Pred R-Squared   0.799      MAE                2.765
## -----
```

```
## RMSE: Root Mean Square Error
```

```
## MSE: Mean Square Error
```

```
## MAE: Mean Absolute Error
```

```
##
```

```
## ANOVA
```

```
## -----
## Sum of
## Squares      DF      Mean Square      F      Sig.
## -----
## Regression    6596.898      1      6596.898    521.627    0.0000
## Residual      1593.492     126      12.647
## Total         8190.390     127
## -----
```

```
##
```

```
## Parameter Estimates
```

```
## -----
## Sig      lower      model      Beta      Std. Error      Std. Beta
## t
## -----
```

```

-----
##              (Intercept)    38.615    1.443
26.763    0.000    35.760    41.470
## Income.composition.of.resources    47.584    2.083    0.897
22.839    0.000    43.461    51.707
## -----
##
##
##
## Forward Selection: Step 2
##
## - Income.composition.of.resources:Adult.Mortality
##
##              Model Summary
## -----
## R              0.933    RMSE              2.907
## R-Squared      0.871    Coef. Var          4.107
## Adj. R-Squared 0.869    MSE              8.450
## Pred R-Squared 0.863    MAE              2.104
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    7134.199      2      3567.099    422.166    0.0000
## Residual      1056.191     125        8.450
## Total         8190.390     127
## -----
##
##              Parameter Estimates
## -----
## -----
##              model      Beta      Std. Error
## Std. Beta      t      Sig      lower      upper
## -----
##              (Intercept)    45.919    1.493
30.751    0.000    42.963    48.874
##              Income.composition.of.resources    43.358    1.783
0.818    24.311    0.000    39.828    46.888
## Income.composition.of.resources:Adult.Mortality    -0.045    0.006
-0.268    -7.974    0.000    -0.056    -0.034
## -----

```

```

##
##
##
## Forward Selection: Step 3
##
## - HIV.AIDS
##
##                               Model Summary
## -----
## R                               0.942           RMSE           2.732
## R-Squared                       0.887           Coef. Var     3.860
## Adj. R-Squared                  0.884           MSE           7.465
## Pred R-Squared                  0.874           MAE           2.009
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7264.780           3      2421.593      324.411      0.0000
## Residual        925.609          124           7.465
## Total          8190.390          127
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               model      Beta      Std. Error
## Std. Beta      t      Sig      lower      upper
## -----
##                               -----
##                               (Intercept)      47.662      1.464
## 32.554      0.000      44.764      50.560
##                               Income.composition.of.resources      40.288      1.830
## 0.760      22.015      0.000      36.665      43.910
##                               HIV.AIDS      -0.828      0.198
## -0.158      -4.183      0.000      -1.220      -0.436
## Income.composition.of.resources:Adult.Mortality      -0.035      0.006
## -0.209      -6.016      0.000      -0.046      -0.023
## -----
##
##
##
## Forward Selection: Step 4
##

```

- Adult.Mortality:HIV.AIDS

##

Model Summary

```
## -----
## R                0.948          RMSE                2.597
## R-Squared        0.899          Coef. Var            3.669
## Adj. R-Squared   0.895          MSE                 6.744
## Pred R-Squared   0.881          MAE                 1.902
## -----
```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

```
## -----
##              Sum of          DF      Mean Square      F      Sig.
##              Squares
## -----
## Regression    7360.922          4      1840.231    272.884    0.0000
## Residual      829.468         123          6.744
## Total        8190.390         127
## -----
```

##

##

Parameter Estimates

```
## -----
##              Std. Beta      t      Sig      lower      upper      model      Beta      Std. Error
## -----
```

```
##              (Intercept)      49.290      1.457
33.833      0.000      46.406      52.174
##              Income.composition.of.resources      39.384      1.756
0.743      22.430      0.000      35.908      42.859
##              HIV.AIDS      -2.359      0.447
-0.450      -5.277      0.000      -3.244      -1.474
## Income.composition.of.resources:Adult.Mortality      -0.043      0.006
-0.258      -7.272      0.000      -0.055      -0.031
##              Adult.Mortality:HIV.AIDS      0.005      0.001
0.329      3.776      0.000      0.002      0.007
## -----
```

##

##

##

Forward Selection: Step 5

##

- Adult.Mortality

##

Model Summary


```

## -----
## R                0.948      RMSE                2.596
## R-Squared        0.900      Coef. Var            3.668
## Adj. R-Squared   0.895      MSE                 6.741
## Pred R-Squared   0.875      MAE                 1.919
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##              Sum of          DF      Mean Square      F      Sig.
##              Squares
## -----
## Regression      7367.949         5      1473.590    218.591    0.0000
## Residual        822.441        122         6.741
## Total          8190.390        127
## -----
##
##                               Parameter Estimates
## -----
## -----
##              Std. Beta      t      Sig      lower      upper      model      Beta      Std. Error
## -----
##              (Intercept)
## 21.822      0.000      46.525      55.807
##              Income.composition.of.resources
## 0.693      11.822      0.000      30.607      42.919
##              HIV.AIDS
## -0.465      -5.375      0.000      -3.338      -1.541
##              Adult.Mortality
## -0.143      -1.021      0.309      -0.034      0.011
## Income.composition.of.resources:Adult.Mortality
## -0.155      -1.456      0.148      -0.061      0.009
##              Adult.Mortality:HIV.AIDS
## 0.364      3.889      0.000      0.002      0.008
## -----
##
##
##
## No more variables to be added.
##
## Variables Entered:
##
## + Income.composition.of.resources
## + Income.composition.of.resources:Adult.Mortality

```

```

## + HIV.AIDS
## + Adult.Mortality:HIV.AIDS
## + Adult.Mortality
##
##
## Final Model Output
## -----
##
##                               Model Summary
## -----
## R                               0.948           RMSE           2.596
## R-Squared                       0.900           Coef. Var     3.668
## Adj. R-Squared                   0.895           MSE           6.741
## Pred R-Squared                   0.875           MAE           1.919
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7367.949           5      1473.590      218.591      0.0000
## Residual        822.441          122        6.741
## Total          8190.390          127
## -----
##
##                               Parameter Estimates
## -----
##                               Beta      Std. Error
## -----
##                               (Intercept)      51.166      2.345
## Std. Beta      t      Sig      lower      upper
## -----
##                               Income.composition.of.resources      36.763      3.110
## 0.693      11.822      0.000      30.607      42.919
##                               HIV.AIDS      -2.439      0.454
## -0.465      -5.375      0.000      -3.338      -1.541
##                               Adult.Mortality      -0.012      0.011
## -0.143      -1.021      0.309      -0.034      0.011
## Income.composition.of.resources:Adult.Mortality      -0.026      0.018
## -0.155      -1.456      0.148      -0.061      0.009
##                               Adult.Mortality:HIV.AIDS      0.005      0.001
## 0.364      3.889      0.000      0.002      0.008
## -----

```

```
life2015_forwardmodel
```

```
##
```

```
##
```

Selection Summary

```
## -----
```

```
##      Variable
```

```
Adj.
```

```
## Step
```

```
Square
```

```
C(p)
```

```
AIC
```

```
Entered
```

```
RMSE
```

```
R-Square
```

```
R-
```

```
## -----
```

```
##      1      Income.composition.of.resources      0.8054
```

```
0.8039      110.8498      692.0198      3.5562
```

```
##      2      Income.composition.of.resources:Adult.Mortality      0.8710
```

```
0.8690      33.6621      641.3787      2.9068
```

```
##      3      HIV.AIDS      0.8870
```

```
0.8843      16.4169      626.4863      2.7321
```

```
##      4      Adult.Mortality:HIV.AIDS      0.8987
```

```
0.8954      4.2475      614.4488      2.5969
```

```
##      5      Adult.Mortality      0.8996
```

```
0.8955      5.2118      615.3598      2.5964
```

```
## -----
```

```
#life2015_forwardmodel$mallows_cp
```

```
paste("Stepwise procedure with FORWARD also indicated to consider only the  
interaction terms: Income.composition.of.resources * Adult.Mortality and  
Adult.Mortality * HIV.AIDS.")
```

```
## [1] "Stepwise procedure with FORWARD also indicated to consider only the  
interaction terms: Income.composition.of.resources * Adult.Mortality and  
Adult.Mortality * HIV.AIDS."
```

```
# (F.3.3) BACKWARD
```

```
life2015_backwardmodel = ols_step_backward_p(life2015_reduced_modelF, prem =  
0.05, details = TRUE)
```

```
## Backward Elimination Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1 . Income.composition.of.resources
```

```
## 2 . Adult.Mortality
```

```
## 3 . HIV.AIDS
```

```
## 4 . Income.composition.of.resources:Adult.Mortality
```

```
## 5 . Income.composition.of.resources:HIV.AIDS
```

```
## 6 . Adult.Mortality:HIV.AIDS
```

```
##
```

```
## We are eliminating variables based on p value...
```

```

##
## - Income.composition.of.resources:HIV.AIDS
##
## Backward Elimination: Step 1
##
## Variable Income.composition.of.resources:HIV.AIDS Removed
##
##                               Model Summary
## -----
## R                               0.948           RMSE           2.596
## R-Squared                       0.900           Coef. Var     3.668
## Adj. R-Squared                   0.895           MSE           6.741
## Pred R-Squared                   0.875           MAE           1.919
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      7367.949           5      1473.590      218.591      0.0000
## Residual        822.441          122           6.741
## Total          8190.390          127
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               model      Beta      Std. Error
## Std. Beta      t      Sig      lower      upper
## -----
##                               -----
##                               (Intercept)      51.166      2.345
## 21.822      0.000      46.525      55.807
##                               Income.composition.of.resources      36.763      3.110
## 0.693      11.822      0.000      30.607      42.919
##                               Adult.Mortality      -0.012      0.011
## -0.143      -1.021      0.309      -0.034      0.011
##                               HIV.AIDS      -2.439      0.454
## -0.465      -5.375      0.000      -3.338      -1.541
## Income.composition.of.resources:Adult.Mortality      -0.026      0.018
## -0.155      -1.456      0.148      -0.061      0.009
##                               Adult.Mortality:HIV.AIDS      0.005      0.001
## 0.364      3.889      0.000      0.002      0.008
## -----
##
##

```

```

##
## - Income.composition.of.resources:Adult.Mortality
##
## Backward Elimination: Step 2
##
## Variable Income.composition.of.resources:Adult.Mortality Removed
##
##                               Model Summary
## -----
## R                               0.948          RMSE          2.608
## R-Squared                       0.898          Coef. Var    3.685
## Adj. R-Squared                   0.895          MSE          6.803
## Pred R-Squared                   0.874          MAE          1.963
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      7353.659              4          1838.415          270.248          0.0000
## Residual         836.731             123              6.803
## Total           8190.390             127
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               t          Sig          lower          upper          Beta          Std. Error          Std. Beta
## -----
##                               (Intercept)          53.350          1.810
## 29.468          0.000          49.766          56.933
## ## Income.composition.of.resources          33.346          2.050          0.629
## 16.270          0.000          29.289          37.403
## ##                               Adult.Mortality          -0.027          0.004          -0.336          -
## 7.166          0.000          -0.035          -0.020
## ##                               HIV.AIDS          -2.510          0.453          -0.478          -
## 5.538          0.000          -3.407          -1.613
## ##                               Adult.Mortality:HIV.AIDS          0.005          0.001          0.395
## 4.322          0.000          0.003          0.008
## -----
##
##
##
##

```

No more variables satisfy the condition of p value = 0.05

##

##

Variables Removed:

##

- Income.composition.of.resources:HIV.AIDS

- Income.composition.of.resources:Adult.Mortality

##

##

Final Model Output

##

Model Summary

## R	0.948	RMSE	2.608
------	-------	------	-------

## R-Squared	0.898	Coef. Var	3.685
--------------	-------	-----------	-------

## Adj. R-Squared	0.895	MSE	6.803
-------------------	-------	-----	-------

## Pred R-Squared	0.874	MAE	1.963
-------------------	-------	-----	-------

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

	Sum of				
	Squares	DF	Mean Square	F	Sig.

## Regression	7353.659	4	1838.415	270.248	0.0000
---------------	----------	---	----------	---------	--------

## Residual	836.731	123	6.803		
-------------	---------	-----	-------	--	--

## Total	8190.390	127			
----------	----------	-----	--	--	--

##

Parameter Estimates

			model	Beta	Std. Error	Std. Beta	
	Sig	lower	upper				

##			(Intercept)	53.350	1.810		
29.468	0.000	49.766	56.933				

## Income.composition.of.resources				33.346	2.050	0.629	
16.270	0.000	29.289	37.403				

## Adult.Mortality				-0.027	0.004	-0.336	-
7.166	0.000	-0.035	-0.020				

## HIV.AIDS				-2.510	0.453	-0.478	-
5.538	0.000	-3.407	-1.613				

## Adult.Mortality:HIV.AIDS				0.005	0.001	0.395	
4.322	0.000	0.003	0.008				

```
## -----
life2015_backwardmodel

##
##
##                                     Elimination Summary
## -----
##      Variable
Adj.
## Step      C(p)      AIC      Removed      R-Square      R-
Square      RMSE
## -----
##      1      Income.composition.of.resources:HIV.AIDS      0.8996
0.8955      5.2118      615.3598      2.5964
##      2      Income.composition.of.resources:Adult.Mortality      0.8978
0.8945      5.3179      615.5647      2.6082
## -----

#life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD to consider only the interaction term
Adult.Mortality and HIV.AIDS.")

## [1] "Stepwise procedure with BACKWARD to consider only the interaction
term Adult.Mortality and HIV.AIDS."

# (F.4) BEST SUBSET
#OPTION 1
ks = ols_step_best_subset(life2015_reduced_modelF, details = TRUE)

#Summary of Selected model based on cp, aic, AdjustedR2
rsquare <- c(ks$rsq)
cp <- c(ks$cp)
aic <- c(ks$aic)
AdjustedR2 <- c(ks$adjr)
cbind(rsquare, cp, aic, AdjustedR2)

##      rsquare      cp      aic AdjustedR2
## [1,] 0.8054437 110.849829 692.0198 0.8038996
## [2,] 0.8710451 33.662094 641.3787 0.8689818
## [3,] 0.8889036 14.104980 624.2984 0.8862158
## [4,] 0.8987781 4.185469 614.3838 0.8954863
## [5,] 0.8995847 5.211830 615.3598 0.8954693
## [6,] 0.8997602 7.000000 617.1359 0.8947896

#OPTION 2
best.subset <- regsubsets(Life.expectancy ~ (Income.composition.of.resources
```

```

+ Adult.Mortality + HIV.AIDS)^2, data = life2015[-c(119, 53), ], nv = 6)
summary(best.subset)

## Subset selection object
## Call: regsubsets.formula(Life.expectancy ~
(Income.composition.of.resources +
##   Adult.Mortality + HIV.AIDS)^2, data = life2015[-c(119, 53),
##   ], nv = 6)
## 6 Variables (and intercept)
##
##                               Forced in Forced out
## Income.composition.of.resources      FALSE      FALSE
## Adult.Mortality                      FALSE      FALSE
## HIV.AIDS                            FALSE      FALSE
## Income.composition.of.resources:Adult.Mortality      FALSE      FALSE
## Income.composition.of.resources:HIV.AIDS              FALSE      FALSE
## Adult.Mortality:HIV.AIDS                          FALSE      FALSE
## 1 subsets of each size up to 6
## Selection Algorithm: exhaustive
##           Income.composition.of.resources Adult.Mortality HIV.AIDS
## 1 ( 1 ) "*"
## 2 ( 1 ) "*"
## 3 ( 1 ) "*"
## 4 ( 1 ) "*"
## 5 ( 1 ) "*"
## 6 ( 1 ) "*"
##           Income.composition.of.resources:Adult.Mortality
## 1 ( 1 ) " "
## 2 ( 1 ) "*"
## 3 ( 1 ) "*"
## 4 ( 1 ) " "
## 5 ( 1 ) "*"
## 6 ( 1 ) "*"
##           Income.composition.of.resources:HIV.AIDS Adult.Mortality:HIV.AIDS
## 1 ( 1 ) " "
## 2 ( 1 ) " "
## 3 ( 1 ) "*"
## 4 ( 1 ) "*"
## 5 ( 1 ) " "
## 6 ( 1 ) "*"

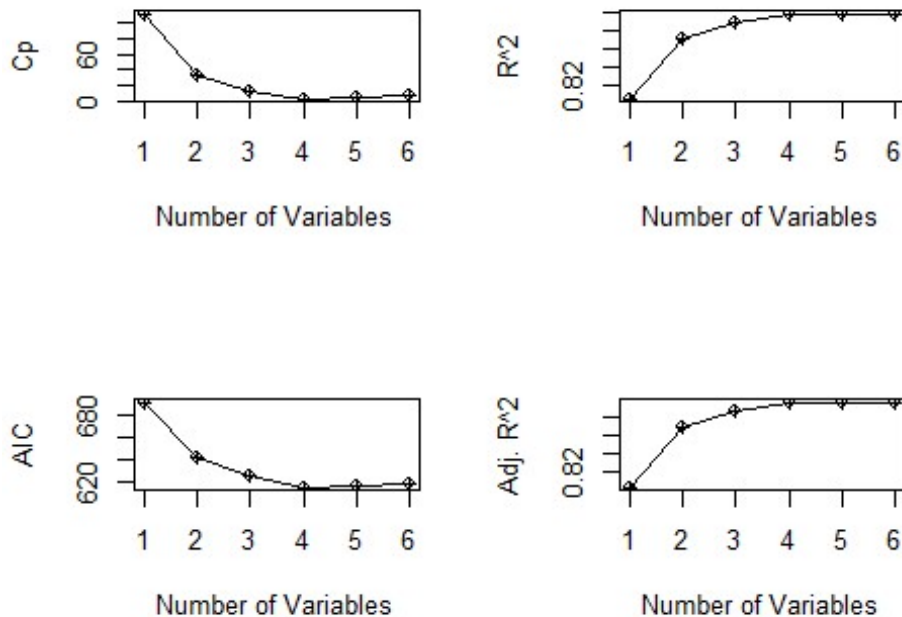
reg.summary <- summary(best.subset)

#Plotting
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"R^2")
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"AIC")

```



```
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Adj. R^2")
```



```
paste("Based on Marlow's Cp the minimum value was obtained with the  
interaction terms Income.composition.of.resources * Adult.Mortality and  
Adult.Mortality * HIV.AIDS, but without main variable HIV.AIDS. So, as the  
main effect should be in the model, we considered the second lowest Marlow's  
Cp that considers only the interaction term Adult.Mortality * HIV.AIDS.")
```

```
## [1] "Based on Marlow's Cp the minimum value was obtained with the  
interaction terms Income.composition.of.resources * Adult.Mortality and  
Adult.Mortality * HIV.AIDS, but without main variable HIV.AIDS. So, as the  
main effect should be in the model, we considered the second lowest Marlow's  
Cp that considers only the interaction term Adult.Mortality * HIV.AIDS."
```

```
#MODEL DIAGNOSTICS
```

```
# (F.5) CHECKING LINEARITY
```

```
#Residuals plot
```

```
plot(life2015_reduced_modelF, which = 1)
```

```
# a Scale Location Plot
```

```
plot(life2015_reduced_modelF, which = 3)
```

```
paste("It is verified some slightly change in the pattern for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts values in the last plot, but in overall the average of residuals almost horizontal and we can say that the residuals are equally spread that we can conclude that the linearity assumption is respected."
```

```
# (F.6) Heteroscedasticity Test - the Breush-Pagan test  
# Ho : heteroscedasticity is NOT presented (homoscedasticity)  
# Ha: heteroscedasticity is presented  
bptest(life2015_reduced_modelF)
```

```
##  
## studentized Breusch-Pagan test  
##  
## data: life2015_reduced_modelF  
## BP = 11.978, df = 6, p-value = 0.06245
```

```
paste("The output displays the Breush-Pagan test that from the model presented the p-value 0.062457 > 0.05, indicating the we should NOT REJECT the null hypthesis at 5 % level and consequently the test provides evidence that heteroscedasticity does not exist.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model presented the p-value 0.062457 > 0.05, indicating the we should NOT REJECT the null hypthesis at 5 % level and consequently the test provides evidence that heteroscedasticity does not exist."
```

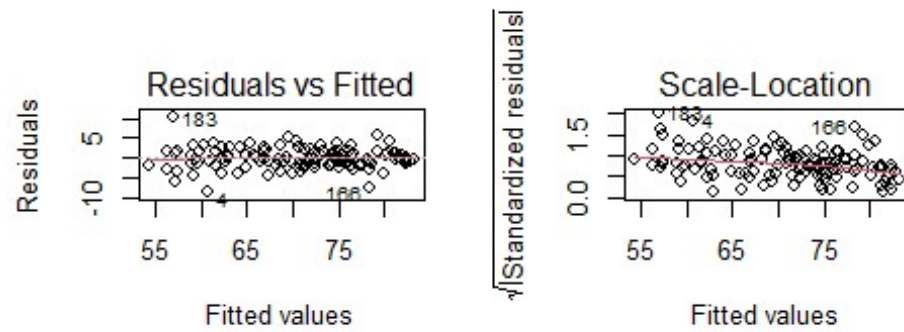
```
# (F.7) Testing for normality  
#NORMALITY - Shapiro-Wilk Test  
# Ho : the sample data is significantly normally distributed  
# Ha : the sample data is NOT significantly normally distributed  
shapiro.test(residuals(life2015_reduced_modelF))
```

```
##  
## Shapiro-Wilk normality test  
##  
## data: residuals(life2015_reduced_modelF)  
## W = 0.97532, p-value = 0.01935
```

```
paste("Shapiro-Wilk normality test presented the p-value = 0.01248 < 0.05 that indicated that we should REJECT the null hypothesis and conclude that the residuals are NOT normally distributed at 5 % level.")
```

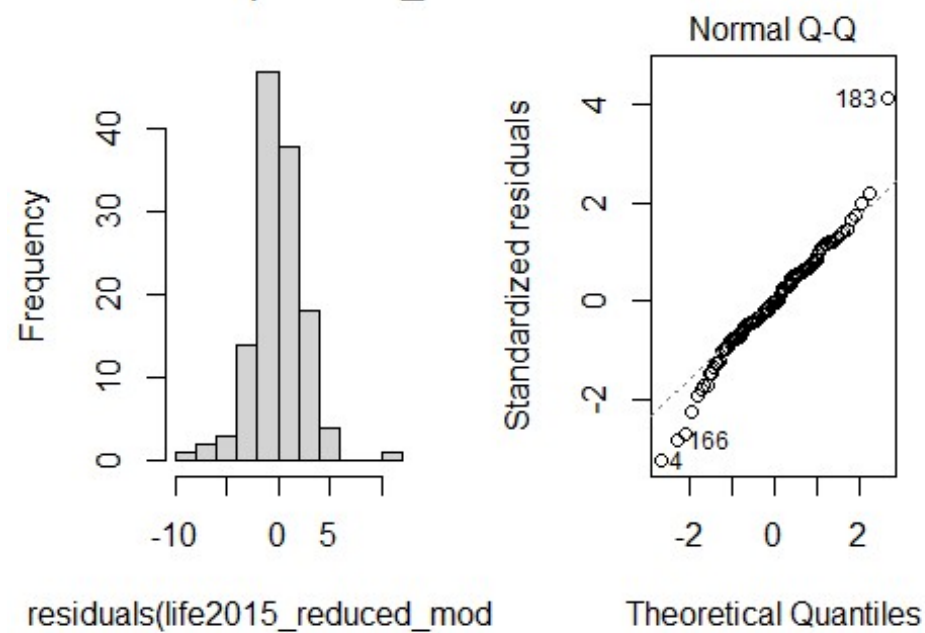
```
## [1] "Shapiro-Wilk normality test presented the p-value = 0.01248 < 0.05 that indicated that we should REJECT the null hypothesis and conclude that the residuals are NOT normally distributed at 5 % level."
```

```
#Plots  
par(mfrow = c(1,2))
```



```
hist(residuals(life2015_reduced_modelF))
plot(life2015_reduced_modelF, which = 2)
```

1 of residuals(life2015_redi



```

paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")

## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."

# (F.8) Multicollinearity test
#VIF
imcdiag(life2015_reduced_modelF, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_reduced_modelF, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##
##                                     VIF detection
## Income.composition.of.resources      4.2569          0
## Adult.Mortality                     32.9464          1
## HIV.AIDS                           128.4596          1
## Income.composition.of.resources:Adult.Mortality  20.6179          1
## Income.composition.of.resources:HIV.AIDS        85.0559          1
## Adult.Mortality:HIV.AIDS                 12.6316          1
##
## Multicollinearity may be due to Adult.Mortality HIV.AIDS
Income.composition.of.resources:Adult.Mortality
Income.composition.of.resources:HIV.AIDS Adult.Mortality:HIV.AIDS regressors
##
## 1 --> COLLINEARITY is detected by the test
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("The preliminary Multicollinearity test with all the interaction terms
indicated several redundant variables. ")

## [1] "The preliminary Multicollinearity test with all the interaction terms
indicated several redundant variables. "

# (F.9) CHECKING FOR INFLUENTIAL OUTLIERS
#Plotting Cook's distance versus
plot(life2015_reduced_modelF, which = 5)

paste("It is possible to verify in the standardized residuals plot that the
data point #93 is beyond Cook's distance lines. However, points #183 and #53
are near these lines.")

## [1] "It is possible to verify in the standardized residuals plot that the
data point #93 is beyond Cook's distance lines. However, points #183 and #53
are near these lines."

```

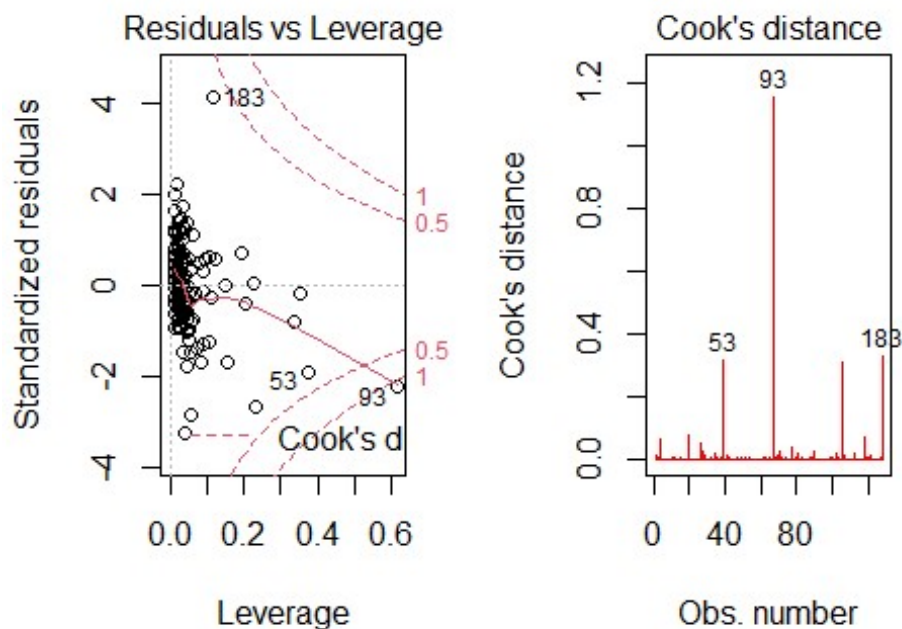
```

# Cook's Distance
#Checking if there is Cook statistics Larger than 0.5
life2015[cooks.distance(life2015_reduced_modelF) > 0.5, ]

##      Status Life.expectancy Adult.Mortality infant.deaths Hepatitis.B
Measles
## 92 Developing                74.9                98                1                81
39
##      BMI under.five.deaths Polio Diphtheria HIV.AIDS      GDP Population
## 92 66.1                1      75                81      0.1 846.6329      5851479
##      thinness..1.19.years thinness.5.9.years Income.composition.of.resources
## 92                4.9                4.9                0.763
##      Schooling
## 92                13.3

plot(life2015_reduced_modelF, pch = 18, col = "red", which = c(4))

```



```

paste("Based on the first interaction model, we can consider only the
interaction term Adult.deaths * HIV.AIDS seeing based on Best Subset,
Stepwise procedure with Backward and t-test and to remove data point #93.")

## [1] "Based on the first interaction model, we can consider only the
interaction term Adult.deaths * HIV.AIDS seeing based on Best Subset,
Stepwise procedure with Backward and t-test and to remove data point #93."

##### MODEL G - Model E with only the interaction term Adult * HIV and
without Data points #119, #53 and #93 #####
paste("Stepwise indicated to remove infant.deaths and under.five.deaths.")

```

```
## [1] "Stepwise indicated to remove infant.deaths and under.five.deaths."

# (G.1) F test to verify if at least one variable is related to Life
Expectancy
#The Analysis of Variance for Multiple Linear Regression
# Ho : Reduced model B without under.five.deaths
# Ha : Full model

#Reduced mode
life2015_reduced_modelG <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
Adult.Mortality * HIV.AIDS, data = life2015)

#Comparing the NULL model with the full model
anova(life2015_reduced_modelG, life2015_full_modelA)

## Analysis of Variance Table
##
## Model 1: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
## HIV.AIDS + Adult.Mortality * HIV.AIDS
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +
infant.deaths +
## Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
## Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
## thinness.5.9.years + Income.composition.of.resources + Schooling
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 125 854.19
## 2 113 820.58 12 33.606 0.3856 0.9663

paste("The output shows that Fcal = 0.3856 with df 12, 113 (p-value = 0.9963
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in
other words, the reduced model presents better prediction of Life Expectancy
compared with full model.")

## [1] "The output shows that Fcal = 0.3856 with df 12, 113 (p-value = 0.9963
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in
other words, the reduced model presents better prediction of Life Expectancy
compared with full model."

# (G.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_reduced_modelG <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
Adult.Mortality * HIV.AIDS, data = life2015[
c(119, 53, 93), ])

summary(life2015_reduced_modelG)
```

```
##
## Call:
## lm(formula = Life.expectancy ~ Income.composition.of.resources +
##     Adult.Mortality + HIV.AIDS + Adult.Mortality * HIV.AIDS,
##     data = life2015[-c(119, 53, 93), ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -8.2689 -1.4732  0.0081  1.6539 10.0724
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      53.376107    1.815257   29.404 < 2e-16 ***
## Income.composition.of.resources 33.287628    2.056622   16.186 < 2e-16 ***
## Adult.Mortality    -0.027104    0.003790   -7.151 6.92e-11 ***
## HIV.AIDS           -2.506238    0.454388   -5.516 1.98e-07 ***
## Adult.Mortality:HIV.AIDS      0.005434    0.001262    4.306 3.38e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.615 on 122 degrees of freedom
## Multiple R-squared:  0.8976, Adjusted R-squared:  0.8942
## F-statistic: 267.2 on 4 and 122 DF,  p-value: < 2.2e-16

paste("For the interaction model the t-test indicated all the coefficients
with p-value < 0.05 that indicates that all the variables and the interaction
term significant influence in Life Expectancy at 5 % level.")

## [1] "For the interaction model the t-test indicated all the coefficients
with p-value < 0.05 that indicates that all the variables and the interaction
term significant influence in Life Expectancy at 5 % level."

#confidence interval for the coefficients
confint(life2015_reduced_modelG, level = 0.95)

##              2.5 %       97.5 %
## (Intercept)      49.782625315 56.96958891
## Income.composition.of.resources 29.216340098 37.35891546
## Adult.Mortality    -0.034606774 -0.01960134
## HIV.AIDS           -3.405744334 -1.60673210
## Adult.Mortality:HIV.AIDS      0.002935985  0.00793218

paste("For a 95% confidence interval, none of the coefficients with Zero
between upper and lower values!")

## [1] "For a 95% confidence interval, none of the coefficients with Zero
between upper and lower values!"

# (G.3) STEPWISE REGRESSION PROCEDURE
# (G.3.1) BOTH
```

```
life2015_bothmodel = ols_step_both_p(life2015_reduced_modelG, pent = 0.05,
prem = 0.05, details = TRUE)
```

```
## Stepwise Selection Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1. Income.composition.of.resources
```

```
## 2. Adult.Mortality
```

```
## 3. HIV.AIDS
```

```
## 4. Adult.Mortality:HIV.AIDS
```

```
##
```

```
## We are selecting variables based on p value...
```

```
##
```

```
##
```

```
## Stepwise Selection: Step 1
```

```
##
```

```
## - Income.composition.of.resources added
```

```
##
```

```
## Model Summary
```

```
## -----
```

R	0.897	RMSE	3.567
R-Squared	0.805	Coef. Var	5.043
Adj. R-Squared	0.803	MSE	12.721
Pred R-Squared	0.798	MAE	2.773

```
## -----
```

```
## RMSE: Root Mean Square Error
```

```
## MSE: Mean Square Error
```

```
## MAE: Mean Absolute Error
```

```
##
```

```
## ANOVA
```

```
## -----
```

	Sum of Squares	DF	Mean Square	F	Sig.
Regression	6550.577	1	6550.577	514.952	0.0000
Residual	1590.095	125	12.721		
Total	8140.673	126			

```
## -----
```

```
##
```

```
## Parameter Estimates
```

```
## -----
```

	Sig	lower	model upper	Beta	Std. Error	Std. Beta
t						
26.683	0.000	35.781	41.514	38.647	1.448	

```
## -----
```



```

## Income.composition.of.resources    47.514          2.094          0.897
22.693    0.000    43.370    51.658
## -----
##
##
##
## Stepwise Selection: Step 2
##
## - Adult.Mortality added
##
##                               Model Summary
## -----
## R                               0.933          RMSE                2.920
## R-Squared                       0.870          Coef. Var          4.129
## Adj. R-Squared                   0.868          MSE                8.527
## Pred R-Squared                   0.862          MAE                2.128
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      7083.363              2          3541.682          415.364          0.0000
## Residual        1057.309             124              8.527
## Total           8140.673             126
## -----
##
##                               Parameter Estimates
## -----
##                               t          Sig          lower          model          Beta          Std. Error          Std. Beta
##                               t          Sig          lower          upper
## -----
##                               (Intercept)          49.802          1.843
27.019    0.000    46.154    53.450
## Income.composition.of.resources          37.147          2.158          0.701
17.210    0.000    32.874    41.419
##                               Adult.Mortality          -0.026          0.003          -0.322    -
7.905    0.000    -0.032    -0.019
## -----
##
##
##

```

```

##                                     Model Summary
## -----
## R                                0.933          RMSE          2.920
## R-Squared                       0.870          Coef. Var      4.129
## Adj. R-Squared                   0.868          MSE           8.527
## Pred R-Squared                   0.862          MAE           2.128
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##               Sum of          DF      Mean Square      F      Sig.
##               Squares
## -----
## Regression      7083.363           2      3541.682    415.364    0.0000
## Residual        1057.309          124           8.527
## Total           8140.673          126
## -----
##
##                                     Parameter Estimates
## -----
## -----
##               Sig      lower      model      Beta      Std. Error      Std. Beta
##               t               upper
## -----
##               (Intercept)      49.802           1.843
## 27.019      0.000      46.154      53.450
## Income.composition.of.resources      37.147           2.158           0.701
## 17.210      0.000      32.874      41.419
## Adult.Mortality      -0.026           0.003      -0.322      -
## 7.905      0.000      -0.032      -0.019
## -----
##
##
##
## Stepwise Selection: Step 3
##
## - HIV.AIDS added
##
##                                     Model Summary
## -----
## R                                0.939          RMSE          2.795
## R-Squared                       0.882          Coef. Var      3.952
## Adj. R-Squared                   0.879          MSE           7.811
## Pred R-Squared                   0.866          MAE           2.096
## -----

```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

	Sum of Squares	DF	Mean Square	F	Sig.
Regression	7179.967	3	2393.322	306.419	0.0000
Residual	960.706	123	7.811		
Total	8140.673	126			

##

Parameter Estimates

	Sig	lower	model upper	Beta	Std. Error	Std. Beta	
(Intercept)				50.145	1.767		
28.381	0.000	46.647	53.642				
Income.composition.of.resources				36.037	2.090	0.680	
17.245	0.000	31.901	40.174				
Adult.Mortality				-0.020	0.004	-0.244	-
5.455	0.000	-0.027	-0.013				
HIV.AIDS				-0.746	0.212	-0.143	-
3.517	0.001	-1.166	-0.326				

##

##

##

Model Summary

R	0.939	RMSE	2.795
R-Squared	0.882	Coef. Var	3.952
Adj. R-Squared	0.879	MSE	7.811
Pred R-Squared	0.866	MAE	2.096

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

	Sum of Squares	DF	Mean Square	F	Sig.
--	-------------------	----	-------------	---	------

```

## Regression      7179.967          3      2393.322      306.419      0.0000
## Residual        960.706         123          7.811
## Total           8140.673         126
## -----
##
##                                     Parameter Estimates
## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      50.145          1.767
28.381      0.000      46.647      53.642
## Income.composition.of.resources      36.037          2.090          0.680
17.245      0.000      31.901      40.174
##          Adult.Mortality      -0.020          0.004      -0.244      -
5.455      0.000      -0.027      -0.013
##          HIV.AIDS      -0.746          0.212      -0.143      -
3.517      0.001      -1.166      -0.326
## -----
##
##
##
## Stepwise Selection: Step 4
##
## - Adult.Mortality:HIV.AIDS added
##
##                                     Model Summary
## -----
## R          0.947          RMSE          2.615
## R-Squared    0.898          Coef. Var    3.697
## Adj. R-Squared 0.894          MSE          6.836
## Pred R-Squared 0.874          MAE          1.966
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of
##          Squares          DF      Mean Square          F          Sig.
## -----
## Regression    7306.723          4          1826.681      267.228      0.0000
## Residual       833.950         122           6.836
## Total          8140.673         126
## -----
##

```

```

##                                     Parameter Estimates
## -----
##                                     -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##                                     -----
##                                     (Intercept)      53.376      1.815
29.404      0.000      49.783      56.970
## Income.composition.of.resources      33.288      2.057      0.628
16.186      0.000      29.216      37.359
##                                     Adult.Mortality      -0.027      0.004      -0.337      -
7.151      0.000      -0.035      -0.020
##                                     HIV.AIDS      -2.506      0.454      -0.479      -
5.516      0.000      -3.406      -1.607
##                                     Adult.Mortality:HIV.AIDS      0.005      0.001      0.396
4.306      0.000      0.003      0.008
## -----
## -----
##
##
##
##                                     Model Summary
## -----
## R      0.947      RMSE      2.615
## R-Squared      0.898      Coef. Var      3.697
## Adj. R-Squared      0.894      MSE      6.836
## Pred R-Squared      0.874      MAE      1.966
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of      DF      Mean Square      F      Sig.
##          Squares
## -----
## Regression      7306.723      4      1826.681      267.228      0.0000
## Residual      833.950      122      6.836
## Total      8140.673      126
## -----
##
##
##                                     Parameter Estimates
## -----
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
## -----

```

```

##              (Intercept)    53.376      1.815
29.404    0.000    49.783    56.970
## Income.composition.of.resources    33.288      2.057      0.628
16.186    0.000    29.216    37.359
##              Adult.Mortality    -0.027      0.004     -0.337    -
7.151    0.000    -0.035    -0.020
##              HIV.AIDS    -2.506      0.454     -0.479    -
5.516    0.000    -3.406    -1.607
##      Adult.Mortality:HIV.AIDS      0.005      0.001      0.396
4.306    0.000      0.003      0.008
## -----
##
##
##
##
## Final Model Output
## -----
##
##              Model Summary
## -----
## R              0.947      RMSE              2.615
## R-Squared      0.898      Coef. Var          3.697
## Adj. R-Squared 0.894      MSE              6.836
## Pred R-Squared 0.874      MAE              1.966
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    7306.723      4      1826.681    267.228    0.0000
## Residual      833.950     122        6.836
## Total        8140.673     126
## -----
##
##              Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
## t              Sig      lower      upper
## -----
##              (Intercept)    53.376      1.815
29.404    0.000    49.783    56.970
## Income.composition.of.resources    33.288      2.057      0.628

```

```

16.186    0.000    29.216    37.359
##          Adult.Mortality    -0.027          0.004          -0.337    -
7.151    0.000    -0.035    -0.020
##          HIV.AIDS    -2.506          0.454          -0.479    -
5.516    0.000    -3.406    -1.607
##          Adult.Mortality:HIV.AIDS    0.005          0.001          0.396
4.306    0.000    0.003    0.008
## -----

```

```
life2015_bothmodel
```

```

##
##                                     Stepwise Selection Summary
## -----
##
## Step      Variable      Added/      R-Square      Adj.
## Square    C(p)      AIC      RMSE      Removed      R-
## -----
## 1      Income.composition.of.resources      addition      0.805
0.803      109.6180      687.3854      3.5666
## 2      Adult.Mortality      addition      0.870
0.868      33.6760      637.5609      2.9200
## 3      HIV.AIDS      addition      0.882
0.879      21.5430      627.3925      2.7947
## 4      Adult.Mortality:HIV.AIDS      addition      0.898
0.894      5.0000      611.4226      2.6145
## -----

```

```
#Life2015_bothmodel$mallows_cp
```

```
paste("Stepwise procedure with BOTH indicated to consider all the variables
and the interaction term.")
```

```
## [1] "Stepwise procedure with BOTH indicated to consider all the variables
and the interaction term."
```

```
# (G.3.2) FORWARD
```

```
life2015_forwardmodel = ols_step_forward_p(life2015_reduced_modelG, pent =
0.05, details = TRUE)
```

```
## Forward Selection Method
```

```
## -----
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1. Income.composition.of.resources
```

```
## 2. Adult.Mortality
```

```
## 3. HIV.AIDS
```

```

## 4. Adult.Mortality:HIV.AIDS
##
## We are selecting variables based on p value...
##
## Forward Selection: Step 1
##
## - Income.composition.of.resources
##
##                               Model Summary
## -----
## R                               0.897          RMSE              3.567
## R-Squared                       0.805          Coef. Var        5.043
## Adj. R-Squared                   0.803          MSE              12.721
## Pred R-Squared                   0.798          MAE              2.773
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      6550.577           1      6550.577      514.952      0.0000
## Residual        1590.095          125           12.721
## Total           8140.673          126
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               Sig      lower      model      Beta      Std. Error      Std. Beta
##                               t      upper
## -----
##                               (Intercept)      38.647      1.448
## 26.683      0.000      35.781      41.514
## Income.composition.of.resources      47.514      2.094      0.897
## 22.693      0.000      43.370      51.658
## -----
##
##
##
## Forward Selection: Step 2
##
## - Adult.Mortality
##

```



```

##                                     Model Summary
## -----
## R                                0.933          RMSE          2.920
## R-Squared                       0.870          Coef. Var      4.129
## Adj. R-Squared                   0.868          MSE           8.527
## Pred R-Squared                   0.862          MAE           2.128
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##               Sum of          DF      Mean Square      F      Sig.
##               Squares
## -----
## Regression      7083.363           2       3541.682    415.364    0.0000
## Residual        1057.309          124           8.527
## Total           8140.673          126
## -----
##
##                                     Parameter Estimates
## -----
##
##               Sig      lower      model      Beta      Std. Error      Std. Beta
## t               Sig      lower      upper
## -----
##               (Intercept)      49.802          1.843
## 27.019      0.000      46.154      53.450
## Income.composition.of.resources      37.147          2.158          0.701
## 17.210      0.000      32.874      41.419
## Adult.Mortality      -0.026          0.003      -0.322      -
## 7.905      0.000      -0.032      -0.019
## -----
##
##
##
## Forward Selection: Step 3
##
## - HIV.AIDS
##
##                                     Model Summary
## -----
## R                                0.939          RMSE          2.795
## R-Squared                       0.882          Coef. Var      3.952
## Adj. R-Squared                   0.879          MSE           7.811
## Pred R-Squared                   0.866          MAE           2.096
## -----

```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

	Sum of Squares	DF	Mean Square	F	Sig.
Regression	7179.967	3	2393.322	306.419	0.0000
Residual	960.706	123	7.811		
Total	8140.673	126			

##

Parameter Estimates

	Sig	lower	model upper	Beta	Std. Error	Std. Beta	
(Intercept)				50.145	1.767		
28.381	0.000	46.647	53.642				
Income.composition.of.resources				36.037	2.090	0.680	
17.245	0.000	31.901	40.174				
Adult.Mortality				-0.020	0.004	-0.244	-
5.455	0.000	-0.027	-0.013				
HIV.AIDS				-0.746	0.212	-0.143	-
3.517	0.001	-1.166	-0.326				

##

##

##

##

Forward Selection: Step 4

##

- Adult.Mortality:HIV.AIDS

##

Model Summary

R	0.947	RMSE	2.615
R-Squared	0.898	Coef. Var	3.697
Adj. R-Squared	0.894	MSE	6.836
Pred R-Squared	0.874	MAE	1.966

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

```

## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    7306.723        4      1826.681    267.228    0.0000
## Residual      833.950       122        6.836
## Total        8140.673       126
## -----
##
##                               Parameter Estimates
## -----
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)      53.376      1.815
29.404      0.000      49.783      56.970
## Income.composition.of.resources      33.288      2.057      0.628
16.186      0.000      29.216      37.359
## Adult.Mortality      -0.027      0.004      -0.337      -
7.151      0.000      -0.035      -0.020
## HIV.AIDS      -2.506      0.454      -0.479      -
5.516      0.000      -3.406      -1.607
## Adult.Mortality:HIV.AIDS      0.005      0.001      0.396
4.306      0.000      0.003      0.008
## -----
##
##
##
##
## Variables Entered:
##
## + Income.composition.of.resources
## + Adult.Mortality
## + HIV.AIDS
## + Adult.Mortality:HIV.AIDS
##
##
## Final Model Output
## -----
##
##              Model Summary
## -----
## R      0.947      RMSE      2.615
## R-Squared      0.898      Coef. Var      3.697
## Adj. R-Squared      0.894      MSE      6.836
## Pred R-Squared      0.874      MAE      1.966
## -----

```

##

ANOVA

ANOVA					
	Sum of Squares	DF	Mean Square	F	Sig.
Regression	7306.723	4	1826.681	267.228	0.0000
Residual	833.950	122	6.836		
Total	8140.673	126			

##

```
##                                     Parameter Estimates
```

##			model	Beta	Std. Error	Std. Beta
t	Sig	lower	upper			
##						

##	(Intercept)	53.376	1.815
----	-------------	--------	-------

29.404	0.000	49.783	56.970
--------	-------	--------	--------

## Income.composition.of.resources	33.288	2.057	0.628
------------------------------------	--------	-------	-------

16.186	0.000	29.216	37.359
--------	-------	--------	--------

##	Adult.Mortality	-0.027	0.004	-0.337	-
----	-----------------	--------	-------	--------	---

7.151	0.000	-0.035	-0.020
-------	-------	--------	--------

##	HIV.AIDS	-2.506	0.454	-0.479	-
----	----------	--------	-------	--------	---

5.516 0.000 -3.406 -1.607

##	Adult.Mortality:HIV.AIDS	0.005	0.001	0.396
----	--------------------------	-------	-------	-------

4.306	0.000	0.003	0.008
-------	-------	-------	-------

life2015_forwardmodel

##

Selection Summary

##	Variable	Adj.
----	----------	------

##	Step	Entered	R-Square	R-Square	C(p)
----	------	---------	----------	----------	------

	AIC	RMSE
--	-----	------

##	1	Income.composition.of.resources	0.8047	0.8031
----	---	---------------------------------	--------	--------

109.6179	687.3854	3.5666
----------	----------	--------

##	2	Adult.Mortality	0.8701	0.8680
----	---	-----------------	--------	--------

33.6757	637.5609	2.9200
---------	----------	--------

##	3	HIV.AIDS	0.8820	0.8791
----	---	----------	--------	--------

```

21.5434    627.3925    2.7947
##      4      Adult.Mortality:HIV.AIDS          0.8976      0.8942
5.0000    611.4226    2.6145
## -----
-----

#life2015_forwardmodel$allows_cp
paste("Stepwise procedure with FORWARD indicated to consider all the
variables and the interaction term.")

## [1] "Stepwise procedure with FORWARD indicated to consider all the
variables and the interaction term."

# (G.3.3) BACKWARD
life2015_backwardmodel = ols_step_backward_p(life2015_reduced_modelG, prem =
0.05, details = TRUE)

## Backward Elimination Method
## -----
##
## Candidate Terms:
##
## 1 . Income.composition.of.resources
## 2 . Adult.Mortality
## 3 . HIV.AIDS
## 4 . Adult.Mortality:HIV.AIDS
##
## We are eliminating variables based on p value...
##
##
## No more variables satisfy the condition of p value = 0.05
##
##
## Variables Removed:
##
##
##
## Final Model Output
## -----
##
##                               Model Summary
## -----
## R                               0.947          RMSE          2.615
## R-Squared                       0.898          Coef. Var      3.697
## Adj. R-Squared                   0.894          MSE           6.836
## Pred R-Squared                   0.874          MAE           1.966
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##

```

```
##
## ANOVA
## -----
##          Sum of          DF      Mean Square      F      Sig.
##          Squares
## -----
## Regression    7306.723         4        1826.681    267.228    0.0000
## Residual      833.950        122          6.836
## Total         8140.673        126
## -----
##
## Parameter Estimates
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t
## -----
##          (Intercept)      53.376      1.815
29.404    0.000    49.783    56.970
## Income.composition.of.resources      33.288      2.057      0.628
16.186    0.000    29.216    37.359
## Adult.Mortality      -0.027      0.004      -0.337    -
7.151    0.000    -0.035    -0.020
## HIV.AIDS      -2.506      0.454      -0.479    -
5.516    0.000    -3.406    -1.607
## Adult.Mortality:HIV.AIDS      0.005      0.001      0.396
4.306    0.000      0.003      0.008
## -----
-----

life2015_backwardmodel

## [1] "No variables have been removed from the model."

#life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD indicated to consider all the
variables and the interaction term.")

## [1] "Stepwise procedure with BACKWARD indicated to consider all the
variables and the interaction term."

# (G.4) BEST SUBSET
#OPTION 1
ks = ols_step_best_subset(life2015_reduced_modelG, details = TRUE)

#Summary of Selected model based on cp, aic, AdjustedR2
rsquare <- c(ks$rsq)
cp <- c(ks$cp)
aic <- c(ks$aic)
AdjustedR2 <- c(ks$adjr)
cbind(rsquare, cp, aic, AdjustedR2)
```

```
##          rsquare          cp          aic AdjustedR2
## [1,] 0.8046727 109.61791 687.3854 0.8031101
## [2,] 0.8701202 33.67568 637.5609 0.8680253
## [3,] 0.8819869 21.54336 627.3925 0.8791086
## [4,] 0.8975577 5.00000 611.4226 0.8941989
```

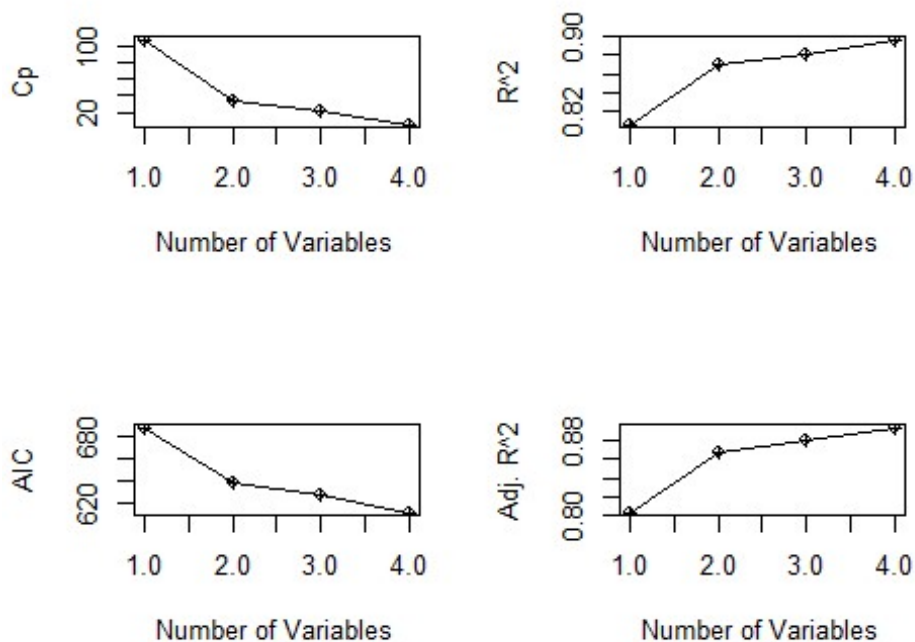
#OPTION 2

```
best.subset <- regsubsets(Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality + HIV.AIDS +
                        Adult.Mortality * HIV.AIDS, data = life2015[-
c(119, 53, 93), ], nv = 4)
summary(best.subset)
```

```
## Subset selection object
## Call: regsubsets.formula(Life.expectancy ~ Income.composition.of.resources
+
##      Adult.Mortality + HIV.AIDS + Adult.Mortality * HIV.AIDS,
##      data = life2015[-c(119, 53, 93), ], nv = 4)
## 4 Variables (and intercept)
##
##              Forced in Forced out
## Income.composition.of.resources FALSE FALSE
## Adult.Mortality FALSE FALSE
## HIV.AIDS FALSE FALSE
## Adult.Mortality:HIV.AIDS FALSE FALSE
## 1 subsets of each size up to 4
## Selection Algorithm: exhaustive
##      Income.composition.of.resources Adult.Mortality HIV.AIDS
## 1 ( 1 ) "*" " " " "
## 2 ( 1 ) "*" " " " "
## 3 ( 1 ) "*" "*" " "
## 4 ( 1 ) "*" "*" "*"
##      Adult.Mortality:HIV.AIDS
## 1 ( 1 ) " "
## 2 ( 1 ) " "
## 3 ( 1 ) " "
## 4 ( 1 ) "*"
reg.summary <- summary(best.subset)
```

#Plotting

```
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"R^2")
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"AIC")
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"Adj. R^2")
```



```
paste("Based on Marlow's Cp the minimum value was obtained with all the
variables and with the interaction term  Adult.Mortality * HIV.AIDS.")
```

```
## [1] "Based on Marlow's Cp the minimum value was obtained with all the
variables and with the interaction term  Adult.Mortality * HIV.AIDS."
```

```
#MODEL DIAGNOSTICS
```

```
# (G.5) CHECKING LINEARITY
```

```
#Residuals plot
```

```
plot(life2015_reduced_modelG, which = 1)
```

```
# a Scale Location Plot
```

```
plot(life2015_reduced_modelG, which = 3)
```

```
paste("It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected."
```

```
# (G.6) Heteroscedasticity Test - the Breush-Pagan test
```

```
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
```



```

# Ha: heteroscedasticity is presented
bptest(life2015_reduced_modelG)

##
## studentized Breusch-Pagan test
##
## data: life2015_reduced_modelG
## BP = 12.055, df = 4, p-value = 0.01695

paste("The output displays the Breush-Pagan test that from the model
presented the p-value 0.01695 < 0.05, indicating the we should REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity exists.")

## [1] "The output displays the Breush-Pagan test that from the model
presented the p-value 0.01695 < 0.05, indicating the we should REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity exists."

# (G.7) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_reduced_modelG))

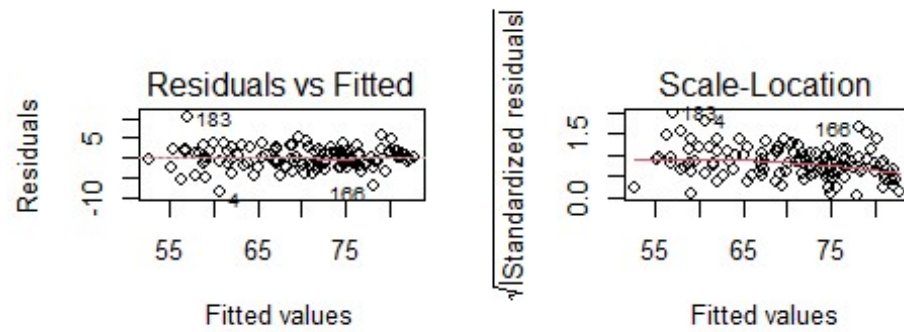
##
## Shapiro-Wilk normality test
##
## data: residuals(life2015_reduced_modelG)
## W = 0.9801, p-value = 0.05849

paste("Shapiro-Wilk normality test presented the p-value = 0.05849 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level.")

## [1] "Shapiro-Wilk normality test presented the p-value = 0.05849 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level."

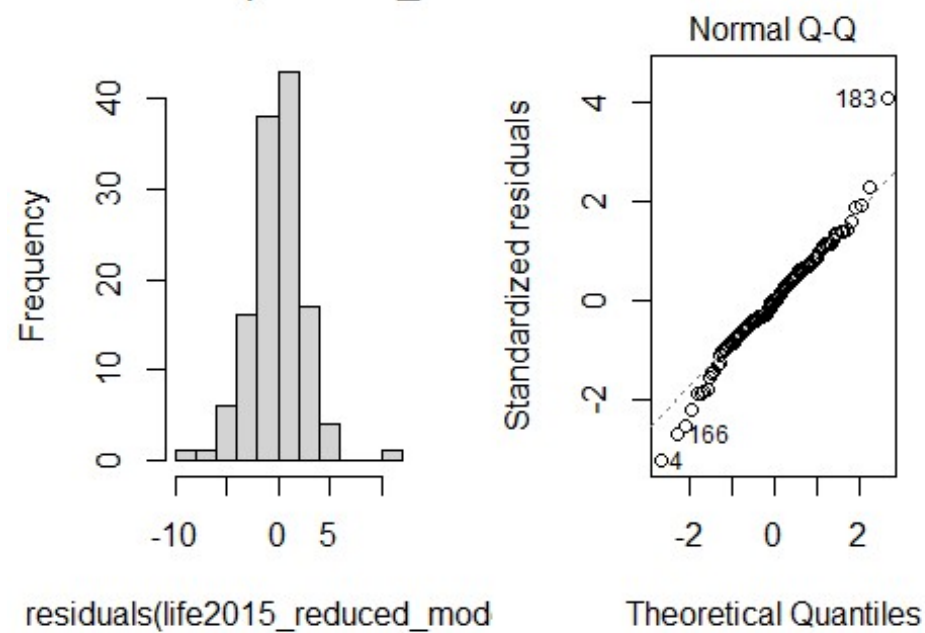
#Plots
par(mfrow = c(1,2))

```



```
hist(residuals(life2015_reduced_modelG))
plot(life2015_reduced_modelG, which = 2)
```

of residuals(life2015_red



```

paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")

## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."

# (G.8) Multicollinearity test
#VIF
imcdiag(life2015_reduced_modelG, method = "VIF")

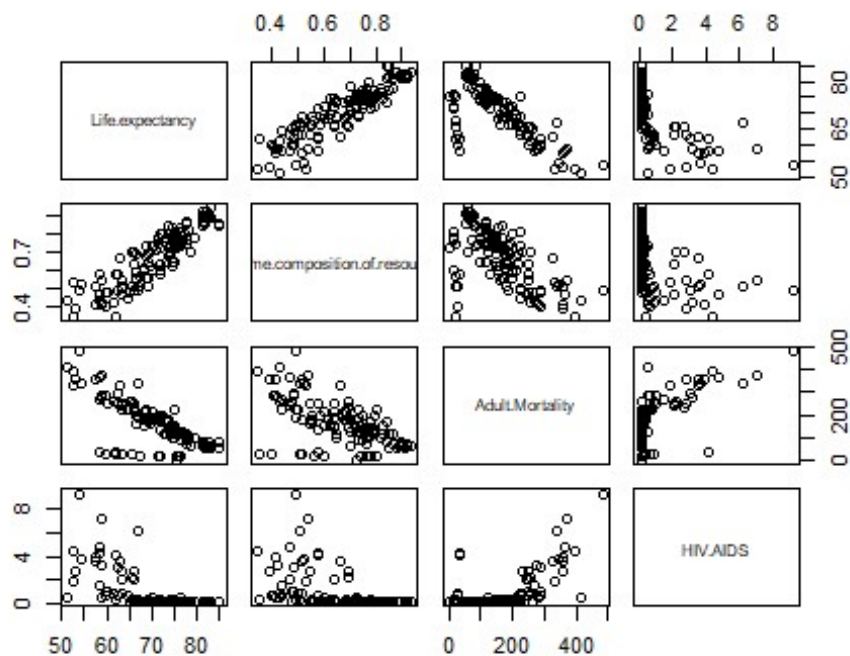
##
## Call:
## imcdiag(mod = life2015_reduced_modelG, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##                               VIF detection
## Income.composition.of.resources  1.7954          0
## Adult.Mortality                  2.6393          0
## HIV.AIDS                        8.9732          0
## Adult.Mortality:HIV.AIDS         10.0602          1
##
## Multicollinearity may be due to Adult.Mortality:HIV.AIDS regressors
##
## 1 --> COLLINEARITY is detected by the test
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity exists between the interaction term and HIV.AIDS
variable.")

## [1] "Multicollinearity exists between the interaction term and HIV.AIDS
variable."

# Plotting only the selected variables
pairs(~Life.expectancy + Income.composition.of.resources + Adult.Mortality +
HIV.AIDS +
                                Adult.Mortality * HIV.AIDS, data = life2015[-
c(119, 53, 93),])

```



```
paste("based on the plot, the variables HIV.AIDS and Income do not have a  
linear impact in Life Expectancy.")
```

```
## [1] "based on the plot, the variables HIV.AIDS and Income do not have a  
linear impact in Life Expectancy."
```

```
##### MODEL H - Model F with QUADRATIC term for HIV.AIDS and without Data  
points #119, #53 and #93 #####
```

```
# (G.1) F test to verify if at least one variable is related to Life  
Expectancy
```

```
#The Analysis of Variance for Multiple Linear Regression
```

```
# Ho : Reduced model B without under.five.deaths
```

```
# Ha : Full model
```

```
#Reduced mode
```

```
life2015_reduced_modelH <- lm(Life.expectancy ~  
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +  
+ I(HIV.AIDS ^ 2) + Adult.Mortality *  
HIV.AIDS, data = life2015)
```

```
#Comparing the NULL model with the full model
```

```
anova(life2015_reduced_modelH, life2015_full_modelA)
```

```
## Analysis of Variance Table
```

```
##
```

```
## Model 1: Life.expectancy ~ Income.composition.of.resources +
```

```

Adult.Mortality +
##      HIV.AIDS + I(HIV.AIDS^2) + Adult.Mortality * HIV.AIDS
## Model 2: Life.expectancy ~ factor(Status) + Adult.Mortality +
infant.deaths +
##      Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
##      Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
##      thinness.5.9.years + Income.composition.of.resources + Schooling
## Res.Df    RSS Df Sum of Sq    F Pr(>F)
## 1      124 854.19
## 2      113 820.58 11      33.606 0.4207 0.9442

```

`paste("The output shows that Fcal = 0.3856 with df 12, 113 (p-value = 0.9442 > alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in other words, the reduced model presents better prediction of Life Expectancy compared with full model.")`

```
## [1] "The output shows that Fcal = 0.3856 with df 12, 113 (p-value = 0.9442 > alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in other words, the reduced model presents better prediction of Life Expectancy compared with full model."
```

(G.2) PARTIAL TEST - Individual Coefficients Test (t-test)

Ho : Beta(i) = 0

Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

```

life2015_reduced_modelH <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
                                + I(HIV.AIDS^2) + Adult.Mortality * HIV.AIDS,
data = life2015[-c(119, 53, 93), ])

```

```
summary(life2015_reduced_modelH)
```

```

##
## Call:
## lm(formula = Life.expectancy ~ Income.composition.of.resources +
##      Adult.Mortality + HIV.AIDS + I(HIV.AIDS^2) + Adult.Mortality *
##      HIV.AIDS, data = life2015[-c(119, 53, 93), ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -8.2946 -1.4693  0.0136  1.6394 10.0905
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    53.376368    1.822693   29.284 < 2e-16 ***
## Income.composition.of.resources 33.306783    2.077819   16.030 < 2e-16 ***
## Adult.Mortality   -0.027272    0.004305   -6.334 4.23e-09 ***
## HIV.AIDS         -2.493582    0.480903   -5.185 8.77e-07 ***
## I(HIV.AIDS^2)    -0.007715    0.092653   -0.083  0.93378
## Adult.Mortality:HIV.AIDS      0.005552    0.001903    2.917 0.00421 **
## ---

```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.625 on 121 degrees of freedom
## Multiple R-squared:  0.8976, Adjusted R-squared:  0.8933
## F-statistic: 212 on 5 and 121 DF,  p-value: < 2.2e-16

paste("For the interaction model the t-test indicated the quadratic
coefficient with p-value = 0.93378 > 0.05 that indicates that this
coeffiecient pratically does not have influence in Life Expectancy at 5 %
level.")

## [1] "For the interaction model the t-test indicated the quadratic
coefficient with p-value = 0.93378 > 0.05 that indicates that this
coeffiecient pratically does not have influence in Life Expectancy at 5 %
level."

#confidence interval for the coefficients
confint(life2015_reduced_modelH, level = 0.95)

##
##                2.5 %          97.5 %
## (Intercept)      49.767867169  56.984869224
## Income.composition.of.resources 29.193192634 37.420373658
## Adult.Mortality  -0.035795200 -0.018748194
## HIV.AIDS          -3.445655721 -1.541508554
## I(HIV.AIDS^2)     -0.191146489  0.175717357
## Adult.Mortality:HIV.AIDS      0.001784496  0.009320143

paste("For a 95% confidence interval, the quadratic coefficient has Zero
between upper and lower values!")

## [1] "For a 95% confidence interval, the quadratic coefficient has Zero
between upper and lower values!"

## MODEL I - Model F with QUADRATIC term for Income.composition.of.resources
and without Data points #119, #53 and #93 ##

# (I.1) F test to verify if at Least one variable is related to Life
Expectancy
#The Analysis of Variance for Multiple Linear Regression
# Ho : Reduced model B without under.five.deaths
# Ha : Full model

#Reduced mode
life2015_reduced_modelI <- lm(Life.expectancy ~
Income.composition.of.resources +
                                I(Income.composition.of.resources ^ 2) +
Adult.Mortality + HIV.AIDS +
                                + Adult.Mortality * HIV.AIDS, data =
life2015)
```

#Comparing the NULL model with the full model

```
anova(life2015_reduced_modelI, life2015_full_modelA)
```

```
## Analysis of Variance Table
```

```
##
```

```
## Model 1: Life expectancy ~ Income.composition.of.resources +  
I(Income.composition.of.resources^2) +
```

```
## Adult.Mortality + HIV.AIDS + +Adult.Mortality * HIV.AIDS
```

```
## Model 2: Life expectancy ~ factor(Status) + Adult.Mortality +  
infant.deaths +
```

```
## Hepatitis.B + Measles + BMI + under.five.deaths + Polio +
```

```
## Diphtheria + HIV.AIDS + GDP + Population + thinness..1.19.years +
```

```
## thinness.5.9.years + Income.composition.of.resources + Schooling
```

```
## Res.Df RSS Df Sum of Sq F Pr(>F)
```

```
## 1 124 853.89
```

```
## 2 113 820.58 11 33.308 0.417 0.946
```

```
paste("The output shows that Fcal = 0.3856 with df 11, 113 (p-value = 0.946 >  
alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in  
other words, the reduced model presents better prediction of Life Expectancy  
compared with full model.")
```

```
## [1] "The output shows that Fcal = 0.3856 with df 11, 113 (p-value = 0.946  
> alpha = 0.05), indicating that we should NOT REJECT the null hypothesis, in  
other words, the reduced model presents better prediction of Life Expectancy  
compared with full model."
```

(I.2) PARTIAL TEST - Individual Coefficients Test (t-test)

Ho : Beta(i) = 0

Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

```
life2015_reduced_modelI <- lm(Life expectancy ~
```

```
Income.composition.of.resources +
```

```
I(Income.composition.of.resources ^ 2) +
```

```
Adult.Mortality + HIV.AIDS +
```

```
+ Adult.Mortality * HIV.AIDS, data =
```

```
life2015[-c(119, 53, 93), ])
```

```
summary(life2015_reduced_modelI)
```

```
##
```

```
## Call:
```

```
## lm(formula = Life expectancy ~ Income.composition.of.resources +
```

```
## I(Income.composition.of.resources^2) + Adult.Mortality +
```

```
## HIV.AIDS + +Adult.Mortality * HIV.AIDS, data = life2015[-c(119,
```

```
## 53, 93), ])
```

```
##
```

```
## Residuals:
```

```
## Min 1Q Median 3Q Max
```

```
## -8.2580 -1.4616 -0.0704 1.6614 10.0905
```

```
##
```

```

## Coefficients:
##
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 54.315126 4.498111 12.075 < 2e-16
***
## Income.composition.of.resources 30.205780 13.653933 2.212 0.0288
*
## I(Income.composition.of.resources^2) 2.359014 10.331285 0.228 0.8198
## Adult.Mortality -0.027011 0.003827 -7.059 1.14e-10
***
## HIV.AIDS -2.497996 0.457589 -5.459 2.59e-07
***
## Adult.Mortality:HIV.AIDS 0.005394 0.001279 4.218 4.79e-05
***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.625 on 121 degrees of freedom
## Multiple R-squared: 0.8976, Adjusted R-squared: 0.8934
## F-statistic: 212.1 on 5 and 121 DF, p-value: < 2.2e-16

paste("For the interaction model the t-test indicated the quadratic
coefficient with p-value = 0.8198 > 0.05 that indicates that this
coeffiecient pratically does not have influence in Life Expectancy at 5 %
level.")

## [1] "For the interaction model the t-test indicated the quadratic
coefficient with p-value = 0.8198 > 0.05 that indicates that this
coeffiecient pratically does not have influence in Life Expectancy at 5 %
level."

#confidence interval for the coefficients
confint(life2015_reduced_modelI, level = 0.95)

##
## 2.5 % 97.5 %
## (Intercept) 45.409928951 63.220322660
## Income.composition.of.resources 3.174217575 57.237342780
## I(Income.composition.of.resources^2) -18.094488962 22.812517109
## Adult.Mortality -0.034586755 -0.019435521
## HIV.AIDS -3.403914671 -1.592076980
## Adult.Mortality:HIV.AIDS 0.002862208 0.007926009

paste("For a 95% confidence interval, the quadratic coefficient has Zero
between upper and lower values!")

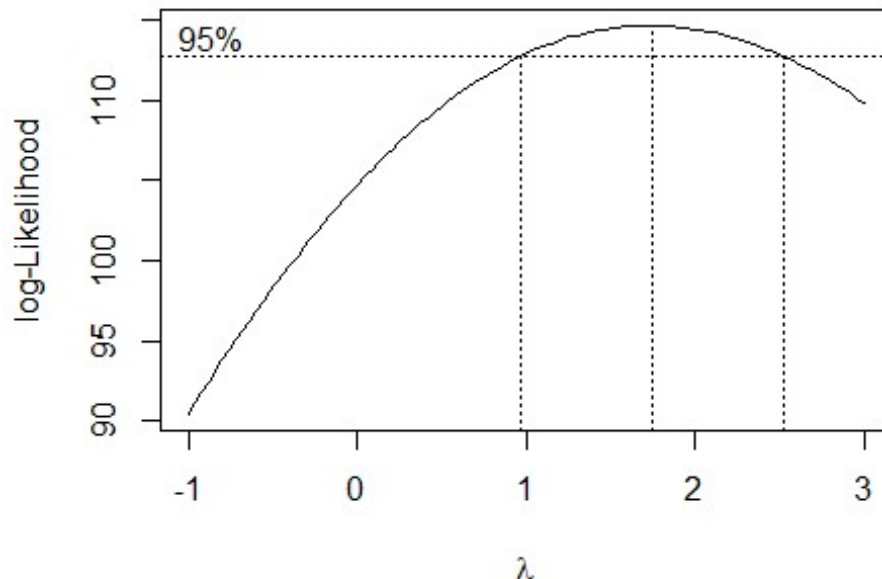
## [1] "For a 95% confidence interval, the quadratic coefficient has Zero
between upper and lower values!"

##### MODEL J - BOX COX TRANSFORMATION of MODEL G #####
#(J.1) BOX-COX TRANSFORMATION
# The necessity of transformation is due to the indication of
hetereosdasticity in model G

```


For this transformation we will be considering only the significant parameters obtained beforehand

```
bc = boxcox(life2015_reduced_modelG, lambda = seq(-1,3))
```



#Extract best Lambda

```
bestlambda = bc$x[which(bc$y == max(bc$y))]
```

```
paste("The best lambda for Box-Cox Transformation is: ", bestlambda)
```

```
## [1] "The best lambda for Box-Cox Transformation is: 1.74747474747475"
```

```
paste("From the output, as the best lambda would be 1.7474.")
```

```
## [1] "From the output, as the best lambda would be 1.7474."
```

#BOX-COX with LAMBDA = 1.7474

```
life2015_box_modelJ = lm((((Life.expectancy^1.7474) - 1)/1.7474) ~  
Income.composition.of.resources + Adult.Mortality +  
HIV.AIDS + Adult.Mortality * HIV.AIDS,  
data = life2015)
```

(J.2) PARTIAL TEST - Individual Coefficients Test (t-test)

Ho : Beta(i) = 0

Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

```
life2015_box_modelJ = lm((((Life.expectancy^1.7474) - 1)/1.7474) ~  
Income.composition.of.resources + Adult.Mortality +
```

```
HIV.AIDS + Adult.Mortality * HIV.AIDS, data = life2015[-
c(119,53,93),])
```

```
summary(life2015_box_modelJ)
```

```
##
## Call:
## lm(formula = (((Life.expectancy^1.7474) - 1)/1.7474) ~
Income.composition.of.resources +
##      Adult.Mortality + HIV.AIDS + Adult.Mortality * HIV.AIDS,
##      data = life2015[-c(119, 53, 93), ])
##
## Residuals:
##      Min        1Q    Median        3Q        Max
## -172.217  -36.176   -0.529   38.586  220.097
##
## Coefficients:
##                                Estimate Std. Error t value Pr(>|t|)
## (Intercept)                560.41500    42.96801   13.043 < 2e-16 ***
## Income.composition.of.resources 804.77614    48.68123   16.532 < 2e-16 ***
## Adult.Mortality              -0.64287     0.08971   -7.166 6.42e-11 ***
## HIV.AIDS                    -59.12437    10.75558   -5.497 2.15e-07 ***
## Adult.Mortality:HIV.AIDS         0.13519     0.02987    4.526 1.41e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 61.89 on 122 degrees of freedom
## Multiple R-squared:  0.8977, Adjusted R-squared:  0.8944
## F-statistic: 267.7 on 4 and 122 DF,  p-value: < 2.2e-16
```

```
paste("The t-test indicated that after the Box-Cox transformation the all the
coefficients selected previously have p-value < 0.05 that indicates all the
selected variables has significant influence in Life Expectancy at 5 %
level.")
```

```
## [1] "The t-test indicated that after the Box-Cox transformation the all
the coefficients selected previously have p-value < 0.05 that indicates all
the selected variables has significant influence in Life Expectancy at 5 %
level."
```

```
#confidence interval for the coefficients
```

```
confint(life2015_box_modelJ, level = 0.95)
```

```
##                                2.5 %        97.5 %
## (Intercept)                475.35554431 645.4744645
## Income.composition.of.resources 708.40678683 901.1455019
## Adult.Mortality              -0.82045870  -0.4652729
## HIV.AIDS                    -80.41612024 -37.8326265
## Adult.Mortality:HIV.AIDS         0.07605862   0.1943210
```

```
paste("For a 95% confidence interval, none coefficients with Zero between  
upper and lower values!")
```

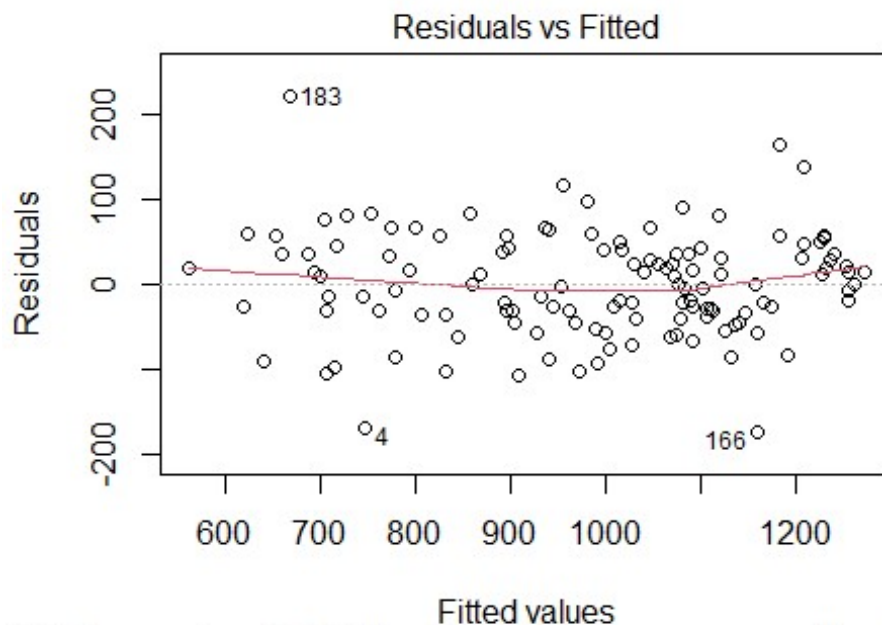
```
## [1] "For a 95% confidence interval, none coefficients with Zero between  
upper and lower values!"
```

```
#MODEL DIAGNOSTICS
```

```
# (J.3) CHECKING LINEARITY
```

```
#Residuals plot
```

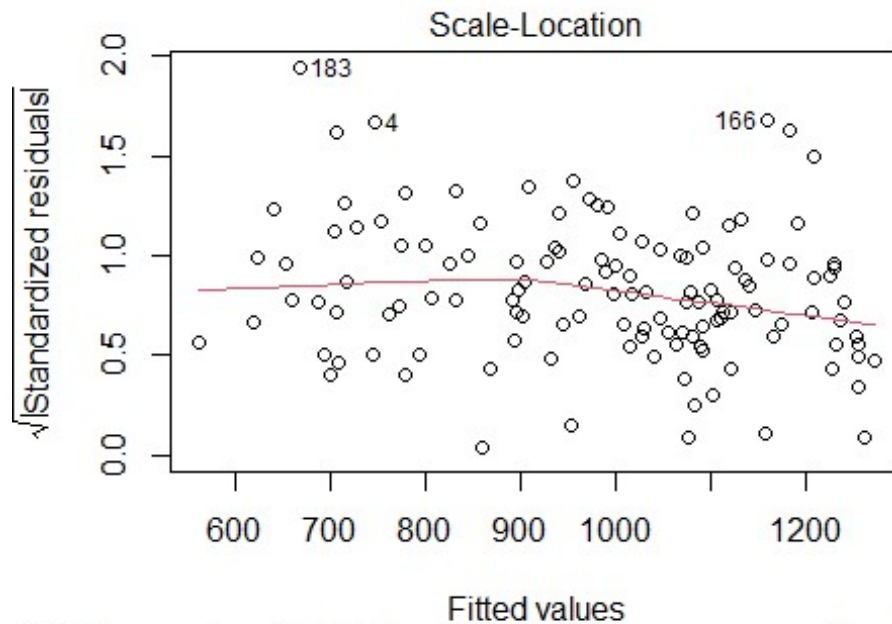
```
plot(life2015_box_modelJ, which = 1)
```



```
1((((Life.expectancy^1.7474) - 1)/1.7474) ~ Income.composition.of.res)
```

```
# a Scale Location Plot
```

```
plot(life2015_box_modelJ, which = 3)
```



```
(((Life.expectancy^1.7474) - 1)/1.7474) ~ Income.composition.of.resc
```

```
paste("It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected."
```

```
# (J.4) Heteroscedasticity Test - the Breush-Pagan test
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
# Ha: heteroscedasticity is presented
bptest(life2015_box_model)
```

```
##
## studentized Breusch-Pagan test
##
## data: life2015_box_model
## BP = 9.9668, df = 4, p-value = 0.04099
```

```
paste("The output displays the Breush-Pagan test that from the model
presented the p-value 0.04099 < 0.05, indicating the we should REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity exists.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model
presented the p-value 0.04099 < 0.05, indicating the we should REJECT the
```

null hypothesis at 5 % level and consequently the test provides evidence that heteroscedasticity exists."

```
# (J.5) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_model))

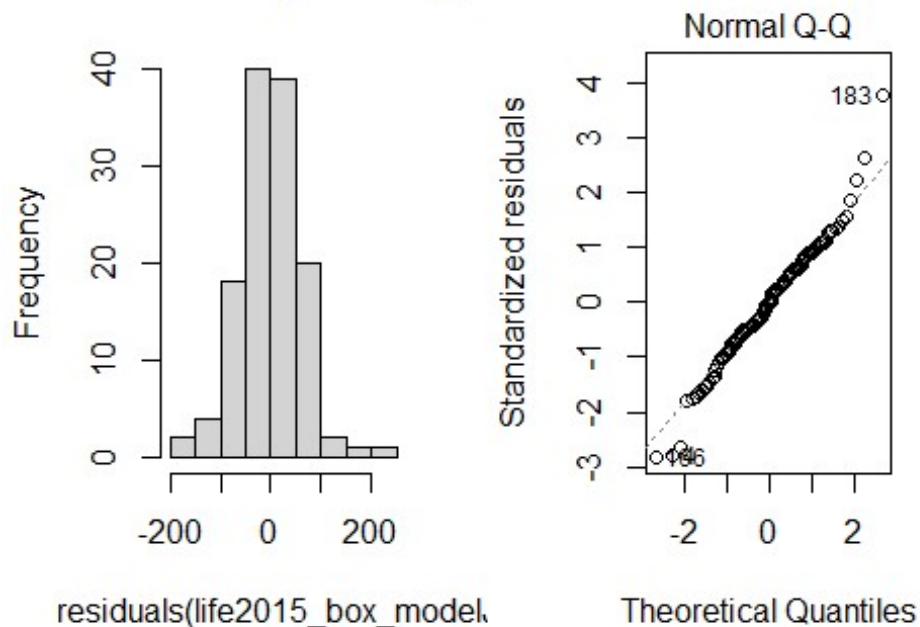
##
##  Shapiro-Wilk normality test
##
## data:  residuals(life2015_box_model)
## W = 0.98446, p-value = 0.1559

paste("Shapiro-Wilk normality test presented the p-value = 0.1559 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are ormallly distributed at 5 % level.")

## [1] "Shapiro-Wilk normality test presented the p-value = 0.1559 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are ormallly distributed at 5 % level."

#Plots
par(mfrow = c(1,2))
hist(residuals(life2015_box_model))
plot(life2015_box_model, which = 2)
```

am of residuals(life2015_b



```

paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")

## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."

# (J.6) Multicollinearity test
#VIF
imcdiag(life2015_box_modelJ, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_box_modelJ, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##                               VIF detection
## Income.composition.of.resources  1.7954          0
## Adult.Mortality                  2.6393          0
## HIV.AIDS                         8.9732          0
## Adult.Mortality:HIV.AIDS         10.0602          1
##
## Multicollinearity may be due to Adult.Mortality:HIV.AIDS regressors
##
## 1 --> COLLINEARITY is detected by the test
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. ")

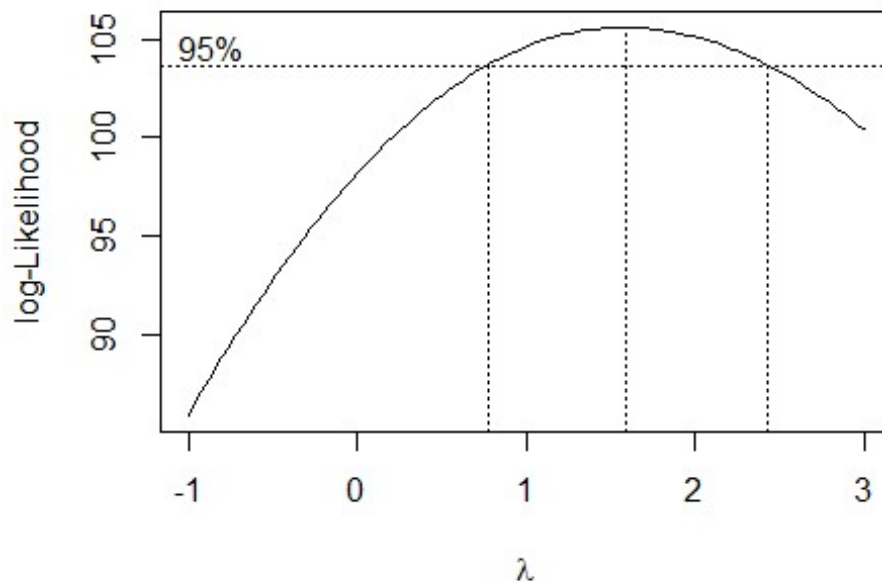
## [1] "Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. "

##### MODEL K - BOX COX TRANSFORMATION of MODEL E (no interaction
term) #####
paste("t-test of previous Box-Cox transformation indicated no significance of
interaction term after this transformation")

## [1] "t-test of previous Box-Cox transformation indicated no significance
of interaction term after this transformation"

#(K.1) BOX-COX TRANSFORMATION
# The necessity of transformation is due to the indication of
heterosdasticity in model E
# For this transformation we will be considering only the significant
parameters obtained beforehand
bc = boxcox(life2015_reduced_modelE, lambda = seq(-1,3))

```



```
#Extract best Lambda
bestlambda = bc$x[which(bc$y == max(bc$y))]
paste("The best lambda for Box-Cox Transformation is: ", bestlambda)

## [1] "The best lambda for Box-Cox Transformation is:  1.58585858585859"

paste("From the output, as the best lambda would be 1.5858.")

## [1] "From the output, as the best lambda would be 1.5858."

#BOX-COX with LAMBDA = 1.7474
life2015_box_modelK = lm((((Life.expectancy^1.5858) - 1)/1.5858) ~
Income.composition.of.resources + Adult.Mortality +
HIV.AIDS, data = life2015)

# (K.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) # 0 (i = 1, 2, ..., p)

life2015_box_modelK = lm((((Life.expectancy^1.5858) - 1)/1.5858) ~
Income.composition.of.resources + Adult.Mortality +
HIV.AIDS, data = life2015[-c(119,53,93),])

summary(life2015_box_modelK)
```

```
##
## Call:
## lm(formula = (((Life.expectancy^1.5858) - 1)/1.5858) ~
Income.composition.of.resources +
##     Adult.Mortality + HIV.AIDS, data = life2015[-c(119, 53, 93),
##     ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -108.705  -18.633    2.053   21.812   109.868
##
## Coefficients:
##                                Estimate Std. Error t value Pr(>|t|)
## (Intercept)                   290.82166    21.16787   13.739 < 2e-16 ***
## Income.composition.of.resources 438.16799    25.03692   17.501 < 2e-16 ***
## Adult.Mortality                 -0.23178     0.04323   -5.362 3.92e-07 ***
## HIV.AIDS                       -7.98789     2.54146   -3.143  0.0021 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 33.48 on 123 degrees of freedom
## Multiple R-squared:  0.8812, Adjusted R-squared:  0.8783
## F-statistic: 304.1 on 3 and 123 DF,  p-value: < 2.2e-16
```

```
paste("The t-test indicated that after the Box-Cox transformation the all the
coefficients selected previously have p-value < 0.05 that indicates all the
selected variables has significant influence in Life Expectancy at 5 %
level.")
```

```
## [1] "The t-test indicated that after the Box-Cox transformation the all
the coefficients selected previously have p-value < 0.05 that indicates all
the selected variables has significant influence in Life Expectancy at 5 %
level."
```

```
#confidence interval for the coefficients
confint(life2015_box_modelK, level = 0.95)
```

```
##                                2.5 %      97.5 %
## (Intercept)                   248.9211609 332.7221525
## Income.composition.of.resources 388.6089394 487.7270394
## Adult.Mortality                 -0.3173503  -0.1462165
## HIV.AIDS                       -13.0185493  -2.9572222
```

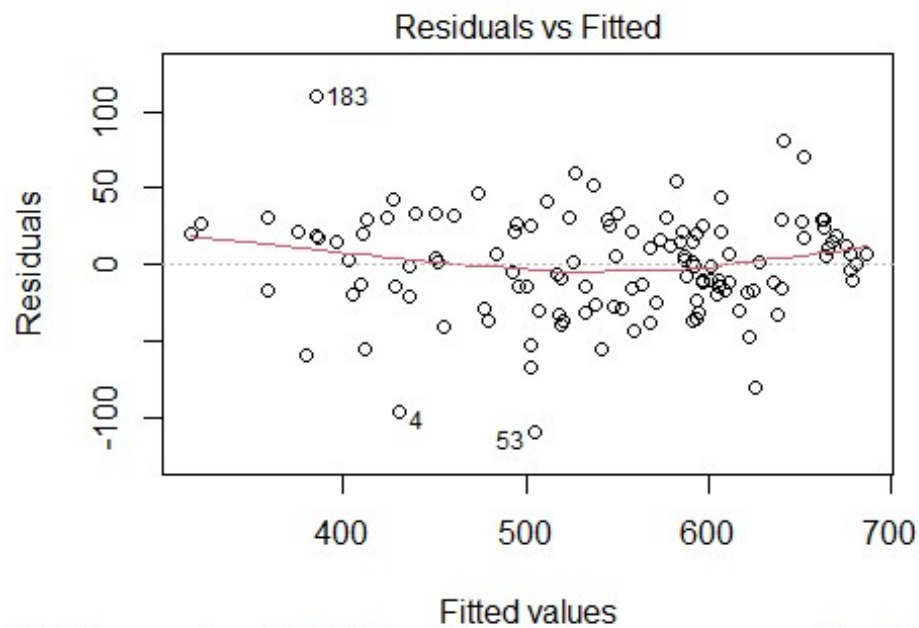
```
paste("For a 95% confidence interval, none coefficients with Zero between
upper and lower values!")
```

```
## [1] "For a 95% confidence interval, none coefficients with Zero between
upper and lower values!"
```

```
#MODEL DIAGNOSTICS
# (J.3) CHECKING LINEARITY
```

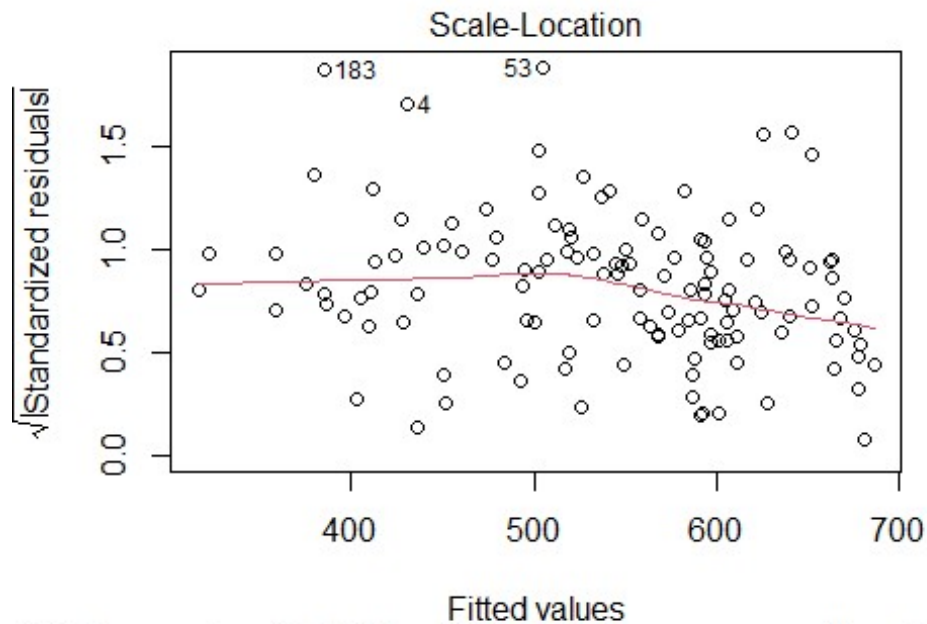


```
#Residuals plot
plot(life2015_box_modelK, which = 1)
```



```
l(((Life.expectancy^1.5858) - 1)/1.5858) ~ Income.composition.of.res
```

```
# a Scale Location Plot
plot(life2015_box_modelK, which = 3)
```



```
l(((Life.expectancy^1.5858) - 1)/1.5858) ~ Income.composition.of.resc
```

```
paste("It is verified some slightly change in the pattern for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected."
```

```
# (J.4) Heteroscedasticity Test - the Breush-Pagan test  
# Ho : heteroscedasticity is NOT presented (homoscedasticity)  
# Ha: heteroscedasticity is presented  
bptest(life2015_box_modelK)
```

```
##  
## studentized Breusch-Pagan test  
##  
## data: life2015_box_modelK  
## BP = 17.617, df = 3, p-value = 0.0005274
```

```
paste("The output displays the Breush-Pagan test that from the model  
presented the p-value 0.0005274 < 0.05, indicating the we should REJECT the  
null hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity exists.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model  
presented the p-value 0.0005274 < 0.05, indicating the we should REJECT the
```

null hypothesis at 5 % level and consequently the test provides evidence that heteroscedasticity exists."

```
# (J.5) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_modelK))

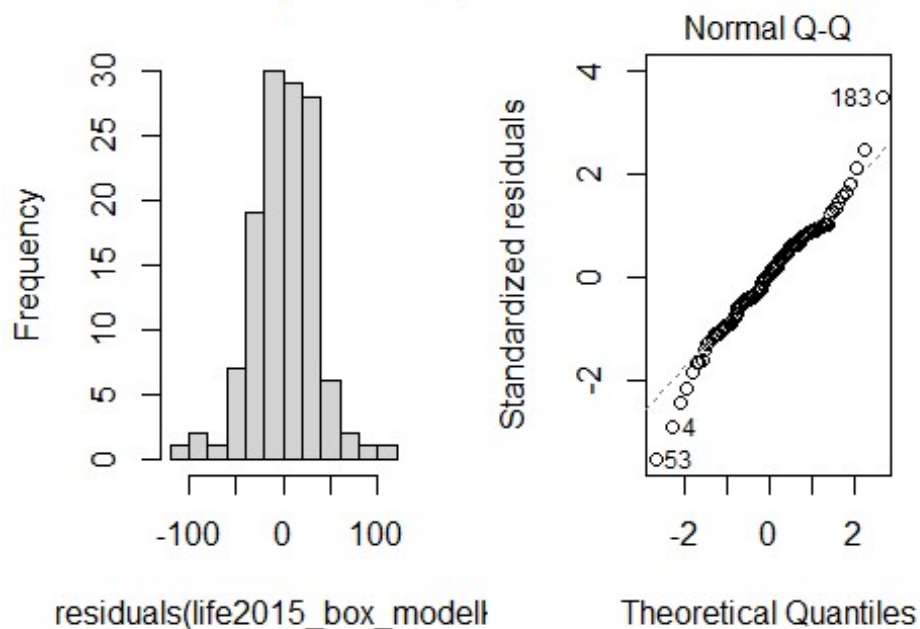
##
##  Shapiro-Wilk normality test
##
## data:  residuals(life2015_box_modelK)
## W = 0.9811, p-value = 0.07322

paste("Shapiro-Wilk normality test presented the p-value = 0.07322 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are ormally distributed at 5 % level.")

## [1] "Shapiro-Wilk normality test presented the p-value = 0.07322 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are ormally distributed at 5 % level."

#Plots
par(mfrow = c(1,2))
hist(residuals(life2015_box_modelK))
plot(life2015_box_modelK, which = 2)
```

am of residuals(life2015_b



```

paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")

## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."

# (J.6) Multicollinearity test
#VIF
imcdiag(life2015_box_modelK, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_box_modelK, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##                               VIF detection
## Income.composition.of.resources 1.6223      0
## Adult.Mortality                 2.0934      0
## HIV.AIDS                        1.7115      0
##
## NOTE: VIF Method Failed to detect multicollinearity
##
##
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. ")

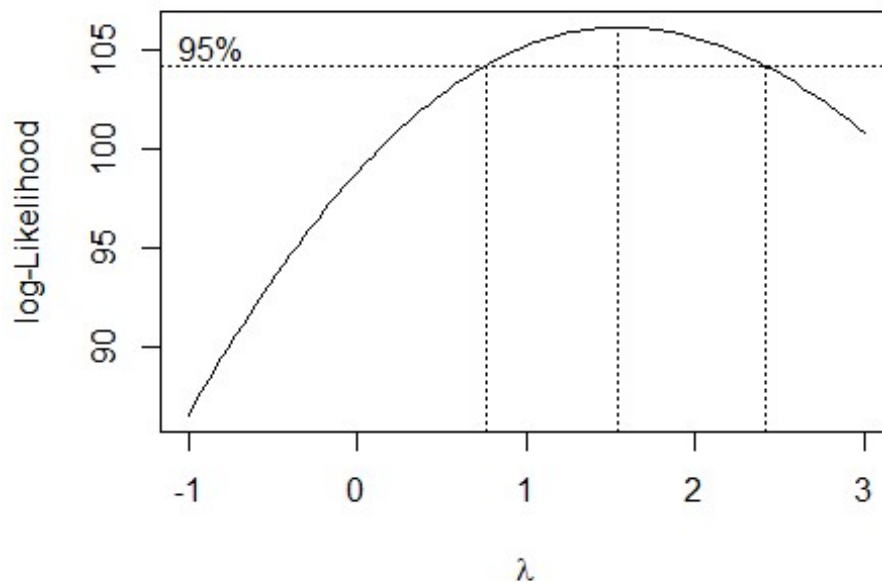
## [1] "Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. "

##### MODEL L - BOX COX TRANSFORMATION of MODEL C (under.five back)
#####
paste("t-test of previous Box-Cox transformation indicated no significance of
interaction term after this transformation")

## [1] "t-test of previous Box-Cox transformation indicated no significance
of interaction term after this transformation"

#(L.1) BOX-COX TRANSFORMATION
# The necessity of transformation is due to the indication of
heterosdasticity in model E
# For this transformation we will be considering only the significant
parameters obtained beforehand
bc = boxcox(life2015_reduced_modelC, lambda = seq(-1,3))

```



```
#Extract best Lambda
bestlambda = bc$x[which(bc$y == max(bc$y))]
paste("The best lambda for Box-Cox Transformation is: ", bestlambda)

## [1] "The best lambda for Box-Cox Transformation is:  1.54545454545455"

paste("From the output, as the best lambda would be 1.5454.")

## [1] "From the output, as the best lambda would be 1.5454."

#BOX-COX with LAMBDA = 1.7474
life2015_box_modelL = lm((((Life.expectancy^1.5454) - 1)/1.5454) ~
Income.composition.of.resources + Adult.Mortality +
HIV.AIDS + under.five.deaths, data = life2015)

# (K.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_box_modelL = lm((((Life.expectancy^1.5454) - 1)/1.5454) ~
Income.composition.of.resources + Adult.Mortality +
HIV.AIDS + under.five.deaths, data = life2015[
c(119,53,93),])

summary(life2015_box_modelL)
```

```
##
## Call:
## lm(formula = (((Life.expectancy^1.5454) - 1)/1.5454) ~
Income.composition.of.resources +
##      Adult.Mortality + HIV.AIDS + under.five.deaths, data = life2015[-
c(119,
##      53, 93), ])
##
## Residuals:
##      Min        1Q    Median        3Q        Max
## -91.997 -15.704   1.861  17.769  92.399
##
## Coefficients:
##                                Estimate Std. Error t value Pr(>|t|)
## (Intercept)                   259.43617   18.00961  14.405 < 2e-16 ***
## Income.composition.of.resources 365.33548   21.32025  17.136 < 2e-16 ***
## Adult.Mortality                 -0.19278    0.03647  -5.286 5.57e-07 ***
## HIV.AIDS                       -6.88956    2.14133  -3.217 0.00166 **
## under.five.deaths              -0.02203    0.02060  -1.070 0.28695
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 28.18 on 122 degrees of freedom
## Multiple R-squared:  0.8824, Adjusted R-squared:  0.8786
## F-statistic: 228.9 on 4 and 122 DF,  p-value: < 2.2e-16

paste("The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level.")

## [1] "The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level."

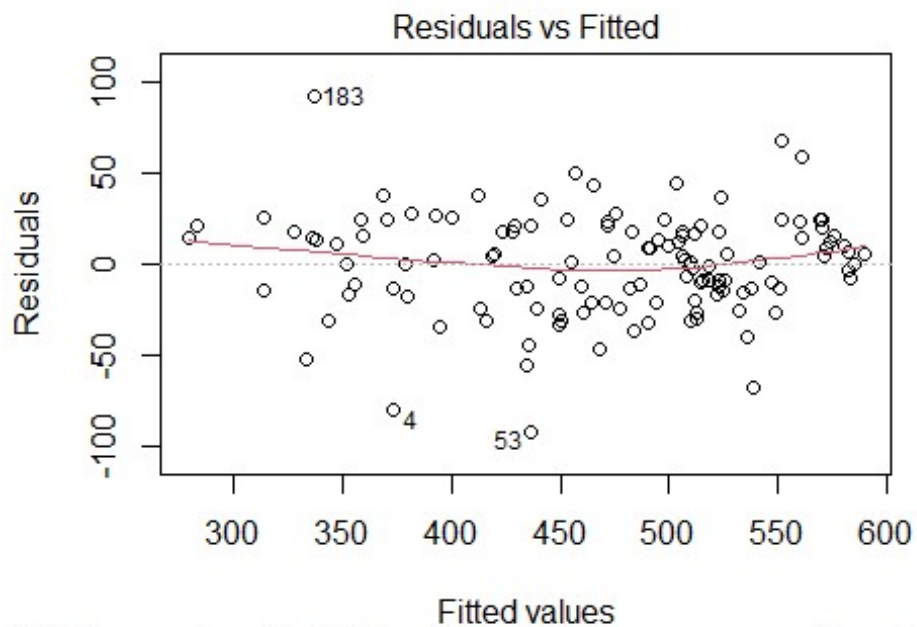
#confidence interval for the coefficients
confint(life2015_box_modelL, level = 0.95)

##                                2.5 %        97.5 %
## (Intercept)                   223.78434183 295.08799419
## Income.composition.of.resources 323.12991879 407.54104233
## Adult.Mortality                 -0.26498350 -0.12058359
## HIV.AIDS                       -11.12852662 -2.65058367
## under.five.deaths              -0.06280438  0.01874567

paste("For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!")

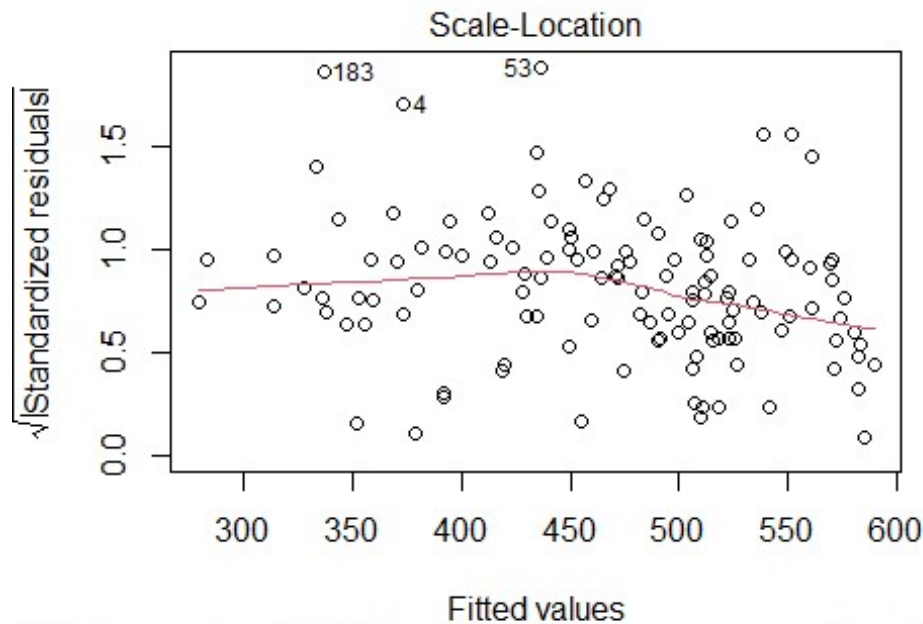
## [1] "For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!"
```

```
#MODEL DIAGNOSTICS
# (L.3) CHECKING LINEARITY
#Residuals plot
plot(life2015_box_modelL, which = 1)
```



$\ln(((\text{Life expectancy}^{1.5454}) - 1)/1.5454) \sim \text{Income.composition.of.res}$

```
# a Scale Location Plot
plot(life2015_box_modelL, which = 3)
```



```
(((Life.expectancy^1.5454) - 1)/1.5454) ~ Income.composition.of.resc
```

```
paste("It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected."
```

```
# (L.4) Heteroscedasticity Test - the Breush-Pagan test
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
# Ha: heteroscedasticity is presented
bptest(life2015_box_modelL)
```

```
##
## studentized Breusch-Pagan test
##
## data: life2015_box_modelL
## BP = 16.483, df = 4, p-value = 0.002435
```

```
paste("The output displays the Breush-Pagan test that from the model
presented the p-value 0.002435 < 0.05, indicating the we should REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity exists.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model
presented the p-value 0.002435 < 0.05, indicating the we should REJECT the
```


null hypothesis at 5 % level and consequently the test provides evidence that heteroscedasticity exists."

```
# (L.5) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_modelL))

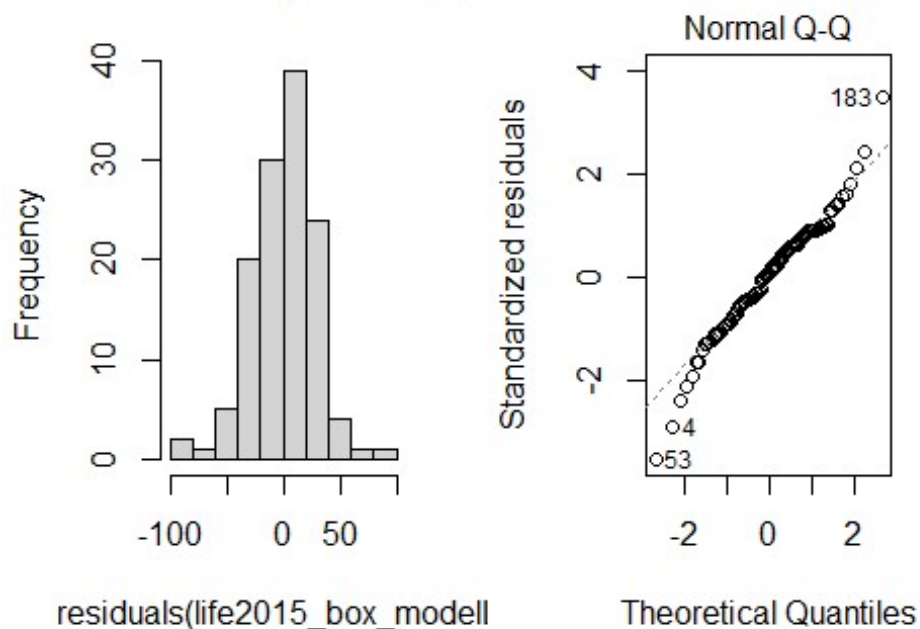
##
##  Shapiro-Wilk normality test
##
## data:  residuals(life2015_box_modelL)
## W = 0.97924, p-value = 0.04811

paste("Shapiro-Wilk normality test presented the p-value = 0.04811 < 0.05
that indicated that we should REJECT the null hypothesis and conclude that
the residuals are NOT normally distributed at 5 % level.")

## [1] "Shapiro-Wilk normality test presented the p-value = 0.04811 < 0.05
that indicated that we should REJECT the null hypothesis and conclude that
the residuals are NOT normally distributed at 5 % level."

#Plots
par(mfrow = c(1,2))
hist(residuals(life2015_box_modelL))
plot(life2015_box_modelL, which = 2)
```

am of residuals(life2015_b



```

paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")

## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."

# (L.6) Multicollinearity test
#VIF
imcdiag(life2015_box_modelL, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_box_modelL, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##                               VIF detection
## Income.composition.of.resources 1.6606      0
## Adult.Mortality                2.1035      0
## HIV.AIDS                       1.7151      0
## under.five.deaths              1.0571      0
##
## NOTE: VIF Method Failed to detect multicollinearity
##
##
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. ")

## [1] "Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. "

##### MODEL M - BOX COX TRANSFORMATION of MODEL G with under.five
back #####
paste("t-test of previous Box-Cox transformation indicated no significance of
interaction term after this transformation")

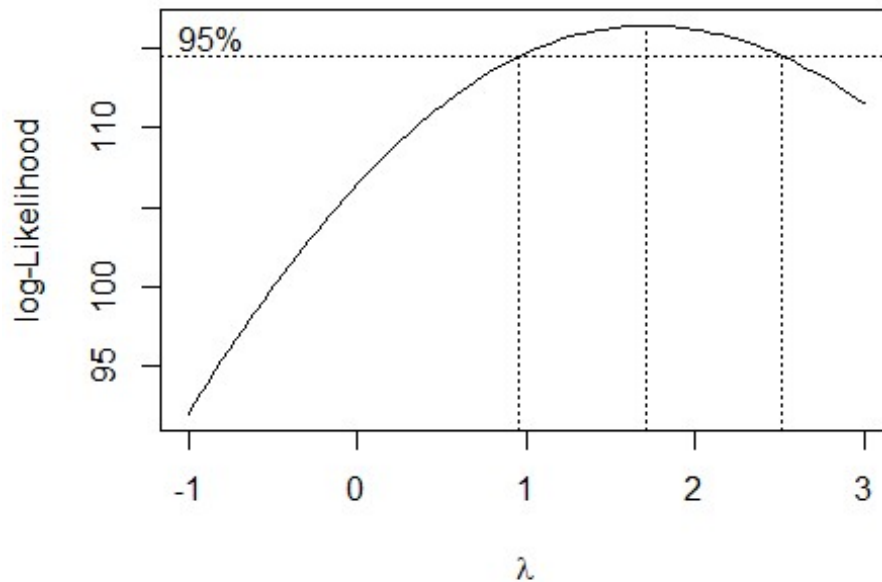
## [1] "t-test of previous Box-Cox transformation indicated no significance
of interaction term after this transformation"

#(M.1) BOX-COX TRANSFORMATION
# The necessity of transformation is due to the indication of
heterosdasticity in model E
# For this transformation we will be considering only the significant
parameters obtained beforehand
life2015_reduced_modelG1<- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
under.five.deaths + Adult.Mortality *)

```

```
HIV.AIDS, data = life2015)
```

```
bc = boxcox(life2015_reduced_modelG1, lambda = seq(-1,3))
```



```
#Extract best Lambda
```

```
bestlambda = bc$x[which(bc$y == max(bc$y))]
```

```
paste("The best lambda for Box-Cox Transformation is: ", bestlambda)
```

```
## [1] "The best lambda for Box-Cox Transformation is: 1.70707070707071"
```

```
paste("From the output, as the best lambda would be 1.70707.")
```

```
## [1] "From the output, as the best lambda would be 1.70707."
```

```
#BOX-COX with LAMBDA = 1.70707
```

```
life2015_box_modelM = lm((((Life.expectancy^1.70707) - 1)/1.70707) ~  
Income.composition.of.resources +  
Adult.Mortality + HIV.AIDS + under.five.deaths +  
Adult.Mortality * HIV.AIDS,  
data = life2015)
```

```
# (M.2) PARTIAL TEST - Individual Coefficients Test (t-test)
```

```
# Ho : Beta(i) = 0
```

```
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)
```

```
life2015_box_modelM = lm((((Life.expectancy^1.70707) - 1)/1.70707) ~
```

```

Income.composition.of.resources +
      Adult.Mortality + HIV.AIDS + under.five.deaths +
Adult.Mortality * HIV.AIDS,
      data = life2015[-c(119,53,93),])

summary(life2015_box_modelM)

##
## Call:
## lm(formula = (((Life.expectancy^1.70707) - 1)/1.70707) ~
Income.composition.of.resources +
##      Adult.Mortality + HIV.AIDS + under.five.deaths + Adult.Mortality *
##      HIV.AIDS, data = life2015[-c(119, 53, 93), ])
##
## Residuals:
##      Min        1Q    Median        3Q        Max
## -144.989  -29.971   -0.406    31.990   185.187
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      496.18857    36.47226   13.605 < 2e-16 ***
## Income.composition.of.resources  671.36142    41.39331   16.219 < 2e-16 ***
## Adult.Mortality      -0.53626     0.07570   -7.084 1.01e-10 ***
## HIV.AIDS            -49.92697     9.05512   -5.514 2.02e-07 ***
## under.five.deaths    -0.04094     0.03808   -1.075  0.285
## Adult.Mortality:HIV.AIDS      0.11330     0.02515    4.505 1.54e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 52.1 on 121 degrees of freedom
## Multiple R-squared:  0.8988, Adjusted R-squared:  0.8946
## F-statistic: 214.9 on 5 and 121 DF,  p-value: < 2.2e-16

paste("The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level.")

## [1] "The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level."

#confidence interval for the coefficients
confint(life2015_box_modelM, level = 0.95)

##              2.5 %          97.5 %
## (Intercept)      423.9821221  568.39502669
## Income.composition.of.resources  589.4124369  753.31039491
## Adult.Mortality    -0.6861274   -0.38638762
## HIV.AIDS          -67.8539694  -31.99997145
## under.five.deaths  -0.1163311    0.03445804
## Adult.Mortality:HIV.AIDS    0.0635141    0.16309373

```

```

paste("For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!")

## [1] "For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!"

# (M.3) STEPWISE REGRESSION PROCEDURE
# (G.3.1) BOTH
life2015_bothmodel = ols_step_both_p(life2015_box_modelM, pent = 0.05, prem =
0.05, details = TRUE)

## Stepwise Selection Method
## -----
##
## Candidate Terms:
##
## 1. Income.composition.of.resources
## 2. Adult.Mortality
## 3. HIV.AIDS
## 4. under.five.deaths
## 5. Adult.Mortality:HIV.AIDS
##
## We are selecting variables based on p value...
##
##
## Stepwise Selection: Step 1
##
## - Income.composition.of.resources added
##
##                               Model Summary
## -----
## R                               0.901          RMSE                69.927
## R-Squared                       0.812          Coef. Var          8.251
## Adj. R-Squared                  0.810          MSE                4889.727
## Pred R-Squared                  0.806          MAE                54.992
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF      Mean Square          F          Sig.
## -----
## Regression      2633488.698              1      2633488.698      538.576      0.0000
## Residual         611215.855             125          4889.727
## Total           3244704.553             126
## -----
##

```

```

##                                     Parameter Estimates
## -----
##                                     -----
##          Sig          lower      model      Beta      Std. Error      Std. Beta
##          t          lower      upper
## -----
##                                     (Intercept)      204.373      28.397
7.197      0.000      148.171      260.575
## Income.composition.of.resources      952.684      41.051      0.901
23.207      0.000      871.439      1033.930
## -----
##
##
##
## Stepwise Selection: Step 2
##
## - Adult.Mortality added
##
##                                     Model Summary
## -----
## R          0.934      RMSE          57.958
## R-Squared    0.872      Coef. Var    6.839
## Adj. R-Squared 0.870      MSE          3359.148
## Pred R-Squared 0.864      MAE          43.254
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of          DF      Mean Square          F          Sig.
##          Squares
## -----
## Regression    2828170.259          2      1414085.129      420.965      0.0000
## Residual      416534.295          124          3359.148
## Total         3244704.553          126
## -----
##
##                                     Parameter Estimates
## -----
##                                     -----
##          Sig          lower      model      Beta      Std. Error      Std. Beta
##          t          lower      upper
## -----
##                                     (Intercept)      417.598      36.585
11.414      0.000      345.186      490.010

```

```

## Income.composition.of.resources    754.502        42.841        0.713
17.611    0.000    669.707    839.297
##                               Adult.Mortality    -0.496        0.065    -0.308    -
7.613    0.000    -0.625    -0.367
## -----
##
##
##
##                               Model Summary
## -----
## R                0.934            RMSE                57.958
## R-Squared        0.872            Coef. Var            6.839
## Adj. R-Squared   0.870            MSE                3359.148
## Pred R-Squared   0.864            MAE                43.254
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2828170.259            2      1414085.129      420.965      0.0000
## Residual        416534.295           124        3359.148
## Total          3244704.553           126
## -----
##
##                               Parameter Estimates
## -----
##                               Sig      lower      model      Beta      Std. Error      Std. Beta
##                               t      upper
## -----
##                               (Intercept)      417.598        36.585
11.414    0.000    345.186    490.010
## Income.composition.of.resources    754.502        42.841        0.713
17.611    0.000    669.707    839.297
##                               Adult.Mortality    -0.496        0.065    -0.308    -
7.613    0.000    -0.625    -0.367
## -----
##
##
##
## Stepwise Selection: Step 3
##

```

- HIV.AIDS added

##

Model Summary

```
## -----
## R                0.938          RMSE                56.096
## R-Squared        0.881          Coef. Var            6.619
## Adj. R-Squared   0.878          MSE                 3146.805
## Pred R-Squared   0.866          MAE                 43.049
## -----
```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

```
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2857647.555      3      952549.185    302.704    0.0000
## Residual      387056.998     123      3146.805
## Total         3244704.553     126
## -----
```

##

##

Parameter Estimates

```
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)      423.584      35.464
11.944    0.000      353.386      493.783
## Income.composition.of.resources      735.124      41.946      0.695
17.526    0.000      652.095      818.153
## Adult.Mortality      -0.387      0.072      -0.240      -
5.337    0.000      -0.530      -0.243
## HIV.AIDS      -13.032      4.258      -0.125      -
3.061    0.003      -21.460      -4.604
## -----
```

##

##

##

##

Model Summary

```
## -----
## R                0.938          RMSE                56.096
## R-Squared        0.881          Coef. Var            6.619
## Adj. R-Squared   0.878          MSE                 3146.805
## Pred R-Squared   0.866          MAE                 43.049
## -----
```


##

ANOVA

ANOVA						
		Sum of Squares	DF	Mean Square	F	Sig.
	Regression	2857647.555	3	952549.185	302.704	0.0000
	Residual	387056.998	123	3146.805		
	Total	3244704.553	126			

##

##

```
##                                     Parameter Estimates
```

##			model	Beta	Std. Error	Std. Beta
t	Sig	lower	upper			
##						

##	(Intercept)	423.584	35.464
----	-------------	---------	--------

11.944	0.000	353.386	493.783
--------	-------	---------	---------

## Income.composition.of.resources	735.124	41.946	0.695
------------------------------------	---------	--------	-------

17.526	0.000	652.095	818.153
--------	-------	---------	---------

##	Adult.Mortality	-0.387	0.072	-0.240	-
----	-----------------	--------	-------	--------	---

5.337	0.000	-0.530	-0.243
-------	-------	--------	--------

##	HIV.AIDS	-13.032	4.258	-0.125	-

3.061	0.003	-21.460	-4.604
-------	-------	---------	--------

```

01.0001      01.0002      222.100      1.0001
## -----

```

##

##

##

```
## Stepwise Selection: Step 4
```

##

```
## - Adult.Mortality:HIV.AIDS added
```

##

```
##                                     Model Summary
```

## R	0.948	RMSE	52.134
------	-------	------	--------

## R-Squared	0.898	Coef. Var	6.152
--------------	-------	-----------	-------

## Adj. R-Squared	0.894	MSE	2717.959
-------------------	-------	-----	----------

## Pred R-Squared	0.874	MAE	40.229
-------------------	-------	-----	--------

```
## -----
```

```
## RMSE: Root Mean Square Error
```

MSE: Mean Square Error

```
## MAE: Mean Absolute Error
```

##

ANOVA

```

## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2913113.591      4      728278.398    267.951    0.0000
## Residual      331590.962     122      2717.959
## Total         3244704.553     126
## -----
##
##                                     Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t
## -----
##              (Intercept)      491.182      36.197
13.570    0.000    419.527    562.837
## Income.composition.of.resources      677.607      41.010      0.641
16.523    0.000    596.424    758.790
## Adult.Mortality      -0.542      0.076      -0.337    -
7.169    0.000    -0.691    -0.392
## HIV.AIDS      -49.853      9.061      -0.477    -
5.502    0.000    -67.789    -31.916
## Adult.Mortality:HIV.AIDS      0.114      0.025      0.415
4.517    0.000      0.064      0.163
## -----
##
##
##
##              Model Summary
## -----
## R      0.948      RMSE      52.134
## R-Squared      0.898      Coef. Var      6.152
## Adj. R-Squared      0.894      MSE      2717.959
## Pred R-Squared      0.874      MAE      40.229
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2913113.591      4      728278.398    267.951    0.0000
## Residual      331590.962     122      2717.959
## Total         3244704.553     126

```

Parameter Estimates						
	Sig.	lower	model upper	Beta	Std. Error	Std. Beta
(Intercept)				491.182	36.197	
Income.composition.of.resources				677.607	41.010	0.641
Adult.Mortality				-0.542	0.076	-0.337
HIV.AIDS				-49.853	9.061	-0.477
Adult.Mortality:HIV.AIDS				0.114	0.025	0.415
Model Summary						
R		0.948		RMSE		52.101
R-Squared		0.899		Coef. Var		6.148
Adj. R-Squared		0.895		MSE		2714.499
Pred R-Squared		0.872		MAE		40.072
ANOVA						
	Sum of Squares		DF	Mean Square	F	Sig.
Regression	2916250.184		5	583250.037	214.865	0.0000
Residual	328454.370		121	2714.499		
Total	3244704.553		126			
Parameter Estimates						

```

## -----
##
##          Sig          lower      model      Beta      Std. Error      Std. Beta
##          t          lower      upper
## -----
##
##          (Intercept)      496.189      36.472
## 13.605      0.000      423.982      568.395
## Income.composition.of.resources      671.361      41.393      0.635
## 16.219      0.000      589.412      753.310
## Adult.Mortality      -0.536      0.076      -0.334      -
## 7.084      0.000      -0.686      -0.386
## HIV.AIDS      -49.927      9.055      -0.478      -
## 5.514      0.000      -67.854      -32.000
## under.five.deaths      -0.041      0.038      -0.032      -
## 1.075      0.285      -0.116      0.034
## Adult.Mortality:HIV.AIDS      0.113      0.025      0.413
## 4.505      0.000      0.064      0.163
## -----
##
##
##
##
## Model Summary
## -----
## R      0.948      RMSE      52.101
## R-Squared      0.899      Coef. Var      6.148
## Adj. R-Squared      0.895      MSE      2714.499
## Pred R-Squared      0.872      MAE      40.072
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
##          Sum of
##          Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2916250.184      5      583250.037      214.865      0.0000
## Residual      328454.370      121      2714.499
## Total      3244704.553      126
## -----
##
##
## Parameter Estimates
## -----
##
##          Sig          lower      model      Beta      Std. Error      Std. Beta
##          t          lower      upper
## -----

```

```

-----
##              (Intercept)    496.189      36.472
13.605    0.000    423.982    568.395
## Income.composition.of.resources    671.361      41.393      0.635
16.219    0.000    589.412    753.310
##              Adult.Mortality    -0.536      0.076     -0.334    -
7.084    0.000     -0.686     -0.386
##              HIV.AIDS    -49.927      9.055     -0.478    -
5.514    0.000    -67.854    -32.000
##              under.five.deaths    -0.041      0.038     -0.032    -
1.075    0.285     -0.116      0.034
##              Adult.Mortality:HIV.AIDS    0.113      0.025      0.413
4.505    0.000      0.064      0.163
## -----
##
##
##
## Stepwise Selection: Step 6
##
## - Adult.Mortality:HIV.AIDS added
##
##
##              Model Summary
## -----
## R              0.939      RMSE              56.070
## R-Squared      0.882      Coef. Var          6.616
## Adj. R-Squared 0.878      MSE              3143.861
## Pred R-Squared 0.864      MAE              42.950
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2861153.535      4      715288.384    227.519    0.0000
## Residual      383551.019     122      3143.861
## Total         3244704.553     126
## -----
##
##
##              Parameter Estimates
## -----
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t
## -----

```

```

##              (Intercept)    429.108        35.831
11.976    0.000    358.177    500.039
## Income.composition.of.resources    728.324        42.418        0.689
17.170    0.000    644.354    812.295
##              Adult.Mortality    -0.381        0.073    -0.237    -
5.254    0.000    -0.525    -0.238
##              HIV.AIDS    -13.237        4.260    -0.127    -
3.107    0.002    -21.670    -4.803
##              under.five.deaths    -0.043        0.041    -0.034    -
1.056    0.293    -0.124    0.038
## -----
##
##
##
##
## Final Model Output
## -----
##
##              Model Summary
## -----
## R                0.939        RMSE                56.070
## R-Squared        0.882        Coef. Var            6.616
## Adj. R-Squared   0.878        MSE                3143.861
## Pred R-Squared   0.864        MAE                42.950
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares        DF        Mean Square        F        Sig.
## -----
## Regression      2861153.535         4        715288.384        227.519        0.0000
## Residual        383551.019        122         3143.861
## Total          3244704.553        126
## -----
##
##              Parameter Estimates
## -----
##
##              Sig        lower        model        Beta        Std. Error        Std. Beta
##              t              upper
## -----
##
##              (Intercept)    429.108        35.831
11.976    0.000    358.177    500.039
## Income.composition.of.resources    728.324        42.418        0.689

```

```

17.170    0.000    644.354    812.295
##          Adult.Mortality    -0.381          0.073    -0.237    -
5.254    0.000    -0.525    -0.238
##          HIV.AIDS    -13.237          4.260    -0.127    -
3.107    0.002    -21.670    -4.803
##          under.five.deaths    -0.043          0.041    -0.034    -
1.056    0.293    -0.124    0.038
## -----

```

```
life2015_bothmodel
```

```

##
##                                     Stepwise Selection Summary
## -----
##                                     Added/      Adj.
## Step      Variable      Removed      R-Square      R-
## Square      C(p)      AIC      RMSE
## -----
## 1      Income.composition.of.resources      addition      0.812
0.810      102.1670      1443.2457      69.9266
## 2      Adult.Mortality      addition      0.872
0.870      32.4480      1396.5436      57.9582
## 3      HIV.AIDS      addition      0.881
0.878      23.5890      1389.2222      56.0964
## 4      Adult.Mortality:HIV.AIDS      addition      0.898
0.894      5.1550      1371.5791      52.1340
## 5      under.five.deaths      addition      0.899
0.895      6.0000      1372.3721      52.1009
## 6      Adult.Mortality:HIV.AIDS      removal      0.882
0.878      24.2970      1390.0666      56.0701
## -----

```

```
#life2015_bothmodel$mallows_cp
```

```
paste("Stepwise procedure with BOTH indicated to consider all the variables
and the interaction term.")
```

```
## [1] "Stepwise procedure with BOTH indicated to consider all the variables
and the interaction term."
```

```
# (G.3.2) FORWARD
```

```
life2015_forwardmodel = ols_step_forward_p(life2015_box_modelM, pent = 0.05,
details = TRUE)
```

```
## Forward Selection Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```

##
## 1. Income.composition.of.resources
## 2. Adult.Mortality
## 3. HIV.AIDS
## 4. under.five.deaths
## 5. Adult.Mortality:HIV.AIDS
##
## We are selecting variables based on p value...
##
##
## Forward Selection: Step 1
##
## - Income.composition.of.resources
##
##                               Model Summary
## -----
## R                               0.901          RMSE                69.927
## R-Squared                       0.812          Coef. Var           8.251
## Adj. R-Squared                   0.810          MSE                4889.727
## Pred R-Squared                   0.806          MAE                54.992
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF      Mean Square          F          Sig.
## -----
## Regression      2633488.698              1      2633488.698      538.576      0.0000
## Residual         611215.855             125           4889.727
## Total           3244704.553             126
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model          Beta      Std. Error      Std. Beta
## t          Sig          lower          upper
## -----
##                               (Intercept)      204.373          28.397
## 7.197      0.000      148.171      260.575
## Income.composition.of.resources      952.684          41.051          0.901
## 23.207      0.000      871.439      1033.930
## -----
##
##
##

```



```

##
## Forward Selection: Step 2
##
## - Adult.Mortality
##
##                               Model Summary
## -----
## R                               0.934          RMSE                57.958
## R-Squared                       0.872          Coef. Var          6.839
## Adj. R-Squared                  0.870          MSE                3359.148
## Pred R-Squared                  0.864          MAE                43.254
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF      Mean Square      F          Sig.
## -----
## Regression      2828170.259          2      1414085.129      420.965      0.0000
## Residual        416534.295          124      3359.148
## Total          3244704.553          126
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               t          Sig          lower          model          Beta      Std. Error      Std. Beta
##                               upper
## -----
##                               (Intercept)      417.598          36.585
## 11.414      0.000      345.186      490.010
## ## Income.composition.of.resources      754.502          42.841          0.713
## 17.611      0.000      669.707      839.297
## ## Adult.Mortality      -0.496          0.065          -0.308
## 7.613      0.000      -0.625      -0.367
## -----
##
##
##
## Forward Selection: Step 3
##
## - HIV.AIDS
##
##                               Model Summary
## -----

```

```

## R                0.938      RMSE                56.096
## R-Squared        0.881      Coef. Var            6.619
## Adj. R-Squared   0.878      MSE                3146.805
## Pred R-Squared   0.866      MAE                43.049
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                Sum of
##                Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2857647.555      3      952549.185      302.704      0.0000
## Residual        387056.998     123        3146.805
## Total          3244704.553     126
## -----
##
##                               Parameter Estimates
## -----
## -----
##                Sig      lower      model      Beta      Std. Error      Std. Beta
##                t                upper
## -----
## -----
##                (Intercept)      423.584      35.464
## 11.944      0.000      353.386      493.783
## Income.composition.of.resources      735.124      41.946      0.695
## 17.526      0.000      652.095      818.153
## Adult.Mortality      -0.387      0.072      -0.240      -
## 5.337      0.000      -0.530      -0.243
## HIV.AIDS      -13.032      4.258      -0.125      -
## 3.061      0.003      -21.460      -4.604
## -----
## -----
##
##
##
## Forward Selection: Step 4
##
## - Adult.Mortality:HIV.AIDS
##
##                               Model Summary
## -----
## R                0.948      RMSE                52.134
## R-Squared        0.898      Coef. Var            6.152
## Adj. R-Squared   0.894      MSE                2717.959
## Pred R-Squared   0.874      MAE                40.229
## -----

```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

	Sum of Squares	DF	Mean Square	F	Sig.
## Regression	2913113.591	4	728278.398	267.951	0.0000
## Residual	331590.962	122	2717.959		
## Total	3244704.553	126			

##

##

Parameter Estimates

	Sig	lower	model upper	Beta	Std. Error	Std. Beta
## (Intercept)				491.182	36.197	
13.570	0.000	419.527	562.837			
## Income.composition.of.resources				677.607	41.010	0.641
16.523	0.000	596.424	758.790			
## Adult.Mortality				-0.542	0.076	-0.337
7.169	0.000	-0.691	-0.392			
## HIV.AIDS				-49.853	9.061	-0.477
5.502	0.000	-67.789	-31.916			
## Adult.Mortality:HIV.AIDS				0.114	0.025	0.415
4.517	0.000	0.064	0.163			

##

##

##

##

##

Forward Selection: Step 5

##

- under.five.deaths

##

##

##

##

##

##

##

##

##

##

##

##

##

##

##

##

##

##

```

##
##
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2916250.184      5      583250.037    214.865    0.0000
## Residual      328454.370     121      2714.499
## Total         3244704.553     126
## -----
##
##              Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)    496.189      36.472
## 13.605    0.000    423.982    568.395
## Income.composition.of.resources    671.361      41.393      0.635
## 16.219    0.000    589.412    753.310
## Adult.Mortality    -0.536      0.076    -0.334    -
## 7.084    0.000    -0.686    -0.386
## HIV.AIDS    -49.927      9.055    -0.478    -
## 5.514    0.000    -67.854    -32.000
## under.five.deaths    -0.041      0.038    -0.032    -
## 1.075    0.285    -0.116      0.034
## Adult.Mortality:HIV.AIDS    0.113      0.025      0.413
## 4.505    0.000      0.064      0.163
## -----
##
##
##
##
## Variables Entered:
##
## + Income.composition.of.resources
## + Adult.Mortality
## + HIV.AIDS
## + Adult.Mortality:HIV.AIDS
## + under.five.deaths
##
##
## Final Model Output
## -----
##
##              Model Summary
## -----

```

```

## R                0.948      RMSE                52.101
## R-Squared        0.899      Coef. Var            6.148
## Adj. R-Squared   0.895      MSE                2714.499
## Pred R-Squared   0.872      MAE                40.072
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                Sum of
##                Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2916250.184      5      583250.037      214.865      0.0000
## Residual        328454.370     121        2714.499
## Total          3244704.553     126
## -----
##
##                               Parameter Estimates
## -----
##
##                               t      Sig      lower      model      upper      Beta      Std. Error      Std. Beta
##                               -----
##                               (Intercept)      496.189      36.472
## 13.605      0.000      423.982      568.395
## ## Income.composition.of.resources      671.361      41.393      0.635
## 16.219      0.000      589.412      753.310
## ## Adult.Mortality      -0.536      0.076      -0.334      -
## 7.084      0.000      -0.686      -0.386
## ## HIV.AIDS      -49.927      9.055      -0.478      -
## 5.514      0.000      -67.854      -32.000
## ## under.five.deaths      -0.041      0.038      -0.032      -
## 1.075      0.285      -0.116      0.034
## ## Adult.Mortality:HIV.AIDS      0.113      0.025      0.413
## 4.505      0.000      0.064      0.163
## -----
## -----
life2015_forwardmodel

##
##                               Selection Summary
## -----
## -----
##                Variable
## Step      Entered      R-Square      Adj.      C(p)
AIC      RMSE

```

```

## -----
##      1      Income.composition.of.resources      0.8116      0.8101
102.1671      1443.2457      69.9266
##      2      Adult.Mortality      0.8716      0.8696
32.4479      1396.5436      57.9582
##      3      HIV.AIDS      0.8807      0.8778
23.5887      1389.2222      56.0964
##      4      Adult.Mortality:HIV.AIDS      0.8978      0.8945
5.1555      1371.5791      52.1340
##      5      under.five.deaths      0.8988      0.8946
6.0000      1372.3721      52.1009
## -----

#life2015_forwardmodel$mallows_cp
paste("Stepwise procedure with FORWARD indicated to consider all the
variables and the interaction term.")

## [1] "Stepwise procedure with FORWARD indicated to consider all the
variables and the interaction term."

# (G.3.3) BACKWARD
life2015_backwardmodel = ols_step_backward_p(life2015_box_modelM, prem =
0.05, details = TRUE)

## Backward Elimination Method
## -----
##
## Candidate Terms:
##
## 1 . Income.composition.of.resources
## 2 . Adult.Mortality
## 3 . HIV.AIDS
## 4 . under.five.deaths
## 5 . Adult.Mortality:HIV.AIDS
##
## We are eliminating variables based on p value...
##
## - under.five.deaths
##
## Backward Elimination: Step 1
##
## Variable under.five.deaths Removed
##
##
## Model Summary
## -----
## R      0.948      RMSE      52.134
## R-Squared      0.898      Coef. Var      6.152
## Adj. R-Squared      0.894      MSE      2717.959
## Pred R-Squared      0.874      MAE      40.229

```

```
##
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of
## Squares DF Mean Square F Sig.
## -----
## Regression 2913113.591 4 728278.398 267.951 0.0000
## Residual 331590.962 122 2717.959
## Total 3244704.553 126
## -----
##
## Parameter Estimates
## -----
##
## Sig. lower model upper Beta Std. Error Std. Beta
## t Sig. lower upper
## -----
## (Intercept) 491.182 36.197
13.570 0.000 419.527 562.837
## Income.composition.of.resources 677.607 41.010 0.641
16.523 0.000 596.424 758.790
## Adult.Mortality -0.542 0.076 -0.337
7.169 0.000 -0.691 -0.392
## HIV.AIDS -49.853 9.061 -0.477
5.502 0.000 -67.789 -31.916
## Adult.Mortality:HIV.AIDS 0.114 0.025 0.415
4.517 0.000 0.064 0.163
## -----
##
##
##
## No more variables satisfy the condition of p value = 0.05
##
##
## Variables Removed:
##
## - under.five.deaths
##
##
## Final Model Output
## -----
##
## Model Summary
## -----
```

```

## R                0.948      RMSE                52.134
## R-Squared        0.898      Coef. Var            6.152
## Adj. R-Squared   0.894      MSE                2717.959
## Pred R-Squared   0.874      MAE                40.229
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                Sum of
##                Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2913113.591      4      728278.398      267.951      0.0000
## Residual        331590.962     122      2717.959
## Total          3244704.553     126
## -----
##
##                               Parameter Estimates
## -----
##
##                               t      Sig      lower      model      upper      Beta      Std. Error      Std. Beta
##                               -----
##                               (Intercept)      491.182      36.197
## 13.570      0.000      419.527      562.837
## ## Income.composition.of.resources      677.607      41.010      0.641
## 16.523      0.000      596.424      758.790
## ## Adult.Mortality      -0.542      0.076      -0.337      -
## 7.169      0.000      -0.691      -0.392
## ## HIV.AIDS      -49.853      9.061      -0.477      -
## 5.502      0.000      -67.789      -31.916
## ## Adult.Mortality:HIV.AIDS      0.114      0.025      0.415
## 4.517      0.000      0.064      0.163
## -----
## -----
life2015_backwardmodel
##
##
##                               Elimination Summary
## -----
## -----
##                Variable      Adj.
##                Removed      R-Square      R-Square      C(p)      AIC
## RMSE
## -----

```



```

-----
##      1      under.five.deaths      0.8978      0.8945      5.1555      1371.5791
52.1340
## -----
-----

#life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD indicated to remove under.five.deaths
variable.")

## [1] "Stepwise procedure with BACKWARD indicated to remove
under.five.deaths variable."

# (M.4) BEST SUBSET
#OPTION 1
ks = ols_step_best_subset(life2015_box_modelM, details = TRUE)

#Summary of Selected model based on cp, aic, AdjustedR2
rsquare <- c(ks$rsq)
cp <- c(ks$cp)
aic <- c(ks$aic)
AdjustedR2 <- c(ks$adjr)
cbind(rsquare, cp, aic, AdjustedR2)

##           rsquare           cp           aic AdjustedR2
## [1,] 0.8116267 102.167102 1443.246 0.8101197
## [2,] 0.8716264 32.447950 1396.544 0.8695559
## [3,] 0.8807112 23.588746 1389.222 0.8778017
## [4,] 0.8978055 5.155496 1371.579 0.8944549
## [5,] 0.8987722 6.000000 1372.372 0.8945892

#OPTION 2
best.subset <- regsubsets((((Life.expectancy^1.70707) - 1)/1.70707) ~
Income.composition.of.resources +
                        Adult.Mortality + HIV.AIDS + under.five.deaths +
Adult.Mortality * HIV.AIDS,
                        data = life2015[-c(119, 53, 93), ], nv = 4)
summary(best.subset)

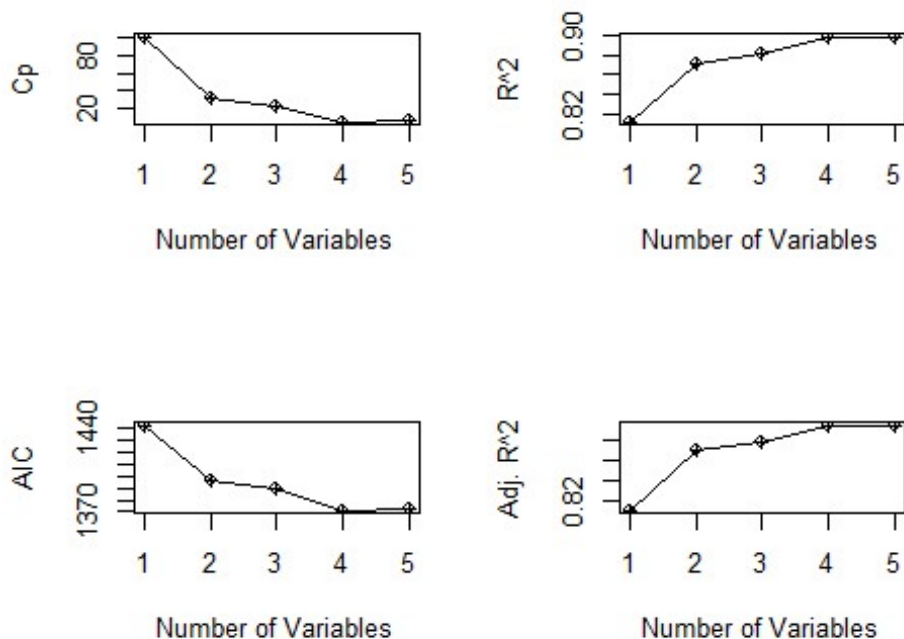
## Subset selection object
## Call: regsubsets.formula((((Life.expectancy^1.70707) - 1)/1.70707) ~
##      Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
##      under.five.deaths + Adult.Mortality * HIV.AIDS, data = life2015[-
c(119,
##      53, 93), ], nv = 4)
## 5 Variables (and intercept)
##
##           Forced in Forced out
## Income.composition.of.resources      FALSE      FALSE
## Adult.Mortality                      FALSE      FALSE
## HIV.AIDS                          FALSE      FALSE
## under.five.deaths                   FALSE      FALSE

```

```
## Adult.Mortality:HIV.AIDS          FALSE      FALSE
## 1 subsets of each size up to 4
## Selection Algorithm: exhaustive
##           Income.composition.of.resources Adult.Mortality HIV.AIDS
## 1  ( 1 ) "*"                        " "      " "
## 2  ( 1 ) "*"                        "*"      " "
## 3  ( 1 ) "*"                        "*"      "*"
## 4  ( 1 ) "*"                        "*"      "*"
##           under.five.deaths Adult.Mortality:HIV.AIDS
## 1  ( 1 ) " "                      " "
## 2  ( 1 ) " "                      " "
## 3  ( 1 ) " "                      " "
## 4  ( 1 ) " "                      "*"

reg.summary <- summary(best.subset)

#Plotting
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab = "R^2")
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab = "AIC")
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Adj. R^2")
```



```
paste("Based on Marlow's Cp the minimum value was obtained without  
under.five.deaths, but the model with this variable presented a slightly high  
value.")
```

```
## [1] "Based on Marlow's Cp the minimum value was obtained without  
under.five.deaths, but the model with this variable presented a slightly high  
value."
```

```
#MODEL DIAGNOSTICS
```

```
# (M.5) CHECKING LINEARITY
```

```
#Residuals plot
```

```
plot(life2015_box_modelM, which = 1)
```

```
# a Scale Location Plot
```

```
plot(life2015_box_modelM, which = 3)
```

```
paste("It is verified some slightly change in the pattern for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts  
values in the last plot, but in overall the average of residuals almost  
horizontal and we can say that the residuals are equally spread that we can  
conclude that the linearity assumption is respected."
```

```
# (M.6) Heteroscedasticity Test - the Breush-Pagan test
```

```
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
```

```
# Ha: heteroscedasticity is presented
```

```
bptest(life2015_box_modelM)
```

```
##
```

```
## studentized Breusch-Pagan test
```

```
##
```

```
## data: life2015_box_modelM
```

```
## BP = 9.6388, df = 5, p-value = 0.08614
```

```
paste("The output displays the Breush-Pagan test that from the model  
presented the p-value 0.08614 > 0.05, indicating the we should NOT REJECT the  
null hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity does not exist.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model  
presented the p-value 0.08614 > 0.05, indicating the we should NOT REJECT the  
null hypthesis at 5 % level and consequently the test provides evidence that  
heteroscedasticity does not exist."
```

```
# (M.7) Testing for normality
```

```
#NORMALITY - Shapiro-Wilk Test
```

```
# Ho : the sample data is significantly normally distributed
```

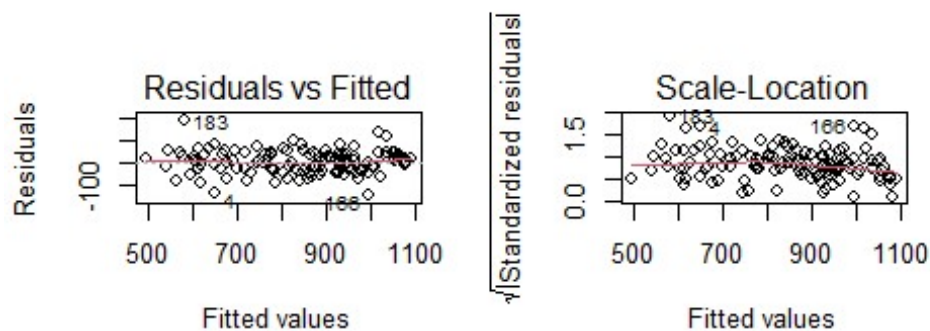
```
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_modelM))

##
## Shapiro-Wilk normality test
##
## data: residuals(life2015_box_modelM)
## W = 0.98406, p-value = 0.1425

paste("Shapiro-Wilk normality test presented the p-value = 0.1425 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level.")

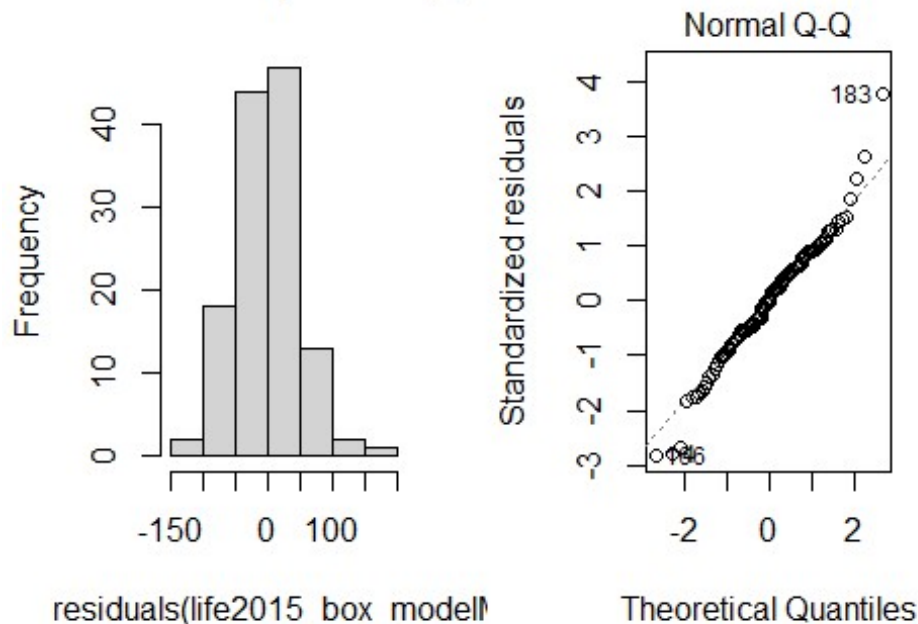
## [1] "Shapiro-Wilk normality test presented the p-value = 0.1425 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level."

#Plots
par(mfrow = c(1,2))
```



```
hist(residuals(life2015_box_modelM))
plot(life2015_box_modelM, which = 2)
```

am of residuals(life2015_b



```
paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")
```

```
## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."
```

```
# (M.8) Multicollinearity test
```

```
#VIF
```

```
imcdiag(life2015_box_modelM, method = "VIF")
```

```
##
```

```
## Call:
```

```
## imcdiag(mod = life2015_box_modelM, method = "VIF")
```

```
##
```

```
##
```

```
## VIF Multicollinearity Diagnostics
```

```
##
```

```
##
```

	VIF	detection
Income.composition.of.resources	1.8315	0
Adult.Mortality	2.6515	0
HIV.AIDS	8.9737	0
under.five.deaths	1.0573	0
Adult.Mortality:HIV.AIDS	10.0621	1

```
##
```

```
## Multicollinearity may be due to Adult.Mortality:HIV.AIDS regressors
```

```
##
```

```
## 1 --> COLLINEARITY is detected by the test
```

```
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. ")

## [1] "Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. "

# (M.9) LEVERAGE OF THE POINTS
# Leverage Points
lev = hatvalues(life2015_box_modelM)
p = length(coef(life2015_box_modelM))
n = nrow(life2015)
outlier = lev[lev > (2 * p/n)]
print(outlier)

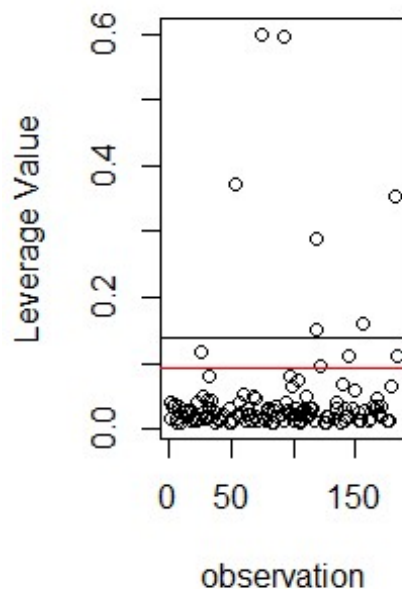
##          26          53          75          93          118          119
122
## 0.11862829 0.37316436 0.59996323 0.59586815 0.15169141 0.28799464
0.09702407
##          144          156          182          183
## 0.11143226 0.16085667 0.35257285 0.11138531

plot(rownames(life2015[-c(119, 53, 93), ]), lev, main = "Leverage in Life2015
Dataset",
      xlab = "observation", ylab = "Leverage Value")
abline(h = 2 * p/n, lty = 1, col = "red")
abline(h = 3 * p/n, lty = 1)

paste("The results shows that that the data points #26, #53, #75, #93, #118,
#119, #122, #144, #156, #152, #156 and #182 presented leverage values higher
than 2.p/n that might be outliers.")

## [1] "The results shows that that the data points #26, #53, #75, #93, #118,
#119, #122, #144, #156, #152, #156 and #182 presented leverage values higher
than 2.p/n that might be outliers."
```

Leverage in Life2015 Data



```
##### MODEL N - BOX COX TRANSFORMATION of MODEL G with infant.deaths  
back #####  
paste("t-test of previous Box-Cox transformation indicated no significance of  
interaction term after this transformation")  
  
## [1] "t-test of previous Box-Cox transformation indicated no significance  
of interaction term after this transformation"  
  
#(N.1) BOX-COX TRANSFORMATION  
# The necessity of transformation is due to the indication of  
heterosdasticity in model E  
# For this transformation we will be considering only the significant  
parameters obtained beforehand  
life2015_reduced_modelG2<- lm(Life.expectancy ~  
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +  
infant.deaths + Adult.Mortality * HIV.AIDS,  
data = life2015)  
  
bc = boxcox(life2015_reduced_modelG2, lambda = seq(-1,3))
```



```
summary(life2015_box_modelN)

##
## Call:
## lm(formula = (((Life.expectancy^1.74747) - 1)/1.74747) ~
Income.composition.of.resources +
##     Adult.Mortality + HIV.AIDS + infant.deaths + Adult.Mortality *
##     HIV.AIDS, data = life2015[-c(119, 53, 93), ])
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -172.553  -36.040   -0.149   37.963  219.337
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      565.41167    43.34029   13.046 < 2e-16 ***
## Income.composition.of.resources 799.09051    49.16044   16.255 < 2e-16 ***
## Adult.Mortality      -0.63794     0.08998   -7.090 9.74e-11 ***
## HIV.AIDS            -59.32336    10.76804   -5.509 2.06e-07 ***
## infant.deaths      -0.05282     0.05791   -0.912  0.364
## Adult.Mortality:HIV.AIDS      0.13500     0.02990    4.515 1.48e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 61.95 on 121 degrees of freedom
## Multiple R-squared:  0.8984, Adjusted R-squared:  0.8942
## F-statistic: 214 on 5 and 121 DF, p-value: < 2.2e-16

paste("The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level.")

## [1] "The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level."

#confidence interval for the coefficients
confint(life2015_box_modelN, level = 0.95)

##              2.5 %       97.5 %
## (Intercept)  479.60813253 651.21521678
## Income.composition.of.resources 701.76445020 896.41657664
## Adult.Mortality   -0.81606645 -0.45980664
## HIV.AIDS         -80.64152260 -38.00518819
## infant.deaths    -0.16745789  0.06182037
## Adult.Mortality:HIV.AIDS  0.07580645  0.19419946

paste("For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!")
```

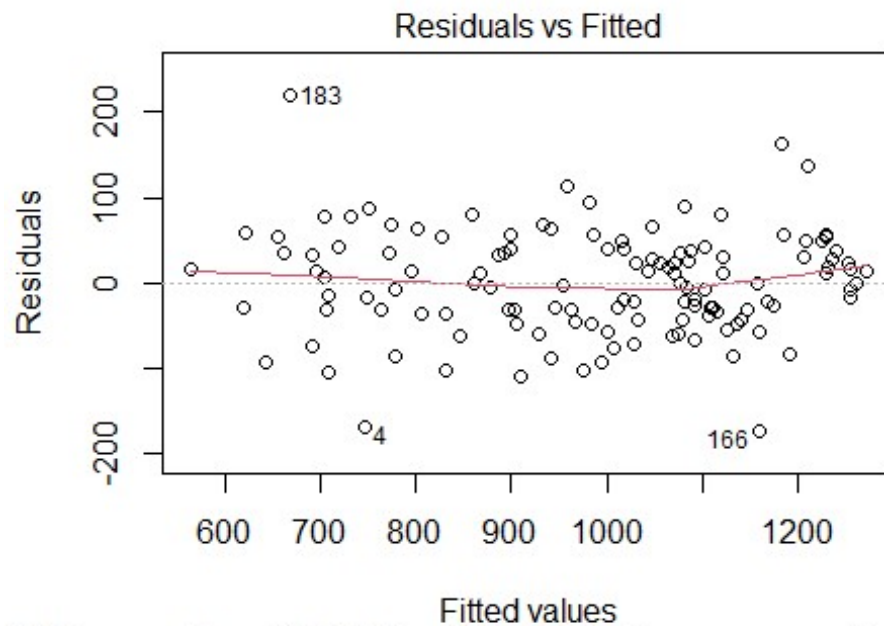
```
## [1] "For a 95% confidence interval, under.five.deaths with Zero between  
upper and lower values!"
```

```
#MODEL DIAGNOSTICS
```

```
# (N.3) CHECKING LINEARITY
```

```
#Residuals plot
```

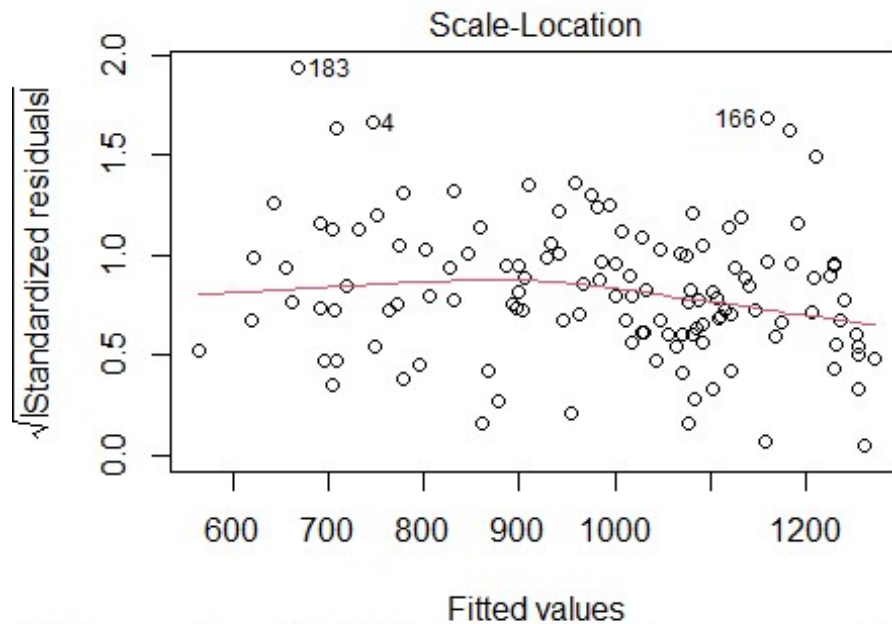
```
plot(life2015_box_modelN, which = 1)
```



```
l((((Life.expectancy^1.74747) - 1)/1.74747) ~ Income.composition.of.re
```

```
# a Scale Location Plot
```

```
plot(life2015_box_modelN, which = 3)
```



`(((Life.expectancy^1.74747) - 1)/1.74747) ~ Income.composition.of.re`

```
paste("It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected."
```

```
# (N.4) Heteroscedasticity Test - the Breush-Pagan test
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
# Ha: heteroscedasticity is presented
bptest(life2015_box_modelN)
```

```
##
## studentized Breusch-Pagan test
##
## data: life2015_box_modelN
## BP = 9.689, df = 5, p-value = 0.08454
```

```
paste("The output displays the Breush-Pagan test that from the model
presented the p-value 0.08454 > 0.05, indicating the we should NOT REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity does not exist.")
```

```
## [1] "The output displays the Breush-Pagan test that from the model
presented the p-value 0.08454 > 0.05, indicating the we should NOT REJECT the
```

null hypothesis at 5 % level and consequently the test provides evidence that heteroscedasticity does not exist."

```
# (N.5) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_modelN))

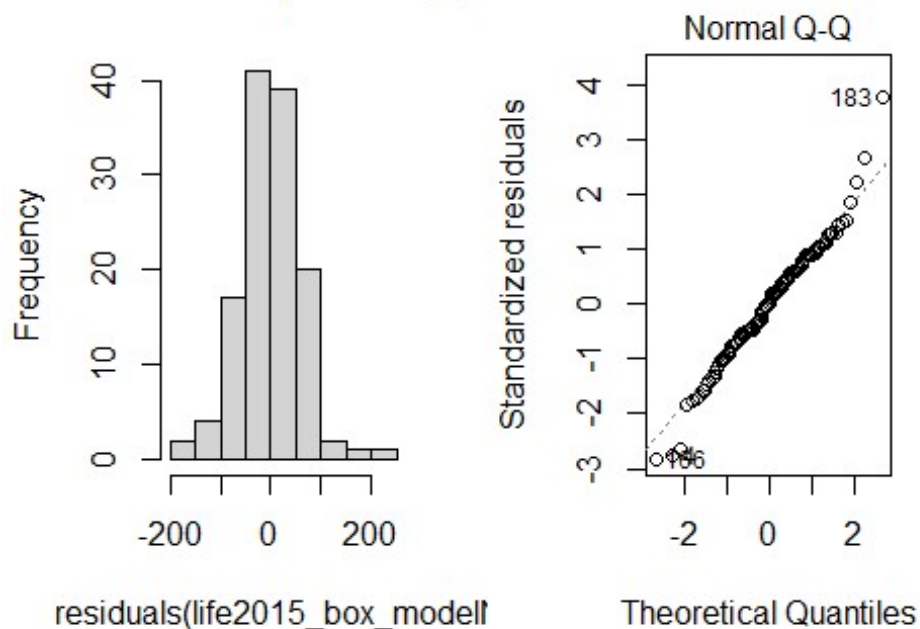
##
##  Shapiro-Wilk normality test
##
## data:  residuals(life2015_box_modelN)
## W = 0.98434, p-value = 0.1517

paste("Shapiro-Wilk normality test presented the p-value = 0.1517 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level.")

## [1] "Shapiro-Wilk normality test presented the p-value = 0.1517 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level."

#Plots
par(mfrow = c(1,2))
hist(residuals(life2015_box_modelN))
plot(life2015_box_modelN, which = 2)
```

am of residuals(life2015_b



```

paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")

## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."

# (N.6) Multicollinearity test
#VIF
imcdiag(life2015_box_modelN, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_box_modelN, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##                               VIF detection
## Income.composition.of.resources 1.8273      0
## Adult.Mortality                 2.6496      0
## HIV.AIDS                        8.9762      0
## infant.deaths                   1.0465      0
## Adult.Mortality:HIV.AIDS        10.0609     1
##
## Multicollinearity may be due to Adult.Mortality:HIV.AIDS regressors
##
## 1 --> COLLINEARITY is detected by the test
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. ")

## [1] "Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. "

##### MODEL P - BOX COX TRANSFORMATION of MODEL G with under.five and
Hepatitis.B #####
paste("t-test of previous Box-Cox transformation indicated no significance of
interaction term after this transformation")

## [1] "t-test of previous Box-Cox transformation indicated no significance
of interaction term after this transformation"

#(P.1) BOX-COX TRANSFORMATION
# The necessity of transformation is due to the indication of
heterosdasticity in model E
# For this transformation we will be considering only the significant
parameters obtained beforehand
life2015_reduced_modelP<- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +

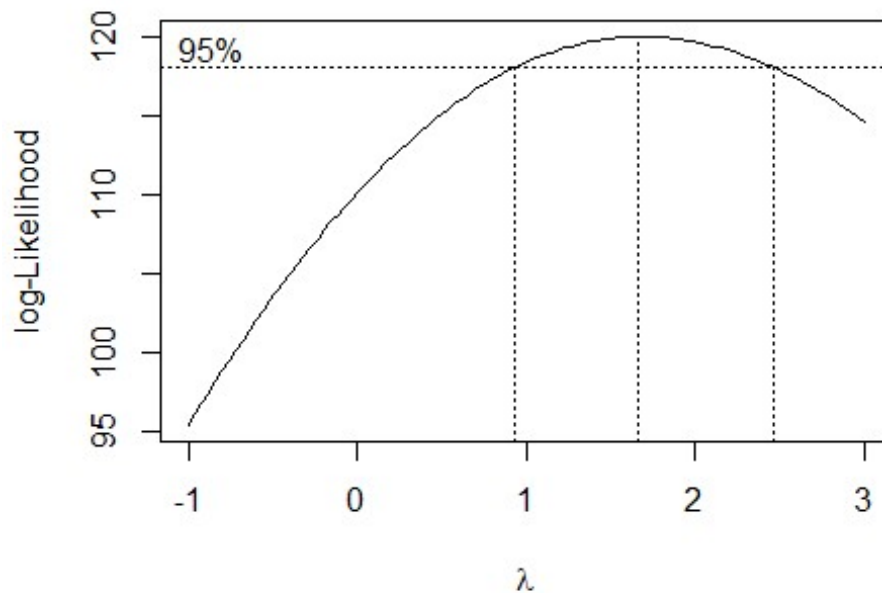
```

```

                                under.five.deaths + Hepatitis.B +
Adult.Mortality * HIV.AIDS, data = life2015)

bc = boxcox(life2015_reduced_modelP, lambda = seq(-1,3))

```



```

#Extract best Lambda
bestlambda = bc$x[which(bc$y == max(bc$y))]
paste("The best lambda for Box-Cox Transformation is: ", bestlambda)

## [1] "The best lambda for Box-Cox Transformation is:  1.66666666666667"

paste("From the output, as the best lambda would be 1.66667.")

## [1] "From the output, as the best lambda would be 1.66667."

#BOX-COX with LAMBDA = 1.70707
life2015_box_modelP = lm((((Life.expectancy^1.66667) - 1)/1.66667) ~
Income.composition.of.resources +
                                Adult.Mortality + HIV.AIDS + under.five.deaths +
Hepatitis.B +
                                Adult.Mortality * HIV.AIDS,
                                data = life2015)

# (P.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

```

```

life2015_box_modelP = lm((((Life.expectancy^1.66667) - 1)/1.66667) ~
Income.composition.of.resources +
                        Adult.Mortality + HIV.AIDS + under.five.deaths +
Hepatitis.B +
                        Adult.Mortality * HIV.AIDS, data = life2015[-
c(119,53,93),])

summary(life2015_box_modelP)

##
## Call:
## lm(formula = (((Life.expectancy^1.66667) - 1)/1.66667) ~
Income.composition.of.resources +
##      Adult.Mortality + HIV.AIDS + under.five.deaths + Hepatitis.B +
##      Adult.Mortality * HIV.AIDS, data = life2015[-c(119, 53, 93),
##      ])
##
## Residuals:
##      Min        1Q    Median        3Q        Max
## -116.902  -25.632    2.449    27.219   141.008
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      401.02873    33.47769   11.979 < 2e-16 ***
## Income.composition.of.resources 560.33975    34.30497   16.334 < 2e-16 ***
## Adult.Mortality      -0.45129     0.06263   -7.206 5.54e-11 ***
## HIV.AIDS            -34.60606     8.15281   -4.245 4.34e-05 ***
## under.five.deaths    -0.03015     0.03156   -0.955 0.341398
## Hepatitis.B           0.41898     0.18023    2.325 0.021769 *
## Adult.Mortality:HIV.AIDS    0.08023     0.02178    3.683 0.000347 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 43.1 on 120 degrees of freedom
## Multiple R-squared:  0.9032, Adjusted R-squared:  0.8984
## F-statistic: 186.7 on 6 and 120 DF,  p-value: < 2.2e-16

paste("The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level.")

## [1] "The t-test indicated that after the Box-Cox transformation the
under.five.deaths with p-value > 0.05 that indicates that this variable does
not have significant influence in Life Expectancy at 5 % level."

#confidence interval for the coefficients
confint(life2015_box_modelP, level = 0.95)

##              2.5 %        97.5 %
## (Intercept)  334.74523791 467.31221303

```

```

## Income.composition.of.resources 492.41829099 628.26121039
## Adult.Mortality -0.57528254 -0.32729041
## HIV.AIDS -50.74805455 -18.46407368
## under.five.deaths -0.09264135 0.03234359
## Hepatitis.B 0.06213597 0.77581510
## Adult.Mortality:HIV.AIDS 0.03709493 0.12335954

paste("For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!")

## [1] "For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!"

# (P.3) STEPWISE REGRESSION PROCEDURE
# (P.3.1) BOTH
life2015_bothmodel = ols_step_both_p(life2015_box_modelP, pent = 0.05, prem =
0.05, details = TRUE)

## Stepwise Selection Method
## -----
##
## Candidate Terms:
##
## 1. Income.composition.of.resources
## 2. Adult.Mortality
## 3. HIV.AIDS
## 4. under.five.deaths
## 5. Hepatitis.B
## 6. Adult.Mortality:HIV.AIDS
##
## We are selecting variables based on p value...
##
## Stepwise Selection: Step 1
##
## - Income.composition.of.resources added
##
##
## Model Summary
## -----
## R 0.901 RMSE 58.964
## R-Squared 0.811 Coef. Var 8.074
## Adj. R-Squared 0.810 MSE 3476.746
## Pred R-Squared 0.806 MAE 46.343
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of

```


##		Squares	DF	Mean Square	F	Sig.
##						
##	Regression	1869033.599	1	1869033.599	537.581	0.0000
##	Residual	434593.203	125	3476.746		
##	Total	2303626.802	126			

```
## -----
##
##                               Parameter Estimates
## -----
```

##			model	Beta	Std. Error	Std. Beta
t	Sig	lower	upper			
##	-----					
##			(Intercept)	188.495	23.945	
7.872	0.000	141.104	235.886			
##			Income.composition.of.resources	802.586	34.615	0.901
23.186	0.000	734.078	871.095			
##	-----					

```
##
##
##
## Stepwise Selection: Step 2
```

```
## - Adult.Mortality added
```

Model Summary			
##	0.934	RMSE	48.832
## R-Squared	0.872	Coef. Var	6.687
## Adj. R-Squared	0.870	MSE	2384.535
## Pred R-Squared	0.864	MAE	36.404

```
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
```

		ANOVA				
		Sum of Squares	DF	Mean Square	F	Sig.
Regression	2007944.409	2	1003972.204	421.035	0.0000	
Residual	295682.393	124	2384.535			
Total	2303626.802	126				

[illegible]

```

-----
##
t      Sig      lower      model      Beta      Std. Error      Std. Beta
## -----
##
##      (Intercept)      368.608      30.824
11.958      0.000      307.598      429.617
## Income.composition.of.resources      635.181      36.095      0.713
17.597      0.000      563.738      706.624
##      Adult.Mortality      -0.419      0.055      -0.309      -
7.632      0.000      -0.527      -0.310
## -----
##
##
##
##      Model Summary
## -----
## R      0.934      RMSE      48.832
## R-Squared      0.872      Coef. Var      6.687
## Adj. R-Squared      0.870      MSE      2384.535
## Pred R-Squared      0.864      MAE      36.404
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##      ANOVA
## -----
##      Sum of
##      Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2007944.409      2      1003972.204      421.035      0.0000
## Residual      295682.393      124      2384.535
## Total      2303626.802      126
## -----
##
##      Parameter Estimates
## -----
-----
##
t      Sig      lower      model      Beta      Std. Error      Std. Beta
## -----
##
##      (Intercept)      368.608      30.824
11.958      0.000      307.598      429.617
## Income.composition.of.resources      635.181      36.095      0.713
17.597      0.000      563.738      706.624
##      Adult.Mortality      -0.419      0.055      -0.309      -
7.632      0.000      -0.527      -0.310

```

```

## -----
##
##
## Stepwise Selection: Step 3
##
## - Hepatitis.B added
##
##                               Model Summary
## -----
## R                               0.942          RMSE                45.789
## R-Squared                       0.888          Coef. Var          6.270
## Adj. R-Squared                   0.885          MSE                2096.664
## Pred R-Squared                   0.877          MAE                35.786
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2045737.074           3      681912.358      325.237      0.0000
## Residual        257889.729          123           2096.664
## Total           2303626.802          126
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               Sig      lower      model      Beta      Std. Error      Std. Beta
##                               t      upper
## -----
##                               (Intercept)      328.105      30.437
## 10.780      0.000      267.856      388.354
## ## Income.composition.of.resources      605.448      34.563      0.679
## 17.517      0.000      537.032      673.864
## ## Adult.Mortality      -0.415      0.051      -0.306
## 8.059      0.000      -0.517      -0.313
## ## Hepatitis.B      0.737      0.174      0.133
## 4.246      0.000      0.393      1.080
## -----
##
##
##
##

```

```

##                               Model Summary
## -----
## R                               0.942          RMSE          45.789
## R-Squared                       0.888          Coef. Var      6.270
## Adj. R-Squared                  0.885          MSE           2096.664
## Pred R-Squared                  0.877          MAE           35.786
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      2045737.074              3          681912.358          325.237          0.0000
## Residual        257889.729             123           2096.664
## Total          2303626.802             126
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model          Beta          Std. Error          Std. Beta
## t          Sig          lower          upper
## -----
##                               (Intercept)          328.105          30.437
## 10.780      0.000          267.856          388.354
## Income.composition.of.resources          605.448          34.563          0.679
## 17.517      0.000          537.032          673.864
## Adult.Mortality          -0.415          0.051          -0.306
## 8.059       0.000          -0.517          -0.313
## Hepatitis.B          0.737          0.174          0.133
## 4.246       0.000          0.393          1.080
## -----
## -----
##
##
## Stepwise Selection: Step 4
##
## - HIV.AIDS added
##
##                               Model Summary
## -----
## R                               0.944          RMSE          45.246
## R-Squared                       0.892          Coef. Var      6.196
## Adj. R-Squared                  0.888          MSE           2047.238

```

```

## Pred R-Squared      0.876      MAE      35.071
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2053863.798      4      513465.950      250.809      0.0000
## Residual        249763.004      122      2047.238
## Total           2303626.802      126
## -----
##
##                               Parameter Estimates
## -----
##
##                               t      Sig      lower      model      Beta      Std. Error      Std. Beta
##                               t      Sig      lower      upper
## -----
##                               (Intercept)      337.520      30.445
## 11.086      0.000      277.250      397.790
## Income.composition.of.resources      599.234      34.296      0.673
## 17.473      0.000      531.342      667.126
## Adult.Mortality      -0.355      0.059      -0.262      -
## 6.016      0.000      -0.472      -0.238
## Hepatitis.B      0.626      0.180      0.113
## 3.469      0.001      0.269      0.983
## HIV.AIDS      -7.195      3.611      -0.082      -
## 1.992      0.049      -14.345      -0.046
## -----
##
##
##
##                               Model Summary
## -----
## R      0.944      RMSE      45.246
## R-Squared      0.892      Coef. Var      6.196
## Adj. R-Squared      0.888      MSE      2047.238
## Pred R-Squared      0.876      MAE      35.071
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA

```

```

## -----
##               Sum of
##               Squares          DF      Mean Square          F          Sig.
## -----
## Regression      2053863.798           4      513465.950      250.809      0.0000
## Residual        249763.004          122       2047.238
## Total          2303626.802          126
## -----
##
##                                     Parameter Estimates
## -----
##
##               Sig      lower      model      Beta      Std. Error      Std. Beta
##               t               upper
## -----
##               (Intercept)      337.520      30.445
## 11.086      0.000      277.250      397.790
## Income.composition.of.resources      599.234      34.296      0.673
## 17.473      0.000      531.342      667.126
## Adult.Mortality      -0.355      0.059      -0.262      -
## 6.016      0.000      -0.472      -0.238
## Hepatitis.B      0.626      0.180      0.113
## 3.469      0.001      0.269      0.983
## HIV.AIDS      -7.195      3.611      -0.082      -
## 1.992      0.049      -14.345      -0.046
## -----
##
##
##
## Stepwise Selection: Step 5
##
## - Adult.Mortality:HIV.AIDS added
##
##               Model Summary
## -----
## R      0.950      RMSE      43.087
## R-Squared      0.902      Coef. Var      5.900
## Adj. R-Squared      0.898      MSE      1856.452
## Pred R-Squared      0.875      MAE      33.269
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##               ANOVA
## -----
##               Sum of
##               Squares          DF      Mean Square          F          Sig.

```

```

## -----
## Regression      2078996.131          5      415799.226      223.975      0.0000
## Residual        224630.672        121        1856.452
## Total           2303626.802        126
## -----
##
##                                     Parameter Estimates
## -----
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t                                     -----
##          -----
##                                     (Intercept)      396.517          33.131
11.968      0.000      330.926      462.108
## Income.composition.of.resources      564.804          33.973          0.634
16.625      0.000      497.546      632.062
## Adult.Mortality      -0.455          0.062      -0.336      -
7.290      0.000      -0.579      -0.332
## Hepatitis.B          0.429          0.180          0.077
2.388      0.018          0.073          0.785
## HIV.AIDS      -34.365          8.146      -0.390      -
4.219      0.000      -50.493      -18.238
## Adult.Mortality:HIV.AIDS      0.080          0.022          0.347
3.679      0.000          0.037          0.123
## -----
## -----
##
##                                     Model Summary
## -----
## R          0.950      RMSE          43.087
## R-Squared    0.902      Coef. Var          5.900
## Adj. R-Squared    0.898      MSE          1856.452
## Pred R-Squared    0.875      MAE          33.269
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of
##          Squares          DF      Mean Square          F          Sig.
## -----
## Regression      2078996.131          5      415799.226      223.975      0.0000
## Residual        224630.672        121        1856.452
## Total           2303626.802        126
## -----

```

```

##
##                                     Parameter Estimates
## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      396.517      33.131
## 11.968      0.000      330.926      462.108
## Income.composition.of.resources      564.804      33.973      0.634
## 16.625      0.000      497.546      632.062
## Adult.Mortality      -0.455      0.062      -0.336      -
## 7.290      0.000      -0.579      -0.332
## Hepatitis.B      0.429      0.180      0.077
## 2.388      0.018      0.073      0.785
## HIV.AIDS      -34.365      8.146      -0.390      -
## 4.219      0.000      -50.493      -18.238
## Adult.Mortality:HIV.AIDS      0.080      0.022      0.347
## 3.679      0.000      0.037      0.123
## -----
##
##
##
## Stepwise Selection: Step 6
##
## - under.five.deaths added
##
##                                     Model Summary
## -----
## R      0.950      RMSE      43.102
## R-Squared      0.903      Coef. Var      5.902
## Adj. R-Squared      0.898      MSE      1857.797
## Pred R-Squared      0.873      MAE      33.137
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of
##          Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2080691.190      6      346781.865      186.663      0.0000
## Residual      222935.612      120      1857.797
## Total      2303626.802      126
## -----
##
##

```


Parameter Estimates						

##			model	Beta	Std. Error	Std. Beta
t	Sig	lower	upper			
##	-----					
##		(Intercept)		401.029	33.478	
11.979	0.000	334.745	467.312			
##	Income.composition.of.resources	560.340		34.305	0.629	
16.334	0.000	492.418	628.261			
##	Adult.Mortality	-0.451		0.063	-0.333	
7.206	0.000	-0.575	-0.327			
##	Hepatitis.B	0.419		0.180	0.076	
2.325	0.022	0.062	0.776			
##	HIV.AIDS	-34.606		8.153	-0.393	
4.245	0.000	-50.748	-18.464			
##	under.five.deaths	-0.030		0.032	-0.028	
0.955	0.341	-0.093	0.032			
##	Adult.Mortality:HIV.AIDS	0.080		0.022	0.347	
3.683	0.000	0.037	0.123			
##	-----					

##						
##						
##	Model Summary					
##	-----					
##	R	0.950		RMSE	43.102	
##	R-Squared	0.903		Coef. Var	5.902	
##	Adj. R-Squared	0.898		MSE	1857.797	
##	Pred R-Squared	0.873		MAE	33.137	
##	-----					
##	RMSE: Root Mean Square Error					
##	MSE: Mean Square Error					
##	MAE: Mean Absolute Error					
##						
##	ANOVA					
##	-----					
##		Sum of				
##		Squares	DF	Mean Square	F	Sig.
##	-----					
##	Regression	2080691.190	6	346781.865	186.663	0.0000
##	Residual	222935.612	120	1857.797		
##	Total	2303626.802	126			
##	-----					
##						
##						
##	Parameter Estimates					
##	-----					

```

##          t          Sig          lower          model          Beta          Std. Error          Std. Beta
##          t          Sig          lower          upper
## -----
##          (Intercept)          401.029          33.478
11.979      0.000      334.745      467.312
## Income.composition.of.resources          560.340          34.305          0.629
16.334      0.000      492.418      628.261
##          Adult.Mortality          -0.451          0.063          -0.333      -
7.206      0.000      -0.575      -0.327
##          Hepatitis.B          0.419          0.180          0.076
2.325      0.022          0.062          0.776
##          HIV.AIDS          -34.606          8.153          -0.393      -
4.245      0.000      -50.748      -18.464
##          under.five.deaths          -0.030          0.032          -0.028      -
0.955      0.341      -0.093          0.032
##          Adult.Mortality:HIV.AIDS          0.080          0.022          0.347
3.683      0.000          0.037          0.123
## -----
##
##
##
## Stepwise Selection: Step 7
##
## - Adult.Mortality:HIV.AIDS added
##
##          Model Summary
## -----
## R          0.945          RMSE          45.284
## R-Squared          0.892          Coef. Var          6.201
## Adj. R-Squared          0.888          MSE          2050.677
## Pred R-Squared          0.875          MAE          35.066
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##          ANOVA
## -----
##          Sum of
##          Squares          DF          Mean Square          F          Sig.
## -----
## Regression          2055494.915          5          411098.983          200.47          0.0000
## Residual          248131.887          121          2050.677
## Total          2303626.802          126
## -----
##
##          Parameter Estimates
## -----

```

```

-----
##          t          Sig          lower          model          Beta          Std. Error          Std. Beta
##          t          Sig          lower          upper          -----
## -----
##          (Intercept)          341.872          30.859
11.078          0.000          280.778          402.966
## Income.composition.of.resources          594.898          34.667          0.668
17.160          0.000          526.265          663.530
##          Adult.Mortality          -0.351          0.059          -0.259          -
5.924          0.000          -0.468          -0.234
##          Hepatitis.B          0.616          0.181          0.111
3.405          0.001          0.258          0.974
##          HIV.AIDS          -7.397          3.622          -0.084          -
2.043          0.043          -14.567          -0.227
##          under.five.deaths          -0.030          0.033          -0.027          -
0.892          0.374          -0.095          0.036
## -----
##
##
##
##
## Final Model Output
## -----
##
##          Model Summary
## -----
## R          0.945          RMSE          45.284
## R-Squared          0.892          Coef. Var          6.201
## Adj. R-Squared          0.888          MSE          2050.677
## Pred R-Squared          0.875          MAE          35.066
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##          ANOVA
## -----
##          Sum of
##          Squares          DF          Mean Square          F          Sig.
## -----
## Regression          2055494.915          5          411098.983          200.47          0.0000
## Residual          248131.887          121          2050.677
## Total          2303626.802          126
## -----
##
##          Parameter Estimates
## -----

```

## t	Sig	lower	model upper	Beta	Std. Error	Std. Beta
##						
##			(Intercept)	341.872	30.859	
11.078	0.000	280.778	402.966			
##			Income.composition.of.resources	594.898	34.667	0.668
17.160	0.000	526.265	663.530			
##			Adult.Mortality	-0.351	0.059	-0.259
5.924	0.000	-0.468	-0.234			
##			Hepatitis.B	0.616	0.181	0.111
3.405	0.001	0.258	0.974			
##			HIV.AIDS	-7.397	3.622	-0.084
2.043	0.043	-14.567	-0.227			
##			under.five.deaths	-0.030	0.033	-0.027
0.892	0.374	-0.095	0.036			

life2015_bothmodel

##	##	Stepwise Selection Summary				
##	##					
##	##	Step	Variable	Added/ Removed	R-Square	Adj. R-
##	##	Square	C(p)	AIC	RMSE	
##	##					
##	1	Income.composition.of.resources		addition	0.811	
0.810		110.9290	1399.9337	58.9639		
##	2	Adult.Mortality		addition	0.872	
0.870		38.1580	1353.0229	48.8317		
##	3	Hepatitis.B		addition	0.888	
0.885		19.8150	1337.6551	45.7893		
##	4	HIV.AIDS		addition	0.892	
0.888		17.4400	1335.5886	45.2464		
##	5	Adult.Mortality:HIV.AIDS		addition	0.902	
0.898		5.9120	1324.1197	43.0866		
##	6	under.five.deaths		addition	0.903	
0.898		7.0000	1325.1577	43.1022		
##	7	Adult.Mortality:HIV.AIDS		removal	0.892	
0.888		18.5620	1336.7565	45.2844		

#life2015_bothmodel\$mallows_cp

paste("Stepwise procedure with BOTH indicated to consider all the variables and the interaction term.")

```
## [1] "Stepwise procedure with BOTH indicated to consider all the variables
and the interaction term."
```

```
# (P.3.2) FORWARD
```

```
life2015_forwardmodel = ols_step_forward_p(life2015_box_modelP, pent = 0.05,
details = TRUE)
```

```
## Forward Selection Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1. Income.composition.of.resources
```

```
## 2. Adult.Mortality
```

```
## 3. HIV.AIDS
```

```
## 4. under.five.deaths
```

```
## 5. Hepatitis.B
```

```
## 6. Adult.Mortality:HIV.AIDS
```

```
##
```

```
## We are selecting variables based on p value...
```

```
##
```

```
##
```

```
## Forward Selection: Step 1
```

```
##
```

```
## - Income.composition.of.resources
```

```
##
```

```
## Model Summary
```

```
## -----
```

## R	0.901	RMSE	58.964
## R-Squared	0.811	Coef. Var	8.074
## Adj. R-Squared	0.810	MSE	3476.746
## Pred R-Squared	0.806	MAE	46.343

```
## -----
```

```
## RMSE: Root Mean Square Error
```

```
## MSE: Mean Square Error
```

```
## MAE: Mean Absolute Error
```

```
##
```

```
## ANOVA
```

```
## -----
```

##	Sum of				
##	Squares	DF	Mean Square	F	Sig.
##					
## Regression	1869033.599	1	1869033.599	537.581	0.0000
## Residual	434593.203	125	3476.746		
## Total	2303626.802	126			

```
## -----
```

```
##
```

```
## Parameter Estimates
```

```
## -----
```

```

##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##          (Intercept)      188.495      23.945
7.872      0.000      141.104      235.886
## Income.composition.of.resources      802.586      34.615      0.901
23.186      0.000      734.078      871.095
## -----
##
##
##
## Forward Selection: Step 2
##
## - Adult.Mortality
##
##          Model Summary
## -----
## R          0.934      RMSE          48.832
## R-Squared      0.872      Coef. Var      6.687
## Adj. R-Squared      0.870      MSE          2384.535
## Pred R-Squared      0.864      MAE          36.404
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##          ANOVA
## -----
##          Sum of
##          Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2007944.409      2      1003972.204      421.035      0.0000
## Residual          295682.393      124          2384.535
## Total          2303626.802      126
## -----
##
##          Parameter Estimates
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##          (Intercept)      368.608      30.824
11.958      0.000      307.598      429.617
## Income.composition.of.resources      635.181      36.095      0.713
17.597      0.000      563.738      706.624
## Adult.Mortality      -0.419      0.055      -0.309      -

```

```

7.632    0.000    -0.527    -0.310
## -----
##
##
##
## Forward Selection: Step 3
##
## - Hepatitis.B
##
##                               Model Summary
## -----
## R                               0.942          RMSE                45.789
## R-Squared                       0.888          Coef. Var          6.270
## Adj. R-Squared                  0.885          MSE                2096.664
## Pred R-Squared                  0.877          MAE                35.786
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      2045737.074                3          681912.358          325.237          0.0000
## Residual        257889.729               123              2096.664
## Total           2303626.802               126
## -----
##
##                               Parameter Estimates
## -----
##                               Sig          lower          model          Beta          Std. Error          Std. Beta
##                               t          upper
## -----
##                               (Intercept)          328.105          30.437
## 10.780          0.000          267.856          388.354
## ## Income.composition.of.resources          605.448          34.563          0.679
## 17.517          0.000          537.032          673.864
## ## Adult.Mortality          -0.415          0.051          -0.306          -
## 8.059          0.000          -0.517          -0.313
## ## Hepatitis.B          0.737          0.174          0.133
## 4.246          0.000          0.393          1.080
## -----
##
##
##

```

```

##
## Forward Selection: Step 4
##
## - HIV.AIDS
##
##                               Model Summary
## -----
## R                               0.944          RMSE                45.246
## R-Squared                       0.892          Coef. Var          6.196
## Adj. R-Squared                  0.888          MSE                2047.238
## Pred R-Squared                  0.876          MAE                35.071
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      2053863.798              4          513465.950          250.809          0.0000
## Residual        249763.004             122           2047.238
## Total          2303626.802             126
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               t          Sig          lower          model          Beta          Std. Error          Std. Beta
##                               upper
## -----
##                               (Intercept)          337.520          30.445
## 11.086          0.000          277.250          397.790
## ## Income.composition.of.resources          599.234          34.296          0.673
## 17.473          0.000          531.342          667.126
## ## Adult.Mortality          -0.355          0.059          -0.262          -
## 6.016          0.000          -0.472          -0.238
## ## Hepatitis.B          0.626          0.180          0.113
## 3.469          0.001          0.269          0.983
## ## HIV.AIDS          -7.195          3.611          -0.082          -
## 1.992          0.049          -14.345          -0.046
## -----
##
##
##
## Forward Selection: Step 5
##

```


- Adult.Mortality:HIV.AIDS

##

Model Summary

```
## -----
## R                0.950      RMSE                43.087
## R-Squared        0.902      Coef. Var            5.900
## Adj. R-Squared   0.898      MSE                1856.452
## Pred R-Squared   0.875      MAE                33.269
## -----
```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

```
## -----
## Sum of
## Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2078996.131      5      415799.226    223.975    0.0000
## Residual      224630.672     121      1856.452
## Total         2303626.802     126
## -----
```

##

Parameter Estimates

```
## -----
## Sig      lower      model      Beta      Std. Error      Std. Beta
## t          upper
## -----
## (Intercept)      396.517      33.131
11.968    0.000    330.926    462.108
## Income.composition.of.resources      564.804      33.973      0.634
16.625    0.000    497.546    632.062
## Adult.Mortality      -0.455      0.062      -0.336    -
7.290    0.000    -0.579    -0.332
## Hepatitis.B      0.429      0.180      0.077
2.388    0.018      0.073      0.785
## HIV.AIDS      -34.365      8.146      -0.390    -
4.219    0.000    -50.493    -18.238
## Adult.Mortality:HIV.AIDS      0.080      0.022      0.347
3.679    0.000      0.037      0.123
## -----
```

##

##

##

Forward Selection: Step 6

##

- under.five.deaths

```

##
##                               Model Summary
## -----
## R                               0.950           RMSE           43.102
## R-Squared                       0.903           Coef. Var       5.902
## Adj. R-Squared                   0.898           MSE           1857.797
## Pred R-Squared                   0.873           MAE           33.137
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares           DF           Mean Square           F           Sig.
## -----
## Regression      2080691.190                6           346781.865           186.663           0.0000
## Residual        222935.612               120             1857.797
## Total          2303626.802               126
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model           Beta           Std. Error           Std. Beta
## t           Sig           lower           upper
## -----
##                               (Intercept)           401.029           33.478
## 11.979      0.000           334.745           467.312
## Income.composition.of.resources           560.340           34.305           0.629
## 16.334      0.000           492.418           628.261
## Adult.Mortality           -0.451           0.063           -0.333      -
## 7.206       0.000           -0.575           -0.327
## Hepatitis.B           0.419           0.180           0.076
## 2.325       0.022           0.062           0.776
## HIV.AIDS           -34.606           8.153           -0.393      -
## 4.245       0.000           -50.748           -18.464
## under.five.deaths           -0.030           0.032           -0.028      -
## 0.955       0.341           -0.093           0.032
## Adult.Mortality:HIV.AIDS           0.080           0.022           0.347
## 3.683       0.000           0.037           0.123
## -----
## -----
##
##
##
## Variables Entered:

```

```

##
## + Income.composition.of.resources
## + Adult.Mortality
## + Hepatitis.B
## + HIV.AIDS
## + Adult.Mortality:HIV.AIDS
## + under.five.deaths
##
##
## Final Model Output
## -----
##
##                               Model Summary
## -----
## R                               0.950          RMSE                43.102
## R-Squared                       0.903          Coef. Var          5.902
## Adj. R-Squared                   0.898          MSE                1857.797
## Pred R-Squared                   0.873          MAE                33.137
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      2080691.190              6          346781.865          186.663          0.0000
## Residual         222935.612             120              1857.797
## Total           2303626.802             126
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               model          Beta          Std. Error          Std. Beta
##                               t          Sig          lower          upper
## -----
##                               (Intercept)          401.029          33.478
## 11.979          0.000          334.745          467.312
## Income.composition.of.resources          560.340          34.305          0.629
## 16.334          0.000          492.418          628.261
## Adult.Mortality          -0.451          0.063          -0.333          -
## 7.206          0.000          -0.575          -0.327
## Hepatitis.B          0.419          0.180          0.076
## 2.325          0.022          0.062          0.776
## HIV.AIDS          -34.606          8.153          -0.393          -
## 4.245          0.000          -50.748          -18.464

```

```
##          under.five.deaths    -0.030      0.032      -0.028      -
0.955      0.341      -0.093      0.032
##          Adult.Mortality:HIV.AIDS    0.080      0.022      0.347
3.683      0.000      0.037      0.123
## -----
```

```
life2015_forwardmodel
```

```
##
##                               Selection Summary
## -----
```

##	Variable				
## Step	Entered		R-Square	Adj.	C(p)
AIC	RMSE			R-Square	
## 1	Income.composition.of.resources		0.8113	0.8098	
110.9294	1399.9337	58.9639			
## 2	Adult.Mortality		0.8716	0.8696	
38.1576	1353.0229	48.8317			
## 3	Hepatitis.B		0.8881	0.8853	
19.8148	1337.6551	45.7893			
## 4	HIV.AIDS		0.8916	0.8880	
17.4404	1335.5886	45.2464			
## 5	Adult.Mortality:HIV.AIDS		0.9025	0.8985	
5.9124	1324.1197	43.0866			
## 6	under.five.deaths		0.9032	0.8984	
7.0000	1325.1577	43.1022			

```
#life2015_forwardmodel$mallows_cp
```

```
paste("Stepwise procedure with FORWARD indicated to consider all the
variables and the interaction term.")
```

```
## [1] "Stepwise procedure with FORWARD indicated to consider all the
variables and the interaction term."
```

```
# (P.3.3) BACKWARD
```

```
life2015_backwardmodel = ols_step_backward_p(life2015_box_modelP, prem =
0.05, details = TRUE)
```

```
## Backward Elimination Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1 . Income.composition.of.resources
```

```
## 2 . Adult.Mortality
```

```

## 3 . HIV.AIDS
## 4 . under.five.deaths
## 5 . Hepatitis.B
## 6 . Adult.Mortality:HIV.AIDS
##
## We are eliminating variables based on p value...
##
## - under.five.deaths
##
## Backward Elimination: Step 1
##
## Variable under.five.deaths Removed
##
##
##                               Model Summary
## -----
## R                               0.950          RMSE                43.087
## R-Squared                       0.902          Coef. Var          5.900
## Adj. R-Squared                  0.898          MSE                1856.452
## Pred R-Squared                  0.875          MAE                33.269
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      2078996.131              5          415799.226          223.975          0.0000
## Residual         224630.672             121             1856.452
## Total           2303626.802             126
## -----
##
##                               Parameter Estimates
## -----
## -----
##                               model          Beta          Std. Error          Std. Beta
## t          Sig          lower          upper
## -----
##                               (Intercept)          396.517          33.131
## 11.968      0.000          330.926          462.108
## Income.composition.of.resources          564.804          33.973          0.634
## 16.625      0.000          497.546          632.062
## Adult.Mortality          -0.455          0.062          -0.336      -
## 7.290      0.000          -0.579          -0.332
## HIV.AIDS          -34.365          8.146          -0.390      -
## 4.219      0.000          -50.493          -18.238
## Hepatitis.B          0.429          0.180          0.077

```

```

2.388      0.018      0.073      0.785
##      Adult.Mortality:HIV.AIDS      0.080      0.022      0.347
3.679      0.000      0.037      0.123
## -----
##
##
##
## No more variables satisfy the condition of p value = 0.05
##
##
## Variables Removed:
##
## - under.five.deaths
##
##
## Final Model Output
## -----
##
##                      Model Summary
## -----
## R                      0.950      RMSE                      43.087
## R-Squared              0.902      Coef. Var                5.900
## Adj. R-Squared         0.898      MSE                      1856.452
## Pred R-Squared         0.875      MAE                      33.269
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                      ANOVA
## -----
##                      Sum of
##                      Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2078996.131      5      415799.226      223.975      0.0000
## Residual        224630.672      121      1856.452
## Total           2303626.802      126
## -----
##
##                      Parameter Estimates
## -----
## -----
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##                      (Intercept)      396.517      33.131
11.968      0.000      330.926      462.108
## Income.composition.of.resources      564.804      33.973      0.634

```

```

16.625    0.000    497.546    632.062
##          Adult.Mortality    -0.455          0.062    -0.336    -
7.290    0.000    -0.579    -0.332
##          HIV.AIDS    -34.365          8.146    -0.390    -
4.219    0.000    -50.493    -18.238
##          Hepatitis.B    0.429          0.180    0.077
2.388    0.018    0.073    0.785
##          Adult.Mortality:HIV.AIDS    0.080          0.022    0.347
3.679    0.000    0.037    0.123
## -----
-----

life2015_backwardmodel

##
##
##          Elimination Summary
## -----
-----
##          Variable          Adj.
## Step          Removed          R-Square          R-Square          C(p)          AIC
RMSE
## -----
-----
##    1    under.five.deaths    0.9025    0.8985    5.9124    1324.1197
43.0866
## -----
-----

#life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD indicated to remove under.five.deaths
variable.")

## [1] "Stepwise procedure with BACKWARD indicated to remove
under.five.deaths variable."

# (P.4) BEST SUBSET
#OPTION 1
ks = ols_step_best_subset(life2015_box_modelP, details = TRUE)

#Summary of Selected model based on cp, aic, AdjustedR2
rsquare <- c(ks$rsq)
cp <- c(ks$cp)
aic <- c(ks$aic)
AdjustedR2 <- c(ks$adjr)
cbind(rsquare, cp, aic, AdjustedR2)

##          rsquare          cp          aic AdjustedR2
## [1,] 0.8113439 110.929357 1399.934 0.8098347
## [2,] 0.8716448 38.157556 1353.023 0.8695746
## [3,] 0.8880506 19.814823 1337.655 0.8853201

```

```
## [4,] 0.8978937    9.609600 1327.967    0.8945459
## [5,] 0.9024883    5.912403 1324.120    0.8984588
## [6,] 0.9032241    7.000000 1325.158    0.8983853
```

#OPTION 2

```
best.subset <- regsubsets((((Life.expectancy^1.66667) - 1)/1.66667) ~
Income.composition.of.resources +
                        Adult.Mortality + HIV.AIDS + under.five.deaths +
Hepatitis.B + Adult.Mortality * HIV.AIDS,
                        data = life2015[-c(119, 53, 93), ], nv = 4)
summary(best.subset)
```

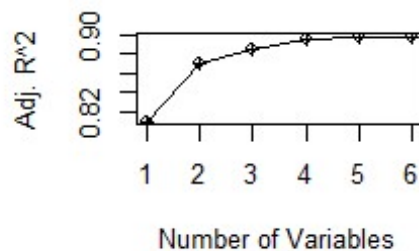
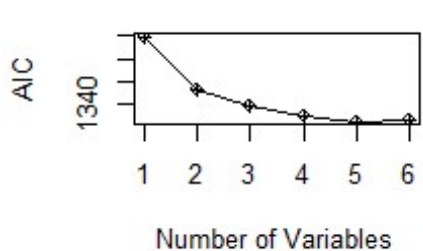
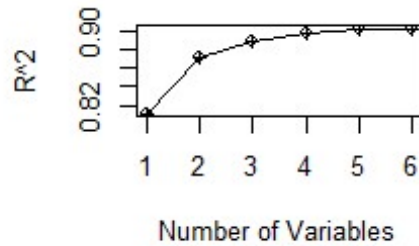
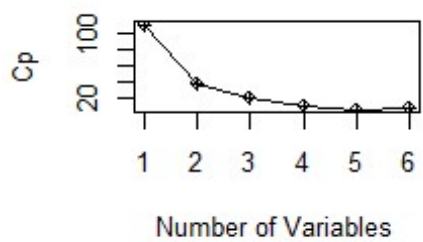
```
## Subset selection object
## Call: regsubsets.formula((((Life.expectancy^1.66667) - 1)/1.66667) ~
##      Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
##      under.five.deaths + Hepatitis.B + Adult.Mortality * HIV.AIDS,
##      data = life2015[-c(119, 53, 93), ], nv = 4)
## 6 Variables (and intercept)
##
##              Forced in Forced out
## Income.composition.of.resources    FALSE    FALSE
## Adult.Mortality                    FALSE    FALSE
## HIV.AIDS                          FALSE    FALSE
## under.five.deaths                 FALSE    FALSE
## Hepatitis.B                      FALSE    FALSE
## Adult.Mortality:HIV.AIDS          FALSE    FALSE
## 1 subsets of each size up to 4
## Selection Algorithm: exhaustive
##      Income.composition.of.resources Adult.Mortality HIV.AIDS
## 1  ( 1 ) "*"                " "                " "
## 2  ( 1 ) "*"                "*"                " "
## 3  ( 1 ) "*"                "*"                " "
## 4  ( 1 ) "*"                "*"                "*"
##      under.five.deaths Hepatitis.B Adult.Mortality:HIV.AIDS
## 1  ( 1 ) " "            " "            " "
## 2  ( 1 ) " "            " "            " "
## 3  ( 1 ) " "            "*"            " "
## 4  ( 1 ) " "            " "            "*"

```

```
reg.summary <- summary(best.subset)
```

#Plotting

```
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"R^2")
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"AIC")
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"Adj. R^2")
```

```
paste("Based on Marlow's Cp the minimum value was obtained without
under.five.deaths, but the model with this variable presented a slightly high
value.")
```

```
## [1] "Based on Marlow's Cp the minimum value was obtained without
under.five.deaths, but the model with this variable presented a slightly high
value."
```

```
#MODEL DIAGNOSTICS
```

```
# (P.5) CHECKING LINEARITY
```

```
#Residuals plot
```

```
plot(life2015_box_modelP, which = 1)
```

```
# a Scale Location Plot
```

```
plot(life2015_box_modelP, which = 3)
```

```
paste("It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected.")
```

```
## [1] "It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected."
```

```

# (P.6) Heteroscedasticity Test - the Breush-Pagan test
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
# Ha: heteroscedasticity is presented
bptest(life2015_box_modelP)

##
## studentized Breusch-Pagan test
##
## data:  life2015_box_modelP
## BP = 12.235, df = 6, p-value = 0.05693

paste("The output displays the Breush-Pagan test that from the model
presented the p-value 0.05693 > 0.05, indicating the we should NOT REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity does not exist.")

## [1] "The output displays the Breush-Pagan test that from the model
presented the p-value 0.05693 > 0.05, indicating the we should NOT REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity does not exist."

# (P.7) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_modelP))

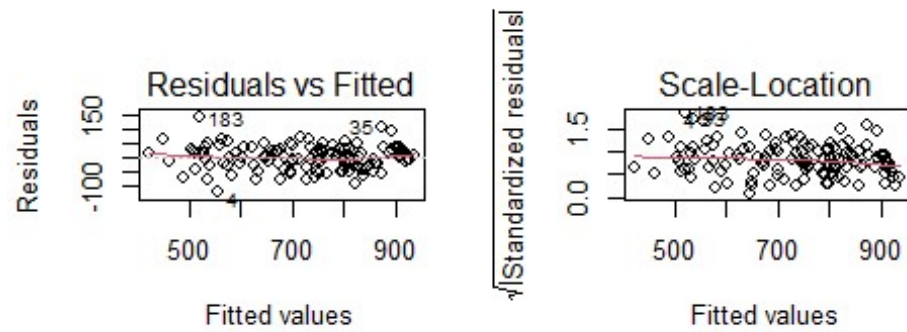
##
## Shapiro-Wilk normality test
##
## data:  residuals(life2015_box_modelP)
## W = 0.99081, p-value = 0.5664

paste("Shapiro-Wilk normality test presented the p-value = 0.5664 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level.")

## [1] "Shapiro-Wilk normality test presented the p-value = 0.5664 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level."

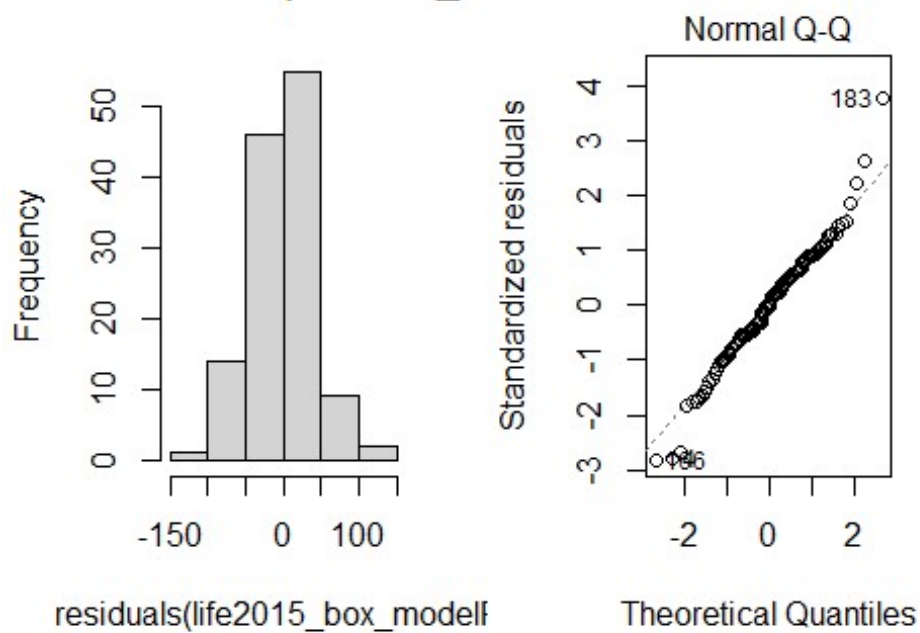
#Plots
par(mfrow = c(1,2))

```



```
hist(residuals(life2015_box_modelP))
plot(life2015_box_modelM, which = 2)
```

am of residuals(life2015_b



```

paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")

## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."

# (P.8) Multicollinearity test
#VIF
imcdiag(life2015_box_modelP, method = "VIF")

##
## Call:
## imcdiag(mod = life2015_box_modelP, method = "VIF")
##
## VIF Multicollinearity Diagnostics
##
##                               VIF detection
## Income.composition.of.resources  1.8380          0
## Adult.Mortality                  2.6515          0
## HIV.AIDS                        10.6289          1
## under.five.deaths                1.0612          0
## Hepatitis.B                     1.3093          0
## Adult.Mortality:HIV.AIDS         11.0314          1
##
## Multicollinearity may be due to HIV.AIDS Adult.Mortality:HIV.AIDS
regressors
##
## 1 --> COLLINEARITY is detected by the test
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. ")

## [1] "Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. "

# (P.9) LEVERAGE OF THE POINTS
# Leverage Points
lev = hatvalues(life2015_box_modelP)
p = length(coef(life2015_box_modelP))
n = nrow(life2015)
outlier = lev[lev > (2 * p/n)]
print(outlier)

##          26          53          75          93         118         119         126
127
## 0.1235814 0.3731765 0.6032480 0.5985183 0.1540768 0.2893001 0.1173017
0.1316946

```

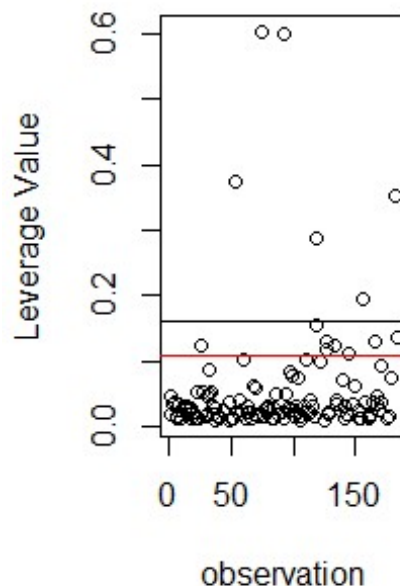
```
##          133          144          156          166          182          183
## 0.1235340 0.1115889 0.1966767 0.1292409 0.3540528 0.1360596

plot(rownames(life2015[-c(119, 53, 93), ]), lev, main = "Leverage in Life2015
Dataset",
      xlab = "observation", ylab = "Leverage Value")
abline(h = 2 * p/n, lty = 1, col = "red")
abline(h = 3 * p/n, lty = 1)

paste("The results shows that that the data points #26, #53, #75, #93, #118,
#119, #126, #127, #166, #182 and #183 presented leverage values higher than
2.p/n that might be outliers.")

## [1] "The results shows that that the data points #26, #53, #75, #93, #118,
#119, #126, #127, #166, #182 and #183 presented leverage values higher than
2.p/n that might be outliers."
```

Leverage in Life2015 Data



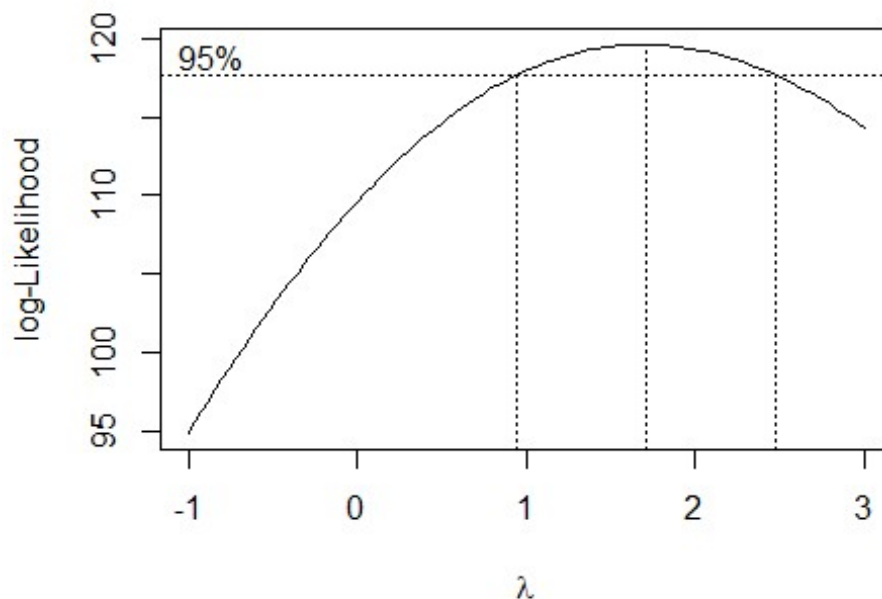
```
##### MODEL Q - BOX COX TRANSFORMATION of MODEL G with Hepatitis.B
#####
paste("t-test of previous Box-Cox transformation indicated no significance of
interaction term after this transformation")

## [1] "t-test of previous Box-Cox transformation indicated no significance
of interaction term after this transformation"

#(Q.1) BOX-COX TRANSFORMATION
# The necessity of transformation is due to the indication of
heterosdasticity in model E
```

```
# For this transformation we will be considering only the significant
parameters obtained beforehand
life2015_reduced_modelQ <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
Hepatitis.B + Adult.Mortality * HIV.AIDS,
data = life2015)

bc = boxcox(life2015_reduced_modelQ, lambda = seq(-1,3))
```



```
#Extract best Lambda
bestlambda = bc$x[which(bc$y == max(bc$y))]
paste("The best lambda for Box-Cox Transformation is: ", bestlambda)

## [1] "The best lambda for Box-Cox Transformation is:  1.70707070707071"

paste("From the output, as the best lambda would be 1.70707.")

## [1] "From the output, as the best lambda would be 1.70707."

#BOX-COX with LAMBDA = 1.70707
life2015_box_modelQ = lm((((Life.expectancy^1.70707) - 1)/1.70707) ~
Income.composition.of.resources +
Adult.Mortality + HIV.AIDS + Hepatitis.B +
Adult.Mortality * HIV.AIDS,
data = life2015)

# (Q.2) PARTIAL TEST - Individual Coefficients Test (t-test)
```

```

# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_box_modelQ = lm((((Life.expectancy^1.70707) - 1)/1.70707) ~
Income.composition.of.resources +
                        Adult.Mortality + HIV.AIDS + Hepatitis.B +
                        Adult.Mortality * HIV.AIDS, data = life2015[-
c(119,53,93),])

summary(life2015_box_modelQ)

##
## Call:
## lm(formula = (((Life.expectancy^1.70707) - 1)/1.70707) ~
Income.composition.of.resources +
##      Adult.Mortality + HIV.AIDS + Hepatitis.B + Adult.Mortality *
##      HIV.AIDS, data = life2015[-c(119, 53, 93), ])
##
## Residuals:
##      Min        1Q    Median        3Q        Max
## -138.153  -30.117    1.544    33.158   167.074
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      450.88454    39.34010   11.461 < 2e-16 ***
## Income.composition.of.resources 671.01036    40.33988   16.634 < 2e-16 ***
## Adult.Mortality      -0.54043     0.07417   -7.287 3.55e-11 ***
## HIV.AIDS            -40.77686     9.67267   -4.216 4.83e-05 ***
## Hepatitis.B         0.50896     0.21354    2.383 0.018706 *
## Adult.Mortality:HIV.AIDS      0.09539     0.02586    3.689 0.000339 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 51.16 on 121 degrees of freedom
## Multiple R-squared:  0.9024, Adjusted R-squared:  0.8984
## F-statistic: 223.7 on 5 and 121 DF,  p-value: < 2.2e-16

paste("The t-test indicated that after the Box-Cox transformation all the
coefficients with p-value < 0.05 that indicates that all the variables have
significant influence in Life Expectancy at 5 % level.")

## [1] "The t-test indicated that after the Box-Cox transformation all the
coefficients with p-value < 0.05 that indicates that all the variables have
significant influence in Life Expectancy at 5 % level."

#confidence interval for the coefficients
confint(life2015_box_modelQ, level = 0.95)

##              2.5 %        97.5 %
## (Intercept)      373.00043645 528.7686394
## Income.composition.of.resources 591.14692861 750.8737869

```

```

## Adult.Mortality          -0.68726236  -0.3935969
## HIV.AIDS                 -59.92646782 -21.6272619
## Hepatitis.B              0.08620761   0.9317125
## Adult.Mortality:HIV.AIDS  0.04419529   0.1465805

paste("For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!")

## [1] "For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!"

# (Q.3) STEPWISE REGRESSION PROCEDURE
# (Q.3.1) BOTH
life2015_bothmodel = ols_step_both_p(life2015_box_modelQ, pent = 0.05, prem =
0.05, details = TRUE)

## Stepwise Selection Method
## -----
##
## Candidate Terms:
##
## 1. Income.composition.of.resources
## 2. Adult.Mortality
## 3. HIV.AIDS
## 4. Hepatitis.B
## 5. Adult.Mortality:HIV.AIDS
##
## We are selecting variables based on p value...
##
##
## Stepwise Selection: Step 1
##
## - Income.composition.of.resources added
##
##
##                               Model Summary
## -----
## R                0.901          RMSE                69.927
## R-Squared        0.812          Coef. Var            8.251
## Adj. R-Squared   0.810          MSE                4889.727
## Pred R-Squared   0.806          MAE                54.992
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF      Mean Square          F          Sig.
## -----
## Regression      2633488.698              1      2633488.698      538.576      0.0000

```



```

## Residual      611215.855      125      4889.727
## Total        3244704.553      126
## -----
##
##                                     Parameter Estimates
## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      204.373      28.397
## 7.197      0.000      148.171      260.575
## Income.composition.of.resources      952.684      41.051      0.901
## 23.207      0.000      871.439      1033.930
## -----
##
##
##
## Stepwise Selection: Step 2
##
## - Adult.Mortality added
##
##                                     Model Summary
## -----
## R      0.934      RMSE      57.958
## R-Squared      0.872      Coef. Var      6.839
## Adj. R-Squared      0.870      MSE      3359.148
## Pred R-Squared      0.864      MAE      43.254
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of      DF      Mean Square      F      Sig.
##          Squares
## -----
## Regression      2828170.259      2      1414085.129      420.965      0.0000
## Residual      416534.295      124      3359.148
## Total      3244704.553      126
## -----
##
##
##                                     Parameter Estimates
## -----
##
##          Sig      lower      model      Beta      Std. Error      Std. Beta
##          t          upper

```

```

## -----
##
## (Intercept) 417.598 36.585
11.414 0.000 345.186 490.010
## Income.composition.of.resources 754.502 42.841 0.713
17.611 0.000 669.707 839.297
## Adult.Mortality -0.496 0.065 -0.308 -
7.613 0.000 -0.625 -0.367
## -----
##
##
##
## Model Summary
## -----
## R 0.934 RMSE 57.958
## R-Squared 0.872 Coef. Var 6.839
## Adj. R-Squared 0.870 MSE 3359.148
## Pred R-Squared 0.864 MAE 43.254
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of
## Squares DF Mean Square F Sig.
## -----
## Regression 2828170.259 2 1414085.129 420.965 0.0000
## Residual 416534.295 124 3359.148
## Total 3244704.553 126
## -----
##
## Parameter Estimates
## -----
##
## Sig lower model upper Beta Std. Error Std. Beta
t Sig lower upper Beta Std. Error Std. Beta
## -----
##
## (Intercept) 417.598 36.585
11.414 0.000 345.186 490.010
## Income.composition.of.resources 754.502 42.841 0.713
17.611 0.000 669.707 839.297
## Adult.Mortality -0.496 0.065 -0.308 -
7.613 0.000 -0.625 -0.367
## -----
##
##

```

```

##
##
## Stepwise Selection: Step 3
##
## - Hepatitis.B added
##
##                               Model Summary
## -----
## R                               0.942          RMSE                54.362
## R-Squared                       0.888          Coef. Var          6.415
## Adj. R-Squared                   0.885          MSE                2955.267
## Pred R-Squared                   0.877          MAE                42.505
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      2881206.757              3          960402.252          324.98          0.0000
## Residual        363497.796             123           2955.267
## Total          3244704.553             126
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               t          Sig          lower          model          Beta          Std. Error          Std. Beta
##                               upper
## -----
##                               (Intercept)          369.617          36.136
## 10.228          0.000          298.088          441.146
## ## Income.composition.of.resources          719.280          41.035          0.680
## 17.529          0.000          638.054          800.506
## ## Adult.Mortality          -0.491          0.061          -0.305
## 8.036          0.000          -0.612          -0.370
## ## Hepatitis.B          0.873          0.206          0.133
## 4.236          0.000          0.465          1.281
## -----
##
##
##
##                               Model Summary
## -----
## R                               0.942          RMSE                54.362

```

```

## R-Squared          0.888      Coef. Var      6.415
## Adj. R-Squared     0.885      MSE           2955.267
## Pred R-Squared     0.877      MAE           42.505
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2881206.757      3      960402.252      324.98      0.0000
## Residual        363497.796     123      2955.267
## Total          3244704.553     126
## -----
##
##                               Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t
## -----
##              (Intercept)      369.617      36.136
## 10.228      0.000      298.088      441.146
## Income.composition.of.resources      719.280      41.035      0.680
## 17.529      0.000      638.054      800.506
## Adult.Mortality      -0.491      0.061      -0.305
## 8.036      0.000      -0.612      -0.370
## Hepatitis.B      0.873      0.206      0.133
## 4.236      0.000      0.465      1.281
## -----
##
##
##
## No more variables to be added/removed.
##
##
## Final Model Output
## -----
##
##                               Model Summary
## -----
## R      0.942      RMSE      54.362
## R-Squared      0.888      Coef. Var      6.415
## Adj. R-Squared      0.885      MSE           2955.267
## Pred R-Squared      0.877      MAE           42.505

```

```

## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2881206.757      3      960402.252    324.98    0.0000
## Residual      363497.796     123      2955.267
## Total         3244704.553     126
## -----
##
##                                Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##
##              (Intercept)      369.617      36.136
## 10.228    0.000    298.088    441.146
## Income.composition.of.resources      719.280      41.035      0.680
## 17.529    0.000    638.054    800.506
## Adult.Mortality      -0.491      0.061      -0.305
## 8.036     0.000    -0.612    -0.370
## Hepatitis.B      0.873      0.206      0.133
## 4.236     0.000      0.465      1.281
## -----
## -----
life2015_bothmodel
##
##                                Stepwise Selection Summary
## -----
##
##              Added/      R-Square      Adj.
##              Removed
## Step      C(p)      Variable      RMSE
## Square
## -----
## 1      Income.composition.of.resources      addition      0.812
## 0.810    110.5090    1443.2457    69.9266
## 2      Adult.Mortality      addition      0.872
## 0.870    38.1330    1396.5436    57.9582
## 3      Hepatitis.B      addition      0.888
## 0.885    19.8710    1381.2467    54.3624

```

```
## -----
## -----

#life2015_bothmodel$mallows_cp
paste("Stepwise procedure with BOTH indicated to consider only Income,
Adult.deaths and Hepatitis.B..")

## [1] "Stepwise procedure with BOTH indicated to consider only Income,
Adult.deaths and Hepatitis.B.."

# (Q.3.2) FORWARD
life2015_forwardmodel = ols_step_forward_p(life2015_box_modelQ, pent = 0.05,
details = TRUE)

## Forward Selection Method
## -----
##
## Candidate Terms:
##
## 1. Income.composition.of.resources
## 2. Adult.Mortality
## 3. HIV.AIDS
## 4. Hepatitis.B
## 5. Adult.Mortality:HIV.AIDS
##
## We are selecting variables based on p value...
##
##
## Forward Selection: Step 1
##
## - Income.composition.of.resources
##
##
## Model Summary
## -----
## R                0.901          RMSE                69.927
## R-Squared        0.812          Coef. Var            8.251
## Adj. R-Squared   0.810          MSE                4889.727
## Pred R-Squared   0.806          MAE                54.992
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of
## Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2633488.698      1      2633488.698    538.576    0.0000
## Residual      611215.855     125      4889.727
## Total        3244704.553     126
```

```

## -----
##
##                                     Parameter Estimates
## -----
##
##                                     -----
##          Sig          lower      model      Beta      Std. Error      Std. Beta
##          t          lower      upper
##          -----
##                                     (Intercept)      204.373          28.397
7.197      0.000      148.171      260.575
## Income.composition.of.resources      952.684          41.051          0.901
23.207      0.000      871.439      1033.930
## -----
##
##
##
## Forward Selection: Step 2
##
## - Adult.Mortality
##
##                                     Model Summary
## -----
## R          0.934          RMSE          57.958
## R-Squared    0.872          Coef. Var    6.839
## Adj. R-Squared 0.870          MSE          3359.148
## Pred R-Squared 0.864          MAE          43.254
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                     ANOVA
## -----
##          Sum of          DF      Mean Square          F          Sig.
##          Squares
## -----
## Regression    2828170.259          2      1414085.129      420.965      0.0000
## Residual      416534.295          124          3359.148
## Total        3244704.553          126
## -----
##
##
##                                     Parameter Estimates
## -----
##
##                                     -----
##          Sig          lower      model      Beta      Std. Error      Std. Beta
##          t          lower      upper
##          -----

```

```

##              (Intercept)    417.598      36.585
11.414    0.000    345.186    490.010
## Income.composition.of.resources    754.502      42.841      0.713
17.611    0.000    669.707    839.297
##              Adult.Mortality    -0.496      0.065      -0.308    -
7.613    0.000    -0.625    -0.367
## -----
##
##
##
## Forward Selection: Step 3
##
## - Hepatitis.B
##
##              Model Summary
## -----
## R              0.942      RMSE              54.362
## R-Squared      0.888      Coef. Var          6.415
## Adj. R-Squared 0.885      MSE              2955.267
## Pred R-Squared 0.877      MAE              42.505
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    2881206.757      3      960402.252      324.98      0.0000
## Residual      363497.796      123      2955.267
## Total         3244704.553      126
## -----
##
##              Parameter Estimates
## -----
## -----
##              model      Beta      Std. Error      Std. Beta
## t              Sig      lower      upper
## -----
##              (Intercept)    369.617      36.136
10.228    0.000    298.088    441.146
## Income.composition.of.resources    719.280      41.035      0.680
17.529    0.000    638.054    800.506
##              Adult.Mortality    -0.491      0.061      -0.305    -
8.036    0.000    -0.612    -0.370
##              Hepatitis.B      0.873      0.206      0.133

```



```

4.236      0.000      0.465      1.281
## -----
##
##
##
## No more variables to be added.
##
## Variables Entered:
##
## + Income.composition.of.resources
## + Adult.Mortality
## + Hepatitis.B
##
##
## Final Model Output
## -----
##
##                               Model Summary
## -----
## R                               0.942          RMSE                54.362
## R-Squared                       0.888          Coef. Var          6.415
## Adj. R-Squared                   0.885          MSE                2955.267
## Pred R-Squared                   0.877          MAE                42.505
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF          Mean Square          F          Sig.
## -----
## Regression      2881206.757              3          960402.252          324.98      0.0000
## Residual        363497.796             123           2955.267
## Total          3244704.553             126
## -----
##
##                               Parameter Estimates
## -----
##                               t          Sig          lower          model          Beta          Std. Error          Std. Beta
##                               t          Sig          lower          upper
## -----
##                               (Intercept)          369.617          36.136
## 10.228      0.000          298.088          441.146
## Income.composition.of.resources          719.280          41.035          0.680
## 17.529      0.000          638.054          800.506

```

```
##          Adult.Mortality    -0.491          0.061          -0.305    -
8.036      0.000      -0.612      -0.370
##          Hepatitis.B        0.873          0.206          0.133
4.236      0.000          0.465          1.281
## -----
```

```
life2015_forwardmodel
```

```
##
##                               Selection Summary
## -----
```

```
##          Variable
## Step          Entered          R-Square    Adj.          C(p)
AIC          RMSE
## -----
```

```
##    1    Income.composition.of.resources    0.8116    0.8101
110.5088    1443.2457    69.9266
##    2    Adult.Mortality    0.8716    0.8696
38.1327    1396.5436    57.9582
##    3    Hepatitis.B    0.8880    0.8852
19.8706    1381.2467    54.3624
## -----
```

```
#Life2015_forwardmodel$mallows_cp
```

```
paste("Stepwise procedure with FORWARD indicated to consider only Income,
Adult.deaths and Hepatitis.B.")
```

```
## [1] "Stepwise procedure with FORWARD indicated to consider only Income,
Adult.deaths and Hepatitis.B."
```

```
# (Q.3.3) BACKWARD
```

```
life2015_backwardmodel = ols_step_backward_p(life2015_box_modelQ, prem =
0.05, details = TRUE)
```

```
## Backward Elimination Method
```

```
## -----
```

```
##
```

```
## Candidate Terms:
```

```
##
```

```
## 1 . Income.composition.of.resources
```

```
## 2 . Adult.Mortality
```

```
## 3 . HIV.AIDS
```

```
## 4 . Hepatitis.B
```

```
## 5 . Adult.Mortality:HIV.AIDS
```

```
##
```

```
## We are eliminating variables based on p value...
```

```
##
```

```

##
## No more variables satisfy the condition of p value = 0.05
##
##
## Variables Removed:
##
##
##
## Final Model Output
## -----
##
##                               Model Summary
## -----
## R                               0.950           RMSE                51.162
## R-Squared                       0.902           Coef. Var          6.037
## Adj. R-Squared                  0.898           MSE                2617.528
## Pred R-Squared                  0.875           MAE                 39.538
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      2927983.668           5      585596.734      223.721      0.0000
## Residual        316720.885          121        2617.528
## Total          3244704.553          126
## -----
##
##                               Parameter Estimates
## -----
##
##                               model      Beta      Std. Error      Std. Beta
## t      Sig      lower      upper
## -----
##                               (Intercept)      450.885      39.340
## 11.461      0.000      373.000      528.769
## Income.composition.of.resources      671.010      40.340      0.635
## 16.634      0.000      591.147      750.874
## Adult.Mortality      -0.540      0.074      -0.336      -
## 7.287      0.000      -0.687      -0.394
## HIV.AIDS      -40.777      9.673      -0.390      -
## 4.216      0.000      -59.926      -21.627
## Hepatitis.B      0.509      0.214      0.077
## 2.383      0.019      0.086      0.932
## Adult.Mortality:HIV.AIDS      0.095      0.026      0.348

```

```

3.689      0.000      0.044      0.147
## -----
-----

life2015_backwardmodel

## [1] "No variables have been removed from the model."

#life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD indicated to consider all the
selected variables.")

## [1] "Stepwise procedure with BACKWARD indicated to consider all the
selected variables."

# (Q.4) BEST SUBSET
#OPTION 1
ks = ols_step_best_subset(life2015_box_modelQ, details = TRUE)

#Summary of Selected model based on cp, aic, AdjustedR2
rsquare <- c(ks$rsq)
cp <- c(ks$cp)
aic <- c(ks$aic)
AdjustedR2 <- c(ks$adjr)
cbind(rsquare, cp, aic, AdjustedR2)

##          rsquare          cp          aic AdjustedR2
## [1,] 0.8116267 110.508815 1443.246 0.8101197
## [2,] 0.8716264 38.132700 1396.544 0.8695559
## [3,] 0.8879720 19.870644 1381.247 0.8852396
## [4,] 0.8978055 9.680962 1371.579 0.8944549
## [5,] 0.9023884 6.000000 1367.752 0.8983548

#OPTION 2
best.subset <- regsubsets((((Life.expectancy^1.70707) - 1)/1.70707) ~
Income.composition.of.resources +
Adult.Mortality + HIV.AIDS + Hepatitis.B +
Adult.Mortality * HIV.AIDS,
data = life2015[-c(119, 53, 93), ], nv = 4)
summary(best.subset)

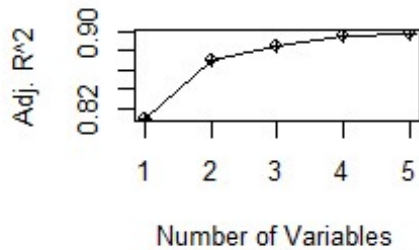
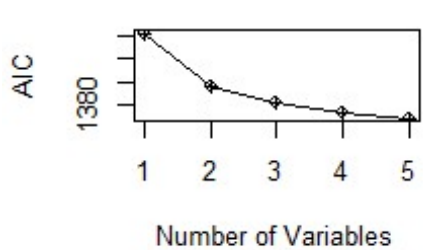
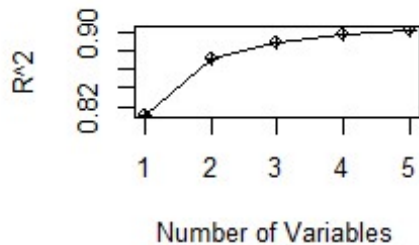
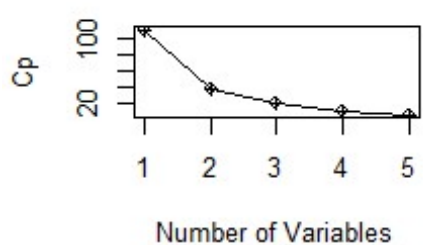
## Subset selection object
## Call: regsubsets.formula((((Life.expectancy^1.70707) - 1)/1.70707) ~
##      Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
##      Hepatitis.B + Adult.Mortality * HIV.AIDS, data = life2015[-c(119,
##      53, 93), ], nv = 4)
## 5 Variables (and intercept)
##
##              Forced in Forced out
## Income.composition.of.resources      FALSE      FALSE
## Adult.Mortality                     FALSE      FALSE
## HIV.AIDS                           FALSE      FALSE
## Hepatitis.B                         FALSE      FALSE

```

```
## Adult.Mortality:HIV.AIDS          FALSE      FALSE
## 1 subsets of each size up to 4
## Selection Algorithm: exhaustive
##           Income.composition.of.resources Adult.Mortality HIV.AIDS
Hepatitis.B
## 1 ( 1 ) "*"
## 2 ( 1 ) "*"
## 3 ( 1 ) "*"
## 4 ( 1 ) "*"
##           Adult.Mortality:HIV.AIDS
## 1 ( 1 ) " "
## 2 ( 1 ) " "
## 3 ( 1 ) " "
## 4 ( 1 ) "*"

reg.summary <- summary(best.subset)

#Plotting
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"R^2")
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"AIC")
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"Adj. R^2")
```



```

paste("Based on Marlow's Cp the minimum value was obtained with all the
variables.")

## [1] "Based on Marlow's Cp the minimum value was obtained with all the
variables."

#MODEL DIAGNOSTICS
# (Q.5) CHECKING LINEARITY
#Residuals plot
plot(life2015_box_modelQ, which = 1)

# a Scale Location Plot
plot(life2015_box_modelQ, which = 3)

paste("It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected.")

## [1] "It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected."

# (Q.6) Heteroscedasticity Test - the Breush-Pagan test
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
# Ha: heteroscedasticity is presented
bptest(life2015_box_modelQ)

##
## studentized Breusch-Pagan test
##
## data:  life2015_box_modelQ
## BP = 12.418, df = 5, p-value = 0.02948

paste("The output displays the Breush-Pagan test that from the model
presented the p-value 0.02948 > 0.05, indicating the we should NOT REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity does not exist.")

## [1] "The output displays the Breush-Pagan test that from the model
presented the p-value 0.02948 > 0.05, indicating the we should NOT REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity does not exist."

# (Q.7) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_modelQ))

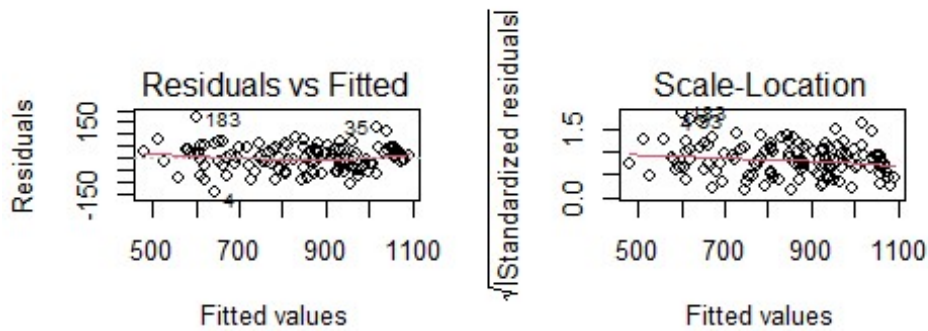
```

```
##
## Shapiro-Wilk normality test
##
## data: residuals(life2015_box_modelQ)
## W = 0.99142, p-value = 0.6263

paste("Shapiro-Wilk normality test presented the p-value = 0.6263 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level.")

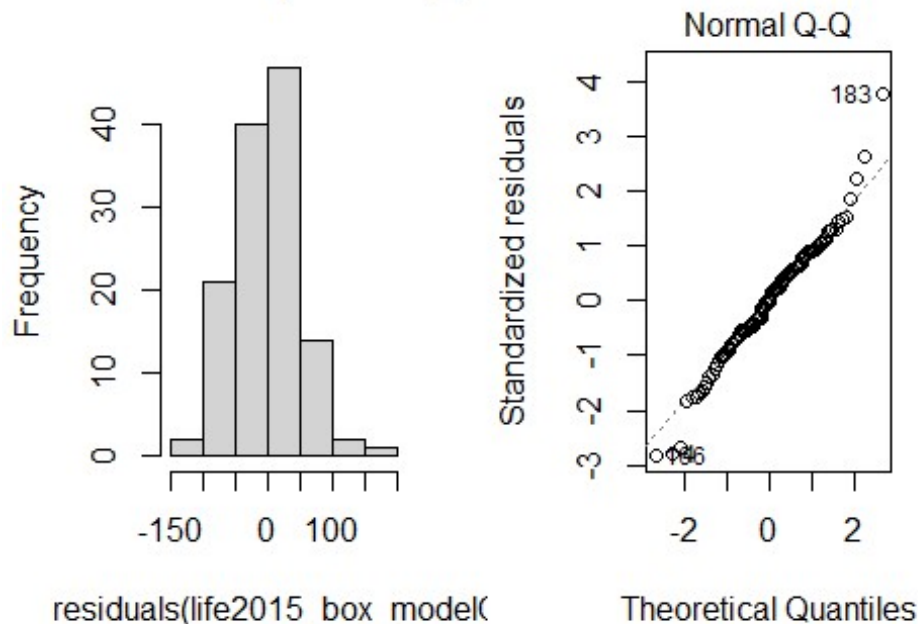
## [1] "Shapiro-Wilk normality test presented the p-value = 0.6263 > 0.05
that indicated that we should NOT REJECT the null hypothesis and conclude
that the residuals are normally distributed at 5 % level."

#Plots
par(mfrow = c(1,2))
```



```
hist(residuals(life2015_box_modelQ))
plot(life2015_box_modelM, which = 2)
```

am of residuals(life2015_b



```
paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")
```

```
## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."
```

```
# (Q.8) Multicollinearity test
```

```
#VIF
```

```
imcdiag(life2015_box_modelQ, method = "VIF")
```

```
##
```

```
## Call:
```

```
## imcdiag(mod = life2015_box_modelQ, method = "VIF")
```

```
##
```

```
##
```

```
## VIF Multicollinearity Diagnostics
```

```
##
```

```
##
```

	VIF	detection
Income.composition.of.resources	1.8039	0
Adult.Mortality	2.6394	0
HIV.AIDS	10.6188	1
Hepatitis.B	1.3045	0
Adult.Mortality:HIV.AIDS	11.0312	1

```
##
```

```
## Multicollinearity may be due to HIV.AIDS Adult.Mortality:HIV.AIDS
regressors
```

```
##
```



```

## 1 --> COLLINEARITY is detected by the test
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. ")

## [1] "Multicollinearity verified between HIV.AIDS and the interaction term
Adult.deaths * HIV.AIDS. "

# (Q.9) LEVERAGE OF THE POINTS
# Leverage Points
lev = hatvalues(life2015_box_modelQ)
p = length(coef(life2015_box_modelQ))
n = nrow(life2015)
outlier = lev[lev > (2 * p/n)]
print(outlier)

##          26          53          60          93          111          118          126
127
## 0.1235762 0.3728366 0.1006973 0.5974663 0.1027386 0.1539996 0.1163033
0.1316548
##          133          144          156          166          182          183
## 0.1225211 0.1083070 0.1943514 0.1287227 0.3540296 0.1359353

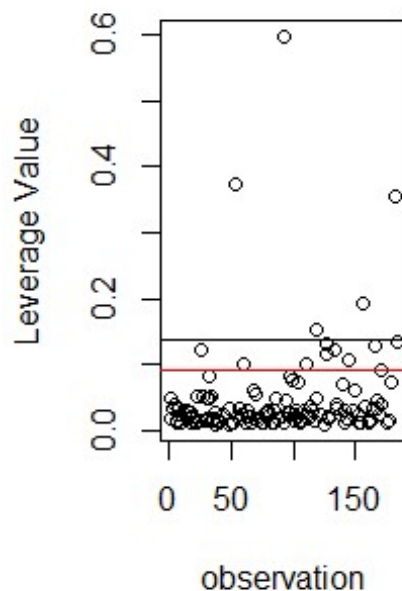
plot(rownames(life2015[-c(119, 53, 93), ]), lev, main = "Leverage in Life2015
Dataset",
      xlab = "observation", ylab = "Leverage Value")
abline(h = 2 * p/n, lty = 1, col = "red")
abline(h = 3 * p/n, lty = 1)

paste("The results shows that that the data points #26, #53, #60, #93, #111,
#118, #126, #127, #166, #182and #183 presented leverage values higher than
2.p/n that might be outliers.")

## [1] "The results shows that that the data points #26, #53, #60, #93, #111,
#118, #126, #127, #166, #182and #183 presented leverage values higher than
2.p/n that might be outliers."

```

Leverage in Life2015 Data

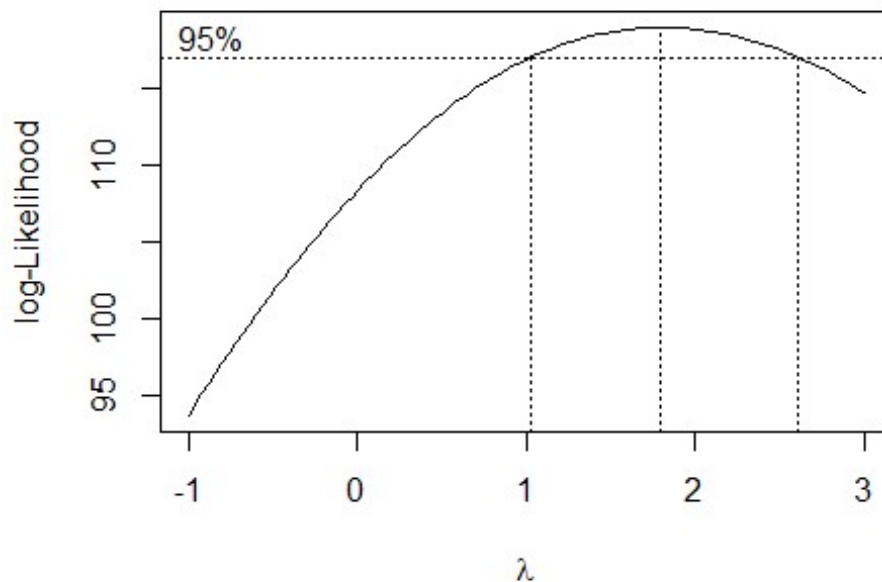


```
##### MODEL R - BOX COX TRANSFORMATION of DANIEL's model
#####
paste("t-test of previous Box-Cox transformation indicated no significance of
interaction term after this transformation")

## [1] "t-test of previous Box-Cox transformation indicated no significance
of interaction term after this transformation"

#(R.1) BOX-COX TRANSFORMATION
# The necessity of transformation is due to the indication of
heterosdasticity in model E
# For this transformation we will be considering only the significant
parameters obtained beforehand
life2015_reduced_modelR <- lm(Life.expectancy ~
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
Hepatitis.B + Adult.Mortality * Hepatitis.B,
data = life2015)

bc = boxcox(life2015_reduced_modelR, lambda = seq(-1,3))
```



```
#Extract best Lambda
bestlambda = bc$x[which(bc$y == max(bc$y))]
paste("The best lambda for Box-Cox Transformation is: ", bestlambda)

## [1] "The best lambda for Box-Cox Transformation is:  1.78787878787879"

paste("From the output, as the best lambda would be 1.78787.")

## [1] "From the output, as the best lambda would be 1.78787."

#BOX-COX with LAMBDA = 1.70707
life2015_box_modelR = lm((((Life.expectancy^1.78787) - 1)/1.78787) ~
Income.composition.of.resources +
                        Adult.Mortality + HIV.AIDS + Hepatitis.B +
                        Adult.Mortality * Hepatitis.B,
                        data = life2015)

# (R.2) PARTIAL TEST - Individual Coefficients Test (t-test)
# Ho : Beta(i) = 0
# Ha : Beta(i) ≠ 0 (i = 1, 2, ..., p)

life2015_box_modelR = lm((((Life.expectancy^1.78787) - 1)/1.78787) ~
Income.composition.of.resources +
                        Adult.Mortality + HIV.AIDS + Hepatitis.B +
                        Adult.Mortality * Hepatitis.B, data = life2015[-
c(119,53,93),])
```

```
summary(life2015_box_modelR)

##
## Call:
## lm(formula = (((Life.expectancy^1.78787) - 1)/1.78787) ~
Income.composition.of.resources +
##      Adult.Mortality + HIV.AIDS + Hepatitis.B + Adult.Mortality *
##      Hepatitis.B, data = life2015[-c(119, 53, 93), ])
##
## Residuals:
##      Min        1Q    Median        3Q        Max
## -208.919  -45.093    1.978    50.646   234.400
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      392.661654   55.300722    7.100 9.24e-11 ***
## Income.composition.of.resources 982.971792   55.356994   17.757 < 2e-16 ***
## Adult.Mortality      -0.010246    0.191137   -0.054 0.957337
## HIV.AIDS            -10.724531    5.795859   -1.850 0.066698 .
## Hepatitis.B          2.485340    0.502321    4.948 2.45e-06 ***
## Adult.Mortality:Hepatitis.B    -0.008320    0.002375   -3.504 0.000644 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 72.55 on 121 degrees of freedom
## Multiple R-squared:  0.9011, Adjusted R-squared:  0.897
## F-statistic: 220.4 on 5 and 121 DF,  p-value: < 2.2e-16

paste("The t-test indicated that after the Box-Cox transformation only
Adult.Mortality with p-value > 0.05 that indicates that this variable might
not have significant influence in Life Expectancy at 5 % level.")

## [1] "The t-test indicated that after the Box-Cox transformation only
Adult.Mortality with p-value > 0.05 that indicates that this variable might
not have significant influence in Life Expectancy at 5 % level."

#confidence interval for the coefficients
confint(life2015_box_modelQ, level = 0.95)

##              2.5 %        97.5 %
## (Intercept)      373.00043645  528.7686394
## Income.composition.of.resources 591.14692861  750.8737869
## Adult.Mortality      -0.68726236   -0.3935969
## HIV.AIDS            -59.92646782 -21.6272619
## Hepatitis.B          0.08620761    0.9317125
## Adult.Mortality:HIV.AIDS      0.04419529    0.1465805

paste("For a 95% confidence interval, under.five.deaths with Zero between
upper and lower values!")
```

```
## [1] "For a 95% confidence interval, under.five.deaths with Zero between  
upper and lower values!"
```

(R.3) STEPWISE REGRESSION PROCEDURE

(Q.3.1) BOTH

```
life2015_bothmodel = ols_step_both_p(life2015_box_modelR, pent = 0.05, prem = 0.05, details = TRUE)
```

Stepwise Selection Method

##

Candidate Terms:

##

1. Income.composition.of.resources

2. Adult.Mortality

3. HIV.AIDS

4. Hepatitis.B

```
## 5. Adult.Mortality:Hepatitis.B
```

##

```
## We are selecting variables based on p value...
```

##

##

```
## Stepwise Selection: Step 1
```

##

```
## - Income.composition.of.resources added
```

##

Model Summary

##

## R	0.901	RMSE	98.363
------	-------	------	--------

## R-Squared	0.812	Coef. Var	8.604
--------------	-------	-----------	-------

## Adj. R-Squared	0.811	MSE	9675.272
-------------------	-------	-----	----------

##	Pred	R-Squared	0.806	MAE	77.442
----	------	-----------	-------	-----	--------

```
## RMSE: Root Mean Square Error
```

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

##

##	Sum of				
##	Squares	DF	Mean Square	F	Sig.

## Regression	5228792.054	1	5228792.054	540.428	0.0000
---------------	-------------	---	-------------	---------	--------

## Residual	1209409.006	125	9675.272
-------------	-------------	-----	----------

```
## Total      6438201.060      126
```

##

```
##                                     Parameter Estimates
```

```

##          Sig          lower      model      Beta      Std. Error      Std. Beta
t          Sig          lower      upper
## -----
##          (Intercept)      237.031      39.945
5.934      0.000      157.974      316.088
## Income.composition.of.resources      1342.406      57.745      0.901
23.247      0.000      1228.122      1456.691
## -----
##
##
##
## Stepwise Selection: Step 2
##
## - Adult.Mortality added
##
##          Model Summary
## -----
## R          0.934      RMSE          81.665
## R-Squared      0.872      Coef. Var      7.143
## Adj. R-Squared      0.869      MSE          6669.126
## Pred R-Squared      0.864      MAE          61.079
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##          ANOVA
## -----
##          Sum of
##          Squares      DF      Mean Square      F      Sig.
## -----
## Regression      5611229.416      2      2805614.708      420.687      0.0000
## Residual      826971.644      124      6669.126
## Total      6438201.060      126
## -----
##
##          Parameter Estimates
## -----
##          Sig          lower      model      Beta      Std. Error      Std. Beta
t          Sig          lower      upper
## -----
##          (Intercept)      535.883      51.549
10.396      0.000      433.852      637.914
## Income.composition.of.resources      1064.638      60.365      0.715
17.637      0.000      945.159      1184.117
## Adult.Mortality      -0.695      0.092      -0.307

```

```

-7.573    0.000    -0.876    -0.513
## -----
##
##
##
##              Model Summary
## -----
## R                0.934          RMSE                81.665
## R-Squared        0.872          Coef. Var            7.143
## Adj. R-Squared   0.869          MSE                6669.126
## Pred R-Squared   0.864          MAE                61.079
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##              ANOVA
## -----
##              Sum of
##              Squares          DF          Mean Square          F          Sig.
## -----
## Regression      5611229.416          2          2805614.708          420.687          0.0000
## Residual        826971.644          124           6669.126
## Total          6438201.060          126
## -----
##
##              Parameter Estimates
## -----
##
##              Sig          lower          model          Beta          Std. Error          Std. Beta
##              t          Sig          lower          upper
## -----
##              (Intercept)          535.883          51.549
## 10.396    0.000    433.852    637.914
## Income.composition.of.resources          1064.638          60.365          0.715
## 17.637    0.000    945.159    1184.117
## Adult.Mortality          -0.695          0.092          -0.307
## -7.573    0.000    -0.876    -0.513
## -----
##
##
##
## Stepwise Selection: Step 3
##
## - Hepatitis.B added
##
##              Model Summary

```

```

## -----
## R                0.942      RMSE                76.643
## R-Squared        0.888      Coef. Var            6.704
## Adj. R-Squared   0.885      MSE                5874.093
## Pred R-Squared   0.877      MAE                59.997
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                Sum of
##                Squares      DF      Mean Square      F      Sig.
## -----
## Regression      5715687.680      3      1905229.227      324.344      0.0000
## Residual        722513.379     123        5874.093
## Total          6438201.060     126
## -----
##
##                               Parameter Estimates
## -----
## -----
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)      468.546      50.946
## 9.197      0.000      367.701      569.392
## Income.composition.of.resources      1015.207      57.853      0.682
## 17.548      0.000      900.691      1129.723
## Adult.Mortality      -0.688      0.086      -0.304
## -7.989      0.000      -0.859      -0.518
## Hepatitis.B      1.225      0.291      0.132
## 4.217      0.000      0.650      1.800
## -----
##
##
##
##                               Model Summary
## -----
## R                0.942      RMSE                76.643
## R-Squared        0.888      Coef. Var            6.704
## Adj. R-Squared   0.885      MSE                5874.093
## Pred R-Squared   0.877      MAE                59.997
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error

```



```

##
##
## ANOVA
## -----
## Sum of
## Squares DF Mean Square F Sig.
## -----
## Regression 5715687.680 3 1905229.227 324.344 0.0000
## Residual 722513.379 123 5874.093
## Total 6438201.060 126
## -----
##
## Parameter Estimates
## -----
##
## Sig lower model upper Beta Std. Error Std. Beta
## t -----
##
## (Intercept) 468.546 50.946
9.197 0.000 367.701 569.392
## Income.composition.of.resources 1015.207 57.853 0.682
17.548 0.000 900.691 1129.723
## Adult.Mortality -0.688 0.086 -0.304
-7.989 0.000 -0.859 -0.518
## Hepatitis.B 1.225 0.291 0.132
4.217 0.000 0.650 1.800
## -----
##
##
##
## Stepwise Selection: Step 4
##
## - Adult.Mortality:Hepatitis.B added
##
## Model Summary
## -----
## R 0.948 RMSE 73.268
## R-Squared 0.898 Coef. Var 6.409
## Adj. R-Squared 0.895 MSE 5368.165
## Pred R-Squared 0.886 MAE 57.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of
## Squares DF Mean Square F Sig.

```

##						
## Regression		5783284.916	4	1445821.229	269.332	0.0000
## Residual		654916.144	122	5368.165		
## Total		6438201.060	126			
## -----						
##						
##						
## Parameter Estimates						
## -----						
##						
##			model	Beta	Std. Error	Std. Beta
t	Sig	lower	upper			
## -----						
##						
		(Intercept)		376.653	55.160	
6.828	0.000	267.459	485.847			
## Income.composition.of.resources				991.729	55.700	0.666
17.805	0.000	881.467	1101.992			
## Adult.Mortality				-0.086	0.189	-0.038
0.458	0.648	-0.459	0.287			
## Hepatitis.B				2.682	0.496	0.289
5.411	0.000	1.701	3.664			
## Adult.Mortality:Hepatitis.B				-0.009	0.002	-0.309
3.549	0.001	-0.013	-0.004			
## -----						
##						
##						
##						
## Model Summary						
## -----						
## R		0.948		RMSE	73.268	
## R-Squared		0.898		Coef. Var	6.409	
## Adj. R-Squared		0.895		MSE	5368.165	
## Pred R-Squared		0.886		MAE	57.090	
## -----						
## RMSE: Root Mean Square Error						
## MSE: Mean Square Error						
## MAE: Mean Absolute Error						
##						
## ANOVA						
## -----						
		Sum of				
		Squares	DF	Mean Square	F	Sig.
## -----						
## Regression		5783284.916	4	1445821.229	269.332	0.0000
## Residual		654916.144	122	5368.165		
## Total		6438201.060	126			
## -----						
##						
##						
## Parameter Estimates						

```

## -----
##
##          Sig          lower      model      Beta      Std. Error      Std. Beta
##          t          upper
## -----
##
##          (Intercept)      376.653      55.160
6.828      0.000      267.459      485.847
## Income.composition.of.resources      991.729      55.700      0.666
17.805      0.000      881.467      1101.992
##          Adult.Mortality      -0.086      0.189      -0.038      -
0.458      0.648      -0.459      0.287
##          Hepatitis.B      2.682      0.496      0.289
5.411      0.000      1.701      3.664
##          Adult.Mortality:Hepatitis.B      -0.009      0.002      -0.309      -
3.549      0.001      -0.013      -0.004
## -----
##
##
##
## Stepwise Selection: Step 5
##
## - HIV.AIDS added
##
##          Model Summary
## -----
## R          0.949      RMSE          72.551
## R-Squared      0.901      Coef. Var      6.346
## Adj. R-Squared      0.897      MSE          5263.588
## Pred R-Squared      0.886      MAE          56.257
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##          ANOVA
## -----
##          Sum of
##          Squares      DF      Mean Square      F      Sig.
## -----
## Regression      5801306.908      5      1160261.382      220.432      0.0000
## Residual      636894.151      121      5263.588
## Total      6438201.060      126
## -----
##
##          Parameter Estimates
## -----
##          model      Beta      Std. Error      Std. Beta

```

t	Sig	lower	upper
## -----			
## (Intercept) 392.662 55.301			
7.100	0.000	283.179	502.144
## Income.composition.of.resources 982.972 55.357 0.660			
17.757	0.000	873.378	1092.566
## Adult.Mortality -0.010 0.191 -0.005 -			
0.054	0.957	-0.389	0.368
## Hepatitis.B 2.485 0.502 0.268			
4.948	0.000	1.491	3.480
## HIV.AIDS -10.725 5.796 -0.073 -			
1.850	0.067	-22.199	0.750
## Adult.Mortality:Hepatitis.B -0.008 0.002 -0.302 -			
3.504	0.001	-0.013	-0.004
## -----			
##			
##			
##			
## Model Summary			
## -----			
## R		0.949	RMSE 72.551
## R-Squared		0.901	Coef. Var 6.346
## Adj. R-Squared		0.897	MSE 5263.588
## Pred R-Squared		0.886	MAE 56.257
## -----			
## RMSE: Root Mean Square Error			
## MSE: Mean Square Error			
## MAE: Mean Absolute Error			
##			
## ANOVA			
## -----			
##		Sum of	
##		Squares	DF Mean Square F Sig.
## -----			
## Regression		5801306.908	5 1160261.382 220.432 0.0000
## Residual		636894.151	121 5263.588
## Total		6438201.060	126
## -----			
##			
## Parameter Estimates			
## -----			
##			model
t	Sig	lower	upper Beta Std. Error Std. Beta
## -----			
## (Intercept) 392.662 55.301			
7.100	0.000	283.179	502.144

```

## Income.composition.of.resources    982.972    55.357    0.660
17.757    0.000    873.378    1092.566
##
##      Adult.Mortality    -0.010    0.191    -0.005    -
0.054    0.957    -0.389    0.368
##
##      Hepatitis.B    2.485    0.502    0.268
4.948    0.000    1.491    3.480
##
##      HIV.AIDS    -10.725    5.796    -0.073    -
1.850    0.067    -22.199    0.750
##      Adult.Mortality:Hepatitis.B    -0.008    0.002    -0.302    -
3.504    0.001    -0.013    -0.004
## -----
##
##
##
## Stepwise Selection: Step 6
##
## - Adult.Mortality:Hepatitis.B added
##
##
##      Model Summary
## -----
## R    0.944    RMSE    75.829
## R-Squared    0.891    Coef. Var    6.633
## Adj. R-Squared    0.887    MSE    5750.031
## Pred R-Squared    0.876    MAE    58.946
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##
##      ANOVA
## -----
##
##      Sum of
##      Squares    DF    Mean Square    F    Sig.
## -----
## Regression    5736697.264    4    1434174.316    249.42    0.0000
## Residual    701503.795    122    5750.031
## Total    6438201.060    126
## -----
##
##
##      Parameter Estimates
## -----
##
##
##      Sig    lower    model    Beta    Std. Error    Std. Beta
##      t
## -----
##
##      (Intercept)    483.685    51.024
9.480    0.000    382.678    584.692
## Income.composition.of.resources    1005.215    57.477    0.675

```

```

17.489    0.000    891.434    1118.996
##          Adult.Mortality    -0.592    0.099    -0.262
-5.987    0.000    -0.788    -0.396
##          Hepatitis.B    1.046    0.302    0.113
3.462    0.001    0.448    1.645
##          HIV.AIDS    -11.569    6.053    -0.079
-1.911    0.058    -23.551    0.412
## -----
##
##
##
##
## Final Model Output
## -----
##
##                      Model Summary
## -----
## R                      0.944    RMSE                      75.829
## R-Squared              0.891    Coef. Var                  6.633
## Adj. R-Squared         0.887    MSE                      5750.031
## Pred R-Squared         0.876    MAE                      58.946
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                      ANOVA
## -----
##          Sum of
##          Squares          DF    Mean Square          F          Sig.
## -----
## Regression    5736697.264          4    1434174.316    249.42    0.0000
## Residual      701503.795         122      5750.031
## Total         6438201.060         126
## -----
##
##                      Parameter Estimates
## -----
##
##          Sig          lower          model          Beta    Std. Error    Std. Beta
##          t          upper
## -----
##          (Intercept)    483.685    51.024
9.480    0.000    382.678    584.692
## Income.composition.of.resources    1005.215    57.477    0.675
17.489    0.000    891.434    1118.996
##          Adult.Mortality    -0.592    0.099    -0.262
-5.987    0.000    -0.788    -0.396

```

```

##              Hepatitis.B          1.046          0.302          0.113
3.462      0.001      0.448      1.645
##              HIV.AIDS          -11.569          6.053          -0.079
-1.911      0.058      -23.551      0.412
## -----
-----

life2015_bothmodel

##
##                                     Stepwise Selection Summary
## -----
-----

##                                     Added/      Adj.
## Step                               Removed      R-Square      R-
## Square                               C(p)          AIC          RMSE
## -----
-----

##      1      Income.composition.of.resources      addition      0.812
0.811      106.7690      1529.9152      98.3630
##      2              Adult.Mortality      addition      0.872
0.869      36.1120      1483.6404      81.6647
##      3              Hepatitis.B      addition      0.888
0.885      18.2660      1468.4910      76.6426
##      4      Adult.Mortality:Hepatitis.B      addition      0.898
0.895      7.4240      1458.0160      73.2678
##      5              HIV.AIDS      addition      0.901
0.897      6.0000      1456.4722      72.5506
##      6      Adult.Mortality:Hepatitis.B      removal      0.891
0.887      16.2750      1466.7433      75.8290
## -----
-----

#life2015_bothmodel$mallows_cp
paste("Stepwise procedure with BOTH indicated to consider all the selected
variables.")

## [1] "Stepwise procedure with BOTH indicated to consider all the selected
variables."

# (R.3.2) FORWARD
life2015_forwardmodel = ols_step_forward_p(life2015_box_modelR, pent = 0.05,
details = TRUE)

## Forward Selection Method
## -----
##
## Candidate Terms:
##
## 1. Income.composition.of.resources
## 2. Adult.Mortality

```

```

## 3. HIV.AIDS
## 4. Hepatitis.B
## 5. Adult.Mortality:Hepatitis.B
##
## We are selecting variables based on p value...
##
## Forward Selection: Step 1
##
## - Income.composition.of.resources
##
##                               Model Summary
## -----
## R                               0.901          RMSE                98.363
## R-Squared                       0.812          Coef. Var          8.604
## Adj. R-Squared                   0.811          MSE                9675.272
## Pred R-Squared                   0.806          MAE                77.442
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares          DF      Mean Square          F          Sig.
## -----
## Regression      5228792.054              1      5228792.054      540.428      0.0000
## Residual        1209409.006             125           9675.272
## Total           6438201.060             126
## -----
##
##                               Parameter Estimates
## -----
##                               -----
##                               model      Beta      Std. Error      Std. Beta
##                               t      Sig      lower      upper
## -----
##                               (Intercept)      237.031      39.945
## 5.934      0.000      157.974      316.088
## Income.composition.of.resources      1342.406      57.745      0.901
## 23.247      0.000      1228.122      1456.691
## -----
##
##
##
## Forward Selection: Step 2
##

```


- Adult.Mortality

##

Model Summary

```
## -----
## R                0.934          RMSE                81.665
## R-Squared        0.872          Coef. Var            7.143
## Adj. R-Squared   0.869          MSE                6669.126
## Pred R-Squared   0.864          MAE                61.079
## -----
```

RMSE: Root Mean Square Error

MSE: Mean Square Error

MAE: Mean Absolute Error

##

ANOVA

```
## -----
##                Sum of
##                Squares      DF      Mean Square      F      Sig.
## -----
## Regression    5611229.416      2      2805614.708    420.687    0.0000
## Residual      826971.644     124        6669.126
## Total        6438201.060     126
## -----
```

##

Parameter Estimates

```
## -----
##                Sig      lower      model      Beta      Std. Error      Std. Beta
## t                Sig      lower      upper
## -----
##                (Intercept)      535.883      51.549
10.396    0.000    433.852    637.914
## Income.composition.of.resources      1064.638      60.365      0.715
17.637    0.000    945.159    1184.117
## Adult.Mortality      -0.695      0.092      -0.307
-7.573    0.000    -0.876    -0.513
## -----
```

##

##

##

Forward Selection: Step 3

##

- Hepatitis.B

##

Model Summary

```
## -----
## R                0.942          RMSE                76.643
## R-Squared        0.888          Coef. Var            6.704
## Adj. R-Squared   0.885          MSE                5874.093
```

```

## Pred R-Squared      0.877      MAE      59.997
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
## ANOVA
## -----
## Sum of
## Squares      DF      Mean Square      F      Sig.
## -----
## Regression      5715687.680      3      1905229.227      324.344      0.0000
## Residual      722513.379      123      5874.093
## Total      6438201.060      126
## -----
##
## Parameter Estimates
## -----
##
## Sig      lower      model      Beta      Std. Error      Std. Beta
## t      upper
## -----
## (Intercept)      468.546      50.946
9.197      0.000      367.701      569.392
## Income.composition.of.resources      1015.207      57.853      0.682
17.548      0.000      900.691      1129.723
## Adult.Mortality      -0.688      0.086      -0.304
-7.989      0.000      -0.859      -0.518
## Hepatitis.B      1.225      0.291      0.132
4.217      0.000      0.650      1.800
## -----
##
##
##
## Forward Selection: Step 4
##
## - Adult.Mortality:Hepatitis.B
##
## Model Summary
## -----
## R      0.948      RMSE      73.268
## R-Squared      0.898      Coef. Var      6.409
## Adj. R-Squared      0.895      MSE      5368.165
## Pred R-Squared      0.886      MAE      57.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error

```

```

##
##                                ANOVA
## -----
##              Sum of
##              Squares          DF      Mean Square          F          Sig.
## -----
## Regression      5783284.916          4      1445821.229      269.332      0.0000
## Residual        654916.144         122          5368.165
## Total          6438201.060         126
## -----
##
##                                Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)      376.653          55.160
## 6.828      0.000      267.459      485.847
## ## Income.composition.of.resources      991.729          55.700          0.666
## 17.805      0.000      881.467      1101.992
## ## Adult.Mortality      -0.086          0.189      -0.038      -
## 0.458      0.648      -0.459          0.287
## ## Hepatitis.B      2.682          0.496          0.289
## 5.411      0.000          1.701          3.664
## ## Adult.Mortality:Hepatitis.B      -0.009          0.002      -0.309      -
## 3.549      0.001      -0.013      -0.004
## -----
##
##
##
## Forward Selection: Step 5
##
## - HIV.AIDS
##
##                                Model Summary
## -----
## R              0.949          RMSE              72.551
## R-Squared        0.901          Coef. Var          6.346
## Adj. R-Squared    0.897          MSE              5263.588
## Pred R-Squared    0.886          MAE              56.257
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                                ANOVA
## -----

```

```

##              Sum of
##              Squares      DF      Mean Square      F      Sig.
## -----
## Regression    5801306.908        5    1160261.382    220.432    0.0000
## Residual      636894.151       121      5263.588
## Total         6438201.060       126
## -----
##
##                                     Parameter Estimates
## -----
##
##              Sig      lower      model      Beta      Std. Error      Std. Beta
##              t              upper
## -----
##              (Intercept)      392.662      55.301
## 7.100      0.000      283.179      502.144
## Income.composition.of.resources      982.972      55.357      0.660
## 17.757      0.000      873.378      1092.566
## Adult.Mortality      -0.010      0.191      -0.005      -
## 0.054      0.957      -0.389      0.368
## Hepatitis.B      2.485      0.502      0.268
## 4.948      0.000      1.491      3.480
## HIV.AIDS      -10.725      5.796      -0.073      -
## 1.850      0.067      -22.199      0.750
## Adult.Mortality:Hepatitis.B      -0.008      0.002      -0.302      -
## 3.504      0.001      -0.013      -0.004
## -----
##
##
##
##
## Variables Entered:
##
## + Income.composition.of.resources
## + Adult.Mortality
## + Hepatitis.B
## + Adult.Mortality:Hepatitis.B
## + HIV.AIDS
##
##
## Final Model Output
## -----
##
##                                     Model Summary
## -----
## R      0.949      RMSE      72.551
## R-Squared      0.901      Coef. Var      6.346
## Adj. R-Squared      0.897      MSE      5263.588

```

```

## Pred R-Squared      0.886      MAE      56.257
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      5801306.908      5      1160261.382      220.432      0.0000
## Residual        636894.151      121      5263.588
## Total           6438201.060      126
## -----
##
##                               Parameter Estimates
## -----
##
##                               Beta      Std. Error      Std. Beta
## t      Sig      lower      model      upper
## -----
##                               (Intercept)      392.662      55.301
## 7.100      0.000      283.179      502.144
## Income.composition.of.resources      982.972      55.357      0.660
## 17.757      0.000      873.378      1092.566
## Adult.Mortality      -0.010      0.191      -0.005      -
## 0.054      0.957      -0.389      0.368
## Hepatitis.B      2.485      0.502      0.268
## 4.948      0.000      1.491      3.480
## HIV.AIDS      -10.725      5.796      -0.073      -
## 1.850      0.067      -22.199      0.750
## Adult.Mortality:Hepatitis.B      -0.008      0.002      -0.302      -
## 3.504      0.001      -0.013      -0.004
## -----
## -----
life2015_forwardmodel
##
##                               Selection Summary
## -----
## -----
##                               Variable
## Step      Entered      R-Square      Adj.      C(p)
## AIC      RMSE
## -----
## 1      Income.composition.of.resources      0.8122      0.8106

```

```

106.7689    1529.9152    98.3630
##      2      Adult.Mortality                0.8716    0.8695
36.1118    1483.6404    81.6647
##      3      Hepatitis.B                  0.8878    0.8850
18.2663    1468.4910    76.6426
##      4      Adult.Mortality:Hepatitis.B    0.8983    0.8949
7.4239     1458.0160    73.2678
##      5      HIV.AIDS                     0.9011    0.8970
6.0000     1456.4722    72.5506
## -----
-----

#life2015_forwardmodel$mallows_cp
paste("Stepwise procedure with FORWARD indicated to consider all the selected
variables.")

## [1] "Stepwise procedure with FORWARD indicated to consider all the
selected variables."

# (R.3.3) BACKWARD
life2015_backwardmodel = ols_step_backward_p(life2015_box_modelR, prem =
0.05, details = TRUE)

## Backward Elimination Method
## -----
##
## Candidate Terms:
##
## 1 . Income.composition.of.resources
## 2 . Adult.Mortality
## 3 . HIV.AIDS
## 4 . Hepatitis.B
## 5 . Adult.Mortality:Hepatitis.B
##
## We are eliminating variables based on p value...
##
## - HIV.AIDS
##
## Backward Elimination: Step 1
##
## Variable HIV.AIDS Removed
##
##
##                               Model Summary
## -----
## R                               0.948      RMSE                73.268
## R-Squared                       0.898      Coef. Var           6.409
## Adj. R-Squared                   0.895      MSE                5368.165
## Pred R-Squared                   0.886      MAE                57.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error

```

MAE: Mean Absolute Error

##

ANOVA

	Sum of Squares	DF	Mean Square	F	Sig.
## Regression	5783284.916	4	1445821.229	269.332	0.0000
## Residual	654916.144	122	5368.165		
## Total	6438201.060	126			

##

Parameter Estimates

	Sig	lower	model upper	Beta	Std. Error	Std. Beta
--	-----	-------	----------------	------	------------	-----------

##			(Intercept)	376.653	55.160	
6.828	0.000	267.459	485.847			
## Income.composition.of.resources			991.729	55.700	0.666	
17.805	0.000	881.467	1101.992			
##			Adult.Mortality	-0.086	0.189	-0.038
0.458	0.648	-0.459	0.287			
##			Hepatitis.B	2.682	0.496	0.289
5.411	0.000	1.701	3.664			
##			Adult.Mortality:Hepatitis.B	-0.009	0.002	-0.309
3.549	0.001	-0.013	-0.004			

##

##

##

No more variables satisfy the condition of p value = 0.05

##

##

Variables Removed:

##

- HIV.AIDS

##

##

Final Model Output

##

Model Summary

## R	0.948	RMSE	73.268
## R-Squared	0.898	Coef. Var	6.409
## Adj. R-Squared	0.895	MSE	5368.165

```

## Pred R-Squared      0.886      MAE      57.090
## -----
## RMSE: Root Mean Square Error
## MSE: Mean Square Error
## MAE: Mean Absolute Error
##
##                               ANOVA
## -----
##                               Sum of
##                               Squares      DF      Mean Square      F      Sig.
## -----
## Regression      5783284.916      4      1445821.229      269.332      0.0000
## Residual        654916.144      122      5368.165
## Total           6438201.060      126
## -----
##
##                               Parameter Estimates
## -----
##
##                               Beta      Std. Error      Std. Beta
## t      Sig      lower      model      upper
## -----
##
##                               (Intercept)      376.653      55.160
## 6.828      0.000      267.459      485.847
## Income.composition.of.resources      991.729      55.700      0.666
## 17.805      0.000      881.467      1101.992
## Adult.Mortality      -0.086      0.189      -0.038      -
## 0.458      0.648      -0.459      0.287
## Hepatitis.B      2.682      0.496      0.289
## 5.411      0.000      1.701      3.664
## Adult.Mortality:Hepatitis.B      -0.009      0.002      -0.309      -
## 3.549      0.001      -0.013      -0.004
## -----
## -----
life2015_backwardmodel
##
##
##                               Elimination Summary
## -----
##                               Variable      Adj.
##                               Removed      R-Square      R-Square      C(p)      AIC      RMSE
## -----
## 1      HIV.AIDS      0.8983      0.8949      7.4239      1458.0160      73.2678
## -----

```



```

#life2015_backwardmodel$mallows_cp
paste("Stepwise procedure with BACKWARD indicated to remove
Adult.Mortality.")

## [1] "Stepwise procedure with BACKWARD indicated to remove
Adult.Mortality."

# (R.4) BEST SUBSET
#OPTION 1
ks = ols_step_best_subset(life2015_box_modelR, details = TRUE)

#Summary of Selected model based on cp, aic, AdjustedR2
rsquare <- c(ks$rsq)
cp <- c(ks$cp)
aic <- c(ks$aic)
AdjustedR2 <- c(ks$adjr)
cbind(rsquare, cp, aic, AdjustedR2)

##          rsquare          cp          aic AdjustedR2
## [1,] 0.8121511 106.768933 1529.915 0.8106483
## [2,] 0.8715524 36.111772 1483.640 0.8694806
## [3,] 0.8981020 5.637419 1456.234 0.8956166
## [4,] 0.9010734 4.002874 1454.475 0.8978299
## [5,] 0.9010758 6.000000 1456.472 0.8969880

#OPTION 2
best.subset <- regsubsets((((Life.expectancy^1.78787) - 1)/1.78787) ~
Income.composition.of.resources +
                        Adult.Mortality + HIV.AIDS + Hepatitis.B +
                        Adult.Mortality * Hepatitis.B,
                        data = life2015[-c(119, 53, 93), ], nv = 4)
summary(best.subset)

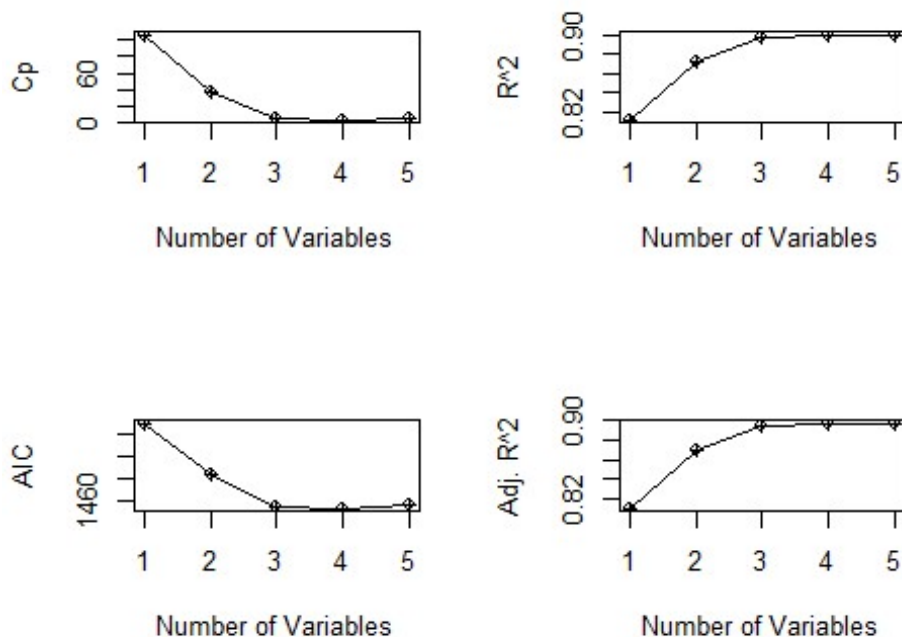
## Subset selection object
## Call: regsubsets.formula((((Life.expectancy^1.78787) - 1)/1.78787) ~
##      Income.composition.of.resources + Adult.Mortality + HIV.AIDS +
##      Hepatitis.B + Adult.Mortality * Hepatitis.B, data = life2015[-
##      c(119,
##      53, 93), ], nv = 4)
## 5 Variables (and intercept)
##
##              Forced in Forced out
## Income.composition.of.resources      FALSE      FALSE
## Adult.Mortality                     FALSE      FALSE
## HIV.AIDS                           FALSE      FALSE
## Hepatitis.B                         FALSE      FALSE
## Adult.Mortality:Hepatitis.B          FALSE      FALSE
## 1 subsets of each size up to 4
## Selection Algorithm: exhaustive
##      Income.composition.of.resources Adult.Mortality HIV.AIDS
Hepatitis.B
## 1 ( 1 ) "*" " " " " " "

```

```
## 2 ( 1 ) "*"
## 3 ( 1 ) "*"
## 4 ( 1 ) "*"
##      Adult.Mortality:Hepatitis.B
## 1 ( 1 ) " "
## 2 ( 1 ) " "
## 3 ( 1 ) "*"
## 4 ( 1 ) "*"

reg.summary <- summary(best.subset)

#Plotting
par(mfrow = c(2,2)) # split the plotting model in 2 x 2 grid
plot(ks$cp, type = "o", pch = 10, xlab = "Number of Variables", ylab = "Cp")
plot(ks$rsq, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"R^2")
plot(ks$aic, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"AIC")
plot(ks$adjr, type = "o", pch = 10, xlab = "Number of Variables", ylab =
"Adj. R^2")
```



```
paste("Based on Marlow's Cp the minimum value was obtained without
Adult.Mortality.")
```

```
## [1] "Based on Marlow's Cp the minimum value was obtained without
Adult.Mortality."
```

```

#MODEL DIAGNOSTICS
# (R.5) CHECKING LINEARITY
#Residuals plot
plot(life2015_box_modelR, which = 1)

# a Scale Location Plot
plot(life2015_box_modelR, which = 3)

paste("It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected.")

## [1] "It is verified some slightly change in the pattern for last fitts
values in the last plot, but in overall the average of residuals almost
horizontal and we can say that the residuals are equally spread that we can
conclude that the linearity assumption is respected."

# (R.6) Heteroscedasticity Test - the Breush-Pagan test
# Ho : heteroscedasticity is NOT presented (homoscedasticity)
# Ha: heteroscedasticity is presented
bptest(life2015_box_modelR)

##
## studentized Breusch-Pagan test
##
## data: life2015_box_modelR
## BP = 8.7085, df = 5, p-value = 0.1213

paste("The output displays the Breush-Pagan test that from the model
presented the p-value 0.1213 > 0.05, indicating the we should NOT REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity does not exist.")

## [1] "The output displays the Breush-Pagan test that from the model
presented the p-value 0.1213 > 0.05, indicating the we should NOT REJECT the
null hypthesis at 5 % level and consequently the test provides evidence that
heteroscedasticity does not exist."

# (R.7) Testing for normality
#NORMALITY - Shapiro-Wilk Test
# Ho : the sample data is significantly normally distributed
# Ha : the sample data is NOT significantly normally distributed
shapiro.test(residuals(life2015_box_modelR))

##
## Shapiro-Wilk normality test
##
## data: residuals(life2015_box_modelR)
## W = 0.99243, p-value = 0.7269

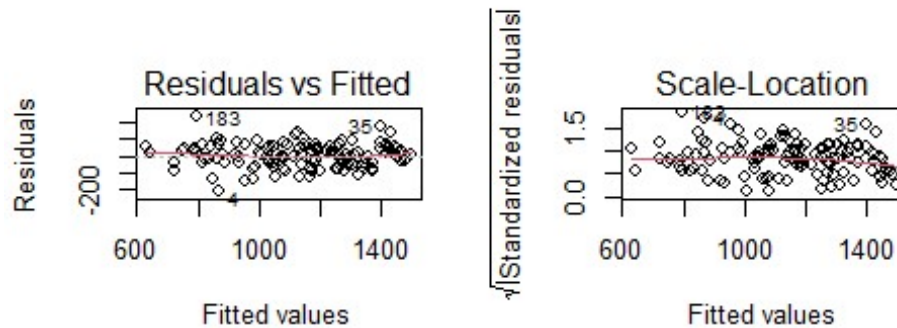
```

```
paste("Shapiro-Wilk normality test presented the p-value = 0.7269 > 0.05  
that indicated that we should NOT REJECT the null hypothesis and conclude  
that the residuals are normally distributed at 5 % level.")
```

```
## [1] "Shapiro-Wilk normality test presented the p-value = 0.7269 > 0.05  
that indicated that we should NOT REJECT the null hypothesis and conclude  
that the residuals are normally distributed at 5 % level."
```

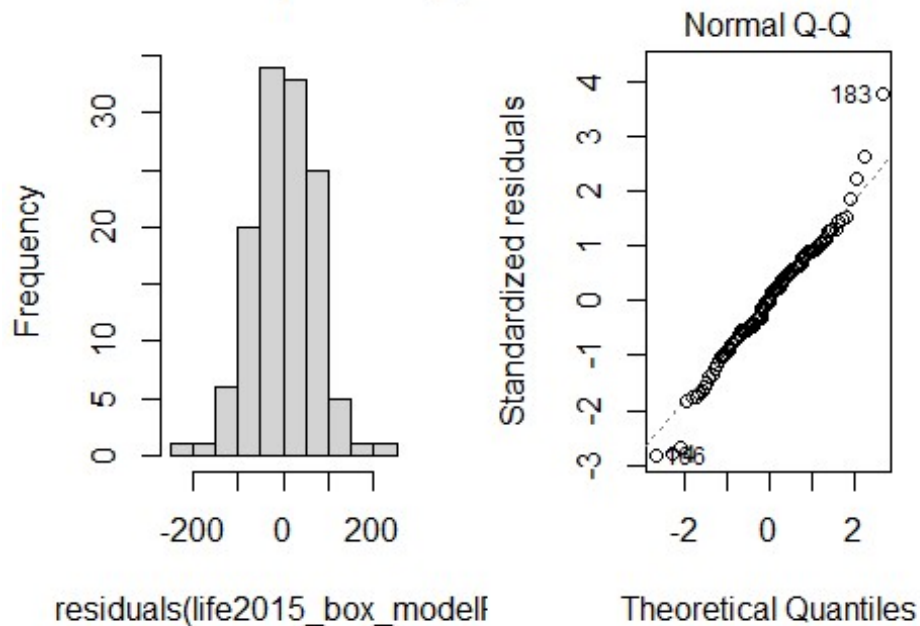
```
#Plots
```

```
par(mfrow = c(1,2))
```



```
hist(residuals(life2015_box_modelR))  
plot(life2015_box_modelM, which = 2)
```

am of residuals(life2015_b



```
paste("Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected.")
```

```
## [1] "Q-Q plot indicates that data points #183 and #4 have significant
discrepancy for what is expected."
```

```
# (R.8) Multicollinearity test
```

```
#VIF
```

```
imcdiag(life2015_box_modelR, method = "VIF")
```

```
##
```

```
## Call:
```

```
## imcdiag(mod = life2015_box_modelR, method = "VIF")
```

```
##
```

```
##
```

```
## VIF Multicollinearity Diagnostics
```

```
##
```

```
## VIF detection
```

```
## Income.composition.of.resources 1.6893 0
```

```
## Adult.Mortality 8.7175 0
```

```
## HIV.AIDS 1.8959 0
```

```
## Hepatitis.B 3.5898 0
```

```
## Adult.Mortality:Hepatitis.B 9.0869 0
```

```
##
```

```
## NOTE: VIF Method Failed to detect multicollinearity
```

```
##
```

```
##
```

```
## 0 --> COLLINEARITY is not detected by the test
##
## =====

paste("Multicollinearity verified between Adult.Mortality and the interaction
terms. ")

## [1] "Multicollinearity verified between Adult.Mortality and the interaction
terms. "

# (R.9) LEVERAGE OF THE POINTS
# Leverage Points
lev = hatvalues(life2015_box_modelR)
p = length(coef(life2015_box_modelR))
n = nrow(life2015)
outlier = lev[lev > (2 * p/n)]
print(outlier)

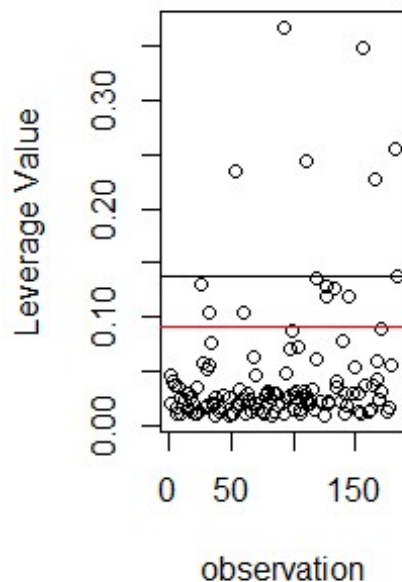
##          26          33          53          60          93          111          118
126
## 0.1310916 0.1051954 0.2352384 0.1036336 0.3664417 0.2436371 0.1353749
0.1195921
##          127          133          144          156          166          182          183
## 0.1290916 0.1262577 0.1187967 0.3482968 0.2271205 0.2555812 0.1372311

plot(rownames(life2015[-c(119, 53, 93), ]), lev, main = "Leverage in Life2015
Dataset",
      xlab = "observation", ylab = "Leverage Value")
abline(h = 2 * p/n, lty = 1, col = "red")
abline(h = 3 * p/n, lty = 1)

paste("The results shows that that the data points #26, #33, #53, #60, #93,
#111, #118, #126, #127, #166, #182and #183 presented leverage values higher
than 2.p/n that might be outliers.")

## [1] "The results shows that that the data points #26, #33, #53, #60, #93,
#111, #118, #126, #127, #166, #182and #183 presented leverage values higher
than 2.p/n that might be outliers."
```

Leverage in Life2015 Data



#Comparing the best models (M, P and R)

```
life2015_reduced_modelM<- lm(Life.expectancy ~  
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +  
                                under.five.deaths + Adult.Mortality *  
HIV.AIDS, data = life2015)  
  
life2015_reduced_modelP<- lm(Life.expectancy ~  
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +  
                                under.five.deaths + Hepatitis.B +  
Adult.Mortality * HIV.AIDS, data = life2015)  
  
life2015_reduced_modelR <- lm(Life.expectancy ~  
Income.composition.of.resources + Adult.Mortality + HIV.AIDS +  
                                Hepatitis.B + Adult.Mortality * Hepatitis.B,  
data = life2015)
```

#Comparing Models P and R without Box Cox Transformation (Anova does not compare with Box Cox)

#Ho: R model

#Ha: P model

```
anova(life2015_reduced_modelR, life2015_reduced_modelP)
```

```
## Analysis of Variance Table
```

```
##
```

```
## Model 1: Life.expectancy ~ Income.composition.of.resources +
```

```

Adult.Mortality +
##      HIV.AIDS + Hepatitis.B + Adult.Mortality * Hepatitis.B
## Model 2: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
##      HIV.AIDS + under.five.deaths + Hepatitis.B + Adult.Mortality *
##      HIV.AIDS
##      Res.Df      RSS Df Sum of Sq      F Pr(>F)
## 1      124 818.20
## 2      123 798.31  1      19.886 3.064 0.08254 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

paste("The output shows that Fcal = 3.064 with df 1, 123 (p-value = 0.0825 >
alpha = 0.05), indicating that we should clearly NOT REJECT the null
hypothesis. In other words, Model R presents better prediction than Model P
at 5 %.")

## [1] "The output shows that Fcal = 3.064 with df 1, 123 (p-value = 0.0825 >
alpha = 0.05), indicating that we should clearly NOT REJECT the null
hypothesis. In other words, Model R presents better prediction than Model P
at 5 %."

#Ho: R model
#Ha: M model
anova(life2015_reduced_modelR, life2015_reduced_modelM)

## Analysis of Variance Table
##
## Model 1: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
##      HIV.AIDS + Hepatitis.B + Adult.Mortality * Hepatitis.B
## Model 2: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
##      HIV.AIDS + under.five.deaths + Adult.Mortality * HIV.AIDS
##      Res.Df      RSS Df Sum of Sq F Pr(>F)
## 1      124 818.20
## 2      124 845.97  0      -27.776

paste("It is not possible to compare two model with the same number of
variables.")

## [1] "It is not possible to compare two model with the same number of
variables."

#Ho: M model
#Ha: P model
anova(life2015_reduced_modelM, life2015_reduced_modelP)

## Analysis of Variance Table
##
## Model 1: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +

```



```
## HIV.AIDS + under.five.deaths + Adult.Mortality * HIV.AIDS
## Model 2: Life.expectancy ~ Income.composition.of.resources +
Adult.Mortality +
## HIV.AIDS + under.five.deaths + Hepatitis.B + Adult.Mortality *
## HIV.AIDS
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 124 845.97
## 2 123 798.31 1 47.662 7.3436 0.007692 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

`paste("The output shows that Fcal = 47.662 with df 1, 123 (p-value = 0.007692 < alpha = 0.05), indicating that we should clearly REJECT the null hypothesis. In other words, Model P presents better prediction than Model M at 5 %.")`

```
## [1] "The output shows that Fcal = 47.662 with df 1, 123 (p-value =
0.007692 < alpha = 0.05), indicating that we should clearly REJECT the null
hypothesis. In other words, Model P presents better prediction than Model M
at 5 %."
```

`paste("based on the analysis, P and R are the best models.")`

```
## [1] "based on the analysis, P and R are the best models."
```

#Prediction Life Expectancy

#Creating a dataset with only the significant variables and removing the significant outliers

```
life2015_onlySign <- life2015[-c(119,53,93),]
```

```
life2015_onlySign$infant.deaths <- NULL
life2015_onlySign$Measles <- NULL
life2015_onlySign$BMI <- NULL
life2015_onlySign$Status <- NULL
life2015_onlySign$Polio <- NULL
life2015_onlySign$Diphtheria <- NULL
life2015_onlySign$GDP <- NULL
life2015_onlySign$Population <- NULL
life2015_onlySign$thinness..1.19.years <- NULL
life2015_onlySign$thinness.5.9.years <- NULL
life2015_onlySign$Schooling <- NULL
```

`summary(life2015_onlySign)`

```
## Life.expectancy Adult.Mortality Hepatitis.B under.five.deaths
## Min. :51.00 Min. : 1.0 Min. : 6.00 Min. : 0.00
## 1st Qu.:65.75 1st Qu.: 80.0 1st Qu.:78.00 1st Qu.: 1.00
## Median :72.00 Median :146.0 Median :91.00 Median : 3.00
## Mean :70.72 Mean :160.3 Mean :81.32 Mean : 37.48
## 3rd Qu.:75.95 3rd Qu.:218.5 3rd Qu.:96.00 3rd Qu.: 28.00
```

```
## Max. :85.00 Max. :484.0 Max. :99.00 Max. :1100.00
## HIV.AIDS Income.composition.of.resources
## Min. :0.1000 Min. :0.347
## 1st Qu.:0.1000 1st Qu.:0.551
## Median :0.1000 Median :0.701
## Mean :0.7984 Mean :0.675
## 3rd Qu.:0.5000 3rd Qu.:0.784
## Max. :9.3000 Max. :0.937
```

`paste("Based on statistics above for each variables, for the selected models P (foundation based on t-test) and R (foundation based in Stepwise) we have the following min, median, mean and max: values:")`

```
## [1] "Based on statistics above for each variables, for the selected
models P (foundation based on t-test) and R (foundation based in Stepwise) we
have the following min, median, mean and max: values:"
```

Income.composition.of.resources

```
meanIncome = mean(life2015_onlySign$Income.composition.of.resources)
medianIncome = median(life2015_onlySign$Income.composition.of.resources)
minIncome = min(life2015_onlySign$Income.composition.of.resources)
maxIncome = max(life2015_onlySign$Income.composition.of.resources)
```

Adult.Mortality

```
meanAdult = mean(life2015_onlySign$Adult.Mortality)
medianAdult = median(life2015_onlySign$Adult.Mortality)
minAdult = min(life2015_onlySign$Adult.Mortality)
maxAdult = max(life2015_onlySign$Adult.Mortality)
```

HIV.AIDS

```
meanHIV = mean(life2015_onlySign$HIV.AIDS)
medianHIV = median(life2015_onlySign$HIV.AIDS)
minHIV = min(life2015_onlySign$HIV.AIDS)
maxHIV = max(life2015_onlySign$HIV.AIDS)
```

Hepatitis.B

```
meanHepatitisB = mean(life2015_onlySign$Hepatitis.B)
medianHepatitisB = median(life2015_onlySign$Hepatitis.B)
minHepatitisB = min(life2015_onlySign$Hepatitis.B)
maxHepatitisB = max(life2015_onlySign$Hepatitis.B)
```

Under.five.deaths

```
meanUnder = mean(life2015_onlySign$Under.five.deaths)
medianUnder = median(life2015_onlySign$Under.five.deaths)
minUnder = min(life2015_onlySign$Under.five.deaths)
maxUnder = max(life2015_onlySign$Under.five.deaths)
```

Selected Models

```
life2015_box_modelP = lm((((Life.expectancy^1.66667) - 1)/1.66667) ~  
Income.composition.of.resources +  
Adult.Mortality + HIV.AIDS + under.five.deaths +  
Hepatitis.B +  
Adult.Mortality * HIV.AIDS, data =  
life2015_onlySign)
```

```
life2015_box_modelR = lm((((Life.expectancy^1.78787) - 1)/1.78787) ~  
Income.composition.of.resources +  
Adult.Mortality + HIV.AIDS + Hepatitis.B +  
Adult.Mortality * Hepatitis.B, data =  
life2015_onlySign)
```

1 - PREDICTION based on Means

```
meanData = data.frame(Income.composition.of.resources = meanIncome,  
Adult.Mortality = meanAdult,  
HIV.AIDS = meanHIV, under.five.deaths = meanUnder,  
Hepatitis.B = meanHepatitisB)
```

MODEL P

```
predict(life2015_box_modelP, meanData, interval = "predict")
```

```
##          fit          lwr          upr  
## 1 722.5345 636.7589 808.31
```

Converting to Life.Expectancy (Box-Cox Lambda)

```
pred_Mean_Life_P_fit = (722.5345 * 1.66667 + 1)^(1/1.66667)  
pred_Mean_Life_P_lwr = (636.7589 * 1.66667 + 1)^(1/1.66667)  
pred_Mean_Life_P_upr = (808.31 * 1.66667 + 1)^(1/1.66667)
```

```
paste("Considering the mean of significant variables of Model P as input,  
on the average Life Expectancy around the World in 2015 based on Model P is  
", pred_Mean_Life_P_fit, " years at 5 %p level. For 95% prediction interval,  
the life expectancy estimated by Model P is between ", pred_Mean_Life_P_lwr,  
" and ", pred_Mean_Life_P_upr)
```

```
## [1] "Considering the mean of significant variables of Model P as input,  
on the average Life Expectancy around the World in 2015 based on Model P is  
70.572822292507 years at 5 %p level. For 95% prediction interval, the life  
expectancy estimated by Model P is between 65.4239214422687 and  
75.4824759745119"
```

```

# MODEL R
predict(life2015_box_modelR, meanData, interval = "predict")

##          fit          lwr          upr
## 1 1139.689 995.4774 1283.9

# Converting to Life.Expectancy (Box-Cox Lambda)
pred_Mean_Life_R_fit = (1139.689 * 1.78787 + 1)^(1/1.78787)
pred_Mean_Life_R_lwr = (995.4774 * 1.78787 + 1)^(1/1.78787)
pred_Mean_Life_R_upr = (1283.9 * 1.78787 + 1)^(1/1.78787)

paste("Considering the mean of significant variables of Model R as input,
on the average Life Expectancy around the World in 2015 based on Model R is
", pred_Mean_Life_R_fit, " years at 5 %p level. For 95% prediction interval,
the life expectancy estimated by Model R is between ", pred_Mean_Life_R_lwr,
" and ", pred_Mean_Life_R_upr)

## [1] "Considering the mean of significant variables of Model R as input,
on the average Life Expectancy around the World in 2015 based on Model R is
70.9566810821455 years at 5 %p level. For 95% prediction interval, the life
expectancy estimated by Model R is between 65.7881166846681 and
75.844148896526"

# 2 - PREDICTION based on Medians
medianData = data.frame(Income.composition.of.resources = medianIncome,
Adult.Mortality = medianAdult,
                        HIV.AIDS = medianHIV, under.five.deaths = medianUnder,
Hepatitis.B = medianHepatitisB)

# MODEL P
predict(life2015_box_modelP, medianData, interval = "predict")

##          fit          lwr          upr
## 1 763.6861 677.8892 849.483

# Converting to Life.Expectancy (Box-Cox Lambda)
pred_Median_Life_P_fit = (763.6861 * 1.66667 + 1)^(1/1.66667)
pred_Median_Life_P_lwr = (677.8892 * 1.66667 + 1)^(1/1.66667)
pred_Median_Life_P_upr = (849.483 * 1.66667 + 1)^(1/1.66667)

paste("Considering the MEDIAN of significant variables of Model P as input,
on the average Life Expectancy in 2015 based on Model P is ",
pred_Median_Life_P_fit, " years at 5 %p level. For 95% prediction interval,
the life expectancy estimated by Model P is between ",
pred_Median_Life_P_lwr, " and ", pred_Median_Life_P_upr)

## [1] "Considering the MEDIAN of significant variables of Model P as
input, on the average Life Expectancy in 2015 based on Model P is
72.9557659000727 years at 5 %p level. For 95% prediction interval, the life
expectancy estimated by Model P is between 67.9253476462157 and
77.7647513548965"

```

```

# MODEL R
predict(life2015_box_modelR, medianData, interval = "predict")

##          fit          lwr          upr
## 1 1194.788 1050.42 1339.156

# Converting to Life.Expectancy (Box-Cox Lambda)
pred_Median_Life_R_fit = (1194.788 * 1.78787 + 1)^(1/1.78787)
pred_Median_Life_R_lwr = (1050.42 * 1.78787 + 1)^(1/1.78787)
pred_Median_Life_R_upr = (1339.156 * 1.78787 + 1)^(1/1.78787)

paste("Considering the Median of significant variables of Model R as input,
on the average Life Expectancy in 2015 based on Model R is ",
pred_Median_Life_R_fit, " years at 5 %p level. For 95% prediction interval,
the life expectancy estimated by Model R is between ",
pred_Median_Life_R_lwr, " and ", pred_Median_Life_R_upr)

## [1] "Considering the Median of significant variables of Model R as
input, on the average Life Expectancy in 2015 based on Model R is
72.8545149528908 years at 5 %p level. For 95% prediction interval, the life
expectancy estimated by Model R is between 67.7938422582918 and
77.6521251437127"

paste("for the mean and median values, both models indicated similar Life
Expectancy predictions, including the 95% prediction interval.")

## [1] "for the mean and median values, both models indicated similar Life
Expectancy predictions, including the 95% prediction interval."

```