

Exercise 1

In this exercise, we will be referring the dataset `Darts.csv`.

- (a) We were asked to find the best linear regression model for the response variable `Width`. First, we will fit two different linear models, which are obtained from forward step and backward step.

```
model0<-lm(Width ~ Length) # fit a minimal model
modelstep0<-step(model0, scope=formula(Width~Length*Thickness*Name)) # fit a forward step model
model1<-lm(Width~Length*Thickness*Name) # fit a full model
modelstep1 <- step(model1) # fit a (backward) step model from a full model
```

where the formula for each model are given as follows

```
> modelstep0
Call:
lm(formula = Width ~ Length + Name)
> modelstep1
Call:
lm(formula = Width ~ Length * Thickness * Name)
```

To choose the best model, we compare the AIC component of the models.

```
> AIC(modelstep0)
[1] 458.9517
> AIC(modelstep1)
[1] 432.4108
```

It follows that we have evidence to choose `modelstep1` over `modelstep0`. Now, we will check the diagnostic plot of the model that we chose.

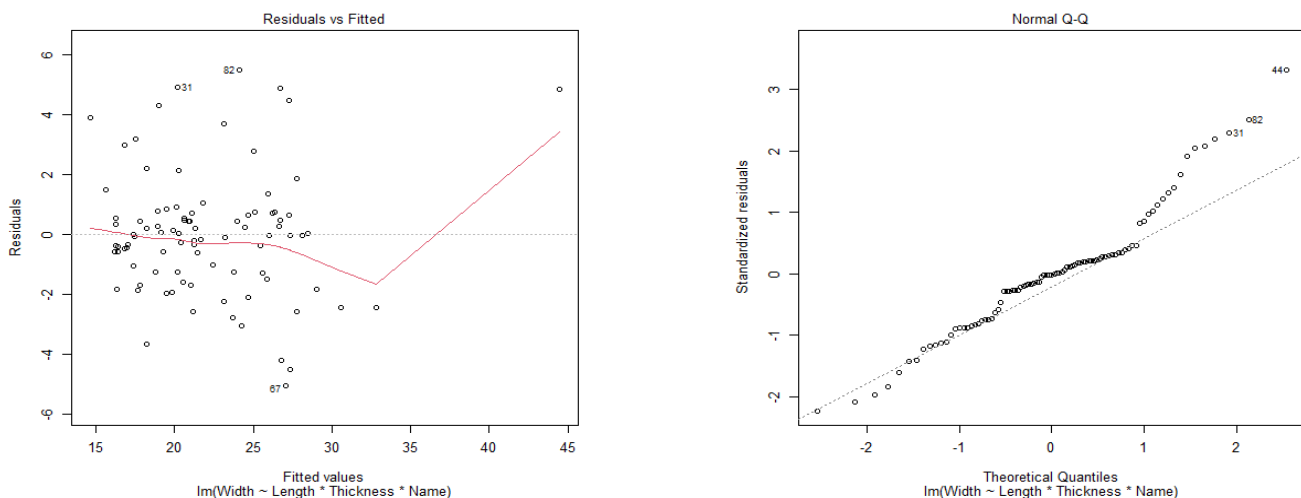


Figure 1: Two of the diagnostic plots of `modelstep1`

One of the obvious issue highlighted here is the violation of the normality of the errors. We also perform a Box-Cox transformation (as shown in Figure 2) and it suggests taking the logarithmic value of the response variable. Thus, we fit a new linear model.

```
> modelfix <- lm(log(Width) ~ Length * Thickness * Name)
```

From the diagnostic plots of `modelfix` which are shown in Figure 3 below, we see some improvements on the scatter of Residuals vs Fitted, however, it is still **not** obvious from the Normal Q-Q Plot that this model follow our assumption on the normality of the errors.

- (b) We will now predict the maximum width in milimetres for the designated input as follows

```
> # predicting a given input form our model fix
> new_dart <- data.frame(Name = "Travis", Thickness = 8, Length = 50)
> width_pred <- predict(modelfix, new_dart)
> exp(width_pred) # obtain the value for width from log(Width)
```

This returns the value **20.25064 mm**.

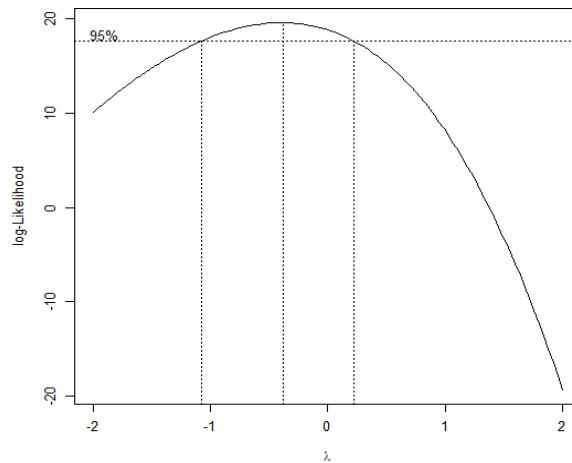


Figure 2: Box-Cox plot for modelstep1

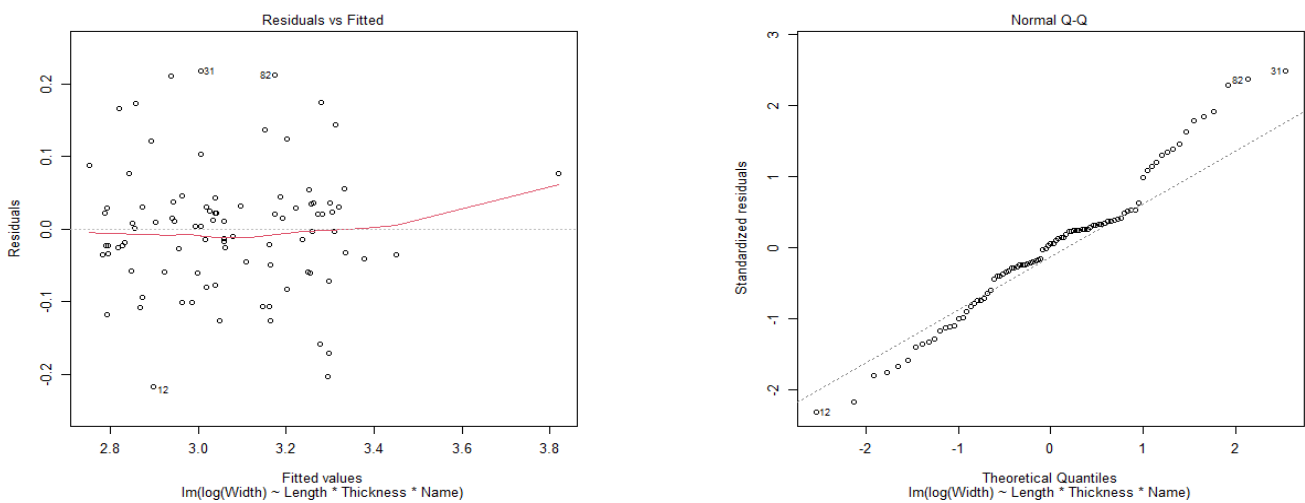


Figure 3: Two of the diagnostic plots of modelfix

Exercise 2

In this exercise we will be referring to the dataset extracted from `optimal.csv`.

- (a) We were asked to find the value of x_1 and x_2 that minimise the response variable y . First, we fit a model that includes the quadratic terms and also the interaction terms.

```
modelPoly <- lm(y ~ x1*x2 + I(x1^2) + I(x2^2)) # fit a LM with quadratic terms
```

Next, we extract the coefficients of the estimators as follows

```
> beta <- modelPoly$coefficients
> beta
(Intercept)      x1      x2      I(x1^2)      I(x2^2)      x1:x2
  0.86770    -2.81388   -0.08523    1.94370    0.19913   -0.00467
> b <- beta[2:3] # vector of coefficient for linear term.
> b
      x1      x2
-2.81387591 -0.08523381
>
> B <- matrix(0, 2, 2) # creating an empty matrix
> B[1,1] <- beta[4]
> B[2,2] <- beta[5]
> B[1,2] <- B[2,1] <- 0.5*beta[6]
> B # obtain the quadratic form
      [,1] [,2]
[1,]  1.94370 -0.00233
[2,] -0.00233  0.19913
```

After computing the vector b and the matrix B , we can now calculate the critical point.

```
x_opt<--0.5*solve(B)%*%b
> x_opt          # minimum point
```

This gives us $x_1 = 0.72411$ and $x_2 = 0.22250$.

- (b) Next, we will provide 95% confidence interval for y when x_1 and x_2 are as in the previous.

```
> new_opt <- data.frame(x1 = x_opt[1], x2 = x_opt[2])
> x_opt_conf <- predict(modelPoly, new_opt, interval = "confidence", level = 0.95)
> x_opt_conf
      fit      lwr      upr
1 -0.16056 -0.26511 -0.056016
```

Exercise 3

We will load the dataset `warpbreaks` and fit a generalised linear model for the dataset.

```
> library(datasets)
> data("warpbreaks")
> glm1<- glm(breaks ~ wool*tension,family=poisson, data = warpbreaks)
```

- (a) Under the function `glm`, we construct a generalised linear model to observe the response variable `breaks` of the dataset `warpbreaks`. Here, we specify `poisson` as the probability distribution together with the log function as the default link function.

Let Y be the number of breaks, then the model `glm1` follows

$$Y_i \sim \text{Pois}(\mu_i)$$

where Y_i independent for $i = 1, 2, \dots, n$, and the Poisson parameter (mean) μ_i is related to the estimators by the link function

$$\log(\mu_i) = \beta_0 + \beta_1 x_{1i} + \beta_2 x_{2i} + \beta_3 x_{3i} + \beta_4 x_{1i} x_{2i} + \beta_5 x_{1i} x_{3i}$$

where

$$x_{1i} = \begin{cases} 1 & , \text{if wool B} \\ 0 & , \text{otherwise} \end{cases}$$

$$x_{2i} = \begin{cases} 1 & , \text{if tension M} \\ 0 & , \text{otherwise} \end{cases}$$

$$x_{3i} = \begin{cases} 1 & , \text{if tension H} \\ 0 & , \text{otherwise.} \end{cases}$$

The estimates of the parameters β_j for $j = 0, \dots, 5$ are given below.

```
> glm1$coefficients
(Intercept)      woolB      tensionM      tensionH woolB:tensionM
   3.79674    -0.45663    -0.61868    -0.59580     0.63818
woolB:tensionH
   0.18836
```

and also the estimate for the dispersion parameter ϕ is equal to 1.

- (b) We were asked to calculate the expected value for the number of breaks when we have the type of wool A and tension M. Firstly, we store a new data frame as follows.

```
try_data <- data.frame(wool = "A", tension = "M")
```

Next, we use the function `predict()` to obtain the value of $\log \mu_i$ for `try_data` and apply the exponential function on the value to yield the expected value of the number of breaks.

```
ypred<-predict(glm1, try_data)
exp(ypred)
```

This returns the value **24**. Note that the result is the same if we calculated manually from the dataset.