

## **Abstract**

In this paper, we investigate the height of fish, and whether there is a relationship between the height and 6 variables. According to our research, we are going to build two models. One model also finds the interaction of category variable and non-category variables. we use ANOVA to choose one of these two models. After that, the one models we retained will be selected through the stepwise regression, transformation, including removing the outliers by studentized residuals method and robust regression to build our final model.

## **Introduction**

Fish is a special animal that they cannot live without water. To learn more about the physical property, we assume whether there is a linear relationship between its height and the other variables that are its length in different degrees, weight, width, and species. According to the dataset we try to investigate. We focus on Bream, Parkki, Perch, Pike, Roach, Smelt, and White fish as sample of an analysis.

## **Data Description**

Dataset:

There are 159 sample with 7 variables in this data, which is collected from the fish market.

Variables:

Species – species name of fish ('Bream', 'Parkki', 'Perch', 'Pike', 'Roach', 'Smelt', 'White fish')

Weight – weight of fish in Gram (g)

Length1 – vertical length in cm

Length2 – diagonal length in cm

Length3 – cross length in cm

Height – height in cm

Width – diagonal width in cm

New addition data point:

Species = Smelt, Weight = 8.0, Length1=10.2, Length2= 11.7

Length3= 11.8, Height = 1.9945, Width= 1.4152

## Methods

According to the data, two kinds of full model we are going to build:

### Model 1:

$$\text{Height} = \beta_0 + \beta_1 * (\text{Length1}) + \beta_2 * (\text{Length2}) + \beta_3 * (\text{Length3}) + \beta_4 * (\text{Weight}) + \beta_5 * (\text{Width}) + \beta_6 * (\text{Species})$$

### Model 2: (variable with interaction)

$$\begin{aligned} \text{Height} = & \beta_0 + \beta_1 * (\text{Length1}) + \beta_2 * (\text{Length2}) + \beta_3 * (\text{Length3}) + \beta_4 * (\text{Weight}) + \beta_5 * (\text{Width}) + \\ & \beta_6 * (\text{Species}) + \beta_7 * (\text{Species}) * (\text{Length1}) + \beta_8 * (\text{Species}) * (\text{Length2}) + \beta_9 * (\text{Species}) \\ & * (\text{Length3}) + \beta_{10} * (\text{Species}) * (\text{Weight}) + \beta_{11} * (\text{Species}) * (\text{Width}) \end{aligned}$$

Where the variable 'Species' is a categorical variable

Due to this regression analysis, the models follow these five assumptions:

1. The relationship between the response variable (Height) and the regressors is linear, at least approximately.
2. The error term  $\varepsilon$  has zero mean.
3. The error term  $\varepsilon$  has constant variance  $\sigma^2$ .
4. the errors are uncorrelated.
5. The errors are normally distributed.

In two model, scatterplot matrices help us to determine whether the response variable is linear relationship with the other variable by looking. Then, we find out whether it is a

significant evidence that there is a linear relationship between the response variable and each the regressors at a 0.05 significant level. Since both models are similar except for the interaction, the next step we will do is to detect if the interaction has a relationship with the response variable. We will choose the one model as our final model by ANOVA test.

Regarding the multicollinearity, we try to center for each regressor which variance inflation factor (VIF) is larger than 10. After that, remake the models with the modified dataset, and using stepwise regression method to retain or reduce the regressor by the minimum AIC in order to find our best model. Then, we use fitted plot and normal Q-Q plot to determine both models are linear and normal. If not, we use the transformation method to improve the model to be more linear and normal. We use studentized residual and cook's distance, and check whether there are influence points and outliers. If we find the influence point and outliers, then remove outlier, but we may keep the influence points because that points may not severely interfere in change of slope. After we remake the model again, make a comparison to the same model with robust regression. We also observe the weight in robust regression to determine the problematic observations and consider if we need do remove it. Through these kinds of method, this model is confirmed as the final model for studying the response variable.

## Result

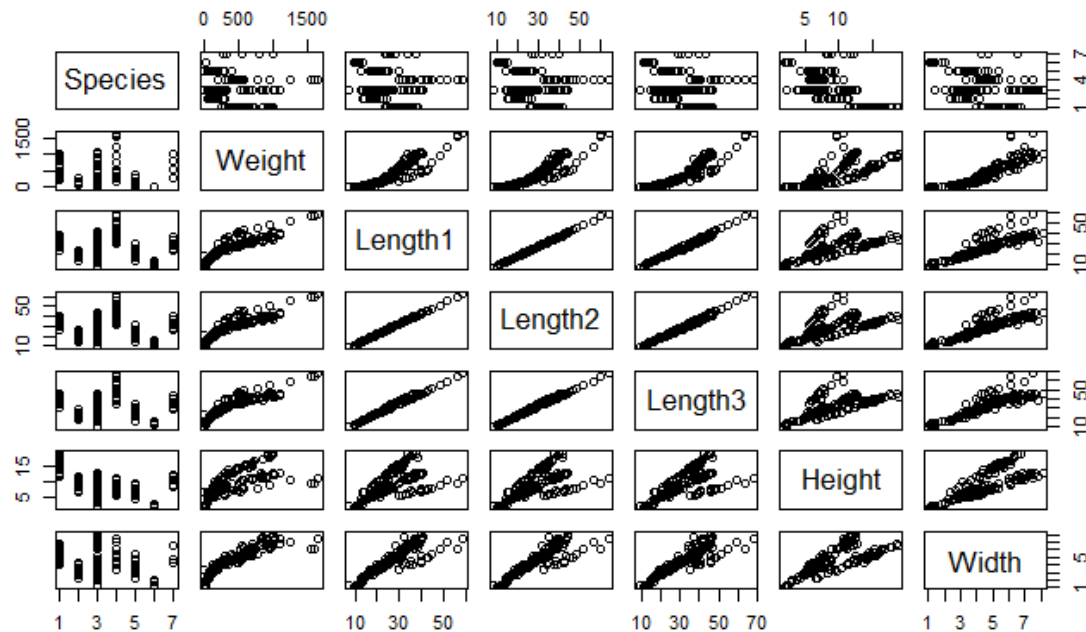


Figure 1: Scatterplot Matrices

In figure 1, we find that the response variable (Height) has a linear relationship among the regressors (Length1, Length2, Length3, Weight, Width,). Since the regressor ‘Species’ is a category variable, we use analysis tools to figure it out soon.

```
Call:
lm(formula = Height ~ Length1 + Length2 + Length3 + weight +
    width + species, data = fish)

Residuals:
    Min       1Q   Median       3Q      Max
-1.93193 -0.26715 -0.00155  0.26923  1.85551

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  4.8072649   0.8457595   5.684 6.85e-08 ***
Length1     -0.3111477   0.2314242  -1.344  0.181
Length2      0.1793095   0.2912636   0.616  0.539
Length3      0.2071286   0.1849836   1.120  0.265
weight       0.0002093   0.0005205   0.402  0.688
width        1.0677290   0.1226400   8.706 5.92e-15 ***
SpeciesParkki -1.8579528   0.4607772  -4.032 8.84e-05 ***
SpeciesPerch  -5.2114885   0.6304401  -8.266 7.50e-14 ***
SpeciesPike   -7.7045958   0.5812054 -13.256 < 2e-16 ***
SpeciesRoach  -4.7923659   0.4230867 -11.327 < 2e-16 ***
SpeciesSmelt  -5.3661495   0.6532114  -8.215 1.01e-13 ***
Specieswhitefish -4.4972697   0.4870806  -9.233 2.69e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.5924 on 147 degrees of freedom
Multiple R-squared:  0.9822,    Adjusted R-squared:  0.9809
F-statistic: 738.5 on 11 and 147 DF,  p-value: < 2.2e-16
```

Figure 2: summary of model 1 (full model)

As we can see from figure 2, the p-value of F-statistic is  $< 2.2e-16$ , we conclude that at least one variable has a linear relationship with the response variable (Height). We also find that 'Width' regressors and all regressors in 'species' has a extremely smaller p-value than 0.001. This result reflects that height is affected by width and species.

```
Call:
lm(formula = Height ~ Length1 + Length2 + Length3 + weight +
    width + Species + Length1 * Species + Length2 * Species +
    Length3 * Species + weight * Species + width * Species, data = fish)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-0.9371 -0.1763  0.0000  0.1863  0.8275
```

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	3.0624043	1.6467754	1.860	0.06545	.
Length1	-0.6350031	0.3417733	-1.858	0.06569	.
Length2	0.3510791	0.3827050	0.917	0.36084	
Length3	0.4296143	0.2975844	1.444	0.15150	
weight	0.0048081	0.0011091	4.335	3.1e-05	***
width	0.0540178	0.3119944	0.173	0.86284	
SpeciesParkki	-0.6994786	3.4643904	-0.202	0.84034	
SpeciesPerch	-2.7356999	1.7045022	-1.605	0.11119	
SpeciesPike	1.4690409	2.7448588	0.535	0.59353	
SpeciesRoach	-4.0407038	1.9901422	-2.030	0.04459	*
SpeciesSmelt	-2.3044835	3.3749411	-0.683	0.49607	
Specieswhitefish	11.4242686	28.9570727	0.395	0.69391	
Length1:SpeciesParkki	-0.5449564	5.7696788	-0.094	0.92491	
Length1:SpeciesPerch	0.6958125	0.4278716	1.626	0.10659	
Length1:SpeciesPike	1.5014187	0.9852743	1.524	0.13024	
Length1:SpeciesRoach	0.0868616	0.5800150	0.150	0.88121	
Length1:SpeciesSmelt	1.2287569	2.2162315	0.554	0.58034	
Length1:Specieswhitefish	2.8207841	7.9874757	0.353	0.72461	
Length2:SpeciesParkki	-4.1544092	7.5854342	-0.548	0.58495	
Length2:SpeciesPerch	-0.3462941	0.5498308	-0.630	0.53004	
Length2:SpeciesPike	-0.4453847	0.9403660	-0.474	0.63665	
Length2:SpeciesRoach	-0.0490672	0.6834118	-0.072	0.94289	
Length2:SpeciesSmelt	-0.7581324	1.4440478	-0.525	0.60057	
Length2:Specieswhitefish	0.5024883	5.3904100	0.093	0.92589	
Length3:SpeciesParkki	4.3013111	3.9666654	1.084	0.28043	
Length3:SpeciesPerch	-0.3396825	0.4007721	-0.848	0.39841	
Length3:SpeciesPike	-1.2049972	0.4138569	-2.912	0.00431	**
Length3:SpeciesRoach	-0.0423051	0.4430139	-0.095	0.92409	
Length3:SpeciesSmelt	-0.5290416	1.1785605	-0.449	0.65434	
Length3:Specieswhitefish	-3.3549908	6.0081443	-0.558	0.57763	
weight:SpeciesParkki	0.0083245	0.0145725	0.571	0.56893	
weight:SpeciesPerch	-0.0034765	0.0012615	-2.756	0.00679	**
weight:SpeciesPike	-0.0024257	0.0017308	-1.401	0.16372	
weight:SpeciesRoach	-0.0055119	0.0029649	-1.859	0.06553	.
weight:SpeciesSmelt	0.0819951	0.1587978	0.516	0.60659	
weight:Specieswhitefish	0.0006414	0.0094918	0.068	0.94624	
width:SpeciesParkki	-1.2269960	2.4304507	-0.505	0.61462	
width:SpeciesPerch	0.5082393	0.3458325	1.470	0.14435	
width:SpeciesPike	1.2694930	0.4641485	2.735	0.00721	**
width:SpeciesRoach	0.6835914	0.6153827	1.111	0.26892	
width:SpeciesSmelt	-0.0943835	1.1461661	-0.082	0.93451	
width:Specieswhitefish	0.5589113	0.4553299	1.227	0.22210	

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

```
Residual standard error: 0.3876 on 117 degrees of freedom
Multiple R-squared:  0.9939,    Adjusted R-squared:  0.9918
F-statistic: 468.4 on 41 and 117 DF,  p-value: < 2.2e-16
```

Figure 3: summary of model 2 (interaction full model)

In figure 3, the p-value of F-statistic is  $< 2.2e-16$ , we conclude that at least one variable has a linear relationship with the response variable (Height) too. We also find that 'Weight', 'SpeciesRoach', 'Length3:SpeciesPike', 'Weight:SpeciesPerch', 'Width:SpeciesPike' has small p-value that is less than 0.05 significant level. Hence, we conclude that height is affected by weight, roach, interaction of length3 and pike, interaction of weight and perch, and interaction of width and pike.

```
Analysis of variance Table

Model 1: Height ~ Length1 + Length2 + Length3 + Weight + Width + Species
Model 2: Height ~ Length1 + Length2 + Length3 + Weight + Width + Species +
  Length1 * Species + Length2 * Species + Length3 * Species +
  Weight * Species + Width * Species
  Res.Df    RSS Df Sum of Sq    F    Pr(>F)
1     147 51.589
2     117 17.577 30     34.012 7.5466 4.889e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Figure 4: Anova test for two model

Since the p-value is extremely small, we conclude that at least one variable which model 1 does not contain has a linear relationship with the response variable. Therefore, we will choose the model 2 to continue the rest of analysis.

```
Step:  AIC=-281.31
Height ~ Length1 + Length3 + weight + width + Species + Length1:Species +
  Length3:Species + weight:Species + width:Species

      Df Sum of Sq    RSS    AIC
<none>      17.805 -281.31
- width:Species    6   1.53882 19.344 -280.05
+ Length2          1   0.07158 17.734 -279.95
- weight:Species    6   1.63594 19.441 -279.24
- Length1:Species    6   2.11088 19.916 -275.38
- Length3:Species    6   2.43175 20.237 -272.82

Call:
lm(formula = Height ~ Length1 + Length3 + weight + width + Species +
  Length1:Species + Length3:Species + weight:Species + width:Species,
  data = fish)
```

Figure 5: stepwise regression



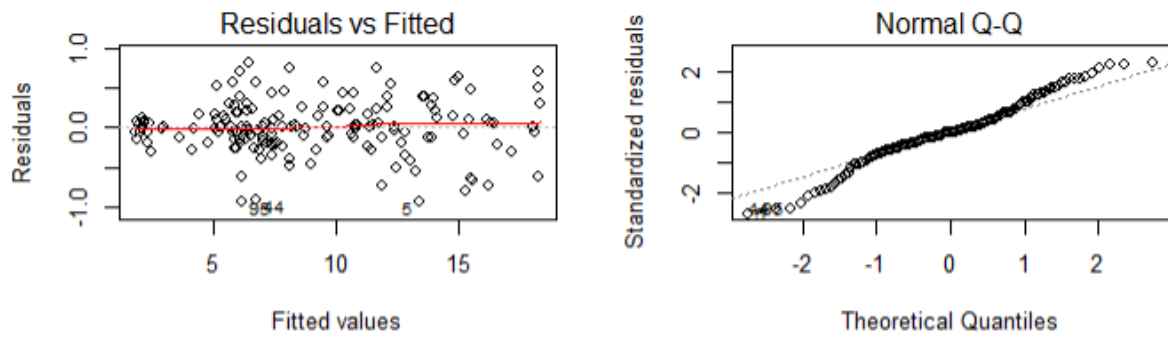


Figure 6: Fitted value plot and Normal Q-Q plot

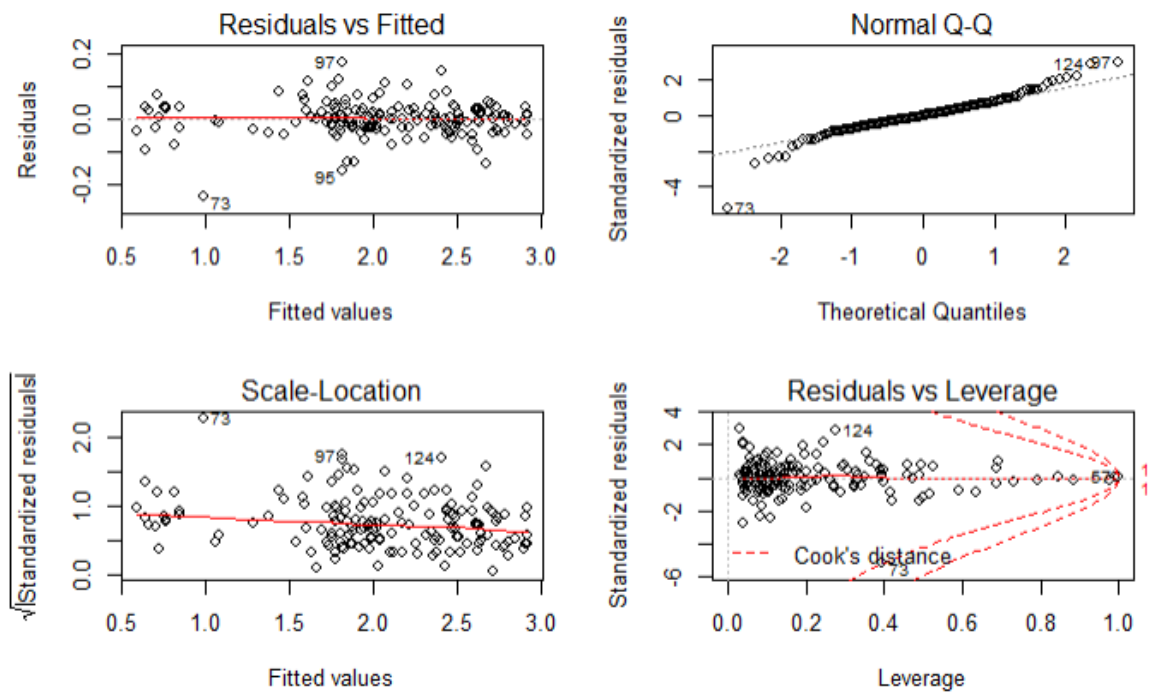


Figure 7: log-transformation for model 2

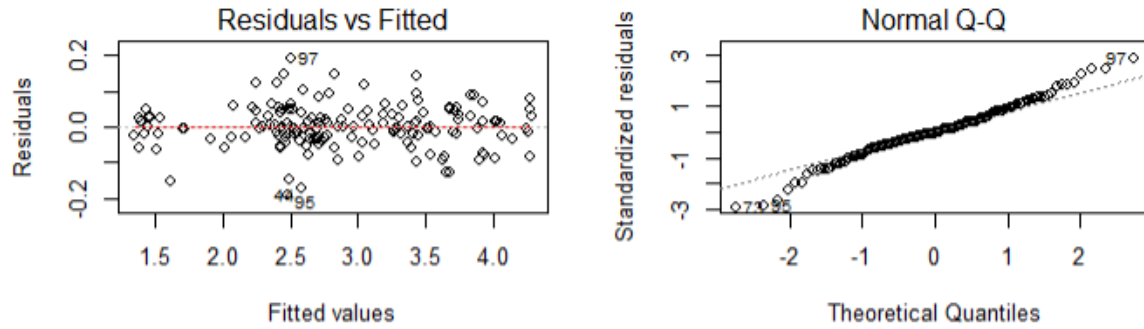


Figure 8: square root transformation for model 2

In figure 5, our variable selection is stepwise regression method. the minimum AIC we get is -281.31. In figure 6, we find that the model is linear, but it seems like non-normal. Hence, we will use transformation to try to make the model look better. Here are two kinds of transformation that I apply in figure 7 & 8. Compare log-transformation model to square root transformation model, we find that both models are linear. However, the model with log-transformation is more normally than the model with square root transformation. Therefore, we choose the log-transformation model as our part of final model.

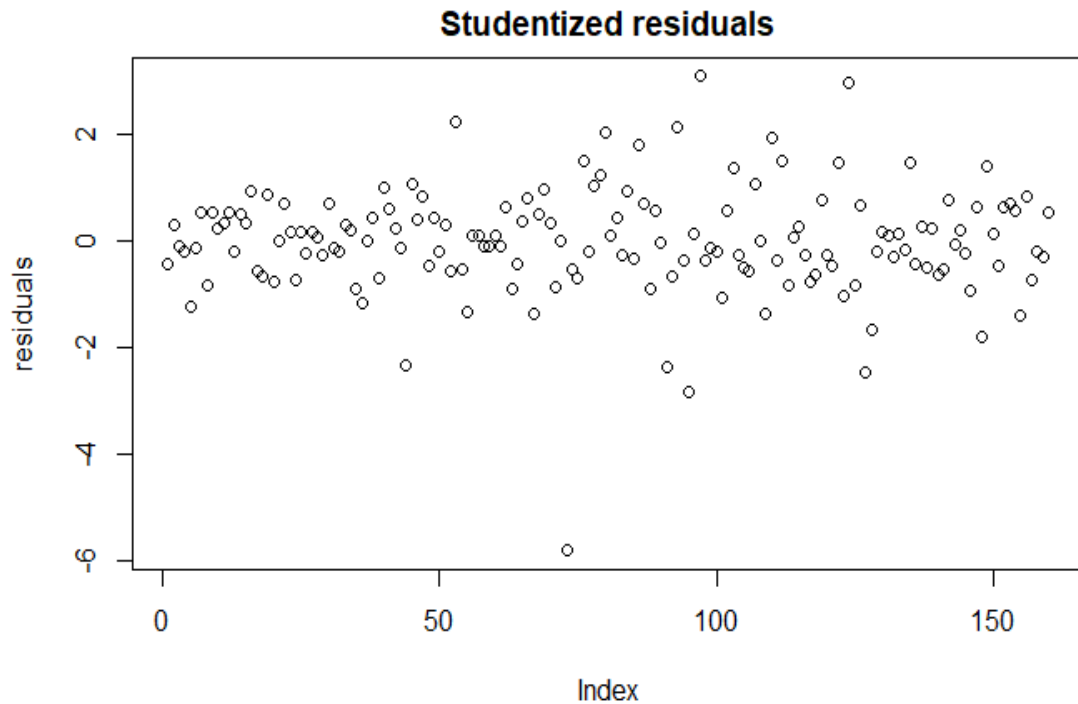


Figure 9: studentized residuals plot

In order to make the linear regression accurately, the next step we are going to do is to detect the outliers and remove them. In figure 9, we find that the residuals are distributed around zero. The mean of residual we acquire almost zero. To detect the outlier, we are considered the point is outlier which the absolute value of residual is larger than 3. After we remove those outliers, we repeat the detection until we cannot detect the outlier anymore. The result of the outliers we detect are 73, 97, 95.

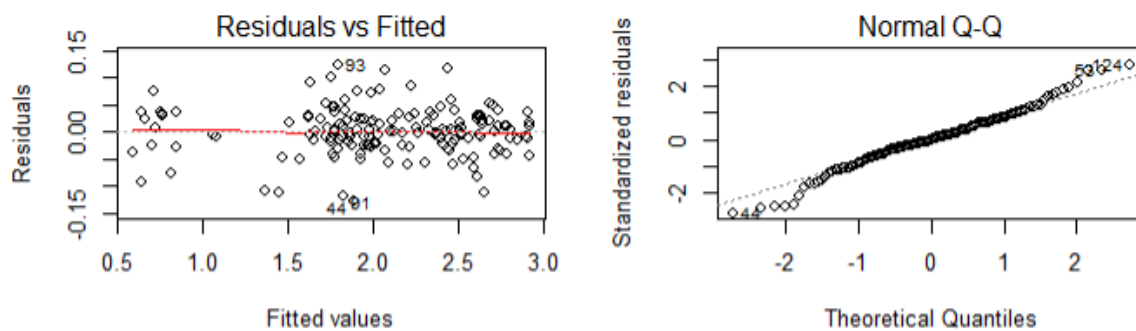


Figure 10: square root transformation for model 2 without some outlier

Since we remove some outlier that we detect, when we diagnostic the model, we still find that the model is linear but still non-normal. Therefore, we use robust regression method to analyze whether any problematic observation exists. According to robust regression model that we build, we want to find out those small weight so that a threshold that we set is 0.4. If the value of weight is less than 0.4, then this point regards as a problematic observation. As a result, observation 44, 53, 74, 75, 91, 93, 124, 127 are problematic observations. To consider making the model normal, we will remove these problematic observations.

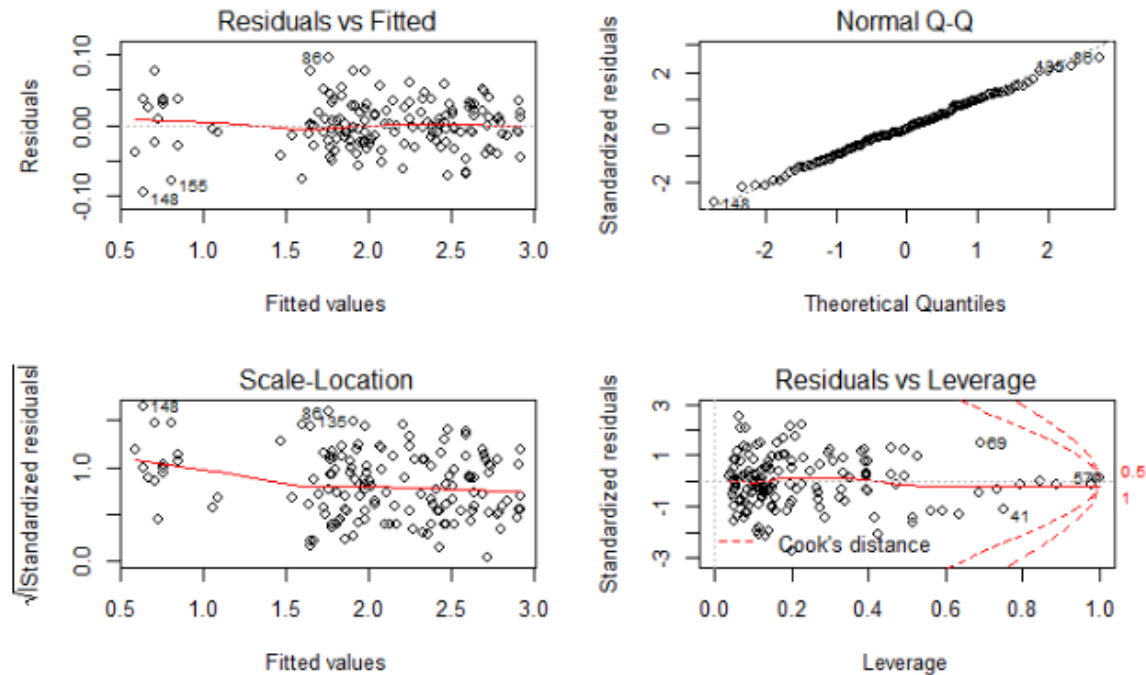


Figure 11: diagnostic plot for final modal of model 2

According to the diagnostic plot in figure 11, the model is linear and normal finally. However, in figure 12, we find that there is a serious multicollinearity for all variables.

	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
Length1	6.743672e+03	1	82.11986
Length3	8.769305e+03	1	93.64457
Weight	1.496073e+02	1	12.23141
Width	2.593501e+02	1	16.10435
Species	2.311278e+16	6	23.10225
Length1:Species	2.913187e+23	6	90.23272
Length3:Species	5.436983e+23	6	95.04877
Weight:Species	4.102141e+13	6	13.62754
Width:Species	4.276361e+15	6	20.07195

Figure 12: variance inflation factor (VIF)

## Conclusion

Based on the ANOVA test of final model (model 2), we conclude that height has a linear relationship among weight of fish in gram (Weight), vertical length in cm (Length1), cross length in cm (Length3), species ('Bream', 'Parkki', 'Perch', 'Pike', 'Roach', 'Smelt', 'White fish'), diagonal width in cm (Width), the interaction of vertical length in cm and species (Length1: Species), the interaction of weight of fish in gram and Species (Weight: Species), and interaction of diagonal width in cm and species (Width:Species).

### Analysis of Variance Table

Response: log(Height)

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
Length1	1	21.7693	21.7693	14519.4242	< 2.2e-16	***
Length3	1	14.6477	14.6477	9769.5444	< 2.2e-16	***
Weight	1	0.0104	0.0104	6.9599	0.0095000	**
Width	1	7.4687	7.4687	4981.3598	< 2.2e-16	***
Species	6	3.9284	0.6547	436.6806	< 2.2e-16	***
Length1:Species	6	0.1449	0.0241	16.1063	2.377e-13	***
Length3:Species	6	0.0191	0.0032	2.1217	0.0560683	.
Weight:Species	6	0.0371	0.0062	4.1185	0.0008847	***
Width:Species	6	0.0318	0.0053	3.5339	0.0030296	**
Residuals	114	0.1709	0.0015			

---

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

### ANOVA test of Final model

According to the summary of final model, our estimator for the linear regression is

$$\begin{aligned} \ln(\text{Height}) = & 2.349 + 0.0003006 * (\text{Weight}) - 0.7966 * (\text{Perch}) - 1.256 * (\text{Pike}) - 0.6401 * (\text{Roach}) \\ & - 0.5860 * (\text{Whitefish}) + 0.1556 * (\text{Pike}) * (\text{Length1}) - 0.1616 * (\text{Pike}) * (\text{Length3}) \\ & - 0.0006385 * (\text{Perch}) * (\text{Weight}) - 0.001192 * (\text{Roach}) * (\text{Weight}) + 0.129 * (\text{Width}) * (\text{Perch}) \\ & + 0.1636 * (\text{Width}) * (\text{Pike}) + 0.1459 * (\text{Width}) * (\text{Roach}) \end{aligned}$$

```

Call:
lm(formula = log(Height) ~ Length1 + Length3 + Weight + Width +
    Species + Length1:Species + Length3:Species + Weight:Species +
    Width:Species, data = fish_infremove2)

Residuals:
    Min       1Q   Median       3Q      Max
-0.094233 -0.022239 -0.001038  0.023741  0.093894

Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept)    2.349e+00  1.042e-01  22.544 < 2e-16 ***
Length1       -3.355e-02  2.611e-02  -1.285  0.201381
Length3        4.449e-02  2.563e-02   1.736  0.085242 .
Weight         3.006e-04  1.092e-04   2.753  0.006881 **
Width        -1.071e-03  3.054e-02  -0.035  0.972075
SpeciesParkki  2.105e-01  5.491e-01   0.383  0.702136
SpeciesPerch   -7.966e-01  1.289e-01  -6.181  1.01e-08 ***
SpeciesPike    -1.256e+00  1.452e-01  -8.648  3.81e-14 ***
SpeciesRoach   -6.401e-01  2.576e-01  -2.485  0.014402 *
SpeciesSmelt   -1.092e+00  5.512e-01  -1.980  0.050067 .
SpeciesWhitefish -5.860e-01  2.743e-01  -2.136  0.034802 *
Length1:SpeciesParkki -3.357e-01  3.326e-01  -1.009  0.315011
Length1:SpeciesPerch  4.680e-02  3.319e-02   1.410  0.161300
Length1:SpeciesPike   1.556e-01  3.721e-02   4.182  5.70e-05 ***
Length1:SpeciesRoach  7.525e-03  3.936e-02   0.191  0.848734
Length1:SpeciesSmelt  1.147e-01  1.323e-01   0.867  0.387951
Length1:SpeciesWhitefish 4.139e-01  6.170e-01   0.671  0.503662
Length3:SpeciesParkki  3.091e-01  2.808e-01   1.101  0.273333
Length3:SpeciesPerch -3.121e-02  3.214e-02  -0.971  0.333596
Length3:SpeciesPike   -1.616e-01  3.847e-02  -4.202  5.29e-05 ***
Length3:SpeciesRoach  1.460e-02  3.728e-02   0.391  0.696186
Length3:SpeciesSmelt  -7.186e-02  1.121e-01  -0.641  0.522665
Length3:SpeciesWhitefish -4.084e-01  5.992e-01  -0.681  0.496954
Weight:SpeciesParkki  3.332e-04  1.400e-03   0.238  0.812295
Weight:SpeciesPerch  -6.385e-04  1.450e-04  -4.402  2.43e-05 ***
Weight:SpeciesPike   -6.064e-05  1.601e-04  -0.379  0.705639
Weight:SpeciesRoach  -1.192e-03  3.078e-04  -3.871  0.000181 ***
Weight:SpeciesSmelt  2.027e-02  1.238e-02   1.637  0.104331
Weight:SpeciesWhitefish 2.482e-04  9.224e-04   0.269  0.788323
Width:SpeciesParkki  -8.575e-02  2.137e-01  -0.401  0.688958
Width:SpeciesPerch   1.290e-01  3.537e-02   3.648  0.000400 ***
Width:SpeciesPike    1.636e-01  4.202e-02   3.893  0.000167 ***
Width:SpeciesRoach   1.459e-01  6.173e-02   2.364  0.019775 *
Width:SpeciesSmelt   7.723e-03  9.588e-02   0.081  0.935940
Width:SpeciesWhitefish 7.343e-02  4.501e-02   1.631  0.105575
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.03872 on 114 degrees of freedom
Multiple R-squared:  0.9965,    Adjusted R-squared:  0.9954
F-statistic: 942.7 on 34 and 114 DF,  p-value: < 2.2e-16

```

According to the formula of linear regression, we find there is a positive relation between height and weight for most of fish except perch and pike. The height of Bream and smelt are affected by their weight. Perch, pike and roach have a positive relation between diagonal width and height. Only the height of pike has negative effect in cross length, but positive in vertical length.

## Appendix

### Summary of model 1:

```
> summary(m1)
```

Call:

```
lm(formula = Height ~ Length1 + Length2 + Length3 + Weight +  
    Width + Species, data = fish)
```

Residuals:

Min	1Q	Median	3Q	Max
-1.93143	-0.25742	-0.01263	0.27204	1.86872

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	4.7537079	0.8364704	5.683	6.81e-08	***
Length1	-0.2783956	0.2209379	-1.260	0.210	
Length2	0.1224601	0.2663392	0.460	0.646	
Length3	0.2323394	0.1771879	1.311	0.192	
Weight	0.0002154	0.0005190	0.415	0.679	
Width	1.0626602	0.1218860	8.718	5.30e-15	***
SpeciesParkki	-1.8203186	0.4531272	-4.017	9.34e-05	***
SpeciesPerch	-5.1389241	0.6111284	-8.409	3.20e-14	***
SpeciesPike	-7.6633827	0.5735744	-13.361	< 2e-16	***
SpeciesRoach	-4.7605520	0.4169724	-11.417	< 2e-16	***
SpeciesSmelt	-5.3437497	0.6499237	-8.222	9.35e-14	***
SpeciesWhitefish	-4.4472738	0.4749903	-9.363	< 2e-16	***

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.5909 on 148 degrees of freedom  
Multiple R-squared: 0.9825, Adjusted R-squared: 0.9812  
F-statistic: 754.9 on 11 and 148 DF, p-value: < 2.2e-16

### ANOVA test of model 1

```
> anova(m1)
```

Analysis of Variance Table

Response: Height

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
Length1	1	1176.62	1176.62	3370.015	< 2.2e-16	***
Length2	1	685.32	685.32	1962.862	< 2.2e-16	***
Length3	1	600.33	600.33	1719.442	< 2.2e-16	***
Weight	1	98.69	98.69	282.651	< 2.2e-16	***
Width	1	203.69	203.69	583.385	< 2.2e-16	***
Species	6	134.76	22.46	64.328	< 2.2e-16	***
Residuals	148	51.67	0.35			

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

### Summary of model 2

```
> summary(m2)
```

Call:

```
lm(formula = Height ~ Length1 + Length2 + Length3 + Weight +  
    Width + Species + Length1 * Species + Length2 * Species +  
    Length3 * Species + Weight * Species + Width * Species, data = fish)
```

Residuals:

Min	1Q	Median	3Q	Max
-0.9371	-0.1739	0.0000	0.1836	0.8275

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	3.0624043	1.6405010	1.867	0.06442	.
Length1	-0.6350031	0.3404711	-1.865	0.06466	.
Length2	0.3510791	0.3812468	0.921	0.35900	
Length3	0.4296143	0.2964506	1.449	0.14994	
weight	0.0048081	0.0011048	4.352	2.89e-05	***
width	0.0540178	0.3108057	0.174	0.86232	
SpeciesParkki	-0.6994786	3.4511908	-0.203	0.83974	
SpeciesPerch	-2.7356999	1.6980080	-1.611	0.10982	
SpeciesPike	1.4690409	2.7344007	0.537	0.59211	
SpeciesRoach	-4.0407038	1.9825596	-2.038	0.04377	*
SpeciesSmelt	-2.4262802	3.3406615	-0.726	0.46910	
SpeciesWhitefish	11.4242686	28.8467439	0.396	0.69280	
Length1:SpeciesParkki	-0.5449564	5.7476959	-0.095	0.92462	
Length1:SpeciesPerch	0.6958125	0.4262414	1.632	0.10525	
Length1:SpeciesPike	1.5014187	0.9815203	1.530	0.12877	
Length1:SpeciesRoach	0.0868616	0.5778051	0.150	0.88076	
Length1:SpeciesSmelt	0.7132029	1.5171580	0.470	0.63916	
Length1:SpeciesWhitefish	2.8207841	7.9570427	0.355	0.72360	
Length2:SpeciesParkki	-4.1544092	7.5565331	-0.550	0.58351	
Length2:SpeciesPerch	-0.3462941	0.5477359	-0.632	0.52846	
Length2:SpeciesPike	-0.4453847	0.9367831	-0.475	0.63535	
Length2:SpeciesRoach	-0.0490672	0.6808080	-0.072	0.94267	
Length2:SpeciesSmelt	-0.3379158	0.6002762	-0.563	0.57455	
Length2:SpeciesWhitefish	0.5024883	5.3698721	0.094	0.92561	
Length3:SpeciesParkki	4.3013111	3.9515521	1.089	0.27859	
Length3:SpeciesPerch	-0.3396825	0.3992452	-0.851	0.39660	
Length3:SpeciesPike	-1.2049972	0.4122801	-2.923	0.00416	**
Length3:SpeciesRoach	-0.0423051	0.4413260	-0.096	0.92380	
Length3:SpeciesSmelt	-0.4401571	1.1410410	-0.386	0.70038	
Length3:SpeciesWhitefish	-3.3549908	5.9852528	-0.561	0.57617	
weight:SpeciesParkki	0.0083245	0.0145170	0.573	0.56744	
weight:SpeciesPerch	-0.0034765	0.0012567	-2.766	0.00658	**
weight:SpeciesPike	-0.0024257	0.0017242	-1.407	0.16211	
weight:SpeciesRoach	-0.0055119	0.0029536	-1.866	0.06450	.
weight:SpeciesSmelt	0.0516065	0.1268345	0.407	0.68483	
weight:SpeciesWhitefish	0.0006414	0.0094557	0.068	0.94603	
width:SpeciesParkki	-1.2269960	2.4211905	-0.507	0.61326	
width:SpeciesPerch	0.5082393	0.3445149	1.475	0.14281	
width:SpeciesPike	1.2694930	0.4623800	2.746	0.00699	**
width:SpeciesRoach	0.6835914	0.6130380	1.115	0.26708	
width:SpeciesSmelt	-0.0203541	1.1183303	-0.018	0.98551	
width:SpeciesWhitefish	0.5589113	0.4535951	1.232	0.22033	

---  
 signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.3861 on 118 degrees of freedom  
 Multiple R-squared: 0.994, Adjusted R-squared: 0.992  
 F-statistic: 479.9 on 41 and 118 DF, p-value: < 2.2e-16

## ANOVA test of model 2

> anova(m2)

Analysis of Variance Table

Response: Height

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
Length1	1	1176.62	1176.62	7892.0267	<2e-16	***
Length2	1	685.32	685.32	4596.7029	<2e-16	***
Length3	1	600.33	600.33	4026.6530	<2e-16	***
weight	1	98.69	98.69	661.9237	<2e-16	***
width	1	203.69	203.69	1366.1938	<2e-16	***
species	6	134.76	22.46	150.6455	<2e-16	***



Length1:Species	6	29.06	4.84	32.4852	<2e-16	***
Length2:Species	6	1.06	0.18	1.1852	0.3189	
Length3:Species	6	1.55	0.26	1.7275	0.1205	
Weight:Species	6	1.15	0.19	1.2865	0.2687	
Width:Species	6	1.27	0.21	1.4142	0.2148	
Residuals	118	17.59	0.15			

---  
 Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

## Model selection by ANOVA

```
> anova(m1,m2)
```

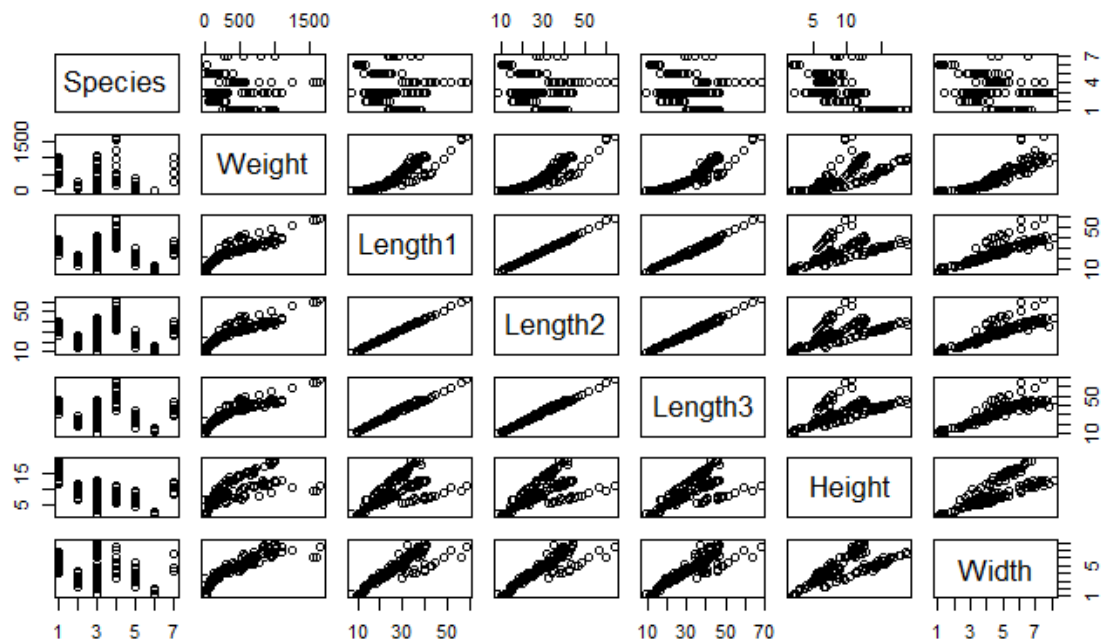
Analysis of Variance Table

Model 1: Height ~ Length1 + Length2 + Length3 + Weight + width + Species  
 Model 2: Height ~ Length1 + Length2 + Length3 + Weight + width + Species +

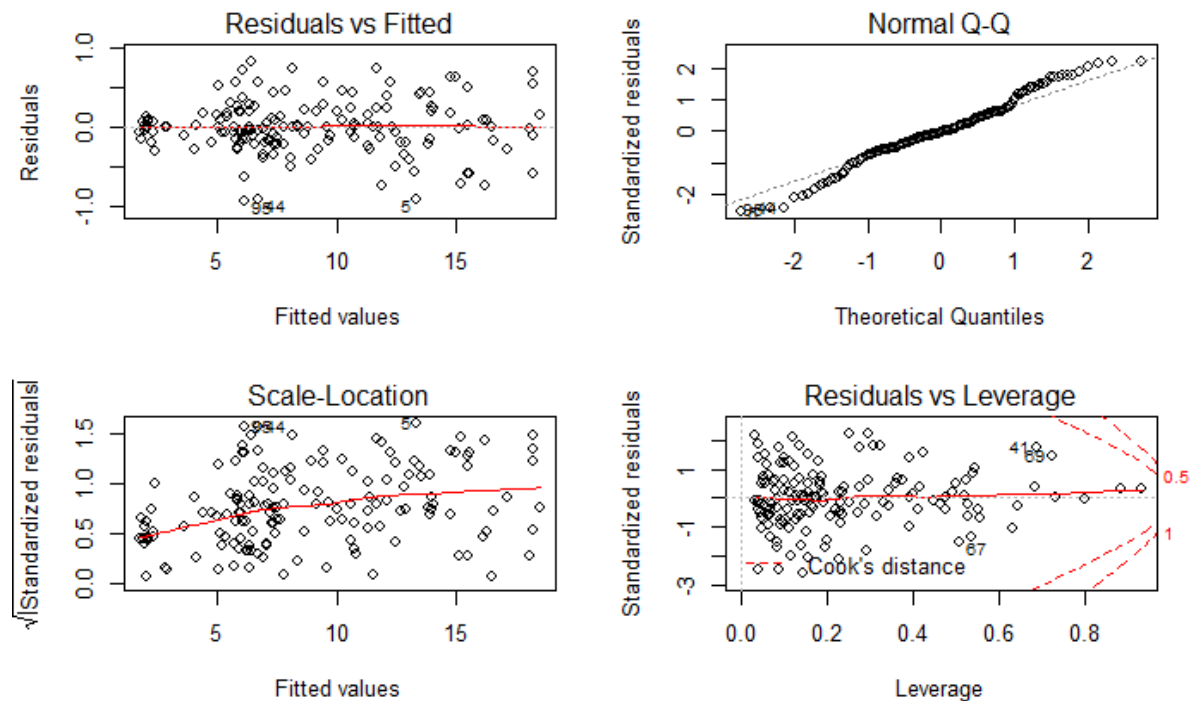
	Length1 * Species + Length2 * Species + Length3 * Species + Weight * Species + width * Species
Res.Df	RSS Df Sum of Sq F Pr(>F)
1	148 51.673
2	118 17.593 30 34.081 7.6197 3.049e-16 ***

---  
 Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

## Scatterplot Matrices



## Diagnostic plot of model 2(full model)



## Stepwise regression of model 2

```
> step(m2, direction = 'both')
```

Start: AIC=-269.23

```
Height ~ Length1 + Length2 + Length3 + Weight + width + Species +
        Length1 * Species + Length2 * Species + Length3 * Species +
        weight * Species + width * Species
```

	Df	Sum of Sq	RSS	AIC
- Length2:Species	6	0.14119	17.734	-279.95
- Length1:Species	6	0.70363	18.296	-274.96
- width:Species	6	1.26507	18.858	-270.12
<none>			17.593	-269.23
- weight:Species	6	1.43702	19.030	-268.67
- Length3:Species	6	1.88083	19.473	-264.98

Step: AIC=-279.95

```
Height ~ Length1 + Length2 + Length3 + Weight + width + Species +
        Length1:Species + Length3:Species + Weight:Species + Width:Species
```

	Df	Sum of Sq	RSS	AIC
- Length2	1	0.07158	17.805	-281.31
<none>			17.734	-279.95
- width:Species	6	1.54194	19.276	-278.61
- weight:Species	6	1.57419	19.308	-278.35
- Length1:Species	6	1.84468	19.578	-276.12
- Length3:Species	6	2.18655	19.920	-273.35
+ Length2:Species	6	0.14119	17.593	-269.23

Step: AIC=-281.31

```
Height ~ Length1 + Length3 + Weight + Width + Species + Length1:Species +
        Length3:Species + Weight:Species + Width:Species
```

	Df	Sum of Sq	RSS	AIC
<none>			17.805	-281.31
- width:Species	6	1.53882	19.344	-280.05
+ Length2	1	0.07158	17.734	-279.95
- weight:Species	6	1.63594	19.441	-279.24
- Length1:Species	6	2.11088	19.916	-275.38
- Length3:Species	6	2.43175	20.237	-272.82

```
Call:
lm(formula = Height ~ Length1 + Length3 + Weight + Width + Species +
    Length1:Species + Length3:Species + Weight:Species + Width:Species,
    data = fish)
```

## Summary of stepwise regression of model 2

```
> summary(m2_new)
```

```
Call:
lm(formula = Height ~ Length1 + Length3 + Weight + Width + Species +
    Length1:Species + Length3:Species + Weight:Species + Width:Species,
    data = fish_new)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-0.9456 -0.1680  0.0009  0.1831  0.8275
```

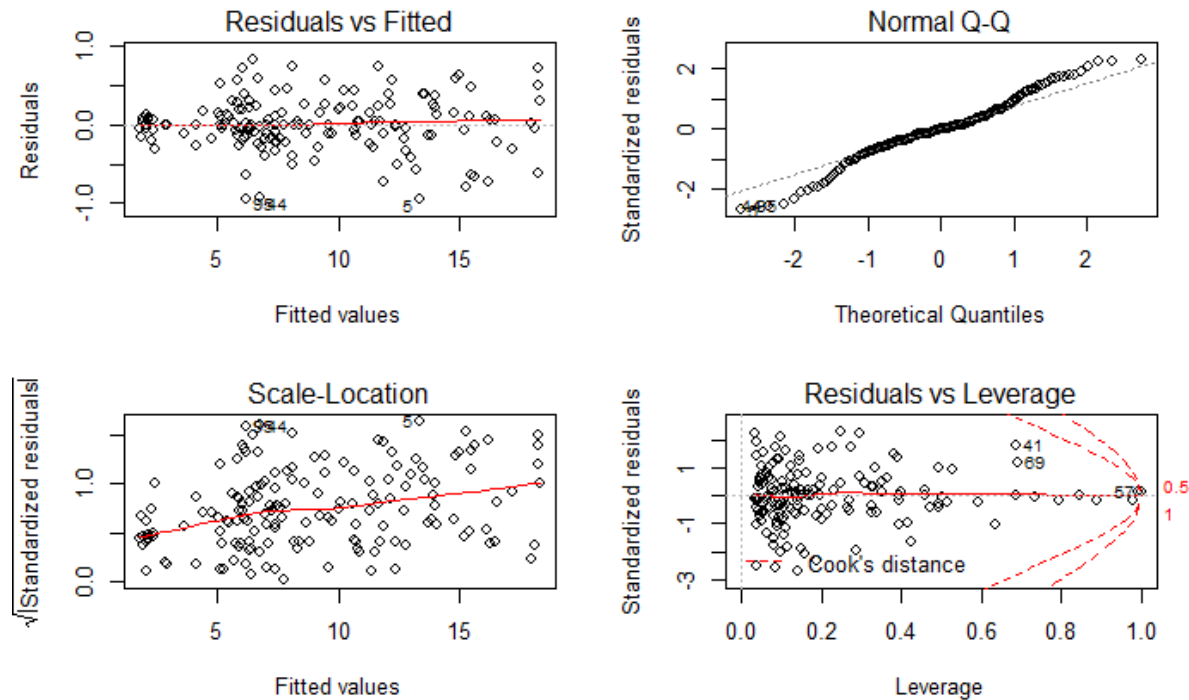
### Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	9.1856824	1.0156923	9.044	2.43e-15	***
Length1	-0.4329889	0.2545065	-1.701	0.091375	.
Length3	0.5679984	0.2497780	2.274	0.024670	*
weight	0.0049808	0.0010642	4.680	7.34e-06	***
width	0.1112336	0.2976662	0.374	0.709271	
SpeciesParkki	4.4560143	5.3525435	0.833	0.406712	
SpeciesPerch	-4.3328021	1.2222135	-3.545	0.000553	***
SpeciesPike	-8.6960663	1.4153659	-6.144	9.91e-09	***
SpeciesRoach	-4.1654081	2.4501147	-1.700	0.091601	.
SpeciesSmelt	-6.4750810	5.3724391	-1.205	0.230387	
SpeciesWhitefish	-4.3019958	2.6738070	-1.609	0.110152	
Length1:SpeciesParkki	-3.1101118	3.2418035	-0.959	0.339221	
Length1:SpeciesPerch	0.4960196	0.3094512	1.603	0.111480	
Length1:SpeciesPike	1.2022204	0.3626636	3.315	0.001200	**
Length1:SpeciesRoach	0.0808139	0.3808095	0.212	0.832284	
Length1:SpeciesSmelt	0.5315168	1.2897011	0.412	0.680954	
Length1:SpeciesWhitefish	3.4220296	6.0142027	0.569	0.570383	
Length3:SpeciesParkki	2.7598087	2.7369509	1.008	0.315234	
Length3:SpeciesPerch	-0.4755035	0.2971774	-1.600	0.112108	
Length3:SpeciesPike	-1.3417852	0.3749942	-3.578	0.000493	***
Length3:SpeciesRoach	-0.0810769	0.3625491	-0.224	0.823410	
Length3:SpeciesSmelt	-0.5835065	1.0923038	-0.534	0.594153	
Length3:SpeciesWhitefish	-3.4448816	5.8408993	-0.590	0.556399	
weight:SpeciesParkki	0.0101556	0.0136459	0.744	0.458138	
weight:SpeciesPerch	-0.0036497	0.0012140	-3.006	0.003198	**
weight:SpeciesPike	-0.0026670	0.0015610	-1.709	0.090022	.
weight:SpeciesRoach	-0.0057025	0.0028810	-1.979	0.049974	*
weight:SpeciesSmelt	0.0506081	0.1206719	0.419	0.675654	
weight:SpeciesWhitefish	0.0008167	0.0089907	0.091	0.927763	
width:SpeciesParkki	-1.8627857	2.0827500	-0.894	0.372833	
width:SpeciesPerch	0.4509554	0.3311808	1.362	0.175755	
width:SpeciesPike	1.1918384	0.4095306	2.910	0.004277	**
width:SpeciesRoach	0.6635906	0.5922474	1.120	0.264665	
width:SpeciesSmelt	-0.0611935	0.9345077	-0.065	0.947895	
width:SpeciesWhitefish	0.5049614	0.4387324	1.151	0.251947	

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

Residual standard error: 0.3774 on 125 degrees of freedom  
Multiple R-squared: 0.994, Adjusted R-squared: 0.9923  
F-statistic: 605.7 on 34 and 125 DF, p-value: < 2.2e-16

## Diagnostic plot of stepwise regression of model 2



## Summary of log-transformation

```
> summary(m2_new_log)
```

Call:

```
lm(formula = log(Height) ~ Length1 + Length3 + weight + width +  
    Species + Length1:Species + Length3:Species + weight:Species +  
    width:Species, data = fish_new)
```

Residuals:

Min	1Q	Median	3Q	Max
-0.238607	-0.025787	-0.000327	0.026715	0.174085

Coefficients:

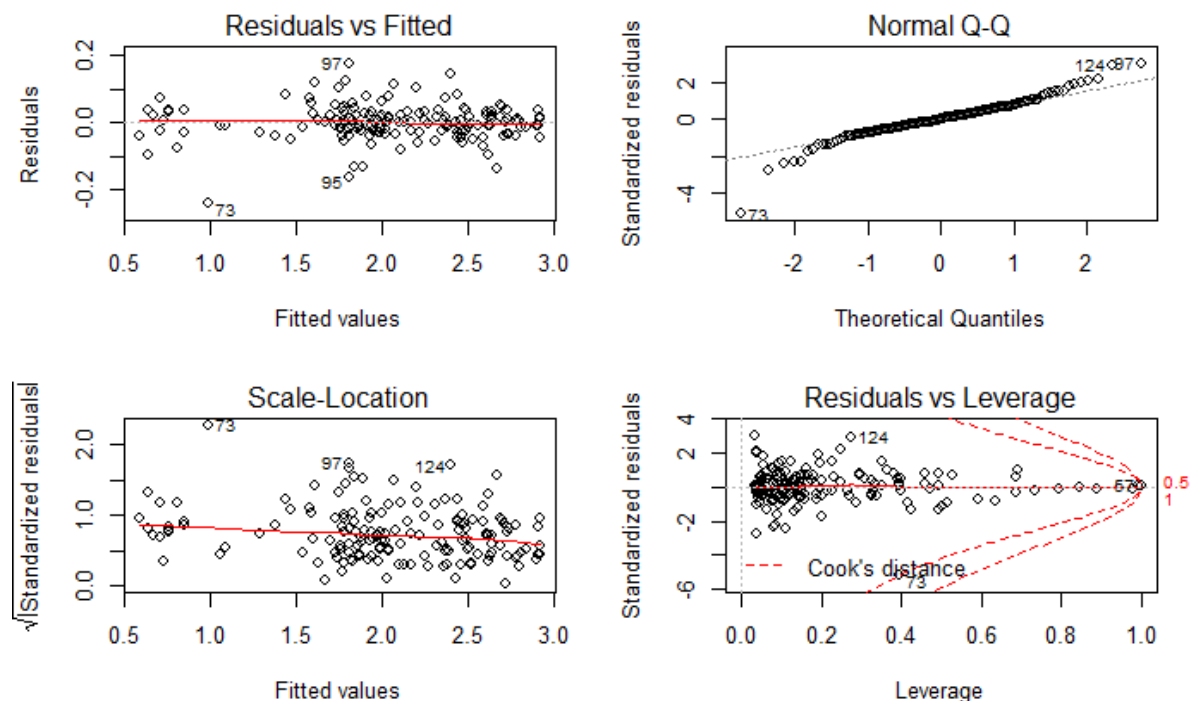
	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	2.349e+00	1.596e-01	14.721	< 2e-16	***
Length1	-3.355e-02	3.999e-02	-0.839	0.40301	
Length3	4.449e-02	3.924e-02	1.134	0.25910	
weight	3.006e-04	1.672e-04	1.797	0.07469	.
width	-1.071e-03	4.677e-02	-0.023	0.98176	
SpeciesParkki	2.105e-01	8.410e-01	0.250	0.80272	
SpeciesPerch	-5.468e-01	1.920e-01	-2.847	0.00515	**
SpeciesPike	-1.256e+00	2.224e-01	-5.647	1.04e-07	***
SpeciesRoach	-6.130e-01	3.850e-01	-1.592	0.11383	
SpeciesSmelt	-1.092e+00	8.441e-01	-1.293	0.19833	
SpeciesWhitefish	-5.860e-01	4.201e-01	-1.395	0.16552	

Length1:SpeciesParkki	-3.357e-01	5.093e-01	-0.659	0.51111
Length1:SpeciesPerch	-5.380e-03	4.862e-02	-0.111	0.91207
Length1:SpeciesPike	1.556e-01	5.698e-02	2.731	0.00724 **
Length1:SpeciesRoach	-6.076e-03	5.983e-02	-0.102	0.91928
Length1:SpeciesSmelt	1.147e-01	2.026e-01	0.566	0.57247
Length1:SpeciesWhitefish	4.139e-01	9.449e-01	0.438	0.66209
Length3:SpeciesParkki	3.091e-01	4.300e-01	0.719	0.47363
Length3:SpeciesPerch	3.156e-02	4.669e-02	0.676	0.50038
Length3:SpeciesPike	-1.616e-01	5.892e-02	-2.744	0.00697 **
Length3:SpeciesRoach	2.561e-02	5.696e-02	0.450	0.65374
Length3:SpeciesSmelt	-7.186e-02	1.716e-01	-0.419	0.67615
Length3:SpeciesWhitefish	-4.084e-01	9.177e-01	-0.445	0.65709
Weight:SpeciesParkki	3.332e-04	2.144e-03	0.155	0.87674
Weight:SpeciesPerch	-9.318e-04	1.908e-04	-4.885	3.10e-06 ***
Weight:SpeciesPike	-6.064e-05	2.453e-04	-0.247	0.80511
Weight:SpeciesRoach	-8.811e-04	4.527e-04	-1.947	0.05383 .
Weight:SpeciesSmelt	2.027e-02	1.896e-02	1.069	0.28708
Weight:SpeciesWhitefish	2.482e-04	1.413e-03	0.176	0.86079
Width:SpeciesParkki	-8.575e-02	3.272e-01	-0.262	0.79372
Width:SpeciesPerch	1.131e-01	5.203e-02	2.174	0.03156 *
Width:SpeciesPike	1.636e-01	6.434e-02	2.542	0.01224 *
Width:SpeciesRoach	1.237e-01	9.305e-02	1.330	0.18610
Width:SpeciesSmelt	7.723e-03	1.468e-01	0.053	0.95814
Width:SpeciesWhitefish	7.343e-02	6.893e-02	1.065	0.28882

---  
 Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.0593 on 125 degrees of freedom  
 Multiple R-squared: 0.9915, Adjusted R-squared: 0.9892  
 F-statistic: 431.1 on 34 and 125 DF, p-value: < 2.2e-16

### Diagnostic plot for log-transformation



## Summary of square root transformation

```
> summary(m2_new_sqrt)
```

Call:

```
lm(formula = sqrt(Height) ~ Length1 + Length3 + Weight + Width +  
    Species + Length1:Species + Length3:Species + Weight:Species +  
    Width:Species, data = fish_new)
```

Residuals:

Min	1Q	Median	3Q	Max
-0.19293	-0.03066	-0.00114	0.03244	0.19189

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	3.1516766	0.1827855	17.242	< 2e-16	***
Length1	-0.0604867	0.0458014	-1.321	0.189037	
Length3	0.0797628	0.0449504	1.774	0.078421	.
Weight	0.0006116	0.0001915	3.193	0.001780	**
Width	0.0059917	0.0535684	0.112	0.911120	
SpeciesParkki	0.5097860	0.9632515	0.529	0.597581	
SpeciesPerch	-0.8197133	0.2199513	-3.727	0.000292	***
SpeciesPike	-1.6786645	0.2547113	-6.590	1.1e-09	***
SpeciesRoach	-0.8322741	0.4409262	-1.888	0.061404	.
SpeciesSmelt	-1.4097205	0.9668320	-1.458	0.147326	
SpeciesWhitefish	-0.8203850	0.4811822	-1.705	0.090690	.
Length1:SpeciesParkki	-0.5136277	0.5833997	-0.880	0.380330	
Length1:SpeciesPerch	0.0494328	0.0556893	0.888	0.376432	
Length1:SpeciesPike	0.2140264	0.0652655	3.279	0.001348	**
Length1:SpeciesRoach	0.0002629	0.0685310	0.004	0.996946	
Length1:SpeciesSmelt	0.1077012	0.2320965	0.464	0.643429	
Length1:SpeciesWhitefish	0.5943708	1.0823247	0.549	0.583874	
Length3:SpeciesParkki	0.4642417	0.4925457	0.943	0.347737	
Length3:SpeciesPerch	-0.0302125	0.0534805	-0.565	0.573137	
Length3:SpeciesPike	-0.2303602	0.0674845	-3.414	0.000865	***
Length3:SpeciesRoach	0.0128315	0.0652449	0.197	0.844408	
Length3:SpeciesSmelt	-0.0930750	0.1965726	-0.473	0.636690	
Length3:SpeciesWhitefish	-0.5919312	1.0511367	-0.563	0.574352	
Weight:SpeciesParkki	0.0011388	0.0024557	0.464	0.643651	
Weight:SpeciesPerch	-0.0008691	0.0002185	-3.978	0.000117	***
Weight:SpeciesPike	-0.0002366	0.0002809	-0.842	0.401335	
Weight:SpeciesRoach	-0.0010488	0.0005185	-2.023	0.045220	*
Weight:SpeciesSmelt	0.0163800	0.0217163	0.754	0.452104	
Weight:SpeciesWhitefish	0.0002786	0.0016180	0.172	0.863558	
Width:SpeciesParkki	-0.2191226	0.3748147	-0.585	0.559860	
Width:SpeciesPerch	0.1193281	0.0595998	2.002	0.047432	*
Width:SpeciesPike	0.2230943	0.0736997	3.027	0.003000	**
Width:SpeciesRoach	0.1472318	0.1065817	1.381	0.169621	
Width:SpeciesSmelt	0.0043668	0.1681754	0.026	0.979326	
Width:SpeciesWhitefish	0.0999413	0.0789549	1.266	0.207938	

---

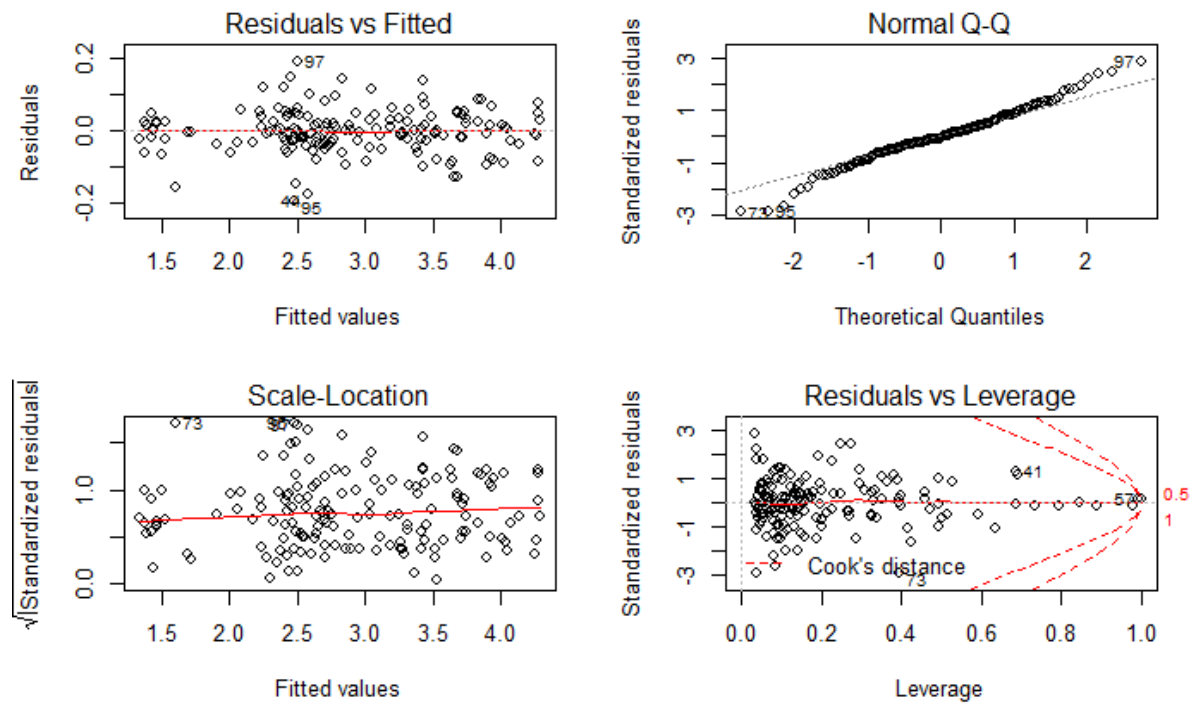
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.06792 on 125 degrees of freedom

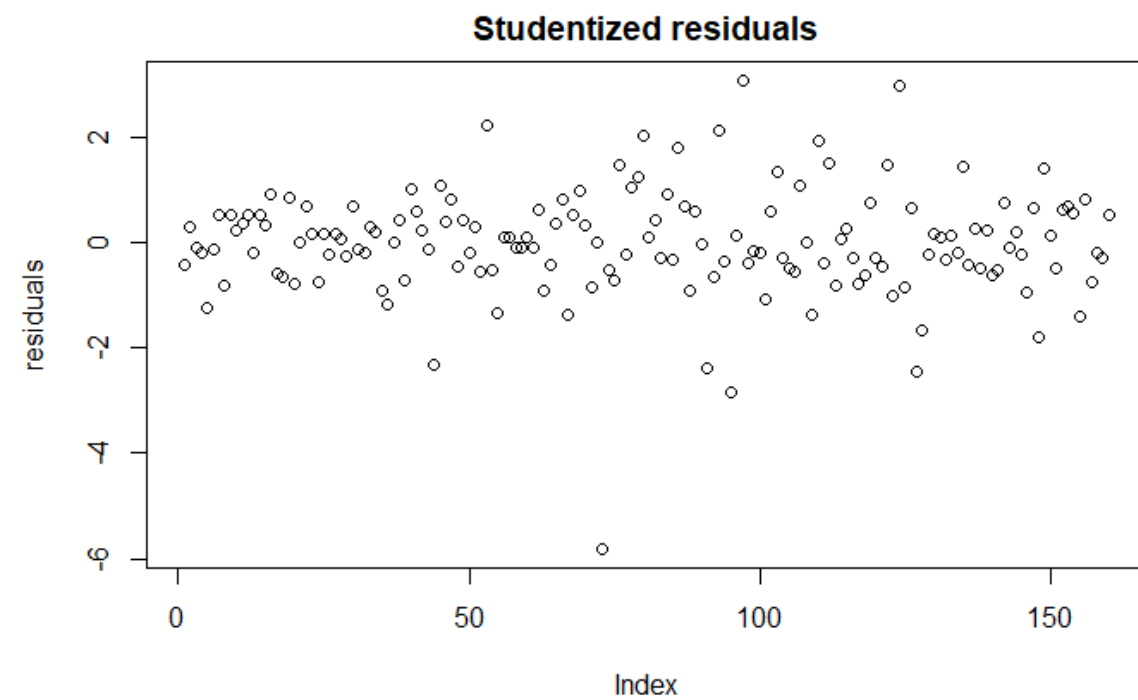
Multiple R-squared: 0.9935, Adjusted R-squared: 0.9918

F-statistic: 565.3 on 34 and 125 DF, p-value: < 2.2e-16

### Diagnostic plot for square root transformation



### Studentized residual plot for log-transformation





## Summary of robust regression with removed outliers and log transformation

```
> summary(rr.m2_new_log2)
```

```
Call: rlm(formula = log(Height) ~ Length1 + Length3 + weight + width +  
  Species + Length1:Species + Length3:Species + weight:Species +  
  width:Species, data = fish_infremove2, psi = psi.huber)
```

```
Residuals:  
      Min       1Q   Median       3Q      Max  
-0.1315655 -0.0204992  0.0001975  0.0209630  0.1272408
```

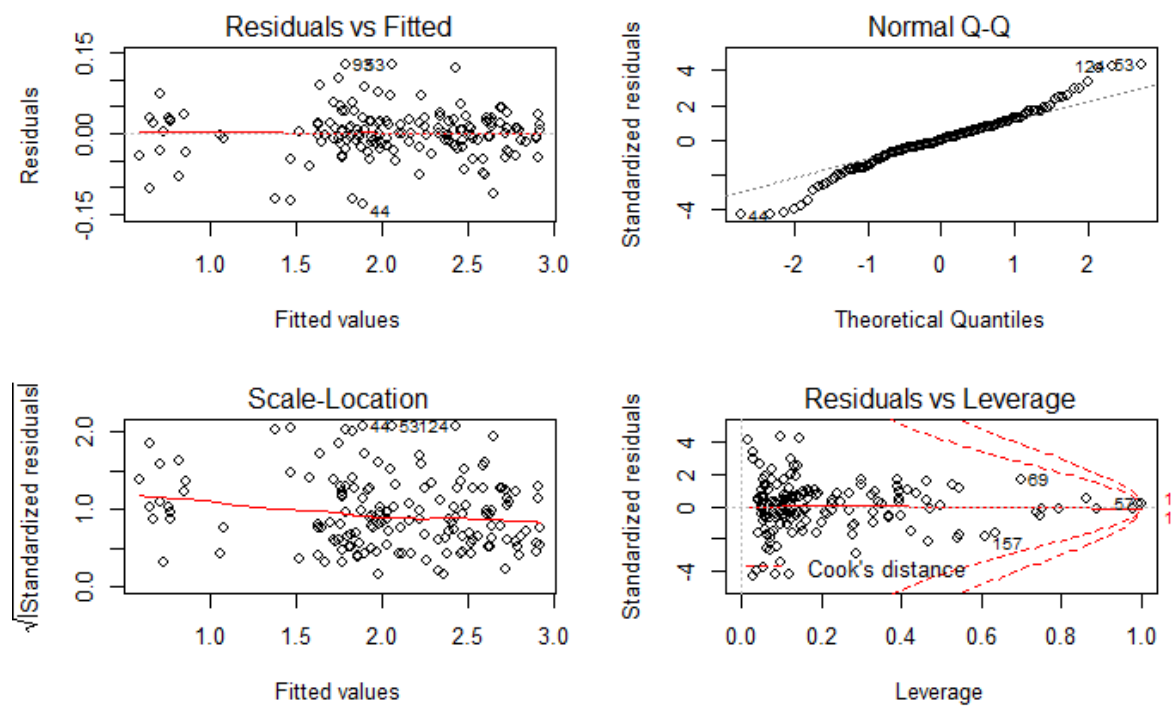
Coefficients:

	Value	Std. Error	t value
(Intercept)	2.3305	0.1184	19.6757
Length1	-0.0309	0.0297	-1.0395
Length3	0.0413	0.0291	1.4187
weight	0.0003	0.0001	2.3421
width	0.0058	0.0347	0.1681
SpeciesParkki	0.1169	0.6242	0.1873
SpeciesPerch	-0.7183	0.1438	-4.9971
SpeciesPike	-1.2689	0.1651	-7.6881
SpeciesRoach	-0.6247	0.2857	-2.1864
SpeciesSmelt	-0.9387	0.6265	-1.4984
SpeciesWhitefish	-0.5672	0.3118	-1.8192
Length1:SpeciesParkki	-0.3176	0.3780	-0.8401
Length1:SpeciesPerch	0.0391	0.0366	1.0666
Length1:SpeciesPike	0.1555	0.0423	3.6757
Length1:SpeciesRoach	-0.0052	0.0444	-0.1162
Length1:SpeciesSmelt	0.1227	0.1504	0.8156
Length1:SpeciesWhitefish	0.4112	0.7013	0.5864
Length3:SpeciesParkki	0.2950	0.3192	0.9244
Length3:SpeciesPerch	-0.0207	0.0354	-0.5842
Length3:SpeciesPike	-0.1634	0.0437	-3.7370
Length3:SpeciesRoach	0.0254	0.0423	0.6001
Length3:SpeciesSmelt	-0.0719	0.1274	-0.5644
Length3:SpeciesWhitefish	-0.4052	0.6811	-0.5949
weight:SpeciesParkki	-0.0002	0.0016	-0.1198
weight:SpeciesPerch	-0.0006	0.0001	-4.2094
weight:SpeciesPike	0.0000	0.0002	-0.0761
weight:SpeciesRoach	-0.0011	0.0003	-3.1452
weight:SpeciesSmelt	0.0176	0.0141	1.2483
weight:SpeciesWhitefish	0.0003	0.0010	0.2462
width:SpeciesParkki	-0.0277	0.2429	-0.1141
width:SpeciesPerch	0.1099	0.0387	2.8389
width:SpeciesPike	0.1661	0.0478	3.4778
width:SpeciesRoach	0.1326	0.0691	1.9194
width:SpeciesSmelt	0.0025	0.1090	0.0226
width:SpeciesWhitefish	0.0665	0.0512	1.3003

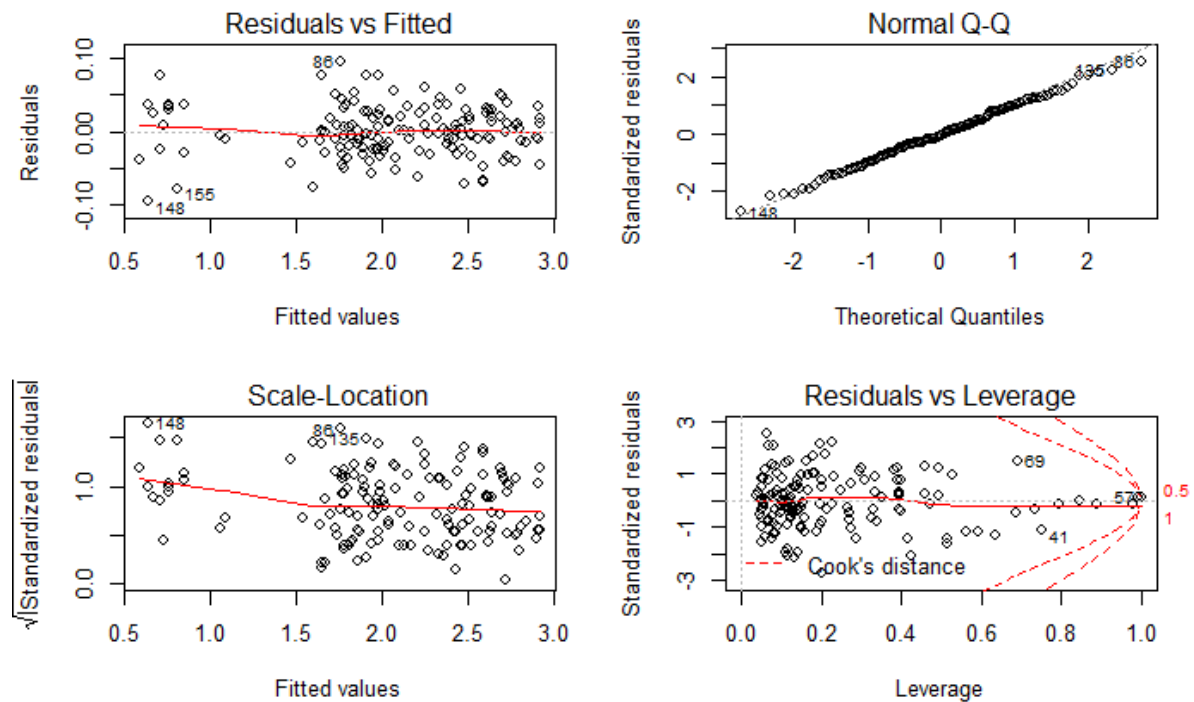
Residual standard error: 0.03108 on 122 degrees of freedom



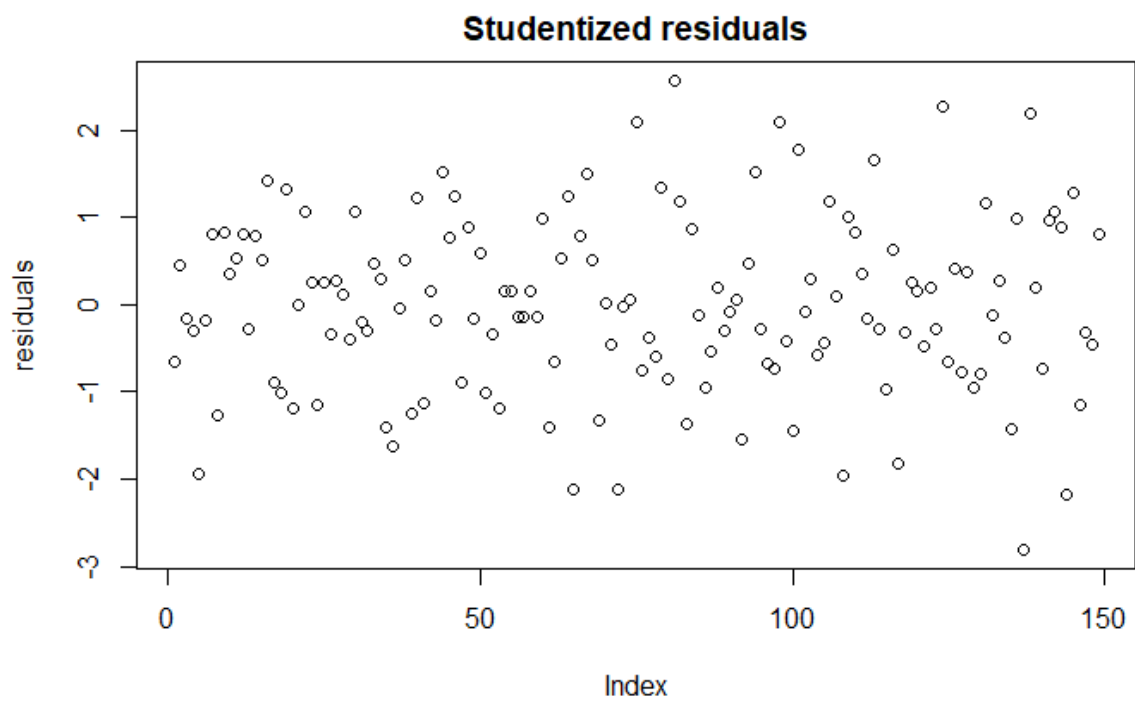
### Diagnostic plot of robust regression with removed outliers and log transformation



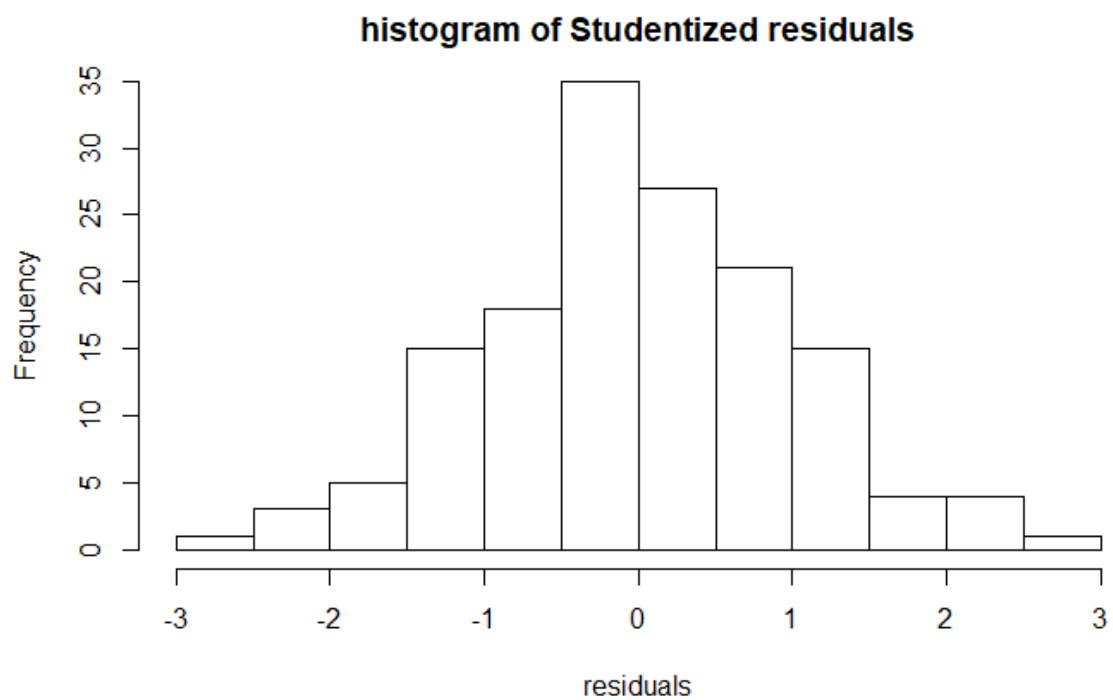
### Diagnostic plot of final model



**Studentized residual plot of final model**



**Histogram of Studentized residual in final model**



## ANOVA of final model

```
> anova(m2_new_log3)
```

Analysis of Variance Table

Response: log(Height)

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
Length1	1	21.7693	21.7693	14519.4242	< 2.2e-16	***
Length3	1	14.6477	14.6477	9769.5444	< 2.2e-16	***
weight	1	0.0104	0.0104	6.9599	0.0095000	**
width	1	7.4687	7.4687	4981.3598	< 2.2e-16	***
Species	6	3.9284	0.6547	436.6806	< 2.2e-16	***
Length1:Species	6	0.1449	0.0241	16.1063	2.377e-13	***
Length3:Species	6	0.0191	0.0032	2.1217	0.0560683	.
weight:Species	6	0.0371	0.0062	4.1185	0.0008847	***
width:Species	6	0.0318	0.0053	3.5339	0.0030296	**
Residuals	114	0.1709	0.0015			

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

## variance inflation factor (VIF)

```
> vif(m2_new_log3)
```

	GVIF	Df	GVIF^(1/(2*Df))
Length1	6.743672e+03	1	82.11986
Length3	8.769305e+03	1	93.64457
weight	1.496073e+02	1	12.23141
width	2.593501e+02	1	16.10435
Species	2.311278e+16	6	23.10225
Length1:Species	2.913187e+23	6	90.23272
Length3:Species	5.436983e+23	6	95.04877
weight:Species	4.102141e+13	6	13.62754
width:Species	4.276361e+15	6	20.07195