

Title: Effects of maternal host versus larval host plant on developmental time

# Methods:

The input data set was in csv format. Two different host plants were investigated for maternal and larval development phases. To investigate the data is normally distributed, aov() function was performed on analysis of variance. Then, ANOVA analysis was done on the input data. There were means, standard deviation of error and mathematics computation done on the data.

Table 1: The head part of the input data set

	LarvalID	LarvalHost	Sex	MaternalHost	MotherID	DevelopmentTime	AdultWeight	GrowthRate
1	14	BarbareaL	M	BarbareaM	9	21	64.2	0.08607
2	22	BarbareaL	M	BarbareaM	9	22	60.8	0.08109
3	23	BarbareaL	F	BarbareaM	9	22	53.0	0.07838
4	24	BarbareaL	M	BarbareaM	9	22	67.8	0.08324
5	25	BarbareaL	F	BarbareaM	9	22	57.1	0.07985
6	30	BarbareaL	F	BarbareaM	9	23	88.9	0.08473

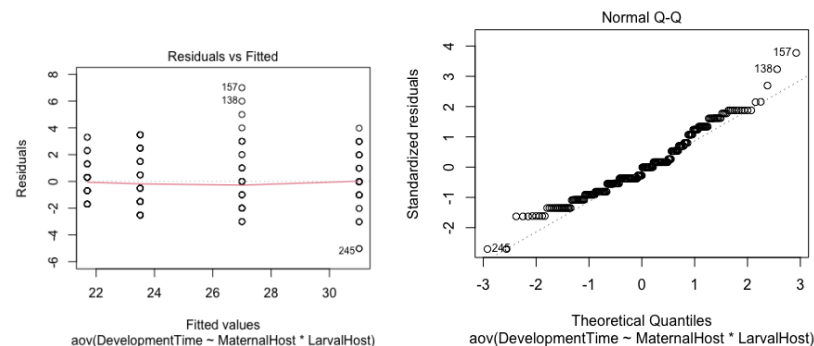


Figure 1. Residuals vs Fitted values(left), Standardized residuals vs Theoretical Quantiles (right)

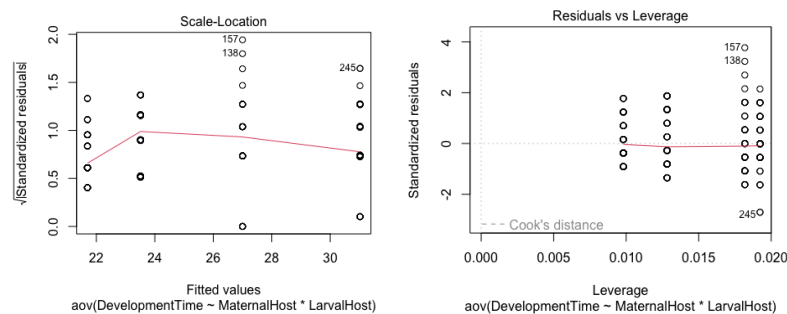


Figure 2. Square root of Standardized residuals vs Fitted values (left), Standardized residuals vs aov(DevelopmentTime~MaternalHost\*LarvalHost (right)

Table 2. Summary of linear model -> lm(formula = DevelopmentTime ~ MaternalHost \* LarvalHost, data = dat)

```

Residuals:
    Min       1Q   Median       3Q      Max
-5.0192 -1.5128 -0.5128  1.0000  7.0000

Coefficients:
            (Intercept)      21.6961      0.1854 117.033 < 2e-16 ***
MaternalHostBerteroM      1.8167      0.2816   6.451 4.80e-10 ***
LarvalHostBerteroL       5.3039      0.3132  16.934 < 2e-16 ***
MaternalHostBerteroM:LarvalHostBerteroL  2.2025      0.4588   4.801 2.56e-06 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 1.872 on 283 degrees of freedom
Multiple R-squared:  0.7734,    Adjusted R-squared:  0.771
F-statistic: 322.1 on 3 and 283 DF,  p-value: < 2.2e-16

```

**Table 3. ANOVA of maternal host versus larval host and interactions between host plant**

Analysis of Variance Table

Response: DevelopmentTime

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
MaternalHost	1	623.61	623.61	177.90	< 2.2e-16 ***
LarvalHost	1	2682.41	2682.41	765.21	< 2.2e-16 ***
MaternalHost:LarvalHost	1	80.80	80.80	23.05	2.561e-06 ***
Residuals	283	992.05	3.51		

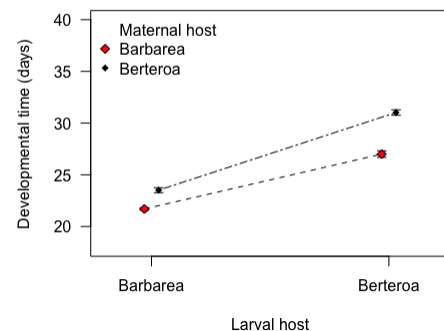
---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

**Table 4. Mean values between maternal host versus larval host plant**

	BarbareaL	BerteroaL
BarbareaM	21.69608	27.00000
BerteroaM	23.51282	31.01923

**Table 5. Column and row means**

	BarbareaL	BerteroaL
	22.60445	29.00962
BarbareaM		
	24.34804	27.26603



**Figure 3. Larval development time depending on larval and maternal host plants**

### Results:

The input data set contained of 8 variables with 287 objects (Table 1). The analysis of variance suggested that the response variable "DevelopmentTime" is influenced by both "MaternalHost" and "LarvalHost," as well as their interaction ("MaternalHost\*LarvalHost"), and the data was normally distributed (Table 1, Figure 1-2). Residual standard error ( $= \sqrt{\frac{\sum(y - \hat{y})^2}{df}}$ ) is small (1.872 on 283 degree of freedom) indicated that the data fitted well with the regression model (Table 2). The sum of square values within the larval host is the largest compared to the maternal host, suggesting a larger variance and there were effects on both the larval and maternal hosts ( $F_{1, 283} = 765.21, 177.9$ ). On the other hand, the variance for the interaction between maternal and larval host is the smallest, indicating that there was a lesser effect on the interaction between two hosts (Table 3). Effect size implied how meaningful the relationship between variables or the difference between groups is. Computation by using the mean (Table 4) by the row and the column were reported (Table 5). Larval grown faster (22.1%) on the Barbarea host and had shorter developmental time (mean development time = 22.6). Similarly larval whose mother were grown on the Barbarea host grown 10.7% faster than the Berteroa host plant (mean development time = 24.3 and 27.3) (Table 5). Barbarea host appeared to be the better oviposit place for the larval, in which the larval's mother had grown either on the Barbarea or Berteroa host. The development time is smaller on Barbarea (19.6%) compared to the Berteroa host (24.1%) (Table 4).