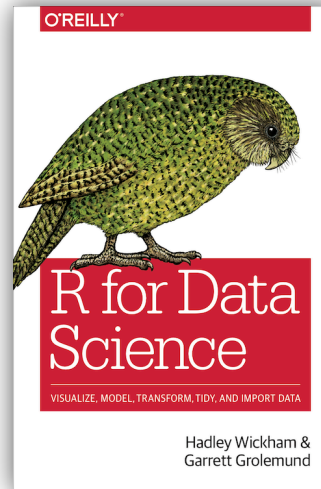
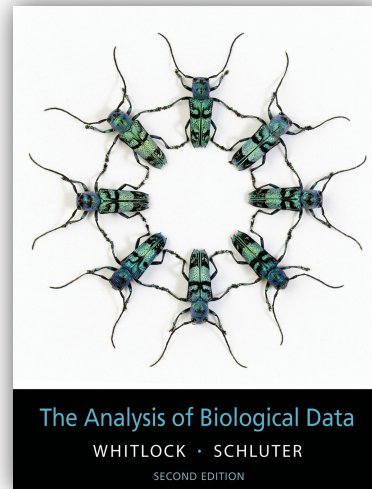


Data Science in Bioinformatics

Palle Villesen & Thomas Bataillon



Outline for week 12

- Chapter 18
 - Mole rats & ANCOVA
 - Exercises wrap up dn/ds and & expression &...
- Thursday session is CANCELLED
- The final assignment is released

Body mass in lazy Mole rats



- Two castes
Workers
Infrequent workers (LAZY)

Regression

$Y \sim \text{Covariate}$

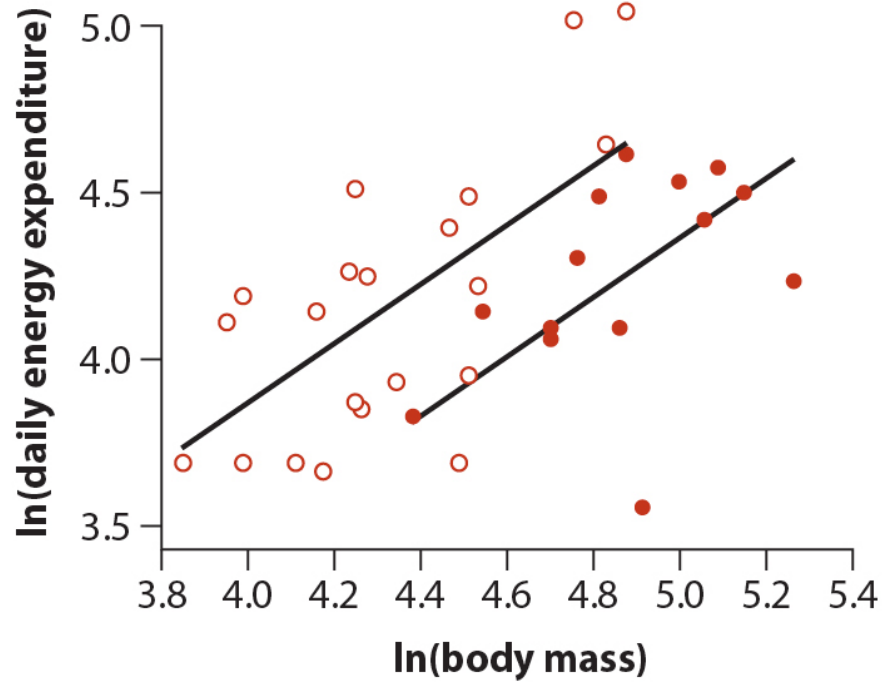
ANOVA

$Y \sim \text{Factor}$

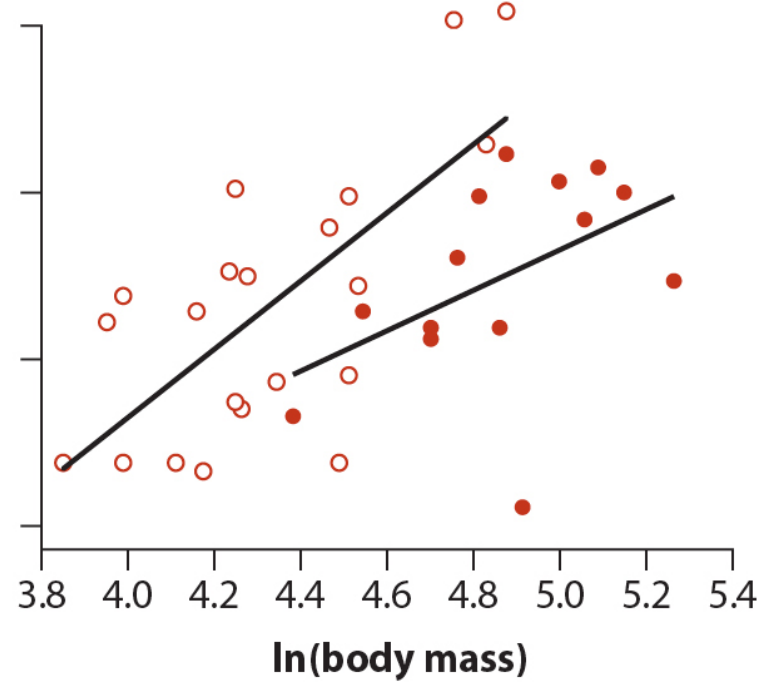
Ancova model

$Y \sim \text{Covariate} + \text{Factor}$

$$\text{ENERGY} = \text{CONSTANT} + \text{CASTE} + \text{MASS}$$

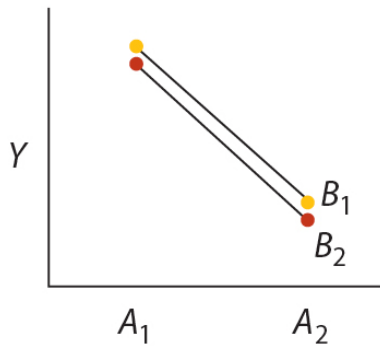


$$\text{ENERGY} = \text{CONSTANT} + \text{CASTE} + \text{MASS} + \text{CASTE} * \text{MASS}$$

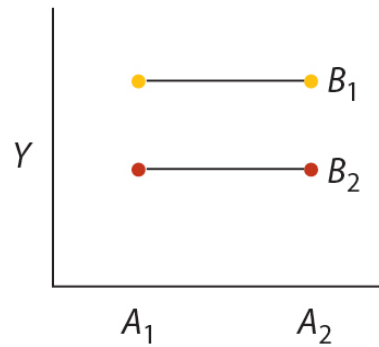


Models with/ without interaction

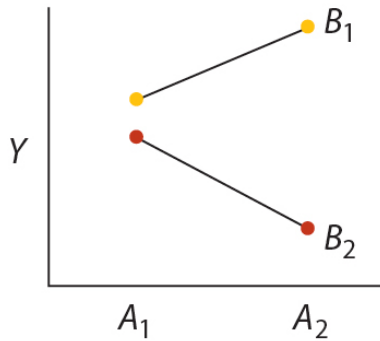
Main effect of A



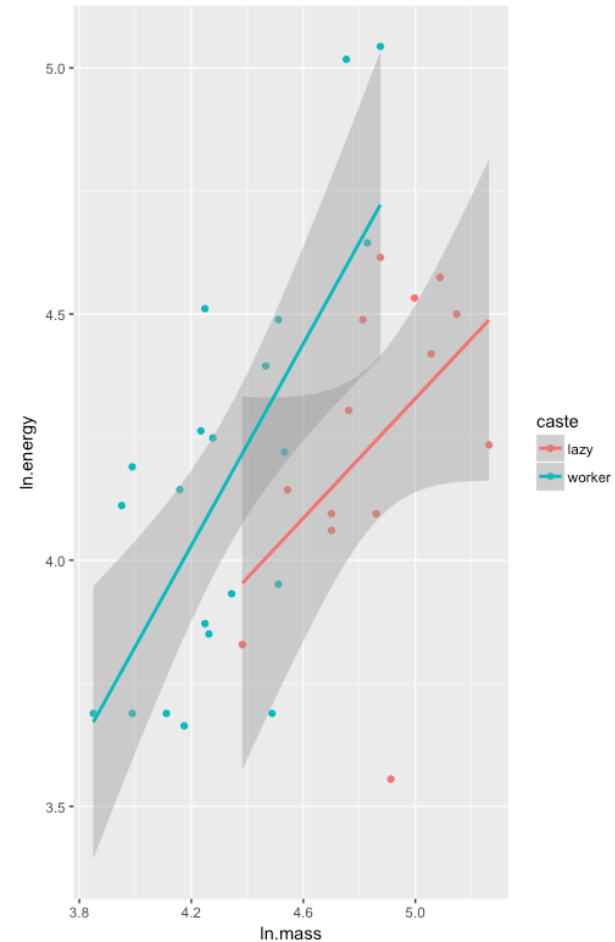
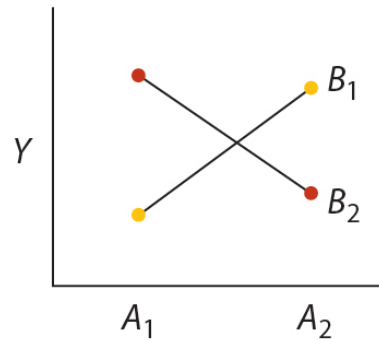
Main effect of B



Main effect of B
Interaction of A and B



Interaction of A and B



Issues when fitting models

Simple models

Easy to test

Easy to interpret

Easy to fool

(underlying extra covariates
not included in the model)

Multiple factors models

How to test effects?

type I, type II, type III

What model should I choose?

Sequential

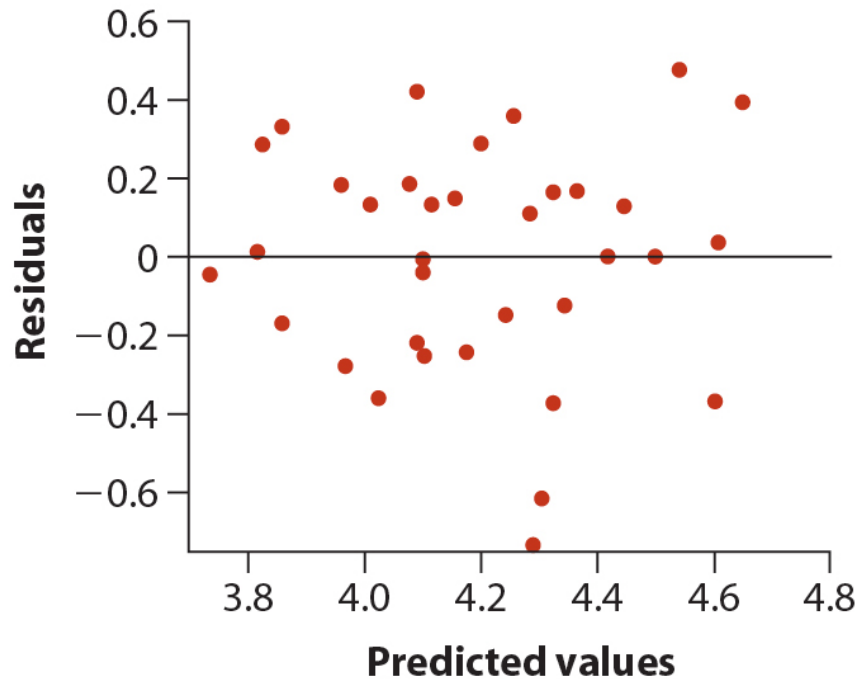
Search all models

Model selection

Model Averaging

→ See Machine Learning in
Bioinformatics

After fitting your model ...



Check your model

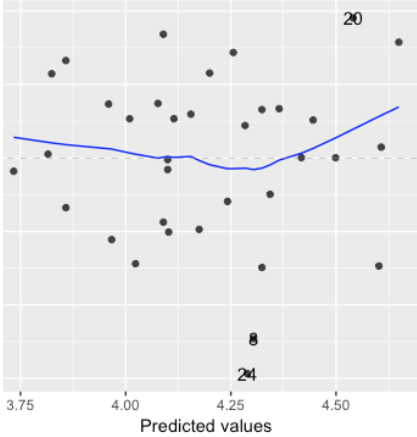
```
fit1 = lm()
```

```
plot(fit1)
```

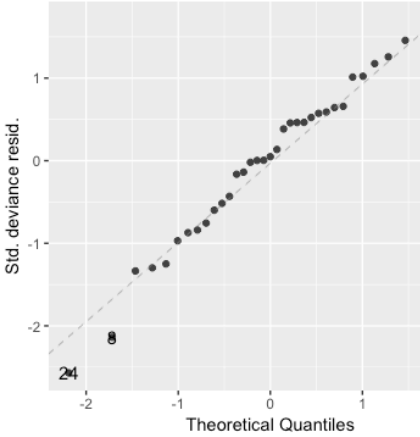
```
ggplot2::autoplot(fit)
```

... refit ... repeat

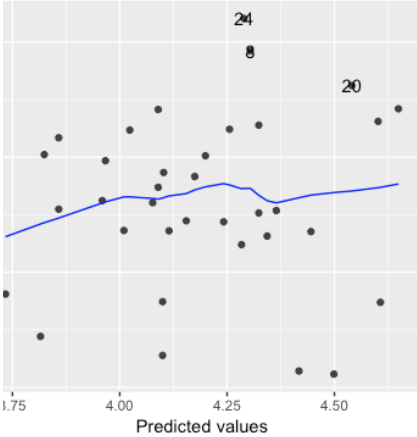
Residuals vs Fitted



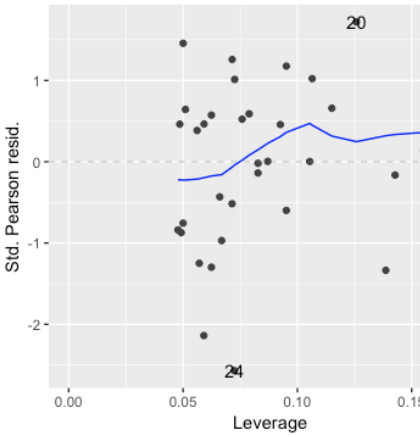
Normal Q-Q



scale-Location



Residuals vs Leverage



The last word ?



“The need for statisticians to reject the role of 'guardian of proven truth', and to resist attempts to provide once-for-all solutions and tidy over-unifications of the subject”

https://en.wikipedia.org/wiki/John_Tukey