

Generalised Linear Models (GLM): Part 1

Lecture Outline

Recap of the course so far

What are GLMs and why do we use them?

Components of a GLM

Maximum likelihood and GLMs

Fitting in R

Lecture Outline

Recap of the course so far

- EX1: Course so far

What are GLMs and why do we use them?

- EX2: Non-normal data

Components of a GLM

- EX3: Examples of non-normal data

Maximum likelihood and GLMs

Fitting in R

- EX4: Fit in R

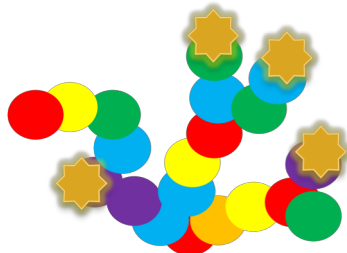
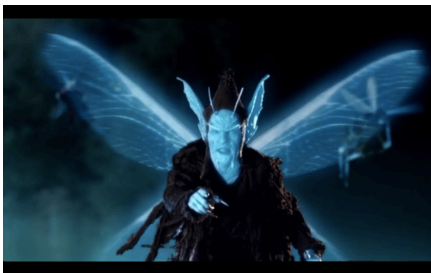
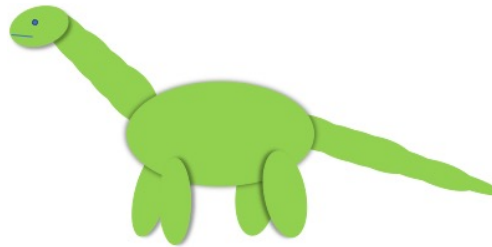
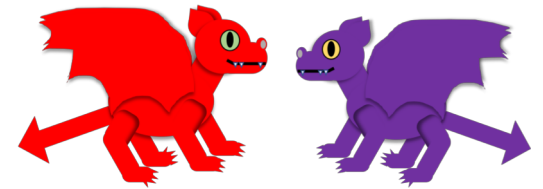
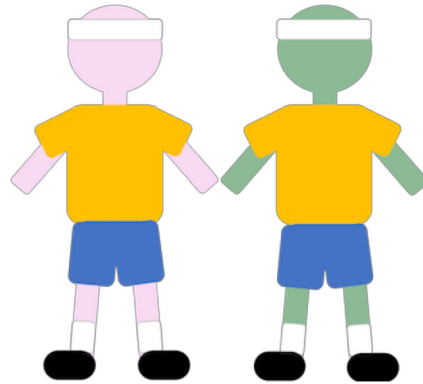
Reading

Chapter 8 – The New Statistics with R

Recap of the
course so far

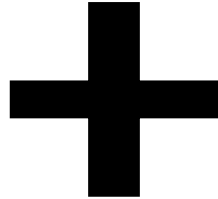
Exercise 1: What have we covered so far?

- Think about the previous weeks, write on your boards some of the topics we have covered.



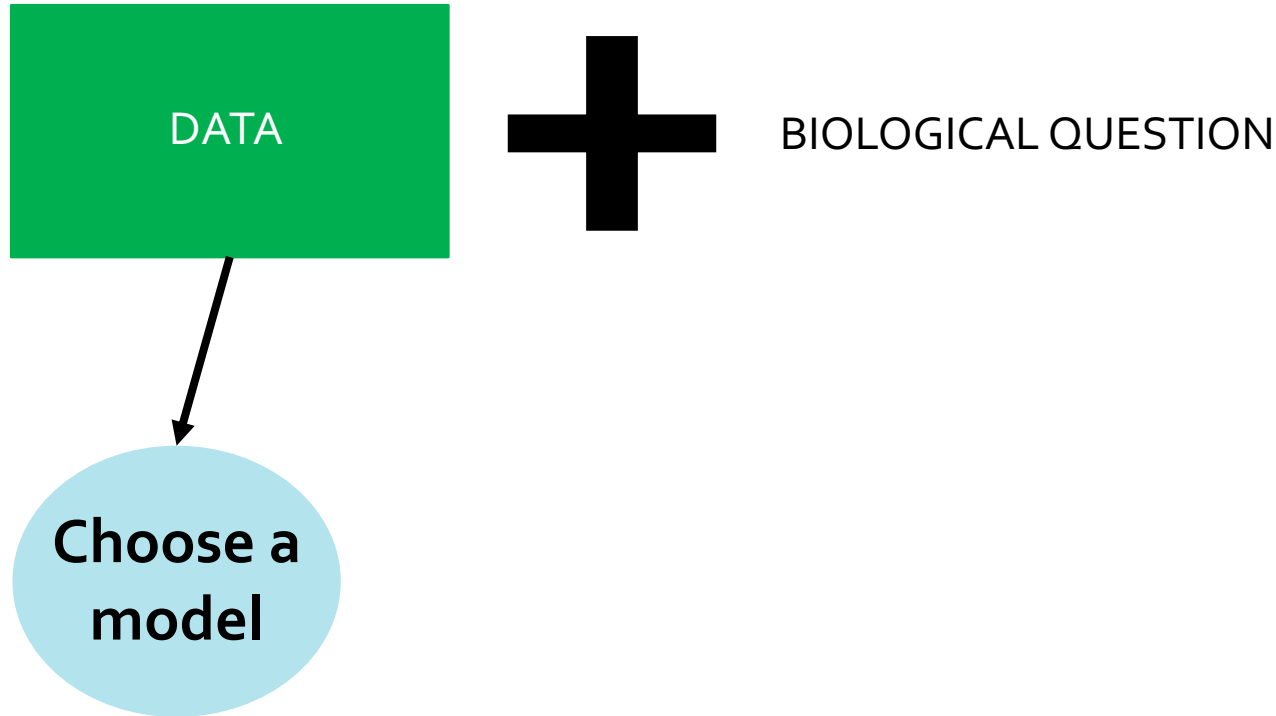
The modelling process

DATA



BIOLOGICAL QUESTION

The modelling process



Mathematical description of how the data were generated.

E.g.

- Distribution
- Linear equation (lines or groups)
- Defined by parameters

The modelling process

Inference

**Choose a
model**

**Get
estimates
of
parameters**

**E.g. Maximum
likelihood estimation**

Find the parameters that
give the highest
likelihood given the
data.

The modelling process

Inference

**Choose a
model**

**Get
estimates
of
parameters**

**Quantify
uncertainty
in
estimates**

The modelling process

**Choose a
model**

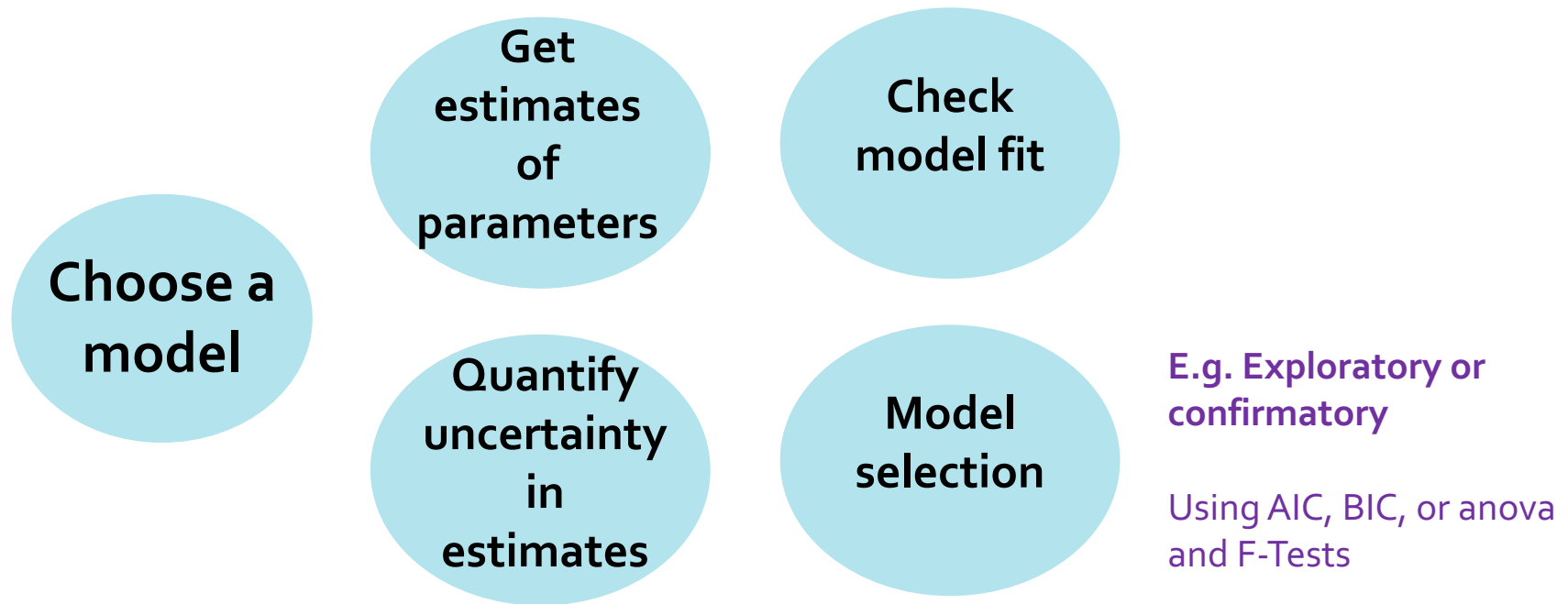
**Get
estimates
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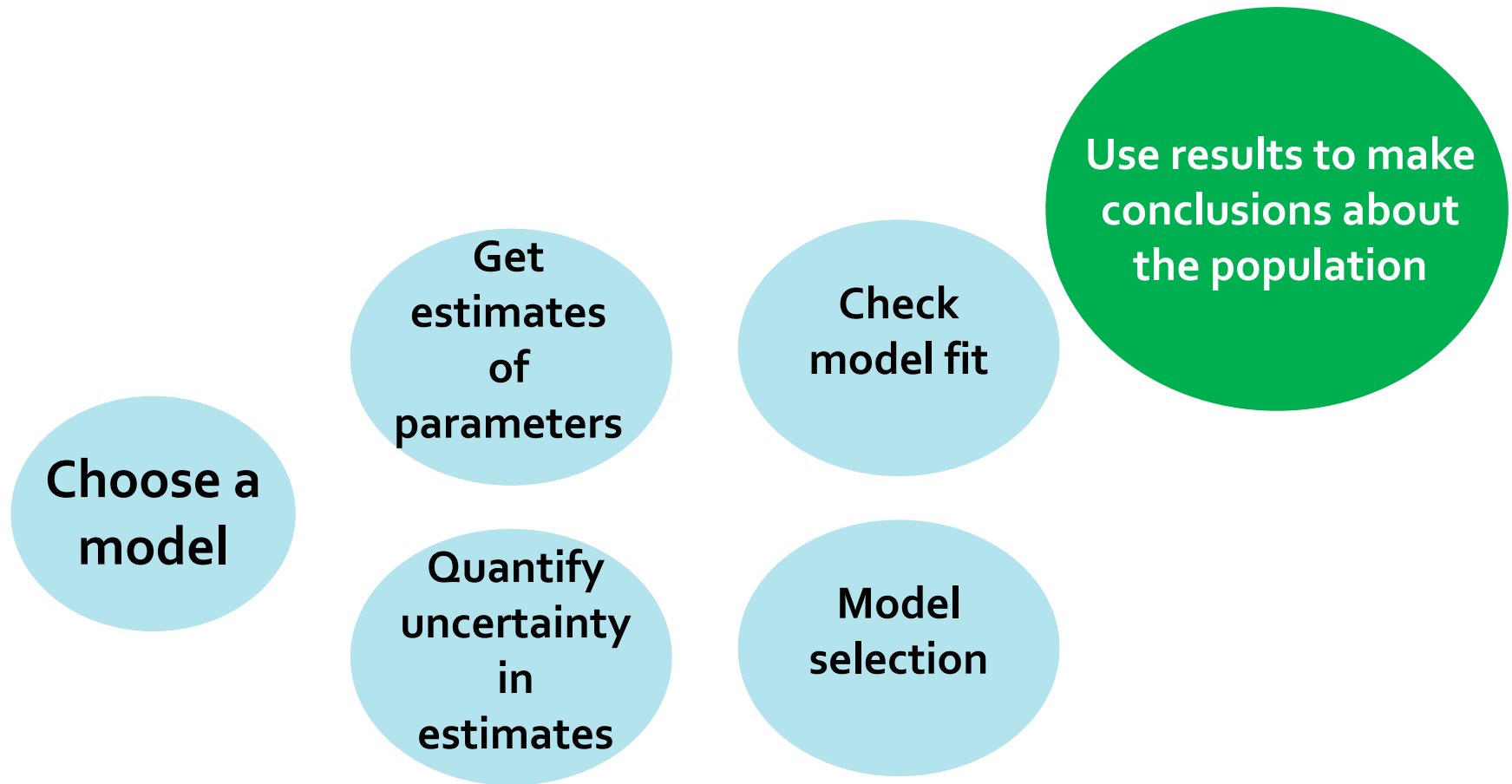
**Check
model fit**

E.g. Check assumptions
have been met

The modelling process



The modelling process



What are GLMs
and why do we
use them?

Linear models

Use linear equations to model a continuous response as a function of explanatory variables


Linear models

Use linear equations to model a continuous response as a function of explanatory variables

$$Y_i = \alpha + \beta X_i + \varepsilon_i$$

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linear predictor

error

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linear predictor

error

Systematic part **Random part**

Linear models

Assumptions:

- straight line (**linearity**)
- errors are independent
- errors have same variance (**homoscedasticity**)
- errors are normally distributed
- errors have zero mean

Exercise 2: Is a linear model appropriate?

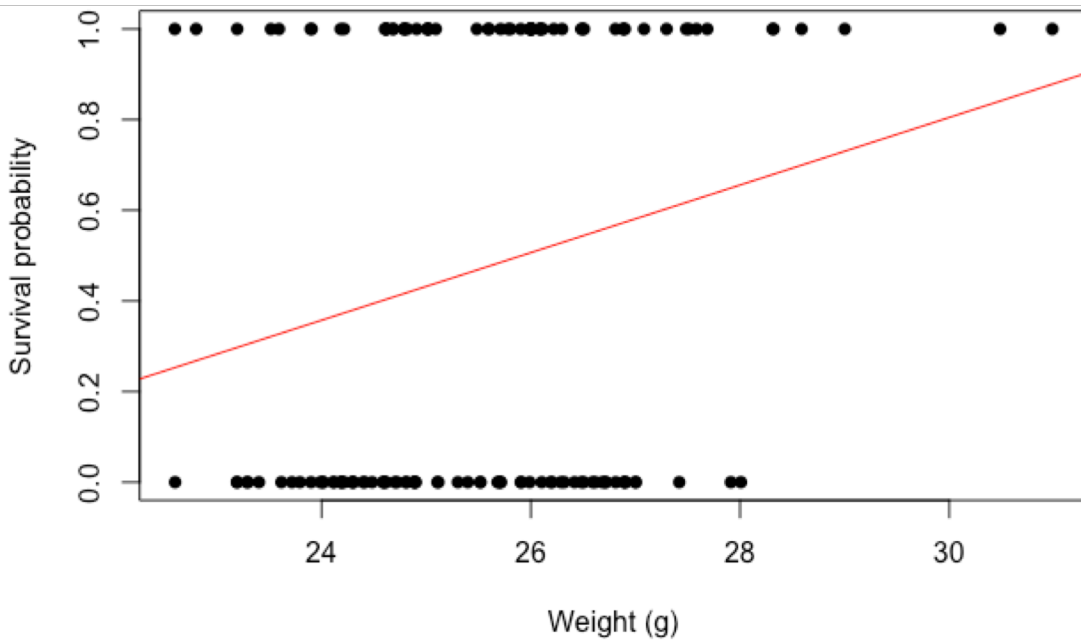
- Take a look at the three datasets on the next slides (you have data plotted with the modelled line from a linear model, residual vs fitted plot, and a Normal Q-Q plot).
- For each, answer the questions:

Is a linear model a suitable model for this data?

If not, why not?

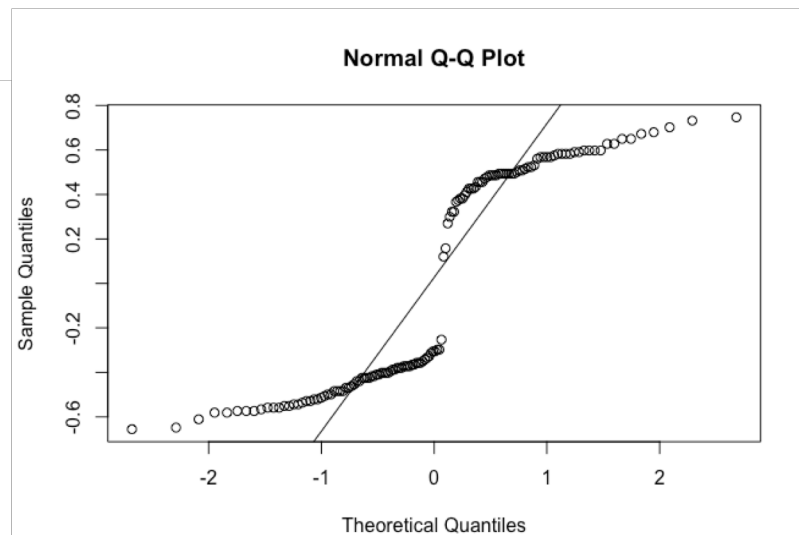
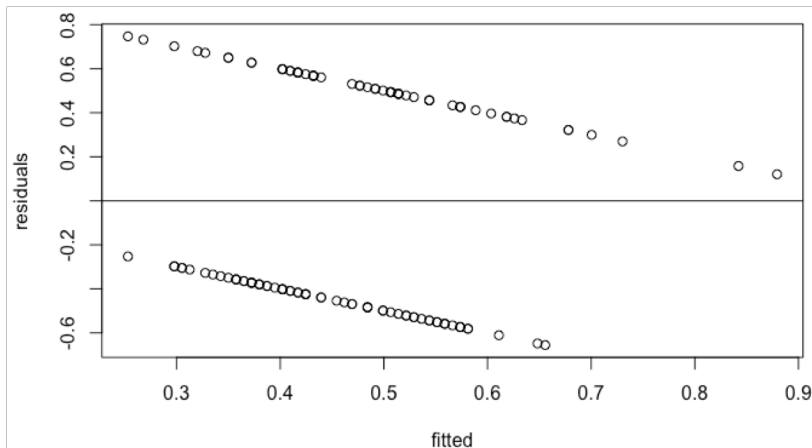
How could you improve it?

Example 1: Survival of sparrows

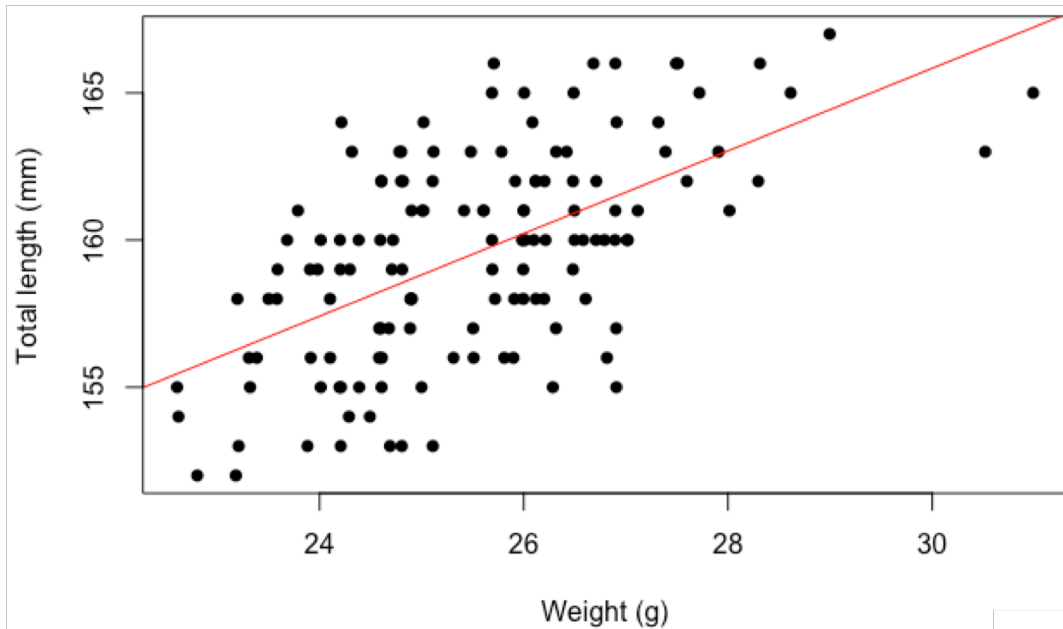


Question: How does body weight influence survival probability in sparrows?

Data: Response = whether the bird survived (1), or not (0).
Explanatory = body weight in grams

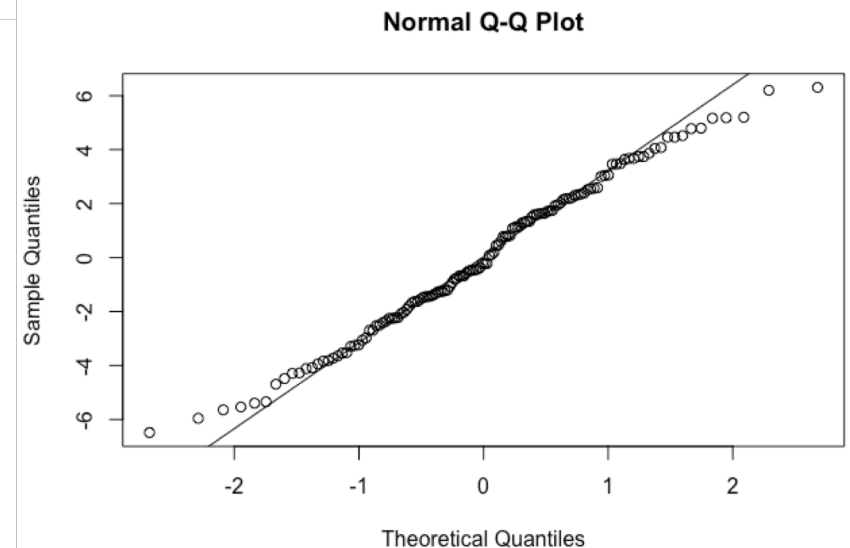
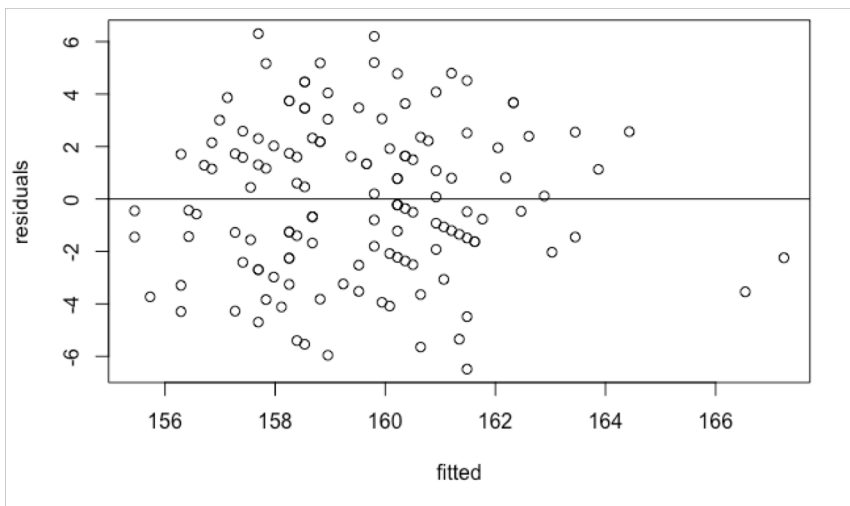


Example 2: Length and weight in sparrows

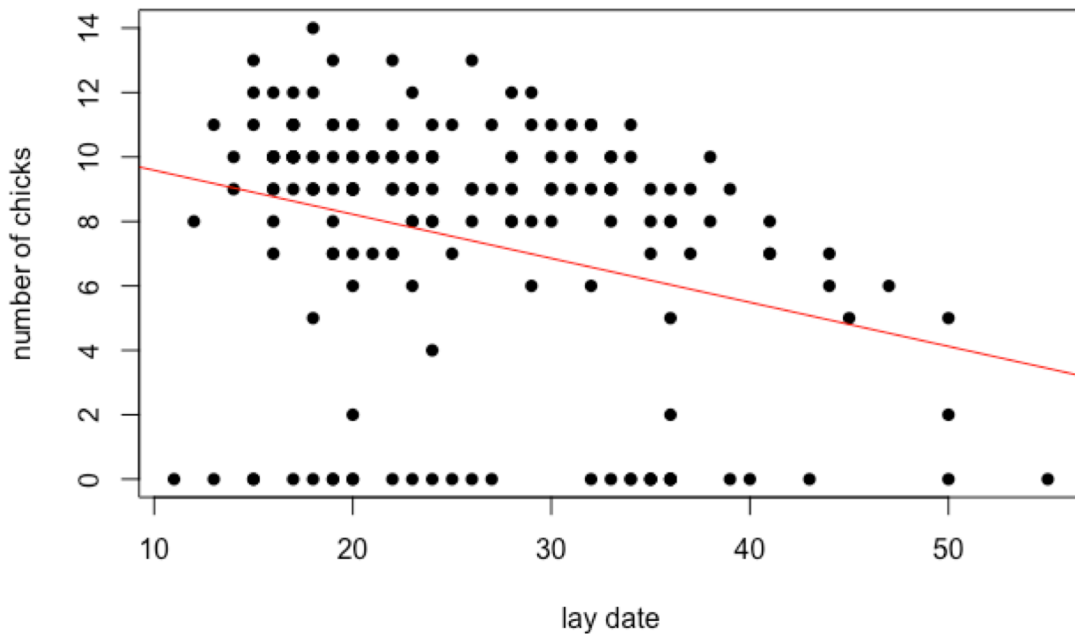


Question: How does body weight influence total length of the sparrows?

Data: Response = total length in mm. Explanatory = body weight in grams

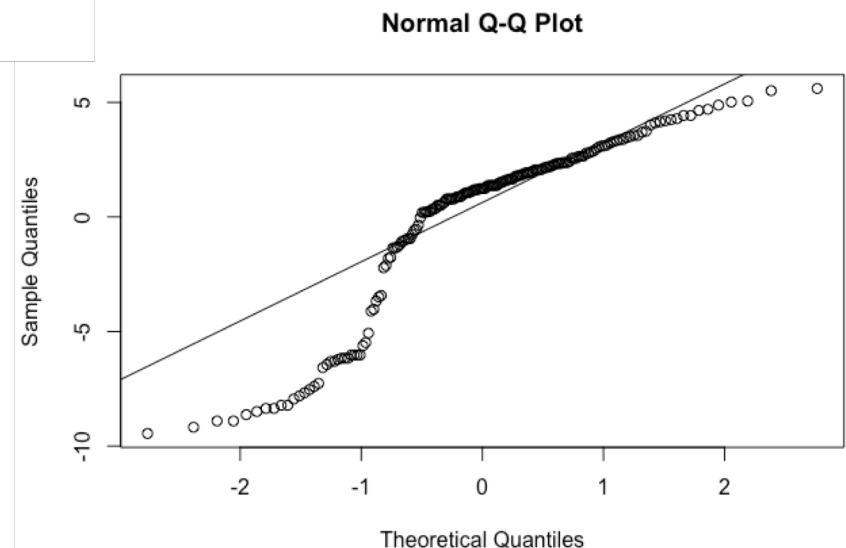
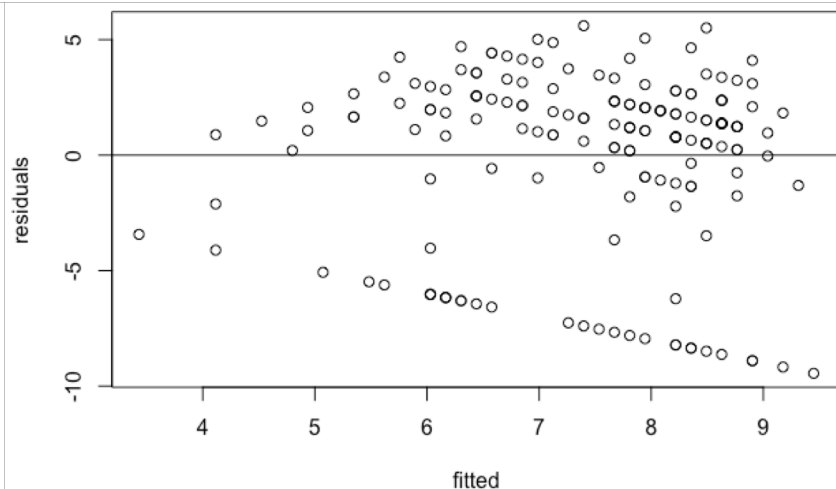


Example 3: Fledge success blue tits



Question: How does lay date influence the number of chicks that leave the nest?

Data: Response = number of chicks that fledge (leave nest alive). Explanatory = lay date (day since 1st April)



What to do with non-normality or non-linearity

Transformation of response?

Different, specialized models?

What to do with non-normality or non-linearity

Transformation of response?

Different, specialized models?

Or

Generalised linear models

A brief intro to Generalised Linear Models

Introduced in 1972 by Nelder and Wedderburn

<https://docs.ufpr.br/~taconeli/CE225/Artigo.pdf>

Can address variance and linearity in single model

Response unchanged

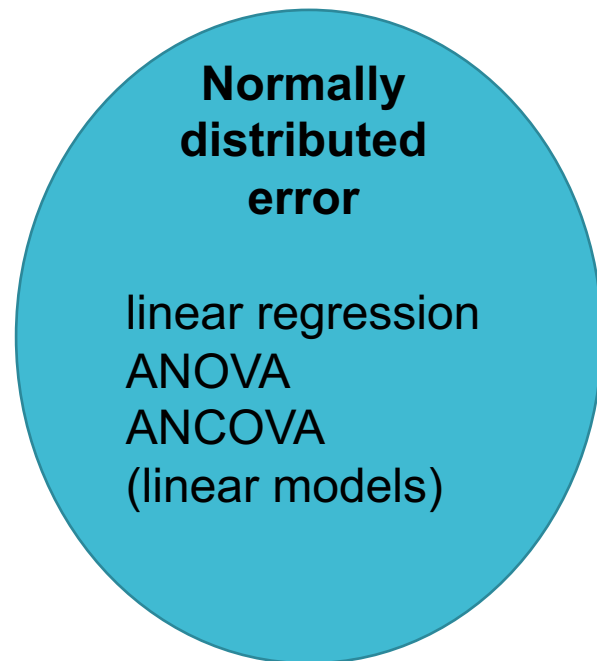
Luckily for us, very similar to `lm()` in R

Basis of many biological models

Key part of modern statistics!

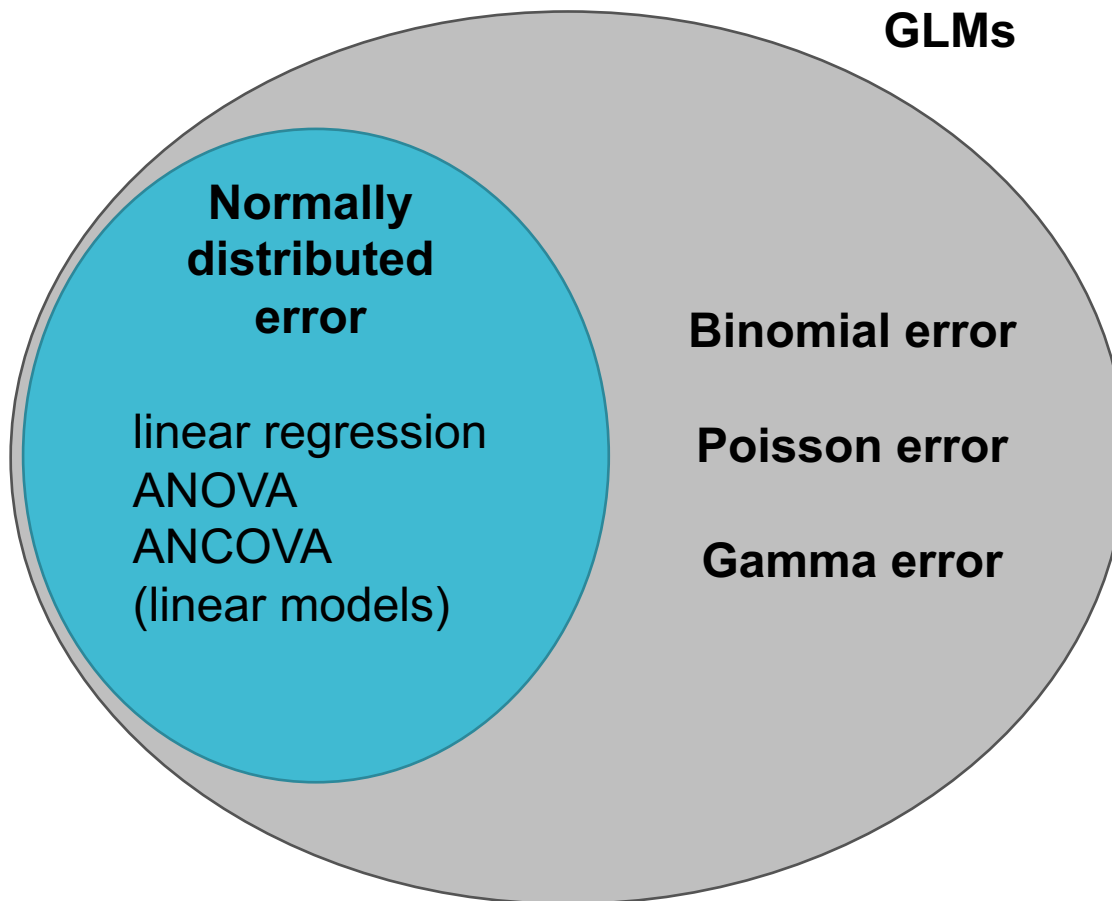
Generalised linear models

Similar to linear models but much more flexible



Generalised linear models

Similar to linear models but much more flexible



Biological examples

Clutch size

Sex ratio

Population size

**Number of plants
in a quadrat**

Two colour morphs

Biological examples

Clutch size

Sex ratio

Population size

**Number of plants
in a quadrat**

Two colour morphs

Counts and binary data

Exercise 3: Think of examples of non-normal data

- In your groups see if you can think of any other biological examples of non-normal data.
- This can be from your practical classes, just things you are interested in or anything else.
- Try and think of 3 examples in each group and write on white boards.
- Present one to the class.

Components of a GLM

Components of a GLM

Three main components of a GLM:

Random part

- the data (with an assumed distribution e.g. Binomial)

Systematic part

- the model for each data point (linear predictor) e.g. $\sum_j X_{ij}\beta_j$

The link function

- transforms the model (linear) onto scale of data e.g. $\log(\sum_j X_{ij}\beta_j)$

Random

Key bits to remember:

Think about the correct distribution for the data

GLM can use Normal, Binomial, Poisson, and Gamma

Different distributions use different link functions

Systematic

Key bits to remember:

This part is the same as a linear model

Key bits to remember:

Different distributions use different link functions

Which you use will alter the interpretation

Connects the Systematic part to the Random data

Describes how the mean depends on the linear predictor

e.g.

$$E(Y_i) = \log\left(\sum_j X_{ij}\beta_j\right)$$

Key bits to remember:


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Expected value of Y_i
(from Poisson
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log link

Maximum likelihood and GLMs

Definitions/synonyms

Explanatory variable = covariate = predictor

Normal distribution = Gaussian distribution

Dispersion = how wide or narrow a distribution is,
measured by variance or standard deviation

Parameter estimation reminder

Use maximum likelihood to estimate parameters

Likelihood is an equation that represents how the data were generated

Likelihood of parameters(θ) given the data (X):

$l(\theta|X)$ = likelihood equation for appropriate distribution

General formulation of likelihoods – not in exam

$$l(\theta|y) = \frac{y\theta - b(\theta)}{a(\phi)} + c(y, \phi)$$

θ is the expected value (e.g. the mean)

y is the data

$l(\theta|y)$ is likelihood of expected value given the data

ϕ is the variance (dispersion)

a , b , and c are functions – will depend on the distribution used

Fitting GLMs in R

100m times data

Previously fit using `lm()` now try with `glm()`

Data are here:

<https://www.math.ntnu.no/emner/ST2304/2019v/Week5/Times.csv>

100m times data

Fit in R using glm()



```
glm(Y ~ X, data, family = gaussian(link=identity))
```

Fit in R using glm()



```
glm(Y ~ X, data, family = gaussian(link=identity))
```

Exactly like lm()

Systematic part

Fit in R using glm()



```
glm(Y ~ X, data, family = gaussian(link=identity))
```

defines the
distribution you are
using for the **random**
part of the glm

today we use
gaussian, aka Normal

Fit in R using glm()



```
glm(Y ~ X, data, family = gaussian(link=identity))
```

defines the **link function** to relate the **systematic** part to the **random** part

Exercise 4: Fit the GLM and interpret

- Take 100m data you used in earlier weeks
- Use code on slides before to fit a `glm()` and an `lm()` for `WomenTimes`
- Stick with gaussian family and identity link
- Compare the results
- Use `coef()` and `confint.lm()` or `summary()`

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General formula for the likelihood that works for all GLMs but exact functions depend on distribution of data

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Fitting in R

Use `glm()`, very similar to `lm()` but with extra arguments for link random part and link function