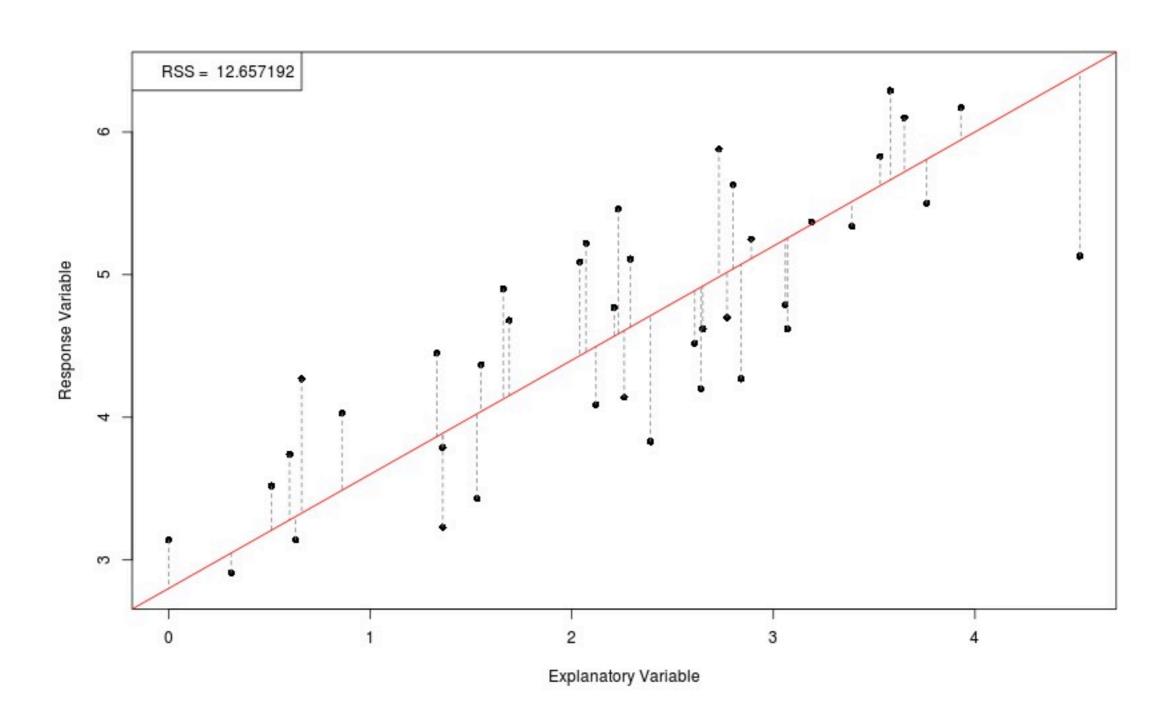
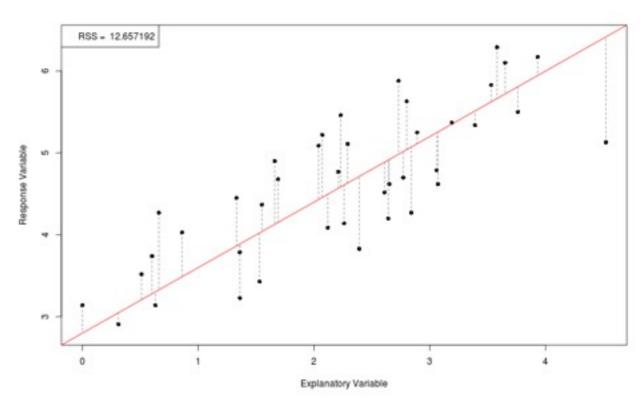
Generalised Linear Models (GLMs)

Owen Jones jones@biology.sdu.dk



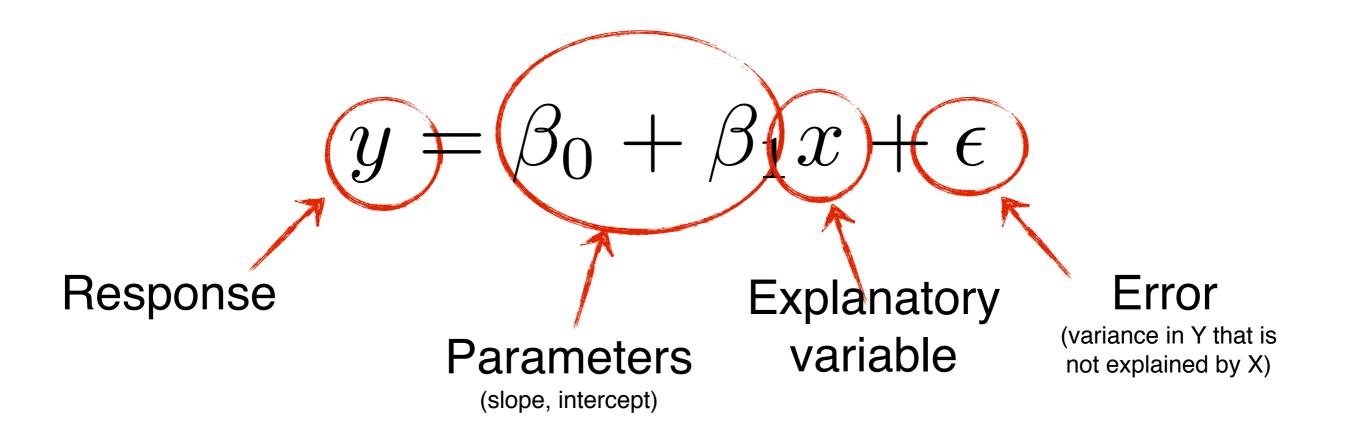
$$y = a + bx$$

intercept slope

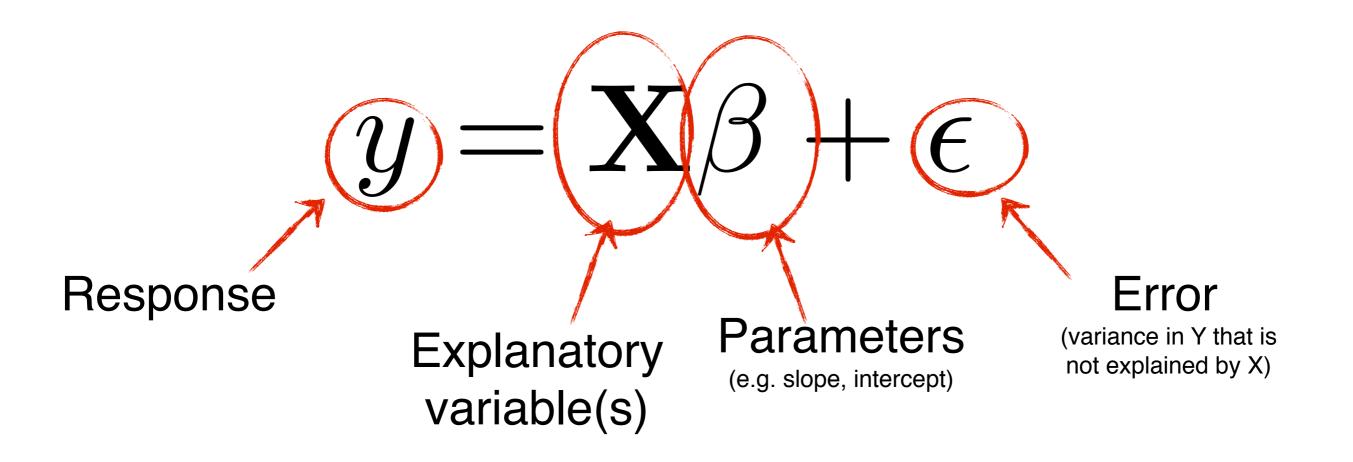


$$y = \alpha + \beta x + \epsilon$$
$$y = \beta_0 + \beta_1 x + \epsilon$$
$$y = \mathbf{X}\beta + \epsilon$$

Textbooks differ in how this is represented: these are identical.

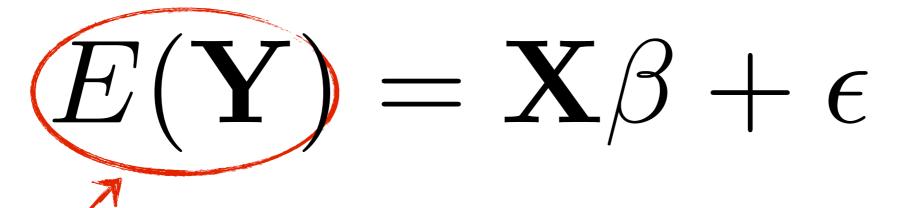


Note: (1) error does not change with explanatory variable (2) change in EV results in linear change in y



Note: (1) error does not change with explanatory variable (2) change in EV results in linear change in y

Remember - we are modeling the *expected* response:

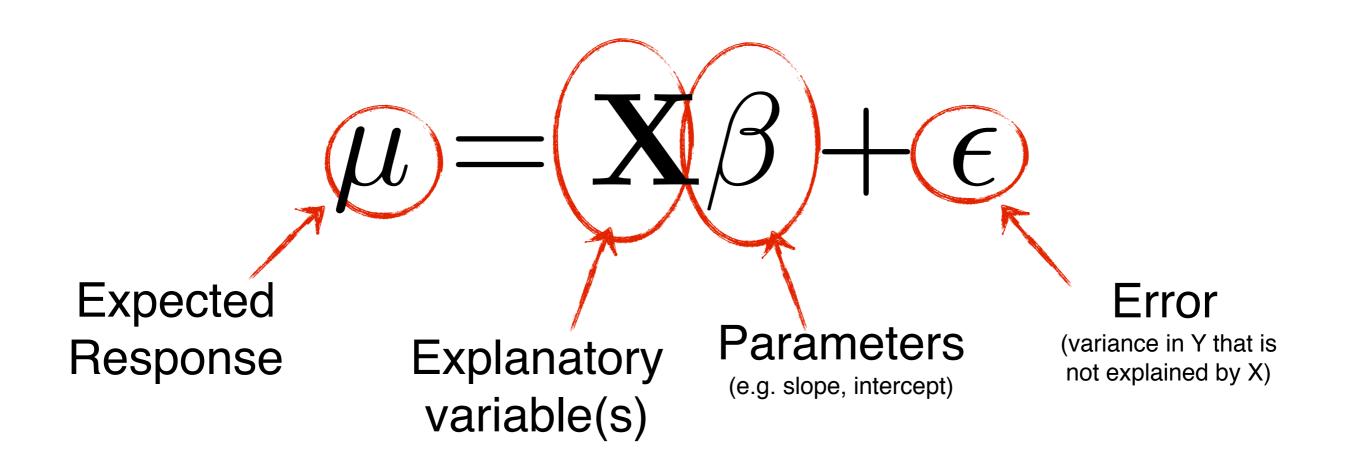


Expected response

and since,
$$E(Y) \equiv \mu$$

we can rephrase to:

$$\mu = \mathbf{X}\beta + \epsilon$$



Includes:

Ordinary regression
Multiple regression
ANOVA/MANOVA
ANCOVA

Assumptions

- linearity of the relationship between explanatory variable(s) and response.
- independence of the errors.
- constant variance (homoscedasticity) vs. the response variable
- normality of the error distribution.

Generalised linear models

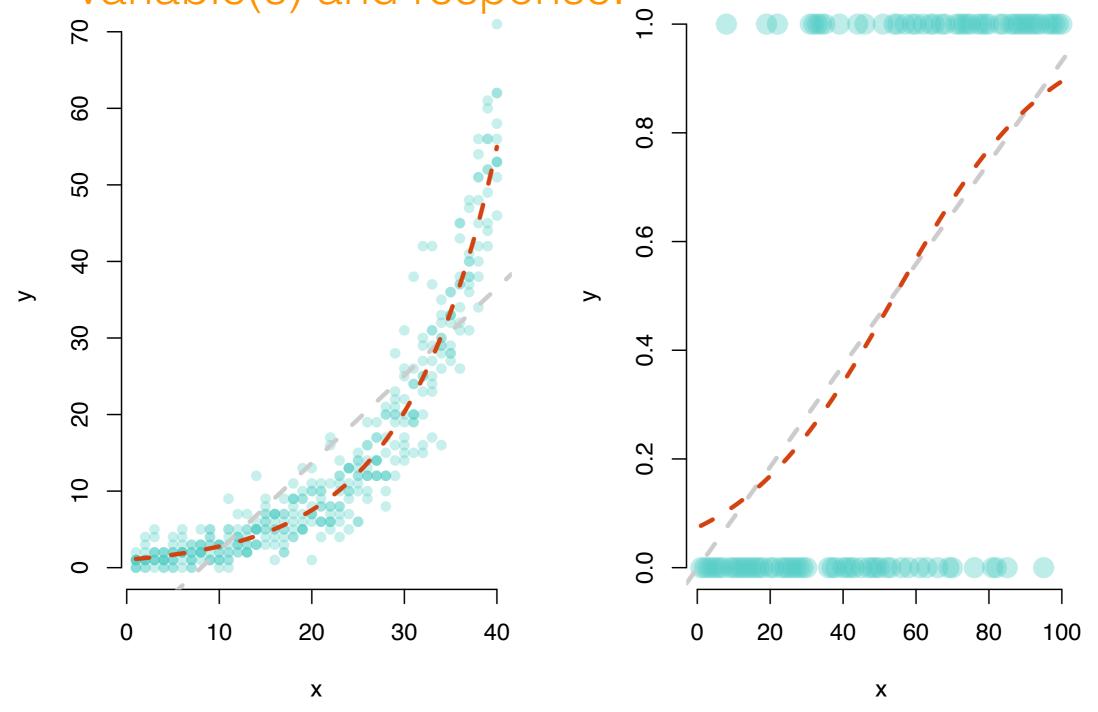
Assumptions

- linearity of the relationship between explanatory variable(s) and response.
- *independence* of the errors.
- constant variance (homoscedasticity) vs.
 the response variable
- normality of the error distribution.

Generalised Linear Models relax these assumptions.

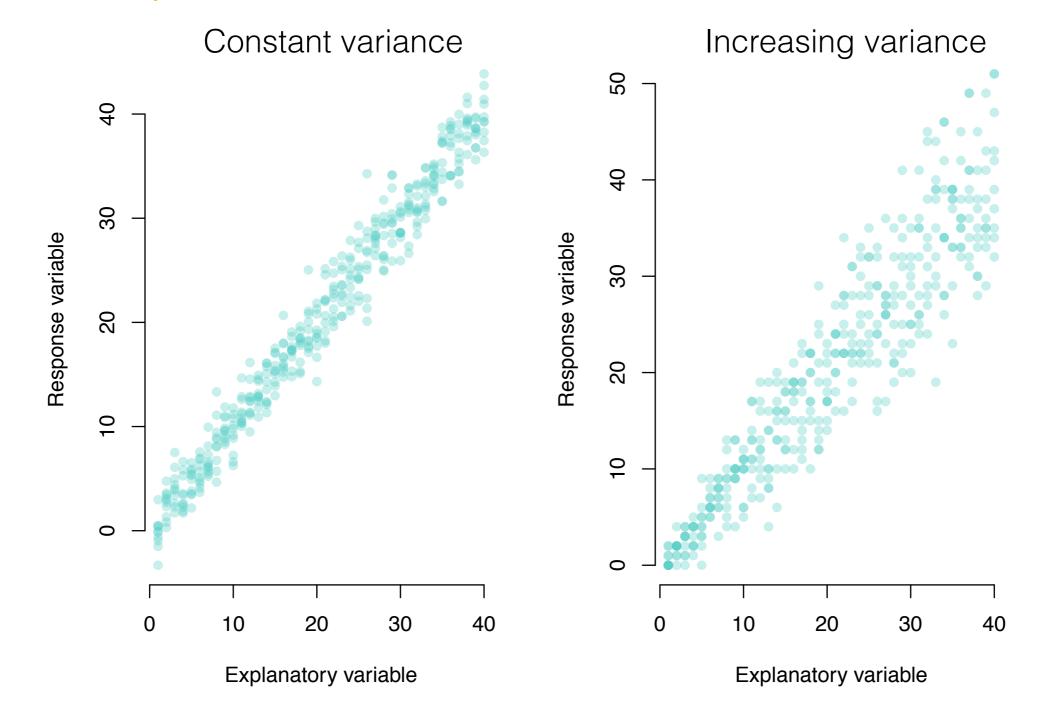
Linearity

linearity of the relationship between explanatory variable(s) and response.



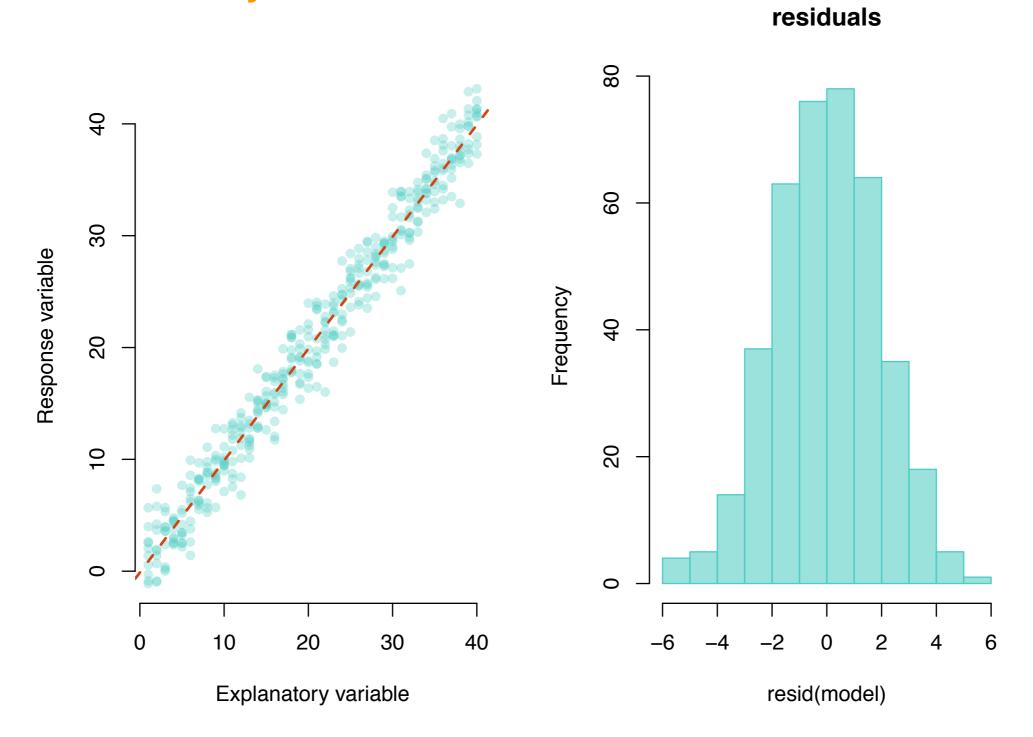
Constant variance

constant variance (homoscedasticity) vs. the response variable



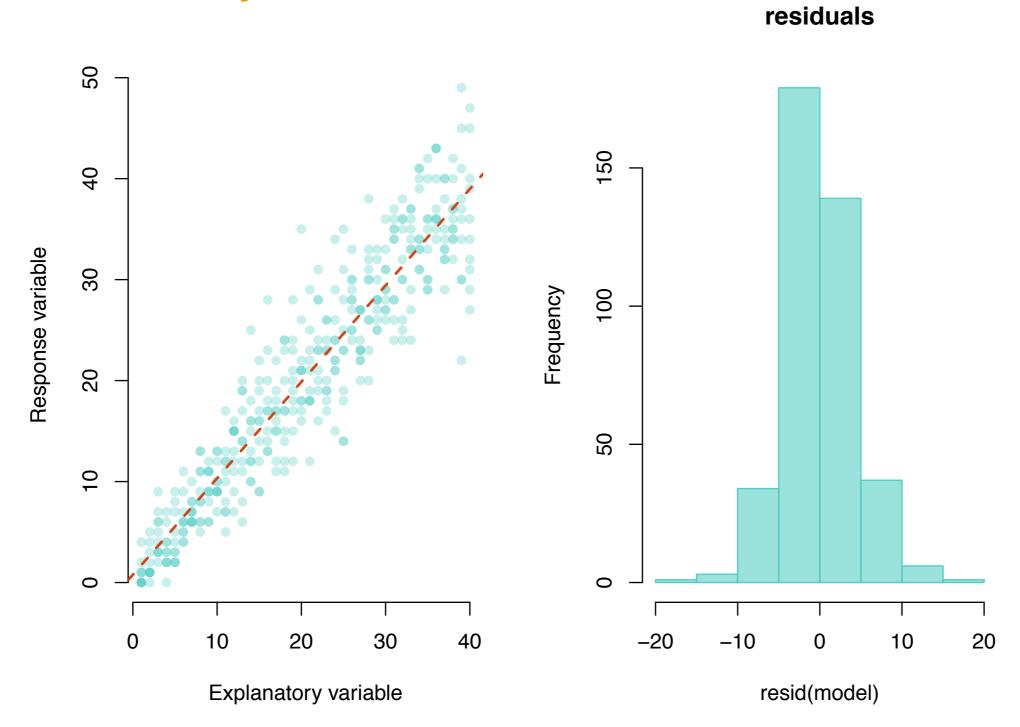
Normality

normality of the error distribution.



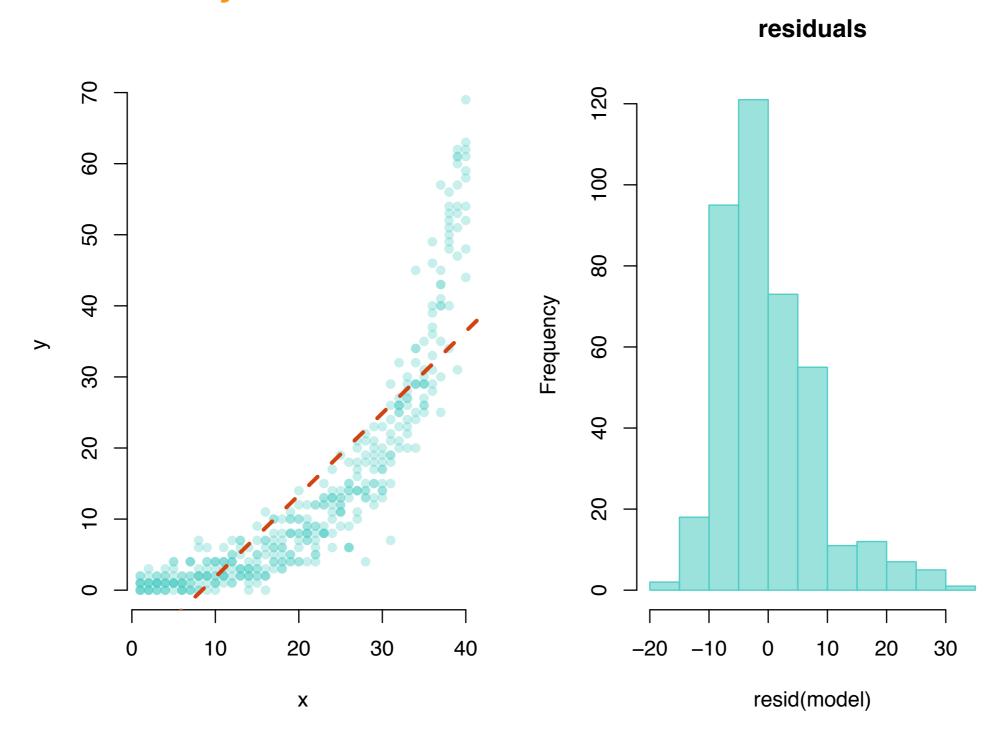
Normality

normality of the error distribution.



Normality

normality of the error distribution.



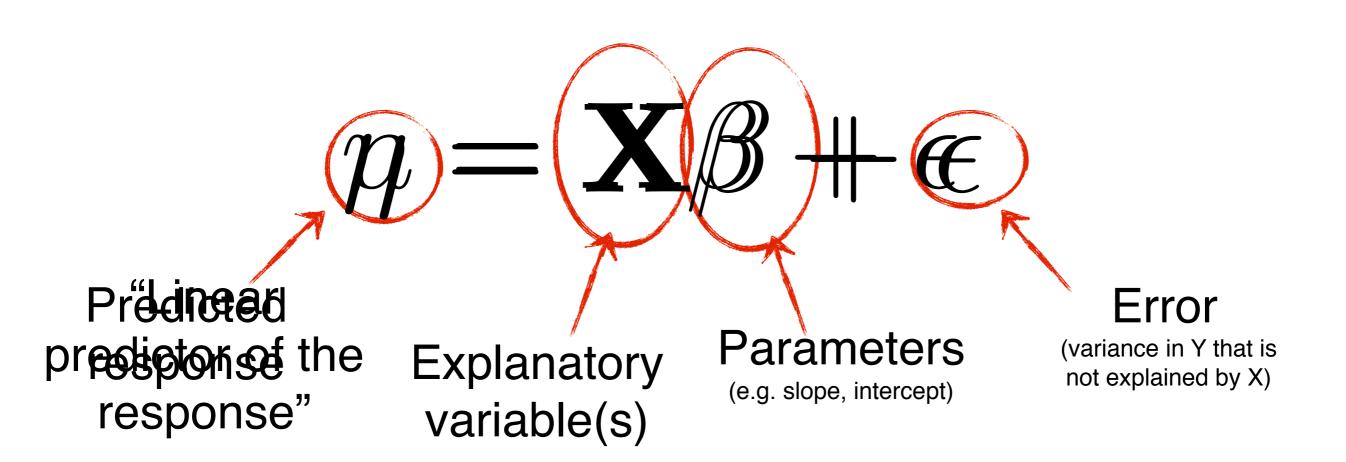
Generalised linear models

GLMs have 3 components

linear predictorlink functionallows to go beyond linear relationship.

variance function - allows non-normal error distributions.

Generalised linear models



Linear predictor & link function

$$\mathfrak{m} = X\beta + \epsilon$$

How do we get back to the predicted response?

$$g(\mu) = \eta = \mathbf{X}\beta + \epsilon$$

If
$$f=g^{-1}$$
 then $f(\eta)=\mu$

Linear predictor & link function

So we have 2 functions, f and g:

link function

$$g(\mu) = \eta$$

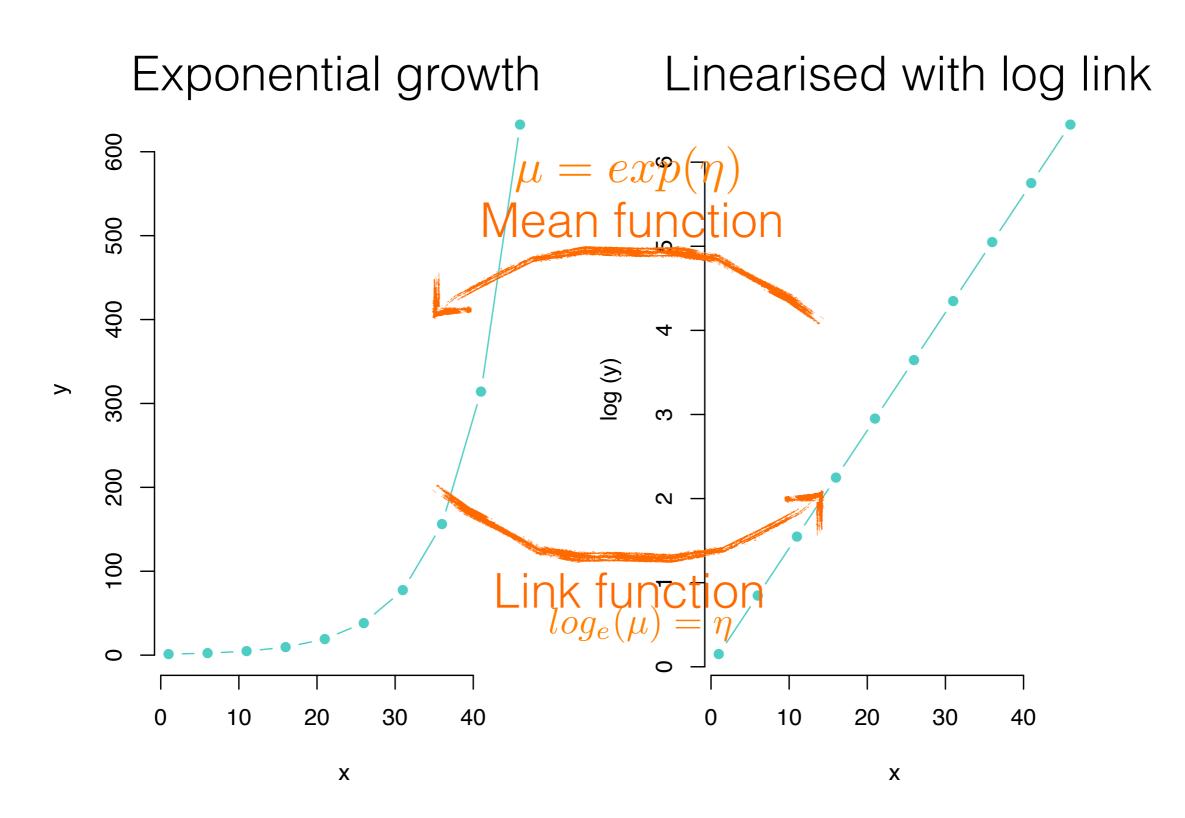
mean function

$$f(\eta) = \mu$$

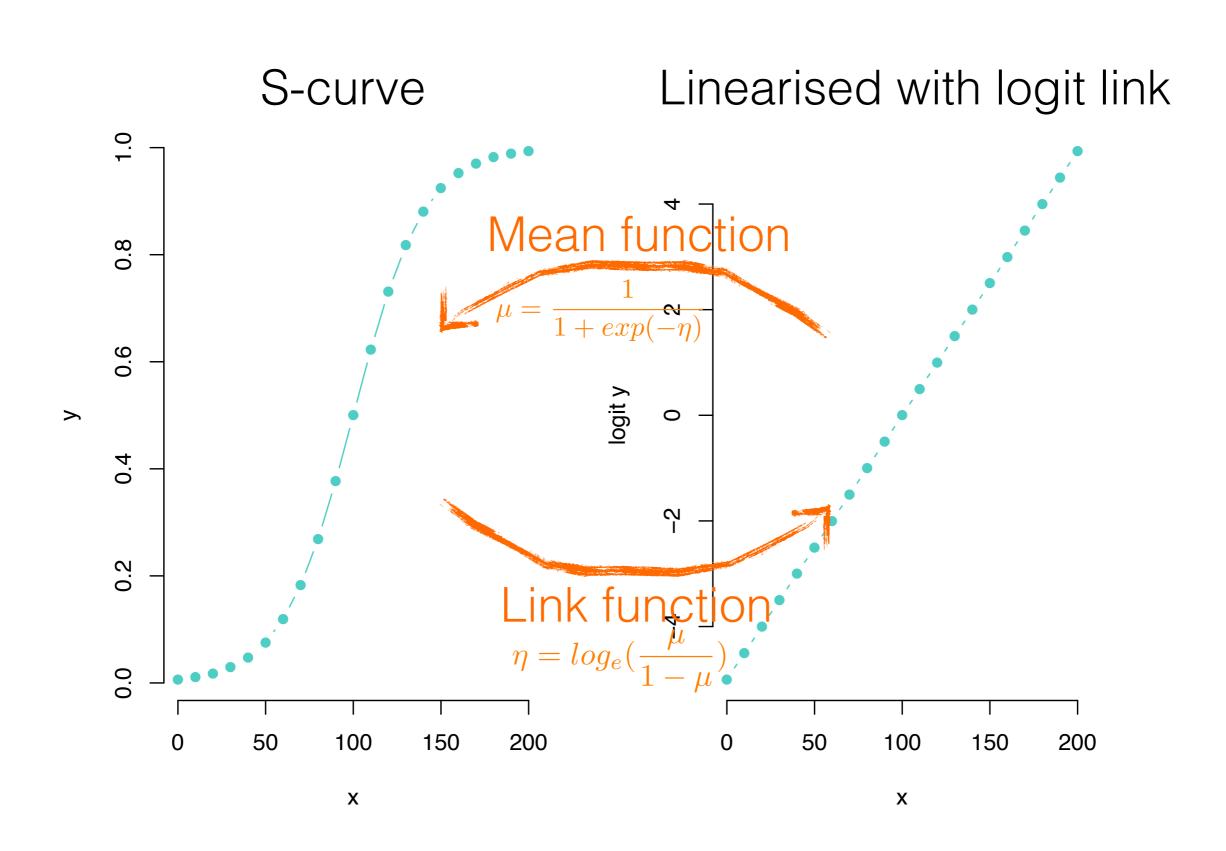
The value of the linear predictor is obtained by transforming the predicted mean using the link function.

The predicted response value is obtained by applying the mean function to the linear predictor.

Log link



Logit link



Variance function

Also called: "Family" or "Error structure"

Describes how variance of response depends on mean - variance is a function of the mean.

- Normal constant variance
- Poisson variance proportional to mean
- Binomial variance proportional to μ(1-μ)
- Gamma variances increases faster than linearly

Choices, choices, choices...

Must choose (1) link (2) error structure:

- Best choice depends on the data.
- Ask yourself:
 - is my response discrete/continuous?
 - what are the bounds?
 - (e.g. can I go less than 0, more than 1)
- To make things easier, there are standard links associated with each error structure.

Choices, choices, choices...

Discrete response (e.g. counts):

- poisson [0,∞], log link.
- binomial [0,N], logit link.

Continuous response (e.g. measurements):

- normal $[-\infty,\infty]$, identity link.
- gamma [0,∞], inverse link.

Standard link and error

Error distribution	Bounds	Link name	Link function	Mean function
Normal	$[-\infty,\infty]$	identity	μ	$\mathbf{X}eta$
Exponential	[0,∞]	inverse	$-\mu^{-1}$	$-(\mathbf{X}\beta)^{-1}$
Gamma	[0,∞]	inverse	$-\mu^{-1}$	$-(\mathbf{X}\beta)^{-1}$
Poisson	[0,∞]	log	$log_e(\mu)$	$exp(\mathbf{X}\beta)$
Binomial	[0,N]	logit	$log_e(\frac{\mu}{1-\mu})$	$\frac{1}{1 + exp(-\mathbf{X}\beta)}$
Bernoulli	[0,1]	logit	$log_e(\frac{\mu}{1-\mu})$	$\frac{1}{1 + exp(-\mathbf{X}\beta)}$

Implementation

Ordinary linear model:

Generalised linear model:

```
model1 = glm(y~x, data=mydata, family = poisson(link = "log"))
```

Handout

The handout runs through:

- Examining data to decide on the error family and link.
- 2. Constructing GLM models for Poisson and Binomial data.
- 3. Interpreting outputs.
- 4. Plotting the model.