(Almost) every technique is a linear model

BIOL2022 - Biology Experimental Design and Analysis (BEDA)

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The general linear model (GLM)

A general (mathematical) framework for expressing relationships between variables.

The basic idea

We are trying to fit a line (or a curve) to the data that best explains the relationship between the variables, essentially a simple equation many of you will be familiar with:

$$y = c + mx$$

where c is the intercept and m is the slope of the line.

Compare this to the general linear model:

$$y = \beta_0 + \beta_1 x + \epsilon$$

$$y = eta_0 + eta_1 x + \epsilon$$

or simply (for your benefit):

$$y \sim x$$

The equation to rule them all, where

- y is the response variable, or the variable we are trying to predict or explain.
- ullet is a predictor variable that can be *anything* continuous, categorical, binary, ordinal, etc.
- ullet eta_i , where i is an integer, are the coefficients/estimates that quantify the relationship between y and x.
- ϵ is the error term, which captures the variability in y that is not explained by x.

In general, common most statistical tests can be expressed as a GLM like above.

$$y=eta_0+eta_1x+\epsilon$$

Why does it matter?

- Under the GLM, the statistical test is identified based on the variables supplied to the model. This follows a model-centric approach:
- If the predictor variable (x) is **binary**, we are performing a GLM *equivalent* to a **t-test** and the slope is the difference in conditional means.
- If the predictor variable (x) is **categorical**, we are performing a GLM *equivalent* to **ANOVA**, and the slope represents the difference in means between the categories.
- If the predictor variable (x) is **continuous**, we are performing a GLM *equivalent* to **regression**. The slope becomes the difference in expected values for a pair of points that differ in x by one unit.
 - \longrightarrow For one unit change in x, the expected change in y is β_1 .
- The GLM takes care of the "statistical test" for you and you don't need to know what the equivalent test is (although it helps).

1. Model the relationship between the response variable and the predictor/explanatory variable(s) as a simple function.

$$response \sim explanation$$

Also read as (recall the previous lecture):

- response is explained by the explanation
- response is a function of the explanation
- response is influenced by the explanation
- ...

1. Model the relationship between the response variable and the predictor/explanatory variable(s) as a simple function.

$$response \sim explanation$$

2. Define the **structure** of the variables in the model, which can be continuous, categorical, binary, etc. – something we can decide if we are the ones designing the study.

$$(continuous) \sim (continuous) \ (continuous) \sim (categorical) \ (binary) \sim (continuous)$$

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$$(continuous) \sim (categorical)$$

3. Fit the model to the data based on the data types – and start work out if the model is a good fit.

$$response = \beta_0 + \beta_1 \times explanation + \epsilon$$

In most statistical software (including R and Jamovi) you only need to specify the model formula (the first part) and the software will take care of the rest to produce the statistical output (the third part).

Your job is to determine if the model is a good fit, and interpret the results.

A good fit?

Checking assumptions – many ways

In previous studies, you may have checked assumptions in different ways depending on the statistical test:

- For a *t*-test, you may have checked for normality by plotting the data and looking for a bell-shaped curve. Equal variances you may have used the F-test or Levene's test.
- For ANOVA, you may have checked for homogeneity of variance using Levene's test or Bartlett's test.
- For regression, you looked at the residuals to check for homoscedasticity and normality.

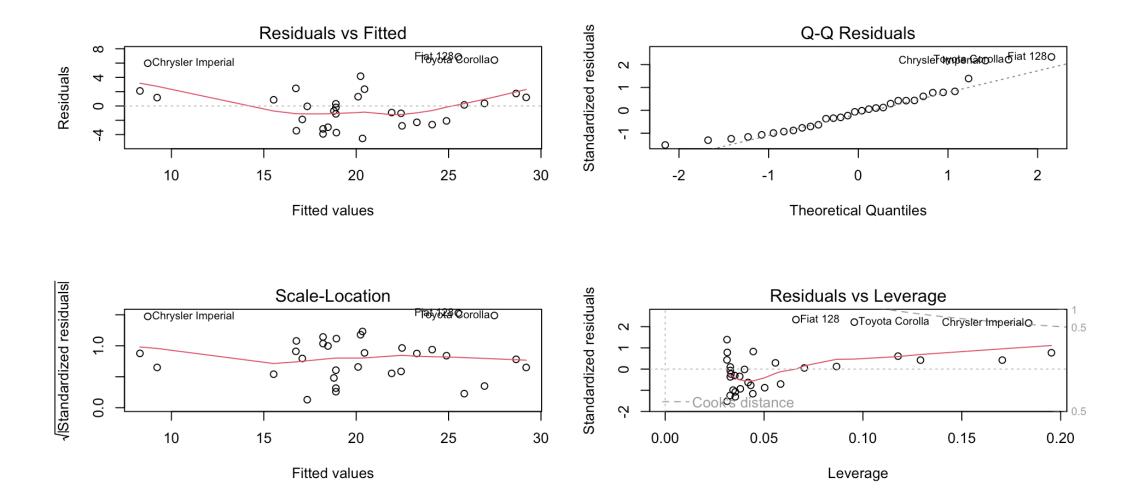
Checking assumptions – the GLM way

A GLM provides a framework for checking model assumptions using residuals to assess any of:

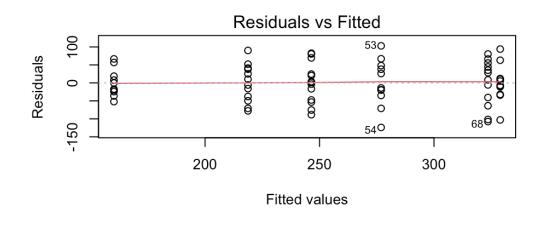
- Homoscedasticity and Linearity: Are the residuals spread evenly across the range of the predictor variable?
- Normality: Are the residuals normally distributed?
- Outliers: Are there any outliers that are affecting the model?
- Influential points: Are there any points that are affecting the model?
- Collinearity: Are the predictor variables correlated with each other?

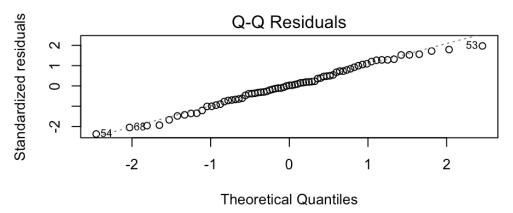
We will cover these in more detail in the next lecture...

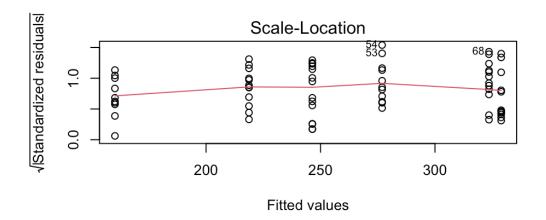
Preview: continuous predictor

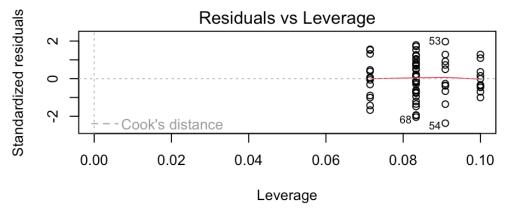


Preview: categorical predictor











Expanding the GLM: More than one variable

So far, we've looked at models with a single explanatory variable. But often, we want to explain the response variable using **more than one** explanatory variable.

For example, perhaps we want to explain the metabolic rate of an organism using both its body mass **and** temperature:

metabolic rate \sim body mass + temperature

What does the plus sign mean?

The plus sign (+) means we are including **both** body mass and temperature as separate, additive predictors of metabolic rate. This is called an **additive model**.

Adding interactions

metabolic rate \sim body mass + temperature

Sometimes, the effect of one variable depends on the value of another. For example, maybe the effect of temperature on metabolic rate changes depending on body mass.

To model this, we include an **interaction**:

metabolic rate \sim body mass \times temperature

This expands to:

metabolic rate \sim body mass + temperature + body mass : temperature

- The: term represents the interaction between body mass and temperature.
- The *shorthand (body mass * temperature) automatically includes both main effects and their interaction.

i Note

There are other types of relationships and model structures, which we will cover as we go along.

Quick exercises

Model the following relationships.

• The number of birds in a forest is influenced by the number of trees and the amount of rainfall.

number of birds \sim number of trees + amount of rainfall

• The number of birds in a forest is influenced by the number of trees and the amount of rainfall, and there is an interaction between the number of trees and the amount of rainfall.

number of birds \sim number of trees + amount of rainfall + number of trees : amount of rainfall

or

number of birds \sim number of trees \times amount of rainfall

Quick exercises

Model the following relationships.

• Whether a turtle is born male or female is influenced by the temperature of the nest.

sex of turtle
$$\sim$$
 temperature of nest

• The height of a seeedling is influenced by temperature and the amount of sunlight it receives.

height of seedling
$$\sim$$
 temperature $+$ amount of sunlight

• air quality is influenced by rainfall, temperature, the number of trees and cloud cover, and there is an interaction between rainfall and temperature.

 $air\ quality \sim rainfall + temperature + number\ of\ trees + cloud\ cover + rainfall:\ temperature$

So how do I use it as a newbie?

You're probably **NOT experienced enough to use it to its full potential** – and that's okay.

- Use the GLM as a **general framework to model relationships** between variables.
- What you *can* do is:
 - 1. Define (model) the relationship between the response and explanatory variables.
 - 2. Identify the **statistical test** the GLM is *related* to.
 - 3. Run the statistical test, but use GLM **as a guide** to diagnose (e.g. assumptions) and interpret the results.

(If there's time) Examples in Jamovi and R

Thanks!

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