

# Random effects and mixed modeling:

General linear models +  
fixed and random effects =  
General linear mixed models

# Outline

- Fixed versus random effects
  - Example with an ANOVA
  - Reasons to use one or the other
- Mixed models
- Lab
  - Going from fixed to random effects

# General Linear Models

## **Linear Regression:**

Continuous response, one continuous explanatory variable

## **T-test:**

Continuous response, one discrete explanatory variable with only two categories

## **One-way ANOVA (Analysis of Variance):**

Continuous response, one discrete explanatory variable with more than two categories

## **Two-way ANOVA:**

Continuous response, two discrete explanatory variable

## **ANCOVA (Analysis of Covariance):**

Continuous response, one discrete explanatory variable and one continuous explanatory variable

# General Linear Models

So far, we assumed that each parameter in our model is estimated separately and independently:

$$y_i = \alpha + \beta x1_i + \delta x2_i + \cdots + \varepsilon_i$$

$$\varepsilon_i \sim \text{Norm}(0, \sigma^2)$$

We refer to each of the parameters as “fixed effects”.

# Fixed and random effects

- ***Fixed effects:*** factors whose levels are experimentally determined or whose interest lies in the specific effects of each level, such as effects of covariates, differences among treatments and interactions.
- ***Random effects:*** factors whose levels are sampled from a larger population, or whose interest lies in the variation among them rather than the specific effects of each level.
- The precise definitions of ‘fixed’ and ‘random’ are controversial; the status of particular variables depends on experimental design and context

# Fixed and random effects

## One-way (factor) ANOVA – fixed effects

- Assume that we measured wing length in five different populations of little owls (*Athene noctua*)
- Is there a difference in wing length among the populations?

# Fixed and random effects

One-way (factor) ANOVA – fixed effects

With a means parameterization, we can write the model:

$$y_i = \alpha_{j(i)} + \varepsilon_i$$
$$\varepsilon_i \sim \text{Norm}(0, \sigma^2)$$

- $y_i$  = observed wing length of owl  $i$  in population  $j$
- $\alpha_{j(i)}$  = expected wing length of an owl in population  $j$
- $\varepsilon_i$  = the random wing deviation of owl  $i$  from its population mean

# Fixed and random effects

- In this model, the levels (groups) of the factor were assumed to be fixed by design; we have a special interest in the particular five little owl populations that we want to compare and have no interest in generalizing to other populations
- But what if aren't actually interested in the differences among the sites? What if instead we considered these five groups a random sample of some larger population of little owl and wanted to generalize our results?



# Fixed and random effects

One-way (factor) ANOVA – random effects

In such a case, we can re-write the model:

$$y_i = \alpha_{j(i)} + \varepsilon_i$$

$$\varepsilon_i \sim \text{Norm}(0, \sigma^2)$$

$$\alpha_{j(i)} \sim \text{Normal}(\mu, \tau^2)$$

- $y_i$ ,  $\alpha_{j(i)}$ , and  $\varepsilon_i$  have the same interpretation but now each of the  $\alpha_{j(i)}$  parameters are no longer assumed to be independent
- Instead, they come from a second normal distribution with a mean of  $\mu$  and a variance of  $\tau^2$

# Fixed and random effects

One-way (factor) ANOVA – random effects

In such a case, we can re-write the model:

$$y_i = \alpha_{j(i)} + \varepsilon_i$$

$$\varepsilon_i \sim \text{Norm}(0, \sigma^2)$$

$$\alpha_{j(i)} \sim \text{Normal}(\mu, \tau^2)$$

What is the interpretation of  $\mu$  and  $\tau^2$  ?

$\mu$  = mean winglength across all the five little owl populations

$\tau^2$  = variance in winglength across populations

# Fixed and random effects

Which to choose?

## **Fixed Effects**

- You have a particular interest in the studied factor levels
- You have included all conceivable levels in a study
- No interest in the variance among levels
- No interest in generalizing to factor levels that you did not study

# Fixed and random effects

Which to choose?

## **Random Effects**

- You don't have a particular interest in the studied factor levels and/or you could not have sampled all levels
- Interested in the variation among levels (but may still want to understand the effects for the observed levels)
- You want to generalize to a larger population (e.g., levels are more like samples)

# Fixed and random effects

## Three reasons to go from fixed to random effects:

- Extrapolation to a wider population of inference
- Improved accounting for system uncertainty

Randomness in  $\varepsilon$  and both  $\tau$ . Acknowledges that repeating our study would result in different parameter estimates

- Efficiency of estimation -> shrinkage

Parameters are no longer independent and will be pulled to the mean, “borrowing strength”

# Mixed models

- In many situations, it is useful to make use of both fixed and random effects
- **Mixed models** contain both fixed and random effects
- Introduced by Fisher when studying correlations in traits among relatives
- Particularly useful for:
  - Repeated measures
  - Missing data

# Mixed models in R

- Can be hard to fit mixed and random effects models using canned functions in most software packages
  - algorithms do not always converge, especially when the number of groups is small

# Mixed models in R

- Can be hard to fit mixed and random effects models using canned functions in most software packages
- In R, can use:
  - **aov** function with “Error” term -- if predictors are all categorical and design is balanced
  - **nlme** package
  - **lme4** package – allows for unbalanced data sets, random effects on parameters (e.g., slope in an ANCOVA), and nonlinear models (not discussed here)



# Mixed models in R

- What to do when those don't work? Suggestions from Bolker:
  1. Fit fixed-effects models instead. May lose power -> conservative results
  2. For nested design (e.g., subsamples in blocks) -> collapse groups data by computing means and do a single level analysis
  3. Fit model ignoring blocks and examine the variation in residuals between blocks -> must show between-group variation in residuals is both statistically and biologically irrelevant
  4. Switch to a Bayesian analysis -> computing/software, while more difficult to learn, is much more flexible

# Fixed and random effects

## *Lab: ANOVA*

- Review the ANOVA lab from last week
- Modify the data simulation with an assumption that means of populations are random drawn from a common population-level grand mean.