Introduction to mixed effects models in R

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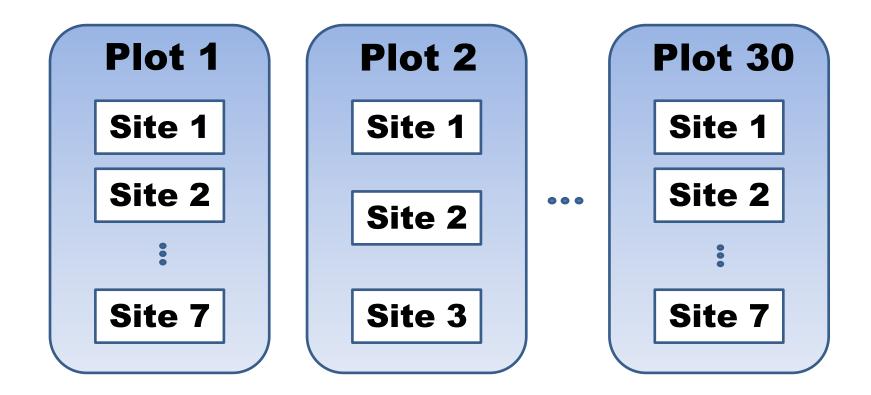
Structural indicators of spider communities across the forest plantation cycle

Anne G. Oxbrough ^a A ⊠, Tom Gittings ^a, John O'Halloran ^a, Paul S. Giller ^a, George F. Smith ^b

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Study sampling design



Linear regression

Shannon_i=
$$\beta_1+\beta_2HerbLayer_i+\varepsilon_i$$

 $\varepsilon_i\sim N(0,\sigma^2)$

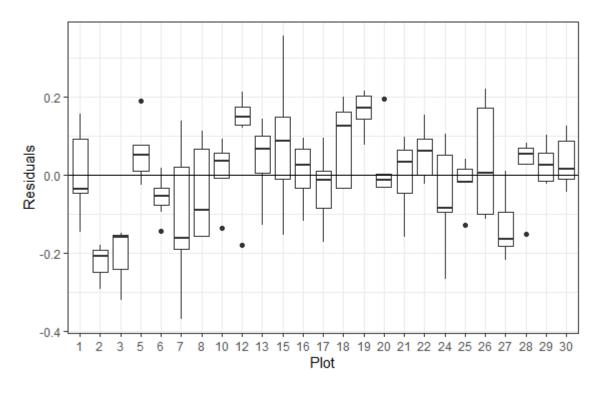
$$Shannon_i \sim N(\mu_i, \sigma^2)$$

$$\mu_i = \beta_1 + \beta_2 Herb Layer_i$$

Model validation

Shannon_i=
$$\beta_1+\beta_2+HerbLayer_i+\varepsilon_i$$

 $\varepsilon_i \sim N(0, \sigma^2)$



There is plot effect in the residuals!!

Alternative model

$$Shannon_{i} = \beta_{1} + \beta_{2} Herb Layer_{i} + Plot_{i} + \varepsilon_{i}$$
$$= (\beta_{1} + Plot_{i}) + \beta_{2} Herb Layer_{i} + \varepsilon_{i}$$

$$\varepsilon_i \sim N(0, \sigma^2)$$

$$Shannon_i \sim N(\mu_i, \sigma^2)$$

$$\mu_i = \beta_1 + \beta_2 HerbLayer_i + Plot_i$$

Complete pooling VS no-pooling

Complete pooling

There is a single intercept for all the observations.

$$Shannon_i = \beta_1 + \beta_2 Herb Layer_i + \varepsilon_i$$



No pooling

There is a single intercept for each site.

$$Shannon_i = (\beta_1 + Plot_i) + \beta_2 Herb Layer_i + \varepsilon_i$$

Complete pooling VS no-pooling

Complete pooling

There is a single intercept for all the observations.

$$Shannon_i = \beta_1 + \beta_2 Herb Layer_i + \varepsilon_i$$

Partial pooling

The intercepts are correlated

No pooling

There is a single intercept for each site.

$$Shannon_i = (\beta_1 + Plot_i) + \beta_2 Herb Layer_i + \varepsilon_i$$

Linear mixed model

Shannon_{ij}=
$$\beta_1+\beta_2HerbLayer_{ij}+a_i+\varepsilon_{ij}$$

 $a_i \sim N(0, \sigma^2_{Plot})$
 $\varepsilon_{ij} \sim N(0, \sigma^2)$

Linear mixed model

$$Shannon_{ij} = \beta_1 + \beta_2 Herb Layer_{ij} + a_i + \varepsilon_{ij}$$

$$a_i \sim N(0, \sigma^2_{Plot})$$
 $\varepsilon_{ij} \sim N(0, \sigma^2)$

$$arepsilon_{i\, m{i}} {\sim} N(0, \sigma^2)$$

Fixed Part: $\beta_1 + \beta_2 Herb Layer_{ij}$

Random Part: $a_i + \varepsilon_{ij}$

A "mixed model" has both fixed and random parts

Random versus Fixed Effects

Random effects are estimated with partial pooling, while fixed effects are not.

(Andrew Gelman and Jennifer Hill)

Varies by group

Variation estimated with **probability model**

Random intercept model

$$Shannon_{ij} = \beta_1 + \beta_2 Herb Layer_{ij} + a_i + \varepsilon_{ij}$$

$$= (\beta_1 + a_i) + \beta_2 Herb Layer_{ij} + \varepsilon_{ij}$$

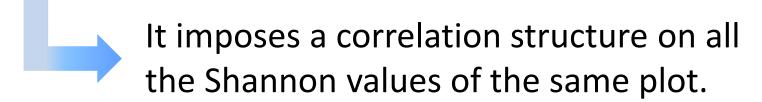
$$\varepsilon_{ij} \sim N(0, \sigma^2)$$

$$a_i \sim N(0, \sigma^2_{Plot})$$

Random intercept model

$$Shannon_{ij} = \beta_1 + \beta_2 Herb Layer_{ij} + a_i + \varepsilon_{ij}$$
$$= (\beta_1 + a_i) + \beta_2 Herb Layer_{ij} + \varepsilon_{ij}$$
$$\varepsilon_{ij} \sim N(0, \sigma^2)$$

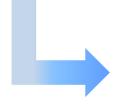
$$a_i \sim N(0, \sigma^2_{Plot})$$



Random intercept model

$$Shannon_{ij} = \beta_1 + \beta_2 Herb Layer_{ij} + a_i + \varepsilon_{ij}$$
$$= (\beta_1 + a_i) + \beta_2 Herb Layer_{ij} + \varepsilon_{ij}$$
$$\varepsilon_{ij} \sim N(0, \sigma^2)$$

$$a_i \sim N(0, \sigma^2_{Plot})$$



$$cor(Shannon_{ij}, Shannon_{ik}) = \varphi = \frac{\sigma^2_{Plot}}{\sigma^2_{Plot} + \sigma^2}$$

Protocol to fit a mixed model

- 1. Based on prior knowledge of the dependency structure in the data, select the random structure (i.e., the random effects) a priori.
- 2. Fit the model and investigate which covariates in the fixed part are important.
- 3. When optimal model has been found, present the numerical output and provide a graphic representation of the model fit.

Zuur, Alain.F, Joseph M. Hilbe, and Elena N Ieno (2013). A Beginner's Guide to GLM and GLMM with R: a frequentist and Bayesian perspective for ecologists, Highland Statistics

Our final random intercept model

$$Shannon_{ij}=0.994+0.034HerbLayer_{ij}$$

$$-0.031LitterCov_{ij}+a_i+\varepsilon_{ij}$$

$$\varepsilon_{ij}\sim N(0,0.106^2)$$

$$a_i\sim N(0,0.063^2)$$

Random intercept and slope model

$$\begin{split} Shannon_{ij} &= \beta_1 + \beta_2 Herb Layer_{ij} + a_i + b_i Herb Layer_{ij} + \varepsilon_{ij} \\ &= (\beta_1 + a_i) + (\beta_1 + b_i) Herb Layer_{ij} + \varepsilon_{ij} \\ \varepsilon_{ij} \sim N(0, \sigma^2) \end{split}$$

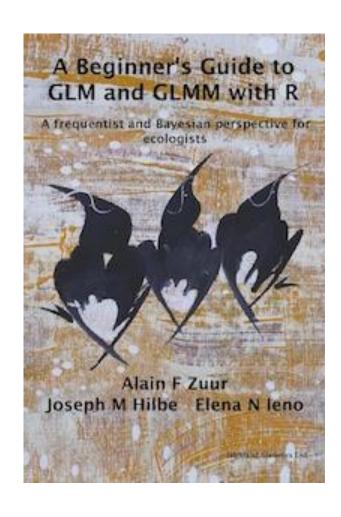
$$a_i \sim N(0, \sigma^2_{1,Plot})$$

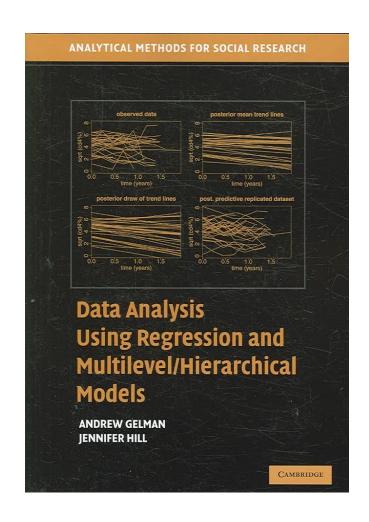
$$b_i \sim N(0, \sigma^2_{2,Plot})$$

Generalized linear mixed models

$$Richness_{ij} \sim Poisson(\mu_{ij})$$
 $E(Richness_{ij}) = \mu_{ij} \quad var(Richness_{ij}) = \mu_{ij}$
 $\mu_{ij} = \beta_1 + \beta_2 HerbLayer_{ij} + a_i$
 $a_i \sim N(0, \sigma^2_{Plot})$

Recommended references





Recommended references



A brief introduction to mixed effects modelling and multi-model inference in ecology

Xavier A. Harrison¹, Lynda Donaldson^{2,3}, Maria Eugenia Correa-Cano², Julian Evans^{4,5}, David N. Fisher^{4,6}, Cecily E.D. Goodwin², Beth S. Robinson^{2,7}, David J. Hodgson⁴ and Richard Inger^{2,4}

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