

GLMM workshop 7 July 2016

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First session 1 PM Room SN2109 Writing the model

Break

Second session 2 PM SN 2018/2025 F-ratios from Expected Mean Squares

Break

Third session 3:30 SN 2067/2071 Executing the analysis

Goal of the first session – Writing Statistical Models

GLM	The General Linear Model	Fixed Effects + Normal Error
GzLM	The Generalized Linear model	Fixed Effects + Non-normal Errors
GLMM	The General Linear Mixed Model	Fixed + Random + Normal
GzLMM	The Generalized Linear Mixed Model	Fixed + Random + Non-normal

Goal of the second session - Writing out the expected mean squares

Forming unambiguous likelihood ratio tests (F , t , χ^2)

Goal of the third session - Executing a GLMM in a statistical package

Interpreting the output

First session

1 PM Room SN2109

Writing the model

Preliminaries

Definitions Nominal, Ordinal, Interval, and Ratio scale variables.

Definitions: GLM General Linear Model

GzLM Generalized Linear Model

GLMM General Linear Mixed Model

GzLMM Generalized Linear Mixed Model

Series of examples to work through.

Distinguish response from explanatory variables

Assign symbols to all variables

Notational conventions Nominal scale variable ALL UPPER CASE

Ratio scale variables Begin with upper case.

β for fixed effect coefficients (slopes and contrasts)

μ for random effect parameters

Write the model, calculate the df, complete the first 2 columns of the ANOVA table

Preview

GLM - Fixed Effects

Single explanatory variable – 3 examples

Write the GLM Fixed Effect model

Write the degrees of freedom below each term in the model - - > Source df table

Two explanatory variables – Crossed - 3 examples

Write the GLM – Fixed * Fixed - - > Source df table

- Factor * Factor
- Factor * Covariate
- Covariate * Covariate

GzLM - Fixed Effects. The first solution to heterogeneous errors - 2 examples

GLM - Random Effects. The second solution to heterogeneous errors.

Definition of Random Effects, Random variables.

Identify explanatory variables as Random or Fixed

β Notation for fixed factors. μ notation for random factors

Single explanatory variable - 1 example

Write the GLMM -- - > Source df table

- Fixed * Random Effects

Two explanatory variables – Nested - 1 example

How to distinguish nested from crossed factors

Write the GLM - - > Source df table

- Random(Random)

Write the GLMM - - > Source df table

- Fixed + Random(Random)

GLMM - Mixed Effects (Fixed and Random)

Two explanatory variables

- Nested Example
- Crossed Example

Three explanatory variables

- Nested Example
- Crossed Example (Latin Square)

GzLMM – Non-normal error and mixed effects (fixed and random factors)

GLM with a single fixed explanatory variable 3 examples.

Write the Fixed Effect GLM, calculate df, fill in the blank columns of the ANOVA table.

1. Pea section growth data, from Box 9.4 in Sokal and Rohlf (1995).

Does length depend on treatment (control versus 4 different sugars with auxin present) ?

10 measurements of length of pea section in each treatment group

Length *Len* Response variable, ratio scale
 Treatment *TRT* Categorical explanatory variable

Write the model $Len = \beta_0 + \beta_{TRT} TRT + \epsilon_{Normal}$
 Calculate df $(10*5) = 1 + (5-1) + 45$

Sketch graph of response vs explanatory

df total = ntot -1 TRT df = number of categories – 1

Fill out first 2 columns of ANOVA table from model

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt3/Ch10_3.pdf

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/PeaSections.csv>

Source	df
<i>TRT</i>	
error	
total	49

2. Example 9.3.1 from Snedecor and Cochran (1989). Quantity of interest is the phosphorus

content of corn (*Pcorn* in ppm), in relation to the phosphorus levels in samples of soils with experimentally fixed levels of phosphorus (*Psoil* in ppm). Does the phosphorus content of corn increase when organic soil phosphorus is increased ? *Pcorn* and *Psoil* are both ratio scale variables. 9 measurements of *Pcorn*, matched with 9 of *Psoil*

Model _____

df _____

Sketch graph of response vs explanatory

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt3/Ch9_1.pdf

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/PCorn.csv>

Source	df
error	7
total	

GLM with a single fixed explanatory variable 3rd example.

3. Does inversion heterozygosity (HZYG) change with elevation above sea level (Hsl) in *Drosophila pseudoobscura*. Data are from Dobzhansky (1948) as reported in Brussard (1984).

One measurement of HZYG at each of 7 different elevations.

Source	df
	5

Response variable with symbol _____

Explanatory variable with symbol _____

Model _____

df _____

GLM with a single fixed explanatory variable Review

Definition of fixed effects:

1. TRT is a fixed effect because we are interested in the contrast among the 5 means.
 β_{TRT} is a set of unknown fixed effect contrasts.
2. P_{soil} is a fixed effect because we are interested in rate of increase in P_{corn} with increase in P_{soil} .
 $\beta_{P_{soil}}$ is the unknown rate.
3. Hsl is a fixed effect because we are interested in the whether $Hzyg$ changes with elevation (altitude above sea level)
 β_{Hsl} is the fixed effect rate. $\hat{\beta}_{Hsl}$ is an estimate of β_{Hsl}

GLM with two fixed explanatory variables 3 examples Factor * Factor
 Factor * Covariate
 Covariate * Covariate

Format for writing a model with two explanatory variables

$$Response = \beta_o + \beta_{V1}V1 + \beta_{V2}V2 + \beta_{V1 \times V2} V1 \times V2 + \epsilon_{Normal}$$

The interactive term is written as the product of two component variables $\beta_{V1 \times V2} V1 \times V2$

Verbal statement: The effect of V1 on the response variable depends on V2

Write the Fixed factor \times Fixed factor GLM, calculate df, fill out the Source df table

$$df \text{ total} = n_{tot} - 1 \quad df \text{ } V1 \times V2 = df(V1) \times df(V2)$$

4. Does oxygen consumption VO_2 depend on salinity (100% 75% and 50% seawater) in two species of limpet (*Acmea digitalis* and *A. scabra*)? Eight measurements at 3 different salinities in each of two species $n_{tot} = 48$. Data from Sokal and Rohlf (1995).

Response variable with symbol _____

Explanatory variable	Symbol	Categorical or Ratio scale
_____	_____	_____
_____	_____	_____

Model _____

df _____

Source	df
	42

Interpret the interactive effect (state this in words)

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt4/Ch13_1.pdf

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/Limpets.csv>

GLM with two fixed explanatory variables

Factor * Factor
2nd example -> Factor * Covariate (aka ANCOVA)
Covariate * Covariate

5. Does inversion heterozygosity (H_{zyg}) change with elevation above sea level (H_{sl}), in 2 species of *Drosophila* (SP = *D. persimilis* or *D. pseudoobscura*). Data are from Dobzhansky (1948) as reported in Brussard (1984). One measurement in each species at 7 different elevations.

Source	df
	10

Model _____

df _____

Complete the Source df table.
Interpret the interactive effect (state it in words)

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt4/Ch14_1.pdf

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/Brussard.csv>

GLM with two fixed explanatory variables

Factor * Factor

Factor * Covariate

3rd example -> Covariate * Covariate (aka multiple regression)

6. Data from Snedecor and Cochran 1980 Table 17.2.1

Does plant available phosphorus content of corn (ppm) from 17 Iowa soils at 20 deg C depend on inorganic and organic phosphorus in the soil?

.

Model _____

df _____

Source	df
	13

Complete the Source df table.

Interpret the interactive effect (state it in words)

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt4/Ch12_1.pdf

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/PAvailable.csv>

GzLM- The first solution to heterogeneous errors - 2 examples (Poisson and Binomial)

The GLM applies to a ratio scale response variable Y with normal error ε_{Normal}

Count data usually violate the assumption of homogeneous residuals.

Ratio scale counts (counts in defined units, ranging from zero upward)

Use Poisson or Negative Binomial error model $\varepsilon_{Poisson}$ or $\varepsilon_{NegBinomial}$

So we use the Generalized Linear Model, which allows us to use a better error model.

The GzLM (which includes the GLM as a special case) has three components

1. The structural model consisting of linear predictors.

For the GLM, the linear predictor is the sum of fixed factors and covariates.

The linear predictor ANCOVA example (Brussard) was $\eta = \beta_o + \beta_{SP}SP + \beta_{Hsl}Hsl + \beta_{SP \cdot Hsl}SP \cdot Hsl$

2. A linkfunction, that links the linear predictor to the response variable.

3. The error

Model equation form: $Y = \beta_o + \beta_x X + \varepsilon_{Normal}$

Probability distribution form: $Y \sim Normal(\beta_o + \beta_x X, \sigma^2)$

This is read as : Y is normally distributed, given the parameters β_o , β_x , and σ^2 (the fixed variance)

The distributional assumption, given the parameters, can only be checked after estimating the parameters

Count data usually violate this assumption.-- > heterogeneous residuals

So we use a better error model (Generalized Linear Model)

Ratio scale counts (counts in defined units, ranging from zero upward)

Use Poisson or Negative Binomial error model $\varepsilon_{Poisson}$ or $\varepsilon_{NegBinomial}$

$$Count = e^\eta + \varepsilon_{Poisson}$$

$$\eta = \beta_o + \beta_{V1}V1 + \dots$$

8. Death by horsekick. The classic example of Poisson data is the number of deaths by horse kick for each of 16 corps in the Prussian army, from 1875 to 1894. Bortkiewicz (1898 *The Law of Small Numbers*) showed that the horsekick data fit a Poisson distribution.

Corps	Deaths
Guard	16
First	16
Second	12
Third	12

Symbol for response variable _____ and for explanatory variable _____

Write the model

$$Odds = e^{\eta} + \varepsilon_{Poisson}$$

(Fit to 1:1:1:1 assumes Poisson error)

$$\eta = \underline{\hspace{10em}}$$

Likelihood Ratio Test: $\Delta G = 1.147$ $df = 3$ $p = 0.7658$ Therefore, cannot reject 1:1:1:1 fit

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt5/Ch17_2.pdf

GzLM with fixed explanatory variables. - 2nd example

The GLM assumes a normal error with fixed (constant variance) = ϵ_{Normal}

Count data usually violate this assumption.-- > heterogeneous residuals

So we use a better error model (Generalized Linear Model)

Nominal scale counts (units scored Y or N) Use binomial error model $\epsilon_{binomial}$

$$\text{Yes/No} = \text{Odds} \quad \text{Odds} = e^{\eta} + \epsilon_{binomial}$$

$$\eta = \beta_o + \beta_{V_1}V_1 + \dots$$

The response variable, Odds, are calculated as $p/(1-p)$, where p is the ratio of success to number of trials.

9. Example – Cancer in cigarette smokers. Data from Cornfield (1951) who established the mathematical basis for using case-control samples to estimate risk in a population.

Odds of tumor for
Heavy smokers _____

Light smokers _____

Odds ratio, heavy relative to light _____

Symbol for response variable _____ and for explanatory variable _____

Write the model

$$\text{Odds} = e^{\eta} + \epsilon_{binomial}$$

(contingency test not correct. it assumes Poisson error instead of binomial)

$$\eta = \underline{\hspace{10em}}$$

The 95% confidence limits are 1.05 to 5.1.

The null hypothesis is OR = 1. (Odds the same for light and heavy smokers)

Do the confidence limits exclude the null? _____

	Lung Tumors		Total
	Present	Absent	
heavy smokers	27	99	126
light smokers	8	72	80

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt5/Ch18_3.pdf

GLM - Random Effects. The second solution to heterogeneous errors.

The GLM assumes a normal error with fixed (constant) variance = ϵ_{Normal}

Grouped data usually violate this assumption.-- > heterogeneous residuals

Examples: Paired data, clustered data, blocked data

Examples: Repeated measures (e.g. 3 samples at one time), longitudinal data (3 samples in sequence)

To capture this heterogeneity, we introduce a random effect variable Z with random coefficients τ (tau).

$$Y = \mu_o + \tau_Z Z + \epsilon_{Normal}$$

$$\mu_o = \text{random intercept}$$

$$\tau_Z Z = \text{random effectd}$$

GLM Single Random Factor

10 The first published ANOVA table was Example 38 in Fisher (1925) *Statistical Methods for Research Workers*.

“In an experiment on the accuracy of counting soil bacteria, a soil sample was divided into four parallel samples and from each of these after dilution seven plates were inoculated. The number of colonies on each plate is shown below in example 12 (Table 41). Do the results from the four samples agree within the limits of random sampling? In other words, is the whole set of 28 values homogeneous, or is there any perceptible intraclass correlation?”

Table 42	Degrees of Freedom	Sum of Squares	Mean Square	F-ratio	R ²	Likelihood Ratio
Between Classes (Soil sample)	3	1446				
Within Classes (Error)	24	94.96				

Assign a symbol to the response variable _____ and explanatory variable _____

Write the model (use μ and τ) _____

Compute both mean squares (= SS/df) and place them in the ANOVA table

Compute the ratio of the two means squares (the F-ratio) and place it in the table

Compute the explained variance $R^2 = \text{Between class SS} / \text{SS}_{total} =$ _____

Do the 4 samples deviate from random sampling? To find out we calculate the likelihood ratio.

$$LR = (1-R^2)^{-n/2} =$$

Likelihood Ratio test: Compare the F-ratio to the 5% p-value of the F-distribution

The 5% probability for the F-distribution (excel code) is: FINV(0.05,3,24) = 3.009

Do the results from the four samples agree within the limits of random sampling? _____

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/FisherEx38.csv>

GLM with two random factors 2 examples

Nested - Random within Random
 Crossed - Random × Random

$$Y = \mu_o + \sum \tau_z Z + \epsilon_{Normal}$$

$\sum \tau_z Z = \text{sum of random effects of random variable } Z$

11. Winglength of 12 mosquitos (3 cages, 4 flies per cage). The left wing of each fly was measured twice.

Source	df	SS	MS	F	----> p
Cage	2	665.68	332.84	1.74	0.23
Fly⊂Cage	9	1720.68	191.19	147.07	<0.0001
<u>Error</u>	<u>12</u>	<u>15.62</u>	1.3017		
Total	23	2401.97			

ANOVA table
 Table 10.1 in Sokal and Rohlf (1995).

Write the model from the Source and df columns in the ANOVA table

Show how each df was calculated: 2 = _____ 9 = _____
 23 = _____ 12 = _____

Note that the Cage F-ratio was not calculated with respect to the MS error.
 The Cage F-ratio was calculated from a random factor, Fly(Cage). Why? Stay tuned.

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt4/Ch13_6.pdf

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/FisherEx38.csv>

GLM with two random factors

2nd example - - >

Nested - Random within Random

Crossed - Random \times Random

12. Fisher's Table 42 (Example 38) shows a nested design.

Plate	Sample			
	I	II	III	IV
1	72	74	78	69
2	69	72	74	67
3	63	70	70	66
4	59	69	58	64
5	59	66	58	62
6	53	58	56	58
7	51	52	56	54
Total	426	461	450	440
Mean	60.86	65.86	64.29	62.86

It ignores the fact that each plate was inoculated with subsamples from each of the four initial samples (Classes). Consequently, we can treat class (*i.e.* sample) as a random factor with 4 levels and cross it with another random factor, plate.

Assign symbols to both explanatory variables and write a two way random effects GLM with an interaction term.

Symbols _____

Model _____

Complete the Source and df columns of the ANOVA table for this model.

The correct model is a saturated model, the error term will have zero degrees of freedom.

We'll use this in the next session.

GLMM with two explanatory variables 2 examples Fixed + Random
Fixed × Random

The GLM assumes a normal error with fixed (constant) variance = ε_{Normal}

Grouped data often violate this assumption.-- > heterogeneous residuals

Paired data, clustered data, blocked data

Repeated measures (e.g. 3 samples at once), longitudinal data (3 sequential samples)

To capture this heterogeneity, we write a General Linear Mixed Model, which has both fixed and random effects.

$$Y = \beta_o + \Sigma\beta_x X + \Sigma\tau_z Z + \varepsilon_{Normal}$$

$\Sigma\beta_x X$ = sum of fixed effects

$\Sigma\tau_z Z$ = sum of heterogeneous random effects

ε_{Normal} = homogeneous normal errors

Random or Fixed? The definition of fixed versus random differs among text books.

Definition from Quinn and Keough (2002)

There are two types of categorical predictor variables in linear models. The most common type is a fixed factor, where all the levels of the factor (*i.e.* all the groups or treatments) that are of interest are included in the analysis.

We cannot extrapolate our statistical conclusions beyond these specific levels to other groups or treatments not in the study. If we repeated the study, we would usually use the same levels of the fixed factor again. Linear models based on fixed categorical predictor variables (fixed factors) are termed fixed effects models (or Model 1 ANOVAs). Fixed effect models are analogous to linear regression models where X is assumed to be fixed.

The other type of factor is a random factor, where we are only using a random selection of all the possible levels (or groups) of the factor and we usually wish to make inferences about all the possible groups from our sample of groups. If we repeated the study, we would usually take another sample of groups from the population of possible groups.

Drawing a branching tree diagram is not a reliable way to distinguish crossed from nested designs.

Why? Because a crossed design can be drawn as a branching tree.

The reliable way to distinguish crossed and nested designs is to write all of the two way tables and fill in the sample size in each cell of each table. If all (or most) of the cells have at least one sample then the two variables are crossed. If not the two factors are nested. For three factors there are three pairs and so three two-way tables.

GLMM with two explanatory variables First example

Fixed + Random

Wheat Yields

13. Wheat Yields from Cornell (1971)

Three pots were assigned to each treatment.

The two-way (Pot × Treatment) table now has 12 cells.

There is 1 sample in each cell.

When we do the cross test the design appears to be crossed.

However, there were 12 pots in the experiment, not 3.

Treatment	Pot Number	Plant number		
		1	2	3
None	1	20.6	22.3	19.8
None	2	23.4	21.9	22.8
None	3	21.8	20.6	21.3
Straw	1	13.6	13.9	14.2
Straw	2	13.7	14.5	13.8
Straw	3	12.9	13.1	13.4
Straw + PO4	1	14.8	14.6	14.9
Straw + PO4	2	14.3	13.9	13.5
Straw + PO4	3	14.4	13.8	14.1
Straw+PO4+lime	1	14.1	13.8	14.3
Straw+PO4+lime	2	14.0	13.9	14.2
Straw+PO4+lime	3	14.4	14.1	13.6

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/WheatYield.csv>

Recode the Pot variable to show that there are 12 pots.

The two-way (Pot × Treatment) table now has 36 cells.

Most of the cells are empty.

We cannot estimate Pot × Treatment.

Pot is nested within treatment Pot(Treatment)

Carry out the cross test for Pot × Plant and Trt × Plant.

Now many cells? _____

How many empty cells? _____

Can Pot × Plant be estimated ? Y/N _____

Can Trt × Plant be estimated ? Y/N _____

Treatment	Pot Number	Plant number		
		1	2	3
None	1	20.6	22.3	19.8
None	2	23.4	21.9	22.8
None	3	21.8	20.6	21.3
Straw	4	13.6	13.9	14.2
Straw	5	13.7	14.5	13.8
Straw	6	12.9	13.1	13.4
Straw + PO4	7	14.8	14.6	14.9
Straw + PO4	8	14.3	13.9	13.5
Straw + PO4	9	14.4	13.8	14.1
Straw+PO4+lime	10	14.1	13.8	14.3
Straw+PO4+lime	11	14.0	13.9	14.2
Straw+PO4+lime	12	14.4	14.1	13.6

GLMM with two explanatory variables 2nd example Fixed × Random

Subject	Drug A	Drug B
1	0.7	1.9
2	-1.6	0.8
3	-0.2	1.1
4	-1.2	0.1
5	-0.1	-0.1
6	3.4	4.4
7	3.7	5.5
8	0.8	1.6
9	0.0	4.6
10	2.0	3.4

14. Sleep data (Cushny and Peebles), used by Student (W. Gossett) to introduce the *t*-test. Data are: hours of extra sleep with two drugs Hyoscyamine (Drug A) and L Hyoscine (Drug B), each administered to 10 subjects. Values reported are averages. The pairing across subject allows us to remove the effects of individual variation.

Assign a symbol to the response variable _____

For each explanatory variable assign a symbol and state reason for assigning it as Fixed or Random

http://www.mun.ca/biology/schneider/b4605/LNotes/Pt4/Ch13_3.pdf

<http://www.mun.ca/biology/schneider/b4605/GLMMworkshop/Data/ExtraSleep.csv>

Crossed or Nested?

There are only two variables, hence only one interaction term.
We can see right away that this is a crossed design.