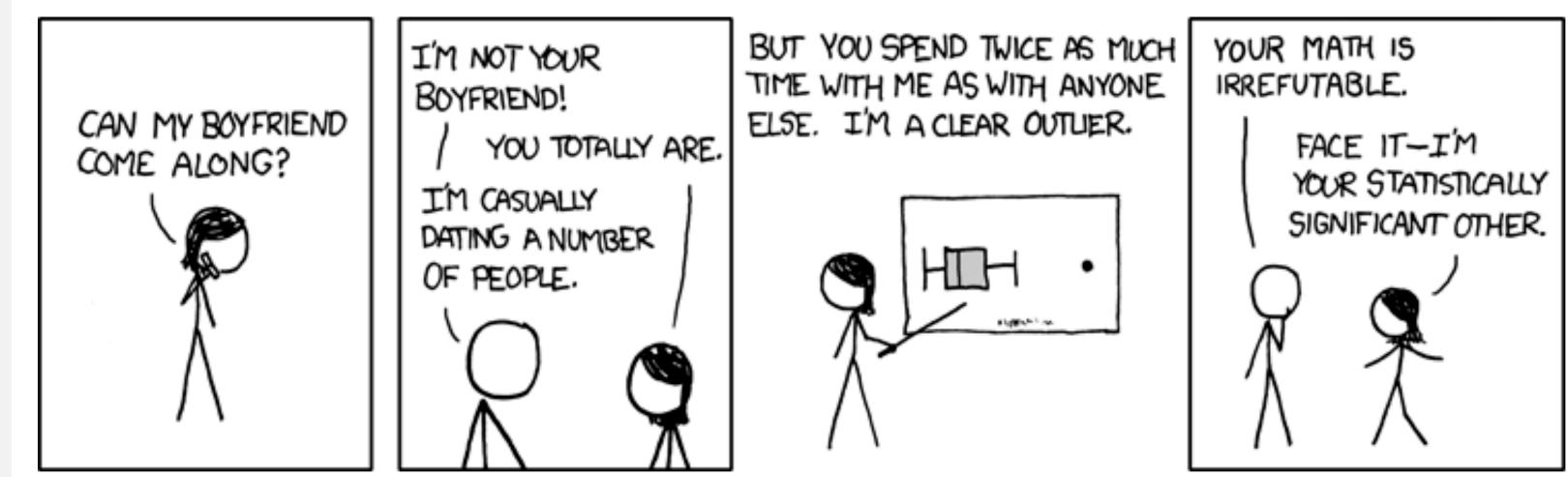


Random vs Fixed Effects Outliers

Biology 683

Lecture 8

Heath Blackmon



Last week

1. Regression
2. General linear model – all fixed effects

Today

- Fixed effects
 - Two-factor ANOVA
- Random effects
- Mixed effects models

What is a fixed effect

These are the variables whose impact we wish to determine

- Characteristics of the media or habitat
- Alternate medical treatments
- Age groups
- Time points

Conclusions that you reach are only applicable to the groups or treatments you include in the study

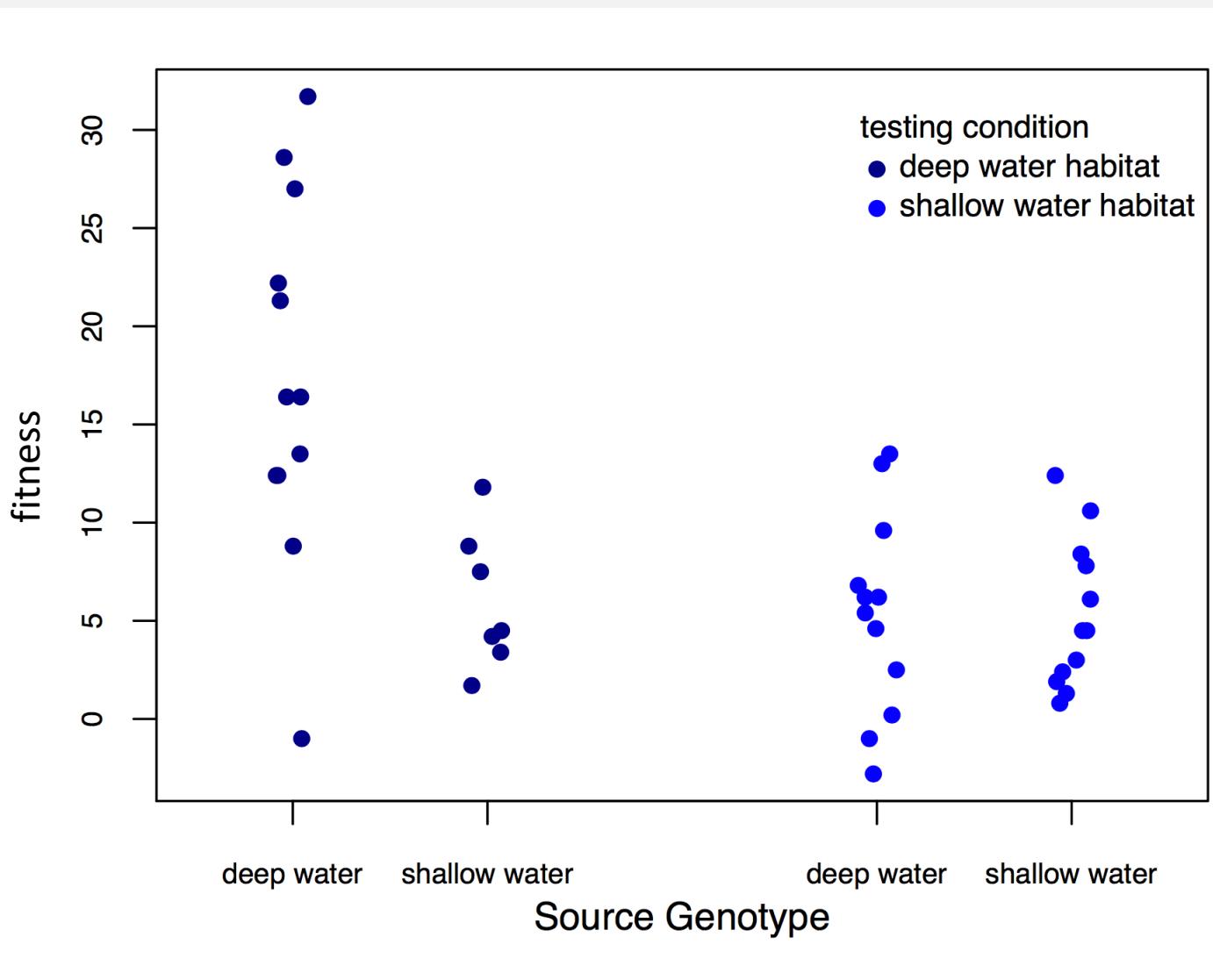
Example of fixed effects

Reciprocal relocation experiment to investigate how genotype and habitat interact to determine the fitness of stickleback fish (Rundle 2002).

		Source habitat	
		Shallow	Deep
Test habitat	Shallow	12 fish	11 fish
	Deep	7 fish	11 fish



Example of fixed effects (two factor ANOVA)



```
> anova(lm(fitness ~ genotype * test.habitat))
```

Analysis of Variance Table

Response: fitness

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
genotype	1	664.01	664.01	17.7278	0.0001452 ***
test.habitat	1	250.03	250.03	6.6753	0.0136378 *
genotype:test.habitat	1	333.58	333.58	8.9059	0.0048864 **
Residuals	39	1460.77	37.46		

What is a random effect

These are randomly sampled categories of a variable that represent groups of individual measurements. Usually random effects are not repeatable.

- Study sites
- Environmental chambers
- Families made up of siblings
- Measurements within individuals

Conclusions that you reach are applicable to the population being studied – beyond the levels included in the experiment.

What is a random effect

Sometimes random effects are a nuisance

- Field sites
- Environmental chambers
- Field plots
- Repeated measures

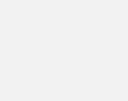
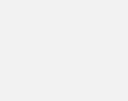
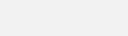
Occasionally random effects are of great interest

- Families - Heritability
- Individuals – Breeding value

Example of random effects

Impact of selective regime on horn size.

Measure both the left and right horn in 25 beetles from two different selective regimes.

Horn size	Beetle	Selective regime	
256	1	High	
276	1	High	
321	2	High	
321	2	High	
423	3	Low	
401	3	Low	
381	4	Low	
409	4	Low	



Example of random effects

Identifying the predictors for the presence or absence of Chrysina beetles.

Number collected	oak	juniper	site	date	trip
8	1	0	21	210	A
2	1	1	13	210	A
1	0	1	31	211	A
5	0	1	15	212	A
4	1	1	21	242	B
6	1	0	13	242	B
0	1	1	31	245	B
7	1	1	15	245	B



Implementing a mixed effects model

Mixed effect models can be fit using the LME function from the package nlme.

```
library(nlme)
fit <- lme(sqrt(beyeri) ~ oaks + jun + elev,
            random = list(~1|site,~1|trip),
            data=dat)
summary(fit)
```

Repeated measures at sites can't
be treated as independent

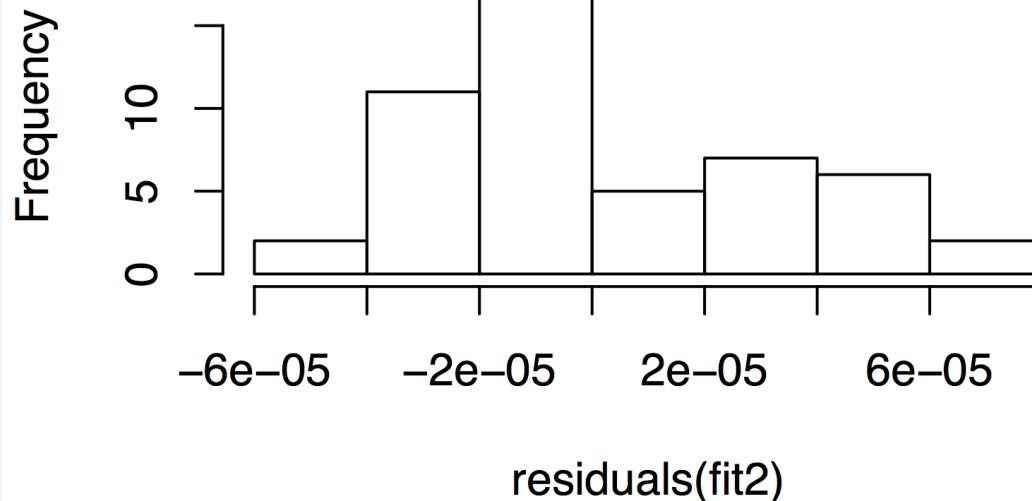
Fixed effects

Random effects

Implementing a mixed effects model

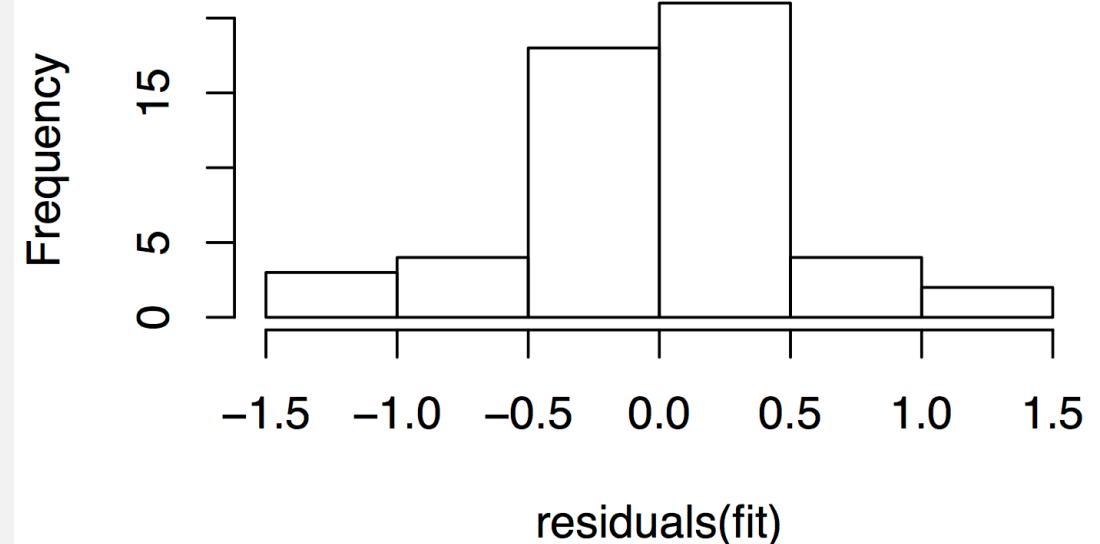
```
fit2 <- lme(beyeri ~ oaks + jun + elev,  
            random = list(~1|site, ~1|trip),  
            data=dat)
```

Histogram of residuals(fit2)



```
fit <- lme(sqrt(beyeri) ~ oaks + jun + elev,  
            random = list(~1|site, ~1|trip),  
            data=dat)
```

Histogram of residuals(fit)



Implementing a mixed effects model

Mixed effect models can be fit using the LME function from the package nlme.

```
fit <- lme(sqrt(beyeri) ~ oaks + jun + elev,  
           random = list(~1|site,~1|trip),  
           data=dat)  
> summary(fit)  
Linear mixed-effects model fit by REML  
Data: dat  
      AIC      BIC      logLik  
 153.6247 166.7231 -69.81233  
  
Random effects:  
 Formula: ~1 | site  
          (Intercept)  
 StdDev: 2.272341e-05  
  
 Formula: ~1 | trip %in% site  
          (Intercept) Residual  
 StdDev:  0.8169537 0.00264004  
  
Fixed effects: sqrt(beyeri) ~ oaks + jun + elev  
              Value Std.Error DF t-value p-value  
(Intercept) -1.6788177 1.5125428 26 -1.109931 0.2772  
oaks         0.9707025 0.2943403 22  3.297892 0.0033  
jun        -0.0860503 0.2460159 22 -0.349775 0.7298  
elev        0.0012513 0.0008072 22  1.550167 0.1354  
Correlation:  
   (Intr) oaks   jun  
oaks  0.530  
jun   0.197 -0.097  
elev -0.993 -0.584 -0.250
```

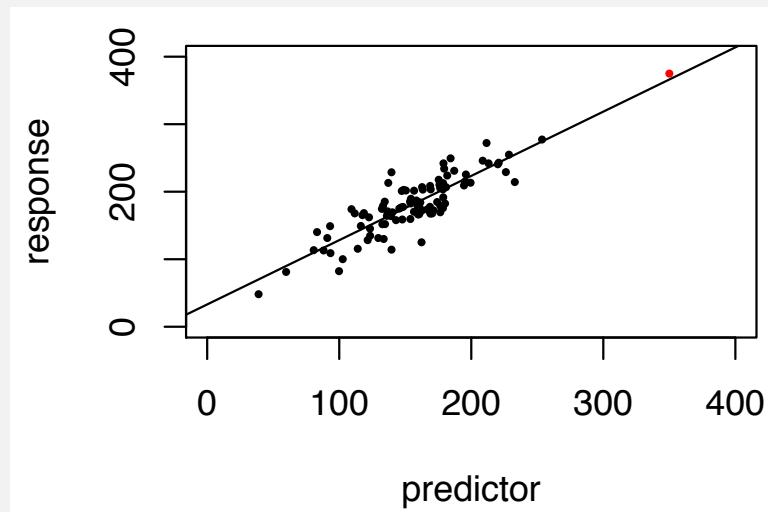
```
fit <- lme(sqrt(beyeri) ~ oaks + elev,  
           random = list(~1|site,~1|trip),  
           data=dat)  
> summary(fit)  
Linear mixed-effects model fit by REML  
Data: dat  
      AIC      BIC      logLik  
 150.7722 162.1231 -69.3861  
  
Random effects:  
 Formula: ~1 | site  
          (Intercept)  
 StdDev: 2.191491e-05  
  
Formula: ~1 | trip %in% site  
          (Intercept) Residual  
 StdDev:  0.8096043 0.002598713  
  
Fixed effects: sqrt(beyeri) ~ oaks + elev  
              Value Std.Error DF t-value p-value  
(Intercept) -1.5744919 1.4695032 26 -1.071445 0.2938  
oaks         0.9607576 0.2903283 23  3.309211 0.0031  
elev        0.0011807 0.0007746 23  1.524404 0.1410  
Correlation:  
   (Intr) oaks  
oaks  0.563  
elev -0.994 -0.631
```

```
fit <- lme(sqrt(beyeri) ~ oaks,  
           random = list(~1|site,~1|trip),  
           data=dat)  
> summary(fit)  
Linear mixed-effects model fit by REML  
Data: dat  
      AIC      BIC      logLik  
 138.5902 148.1504 -64.29512  
  
Random effects:  
 Formula: ~1 | site  
          (Intercept)  
 StdDev: 2.309943e-05  
  
Formula: ~1 | trip %in% site  
          (Intercept) Residual  
 StdDev:  0.8202517 0.002667513  
  
Fixed effects: sqrt(beyeri) ~ oaks  
              Value Std.Error DF t-value p-value  
(Intercept) 0.6514133 0.1674341 26  3.890566 6e-04  
oaks        1.2400606 0.2281742 24  5.434710 0e+00  
Correlation:  
   (Intr)  
oaks -0.734
```

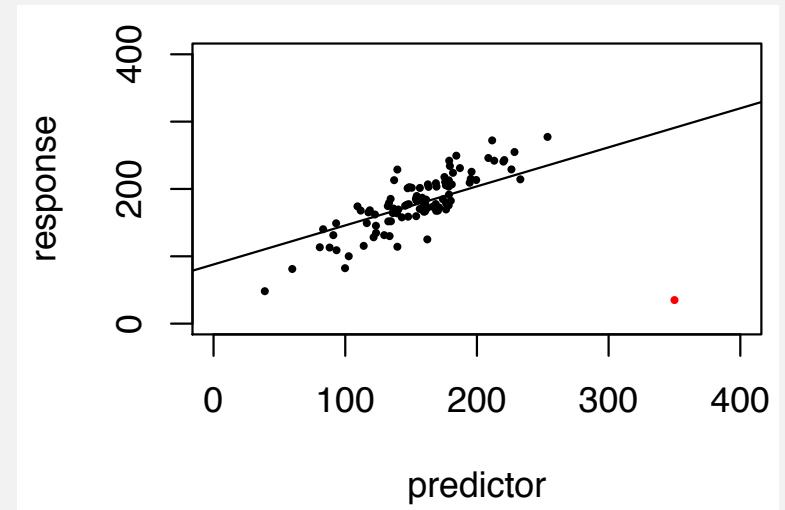
If you report your df with your F-statistic the reviewer will know if you did the right type of model

4 Types of outliers

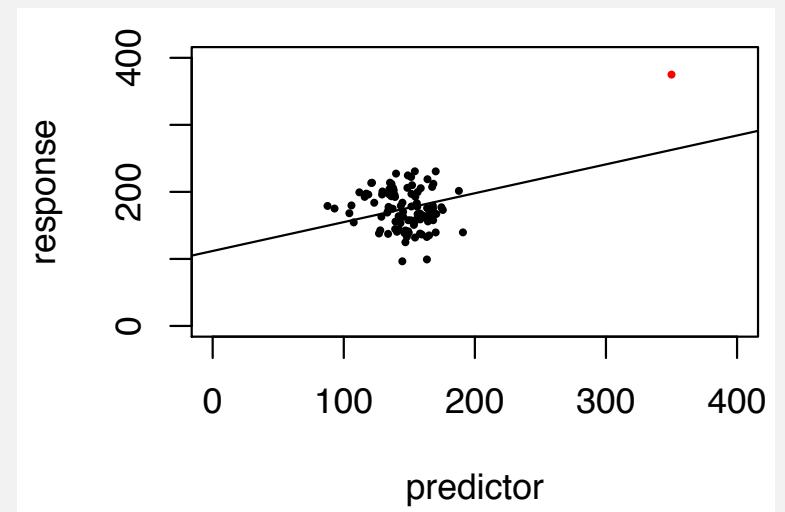
1) Obviously erroneously recorded or measured data



3) Extreme data point that impacts statistic of interest.



2) Extreme data point that doesn't impact statistic of interest.



4) Extreme data point that creates significance.

Considerations for models with random effects

- Most software will assume that all factors are fixed unless you specify them as mixed.
- Designating factors as random effects takes extra work.
- The lm function treats all predictors as fixed effects.
- Treating random effects as fixed effects is fundamentally wrong.

Links

[MCMCMglmm](#): Fit mixed models with phylogenetic or pedigree information in a Bayesian framework.

[Outlier Package](#): Apply outlier tests to identify possible outlier datapoints - I don't recommend this.

For Thursday

Bring laptop to class!

Plans for the rest of the semester.

Heath Blackmon

BSBW 309

coleoguy@gmail.com

@coleoguy

Models

"Remember that all models are wrong; the practical question is how wrong do they have to be to not be useful."

George Box