Statistics with Spa Rows II

Many models, matrices, and magic

Julia Schroeder

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Fixed part

Random part

- Fixed part
- How mean change with predictors

- Random part
- Partitions variances between groups

- Fixed part
- How mean change with predictors

- Random part
- Partitions variances between groups

Mean

Variance

- Fixed part
- How mean change with predictors

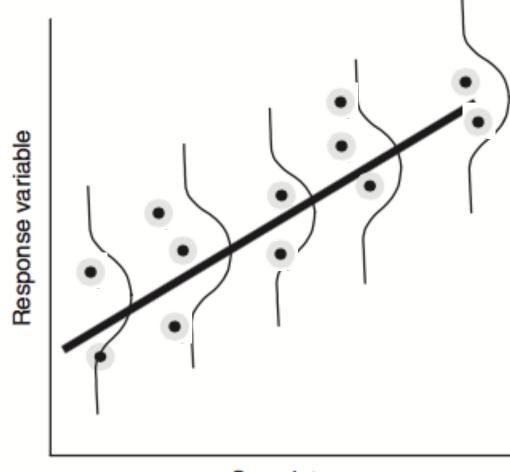
- Random part
- Partitions variances between groups

$$y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}$$

Mean

Variance

 Estimate variance components and fixed parameter estimates simultaneously



Covariate

 You can choose to only estimate one part and set the other one to fixed:

You can choose to only estimate one part

$$y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}$$

You could set the random part to zero (always keep the error!)

You can choose to only estimate one part

$$y_{i,j} = 1 + \alpha_j + \varepsilon_{i,j}$$

 You could set the fixed part to one (always have an intercept is a good idea)

 Or estimate both. Let's look at the components of a LMM just like we did with the lm:

$$y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}$$

• Or estimate both. Let's look at the components of a LMM just like we did with the lm:

```
y i b_0 = ? b_1 = ? x i ? 1 ? 1 ? 2 ? 2 ? 3 1 ? 4 3 ? 4 5
```

• First things first. What's with the j?

```
y \ i \ j \ b_0 = ? \ b_1 = ? \ x \ i \ j \ ? \ 1 \ 1 \ ? \ 2 \ 1 \ ? \ 2 \ 1 \ ? \ 3 \ 1 \ ? \ 3 \ 1 \ ? \ 4 \ 2 \ ? \ 4 \ 2 \ ? \ 5 \ 2
```

- First things first. What's with the j?
- J is a grouping factor. BirdID.

```
y i j b_0 = ? b_1 = ? x i j ? 1 1 ? 1 1 ? 2 1 ? 3 1 1 ? 3 1 1 ? 3 1 1 ? 4 3 1 ? 4 2 ? 4 5 2 ? 5 2
```

- First things first. What's with the j?
- J is a grouping factor. BirdID. Observer.

- First things first. What's with the j?
- J is a grouping factor. BirdID. Observer. NestID.

- First things first. What's with the j?
- J is a grouping factor. BirdID. Observer. NestID. It's a factor in your data. Year.

- First things first. What's with the j?
- J is a grouping factor. BirdID. Observer. NestID. It's a factor in your data. Year.
- It's categorical

$$y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}$$

• Ok. That only leaves alpha-j. What's that?

```
y \ i \ j \ 5 \ 1 \ 1 \ 3 \ 2 \ 1 \ 6 \ 3 \ 1 \ 1 \ 2 \ 1 \ 4 \ 3 \ 1 \ 1 \ 2 \ 2 \ 1 \ 10 \ 4 \ 2 \ 2 \ 5 \ 2
```

• Ok. That only leaves alpha-j. What's that?

```
Variance of
                                  b_1 = ?
                b_0 = ?
                                                                                 ? 1 1
                                             3 1 1
                                                               data
                                                                                 ? 2 1
                                             1 2 1
                                                              grouped
                                                                                 ? 3 1
                                             4 3 1
                                                               by j
                                                                                 ? 4 2
                                             8 4 2
10 4 2
                                                                                   5 2
                                             2 5 2
4 5 2
```

- Ok. That only leaves alpha-j.
- So it's one number, with a measure of precision, like the bs!

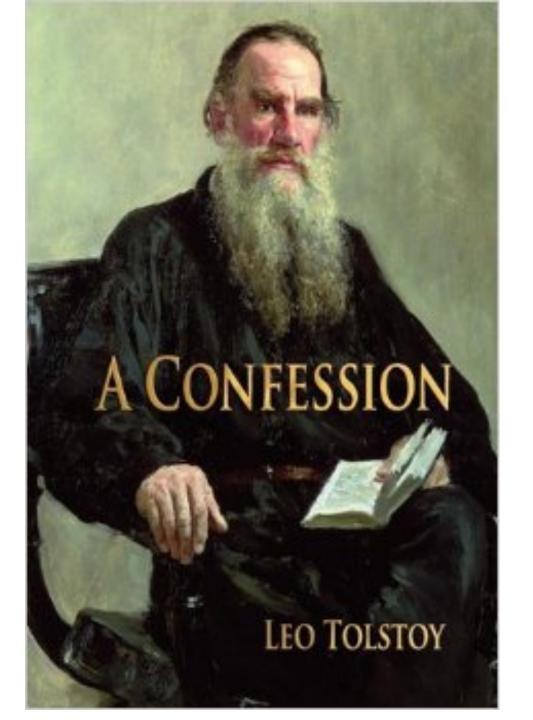
```
y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}
```

```
Variance of
                                   b_1 = ?
                 b_0 = ?
                                                                                   ? 1 1
                                                                data
                                                                                   ? 2 1
                                              1 2 1
                                                                grouped
                                                                                   ? 3 1
                                              4 3 1
                                                                by j
                                                                                   ? 4 2
                                              8 4 2
10 4 2
                                                                                     5 2
                                              2 5 2
```

- Ok. That only leaves alpha-j.
- So it's one number, with a measure of precision, like the bs!
- And we color it red because we want to estimate it!

```
y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}
```

```
y j Group
                             b_1 = ?
              b_0 = ?
                                                      5 1 variance of
                                                                     ? 1 1
                                      3 1 1
                                                      3 1 data y
                                                                     ? 2 1
                                      1 2 1
                                                      6 1 grouped
                                                                    ? 3 1
                                      4 3 1
                                                      10 2 by j
                                                                     ? 4 2
                                      8 4 2
10 4 2
                                                      4 2
                                                                     ? 5 2
                                      2 5 2
```



LMMs – a confession

• In this equation, e-ij is also a variance. The variance of the residuals. We've somewhat ignored this so far. But really, it's a number.

$$y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}$$

```
y j Group
                                                                              Variance of
                              b_1 = ?
              b_0 = ?
                                                       5 1 variance of ? 1 1
                                       3 1 1
                                                                              residuals
                                                       3 1 data y ? 2 1
                                       1 2 1
                                                       6 1 grouped ? 3 1
                                       4 3 1
                                                                      ? 4 2
                                                       10 2 by j
                                       8 4 2
10 4 2
                                       2 5 2
```

LMMs – a confession

• Ok, cool. Let's revisit that group variance thing.

```
y j Group
                                                                             Variance of
                                      хіј
                             b_1 = ?
              b_0 = ?
                                                      5 1 variance of ? 1 1
                                      3 1 1
                                                                            residuals
                                                      3 1 data y ? 2 1
                                      1 2 1
3 2 1
                                                      6 1 grouped ? 3 1
                                      4 3 1
                                                      10 2 by j ? 4 2
                                      8 4 2
10 4 2
                                      2 5 2
4 5 2
```

1,1,2,1,1,15,5,4,5,5,13,3,4,3,3,3

1,1,2,1,1,1 5,5,4,5,5,1 3,3,4,3,3,3

k = number of groups

n = sample size in group

N = total sample size

i = row

j = column

 \bar{x} = group mean

1,1,2,1,1,1 5,5,4,5,5,1 3,3,4,3,3,3

3 = k = number of groups

 $\hat{b} = n = \text{sample size in group}$

18 = N = total sample size

1:18 i = individual counter

1:3 j = group counter

 \bar{x} = group mean

j: 1,1,1,1,1,1,2,2,2,2,2,2,3,3,3,3,3,3,3

1,1,2,1,1,1 5,5,4,5,5,1 3,3,4,3,3,3

3 = k = number of groups

6 = n = sample size in group

18 = N = total sample size

1:18 i = individual counter

1:3 j = group counter \bar{x} = group mean

```
3 = k = number of groups
```

6 = n = sample size in group

18 = N = total sample size

1:18 i = individual counter

1:3 j = group counter \bar{x} = group mean

$$3 = k = number of groups$$

 $6 = n = sample size in group$
 $18 = N = total sample size$
 $1:18 i = individual counter$
 $1:3 j = group counter$
 $\bar{x} = group mean$

 $\bar{\bar{x}}$ = grand total mean

$$Y_{3.2} = 4$$

51

3 = k = number of groups 6 = n = sample size in group 18 = N = total sample size 1:18 i = individual counter 1:3 j = group counter $\bar{x} = group mean$

 $\bar{\bar{x}}$ = grand total mean

$$Y_{3.2} = 4$$

51

$$Y_{i,j}$$
: 1,1,2,1,1,15,5,4,5,5,1 3,3,4,3,3,3

$$Between-group\ variance = \frac{SSG}{k-1}$$

- 3 = k = number of groups
- 6 = n_i = sample size in group
- 18 = N = total sample size
- 1:6 i = individual counter
- 1:3 j = group counter \bar{x}_i = group mean
- 51 $\bar{\bar{x}}_{i,j}$ = grand total mean

$$Y_{i,j}$$
: 1,1,2,1,1,15,5,4,5,5,1 3,3,4,3,3,3

Between - group variance =
$$\frac{SSG}{k-1} = \frac{\sum_{j=1}^{k} n_j (\overline{x_j} - \overline{\overline{x_{i,j}}})^2}{k-1}$$

- 3 = k = number of groups
- $6 = n_i = \text{sample size in group}$
- 18 = N = total sample size
- 1:6 i = individual counter
- 1:3 j = group counter \bar{x}_i = group mean
- $\bar{x}_{i,i}$ = grand total mean

$$Y_{i,j}$$
: 1,1,2,1,1,15,5,4,5,5,13,3,4,3,3,3

$$Within-group\ variance = \frac{SSE}{N-k}$$

- 3 = k = number of groups
- 6 = n_i = sample size in group
- 18 = N = total sample size
- 1:6 i = individual counter
- 1:3 j = group counter \bar{x}_i = group mean
- 51 $\bar{\bar{x}}_{i,i}$ = grand total mean

1:3 j = group counter
$$\bar{x}_j$$
 = group mean

51
$$\bar{\bar{x}}_{i,j}$$
 = grand total mean

Within – group variance =
$$\frac{SSE}{N-k} = \frac{\sum_{j=1}^{k} \sum_{j=1}^{n_j} (x_{i,j} - x_j)^2}{N-k}$$

$$3 = k = number of groups$$

$$6 = n_i = \text{sample size in group}$$

1:3 j = group counter
$$\bar{x}_i$$
 = group mean

51
$$\bar{\bar{x}}_{i,i}$$
 = grand total mean

$$Within - group \ variance = \frac{SSE}{N-k} = \frac{\sum_{j=1}^{k} \sum_{j=1}^{n_j} (x_{i,j} - x_j)^2}{N-k}$$

$$Between - group \ variance = \frac{SSG}{k-1} = \frac{\sum_{j=1}^{k} n_j (\overline{x_j} - \overline{x_{i,j}})^2}{k-1}$$

$$Total\ variance = \frac{SST}{n-1}$$

Variances and groups

$$Y_{i,j}$$
: 1,1,2,1,1,15,5,4,5,5,1 3,3,4,3,3,3

$$3 = k = number of groups$$

$$6 = n_i = \text{sample size in group}$$

1:3 j = group counter
$$\bar{x}_i$$
 = group mean

51
$$\bar{\bar{x}}_{i,i}$$
 = grand total mean

$$Within - group \ variance = \frac{SSE}{N-k} = \frac{\sum_{j=1}^{k} \sum_{j=1}^{n_j} (x_{i,j} - x_j)^2}{N-k}$$

$$Between - group \ variance = \frac{SSG}{k-1} = \frac{\sum_{j=1}^{k} n_j (\overline{x_j} - \overline{x_{i,j}})^2}{k-1}$$

$$Total \ variance = \frac{SST}{n-1}$$

$$SST = SSG + SSE$$

Between-group variance

$y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}$

```
y i j
5 1 1
3 2 1
6 3 1
10 4 2
4 5 2
```

```
b_0 = ?
```

```
b<sub>1</sub> = ?

X i j
3 1 1
1 2 1
4 3 1
8 4 2
2 5 2
```

```
y j Group
5 1 variance of
3 1 data y
6 1 grouped
10 2 by j
4 2
```

```
ε i j Variance of ? 1 1 residuals ? 2 1 ? 3 1 ? 4 2 ? 5 2
```

Within-group variance

Variance of

residuals

Between-group variance



 $b_0 = ?$

b₁ = ?

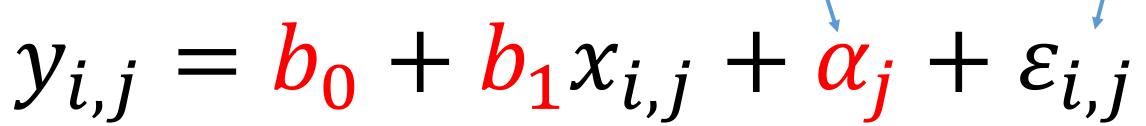
y j Group
5 1 variance of
3 1 data y
6 1 grouped
10 2 by j

4 2

2 1

Between-group variance

Within-group variance Residual variance



 $b_0 = ?$

b₁ = ?

y j Group
5 1 variance of
3 1 data y
6 1 grouped
10 2 by j
4 2

ε i j Variance of ? 1 1 residuals? 2 1? 3 1? 4 2? 5 2

Between-group variance Residual variance Error variance $= b_0 + b_1 x_{i,i} + \alpha_i + \varepsilon_{i,i}$ Group **Variance of** $b_1 = ?$ $b_0 = ?$ variance of residuals data y 2 1

4 2

5 2

10 4 2

Within-group variance

3 1

4 2

5 2

grouped

by j

10 2

4 2

Between-group variance

Within-group variance
Residual variance
Error variance

Total variance =
$$\alpha_i + \varepsilon_{i,j}$$

$y_{i,j} = b_0 + b_1 x_{i,j} + \alpha_j + \varepsilon_{i,j}$

```
y i j
5 1 1
3 2 1
6 3 1
10 4 2
4 5 2
```

 $b_0 = ?$

 $b_1 = ?$

 γ j
 Group
 ε i j

 5 1
 variance of
 ? 1 1

 3 1
 data y
 ? 2 1

 6 1
 grouped
 ? 3 1

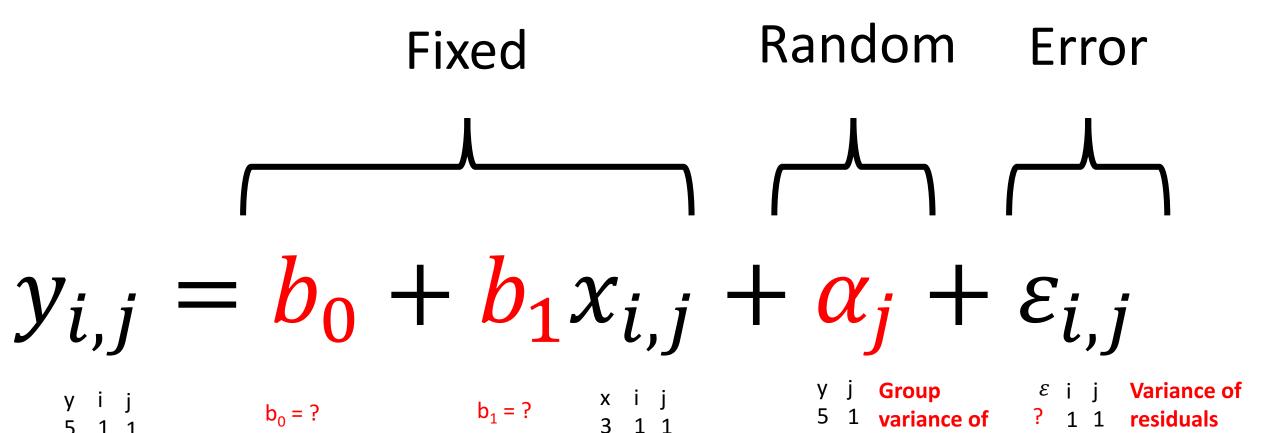
 10 2
 by j
 ? 4 2

ε i j Variance of ? 1 1 residuals ? 2 1 ? 3 1 ? 4 2 ? 5 2

Between-group variance

4 2

Residual variance



1 2 1

4 3 1

8 4 2

2 5 2

10 4 2

4 5 2

Between-group variance

4 2

3 1 **data y**

10 2 **by** j

6 1 grouped

Residual variance

? 2 1

? 3 1

? 4 2

5 2

• Estimate variance components simultaneously to fixed terms

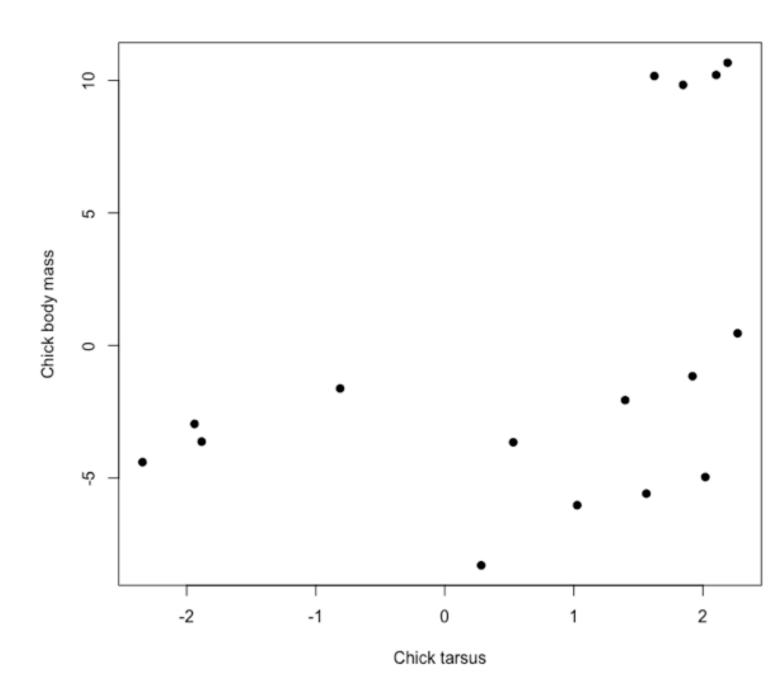
- Estimate variance components simultaneously to fixed terms
- Allow to account for nested structure in data

- Estimate variance components simultaneously to fixed terms
- Allow to account for nested structure in data
- Some things are more similar than others

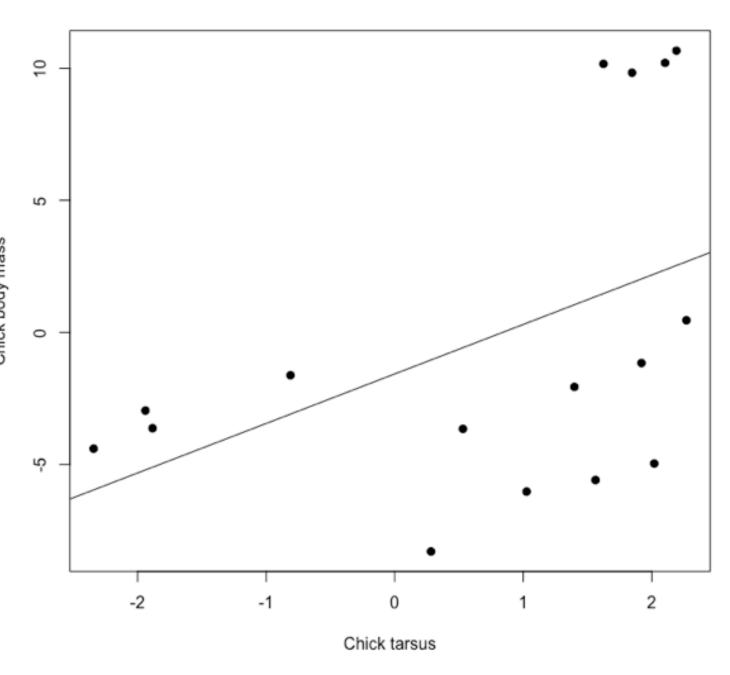




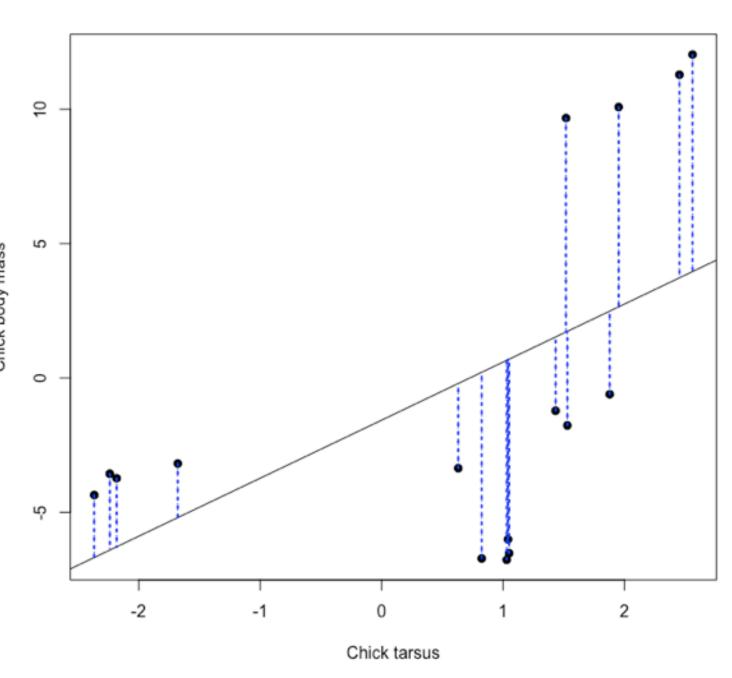
- Estimate variance components simultaneously to fixed terms
- Allow to account for nested structure in data
- Some things are more similar than others



```
> m<-(lm(bm~tarsus))
> summary(m)
Call:
lm(formula = bm \sim tarsus)
                                                            Chick body mass
Residuals:
    Min
             10 Median
-7.2535 -3.9865 -0.3775 3.6447 8.6954
Coefficients:
            Estimate Std. Error t value Pr(>ItI)
(Intercept) -1.570
                                 -0.944
                          1.662
                                          0.3610
               1.875
                          0.965
                                  1.943
                                          0.0725 .
tarsus
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '
Residual standard error: 6.008 on 14 degrees of freedom
Multiple R-squared: 0.2123, Adjusted R-squared: 0.1561
F-statistic: 3.774 on 1 and 14 DF, p-value: 0.07246
>
```



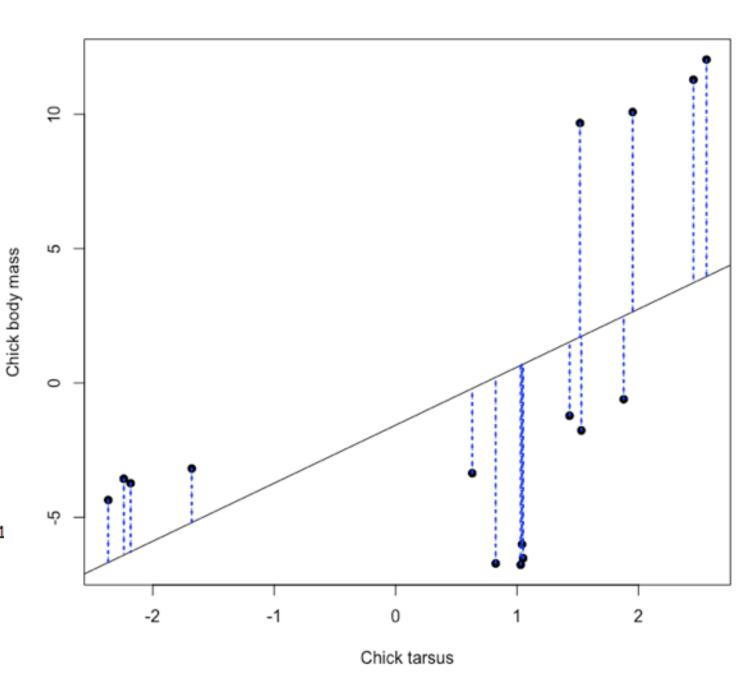
```
> m<-(lm(bm~tarsus))
> summary(m)
Call:
lm(formula = bm \sim tarsus)
                                                           Chick body mass
Residuals:
   Min
             10 Median
-7.2535 -3.9865 -0.3775 3.6447 8.6954
Coefficients:
            Estimate Std. Error t value Pr(>ItI)
(Intercept) -1.570
                          1.662
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                                          0.3610
               1.875
                          0.965
                                 1.943
                                          0.0725 .
tarsus
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>
```

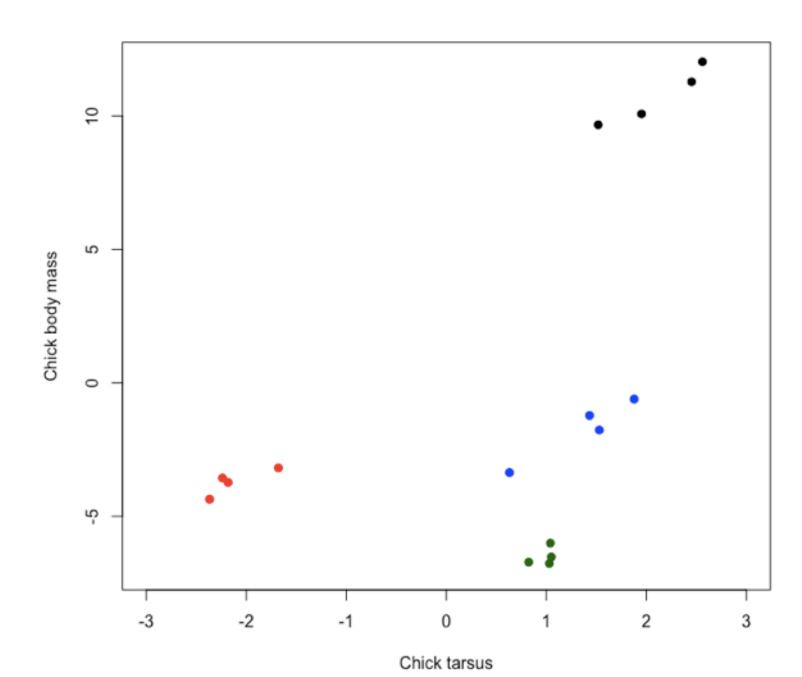


```
> m<-(lm(bm~tarsus))
> summary(m)
Call:
lm(formula = bm \sim tarsus)
Residuals:
    Min
             10 Median
-7.2535 -3.9865 -0.3775 3.6447 8.6954
Coefficients:
            Estimate Std. Error t value Pr(>ItI)
(Interc
               -1.5/0
               1.875
                                          0.0725 .
tarsus
                          0.965
                                  1.943
                0 (***, 0.001 (**, 0.01 (*, 0
```

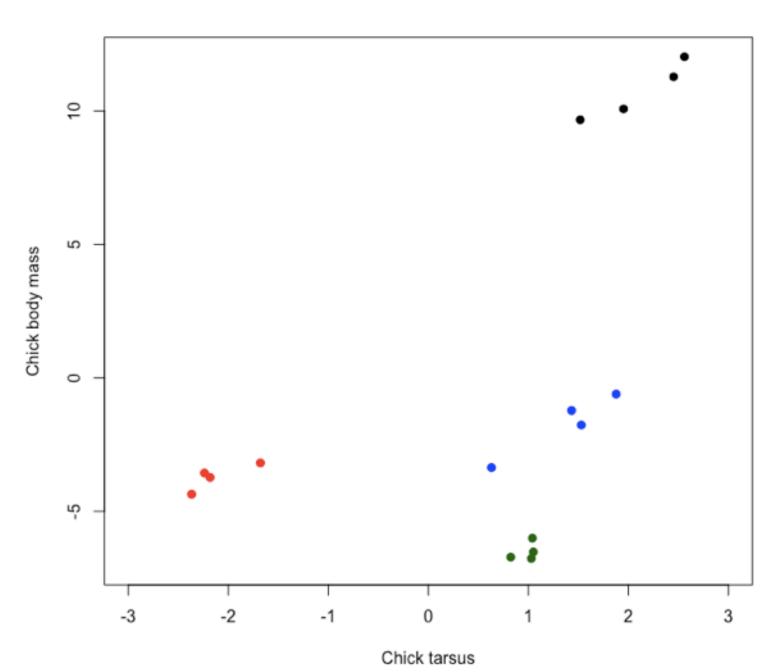
Residual standard error: 6.008 on 14 degrees of freedom Multiple R-squared: 0.2123, Adjusted R-squared: 0.1561 F-statistic: 3.774 on 1 and 14 DF, p-value: 0.07246

>

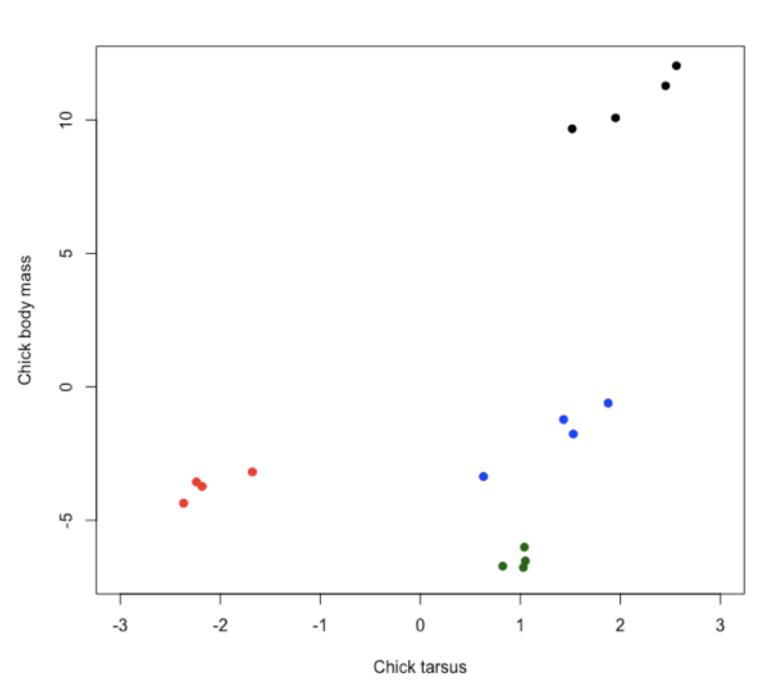




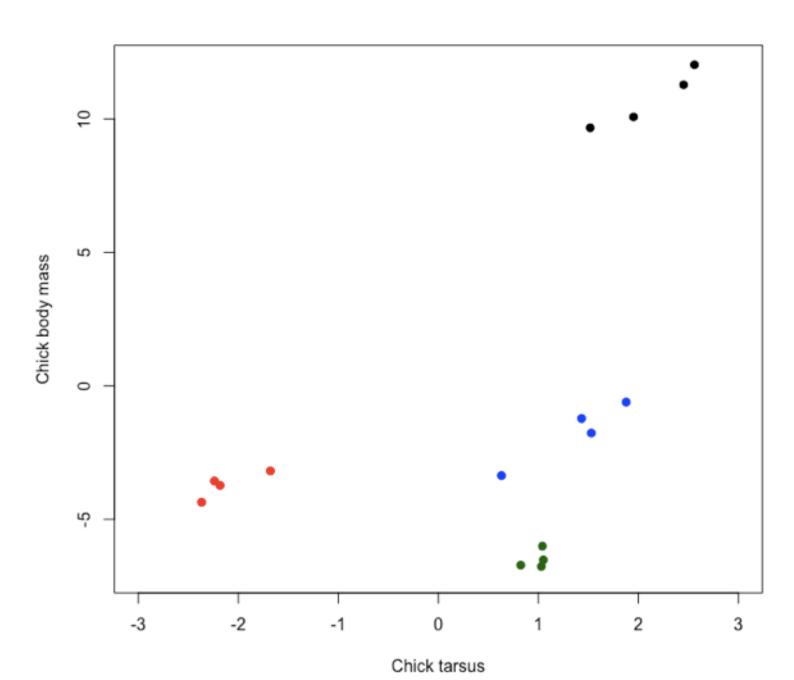
```
> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
```



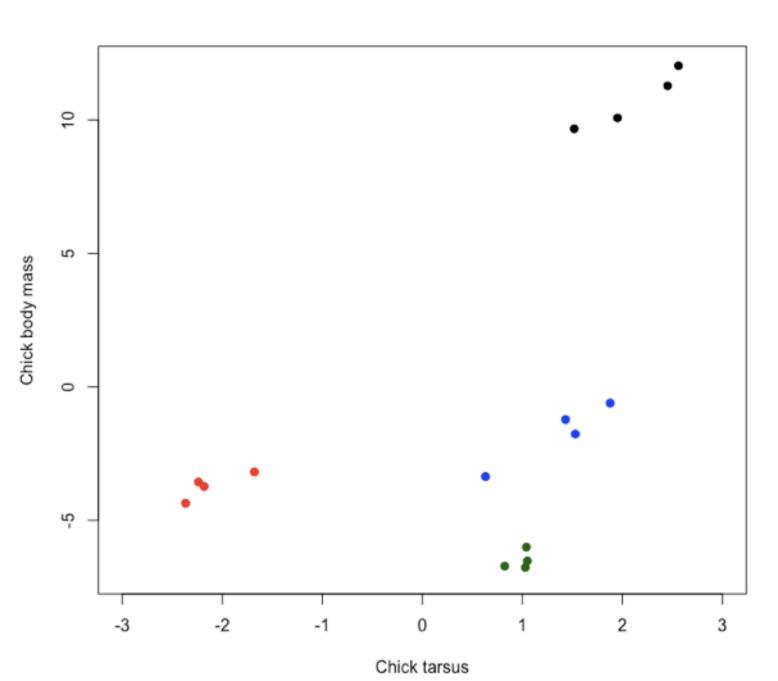
```
> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
> summary(mm)
Linear mixed model fit by REML ['lmerMod']
Formula: bm ~ tarsus + (1 | nest)
REML criterion at convergence: 34.4
Scaled residuals:
     Min
               10
                   Median
                                 30
                                          Max
-1.08739 -0.60062 -0.05266 0.58491 1.18030
Random effects:
                      Variance Std.Dev.
Groups
          Name
nest
          (Intercept) 42.1465 6.492
Residual
                       0.1115 0.334
Number of obs: 16, groups: nest, 4
Fixed effects:
            Estimate Std. Error t value
(Intercept)
             -1.4939
                         3.2503
                                  -0.46
              2.0325
                         0.2446
                                   8.31
tarsus
```



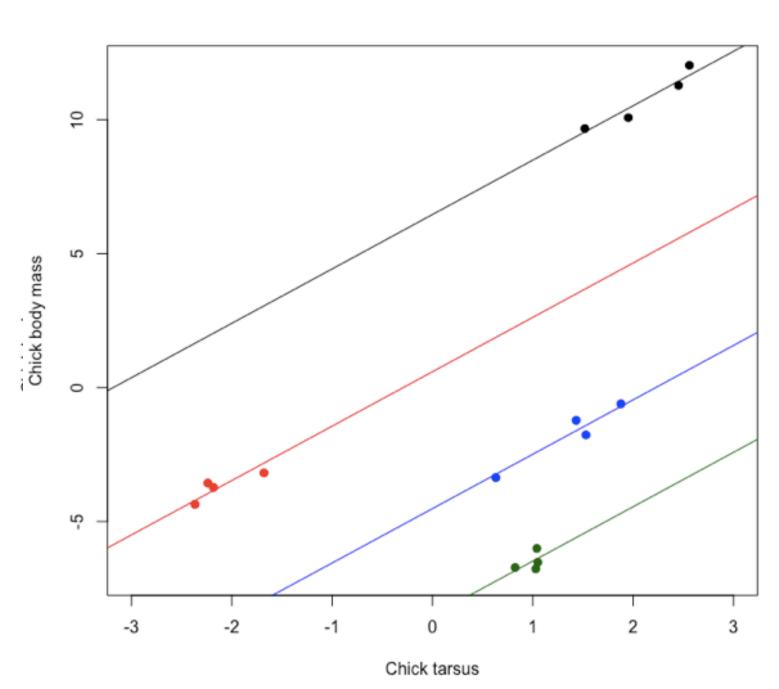
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> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
> summary(mm)
Linear mixed model fit by REML ['lmerMod']
Formula: bm ~ tarsus + (1 | nest)
REML criterion at convergence: 34.4
Scaled residuals:
     Min
               10
                    Median
                                  30
                                          Max
-1.08739 -0.60062 -0.05266 0.58491 1.18030
Random effects:
                      Variance Std.Dev.
Groups
          Name
nest
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Residual
                       0.1115 0.334
Number of obs: 16, groups: nest, 4
Fixed effects:
            Estimate Std. Error t value
(Intercept)
             -1.4939
                         3.2503
                                   -0.46
              2.0325
                         0.2446
                                   8.31
tarsus
```



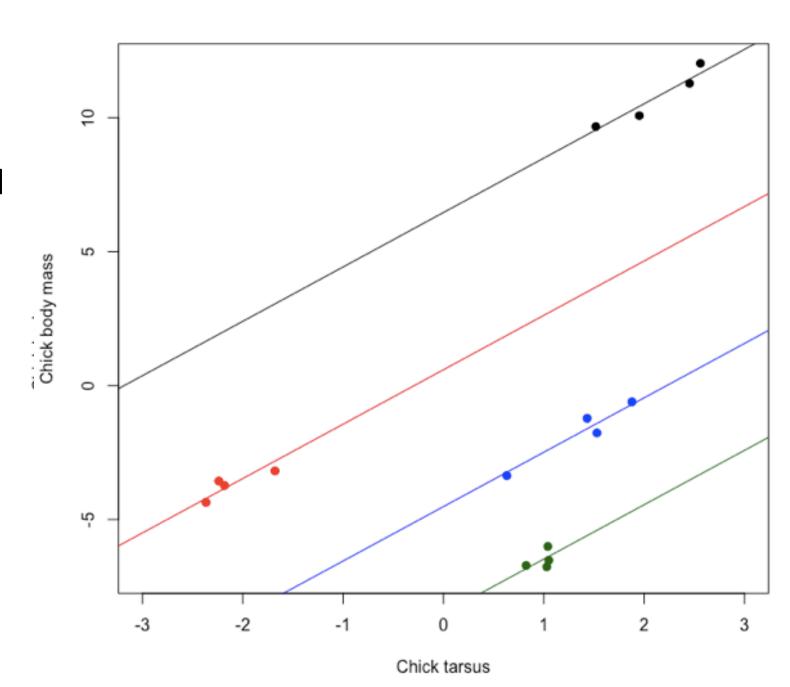
```
> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
> summary(mm)
Linear mixed model fit by REML ['lmerMod']
Formula: bm ~ tarsus + (1 | nest)
REML criterion at convergence: 34.4
Scaled residuals:
     Min
               10
                    Median
                                  30
                                          Max
-1.08739 -0.60062 -0.05266 0.58491 1.18030
Random effects:
                      Variance Std.Dev.
Groups
          Name
nest
          (Intercept) 42.1465 6.492
Residual
                       0.1115 0.334
Number of abs: 16, groups: nest, 4
Fixed effects:
            Estimate Std. Error t value
(Intercept)
             -1.4939
                         3.2503
                                   -0.46
              2.0325
                         0.2446
                                   8.31
tarsus
```



```
> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
> summary(mm)
Linear mixed model fit by REML ['lmerMod']
Formula: bm ~ tarsus + (1 | nest)
REML criterion at convergence: 34.4
Scaled residuals:
    Min
               10 Median
-1.08739 -0.60062 -0.05266 0.58491 1.18030
Random effects:
                      Variance Std.Dev.
Groups
         Name
nest
          (Intercept) 42.1465 6.492
Residual
                       0.1115 0.334
Number of obs: 16, groups: nest, 4
Fixed effects:
            Estimate Std. Error t value
(Intercept)
             -1.4939
                         3.2503
                                  -0.46
              2.0325
                         0.2446
                                   8.31
tarsus
```



 Ok, so why don't we add nest as fixed factor instead?



Random

```
> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
> summary(mm)
Linear mixed model fit by REML ['lmerMod']
Formula: bm ~ tarsus + (1 | nest)
REML criterion at convergence: 34.4
Scaled residuals:
    Min
              10 Median
                                30
                                        Max
-1.08739 -0.60062 -0.05266 0.58491 1.18030
Random effects:
                    Variance Std.Dev.
Groups Name
nest (Intercept) 42.1465 6.492
Residual
                      0.1115 0.334
Number of obs: 16, groups: nest, 4
Fixed effects:
           Estimate Std. Error t value
(Intercept) -1.4939 3.2503 -0.46
             2.0325
                        0.2446 8.31
tarsus
```

Fixed

```
> summary(lm(bm~tarsus+nest))
Call:
lm(formula = bm \sim tarsus + nest)
Residuals:
    Min
             10 Median
                                     Max
-0.36367 -0.20454 -0.01581 0.18941 0.39257
Coefficients:
           Estimate Std. Error t value Pr(>ItI)
(Intercept) 6.4606
                      0.5490 11.767 1.42e-07 ***
tarsus
      2.0305 0.2466 8.234 4.96e-06 ***
nestB -5.8707 1.0715 -5.479 0.000192 ***
nestC -10.9770 0.3005 -36.525 7.81e-13 ***
nestD -14.9654 0.3663 -40.858 2.29e-13 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.3342 on 11 degrees of freedom
Multiple R-squared: 0.9983, Adjusted R-squared: 0.9976
F-statistic: 1581 on 4 and 11 DF, p-value: 4.275e-15
```

Random

```
> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
> summary(mm)
Linear mixed model fit by REML ['lmerMod']
Formula: bm ~ tarsus + (1 | nest)
REML criterion at convergence: 34.4
Scaled residuals:
    Min
              10 Median
                                30
                                        Max
-1.08739 -0.60062 -0.05266 0.58491 1.18030
Random effects:
                     Variance Std.Dev.
Groups Name
nest (Intercept) 42.1465 6.492
Residual
                      0.1115 0.334
Number of obs: 16, groups: nest, 4
Fixed effects:
            Estimate Std. Error t value
(Intercent)
            _1_/020
                      3 2503 -0.46
             2.0325
                        0.2446
                                  8.31
tarsus
```

Fixed

```
> summary(lm(bm~tarsus+nest))
Call:
lm(formula = bm \sim tarsus + nest)
Residuals:
     Min
              10 Median
                                       Max
-0.36367 -0.20454 -0.01581 0.18941 0.39257
Coefficients:
            Estimate Std. Error t value Pr(>ItI)
(Total copt)
                        0.5490 11.767 1.42e-07 ***
             b.460b
                        0.2466 8.234 4.9 e-06 ***
tarsus
             2.0305
             5 9707 1 0715 3.479 0.000192 ***
nestb
          -10.9770 0.3005 -36.525 7.81e-13 ***
nestC
       -14.9654 0.3663 -40.858 2.29e-13 ***
nestD
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.3342 on 11 degrees of freedom
Multiple R-squared: 0.9983, Adjusted R-squared: 0.9976
```

F-statistic: 1581 on 4 and 11 DF, p-value: 4.275e-15

Random

```
> require(lme4)
> mm<-(lmer(bm~tarsus+(1Inest)))</pre>
> summary(mm)
Linear mixed model fit by REML ['lmerMod']
Formula: bm ~ tarsus + (1 | nest)
REML criterion at convergence: 34.4
Scaled residuals:
    Min
              10 Median
                                30
                                        Max
-1.08739 -0.60062 -0.05266 0.58491 1.18030
Random effects:
                     Variance Std.Dev.
Groups Name
nest (Intercept) 42.1465 6.492
Residual
                      0.1115 0.334
Number of obs: 16, groups: nest, 4
Fixed effects:
           Estimate Std. Error t value
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             2.0325
                        0.2446
                                  8.31
tarsus
```

Fixed

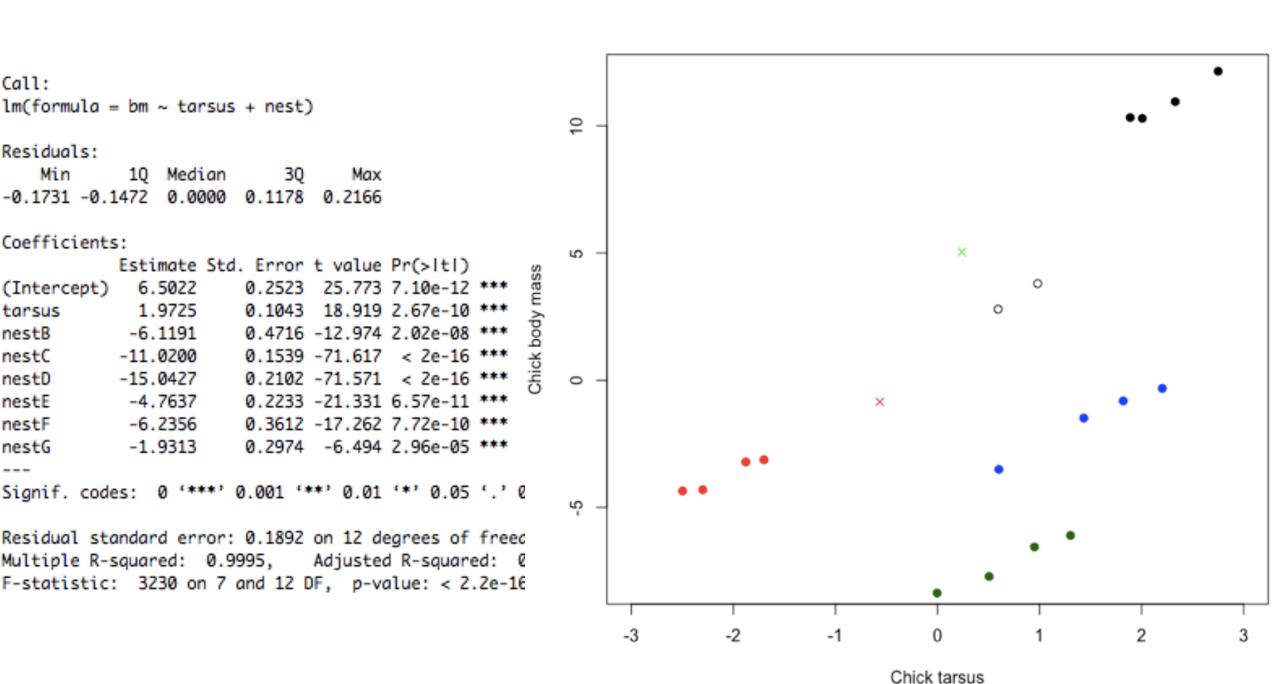
```
> summary(lm(bm~tarsus+nest))
Call:
lm(formula = bm \sim tarsus + nest)
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    Min
             10 Median
                                     Max
-0.36367 -0.20454 -0.01581 0.18941 0.39257
Coefficients:
           Estimate Std. Error t value Pr(>ItI)
(Intercept) 6.4606
                      0.5490 11.767 1.42e-07 ***
tarsus
        2.0305 0.2466 8.234 4.96e-06 ***
nestB -5.8707 1.0715 -5.479 0.000192 ***
      -10.9770 0.3005 -36.525 7.81e-13 ***
nestC
nestD
      -14.9654 0.3663 -40.858 2.29e-13 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.33 2 on 11 degrees of freedom
Multiple R-squared: 0.9983, Adjusted R squared. 0.9976
F-statistic: 1581 on 4 and 11 DF, p-value: 4.275e-15
```

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Fixed

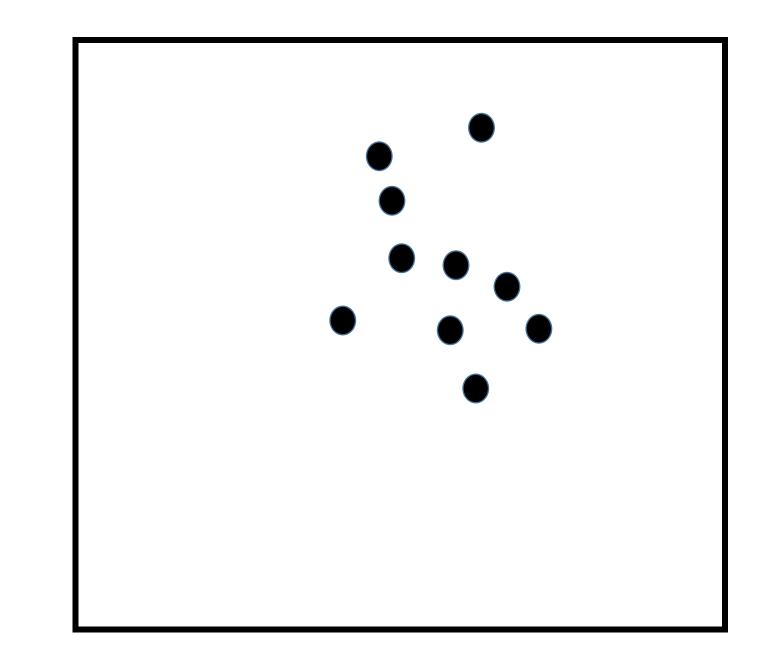
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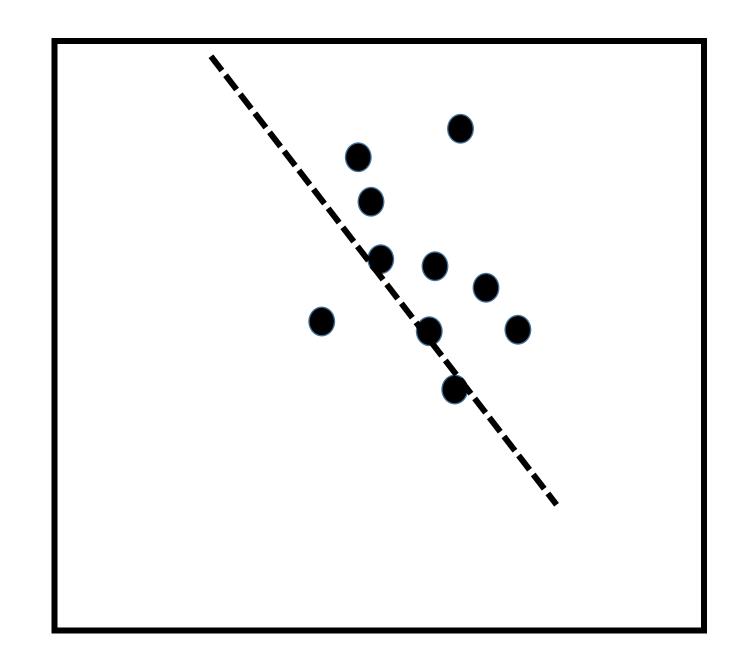


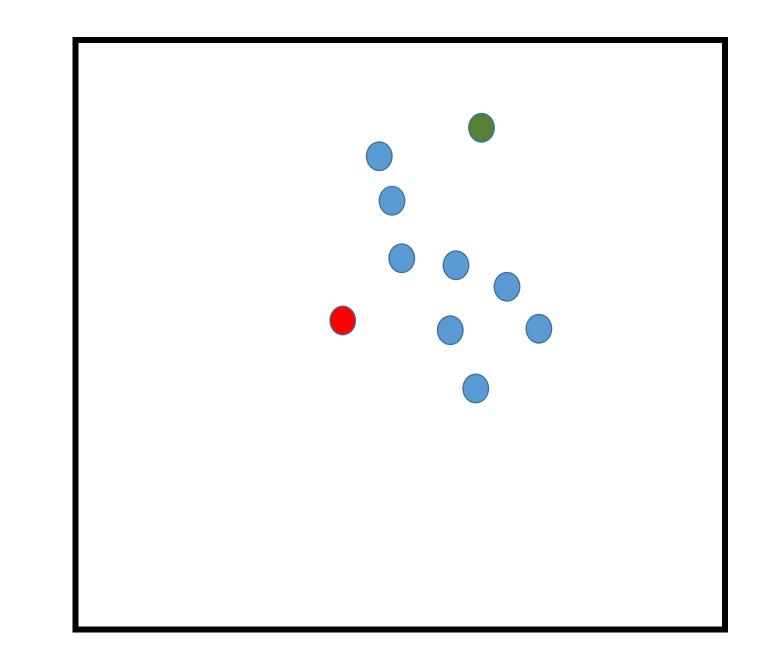
- Estimates variance components simultaneously to fixed terms
- Allow to account for nested structure in data with lots of levels

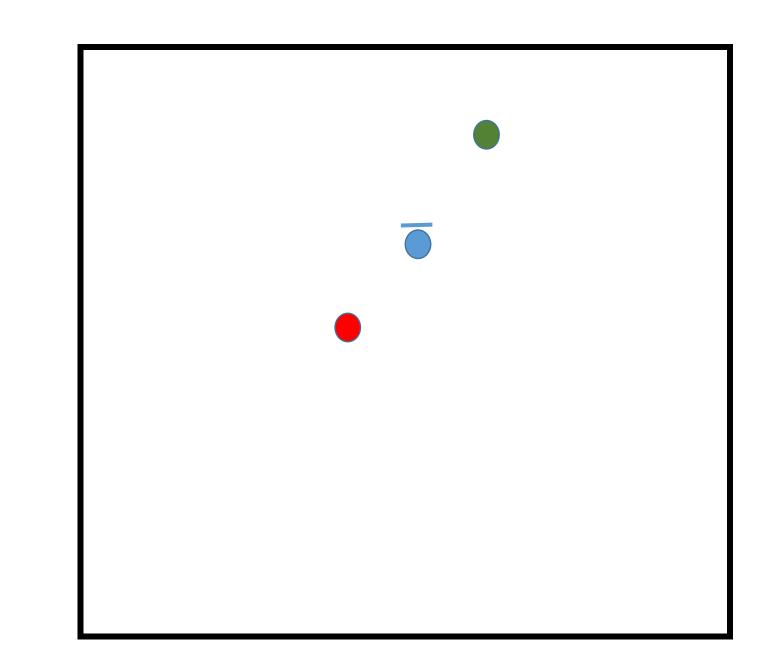
- Estimates variance components simultaneously to fixed terms
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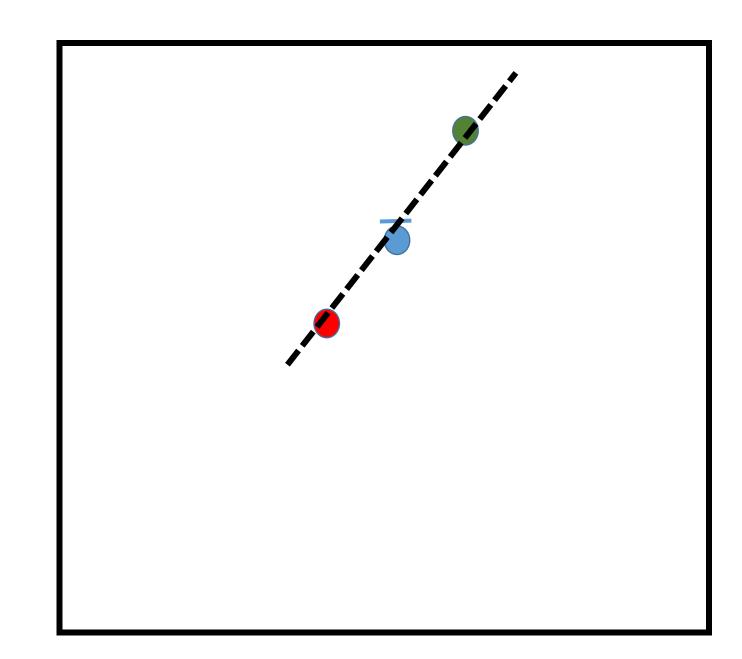
- Estimates variance components simultaneously to fixed terms
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- Allow to account for pseudoreplication

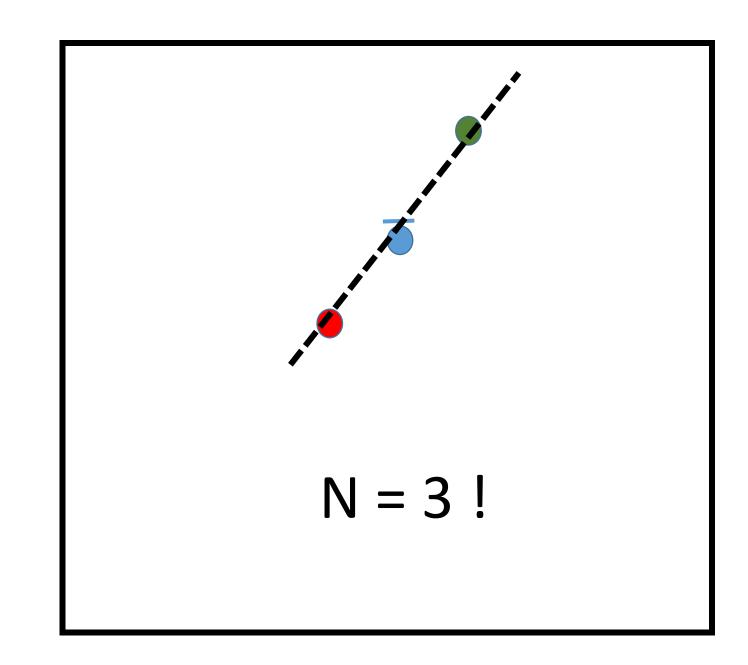


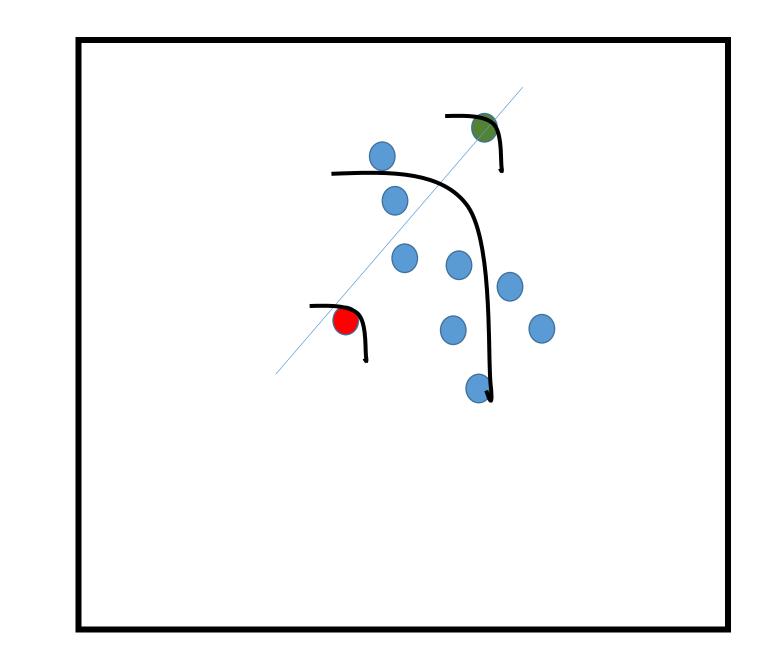












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 Get information about variance components (e.g. to inform about repeatability)

When is something random factor and when fixed?

- Rules of thumb:
- Random effects are factors!
- Are you interested in means (fixed) or variance (random)?
- Do you want to correct for a factorial effect but it's not in your questions specifically? -> random
- More than 5 levels: random
- LMM use only with large N (>50)

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- Run model with all fixed + random effects
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- Use likelihood ratio test or DIC

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2(logLmodel - logLreduced)**Df for chi square = 1**

```
> m0<-lm(y~1)
```

```
> m0<-lm(y~1)
> m1<-lm(y~x)
```

```
> m0<-lm(y~1)
> m1<-lm(y~x)
> lrtest(m0,m1)
```

```
> m0<-lm(y~1)
> m1<-lm(y~x)
> lrtest(m0,m1)
Likelihood ratio test
Model 1: y ~ 1
Model 2: y \sim x
 #Df LogLik Df Chisq Pr(>Chisq)
1 2 -17.488
2 3 -5.684 1 23.609 1.181e-06 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

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- Assume you want to know whether or not to include random effects
- Decide on fixed effects structure (best keep it to what's known)
- Add all the random effects
- Compare with reduced model (always only drop one!), use LRT
- OR
- When using Bayesian (MCMCglmm) use BIC (equiv. to AIC)

Are hard

- Are hard
- Not straightforward

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- But very useful
- Ask for help

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Mixed models - resources

- http://glmm.wikidot.com
- Gelman and Hill 2006 Data Analysis Using Regression and Multilevel/Hierarchical Models
- Zuur et al.: Mixed Effects Models and Extensions in Ecology 2009
- Bolker et al. Trends Ecol Evol 2009 Generalized linear mixed models: a practical guide for ecology and evolution

