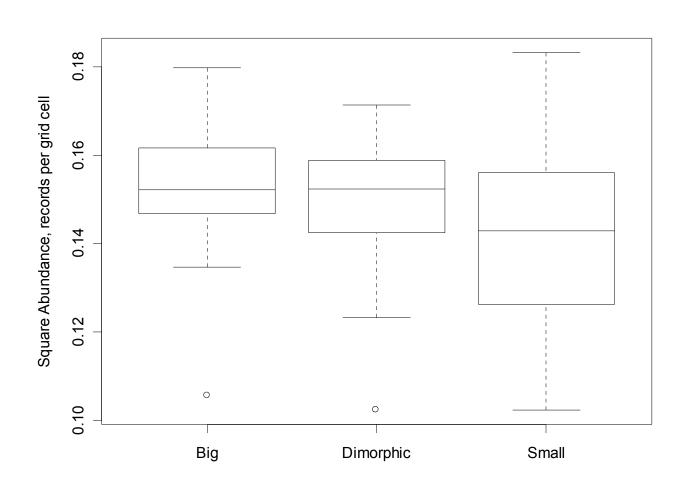
# Lecture 3: ANOVA and Regression I

Bob O'Hara

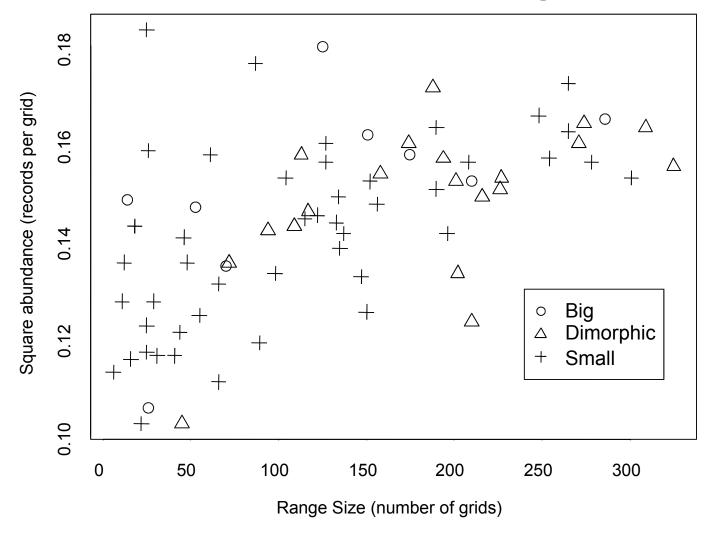
#### The Problem

- Is the abundance of a species related to
  - range size
  - wing type
- Data: from the 'many' habitat type
  - make it simpler
- Split the problem into small bits
  - easier to teach!

# The Data I Abundance of Wing Forms



# The Data II Abundance and Wing Form



### Simple Problem

- Does wing form affect abundance?
  - are their mean abundances different?
- One way ANOVA, 3 groups
- Generalise m groups, each with  $n_i$  observations

### Model

- Start with a model
  - Each observation *j*, from group *i* has a value:

$$-y_{ij} = \mu_i + \varepsilon_{ij} \qquad (j=1...n_i \text{ for } i=1, m)$$

- $-\mu_i$  is the group mean, **systematic** component
- $\varepsilon_{ij}$  is the error, the **random** component
- We can estimate the systematic part, but not perfectly because of the random part

# Different Ways of Coding

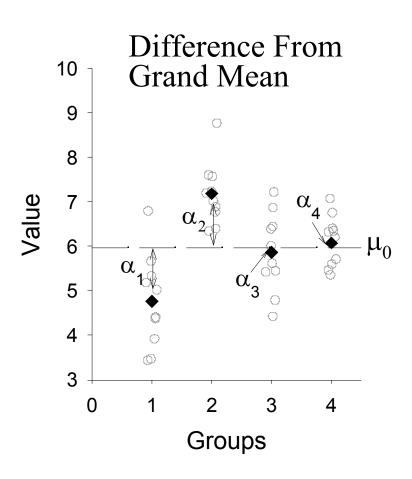
- At the moment, all  $\mu_i$ 's are different
  - no structure
- Can add it in several ways
  - Deviations from grand mean:

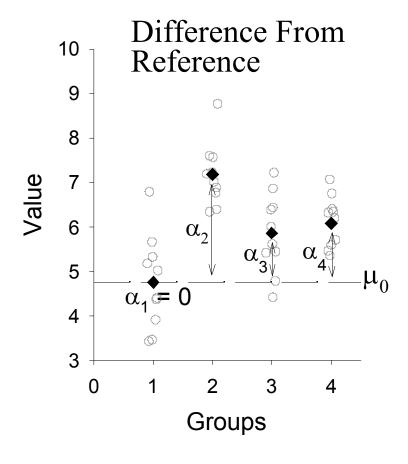
$$\mu_i = \mu_0 + a_i$$
 Constraint:  $\sum_{i=1}^{m} \alpha_i = 0$ 

Deviations from a reference

$$\mu_1 = \mu_0, \ \mu_i = \mu_0 + a_i$$
 (*i*=2...*m*)

# Coding in Pictures





### Estimates

- Assume the  $\varepsilon_{ij}$ 's are normally distributed with mean 0 and variance  $\sigma^2$ .
  - notation:  $\boldsymbol{\varepsilon}_{ij} \sim N(0, \boldsymbol{\sigma}^2)$
- Best estimates of  $\mu_i$ 's are their means  $\bar{y}_i$
- Best estimate of the variance  $\sigma^2$  is the sample variance,  $s^2$

$$s^{2} = \frac{1}{\sum_{j=1}^{m} n_{j} - m} \sum_{j=1}^{m} \sum_{i=1}^{n} (y_{ij} - \overline{y}_{j})^{2}$$

#### **Tests**

- Test whether  $\mu_i$ 's are the same
- Compare likelihoods
- Likelihoods are sums of squares

$$SS_{E} = \sum_{i=1}^{n} \sum_{j=1}^{m_{j}} y_{ij}^{2} - \sum_{i=1}^{n} \bar{y}_{i}^{2} SS_{E} = \sum_{i=1}^{m} n_{i} \bar{y}_{i}^{2} - m \bar{y}_{i}^{2}$$

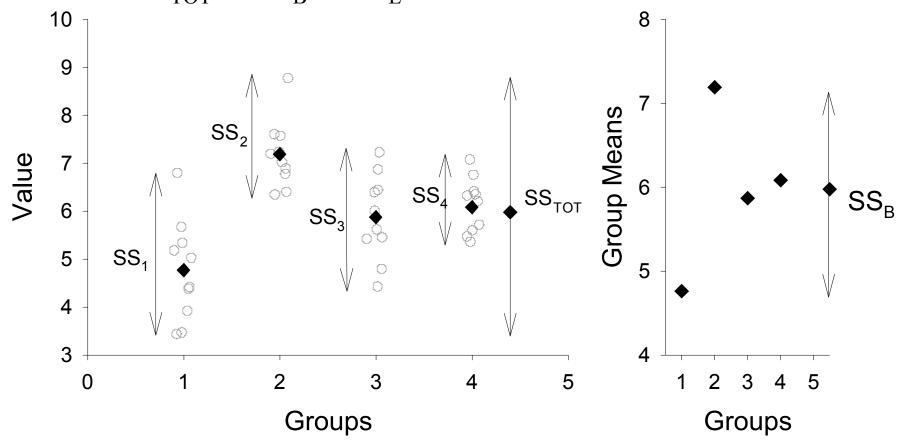
Summarise in an ANOVA table

### ANOVA Table

	df	SS	MS	F
Between	<i>m</i> -1	$SS_{B}$	SS <sub>B</sub> /m	$MS_B/MS_E$
Within (error)	n-m	$SS_{E}$	SS <sub>E</sub> /(n-m)	
Total	<i>n</i> -1	$SS_{T}$		

### What does this mean?

$$SS_{W} = SS_1 + SS_2 + SS_3 + SS_4$$
$$SS_{TOT} = SS_B + SS_E$$



### What it means II

• If the means are the same, then the differences in the means (as measured by  $SS_B$ ) can all be explained by the error (as measured by  $SS_W$ )

### Wing Size and Abundance

- 3 groups (Small, Dimorphic, Big)
- Are their abundances different?

### ANOVA Table

	df	SS	MS	F	Pr(F)
Wings	2	1.12	0.56	1.69	0.19
Residuals (error)	73	24.4	0.33		
Total	75	25.5	_		

(Sums of Squares multiplied by 1000)

# **Summary Statistics**

	Estimate	Std. Error	t value	$e \Pr(> t )$
(Intercept)	0.150	0.0061	24.6	<2e-16
Dimorphic	-0.0015	0.00734	-0.21	0.83
Small	-0.0089	0.00665	-1.34	0.18

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Residual standard error: 0.0183 on 73 degrees of freedom

- Level Big = Intercept, others give difference from this class
- *t*-tests are for difference between level and control level (in this case Big)

Mean for Small is 0.150 - 0.0089 = 0.159

Variance for all is 0.0183

### post hoc tests

- If there is a difference, where is it?
- Can use "orthogonal contrasts" for preplanned tests
  - e.g. control vs treatments
- Most of the time: test everything
  - use *t*-tests

#### One Test

- Compare Dimorphic and Small
- Estimates:

	Mean	Standard Error
Dimorphic	-0.0015	0.00734
Small	-0.0089	0.00665
Residual Mear	Square $(\sigma^2)$ :	0.33 (73 df)

• t-test:

$$t = \frac{x_1 - x_2}{\sqrt{\sigma^2 \left(\frac{1}{m_1} + \frac{1}{m_2}\right)}}$$

# Multiple Testing

- Do many tests at 5%
- If we do 20 tests where the null hypothesis is true, we would expect 1 to be rejected by chance
- We need to adjust the significance level
- Bonferroni: k tests at a%
  - change test level from a to a/k%
  - works OK if k not too large

# Significance Tests are Evil

- A *t* statistic increases by  $n^{1/2}$  as the sample size (n) increases
- For example: Correlation

$$n=10$$
,  $\rho=0.39$ ,  $p=0.26$   
 $n=30$ ,  $\rho=0.39$ ,  $p=0.033$ 

- With very large sample sizes, almost *p* values will be <5%, but the effect sizes could be tiny
- You have been warned....

# Simple Regression

- Similar to ANOVA
- Also use an ANOVA table to test hypotheses
- Estimate of slope and intercept:
  - intercept:  $\hat{\beta}_0 = \bar{y} \hat{\beta}_1 \bar{x}$
  - slope:  $\hat{\beta}_1 = \frac{Cov(x, y)}{Var(x)}$

### Regression Picture

$$SS_{TOT} = SS_{REG} + SS_{ERR}$$

120

100

80

 $SS_{ERR}$ 
 $SS_{REG}$ 
 $SS_{REG}$ 

### ANOVA Table

	df	SS	MS	F
Regression	1	SS <sub>REG</sub>	SS <sub>REG</sub> /1	MS <sub>REG</sub> /MS <sub>ERR</sub>
Error	<i>n</i> -2	SS <sub>ERR</sub>	$SS_{ERR}/(n-2)$	
Total	<i>n</i> -1	$SS_{TOT}$	_	

# Does Range Size Predict Abundance?

	df	SS	MS	F	Pr(>F)
Regression	1	8.20	8.20	35.09	<10 <sup>-7</sup>
Error	74	17.3	0.233		
Total	75	25.5 (S	ums of Squ	uares multip	lied by 1000)

### Range Size - Abundance II

	Estimate	Std. Error	t value Pr(> t )
Intercept	0.029	$3.14x10^{-3}$	$41.0 < 2 \times 10^{-16}$
Range Size	$1.18 \times 10^{-4}$	$1.99 \times 10^{-5}$	$5.92 < 10^{-7}$

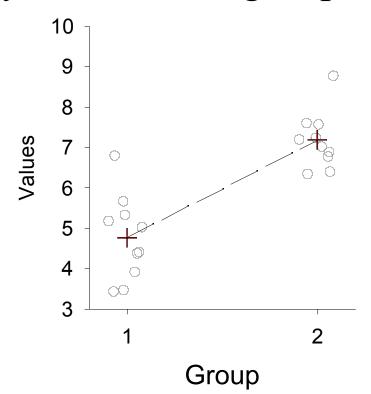
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Residual standard error: 0.0153 on 74 degrees of freedom Multiple R-Squared: 32%, Adjusted R-squared: 31%

- R<sup>2</sup>: Percent of variation explained by the regression
- Adjusted R<sup>2</sup>: adjusts for the number of degrees of freedom
  - easier to compare different sized models

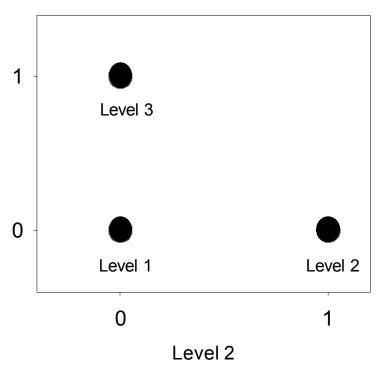
### Regression and ANOVA

- Actually the same
- One-way ANOVA, 2 groups:



### Regression = ANOVA

- One way ANOVA, 3 groups
  - Regression with 2 regressors
  - View Level 1 as a reference
  - Levels 2 and 3 change the mean from Level 1

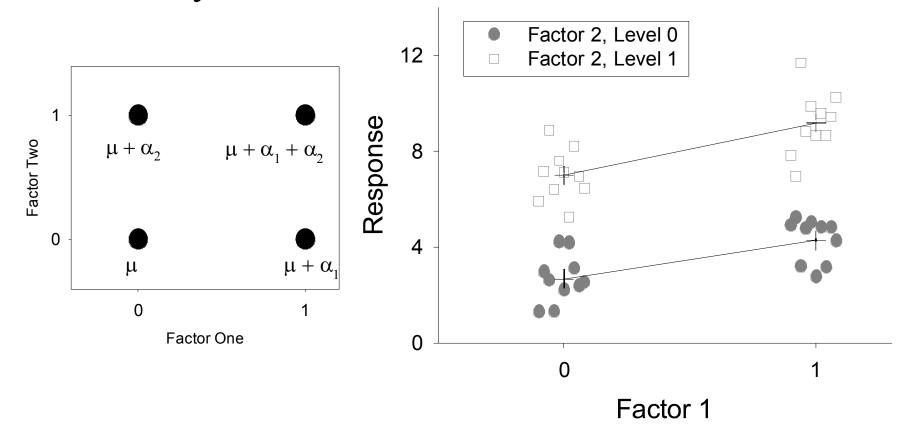


### ANOVA - more parameters

- In 1 way ANOVA, adding each group adds another regression
  - number of parameters builds up
- This is why replication is such an issue
- Makes design of experiments important

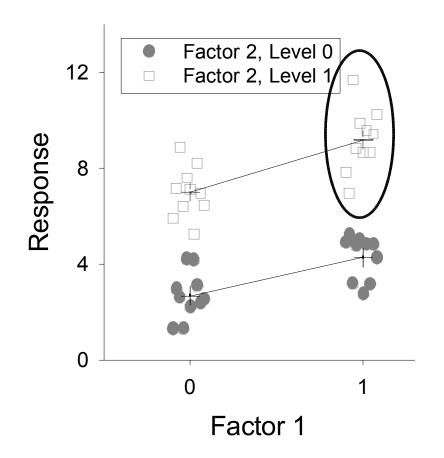
# Into something more difficult

• 2 way ANOVA: no interactions



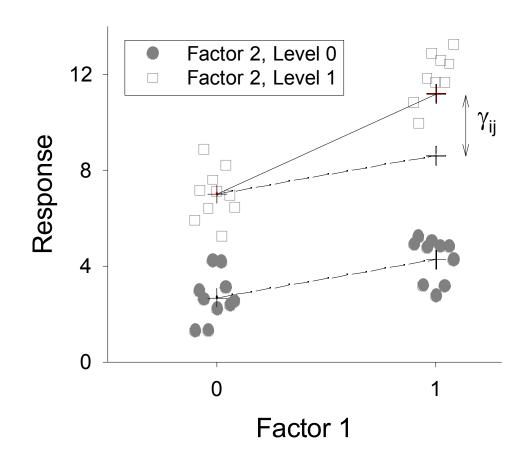
### Interactions

- Here the mean of the ringed group is determined by the 3 other parameters
  - $-\mu$ ,  $a_1$  and  $a_2$
- What if it is different?



### Interaction

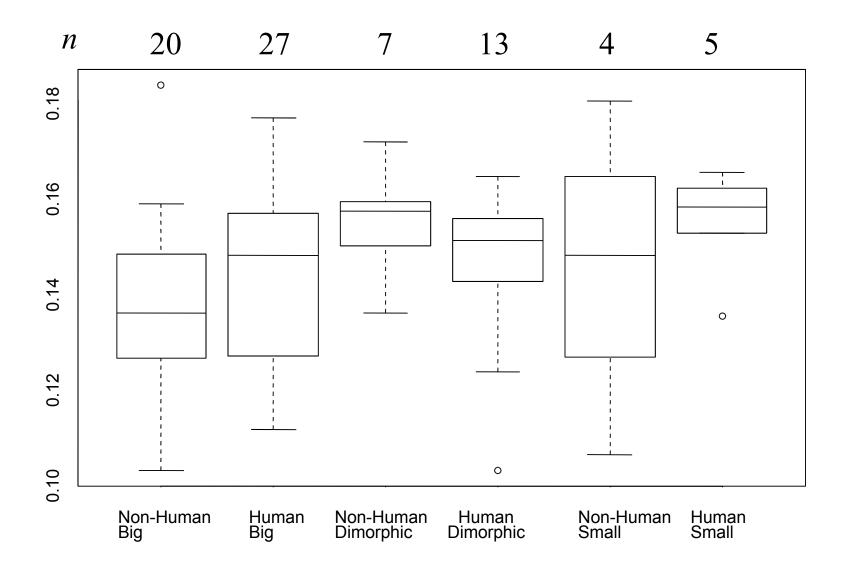
• 
$$y_{ijk} = \mu_0 + \alpha_1 + \alpha_2 + \gamma_{12} + \epsilon_{12k}$$



### Back to Beetles

- As well as wing form, we also know whether they like human habitats
- Does this have an effect?
- Does any effect vary with wing form?

# Box plots



The ANOVA
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	df	SS	MS	F	Pr(F)	
Wings	2	1.12	0.56	1.69	0.19	
Human	1	0.14	0.14	0.42	0.52	
Wings by Human	2	0.82	0.41	1.23	0.30	
Residuals	70	23.4	0.33			
Total	75	25.5	_			

The ANOVA - change order

The first ondings of def						
	df	SS	MS	F	Pr(F)	
Human	1	0.17	0.17	0.51	0.48	
Wings	2	1.09	0.54	1.63	0.20	
Wings by Human	2	0.82	0.41	1.23	0.30	
Residuals	70	23.4	0.33			
Total	75	25.5	_			

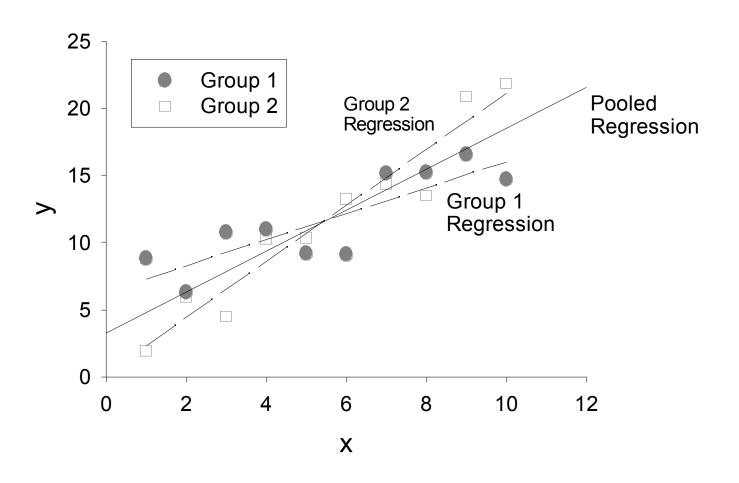
### Comments on the ANOVA

- The total sums of squares and total df will be the same for different models
  - total amount of variation in the model is the same
- Terms are added sequentially
  - Terms are really (Wings), (Human | Wings), (Wings by Human | Human & Wings)
- Order in which they are added makes a slight difference

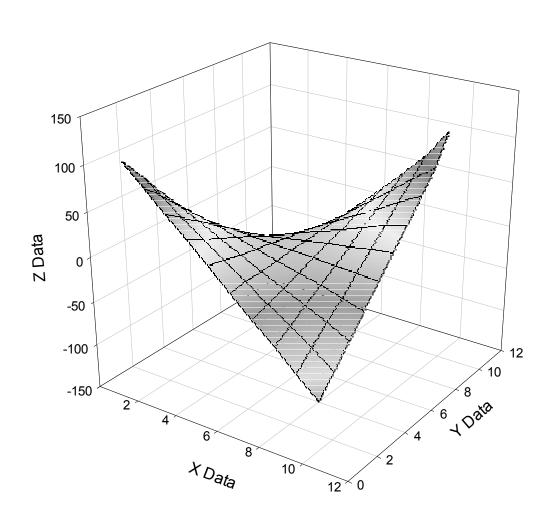
### Regression + ANOVA

- As they are both the same, can do together
- And then can add lots of other factors...
- What does an interaction mean?
  - slopes are different for different levels of a factor
- What would an interaction between 2 slopes mean?
  - the model would include  $\beta_1 x_1 + \beta_2 x_2 + \beta_{12} x_1 x_2$

### Different Slopes



# Regression Interaction



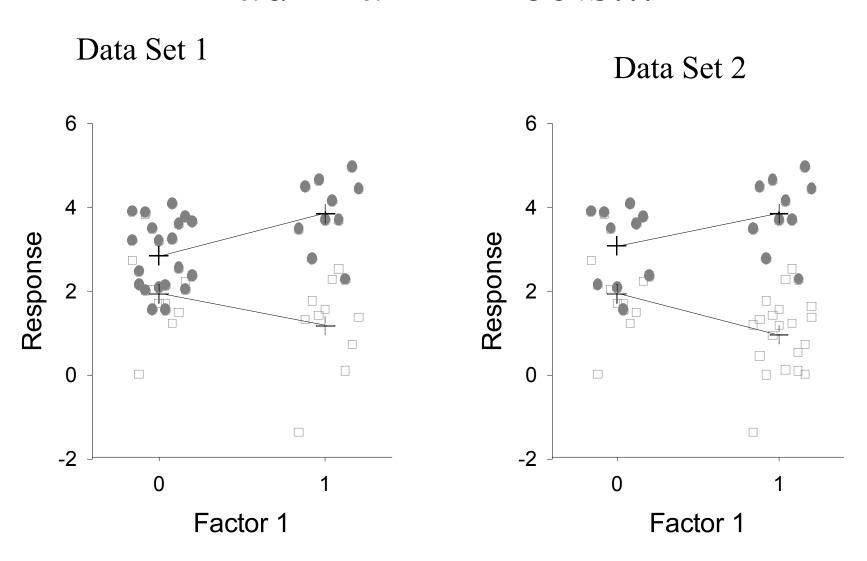
# Getting much more complicated

- As well as fitting linear terms for slopes, we can also fit polynomials
  - $-x^{2}, x^{3}, \text{ etc.}$
- Interactions are a natural extension
  - Model:  $(1+x_1)(1+x_2) = 1+x_1+x_2+x_1x_2+x_1^2+x_2^2$
- When fitting polynomial terms, keep lower order terms
  - unless you've got a good reason to drop them!

### Interactions

- The order they are added can be important for the ANOVA
  - Unless the experiment is balanced
- When adding interactions, keep main effects
  - unless you've got a good reason to drop them!
- In the presence of an interaction, the main effect is normally meaningless

### Bad Main Effects...



### ANOVA tables

		Data S	Data Set 1			Data Set2		
	df	SS	F	Pr(>F)	SS	F	Pr(>F)	
Factor2	1	31.37	35.8	$3x10^{-7}$	11.9	8.41	0.0057	
Factor1	1	0.75	0.85	0.36	7.17	5.05	0.029	
Factor1 by 2	1	8.93	10.2	0.0025	20.8	14.66	$3x10^{-4}$	
Error	46	40.26			65.30			

### With a significant interactions

- Main effects can be informative when
  - All changes are in the same direction
  - from observational studies, the overall direction may be interesting
- If the interaction can be considered a random effect
  - then change the ANOVA to test against the correct error term