

Introduction to Nested (hierarchical) ANOVA

Partitioning variance hierarchically

Two factor nested ANOVA

- Factor A with p groups or levels
 - fixed or random but usually fixed
- Factor B with q groups or levels within each level of A
 - usually random
- Nested design:
 - different (randomly chosen) levels of Factor B in each level of Factor A
 - often one or more levels of subsampling

Sea urchin grazing on reefs

- Andrew & Underwood (1997)
- Factor A - fixed
 - sea urchin density
 - four levels (0% original, 33%, 66%, 100%)
- Factor B - random
 - randomly chosen patches
 - four (3 to 4m²) within each treatment

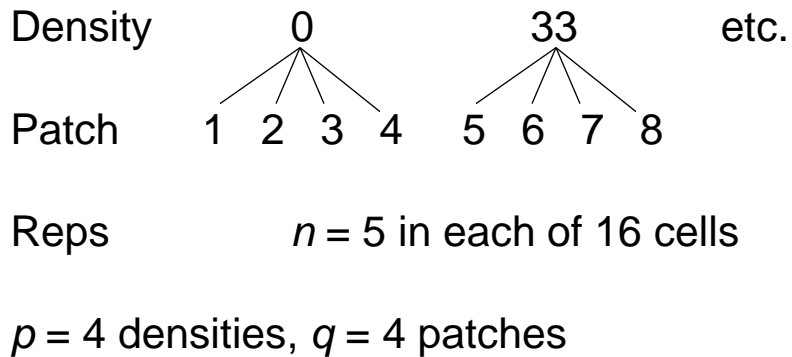


Sea urchin grazing on reefs

- Residual:
 - 5 replicate quadrats within each patch within each density level
- Response variable:
 - % cover of filamentous algae



Worked example



Data layout

| | | | | |
|-----------|----------------|----------------|-------|-----------------|
| Factor A | 1 | 2 | | i |
| A means | \bar{y}_1 | \bar{y}_2 | | \bar{y}_i |
| Factor B | 1... j4 | 5... j8 | | 9... j12 |
| B means | \bar{y}_{11} | | | \bar{y}_{ij} |
| ($q=4$) | | | | |
| Reps | y_{111} | | | y_{ij1} |
| | y_{112} | | | y_{ij2} |
| | ... | | | ... |
| | y_{11k} | | | y_{ijk} |

Linear model

$$y_{ijk} = \mu + \alpha_i + \beta_{j(i)} + \varepsilon_{ijk}$$

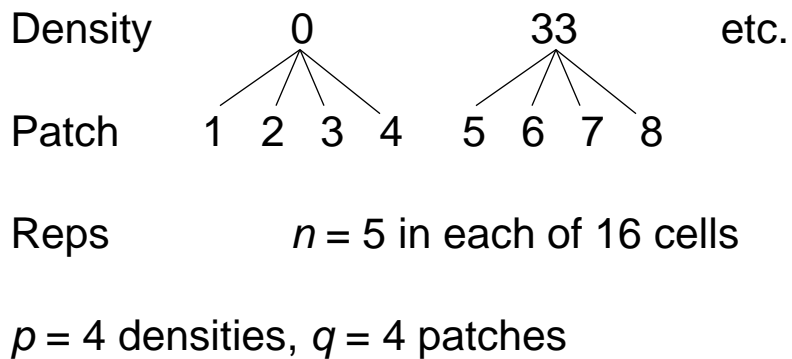
where

| | |
|---------------------|--|
| μ | overall mean |
| α_i | effect of factor A ($\mu_i - \mu$) |
| $\beta_{j(i)}$ | effect of factor B within each level of A ($\mu_{ij} - \mu_i$) |
| ε_{ijk} | unexplained variation (error term) - variation within each cell |

Linear model

$$(\% \text{ cover algae})_{ijk} = \mu + (\text{sea urchin density})_i + (\text{patch within sea urchin density})_{j(i)} + \varepsilon_{ijk}$$

Worked example



Effects

- Main effect:
 - effect of factor A
 - variation between factor A marginal means
- Nested (random) effect:
 - effect of factor B within each level of factor A
 - variation between factor B means within each level of A

Null hypotheses

- H_0 : no difference between means of factor A
 - $\mu_1 = \mu_2 = \dots = \mu_i = \mu$
- H_0 : no main effect of factor A:
 - $\alpha_1 = \alpha_2 = \dots = \alpha_i = 0$
 - $\alpha_i = (\mu_i - \mu) = 0$

Sea urchin example

- No difference between urchin density treatments
- No main effect of density

Null hypotheses

- H_0 : no difference between means of factor B within any level of factor A
 - $\mu_{11} = \mu_{12} = \dots = \mu_{1j}$
 - $\mu_{21} = \mu_{22} = \dots = \mu_{2j}$
 - etc.
- H_0 : no variance between levels of nested random factor B within any level of factor A:
 - $\sigma_\beta^2 = 0$

Sea urchin example

- No difference between mean filamentous algae cover for patches within any urchin density treatment
- No variance between patches within each density treatment

Residual variation

- Variation between replicates within each cell
- Pooled across cells if homogeneity of variance assumption holds

$$\sum (y_{ijk} - \bar{y}_{ij})^2$$

Partitioning total variation

$$SS_{\text{Total}} = SS_A + SS_{B(A)} + SS_{\text{Residual}}$$

SS_A

variation between A marginal means

$SS_{B(A)}$

variation between B means within each level of A

SS_{Residual}

variation between replicates within each cell

Nested ANOVA table

| Source | SS | df | MS |
|-------------|-----------------|-----------|---------------------------------|
| Factor A | SS_A | $p-1$ | $\frac{SS_A}{p-1}$ |
| Factor B(A) | $SS_{B(A)}$ | $p(q-1)$ | $\frac{SS_{B(A)}}{p(q-1)}$ |
| Residual | $SS_{Residual}$ | $pq(n-1)$ | $\frac{SS_{Residual}}{pq(n-1)}$ |

Expected mean squares

A fixed, B random:

- MS_A $\sigma^2 + n\sigma_\beta^2 + \frac{nq \sum \alpha_i^2}{p-1}$
- $MS_{B(A)}$ $\sigma^2 + n\sigma_\beta^2$
- $MS_{Residual}$ σ^2

Testing null hypotheses

- If no main effect of factor A:

- $H_0: \mu_1 = \mu_2 = \dots = \mu_i = \mu$ ($\alpha_i = 0$) is true

- $F\text{-ratio } MS_A / MS_{B(A)} \leq 1$

$$MS_A = \frac{\sigma^2 + n\sigma_\beta^2 + \frac{nq \sum \alpha_i^2}{p-1}}{p-1}$$

- If no nested effect of random factor B:

- $H_0: \sigma_\beta^2 = 0$ is true

- $F\text{-ratio } MS_{B(A)} / MS_{\text{Residual}} \leq 1$

$$MS_{B(A)} = \frac{\sigma^2 + n\sigma_\beta^2}{n}$$

$$MS_{\text{Residual}} = \sigma^2$$

Additional tests

- Main effect:

- planned contrasts & trend analyses as part of design

- unplanned multiple comparisons if main F -ratio test significant

- Nested effect:

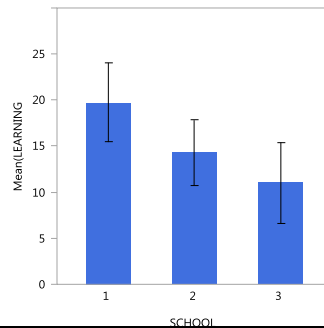
- usually random factor

- Sometimes little interest in further tests

- *Often can provide information on the characteristic spatial signal of a population*

Worked example

- What is the effect of schools on standardized tests?
- Is the effect of school driven in part by differences in teachers



Collected data (three schools, two teachers at each schools, two scores per teacher

| | Teacher 1 | Teacher 2 |
|----------|-----------|-----------|
| School 1 | 25 | 14 |
| | 29 | 11 |
| School 2 | 11 | 22 |
| | 6 | 18 |
| School 3 | 17 | 5 |
| | 20 | 2 |

True data matrix, accounts for teachers not being the same at each school

| | Teacher 1 | Teacher 2 | Teacher 3 | Teacher 4 | Teacher 5 | Teacher 6 |
|----------|-----------|-----------|-----------|-----------|-----------|-----------|
| School 1 | 25 | 14 | | | | |
| | 29 | 11 | | | | |
| School 2 | | | 11 | 22 | | |
| | | | 6 | 18 | | |
| School 3 | | | | | 17 | 5 |
| | | | | | 20 | 2 |

Data format for statistics

| School | Teacher | Score |
|--------|---------|-------|
| 1 | 1 | 25 |
| 1 | 1 | 29 |
| 1 | 2 | 14 |
| 1 | 2 | 11 |
| 2 | 3 | 11 |
| 2 | 3 | 6 |
| 2 | 4 | 22 |
| 2 | 4 | 18 |
| 3 | 5 | 17 |
| 3 | 5 | 20 |
| 3 | 6 | 5 |
| 3 | 6 | 2 |

No obvious effect of School

- After accounting for teacher effect

| Analysis of Variance | | | | |
|----------------------|----|----------------|-------------|--------------------|
| Source | DF | Sum of Squares | Mean Square | F Ratio |
| Model | 5 | 724.00000 | 144.800 | 20.6857 |
| Error | 6 | 42.00000 | 7.000 | Prob > F |
| C. Total | 11 | 766.00000 | | 0.0010 * |

| Variance Component Estimates | | |
|------------------------------|--------------|------------------|
| Component | Var Comp Est | Percent of Total |
| TEACHER[SCHOOL]&Rando | 91.08333 | 92.863 |
| Residual | 7 | 7.137 |
| Total | 98.08333 | 100.000 |

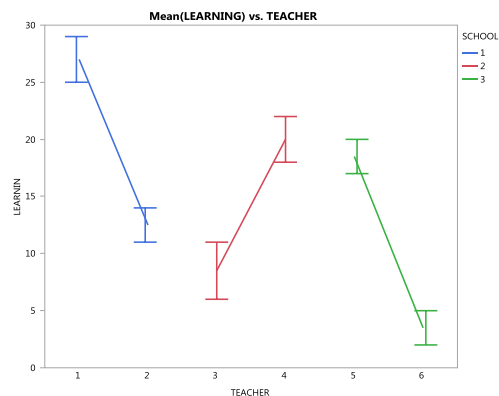
These estimates based on equating Mean Squares to Expected Value.

| Tests wrt Random Effects | | | | | |
|--------------------------|-------|---------|--------|---------|----------|
| Source | SS | MS Num | DF Num | F Ratio | Prob > F |
| SCHOOL | 156.5 | 78.25 | 2 | 0.4137 | 0.6940 |
| TEACHER[SCHOOL]&Rando | 567.5 | 189.167 | 3 | 27.0238 | 0.0007 * |

| Test Denominator Synthesis | | | | |
|----------------------------|---------|--------|-----------------------|--|
| Source | MS Den | DF Den | Denom MS Synthesis | |
| SCHOOL | 189.167 | 3 | TEACHER[SCHOOL]&Rando | |
| TEACHER[SCHOOL]&Rando | 7 | 6 | Residual | |

Big effect of teacher!!

- What about school effect
 - Test $MS_{\text{school}} / MS_{\text{teacher(school)}}$

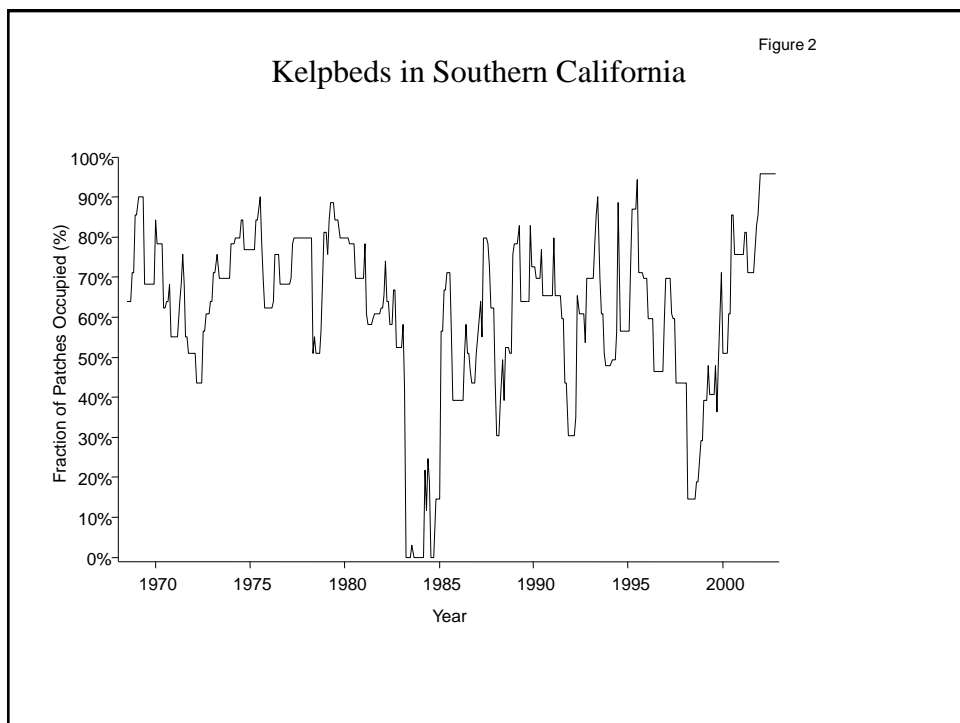


Spatially nested designs

- Used to provide information on the characteristic spatial signal of populations
- Other techniques (geostatistical models) also can do this but nested models are very efficient
- Variance component models (part of nested) can provide the percent of variation that is associated with particular spatial scales.

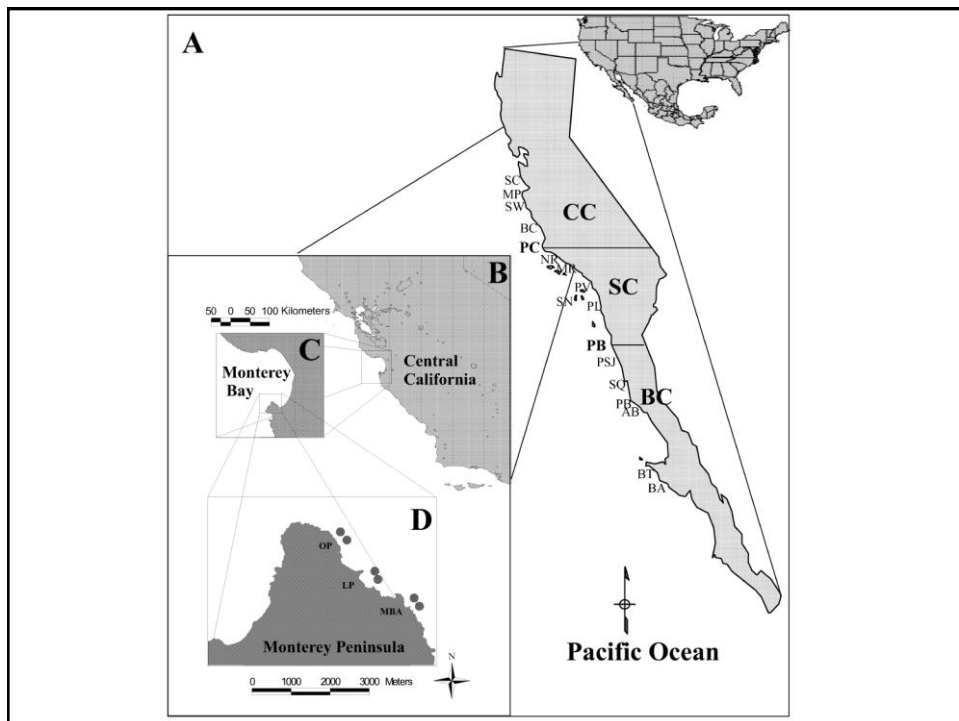
Kelp Forests





Matthew S. Edwards

Estimating scale-dependency in disturbance impacts: El Niños and giant kelp forests in the northeast Pacific



Scale: # Levels Spatial Inference

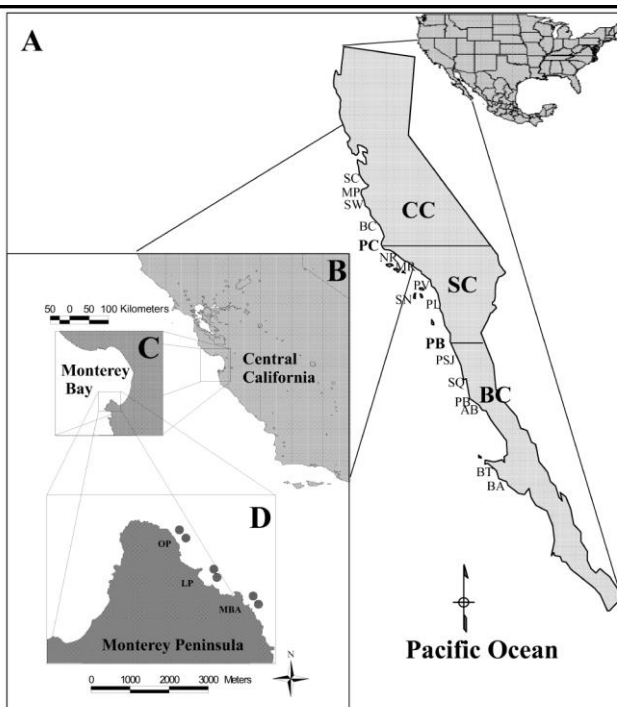
Region: 1 2 3 500 - 1500 km

Location: 1 2 3 4 5 10 - 100 km

Area: 1 2 3 1 - 5 km

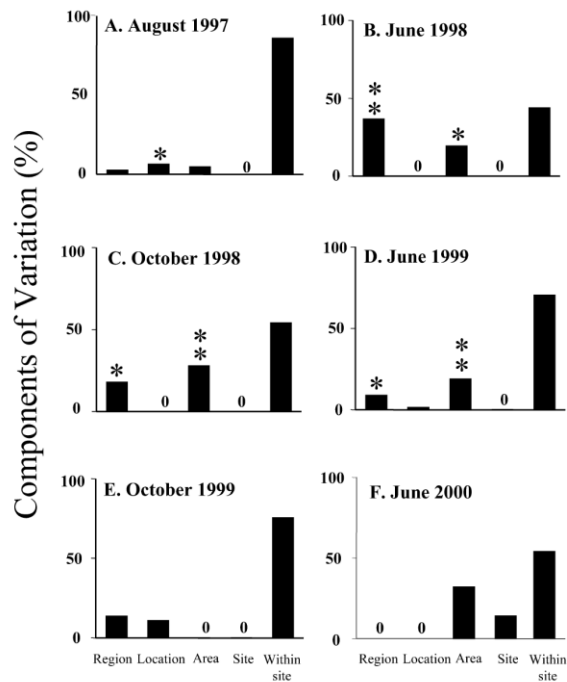
Site: 1 2 100 - 300 m

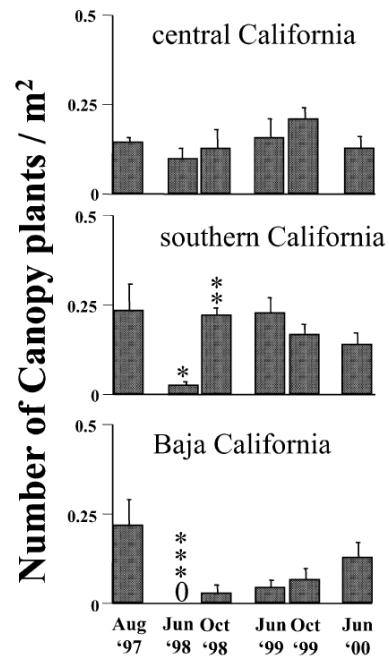
Transect: 1 2 3 < 40 m
(within site)



Complex nested designs

| Source | Expected mean square | Variance Component | F-ratio |
|-------------------------------------|--|---|---|
| Region: A | $\sigma_\epsilon^2 + n\sigma_\delta^2 + nk\sigma_\gamma^2 + nkr\sigma_\beta^2 + nkrq\sigma_\alpha^2$ | $\frac{MS_A - MS_{B(A)}}{nkrq}$ | $\frac{MS_A}{MS_{B(A)}}$ |
| Location: B(A) | $\sigma_\epsilon^2 + n\sigma_\delta^2 + nk\sigma_\gamma^2 + nkr\sigma_\beta^2$ | $\frac{MS_{B(A)} - MS_{C(B(A))}}{nkr}$ | $\frac{MS_{B(A)}}{MS_{C(B(A))}}$ |
| Area: (C(B(A))) | $\sigma_\epsilon^2 + n\sigma_\delta^2 + nk\sigma_\gamma^2$ | $\frac{MS_{C(B(A))} - MS_{D(C(B(A)))}}{nk}$ | $\frac{MS_{C(B(A))}}{MS_{D(C(B(A)))}}$ |
| Site: D(C(B(A))) | $\sigma_\epsilon^2 + n\sigma_\delta^2$ | $\frac{MS_{D(C(B(A)))} - MS_{Residual}}{n}$ | $\frac{MS_{D(C(B(A)))}}{MS_{Residual}}$ |
| Transect: Residual E(D(C(B(A)))) | σ_ϵ^2 | | |





Introduction to Repeated Measures designs

Two major types of repeated measures ANOVA

- Subjects used repeatedly but performance is unlikely to be linked to order (timing)
 - Same subjects used for a series of treatments, treatment order randomized among subjects
- Subjects used repeatedly and performance is likely to be linked to order (timing)
 - Performance = growth, size, etc

Subjects used repeatedly but performance is unlikely to be linked to order (timing)

- Example: the effect of four types of drugs on blood pressure compared between men and women
 - Gender is fixed effect (consider between subject effect)
 - Each subject (within a gender) receives all four drugs (within subject effects)
 - Drug order is:
 - Random and
 - Separation between drugs is assumed to be long enough that there are no carryover effects

Subjects used repeatedly and performance is likely to be linked to order (timing)

- Example: effect of 4 hormones on individual size of fish. Measurements taken repeatedly over time
 - Hormone effect is ‘between subject’ effect
 - Time and Time*Hormone levels are ‘within subject’ effects
 - Separate error terms for between and within subject effects
 - Between subject effects are estimated using (eq.) of means of all temporal measurements (one estimate per individual)
 - Within subject effects are estimated using all measurements (temporal replicates within individuals are used)

Repeated measures designs

- Each whole plot (or individual) measured repeatedly under different treatments and/or times
- Within plots (individual) factor often time, or at least treatments applied through time
- Plots (individuals) termed “subjects” in repeated measures terminology

Cane toad breathing and hypoxia

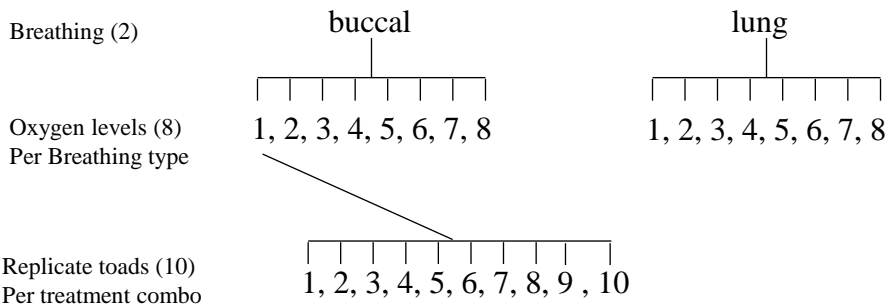


Cane toad breathing and hypoxia

- How do cane toads respond to conditions of hypoxia?
 - Mullens (1993)
- Two factors:
 - Breathing type (buccal vs lung breathers)
 - O_2 concentration (8 different $[O_2]$)
- 10 replicates per breathing type and $[O_2]$ combination
- Response variable is breathing rate

Completely randomized design

- 2 factor design (2 x 8) with 10 replicates
 - total number of toads = 160
- Toads are expensive
 - reduce number of toads?
- Lots of variation between individual toads
 - reduce between toad variation?



160 reps in a completely randomized design

Repeated measures designs

- Factor A:
 - units of replication termed “subjects”
- Factor B (subjects) nested within A
- Factor C:
 - repeated recordings on each subject

Repeated measures design

| Breathing | Toad | [O ₂] | | | | | | | |
|-----------|------|-------------------|-----|-----|-----|-----|-----|-----|-----|
| type | | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |
| Lung | 1 | x | x | x | x | x | x | x | x |
| Lung | 2 | x | x | x | x | x | x | x | x |
| ... | ... | ... | ... | ... | ... | ... | ... | ... | ... |
| Lung | 9 | x | x | x | x | x | x | x | x |
| Buccal | 10 | x | x | x | x | x | x | x | x |
| Buccal | 12 | x | x | x | x | x | x | x | x |
| ... | ... | ... | ... | ... | ... | ... | ... | ... | ... |
| Buccal | 21 | x | x | x | x | x | x | x | x |

Repeated measures design

- Factor A is breathing type:
 - lung vs buccal
 - applied to toads = subjects = plots
- Factor B is subjects (i.e. toads) nested within A
- Factor C is [O₂] treatment
 - 0, 5, 10, 15, 20, 30, 40, 50%
 - applied to toads (subjects) repeatedly

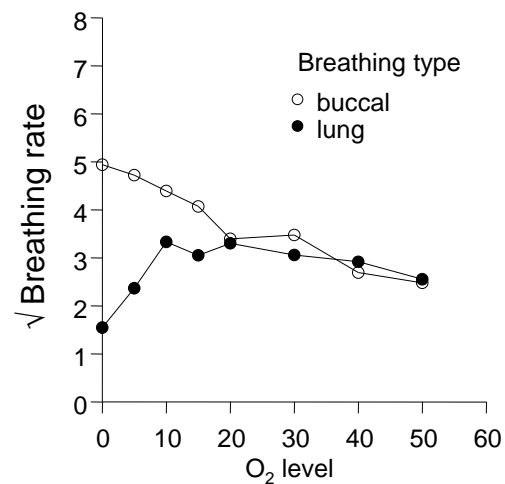
ANOVA

| Source of variation | df |
|--|-----|
| <i>Between subjects (toads)</i> | |
| Breathing type | 1 |
| Toads within breathing type (Residual 1) | 19 |
| <i>Within subjects (toads)</i> | |
| [O ₂] | 7 |
| Breathing type x [O ₂] | 7 |
| Toads (Breathing type) x [O ₂] | |
| (Residual 2) | 133 |
| Total | 167 |

ANOVA toad example

| Source of variation | df | MS | F | P |
|--|-----|-------|-------|--------|
| <i>Between subjects (toads)</i> | | | | |
| Breathing type | 1 | 39.92 | 5.76 | 0.027 |
| Toads (breathing type) | 19 | 6.93 | | |
| <i>Within subjects (toads)</i> | | | | |
| [O ₂] | 7 | 3.68 | 4.88 | <0.001 |
| Breathing type x [O ₂] | 7 | 8.05 | 10.69 | <0.001 |
| Toads (Breathing type) x [O ₂] | 133 | 0.75 | | |
| Total | 167 | | | |

Mullens (1993)



Gange (1995)

- Factor A is tree species:
 - alder 1 vs alder 2
 - applied to trees = subjects = plots
- Factor B is subjects (i.e. trees) nested within A
- Factor C is date
 - 20 dates between May and September
 - recorded from trees (subjects) repeatedly
- Response variable is aphid abundance



Assumptions

- Normality & homogeneity of variance:
 - affects between-plots (between-subjects) tests
 - boxplots, residual plots, variance vs mean plots etc. for average of within-plot (within-subjects) levels

- No “carryover” effects:
 - results on one subplot do not influence results on another subplot.
 - time gap between successive repeated measurements long enough to allow recovery of “subject”

Sphericity

- Sphericity of variance-covariance matrix
 - variances of paired differences between levels of within-plots (or subjects) factor equal within and between levels of between-plots (or subjects) factor
 - variance of differences between $[O_2] 1$ and $[O_2] 2$ = variance of differences between $[O_2] 1$ and $[O_2] 3$ etc.

Sphericity assumption

Toad $O_21 - O_22$ $O_22 - O_23$ $O_21 - O_23$ etc.

| | | | |
|------|-------------------|-------------------|-------------------|
| 1 | $y_{11} - y_{21}$ | $y_{21} - y_{31}$ | $y_{11} - y_{31}$ |
| 2 | $y_{12} - y_{22}$ | $y_{22} - y_{32}$ | $y_{12} - y_{32}$ |
| 3 | $y_{13} - y_{23}$ | $y_{23} - y_{33}$ | $y_{13} - y_{33}$ |
| etc. | | | |

$$\text{Var}(\text{diff}(1-2)) = \text{Var}(\text{diff}(2-3)) = \text{Var}(\text{diff}(1-3))$$

Sphericity (compound symmetry)

- OK for split-plot designs
 - within plot treatment levels randomly allocated to subplots
- OK for repeated measures designs
 - if order of within subjects factor levels randomised
- Not OK for repeated measures designs when within subjects factor is time
 - order of time cannot be randomised

ANOVA options

- Standard univariate partly nested analysis
 - only valid if sphericity assumption is met
 - OK for some repeated measures designs (those where performance is not assumed to change with time)

ANOVA options

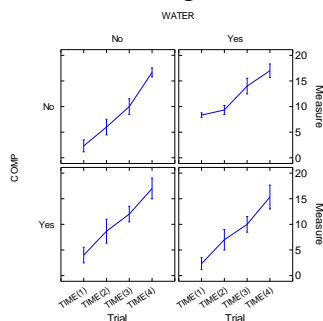
- Adjusted univariate F -tests for within-subjects factors and their interactions
 - conservative tests when sphericity is not met
 - Greenhouse-Geisser better than Huynh-Feldt

ANOVA options

- Multivariate (MANOVA) tests for within subjects or plots factors
 - responses from each subject used in MANOVA
 - doesn't require sphericity
 - sometimes more powerful than GG adjusted univariate, sometimes not

Subjects used repeatedly and performance is likely to be linked to order (timing)

- Effect of Competition (2 levels), Watering (2 levels) and Time (4 levels) on growth of Oak Seedlings
- 3 replicate seedlings for each combination of competition and watering. Each seedling followed over time



Repeated measures seedling growth as a function of competition and watering

Completely randomized

Competition (2) Watering (2) Time (4) Replicate (3)

Yes
No

Yes
No

1
2
3

1
2
3

1
2
3

1
2
3

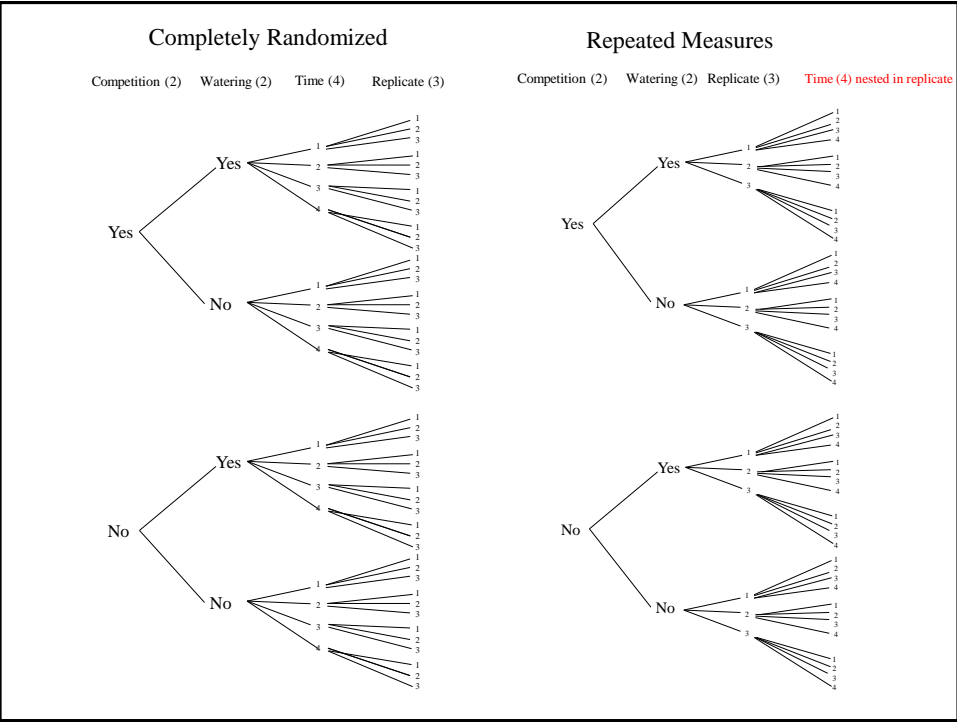
Replication
 $2 \times 2 \times 4 \times 3 = 48$
Each sampled 1 time

Repeated Measures

Competition (2) Watering (2) Replicate (3) Time (4) – note time is nested in replicate

```
graph LR; A[Yes] --> B[Yes]; A --> C[No]; B --> D[1]; B --> E[2]; B --> F[3]; D --> G[1]; D --> H[2]; D --> I[3]; D --> J[4]; E --> K[1]; E --> L[2]; E --> M[3]; E --> N[4]; F --> O[1]; F --> P[2]; F --> Q[3]; F --> R[4]; C --> S[1]; C --> T[2]; C --> U[3]; S --> V[1]; S --> W[2]; S --> X[3]; S --> Y[4]; T --> Z[1]; T --> AA[2]; T --> AB[3]; T --> AC[4]; U --> AD[1]; U --> AE[2]; U --> AF[3]; U --> AG[4]
```

Replication
 $2 \times 2 \times 3 = 12$
Each sampled 4 times



Between subject effects (No time effects)

| COMP\$ | | | | | |
|--------|-----------|---------|-------|-------|--------|
| Test | Value | Exact F | NumDF | DenDF | Prob>F |
| F Tes | 0.1222222 | 0.9778 | 1 | 8 | 0.3517 |

| WATER\$ | | | | | |
|---------|-----------|---------|-------|-------|--------|
| Test | Value | Exact F | NumDF | DenDF | Prob>F |
| F Tes | 0.1010101 | 0.8081 | 1 | 8 | 0.3949 |

| COMP\$*WATER\$ | | | | | |
|----------------|-----------|---------|-------|-------|----------|
| Test | Value | Exact F | NumDF | DenDF | Prob>F |
| F Tes | 0.9707071 | 7.7657 | 1 | 8 | 0.0237 * |

Note degrees of freedom – the denominator DF does note include temporal replication

Within subject effects (includes time effects)

| Time | | | | | |
|----------------------|-----------|----------|--------|--------|----------|
| Test | Value | Exact F | NumDF | DenDF | Prob>F |
| F Test | 63.843216 | 127.6864 | 3 | 6 | <.0001 * |
| Univar unadj Epsilon | 1 | 152.0511 | 3 | 24 | <.0001 * |
| Univar G-G Epsilon | 0.5361496 | 152.0511 | 1.6084 | 12.868 | <.0001 * |
| Univar H-F Epsilon= | 0.9023049 | 152.0511 | 2.7069 | 21.655 | <.0001 * |

| Time*COMP\$ | | | | | |
|----------------------|-----------|---------|--------|--------|----------|
| Test | Value | Exact F | NumDF | DenDF | Prob>F |
| F Test | 3.0913235 | 6.1826 | 3 | 6 | 0.0289 * |
| Univar unadj Epsilon | 1 | 1.2907 | 3 | 24 | 0.3003 |
| Univar G-G Epsilon | 0.5361496 | 1.2907 | 1.6084 | 12.868 | 0.3002 |
| Univar H-F Epsilon= | 0.9023049 | 1.2907 | 2.7069 | 21.655 | 0.3015 |

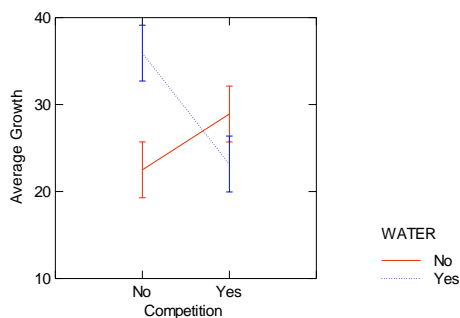
| Time*WATER\$ | | | | | |
|----------------------|-----------|---------|--------|--------|--------|
| Test | Value | Exact F | NumDF | DenDF | Prob>F |
| F Test | 1.7731767 | 3.5464 | 3 | 6 | 0.0875 |
| Univar unadj Epsilon | 1 | 1.8658 | 3 | 24 | 0.1624 |
| Univar G-G Epsilon | 0.5361496 | 1.8658 | 1.6084 | 12.868 | 0.1967 |
| Univar H-F Epsilon= | 0.9023049 | 1.8658 | 2.7069 | 21.655 | 0.1693 |

| Time*COMP\$*WATER\$ | | | | | |
|----------------------|-----------|---------|--------|--------|--------|
| Test | Value | Exact F | NumDF | DenDF | Prob>F |
| F Test | 2.0496341 | 4.0993 | 3 | 6 | 0.0669 |
| Univar unadj Epsilon | 1 | 1.9553 | 3 | 24 | 0.1477 |
| Univar G-G Epsilon | 0.5361496 | 1.9553 | 1.6084 | 12.868 | 0.1847 |
| Univar H-F Epsilon= | 0.9023049 | 1.9553 | 2.7069 | 21.655 | 0.1550 |

Note:

- 1) F-Test is Pillai Trace multivariate F test
- 2) Univar unadj – is regular F-test but subject to sphericity violations
- 3) Univar G-G and H-F are corrected univariate tests (account for sphericity)
- 4) Degrees of freedom for univar tests use temporal replication
- 5) G-G and H-F use adjusted degrees of freedom

Interaction between
Competition and Watering



Effect of Time on Growth
varies as a function of
competition

