Odds and ends

Observational vs. experimental key is whether researcher assigns treatment or not

Goals experiment

- 1) Eliminating bias
- 2) Reduce sampling error

Reduce bias

- Controls
- Random assignment to treatments
- Blinding

Reducing sampling error

$$t = \frac{\overline{Y_1} - \overline{Y_2}}{\sqrt{s_p^2 \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}}$$
 "Noise"

Can be achieved with smaller s or larger n.

Reducing sampling error

- Replication
- Balance (equal sample size)
- Blocking
- Extreme treatments

$$SE_{\bar{Y}_1 - \bar{Y}_2} = \sqrt{s_p^2 \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}.$$

Comparing the means of more than two groups

Analysis of variance (ANOVA)

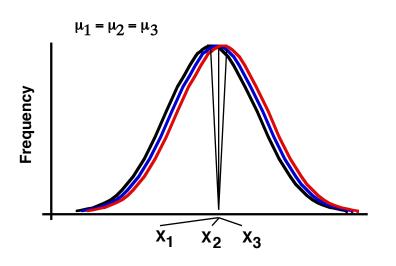
- Like a t-test, but can compare more than two groups
- Asks whether any of two or more means is different from any other.
- In other words, is the variance among groups greater than zero?

Null hypothesis for simple ANOVA

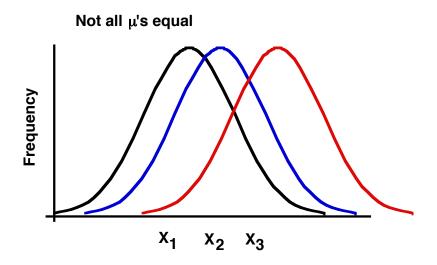
H₀: Variance among groups = 0

OR

• H_0 : $\mu_1 = \mu_2 = \mu_3 = \mu_4 = \dots \mu_k$

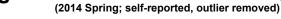


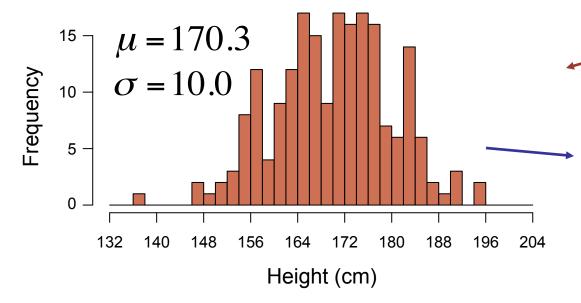
H₀: all populations have equal means



H_A: at least one population mean is different.

Heights of BIOL300 students (N = 202)

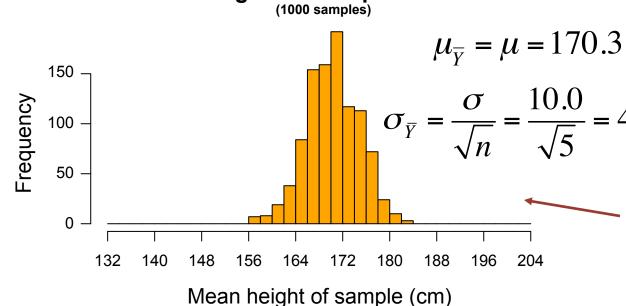




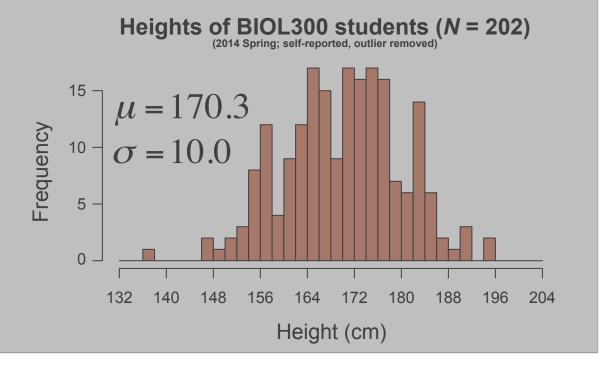
Sampling distribution of Y

If we draw multiple samples from the same population, we are also drawing sample means from a distribution.

Mean heights of samples of size 5



Sampling distribution of Y

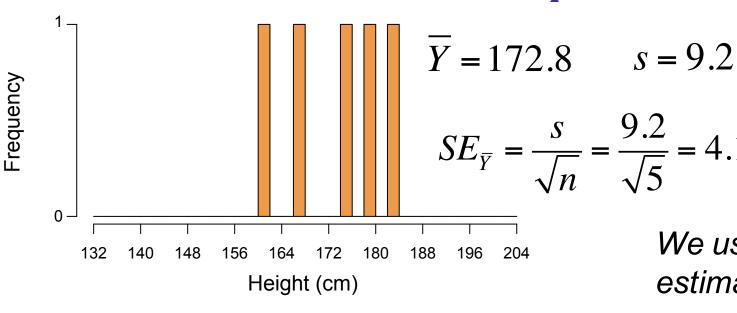


In most cases, we don't know the real population distribution.

We only have a

✓ sample.

Heights of a sample of students
$$(n = 5)$$



We use this as an estimate of
$$\sigma_{\overline{v}}$$

Under the null hypothesis, the sample mean of each group should vary because of sampling error.

The standard deviation of sample means, when the true mean is constant, is the standard error:

$$\sigma_{\bar{x}} = \frac{\sigma_{x}}{\sqrt{n}}$$

Note that we used standard errors in t-tests, e.g.: $t = \frac{Y_1 - Y_2}{SE_{\bar{Y}_1 - \bar{Y}_2}}$

Squaring the standard error, the variance among sample means due to sampling error should be:

$$\sigma_{\bar{x}}^2 = \frac{\sigma_x^2}{n}$$

In ANOVA, we work with variances rather than standard deviations.

Alternatively, if the null is FALSE:

$$\sigma_{\bar{x}}^2 = \frac{\sigma_x^2}{n} + Variance[\mu_i]$$

If the null hypothesis is <u>not</u> true, the variance among sample means should be equal to the variance due to sampling error *plus* the real variance among population means.

With ANOVA, we test whether the variance among true group means is greater than zero.

$$Variance[\mu_i] > 0?$$

We do this by asking whether the variance among sample means is greater than expected under the null hypothesis:

$$\sigma_{\bar{x}}^2 > \frac{\sigma_x^2}{n}?$$

But, our data is affected by sampling error, such that we only have estimates of the above.

$$\sigma_{\bar{x}}^{2} > \frac{\sigma_{x}^{2}}{n}?$$

$$n\sigma_{\bar{x}}^{2} > \sigma_{x}^{2}?$$

Population parameters

Estimates from sample

$$n\sigma_{\bar{x}}^2$$

 $n\sigma_{\overline{x}}^2$ is estimated by the "Mean Squares Group" MS_{group}

'n times variance among groups'

$$\sigma_x^2$$

is the variance within groups, O_{v}^{2} estimated by the "Mean Squares Error"

'variance within groups'

Mean squares group

Abbreviation: MS_{group}

Variance in mean due to sampling error

Estimates this parameter: $n\left(\frac{\sigma_x^2}{n} + Variance[\mu_i]\right)$

Variance in mean due to differences in means

Formula:
$$MS_{groups} = \frac{33_{groups}}{df_{groups}}$$

Mean squares group

$$SS_{group} = \sum n_i (\overline{X}_i - \overline{X})^2$$

i = groupj = indiv.

 \overline{X}_i is the mean of group i, and

$$\overline{X} = \sum_{i} \sum_{j} X_{ij} / N$$
 is the overall mean.

$$df_{\text{groups}} = k-1$$
 (where

(where k is the number of groups)

$$MS_{groups} = \frac{SS_{groups}}{df_{groups}}$$

Mean squares error

Abbreviation:

$$MS_{error}$$

Estimates this parameter:

$$\sigma_x^2$$

Formula:
$$MS_{error} = \frac{SS_{error}}{df_{error}}$$

Mean squares error

Error sum of squares =

$$SS_{error} = \sum df_i s_i^2 = \sum s_i^2 (n_i - 1)$$

Error degrees of freedom =

$$df_{error} = \sum df_i = \sum (n_i - 1) = N - k$$

$$MS_{error} = \frac{SS_{error}}{df_{error}} = \frac{\sum s_i^2(n_i - 1)}{N - k}$$

MS_{error} is like the pooled variance in a 2-sample *t*-test:

$$s_p^2 = \frac{df_1 s_1^2 + df_2 s_2^2}{df_1 + df_2}$$

Test statistic: F

If
$$H_0$$
 is true, then $n \sigma_{\bar{x}}^2 = \sigma_x^2$

In other words:
$$F = \frac{n \sigma_{\overline{x}}^2}{\sigma_x^2} = 1$$

But, the above refer to population parameters. We must estimate F from samples with: $MS_{\text{group}} / MS_{\text{error}}$

F if null hypothesis is false:

We test whether the F ratio is greater than one, as it would be if H_0 is false:

$$F = \frac{n\left(\frac{\sigma_x^2}{n} + Variance[\mu_i]\right)}{\sigma_x^2} > 1$$

But we must take into account sampling error. Often, F calculated from data will be greater than one even when the null is true. Hence we must compare F to a null distribution.

ANOVA table

Source	SS	df	MS	F	P
Group					
Error					
Total					

An ANOVA table is a convenient way to keep track of the important calculations.

Scientific papers often report ANOVA results with ANOVA tables.

Example: Body temperature of squirrels in low, medium and hot environments



Wooden & Walsberg (2004) **Body temperature and locomotor capacity in a heterothermic rodent.** *Journal of Experimental Biology* 207:41-46.

Squirrel body temperature data (° C)

Cold: 30.4, 31.0, 31.2, 31.0, 31.5, 30.4, 30.6, 31.1, 31.3, 31.9, 31.4, 31.6, 31.5, 31.4, 30.3, 30.5, 30.3, 30.0, 30.8, 31.0

Warm: 36.3, 37.5, 36.9, 37.2, 37.5, 37.7, 37.5, 37.7, 38.0, 38.0, 37.6, 37.4, 37.9, 37.2, 36.3, 36.2, 36.4, 36.7, 36.8, 37.0, 37.7

Hot: 40.7, 40.6, 40.9, 41.1, 41.5, 40.8, 40.5, 41.0, 41.3, 41.5, 41.3, 41.2, 40.7, 40.3, 40.2, 41.3, 40.7, 41.6, 41.5, 40.5

Hypotheses

H₀: Mean body temperature is the same for all three groups of squirrels.

H_A: At least one of the three is different from the others.

Summary data

Group	$\overline{\mathcal{X}}$	S	n	
Cold	31.0	0.551	20	
Warm	37.2	0.582	21	
Hot	41.0	0.430	20	

Total sample size: $N = \sum n = 20 + 21 = 20 = 61$

Error mean square for squirrels

$$SS_{error} = \sum df_i s_i^2$$

= $19(0.551)^2 + 20(0.582)^2 + 19(0.430)^2$
= 16.1

$$df_{error} = 19 + 20 + 19 = 58$$

$$MS_{error} = \frac{16.1}{58} = 0.277$$

Squirrel mean squares group:

$$\overline{X} = \frac{20(31.0) + 21(37.2) + 20(41.0)}{20 + 21 + 20} = 36.4$$

$$SS_{group} = \sum n_i (\overline{X}_i - \overline{X})^2$$

$$SS_{group} = 20(31.0-36.4)^2 + 21(37.2-36.4)^2 + 20(41.0-36.4)^2$$

= 1015.7

Squirrel mean squares group:

$$df_{group} = k - 1 = 3-1=2$$

$$MS_{groups} = \frac{SS_{groups}}{df_{groups}} = \frac{1015.7}{2} = 507.9$$

The test statistic for ANOVA is *F*

$$F = \frac{MS_{group}}{MS_{error}} = \frac{507.9}{0.277} = 1834.7$$

 MS_{group} is always in the numerator; MS_{error} is always in the denominator.

Compare to $F_{\alpha(1),df_group,df_error}$

$$F_{0.05(1),2,58} = 3.15.$$

Since 1835 > 3.15, we know P < 0.05 and we can reject the null hypothesis.

The variance in sample group means is bigger than expected given the variance within sample groups.

Therefore, at least one of the groups has a population mean different from another group.

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Source	SS	df	MS	F	P
Group	1015.7	2	507.9	1834.7	<0.0001
Error	16.1	58	0.277		
Total	1031.8	60			

Variability explained: R²

• R² is the fraction of variation that is explained by group differences:

$$R^2 = \frac{SS_{groups}}{SS_{total}}$$

For Squirrel study:

$$R^2 = 1015.7 / 1031.8 = 0.98$$

Assumptions of ANOVA

- Random samples.
- Normal distributions for each population.
 (Small deviations are OK)
- Equal variances for all populations.
 (Homoscedasticity)
 (But OK if balanced samples, and variance difference less than factor of 10)

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A non-parametric test similar to a single factor ANOVA

Uses the ranks of the data points

Conceptually similar to Mann-Whitney U test.

Multiple-factor ANOVA (an advanced topic)

- A factor is a categorical variable
- ANOVAs can be generalized to look at more than one categorical variable at a time
- Not only can we ask whether each categorical variable affects a numerical variable, but also do they interact in affecting the numerical variable.

2-factor ANOVA: Example





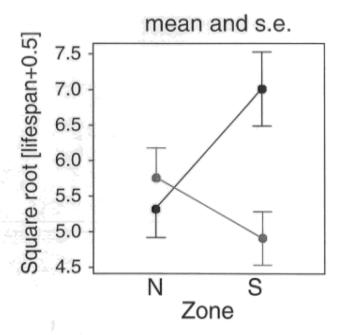
"rayed"

Heliconius erato

"postman"

This experiment uses two "morphs":
the rayed morph from the "north,
and the postman morph from the "south."

Testing multiple hypotheses



"postman" —— "rayed" —— H₀: Mean lifespans are the same in both geographical zones.

H₀: Mean lifespans are the same for both morphs.

H₀: There is no interaction between geographical zone and morph.

Heliconius ANOVA table

Source of variation	SS	df	MS	F	Р
Zone	9.1	1	9.1	0.96	0.327
Morph	34.6	1	34.6	3.68	0.056
Zone*Morph	80.5	1	80.5	8.59	0.004
Error	1837.9	196	9.38		

Fixed vs. random effects

1. Fixed effects: With fixed effects, the treatments are chosen by the experimenter. They are not a random subset of all possible treatments.

```
(e.g., specific drug treatments, specific diets, season...)
```

2. Random effects: With random effects, the groups are a random sample from all possible groups.

```
(e.g., family, location, ...)
Useful for analysis of repeatability of measurements.
```

For single-factor ANOVAs, there is no difference in the statistics for fixed or random effects.

ANOVA's v. t-tests

An ANOVA with 2 groups is mathematically equivalent to a two-tailed 2-sample *t*-test.

Multiple comparisons

Probability of a Type I error in N tests = $1-(1-\alpha)^N$

For 20 tests, the probability of at least one Type I error is ~65%.

"Bonferroni correction" for multiple comparisons

Uses a smaller α value:

$$\alpha' = \frac{\alpha}{\text{number of tests}}$$

Which groups are different?

After finding evidence for differences among means with ANOVA, sometimes we want to know:

Which groups are different from which others?

Planned comparisons: similar to two-sample t-test, but using all samples to estimate SE.

Unplanned comparisons: Need to adjust for inflated Type 1 error rate.

One method: the Tukey-Kramer test

The Tukey-Kramer test

Done after finding variation among groups with single-factor ANOVA.

Compares all group means to all other group means

The wood-wide web

Trees (and other plants) are often connected by roots via mycorrhizae, which allow the exchange of resources.



Test for carbon transfer between birch and Douglas fir; Comparing effects of shading on fir

Net amount of carbon transferred from birch to fir

Shade treatment	Sample mean	Sample standard deviation	Sample size
Deep shade	18.33	6.98	5
Partial shade	8.29	4.76	5
No shade	5.21	3.00	5

ANOVA results

Source of variation	SS	df	MS	F	Р
Groups (treatments)	470.704	2	235.352	8.784	0.004
Error	321.512	12	26.792		
Total	792.216	14			

Order the group means

No shade	Partial shade	Deep shade
\overline{Y}_3	\overline{Y}_2	\overline{Y}_1
5.21	8.29	18.33

Null hypotheses for Tukey-Kramer

$$H_0: \ \mu_1 = \mu_2$$

$$H_0: \mu_1 = \mu_3$$

$$H_0: \mu_2 = \mu_3$$

Why not use a series of two-sample *t*-tests?

Multiple comparisons would cause the *t*-tests to reject too many true null hypotheses.

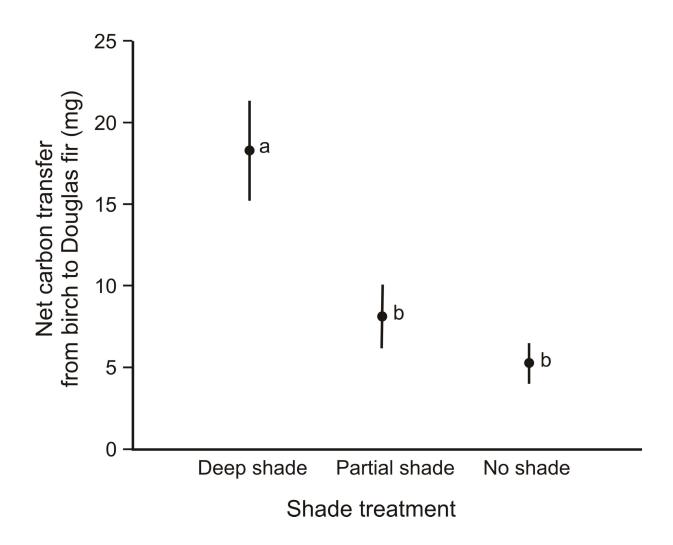
Tukey-Kramer adjusts for the number of tests.

Tukey-Kramer also uses information about the variance within groups from all the data, so it has more power than a *t*-test with a Bonferroni correction.

Tukey-Kramer Results

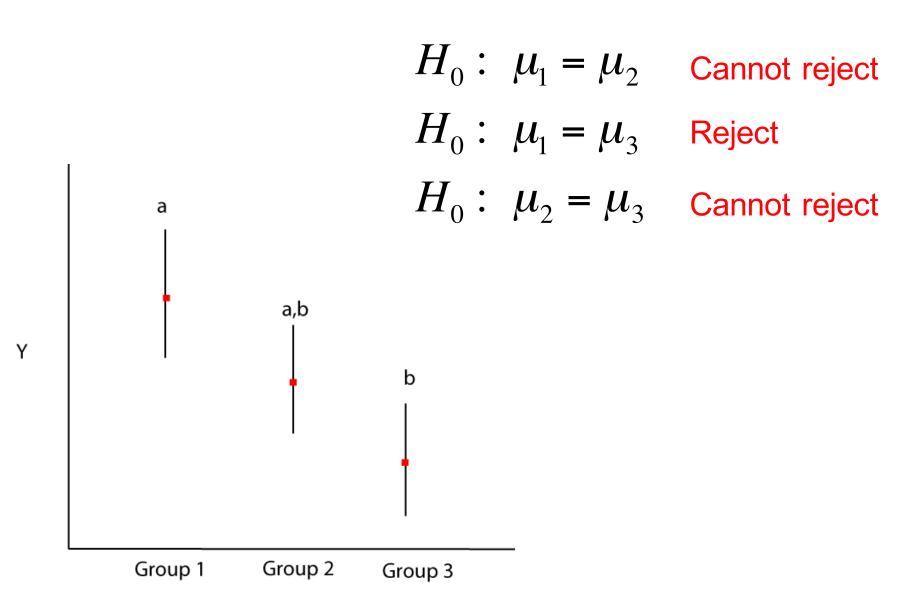
				Test statistic	Critical value	
Group i	Group j	$Y_i - Y_j$	SE	q	$q_{0.05,3,12}$	Conclusion
Deep	No	13.12	3.273693	4.008	2.67	Reject H_0
Deep	Partial	10.04	3.273693	3.067	2.67	Reject H_0
Partial	No	3.08	3.273693	0.941	2.67	Do not reject H_0

$$SE = \sqrt{MS_{error} \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}$$

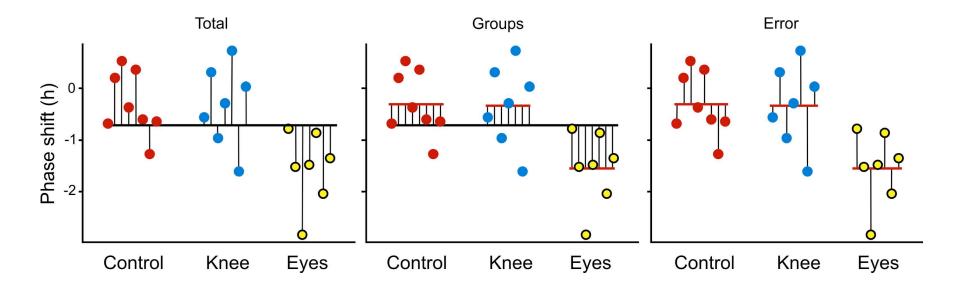


Groups which cannot be distinguished share the same letter.

Another (imaginary) example:



With the Tukey-Kramer method, the probability of making at least one Type 1 error throughout the course of testing all pairs of means is no greater than the significance level α .



$$SS_{total} = SS_{groups} + SS_{error}$$

Announcement

- lab report due today
- new assignment on Friday
- the sun is shining

Comparing the means of more than two groups

Chapter 15

Analysis of variance (ANOVA)

- Like a t-test, but can compare more than two groups
- Asks whether any of two or more means is different from any other.
- In other words, is the variance among groups greater than zero?

Null hypothesis for simple ANOVA

H₀: Variance among groups = 0

OR

• H_0 : $\mu_1 = \mu_2 = \mu_3 = \mu_4 = \dots \mu_k$

Under H₀, the sample mean of each group should vary because of sampling error.

standard deviation of sample means

standard error (when the true mean is constant)

$$\sigma_{\bar{x}} = \frac{\sigma_{x}}{\sqrt{n}}$$

Squaring the standard error, the variance among sample means due to sampling error should be:

$$\sigma_{\bar{x}}^2 = \frac{\sigma_x^2}{n}$$

Under H₀ FALSE:

var(sample mean) = var(sample) + var(real means)



present of means of groups are different $(H_0 \text{ is } \textbf{false})$

Under H₀ FALSE:

$$\sigma_{\bar{x}}^2 = \frac{\sigma_x^2}{n} + Variance[\mu_i]$$

variance among sample means = variance due to sampling error

real variance among population means

With ANOVA, we test whether the variance among true group means is greater than zero.

$$Variance[\mu_i] > 0?$$

With ANOVA, we test whether the variance among true group means is greater than zero.

$$Variance[\mu_i] > 0?$$

We do this by asking whether the variance among sample means is greater than expected under the null hypothesis:

$$\sigma_{\bar{x}}^2 > \frac{\sigma_x^2}{n}?$$

Above is based on population parameters

but.....

we only have *estimates* through our samples.

Variance among groups
$$\sigma_{\bar{x}}^2 > \frac{\sigma_x^2}{n}$$
? Variance within groups n

$$n\sigma_{\bar{x}}^2 > \sigma_x^2?$$

Population parameters

Estimates from sample

$$\sigma_{\bar{x}}^{2} > \frac{\sigma_{x}^{2}}{n}?$$

$$n\sigma_{\bar{x}}^{2} > \sigma_{x}^{2}?$$

Population parameters

Estimates from sample

$$n\sigma_{\bar{x}}^2$$

 $n\sigma_{\overline{x}}^2$ is estimated by the "Mean Squares Group" MS_{group}

'n times variance among groups'

$$\sigma_x^2$$

is the variance within groups, O_{v}^{2} estimated by the "Mean Squares Error"

'variance within groups'

How to calculate the MS_{group} and MS_{error} from sample

Mean squares group

Abbreviation: MS_{group}

Variance in mean due to sampling error

Estimates this parameter: $n\left(\frac{\sigma_x^2}{n} + Variance[\mu_i]\right)$

Variance in mean due to differences in means

Formula:
$$MS_{groups} = \frac{33_{groups}}{df_{groups}}$$

Mean squares group

$$SS_{group} = \sum n_i (\overline{X}_i - \overline{X})^2$$

i = group j = indiv.

 \bar{X}_i is the mean of group i, and

$$\bar{X} = \sum_{i=1}^{\kappa} \sum_{j=1}^{n_i} \frac{X_{ij}}{N}$$

 $df_{groups} = k - 1$ (where k is the number of groups)

$$MS_{groups} = \frac{SS_{groups}}{df_{groups}}$$

Mean squares error

Abbreviation:

$$MS_{error}$$

Estimates this parameter:

$$\sigma_x^2$$

Formula:
$$MS_{error} = \frac{SS_{error}}{df_{error}}$$

Mean squares error

Error sum of squares =

$$SS_{error} = \sum df_i s_i^2 = \sum s_i^2 (n_i - 1)$$

Error degrees of freedom =

$$df_{error} = \sum df_i = \sum (n_i - 1) = N - k$$

$$MS_{error} = \frac{SS_{error}}{df_{error}} = \frac{\sum s_i^2(n_i - 1)}{N - k}$$

Test statistic: F

If
$$H_0$$
 is true, then $n \sigma_{\bar{x}}^2 = \sigma_x^2$

In other words:
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But, refers to population parameters. We must estimate F from samples with: MS_{group} / MS_{error}

F if null hypothesis is false:

We test whether the F ratio is greater than one, as it would be if H_0 is false:

$$F = \frac{n\left(\frac{\sigma_x^2}{n} + Variance[\mu_i]\right)}{\sigma_x^2} > 1$$

Sampling error

Take into account sampling error

 Often, F calculated from data will be greater than one even when the null is true

 Hence we must compare F to a null distribution.

Using sample to get F

$$F = \frac{MS_{groups}}{MS_{errors}}$$

ANOVA table

Source	SS	df	MS	F	P
Group					
Error					
Total					

An ANOVA table is a convenient way to keep track of the important calculations.

Scientific papers often report ANOVA results with ANOVA tables.

Example: Body temperature of squirrels in low, medium and hot environments



Hypotheses

H₀: Mean body temperature is the same for all three groups of squirrels.

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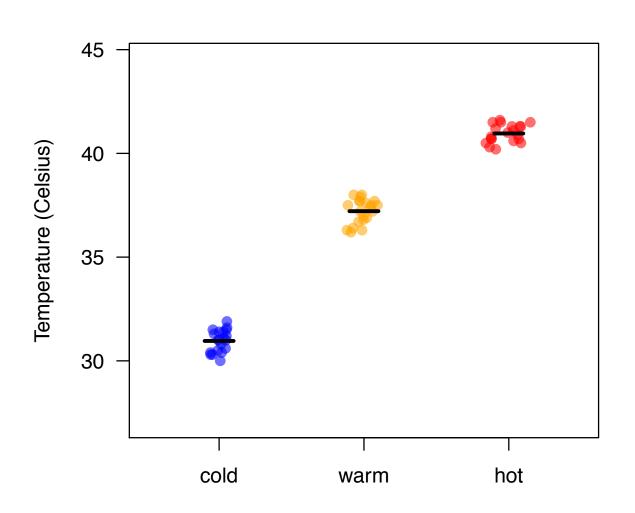
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Plot the data



Summary data

Group	$\overline{\mathcal{X}}$	S	n	
Cold	31.0	0.551	20	
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Total sample size:
$$N = \sum n = 20 + 21 = 20 = 61$$

Next steps

Calculate:

- MS_{error}
- MS_{group}
- F value

MS_{error} for squirrels

$$SS_{error} = \sum df_i s_i^2$$

= $19(0.551)^2 + 20(0.582)^2 + 19(0.430)^2$
= 16.1

$$df_{error} = 19 + 20 + 19 = 58$$

$$MS_{error} = \frac{16.1}{58} = 0.277$$

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$$SS_{group} = 20(31.0-36.4)^2 + 21(37.2-36.4)^2 + 20(41.0-36.4)^2$$

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MS_{group} squares group:

$$df_{group} = k - 1 = 3-1=2$$

$$MS_{groups} = \frac{SS_{groups}}{df_{groups}} = \frac{1015.7}{2} = 507.9$$

F statistic

$$F = \frac{MS_{group}}{MS_{error}} = \frac{507.9}{0.277} = 1834.7$$

MS_{group} is always in the <u>numerator</u>; MS_{error} is always in the <u>denominator</u>.

Compare to $F_{\alpha(1),df_group,df_error}$

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Error	16.1	58	0.277		
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• R² is the fraction of variation that is explained by group differences:

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For Squirrel study:

$$R^2 = 1015.7 / 1031.8 = 0.98$$

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- Random samples.
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A non-parametric test similar to a single factor ANOVA

Uses the ranks of the data points

Conceptually similar to Mann-Whitney U test.

What's up next

- Multi factor anova
- Testing multiple hypotheses
- Correct for multiple testing
- Tukey-Kramer test

Multiple-factor ANOVA (an advanced topic)

- A factor is a categorical variable
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2-factor ANOVA: Example





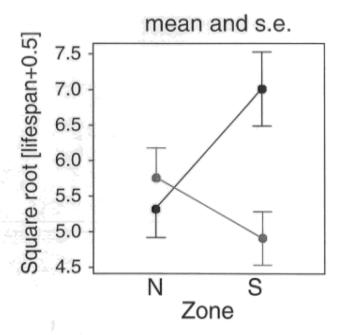
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Heliconius erato

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"postman" —— "rayed" —— H₀: Mean lifespans are the same in both geographical zones.

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The Tukey-Kramer test

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Compares all group means to all other group means

The wood-wide web

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Source of variation	SS	df	MS	F	Р
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Error	321.512	12	26.792		
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Order the group means

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Null hypotheses for Tukey-Kramer

$$H_0: \mu_1 = \mu_2$$

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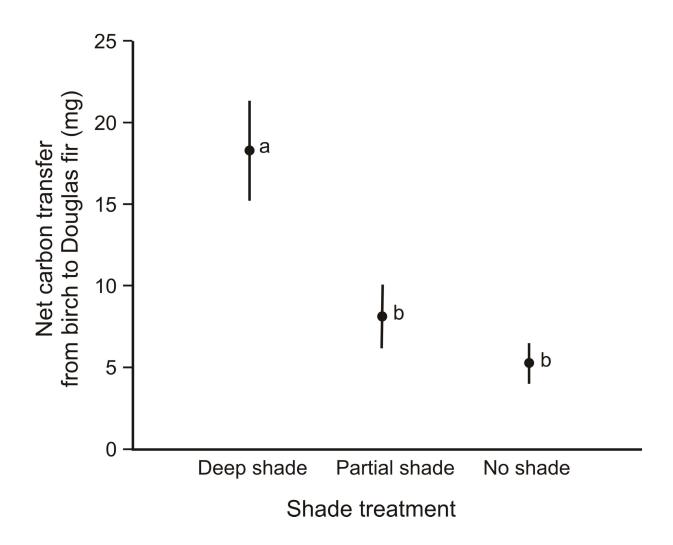
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Tukey-Kramer Results

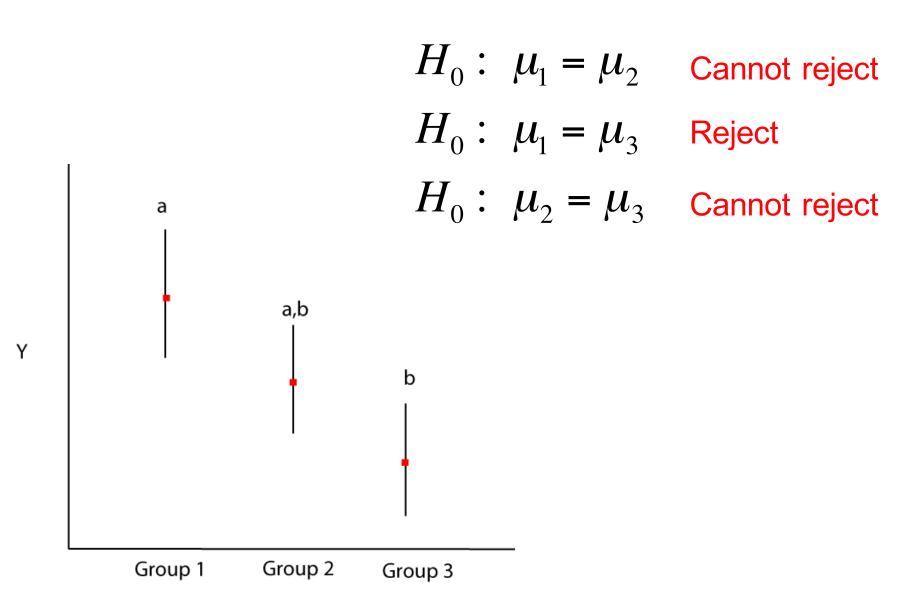
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Group i	Group j	$Y_i - Y_j$	SE	q	$q_{0.05,3,12}$	Conclusion
Deep	No	13.12	3.273693	4.008	2.67	Reject H_0
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$$SE = \sqrt{MS_{error} \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}$$

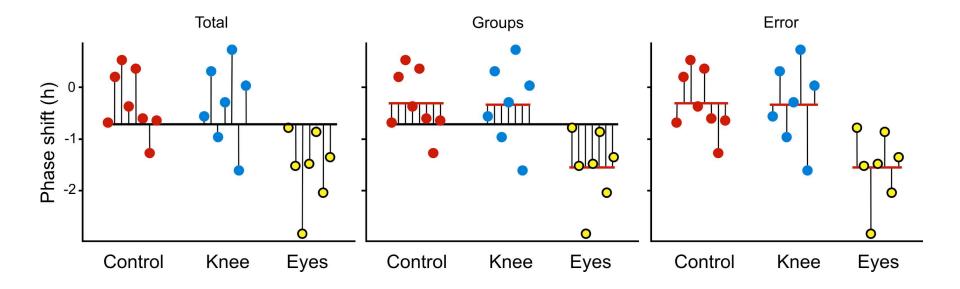


Groups which cannot be distinguished share the same letter.

Another (imaginary) example:



With the Tukey-Kramer method, the probability of making at least one Type 1 error throughout the course of testing all pairs of means is no greater than the significance level α .



$$SS_{total} = SS_{groups} + SS_{error}$$

Announcement

- new assignment available after class on Connect
- Next week your TA will discuss during lab the lab report analyses
- Labs & Easter:
 - -Lab 9: March 21-24 for Mon-Thur sections and April 1st for the Friday section.
 - -Lab 10: everyone April 4-8
- the sun is still shining

Comparing the means of more than two groups

Chapter 15

Analysis of variance (ANOVA)

 Asks whether any of two or more means is different from any other.

 In other words, is the variance among groups greater than zero?

Null hypothesis for simple ANOVA

• H_0 : Variance among groups = 0

OR

• H_0 : $\mu_1 = \mu_2 = \mu_3 = \mu_4 = \dots \mu_k$

Under H₀:

var(sample mean) = var(sample)



expected due to sampling error

Squaring the standard error, the variance among sample means due to sampling error should be:

$$\sigma_{\bar{x}}^2 = \frac{\sigma_x^2}{n}$$

Under H₀ FALSE:

var(sample mean) = var(sample) + var(real means)



present of means of groups are different $(H_0 \text{ is } \textbf{false})$

Under H₀ FALSE:

$$\sigma_{\bar{x}}^2 = \frac{\sigma_x^2}{n} + Variance[\mu_i]$$

variance among sample means = variance due to sampling error

real variance among population means

With ANOVA, we test whether the variance among true group means is greater than zero.

$$Variance[\mu_i] > 0?$$

With ANOVA, we test whether the variance among true group means is greater than zero.

$$Variance[\mu_i] > 0?$$

We do this by asking whether the variance among sample means is greater than expected under the null hypothesis:

$$\sigma_{\bar{x}}^2 > \frac{\sigma_x^2}{n}?$$

$$\sigma_{\bar{x}}^{2} > \frac{\sigma_{x}^{2}}{n}?$$

$$n\sigma_{\bar{x}}^{2} > \sigma_{x}^{2}?$$

Population parameters

Estimates from sample

$$n\sigma_{\bar{x}}^2$$

 $n\sigma_{\overline{x}}^2$ is estimated by the "Mean Squares Group" MS_{group}

'n times variance among groups'

$$\sigma_x^2$$

is the variance within groups, O_{v}^{2} estimated by the "Mean Squares Error"

'variance within groups'

Test statistic: F

If
$$H_0$$
 is true, then $n \sigma_{\bar{x}}^2 = \sigma_x^2$

In other words:
$$F = \frac{n \sigma_{\overline{x}}^2}{\sigma_x^2} = 1$$

F if null hypothesis is false:

We test whether the F ratio is greater than one, as it would be if H_0 is false:

$$F = \frac{n\left(\frac{\sigma_x^2}{n} + Variance[\mu_i]\right)}{\sigma_x^2} > 1$$

Using sample to get F

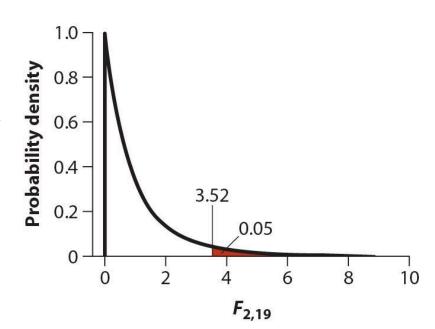
$$F = \frac{MS_{groups}}{MS_{errors}}$$

Compare to $F_{\alpha(1),df_group,df_error}$

$$F_{0.05(1),2,58} = 3.15.$$



Means we are looking at the right hand tail of the F distribution



ANOVA table – squirrel data

Source	SS	df	MS	F	P
Group	1015.7	2	507.9	1834.7	<0.0001
Error	16.1	58	0.277		
Total	1031.8	60			

Variability explained: R²

• R² is the fraction of variation that is explained by group differences:

$$R^2 = \frac{SS_{groups}}{SS_{total}}$$

Assumptions of ANOVA

- Random samples.
- Normal distributions for each population.
 (Small deviations are OK)
- Equal variances for all populations.
 (Homoscedasticity)
 (But OK if balanced samples, and variance difference less than factor of 10)

Kruskal-Wallis test

A non-parametric test similar to a single factor ANOVA

Uses the ranks of the data points

Conceptually similar to Mann-Whitney U test.

Today

- Multi factor anova
- Testing multiple hypotheses
- Correct for multiple testing
- Tukey-Kramer test

Multiple-factor ANOVA (an advanced topic)

- A factor is a categorical variable
- ANOVAs can be generalized to look at more than one categorical variable at a time
- Not only can we ask whether each categorical variable affects a numerical variable, but also do they interact in affecting the numerical variable.

2-factor ANOVA: Example





"rayed"

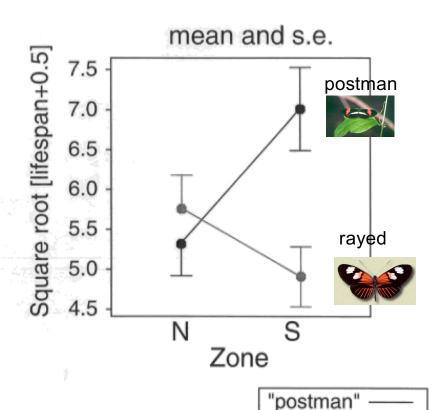
Heliconius erato

"postman"

This experiment uses two "morphs":

- rayed morph from the "north,
- postman morph from the "south."

Testing multiple hypotheses



"rayed"

H₀: Mean lifespans are the same in both geographical zones.

H₀: Mean lifespans are the same for both morphs.

H₀: There is no interaction between geographical zone and morph.

Heliconius ANOVA table

Source of variation	SS	df	MS	F	Р
Zone	9.1	1	9.1	0.96	0.327
Morph	34.6	1	34.6	3.68	0.056
Zone*Morph	80.5	1	80.5	8.59	0.004
Error	1837.9	196	9.38		

Fixed vs. random effects

1. Fixed effects: With fixed effects, the treatments are chosen by the experimenter. They are not a random subset of all possible treatments.

```
(e.g., specific drug treatments, specific diets, season...)
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```
(e.g., family, location, ...)
Useful for analysis of repeatability of measurements.
```

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Probability you don't make a Type I error N times in a row.

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pooled sample variance (MS_{error}) based on all k groups

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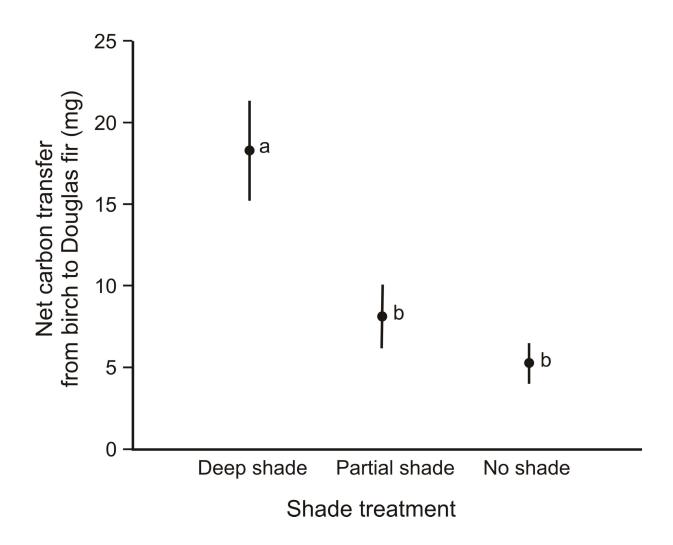
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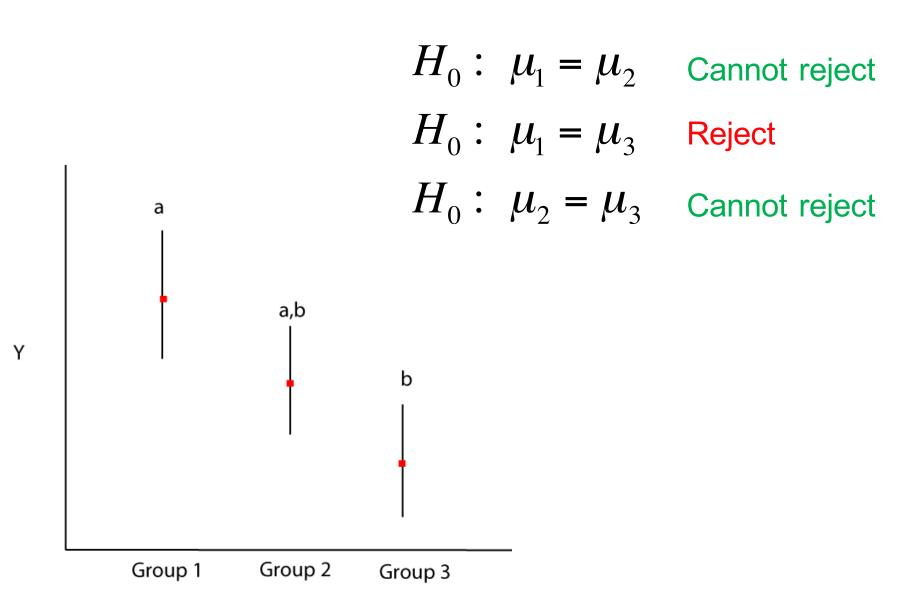
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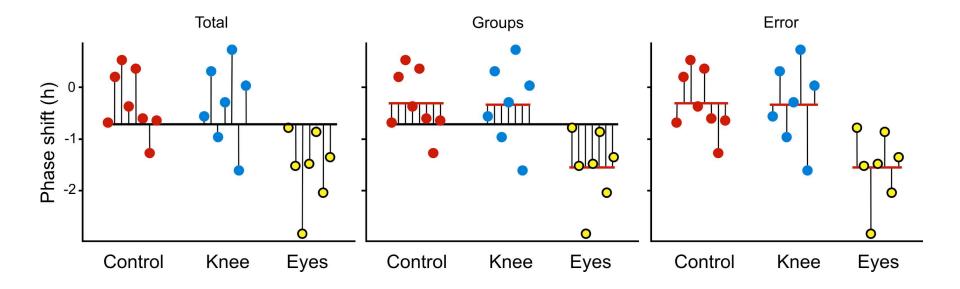


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