

<div> <div>Chapter 5: Comparisons among several samples, One-way ANOVA</div> <div>Class 8: 2/25/09 W</div> </div>	<div>Slide 1 Chapter 5: Comparisons among several samples, One-way ANOVA</div> <div>NOTES:</div>
<div> <div>HW 6 due Monday 3/1/09 9:50</div> <div>Submit as Myname-HW6.doc (or *.rtf)</div> <ul style="list-style-type: none"> <li>Read Chapter 5 <b>Comparisons among several samples</b></li> <li>Comment on Chapter 5 conceptual problems in Blackboard Vista4</li> <li>Computation Problem 6 <ul style="list-style-type: none"> <li>Problem 4.30 Sunlight protection factor</li> </ul> </li> </ul> <div>EEOS611</div> </div>	<div>Slide 2 HW 6 due Monday 3/1/09 9:50</div> <div>NOTES:</div>
<div> <div>HW 7 due Thursday 3/4/09 Noon</div> <div>Submit as Myname-HW7.doc (or *.rtf)</div> <ul style="list-style-type: none"> <li>Read Chapter 6 <b>Comparisons among several samples</b></li> <li>Comment on Chapter 6 conceptual problems in Blackboard Vista4</li> <li>Computation Problem 7 <ul style="list-style-type: none"> <li>Problem 5.25 Duodenal ulcers</li> </ul> </li> </ul> <div>EEOS611</div> </div>	<div>Slide 3 HW 7 due Thursday 3/4/09 Noon</div> <div>NOTES:</div>

<div data-bbox="350 168 665 199" data-label="Section-Header"> <h3>Student Presentations</h3> </div> <div data-bbox="355 210 654 233" data-label="Text"> <p>Starting at 10:50 (8 minutes each)</p> </div> <div data-bbox="263 243 584 430" data-label="List-Group"> <ul style="list-style-type: none"> <li>• Keith Cialino for HW 2 <ul style="list-style-type: none"> <li>▸ Ex 1.21</li> </ul> </li> <li>• Seth Sheldon for HW 3 <ul style="list-style-type: none"> <li>▸ 2.21 Bumpus's data: weights of Bumpus's birds</li> </ul> </li> <li>• Barry Fradkin for HW 4. <ul style="list-style-type: none"> <li>▸ 3.28 Pollen removal</li> </ul> </li> </ul> </div> <div data-bbox="657 506 779 533" data-label="Text"> <p>EEOS611</p> </div>	<div data-bbox="820 130 1234 163" data-label="Section-Header"> <h3>Slide 4 Student Presentations</h3> </div> <div data-bbox="820 254 940 285" data-label="Text"> <p>NOTES:</p> </div>
<div data-bbox="388 655 612 690" data-label="Section-Header"> <h3>Chapter 4 (End)</h3> </div> <div data-bbox="277 697 745 741" data-label="Text"> <p>Rank-based, nonparametric analogues to the paired t test: Signed rank &amp; Sign tests</p> </div> <div data-bbox="246 747 363 867" data-label="Image"> </div> <div data-bbox="376 745 620 848" data-label="Text"> <p><b>Wilcoxon's signed rank test</b> Frank Wilcoxon of American Cyanamide [See Salsburg, 2001, The Lady Tasting Tea, for biographical sketch]</p> </div> <div data-bbox="631 753 779 858" data-label="Text"> <p>Asymptotic relative efficiency &gt;0.864, 95.5% for normally distributed data</p> </div> <div data-bbox="250 882 360 1014" data-label="Image"> </div> <div data-bbox="376 915 579 978" data-label="Text"> <p><b>Fisher's signed test</b> [See Salsburg, 2001 for biographical sketch]</p> </div> <div data-bbox="610 886 768 991" data-label="Text"> <p>Asymptotic relative efficiency 63.7% for normally distributed data</p> </div> <div data-bbox="657 995 779 1020" data-label="Text"> <p>EEOS611</p> </div>	<div data-bbox="820 621 1159 655" data-label="Section-Header"> <h3>Slide 5 Chapter 4 (End)</h3> </div> <div data-bbox="820 741 940 772" data-label="Text"> <p>NOTES:</p> </div>
<div data-bbox="324 1213 698 1289" data-label="Section-Header"> <h3>4.4 Alternatives to the paired t test</h3> </div> <div data-bbox="308 1299 717 1327" data-label="Text"> <p>Wilcoxon sign-rank and Fisher Sign tests</p> </div>	<div data-bbox="820 1110 1408 1144" data-label="Section-Header"> <h3>Slide 6 4.4 Alternatives to the paired t test</h3> </div> <div data-bbox="820 1232 940 1264" data-label="Text"> <p>NOTES:</p> </div>

## Anatomical abnormalities & schizophrenia

Case 2.2 (Sleuth p 30): 15 pairs of twins, paired  $t$  test

Display 2.2

Differences in volumes ( $\text{cm}^3$ ) of left hippocampus in fifteen sets of monozygotic twins where one twin is affected by schizophrenia

Pair #	Unaffected	Affected	Difference	Differences	
1	1.94	1.27	0.67	-2	9
2	1.44	1.63	-0.19	-1	9
3	1.56	1.47	0.09	-0	9
4	1.58	1.39	0.19	0	23479
5	2.06	1.93	0.13	1	0139
6	1.66	1.26	0.40	2	3
7	1.75	1.71	0.04	3	0
8	1.77	1.67	0.10	4	0
9	1.78	1.28	0.50	5	09
10	1.92	1.85	0.07	6	7
11	1.25	1.02	0.23	7	7
12	1.93	1.34	0.59		
13	2.04	2.02	0.02		
14	1.62	1.59	0.03		
15	2.08	1.97	0.11		

Average: 0.199  
Sample SD: 0.238  
n: 15

Legend: | 6 | 7 represents 0.67  $\text{cm}^3$

## Slide 7 Anatomical abnormalities & schizophrenia

NOTES:

## Case 2.2 Statistical Summary

Sleuth, p. 31

There is substantial evidence that the mean difference in the left hippocampus volumes between schizophrenic individuals and their nonschizophrenic twins is nonzero (two-sided  $p$ -value = 0.006, from a paired  $t$  test). It is estimated that the mean volume is 0.20  $\text{cm}^3$  smaller for those with schizophrenia (about 11% smaller). A 95% confidence interval for the difference is from 0.07 to 0.33  $\text{cm}^3$ .

## Slide 8 Case 2.2 Statistical Summary

NOTES:

## Wilcoxon signed rank test

Display 4.12

Signed-rank test statistic computations; schizophrenia study

Pair	Unaffected	Affected	Difference	Ordered Magnitude	Order	Rank	+Ranks	-Ranks
1	1.94	1.27	.67	.02 (+)	1	1	1	
2	1.44	1.63	-.19	.03 (+)	2	2		2
3	1.56	1.47	.09	.04 (+)	3	3	3	
4	1.58	1.39	.19	.07 (+)	4	4	4	
5	2.06	1.93	.13	.09 (+)	5	5	5	
6	1.66	1.26	.40	.10 (+)	6	6	6	
7	1.75	1.71	.04	.11 (+)	7	7	7	
8	1.77	1.67	.10	.13 (+)	8	8	8	
9	1.78	1.28	.50	.19 (+)	9	9	9.5	
10	1.92	1.85	.07	.19 (-)	10	9.5		9.5
11	1.25	1.02	.23	.23 (+)	11	11	11	
12	1.93	1.34	.59	.40 (+)	12	12	12	
13	2.04	2.02	.02	.50 (+)	13	13	13	
14	1.62	1.59	.03	.59 (+)	14	14	14	
15	2.08	1.97	.11	.67 (+)	15	15	15	

SPSS dscards pairs with equal values\*

③ Correct the standard deviation SD(T), based on the pattern of ties [2, 5]

① Order the absolute differences and assign ranks to them

② Signed rank statistics = sum of ranks for positive differences:

## Slide 9 Wilcoxon signed rank test

NOTES:

### Ties in signed rank tests

#### Two sorts of ties in the signed rank test, 1 of 2

1) If you have identical values in in both pairs, Wilcoxon recommended that those paired observations be dropped. That is still the standard recommendation, and SPSS uses this recommendation.

Hollander and Wolfe's Nonparametric statistics, 2nd ed (p. 46) covers the problem of dropping ties of the first sort. If there are many ties, H & W recommend using another test. They also state that you could leave the tied samples in, and use a random number generator to randomly assign positive or negative signs for the zeros. If you want a more conservative 1-sided test, assign all of the tied differences to the group that would make it less likely to reject the null. For example, if you are testing lipitor's effects on cholesterol and a patient had identical cholesterol levels before and after, then assign that difference as if the lipitor blood sample had the higher cholesterol. If you still reject the null, your conservative test would be less likely to result in a Type I error, but of course the probability of Type II error (failing to reject a false null), Pratt (1959), cited in both Lehmann and Hollander & Wolfe, provides a more thorough review of ties. Lehmann cites more recent papers on dealing with the 1st sort of ties in signed rank tests.

### Slide 10 Ties in signed rank tests

NOTES:

### Dealing with tied pairs

#### Two sorts of ties in the signed rank test, 2 of 2

The second sort of ties occurs after the absolute values of the differences are ranked

2) Ties may result after the absolute values of the differences between paired observations are ranked. Two or more differences may have the same absolute value. Those ties are not discarded, and the variance formula is adjusted to take into account the number of tied groups [See next slide]

### Slide 11 Dealing with tied pairs

NOTES:

### SPSS algorithms, signed rank test

There are exact tests if no tied ranks

$$Z = \frac{\min(S_p, S_n) - (n(n+1)/4)}{\sqrt{n(n+1)(2n+1)/24 - \sum_{j=1}^l (t_j^3 - t_j)/48}}$$

L = tied groups  
t<sub>j</sub> = items in each tied group

where

Asymptotic relative efficiency > 0.864,  
95.5% for normally distributed data

$n$  Number of cases with non-zero differences

$l$  Number of ties

$t_j$  Number of elements in the  $j$ -th tie,  $j = 1, \dots, l$

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### Slide 12 SPSS algorithms, signed rank test

NOTES:

## Fisher's sign test

### Straightforward application of the 1-sample binomial test

- Given that the probability of a + sign = probability of a minus sign = 0.5,
- What is the probability of observing exactly  $k$  positive signs in  $n$  Bernoulli (binomial) trials?
  - $P(X=k) = n \text{ Choose } k * p^k (1-p)^{n-k}$ 
    - $X$  has a binomial distribution
    - Must sum probability for observed value of  $k$ , and all more extreme values of  $k$ ,
- Statistical sleuth provides only the normal approximation to the binomial, but SPSS will provide the exact test for  $n < 30$ .

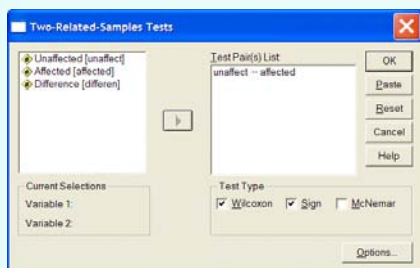
Frequencies		
AFFECTED - UNAFFECT	Negative Differences	14
	Positive Differences	1
	Ties <sup>a</sup>	0
	Total	15
Test Statistics <sup>b</sup>		
	AFFECTED - UNAFFECT	
Exact Sig. (2-tailed)		.0010 <sup>a</sup>
a. Binomial distribution used.		
b. Sign Test		

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## Slide 13 Fisher's sign test

NOTES:

## Sign test in SPSS



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## Slide 14 Sign test in SPSS

NOTES:

## Schizophrenia data

Sleuth (Ch 4.4.1 p. 99) Sign test: Sleuth p. 99  
Sleuth presents the large sample approximation; exact test possible with binomial distribution (used by SPSS)

Display 2.2  
Differences in volumes (cm<sup>3</sup>) of left hippocampus in fifteen sets of monozygotic twins where one twin is affected by schizophrenia

Pair	Unaffected	Affected	Difference
1	1.04	1.27	0.67
2	1.44	1.63	-0.19
3	1.56	1.47	0.09
4	1.58	1.39	0.19
5	2.08	1.93	0.13
6	1.66	1.26	0.40
7	1.75	1.71	0.04
8	1.77	1.87	-0.10
9	1.78	1.28	0.50
10	1.92	1.85	0.07
11	1.25	1.62	-0.23
12	1.93	1.34	0.59
13	2.04	2.62	-0.62
14	1.62	1.59	0.03
15	2.08	1.97	0.11

Legend: 6 7 represents 0.67 cm<sup>3</sup>

Paired  $t$  test for equal difference, two-tailed  $p=0.006$   
[See sleuth p. 31]

Frequencies		
AFFECTED - UNAFFECT	Negative Differences	14
	Positive Differences	1
	Ties <sup>a</sup>	0
	Total	15
Test Statistics <sup>b</sup>		
	AFFECTED - UNAFFECT	
Exact Sig. (2-tailed)		.0010 <sup>a</sup>
a. Binomial distribution used.		
b. Sign Test		

## Slide 15 Schizophrenia data

NOTES:

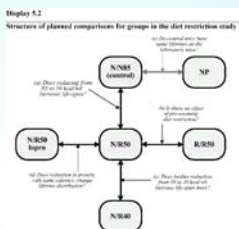
<div data-bbox="305 161 722 224" data-label="Section-Header"> <h3>Assumptions of the Wilcoxon signed rank test</h3> </div> <div data-bbox="235 231 735 541" data-label="List-Group"> <ul style="list-style-type: none"> <li>• Underlying distribution of the differences are continuous and symmetric about zero [Hollander &amp; Wolfe p. 43]</li> <li>• Differences within &amp; between pairs on an ordinal scale</li> <li>• Siegel: use large sample approximation if number of pairs exceeds 25, otherwise use tabulated values of the Wilcoxon signed rank statistic <ul style="list-style-type: none"> <li>▸ Note that there are exact tests for Wilcoxon signed rank tests if there are ties (described in Hollander &amp; Wolfe p 46-47)</li> </ul> </li> </ul> </div>	<div data-bbox="820 130 1349 205" data-label="Section-Header"> <h3>Slide 16 Assumptions of the Wilcoxon signed rank test</h3> </div> <div data-bbox="820 291 940 323" data-label="Text"> <p>NOTES:</p> </div>
<div data-bbox="363 693 649 728" data-label="Section-Header"> <h3>Conclusions (1 of 2)</h3> </div> <div data-bbox="350 735 665 758" data-label="Section-Header"> <h4>Chapter 4 Alternatives to the t tools</h4> </div> <div data-bbox="241 756 716 1050" data-label="List-Group"> <ul style="list-style-type: none"> <li>• Consider using alternatives to the t tools if <ul style="list-style-type: none"> <li>▸ The assumptions are grossly violated or</li> <li>▸ The sample sizes are too small to test distributional assumptions</li> </ul> </li> <li>• Wilcoxon rank sum test <ul style="list-style-type: none"> <li>▸ Appropriate for small sample sizes, but use the exact tests not the normal approximation</li> <li>▸ Appropriate in the presence of outliers</li> <li>▸ Ties are not a problem if the ties-correction used</li> <li>▸ Not appropriate for samples with unequal variances (try Fligner-Policello only if the sample sizes are large)</li> </ul> </li> </ul> </div> <div data-bbox="657 1031 779 1058" data-label="Text"> <p>EEOS611</p> </div>	<div data-bbox="820 657 1221 693" data-label="Section-Header"> <h3>Slide 17 Conclusions (1 of 2)</h3> </div> <div data-bbox="820 779 940 810" data-label="Text"> <p>NOTES:</p> </div>
<div data-bbox="363 1182 649 1218" data-label="Section-Header"> <h3>Conclusions (2 of 2)</h3> </div> <div data-bbox="350 1224 665 1247" data-label="Section-Header"> <h4>Chapter 4 Alternatives to the t tools</h4> </div> <div data-bbox="241 1253 730 1512" data-label="List-Group"> <ul style="list-style-type: none"> <li>• Permutation test <ul style="list-style-type: none"> <li>▸ Appropriate for small sample sizes, when the Student's <i>t</i> distribution might not be appropriate</li> <li>▸ Does not protect against the problem of unequal variances (the Fisher-Behrens problem) <ul style="list-style-type: none"> <li>■ Note that the solution to Case Study 4.1 is based on the equal variance <i>t</i> test.</li> </ul> </li> </ul> </li> <li>• Paired data: tests based on ranks <ul style="list-style-type: none"> <li>▸ Wilcoxon signed rank test: high power efficiency</li> <li>▸ Sign test, simple application of the 1-sample binomial test</li> </ul> </li> </ul> </div> <div data-bbox="657 1520 779 1547" data-label="Text"> <p>EEOS611</p> </div>	<div data-bbox="820 1146 1221 1180" data-label="Section-Header"> <h3>Slide 18 Conclusions (2 of 2)</h3> </div> <div data-bbox="820 1268 940 1299" data-label="Text"> <p>NOTES:</p> </div>

<div data-bbox="300 235 719 315" data-label="Section-Header"> <h2>Chapter 5: Comparisons among several samples</h2> </div>	<div data-bbox="820 132 1390 207" data-label="Section-Header"> <h3>Slide 19 Chapter 5: Comparisons among several samples</h3> </div> <div data-bbox="820 291 941 323" data-label="Text"> <p>NOTES:</p> </div>
<div data-bbox="253 686 745 753" data-label="Section-Header"> <h2>ANOVA, Analysis of Variance, the foundation of experimental design</h2> </div> <div data-bbox="241 777 547 1052" data-label="List-Group"> <ul style="list-style-type: none"> <li>• Most experimental design is based on an ANOVA framework</li> <li>• One can't really appreciate the need for proper replication without considering the Implications for testing treatment effects with ANOVA <ul style="list-style-type: none"> <li>• Hurlbert's (1984) monograph criticizing statistics in ecological papers is largely a criticism of inappropriate ANOVA design</li> <li>• Hurlbert's pseudoreplication is Underwood's 'model misspecification' and both are largely based on using an inappropriate ANOVA model</li> </ul> </li> <li>• While ANOVA is a proper subset of the general linear model (GLM) and regression, as we'll see, the concepts involving design and partitioning degrees of freedom are more evident in ANOVA models</li> </ul> </div> <div data-bbox="570 779 730 976" data-label="Image"> </div> <div data-bbox="581 978 714 1056" data-label="Caption"> <p>R.A. Fisher, inventor of ANOVA</p> </div>	<div data-bbox="820 659 1375 732" data-label="Section-Header"> <h3>Slide 20 ANOVA, Analysis of Variance, the foundation of experimental design</h3> </div> <div data-bbox="820 819 941 850" data-label="Text"> <p>NOTES:</p> </div>
<div data-bbox="292 1287 734 1367" data-label="Section-Header"> <h2>Case 5.1 Diet restriction &amp; longevity</h2> </div>	<div data-bbox="820 1186 1312 1260" data-label="Section-Header"> <h3>Slide 21 Case 5.1 Diet restriction &amp; longevity</h3> </div> <div data-bbox="820 1346 941 1377" data-label="Text"> <p>NOTES:</p> </div>

## Planned comparisons

These are *a priori* contrasts, not *a posteriori*

- If hypotheses are set in advance, then you can test at a pre-set alpha level, without a *posteriori* (or *post hoc*, multiple comparison) adjustment
  - Recall that  $\alpha = P(\text{Type I error})$
  - See Cook & Farewell (1996, J. Roy. Stat. Soc. A). In dose-response studies, no need to adjust for number of dose treatments.
- One large design allows the use of a more precise estimate of the error variance
  - Separate control vs. treatment t tests are powerful
  - If interaction effects are evident, separate tests can be misleading. They can miss interaction effects.

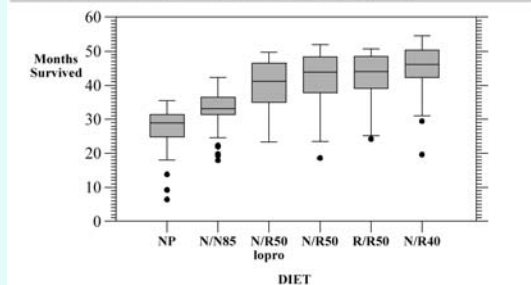


## Slide 22 Planned comparisons

NOTES:

### Display 5.1

### Lifetimes of female mice fed on six different diet regimens



## Slide 23

NOTES:

## Detonator plots

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## Slide 24 Detonator plots

NOTES:



## Summary statistics for lifetimes of mice

Display 5.2, Sleuth 2nd edition page 115

Summary statistics for lifetimes of mice on six different diet regimens

Group	n	Range (mo)	Average	SD	95% CI for Mean
NP	49	6.4 - 35.5	27.4	6.1	25.6 - 29.2
N/N85	57	17.9 - 42.3	32.7	5.1	31.3 - 34.1
N/R50	71	18.6 - 51.9	42.3	7.8	40.5 - 44.1
R/R50	56	24.2 - 50.7	42.9	6.7	41.1 - 44.7
N/R50 lopro	56	23.4 - 49.7	39.7	7.0	37.8 - 41.6
N/R40	60	19.6 - 54.6	45.1	6.7	43.4 - 46.8

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## Slide 25 Summary statistics for lifetimes of mice

NOTES:

## Summary of statistical findings

Case Study 5.1: mouse longevity, 1 of 2

- There is overwhelming evidence that mean lifetimes in the six groups are different (p-value < 0.001); analysis of variance F-test).
- Analysis of the 5 particular questions are
  - (1) There is convincing evidence that lifetime increases as a result of restricting the diet from 85 kcal/wk to 50 kcal/wk (1-sided p-value < 0.0001; t test)
  - (2) There is no evidence that reducing the calories before weaning increased lifetime, when the calorie intake after weaning is 50 kcal/wk (1-sided p value = 0.32, t test). A 95% CI for the amount by which the lifetime under the R/R50 diet exceeds the lifetime under the N/R50 diet is - 1.7 to 2.9 months.

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## Slide 26 Summary of statistical findings

NOTES:

## Summary of statistical findings

Case Study 5.1: mouse longevity, 2 of 2

- Analysis of the 5 particular questions (continued)
  - (3) Further restriction of the diet from 50 to 40 kcal/wk increases lifetime by an estimated 2.8 months (95% CI: 0.5 to 5.1 months). The evidence that this effect is greater than zero is moderate (p=0.017, t test)
  - (4) There was moderate evidence that lifetime was decreased by the lowering of protein in addition to the 50 kcal/wk diet (2-sided p value = 0.024; t-test)
  - (5) There is convincing evidence that the control mice live longer than the mice on the non-purified diet (1-sided p-value < 0.0001)

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## Slide 27 Summary of statistical findings

NOTES:

## Case study 5.2: The Spock trial

### Slide 28 Case study 5.2: The Spock trial

NOTES:

## Case 5.2 The Spock trial

Sleuth, page 117: Dr. Spock's *venire* contained only 1 woman, who was released by the prosecution

Display 5.4

Percents of women in 30-juror *venires* for Boston area U.S. District Court trials, grouped according to the judge presiding

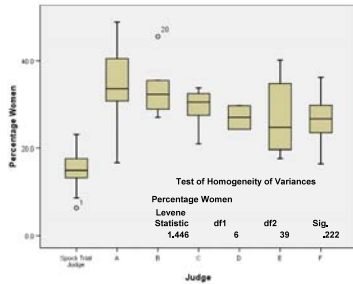
Spock Trial Judge	A	B	C	D	E	F
0	64.87				77.97	65
1	33,36,50,52,77,86	68				
2	31	70.89	10,34,75,75	43.97	15.79	07,35,64,67,95,98
3		08,36	20,27,55		48	19,62
4		05,89	56		02	

Legend: 4|89 represents a *venire* with 48.9% women

1) Is there evidence that women were underrepresented on the Spock judge's *venires*, and 2) Is there evidence that there are differences in women's representation on the other juries?

### Slide 29 Case 5.2 The Spock trial

NOTES:



Is there evidence of a judge effect? There are no true replicates for the Spock-judge effect.

- The percentage of women on the Spock judge's *venires* were substantially lower than the other judges (t test of Spock judge vs. 'Other judges')
- There is little evidence to reject the null hypothesis of no difference in female representation among the other judges  $p=0.32$  (1-way ANOVA)
- The percentage of women is 15% less on the Spock judge's *venires* (95% CI: 10% to 20%)
- Gallagher note: this pooling of judges could be called pseudoreplication, but can be justified as a fixed-effect NESTED ANOVA

### Slide 30

NOTES:

## 5.2 Comparing any two of several means

### 5.2.1 An ideal model for several-sample comparisons

*Gallagher note: Comparisons among means in ANOVA can be analyzed using  $t$  statistics, with a new, more precise estimate of pooled error. It is that pooling, with higher  $df$ , that makes ANOVA a more powerful method than multiple  $t$  tests.*

### Slide 31 5.2 Comparing any two of several means

NOTES:

## 5.2.2 The pooled estimate of the standard deviation, $s_p$

Display 5.6, Sleuth page 120

Pooled estimate of standard deviation; diet restriction data

Group	n	Sample SD
NP	49	6.1
N/NR5	57	5.1
N/R50	71	7.8
R/R50	56	6.7
N/R50 loopro	56	7.0
N/R40	60	6.7

$s_p$  assumes equal variances among groups

Calculate the pooled estimate of variance,  $s_p^2$

$$s_p^2 = \frac{(49-1)(6.1)^2 + (57-1)(5.1)^2 + (71-1)(7.8)^2 + (56-1)(6.7)^2 + (56-1)(7.0)^2 + (60-1)(6.7)^2}{(49-1) + (57-1) + (71-1) + (56-1) + (56-1) + (60-1)}$$

$$= \frac{15,313.90}{343} = 44.647; s_p = \sqrt{44.647} = 6.68$$

$s_p$  is the square root

$df$  is the denominator

### Slide 32 5.2.2 The pooled estimate of the standard deviation, $s_p$

NOTES:

## Pooled sd ( $s_p$ ) in $t$ -tests & ANOVA

$s_p$  In  $t$  tests covered in Chapter 3

$$s_p = \sqrt{\frac{(n_1-1)s_1^2 + (n_2-1)s_2^2}{(n_1 + n_2 - 2)}} \quad d.f. = n_1 + n_2 - 2.$$

New  $s_p$  equation is just an extension of the  $t$ -test formula

$$SE(\bar{Y}_2 - \bar{Y}_1) = s_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$$

100(1 -  $\alpha$ )% Confidence Limits for the Difference Between Means

$$(\bar{Y}_2 - \bar{Y}_1) \pm t_{df}(1 - \alpha/2)SE(\bar{Y}_2 - \bar{Y}_1).$$

Using ANOVA, obtain a more precise  $s_p$ ,  $\sqrt{\text{within groups MS}}$ , with more  $df$  for  $p$  values & CI's

### Slide 33 Pooled sd ( $s_p$ ) in $t$ -tests & ANOVA

NOTES:

### Pooled sd, $s_p$ , for Case 5.1

$$s_p = \sqrt{\text{Error Mean Square}} = \sqrt{\text{Within Groups MS}}$$

Group	n	Sample SD
NP	49	6.1
N/N85	57	5.1
N/R50	71	7.8
R/R50	56	6.7
N/R50 lopro	56	7.0
N/R40	60	6.7

Calculate the pooled estimate of variance,  $s_p^2$

$$s_p^2 = \frac{(49-1)(6.1)^2 + (57-1)(5.1)^2 + (71-1)(7.8)^2 + (56-1)(6.7)^2 + (56-1)(7.0)^2 + (60-1)(6.7)^2}{(49-1) + (57-1) + (71-1) + (56-1) + (56-1) + (60-1)}$$

$$= \frac{15,313.90}{343} = 44.647; \quad s_p = \sqrt{44.647} = 6.68$$

$s_p$  is the square root

$df$  is the denominator

### Slide 34 Pooled sd, $s_p$ , for Case 5.1

NOTES:

① Get averages, sample sizes, and pooled estimate of standard deviation

Group	$\bar{x}$	N/R50	$\bar{x}$	N/N85
Sample size	71	57		
Average (mos.)	42.3	32.7		

Pooled estimate of  $\sigma$ :  $s_p = 6.68$  mos.;  $df = 343$  (from )

② Compute the estimate of  $\mu_3 - \mu_2$  and its standard error

$$\text{Estimate: } \bar{Y}_3 - \bar{Y}_2 = 42.3 - 32.7 = 9.6 \text{ months}$$

$$SE(\bar{Y}_3 - \bar{Y}_2) = 6.68 \sqrt{\frac{1}{71} + \frac{1}{57}} = 1.2 \text{ months}$$

③ 95% confidence interval for  $\mu_3 - \mu_2$

$$t_{343}(.975) = 1.96$$

$$95\% \text{ CI: } 9.6 \pm (1.96)(1.2) = \boxed{7.3 \text{ months}} \rightarrow 11.9 \text{ months}$$

④ Test the hypothesis that  $\mu_3 - \mu_2 = 0$

$$t\text{-stat} = \frac{9.6}{1.2} = 8.08 \rightarrow 2\text{-sided } p\text{-value} < .0001$$

Pooled estimate of standard error, from all treatments

For a priori hypotheses, the pooled sd,  $s_p$ , can be used for p values and confidence limits to compare means 2 at a time

### Slide 35

NOTES:

### ANOVA Tables from syntax

[http://www.spsstools.net/Syntax/T-Test/ANOVA\\_TablesUsing4Methods.txt](http://www.spsstools.net/Syntax/T-Test/ANOVA_TablesUsing4Methods.txt)

Summary statistics for lifetimes of mice on six different diet regimens

Group	n	Range (mo)	Average	SD	95% CI for Mean
NP	49	6.4 - 35.5	27.4	6.1	25.6 - 29.2
N/N85	57	17.9 - 42.3	32.7	5.1	31.3 - 34.1
N/R50	71	18.6 - 51.9	42.3	7.8	40.5 - 44.1
R/R50	56	24.2 - 50.7	42.9	6.7	41.1 - 44.7
N/R50 lopro	56	23.4 - 49.7	39.7	7.0	37.8 - 41.6
N/R40	60	19.6 - 54.6	45.1	6.7	43.4 - 46.8

The entire ANOVA table, including the mean square error, can be constructed from averages, sd's and n's by hand or using SPSS syntax

### Slide 36 ANOVA Tables from syntax

NOTES:

## Slide 37

Pooled estimate of standard deviation; diet restriction data

Group	n	Sample SD
NP	49	6.1
N/R85	57	5.1
N/R50	71	7.8
R/R50	56	6.7
N/R50 lopro	56	7.0
N/R40	60	6.7

$$s_p^2 = \frac{(49-1)(6.1)^2 + (57-1)(5.1)^2 + (71-1)(7.8)^2 + (56-1)(6.7)^2 + (56-1)(7.0)^2 + (60-1)(6.7)^2}{(49-1) + (57-1) + (71-1) + (56-1) + (56-1) + (60-1)}$$

$$= \frac{15,313.90}{343} = 44.647; s_p = \sqrt{44.647} = 6.68$$

$$\Rightarrow df = 343$$

Never report significance values this low!

ANOVA					
	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	12727	5	2545.4	57	5E-043
Within Groups	15314	343	44.647		
Total	28041	348			

NOTES:

## Slide 38 ANOVA from summaries

## ANOVA from summaries

See Movie in Prometheus, Microsoft Excel

Case Study 5.1

n_i	Ave_i	SD_i	n_i-1	(n_i-1)*sd_i^2	n*Ave_i	n_i*Ave_i^2
49	27.4	6.1	48	1786.08	1342.6	36787.24
57	32.7	5.1	56	1456.56	1863.9	60949.53
71	42.3	7.8	70	4258.8	3003.3	127039.59
56	42.9	6.7	55	2468.95	2402.4	103062.96
56	39.7	7	55	2695	2223.2	88261.04
60	45.1	6.7	59	2648.51	2706	122040.6

Sum 349 343 15313.9 13541.4 538140.96

Grand Mean = 38.80057

ANOVA Table

	SS	df	MS	F	p_value
Between Group	12727	5	2545.4	57	< 0.0000001
Within Group	15314	343	44.6		

s\_p 6.681836

See 'Between groups SS formula', Sleuth p 144 (Problem 19)

NOTES:

## Slide 39

DATA LIST LIST /n(F2.0) m(F5.1) sd(F4.1).

BEGIN DATA.

49 27.4 6.1

57 32.7 5.1

71 42.3 7.8

56 42.9 6.7

56 39.7 7.0

60 45.1 6.7

END DATA.

COMPUTE iv=\$CASENUM.

LOOP id=1 TO n.

XSAVE OUTFILE=XOUT1.

END LOOP.

EXECUTE.

GET FILE=XOUT1.

COMPUTE dv=.

COMPUTE k=SQR((sd\*\*2\*(N-1))/2).

IF (id=1) dv=m+k.

IF (id=2) dv=m-k.

EXECUTE.

SUMMARIZE/TABLES=dv BY iv/FORMAT=NOLIST TOTAL

/TITLE='Case Summaries'/CELLS=COUNT MEAN STDDEV

VAR.

ONEWAY dv BY iv.

ANOVA					
	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	12727	5	2545.4	57	5E-043
Within Groups	15314	343	44.647		
Total	28041	348			

The pooled sd is  $\sqrt{(\text{Within Groups Mean Square})} = \sqrt{\text{MSE}}$

This calls the SPSS's ONEWAY

NOTES:

### 5.3 The One-Way Analysis of Variance F-test (Spock data)

#### 5.3.1 Extra-Sum-of Squares principle & equal means model

- Display 5.8, Sleuth page 124

- The extra sum of squares is the single number that summarizes the difference in the sizes of residuals from the full and reduced models, p. 124

- This sum of squares, when divided by the appropriate *df*, estimates a variance
  - The F statistic assess the p-value of the equality of two variance estimates

Estimated means and residuals from two models for mean percentage of women (%W) in venters, from the Spock trial data

Subject	Full model		Reduced model	
	MEAN	RESIDUAL	MEAN	RESIDUAL
1	1.00	0.00	1.00	0.00
2	1.00	0.00	1.00	0.00
3	1.00	0.00	1.00	0.00
4	1.00	0.00	1.00	0.00
5	1.00	0.00	1.00	0.00
6	1.00	0.00	1.00	0.00
7	1.00	0.00	1.00	0.00
8	1.00	0.00	1.00	0.00
9	1.00	0.00	1.00	0.00
10	1.00	0.00	1.00	0.00
11	1.00	0.00	1.00	0.00
12	1.00	0.00	1.00	0.00
13	1.00	0.00	1.00	0.00
14	1.00	0.00	1.00	0.00
15	1.00	0.00	1.00	0.00
16	1.00	0.00	1.00	0.00
17	1.00	0.00	1.00	0.00
18	1.00	0.00	1.00	0.00
19	1.00	0.00	1.00	0.00
20	1.00	0.00	1.00	0.00
21	1.00	0.00	1.00	0.00
22	1.00	0.00	1.00	0.00
23	1.00	0.00	1.00	0.00
24	1.00	0.00	1.00	0.00
25	1.00	0.00	1.00	0.00
26	1.00	0.00	1.00	0.00
27	1.00	0.00	1.00	0.00
28	1.00	0.00	1.00	0.00
29	1.00	0.00	1.00	0.00
30	1.00	0.00	1.00	0.00
31	1.00	0.00	1.00	0.00
32	1.00	0.00	1.00	0.00
33	1.00	0.00	1.00	0.00
34	1.00	0.00	1.00	0.00
35	1.00	0.00	1.00	0.00
36	1.00	0.00	1.00	0.00
37	1.00	0.00	1.00	0.00
38	1.00	0.00	1.00	0.00
39	1.00	0.00	1.00	0.00
40	1.00	0.00	1.00	0.00
41	1.00	0.00	1.00	0.00
42	1.00	0.00	1.00	0.00
43	1.00	0.00	1.00	0.00
44	1.00	0.00	1.00	0.00
45	1.00	0.00	1.00	0.00
46	1.00	0.00	1.00	0.00
47	1.00	0.00	1.00	0.00
48	1.00	0.00	1.00	0.00
49	1.00	0.00	1.00	0.00
50	1.00	0.00	1.00	0.00

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### Slide 40 5.3 The One-Way Analysis of Variance F-test (Spock data)

NOTES:

### 'Extra sum of squares' F statistic

Sleuth Section 5.3.1

**Extra sum of squares =**

Residual sum of squares (reduced model) -  
Residual sum of squares (full model)

**F statistic =**

{(Extra sum of squares)/(Extra degrees of freedom)}

 $\sigma^2_{\text{full model}}$ 

This variance is often the 'within groups' mean square

Tested with  $F_{\{\text{Extra df, Error df full model}\}}$

### Slide 41 'Extra sum of squares' F statistic

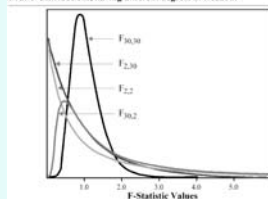
NOTES:

### F distribution

Snedecor's named the F distribution to honor Ronald Fisher

The F distribution can be regarded as the expected ratios of variances from samples drawn from the same normal distribution

Four F-distributions, having different degrees of freedom



These are probability density functions, with area 1.0. Table A.4 (p. 720) provides the area to the left of the F statistic for *df*<sub>1</sub> & *df*<sub>2</sub>

### Slide 42 F distribution

NOTES:

### ANOVA Table for Spock data

#### Partition the sum of squares

Analysis of variance table: a test for equal mean percents of women in venires of seven judges; Spock data

Source of Variation	Sum of Squares	df	Mean Square	F-Statistic	p-value
Between Groups	1,927.08	6	321.18	6.72	.000061
Within Groups	1,864.45	39	47.81		
Total	3,791.53	45			

1 Sum of squared residuals from fitting the full (separate-means) model  
 2 Sum of squared residuals from fitting the reduced (equal-means) model  
 3 degrees of freedom  
 4 Subtract the "Within" from the "Total"  
 5 A mean square is the ratio of a sum-of-squares to its degrees of freedom  
 6 The F-statistic is the ratio of the Between MS to the Within MS  
 7 The p-value comes from an F-distribution with 6 and 39 df  
 NOTE: This is  $s_p^2$

### Slide 43 ANOVA Table for Spock data

NOTES:

### SPSS ANOVA Table

#### Case 5.2 Spock trial

#### ANOVA

Percentage Women

	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	1927	6	321	6.7	.00006
Within Groups	1864	39	48		
Total	3792	45			

- Three ways to do 1-way ANOVA's in SPSS
  - Analyze\compare means\One-way ANOVA
  - Analyze\General Linear Model\Univariate
  - Analyze\Regression\Linear
- Each method has its strengths. All produce identical p values. ANOVA the simplest but least flexible

### Slide 44 SPSS ANOVA Table

NOTES:

### Case 5.2 ANOVA table, p. 130

Do as separate one-way ANOVAs, t tests with appropriate  $s_p$  for p-values or as linear contrast (next chapter)

Complete analysis of variance table for three tests involving the mean percents of women in venires of seven judges

Source of Variation	Sum of Squares	df	Mean Square	F-Statistic	p-value
Between Groups	1,927.08	6	321.18	6.72	0.000061
Spock v. Others	1,600.63	1	1,600.63	32.14	0.000001
Among Others	326.45	5	65.29	1.37	0.26
Within Groups	1,864.45	39	47.81		
Total	3,791.53	45			

Warning This ANOVA table only appropriate if 'judges' is regarded as a fixed effect, producing a fixed effect hierarchic (nested) ANOVA (Chapter 16 & Neter *et al.*) The Spock judge effect is nested within the judge effect (Between Groups)

DOOK, corrected on the Sleuth errata web site

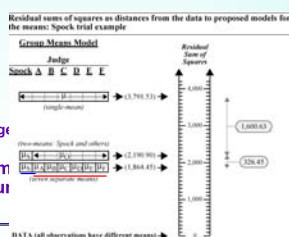
### Slide 45 Case 5.2 ANOVA table, p. 130

NOTES:

## 'Spock judge' vs. other judges

Display 5.11, page 129

- Calculate the mean for the other 6 judges (A-F)
  - Find and sum the squared residuals from that new 'other-judge' mean
    - This pooling may NOT be appropriate if there is large judge-to-judge variability
  - Then, test that residual sum of squares with an extra sum of squares F test
- 
- Residual sum of squares as distances from the data to proposed models for the judges. Speech test example
- Grand Mean Model**
- | Judge        | Speech            | Other             | Mean              | Residual  | Squared Residual    |
|--------------|-------------------|-------------------|-------------------|-----------|---------------------|
| A            | 1,791,511         | 1,791,511         | 1,791,511         | -1,000.63 | 1,000,000.00        |
| B            | 1,791,511         | 1,791,511         | 1,791,511         | 320.67    | 102,830.49          |
| C            | 1,791,511         | 1,791,511         | 1,791,511         | 1,000.63  | 1,000,000.00        |
| D            | 1,791,511         | 1,791,511         | 1,791,511         | -320.67   | 102,830.49          |
| E            | 1,791,511         | 1,791,511         | 1,791,511         | 1,000.63  | 1,000,000.00        |
| F            | 1,791,511         | 1,791,511         | 1,791,511         | -320.67   | 102,830.49          |
| <b>Total</b> | <b>10,749,066</b> | <b>10,749,066</b> | <b>10,749,066</b> | <b>0</b>  | <b>4,225,690.96</b> |
- Sum of squared residuals = 4,225,690.96



## Slide 46 'Spock judge' vs. other judges

NOTES:

## ANOVA: robustness to assumptions

- Normality is not critical. Extremely long-tailed distributions or skewed distributions, coupled with different sample sizes present the only serious distributional problems
- The assumptions of independence within and across groups is critical
- The assumption of equal standard deviations in the populations is crucial. Also called the equal variance assumption, homoscedasticity assumption (vs. Heteroscedasticity)
- The tools are **not** resistant to severely outlying observations.

## Slide 47 ANOVA: robustness to assumptions

NOTES:

## Assumptions of ANOVA

Not robust to heteroscedasticity! (But Winer et al. argue that p values are robust if sample sizes equal — Sleuth appears to have a counterargument)

Display 5.13, page 131

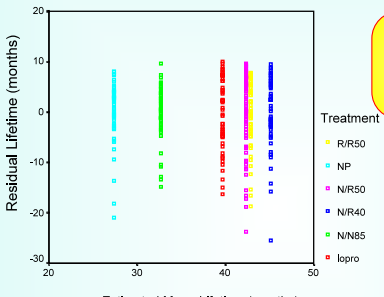
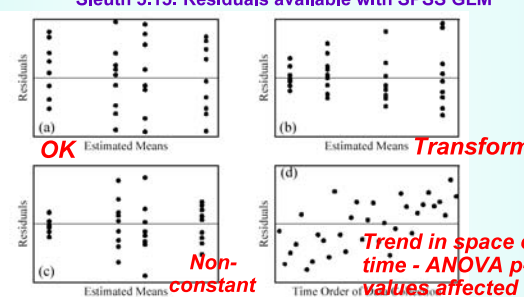
Success rates for 95% confidence intervals for  $\mu_1 - \mu_2$  from samples simulated from normal populations with possibly different SDs

			$\sigma_2=\sigma_1$			$\sigma_2=2\sigma_1$		
$n_1$	$n_2$	$n_3$	$\sigma_3=\sigma_1$	$\sigma_3=2\sigma_1$	$\sigma_3=4\sigma_1$	$\sigma_3=\sigma_1$	$\sigma_3=2\sigma_1$	$\sigma_3=4\sigma_1$
10	10	10	95.4	98.9	99.9	91.9	96.8	99.6
10	10	10	95.5	98.7	99.8	84.8	91.7	98.9
10	20	10	94.1	98.7	99.9	97.0	98.8	99.8
10	10	20	95.6	99.6	99.9	90.4	97.5	99.9

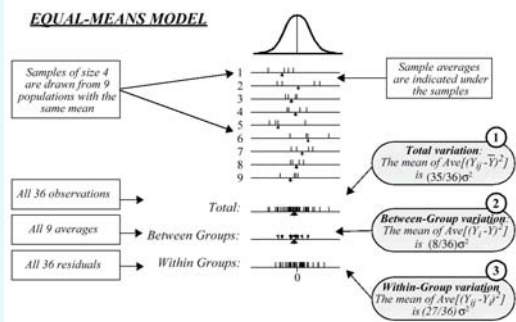
## Slide 48 Assumptions of ANOVA

NOTES:



<div data-bbox="207 132 781 554"> <h3 style="text-align: center;">Diagnostics using residuals</h3> <p style="text-align: center;">Use univariate General Linear Model</p>  <p style="text-align: right;">Examine deviations from the separate mean model</p> </div>	<div data-bbox="813 132 1398 170" style="background-color: #cccccc;">Slide 49 Diagnostics using residuals</div> <div data-bbox="813 254 1398 291">NOTES:</div>
<div data-bbox="207 590 781 1037"> <h3 style="text-align: center;">Detecting problems with residuals</h3> <p style="text-align: center;">Sleuth 5.15: Residuals available with SPSS GLM</p>  <p style="text-align: center;">(a) OK (b) Transform (c) Non-constant variance (d) Trend in space or time - ANOVA p-values affected</p> </div>	<div data-bbox="813 590 1398 630" style="background-color: #cccccc;">Slide 50 Detecting problems with residuals</div> <div data-bbox="813 714 1398 751">NOTES:</div>
<div data-bbox="207 1079 781 1535"> <h3 style="text-align: center;">5.6.1 Further illustration of different sources of variability</h3> <p style="text-align: center;">Where the Sleuth authors use graphic displays to display what those sums of squares represent and to convince you that most analyses of variance are really tests for the difference in means*</p> <p>*Random effects ANOVA: use ANOVA to test whether factors, like judges, increase variance in the response</p> </div>	<div data-bbox="813 1079 1398 1119" style="background-color: #cccccc;">Slide 51 5.6.1 Further illustration of different sources of variability</div> <div data-bbox="813 1304 1398 1341">NOTES:</div>

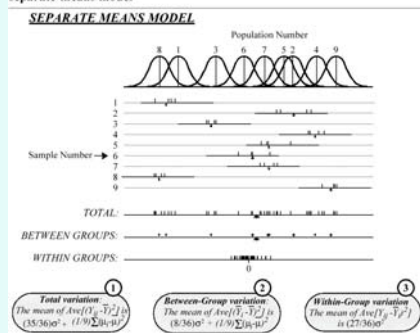
Three sources of variation for data simulated from the equal-means model, and mean values of averages of squares, from statistical theory



## Slide 52

NOTES:

Variations in the several group problem for data simulated from the separate-means model



## Slide 53

NOTES:

## 5.6.2 Kruskal-Wallis Nonparametric ANOVA

Available in SPSS Non-parametric tests  
Resistant to outliers – but susceptible to unequal variance

Spock trial data, rank-transformed

Judge	Rank of venire from smallest (1) to largest (46) percent women															
Spock's	1	2	3	4	5	6	9.5	11	16							
A	8	31	37	44	46											
B	22	26	34	36	41	45										
C	14	17	23.5	23.5	30	32.5	35	38.5	38.5							
D	19	28														
E	9.5	12	15	25	40	43										
F	7	13	18	20	21	27	29	32.5	42							

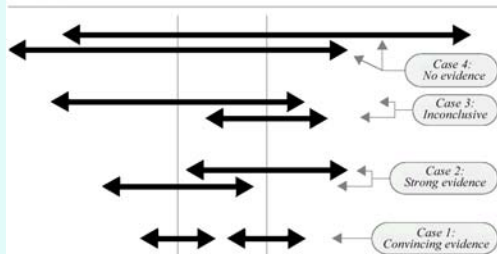
K W ANOVA does not permit analysis of any but the simplest designs

## Slide 54 5.6.2 Kruskal-Wallis Nonparametric ANOVA

NOTES:

### Confidence limits & significant differences

Separate confidence intervals for two group means: are the means different?



### Slide 55 Confidence limits & significant differences

NOTES:

### Statistical vs. Scientific significance

Always report the effect size (don't just report 'significant' or NS)

- Deming: report effect sizes for tests
- Many **statistically significant** results are trivial ecologically (or chemically or socially).
  - All null hypotheses are wrong:  $\mu_1 = \mu_2$  and the p-value is often dependent on the sample size
    - A p value of 0.00001 may not be ecologically meaningful if there is only a 1% difference in effects and at least a 5% difference causes changes in the ecosystem
- Tests with large p values may be consistent with important ecological effects
  - What is the probability of Type II error?

### Slide 56 Statistical vs. Scientific significance

NOTES:

### When is an effect 'random'?

See Sleuth Page 136-138: 'The Random Effects model'

- The differences among subgroup means is NOT of intrinsic interest.
  - You may be interested in whether the effect changes from day to day – i.e. estimating day-to-day or 'among day' variance – but you are not interested specifically in the differences on any pair of days
- If the number of levels of a factor is small relative to the total possible levels of a factor (not the case with district Judges since ALL were sampled)
- Are the subgroups a representative or random sample of some larger group?

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### Slide 57 When is an effect 'random'?

NOTES:

### Quinn & Keough (2002, p. 176) on Fixed vs. Random Factors

Random effects models allow inferences to a larger population

- Investigators use only a random subset of the possibly causal levels of a factor (or factors) and wish to make inferences to all possible levels of the factor
  - e.g., EPA selects a random subsample of zinc-contaminated streams and analyzes the data with a random-effects model
- Q & K: random or at least haphazard selection of experimental or observational units is essential



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### Slide 58 Quinn & Keough (2002, p. 176) on Fixed vs. Random Factors

NOTES:

### Comparing Spock with the other judges: Fixed or random effects?

Is the judge effect fixed or random?

- Type I ANOVA: Fixed effects ANOVA: test for differences in the averages among groups
- Type II ANOVA: Random effects ANOVA: test differences in variances due to the group classification
- Mixed model: Fixed & random factors
- Note
  - The calculations are often identical for random and fixed-effects ANOVA, but the interpretations are different
  - Factorial ANOVA (>1 factor), the F statistics differ among models, with a different denominator mean square for random factors
  - The inference allowed differs among models

### Slide 59 Comparing Spock with the other judges: Fixed or random effects?

NOTES:

### 5.17 Reproduce Display 5.9

Solution as a one-way ANOVA problem

Type I: There is at least 1 difference in the average percentage women jurors that is greater than expected by chance

Type II: There is more judge-to-judge variability in % female voters than expected by chance

Display 5.9

Analysis of variance table: a test for equal mean percents of women in verdicts of seven judges: Spock data

Source of Variation	Sum of Squares	df	Mean Square	F-Statistic	p-value
Between Groups	1,927.08	6	321.18	6.72	.000061
Within Groups	1,864.43	39	47.81		
Total	3,791.53	45			

NOTE: This is  $s_p^2$

Annotations:

- ① Subtracted the "Within" from the "Total"
- ② A mean square is the ratio of a sum of squares to its degrees of freedom
- ③ The F-Statistic is the ratio of the Between MS to the Within MS
- ④ Sum of squared residuals from fitting the full (unrestricted) means model
- ⑤ Sum of squared residuals from fitting the reduced (equal means) model
- ⑥ The p-value comes from an F distribution with 6 and 39 df
- ⑦ Degrees of freedom

### Slide 60 5.17 Reproduce Display 5.9

NOTES:

### District judges: Random?

If the judge effect is a random factor, this design is pseudoreplicated and invalid. But, the judges are NOT a random subset of a larger class of judges. These 7 judges represent all of the judges. The model is a fixed effect design

Complete analysis of variance table for three tests involving the mean percents of women in venires of seven judges

Source of Variation	Sum of Squares	df	Mean Square	F-Statistic	p-value
Between Groups	1,927.08	6	321.18	6.72	0.000061
Spock v. Others	1,600.63	1	1,600.63	32.14	0.000001
Among Others	326.45	5	65.29	1.37	0.26
Within Groups	1,864.45	39	47.81		
Total	3,791.53	45			

ANOVA uses an inappropriate denominator mean square for the Spock judge effect

### Slide 61 District judges: Random?

NOTES:

### Fixed vs. Random effects

Underwood (1997): Fixed effects 1-way ANOVA

$$X_{ij} = \mu + A_i + e_{ij}$$

where  $X_{ij}$  is  $j$ th replicate in  $i$ th treatment ( $i$ th level of factor  $A$ ;  $i = 1 \dots a$ ).

$A_i$  is difference between  $i$ th level of factor  $A$  and overall mean of all levels ( $\mu$ ),  $e_{ij}$  is the deviation of replicate  $j$  in  $i$ th sample from the mean of that population.

Fixed factor:

By definition:

$$\sum_{i=1}^a A_i = 0$$

(see Section 7.6).

Among treatments

Within treatments

where  $k_A^2$  indicates fixed differences, all sampled in the experiment.

$$\frac{n \sum_{i=1}^a (A_i - \bar{A})^2}{(a-1)} \text{ or } \sigma_e^2 + nk_A^2$$

### Slide 62 Fixed vs. Random effects

NOTES:

### Fixed vs. Random factors

Underwood (1997): Random factor (Model II) 1-way ANOVA

Random factor:

$$E\left(\sum_{i=1}^a A_i\right) = 0$$

Meaning you expect  $\sum_{i=1}^a A_i = 0$  on average, over many experiments, but in a single experiment,  $A_i$  values as sampled may not sum to zero.

Analysis of variance	Mean square estimates
Among treatments	$\sigma_e^2 + n\sigma_A^2$
Within treatments	$\sigma_e^2$

where  $\sigma_A^2$  is the variance of the population of  $A_i$  values sampled in your experiment.

### Slide 63 Fixed vs. Random factors

NOTES:

## Factorial ANOVA

Tables from Underwood (will be covered in Sleuth 4.4.5.2)

(a) Both factors fixed				
Source of variation	Sum of squares	Degrees of freedom	Mean square estimates	F-ratio versus
Among levels of A = A	$(a-1)\sigma_A^2 + b\sum_{j=1}^b (\bar{X}_{.j} - \bar{X})^2$	$a-1$	$\sigma_A^2 + b\sigma_B^2$	Residual
Among levels of B = B	$(b-1)\sigma_B^2 + a\sum_{j=1}^a (\bar{X}_{.j} - \bar{X})^2$	$b-1$	$\sigma_B^2 + a\sigma_A^2$	Residual
A x B	$(a-1)(b-1)\sigma_{AB}^2 + \sum_{j=1}^b \sum_{k=1}^a (\bar{X}_{jk} - \bar{X}_{.j} - \bar{X}_{.k} + \bar{X})^2$	$(a-1)(b-1)$	$\sigma_{AB}^2 + a\sigma_A^2$	Residual
Residual	$ab(a-1)\sigma^2$	$ab(a-1)$	$\sigma^2$	
(b) A fixed, B random				
Source of variation	Sum of squares	Degrees of freedom	Mean square estimates	F-ratio versus
Among levels of A = A	$(a-1)\sigma_A^2 + (a-1)b\sigma_{AB}^2 + b\sum_{j=1}^b (\bar{X}_{.j} - \bar{X})^2$	$a-1$	$\sigma_A^2 + b\sigma_{AB}^2 + b\sigma^2$	$\frac{A = A}{A = B}$
Among levels of B = B	$(b-1)\sigma_B^2 + (b-1)a\sigma_{AB}^2$	$b-1$	$\sigma_B^2 + a\sigma_{AB}^2$	Residual
A x B	$(a-1)(b-1)\sigma_{AB}^2 + (a-1)(b-1)a\sigma^2$	$(a-1)(b-1)$	$\sigma_{AB}^2 + a\sigma^2$	Residual
Residual	$ab(a-1)\sigma^2$	$ab(a-1)$	$\sigma^2$	
(c) Both factors random				
Source of variation	Sum of squares	Degrees of freedom	Mean square estimates	F-ratio versus
Among levels of A = A	$(a-1)\sigma_A^2 + (a-1)b\sigma_{AB}^2 + b\sum_{j=1}^b (\bar{X}_{.j} - \bar{X})^2$	$a-1$	$\sigma_A^2 + b\sigma_{AB}^2 + b\sigma^2$	$\frac{A = A}{A = B}$
Among levels of B = B	$(b-1)\sigma_B^2 + (b-1)a\sigma_{AB}^2$	$b-1$	$\sigma_B^2 + a\sigma_{AB}^2$	$\frac{A = B}{A = B}$
A x B	$(a-1)(b-1)\sigma_{AB}^2 + (a-1)(b-1)a\sigma^2$	$(a-1)(b-1)$	$\sigma_{AB}^2 + a\sigma^2$	Residual
Residual	$ab(a-1)\sigma^2$	$ab(a-1)$	$\sigma^2$	

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## Slide 64 Factorial ANOVA

NOTES:

## Mixed Model Nested ANOVA

A 1 in 67 chance of observing such a difference by chance

Tests of Between-Subjects Effects

Dependent Variable: Percentage Women

Source		Type III Sum of Squares	df	Mean Square	F	Sig.
Intercept	Hypothesis	20003	1	20003	293	
	Error	273	4,003	68		
SPOCK	Hypothesis	1537	1	1537	21.6	.015
	Error	236	3,311	71 <sup>b</sup>		
CODE(SPOCK)	Hypothesis	326	5	65	1.4	.258
	Error	1864	39	48 <sup>c</sup>		

a. 1.167 MS(CODE(SPOCK)) - .167 MS(Error)

b. 1.337 MS(CODE(SPOCK)) - .337 MS(Error)

c. MS(Error)

This model is not appropriate because the judges are not a random subset of judges

UNIANOVA  
percent BY spock code  
/METHOD = SSTYPE(3)  
/INTERCEPT = INCLUDE  
/CRITERIA = ALPHA(.05)  
/RANDOM CODE  
/DESIGN = spock code(spock) .

## Slide 65 Mixed Model Nested ANOVA

NOTES:

## Counterfactual conditionals

*Modus tollens*

Hypothesis:

If A then B

Observe 'Not B'

Then conclude: 'Not A' [Reject null]

**Counterfactual conditional**

If A and C then B

Not C

No inference possible about the truth or falsity of A can be inferred from observing either 'B' or 'Not B'

*Modus tollens*

If the Spock judge's venire were due to chance, then  $F=1.0$   
Observe  $F=22$  ( $p=0.015$ ,  $F_{1,3.3}$ )

**Counterfactual conditional**  
If the juries were chosen by chance and the judges were a random subset of judges AND we know that the judges were NOT a random subset of judges. Then  $F=1$   
Observing  $F>1$  doesn't allow us to conclude anything about the fairness of the jury selection

## Slide 66 Counterfactual conditionals

NOTES:

<div style="text-align: center;"> <b>Conclusions</b>  (1 of 3) </div> <ul style="list-style-type: none"> <li>ANOVA tables can be created from summary statistics</li> <li>Assumptions: <ul style="list-style-type: none"> <li>Homoscedasticity <ul style="list-style-type: none"> <li>Levene's test a rough guide</li> <li>Boxplots or residual plots are the standard tools for assessing homoscedasticity (equal variance among groups)</li> <li>Spread vs. Level plots</li> </ul> </li> <li>Independence of errors among groups a key ANOVA assumption</li> <li>Normally distributed errors (not underlying data) not crucial</li> </ul> </li> </ul> <p style="text-align: right;"><b>EEOS611</b></p>	<div style="background-color: #cccccc; text-align: center;"> <b>Slide 67 Conclusions</b> </div> <div style="border: 1px solid black; height: 100px; margin-top: 10px;"></div>
<div style="text-align: center;"> <b>Conclusions</b>  (2 of 3) </div> <ul style="list-style-type: none"> <li>An ANOVA is more efficient &amp; powerful than multiple, separate <math>t</math> tests <ul style="list-style-type: none"> <li>The ANOVA error MS (=within groups MS) provides a more <b>precise</b> estimate of the population standard deviation [It is not a smaller estimate of error {it is an unbiased estimator}]</li> </ul> </li> <li>Kruskal-Wallis ANOVA is the rank-based analogue of 1-way ANOVA and is resistant to outliers but not unequal spread <ul style="list-style-type: none"> <li>Ties correction must be used</li> <li>Effect sizes, hierarchic structure, and covariates difficult to handle</li> </ul> </li> </ul>	<div style="background-color: #cccccc; text-align: center;"> <b>Slide 68 Conclusions</b> </div> <div style="border: 1px solid black; height: 100px; margin-top: 10px;"></div>
<div style="text-align: center;"> <b>Conclusions</b>  (3 of 3) </div> <ul style="list-style-type: none"> <li>ANOVA tests for difference in means (fixed effect) or whether <math>\sigma_i^2 = 0</math> (random effect) or both (mixed model)</li> <li>Fixed vs. random effects <ul style="list-style-type: none"> <li>The choice of fixed vs. random effects is often crucial and depends on whether the factor levels (judges in the Spock example) represent a random or representative sample from some larger statistical population</li> <li>The <math>F</math> statistics and interpretation of the results sometimes change depending on whether fixed or random effects are chosen</li> </ul> </li> </ul> <p style="text-align: right;"><b>EEOS611</b></p>	<div style="background-color: #cccccc; text-align: center;"> <b>Slide 69 Conclusions</b> </div> <div style="border: 1px solid black; height: 100px; margin-top: 10px;"></div>