


Hypothesis Test: Comparing Multiple Groups (ANOVA)


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Review

- ◆ One-sample hypothesis test:
 - $H_0: \mu = \text{constant}$, $H_1: \mu \neq \text{constant}$
 - $H_0: \pi = \text{constant}$, $H_1: \pi \neq \text{constant}$
- ◆ Two-sample hypothesis test:
 - $H_0: \mu_1 = \mu_2$, $H_1: \mu_1 \neq \mu_2$
 - $H_0: \pi_1 = \pi_2$, $H_1: \pi_1 \neq \pi_2$
 - Dependent samples $H_0: \mu_D = 0$
- ◆ One-tailed vs. two-tailed tests


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Issues

- ◆ What if we have more than two groups?
 - different ethnic groups;
 - difference classes in a school;
 - multiple years of data
- ◆ H_0 : All groups are identical
 - E.g. $\mu_1 = \mu_2 = \mu_3 = \mu_4$
- ◆ H_1 : One or more groups differ


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Option #1

- ◆ Two-sample t-test for every combination of groups
 - $\mu_1 = \mu_2$, $\mu_1 = \mu_3$, $\mu_1 = \mu_4$, $\mu_2 = \mu_3$, $\mu_2 = \mu_4$, $\mu_3 = \mu_4$
- ◆ But, the possibility of a Type I error proliferates... 5% for each test.
 - With only 4 groups, 6 two-sample tests, chance of error reaches $6 * 5\% = 30\%$

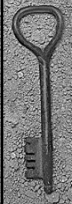
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Option #2: ANOVA

- ◆ ANOVA = “ANalysis Of VAriance”
 - “Oneway ANOVA” : The simplest form
- ◆ Only one test is needed, test whether all groups are the same ($\mu_1 = \mu_2 = \mu_3 = \mu_4$)
 - But, doesn’t distinguish which specific group(s) differ
 - Maybe only μ_2 differs, or maybe all differ from others


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ANOVA: Example

- ◆ Suppose you suspect that a firm is engaging in wage discrimination based on ethnicity
 - Certain ethnic groups might be getting paid more...
- ◆ The company counters: “We pay entry-level workers all **about** the same amount of money. No group gets preferential treatment.”
- ◆ Given data on a sample of employees, ANOVA lets you test this hypothesis.
 - Are observed group differences just due to chance?
 - Or do they reflect differences in the underlying population? (i.e., the whole company)

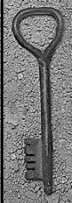
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ANOVA: Example

- ◆ The company has workers of three ethnic groups:
 - Whites, African-Americans, Asian-Americans
- ◆ Based on a sample of workers:
 - $Y\text{-bar}_{\text{White}} = \$8.78 / \text{hour}$
 - $Y\text{-bar}_{\text{AfAm}} = \$8.52 / \text{hour}$
 - $Y\text{-bar}_{\text{AsianAm}} = \$8.91 / \text{hour}$
- ◆ What can we conclude?
 - Nothing! Sample means differ randomly even if all groups had the same population mean ($\mu_{\text{White}} = \mu_{\text{AfAm}} = \mu_{\text{AsianAm}}$).
- ◆ Q: Are the observed differences so large it is unlikely that they are due to random error?
 - Thus, it is unlikely that: $\mu_{\text{White}} = \mu_{\text{AfAm}} = \mu_{\text{AsianAm}}$

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ANOVA: Concepts & Definitions

- ◆ Previously: μ , $Y\text{-bar}$; μ_1, μ_2 , $Y\text{-bar}_1, Y\text{-bar}_2$
- ◆ The **grand mean** is the mean of all groups/cases
 - ex: mean of all entry-level workers = \$8.70/hour
- ◆ The **group mean** is the mean of a particular subgroup of the population
- ◆ We hope to make inferences about **population** grand mean and group means, even though we only have **sample** grand mean and group means
 - We know $Y\text{-bar}$, $Y\text{-bar}_{\text{White}}$, $Y\text{-bar}_{\text{AfAm}}$, $Y\text{-bar}_{\text{AsianAm}}$
 - We want to infer about: μ , μ_{White} , μ_{AfAm} , μ_{AsianAm}

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ANOVA: Concepts & Definitions

- ◆ Recall: variance, standard deviation are based on deviations, which is the distance of a point from the grand mean: $d_i = Y_i - \bar{Y}$
- ◆ ANOVA is based on partitioning deviation into different components

Grand Mean Group Mean Y₁

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ANOVA: Concepts & Definitions

- ◆ The deviation of **any case** is determined by:
 - the distance between a group mean and the grand mean: the “group effect” (α), common to group members
 - the distance from group mean to a case’s value: the within-group deviation (e): called “error”,

Grand Mean Group Mean Y₁

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ANOVA: Concepts & Definitions

- ◆ Initially we calculated deviation as the distance of a point from the grand mean: $d_i = Y_i - \bar{Y}$
- ◆ The total deviation can be partitioned into α_j (group effect) and e_{ij} (case errors, case i in group j)

$$d_{ij} = \alpha_j + e_{ij} = (\bar{Y}_j - \bar{Y}) + (Y_{ij} - \bar{Y}_j) = Y_{ij} - \bar{Y}$$

Grand Mean Group Mean Y₁


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Sum of Squared Deviation

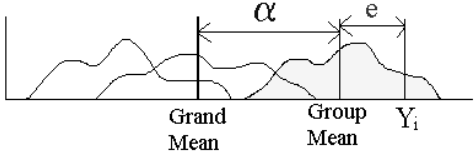
- ◆ The group effects: α_j $\alpha_j = \bar{Y}_j - \bar{Y}$
 - Deviation of the group from the grand mean
- ◆ Individual case error: e_{ij} $e_{ij} = Y_{ij} - \bar{Y}_j$
 - Deviation of the individual from the group mean
- ◆ Each are deviations that can be squared, and “summed up” -> sum of squared deviation
 - Recall: variance is sum of squared deviation

$$s_Y^2 = \frac{\sum_{i=1}^n d_i^2}{n-1} = \frac{\sum_{i=1}^n (Y_i - \bar{Y})^2}{n-1}$$

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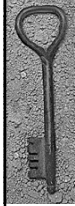


Sum of Squared Deviation



- The total variance (SS_{total}) is made up of:
 - **between group** variance ($SS_{between}$)
 - **within group** variance (SS_{within})
 - $SS_{total} = SS_{between} + SS_{within}$

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Sum of Squared Deviation

- Given a sample with j sub-groups:
 - Grand Mean = \bar{Y}
 - Group Means = $\bar{Y}_1, \bar{Y}_2, \bar{Y}_3, \dots, \bar{Y}_j$
- “Total Sum of Squares” (SS_{total})

$$SS_{total} = \sum_{i=1}^N (Y_i - \bar{Y})^2 = \sum_{j=1}^J \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y})^2$$

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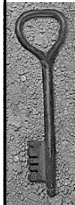
Sum of Squared Deviation

- The **between group** variance is the distance from the grand mean to each group mean (summed for all cases):

$$SS_{Between} = \sum_{j=1}^J n_j (\bar{Y}_j - \bar{Y})^2$$
- The **within group** variance is the distance from each case to its group mean (summed):

$$SS_{Within} = \sum_{j=1}^J \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_j)^2$$

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Sum of Squared Variance

- The sum of squares grows as n gets larger.
 - To derive a more comparable measure, we “average” it, just as with the variance: i.e., divided by n-1

$$s_y^2 = \frac{\sum_{i=1}^n (Y_i - \bar{Y})^2}{n - 1}$$

- For similar reasons, it is desirable to “average” the between/within Sum of Squares
- Result the “Mean Square” variance
 - $MS_{between}$ and MS_{within}

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Sum of Squared Variance

- ◆ Divide Sum of Squares by degree of freedom:

$$MS_{\text{Between}} = \frac{\sum_{j=1}^J n_j (\bar{Y}_j - \bar{Y})^2}{J - 1} \quad (\text{df}=J-1)$$

$$MS_{\text{Within}} = \frac{\sum_{j=1}^J \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_j)^2}{N - J}$$

(df=(n₁-1)+(n₂-1)+ ... + (n_j-1)=N-J)

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Mean Squares and Group Differences

- ◆ Q: Which suggests that group means are quite different?

$MS_{\text{between}} > MS_{\text{within}}$ or $MS_{\text{between}} < MS_{\text{within}}$

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Mean Squares and Group Differences

$MS_{\text{between}} > MS_{\text{within}}$:

$MS_{\text{between}} < MS_{\text{within}}$:

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Mean Squares and Group Differences

- ◆ Q: Which suggests that group means are quite different:

 - $MS_{\text{between}} > MS_{\text{within}}$ or $MS_{\text{between}} < MS_{\text{within}}$

- ◆ Answer: If between group variance is greater than within, the groups are quite distinct
 - It is unlikely that they came from a population with the same mean
- ◆ If within is greater than between, the groups aren't very different – they overlap a lot
 - It is plausible that $\mu_1 = \mu_2 = \mu_3 = \mu_4$

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The F Ratio

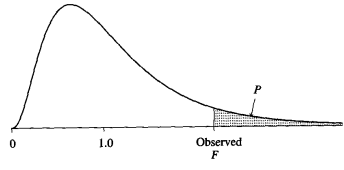
$$F_{J-1, N-J} = \frac{MS_{Between}}{MS_{Within}}$$

- If $MS_{between} > MS_{within}$ then $F > 1$
- If $MS_{between} < MS_{within}$ then $F < 1$
- Larger F indicates that groups are more separate

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The F Ratio

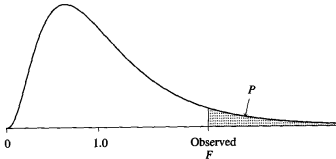
- ◆ The F ratio has a sampling distribution (F-distribution)
- ◆ Again, this sampling distribution has known properties that can be looked up in a table
 - So, we can test hypotheses



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F- Distribution

- ◆ Assume only positive values
- ◆ Skewed to the right
- ◆ Shape is determined by two degrees of freedom:
 - J-1, one for number of groups
 - N-J, one for total sample size N



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The F-test

- ◆ Assumptions required for hypothesis testing using an F-statistic
 - Population distributions for groups are normal
 - Variance for population groups are equal
 - Independent random samples from population groups

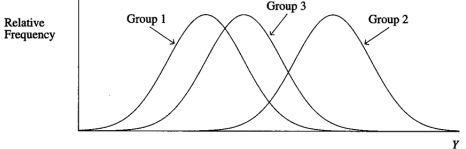



Figure 12.1 Assumptions About Population Distributions: Normal with Equal Standard Deviations

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The F-test

- If these assumptions hold, the F statistic can be looked up in an F-distribution table

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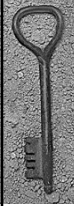
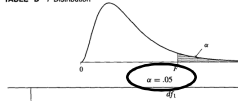


TABLE D. F Distribution




d_f1	1	2	3	4	5	6	8	12	24	∞
1	161.4	199.5	215.7	224.5	230.2	234.0	238.9	243.9	248.9	254.1
2	18.51	19.00	19.16	19.25	19.30	19.33	19.37	19.41	19.45	19.50
3	10.13	9.55	9.28	9.12	9.01	8.94	8.84	8.74	8.64	8.53
4	7.71	6.94	6.59	6.29	6.20	6.16	6.04	5.91	5.77	5.63
5	6.61	5.79	5.41	5.19	5.05	4.95	4.82	4.68	4.53	4.36
6	5.95	5.14	4.76	4.53	4.29	4.28	4.15	4.00	3.84	3.67
8	5.12	4.46	4.07	3.84	3.69	3.58	3.44	3.28	3.12	2.91
10	4.52	4.26	3.86	3.63	3.48	3.37	3.23	3.07	2.90	2.71
12	4.06	4.10	3.71	3.48	3.33	3.22	3.07	2.91	2.74	2.54
14	3.71	3.88	3.49	3.26	3.11	3.00	2.85	2.69	2.50	2.30
16	3.42	3.60	3.21	2.98	2.83	2.72	2.57	2.40	2.21	2.01
18	3.21	3.39	3.00	2.77	2.62	2.51	2.36	2.19	2.00	1.80
20	3.04	3.22	2.83	2.60	2.45	2.34	2.19	2.02	1.83	1.63
25	2.70	2.88	2.49	2.26	2.11	2.00	1.85	1.68	1.49	1.29
30	2.50	2.68	2.29	2.06	1.91	1.80	1.65	1.48	1.29	1.09
40	2.20	2.38	1.99	1.76	1.61	1.50	1.35	1.18	0.99	0.79
50	2.05	2.23	1.84	1.61	1.46	1.35	1.20	1.03	0.84	0.64
60	1.95	2.13	1.74	1.51	1.36	1.25	1.10	0.93	0.74	0.54
80	1.80	2.00	1.61	1.38	1.23	1.12	0.97	0.80	0.61	0.41
100	1.70	1.90	1.51	1.28	1.13	1.02	0.87	0.70	0.51	0.31
120	1.65	1.85	1.46	1.23	1.08	0.97	0.82	0.65	0.46	0.26
140	1.60	1.80	1.41	1.18	1.03	0.92	0.77	0.60	0.41	0.21
160	1.57	1.77	1.38	1.15	1.00	0.89	0.74	0.57	0.38	0.18
180	1.55	1.75	1.36	1.13	0.98	0.87	0.72	0.55	0.36	0.16
200	1.53	1.73	1.34	1.11	0.96	0.85	0.70	0.53	0.34	0.14
250	1.49	1.69	1.30	1.07	0.92	0.81	0.66	0.49	0.30	0.10
300	1.46	1.66	1.27	1.04	0.89	0.78	0.63	0.46	0.27	0.07
400	1.42	1.62	1.23	1.00	0.85	0.74	0.59	0.42	0.23	0.03
500	1.39	1.59	1.20	0.97	0.82	0.71	0.56	0.39	0.20	0.01
600	1.37	1.57	1.18	0.95	0.80	0.69	0.54	0.37	0.18	0.00
800	1.34	1.54	1.15	0.92	0.77	0.66	0.51	0.34	0.15	0.00
1000	1.32	1.52	1.13	0.90	0.75	0.64	0.49	0.32	0.13	0.00

Source: From Table V of R. A. Fisher and F. Yates, *Statistical Tables for Biological, Agricultural and Medical Research*, published by Longman Group Ltd., London, 1974. (Previously published by Oliver & Boyd, Edinburgh.) Reprinted by permission of the authors and publishers.

One table for each significance level

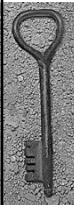
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Example

- Wage discrimination in a firm, 3 groups of workers
 - Whites, African-Americans, Asian-Americans
- You observe in a sample of 300 employees:
 - $n_1=100, Y\text{-bar}_{\text{White}} = \$8.78/\text{hour}, s_1=1.5$
 - $n_2=100, Y\text{-bar}_{\text{AfAm}} = \$8.52/\text{hour}, s_2=1.2$
 - $n_3=100, Y\text{-bar}_{\text{AsianAm}} = \$8.91/\text{hour}, s_3=0.9$
 - $Y\text{-bar} = \$8.74/\text{hour}$ (grand mean)

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Example

- Assumption
 - Wage for each group is normally distributed
 - Assume same variance for each group
 - Independent random sample from each ethnic group
- $H_0: \mu_{\text{white}} = \mu_{\text{AfAm}} = \mu_{\text{AsianAm}}$
 $H_1: \text{one or more group mean is different}$
- Calculate F-statistic

$$F_{J-1, N-J} = \frac{MS_{\text{Between}}}{MS_{\text{Within}}}$$

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Calculating Mean Sum of Squares

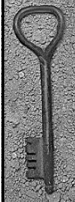


$$MS_{\text{Between}} = \frac{\sum_{j=1}^J n_j (\bar{Y}_j - \bar{Y})^2}{J - 1}$$

$$MS_{\text{Between}} = \frac{100(8.78 - 8.74)^2 + 100(8.52 - 8.74)^2 + 100(8.91 - 8.74)^2}{3 - 1} = 3.95$$

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Calculating Mean Sum of Squares



$$MS_{\text{Within}} = \frac{\sum_{j=1}^J \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_j)^2}{N - J} = \frac{\sum_{j=1}^J (n_j - 1) S_j^2}{N - J}$$


S: Group standard deviation

(Because $s_j^2 = \frac{\sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_j)^2}{n_j - 1}$)

$$MS_{\text{Within}} = \frac{(100 - 1)1.5^2 + (100 - 1)1.2^2 + (100 - 1)0.9^2}{300 - 3} = 1.5$$

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Example

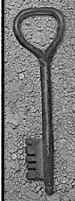
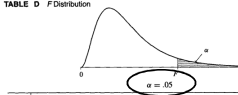


$$F_{J-1, N-J} = \frac{MS_{\text{Between}}}{MS_{\text{Within}}} = \frac{3.95}{1.5} = 2.63$$

- Recall that $N = 300, J = 3$
 - $df_1 = J - 1 = 2, df_2 = N - J = 297$
- If $\alpha = .05$, the critical F value=?

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TABLE D F Distribution

df ₁	1	2	3	4	5	6	8	12	24	∞
1	161.4	199.5	225.7	242.4	255.2	265.8	281.9	305.9	342.2	385.6
2	18.51	19.00	19.16	19.25	19.30	19.33	19.37	19.41	19.45	19.50
3	10.13	9.55	9.28	9.12	9.01	8.94	8.84	8.74	8.64	8.53
4	7.71	6.94	6.59	6.39	6.26	6.16	6.04	5.91	5.77	5.63
5	6.61	5.79	5.41	5.19	5.05	4.95	4.82	4.68	4.53	4.36
6	5.96	5.14	4.76	4.53	4.39	4.28	4.15	4.00	3.84	3.67
7	5.59	4.74	4.35	4.12	3.97	3.87	3.73	3.57	3.41	3.23
8	5.32	4.46	4.07	3.84	3.69	3.58	3.44	3.28	3.12	2.93
9	5.12	4.26	3.86	3.63	3.48	3.37	3.23	3.07	2.90	2.71
10	4.96	4.10	3.71	3.48	3.33	3.22	3.07	2.91	2.74	2.54
11	4.84	3.98	3.59	3.36	3.20	3.09	2.95	2.79	2.61	2.40
12	4.75	3.88	3.49	3.26	3.11	3.00	2.85	2.69	2.50	2.29
15	4.67	3.80	3.41	3.18	3.02	2.91	2.76	2.60	2.41	2.21
16	4.66	3.78	3.39	3.16	3.01	2.90	2.75	2.59	2.40	2.20
17	4.65	3.78	3.39	3.16	3.01	2.90	2.75	2.59	2.40	2.20
18	4.64	3.77	3.38	3.15	2.99	2.88	2.73	2.57	2.38	2.18
19	4.64	3.77	3.38	3.15	2.99	2.88	2.73	2.57	2.38	2.18
20	4.63	3.76	3.37	3.14	2.98	2.87	2.72	2.56	2.37	2.17
21	4.63	3.76	3.37	3.14	2.98	2.87	2.72	2.56	2.37	2.17
22	4.62	3.75	3.36	3.13	2.97	2.86	2.71	2.55	2.36	2.16
23	4.62	3.75	3.36	3.13	2.97	2.86	2.71	2.55	2.36	2.16
24	4.61	3.74	3.35	3.12	2.96	2.85	2.70	2.54	2.35	2.15
25	4.61	3.74	3.35	3.12	2.96	2.85	2.70	2.54	2.35	2.15
26	4.60	3.73	3.34	3.11	2.95	2.84	2.69	2.53	2.34	2.14
27	4.60	3.73	3.34	3.11	2.95	2.84	2.69	2.53	2.34	2.14
28	4.59	3.72	3.33	3.10	2.94	2.83	2.68	2.52	2.33	2.13
29	4.59	3.72	3.33	3.10	2.94	2.83	2.68	2.52	2.33	2.13
30	4.58	3.71	3.32	3.09	2.93	2.82	2.67	2.51	2.32	2.12
40	4.58	3.71	3.32	3.09	2.93	2.82	2.67	2.51	2.32	2.12
60	4.57	3.70	3.31	3.08	2.92	2.81	2.66	2.50	2.31	2.11
120	4.56	3.69	3.30	3.07	2.91	2.80	2.65	2.49	2.30	2.10
∞	4.55	3.68	3.29	3.06	2.90	2.79	2.64	2.48	2.29	2.09

Source: From Table V of R. A. Fisher and F. Yates, *Statistical Tables for Biological, Agricultural and Medical Research*, published by Longman Group Ltd., London, 1974. (Previously published by Oliver & Boyd, Edinburgh.) Reprinted by permission of the authors and publishers.

$df_1=2$
 $df_2=297$

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Example

$$F_{J-1, N-J} = \frac{MS_{Between}}{MS_{Within}} = \frac{3.95}{1.5} = 2.63$$

- ◆ Recall that $N = 300, J = 3$
 - $df_1 = J - 1 = 2, df_2 = N - J = 297$
- ◆ 5. If $\alpha = .05$, the critical F value for 2, 297 is about 3.00
- ◆ 6. Conclusion: the observed $F < 3$, so we fail to reject H_0 ; we can conclude that the groups have the same population mean \rightarrow no racial discrimination in wage

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Summary

- ◆ Concepts:
 - Grand vs. group means
 - Between vs. within group sum of squared deviation (variance) ($SS_{between} / SS_{within}$)
 - Between vs. within group mean squares ($MS_{between} / MS_{within}$)
- ◆ ANOVA (F-test)
 - Assumptions
 - H_0 : all group means are the same
 - H_1 : one or more group means are different
 - F statistic: $F = MS_{between} / MS_{within}$
 - Critical value from F-distribution table, $df_1 = J - 1, df_2 = N - J$
 - Conclusion: if $F > C.V.$, reject H_0 ; if $F < C.V.$, fail to reject H_0

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Conducting ANOVA in SPSS

The screenshot shows the SPSS One-Way ANOVA dialog box. The dependent list contains 'PainLevel' and the factor list contains 'Doseage'. The 'Display' section has 'Display Descriptives' checked. The 'Display' section also has 'Display Homogeneity of Variance Test' checked. The 'Display' section also has 'Display Residuals' checked. The 'Display' section also has 'Display Unadjusted Residuals' checked. The 'Display' section also has 'Display Unadjusted Total' checked. The 'Display' section also has 'Display Unadjusted Error' checked. The 'Display' section also has 'Display Unadjusted Total Error' checked. The 'Display' section also has 'Display Unadjusted Total Error' checked.

Output

Descriptives

PainLevel	N	Mean	Std. Deviation	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
1	3	8.33	.577	.333	6.90	9.77	8	9
2	3	5.00	1.000	.577	2.52	7.48	4	6
3	4	2.25	.957	.479	.73	3.77	1	3
Total	10	4.90	2.767	.875	2.92	6.88	1	9

ANOVA

PainLevel	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	63.483	2	31.742	41.020	.000
Within Groups	5.417	7	.774		
Total	68.900	9			

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