Chi-Square Test of Independence (based on a Contingency Table)

$$\chi^{2} = \sum \frac{(Observed - E \text{ xp ected})^{2}}{Expected}$$

$$df = (r-1)(c-1)$$

In the following contingency table estimate the proportion of drinkers among those who develop Lung Cancer and those without the disease?

		Lung Cancer		Total
		Case	Control	
Drinker	Yes	O11=33	O12=27	R1=60
	No	O21=1667	O22 = 2273	R2=3940

Total

$$C1 = 1700$$

$$C2 = 2300 \quad n = 4000$$

E11=1700(60)/4000=25.5 E12=34.5 E21=1674.5 E22=2265.5

$$\chi_{obs}^{2} \sum_{k=1}^{k=4} \frac{(Observed - E \text{ xp ected})^{2}}{Expected} = \frac{(33 - 25.5)^{2}}{25.5} + \frac{(27 - 34.5)^{2}}{34.5} + \frac{(1667 - 1674.5)^{2}}{1674.5} + \frac{(2273 - 2265.5)^{2}}{2265.5} = 4.0$$

- H_0 : $\mu_1 = \mu_2 = ... = \mu_k$
- Say we reject H₀... what now?
- Need additional tests to find where the differences lie.
- One approach: series of (k choose 2) two-sample t-tests. → increased probability of making Type I error.
- Solution: be more conservative in the individual comparisons.

- As the number of tests \spadesuit , individual $\alpha \Psi$
- <u>Bonferroni correction</u>: to set overall Type I error probability at 0.05,
 - $\alpha^* = 0.05/(k \text{ choose 2})$ for indiv. comparisons
- k=3, total of (3 choose 2)=3 tests
 - $\rightarrow \alpha^* = 0.05/3 = 0.0167$

- H_0 : $\mu_i = \mu_j$
- Calculate $t_{ij} = \frac{\overline{x}_i \overline{x}_j}{sqrt(s_w^2(1/n_i + 1/n_i))}$
- Note that we use all the additional info from the k samples to estimate the common variance σ^2 .
- Under H_0 , t_{ii} has a t-distribution with n k df.
- One drawback of Bonferroni correction: highly conservative, lacks statistical power.

- Example: Follow 3 groups of overweight males for 1 year.
- Group 1: diet, no exercise program
- Group 2: regular exercise, no diet
- Group 3: control, neither diet nor exercise.
- After 1 year, total change in body weight measured for each individual.

	Group 1	Group 2	Group 3	
n _i	42	47	42	
$\overline{X_i}$	-7.2	-4.0	0.6	in kg
Si	3.7	3.9	3.7	in kg

- Assume independent, normally distributed data.
- Null hypothesis H_0 : $\mu_1 = \mu_2 = \mu_3$
- Assume underlying population variances are equal (looks like a good assumption).

First, estimate within-groups variance:

$$s_w^2 = \frac{(n_1 - 1)s_1^2 + (n_2 - 1)s_2^2 + (n_3 - 1)s_3^2}{n_1 + n_2 + n_3 - 3}$$
$$= 14.24 \text{ kg}^2$$

Next, the grand mean of the data:

$$\overline{x} = n_1 \overline{x}_1 + n_2 \overline{x}_2 + n_3 \overline{x}_3$$

$$= n_1 + n_2 + n_3$$

$$= -3.55 \text{ kg}$$

• Then, estimate the between-groups variance:

$$s_{B}^{2} = \frac{n_{1}(\overline{x}_{1} - \overline{x})^{2} + n_{2}(\overline{x}_{2} - \overline{x})^{2} + n_{3}(\overline{x}_{3} - \overline{x})^{2}}{3 - 1}$$
$$= 646.20 \text{ kg}^{2}$$

• So, the test statistic is:

$$F = s_B^2/s_w^2 = 45.38$$

- For an F-distribution with k 1 = 2 and n k = 128 degrees of freedom, p < 0.001
- Therefore, reject H₀ and conclude that the mean changes in weight for the 3 populations are not identical.

- Where are the specific differences?
- Bonferroni multiple comparisons.
- Set the overall probability for making a Type I error at 0.05: a* = 0.05/3 = 0.0167
- Diet vs. Exercise:

H₀:
$$\mu_1 = \mu_2$$

$$t_{12} = \overline{x_1} - \overline{x_2}$$

$$sqrt(s_w^2(1/n_1 + 1/n_2)) = -3.99$$

- For a t-distribution with n k = 128 df, p < 0.001. \rightarrow reject H_0 at the 0.0167 level of significance, conclude that μ_1 differs from μ_2 .
- Diet vs. No Plan:

$$t_{13} = \overline{x}_1 - \overline{x}_3$$

 $sqrt(s_w^2(1/n_1 + 1/n_3)) = -9.47$

• p < 0.001 \rightarrow reject H₀, conclude that $\mu_1 \neq \mu_3$

Exercise vs. No Plan:

$$t_{23} = \frac{\overline{x}_2 - \overline{x}_3}{sqrt(s_w^2(1/n_2 + 1/n_3))} = -5.74$$

- p < 0.001 \rightarrow conclude that $\mu_2 \neq \mu_3$
- Therefore, all 3 population means are different from each other.

$$\mu_1 < \mu_2 < \mu_3$$

Q1: Decreasing the significance level α decreases the probability of making a Type II Error.

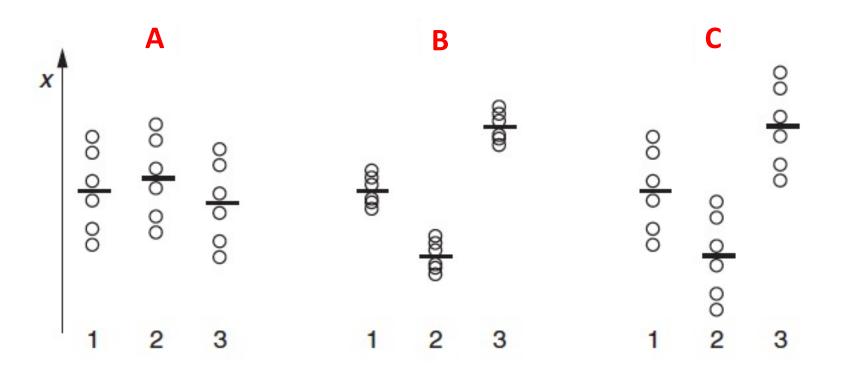
A. True

B. False

Q2: The ANOVA test produces reliable results when the following conditions hold (pick the <u>incorrect</u> one):

- A. The random samples are drawn from normally distributed populations.
- B. All data points are independent of each other.
- C. Each of the populations have the same variance σ^2 .
- D. Each of the populations have the same mean μ .

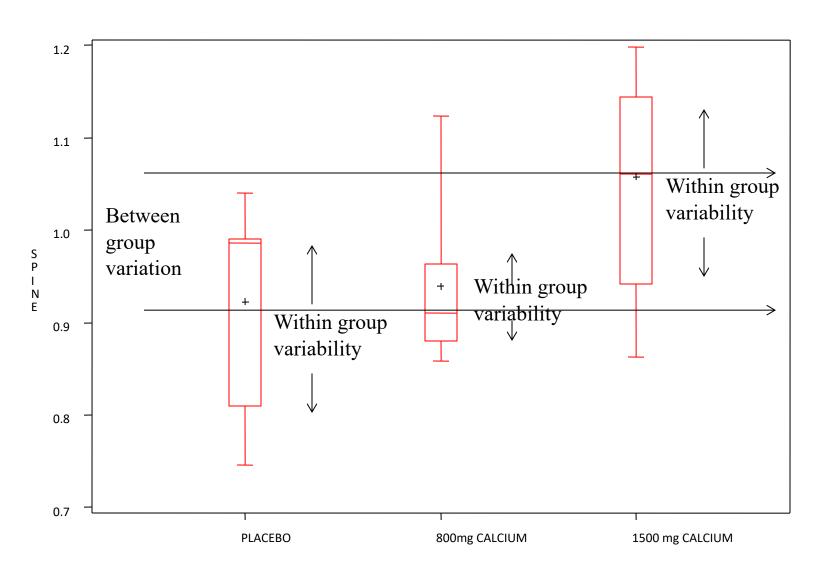
Q3: Which data set will most likely result in a rejected H₀ in an ANOVA test?



ANOVA example

- Randomize 33 subjects to three groups: 800 mg calcium supplement vs. 1500 mg calcium supplement vs. placebo.
- Compare the spine bone density of all 3 groups after 1 year.

Spine bone density vs. treatment



Group means and standard deviations

- Placebo group (n=11):
 - Mean spine BMD = $.92 \text{ g/cm}^2$
 - standard deviation = .10 g/cm²
- 800 mg calcium supplement group (n=11)
 - Mean spine BMD = $.94 \text{ g/cm}^2$
 - standard deviation = .08 g/cm²
- 1500 mg calcium supplement group (n=11)
 - Mean spine BMD =1.06 g/cm²
 - standard deviation = .11 g/cm²

Between-group variation.

The size of the groups.

The F-Test

The difference of each group's mean from the overall mean.

$$s_{between}^2 = ns_{\bar{x}}^2 = 11*\left(\frac{(.92 - .97)^2 + (.94 - .97)^2 + (1.06 - .97)^2}{3 - 1}\right) = .063$$

$$s_{within}^2 = avg \ s^2 = \frac{1}{3}(.10^2 + .08^2 + .11^2) = .0095$$

$$F_{2,30} = \frac{s_{between}^2}{s_{within}^2} = \frac{.063}{.0095} = 6.6$$

The average amount of variation within groups.

Each group's varian

Large F value indicates that the between group variation exceeds the within group variation (=the background noise).

Q4: Pick the most appropriate statistical test...

Are three sample means different from one another?

$$H_0$$
: $\mu_1 = \mu_2 = \mu_3$

- A. Two-sample z test
- B. Paired t test
- C. Two-sample t test
- D. χ^2 test
- E. Analysis of variance

Q5: Pick the most appropriate statistical test...

Compare the mean rolling velocity of 2500 leukocytes at a low fluid shear rate, with the mean rolling velocity of 3000 leukocytes at a high fluid shear rate.

- A. Two-sample z test
- B. Paired t test
- C. Two-sample t test
- D. χ^2 test
- E. Analysis of variance

Q6: Pick the most appropriate statistical test...

Compare the red blood cell counts of 20 patients before and after treatment with erythropoietin.

- A. Two-sample z test
- B. Paired t test
- C. Two-sample t test
- D. χ^2 test
- E. Analysis of variance

Q7: Pick the most appropriate statistical test...

When do fibroblasts stop spreading on fibronectin? Compare the average area of 12 cells at t = 30 min with the average area of 10 cells at t = 120 min.

- A. Two-sample z test
- B. Paired t test
- C. Two-sample t test
- D. χ^2 test
- E. Analysis of variance

Q8: Pick the most appropriate statistical test...

Mortality rates of different groups of leukemic mice are presented in a 4 x 2 contingency table. Test for differences among the population.

- A. Two-sample z test
- B. Paired t test
- C. Two-sample t test
- D. χ^2 test
- E. Analysis of variance

- <u>Parametric tests</u>: (e.g., t-test and ANOVA)
 assumes various parameters and <u>normal</u>
 distribution.
- <u>Nonparametric tests</u>: no assumption of population distribution is made.
- Testing if your data meets normality assumptions...

• Step 1:

- In decreasing order of robustness, either:
 - 1. Test that your data is not significantly deviant from a normal distribution using a test designed to do this (Kolmogorov-Smirnov, Anderson-Darling, or Shapiro-Wilks tests)
 - 2. Plot the data on a <u>normal probability plot</u>. Perfectly normal data will lie on a straight line.

Normal probability plot:

- a. Arrange data from smallest to largest.
- b. Percentile of each data value is determined.
- c. From these percentiles, normal calculations are done to determine their corresponding z-scores.
- d. Each z-score is plotted against its corresponding data value.
 - $z = (\overline{x}_1 \overline{x}_2)/SE(\overline{x}_1 \overline{x}_2)$ where z is distributed as a Normal distribution with μ =0 and σ =1.

Percentile
$$\rightarrow z \rightarrow \overline{x}_1$$

- In decreasing order of robustness, either...
 - 1. Test the data (specialized tests)
 - 2. Plot on <u>normal prob plot</u>
 - 3. Construct a frequency histogram of your data to see whether it looks normal, i.e., symmetrical and monomodal.

• Step 2:

- If the data are <u>not</u> deviant from normality according to test in Step 1, you can now progress to using parametric statistics.
- If data are deviant from normality you must choose between the following:
 - 1. Use a nonparametric test. These are less sensitive, avoid if possible.
 - 2. Transform the data. This can help your data conform better to a normal distribution.

- 3. Use a test based on another distribution (e.g., Weibull). This is an advanced technique, used in some statistical programs.
- 4. Use a parametric test anyway. The test may be inaccurate so be wary of accepting results very close to p = 0.05. Can increase the alpha-level to 0.01 to compensate. ANOVA is a particularly robust test, resistant to distortion by non-normality.

- Data transformation aims to either:
 - 1. Linearize (e.g., power laws, exponential functions)
 - 2. Normalize (make non-normal distributions appear more normal)
 - Equalize variances (make data conform to the assumption of equal variances, "homoscedatic").
 ANOVA assumes that sample variances are equal.

Transformation

log Y or log(Y+1)

Type of data applied to

contagious (=aggregated) or clumped distribution, or where factors in an ANOVA are synergistic and

(use +1 if data contains zeros

apparently multiplicative.

or numbers near 0)

Transformation	Type of data applied to			
arcsin(Y½)	Percentage (0–100%) or			
	proportional (0 – 1) data.			
sqrt(Y) or	Distributions where the			
sqrt(Y+0.375)	variances are proportional to			
	the means, i.e., they increase			
	in unison. (Use the +0.375			
	form if data contain zeros or			
	or numbers near zero)			

- Example of square root transformation on data where the variance increases as the mean increases.
- Data are given for the growth of tumors in three drug treatments.

Control		Tumostat		Inhibin 4	
Original	Sqr. Root	Original	Sqr. Root	Original	Sqr. Root
17	4.12	9	3.00	5	2.24
16	4	8	2.83	4	2.00
2	1.41	3	1.73	2	1.41
1	1.00	2	1.41	1	1.00
s ² 75.34	6.92	30.25	5.02	9.00	2.75

- Original data show gross heteroscedasticity, ratio of largest variance to smallest = 8.4:1.
- Square root transformation reduces this ratio to 2.5:1.
- (for t-test, can pool s_d² when variances differ by ~2, more statistical power.)