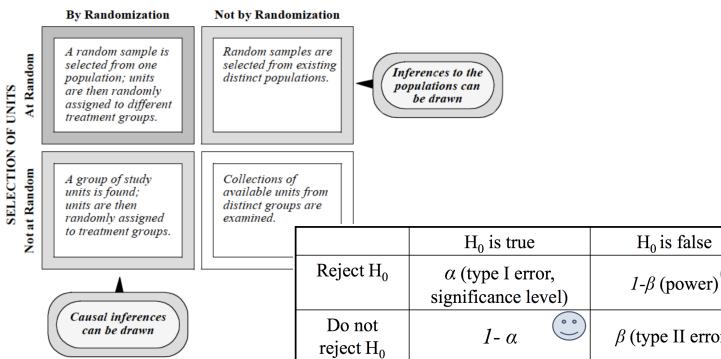


## Definitions:

<p><b>Study (experimental) unit (subject)</b> - one member of a set of entities being studied.</p> <p><b>Parameter (also, estimand)</b> - proportion of childless households in the population.</p> <p><b>Estimate</b> - proportion of childless households in the sample.</p> <p><b>Parameter</b> is a population characteristic.</p> <p><b>Estimand</b> is a parameter that is being estimated (<math>\mu</math>).</p> <p><b>Statistic</b> is a function of the data (therefore, it is random).</p> <p><b>Statistic</b>: A function of the data, <math>\mu = \mu(y)</math>. Test Statistic: Statistic used to weigh evidence supporting and contradicting the null hypothesis.</p> <p><b>Reference Distribution</b>: Probability distribution of the test statistic, assuming that the null hypothesis is true, <math>f(\mu(Y) H_0)</math>.</p> <p><b>Estimator</b> is a statistic used as a guess for the value of the estimand ( ).</p> <p><b>Estimate</b> is a quantity, which is a particular realization of the estimator (4/12 = 0.33).</p> <p><b>Target Population</b>: A collection of units a researcher is interested in; a group about which the researcher wishes to draw conclusions.</p> <p><b>Sampling frame</b>: Collection of units that are potential members of the sample.</p> <p><b>Oversampling</b>: selecting units that are outside your sampling frame.</p> <p><b>Undersampling</b>: excluding units that are within your sampling frame.</p> <p><b>Sample</b>: A [randomly selected] subset of a sampling frame.</p> <p><b>Simple Random Sampling (SRS)</b> - every subset of <math>n</math> units has equal chance to be selected</p> <p><b>Stratified Sampling</b> - split the population into homogeneous subpopulations and use SRS (or another method) within a sampling frame of each subpopulation.</p> <p><b>Systematic Random Sampling</b> - select every <math>k</math>th unit from the ordered sampling frame, starting randomly from the first <math>k</math> positions.</p> <p><b>Variable probability sampling</b> - allow units to have unequal probabilities of being sampled.</p> <p><b>p-value</b> - probability that the test statistic would be at least as extreme as observed, under the null hypothesis. (or "strength of evidence against the <math>H_0</math>")</p> <p><b>Significance level (<math>\alpha</math>)</b> is the criterion compared against the p-value. The null hypothesis is rejected if p-value is lower than <math>\alpha</math>.</p> <p><math>\alpha</math> reflects the probability of rejecting the null hypothesis given that it is true (<b>Type I error</b>).</p> <p><b>Scope of inference</b></p> <ul style="list-style-type: none"> <li>- Internal validity: assumptions of the test satisfied?</li> <li>- External validity: possible to make inference to a broader population?</li> </ul> <p><b>Sampling distribution</b> of a statistic is a (reference) distribution that arises from a chance mechanism used to select a random sample from a population.</p> <p><b>Type I Error</b> (or significance level <math>\alpha</math>): Probability of rejecting the null hypothesis, when the null hypothesis is true.</p> <p><b>Type II Error</b> (<math>\beta</math>): Probability of failing to reject the null hypothesis, when the null hypothesis is false.</p> <p><b>Power (1-<math>\beta</math>)</b>: Probability of rejecting the null hypothesis, when a particular alternative hypothesis is true.</p> <p>If <math>\alpha = 0</math>, <math>\beta = 1</math>; if <math>\alpha = 1</math>, <math>\beta = 0</math>; if alpha increases, beta decreases (power increases).</p> <p>Increase power by: increase alpha; do one-tailed test; increase effect size; increase sample size</p> <p>Most to least power, if assumption of t-tools is met</p> <ul style="list-style-type: none"> <li>- For small samples:             <ol style="list-style-type: none"> <li>1. T-tools</li> <li>2. Permutation test</li> <li>3. Rank-sum test (or signed-rank test for paired data)</li> <li>4. Sign test for paired data)</li> </ol> </li> </ul> <p>Large samples rejection rates are basically equivalent.</p>	<p>Population Variance: <math>\sigma^2 = E((Y - \mu)^2)</math></p> <p>Sample Variance:</p> $S^2 = \frac{1}{n-1} \sum_{i=1}^n (Y_i - \bar{Y})^2 \sim \frac{\sigma^2}{n-1} \chi^2_{n-1}$ <p>Sample variance is chi-sq distributed:</p> $\frac{Z_0}{\sqrt{X^2/n}} \sim t_n$ <p>Let <math>Z_0 \sim N(0,1)</math>, independent of variance <math>X^2 \sim \chi^2_{n-1}</math>, then</p> $X^2 = \sum_{i=1}^n Z_i^2 \sim \chi^2_n$ <p>Chi-sq distribution with <math>n</math> df:</p> $S_p^2 = \frac{(n_X - 1)S_X^2 + (n_Y - 1)S_Y^2}{n_X + n_Y - 2}$ <p>Pooled sample variance: where</p> $S_p^2 = \frac{\sum_{i=1}^{n_X} (X_i - \bar{X})^2}{n_X - 1}, \quad S_p^2 = \frac{\sum_{i=1}^{n_Y} (Y_i - \bar{Y})^2}{n_Y - 1}$ $S_p^2 \sim \frac{\sigma^2}{n_X + n_Y - 2} \chi^2_{n_X + n_Y - 2}$ <p>Sampling distribution of pooled sample variance:</p> $X \sim \chi^2_{n_X} \text{ and } Y \sim \chi^2_{n_Y}$ <p>F-distribution: if <math>X \sim \chi^2_{n_X}</math> and <math>Y \sim \chi^2_{n_Y}</math>, independent of each other, then</p> $\frac{X/n_X}{Y/n_Y} \sim F_{n_X, n_Y}$ <p>(F-distribution with <math>n_X</math> and <math>n_Y</math> d.f.)</p>
<p>No</p>	

## ALLOCATION OF UNITS TO GROUPS



Sum of Squared Residuals	General Name	Calculation	Degrees of freedom	Distribution	Corresponding variance (also called "Mean Square")
SSR <sub>full</sub> Residual sum of squares (full)	SSW (Within Groups)	$\sum_{i=1}^I \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y}_i)^2$	n-l	$\sigma^2 \chi^2_{n-l}$	$S_p^2$ MSR <sub>full</sub>
ESS (Extra sum of squares)	SSB (Between Groups)	$\sum_{i=1}^I n_i (\bar{Y}_i - \bar{Y})^2 = \sum_{i=1}^I n_i (\bar{Y}_i)^2 - n(\bar{Y})^2$	l-1	$\sigma^2 \chi^2_{l-1}$	MSB
SSR <sub>reduced</sub> Residual sum of squares (reduced)	SST (Total)	$\sum_{i=1}^I \sum_{j=1}^{n_i} (Y_{ij} - \bar{Y})^2$	n-1	$\sigma^2 \chi^2_{n-1}$	Variance of the entire sample as one group

**(Fisher's) Randomization Test**: is a distribution-free test for treatment effect in **randomized experiments**.

**H0**: Zero treatment effect for all units,  $\delta=0$ . Each unit's outcome is the same, regardless of the treatment assigned. Consequently, the distribution of outcomes is identical in two groups

**Ha**: Non-zero treatment effect for ALL units,  $\delta \neq 0$ .

Assumptions:

1. Random assignment to groups.
2. Under the H0, independence of study units.

**Test statistic**: Difference between average outcomes

$$\bar{Y}_c - \bar{Y}_v$$

in the two groups\*.

\*In fact, any summary of the data can represent a test statistic.

**Randomization distribution** is a reference distribution of a test statistic in a randomization test, where variation is due to random assignment of the treatment.

**Permutation test** is a distribution-free nonparametric test for association between group status and outcome in **observational studies**.

**HO** : Outcomes not related to group status.

**Ha** : Outcomes are related to (associated with) group status.

Assumptions: Independence of study units.

**Test statistic**: Difference in average observed outcomes between the two groups (or, any other statistic).

Permutation distribution is a reference distribution of test statistics in a permutation test.

**NOTE**: Permutation tests are actually evaluating whether the distributions are equivalent, not just the

### How does CLT apply?

#### # checking assumptions:

**Independence**: plot subgroups, and see if relationship still holds (cluster effect)

Plot data over time (serial effects)

Plot data vs space (spatial effect)

**Normality**: Histograms should look normal – overlay with normal curve and kernel density plot

Boxplot or Q-Q plot will show outliers

**Equal Variance**: divide variances to see if ratio is  $<0.5$  or  $>2$

#### T-test is robust:

-if populations are symmetrically skewed

-if sample sizes are equal (and large)

-when sample sizes are equal, pooled-ttest is robust to unequal variances

**Sensitive**: When the sample sizes are not equal, t-tests are more sensitive to skewedness and long-tailedness.

For small samples, t-tests are somewhat sensitive to markedly different skewedness in two groups.

Watch out for outliers.

#### Robustness of F-tests:

Not resistant to outliers. Fairly robust to non-normality. Independence w/l and b/w groups is necessary. Equal variance is necessary.

#### F-Test Assumptions:

Equal variances is crucial

Not resistant to outliers

Normality is not critical.

#### Confidence intervals: $Y_i \sim \text{iid } N(\mu, \sigma^2)$ .

$$\left( \bar{Y} - t_{1-\alpha/2, n-1} \frac{S}{\sqrt{n}}, \bar{Y} + t_{1-\alpha/2, n-1} \frac{S}{\sqrt{n}} \right)$$

One Sample t-test:

### Wilcoxon Signed-Rank Test for Paired Data:

(more assumptions, but higher power than sign test)

#### Hypotheses: General formulation:

**H0**: The rank of the magnitude of within-pair difference is unrelated to the sign of the difference.

**Ha**: The rank of the magnitude of within-pair difference is related to the sign of the difference.

#### If differences are continuous r.v.'s:

**H0**: Median(Di)=0

**Ha**: Median(Di) ≠ 0 (or >0, or <0)

#### If the distribution of differences is symmetric:

**H0**: E(Di)=0

**Ha**: E(Di) ≠ 0 (or >0, or <0)

1. Calculate differences (+ or -).

2. Rank the absolute values of differences |Di|

$$Z_i = \text{Rank}(|D_i|), i = 1, \dots, n.$$

Test Statistic: (sum of ranks for positive differences)

$$S = \sum_{i=1}^n Z_i |D_i| > 0$$

**Exact sampling distribution**: do permutation test, and switch group status for each pair.

**Normal approximation** (for large sample sizes, m is  $\geq 20$ , where m is the number of non-zero differences):

$$Z = \frac{S - m(m+1)/4}{\sqrt{m(m+1)(2m+1)/24}} \sim N(0,1)$$

Assumptions:

1. Data are paired and come from the same

<p><i>equality of the means:</i></p> <p><b>One Sample z-Test</b>  <math>H_0: \mu = \mu_0</math>  <math>HA: \mu \neq \mu_0</math></p> $Z = \frac{\bar{Y} - \mu_0}{(\sigma / \sqrt{n})}$ <p><b>Test Statistic:</b> <math>Z = \frac{\bar{Y} - \mu_0}{(\sigma / \sqrt{n})}</math> (make sure to double p-value if doing 2-sided test, of <math>\frac{1}{2}</math> alpha)  Exact sampling distribution of Z under <math>H_0: Z \sim N(0,1)</math></p> <p><b>Assumptions:</b>  1. Independence  2. Known population variance, <math>\sigma^2</math>  3. Normality (CLT allows for deviations!)  4. Random sampling from population</p> <p><b>One Sample t-Test</b>  <math>H_0: \mu = \mu_0</math>  <math>HA: \mu \neq \mu_0</math></p> $T = \frac{\bar{Y} - \mu_0}{(S / \sqrt{n})}$ <p><b>Test Statistic:</b> Exact sampling distribution of T under <math>H_0: T \sim t_{n-1}</math> (t-distribution with <math>df = n-1</math>)</p> <p><b>Assumptions:</b>  1. Independence  2. Normality (CLT allows for deviations!)  3. Random sampling from population</p> <p><b>Two Sample z-Test</b>  <math>H_0: \mu_x - \mu_y = \mu_0</math>  <math>HA: \mu_x - \mu_y \neq \mu_0</math></p> $Z = \frac{(\bar{X} - \bar{Y} - \mu_0)}{\sqrt{\frac{s_x^2}{n_x} + \frac{s_y^2}{n_y}}} \sim N(0,1)$ <p><b>Test Statistic:</b> Exact sampling distribution of T under <math>H_0: T \sim t_{n_x+n_y-2}</math> (t-distribution with <math>df = nx+ny-2</math>)</p> <p><b>Note:</b> <math>(\bar{X} - \bar{Y}) \sim N\left[\mu_x - \mu_y, \sigma^2\left(\frac{1}{n_x} + \frac{1}{n_y}\right)\right]</math>, <math>S_x^2 + S_y^2</math></p> <p><math>\frac{s_x^2}{n_x} + \frac{s_y^2}{n_y}</math> is an unbiased estimator of <math>\frac{\sigma_x^2}{n_x} + \frac{\sigma_y^2}{n_y}</math></p> <p><b>Assumptions:</b> same as Welch's, but If using pooled variance – assume equal variances between populations.</p> <p><b>Unpooled (Welch) Two-Sample t-Test</b>  <math>H_0: \mu_x - \mu_y = \mu_0</math>  <math>HA: \mu_x - \mu_y \neq \mu_0</math></p> $T = \frac{(\bar{X} - \bar{Y} - \mu_0)}{\sqrt{\frac{S_x^2}{n_x} + \frac{S_y^2}{n_y}}}$ <p><b>Test Statistic:</b> Approximate Sampling distribution:  <math>T \sim t_{\bar{n}}</math>, where <math>\bar{n} = \left(\frac{s_x^2}{n_x} + \frac{s_y^2}{n_y}\right)^{-1} / \left(\frac{s_x^2}{n_x^2(n_x-1)} + \frac{s_y^2}{n_y^2(n_y-1)}\right)</math></p> <p><b>Assumptions:</b>  1. Independence between units:  – within each population (<math>Y_i</math> with <math>Y_j</math> and <math>X_i</math> with <math>X_j</math>)  – between populations (<math>Y_i</math> with <math>X_j</math>)  2. Homogeneity of units within each population:  – Equal means within each population;  – Equal variances within each population;  – If using pooled variance – assume equal variances between populations.  3. Population distributions are Normal:  – CLT allows for deviations.  3. Random sampling from populations.</p> <p><b>Paired t-Test</b>  <i>i.i.d.</i></p> $X_i - Y_i = D_i \sim N(\mu_D, \sigma^2)$ <p><math>H_0: \mu_D = \mu_0</math>  <math>HA: \mu_D \neq \mu_0</math></p> $T = \frac{\bar{D} - \mu_0}{(S_D / \sqrt{n})}$ <p><b>Test Statistic:</b> Exact sampling distribution of T under <math>H_0: T \sim t_{n-1}</math> (where <math>n</math> is # of pairs)</p> <p><b>Assumptions same as one-sample t-test</b></p> <p><b>F-test for equal variance</b> (could also use Levene's test): (caution: will reject for large samples when ratio is close to 1)</p>	<p><b>Two Sample t-test:</b></p> <ul style="list-style-type: none"> <li><b>Two-sample t-tests</b> <ul style="list-style-type: none"> <li><math>(\bar{X} - \bar{Y} - t_{1-\alpha/2, df}) \cdot SE, \quad \bar{X} - \bar{Y} + t_{1-\alpha/2, df} \cdot SE)</math></li> <li>Pooled t-test: <math>SE = S_p \sqrt{\frac{1}{n_x} + \frac{1}{n_y}}, \quad df = n_x + n_y - 2</math></li> <li>Unpooled t-test: <math>SE = \sqrt{\frac{S_x^2}{n_x} + \frac{S_y^2}{n_y}}, \quad df = \tilde{n}</math>  <math>\tilde{n} = \left(\frac{s_x^2}{n_x} + \frac{s_y^2}{n_y}\right)^2 / \left(\frac{s_x^2}{n_x^2(n_x-1)} + \frac{s_y^2}{n_y^2(n_y-1)}\right)</math></li> </ul> </li> </ul> <p><b>Two-Sample t-test for I samples (<math>I &gt; 2</math>)</b></p> <ul style="list-style-type: none"> <li><b>Random sampling.</b> <ul style="list-style-type: none"> <li>Confidence intervals: <math>SE = S_p \sqrt{\frac{1}{n_m} + \frac{1}{n_k}}, \quad df = n_1 + n_2 + \dots + n_I - I</math></li> <li><math>(\bar{Y}_m - \bar{Y}_k - t_{1-\alpha/2, df} \cdot SE, \quad \bar{Y}_m - \bar{Y}_k + t_{1-\alpha/2, df} \cdot SE)</math></li> </ul> </li> </ul> <p><b>Transformations:</b>  Log Transformation: <ol style="list-style-type: none"> <li><b>Randomized Experiments</b> (T=Treatment, C=Control) <math>Z_{T,i} = \ln(Y_{T,i}), \quad i = 1, \dots, n_T</math> <math>Z_{C,i} = \ln(Y_{C,i}), \quad i = 1, \dots, n_C</math> Responses: <math>Z_{T,i} = Z_{C,i} + \delta \Leftrightarrow \frac{Y_{T,i}}{Y_{C,i}} = e^\delta</math> <math>\exp(\bar{Z}_T - \bar{Z}_C)</math> estimates <math>\frac{Y_{T,i}}{Y_{C,i}}</math> </li> <li>Observational studies with random sampling: <math>\exp(\bar{Z}_2 - \bar{Z}_1)</math> estimates <math>\frac{m_2}{m_1}</math> </li> <li>In paired t-test: <b>Randomized Experiments:</b> <math>\exp(\bar{Z})</math> estimates a multiplicative treatment effect <math>\frac{Y_{2,i}}{Y_{1,i}}</math> </li> </ol> <p><b>Observation studies w/ rand. Sampling:</b>  Median(<math>Y_{2,i} / Y_{1,i}</math>) = <math>m</math>, then <math>\exp(\bar{Z})</math> estimates <math>m</math>.</p> <p>Square Root Transformation <math>\sqrt{Y}</math>: Good for moderately skewed data, but hard to interpret results</p> <p>Reciprocal Transformation <math>1/Y</math>: good for severely skewed data, can be used with negative data</p> <p>Logit <math>\log(Y/(1-Y))</math> or <math>\arcsin(2Y-1)</math>: good for proportions</p> <p><b>Nonparametric alternatives to t-tools:</b></p> <p><b>Rank Sum test:</b>  Hypotheses:  <math>H_0</math>: (if treatments are randomized): Same as Fisher's randomization test  <math>H_0</math>: (if treatments are not randomized): same as permutation test  * If shapes and spreads of the two populations are similar: HA: There is a difference in medians between the two populations.  Transform data:  <math>Z_{1,i} = \text{Rank}(X_i   X_1, \dots, X_{n_x}, Y_1, \dots, Y_{n_y}), \quad i = 1, \dots, n_x</math>  <math>Z_{2,i} = \text{Rank}(Y_i   X_1, \dots, X_{n_x}, Y_1, \dots, Y_{n_y}), \quad i = 1, \dots, n_y</math>  Ties are averaged  <math>T = \sum_{i=1}^{n_x} Z_{1,i}</math>, assuming <math>n_x \leq n_y</math>  Test statistic:  Assumptions: Independence</p> <p>Normal approximation to rank Sum test (if no ties, and samples greater than 10):  <math>T \sim N\left(\frac{n_x(n_x + n_y + 1)}{2}, \frac{n_x n_y (n_x + n_y + 1)}{12}\right)</math></p> <p>With ties, Rbar is sample mean (mean of all ranks) and <math>S_r^2</math> is sample variance for combined set of <math>nx + ny</math> ranks:  <math>T \sim N\left(n_x \bar{R}, S_r^2 \frac{n_x n_y}{n_x + n_y}\right)</math></p> <p>We can use this Z-statistic:  <math>Z = \frac{T - n_x \bar{R}}{S_r \sqrt{\frac{n_x n_y}{n_x + n_y}}}</math></p> <p><b>Sign test for paired data</b> (lower power than Wilcoxon signed-rank test)  <math>H_0: P(D_i &gt; 0) = P(Y_i &gt; X_i) = 0.5</math>,</p> <p><b>General formulation:</b> <math>H_a: P(D_i &gt; 0) \neq 0.5</math> (or <math>&gt; 0.5</math>, or <math>&lt; 0.5</math>).</p> </p>	<p>2. population.  Each pair is chosen randomly and independently.</p> <p>3. The data are measured at least on an <a href="#">ordinal scale</a>, but need not be normal.</p> <p><b>Two-Sample t-Test for I samples</b>  <math>H_0: \mu_m - \mu_k = \mu_0</math>  <math>HA: \mu_m - \mu_k \neq \mu_0</math></p> <p><b>Test Statistic:</b></p> $T = \frac{(\bar{Y}_m - \bar{Y}_k - \mu_0)}{S_p \sqrt{\frac{1}{n_m} + \frac{1}{n_k}}}, \quad S_p^2 = \frac{(n_1-1)S_1^2 + (n_2-1)S_2^2 + \dots + (n_I-1)S_I^2}{n_1 + n_2 + \dots + n_I - I}$ <p>Exact sampling distribution of T under <math>H_0: T \sim t_{n_1+n_2+\dots+n_I-1}</math>, where I is number of groups.</p> <p><b>Assumptions:</b>  Independence of units within each group;  Independence of units between groups;  Normality of all I populations;  Equal variances for all I populations;  Random sampling.</p> <p><b>Anova F-test for Equality of All means in I samples</b>  <math>H_0: \mu_1 = \mu_2 = \dots = \mu_I</math> (reduced model – equal means model)  Ha: at least one mean is different from the others (full model – separate means model)</p> <p><math>SS_{\text{full}} = SSW</math> (within groups)  <math>S_p^2 = \frac{S^2}{n-I}</math>  <math>S^2 = \frac{\sigma^2}{n-I} \chi_{n-I}^2 \Rightarrow SS_{\text{full}} \sim \sigma^2 \chi_{n-I}^2</math>, d.f. = <math>n-I</math></p> <p>Exact sampling distribution of T under <math>H_0: T \sim t_{n_1+n_2+\dots+n_I-1}</math>, where I is number of groups.</p> <p><b>Assumptions:</b>  Independence of units within each group;  Independence of units between groups;  Normality of all I populations;  Equal variances for all I populations;  Random sampling.</p> <p><b>ESS</b> (SSB (between groups))  <math>ESS = SS_{\text{reduced}} - SS_{\text{full}}</math>  <math>H_0</math>  <math>ESS \sim \sigma^2 \chi_{I-1}^2</math>, d.f. = <math>I-1</math></p> <p><b>Test Statistic:</b></p> $R = \frac{ESS / (I-1)}{SS_{\text{full}} / (n-I)} = \frac{MSB}{MSR_{\text{full}}}$ <p>MSB = Mean square between  MSR<sub>full</sub> = Mean square residuals (within groups)</p> <p>Easy way to calculate: <math>MSR_{\text{full}} = S_p^2</math>  Exact sampling distribution of R under <math>H_0: R \sim F_{I-1, n-I}</math></p> <p><b>Kruskal-Wallis Test</b> (Nonparametric ANOVA)  <math>H_0: \text{Median}(Y_1) = \text{Median}(Y_2) = \dots = \text{Median}(Y_I)</math>  Ha: at least one median is different from the others</p> <p>Test Statistic – convert to ranks, and do ANOVA on ranks:  <math>KW = \frac{ESS \text{ (on ranks)}}{S_R^2} \sim \chi_{I-1}^2</math></p> <p><math>S_R^2</math> is the sample variance of all ranks (not pooled)</p> <p><b>Assumptions:</b>  1. Independence.  2. Random Sampling</p> <p><b>F-test for Equality in a subset of groups</b></p>
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$H_0: \sigma_x^2 = \sigma_y^2$ $H_a: \sigma_x^2 \neq \sigma_y^2 \Leftrightarrow \sigma_x^2 / \sigma_y^2 \neq 1$ Test Statistic: $F = \frac{S_x^2}{S_y^2} \sim F_{n_x-1, n_y-1}$ Same assumptions as t-test: <b>In testing the equality of population variances, two assumptions are required: independent samples and normally distributed populations.</b> 2-sided p-value is just 2*the one-sided value. Multiple Comparisons	$H_0: \text{Median}(D_i) = 0,$ $H_a: \text{Median}(D_i) \neq 0 \text{ (or } >0, \text{ or } <0\text{)}$ <b>If differences are continuous r.v's:</b> $H_0: E(D_i) = 0,$ <b>If differences are symmetric:</b> $H_a: E(D_i) \neq 0 \text{ (or } >0, \text{ or } <0\text{)}$ Test Statistic: K = number of positive differences. Exact distribution of K: $K \sim \text{Binom}(m, 0.5)$ , where m is final number of pairs with nonzero $D_i$ . Normal approximation (for large samples): $Z = \frac{K - m/2}{\sqrt{m/4}} \sim N(0,1)$ Assumptions: 1. The differences $D_i$ are assumed to be independent. 2. Each $D_i$ comes from the same continuous population. 3. The values $X_i$ and $Y_i$ represent are ordered (at least the <a href="#">ordinal scale</a> ), so the comparisons "greater than", "less than", and "equal to" are meaningful.	~~~~~ Linear Combinations of Means
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Hypotheses for nonparametric tests		
By Randomization Not by Randomization		
$H_0: \text{Each unit in the population would have the same outcome regardless of the treatment assigned.}$ $H_a: \text{Treatment effect is non-zero, } \delta \neq 0, \text{ in the population.}$ <i>If similar shapes and spreads :  <math>H_0: \text{Median}(X) \neq \text{Median}(Y) \text{ (or } \geq, \text{ or } \leq\text{).}</math></i>	$H_0: \text{Outcome distributions in the two groups within the population are the same.}$ $H_a: \text{There is an association between group status and outcomes in the population.}$ <i>If similar shapes and spreads :  <math>H_0: \text{Median}(X) \neq \text{Median}(Y) \text{ (or } \geq, \text{ or } \leq\text{).}</math></i>	
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Derive sampling distribution: Q2d on Hw 3

Review "Overview" in lecture 13

Central Limit Theorem

Law of large numbers

Internal vs. external validity

Study design concepts (units, target population, SRS vs. stratified

random sampling, etc.);

} Sample statistics vs. population parameters;

} Rules for expectations and variances of r.v.;

} Mechanics of inference using randomization (i.e., random

permutation) vs. random sampling;

} Tests for 1, 2, and 3+ samples:

} Hypotheses that they are testing (depending on a question of interest,

study design, and data transformation used, if any);

} Assumptions for each test and methods for checking them;

} Test-statistics for each test and their sampling distributions, looking up

p-values using a table, calculating confidence intervals;

} Ability to interpret output of R functions that perform these tests;

} Ability to state conclusions and comment on a scope of inference.

} Definition of distributions by representation;  
Conceptual, derivations, other non-simulation questions,

similar to those on homeworks.

} Motivation for and interpretation of simulations on  
homeworks.

} Tests and confidence-interval calculation using  
distribution tables, comments on assumptions.

} Explanation of what happens behind any computer  
function you've used.

} Suggestion: verify the calculations behind t-tests and  
other tests for some of the homework problems.

} Ramsey and Schafer has lots of problems to work  
through (including conceptual ones)

<p><b>Study (experimental) unit (subject)</b> - one member of a set of entities being studied.</p> <p><b>Parameter (also, estimand)</b> - proportion of childless households in the population.</p> <p><b>Estimate</b> - proportion of childless households in the sample.</p> <p><b>Parameter</b> is a population characteristic.</p> <p><b>Estimand</b> is a parameter that is being estimated (<math>\mu</math>).</p> <p><b>Statistic</b> is a function of the data (therefore, it is random).</p> <p><b>Statistic</b>: A function of the data, <math>\mu = \mu(Y)</math>. Test Statistic: Statistic used to weigh evidence supporting and contradicting the null hypothesis.</p> <p><b>Reference Distribution</b>: Probability distribution of the test statistic, assuming that the null hypothesis is true, <math>f(\mu(Y) H_0)</math>.</p> <p><b>Estimator</b> is a statistic used as a guess for the value of the estimand () .</p> <p><b>Estimate</b> is a quantity, which is a particular realization of the estimator (4/12 = 0.33).</p> <p><b>Target Population</b>: A collection of units a researcher is interested in; a group about which the researcher wishes to draw conclusions.</p> <p><b>Sampling frame</b>: Collection of units that are potential members of the sample.</p> <p><b>Overcoverage</b>: selecting units that are outside your sampling frame.</p> <p><b>Undercoverage</b>: excluding units that are within your sampling frame.</p> <p><b>Sample</b>: A [randomly selected] subset of a sampling frame</p> <p><b>Simple Random Sampling (SRS)</b> - every subset of <math>n</math> units has equal chance to be selected</p> <p><b>Stratified Sampling</b> - split the population into homogeneous subpopulations and use SRS (or another method) within a sampling frame of each subpopulation.</p> <p><b>Systematic Random Sampling</b> - select every <math>k^{\text{th}}</math> unit from the ordered sampling frame, starting randomly from the first <math>k</math> positions.</p> <p><b>Variable probability sampling</b> - allow units to have nonequal probabilities of being sampled.</p> <p><b>p-value</b> - probability that the test statistic would be at least as extreme as observed, under the null hypothesis. (or "strength of evidence against the <math>H_0</math>".)</p> <p><b>Significance level (<math>\alpha</math>)</b> is the criterion compared against the p-value. The null hypothesis is rejected if p-value is lower than <math>\alpha</math>. <math>\alpha</math> reflects the probability of rejecting the null hypothesis given that it is true (<b>Type I error</b>).</p> <p><b>Scope of inference</b></p> <ul style="list-style-type: none"> <li><b>Internal validity</b>: are assumptions of the test satisfied?</li> <li><b>External validity</b>: possible to make inference to a broader population?</li> </ul> <p><b>Sampling distribution</b> of a statistic is a (reference) distribution that arises from a chance mechanism used to select a random sample from a population.</p> <p><b>Type I Error</b> (or significance level <math>\alpha</math>): Probability of rejecting the null hypothesis, when the null hypothesis is true.</p> <p><b>Type II Error</b> (<math>\beta</math>): Probability of failing to reject the null hypothesis, when the null hypothesis is false.</p> <p><b>Power (1-<math>\beta</math>)</b>: Probability of rejecting the null hypothesis, when a particular alternative hypothesis is true.</p> <p>If <math>\alpha = 0</math>, beta = 1; if <math>\alpha = 1</math>, beta = 0; if alpha increases, beta decreases (power increases).</p> <p>Increase power by: increase alpha; do one-tailed test; increase effect size; increase sample size</p> <p>Most to least power, if assumption of t-tools is met</p> <ul style="list-style-type: none"> <li>For small samples:</li> <li>1. T-tools</li> <li>2. Permutation test</li> <li>3. Rank-sum test (or signed-rank test for paired data)</li> <li>4. (Sign test for paired data)</li> </ul> <p>Large samples rejection rates are basically equivalent</p>	<p><b>Individual confidence level</b> is a probability that a single confidence interval covers the true value.</p> <p><b>Familwise confidence level</b> is a probability that all confidence intervals cover the corresponding true values.</p> <p><b>Common distributions and calculations:</b></p> <p>Population Variance: <math>\sigma^2 = E(Y - \mu)^2</math></p> <p>Sample Variance:</p> $S^2 = \frac{1}{n-1} \sum_{i=1}^n (Y_i - \bar{Y})^2 \sim \frac{\sigma^2}{n-1} \chi^2_{n-1}$ <p>Sample variance is chi-sq distributed:</p> $\frac{Z_0}{\sqrt{X^2/n}} \sim t_n$ <p>Let <math>Z_0 \sim N(0,1)</math>, independent of variance <math>X \sim \chi^2_{n-1}</math>, then</p> $X^2 = \sum_{i=1}^n Z_i^2 \sim \chi^2_n$ <p>Chi-sq distribution with <math>n</math> d.f.:</p> $S_p^2 = \frac{(n_x-1)S_x^2 + (n_y-1)S_y^2}{n_x+n_y-2}$ <p>Pooled sample variance: , where</p> $S_x^2 = \frac{\sum_{i=1}^{n_x} (X_i - \bar{X})^2}{n_x-1}, \quad S_y^2 = \frac{\sum_{i=1}^{n_y} (Y_i - \bar{Y})^2}{n_y-1}$ $S_p^2 \sim \frac{\sigma^2}{n_x+n_y-2} \chi^2_{n_x+n_y-2}$ <p>Sampling distribution of pooled sample variance:</p> <p>F-distribution: if <math>X \sim \chi^2_{n_x}</math> and <math>Y \sim \chi^2_{n_y}</math>, independent of each other, then</p> $\frac{X/n_x}{Y/n_y} = R \sim F_{n_x, n_y}$ <p>(F-distribution with <math>n_x</math> and <math>n_y</math> d.f.)</p>																																													
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B[By Randomization]     A --&gt; C[Not by Randomization]     B --&gt; D[Random sample is selected from one population; units are then randomly assigned to different treatment groups.]     C --&gt; E[Random samples are selected from existing distinct populations.]     E --&gt; F[Inferences to the populations can be drawn]     D --&gt; G[A group of study units is found; units are randomly assigned to treatment groups.]     G --&gt; H[Collections of available units from distinct groups are examined.]     H --&gt; I[Causal inferences can be drawn]   </pre> <pre> graph TD     J[SELECTION OF UNITS]     J --&gt; K[At Random]     J --&gt; L[Not At Random]     K --&gt; M[Random sample is selected from one population; units are then randomly assigned to different treatment groups.]     K --&gt; N[Random samples are selected from existing distinct populations.]     L --&gt; O[A group of study units is found; units are randomly assigned to treatment groups.]     L --&gt; P[Collections of available units from distinct groups are examined.]     O --&gt; 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<p><b>(Fisher's) Randomization Test</b>: is a distribution-free test for treatment effect in <b>randomized experiments</b>.</p> <p><b>H<sub>0</sub></b>: Zero treatment effect for all units, <math>\delta=0</math>. Each unit's outcome is the same, regardless of the treatment assigned. Consequently, the distribution of outcomes is identical in two groups</p> <p><b>Ha</b>: Non-zero treatment effect for ALL units, <math>\delta \neq 0</math>.</p> <p><b>Assumptions</b>:</p> <ol style="list-style-type: none"> <li>1. Random assignment to groups.</li> <li>2. Under the H<sub>0</sub>, independence of study units.</li> </ol> <p><b>Test statistic</b>: Difference between average outcomes in the two groups*. <math>\bar{Y}_c - \bar{Y}_v</math></p> <p>*In fact, any summary of the data can represent a test statistic.</p> <p><b>Randomization distribution</b> is a reference distribution of a test statistic in a randomization test, where variation is due to random assignment of the treatment.</p> <p><b>Permutation test</b> is a distribution-free nonparametric test for association between group status and outcome in <b>observational studies</b>.</p> <p><b>H<sub>0</sub></b> : Outcomes not related to group status.</p> <p><b>Ha</b> : Outcomes are related to (associated with) group status.</p> <p>Assumptions: Independence of study units.</p> <p><b>Test statistic</b>: Difference in average observed outcomes between the two groups (or, any other statistic), Permutation distribution is a reference distribution of a test statistic in a permutation test.</p> <p><b>NOTE</b>: Permutation tests are actually evaluating whether the distributions are equivalent, not just the equality of the means:</p>	<p><b>One Sample z-Test</b></p> <p><b>H<sub>0</sub></b>: <math>\mu = \mu_0</math> <b>HA</b>: <math>\mu \neq \mu_0</math></p> $Z = \frac{\bar{Y} - \mu_0}{(\sigma/\sqrt{n})}$ <p><b>Test Statistic</b>: (make sure to double p-value if doing 2-sided test, of <math>\frac{1}{2}\alpha</math>) Exact sampling distribution of Z under H<sub>0</sub>: <math>Z \sim N(0,1)</math></p> <p><b>Assumptions</b>:</p> <ol style="list-style-type: none"> <li>1. Independence</li> <li>2. Known population variance, <math>\sigma^2</math></li> <li>3. Normality (CLT allows for deviations!)</li> <li>4. Random sampling from population</li> </ol> <p><b>One Sample t-Test</b></p> <p><b>H<sub>0</sub></b>: <math>\mu = \mu_0</math> <b>HA</b>: <math>\mu \neq \mu_0</math></p> $T = \frac{\bar{Y} - \mu_0}{(S/\sqrt{n})}$ <p><b>Test Statistic</b>: Exact sampling distribution of T under H<sub>0</sub>: <math>T \sim t_{n-1}</math> (t-distribution with df = n-1)</p> <p><b>Assumptions</b>:</p> <ol style="list-style-type: none"> <li>1. Independence</li> <li>2. Known population variance, <math>\sigma^2</math></li> <li>3. Normality (CLT allows for deviations!)</li> <li>4. Random sampling from population</li> </ol> <p><b>Two Sample z-Test</b></p> <p><b>H<sub>0</sub></b>: <math>\mu_X - \mu_Y = \mu_0</math> <b>HA</b>: <math>\mu_X - \mu_Y \neq \mu_0</math></p> $Z = \frac{(\bar{X} - \bar{Y} - \mu_0)}{\sqrt{\frac{\sigma_X^2}{n_X} + \frac{\sigma_Y^2}{n_Y}}} \sim N(0,1)$ <p><b>Test Statistic</b>:</p> <p><b>Pooled Two-Sample t-Test</b></p> <p><b>H<sub>0</sub></b>: <math>\mu_X - \mu_Y = \mu_0</math> <b>HA</b>: <math>\mu_X - \mu_Y \neq \mu_0</math></p> $T = \frac{(\bar{X} - \bar{Y} - \mu_0)}{S_p \sqrt{\frac{1}{n_X} + \frac{1}{n_Y}}}$ <p><b>Test Statistic</b>: Exact sampling distribution of T under H<sub>0</sub>: <math>T \sim t_{n_X+n_Y-2}</math> (t-distribution with df = n<sub>x</sub> + n<sub>y</sub> - 2)</p> <p><b>Note:s</b> <math>\frac{S_x^2 + S_y^2}{n_x + n_y}</math> is an unbiased estimator of <math>\frac{\sigma_x^2 + \sigma_y^2}{n_x + n_y}</math></p> <p><b>Assumptions</b>: same as Welch's, but If using pooled variance - assume equal variances between populations.</p> <p><b>Unpooled (Welch) Two-Sample t-Test</b></p> <p><b>H<sub>0</sub></b>: <math>\mu_X - \mu_Y = \mu_0</math> <b>HA</b>: <math>\mu_X - \mu_Y \neq \mu_0</math></p> <p><b>Test Statistic</b>:</p> <p><b>Approximate Sampling distribution</b>: <math>T \sim t_{\bar{n}}</math>, where <math>\bar{n} = \left( \frac{s_X^2}{n_X} + \frac{s_Y^2}{n_Y} \right)^2 / \left( \frac{s_X^4}{n_X^2(n_X-1)} + \frac{s_Y^4}{n_Y^2(n_Y-1)} \right)</math></p> <p><b>Assumptions</b>:</p> <ol style="list-style-type: none"> <li>1. Independence between units: (within each population (Y<sub>i</sub> with Y<sub>j</sub> and X<sub>i</sub> with X<sub>j</sub>)) And )between populations (Y<sub>i</sub> with X<sub>j</sub>)</li> <li>2. Homogeneity of units within each population:(Equal means within each population; Equal variances within each population; If using pooled variance - assume equal variances between populations.</li> <li>3. Population distributions are Normal:CLT allows for deviations.</li> <li>4. Random sampling from populations.</li> </ol> <p><b>Paired t-Test</b></p> <p><b>H<sub>0</sub></b>: <math>\mu_D = \mu_0</math> <b>HA</b>: <math>\mu_D \neq \mu_0</math></p> <p><b>Test Statistic</b>: Exact sampling distribution of T under H<sub>0</sub>: <math>T \sim t_{n-1}</math> (where n is # of pairs)</p> <p><b>Assumptions</b> same as one-sample t-test</p>																																													

<p><b>F-test for equal variance</b> (could also use levene's test): (caution: will reject for large samples when ratio is close to 1)  <math>H_0: \sigma_x^2 = \sigma_y^2</math>  <math>H_A: \sigma_x^2 \neq \sigma_y^2 \Leftrightarrow \sigma_x^2 / \sigma_y^2 \neq 1</math></p> <p>Test Statistic: <math>F = \frac{S_x^2}{S_y^2} \sim F_{n_x-1, n_y-1}</math></p> <p>Same assumptions as t-test: In testing the equality of population variances, two assumptions are required: independent samples and normally distributed populations.</p> <p>2-sided p-value is just 2*the one-sided value.</p> <p><b>Test Normality:</b> Shapiro-wilk test, Anderson-Darling test, Kolmogorov-Smirnov Test</p>	<p><b>How does CLT apply?</b>  <b># checking assumptions:</b>  <b>Independence:</b> plot subgroups, and see if relationship still holds (cluster effect)  Plot data over time (serial effects)  Plot data vs space (spatial effect)</p> <p><b>Normality:</b> Histograms should look normal – overlay with normal curve and kernel density plot  Boxplot or Q-Q plot will show outliers</p> <p><b>Equal Variance:</b> divide variances to see if ratio is &lt;0.5 or &gt; 2</p> <p><b>T-test is robust:</b>  -if populations are symmetrically skewed  -if sample sizes are equal (and large)  -when sample sizes are equal, pooled-ttest is robust to unequal variances</p> <p><b>Sensitive:</b> When the sample sizes are not equal, t-tests are more sensitive to skewedness and long-tailedness.  For small samples, t-tests are somewhat sensitive to markedly different skewedness in two groups.  Watch out for outliers.</p> <p><b>Robustness of F-tests:</b>  Not resistant to outliers. Fairly robust to non-normality. Independence w/I and b/w groups is necessary. Equal variance is necessary.</p> <p><b>F-Test Assumptions:</b>  Equal variances is crucial  Not resistant to outliers  Normality is not critical.</p>	<p><b>How does CLT apply?</b>  <b># checking assumptions:</b>  <b>Independence:</b> plot subgroups, and see if relationship still holds (cluster effect)  Plot data over time (serial effects)  Plot data vs space (spatial effect)</p> <p><b>Normality:</b> Histograms should look normal – overlay with normal curve and kernel density plot  Boxplot or Q-Q plot will show outliers</p> <p><b>Equal Variance:</b> divide variances to see if ratio is &lt;0.5 or &gt; 2</p> <p><b>T-test is robust:</b>  -if populations are symmetrically skewed  -if sample sizes are equal (and large)  -when sample sizes are equal, pooled-ttest is robust to unequal variances</p> <p><b>Sensitive:</b> When the sample sizes are not equal, t-tests are more sensitive to skewedness and long-tailedness.  For small samples, t-tests are somewhat sensitive to markedly different skewedness in two groups.  Watch out for outliers.</p> <p><b>Robustness of F-tests:</b>  Not resistant to outliers. Fairly robust to non-normality. Independence w/I and b/w groups is necessary. Equal variance is necessary.</p> <p><b>F-Test Assumptions:</b>  Equal variances is crucial  Not resistant to outliers  Normality is not critical.</p>
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<p><b>Confidence intervals:</b> <math>Y_i \sim \text{iid } N(\mu, \sigma^2)</math></p> $\left( \bar{Y} - t_{1-\alpha/2, n-1} \frac{S}{\sqrt{n}}, \bar{Y} + t_{1-\alpha/2, n-1} \frac{S}{\sqrt{n}} \right)$ <p>One Sample t-test:  Two Sample t-test:</p> <ul style="list-style-type: none"> <li>Two-sample t-tests <ul style="list-style-type: none"> <li><math>\bar{X} - \bar{Y} - t_{1-\alpha/2, df} \cdot SE, \bar{X} - \bar{Y} + t_{1-\alpha/2, df} \cdot SE</math></li> </ul> </li> <li>Pooled t-test: <math>SE = S_p \sqrt{\frac{1}{n_x} + \frac{1}{n_y}}, df = n_x + n_y - 2</math></li> <li>Unpooled t-test: <math>SE = \sqrt{\frac{S_x^2}{n_x} + \frac{S_y^2}{n_y}}, df = \tilde{n}</math>  <math>\tilde{n} = \left( \frac{s_x^2}{n_x} + \frac{s_y^2}{n_y} \right)^2 / \left( \frac{s_x^4}{n_x^2(n_x-1)} + \frac{s_y^4}{n_y^2(n_y-1)} \right)</math></li> </ul> <p><b>Two-Sample t-test for I samples (<math>I &gt; 2</math>) Conf int</b></p> <ul style="list-style-type: none"> <li>Random sampling. <ul style="list-style-type: none"> <li><math>\bar{Y}_m - \bar{Y}_k - t_{1-\alpha/2, df} \cdot SE, \bar{Y}_m - \bar{Y}_k + t_{1-\alpha/2, df} \cdot SE</math></li> </ul> </li> <li>Confidence intervals: <math>SE = S_p \sqrt{\frac{1}{n_m} + \frac{1}{n_k}}, df = n_1 + n_2 + \dots + n_I - I</math></li> </ul>	<p><b>Transformations:</b>  Log Transformation:</p> <ol style="list-style-type: none"> <li>1. <b>Randomized Experiments</b> (T=Treatment, C = Control)  <math>Z_{T,i} = \ln(Y_{T,i}), i = 1, \dots, n_T</math>  <math>Z_{C,i} = \ln(Y_{C,i}), i = 1, \dots, n_C</math> Responses:  <math>Z_{T,i} = Z_{C,i} + \delta \Leftrightarrow \frac{Y_{T,i}}{Y_{C,i}} = e^\delta</math>  <math>\exp(\bar{Z}_T - \bar{Z}_C)</math> estimates <math>\frac{Y_{T,i}}{Y_{C,i}}</math>  Interpretation: The response of an experimental unit to the treatment will be <math>\exp(\bar{Z}_{\text{Treatment}} - \bar{Z}_{\text{Control}})</math> times as large as its response to the control</li> <li>2. Observational studies with <b>random sampling</b>:  <math>\exp(\bar{Z}_2 - \bar{Z}_1)</math> estimates <math>\frac{m_2}{m_1}</math>  Interpretation: <math>\exp(\bar{Z}_{\text{Second Population}} - \bar{Z}_{\text{First Population}})</math> times as large as the median of the second population is</li> <li>3. In paired t-test:  <b>Randomized Experiments</b>:  <math>\exp(\bar{Z})</math> estimates a multiplicative treatment effect <math>\frac{Y_{2,i}}{Y_{1,i}}</math>  <b>Observation studies w/ rand. Sampling</b>:  <math>\text{Median}(Y_{2,i} / Y_{1,i}) = m</math>, then <math>\exp(\bar{Z})</math> estimates <math>m</math>.</li> </ol> <p>Square Root Transformation <math>\text{sqrt}(Y)</math>: Good for moderately skewed data, but hard to interpret results  Reciprocal Transformation <math>1/Y</math>: good for severely skewed data, can be used with negative data  Logit <math>\log(Y/(1-Y))</math> or <math>\text{arcsin}(2Y-1)</math>: good for proportions</p>
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<p><b>Nonparametric alternatives to t-tools:</b></p> <p><b>Rank Sum test:</b>  Hypotheses:  <math>H_0</math>: (if treatments are randomized): Same as fisher's randomization test  <math>H_0</math>: (if treatments are not randomized): same as permutation test</p> <p>* If shapes and spreads of the two populations are similar: HA: There is a difference in medians between the two populations. <math>Z_{1,i} = \text{Rank}(X_i   X_1, \dots, X_{n_x}, Y_1, \dots, Y_{n_y}), i = 1, \dots, n_x</math></p> <p>Transform data:  <math>Z_{2,i} = \text{Rank}(Y_i   X_1, \dots, X_{n_x}, Y_1, \dots, Y_{n_y}), i = 1, \dots, n_y</math></p> <p>Ties are averaged</p> $T = \sum_{i=1}^{n_x} Z_{1,i}, \text{ assuming } n_x \leq n_y$ <p>Test statistic:  Assumptions: Independence</p> <p>Normal approximation to rank Sum test (if no ties, and samples greater than 10):</p> $T \sim N\left(\frac{n_x(n_x + n_y + 1)}{2}, \frac{n_x n_y (n_x + n_y + 1)}{12}\right)$ <p>With ties, Rbar is sample mean (mean of all ranks) and Sr^2 is sample variance for combined set of nx + ny ranks:  <math display="block">T \sim N\left(n_x \bar{R}, S_R^2 \frac{n_x n_y}{n_x + n_y}\right) \quad Z = \frac{T - n_x \bar{R}}{S_R \sqrt{\frac{n_x n_y}{n_x + n_y}}}</math></p> <p>We can use this Z-statistic:</p>	<p><b>Sign test for paired data</b> (lower power than Wilcoxon signed-rank test)</p> $H_0: P(D_i > 0) = P(Y_i > X_i) = 0.5$ <p><b>General formulation:</b> <math>H_a: P(D_i &gt; 0) \neq 0.5</math> (or &gt;0.5, or &lt;0.5).  <math>H_0: \text{Median}(D_i) = 0</math>,  <math>H_a: \text{Median}(D_i) \neq 0</math> (or &gt;0, or &lt;0).</p> <p><b>If differences are continuous rv's:</b> <math>H_0: E(D_i) = 0</math>,  <math>H_a: E(D_i) \neq 0</math></p> <p><b>If differences are symmetric:</b> <math>H_a: E(D_i) \neq 0</math> (or &gt;0, or &lt;0)</p> <p><b>Test Statistic:</b> K = number of positive differences.</p> <p><b>Exact distribution of K:</b> <math>K \sim \text{Binom}(m, 0.5)</math>, where m is final number of pairs with nonzero Di.</p> <p><b>Normal approximation (for large samples):</b></p> <p><b>Assumptions:</b></p> <ol style="list-style-type: none"> <li>The differences <math>D_i</math> are assumed to be independent.</li> <li>Each <math>D_i</math> comes from the same continuous population.</li> <li>The values <math>X_i</math> and <math>Y_i</math> represent are ordered (at least the <a href="#">ordinal scale</a>), so the comparisons "greater than", "less than", and "equal to" are meaningful.</li> </ol>
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<p><b>Wilcoxon Signed-Rank Test for Paired Data:</b> (more assumptions, but higher power than sign test)</p> <p><b>Hypotheses: General formulation:</b></p> <p><b>HO:</b> The rank of the magnitude of within-pair difference is unrelated to the sign of the difference.</p> <p><b>Ha:</b> The rank of the magnitude of within-pair difference is related to the sign of the difference.</p> <p><b>If differences are continuous r.v.'s:</b></p> <p><b>HO:</b> Median(Di)=0</p> <p><b>Ha:</b> Median(Di) ≠ 0 (or &gt;0, or &lt;0)</p> <p><b>If the distribution of differences is symmetric :</b></p> <p><b>HO:</b> E(Di)=0</p> <p><b>Ha:</b> E(Di) ≠ 0 (or &gt;0, or &lt;0)</p> <p>1. Calculate differences (+ or -).</p> <p>2. Rank the absolute values of differences  Di </p> <p>3. <math>Z_i = \text{Rank}( D_i ), i = 1, \dots, n.</math></p> <p>Test Statistic: (sum of ranks for positive differences)</p> $S = \sum_{i=1}^n Z_{i D_i>0}$ <p><b>Exact sampling distribution:</b> do permutation test, and switch group status for each pair.</p> <p><b>Normal approximation</b> (for large sample sizes, m is ≥ 20, where m is the number of non-zero differences):</p> $Z = \frac{S - m(m+1)/4}{\sqrt{m(m+1)(2m+1)/24}} \sim N(0,1)$ <p>Assumptions:</p> <ol style="list-style-type: none"> <li>1. Data are paired and come from the same population.</li> <li>2. Each pair is chosen randomly and independently.</li> <li>3. The data are measured at least on an <u>ordinal scale</u>, but need not be normal.</li> </ol> <hr/> <p><b>Kruskal-Wallis Test</b> (Nonparametric ANOVA)</p> <p><b>HO:</b> Median(Y1) = Median(Y2) = ... = Median(YI)</p> <p><b>HA:</b> at least one median is different from the others</p> <p>Test Statistic – convert to ranks, and do ANOVA on ranks:</p> $KW = \frac{\text{ESS (on ranks)}}{S_R^2} \stackrel{H_0}{\sim} \chi^2_{I-1}$ <p><math>S_R^2</math> is the sample variance of all ranks (not pooled)</p> <p>Assumptions:</p> <ol style="list-style-type: none"> <li>1. Independence.</li> <li>2. Random Sampling</li> </ol>	<p><b>Two-Sample t-Test for I samples</b></p> <p><b>HO:</b> <math>\mu_m - \mu_k = \mu_0</math></p> <p><b>HA:</b> <math>\mu_m - \mu_k \neq \mu_0</math></p> <p><b>Test Statistic:</b></p> $T = \frac{(\bar{Y}_m - \bar{Y}_k - \mu_0)}{S_p \sqrt{\frac{1}{n_m} + \frac{1}{n_k}}}; \quad S_p^2 = \frac{(n_1-1)S_1^2 + (n_2-1)S_2^2 + \dots + (n_I-1)S_I^2}{n_1+n_2+\dots+n_I-I}$ <p>Exact sampling distribution of T under H0 is: <math>T \sim t_{n_1+n_2+\dots+n_I-1}</math>, where I is number of groups.</p> <p><b>Assumptions:</b></p> <ul style="list-style-type: none"> <li>Independence of units within each group;</li> <li>Independence of units between groups;</li> <li>Normality of all I populations;</li> <li>Equal variances for all I populations;</li> <li>Random sampling.</li> </ul> <hr/> <p><b>Anova F-test for Equality of All means in I samples</b></p> <p><b>HO:</b> <math>\mu_1 = \mu_2 = \dots = \mu_I</math> (reduced model – equal means model)</p> <p><b>Ha:</b> at least one mean is different from the others (full model – separate means model)</p> <p><math>SSR_{\text{full}} = SSW</math> (within groups)</p> $S_p^2 = \frac{SSR_{\text{full}}}{n-I}$ $S_p^2 \sim \frac{\sigma^2}{n-I} \chi^2_{n-I} \Rightarrow SSR_{\text{full}} \sim \sigma^2 \chi^2_{n-I}, \text{ d.f.} = n-I$ $S_p^2 = \frac{(n_1-1)S_1^2 + (n_2-1)S_2^2 + \dots + (n_I-1)S_I^2}{n-I}$ <p><math>SSR_{\text{reduced}} = SST</math> (total error)</p> <p><math>S^2</math> is the sample variance of the entire sample, taken as one group:</p> $S^2 = \frac{SSR_{\text{reduced}}}{n-1}$ $S^2 \sim \frac{\sigma^2}{n-1} \chi^2_{n-1} \Rightarrow SSR_{\text{reduced}} \sim \sigma^2 \chi^2_{n-1}, \text{ d.f.} = n-1$ <p><math>ESS = SSB</math> (between groups)</p> <p><math>ESS = SSR_{\text{reduced}} - SSR_{\text{full}}</math></p> $H_0: ESS \sim \sigma^2 \chi^2_{I-1}, \text{ d.f.} = I-1$ <p>Test Statistic:</p> $R = \frac{ESS / (I-1)}{SSR_{\text{full}} / (n-I)} = \frac{MSB}{MSR_{\text{full}}}$ <p>MSB = Mean square between MSR<sub>full</sub> = Mean square residuals (within groups)</p> $S_p^2$ <p>Easy way to calculate: <math>MSR_{\text{full}} = S_p^2</math></p> <p>Exact sampling distribution of R under H0:</p> $F_{(I-1, n-I)}$ <p><b>Assumptions (check w/ Q-Q or residual plot):</b></p> <ol style="list-style-type: none"> <li>1. Normality of populations;</li> <li>2. Equal population variances for all groups;</li> <li>3. Independence within each group;</li> <li>4. Independence between groups;</li> <li>5. Homogeneity within each group;</li> <li>6. Random sampling</li> </ol>	<p><b>F-test for Equality in a subset of groups:</b> "Are groups 2-I all the same?"</p> <p><b>Reduced Model:</b> <math>H_0: \mu_1 = \mu_2 = \mu_3 = \dots = \mu_I</math> (Reduced Model)</p> <p><b>Full Model:</b> <math>H_A: \mu_1, \mu_2, \mu_3, \dots, \mu_I</math> (Full Model)</p> <p><b>Reduced Model:</b> <math>H_0: \text{Equality of means in } k_0 \text{ groups of } I \text{ samples}</math></p> <p><b>Full Model:</b> <math>H_A: \text{Equality of means in } k_s \text{ groups of } I \text{ samples, } k_s &gt; k_0, \text{ such that groups for } H_A \text{ are nested within } H_0 \text{ groups.}</math></p> <p><b>Test statistic:</b></p> $R = \frac{\text{ESS}_{(k_s-k_0)}}{\text{SSR}_{\text{full}} / (n - k_s)} \sim F_{(k_s-k_0, n - k_s)}$ <table border="1" style="width: 100%; border-collapse: collapse;"> <thead> <tr> <th style="text-align: center;">Sum of Squared Residuals</th> <th style="text-align: center;">General Name</th> <th style="text-align: center;">Degrees of freedom</th> <th style="text-align: center;">Distribution</th> <th style="text-align: center;">Corresponding variance (also called "Mean Square")</th> </tr> </thead> <tbody> <tr> <td style="text-align: center;"><math>SSR_{\text{full}}</math> Residual sum of squares (full)</td> <td style="text-align: center;"><math>SSW</math> (Within Groups)</td> <td style="text-align: center;"><math>n - k_A</math></td> <td style="text-align: center;"><math>\sigma^2 \chi^2_{n - k_A}</math></td> <td style="text-align: center;"><math>MSR_{\text{full}}</math> Pooled variance for <math>k_A</math> groups.</td> </tr> <tr> <td style="text-align: center;"><math>ESS</math> Extra sum of squares</td> <td style="text-align: center;"><math>SSB</math> (Between Groups)</td> <td style="text-align: center;"><math>k_A - k_0</math></td> <td style="text-align: center;"><math>\sigma^2 \chi^2_{k_A - k_0}</math></td> <td style="text-align: center;">MSB</td> </tr> <tr> <td style="text-align: center;"><math>SSR_{\text{reduced}}</math> Residual sum of squares (reduced)</td> <td style="text-align: center;"><math>SST</math> (Total)</td> <td style="text-align: center;"><math>n - k_0</math></td> <td style="text-align: center;"><math>\sigma^2 \chi^2_{n - k_0}</math></td> <td style="text-align: center;"><math>MSR_{\text{reduced}}</math> Pooled variance for <math>k_0</math> groups.</td> </tr> </tbody> </table> <p><b>ANOTHER VERSION: F-test for Equality in a subset of groups:</b> "Is group 1 different from all other groups?"</p> <p><b>Reduced Model:</b> <math>H_0: \mu_1 = \mu_2 = \mu_3 = \dots = \mu_I = \mu</math> (Reduced Model)</p> <p><b>Full Model:</b> <math>H_A: \mu_1, \dots, \mu_s = \mu_3 = \mu_4 = \dots = \mu_I = \mu_{\text{Other}}</math> (Full Model)</p> <p>Note: <math>SSR_{\text{full}}</math> now is equal to <math>SSR_{\text{reduced}}</math> from above</p> <hr/> <p><b>Hypothesis test for linear combinations of means:</b> "Is one group mean equal to the average of a combination of other group means?"</p> <p>A linear combination of group means is : <math>\gamma = C_1\mu_1 + C_2\mu_2 + \dots + C_I\mu_I</math>, where C's are fixed coefficients chosen by the researcher</p> <p><math>H_0: \gamma = \gamma_0</math></p> <p><math>H_A: \gamma \neq \gamma_0</math></p> <p>If all coefficients sum to 0, then the linear combination is called a <b>contrast</b>.</p> <p>Estimate of linear combination:</p> $G = C_1\bar{Y}_1 + C_2\bar{Y}_2 + \dots + C_I\bar{Y}_I$ <p>Sampling distribution of G under H0:</p> $G \sim N\left(\gamma_0, \sigma^2 \left( \frac{C_1^2}{n_1} + \frac{C_2^2}{n_2} + \dots + \frac{C_I^2}{n_I} \right) \right)$ <p><b>T-test for Linear combination of means:</b></p> $T = \frac{g - \gamma_0}{SE(g)} = \frac{C_1\bar{Y}_1 + C_2\bar{Y}_2 + \dots + C_I\bar{Y}_I - \gamma_0}{S_p \sqrt{\frac{C_1^2}{n_1} + \frac{C_2^2}{n_2} + \dots + \frac{C_I^2}{n_I}}}$ $S_p^2 = \frac{(n_1-1)S_1^2 + (n_2-1)S_2^2 + \dots + (n_I-1)S_I^2}{n-I}$ <p>(same as F-test s-pooled)</p> <p><b>Approximate sampling distribution:</b> Assumption: Same as ANOVA F-test</p> $T \sim t_{n_1+n_2+\dots+n_I-I}$	Sum of Squared Residuals	General Name	Degrees of freedom	Distribution	Corresponding variance (also called "Mean Square")	$SSR_{\text{full}}$ Residual sum of squares (full)	$SSW$ (Within Groups)	$n - k_A$	$\sigma^2 \chi^2_{n - k_A}$	$MSR_{\text{full}}$ Pooled variance for $k_A$ groups.	$ESS$ Extra sum of squares	$SSB$ (Between Groups)	$k_A - k_0$	$\sigma^2 \chi^2_{k_A - k_0}$	MSB	$SSR_{\text{reduced}}$ Residual sum of squares (reduced)	$SST$ (Total)	$n - k_0$	$\sigma^2 \chi^2_{n - k_0}$	$MSR_{\text{reduced}}$ Pooled variance for $k_0$ groups.				
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$ESS$ Extra sum of squares	$SSB$ (Between Groups)	$k_A - k_0$	$\sigma^2 \chi^2_{k_A - k_0}$	MSB																						
$SSR_{\text{reduced}}$ Residual sum of squares (reduced)	$SST$ (Total)	$n - k_0$	$\sigma^2 \chi^2_{n - k_0}$	$MSR_{\text{reduced}}$ Pooled variance for $k_0$ groups.																						
<p><b>Mulitple Comparisons:</b> Familywise Type-I error</p> $\alpha \leq 1 - (1 - \alpha)^N \leq N\alpha$ <p>N perfectly dependent tests <math>\leq</math> probability of at least one type-I error among N indep. Tests <math>\leq</math> max familywise error</p> <p><b>Bonferroni:</b> (When all else fails) Set individual significance levels at alpha/N, where N is the # of tests For pairwise mean comparisons, half width of CI</p> $= t_{n-I, 1-\alpha/(2N)} SE$	<p><b>Tukey HSD (Honest Significant Difference): (when interested in all pairs of groups)</b></p> <ul style="list-style-type: none"> <li>- Considers the largest difference between any two sample means for I groups</li> <li>- ensures familywise conf leve = 1-alpha</li> </ul> $Q = \frac{\bar{Y}_{\max} - \bar{Y}_{\min}}{S_p / \sqrt{n}} \sim q(I, n-I), \text{ where } n = \bar{n}I$ <p><b>Studentized Range Distribution</b>, nbar is # in each sample, I is # of groups Half-width for CI:</p> $\frac{q(I, n-I, 1-\alpha)}{\sqrt{2}} SE, \text{ where } SE = S_p \sqrt{2/n}$ <p>Assumptions: Normality, equal variances, equal sample sizes</p> <p><b>Tukey-Kramer Procedure</b> (like Tukey HSD, but no need for equal sample sizes):</p> $\frac{q(I, n-I, 1-\alpha)}{\sqrt{2}} SE, \text{ where } SE = S_p \sqrt{\frac{1}{n_i} + \frac{1}{n_j}}$	<p>Formulas:</p> $N \text{ Choose } k: \frac{n!}{k!(n-k)!}$ <p><b>Source of Variation</b>      <b>Sum of Squares</b>      <b>df</b>      <b>Mean Square</b>      <b>F-Statistic</b>      <b>p-value</b></p> <table border="1" style="width: 100%; border-collapse: collapse;"> <thead> <tr> <th style="text-align: center;">Source of Variation</th> <th style="text-align: center;">Sum of Squares</th> <th style="text-align: center;">df</th> <th style="text-align: center;">Mean Square</th> <th style="text-align: center;">F-Statistic</th> <th style="text-align: center;">p-value</th> </tr> </thead> <tbody> <tr> <td style="text-align: center;">Between Groups</td> <td style="text-align: center;">1,927.08</td> <td style="text-align: center;">6</td> <td style="text-align: center;">321.18</td> <td style="text-align: center;">6.72</td> <td style="text-align: center;">.000061</td> </tr> <tr> <td style="text-align: center;">Within Groups</td> <td style="text-align: center;">1,864.45</td> <td style="text-align: center;">39</td> <td style="text-align: center;">47.81</td> <td></td> <td></td> </tr> <tr> <td style="text-align: center;">Total</td> <td style="text-align: center;">3,791.53</td> <td style="text-align: center;">45</td> <td></td> <td></td> <td></td> </tr> </tbody> </table> <p><b>Notes:</b></p> <ul style="list-style-type: none"> <li>① Sum of squared residuals from fitting the full (separate-means) model</li> <li>② Sum of squared residuals from fitting the reduced (equal-means) model</li> <li>③ degrees of freedom</li> <li>④ Subtract the "Within" from the "Total"</li> <li>⑤ A mean square is the ratio of a sum-of-squares to its degrees of freedom</li> <li>⑥ The F-statistic is the ratio of the Between MS to the Within MS</li> <li>⑦ The p-value comes from an F-distribution with 6 and 39 df</li> </ul> <p><b>NOTE:</b> This is <math>s_p^2</math></p>	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			
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<p><b>Log-transformed, observational t-test</b></p> <p>Two Sample t-test</p> <pre>data: log(Salary) by Sex t = -6.1715, df = 91, p-value = 1.849e-08 alternative hypothesis: true difference in means is not equal to 0 95 percent confidence interval: -0.19423950 -0.09964777 sample estimates: mean in group Female    mean in group Male 8.539048                  8.685992</pre> <p>What is the 95% confidence interval for the ratio of population median salaries?  <b>Solution:</b> The 95% CI for the ratio is <math>(e^{-0.194}, e^{-0.0996}) = (0.824, 0.905)</math>.</p>	<p><b>When does pooled t-test rejection rate differ:</b></p> <p>If the variances are actually different, the pooled t-test may be invalid. If the larger variance comes from the group with smaller sample size, the t-test tends to reject the null too often. (One point was taken off if you didn't explain that the rejection rate tends to be too high when the larger variance is in the group with the smaller sample size; the rejection rate tends to be too low when the larger variance is in the group with the larger sample size).</p>	<p>Derive the mean and variance of the signed-rank test statistic under its null hypothesis in a study with <math>N</math> pairs. Show your work/explain your answer: just writing down the correct mean and variance is not sufficient for full credit. [6 points]</p> <p><b>Answer:</b></p> <p>If there are <math>N</math> pairs, for each pair <math>i</math>, denote <math>D_i = 1</math> if 1st observation is larger, and <math>D_i = 0</math> if 2nd observation is larger.</p> <p><math>D_i \sim Ber(1/2)</math> i.i.d.</p> <p>Let <math>T = \text{test stat} = \sum_{i=1}^N D_i * i</math>.</p> $E(T) = E\left(\sum_{i=1}^N D_i * i\right) = \sum_{i=1}^N i * E(D_i) = \frac{1}{2} \sum_{i=1}^N i = \frac{N(N+1)}{4}$ <p>Because all the <math>D_i</math>s are independent of each other,</p> $Var(T) = Var\left(\sum_{i=1}^N D_i * i\right) = \sum_{i=1}^N i^2 * Var(D_i) = \frac{1}{4} \sum_{i=1}^N i^2 = \frac{N(N+1)(2N+1)}{24}$
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$$\frac{\bar{X} - \mu_X}{\frac{\sigma}{\sqrt{n_X}}} \sim N(0, 1), \frac{S_p^2}{\sigma^2} \sim \frac{1}{n_X + n_Y - 2} \chi_{n_X + n_Y - 2}^2.$$

$\frac{\bar{X} - \mu_X}{\frac{\sigma}{\sqrt{n_X}}}$  and  $\frac{S_p^2}{\sigma^2}$  are independent. By definition,  $t_n \sim \frac{N(0,1)}{\sqrt{\chi^2/n}}$ , where the numerator and the denominator are independent. Therefore,

$$\frac{\bar{X} - \mu_X}{S_p / \sqrt{n_X}} = \frac{\bar{X} - \mu_X}{\frac{\sigma}{\sqrt{n_X}}} \frac{1}{\sqrt{\frac{S_p^2}{\sigma^2}}} \sim t_{n_X + n_Y - 2}.$$

Derive test stat for paired t-test

Name	Param.	PMF or PDF	Mean	Variance
Bernoulli	$p$	$P(X = 1) = p, P(X = 0) = q$	$p$	$pq$
Binomial	$n, p$	$\binom{n}{k} p^k q^{n-k}$ , for $k \in \{0, 1, \dots, n\}$	$np$	$npq$
FS	$p$	$pq^{k-1}$ , for $k \in \{1, 2, \dots\}$	$1/p$	$q/p^2$
Geom	$p$	$pq^k$ , for $k \in \{0, 1, 2, \dots\}$	$q/p$	$q/p^2$
NBinom	$r, p$	$\binom{r+n-1}{r-1} p^r q^n, n \in \{0, 1, 2, \dots\}$	$rq/p$	$rq/p^2$
HGeom	$w, b, n$	$\frac{\binom{w}{k} \binom{b}{n-k}}{\binom{w+b}{n}}, \text{ for } k \in \{0, 1, \dots, n\}$	$\mu = \frac{nw}{w+b}$	$(\frac{w+b-n}{w+b-1}) n \frac{\mu}{n} (1 - \frac{\mu}{n})$
Poisson	$\lambda$	$\frac{e^{-\lambda} \lambda^k}{k!}$ , for $k \in \{0, 1, 2, \dots\}$	$\lambda$	$\lambda$
Uniform	$a < b$	$\frac{1}{b-a}$ , for $x \in (a, b)$	$\frac{a+b}{2}$	$\frac{(b-a)^2}{12}$
Normal	$\mu, \sigma^2$	$\frac{1}{\sigma \sqrt{2\pi}} e^{-(x-\mu)^2/(2\sigma^2)}$	$\mu$	$\sigma^2$
Expo	$\lambda$	$\lambda e^{-\lambda x}$ , for $x > 0$	$1/\lambda$	$1/\lambda^2$
Gamma	$a, \lambda$	$\Gamma(a)^{-1} (\lambda x)^a e^{-\lambda x} x^{-1}$ , for $x > 0$	$a/\lambda$	$a/\lambda^2$
Beta	$a, b$	$\frac{\Gamma(a+b)}{\Gamma(a)\Gamma(b)} x^{a-1} (1-x)^{b-1}$ , for $0 < x < 1$	$\mu = \frac{a}{a+b}$	$\frac{\mu(1-\mu)}{a+b+1}$
$\chi^2$	$n$	$\frac{1}{2^{n/2} \Gamma(n/2)} x^{n/2-1} e^{-x/2}$ , for $x > 0$	$n$	$2n$
Student-t	$n$	$\frac{\Gamma((n+1)/2)}{\sqrt{n\pi}\Gamma(n/2)} (1+x^2/n)^{-(n+1)/2}$	0 if $n > 1$	$\frac{n}{n-2}$ if $n > 2$

The function  $\Gamma$  is given by  $\Gamma(a) = \int_0^\infty x^a e^{-x} \frac{dx}{x}$  for all  $a > 0$ . For any  $a > 0$ ,  $\Gamma(a+1) = a\Gamma(a)$ . We have  $\Gamma(n) = (n-1)!$  for  $n$  a positive integer, and  $\Gamma(\frac{1}{2}) = \sqrt{\pi}$ .

Derive distribution for  $T^2$

(5 pts) Let  $T \sim t_{n-1}$ . Derive the distribution of  $T^2$ .

(2 pts for a precise, neat RV set-up) Let us define two independent random variables  $Z$  and  $S^2$ , where  $Z \sim N(0, 1)$ , and  $S^2 \sim \chi_{n-1}^2$ . Then, using the definition by representation,  $Z^2 \sim \chi_1^2$ . The representation of a random variable  $T \sim t_{n-1}$  is

$$T = \frac{Z}{S/\sqrt{n-1}}$$

$$(3 \text{ pts}) \Rightarrow T^2 = \frac{Z^2}{S^2/(n-1)} \sim \frac{\chi_1^2/1}{\chi_{n-1}^2/(n-1)} \sim F_{1,n-1}$$

because  $Z^2$  and  $S^2$  are independent.