Possible Course Flow

- Estimating success probabilities
- Single location: estimates, tests, intervals
- Two locations: testing, estimating differences between locations

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• Scale comparisons and others

- Multiple locations and factors
- Independence
- Nonparametric regression
- Other topics ...

Two-Sample Problem

- Compare two population centers via locations (medians)
- Now, compare scale parameters
- Perhaps same location, perhaps not
- Even more generally, compare two distributions in all respects

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Assumptions

- *X_i*, *i* = 1, 2, . . . , *m* iid
- $Y_i, i = 1, 2, ..., n$ iid
- N = m + n observations
- X_i 's and Y_i 's are independent
- Continuous populations
- F is distribution of X, population 1
- G is distribution of Y, population 2

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- Distribution-free
- Ranks again
- Null:

$$H_0: F(t) = G(t)$$
 for all t

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Same distribution (but no specified)

• Assume same median $\theta_1 = \theta_2$

- Interested in knowing if one distribution has different variability than the other
- Suppose

•
$$F(t) = H\left(\frac{t-\theta_1}{\eta_1}\right)$$

• $G(t) = H\left(\frac{t-\theta_2}{\eta_2}\right)$
• Equivalently: $X \stackrel{d}{=} \eta_1 Z + \theta_1$, $Y \stackrel{d}{=} \eta_2 Z + \theta_2$, with $Z \sim H$

• *H* is continuous with median $0 \Rightarrow F(\theta_1) = G(\theta_2) = 1/2$

- Further assumption: $\theta_1 = \theta_2$ (common median)
- In summary, $\frac{X-\theta}{\eta_1} \stackrel{d}{=} \frac{Y-\theta}{\eta_2}$
- If $\theta_1 \neq \theta_2$, but both are **known**, shift each sample: $X'_i = X_i - \theta_1$, $Y'_i = Y_i - \theta_2$. Now have common median 0

- Look at ratio of scales: $\gamma = \eta_1/\eta_2$
- If variances exist for X and Y, then

$$\gamma^2 = \frac{\operatorname{var}(X)}{\operatorname{var}(Y)}$$

• Write null as

$$H_0: \gamma^2 = 1$$

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- Order the N combined sample values
- Assign 1 to smallest and largest
- Assign 2 to next smallest and next largest

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- Continue ...
- R_j = score assigned to Y_j

•
$$C = \sum_{j=1}^{n} R_j$$
 is the test statistic

One-tail alternative

$$H_1: \gamma^2 > 1$$

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- Reject H_0 if $C \ge c_{\alpha}$
- Table A.8
- Assumes Y is the smaller sample size $(n \le m)$

One-tail alternative

$$H_1:\gamma^2<1$$

- Reject H_0 if $C \leq c_{1-lpha} 1$
- Two-tail alternative

$$H_1: \gamma^2 \neq 1$$

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- Reject H_0 if $C \geq c_{lpha_1}$ or $C \leq c_{1-lpha_2}-1$
- Typically set $\alpha_1 = \alpha_2 = \alpha/2$ (valid for even N due to symmetry)

- Large sample approximation
- When N even:

•
$$E(C) = \frac{n(N+2)}{4}$$

• $var(C) = \frac{mn(N+2)(N-2)}{48(N-1)}$

Null distribution is symmetric $\Rightarrow c_{1-lpha} - 1 = rac{n(N+2)}{2} - c_{lpha}$

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• Otherwise:

•
$$E(C) = \frac{n(N+1)^2}{4N}$$

• $var(C) = \frac{mn(N+1)(N^2+3)}{48N^2}$

• CLT, standardize

$$\mathcal{C}^* = rac{\mathcal{C} - \mathcal{E}(\mathcal{C})}{\sqrt{\mathsf{var}(\mathcal{C})}} \sim \; \mathsf{standard \; \mathsf{normal}}$$

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- Need smaller of *n* and *m* large
- Use $z_{\alpha}, z_{\alpha/2}$

- Continuous \Rightarrow No ties, strictly increasing ranks
- Ties will occur in practice
- Give each group in tie the average of the scores
- Approximately a level- α test
- In large sample approximation, have different value for variance
- If N even,

$$var(C) = rac{mn\left[16\sum_{j=1}^{g}t_{j}r_{j}^{2}-N(N+2)^{2}
ight]}{16N(N-1)}$$

• g is number of groups, t_j is size of group, r_j is average in group

Assumptions

- E(X) E(Y) may not exist
- Only scale difference
- Common median assumption is essential

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• Test $H_0: \gamma^2 = \gamma_0$ with common median θ_0 (known)

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- Use $X'_i = (X_i \theta_0)/\gamma_0$ and $Y'_i = (Y_i \theta_0)$
- Perform test with Y'_i and X'_i

- R
- ansari.test(x, y, exact, conf.int, conf.level)
- Confidence interval (Bauer, Comment 12)
- Estimates the ratio of the scales
- R uses different method when ties cross the center point

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- Medians not equal (or known)
- Previous location-scale model assumption holds (Ansari Bradley)

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- Also assume: $E(V^4) < \infty$ where $V \sim H$
- This assumption implies that γ^2 is ratio of variances
- Uses jackknife method
- More applicable

- Jackknife is a **resampling** method
- Sample the data repeatedly (*without* replacement), form estimates for each sample
- Combine these estimates; solutions are functions of the estimates
- In jackknife, each sample leaves out one particular piece of data
- If there are *n* pieces of data, then there are *n* jackknife samples

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• Sometimes referred to as "leave one out" method

General Jackknife Procedure

- Let $\hat{\theta}$ be the estimate using all the data
- For the *i*-th sample (without using piece *i*), calculate the estimate θ̂_(i) in the same way, *i* = 1,..., n

• Form
$$\tilde{\theta}_i = n\hat{\theta} - (n-1)\hat{\theta}_{(i)}, \ i = 1, \dots, n$$

- The jackknife estimate $\hat{\theta}_J$ is the mean of $\tilde{\theta}_i$, namely, $\sum \tilde{\theta}_i/n$.
- The standard error of this estimate is given by $\sqrt{}$

$$\frac{\sqrt{\sum_{i=1}^{n} (\tilde{\theta}_i - \hat{\theta}_J)^2}}{(n-1)n}$$

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• Why there's an extra factor *n*?

- Get \overline{X}_i , \overline{Y}_j for each jackknife sample
- \overline{X}_i is the sample mean without using data piece *i*
- Also get sample variances for X and Y, leaving out one data piece each time

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• The Miller test will also use full sample mean and sample variance to construct test statistic

$$\overline{X}_i = \frac{1}{m-1} \sum_{s \neq i}^m X_s$$

$$D_i^2 = \frac{1}{m-2} \sum_{s \neq i}^m (X_s - \overline{X}_i)^2$$

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Get \overline{Y}_j and E_j^2 for Y

Set

$$S_i = \ln D_i^2, \quad i = 1, 2, \dots, m$$

$$T_j = \ln E_j^2, \quad j = 1, 2, \dots, n$$

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Also get S_0 and T_0 using all data In: stable the variance, make the statistic more normal

Set

$$A_i = mS_0 - (m-1)S_i, \quad i = 1, 2, \dots, m$$

$$B_j = nT_0 - (n-1)T_j, \quad j = 1, 2, \dots, n$$

$$V_1 = \sum_{i=1}^m \frac{(A_i - \overline{A})^2}{m(m-1)} (\text{estimate var}\overline{A}), \quad V_2 = \sum_{i=1}^m \frac{(B_j - \overline{B})^2}{n(n-1)} (\text{estimate var}\overline{A}).$$

$$Q = \frac{\overline{A} - \overline{B}}{\sqrt{V_1 + V_2}}$$

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• Write null as

$$H_0:\gamma^2=1$$

• One-tail alternative

$$H_1:\gamma^2>1$$

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- Reject H_0 if $Q \ge z_{\alpha}$
- Table A.1 or qnorm

• One-tail alternative

$$H_1:\gamma^2<1$$

- Reject H_0 if $Q \leq -z_{lpha}$
- One-tail alternative

$$H_1:\gamma^2
eq 1$$

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• Reject H_0 if $Q \leq -z_{lpha/2}$ or $Q \geq z_{lpha/2}$

- Asymptotically distribution free
 - F-test: extremely nonrobust
- $\bullet\,$ These are approximately α significance tests
- ullet Asymptotic tests good as sample size $\rightarrow\infty$

- Not a rank test
- No ties

• Estimate of ratio:

$$\tilde{\gamma}^2 = e^{\left\{\overline{A} - \overline{B}
ight\}}$$

• $(1 - \alpha)$ confidence intervals (approximately) • Two-sided

$$\begin{split} \gamma_U^2 &= e^{\left\{\overline{A} - \overline{B} + z_{\alpha/2}\sqrt{V_1 + V_2}\right\}} \\ \gamma_L^2 &= e^{\left\{\overline{A} - \overline{B} - z_{\alpha/2}\sqrt{V_1 + V_2}\right\}} \end{split}$$

• One-sided (lower)

$$\gamma_L^2 = e^{\left\{\overline{A} - \overline{B} - z_\alpha \sqrt{V_1 + V_2}\right\}}$$

• One-sided (upper)

$$\gamma_U^2 = e^{\left\{\overline{A} - \overline{B} + z_\alpha \sqrt{V_1 + V_2}\right\}}$$

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Lepage Test

- Test for either scale or location differences
- Null:

$$H_0: F(t) = G(t)$$
 for all t

Alternative:

$$H_1: \theta_1 \neq \theta_2 \text{ and/or } \eta_1 \neq \eta_2$$

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- Rank test
- Large sample approximation
- Section 5.3 (skip)

- Test for differences in two populations
- Not location, not scale specific
- Assume X and Y independent (within and between samples)

- H_1 : any difference, $F(t) \neq G(t)$ for at least one t
- Commonly used test for Goodness of fit

Set

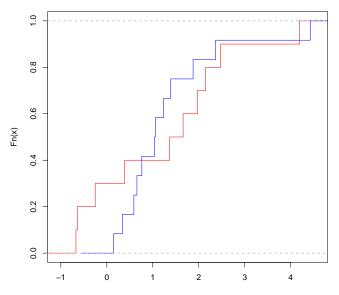
$$F_m(t) = rac{\text{number of sample } X's \leq t}{m}$$

$$G_n(t) = rac{\text{number of sample } Y's \leq t}{n}$$

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 $F_m(t)$ and $G_n(t)$ are empirical distribution functions which are non-decreasing, step functions

ecdf(x)



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$$J = \frac{mn}{d} \max_{-\infty < t < \infty} \{|F_m(t) - G_n(t)|\}$$

where d = greatest common divisor of n and m. Or

$$\mathcal{K}_{m,n}(J^*) = \sqrt{\frac{mn}{m+n}} D_{m,n} = \sqrt{\frac{mn}{m+n}} \max_{-\infty < t < \infty} |\mathcal{F}_m(t) - \mathcal{G}_n(t)|$$

- Order the m + n values as $Z_{(1)}, Z_{(2)}, \ldots, Z_{(m+n)}$
- Sufficient to consider these (finitely many) differences

$$D_{m,n} = \max_{i} |F_m(Z_{(i)}) - G_n(Z_{(i)})|$$

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Null:

$$H_0: F(t) = G(t)$$
 for all t

Alternative:

 $H_1: F(t) \neq G(t)$ for at least one t

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- Reject H_0 if $J \ge j_{\alpha}$
- Table A.10 (X smaller)

• Large sample approximation $J^* = \frac{Jd}{\sqrt{mnN}}$, or in fact

$$K_{m,n} = \sqrt{\frac{mn}{m+n}} D_{m,n}$$

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• Reject
$$H_0$$
 if $J^* \ge q^*_{lpha}$

•
$$P(J^* \ge q^*_\alpha) = \alpha$$

- Limiting distribution: *Kolmogorov distribution* (cf. (5.74) P179)
- Table A.11
- Not normal
- No ties

- R
- ks.test
- Can test X and Y, or,
- Test X against a particular distribution
- pnorm(mean, sd), pexp(mean), etc.

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(Skip 5.5)