

Comparing two samples (Chapter 11)

• Comparing two independent samples

• Problem formulation and statistical modeling

Data

| \underline{U} | \underline{V} |
|-----------------|-----------------|
| 1 | X_1 |
| \vdots | \vdots |
| 1 | X_n |
| 2 | Y_1 |
| \vdots | \vdots |
| 2 | Y_m |

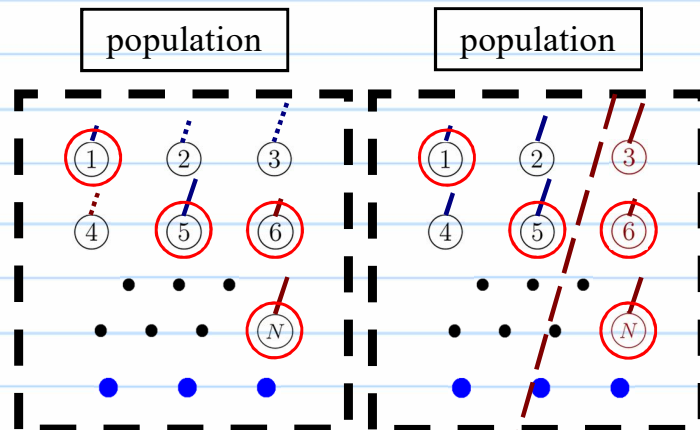
observed data from s.r.s
(random variables)

$$\{X_1, \dots, X_n\}$$

$$\{Y_1, \dots, Y_m\}$$

- X_i 's, Y_j 's are continuous quantities of same characteristic
- $X-Y$ is meaningful

s.r.s., $N \rightarrow \infty$:
without replacement
 \approx with replacement
(\Rightarrow i.i.d.)



For example, in
medical study,

- X_i 's: treatment
- Y_j 's: control

For example, in human
population,

- X_i 's: heights of males
- Y_j 's: heights of females

Why?

- $X_1, \dots, X_n \sim$ i.i.d. with a common continuous distribution F
- $Y_1, \dots, Y_m \sim$ i.i.d. from a common continuous distribution G
- $\{X_1, \dots, X_n\}$ and $\{Y_1, \dots, Y_m\}$ are independent

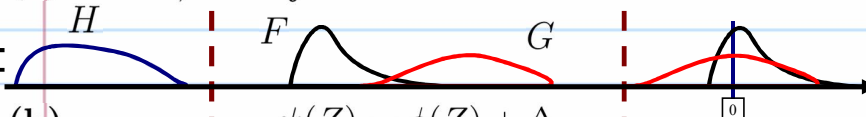


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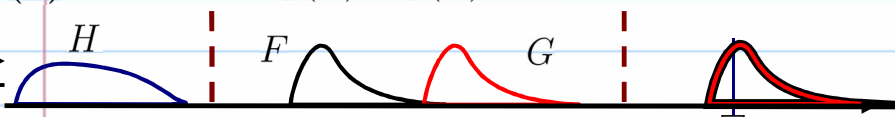
Ch 11, p. 2

- Let random variables Z_1, \dots, Z_n and Z_{n+1}, \dots, Z_{n+m} represent the variability of the $n + m$ members sampled from the population.
- Assume Z_1, \dots, Z_{n+m} are i.i.d. from a population distribution H .
- Let F and G be the distributions of $X = \phi(Z)$ and $Y = \psi(Z)$, respectively.
- The transformations ϕ and ψ might contain random components, e.g., $\phi(Z) = \phi^*(Z) + \delta$, where ϕ^* : a fixed function and δ : a random variable.
- Let μ_X and μ_Y be the means of F and G , respectively.

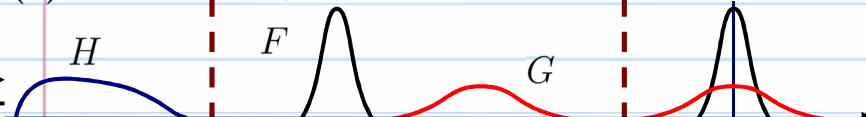
(a) F, G : any continuous distribution



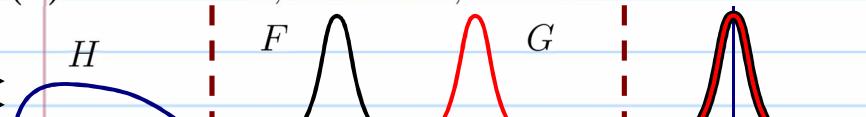
(b) $\psi(Z) = \phi(Z) + \Delta$



(c) F, G : normal



(d) F, G : normal, same σ^2



- Let $X_i = \phi(Z_i)$, $i = 1, \dots, n$,
 $\Rightarrow X_i \sim F$
- Let $Y_j = \psi(Z_{n+j})$, $j = 1, \dots, m$,
 $\Rightarrow Y_j \sim G$
- $X_1, \dots, X_n, Y_1, \dots, Y_m$ are independent

- $X_i = \mu_X + \epsilon_{1,i}$, $i = 1, \dots, n$,
- $Y_j = \mu_Y + \epsilon_{2,j}$, $j = 1, \dots, m$,
- $E(\epsilon) = 0$ and ϵ 's are independent

Note 1 (Some notes about comparing several samples)

- The samples are drawn under different conditions, and inferences must be made about possible effects of these conditions, e.g., treatment and control groups.
- Two-sample comparison (and ANOVA, multiple comparison):
 - methods for comparing samples from distributions that may be different
 - methods for making inference about how the distributions differ
- This chapter (and next chapter) will be concerned with analyzing measurements that are continuous in nature, e.g., temperature.

Example 1 (heat of fusion of ice, Natrella, 1963)

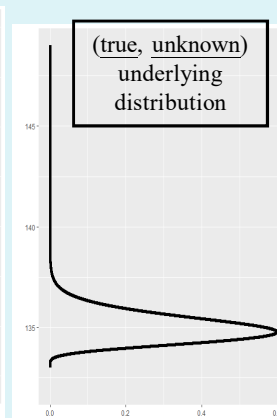
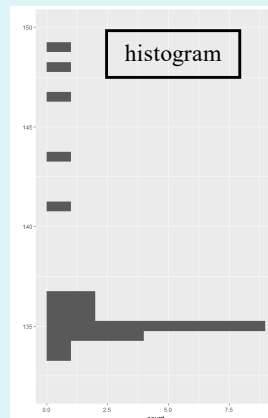
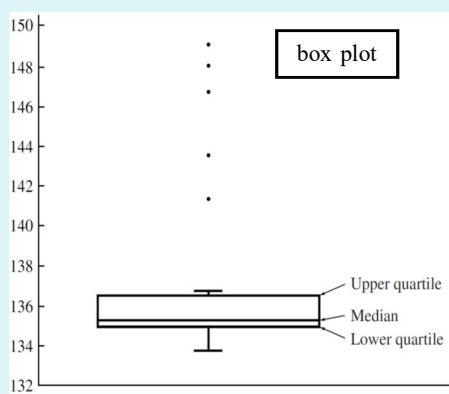
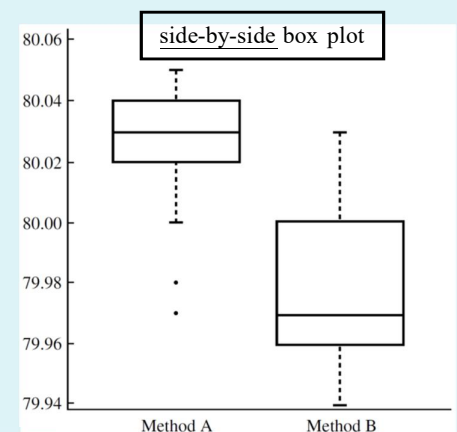
- Two methods, *A* and *B*, were used in a determination of the latent heat of fusion of ice.
- The investigators wished to find out how the methods differed.
- The following table gives the change in total heat from ice at -0.72°C to water 0°C in calories per gram of mass:

| | | | | | | | | |
|-----------------|-------|-------|-------|-------|-------|-------|-------|-------|
| Method <i>A</i> | 79.98 | 80.04 | 80.02 | 80.04 | 80.03 | 80.03 | 80.04 | 79.97 |
| | 80.05 | 80.03 | 80.02 | 80.00 | 80.02 | | | |
| Method <i>B</i> | 80.02 | 79.94 | 79.98 | 79.97 | 79.97 | 80.03 | 79.95 | 79.97 |

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Definition 1 (box plot)

- horizontal lines
 - at median, upper and lower quartiles (50%, 75%, 25% quantiles)
 - IQR = upper quartile – lower quartile
- vertical lines: from upper (or lower) quartile to the most extreme data point that is within a distance of $1.5 \times IQR$ of the upper (or lower) quartile
- each data point beyond the ends of the vertical lines is marked with an asterisk or dot (might be regarded as possible outliers)



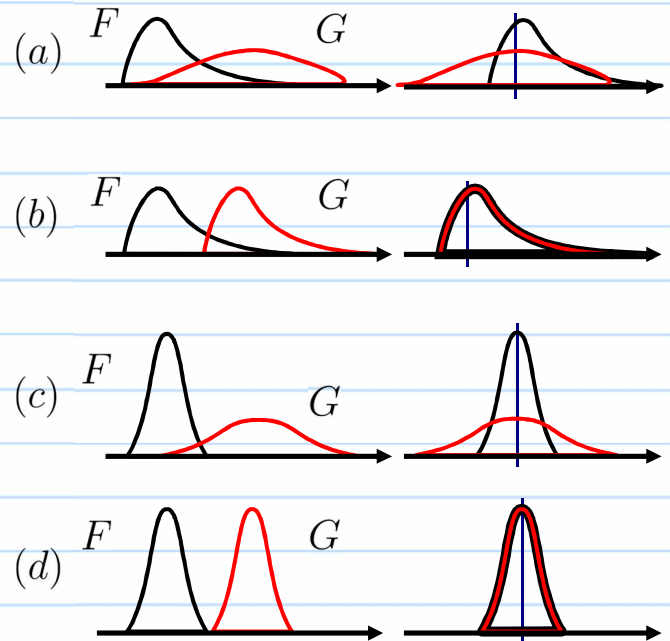
Question 1.

Q: How to define “two samples are identical” or “two samples are different”?

- Notice the distinction between “two identical random variables” ($X = Y$) and “two random variables with an identical distribution” ($X \sim Y$)
- In the statistical modeling of two-sample problem, $X \sim F$ and $Y \sim G$

Consider the different modelings in LNp.2,

- For (a), $F = G$ vs. $F \neq G$
- For (a), $\mu_X = \mu_Y$ vs. $\mu_X \neq \mu_Y$
- For (a), $\tilde{\mu}_X = \tilde{\mu}_Y$ vs. $\tilde{\mu}_X \neq \tilde{\mu}_Y$
($\tilde{\mu}$: median)
- For (b), $\Delta = 0$ vs. $\Delta \neq 0$
- For (c), $\mu_X = \mu_Y$ and $\sigma_X^2 = \sigma_Y^2$ vs.
 $\mu_X \neq \mu_Y$ or $\sigma_X^2 \neq \sigma_Y^2$
- For (c), $\mu_X = \mu_Y$ (i.e., different variances are allowed) vs. $\mu_X \neq \mu_Y$
- For (c), $\sigma_X^2 = \sigma_Y^2$ (i.e., different means are allowed) vs. $\sigma_X^2 \neq \sigma_Y^2$
- For (d), $\mu_X = \mu_Y$ vs. $\mu_X \neq \mu_Y$



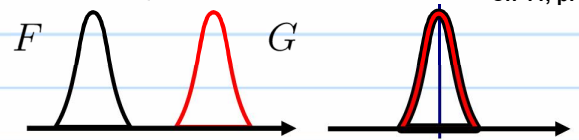
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• **Methods based on normality assumptions**

- Assume that (1) F and G are normal, and (2) F and G have same variance.
- Thus, the statistical model is:

$$\left. \begin{array}{l} \text{1st sample: } X_1, \dots, X_n \sim \text{i.i.d. } N(\mu_X, \sigma^2) \\ \text{2nd sample: } Y_1, \dots, Y_m \sim \text{i.i.d. } N(\mu_Y, \sigma^2) \end{array} \right\} \Leftarrow \text{independent } (*)$$

- This model contains three parameters: $\mu_X (\in \mathbb{R})$, $\mu_Y (\in \mathbb{R})$, $\sigma^2 (> 0)$.
- Under this model, the “difference” between F and G is simplified to be the difference between μ_X and μ_Y , i.e., $\Delta \equiv \mu_X - \mu_Y$ (\Leftarrow called “effect”), and
 $\mu_X - \mu_Y = 0 \Leftrightarrow$ no difference or no effect



Review 1 (estimation of the parameters in one-sample normal model)

Consider $X_1, \dots, X_n \sim \text{i.i.d. } N(\mu, \sigma^2)$, and the statistics

$$\bar{X} = \frac{1}{n} \sum_{i=1}^n X_i \xrightarrow{e} \mu \quad \text{and} \quad s_X^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2 \xrightarrow{e} \sigma^2$$

- distribution (**exercise**)
 - \bar{X} and s_X^2 are independent
 - $\bar{X} \sim N(\mu, \sigma^2/n) \Rightarrow \sqrt{n}(\bar{X} - \mu)/\sigma \sim N(0, 1)$
 - $(n-1)s_X^2 \sim \sigma^2 \chi_{n-1}^2 \Rightarrow (n-1)s_X^2/\sigma^2 \sim \chi_{n-1}^2$; $n-1$: degrees of freedom

- $(T_1 = \sum_{i=1}^n X_i, T_2 = \sum_{i=1}^n X_i^2)$ is a sufficient and complete statistic (**exercise**, **Hint**. 2-parameter exponential family)
- Optimality
 - \bar{X} is the uniformly minimum variance unbiased estimator (UMVUE) of μ (**exercise**, **Hint**. Lehmann-Scheffe Thm)
 - \bar{X} is the maximum likelihood estimator (MLE) of μ (**exercise**, **Hint**.

$$\text{log-likelihood} \propto -\frac{n}{2} \log(\sigma^2) - \frac{\sum_{i=1}^n (X_i - \mu)^2}{2\sigma^2}$$
)
 - s_X^2 is the UMVUE of σ^2 (**exercise**, **Hint**. Lehmann-Scheffe Thm)
 - The MLE of σ^2 is $\frac{n-1}{n} s_X^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \bar{X})^2$ (**exercise**)

Definition 1 (estimators of the parameters in the 2-sample normal model)

Under the two-sample normal model (*) in LNp.6,

- an intuitive estimator of μ_X is $\bar{X} = \frac{1}{n} \sum_{i=1}^n X_i$,
- an intuitive estimator of μ_Y is $\bar{Y} = \frac{1}{m} \sum_{j=1}^m Y_j$,
- since $s_X^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$ and $s_Y^2 = \frac{1}{m-1} \sum_{j=1}^m (Y_j - \bar{Y})^2$ estimate the same parameter σ^2 , we can pool them to get a better estimator:

$$s_p^2 = \frac{(n-1)}{(n-1) + (m-1)} s_X^2 + \frac{(m-1)}{(n-1) + (m-1)} s_Y^2 = \frac{(n-1)s_X^2 + (m-1)s_Y^2}{m+n-2}.$$

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Note 1 (Some notes about the estimator of σ^2)

- s_p^2 is called the **pooled sample variance**
- s_p^2 is a weighted average of the sample variances of the X_i 's and Y_j 's, where
 - the weights are proportional to the degrees of freedom, it is appropriate since if one sample is of much larger size than the other, the estimate of σ^2 from that sample is more reliable \Rightarrow it receives greater weight
 - since $E(s_X^2) = \sigma^2$ and $E(s_Y^2) = \sigma^2 \Rightarrow s_p^2$: an unbiased estimator of σ^2

Theorem 1 (distributions of the parameter estimators, 2-sample normal model)

- Since $(X_1, \dots, X_n), (Y_1, \dots, Y_m)$ are independent random variables
 $\Rightarrow (\bar{X}, s_X^2, \bar{Y}, s_Y^2)$ are independent random variables
 $\Rightarrow (\bar{X}, \bar{Y}, s_p^2)$ are independent random variables
- $\bar{X} \sim N(\mu_X, \sigma^2/n) \Rightarrow \sqrt{n}(\bar{X} - \mu_X)/\sigma \sim N(0, 1)$
- $\bar{Y} \sim N(\mu_Y, \sigma^2/m) \Rightarrow \sqrt{m}(\bar{Y} - \mu_Y)/\sigma \sim N(0, 1)$
- $\bar{X} - \bar{Y} \sim N\left(\mu_X - \mu_Y, \frac{\sigma^2}{n} + \frac{\sigma^2}{m}\right) \Rightarrow \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sigma \sqrt{\frac{1}{n} + \frac{1}{m}}} \sim N(0, 1)$
- Since (i) $(n-1)s_X^2/\sigma^2 \sim \chi_{n-1}^2$, (ii) $(m-1)s_Y^2/\sigma^2 \sim \chi_{m-1}^2$, and (iii) s_X^2 and s_Y^2 are independent,

$$\frac{(n-1)s_X^2 + (m-1)s_Y^2}{\sigma^2} = \frac{(m+n-2)s_p^2}{\sigma^2} \sim \chi_{m+n-2}^2$$

Theorem 2 (log-likelihood, 2-sample normal model)

Under the two-sample normal model (*) in LNp.6, the log-likelihood is proportional to (exercise)

$$\begin{aligned}
 l(\mu_X, \mu_Y, \sigma^2) &\propto -\frac{m+n}{2} \log(\sigma^2) - \frac{\sum_{i=1}^n (X_i - \mu_X)^2}{2\sigma^2} - \frac{\sum_{j=1}^m (Y_j - \mu_Y)^2}{2\sigma^2} \\
 &= -\frac{1}{2\sigma^2} \left(\sum_{i=1}^n X_i^2 + \sum_{j=1}^m Y_j^2 \right) + \frac{\mu_X}{\sigma^2} \left(\sum_{i=1}^n X_i \right) + \frac{\mu_Y}{\sigma^2} \left(\sum_{j=1}^m Y_j \right) \\
 &\quad - [(m+n)/2] \log(\sigma^2) - (n\mu_X^2)/(2\sigma^2) - (m\mu_Y^2)/(2\sigma^2) \\
 &\in \text{3-parameter exponential family}
 \end{aligned}$$

From the log-likelihood, we have

- $\frac{\partial l}{\partial \mu_X} = \frac{1}{\sigma^2} [(\sum_{i=1}^n X_i) - n \times \mu_X]$
- $\frac{\partial l}{\partial \mu_Y} = \frac{1}{\sigma^2} [(\sum_{j=1}^m Y_j) - m \times \mu_Y]$
- $\frac{\partial l}{\partial \sigma^2} = -\frac{m+n}{2\sigma^2} + \frac{\sum_{i=1}^n (X_i - \mu_X)^2}{2\sigma^4} + \frac{\sum_{j=1}^m (Y_j - \mu_Y)^2}{2\sigma^4}$

Theorem 3 (UMVUE and MLE of the parameters in the 2-sample normal model)

- $(R_1 = \sum_{i=1}^n X_i, R_2 = \sum_{j=1}^m Y_j, R_3 = \sum_{i=1}^n X_i^2 + \sum_{j=1}^m Y_j^2)$ is a sufficient and complete statistic (**Hint.** 3-parameter exponential family)
- $\bar{X} (= R_1/n)$ is the UMVUE (by Lehmann-Scheffe Thm) and MLE of μ_X
- $\bar{Y} (= R_2/m)$ is the UMVUE (by Lehmann-Scheffe Thm) and MLE of μ_Y

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- (by Lehmann-Scheffe Thm) The pooled sample variance s_p^2 is the UMVUE of σ^2 , since (i) s_p^2 is unbiased, and (ii)

$$\begin{aligned}
 (m+n-2)s_p^2 &= (n-1)s_X^2 + (m-1)s_Y^2 \\
 &= \sum_{i=1}^n (X_i - \bar{X})^2 + \sum_{j=1}^m (Y_j - \bar{Y})^2 \\
 &= \left(\sum_{i=1}^n X_i^2 \right) - n\bar{X}^2 + \left(\sum_{j=1}^m Y_j^2 \right) - m\bar{Y}^2 = R_3 - (R_1^2/n) - (R_2^2/m)
 \end{aligned}$$
- The MLE of σ^2 is $\frac{m+n-2}{m+n} s_p^2 = \frac{(n-1)s_X^2 + (m-1)s_Y^2}{m+n}$

Question 2 (how to claim $\Delta=0$ or $\Delta \neq 0$?)

Under the two-sample normal model (*) in LNp.6, consider the parameter

$$\Delta = \mu_X - \mu_Y.$$

Notice that

$$\Delta = 0 \Leftrightarrow \text{no difference in the two samples}$$

- The UMVUE (by Lehmann-Scheffe Thm and $\hat{\Delta} = R_1/n - R_2/m$) and MLE of Δ is $\hat{\Delta} = \bar{X} - \bar{Y}$.
- But, $\hat{\Delta} \neq 0$ is not a strong enough evidence to reject $\Delta = 0$ (**Note.** $P(\hat{\Delta} \neq 0) = 1$). A better way is to examine if a C.I. of Δ contains 0.
- **Q:** how to construct an interval estimator for Δ ?

Review 2 (pivotal quantity of θ)

A **pivotal quantity** for θ is a function of data X_1, \dots, X_n and the parameter θ , denoted by

$$Q(\mathbf{X}, \theta) = Q(X_1, \dots, X_n, \theta),$$

if the distribution of $Q(\mathbf{X}, \theta)$ is irrelevant to *all* parameters.

Theorem 4 (confidence interval of Δ , 2-sample normal model)

Under the two-sample normal model (*) in LNp.6,

- σ^2 known (σ^2 is not a parameter)

– a pivotal quantity of Δ is

$$Q_{Z,\Delta} = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sigma \sqrt{\frac{1}{n} + \frac{1}{m}}} = \frac{(\bar{X} - \bar{Y}) - \Delta}{\sigma \sqrt{\frac{1}{n} + \frac{1}{m}}} \sim N(0, 1)$$

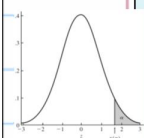
– a $100(1 - \alpha)\%$ C.I. for Δ is $(\bar{X} - \bar{Y}) \pm z(\alpha/2) \times (\sigma \sqrt{\frac{1}{n} + \frac{1}{m}})$ since
 $1 - \alpha = P(|Q_{Z,\Delta}| < z(\alpha/2))$

$$= P\left((\bar{X} - \bar{Y}) - z(\alpha/2)\sigma \sqrt{\frac{1}{n} + \frac{1}{m}} < \Delta < (\bar{X} - \bar{Y}) + z(\alpha/2)\sigma \sqrt{\frac{1}{n} + \frac{1}{m}}\right)$$

- σ^2 unknown (σ^2 is a parameter)

– a pivotal quantity of Δ is

$$Q_{T,\Delta} = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{s_p \sqrt{\frac{1}{n} + \frac{1}{m}}} = \frac{[(\bar{X} - \bar{Y}) - \Delta] / (\sigma \sqrt{\frac{1}{n} + \frac{1}{m}})}{\sqrt{\left[\frac{(m+n-2)s_p^2}{\sigma^2}\right] \frac{1}{m+n-2}}} \sim t_{m+n-2}$$



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– a $100(1 - \alpha)\%$ C.I. for Δ is $(\bar{X} - \bar{Y}) \pm t_{m+n-2}(\alpha/2) \times \left(s_p \sqrt{\frac{1}{n} + \frac{1}{m}}\right)$

Note 2 (A note about the confidence intervals of Δ)

These confidence intervals are of the form

$$(\text{estimate}) \pm (\text{critical value}) \times [(\text{estimated}) \text{ standard error}],$$

where the (estimated) standard error of $\bar{X} - \bar{Y}$ is $\sigma_{\bar{X}-\bar{Y}} = \sigma \sqrt{\frac{1}{n} + \frac{1}{m}}$ when σ^2 is known, and is $s_{\bar{X}-\bar{Y}} = s_p \sqrt{\frac{1}{n} + \frac{1}{m}}$ when σ^2 is unknown.

Example 2 (confidence interval of Δ , heat of fusion of ice, cont. Ex.1 in LNp.3)

- $n = 13, \bar{X}_A = 80.02, s_A = 0.024; \quad m = 8, \bar{X}_B = 79.98, s_B = 0.031$

$$\bullet \quad s_p = \sqrt{\frac{12}{19} s_A^2 + \frac{7}{19} s_B^2} = 0.027, \quad s_{\bar{X}_A - \bar{X}_B} = s_p \sqrt{\frac{1}{13} + \frac{1}{8}} = 0.012$$

- A 95% confidence interval for $\Delta = \mu_A - \mu_B$ is

$$(\bar{X}_A - \bar{X}_B) \pm t_{19}(0.025) \times s_{\bar{X}_A - \bar{X}_B} = (0.04) \pm (2.093) \times (0.012) = (0.015, 0.065).$$

Question 3 (how to perform testing of $\Delta=0$?)

- **Recall.** duality between confidence interval and hypothesis testing
- **Q:** What are the hypothesis testings corresponding to these confidence intervals of Δ ?

Theorem 5 (z -test and t -test for $\Delta = \Delta_0$, 2-sample normal model)

Under the two-sample normal model (*) in LNp.6, consider the null and alternative hypotheses: $H_0 : \mu_X - \mu_Y = \Delta = \Delta_0$ vs. $H_A : \mu_X - \mu_Y = \Delta \neq \Delta_0$

where Δ_0 is a known constant (**Note.** if $\Delta_0 = 0$, $H_0 : \mu_X = \mu_Y$ vs. $H_A : \mu_X \neq \mu_Y$), and H_A is a **two-sided alternative**. From the duality between C.I. and testing,

$$|Q_{Z_0, \Delta}| < z(\alpha/2) \quad \xleftrightarrow{\text{cf.}} \quad |Q_{Z, \Delta_0}| < z(\alpha/2)$$

$$|Q_{T_0, \Delta}| < t_{m+n-2}(\alpha/2) \quad \xleftrightarrow{\text{cf.}} \quad |Q_{T, \Delta_0}| < t_{m+n-2}(\alpha/2)$$

the corresponding test of these confidence intervals are:

- test statistic

$$- \sigma^2 \text{ known: } Z = \frac{(\bar{X} - \bar{Y}) - \Delta_0}{\sigma \sqrt{\frac{1}{n} + \frac{1}{m}}} \quad \left(\xleftrightarrow{\text{cf.}} Q_{Z, \Delta} \text{ in LNp.11} \right)$$

$$- \sigma^2 \text{ unknown: } T = \frac{(\bar{X} - \bar{Y}) - \Delta_0}{s_p \sqrt{\frac{1}{n} + \frac{1}{m}}} \quad \left(\xleftrightarrow{\text{cf.}} Q_{T, \Delta} \text{ in LNp.11} \right)$$

- null distribution

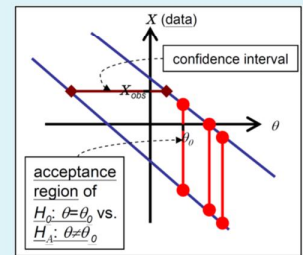
$$- \sigma^2 \text{ known: under } H_0, Z \sim N(0, 1)$$

$$- \sigma^2 \text{ unknown: under } H_0, T \sim t_{m+n-2}$$

- level- α rejection region

$$- \sigma^2 \text{ known: } |Z| > z(\alpha/2), \text{ called } z\text{-test} \quad (\text{reasonable?})$$

$$- \sigma^2 \text{ unknown: } |T| > t_{m+n-2}(\alpha/2), \text{ called } t\text{-test} \quad (\text{reasonable?})$$



Note. The t -test (or z -test) rejects H_0 if and only if its corresponding C.I. does not include Δ_0 .

Note 3 (Some notes about z - and t -tests)

- For the null and alternative hypotheses:

$$\text{or } H_0 : \Delta = \Delta_0 \text{ (or } \Delta \leq \Delta_0) \text{ vs. } H_A^* : \Delta > \Delta_0 \text{ (need domain knowledge)}$$

$$\text{or } H_0 : \Delta = \Delta_0 \text{ (or } \Delta \geq \Delta_0) \text{ vs. } H_A^{**} : \Delta < \Delta_0 \text{ (need domain knowledge)}$$

where H_A^* and H_A^{**} are **one-sided alternatives**, the z - and t -tests are

$$- \sigma^2 \text{ known: } Z > z(\alpha) \text{ for } H_A^*, \text{ and } Z < -z(\alpha) \text{ for } H_A^{**}$$

$$- \sigma^2 \text{ unknown: } T > t_{m+n-2}(\alpha) \text{ for } H_A^*, \text{ and } T < -t_{m+n-2}(\alpha) \text{ for } H_A^{**}$$

- FYI.** All the tests presented in LNp.13-14 are uniformly most powerful unbiased (UMPU) tests. (**Note.** Its proof follows a theorem of UMPU tests for exponential family with *nuisance parameters*)

- The test statistics are of the form:

$$\frac{(\bar{X} - \bar{Y}) - \Delta_0}{s_{\bar{X} - \bar{Y}}}$$

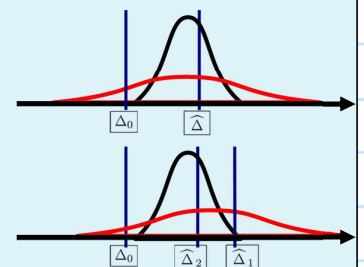
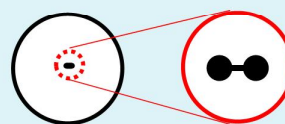
$$- \text{In the numerator, } (\bar{X} - \bar{Y}) - \Delta_0 \text{ estimates } \Delta - \Delta_0.$$

$$- \text{Q: why is this estimate divided by } s_{\bar{X} - \bar{Y}} \text{ (} s_{\bar{X} - \bar{Y}} \downarrow \text{ when } m \uparrow \text{ and/or } n \uparrow \text{)?}$$

- Q:** if H_0 not rejected, do we really accept $\Delta = \Delta_0$, say $\mu_X = \mu_Y$? (better to claim “sample size is not large enough to reject H_0 .”)

- statistically significant difference vs. physically significant difference (example?)

statistical standard \longleftrightarrow physical standard



Theorem 6 (likelihood ratio tests for $\Delta=\Delta_0$, 2-sample normal model)

All the tests presented in LNp.13-14 are likelihood ratio tests.

Proof: We only prove the case of two-sided hypothesis. For the case of one-sided hypothesis, its proof is similar (**exercise**).

• Recall that

– the log-likelihood is

$$l = \log(\mathcal{L}) \propto -\frac{m+n}{2} \log(\sigma^2) - \frac{\sum_{i=1}^n (X_i - \mu_X)^2}{2\sigma^2} - \frac{\sum_{j=1}^m (Y_j - \mu_Y)^2}{2\sigma^2},$$

– the test statistic of likelihood ratio test is

$$\Lambda = \frac{\sup_{\omega} \mathcal{L}}{\sup_{\Omega} \mathcal{L}} \quad \text{or} \quad \log(\Lambda) = \sup_{\omega} \log(\mathcal{L}) - \sup_{\Omega} \log(\mathcal{L}) = \sup_{\omega} l - \sup_{\Omega} l,$$

where $\Omega = H_0 \cup H_A$ and $\omega = H_0$,

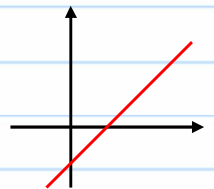
– a likelihood ratio test rejects H_0 for small values of Λ (or $\log \Lambda$).

• σ^2 known

– The parameter spaces Ω and ω are

$$\Omega = \{(\mu_X, \mu_Y) \mid \mu_X \in \mathbb{R}, \mu_Y \in \mathbb{R}\}$$

$$\omega = \{(\mu_X, \mu_Y) \mid \mu_X \in \mathbb{R}, \mu_Y = \mu_X - \Delta_0\}$$



– Under Ω , the MLE's of (μ_X, μ_Y) are

$$\hat{\mu}_{X,\Omega} = \bar{X}, \quad \hat{\mu}_{Y,\Omega} = \bar{Y},$$

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$$\Rightarrow \sup_{\Omega} l = l(\hat{\mu}_{X,\Omega}, \hat{\mu}_{Y,\Omega}) \propto -\frac{\sum_{i=1}^n (X_i - \bar{X})^2 + \sum_{j=1}^m (Y_j - \bar{Y})^2}{2\sigma^2} - \frac{m+n}{2} \log(\sigma^2)$$

– Under ω , the log-likelihood is proportional to

$$-\frac{m+n}{2} \log(\sigma^2) - \frac{\sum_{i=1}^n (X_i - \mu_X)^2}{2\sigma^2} - \frac{\sum_{j=1}^m [Y_j - (\mu_X - \Delta_0)]^2}{2\sigma^2},$$

and the MLE's of (μ_X, μ_Y) are

$$\hat{\mu}_{X,\omega} = \frac{n}{m+n} \bar{X} + \frac{m}{m+n} (\bar{Y} + \Delta_0),$$

$$\hat{\mu}_{Y,\omega} = \hat{\mu}_{X,\omega} - \Delta_0 = \frac{n}{m+n} (\bar{X} - \Delta_0) + \frac{m}{m+n} \bar{Y},$$

$$\Rightarrow \sup_{\omega} l = l(\hat{\mu}_{X,\omega}, \hat{\mu}_{Y,\omega}) \propto -\frac{\sum_{i=1}^n (X_i - \hat{\mu}_{X,\omega})^2 + \sum_{j=1}^m (Y_j - \hat{\mu}_{Y,\omega})^2}{2\sigma^2}$$

– Therefore, the log-likelihood-ratio is

$$\log(\Lambda) = l(\hat{\mu}_{X,\omega}, \hat{\mu}_{Y,\omega}) - l(\hat{\mu}_{X,\Omega}, \hat{\mu}_{Y,\Omega})$$

$$= -\frac{1}{2\sigma^2} \left[\left(\sum_{i=1}^n X_i^2 - 2n\bar{X}\hat{\mu}_{X,\omega} + n\hat{\mu}_{X,\omega}^2 + \sum_{j=1}^m Y_j^2 - 2m\bar{Y}\hat{\mu}_{Y,\omega} + m\hat{\mu}_{Y,\omega}^2 \right) - \left(\sum_{i=1}^n X_i^2 - n\bar{X}^2 + \sum_{j=1}^m Y_j^2 - m\bar{Y}^2 \right) \right]$$

$$= -\frac{1}{2\sigma^2} [n(\bar{X} - \hat{\mu}_{X,\omega})^2 + m(\bar{Y} - \hat{\mu}_{Y,\omega})^2]$$

$$= -\frac{1}{2\sigma^2} \left(\frac{mn}{m+n} \right) (\bar{X} - \bar{Y} - \Delta_0)^2$$

$$\bar{X} - \hat{\mu}_{X,\omega} = \frac{m}{m+n} (\bar{X} - \bar{Y} - \Delta_0)$$

$$\bar{Y} - \hat{\mu}_{Y,\omega} = \frac{-n}{m+n} (\bar{X} - \bar{Y} - \Delta_0)$$

- The likelihood ratio test rejects H_0 for

$$\text{small values of } \log(\Lambda) \Leftrightarrow \text{large values of } |(\bar{X} - \bar{Y}) - \Delta_0|,$$

which is the z -test apart from constants that do not depend on the data.

- σ^2 unknown

- The parameter spaces Ω and ω are

$$\Omega = \{(\mu_X, \mu_Y, \sigma^2) \mid \mu_X \in \mathbb{R}, \mu_Y \in \mathbb{R}, \sigma^2 > 0\}$$

$$\omega = \{(\mu_X, \mu_Y, \sigma^2) \mid \mu_X \in \mathbb{R}, \mu_Y = \mu_X - \Delta_0, \sigma^2 > 0\}$$

- Under Ω , the MLE's of (μ_X, μ_Y, σ^2) are

$$\hat{\mu}_{X,\Omega} = \bar{X}, \quad \hat{\mu}_{Y,\Omega} = \bar{Y},$$

$$\hat{\sigma}_\Omega^2 = \frac{1}{m+n} \left[\sum_{i=1}^n (X_i - \hat{\mu}_{X,\Omega})^2 + \sum_{j=1}^m (Y_j - \hat{\mu}_{Y,\Omega})^2 \right] = \frac{(n-1)S_X^2 + (m-1)S_Y^2}{m+n}$$

$$\begin{aligned} \Rightarrow l(\hat{\mu}_{X,\Omega}, \hat{\mu}_{Y,\Omega}, \hat{\sigma}_\Omega^2) &\propto -\frac{m+n}{2} \log(\hat{\sigma}_\Omega^2) - \frac{\sum_{i=1}^n (X_i - \bar{X})^2 + \sum_{j=1}^m (Y_j - \bar{Y})^2}{2\hat{\sigma}_\Omega^2} \\ &= -\frac{m+n}{2} \log(\hat{\sigma}_\Omega^2) - \frac{m+n}{2} \end{aligned}$$

- Under ω , the log-likelihood is proportional to

$$-\frac{m+n}{2} \log(\sigma^2) - \frac{\sum_{i=1}^n (X_i - \mu_X)^2}{2\sigma^2} - \frac{\sum_{j=1}^m [Y_j - (\mu_X - \Delta_0)]^2}{2\sigma^2},$$

and the MLE's of (μ_X, μ_Y, σ^2) are

$$\hat{\mu}_{X,\omega} = \frac{n}{m+n} \bar{X} + \frac{m}{m+n} (\bar{Y} + \Delta_0),$$

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$$\hat{\mu}_{Y,\omega} = \hat{\mu}_{X,\omega} - \Delta_0 = \frac{n}{m+n} (\bar{X} - \Delta_0) + \frac{m}{m+n} \bar{Y},$$

$$\hat{\sigma}_\omega^2 = \frac{1}{m+n} \left[\sum_{i=1}^n (X_i - \hat{\mu}_{X,\omega})^2 + \sum_{j=1}^m (Y_j - \hat{\mu}_{Y,\omega})^2 \right].$$

$$\begin{aligned} \Rightarrow l(\hat{\mu}_{X,\omega}, \hat{\mu}_{Y,\omega}, \hat{\sigma}_\omega^2) &\propto -\frac{m+n}{2} \log(\hat{\sigma}_\omega^2) - \frac{\sum_{i=1}^n (X_i - \hat{\mu}_{X,\omega})^2 + \sum_{j=1}^m (Y_j - \hat{\mu}_{Y,\omega})^2}{2\hat{\sigma}_\omega^2} \\ &= -\frac{m+n}{2} \log(\hat{\sigma}_\omega^2) - \frac{m+n}{2} \end{aligned}$$

- Therefore, the log-likelihood-ratio is

$$\log(\Lambda) = l(\hat{\mu}_{X,\omega}, \hat{\mu}_{Y,\omega}, \hat{\sigma}_\omega^2) - l(\hat{\mu}_{X,\Omega}, \hat{\mu}_{Y,\Omega}, \hat{\sigma}_\Omega^2) = -\frac{m+n}{2} \log \left(\frac{\hat{\sigma}_\omega^2}{\hat{\sigma}_\Omega^2} \right) \text{ and}$$

$$\begin{aligned} \frac{\hat{\sigma}_\omega^2}{\hat{\sigma}_\Omega^2} &= \frac{\sum_{i=1}^n (X_i - \hat{\mu}_{X,\omega})^2 + \sum_{j=1}^m (Y_j - \hat{\mu}_{Y,\omega})^2}{\sum_{i=1}^n (X_i - \bar{X})^2 + \sum_{j=1}^m (Y_j - \bar{Y})^2} \\ &= \frac{\sum_{i=1}^n (X_i - \bar{X})^2 + \sum_{j=1}^m (Y_j - \bar{Y})^2 + \frac{mn}{m+n} (\bar{X} - \bar{Y} - \Delta_0)^2}{\sum_{i=1}^n (X_i - \bar{X})^2 + \sum_{j=1}^m (Y_j - \bar{Y})^2} \\ &= 1 + \frac{mn}{m+n} \times \frac{(\bar{X} - \bar{Y} - \Delta_0)^2}{\sum_{i=1}^n (X_i - \bar{X})^2 + \sum_{j=1}^m (Y_j - \bar{Y})^2} \propto \frac{(\bar{X} - \bar{Y} - \Delta_0)^2}{(m+n-2)s_p^2} \end{aligned}$$

- The likelihood ratio test rejects H_0 for

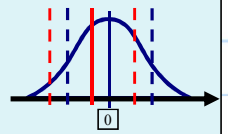
$$\text{small values of } \log(\Lambda) \Leftrightarrow \text{large values of } \frac{|(\bar{X} - \bar{Y}) - \Delta_0|}{s_p},$$

which is the t -test apart from constants that do not depend on the data.

Theorem 7 (power of z -test, 2-sample normal model)

When σ^2 is known, the power function of a level- α z -test for $H_0 : \Delta = \mu_X - \mu_Y = \Delta_0$ vs. $H_A : \Delta = \mu_X - \mu_Y \neq \Delta_0$ is

$$\beta_{\Delta} = 1 - \Phi\left(z(\alpha/2) - \frac{\Delta - \Delta_0}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}}\right) + \Phi\left(-z(\alpha/2) - \frac{\Delta - \Delta_0}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}}\right),$$



where $\Delta \in H_A$, Φ is the cdf of $N(0, 1)$, and $z(\alpha/2)$ is the $(1 - \alpha/2)$ -quantile of $N(0, 1)$.

Proof. power $\beta_{\Delta} = P(\text{rejection region} \mid \mu_X - \mu_Y = \Delta)$

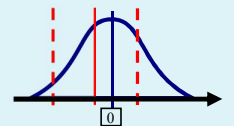
$$\begin{aligned} &= P\left(\left|\frac{(\bar{X} - \bar{Y}) - \Delta_0}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}}\right| > z(\alpha/2) \mid \mu_X - \mu_Y = \Delta\right) \\ &= P\left(\frac{(\bar{X} - \bar{Y}) - \Delta}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}} > z(\alpha/2) - \frac{\Delta - \Delta_0}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}} \mid \mu_X - \mu_Y = \Delta\right) \\ &\quad + P\left(\frac{(\bar{X} - \bar{Y}) - \Delta}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}} < -z(\alpha/2) - \frac{\Delta - \Delta_0}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}} \mid \mu_X - \mu_Y = \Delta\right), \end{aligned}$$

where $[(\bar{X} - \bar{Y}) - \Delta] / \left(\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}\right) \sim N(0, 1)$ when $\mu_X - \mu_Y = \Delta$.

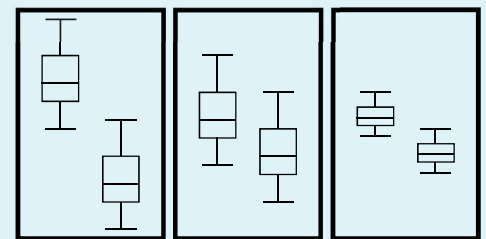
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Note 4 (Some notes about the power function of z - and t -tests)

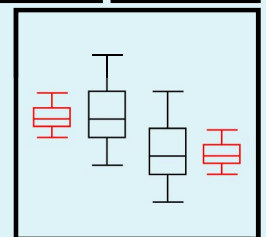
- The power $\beta_{\Delta} \uparrow 1$ as $\alpha \uparrow 1$ (reasonable?) or $\frac{|\Delta - \Delta_0|}{\sigma\sqrt{\frac{1}{n} + \frac{1}{m}}} \uparrow \infty$, i.e.,
 - as $|\Delta - \Delta_0|$ increases (reasonable?), or
 - as σ decreases (reasonable?), or
 - as n, m increase (reasonable?).



- When σ^2 is unknown, the exact power of the t -test can be similarly calculated. But, this calculation requires the use of *noncentral t* distribution.



- Sample size determination** using power.
 - The necessary sample sizes can be determined from α , σ , Δ , and β_{Δ} .
 - For example, when σ^2 is known and $n = m$,



$$\beta_{\Delta} = 1 - \underbrace{\Phi\left(z\left(\frac{\alpha}{2}\right) - \frac{\Delta - \Delta_0}{\sigma}\sqrt{\frac{n}{2}}\right)}_{\text{first term}} + \underbrace{\Phi\left(-z\left(\frac{\alpha}{2}\right) - \frac{\Delta - \Delta_0}{\sigma}\sqrt{\frac{n}{2}}\right)}_{\text{second term}}$$

* Usually, one of these terms is negligible with respect to the other.

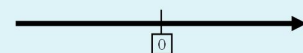
* If $\Delta - \Delta_0 > 0$, the first term will be dominant and

$$\begin{aligned}\beta_\Delta &\approx 1 - \Phi\left(z\left(\frac{\alpha}{2}\right) - \frac{\Delta - \Delta_0}{\sigma}\sqrt{\frac{n}{2}}\right) \\ \Rightarrow z(\beta_\Delta) &\approx z\left(\frac{\alpha}{2}\right) - \frac{\Delta - \Delta_0}{\sigma}\sqrt{\frac{n}{2}} \\ \Rightarrow n &\approx [z(\alpha/2) - z(\beta_\Delta)]^2 \frac{2\sigma^2}{(\Delta - \Delta_0)^2}\end{aligned}$$

Q: What if $\Delta - \Delta_0 < 0$? (**exercise**)

- Determining sample sizes using power is equivalent to using length of C.I. For example, consider the case of σ^2 known.
 - Suppose that m, n are such that the half-length of the C.I. for Δ is $L_{n,m}$.
 - From Thm 4 (LNp.11), $L_{n,m} = z(\alpha/2) \sigma \sqrt{\frac{1}{n} + \frac{1}{m}}$.
 - From Thm 7 (LNp.19), the corresponding power function of $L_{n,m}$ is

$$\beta_\Delta = 1 - \Phi\left(z(\alpha/2) - \frac{\Delta - \Delta_0}{L_{n,m}} z(\alpha/2)\right) + \Phi\left(-z(\alpha/2) - \frac{\Delta - \Delta_0}{L_{n,m}} z(\alpha/2)\right)$$
 - This property could be used to suggest sample sizes m, n under which the statistical standard (LNp.14) is more consistent with the physical standard.



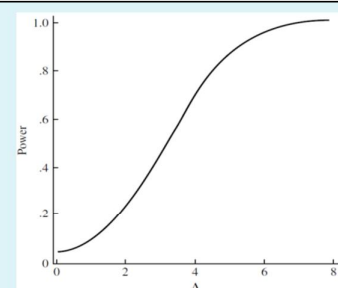
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Example 3 (power function and sample size determination)

- Figure 11.6 (textbook, p.435) gives the power function β_Δ when $n = m = 18$, $\sigma = 5$, $\Delta_0 = 0$, $\alpha = 0.05$ ($\Rightarrow z(\alpha/2) = 1.96$).
- Suppose we want to detect a difference of $\Delta = 1$ with probability (power β_Δ) 0.9. The sample size should be such that

$$0.1 = 1 - \beta_\Delta \approx \Phi\left(1.96 - (\Delta/\sigma)\sqrt{n/2}\right)$$

- Solving for n , we find that the necessary sample size would be 525!
- This is a consequence of a large $\sigma = 5$ relative to the difference $\Delta = 1$.
- If the experimenters want to detect such a difference with a smaller sample size, some modification of the experimental technique to reduce σ would be necessary.



Note 5 (Some notes about z- and t-tests when 2-sample normal model does not hold)

- **Q:** Can we use z- or t-tests (or their corresponding C.I.) when the underlying distributions F, G of X, Y are not normal?

Ans: Yes, if the sample sizes m, n are large. But, **why?** Consider the model:

$$\left. \begin{array}{l} \text{1st sample: } X_1, \dots, X_n \sim \text{i.i.d. from } F \\ \text{2nd sample: } Y_1, \dots, Y_m \sim \text{i.i.d. from } G \end{array} \right\} \Leftarrow \text{independent}$$

where F and G can be *any* continuous distributions with *same finite* variance.

– Denote the means of F, G by μ_X, μ_Y , respectively, and their (identical) variance by $\sigma^2 (< \infty)$. Consider testing $H_0 : \mu_X = \mu_Y$ vs. $H_A : \mu_X \neq \mu_Y$.

– By CLT and LLN, when $m \rightarrow \infty$ and $n \rightarrow \infty$,

$$\bar{X} \stackrel{D}{\approx} N\left(\mu_X, \frac{\sigma^2}{n}\right), \bar{Y} \stackrel{D}{\approx} N\left(\mu_Y, \frac{\sigma^2}{m}\right) \Rightarrow \bar{X} - \bar{Y} \stackrel{D}{\approx} N\left(\mu_X - \mu_Y, \sigma^2 \left(\frac{1}{n} + \frac{1}{m}\right)\right)$$

$$s_X^2 \xrightarrow{P} \sigma^2, s_Y^2 \xrightarrow{P} \sigma^2 \Rightarrow s_p^2 = \frac{n-1}{m+n-2} s_X^2 + \frac{m-1}{m+n-2} s_Y^2 \xrightarrow{P} \sigma^2$$

– Thus,

$$\begin{aligned} & \text{* when } \sigma^2 \text{ is known, } Q_{Z,\Delta} = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sigma \sqrt{\left(\frac{1}{n} + \frac{1}{m}\right)}} \stackrel{D}{\approx} N(0, 1) \\ & \text{* when } \sigma^2 \text{ is unknown,} \end{aligned}$$

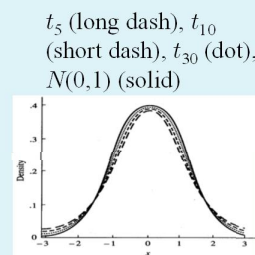
$$Q_{T,\Delta} = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{s_p \sqrt{\left(\frac{1}{n} + \frac{1}{m}\right)}} = \frac{[(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)] / \sqrt{\sigma^2 \left(\frac{1}{n} + \frac{1}{m}\right)}}{\sqrt{s_p^2 / \sigma^2}} \stackrel{D}{\approx} N(0, 1) \quad (\text{by Slutsky's Thm})$$

and t_{m+n-2} tends to $N(0, 1)$ as $m, n \rightarrow \infty$.

- **Q:** How to modify the z - and t -tests (or their corresponding C.I.) when the equal variance assumption in the 2-sample normal model does not hold? Consider the model:

$$\left. \begin{aligned} \text{1st sample: } X_1, \dots, X_n &\sim \text{i.i.d. } N(\mu_X, \sigma_X^2) \\ \text{2nd sample: } Y_1, \dots, Y_m &\sim \text{i.i.d. } N(\mu_Y, \sigma_Y^2) \end{aligned} \right\} \Leftarrow \text{independent}$$

where σ_X^2 and σ_Y^2 can be different. Consider $H_0 : \mu_X = \mu_Y$ vs. $H_A : \mu_X \neq \mu_Y$.



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– This model has 4 parameters: $\mu_X, \mu_Y, \sigma_X^2, \sigma_Y^2$, and their intuitive estimators are

$$\bar{X} \xrightarrow{e} \mu_X, \bar{Y} \xrightarrow{e} \mu_Y, s_X^2 \xrightarrow{e} \sigma_X^2, s_Y^2 \xrightarrow{e} \sigma_Y^2.$$

– Under this model,

$$\text{* } \bar{X} - \bar{Y} \xrightarrow{e} \Delta = \mu_X - \mu_Y \text{ and } \bar{X} - \bar{Y} \sim N\left(\mu_X - \mu_Y, \frac{\sigma_X^2}{n} + \frac{\sigma_Y^2}{m}\right)$$

$$\text{* when } \sigma_X^2, \sigma_Y^2 \text{ are known, } Q_{Z,\Delta}^* = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sqrt{\frac{\sigma_X^2}{n} + \frac{\sigma_Y^2}{m}}} \sim N(0, 1),$$

* when σ_X^2, σ_Y^2 are unknown,

$$\cdot \text{ } Var(\bar{X} - \bar{Y}) = \frac{\sigma_X^2}{n} + \frac{\sigma_Y^2}{m} \text{ can be estimated by } s_{\bar{X}-\bar{Y}}^2 = \frac{s_X^2}{n} + \frac{s_Y^2}{m}$$

• it has been shown (Welch, 1938) the distribution of $\frac{(s_X^2/n) + (s_Y^2/m)}{(\sigma_X^2/n) + (\sigma_Y^2/m)}$ can be approximated by χ_ν^2/ν where

$$\nu = \frac{[(\sigma_X^2/n) + (\sigma_Y^2/m)]^2}{\frac{(\sigma_X^2/n)^2}{n-1} + \frac{(\sigma_Y^2/m)^2}{m-1}},$$

• the degrees of freedom ν can be estimated by $\hat{\nu} = \frac{[(s_X^2/n) + (s_Y^2/m)]^2}{\frac{(s_X^2/n)^2}{n-1} + \frac{(s_Y^2/m)^2}{m-1}}$ and then rounded to the nearest integer,

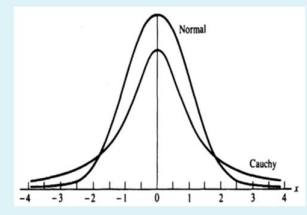
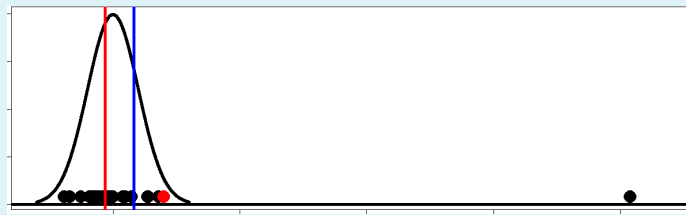
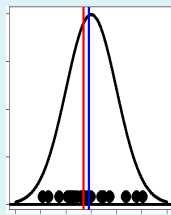
• thus,

$$Q_{T,\Delta}^* = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sqrt{\frac{s_X^2}{n} + \frac{s_Y^2}{m}}} = \frac{[(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)] / \sqrt{\frac{\sigma_X^2}{n} + \frac{\sigma_Y^2}{m}}}{\sqrt{\left(\frac{s_X^2}{n} + \frac{s_Y^2}{m}\right) / \left(\frac{\sigma_X^2}{n} + \frac{\sigma_Y^2}{m}\right)}} \stackrel{D}{\approx} t_\nu$$

and ν can be substituted by its integer estimate.

Note 6 (The circumstances under which z- and t-tests may be invalid)

- The distributions F, G of X, Y are not normal, and sample sizes are small (\Rightarrow the null distribution of the statistic T (or Z) might not be close to t (or $N(0, 1)$) distribution.)
- Data contains *outliers* (extreme values)
 - Problem of averaging data, such as $\bar{X}, \bar{Y}, \hat{\sigma}^2$: sensitive to extreme values

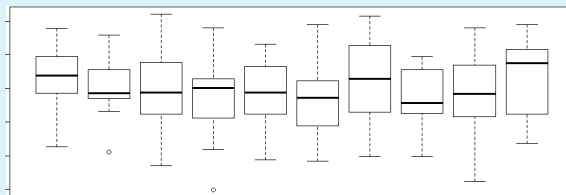


In contrast, median of data is insensitive to extreme values.

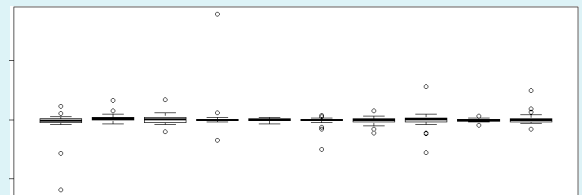
- **Q:** In what distributions, observing extreme values is often expected?

Ans. heavy-tail distributions, such as Cauchy $C(\mu, \sigma)$.

$X_1, \dots, X_{20} \sim \text{i.i.d. } N(0, 1)$ (repeat 10 times)



$X_1, \dots, X_{20} \sim \text{i.i.d. } C(0, 1)$ (repeat 10 times)



- Some properties of Cauchy distribution
 - * it does not have finite moments of any order (a consequence of heavy tail)

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* if X_1, \dots, X_n are i.i.d. $\sim C(\mu, \sigma)$, then $\bar{X} \sim C(\mu, \sigma)$.

($\xleftrightarrow{\text{cf.}}$ if X_1, \dots, X_n are i.i.d. from a distribution with *finite* variance σ^2 , then $\text{Var}(\bar{X}) = \sigma^2/n \rightarrow 0$ when $n \rightarrow \infty$.)

\Rightarrow For 2-sample data, when F, G are Cauchy, even though the sample sizes m, n are large, the property " $\bar{X}, \bar{Y} \stackrel{D}{\approx}$ normal" does not hold.

(**Q:** Why do LLN and CLT not apply to Cauchy?)

Question 4.

How to develop statistical methods for the circumstances under which z- and t-tests are inappropriate?

Q: What limits the validity of the tests?

Ans. Statistical models (i.e., joint distributions of data) covered in the 2-sample normal model (a *model space* of dimension *three*) are still not flexible enough to reflect the pattern of data

- \Rightarrow should include more joint distributions into the model space, say enlarge the model space to allow for F, G being *any* distributions
- \Rightarrow develop statistical methods under this enlarged model space
- \Rightarrow such statistical methods should be suitable for data of any patterns

❖ **Reading:** textbook, 11.1, 11.2.1, 11.2.2

• A nonparametric method for 2-sample problem --- Mann-Whitney Test

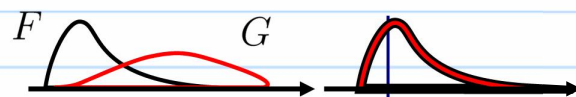
Question 5.

- In the materials taught before, we usually assume, in the statistical modeling, that the data follows a *particular* joint distribution which contains some unknown parameters of *finite* dimension.
- The statistical inferences, estimation and testing, are then based on a formulation of these parameters.

Q: What if we do not have any knowledge about the *particular* form of the joint distribution of data?

Consider the problem of 2-sample comparison.

- Let Ω be the collection of *all* continuous distributions
- Only assume that $F, G \in \Omega$
- Thus, the statistical model is:



$$\left. \begin{array}{l} \text{1st sample: } X_1, \dots, X_n \sim \text{i.i.d. from } F \\ \text{2nd sample: } Y_1, \dots, Y_m \sim \text{i.i.d. from } G \end{array} \right\} \Leftarrow \text{independent} \quad (\square)$$

- This model contains parameters of *infinitely many* dimension because

$$\dim(\Omega) = \infty \quad (\text{why?})$$

- Under this model, a 2-sample comparison examines the null and alternative hypotheses:

$$H_0 : F = G \quad \text{vs.} \quad H_A : F \neq G.$$

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Definition 2 (nonparametric models and nonparametric methods)

- Nonparametric models do not assume any *particular* distributional form. Nonparametric models can be viewed as having *infinitely many* parameters. ($\xleftrightarrow{\text{cf.}}$ parametric models: parameters are of *finite* dimension)
- Statistical methods developed under nonparametric models are called *nonparametric methods*.

Review 3 (order statistics and ranks)

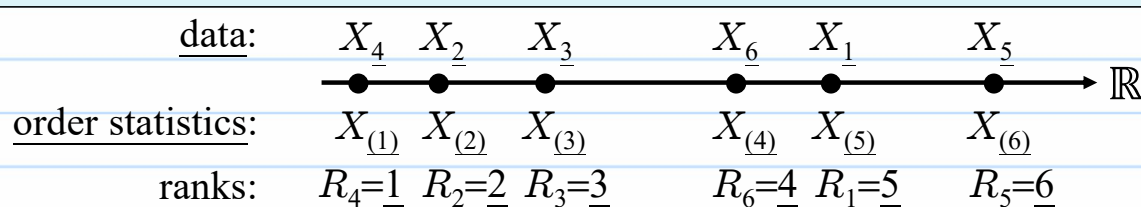
- Let X_1, X_2, \dots, X_n be random variables. We sort the X_i 's and denote by $X_{(1)} \leq X_{(2)} \leq \dots \leq X_{(n)}$ the *order statistics*. Using the notation,

$$X_{(1)} = \min(X_1, X_2, \dots, X_n) \quad \text{is the } \textit{minimum},$$

$$X_{(n)} = \max(X_1, X_2, \dots, X_n) \quad \text{is the } \textit{maximum}.$$

- Let $R(X_1, X_2, \dots, X_n) = (R_1, R_2, \dots, R_n)$ such that $X_i = X_{(R_i)}$, $i = 1, \dots, n$. Then, (R_1, R_2, \dots, R_n) is called the *ranks* of X_1, X_2, \dots, X_n . Notice that

$$R_i = \sum_{j=1}^n \delta(X_i - X_j), \quad \text{where } \delta(t) = \begin{cases} 1, & \text{if } t \geq 0, \\ 0, & \text{if } t < 0. \end{cases}$$



Theorem 8 (sufficient and complete statistics for nonparametric models)

Let X_1, \dots, X_n be i.i.d. from F , where $F \in \Omega$.

Then, $(X_{(1)}, X_{(2)}, \dots, X_{(n)})$ is sufficient and complete.

($\xleftrightarrow{\text{cf.}}$ $X_1, \dots, X_n \sim \text{i.i.d. } N(\mu, \sigma^2) \Rightarrow (\bar{X}, s_X^2)$ is sufficient and complete.)

Proof. Denote the pdf of F by f . The joint pdf of $X_{(1)}, X_{(2)}, \dots, X_{(n)}$ is

$$f_{X_{(1)}, X_{(2)}, \dots, X_{(n)}}(x_1, x_2, \dots, x_n) = n! \times f(x_1)f(x_2) \cdots f(x_n),$$

for $x_1 < x_2 < \cdots < x_n$ and zero, otherwise.

The proof of sufficiency follows from the fact that the conditional probability of X_1, \dots, X_n given $X_{(1)}, \dots, X_{(n)}$ is $\frac{1}{n!}$, which is irrelevant to F .

The proof of completeness is omitted (out of the scope of this course).

Note 7 (Some notes about order statistics and ranks)

- Order statistics and ranks are defined precisely, i.e., **no ties**, under the condition $P(X_i = X_j) = 0, i \neq j$ (**Note.** this condition holds when $X_1, \dots, X_n \sim \text{i.i.d. from } F$ and F is a continuous distribution).
- Under Ω , the dimension of data (i.e., n) cannot be reduced without losing the information about $F (\in \Omega)$.
- Under 1-sample model, ranks + order statistics = complete data
- Order statistics are intuitive estimator of quantiles, e.g., median.

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- Ranks are invariant under any monotonic transformation of data, i.e.,

$$R(X_1, \dots, X_n) = R(H(X_1), \dots, H(X_n)),$$

if H is a monotone increasing function and

$$R(X_1, \dots, X_n) = (n+1) - R(H(X_1), \dots, H(X_n)),$$

if H is a monotone decreasing function. ($\xleftrightarrow{\text{cf.}}$ z - or t -tests may change significantly under monotonic transformations of data).

- Replacing the data by their ranks also has the effect of moderating the influence of outliers.
- Many nonparametric methods are based on order statistics and/or ranks.
- Q:** Why are many nonparametric methods based on replacement of the data by ranks? What information of data are contained in their ranks?

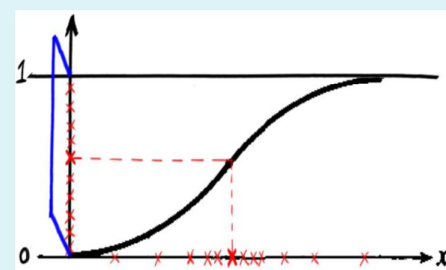
(exercise) – **Recall.** Let X_1, \dots, X_n be i.i.d. from a continuous cdf F , and let $U_i = F(X_i)$, $i = 1, \dots, n$. Then, U_1, \dots, U_n are i.i.d. from $U(0, 1)$.

(exercise) – **Recall.** If $U_1, \dots, U_n \sim \text{i.i.d. } U(0, 1)$, the pdf of the i th-order statistic $U_{(i)}$ is

$$f_{U_{(i)}}(u) = \frac{n!}{(i-1)!(n-i)!} u^{i-1} (1-u)^{n-i},$$

for $0 < u < 1$ and zero, otherwise.

Note that $E(U_{(i)}) = i/(n+1)$.



– $U_i = F(X_i)$ is not a statistic because F is an unknown function.

– But,

$$X_i = X_{(R_i)} \rightarrow U_{(R_i)} = F(X_{(R_i)}) \rightarrow R_i = (n+1) \frac{R_i}{n+1} \leftrightarrow (n+1)E[U_{(R_i)} | R_i].$$

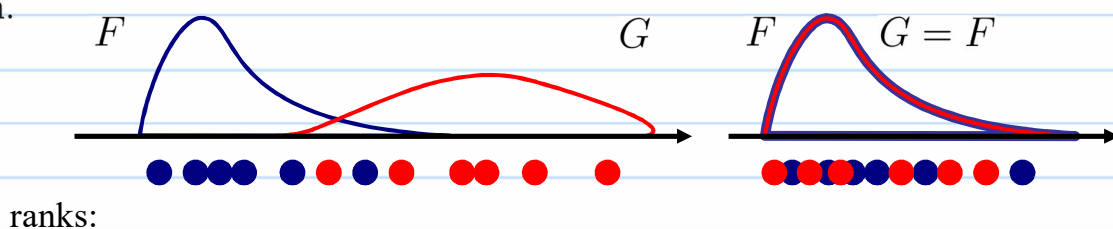
Question 6.

How to use ranks to compare two samples? Under the nonparametric model (\square) in LNp.27, for the null and alternative hypotheses:

$$H_0 : F = G \quad \text{vs.} \quad H_A : F \neq G$$

what data are “more extreme,” i.e., cast more doubts on H_0 ?

Intuition.



Theorem 9 (Mann-Whitney test or Wilcoxon rank sum test)

Consider the nonparametric model (\square) in LNp.27.

- Pool all $m+n$ observations (i.e., $X_1, \dots, X_n, Y_1, \dots, Y_m$) together and rank them in order of increasing size, i.e.,

$$R(X_1, \dots, X_n, Y_1, \dots, Y_m) = (R_1, \dots, R_n, R_{n+1}, \dots, R_{m+n}).$$

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- Test statistic W_X (or W_Y)

– Let $W_X = \sum_{i=1}^n R_i$ and $W_Y = \sum_{j=1}^m R_{n+j}$. They are respectively the sums of the ranks of X_i 's and Y_j 's in the pooled data. Notice that

$$W_X + W_Y = 1 + 2 + \dots + (m+n) = \frac{(m+n)(m+n+1)}{2}$$

$$\Rightarrow W_Y = \frac{(m+n)(m+n+1)}{2} - W_X.$$

– Data with larger or smaller W_X are more extreme \Rightarrow tend to reject H_0

- Null distribution of W_X

– Under H_0 ($F = G$),

$$\begin{array}{ccccccc} X_1, & \dots, & X_n, & Y_1, & \dots, & Y_m & \sim \text{i.i.d. } F \\ \downarrow & & \downarrow & \downarrow & & \downarrow & \\ R_1, & \dots, & R_n, & R_{n+1} & \dots, & R_{m+n} & \sim ? \end{array}$$

– Any assignments of the ranks $\{1, \dots, m+n\}$ to the pooled $m+n$ data are equally likely, and the total number of different assignments is $(m+n)!$.

– Joint distribution of R_1, \dots, R_n :

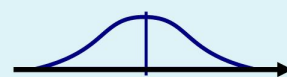
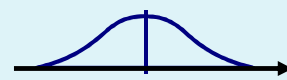
* Consider an urn containing $m+n$ balls, labelled by $1, 2, \dots, m+n$, respectively.

* Sequentially draw n balls without replacement from the urn \Rightarrow there are $\binom{m+n}{n} \times n!$ different outcomes, each with equal probability



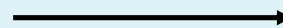
- * Let r_1, r_2, \dots, r_n be the numbers on the 1st, 2nd, \dots , n th balls drawn, respectively. Then,

$$P(R_1 = r_1, \dots, R_n = r_n) = \frac{1}{\binom{m+n}{n} \times n!} = \frac{m!}{(m+n)!}.$$
- The null distribution of $W_X = R_1 + \dots + R_n$ (W_X is the sum of the numbers on the n balls) can be obtained from the joint distribution of R_1, \dots, R_n .
- Rejection region
 - Let $n_1 = \min(n, m)$ be the smaller sample size, and W be the rank sum from that sample (i.e., $W = W_X$ if $n \leq m$ and $W = W_Y$ if $n > m$).
 - * Note that under H_0 ,
 - $E(W) = \begin{cases} E(R_1) + \dots + E(R_n), & \text{if } n \leq m \\ E(R_{n+1}) + \dots + E(R_{n+m}), & \text{if } n > m \end{cases} = \frac{n_1(m+n+1)}{2}.$
 - the null distribution of W is symmetric around $E(W)$ (exercise).
 - * Let $W' = n_1(m+n+1) - W$.
 - * Let $W^* = \min(W, W')$.
 - Reject H_0 when W^* is small, i.e., $W^* \leq w$.
 - Table 8 of Appendix B in the textbook gives critical values w for W^* .



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- We have assumed here that there are no ties among the observations. If there are only a small number of ties; tied observations are assigned average ranks.



Example 4 (Mann-Whitney test, heat of fusion of ice, cont. Ex.1 in LNp.3)

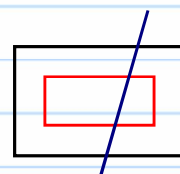
- The ranks are (ties \Rightarrow average rank)

| | | | | | | | | |
|----------|------|------|------|------|------|------|------|-----|
| Method A | 7.5 | 19.0 | 11.5 | 19.0 | 15.5 | 15.5 | 19.0 | 4.5 |
| | 21.0 | 15.5 | 11.5 | 9.0 | 11.5 | | | |
| Method B | 11.5 | 1.0 | 7.5 | 4.5 | 4.5 | 15.5 | 2.0 | 4.5 |

- $n_1 = 8$, $W = W_B = 51$, $W' = 8(8+13+1) - W = 125$, $W^* = \min(W, W') = 51$
- two-sided test at level $\alpha = 0.01$, critical value = 53
- two-sided test at level $\alpha = 0.05$, critical value = 60
- Therefore, the Mann-Whitney test rejects the null hypothesis at $\alpha = 0.01$.

a comparison of parametric and nonparametric models

| | model | data | | power on | power on |
|----------------------|-------|-----------|------------|----------|----------------------------|
| | space | reduction | robustness | H_A^p | $H_A^{np} \setminus H_A^p$ |
| parametric models | small | low-dim | worse | higher | (usually) lower |
| nonparametric models | large | high-dim | better | lower | (usually) higher |

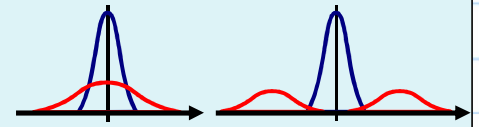


Question 7.

Does Mann-Whitney test have reasonably good powers over the whole $H_A : F \neq G$? Note that

$$H_0 \cup H_A = \{(F, G) \mid F, G \in \Omega\},$$

$$H_0 = \{(F, G) \mid F \in \Omega, G = F\}.$$



- Assume that the distributions (cdfs)

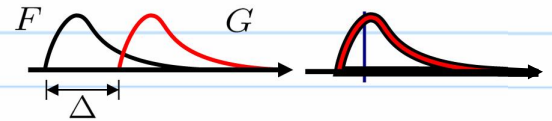
$F, G \in \Omega$ and F, G have same shape.

- If $X \sim F$ and $Y = X + \Delta$, where Δ is an unknown constant, then for the cdf $G(y)$ of Y , we have

$$G(y) \equiv P(Y \leq y) = P(X + \Delta \leq y) = P(X \leq y - \Delta) = F(y - \Delta),$$

and for the pdfs $f(x)$ of X and $g(y)$ of Y , we have

$$g(y) = \frac{d}{dy} G(y) = \frac{d}{dy} F(y - \Delta) = f(y - \Delta).$$



- Thus, the statistical model is:

$$\left. \begin{array}{l} \text{1st sample: } X_1, \dots, X_n \sim \text{i.i.d. from } F \\ \text{2nd sample: } Y_1, \dots, Y_m \sim \text{i.i.d. from } G \end{array} \right\} \Leftarrow \text{independent} \quad (\diamond)$$

where $F \in \Omega$ and $G(x) = F(x - \Delta)$.

- This model contains *infinitely many* parameters because $\dim(\Omega) = \infty$.

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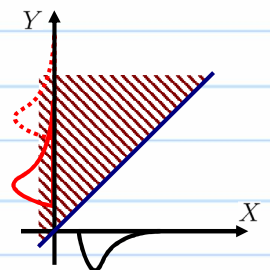
- Under this model, the null $H_0 : F = G$ become $H_0 : \Delta = 0$, and the alternative $H_A : F \neq G$ becomes $H_A : \Delta \neq 0$, i.e.,

$$* H_0 \cup H_A = \{(F, G) \mid F \in \Omega, G(y) = F(y - \Delta), \Delta \in \mathbb{R} \text{ (or } \pi_\Delta \in [0, 1])\}$$

$$* H_0 = \{(F, G) \mid F \in \Omega, G(y) = F(y - \Delta), \Delta = 0 \text{ (or } \pi_\Delta = 1/2)\}$$

Theorem 10 (An alternative formulation of H_0 and H_A)

- Suppose that (1) $X \sim F \in \Omega$, (2) $Y \sim G$, where $G(x) = F(x - \Delta)$, and (3) X, Y are independent. The joint pdf of (X, Y) is $f(x)g(y) = f(x)f(y - \Delta)$.
- Define $\pi_\Delta = P_\Delta(X < Y)$. Clearly, $0 \leq \pi_\Delta \leq 1$.
- Then, $\pi_\Delta = 1/2$ if and only if $\Delta = 0$.

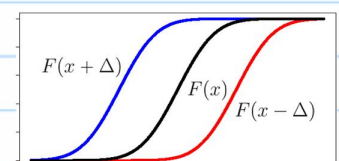


Proof.

$$P_\Delta(X < Y) = \int_{-\infty}^{\infty} \int_x^{\infty} f(x) f(y - \Delta) dy dx$$

$$= \int_{-\infty}^{\infty} f(x) [F(y - \Delta)]_x^{\infty} dx$$

$$= \int_{-\infty}^{\infty} [1 - F(x - \Delta)] f(x) dx = 1 - \int_{-\infty}^{\infty} F(x - \Delta) f(x) dx$$



- If $\Delta > 0$, then $F(x - \Delta) \leq F(x) \leq F(x + \Delta)$, $\forall x$, and there must exist a region A of x in which the inequalities are strict and $\int_A f(x) dx > 0$.

- Thus, for $\Delta > 0$,

$$\underbrace{\int_{-\infty}^{\infty} F(x - \Delta) f(x) dx}_{1 - P_{\Delta}(X < Y)} < \underbrace{\int_{-\infty}^{\infty} F(x) f(x) dx}_{1 - P_{\Delta=0}(X < Y)} < \underbrace{\int_{-\infty}^{\infty} F(x + \Delta) f(x) dx}_{1 - P_{-\Delta}(X < Y)}$$

- Then, the results follow from: $\int_{-\infty}^{\infty} F(x) f(x) dx = \int_0^1 z dz = \frac{1}{2} z^2 \Big|_0^1 = \frac{1}{2}.$

Theorem 11 (An alternative view of Mann-Whitney test)

Consider the nonparametric model (\diamond) in LNp.35.

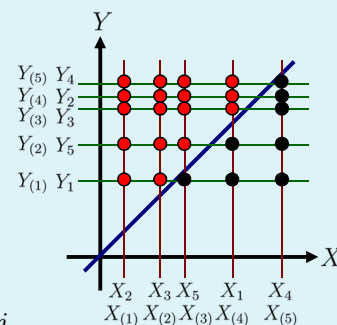
- Estimation of π_{Δ} : the parameter $\pi_{\Delta} = P_{\Delta}(X < Y)$ can be estimated by the proportion of the comparisons for which X was less than Y , i.e.,

– consider any pairs (X_i, Y_j) , $1 \leq i \leq n$, $1 \leq j \leq m$,

– let $Z_{ij} = \begin{cases} 1, & \text{if } X_i < Y_j, \\ 0, & \text{otherwise,} \end{cases} \Rightarrow \hat{\pi}_{\Delta} = \frac{1}{mn} \sum_{i=1}^n \sum_{j=1}^m Z_{ij}$

– an alternative expression: consider the mn pairs $(X_{(i)}, Y_{(j)})$, and let

$$V_{ij} = \begin{cases} 1, & \text{if } X_{(i)} < Y_{(j)}, \\ 0, & \text{otherwise,} \end{cases} \Rightarrow \hat{\pi}_{\Delta} = \frac{1}{mn} \sum_{i=1}^n \sum_{j=1}^m V_{ij}$$



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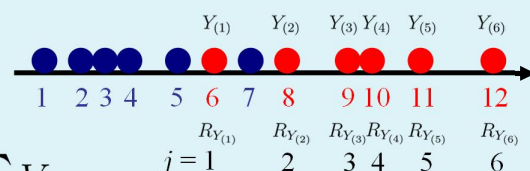
- Test $H_0 : \pi_{\Delta} = 1/2 (\Leftrightarrow \Delta = 0)$ vs. $H_A : \pi_{\Delta} \neq 1/2 (\Leftrightarrow \Delta \neq 0)$

– intuitively, should reject H_0 if $\hat{\pi}_{\Delta}$ is too small (closer to 0) or too large (closer to 1)

– test statistic U_Y (or U_X)

* Define

$$U_Y \equiv (mn) \hat{\pi}_{\Delta} = \sum_{i=1}^n \sum_{j=1}^m Z_{ij} = \sum_{i=1}^n \sum_{j=1}^m V_{ij}.$$



• Reject H_0 if U_Y is too small or too large (closer to 0 or mn).

* Let $R_{Y_{(j)}}$ be the rank of $Y_{(j)}$ in the *pooled* sample. Then,

$$\sum_{j=1}^m R_{Y_{(j)}} = \text{rank sum of } Y_j\text{'s (or } Y_{(j)}\text{'s)} = R_{n+1} + \cdots + R_{n+m} = W_Y.$$

* Notice that

$$U_Y = \sum_{j=1}^m \left(\sum_{i=1}^n V_{ij} \right) = \sum_{j=1}^m \underbrace{(R_{Y_{(j)}} - j)}_{\#\{X_{(i)} < Y_{(j)}\}} = \left(\sum_{j=1}^m R_{Y_{(j)}} \right) - \frac{m(m+1)}{2} = W_Y - [m(m+1)]/2.$$

* Similarly, U_X can be defined by changing " $X_{(i)} < Y_{(j)}$ " in V_{ij} to " $X_{(i)} > Y_{(j)}$ ", and

$$\cdot \frac{1}{mn} U_X \xrightarrow{e} 1 - \pi_{\Delta} = P_{\Delta}(X > Y)$$

- $U_X = mn - U_Y$
- $U_X = W_X - \frac{1}{2}n(n+1)$
- reject H_0 if U_X is too small or too large
- null distribution of U_Y : the pmf of U_Y under H_0 can be obtained from the null distribution of W_Y by

$$P(U_Y = u) = P\left(W_Y - \frac{m(m+1)}{2} = u\right) = P\left(W_Y = u + \frac{m(m+1)}{2}\right).$$
- The tests based on U_Y and W_Y (or U_X and W_X) are actually equivalent.

Note 8 (A comparison of t-test and Mann-Whitney (M-W) test)

- Unlike t -test, the M-W test does not depend on normality assumption.
- The M-W test is insensitive to outliers, where as the t -test is sensitive.
- When the normality assumption holds, the t -test is more powerful.
- However, under normality assumption, the M-W test is nearly as powerful as the t -test. It has been shown that to attain the same power
 - the total sample size required for the t -test is approximately 0.95 times the total sample size required for the M-W test.
- The M-W test is generally preferable, especially for small sample sizes.

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Theorem 12 (means and variances of U_Y and W_Y under H_0)

Consider the nonparametric model (\diamond) in LNp.35. If $\Delta = 0$ ($\Leftrightarrow \pi_\Delta = 1/2$),

- $E(W_Y) = [m(m+n+1)]/2$ and $Var(W_Y) = [mn(m+n+1)]/12$
 ($\Leftrightarrow E(W_X) = [n(m+n+1)]/2$ and $Var(W_X) = [mn(m+n+1)]/12$
 since $W_X = [(m+n)(m+n+1)]/2 - W_Y$)
- $E(U_Y) = mn/2$ and $Var(U_Y) = [mn(m+n+1)]/12$
 ($\Leftrightarrow E(U_X) = mn/2$ and $Var(U_X) = [mn(m+n+1)]/12$
 since $U_X = mn - U_Y$)

Proof. It is enough to prove the case of W_Y .

- Note that $W_Y = R_{n+1} + \cdots + R_{m+n}$.

Under $H_0 : \Delta = 0$, $(R_{n+1}, \dots, R_{m+n})$ can be viewed as a without-replacement simple random sample from the population

$$\{1, \dots, n, n+1, \dots, m+n\}.$$

- Let $N = m+n$. Since

$$\sum_{k=1}^N k = \frac{N(N+1)}{2} \quad \text{and} \quad \sum_{k=1}^N k^2 = \frac{N(N+1)(2N+1)}{12},$$

the population mean μ and variance σ^2 of this population distribution are

$$\mu = \frac{1}{N} \left(\sum_{k=1}^N k \right) = \frac{N+1}{2} \quad \text{and} \quad \sigma^2 = \frac{1}{N} \left(\sum_{k=1}^N k^2 \right) - \mu^2 = \frac{N^2 - 1}{12}.$$

- Let $\bar{R} (= W_Y/m)$ be the average of this without-replacement sample $(R_{n+1}, \dots, R_{m+n})$. Then (by Thms 1 & 3 in LN, Ch7, p.16-18),

$$E(\bar{R}) = \mu \quad \text{and} \quad Var(\bar{R}) = (\sigma^2/m)[(N-m)/(N-1)].$$
- The results follows from $E(W_Y) = m E(\bar{R})$ and $Var(W_Y) = m^2 Var(\bar{R})$.

Theorem 13 (Asymptotic null distribution of U_Y)

Consider the nonparametric model (\diamond) in LNp.35 and the null $H_0 : \Delta = 0$ ($\Leftrightarrow \pi_\Delta = 1/2$). For m, n both greater than 10, the null distribution of U_Y (or U_X) is well approximated by a normal distribution, i.e.,

$$\frac{U_Y - E(U_Y)}{\sqrt{Var(U_Y)}} \stackrel{D}{\approx} N(0, 1) \quad \left(\text{or} \quad \frac{U_X - E(U_X)}{\sqrt{Var(U_X)}} \stackrel{D}{\approx} N(0, 1) \right).$$

The proof is omitted, but some **notes** are given below.

- This Thm does not follow immediately from the ordinary CLT although

$$U_Y = \sum_i \sum_j Z_{ij} \quad \text{and} \quad Z_{ij} \sim \text{binomial}(1, \pi_\Delta).$$

But, Z_{ij} 's are not independent.

- Similarly, the null distribution of W_Y (or W_X) can be approximated by normal, i.e.,

$$\frac{W_Y - E(W_Y)}{\sqrt{Var(W_Y)}} \stackrel{D}{\approx} N(0, 1) \quad \left(\text{or} \quad \frac{W_X - E(W_X)}{\sqrt{Var(W_X)}} \stackrel{D}{\approx} N(0, 1) \right).$$

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Example 5 (Asymptotic null dist. of W_Y , heat of fusion of ice, cont. Ex.4 in LNp.34)

- $n = 13$ (method A), $m = 8$ (method B), $W_B = 51$.
- Under the null, $\mu_{W_B} = E(W_B) = [8(8 + 13 + 1)]/2 = 88$,
 $\sigma_{W_B} = \sqrt{Var(W_B)} = \sqrt{[(8 \times 13)(8 + 13 + 1)]/12} = 13.8$.
- Because
$$\frac{W_B - \mu_{W_B}}{\sigma_{W_B}} = \frac{51 - 88}{13.8} = -2.68,$$

the approximate p -value is $P(|N(0, 1)| > 2.68) = 2 \times [1 - \Phi(2.68)] = 0.0074$
 $(\Rightarrow \text{reject } H_0 \text{ at } \alpha = 0.01 \Rightarrow \text{consistent with the testing result using exact null distribution in Ex.4})$

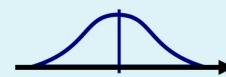
Theorem 14 (Nonparametric confidence interval for Δ)

Consider the nonparametric model (\diamond) in LNp.35.

- Q:** How can we test $H_0^* : \Delta = \Delta_0$ vs. $H_A^* : \Delta \neq \Delta_0$,
 where Δ_0 is a known constant?
 - Under H_0^* , we have (1) $X_i \sim F$, (2) $Y_j \sim G$, and (3) $G(x) = F(x - \Delta_0)$.
 Then,

$$X_1, \dots, X_n, Y_1 - \Delta_0, \dots, Y_m - \Delta_0 \sim \text{i.i.d. } F$$
 - The test of H_0^* vs. H_A^* using the data X_i 's and Y_j 's is equivalent to testing $H_0 : \Delta = 0$ vs. $H_A : \Delta \neq 0$ using the data X_i 's and $(Y_j - \Delta_0)$'s.
 - To test $H_0^* : \Delta = \Delta_0$, can use

- * the test statistic: $U_Y(\Delta_0) = \#\{X_i < Y_j - \Delta_0\} = \#\{Y_j - X_i > \Delta_0\}$,
- * the acceptance region: $k(\alpha) \leq U_Y(\Delta_0) \leq mn - k(\alpha)$,
where $k(\alpha)$ is the critical value determined by the significance level α
(**Note.** The null distribution of $U_Y(\Delta_0)$ is symmetric about $mn/2$.)

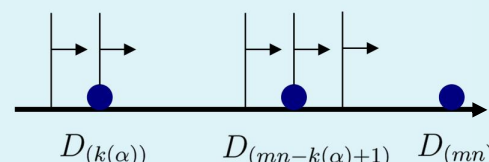


- By the duality of test and C.I., a $100(1 - \alpha)\%$ confidence interval for Δ is

$$C = \{ \Delta \mid k(\alpha) \leq U_Y(\Delta) \leq mn - k(\alpha) \}.$$

- Let $D_{(1)}, D_{(2)}, \dots, D_{(mn)}$ denote the ordered mn differences $(Y_j - X_i)$'s.

Then, $C = [D_{(k(\alpha))}, D_{(mn-k(\alpha)+1)}]$.



- To see this,

- * if $\Delta_0 = D_{(k(\alpha))}$, then $U_Y(\Delta_0) = \#\{Y_j - X_i > \Delta_0\} = mn - k(\alpha)$,
if $\Delta_0 < D_{(k(\alpha))}$, then $U_Y(\Delta_0) = \#\{Y_j - X_i > \Delta_0\} \geq mn - k(\alpha) + 1$,
thus, $D_{(k(\alpha))}$ is the leftmost point of the confidence interval C ,
- * if $\Delta_0 < D_{(mn-k(\alpha)+1)}$, then $U_Y(\Delta_0) = \#\{Y_j - X_i > \Delta_0\} \geq k(\alpha)$,
if $\Delta_0 \geq D_{(mn-k(\alpha)+1)}$, then $U_Y(\Delta_0) = \#\{Y_j - X_i > \Delta_0\} \leq k(\alpha) - 1$,
thus, $D_{(mn-k(\alpha)+1)}$ is the rightmost point of the confidence interval C .

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Example 6 (C.I. for Δ , heat of fusion of ice, cont. Ex.4 in LNp.34 & Ex.5 in LNp.42)

- $n = 13$ (method A), $m = 8$ (method B), $W_B = 51$. Under null, $E(W_B) = 88$.
- Under the significant level $\alpha = 0.05$, the critical value for W_B^* is 60 (Ex.4, LNp.34) \Rightarrow acceptance region: $61 \leq W_B \leq 88 + (88 - 61) = 115$
 $\Leftrightarrow 25 \leq U_B = W_B - [8(8 + 1)]/2 = W_B - 36 \leq 79$.
- After sorting the $mn = 8 \times 13 = 104$ differences $(Y_j - X_i)$'s, we get
 $D_{(k(\alpha)=25)} = -0.07$ and $D_{(mn-k(\alpha)+1=80)} = -0.01$.
A 95% confidence interval for Δ is $(-0.07, -0.01)$, which does not contain 0.
[$\xleftrightarrow{\text{cf.}}$ the C.I. $(0.015, 0.065)$ given in Ex.2 (LNp.12)
– Note that the Δ here is the $-\Delta$ in Ex.2.
– In this case, the C.I. based on the nonparametric model is slightly wider than the one based on the normal model.
– But, the latter C.I. relies on the validity of normality assumption.]

Theorem 15 (Bootstrap confidence interval for π_Δ ($\leftrightarrow \Delta$))

Consider the nonparametric model (\diamond) in LNp.35 or the nonparametric model (\square) in LNp.27. (**Note.** (1) (\square) has more models than (\diamond) (2) $\pi_\Delta = P(X < Y)$ is well-defined in (\diamond) and (\square) (3) Δ is well-defined only in (\diamond))

- Bootstrapping is a numerical method that can be used to gain information about the sampling distribution of $\hat{\pi}_\Delta = \frac{1}{mn}(\#\{X_i < Y_j\}) \xrightarrow{e} \pi_\Delta$, and the estimated standard error of $\hat{\pi}_\Delta$.

- In bootstrap, we

$$\left. \begin{array}{l} X_1, \dots, X_n \sim \text{i.i.d. from } F \\ Y_1, \dots, Y_m \sim \text{i.i.d. from } G \end{array} \right\} \Leftarrow \text{independent}$$

- replace the true cdf F (unknown) by the empirical cdf \hat{F}_n (known) of $(X_1, \dots, X_n) = (x_1, \dots, x_n)$ [\hat{F}_n : assigns x_i 's equal probabilities $1/n$]
- replace the true cdf G (unknown) by the empirical cdf \hat{G}_m (known) of $(Y_1, \dots, Y_m) = (y_1, \dots, y_m)$ [\hat{G}_m : assigns y_j 's equal probabilities $1/m$]
- Re-sample (generate data $X'_1, \dots, X'_n, Y'_1, \dots, Y'_m$ using simulation) from this model:

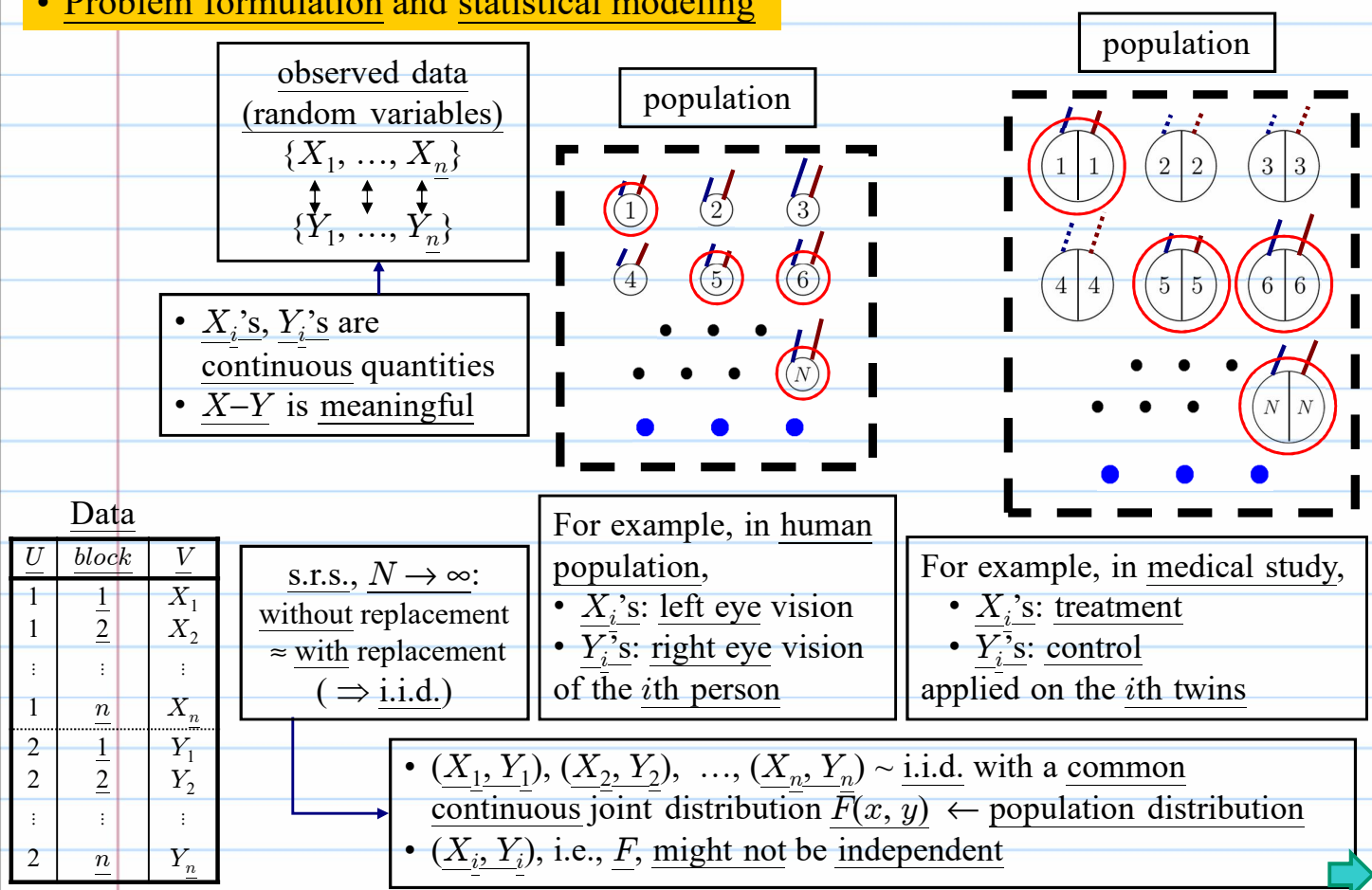
$$\left. \begin{array}{l} X'_1, \dots, X'_n \sim \text{i.i.d. from } \hat{F}_n \\ Y'_1, \dots, Y'_m \sim \text{i.i.d. from } \hat{G}_m \end{array} \right\} \Leftarrow \text{independent}$$
 - X'_1, \dots, X'_n is a with-replacement sample from the population $\{x_1, \dots, x_n\}$,
 - Y'_1, \dots, Y'_m is a with-replacement sample from the population $\{y_1, \dots, y_m\}$.
- Repeat the re-sampling procedure many times, say B times, and
 - at each time, compute $\hat{\pi}'_{\Delta} = \frac{1}{mn} \#\{X'_i < Y'_j\}$ from $(X'_1, \dots, X'_n, Y'_1, \dots, Y'_m)$
 - this produces a bootstrap sample: $(\hat{\pi}'_{\Delta,1}, \dots, \hat{\pi}'_{\Delta,B})$
- A histogram of $(\hat{\pi}'_{\Delta,1}, \dots, \hat{\pi}'_{\Delta,B})$ offers an indication of the sampling distribution of $\hat{\pi}_{\Delta}$ (\Rightarrow a $100(1 - \alpha)\%$ C.I. of π_{Δ} is $[\hat{\pi}'_{\Delta,(B(\alpha/2))}, \hat{\pi}'_{\Delta,(B(1-\alpha/2))}]$),
- the standard deviation of $(\hat{\pi}'_{\Delta,1}, \dots, \hat{\pi}'_{\Delta,B}) \xrightarrow{e}$ the standard error of $\hat{\pi}_{\Delta}$.

❖ Reading: textbook, 11.2.3

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• Comparing paired samples \longleftrightarrow Independent samples

• Problem formulation and statistical modeling

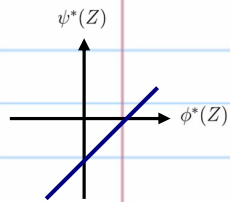


- Let random variables Z_1, \dots, Z_n represent the variability of the n members sampled from the population.
- Assume Z_1, \dots, Z_n are i.i.d. from a population distribution $H(z)$.
- Let $X = \phi(Z)$ and $Y = \psi(Z)$, where ϕ, ψ contain random components, and denote
 - $F(x, y)$: the joint distributions of (X, Y) ,
 - μ_X and μ_Y : the means of X and Y , respectively,
 - $\Delta = \mu_X - \mu_Y$.

$$(\phi, \psi): \begin{matrix} Z & \rightarrow & (X, Y) \\ (H(z)) & & (F(x, y)) \end{matrix}$$

$$\begin{aligned} &\text{Because } Z_1, \dots, Z_n \sim \text{i.i.d. } H(z), \\ &(X_1, Y_1), \dots, (X_n, Y_n) \\ &\sim \text{i.i.d. } F(x, y) \end{aligned}$$

- Then, for $1 \leq i \leq n$, $\begin{cases} X_i = \phi(Z_i) \\ Y_i = \psi(Z_i) \end{cases}$



$$\begin{cases} X_i = \phi(Z_i) = \mu_X + \epsilon_{1i}, \\ Y_j = \psi(Z_{n+j}) = \mu_Y + \epsilon_{2j}, \end{cases}$$

in two independent samples case.

- Further assume that

- $\phi(Z) = \phi^*(Z) + \delta_1$ and $\psi(Z) = \psi^*(Z) + \delta_2$, where ϕ^*, ψ^* are fixed functions and δ_1, δ_2 are independent random variables with mean 0
- Z, δ_1, δ_2 are independent
- $\psi^*(Z) = \phi^*(Z) - \Delta \Rightarrow \Delta = \phi^*(Z) - \psi^*(Z)$

$$\begin{aligned} \mu_X &= E[\phi^*(Z)] \\ \mu_Y &= E[\psi^*(Z)] \end{aligned}$$

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Then,

$$\begin{cases} X_i = \phi^*(Z_i) + \delta_{1i} = \phi^*(Z_i) + \delta_{1i} = \mu_X + [\phi^*(Z_i) - \mu_X] + \delta_{1i} \\ Y_i = \psi^*(Z_i) + \delta_{2i} = \phi^*(Z_i) - \Delta + \delta_{2i} = \mu_Y + [\phi^*(Z_i) - \mu_X] + \delta_{2i} \end{cases}$$

same marginal distribution

- Q:** What are the sources of variation in ϵ 's and δ 's? If we apply the above formulation to the case of two independent samples, then

$$\begin{cases} X_i = \mu_X + \epsilon_{1i} = \phi^*(Z_i) + \delta_{1i} = \mu_X + (\phi^*(Z_i) - \mu_X) + \delta_{1i} \\ Y_j = \mu_Y + \epsilon_{2j} = \psi^*(Z_{n+j}) + \delta_{2j} = \mu_Y + (\phi^*(Z_{n+j}) - \mu_X) + \delta_{2j} \end{cases}$$

- A comparison

- Increase sample sizes: increase information about μ_X and μ_Y (signal)
- 2 independent \rightarrow paired: suppress the variation of error (noise)

Theorem 16 (A brief variance comparison of paired and independent samples)

Consider the models in the *dashed* frames. Under the two models,

- $\epsilon = [\phi^*(Z) - \mu_X] + \delta \Rightarrow \text{Var}(\epsilon) = \text{Var}[\phi^*(Z)] + \text{Var}(\delta) \geq \text{Var}(\delta)$
- 2 independent samples ($n = m$)
 - $X_i - Y_j = (\mu_X - \mu_Y) + (\epsilon_{1i} - \epsilon_{2j}) \equiv \sigma_\epsilon^2$
 - $= (\mu_X - \mu_Y) + [\phi^*(Z_i) - \phi^*(Z_{n+j})] + (\delta_{1i} - \delta_{2j})$
 - $\bar{X} - \bar{Y} = (\mu_X - \mu_Y) + (\bar{\epsilon}_1 - \bar{\epsilon}_2)$
 - $\Rightarrow \bar{X} - \bar{Y} \xrightarrow{e} \Delta$ and $\text{Var}(\bar{X} - \bar{Y}) = (\sigma_{\epsilon_1}^2/n) + (\sigma_{\epsilon_2}^2/n)$

- paired samples

- $D_i \equiv X_i - Y_i = (\mu_X - \mu_Y) + (\delta_{1i} - \delta_{2i})$

- $\bar{D} = \bar{X} - \bar{Y} = (\mu_X - \mu_Y) + (\bar{\delta}_1 - \bar{\delta}_2)$

$$\Rightarrow \bar{X} - \bar{Y} \xrightarrow{e} \Delta = \mu_X - \mu_Y \quad \text{and}$$

$$Var(\bar{X} - \bar{Y}) = (\sigma_{\delta_1}^2 / n) + (\sigma_{\delta_2}^2 / n) \longleftrightarrow$$

$Var(\bar{X} - \bar{Y})$ under the 2-independent-sample model with the sample size

$$n' = \frac{\sigma_{\epsilon}^2}{\sigma_{\delta}^2} n \quad (\geq n).$$

- Paired sample is more effective than independent samples in this case.

Theorem 17 (Conditions under which paired sample is more effective)

Consider the models in the *dotted* frames of LNp.48. Under the two models,

- $E(X) = E[\phi^*(Z) + \delta_1] = E[\phi^*(Z)] = \mu_X$

$$E(Y) = E[\psi^*(Z) + \delta_2] = E[\psi^*(Z)] = \mu_Y$$

- $Var(X) = Var[\phi^*(Z) + \delta_1] = Var[\phi^*(Z)] + Var(\delta_1) \equiv \sigma_X^2 (= \sigma_{\epsilon_1}^2)$

$$Var(Y) = Var[\psi^*(Z) + \delta_2] = Var[\psi^*(Z)] + Var(\delta_2) \equiv \sigma_Y^2 (= \sigma_{\epsilon_2}^2)$$

- 2 independent samples ($n = m$)

- $Cov(X_i, Y_j) = Cov[\phi^*(Z_i) + \delta_{1i}, \psi^*(Z_{n+j}) + \delta_{2j}] = 0$

- $E(\bar{X} - \bar{Y}) = \mu_X - \mu_Y = \Delta$

- $Var(\bar{X} - \bar{Y}) = (\sigma_X^2 + \sigma_Y^2) / n$

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- paired samples

- $Cov(X_i, Y_i) = Cov[\phi^*(Z_i) + \delta_{1i}, \psi^*(Z_i) + \delta_{2i}]$

$$= Cov[\phi^*(Z_i), \psi^*(Z_i)] \equiv \sigma_{XY}$$

* **Note.** We do not observe $(\phi^*(Z_i), \psi^*(Z_i))$'s. But, σ_{XY} can be estimated using (X_i, Y_i) 's data.

* Denote the correlation of (X_i, Y_i) by $\rho_{XY} = \frac{\sigma_{XY}}{\sigma_X \sigma_Y}$

* Notice that $\rho_{XY} \neq$

$$Cor[\phi^*(Z), \psi^*(Z)] = \frac{\sigma_{XY}}{\sigma_{\phi^*(Z)} \sigma_{\psi^*(Z)}}$$

- Let $D_i = X_i - Y_i, i = 1, \dots, n$. Then,

- * D_1, \dots, D_n are i.i.d.

- * $E(D_i) = \mu_X - \mu_Y$

- * $Var(D_i) = Var(X_i) + Var(Y_i) - 2Cov(X_i, Y_i) = \sigma_X^2 + \sigma_Y^2 - 2\sigma_{XY}$

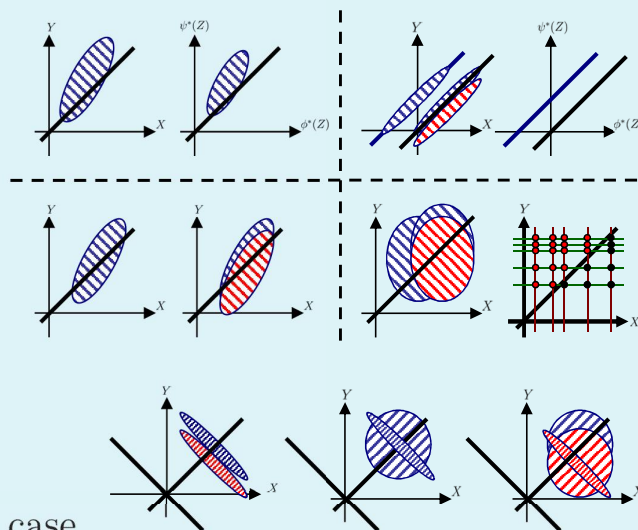
- Since $\bar{D} = \bar{X} - \bar{Y} \xrightarrow{e} \Delta$

- * $E(\bar{D}) = \mu_X - \mu_Y = \Delta$

- * $Var(\bar{D}) = Var(\bar{X} - \bar{Y}) = (\sigma_X^2 + \sigma_Y^2 - 2\sigma_{XY}) / n$

$$= (\sigma_X^2 + \sigma_Y^2 - 2\rho_{XY} \sigma_X \sigma_Y) / n$$

- If $\rho_{XY} > 0$ ($\Leftrightarrow \sigma_{XY} > 0 \Leftrightarrow \text{Cov}[\phi^*(Z), \psi^*(Z)] > 0$), then paired sample is more effective than independent samples.



- When $\psi^*(Z) = \phi^*(Z) - \Delta$,

$$\begin{aligned}\text{Cov}[\phi^*(Z), \psi^*(Z)] \\ &= \text{Cov}[\phi^*(Z), \phi^*(Z) - \Delta] \\ &= \text{Var}[\phi^*(Z)] > 0.\end{aligned}$$

- **Q:** Why are independent samples more effective than paired samples when $\sigma_{XY} < 0$?

- If $\sigma_X^2 = \sigma_Y^2 = \sigma^2$, then in the paired case

$$\sigma_D^2 = \text{Var}(\bar{D}) = [2\sigma^2(1 - \rho_{XY})]/n \quad \text{and} \quad \sigma_{\bar{X}-\bar{Y}}^2 = \text{Var}(\bar{X}-\bar{Y}) = 2\sigma^2/n$$

in the unpaired case. The relative efficiency is $\sigma_D^2/\sigma_{\bar{X}-\bar{Y}}^2 = 1 - \rho_{XY}$.

- If $\rho_{XY} = 0.5$, a paired design with n pairs of subjects yields the same precision as an unpaired design with $2n$ subjects per treatment.

- From now on, the analyses of paired data are based on

$$D_i = X_i - Y_i, \quad i = 1, \dots, n.$$

- Statistical modeling for D_i 's: $D_1, \dots, D_n \sim \text{i.i.d. } F \Leftarrow \text{one-sample model}$

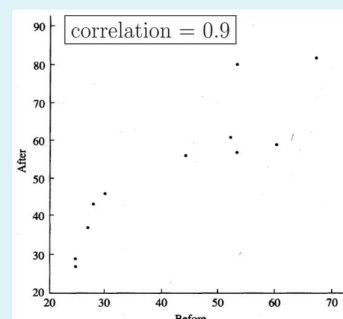
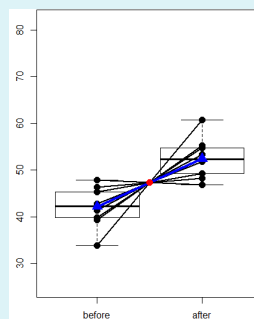
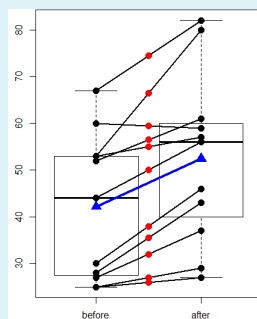
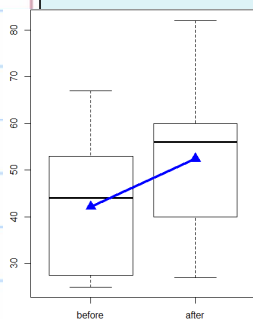
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Example 7 (Effect of cigarette smoking on platelet aggregation, Levine, 1973)

- Blood samples were drawn from 11 individuals before and after they smoked a cigarette to measure the extent to which the blood platelets aggregated.
- data: maximum percentage of all platelets that aggregated after being exposed to a stimulus.

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 |
|----------------|----|----|----|----|----|----|----|----|----|----|----|
| before (Y) | 25 | 25 | 27 | 44 | 30 | 67 | 53 | 53 | 52 | 60 | 28 |
| after (X) | 27 | 29 | 37 | 56 | 46 | 82 | 57 | 80 | 61 | 59 | 43 |
| difference (D) | 2 | 4 | 10 | 12 | 16 | 15 | 4 | 27 | 9 | -1 | 15 |

- **Q:** Do the differences D_i 's indicate a clear pattern of $\Delta = \mu_X - \mu_Y \neq 0$?
- The two-sample (unpaired) t -test for the before and after data gives a p -value = 0.1721 \Rightarrow the null $H_0 : \mu_X = \mu_Y$ is not rejected under $\alpha = 0.1$ ($s_p^2 = 289.34$).
 - **Q:** Why did the 2-sample t -test not reject H_0 when the differences showed such a clear pattern of $\mu_X > \mu_Y$?
 - Note that in 2-sample t -test, the test statistic is $|T| = \frac{|\bar{X} - \bar{Y}|}{s_p \sqrt{\frac{1}{n} + \frac{1}{m}}}$, where s_p^2 estimates σ_ϵ^2 , rather than σ_δ^2 .



- Figure 11.7 (textbook, p.447) plots after-values vs. before-values. They are positively correlated with a sample correlation coefficient 0.9. Pairing was a natural and effective experimental design in this case: relative efficiency = 0.1.

• Methods based on normality assumptions

- **Recall.** $D_i = X_i - Y_i, i = 1, \dots, n$, and $D_1, \dots, D_n \sim \text{i.i.d. } F$.
- Assume that F is Normal.
- Thus, the statistical model for D_i 's is

$$D_1, \dots, D_n \sim \text{i.i.d. } N(\mu_D, \sigma_D^2), \quad (\Delta)$$

where $\mu_D = \mu_X - \mu_Y$.

- This model contains two parameters: $\mu_D (\in \mathbb{R})$ and $\sigma_D^2 (> 0)$.
- Under this model, we can only examine whether there exists “difference” between the means of the two paired samples, i.e.,

$$\mu_D = \mu_X - \mu_Y = 0 \Rightarrow \text{no difference or no effect}$$

Theorem 18 (test and confidence interval for μ_D , 1-sample normal model, paired data)

Consider the model (Δ) .

- **Recall** (Review 1 in LNp.6-7).
 - $\bar{D} = \frac{1}{n} \sum_{i=1}^n D_i \xrightarrow{e} \mu_D$, and $\bar{D} \sim N(\mu_D, \sigma_D^2/n) \Rightarrow \sqrt{n}(\bar{D} - \mu_D)/\sigma_D \sim N(0, 1)$
 - $s_D^2 = \frac{\sum_{i=1}^n (D_i - \bar{D})^2}{n-1} \xrightarrow{e} \sigma_D^2$, and $(n-1)s_D^2 \sim \sigma_D^2 \chi_{n-1}^2 \Rightarrow (n-1)s_D^2/\sigma_D^2 \sim \chi_{n-1}^2$
 - \bar{D} and s_D^2 are independent

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- Pivotal quantity for μ_D
 - σ_D known: $Q_{Z, \mu_D} \equiv \frac{\bar{D} - \mu_D}{\sigma_{\bar{D}}} = \frac{\bar{D} - \mu_D}{\sigma_D/\sqrt{n}} = \frac{\sqrt{n}(\bar{D} - \mu_D)}{\sigma_D} \sim N(0, 1)$
 - σ_D unknown: $Q_{T, \mu_D} \equiv \frac{\bar{D} - \mu_D}{s_{\bar{D}}} = \frac{\bar{D} - \mu_D}{s_D/\sqrt{n}} = \frac{\sqrt{n}(\bar{D} - \mu_D)/\sigma_D}{\sqrt{\frac{(n-1)s_D^2/\sigma_D^2}{n-1}}} \sim t_{n-1}$
- Test the null and alternative hypotheses at significance level α :

$$H_0 : \mu_D = \mu_{D,0} \quad \text{vs.} \quad H_A : \mu_D \neq \mu_{D,0}$$

where $\mu_{D,0}$ is a known constant.

(**Note.** if $\mu_{D,0} = 0$, this is equivalent to $H_0 : \mu_X = \mu_Y$ vs. $H_A : \mu_X \neq \mu_Y$.)

 - σ_D known: reject H_0 if

$$\left| Z \equiv \frac{\bar{D} - \mu_{D,0}}{\sigma_{\bar{D}}} \right| > z(\alpha/2) \Leftrightarrow |\bar{D} - \mu_{D,0}| > z(\alpha/2) \sigma_{\bar{D}} = z(\alpha/2) \frac{\sigma_D}{\sqrt{n}}$$
 - σ_D unknown: reject H_0 if

$$\left| T \equiv \frac{\bar{D} - \mu_{D,0}}{s_{\bar{D}}} \right| > t_{n-1}(\alpha/2) \Leftrightarrow |\bar{D} - \mu_{D,0}| > t_{n-1}(\alpha/2) s_{\bar{D}} = t_{n-1}(\alpha/2) \frac{s_D}{\sqrt{n}}$$
- A $100(1 - \alpha)\%$ confidence interval for μ_D is
 - σ_D known: $\bar{D} \pm z(\alpha/2) \times \sigma_{\bar{D}} \Leftrightarrow \bar{D} \pm z(\alpha/2) \times (\sigma_D/\sqrt{n})$
 - σ_D unknown: $\bar{D} \pm t_{n-1}(\alpha/2) \times s_{\bar{D}} \Leftrightarrow \bar{D} \pm t_{n-1}(\alpha/2) \times (s_D/\sqrt{n})$

Example 8 (Effect of smoking, t -test for paired data, cont. Ex.7 In LNp.52)

- $n = 11$, $D_i = \text{after}_i - \text{before}_i$
- $\bar{D} = 10.27$, $s_{\bar{D}} = 2.405$ ($\Rightarrow s_D^2 = 11 \times 2.405^2 = 63.62$
 $\Rightarrow 63.62/2 = 31.81 \xrightarrow{e} \sigma_{\delta}^2 \xleftarrow{\text{cf.}} s_p^2 = 289.34 \xrightarrow{e} \sigma_{\epsilon}^2$ in Ex.7)
- A 90% confidence interval for μ_D is
 $\bar{D} \pm t_{10}(0.05) s_{\bar{D}} = 10.27 \pm 1.812 \times 2.40 = (5.9, 14.6)$,
 which does not contain zero ($\xleftarrow{\text{cf.}} H_0$ not rejected in Ex.7 using 2-sample t -test)
- The (one-sample) t -statistic is $T = (10.27 - 0)/2.40 = 4.28 > t_{10}(0.005) = 3.169$.
 The p -value of a two-sided test is less than 0.01. There is little doubt that smoking increases platelet aggregation.

If
 $\sigma_{\delta_1}^2 = \sigma_{\delta_2}^2 = \sigma_{\delta}^2$,
 then
 $\text{Var}(D_i) = 2\sigma_{\delta}^2$

Note 9 (Some notes about one-sample t -test when normality assumption does not hold)

- Consider the model: $D_1, \dots, D_n \sim \text{i.i.d. } F$,
 where F can be *any* continuous distributions with *finite* variance.
 - By CLT and LLN, when $n \rightarrow \infty$ (sample size is large),
 $\bar{D} \stackrel{D}{\approx} N(\mu_D, \sigma_D^2/n)$ and $s_D^2 \xrightarrow{P} \sigma_D^2$.
 - Thus, by Slutsky's Thm,
 $Q_{T, \mu_D} = \frac{(\bar{D} - \mu_D)/(\sigma_D/\sqrt{n})}{\sqrt{s_D^2/\sigma_D^2}} \stackrel{D}{\approx} N(0, 1)$
 and t_{n-1} tends to $N(0, 1)$ as $n \rightarrow \infty$.
- **Q:** What if the sample size n is small or population variance $= \infty$?

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A nonparametric method --- the signed rank test

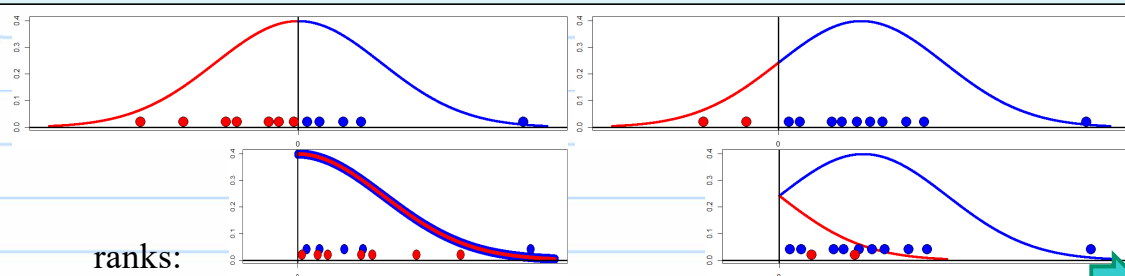
- Let Ω be the collection of *all* continuous distributions $\Rightarrow \dim(\Omega) = \infty$
- Consider the nonparametric statistical model: $D_1, \dots, D_n \sim \text{i.i.d. } F$, (∇)
 where $F \in \Omega$.
- Let $\Omega_0 = \{F \mid F \in \Omega \text{ and } F \text{ is symmetric about } 0\}$
 - $\Omega_0 \subset \Omega$ and $\dim(\Omega_0) = \infty$
 - If $F \in \Omega_0$, then the median of F is 0. But, F with median zero is not necessary a distribution being symmetric about 0.
- Under the model (∇) , we want to test the null and alternative hypotheses:
 $H_0 : F \in \Omega_0$ vs. $H_A : F \in \Omega \setminus \Omega_0$

Q: Why add the “symmetric” condition in the null?

Question 8.

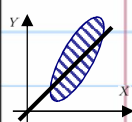
How to use ranks to examine “symmetric about 0”? What data are “more extreme,” i.e., cast more doubts on H_0 ?

Intuition.





- A brief comparison to (2-sample) rank sum test (assume no $D_i = 0$)
 - similarity: if in the paired case,
 - * the data $\{-D_i \mid D_i < 0\}$ is treated as the 1st sample
 - * the data $\{D_i \mid D_i > 0\}$ is treated as the 2nd sample
 - * then, the calculation for the paired case is equivalent to the rank-sum statistic in the unpaired case
 - difference
 - * In 2-sample unpaired cases, the sample sizes m, n are fixed numbers.
 - * In the paired case, the sizes



$$N_- = \#\{D_i < 0\} \quad \text{and} \quad N_+ = \#\{D_i > 0\}$$

(Note. $N_- + N_+ = n$) are random variables.

- * Under H_0 ,
 - $I_{[D_1 > 0]}, \dots, I_{[D_n > 0]} \sim \text{i.i.d. Bernoulli}(1/2)$,
 - $N_+ = \sum_{i=1}^n I_{[D_i > 0]} \sim \text{bin}(n, 1/2)$ and $N_- = n - N_+ \sim \text{bin}(n, 1/2)$
- * When conditioned on N_+ (or N_-), the null distribution of the test statistic in the paired case is identical to the null distribution of rank-sum statistic in the unpaired case.
- Alternative test: sign test (TBp.461, problem 12)
 - Consider the model (∇) in LNp.56.

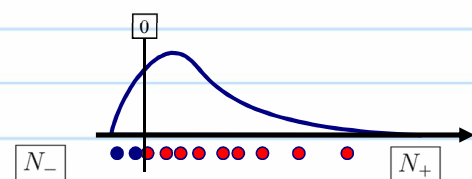
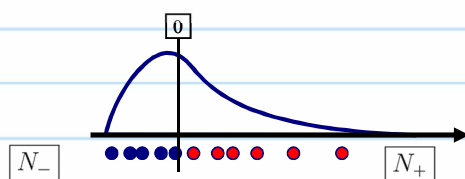
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- Let $\Omega_0^* = \{F \mid F \in \Omega \text{ and } F \text{ has median } 0\}$
 - * $\Omega_0 \subset \Omega_0^* \subset \Omega$ and $\dim(\Omega_0^*) = \infty$
- Under the model (∇) , test the null and alternative hypotheses:

$$H_0^* : F \in \Omega_0^* \quad \text{vs.} \quad H_A^* : F \in \Omega \setminus \Omega_0^*$$

– **Intuition.**

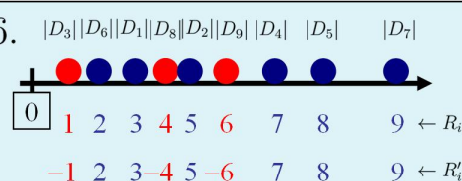


- Reject H_0^* if N_+ (or N_-) is small (close to 0) or large (close to n)
- Null distribution of N_+ (or N_-): $\text{bin}(n, 1/2)$

Theorem 19 (Wilcoxon signed rank test)

Consider the nonparametric model (∇) in LNp.56.

- test statistic W_+ (or $W_- = \frac{n(n+1)}{2} - W_+$)



- (1) Let $R_i = \text{rank of } |D_i|, i = 1, \dots, n$.
 - (2) Restore the signs of D_i 's to the ranks R_i 's, i.e., let $R'_i = \text{sign}(D_i) \times R_i$.
 - (3) $W_+ = \sum_{i=1}^n I_{[D_i > 0]} R'_i$, i.e., sum of the ranks R'_i 's that have positive signs.
- **Q:** What values of W_+ are more extreme? If there is no difference between the two paired conditions, we expect



- about *half* the D_i 's to be positive and *half* negative (median=0?)
 - positive R_i 's and negative R_i 's *similarly* distributed (symmetric?)
- and W_+ will not be too small or too large
 \Rightarrow data with larger or smaller W_+ are more extreme \Rightarrow tend to reject H_0

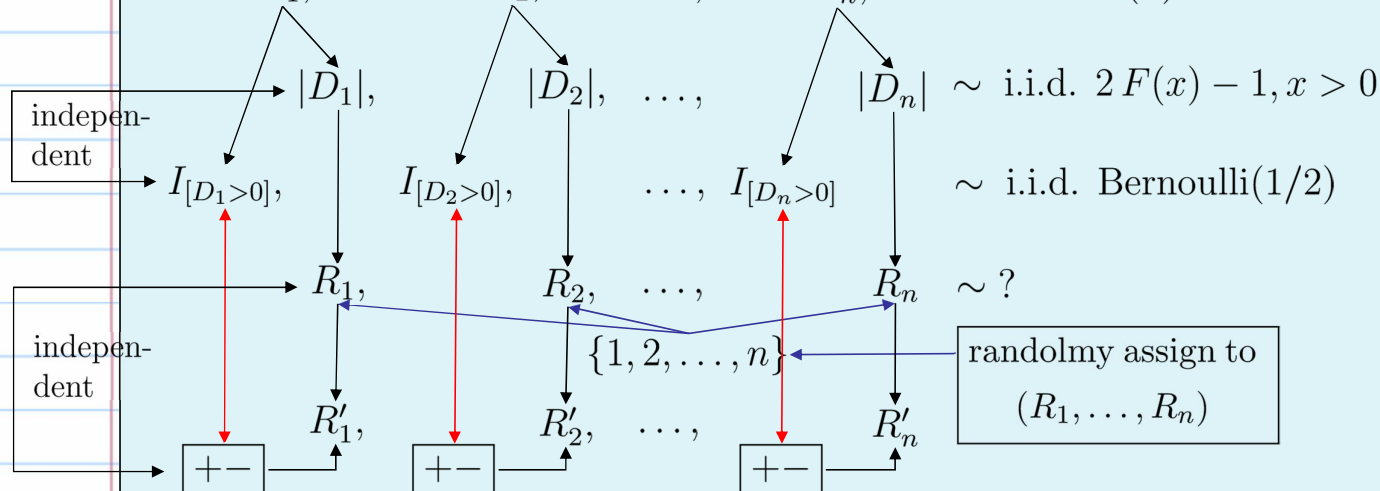
- Null distribution of W_+

- $W_+ \in \{0, 1, 2, \dots, \frac{n(n+1)}{2}\}$

- Under the null H_0 (F is symmetric about 0)

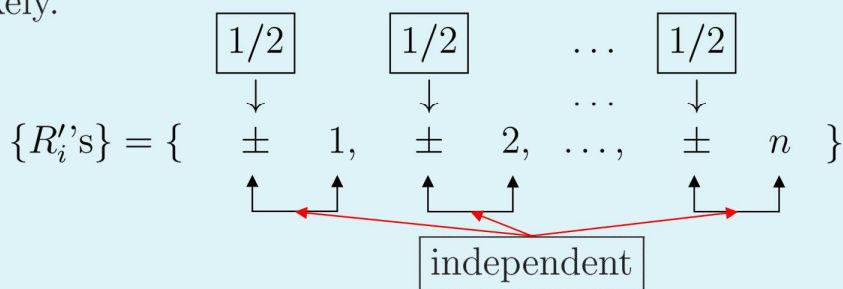
- * $D_i \Leftrightarrow (I_{[D_i>0]}, |D_i|)$ and D_i has the same distribution as $-D_i$

- * $D_1, D_2, \dots, D_n \sim \text{i.i.d. } F(x)$



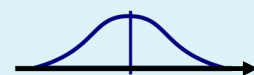
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- * Any particular assignment of $\{-, +\}$ signs to the integers $1, \dots, n$ (the ranks) is equally likely.



- * There are 2^n such assignments and for each we can calculate $W_+ \Rightarrow$ obtain 2^n values (not all distinct) of W_+ , each with probability $1/2^n$.

- * The probability of each distinct value of W_+ may thus be calculated, giving the desired null distribution.



- (Two-sided) rejection region

- (exercise) – The null distribution of W_+ is symmetric around $E(W_+)$
- Reject H_0 when $\min(W_+, W_-)$ is small, i.e., $\min(W_+, W_-) \leq w$
- Table 9 of Appendix B in textbook (TBp.A24) gives critical values w

- Ties

- Tie between (X_i, Y_i) : If some of the differences D_i 's are zero, the most common technique is to discard those observations.

- Tie between $|D_i|$'s: If there are ties, each $|D_i|$ is assigned the average value of the ranks for which it is tied.
- If there are a large number of ties, modifications must be made. See Hollander and Wolfe (1973) or Lehmann (1975).

Example 9 (Smoking effect, signed-rank test for paired data, cont. Ex.7 In LNp.52)

- $n = 11$, $W_- = 1$ and $W_+ = [11(11 + 1)]/2 - W_- = 65 \Rightarrow \min(W_-, W_+) = 1$
- From Table 9 of Appendix B (TBp.A24), the critical value for two-sided test with significant level $\alpha = 0.01$ is 5.
- Since $\min(W_-, W_+) < 5$, reject H_0 at $\alpha = 0.01$ (consistent with the test result in Ex.8, LNp.55).

Note 10 (A comparison of one-sample t -test and signed rank test for paired data)

- Unlike (one-sample) t -test, the signed-rank test does not depend on normality assumption.
- The signed-rank test is insensitive to outliers, whereas the t -test is sensitive.
- When the normality assumption holds, the t -test is more powerful.
- However, it has been shown that even when normality assumption holds, the signed-rank test is nearly as powerful as the t -test (relative efficiency of signed-rank test statistic to (one-sample) t -test statistic ≈ 0.95).
- The signed-rank test is generally preferable, especially for *small* sample sizes.

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Theorem 20 (means and variances of W_+ under H_0)

- Consider the nonparametric model (∇) in LNp.56.
- Under the null hypothesis H_0 : F is symmetric about 0,

$$E(W_+) = \frac{n(n+1)}{4} \quad \text{and} \quad \text{Var}(W_+) = \frac{n(n+1)(2n+1)}{24}.$$

$$(\Leftrightarrow E(W_-) = [n(n+1)]/4 \quad \text{and} \quad \text{Var}(W_-) = [n(n+1)(2n+1)]/24 \\ \text{since } W_- = [n(n+1)]/2 - W_+)$$

Proof.

- For $k = 1, \dots, n$, let $I_k = \begin{cases} 1, & \text{if the } k\text{th largest } |D_i| \text{ has } D_i > 0, \\ 0, & \text{otherwise.} \end{cases}$

- Under H_0 ,

$$- I_1, \dots, I_n \sim \text{i.i.d. Bernoulli}(1/2),$$

$$- E(I_k) = 1/2 \quad \text{and} \quad \text{Var}(I_k) = 1/4.$$

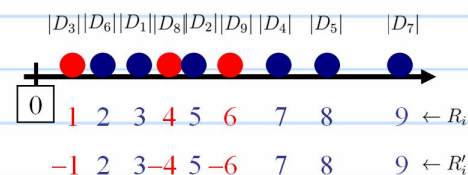
- Write

$$W_+ = \sum_{i=1}^n I_{[D_i > 0]} R'_i = \sum_{k=1}^n k I_k.$$

- Thus,

$$E(W_+) = \sum_{k=1}^n k E(I_k) = \frac{1}{2} \left(\sum_{k=1}^n k \right) = \frac{n(n+1)}{4}$$

$$\text{Var}(W_+) = \sum_{k=1}^n k^2 \text{Var}(I_k) = \frac{1}{4} \left(\sum_{k=1}^n k^2 \right) = \frac{n(n+1)(2n+1)}{24}$$



Theorem 21 (Asymptotic null distribution of W_+)

- Consider the nonparametric model (∇) in LNP.56.
- Under the null $H_0 : F$ is symmetric about 0, if the sample size n is greater than 20, the null distribution of W_+ is well approximated by a normal distribution, i.e.,

$$\frac{W_+ - E(W_+)}{\sqrt{Var(W_+)}} \stackrel{D}{\approx} N(0, 1) \quad \left(\text{or } \frac{W_- - E(W_-)}{\sqrt{Var(W_-)}} \stackrel{D}{\approx} N(0, 1) \right).$$

Hint for Proof. Use the expression $W_+ = \sum_{k=1}^n k I_k$ to find the moment generating function of W_+ , and show it converges (after standardization) to the moment generating function of $N(0, 1)$, which is $e^{t^2/2}$.

❖ **Reading:** textbook, 11.3