

Statistic take home exam

Z test : $X_1, \dots, X_n \sim N(\mu_x, 1)$ i.i.d.
 $Y_1, \dots, Y_n \sim N(\mu_y, 1)$ i.i.d. } independently from each other

def $\bar{X}_n = \frac{1}{n} \sum_{i=1}^n X_i$, $\bar{Y}_n = \frac{1}{n} \sum_{i=1}^n Y_i$

def test statistic $S = \frac{(\bar{Y}_n - \bar{X}_n)}{\sqrt{\frac{2}{n}}}$

Valid p-values : $\cdot p_A := 1 - \Phi(S)$ for $\underline{H_0 : \mu_y = \mu_x}$
 against $\underline{H_1^A : \mu_y > \mu_x}$

$\cdot p_B := 1 - \Phi(-S)$ for $\underline{H_0 : \mu_y = \mu_x}$
 against $\underline{H_1^B : \mu_y < \mu_x}$

Obj. test $H_0 : \mu_x = \mu_y$ against $H_1 : \mu_y \neq \mu_x$

$\left\{ \begin{array}{ll} \text{if } \bar{Y}_n > \bar{X}_n & \text{report } p_A \\ \text{if } \bar{Y}_n < \bar{X}_n & \text{report } p_B \end{array} \right.$ [1]

a) Is this a valid p-value?

We can write [1] as:

$$p(x) = \begin{cases} p_A = 1 - \Phi(S(x)) & \text{if } \bar{Y}_n > \bar{x}_n \\ p_B = 1 - \Phi(-S(x)) & \text{if } \bar{Y}_n < \bar{x}_n \end{cases}$$

$$= \begin{cases} 1 - \Phi \left[\frac{(\bar{Y}_n - \bar{x}_n)}{\sqrt{\frac{\sigma^2}{n}}} \right] & \text{if } \bar{Y}_n < \bar{x}_n \\ 1 - \Phi \left[\frac{\bar{x}_n - \bar{Y}_n}{\sqrt{\frac{\sigma^2}{n}}} \right] & \text{if } \bar{Y}_n > \bar{x}_n \end{cases}$$

Since we are changing the sign we can use the abs. value

$$p(x) = 1 - \Phi \left[\left| \bar{Y}_n - \bar{x}_n \right| \cdot \sqrt{\frac{n}{\sigma^2}} \right]$$

We know that a valid p-value satisfies:

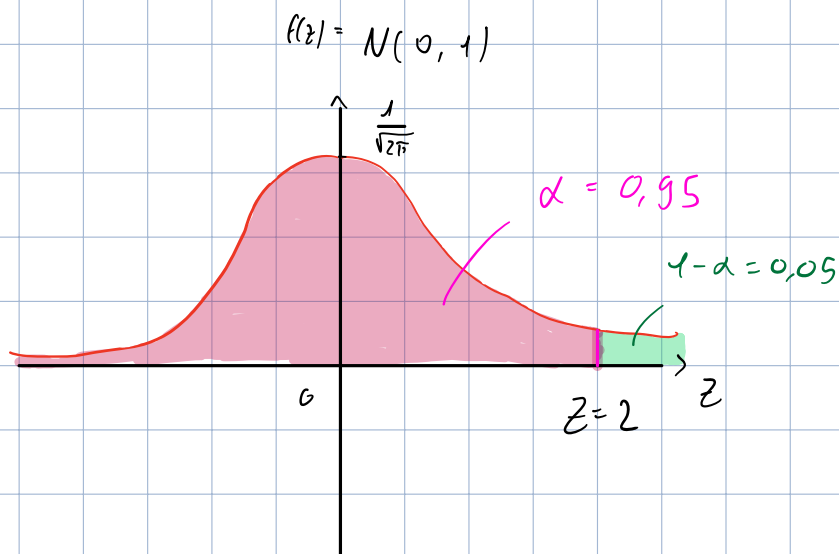
$$\alpha = P_{H_0} (p(x) < \alpha)$$

$$= P_{H_0} \left[1 - \Phi \left(\left| \bar{Y}_n - \bar{x}_n \right| \sqrt{\frac{n}{\sigma^2}} \right) < \alpha \right] =$$

$$= P_{H_0} \left[\Phi \left(\left| \bar{Y}_n - \bar{x}_n \right| \sqrt{\frac{n}{\sigma^2}} \right) > 1 - \alpha \right] =$$

$$= P_{H_0} \left[\left| \bar{Y}_n - \bar{x}_n \right| \sqrt{\frac{n}{\sigma^2}} > \Phi^{-1}(1 - \alpha) \right] =$$

NOTE: $\alpha \in [0, 1]$



this is the valid p-value for test A

$$P_{H_0} (1 - \Phi(z) > \alpha) = \alpha$$

$$= P_{H_0} \left[\left(\bar{Y}_n - \bar{X}_n \right) \sqrt{\frac{n}{2}} > \Phi^{-1}(1-\alpha) \right] +$$

$$+ P_{H_0} \left[\left(\bar{Y}_n - \bar{X}_n \right) \sqrt{\frac{n}{2}} < -\Phi^{-1}(1-\alpha) \right] =$$

this is the valid p-value for test B

$$P_{H_0} (1 - \Phi(-z) > \alpha) = \alpha$$

Since p_A and p_B
are valid p-values

$$= \alpha + \alpha = 2\alpha$$

We get $\alpha = 2\alpha$ which is valid only for $\alpha = 0$.

$$\Rightarrow P(x) = 1 - \Phi \left[\left| \bar{Y}_n - \bar{X}_n \right| \sqrt{\frac{n}{2}} \right] \text{ is not a valid p-value.}$$

b) Correct the p-value:

I can correct the p value considering

$$p(x) = 2 \left[1 - \Phi \left(|\bar{Y}_n - \bar{X}_n| \sqrt{\frac{n}{2}} \right) \right]$$

And we can verify it:

$$\alpha = P_{H_0} (p(x) < \alpha)$$

$$= P_{H_0} \left(2 \left[1 - \Phi \left(|\bar{Y}_n - \bar{X}_n| \sqrt{\frac{n}{2}} \right) \right] < \alpha \right)$$

$$= P_{H_0} \left(\Phi \left(|\bar{Y}_n - \bar{X}_n| \sqrt{\frac{n}{2}} \right) > 1 - \frac{\alpha}{2} \right) =$$

$$= P_{H_0} \left(|\bar{Y}_n - \bar{X}_n| \sqrt{\frac{n}{2}} > \Phi^{-1} \left(1 - \frac{\alpha}{2} \right) \right) +$$

$$+ P_{H_0} \left(|\bar{X}_n - \bar{Y}_n| \sqrt{\frac{n}{2}} > \Phi^{-1} \left(1 - \frac{\alpha}{2} \right) \right)$$

this is the valid
p-value for test A

$$P_{H_0} (1 - \Phi(t) > \beta) = \beta$$

I can substitute $\frac{\alpha}{2} = \beta$

this is the valid
p-value for test B

$$P(1 - \Phi(-s) > \beta) = \beta$$

$$= \beta + \beta = \alpha$$

This proves that $p(x) = 2 \left[1 - \Phi \left(|\bar{Y}_n - \bar{X}_n| \sqrt{\frac{n}{2}} \right) \right]$

is a valid p-value.

c) $\Delta_{\mu} := (\mu_Y - \mu_X) \sqrt{\frac{2}{n}} > 0$

See Exercise_1_C.R file in the submission folder to run the code.

Exercise 2

See the file: Exercise_2.v

It prints all the required graphs

NOTE: When we order our data, the median is the point in the $(n//2)+1$ position.

Since we want to compute which is the maximum local sensitivity I want to achieve it I change K entries of my data.

This can be visualize as moving the middle value to 0 or to the max value. This means that the new middle point is one of the adjacent values.

The function `get_indices` provides the indices of the potential new middle point.

NOTE: For each potential new middle point, the local sensitivity is computed. Subsequently we store the maximum.

Exercise 3

We have the population graph $G = (V, E)$

and we consider an estimation

$$\tau = \sum_{e \in E} y_e$$

Using the Horowitz Thompson approach :

$$\hat{\tau} = \sum_{e \in E^*} \frac{y_e}{\pi_e^{(1)}}$$

probability that
the edge e is sampled

$$\pi_e^{(1)} = P(e \in E^*) \quad [7]$$

We already computed the vertex pair inclusion probability

$$\pi_{\{i,j\}}^{(2)} = P(\text{"vertex pair } \{i,j\} \text{ is sampled"})$$

for all the $\{i,j\} \in V^{(2)}$.

Consider $\hat{\tau} = \sum_{e \in E^*} \frac{y_e}{\pi_e^{(2)}}$ as the estimator for τ

Prove that it is in general unbiased

pf. to prove that it is not unbiased let's consider its value in expectation:

$$E(\hat{\tau}) = E\left(\sum_{e \in E^*} \frac{Y_e}{\hat{\pi}_e^{(2)}}\right) =$$

$$= E\left(\sum_{e \in E} \frac{Y_e}{\hat{\pi}_e^{(2)}} \mathbb{I}_{E^*}\right) =$$

I can consider /
the sum of all
the $e \in E$ and
add the indicator
function

$$\mathbb{I}_{E^*} = \begin{cases} 0 & e \notin E^* \\ 1 & \text{else} \end{cases}$$

in expectation we
have that this is
equal to $P(e \in E^*)$

$$\stackrel{\text{linearity}}{=} \sum_{e \in E} \frac{Y_e}{\hat{\pi}_e^{(2)}} E(\mathbb{I}_{E^*})$$

$$= \sum_{e \in E} \frac{Y_e}{\hat{\pi}_e^{(2)}} P(e \in E^*)$$

$$\stackrel{[1]}{=} \sum_{e \in E} \frac{Y_e}{\hat{\pi}_e^{(1)}} \hat{\pi}_e^{(1)}$$

To have an unbiased estimator we require:

$$\tau := \sum_{e \in E} Y_e = \sum_{e \in E} \frac{Y_e}{\hat{\pi}_e^{(2)}} \hat{\pi}_e^{(1)} := \hat{\tau}$$

which is true only for

$$\hat{\pi}_e^{(1)} = \hat{\pi}_e^{(2)}$$

Let's consider an example where $\tilde{\pi}_e^{(1)} \neq \tilde{\pi}_e^{(2)}$.

I choose to consider the unlabeled star sampling:

$$\circ \quad \tilde{\pi}_e^{(1)} = 1 - \binom{N_v - 2}{n}$$

$$\circ \quad \tilde{\pi}_e^{(2)} = \frac{n(n-1)}{N_v(N_v-1)}$$

where N_v is the number of vertex.

$$1 - \binom{N_v - 2}{n} \neq \frac{n(n-1)}{N_v(N_v-1)}$$

We can see that, in a general setting:

$$\tilde{\pi}_e^{(1)} \neq \tilde{\pi}_e^{(2)}$$

