

1 Multiple Testing

A *multiple testing problem* is one where we test > 1 hypotheses using same data set.

Example 1.1. $X_i \stackrel{\text{iid}}{\sim} N(\mu_i, 1), i = 1, \dots, n$
 $H_{0,i} : \mu_i = 0, H_{1,i} : \mu_i \neq 0$

Common setup:

- Measuring crop yields / patient outcomes under different treatments
- Gene expression of n genes in cancer vs controls / FMRI meas. in n voxels

Example 1.2. $p_i \in [0, 1], i = 1, \dots, n (i \in \mathcal{I})$
 $H_{0,i} : p_i \sim U[0, 1]$ [or p_i stoch. larger than $U[0, 1]$]
 $H_{1,i} : p_i$ “smaller”
 Genome-wide association study (GWAS), for SNP i

	healthy	diseased
wild-type	·	·
mutant	·	·

Observe $X \sim P \in \mathcal{P}$, return set of rejected hypotheses.

[Ex. 1] Suppose $n = 100, \mu_i = 0, \forall i = 1, \dots, n$,

$\mathbb{E}[\# \text{Rejections}] = n\alpha = 5 (\alpha = 0.05)$

$\mathbb{P}[\text{at least one false rejection}] \approx 1$

Classic proposal: control *familywise error rate* (FWER)

$$FWER = \sup_{P \in \mathcal{P}} \mathbb{P}_P[\text{any true } H_{0,i} \text{ rejected}] \quad (1.1)$$

1.1 Bonferroni / Šidák Correction

First rule (Bonferroni): Reject $H_{0,i}$ if $p_i \leq \frac{\alpha}{n}$

$$\mathbb{P}[\text{any false rejection}] = \mathbb{P}[\cup_{i \in \mathcal{H}_0} H_{0,i} \text{ reject}] \quad (1.2)$$

$$\leq \sum_{i \in \mathcal{H}_0} \mathbb{P}[H_{0,i} \text{ rejected}] \leq |\mathcal{H}_0| \frac{\alpha}{n} \leq \alpha \quad (1.3)$$

where $\mathcal{H}_0 : \{i : H_{0,i} \text{ true}\}$.

More liberal (Šidák): If p_i independent, can use the rule: Reject $H_{0,i}$ if $p_i \leq \tilde{\alpha}_n$ where $\tilde{\alpha}_n = 1 - (1 - \alpha)^{1/n} \approx \frac{\alpha}{n}$. Improvement in power over Bonferroni: $\tilde{\alpha}_n > \frac{\alpha}{n}$

1.2 Correlated Statistics

Often test stats are dependent

Example 1.3 (Pairwise comparisons). $X_i \stackrel{\text{ind.}}{\sim} N(\mu_i, 1) = \mu_i + \varepsilon_i$.

$H_{0,ij} : \mu_i = \mu_j, H_{1,ij} : \mu_i \neq \mu_j$

$\binom{n}{2}$ hypotheses.

$X_i - X_j = \mu_i - \mu_j + (\varepsilon_i - \varepsilon_j) \sim N(\mu_i - \mu_j, 2)$

Could use Bonferroni: Reject if $\frac{|X_i - X_j|}{\sqrt{2}} > z_{\frac{\alpha}{2\binom{n}{2}}}$

More powerful: Reject $H_{0,ij}$ if $|X_i - X_j| > r_\alpha$ where

$$\mathbb{P} \left[\max_{i,j} |\varepsilon_i - \varepsilon_j| > r_\alpha \right] = \alpha \quad (1.4)$$

This is less conservative than the Bonferroni correction: apply the union bound to get

$$\mathbb{P} \left[\max_{i,j} |\varepsilon_i - \varepsilon_j| > z_{\frac{\alpha}{2\binom{n}{2}}} \sqrt{2} \right] \leq \binom{n}{2} \mathbb{P} \left[|\varepsilon_i - \varepsilon_j| > z_{\frac{\alpha}{2\binom{n}{2}}} \sqrt{2} \right] \quad (1.5)$$

$$= \binom{n}{2} \frac{\alpha}{\binom{n}{2}} = \alpha \quad (1.6)$$

This is called *Tukey's Honestly Significant Difference* procedure

Example 1.4. Comparing all linear combos, $X_i \sim \text{ind.} N(\mu_i, 1)$.

$H_{0,v} : v' \mu = 0, v \in \mathbb{R}^n$

Reject when $\frac{|X'v|}{\|v\|} > \chi_n(\alpha)$.

This is because

$$\mathbb{P}[\text{any false rej.}] = \mathbb{P} \left[\frac{|X'v|}{\|v\|} > \chi_n(\alpha), \text{ any } v \text{ s.t. } v' \mu = 0 \right] \quad (1.7)$$

$$\leq \mathbb{P} \left[\max_{\|v\|=1} |\varepsilon'v| > \chi_n(\alpha) \right] \quad (1.8)$$

$$= \mathbb{P} [\|\varepsilon\| > \chi_n(\alpha)] = \alpha \quad (1.9)$$

called *Scheffé's S-method*

$$\chi_n(\alpha) \approx \sqrt{n} \gg z_{\frac{\alpha}{2\binom{n}{2}}} \sqrt{2} \quad (1.10)$$

$$\approx \sqrt{2 \cdot 2 \log \binom{n}{2}} \approx 2\sqrt{2 \log n} \quad (1.11)$$

More generally $X_i \sim N(\mu_i, \sigma^2)$ where σ^2 is unknown, $\hat{\sigma}^2 \sim \sigma^2 \chi_d^2 \perp X$.

Test $H_{0,v} : \mu'v = 0, \forall v \in \Xi \subset \mathbb{R}^n$.

Reject $H_{0,v}$ if $\frac{|X'_i v|}{\hat{\sigma} \|v\|} \geq c_\alpha$

$$\mathbb{P} \left[\sup_{v \in \Xi} \frac{|\varepsilon'_i v|}{\hat{\sigma} \|v\|} > c \right] = \quad (1.12)$$

2 Simultaneous Intervals

Construct CI for ...