STA 35C: Statistical Data Science III

Lecture 18: Multiple Hypotheses Testing

Dogyoon Song

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Announcement

Midterm 2 on Fri, May 16 (12:10 pm-1:00 pm in class)

- Arrive early: The exam starts at 12:10 pm and ends at 1:00 pm sharp
- One hand-written cheat sheet: Letter-size (8.5"×11"), double-sided, brief formulas/notes
- Calculator: A simple (non-graphing) scientific calculator is allowed
- No other materials beyond the single cheat sheet (no textbooks, etc.)
- SDC accommodations: Confirm scheduling with AES online ASAP

Preparation tips:

- Primary coverage: Lectures 12–19 (including Wed)
- A practice midterm and answer key are available on the course webpage
- Office hours this week:
 - Instructor: Wed, 4-6pm (extended); no OH Thu
 - TA: Mon/Thu 1–2pm

Today's topics

Multiple hypotheses testing

- Recap: Motivation & challenges
 - Issues arising with multiple tests
 - Real-world concerns: p-hacking & data dredging
- Family-wise error rate (FWER)
 - Definition & intuition
 - Controlling FWER: Bonferroni correction & Holm's step-down
- False discovery rate (FDR)
 - Definition & intuition
 - Controlling FDR: Benjamini-Hochberg procedure

Recap: Multiple testing

Single-hypothesis test:

- Typically set up H_0 , and gather data to reject it if there is significant evidence
- Type I error = false positive; Type II error = false negative
- Each test has Type I error at most α (e.g. 0.05)

Modern data analysis: multiple tests simultaneously

- E.g. Testing thousands of predictors or biomarkers to discovery significant ones
- If m is large, false rejections can occur easily by chance
- ullet On average, lpha imes m false positives if each test is at level lpha

Key challenge: Address the inflation of false positives as *m* grows

Related issues: p-hacking and data dredging

Real danger: Searching for "significant" results in many ways until something "works"

- Repeatedly testing different hypotheses/subgroups
- Eventually, some test may yield p < 0.05 by chance

Outcome: Spurious "discoveries"

- Published claims may fail to replicate
- True findings can be overshadowed by noise

Conclusion: Systematic multiple-testing corrections are crucial, especially for large m

Articles warning about misused statistical significance





Figure: Many reproducibility crises trace back to undisclosed multiple testing or selective reporting. Proper adjustments can help mitigate these issues.

Family-wise error rate (FWER): Definition

Single test:

- H_0 : "no signal" vs. H_a : "signal"
- Reject H₀: "Discovery" of "signal"

	H_0 is true	H_0 is not true
Reject H ₀	Type-I error (FP)	Correct (TP)
Not reject H_0	Correct (TN)	Type-II error (FN)

- \implies Pr(Type I error) = Pr(reject a true null)
 - By setting threshold α , we want to control Pr(Type I error) below α

Family-wise error rate (FWER): Recall single test

Single test: $Pr(Type \mid error) = Pr(reject a true null)$

Multiple tests (*m* hypotheses):

$$\begin{aligned} \mathrm{FWER} &= \mathsf{Pr}(\mathsf{reject} \ \mathsf{at} \ \mathsf{least} \ \mathsf{one} \ \mathsf{true} \ \mathcal{H}_0) \\ &= \mathsf{Pr}(\# \ \mathsf{Type-I} \ \mathsf{error} \geq 1), \end{aligned}$$

i.e. the probability of any false positive among m tests

If tests are independent, and each are at level α :

$$FWER = 1 - (1 - \alpha)^m,$$

- When m = 1, FWER = $1 (1 \alpha)^m = 1 (1 \alpha) = \alpha$
- Grows quickly with *m*
 - E.g. m = 20, $\alpha = 0.05 \implies \text{FWER} \approx 0.64 \gg 0.05$

Visualization: FWER grows as *m* increases

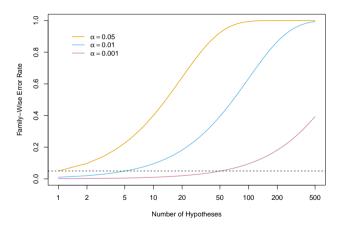


Figure: FWER vs. number of tests m (log scale) for $\alpha=0.05$ (orange), 0.01 (blue), 0.001 (purple). The dashed line is 0.05. For m=50 and target FWER=0.05, each test must be at $\alpha=0.001$ [JWHT21, Figure 13.2].

The Bonferroni correction

Key idea: Observe that

$$\text{FWER} = \Pr\left(\sum_{j=1}^{m} \left\{ \text{Reject } H_j \right\} \right) \leq \sum_{j=1}^{m} \Pr\left(\left\{ \text{Reject } H_j \right\} \right)$$

• Each test is done at level $\alpha/m \implies \Pr(\{\text{Reject } H_j\}) \leq \alpha/m \implies \text{FWER} \leq \alpha$

The Bonferroni method (Bonferroni correction):

• For each hypothesis H_1, \ldots, H_m , reject H_j if only if $p_j < \frac{\alpha}{m}$

Pros & Cons:

- Pros: Simple & widely used
- Cons: Often very conservative \implies few rejections (=discoveries) & lower power¹

 $^{^{1}\}mathsf{Power} = \mathsf{TPR} = \mathsf{the}$ fraction of false null hypotheses that are successfully rejected

Example: Bonferroni correction

Example

Let m = 6 hypotheses with p-values:

$$p_1 = 0.0018$$
, $p_2 = 0.009$, $p_3 = 0.021$, $p_4 = 0.034$, $p_5 = 0.045$, $p_6 = 0.070$.

At $\alpha=$ 0.05, threshold $=\frac{\alpha}{m}=\frac{0.05}{6}\approx$ 0.00833.

Reject
$$H_j$$
 if $p_j < 0.00833$.

Hence:

$$p_1 = 0.0018 < 0.00833 \implies \text{reject } H_1,$$

but $p_2 = 0.009 > 0.00833$ and the rest are larger. So Bonferroni rejects only H_1 .

Conclusion: 1 rejection using Bonferroni, whereas naive p < 0.05 would reject 5 of them (p_1, \ldots, p_5) .

Holm's step-down procedure

Holm's method refines Bonferroni to be less conservative:

Holm's method

- 1 Specify α , the level at which to control the FWER
- 2 Compute the *p*-values for the *m* null hypotheses, H_{01}, \ldots, H_{0m}
- 3 Sort p-values so that $p_{(1)} \leq p_{(2)} \leq \cdots \leq p_{(m)}$
- 4 Define

$$L = \min \left\{ j : p_{(j)} > \frac{\alpha}{m+1-j} \right\}$$

5 Reject all null hypotheses H_{0j} for which $p_j < p_{(L)}$

Properties:

- Ensures $FWER \leq \alpha$
- Rejects at least as many hypotheses as Bonferroni

Example: Holm's step-down procedure

Example

- **Step 1**: Set $\alpha = 0.05$
- **Step 2:** $p_1 = 0.0018, p_2 = 0.009, p_3 = 0.021, p_4 = 0.034, p_5 = 0.045, p_6 = 0.070.$
- **Step 3:** Sort *p*-values $p_{(1)} = 0.0018$, $p_{(2)} = 0.009$, $p_{(3)} = 0.021$, $p_{(4)} = 0.034$, $p_{(5)} = 0.045$, $p_{(6)} = 0.070$.
- **Step 4:** Find L = 3 because

$$p_{(1)} = 0.0018 ? 0.0018 \le \frac{0.05}{6+1-1} = \frac{0.05}{6} \approx 0.00833$$
 \implies reject $H_{(1)}$, continue $p_{(2)} = 0.009 ? 0.009 \le \frac{0.05}{6+1-2} = \frac{0.05}{5} = 0.01$ \implies reject $H_{(2)}$, continue $p_{(3)} = 0.021 ? 0.021 \le \frac{0.05}{6+1-3} = \frac{0.05}{4} = 0.0125?$ No \implies stop; $L = 3$

Step 5: We reject $H_{(1)}$, $H_{(2)}$ total 2 rejections. The rest are not rejected.

Conclusion: Holm's method rejects 2, whereas Bonferroni rejected only 1.

Visualization: Bonferroni vs. Holm

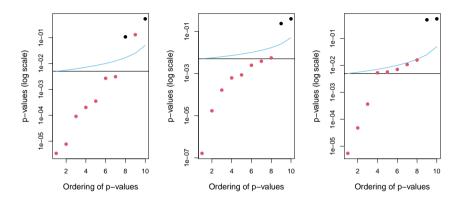


Figure: Each panel shows sorted p-values from a separate simulation of m=10 null hypotheses, with the two true nulls in black and the others in red. Controlling the FWER at 0.05, Bonferroni rejects all points below the **black** line, while Holm rejects all below the **blue** line. The gap between these lines indicates the additional hypotheses Holm rejects but Bonferroni does not. In the middle panel, Holm rejects one more null than Bonferroni; in the right panel, it rejects five more [JWHT21, Figure 13.3].

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FWER vs. power trade-off

FWER demands *no* false rejections with probability at least $1 - \alpha$:

- Very stringent if *m* is large
- Tends to reduce power (fewer true positives found)

In modern "exploratory" studies:

- We may tolerate a small fraction of false positives to discover more true ones
- This leads to the false discovery rate (FDR) approach

	H_0 is true	H_0 is not true
Reject H_0 Not reject H_0	Type-I error (FP) Correct (TN)	Correct (TP) Type-II error (FN)

Illustration: power vs. FWER

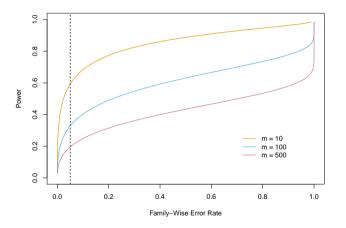


Figure: In a simulation with 90% of m nulls true, the power is displayed against FWER. Colors of the curves: m=10 (orange), m=100 (blue), m=500 (purple). Larger m reduces power. The vertical dashed line marks FWER=0.05 [JWHT21, Figure 13.5].

False discovery rate (FDR): Definition and motivation

Motivation: Controlling FWER can be too conservative for large m

Instead: control the fraction of rejected hypotheses that are *false positives*

$$FDP = \frac{\# \text{ false positives}}{\# \text{ total rejections}} = \frac{\# FP}{\# FP + \# TP}$$

ullet Controlling FDP is impossible because we never know which H_{0j} are true/false

False discovery rate $(FDR) = \mathbb{E}[FDP]$

- Allow up to fraction q of false positives on average among the "claimed discoveries"
- ullet The choice of q is context- and dataset-dependent (no gold standard like lpha=0.05)

Properties:

- Accept a small fraction of false positives, in exchange for more total discoveries
- Typically yields more rejections ("discoveries") than FWER-based methods

Controlling FDR: Benjamini-Hochberg procedure

Benamini-Hochberg procedure

- 1 Specify q, the level at which to control the FDR
- 2 Compute the *p*-values for the *m* null hypotheses, H_{01}, \ldots, H_{0m}
- 3 Sort p-values so that $p_{(1)} \leq p_{(2)} \leq \cdots \leq p_{(m)}$
- 4 Define

$$L = \max\left\{j: p_{(j)} < \frac{qj}{m}\right\}$$

5 Reject all null hypotheses H_{0j} for which $p_j \leq p_{(L)}$

Result:

- Ensures FDR $\leq q$, but but not necessarily small FWER
- Typically more powerful, yielding more rejections, than Bonferroni/Holm if *m* is large

Example: Benjamini-Hochberg procedure

Example

Step 1: Set
$$q = 0.05$$

Step 2:
$$p_1 = 0.0018$$
, $p_2 = 0.009$, $p_3 = 0.021$, $p_4 = 0.034$, $p_5 = 0.045$, $p_6 = 0.070$.

Step 3: Sort *p*-values
$$p_{(1)} = 0.0018$$
, $p_{(2)} = 0.009$, $p_{(3)} = 0.021$, $p_{(4)} = 0.034$, $p_{(5)} = 0.045$, $p_{(6)} = 0.070$.

Step 4: Find L = 3 because

$$k = 1: \quad 0.0018 \le 0.05 \times \frac{1}{6} \approx 0.0083? \checkmark$$
 $k = 2: \quad 0.009 \le 0.05 \times \frac{2}{6} \approx 0.0167? \checkmark$
 $k = 3: \quad 0.021 \le 0.05 \times \frac{3}{6} = 0.025? \checkmark$
 $k = 4: \quad 0.034 \le 0.05 \times \frac{4}{6} \approx 0.0333? \text{ No } (0.034 > 0.0333)$

Step 5: Reject $H_{(1)}, H_{(2)}, H_{(3)}$.

Conclusion: BH rejects 3, while Holm rejects 2, Bonferroni rejects 1.

Visual comparison: Bonferroni vs. Benjamini-Hochberg

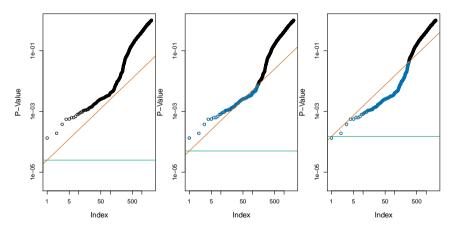


Figure: Panels: same set of m=2000 sorted p-values for the Fund dataset. Green lines: thresholds for FWER control (Bonferroni) at $\alpha=0.05,\,0.1,\,0.3$ (left to right). Orange lines: thresholds for FDR control (Benjamini-Hochberg) at $q=0.05,\,0.1,\,0.3$ (left to right). E.g., When the FDR is controlled at $q=0.1,\,146$ nulls are rejected (center, blue points). At $q=0.3,\,279$ nulls are rejected (right, blue points) [JWHT21, Figure 13.6].

Wrap-up

- **FWER** (Bonferroni/Holm):
 - Strictly ensures $Pr(any false positive) \le \alpha$
 - Conservative for large m, leading to fewer rejections & reduced power
- **FDR** (Benjamini–Hochberg):
 - Controls the expected fraction of false positives among rejections
 - ullet Typically yields more rejections than FWER, especially for large m
- Practical consideration:
 - Use FWER for strict confirmatory analyses needing minimal Type I error
 - Use FDR for exploratory, large-scale studies, tolerating some false positives to gain more discoveries

References



Gareth James, Daniela Witten, Trevor Hastie, and Robert Tibshirani.

An Introduction to Statistical Learning: with Applications in R, volume 112 of Springer Texts in Statistics.

Springer, New York, NY, 2nd edition, 2021.