

Controlling the False Discovery Rate (Benjamini & Hochberg, 1995)

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Presenting for Stats 319 (Journal Club)

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Acknowledgment: Our presentation is largely inspired by the 300C lectures by Prof. Emmanuel Candès

A review of the hypothesis testing framework

A Decision Problem

Data are sampled from some distribution parameterized by θ :

- $X \sim \mathbb{P}_\theta$
- $\theta \in \Omega$

Furthermore, the parameter space Ω can be split into disjoint subclasses known as "hypotheses":

$$H_0 : \theta \in \Omega_0 \subset \Omega \quad (\text{null hypothesis})$$

$$H_1 : \theta \in \Omega_1 = \Omega \setminus \Omega_0 \quad (\text{alternative hypothesis})$$

Our goal is to infer which hypothesis is correct.

The Neyman-Pearson Paradigm

	Reject H_0	Retain H_0
$\theta \in \Omega_0$	Type I error	Good
$\theta \in \Omega_1$	Good	Type II error

- **Level of significance:** A level- α test guarantees that $\mathbb{P}_{H_0}(\text{Type I error}) \leq \alpha$.
- **power** = $1 - \mathbb{P}_{H_1}(\text{Type II error})$

Under the Neyman-Pearson paradigm, a test procedure maximizes the power subject to the level of significance. **The only guarantee is a type I error rate less than α .**

Typically, θ is an unobservable state of the universe which interests us, and H_0 represents our default state of belief:

- H_0 : Male and female births are equally likely.
- H_0 : No difference in expected blood pressure after treatment.
- H_0 : The true regression coefficient β_1 is zero.

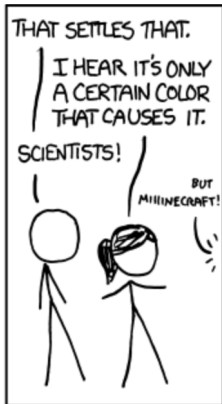
But what if we perform multiple tests?

- H_{0j} : No difference in expected blood pressure after treatment j .
- H_{0j} : The true j th regression coefficient β_j is zero.

Recall that we only control the type I error rate, typically at level $\alpha = 0.05$.

What does this mean for the state of science?

Testing multiple hypotheses



WE FOUND NO LINK BETWEEN PURPLE JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN BROWN JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN PINK JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN BLUE JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN TEAL JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN SALMON JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN RED JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN TURQUOISE JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN MAGENTA JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN YELLOW JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN GREY JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN TAN JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN CYAN JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND A LINK BETWEEN GREEN JELLY BEANS AND AONE ($P < 0.05$).



WE FOUND NO LINK BETWEEN MAUVE JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN BEIGE JELLY BEANS AND AONE ($P > 0.05$).



WE FOUND NO LINK BETWEEN LILAC JELLY BEANS AND AONE ($P > 0.05$).



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NEWS

GREEN JELLY BEANS LINKED TO ACNE!

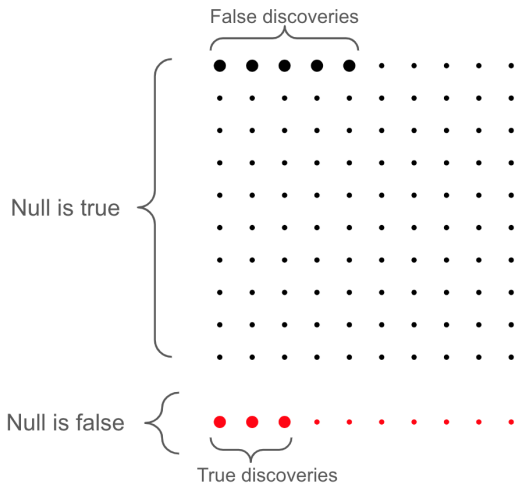
95% CONFIDENCE

ONLY 5% CHANCE OF COINCIDENCE!



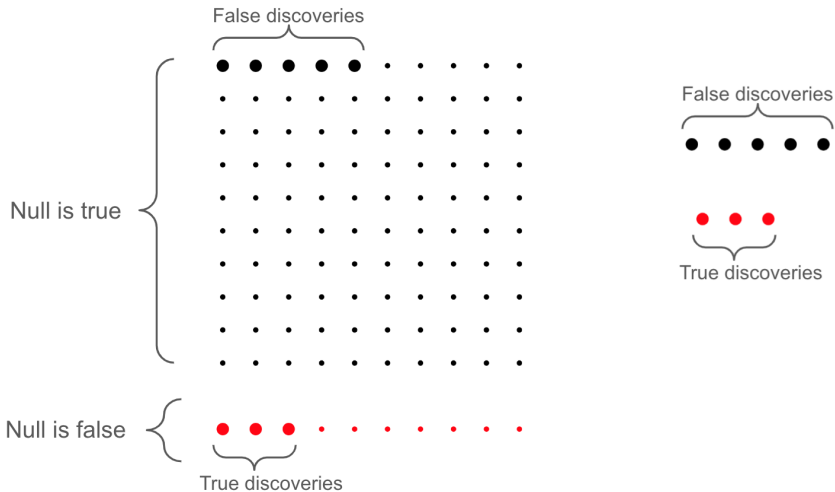
SCIENTISTS...

Traditional type-I error control

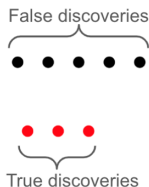


- Each dot represents a hypothesis being tested. A bold dot represents rejecting the null hypothesis (declaring a discovery).
- We imagine an army of scientists all around the world, all testing their own hypotheses.
- Type-I error control says: out of all the dots, all the hypotheses tested around the world, at most 5% are *bold black dots* (false discoveries).
- But if only the discoveries are published, we don't get to see all the dots!

The file-drawer effect: what do we actually see?



FDR lets us control the proportion of false discoveries out of all *discoveries*, not out of all *hypotheses tested*



FDR control puts an upper bound on

$$E \left(\frac{\text{false discoveries}}{\text{false discoveries} + \text{true discoveries}} \right)$$

Formal introduction to FDR control

	declared non-signif.	declared significant	Total
H_0 true	U	V	n_0
H_0 false	T	S	$n - n_0$
	$n - R$	R	n

- Familywise error rate (FWER) = $\mathbb{P}(V \geq 1)$
- False discovery proportion (FDP):

$$\text{FDP} = \frac{V}{\max(R, 1)} = \begin{cases} V/R & \text{if } R \geq 1 \\ 0 & \text{if } R = 0 \end{cases}$$

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

- If all the hypotheses are true, then

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Control FDR instead of controlling FWER?

FWER vs. FDR (contd.)

- Small # hypotheses \rightarrow FWER control \checkmark (but, may lack power)
- Large-scale studies \rightarrow FWER control may miss important findings

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- Small # hypotheses \rightarrow FWER control \checkmark (but, may lack power)
- Large-scale studies \rightarrow FWER control may miss important findings
- FDR control sacrifices some stringency to permit exploration with a few false positives
- FDR control does not assure a specific study, but ensures that science as a whole will be alright!

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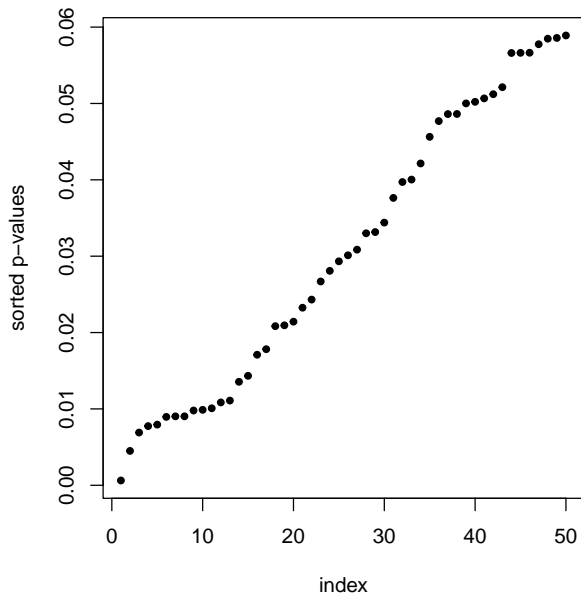
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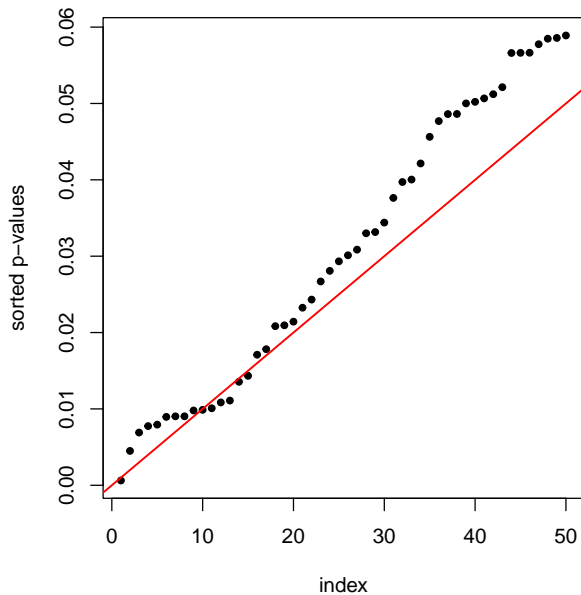
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- Compute p -values p_1, \dots, p_n for the n hypotheses H_1, \dots, H_n
- Sort the p -values: $p_{(1)} \leq p_{(2)} \leq \dots \leq p_{(n)}$
- BH_α procedure: Reject $H_{(1)}, \dots, H_{(i_0)}$ where

$$i_0 = \max\{i : p_{(i)} \leq i\alpha/n\}$$

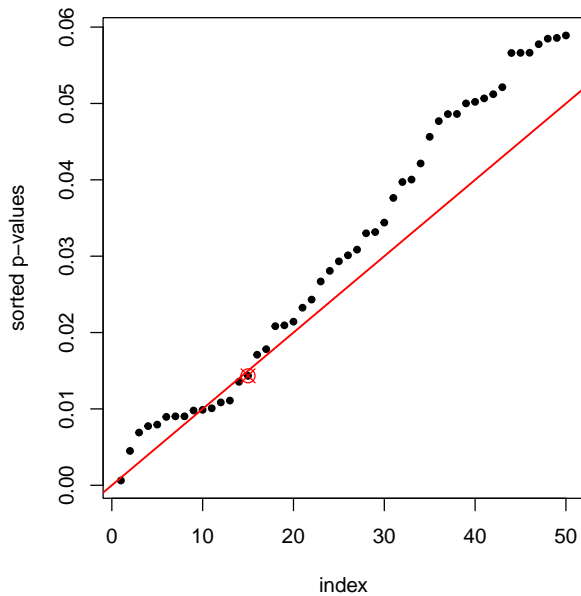
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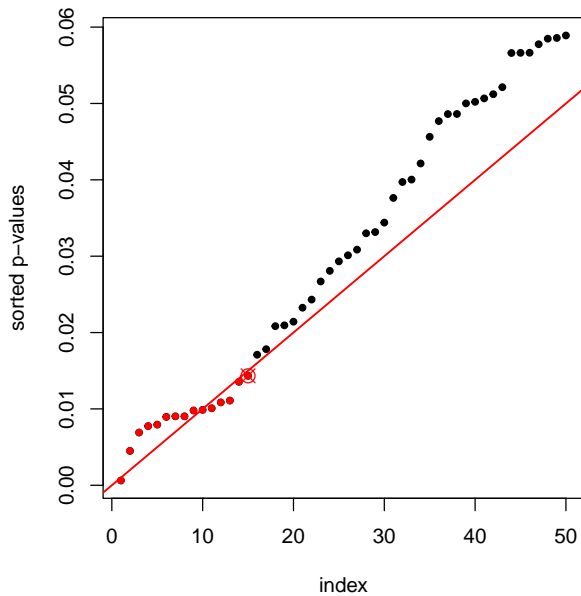
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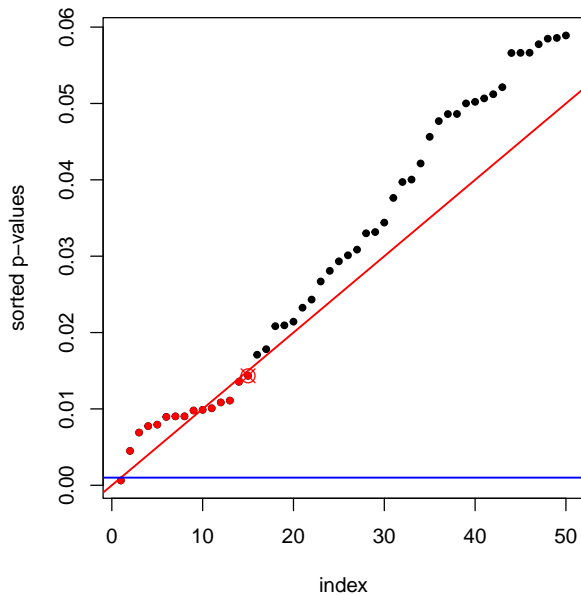
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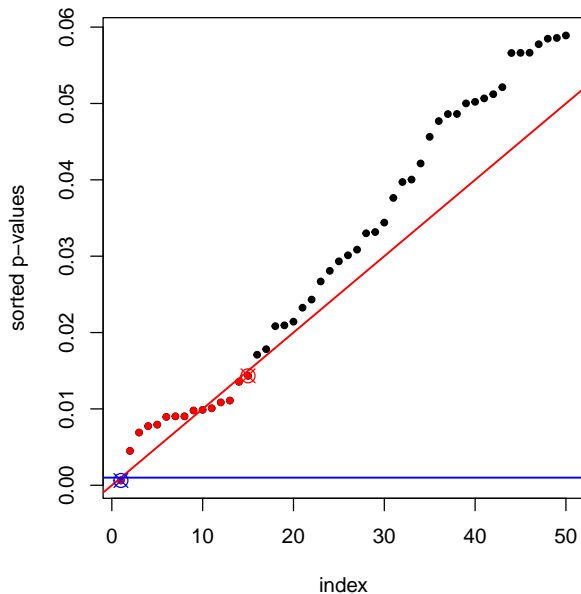
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- Hochberg (1988) gives a procedure for strong FWER control

$$i_0 = \max \left\{ i : p_{(i)} \leq \frac{\alpha}{n+1-i} \right\} \quad \text{vs} \quad i_0 = \max \left\{ i : p_{(i)} \leq \frac{i\alpha}{n} \right\}$$

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- BH argue how their procedure rejects more than the above one

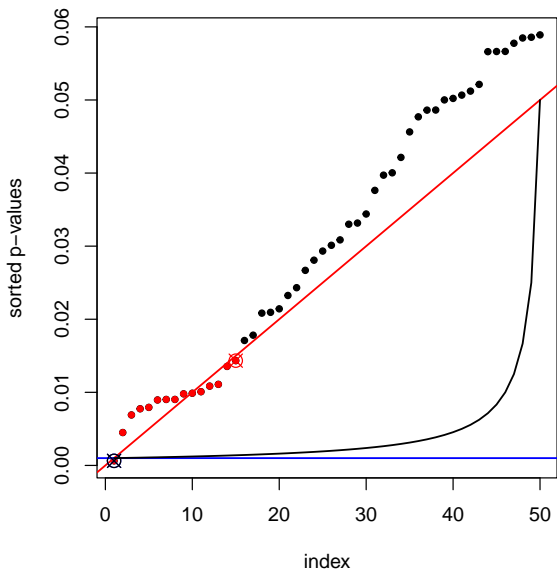


Figure 1: FDR control makes more rejections (and has more power) than FWER control

Theoretical guarantees

Theorem (Benjamini & Hochberg, 1995). The BH_α procedure controls the FDR at level α if the p-values are independent:

$$\text{FDR} = \frac{n_0}{n} \alpha \leq \alpha.$$

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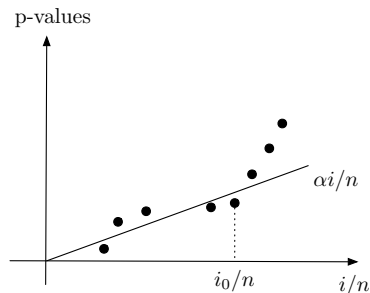
Numerous proofs, see our Stats 300C lecture notes for a couple of them

BH has more power

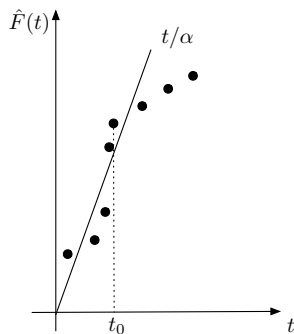
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(a) p -values on the y axis, indices on x



(b) p -values on the x axis, indices on y

Proof of FDR control by Martingale theory (Storey et al., 2004)

Consider rejecting all H_i with p-values $p_i \leq t$, where $t \in (0, 1)$

	H_0 not rejected	H_0 rejected	Total
H_0 true	$U(t)$	$V(t)$	n_0
H_0 false	$T(t)$	$S(t)$	$n - n_0$
	$n - R(t)$	$R(t)$	n

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$V(t)/t$ is a backwards martingale $\mathbb{E}\left[\frac{V(s)}{s} \mid \mathcal{F}_{\geq t}\right] = \frac{1}{s} \frac{s}{t} V(t)$ for $s \leq t$

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$$\text{FDR}(\tau) = \mathbb{E} \left[\frac{V(\tau)}{R(\tau) \vee 1} \right] \stackrel{\text{pic}}{\leq} \frac{\alpha}{n} \mathbb{E} \left[\frac{V(\tau)}{\tau} \right] \stackrel{\text{OST}}{=} \frac{\alpha}{n} \mathbb{E} \left[\frac{V(1)}{1} \right] \stackrel{\text{def}}{=} \alpha \frac{n_0}{n} \leq \alpha$$

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Storey's procedure improves upon BH, by doing better than $\frac{n_0}{n} \leq 1$

Theorem (Benjamini & Yekutieli, 2001). Under arbitrary dependence of the p-values, the BH_α procedure has the following guarantee

$$\text{FDR} = \frac{n_0}{n} \alpha H(n) \leq \alpha H(n)$$

where $H(n) = 1 + \frac{1}{2} + \cdots + \frac{1}{n} \approx \log n + 0.577$.

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Theorem (Guo & Rao, 2008). There are joint distributions of p-values for which FDR of the BH procedure is at least $\min\{\alpha H(n), 1\}$.

The e-BH procedure (Wang & Ramdas, 2020)

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- **e-BH** procedure: apply BH to a bunch of (e-values)⁻¹
- **Theorem** (Wang & Ramdas, 2020). The e-BH procedure has FDR at most $\alpha n_0/n \leq \alpha$ (same guarantee as for the usual BH procedure with independent p -values)

- Traditional type-I error control fails when you test multiple hypotheses but suppress null findings.
- FDR is a *statistical* fix. But we also need *sociological* or *cultural* fixes: change the incentives in science so we can see more of the null findings.
 - Preregistration
 - Journals for null results
 - Evaluation criteria for job candidates, tenure, prestigious awards: do we value shocking results, or careful study design?



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Journal of Negative Results in Biomedicine

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Thank You!

Questions?