## **FWER**



	True H <sub>i</sub> s	False H <sub>i</sub> s	Total
Accepted $H_i$ s	U	T	M-R
Rejected $H_i$ s	V	S	R
Total	$m_0$	$m-m_0$	m

### Familywise (type I) error rate:

$$FWER = P(V > 0)$$

Controlling FWER at level  $\alpha$ :

$$FWER = P(V > 0) \le \alpha$$

How?

## **FWER**

#### Familywise (type I) error rate:

$$FWER = P(V > 0)$$

introlling FWFR at level  $\alpha$ :

$$FWER = P(V > 0) \le \alpha$$

How?

 $\alpha_1, \dots, \alpha_m$  – significance levels for  $H_1, \dots, H_m$ 

We have to select them to ensure FWER  $\leq \alpha$ .

## **Bonferroni correction**

#### **Bonferroni method:**

$$\alpha_1 = \dots = \alpha_m = \frac{\alpha}{m}$$

Comparing  $\alpha_i$  and  $p_i$  is the same as comparing original  $\alpha$  and adjusted p-value

$$\tilde{p}_i = \min(1, mp_i)$$

 $H_i$  is rejected when  $\tilde{p}_i \leq \alpha$ .

### **Bonferroni** correction

**Theorem**. If  $H_i$  is rejected when  $p_i \leq \alpha/m$ , then FWER  $\leq \alpha$ .

Proof.

$$\begin{aligned} \text{FWER} &= P(V > 0) = P\left(\bigcup_{i \in M_0} \left\{ p_i \le \frac{\alpha}{m} \right\} \right) \le \\ &\le \sum_{i \in M_0} P\left( p_i \le \frac{\alpha}{m} \right) \le \\ &\le \sum_{i \in M_0} \frac{\alpha}{m} = \frac{m_0}{m} \alpha \le \alpha \end{aligned}$$

50 samples from N(1,1), 150 samples from N(0,1), n=20  $H_i$ :  $\mathbb{E}X_i = 0$ ,  $H'_i$ :  $\mathbb{E}X_i \neq 0$ , one sample t-test

#### No corrections:

	True $H_i$ s	False H <sub>i</sub> s	Total
Accepted $H_i$ s	143	0	143
Rejected <i>H</i> <sub>i</sub> s	7	50	57
Total	150	50	200

#### Bonferroni correction:

	True $H_i$ s	False $H_i$ s	Total
Accepted $H_i$ s	150	19	169
Rejected <i>H</i> <sub>i</sub> s	0	31	31
Total	150	50	200

### Can we do better?

Bonferroni method:

$$\alpha_1 = \dots = \alpha_m = \frac{\alpha}{m}$$

A more powerful method is possible if we allow  $\alpha_i$ s to vary.

## Step-down methods

Sorted p-values:

$$p_{(1)} \leq \cdots \leq p_{(m)}$$

 $H_{(1)}, \dots, H_{(m)}$  – corresponding hypotheses

#### Step-down procedure:

- 1. If  $p_{(1)} > \alpha_1$ , accept  $H_{(1)}, \dots, H_{(m)}$  and stop; otherwise reject  $H_{(1)}$  and continue
- 2. If  $p_{(2)} > \alpha_2$ , accept  $H_{(2)}, \dots, H_{(m)}$  and stop; otherwise reject  $H_{(2)}$  and continue
- 3. ...

## Holm's method

Holm's method – a step-down procedure with

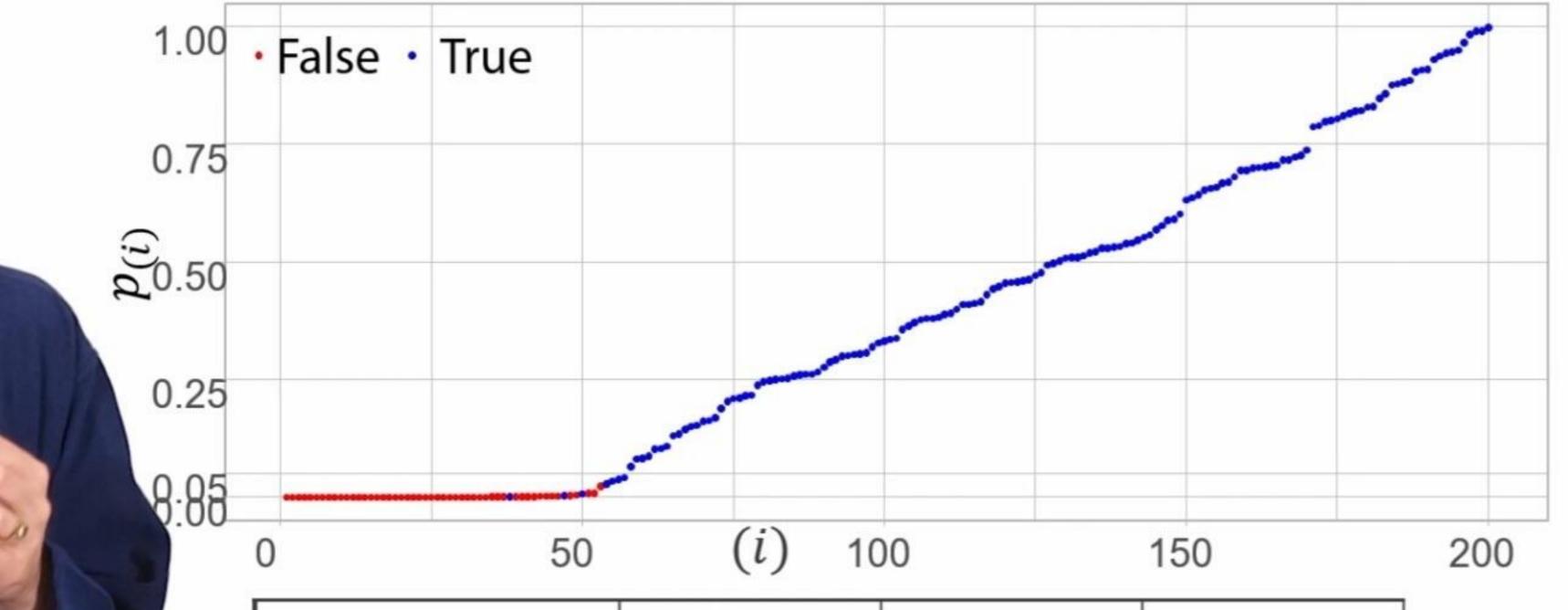
$$\alpha_1 = \frac{\alpha}{m}, \alpha_2 = \frac{\alpha}{m-1}, \dots, \alpha_i = \frac{\alpha}{m-i+1}, \dots, \alpha_m = \alpha$$

Adjusted p-values:

$$\tilde{p}_{(i)} = \min\left(1, \max\left((m-i+1)p_{(i)}, \tilde{p}_{(i-1)}\right)\right)$$

• FWER  $\leq \alpha$  is guaranteed

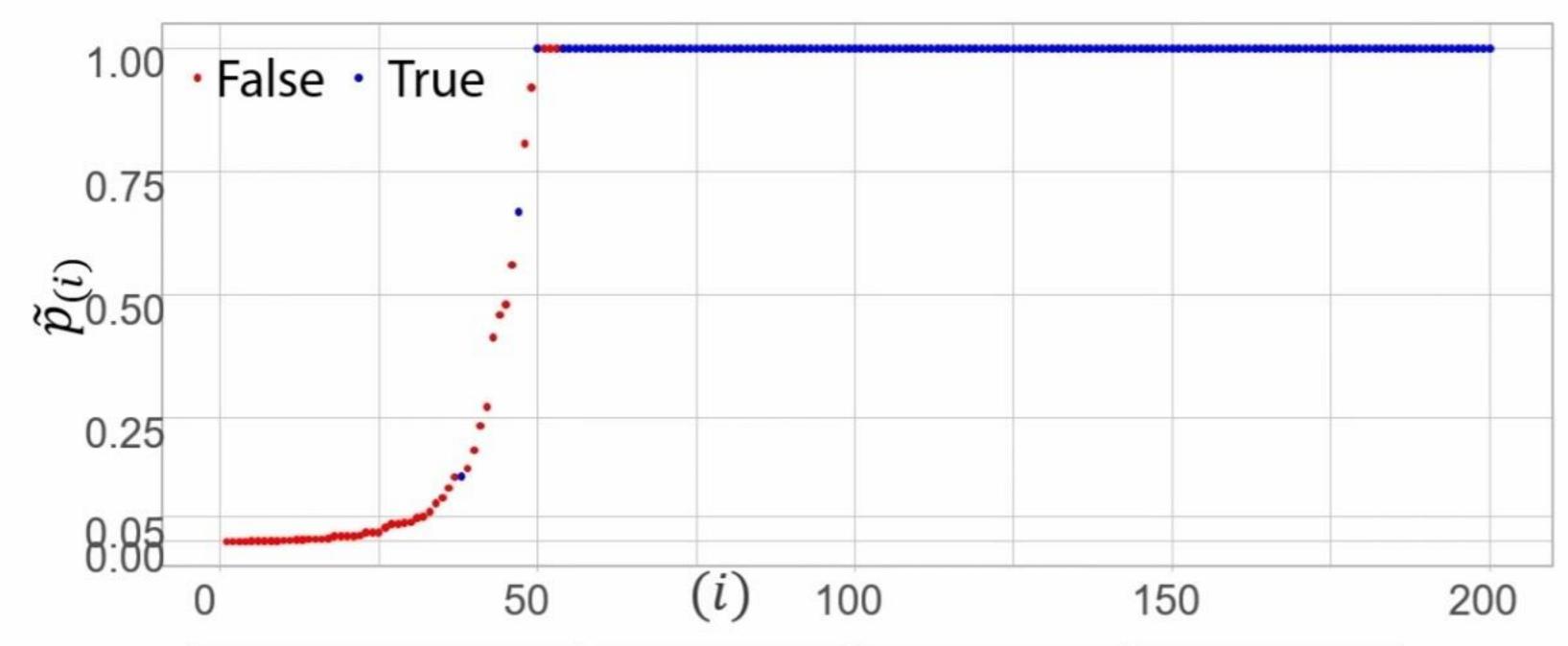
No corrections:



	True $H_i$ s	False H <sub>i</sub> s	Total
Accepted $H_i$ s	143	0	143
Rejected <i>H</i> <sub>i</sub> s	7	50	57
Total	150	50	200



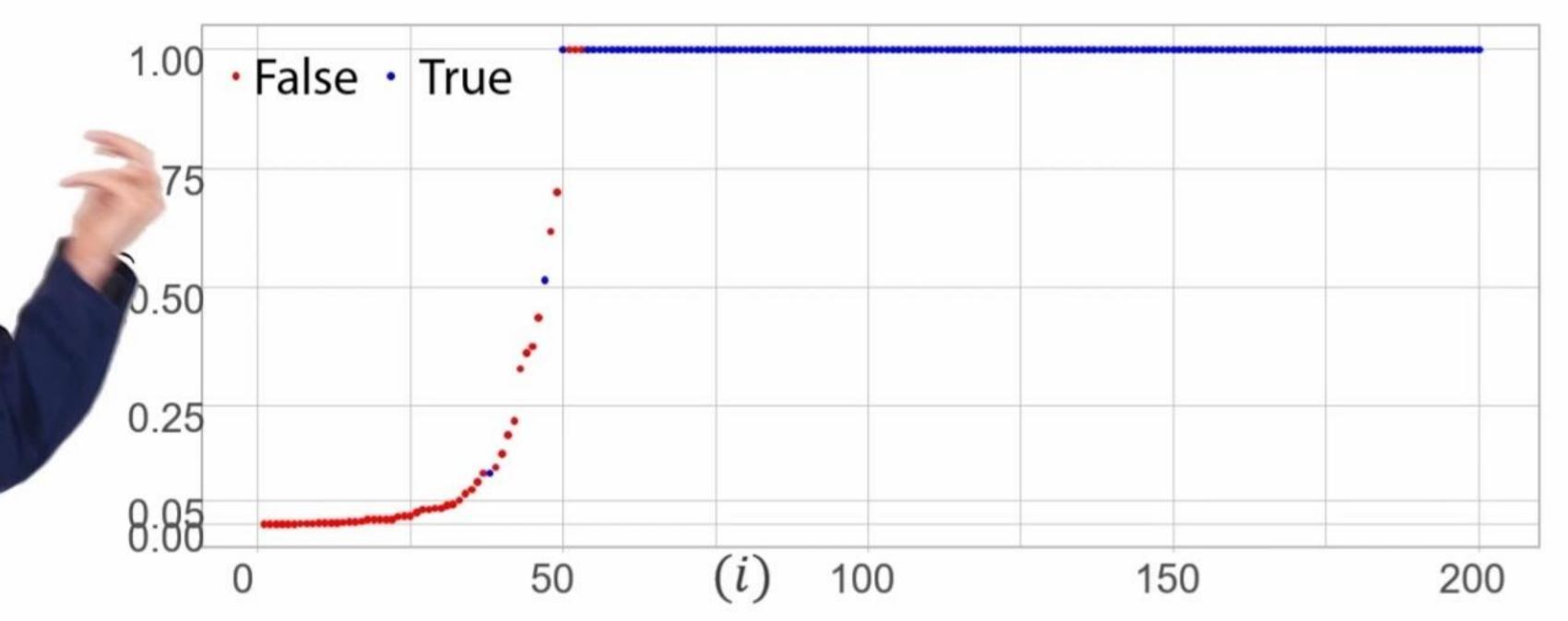
Bonferroni correction:



	True H <sub>i</sub> s	False H <sub>i</sub> s	Total
Accepted $H_i$ s	150	19	169
Rejected <i>H</i> <sub>i</sub> s	0	31	31
Total	150	50	200



Holm's method:



	True H <sub>i</sub> s	False H <sub>i</sub> s	Total
Accepted $H_i$ s	150	18	168
Rejected <i>H</i> <sub>i</sub> s	0	32	32
Total	150	50	200



#### Bonferroni correction:

	True H <sub>i</sub> s	False H <sub>i</sub> s	Total
Accepted $H_i$ s	150	19	169
Rejected <i>H</i> <sub>i</sub> s	0	31	31
Total	150	50	200

#### Holm's method:

	True <i>H</i> <sub>i</sub> s	False H <sub>i</sub> s	Total
Accepted $H_i$ s	150	18	168
Rejected $H_i$ s	0	32	32
Total	150	50	200

# **Takeaways about FWER**

Control FWER if it's very important not to make ANY type I error

 Use Holm's method instead of Bonferroni to reject more hypotheses FOR FREE