

Hypothesis testing. $\Theta_0 \cap \Theta_1 = \emptyset$.

Decide: $H_0: \theta \in \Theta_0$ v.s. $H_1: \theta \in \Theta_1$.

based on observed data.

Usually, we compute some test statistics $T(X) \in \mathbb{R}$
and reject when $T(X) > c$

for some pre-specified (determinate) $c \in \mathbb{R}$.
"critical value".

Type - I error: $\theta \in \Theta_0$, but reject

Type - II error: $\theta \in \Theta_1$, but cannot reject.

Power function $\beta(\theta) = P_{\theta}(\text{reject}) \quad (\forall \theta \in \Theta)$.

size / level of the test

$$\alpha = \sup_{\theta \in \Theta_0} \beta(\theta).$$

"worse-case
type-I error"

Goal of testing

$$\text{maximize } \left\{ \beta(\theta) \text{ for } \theta \in \Theta_1 \right\}$$

Still a multi-objective optimization problem.

s.t. $\alpha = \sup_{\theta \in \Theta_0} \beta(\theta) \leq \alpha_0$ (e.g. 5%).

A valid test: type-I error constraint is satisfied.

(Asymptotic version: $X_1, X_2, \dots, X_n \stackrel{iid}{\sim} P_\theta$.

$$\text{level} = \sup_{\theta \in \Theta_0} \lim_{n \rightarrow \infty} P_{\theta}(\text{reject})$$

Concrete instantiation:

1-dim problem:

. Two-sided testing:

$$H_0: \theta = \theta_0 \text{ v.s. } H_1: \theta \neq \theta_0$$

(need to ensure $P_{\theta_0}(\text{reject}) \leq \alpha$).

. One-sided testing

$$H_0: \theta \leq \theta_0 \text{ v.s. } H_1: \theta > \theta_0$$

(or the other way around)

Need to ensure $\forall \theta \leq \theta_0$, $P_\theta(\text{reject}) \leq \alpha$

e.g. $X_1, X_2, \dots, X_n \stackrel{\text{iid}}{\sim} N(\theta, 1)$

$$T(X) = \frac{1}{n} \sum_{i=1}^n X_i$$

- One-sided testing: $H_0: \theta \leq \theta_0$, $H_1: \theta > \theta_0$.

Reject when $T(X) > c$

under θ , $T(X) \sim N\left(\theta, \frac{1}{n}\right)$

$$P_\theta(\text{reject}) = 1 - \Phi\left(\sqrt{n}(c - \theta)\right)$$

(where Φ is cdf of $N(0, 1)$).

So we choose $c = \theta_0 + \frac{1}{\sqrt{n}} Z_{1-\alpha}$

(($1-\alpha$) quantile of $N(0, 1)$).

Similarly, for two-sided tests.

reject when $T(X) > c_2$
or
 $T(X) < c_1$

$$P_{\theta_0}(\text{rejection}) = 1 - \Phi\left(\sqrt{n}(C_2 - \theta_0)\right) + \Phi\left(\sqrt{n}(\theta_0 - C_1)\right)$$

Need to choose (C_1, C_2) pair s.t. $P_{\theta_0}(\text{rejection}) \leq \alpha$.

Commonly used choice:

$$\begin{cases} C_2 = \theta_0 + \frac{1}{\sqrt{n}} Z_{1-\alpha/2} \\ C_1 = \theta_0 - \frac{1}{\sqrt{n}} Z_{1-\alpha/2} \end{cases}$$

"unbiased test"

most powerful
among unbiased
tests.

For any $r \in (0, 1)$,

$$\begin{cases} C_2 = \theta_0 + \frac{1}{\sqrt{n}} Z_{1-r\alpha} \\ C_1 = \theta_0 - \frac{1}{\sqrt{n}} Z_{1-(1-r)\alpha} \end{cases}$$

is also a valid choice.

Wald test:

$H_0: \theta = \theta_0$ vs. $H_1: \theta \neq \theta_0$.

Suppose that we have estimator $\hat{\theta}_n$

s.t. under P_{θ_0} , we have $\frac{\hat{\theta}_n - \theta_0}{\widehat{\text{se}}} \xrightarrow{d} N(0, 1)$

then we reject when

$$\left| \frac{\hat{\theta}_n - \theta_0}{\widehat{se}} \right| \geq Z_{1-\alpha}$$

By def of " \xrightarrow{d} ", we have

$$\lim_{n \rightarrow \infty} P_{\theta_0}(\text{rejection}) = \alpha.$$

Similarly, we have one-sided test:

$$H_0: \theta \leq \theta_0, \quad H_1: \theta > \theta_0$$

reject when $\frac{\hat{\theta}_n - \theta_0}{\widehat{se}} > Z_{1-\alpha}$.

χ^2 tests.

Suppose that $\theta \in \mathbb{R}^d$

$$H_0: \theta = \theta_0 \quad \text{vs.} \quad H_1: \theta \neq \theta_0.$$

Estimator $\hat{\theta}_n: (\sum)^{-\frac{1}{2}} (\hat{\theta}_n - \theta_0) \xrightarrow{d} N(0, \text{Id})$

($\hat{\Sigma}$ computed from data, estimator for asympt cov).

If $\hat{\beta} \sim N(0, \text{Id})$, then $\|\hat{\beta}\|_2^2 \sim \chi_d^2$

χ^2 test: reject when

$$\left\| \hat{\Sigma}_n^{1/2} (\hat{\theta}_n - \theta_0) \right\|_2^2 \geq (1-\alpha) \text{ quantile of } \chi_d^2.$$

Remark: $\hat{s}\hat{e}$ and $\hat{\Sigma}$ can also be replaced by exact expressions of asympt. var. under P_{θ_0} .

The results will still be valid.

(we use $\hat{s}\hat{e}$ and $\hat{\Sigma}$ for computational reasons)

Special case:

$$X \sim \text{Multinomial}(n, p) \quad \text{for } p \in \mathbb{R}^k$$
$$p_1, p_2, \dots, p_k > 0$$
$$\sum_{j=1}^k p_j = 1$$

Want to test

$$H_0: p = p_0 \quad \text{Under } P_0.$$

$$H_1: p \neq p_0. \quad \sqrt{n} \left(\frac{X}{n} - p_0 \right) \xrightarrow{d} \text{degenerate normal.}$$

Pearson's χ^2 test.

under H_0

$$T = \sum_{j=1}^k \frac{(X_j - np_{0j})^2}{np_{0j}} \xrightarrow{d} \chi_{k-1}^2$$

Student t-test.

$$X_1, X_2, \dots, X_n \stackrel{iid}{\sim} N(\mu, \sigma^2)$$

μ, σ^2 are unknown, we want to test

$$H_0: \mu = \mu_0, \text{ vs. } H_1: \mu \neq \mu_0.$$

$$T = \frac{\sqrt{n}(\bar{X}_n - \mu_0)}{\hat{s}\epsilon_n} \quad (\xrightarrow{d} N(0, 1), \text{ as } n \rightarrow \infty)$$

"t-distribution w/ def $n-1$ "

:= distribution of

$$\frac{\frac{1}{\sqrt{n}} \sum_{i=1}^n \xi_i}{\hat{s}\epsilon(\xi_1, \dots, \xi_n)}$$

where $\xi_1, \xi_2, \dots, \xi_n \stackrel{iid}{\sim} N(0, 1)$.

reject based on quantile of t-distribution
 \Rightarrow exact level- α test.

p-values.

Defn. Suppose $T(X)$ is the test statistic.

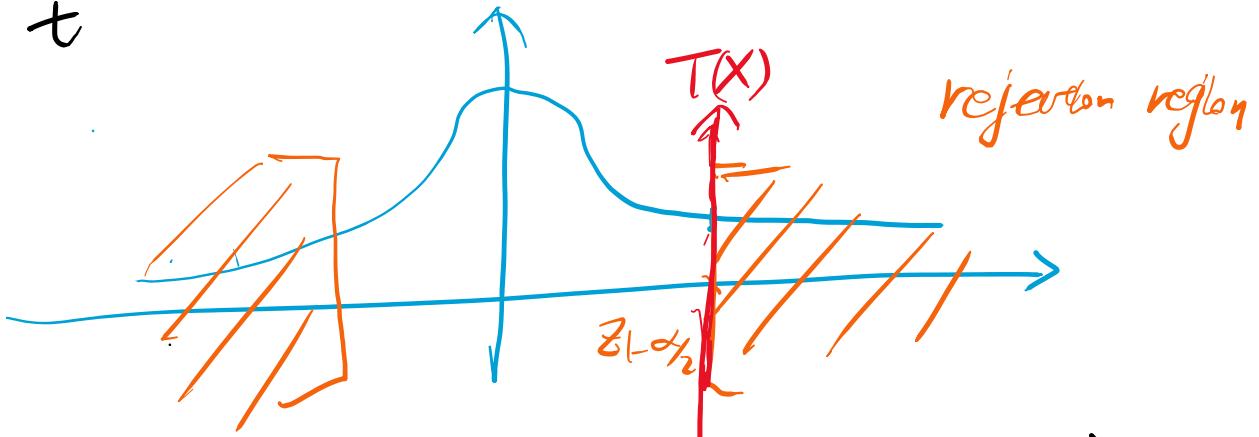
Rejection region R_α (reject when $T(X) \in R_\alpha$)

$$p\text{-value} := \inf \{\alpha : T(X) \in R_\alpha\}.$$

(read the warnings from textbook)

Test becomes: reject when p-value $\leq \alpha$.

e.g. $T(X) = \frac{\hat{\theta} - \theta_0}{\hat{s}_\theta}$ $R_\alpha = \{t \in \mathbb{R} : |t| > z_{1-\alpha/2}\}$.



$$p\text{-value}(t) = P_{\theta_0}(|T(X)| \geq |t|) = 2 \Phi(-|t|)$$

So the p-value is given by $2 \Phi(-|T(X)|)$.

(in other words, we can solve for p-value by

$$Z_{1-\alpha/2} = T(X)$$

Fact: If $T(X)$ has a cts distribution under P_0 ,
then the p-value $\sim \text{Unif}(0, 1)$ under P_0 .

(This is a direct corollary of the fact:

(If X has cts distribution, F is its cdf
then $F(X) \sim \text{Unif}(0, 1)$).

Multiple testing.

A collection of testing problems

H_0 : vs. $H_{1,i}$ for $i=1, 2, \dots, m$.

p_1, p_2, \dots, p_m are p-values computed

for each testing problem

Naive idea: reject p_j when $p_j \leq \alpha$
for each $j \in \{1, 2, \dots, m\}$.

e.g. If p_j 's are all independent,
and test stats have cts distributions.

$$P_{H_0}(\text{make any false discovery}) = (1 - \alpha)^m$$

$$\text{when } m \approx \frac{1}{\alpha}, P(\dots) \approx 1 - \frac{1}{e}$$

Bonferroni correction.

Idea: (no matter what dependence structures.)

$$P_{H_0}(\text{false discovery}) \leq \sum_{j=1}^m P_{H_0}(\text{reject } H_0j) \leq \alpha$$

We choose to test each H_0j at level $\frac{\alpha}{m}$.

Relaxation : False Discovery Rate (FDR).

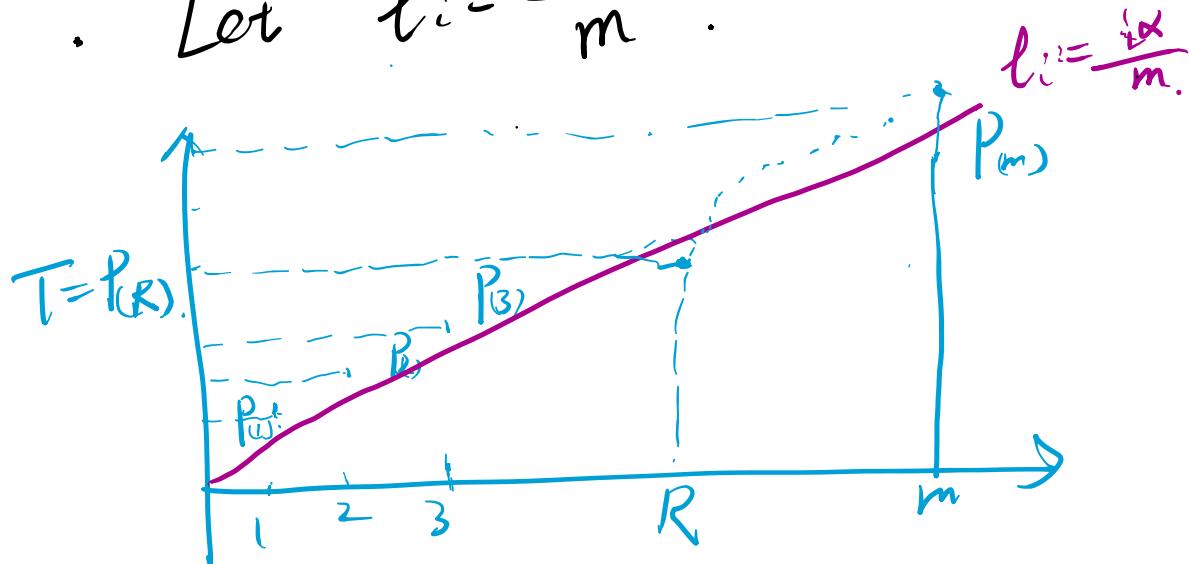
False discovery proportion.

$$\text{FDP} := \begin{cases} \frac{\# \text{ false rejections}}{\# \text{ rejections.}} & (\# \text{ rej} > 0) \\ 0 & (\# \text{ rej} = 0) \end{cases}$$

$$\text{FDR} = \mathbb{E}[\text{FDP}].$$

Benjamini-Hochberg (BH) procedure:

- $P_{(1)} < P_{(2)} < \dots < P_{(m)}$ (sorted p-values)
- Let $t_i = \frac{i\alpha}{m}$.



- $R := \max\{i : P_{(i)} \leq t_i\}$

we let $T = P_{(R)}$.

and we reject when $P_i \leq T$

(i.e. we reject the R smallest p-values)

Permutation tests.

Suppose $X_1, X_2, \dots, X_n \stackrel{iid}{\sim} P_X$

$Y_1, Y_2, \dots, Y_m \stackrel{iid}{\sim} P_Y$

$$H_0: P_X = P_Y$$

$$H_1: P_X \neq P_Y$$

Ideas under H_0 , $(n+m)$ samples can be mixed together, permutation does not make much difference

Given a test statistic

$$T(X_1, X_2, \dots, X_n, Y_1, Y_2, \dots, Y_m)$$

(e.g. $\left| \frac{1}{n} \sum_i^n X_i - \frac{1}{m} \sum_j^m Y_j \right|$,

$$\sup_{g \in G} \left| \frac{1}{n} \sum_i^n g(X_i) - \frac{1}{m} \sum_j^m g(Y_j) \right|$$

Let $N = n+m$, consider all possible

$N!$ permutations of $(X_1, \dots, X_n, Y_1, \dots, Y_m)$.

(for notation convenience, we let $X_{n+i} = Y_i$
 $i=1, \dots, m$)

For each permutation σ

Let

$T_\sigma :=$ test statistic computed on
 $X_{\sigma(1)}, X_{\sigma(2)}, \dots, X_{\sigma(k)}, X_{\sigma(k+1)} \dots X_{\sigma(n+m)}$
as X 's as Y 's.

p-value := $\frac{1}{N!} \sum_{\sigma} \{ \text{If } T_\sigma > T(X; Y) \}$.

(Percentile of observed $T(X; Y)$ among $N!$
hypothetical worlds)

Monte-Carlo simulation: (similar to bootstrap).

For $b = 1, 2, \dots, B$,

sample σ_b uniformly among permutations.

Compare T_{σ_b} .

p-value $\approx \frac{1}{B} \sum_{b=1}^B \{ \text{If } T_{\sigma_b} > T(X; Y) \}$.