

Global tests in high-dimensions

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High-dimensional statistics

Classical theory

- It concerns the behaviour when the *sample size* $n \rightarrow \infty$
- Suppose $Y_1, \dots, Y_n \stackrel{i.i.d.}{\sim} Y$ with mean $\mu = \mathbb{E}(Y)$ and finite variance $\Sigma = \mathbb{V}\text{ar}(Y)$
- *Law of large numbers*: the sample mean $\hat{\mu}_n = \frac{1}{n} \sum_{i=1}^n Y_i$ converges in probability to μ
- *Central limit theorem*: the rescaled deviation $\sqrt{n}(\hat{\mu}_n - \mu)$ converges in distribution to a centered Gaussian with covariance matrix Σ
- *Consistency of maximum likelihood estimation*
- Etc.

Suppose that we are given $n = 1000$ samples from a statistical model in $m = 500$ dimensions

Will theory that requires $n \rightarrow \infty$ with the dimension m remaining fixed provide useful predictions?

High-dimensional data

- The data sets arising in many parts of modern science have a “high-dimensional flavor”, with m on the same order as, or possibly larger than n

$$m \gg n$$

- Classical “large n , fixed m ” theory fails to provide useful predictions
- Classical methods can break down dramatically in high-dimensional regimes

Reference

Wainwright (2019)

High-Dimensional Statistics: A Non-Asymptotic Viewpoint

Cambridge University Press

Sections 1.1.1, 1.2.1, 1.3.1

Linear discriminant analysis in high-dimensions

Two classes

- Hypothesis testing can be used to determine whether an observed vector $x = (x_1, \dots, x_m)^T \in \mathbb{R}^m$ has been drawn from one of two possible densities $f_A \equiv P(X|Y = A)$ and $f_B \equiv P(X|Y = B)$, corresponding to two possible classes A and B
- Consider testing $H_A : X_A \sim f_A$ vs $H_B : X_B \sim f_B$, where $X_A \equiv (X|Y = A)$ and $X_B \equiv (X|Y = B)$
- When these two distributions are known, then the Neyman-Pearson lemma says that the optimal decision rule is based on thresholding the log-likelihood ratio

$$\log \frac{f_B(x)}{f_A(x)}$$

- By testing H_A vs H_B and H_B vs H_A the conclusion is that the observed data x is consistent with A (H_B rejected), with B (H_A rejected), with both (no rejections), or with neither (both rejected)

Classification problem

- Let's turn to the classification problem involving the allocation of the observed unit x to one of two classes A and B
- For a Bayesian analysis suppose that the prior probabilities are $\pi_A \equiv P(Y = A)$ and $\pi_B \equiv P(Y = B)$ with $\pi_A + \pi_B = 1$. Then the posterior probabilities satisfy

$$\frac{P(Y = B|X = x)}{P(Y = A|X = x)} = \frac{\pi_B f_B(x)}{\pi_A f_A(x)}$$

giving the class with the higher posterior probability

- As a special case, suppose that the two classes are distributed as multivariate Gaussians $X_A \sim N(\mu_A, I_m)$ and $X_B \sim N(\mu_B, I_m)$, with $\pi_A = \pi_B = 1/2$

Optimal decision

- The optimal decision rule is to threshold the log-likelihood ratio

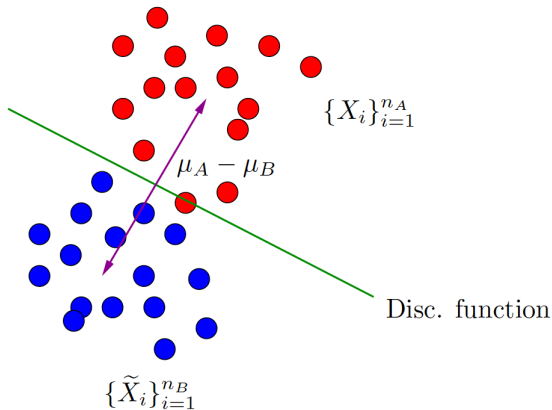
$$\Psi(x) = \langle \mu_A - \mu_B, \left(x - \frac{\mu_A + \mu_B}{2} \right) \rangle$$

where $\langle x, z \rangle = x^\top z = \sum_{j=1}^m x_j z_j$ denotes the Euclidean inner product in \mathbb{R}^m

- If $\Psi(x) > 0$ then classify A , otherwise B
- Error probability of the optimal rule:

$$\text{Err}(\Psi) = \frac{1}{2} \text{P}(\Psi(X_A) < 0) + \frac{1}{2} \text{P}(\Psi(X_B) \geq 0) = \Phi\left(-\frac{\gamma}{2}\right)$$

where $\gamma = \|\mu_A - \mu_B\|_2$, $\|\mu\|_2 = \sqrt{\mu^\top \mu}$, and Φ is the cdf of a standard normal variable



$$\langle \mu_A - \mu_B, \left(x - \frac{\mu_A + \mu_B}{2} \right) \rangle = 0$$

source: Wainwright

Linear Discriminant Analysis

- Fisher's LDA: uses the plug-in principle based on n_A samples from class A and n_B samples from class B

$$\hat{\Psi}(x) = \langle \hat{\mu}_A - \hat{\mu}_B, x - \frac{\hat{\mu}_A + \hat{\mu}_B}{2} \rangle$$

- Error probability of LDA (is itself a random variable)

$$\text{Err}(\hat{\Psi}) = \frac{1}{2}P(\hat{\Psi}(X_A) < 0) + \frac{1}{2}P(\hat{\Psi}(X_B) \geq 0)$$

- Classical theory: if $(n_A, n_B) \rightarrow \infty$ and m remains fixed, then $\hat{\mu}_A \xrightarrow{prob.} \mu_A$, $\hat{\mu}_B \xrightarrow{prob.} \mu_B$ and the asymptotic error probability is $\text{Err}(\hat{\Psi}) \xrightarrow{prob.} \text{Err}(\Psi) = \Phi(-\gamma/2)$

High-Dimensional Theory

- What happens if $(n_A, n_B, m) \rightarrow \infty$ with
 - $m/n_A \rightarrow \delta$ with $\delta \geq 0$
 - $m/n_B \rightarrow \delta$
 - $\|\mu_A - \mu_B\|_2 \rightarrow \gamma > 0$
- Kolmogorov (1960) showed that

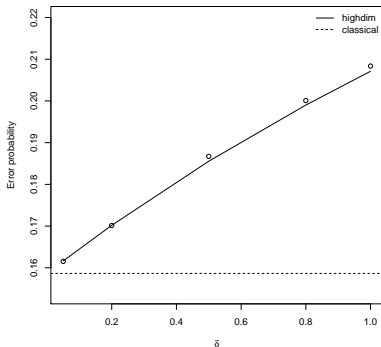
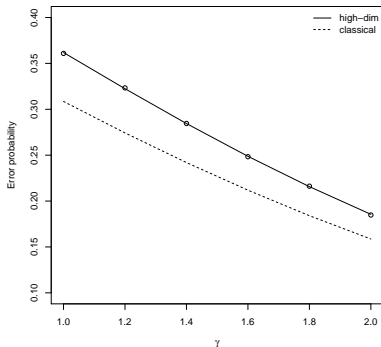
$$\text{Err}(\hat{\Psi}) \xrightarrow{\text{prob.}} \Phi\left(-\frac{\gamma^2}{2\sqrt{\gamma^2 + 2\delta}}\right)$$

- If $m/n \rightarrow 0$, then the asymptotic error probability is $\Phi(-\gamma/2)$ as is predicted by classical theory
- If $m/n \rightarrow \delta > 0$, then the asymptotic error probability is strictly larger than $\Phi(-\gamma/2)$

The error probability of $\hat{\Phi}$, for the finite triple

$$(m, n_A, n_B) = (400, 800, 800)$$

is better described by the classical $\Phi(-\gamma/2)$, or the high-dimensional analog $\Phi(-\gamma^2/(2\sqrt{\gamma^2 + 2\delta}))$?



circles correspond to the empirical error probabilities, averaged over 10 trials

What can help us in high dimensions?

- An important fact is that high-dimensional phenomena are unavoidable
- If the ratio m/n stays bounded strictly above zero, then it is not possible to achieve the optimal classification rate
- Our only hope is that the data is endowed with some form of *low-dimensional structure*

- What is the underlying cause of the inaccuracy of the prediction for the LDA in high-dimensions?
- The squared Euclidean error

$$\|\hat{\mu} - \mu\|_2^2 = \sum_{j=1}^m (\hat{\mu}_j - \mu_j)^2$$

concentrates sharply around m/n , i.e. for $t \in (0, 1)$

$$\mathbb{P} \left(\left| \|\hat{\mu} - \mu\|_2^2 - \frac{m}{n} \right| \geq \frac{m}{n} t \right) = \mathbb{P} \left(\left| \frac{1}{m} \sum_{j=1}^m Z_j^2 - 1 \right| \geq t \right) \leq 2e^{-\frac{mt^2}{8}}$$

where $Z_j = \sqrt{n}(\hat{\mu}_j - \mu_j) \sim N(0, 1)$; for the upper bound see Wainwright (2019), Example 2.11

Sparsity

- Suppose that the m -vector μ is *sparse*, with only s of its m entries being nonzero, for some sparsity parameter $s \ll m$
- If sparsity holds, we can obtain a better estimator by thresholding the sample means

$$\tilde{\mu}_j = \hat{\mu}_j \mathbb{1}\{|\hat{\mu}_j| > \lambda\}$$

where

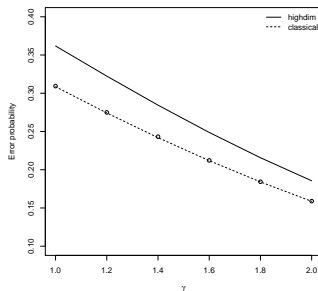
$$\lambda = \sqrt{\frac{2 \log m}{n}}$$

Thresholded mean

Suppose to replace $\hat{\mu}$ by the thresholded mean $\tilde{\mu}$, then

$$\tilde{\Psi}(x) = \langle \tilde{\mu}_A - \tilde{\mu}_B, x - \frac{\tilde{\mu}_A + \tilde{\mu}_B}{2} \rangle$$

approaches the optimal $\text{Err}(\Psi)$ if $\log \binom{m}{s} / n \rightarrow 0$. For $s = 5$:



circles correspond to the empirical error probabilities, averaged over 10 trials

Inference for the mean vector

- Random sample of n observations from $y \sim N_m(\mu, \Sigma)$

$$\begin{pmatrix} y_1 \\ \vdots \\ y_m \end{pmatrix} \sim N_m \left(\begin{pmatrix} \mu_1 \\ \vdots \\ \mu_m \end{pmatrix}, \begin{bmatrix} \sigma_1^2 & \sigma_{12} & \cdot & \sigma_{1m} \\ \sigma_{21} & \sigma_2^2 & \cdot & \sigma_{2m} \\ \cdot & \cdot & \cdot & \cdot \\ \sigma_{m1} & \cdot & \cdot & \sigma_m^2 \end{bmatrix} \right)$$

- The parameter of interest is $\mathbb{E}(y) = \mu$, where $\mu_j = 0$ means “no effect” and $\mu_j \neq 0$ means “effect” in the j th component
- The nuisance parameter is the variance/covariance matrix $\text{Var}(y) = \Sigma$

Three questions

1. *Detecting effects*: There is at least one μ_j different from μ_0 ?
2. *Counting effects*: How many μ_j are different from μ_0 ?
3. *Identifying effects*: Which μ_j are different from μ_0 ?

Global null hypothesis

Testing the global null hypothesis aims at detecting any effect

$$H_0 : \mu = 0, \text{ i.e. } \bigcap_{j=1}^m \{\mu_j = 0\}$$

$$H_1 : \mu \neq 0, \text{ i.e. } \bigcup_{j=1}^m \{\mu_j \neq 0\}$$

One-sided alternative

$$H_0 : \bigcap_{j=1}^m \{\mu_j = 0\}$$

$$H_1 : \bigcup_{j=1}^m \{\mu_j > 0\}$$

MaxT and SumT

- For simplicity, assume $\Sigma = I_m$ and $n = 1$ and consider the one-sided alternative
- $T_j = y_j \sim N(\mu_j, 1)$ for $j = 1, \dots, m$
- $(T_1, \dots, T_m)' \stackrel{H_0}{\sim} N_m(0, I_m)$
- MaxT

$$T_{\max} = \max(T_1, \dots, T_m)$$

- SumT

$$T_{\text{sum}} = \sum_{j=1}^m T_j$$

MaxT

- The critical value $t_{1-\alpha}$ of T_{\max} is

$$P_0(T_{\max} \geq t_{1-\alpha}) = \alpha$$

where $t_{1-\alpha}$ is the $1 - \alpha$ quantile of the distribution of the maximum of m independent standard normal variables

$$\int_{t_{1-\alpha}}^{\infty} m\phi(y)\Phi(y)^{m-1}dy = \alpha$$

where ϕ and Φ are the density and cdf of $N(0, 1)$

Critical value approximation

- We can replace $t_{1-\alpha}$ by $z_{1-\frac{\alpha}{m}}$

$$\begin{aligned} P_0(T_{\max} \geq z_{1-\frac{\alpha}{m}}) &= P_0\left(\bigcup_{j=1}^m \{T_j \geq z_{1-\frac{\alpha}{m}}\}\right) \\ &\leq \sum_{j=1}^m P_0(T_j \geq z_{1-\frac{\alpha}{m}}) = m \frac{\alpha}{m} = \alpha \end{aligned}$$

- The union bound might seem crude, but with independent T_j s the size of the test is very near α

$$\begin{aligned} P_0(T_{\max} \geq z_{1-\frac{\alpha}{m}}) &= 1 - \prod_{j=1}^m P_0(T_j < z_{1-\frac{\alpha}{m}}) \\ &= 1 - \left(1 - \frac{\alpha}{m}\right)^m \xrightarrow{m \rightarrow \infty} 1 - e^{-\alpha} \end{aligned}$$

For $\alpha = 0.05$, $1 - e^{-\alpha} = 0.0487$

Magnitude of the critical value

- How large is the threshold $z_{1-\frac{\alpha}{m}}$? For large m

$$\begin{aligned} z_{1-\frac{\alpha}{m}} &\approx \sqrt{2 \log m} - \frac{\log(2 \log m) + \log 2\pi}{2\sqrt{2 \log m}} \\ &\approx \sqrt{2 \log m} \end{aligned}$$

with no dependence on α

Proof

$$\frac{\phi(t)}{t} \left(\frac{t^2}{t^2 + 1} \right) \leq P(N(0, 1) > t) \leq \frac{\phi(t)}{t}$$

where $\phi(t)$ is the probability density function of $N(0, 1)$. This result implies that for large t , $\frac{\phi(t)}{t}$ is a good approximation to the normal tail probability. Let $z^* = z_{1-\frac{\alpha}{m}}$. We have

$$\frac{\alpha}{m} = P(N(0, 1) > z_{1-\frac{\alpha}{m}}) \approx \frac{\phi(z^*)}{z^*}, \text{ which implies}$$

$$\alpha/m \approx \frac{1}{z^* \sqrt{2\pi}} e^{-\frac{(z^*)^2}{2}}. \text{ Taking the logarithm}$$

$$\log m \approx \frac{1}{2} \log(2\pi) + \frac{1}{2}(z^*)^2 + \log(z^*) + \log(\alpha)$$

Note that z^* is increasing in m , i.e. $m \rightarrow \infty$ induces $z^* \rightarrow \infty$. As $\frac{1}{2} \log(2\pi) + \log(z^*) + \log(\alpha)$ is negligible compared to $(z^*)^2$ when m goes to ∞ , it gives

$$z_{1-\frac{\alpha}{m}} \approx \sqrt{2 \log m}$$

Needle in a haystack problem

$$H_0 : \mu_j = 0 \text{ for all } j = 1, \dots, m$$

$$H_1 : \mu_j = c_m > 0, \mu_k = 0 \text{ for } k \neq j$$

- What is the limiting power of the MaxT test?

$$\lim_{m \rightarrow \infty} P_1(T_{\max} > z_{1-\frac{\alpha}{m}})$$

It depends on the limiting ratio

$$\lim_{m \rightarrow \infty} \frac{c_m}{\sqrt{2 \log m}} \leq 1$$

where c_m is the value of the single non-zero mean, which depends on m

Two cases:

- Assume without loss of generality that $\mu_1 = c_m$. Suppose $c_m > (1 + \epsilon)\sqrt{2 \log m}$. Then, for $m \rightarrow \infty$

$$P_1(T_{\max} > z_{1-\frac{\alpha}{m}}) \geq P_1(T_1 > z_{1-\frac{\alpha}{m}}) = P(N(0, 1) > z_{1-\frac{\alpha}{m}} - c) \rightarrow 1$$

- Suppose $c_m < (1 - \epsilon)\sqrt{2 \log m}$. Then for $m \rightarrow \infty$

$$\begin{aligned} P_1(T_{\max} > z_{1-\frac{\alpha}{m}}) &\leq P(T_1 > z_{1-\frac{\alpha}{m}}) + P(\max_{j>1} T_j > z_{1-\frac{\alpha}{m}}) \\ &= P(N(0, 1) > z_{1-\frac{\alpha}{m}} - c) + P(\max_{j>1} T_j > z_{1-\frac{\alpha}{m}}) \\ &\rightarrow 0 + (1 - e^{-\alpha}) \end{aligned}$$

and the MaxT test has no power

SumT

$$T_{\text{sum}} = \sum_{j=1}^m T_j \sim N\left(\sum_{j=1}^m \mu_j, m\right)$$

$$- \frac{T_{\text{sum}}}{\sqrt{m}} \stackrel{H_0}{\approx} N(0, 1); \frac{T_{\text{sum}}}{\sqrt{m}} \stackrel{H_1}{\approx} N(\theta_m, 1) \text{ with}$$

$$\theta_m = \frac{\sum_{j=1}^m \mu_j}{\sqrt{m}}$$

but if $\theta_m \rightarrow 0$ when $m \rightarrow \infty$, then T_{sum} has no power

- By the Neyman-Pearson lemma, T_{sum} is the UMP test for

$$H_0 : \mu_j = 0 \text{ for all } j$$

$$H_1 : \mu_j = c_m > 0 \text{ for all } j$$

where $\theta_m = \sqrt{m}c_m$, but if $c_m = \frac{1}{m}$ the UMP test has no power

Comparison

- **Few strong effects:**

$m^{1/4}$ of the μ_j s are equal to $\sqrt{2 \log m}$, the rest 0.

E.g. when $m = 10^6$, $m^{1/4} \approx 36$ and $\sqrt{2 \log m} \approx 5.3$. In this setting T_{\max} has full power, but T_{sum} has no power because

$$\theta_m = \frac{m^{1/4} \sqrt{2 \log m}}{\sqrt{m}} \rightarrow 0$$

- **Small, distributed effects:**

$\sqrt{2m}$ of the μ_j s are equal to 3, the rest 0.

The T_{sum} has (almost) full power, but T_{\max} has no power because when m is large it's very likely that the largest y_j value comes from a null μ_j

MinP

- Let $p_j = 1 - \Phi(T_j)$ be the j th p -value
- $p_1, \dots, p_m \stackrel{i.i.d.}{\sim} U(0, 1)$ under H_0
- The MinP test is based on the minimum p -value

$$p_{\min} = \min(p_1, \dots, p_m) \stackrel{H_0}{\sim} \text{Beta}(1, m)$$

- The MinP test rejects H_0 if $p_{\min} \leq 1 - (1 - \alpha)^{\frac{1}{m}}$ and has size α :

$$\begin{aligned} \mathbb{P}_0(p_{\min} \leq 1 - (1 - \alpha)^{\frac{1}{m}}) &= 1 - \mathbb{P}_0\left(\bigcap_{i=1}^m \{p_i > 1 - (1 - \alpha)^{\frac{1}{m}}\}\right) \\ &= 1 - [(1 - \alpha)^{\frac{1}{m}}]^m = \alpha \end{aligned}$$

Simes test

- $p_1, \dots, p_m \stackrel{i.i.d.}{\sim} U(0, 1)$ under H_0
- Sort the p -values

$$p_{(1)} \leq p_{(2)} \leq \dots \leq p_{(m)}$$

- The null distribution of j th ordered p -value is

$$p_{(j)} \stackrel{H_0}{\sim} \text{Beta}(j, m - j + 1)$$

- The Simes test p -value

$$p_s = \min_{j=1, \dots, m} \left\{ p_{(j)} \frac{m}{j} \right\} \stackrel{H_0}{\sim} U(0, 1)$$

- It rejects H_0 if

$$\exists j : p_{(j)} \leq \frac{\alpha j}{m}$$

Fisher combination

- $p_1, \dots, p_m \stackrel{i.i.d.}{\sim} U(0, 1)$ under H_0
- Fisher's method of combining p -values

$$T_f = \sum_{j=1}^m 2 \log \left(\frac{1}{p_j} \right) \stackrel{H_0}{\sim} \chi_{2m}^2$$

Higher criticism

- Empirical cdf $\hat{F}_m(t) = \frac{\sum_{j=1}^m \mathbb{1}\{p_j \leq t\}}{m}$ for $t \in [0, 1]$.
- Since $p_1, \dots, p_m \stackrel{i.i.d.}{\sim} U(0, 1)$ under H_0 , then

$$m\hat{F}_m(t) \stackrel{H_0}{\sim} \text{Binomial}(m, t)$$

- The higher criticism test is

$$T_{\text{hc}} = \sup_{t \in [0, 1]} \frac{\hat{F}_m(t) - t}{\sqrt{t(1-t)/m}}$$

or equivalently

$$T_{\text{hc}} = \max_{j=1, \dots, m} \sqrt{m} \frac{(j/m) - p_{(j)}}{\sqrt{p_{(j)}(1 - p_{(j)})}}$$

Reference

Donoho, Jin (2015)

Higher Criticism for Large-Scale Inference, Especially for Rare and Weak Effects
Statistical Science, 30:1–25

- For $m \rightarrow \infty$, $b_m T_{\text{hc}} - c_m$ converges weakly to the standard Gumbel distribution, where $b_m = \sqrt{2 \log \log m}$ and $c_m = \frac{1}{2}(\log \log \log(m) - 4\pi)$
- For any fixed α and $m \rightarrow \infty$, its critical value is

$$t_{1-\alpha} \approx (1 + a) \sqrt{2 \log \log m}$$

for some $a > 0$, e.g. $a = 1.08$ for $m \approx 10^6$ and $\alpha = 0.05$

Mixture distribution

- We assume that our samples follow a mixture of $N(0, 1)$ and $N(\mu, 1)$ distributions

$$H_0 : y_j \stackrel{i.i.d}{\sim} N(0, 1)$$

$$H_1 : y_j \stackrel{i.i.d}{\sim} \pi_0 N(0, 1) + \pi_1 N(\mu, 1)$$

where $\pi_1 = 1 - \pi_0$

- To carry out asymptotic analysis, we must specify the dependence scheme of $\pi_1 = \pi_1(m)$ and $\mu = \mu(m)$ on m :

$$\begin{aligned} \pi_1 &= m^{-\beta} & \frac{1}{2} < \beta < 1 \\ \mu &= \sqrt{2r \log m} & 0 < r < 1 \end{aligned}$$

- The needle in a haystack problem: $\beta = 1$ and $r = 1$; small distributed effects: $\beta = 1/2$

Threshold curve

Consider the following threshold curve for r

$$\rho_{\text{hc}}(\beta) = \begin{cases} \beta - \frac{1}{2} & \text{if } \frac{1}{2} < \beta \leq \frac{3}{4} \\ (1 - \sqrt{1 - \beta})^2 & \text{if } \frac{3}{4} \leq \beta \leq 1 \end{cases}$$

- If $r > \rho_{\text{hc}}(\beta)$ the Neyman-Pearson optimal test achieves

$$P_0(\text{Type I Error}) + P_1(\text{Type II Error}) \rightarrow 0$$

The Higher Criticism is asymptotically equivalent to the optimal test without knowledge of π_1 and/or μ

- If $r < \rho_{\text{hc}}(\beta)$ then for *any* test

$$\liminf_{m \rightarrow \infty} P_0(\text{Type I Error}) + P_1(\text{Type II Error}) \geq 1$$

Detectable region

