Theory

- Fundamental Ideas of Statistical Learning
- Multivariate variability
- Principal Component Analysis
- Gaussian Model
- Estimator of the mean and variance of a Gaussian
- Inference for the mean
- Inference for linear combination of \$\underline{\mu}\$
- Large scale hypothesis testing and FDR
- · Comparing means of multivariate Gaussian Distr.
- Extension to two-way ANOVA.
- Classification
- · Evaluating a classifier
- Regression
- Ensemble methods

Latex new feature: ⊥

Recall orthogonal projection

Remember:

$$||\underline{x}|| = \sqrt{\sum_i u_i^2} = \underline{u}' \cdot \underline{u} = < u, u> \quad \Rightarrow \quad cos\theta = \frac{\underline{u}' \cdot \underline{v}}{||\underline{u}|| \cdot ||\underline{v}||} = \frac{<\underline{u},\underline{v}>}{||\underline{u}|| \cdot ||\underline{v}||}$$

From this you can derive the orthogonal projection of u on v as:

$$\pi_{u|v} = rac{\overbrace{v \cdot v'}^{\in \mathbb{R}^{n imes n}}}{\underbrace{v' \cdot v'}_{\in \mathbb{R}}} \cdot \underline{u}$$

Operator $T = \frac{\underline{v} \cdot \underline{v'}}{\underline{v'} \cdot \underline{v}}$ has this property:

- linearity;
- symmetric (self-adjoint);
- idempotent $\leftrightarrow T = T \cdot T = T^2$

So T thanks to this property it is an #orthogonal_projection_operator .

Sample Mean

Usually we have as columns each feature and as rows each observation.

#Sample_mean of the first column \underline{c}_1 is the projection of it on the $\underline{1}$ (which is the subspace of \mathbb{R}^n without variability).



$$\text{mean of }\underline{c}_1 = \overline{x}_1 \cdot \underline{1} = \frac{1}{n} \sum_i x_{i,1} \cdot 1 = \frac{\underline{1}' \cdot \underline{c}_1}{\underline{1}' \cdot \underline{1}} \cdot \underline{1} = \frac{\underline{1} \cdot \underline{1}'}{\underline{1}' \cdot \underline{1}} \cdot \underline{c}_1 = \pi_{\underline{c}_1 | \underline{1}}$$

Remark: \underline{c}_1 and $\underline{1}$ are column vectors

Remark: Could be more than a vector with the same mean so we create \underline{d}_1

Sample variance and standard deviation

#Sample_standard_deviation =
$$\sqrt{s_{1,1}}=[\frac{1}{n-1}\sum_i(x_{i,1}-\overline{x}_1)^2]^{\frac{1}{2}}$$
 #Sample_variance = $s_{1,1}=\frac{1}{n-1}\sum_i(x_{i,1}-\overline{x}_1)^2$

From this we can create the | #deviation | vector = $\underline{d}_1 = \underline{c}_1 - \overline{x}_1 \cdot \underline{1}$

 $\Rightarrow \quad ||d_1|| = \text{how good is the approximation done by } \pi_{\underline{c}_1|\underline{1}} \text{ of } \underline{c}_1 = \sqrt{n-1} \cdot \sqrt{s_{1,1}}$

If our data follow a Gaussian distribution we know that:

- 68% of our data are in the interval $[\overline{x}_1 \pm \sqrt{s_{1,1}}]$
- 95% of our data are in the interval $[\overline{x}_1 \pm s_{1,1}]$

Otherwise, try to reach Gaussianity with:

- transformation of our data (example: doing log(data))
- using <code>#Chebyscev_inequality</code> : $F_k\{x_i\in[\overline{x}_1\pm k\sqrt{s_{1,1}}]\}<1-\frac{1}{k^2};$ if $k=2\Rightarrow$ we are covering about 75%

Interpretation: we are paying the unknown distribution with a larger interval

A different ways to calculate $\cos(\theta_{1,2})$:

$$\cos(\theta_{1,2}) = \frac{< d_1, d_2>}{||d_1||\cdot||d_2||} = \frac{d_1'd_2}{\sqrt{n-1}\sqrt{s_{1,1}}\cdot\sqrt{n-1}\sqrt{s_{2,2}}} = \frac{\sum_i (x_{i,1}-\overline{x}_1)(x_{i,2}-\overline{x}_2)}{(n-1)\sqrt{s_{1,1}\cdot s_{2,2}}} = \frac{cov(\underline{x}_1,\underline{x}_2)}{\sqrt{s_{1,1}\cdot s_{2,2}}}$$

So we reach that:

$$\cos(\theta_{1,2}) = \frac{s_{1,2}}{\sqrt{s_{1,1} \cdot s_{2,2}}} = \rho_{1,2} = corr(\underline{x}_1,\underline{x}_2) = \text{ correlation between } \underline{x}_1 \text{ and } \underline{x}_2$$

Given two column $\underline{c}_1,\underline{c}_2$ with their respective $\overline{x}_1,\overline{x}_2,d_1,d_2$ we have two limit case:

- $\begin{array}{l} \bullet \;\; \theta_{1,2} = 0 \colon \\ \qquad d_2 \in span\{d_1\} \Rightarrow \exists \beta : d_2 = d_1 \cdot \beta \Leftrightarrow x_{i,2} \overline{x}_2 = \beta \cdot (x_{i,1} \overline{x}_1) \Rightarrow \; \text{perfect correlation} \end{array}$
- $heta_{1,2}=rac{\pi}{2}$: \underline{c}_1 give us none information about \underline{c}_2 and viceversa $\Rightarrow
 ho_{1,2}=0={
 m \ zero \ correlation}$

To summarize $X \in \mathbb{R}^{n \times p}$, where n is the number of observation while p is the number of features for each observation, we can create:

- \overline{x} which is the #mean_vector of each features $(\in \mathbb{R}^p)$;
- #covariance_matrix $S \in \mathbb{R}^{p \times p}$ s.t. $s_{j,k} = \frac{1}{n-1} \sum_i (x_{i,j} \overline{x}_j)(x_{i,k} \overline{x}_k)$ From S we can deduce the correlation matrix $\rho \in \mathbb{R}^{p \times p}$.



Remark: standardize a variable means how far each element is from the mean in terms of standard deviation and implies that $S = \rho$

Fundamental Ideas of Statistical Learning

Goal: explain variability of y as a function of \underline{x}

Formally: find $\hat{f}:\mathbb{R}^p o\mathbb{R}$ s.t. $\hat{f}(\underline{x})=\mathop{argmin}_{f(x)}\{\mathbb{E}[|y-f(\underline{x})|^2]\}$

Solution: $\pi_{Y|\sigma(x)} = \hat{f}(\underline{x}) = \mathbb{E}[y|\underline{x}]$ which is called <code>#regression_function</code>

Rmk - 1: it is the Radon-Nykodyn derivative, because it allows to describe the distribution of y using distribution of features.

Proof:

$$egin{aligned} \mathbb{E}[|y-f(\underline{x})|^2] &= \mathbb{E}[|y-\mathbb{E}[y|\underline{x}] + \mathbb{E}[y|\underline{x}] - f(\underline{x})|^2] = \ &= \mathbb{E}[|y-\mathbb{E}[y|x]|^2] + \mathbb{E}[|\mathbb{E}[y|x] - f(x)|^2] + 2\mathbb{E}[(y-\mathbb{E}[y|x])(\mathbb{E}[y|x] - f(x))] \end{aligned}$$

But we notice that:

$$2\mathbb{E}[(y-\mathbb{E}[y|\underline{x}])(\mathbb{E}[y|\underline{x}]-f(\underline{x}))] = 2\mathbb{E}[\mathbb{E}[(y-\mathbb{E}[y|\underline{x}])(\mathbb{E}[y|\underline{x}]-f(\underline{x}))|\underline{x}]] = \\ = 2\mathbb{E}[(\mathbb{E}[y|\underline{x}]-f(\underline{x}))\cdot\underbrace{\mathbb{E}[(y-\mathbb{E}[y|\underline{x}])|\underline{x}]}_{\mathbb{E}[y|x]-\mathbb{E}[y|x]=0}] = 0$$

Since we remain with:

 $\mathbb{E}[|y - \mathbb{E}[y|x]|^2] + \mathbb{E}[|\mathbb{E}[y|x] - f(x)|^2]$

to minimize it we need that $f(x) = \mathbb{E}[y|x]$

П

Since y is a random variable we know that our general model is:

$$y = \mathbb{E}[y|\underline{x}] + \epsilon \Rightarrow \mathbb{E}[y] = \mathbb{E}[\mathbb{E}[y|\underline{x}]] + \mathbb{E}[\epsilon] = \mathbb{E}[y] + \mathbb{E}[\epsilon] \Rightarrow \mathbb{E}[\epsilon] = 0$$

So we found that: $\epsilon \perp \sigma(\underline{x})$

Problem: how to build an operator $F: \mathbb{X} \to \hat{f}(\underline{x})$ s.t. given a dataset \mathbb{X} it give us the best regression function.

Simple idea: check neighbourhood of the input point.

K-Nearest Neighbors - #knn

Given $x\in\mathbb{R}$ we define as $N_x=\{x_i \text{ in the training set }: \text{ close to } x\}$, $dim\{N_x\}=k$ From that we can estimate $\hat{f}(x)=rac{1}{k}\sum_{x_i\in N_x}f(x_i)$

Curse of dimensionality

This method isn't always good since it suffer from the problem called $\#curse_of_dimensionality$. Explanatory example: find r to navigate 10% of an area centered in 0 with radius 1.

$$p = 1$$
:

We are working on $\mathbb{R} \Rightarrow rac{2 \cdot r}{2} = 0, 1 \rightarrow r = 0, 1$

p = 2:



We are working on $\mathbb{R}^2 \Rightarrow rac{\pi \cdot r^2}{\pi} = 0, 1 o r = 0, 31$

p = 3:

We are working on $\mathbb{R}^3 \Rightarrow rac{rac{4}{3}\pi\cdot r^3}{rac{4}{3}\pi} = 0, 1 o r = 0, 46$

p = 100:

We are working on $\mathbb{R}^{100} \Rightarrow r = 0, 1^{rac{1}{100}} = 0, 97$

To solve this problem we can:

- Reduce the number of features p using Principal Component Analysis #PCA;
- Use structured model to reduce the dimension of \hat{f} so it depends from a parameter $\theta \in \mathbb{R}^k$;

Bias-Variance Tradeoff

Explanatory example

Framework: linear model: $f_{\theta}(\underline{x}) = \beta_0 + \sum_{i \in \{1,...,p\}} \beta_i \cdot x_i$ with $\theta = \{\beta_i\}_{i \in \{1,...,p\}}$

Knowing that $y = f(\underline{x}) + \epsilon$

We define
$$\#$$
generalization_error $= \mathbb{E}_{\cdot \mid \mathbb{X}}[(y - \hat{f}(\underline{x}))^2] = \underbrace{(f(\underline{x}) - \hat{f}(\underline{x}))^2}_{\text{reducible error}} + \underbrace{var(\epsilon)}_{\text{irreducible error}}$

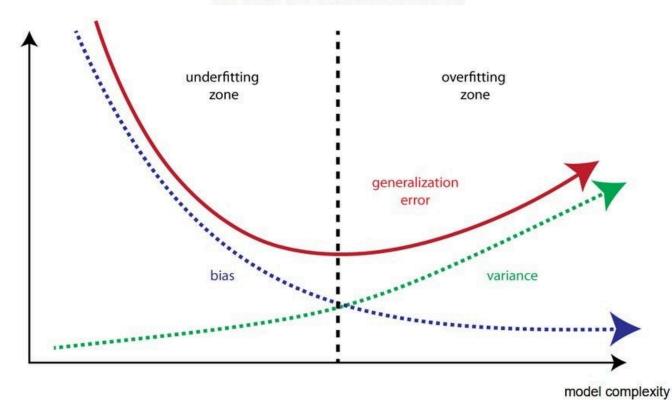
There could be 2 causes if the generalization error is 0:

- · phenomena is deterministic;
- test data are data from the training set, so we are not learning anything.

So we can create the #expected generalization error as:

$$\mathbb{E}[\mathbb{E}_{\cdot|\mathbb{X}}[(y-\hat{f}(\underline{x}))^2]] = \mathbb{E}[(f(\underline{x})-\hat{f}(\underline{x}))^2] + var(\epsilon) = \underbrace{(f(\underline{x})-\mathbb{E}[\hat{f}(\underline{x})])^2}_{bias^2} + \underbrace{\mathbb{E}[(\hat{f}(\underline{x})-\mathbb{E}[\hat{f}(\underline{x})])^2]}_{var(\hat{f}(x))} + var(\epsilon)$$

the bias vs. variance trade-off





Recall

We work with $\underline{x}_1,\dots,\underline{x}_n\in\mathbb{R}^p$ $\mathrm{iid}\ \sim\underline{x}$ with $\mathbb{E}[\underline{x}]=\underline{\mu}$ and $cov(\underline{x})=\Sigma$

We have the following sample estimator:

•
$$\underline{\mu} \rightarrow \frac{1}{n} \sum_{i=1}^{n} \underline{x}_{i} = \overline{\underline{x}}$$

•
$$\Sigma
ightarrow rac{1}{n} \sum_{i}^{n} (\underline{x}_{i} - \overline{\underline{x}}) (\underline{x}_{i} - \overline{\underline{x}})' = S_{n}$$

Prop - properties of sample mean and sample variance

- $\bullet \ \ \mathbb{E}[\overline{\underline{x}}] = \mu \leftarrow \mathsf{unbiased}$
- $cov(\overline{\underline{x}}) = \frac{\Sigma}{n}$
- $\mathbb{E}[S_n] = \frac{n}{n-1}\Sigma \Rightarrow \mathbb{E}[\frac{n-1}{n}S_n] = \Sigma \leftarrow \text{we need } n-1 \text{ because } S_n \text{ must be orthogonal to } span\{1\} \text{ so the linear subspace of means.}$

Proof - previous Prop

First point:

$$\mathbb{E}[\overline{x}] = \mathbb{E}egin{array}{c} \left[\overline{x}_1
ight] &= \left[\mathbb{E}[\overline{x}_1]
ight] \ dots & \left[\mathbb{E}[\overline{x}]
ight] & \Longrightarrow orall k \in \{1,\ldots,p\} & \mathbb{E}[\overline{x}_k] = rac{1}{n}\sum_i^n \mathbb{E}[x_{i,k}] = \mu_k \end{cases}$$

Second one:

$$\begin{split} &cov(\overline{\underline{x}}) = & \mathbb{E}[(\overline{\underline{x}} - \underline{\mu})(\overline{\underline{x}} - \underline{\mu})'] = \\ &= & \mathbb{E}[\{\frac{1}{n}\sum_{i}^{n}(\underline{x}_{i} - \underline{\mu})\}\{\frac{1}{n}\sum_{j}^{n}(\underline{x}_{j} - \underline{\mu})\}] \\ &= & \frac{1}{n^{2}}\sum_{i,j}\mathbb{E}[(\underline{x}_{i} - \underline{\mu})(\underline{x}_{j} - \underline{\mu})] = \begin{cases} 0 & i \neq j \leftarrow \text{ independent} \\ \sigma & i = j \leftarrow \text{ identically distribuited} \end{cases} \\ &= & \frac{1}{n^{2}} \cdot n\Sigma = \frac{\Sigma}{n} \end{split}$$

Take \underline{c}_i as a column of \mathbb{X} , we find that $\text{\#deviation} = \underline{d}_j = \underline{c}_j - \overline{x}_j \underline{1} = \underline{c}_j - \frac{\underline{1}\underline{1}'}{\underline{1}'\underline{1}} \underline{c}_j = [I - \frac{\underline{1}\underline{1}'}{\underline{1}'\underline{1}}]\underline{c}_j$ Rmk: With $[I - \frac{\underline{1}\underline{1}'}{\underline{1}'\underline{1}}]$ be the $\text{\#orthogonal_projection_operator}$ on $\mathcal{L}^\perp(\underline{1})$

$$\Rightarrow S = \frac{1}{n-1}\begin{bmatrix}\underline{d}_1'\underline{d}_1 & \underline{d}_1'\underline{d}_2 & \dots \\ \dots & \dots & \dots \\ \dots & \dots & \underline{d}_p'\underline{d}_p\end{bmatrix} = \frac{1}{n-1}d'd = \frac{1}{n-1}\mathbb{X}'[I - \frac{11'}{\underline{1}'\underline{1}}]'[I - \frac{11'}{\underline{1}'\underline{1}}]\mathbb{X}$$

But since we know that the orthogonal projection operator is symmetric and idempotent we reach:

$$S = \frac{1}{n-1} \mathbb{X}' [I - \frac{\underline{1}\underline{1}'}{\underline{1}'\underline{1}}] \mathbb{X}$$



Multivariate variability

We define:

- #generalized_variance = det(S);
- #total variance = trace(S)

Geometric interpretation - p = 2

$$S = rac{1}{n-1}egin{bmatrix} \underline{d}_1' \underline{d}_1 & \underline{d}_1' \underline{d}_2 \ \underline{d}_2' \underline{d}_1 & \underline{d}_2' \underline{d}_2 \end{bmatrix} \Rightarrow det(S) = rac{||\underline{d}_1||^2 ||\underline{d}_2||^2 sin^2 heta_{1,2}}{(n-1)^2}$$

So:

- det(S): catch the squared area of the parallelogram with as sides \underline{d}_1 and \underline{d}_2 .
- trace(S): catch the perimeter of the parallelogram with as sides \underline{d}_1 and \underline{d}_2 , so don't catch changes of $\theta_{1,2}$

Prop

 $det(S) = 0 \Leftrightarrow \underline{d}_1, \dots, \underline{d}_p$ are linearly dependent

Dim - previous Prop

Corollary

If
$$p \ge n \Rightarrow det(S) = 0$$

Interpretation: more features to analyze than data available

Proof - previous Corollary

Take $\underline{d}_1,\ldots,\underline{d}_p$ with $p\geq n$, with $\underline{d}_i\in\mathbb{R}^n$ and $\underline{d}_i\in\mathcal{L}^\perp(\underline{1})$ so we have that $dim(\mathcal{L}^\perp)\leq n-1$ Since $p\geq n$ we have more vectors than degree of freedom \Rightarrow some are linearly dependent $\Rightarrow det(S)=0$

Prop



$$\left\{ egin{aligned} ext{S is positive semi-defined} \ & \Rightarrow S ext{ is positive defined} \end{aligned}
ight.
ight.$$

Proof - previous Prop

Since it is positive semi-defined we know that:

- $\forall \underline{c} \in \mathbb{R}^p$ $\underline{c}' S \underline{c} = \frac{1}{n-1} \underline{c}' d' d \underline{c} = \frac{1}{n-1} ||d \underline{c}'||^2 \geq 0;$
- $\forall i \quad \lambda_i \geq 0;$

By contradiction:

From hypothesis we know that $det(S) \neq 0$ if:

$$\exists \underline{c} \neq \underline{0} : \underline{c}' S \underline{c} \Rightarrow ||\underline{d}\underline{c}||^2 = 0 \Rightarrow \underline{d}\underline{c} = 0 \Rightarrow \underline{d}_1, \dots, \underline{d}_p \text{ lin. dep. } \Rightarrow \det(S) = 0 \text{ (contraddiction)}$$

Since $\not\exists \underline{c} : \underline{c}' S \underline{c} = 0$ then S is positive defined.

Since S is positive semi-defined so $\forall i \quad \lambda_i \geq 0 \Rightarrow S = P\Lambda P'$

With:

- $P = \text{eigenvectors matrix } (\underline{e}_i \text{ is the } i^{th} \text{ eigenvector});$
- $\Lambda=$ ordered eigenvalues matrix so $\lambda_1\geq \lambda_2\geq \cdots \geq \lambda_p\geq 0$

Rmk: if $det(S)
eq 0 \Rightarrow S^{-1} = \sum_i \frac{1}{\lambda_i} \underline{e_i} \underline{e_i'}$

Def - #Mahalanobis distance

Mahalanobis distance $=\underline{d}_{S^{-1}}^2(\underline{x},y)=(\underline{x}-y)'S^{-1}(\underline{x}-y)$

Interpretation: $\underline{d}_{S^{-1}}^2(\underline{x},\underline{y})=3$ means that \underline{x} and \underline{y} are 3 standard deviation apart w.r.t. their joint distribution.

We call $\epsilon_r(\overline{\underline{x}})=$ sphere centered in $\overline{\underline{x}}$ of radius r w.r.t. Mahalanobis distance

In \mathbb{R}^n this is an ellipse centered in $\underline{\overline{x}}$ with axis in the same direction of \underline{e}_i and semi-axis equal to $\alpha \frac{1}{\sqrt{\lambda_i}} r$ with $(\lambda_i, \underline{e}_i)$ eigenvalues and eigenvectors of S.

 $\Rightarrow \text{ volume of this ellipse} = lpha \cdot \sqrt{\lambda}_1 \cdot \sqrt{\lambda}_2 \cdot \dots \cdot \sqrt{\lambda}_p \cdot r^p = \sqrt{\overline{det(S)}} \cdot r^p$

Principal Component Analysis

Intuition behind Principal Component Analisys #PCA

We can always decompose S with the spectral theorem, so it can be written as $S = P\Lambda P'$.

Given X we can change the coordinates of each vector composing our dataset using P so:

$$orall x \in \mathbb{X} \quad x o ilde{x} = P'x \in ilde{\mathbb{X}}$$

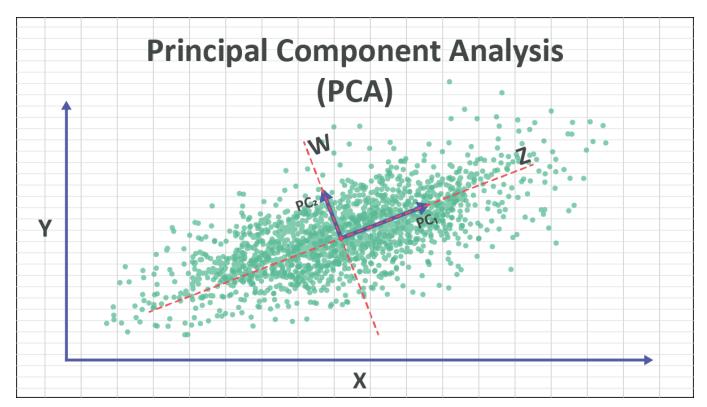
After this transformation we need to re-calculate the $\# covariance_matrix$ \tilde{S} and we find that:

$$ilde{S}=rac{1}{n-1} ilde{\mathbb{X}}'[I-rac{11'}{1'1}] ilde{\mathbb{X}}=P'SP=P'P\Lambda P'P=\Lambda$$

Important:

Using as axis the eigenvector of the covariance matrix, covariances between features become 0. Idea: covariances depends only by the reference system used.





We call as **Principal Component** the linear subspace which maximize the variance of our data. Given a random variable $\underline{x} \in \mathbb{R}^p$ s.t. $\underline{x} \sim \mu, \Sigma$ we want p vectors $\underline{a}_1, \ldots, \underline{a}_p$ such that:

- 1. Find $\underline{a}_1 \in \mathbb{R}^p$ with $||\underline{a}_1|| = 1$ s.t. $Var(\underline{a}_1'\underline{x})$ is maximized;
- 2. Find $\underline{a}_2 \in \mathbb{R}^p$ with $||\underline{a}_2|| = 1$ s.t. $Var(\underline{a}_2'\underline{x})$ is maximized and $cov(\underline{a}_1\underline{x},\underline{a}_2\underline{x}) = 0$;
- 3. Find $\underline{a}_3 \in \mathbb{R}^p$ with $||\underline{a}_3|| = 1$ s.t. $Var(\underline{a}_3'\underline{x})$ is maximized and: $cov(\underline{a}_1\underline{x},\underline{a}_3\underline{x}) = 0$, $cov(\underline{a}_2\underline{x},\underline{a}_3\underline{x}) = 0$;
- 4. . . . ;

Recall: $var(\mathbb{X}) = S$ so $var(\underline{a}'\mathbb{X}) = \underline{a}'S\underline{a}$ and $\underline{x}'\underline{x} = ||x||$

Lemma

Take the matrix $B \in p \times p$ symmetric and positive defined s.t. $B = \sum_{i=1}^{p} \lambda_i \underline{e}_i \underline{e}_i'$ we know that:

- $\bullet \ \ \max_{\underline{x} \in \mathbb{R}^p} \tfrac{\underline{x'B\underline{x}}}{\underline{x'\underline{x}}} = \lambda_1 \ \text{and} \ \arg\max_{\underline{x} \in \mathbb{R}^p} \tfrac{\underline{x'B\underline{x}}}{\underline{x'\underline{x}}} = \underline{e}_1;$
- $\bullet \ \ \max_{\underline{x} \in \mathbb{R}^p, \underline{x} \perp \underline{e}_1} \frac{\underline{x'Bx}}{\underline{x'x}} = \lambda_2 \ \text{and} \ \arg\max_{\underline{x} \in \mathbb{R}^p, \underline{x} \perp \underline{e}_1} \frac{\underline{x'Bx}}{\underline{x'x}} = \underline{e}_2$
- ... so on ...

Proof - previous Lemma

First principal component:

Taking $x \neq 0$:

$$rac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = rac{\underline{x}'P\Lambda P'\underline{x}}{\underline{x}'\underline{x}} = rac{\underline{x}'P\Lambda P'\underline{x}}{\underline{x}'PP'\underline{x}} \stackrel{\underline{y}=P'\underline{x}}{=} rac{\underline{y}'\Lambda\underline{y}}{\underline{y}'\underline{y}} \leq \lambda_1 rac{\sum_i^p y_i^2}{\sum_i^p y_i^2} = \lambda_1$$

If $\underline{x} = \underline{e}_1 o rac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = \lambda_1 \Rightarrow \max_{\underline{x} \in \mathbb{R}^p} rac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = \lambda_1 ext{ and } rg \max_{\underline{x} \in \mathbb{R}^p} rac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = \underline{e}_1$



Second principal component:

Taking $\underline{x} \neq 0$:

$$\frac{\underline{x'B}\underline{x}}{\underline{x'x}} = \frac{\underline{x'P}\Lambda P'\underline{x}}{\underline{x'x}} = \frac{\underline{x'P}\Lambda P'\underline{x}}{\underline{x'P}P'\underline{x}} \stackrel{\underline{y=P'\underline{x}, \perp \underline{e}_1}}{=} \frac{\underline{y'}\Lambda\underline{y}}{\underline{y'y}} \leq \lambda_2$$

So we reach that: $\max_{\underline{x} \in \mathbb{R}^p, \underline{x} \perp \underline{e}_1} \frac{\underline{x'Bx}}{\underline{x'x}} = \lambda_2 \text{ and } \arg\max_{\underline{x} \in \mathbb{R}^p, \underline{x} \perp \underline{e}_1} \frac{\underline{x'Bx}}{\underline{x'x}} = \underline{e}_2$

It's curious that to satisfy $\underline{y} \perp \underline{e}_1$ we reach that $\underline{y} = P'\underline{x} = \begin{bmatrix} \underline{e}_2\underline{x} \\ \vdots \\ \underline{e}_xx \end{bmatrix}$

p principal component:

Taking $\underline{x} \neq 0$:

$$\frac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = \frac{\underline{y}'\Lambda\underline{y}}{\underline{y}'\underline{y}} = \frac{\sum_{i}^{p}\lambda_{i}y_{i}^{2}}{\sum_{i}^{p}y_{i}^{2}} \geq \lambda_{p}\frac{\sum_{i}^{p}y_{i}^{2}}{\sum_{i}^{p}y_{i}^{2}} = \lambda_{p}$$

$$\text{If } \underline{x} = \underline{e}_{p} \rightarrow \frac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = \lambda_{p} \Rightarrow \min_{\underline{x} \in \mathbb{R}^{p}} \frac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = \lambda_{p} \text{ and } \arg\min_{\underline{x} \in \mathbb{R}^{p}} \frac{\underline{x}'B\underline{x}}{\underline{x}'\underline{x}} = \underline{e}_{p}$$

Rmk: $\underline{a} \perp \underline{b} \Rightarrow cov(\underline{a},\underline{b}) = \underline{a}'\Sigma\underline{b} = \underline{b}'\Sigma\underline{a} = 0$

We define:

- #scores = $y_i = \underline{e}'_i \underline{x}$ = score of i^{th} principal component, it's the coordinate of the projection of \underline{x} on the \underline{e}_i axis;
- $\# loadings = \underline{e}_i = vector describing the <math>i^{th}$ principal component;

 (y_i,\underline{e}_i) is the i^{th} principal component.

It's useful to centered our data doing this:

$$y_1 = \underline{e}_1'(\underline{x} - \mu)$$

Corollary

$$cov(y_i,y_j) = \lambda_{i,j} \cdot \delta_{i,j} = egin{cases} \lambda_i & i = j \ 0 & else \end{cases}$$

Proof - previous Corollary

$$cov(y_i,y_j) = cov(\underline{e}_i'\underline{x},\underline{e}_j'\underline{x}) = \underline{e}_i'\Sigma\underline{e}_j' = \lambda_i\underline{e}_i\underline{e}_j' = egin{cases} \lambda_i & i=j \ 0 & ext{else since } \underline{x}_i \perp \underline{x}_j \end{cases}$$

Framework:

- take $\underline{x}\in\mathbb{R}^p$ s.t. $\mathbb{E}[\underline{x}]=\mu$ and $cov(\underline{x})=\Sigma=P\Lambda P'$ with $det(\Sigma)\neq 0$
- Projection into eigen-space = Principal component $= y = P'(\underline{x} \mu) = \begin{bmatrix} \underline{e}_1'(\underline{x} \underline{\mu}) & \dots & \underline{e}_p'(\underline{x} \underline{\mu}) \end{bmatrix}'$

$$egin{array}{cccc} - & - & - & - & - & - \ ullet & \Rightarrow cov(y) = \Lambda ext{ and } \mathbb{E}[y] = 0 \end{array}$$

Sometimes is useful to give the proportion of variance explained by the i^{th} principal component $y_i = \underline{e}_i(\underline{x} - \underline{\mu})$. To do so we calculate $\frac{\lambda_i}{\sum_i \lambda_j}$.



#PCA 's problem: interpretation of loading

 $orall i \quad y_i = \underline{e}_i'(\underline{x} - \mu) = \sum_{j=1}^p e_{i,j}(x_j - \mu_j)$ we call j^{th} the component of the i^{th} #loadings \underline{e}_i as $e_{i,j}$.

Prop

$$orall i, k = 1, \dots, p \quad corr(y_i, x_k) = e_{k,i} rac{\sqrt{\lambda_i}}{\sqrt{\sigma_{kk}}}$$

Proof - previous Prop

Recall:
$$corr(y_i, x_k) = rac{cov(y_i, x_k)}{\sqrt{\lambda_i} \sqrt{\sigma_{kk}}}$$

$$cov(\underline{y}_{i}, x_{k}) = cov(\underline{e}_{i}'(\underline{x} - \underline{\mu}), \underline{u}_{k}'\underline{x}) = cov(\underline{e}_{i}'(\underline{x}, \underline{u}_{k}'\underline{x})) = cov(\underline{e}_{i}'\underline{x}, \underline{u}_{k}'\underline{x}) = \underline{e}_{i}'\Sigma u_{k} = \underline{u}_{k}'\Sigma e_{i} = \lambda_{i}\underline{u}_{k}'\underline{e}_{i} = \lambda_{i}e_{i,k}$$

$$\Rightarrow corr(\underline{y}_{i}, \underline{x}_{k}) = \frac{\lambda_{i}e_{i,k}}{\sqrt{\lambda_{i}}\sqrt{\sigma_{kk}}} = e_{i,k}\frac{\sqrt{\lambda_{i}}}{\sqrt{\sigma_{kk}}}$$

So we find that if σ_{kk} are of the same order of magnitude we can interpret component of loadings as related to the correlation.

Otherwise, if σ_{kk} have different order of magnitude the first component of the PCA take the component with the biggest variance (biggest as number not as interpretation).

We define V as a diagonal matrix with as elements of the main diagonal each variance, then we call standardize vector $\underline{Z} = V^{-\frac{1}{2}}(\underline{x} - \underline{\mu})$. It has $\mathbb{E}[\underline{Z}] = \underline{0}$ and $cov(\underline{Z}) = \underbrace{V^{-\frac{1}{2}}\Sigma(V^{-\frac{1}{2}})'}_{\rho}$

So we apply PCA to the covariance/correlation matrix $\underline{\rho} \Rightarrow \underline{y} = \underbrace{P'}_{\text{eigenvectors of } \rho} \underline{Z} = P'V^{-\frac{1}{2}}(\underline{x} - \underline{\mu})$

 $tr(\Lambda)=tr(
ho)=\sum_i \lambda_i=p \Rightarrow ext{rule of thumb:}$ select components with $\lambda_i\geq 1$

Remark: usually we estimate Σ with S so we apply PCA to eigenvectors and eigenvalues of S

A different perspective of PCA

Framework: our data are $\underline{x}_1, \dots, \underline{x}_n \in \mathbb{R}^p$

Question: find the linear subspace of dimension h in \mathbb{R}^p closest to our data

We want to find $\underline{\varphi}_1, \dots, \underline{\varphi}_k$ orthonormal s.t. $\mathcal{L} = span\{\underline{\varphi}_1, \dots, \underline{\varphi}_k\}$ is the closest to $\underline{x}_1, \dots, \underline{x}_n$

Algorithm's step:

- 1. Center our data: $\forall i \in \{1,\ldots,n\}$ $\underline{x}_1 \overline{\underline{x}},\ldots,\underline{x}_n \overline{\underline{x}}$
- 2. Project $\underline{x}_1, \dots, \underline{x}_n$ on $\mathcal{L} : \pi_{\underline{x}_i \mid \mathcal{L}} = \sum_{j=1}^k \underline{\varphi}_j \underline{\varphi}_j' (\underline{x}_i \overline{\underline{x}}) \quad \forall i$
- 3. Find \mathcal{L} s.t. $min\{\sum_{i=1}^n||(\underline{x}_i-\overline{\underline{x}})-\sum_{j=1}^k\underline{\varphi}_j\underline{\varphi}_j'(\underline{x}_i-\overline{\underline{x}})||^2\}$

How to find \mathcal{L} ?



$$\begin{split} \sum_{i=1}^{n} ||(\underline{x}_{i} - \overline{\underline{x}}) - \sum_{j=1}^{k} \underline{\varphi}_{j} \underline{\varphi}_{j}'(\underline{x}_{i} - \overline{\underline{x}})||^{2} &= \sum_{i=1}^{n} ||\underline{v}_{i} - \sum_{j=1}^{k} \underline{\varphi}_{j} \underline{\varphi}_{j}' \underline{v}_{i}||^{2} \\ &= \sum_{i=1}^{n} (\underline{v}_{i} - \sum_{j}^{k} \underline{\varphi}_{j} \underline{\varphi}_{j}' \underline{v}_{i})'(\underline{v}_{i} - \sum_{j}^{k} \underline{\varphi}_{j} \underline{\varphi}_{j}' \underline{v}_{i}) = \sum_{i=1}^{n} \underline{v}_{i} \underline{v}_{i}' - \sum_{i=1}^{n} \sum_{j=1}^{k} \underline{v}_{i}' \underline{\varphi}_{j} \underline{\varphi}_{j}' \underline{v}_{i} \\ &= \underline{v}_{i}' \underline{v}_{i} - 2 \sum_{j}^{k} \underline{v}_{i}' \underline{\varphi}_{j} \underline{\varphi}_{j}' \underline{v}_{i} + (\underbrace{\sum_{j} \underline{\varphi}_{j} \underline{\varphi}_{j}' \underline{v}_{i}})'(\underbrace{\sum_{t} \underline{\varphi}_{t} \underline{\varphi}_{t}' \underline{v}_{i}}) = \underline{v}_{i} \underline{v}_{i}' - \sum_{j=1}^{k} \underline{v}_{i}' \underline{\varphi}_{j} \underline{\varphi}_{j}' \underline{v}_{i} \\ &= \underline{0} \text{ if } \underline{j} \neq t \text{ else } = \sum_{j}^{k} \underline{v}_{i}' \underline{\varphi}_{i} \underline{\varphi}_{i}' \underline{v}_{i} \end{split}$$

So we reach that:

- $\sum_{i}^{n} \underline{v}_{i} \underline{v}_{i}'$ doesn't depend from \underline{arphi}_{j}
- $\sum_{i=1}^{n} \sum_{j=1}^{k} \underline{v}_{i}' \underline{\varphi}_{j} \underline{\varphi}'_{j} \underline{v}_{i}$ has to be maximized (since there is a minus before it)

Maximum of $\sum_{i=1}^{n}\sum_{j=1}^{k}\underline{v}_{i}'\underline{\varphi}_{j}\underline{\varphi}_{j}'\underline{v}_{i}$:

$$egin{aligned} \sum_{i}^{n} \sum_{j=1}^{k} \underline{v}_{i}' \underline{arphi}_{j} \underline{arphi}_{i}' &= \sum_{j=1}^{k} \sum_{i}^{n} \underline{arphi}_{j}' \underline{v}_{i} \underline{v}_{i}' \underline{arphi}_{j} &= \sum_{j=1}^{k} \underline{arphi}_{j}' (\sum_{i}^{n} \underline{v}_{i} \underline{v}_{i}') \underline{arphi}_{j} &= \\ &= \sum_{j=1}^{k} \underline{arphi}_{j}' (n-1) S \underline{arphi}_{j} &= (n-1) \sum_{j=1}^{k} \underline{arphi}_{j}' S \underline{arphi}_{j} \end{aligned}$$

Using a previous lemma we know that to maximize: k=1 $\underline{\varphi}_j=\underline{e}_1; \quad k=2$ $\underline{\varphi}_j=\underline{e}_2;\ldots$ So we find that $\mathcal{L}=span\{\underline{e}_1,\ldots,\underline{e}_k\}$ and the approximation error is: $(n+1)\sum_{i=k+1}^p \lambda_i$

Gaussian Model

Framework: $\mu \in \mathbb{R}^p; \Sigma = [\sigma_{i,j}] \in p imes p$ positive defined

Recall

$$p = 1$$
:

 $\mu\in\mathbb{R},\sigma_{1,1}>0\Rightarrow x_1\sim N_1(\mu,\sigma_{1,1}) ext{ if the density of } x_1 ext{ is }\phi(t)=rac{1}{\sqrt{2\pi\sigma_{1,1}}}exp\{-rac{(t-\mu)^2}{\sigma_{1,1}}\} ext{ with } t\in\mathbb{R}$ p
eq 1:

$$\underline{x} \sim N_p(\mu, \Sigma) \Leftrightarrow \phi(\underline{t}) = \frac{1}{\sqrt{(2\pi)^p det(\Sigma)}} exp\{-\frac{1}{2}(\underline{t} - \underline{\mu})'\Sigma^{-1}(\underline{t} - \underline{\mu})\}$$

Prop

$$\underline{x} \sim N_p(\mu, \Sigma) \Rightarrow orall \underline{a} \in \mathbb{R}^p \quad \underline{a}'\underline{x} \sim N_1(\underline{a}'\mu, \underline{a}'\Sigma\underline{a})$$

Corollary

$$\underline{x} \sim N_p(\underline{\mu}, \Sigma) \Rightarrow orall i \in \{1, \dots, p\} \quad x_i \sim N_1(\mu_i, \sigma_{i,i})$$

Proof - previous Corollary



$$ext{Take } \underline{u}_i = [0,\ldots,0,\overbrace{1}^{i^{th} ext{ position}},0,\ldots,0]' \stackrel{ ext{prev. prop}}{=\!=\!=\!=} \underline{u}_i \underline{x} \sim_1 (\underline{u}_i' \underline{\mu},\underline{u}_i' \Sigma \underline{u}_i) = N_1(\mu_i,\sigma_{i,i})$$

Corollary

$$egin{cases} rac{x}{A} \sim N_p(\underline{\mu}, \Sigma) \ A \in q imes p \end{cases} \, \Rightarrow A \underline{x} \sim N_q(A \underline{\mu}, A \Sigma A')$$

Proof

Thesis:
$$\forall \underline{a} \in \mathbb{R}^q \quad \underline{a}'(A\underline{x}) \sim N_1(\underline{a}'A\mu,\underline{a}'A\Sigma A'\underline{a})$$

$$\begin{array}{ll} \textit{Thesis:} \ \forall \underline{a} \in \mathbb{R}^q & \underline{a}'(A\underline{x}) \sim N_1(\underline{a}'A\underline{\mu},\underline{a}'A\Sigma A'\underline{a}) \\ \\ \underline{a}'(A\underline{x}) = (\underline{a}'A)\underline{x} = \overbrace{(A'\underline{a})'\underline{x}}^{\in \mathbb{R}^p} \sim N_1((A'\underline{a})'\underline{\mu},(A'\underline{a})'\Sigma (A'\underline{a})) = N_1(\underline{a}'A\underline{\mu},\underline{a}'A\Sigma A'\underline{a}) \end{array}$$

From that we conclude that:

$$\Rightarrow A\underline{x} \sim N_q(A\underline{\mu}, A\Sigma A')$$

Corollary

$$\underline{d} \in \mathbb{R}^p, \underline{x} \sim N(\mu, \Sigma) \quad \Rightarrow \quad \underline{x} + \underline{d} \sim N_p(\mu + \underline{d}, \Sigma)$$

Observation

$$Z_1, \ldots, Z_p \text{ iid } \sim N_1(0,1) \Rightarrow \underline{Z} \sim N_p(\underline{0},I)$$

So using the last observation and the previous corollaries we reach:

$$egin{aligned} \underline{x} \sim N_p(\underline{\mu}, \Sigma) &\Rightarrow \underline{Z} = \Sigma^{-rac{1}{2}}(\underline{x} - \underline{\mu}) \sim N_p(\underline{0}, I) \ \underline{Z} \sim N_p(\underline{0}, I) &\Rightarrow \underline{x} = \Sigma^{rac{1}{2}}\underline{Z} + \underline{\mu} \sim N_p(\underline{\mu}, \Sigma) \end{aligned}$$

Observation

Given $\underline{x} \sim N_p(\underline{\mu}, \Sigma)$ we can find the <code>#Mahalanobis_distance</code> W:

Given
$$\underline{x} \sim N_p(\underline{\mu}, \Sigma)$$
 we can find the `#Mahalanobis_distance` W :
$$W = d_{\Sigma^{-1}}^2(\underline{x}, \underline{\mu}) = (\underline{x} - \underline{\mu})'\Sigma^{-1}(\underline{x} - \underline{\mu}) = (\underline{x} - \underline{\mu})'\Sigma^{-\frac{1}{2}}\Sigma^{-\frac{1}{2}}(\underline{x} - \underline{\mu}) = \underline{Z}'\underline{Z} = \sum_i^p Z_i^2 \sim \chi^2(p)$$

$$\mathbb{P}[d_{\Sigma^{-1}}^2(\underline{x}, \underline{\mu}) \leq \chi_{1-\alpha}^2(p)] = 1 - \alpha$$

$$\mathbb{P}[d^2_{\Sigma^{-1}}(\underline{x},\underline{\mu}) \leq \chi^2_{1-lpha}(p)] = 1-lpha$$

Corollary

$$\mathsf{Take}\ \underline{x} \in \mathbb{R}^p \ \mathsf{compose}\ \mathsf{as}\ \underline{x} = \begin{bmatrix} \underline{x}_1 \in \mathbb{R}^q \\ \underline{x}_2 \in \mathbb{R}^{p-q} \end{bmatrix} \sim N_p(\begin{bmatrix} \underline{\mu}_1 \\ \underline{\mu}_2 \end{bmatrix}, \begin{bmatrix} \Sigma_{1,1} & \Sigma_{1,2} \\ \Sigma_{2,1} & \Sigma_{2,2} \end{bmatrix}) \Rightarrow \underline{x}_1 \sim N_q(\underline{\mu}, \Sigma_{1,1})$$

Proof -previous Corollary

Take
$$A=[I_{q imes q}\quad 0_{q imes (p-q)}]\in q imes p\Rightarrow A\underline{x}=\underline{x}_1\sim N_q(\underline{\mu}_1,A\Sigma A'=\Sigma_{1,1})$$
 \square

Prop



$$\underline{x} = egin{bmatrix} \underline{x}_1 \in \mathbb{R}^q \ \underline{x}_2 \in \mathbb{R}^{p-q} \end{bmatrix} \sim N_p(egin{bmatrix} \underline{\mu}_1 \ \underline{\mu}_2 \end{bmatrix}, egin{bmatrix} \Sigma_{1,1} & 0 \ 0 & \Sigma_{2,2} \end{bmatrix}) \Rightarrow \underline{x}_1 \perp \!\!\! \perp \underline{x}_2$$

Theo

$$\underline{x} = egin{bmatrix} \underline{x}_1 \in \mathbb{R}^q \ \underline{x}_2 \end{bmatrix} \in \mathbb{R}^q \sim N_p(egin{bmatrix} \underline{\mu}_1 \ \underline{\mu}_2 \end{bmatrix}, egin{bmatrix} \Sigma_{1,1} & \Sigma_{1,2} \ \Sigma_{2,1} & \Sigma_{2,2} \end{bmatrix}) \Rightarrow \underline{x}_1 | \underline{x}_2 \sim N_q(\underline{\mu}_1 + \Sigma_{1,2}\Sigma_{2,2}^{-1}(\underline{x}_2 - \underline{\mu}_2), \Sigma_{1,1} - \Sigma_{1,2}\Sigma_{2,2}^{-1}\Sigma_{2,1})$$

Proof - previous Theo

Remark: $\Sigma_{1,1} - \Sigma_{1,2} \Sigma_{2,2}^{-1} \Sigma_{2,1}$ is called partial covariance

Estimator of the mean and variance of a Gaussian

Framework: we have $\underline{x}_1,\ldots,\underline{x}_n \ \mathrm{iid} \ \sim N_p(\underline{\mu},\Sigma)$ with $\underline{\mu} \ \mathrm{and} \ \Sigma$ unknown.

Recall - 1

Estimator:

- $\overline{\underline{x}} = \frac{1}{n} \sum_{i=1}^{n} \underline{x}_i =$ sample mean
- $S=rac{1}{n-1}\sum_{i=1}^n(\underline{x}_i-\overline{\underline{x}})(\underline{x}_i-\overline{\underline{x}})'=\mathsf{sample}\;\Sigma$

Recall - 2

$$\begin{array}{l} \text{Likehood} = L(\underline{\mu}, \Sigma | \underline{x}_1, \dots, \underline{x}_n) = \prod_{i=1}^n \frac{1}{\sqrt{(2\pi)^p det(\Sigma)}} \exp(-\frac{1}{2}(\underline{x}_i - \underline{\mu})' \Sigma^{-1}(\underline{x}_i - \underline{\mu})) \\ \text{Log-Likehood} = l(\underline{\mu}, \Sigma | \underline{x}_1, \dots, \underline{x}_n) = n \cdot ln(\frac{1}{\sqrt{(2\pi)^p det(\Sigma)}}) - \frac{1}{2} \sum_{i}^n (\underline{x}_i - \underline{\mu})' \Sigma^{-1}(\underline{x}_i - \underline{\mu}) \end{array}$$

Interpretation: likelihood function $\mathcal{L}(\theta|\underline{x})$ is the probability of observing parameter θ assuming \underline{x} as data.

Theo

If
$$\underline{x}_1, \ldots, \underline{x}_n$$
 iid $\sim N_p(\mu, \Sigma)$

Then the Maximum Likehood Estimator (#MLE) estimator for:

- μ is $\hat{\mu} = \frac{1}{n} \sum_{i=1}^n \underline{x}_i$ which is unbiased;
- Σ is $\hat{\Sigma}=rac{1}{n}\sum_{i=1}^n(\underline{x}_i-\overline{\underline{x}})(\underline{x}_i-\overline{\underline{x}})'$ which is biased

Prop



If
$$\underline{x}_1,\dots,\underline{x}_n$$
 iid $\sim N_p(\underline{\mu},\Sigma)$
Then: $\overline{\underline{x}}\sim N_p(\mu,\frac{1}{n}\Sigma)$

Proof - previous Prop

Which is a squared identity matrix of dimension p one against others

$$egin{aligned} \overline{\underline{x}} = rac{1}{n}A ilde{\underline{x}} = egin{bmatrix} rac{1}{n}\sum_{i=1}^n x_{i,1} \ drawtriangle & drawtriangle & N_p(rac{1}{n}Aegin{bmatrix} rac{\mu}{drawtriangle} \ drawtriangle & drawtriangle & N_p(rac{1}{n}Aegin{bmatrix} rac{\mu}{drawtriangle} \ drawtriangle & drawtriangle & N_p(rac{1}{n}Aegin{bmatrix} rac{\mu}{drawtriangle} \ drawtriangle & drawtriangle & N_p(rac{\mu}{n}Aegin{bmatrix} rac{\mu}{drawtriangle} \ drawtriangle & N_p(rac{\mu}{n}Aegin{bmatrix} rac{\mu}{drawtriangle} \ drawtriangle & N_p(rac{\mu}{n}Aegin{bmatrix}
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#Wishart_distribution - introduced in 1928

Let $\underline{Z}_1,\ldots,\underline{Z}_n$ iid $\sim N_p(\underline{0},\Sigma)\Rightarrow A=\sum_{i=1}^n\underline{Z}_i\underline{Z}_i'\sim Wish(\Sigma,m)$ where m is the number of independent addends, in this case m=n.

Spoiler: it's a multidimensional χ^2 .

Prop - 1

 $\mathsf{Suppose}\ A_1 \sim Wish(\Sigma, m_1), A_2 \sim Wish(\Sigma, m_2), A_1 \perp\!\!\!\perp A_2 \Rightarrow A_1 + A_2 \sim Wish(\Sigma, m_1 + m_2)$

Proof - previous Prop

$$A_1 = \sum_{i=1}^{m_1} oldsymbol{\underline{Z}}_i oldsymbol{\underline{Z}}_i^\prime; \quad A_2 = \sum_{i=1}^{m_2} oldsymbol{\underline{\widetilde{Z}}}_i oldsymbol{\underline{\widetilde{Z}}}_i^\prime \quad ext{with } oldsymbol{\underline{Z}}_i, oldsymbol{\underline{\widetilde{Z}}}_j \sim N_p(oldsymbol{\underline{0}}, \Sigma)$$

 $\Rightarrow ext{we call } \underline{Z}_i ext{ as } \underline{w}_{j \in \{1, \dots, m_1\}} ext{ and } \underline{\tilde{Z}}_i ext{ as } \underline{w}_{j \in \{m_1, \dots, m_1 + m_2\}}$

$$A_1 + A_2 = \sum_{i=1}^{m_1+m_2} \underline{w}_i \underline{w}_i' \sim Wish(\Sigma, m_1+m_2)$$

Prop - 2

 $A \sim Wish(\Sigma,m); C \in k imes p \Rightarrow CAC' \sim Wish(C\Sigma C',m)$

Proof - previous Prop



$$A = \sum_{i=1}^m \underline{Z}_i \underline{Z}_i' \quad ext{with } \underline{Z}_i \sim N_p(\underline{0}, \Sigma) ext{ iid } \ CAC' = \sum_{i=1}^m \underbrace{CZ_i \underline{Z}_i'C}_{\underline{w}_i} ext{with } \underline{w}_i \sim N_k(\underline{0}, C\Sigma C') \Rightarrow CAC' \sim Wish(C\Sigma C', m)$$

Prop - 3

$$A \sim Wish(\Sigma,m); \sigma^2 \in \mathbb{R}^+ ackslash \{0\} \Rightarrow \sigma^2 A \sim Wish(\sigma^2 \Sigma,m)$$

Proof - previous Prop

$$\sigma^2 A = \sigma^2 \sum_{i=1}^m \underline{Z_i} \underline{Z_i'} = \sum_{i=1}^m \underbrace{\sigma \underline{Z_i} \sigma \underline{Z_i'}}_{\underline{w_i'}} ext{with } \underline{w}_i \sim N_p(\underline{0}, \sigma^2 \Sigma) \Rightarrow \sigma^2 A \sim Wish(\sigma^2 \Sigma, m)$$

Prop - 4

Let
$$p=1\Rightarrow Wish(\overbrace{\Sigma}^{\sigma^2},m)=\sigma^2\chi^2(m)$$

Proof - previous Prop

If
$$p=1 o \Sigma=\sigma^2\Rightarrow rac{1}{\sigma^2}A=\sum_i^m(rac{Z_i}{\sigma})^2$$
 with $rac{Z_i}{\sigma}\sim N_1(0,1)$ $\Rightarrow rac{1}{\sigma^2}A\sim \chi^2(m)$

Theo

Take
$$\underline{x}_1,\ldots,\underline{x}_n$$
 iid $\sim N_p(\mu,\Sigma)$

Then:
$$\sum_{i}^{n}(\underline{x}_{i}-\overline{\underline{x}})(\underline{x}_{i}-\overline{\underline{x}})'\sim Wish(\Sigma,n-1)$$

Note: The second parameter is n-1 and not n since we are removing the subspace of the mean $span\{\underline{1}\}$.

Corollary

Take
$$\underline{x}_1, \ldots, \underline{x}_n$$
 iid $\sim N_p(\mu, \Sigma)$

Then:
$$\hat{\Sigma} = \frac{1}{n} \sum_{i}^{n} (\underline{x}_{i} - \overline{\underline{x}}) (\underline{x}_{i} - \overline{\underline{x}})' \sim Wish(\frac{1}{n}\Sigma, n-1) \Rightarrow S \sim Wish(\frac{1}{n-1}\Sigma, n-1)$$

Theo

Take
$$\underline{x}_1,\ldots,\underline{x}_n$$
 iid $\sim N_p(\mu,\Sigma)$

Then:

1.
$$\underline{\overline{x}} \sim N_p(\mu, \frac{1}{n}\Sigma)$$

2.
$$(n-1)S \sim Wish(\Sigma, n-1)$$

3.
$$\overline{\underline{x}} \perp \!\!\! \perp S$$

Recall - Central Limit Theorem #CLN

Take $\underline{x}_1,\ldots,\underline{x}_n$ iid $\sim F(\underline{\mu},\Sigma)$ with F as generic distribution

Then:



$$ullet \sqrt{n}(\overline{x}-\mu) \sim AN_p(0,\Sigma)$$

• if
$$n o +\infty$$
 $\overline{\underline{x}} \sim N_p(\mu, rac{1}{n}\Sigma)$

IMPORTANT:

It's importante to notice that in this theorem is involed the sample mean!!

Recall - Law Large Number #LNN

Take
$$\underline{x}_1, \dots, \underline{x}_n$$
 iid $\sim F(\mu, \Sigma)$

$$\underline{\overline{x}} \overset{\mathbb{P}}{ o} \mu, \quad S \overset{\mathbb{P}}{ o} \Sigma \quad ext{ as } n o + \infty$$

Inference for the mean

n >> p

Rule of Thumb: is valid if $n > 30p^2$

Framework: Take $\underline{x}_1,\dots,\underline{x}_n$ $\mathrm{iid}\ \sim F(\underline{\mu},\Sigma)$ with F as generic distribution

By the #CLN we know that:

$$\sqrt{n}(\overline{\underline{x}}-\mu) \sim AN_p(\overline{\underline{0}},\Sigma)
ightarrow \overline{\underline{x}} \sim AN_p(\mu, rac{1}{n}\Sigma)$$

From this we can say, approximating, that:

$$(\overline{\underline{x}}-\underline{\mu})'(\frac{1}{n}\Sigma)^{-1}(\overline{\underline{x}}-\underline{\mu}) = \boxed{n(\overline{\underline{x}}-\underline{\mu})'\Sigma^{-1}(\overline{\underline{x}}-\underline{\mu}) \sim \chi^2(p)}$$

Rmk: if Σ is known it is a pivotal quantity for $\underline{\mu}$

If we put $n \to +\infty$ we can say that:

$$S \stackrel{\mathbb{P}}{\underset{LLN}{\longrightarrow}} \Sigma \Rightarrow n(\overline{\underline{x}} - \underline{\mu})' S^{-1}(\overline{\underline{x}} - \underline{\mu}) = d^2_{(rac{1}{n}S)^{-1}}(\overline{\underline{x}}, \underline{\mu}) \sim \chi^2(p)$$

which is pivotal quantity of μ so:

$$\mathbb{P}[d^2_{(rac{1}{2}S)^{-1}}(\overline{\underline{x}},\underline{\mu}) \leq \chi^2_{1-lpha}(p)] = 1-lpha$$

We can create 2 ellipsoid:

$$\bullet \ \ \epsilon^{\alpha}_{\chi^{2}_{1-\alpha}(p)}(\underline{\mu}) = \{\underline{x} \in \mathbb{R}^{p} : d^{2}_{(\frac{1}{n}S)^{-1}}(\underline{x},\underline{\mu}) \leq \chi^{2}_{1-\alpha}(p)\}$$

$$\bullet \ \ \epsilon^{\alpha}_{\chi^{2}_{1-\alpha}(p)}(\overline{\underline{x}}) = \{\underline{\eta} \in \mathbb{R}^{p} : d^{2}_{(\frac{1}{n}S)^{-1}}(\underline{\eta}, \overline{\underline{x}}) \leq \chi^{2}_{1-\alpha}(p)\}$$

So we can see, geometrically, as: $\mathbb{P}[\overline{\underline{x}} \in \epsilon^{\alpha}_{\chi^2_{1-\alpha}(p)}(\underline{\mu})] = 1 - \alpha = \mathbb{P}[\underline{\mu} \in \epsilon^{\alpha}_{\chi^2_{1-\alpha}(p)}(\overline{\underline{x}})]$

From this we can create the confidence region of μ at level $1-\alpha$:

$$\boxed{\text{Hotelling statistic} = T_0^2 = n(\overline{\underline{x}} - \underline{\mu}_0)'S^{-1}(\overline{\underline{x}} - \underline{\mu}_0) \quad \Rightarrow \quad CR_{1-\alpha}(\underline{\mu}) = \{\underline{\eta} \in \mathbb{R}^p : T_0^2 \leq \chi_{1-\alpha}^2(p)\}}$$

where $\underline{\mu}_0$ is the hypothesis H_0 .

We reject H_0 if $T_0^2 > \chi_{1-\alpha}^2(p)$ if we have a confidence level α .

Otherwise, instead of fixing α we can evaluate p-value so at level α

we reject our H_0 if p-value $< \alpha$

Note: our $CR_{1-\alpha}(\underline{\mu})$ identifies values of $\underline{\mu}_0$ for which we cannot reject H_0



$n \approx p$

Framework: Take $\underline{x}_1, \dots, \underline{x}_n \ \mathrm{iid} \ \sim N_p(\mu, \Sigma)$

Rmk: we need Gaussianity hypothesis since it is the cost of not having so much data

#Fisher_distribution - introduced in 1928

Let
$$Y \sim \chi^2(n), \quad W \sim \chi^2(m), \quad Y \perp\!\!\!\perp W$$

Then:
$$rac{rac{Y}{n}}{rac{W}{m}} \sim F(n,m)$$

Rmk:
$$t \stackrel{^m}{=} rac{Z}{\sqrt{rac{W}{m}}} ext{ with } Z \sim N(0,1), W \sim \chi^2(m), Z \perp \!\!\! \perp W \Rightarrow t^2 \sim F(1,m)$$

Rmk:
$$F(n,m) \xrightarrow{m \to +\infty} rac{1}{n} \chi^2(n)$$
 since $rac{W}{m} = rac{1}{m} \sum_i^m Z_i^2 \xrightarrow{LLN} 1$ with $Z_i \sim N(0,1)$

Theo - Hotelling (1931)

Assume:
$$\underline{x} \sim N_p(\mu, \Sigma)$$
 with $det(\Sigma) > 0$; $W \sim Wish(\Sigma, m)$ with $W \perp \!\!\! \perp \underline{x}$

Then:
$$rac{m-p+1}{mp}(\underline{x}-\underline{\mu})'W^{-1}(\underline{x}-\underline{\mu})\sim F(p,m-p+1)$$

Corollary

$$\underline{x}_1,\dots,\underline{x}_n ext{ iid } \sim N_p(\underline{\mu},\Sigma) \Rightarrow \boxed{n(\overline{\underline{x}}-\underline{\mu})'S^{-1}(\overline{\underline{x}}-\underline{\mu}) = d_{(\frac{1}{n}S)^{-1}}^2(\overline{\underline{x}},\underline{\mu}) \sim rac{n-1}{n-p}p \cdot F(p,n-p)}$$

Rmk: is a pivotal quantity

Proof - previous Corollary

$$egin{aligned} \sqrt{n}(\overline{\underline{x}}-\underline{\mu}) &\sim N_p(\underline{0},\Sigma); \quad (n-1)S \sim Wish(\Sigma,n-1); \quad \sqrt{n}(\overline{\underline{x}}-\underline{\mu}) \perp\!\!\!\perp S \ \Rightarrow n(\overline{\underline{x}}-\underline{\mu})'S^{-1}(\overline{\underline{x}}-\underline{\mu}) &= d_{rac{1}{n}S^{-1}}^2(\overline{\underline{x}},\underline{\mu}) \sim rac{n-1}{n-p}p\cdot F(p,n-p) \end{aligned}$$

Fixed
$$\alpha \in [0,1]$$
 $CR_{1-\alpha}(\underline{\mu}) = \{\underline{\eta} \in \mathbb{R}^p : n(\overline{\underline{x}} - \underline{\mu})'S^{-1}(\overline{\underline{x}} - \underline{\mu}) \leq \frac{n-1}{n-p}p \cdot F_{1-\alpha}(p,n-p)\}$

So in a test of hypothesis we reject H_0 if $T_0^2 > rac{n-1}{n-p} p \cdot F_{1-lpha}(p,n-p)$

Rmk:

Squared radius of the ellipse:

• n large:
$$\chi^2_{1-\alpha}(p)$$
;

• n small:
$$\frac{n-1}{n-p}p\cdot F_{1-lpha}(p,n-p)$$

Since:
$$\frac{n-1}{n-p}p\cdot F_{1-lpha}(p,n-p) \xrightarrow{n o +\infty} \chi^2_{1-lpha}(p)$$

Inference for linear combination of μ

Framework: we have $\underline{x}_1, \ldots, \underline{x}_n$ iid $\sim N_p(\mu, \Sigma)$

 $\textbf{Goal:} \ \text{we want to do inference on linear combination of } \mu$

Take a generic $\underline{a} \in \mathbb{R}^p \Rightarrow \text{ the estimator of } \underline{a}'\mu \text{ is } \underline{a}'\overline{\underline{x}}$

Because we know that:



$$1.\ \underline{a}' \overline{\underline{x}} \sim N_1(\underline{a}' \underline{\mu}, \frac{1}{n} \underline{a}' \Sigma \underline{a}) \Leftrightarrow \underbrace{\frac{\sqrt{n}(\underline{a}' \overline{\underline{x}} - \underline{a}' \underline{\mu})}{\sqrt{\underline{a}' \Sigma \underline{a}}}}_{\mathbb{R}} \sim N_1(0, 1)$$

2. Starting from
$$(n-1)S \sim Wish(\Sigma, n-1)$$
 we can find:
$$\Rightarrow (n-1)\underline{a}'S\underline{a} \sim Wish(\underline{\underline{a}'\Sigma\underline{a}}, n-1) = (\underline{a}'\Sigma\underline{a})\chi^2(n-1)$$

$$\Rightarrow \underbrace{\frac{(n-1)\underline{a}'S\underline{a}}{(\underline{a}'\Sigma\underline{a})}} \sim \chi^2(n-1)$$

Since
$$(\mathbb{R} \perp \!\!\! \perp \mathbb{S}) \Rightarrow \frac{(\mathbb{R})}{\sqrt{\frac{(\mathbb{S})}{n-1}}} \sim t(n-1)$$

$$\Rightarrow \boxed{rac{\sqrt{n}(\underline{a}'\overline{\underline{x}}-\underline{a}'\underline{\mu})}{\sqrt{\underline{a}'S\underline{a}}} \sim t(n-1) ext{ which is a pivotal quantity of } \underline{a}'\underline{\mu}}.$$

Fixed $\alpha \in [0.$

$$\mathbb{P}[\frac{\sqrt{n}|\underline{a'}\overline{x}-\underline{a'}\underline{\mu}|}{\sqrt{\underline{a'}S\underline{a}}} < t_{1-\frac{\alpha}{2}}(n-1)] = 1-\alpha \Leftrightarrow \mathbb{P}[|\underline{a'}\overline{x}-\underline{a'}\underline{\mu}| < t_{1-\frac{\alpha}{2}}(n-1)\sqrt{\frac{\underline{a'}S\underline{a}}{n}}] = 1-\alpha$$

So we can create:

- Confidence Interval $=CI_{1-lpha}(\underline{a}'\underline{\mu})=[\underline{a}'\overline{\underline{x}}\pm t_{1-rac{lpha}{2}}(n-1)\sqrt{rac{\underline{a}'}{2}}$
- Hypothesis tests with as testing statistics $t_0 = \frac{\sqrt{n}(\underline{a}'\overline{x} \delta_0)}{\sqrt{a'Sa}}$;

For hypothesis tests we can deal with:

1.
$$H_0: \underline{a}'\mu \leq \delta_0 \text{ vs } H_1: \underline{a}'\mu \geq \delta_0 \quad \Rightarrow \quad \text{Reject } H_0 \text{ if } t_0 > t_{1-\alpha}(n-1)$$

1.
$$H_0: \underline{a'}\underline{\mu} \leq \delta_0 \text{ vs } H_1: \underline{a'}\underline{\mu} \geq \delta_0 \quad \Rightarrow \quad \text{Reject } H_0 \text{ if } t_0 > t_{1-\alpha}(n-1)$$
2. $H_0: \underline{a'}\underline{\mu} = \delta_0 \text{ vs } H_1: \underline{a'}\underline{\mu} \neq \delta_0 \quad \Rightarrow \quad \text{Reject } H_0 \text{ if } |t_0| > t_{1-\frac{\alpha}{2}}(n-1)$

Prop

Take $\alpha \in [0,1]$:

$$\forall \underline{a} \in \mathbb{R}^p \quad CI_{1-\alpha}(\underline{a'}\underline{\mu}) = [\underline{a'}\underline{\overline{x}} + t_{1-\frac{\alpha}{2}}(n-1)\sqrt{\frac{\underline{a'Sa}}{n}}] \quad \Leftrightarrow \quad \forall \underline{a} \in \mathbb{R}^p \quad \mathbb{P}[\underline{a'}\underline{\mu} \in CI_{1-\alpha}(\underline{a'}\underline{\mu})] = 1-\alpha$$

$$\textit{Rmk:} \text{ which is not equal to } \mathbb{P}[\underline{a'}\underline{\mu} \in CI_{1-\alpha}(\underline{a'}\underline{\mu}), \quad \forall \underline{a} \in \mathbb{R}^p] = 1-\alpha$$

Recall

Take $\underline{b},\underline{d} \in \mathbb{R}^p$ we know that $\cos(\theta) = \frac{\underline{b'd}}{||\underline{b}|||\underline{d}||}$ so we deduce that $(\underline{b'd})^2 \leq ||\underline{b}||^2||\underline{d}||^2$

Theo - Extended Cauchy-Schwartz

Take $B \in p \times p$ positive defined and symmetric.

Then: $\forall b, d \in \mathbb{R}^p \text{ holds } (b'd)^2 \leq (b'Bb)(d'B^{-1}d)$

Rmk: the equality holds if $\underline{b} \in \mathcal{L}(B^{-1}\underline{d})$

Proof - previous Theo

$$(\underline{b'}\underline{d})^2 = (\underline{b'}B^{\frac{1}{2}}B^{-\frac{1}{2}}\underline{d})^2 \leq ||B^{\frac{1}{2}}\underline{b}||^2||B^{-\frac{1}{2}}\underline{d}||^2 = (\underline{b'}B^{\frac{1}{2}}B^{\frac{1}{2}}\underline{b})(\underline{d'}B^{-\frac{1}{2}}B^{-\frac{1}{2}}\underline{d}) = (\underline{b'}B\underline{b})(\underline{d'}B^{-1}\underline{d})$$



Lemma

Take $B \in p imes p$ positive defined and symmetric and $\underline{d} \in \mathbb{R}^p$

Then:
$$\max_{\underline{x}\in\mathbb{R}^p,||\underline{x}||
eq 0} rac{(\underline{x}'\underline{d})^2}{\underline{x}'B\underline{x}} = \underline{d}'B^{-1}\underline{d}$$

Proof - previous Lemma

$$(\underline{x}'\underline{d})^2 \le (\underline{x}'B\underline{x})(\underline{d}'B^{-1}\underline{d}) \text{ by Cauchy-Schwartz } \Rightarrow \frac{(\underline{x}'\underline{d})^2}{\underline{x}'B\underline{x}} \le \underline{d}'B^{-1}\underline{d} \quad \forall \underline{x} \ne \underline{0}$$

$$\Rightarrow \max_{\underline{x} \in \mathbb{R}^p, ||\underline{x}|| \ne 0} \frac{(\underline{x}'\underline{d})^2}{\underline{x}'B\underline{x}} = \underline{d}'B^{-1}\underline{d}$$

So we find that:

$$\max_{\underline{a}\in\mathbb{R}^p, ||\underline{a}||
eq 0} nrac{[\underline{a}'(\overline{\underline{x}}-\underline{\mu})]^2}{a'Sa} = n(\overline{\underline{x}}-\underline{\mu})'S^{-1}(\overline{\underline{x}}-\underline{\mu}) \sim prac{n-1}{n-p}F(p,n-p)$$

This is helpful in addressing the problem of resolving $\mathbb{P}[\underline{a'}\underline{\mu} \in CI_{1-\alpha}(\underline{a'}\underline{\mu}), \quad \forall \underline{a} \in \mathbb{R}^p] = 1 - \alpha$ because:

$$\mathbb{P}[\underline{a'}\underline{\mu} \in CI_{1-lpha}(\underline{a'}\underline{\mu}), \quad orall \underline{a} \in \mathbb{R}^p] = 1 - lpha \overset{ ext{as before}}{\Longleftrightarrow} \mathbb{P}[nrac{[\underline{a'}\overline{x} - \underline{a'}\underline{\mu}]^2}{a'Sa} \leq \overbrace{c^2}^{unknown} = c, \quad orall \underline{a} \in \mathbb{R}^p]$$

Finding c is the same of maximize $n \frac{[\underline{a'}(\overline{x} - \underline{\mu})]^2}{\underline{a'}S\underline{a}}$ so we reach that

 $CI_{1-lpha}(\underline{a'}\underline{\mu}) = [\underline{a'}\overline{\underline{x}} \pm \sqrt{prac{n-1}{n-p}}F_{1-lpha}(p,n-p)\cdot\sqrt{rac{\underline{a'}S\underline{a}}{n}}]$ and we have finally achieve:

$$\mathbb{P}[\underline{a}'\mu \in CI_{1-lpha}(\underline{a}'\mu), \quad orall \underline{a} \in \mathbb{R}^p] = 1-lpha$$

To summarize:

$$- ext{ t-test (one at time): } \hspace{0.2cm} orall \underline{a} \in \mathbb{R}^p \hspace{0.2cm} CI_{1-lpha}(\underline{a}'\underline{\mu}) = [\underline{a}'\overline{\underline{x}} + t_{1-rac{lpha}{2}}(n-1)\sqrt{rac{\underline{a}'S\underline{a}}{n}}]$$

$$- ext{ F-test (multiple at time): } CI_{1-lpha}(\underline{a}'\underline{\mu}) = [\underline{a}'\overline{\underline{x}} \pm \sqrt{prac{n-1}{n-p}}F_{1-lpha}(p,n-p)\cdot \sqrt{rac{\underline{a}'S\underline{a}}{n}} \quad orall \underline{a} \in \mathbb{R}^p]$$

Interpretation: in the first case we want to have confidence values of an interval in \mathbb{R} , in the second case we want to have confidence values of each possible direction (like an area in \mathbb{R}^p) so caused by #curse_of_dimensionality we need to enlarge the interval to keep the same confidence level.

Geometrically: in the first case you are projecting the elliptic $CI_{1-\alpha}(\underline{a'\mu})$ on one direction (which is \underline{a}) and this projection is called shadow.

Q: How to not enlarge our CI as in F-test?



R: Bonferroni solution

Goal: we want to take k tests simultaneously and also keep "small" our CI

Take $\underline{a}_1,\ldots,\underline{a}_k\in\mathbb{R}^p$ to test; which is a correct $\beta\in[0,1]$ s.t. $\mathbb{P}\{\cap_{i=1}^k[\underline{a}_i'\underline{\mu}\in CI_{1-\beta}(\underline{a}_i'\underline{\mu})]\}=1-\alpha$ with $CI_{1-\beta}(\underline{a}_i'\mu)$ t-tests?

Bonferroni Idea: $\beta = \frac{\alpha}{k}$

We can arrive to this with simple calculations:

$$\begin{split} \mathbb{P}\{\cap_{i=1}^{k}[\underline{a_{i}'\underline{\mu}} \in CI_{1-\beta}(\underline{a_{i}'\underline{\mu}})]\} &= 1 - \alpha = \\ &= 1 - \mathbb{P}\{\cup_{i=1}^{k}[\underline{a_{i}'\underline{\mu}} \not\in CI_{1-\beta}(\underline{a_{i}'\underline{\mu}})]\} = \\ &= 1 - \sum_{i=1}^{k} \underbrace{\mathbb{P}\{\underline{a_{i}'\underline{\mu}} \not\in CI_{1-\beta}(\underline{a_{i}'\underline{\mu}})\}}_{=\beta} = \\ &= 1 - k\beta \\ &\Rightarrow \alpha = k\beta \end{split}$$

From this we can create Bonferroni Confidence Interval (one at time for each \underline{a}_i):

$$BCI_{1-rac{lpha}{k}}(\underline{a}_i'\underline{\mu})=[\underline{a}_i'\overline{\underline{x}}+t_{1-rac{lpha}{2k}}(n-1)\sqrt{rac{\underline{a}_i'S\underline{a}_i}{n}}]$$

So we can also expand this theory to testing:

$$H_0 = egin{cases} rac{lpha_1' \mu = \delta_1}{lpha_2' \mu = \delta_2} \ dots \ rac{lpha_2' \mu = \delta_2}{dots} \end{cases} ext{ vs } H_1 = ext{at least 1 fails} \ rac{lpha_k' \mu = \delta_k} \end{cases}$$

So we reject H_0 at level lpha if for at least one $i\in\{1,\dots,k\}$ $\frac{\sqrt{n}|\underline{a_i'}\overline{x}-\delta_i|}{\sqrt{\underline{a_i'}Sa_i}}>t_{1-\frac{lpha}{2k}}(n-1)$

Taking as p_i the p-value of the test $H_{0,i}$ vs $H_{1,i}$ we know that we reject $H_{0,i}$ if $p_i < \dfrac{\alpha}{k}$.

So we find:

$$egin{aligned} \mathbb{P}[ext{Reject } H_0|H_0 ext{ holds}] &= \mathbb{P}[\cup_{i=1}^k \{ ext{Reject } H_{0,i}\}| \cap_{i=1}^k H_{0,i}] \ &\leq \sum_{i=1}^k \mathbb{P}[ext{Reject } H_{0,i}|H_{0,i} ext{ holds}] = \sum_{i=1}^k rac{lpha}{k} = lpha \end{aligned}$$

Large scale hypothesis testing and FDR

The solution reported above has a huge limit, if $k \uparrow$ then $\beta \to 0$. This is a problem in cases like large scale hypothesis testing, for example in genomics.

Solution: False Discovery Rate (#FDR) by Benjamini&Hochberg (1995)

We want to test k hypothesis with a strategy D (for example Bonferroni test). We can construct this table:



Where:

- *V* = false discoveries;
- T =missed discoveries;
- *S* = true discoveries;
- R = rejections (which are the only things observable)

We create $I_0 = \{i \in \{1, \dots, k\} : H_{0,i} \text{ holds}\}.$

In the following calculation we use Bonferroni as strategy D:

$$egin{aligned} \mathbb{P}[V \geq 1] &= \mathbb{P}[ext{ at least 1 false discovery}] = \mathbb{P}[\cup_{j \in I_0} \{ ext{Reject } H_{0,j}\} | \cap_{j \in I_0} H_{0,j}] \ &\leq \sum_{j \in I_0} \mathbb{P}[ext{Reject } H_{0,j} | H_{0,j} ext{ holds}] \ &= \sum_{j \in I_0} rac{lpha}{k} = rac{k_0}{k} lpha \ &\leq lpha \end{aligned}$$

The previous probability is called *Family-Wise Error Rate* (#FWER) and for Bonferroni hold $FWER < \alpha$.

So we can define the proportion of false discoveries among discoveries $=Q=\begin{cases} 0 & R=0 \\ \frac{V}{R} & R>0 \end{cases}$ From above we create the *False Discovery Rate* $=FDR=\mathbb{E}[Q]$

Observation - 1 ($k_0 = k$)

We don't have any discovery so $Q = egin{cases} 0 & V = 0 \\ 1 & V > 0 \end{cases} \Rightarrow FDR = FWER$

Observation - 2 ($k_0 < k$)

$$Q = \begin{cases} 0 & V = 0 \\ \frac{V}{R} \leq 1 & V > 0 \end{cases} \Rightarrow Q \leq \mathbb{1}_{V > 0} \Rightarrow FDR = \mathbb{E}[Q] \leq \mathbb{E}[\mathbb{1}_{V > 0}] = \mathbb{P}[V > 0] = FWER$$

So we reach that controlling FDR is less restricting that controlling FWER.

How to control FDR?

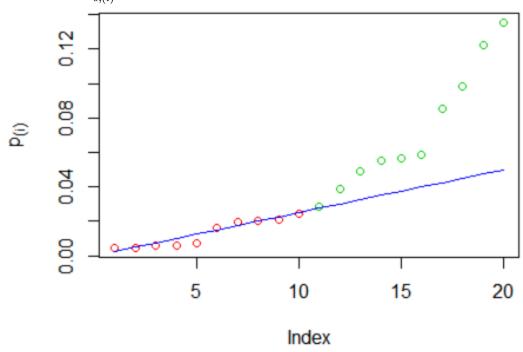
Framework: p_i is the p-value of $H_{0,i}$ vs $H_{1,i}$

- 1. Ordering the p-value such that the biggest one is $p_{(1)}$ while the smallest one is $p_{(k)}$.
- 2. Taking $\alpha \in (0,1)$, find m which is the greatest index s.t. $p_{(m)} < m \frac{\alpha}{k}$, same as: $m = \arg\max_{i \in \{1,\ldots,k\}} \{p_{(i)} \le i \frac{\alpha}{k}\}$
- 3. Reject $H_{0,(1)}, \ldots, H_{0,(m)}$

In the following picture we can see that the blue line is $i\frac{\alpha}{k}$ while the point are p_i . The Bonferroni threshold will be a horizontal line at level $\frac{\alpha}{k}$ so if k increase we are subject to accepting more



and more $H_{0,(i)}$.



Theo

If p_i, \ldots, p_k are independent, this strategy control FDR at level α .

Rmk: each p-value p_i is a function of $\underline{x}_1,\dots,\underline{x}_n$

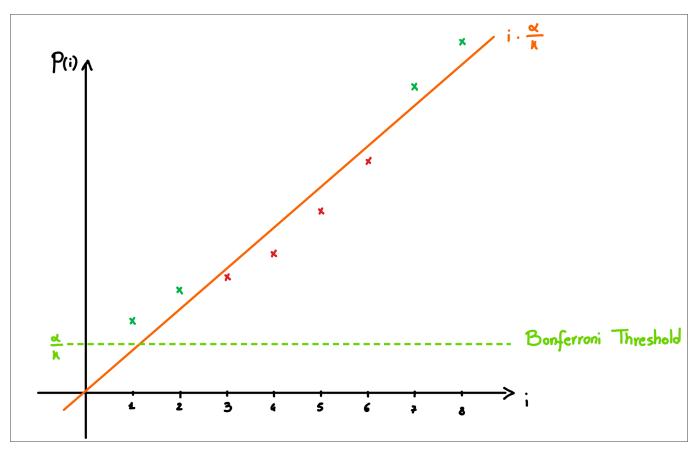
Q: What if p_i aren't independent?

Theo - Benjamini ad Yekuteli (2001)

- p_1, \ldots, p_k are positively correlated \Rightarrow holds B&H (1995) strategy
- p_1,\ldots,p_k are negatively correlated $\Rightarrow m^*=rg\max_{i\in\{1,\ldots,k\}}\{p_i\leq irac{lpha}{k\cdot C(k)}\}$ and $C(k)=\sum_{j=1}^krac{i}{j}$

Observation





In a situation like this, how different strategies works:

- Benjamini&Hochberg: Since m=6 reject $H_{0,(1)},\dots,H_{0,(6)}$
- **Efron (2010):** which is a strong version of B&H, reject $H_{0,(i)}$ if $p_{(i)} \leq i \frac{\alpha}{k}$ so reject $H_{0,(3)},\ldots,H_{0,(6)}$
- Bonferroni: doesn't reject

Comparing means of multivariate Gaussian Distr.

Framework:

We have paired data of n statistical units observed twice: $\underline{x}_{1,i}, \underline{x}_{2,i} \in \mathbb{R}^p$ with $i \in \{1, \dots, n\}$ So we have:

$$egin{pmatrix} \left(\underline{x}_{1,1} \\ \underline{x}_{2,1} \end{pmatrix}, \left(\underline{x}_{1,2} \\ \underline{x}_{2,2} \end{pmatrix}, \ldots, \left(\underline{x}_{1,n} \\ \underline{x}_{2,n} \end{pmatrix} ext{ iid } \sim N_{2p}(egin{pmatrix} \underline{\mu}_1 \\ \underline{\mu}_2 \end{pmatrix}, \overbrace{\Sigma} \end{pmatrix}$$

Rmk: independence is between vectors not between components.

Problem to solve: $H_0: \underline{\mu}_1 - \underline{\mu}_2 = \underline{\delta} \quad {
m vs} \quad H_1: \underline{\mu}_1 - \underline{\mu}_2
eq \underline{\delta}$

We create:

$$\underline{D}_i \in \mathbb{R}^p ext{ s.t. } \underline{D}_i = \underline{x}_{1,i} - \underline{x}_{2,i} ext{ iid } \sim N_p(\underline{\delta}, \Sigma_D)$$

$$\Rightarrow \overline{\underline{D}} = \frac{1}{n} \sum_{i=1}^n \underline{D}_i \text{ and } S_D = \frac{1}{n-1} \sum_{i=1}^n (\underline{D}_i - \overline{\underline{D}}) (\underline{D}_i - \overline{\underline{D}})'$$

So we reach that: $n(\overline{\underline{D}} - \underline{\delta})'S^{-1}(\overline{\underline{D}} - \underline{\delta}) \sim p \frac{n-1}{n-p} F(p,n-p)$ which is a pivotal quantity and

useful to conclude, using theory developed in previous chapters, that taking $\alpha \in (0,1)$:



$$CR_{1-lpha}(\underline{\mu}_1-\underline{\mu}_2)=CR_{1-lpha}(\underline{\delta})=\{\underline{\delta}\in\mathbb{R}^p:n(\overline{\underline{D}}-\underline{\delta})'S^{-1}(\overline{\underline{D}}-\underline{\delta})\leq prac{n-1}{n-p}F_{1-lpha}(p,n-p)\}$$
 $CR_{1-lpha}(\underline{a}'\underline{\delta})=\{\underline{a}'\overline{\underline{D}}\pm\sqrt{prac{n-1}{n-p}F_{1-lpha}(p,n-p)}\sqrt{rac{\underline{a}'S_D\underline{a}}{n}}\}$

Univariate case

Framework: we repeat q times each measure.

Take $\underline{x}_1, \ldots, \underline{x}_n$ iid $\sim N_q(\mu, \Sigma)$ so each

 $\underline{x}_i = \left[x_{i,1}, \dots, x_{i,q}\right]' \in \mathbb{R}^q \leftarrow q \text{ measurements of the same quantity}$

Goal: test $H_0: \mu_1 = \cdots = \mu_q \quad ext{ vs } \quad H_1: \exists i, j ext{ s.t. } \mu_i
eq \mu_j$

We can define the $\#Contrast_matrix$ C as:

 $C \in (q-1) imes q$ is a contrast matrix if $C = egin{bmatrix} \underline{c}_1' \\ \vdots \\ \underline{c}_{q-1}' \end{bmatrix}$ with $\underline{c}_1, \dots, \underline{c}_{q-1}$ are linearly independent and

 $\forall i \quad \underline{c}_i' \underline{1} = 0.$

We can rewrite this definition as:

$$oxed{C ext{ is a contrast matrix if } \mathcal{L}^{\perp}(\underline{1}) = span(\underline{c}_1, \dots, \underline{c}_{q-1})}$$

So we can rewrite our test as: $H_0: C\mu = \underline{0} \quad ext{ vs } \quad H_1: C\mu
eq \underline{0}$

Interpretation: a contrast matrix is used to specify comparisons among group means. It allows to test hypotheses about specific combinations of group means beyond simple pairwise comparisons.

Recall

We already know that:

- $C\underline{x}$ is unbiased for $C\mu$;
- $\overline{\underline{x}} \sim N_q(\mu, \frac{1}{n}\Sigma) \Rightarrow C\overline{\underline{x}} \sim N_{q-1}(C\mu, \frac{1}{n}C\Sigma C')$
- S is an estimator for $\Sigma\Rightarrow (n-1)CSC'\sim Wish(C\Sigma C',n-1)\perp\!\!\!\perp C\overline{\underline{x}}$
 - Hotelling Theorem $n(C\overline{\underline{x}}-C\underline{\mu})'(CSC')^{-1}(C\overline{\underline{x}}-C\underline{\mu})\sim \frac{(n-1)(q-1)}{n-q+1}F(q-1,n-q+1)$ which is pivotal

We define the test statistics as $T_0^2=n(C\overline{\underline{x}})'(CSC')^{-1}C\overline{\underline{x}}$ so for a fixed α we reject H_0 if $T_0^2>\frac{(n-1)(q-1)}{n-q+1}F_{1-\alpha}(q-1,n-q+1)$

Obs

Let C, \tilde{C} two contrast matrices, take $B \in (q-1) \times (q-1)$ with $det(B) \neq 0$ such that $C = B\tilde{C}$ which is the operator to change basis. So:

$$T_0^2 = n(C\overline{\underline{x}})'(CSC')^{-1}C\overline{\underline{x}} = n(B\tilde{C}\overline{\underline{x}})'(B\tilde{C}S\tilde{C}'B')^{-1}B\tilde{C}\overline{\underline{x}} = n(\tilde{C}\overline{\underline{x}})'(\tilde{C}S\tilde{C}')^{-1}\tilde{C}\overline{\underline{x}} = \tilde{T}_0^2$$

Obs



$$\mathsf{Let}\ \underline{x}_1,\dots,\underline{x}_n\ \mathsf{iid}\ \sim N_q(\underline{\mu},\Sigma),\ \mathsf{have}\ \mathcal{L}^\perp = span(\underline{c}_1,\dots,\underline{c}_{p-k})\ \mathsf{so}\ \mathsf{our}\ C = \begin{bmatrix} \ \underline{c}_1' \ \\ \vdots \ \\ \underline{c}_{p-k}' \end{bmatrix} \ \mathsf{and}\ \mathsf{we}\ \mathsf{can}\ \mathsf{also}$$

test:
$$H_0: C\underline{\mu} = \underline{0} \quad ext{ vs } \quad H_1: C\underline{\mu}
eq \underline{0}$$

So our test statistics
$$T_0^2=n(C\overline{\underline{x}})'(CSC')^{-1}C\overline{\underline{x}}\sim \frac{(n-1)(p-k)}{n-p+k}F(p-k,n-p+k)$$

Multivariate case

Each unit isn't a number but become a vector of dimension l that is observed q times.

So to construct the test we made:

$$H_0: egin{cases} \mu_{1,1} = \cdots = \mu_{q,1} \ dots & dots & dots \ \mu_{1,l} = \cdots = \mu_{q,l} \end{cases} ext{ vs } H_1: \exists i,j,k ext{ s.t. } \mu_{i,k}
eq \mu_{j,k}$$

tips: change rappresentation to achieve

 $\underline{x}_i = (x_{i,1,1}, \dots, x_{i,1,l}, x_{i,2,1}, \dots, x_{i,2,l}, \dots, x_{i,q,1}, \dots, x_{i,q,l}) \in \mathbb{R}^{lq}$ in this way C is a block matrix where each block is a contrast matrix.

Multivariate ANalysis Of VAriance #Manova

Framework: To complicate things we can take sample from different Gaussian distribution. So our sample become: $\underline{x}_{1,1},\ldots,\underline{x}_{1,n_1}$ iid $\sim N_p(\underline{\mu}_1,\Sigma);\ldots;\underline{x}_{g,1},\ldots,\underline{x}_{g,n_g}$ iid $\sim N_p(\underline{\mu}_q,\Sigma)$

$$p \ge 1$$
 and $g = 2$:

$$\underline{x}_{1,1},\ldots,\underline{x}_{1,n_1} ext{ iid } \sim N_p(\underline{\mu}_1,\Sigma) \perp\!\!\!\perp \underline{x}_{2,1},\ldots,\underline{x}_{2,n_2} ext{ iid } \sim N_p(\underline{\mu}_2,\Sigma)$$

Goal: inference of $\underline{\mu}_1 - \underline{\mu}_2$

We have:

$$\bullet \ \ \overline{\underline{x}}_1 - \overline{\underline{x}}_2 \sim N_p(\underline{\mu}_1 - \underline{\mu}_2, (\tfrac{1}{n_1} + \tfrac{1}{n_2})\Sigma) \Rightarrow (\tfrac{1}{n_1} + \tfrac{1}{n_2})^{-\frac{1}{2}}[(\overline{\underline{x}}_1 - \overline{\underline{x}}_2) - (\underline{\mu}_1 - \underline{\mu}_2)] \sim N_p(\underline{0}, \Sigma)$$

• S_1,S_2 both estimate Σ so we create $S_{pooled}=rac{(n_1-1)S_1+(n_2-1)S_2}{n_1+n_2-2}\Rightarrow (n_1+n_2-2)S_{pooled}\sim Wish(\Sigma,n_1+n_2-2)$

So we can develop Hotelling theorem obtaining:

$$oxed{ egin{aligned} & (rac{1}{n_1} + rac{1}{n_2})^{-1}[(\overline{\underline{x}}_1 - \overline{\underline{x}}_2) - (\underline{\mu}_1 - \underline{\mu}_2)]'S_{pooled}^{-1}[(\overline{\underline{x}}_1 - \overline{\underline{x}}_2) - (\underline{\mu}_1 - \underline{\mu}_2)] = \ & = d_{[(rac{1}{n_1} + rac{1}{n_2})S]^{-1}}(\overline{\underline{x}}_1 - \overline{\underline{x}}_2, \underline{\mu}_1 - \underline{\mu}_2) \sim rac{(n_1 + n_2 - 2)p}{n_1 + n_2 - 1 - p}F(p, n_1 + n_2 - 1 - p) \end{aligned}$$

Which is a pivotal quantity.

Take $\alpha \in (0,1)$:

$$CR_{1-\alpha}(\underline{\mu}_1-\underline{\mu}_2)=\{\underline{\nu}=\underline{\mu}_1-\underline{\mu}_2:d_{[(\frac{1}{n_1}+\frac{1}{n_2})S]^{-1}}(\overline{\underline{x}}_1-\overline{\underline{x}}_2,\underline{\nu})\leq \tfrac{(n_1+n_2-2)p}{n_1+n_2-1-p}F_{1-\alpha}(p,n_1+n_2-1-p)\}$$

Anova

Framework: we have unit of p=1 features and we collect g different unit each one $n_{i\in\{1,\ldots,g\}}$ times.



Our sample is made by:

$$x_{1,1},\ldots,x_{1,n_1} ext{ iid } \sim N_1(\mu_1,\sigma^2); \ x_{2,1},\ldots,x_{2,n_2} ext{ iid } \sim N_1(\mu_2,\sigma^2); \ \ldots; x_{g,1},\ldots,x_{g,n_g} ext{ iid } \sim N_1(\mu_g,\sigma^2)$$
 Where $x_{i,k} \perp \!\!\! \perp x_{j,l} \quad orall i, k,j,l$

Firstly, we reparameter the problem by breaking down $\mu_i = \mu + \tau_i$ so we can express each $x_{i,j} = \mu + \tau_i + \epsilon_{i,j}$ with $\epsilon_{i,j} \in N(0,\sigma^2)$.

Problem: we pass from g-parameters (μ_i) to (g+1)-parameters (μ, τ_i) .

Goal - 1: Find estimator for μ

$$\overline{x} = rac{1}{n} \sum_{i=1}^g \sum_{j=1}^{n_i} x_{i,j}$$
 with $n = n_1 + n_2 + \dots + n_g$

Is it unbiased?

$$\mathbb{E}[\overline{x}] = rac{1}{n} \sum_{i=1}^g \sum_{j=1}^{n_i} \mathbb{E}[x_{i,j}] = rac{1}{n} \sum_{i=1}^g \sum_{j=1}^{n_i} (\mu + au_i) = \mu + rac{\sum_i n_i au_i}{n}$$

Idea: To unbiased the estimator of μ and to resolve the problem of too much parameters we add a constrain: $\sum_i n_i \tau_i = 0$

Goal - 2: Find estimator for τ_i

$$egin{aligned} \overline{x}_i &= rac{1}{n} \sum_{j=1}^{n_i} x_{i,j} ext{ with } n=n_1+n_2+\cdots+n_g \ \Rightarrow \mathbb{E}[\overline{x}_i-\overline{x}] &= \mu+ au_i-\mu= au_i \end{aligned}$$

Recall - Decomposition of variance

By construction we know that the following properties holds for u_i :

- $\underline{u}_1, \dots, \underline{u}_q$ are linearly independent;
- $ullet < \underline{u}_i, \underline{u}_i > = 0 \quad orall i
 eq j ext{ so they are orthogonal each others;}$
- $\underline{1} \in span\{\underline{u}_1, \dots, \underline{u}_q\}$

So we reach:

$$\underline{x} = \underbrace{\overline{x} \cdot \underline{1}}_{\mu} + \underbrace{\sum_{i=1}^g (\overline{x}_i - \overline{x}) \underline{u}_i}_{\sum_i (\mu_i - \mu) = \sum_i \tau_i = au} + (\underline{x} - \sum_g^{i=1} \overline{x}_i \cdot \underline{u}_i)$$

Using Pitagora's theorem can write:

$$\begin{split} ||\underline{x}||^2 &= ||\overline{x} \cdot \underline{1}||^2 + ||\sum_{i=1}^g (\overline{x}_i - \overline{x})\underline{u}_i||^2 + ||\underline{x} - \sum_g^{i=1} \overline{x}_i \cdot \underline{u}_i||^2 \\ \Rightarrow \underbrace{\sum_{i=1}^g x_i^2}_{SS_{TOT}} + \underbrace{\sum_{i=1}^g n_i (\overline{x}_i - \overline{x})^2}_{SS_{treatment}} + \underbrace{\sum_{i=1}^g \sum_{j=1}^{n_i} (x_{i,j} - \overline{x}_i)^2}_{SS_{residual}} \leftrightarrow \#1_decomposition_of_variance \\ \Rightarrow \underbrace{||\underline{x} - \overline{x} \cdot \underline{1}||^2}_{SS_{treatment}} + \underbrace{||\underline{x} - \sum_g^{i=1} \overline{x}_i \cdot \underline{u}_i||^2}_{SS_{residual}} \leftrightarrow \#2_decomposition_of_variance \end{split}$$



Rmk: SS stands for Sum of Squares

$$\pi_{\Sigma|\underline{u}_{1},...,\underline{u}_{2}} = \frac{1}{\underline{u}_{1}^{2}} \underbrace{x}_{1} = \dots = \underbrace{x}_{1} \cdot \underline{u}_{1}^{2}$$

$$\pi_{\Sigma|\underline{u}_{1},...,\underline{u}_{2}} = \frac{1}{\underline{u}_{1}^{2}} \underbrace{x}_{1} = \dots = \underbrace{x}_{1} \cdot \underline{u}_{1}^{2}$$

$$\pi_{\Sigma|\underline{u}_{1}|\underline{u}_{1}^{2}} = \frac{1}{\underline{u}_{1}^{2}} \underbrace{x}_{1} = \dots = \underbrace{x}_{1} \cdot \underline{u}_{1}^{2}$$

$$\pi_{\Sigma|\underline{u}_{1}|\underline{u}_{1}^{2}} = \underbrace{1}_{\underline{u}_{1}^{2}} \underbrace{x}_{1} = \dots = \underbrace{x}_{1} \cdot \underline{u}_{1}^{2}$$

$$\pi_{\Sigma|\underline{u}_{1}|\underline{u}_{1}^{2}} = \underbrace{1}_{\underline{u}_{1}^{2}} \underbrace{x}_{1} = \dots = \underbrace{x}_{1} \cdot \underline{u}_{1}^{2}$$

Goal - 3: Test $H_0: \mu_1 = \cdots = \mu_g$ vs $H_1: H_0^c$ equivalent to $H_0: au_1 = \cdots = au_g$ **Idea:** reject H_0 if $SS_{treatment}$ is large w.r.t. $SS_{residual}$ so we take the ratio $\frac{SS}{SS}$

 $SS_{residuals} = \sum_{i=1}^g \sum_{j=1}^{n_i} (x_{i,j} - \overline{x}_i)^2 = \sum_{i=1}^g (n_i - 1)S_i^2 \xrightarrow{ ext{since groups are } \mathbb{H}} SS_{residuals} \sim \sigma^2 \chi^2(n-g)$ If H_0 is true:

- $\sum_{i=1}^g \sum_{j=1}^{n_i} (x_{i,j}-\overline{x})^2 = (n-1)S \sim \sigma^2 \chi^2 (n-1)$
- $SS_{treatment} \sim \sigma^2 \chi^2 (q-1)$

Remark: \overline{x}_i is the mean of n_i measurements of a unit while \overline{x} is the overall mean

So
$$rac{SS_{treatment}}{g-1} rac{n-g}{SS_{residuals}} \sim F(g-1,n-g)$$

So
$$\frac{SS_{treatment}}{g-1} \frac{n-g}{SS_{residuals}} \sim F(g-1,n-g)$$
 \Rightarrow reject H_0 at level α if $\frac{SS_{treatment}}{g-1} \frac{n-g}{SS_{residuals}} > F_{1-\alpha}(g-1,n-g)$

Manova

Framework: we have unit of $p \ge 1$ features and we collect g different unit each one $n_{i \in \{1, \dots, q\}}$ times.

Take $x_{i,j} ext{ iid } \sim N_p(\mu,\Sigma)$ with $i \in \{1,\ldots,g\}$ and $j \in \{1,\ldots,n_i\}$ We reparametrize so reach: $\boxed{\underline{x}_{i,j} = \underline{\mu} + \underline{\tau}_i + \underline{\epsilon}_{i,j}}$ with $\underline{\epsilon}_{i,j} \sim N_p(\underline{0}, \Sigma)$ and as constrain $\sum_{i} n_{i} \underline{\tau}_{i} = 0$

Decomposition of covariance

Taking $\overline{\underline{x}} = \frac{1}{n} \sum_{i=1}^g \sum_j^{n_i} \underline{x}_{i,j}$ we deduce:

$$\sum_{i=1}^g \sum_{j=1}^{n_i} (\underline{x}_{i,j} - \overline{\underline{x}}) (\underline{x}_{i,j} - \overline{\underline{x}})' = \sum_{i=1}^g n_i (\overline{\underline{x}}_i - \overline{\underline{x}}) (\overline{\underline{x}}_i - \overline{\underline{x}})' + \sum_{i=1}^g \sum_{j=1}^{n_i} (\underline{x}_{i,j} - \overline{\underline{x}}_i) (\underline{x}_{i,j} - \overline{\underline{x}}_i)' = B + W$$

Goal: Test $H_0: \underline{\mu}_1=\dots=\underline{\mu}_q$ vs $H_1: H_0^c$ equivalent to $H_0: \underline{\tau}_1=\dots=\underline{\tau}_q$ vs $H_1: H_0^c$ Test statistics proposals:

- #Wilks $\Lambda_W = \frac{det(W)}{det(W+B)} \Rightarrow \text{reject } H_0 \text{ if it is too small}$
- #Pillai $\Lambda_P = trace(B(B+W)^{-1}) \Rightarrow \text{reject } H_0 \text{ if it is too big}$
- #Hotelling-Lawley $\Lambda_{HL} = trace(BW^{-1}) \Rightarrow$ reject H_0 if it is too big



Obs: each test statistics described before could be expressed in terms of the eigenvalues of BW^{-1}

Question: Which is the distribution of #Wilks statistics Λ_W if H_0 holds?

Theo - Bartlett asymptotic approximation

If H_0 holds then: $-(n-1-rac{p+g}{2})\ln(\Lambda_W)\sim \chi^2(p(g-1))$

So reject H_0 at level α if $-(n-1-\frac{p+g}{2})\ln(\Lambda_W)>\chi^2_{1-\alpha}(p(g-1))$

If we reject H_0 we want to create a $CI(\tau_{i,l}-\tau_{k,l})$ with $i,k\in\{1,\ldots,g\}$ and $l\in\{1,\ldots,p\}$.

To do so firstly we need a point estimator for $\tau_{i,l} - \tau_{k,l}$.

Since $\underline{\tau}_i$ is estimated by $\overline{\underline{x}}_i - \underline{\overline{x}}$ we conclude that $\tau_{i,l}$ is estimated by $\overline{x}_{i,l} - \overline{x}_l$.

 $r_{i,l}- au_{k,l}$ is estimated by $\overline{x}_{i,l}-\overline{x}_{k,l}\sim N(au_{i,l}- au_{k,l},\sigma_{l,l}(rac{1}{n_i}+rac{1}{n_k}))$ but we don't know $\sigma_{l,l}$ so we need to estimate it.

 \Rightarrow since Σ is estimated by $S_{pooled}=rac{1}{n-g}\sum_{i=1}^g(n_i-1)S_i=rac{W}{n-g}$ with

$$S_i = rac{1}{n_i-1}\sum_{i=1}^{n_i}(\underline{x}_{i,j}-\overline{x}_i)(\underline{x}_{i,j}-\overline{x}_i)'$$
 we use as estimator of $\sigma_{l,l}$ the $S_{pooled_{l,l}}=rac{w_{l,l}}{n-g}$

Now we can create Bonferroni $CI_{1-lpha}(au_{i,l}- au_{k,l})=[\overline{x}_{i,l}-\overline{x}_l\pm t_{1-rac{lpha}{2\$}}(n-g)\sqrt{rac{w_l}{n-a}(rac{1}{n_i}+rac{1}{n_s})}]$ with as (s) = $p^{\frac{g(g-1)}{2}}$

Extension to two-way ANOVA.

Framework: we have 2 treatment (factors), the first treatment is made by g levels while the second one has b levels.

We have two possible models for our $\mu_{i,j}$:

- 1. Complete model: $\mu_{i,j} = \mu + \tau_i + \beta_j + \gamma_{i,j} \leftarrow \gamma_{i,j}$ model the relation between first and second treatment level.
- 2. Additive model: $\mu_{i,j} = \mu + \tau_i + \beta_j \leftarrow$ effect of first treatment are independent from the second treatment level

Remark: link useful to understand

Remark: $\overline{x}_{i,j}=rac{1}{n}\sum_{k=1}^n x_{i,j,k}$ Remark: $\overline{x}=rac{1}{n\cdot g\cdot b}\sum_{i=1}^g \sum_{j=1}^b \sum_{k=1}^n x_{i,j,k}$

Our estimators are:

- $\mu
 ightarrow \overline{x}$
- ullet $au_i
 ightarrow \overline{x}_{i,\cdot} \overline{x}_i$
- $\beta_i \to \overline{x}_{\cdot,i} \overline{x}$
- $\bullet \ \ \gamma_{i,j} \to \overline{x}_{i,j} (\overline{x}_{\cdot,j} \overline{x}) (\overline{x}_{i,\cdot} \overline{x}) \overline{x} = \overline{x}_{i,j} \overline{x}_{\cdot,j} \overline{x}_{i,\cdot} + \overline{x}$

Our constrains are:

- $\sum_i \tau_i = 0$
- $\sum_{i} \beta_{i} = 0$
- $\sum_{i} \gamma_{i,j} = 0 \quad \forall j$
- $\sum_{i} \gamma_{i,j} = 0 \quad \forall i$

$$\sum_{i}^{g} \sum_{j}^{b} \sum_{k}^{n} (x_{i,j,k} - \overline{x})^{2} = \sum_{i}^{g} nb(\overline{x}_{i,\cdot} - \overline{x})^{2} + \sum_{j}^{b} ng(\overline{x}_{\cdot,j} - \overline{x})^{2} + \sum_{i}^{g} \sum_{j}^{b} n(\overline{x}_{i,j} - \overline{x}_{i,\cdot} - \overline{x}_{\cdot,j} + \overline{x})^{2} + \sum_{i}^{g} \sum_{j}^{b} \sum_{k}^{n} (x_{i,j,k} - \overline{x}_{i,j})^{2} + \sum_{i}^{g} \sum_{j}^{h} \sum_{k}^{n} (x_{i,j,k} - \overline{x}_{i,j})^{2} + \sum_{i}^{g} \sum_{j}^{h} \sum_{k}^{n} (x_{i,j,k} - \overline{x}_{i,j})^{2} + \sum_{i}^{g} \sum_{k}^{h} \sum_{k}$$

With:

- $SS_{
 m treatment \, 1} =$ variability explain by treatment 1 has g-1 degree of freedom
- $SS_{
 m treatment~2} =$ variability explain by treatment 2 has b-1 degree of freedom
- $SS_{
 m interactions}=$ variability explain by treatment 1 and treatment 2 interactions has (g-1)(b-1) degree of freedom
- $SS_{
 m residuals} =$ variability remaining has gb(n-1) degree of freedom

How to test if we need to model interactions: $H_0: \gamma_{i,j} = 0 \quad orall i, j \quad ext{ vs } \quad H_1: \exists \gamma_{i,j}
eq 0$

We reject
$$H_0$$
 if $\boxed{rac{SS_{interact}}{(g-1)(b-1)}rac{gb(n-1)}{SS_{res}}}>F_{1-lpha}((g-1)(b-1),gb(n-1))$

If H_0 holds we can use additive model

Remark: testing treatment one, similar to one-way ANOVA, is equal to check

$$rac{SS_{ ext{treatment 1}}}{g-1} rac{gb(n-1)}{SS_{res}} > F_{1-lpha}(g-1,gb(n-1))$$

Classification

Framework: each unit is represented by (\underline{x}', L) where L is the label while \underline{x}' are features **Goal:** learning $\delta: X \to \text{Label-space}$

There are 2 possible ways to approaches to this problem:

- 1. **Discriminant Analysis Supervised Learning:** we train our model starting from a training set with features x' and labels L;
- Cluster Analysis Unsupervised Learning: we don't know a-priori labels so we cluster together units with similar features.

#Discriminant_Analysis - Supervised Learning

To use this approach we need 3 ingredients:

- Distribution of features: $\underline{x}|L=i\sim f_i(\underline{x})\leftarrow$ within-class density of features given the class label, used into calculation of likelihood;
- Prior distribution: $\mathbb{P}[L=i]=p_i$ with $i\in\{1,\ldots,g\}\leftarrow$ represents the initial belief about the probability of each class occurring before observing any data, used into calculation of posterior probabilities;
- Cost of mis-classification: $c(i|j) = \cos t$ of attributing unit to group i while it belongs to group j

Remark: we usually have a costs matrix C and it would be desirable if diag(C)=c(i,i)=0Remark: learning a classification function δ is equivalent to a learn a partition $\{R_1,\ldots,R_q\}$ of X



s.t. $R_i \cap R_j = \emptyset$ if $i \neq j$ and $\cup_i^g R_i = X$

Remark: usually we want to minimize the #ECM (Expected Cost of Misclassification)

Example - dichotomous classifier

 $\delta o \{R_1,R_2\}$ so if $\delta(\underline{x})=1$ implies that $\underline{x} \in R_1$

$$egin{aligned} ECM(\delta) &= \int_{R_2} c(2ert 1) f_1(\underline{x}) p_1 d\underline{x} + \int_{R_1} c(1ert 2) f_2(\underline{x}) p_2 d\underline{x} \ &= \underbrace{\int_{X} c(2ert 1) f_1(\underline{x}) p_1 d\underline{x}}_{c(2ert 1) p_1} - \int_{R_1} c(2ert 1) f_1(\underline{x}) p_1 d\underline{x} + \int_{R_1} c(1ert 2) f_2(\underline{x}) p_2 d\underline{x} \ &= c(2ert 1) p_1 + \int_{R_1} (-c(2ert 1) f_1(\underline{x}) p_1 + c(1ert 2) f_2(\underline{x}) p_2) d\underline{x} \end{aligned}$$

 δ optimal is the one which minimize $ECM(\delta)$ so:

$$egin{aligned} R_1 &= \{ \underline{x} \in X : c(2|1) f_1(\underline{x}) p_1 \geq c(1|2) f_2(\underline{x}) p_2 \} \ R_2 &= \{ \underline{x} \in X : c(2|1) f_1(\underline{x}) p_1 \leq c(1|2) f_2(\underline{x}) p_2 \} \end{aligned}$$

Idea:

$$\mathbb{P}(X=\underline{x},cls=1)=f_1(\underline{x})p_1\\ =>\int_{R_2}c(2|1)f_1(\underline{x})p_1d\underline{x}=\int_{R_2}c(2|1)\mathbb{P}(X=\underline{x},cls=1)d\underline{x}=\text{probs of misclassify all the obs of the class 1 as class 2}$$

Obs:

$$\delta(\underline{x}) = t \in \{1, \dots, g\} \Leftrightarrow rac{1}{\sum_{l=1}^g f_l(\underline{x}) p_l} \sum_{k
eq t} c(t|k) f_k(\underline{x}) p_k \leq rac{1}{\sum_{l=1}^g f_l(\underline{x}) p_l} \sum_{k
eq j} c(j|k) f_k(\underline{x}) p_k \quad orall j
eq t$$

Idea:

Recalling that: $\frac{f_k(\underline{x})p_k}{\sum_{j=1}^g f_j(\underline{x})p_j} = \frac{\mathbb{P}[\underline{x}|L=k]\mathbb{P}(L=k)}{\sum_{j=1}^g \mathbb{P}[\underline{x}|L=j]\mathbb{P}(L=j)} = \mathbb{P}[L=k|\underline{x}] = \text{ posterior probability So, we reach:}$

$$\delta(\underline{x}) = t \Leftrightarrow \sum_{k
eq t} c(t|k) \mathbb{P}[L = k|\underline{x}] \leq \sum_{k
eq j} c(j|k) \mathbb{P}[L = k|\underline{x}] \quad orall j
eq t$$

We are minimize ECM (Excpected Cost of Misclassification)

Example - Bayes classifier

$$c(i,j)=d=constant\geq 0 \quad orall i
eq j ext{ and } c(i,i)=0 \quad orall i$$
 Optimal δ :

$$\delta(\underline{x}) = t \Leftrightarrow \sum_{k
eq t} \mathbb{P}[L = k | \underline{x}] \leq \sum_{k
eq j} \mathbb{P}[L = k | \underline{x}] \quad orall j
eq t$$

ldea:

We are minimizing the sum of the posterior probabilities of assign a datum to a wrong cluster.

Example - Maximum Likelihood classifier



$$c(i,j)=d=constant \geq 0 \quad orall i
eq j; \ c(i,i)=0 \quad orall i; \ p_1=p_2=\cdots=rac{1}{g}$$
 Optimal δ : $\delta(\underline{x})=t \Leftrightarrow rac{f_t(\underline{x})p_t}{\sum_{i=1}^g f_i(\underline{x})p_i} \geq rac{f_k(\underline{x})p_k}{\sum_{i=1}^g f_i(\underline{x})p_i} \Leftrightarrow f_t(\underline{x}) \geq f_k(\underline{x}) \quad orall k
eq t$

If the features are Gaussian [so $\underline{x}|L \sim N_p(\mu_i, \Sigma_i)]$ implies that the Bayes classifier become:

$$\begin{split} \delta(\underline{x}) &= t \Leftrightarrow \sum_{k \neq t} \mathbb{P}[L = k | \underline{x}] \leq \sum_{k \neq j} \mathbb{P}[L = k | \underline{x}] \quad \forall j \neq t \\ & \to 1 - \sum_{k \neq t} \mathbb{P}[L = k | \underline{x}] \geq 1 - \sum_{k \neq j} \mathbb{P}[L = k | \underline{x}] \quad \forall j \neq t \\ & \to \mathbb{P}[L = t | \underline{x}] \geq \mathbb{P}[L = j | \underline{x}] \quad \forall j \neq t \\ & \to f_t(\underline{x}) p_t \geq f_j(\underline{x}) p_j \quad \forall j \in \{1, \dots, g\} \\ & \to p_t \frac{\exp\{-\frac{1}{2}(\underline{x} - \underline{\mu}_t)' \Sigma_t^{-1}(\underline{x} - \underline{\mu}_t)\}}{[(2\pi)^p \det(\Sigma_t)]^{\frac{1}{2}}} \geq p_j \frac{\exp\{-\frac{1}{2}(\underline{x} - \underline{\mu}_j)' \Sigma_j^{-1}(\underline{x} - \underline{\mu}_j)\}}{[(2\pi)^p \det(\Sigma_j)]^{\frac{1}{2}}} \\ & \to \ln(p_t) - \frac{1}{2} \ln(\det(\Sigma_t)) - \frac{1}{2} d_{\Sigma_t^{-1}}(\underline{x}, \underline{\mu}_t) \geq \ln(p_j) - \frac{1}{2} \ln(\det(\Sigma_j)) - \frac{1}{2} d_{\Sigma_j^{-1}}(\underline{x}, \underline{\mu}_j) \end{split}$$

Obs:

If $p_1=p_2=\cdots=p_g=rac{1}{g}$ and $\Sigma_1=\Sigma_2=\cdots=\Sigma_g$ then our classifier become: $\delta(\underline{x})=t\Leftrightarrow \underline{x}\in R_t\Leftrightarrow d_{\Sigma^{-1}}(\underline{x},\underline{\mu}_t)\leq d_{\Sigma^{-1}}(\underline{x},\underline{\mu}_i)$

Interpretation: we are assigning a point to the group with the closest mean in a distribution sense, since the distance cited above is the #Mahalanobis distance.

QDA

Hypothesis: Gaussianity

We call **Quadratic Discriminant Function** $=d_t^Q(\underline{x})=\ln(p_t)-\frac{1}{2}\ln(\det(\Sigma_t))-\frac{1}{2}d_{\Sigma_t^{-1}}(\underline{x},\underline{\mu}_t)$ So we can rewrite our Bayes classifier with Gaussian features using the previous definition: $\delta(\underline{x})=t\Leftrightarrow \underline{x}\in R_t\Leftrightarrow d_t^Q(\underline{x})\geq d_j^Q(\underline{x})\leftarrow \text{which is } \textit{Quadratic Discriminant Analysis}$ (#QDA) **Interpretation:** we are maximizing log-likelihood

LDA

Hypothesis: Gaussianity and Homoscedasticity

$$\underline{x} \underline{\Sigma}^{-1} \underline{\mu}_t + \ln(p_t) - \tfrac{1}{2} \underline{\mu}_t \underline{\Sigma}^{-1} \underline{\mu}_t \geq \underline{x} \underline{\Sigma}^{-1} \underline{\mu}_j + \ln(p_j) - \tfrac{1}{2} \underline{\mu}_j \underline{\Sigma}^{-1} \underline{\mu}_j \quad \forall j \neq t$$
 We call **Linear Discriminant Function**
$$= d_t(\underline{x}) = \underline{x} \underline{\Sigma}^{-1} \underline{\mu}_t + \ln(p_t) - \tfrac{1}{2} \underline{\mu}_t \underline{\Sigma}^{-1} \underline{\mu}_t$$

So we can rewrite our Bayes classifier with Gaussian features using the previous definition:

$$\delta(\underline{x}) = t \Leftrightarrow \underline{x} \in R_t \Leftrightarrow d_t(\underline{x}) \geq d_j(\underline{x}) \leftarrow ext{which is } \textit{Linear Discriminant Analysis}$$
 (#LDA)

Remark: We use the training set to estimate $f_i(\underline{x}) \quad \forall i$

#LDA : $\mu_1,\ldots,\mu_g,\Sigma$ should be estimate from data



#QDA: $\mu_1, \ldots, \mu_g, \Sigma_1, \ldots, \Sigma_g$ should be estimate from data

So we estimate:

• μ_i with $\overline{\underline{x}}_i = \frac{1}{n_i} \sum_{\{j: e_j = i\}} \underline{x}_j$

•
$$\Sigma_i$$
 with $S_i=rac{1}{n_i-1}\sum_{\{j:e_j=i\}}(\underline{x}_j-\overline{\underline{x}}_i)(\underline{x}_j-\overline{\underline{x}}_i)' o$ for QDA

•
$$\Sigma$$
 with $S_{
m pooled}=rac{1}{n-g}\sum_i(n_i-1)S_i o$ for LDA with $n=n_1+n_2+\cdots+n_g$

Since we have a lot of things to estimate we need that n_1, n_2, \ldots, n_g is large with respect to p.

Naive Bayes Classifier

A classifier which not require a large sample size training set is **Naive Bayes Classifier** which parameterize Σ_i as a diagonal matrix so $d_t^Q(x)$ become:

$$d_t^Q(\underline{x}) = \ln(p_t) - rac{1}{2} \sum_{i=1}^p \ln(\sigma_{ii}^{(t)}) - rac{1}{2} \sum_{i=1}^p rac{(x_i - \overline{x}_{t,i})^2}{\sqrt{\sigma_{i,i}^{(t)}}} ext{ with } \sigma_{i,i}^{(t)} = rac{1}{n_t - 1} \sum_{\{j: e_j = i\}} (\underline{x}_{j,i} - \overline{x}_{t,i})^2$$

Example #knn - classifier

Fix $k \geq 1$ then $N_k(\underline{x}) = \{$ k units \underline{x}_i which are closest to $\underline{x}\}$

We assign the label t to \underline{x} if it is the most frequent label in $N_k(\underline{x})$

Fisher argument for #LDA

Goal: We develop the #LDA classifier framework without assuming Gaussinity.

Idea: It is a method designed to find a linear combination of features that separates two or more classes of objects or events. The core idea behind Fisher's argument for LDA is to project high-dimensional data onto a lower-dimensional space in such a way that maximizes the separability among known categories.

Framework: LDA framework

Define B= Variance matrix between group means $=\frac{1}{g-1}\sum_{i=1}^g(\underline{\mu}_i-\overline{\underline{\mu}})(\underline{\mu}_i-\overline{\underline{\mu}})'$ with $\underline{\overline{\mu}}=\frac{1}{g}\sum_{i=1}^g\underline{\mu}_i$ and as $\underline{\mu}_i=\mathbb{E}[\underline{x}|L=i]$

So with $a \in \mathbb{R}^p$:

- $\mathbb{E}[\underline{a}'\underline{x}|L=i] = \underline{a}'\mu_i$
- $var[a'x|L=i] = a'\Sigma a$

 $\textit{Problem:} \ \mathsf{find} \ \underset{a \in \mathbb{R}^p}{arg\, max} \ \tfrac{\underline{a'B\underline{a}}}{\underline{a'\Sigma\underline{a}}} = \underset{a \in \mathbb{R}^p}{arg\, max} \ \tfrac{\frac{1}{g-1} \sum_{i=1}^g (\underline{a'}\underline{\mu}_i - \underline{a'}\overline{\mu})^2}{\underline{a'\Sigma\underline{a}}}$

Solution: same reasoning as <code>#PCA</code> we find that $arg \max_{\underline{a} \in \mathbb{R}^p} \frac{\underline{a}'B\underline{a}}{\underline{a}'\Sigma\underline{a}} = \Sigma^{-\frac{1}{2}}\underline{e}_1$ with \underline{e}_i are the eigenvectors of $\Sigma^{-\frac{1}{2}}B\Sigma^{-\frac{1}{2}}$.

We can rewrite \underline{x} using \underline{a}_i as reference system so $\underline{x} \to \underline{\tilde{x}} \Rightarrow \underline{a}_i' \underline{x}$ are called *Fisher Discriminant*

Scores and have a useful feature: $cov(\underline{a}_i\underline{x},\underline{a}_j\underline{x}) = \begin{cases} 0 & i \neq j \\ 1 & \mathrm{else} \end{cases} \Rightarrow cov(A\underline{x}) = I$

Interpretation: how well-separated the classes are along the FLD axis.



So, how to build a classifier?

We estimate:

•
$$\mu_i$$
 with $\overline{\underline{x}}_i = \frac{1}{n_i} \sum_{\{j: e_j = i\}} \underline{x}_j$

•
$$\Sigma_i$$
 with $S_i = rac{1}{n_i-1}\sum_{\{j:e_j=i\}} (\underline{x}_j - \overline{\underline{x}}_i)(\underline{x}_j - \overline{\underline{x}}_i)'$

•
$$\Sigma$$
 with $S_{
m pooled}=rac{1}{n-g}\sum_i(n_i-1)S_i o$ for LDA with $n=n_1+n_2+\cdots+n_g$

So our
$$\hat{B} = \frac{1}{g-1} \sum_{i=1}^g (\underline{x}_i - \overline{\underline{x}}) (\underline{x}_i - \overline{\underline{x}})'$$

So: $\delta(\underline{x}) = t \Leftrightarrow \sum_{i=1}^k (\tilde{x}_i - \tilde{x}_{t,i})^2 \leq \sum_{i=1}^k (\tilde{x}_i - \tilde{x}_{j,i})^2$

Interpretation: we assign x to the closest projection on the Fisher Discriminant Scores.

Evaluating a classifier

$$\textit{framework:} \ \mathbb{X} = \text{training set} = \begin{bmatrix} \vdots & \vdots \\ \underline{x}_k & l_k \\ \vdots & \vdots \end{bmatrix} \text{ with } \underline{x}_k \in \mathbb{R}^p \text{ and } l_k \in \{1,2,\ldots,g\}$$

Recall:

$$AER(\delta) = ext{Actual Error Rate} = \sum_{k
eq 1} \int_{\mathbb{R}_k} f_1(\underline{x}) p_1 d\underline{x} + \sum_{k
eq 2} \int_{\mathbb{R}_k} f_2(\underline{x}) p_2 d\underline{x} + \dots + \sum_{k
eq g} \int_{\mathbb{R}_k} f_g(\underline{x}) p_g d\underline{x}$$

Q: How to evaluate a classifier? How to estimate $AER(\delta)$ from data?

For sake of simplicity our reasoning are in the framework of g=2. So, the correct

$$AER(\delta) = \sum_{k
eq 1} \int_{\mathbb{R}_k} f_1(\underline{x}) p_1 d_{\underline{x}} + \sum_{k
eq 2} \int_{\mathbb{R}_k} f_2(\underline{x}) p_2 d_{\underline{x}}$$

How to estimate it from data?

- Apply δ to \mathbb{X} ;
- · Construct the confusion matrix;
- $A\hat{ER}(\delta) = APER(\delta) = ext{APpearent Error Rate} = rac{n_{1,2} + n_{2,1}}{n} \leftarrow ext{spoiler: is too optimistic}$

Remember that the confusion matrix is construct as follow:

\downarrow truth \ estimated $ ightarrow$	1	2	
1	$n_{1,1}$	$n_{1,2}$	$n_{1,\cdot}$
2	$n_{2,1}$	$n_{2,2}$	$n_{2,\cdot}$
	$n_{\cdot,1}$	$n_{\cdot,2}$	

With a bit of computation we reach that:

$$APER(\delta) = \hat{p_1} \int_{\mathbb{R}_2} \widehat{f_1(\underline{x})} d\underline{x} + \hat{p_2} \int_{\mathbb{R}_1} \widehat{f_2(\underline{x})} d\underline{x} = rac{n_{1,\cdot}}{n} rac{n_{1,2}}{n_{1,\cdot}} + rac{n_{2,\cdot}}{n} rac{n_{2,1}}{n_{2,\cdot}}$$

Obs

In case of dichotomous $\{0,1\}$ -classifier we know this metrics coming from the confusion matrix:

- precision = PPV = positive predicted value: $\frac{n_{1,1}}{n_{1,1}} \Rightarrow \text{precision} \uparrow \text{ false positive } \downarrow$
- APER= $\frac{n_{1,0}+n_{0,1}}{n}$
- recall = sensitivity = $\frac{n_{1,1}}{n_{1,\cdot}} \Rightarrow \text{ recall } \uparrow \text{ false negative } \downarrow$
- specificity = $\frac{n_{0,0}}{n_{\cdot,0}}$



To improve quality of $\hat{AER}(\delta)$ we can:

- · Construct confusion matrix on a test set;
- Use cross-validation on $\mathbb X$ to extract from it the test set (Leave-One-Out) and repeat m-times;
- k-fold cross validation: split $\mathbb X$ into k subset and use cross validation inside each subset, repeat it B-times.

Using k-fold cross validation we are decreasing the correlation between models trained so the variance of estimators given by this technique is lower than the one given by the cross-validation Leave-One-Out.

Support Vector Machine #SVM

Main contributor to this theory was Vapnick in '90.

Framework: We have a dychotomous problem and X is our dataset.

Idea: Find an hyperplane s.t. separate points which belongs into group 1 to ones belonging group 2.

What is an hyperplane?

 \mathcal{L} := hyperplane in \mathbb{R}^p := affine subspace of dimension p-1

So $\exists \beta \in \mathbb{R}^p$ with $||\beta|| = 1$ s.t. $\beta \perp \mathcal{L}$

Let be $\underline{x}_0 = span\{\underline{\beta}\} \cap \mathcal{L}$ and $\beta_0 = ||\underline{x}_0|| \leftarrow \text{so where } \beta \text{ and } \mathcal{L} \text{ meet}$

Given $\underline{x} \in \mathbb{R}^p$ we know that:

$$\underline{x} \in \mathcal{L} \quad \Leftrightarrow \quad \pi_{\underline{x}|\underline{eta}} = \underline{x}_0 = rac{\underline{eta} \cdot \underline{eta}'}{\underline{eta}' \cdot \underline{eta}} \cdot \underline{x} \quad \leftrightarrow \quad \underline{eta}' \cdot \underline{x} = eta_0$$

Question 1: Given $A,B\in\mathbb{R}^p$ when can I separate them with an hyperplane?

Let CH(A), CH(B) be the convex-hull generated by A and B.

Assumptions (Geometric Hahn-Banach theorem):

- $CH(A) \neq \emptyset$, $CH(B) \neq \emptyset$;
- $CH(A) \cap CH(B) = \emptyset$
- Either CH(A) or CH(B) is open
 - \Rightarrow \exists separating hyperplane

Obs:

 3^{rd} assumption could be rewrite as:

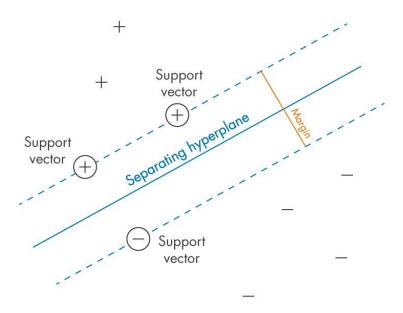
CH(A) and CH(B) are closed and at least one of them is compact

Question 2: Find the best hyperplane \mathcal{L}

Assume \mathbb{X} is s.t. $\exists \mathcal{L}$ separating classes 1 from classes 2.

We define \mathbb{R}^+ and \mathbb{R}^- the half planes divided by $\mathcal{L}.$





Let $\underline{y}_i = \pm 1$ and if it positive \underline{x}_i belongs to class 1 otherwise belongs to class 2. Define $\mathrm{Margin} = M_1 := \min_{\forall i=1,\dots,n} \{\underline{y}_i \cdot (\underline{\beta}' \cdot \underline{x}_i - \underline{\beta}_0)\}$

The best \mathcal{L} is that one which maximize M_1 .

The problem to solve is:

$$egin{cases} oldsymbol{max}_{\underline{eta},\underline{eta}_0} M \ igg| |\underline{eta}|| = 1 \ \underline{y}_i \cdot (\underline{eta}' \cdot \underline{x}_i - \underline{eta}_0) \geq M \end{cases} \quad orall i$$

This problem is the most strict version of our problem so is called *hard problem*.

Solution:

$$\hat{eta} = \sum_{i=1}^n \hat{\lambda}_i y_i \underline{x}_i, \quad \hat{eta}_0 \in \mathbb{R} \quad \Rightarrow \quad \hat{f}(\underline{x}) = \hat{\underline{eta}}' \underline{x} - \hat{eta}_0 = \sum_{i=1}^n \hat{\lambda}_i y_i \underline{x}_i' \cdot \underline{x}_i - \hat{eta}_0$$

So our classifier became $c(x) = sign\{\hat{f}(x)\}$

For that we define $support := \{points which find the hyperplane\} \subset X$

If our data are so mixed we have 3 approach to try to find a classifier:

• Using soft problem, so the last request in the hard problem becomes

$$\underline{y}_i \cdot (\underline{eta}' \cdot \underline{x}_i - \underline{eta}_0) \geq (1 - \epsilon_i) M \qquad orall i$$

with $\epsilon_i \geq 0$ and $\sum_{i=0}^n \leq c$ where c is called *budget constrain*.

Interpretation: we are admitting that some point of one group could be in the half space of the other and viceversa.

- Transforming our data, or our axis, in such a way that the hyperplane become clear.
- Instead of using $\underline{x}_i' \cdot \underline{x}_i$ we can define a new kernel $k(\underline{x},\underline{w}) : \mathbb{R}^p \times \mathbb{R}^p \to \mathbb{R}$ so we can write $\hat{f}(\underline{x}) = \sum_{i=1}^n \hat{\lambda}_i y_i k(\underline{x}_i,\underline{x}_i) \hat{\beta}_0$

Interpretation: kernel function are the tool with we transform our data such that is more clear the dividing hyperplane.



Example - kernels

- d-degree polynomials = $k(x, w) = (1 + x'w)^d$
- radial-basis function = $k(\underline{x}, \underline{w}) = \exp\{-\gamma ||\underline{x} \underline{w}||^2\}$
- $k(\underline{x},\underline{w}) = tanh\{k_1 \cdot \underline{x}'\underline{w} + k_2\} \leftarrow$ common in neural networks

Unsupervised Learning - #Cluster-analysis

It is useful when given \mathbb{X} neither labels associate or the number of groups in the dataset.

Idea: unit in the same cluster are more similar than units in different clusters.

Firstly, we need a measure of similarity/dissimilarity. It should have the following properties:

- $d(\underline{x},\underline{x}) = 0$ $\forall \underline{x} \in \mathbb{R}^p$
 - Stronger version: $d(\underline{x},y)=0 \Leftrightarrow \underline{x}=\underline{y}$
- $d(\underline{x},y)=d(y,\underline{x})$ $orall \underline{x},y\in\mathbb{R}^p$
- $d(\underline{x},\underline{y}) \leq d(\underline{x},\underline{z}) + d(\underline{z},\underline{y}) \qquad orall \underline{x},\underline{y},\underline{z} \in \mathbb{R}^p$
 - $\bullet \ \ \text{Stronger version:} \ d(\underline{x},\underline{y}) \leq max\{d(\underline{x},\underline{z}),d(\underline{z},\underline{y})\} \qquad \forall \underline{x},\underline{y},\underline{z} \in \mathbb{R}^p$

If our similarity function has all the stronger version of each condition it is an **ultra metrics**. If our similarity function has all conditions and the first one in the stronger version it is a **metrics**.

If our similarity function has all conditions in their base form it is a **pseudo metrics**.

Example - Ultrametric

Let
$$\underline{x} = \{\alpha, \beta, \gamma, \sigma, \epsilon, \theta\}, \quad y = \{\alpha, \beta, \gamma, \omega, \nu, \mu\}$$

Define $d(\underline{x},\underline{y}) = \frac{1}{2^n}$ where n is the index of the first element which differs in the objects. In this case is 4.

Some trivial distances with $\underline{x},y\in\mathbb{R}^p$:

- <code>#Euclidean_distance</code> : $d(\underline{x},\underline{y}) = \sqrt{(x-y)'(x-y)}$
- <code>#Mahalanobis_distance</code> : $d(\underline{x},\underline{y}) = \sqrt{(x-y)'\Sigma^{-1}(x-y)}$
- <code>#Minkowsky_distance</code> : $d^r(\underline{x},y) = \sum_{i=1}^p |x_i y_i|^r$
- ullet #Camberra_distance ullet : $d(ar{x},ar{y}) = \sum_{i=1}^p rac{|x_i-y_i|}{x_i+y_i}$

Some trivial distances with $\underline{x},\underline{y} \in \{0,1\}^p$:

• #Euclidean_distance : $d(\underline{x},\underline{y})=\sqrt{(x-y)'(x-y)}$ (which indicates the number of discordances)

From contingency matrix:

$$egin{array}{cccc} 0 & 1 \\ 0 & a & b \\ 1 & c & d \end{array}$$

• $d(\underline{x},\underline{y})=1-\frac{a}{p}$ where p is the total number of observations



```
• d(\underline{x}, y) = \sqrt{2 \cdot (1 - corr(x, y))}
```

It's very useful to create the D matrix with distances between each x_i and all other x_i .

- \Rightarrow *D* has the following properties:
 - Each element on the diagonal is 0;
 - It's symmetric if d is symmetric and reflexive.

Distances between finite subset of \mathbb{R}^p

Let U, V be two finite subset (cluster) of \mathbb{R}^p .

How to calculate d(U, V)?

Possible solutions are:

- #Single_linkage : $d(U,V) = \min\{d(\underline{x},y): \underline{x} \in U, y \in V\}$
- #Complete_linkage : $d(U,V) = \max\{d(\underline{x},y): \underline{x} \in U, y \in V\}$
- #Average_linkage : $d(U,V) = rac{1}{\#U\cdot\#V} \sum_{\underline{x}\in U, y\in V} d(\underline{x},\underline{y})$
- #Centroid_distance : $d(U,V)=d(\underline{c}_U,\underline{c}_V)$ with \underline{c}_U and \underline{c}_V the centroid of U and V

Hierarchical Agglomerative Cluster Algorithms

Let D be the distances matrix and chosen a linkage $\#Hierarchical_Agglomerative_Cluster$ works in this way:

```
Initialization: each unit is a Cluster
while(!convergence condition reached){
     cluster together 2 closest clusters;
     update D;
}
```

Let $\mathbb X$ with $\underline x_i\in\mathbb R^p$ be the training set and choose $d:\mathbb R^p\times\mathbb R^p\to[0,+\infty)$ as pseudo-metric, metric and ultra-metric.

Hierarchical Clustering - #Ward_method

Hypothesis: d must be the <code>#Euclidean_distance</code>, so we use: $d^2(\underline{x},\underline{y}) = (\underline{x}-\underline{y})'(\underline{x}-\underline{y})$ We call C a cluster of finite point in \mathbb{R}^p , it is the area where we classify the point of \mathbb{X} with a certain label.

Since we want to find: $\forall i \quad \underline{\overline{x}}_i = argmin_{\underline{x} \in \mathbb{R}^p} \sum_{\underline{x}_i \in C_i} d^2(\underline{x}, \underline{x}_i) = \frac{1}{|C_i|} \sum_{\underline{x}_i \in C_i} \underline{x}_i = \text{barycentre of } C_i$ From that we can define the $\text{\#error_sum_squares}$ $ESS_i = \sum_{\underline{x} \in C_i} d^2(\underline{x}, \overline{\underline{x}}_i)$ which describe the variability of all the data \underline{x} in our cluster C_i around the barycentre $\overline{\underline{x}}_i$.

So we can create a cumulative index called $ESS = \sum_i ESS_i$

```
Initialization: each unit is a cluster so ESS = 0
while(!convergence condition reached){
```



```
merge together 2 clusters which imply the minimum ESS increase;
}
```

Not Hierarchical Clustering

Goal: cluster $\mathbb X$ in k clusters s.t. $\mathbb X=\cup_i^k C_i$ and $orall i\neq j$ $C_i\cap C_j=\emptyset$

#K-means

It's a generalization of #Ward method with d different from the euclidean.

Calculate $\overline{\underline{x}}_i = argmin_{\underline{x} \in \mathbb{R}^p} \sum_{x_i \in C_i} d^2(\underline{x},\underline{x}_i) = \text{centroid of } C_i$

Goal: find C_i s.t. $min\{\sum_i^k \sum_{\underline{x}_i \in C_i} d^2(\underline{x}, \overline{\underline{x}}_i)\} = min\{\sum_i^k ESS_i\}$

```
Initialization: create k random centroid
or
Initialization: create k random subset

While(centroid change){
        Compute centroid for each cluster;
        Assign every unit to cluster with the nearest centroid;
}
```

Since the computation of centroid is a difficult problem we can simplify this step calculating medoids so become $\underline{x}_i = argmin_{\underline{x} \in \mathbb{X}} \sum_{\underline{x}_i \in C_i} d^2(\underline{x}, \underline{x}_i)$ and the algorithm change also names: k-means \rightarrow k-medoids

How to choose k?

Secchi's tips:

- If you have a dendrogram choose a cut-point s.t. clusters are robust to some oscillation of it, don't choose it near a split;
- Express the cost function $ESS = \sum_i^k \sum_{\underline{x}_i \in C_i} d^2(\underline{x}, \overline{\underline{x}}_i)$ in function of k and choose the k with the elbow rule (if the increasing of k is longer than the decrease of the cost function it's time to stop)

Problems

Using Hierarchical or Not-Hierarchical cluster we have some problems:

- #Complete_linkage , #Average_linkage , #K-means , #Ward_method tend to create ellipsoidal clusters in the metric choosen;
- #Single_linkage has the problem of the chain effect;

Non-parametric density-based clustering

What is neighbourhood of x in \mathbb{R}^p ?



We call:

 $\epsilon ext{-neighbourhood of }\underline{x}=N_\epsilon(\underline{x})=\{y\in\mathbb{R}^p:d(\underline{x},y)<\epsilon\}$

Which is a sphere in the sense of d chosen.

We also define $|N_{\epsilon}(\underline{x})| = \#\{ ext{points } \in \mathbb{X}, \in N_{\epsilon}(\underline{x}) \}$

Density-Based Spatial Clustering of Applications with Noise

#DBSCAN

Idea: cluster are region with high density of points, low density areas are boundaries between clusters, noises or outliers.

We have to fix in advance:

- Radius $\epsilon > 0$;
- $minPts \in \mathbb{N} \setminus \{0\}$

After that we have to define some names:

- x is called *core-point* if $|N_{\epsilon}(x)| \geq minPts$;
- x is called *border-point* if $|N_{\epsilon}(\underline{x})| < minPts$ but $\exists \underline{x}_i$ core-point s.t. $\underline{x}_i \in N_{\epsilon}(\underline{x}_i)$;
- *x* is called *outlier* in every other cases.

As last definitions we need:

- \underline{x}_i is directly density reachable from \underline{x}_i if: $\underline{x}_i \in N_{\epsilon}(\underline{x}_i)$ with \underline{x}_i a core-point;
- \underline{x}_j is *density reachable* from \underline{x}_i if there are $\underline{y}_{i\in\{1,\dots,k\}}$ with $k\geq 0$ s.t.:
 - $\underline{y}_{i\in\{1,\dots,k-1\}}$ are core points;
 - $\underline{y}_{\scriptscriptstyle 1} = \underline{x}_i$ and $\underline{y}_k = \underline{x}_j$
- \underline{x}_j is density connected from \underline{x}_i if there is an \underline{x} s.t. both \underline{x}_i and \underline{x}_j are density reachable from \underline{x}

DBSCAN identifies a cluster $C \subset \mathbb{X}$ s.t.:

- 1. if \underline{x}_j is density reachable from $\underline{x}_i \in C \Rightarrow \underline{x}_j \in C$;
- 2. $\forall \underline{x}_i, \underline{x}_j \in C \quad \underline{x}_i, \underline{x}_j$ must be density connected.

MultiDimensional Scaling #MDS

Create the distance matrix D

Goal: create a new dataset $\tilde{\mathbb{X}}$ with $\underline{y}_i \in \mathbb{R}^q$ and q << p where we can use <code>#Euclidean_distance</code> and $d_{euclidean}(\underline{y}_i,\underline{y}_j) = \delta_{i,j} \simeq d_{ij} \in D$

There are two approches to solve this problem:

- Classical MDS: find $\underline{y}_{i \in \{1,...,k\}} = argmin \sum_{i
 eq j} (d_{i,j} \delta_{i,j})^2$
- $\bullet \ \textit{Kruskal MDS:} \ \text{find} \ \underline{\underline{y}}_{i \in \{1, \dots, k\}} = \underset{\text{stress function}}{\underbrace{\sum_{i \neq j} (\theta(d_{i,j}) \delta_{i,j})^2}} \ \text{with} \ \theta : \mathbb{R} \to \mathbb{R} \ \text{monotone}$



Remark: in the classical MDS, if D is done with euclidian distance we can use #PCA to capture the first q variables which better describe the variability of our data. So become the same thing.

Regression

Recall

We call target variable $y \in \mathbb{R}$ while vector of features $\underline{x} \in \mathbb{R}^p$.

Goal: explain y variability as f(x).

Remember that: $\# regression_function = \mathbb{E}[y|\underline{x}] : \mathbb{R}^p \to \mathbb{R}$

Rmk: if $(y,\underline{x}) \sim N_{p+1} \Rightarrow \mathbb{E}[y|\underline{x}] = \beta_0 + \beta_1 x_1 + \ldots$ is linear to the regressor;

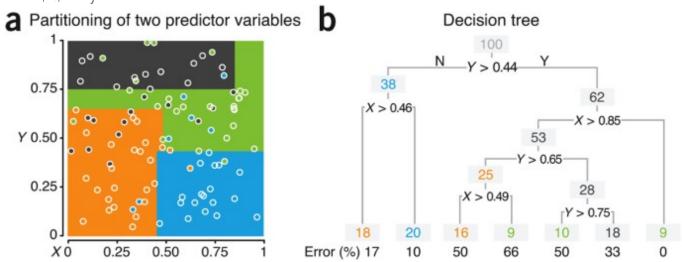
The general model is $y = \mathbb{E}[y|\underline{x}] + \epsilon$ and we want to find $\hat{f}(\underline{x})$ to estimate the regression function.

There main categories of method to do it:

- totally data driven: we haven't a model of f, is very good for prediction but really bad to interpret and is often used in data-mining and machine learning approaches. (CART)
- *structured model:* we know how looks like *f* and we use data to find unknown parameters of the model chosen.

Classification And Regression Trees #CART

Idea: split features space \mathbb{R}^p into j partitions R_1,R_2,\ldots,R_j and predict y in R_i as $\overline{y}_i=\frac{1}{|R_i|}\sum_{\underline{x}_j\in R_i}y_i$



The main difficult is to find partitions R_1, R_2, \dots, R_j with following properties:

- $\cup_{k=1}^j R_k = \mathbb{R}^p$;
- $R_i \cap R_j = \emptyset \quad \forall i \neq j;$
- $R_i \cap \mathbb{X}
 eq \emptyset \quad orall i o$ to ensure the existence of the mean \overline{y}_i

Optimal Criterion: find j and $R_1, R_2, \ldots, R_j = argmin\{\sum_{k=1}^j \sum_{\underline{x}_i \in R_k} (y_i - \overline{y}_i)^2\}$

This optimal criteria is NP-complete so CART suppose that R_1,R_2,\ldots,R_j are rectangles.

We have to introduce the following function depending of the split (s_k^*) :



$$u(s_k^*) = \underbrace{\sum_{i=1}^n (y_i - \overline{y}_k)^2}_{ ext{variance pre-split}} - \underbrace{[\sum_{x_{k,i} > s_k^*} (y_i - \overline{y}_k^+)^2 + \sum_{x_{k,i} < s_k^*} (y_i - \overline{y}_k^-)^2]}_{ ext{variance post-split}}$$

```
foreach feature i{
        find split associate to feature i which maximize u;
}
choose best split, the one that maximizes u;
split feature space in 2 partition;
repeat for each partition if stopping criteria isn't reached.
```

Typical stopping criteria: $|R_i| < threshold$ where R_i is the partition where the algorithm restart in.

This method is very good because can be read by anyone since create a binomial tree. A problem of this algorithm is that it's easy to overfit our training set. To avoid this problem it's useful to modify $u(s_k^*)$ adding a penalization to the number of partitions j:

$$u(s_k^*) = \sum_{k=1}^j \sum_{i=1}^n (y_i - \overline{y}_k)^2 + lpha \cdot j$$

where α is choose with cross validation.

Rmk: without any constrain to j the model will overfit training data putting $j = |\mathbb{X}|$ and creating a partition with only 1 point;

Rmk: work also for categorical value simply our split from a threshold become a subset of categorical variables analyzed.

Linear Model for Regression

We call each feature as x_i where i indicate the column number representing that feature. Recall that:

$$\mathbb{X} = egin{bmatrix} ar{x_1'} & y_1 \ \dots & \dots \ ar{x_n'} & y_n \end{bmatrix} ext{with } ar{x_i'} = (x_{i,1}, x_{i,2}, \dots, x_{i,p})$$

From that we can create a matrix to store each transformation of the raw data contained in \mathbb{X} , it is called $\#\text{design_matrix} \in n \times (r+1)$:

$$\mathbb{Z} = egin{bmatrix} 1 & z_{1,1} & \dots & z_{1,r} \ \dots & \dots & \dots & \dots \ 1 & z_{n,1} & \dots & z_{n,r} \end{bmatrix}$$



We call each feature of our design matrix as $z_i = h_i(x_1, x_2, \dots, x_p)$ where $i \in \{1, \dots, r\}$ indicate the column number representing that feature and $h_i(x_1, x_2, \dots, x_p)$ is the transformation of the raw-features.

Without supposing Gaussianity, this type of models is linear in z_i so we find that the #regression_function is:

$$\mathbb{E}[y|x_1,\ldots,x_p] = eta_0 + eta_1 h_1(x_1,x_2,\ldots,x_p) + eta_2 h_2(x_1,x_2,\ldots,x_p) + \cdots + eta_r h_r(x_1,x_2,\ldots,x_p)$$

From that we assume the general model of y is:

$$y = eta_0 + eta_1 z_1 + \dots + eta_r z_r + \epsilon \quad \Rightarrow \quad \boxed{\underline{y} = (y_1, \dots, y_n)' = \mathbb{Z} \underline{eta} + \underline{\epsilon}}$$

We know that $\mathbb{E}[\underline{\epsilon}] = \underline{0}$ and $cov(\underline{\epsilon}) = \sigma^2 I$ with I the identity matrix so each ϵ_i is uncorrelated to others.

Example - #ANOVA

Our starting point is the ANOVA system of equations:

$$\left\{egin{array}{ll} x_{1,1},\ldots,x_{1,n_1} & iid \sim N(\mu_1,\sigma^2) \ x_{2,1},\ldots,x_{2,n_2} & iid \sim N(\mu_2,\sigma^2) \ \ldots \ x_{g,1},\ldots,x_{g,n_g} & iid \sim N(\mu_g,\sigma^2) \end{array}
ight.$$

Where each row is \bot to others and $n=n_1+n_2+\cdots+n_g$ We know that $\underline{y}=(x_{1,1},\ldots,x_{1,n_1},\ldots,x_{g,1},\ldots,x_{g,n_g})\in\mathbb{R}^n$ So our design matrix become:

$$egin{bmatrix} egin{bmatrix} 1 & 1 & 0 & 0 & \dots \ 1 & \dots & 0 & \dots & \dots \ 1 & 1 & 0 & 0 & \dots \ \end{bmatrix} \ egin{matrix} 1 & 1 & 0 & 0 & \dots \ 1 & 0 & 1 & 0 & \dots \ 1 & 0 & 1 & 0 & \dots \ \end{bmatrix} \in n imes (g+1) \ egin{matrix} 1 & 0 & 0 & 1 & \dots \ 1 & \dots & \dots & \dots \end{bmatrix}$$

Where the ones in the first column are exactly n_1 , in the second n_2 and so on. We can also define $\underline{\beta}=(\mu,\tau_1,\ldots,\tau_g)$ and $\underline{\epsilon}\sim N(\underline{0},\sigma^2I)$ with I the identity matrix. So we reach that:

$$\underline{y} = \mathbb{Z} \underline{eta} + \underline{\epsilon} \Leftrightarrow egin{cases} x_{i,j} = \mu + au_i + \epsilon_{i,j} \ \epsilon_{i,j} \quad iid \sim N(0,\sigma^2) \end{cases}$$

We know that this problem is over-parametrized, due to the fact that $\mathbb Z$ has not full rank, so we have to add the following constrain: $\sum_{i=1}^g n_i \tau_i = 0$

From the constrained we deduce that $au_g = -\sum_{i=1}^{g-1} rac{n_i}{n_g} au_i$ so we can rewrite the design matrix as:



$$egin{bmatrix} 1 & 1 & 0 & 0 & \dots \ 1 & \dots & 0 & \dots & \dots \ 1 & 1 & 0 & 0 & \dots \ 1 & 0 & 1 & 0 & \dots \ 1 & \dots & \dots & \dots & \dots \ 1 & 0 & 1 & 0 & \dots \ 1 & 0 & 0 & 1 & \dots \ 1 & \dots & \dots & \dots & \dots \ 1 & \dots & \dots & \dots & \dots \ 1 & \dots & \dots & \dots & \dots \ 1 & rac{n_2}{n_g} & rac{n_3}{n_g} & rac{n_4}{n_g} & \dots \ 1 & \dots & \dots & \dots \ 1 & rac{n_2}{n_g} & rac{n_3}{n_g} & rac{n_4}{n_g} & \dots \ \end{bmatrix}$$

Estimate β and σ^2 :

Remember: #ordinary_least_squared
$$:=\hat{\underline{\beta}} = rgmin ||\underline{y} - \mathbb{Z}\underline{\beta}||^2$$

Prop

If \mathbb{Z} is full rank ($\leftrightarrow \operatorname{rank}(\mathbb{Z}) = r + 1 \le n$)

Then:

Proof - previous Prop

Starting points:

- $\mathbb{Z}'\mathbb{Z} \in (r+1) imes (r+1)$
- $\mathbb{Z}'\mathbb{Z}=\sum_{i=1}^{r+1}\lambda_i\underline{e}_i\underline{e}_i'$ with $\lambda_1\geq\lambda_2\geq\cdots\geq\lambda_{r+1}>0$
- If $\mathbb Z$ isn't full rank else $\lambda_{r+1}=0$

$$\begin{array}{l} \Rightarrow (\mathbb{Z}'\mathbb{Z})^{-1} = \sum_{i=1}^{r+1} \frac{1}{\lambda_i} \underline{e}_i \underline{e}_i' \\ \text{Define } \forall i \in \{1, \dots, r+1\} \quad \underline{q}_i = \frac{1}{\sqrt{\lambda_i}} \mathbb{Z} \underline{e}_i \end{array}$$

So:

$$\begin{cases} \forall i \quad \underline{q}_i \in \mathcal{L}(\mathbb{Z}) \\ \\ \underline{q}_i' \underline{q}_j = \frac{1}{\sqrt{\lambda_i \lambda_j}} \underline{e}_i' \mathbb{Z}' \mathbb{Z} \underline{e}_j & \stackrel{\mathbb{Z}'\mathbb{Z}}{\underline{e}_j = \lambda_j \underline{e}_j} \\ \\ 1 \quad i = j \end{cases} \Rightarrow \underline{q}_1, \dots, \underline{q}_{r+1} \text{ is an orthonormal basis of } \mathcal{L}(\mathbb{Z})$$

$$\Rightarrow \pi_{\underline{y}|\mathbb{Z}} = \sum_{i=1}^{r+1} \underbrace{\frac{\underline{q}_i q_i'}{\underline{q}_i' \underline{q}_i}}_{\underline{q}_i \underline{q}_i'} \underline{y} = \sum_{i=1}^{r+1} \underline{q}_i \underline{q}_i' \underline{y} = \sum_{i=1}^{r+1} \frac{1}{\lambda_i} \mathbb{Z} \underline{e}_i \underline{e}_i' \mathbb{Z}' \underline{y} = \mathbb{Z} (\sum_{i=1}^{r+1} \frac{1}{\lambda_i} \underline{e}_i \underline{e}_i') \mathbb{Z}' \underline{y} = \mathbb{Z} \underline{\hat{\beta}} = \underbrace{\mathbb{Z}} (\mathbb{Z}'\mathbb{Z})^{-1} \mathbb{Z}' \underline{y}$$

$$\Rightarrow \underline{\hat{y}} = H \underline{y} \Rightarrow \underline{\hat{\beta}} = (\mathbb{Z}'\mathbb{Z})^{-1} \mathbb{Z}' \underline{y}$$



Remark:

- $rank(\mathbb{Z}) = k < r+1 \le n : \lambda_k > 0$ while $\lambda_{k+1} = \cdots = \lambda_{r+1} = 0$ so $\not\exists (\mathbb{Z}'\mathbb{Z})^{-1}$ but we can use Moore-Pensore Inverse $(\mathbb{Z}'\mathbb{Z})^-$ called also Generalized Inverse
- $rank(\mathbb{Z})=n=r+1:\mathcal{L}(\mathbb{Z})=\mathbb{R}^n$ so $y=\hat{y}\Rightarrow H=I\Rightarrow$ interpolating data

framework: \mathbb{Z} is full rank

We know that $H = \mathbb{Z}(\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'$ is the <code>#orthogonal_projection_operator</code> on $\mathcal{L}(\mathbb{Z})$. So I - H is the orthogonal projection operator on $\mathcal{L}^{\perp}(\mathbb{Z})$ where $dim(\mathcal{L}^{\perp}(\mathbb{Z})) = n - (r+1)$

We find that: $\underline{\hat{\epsilon}} = (I - H) y \Rightarrow \underline{\hat{\epsilon}} \perp \hat{y}$

So we reach the #1_decomposition_of_variance :

$$||y||^2 = SS_{tot} = ||\hat{y}||^2 + ||\hat{\underline{\epsilon}}||^2 = SS_{regression} + SS_{residuals}$$

recall: $\pi_{y|\underline{1}} = \overline{y} \cdot \underline{1}$

Starting from: $\pi_{\hat{y}|\underline{1}}=\frac{\underline{1}\underline{1}'}{\underline{1}'\underline{1}}\hat{\underline{y}}=\frac{\underline{1}\underline{1}'}{\underline{1}'\underline{1}}H\underline{y}$

Remember that orth. proj. oper. is symmetric so: $\underline{1}'H = (H'\underline{1})' = (H\underline{1})' = \underline{1}'$

We reach that: $\pi_{\hat{y}|\underline{1}} = \frac{\underline{1}\underline{1'}}{\underline{1'}\underline{1}}\underline{y} = \overline{y} \cdot \underline{1} = \pi_{y|\underline{1}}$ so \underline{y} and $\hat{\underline{y}}$ have the same mean

From the previous result we arrive at $\#2_decomposition_of_variance$ using the Corrected Sum of Squares (CSS):

$$||\underline{y} - \overline{y} \cdot \underline{1}||^2 = CSS = ||\hat{\underline{y}} - \overline{y} \cdot \underline{1}||^2 + ||\hat{\underline{\epsilon}}||^2 = CSS_{regression} + CSS_{residuals}$$

So we can define #coefficient_of_determination as:

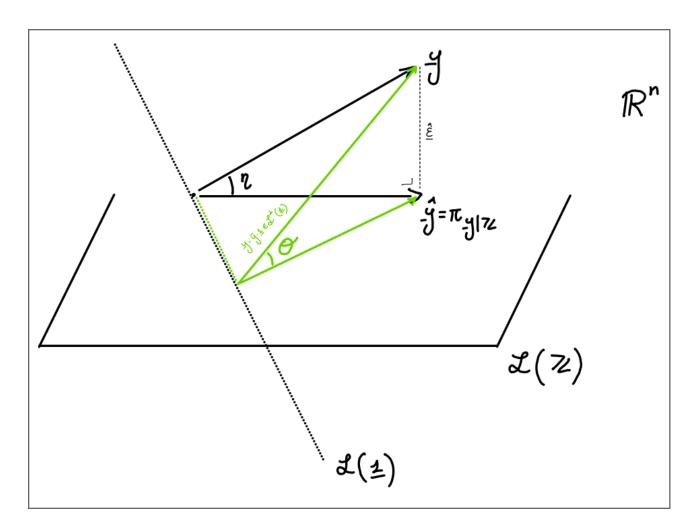
$$R^2 = 1 - rac{\sum_i \hat{oldsymbol{arepsilon}}_i^2}{\sum_i (oldsymbol{y}_i - \hat{oldsymbol{y}}_i)^2} = 1 - rac{||\hat{oldsymbol{arepsilon}}||^2}{||\hat{oldsymbol{y}}||^2} = 1 - rac{SS_{residuals}}{SS_{regression}} = 1 - \sin^2 heta = \cos^2 heta$$

Interpretation: portion of variability explained by \hat{y}

Remarks:

- All this deduction are valid since the first column of \mathbb{Z} is $1\Rightarrow \mathbb{E}[\hat{y}]=\mathbb{E}[y]$
- From 1^{st} decomposition of variance we can deduce $ilde{R}^2=1-rac{||\hat{\epsilon}||^2}{||y||^2}=\cos^2\eta$
- $R^2_{adjusted}=1-rac{\sum_i\hat{\epsilon}_i^2}{n-(r+1)}:rac{\sum_i(\underline{y}_i-\hat{\underline{y}}_i)^2}{n-1}$ is R^2 adjusted by the degree of freedom, it's a measure of how good is a model fitted





Prop

- $R^2=1\Rightarrow heta=0\Rightarrow y=\hat{y}\Rightarrow$ Interpolation;
- $R^2=0\Rightarrow heta=rac{\pi}{2}\Rightarrow \underline{y}\perp \mathcal{L}(\mathbb{Z})\Rightarrow \hat{\underline{y}}=\overline{y}\cdot \underline{y}$

Properties β and $\hat{\epsilon}$:

Framework:

- \mathbb{Z} full rank $\Leftrightarrow rank(\mathbb{Z}) = r + 1$
- $\hat{eta}=(\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'y$
- $\hat{\underline{\epsilon}} = (I H)\underline{y} \Rightarrow \hat{\underline{\epsilon}} \perp \hat{\underline{y}}$

Prop

- 1. $\mathbb{E}[\hat{\underline{\beta}}] = \underline{\beta} \leftarrow \text{unbiased}$
- 2. $cov(\hat{\beta}) = \sigma^2(\mathbb{Z}'\mathbb{Z})^{-1}$
- 3. $\mathbb{E}[\hat{\epsilon}] = \underline{0}$
- 4. $cov(\hat{\epsilon}) = \sigma^2(I H) \neq \sigma^2 I = cov(\hat{\epsilon})$
- $5. \ \mathbb{E}[\underline{\hat{\epsilon}}'\underline{\hat{\epsilon}}] = \mathbb{E}[||\underline{\hat{\epsilon}}||^2] = [n-(r+1)]\sigma^2 \Rightarrow \mathbb{E}[\tfrac{\hat{\epsilon}'\hat{\epsilon}}{n-(r+1)}] = \sigma^2 \leftarrow \text{unbiased}$

Proof - previous Prop



1.
$$\begin{cases} \mathbb{E}[\hat{\underline{\beta}}] = \mathbb{E}[(\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'\underline{y}] = (\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'\mathbb{E}[\underline{y}] \\ \mathbb{E}[\underline{y}] = \mathbb{E}[\mathbb{Z}\underline{\beta} + \underline{\epsilon}] = \mathbb{E}[\mathbb{Z}\underline{\beta}] \end{cases} \Rightarrow \mathbb{E}[\hat{\underline{\beta}}] = (\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'\mathbb{Z}\underline{\beta} = \underline{\beta}$$

$$2. \ cov(\underline{\hat{\beta}}) = (\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'\underbrace{cov(\underline{y})}_{=\sigma^2I}\mathbb{Z}(\mathbb{Z}'\mathbb{Z})^{-1} = \sigma^2(\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'\mathbb{Z}(\mathbb{Z}'\mathbb{Z})^{-1} = \sigma^2(\mathbb{Z}'\mathbb{Z})^{-1}$$

3.
$$\mathbb{E}[\hat{\underline{\epsilon}}] = \mathbb{E}[(I - H)\underline{y}] = \overbrace{(I - H)}^{\in \mathcal{L}^{\perp}(\mathbb{Z})} \underbrace{\mathcal{E}(\mathbb{Z})}_{\mathcal{Z}} \underline{\beta} = 0$$

4.
$$cov(\hat{\underline{\epsilon}}) = (I - H)cov(\underline{y})(I - H)' = \sigma^2(I - H)(I - H)' \xrightarrow{\text{symmetric} + \text{idempotent}} = \sigma^2(I - H) \text{ with } det(I - H) = 0 \text{ since is an } n \times n \text{ matrix which project on a space of dimension } n - (r + 1)$$

5. Recall:

1)
$$\hat{\epsilon} = (I - H)y = (I - H)(\mathbb{Z}\beta + \epsilon) = (I - H)\epsilon$$

2)
$$tr(H) = tr(\mathbb{Z}'(\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}) = tr(\mathbb{Z}'\mathbb{Z}(\mathbb{Z}'\mathbb{Z})^{-1}) = tr(I_{(r+1)\times(r+1)}) = r+1$$

We can reach:

$$\mathbb{E}[\underline{\hat{\epsilon}}'\underline{\hat{\epsilon}}] = tr(\mathbb{E}[\underline{\hat{\epsilon}}'\underline{\hat{\epsilon}}]) = \mathbb{E}[tr(\underline{\hat{\epsilon}}'\underline{\hat{\epsilon}})] = \mathbb{E}[tr(\underline{\hat{\epsilon}}\underline{\hat{\epsilon}}')]$$

$$\overset{1^{st} \text{ recall}}{\Rightarrow} \ \mathbb{E}[tr((I-H)\underline{\epsilon}\underline{\epsilon}'(I-H)')] = tr((I-H)\mathbb{E}[\underline{\epsilon}\underline{\epsilon}'](I-H)) = \overbrace{tr(\sigma^2(I-H))}^{= \dim(\text{space into proj})} = \sigma^2(n-tr(H))$$

Obs

- $cov(\hat{\underline{\beta}}) = \sigma^2(\mathbb{Z}'\mathbb{Z})^{-1} o \text{usually } \hat{\underline{\beta}}_i \text{ aren't uncorrelated unless } \mathbb{Z}'\mathbb{Z} = \alpha I \text{ with } \alpha \in \mathbb{R}$
- When designing an experiment we can control variability of \hat{eta} controlling $\mathbb Z$

Change of paradigm $\epsilon \sim N(0,\sigma^2 I)$

Prop

Adding gaussianity assumption to ϵ bring us some nice properties:

1.
$$\hat{\underline{\beta}}$$
 and $\hat{\sigma}^2 = \frac{\hat{\underline{\epsilon}}'\hat{\underline{\epsilon}}}{n}$ are **MLE** estimator of $\underline{\epsilon}$ and σ^2 ;

2.
$$\hat{eta} \sim N_{r+1}(eta, \sigma^2(\mathbb{Z}'\mathbb{Z})^{-1})$$

3.
$$\hat{\underline{\epsilon}} \sim N_n(\underline{0}, \sigma^2(I-H))$$

4.
$$\hat{\epsilon} \perp \!\!\!\perp \hat{\beta}$$

5.
$$\hat{\underline{\epsilon}}'\hat{\underline{\epsilon}} \sim \sigma^2 \chi(n-(r+1))$$

Proof - previous Prop

Since as assumption we know that $\underline{\epsilon} \sim N(\underline{0}, \sigma^2 I) \Rightarrow y \sim N(\mathbb{Z}\beta, \sigma^2 I)$

1. Write likehood and do differentiation;



$$\begin{bmatrix} \underline{\hat{\beta}} \\ \underline{\hat{\epsilon}} \end{bmatrix} = \underbrace{\begin{bmatrix} (\mathbb{Z}'\mathbb{Z})\mathbb{Z} \\ I-H \end{bmatrix}}_{A} \underline{y} \sim N(\begin{bmatrix} \underline{\beta} \\ \underline{0} \end{bmatrix}, \begin{bmatrix} (\mathbb{Z}'\mathbb{Z})^{-1} & \underline{0} \\ \underline{0} & I-H \end{bmatrix}) = N(A\mathbb{Z}\underline{\beta}, \sigma^2AIA')$$

$$2. \ \underline{\hat{\epsilon}} \sim N_n(\underline{0}, \sigma^2(I-H)) \Rightarrow \underline{\hat{\epsilon}}' \underline{\hat{\epsilon}} = d^2(\underline{\hat{\epsilon}}, \underline{0}) = \frac{1}{\sigma^2} (\underline{\hat{\epsilon}} - \underline{0})' \underbrace{(I-H)^-}_{\text{Penrose Inverse}} (\underline{\hat{\epsilon}} - \underline{0}) \sim \chi^2(n - (r+1))$$

Framework: \mathbb{Z} is full rank and $\underline{\epsilon} \sim N(0, \sigma^2 I)$

Recall: $S^2=rac{\hat{\epsilon}'\hat{\epsilon}}{n-(r+1)}$

Goal: find $CI(\beta)$ one-at-time

Obs

• (R) ∐ (S)

From the previous observation we find:

$$rac{(\hat{eta}-eta)'(\mathbb{Z}'\mathbb{Z})(\hat{eta}-eta)}{r+1}rac{n-(r+1)}{\hat{\epsilon}'\hat{\epsilon}}\sim F(r+1,n-(r+1)) \ \Leftrightarrow rac{1}{S^2}(\hat{eta}-eta)'(\mathbb{Z}'\mathbb{Z})(\hat{eta}-eta)\sim (r+1)F(r+1,n-(r+1)) \ \end{cases}$$

From this we can construct the confidence interval at level α as:

$$CI_{1-lpha}(eta)=\{\mu\in\mathbb{R}^{r+1}: (\hat{eta}-\mu)'(\mathbb{Z}'\mathbb{Z})(\hat{eta}-\mu)\leq (r+1)S^2F_{1-lpha}(r+1,n-(r+1))\}$$

So we can say that taking $\underline{a} \in \mathbb{R}^{r+1}$ we reach: $\underline{a}' \hat{\underline{\beta}} \sim N_1(\underline{a}' \underline{\beta}, \sigma^2 \underline{a}' (\mathbb{Z}' \mathbb{Z})^{-1} \underline{a})$ This implies that:

$$\boxed{\frac{\underline{a'(\hat{\beta}-\underline{\beta})}}{\sigma\sqrt{\underline{a'}\mathbb{Z'}\mathbb{Z}\underline{a}}}\cdot\frac{\sqrt{\sigma^2}}{\sqrt{S^2}} = \frac{\underline{a'(\hat{\beta}-\underline{\beta})}}{S\sqrt{\underline{a'}\mathbb{Z'}\mathbb{Z}\underline{a}}} \sim t(n-(r+1)) \leftarrow \text{ Pivotal quantity}}$$

$$\Rightarrow CI_{1-\alpha}(\underline{a}'\underline{\beta}) = \{\underline{a}'\underline{\hat{\beta}} \pm S\sqrt{\underline{a}'\mathbb{Z}'\mathbb{Z}\underline{a}} \cdot t_{1-\frac{\alpha}{2}}(n-(r+1))\} \text{ which is one-at-time} \\ \Rightarrow CI_{1-\alpha}(\beta_i) = \{\hat{\beta}_i \pm S\sqrt{diag_i(\mathbb{Z}'\mathbb{Z})} \cdot t_{1-\frac{\alpha}{2}}(n-(r+1))\} \text{ with } \underline{a} = [0,\ldots,0,1,0,\ldots,0] \text{ which is one-at-time}.$$

WARNING: not use this to create a $CI(\beta)$ because aren't simultaneously



Goal: find $CI(\beta)$ simultaneously

$$\max_{\underline{a} \in \mathbb{R}^{r+1}} rac{[\underline{a}'(\hat{eta}-eta)]^2}{S^2(a'\mathbb{Z}'\mathbb{Z}a)} = rac{1}{S^2}(\hat{eta}-eta)'(\mathbb{Z}'\mathbb{Z})(\hat{eta}-eta) \sim (r+1)F(r+1,n-(r+1))$$

So we can construct the confidence interval at level α as:

$$CI_{1-lpha}(\underline{a}'\underline{eta}) = [\underline{a}'\hat{\underline{eta}} \pm S(\underline{a}'\mathbb{Z}'\mathbb{Z}\underline{a})\sqrt{(r+1)F_{1-lpha}(r+1,n-(r+1))}]$$

Goal: find $CI(\sigma^2)$

$$rac{1}{\sigma^2} \hat{\underline{\epsilon}}' \hat{\underline{\epsilon}} = rac{[n-(r+1)]S^2}{\sigma^2} \sim \chi^2(n-(r+1))$$

So we can construct the confidence interval at level α as:

$$CI_{1-lpha}(\sigma^2) = \left[rac{[n-(r+1)]S^2}{\chi^2_{1-rac{lpha}{2}}(n-(r+1))}; rac{[n-(r+1)]S^2}{\chi^2_{rac{lpha}{2}}(n-(r+1))}
ight]$$

Framework: $C \in p imes (r+1)$, contains linear combinations of eta

Goal: test $H_0: C\underline{\beta} = 0$ vs $H_1: C\beta \neq 0$

Since we know that: $C\hat{eta} \sim N_p(Ceta, \sigma^2C(\mathbb{Z}'\mathbb{Z})^{-1}C)$

We reach:

$$rac{rac{1}{\sigma^2}(C\hat{eta})'(C(\mathbb{Z}'\mathbb{Z})^{-1}C)^{-1}(C\hat{eta})}{p}\cdotrac{\sigma^2}{S^2}\sim F(p,n-(r+1))$$

So we can construct our test statistics F such that:

$$F = rac{(C\hat{eta})'(C(\mathbb{Z}'\mathbb{Z})^{-1}C)^{-1}(C\hat{eta})}{S^2} \sim p \cdot F(p,n-(r+1))$$

Reject H_0 if at level lpha if $F>p\cdot F_{1-lpha}(p,n-(r+1))$

Goal: test $H_0: \beta_r = \beta_{r-1} = \cdots = \beta_{r-(p-1)} = 0$ vs $H_1: else$

This test con be rewritten as the case before using as C:



$$C = egin{bmatrix} 0 & \dots & 0 & 1 & 0 & 0 & \dots & 0 \ 0 & \dots & 0 & 0 & 1 & 0 & \dots & 0 \ 0 & \dots & 0 & 0 & 0 & 1 & \dots & 0 \ & dots \ \end{bmatrix} = [0 \quad I_{p imes p}] \in p imes (r+1)$$

With this is like to say that $\mathbb{Z} = [\mathbb{Z}_1, \mathbb{Z}_2]$ and we want to test our reduced model \mathbb{Z}_1 against the full model \mathbb{Z} , since we are testing that betas of \mathbb{Z}_2 are zeros.

**We reject H_0 if $SS_{res}(\mathbb{Z}_1) - SS_{res}(\mathbb{Z})$ is large knowing that

$$rac{SS_{res}(\mathbb{Z}_1) - SS_{res}(\mathbb{Z})}{S^2p} \sim F(p, n - (r+1))$$

Special case: when we want to test: $H_0: eta_1=eta_2=\cdots=eta_r=0$ vs $H_1:else$

So our

$$\mathbb{Z}_1 = egin{bmatrix} 1 \ dots \ 1 \end{bmatrix} \Rightarrow SS_{res}(\mathbb{Z}_1) = \sum_1^n (y_i - \overline{y})^2 \Rightarrow rac{SS_{res}(\mathbb{Z}_1) - SS_{res}(\mathbb{Z})}{S^2 r} = rac{\sum_1^n (y_i - \overline{y})^2 - \sum \hat{\epsilon_i}^2}{S^2 r} \sim F(r, n - (r+1))$$

This test is done by default as F-test in R packages.

Goal: Prediction of y

We have a model for the phenomenon which is $y_0 = \underline{Z}_0' \underline{\beta} + \epsilon_0$ with \underline{Z}_0 the vector collecting the "weights" of each regressor.

We know that the prediction $\mathbb{E}[y_0|\underline{Z}_0]=\underline{Z}_0'\underline{\beta}$ which we know its unbiased predictor is $\underline{Z}_0'\underline{\hat{\beta}}$

Theo - Gauss-Markov

 $\underline{Z'_0}\hat{\underline{\beta}}$ is the Best Linear Unbiased Estimator (<code>#BLUE</code>) of $\underline{Z'_0}\underline{\beta}$

Framework: we change the paradigm, now $cov(\underline{\epsilon})=\sigma^2\Sigma$ with $\Sigma\in n\times n\to to$ model heteroscedasticity + correlations between ϵ_i and ϵ_j

Case 1 - Σ known but not σ^2

We use #Generalized_least_squared to find $\hat{\beta}$.

$$\underline{\hat{\beta}} = \underset{\underline{\beta}}{argmin}[(\underline{y} - \mathbb{Z}\underline{\beta})'\Sigma^{-1}(\underline{y} - \mathbb{Z}\underline{\beta})] \leftarrow \text{ minimize Mahalanobis distance}$$

We do the following change of variables:

$$\begin{split} &(\underline{y} - \mathbb{Z}\underline{\beta})'\Sigma^{-1}(\underline{y} - \mathbb{Z}\underline{\beta}) = (\underline{y} - \mathbb{Z}\underline{\beta})'\Sigma^{-\frac{1}{2}}\Sigma^{-\frac{1}{2}}(\underline{y} - \mathbb{Z}\underline{\beta}) = ||\underline{\tilde{y}} - \tilde{\mathbb{Z}}\underline{\beta}||^2 & \text{with } \underline{\tilde{y}} = \Sigma^{-\frac{1}{2}}\underline{y} \text{ and } \widetilde{\mathbb{Z}} = \Sigma^{-\frac{1}{2}}\mathbb{Z} \\ &\Rightarrow \hat{\underline{\beta}} = \underset{\underline{\beta}}{argmin}||\underline{\tilde{y}} - \tilde{\mathbb{Z}}\underline{\beta}||^2 \\ &\Rightarrow \hat{\beta} = (\widetilde{\mathbb{Z}}'\widetilde{\mathbb{Z}})^{-1}\widetilde{\mathbb{Z}}'\tilde{y} = (\mathbb{Z}'\Sigma^{-1}\mathbb{Z})^{-1}\mathbb{Z}'\Sigma^{-1}y \end{split}$$

Idea: \tilde{y} is a change of the reference system in a way such that $cov(\underline{\tilde{\epsilon}}) = \Sigma^{-\frac{1}{2}}cov(\underline{\epsilon})\Sigma^{-\frac{1}{2}} = \sigma^2 I$



Case 2 - Σ and σ^2 unknown

Possible solutions:

- 1. parameterize Σ , we will see this idea in Linear Mixed Model;
- 2. Estimate Σ iteratively from residual using as starting point $\Sigma = I$;
- 3. Use appropriate transformation of y and/or \mathbb{Z} ;

Example 1 - Σ known

 $y_i = \mathrm{mean} \; \mathrm{of} \; n_i$ independent observations with the same variance σ^2

So we know that
$$var(y_i)=rac{\sigma^2}{n_i}$$
 and we can model $\Sigma=egin{bmatrix} rac{1}{n_1} & 0 & \dots & 0 \\ 0 & rac{1}{n_2} & \dots & 0 \\ \dots & \dots & \dots \\ 0 & 0 & \dots & rac{1}{n_n} \end{bmatrix}$

Someone called this Weighted Least Squared (#WLS).

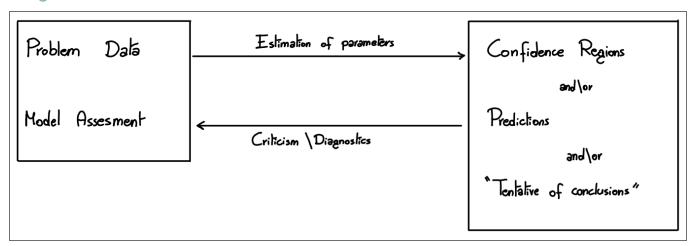
Example 2 - Σ known

 $y_i = \text{sum of } n_i$ independent observations with the same variance σ^2 (like GDP of each states)

So we know that
$$var(y_i) = \sigma^2 n_i$$
 and we can model $\Sigma = egin{bmatrix} n_1 & 0 & \dots & 0 \\ 0 & n_2 & \dots & 0 \\ \vdots & \ddots & \ddots & \ddots \\ 0 & 0 & \dots & n_n \end{bmatrix}$

Another example of #WLS

Diagnostics for a linear model



General way to do diagnostic is to control the following stuff:

- residuals analysis, outliers, heteroscedasticity, normality test, auto-correlation;
- Influential cases;
- collinearity;

Residuals analysis

We have:

• Abstract model $o \underline{y}=\mathbb{Z} \underline{\beta}+\underline{\epsilon}$ with $\mathbb{E}[\underline{\epsilon}]=0$ and $cov(\epsilon)=\sigma^2 I$



• Fitted model $o \hat{y} = \mathbb{Z}\hat{eta} + \hat{\underline{\epsilon}}$ with $\mathbb{E}[\hat{\epsilon}] = 0$ and $cov(\hat{\underline{\epsilon}}) = \sigma^2(I - H)$

Problem: If in the abstract model $\underline{\epsilon} \sim N_n(\underline{0}, \sigma^2 I)$, which is a distribution on \mathbb{R}^n then in the fitted model $\underline{\epsilon} \sim N(0, \sigma^2 (I - H))$ which is a distribution in $\mathcal{L}^{\perp}(\mathbb{Z})$.

Someone prefers to do analysis on the studentized residuals of unit ${\sf i}=rac{\hat{\epsilon}_i}{S\sqrt{1-h_{i,i}}}$ with

 $h_{i,i} = diag(H)_{i,i} = ext{ leverages } \in (0,1) ext{ since } H ext{ is a projector matrix.}$

So if $h_{i,i} \uparrow \Rightarrow var(\hat{\epsilon}_i) \downarrow 0$ and also if $\mathbb{E}[\hat{\epsilon}_i] = 0$ then $\hat{\epsilon}_i = 0$

#Cook_distance and influential cases

We call \mathbb{Z}_{-i} which is \mathbb{Z} without the i-row \Rightarrow we can write the abstract model as $\underline{y}_{-i} = \mathbb{Z}_{-i}\underline{\beta}_{-i} + \underline{\epsilon}_{-i}$ and also we can estimate with $\underline{\hat{\beta}}_{-i}$.

To see if a unit is influential we can compare $\hat{\beta}$ and $\hat{\beta}_{-i}$ and to compare it we use the Cook distance:

$$D_i = rac{(\hat{eta} - \hat{eta}_{-i})(\mathbb{Z}'\mathbb{Z})(\hat{eta} - \hat{eta}_{-i})}{S^2(r+1)} = (rac{\hat{\epsilon_i}}{S\sqrt{1-h_{i,i}}})^2 rac{h_{i,i}}{1-h_{i,i}} rac{1}{r+1} \sim F(r+1,n-(r+1))$$

Rule of thumb: delete unit with $D_i > 1$

Model selection

Since with r regressors we can have 2^r models to find the best one we can:

```
for(k = 1:r){
    fit all possible models with k regressors;
    choose best one (using GOF indicator)
}
```

Why collinearity is a problem?

Since from <code>#ordinary_least_squared</code> we have found that:

•
$$\hat{\underline{\beta}}=(\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}'\underline{y}$$
 if \mathbb{Z} is full rank

If there are collinear regressors, tend to be $\mathbb{Z}'\mathbb{Z}$ singular so the inversion of it explode

$$\Rightarrow \quad var(\hat{eta}) = \sigma^2(\mathbb{Z}'\mathbb{Z})^{-1} ext{ significantly increases}$$

Curios is to find that:

$$var(\hat{eta}_j) = rac{\sigma^2}{\sum_{i=1}^n (z_{i,j} - \overline{z}_j)^2} \cdot rac{1}{1 - R_j^2}$$
 with R_j^2 as `#coefficient_of_determination` when z_j is regressed from $z_{i \in \{1, \ldots, r\} \setminus \{j\}}$

Remark:
$$\sum_{i=1}^n (z_{i,j} - \overline{z}_j)^2 \uparrow \Rightarrow var(\hat{\beta}_j) \downarrow$$

Remark: If regressors are orthogonal than $R_j^2=0$ $\forall j$ otherwise $R_j^2\uparrow 1\Rightarrow var(\hat{\beta}_j)\uparrow$ caused by collinearity.



Tips: red alarm when $VIF = \text{variance inflation factor} = \frac{1}{1-R_s^2} \geq 5$

Solution: To limit this problem we adding the constrain of keep $var(\hat{\beta})$ low

New nomenclature

We centered our data, so we reformulate our #ordinary_least_squared as:

$$\begin{array}{lll} \bullet & \underline{y} \rightarrow \underline{y}^* = \underline{y} - \overline{y} \cdot \underline{1} \\ & & \begin{bmatrix} z_{1,1} - \overline{z}_1 & z_{1,2} - \overline{z}_2 & \dots & z_{1,r} - \overline{z}_r \\ z_{2,1} - \overline{z}_1 & z_{2,2} - \overline{z}_2 & \dots & z_{2,r} - \overline{z}_r \end{bmatrix} \\ \bullet & \mathbb{Z} \rightarrow \mathbb{Z}^* = & \vdots & \vdots & \vdots \\ & \vdots & \vdots & \vdots & \vdots \\ & z_{n,1} - \overline{z}_1 & z_{n,2} - \overline{z}_2 & \dots & z_{n,r} - \overline{z}_r \end{bmatrix} \in n \times r \\ \bullet & & \\ & \hat{\beta}^* = \arg \min_{\underline{\beta} \in \mathbb{R}^r} ||\underline{y}^* - \mathbb{Z}^* \underline{\beta}||^2 \\ \bullet & & \\ & \hat{\beta}_0 = \overline{y} - \sum_{i=1}^r \hat{\beta}_i^* \overline{z}_i \end{array}$$

CAUTION: So from now on we will call y^* as y and \mathbb{Z}^* as \mathbb{Z} .

#Ridge_regression

Useful reference: link

Add a parameter and change (§) into:

$$egin{cases} rac{\hat{eta}_{Ridge}}{||ar{eta}||^2} = rg \ min_{ar{eta} \in \mathbb{R}^r} ||ar{y} - \mathbb{Z} \underline{eta}||^2 \ ||ar{eta}||^2 \leq s \end{cases}$$

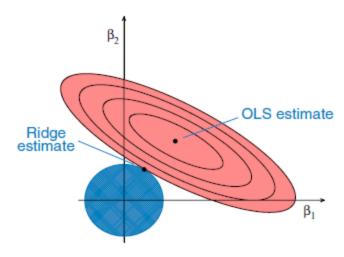
Relation Between OLS and Ridge Regression

But knowing that $\underline{\hat{y}} = H\underline{y} = \mathbb{Z}(\mathbb{Z}'\mathbb{Z})^{-1}\mathbb{Z}\underline{y} = \mathbb{Z}\underline{\hat{\beta}}_{OLS}$ we can simplify find that:

$$\begin{split} ||\underline{y} - \mathbb{Z}\underline{\beta}||^2 &= ||\underline{y} - \underline{\hat{y}} + \underline{\hat{y}} - \mathbb{Z}\underline{\beta}||^2 \\ &= ||\underline{y} - \underline{\hat{y}} - \mathbb{Z}(\underline{\beta} - \underline{\hat{\beta}}_{OLS})||^2 \\ &= ||\underbrace{\hat{\epsilon}}_{\underline{\hat{\epsilon}}} - \mathbb{Z}(\underline{\beta} - \underline{\hat{\beta}}_{OLS})||^2 \\ &= ||\underline{\hat{\epsilon}}||^2 - ||\mathbb{Z}(\underline{\beta} - \underline{\hat{\beta}}_{OLS})||^2 \\ &\Rightarrow \begin{cases} \underline{\hat{\beta}}_{Ridge} &= \arg\min_{\underline{\beta} \in \mathbb{R}^r} |||\mathbb{Z}(\underline{\beta} - \underline{\hat{\beta}}_{OLS})||^2 \\ ||\underline{\beta}||^2 \leq s \end{split}$$



So we can see it as:



How to find Ridge regressors

To find it we have to use lagrangian: $\arg \ min_{eta \in \mathbb{R}^r} ||\underline{y} - \mathbb{Z}\underline{\beta}||^2 + \lambda ||\underline{\beta}||^2$

So we reach: $\hat{\underline{eta}}_{Ridge} = (\mathbb{Z}'\mathbb{Z} - \lambda I)^{-1}\mathbb{Z}'\underline{y}$

Problem: $\hat{\underline{\beta}}_{Ridge}$ is biased because of λ in the lagrangian

But Hoerl&Kennard in 1970 find that:

 $\forall \text{ regression problem } \exists \lambda^* : \mathbb{E}[||\hat{\underline{\beta}}_{Ridge} - \underline{\beta}||^2] \leq \mathbb{E}[||\hat{\underline{\beta}}_{OLS} - \underline{\beta}||^2] \text{ with } \lambda^* \text{ found using cross validation}$

#PCA_regression

Another way to manage collinearity is to use #PCA on \mathbb{Z} to find $k \leq r$ orthogonal regressors **Problem:** PCA regression and Ridge regression doesn't support sparse solution in term of z_1, \ldots, z_r , so they only limit the effect of some regressor but not eliminate it

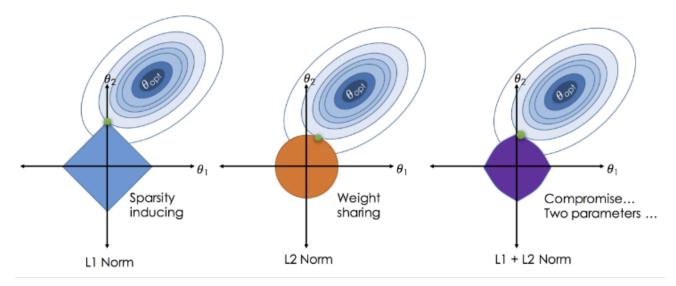
Intuition Tibshirami (1996)

Use Ridge regression but change norm on which limit the $\underline{\beta}$. In this way, we are also selecting regressors since someones become 0.

So it become:

$$egin{cases} \hat{eta} = rg \ min_{oldsymbol{eta} \in \mathbb{R}^r} || \underline{y} - \mathbb{Z} \underline{eta} ||^2 \ || \underline{eta} ||_1 \leq s \end{cases}
ightarrow ext{which is called LASSO}$$





From this we can spike the area of search of our $\hat{\beta}$. Usually, much is spike our area and more difficult is the resolution of the lagrangian.

Note: $\arg min_{eta \in \mathbb{R}^r} ||y - \mathbb{Z}\beta||^2 + \lambda_1 ||eta||_1 + \lambda_2 ||eta||_2^2$ is called Elastic Net

Since now we have studied that, to do regression\classification we have to:

- Take a training set X;
- Fit a model on that training set $M_{\mathbb{X}}$;
- Give a new features vector \underline{x} predict with $M_{\mathbb{X}}(\underline{x})$

Ensemble methods

Problem: we have to deal with the bias-variance trade off

Q: how to reduce variance without dealing with bias?

If is true that $\mathbb{E}[\epsilon] = 0$ we can take B independent measurements y and use the mean of these to find the real value μ since we know that:

$$\overline{y} = rac{1}{B} \sum_{i=1}^B y_i \quad \Rightarrow \quad \mathbb{E}[\overline{y}] = \mu ext{ and } var(\overline{y}) = rac{1}{B} \sigma^2 \quad \Rightarrow \quad B \uparrow ext{ then } var(\overline{y}) \downarrow$$

So we are reducing variance without adding a bias.

Idea: generate B training set \mathbb{X}_i and fit B models $M_{\mathbb{X}_i}$ and take as final model $M = \frac{1}{B} \sum_{i=1}^B M_{\mathbb{X}_i}$ **Cons:**

- Boring and costly to manually fit ${\it B}$ models;
- If I have access to B training set why to not use it fit a unique big model?

So, usually we have a single training set and we want to build multiples of it.

Idea: Use #Bagging which is the composition of #bootstrap and aggregate the data obtained in dataset.

What is bootstrap?

It is a simple algorithm:



```
Initialization: training set of n units
for(i in 1:n){
     sample randomly one unit from the training set;
}
```

What is bagging?

It is very similar to bootstrap but we aggregate samples into a new training set.

```
Initialization: training set of n units
for(i in 1:n){
     sample randomly one unit from the training set;
     add the sampled unit to a new training;
}
```

In this way, the new training set \mathbb{X}^* contains some of the units of the original \mathbb{X} but with some copies, so $\mathbb{X}^* \subset \mathbb{X}$.

```
\mathbb{P}[u 
otin \mathbb{X}^*]? \mathbb{P}[u 
otin \mathbb{X}^*] = (1 - \frac{1}{n})^n \xrightarrow{n \to +\infty} e^{-1} \approx \frac{1}{3}
```

So we have approximately $\frac{2}{3}$ of the original units of \mathbb{X} in \mathbb{X}^* , remaining observations are called Out-Of-Bag observations #OOB.

This causes that $\mathbb{X}^* \not\perp \mathbb{X}$ so we reach that using the strategy of taking the mean model M we reach that: $var(M) \in (\frac{\sigma^2}{B}, \sigma^2)$

Framework: we use #Regression_tree as model

Idea: To increase independence between models we don't search among each features and cut the best one but for each model we sample randomly k < p features and do the best cut in that k features. This increase independence between model so we reduce variance.

From that come out #random forest

And to see performance of each model we can use its set of #OOB observations.

Boosting

This technique work fitting sequentially decision trees each one on the residual of the previous tree.

