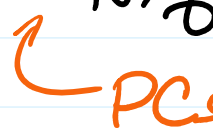


Principal Components Regression (PCR)

Instead of regressing Y onto $X_{N \times p}$,

regress Y onto $Z_{N \times q}$


Steps for PCR

① mean center X

$$X_c = \begin{bmatrix} X_1 - \hat{\mu}_1 & X_2 - \hat{\mu}_2 & \dots & X_p - \hat{\mu}_p \end{bmatrix}$$

$$\hat{\mu}_i = \text{mean}(X_i)$$

② do PCA on $X_c = U D V^T$

$$Z = X_c V_{1:q}$$

② regress Y onto Z
(typically include intercept)

$$D = \begin{bmatrix} 1 & Z \\ 1 & 1 \end{bmatrix}$$

$$\hat{\beta}^{(PCR)} = (D^T D)^{-1} D^T Y \in \mathbb{R}^{q+1}$$

What about new data?

let X^{test} be $M \times P$

For training: $\hat{Y} = D \hat{\beta}^{(PCR)}$

Need to calc. D^{test} by apply some procedure to X^{test}

① mean center X^{test}

$$X_c^{\text{test}} = \begin{bmatrix} X_1^{\text{tot}} - \hat{\mu}_1 & \dots & X_p^{\text{tot}} - \hat{\mu}_p \end{bmatrix}$$

$$\hat{\Lambda}_c = [\hat{\Lambda}_1 - \mu_1$$

$$, \dots, \mu_p]$$

from
training data

① apply PCA:

$$Z^{\text{test}} = X_c^{\text{test}} V_{1:q}$$

from
training

$$\textcircled{2} D^{\text{test}} = \begin{bmatrix} 1 & Z^{\text{test}} \\ 1 & \end{bmatrix}$$

$$\textcircled{3} \hat{y}^{\text{test}} = D^{\text{test}} \hat{\beta}^{\text{(PCR)}}$$

from train.

Compare PCR w/ ridge regression

$$\hat{y}^{\text{(ridge)}} = X \hat{\beta}^{\text{(ridge)}}$$

\vdots

1, 2, ...

T...

up scaling
lots of Z

$$= \sum_{j=1}^p \underbrace{\left(\frac{\sigma_j^2}{\sigma_j^2 + \lambda} \right)}_{\Delta_j} u_j u_j^T \gamma$$

Can also show:

$$\hat{Y}^{(PCR)} = Z \hat{\beta}^{(PCR)} = \dots = \sum_{j=1}^p \Delta_j u_j u_j^T \gamma$$

$$\Delta_j = \begin{cases} 1, & j \leq q \\ 0 & j > q \end{cases}$$

$$\text{OLS: } \Delta_j = 1$$

$$\text{ridge: } \Delta_j = \frac{\sigma_j^2}{\sigma_j^2 + \lambda}$$

$$\text{PCR: } \Delta_j = \mathbb{1}(j \leq q)$$

Consider X to be full rank

Then as $\lambda \rightarrow 0$ we have

Then as $\lambda \rightarrow 0$ we have

$$\hat{\beta}^{(\text{ridge})} \rightarrow \hat{\beta}^{(\text{OLS})}$$

Similarly, as $q \rightarrow P$ then

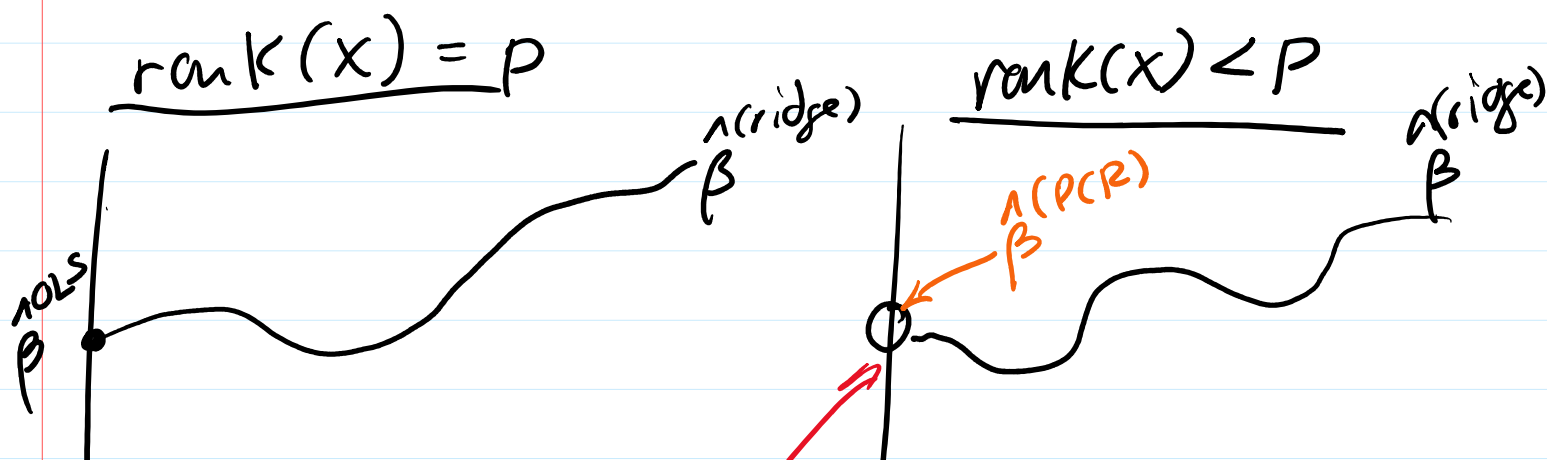
$$\hat{\beta}^{(\text{PCR})} \rightarrow \hat{\beta}^{(\text{OLS})}$$

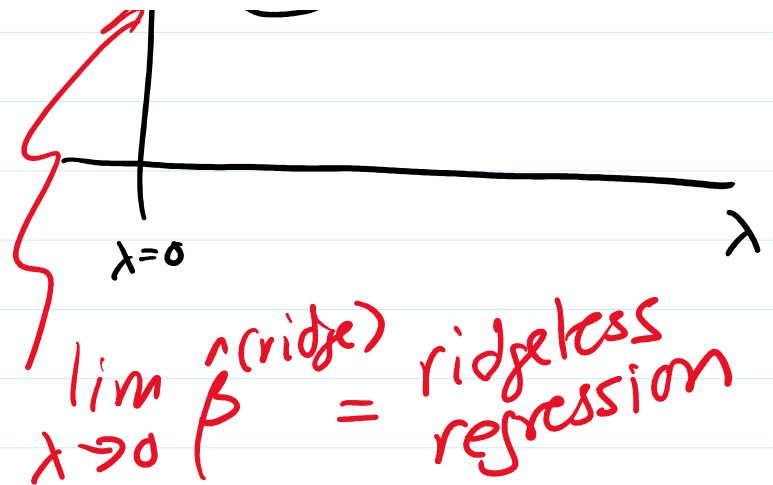
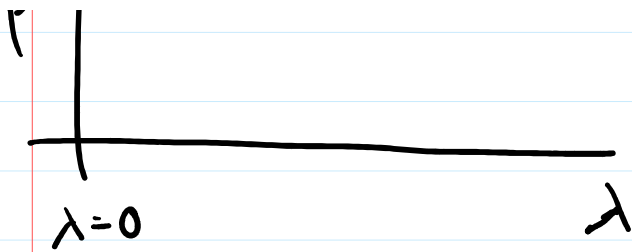
If $\text{rank}(X) < P$

Then OLS doesn't exist.

(Equiv. ridge w/ $\lambda = 0$ doesn't exist)

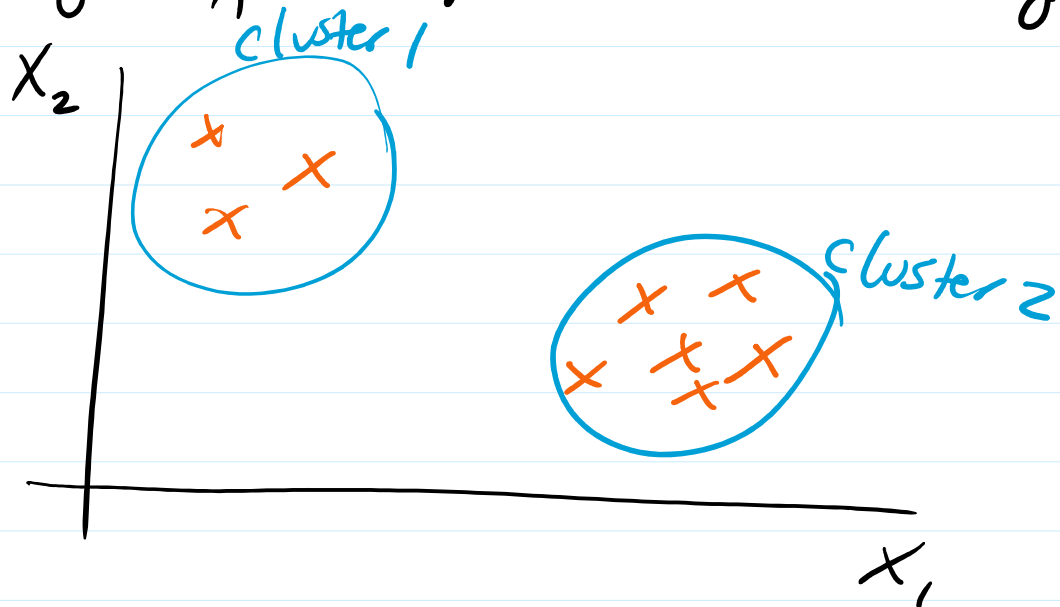
However $\lim_{\lambda \rightarrow 0} \hat{\beta}^{(\text{ridge})} = \hat{\beta}^{(\text{PCR})}$
w/ $q = \text{rank}(X)$





Back to unsupervised

Second type of unsupervised: clustering



To find clusters I need to define how pts are similar/dissim to each other

Dissimilarity Measure

Dissimilarity Measure

If I have N obs then I can define a dissim mtrx

$$D \quad (N \times N)$$

where

$D_{ii'}$ = dissim meas betwn
obs. i and i' .

For many clustering methods - don't
need X but only D .

Properties of D :

- ① $D_{ii} = 0$
 - ② $D_{ii'} \geq 0$
 - ③ $D = D^T$
-

K-means clustering

Assume that each data pt belongs to
one of K clusters (or groups):

Assume that each data pt belongs to one of K clusters (or groups):

$$G_1, \dots, G_K$$

want: assign each point i to some cluster G_k

how: want to make assignments so that we minimize some measure of not being well clustered (loss)

classic way to measure this is

$W =$ total within-cluster dissimilarity

$$= \sum_{k=1}^K \sum_{i, i' \in G_k} D_{ii'}$$

↑ small if clustering is good
large // bad

$$T = \text{total dissim} = \sum_{i, i'} D_{ii'}$$

$B =$ total between
cluster dissim

$$= \sum_{k \neq k'} \sum_{i \in G_k} \sum_{i' \in G_{k'}} D_{ii'}$$

Can show: $T = B + W$

So, to find G_1, \dots, G_K

① minimize W

or ② maximize B

Ideally: try all possible cluster
assignments

practically: not possible —
too computationally
demanding

Assume, all data is numeric: $x_i \in \mathbb{R}^p$

Assume all data is numeric: $x_n \in \mathbb{R}^p$

define

$$D_{ii'} = \|x_i - x_{i'}\|_2^2$$

If I do this then

$$W = \sum_{k=1}^K 2N_k \sum_{i \in G_k} \|x_i - \bar{x}_k\|^2$$

dist of pt to mean of cluster

$N_k = |G_k|$

mean of pts in cluster k

Goal: try to minimize this W

Lloyd's Algorithm

① Initialization Step

make some random initializations of cluster means:

$$\mu_1^{(0)}, \mu_2^{(0)}, \dots, \mu_K^{(0)}$$

For $t = 1, 2, 3, \dots$

at the t^{th} iteration :

① Assignment Step:

assign i (or x_i) to cluster G_k
w/ the closest mean to x_i

$$G_k^{(t)} = \{x_i : \|x_i - \mu_k^{(t)}\| \leq \|x_i - \hat{\mu}_{k'}^{(t)}\| \quad \forall k' \neq k\}$$

② Update Step: re-calculate μ_s
given current assignments

$$\hat{\mu}_k^{(t+1)} = \text{mean } x_i \quad i \in G_k^{(t)}$$
