PCA

Two ways to derive PCA

- D Maximize variance after projection
- 1 Minimize Reconstruction error

We show D and D are the same

D X E IR " > data matrix, each row a data point

Assume data points are centered, i.e.

 $\overline{X} = \overline{1} \overline{1} \overline{X} = \overline{0}$, 1: a vector of ones

Projection of X onto $\vec{V} \in \mathbb{R}^p$: $X\vec{V} \in \mathbb{R}^n$

Sample mean of $X\vec{V}$: $\frac{1}{N}\vec{V} = \vec{X}\vec{V} = \vec{0}\vec{V} = 0$

Sample variance of $X\vec{v} : \frac{1}{n-1}(X\vec{v})^T(X\vec{v}) = \frac{1}{n-1}\vec{v}^TX^TX\vec{v}$

Find I that maximizes the variance.

 $\max_{V} \overrightarrow{V} \overrightarrow{V} \times \overrightarrow{V} s.t. \overrightarrow{V} \overrightarrow{V} = 1$

Solution: solve eigenvalue problem $X^T \times V = \lambda V - \omega$ Among all (λ, V) that satisfies D, choose the one whose λ is the largest since $V^T \times X^T = \lambda V^T = \lambda$

$$X = \begin{bmatrix} \vec{X}_1 \\ \vec{X}_1 \end{bmatrix} \Rightarrow \text{ that data point}$$

$$X = \begin{bmatrix} \vec{X}_1 \\ \vec{X}_2 \end{bmatrix} \Rightarrow \text{ nth data point}$$

$$(X\overrightarrow{V})_{i} = \overrightarrow{X_{i}}\overrightarrow{V}$$
 is the projection of $\overrightarrow{X_{i}}$ ento \overrightarrow{V} $(\overrightarrow{X}\overrightarrow{V})_{i}\overrightarrow{V}$ is the reconstruction of $\overrightarrow{X_{i}}$ using \overrightarrow{V} Reconstruction error: $\sum_{i=1}^{n} ||\overrightarrow{X_{i}} - (\overrightarrow{X}\overrightarrow{V})_{i}\overrightarrow{V}||^{2}$

$$=\sum_{i=1}^{N}\left(\overline{x_{i}}^{T}\overline{X_{i}}-2(\overline{x_{i}}^{T})\overline{x_{i}}^{T}\overline{V}+((\overline{x_{i}}^{T})\overline{y_{i}}^{T}\overline{V}\right)$$

We require

$$\frac{n}{\sum |\vec{x}|} = \frac{n}{\sum |$$

So, minimizing the Reconstruction error is the same

as
$$\lim_{N \to \infty} - \sum_{i=1}^{n} ((x\vec{r})_i)^2 = -\|x\vec{r}\|^2$$
$$= -\|x\vec{r}\|^2$$
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Which is equivalente to

Laplacian Eigenmap and Speitral Clustering n data points xi, xi, ..., xr

A & IR "x", AT = A an affinity matrix

Aij: similarity between Xi and Xz

D: a diagenal matrix such that Dir = \(\overline{\pi_1} \) Aij

Laplacian L = D-A

Laplacian Figenmap

min $\sum A_{ij}(t_i-t_j) = f^T L f$ one dimensional

S.t. + TD + = 1 representation of data points

Spectral Clustering (Relaxed Normalized Cut)

 $\min \frac{f^T L f}{f^T D f} = 0$

Obtain cluster assignments by thresholding f

Require FTDf=1 (remove arbitrary scaling)

Require FDI=0 (f=I is a trivial solution)

Symetrically normalized Laplacian:

$$\mathcal{L} = D^{-\frac{1}{2}} L D^{\frac{1}{2}}$$
 (Assume D'exists)

min yTy sit. yTy=1

Eet

Vorable transformation:

$$D^{-\frac{1}{2}}y = f \implies y = D^{\frac{1}{2}}f$$

$$Well-cletined if D^{-1} exists.$$

Then

$$y^{T} \sum_{j=1}^{N} y^{j} = f^{D^{\frac{1}{2}}} p^{\frac{1}{2}} \sum_{j=1}^{N} p^{\frac{1}{2}} p^{\frac{1}{2}} = f^{T} p^{\frac{1$$

So, when D'exists, there is a one-to-one mapping between the unnormalized and normalized formulations.