

PCA & SVD

Note Title

Recall the centered data matrix

$$\tilde{X} := [\tilde{x}_1 \ \cdots \ \tilde{x}_n] \in \mathbb{R}^{d \times n}$$

$$\tilde{x}_j := x_j - \bar{x}, \quad \bar{x} := \frac{1}{n} \sum_{i=1}^n x_i,$$

and the sample covariance matrix

$$S := \frac{1}{n} \tilde{X} \tilde{X}^T$$

Then, PCA is nothing but the eigen decomposition of S

$$S = \Phi \Lambda \Phi^T, \quad \Lambda = \text{diag}(\lambda_1, \dots, \lambda_d)$$

$$\lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_d \geq 0.$$

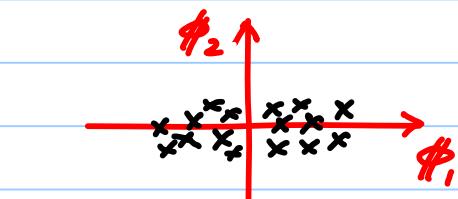
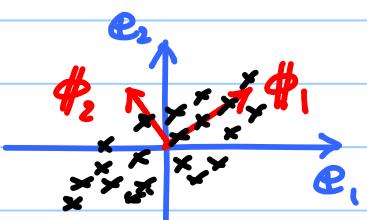
$\Phi := [\phi_1, \dots, \phi_d] \in \mathbb{R}^{d \times d}$ is

an ortho. matrix, and $\{\phi_1, \dots, \phi_d\}$ form an ONB of \mathbb{R}^d .

$\phi_j^T \tilde{X}$ is said to be the j th principal components of \tilde{X} .

These are nothing but the expansion coefficients of \tilde{X} w.r.t. the ONB vector ϕ_j .

If \tilde{X} forms a "cigar" shape, then $\phi_j^T \tilde{X}$ are the coordinate values of \tilde{X} under the rotated axes



- Hence viewing the given dataset under the principal axes Φ_1, Φ_2, \dots , provides us better interpretations of the data than viewing them under the original axes $\epsilon_1, \epsilon_2, \dots$.
- PCA is also often used as a tool to do **dimension reduction** and **feature extraction** by keeping only **top k** PCA coordinates where $k \ll d$, i.e.,

$$\Phi_k := [\Phi_1, \dots, \Phi_k] \in \mathbb{R}^{d \times k}$$

$$\mathbb{R}^d \ni \tilde{x}_j \mapsto \underbrace{\Phi_k^\top}_{\text{top } k \text{ PCA coordinates}} \tilde{x}_j \in \mathbb{R}^k$$

or top k Principal components of \tilde{x}_j .

Note that using these **top k** principal components, we can approximate the original data x_j by

$$x_j \approx \bar{x} + \Phi_k \Phi_k^\top \tilde{x}_j$$

Of course the approximation gets better and better as k increases. In fact, if $k=d$, then \tilde{x}_j is recovered exactly (within machine ϵ).

Now we'll face the problem when we compute the eigendecomposition of $S = \Phi \Lambda \Phi^T$:

- (1) If d is large, we cannot compute this eigendecomposition because we cannot hold $\Phi \in \mathbb{R}^{d \times d}$ in computer memory, and its computational cost is $O(d^3)$, i.e., too expensive to compute.
- (2) Fortunately, we often do not need all d eigenvectors, most likely, only first k eigenvectors $k \ll d$.
- (3) Moreover if $d > n$, then $\text{rank}(S) = n-1$ if $\tilde{\mathbf{x}}_j$'s are linearly indep. So, after the first $n-1$ eigenvectors are useless!

Why? $S = \frac{1}{n} \tilde{\mathbf{X}} \tilde{\mathbf{X}}^T = \frac{1}{n} \left\{ \underbrace{\tilde{\mathbf{x}}_1 \tilde{\mathbf{x}}_1^T + \cdots + \tilde{\mathbf{x}}_n \tilde{\mathbf{x}}_n^T}_{\text{rank 1}} \right\}$

So looks like $\text{rank}(S) = n$.
 But since $\tilde{\mathbf{x}}_1 + \cdots + \tilde{\mathbf{x}}_n = \mathbf{0}$ because the mean $\bar{\mathbf{x}}$ is subtracted from each data vector $\tilde{\mathbf{x}}_j$ (i.e., $\tilde{\mathbf{x}}_j = \mathbf{x}_j - \bar{\mathbf{x}}$)
 Hence, S loses 1 rank.
 So, $\text{rank}(S) = n-1$.

Now, let's consider the **reduced SVD** of \tilde{X} :

$$\tilde{X} = \hat{U} \hat{\Sigma} \hat{V}^T$$

$$\begin{array}{c} \text{d} \geq n \\ \tilde{X} = \hat{U} \hat{\Sigma} \hat{V}^T \end{array} \quad \begin{array}{c} \text{d} < n \\ \tilde{X} = \hat{U} \hat{\Sigma} \hat{V}^T \end{array}$$

Just consider the "neo-classical" setting, i.e., $d \geq n$ (e.g., the face image database)

Then consider the sample covariance matrix S using the above SVD:

$$S = \frac{1}{n} \tilde{X} \tilde{X}^T = \frac{1}{n} \hat{U} \hat{\Sigma} \hat{V}^T \hat{V} \hat{\Sigma}^T \hat{U}^T$$

$$= \frac{1}{n} \hat{U} \hat{\Sigma} \hat{\Sigma}^T \hat{U}^T = \frac{1}{n} \hat{U} \hat{\Sigma}^2 \hat{U}^T$$

Now $\hat{\Sigma} = \text{diag}(\sigma_1, \dots, \sigma_{n-1}, 0)$

if $\tilde{x}_1, \dots, \tilde{x}_n$ are linearly indep.

So, $\hat{\Sigma}^2 = \text{diag}(\sigma_1^2, \dots, \sigma_{n-1}^2, 0)$.

Finally, S can be written as

$$S = \hat{U} \left(\frac{1}{n} \hat{\Sigma}^2 \right) \hat{U}^T$$

$\uparrow = \text{diag}(\sigma_1^2/n, \dots, \sigma_{n-1}^2/n, 0)$

columns are orthonormal.

Comparing this with the eigende decomposition

$S = \Phi \Lambda \Phi^T$, we can conclude that

$$\begin{cases} \Phi(:, 1:n) = \hat{U} \\ \Lambda(1:n, 1:n) = \frac{1}{n} \hat{\Sigma}^2 = \text{diag}(\sigma_1^2, \dots, \sigma_{n-1}^2, 0) \end{cases}$$

In fact, only the $1:n-1$ portion is useful since $\sigma_n = 0$.

Hence, we should use the reduced SVD of \tilde{X} (not S) for computing PCA!!
Do not use the eigendecomposition of S unless d is small.

Note: $\tilde{X} V = \hat{U} \hat{\Sigma} V^T V = \hat{U} \hat{\Sigma}$
 $= [\sigma_1 u_1, \dots, \sigma_{n-1} u_{n-1}, 0]$
 $= [\tilde{X} v_1, \dots, \tilde{X} v_n]$
So, $u_j = \frac{1}{\sigma_j} \tilde{X} v_j$, $j=1, \dots, n-1$.

In other words, each principal axis u_j is just a linear combination of the (centered) input vectors $\tilde{x}_1, \dots, \tilde{x}_n$!

Now let's do MATLAB experiments using the face image database consisting of 143 faces each of which has $128 \times 128 = 16384$ pixels, i.e., $d=16384$, $n=143$.