











**ICA**

Given  $Z = (\zeta_1 \dots \zeta_n)$ ,  $k \leq \text{rank}(Z)$

Eigenvector decomposition of Empirical Correlation Matrix:

$$Z^T Z = U \Lambda U^T \quad U = (\zeta_1 \dots \zeta_k), \quad \Lambda^{-1} = U^T$$

$$\Lambda = \begin{pmatrix} \lambda_1 & & \\ & \ddots & \\ & & \lambda_k \end{pmatrix}, \quad \lambda_1 \geq \lambda_2 \dots \geq \lambda_k > 0$$

Best k-dimensional Subspace:  $V_k = \text{span}(\zeta_1, \dots, \zeta_k)$

$$F = (\zeta_1 \dots \zeta_k) \quad F^T F = I_k$$

Encoder/Encoded Test Sample:  
 $\tilde{x} \in \mathbb{R}^d \rightarrow y = F^T \tilde{x} \in \mathbb{R}^k$

Reconstruct/Decode:  $y \rightarrow \hat{x} = Fy = F\Lambda^{-1}x \in \mathbb{R}^d$   
 $= \text{Proj}_{V_k}(x)$

Approximation Error:  $\|x - \hat{x}\|^2 = \|x\|^2 - \|\Lambda^{-1}x\|^2$   
 $= \|x\|^2 - \|U\Lambda^{-1}U^T x\|^2$   
 $= \|x\|^2 - \|U\Lambda^{-1}\|^2 \|U^T x\|^2$

Total Approx Error:  $\sum_j \|x_j - \hat{x}_j\|^2 = \sum_j \lambda_j$

Dual/kernel PCA

Given  $Z, k$

Gram/Kernel Matrix:  $Z^T Z = V \Lambda V^T, V^T = V^T$

Eigen Decomp:  $\Lambda = (\lambda_1 \dots \lambda_n), \lambda_i \geq \dots \geq \lambda_k > 0$

Best k-dimensional Subspace:  $V_k = \text{span}(\zeta_1 \dots \zeta_k)$

$F = Z^T Z \Lambda^{-1/2}$  or  $(V \Lambda V^T)^{-1/2}$

$F^T F = I_k, V_k$  bests  $\text{span}(\zeta_1 \dots \zeta_k)$

Embed/Encode:  
 $x \in \mathbb{R}^d \rightarrow y = F^T x = \frac{1}{\sqrt{k}} \begin{pmatrix} -1 & \dots & -1 \\ \vdots & \ddots & \vdots \\ -1 & \dots & -1 \end{pmatrix} (Z^T x)$

Reconstruction: only if kernel feature mapping explicitly known  
 $\text{Proj}_{V_k}(x) = \hat{x} = Fy = F^T x$

Approx Error:  $\|x - \hat{x}\|^2 = \|x\|^2 - \|\Lambda^{-1}x\|^2$   
 $= \|x\|^2 - \|U\Lambda^{-1}U^T x\|^2$   
 $= \|x\|^2 - \|U\Lambda^{-1}\|^2 \|U^T x\|^2$

Total Approx:  $\sum_j \|x_j - \hat{x}_j\|^2 = \sum_j \lambda_j$

Encoding matrix Path Set:  $F^T Z = \Lambda^{-1/2} V_k^T$

**HW #9 Clustering**

1) Consider a 1-dimensional dataset of  $n=2m+1$  points ( $m \in \mathbb{N}$ ) spread on the real line with one point at  $0$ ,  $m$  points at  $a$ , and  $m$  points at  $-a$ , where  $0 < a \leq \frac{1}{2}$ . Now suppose we run k-means on the data points  $w_1, w_2, \dots, w_n$  and all centroid choices other than the random  $(0, a, -a)$  perform all distinct clusterings so that the k-means algorithm could converge to:  
 clustering 1:  $\{0, a, -a, \dots, a\}, \{c_1, c_2, \dots, c_{m-1}\}$   
 clustering 2:  $\{0\}, \{a, -a, \dots, a\}, \{c_1, c_2, \dots, c_{m-1}\}$

2) For each solution, provide expressions for centroids and WCSS in terms of  $m, a$ .

clustering 3:  $w_1 = 0, w_2 = \frac{a}{m+1} + a \frac{m}{m+1}$

$$\mu_1 = a$$

$$\text{WCSS}_1 = a((0-\mu_1)^2 + m(a-\mu_1)^2 + m(a+\mu_1)^2) = a + ma^2 + \frac{ma^2}{m+1} + \frac{ma^2}{m+1} = ma^2$$

clustering 2:  $w_1 = 0, w_2 = \frac{a}{2}$

$$\text{WCSS}_2 = a((0-\mu_1)^2 + m(a-\mu_1)^2 + m(a+\mu_1)^2) = a + ma^2 + ma^2 = 2a + 2ma^2$$

3) Compute the ratio of the largest and smallest WCSS as a function of  $m, a$  and comment on the implications.

Clustering 1)  $\text{WCSS}_1 \leq \text{WCSS}_2$

$$\text{WCSS}_1 = \frac{ma^2}{m+1} \leq \frac{ma^2}{2} \leq \frac{mb^2}{2} = \text{WCSS}_2$$

$$\text{Ratio } \frac{\text{WCSS}_1}{\text{WCSS}_2} = \frac{1}{m+1}$$

This ratio can become arbitrarily large for sufficiently large  $m$ .

So, if a dataset, for which k-means can converge to a solution whose WCSS is arbitrarily higher than the best solution.

Take away: necessary to run k-means w/ multiple initializations of centroids. Exchange solutions resulting in the smallest WCSS.

4) Suppose two initial centroids are chosen independently uniformly at random over  $[0, 1-a]$ . Compute the probability of converging to each possible solution. This is determined by whether the mid-point of the initial centroids lies to the left or right of  $a$ .

(Cluster 2) =  $P\left(\frac{\mu_1 + \mu_2}{2} < a\right)$

where  $\mu_1, \mu_2 \sim \text{IID Uniform}[0, 1-a]$

$$\frac{1}{2} \times 2a \times 2a \times \frac{1}{1-a} = \frac{2a^2}{1-a} < \frac{1}{2}$$

So,  $P(\text{Cluster 1}) = 1 - \frac{2a^2}{1-a}$

$\mu_2(0) \rightarrow \mu_1(1-a) < a$

$\text{Cov}(X, Y) = \sum_{i=1}^n (X_i - \bar{X})(Y_i - \bar{Y})$

**e.g. Find eigenvalues & eigenvectors of a  $2 \times 2$  matrix**

$A = \begin{bmatrix} 0 & 1 \\ -2 & 1 \end{bmatrix}$

$$|A - \lambda I| = \begin{vmatrix} 0-\lambda & 1 \\ -2 & 1 \end{vmatrix} = \begin{vmatrix} 0 & 1 \\ -2 & 1 \end{vmatrix} = 0$$

$$\begin{vmatrix} 0 & 1 \\ -2 & 1 \end{vmatrix} = 1^2 + 3\lambda + 2 = 0 \rightarrow \lambda_1 = -1, \lambda_2 = -2$$

$$A - \lambda_1 I = A + I = \begin{bmatrix} 1 & 1 \\ -1 & 0 \end{bmatrix}$$

$$(A - \lambda_1 I) \cdot v_1 = 0$$

$$\begin{bmatrix} 1 & 1 \\ -1 & 0 \end{bmatrix} \begin{bmatrix} 1 & 1 \\ -1 & 0 \end{bmatrix} = 0$$

$$\text{(stop now)}$$

$$\begin{bmatrix} 1 & 1 \\ -2 & 1 \end{bmatrix} \begin{bmatrix} v_{11} \\ v_{12} \end{bmatrix} = 0 \rightarrow \begin{bmatrix} v_{11} \\ v_{12} \end{bmatrix} = \begin{bmatrix} 1 \\ 1 \end{bmatrix}$$

(same procedure for second eigenvalue)

proportion of variance for 1st:  $\frac{1}{2}$

**PCA**

- 1) Subtract mean from each data dimension (get  $\tilde{X}$ ) → center data to get  $\tilde{X}$
- 2) Calculate covariance matrix
- 3) Calculate eigenvals/evecs
- 4) Eigenvect w/ highest eigenval is the PC of the data set
- 5) Final data =  $(eig_1 \ eig_2 \dots)^T (x_1 \ x_2 \dots)$
- 6) Back to orig:  $x_i = (eig_1 \ eig_2 \dots)(\text{Final data})$

def: # of features  
 $n$ : # of samples  
 $X, X_i \rightarrow \tilde{X} \rightarrow \text{new transformed matrix}$   
 $P$ : transformation mat  
 $V = (eig_1 \ eig_2)$   
 $PX = \tilde{X} \rightarrow \text{goal}$

$S_k = \frac{1}{n-1} \sum_{i=1}^n \tilde{x}_i \tilde{x}_i^T = \text{covmat}$   
 choose  $P = VT$

Final Data: data rows in columns  
 dims on rows

USE UNIT EIGENVECS

Orig data =  $(eig_1 \ eig_2 \dots) \times (\text{Final data}) + \text{orig means}$

1	2	3	4	5	6	
x	10	11	8	3	2	1
y	6	4	5	3	2	1

avg. measure for grade 1 =  $\frac{35}{6}$   
 grade 2 =  $\frac{10}{6}$   
 "fit" = project data onto it, minimize distances

Analy:

$n_{ij} = \# \text{ of points in cluster } i \text{ belonging to class } j$

$n_j = \sum_{i=1}^k n_{ij} = \text{total # of points in cluster } i$

purity of a cluster =  $p_i = \frac{\max_j n_{ij}}{n_i}$

purity of clustering:  $P = \frac{1}{k} \sum_{i=1}^k p_i$

$= \frac{1}{6} \sum_{i=1}^6 \max_{j=1}^2 n_{ij}$