# Removing Redundancies from Classification Dataset: Fisher Discriminant Analysis



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**Ronald Fisher** 

# Objectives

- •Understand how to preserve and highlight class information when reducing dimensionality of dataset.
  - Good embedding strategies for classification tasks
- Know how to perform Fisher Discriminant Analysis (FDA)
  - Difference between FDA and PCA

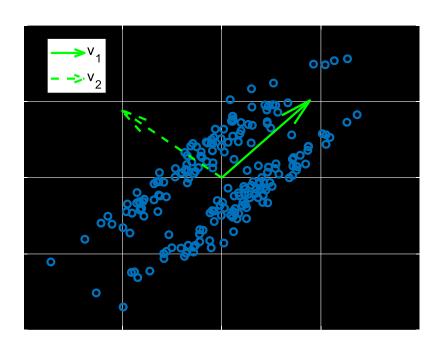
# Principle Component Analysis

- •PCA embed data points onto a lower dimensional surface, where they spread out the most.
  - •By a trace maximization problem.

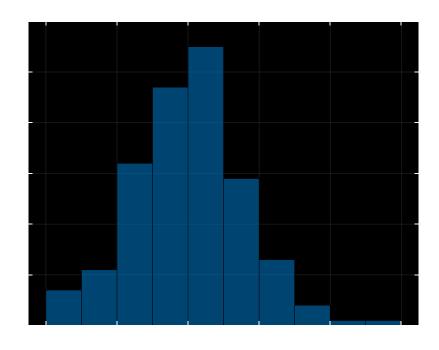
•PCA is performed by looking at eigenvectors corresp. to largest eigenvalues.

#### Problem of PCA

PCA ignores class/cluster information in the dataset!



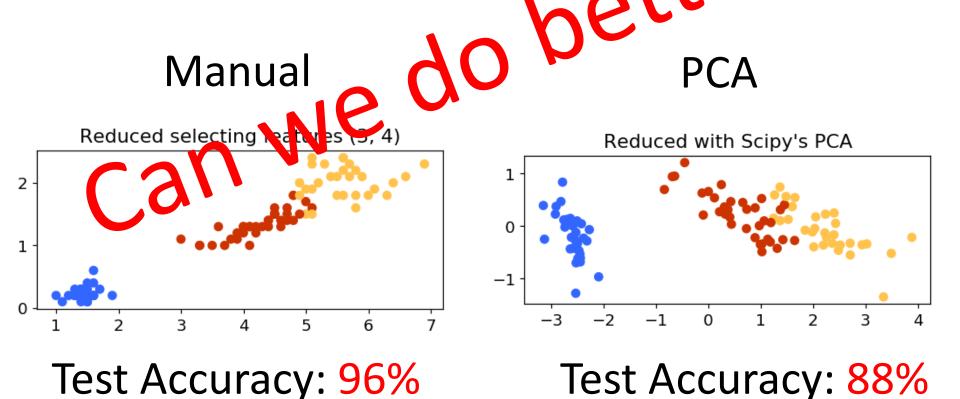
Eigenvecs



**Embedding** 

#### Problem of PCA

•Although, by maximizing the spread, PCA still does an respectable job.



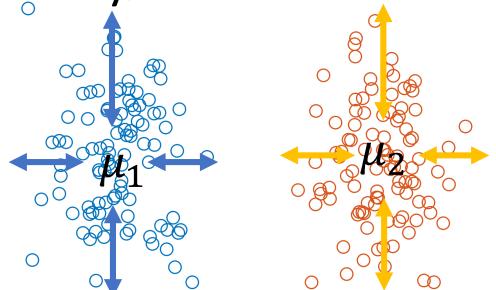
# **Problem Setting**

- Consider a classification dataset:
- • $D = \{(y_i, x_i)\}_{i=1}^n, x \in \mathbb{R}^d, y \in \{1 \dots k\}.$

- •Find feature transform function  $f(x) \in \mathbb{R}^m$  to reduce dimensionality of dataset.
  - •while preserving distinct class separation.

# What is a Good Embedding for a Classification Dataset?

- Points within the same class are close to each other.
  - •Within classes **scatterness** can be measured by distances to class center.



# What is a Good Embedding for a Classification Dataset?

- •Points **between** different classes are far apart from each other.
  - •Between classes scatterness can be measured by distances between class centers and dataset centers.

#### Within-class Scatterness

- •Embedding is  $Bx^{\top}$ .
- •Embedded center for class k:

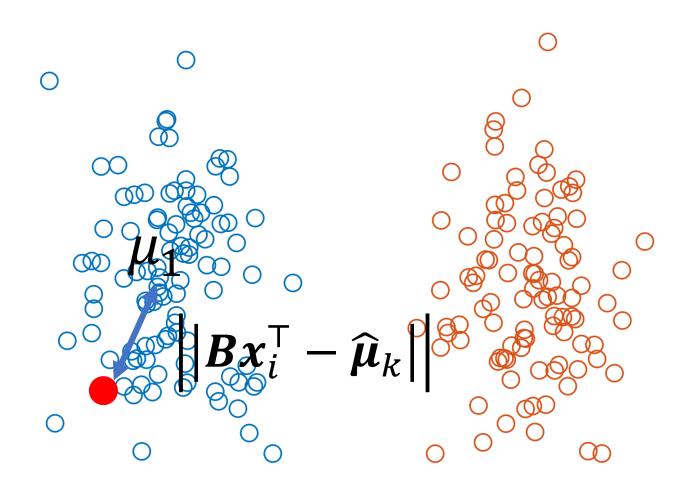
$$\bullet \widehat{\boldsymbol{\mu}}_k = \frac{1}{n_k} \sum_{i, y_i = k} \boldsymbol{B} \boldsymbol{x}_i^{\top}$$

•Within class scatterness of class k:

$$s_{w,k} = \sum_{i,y_i=k} \left| \left| B x_i^{\mathsf{T}} - \widehat{\mu}_k \right| \right|^2$$

•Sum over points in individual classes.

### Within-class Scatterness



#### **Between-class Scatterness**

•Embedded dataset centroid:

$$\bullet \widehat{\boldsymbol{\mu}} = \frac{1}{n} \sum_{i=1}^{n} \boldsymbol{B} \boldsymbol{x}_{i}^{\mathsf{T}}$$

Between-class scatterness

$$\bullet s_{b,k} = n_k ||\widehat{\boldsymbol{\mu}}_k - \boldsymbol{\mu}||_{\circ}^2$$

• $n_k$  is needed to make  $s_{b,k}$  at the same scale with  $s_{w,k}$ .

# Objective

- •Maximizing between class scatterness  $\forall_k$ .
  - •Minimize within class scatterness  $\forall_k$ .

$$\max_{\mathbf{B}} \sum_{k} s_{b,k} - \sum_{k} s_{w,k}$$

$$\bullet \sum_{k} s_{b,k} = \operatorname{tr} \{ \boldsymbol{B} \big[ \sum_{k} n_{k} (\widehat{\boldsymbol{\mu}}_{k} - \widehat{\boldsymbol{\mu}})^{\top} (\widehat{\boldsymbol{\mu}}_{k} - \widehat{\boldsymbol{\mu}}) \big] \boldsymbol{B}^{\top} \}$$

$$\bullet \sum_{k} s_{w,k} = \operatorname{tr} \{ \boldsymbol{B} \big[ \sum_{k} \sum_{i} (\boldsymbol{x}_{i} - \widehat{\boldsymbol{\mu}}_{k})^{\top} (\boldsymbol{x}_{i} - \widehat{\boldsymbol{\mu}}_{k}) \big] \boldsymbol{B}^{\top} \}$$

Live demonstration

### Objective

•Let 
$$S_w \coloneqq \sum_k \sum_i (x_i - \widehat{\mu}_k)^{\top} (x_i - \widehat{\mu}_k)$$

•Let 
$$S_b \coloneqq \sum_k n_k (\widehat{\boldsymbol{\mu}}_k - \widehat{\boldsymbol{\mu}})^{\top} (\widehat{\boldsymbol{\mu}}_k - \widehat{\boldsymbol{\mu}})$$

• 
$$\max_{\boldsymbol{B}} \sum_{k} s_{b,k} - \sum_{k} s_{w,k}$$
  
=  $\max_{\boldsymbol{B}} \operatorname{tr}[\boldsymbol{B}\boldsymbol{S}_{b}\boldsymbol{B}^{\top}] - \operatorname{tr}[\boldsymbol{B}\boldsymbol{S}_{w}\boldsymbol{B}^{\top}]$ 

### Objective

- However, the above problem is very hard to solve!
  - •Like PCA, we make the problem easier by introducing a constraint on  $\boldsymbol{B}$ .

Constrained Objective:

$$\max_{\boldsymbol{B}, \boldsymbol{B} \boldsymbol{S}_{\boldsymbol{W}} \boldsymbol{B}^{\top} = \boldsymbol{I}} \operatorname{tr} \left[ \boldsymbol{B} \boldsymbol{S}_{\boldsymbol{b}} \boldsymbol{B}^{\top} \right] - \operatorname{tr} \left[ \boldsymbol{B} \boldsymbol{S}_{\boldsymbol{W}} \boldsymbol{B}^{\top} \right]$$

$$\max_{\boldsymbol{B},\boldsymbol{B}\boldsymbol{S}_{\boldsymbol{W}}\boldsymbol{B}^{\top}=\boldsymbol{I}}\operatorname{tr}[\boldsymbol{B}\boldsymbol{S}_{\boldsymbol{b}}\boldsymbol{B}^{\top}]$$

#### Solution

- Eigenvalue/eigenvectors of A
  - $\bullet A \boldsymbol{v}_i = \lambda_i \boldsymbol{v}_i$
- •Generalized eigenvalue/eigenvectors of  $m{A}$  and  $m{B}$ 
  - $\bullet A \boldsymbol{v}_i = \lambda_i \boldsymbol{B} \boldsymbol{v}_i$
  - •MATLAB: [V,LABMDA] = eig(A,B)
  - Python: scipy.linalg.eig(A,B)

#### Solution

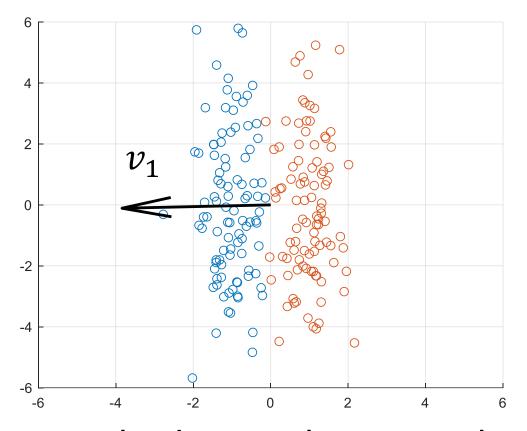
- $\max_{B,BS_wB^\top=I} \operatorname{tr}[BS_bB^\top]$
- •The embedding matrix  $\widehat{m{B}}$  can be constructed by
- $m{\cdot}\widehat{\pmb{B}} = [m{v}_1, m{v}_2, ... m{v}_m]^{ op}$ 
  - • $(\lambda_1, v_1), \dots, (\lambda_m, v_m)$  are m largest generalized eigenval. and eigenvec. of
  - $\bullet S_b v_i = \lambda_i S_w v_i$

#### Solution

- •Unfortunately, m < c 1.
  - •For a binary classification dataset, the embedding has to be 1D.
  - •rank( $S_b$ ) = c-1

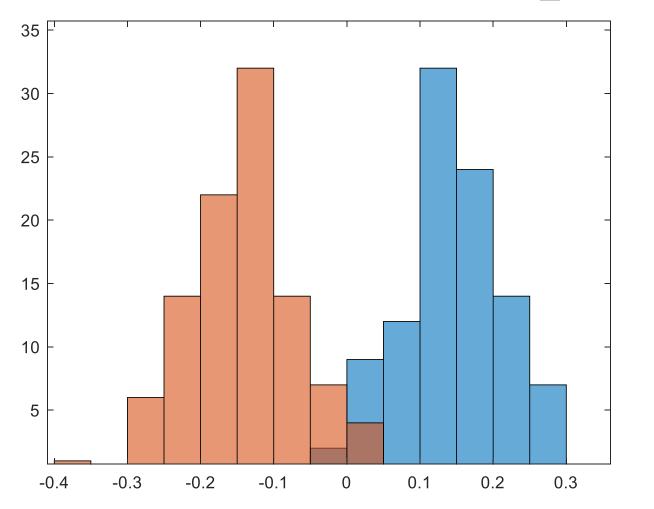
•The process of computing embedding using eigenvec. of  $S_b$  and  $S_w$  is called **Fisher Discriminant** Analysis (FDA).

# Example: Binary Classification Dataset



FDA embeds samples to a subspace that is the most **linearly** separable.

# Example: embedding, $\boldsymbol{v}_1^{\mathsf{T}} \boldsymbol{x}^{\mathsf{T}}$

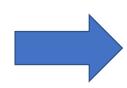


Class separation is preserved after embedding.

# Eigenfaces

$$X = \{x_i'\}, x_i' \in R^{d' \times d'}$$





$$x_i \in R^{d'd'}$$

#### Perform PCA -> Eigenvectors

$$\bullet [\boldsymbol{v}_1 \dots \boldsymbol{v}_m], \boldsymbol{v}_i \in R^{d'd' \times 1}$$

# Eigenfaces

$$\mathbf{v}_i \in R^{d'd'}$$

$$\mathbf{v}_i' \in R^{d' \times d'}$$

•Eigenfaces have huge applications in facial recognition.

#### Conclusion

- Good embedding of a classification dataset should have:
  - Small within class scatter
  - Large between class scatter

- •FDA maximizes between class scatter and minimizes within class scatter
  - Preserves class separation on datasets.