CS559 Machine Learning Fisher Linear Discriminant Principal Component Analysis

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Week 6

Outline

- Fisher's Linear Discriminant
- Principal Component Analysis
 - High-dimensional data, Eigenfaces
 - PCA vs FLD

Binary Classification

In previous lecture, we study **linear discriminant function** and use it to solve the binary classification problem. Specifically, we have:

$$y = \begin{cases} +1, \text{ if } \mathbf{w}^T \mathbf{x} > \theta \\ -1, \text{ if } \mathbf{w}^T \mathbf{x} < \theta \end{cases}$$

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We could view classification in another way.....

Classification through projection

• A linear function: $f(\mathbf{x}) = \mathbf{w}^T \mathbf{x} + w_0$ assuming in 2D, projects each point $\mathbf{x} = [x_1, x_2]^T$ to a line parallel to \mathbf{w} :

$$\begin{array}{c|c} \text{point in } \mathcal{R}^D & \text{projected point in } \mathcal{R} \\ \mathbf{x}_1 & z_1 = \mathbf{w}^T \mathbf{x}_1 \\ \mathbf{x}_2 & z_2 = \mathbf{w}^T \mathbf{x}_2 \\ \dots & \dots \\ \mathbf{x}_n & z_n = \mathbf{w}^T \mathbf{x}_n \end{array}$$

• We can study how well the projected points $z_1, ..., z_n$, viewed as functions of \mathbf{w} , are separated across the classes.

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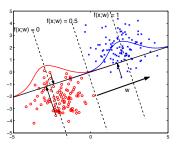


Figure: [N.Yue, CS559 S19]

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Classification through projection

 By varying w we get different levels of separation between the projected points

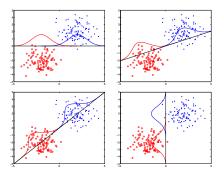


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Find the good projection

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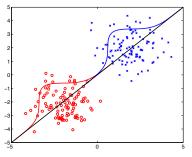


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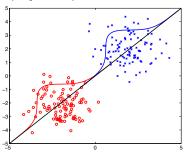


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 We can quantify the separation (overlap) in terms of means and variances of the resulting 1-dimensional class distributions

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• Decision stage: select a proper threshold y_0 , then

if
$$y \ge y_0$$
 assign \mathbf{x} to C_1 otherwise assign \mathbf{x} to C_2

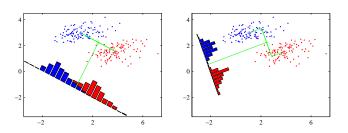


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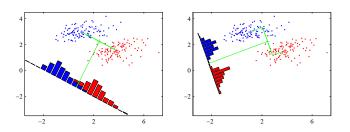


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- Find an direction along which the projected samples are well separated;
- We are looking for the linear projection that best separates the data, i.e. best discriminates data of different classes.

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- Assume $||\mathbf{w}|| = 1$, then $\mathbf{w}^T \mathbf{x}$ is the projection of \mathbf{x} onto \mathbf{w} .

Two goals

After projection, in order to obtain the best separation of the data, we need to satisfy two goals:

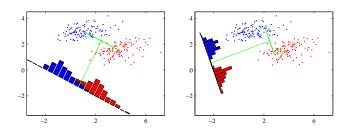


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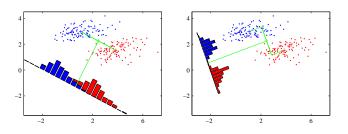


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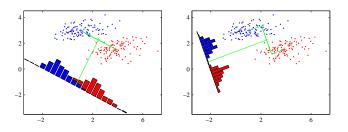


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 - $(\mathbf{w}^T \mu_1 \mathbf{w}^T \mu_2)^2 = \mathbf{w}^T (\mu_1 \mu_2) (\mu_1 \mu_2)^T \mathbf{w} = \mathbf{w}^T S_B \mathbf{w}$ (S_B : between class covariance matrix)

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 - More compactly, we have:

$$\max_{\mathbf{w}} \frac{\mathbf{w}^T S_B \mathbf{w}}{\mathbf{w}^T S_W \mathbf{w}}$$

Transform to constrained optimization problem:

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Take $\frac{\partial L(\mathbf{w},\lambda)}{\partial \mathbf{w}} = 0$, we have the generalized eigenvalue problem:

$$S_B \mathbf{w} = \lambda S_W \mathbf{w}$$

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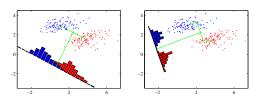


Figure: [C.Bishop PRML]

FLD procedure for two class classification:

Input: N_1 points from C_1 and N_2 points from C_2 .

- 1. Compute sample means for each class, i.e., μ_1 , μ_2 , and compute the S_W .
- 2. Compute $\mathbf{w} \leftarrow S_W^{-1}(\mu_1 \mu_2)$
- 3. Normalize: $\mathbf{w} \leftarrow \frac{\mathbf{w}}{||\mathbf{w}||}$
- 4. Select a suitable threshold θ , then classify \mathbf{x} to be C_1 if $\mathbf{w}^T \mathbf{x} > \theta$, otherwise, classify it to be C_2 .

Note:

• In the literature, people always use within class scatter matrix for S_W , which is:

$$S_W = \sum_{\mathbf{x} \in C_1} (\mathbf{x} - \mu_1)(\mathbf{x} - \mu_1)^T + \sum_{\mathbf{x} \in C_2} (\mathbf{x} - \mu_2)(\mathbf{x} - \mu_2)^T$$

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- What about other dimensionality reduction techniques?
 What if we do not have class label information?

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- In many applications, the observed data has very high dimensionality, e.g., images, videos, DNA sequences...
- **Assumption:** the data points lie close to a subspace of much *lower dimensionality* than that of the original data space.

Low-dimensional subspace

• E.g., the points in 3D space may form a line or plane.

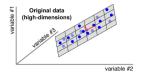


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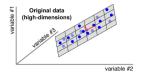


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• E.g., a small 100×100 gray-scaled image has 10,000 dimensions! While the intrinsic dimensionality might be low.



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- Useful tools in many applications such as face recognition, data compression, feature extraction etc.

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- Normally, not all D PCs are used but rather a subset of K "most important" PCs, $\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_K$.
- Key assumption: the direction within the data that shows the most variance contains the most information and therefore, is likely the most important.

First principal component

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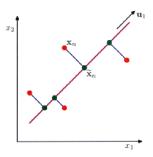


Figure: [C.Bishop, PRML]

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• The variance of the projected data:

$$\frac{1}{N} \sum_{n=1}^{N} [\mathbf{u}_{1}^{T} \mathbf{x}_{n} - \mathbf{u}_{1}^{T} \bar{\mathbf{x}}]^{2} = \mathbf{u}_{1}^{T} S \mathbf{u}_{1}$$

$$S = \frac{1}{N} \sum_{n=1}^{N} (\mathbf{x}_{n} - \bar{\mathbf{x}}) (\mathbf{x}_{n} - \bar{\mathbf{x}})^{T}$$

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• Using Lagrange Multiplier:

$$L(\mathbf{u}_1, \lambda) = \mathbf{u}_1^T S \mathbf{u}_1 - \lambda_1 (\mathbf{u}_1^T \mathbf{u}_1 - 1)$$

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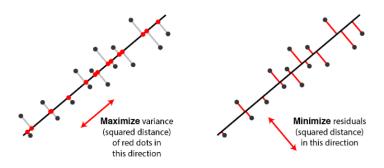


Figure: [Source]

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 - \mathbf{u}_3 is the eigenvector of S having the third largest eigenvalue λ_3 .
- In general, we get D eigenvalues of covariance matrix S, ordered from largest to smallest:

$$\lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_D$$

Then the corresponding eigenvectors $\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_D$ are **principal components**.

Dimension reduction

The principal components (PCs) $\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_D$ form the new coordinate system, so the original data \mathbf{x}_n can be projected onto new system:

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- If we keep all D PCs, then there is no dimensionality reduction but simply a rotation of the coordinate axes to align with the principal components.
- If keep K(< D) PCs, then we have *lower dimensional* representation, and we could approximate \mathbf{x}_n by:

$$\mathbf{x}_n pprox \sum_{i=1}^K (\mathbf{x}_n^T \mathbf{u}_i) \mathbf{u}_i$$

PCA for High-dimensional Data

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- N points in a D-dimensional space defines a subspace whose dimensionality is at most N-1, therefore, apply PCA, we will find at least D-N+1 of the eigenvalues to be 0.
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• The eigenvectors of S, i.e., $\mathbf{u}_1, \mathbf{u}_2, \ldots$ forms the **eigenfaces**.

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• Let $\mathbf{v}_i = \mathbf{X}\mathbf{u}_i$:

$$\frac{1}{N} \underbrace{\mathbf{X} \mathbf{X}^T}_{N \times N} \mathbf{v}_i = \lambda_i \mathbf{v}_i$$

• After we obtained the eigenvectors $\mathbf{v}_1, \mathbf{v}_2, \ldots$, we can use $(\mathbf{X}^T \mathbf{v}_i)$ to get eigenvector of S with eigenvalue λ_i :

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- Rescale $\mathbf{u}_i \propto \mathbf{X}^T \mathbf{v}_i$ such that $||\mathbf{u}_i|| = 1$.
- Keep top-K eigenvectors $\mathbf{u}_1, \mathbf{u}_2, \ldots, \mathbf{u}_K$ to form K eigenfaces. Reshape the D-dimensional \mathbf{u}_i to [h,w] to visualize the eigenfaces.

General procedures of obtaining eigenfaces

- 1. Construct the data matrix **X**, each row contains one centred face image represented by long-vector.
- 2. Consider $\mathbf{X}\mathbf{X}^T$ ($N \times N$), and find its K(< N) eigenvectors $\mathbf{v}_1, \dots, \mathbf{v}_K$.
- 3. Get K eigenfaces by normalizing the corresponding $\mathbf{X}^T \mathbf{v}_1, \dots, \mathbf{X}^T \mathbf{v}_K$.

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- Approximate the centred test/new face image using eigenfaces: $(\mathbf{x} \bar{\mathbf{x}})WW^T$.
- Finally, add back the mean image to obtain the approximation/reconstruction for test/new image: $\bar{\mathbf{x}} + (\mathbf{x} \bar{\mathbf{x}})WW^T$

Eigenface representation



Figure: [Source]

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- Principal Component Analysis depends only on data x without label information which is unsupervised.

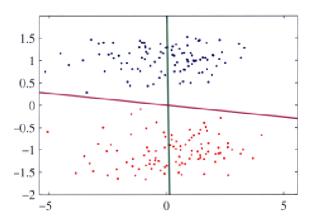


Figure: Red line: PCA. Green line: FLD [C.Bishop, PRML]

Acknowledgement and Further Reading

First few slides of Fisher Linear Discriminant are taken from Dr. Y. Ning's Spring 19 offering of CS-559.

Part of the discussion on different views of PCA is inspired by [bioramble]

Further Reading:

Chapter 4.1 and 12.1 of *Pattern Recognition and Machine Learning* by C. Bishop.