

Optimization and Data Analytics

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Given a set of N samples, each represented by a vector $\mathbf{x}_i \in \mathbb{R}^D$, and the corresponding labels l_i , we can define the class mean vectors:

$$\mu_k = \frac{1}{N_k} \sum_{i,l_i=k} \mathbf{x}_i, \ k = 1, \dots, K$$

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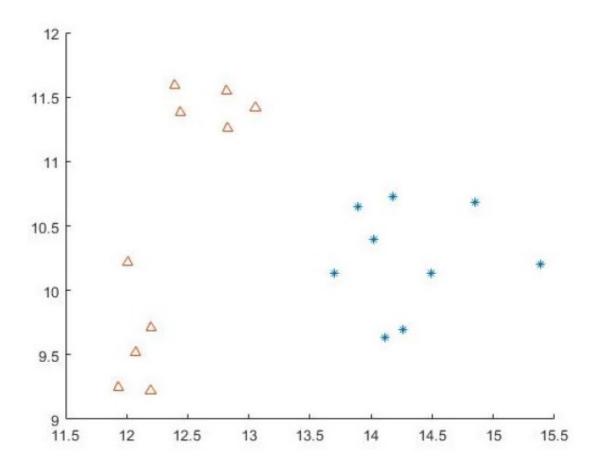
We use μ_k , k=1,...,K to represent the K classes.

Then, a new vector \mathbf{x}_* can be classified based on the minimal distance from $\mathbf{\mu}_k$

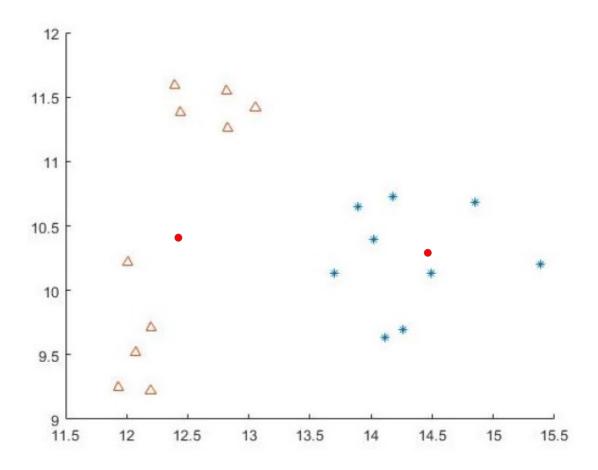
$$d(\mathbf{x}_*, \boldsymbol{\mu}_k) = \|\mathbf{x}_* - \boldsymbol{\mu}_k\|_2^2$$

This can be interpreted as a probalistic classification if we say that $p(x|c_k)$ is defined in such a way that the Sigma is identity and all prior probabilities $P(c_k)$ are the same. If $P(c_k)$ is larger than any of the other classes then naturally it will have more influence which moves the decision boundary!

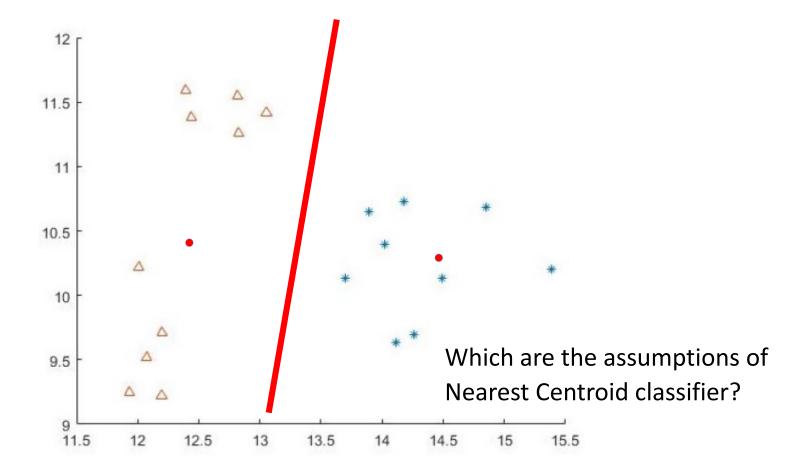














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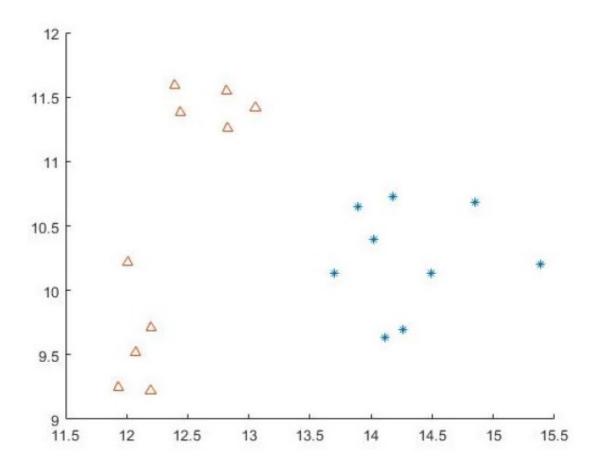
We obtain multiple prototypes for each class

$$\mu_{km} = \frac{1}{N_{km}} \sum_{i,l_i=k,q_i=m} \mathbf{x}_i$$

Then, a new vector \mathbf{x}_* can be classified based on the minimal distance from $\mathbf{\mu}_{\mathsf{km}}$

$$d(\mathbf{x}_*, \boldsymbol{\mu}_{km}) = \|\mathbf{x}_* - \boldsymbol{\mu}_{km}\|_2^2$$

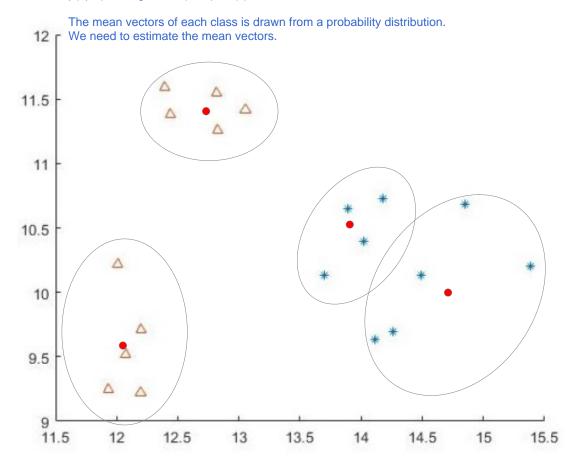






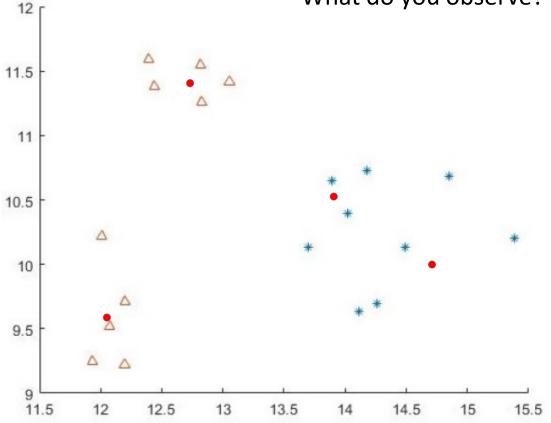
Example

Interpreting this as probability-based classifier is complicated because we do not know the mean vectors. The mean vectors are found by applying k-means. Applying k-means more than once, we get different resuts. p(x|c1) = integral of N(mu,) * P(x)

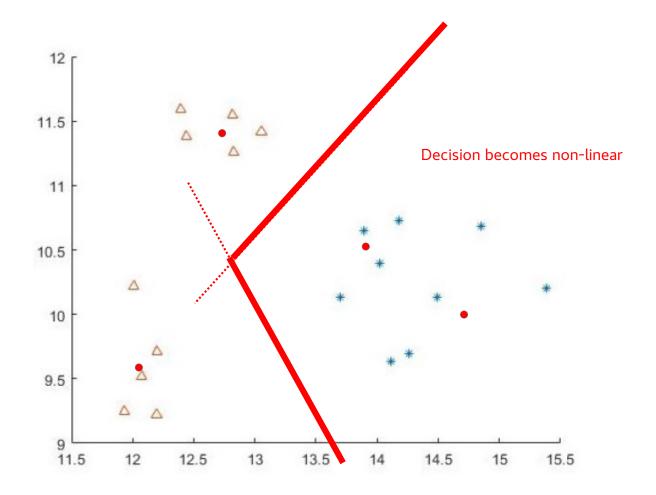




- Which will be the decision hyperplane?
- What do you observe?









Nearest Neighbor-based Classification

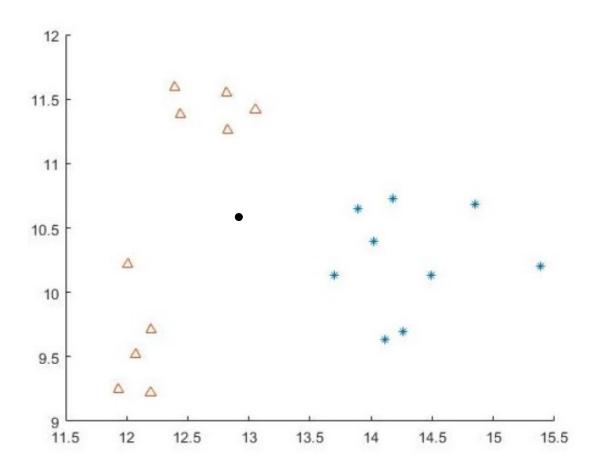
In the limit case where we assume that each training sample is a prototype, we end up calculating the distance of \mathbf{x}_* with all training vectors \mathbf{x}_i , i=1,...,N and classify it to the class of the closest training sample.

How can we use multiple nearest neighbors for classification?

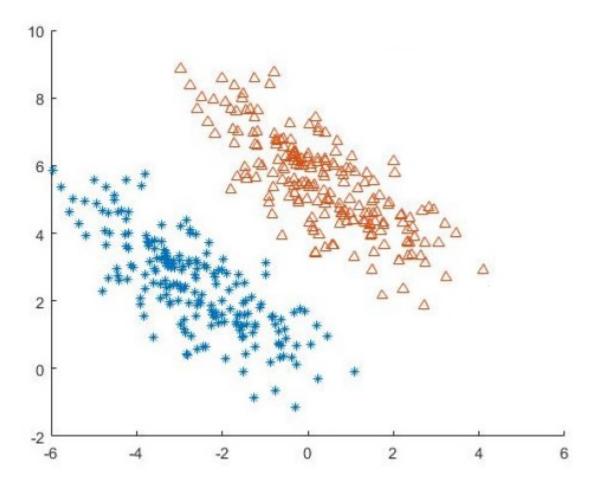
Demo: http://vision.stanford.edu/teaching/cs231n-demos/knn/



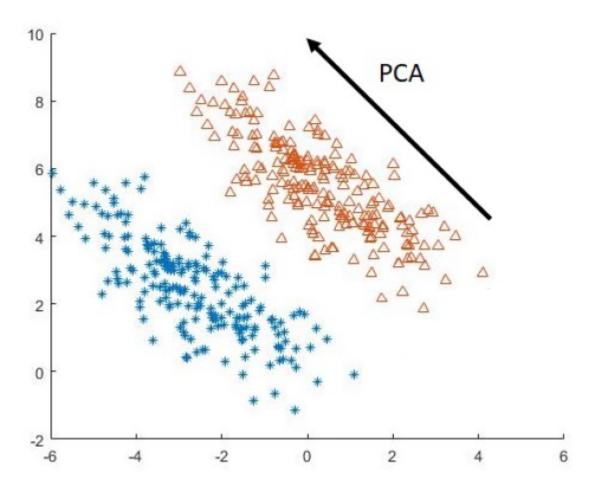
Nearest Neighbor-based Classification



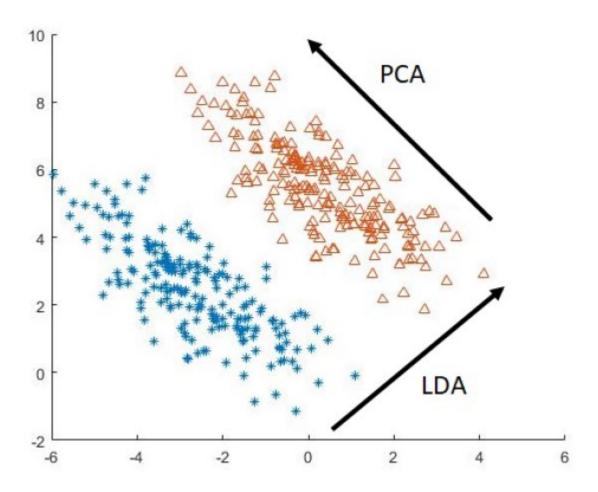














Given a set of N samples, each represented by a vector $\mathbf{x}_i \in \mathbb{R}^D$, and the corresponding labels $l_i = \{1,2\}$ we can define a linear projection of the form

$$y_i = \mathbf{w}^T \mathbf{x}_i$$

where $\mathbf{w} \in \mathbb{R}^D$ is a (projection) vector mapping the D-dimensional space to a line.

Demo: https://calerga.com/projects/fm20170202/lda.html



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Assuming that each class is unimodal and follows a Normal Distribution, how can we define the optimal vector \mathbf{w} ? The mean vector



We define the class mean vectors $\mathbf{\mu}_{k} \in \mathbb{R}^{D}, k=1,...,K$

$$\mu_k = \frac{1}{N_k} \sum_{i,l_i = k} \mathbf{x}_i$$

Then, the mean values of each class in the projection space (line) are

$$m_k = \frac{1}{N_k} \sum_{i,l_i=k} y_i = \frac{1}{N_k} \sum_{i,l_i=k} \mathbf{w}^T \mathbf{x}_i = \mathbf{w}^T \boldsymbol{\mu}_k$$



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The variance of each class in the line is

$$\sigma_k^2 = \frac{1}{N_k} \sum_{i,l_i = k} (y_i - m_k)^2$$



Since classes are unimodal and follow a Normal Distribution, they are better discriminated when:

- 1. The two mean values are as far as possible, i.e. their distance is as large as possible
- 2. The variances of the classes in the line are as small as possible



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The distance of the centers can be expressed as a function of **w**Similar to the expressing the objective function in PCA

$$(m_1 - m_2)^2 = (\mathbf{w}^T \boldsymbol{\mu}_1 - \mathbf{w}^T \boldsymbol{\mu}_2)^2$$
$$= \mathbf{w}^T (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2) (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)^T \mathbf{w}$$
$$= \mathbf{w}^T \mathbf{S}_b \mathbf{w}$$



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The variance can be written as $\sigma^2 = \sigma_1^2 + \sigma_2^2 = \sum_{k=1}^{\infty} \sum_{i,l,k} (\mathbf{w}^T \mathbf{x}_i - \mathbf{w}^T \boldsymbol{\mu}_k)^2$

$$= \sum_{k=1}^{2} \sum_{i,l,i=k} \mathbf{w}^{T} (\mathbf{x}_{i} - \boldsymbol{\mu}_{k}) (\mathbf{x}_{i} - \boldsymbol{\mu}_{k})^{T} \mathbf{w} = \mathbf{w}^{T} \mathbf{S}_{w} \mathbf{w}$$



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After expressing the two objectives above as functions of w, we can formulate an optimization problem which is a function of w

$$\mathcal{J}(\mathbf{w}) = rac{\mathbf{w}^T \mathbf{S}_b \mathbf{w}}{\mathbf{w}^T \mathbf{S}_w \mathbf{w}}$$
 maximise the numerator while minimising the denominator

maximise the numerator



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The above optimization problem is equivalent to the following problem

We can arrive at the next expressiong if we set the derivative of J(w) = 0. Once we take the derivative then the denominator cannot be zero. Look at the camera picture. $S_{\bf k} {\bf w} = \lambda S_{\bf w} {\bf w}$



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The above optimization problem is equivalent to the following problem

$$\mathbf{S}_b \mathbf{w} = \lambda \mathbf{S}_w \mathbf{w}$$

Notice that w has not the same scale here.

Therefore, w must be normalised

Assuming that S_w is non-singular

$$\mathbf{S}_w^{-1}\mathbf{S}_b\mathbf{w} = \lambda\mathbf{w}$$
 $\mathbf{w} = \mathbf{S}_w^{-1}(\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)$



Linear Discriminant Analysis (LDA) is the extension of Fisher Discriminant Analysis for the case where K > 2.



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In LDA, S_w is a straightforward extension of the one used in FDA

$$\mathbf{S}_k = \sum_{i,l_i=k} (\mathbf{x}_i - \boldsymbol{\mu}_k) (\mathbf{x}_i - \boldsymbol{\mu}_k)^T$$

$$\mathbf{S}_w = \sum_{k=1}^{\mathsf{K}} \sum_{i,l_i = k} (\mathbf{x}_i - \boldsymbol{\mu}_k) (\mathbf{x}_i - \boldsymbol{\mu}_k)^T$$



In order to define the between-class scatter, we have

Total scatter matrix? Scatter matrix of all samples!

$$\begin{split} \mathbf{S}_T &= \sum_{k=1}^K \sum_{i,l_i=k} (\mathbf{x}_i - \boldsymbol{\mu}) (\mathbf{x}_i - \boldsymbol{\mu})^T \\ &= \sum_{k=1}^K \sum_{i,l_i=k} (\mathbf{x}_i - \boldsymbol{\mu}_k + \boldsymbol{\mu}_k - \boldsymbol{\mu}) (\mathbf{x}_i - \boldsymbol{\mu}_k + \boldsymbol{\mu}_k - \boldsymbol{\mu})^T \\ &= \sum_{k=1}^K \sum_{i,l_i=k} (\mathbf{x}_i - \boldsymbol{\mu}_k) (\mathbf{x}_i - \boldsymbol{\mu}_k)^T + \sum_{k=1}^K \sum_{i,l_i=k} (\boldsymbol{\mu}_k - \boldsymbol{\mu}) (\boldsymbol{\mu}_k - \boldsymbol{\mu})^T \\ &= \mathbf{S}_w + \sum_{k=1}^K N_k (\boldsymbol{\mu}_k - \boldsymbol{\mu}) (\boldsymbol{\mu}_k - \boldsymbol{\mu})^T \quad \text{N_k is because we have double sum here Essentially,} \\ &= \mathbf{S}_w + \mathbf{S}_b. \end{split}$$



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Thus, the within-class and between-class scatter matrices are defined as

$$\mathbf{S}_k = \sum_{i,l_i=k} (\mathbf{x}_i - \boldsymbol{\mu}_k) (\mathbf{x}_i - \boldsymbol{\mu}_k)^T$$

$$\mathbf{S}_w = \sum_{k=1}^{\mathsf{K}} \sum_{i,l_i = k} (\mathbf{x}_i - \boldsymbol{\mu}_k) (\mathbf{x}_i - \boldsymbol{\mu}_k)^T$$

$$\mathbf{S}_b = \sum_{k=1}^K N_k (\boldsymbol{\mu}_k - \boldsymbol{\mu}) (\boldsymbol{\mu}_k - \boldsymbol{\mu})^T$$



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The optimization problem of LDA is

$$\mathcal{J}(\mathbf{W}) = \frac{Tr(\mathbf{W}^T \mathbf{S}_b \mathbf{W})}{Tr(\mathbf{W}^T \mathbf{S}_w \mathbf{W})}$$

What are the dimensions of **W**?



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Solving this eigenproblem we get K-1 eigenvectors because the rank of S_b is K-1

We usually add a constraint $W^TW = I$, why?

Because we need the vectors in W to be orthonormal. The practical problem that we are solving is we don't want to have irrelavent information.