

Prof. Marios Savvides

# Pattern Recognition Theory

**Lecture 6 : Principal Component Analysis (PCA) - I**

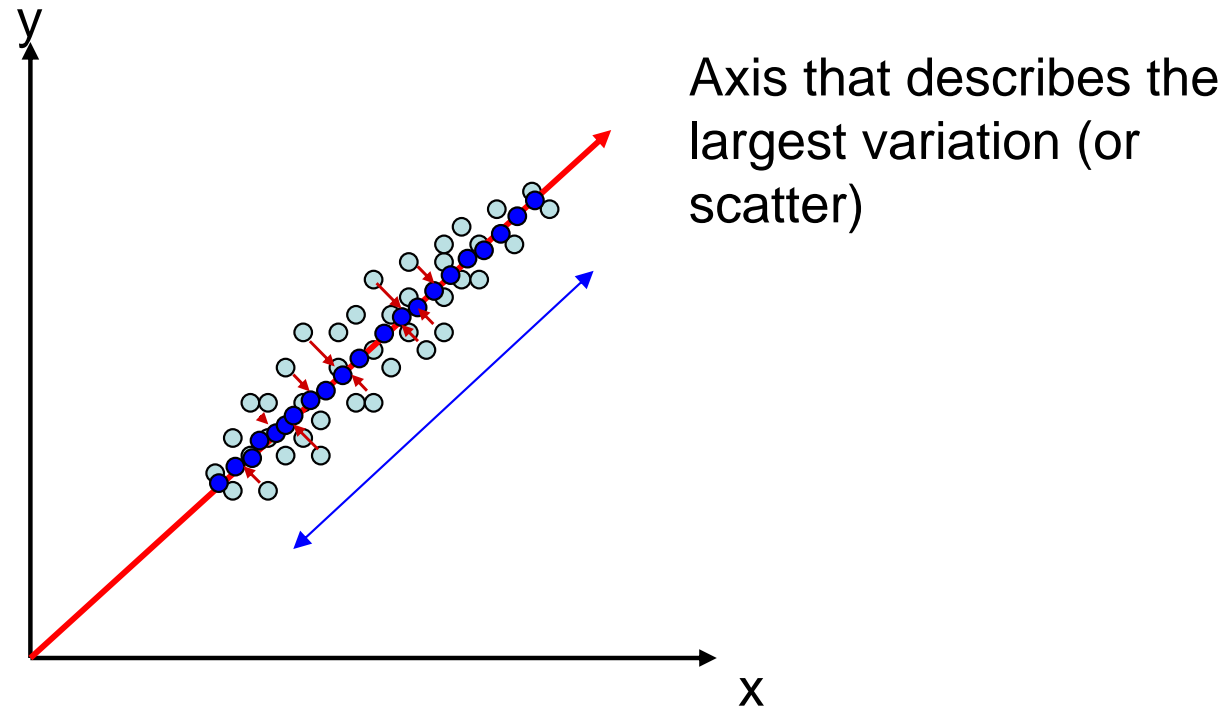
# PCA

- Also known as Hotelling Transform or Karhunen Loeve Transform
- Very popular pattern recognition tool...also known as “Eigenfaces” by Turk & Pentland
- Commonly used baseline benchmark when testing your new pattern recognition algorithm..i.e. given same training data and testing configuration, can I do better than PCA?
- So....you must DEFINITELY LEARN PCA!

# What is PCA?

## What are we trying to do?

- We want to find projections of data (i.e. direction vectors that we can project the data on to) that describe the maximum variation.



# PCA formulation

- We want projection vectors  $\omega$  that when we project the data onto these directions we maximize the scatter or variance.

- i.e. we want to maximize

$$\text{Var} (\omega^T \mathbf{x})$$

Subject to the constraint that  $\omega$  is a direction vector of unit norm i.e.  $\|\omega\|=1$

# How is PCA derived?

- Maximize the following objective function:

$$\begin{aligned} J(\omega) &= \text{Var}(\omega^T \mathbf{x}) \\ &= E(\omega^T \mathbf{x} - \omega^T \mu)^2 \\ &= E(\omega^T \mathbf{x} - \omega^T \mu)(\mathbf{x}^T \omega - \mu^T \omega) \\ &= \omega^T E(\mathbf{x} - \mu)(\mathbf{x} - \mu)^T \omega \\ &= \omega^T \Sigma \omega \end{aligned}$$

Where covariance matrix  $\Sigma = E(\mathbf{x} - \mu)(\mathbf{x} - \mu)^T$

# Derivation

- We want to maximize  $J(\omega)$  subject to the projection vectors  $\omega$  be unit norm i.e.
- Maximize quadratic term  $\omega^T \Sigma \omega$  subject to  $\|\omega\|=1$ .
- Solution: Form Lagrangian optimization to take care of constraints, take derivative and set to zero to find  $\omega$  vectors.

# Derivation

Solve:  $L(\omega, \lambda) = \omega^T \Sigma \omega - \lambda(\omega^T \omega - 1)$

Take derivative w.r.t  $\omega$ , set it to zero

$$\frac{\partial L(\omega, \lambda)}{\partial \omega} = 2\Sigma\omega - 2\lambda\omega = 0$$

$$\Sigma\omega = \lambda\omega$$

Standard Eigenvalue/Eigenvector problem ( $Ax = \lambda x$ ). i.e.

the vectors  $\omega$  are the eigenvectors of the covariance matrix  $\Sigma$ .

# PCA

- Ok..do eigenanalysis...find eigenvectors  $\omega_i$  and corresponding eigenvalues  $\lambda_i$ . Which to use though? All of them?
- A  $d \times d$  covariance matrix contains max  $d$  eigenvector/eigenvalue pairs. Do we need to compute all of them?
- Which  $\omega_i$  vector and eigenvalue  $\lambda_i$  pairs to use?



# PCA

- Lets re-visit our original objective of PCA....
- We want to find projections which describe the biggest variance (or have the maximum scatter of data).
- Remember the variance of projected data is

$$\omega^T \Sigma \omega. \quad (1)$$

- And our solution yielded  $\Sigma \omega = \lambda \omega \quad (2)$

- Plug (2) in (1) and we get

$$\begin{aligned} \text{projected variance} &= \omega^T \Sigma \omega = \omega^T \lambda \omega \\ &= \lambda \omega^T \omega \quad (\text{remember } \|\omega\|=1) \\ &= \lambda \end{aligned}$$

# PCA

- So this means that the direction vector  $\omega_i$  has captures  $\lambda_i$  variance.
- So we want to first represent the data using projections with largest variance, so order and keep the  $\omega_i$ 's with largest  $\lambda_{\max}$ .
- In fact the  $d \times d$  covariance matrix will not be full rank. Assume you  $N$  training images of size  $M \times M = d$ , then you have AT MOST  $N-1$  non-zero eigenvalues!

# Properties of PCA vectors

- Remember we computed eigenvectors  $\omega$  from covariance matrix  $\Sigma$ .
- IMPORTANT NOTE:  $\Sigma$  matrix is symmetric, i.e.  $\Sigma = \Sigma^T$  is how you test for symmetry.

Proof:

$$\begin{aligned}\Sigma &= E(x - \mu)(x - \mu)^T = [E(x - \mu)(x - \mu)^T]^T \\ &= E(x - \mu)(x - \mu)^T = \Sigma^T\end{aligned}$$

# So what?

- Symmetric Matrix, has ORTHOGONAL eigenvectors!
- What does this mean?

$$\begin{aligned}\omega_i^T \omega_k &= 0 \quad \text{for all } i \neq k \\ \omega_i^T \omega_k &= 1 \quad \text{for all } i = k \quad (\omega^T \omega = 1)\end{aligned}$$

(i.e. the inner-product or dot-product between different eigenvectors is 0.) They are un-correlated, i.e. information about how data varies in the direction of one of the eigenvectors tells you nothing about how data varies in the directions of the eigenvectors.

- This also means we have an ORTHOGONAL basis that we can use to represent our data!

# Expanding a signal using a basis

From this slide on: we refer to principal direction vectors  $\omega$  as eigenvectors  $\mathbf{v}$

- Assume you have a discrete vector signal  $\mathbf{x}$ .
- You have a set of  $N$  basis vectors  $\mathbf{v}_i$  which can use to represent a signal as follows:

$$\mathbf{x} = \sum_{i=1}^N p_i \mathbf{v}_i = p_1 \begin{bmatrix} | \\ \mathbf{v}_1 \\ | \end{bmatrix} + p_2 \begin{bmatrix} | \\ \mathbf{v}_2 \\ | \end{bmatrix} + \dots + p_n \begin{bmatrix} | \\ \mathbf{v}_n \\ | \end{bmatrix} = \mathbf{V} \mathbf{p}$$

i.e. the signal is a linear combination of the basis vectors where the  $p_i$  scalars are weight coefficients.

$$\mathbf{V} = \begin{bmatrix} | & | & | & | \\ \mathbf{v}_1 & \mathbf{v}_2 & \mathbf{v}_3 & \dots & \mathbf{v}_N \\ | & | & | & | \end{bmatrix}$$

$$\mathbf{p} = \begin{bmatrix} p_1 \\ p_2 \\ | \\ p_N \end{bmatrix}$$

# Computing the weight coefficients

$$\mathbf{x} = \sum_{i=1}^N p_i \mathbf{v}_i = p_1 \begin{bmatrix} | \\ \mathbf{v}_1 \\ | \end{bmatrix} + p_2 \begin{bmatrix} | \\ \mathbf{v}_2 \\ | \end{bmatrix} + \dots + p_n \begin{bmatrix} | \\ \mathbf{v}_n \\ | \end{bmatrix} = \mathbf{V} \mathbf{p}$$

- Since this is an orthogonal basis, we can easily find the coefficients  $p$ . These are just the projections of the signal on to each basis.

- i.e.  $p_1 = \mathbf{x}^T \mathbf{v}_1$  or  $\mathbf{p} = \mathbf{V}^T \mathbf{x}$

$$\mathbf{V}^T \mathbf{x} = \begin{bmatrix} - & \mathbf{v}_1 & - \\ - & \mathbf{v}_2 & - \\ - & \mathbf{v}_3 & - \end{bmatrix} \begin{bmatrix} | \\ \mathbf{x} \\ | \end{bmatrix} = \begin{bmatrix} p_1 \\ p_2 \\ p_3 \end{bmatrix}$$

# In PCA we model the variations about the mean.

- Don't forget that you model the variance about the mean!
- i.e. Don't forget to subtract the global mean before you analyze your data.

$$\mathbf{p} = \mathbf{V}^T (\mathbf{x} - \mathbf{m})$$

- By Mean we mean the sample mean image/vector of all your training data samples.
- Don't forget to add the mean back before you reconstruct the data!

$$\mathbf{x} = \sum_{i=1}^N p_i \mathbf{v}_i + \mathbf{m} = p_1 \begin{bmatrix} | \\ \mathbf{v}_1 \\ | \end{bmatrix} + p_2 \begin{bmatrix} | \\ \mathbf{v}_2 \\ | \end{bmatrix} + \dots + p_n \begin{bmatrix} | \\ \mathbf{v}_n \\ | \end{bmatrix} + \mathbf{m} = \mathbf{V} \mathbf{p} + \mathbf{m}$$

# PCA – Properties of

- These projection vectors form a basis for describing the training data in the Minimum Squared Error (MSE) sense.
- I.e. the eigenvectors with most largest eigenvectors capture most of the signal energy and reconstruction ability.
- $N$  samples of data will have a covariance matrix of rank at most  $N-1$ . That means you have at MOST  $N-1$  NON-ZERO eigenvalues. And you ONLY care about these corresponding  $N-1$  eigenvectors.



# How do I compute these in Matlab?

- Use MATLAB command “eigs” and “eig”
- “Eigs” is nice because it can find only a few eigenvectors,
- $[v,d]=\text{eigs}(\Sigma,N)$  which finds the  $N$  eigenvectors with the largest  $N$  eigenvalues.

# Practical Issues in computing PCA

- Assume we have image data of size 200x200 pixels.
- This means that our data is 40,000 dimensions!
- What is the size of Covariance Matrix?  
$$\Sigma = E(x - \mu)(x - \mu)^T \Rightarrow 40,000 \times 40,000$$
- This is a BIG MATRIX!..infact you CAN NOT create this matrix in MATLAB, no matter how much memory you have!

# Also...

- Well we only have  $N$  training samples, where  $N \ll d$  ( $d$ =dimensionality of our problem=40,000).
- Since  $N$  is much smaller than  $d$  and we will have AT MOST  $N-1$  eigenvectors (assuming  $N-1$  linearly independent data samples).
- It's not computationally feasible or possible to work with large covariance matrices.

# Instead...I show you the Gram Matrix Trick!

- We know that  $\Sigma = E(x - \mu)(x - \mu)^T = XX^T$

Must solve  $\Sigma v = \lambda v$

$$XX^T v = \lambda v \quad (\text{pre-mult by } X^T) \quad (1)$$

$$X^T X X^T v = \lambda X^T v \quad (v' = X^T v) \quad (2)$$

Solve eigenvalue problem  $X^T X v' = \lambda v'$

$X^T X$  is a gram or inner-product matrix, its size is not dependent on dimensionality of data but rather on the number of data samples  $N$ . It is of size  $N \times N$  and hence easier to compute if  $N \ll d$ .

# Ok..but how do I compute $v$ from $v'$ ?

- Remember Eq(1) and Eq (2):

$$XX^T v = \lambda v \quad (1)$$

$$v' = X^T v \quad (2)$$

Recognizing and subst (2) in (1) we get

$$Xv' = \lambda v$$

Thus  $v = Xv'$ . We don't care about scaling term because we will anyway make eigenvector orthonormal i.e.  $\|v\|=1$ .

# Lets see an example on real face data.

- We use CMU's AMP Lab facial Expression database.
- 13 people.
- Images are 64x64 cropped and centered facial images.
- Variations are due to varying expressions in the video sequence.
- 75 images in each person's video sequence

# AMP Lab Facial Expression DB



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# Experiment:

- Take the first 5 images per person and use them as training data
- Total of  $5 \times 13 = 65$  training images
- Do PCA and compute basis eigenvectors.
- Measure Reconstruction ability
- Perform Dimensionality Reduction (to 3d, i.e. only use a few eigenvectors and look at how data clusters in this 3D linear subspace).



# What do the eigenvectors look like?

Mean



V1



V2



V3



V4



V5



V6



V7



V8



V9



V10



V11



V12



V13



V14



V15



# How much reconstruction?

- If you want to say reconstruct an image with only 20% MSE (or 80% reconstruction)
- Then use the eigenvectors with largest M eigenvalues from the total of N non-zero eigenvalues satisfying this ratio:

$$reconstruction\% = \frac{\sum_{i=1}^M \lambda_i}{\sum_{i=1}^N \lambda_i} * 100$$

Sum of the eigenvalues of the kept eigenvectors

Sum of all eigenvalues

# All 64 eigenvectors..do we need all?



Noisy..not much info..will not contribute to image reconstruction

# See how it works...(use only the 3 most dominant eigenvectors)

Mean



+ 230

v1



- 917

v2



+ 1050

v3



MSE=758.13

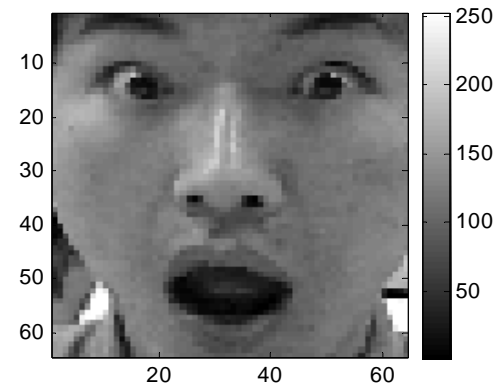
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reconstructed

$\tilde{\mathbf{X}}$

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Original

$\mathbf{X}$

# How do we compute projection coefficients p?

Mean (m)

v1

v2

v3



+ 230



- 917



+ 1050



x

m

c



-



=



## Step 1:

Subtract mean image

## Step 2:

Project onto eigenvector

$$\sum_{x=1}^X \sum_{y=1}^Y$$



\*

(element multip



= 230

# Projections in vector form

Step 2:  
Project onto  
eigenvector

$$\sum_{x=1}^X \sum_{y=1}^Y$$



$*$   
.  
(element multip



=230



$$\Rightarrow \begin{bmatrix} | \\ | \\ | \\ | \\ | \\ | \\ | \\ | \end{bmatrix} = \mathbf{x}$$



$$\Rightarrow \begin{bmatrix} | \\ | \\ | \\ | \\ | \\ | \\ | \\ | \end{bmatrix} = \mathbf{v}_1$$

$$\mathbf{p}_1 = \mathbf{x}^T \mathbf{v}_1 = 230$$

MSE=1233.16



- If we use only 1 eigenvector, MSE=1233  
(max Eigenvalue)

MSE=1027.63



- If we use 2 eigenvectors, MSE=1027



MSE=758.13



- If we use 3 eigenvectors, MSE=758

MSE=634.54



- If we use 4 eigenvectors, MSE=634

MSE=399.08



- If we use 7 eigenvectors, MSE=399

MSE=285.08



- If we use 8 eigenvectors, MSE=285

MSE=216.88



- If we use 13 eigenvectors, MSE=216

MSE=87.93



- If we use 20 eigenvectors, MSE=87

MSE=20.55



- If we use 30 eigenvectors, MSE=20

MSE=6.84



- If we use 45 eigenvectors, MSE=6.8



MSE=2.14



- If we use 50 eigenvectors, MSE=2.1

MSE=0.06



- If we use 60 eigenvectors,  $MSE=0.06$

MSE=0.00



- If we use 64 eigenvectors, MSE=0

(NOTE: We are using ALL NON-ZERO EigenVALUE Eigenvectors!)

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# Feature Extraction via Dimensionality Reduction

- Lets throw away many eigenvectors and only use a couple most dominant ones (this will perform dimensionality reduction)
- We project our data samples onto these vectors and store the projection coefficient for each projected data sample.

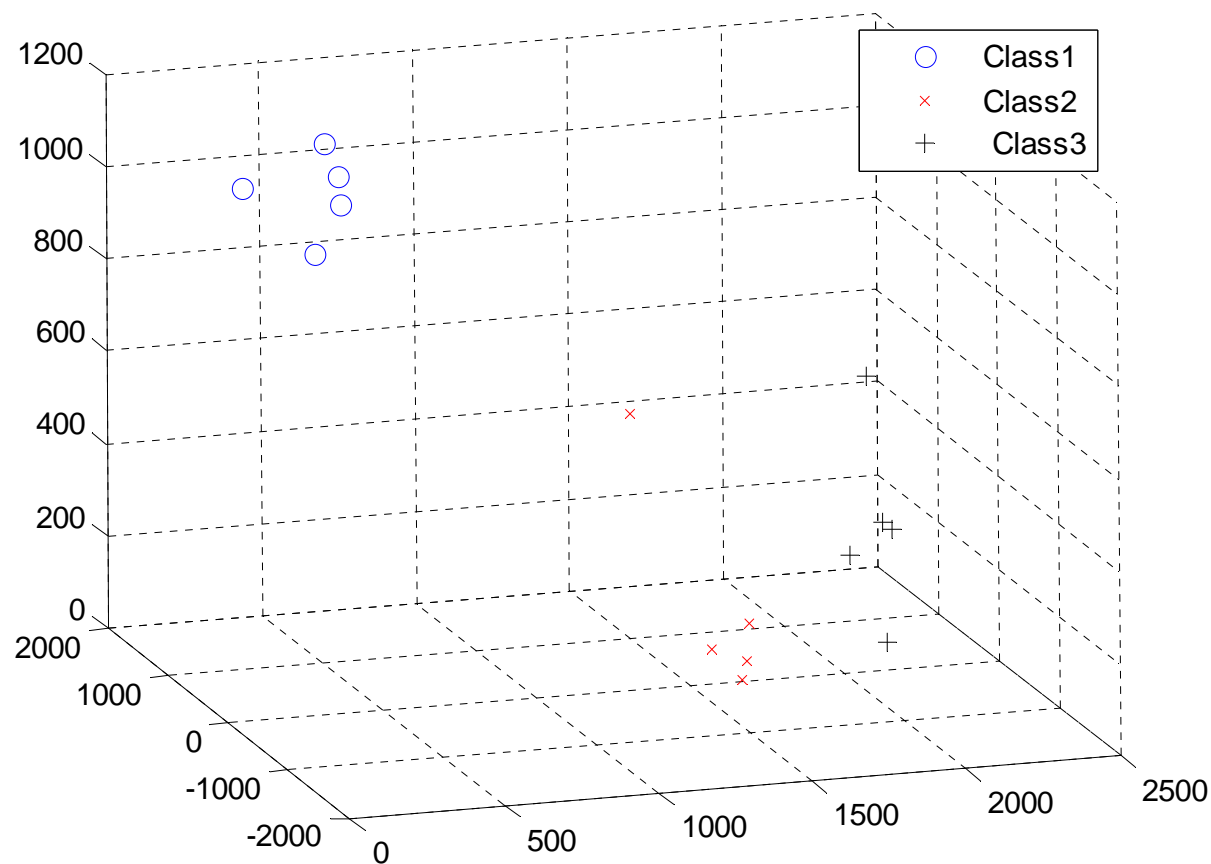
$$\mathbf{p} = \mathbf{V}^T(\mathbf{x} - \mathbf{m})$$

- So if we have N classes with k images per class then we will have kxN projection vectors  $\mathbf{p}$ . We only need to store these vectors and not the original training data to do classification.

# E.g.

- In our example we have 64x64 images, however in this toy example we performed PCA (global PCA since it uses all classes) and then only kept the 3 most dominant eigenvectors (i.e. with the 3 largest eigenvalues).
- Project each  $64 \times 64 = 4096$  dimensional training image onto this basis to get a 3-dimensional vector. That 3d vector represents each training sample in a 3d subspace.

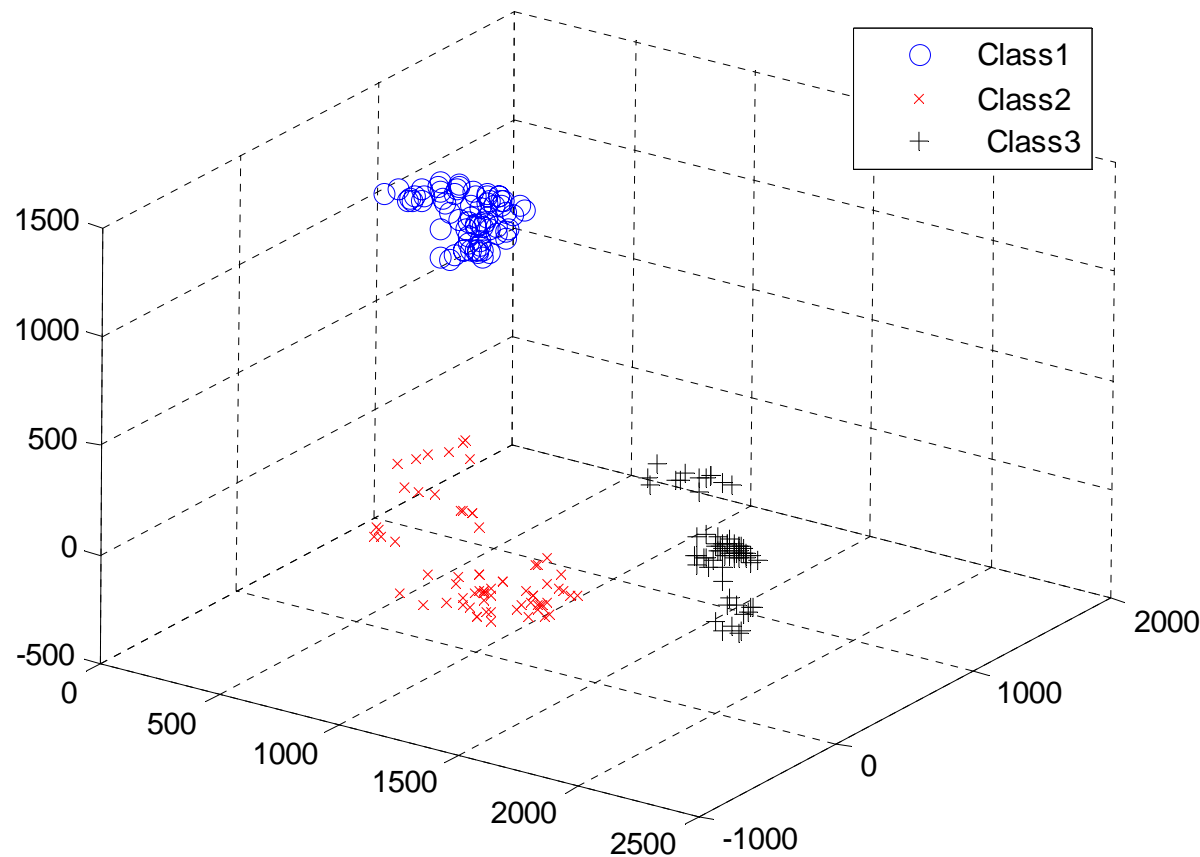
# The 3D linear subspace containing the projected training samples from class 1, 2 and 3



# How do we classify?

- Given a test face, we project onto this 3D linear subspace (the 3 eigenvectors) to get a 3d-vector.
- Then we can do a simple Nearest Neighbor search (euclidean or L2) distance to all the stored training vectors to find which is closest to and label the data.

Test: We projected all 75 images per person in the 3D PCA subspace



- We can see the 3 data classes clusters.



# PCA Variants

- What I described here was “Global PCA”  
i.e. PCA was performed on all classes to find a universal linear subspace.
- However we can also do Individual PCA (IPCA),  
i.e. build a subspace for each class using the images from that class.
  - How do we test?
  - We measure the reconstruction error between the test image and the reconstructed image from each subspace.
  - We label the test image with the subspace/basis that yielded the smallest reconstruction error.

# Pros & Cons?

- IPCA you need to have more data for each person (won't work with just 1 image/person).
- However, you don't need to re-train the system everytime you add a person in the database (as with Global PCA).